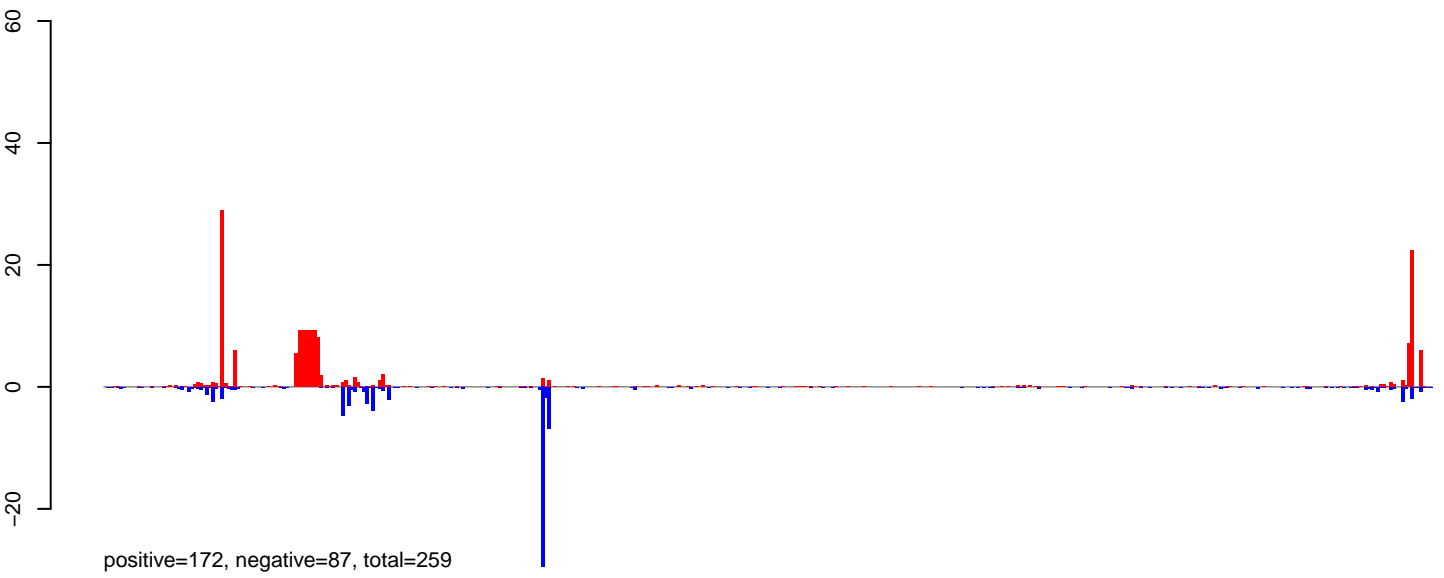
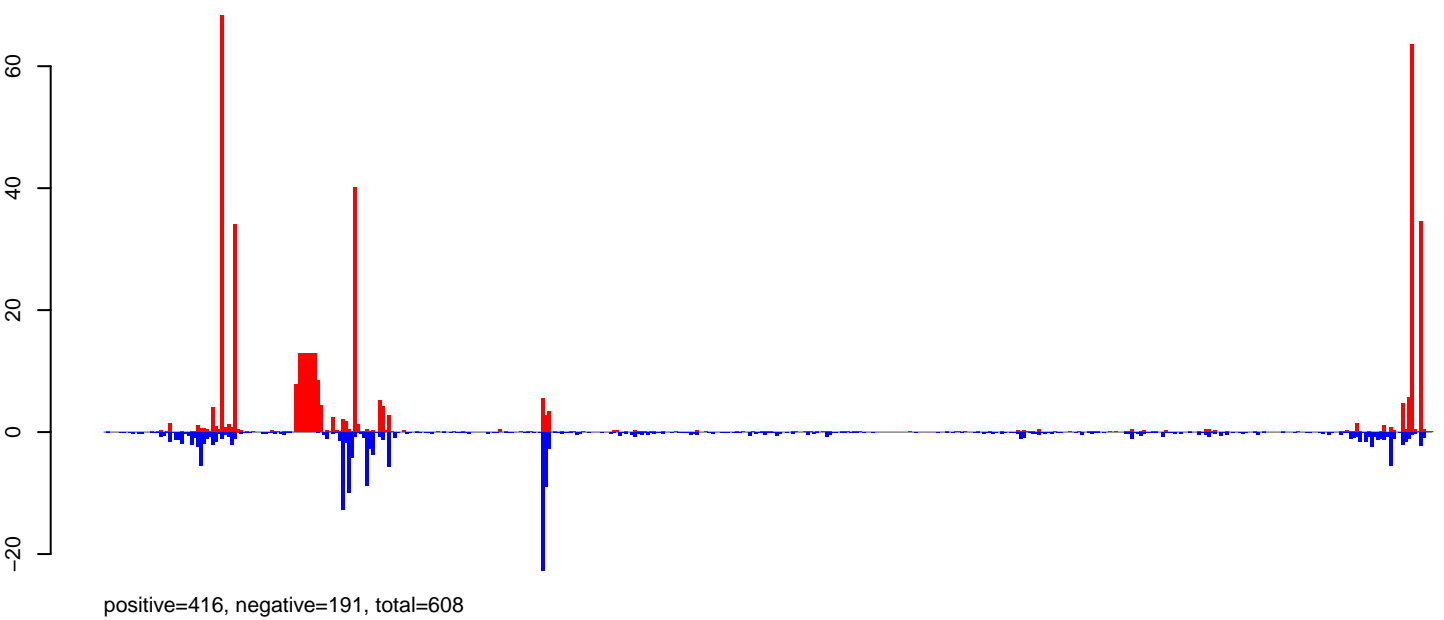


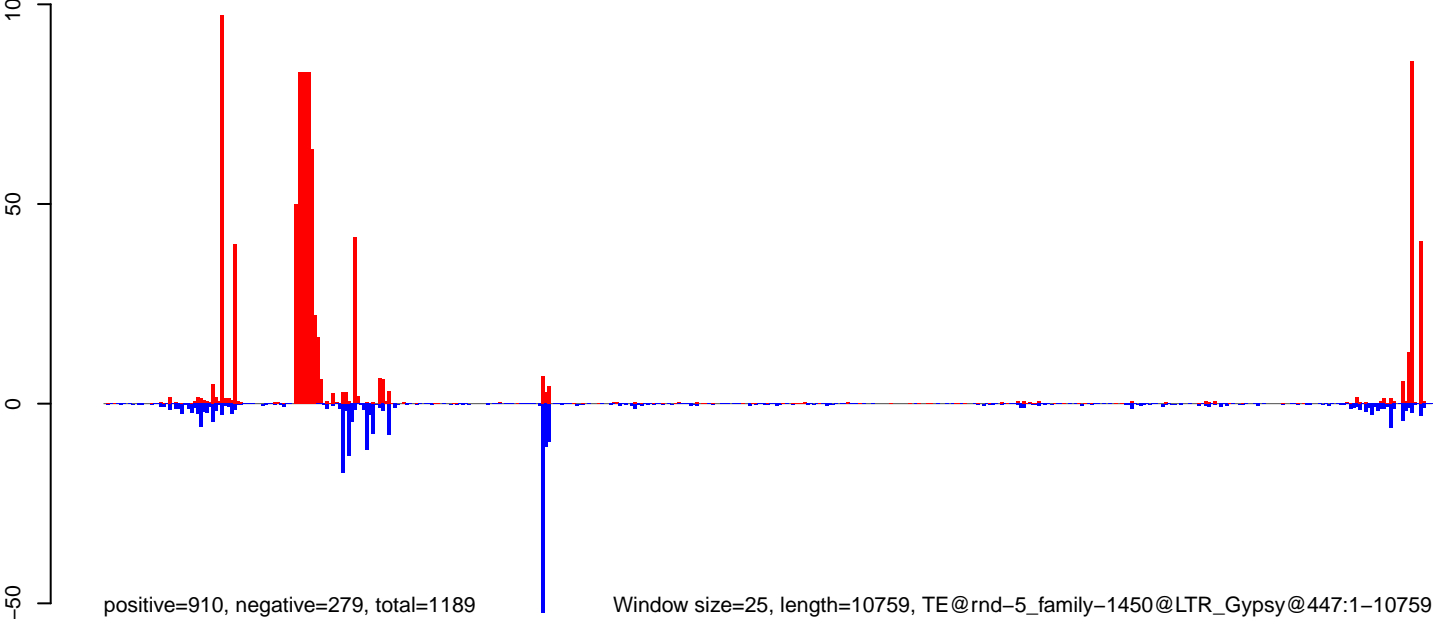
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

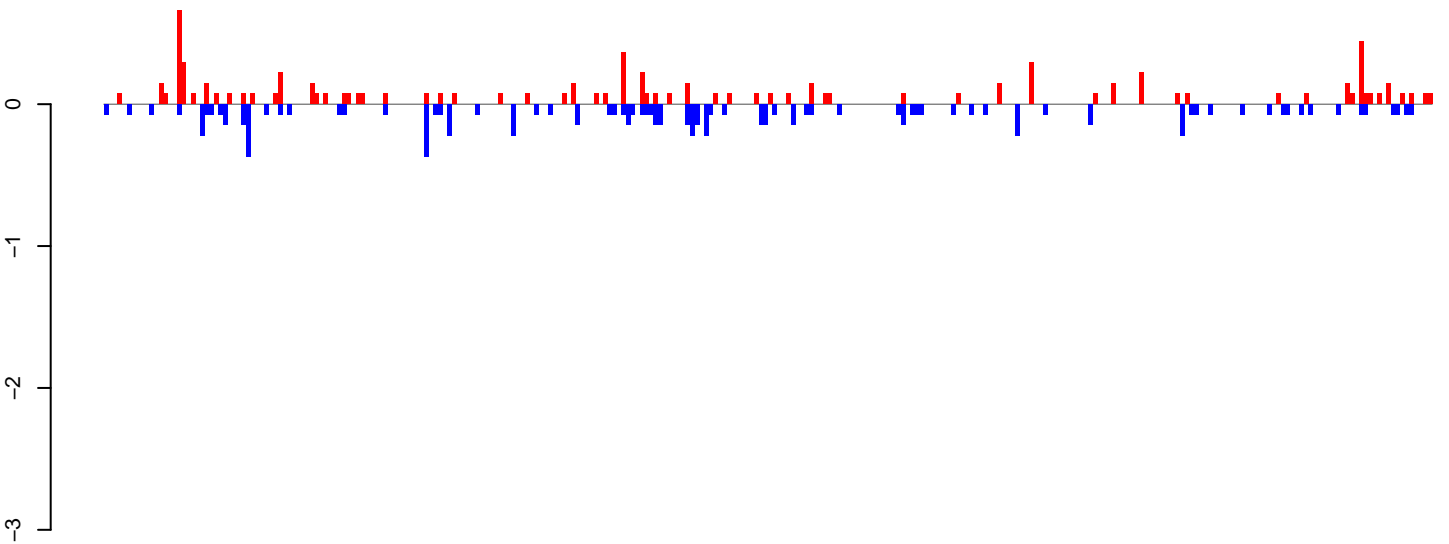


CuQuin_Testes_NL.rep



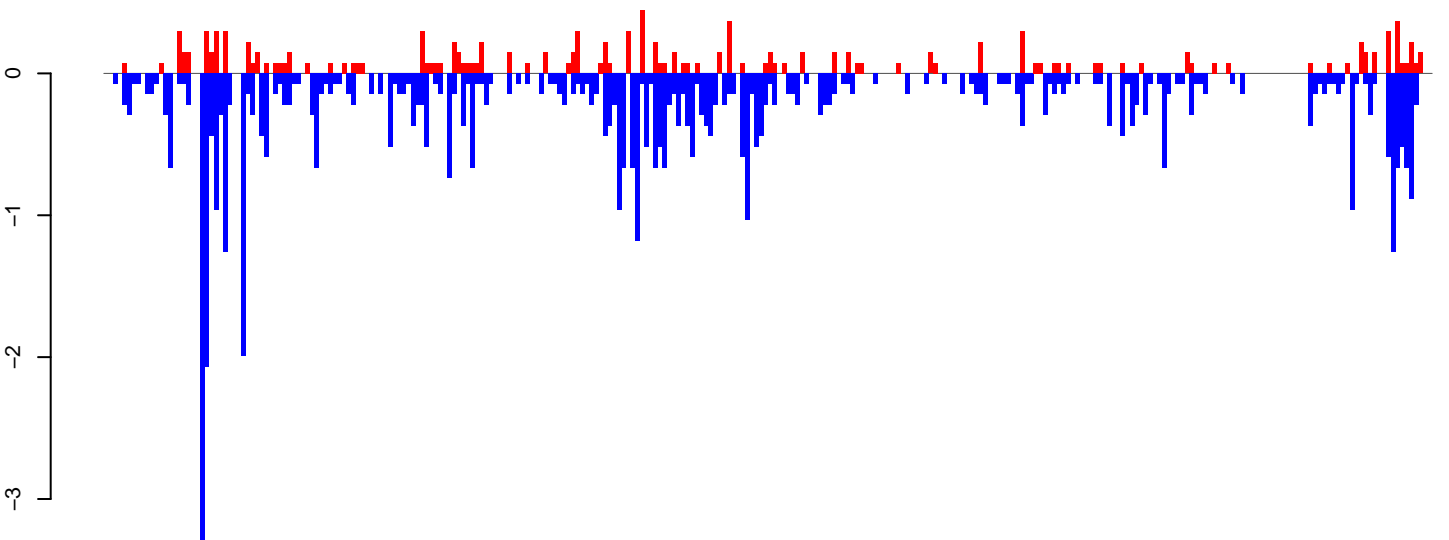
0 2000 4000 6000 8000 10000

CuQuin_Testes_NL.18_23.rep



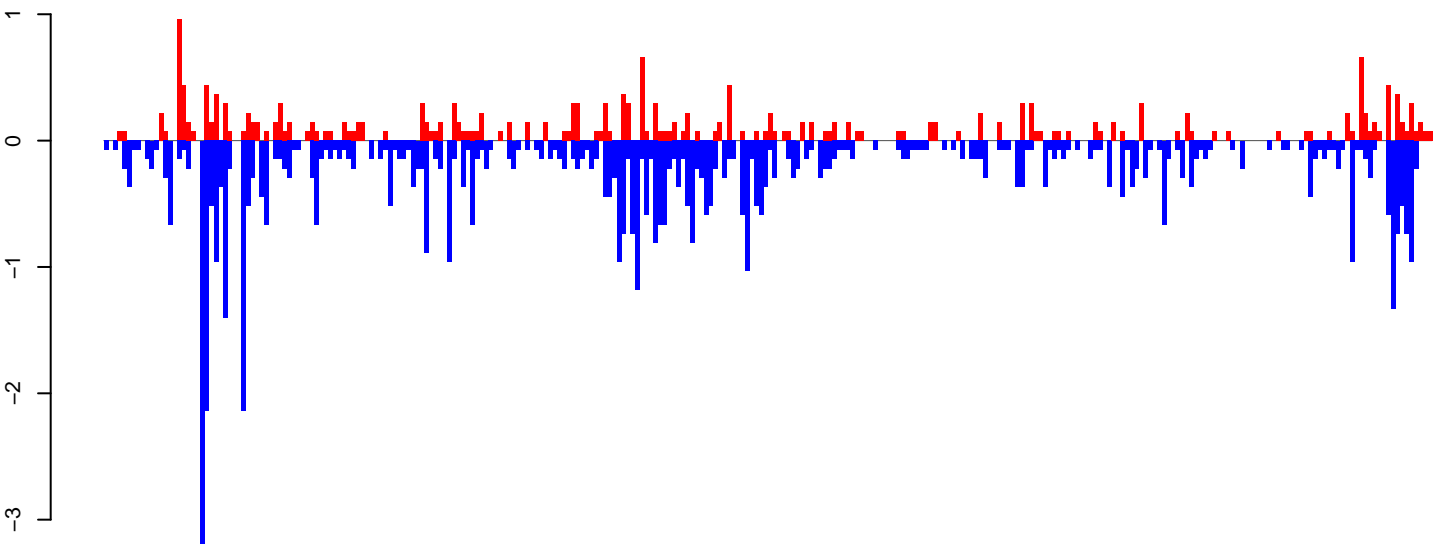
positive=8, negative=8, total=16

CuQuin_Testes_NL.24_35.rep



positive=13, negative=54, total=67

CuQuin_Testes_NL.rep

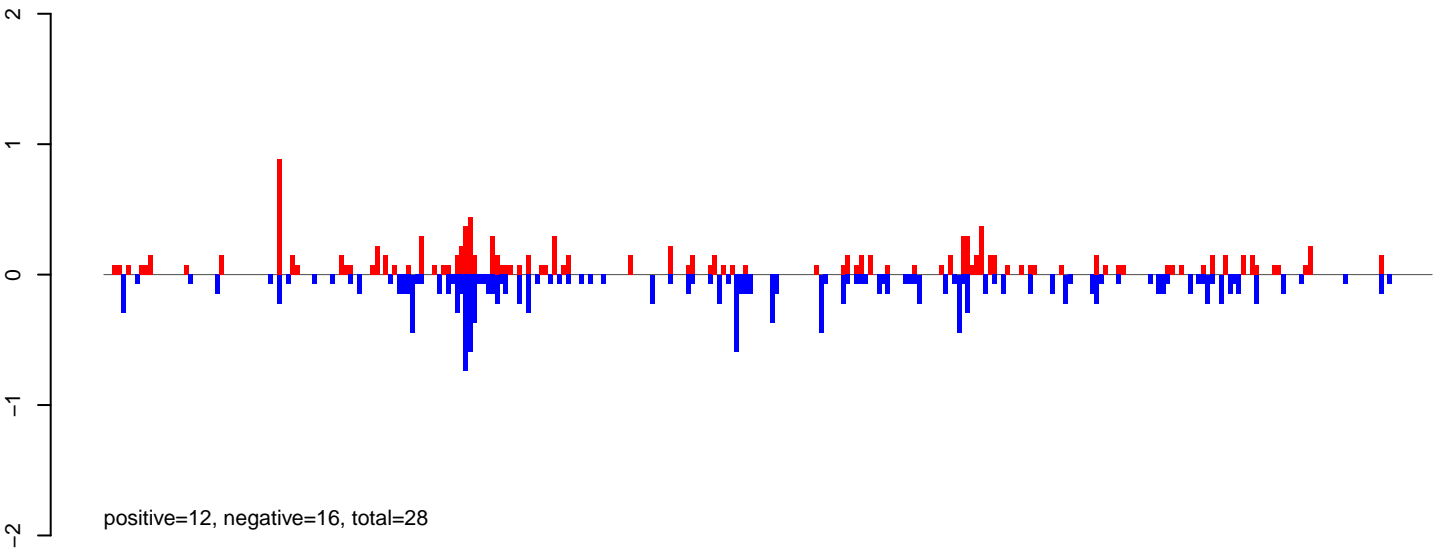


positive=21, negative=63, total=83

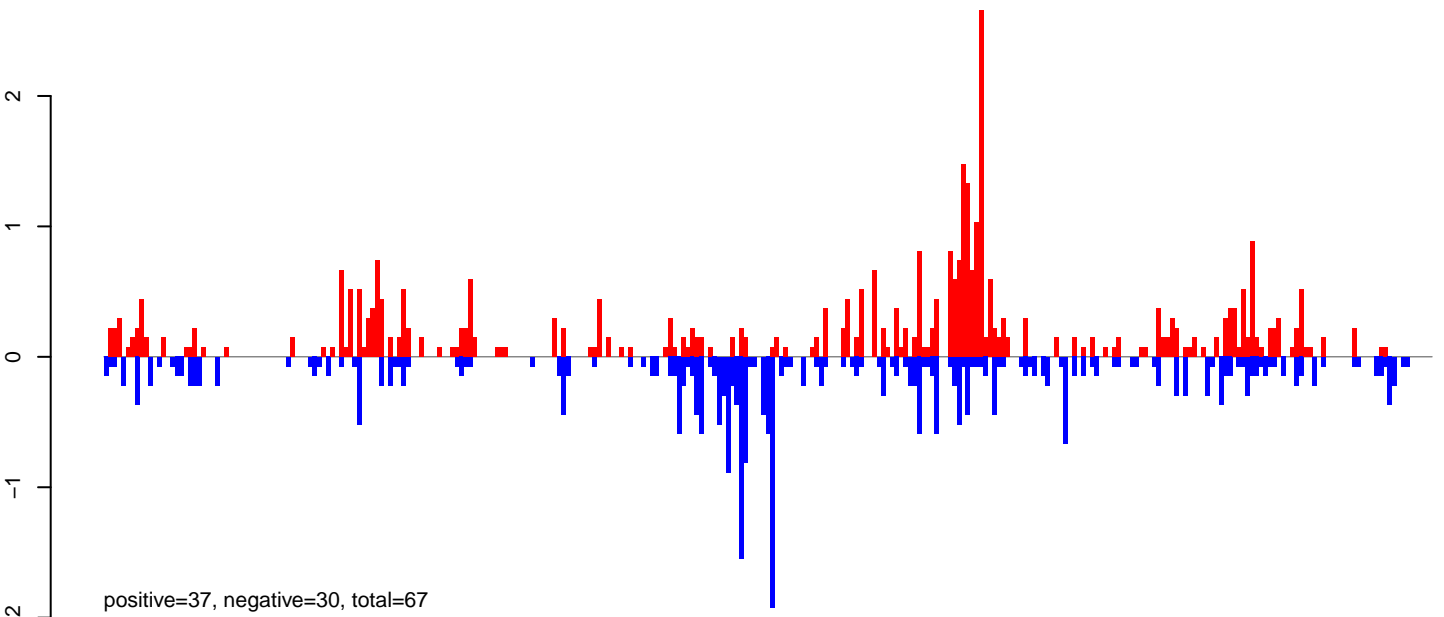
Window size=25, length=7249, TE@rnd-5_family-173@LTR_Pao@2997:1-7249

0 2000 4000 6000

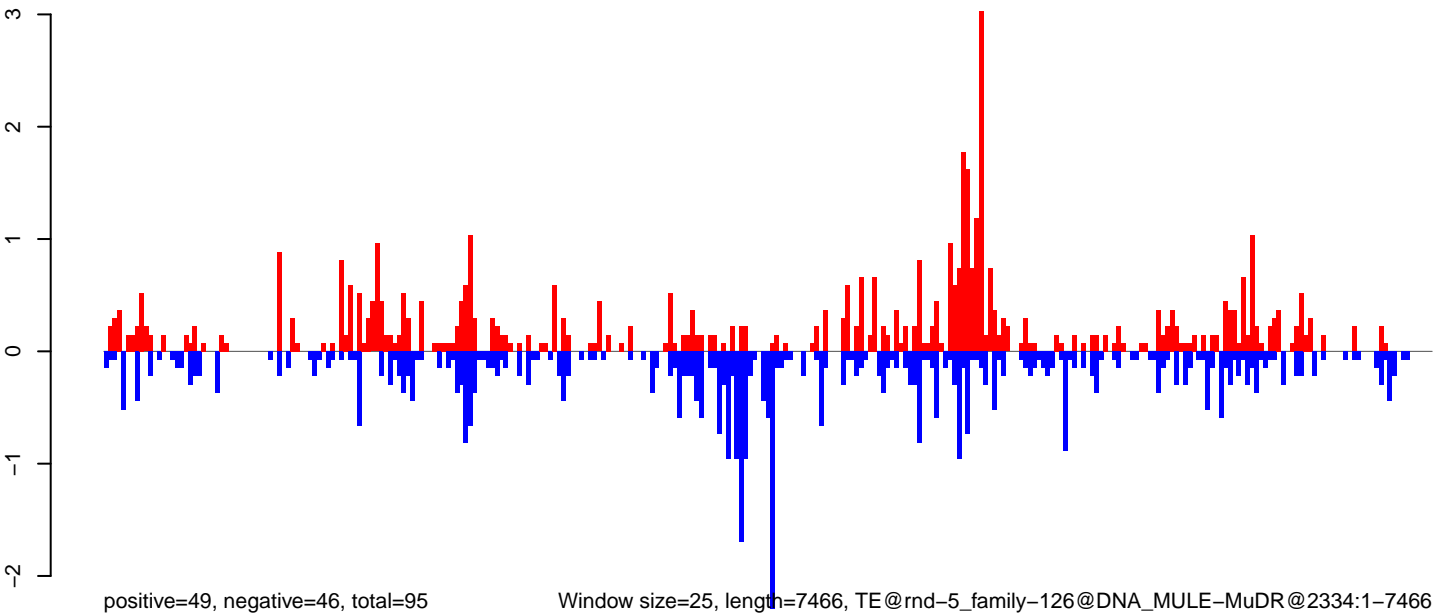
CuQuin_Testes_NL.18_23.rep



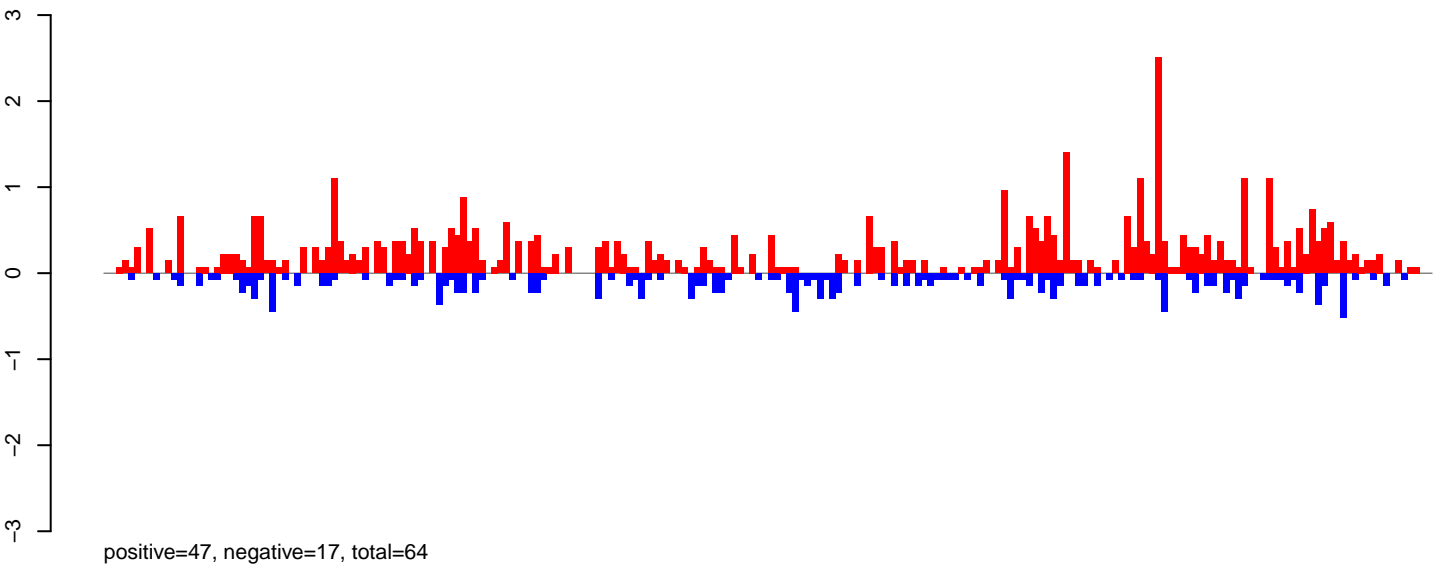
CuQuin_Testes_NL.24_35.rep



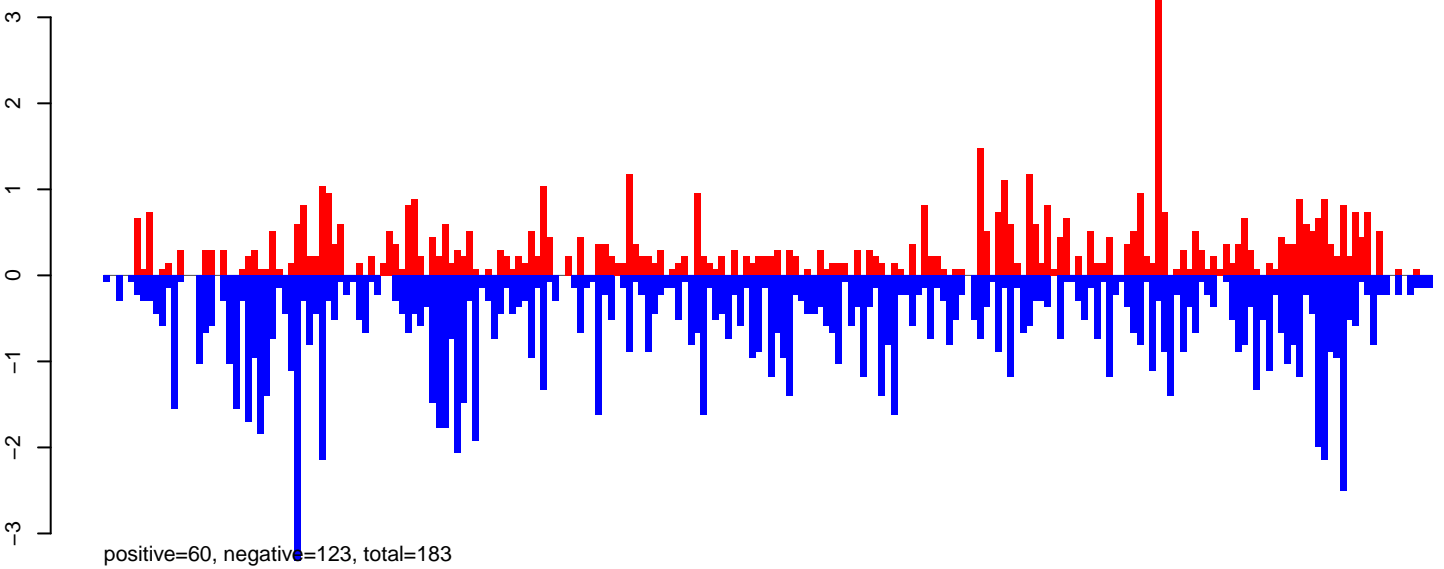
CuQuin_Testes_NL.rep



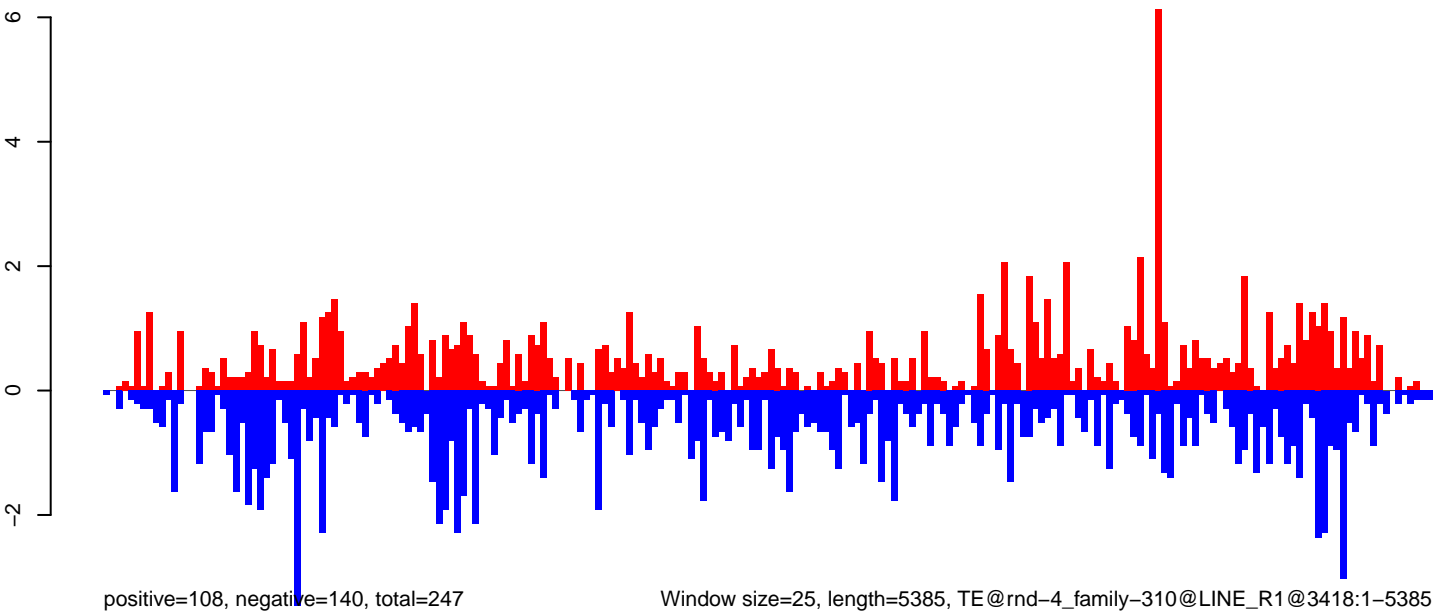
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

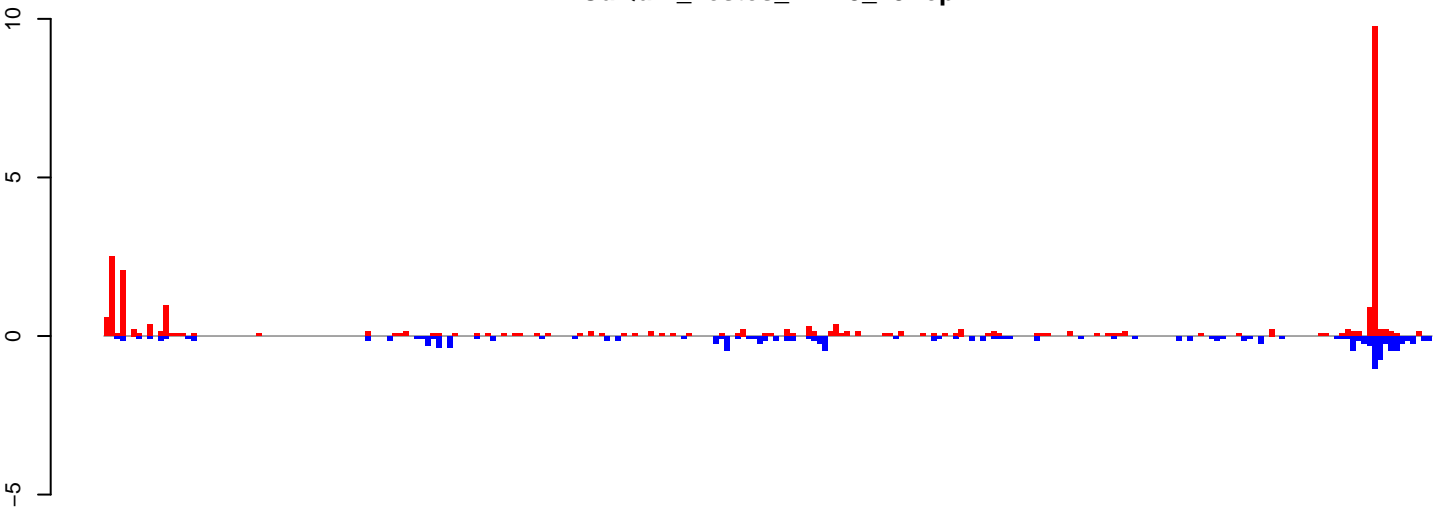


CuQuin_Testes_NL.rep



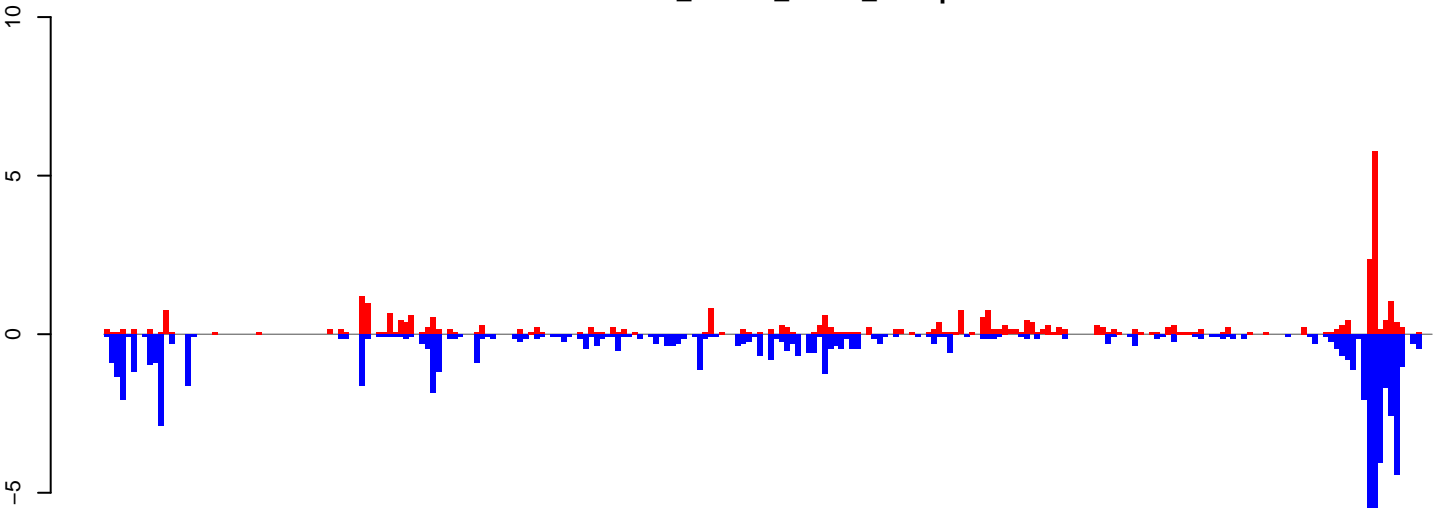
0 1000 2000 3000 4000 5000

CuQuin_Testes_NL.18_23.rep



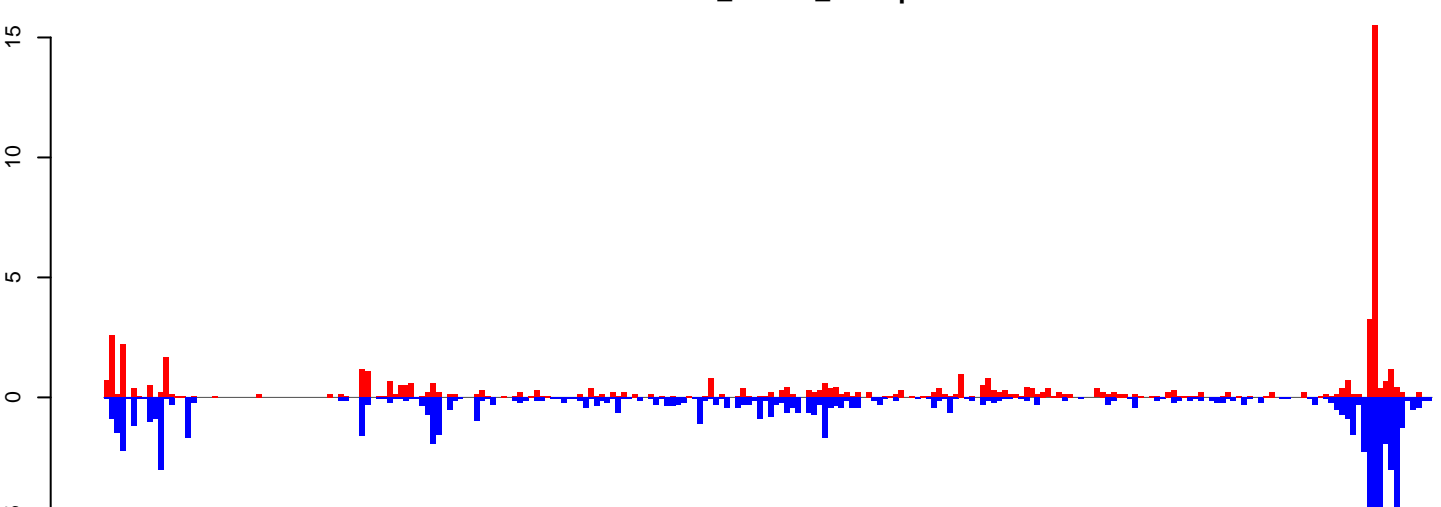
positive=26, negative=13, total=39

CuQuin_Testes_NL.24_35.rep



positive=34, negative=76, total=110

CuQuin_Testes_NL.rep

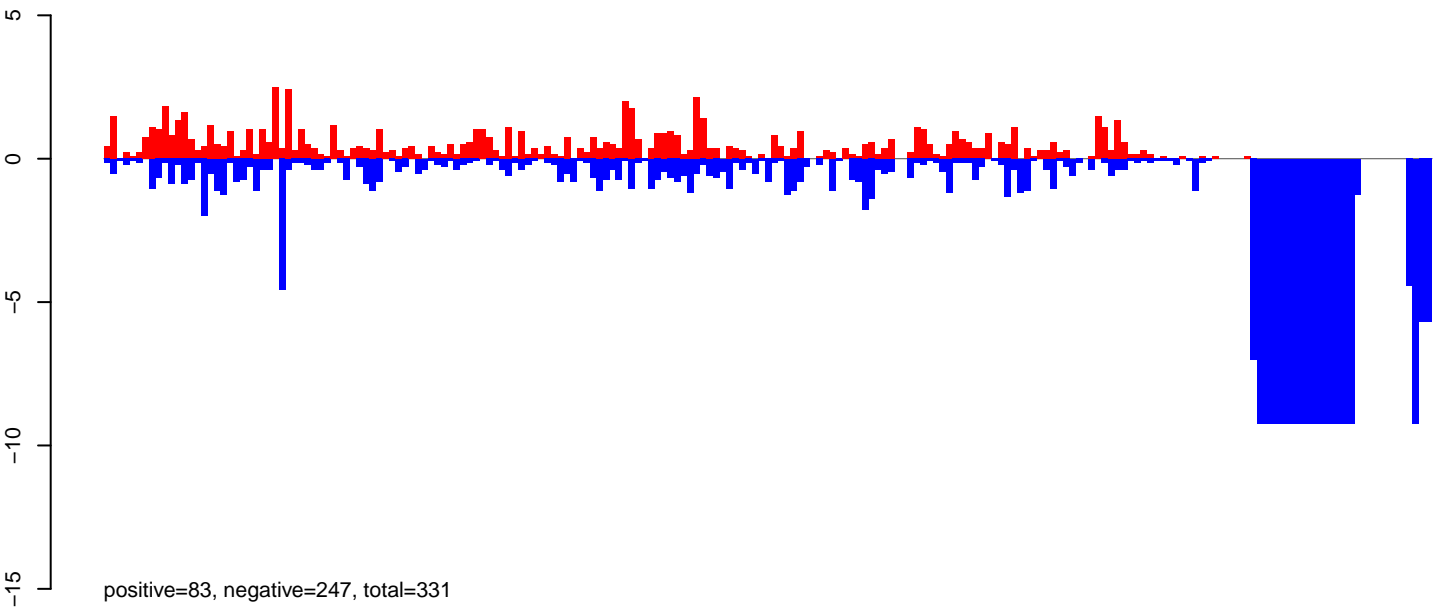


positive=60, negative=90, total=150

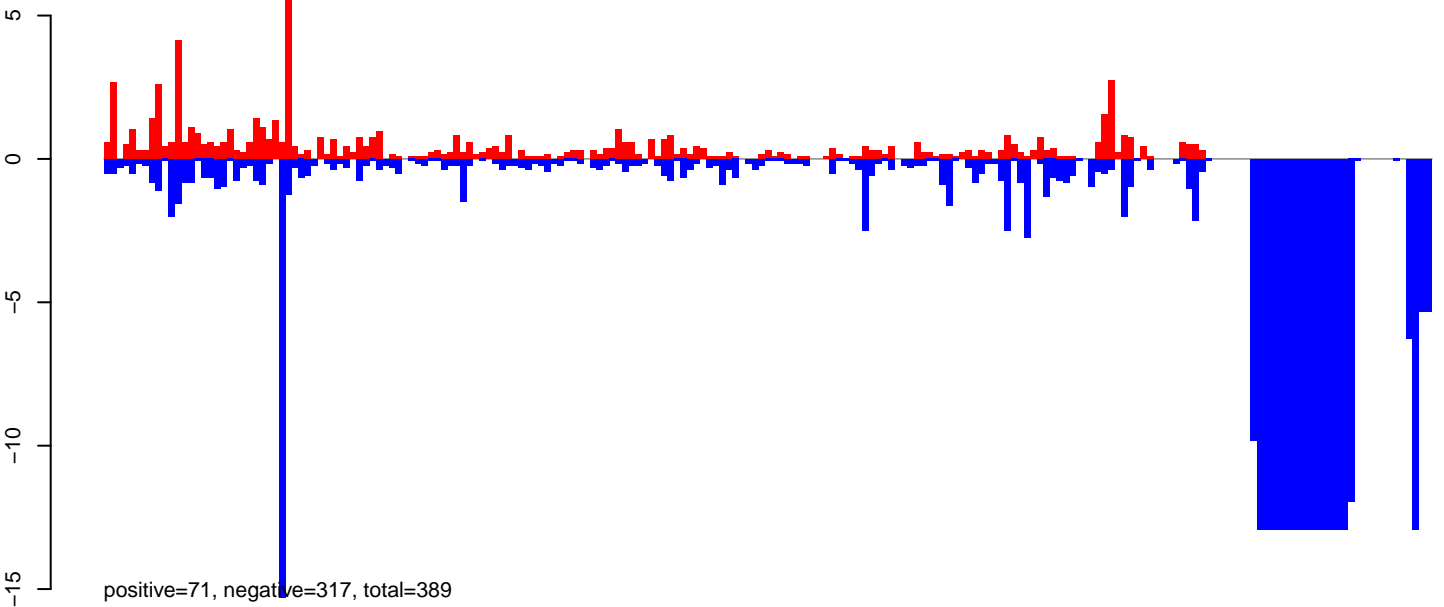
Window size=25, length=6077, TE@rnd-5_family-378@LINE_I@1063:1-6077

0 1000 2000 3000 4000 5000 6000

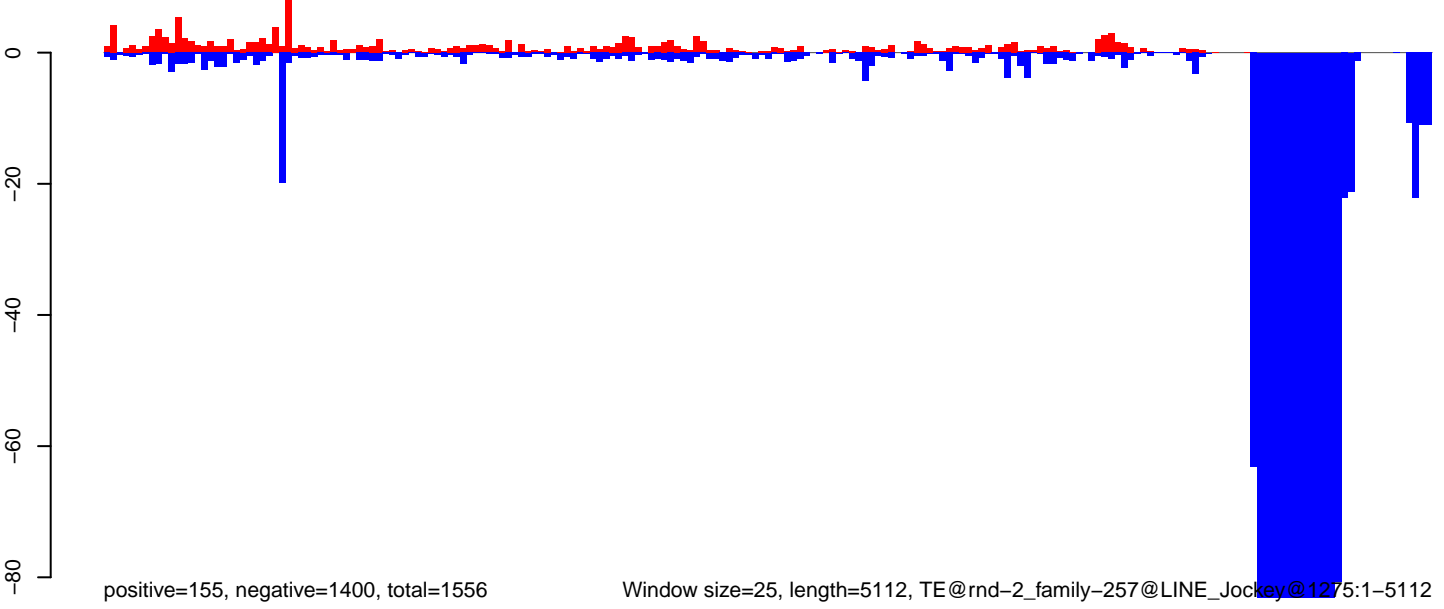
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



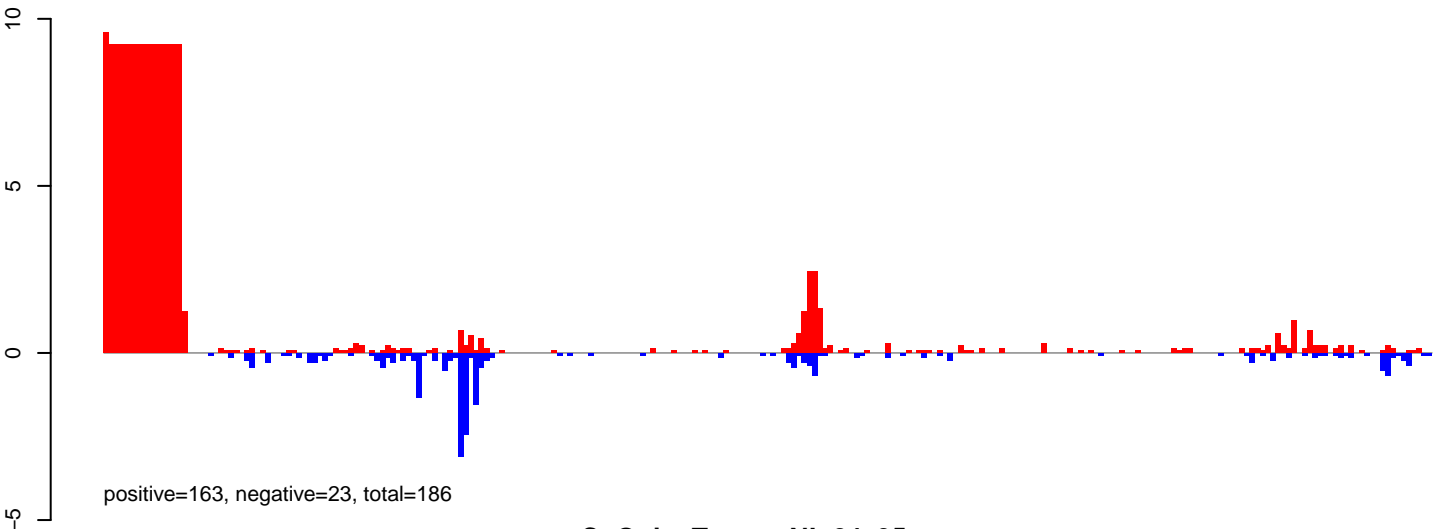
CuQuin_Testes_NL.rep



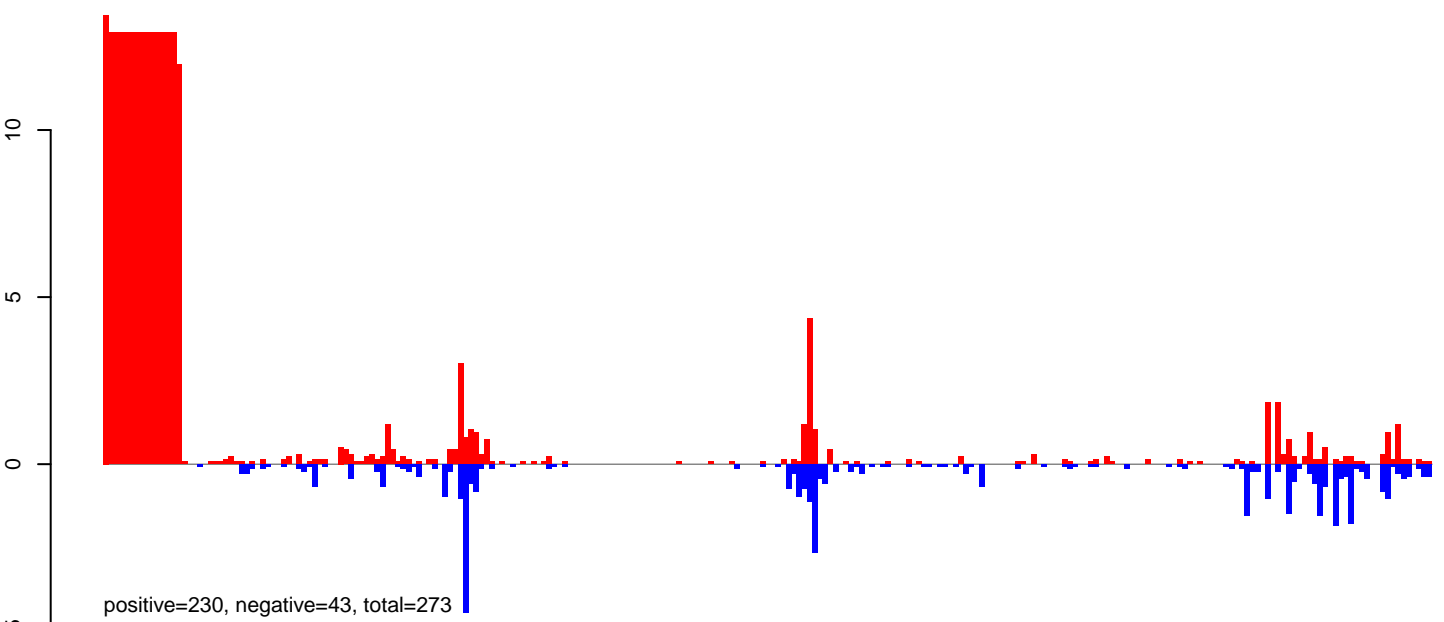
Window size=25, length=5112, TE@rnd-2_family-257@LINE_Jockey@1275:1-5112

0 1000 2000 3000 4000 5000

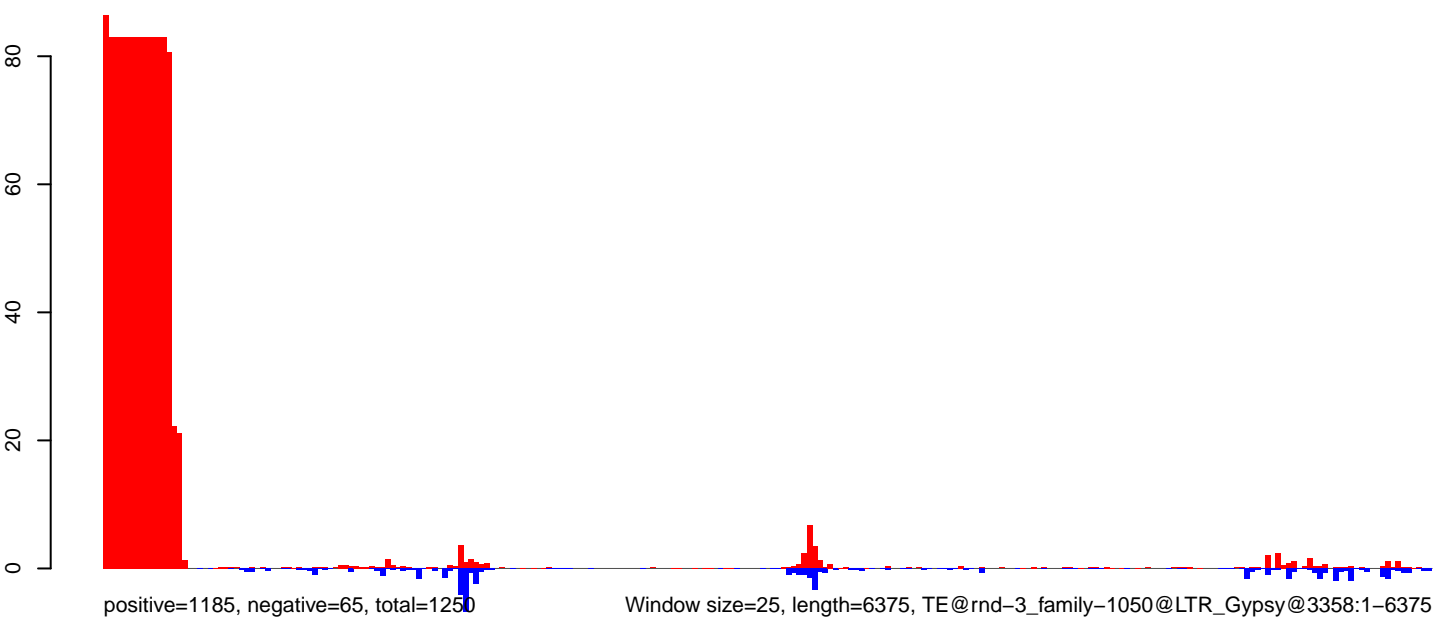
CuQuin_Testes_NL.18_23.rep



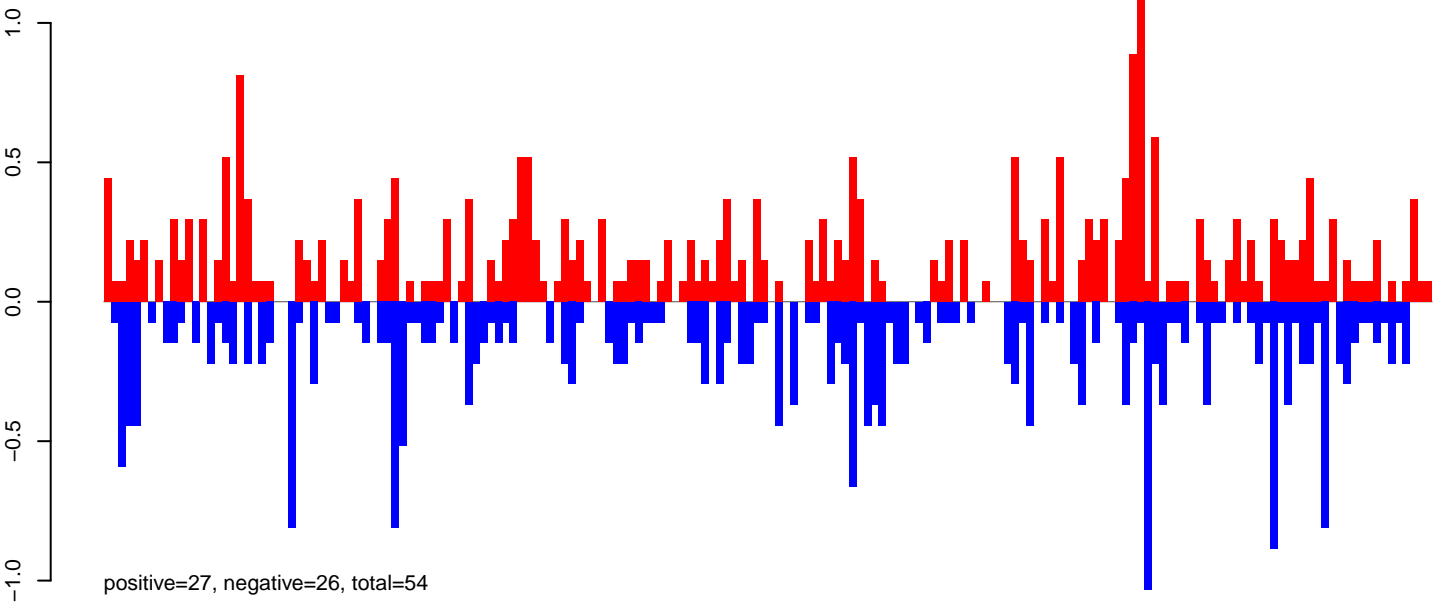
CuQuin_Testes_NL.24_35.rep



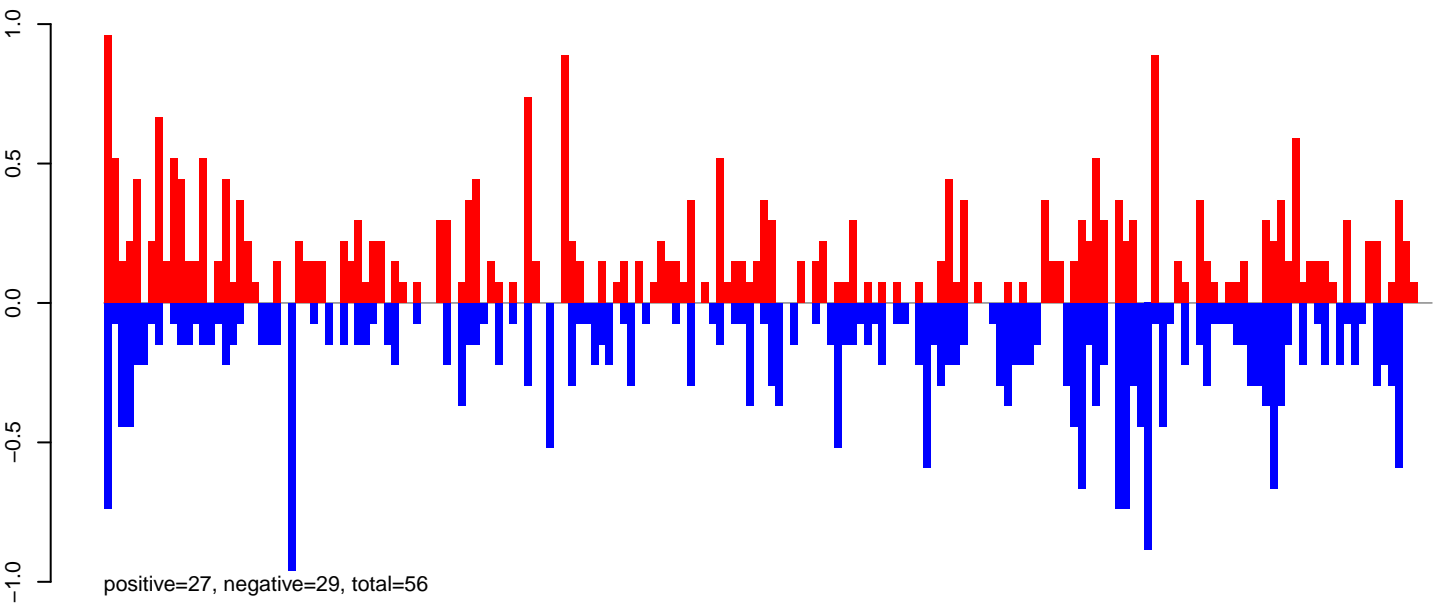
CuQuin_Testes_NL.rep



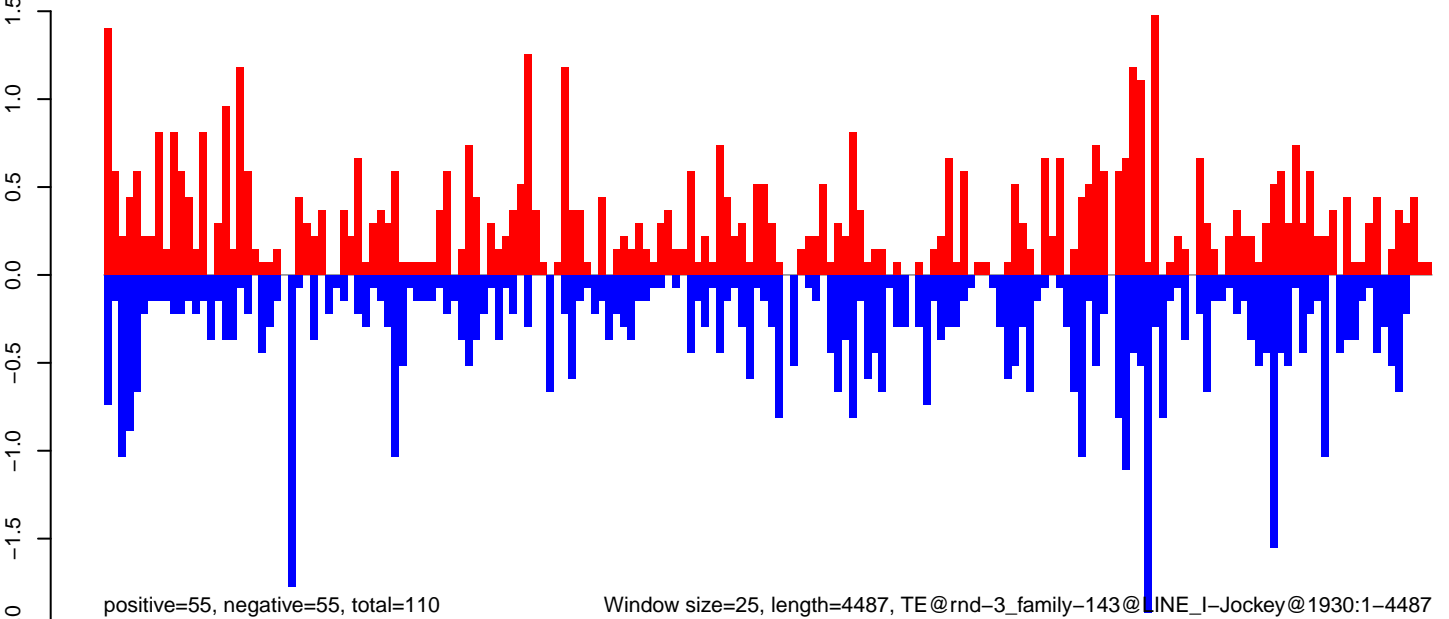
CuQuin_Testes_NL.18_23.rep



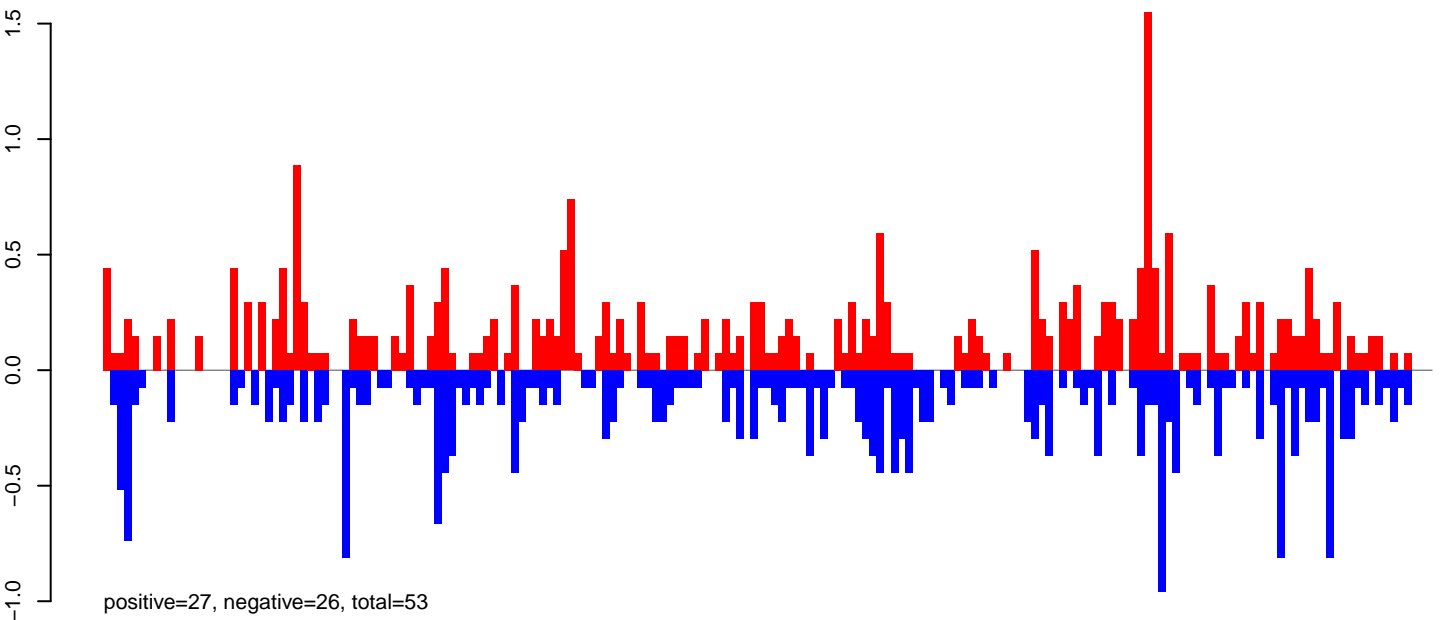
CuQuin_Testes_NL.24_35.rep



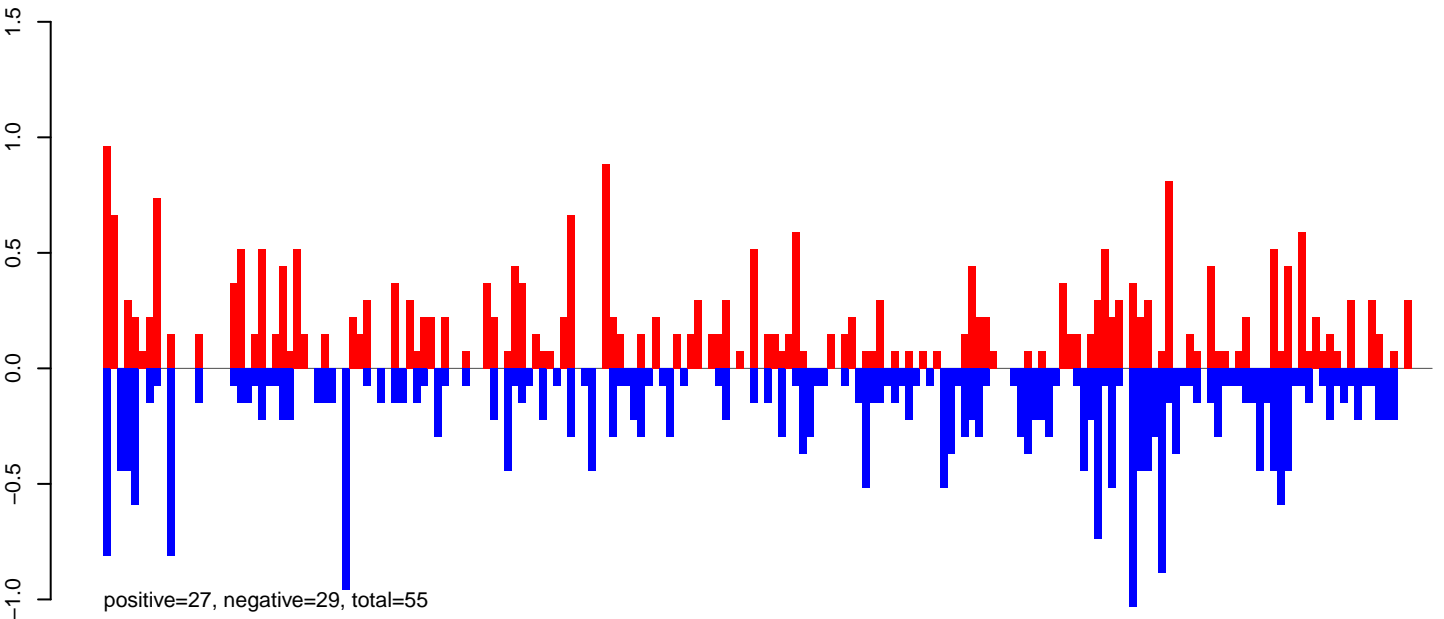
CuQuin_Testes_NL.rep



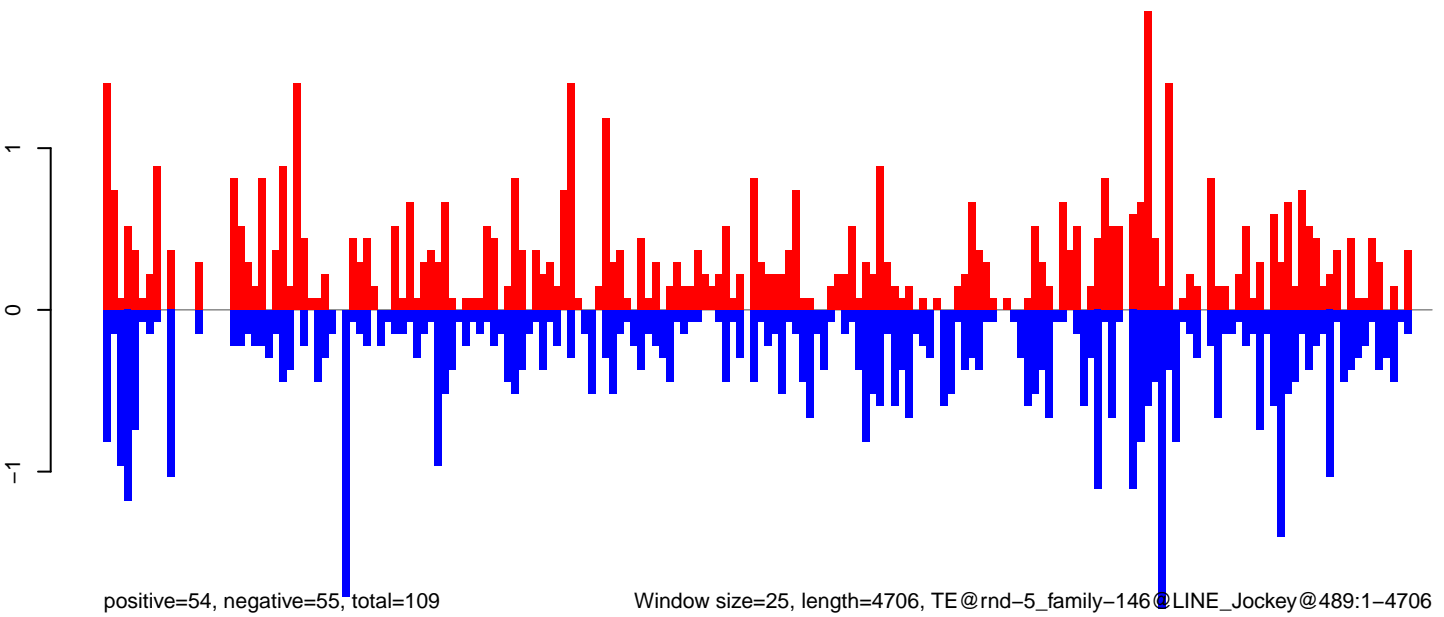
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



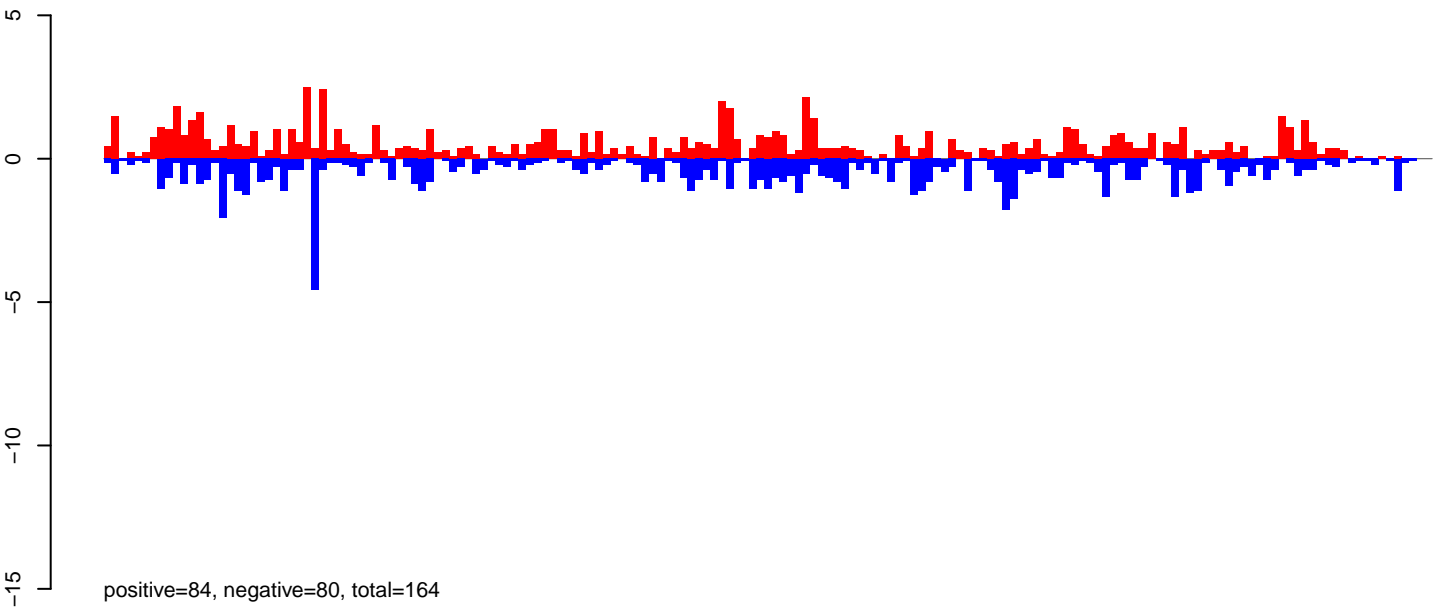
CuQuin_Testes_NL.rep



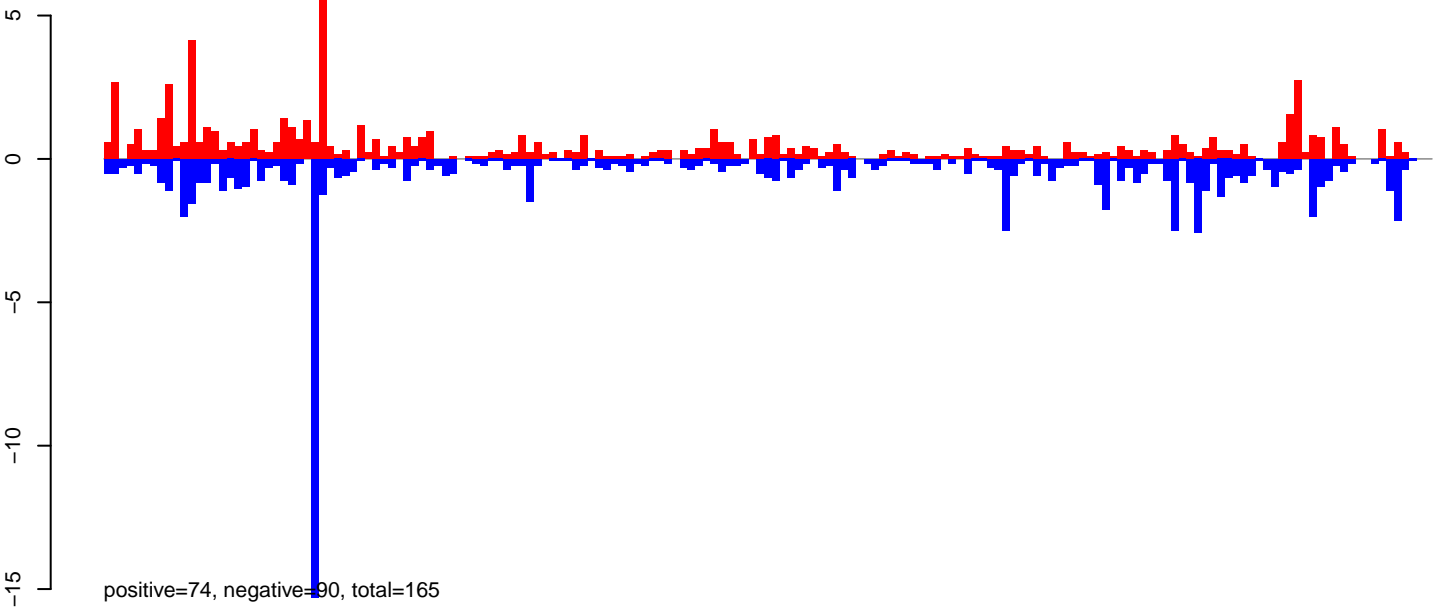
Window size=25, length=4706, TE@rnd-5_family-146@LINE_Jockey@489:1-4706

0 1000 2000 3000 4000

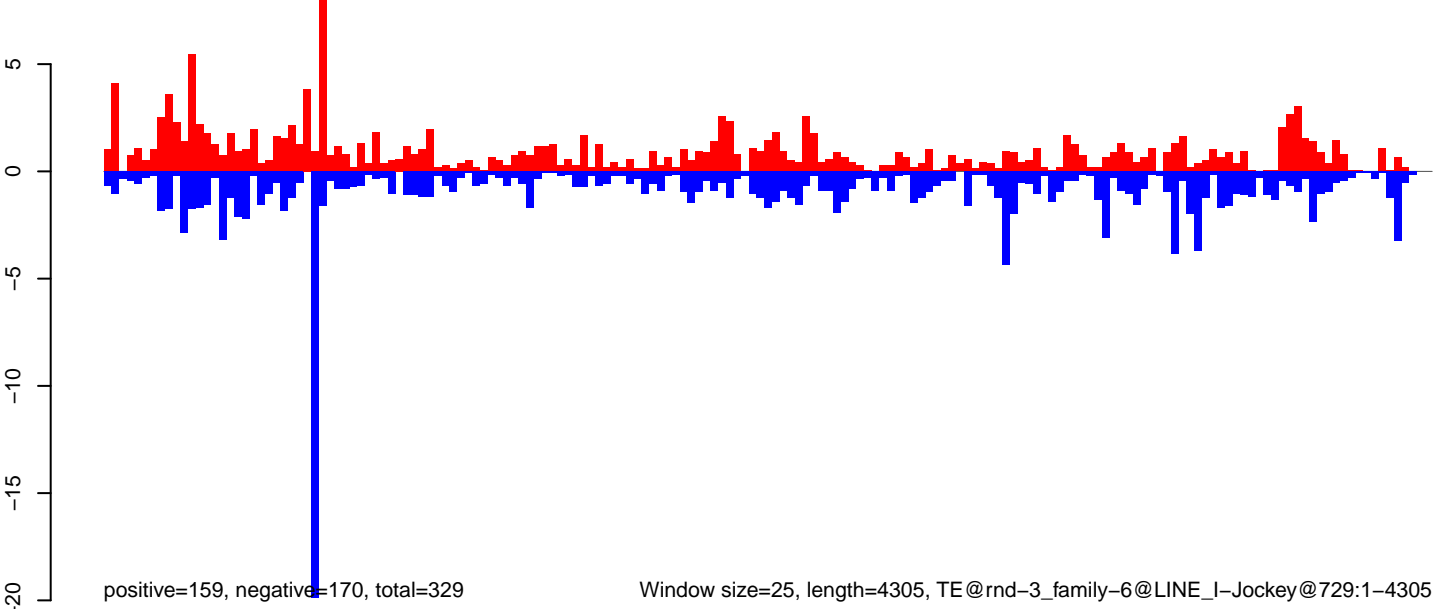
CuQuin_Testes_NL.18_23.rep



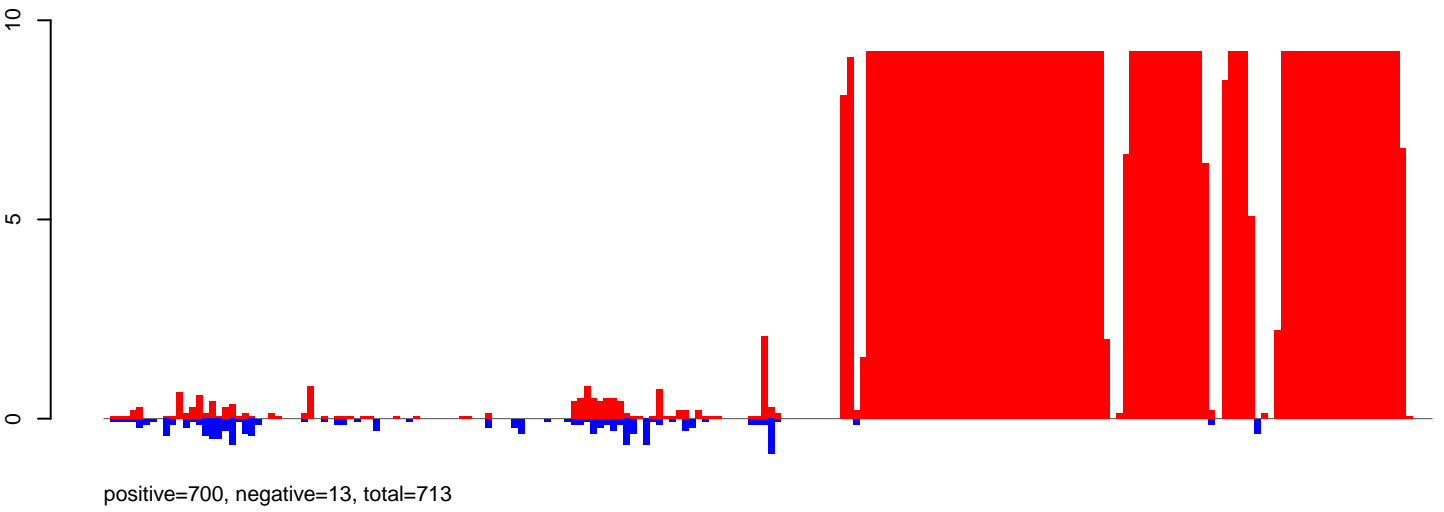
CuQuin_Testes_NL.24_35.rep



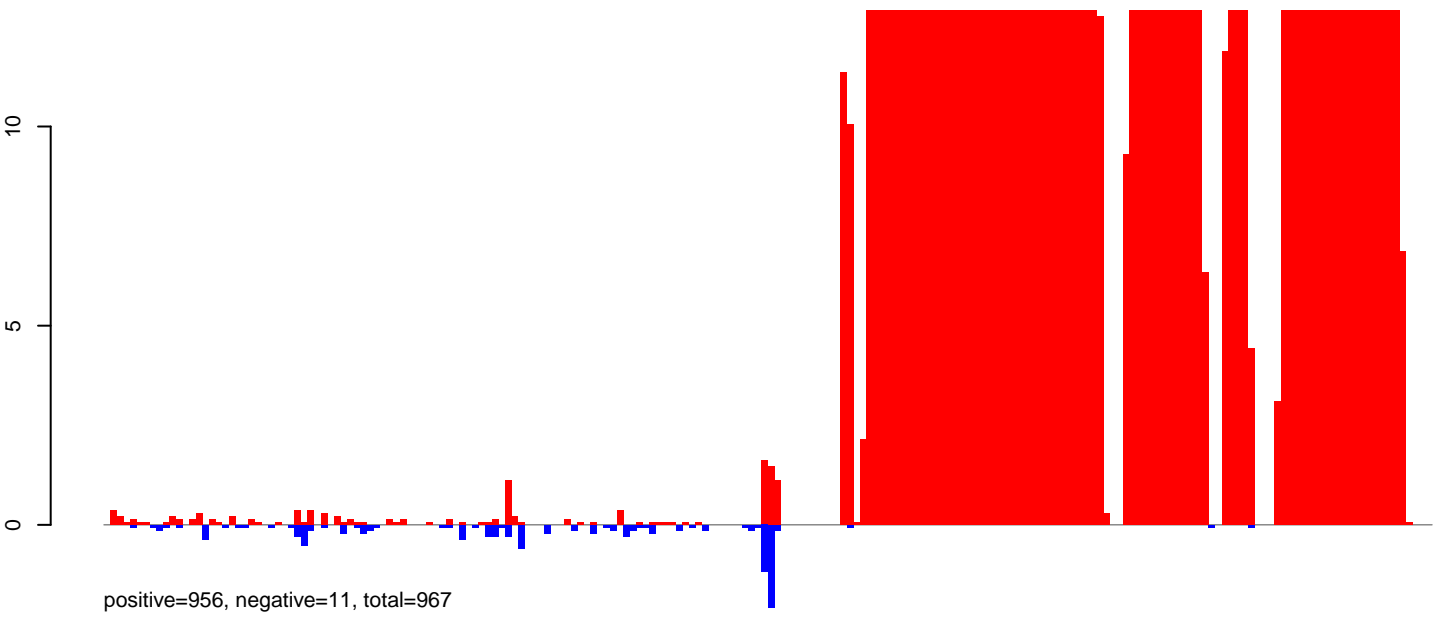
CuQuin_Testes_NL.rep



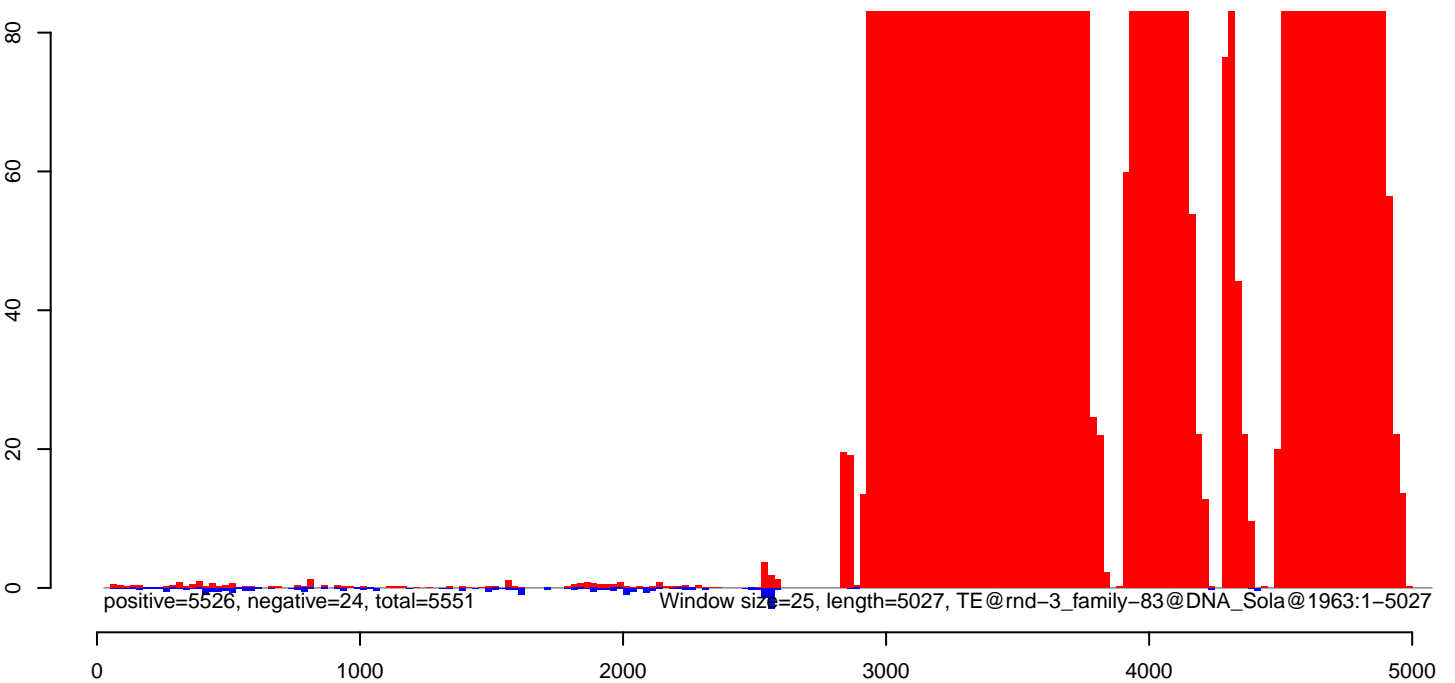
CuQuin_Testes_NL.18_23.rep



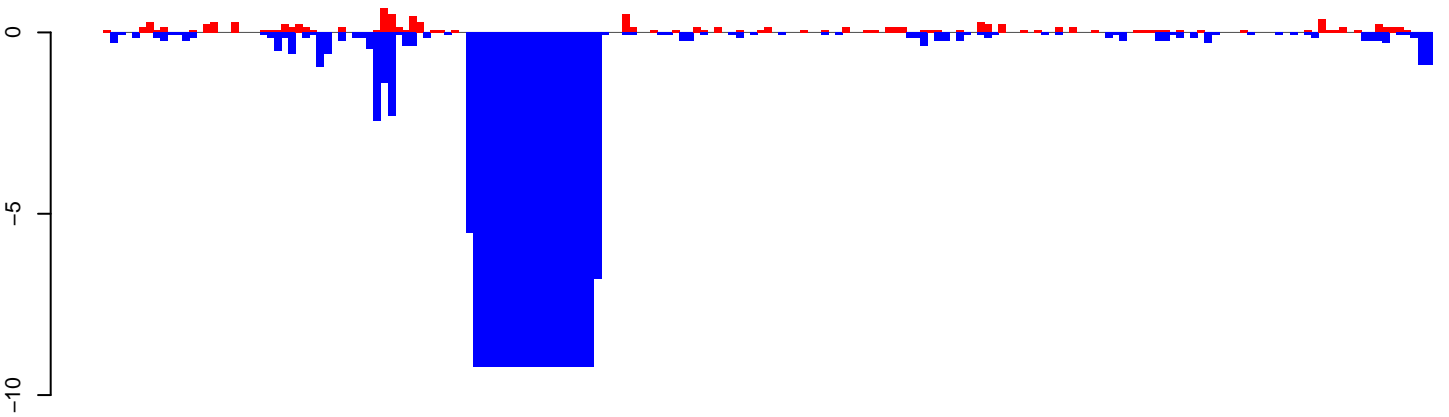
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

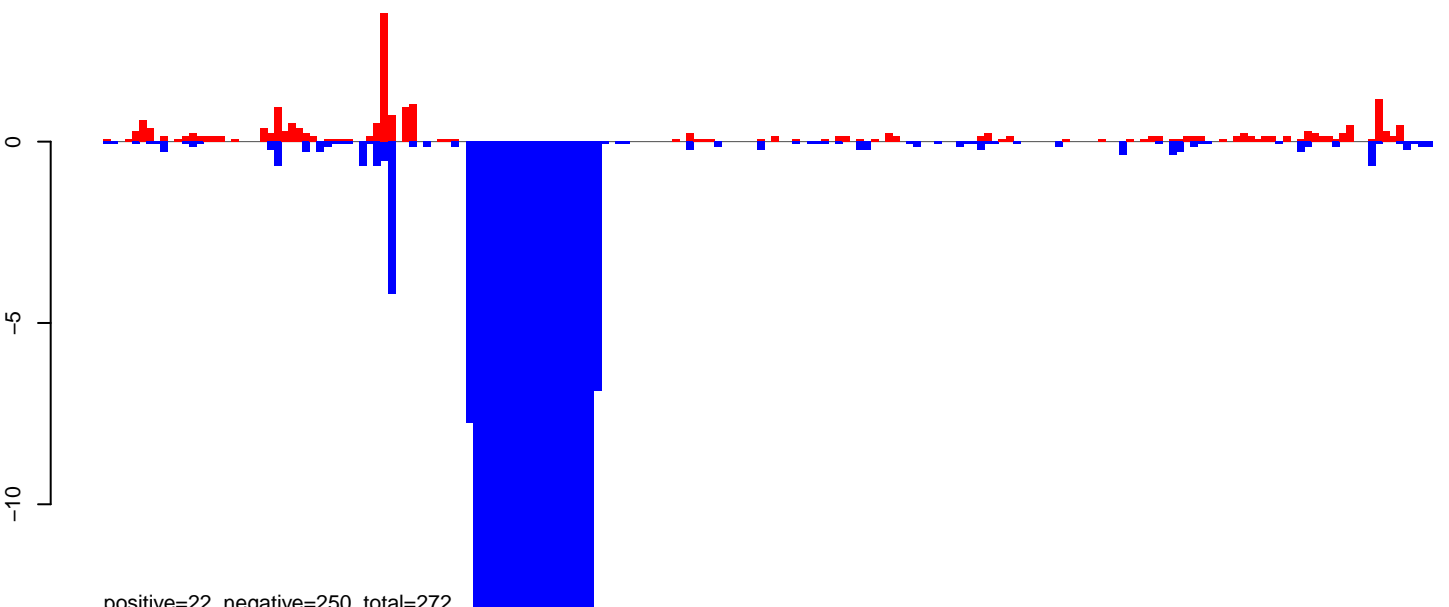


CuQuin_Testes_NL.18_23.rep



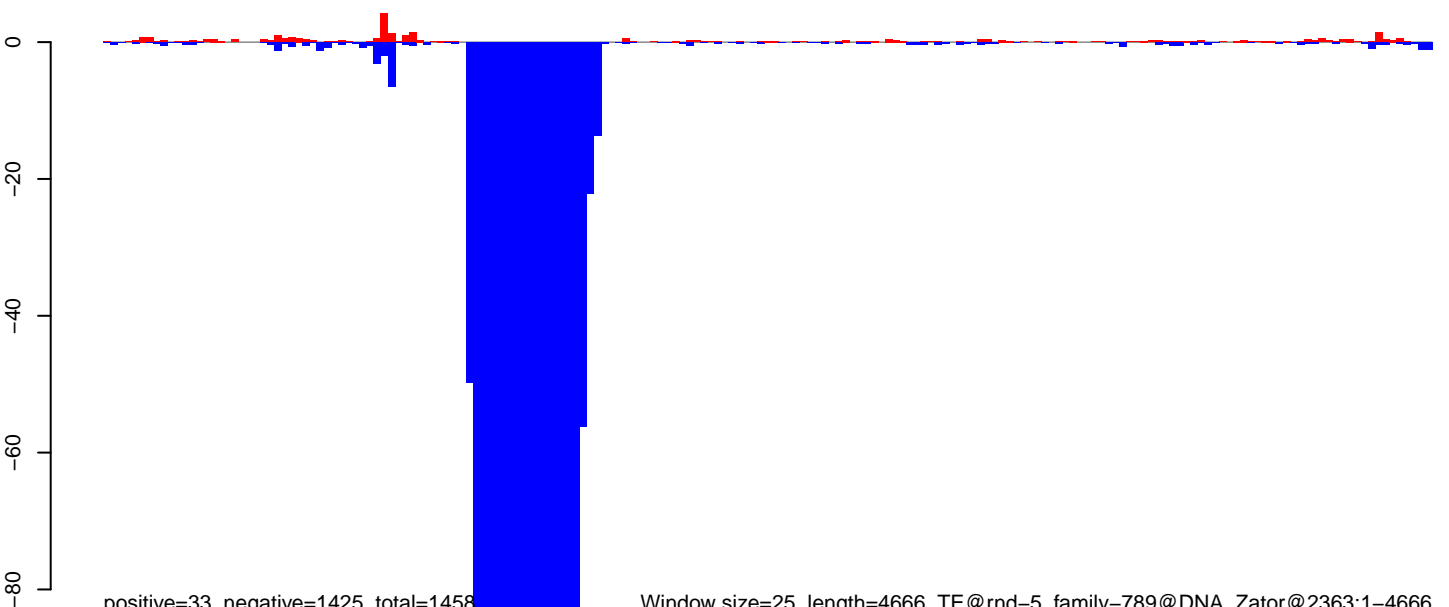
positive=11, negative=190, total=202

CuQuin_Testes_NL.24_35.rep



positive=22, negative=250, total=272

CuQuin_Testes_NL.rep

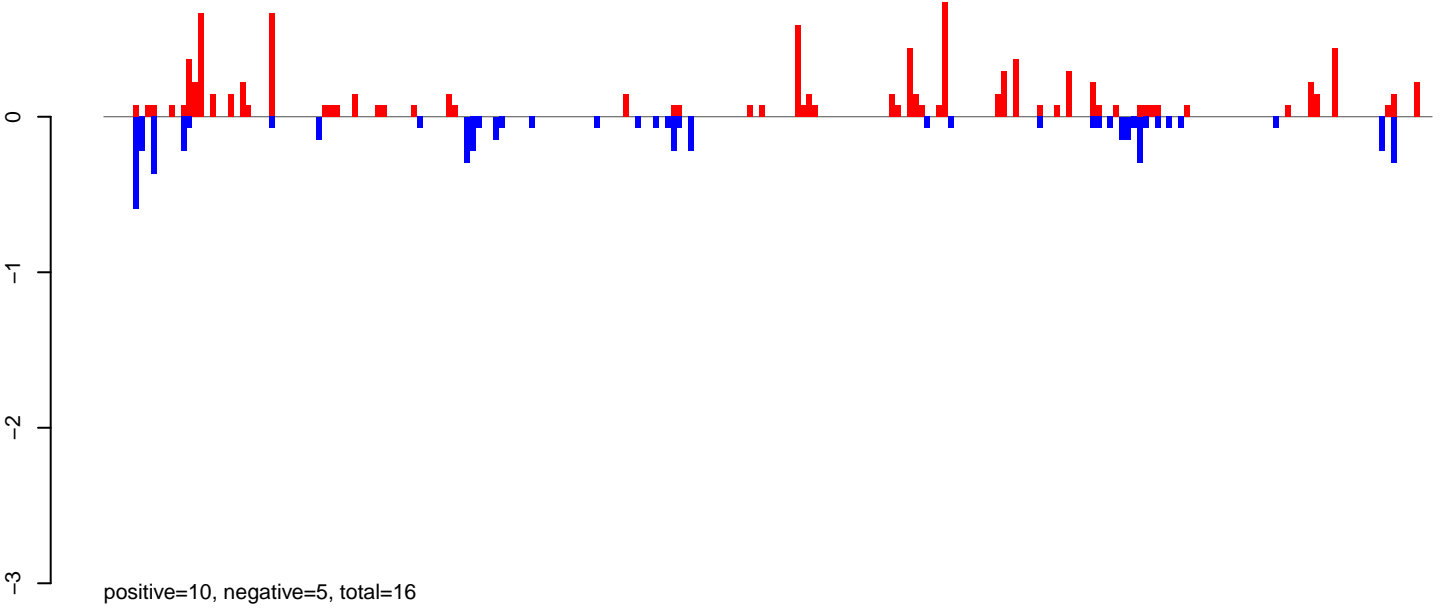


positive=33, negative=1425, total=1458

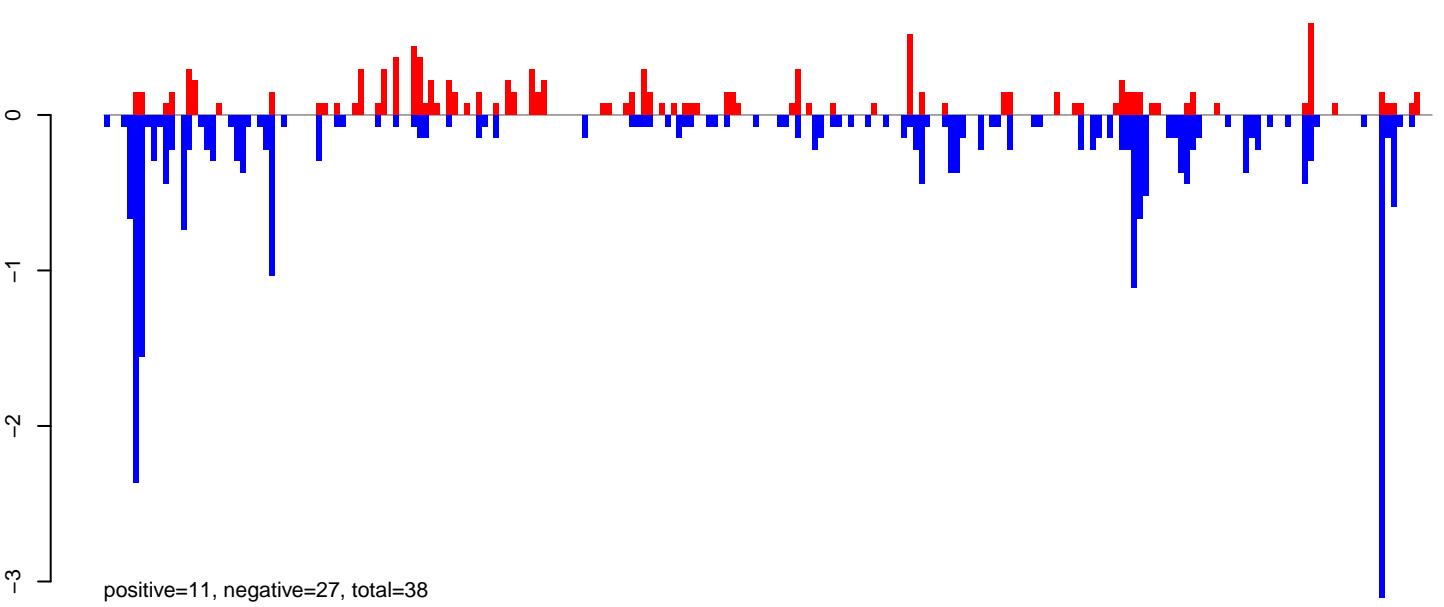
Window size=25, length=4666, TE@rnd-5_family-789@DNA_Zator@2363:1-4666

0 1000 2000 3000 4000

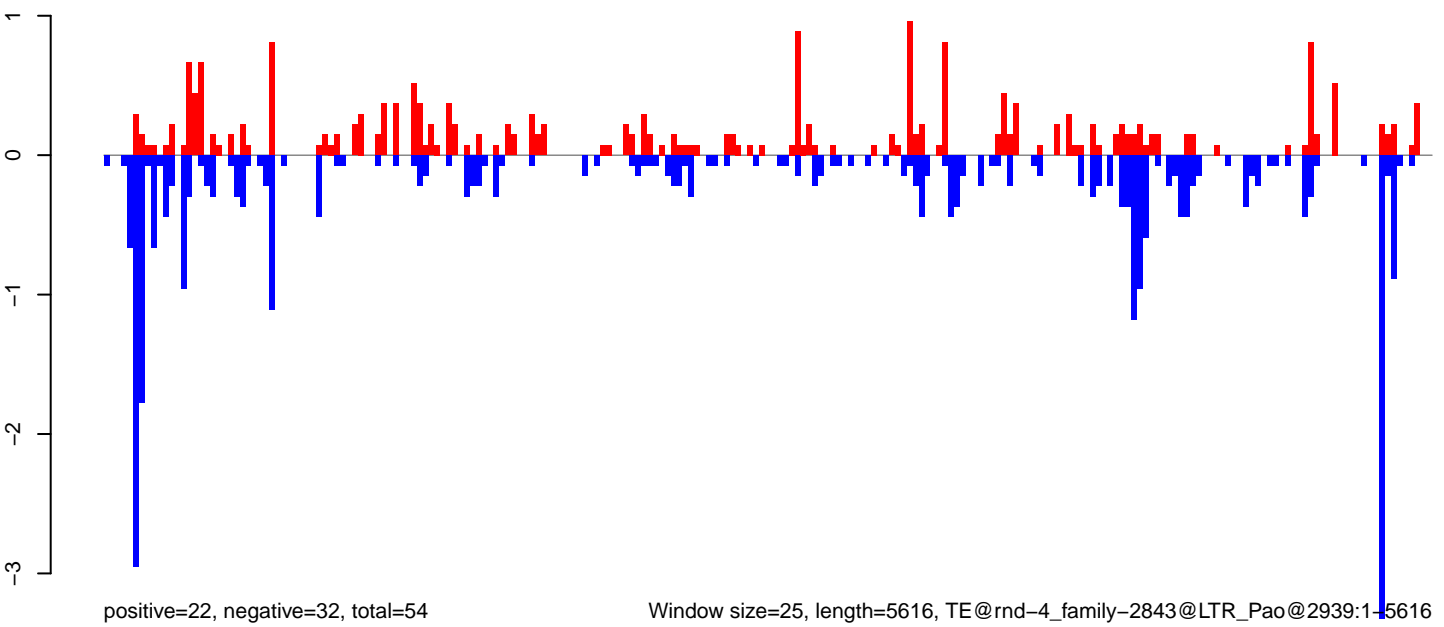
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



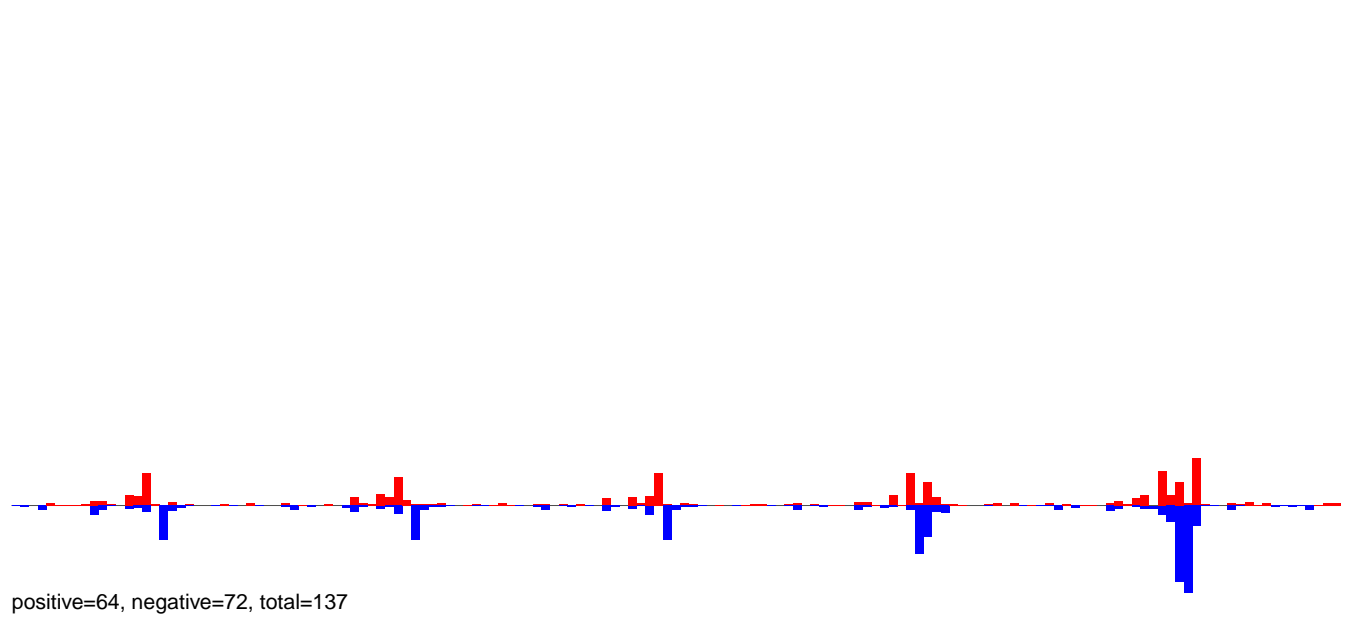
CuQuin_Testes_NL.rep



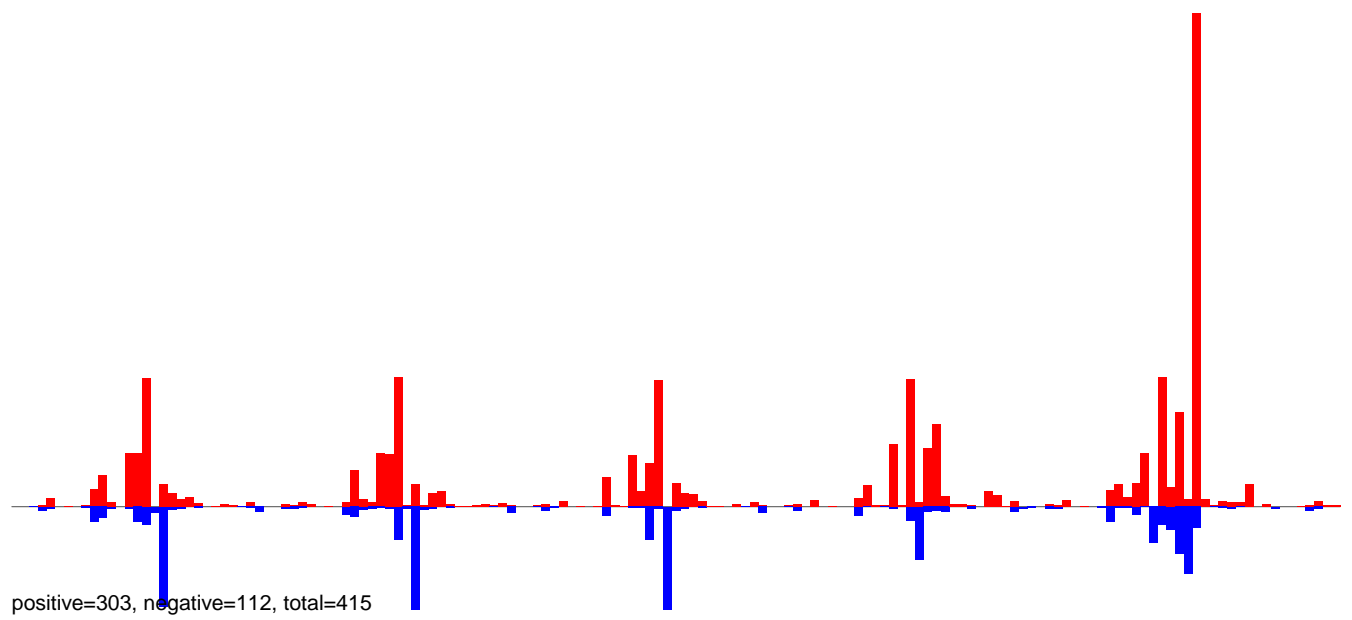
Window size=25, length=5616, TE@rnd-4_family-2843@LTR_Pao@2939:1-5616

0 1000 2000 3000 4000 5000

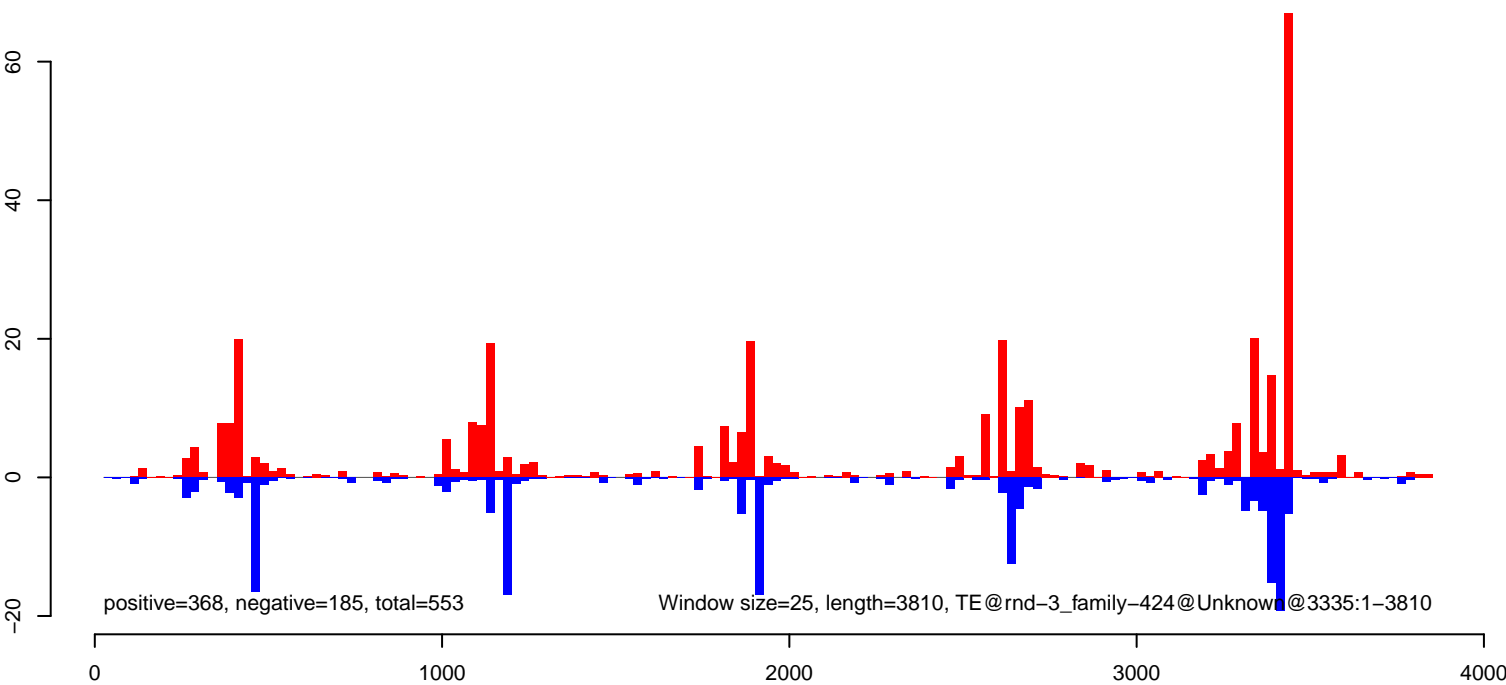
CuQuin_Testes_NL.18_23.rep



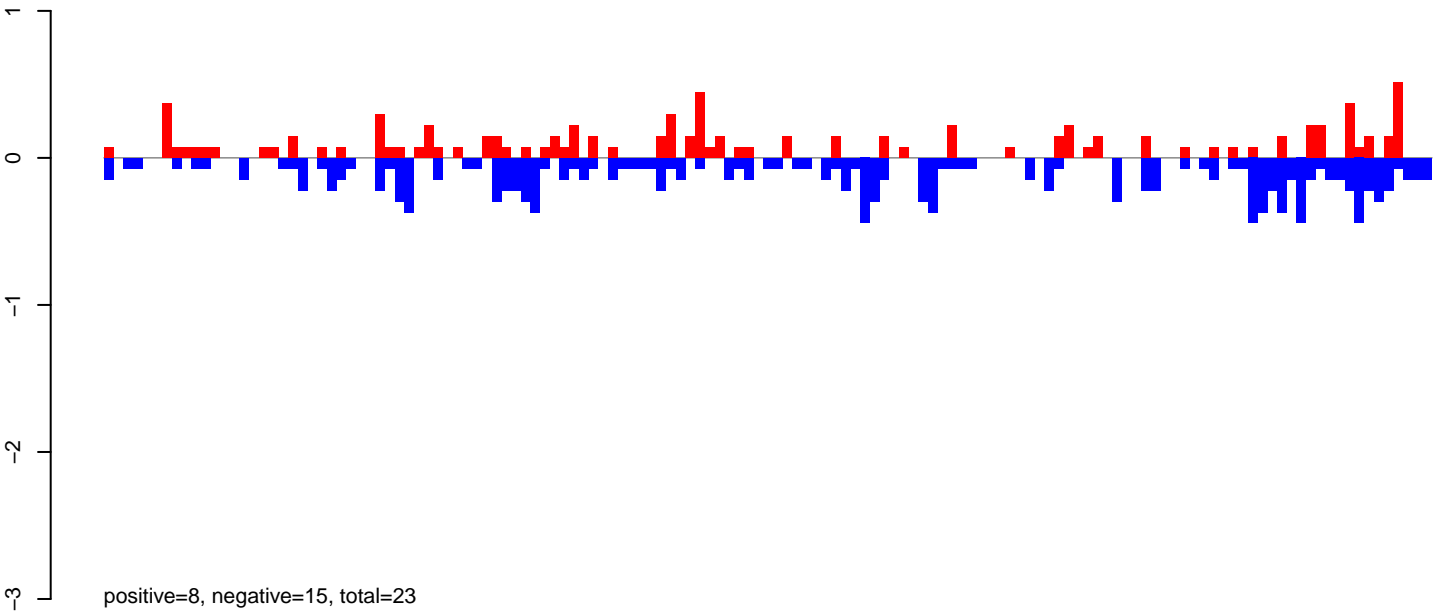
CuQuin_Testes_NL.24_35.rep



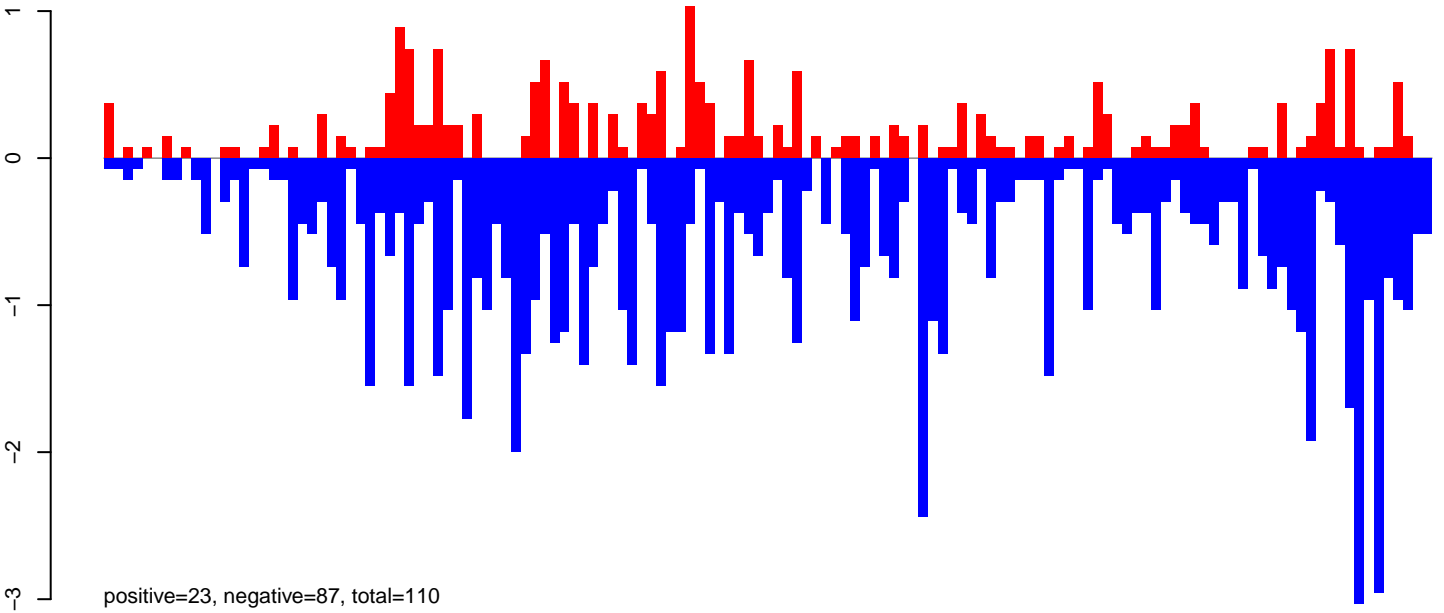
CuQuin_Testes_NL.rep



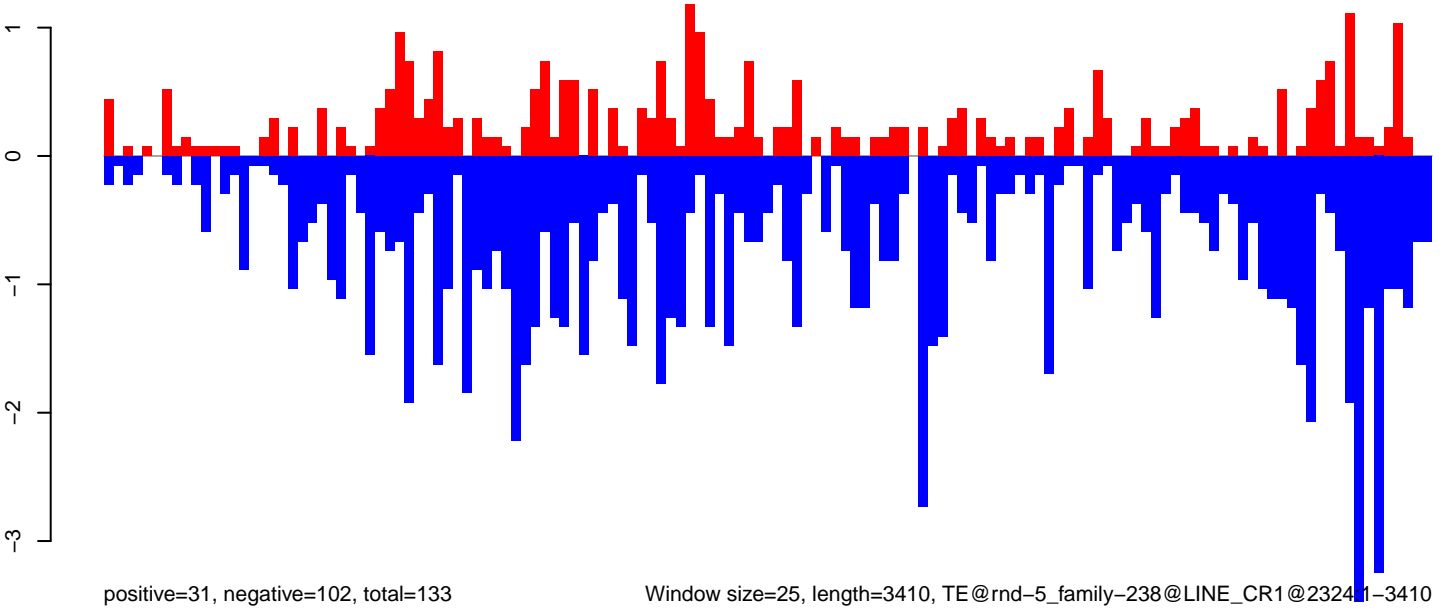
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



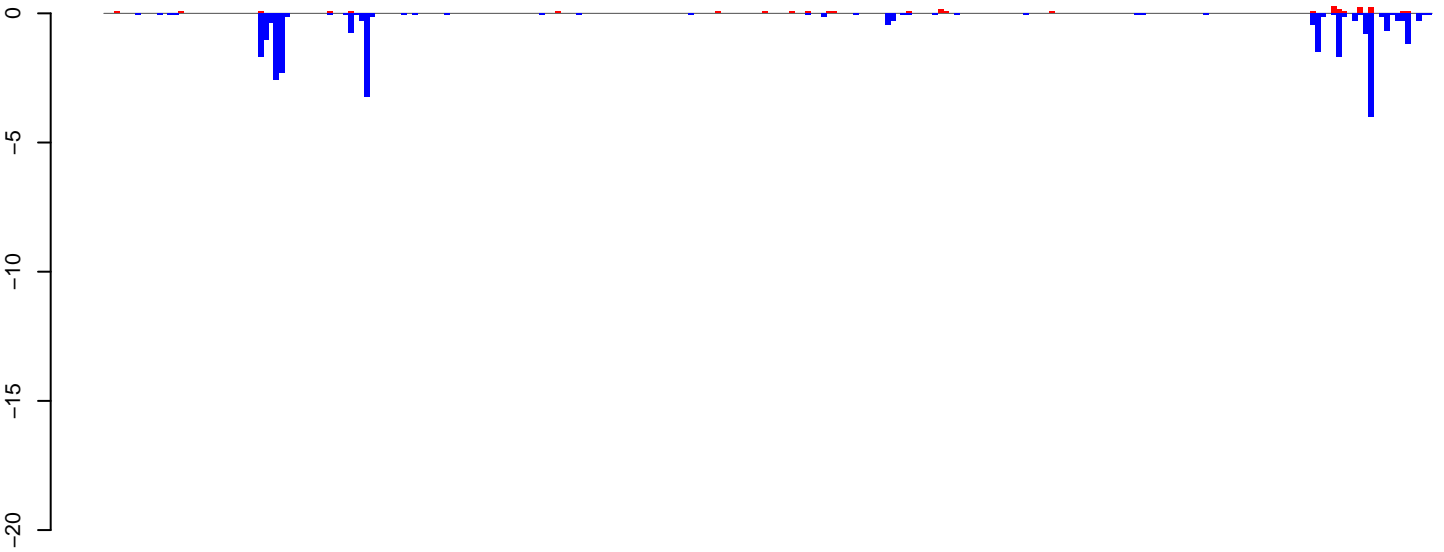
CuQuin_Testes_NL.rep



Window size=25, length=3410, TE@rnd-5_family-238@LINE_CR1@23241-3410

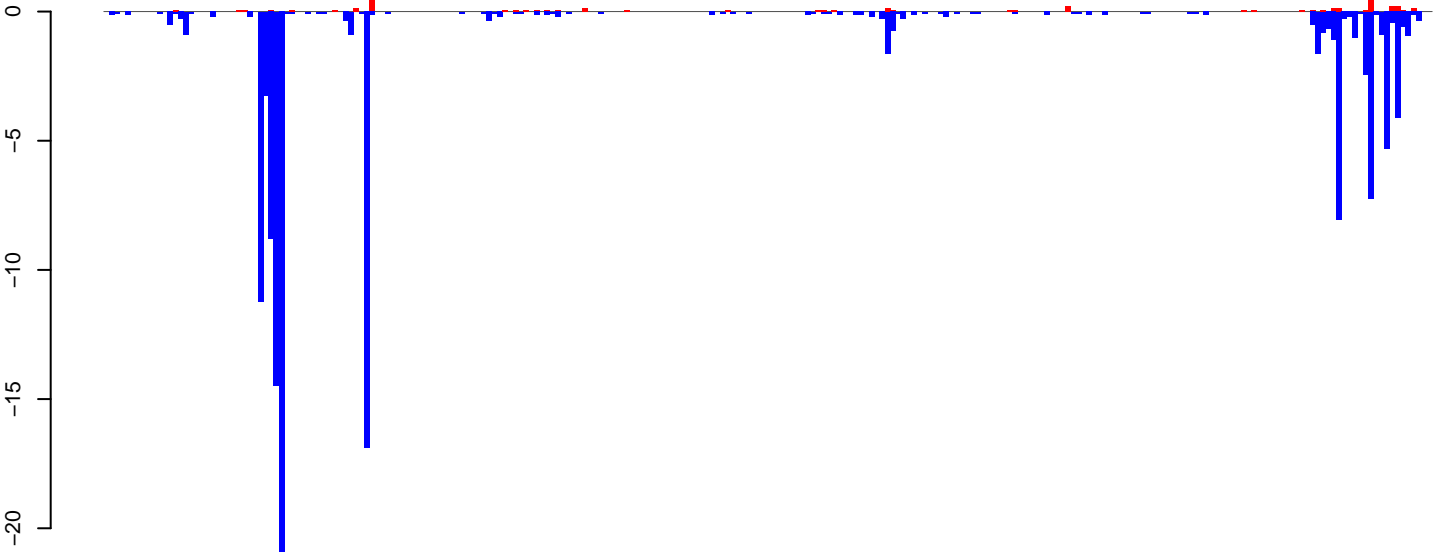
0 500 1000 1500 2000 2500 3000 3500

CuQuin_Testes_NL.18_23.rep



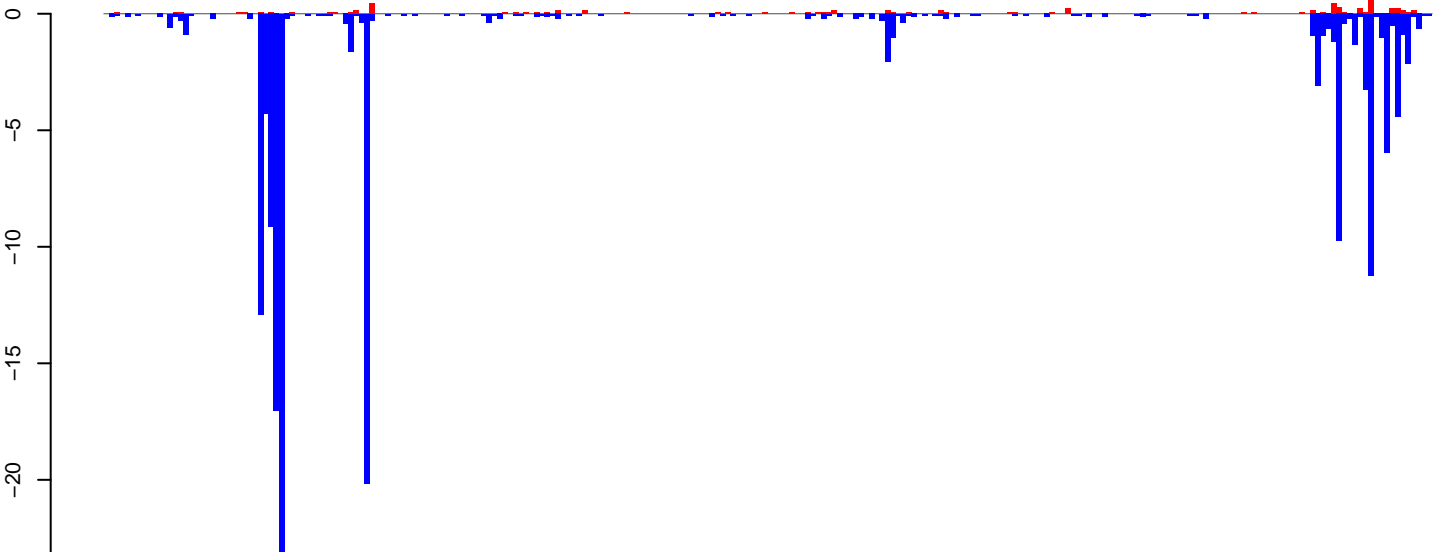
positive=2, negative=27, total=30

CuQuin_Testes_NL.24_35.rep



positive=5, negative=126, total=131

CuQuin_Testes_NL.rep

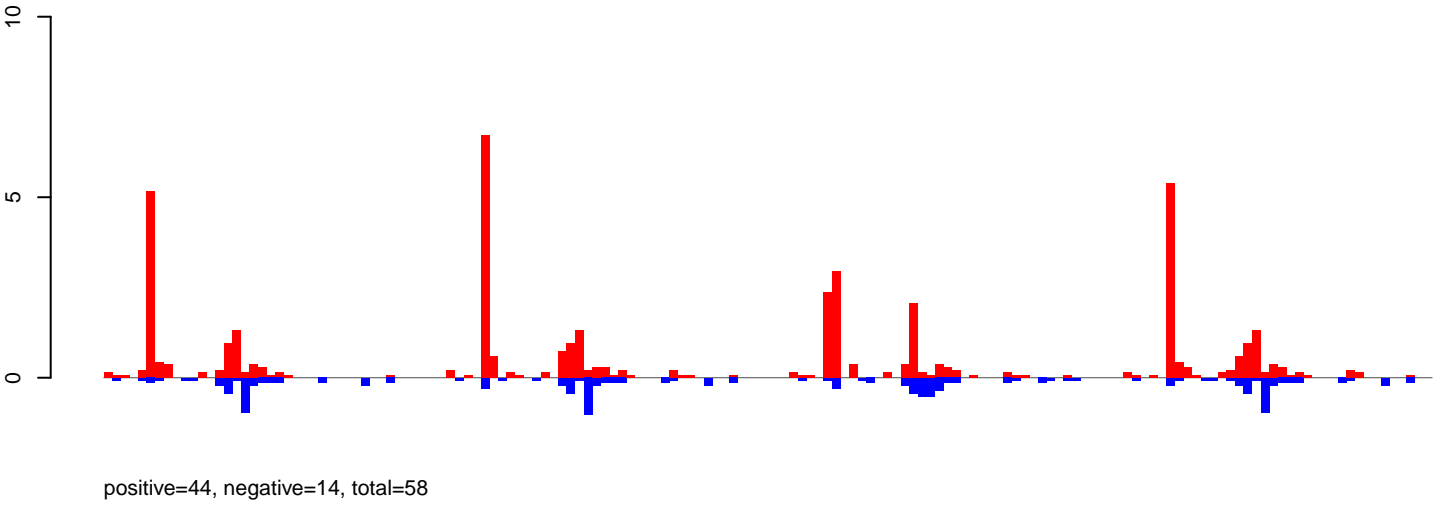


positive=8, negative=154, total=161

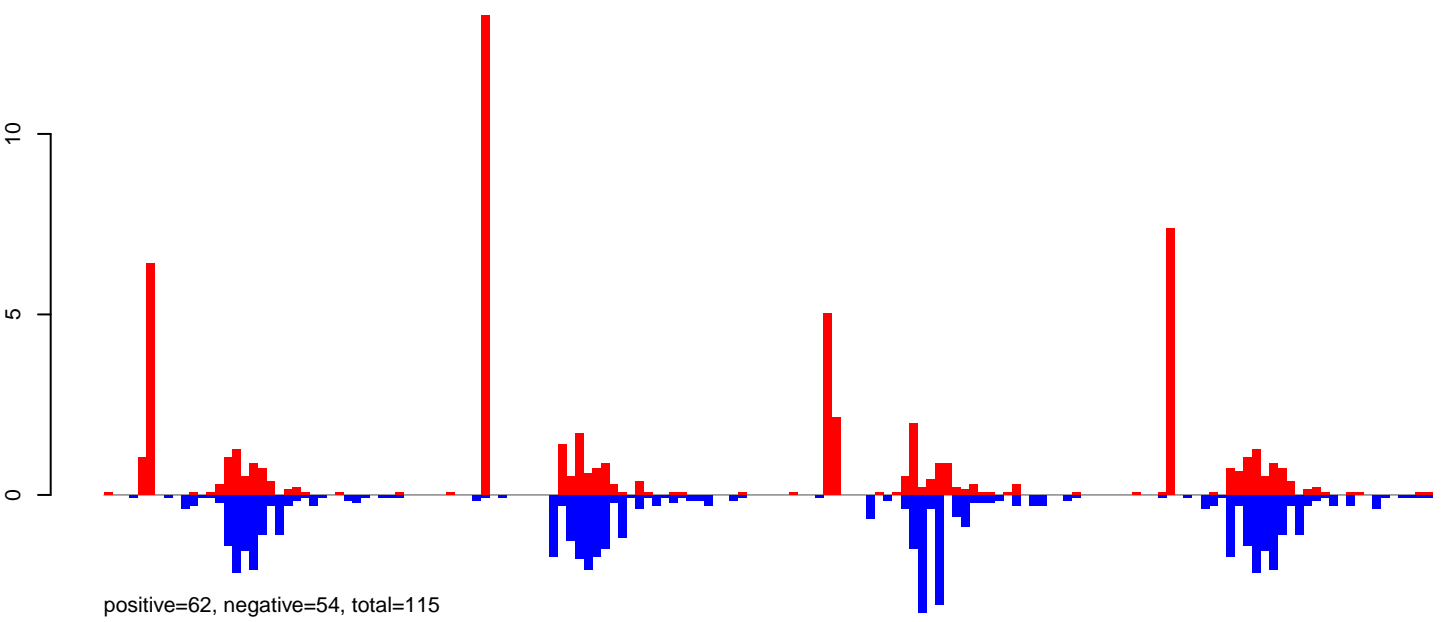
Window size=25, length=6240, TE@rnd-4_family-49@LTR_Gypsy@3453:1-6240

0 1000 2000 3000 4000 5000 6000

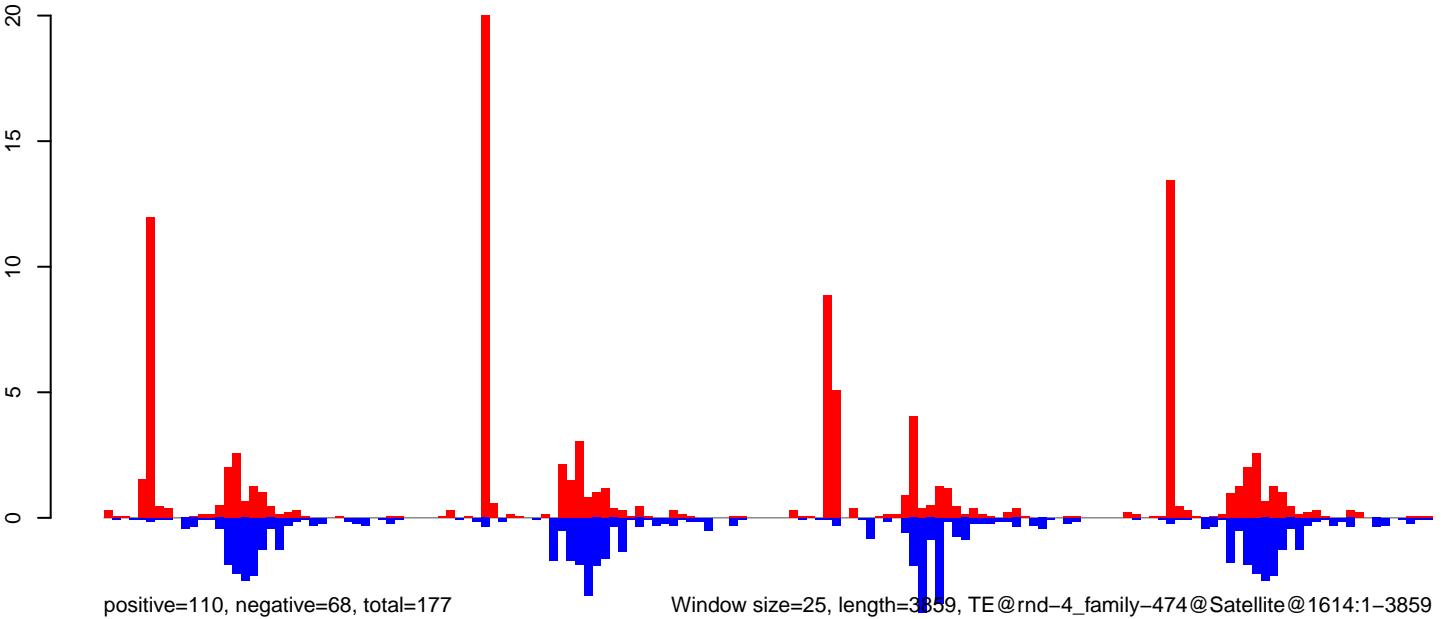
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



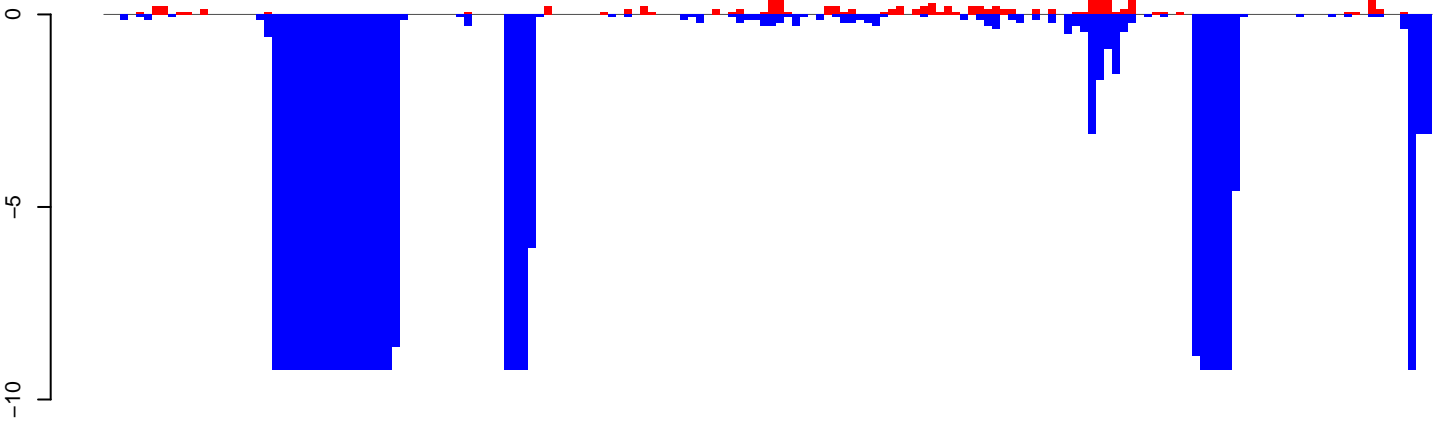
CuQuin_Testes_NL.rep



Window size=25, length=3859, TE@rnd-4_family-474@Satellite@1614:1-3859

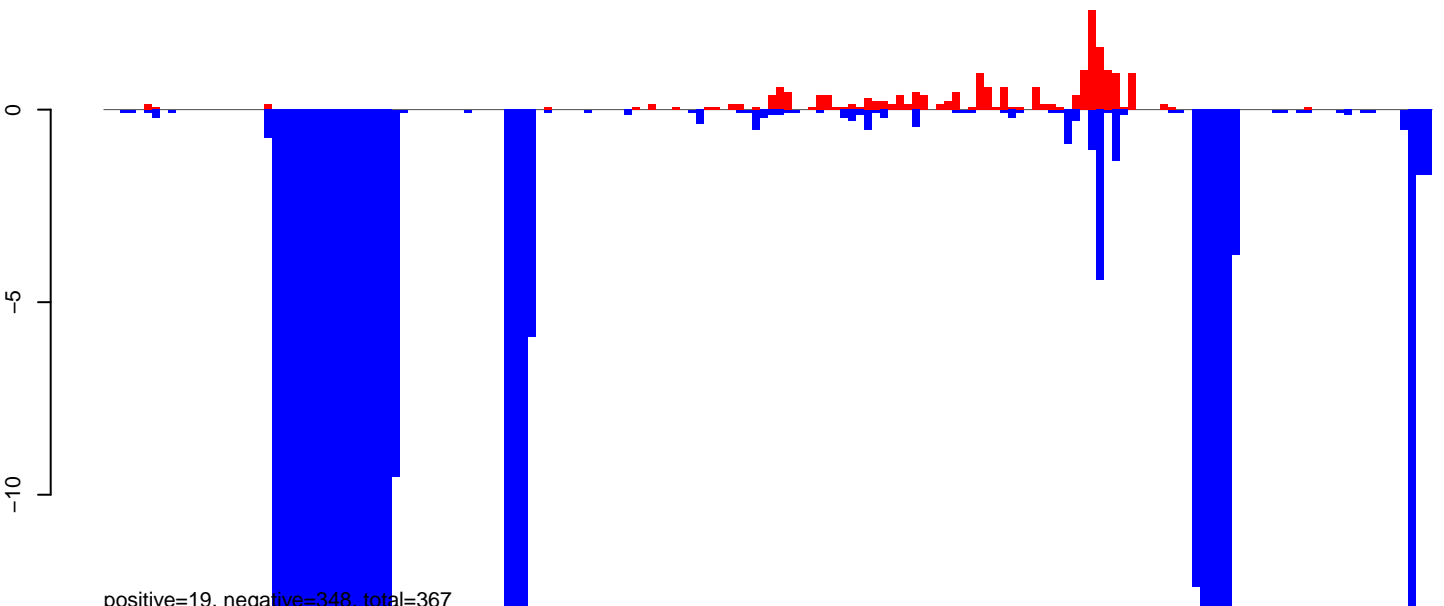
0 1000 2000 3000 4000

CuQuin_Testes_NL.18_23.rep



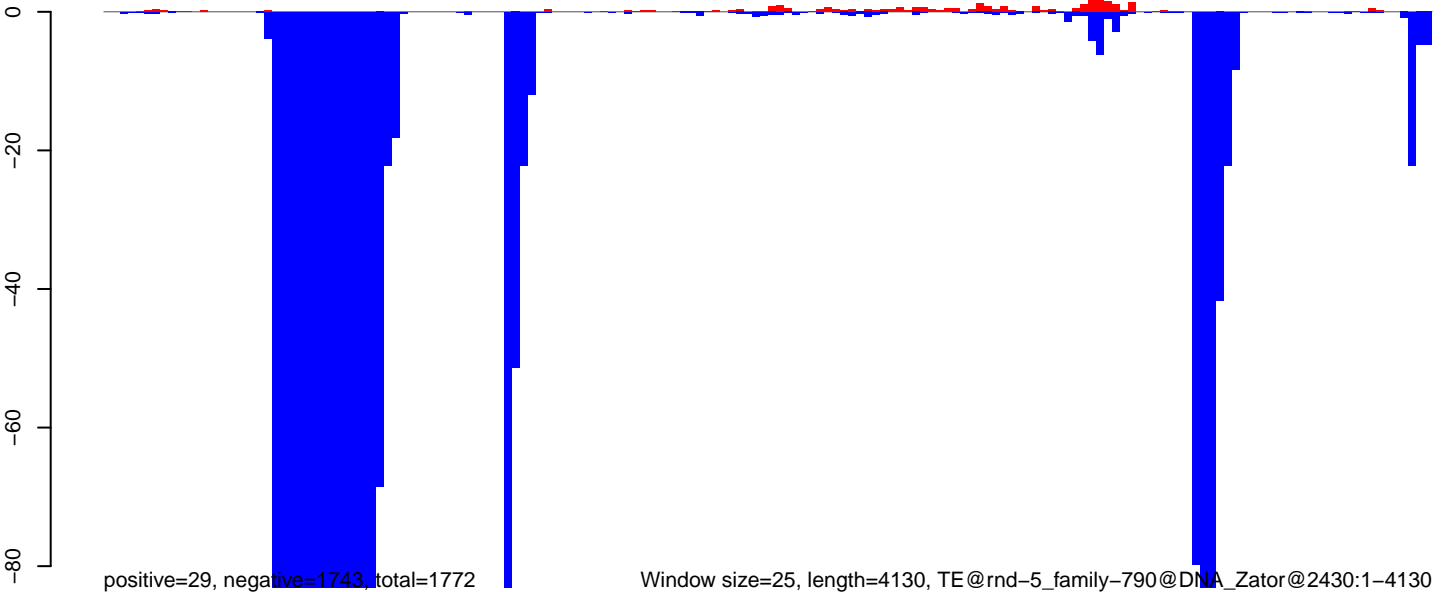
positive=10, negative=264, total=274

CuQuin_Testes_NL.24_35.rep



positive=19, negative=348, total=367

CuQuin_Testes_NL.rep

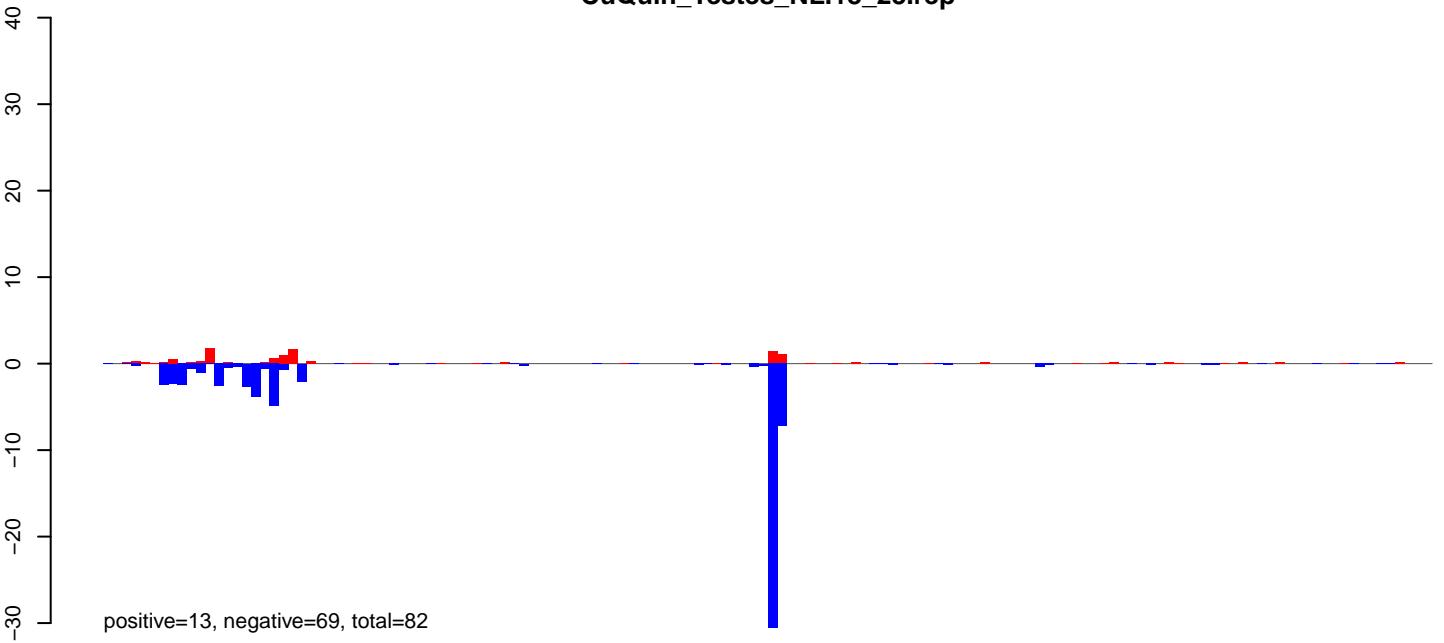


positive=29, negative=1743, total=1772

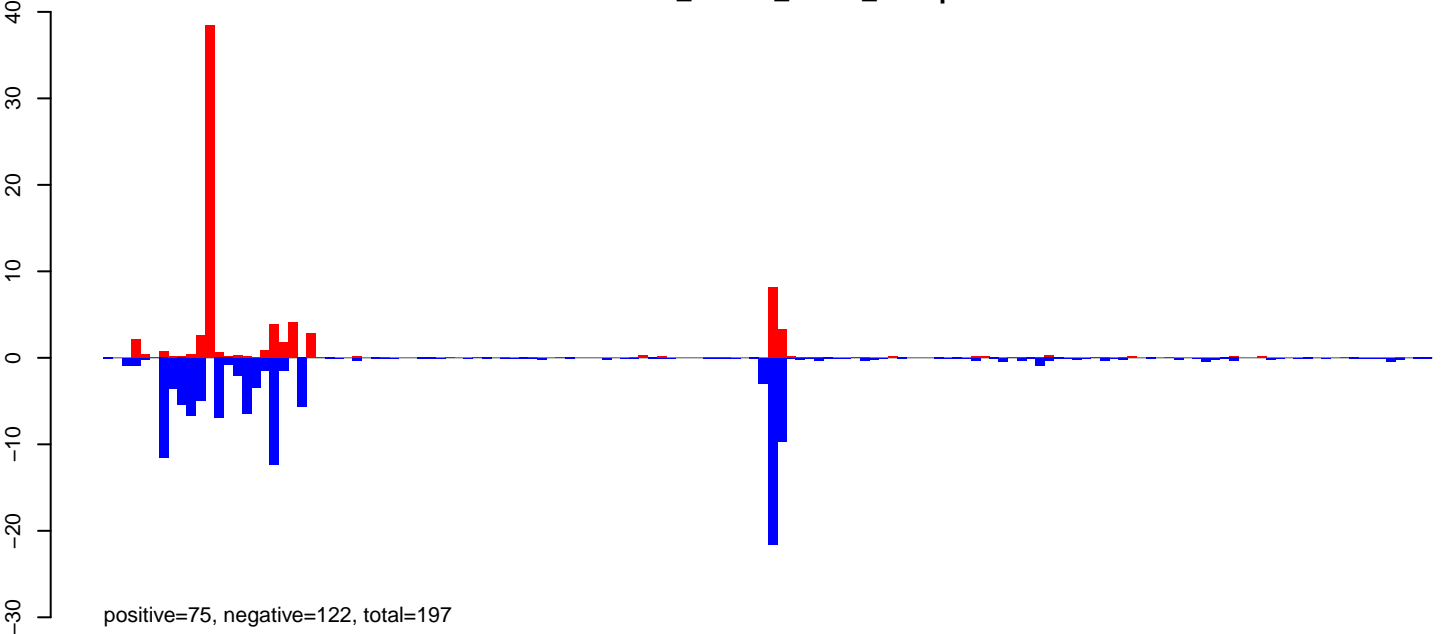
Window size=25, length=4130, TE@rnd-5_family-790@DNA_Zator@2430:1-4130

0 1000 2000 3000 4000

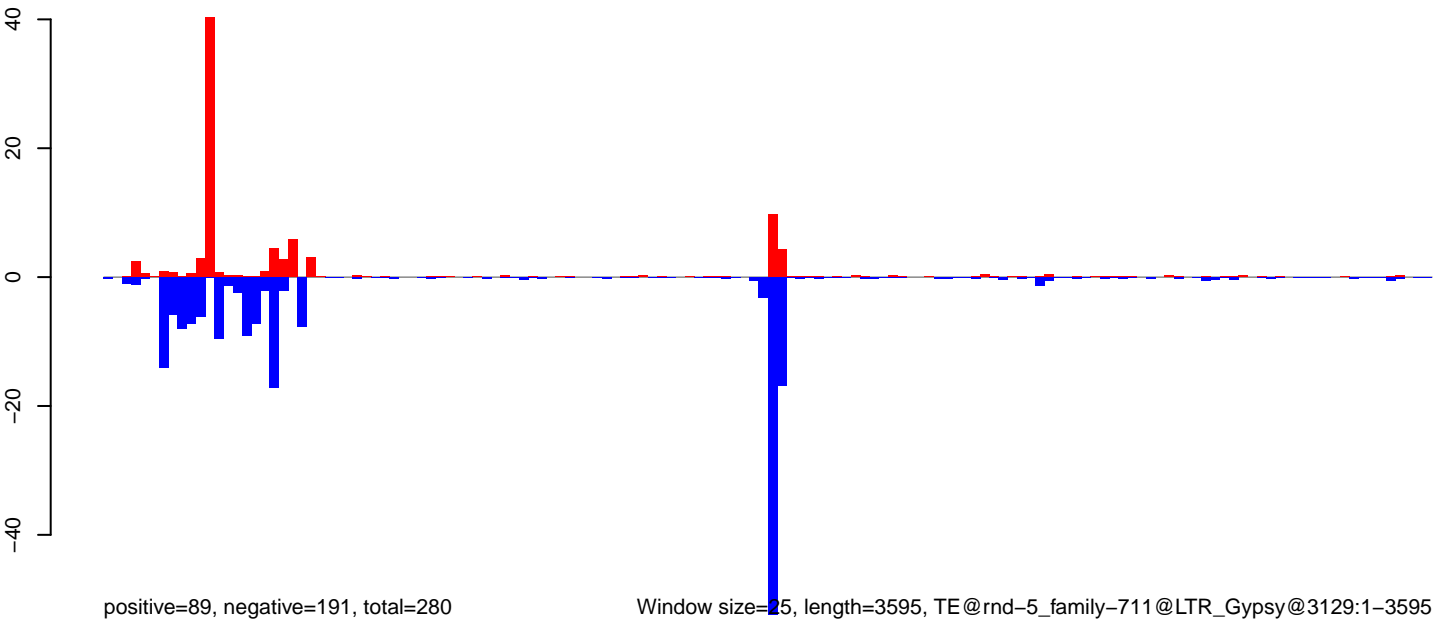
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

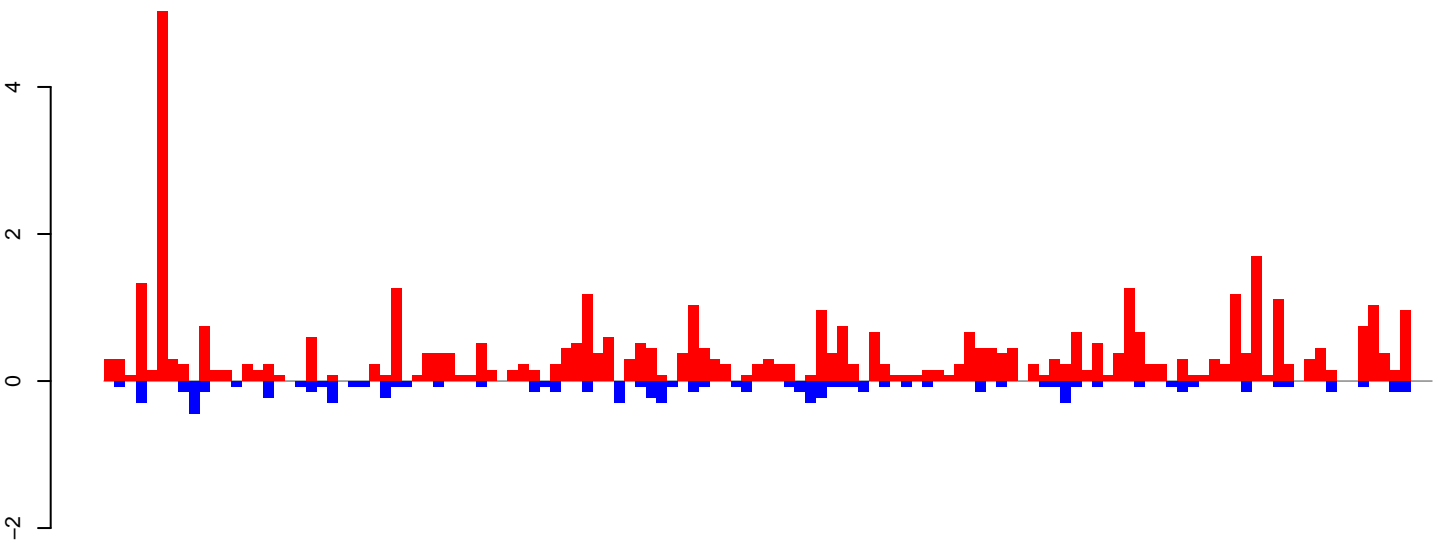


CuQuin_Testes_NL.rep



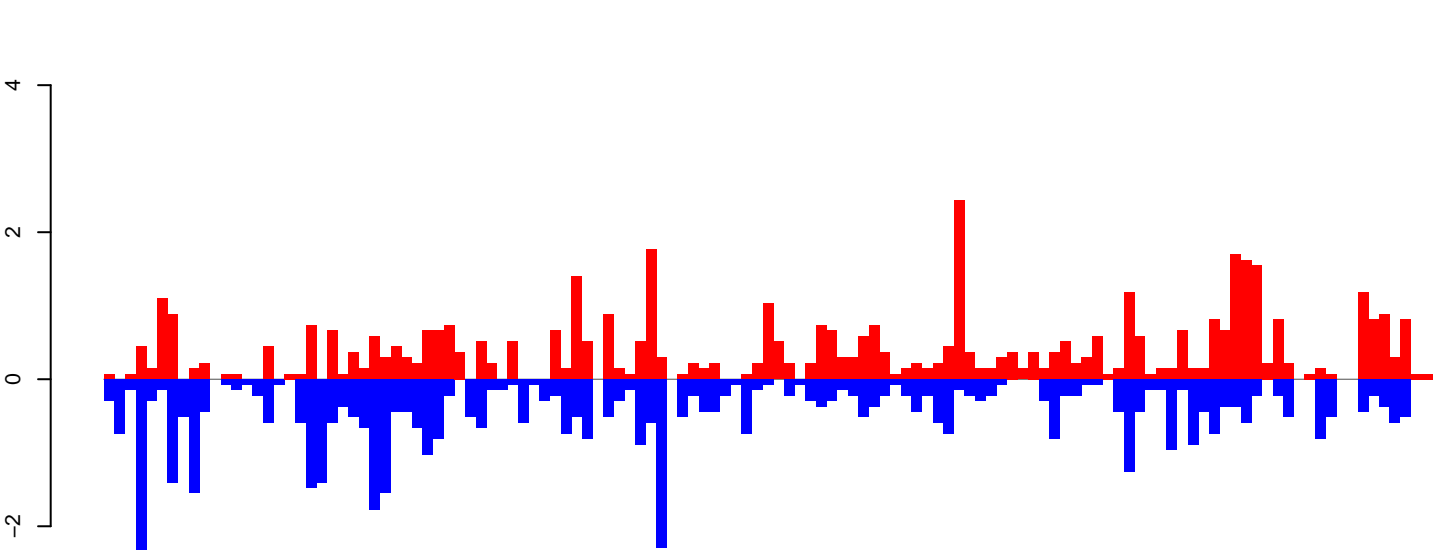
Window size=25, length=3595, TE@rnd-5_family-711@LTR_Gypsy@3129:1-3595

CuQuin_Testes_NL.18_23.rep



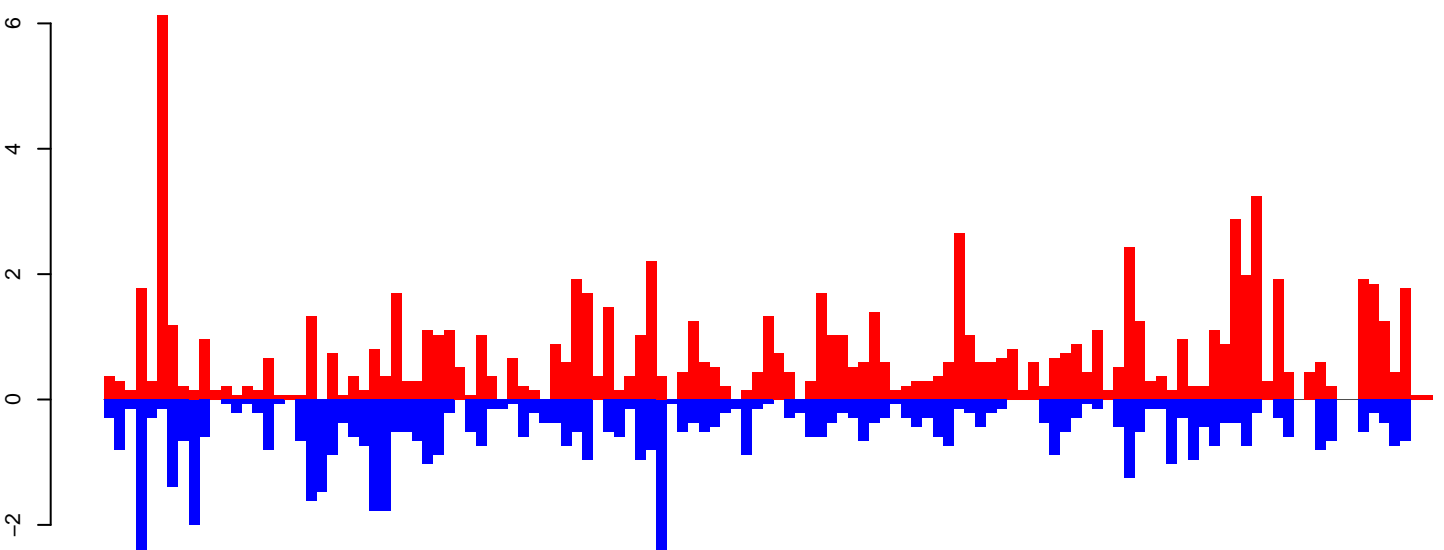
positive=43, negative=8, total=51

CuQuin_Testes_NL.24_35.rep



positive=47, negative=54, total=101

CuQuin_Testes_NL.rep

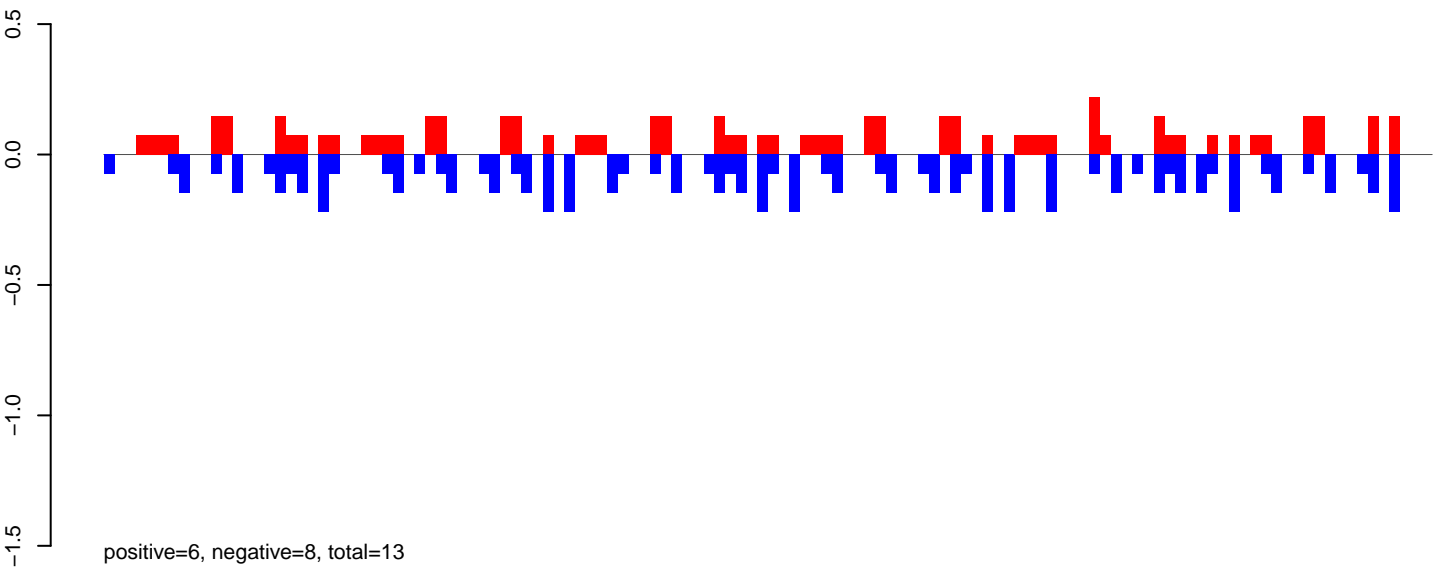


positive=91, negative=62, total=153

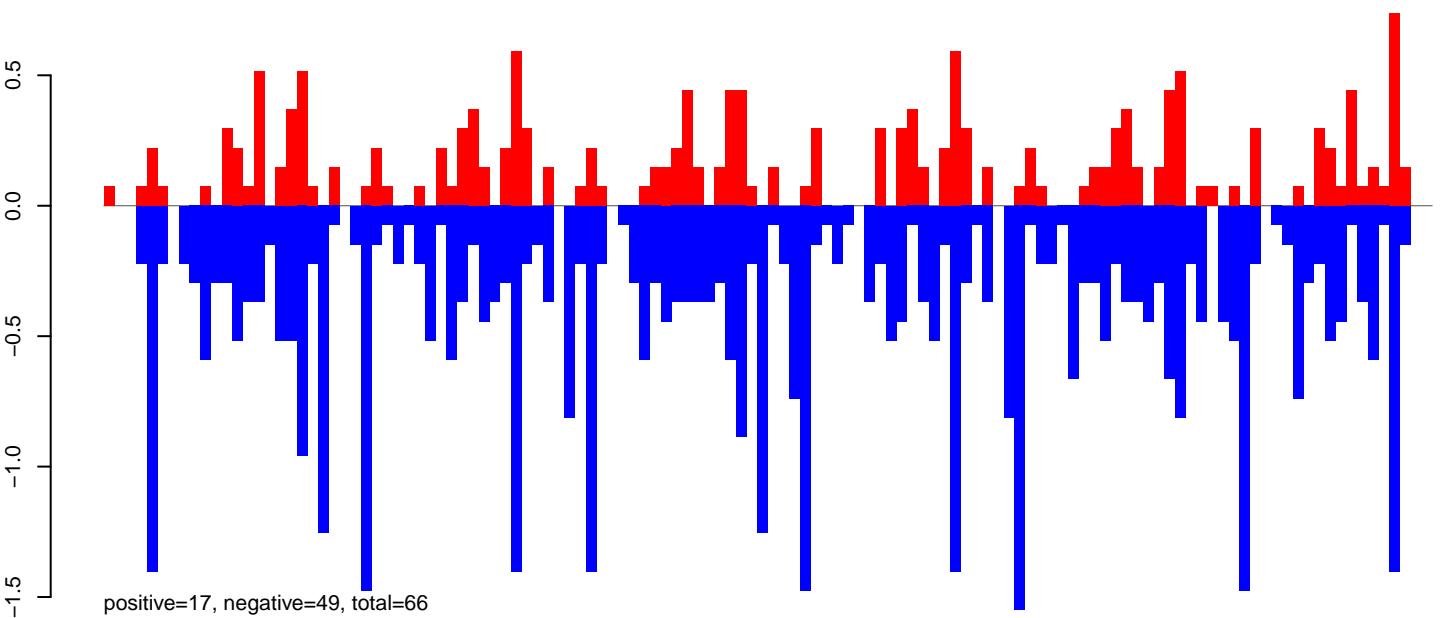
Window size=25, length=3109, TE@rnd-2_family-1@LINE_RTE-BovB@1921:1-3109

0 500 1000 1500 2000 2500 3000

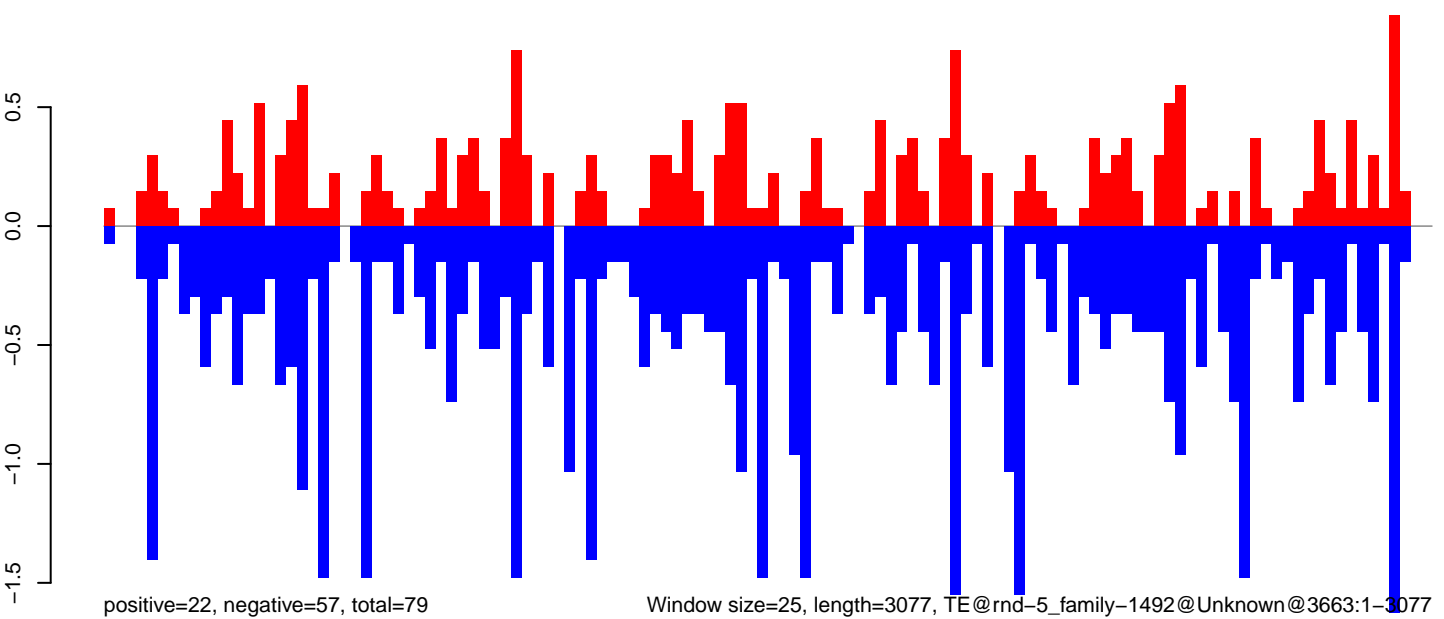
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



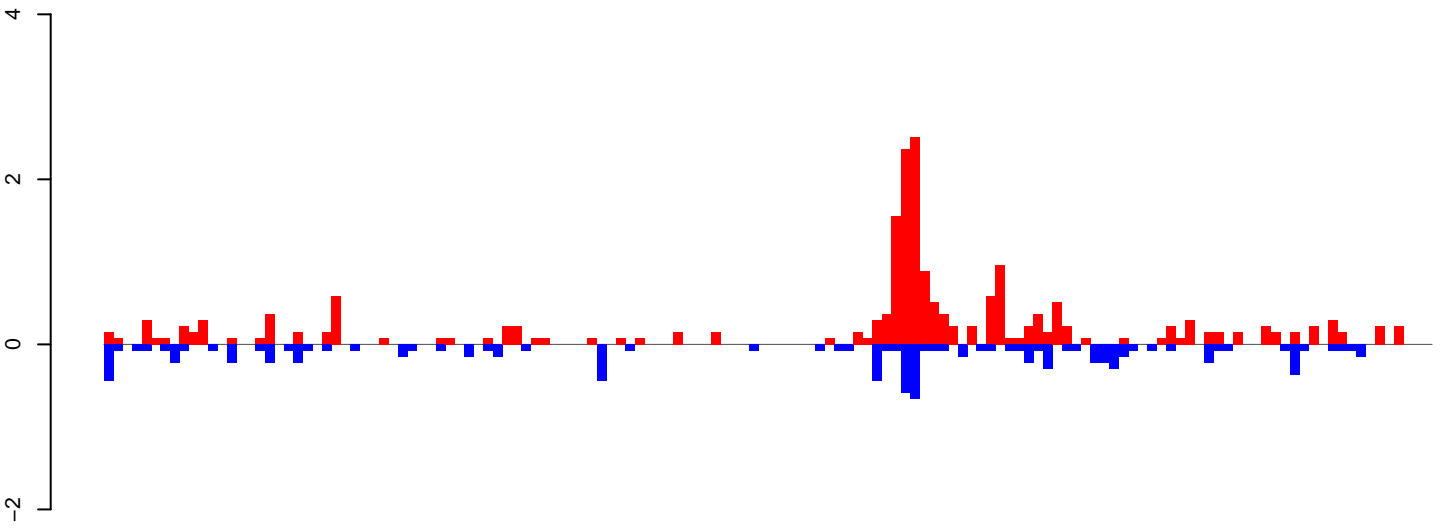
CuQuin_Testes_NL.rep



Window size=25, length=3077, TE@rnd-5_family-1492@Unknown@3663:1-3077

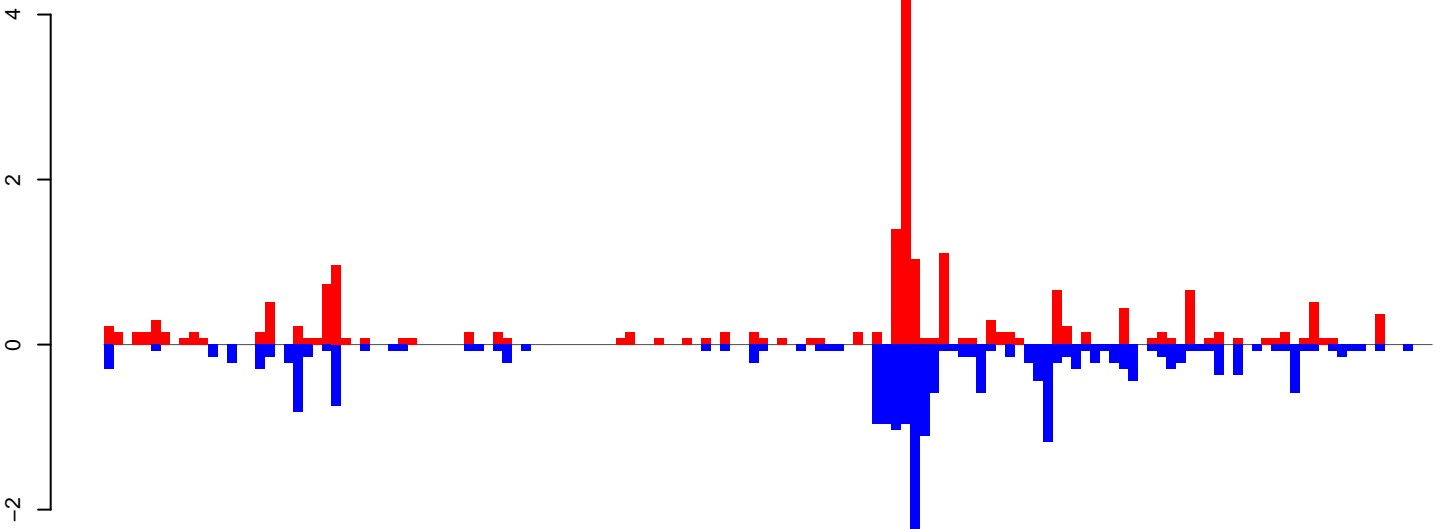
0 500 1000 1500 2000 2500 3000

CuQuin_Testes_NL.18_23.rep



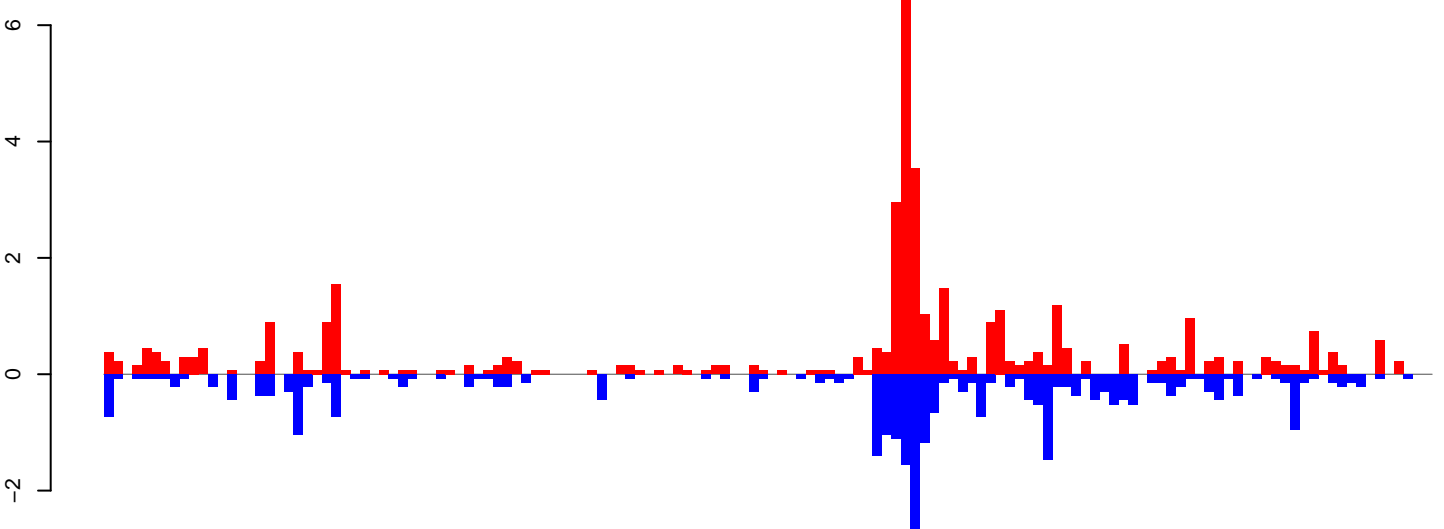
positive=20, negative=9, total=29

CuQuin_Testes_NL.24_35.rep



positive=19, negative=21, total=41

CuQuin_Testes_NL.rep

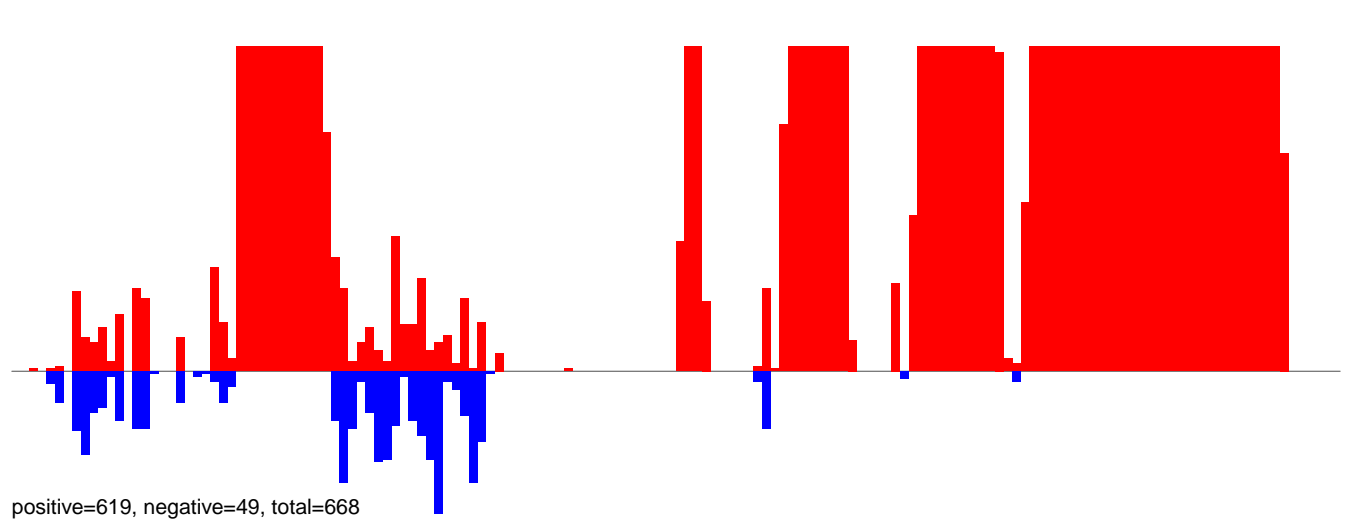


positive=39, negative=31, total=70

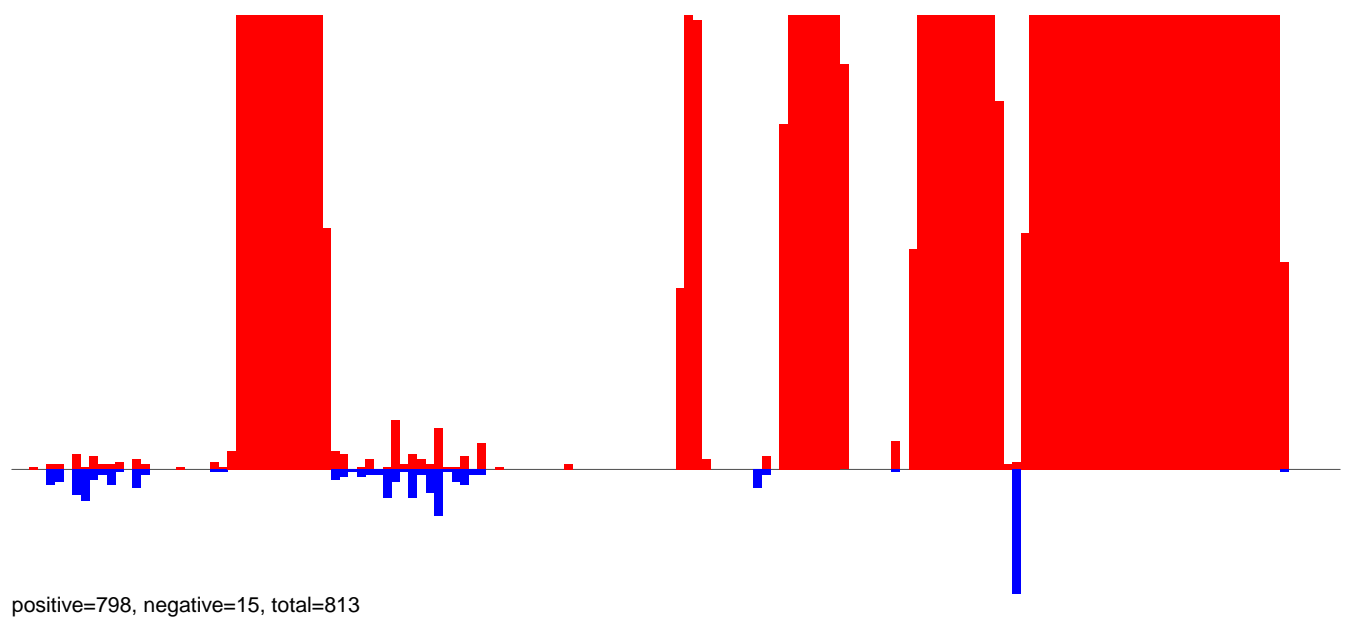
Window size=25, length=3495, TE@rnd-3_family-471@Unknown@3294:1-3495

0 500 1000 1500 2000 2500 3000 3500

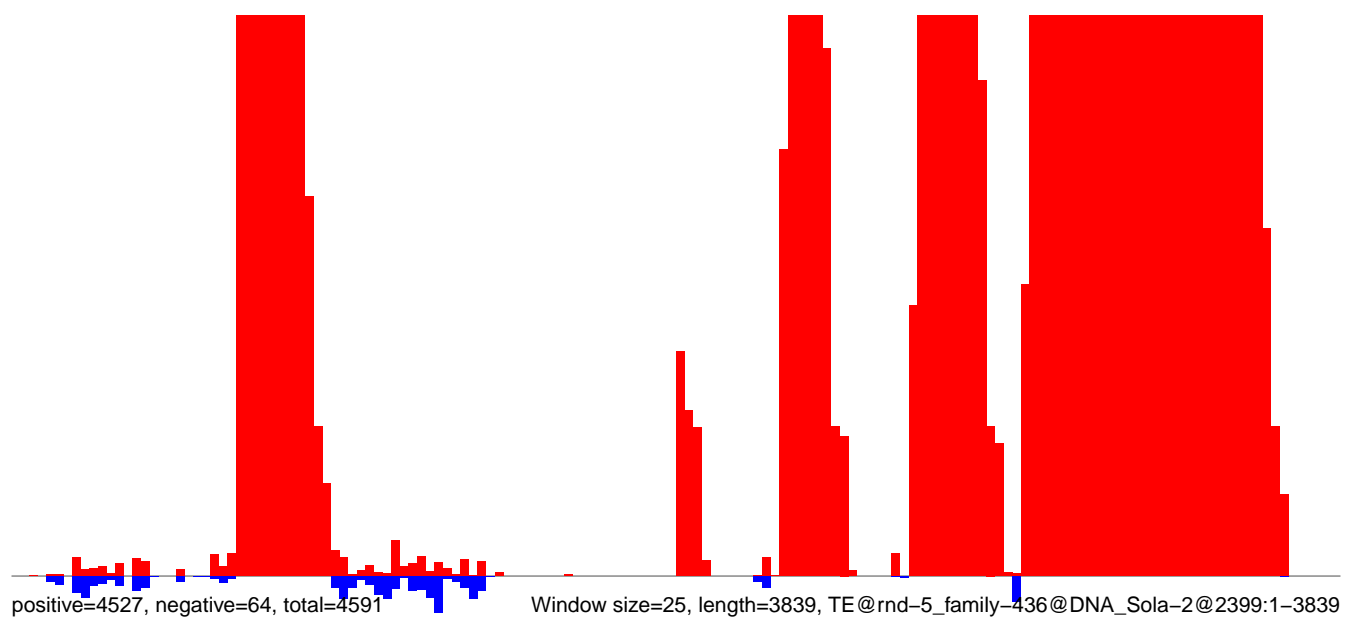
CuQuin_Testes_NL.18_23.rep



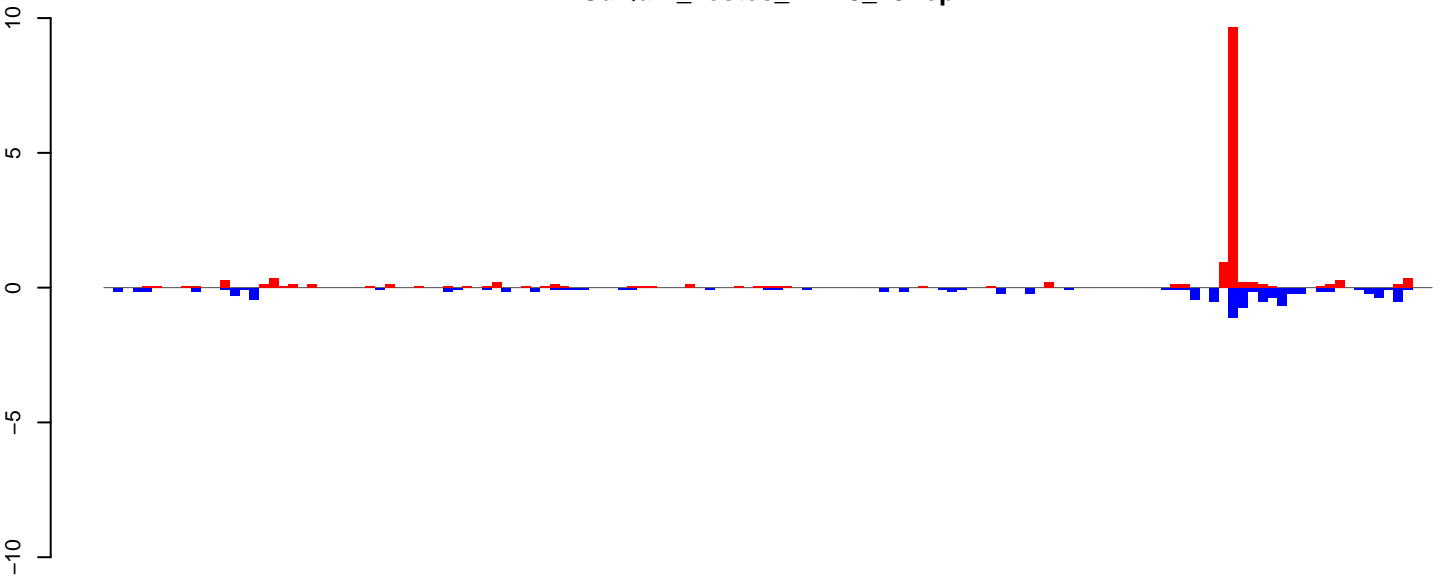
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

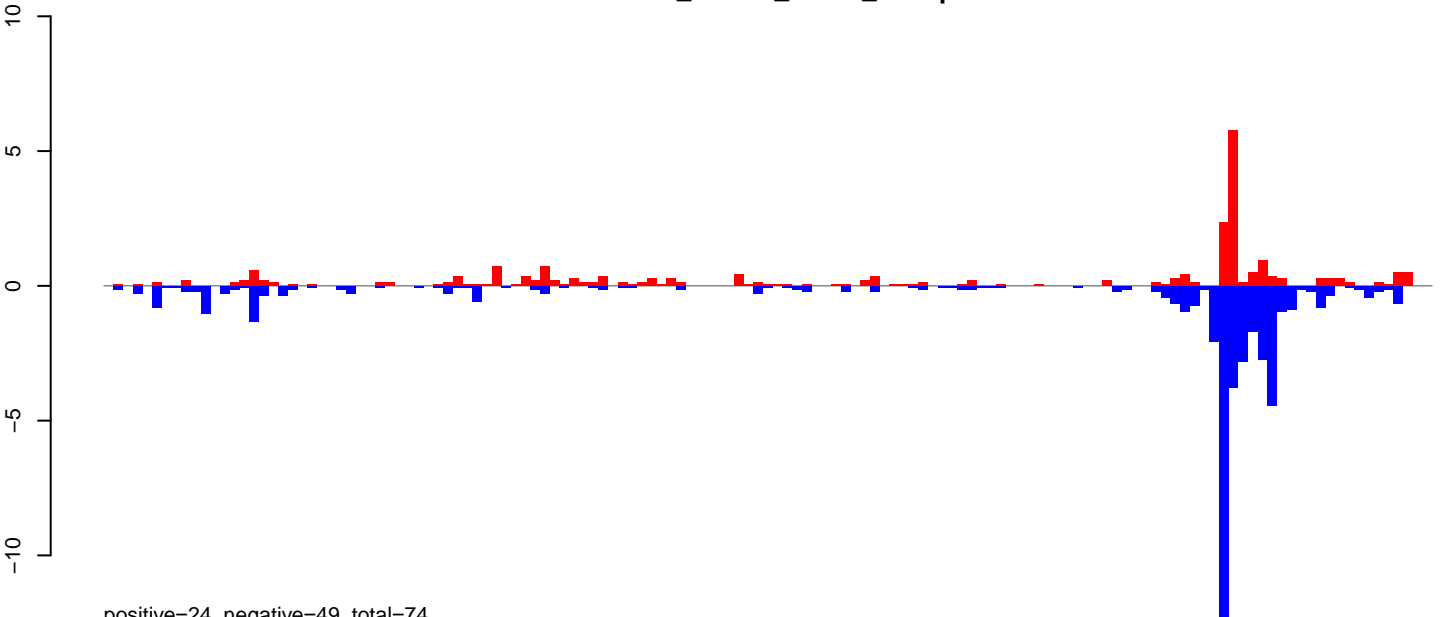


CuQuin_Testes_NL.18_23.rep



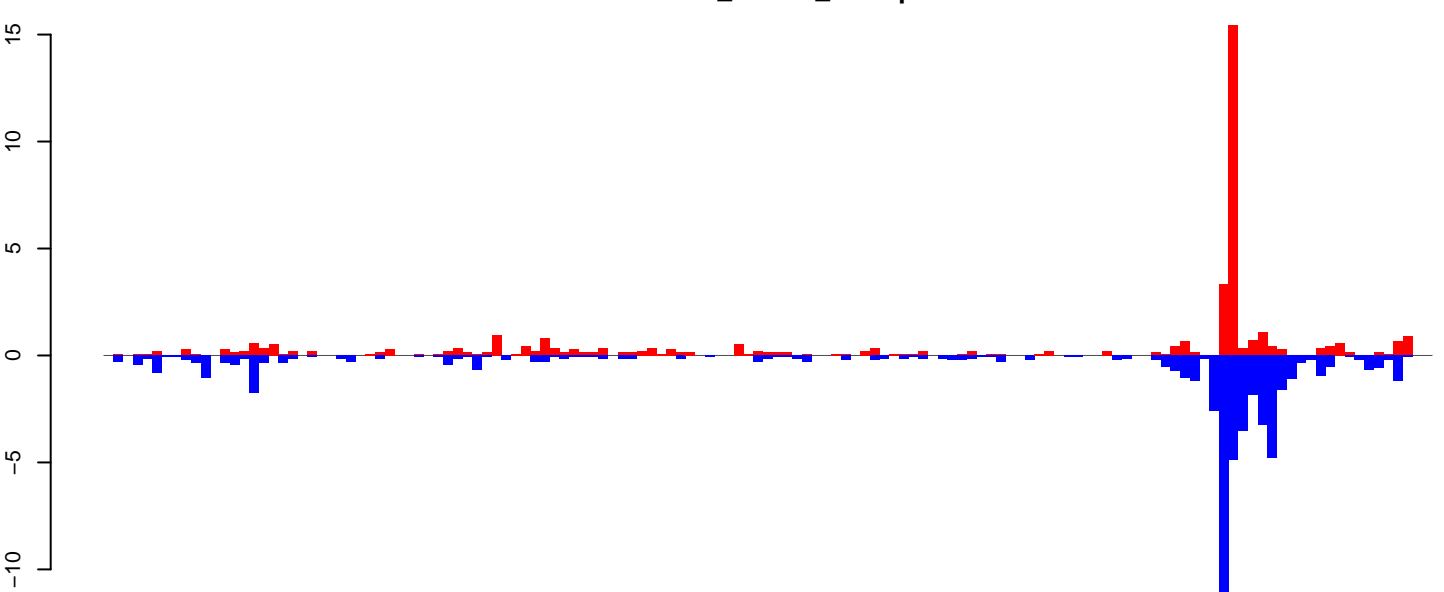
positive=16, negative=11, total=27

CuQuin_Testes_NL.24_35.rep



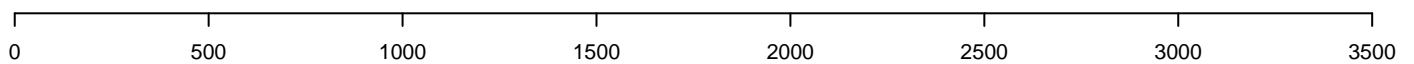
positive=24, negative=49, total=74

CuQuin_Testes_NL.rep

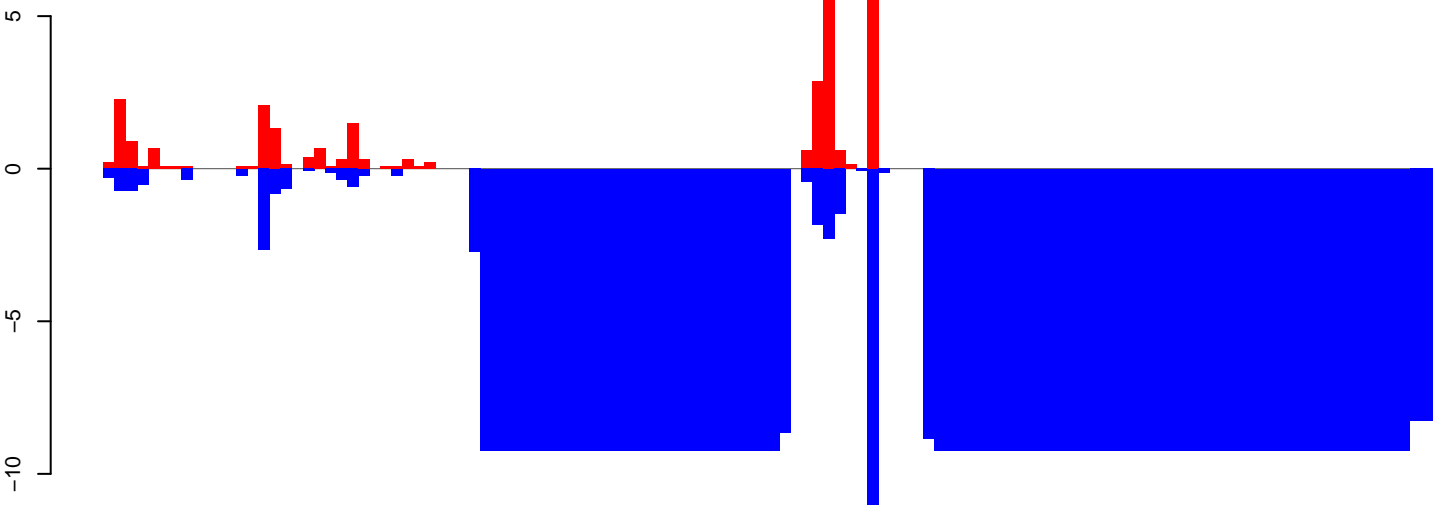


positive=41, negative=60, total=101

Window size=25, length=3414, TE@rnd-4_family-588@LINE_I@183:1-3414

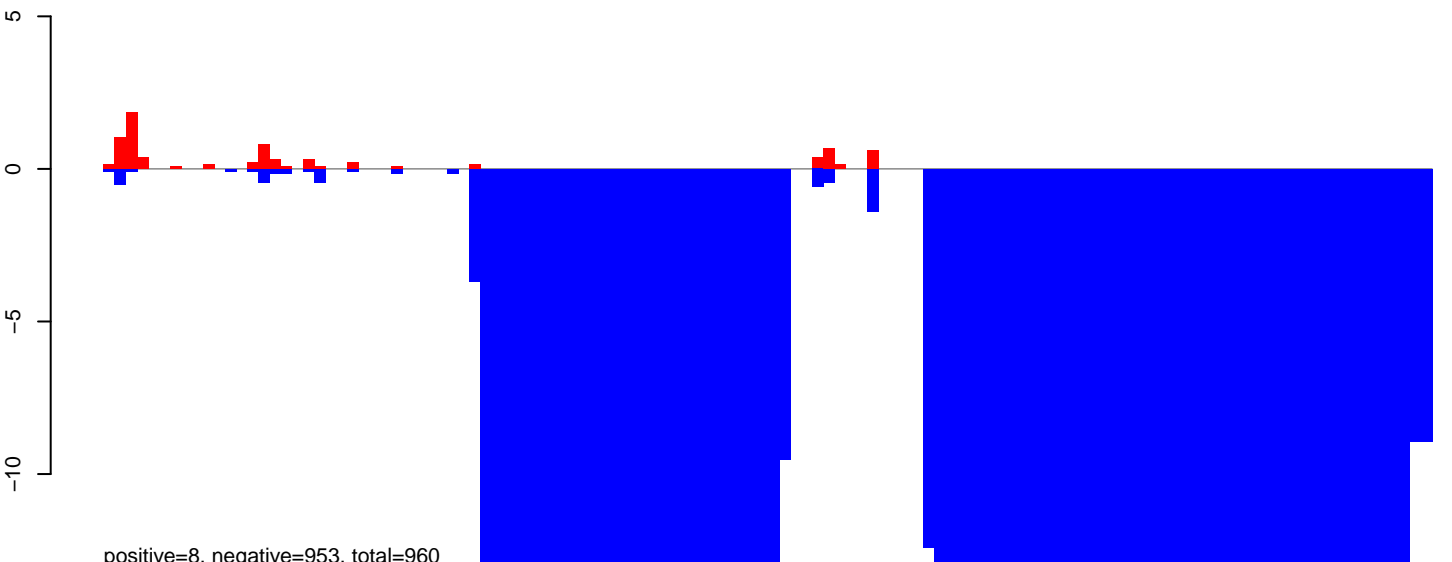


CuQuin_Testes_NL.18_23.rep



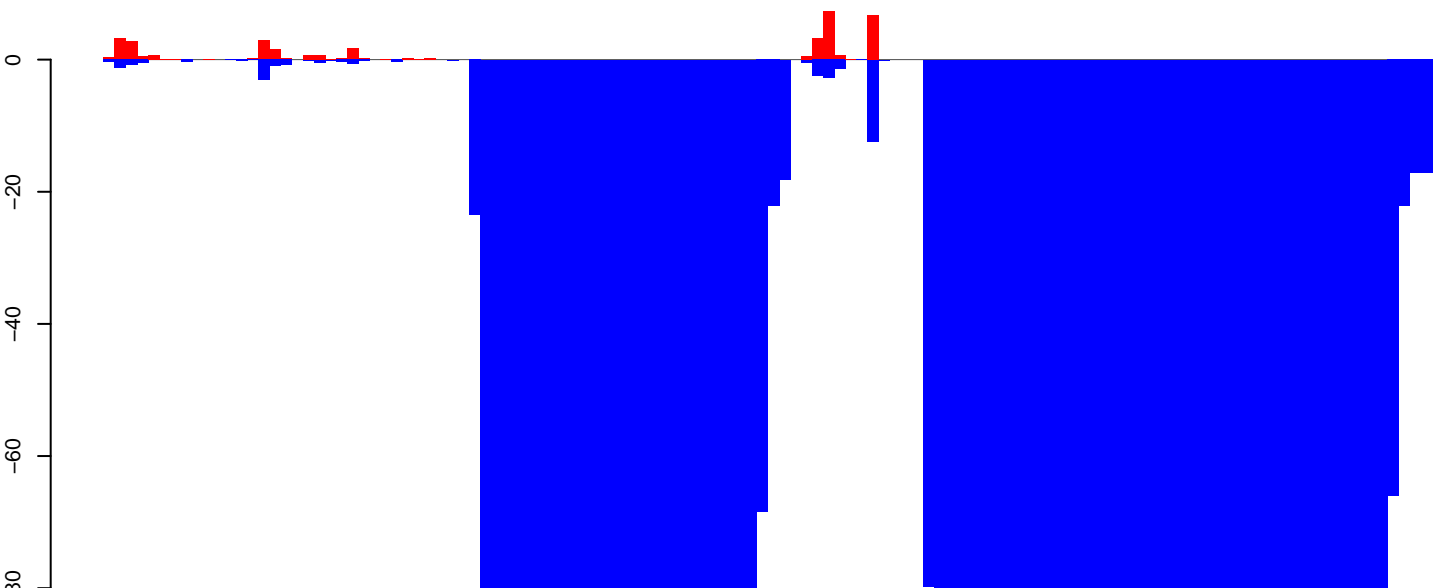
positive=29, negative=709, total=738

CuQuin_Testes_NL.24_35.rep



positive=8, negative=953, total=960

CuQuin_Testes_NL.rep

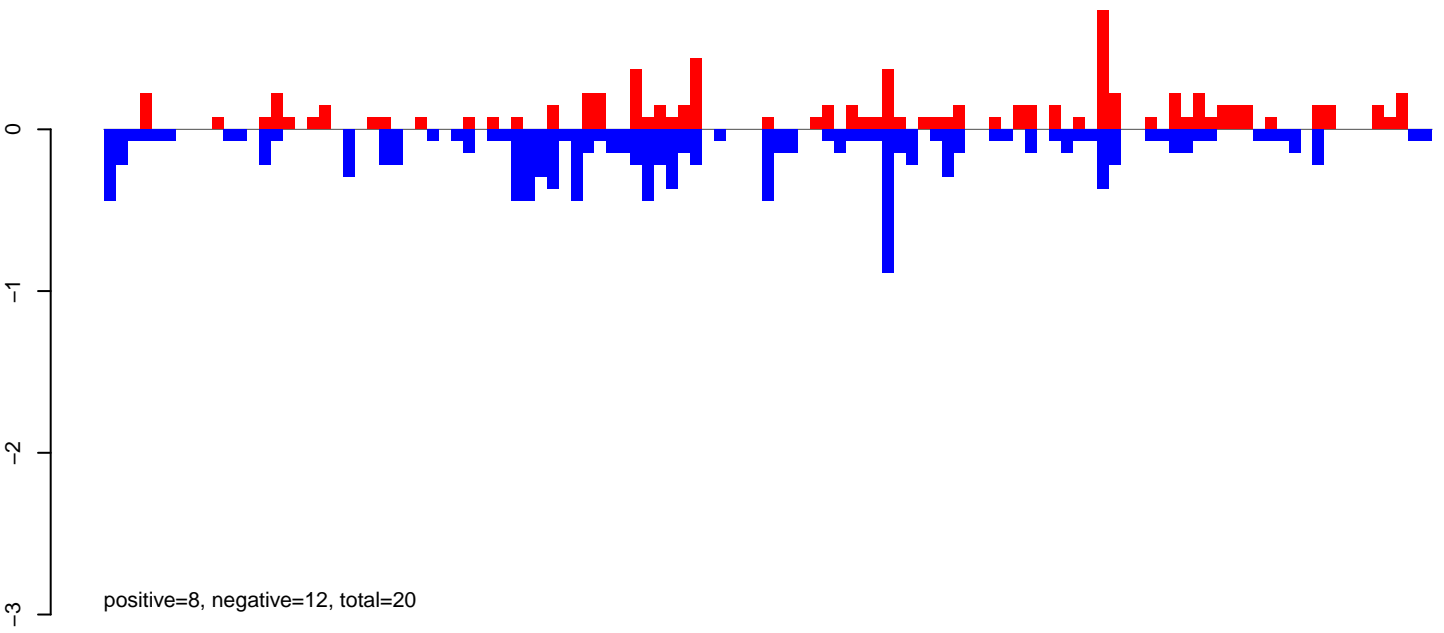


positive=37, negative=5848, total=5885

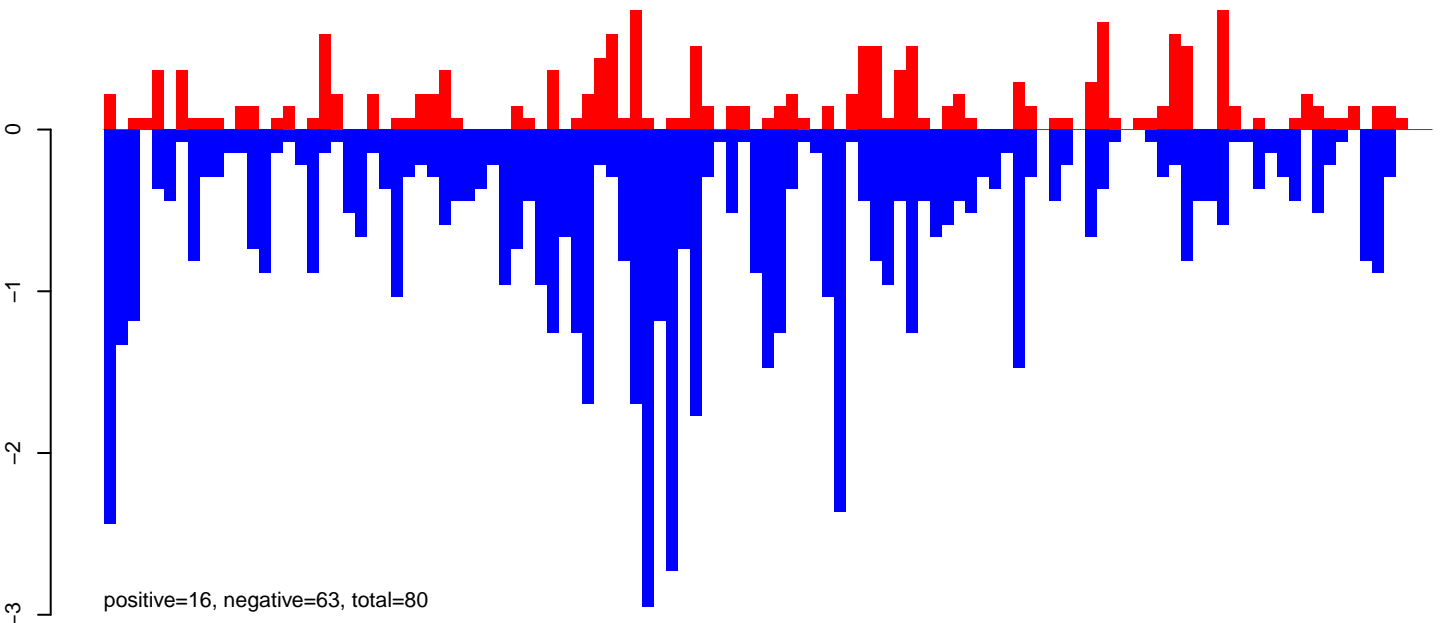
Window size=25, length=2994, TE@rnd-5_family-215@Unknown@3059:1-2994

0 500 1000 1500 2000 2500 3000

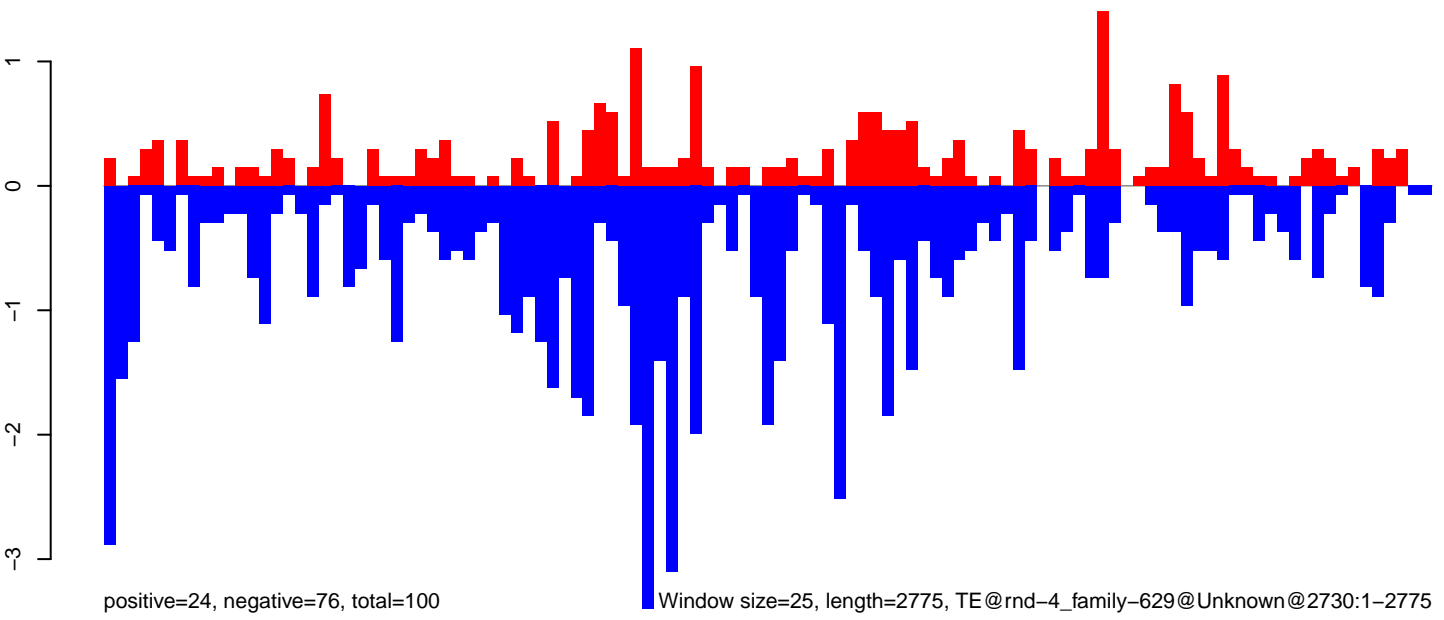
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

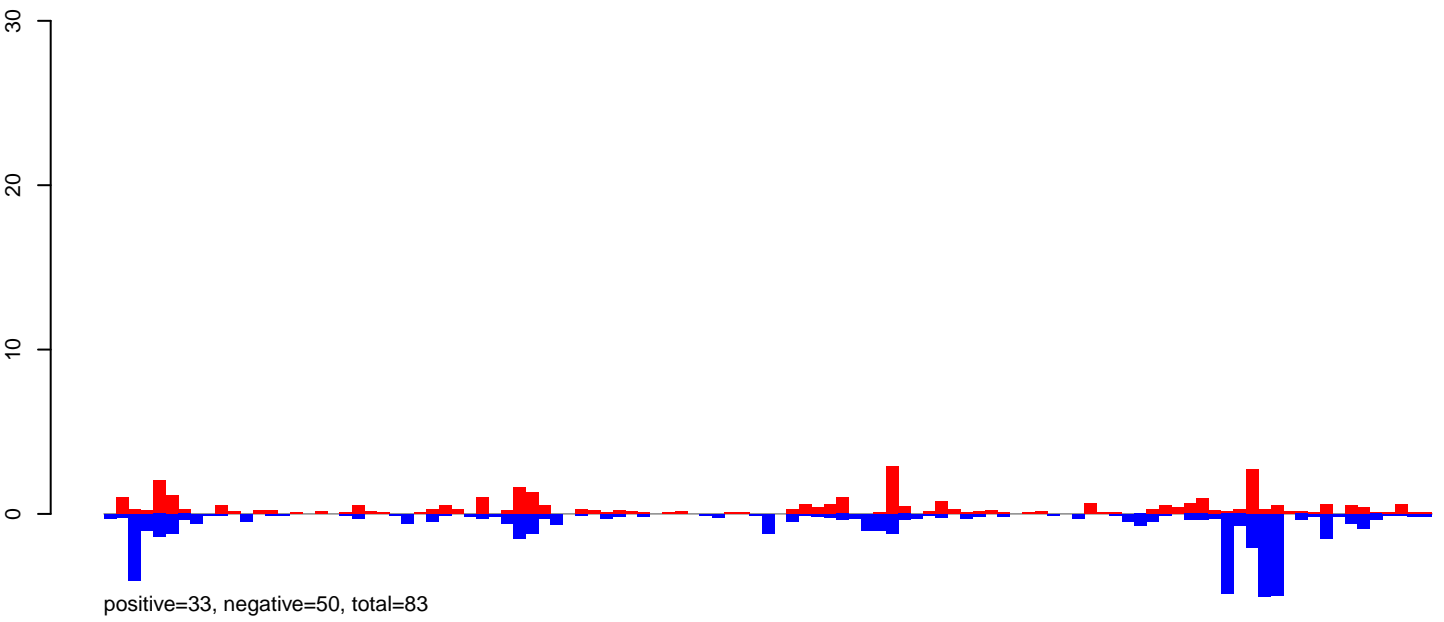


CuQuin_Testes_NL.rep

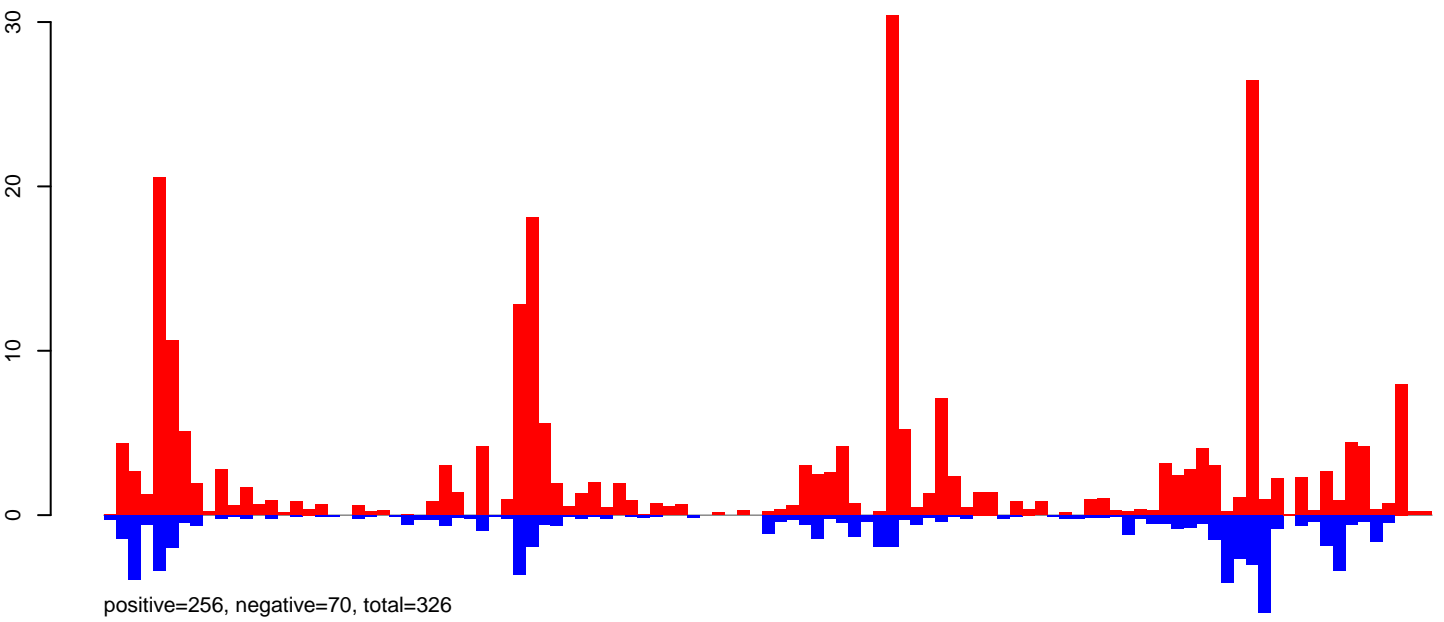


0 500 1000 1500 2000 2500

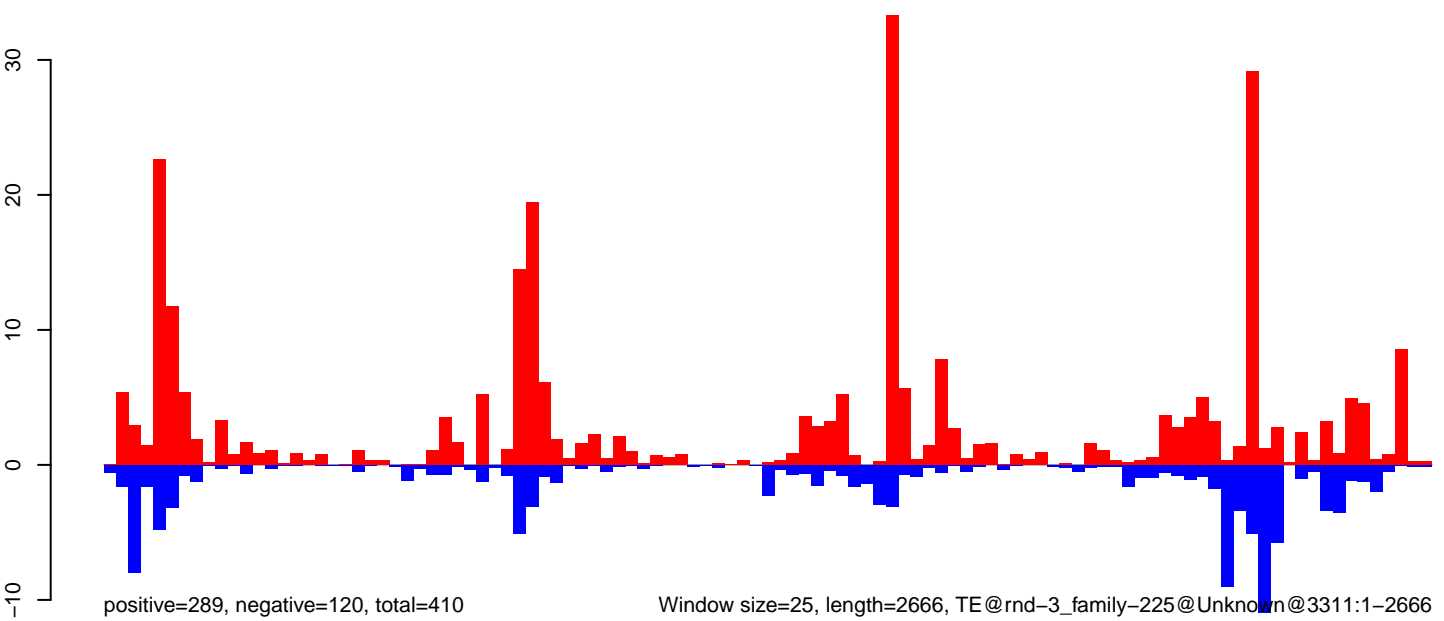
CuQuin_Testes_NL.18_23.rep



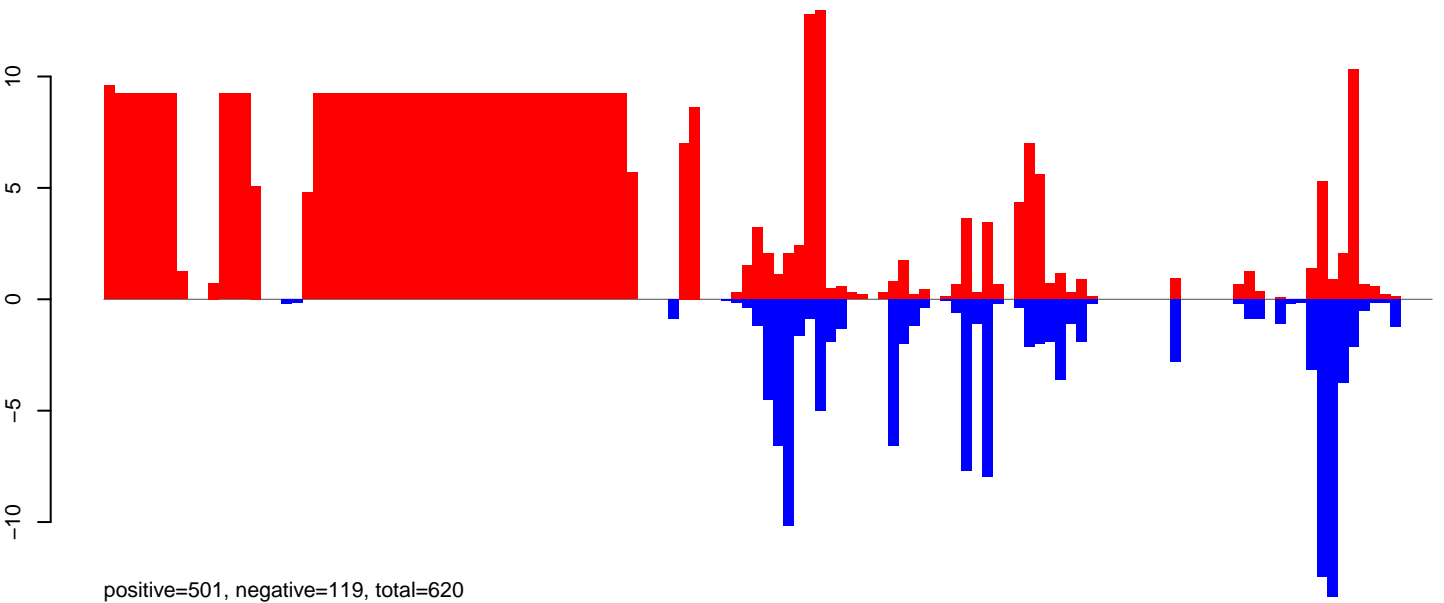
CuQuin_Testes_NL.24_35.rep



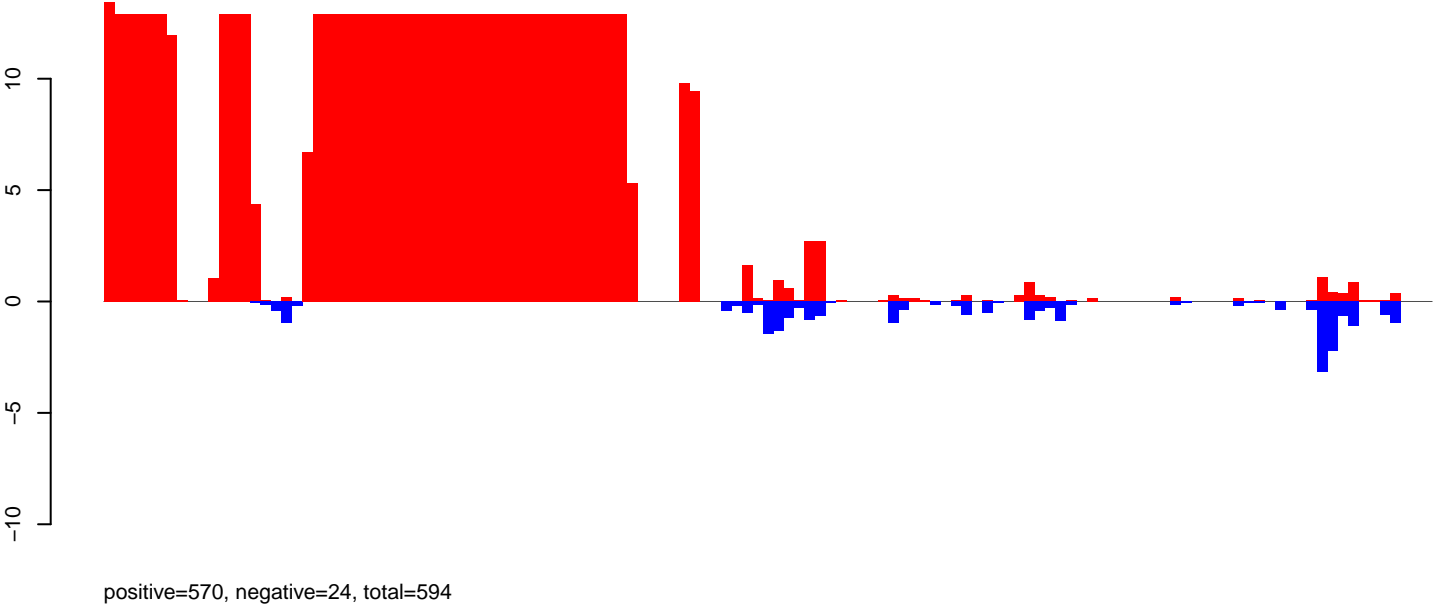
CuQuin_Testes_NL.rep



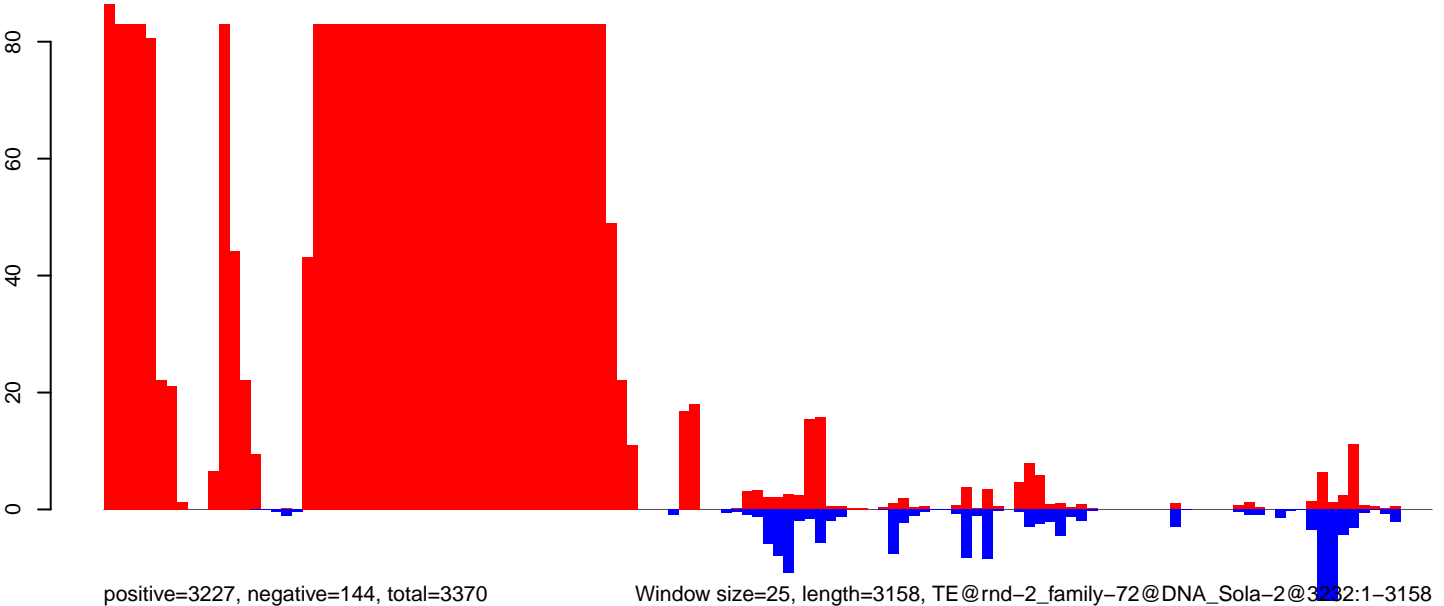
CuQuin_Testes_NL.18_23.rep



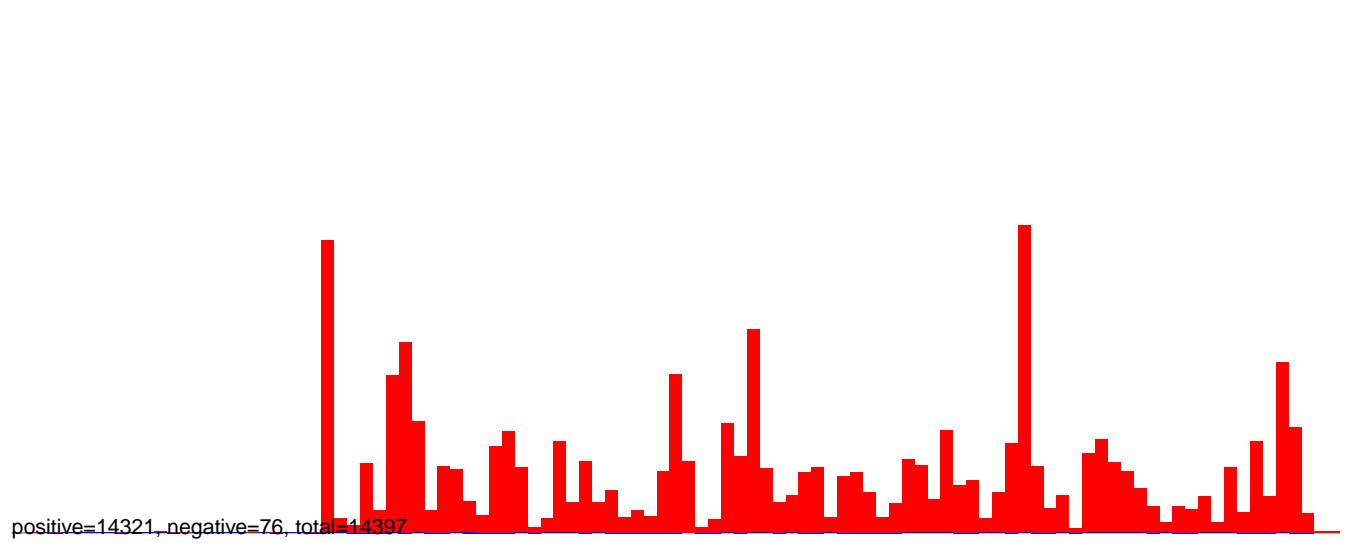
CuQuin_Testes_NL.24_35.rep



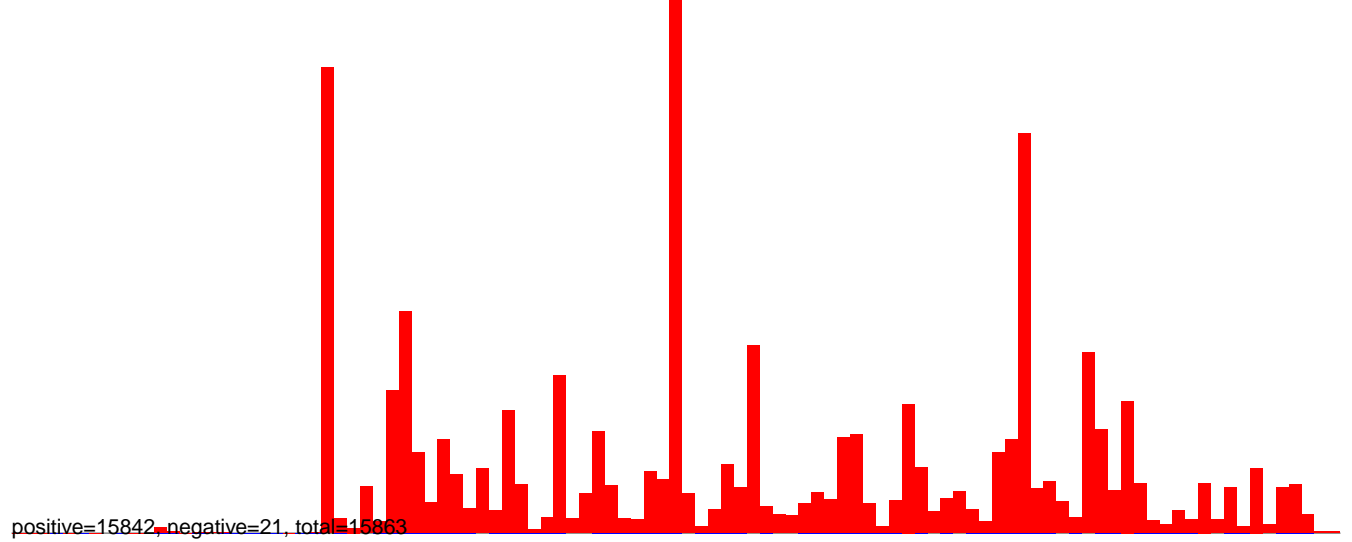
CuQuin_Testes_NL.rep



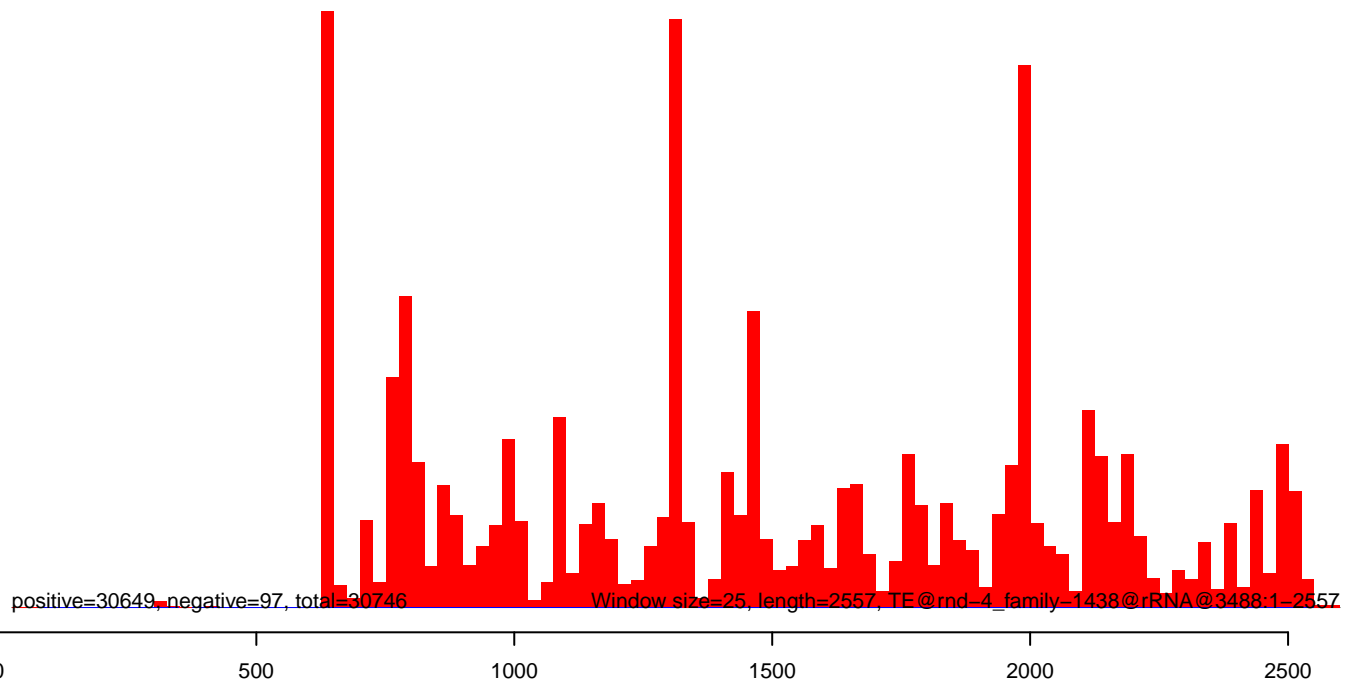
CuQuin_Testes_NL.18_23.rep



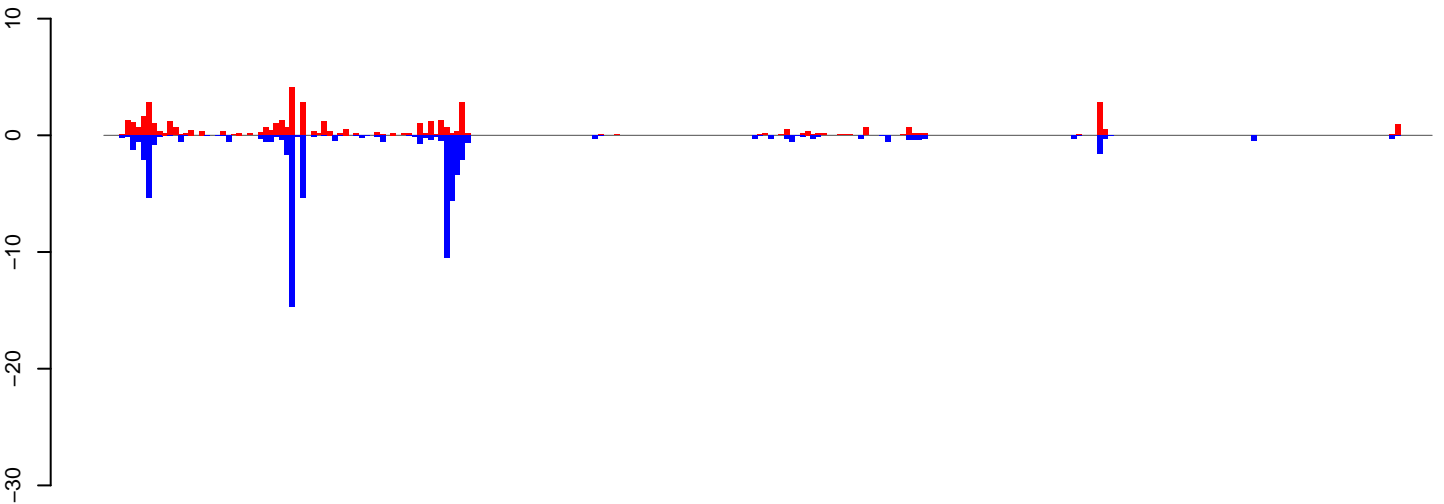
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

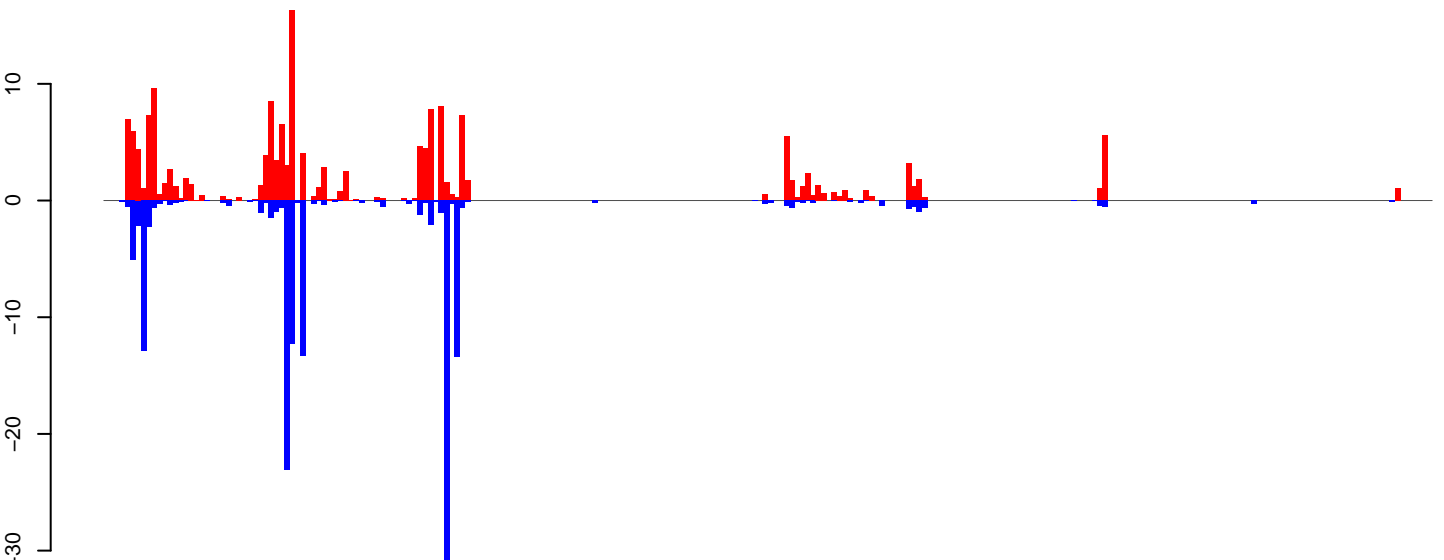


CuQuin_Testes_NL.18_23.rep



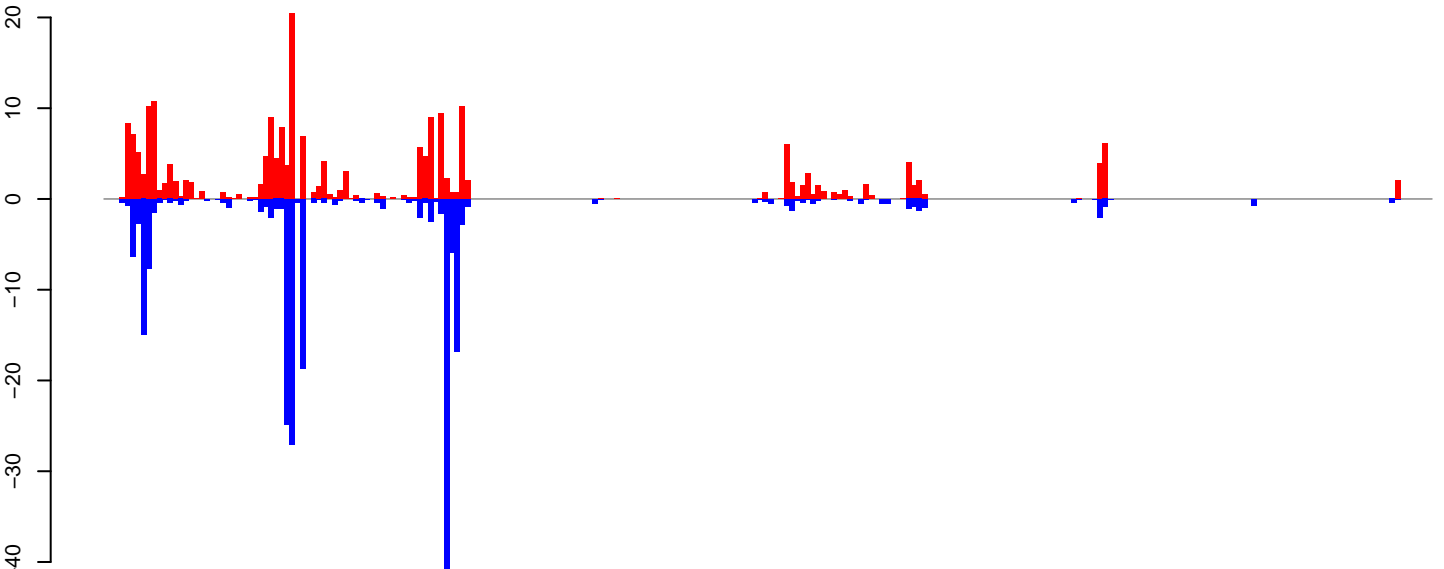
positive=45, negative=70, total=115

CuQuin_Testes_NL.24_35.rep



positive=172, negative=144, total=316

CuQuin_Testes_NL.rep

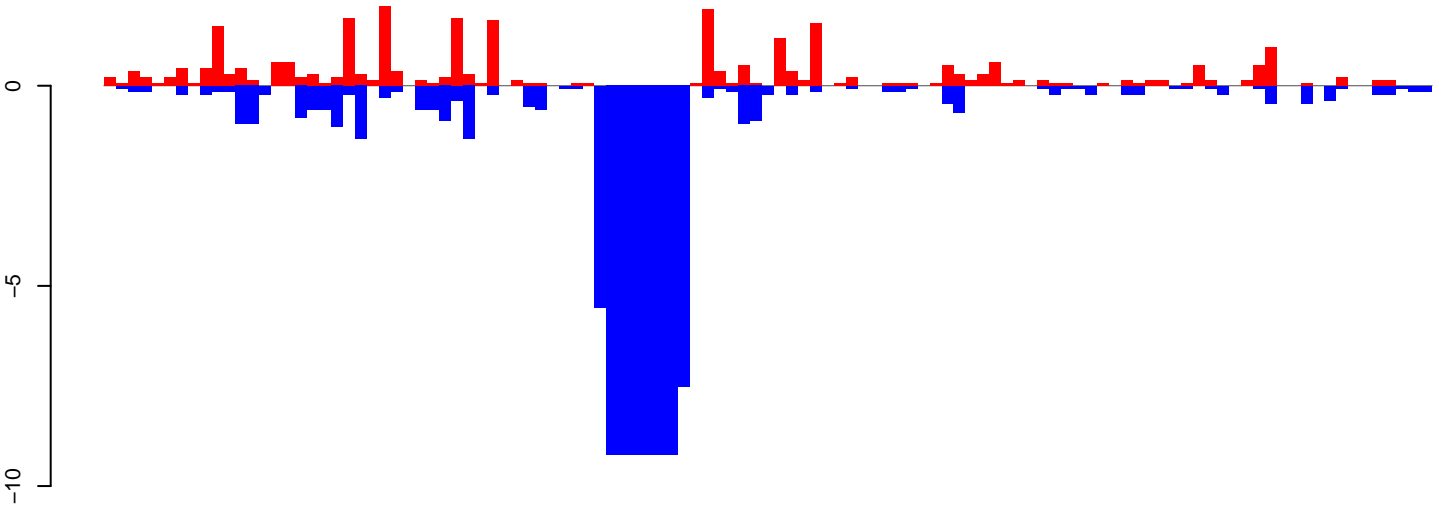


positive=217, negative=214, total=431

Window size=25, length=6247, TE@rnd-4_family-567@Unknown@3608:1-6247

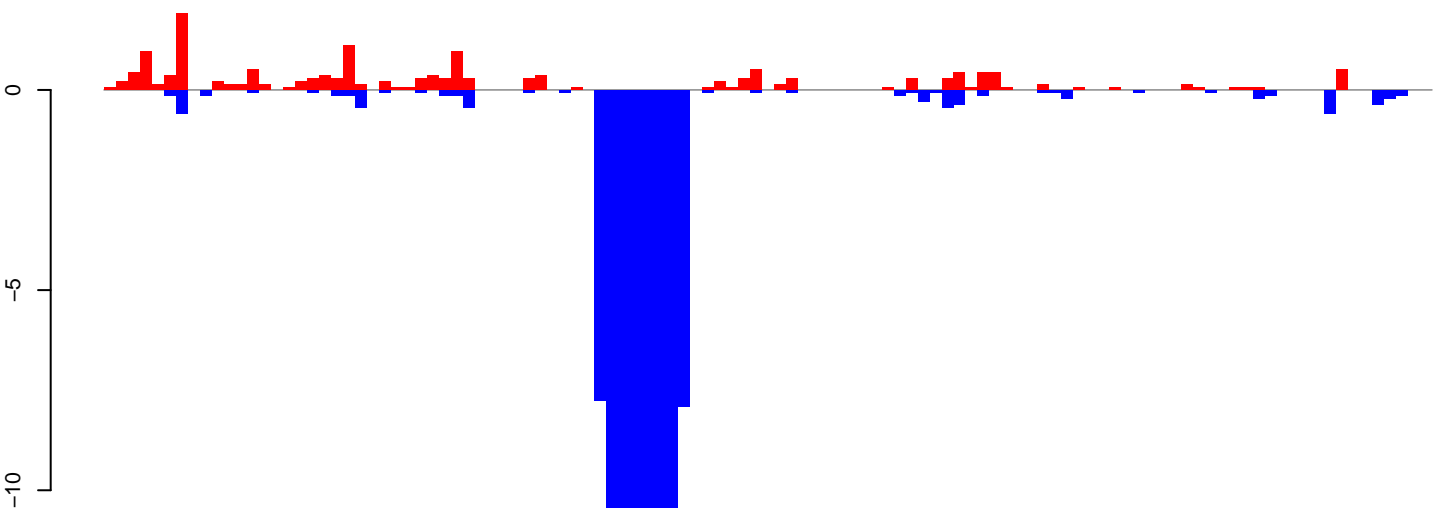
0 1000 2000 3000 4000 5000 6000

CuQuin_Testes_NL.18_23.rep



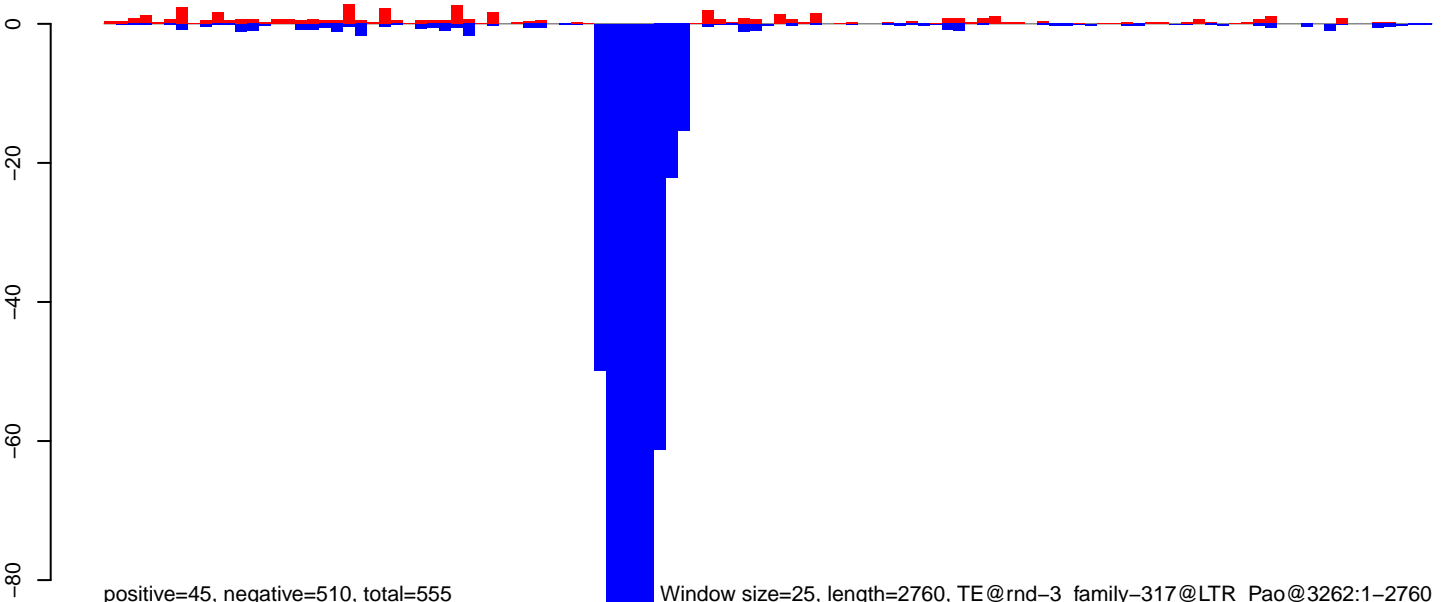
positive=28, negative=90, total=119

CuQuin_Testes_NL.24_35.rep



positive=16, negative=100, total=116

CuQuin_Testes_NL.rep

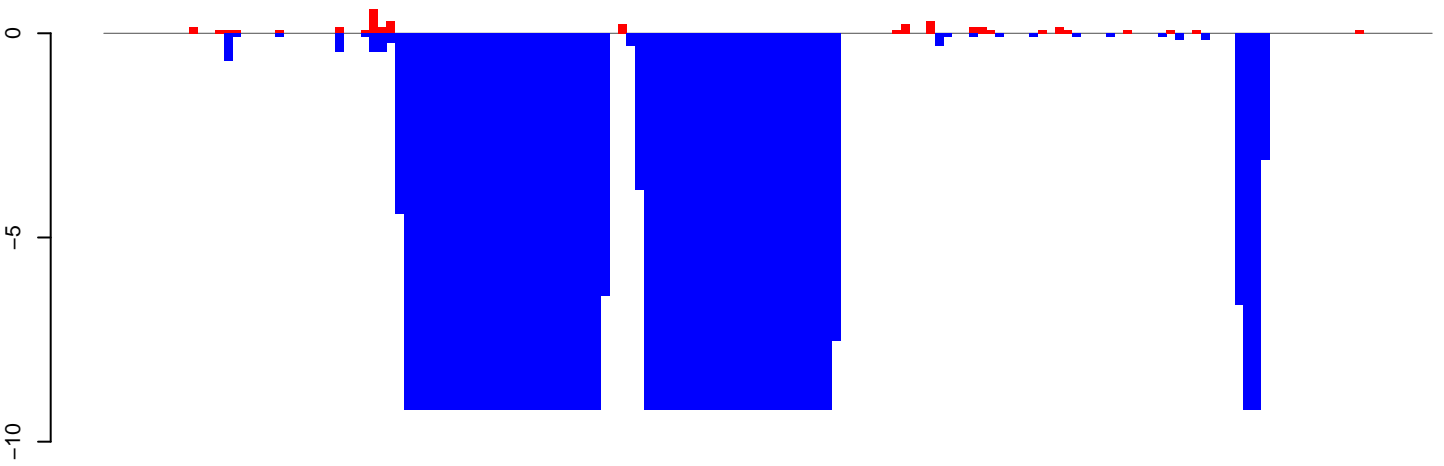


positive=45, negative=510, total=555

Window size=25, length=2760, TE@rnd-3_family-317@LTR_Pao@3262:1-2760

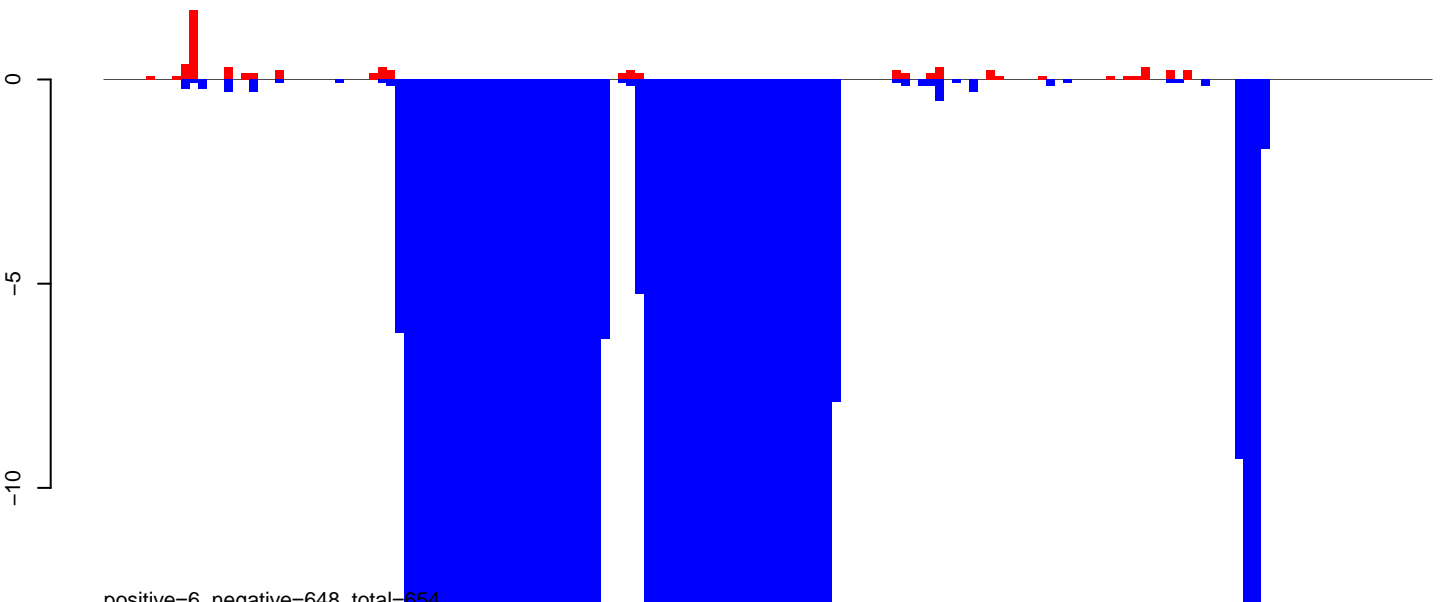
0 500 1000 1500 2000 2500

CuQuin_Testes_NL.18_23.rep



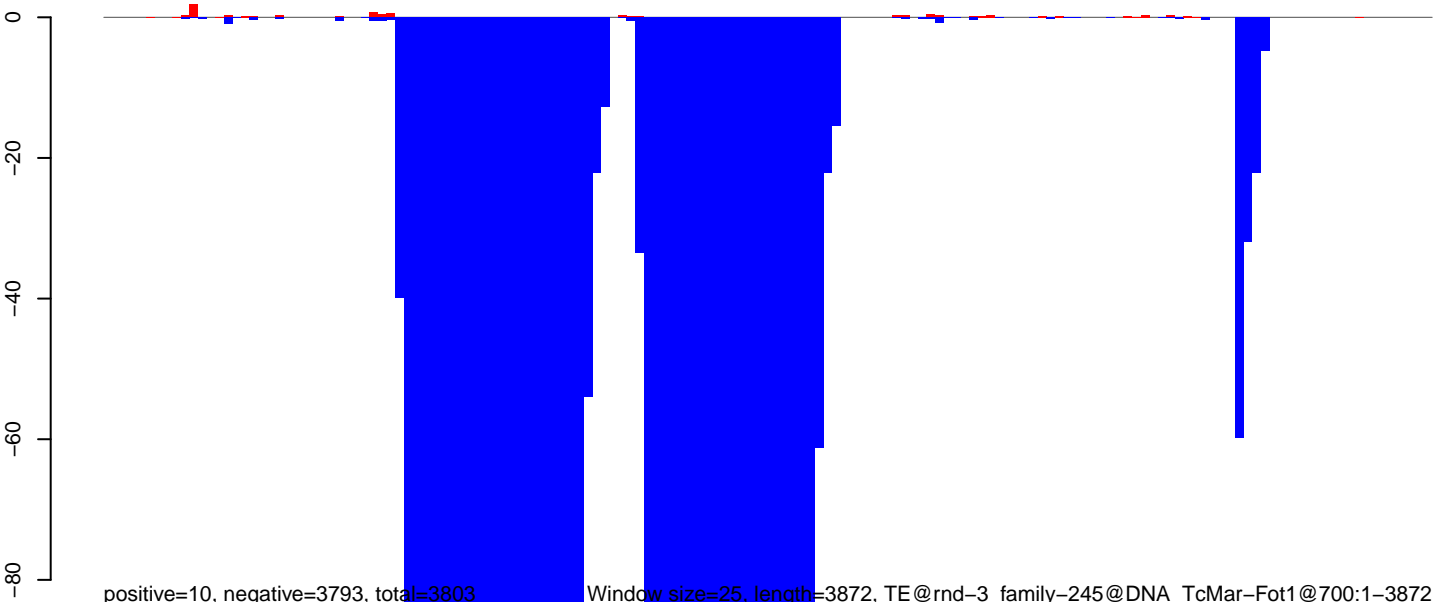
positive=3, negative=470, total=473

CuQuin_Testes_NL.24_35.rep



positive=6, negative=648, total=654

CuQuin_Testes_NL.rep

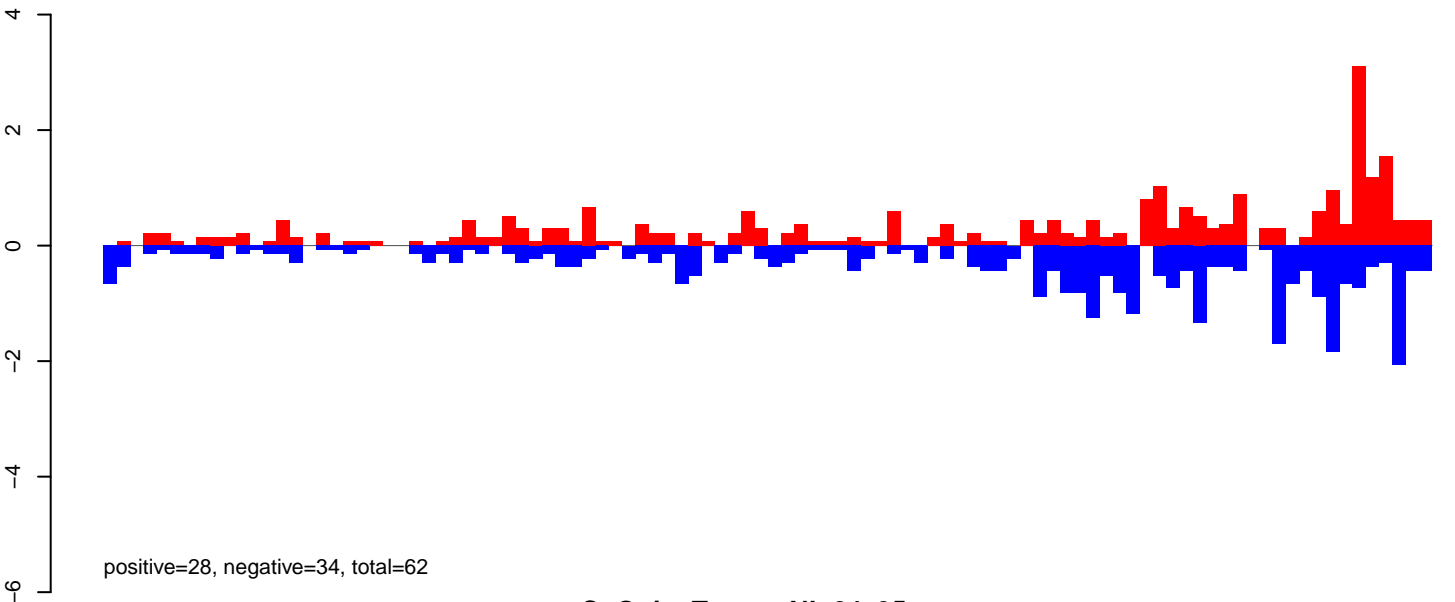


positive=10, negative=3793, total=3803

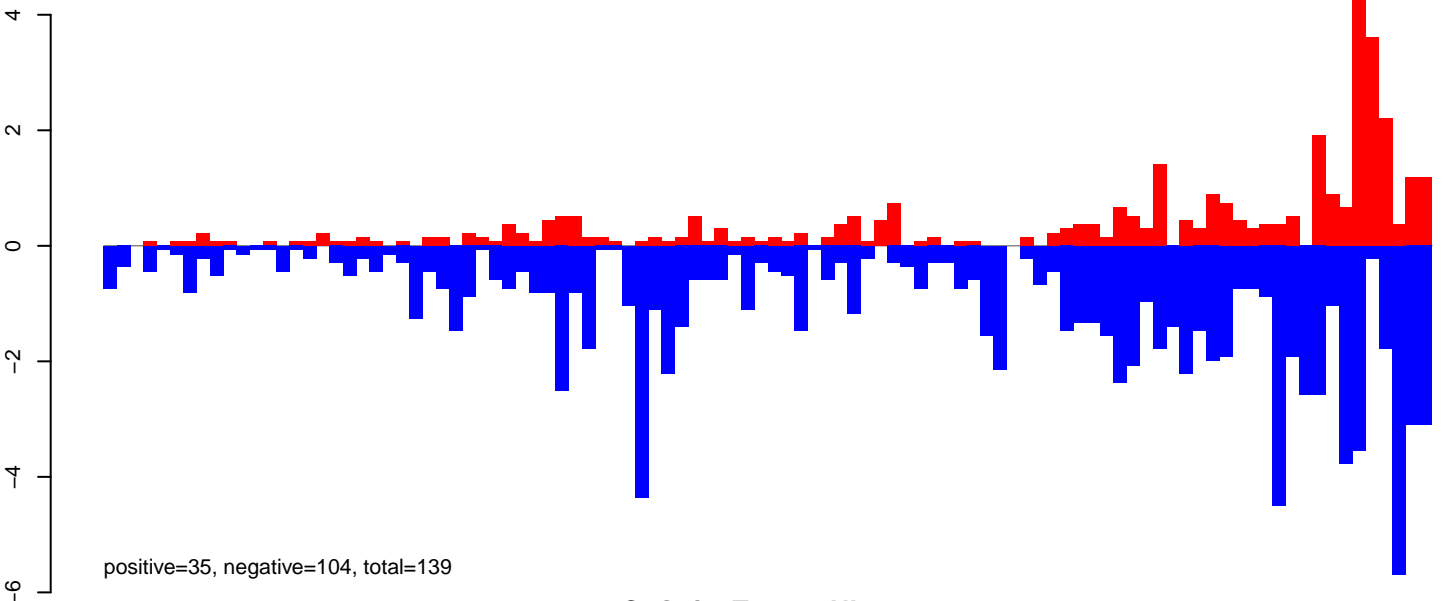
Window size=25, length=3872, TE@rnd-3_family-245@DNA_TcMar-Fot1@700:1-3872

0 1000 2000 3000 4000

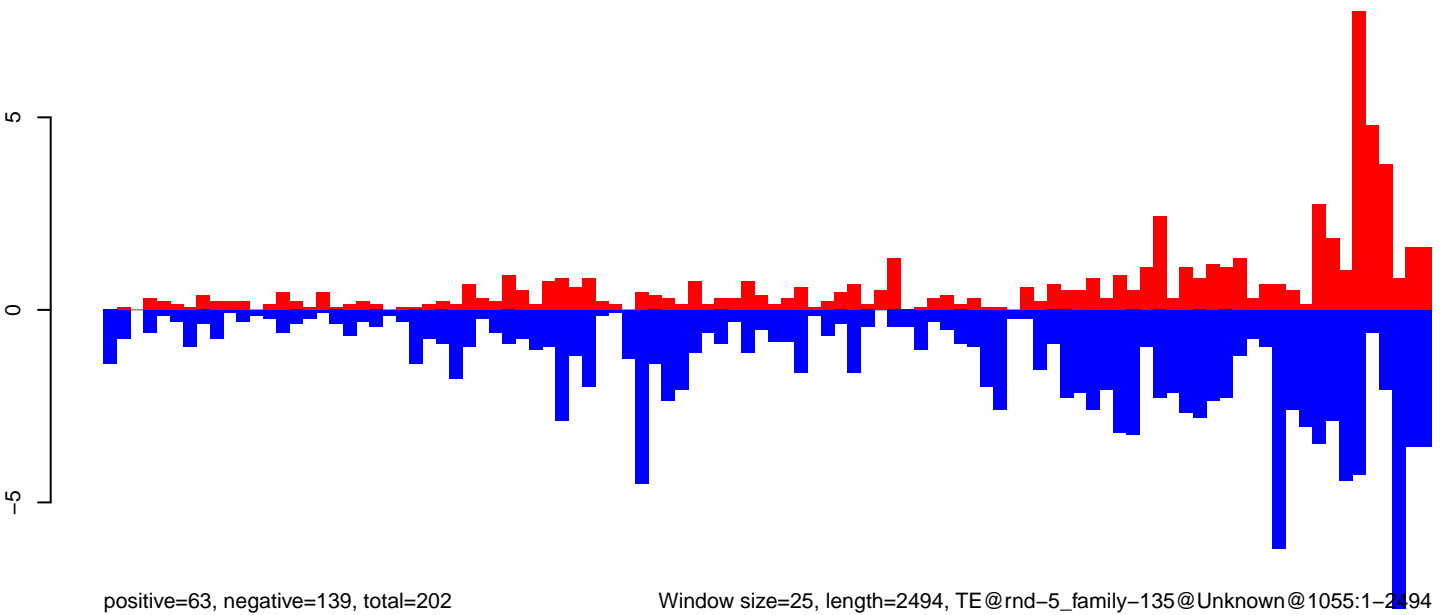
CuQuin_Testes_NL.18_23.rep



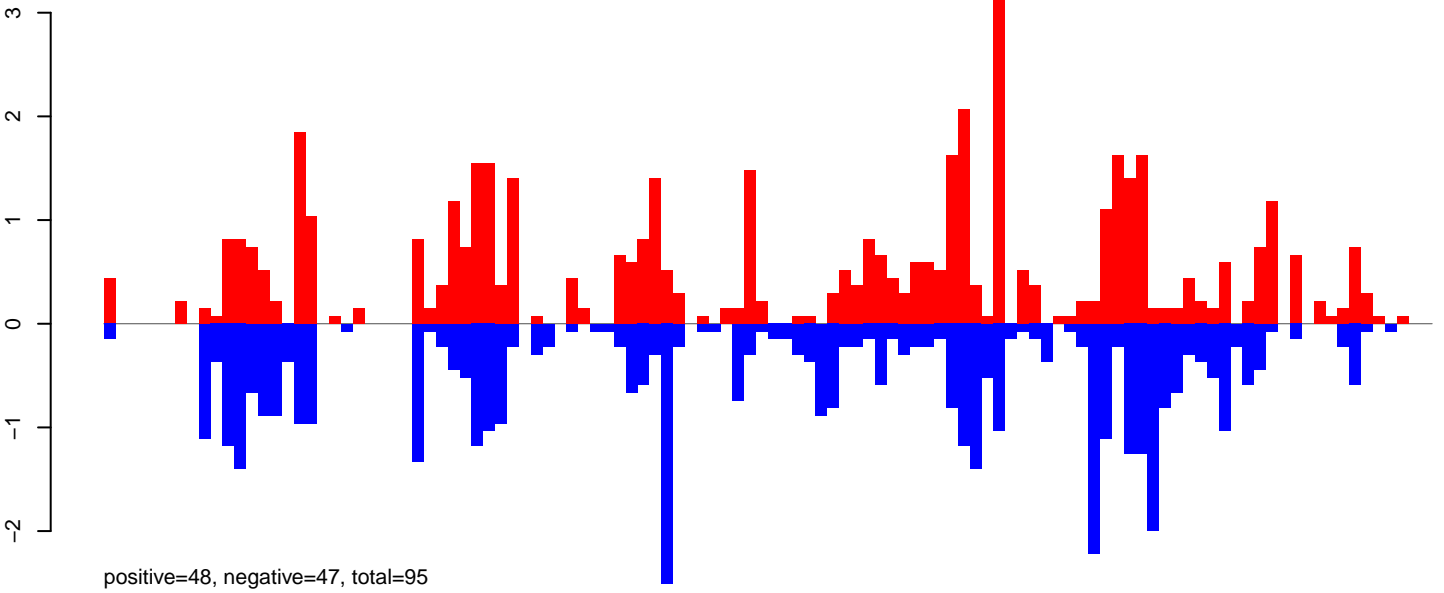
CuQuin_Testes_NL.24_35.rep



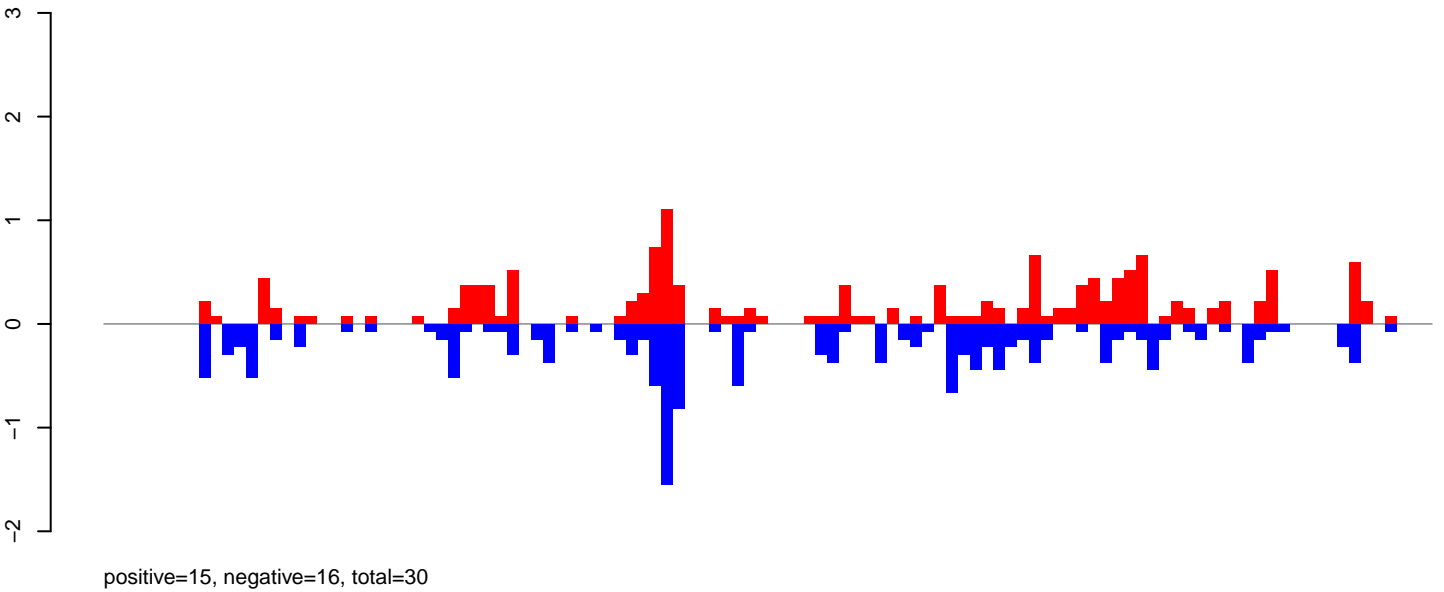
CuQuin_Testes_NL.rep



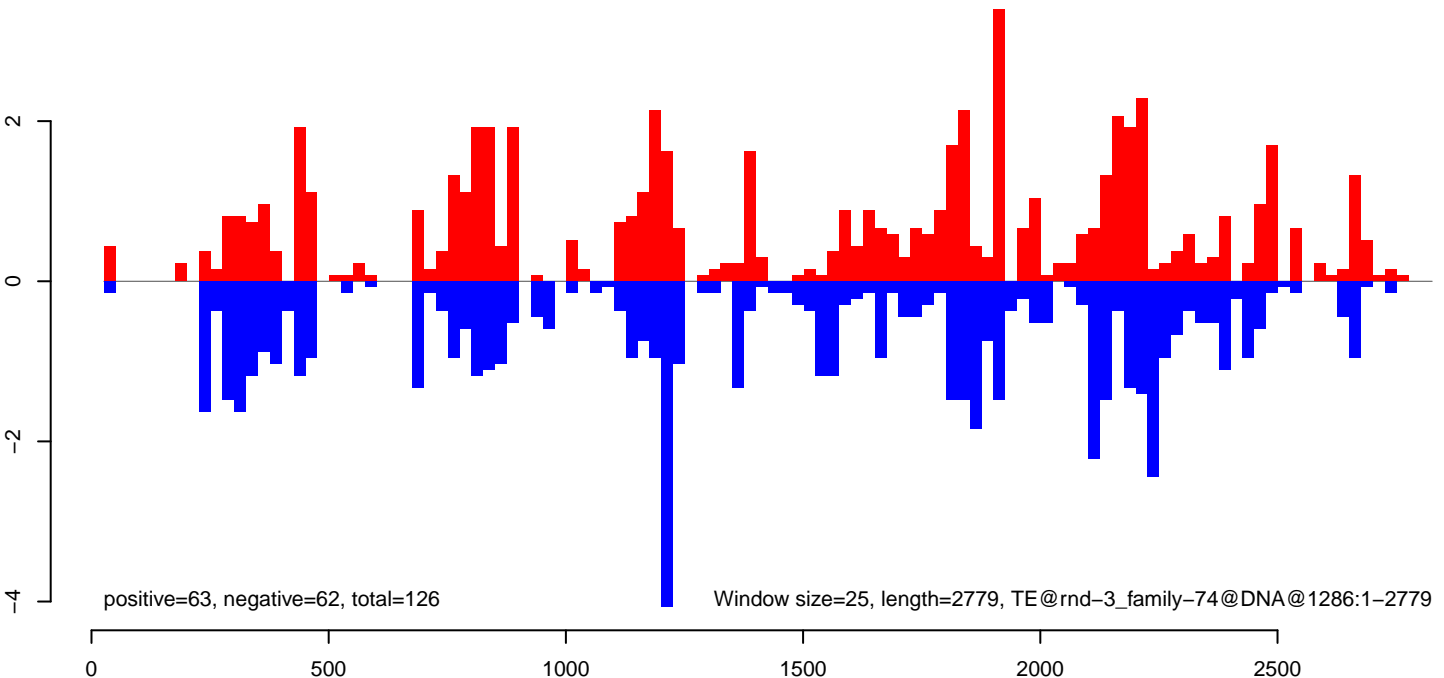
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



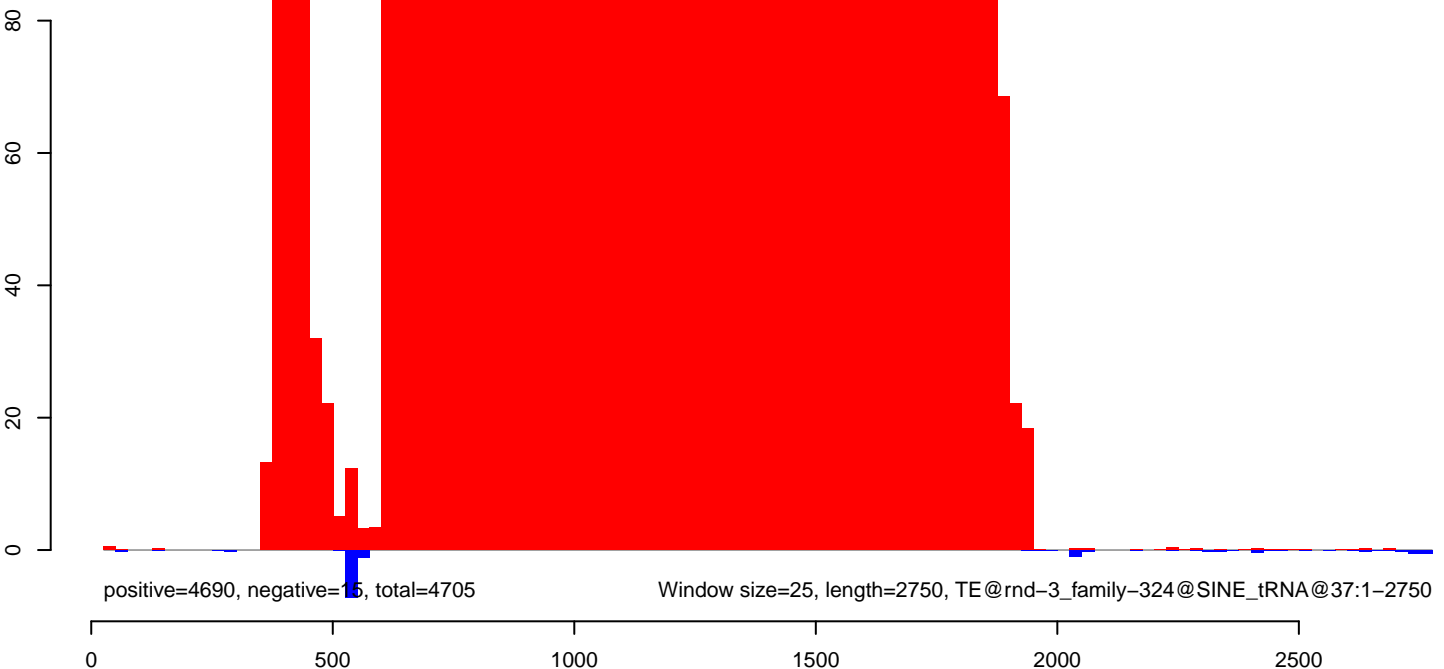
CuQuin_Testes_NL.18_23.rep



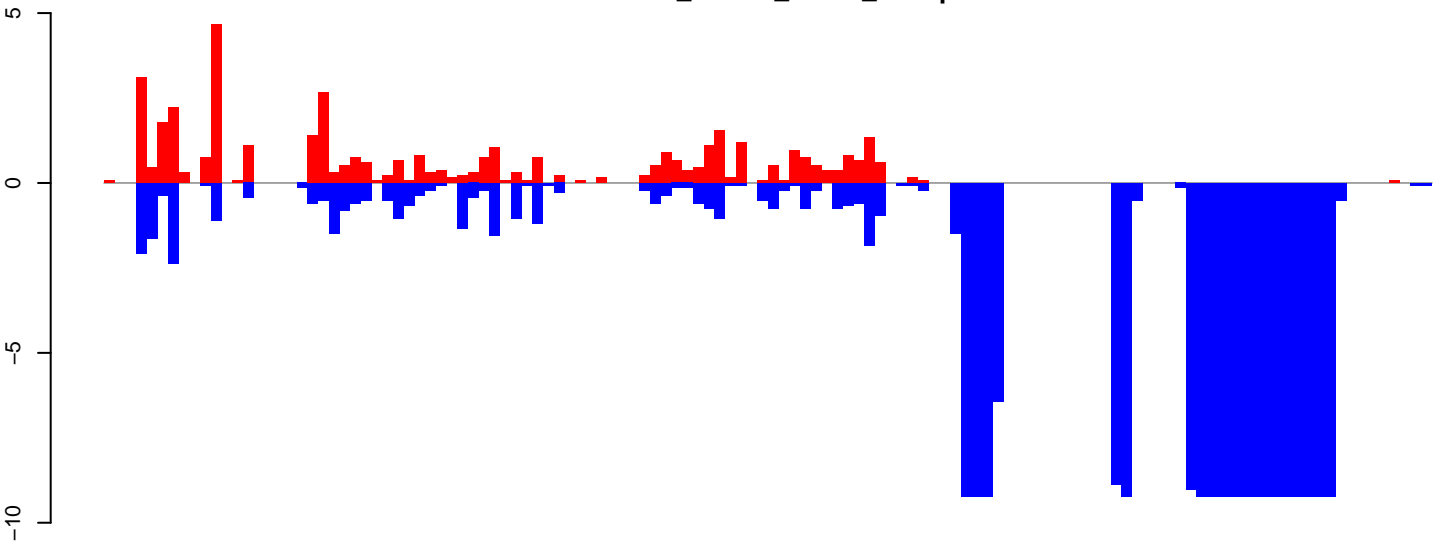
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

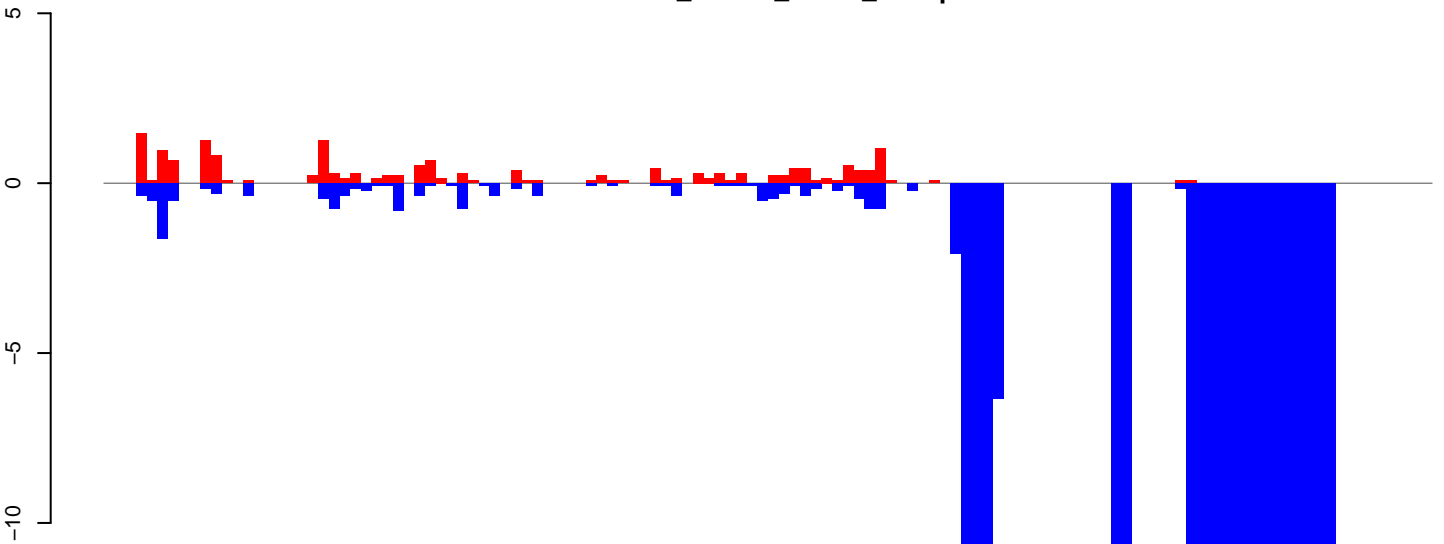


CuQuin_Testes_NL.18_23.rep



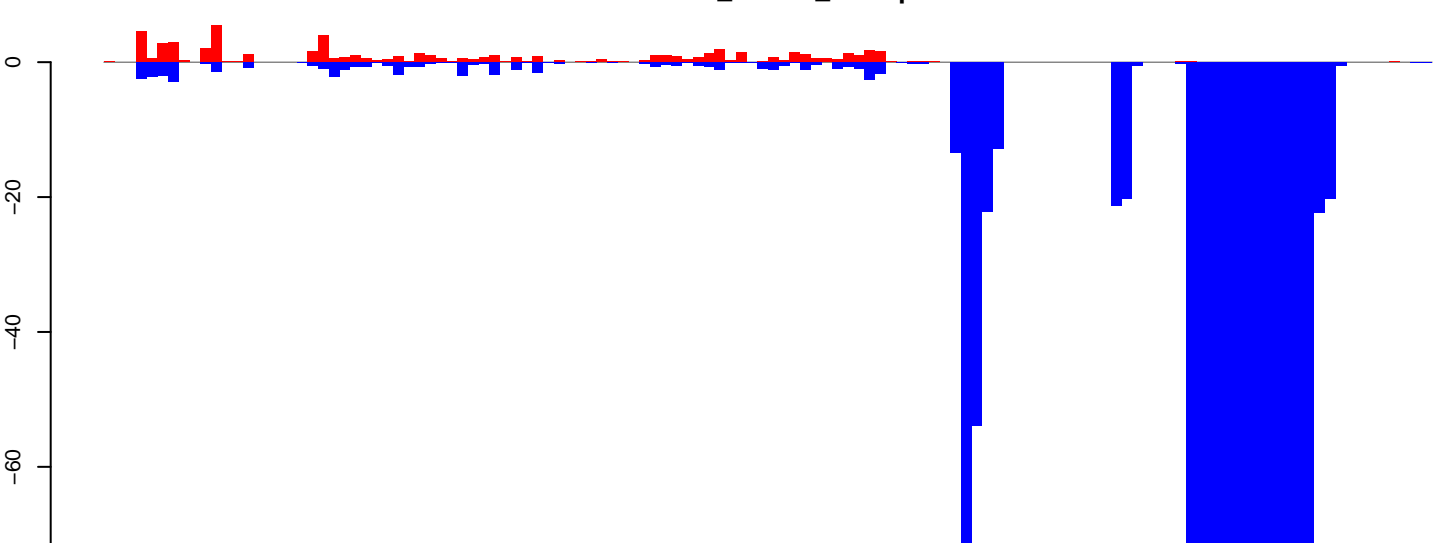
positive=42, negative=217, total=259

CuQuin_Testes_NL.24_35.rep



positive=17, negative=263, total=280

CuQuin_Testes_NL.rep

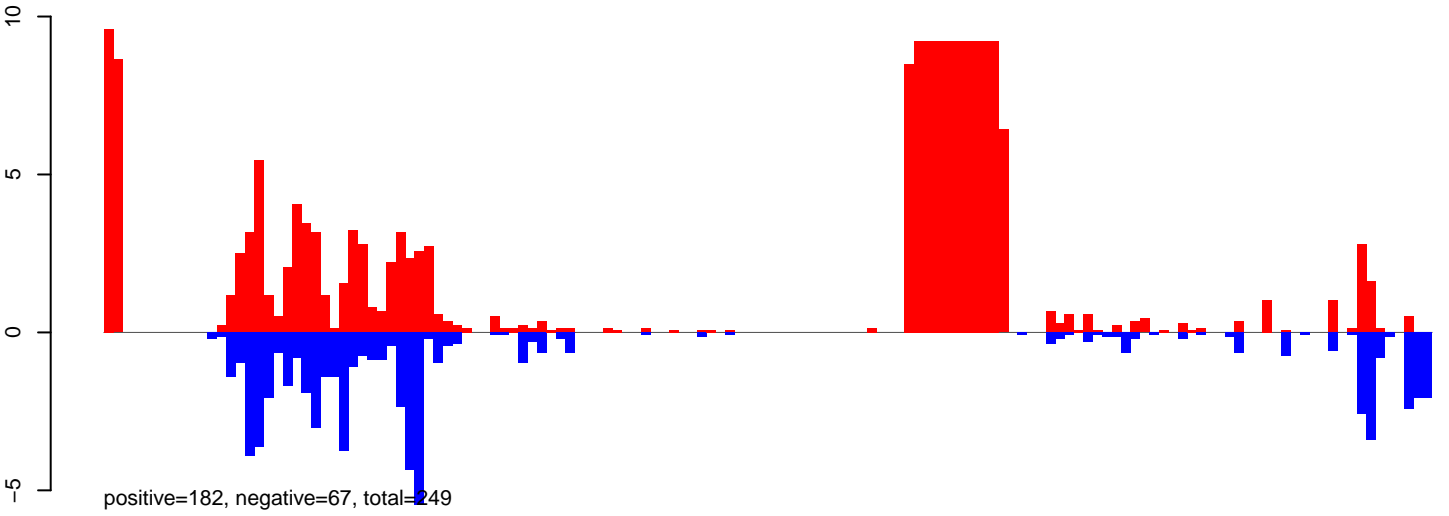


positive=59, negative=1305, total=1363

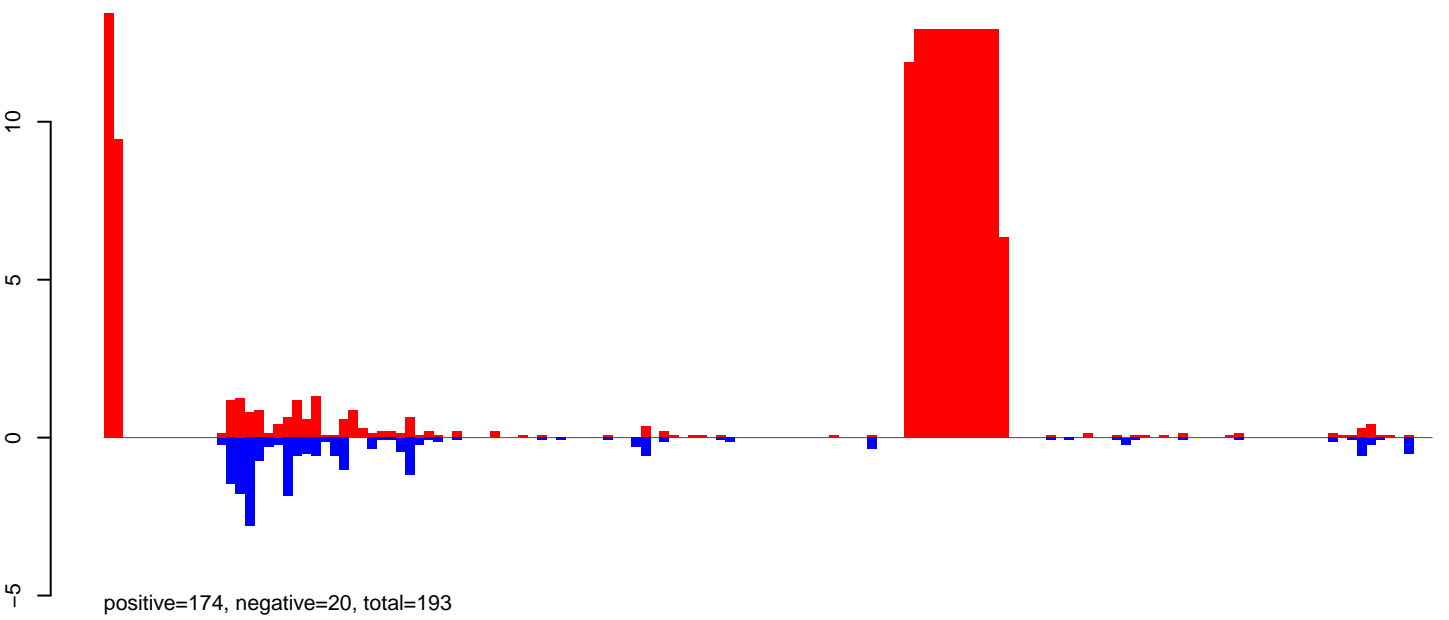
Window size=25, length=3086, TE@rnd-3_family-210@Unknown@38:1-3086

0 500 1000 1500 2000 2500 3000

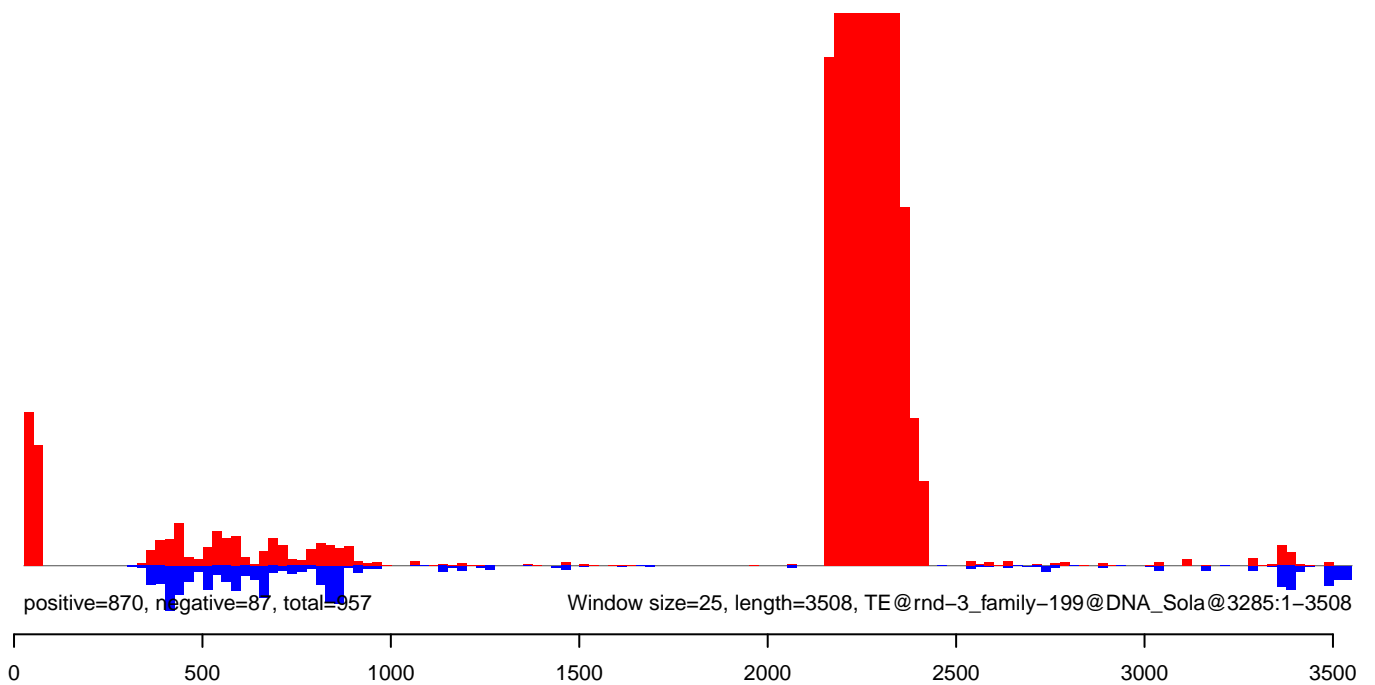
CuQuin_Testes_NL.18_23.rep



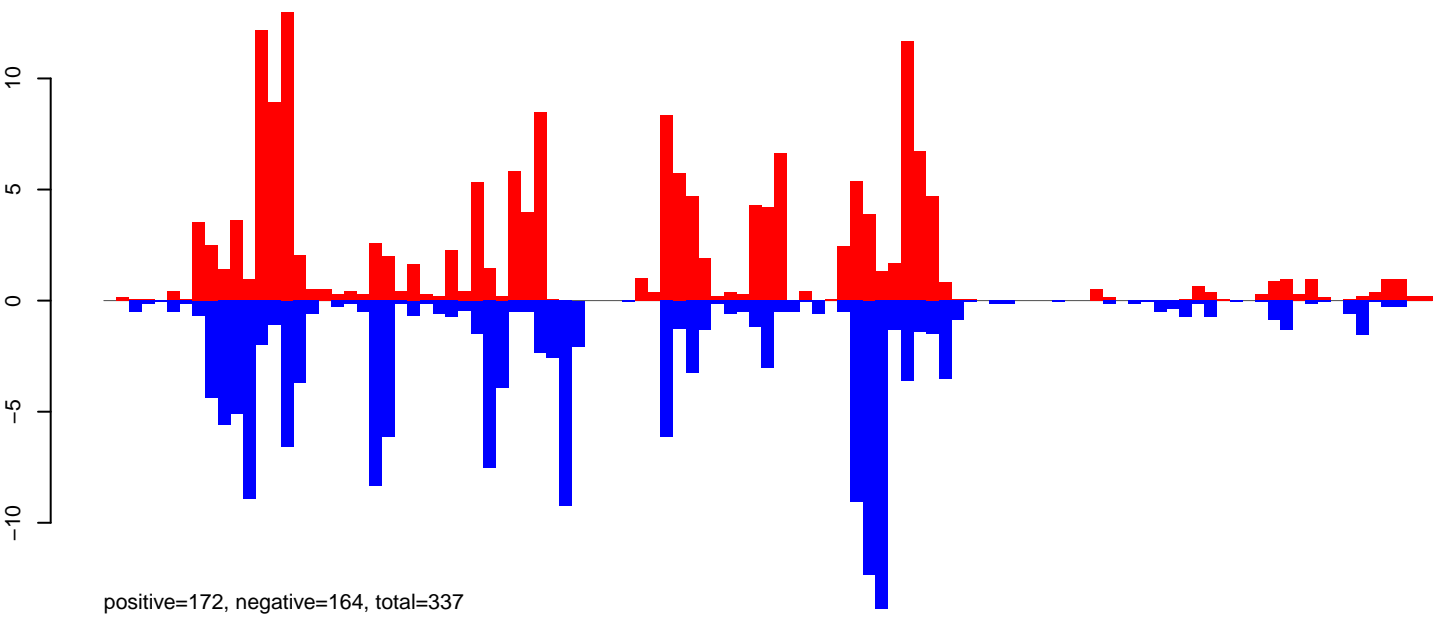
CuQuin_Testes_NL.24_35.rep



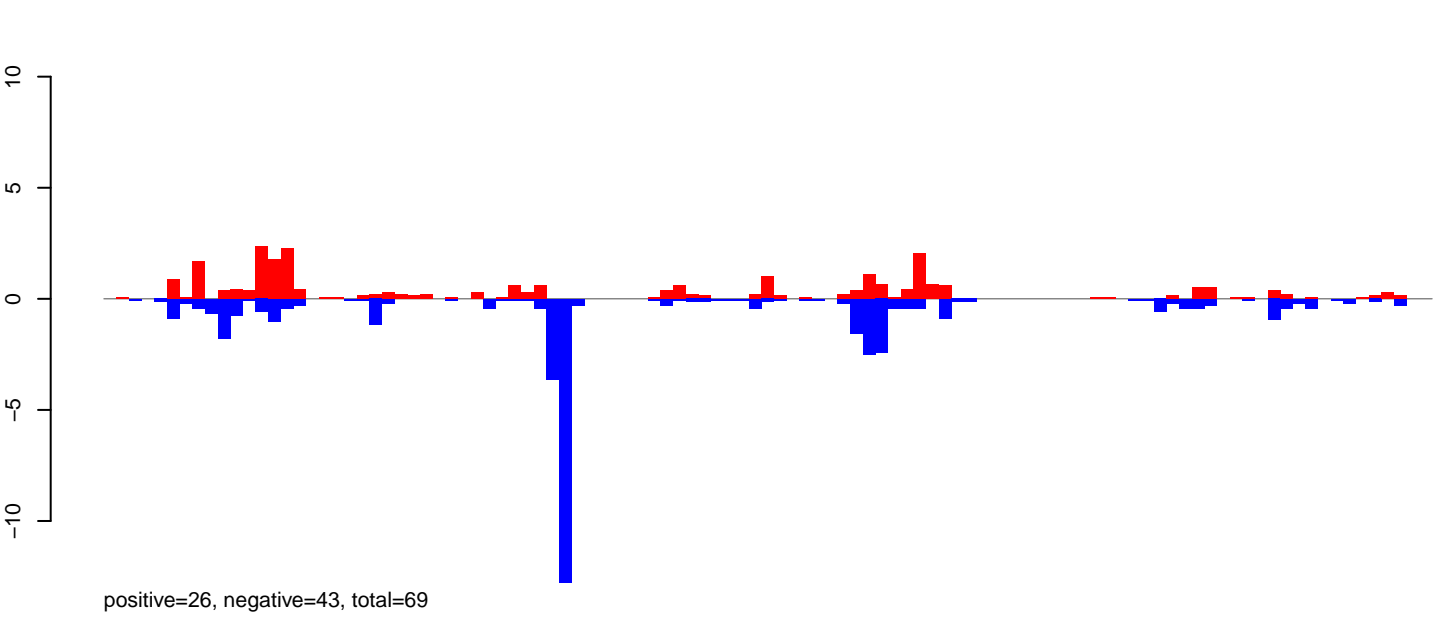
CuQuin_Testes_NL.rep



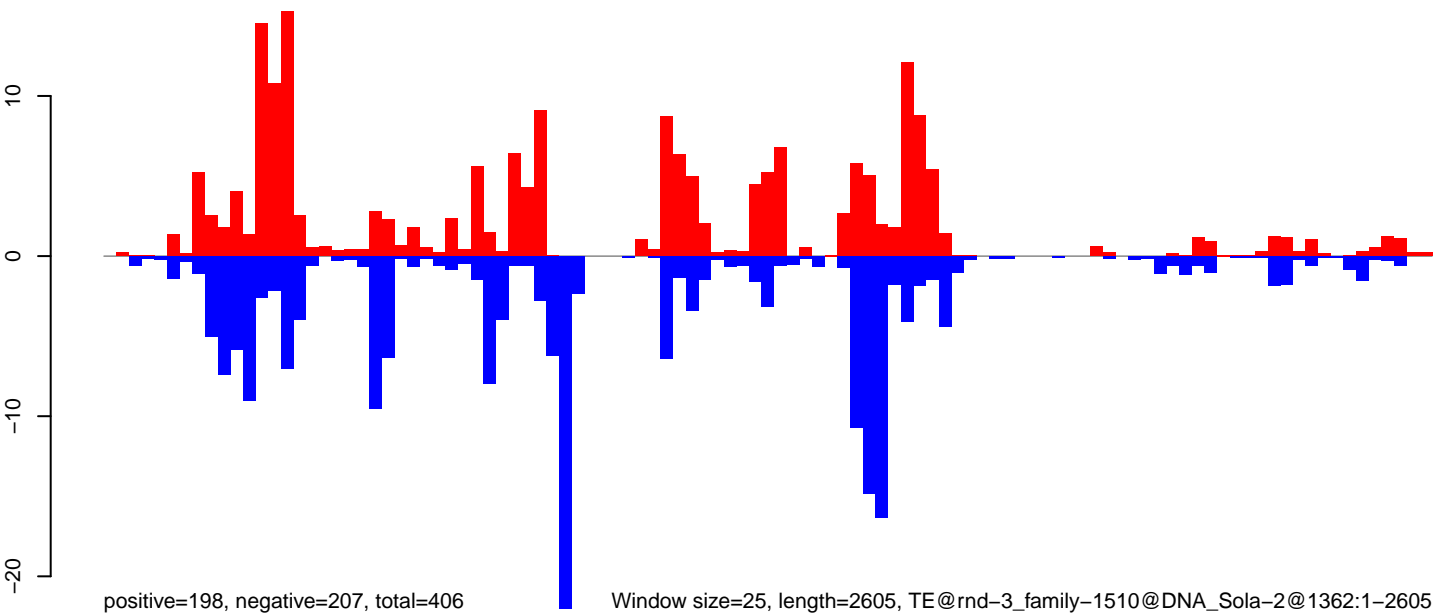
CuQuin_Testes_NL.18_23.rep



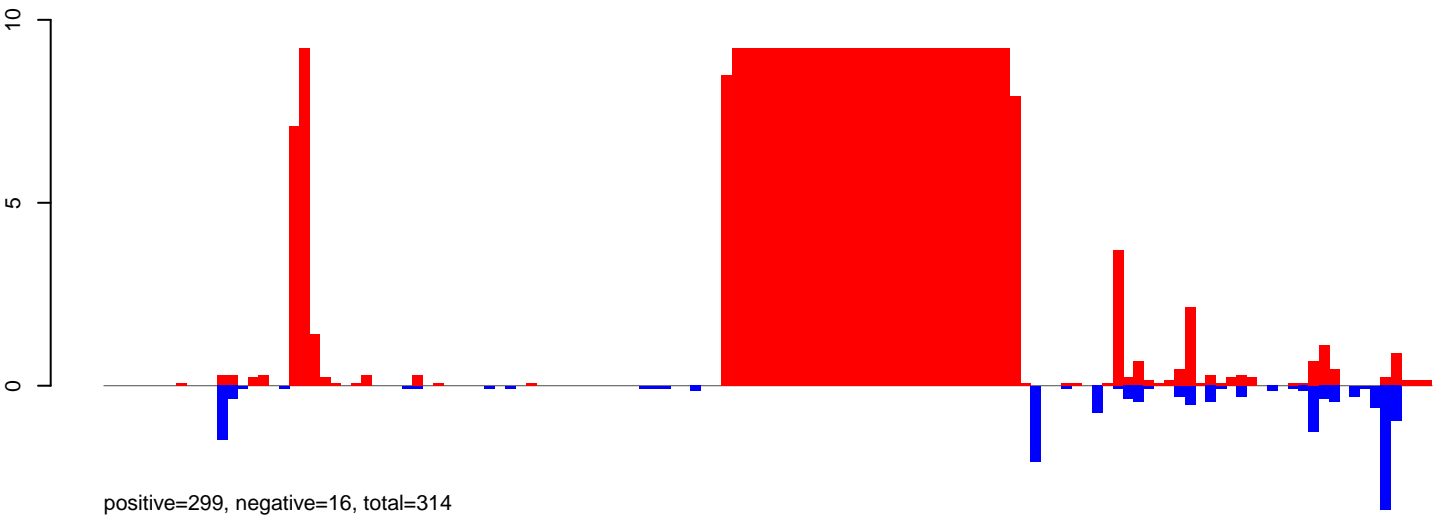
CuQuin_Testes_NL.24_35.rep



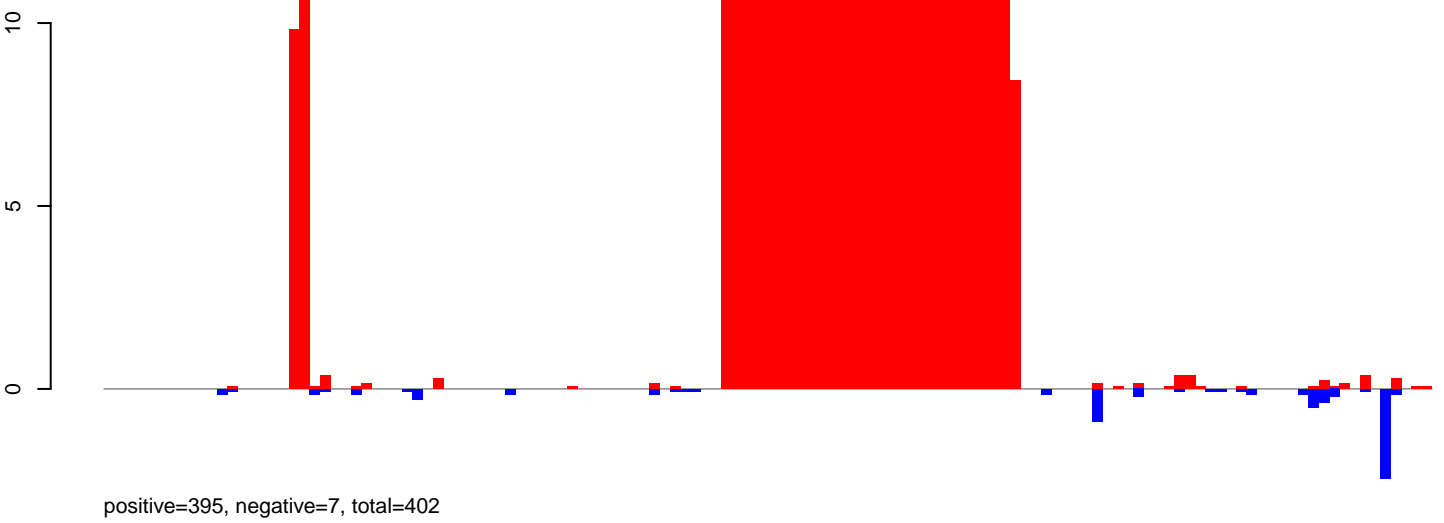
CuQuin_Testes_NL.rep



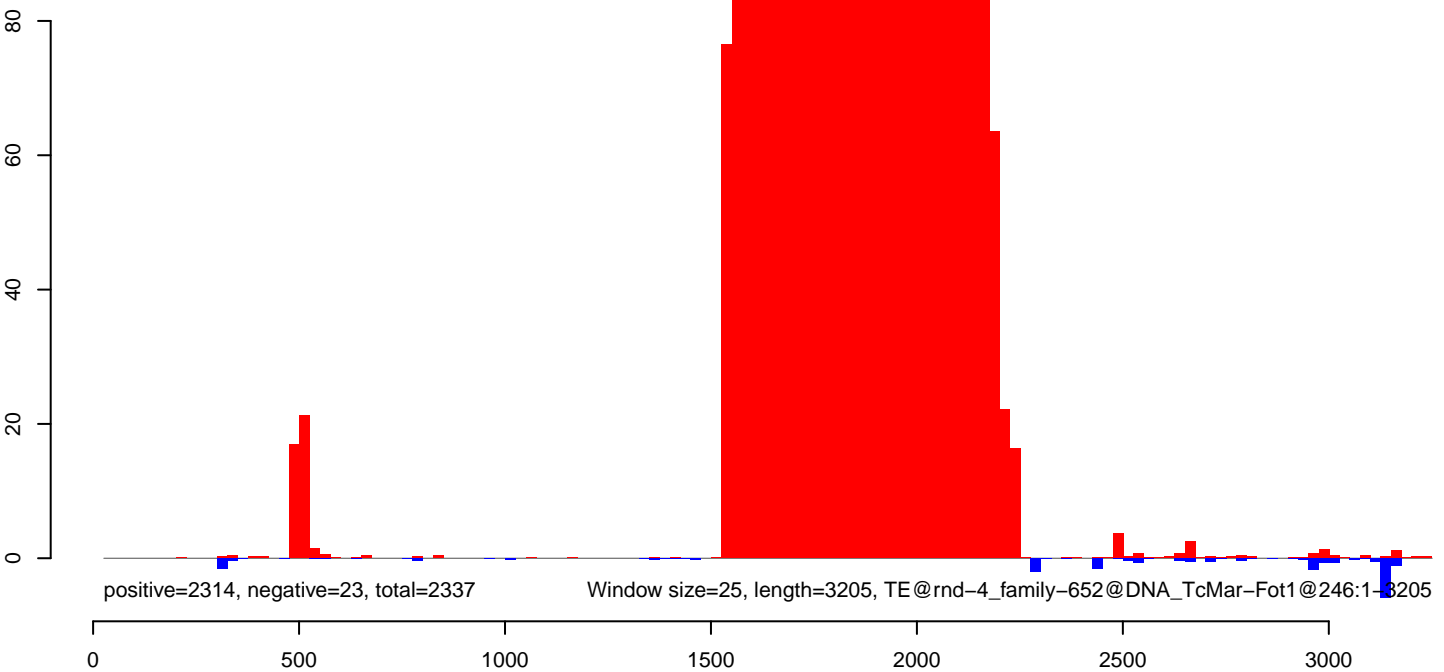
CuQuin_Testes_NL.18_23.rep



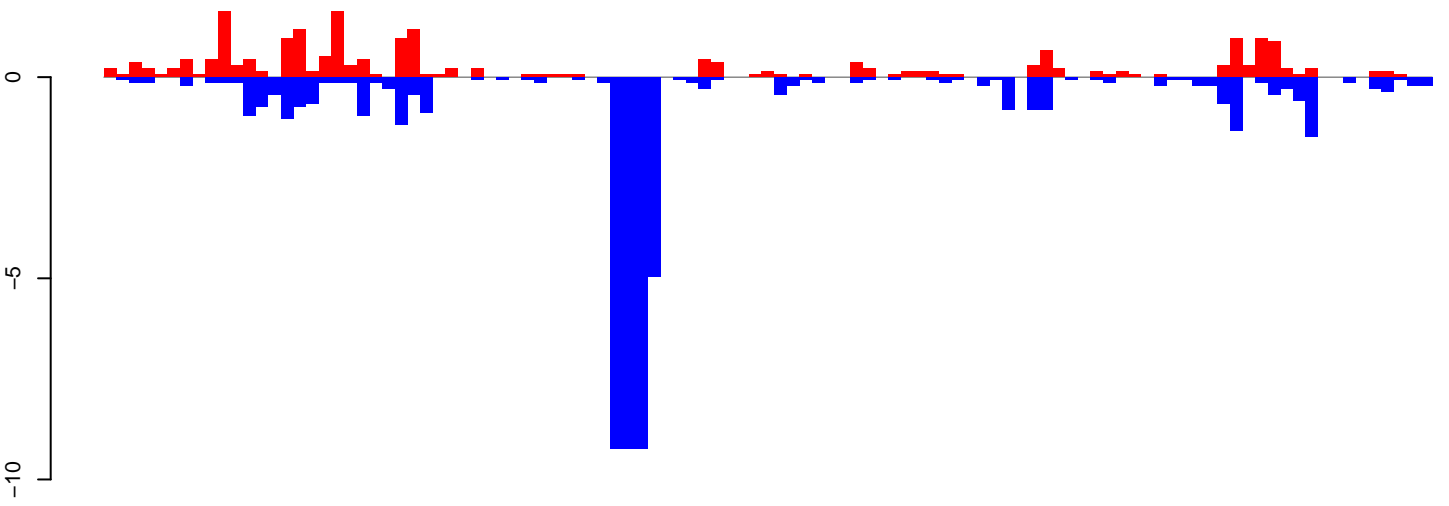
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

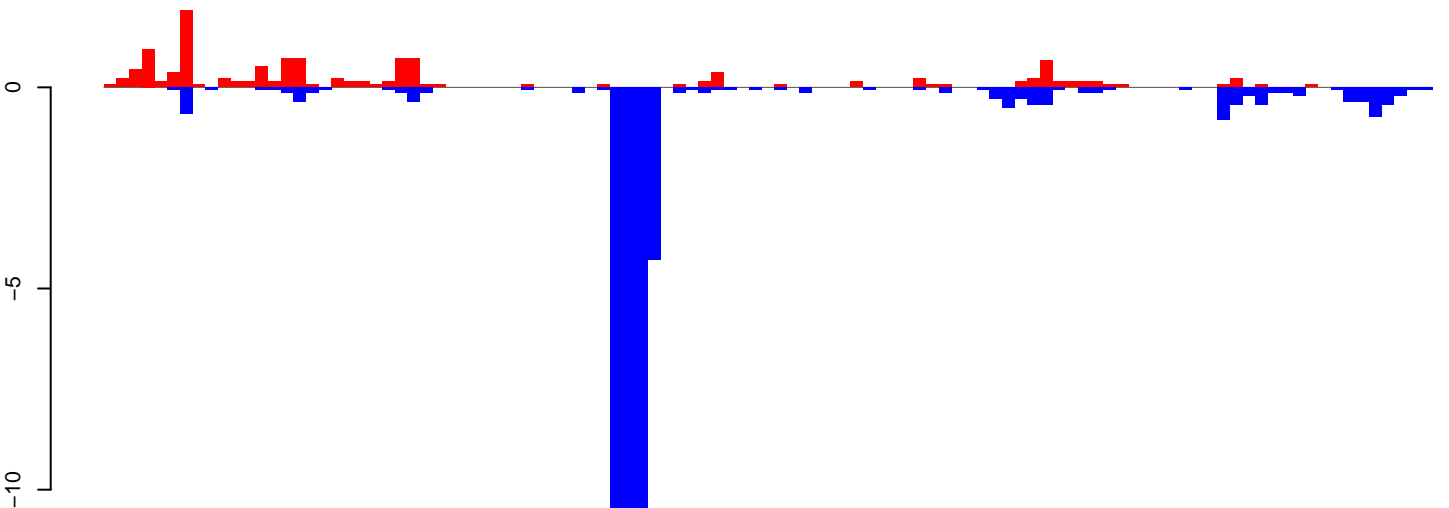


CuQuin_Testes_NL.18_23.rep



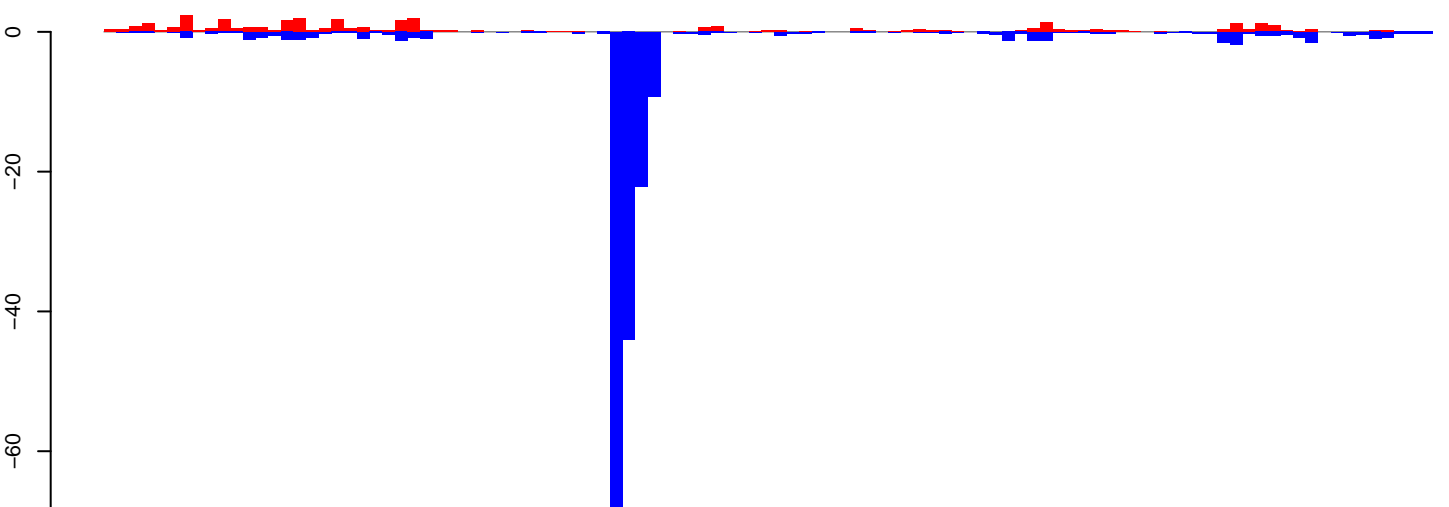
positive=21, negative=55, total=77

CuQuin_Testes_NL.24_35.rep



positive=13, negative=54, total=67

CuQuin_Testes_NL.rep

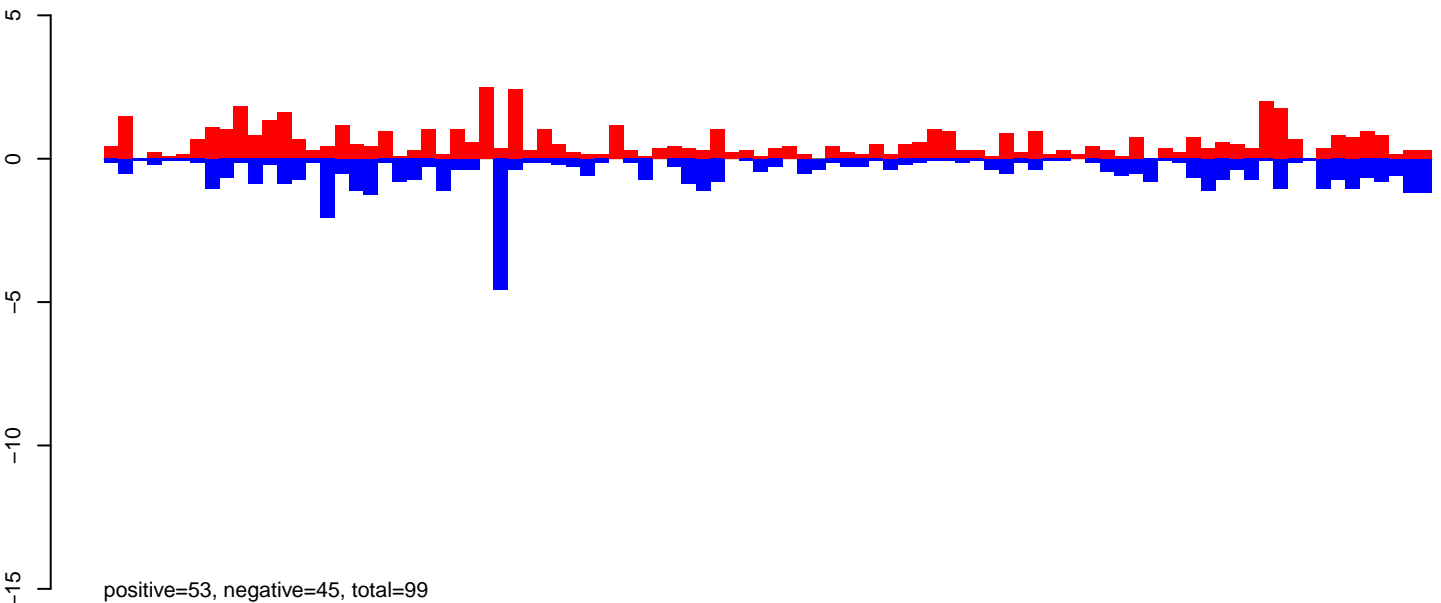


positive=34, negative=193, total=227

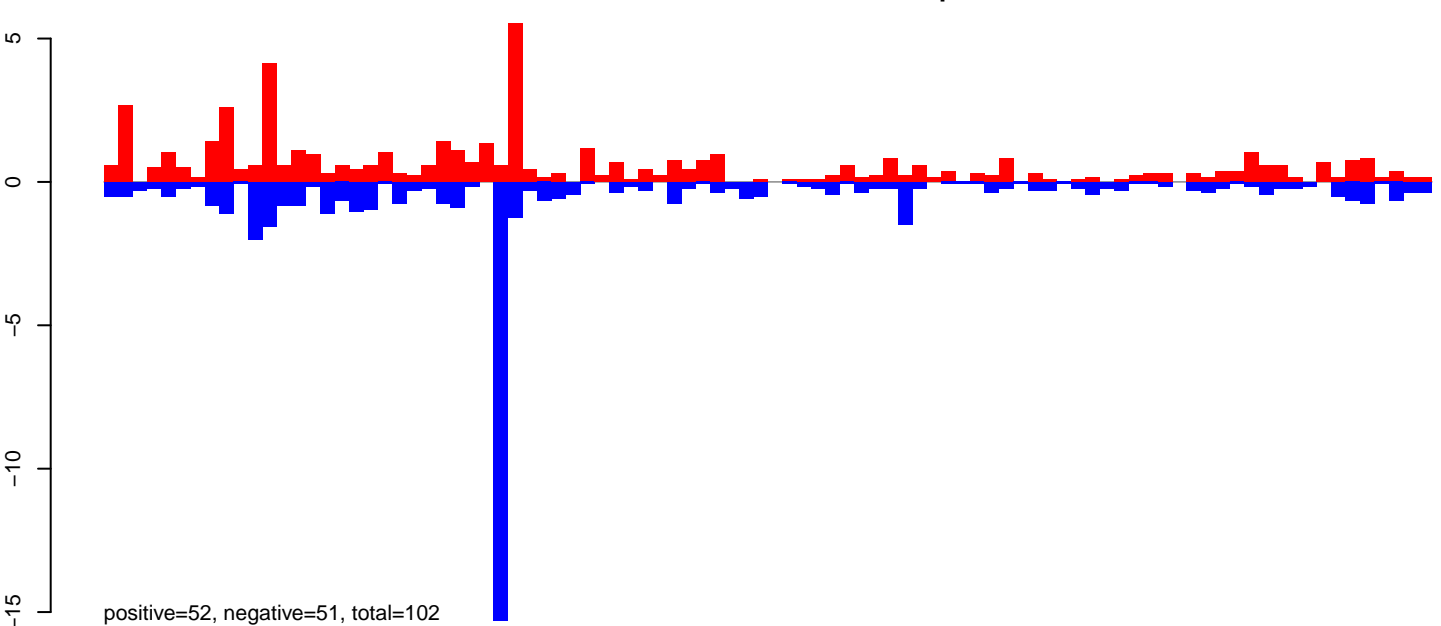
Window size=25, length=2625, TE@rnd-2_family-5@LTR_Pao@2567:1-2625

0 500 1000 1500 2000 2500

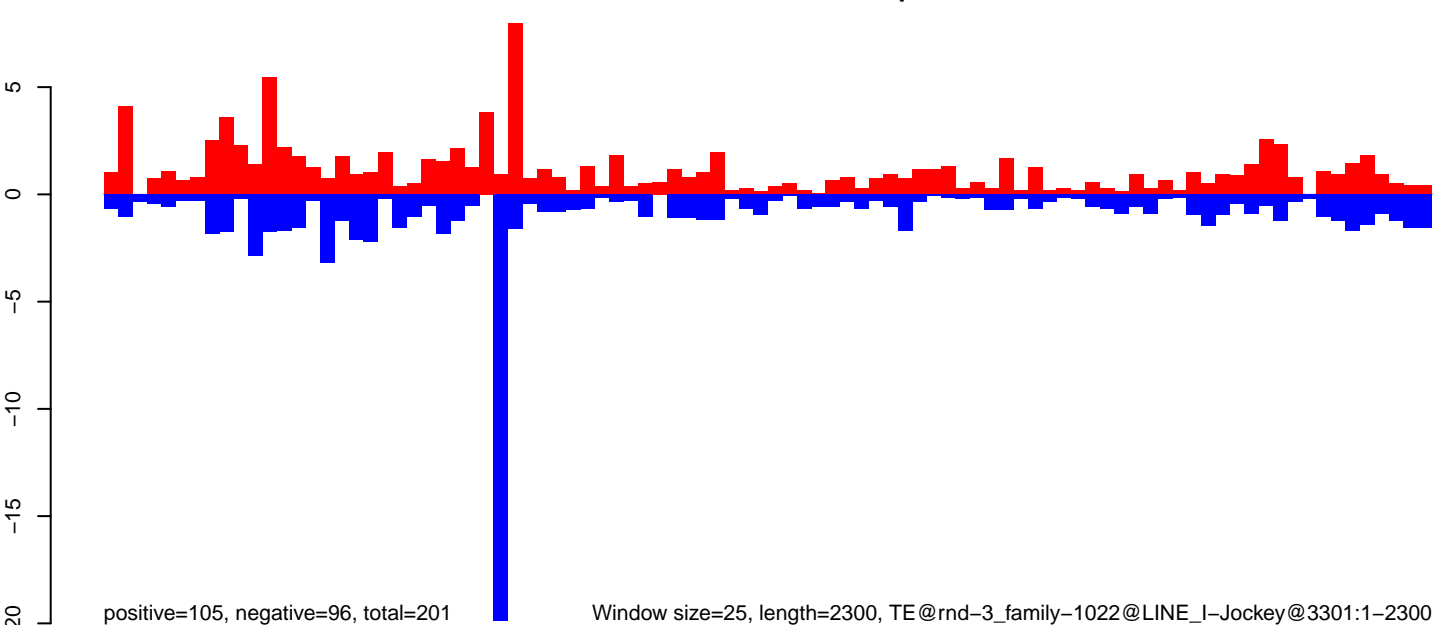
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

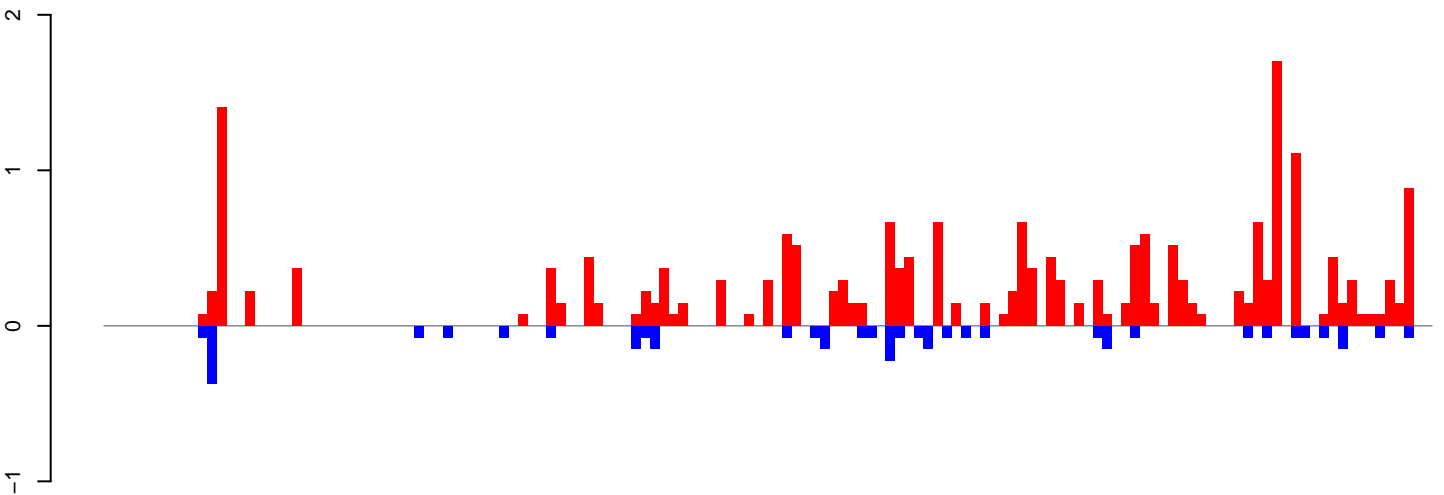


CuQuin_Testes_NL.rep



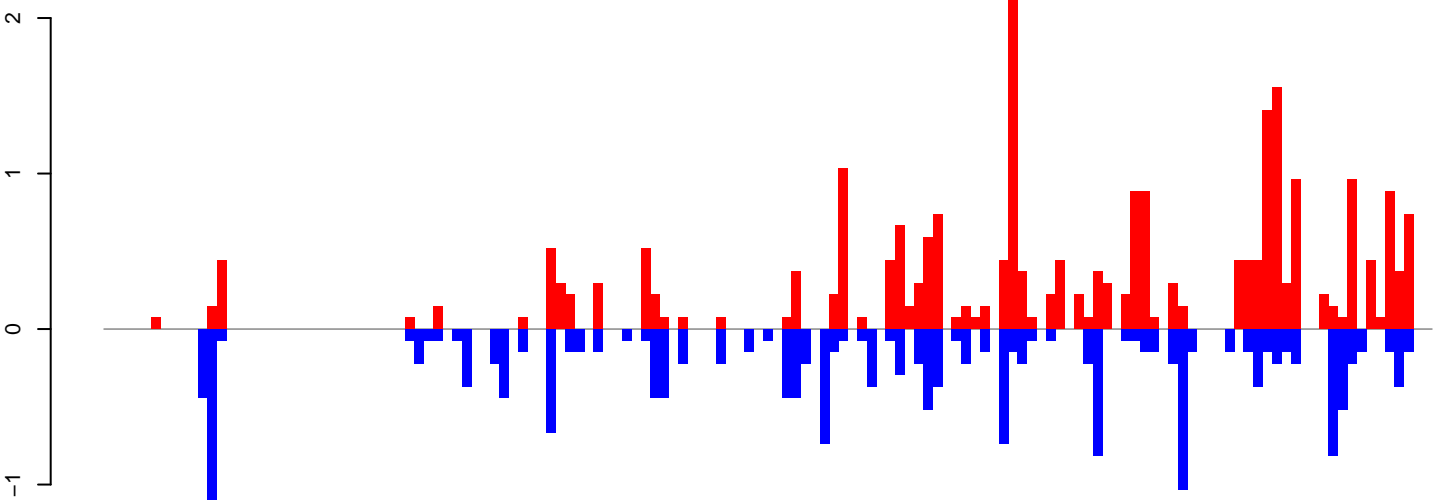
Window size=25, length=2300, TE@rnd-3_family-1022@LINE_I-Jockey@3301:1-2300

CuQuin_Testes_NL.18_23.rep



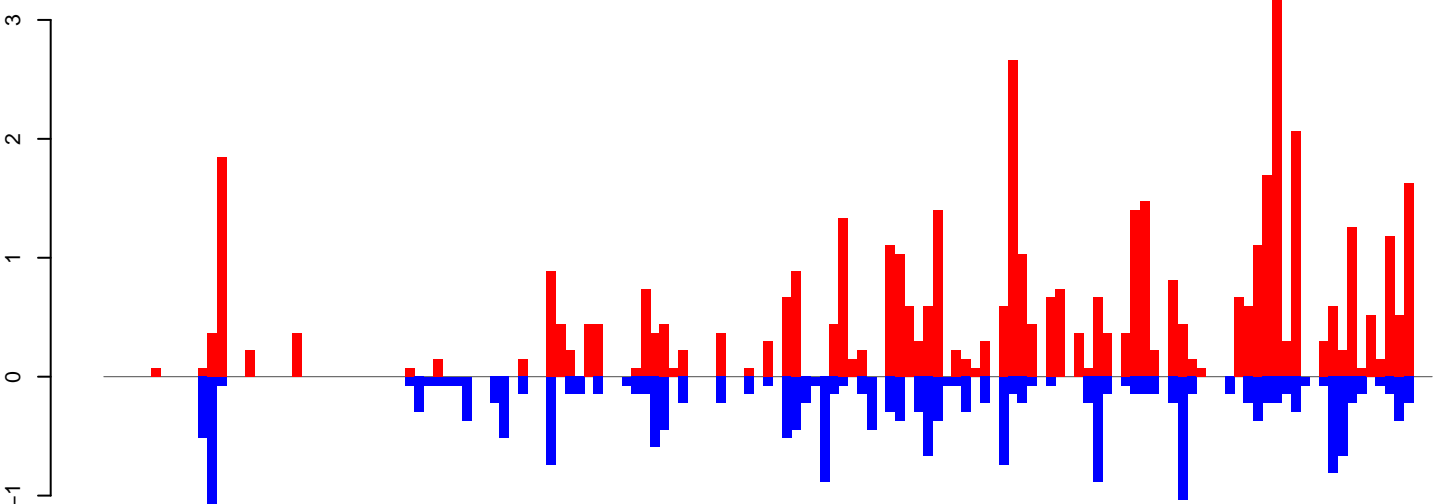
positive=21, negative=3, total=24

CuQuin_Testes_NL.24_35.rep



positive=25, negative=19, total=44

CuQuin_Testes_NL.rep

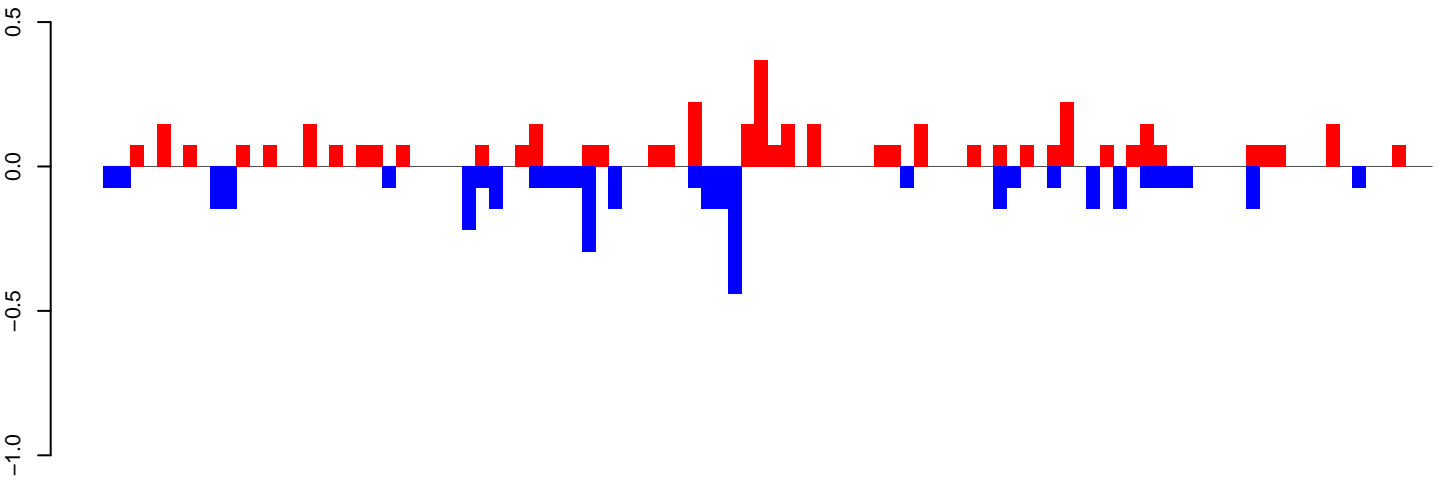


positive=47, negative=22, total=69

Window size=25, length=3513, TE@rnd-3_family-577@LINE_RTE-BovB@1330:1-3513

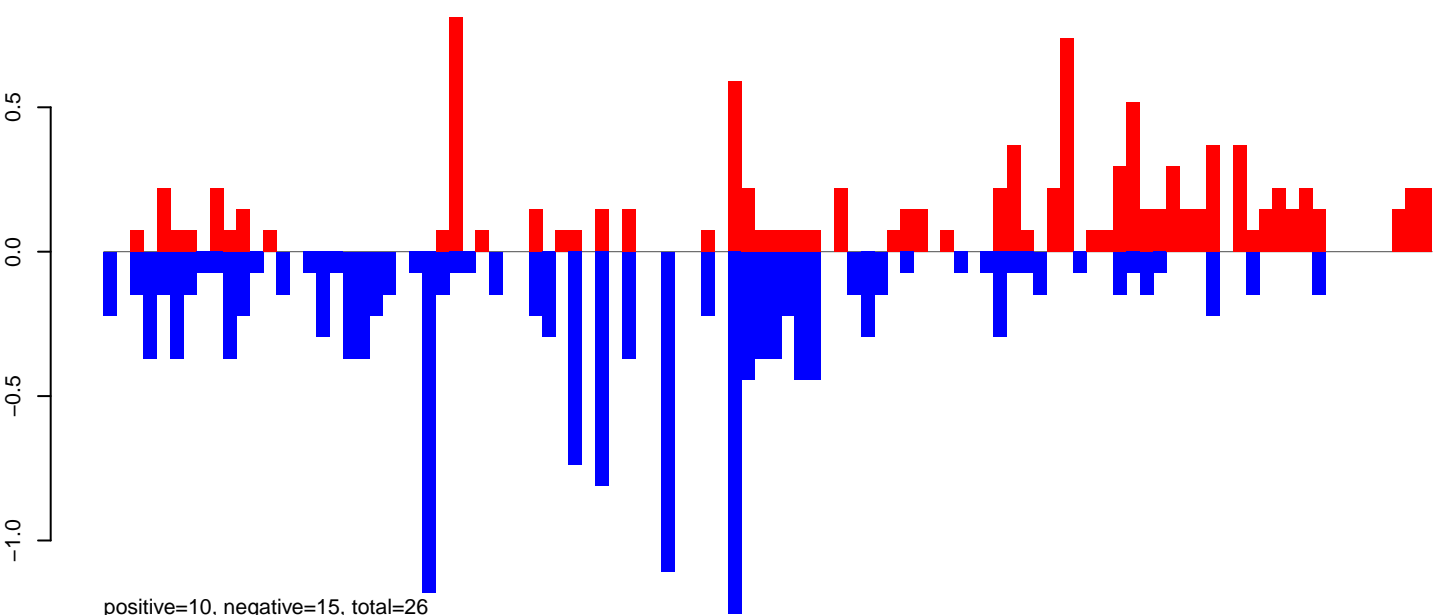
0 500 1000 1500 2000 2500 3000 3500

CuQuin_Testes_NL.18_23.rep



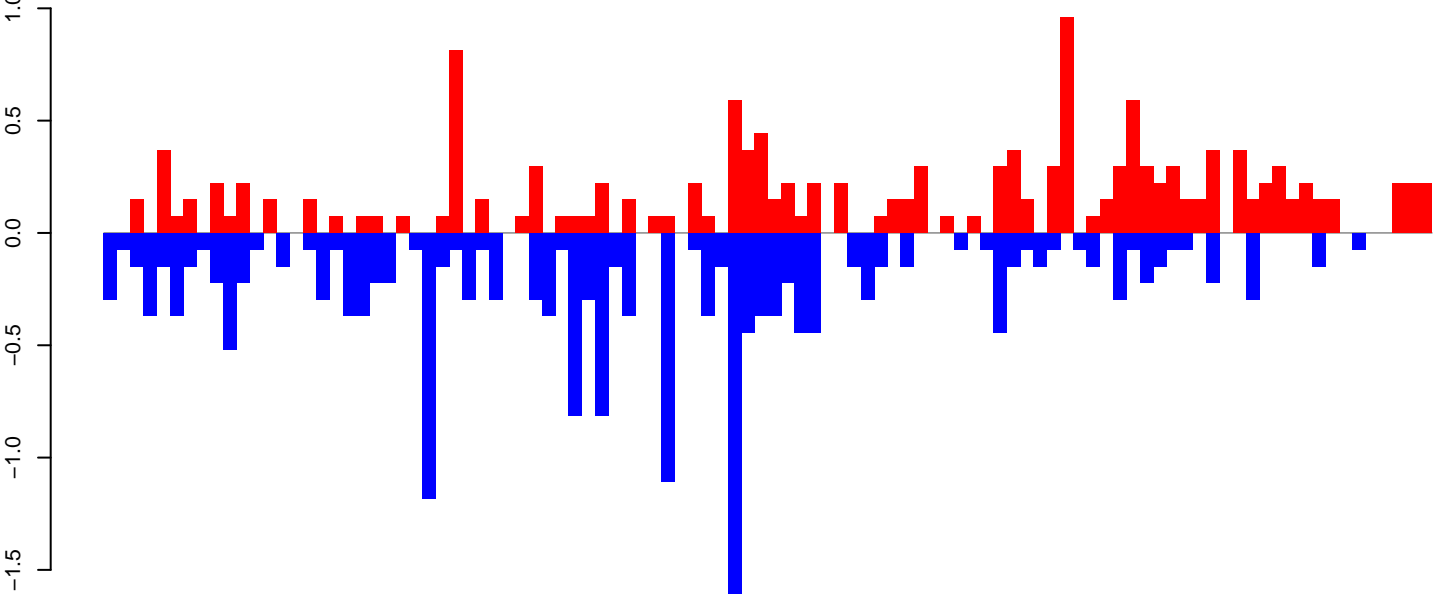
positive=4, negative=4, total=8

CuQuin_Testes_NL.24_35.rep



positive=10, negative=15, total=26

CuQuin_Testes_NL.rep

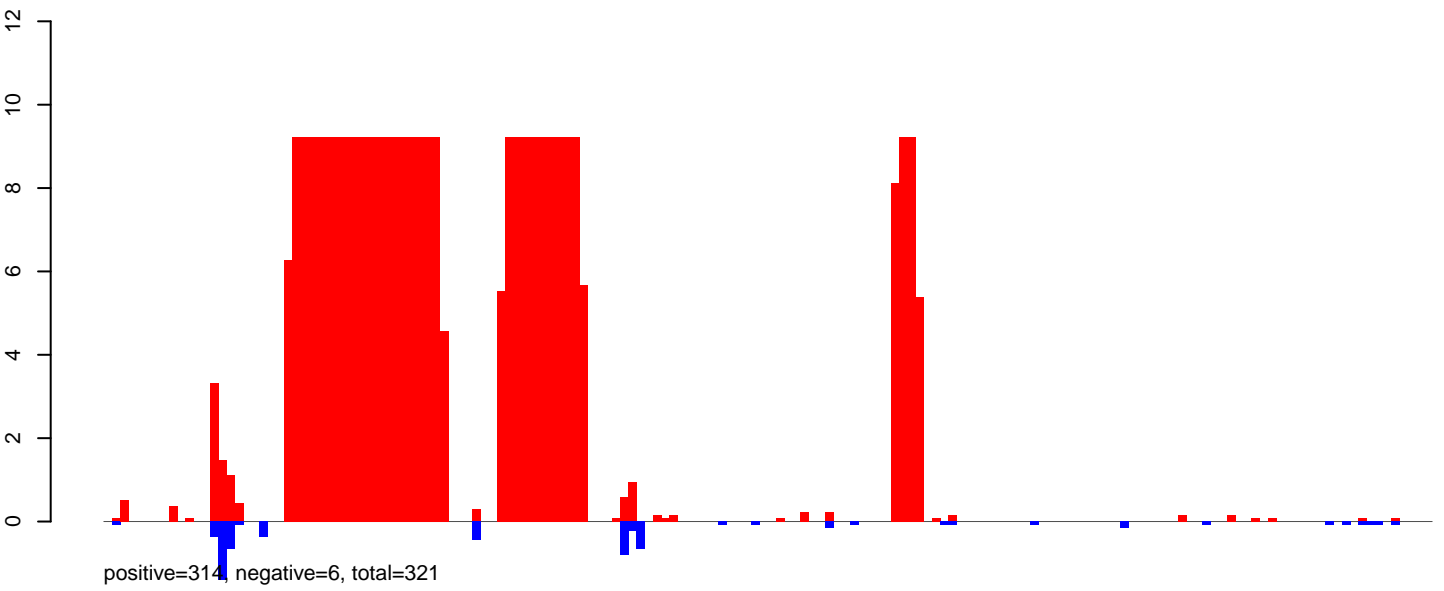


positive=14, negative=19, total=33

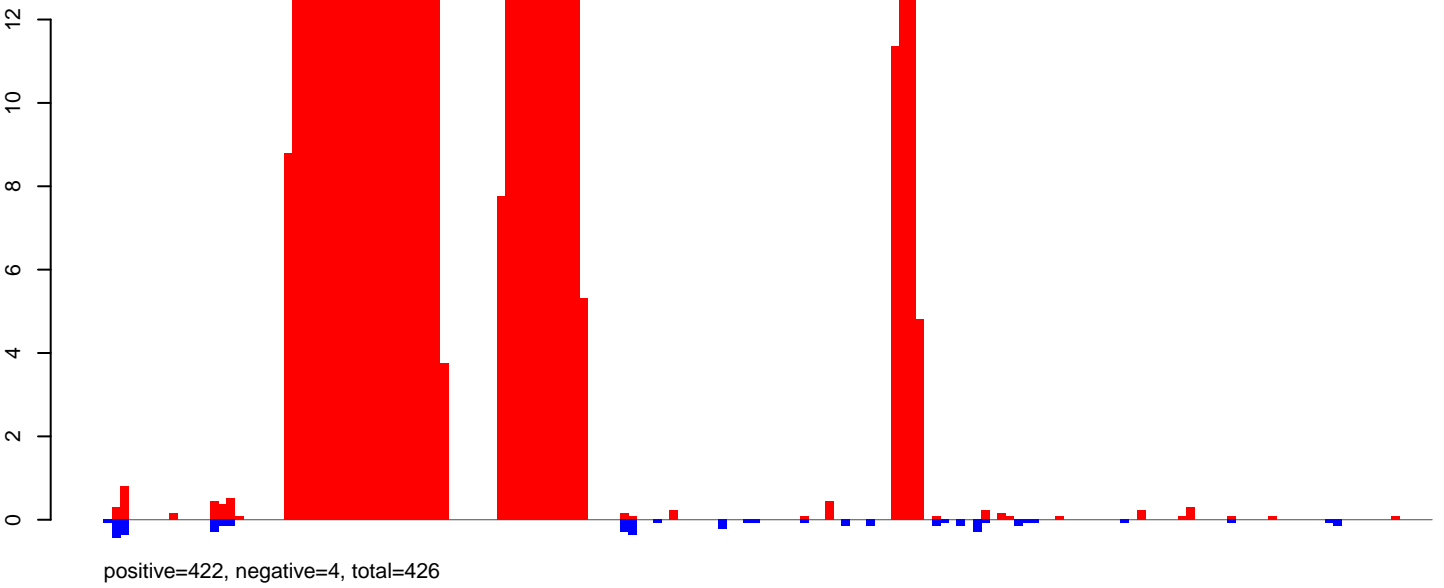
Window size=25, length=2495, TE@rnd-5_family-66@LINE_I@2559:1-2495

0 500 1000 1500 2000 2500

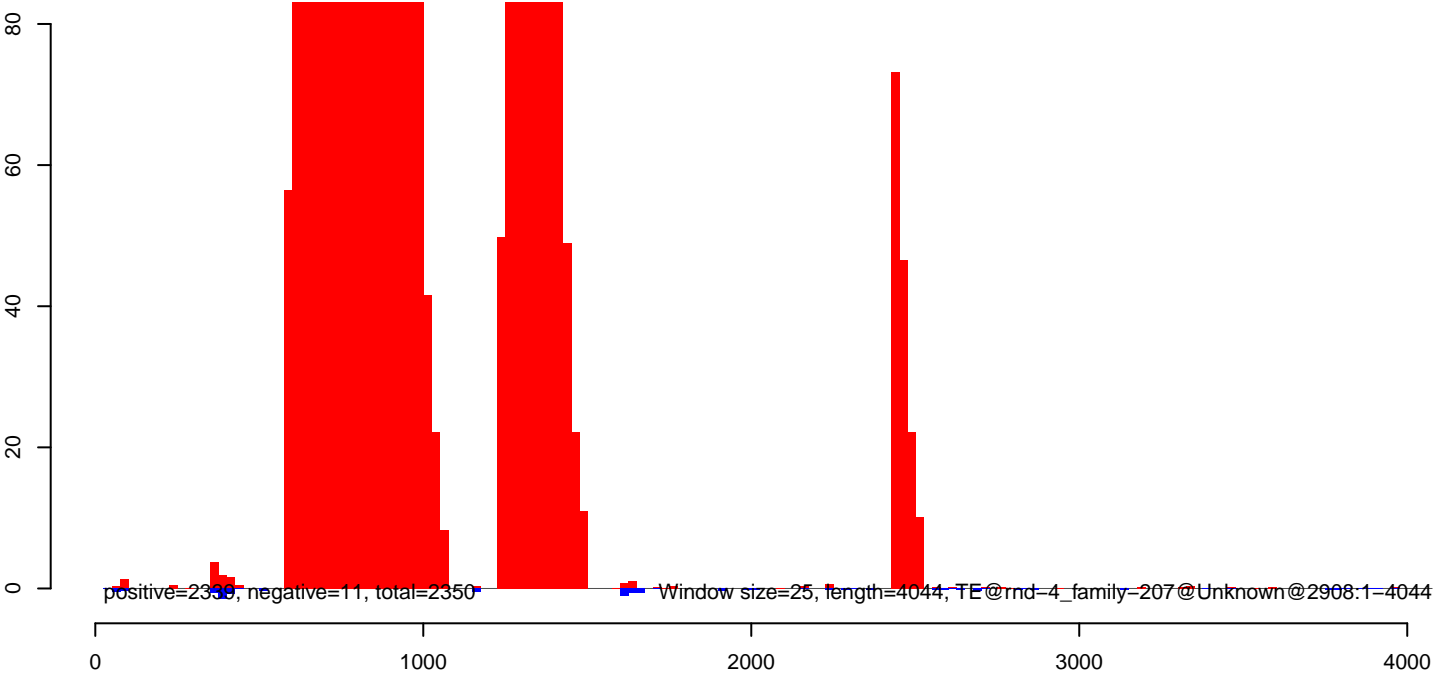
CuQuin_Testes_NL.18_23.rep



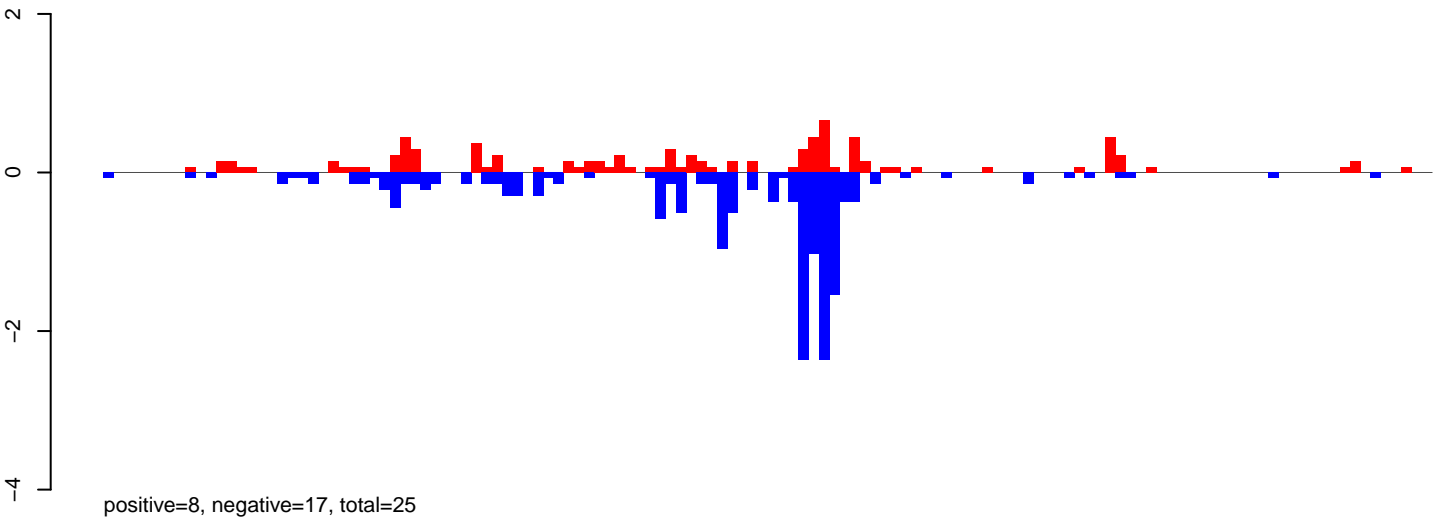
CuQuin_Testes_NL.24_35.rep



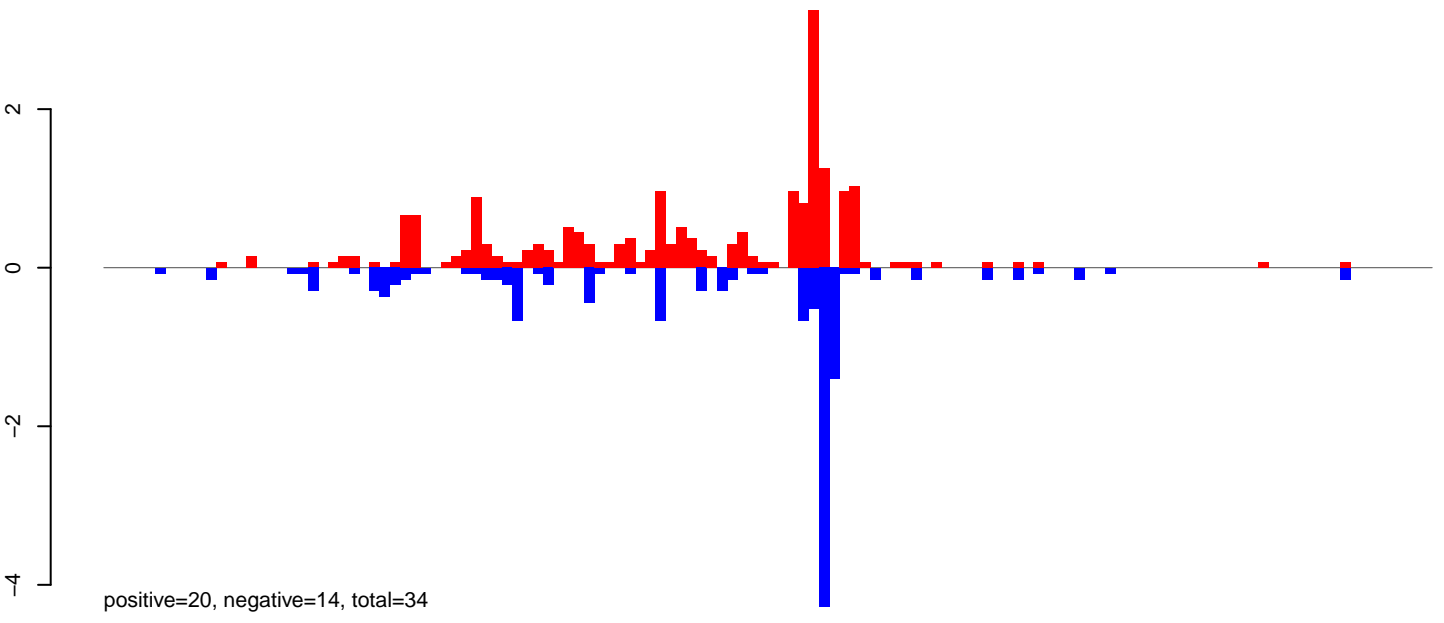
CuQuin_Testes_NL.rep



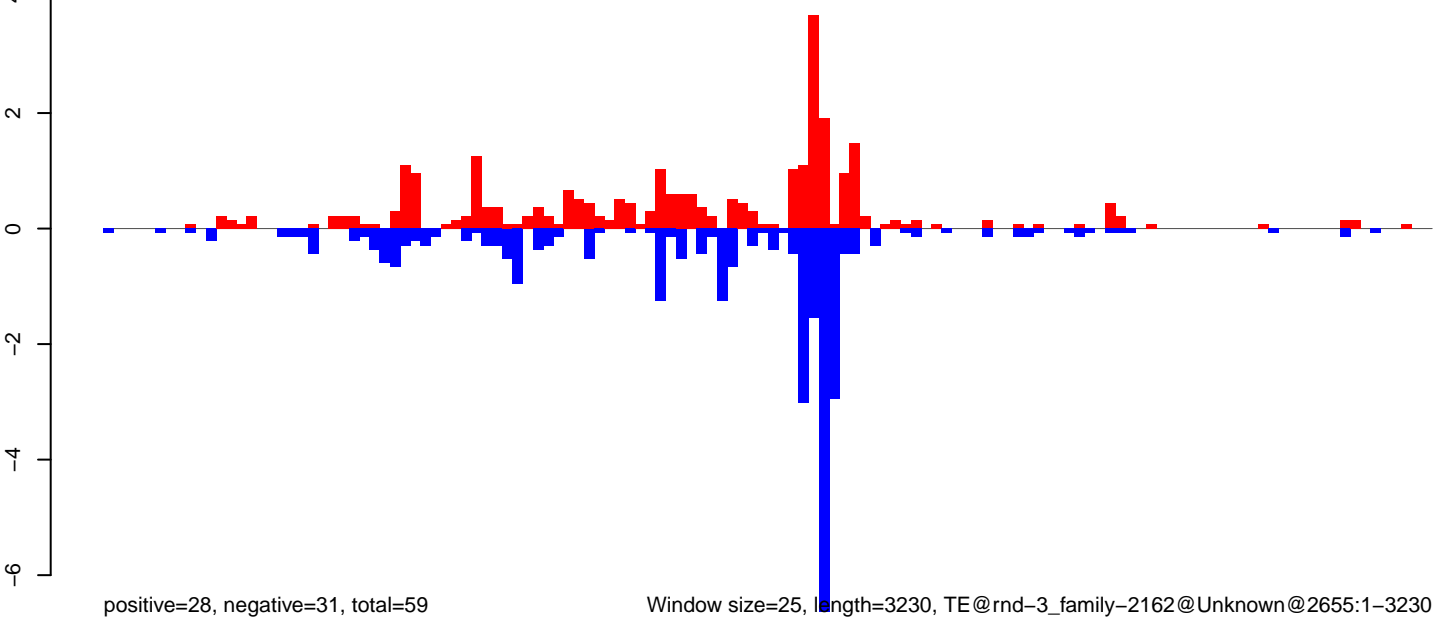
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

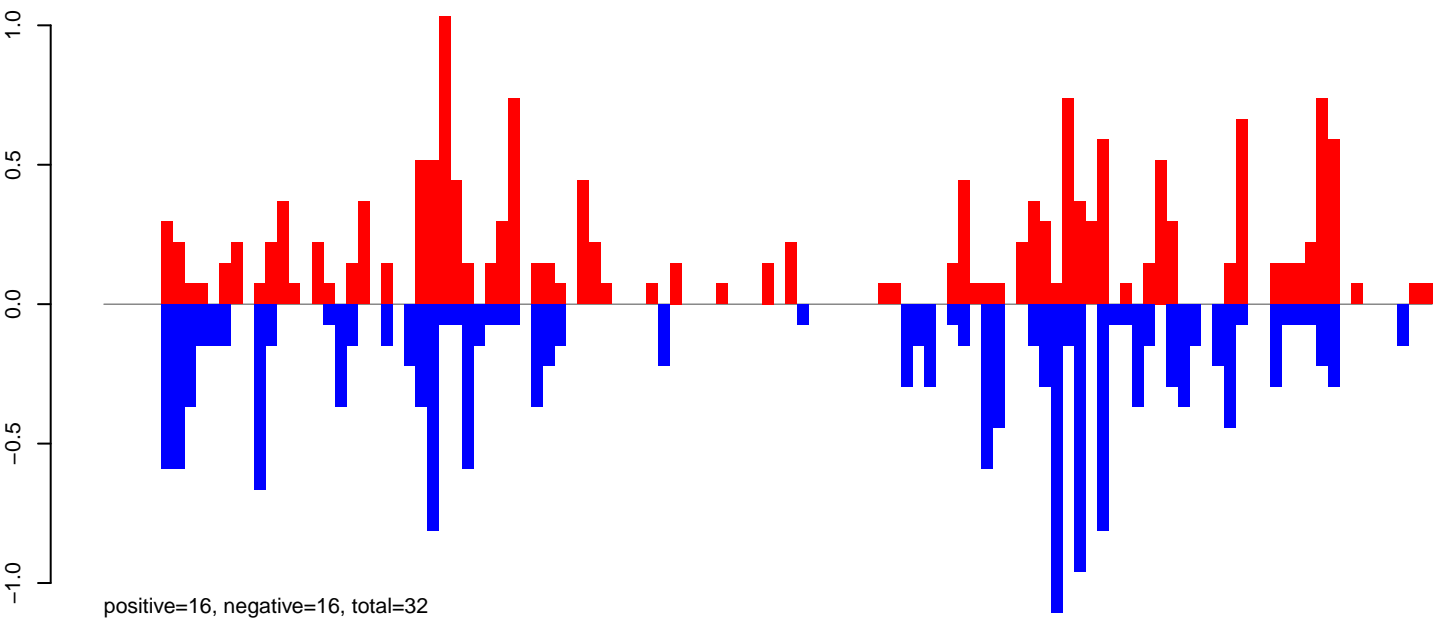


CuQuin_Testes_NL.rep

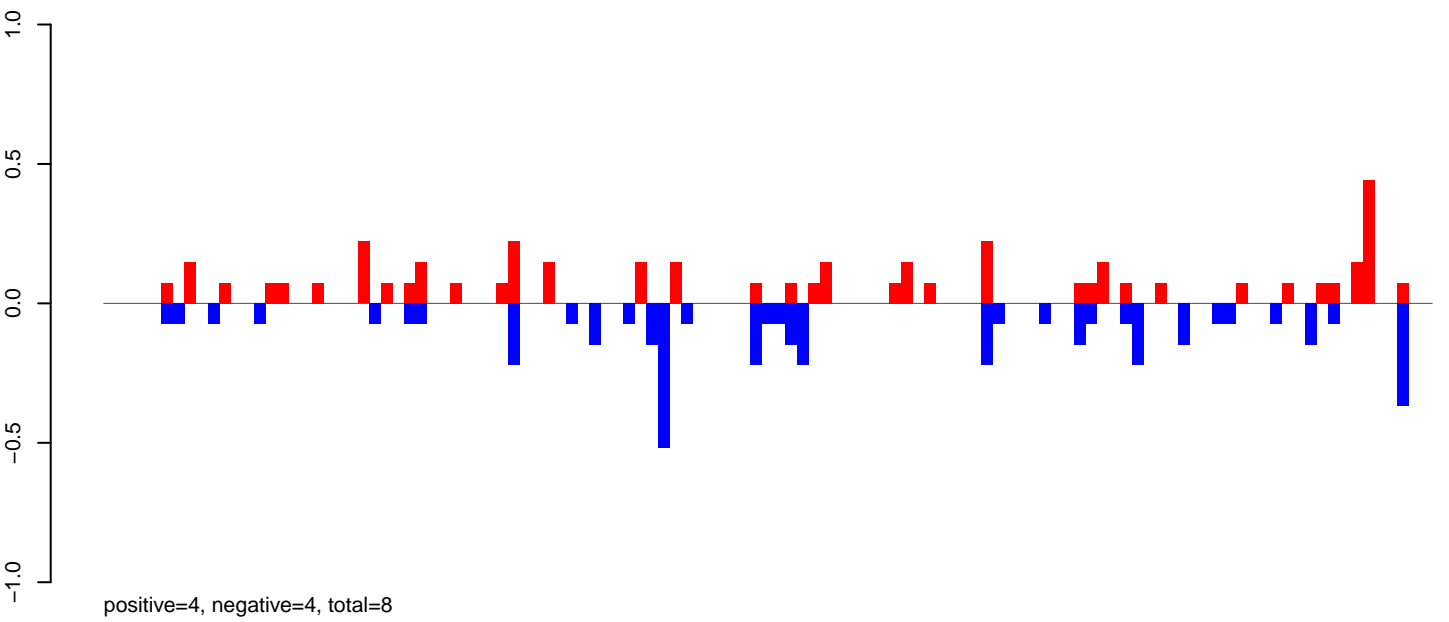


Window size=25, length=3230, TE@rnd-3_family-2162@Unknown@2655:1-3230

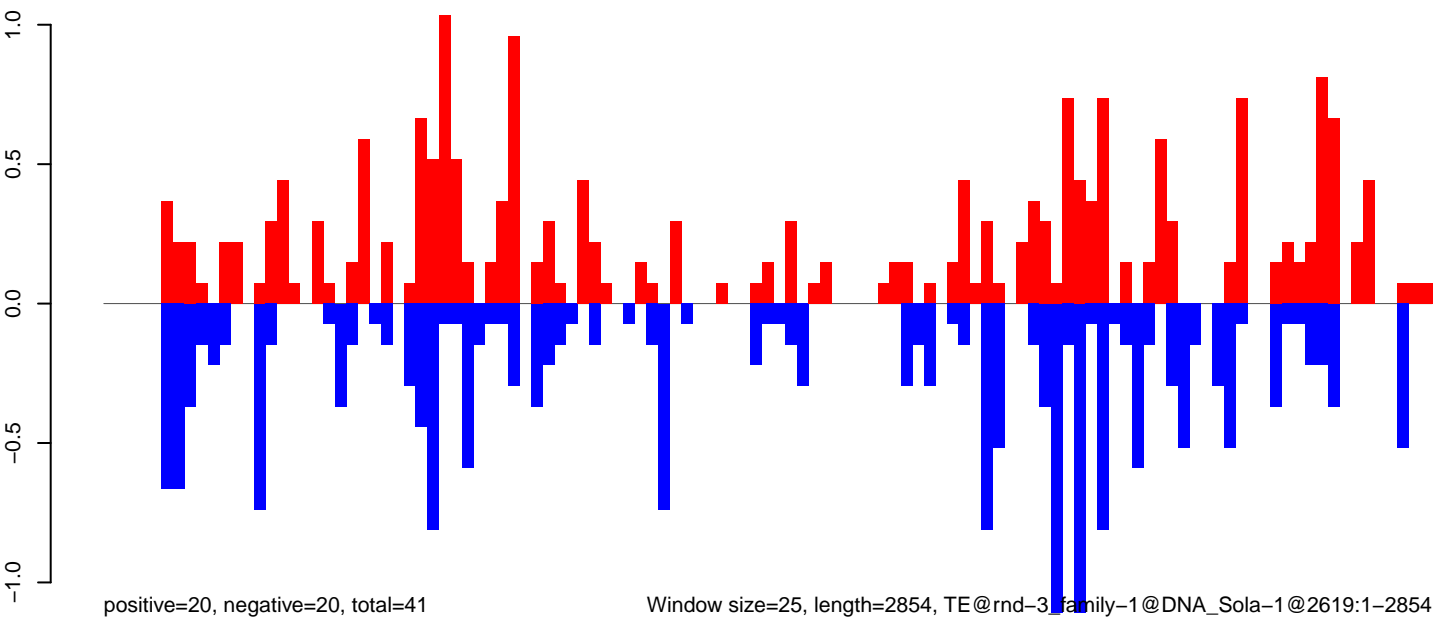
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

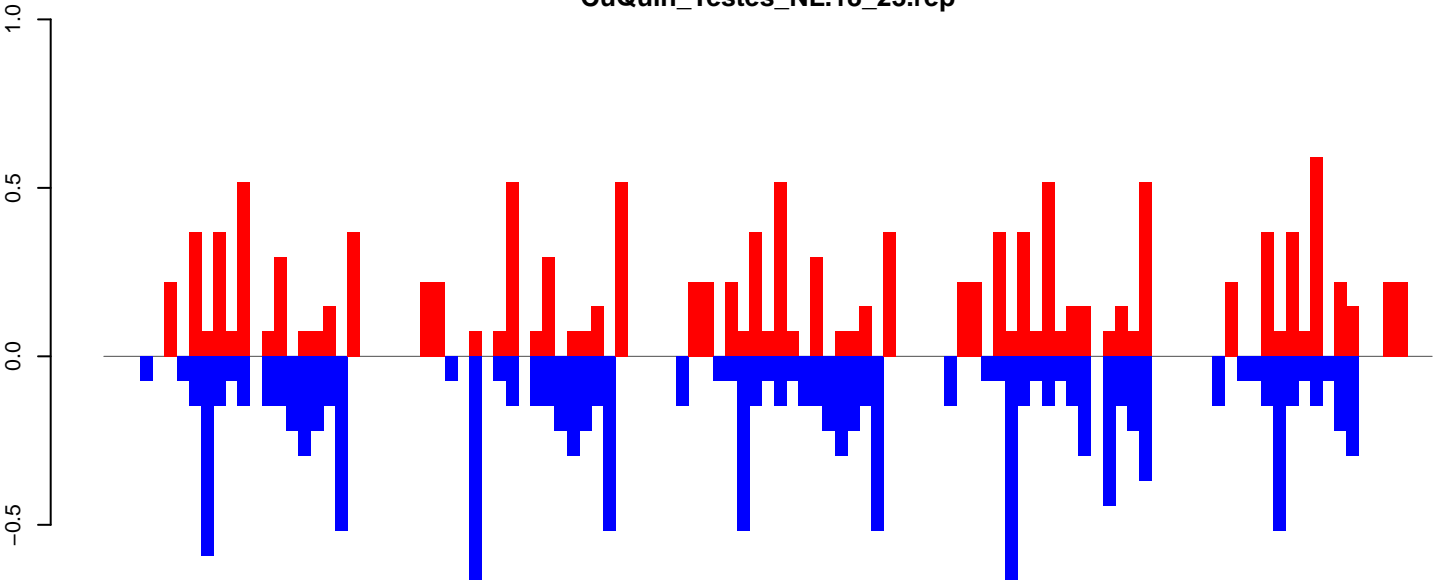


CuQuin_Testes_NL.rep



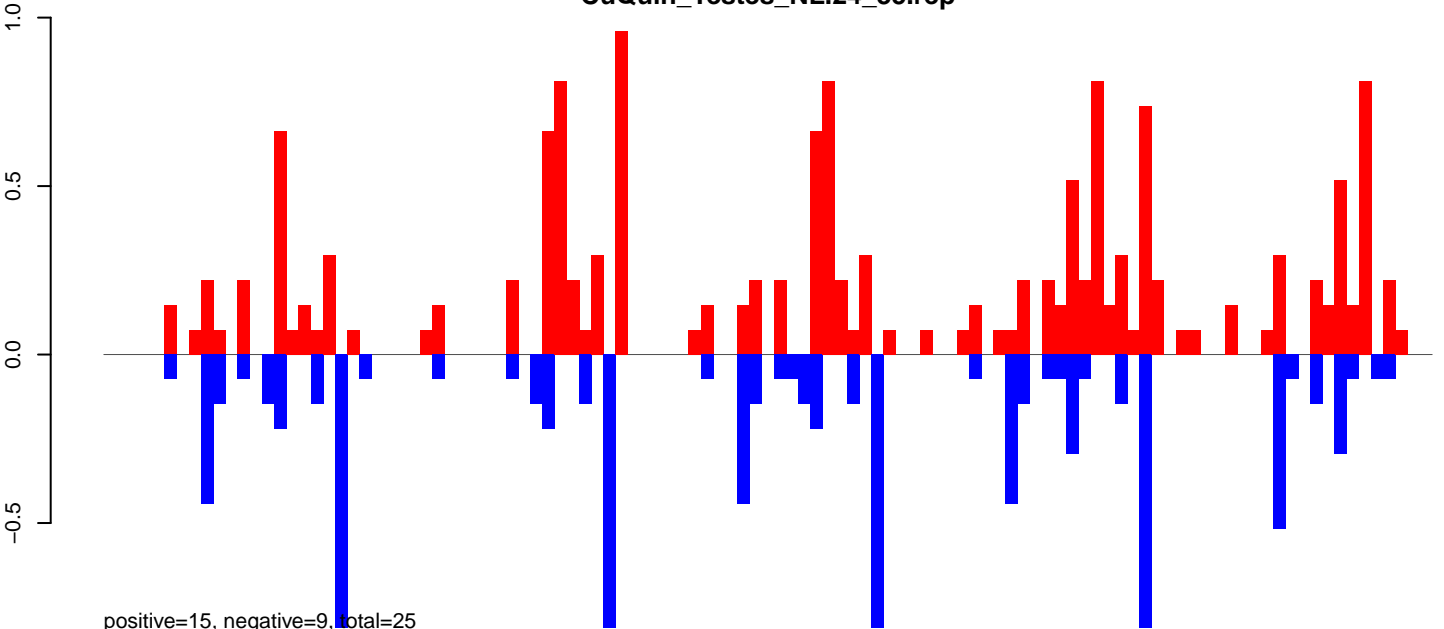
0 500 1000 1500 2000 2500 3000

CuQuin_Testes_NL.18_23.rep



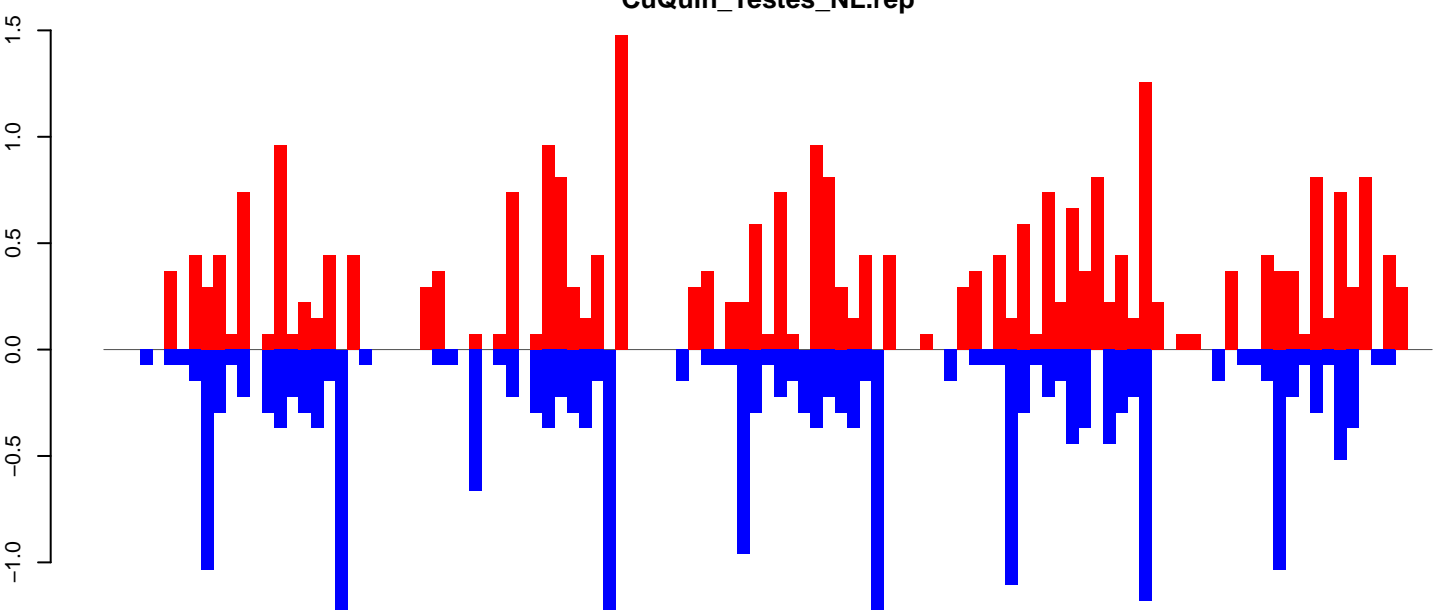
positive=13, negative=14, total=27

CuQuin_Testes_NL.24_35.rep



positive=15, negative=9, total=25

CuQuin_Testes_NL.rep

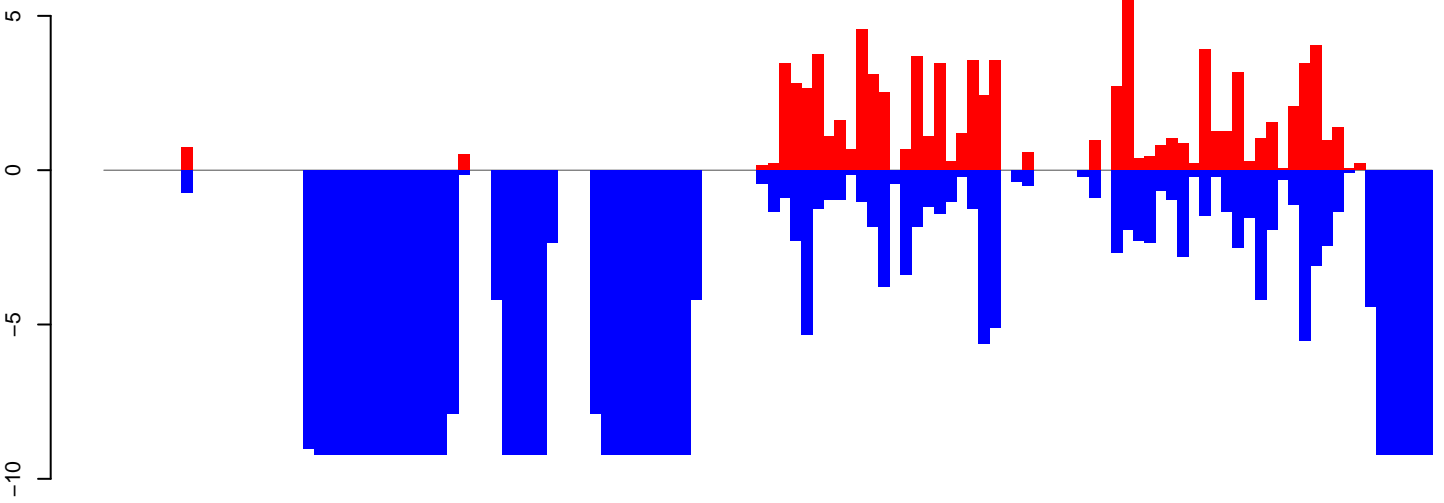


positive=29, negative=23, total=51

Window size=25, length=2707, TE@rnd-5_family-93@Unknown@3078:1-2707

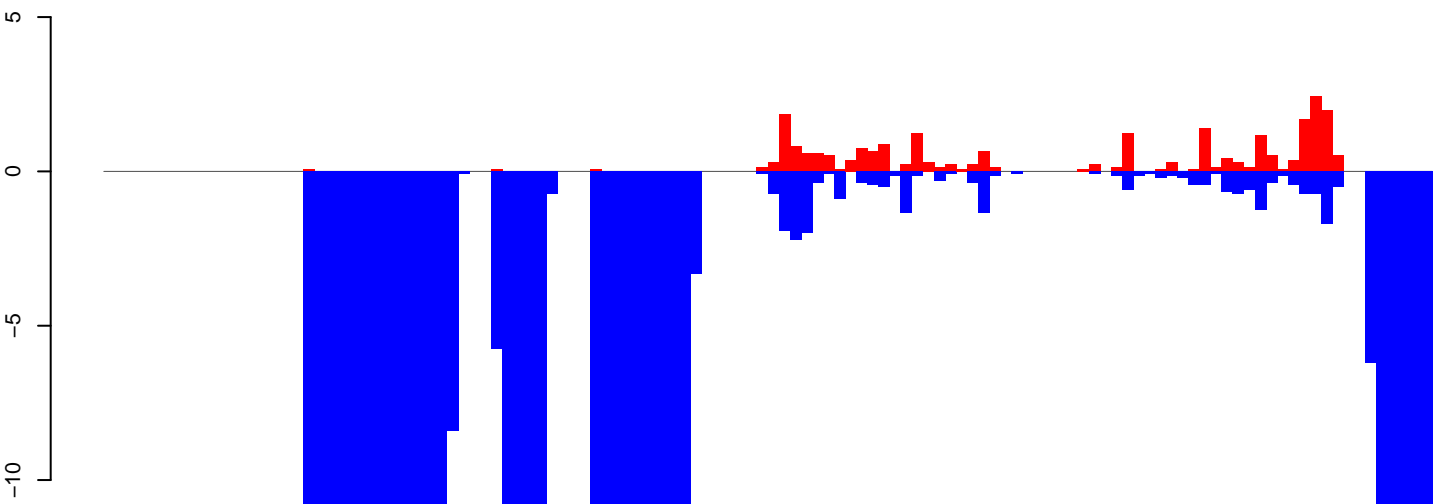
0 500 1000 1500 2000 2500

CuQuin_Testes_NL.18_23.rep



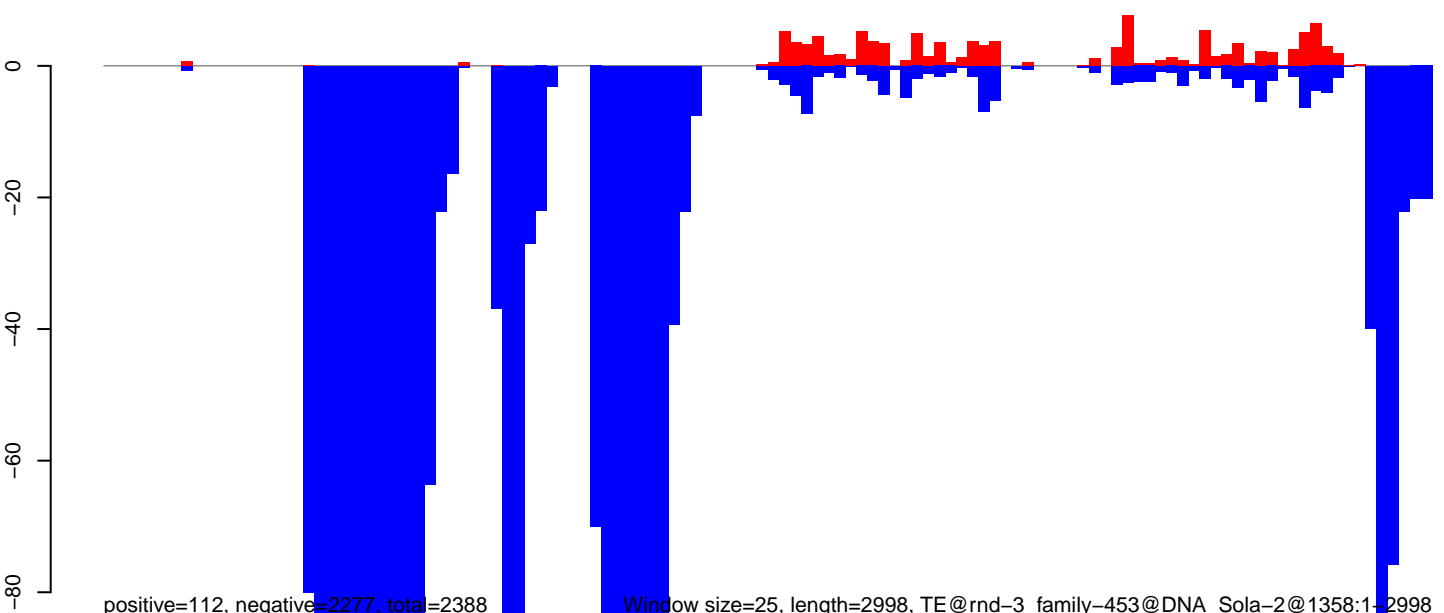
positive=87, negative=393, total=480

CuQuin_Testes_NL.24_35.rep



positive=24, negative=443, total=467

CuQuin_Testes_NL.rep

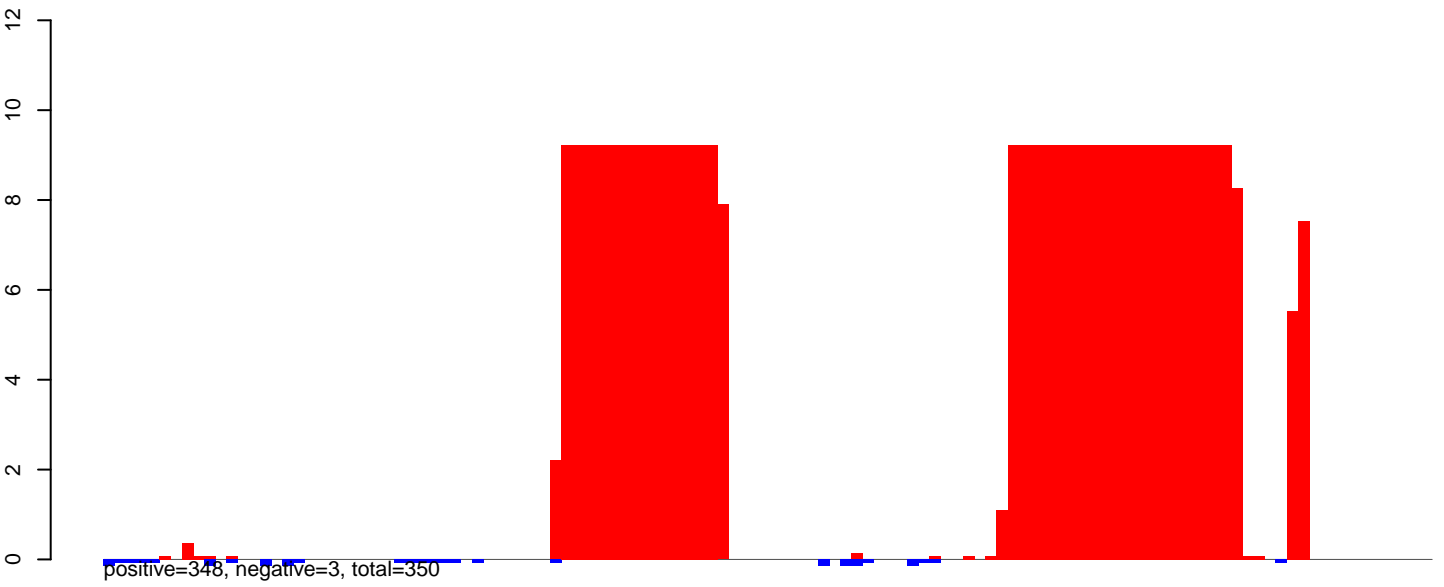


positive=112, negative=2277, total=2388

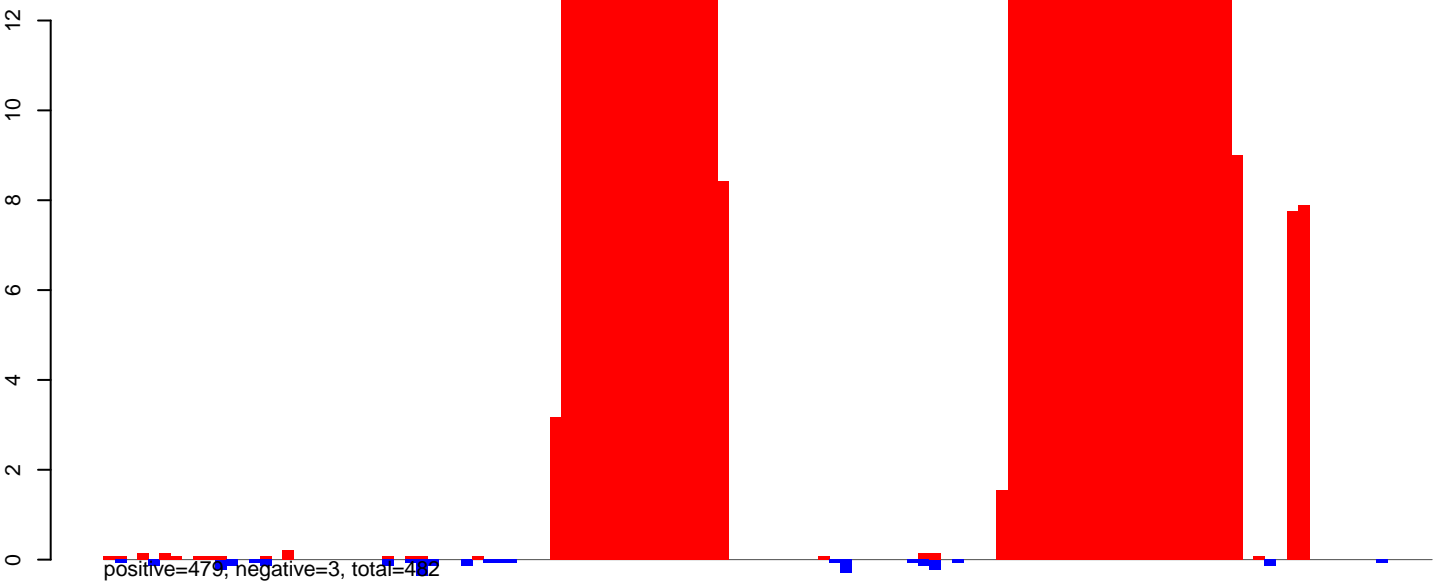
Window size=25, length=2998, TE@rnd-3_family-453@DNA_Sola-2@1358:1-2998

0 500 1000 1500 2000 2500 3000

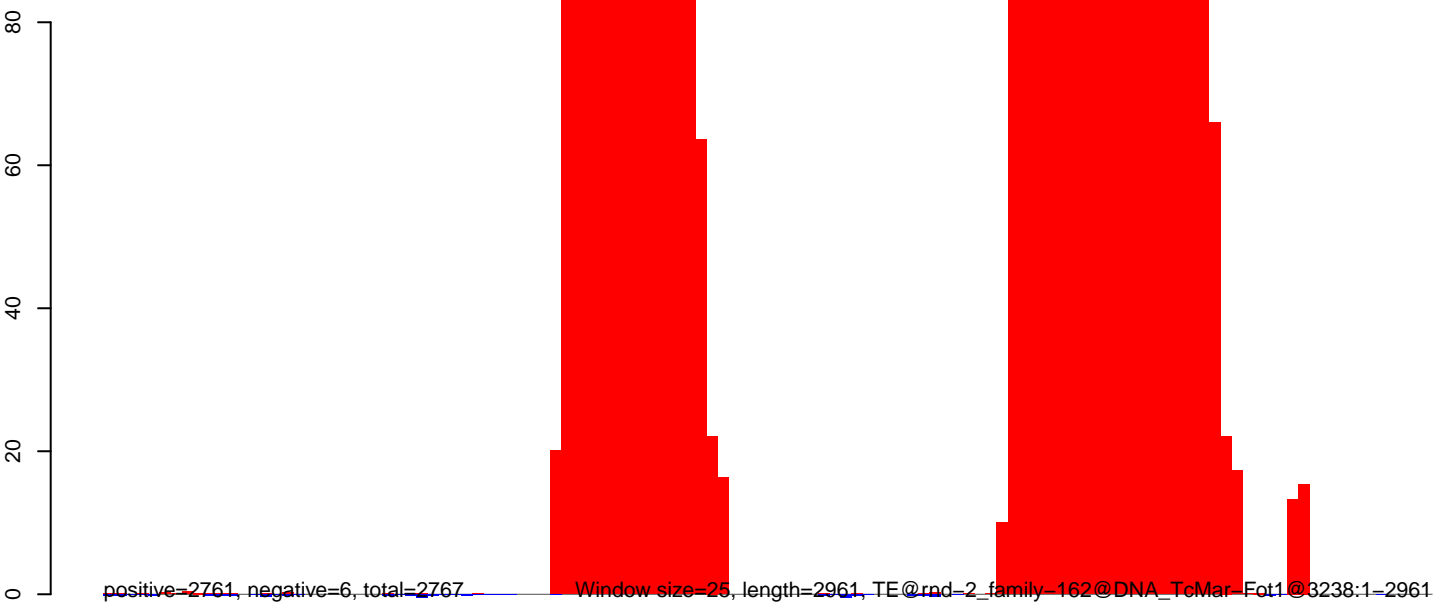
CuQuin_Testes_NL.18_23.rep



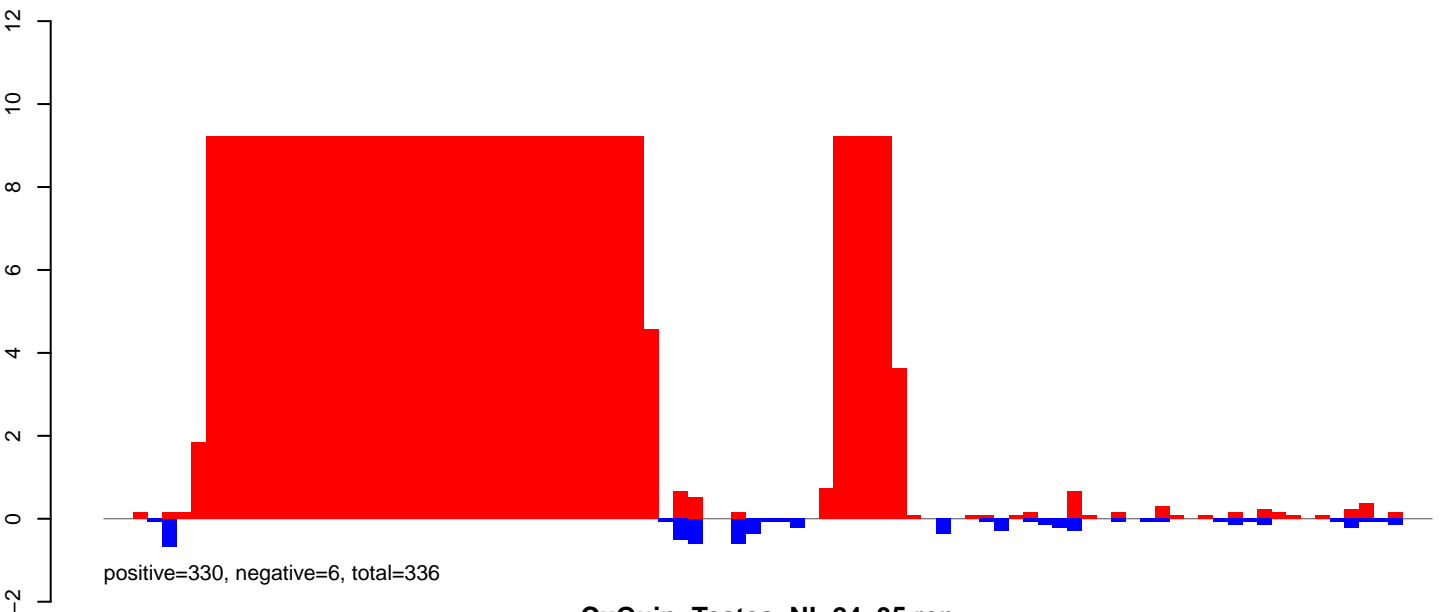
CuQuin_Testes_NL.24_35.rep



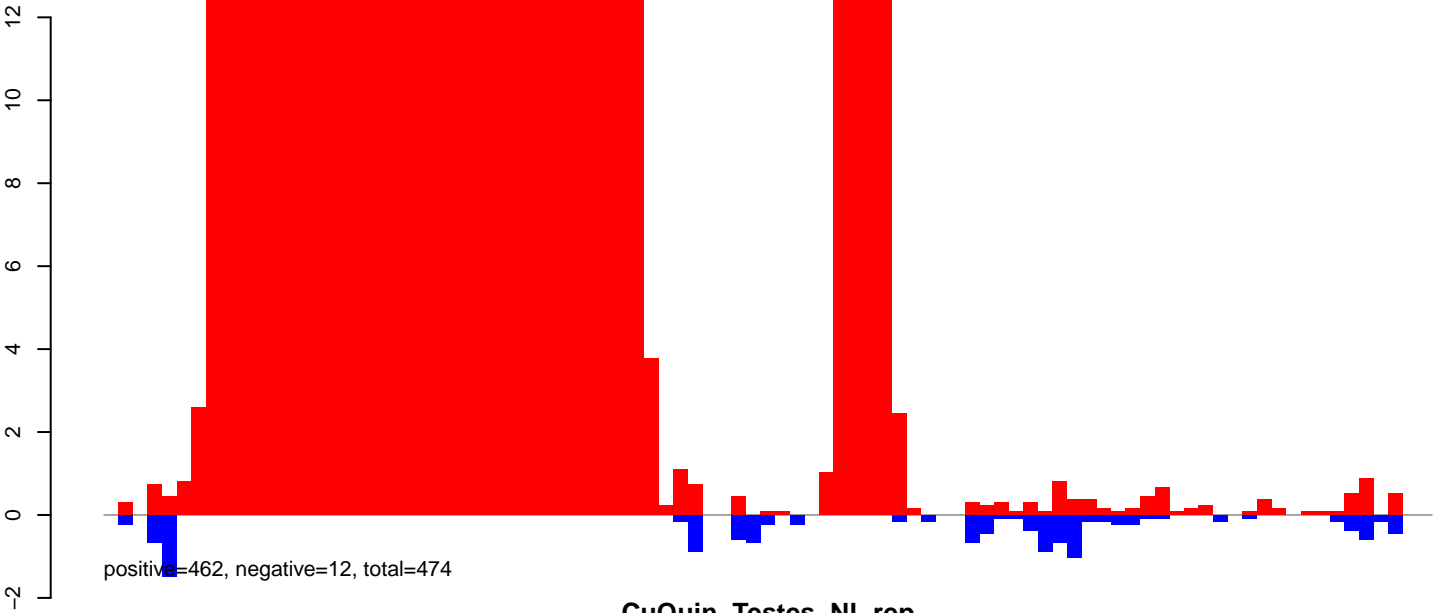
CuQuin_Testes_NL.rep



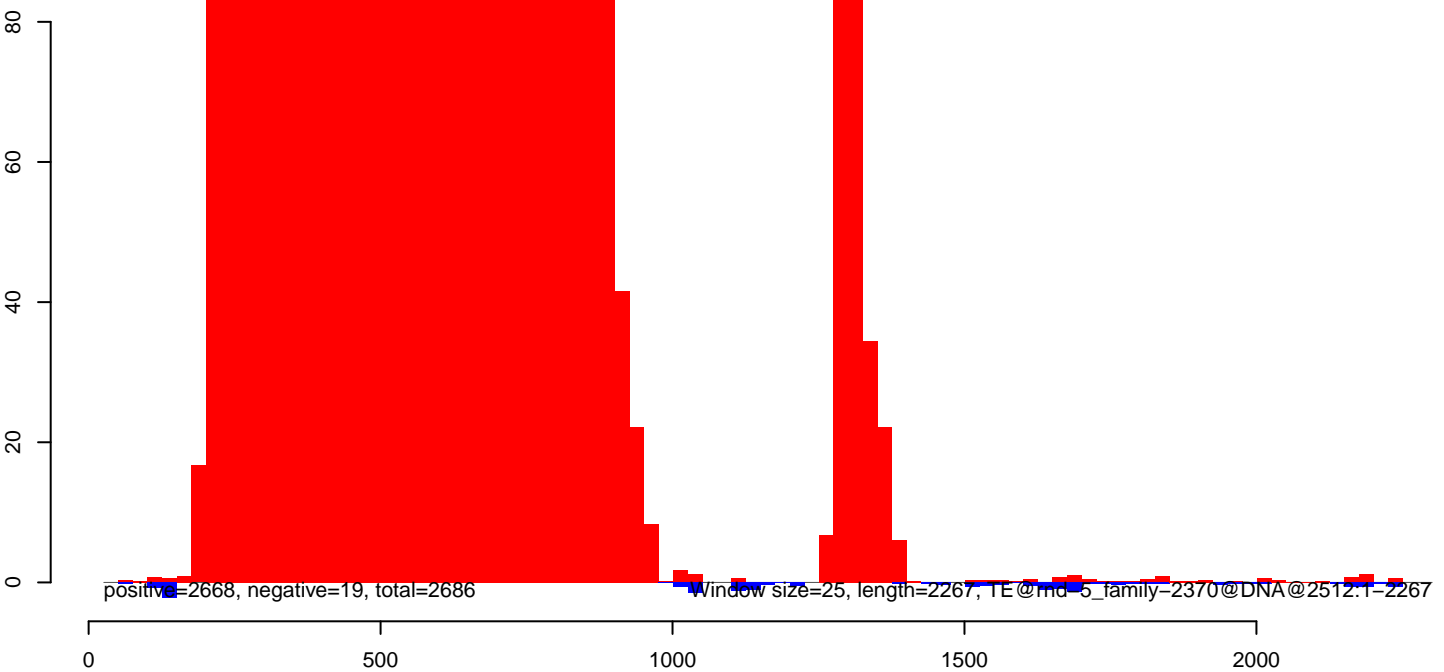
CuQuin_Testes_NL.18_23.rep



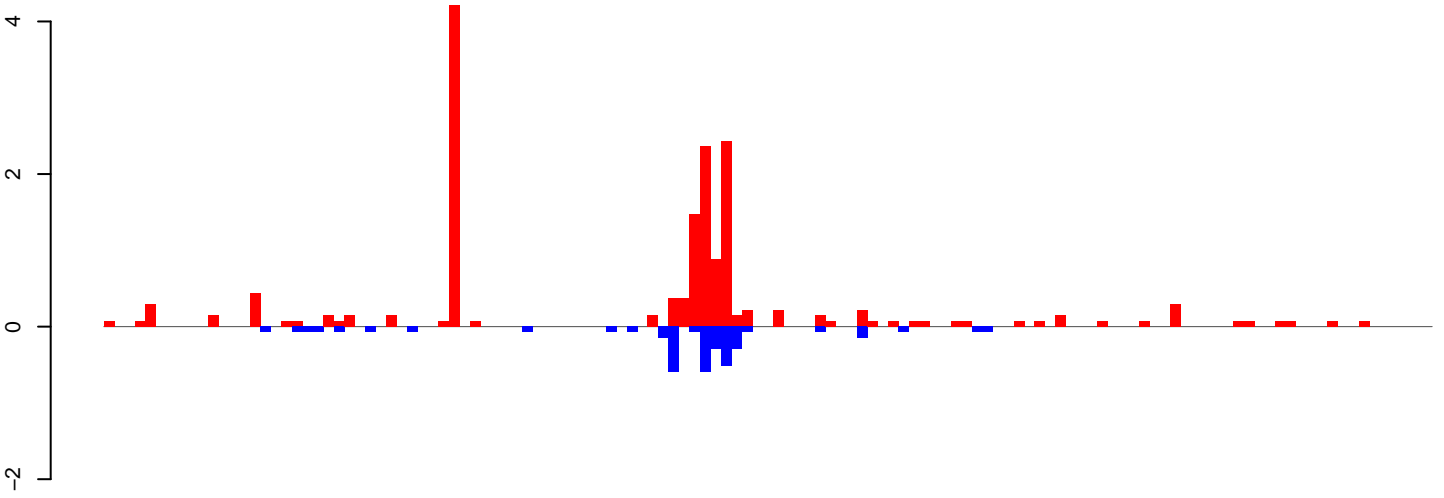
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

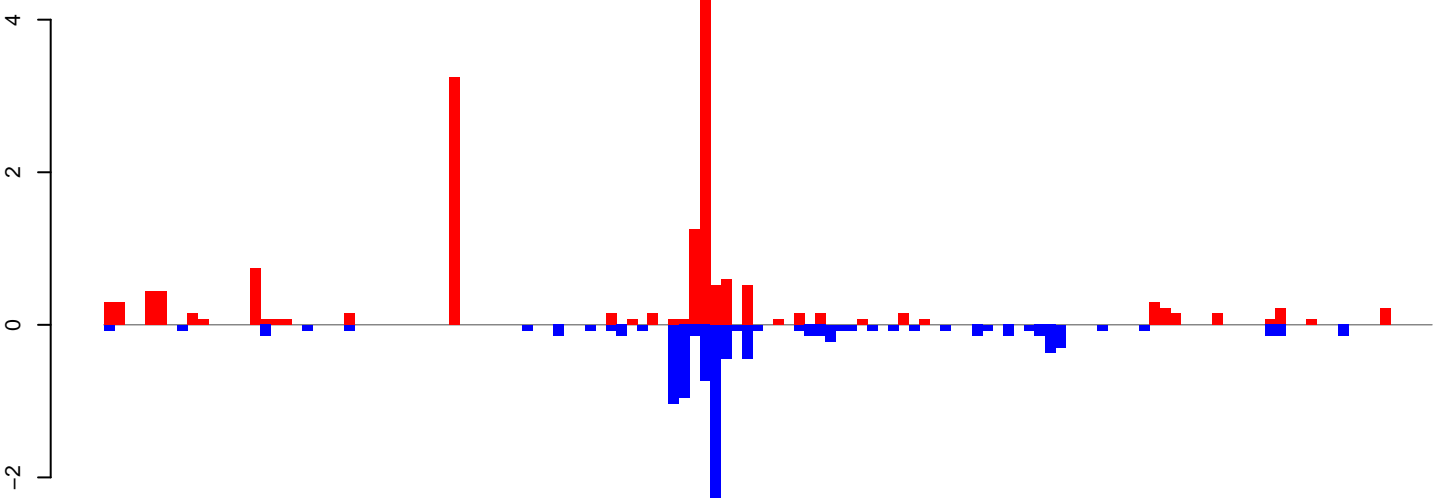


CuQuin_Testes_NL.18_23.rep



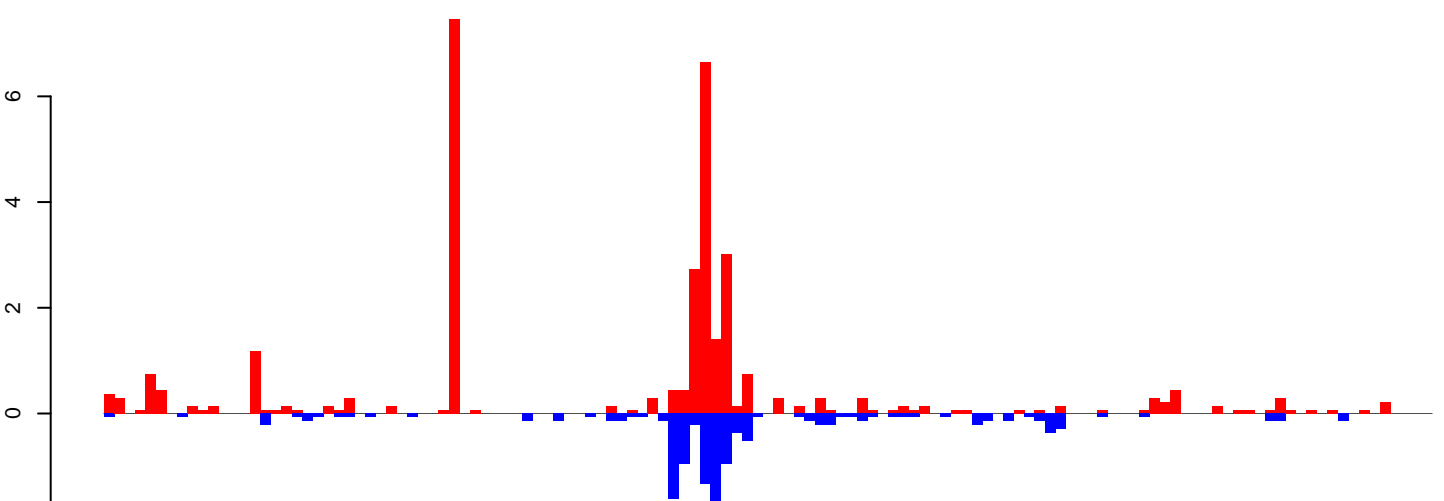
positive=17, negative=4, total=21

CuQuin_Testes_NL.24_35.rep



positive=16, negative=11, total=27

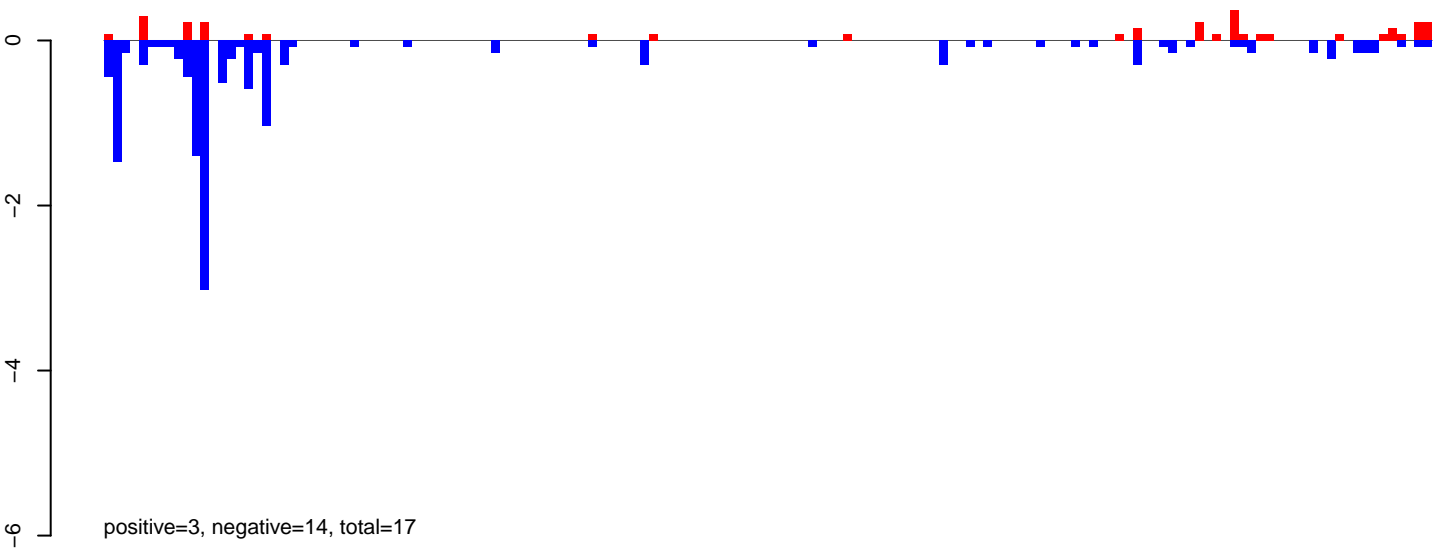
CuQuin_Testes_NL.rep



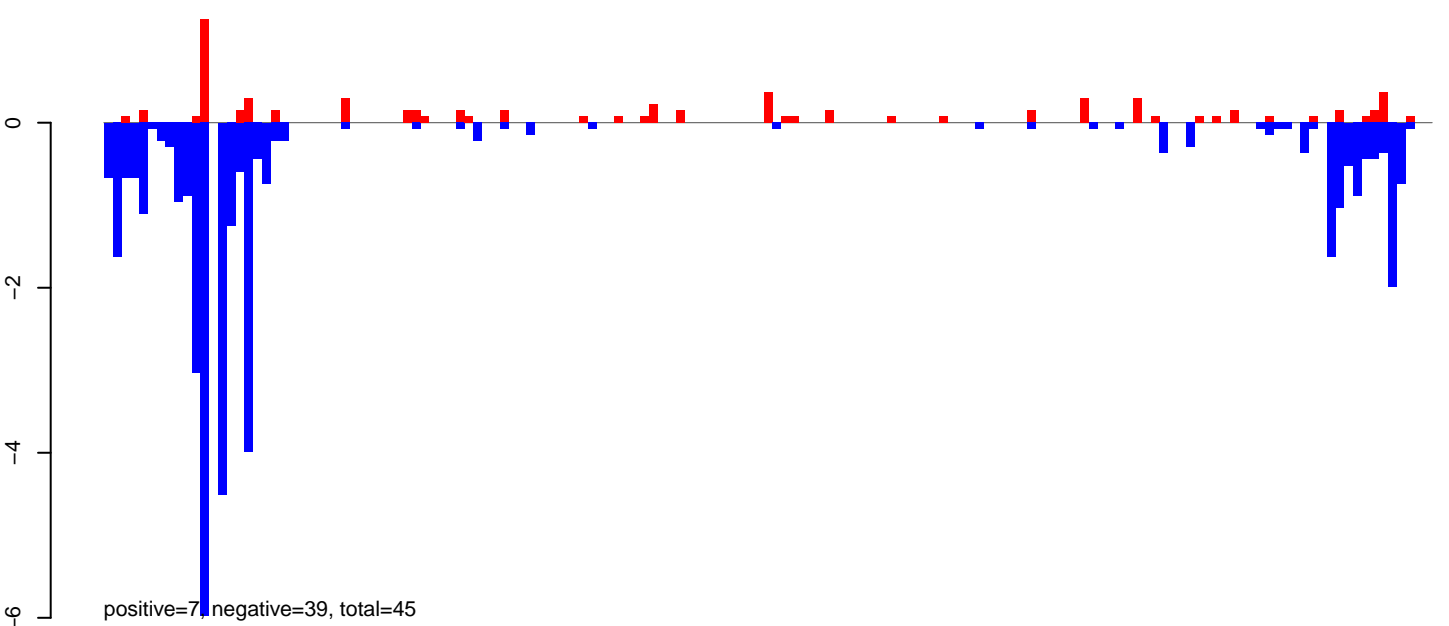
positive=33, negative=15, total=48

Window size=25, length=3169, TE@rnd-5_family-317@LTR_Copia@2420:1-3169

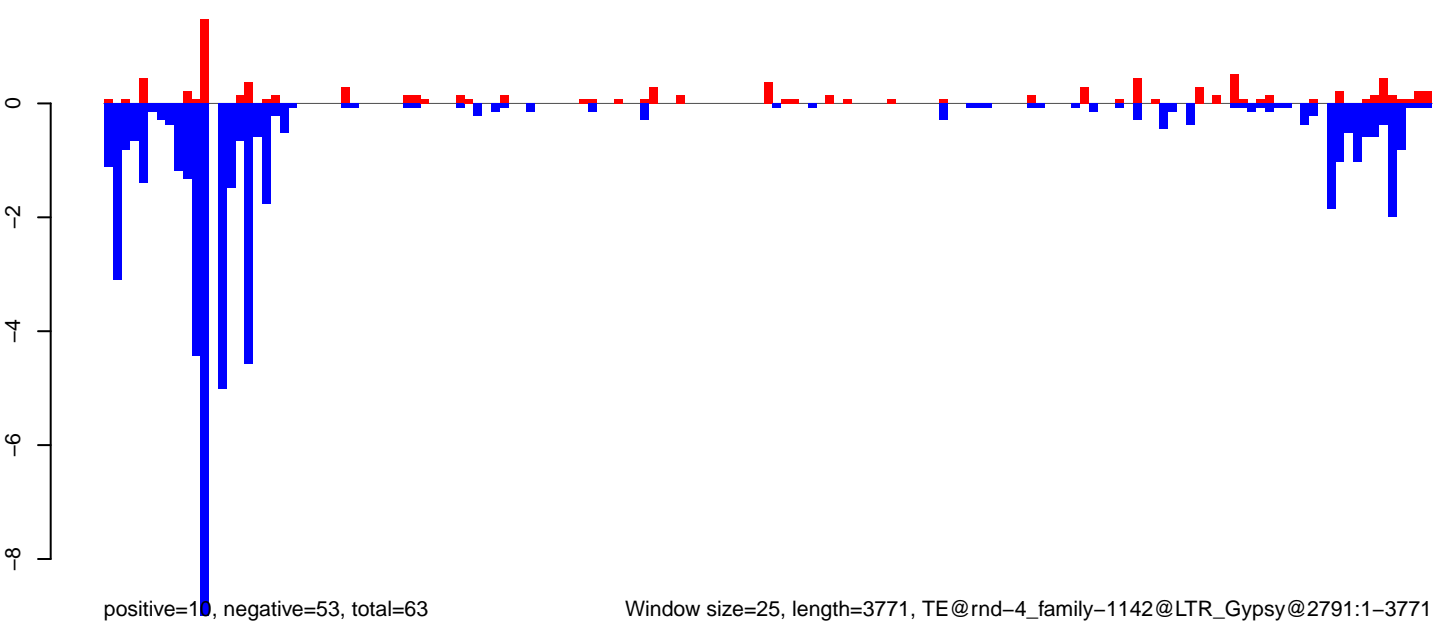
CuQuin_Testes_NL.18_23.rep



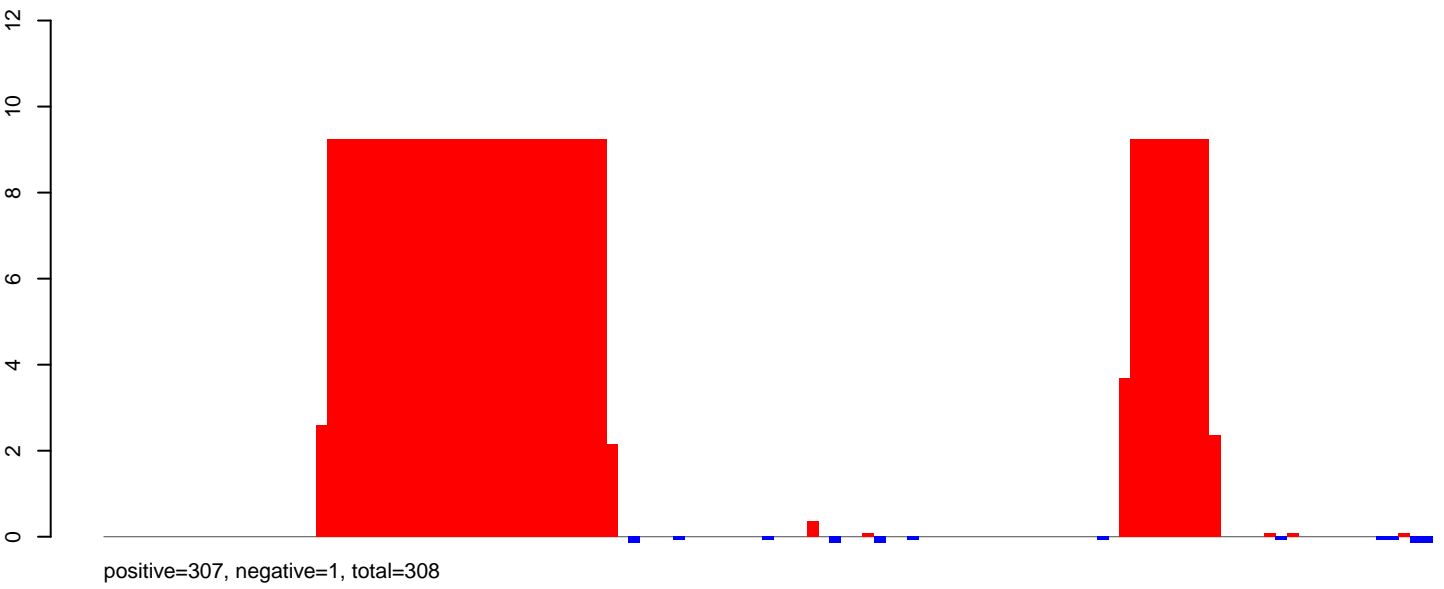
CuQuin_Testes_NL.24_35.rep



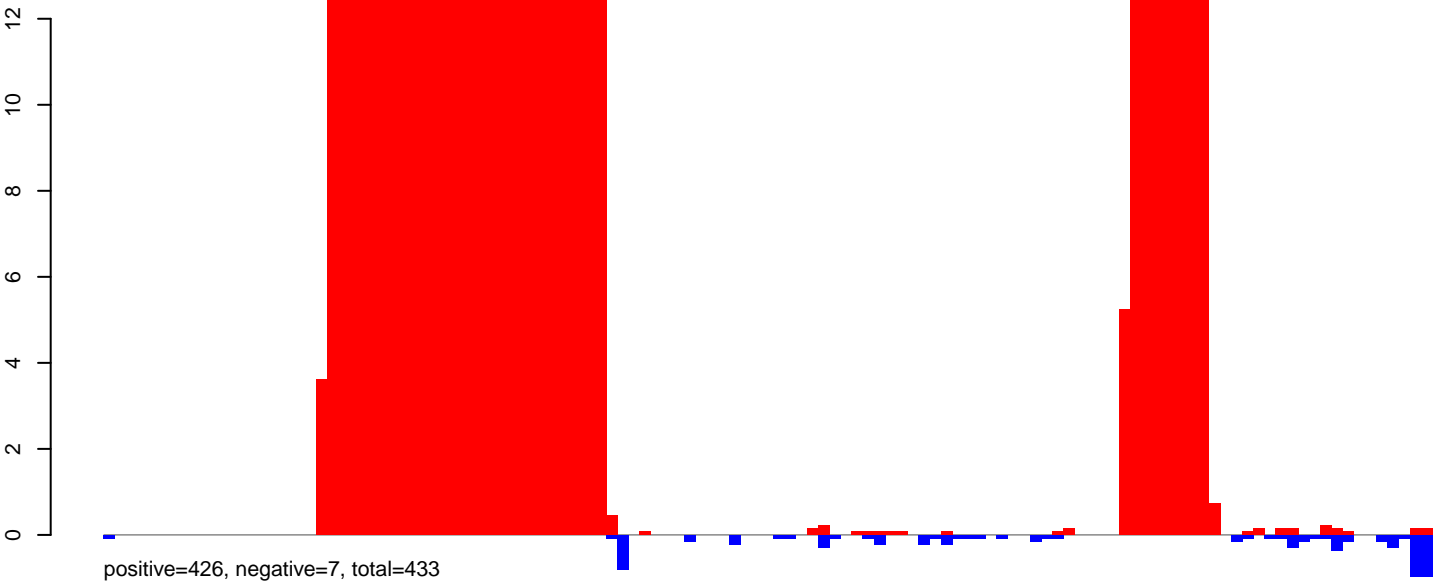
CuQuin_Testes_NL.rep



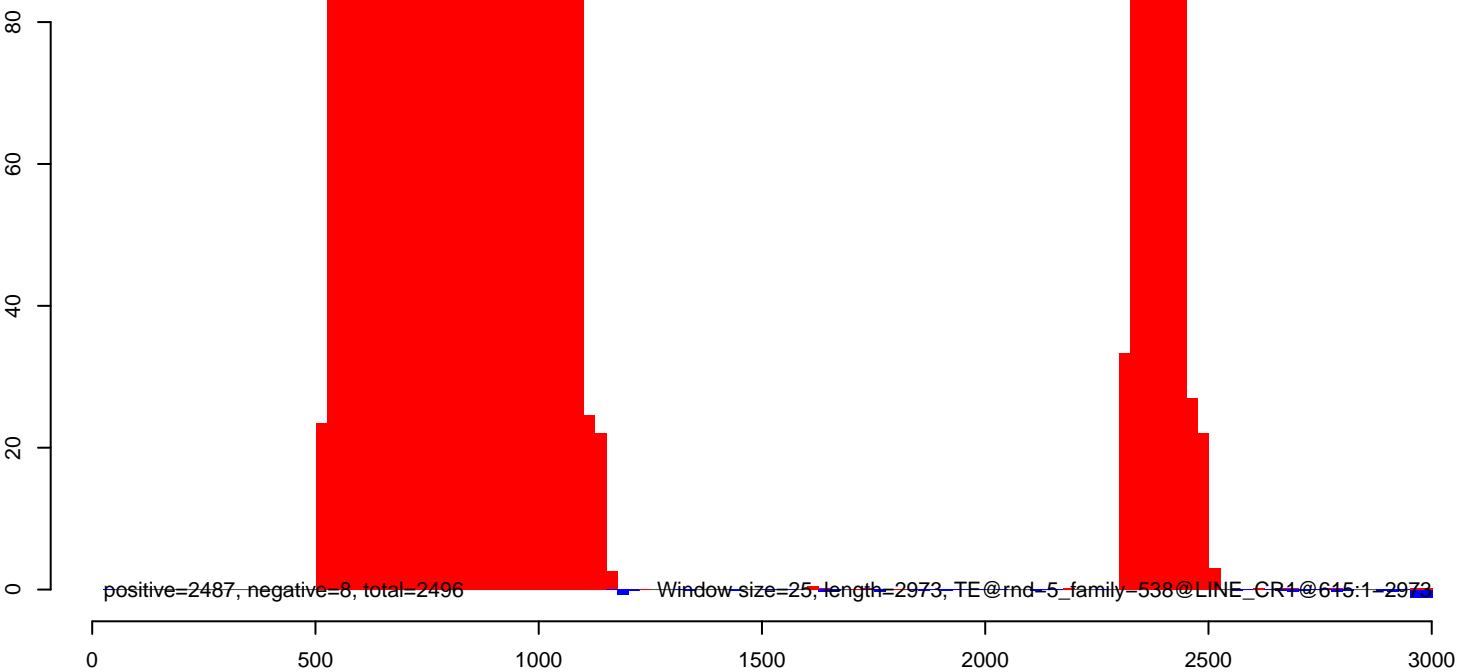
CuQuin_Testes_NL.18_23.rep



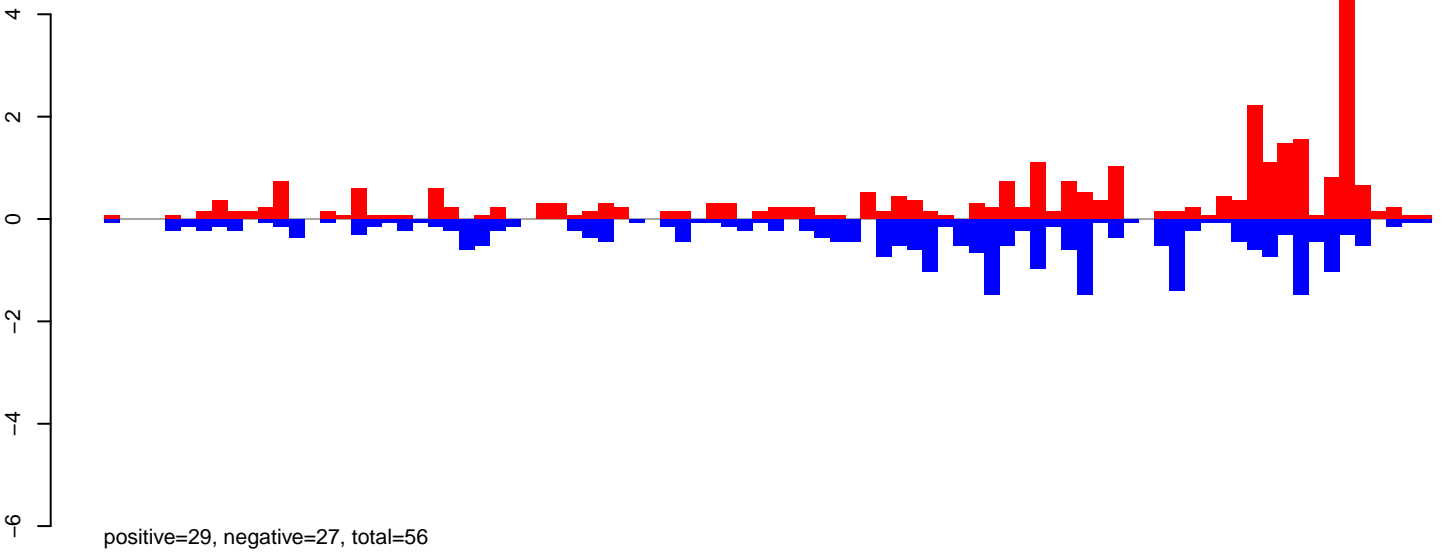
CuQuin_Testes_NL.24_35.rep



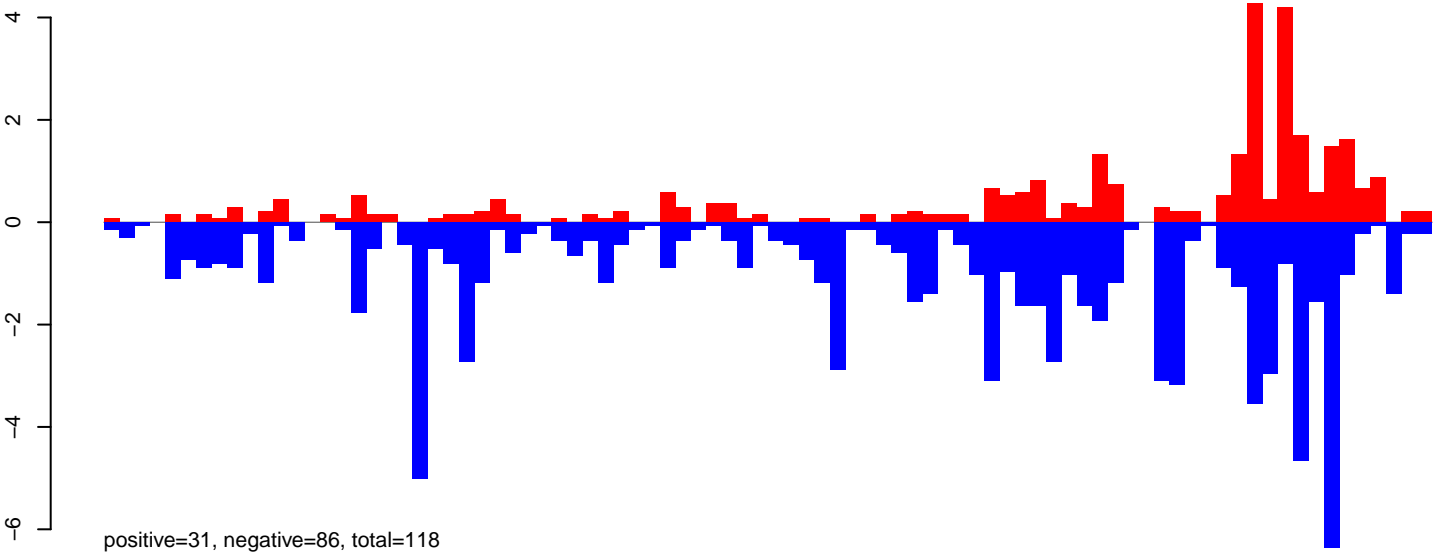
CuQuin_Testes_NL.rep



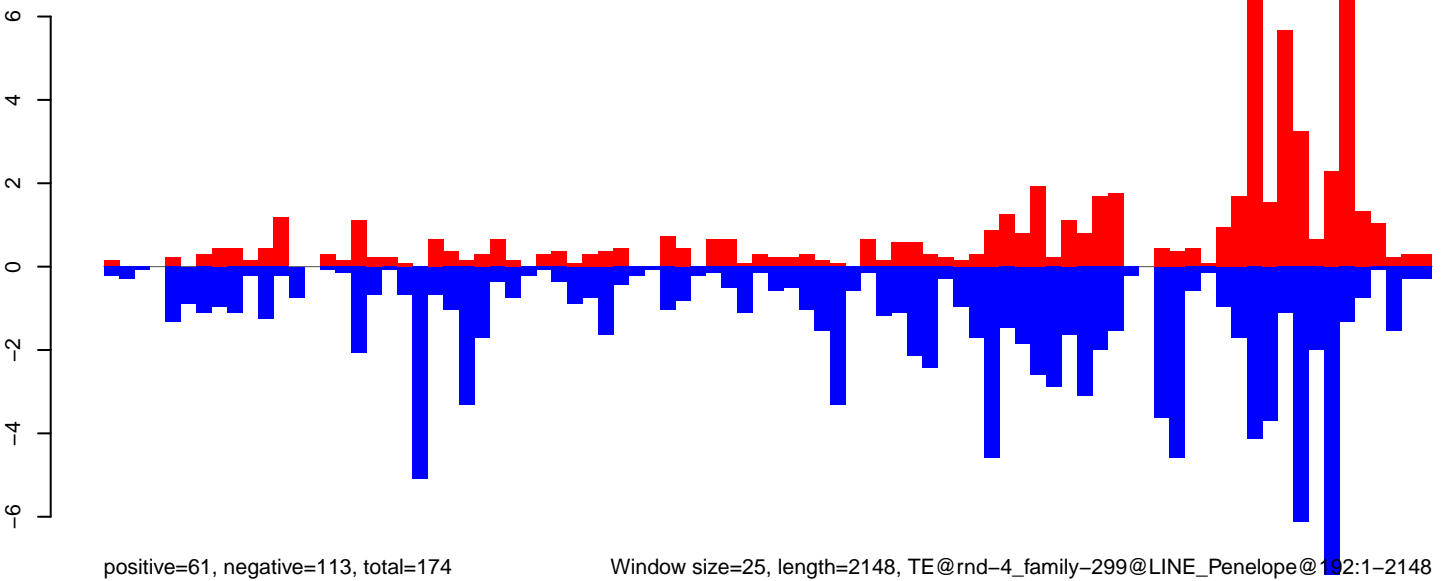
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

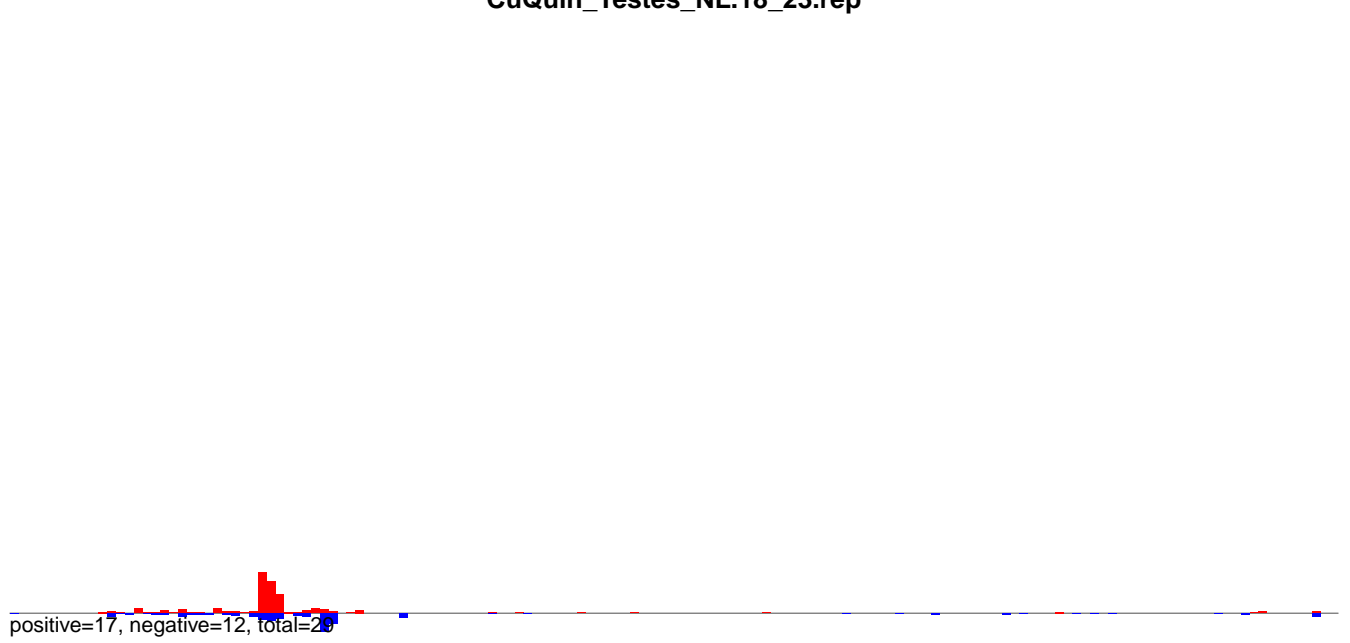


CuQuin_Testes_NL.rep

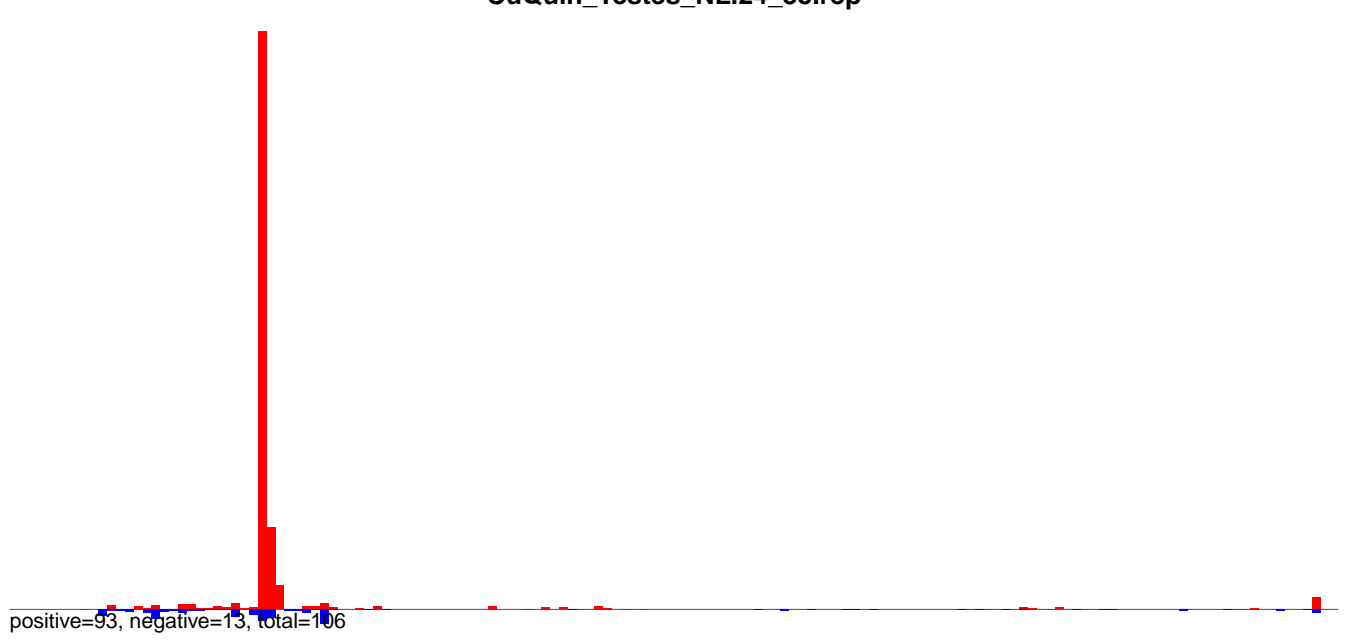


0 500 1000 1500 2000

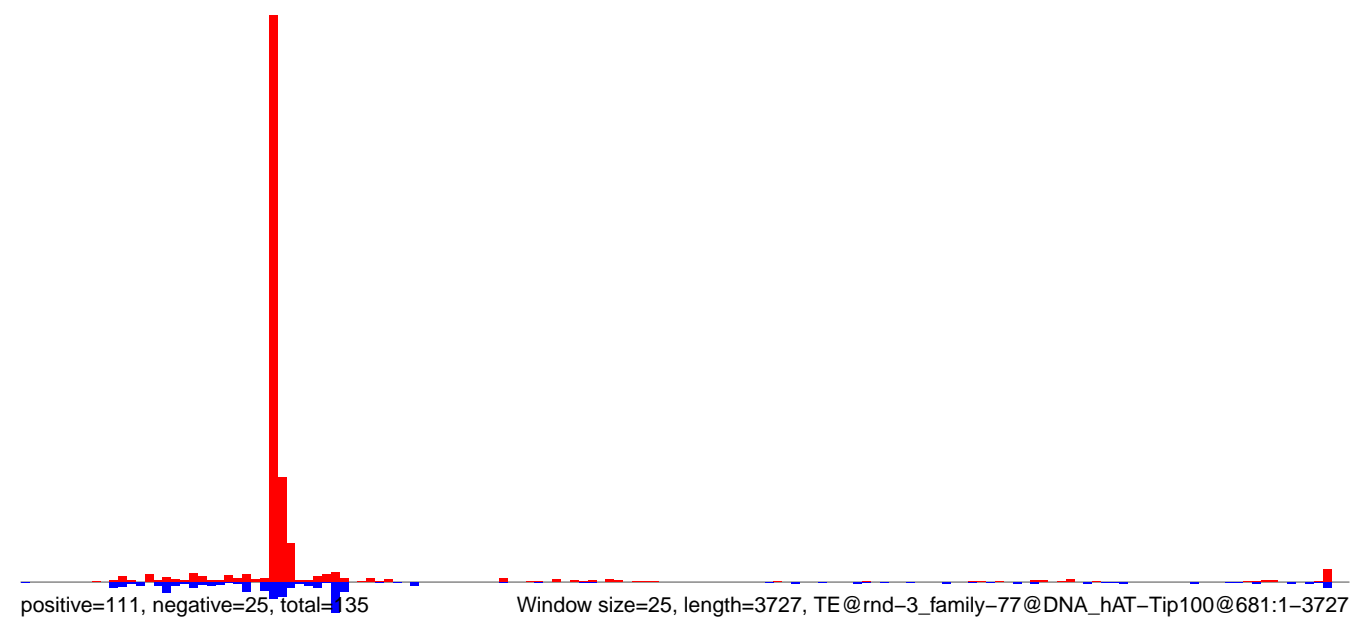
CuQuin_Testes_NL.18_23.rep



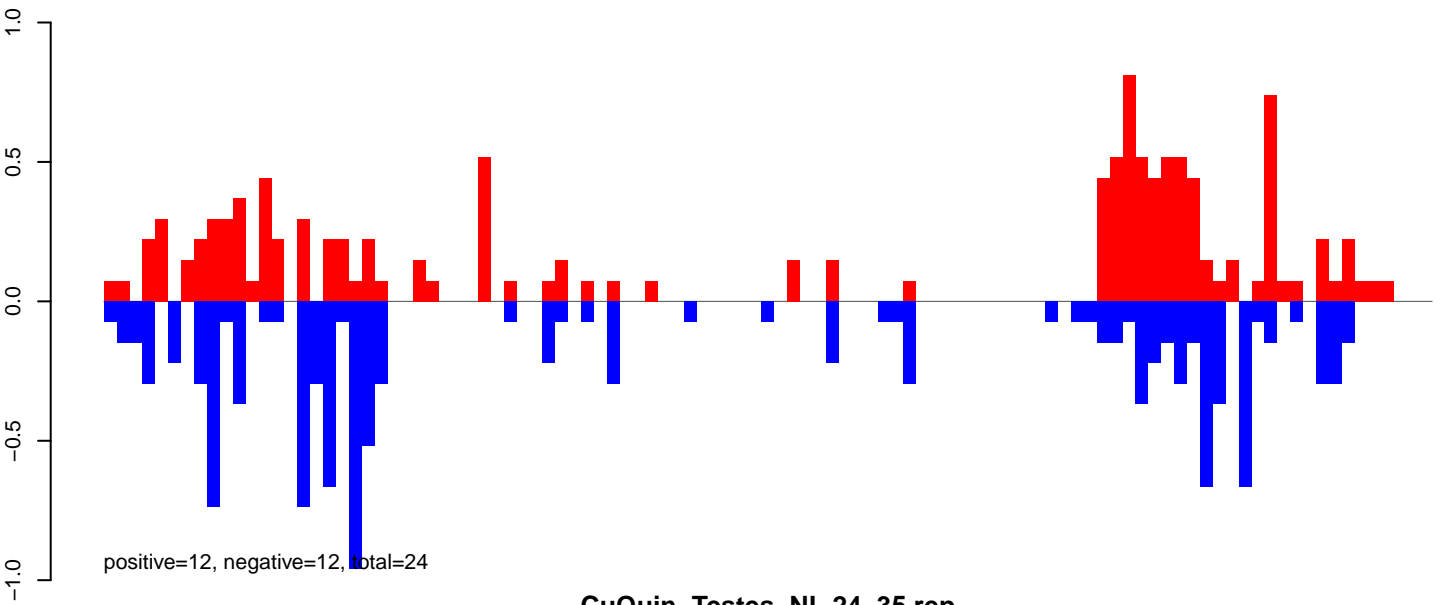
CuQuin_Testes_NL.24_35.rep



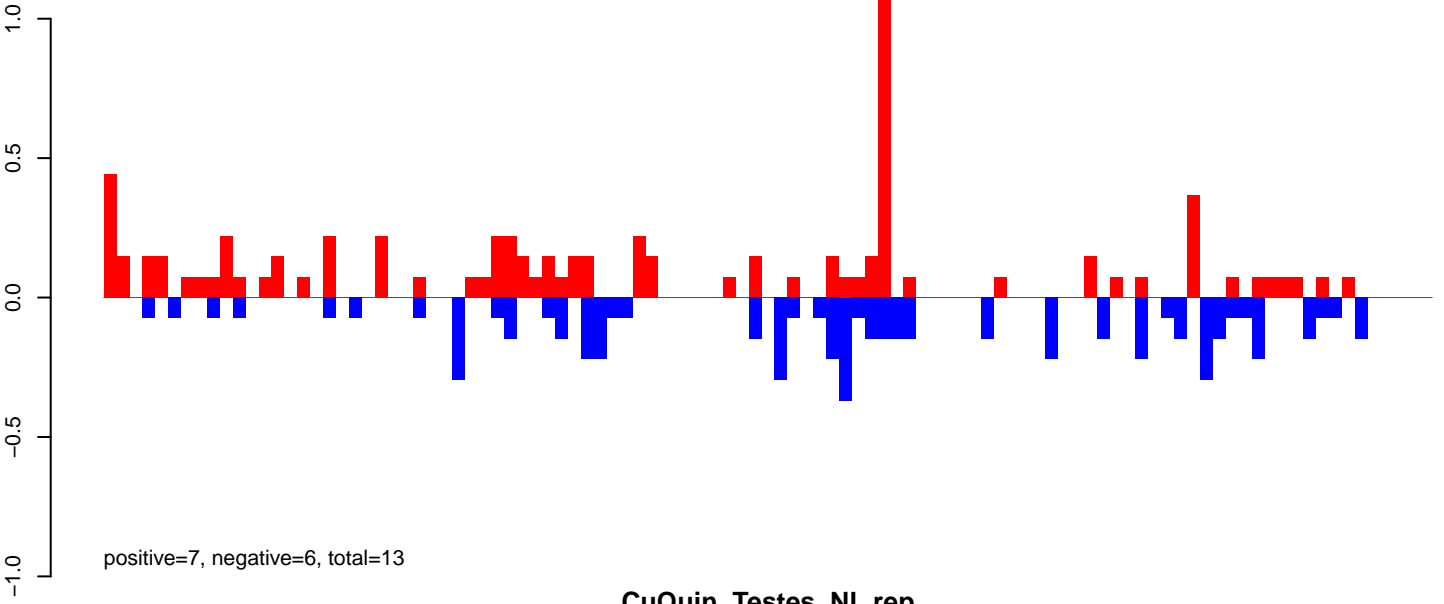
CuQuin_Testes_NL.rep



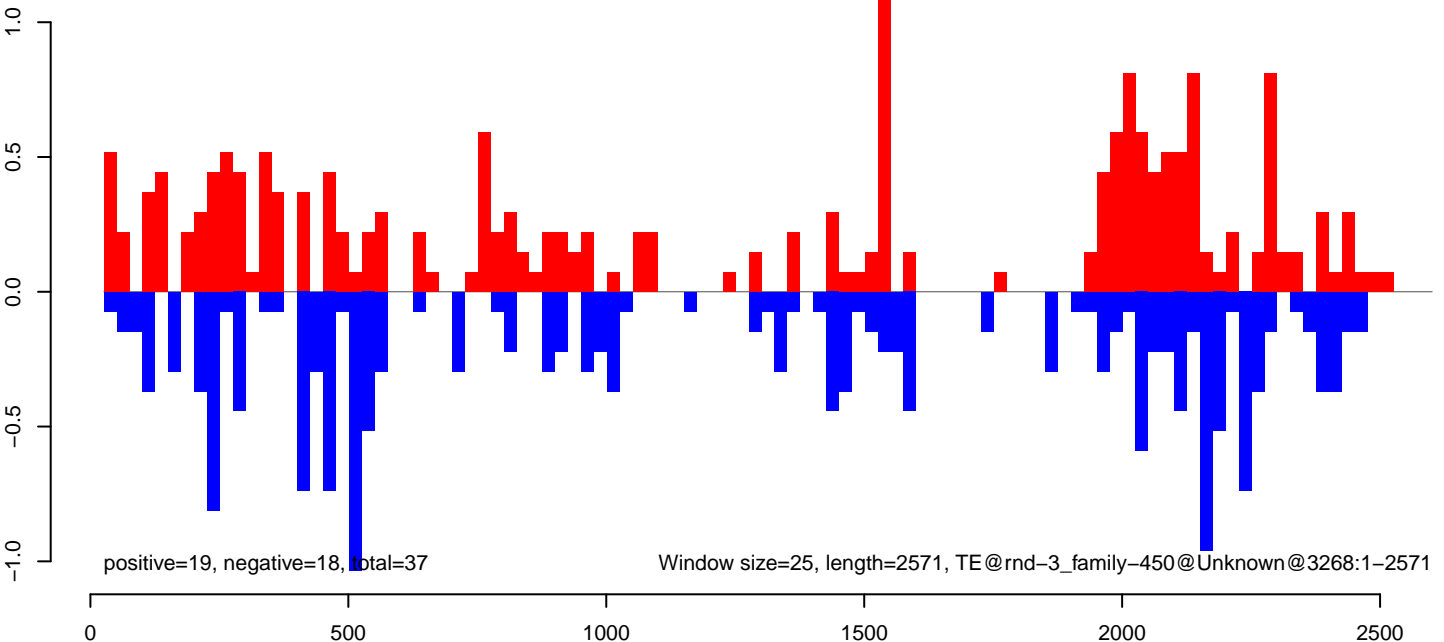
CuQuin_Testes_NL.18_23.rep



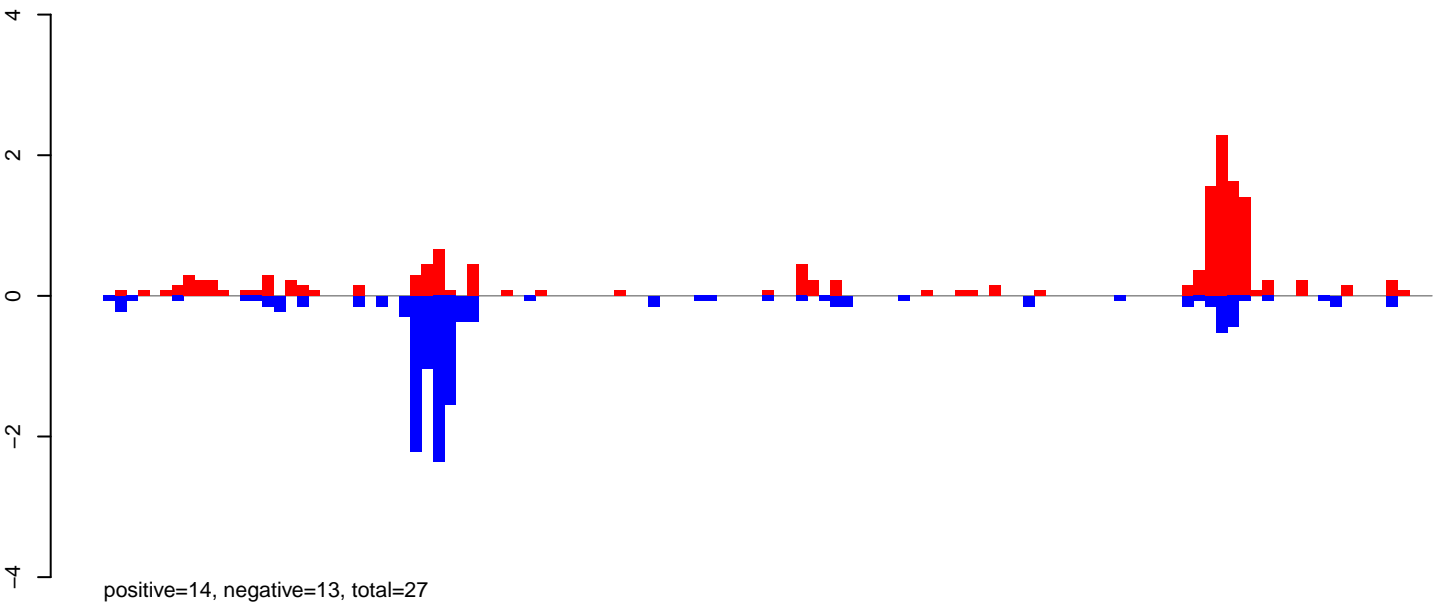
CuQuin_Testes_NL.24_35.rep



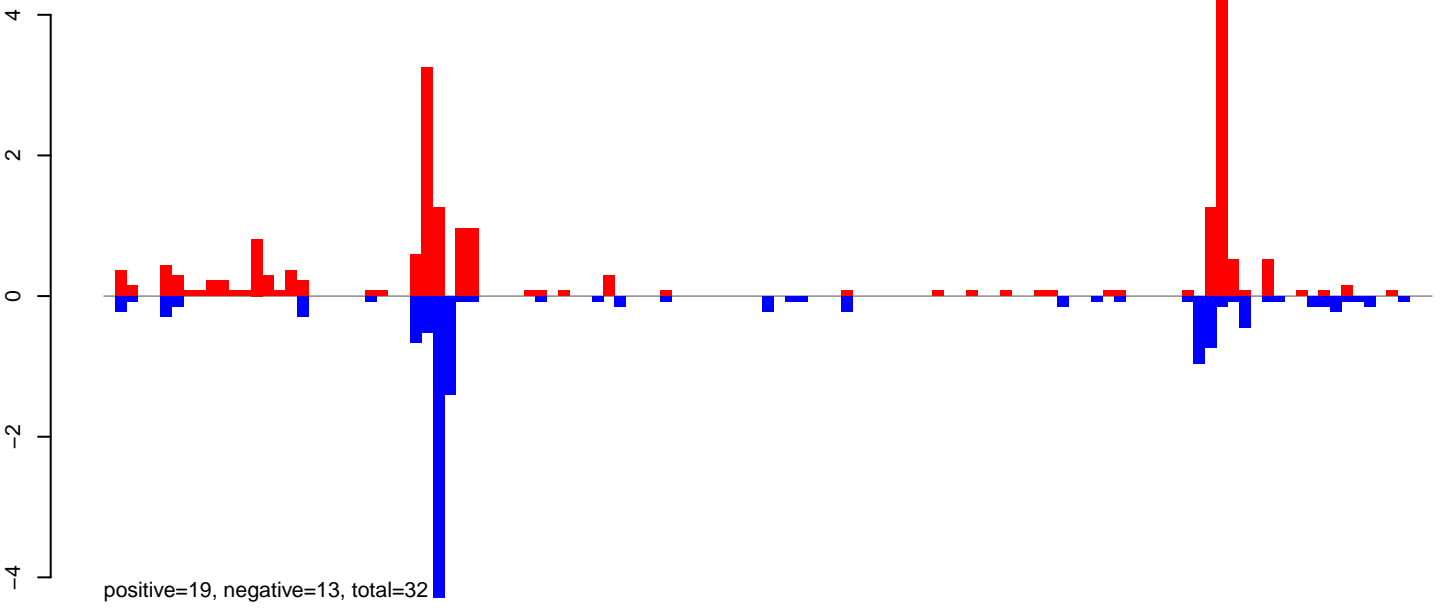
CuQuin_Testes_NL.rep



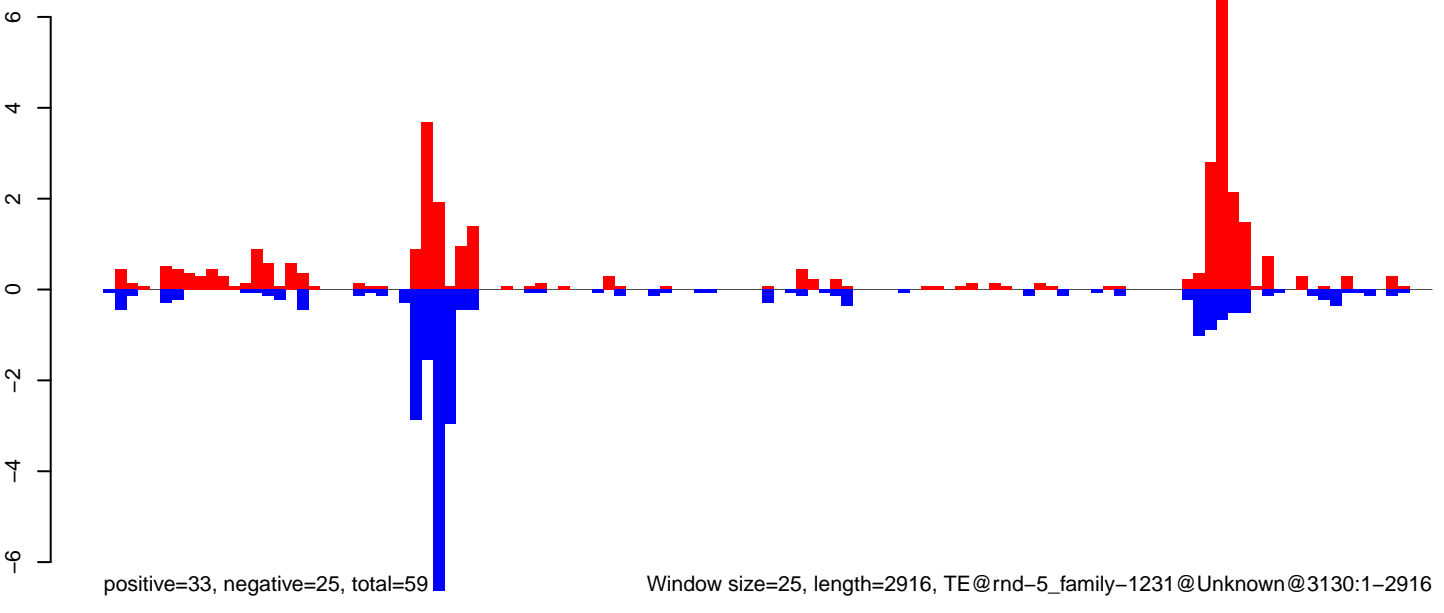
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

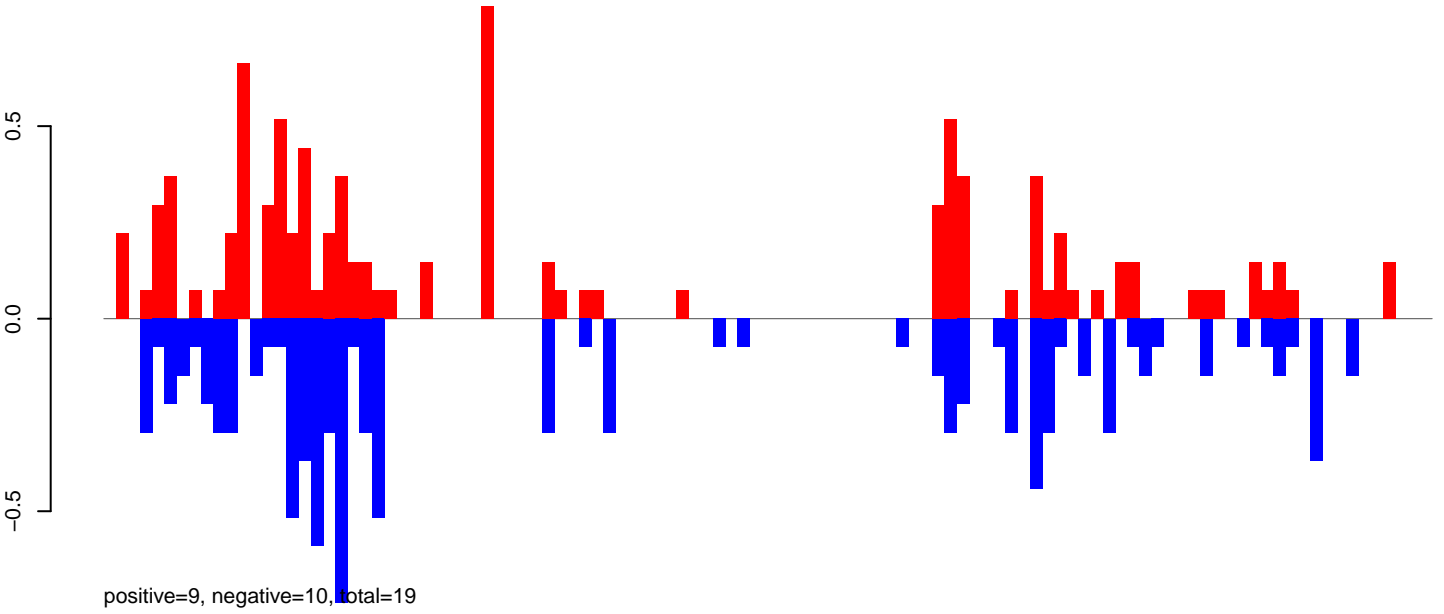


CuQuin_Testes_NL.rep

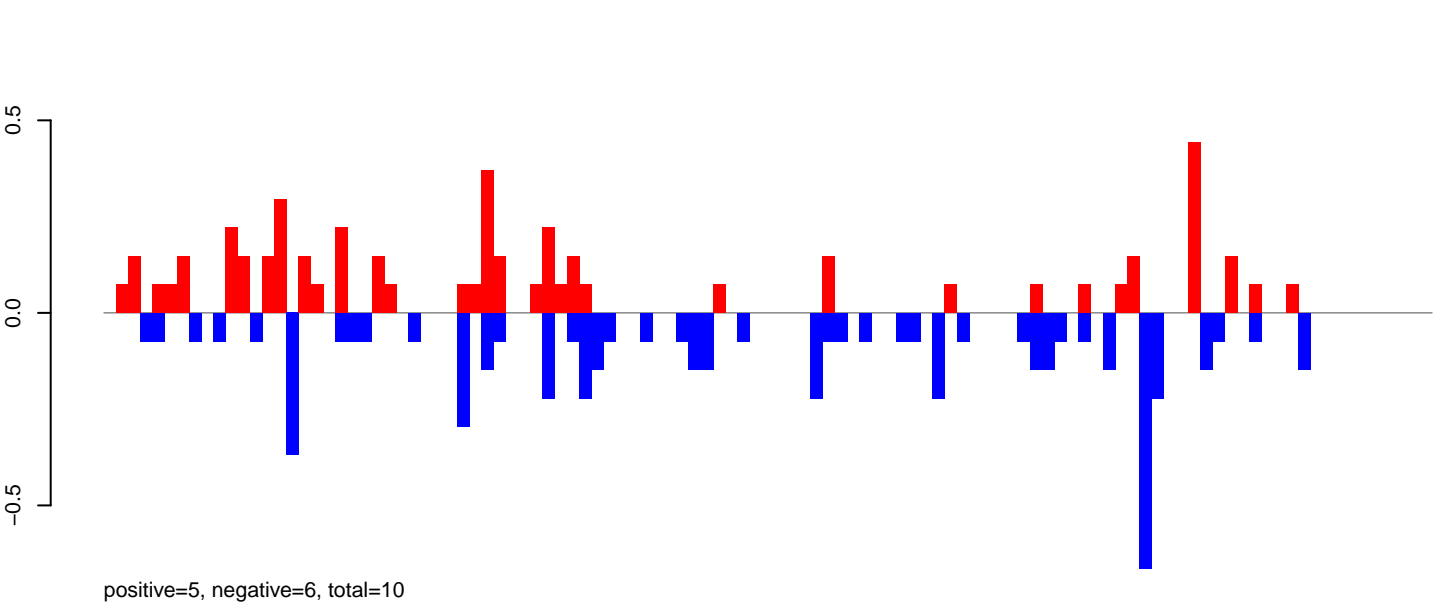


Window size=25, length=2916, TE@rnd-5_family-1231@Unknown@3130:1-2916

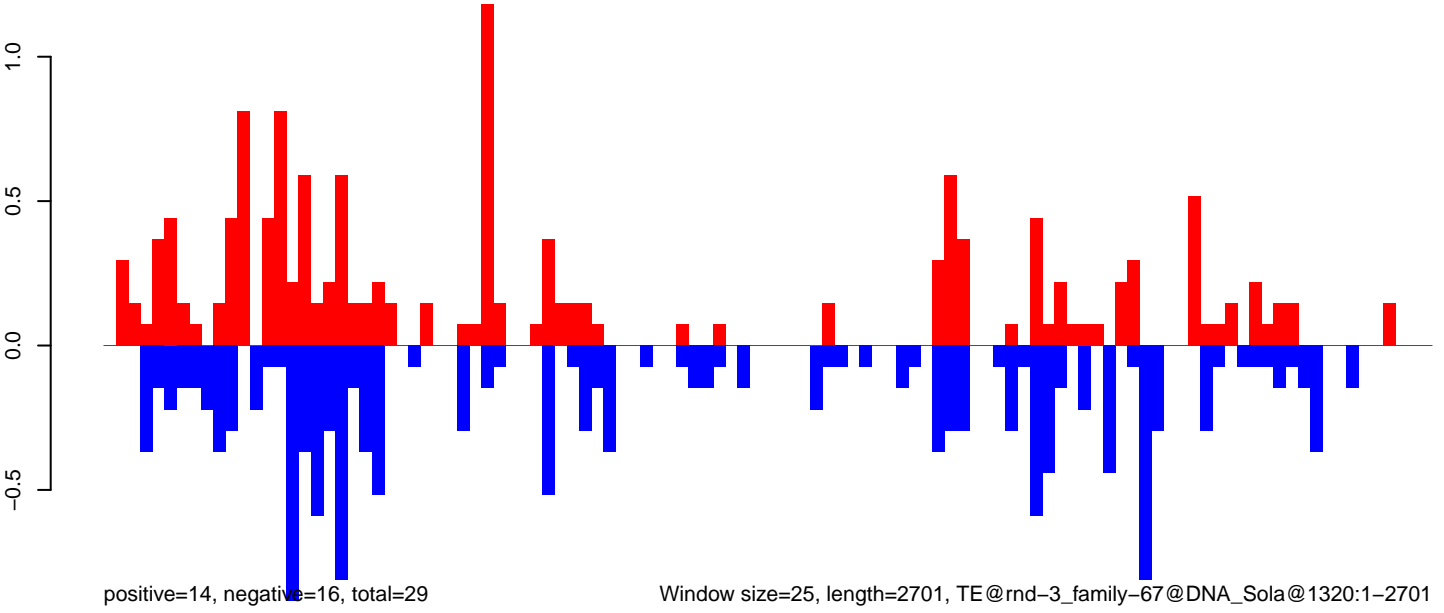
CuQuin_Testes_NL.18_23.rep



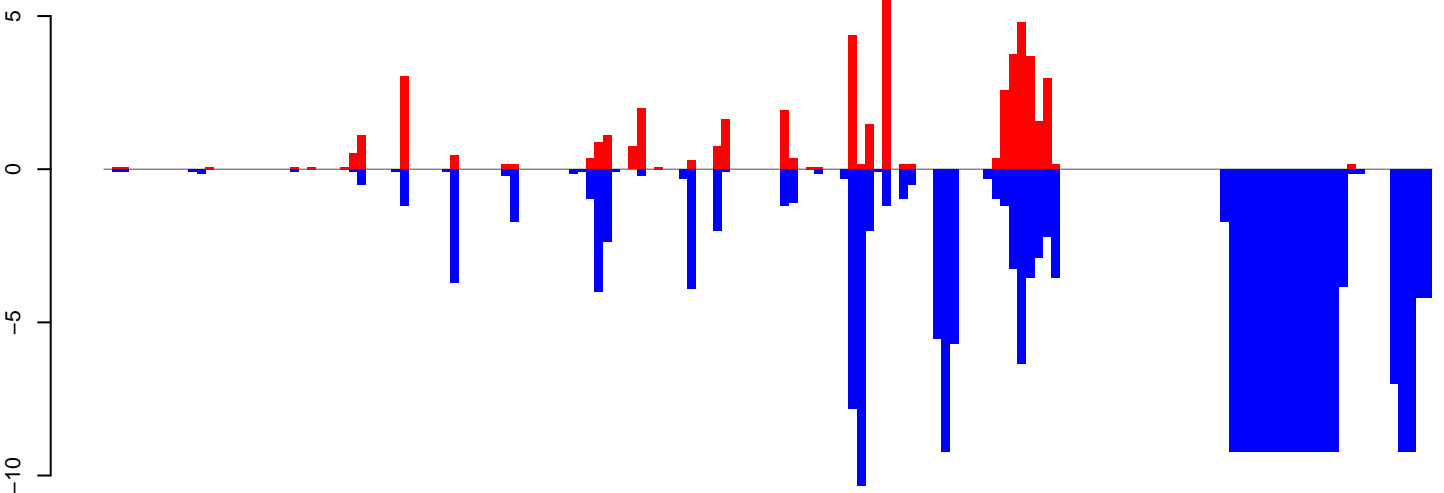
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

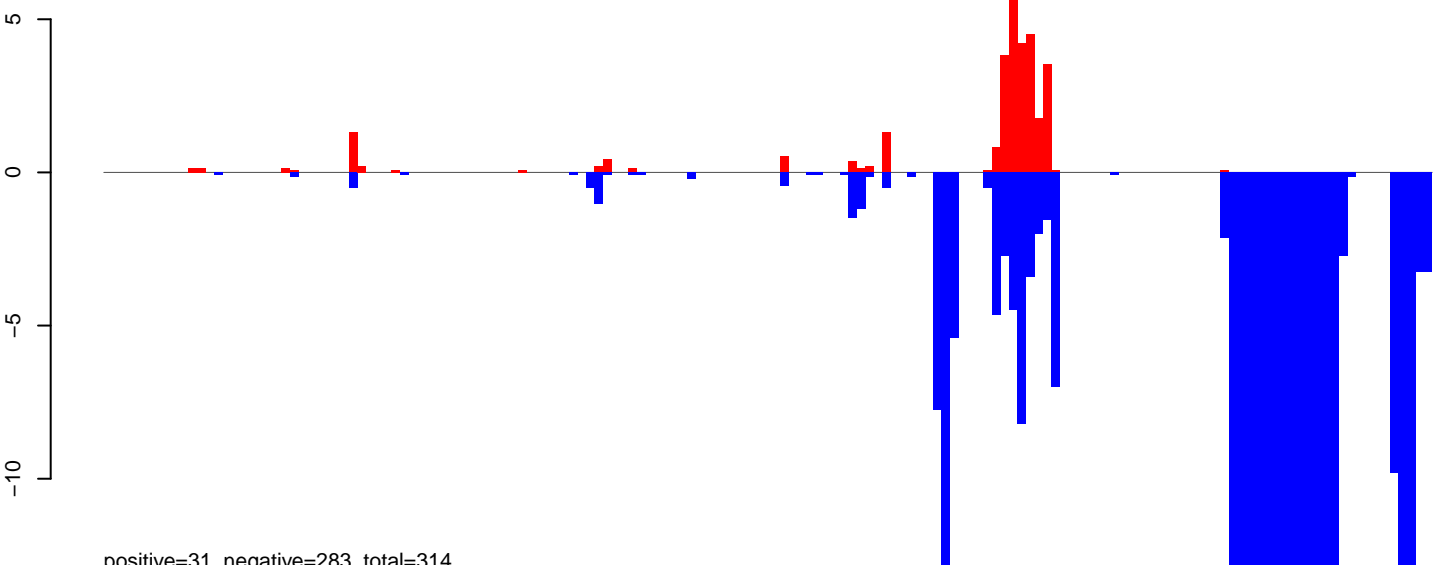


CuQuin_Testes_NL.18_23.rep



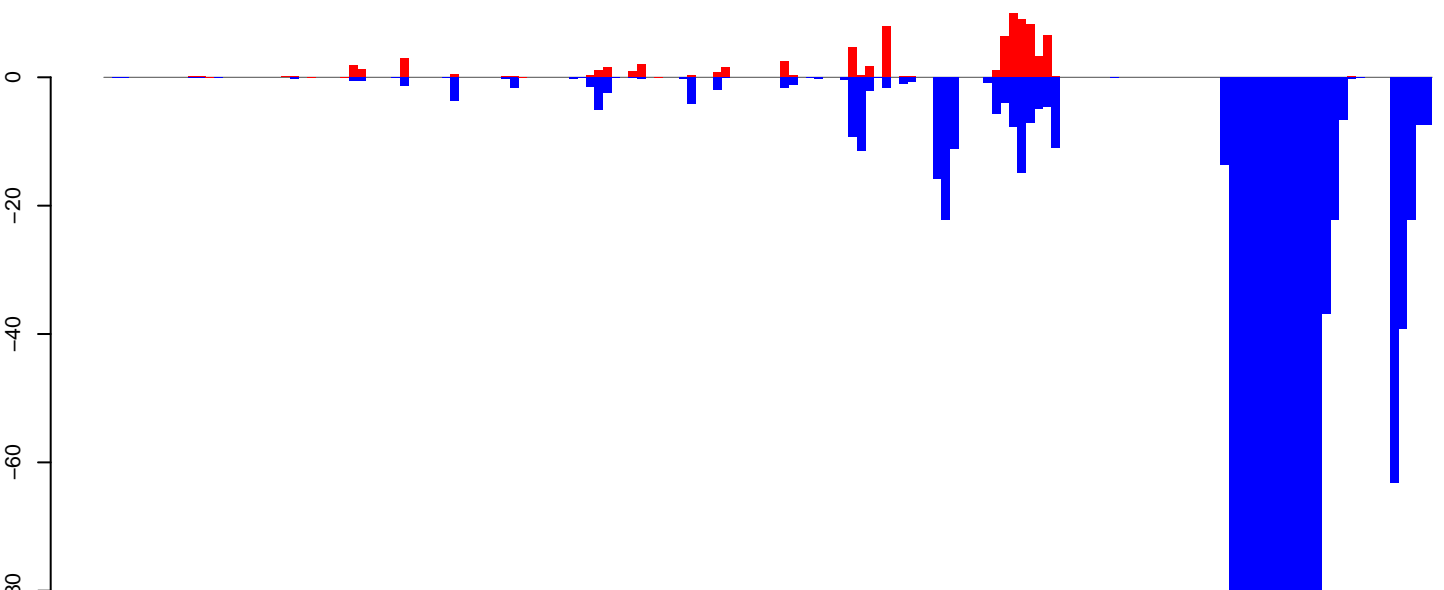
positive=49, negative=252, total=301

CuQuin_Testes_NL.24_35.rep



positive=31, negative=283, total=314

CuQuin_Testes_NL.rep

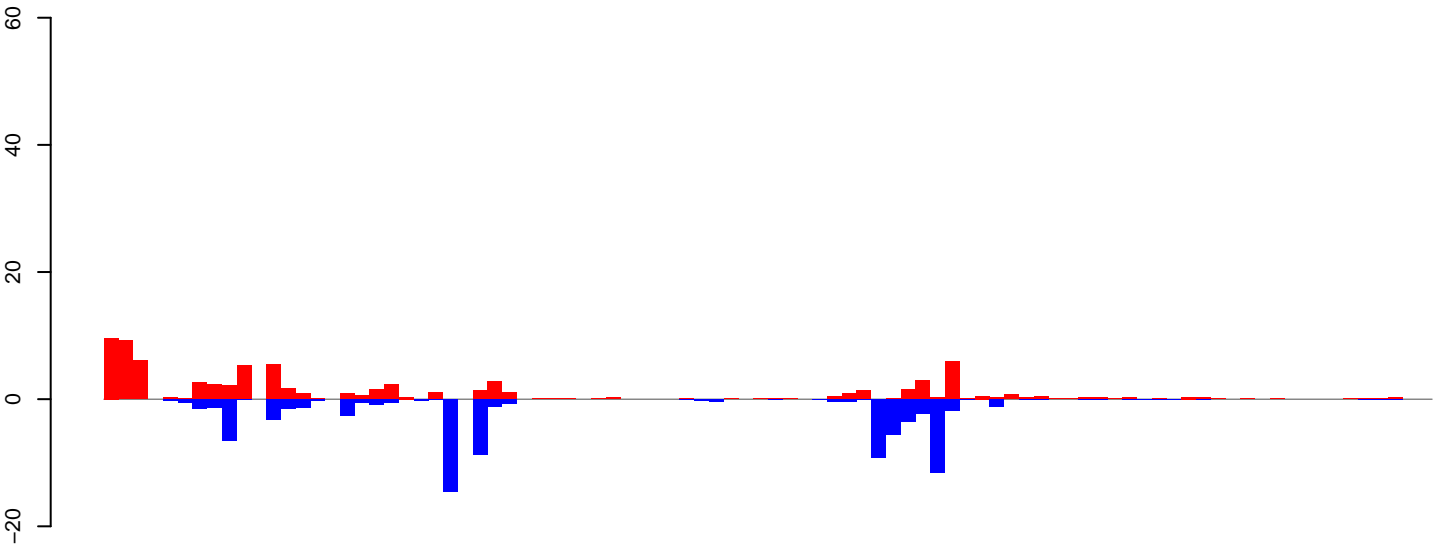


positive=80, negative=1297, total=1377

Window size=25, length=3908, TE@rnd-3_family-461@DNA_Sola-2@683:1-3908

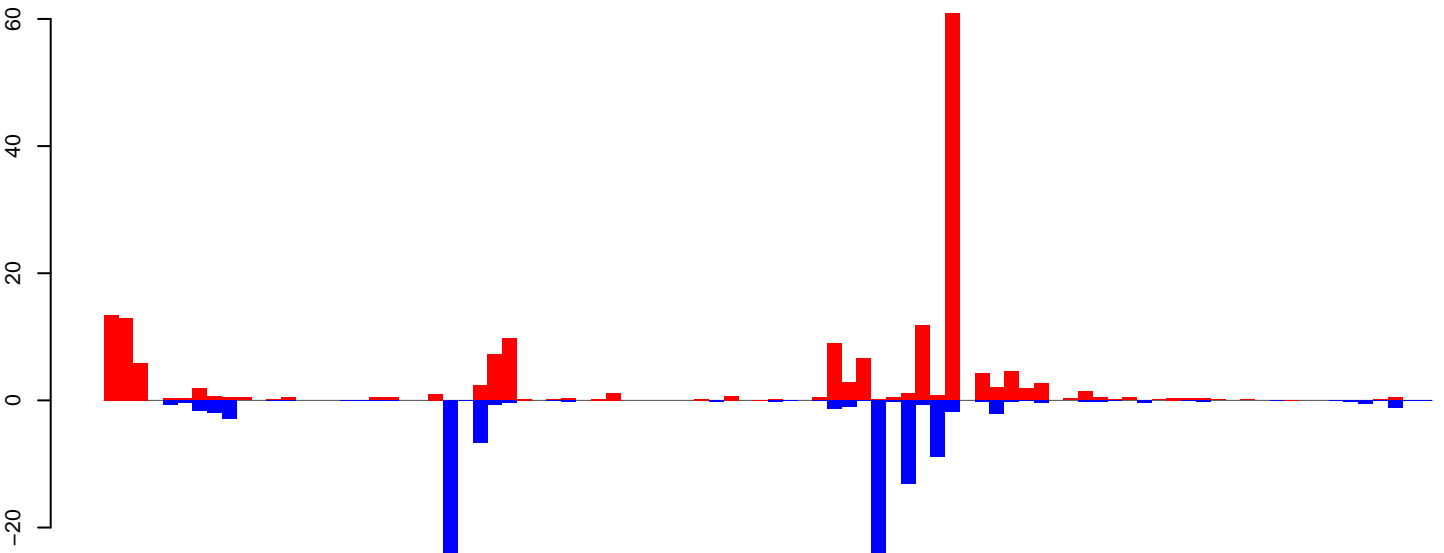
0 1000 2000 3000 4000

CuQuin_Testes_NL.18_23.rep



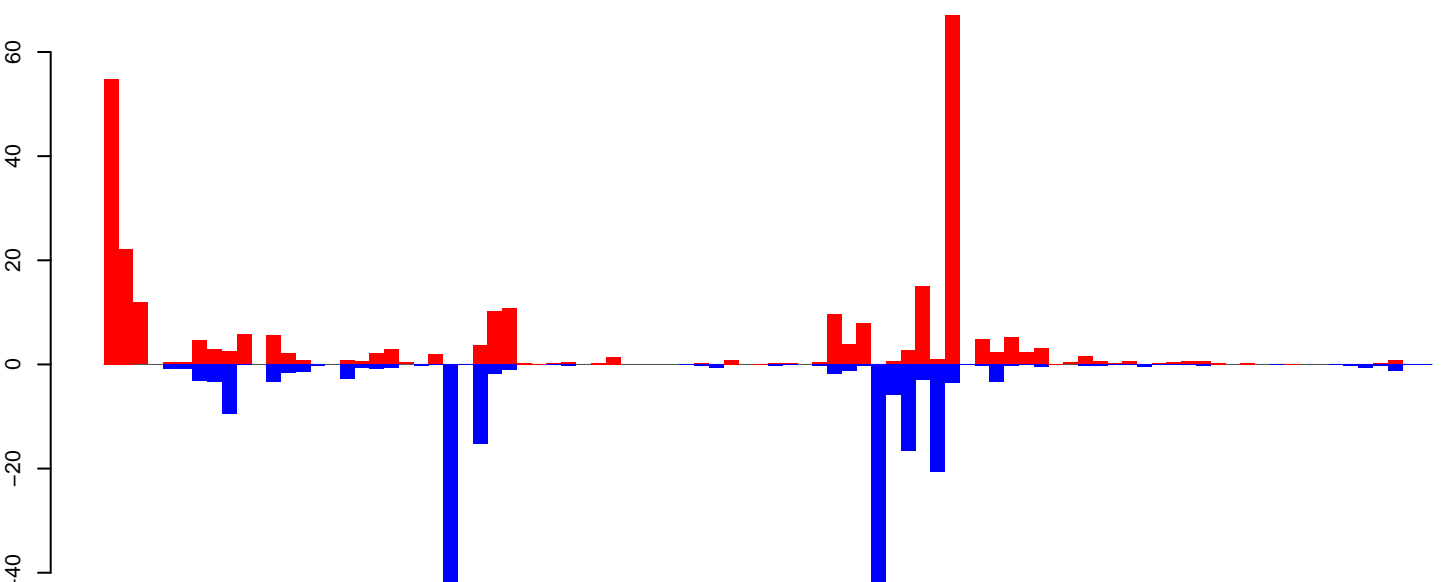
positive=78, negative=84, total=162

CuQuin_Testes_NL.24_35.rep



positive=178, negative=117, total=295

CuQuin_Testes_NL.rep



positive=287, negative=202, total=489

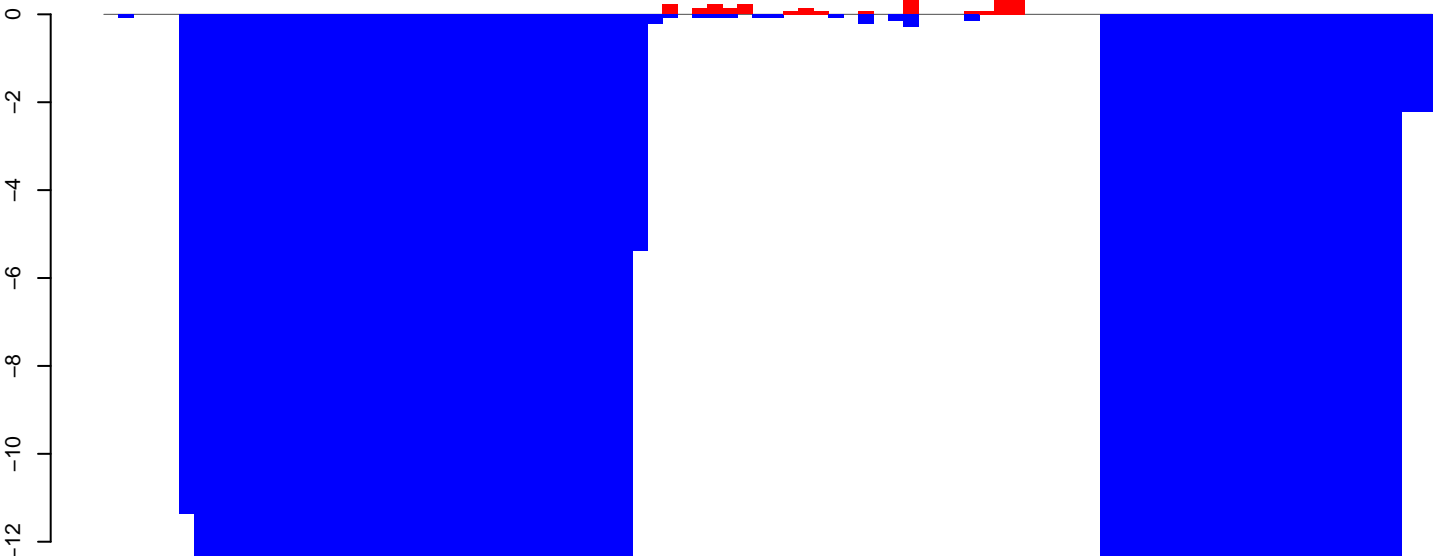
Window size=25, length=2234, TE@rnd-5_family-517@Unknown@415:1-2234

0 500 1000 1500 2000

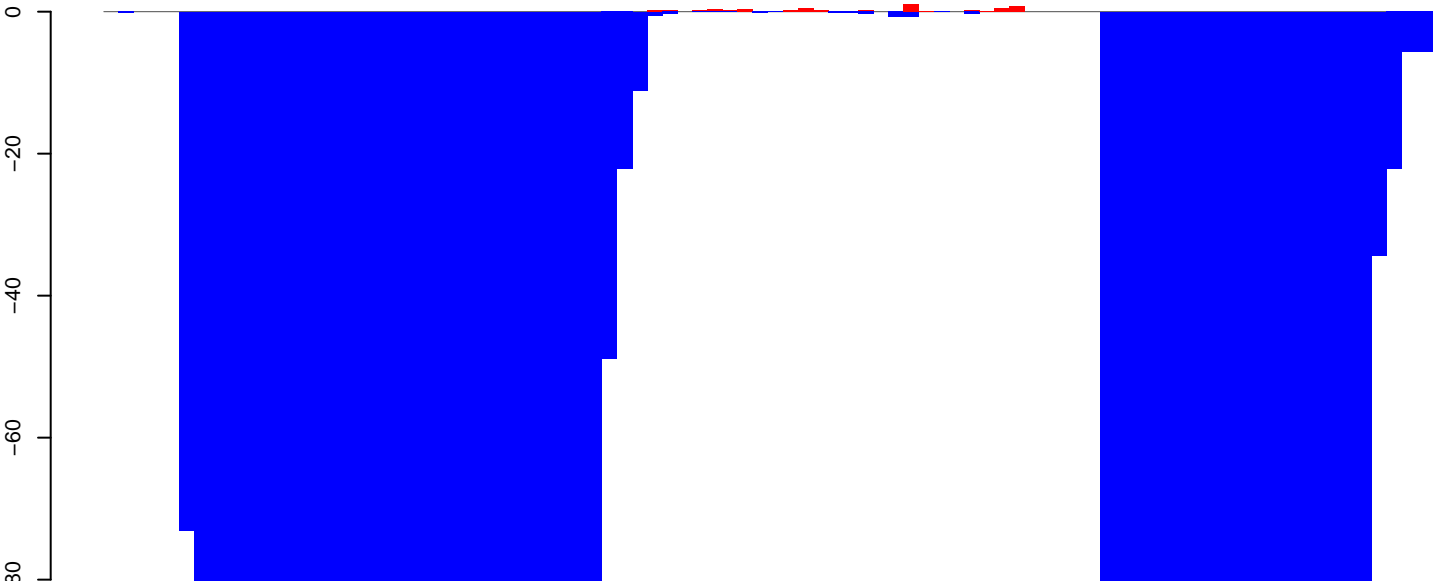
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



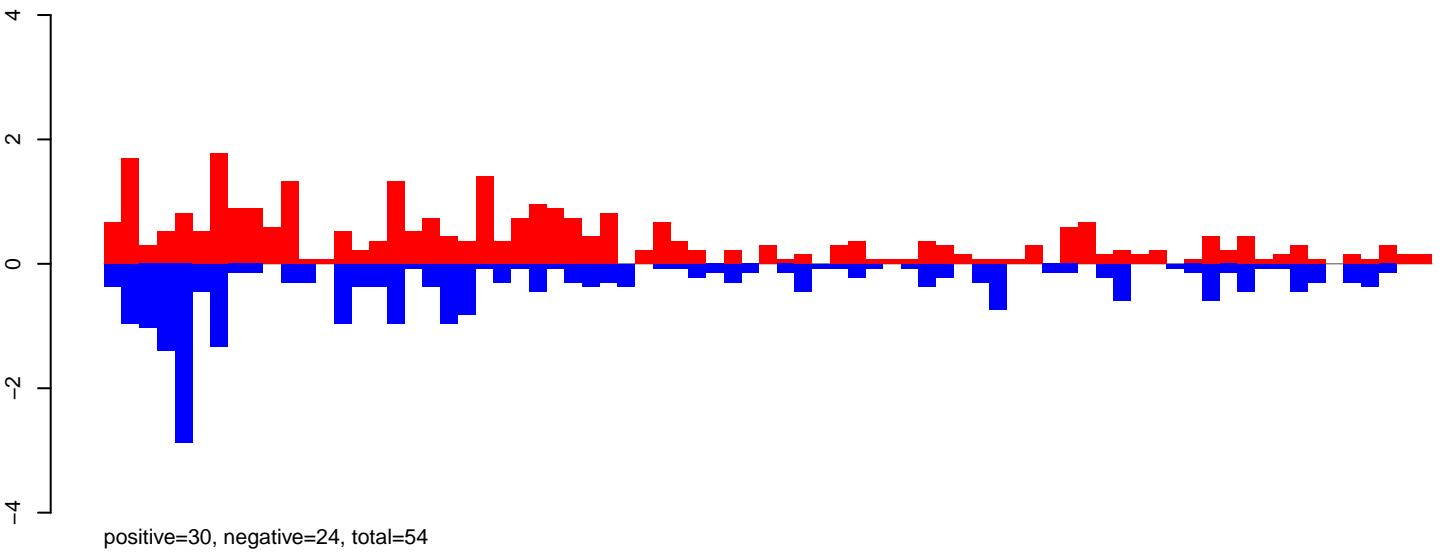
CuQuin_Testes_NL.rep



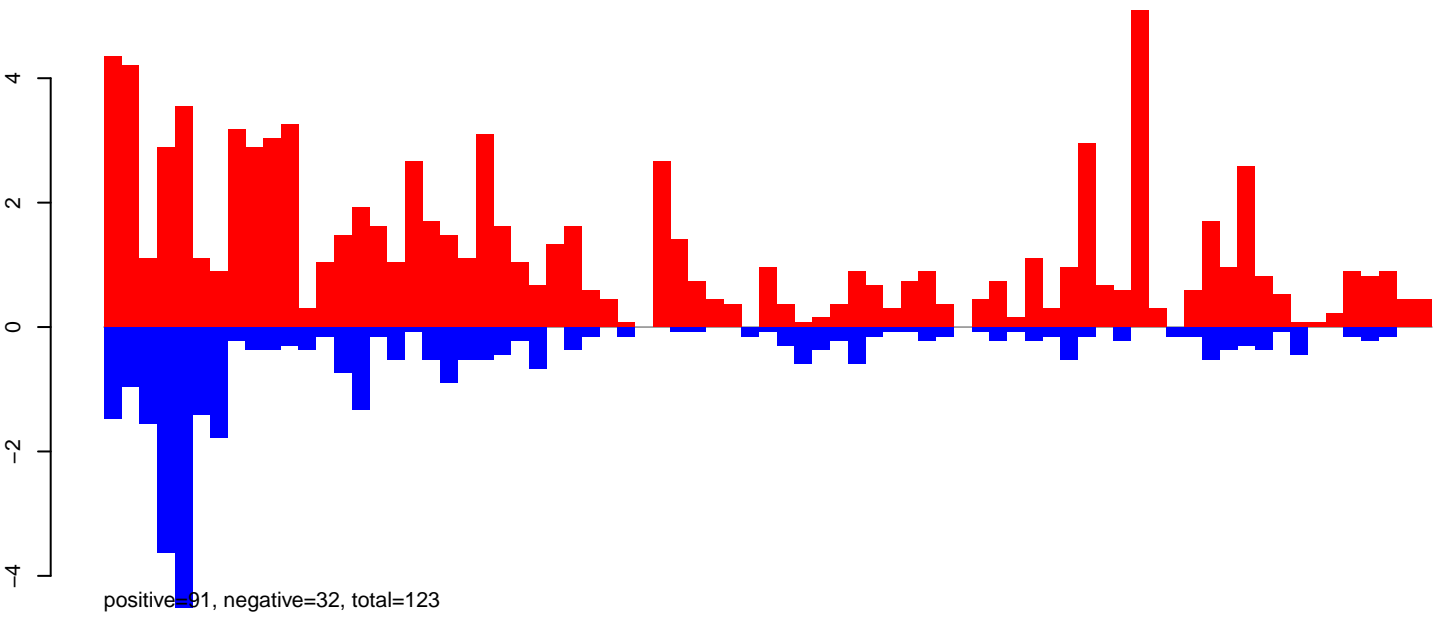
Window size=25, length=2181, TE@rnd-2_family-273@DNA_Sola-1@640:1-2181

0 500 1000 1500 2000

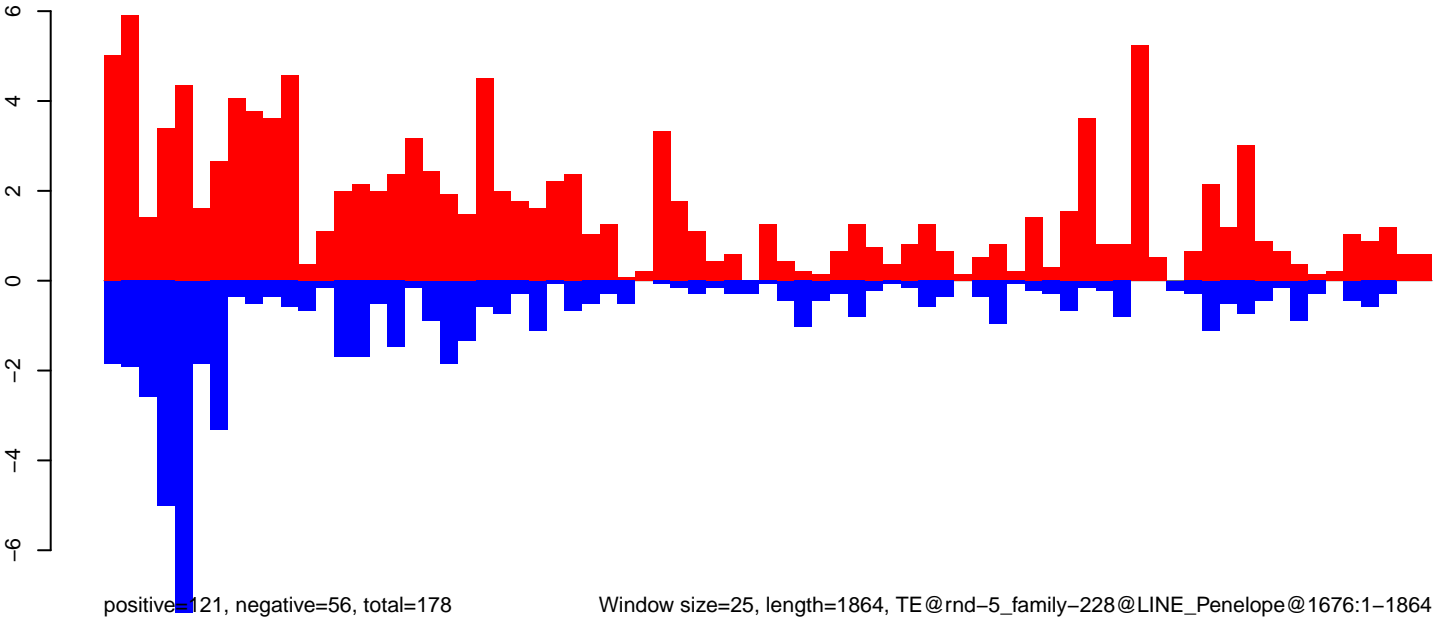
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



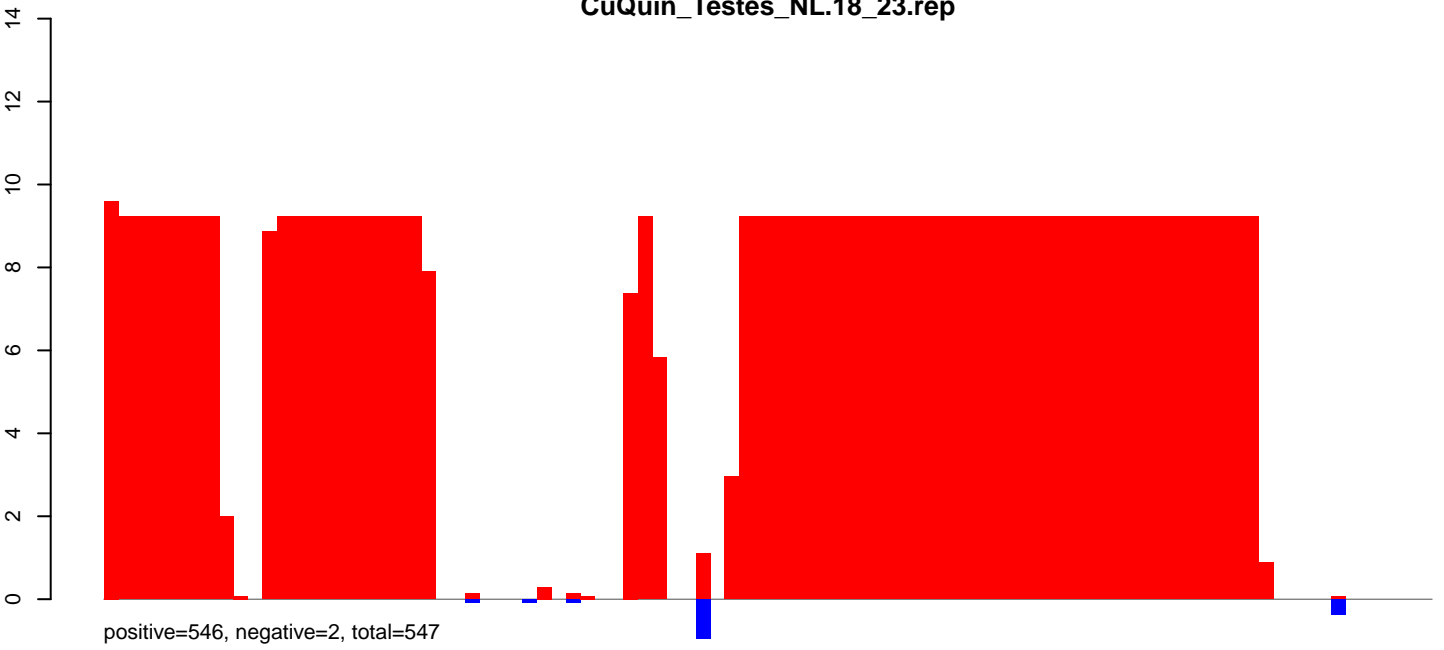
CuQuin_Testes_NL.rep



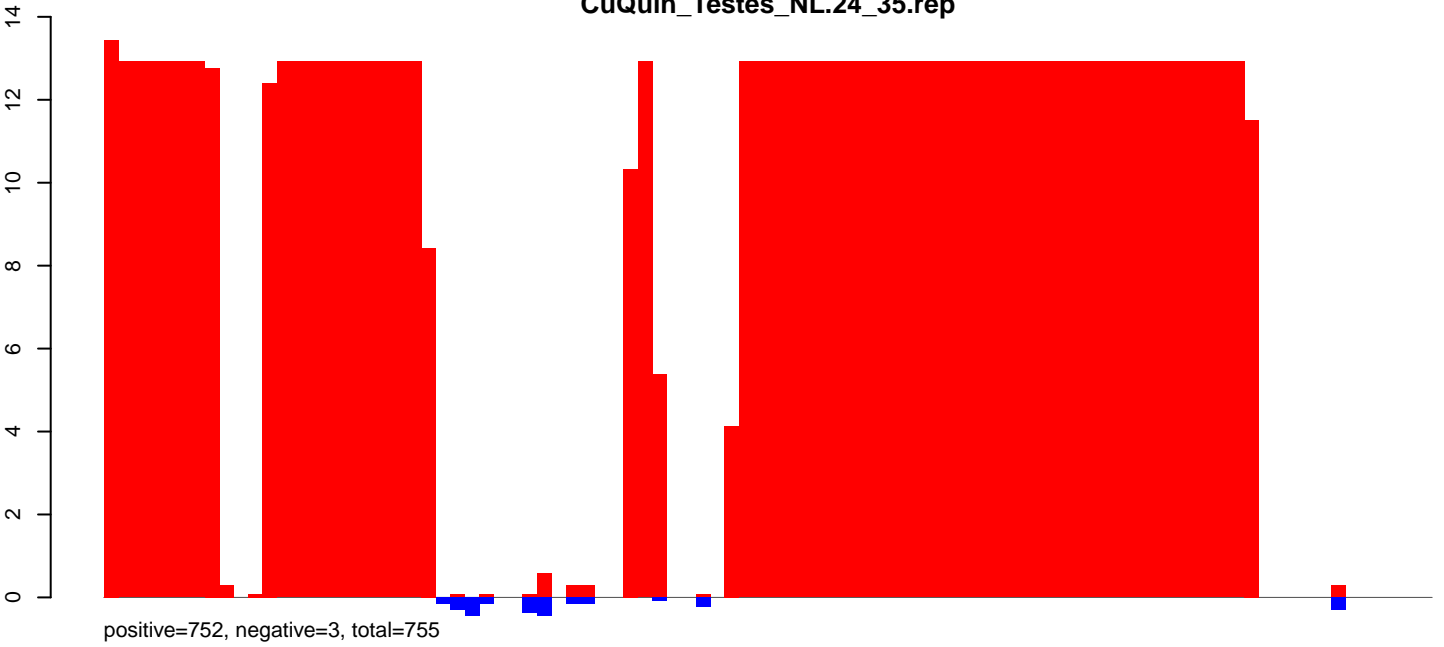
Window size=25, length=1864, TE@rnd-5_family-228@LINE_Penelope@1676:1-1864

0 500 1000 1500

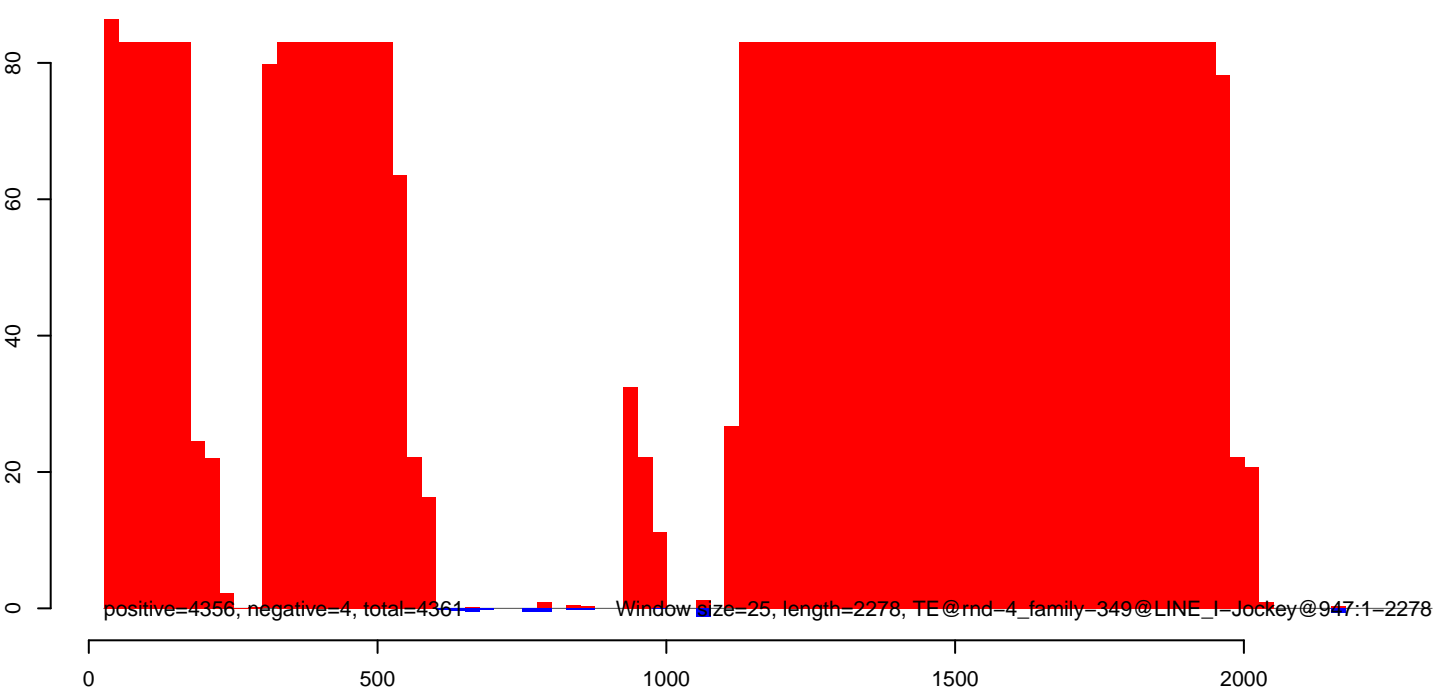
CuQuin_Testes_NL.18_23.rep



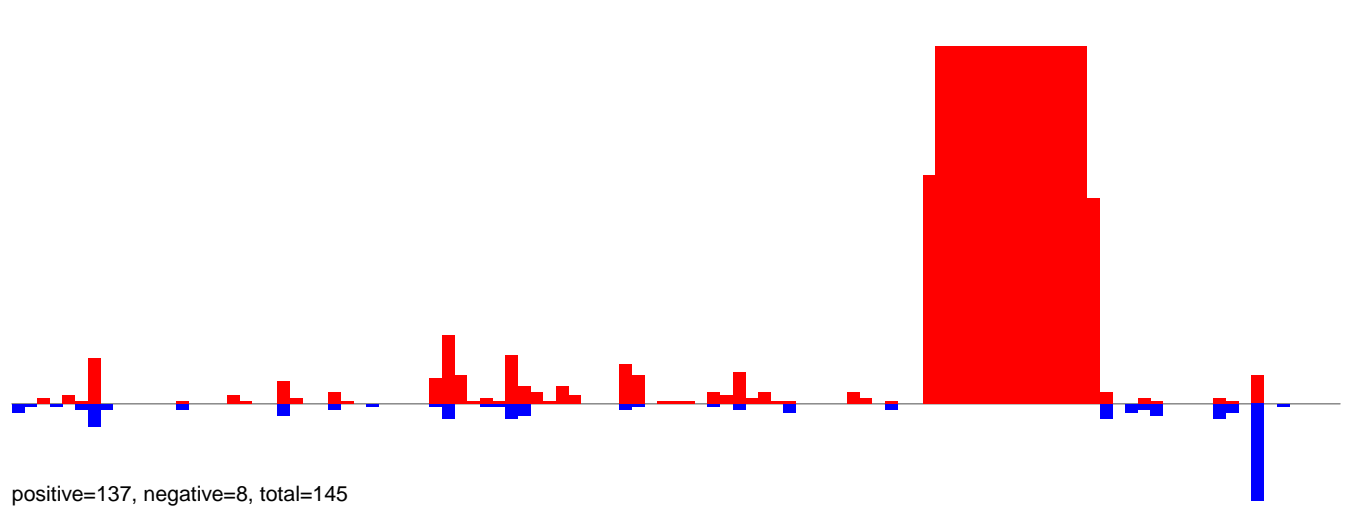
CuQuin_Testes_NL.24_35.rep



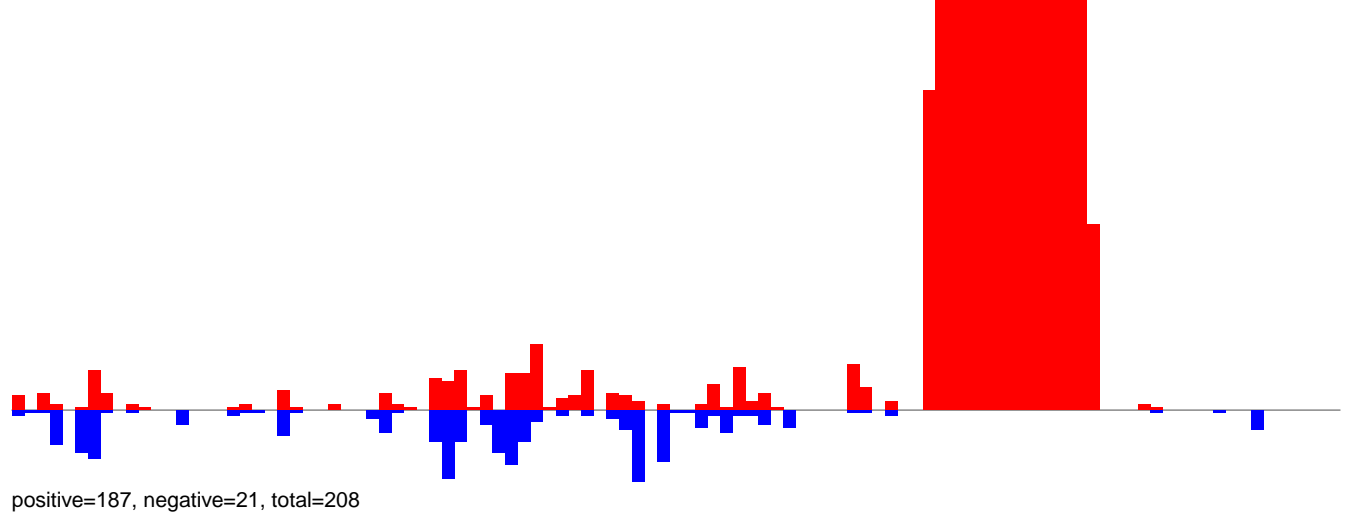
CuQuin_Testes_NL.rep



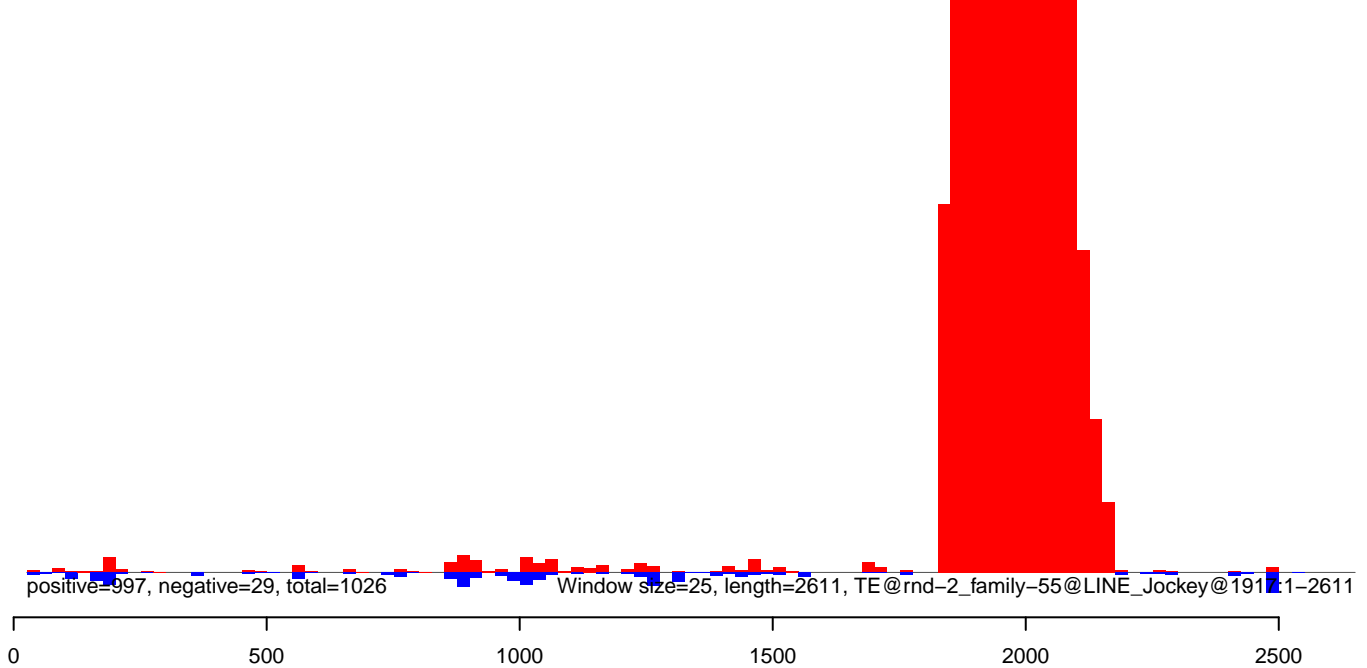
CuQuin_Testes_NL.18_23.rep



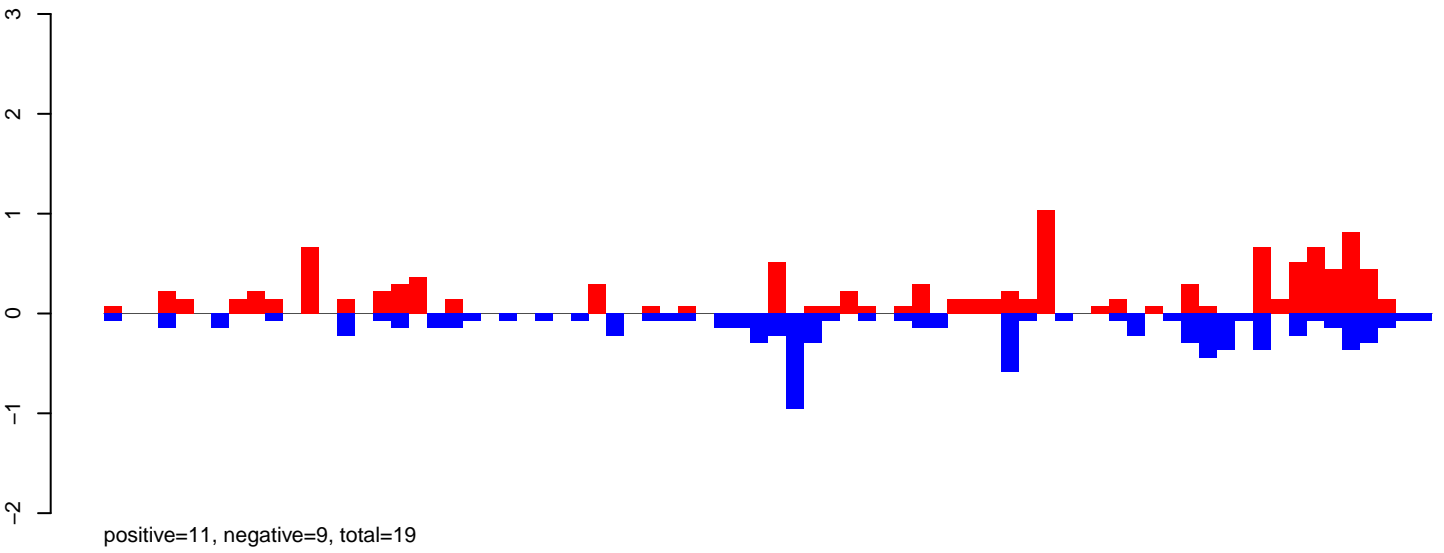
CuQuin_Testes_NL.24_35.rep



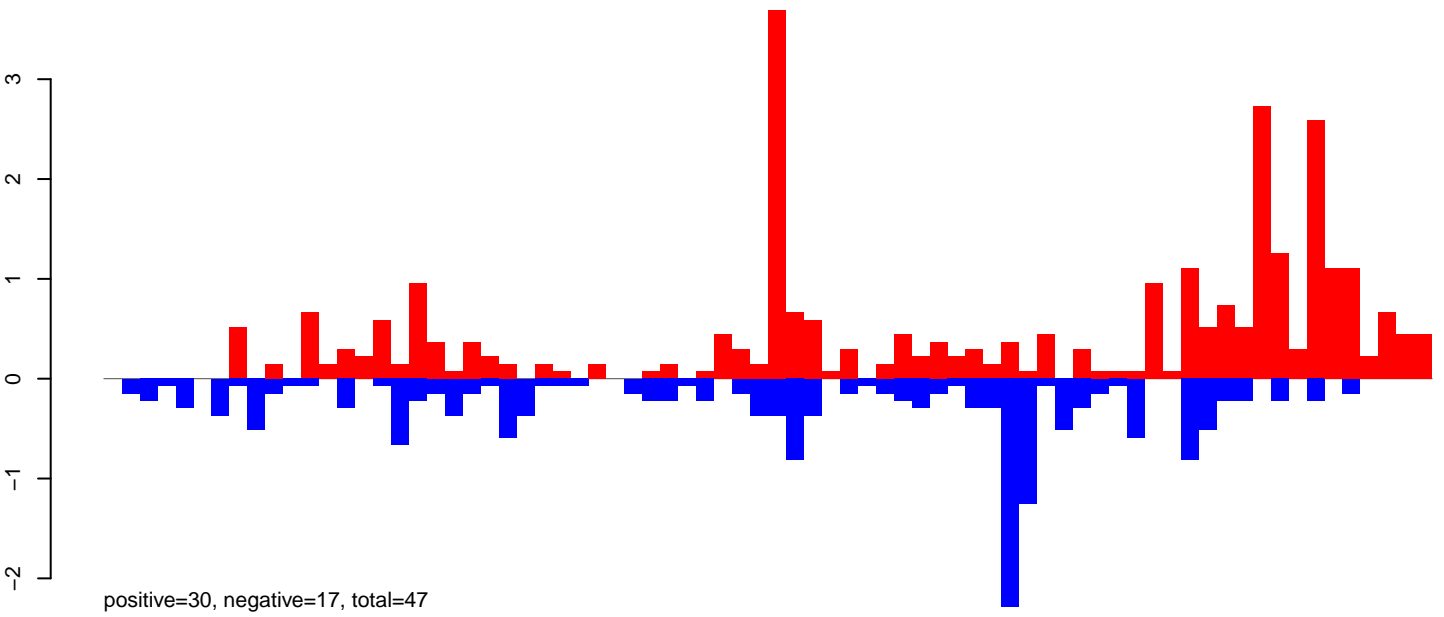
CuQuin_Testes_NL.rep



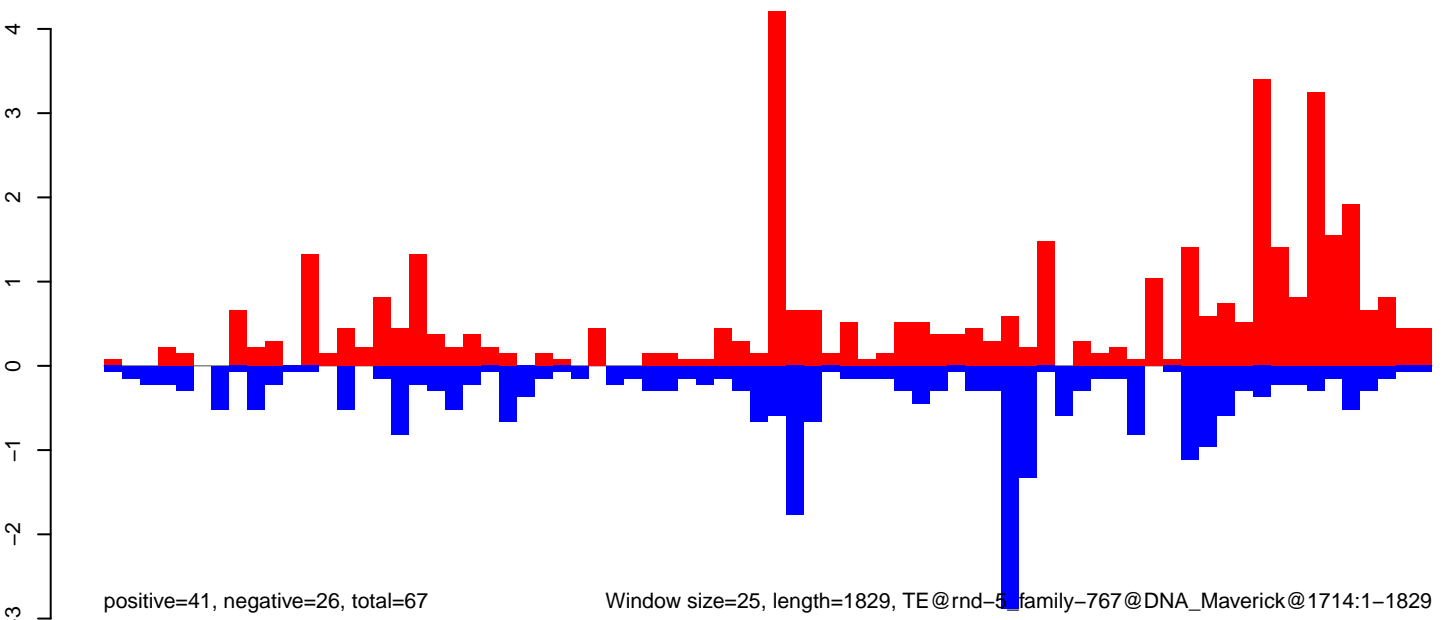
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

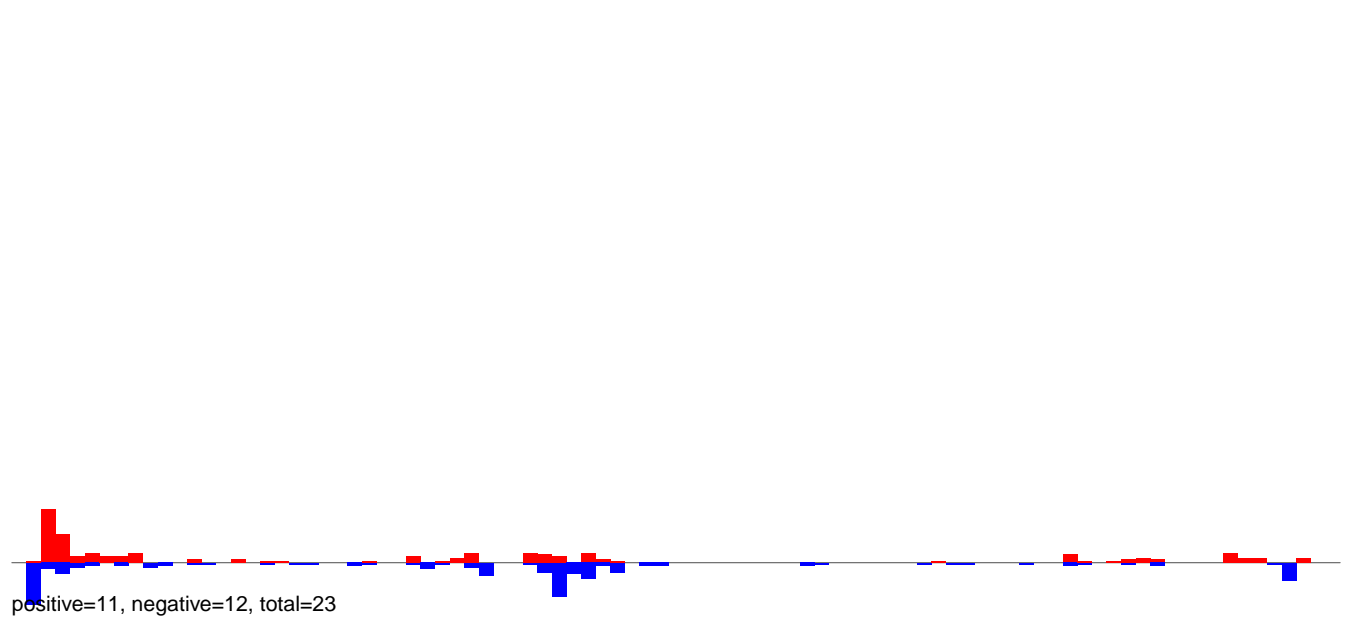


CuQuin_Testes_NL.rep

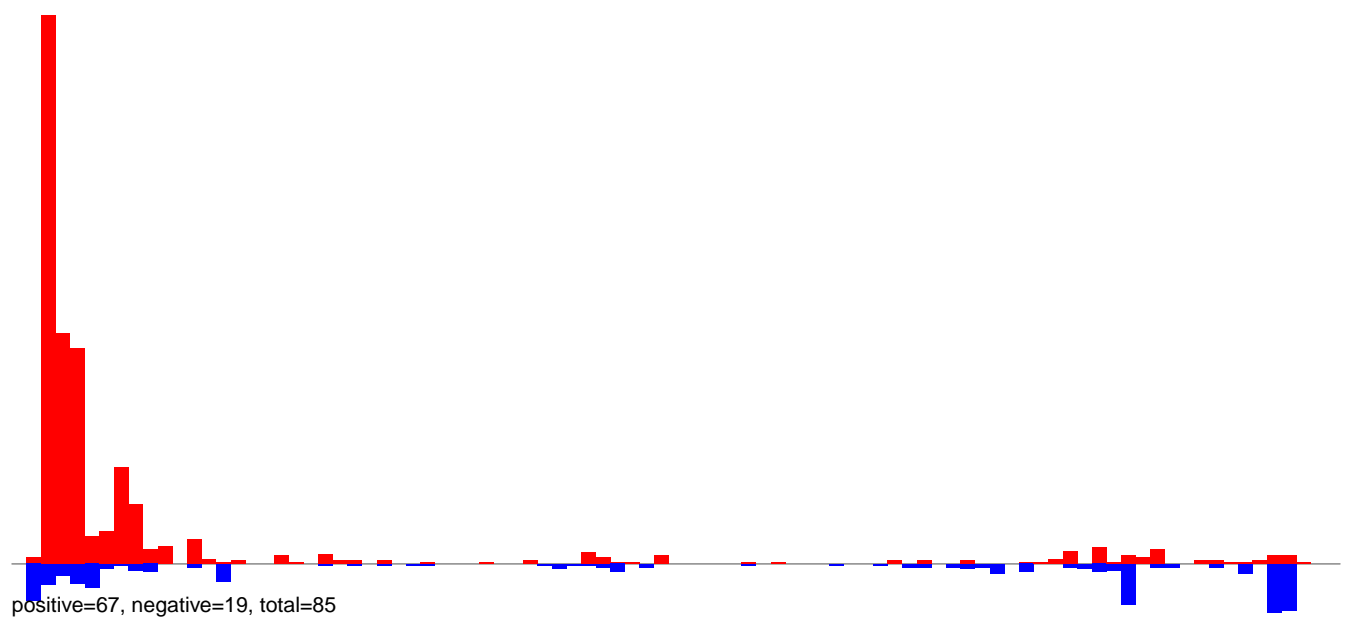


Window size=25, length=1829, TE@rnd-5 family-767@DNA_Maverick@1714:1-1829

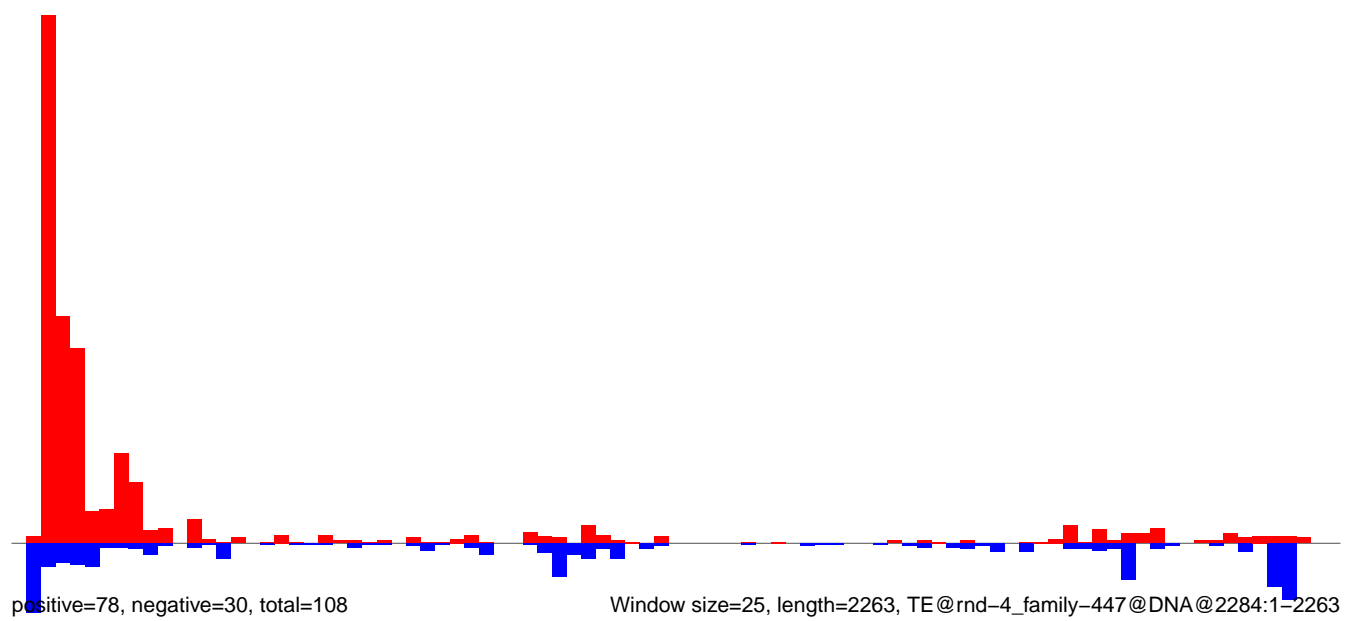
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

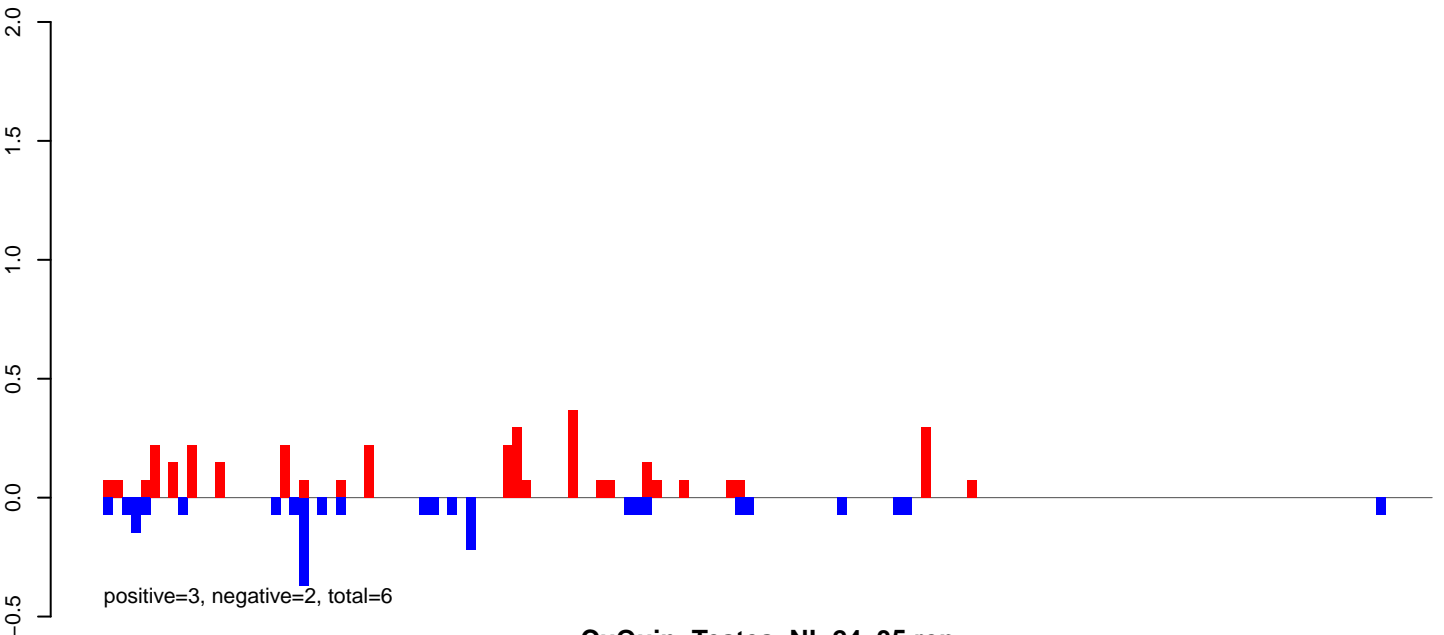


CuQuin_Testes_NL.rep

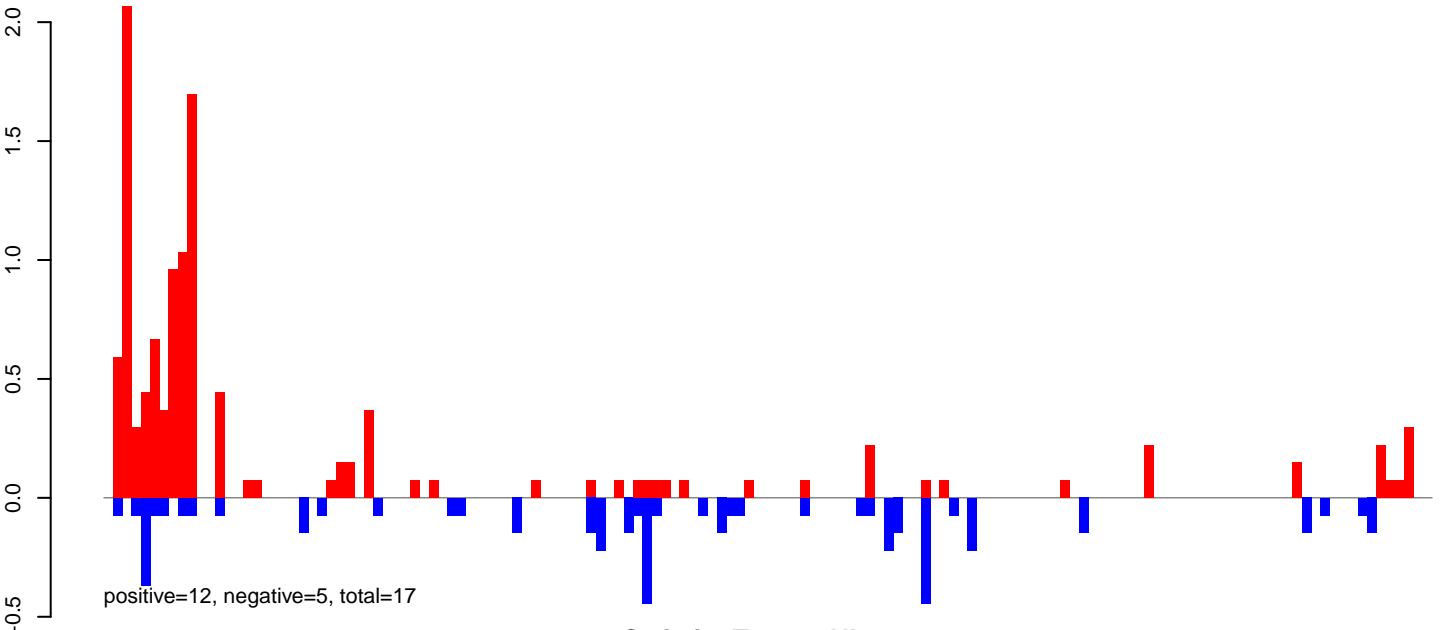


Window size=25, length=2263, TE@rnd-4_family-447@DNA@2284:1-2263

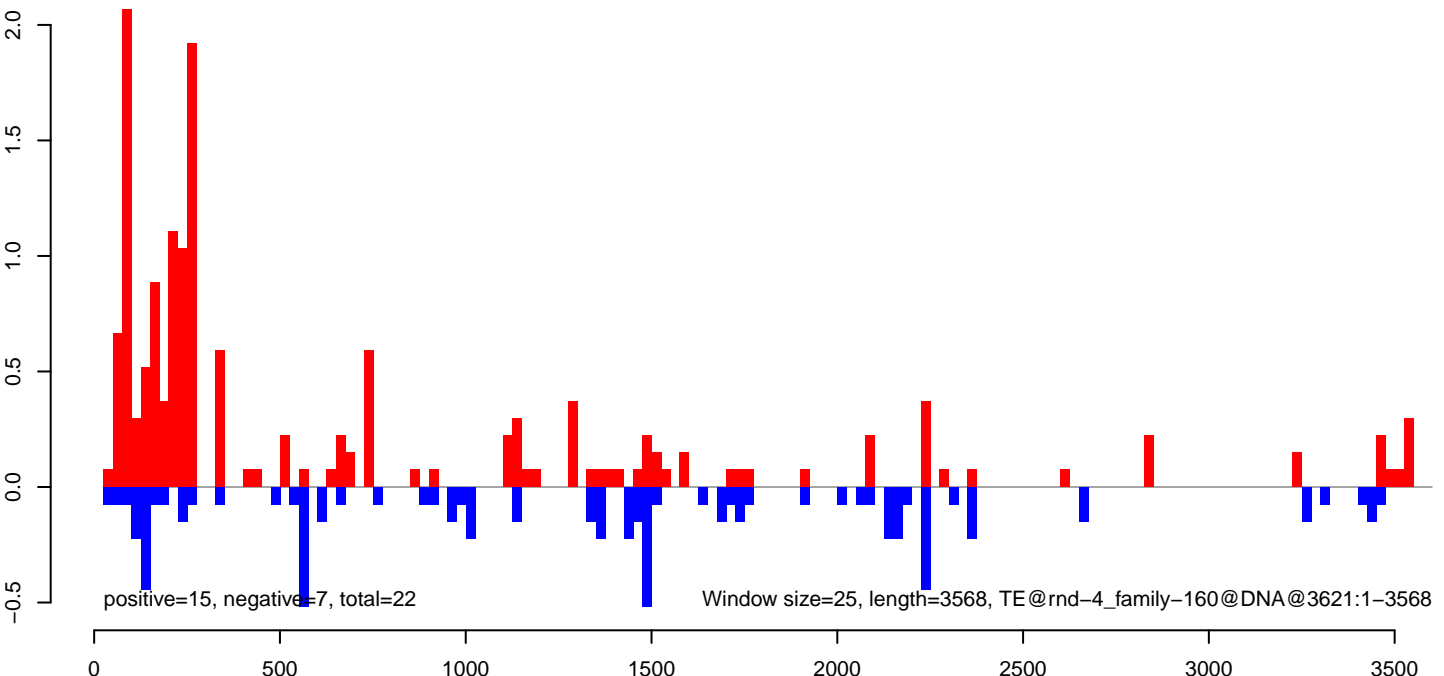
CuQuin_Testes_NL.18_23.rep



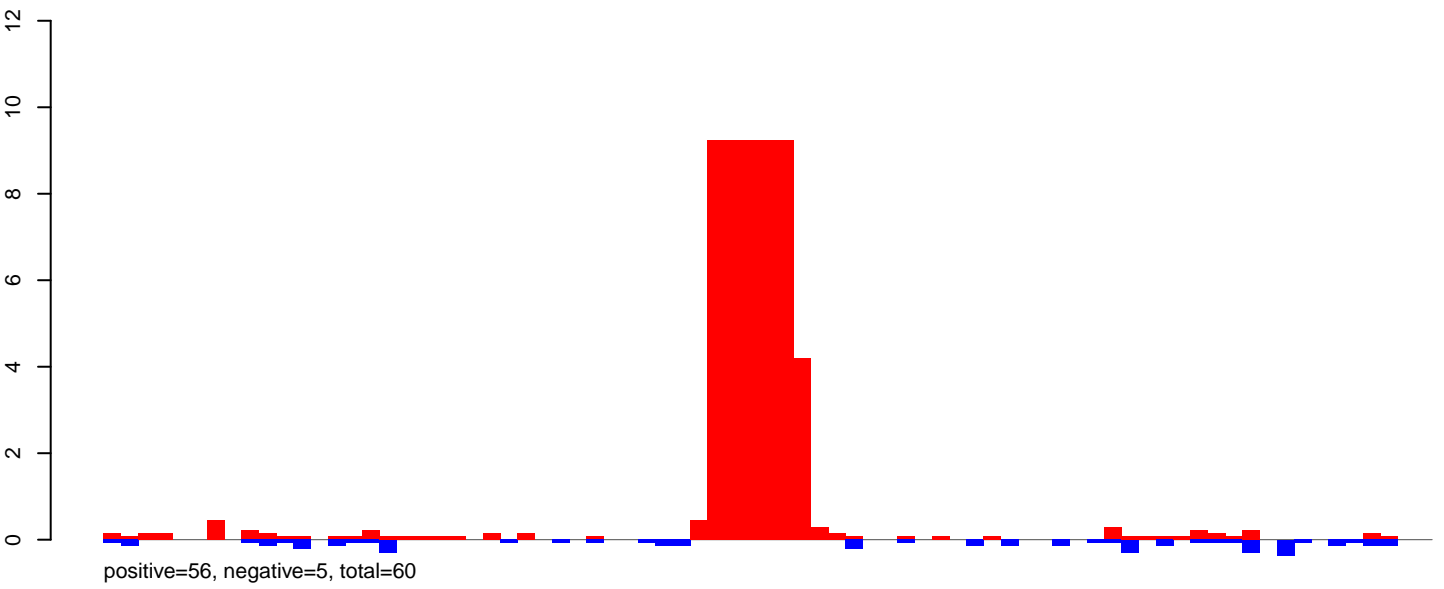
CuQuin_Testes_NL.24_35.rep



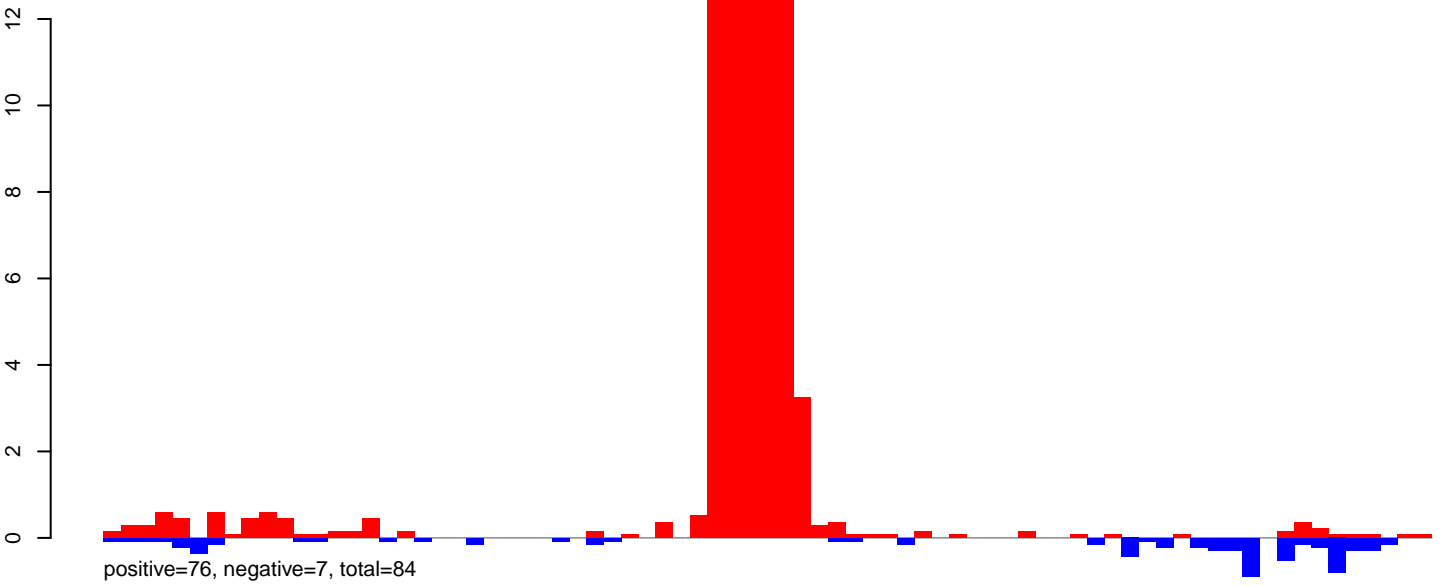
CuQuin_Testes_NL.rep



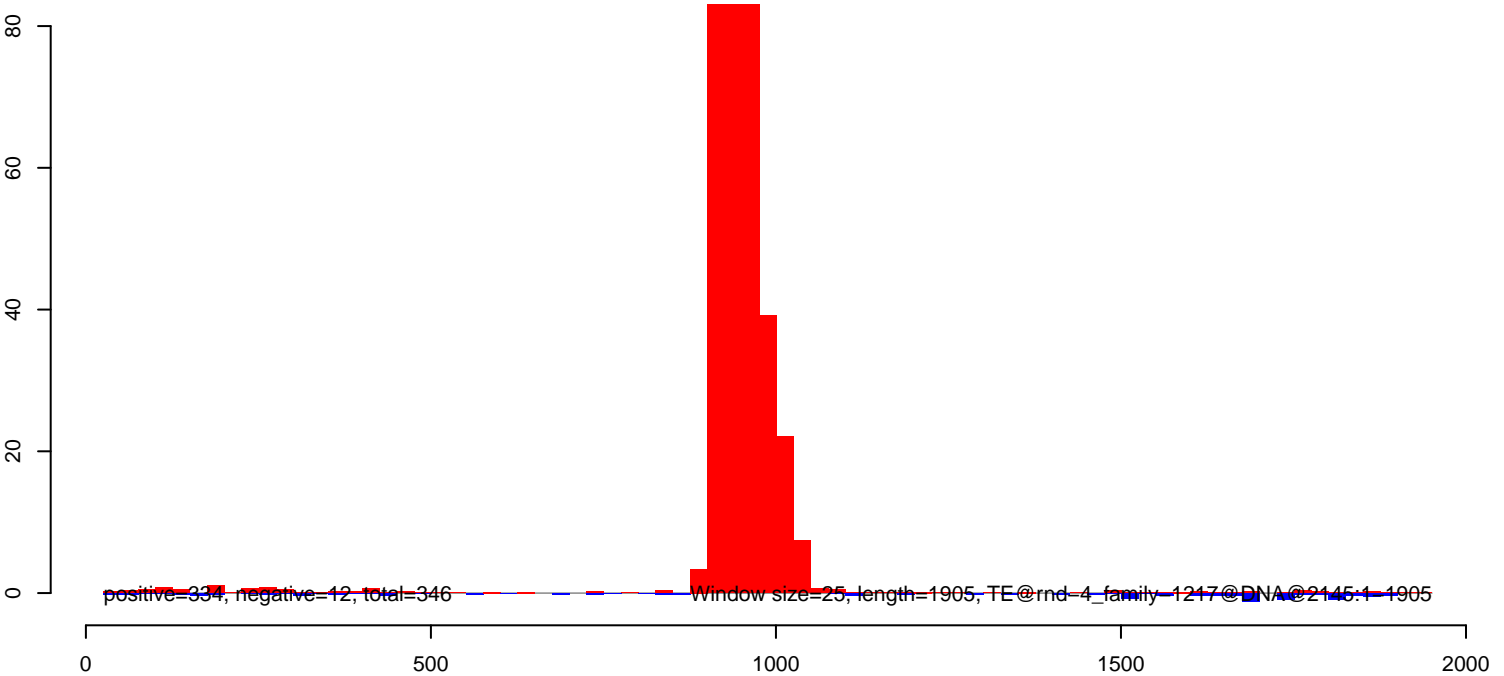
CuQuin_Testes_NL.18_23.rep



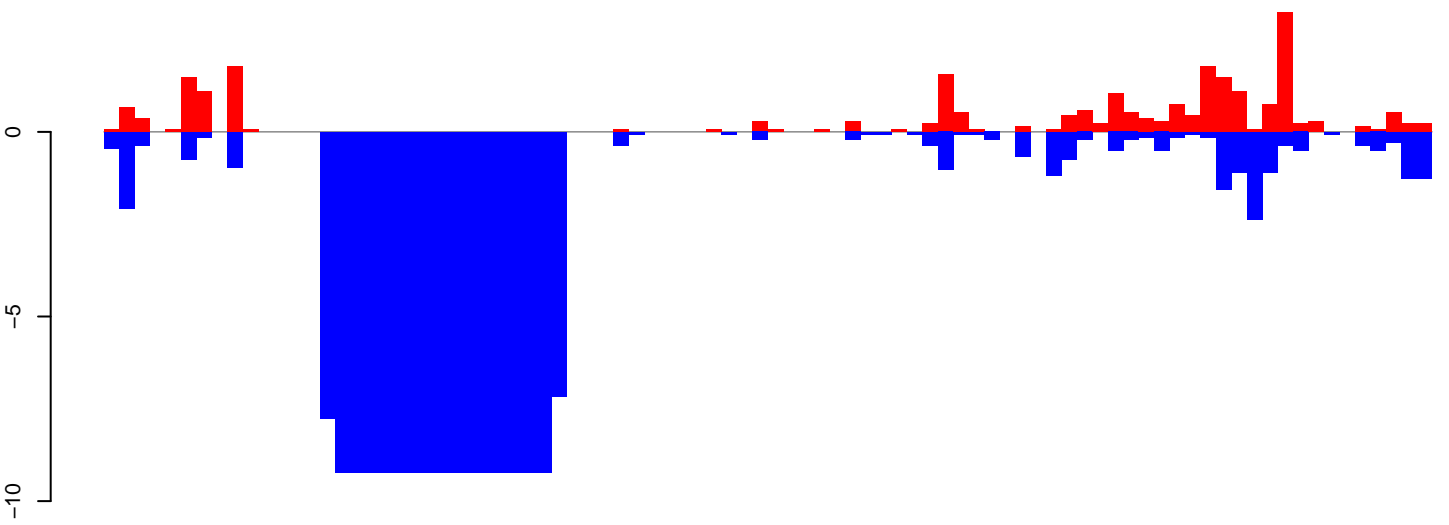
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

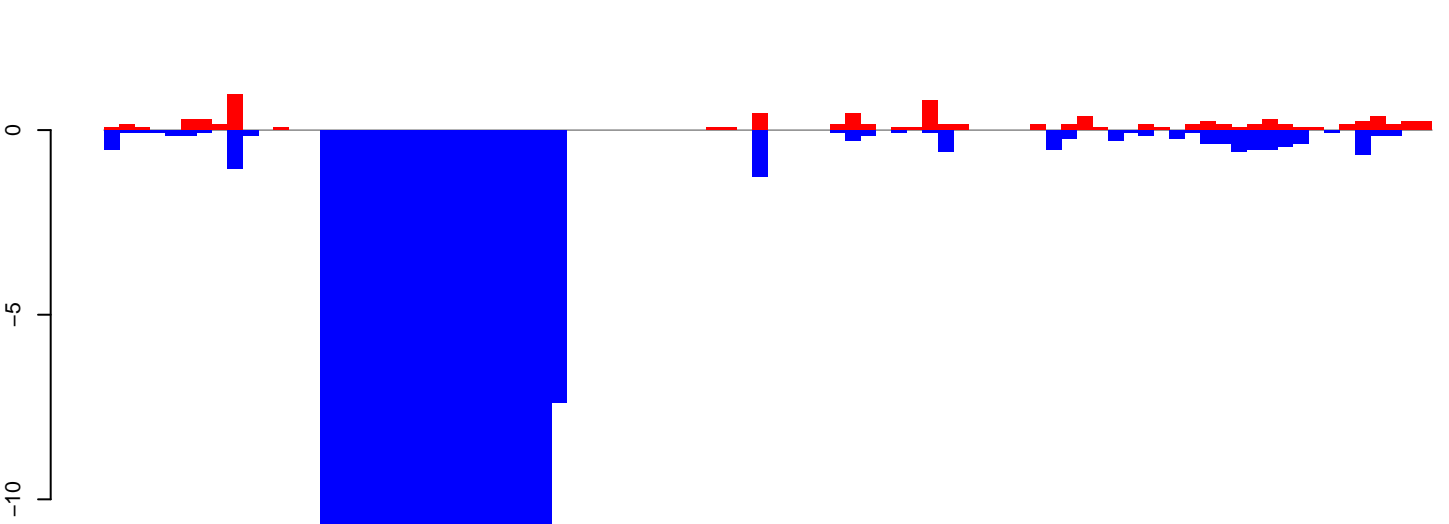


CuQuin_Testes_NL.18_23.rep



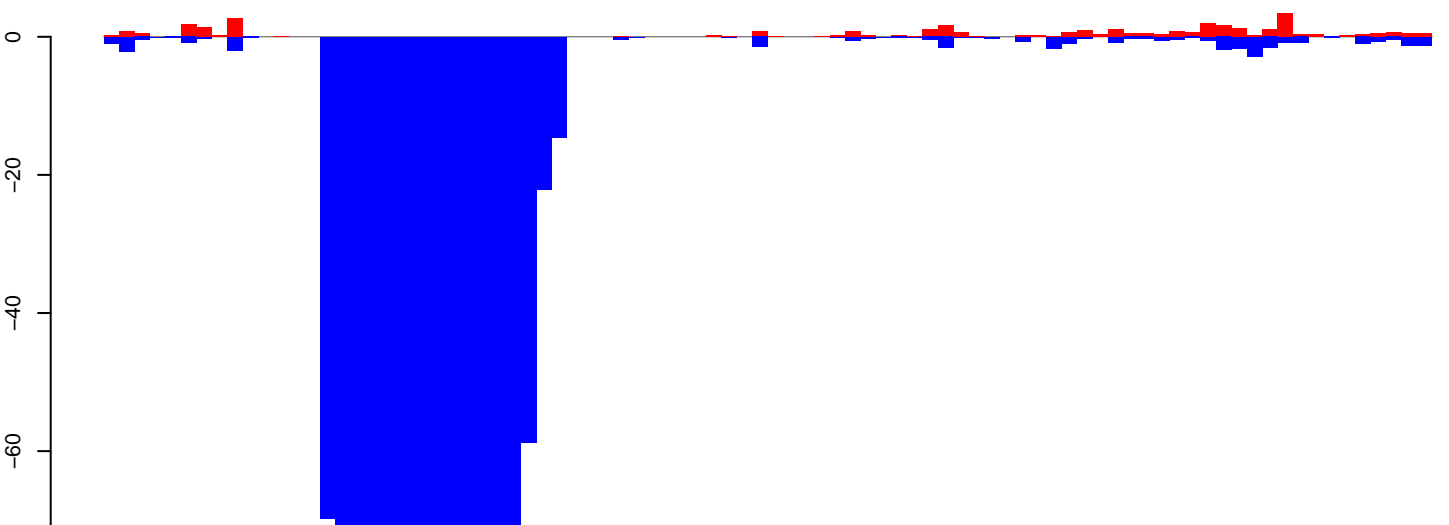
positive=24, negative=167, total=191

CuQuin_Testes_NL.24_35.rep



positive=8, negative=210, total=218

CuQuin_Testes_NL.rep

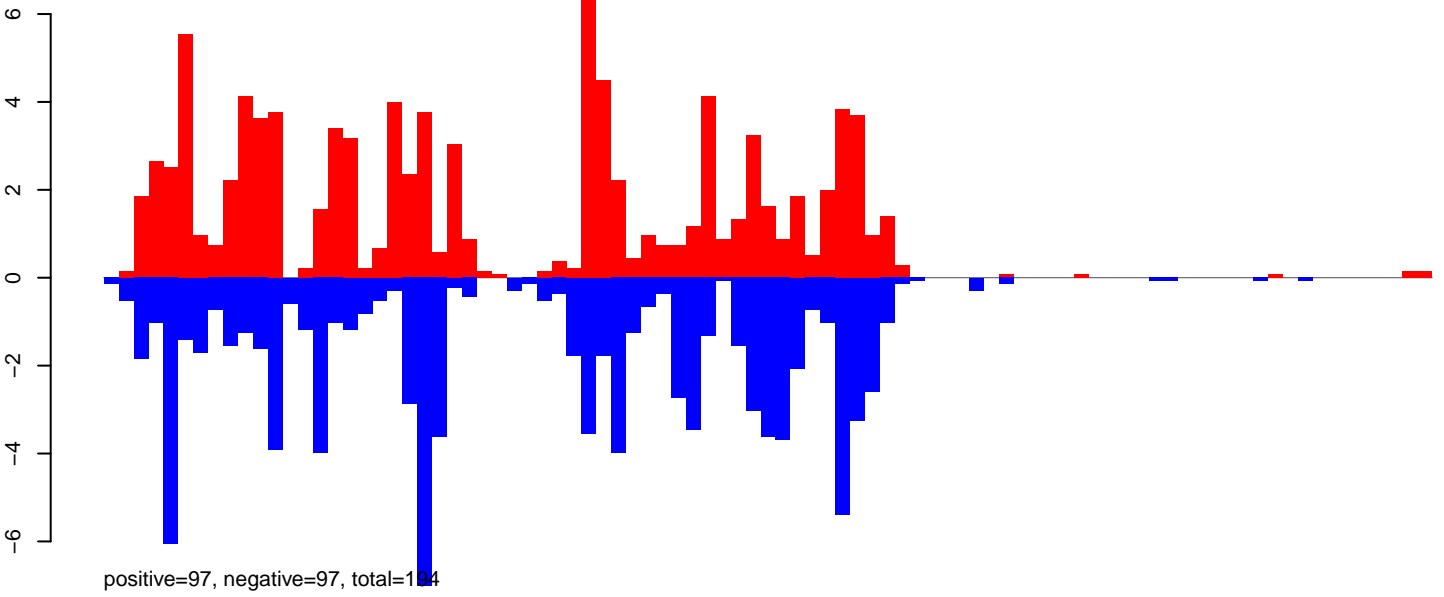


positive=32, negative=1196, total=1228

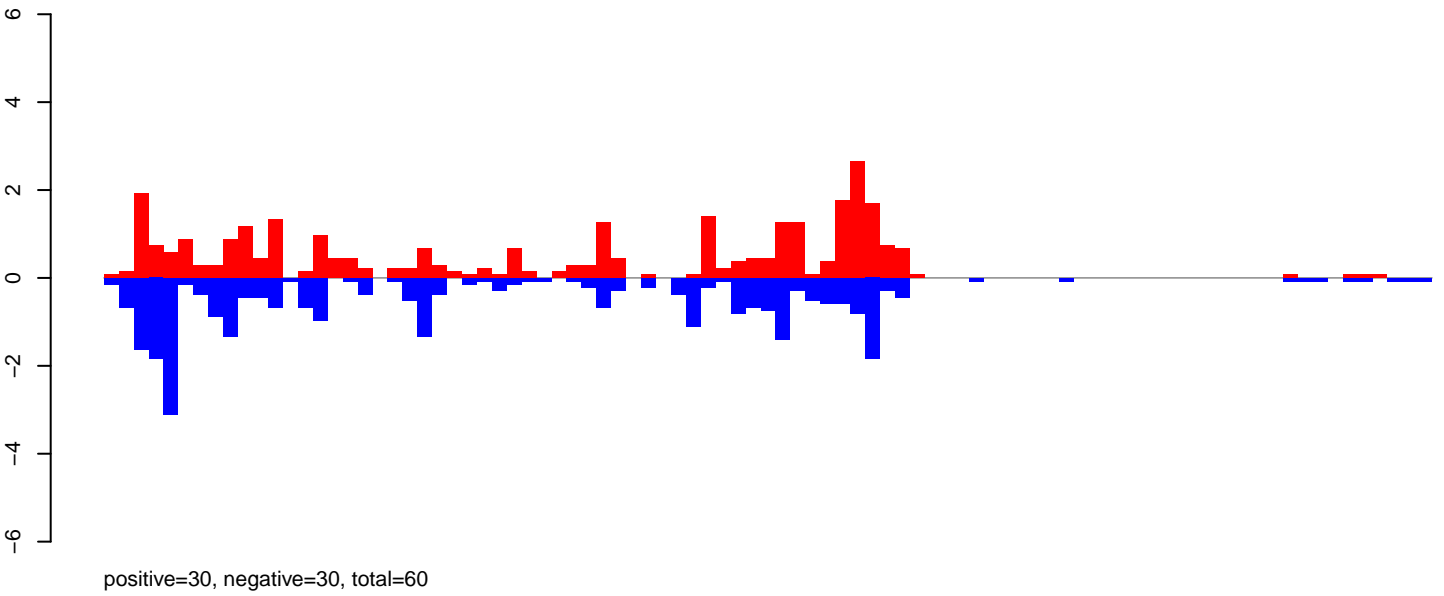
Window size=25, length=2130, TE@rnd-4_family-100@DNA_Sola-2@2037:1-2130

0 500 1000 1500 2000

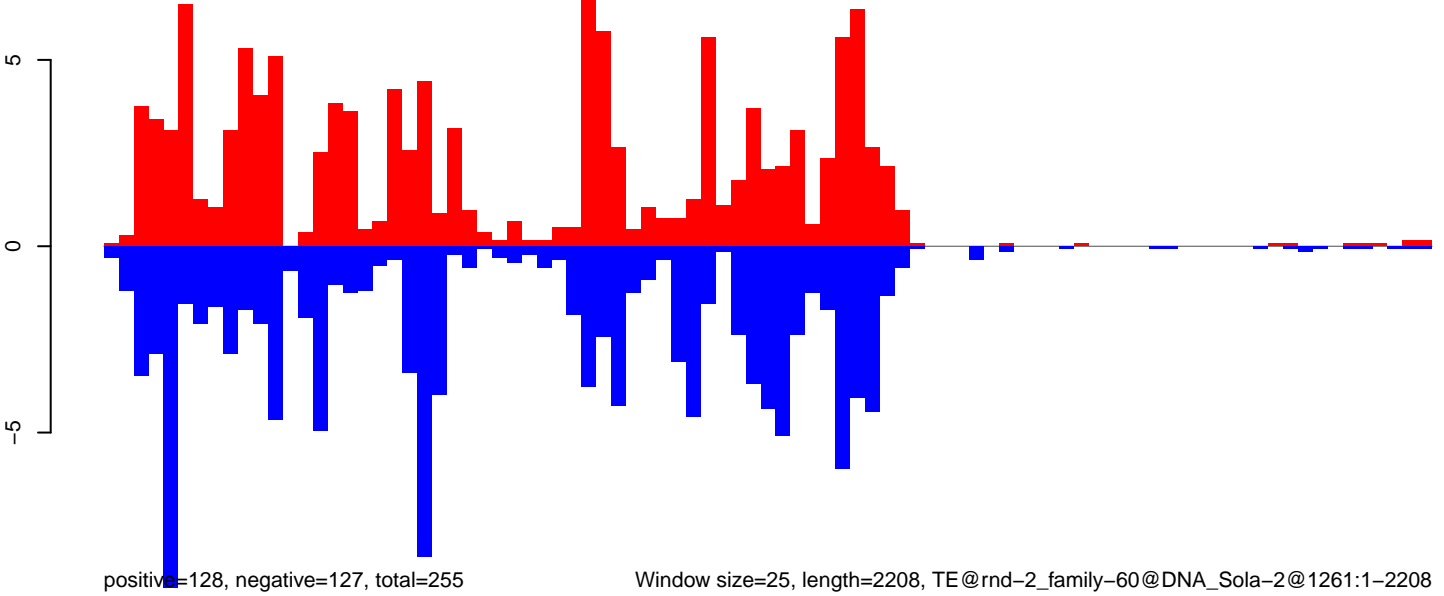
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



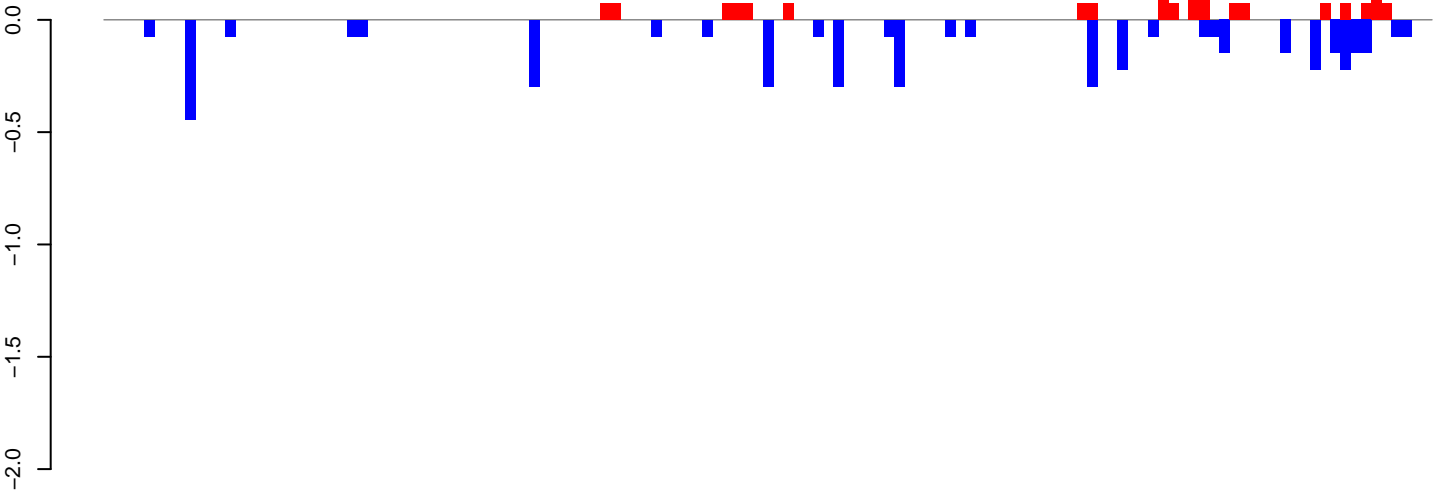
CuQuin_Testes_NL.rep



Window size=25, length=2208, TE@rnd-2_family-60@DNA_Sola-2@1261:1-2208

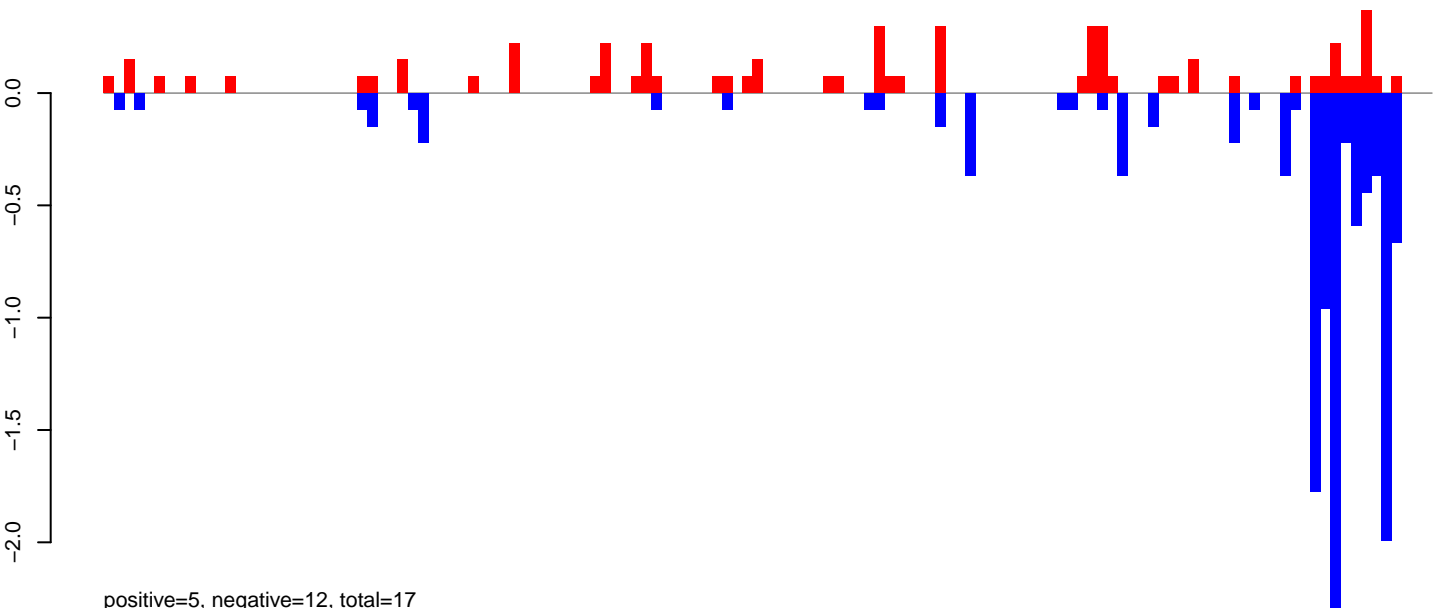
0 500 1000 1500 2000

CuQuin_Testes_NL.18_23.rep



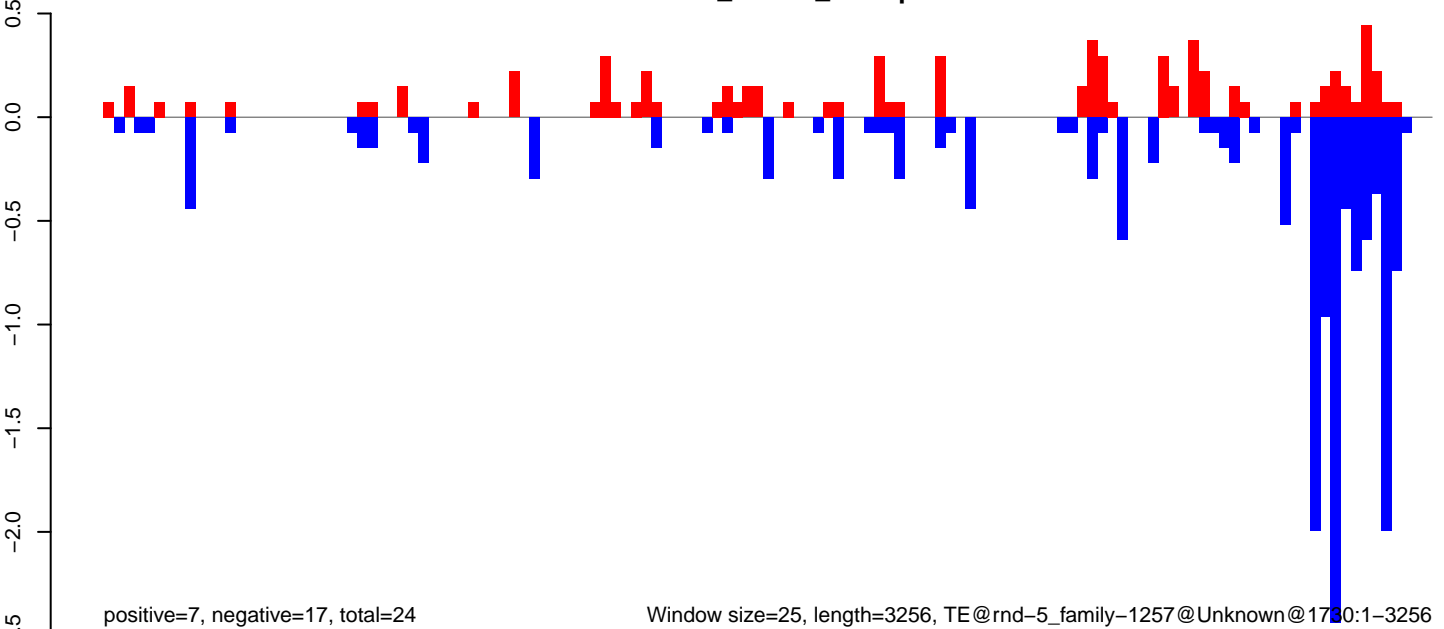
positive=2, negative=4, total=6

CuQuin_Testes_NL.24_35.rep



positive=5, negative=12, total=17

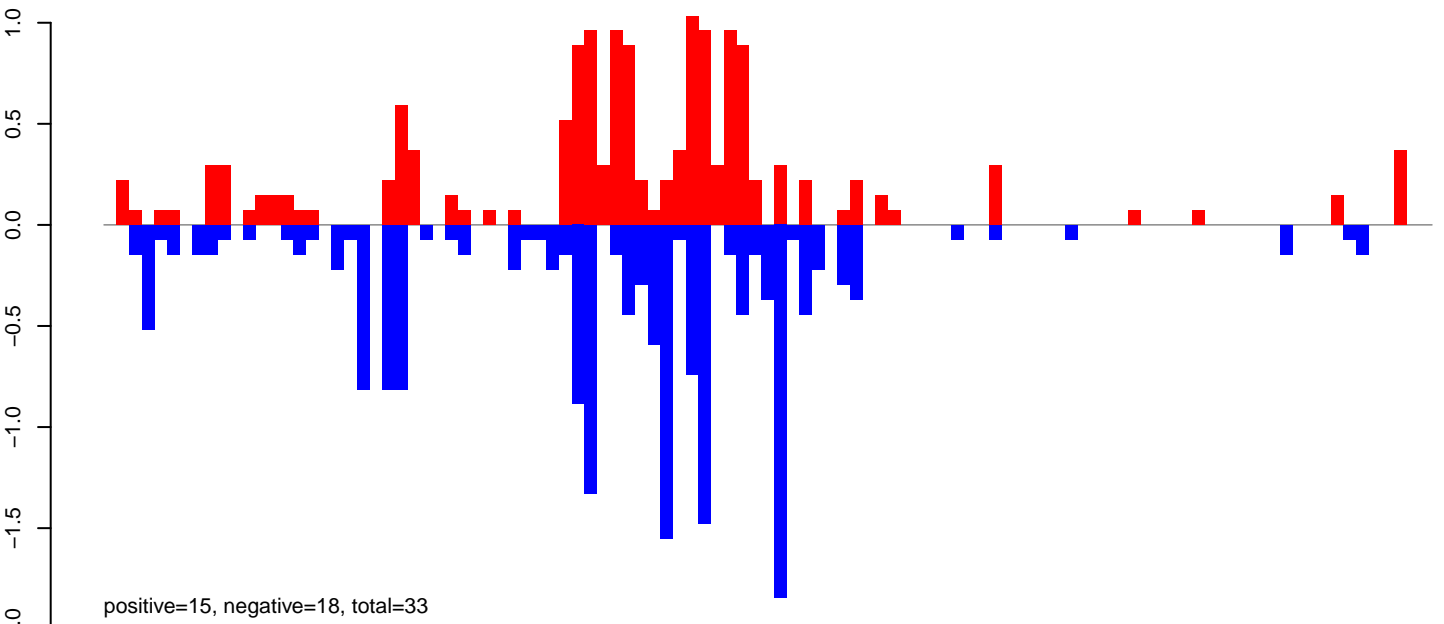
CuQuin_Testes_NL.rep



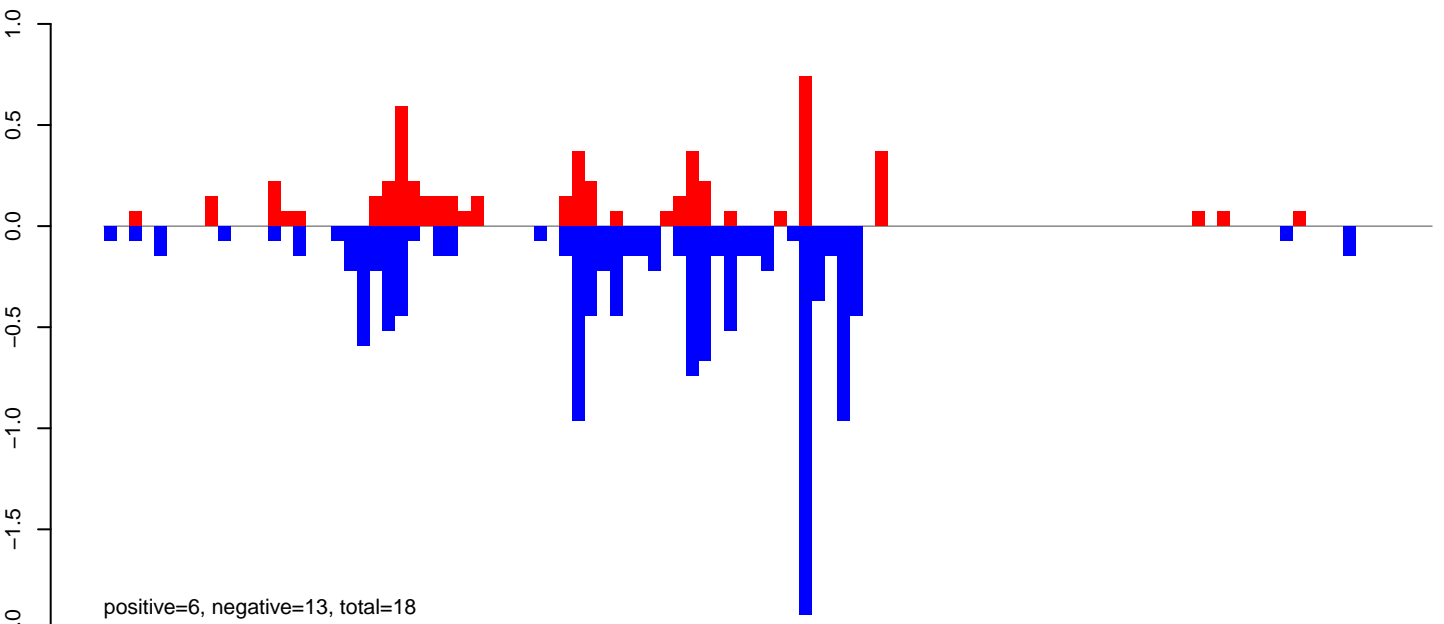
positive=7, negative=17, total=24

Window size=25, length=3256, TE@rnd-5_family-1257@Unknown@1730:1-3256

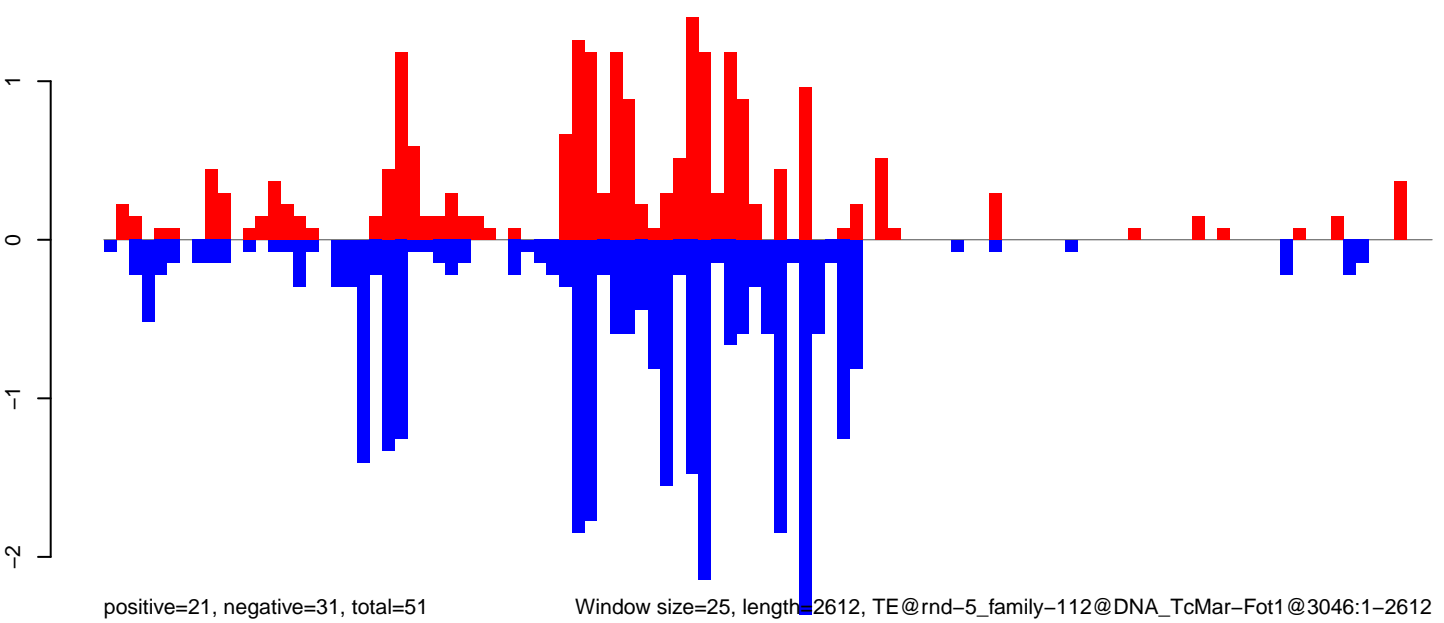
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

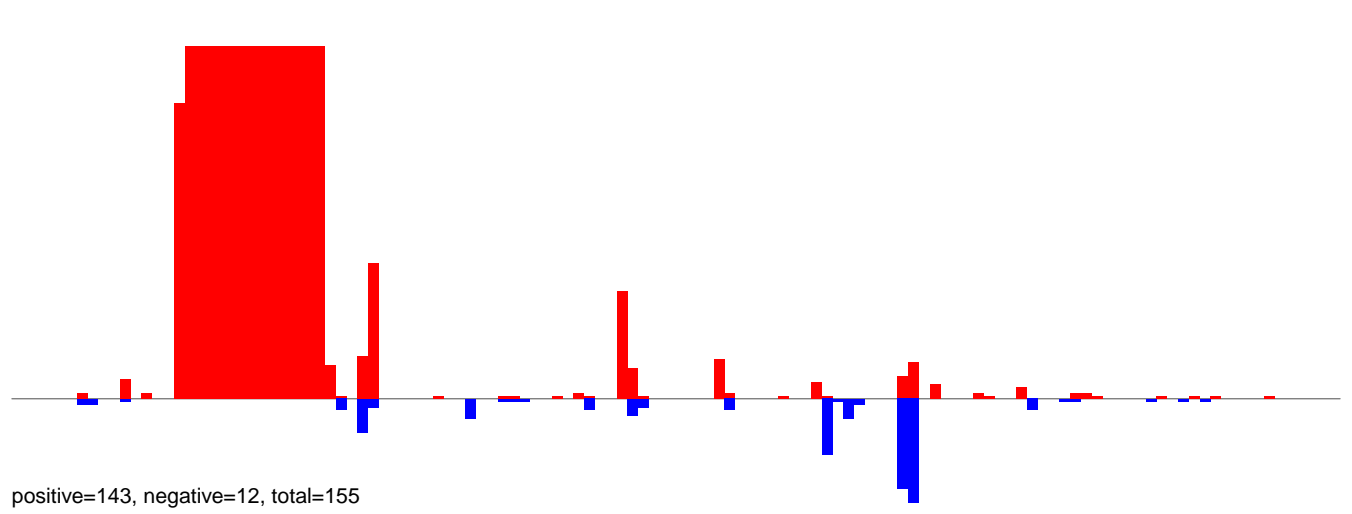


CuQuin_Testes_NL.rep

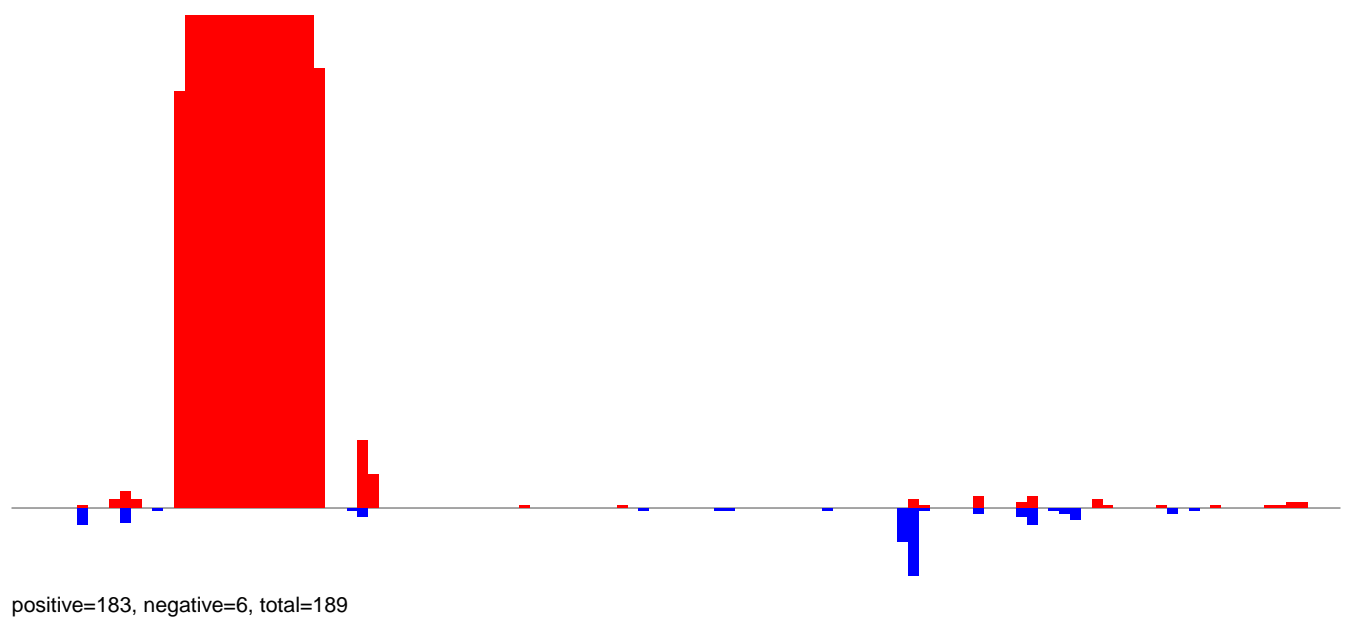


0 500 1000 1500 2000 2500

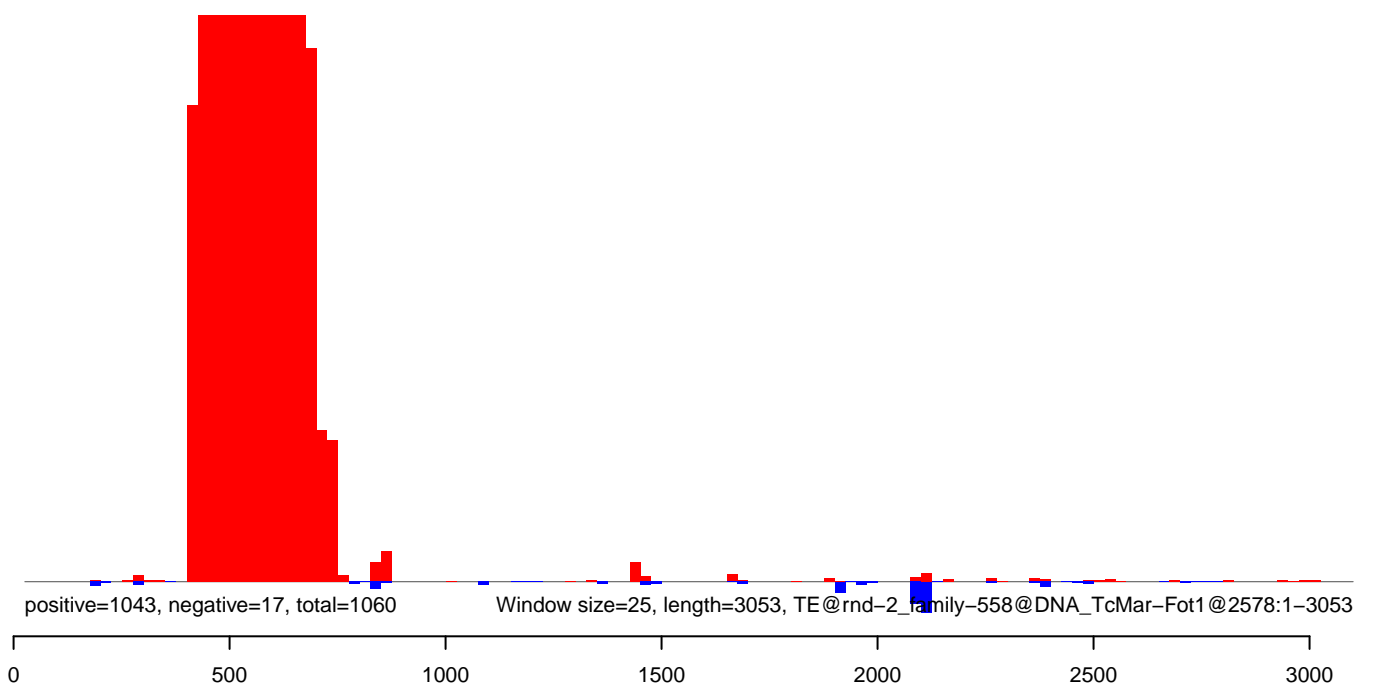
CuQuin_Testes_NL.18_23.rep



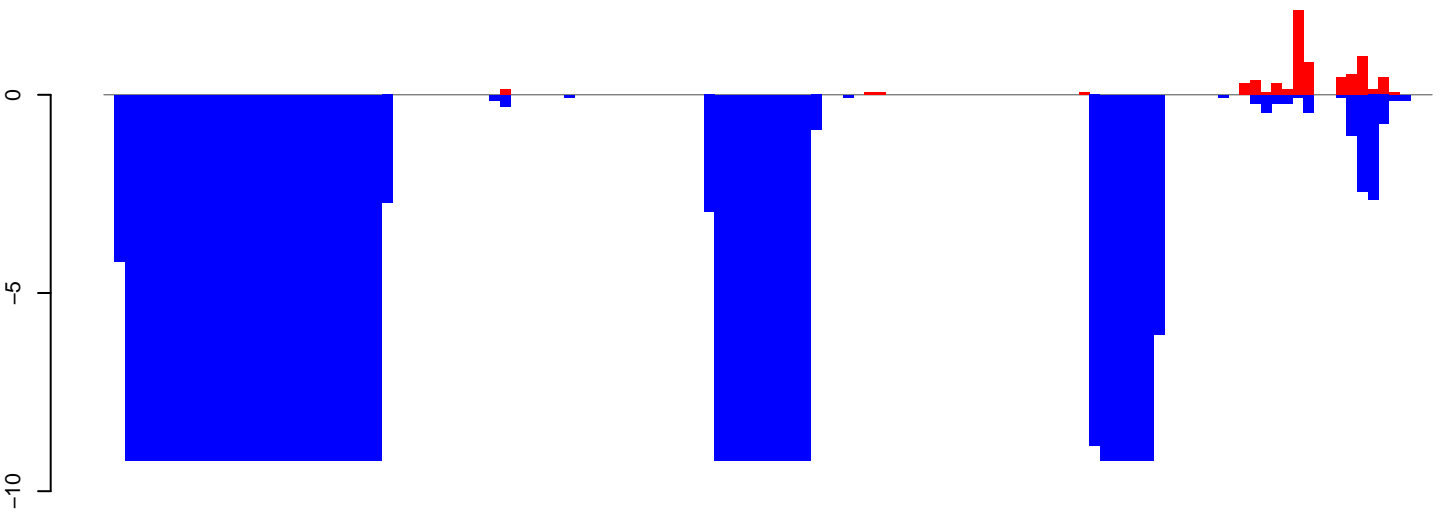
CuQuin_Testes_NL.24_35.rep



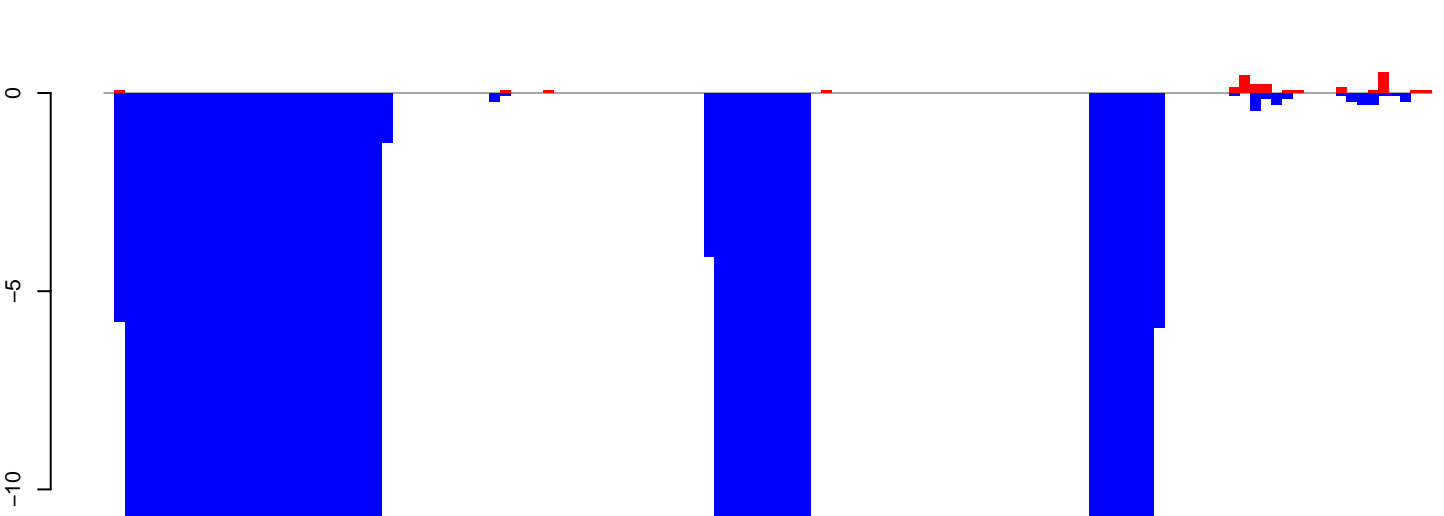
CuQuin_Testes_NL.rep



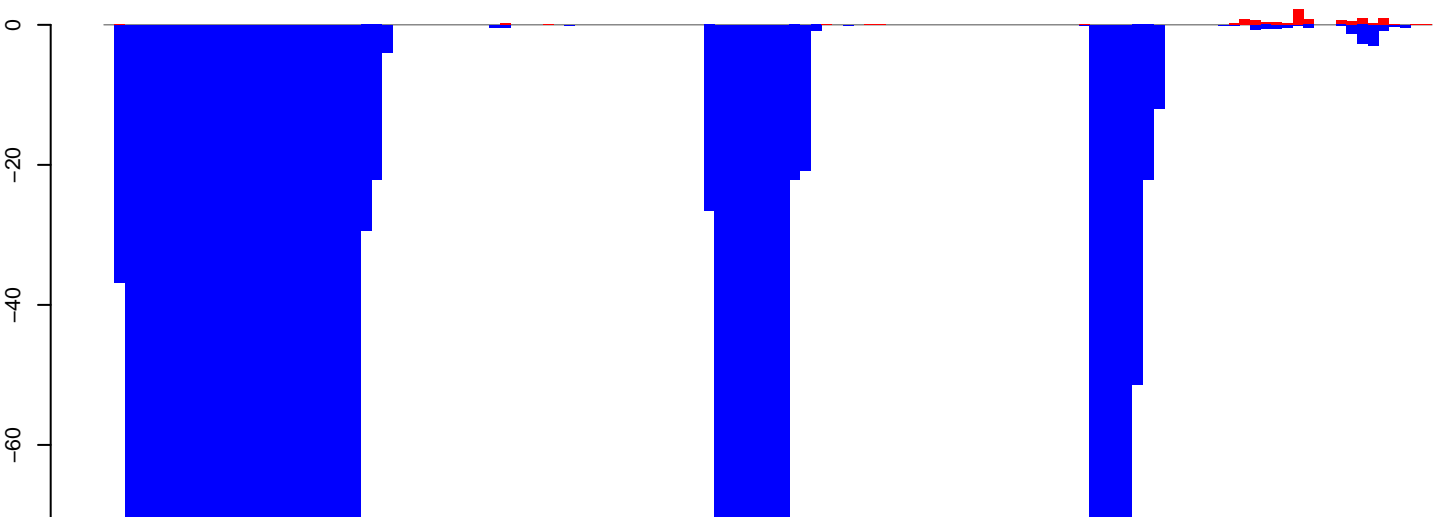
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

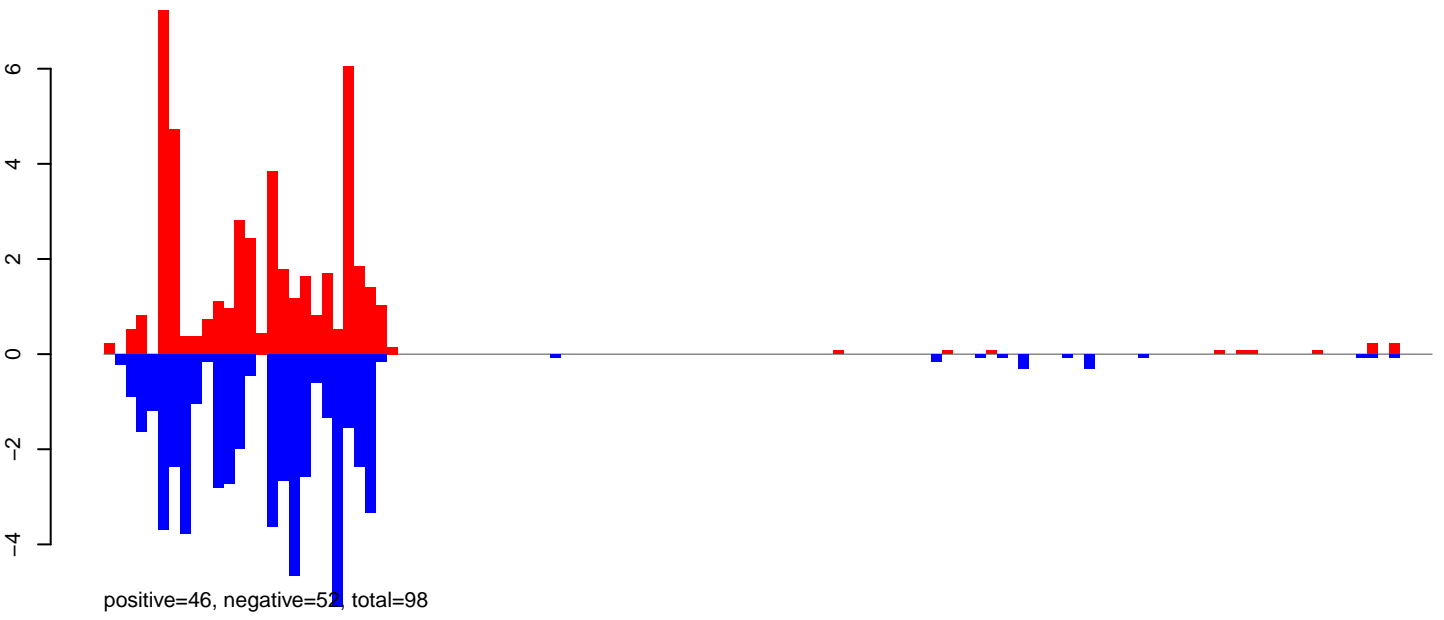


CuQuin_Testes_NL.rep

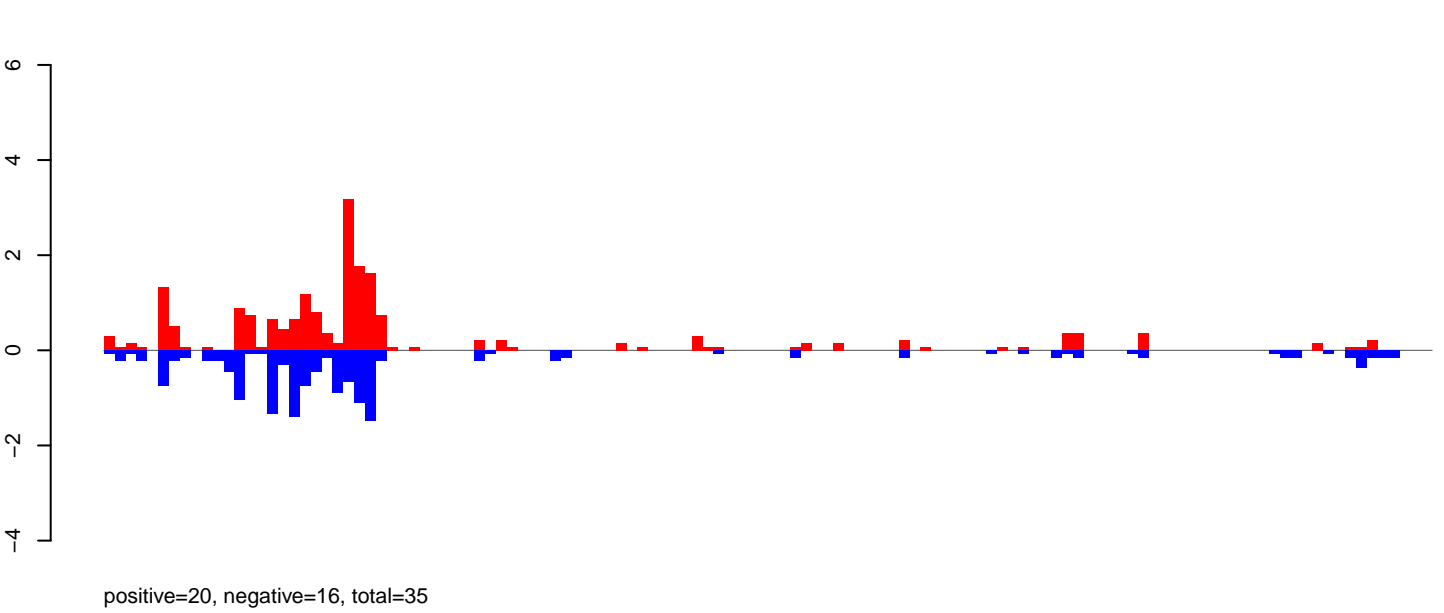


Window size=25, length=3089, TE@rnd-2_family-302@DNA_Zator@2573:1-3089

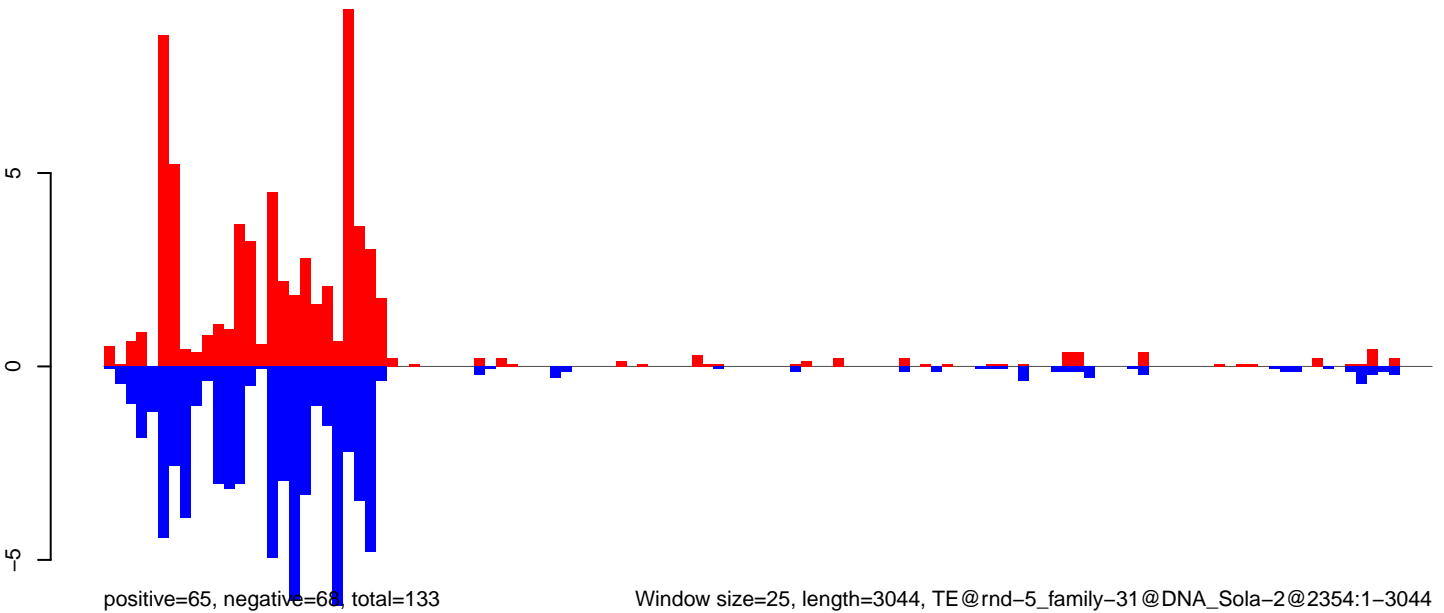
CuQuin_Testes_NL.18_23.rep



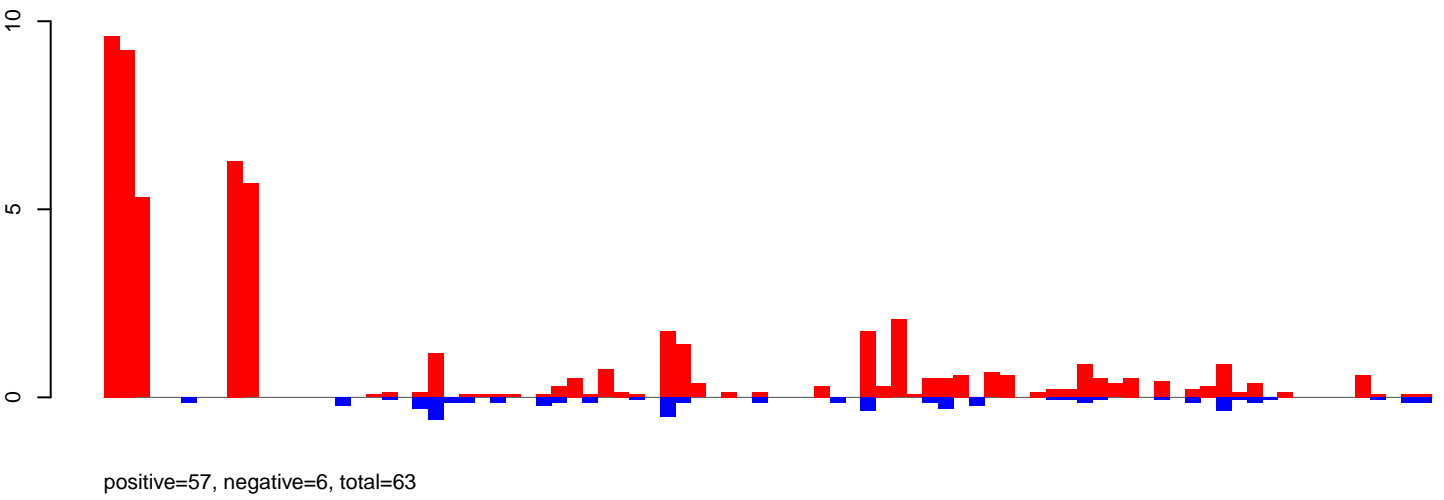
CuQuin_Testes_NL.24_35.rep



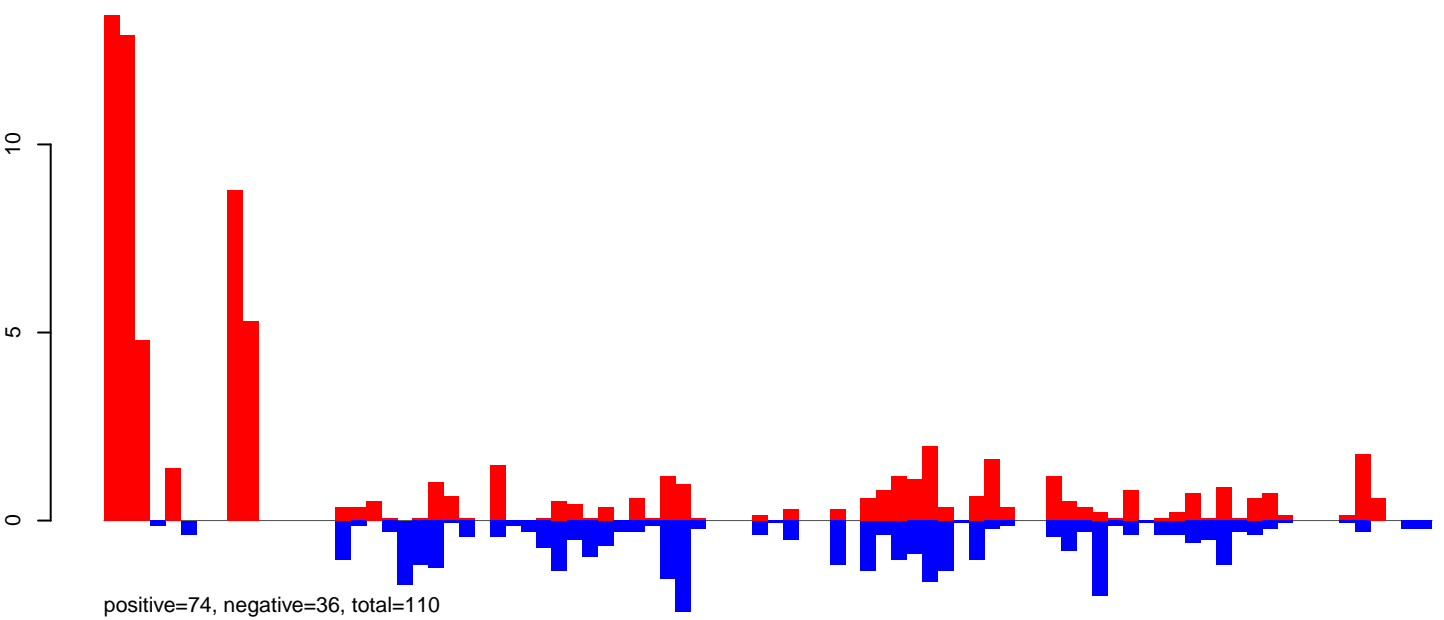
CuQuin_Testes_NL.rep



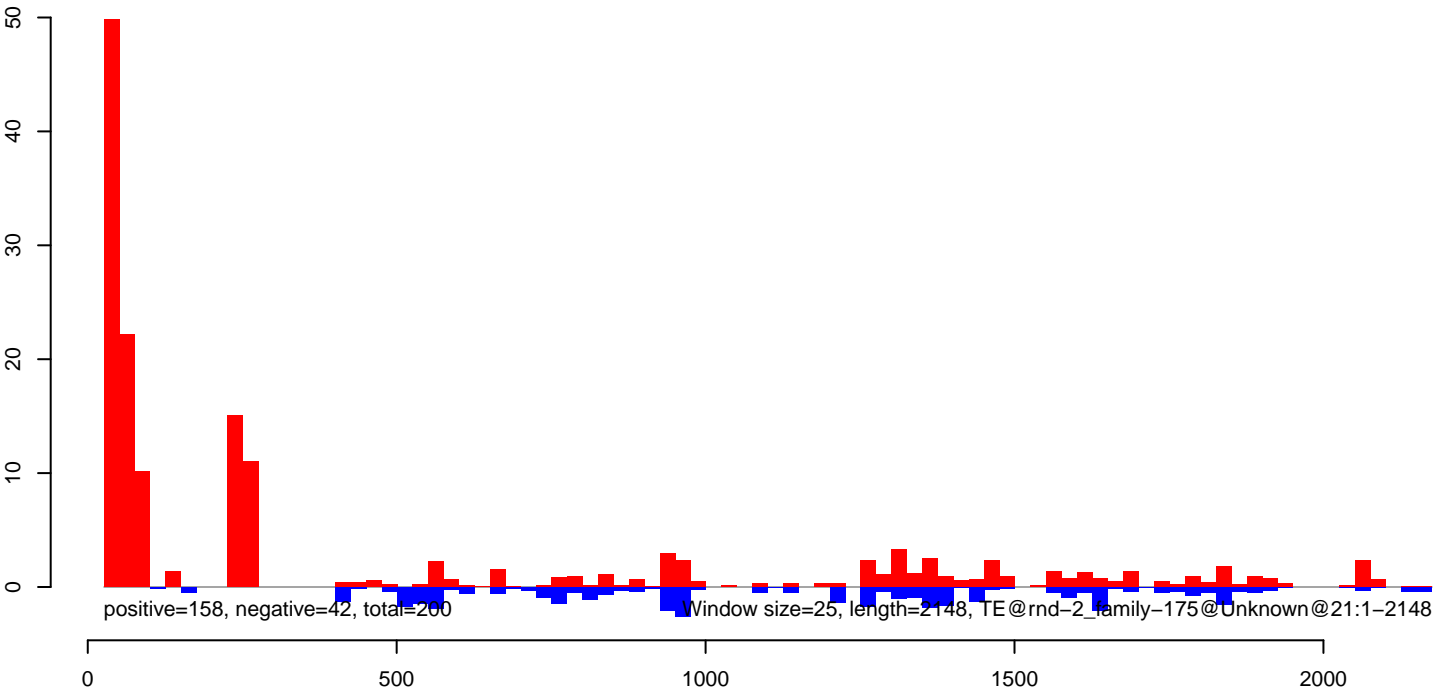
CuQuin_Testes_NL.18_23.rep



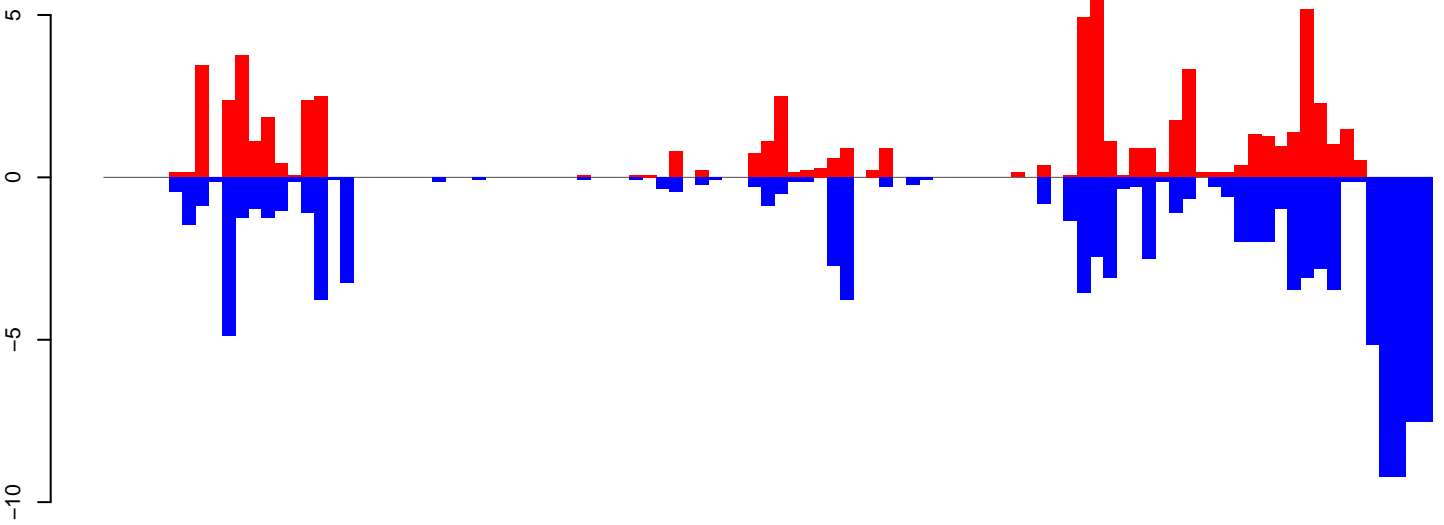
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

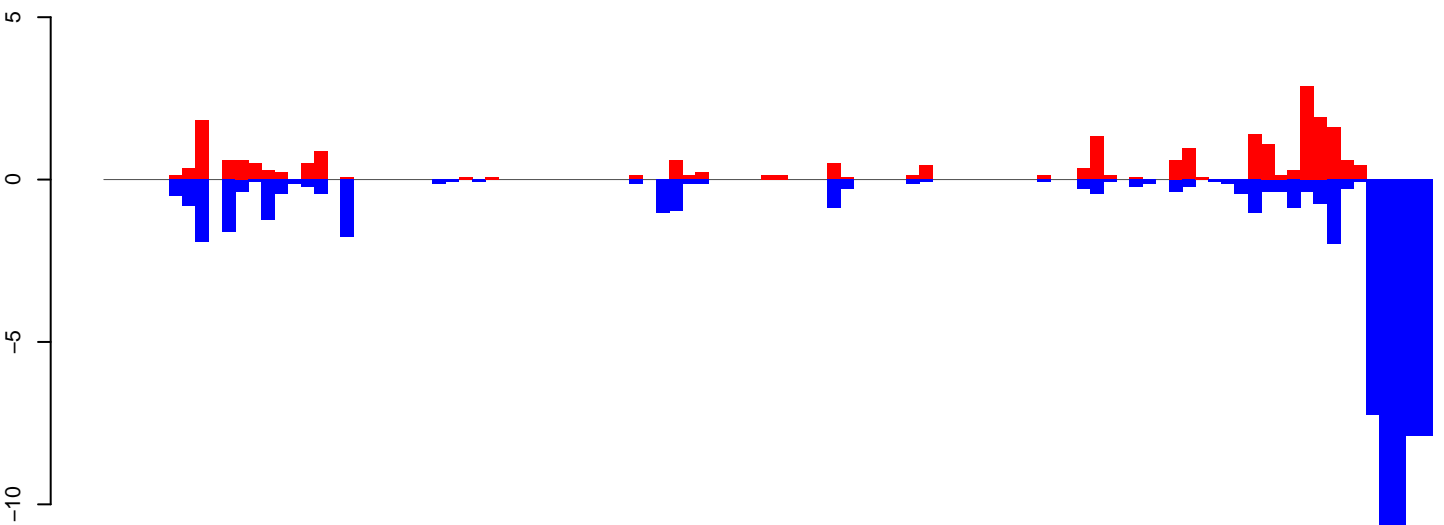


CuQuin_Testes_NL.18_23.rep



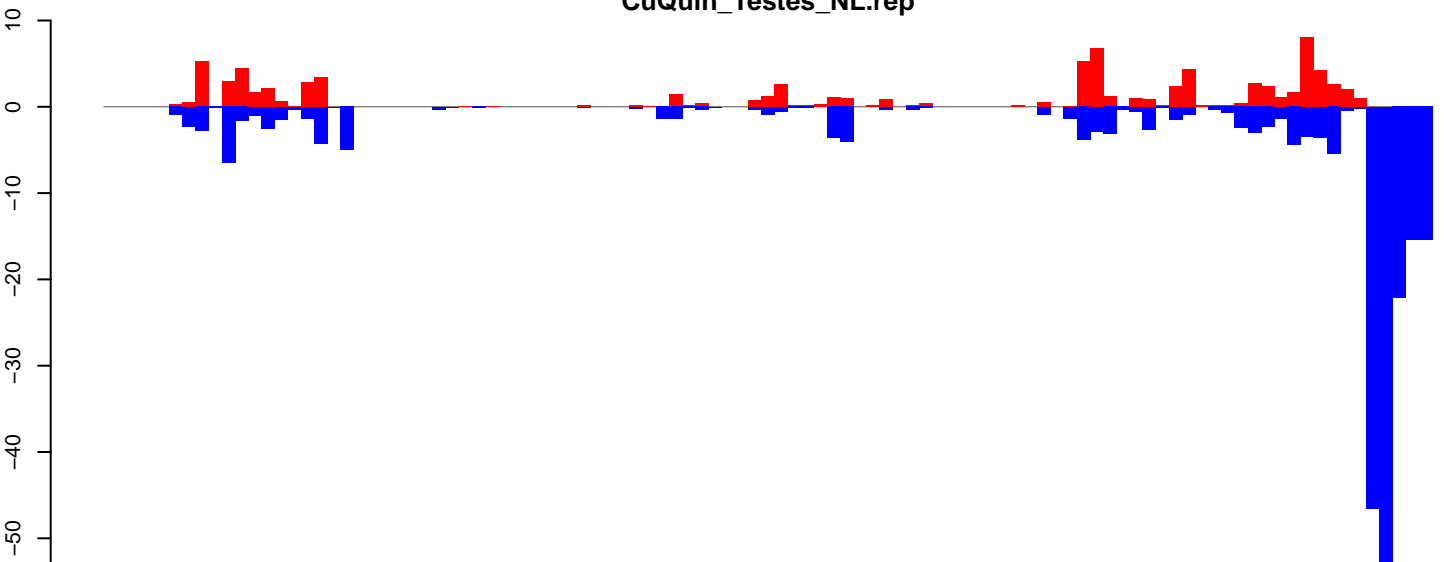
positive=63, negative=107, total=170

CuQuin_Testes_NL.24_35.rep



positive=23, negative=71, total=94

CuQuin_Testes_NL.rep

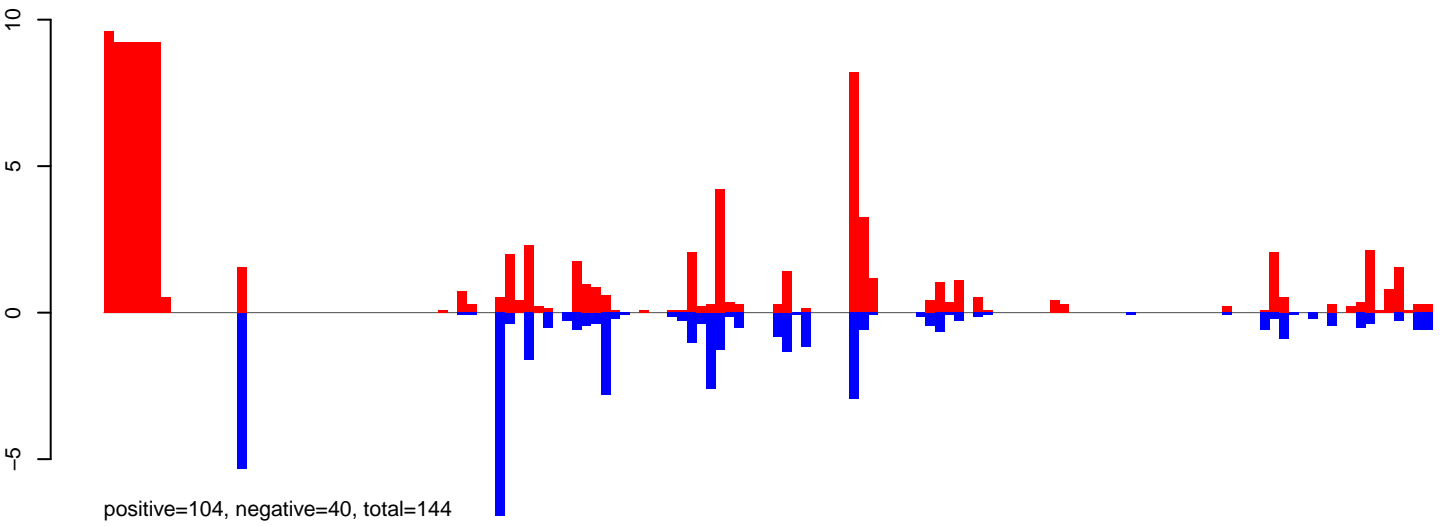


positive=86, negative=252, total=337

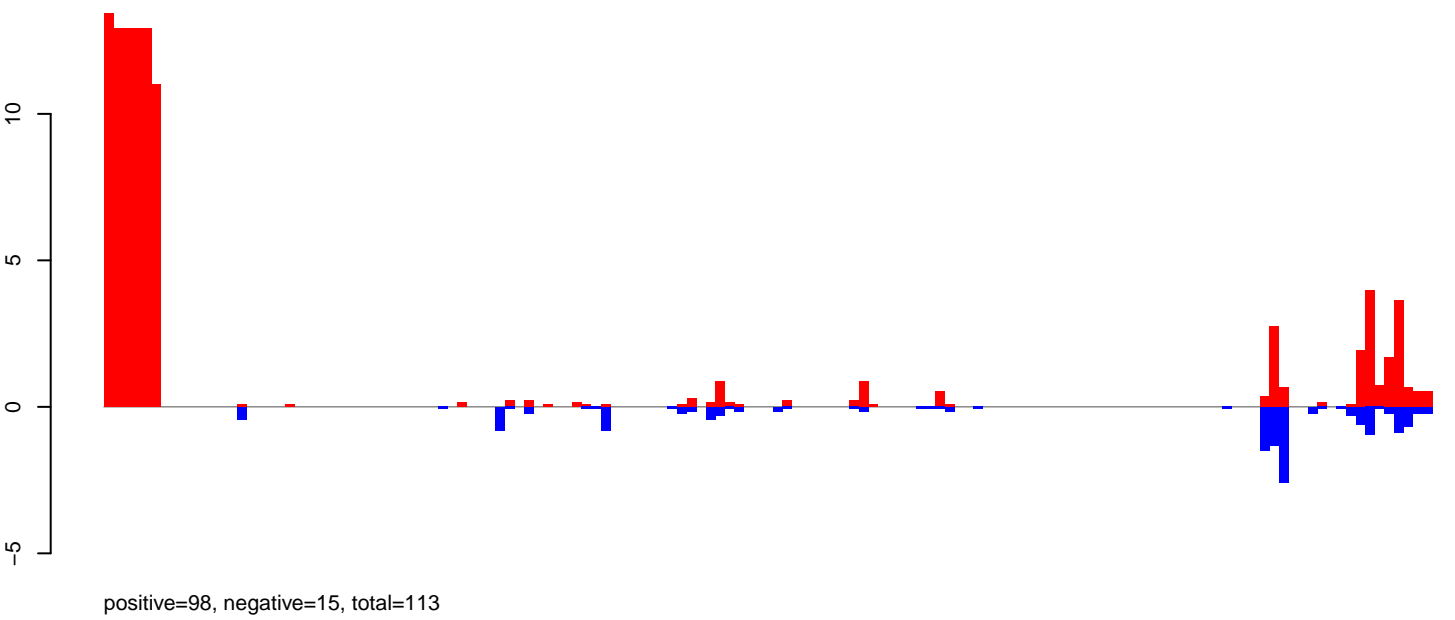
Window size=25, length=2517, TE@rnd-5_family-254@DNA_Sola-2@553:1-2517

0 500 1000 1500 2000 2500

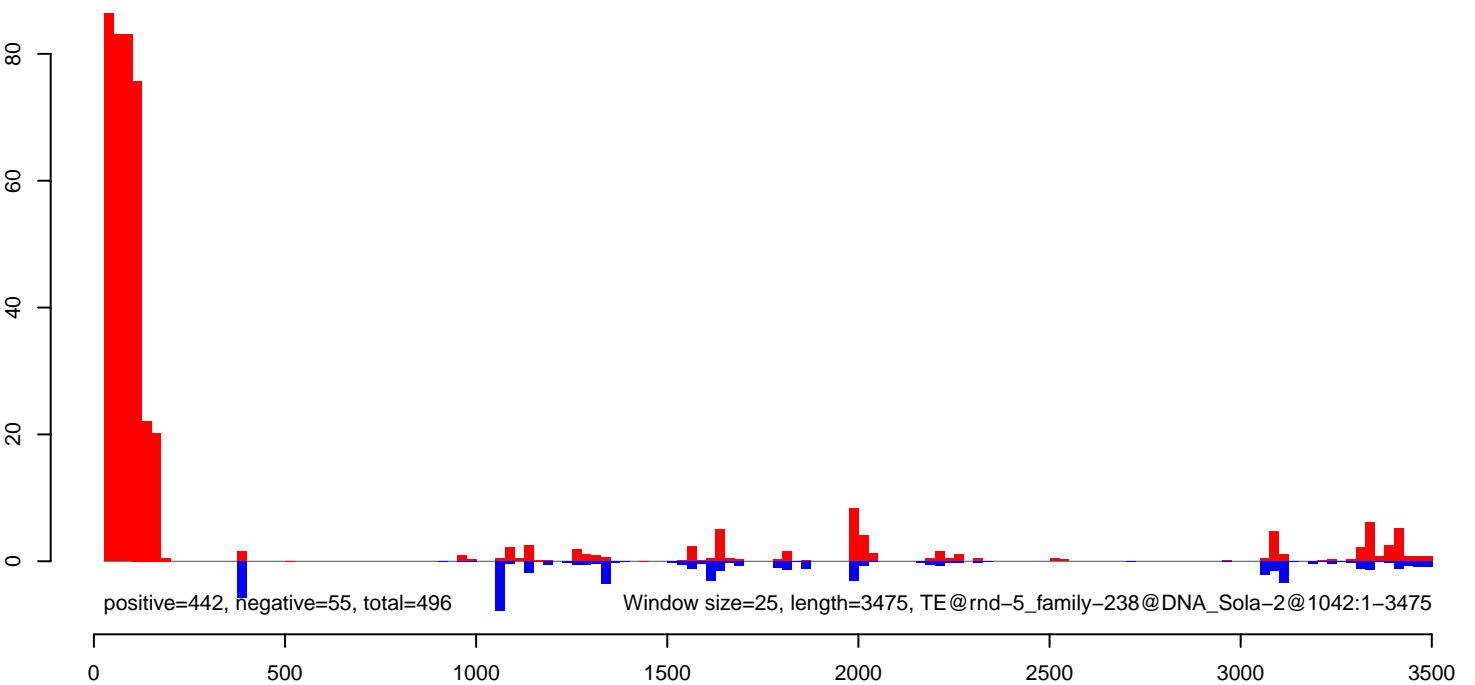
CuQuin_Testes_NL.18_23.rep



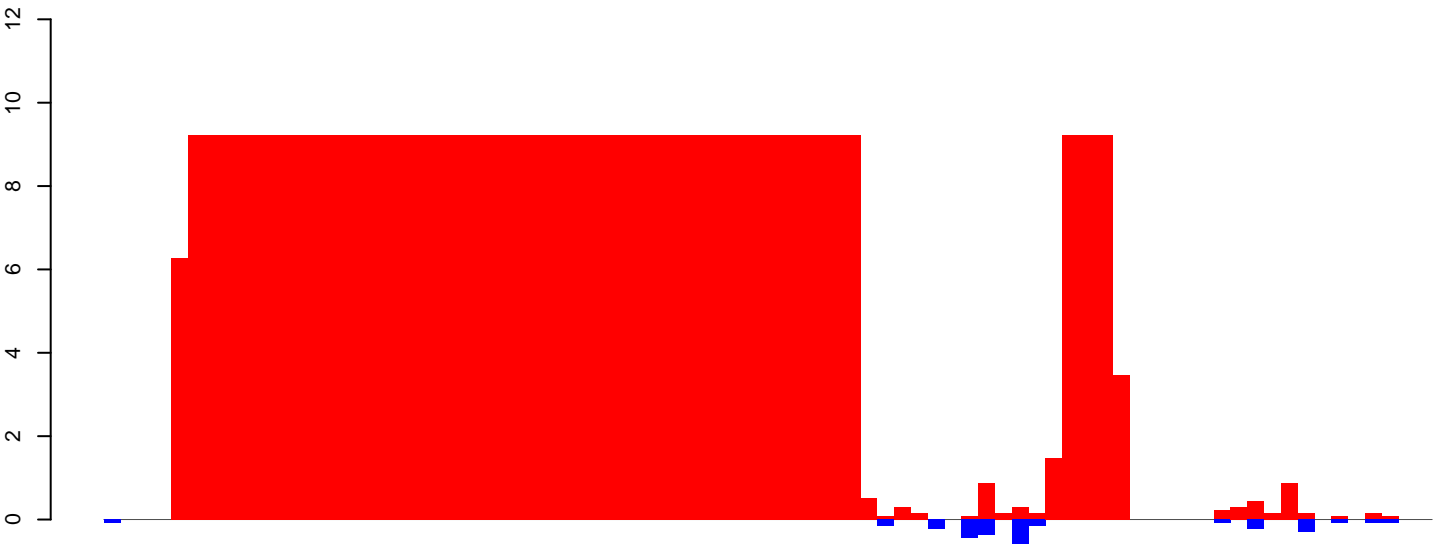
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

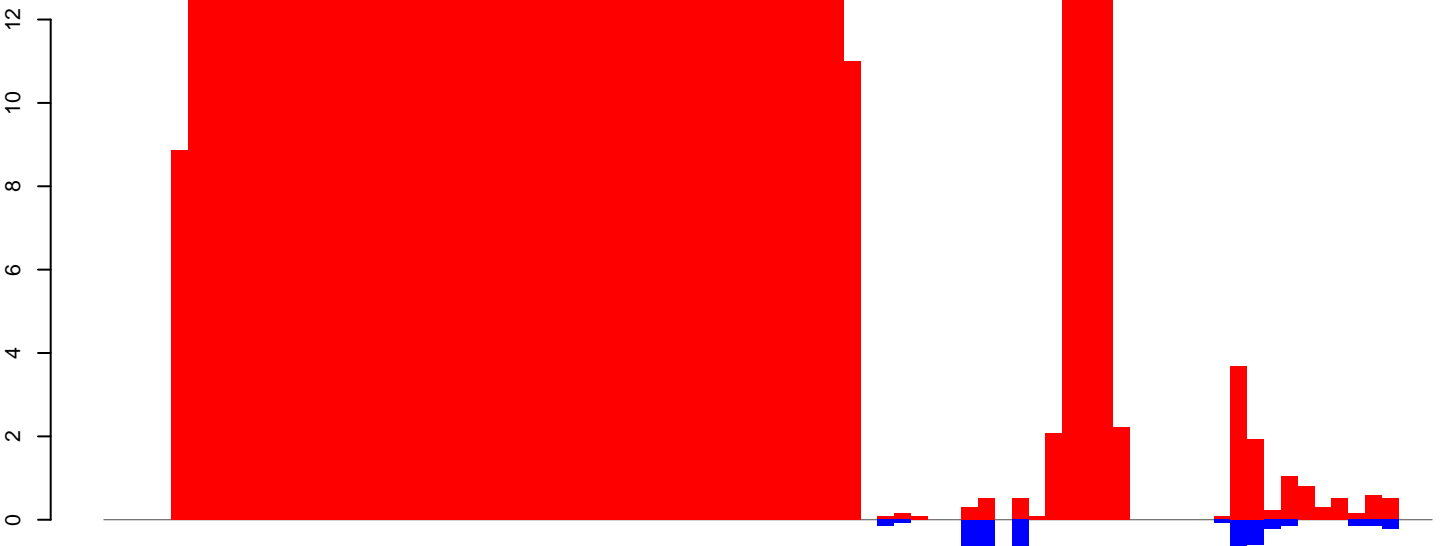


CuQuin_Testes_NL.18_23.rep



positive=413, negative=3, total=416

CuQuin_Testes_NL.24_35.rep



positive=578, negative=5, total=584

CuQuin_Testes_NL.rep

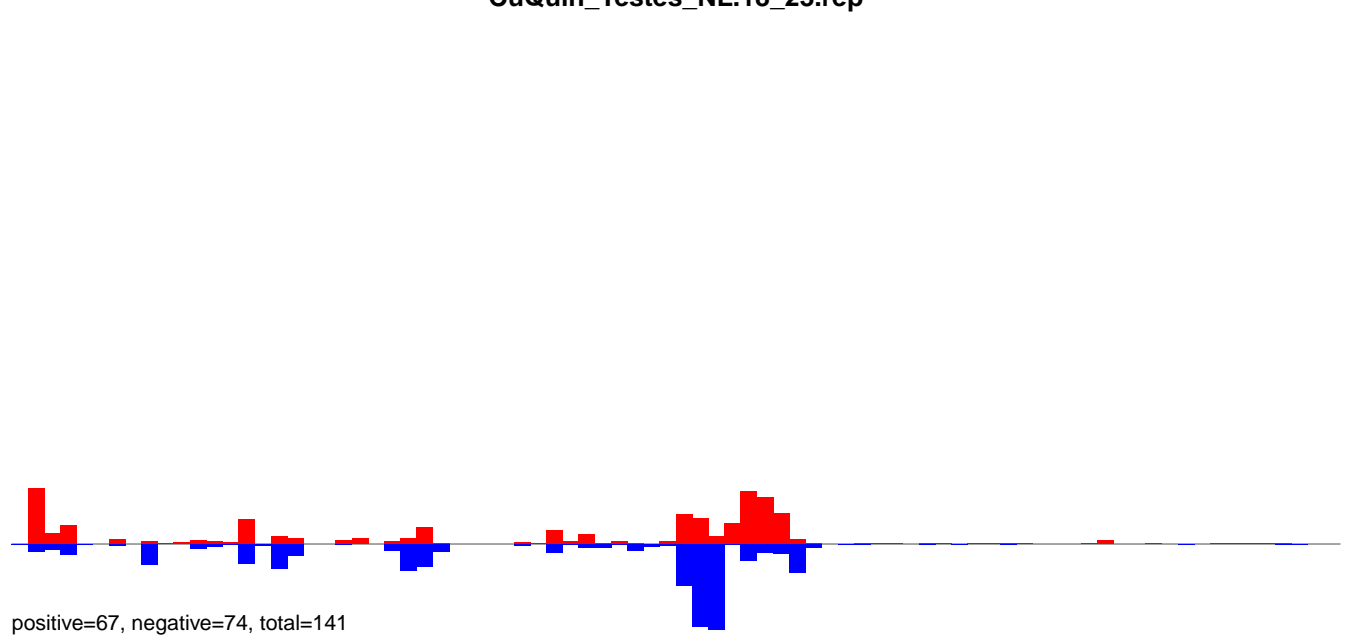


positive=3428, negative=8, total=3437

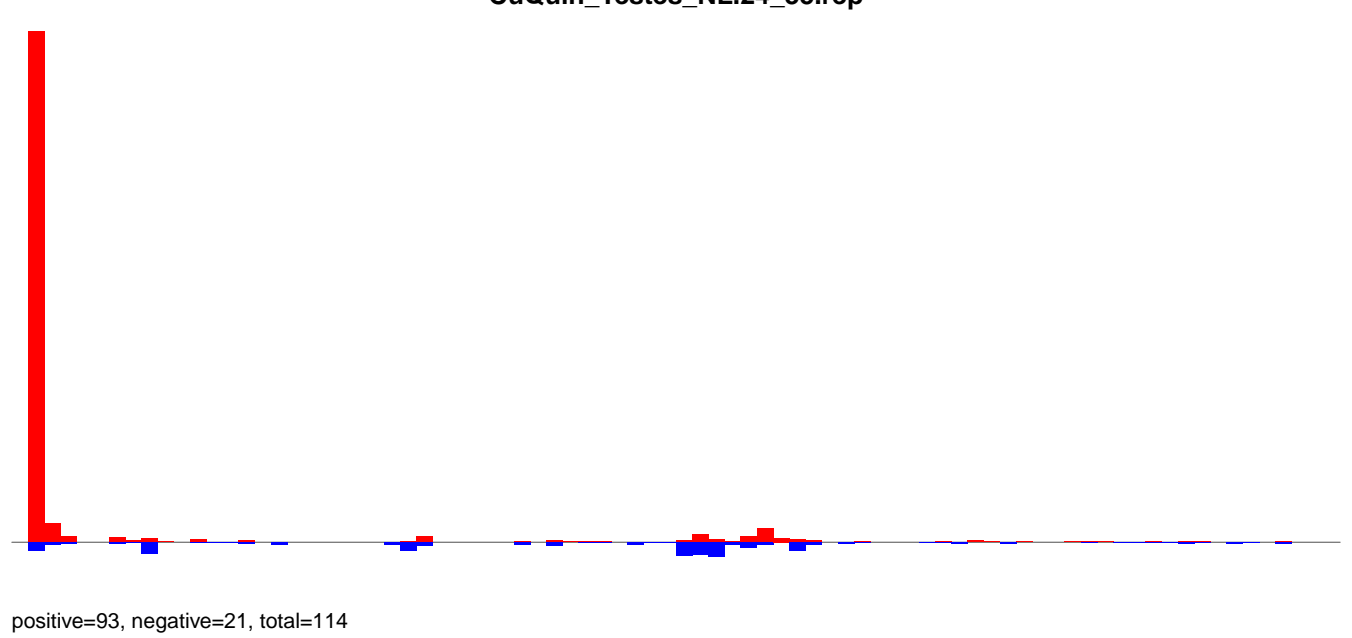
Window size=25, length=1966, TE@rn-4_family-1545@Unknown@2889:1-1966

0 500 1000 1500 2000

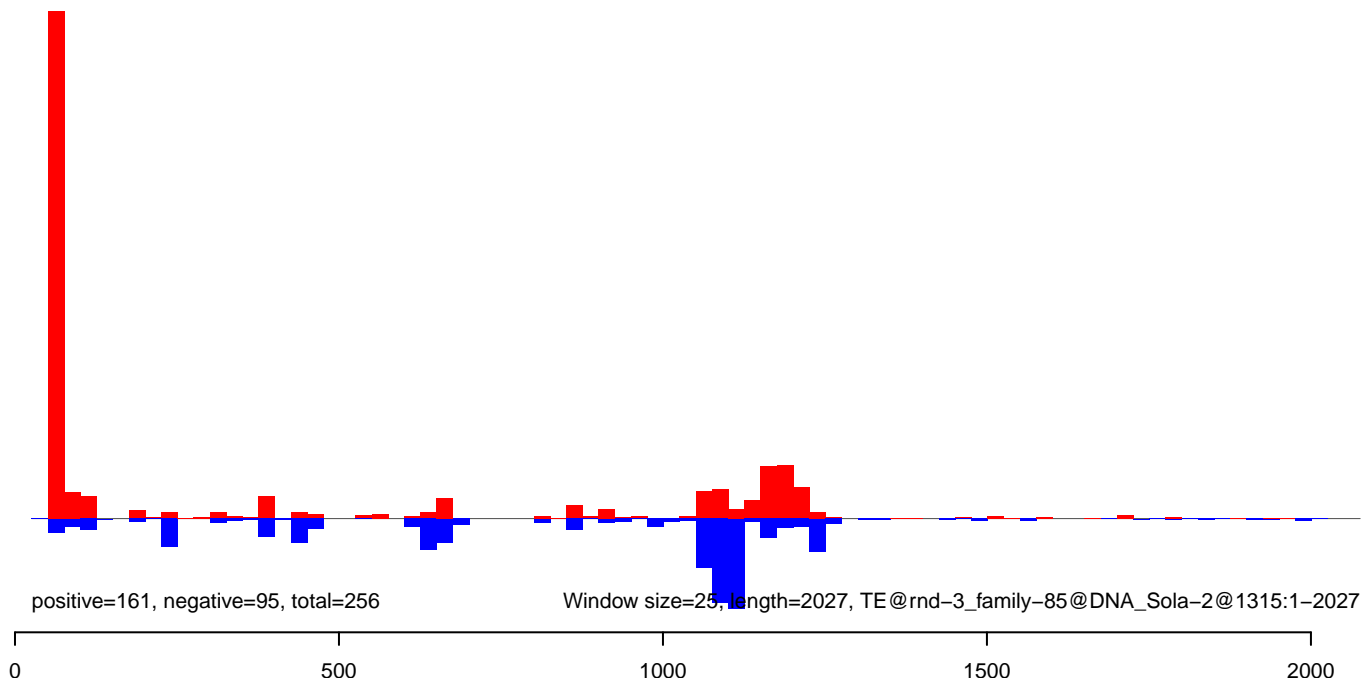
CuQuin_Testes_NL.18_23.rep



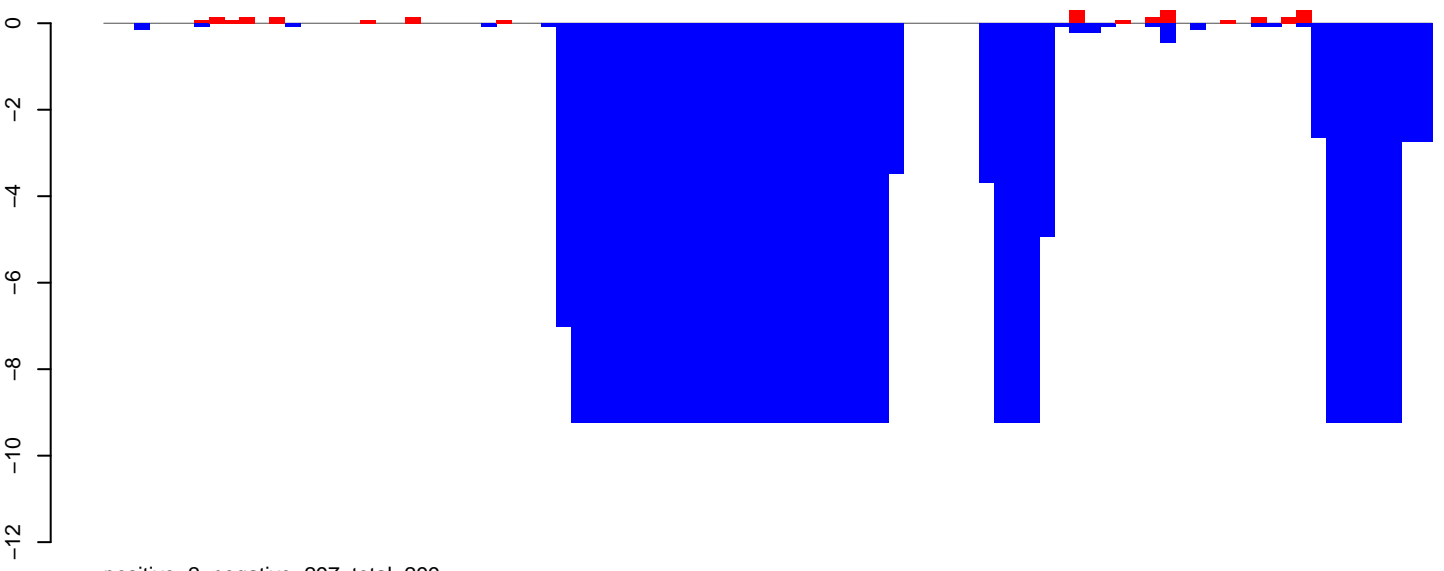
CuQuin_Testes_NL.24_35.rep



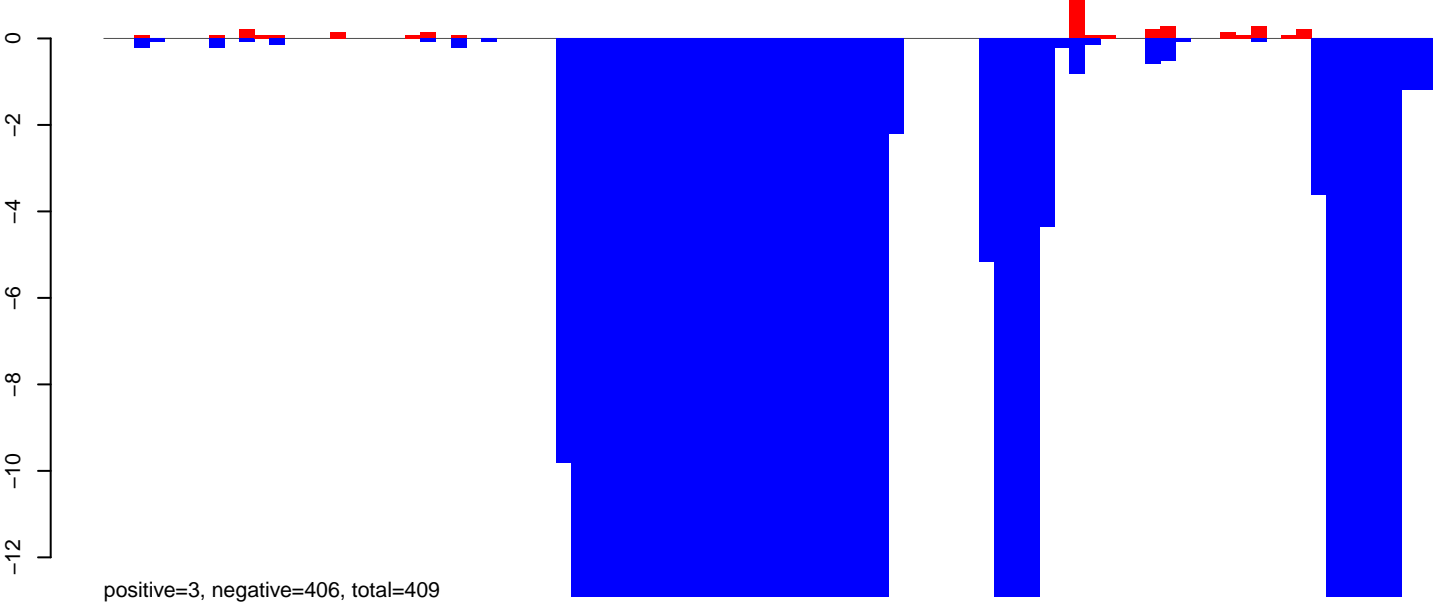
CuQuin_Testes_NL.rep



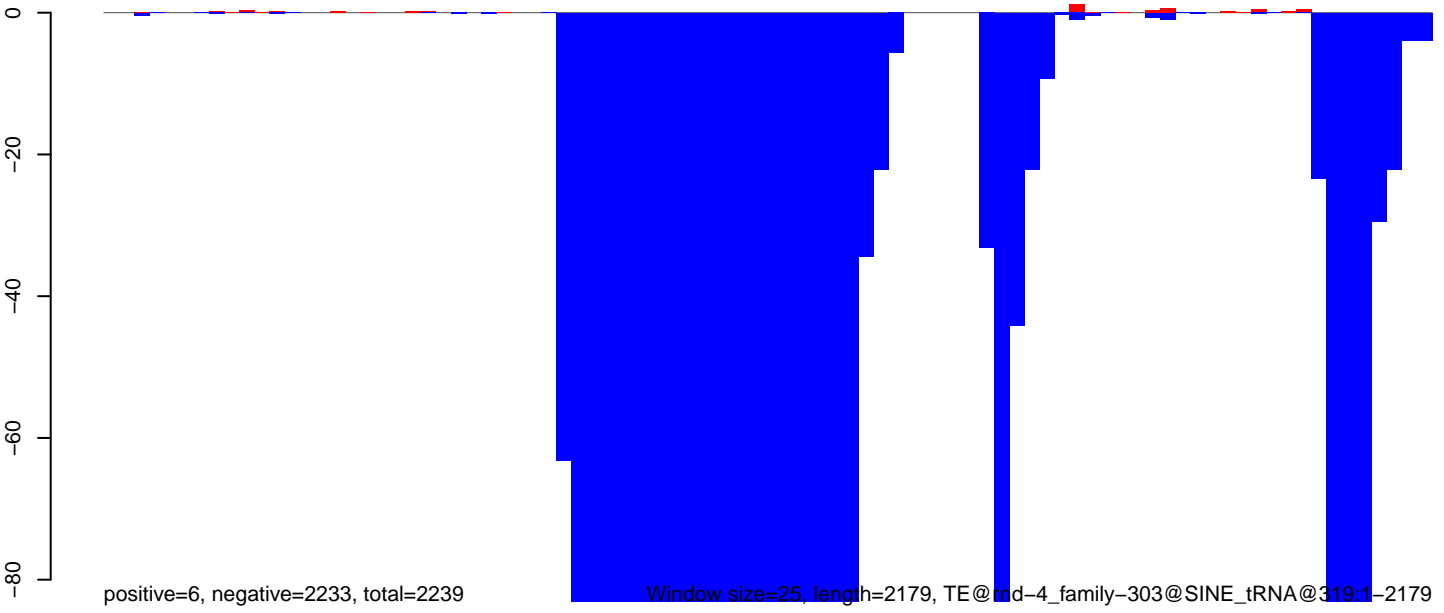
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

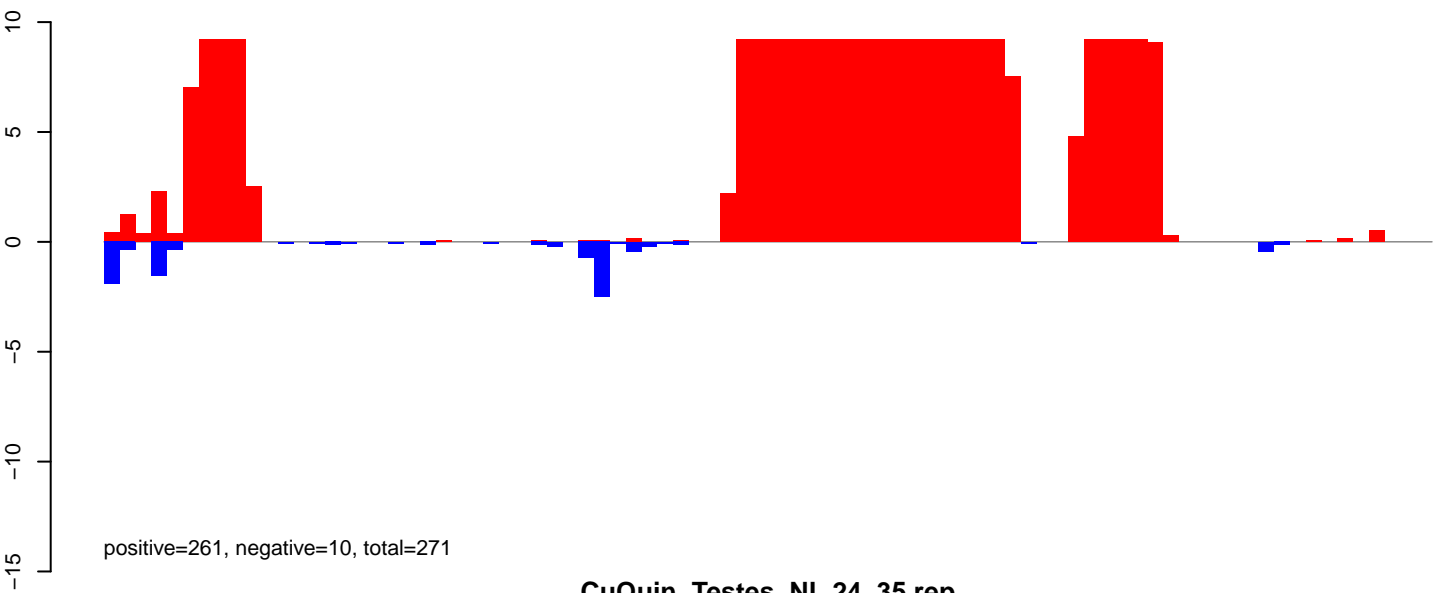


CuQuin_Testes_NL.rep

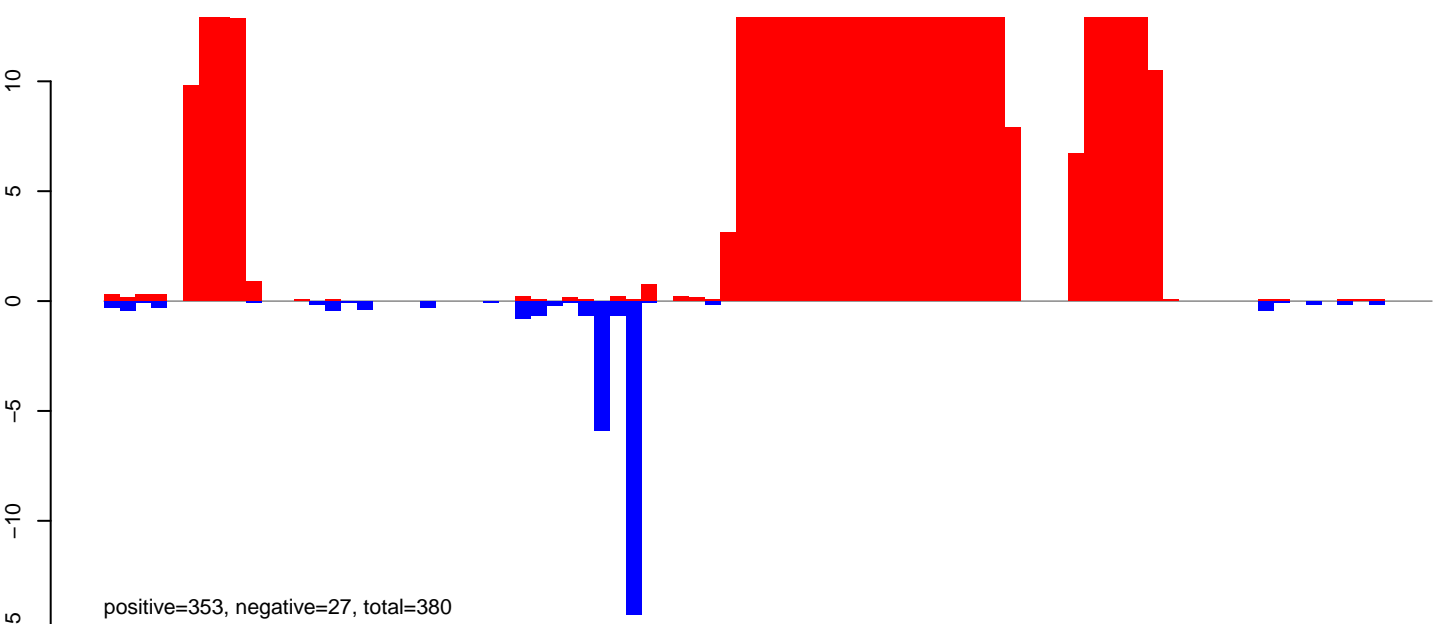


Window size=25, length=2179, TE@mid-4_family-303@SINE_tRNA@319:1-2179

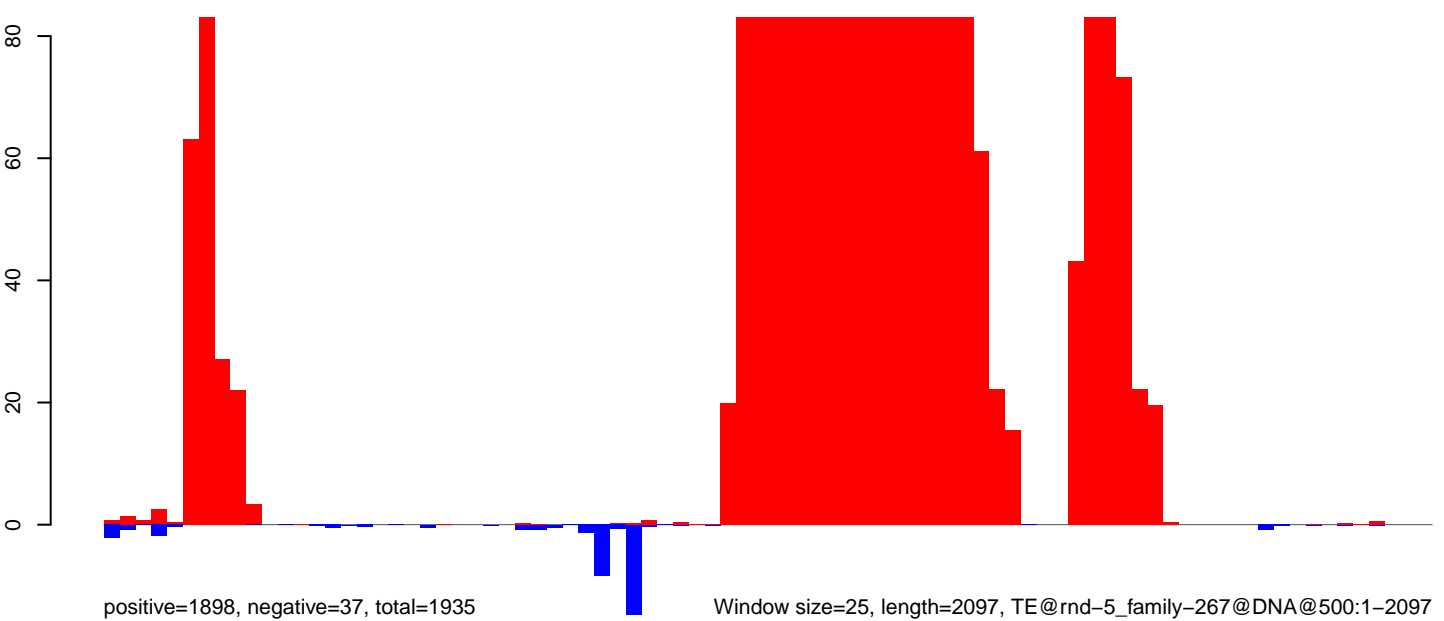
CuQuin_Testes_NL.18_23.rep



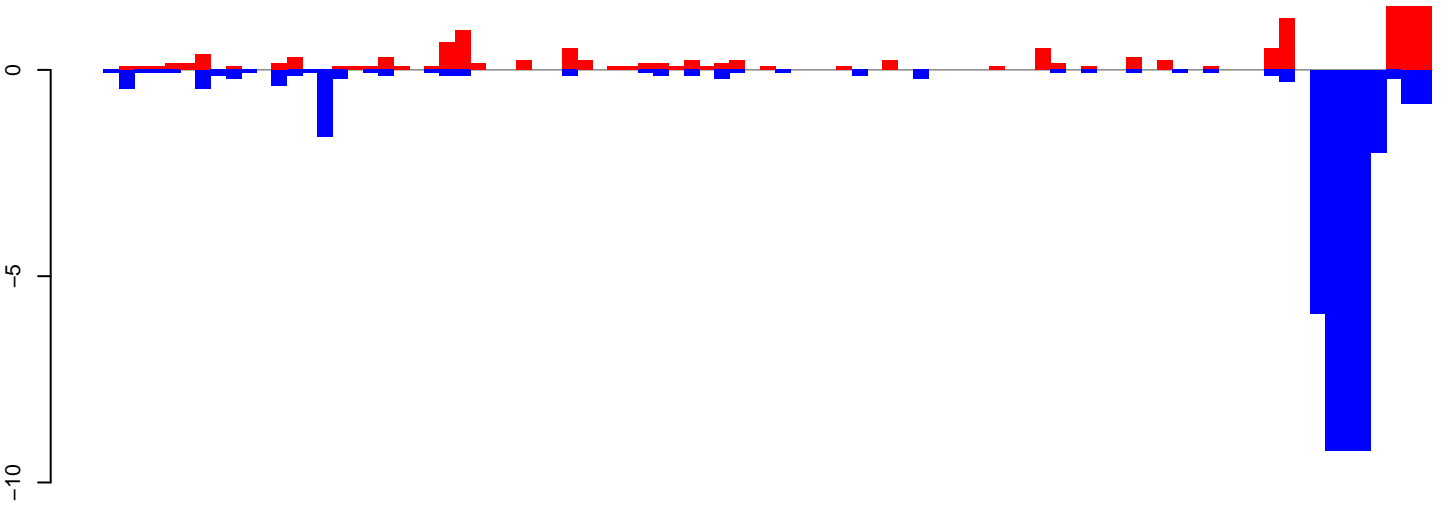
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

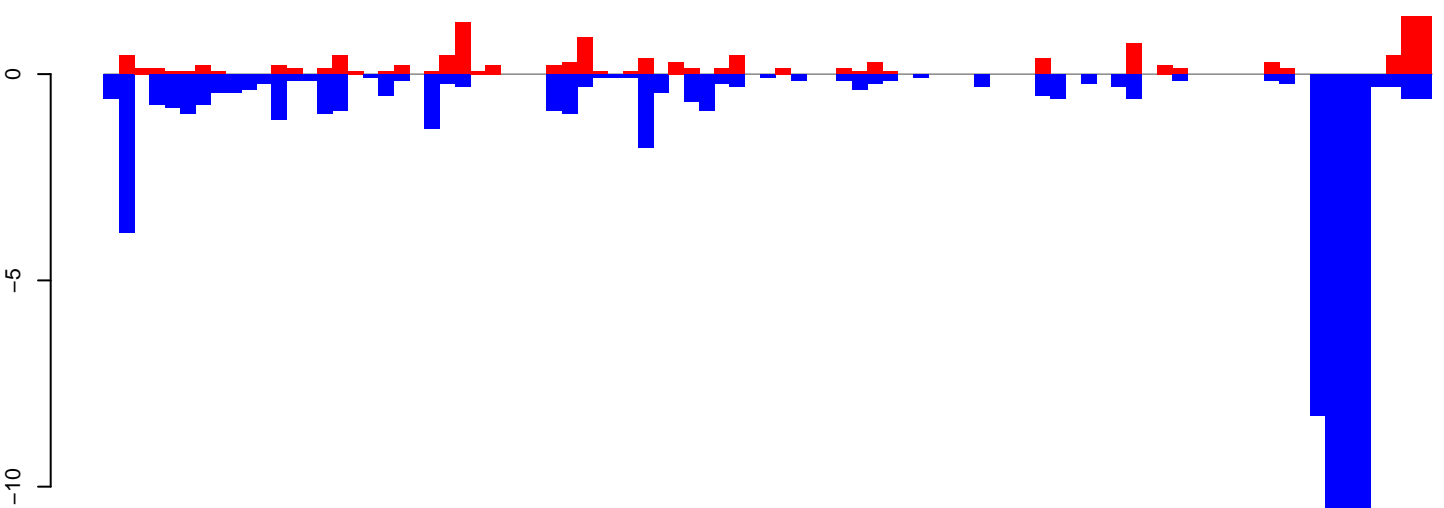


CuQuin_Testes_NL.18_23.rep



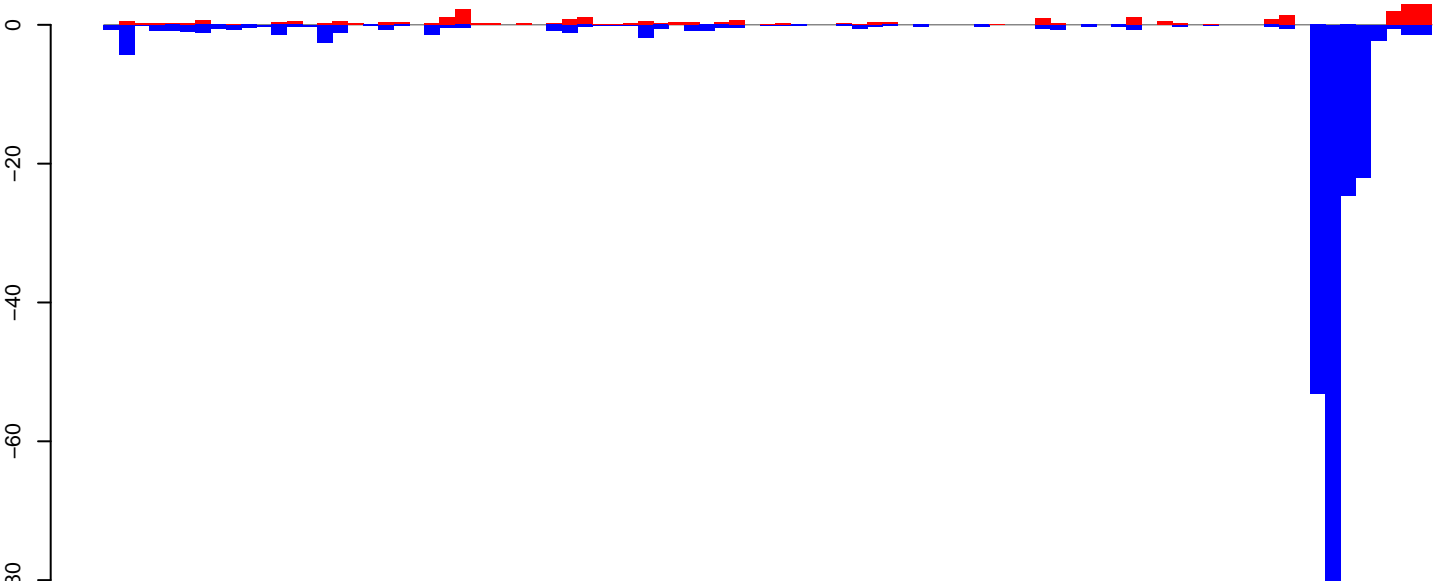
positive=14, negative=44, total=58

CuQuin_Testes_NL.24_35.rep



positive=13, negative=75, total=88

CuQuin_Testes_NL.rep

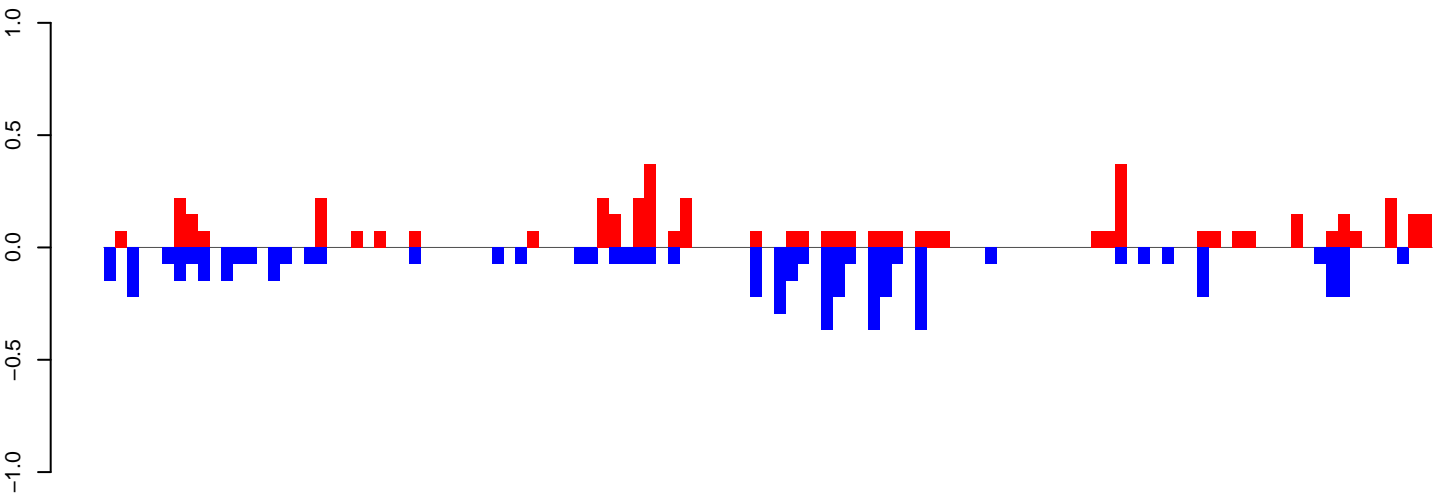


positive=28, negative=221, total=249

Window size=25, length=2172, TE@rnd-5_family-25@LINE_I-Jockey@2342:1-2172

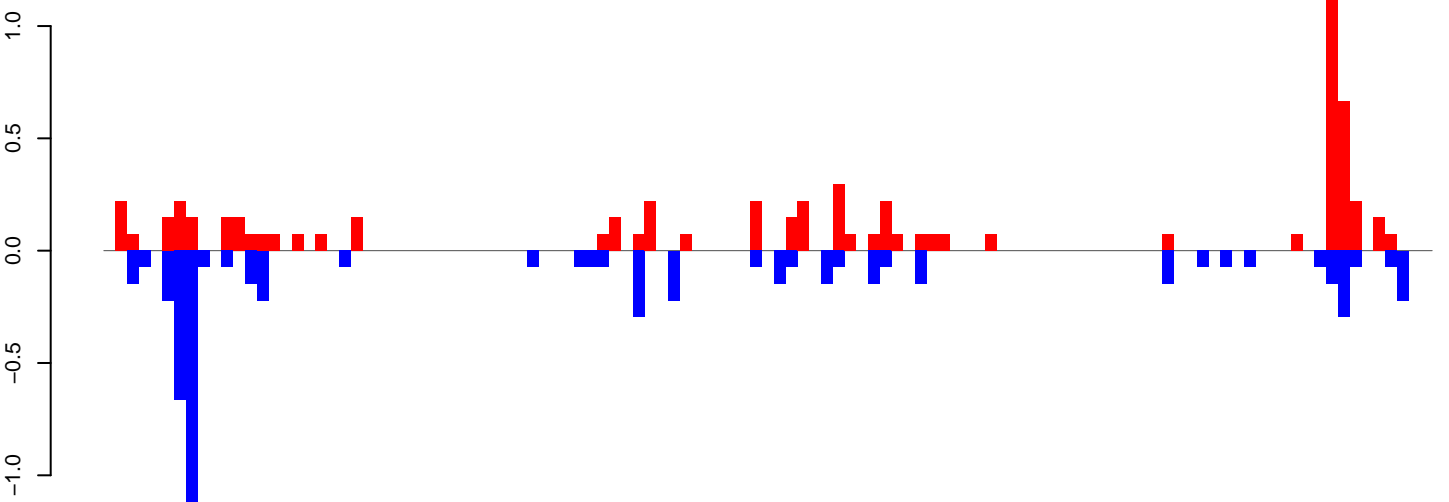
0 500 1000 1500 2000

CuQuin_Testes_NL.18_23.rep



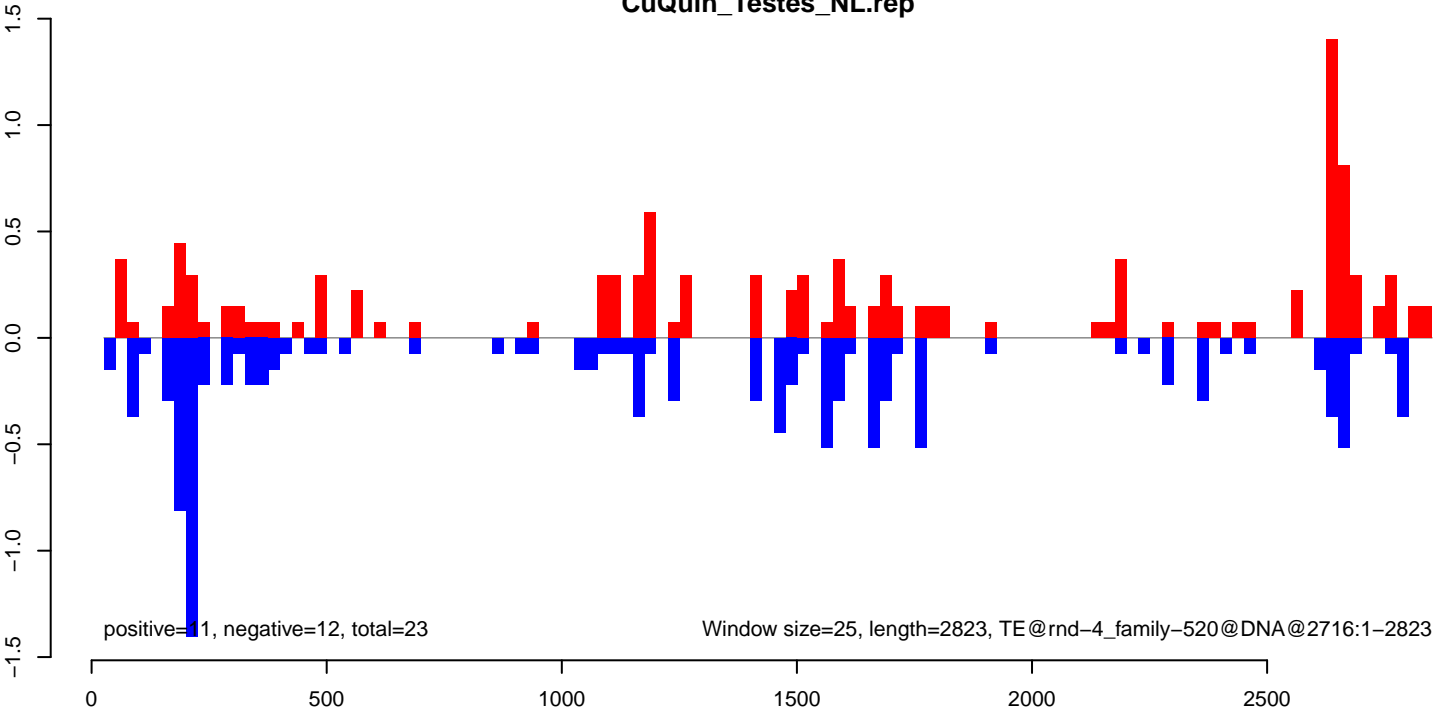
positive=5, negative=6, total=11

CuQuin_Testes_NL.24_35.rep



positive=6, negative=6, total=12

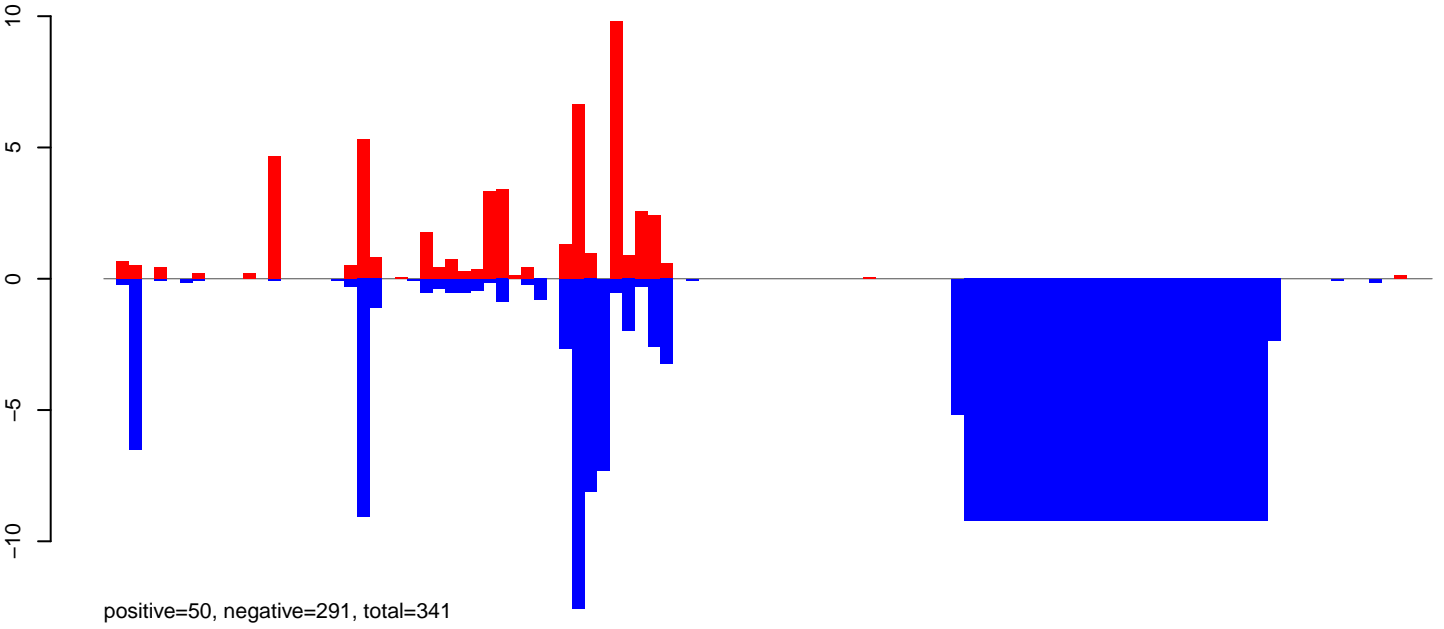
CuQuin_Testes_NL.rep



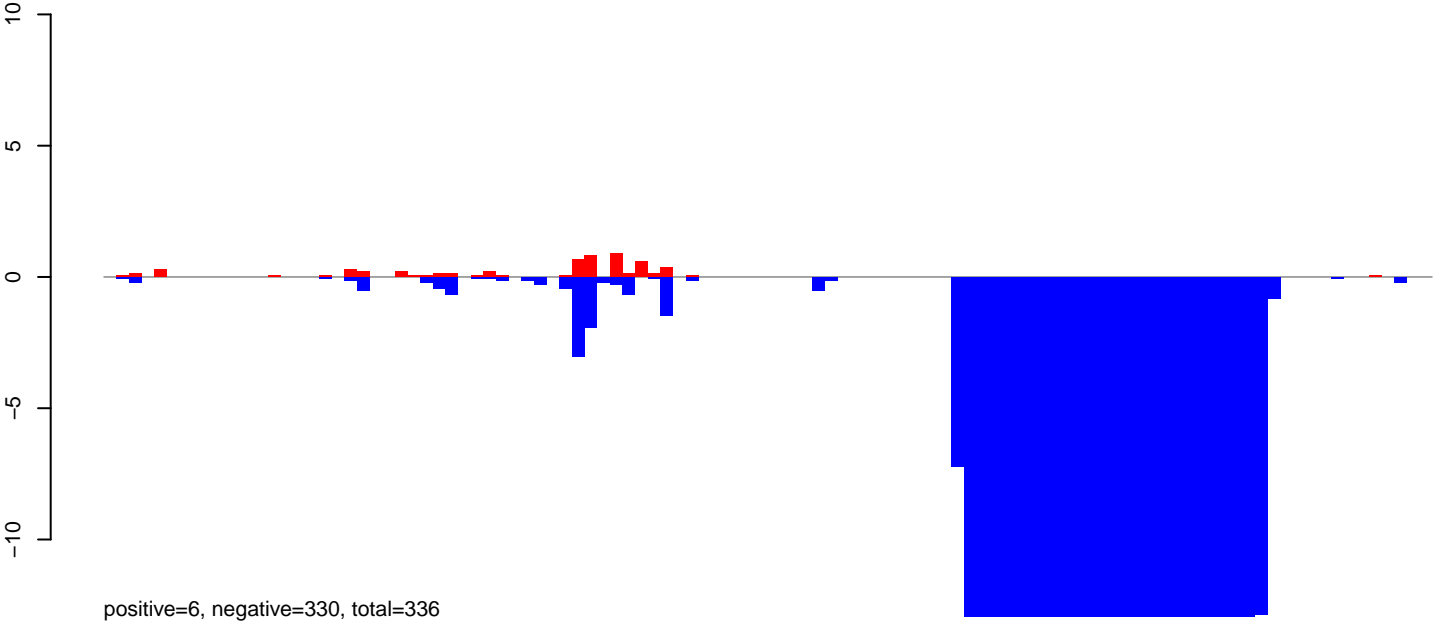
positive=11, negative=12, total=23

Window size=25, length=2823, TE@rnd-4_family-520@DNA@2716:1-2823

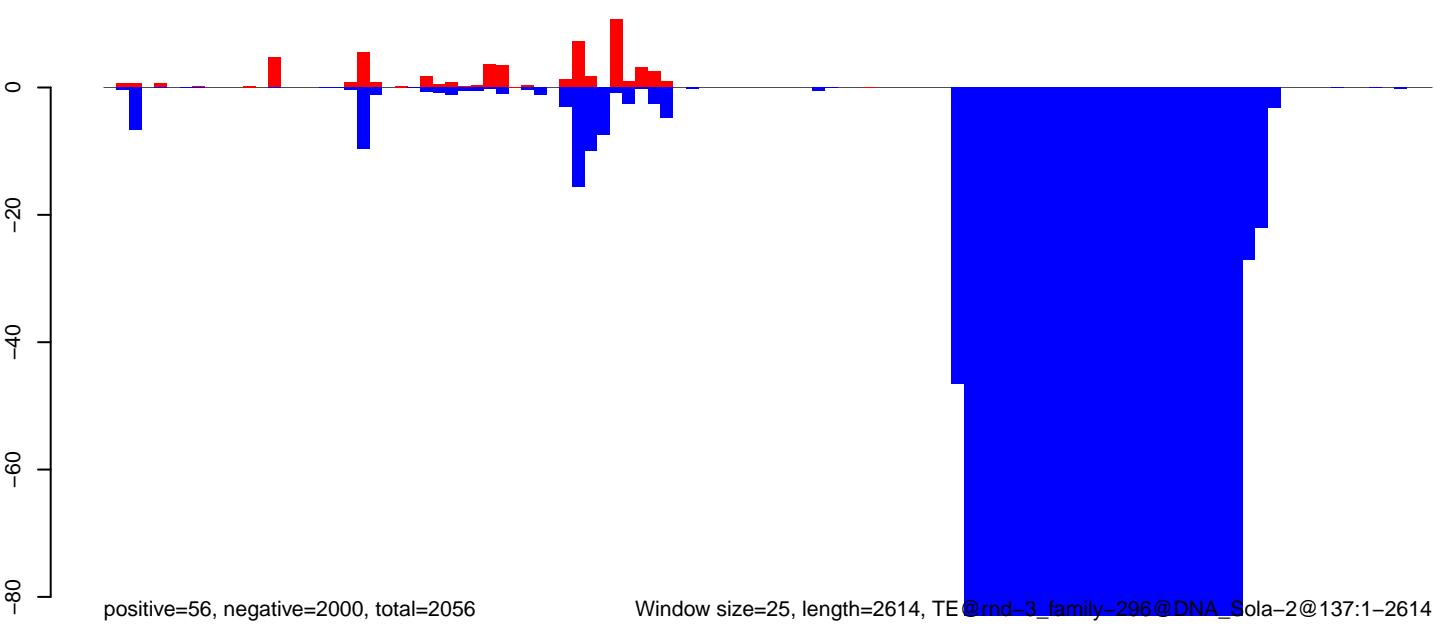
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

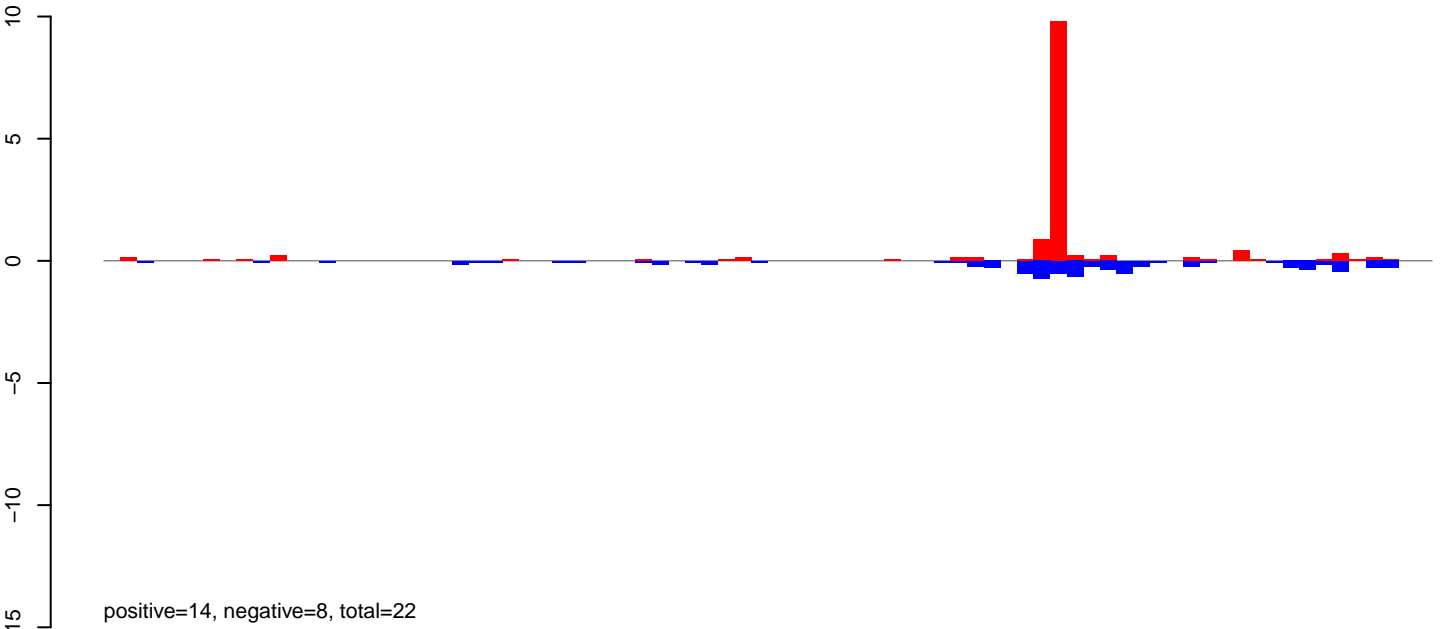


CuQuin_Testes_NL.rep

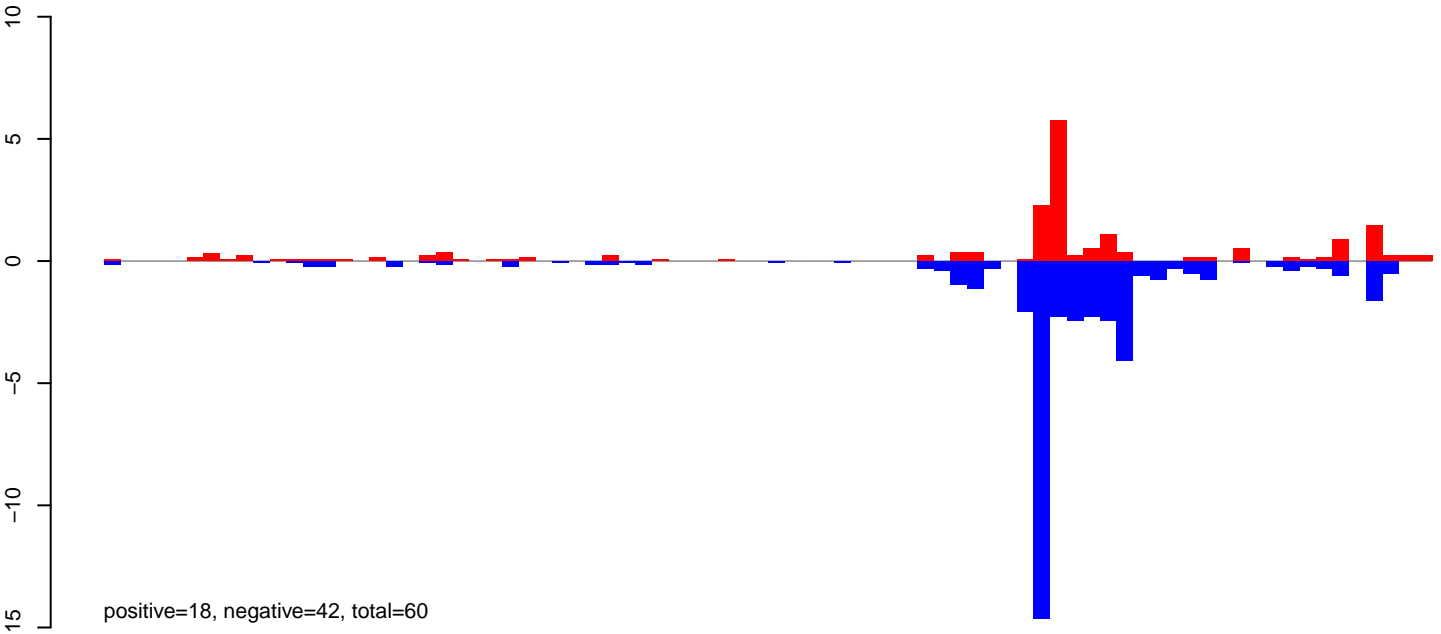


Window size=25, length=2614, TE @rnd-3_family-296@DNA_Sola-2@137:1-2614

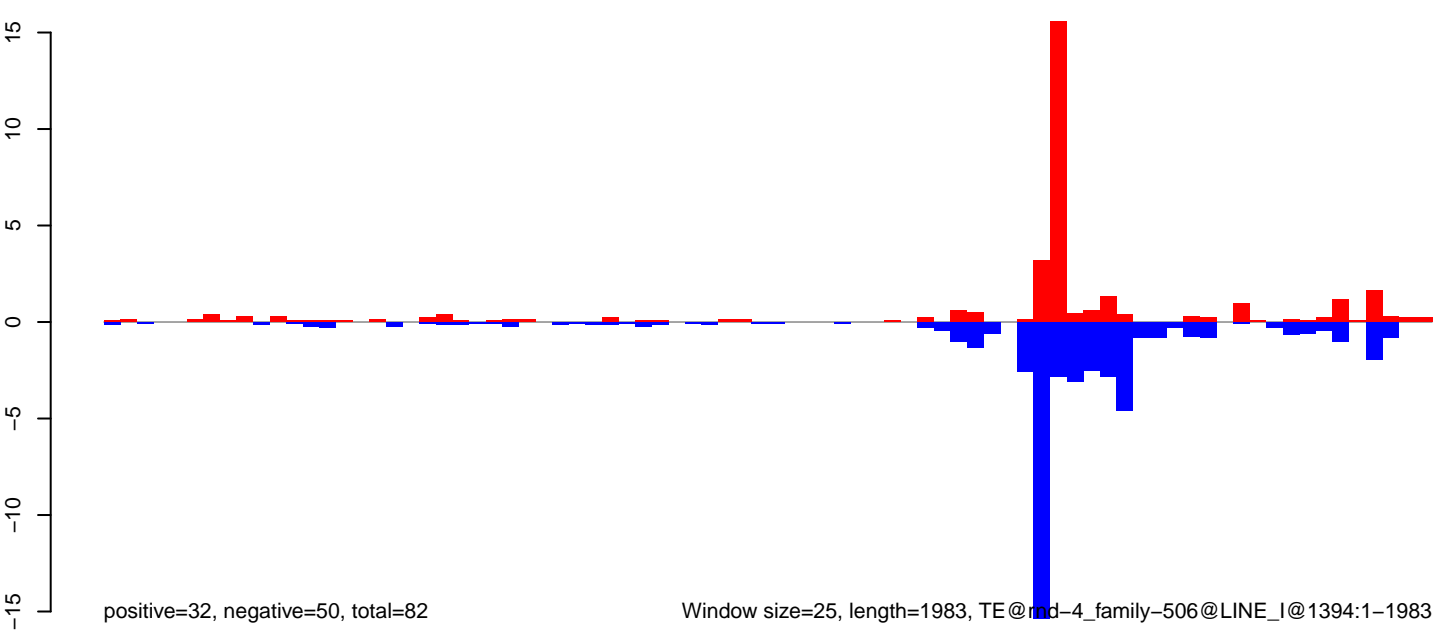
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

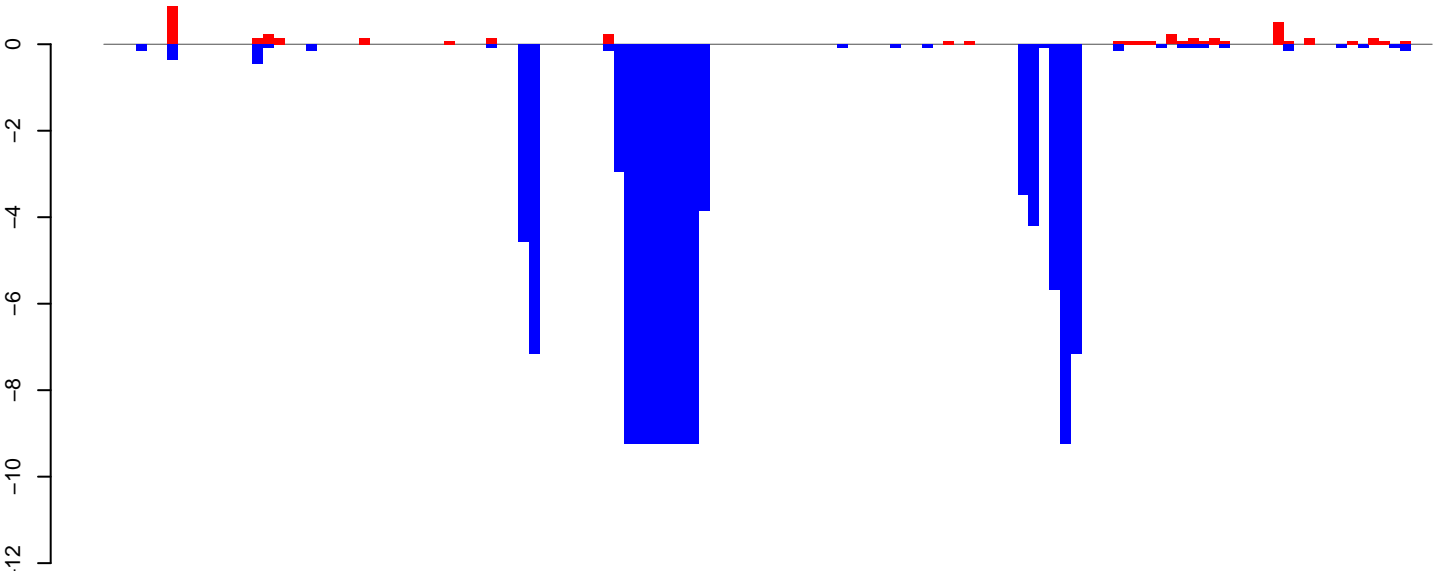


CuQuin_Testes_NL.rep



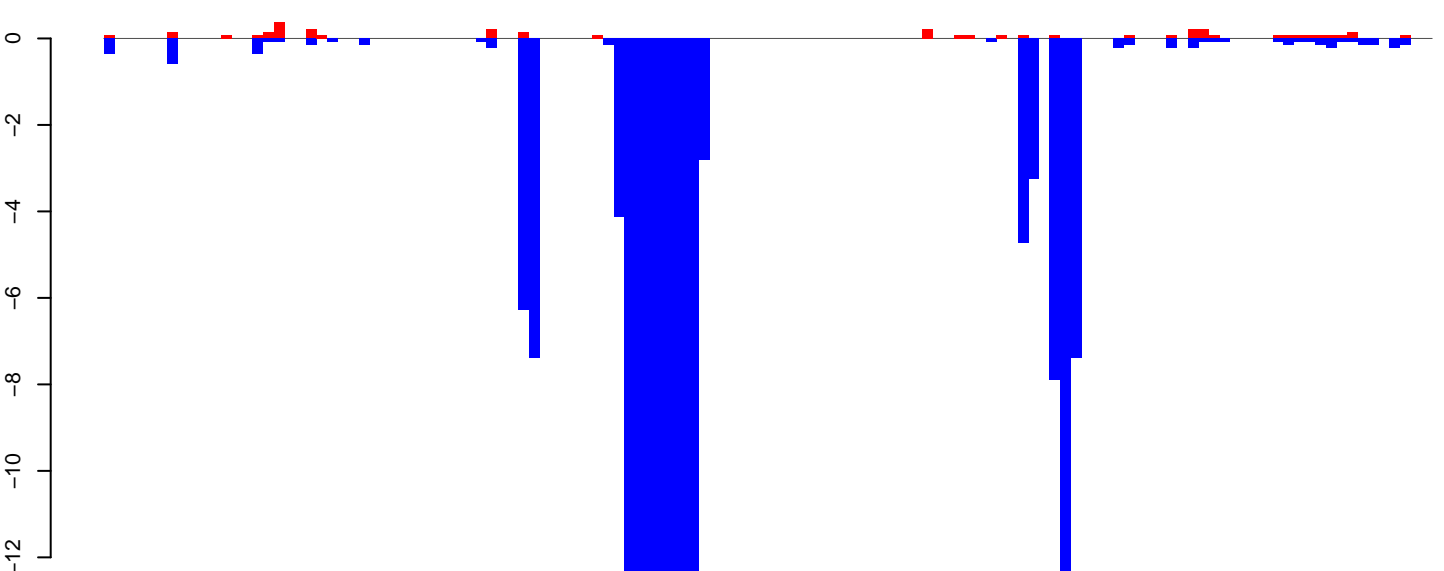
Window size=25, length=1983, TE@md-4_family-506@LINE_I@1394:1-1983

CuQuin_Testes_NL.18_23.rep



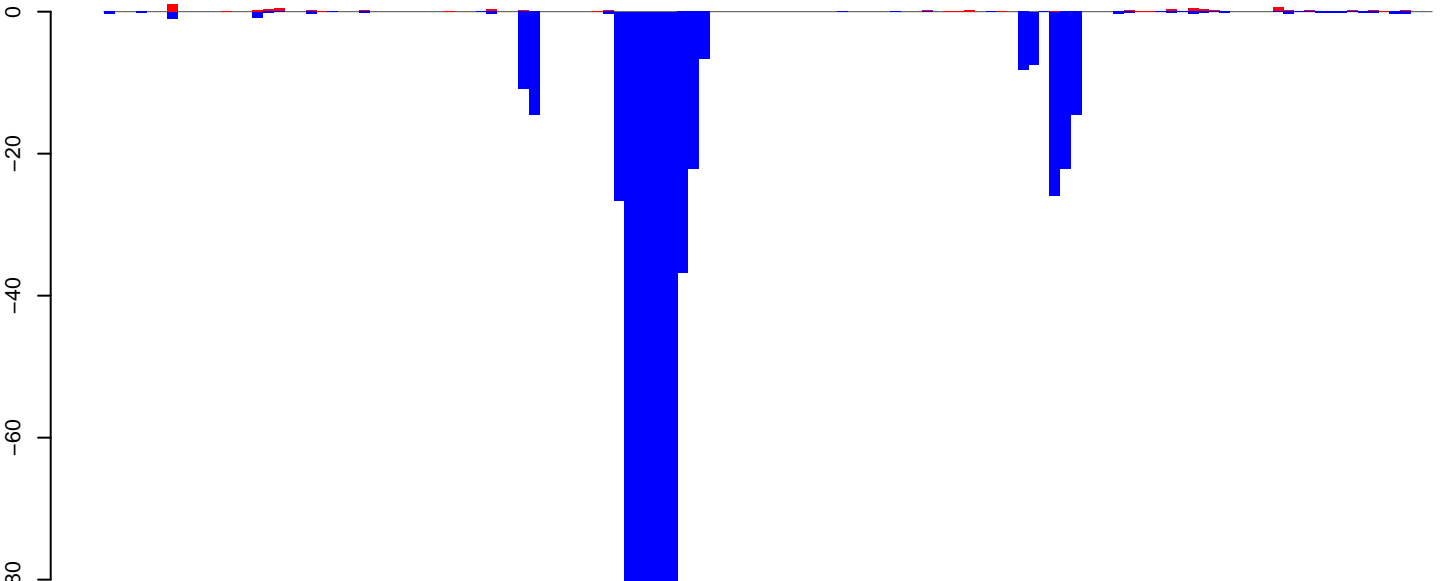
positive=4, negative=116, total=120

CuQuin_Testes_NL.24_35.rep



positive=4, negative=152, total=156

CuQuin_Testes_NL.rep

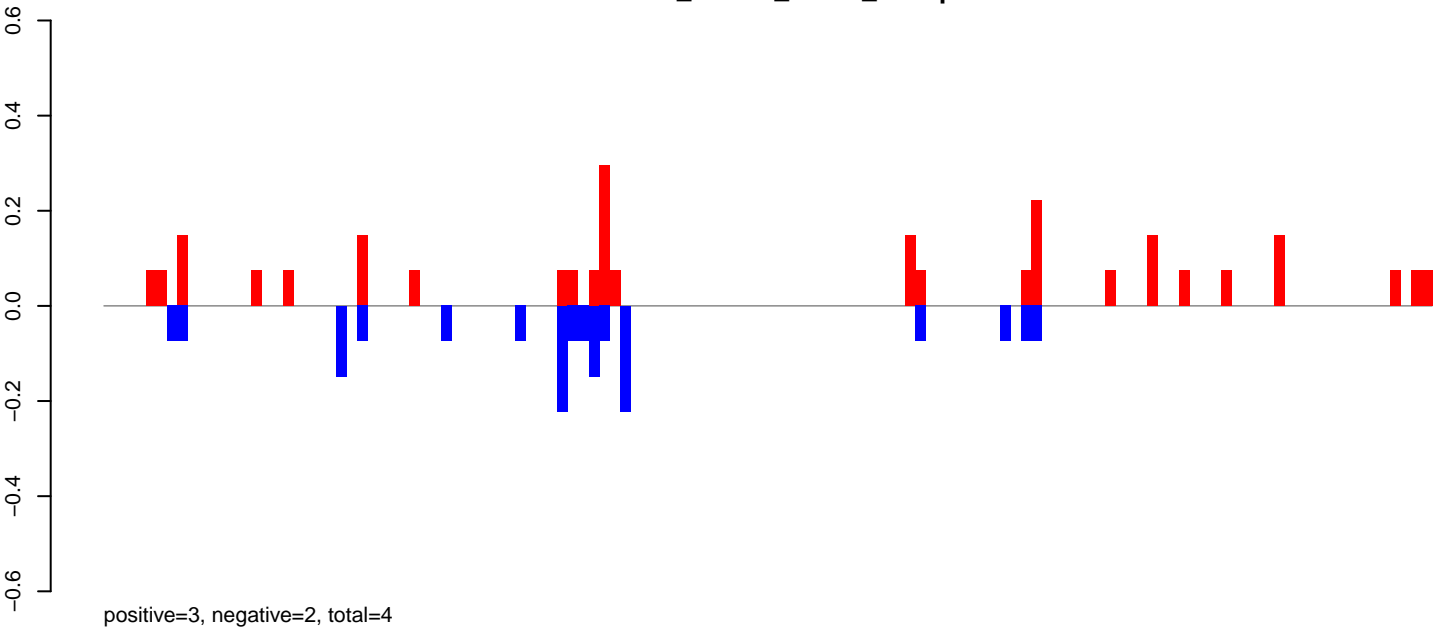


positive=8, negative=619, total=627

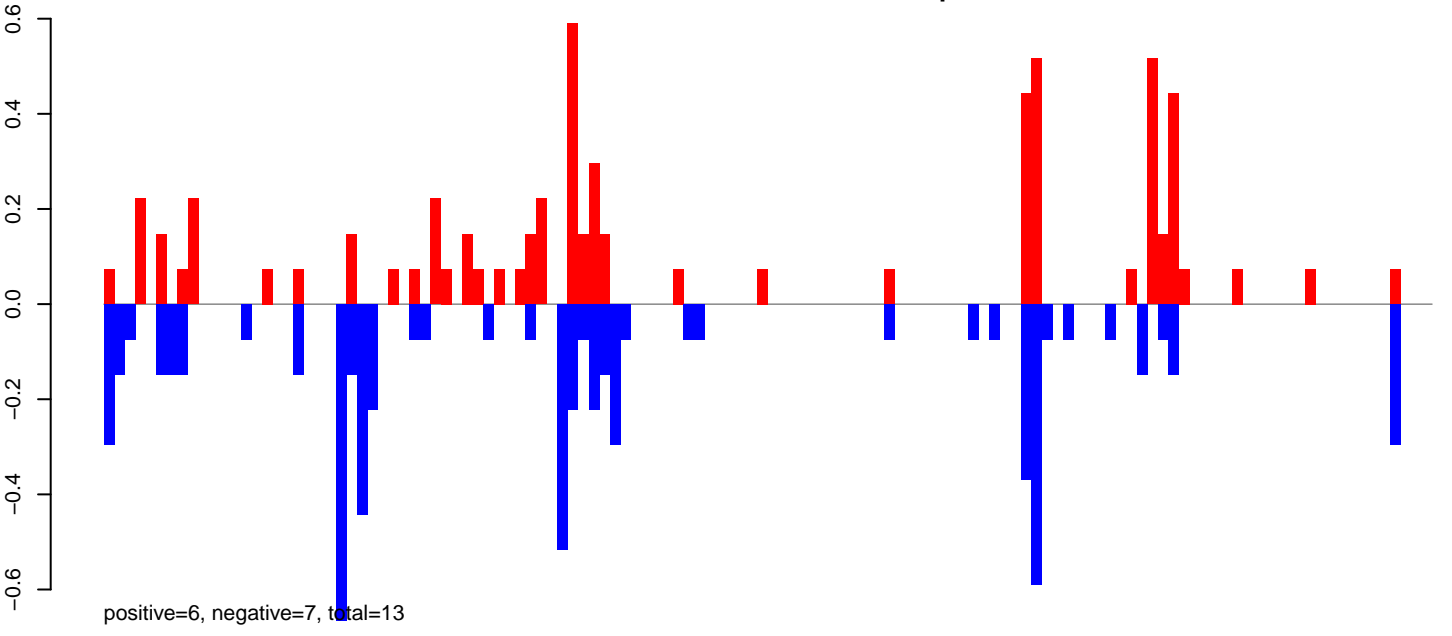
Window size=25, length=3107, TE@rnd-3_family-11@DNA@658:1-3107

0 500 1000 1500 2000 2500 3000

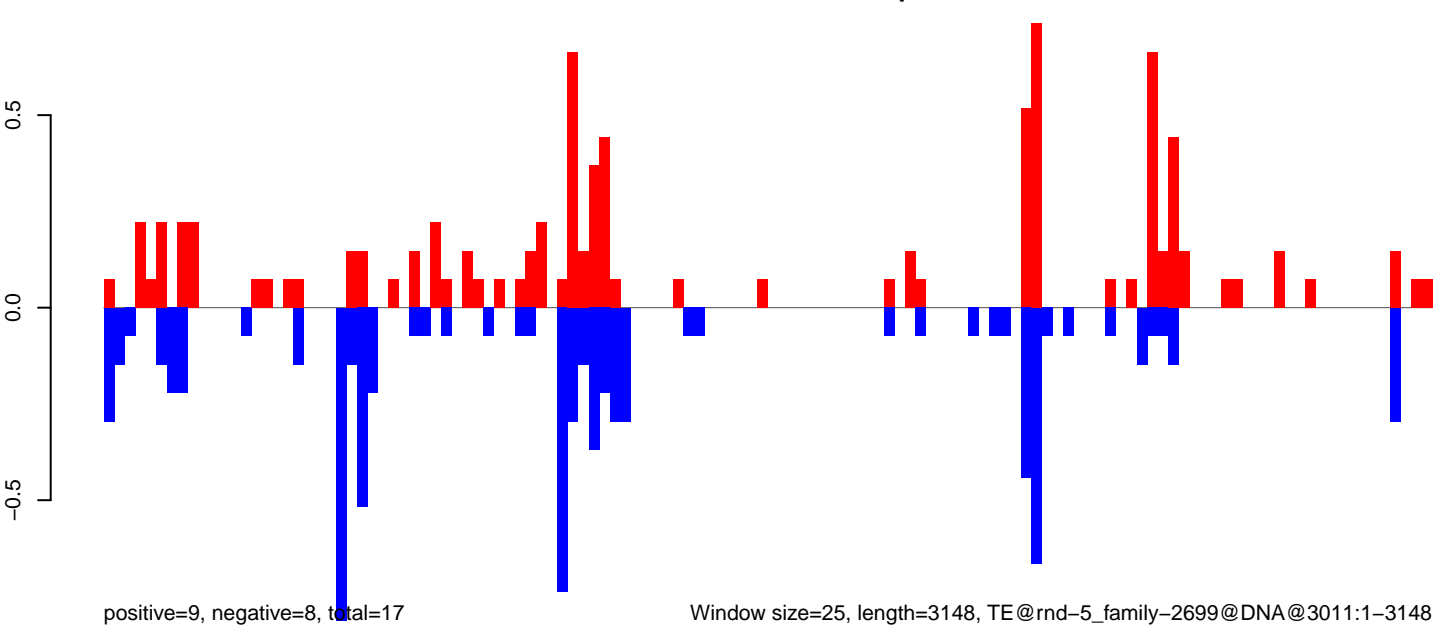
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

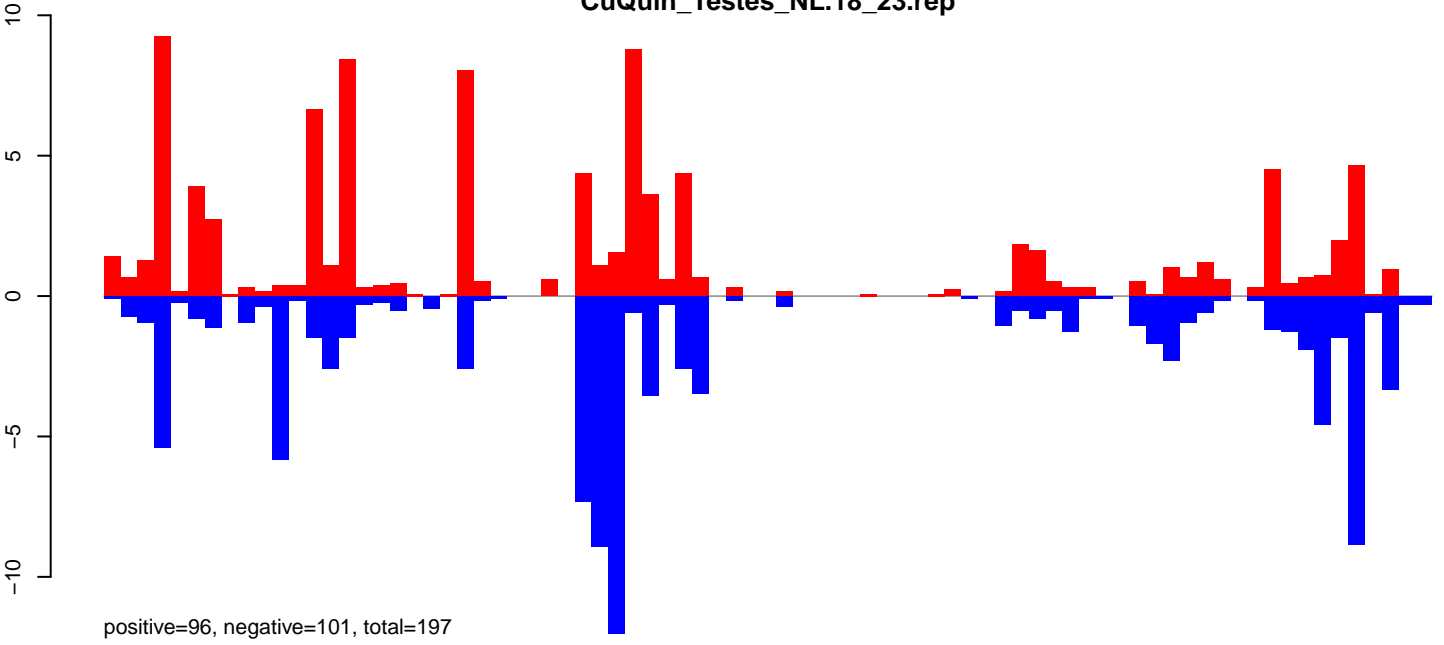


CuQuin_Testes_NL.rep

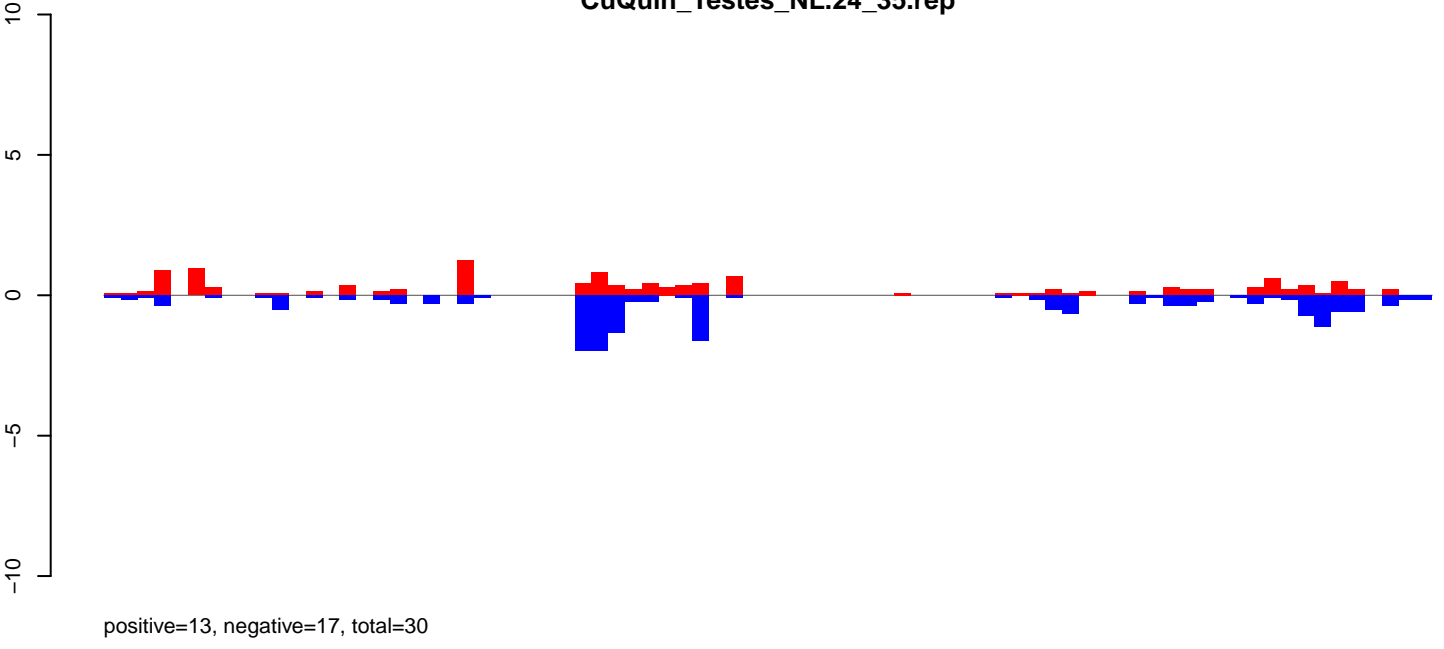


Window size=25, length=3148, TE@rnd-5_family-2699@DNA@3011:1-3148

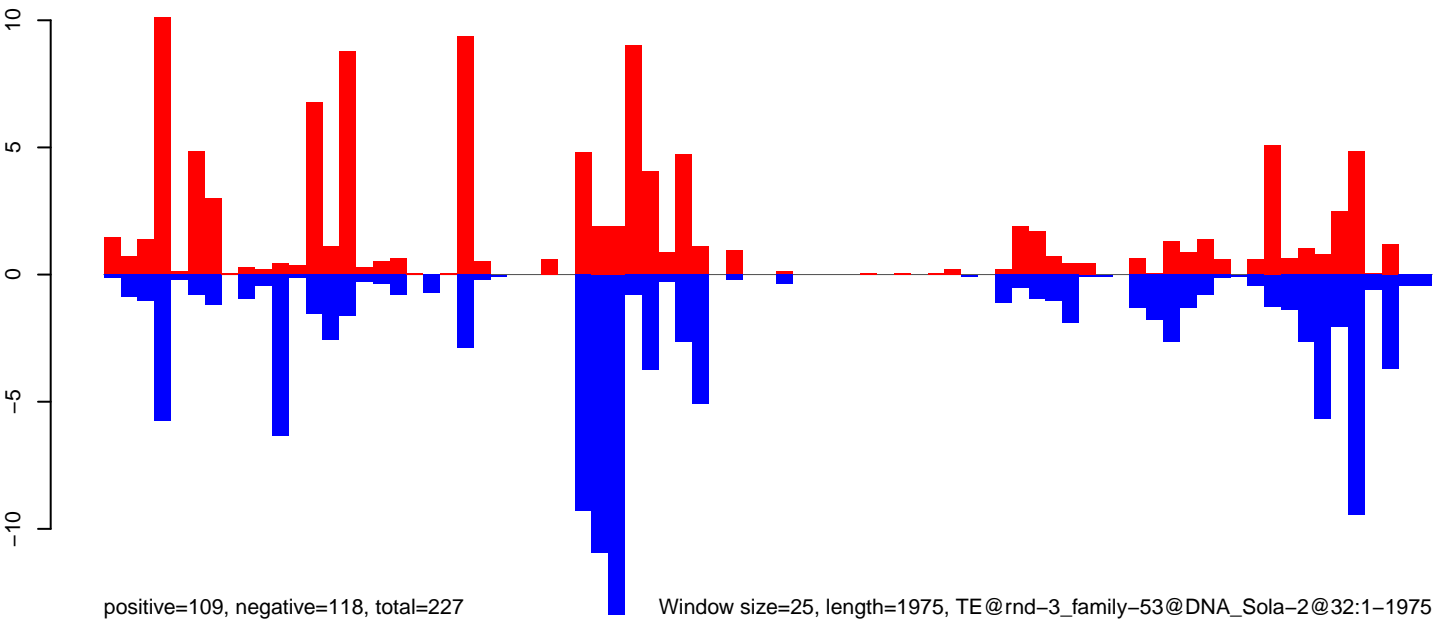
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

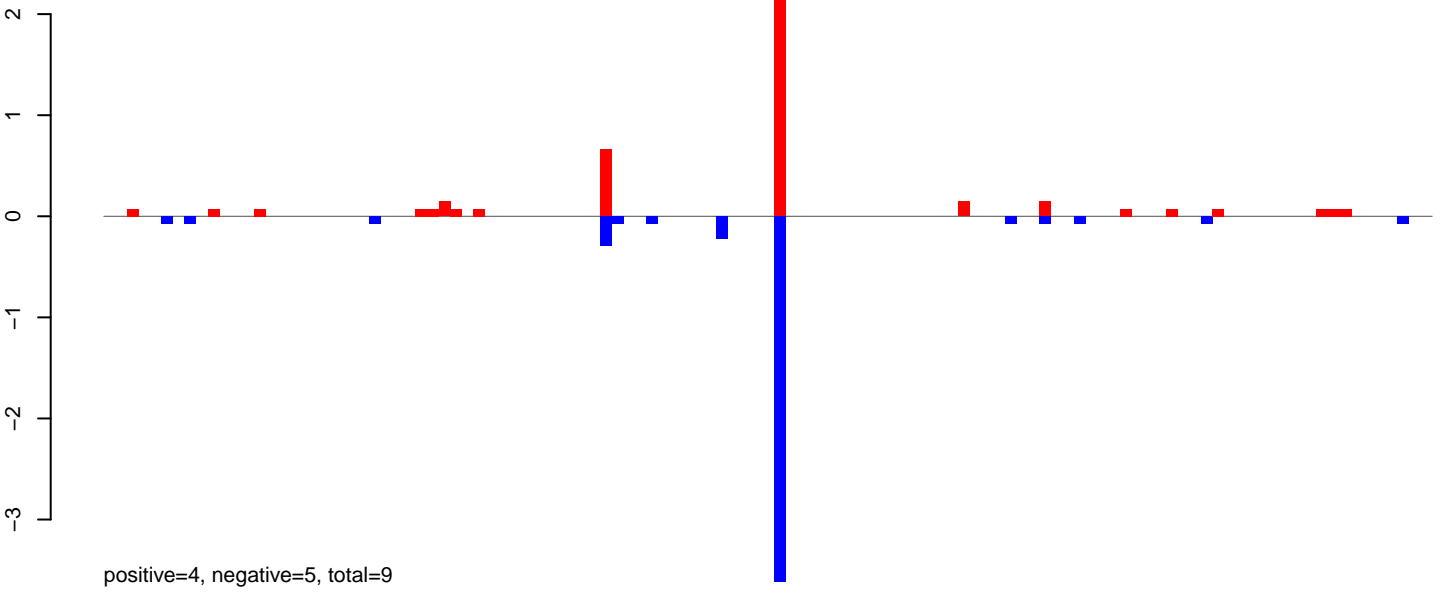


CuQuin_Testes_NL.rep

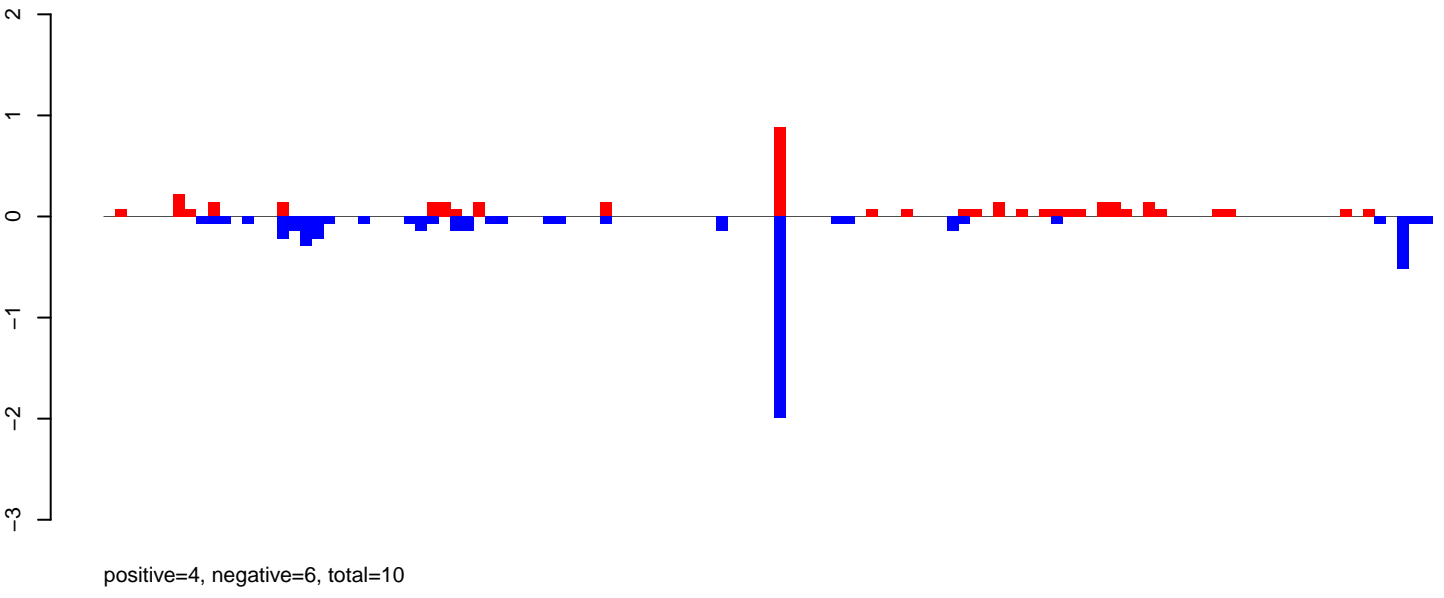


Window size=25, length=1975, TE@rnd-3_family-53@DNA_Sola-2@32:1-1975

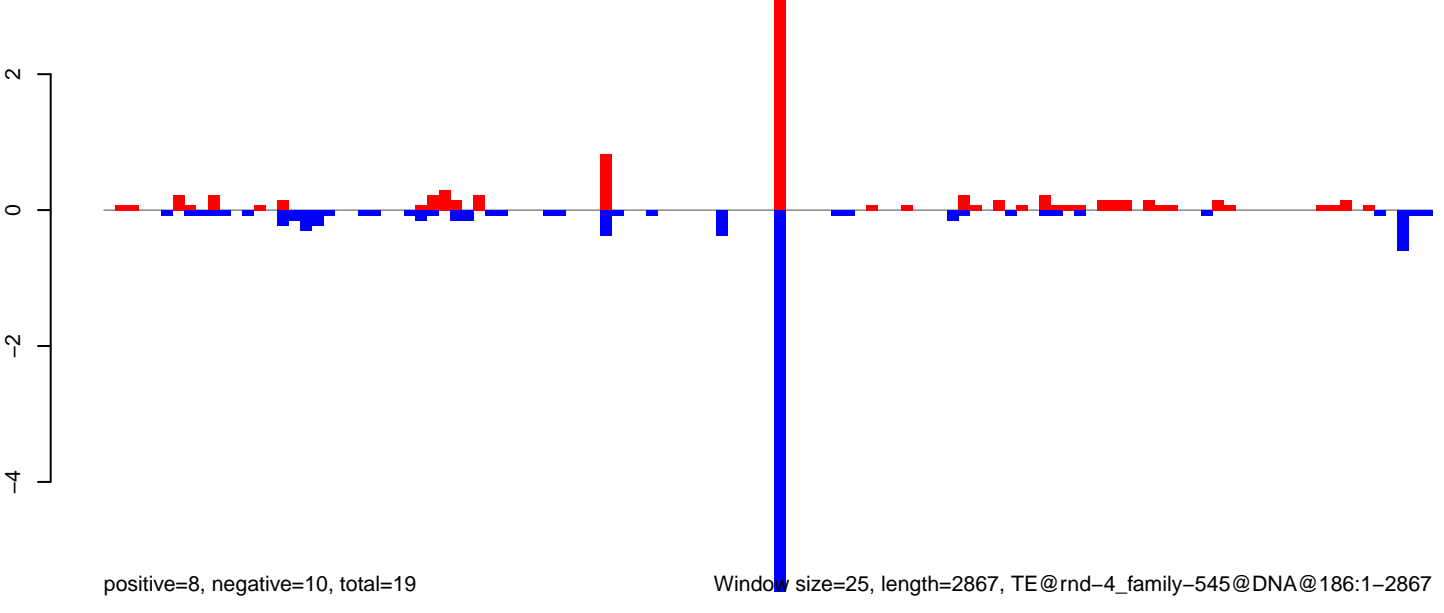
CuQuin_Testes_NL.18_23.rep



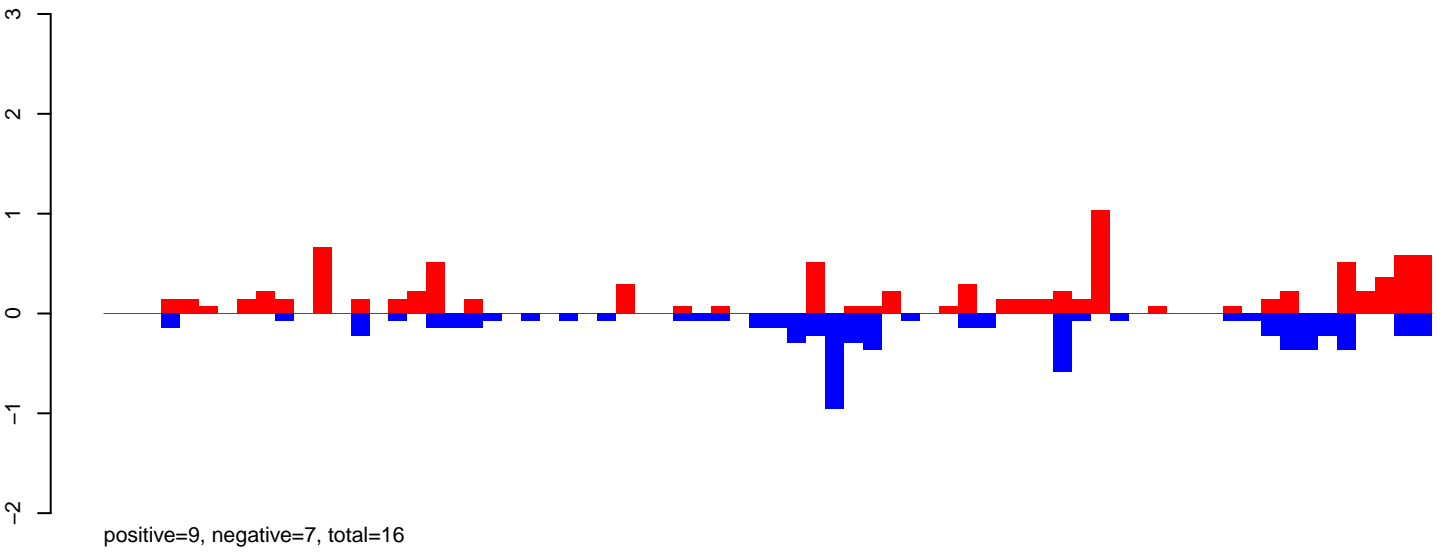
CuQuin_Testes_NL.24_35.rep



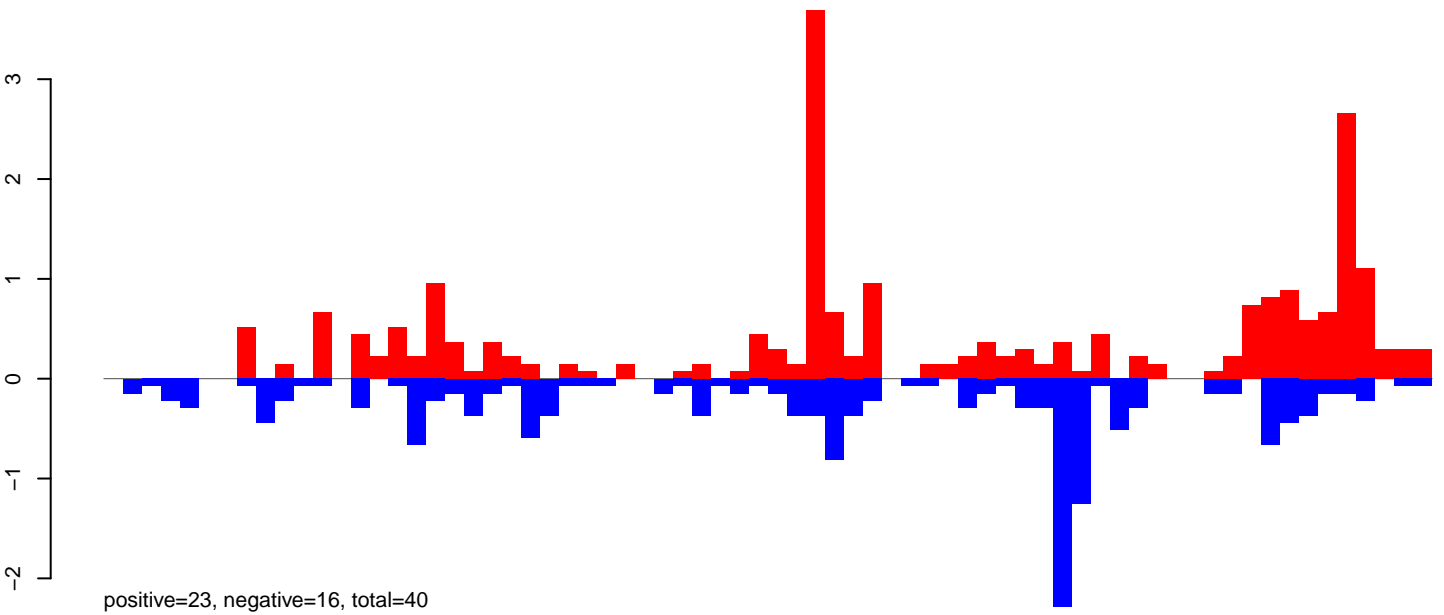
CuQuin_Testes_NL.rep



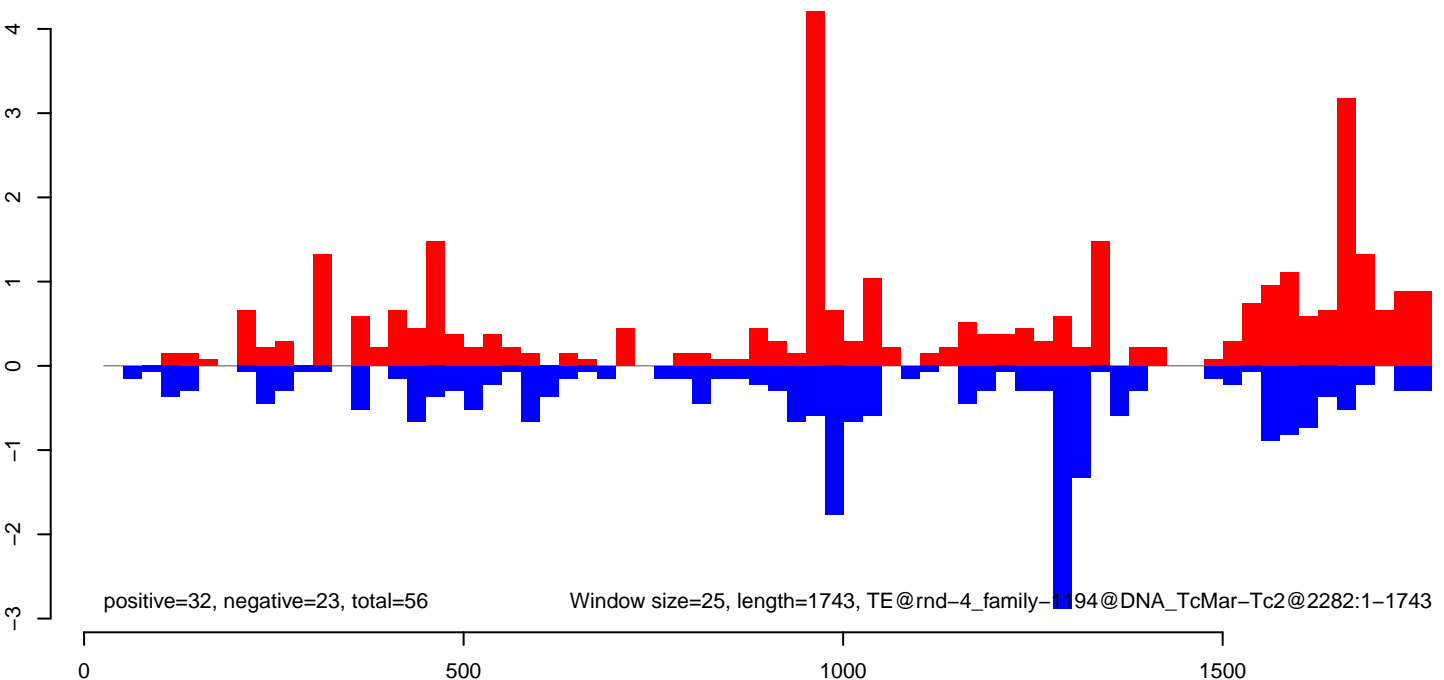
CuQuin_Testes_NL.18_23.rep



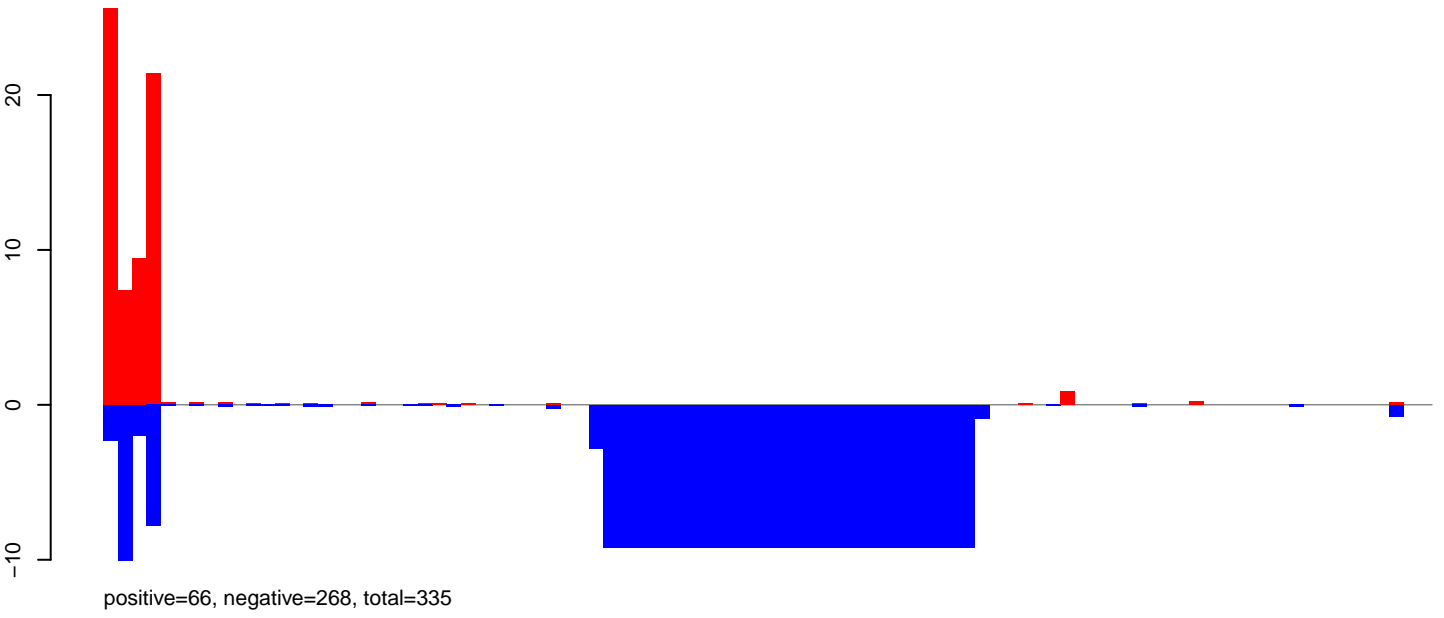
CuQuin_Testes_NL.24_35.rep



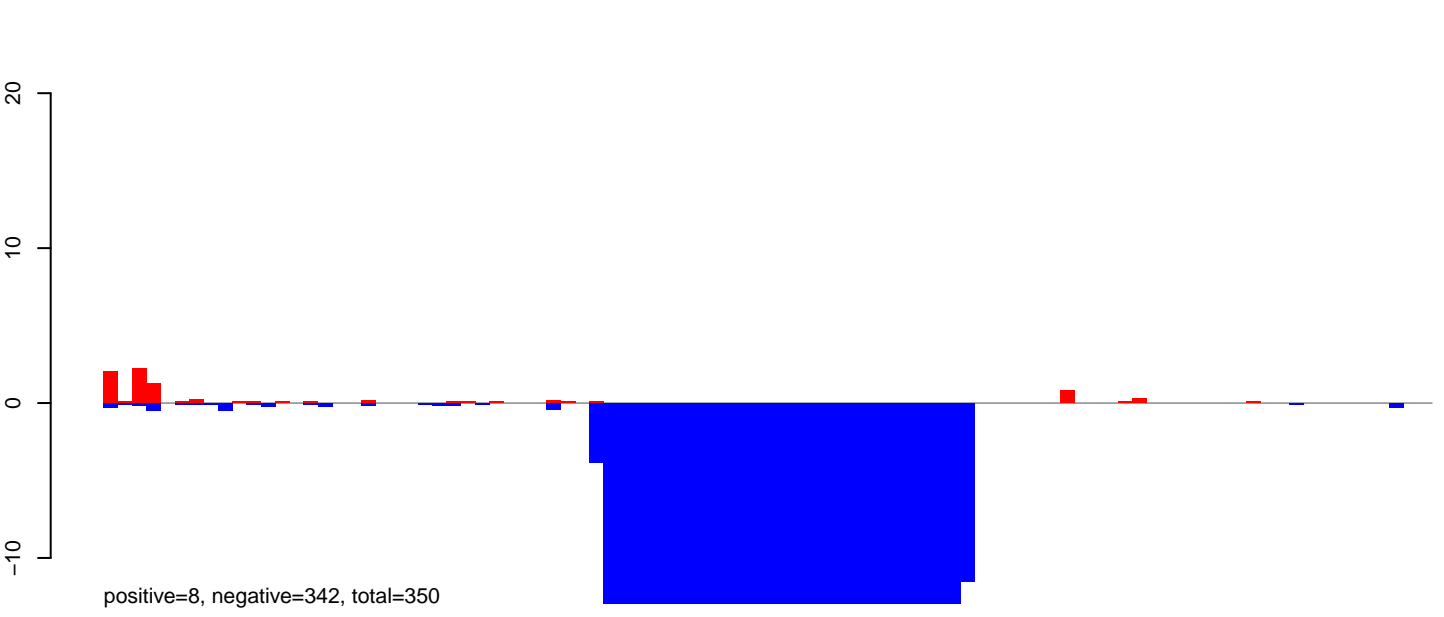
CuQuin_Testes_NL.rep



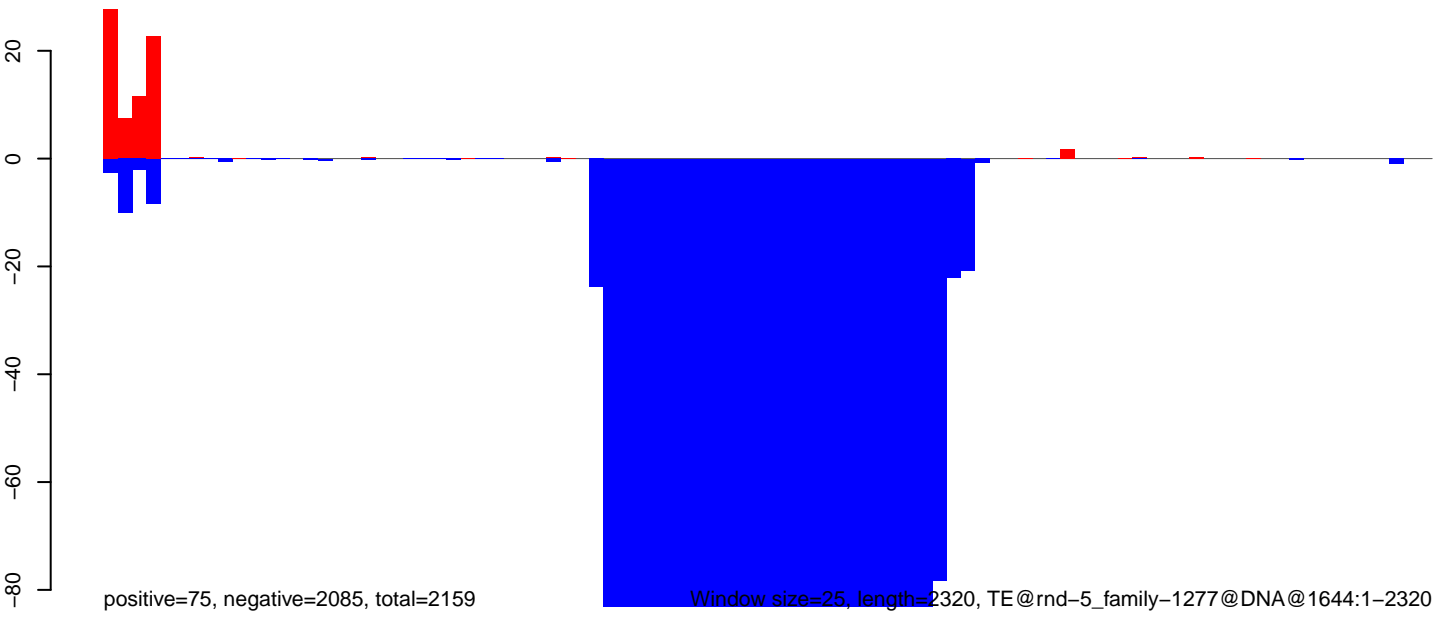
CuQuin_Testes_NL.18_23.rep



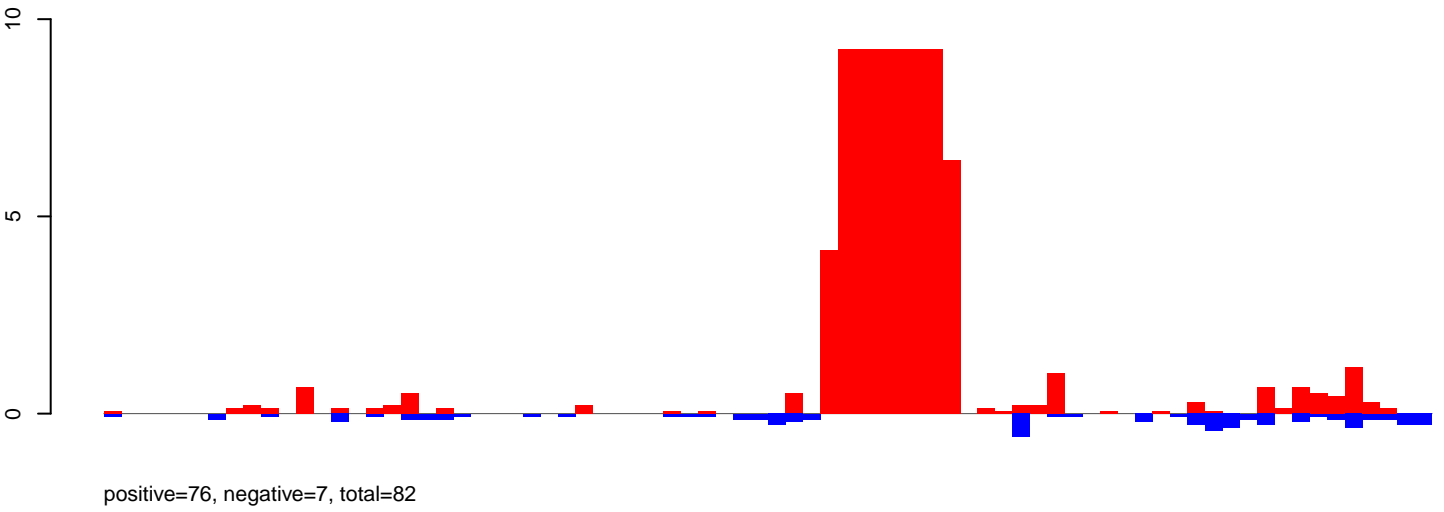
CuQuin_Testes_NL.24_35.rep



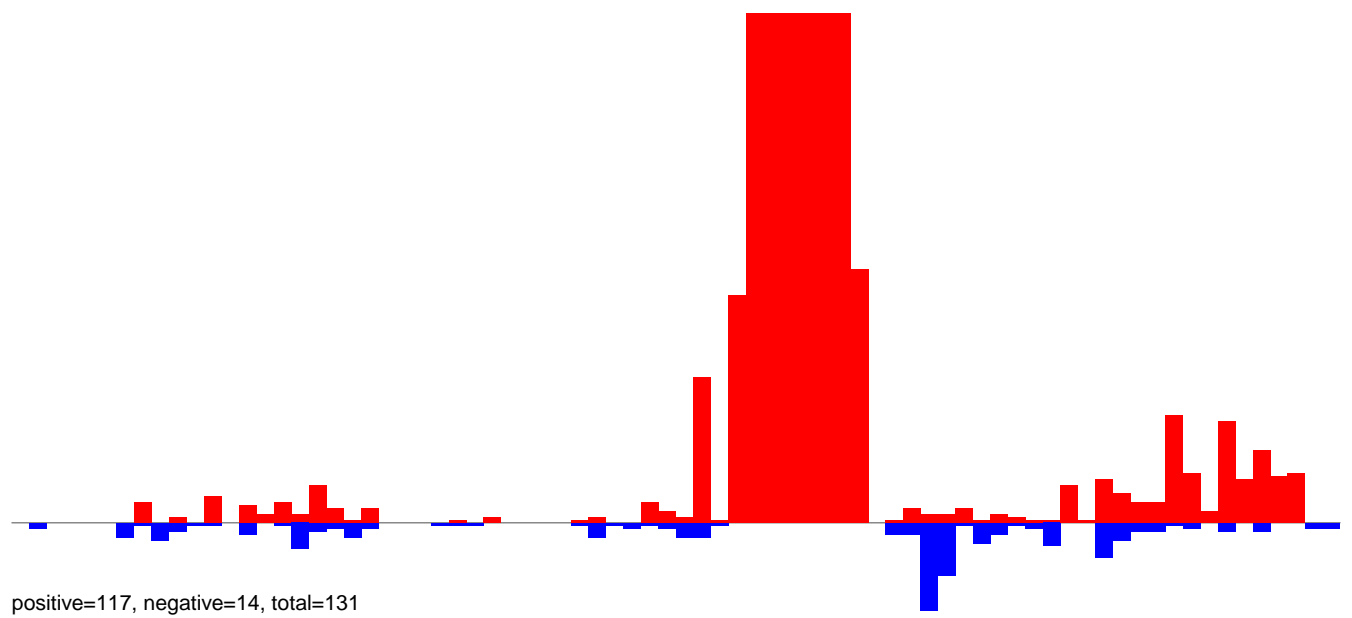
CuQuin_Testes_NL.rep



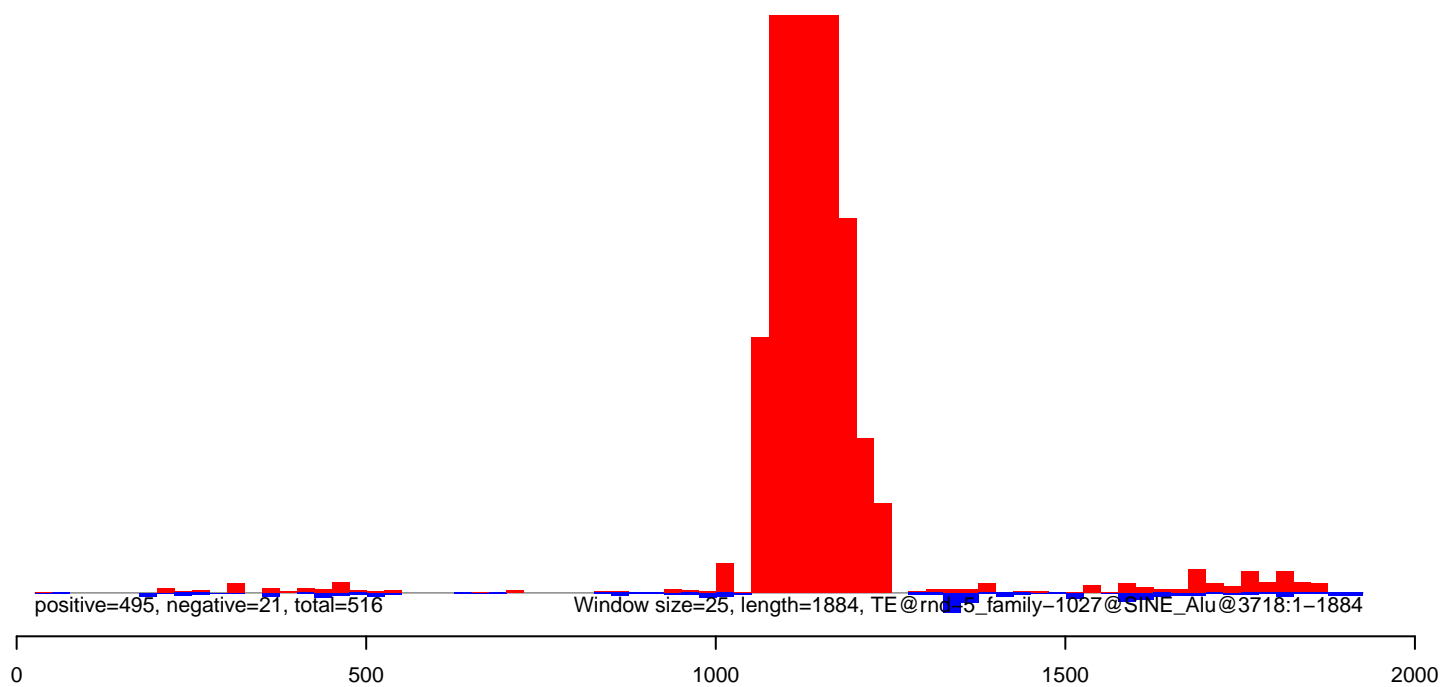
CuQuin_Testes_NL.18_23.rep



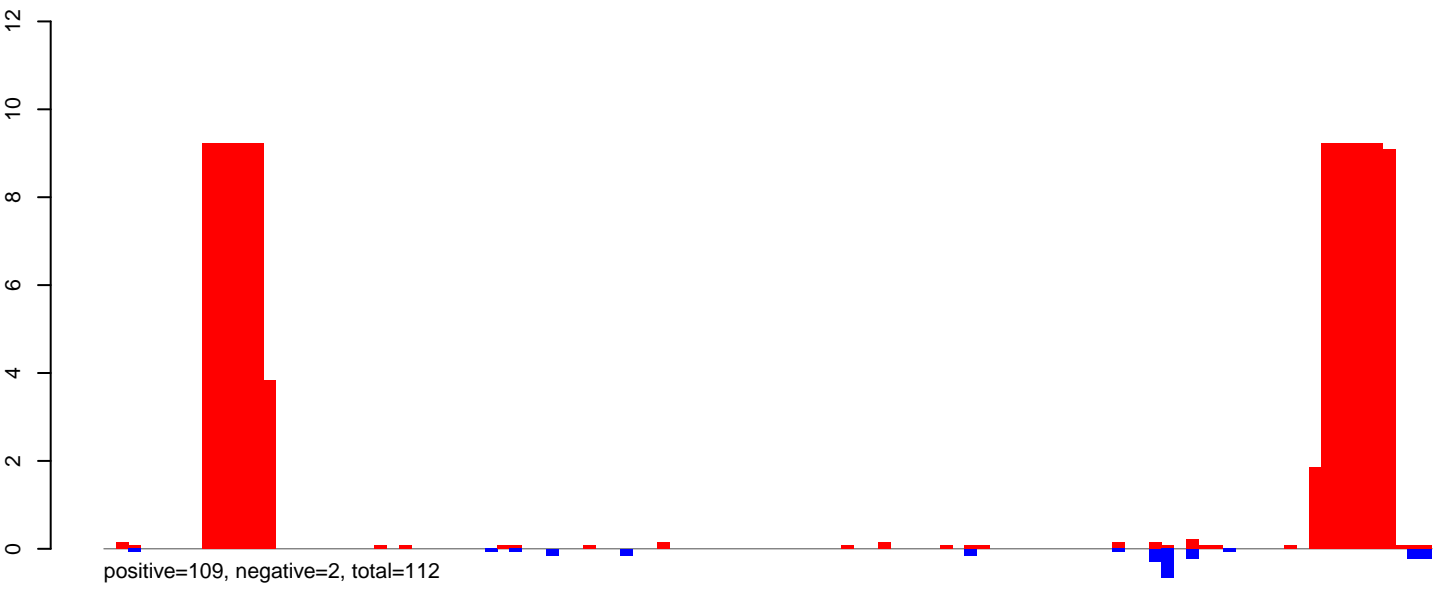
CuQuin_Testes_NL.24_35.rep



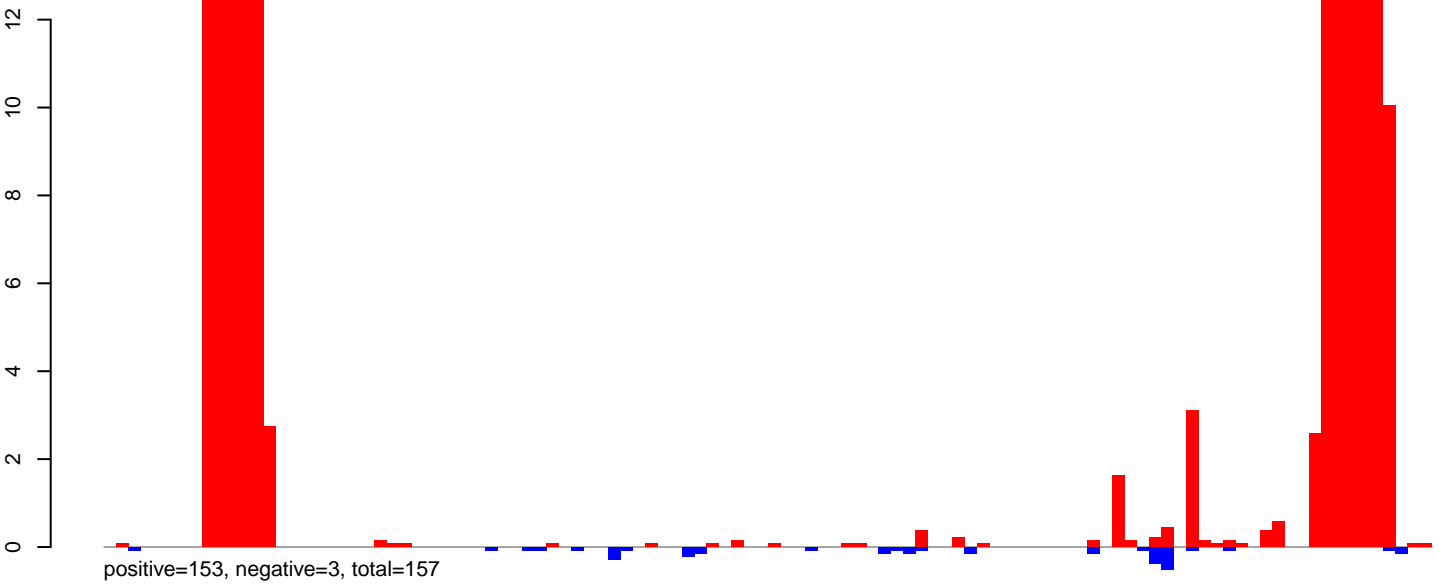
CuQuin_Testes_NL.rep



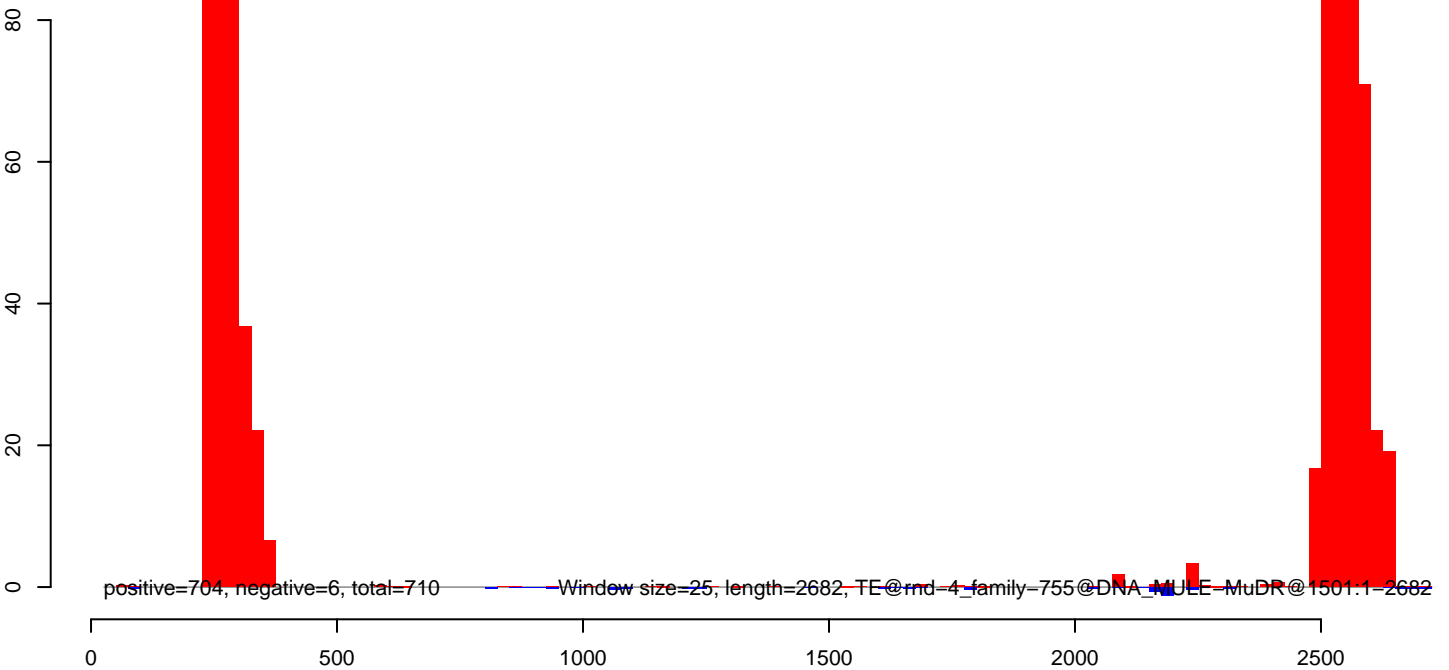
CuQuin_Testes_NL.18_23.rep



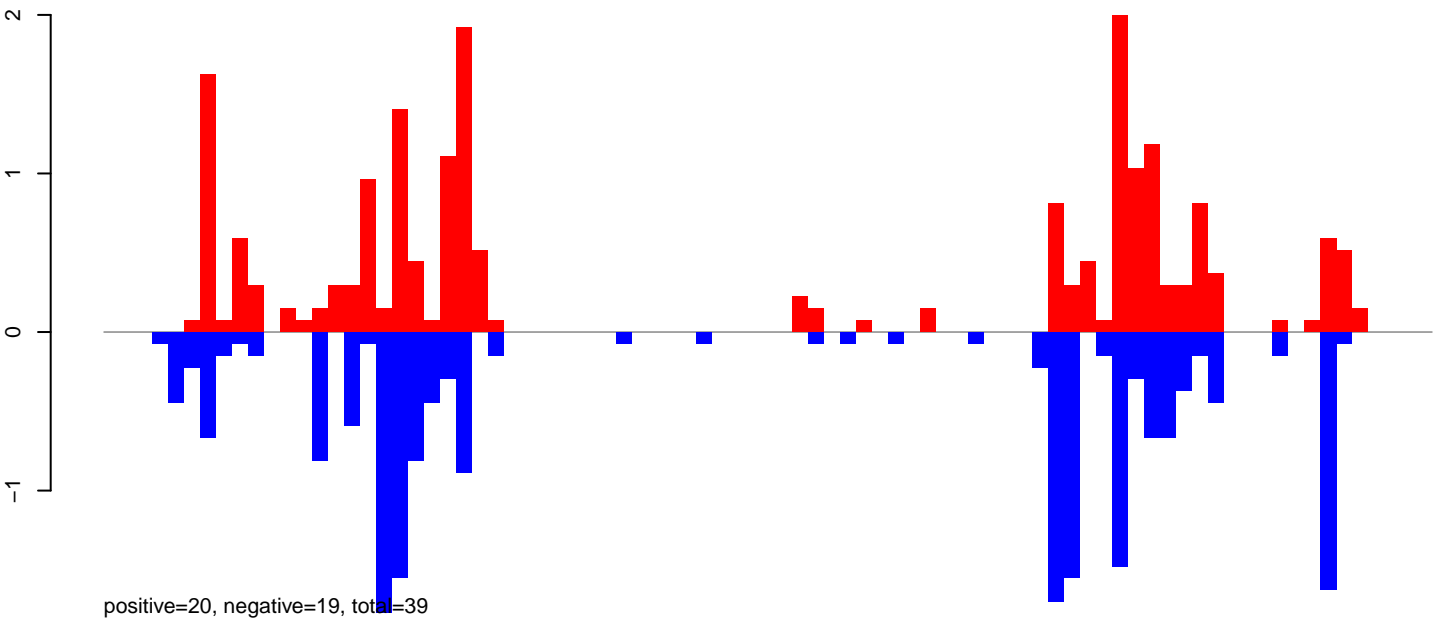
CuQuin_Testes_NL.24_35.rep



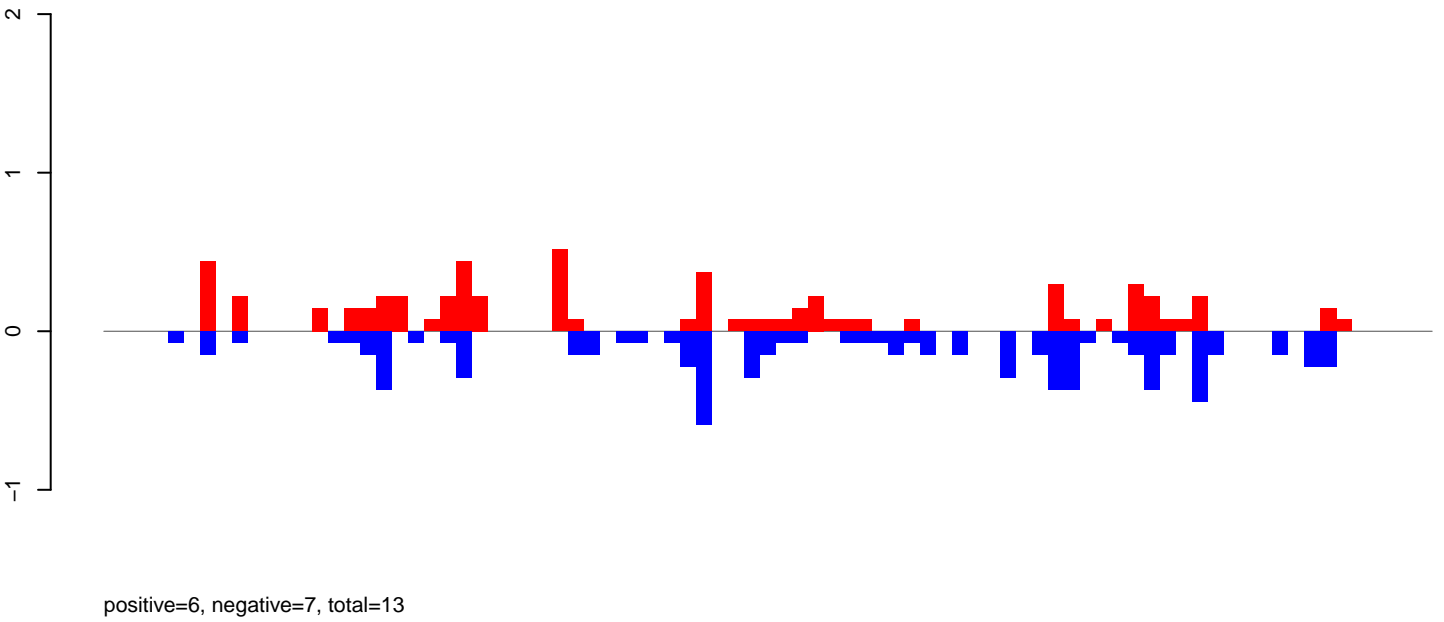
CuQuin_Testes_NL.rep



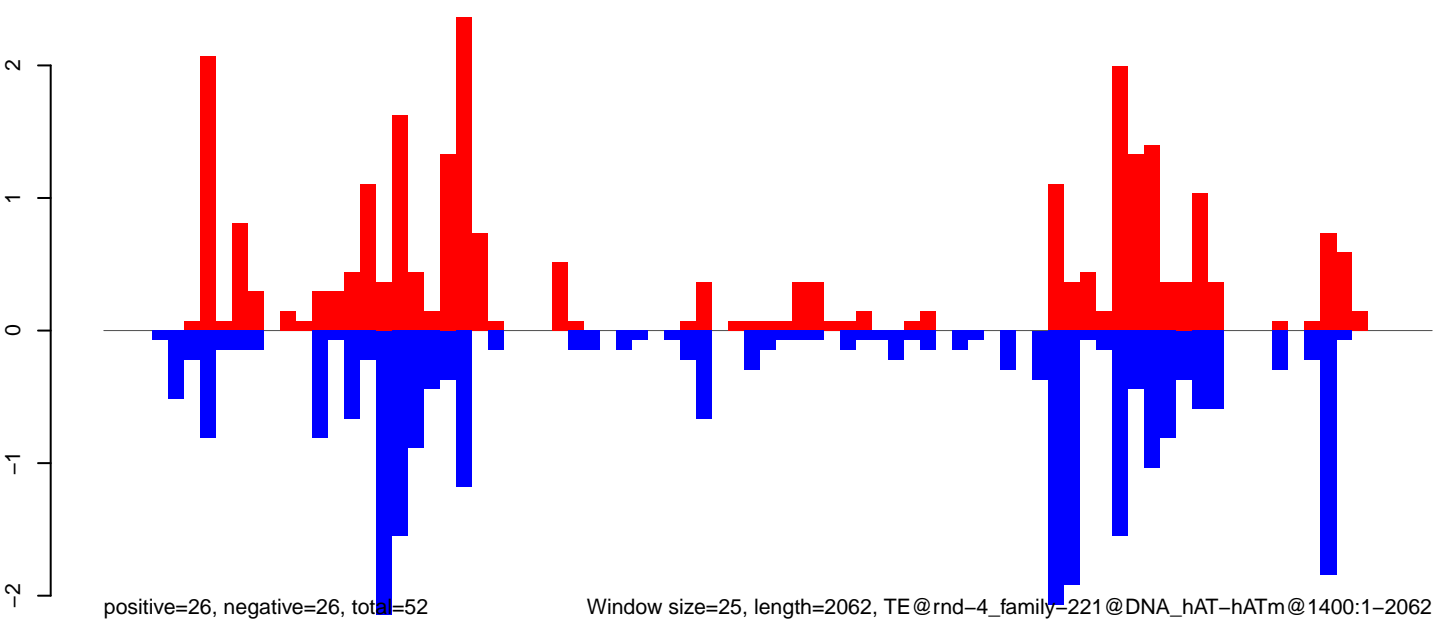
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



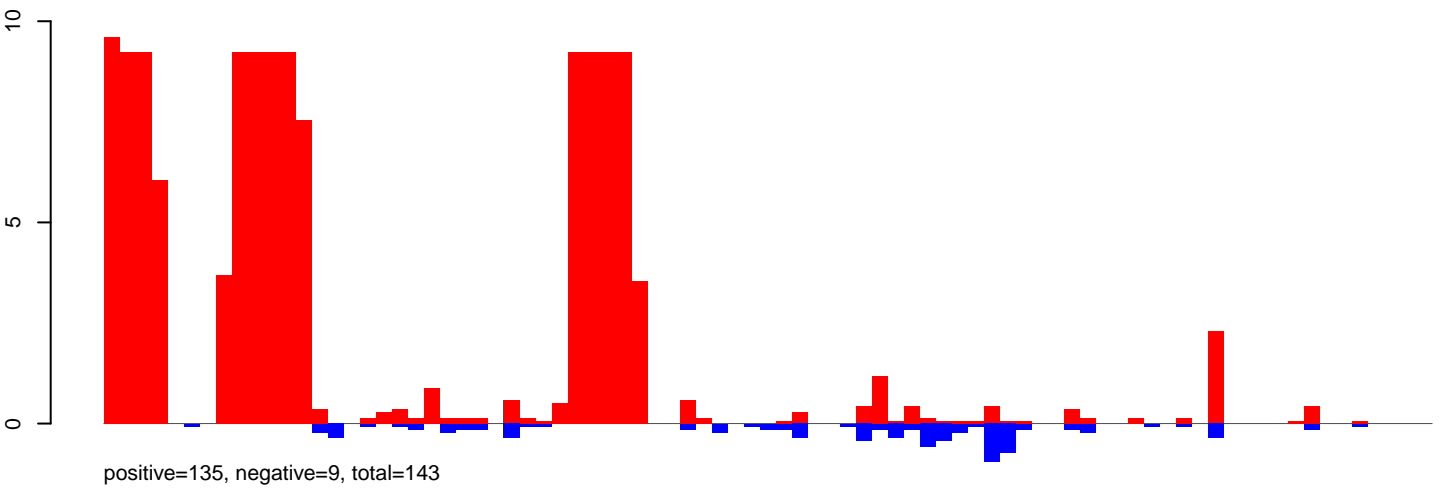
CuQuin_Testes_NL.rep



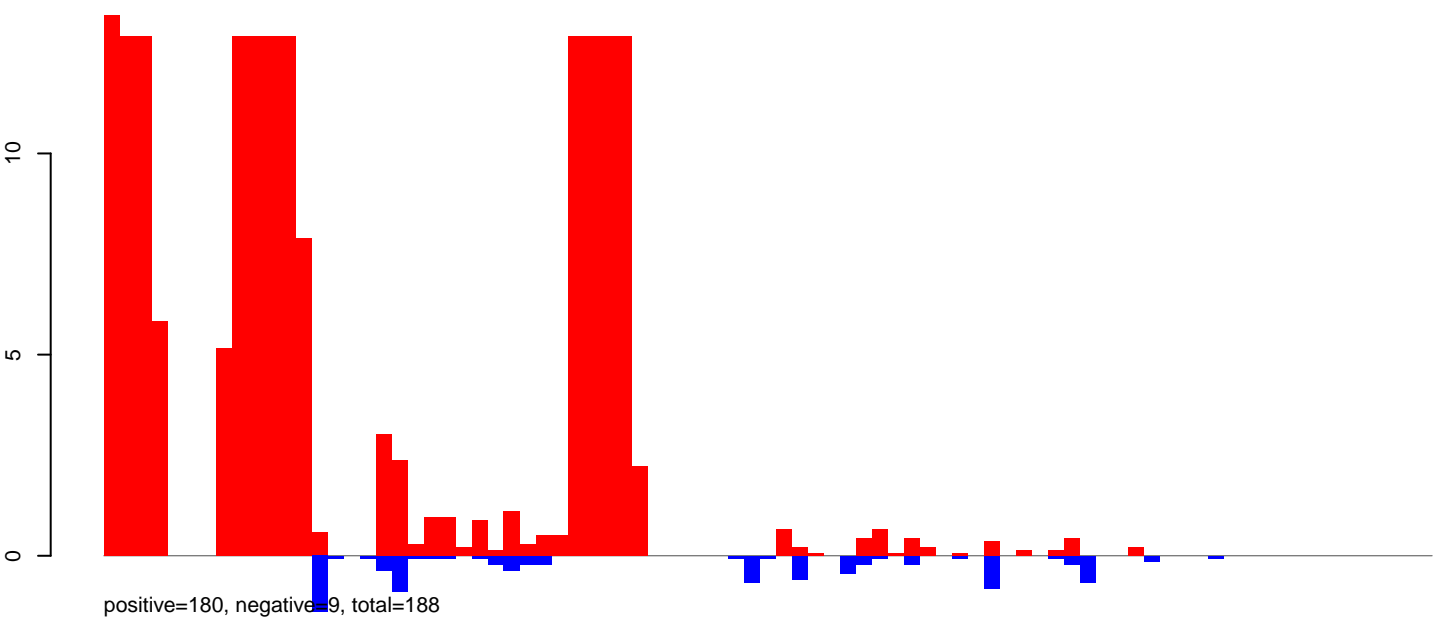
Window size=25, length=2062, TE@rnd-4_family-221@DNA_hAT-hATm@1400:1-2062

0 500 1000 1500 2000

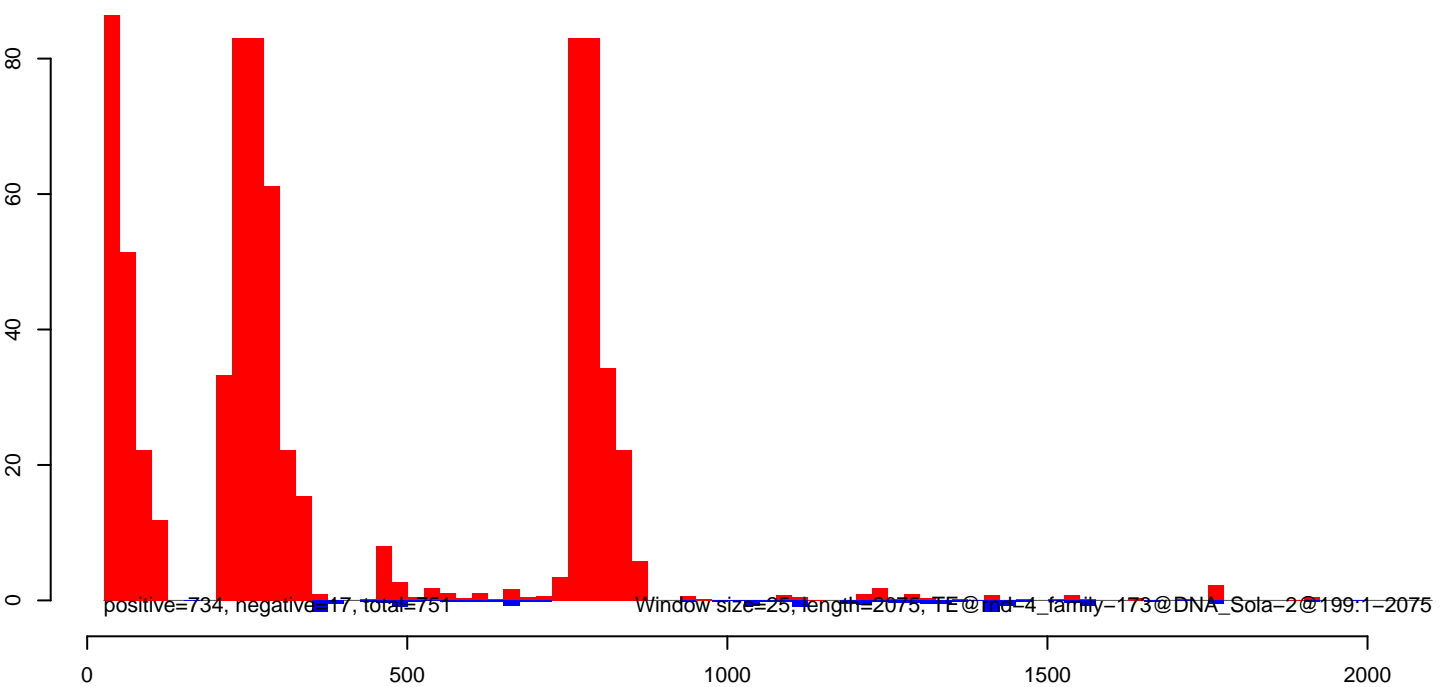
CuQuin_Testes_NL.18_23.rep



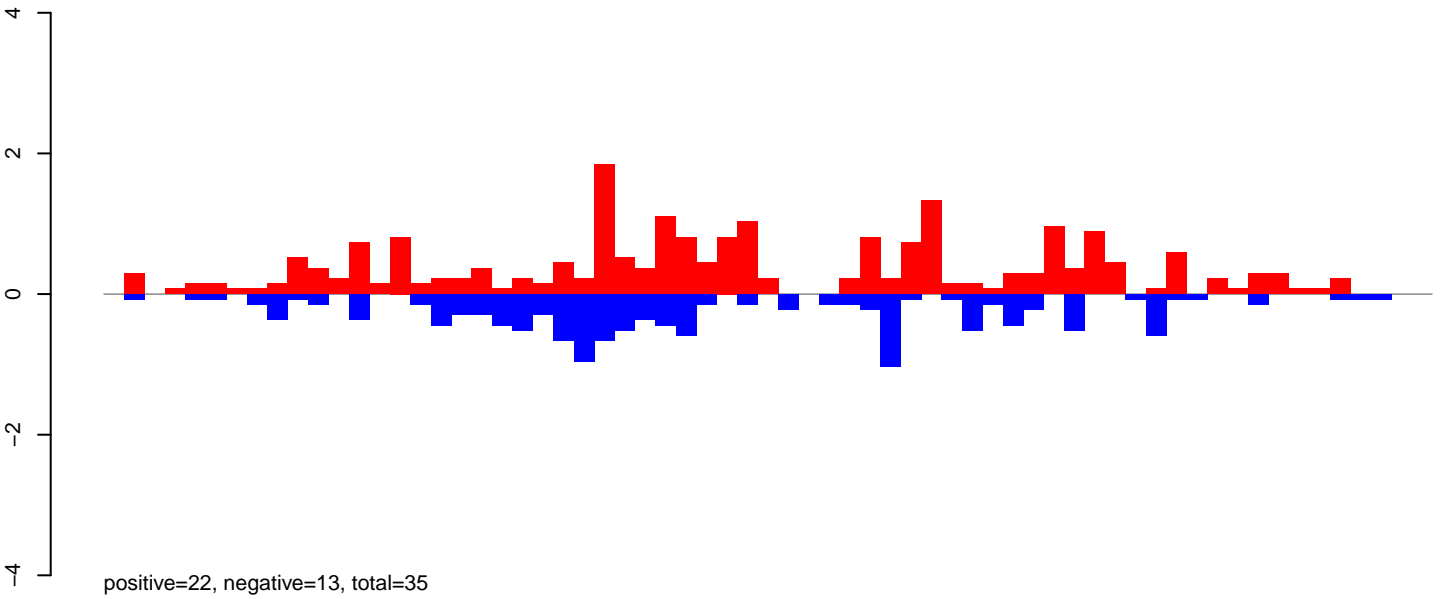
CuQuin_Testes_NL.24_35.rep



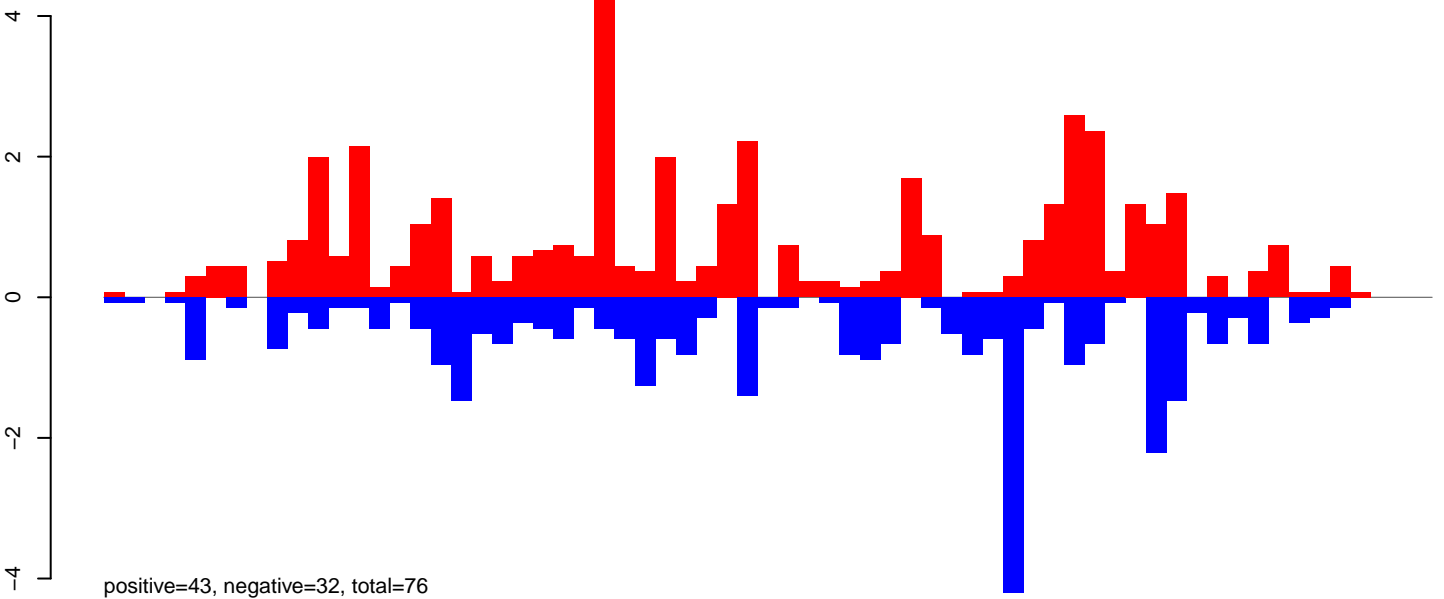
CuQuin_Testes_NL.rep



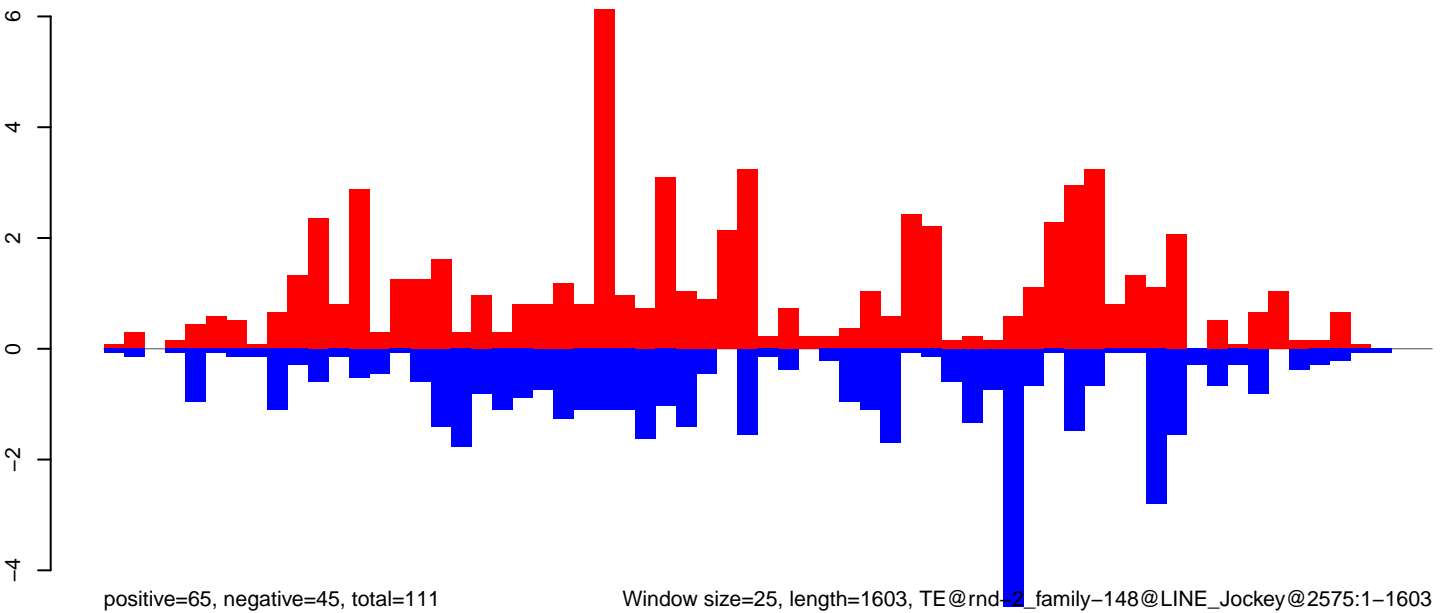
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



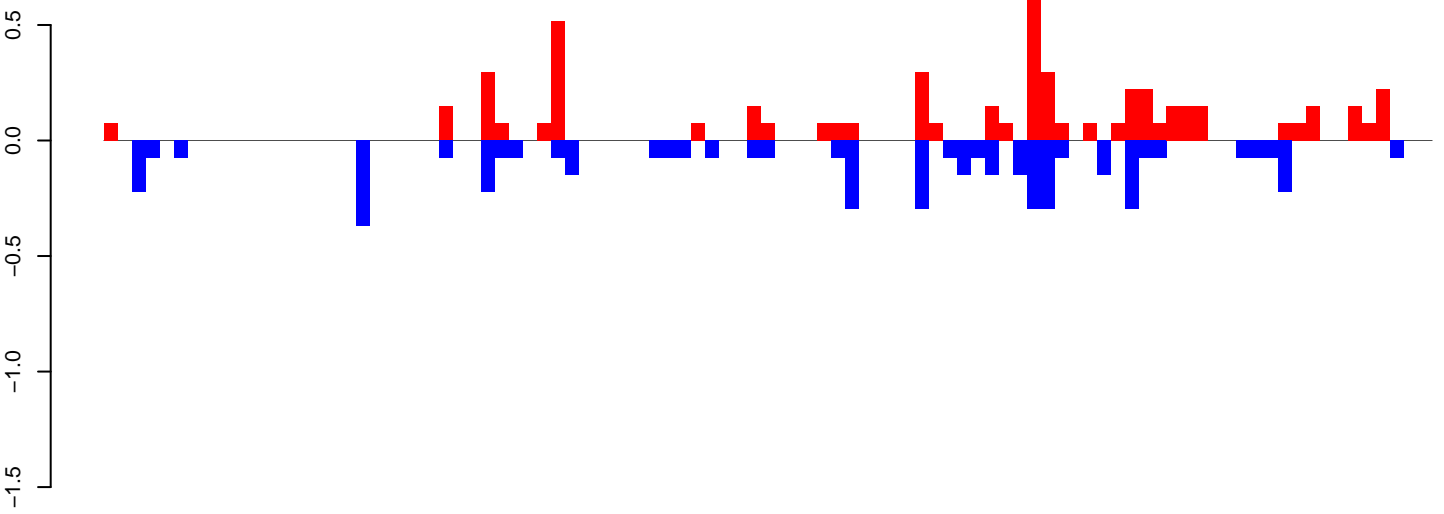
CuQuin_Testes_NL.rep



Window size=25, length=1603, TE@rnd-2_family-148@LINE_Jockey@2575:1-1603

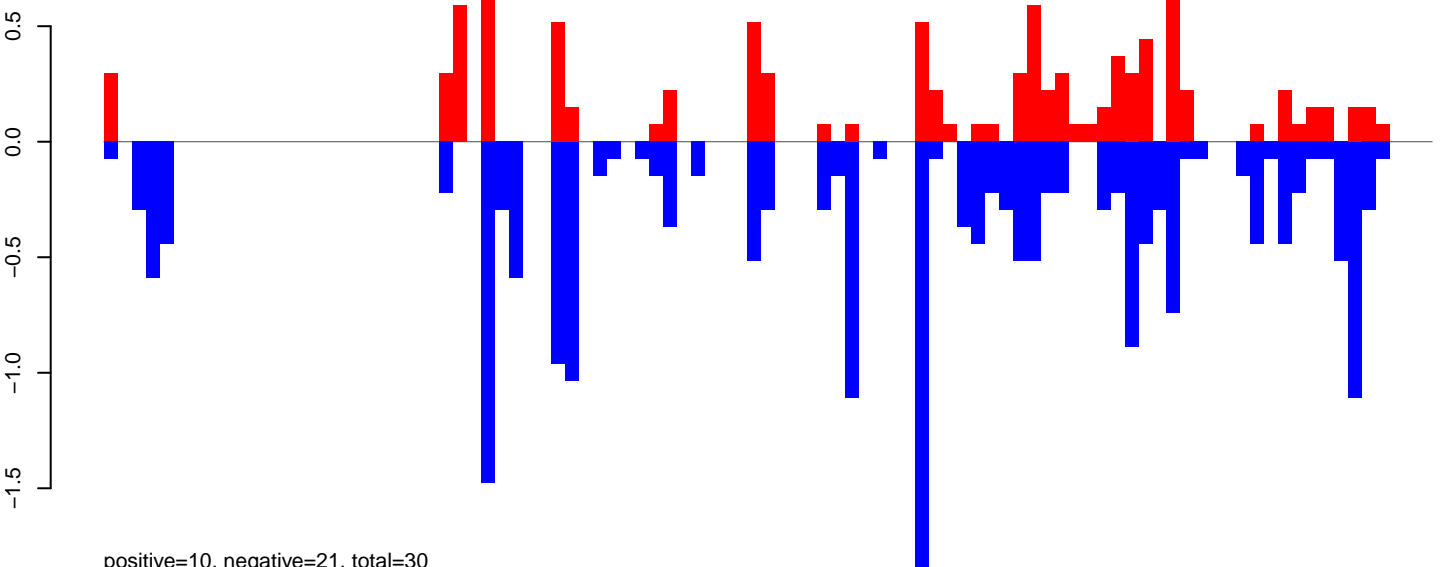
0 500 1000 1500

CuQuin_Testes_NL.18_23.rep



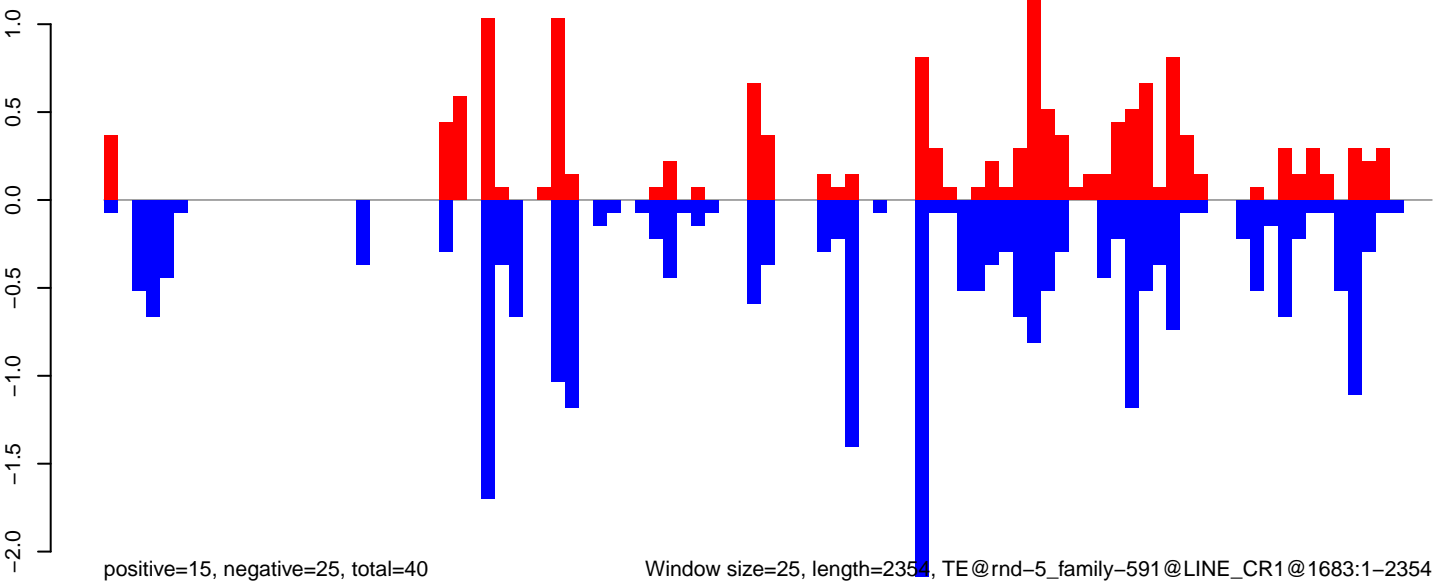
positive=5, negative=5, total=10

CuQuin_Testes_NL.24_35.rep



positive=10, negative=21, total=30

CuQuin_Testes_NL.rep

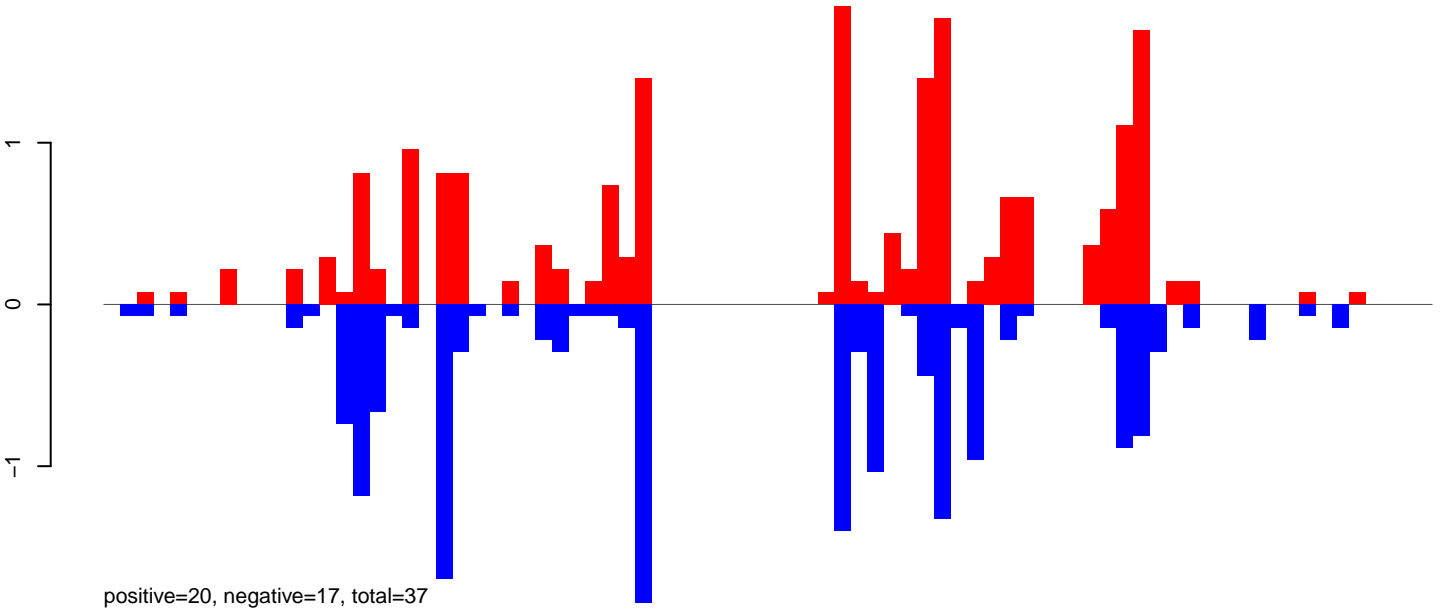


positive=15, negative=25, total=40

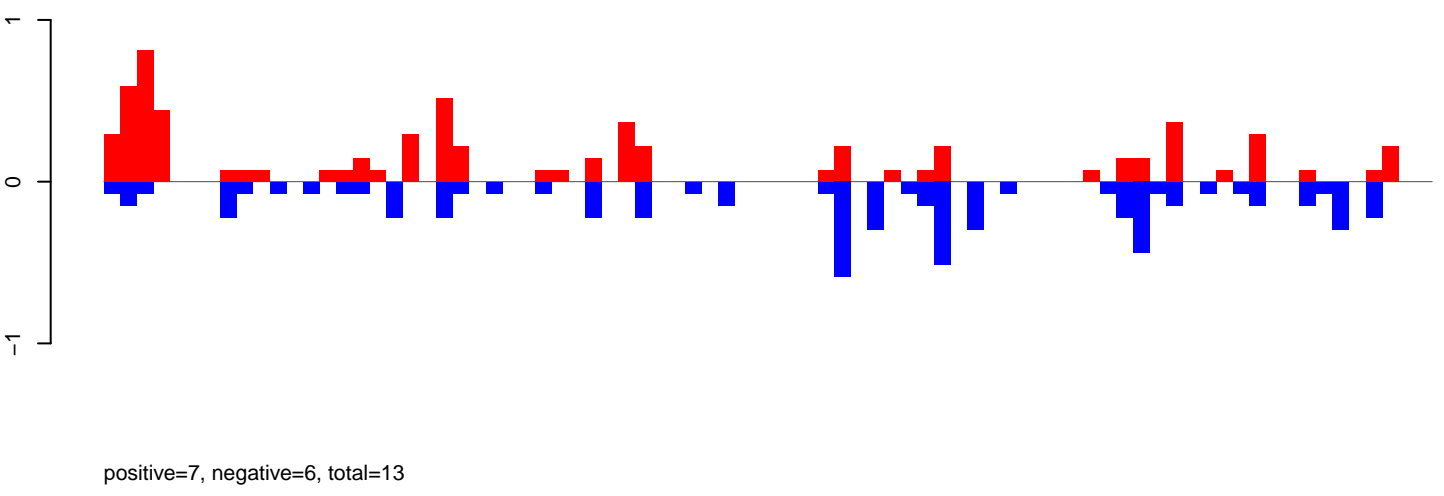
Window size=25, length=2354, TE@rnd-5_family-591@LINE_CR1@1683:1-2354

0 500 1000 1500 2000

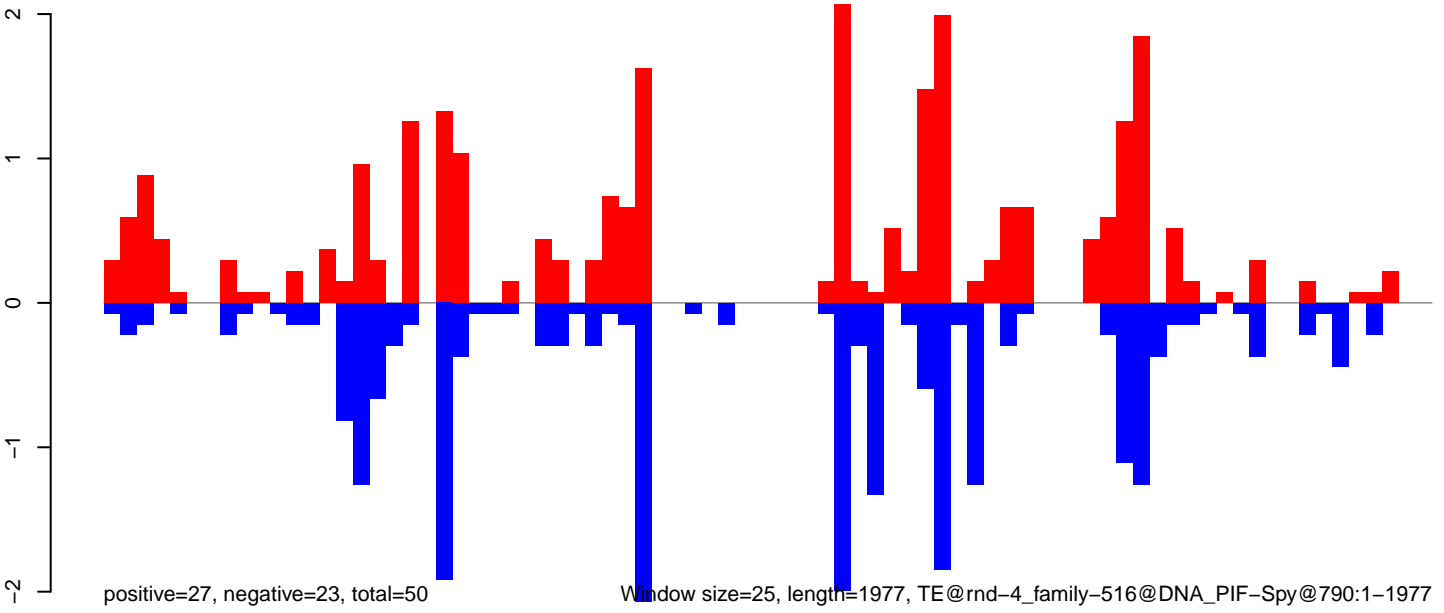
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

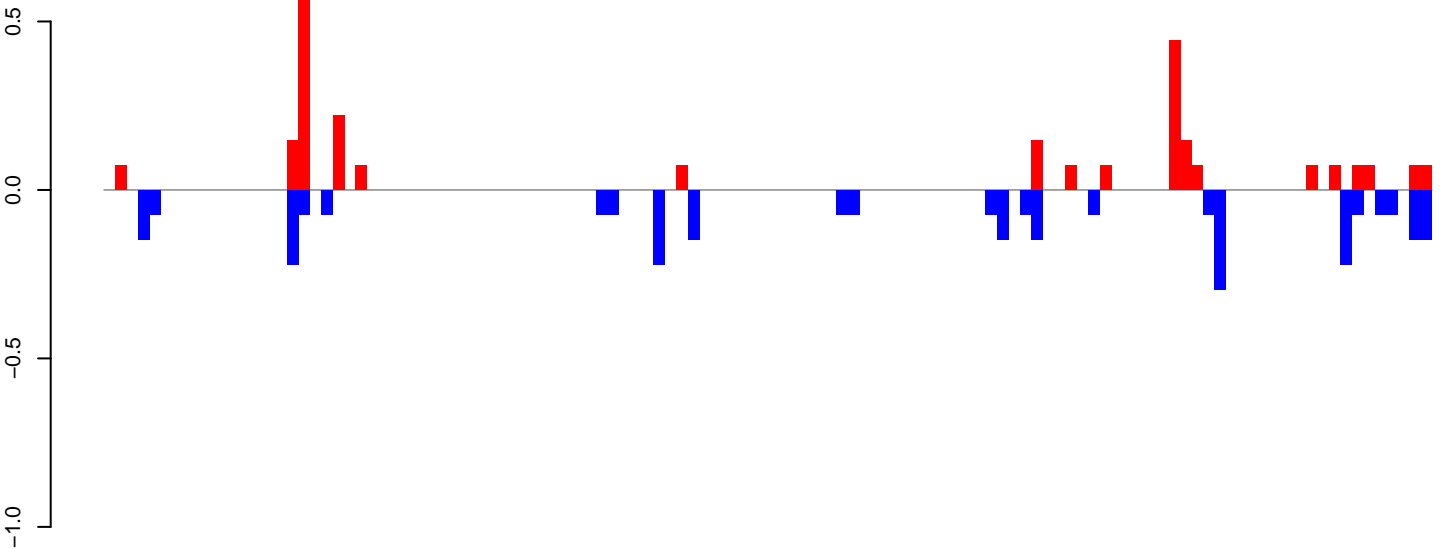


CuQuin_Testes_NL.rep



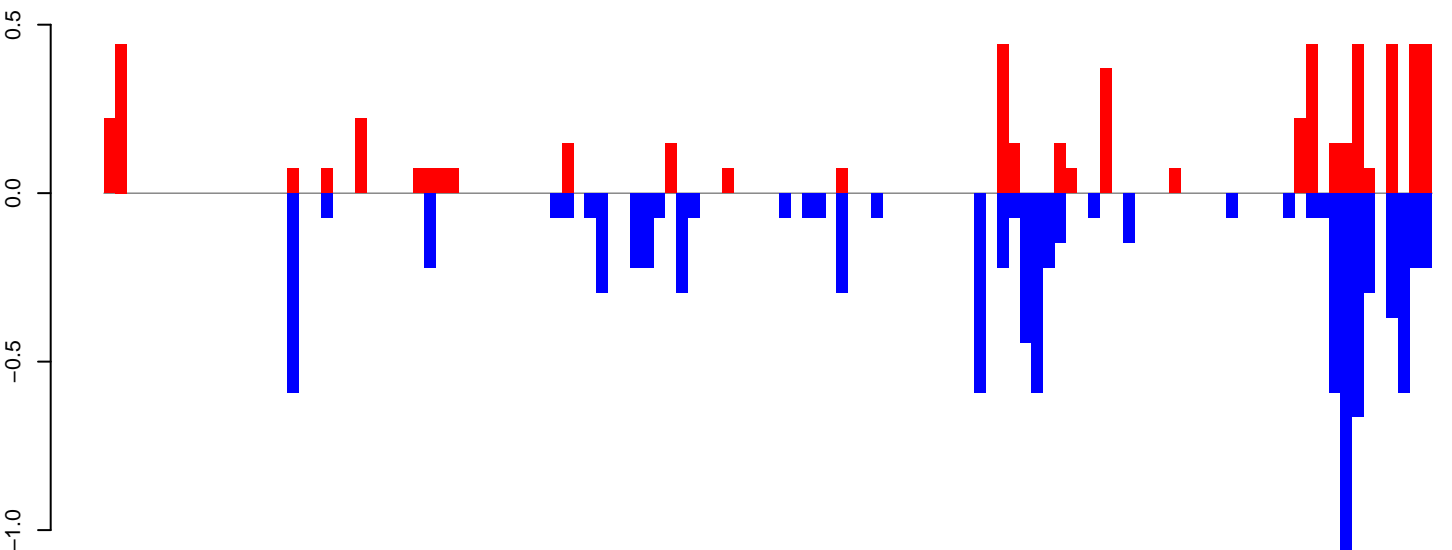
Window size=25, length=1977, TE@rnd-4_family-516@DNA_PIF-Spy@790:1-1977

CuQuin_Testes_NL.18_23.rep



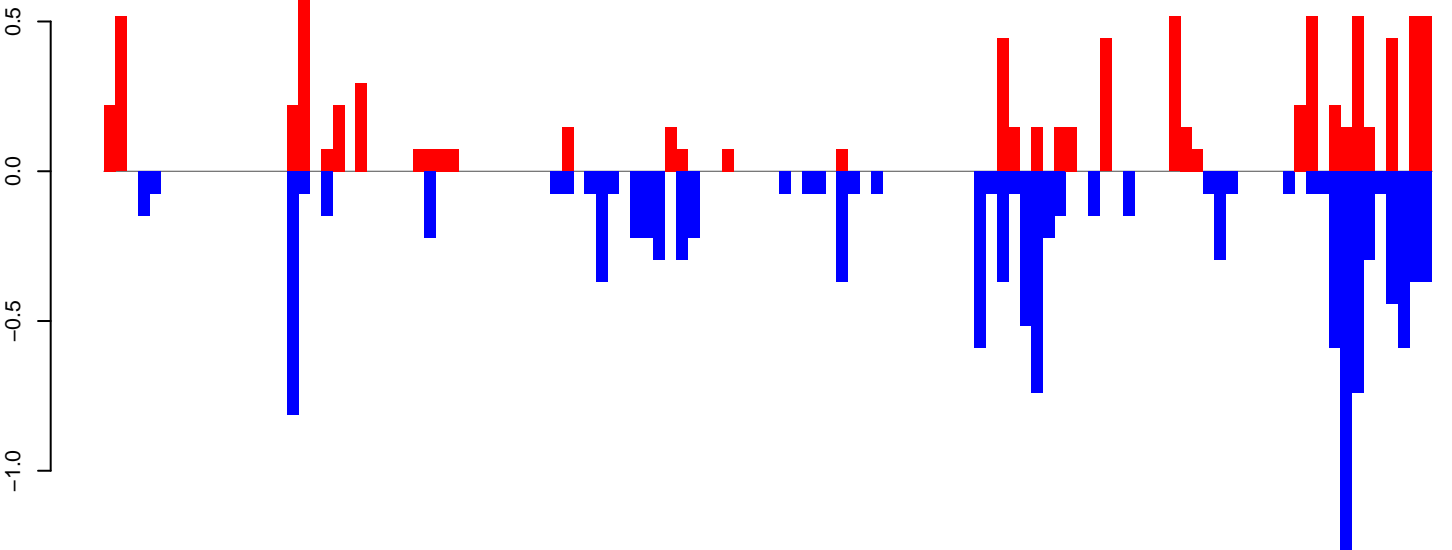
positive=3, negative=3, total=5

CuQuin_Testes_NL.24_35.rep



positive=6, negative=10, total=16

CuQuin_Testes_NL.rep

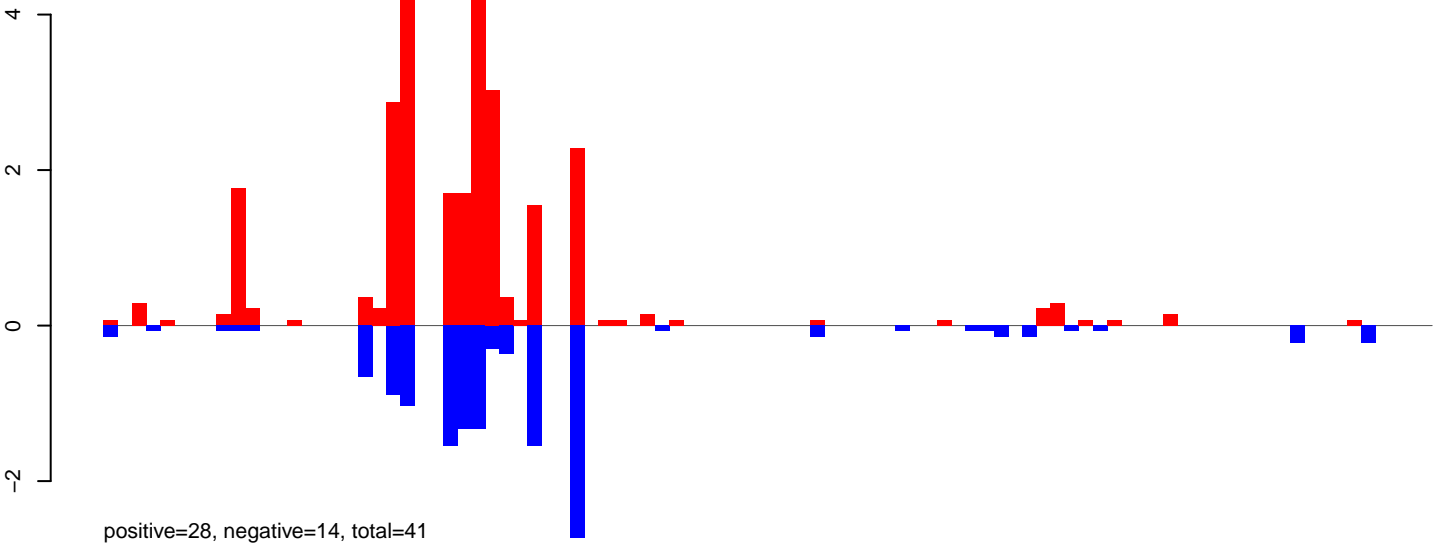


positive=8, negative=13, total=21

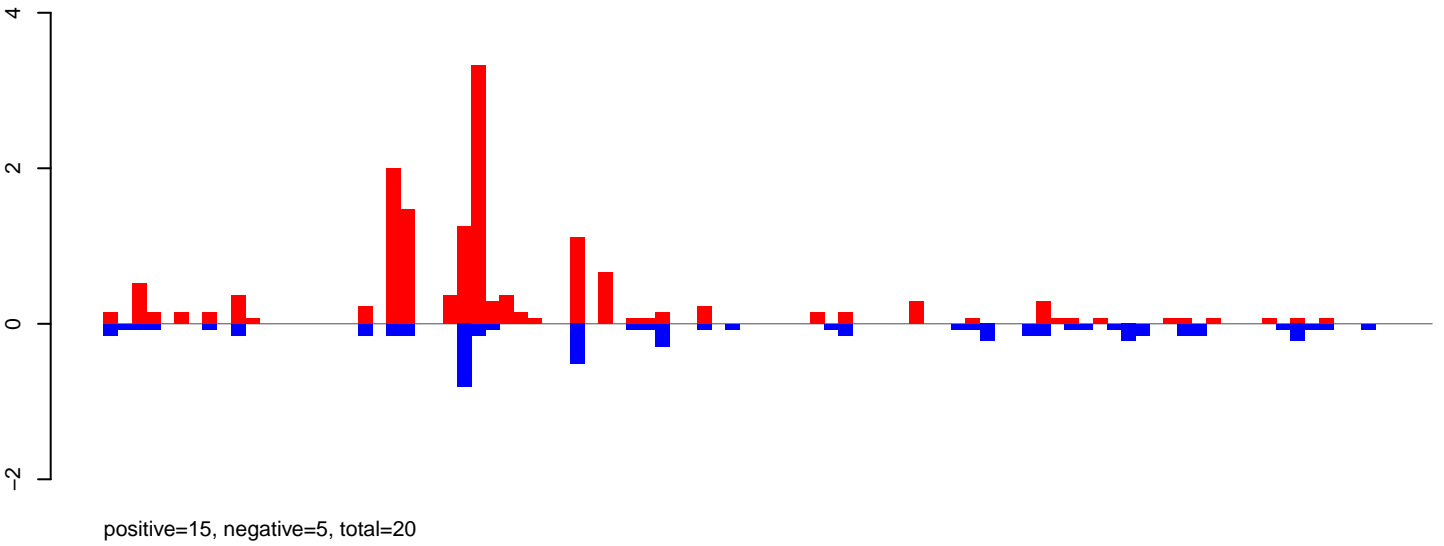
Window size=25, length=2887, TE@rnd-3_family-82@DNA@720:1-2887

0 500 1000 1500 2000 2500 3000

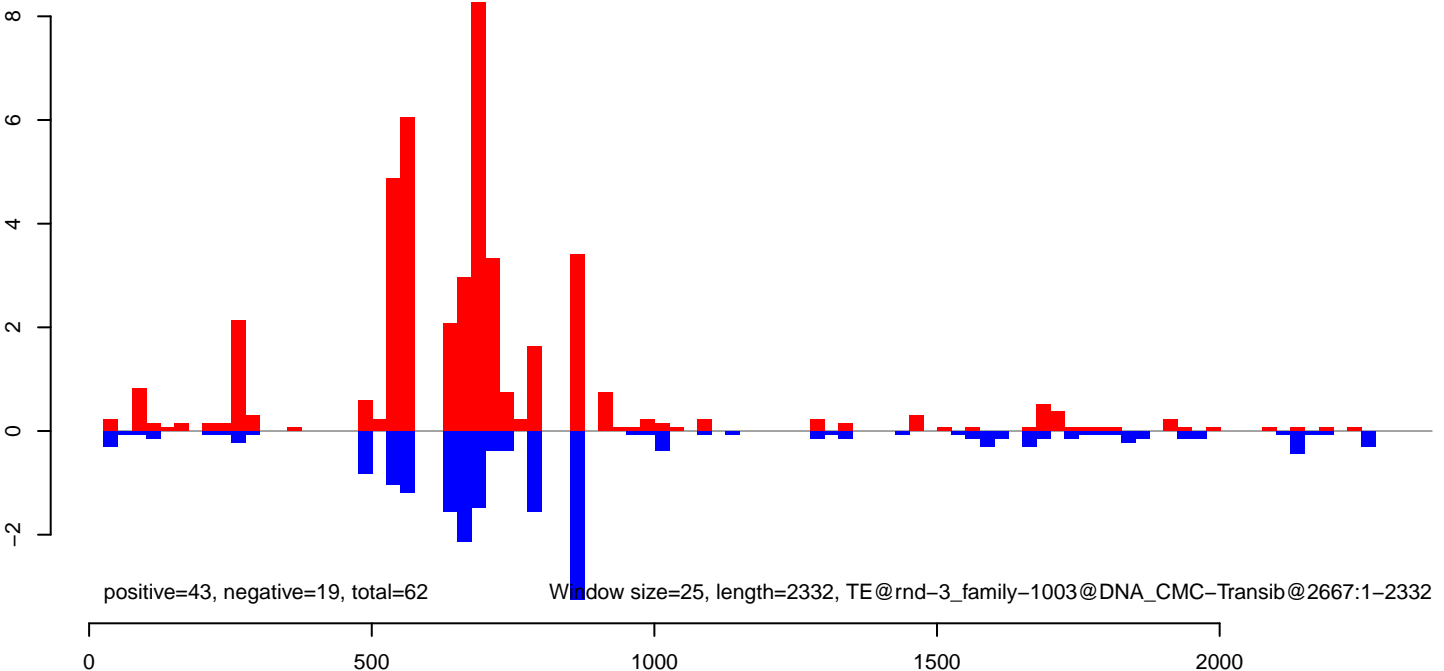
CuQuin_Testes_NL.18_23.rep



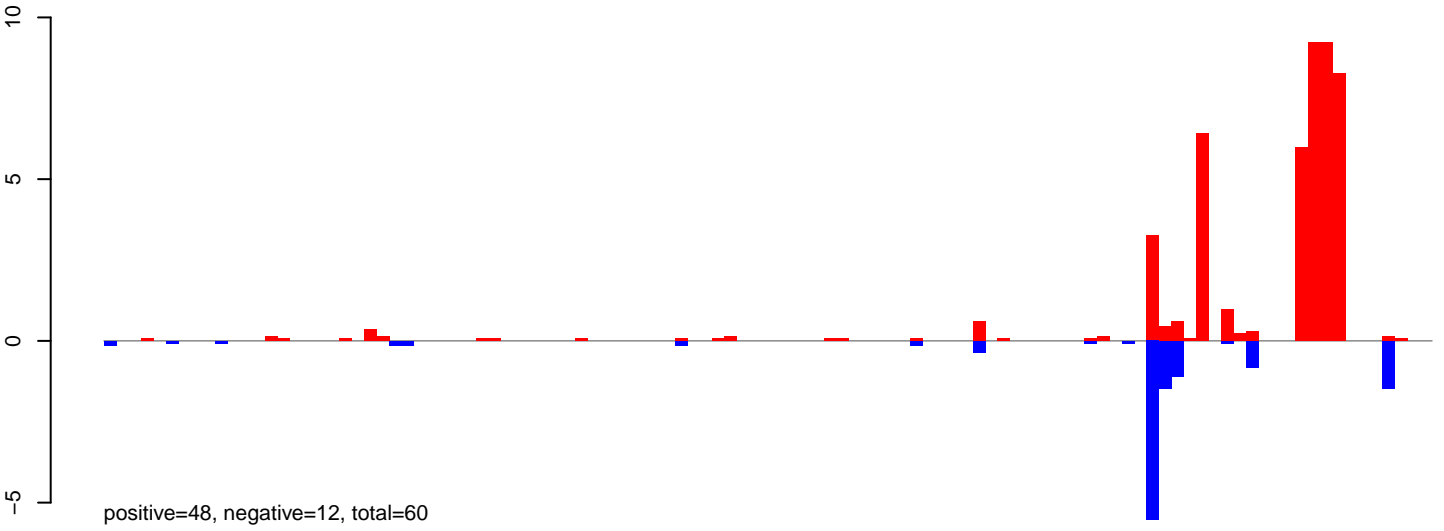
CuQuin_Testes_NL.24_35.rep



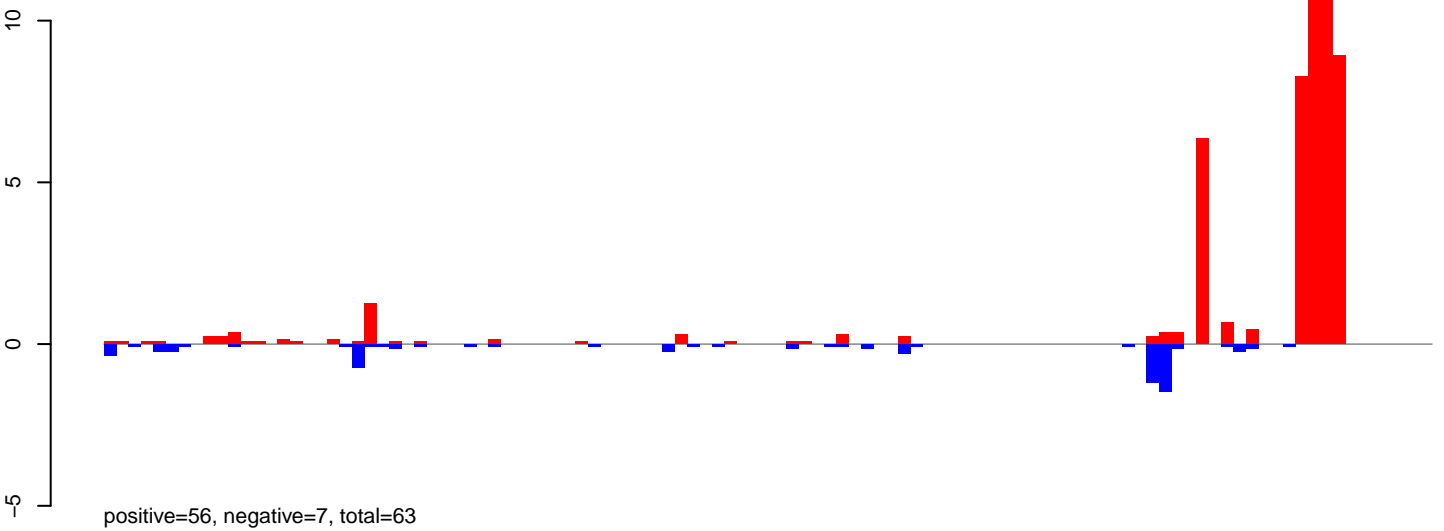
CuQuin_Testes_NL.rep



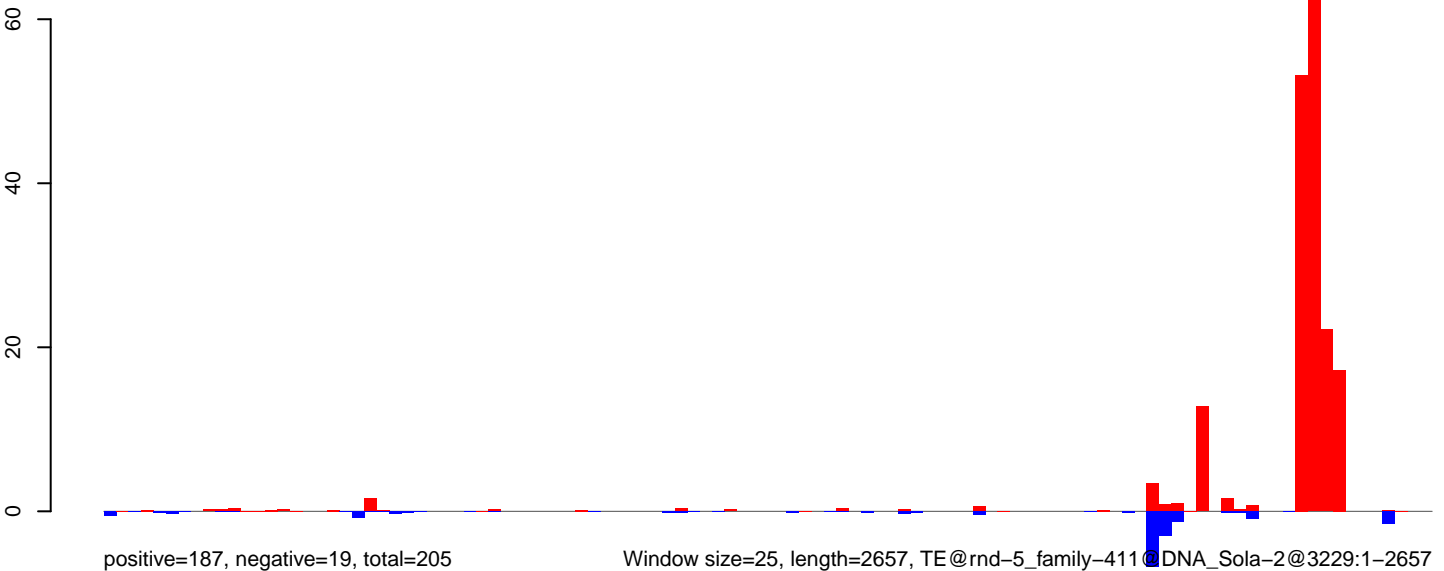
CuQuin_Testes_NL.18_23.rep



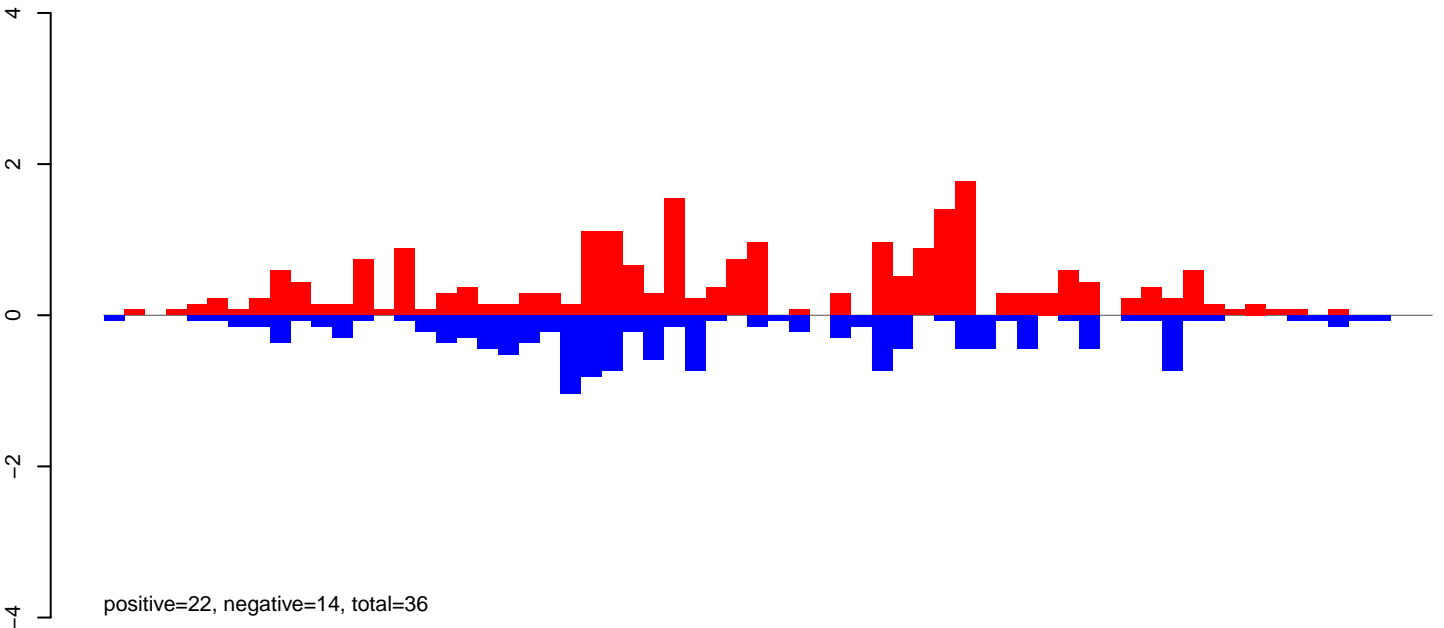
CuQuin_Testes_NL.24_35.rep



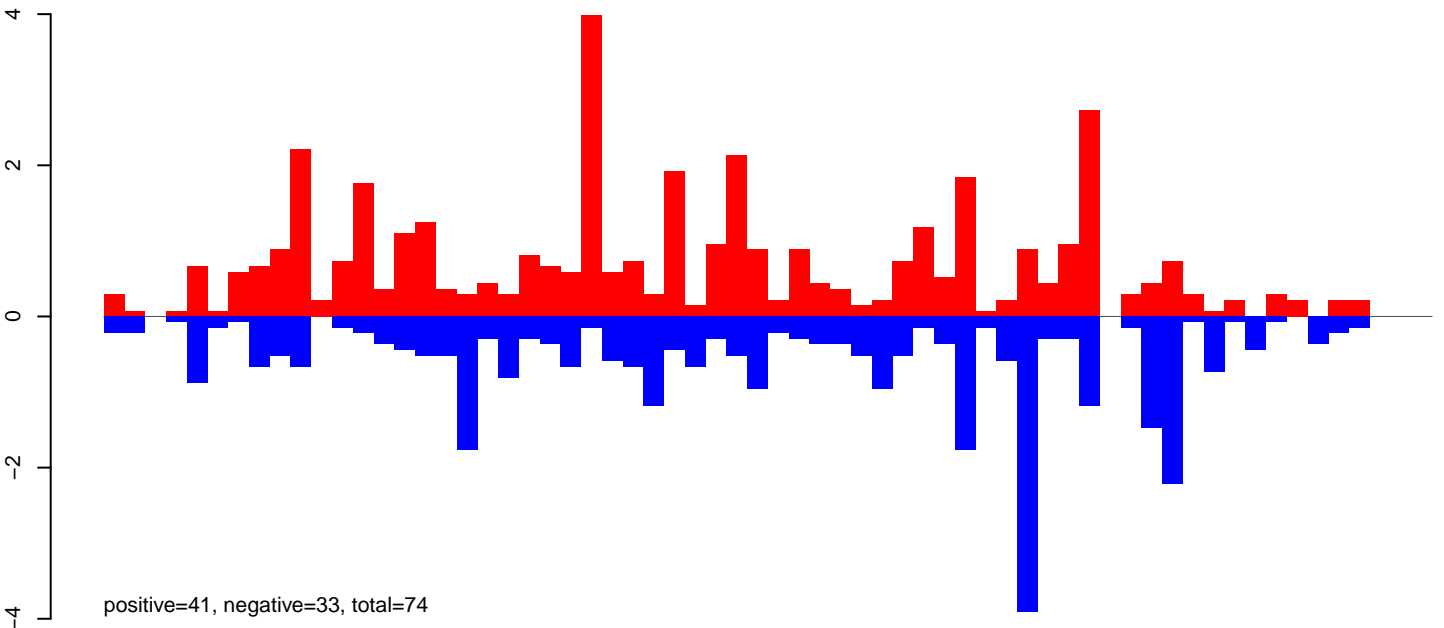
CuQuin_Testes_NL.rep



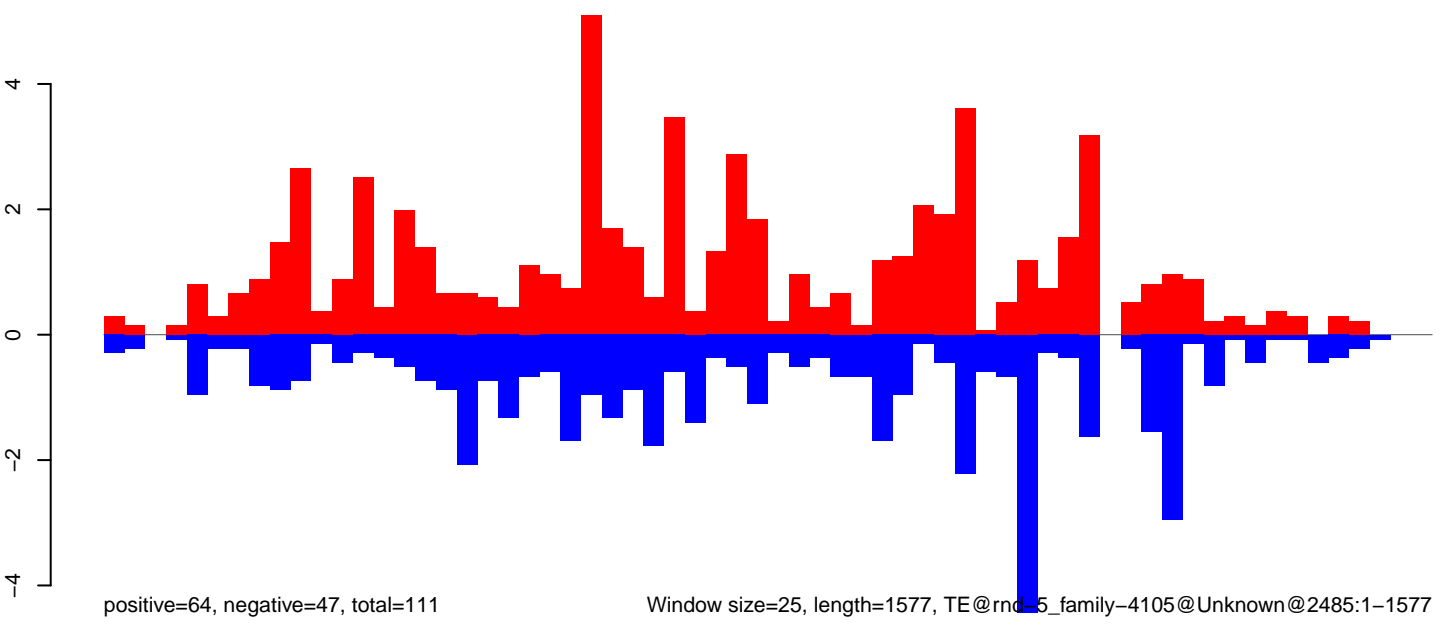
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



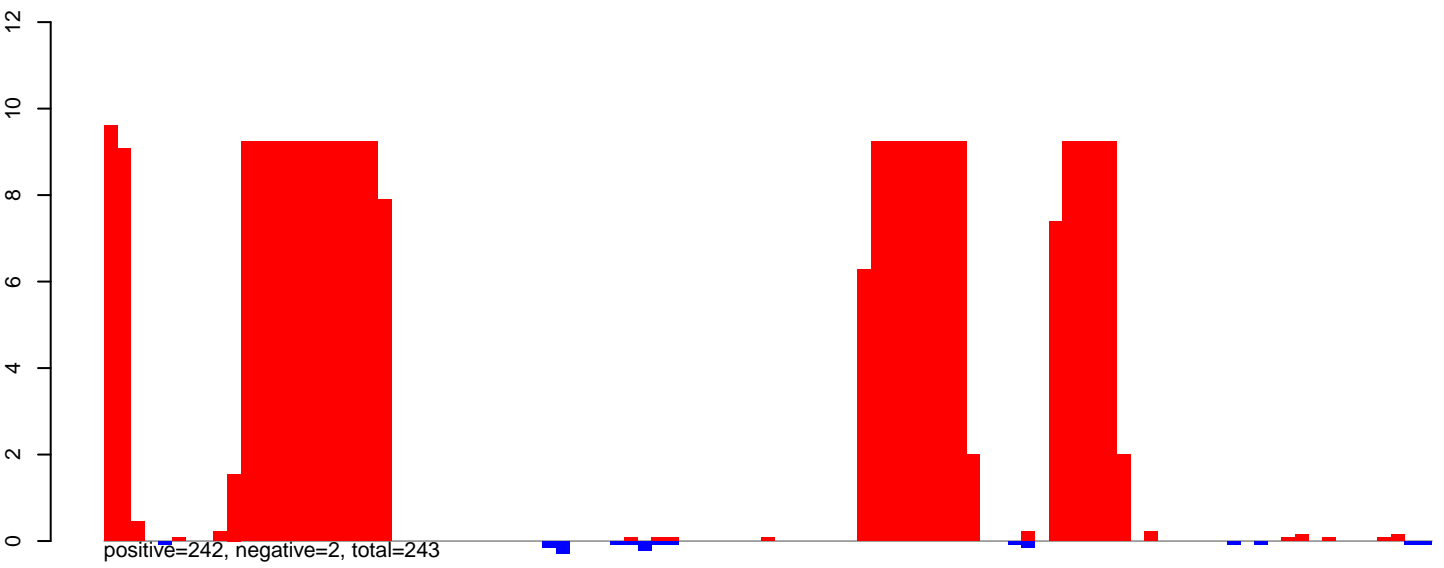
CuQuin_Testes_NL.rep



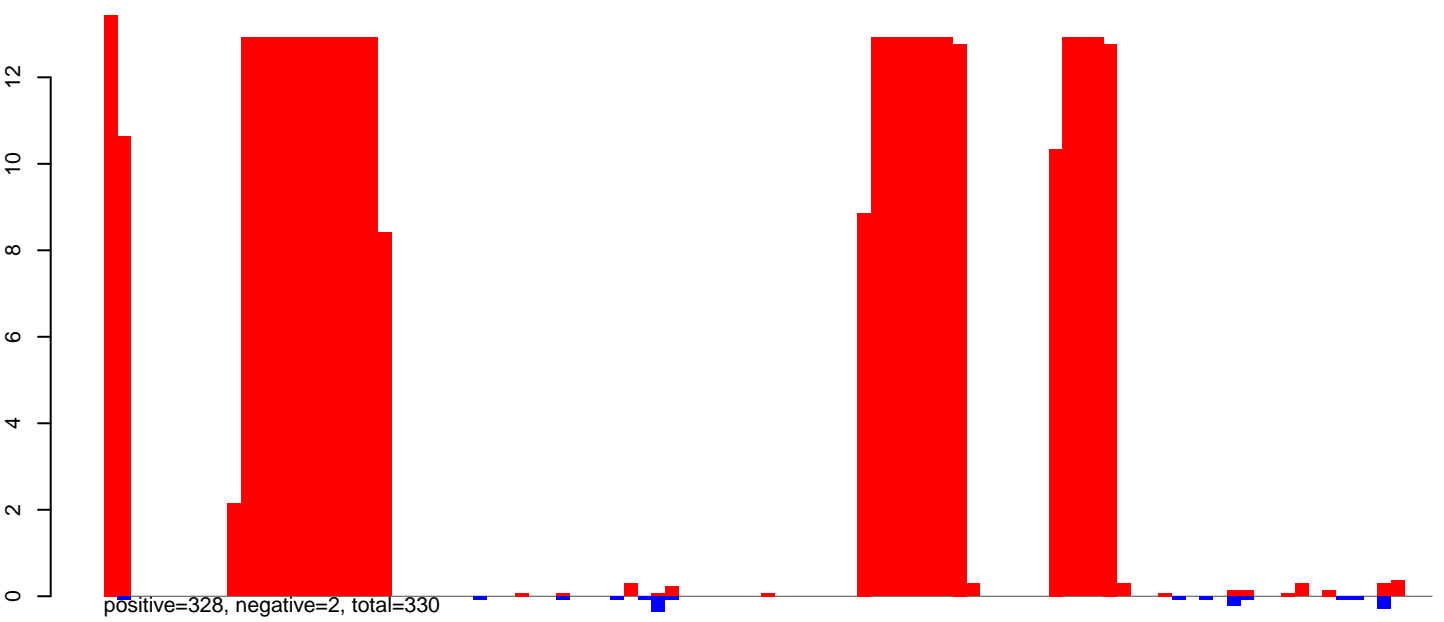
Window size=25, length=1577, TE@rnd_5_family-4105@Unknown@2485:1-1577

0 500 1000 1500

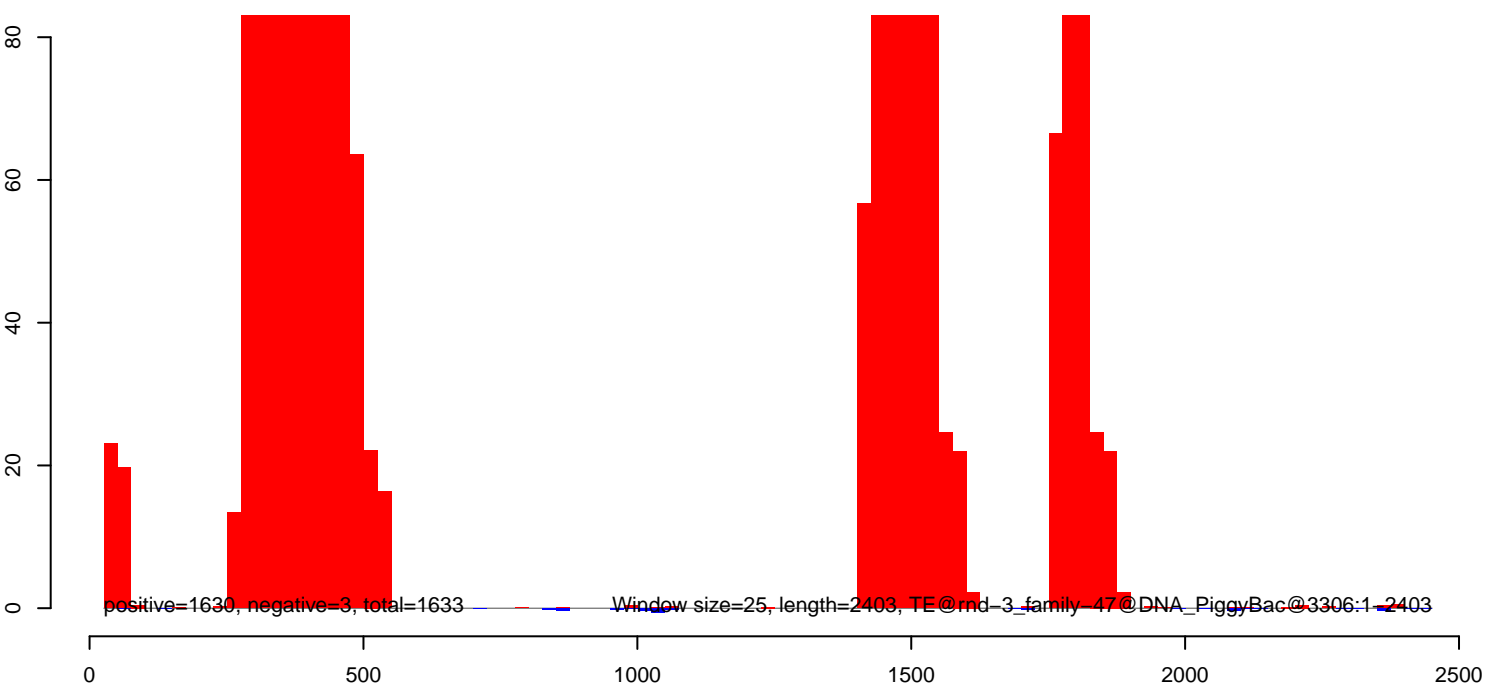
CuQuin_Testes_NL.18_23.rep



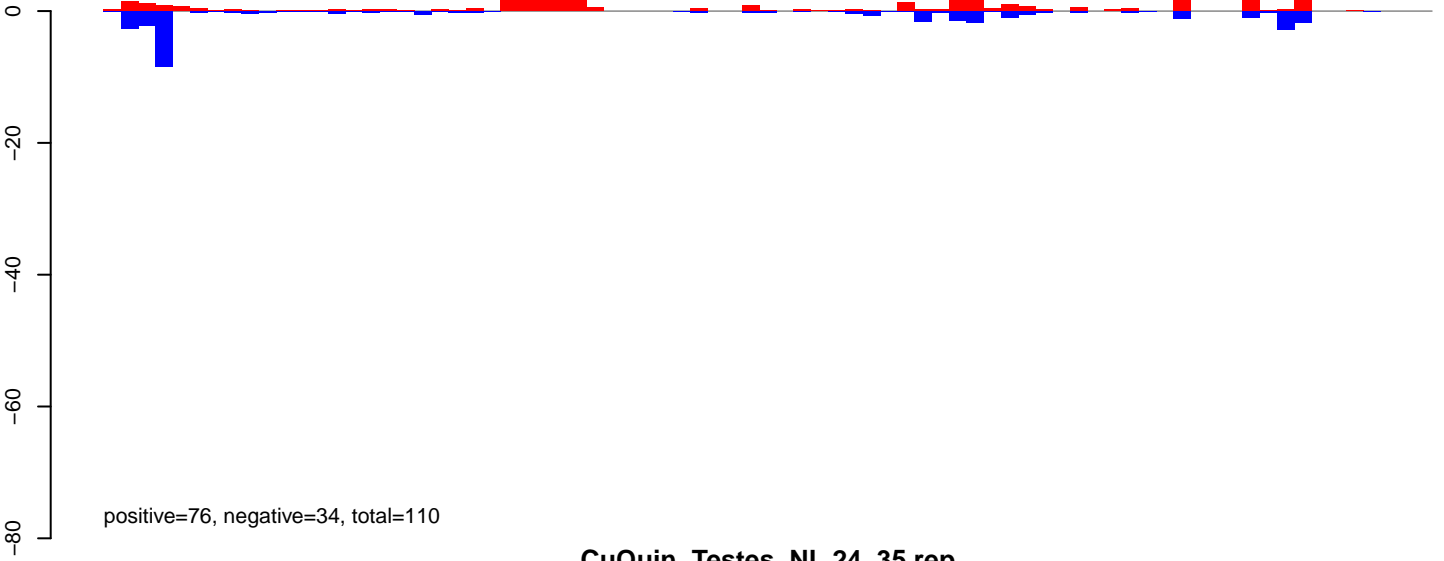
CuQuin_Testes_NL.24_35.rep



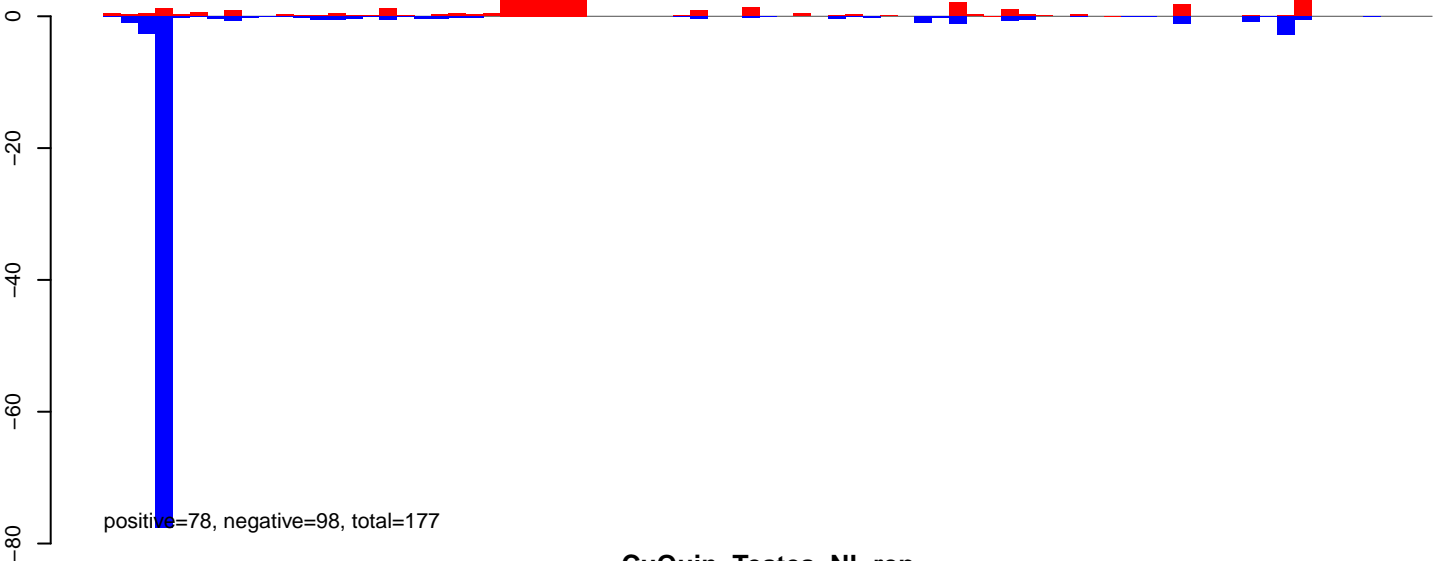
CuQuin_Testes_NL.rep



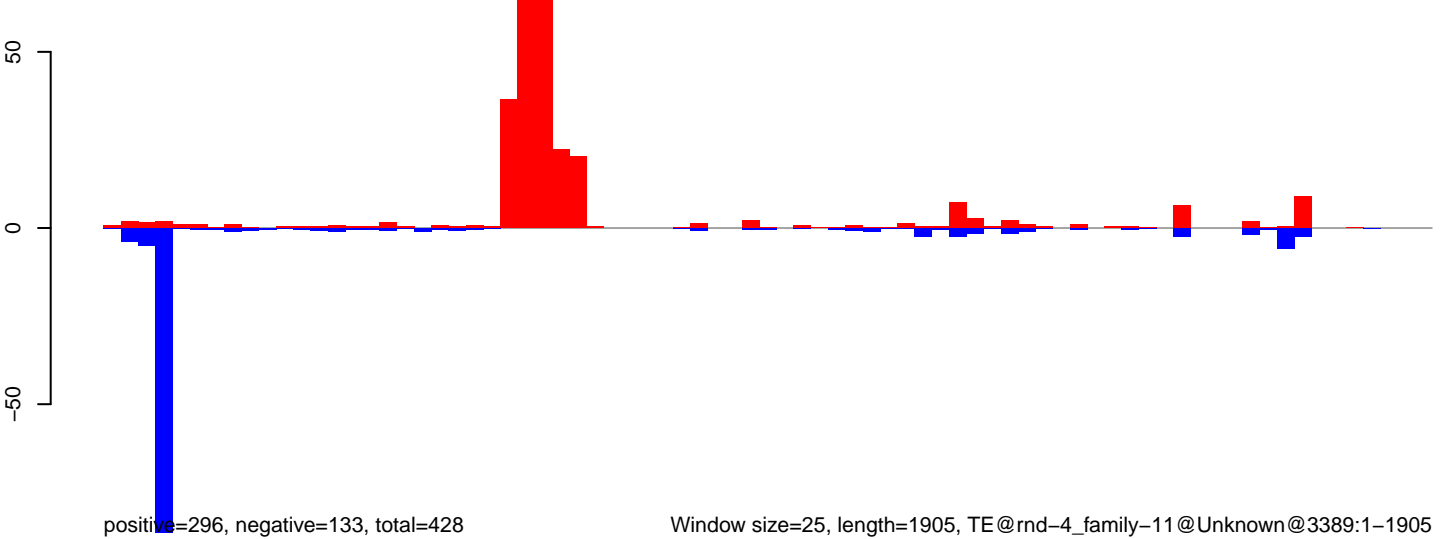
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



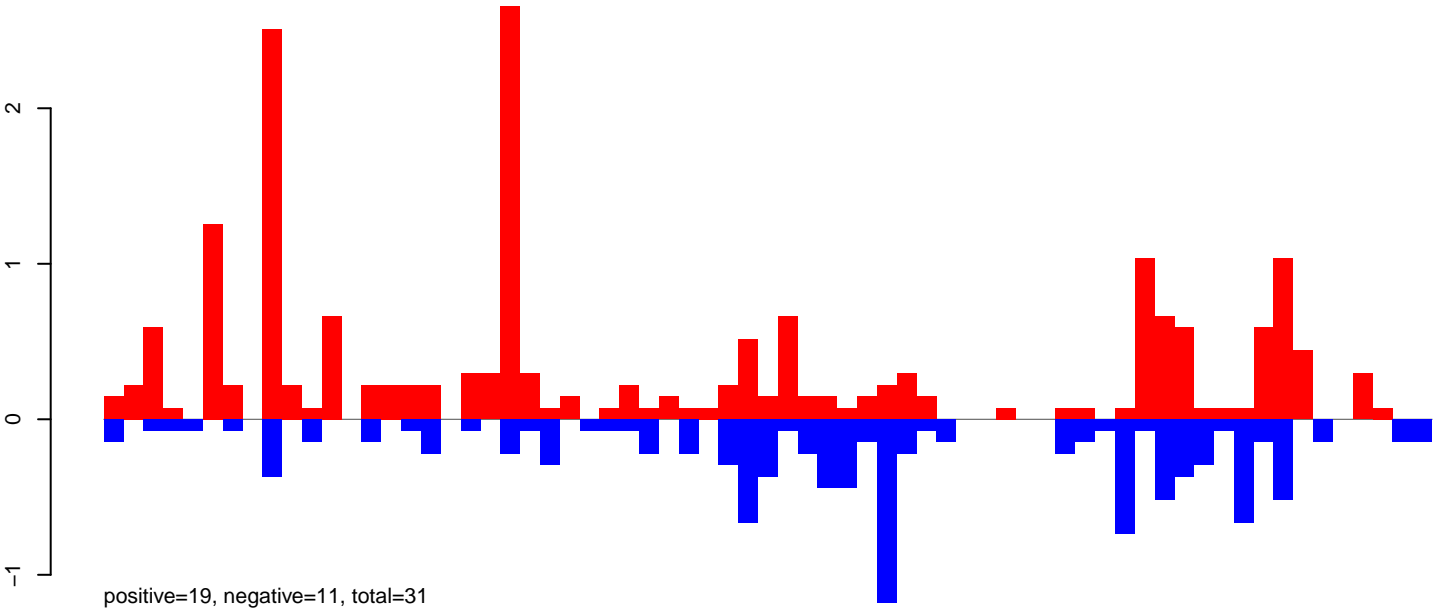
CuQuin_Testes_NL.rep



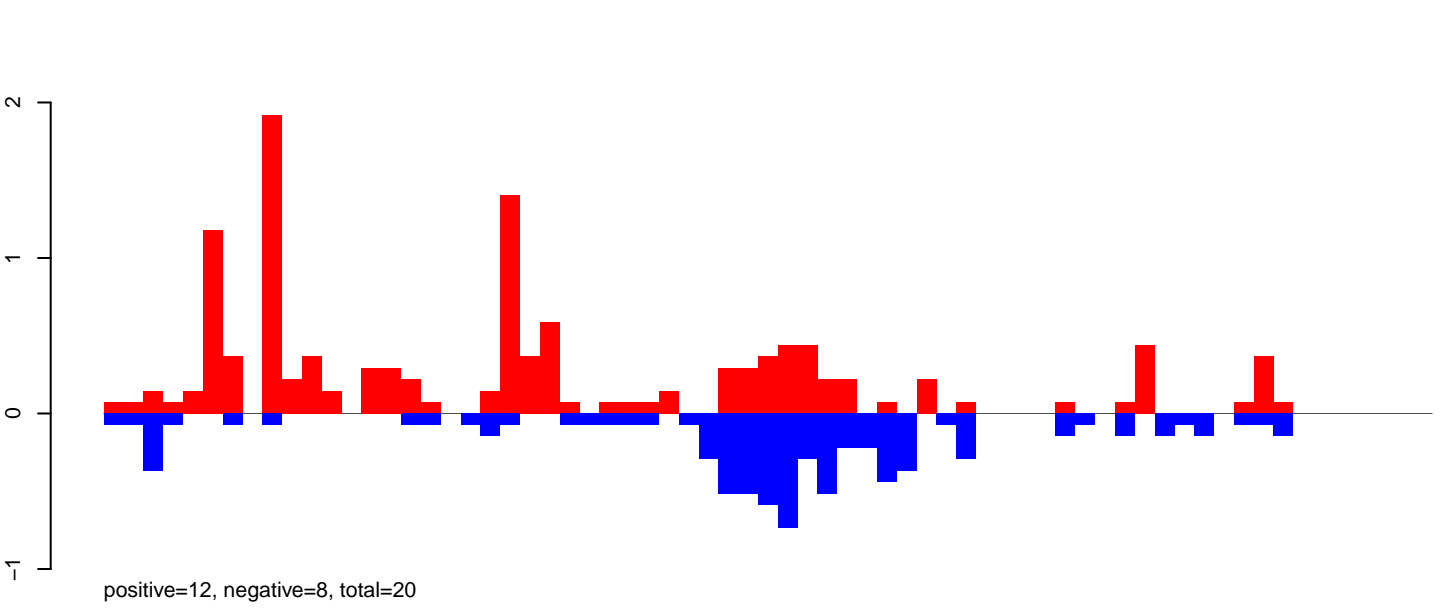
Window size=25, length=1905, TE@rnd-4_family-11@Unknown@3389:1-1905

0 500 1000 1500 2000

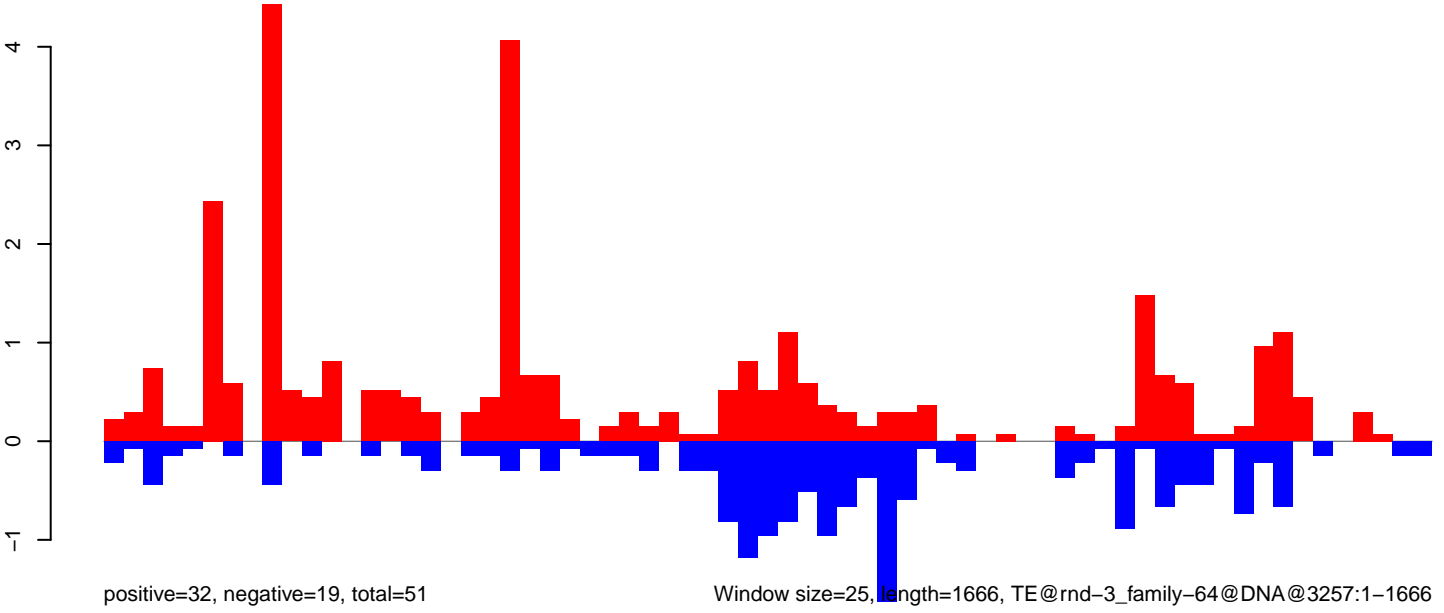
CuQuin_Testes_NL.18_23.rep



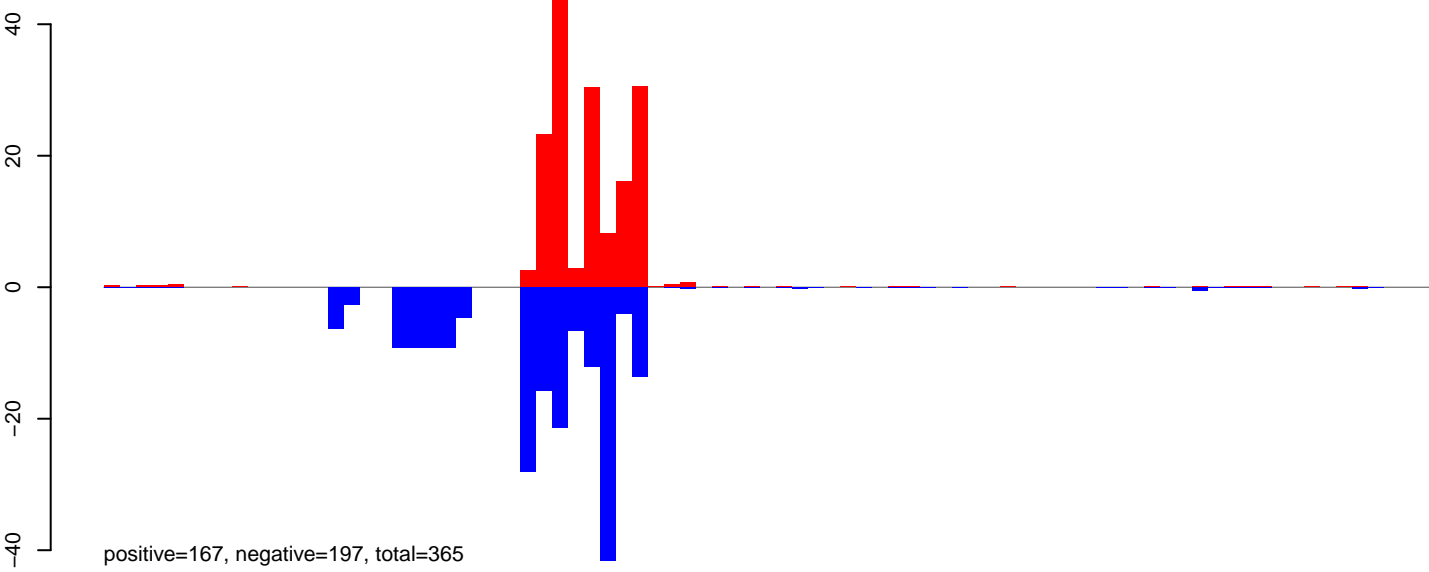
CuQuin_Testes_NL.24_35.rep



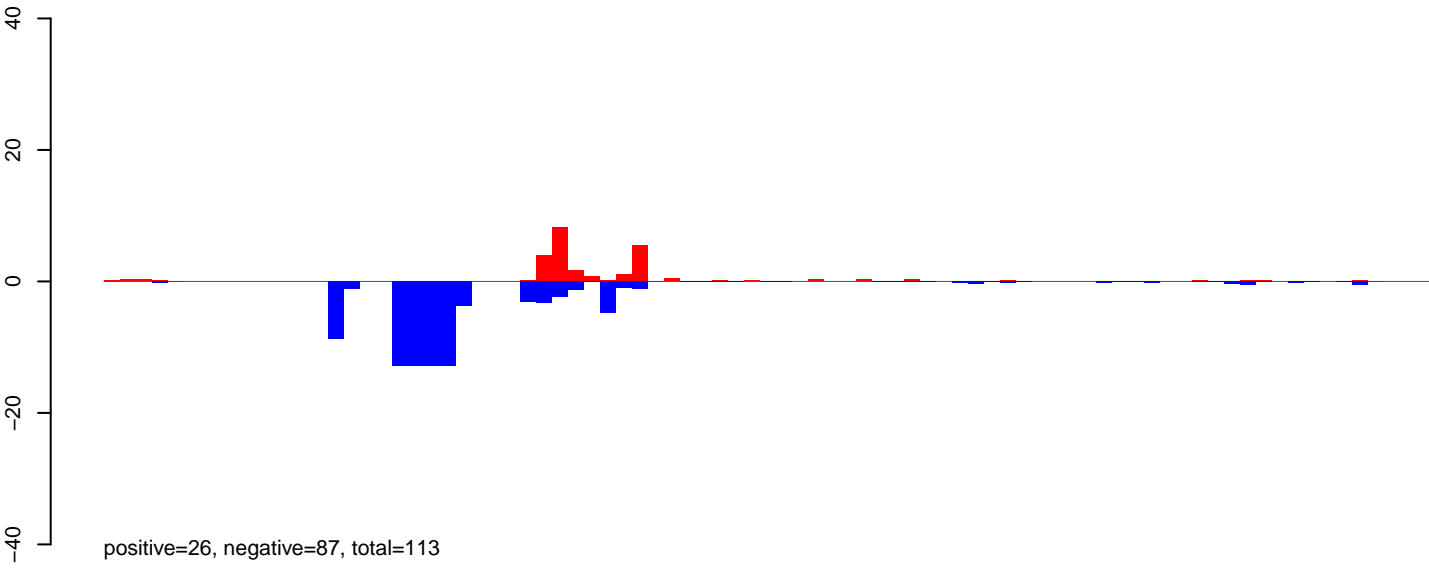
CuQuin_Testes_NL.rep



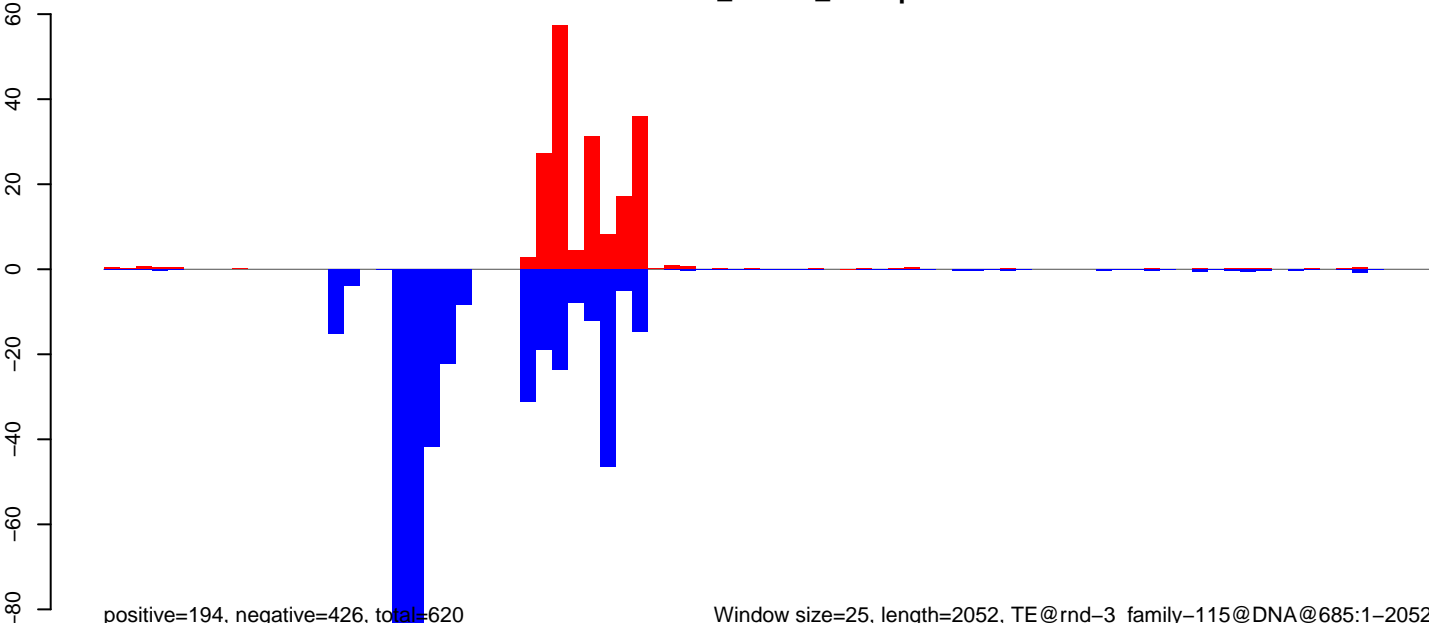
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



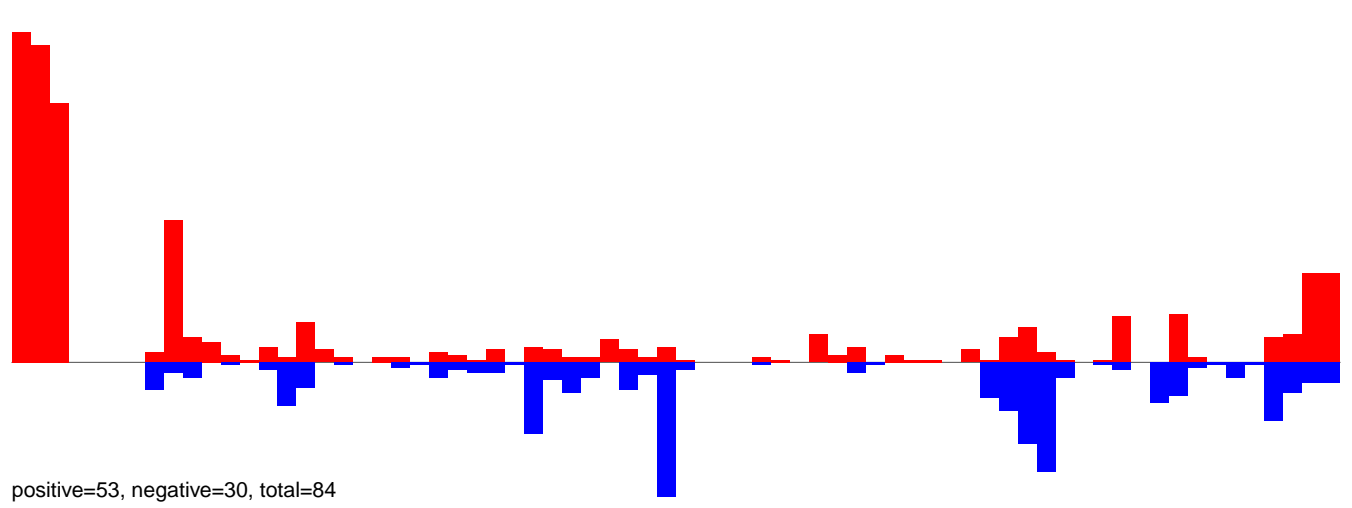
CuQuin_Testes_NL.rep



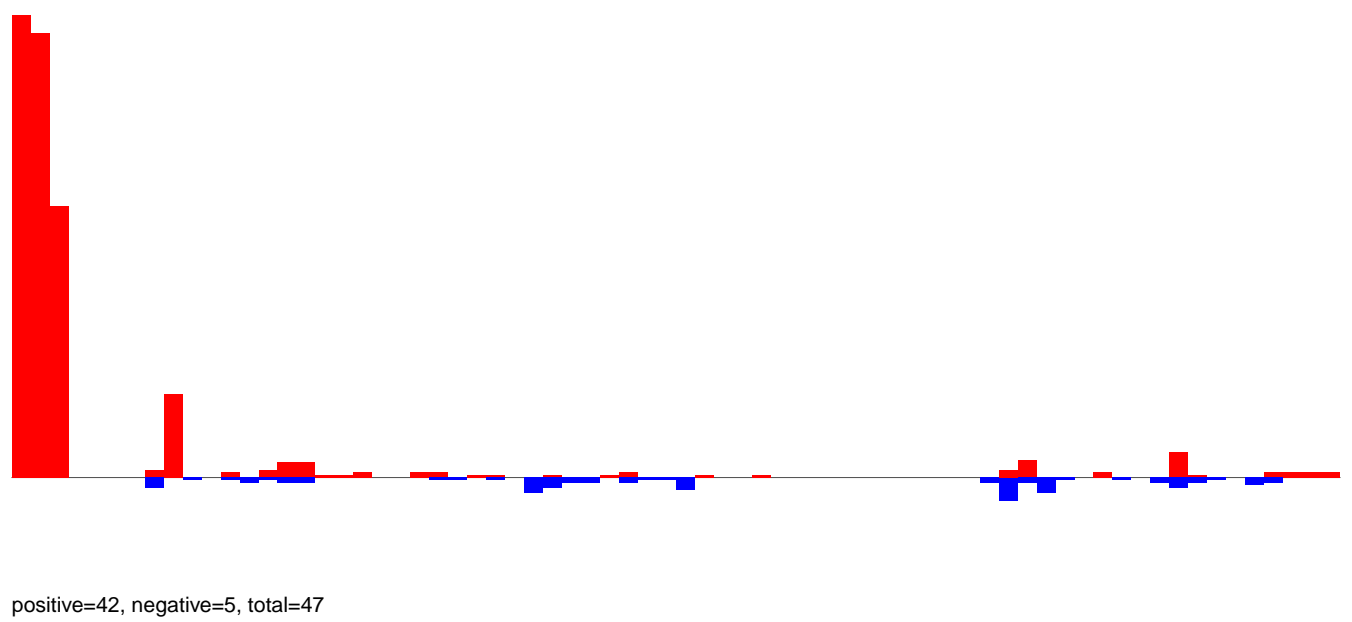
Window size=25, length=2052, TE@rnd-3_family-115@DNA@685:1-2052

0 500 1000 1500 2000

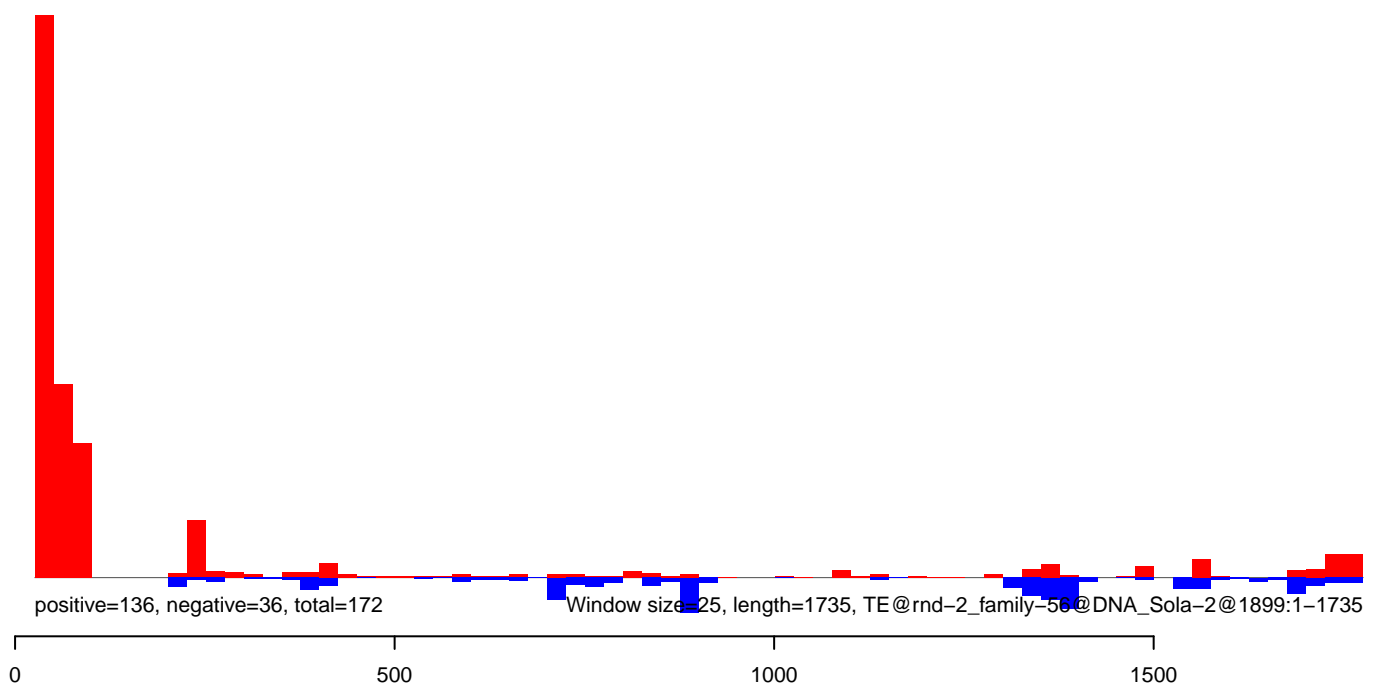
CuQuin_Testes_NL.18_23.rep



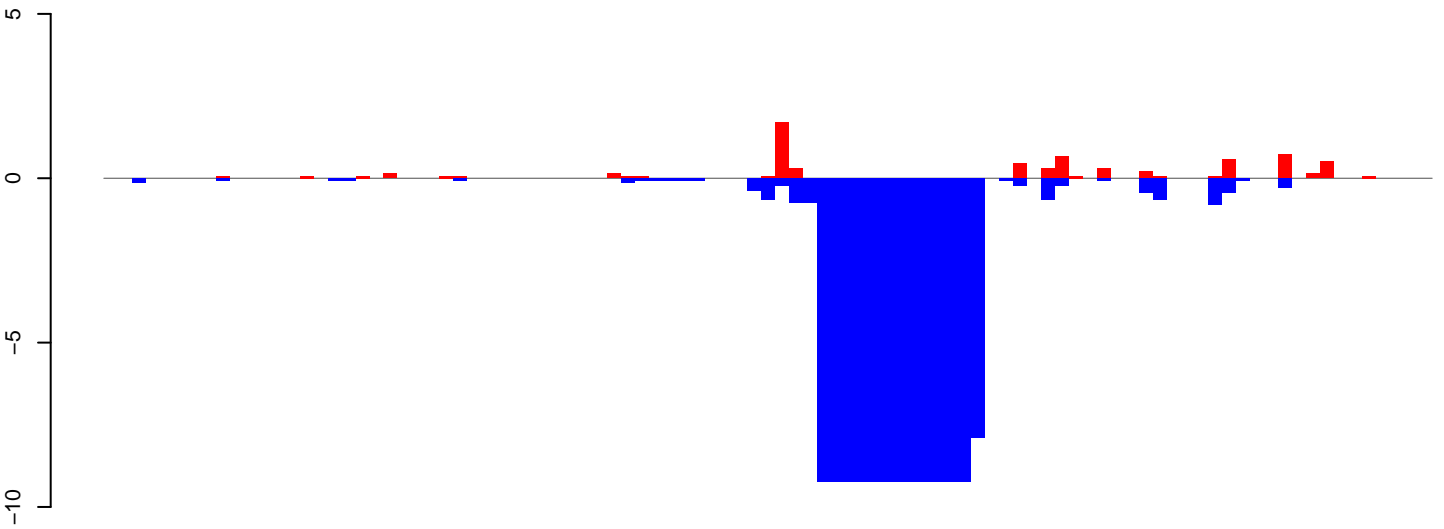
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

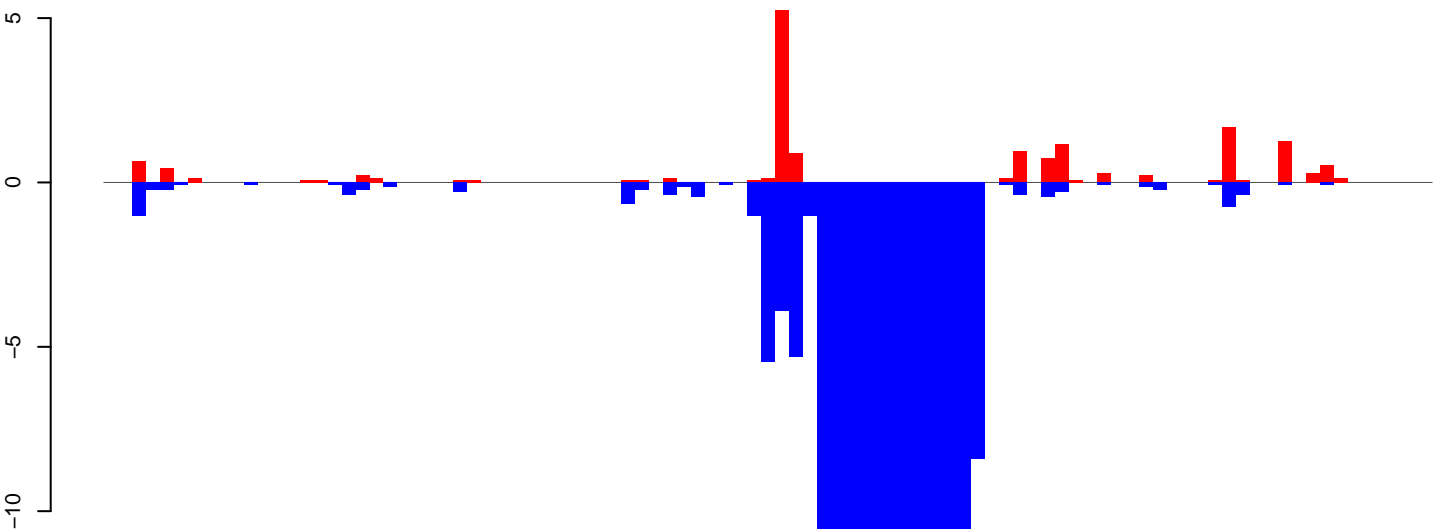


CuQuin_Testes_NL.18_23.rep



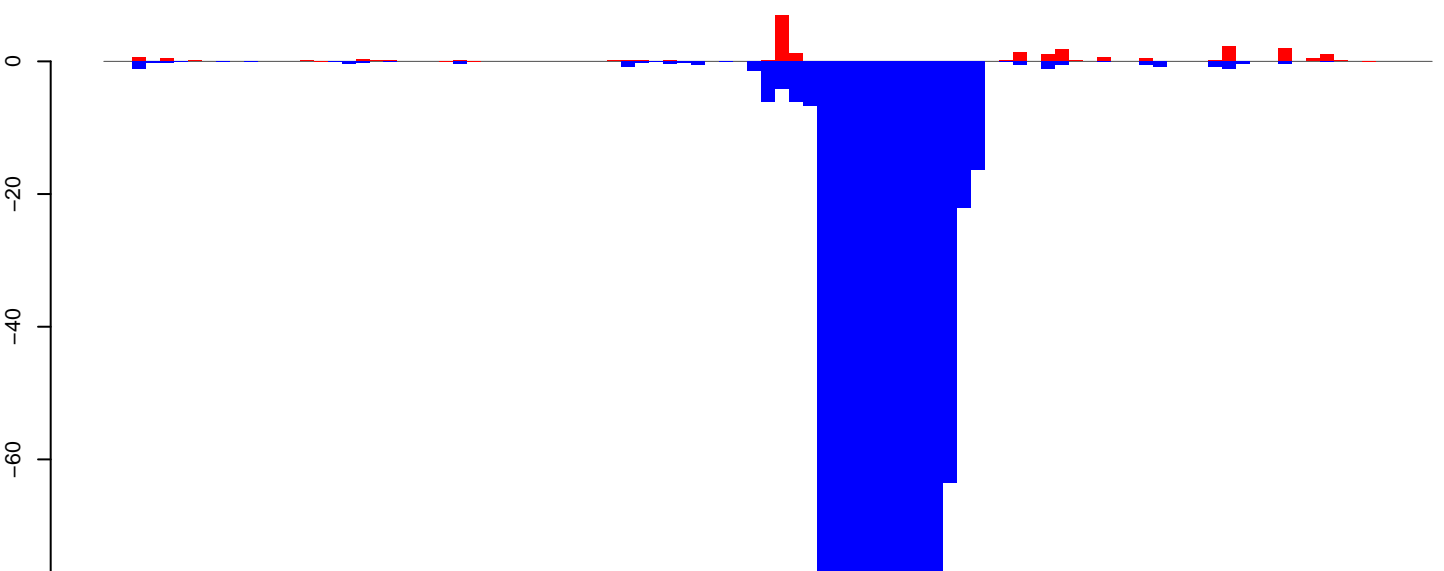
positive=7, negative=117, total=124

CuQuin_Testes_NL.24_35.rep



positive=16, negative=175, total=191

CuQuin_Testes_NL.rep

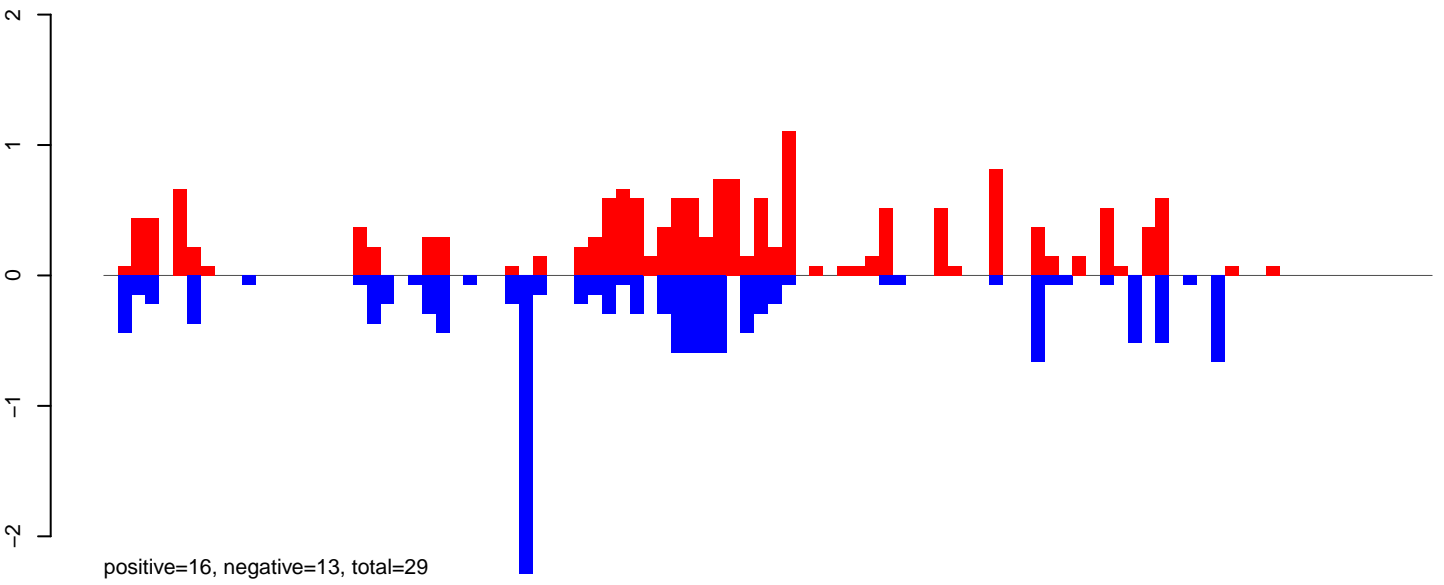


positive=23, negative=887, total=910

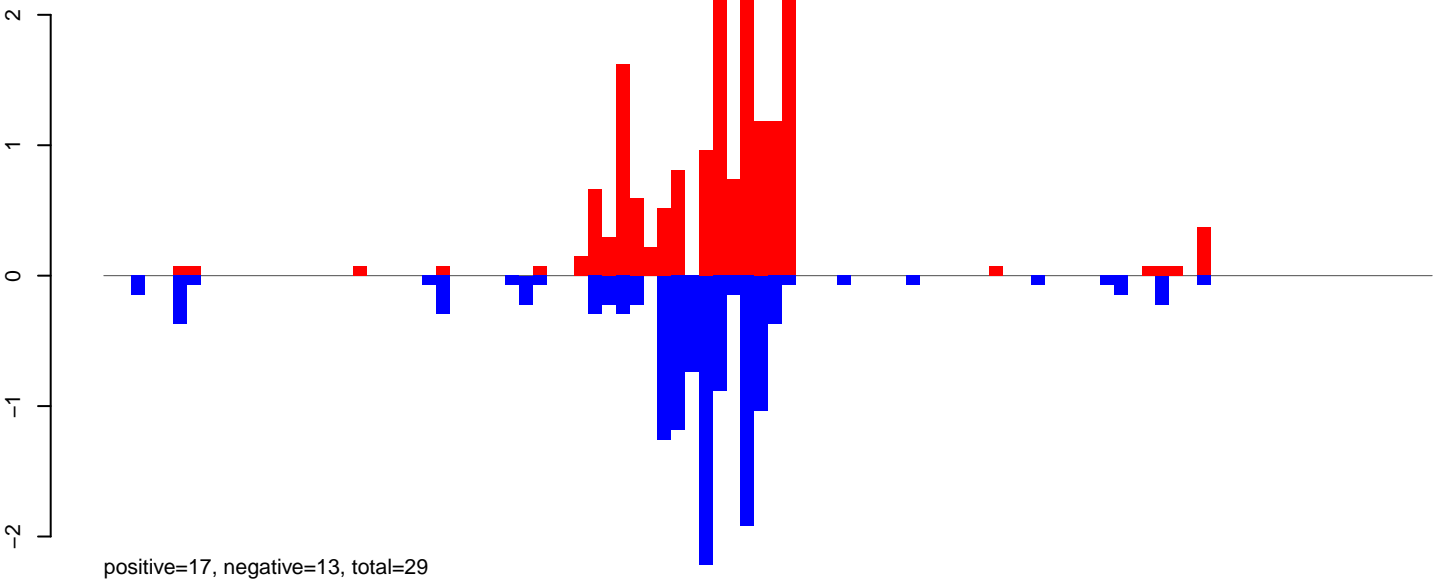
Window size=25, length=2371, TE@rnd-5_family-898@Unknown@2326:1-2371

0 500 1000 1500 2000

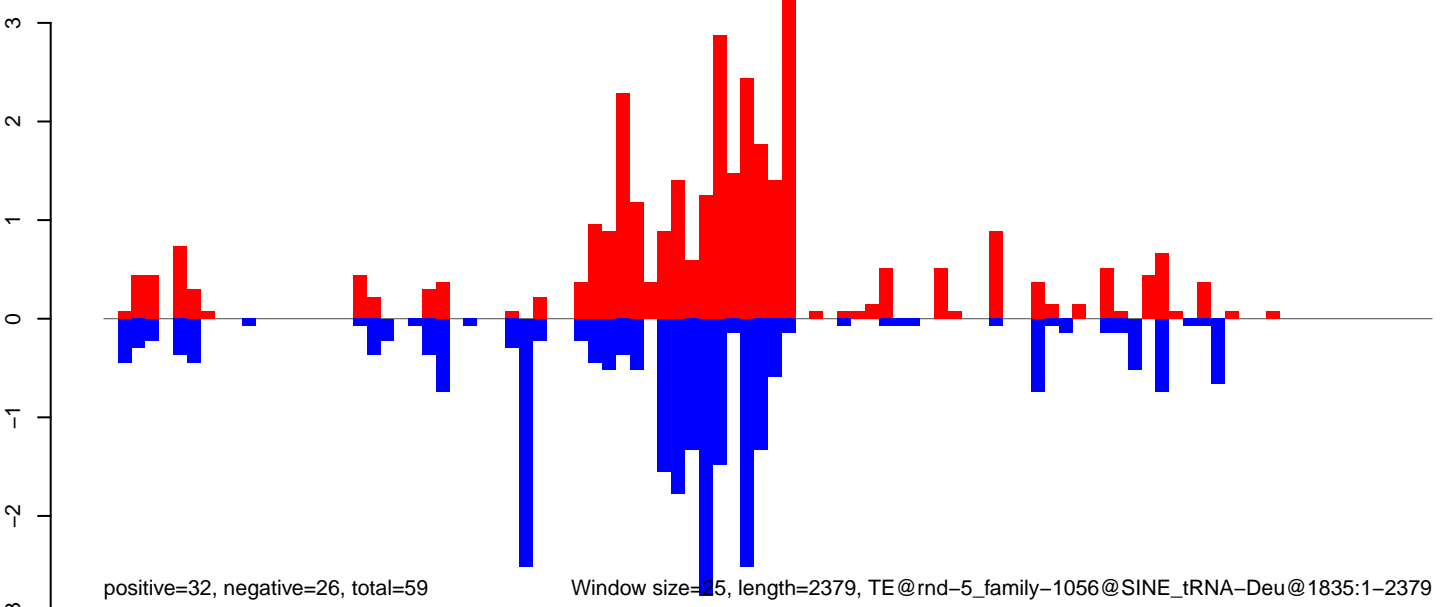
CuQuin_Testes_NL.18_23.rep



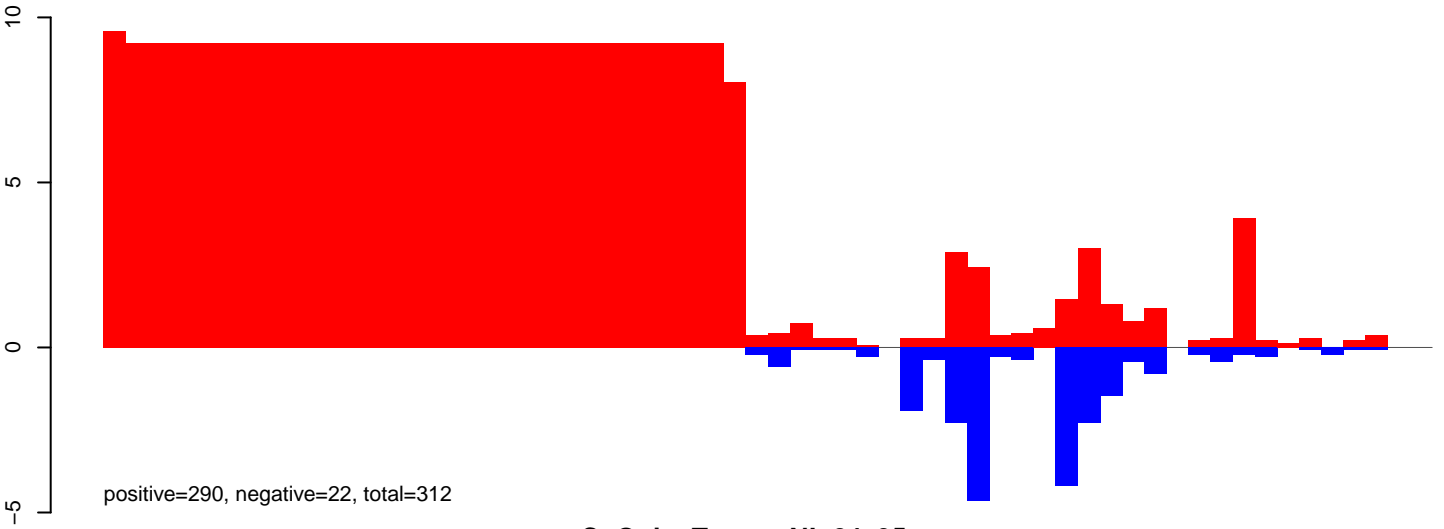
CuQuin_Testes_NL.24_35.rep



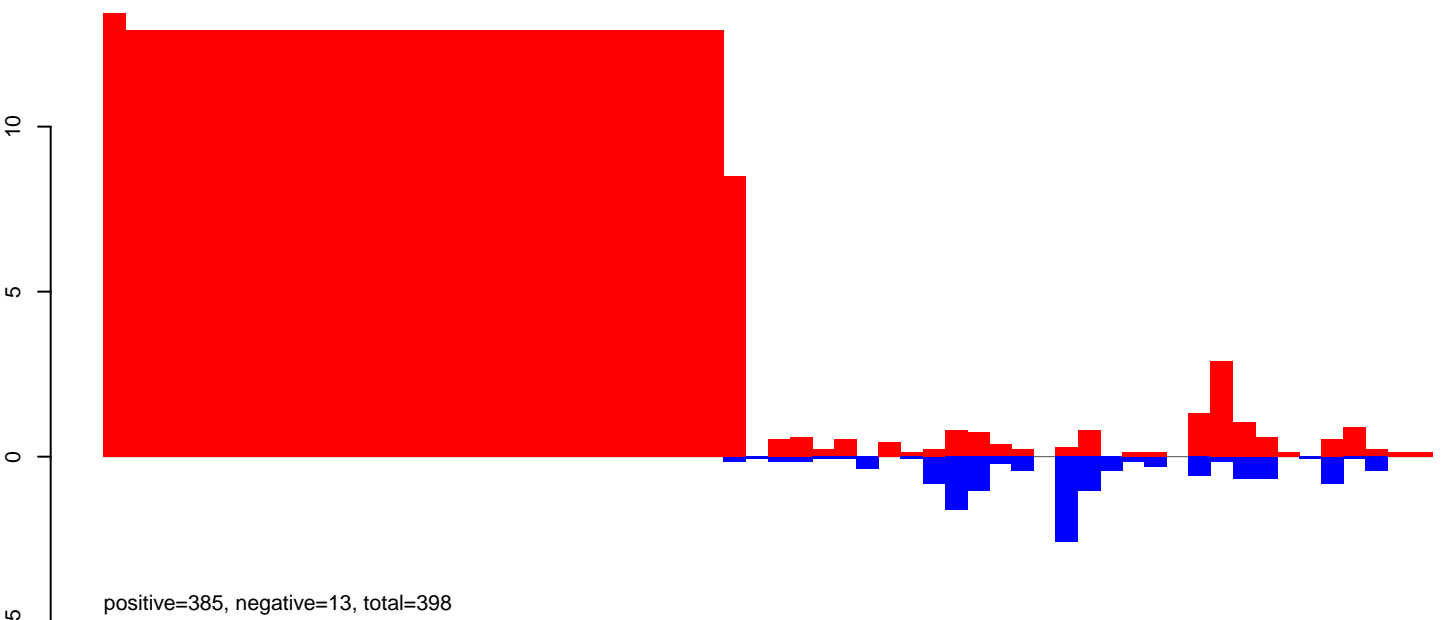
CuQuin_Testes_NL.rep



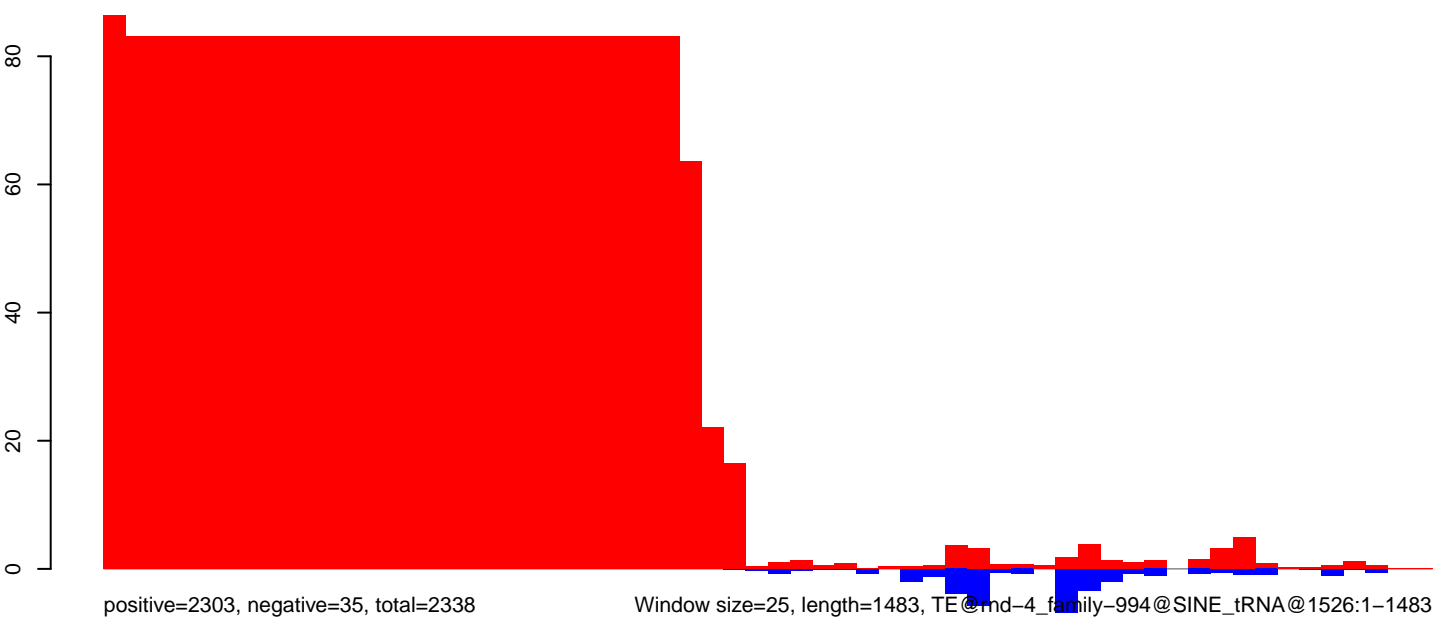
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



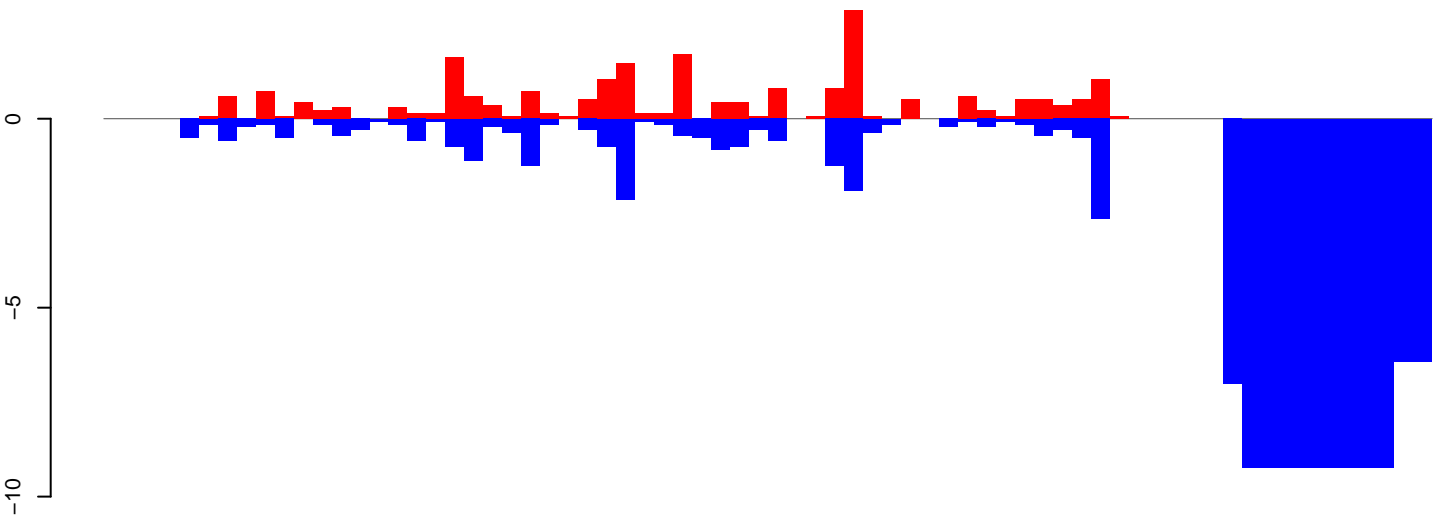
CuQuin_Testes_NL.rep



Window size=25, length=1483, TE@md-4_family-994@SINE_tRNA@1526:1-1483

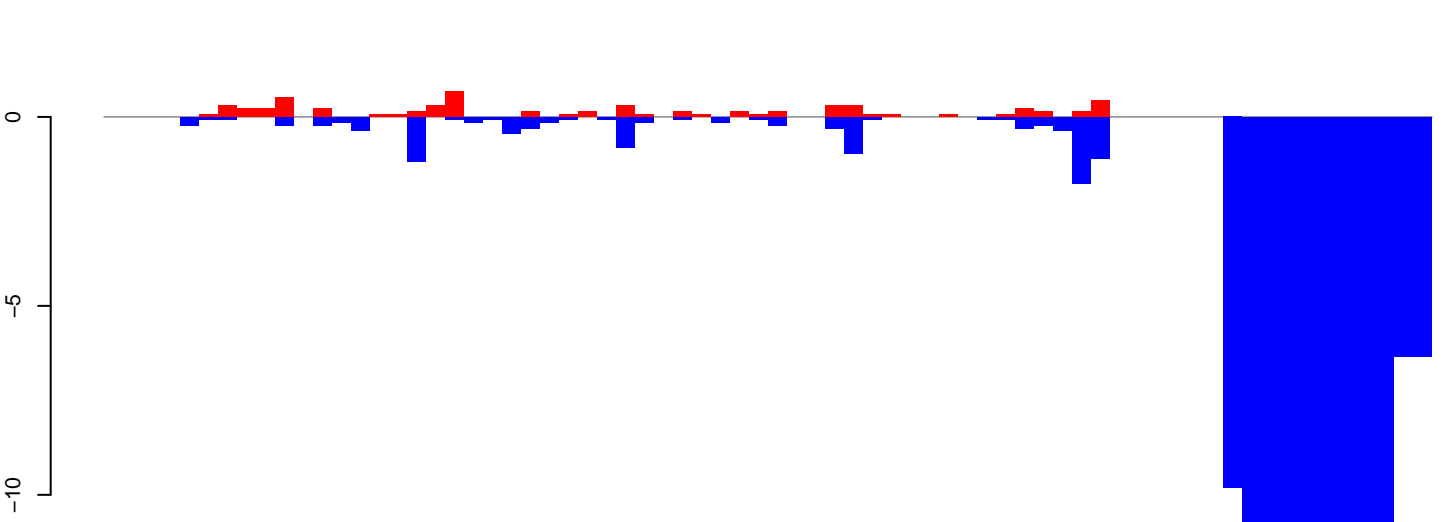
0 500 1000 1500

CuQuin_Testes_NL.18_23.rep



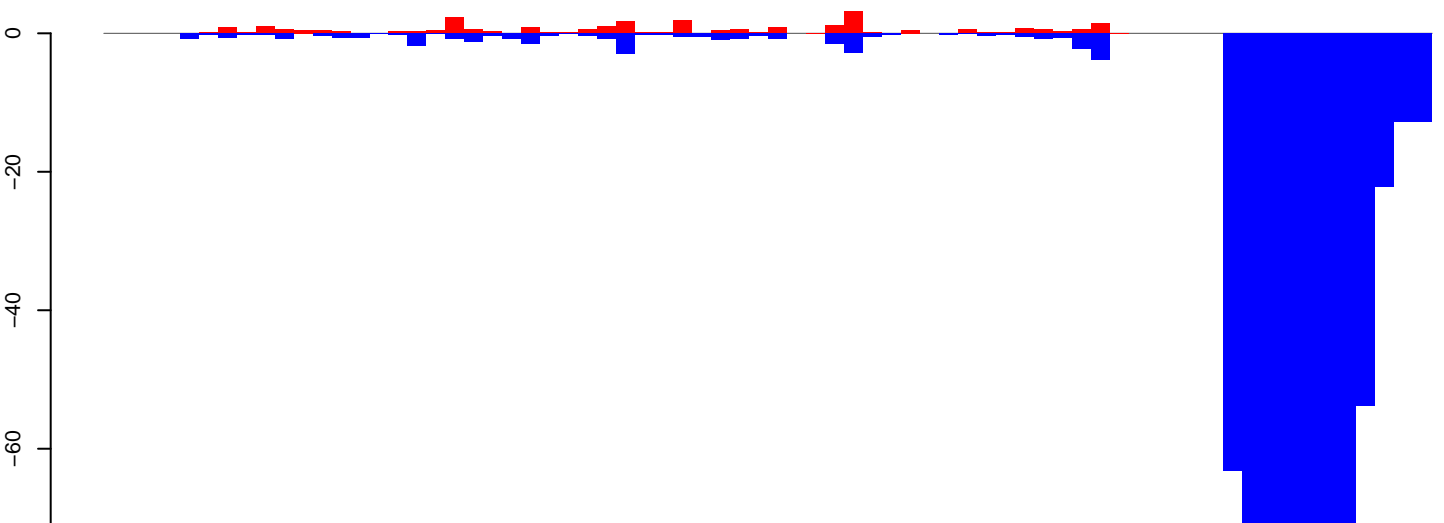
positive=22, negative=117, total=138

CuQuin_Testes_NL.24_35.rep



positive=6, negative=136, total=142

CuQuin_Testes_NL.rep

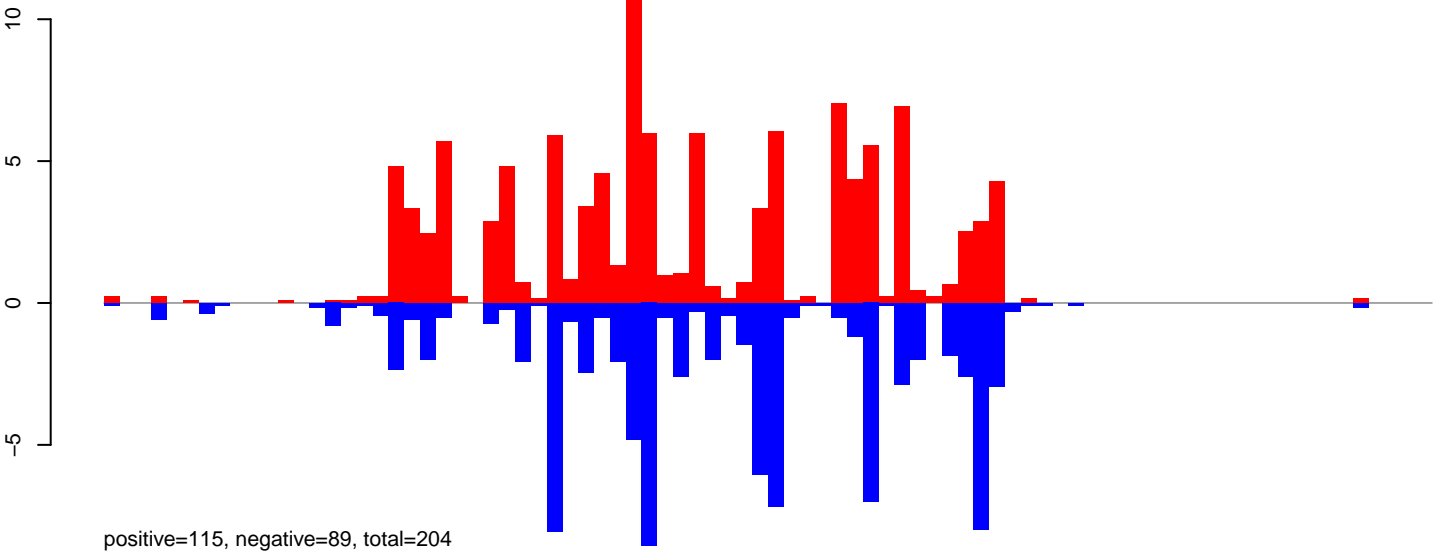


positive=28, negative=697, total=724

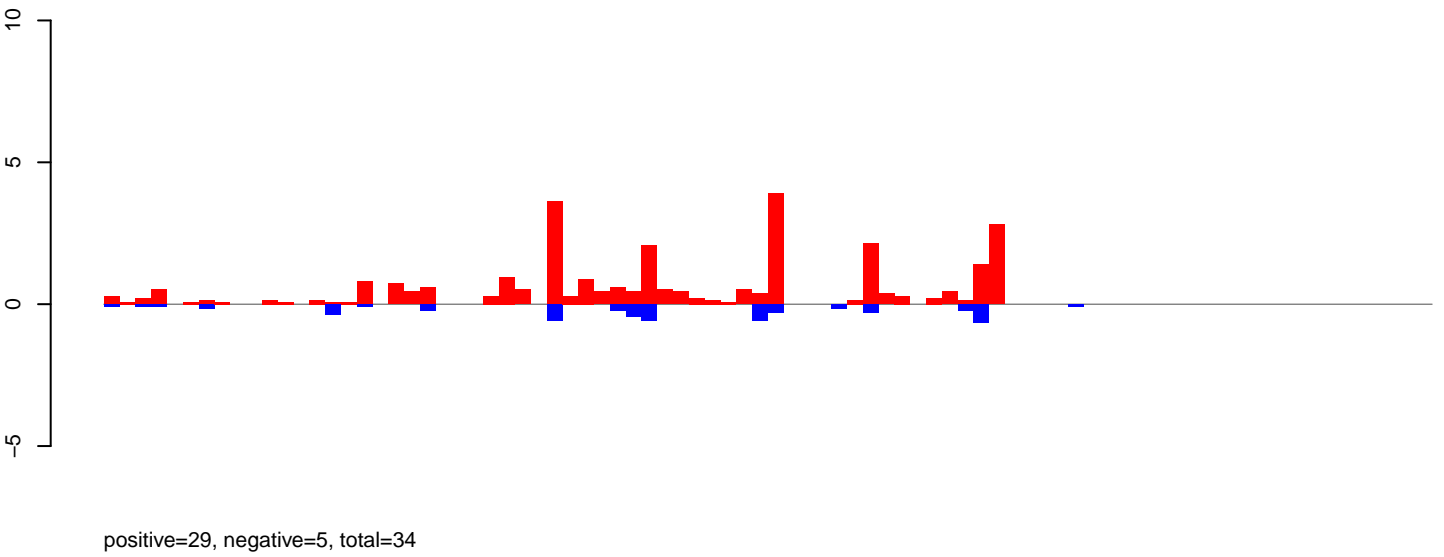
Window size=25, length=1739, TE@rnd-3_family-105@Unknown@1283:1-1739

0 500 1000 1500

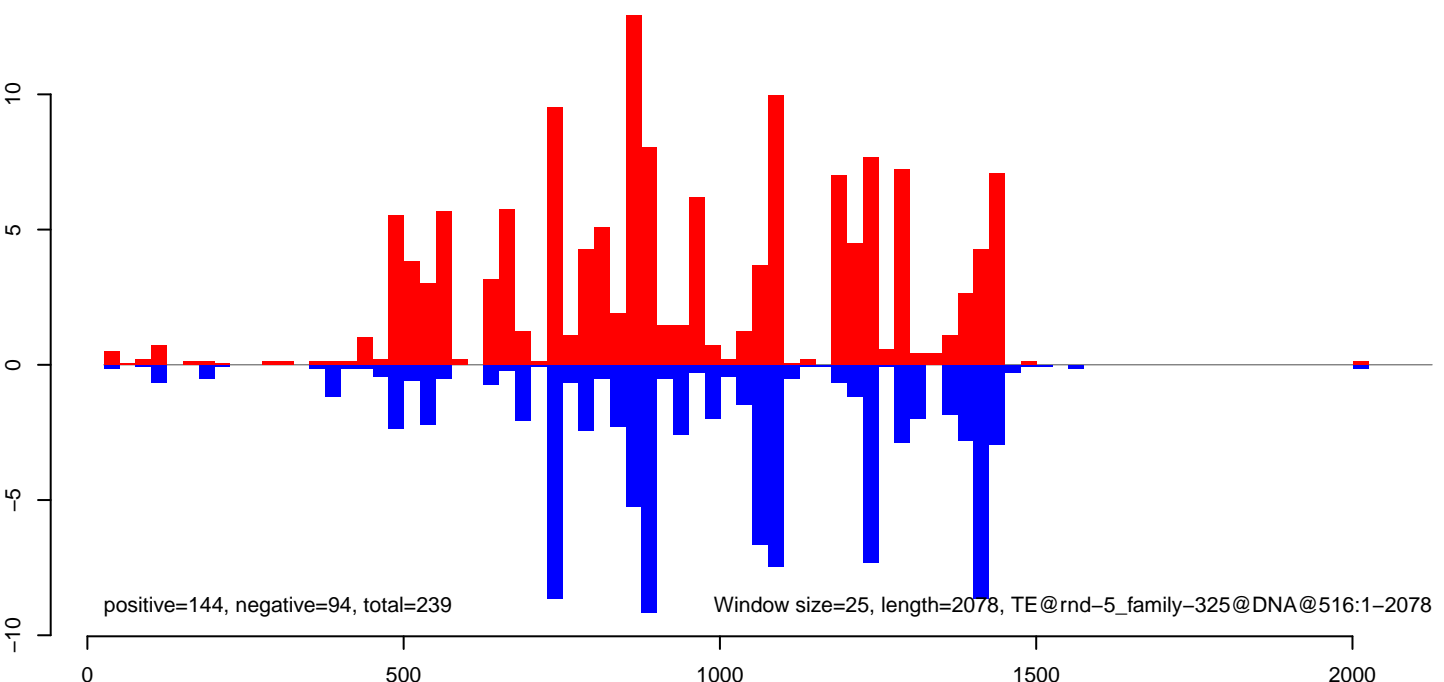
CuQuin_Testes_NL.18_23.rep



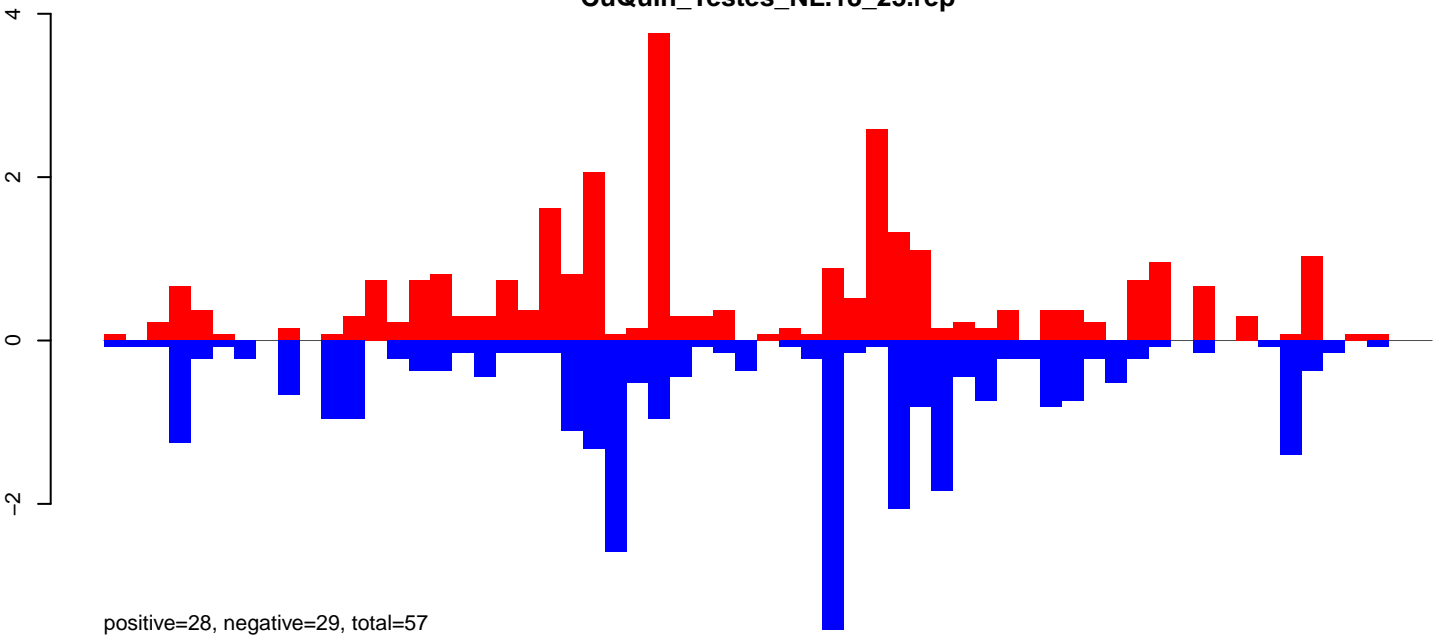
CuQuin_Testes_NL.24_35.rep



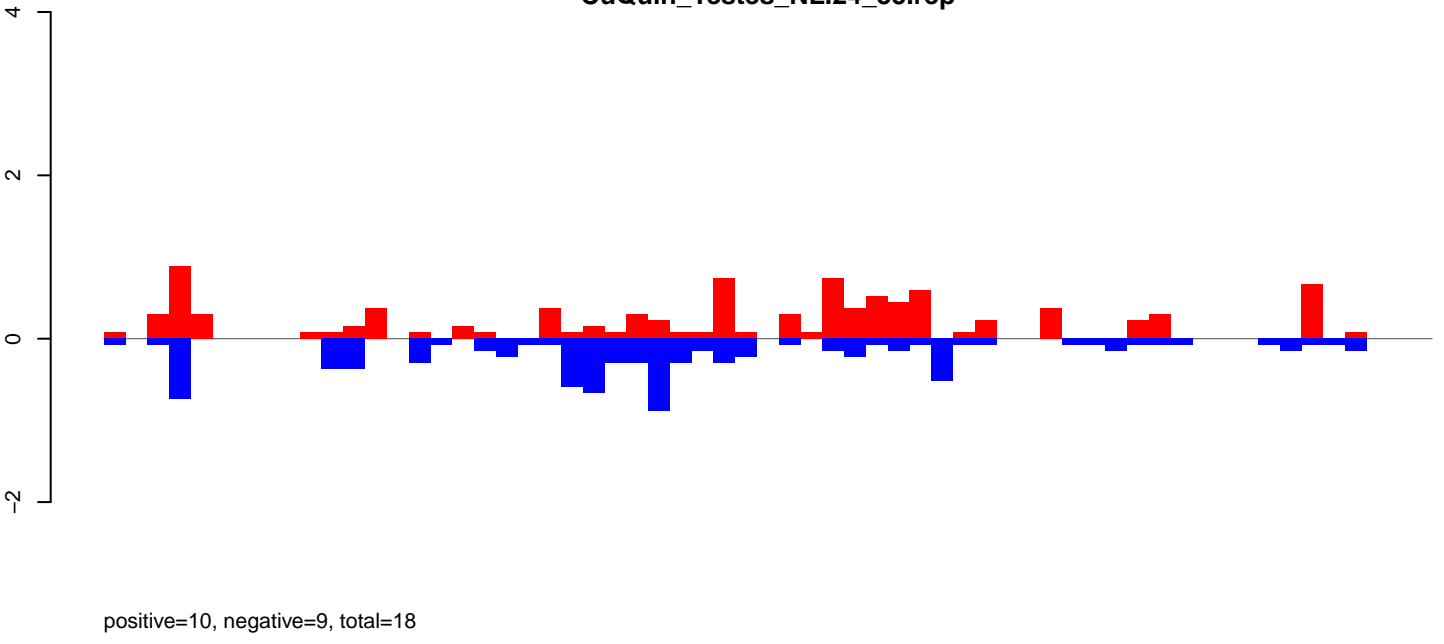
CuQuin_Testes_NL.rep



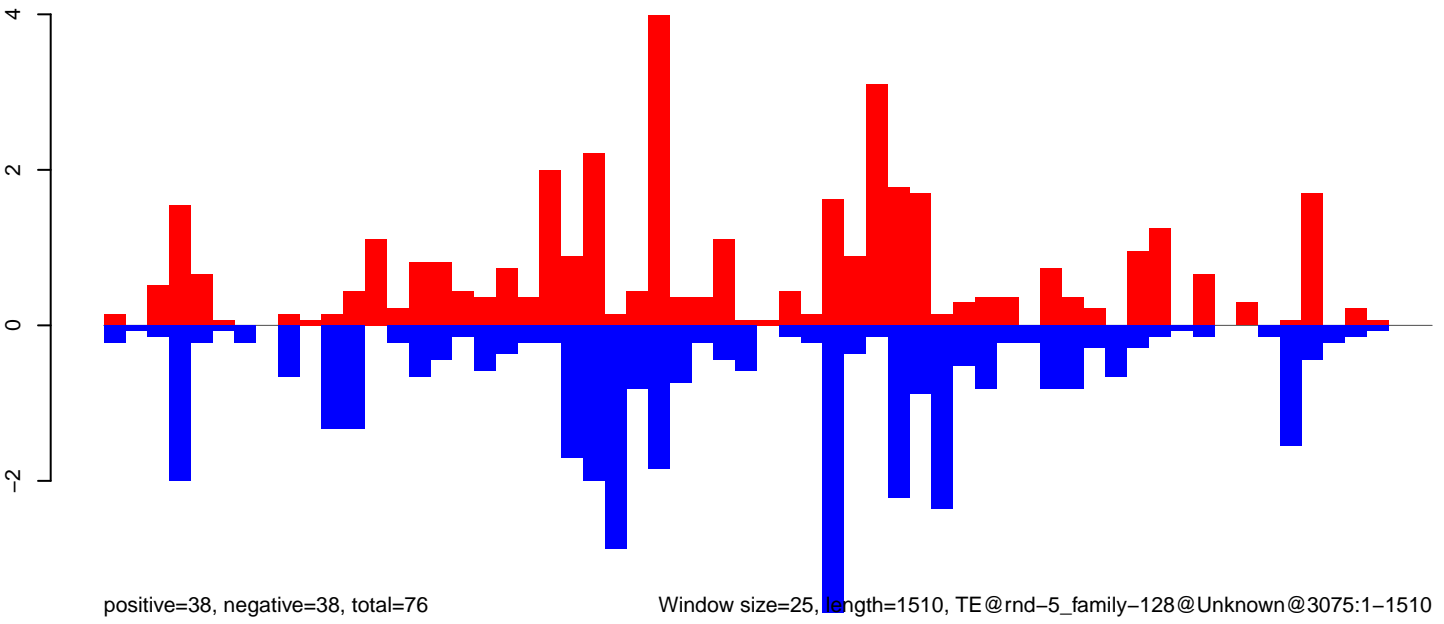
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

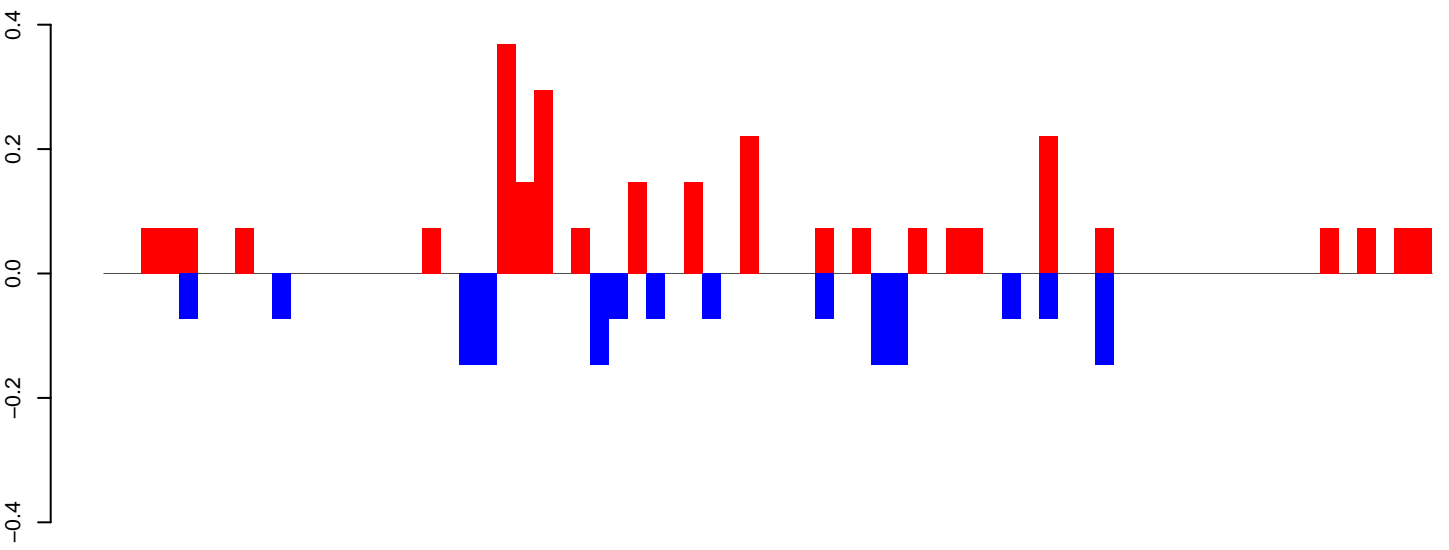


CuQuin_Testes_NL.rep



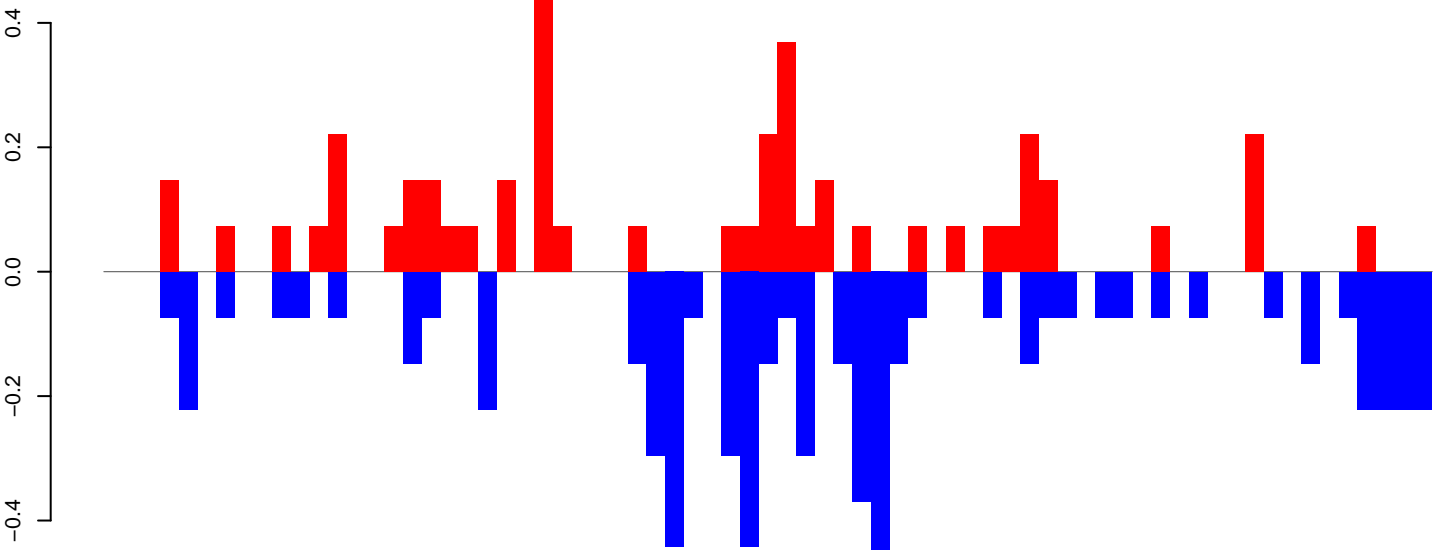
Window size=25, length=1510, TE@rnd-5_family-128@Unknown@3075:1-1510

CuQuin_Testes_NL.18_23.rep



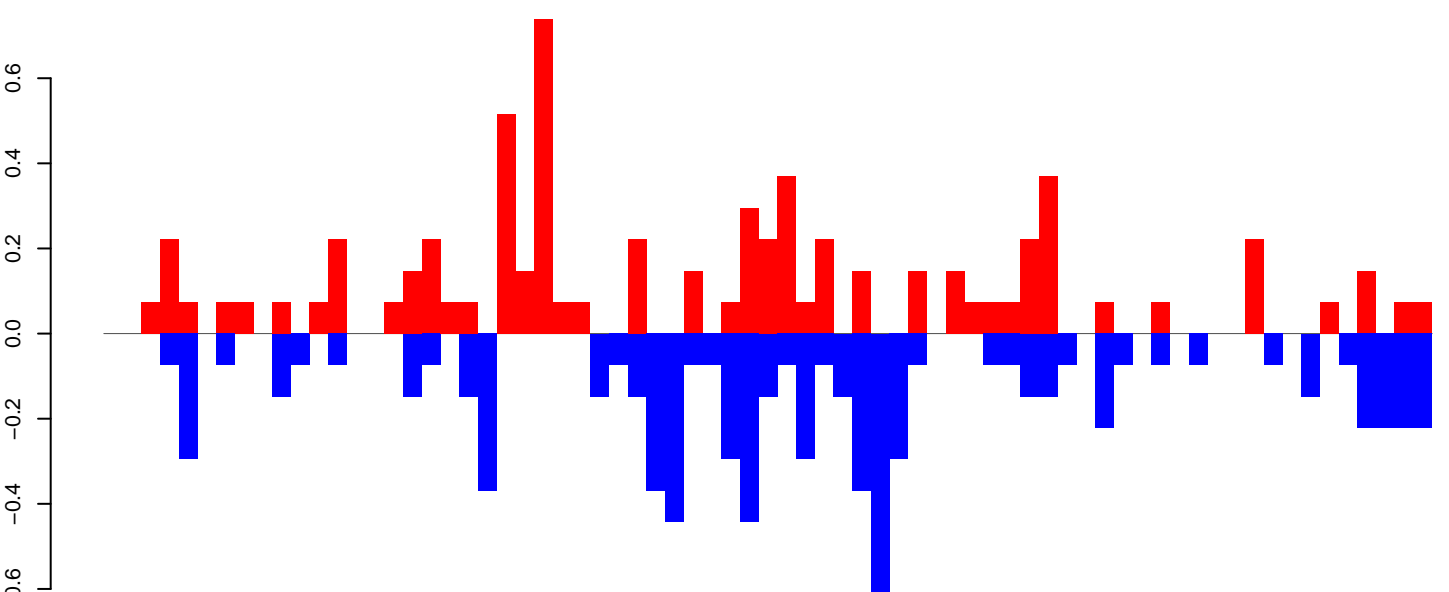
positive=3, negative=1, total=4

CuQuin_Testes_NL.24_35.rep



positive=4, negative=6, total=10

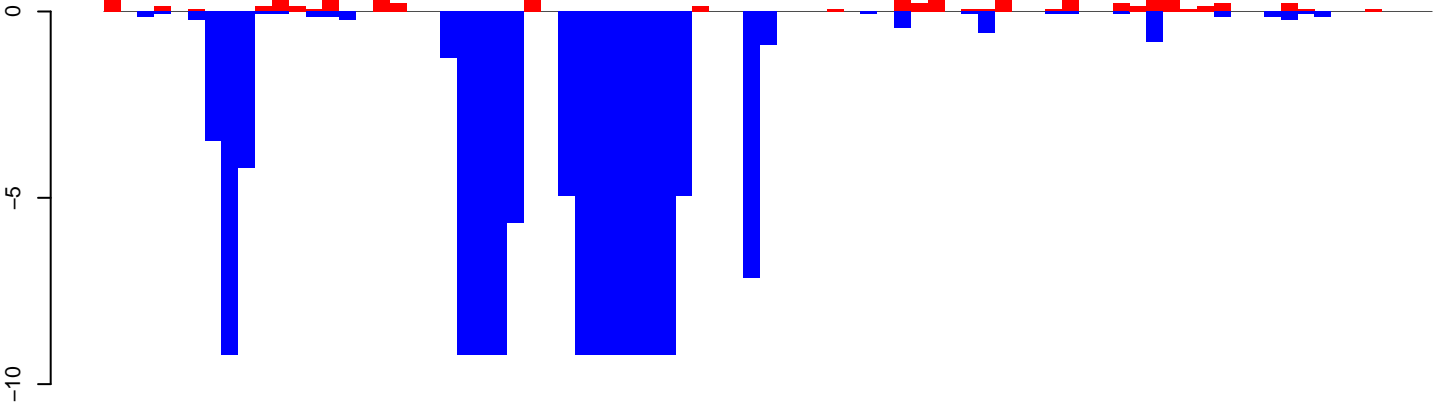
CuQuin_Testes_NL.rep



positive=7, negative=8, total=14

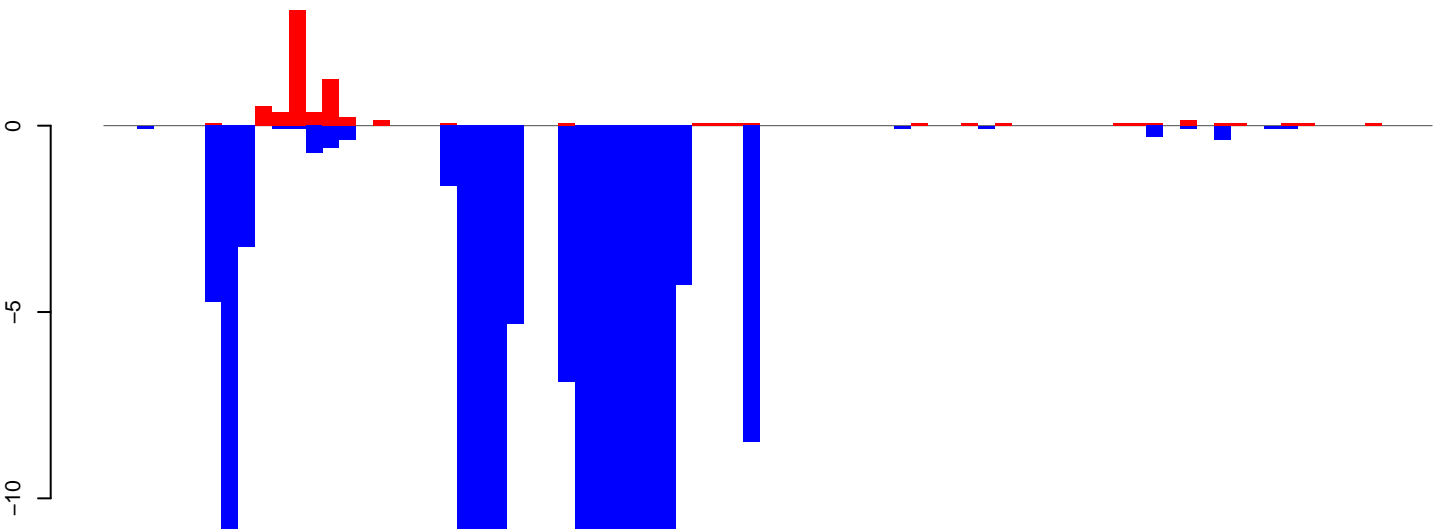
Window size=25, length=1775, TE@rnd-4_family-574@Unknown@1445:1-1775

CuQuin_Testes_NL.18_23.rep



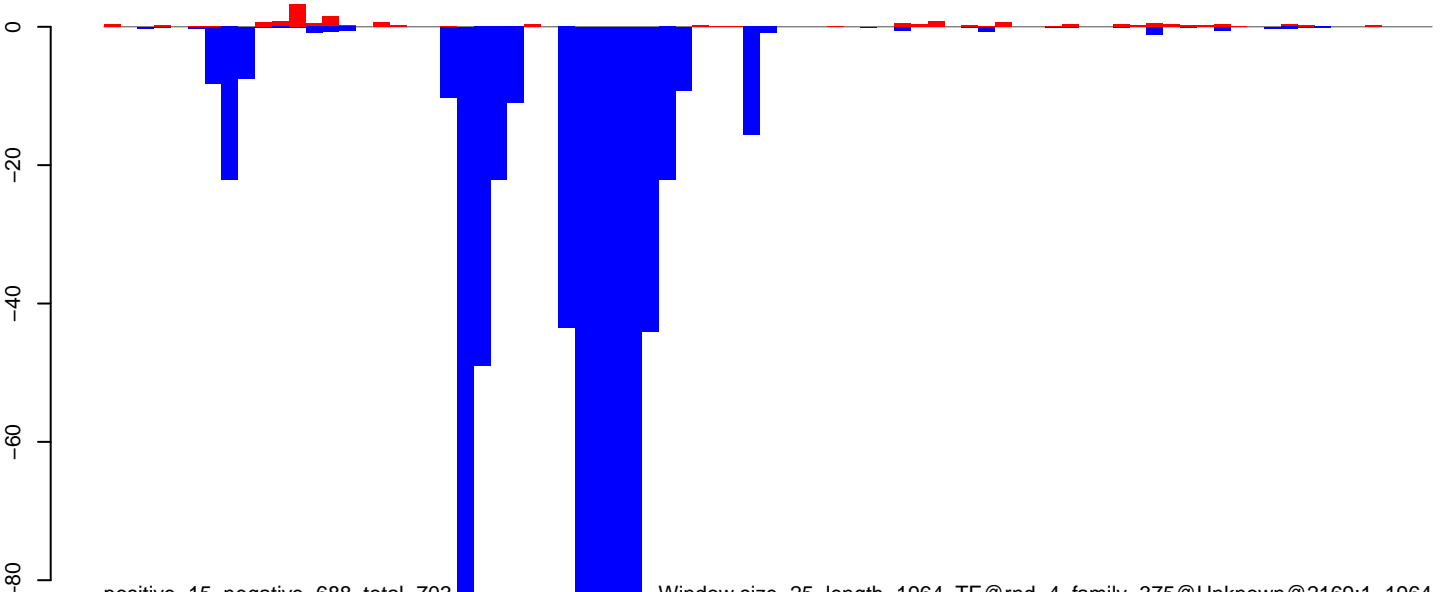
positive=7, negative=129, total=136

CuQuin_Testes_NL.24_35.rep



positive=7, negative=167, total=174

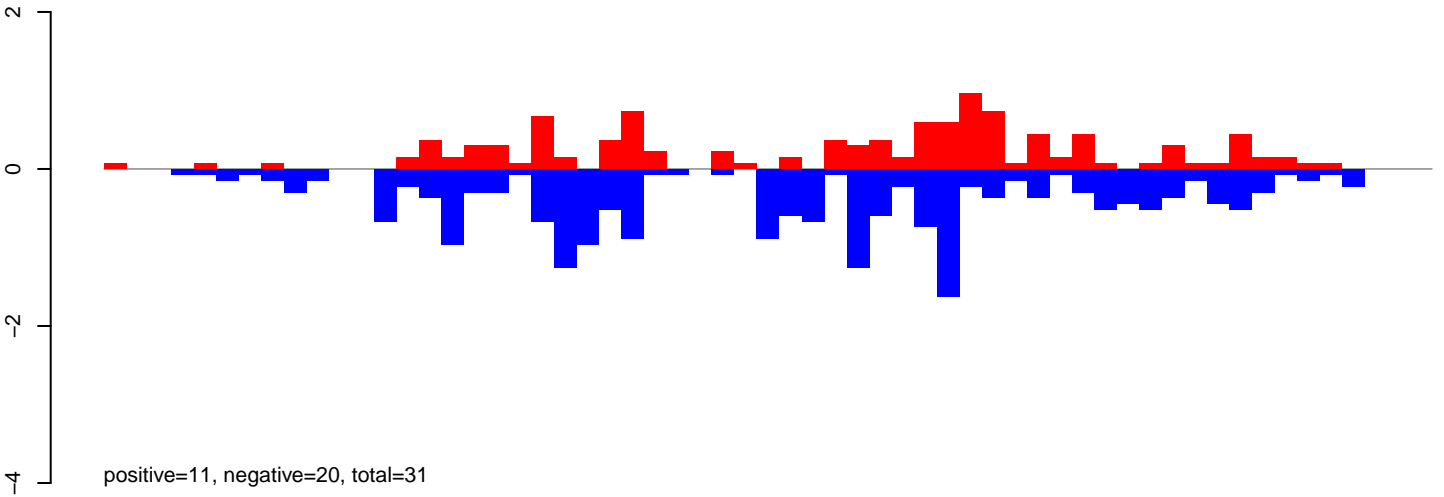
CuQuin_Testes_NL.rep



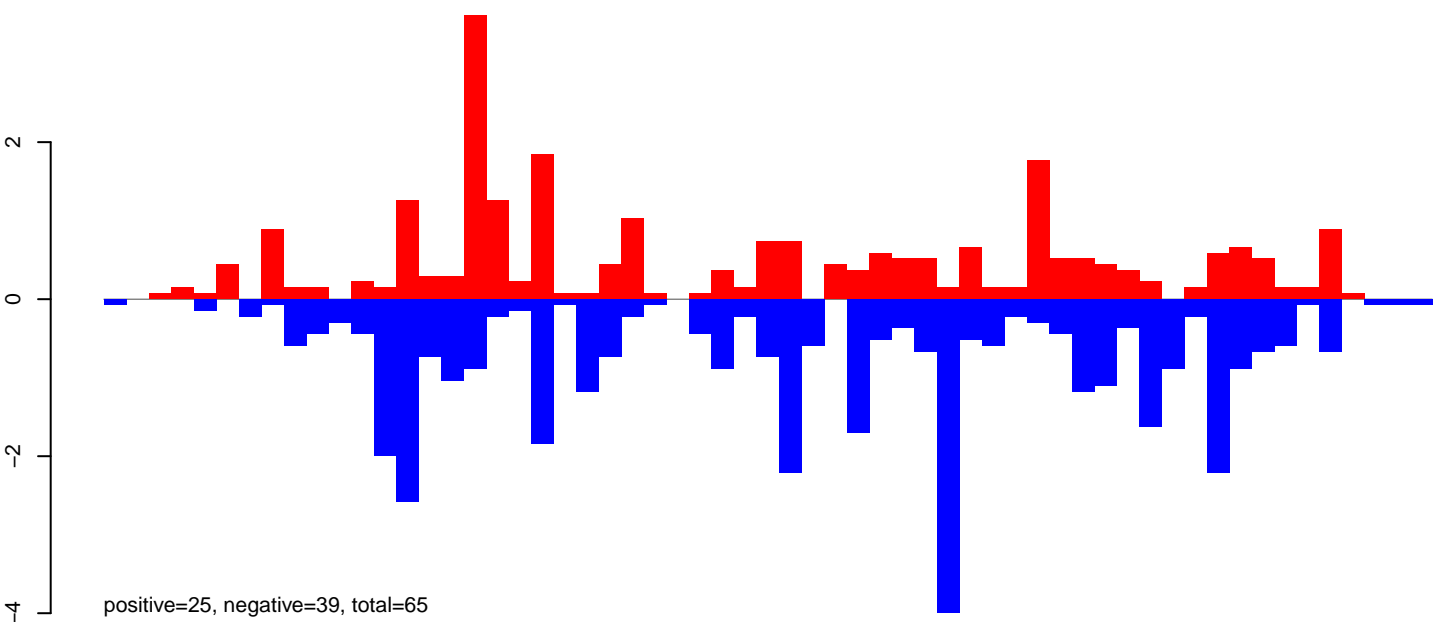
positive=15, negative=688, total=703

Window size=25, length=1964, TE@rnd-4_family-375@Unknown@2169:1-1964

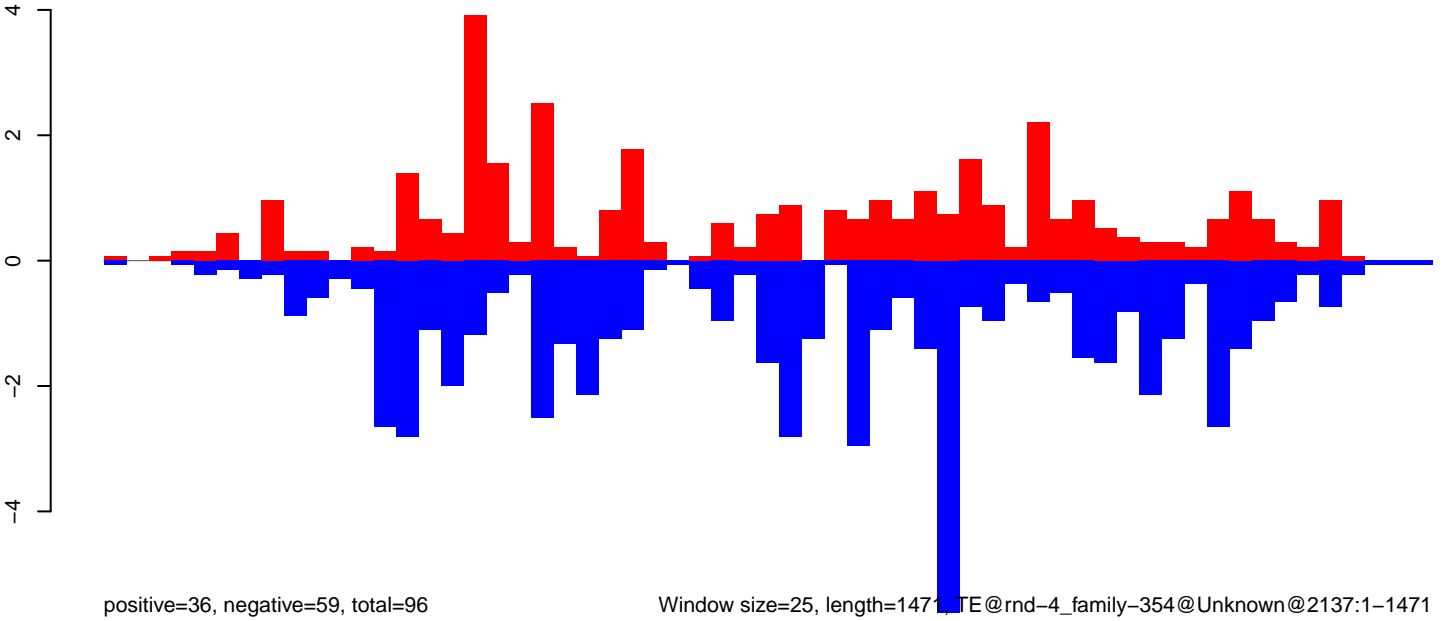
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

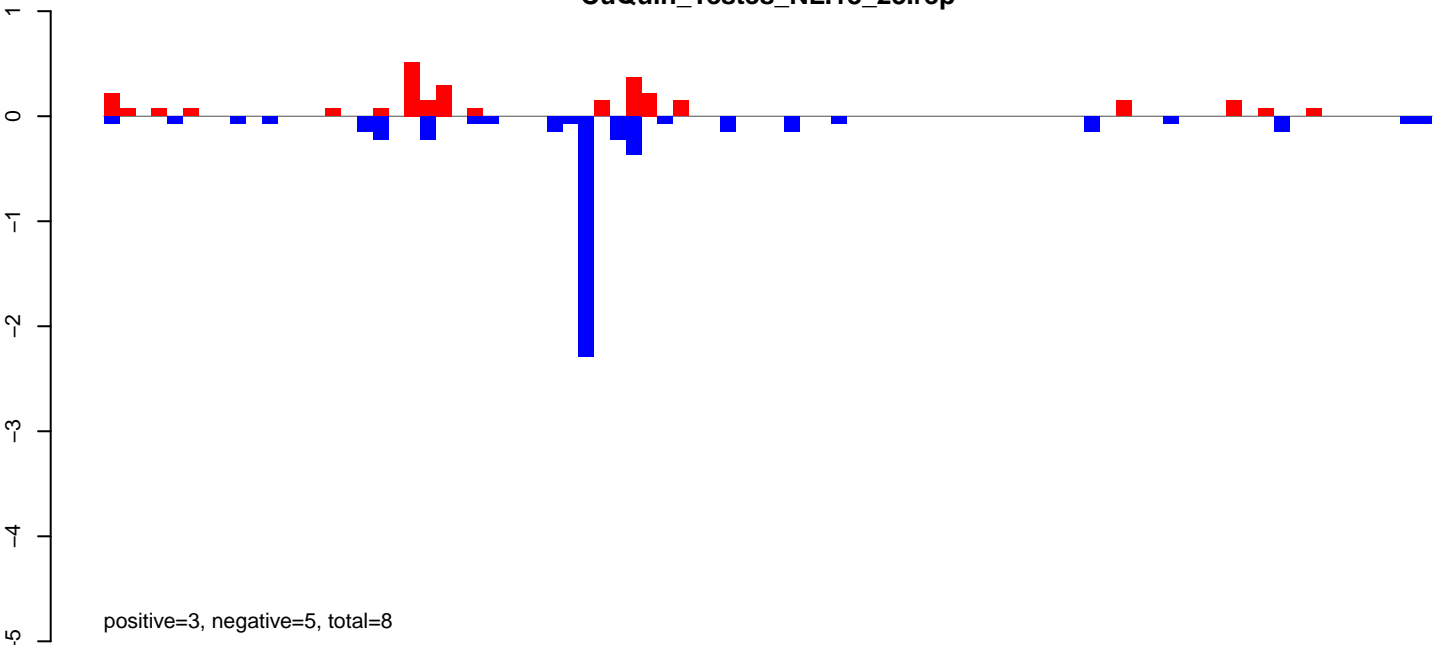


CuQuin_Testes_NL.rep

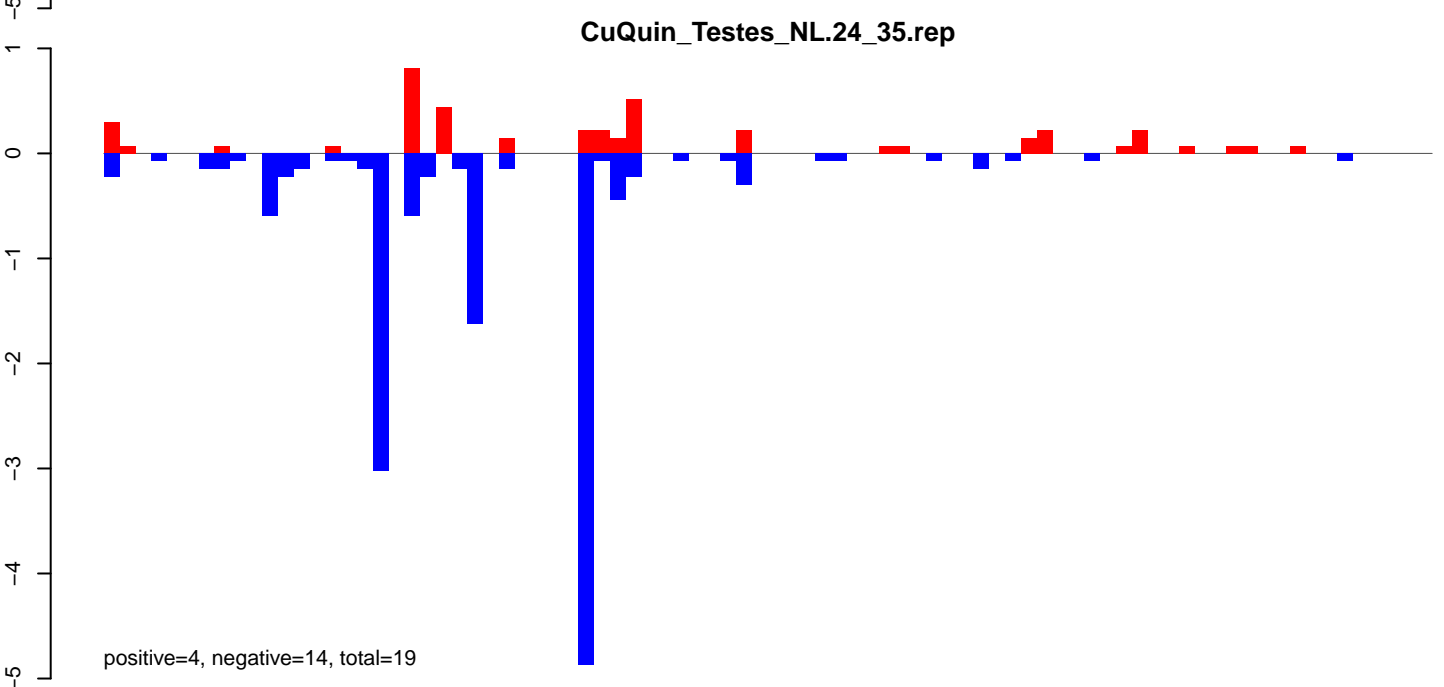


Window size=25, length=1471 TE@rnd-4_family-354@Unknown@2137:1-1471

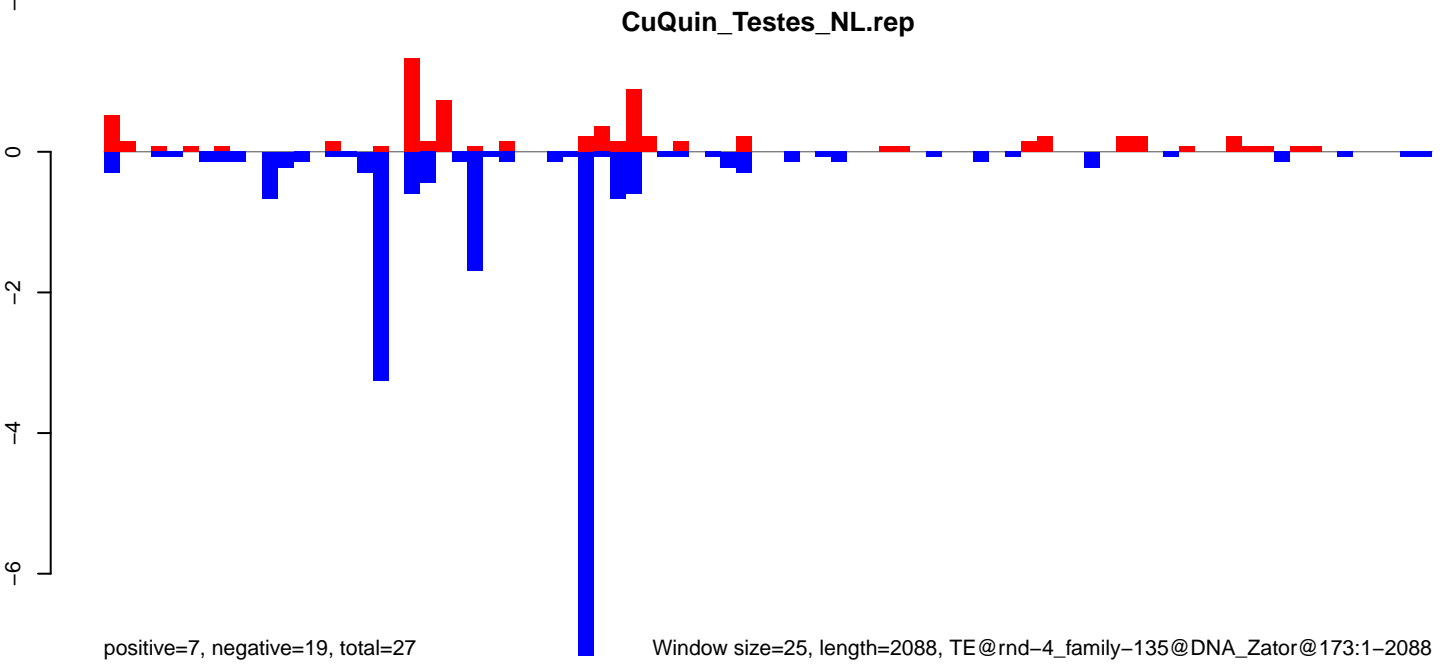
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

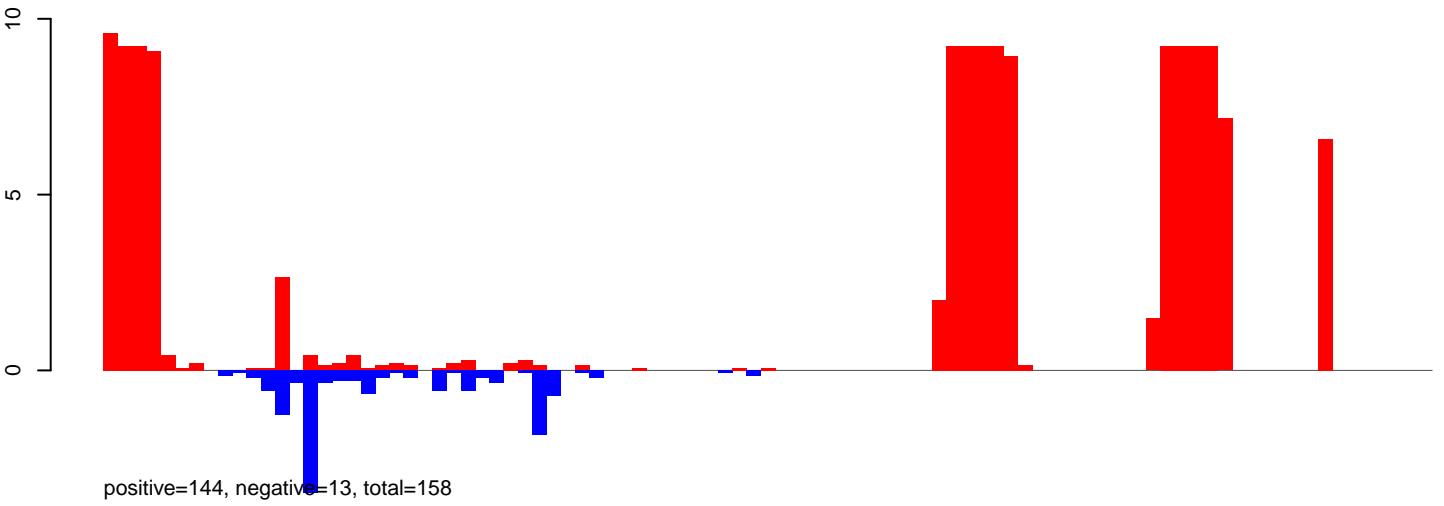


CuQuin_Testes_NL.rep

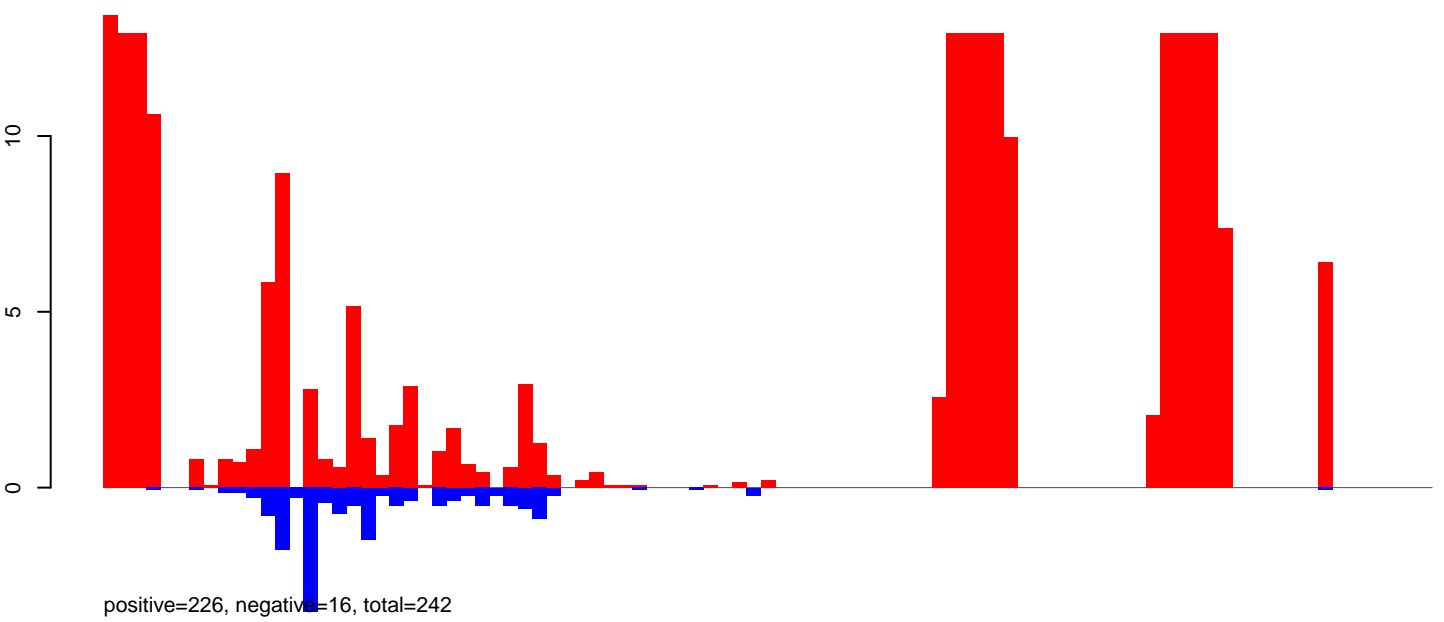


Window size=25, length=2088, TE@rnd-4_family-135@DNA_Zator@173:1-2088

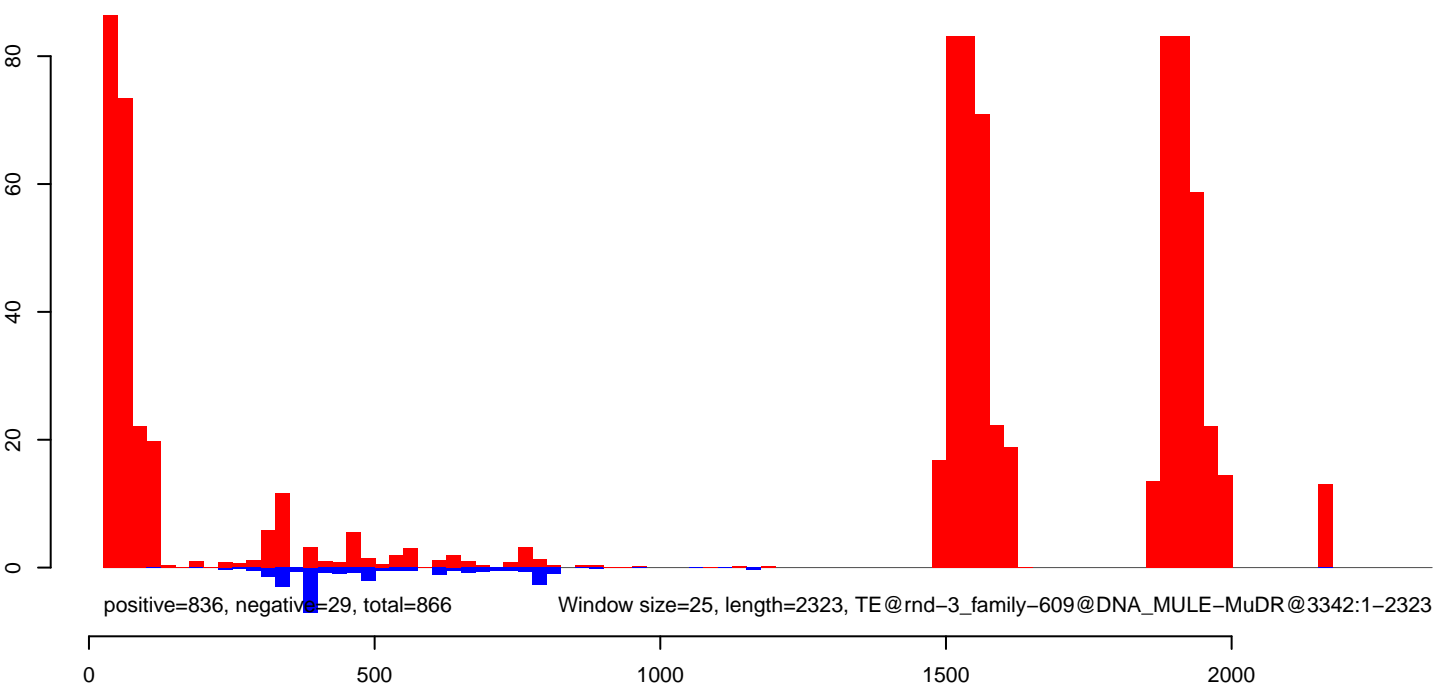
CuQuin_Testes_NL.18_23.rep



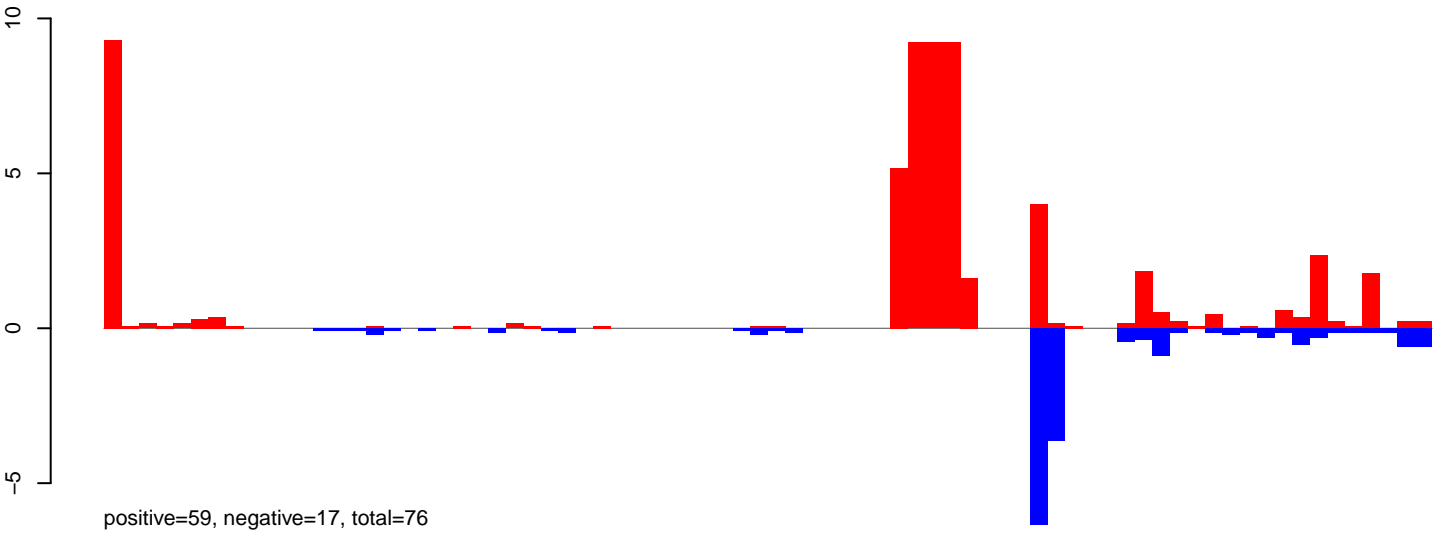
CuQuin_Testes_NL.24_35.rep



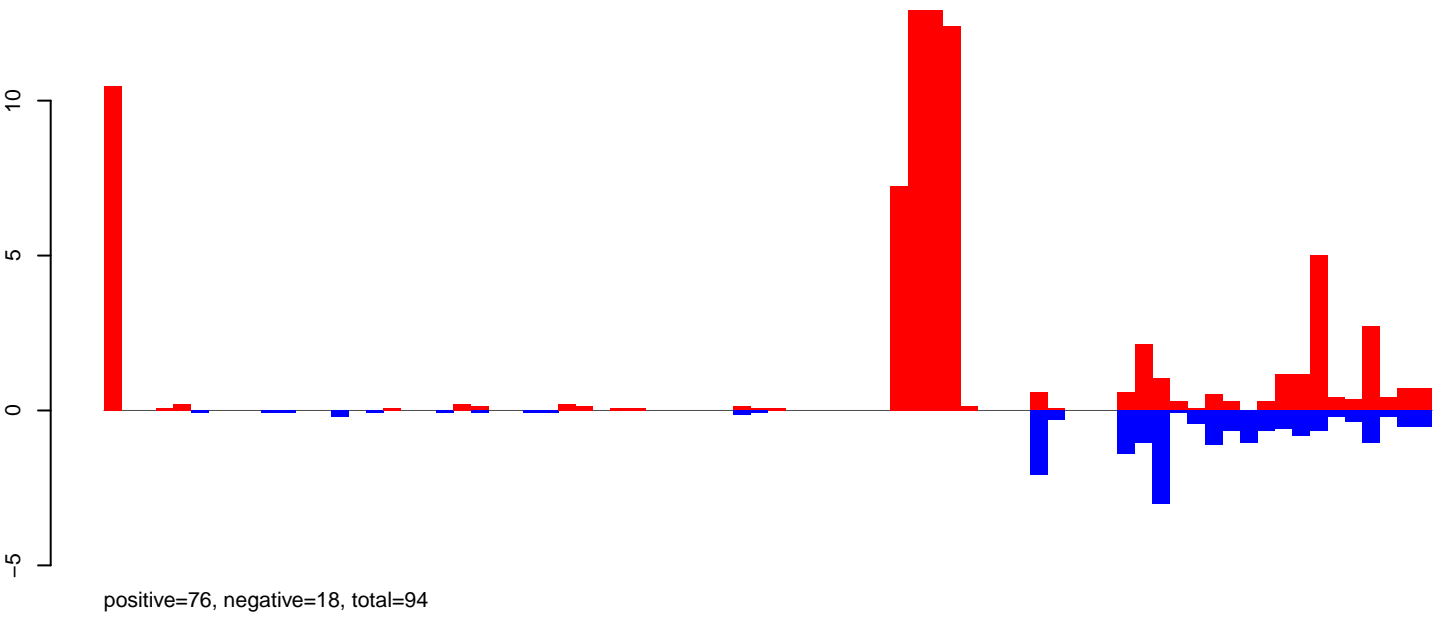
CuQuin_Testes_NL.rep



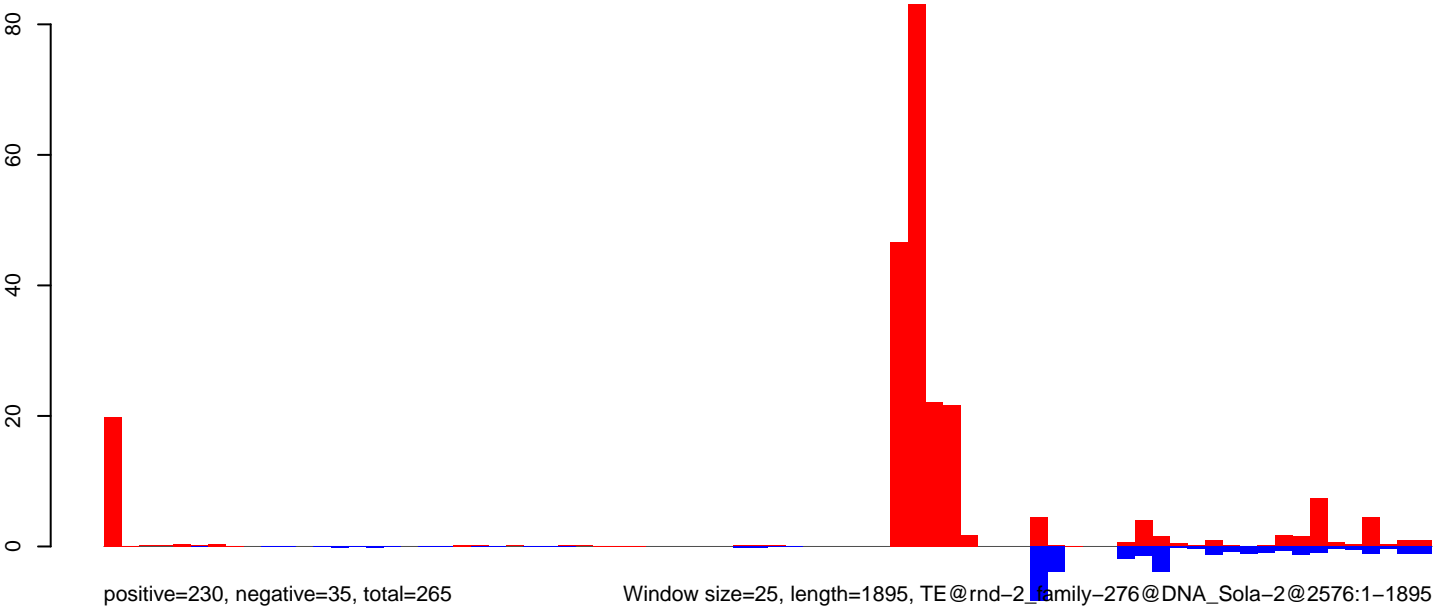
CuQuin_Testes_NL.18_23.rep



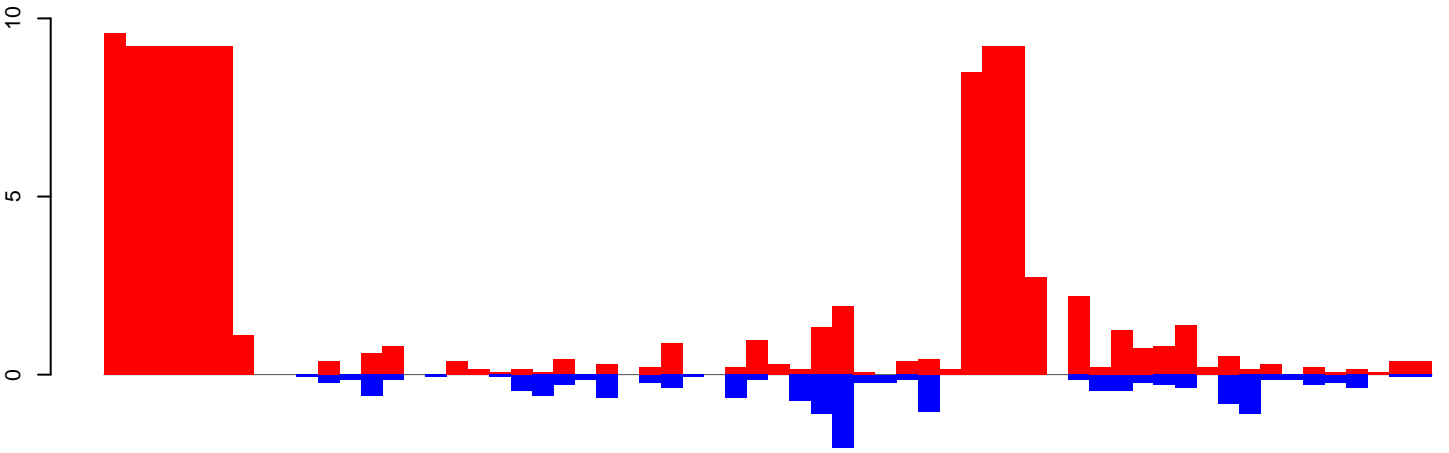
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

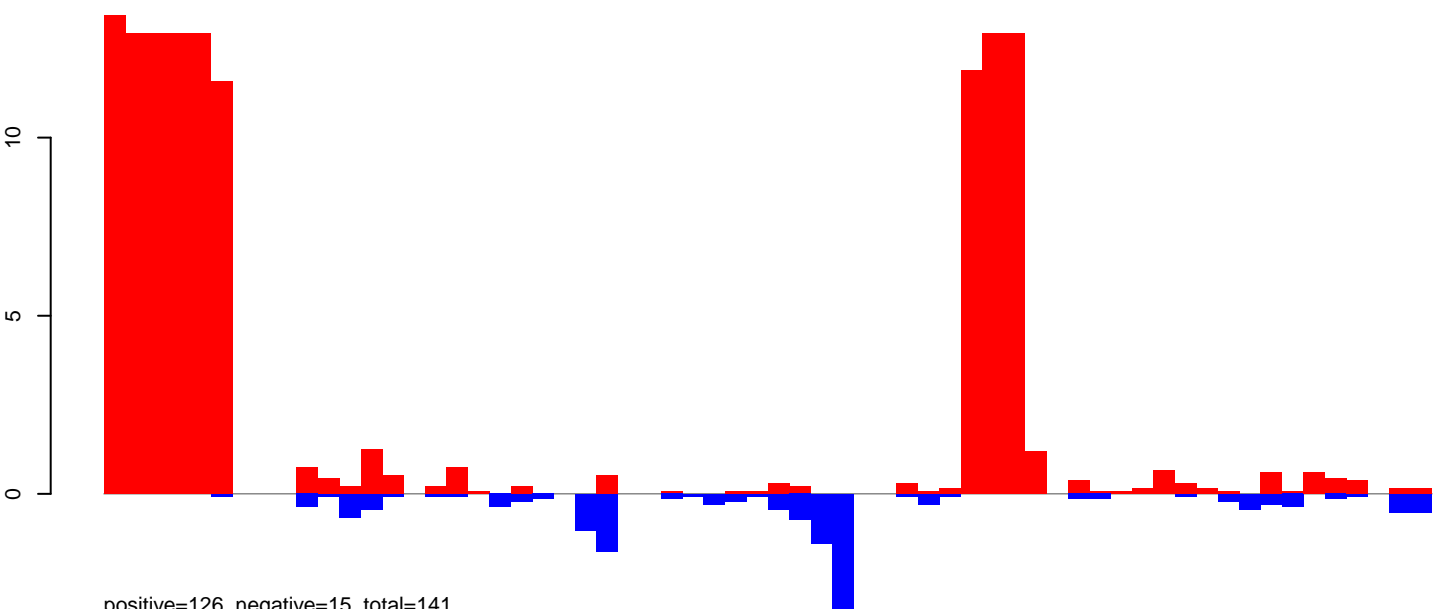


CuQuin_Testes_NL.18_23.rep



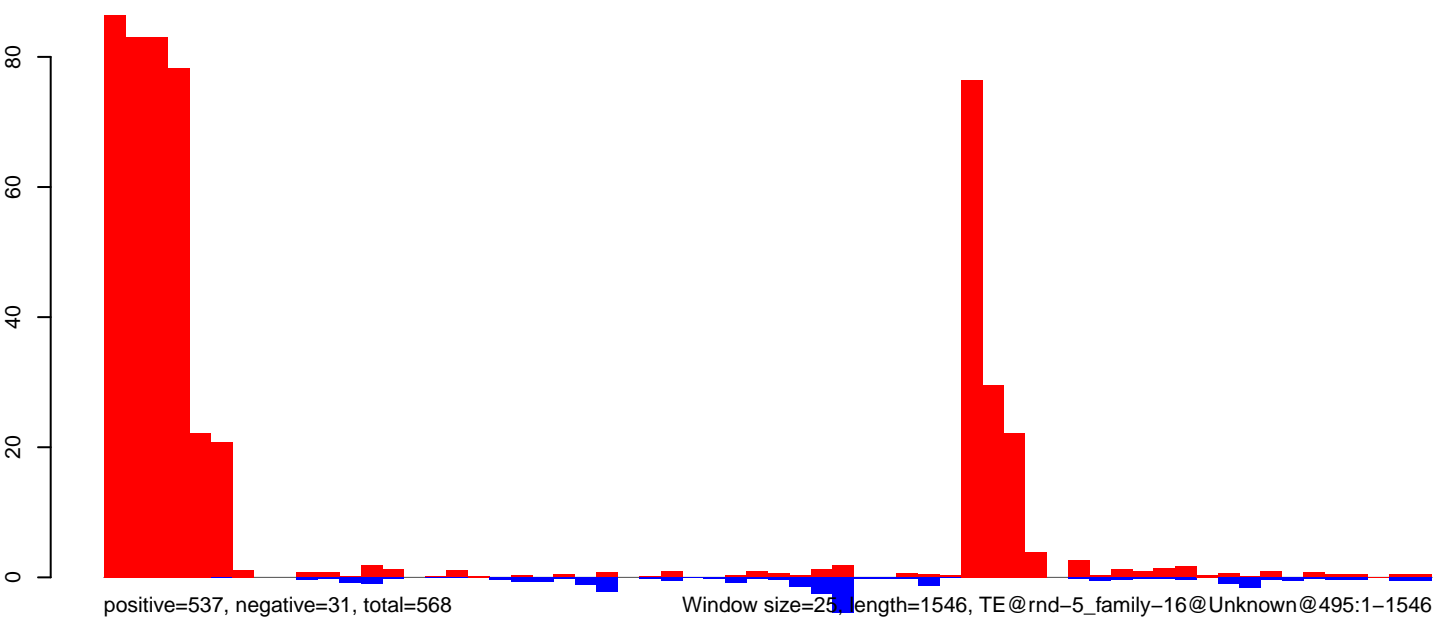
positive=106, negative=16, total=122

CuQuin_Testes_NL.24_35.rep



positive=126, negative=15, total=141

CuQuin_Testes_NL.rep

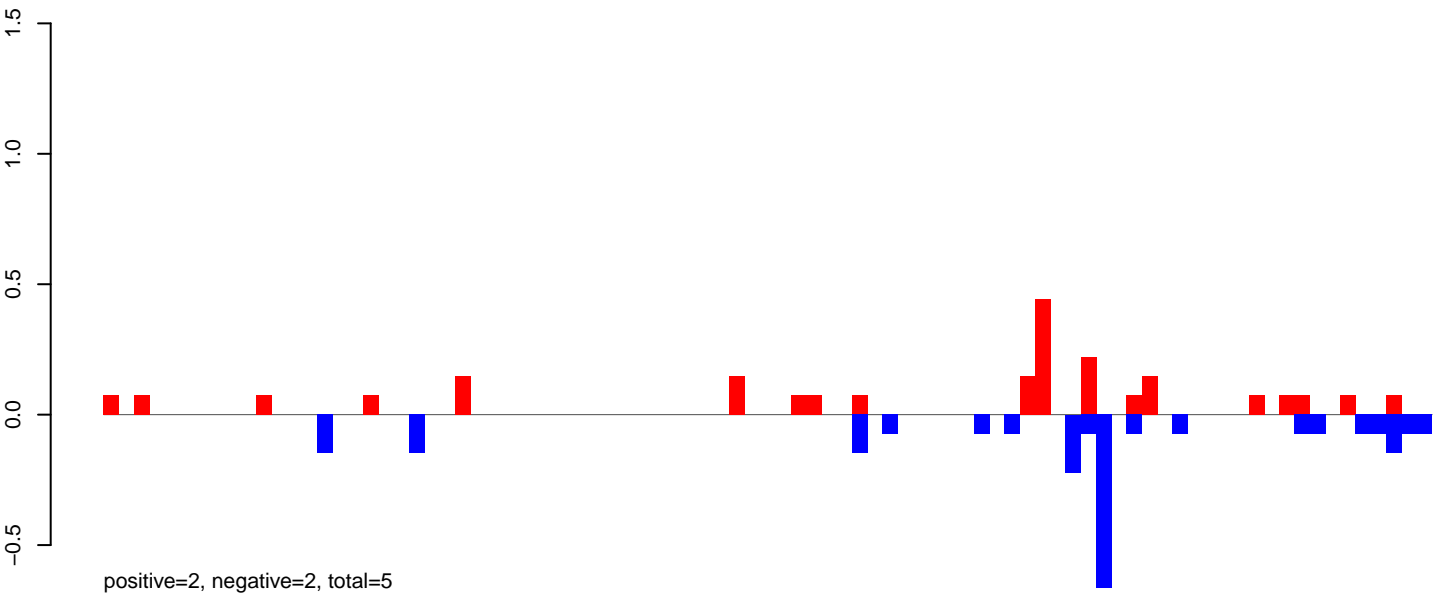


positive=537, negative=31, total=568

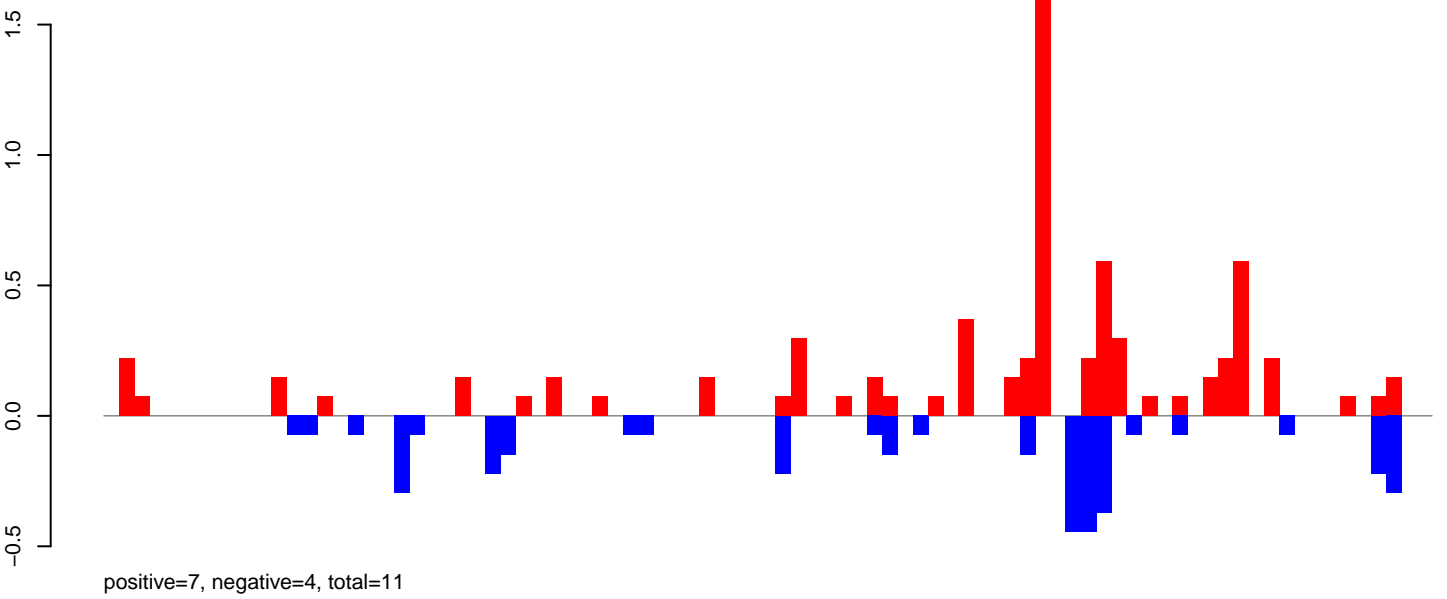
Window size=25, length=1546, TE@rnd-5_family-16@Unknown@495:1-1546

0 500 1000 1500

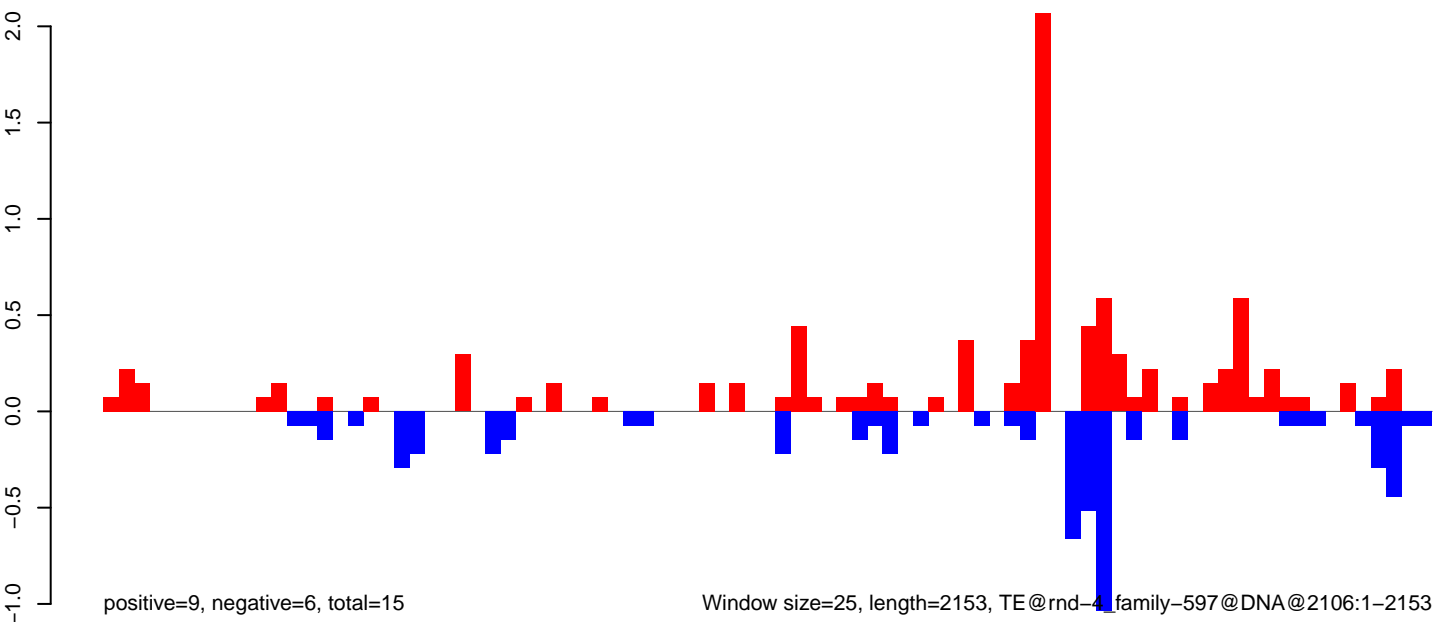
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

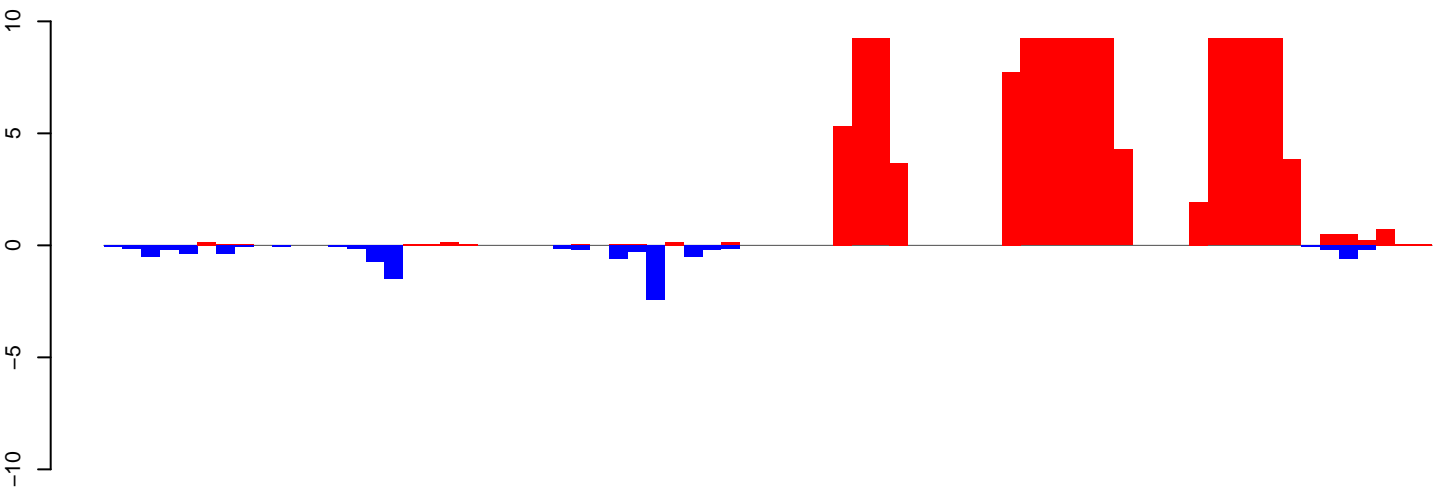


CuQuin_Testes_NL.rep



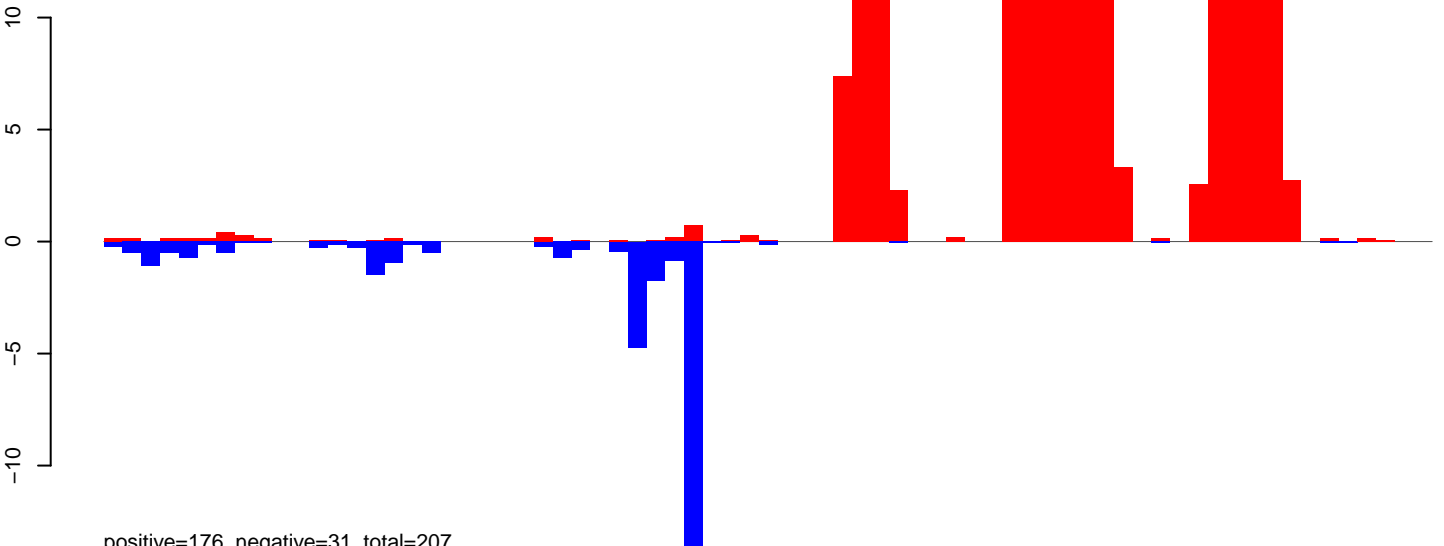
Window size=25, length=2153, TE@rnd-4, family-597@DNA@2106:1-2153

CuQuin_Testes_NL.18_23.rep



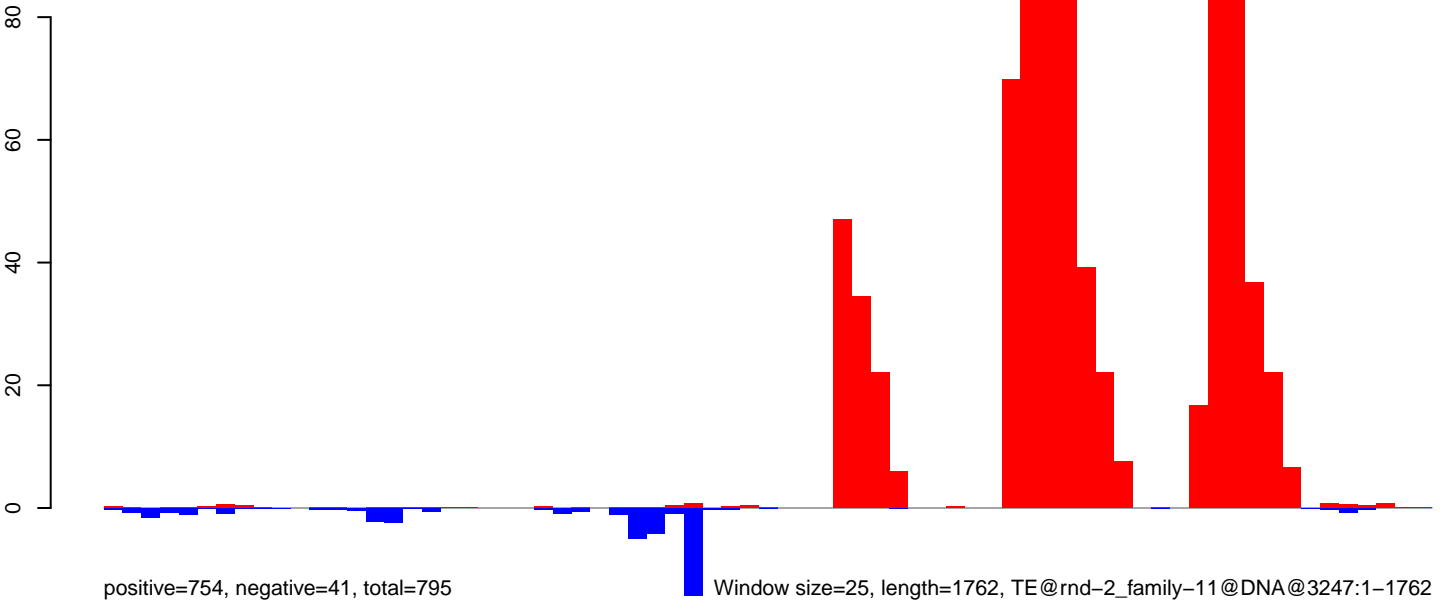
positive=132, negative=10, total=142

CuQuin_Testes_NL.24_35.rep



positive=176, negative=31, total=207

CuQuin_Testes_NL.rep

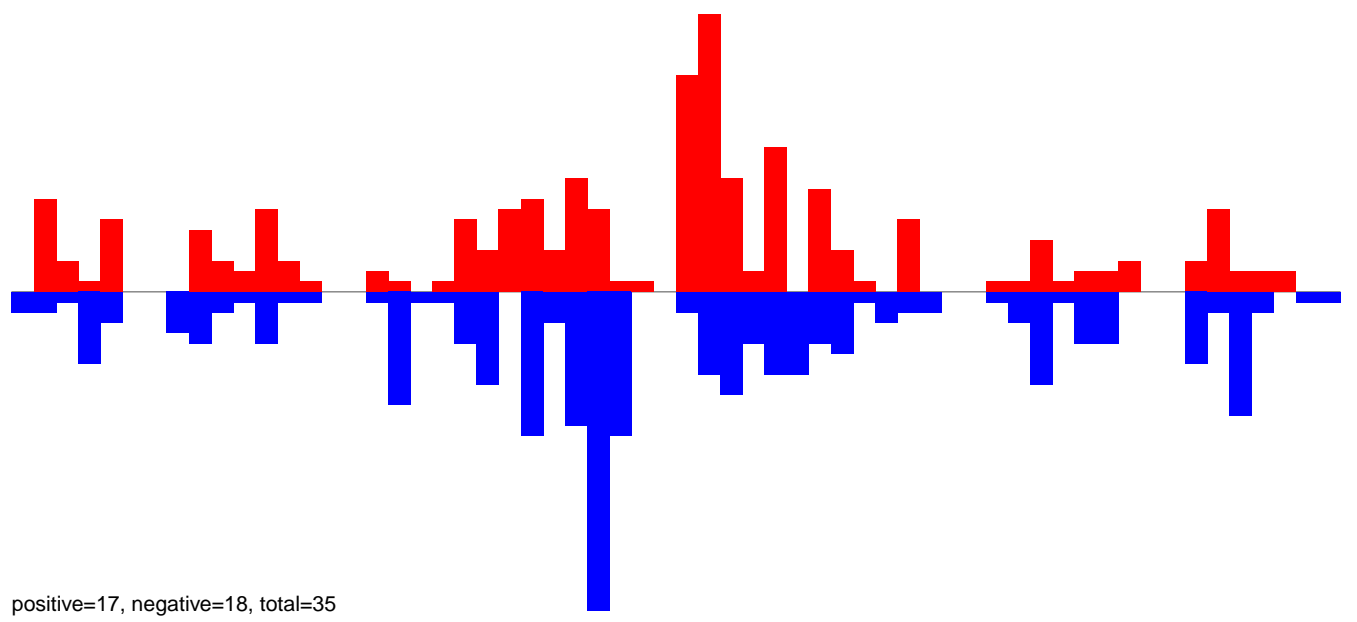


positive=754, negative=41, total=795

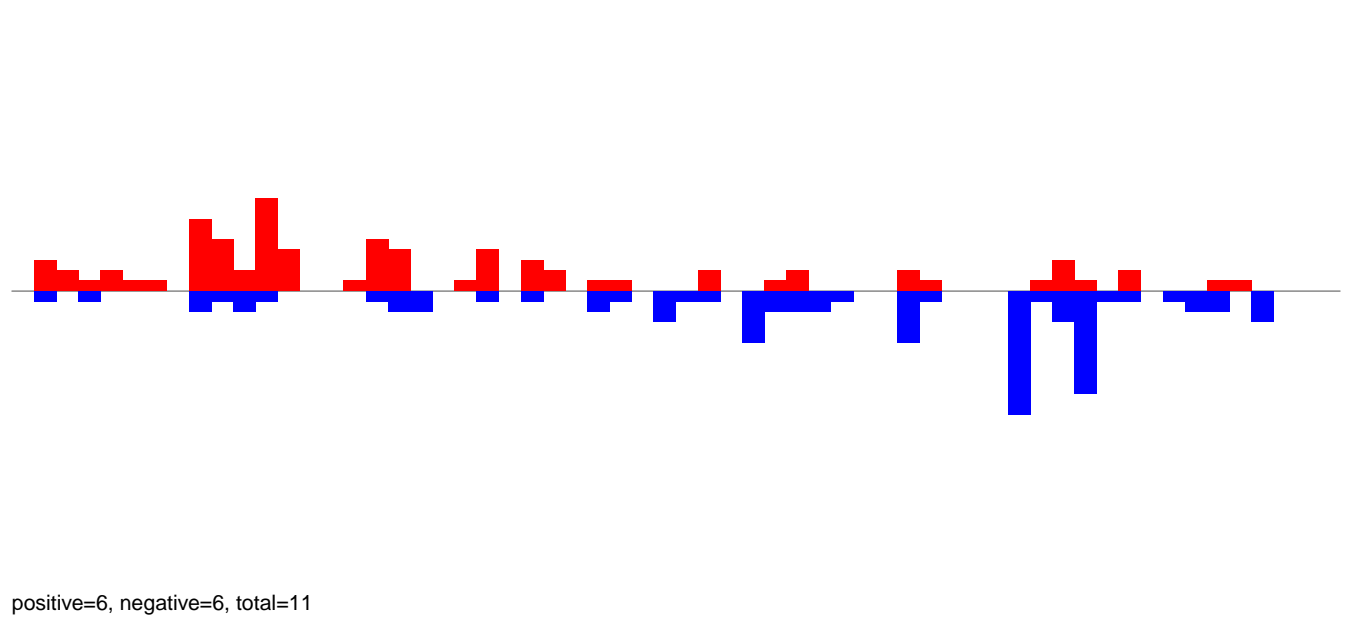
Window size=25, length=1762, TE@rnd-2_family-11@DNA@3247:1-1762

0 500 1000 1500

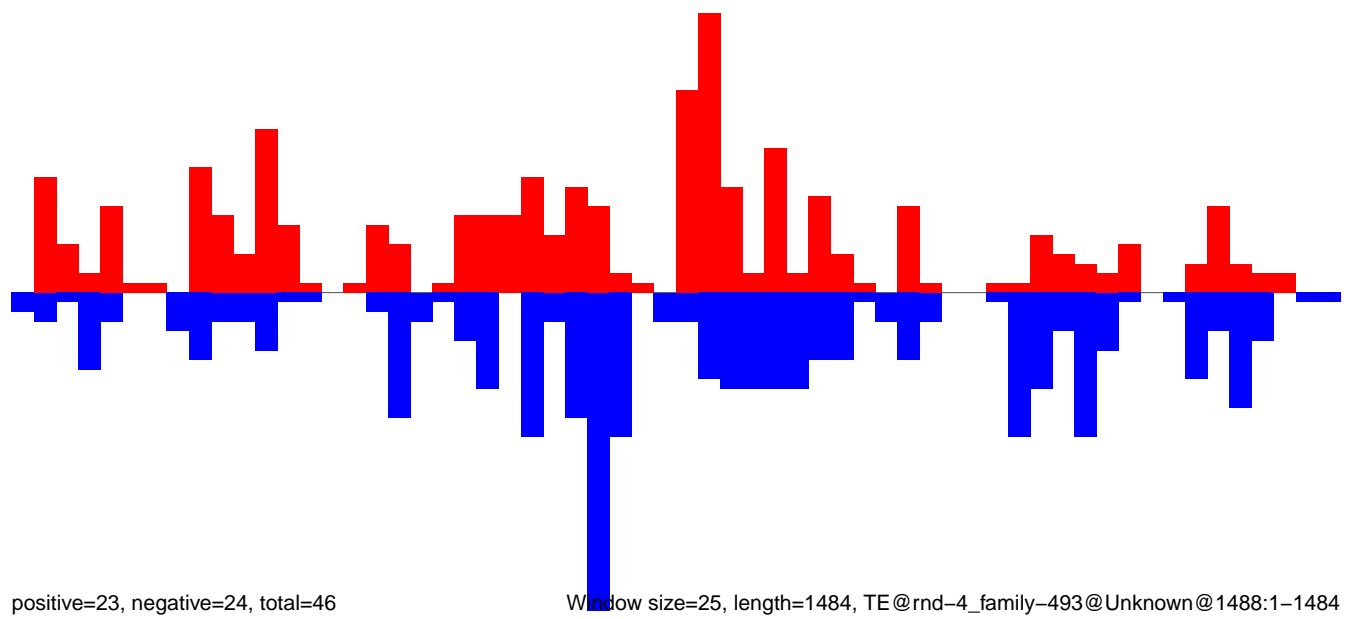
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

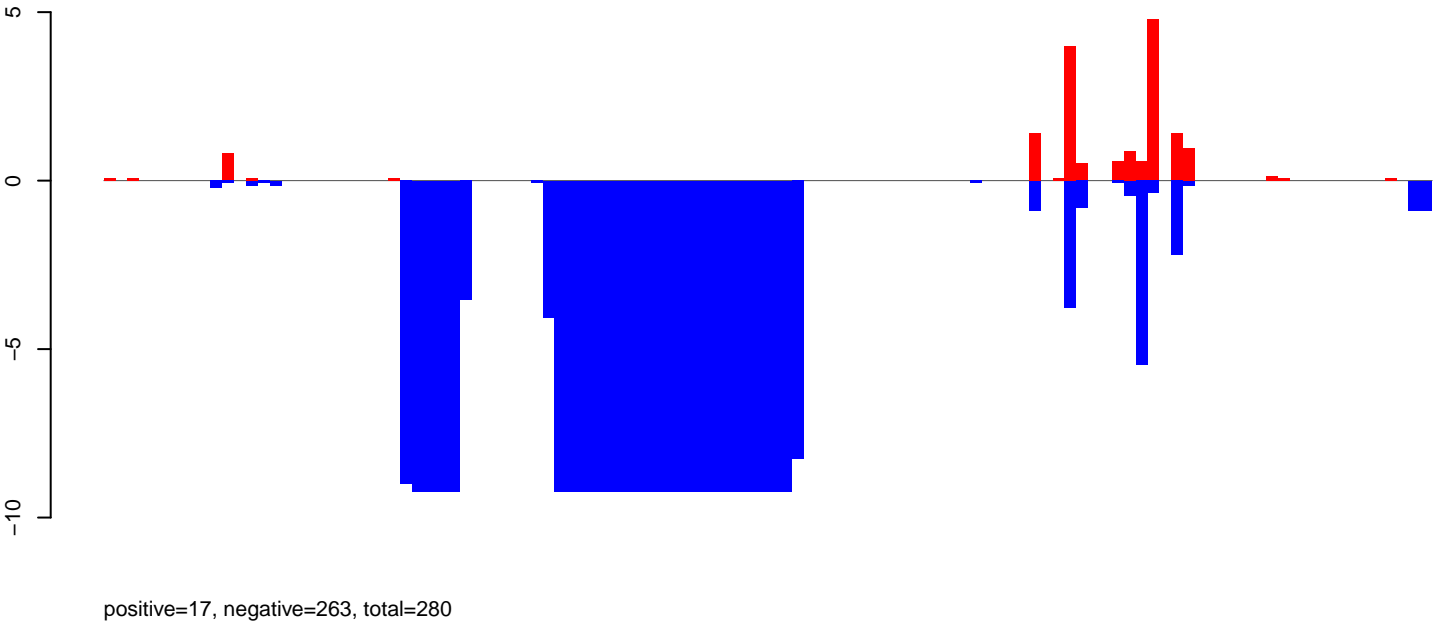


CuQuin_Testes_NL.rep

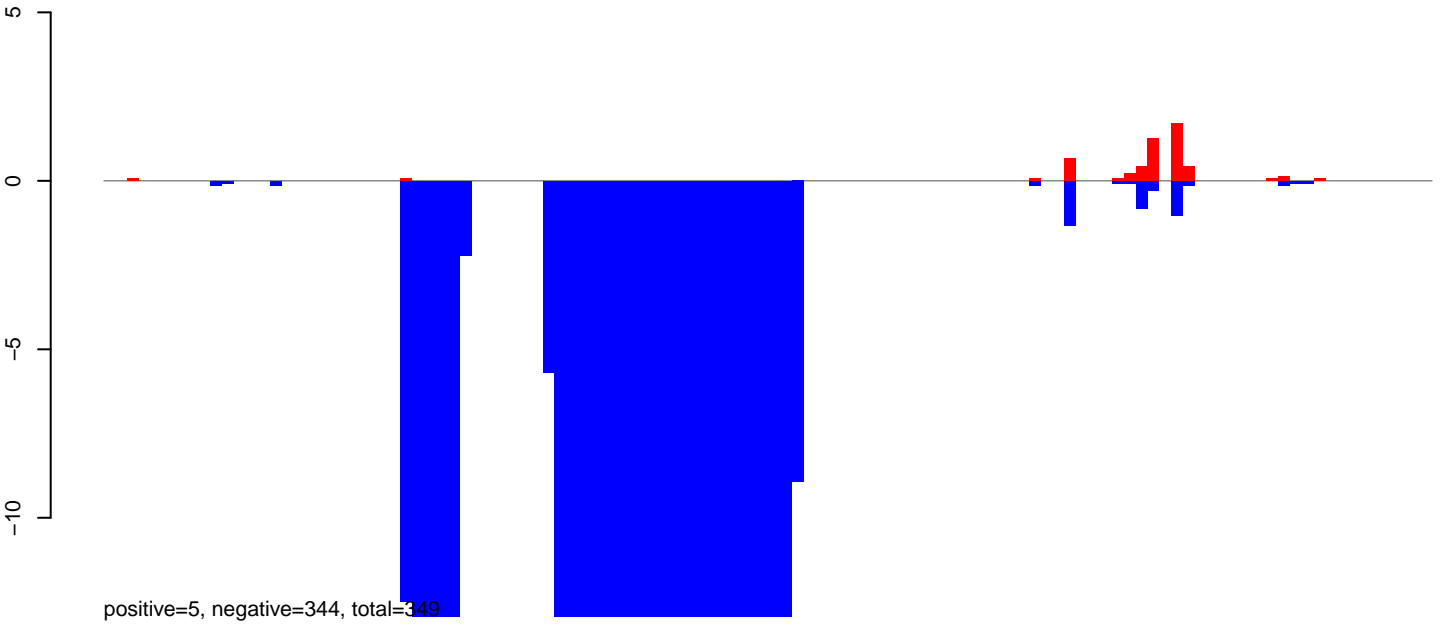


Window size=25, length=1484, TE@rnd-4_family-493@Unknown@1488:1-1484

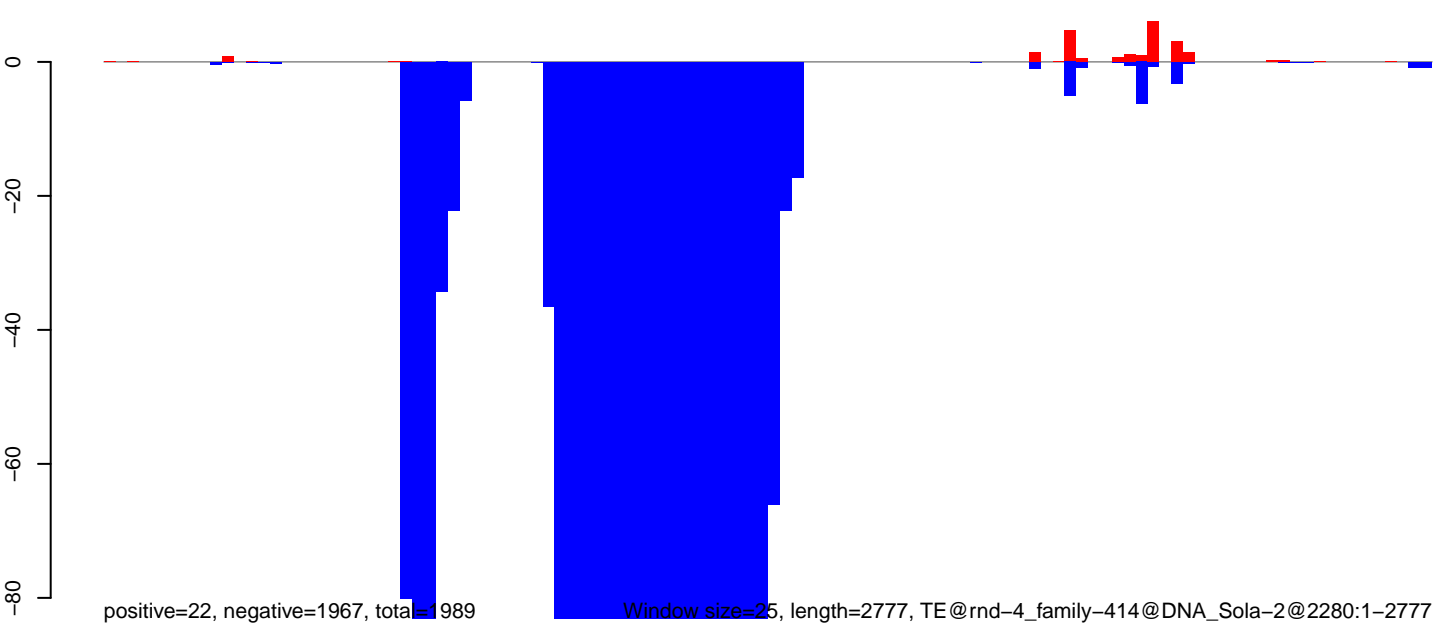
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

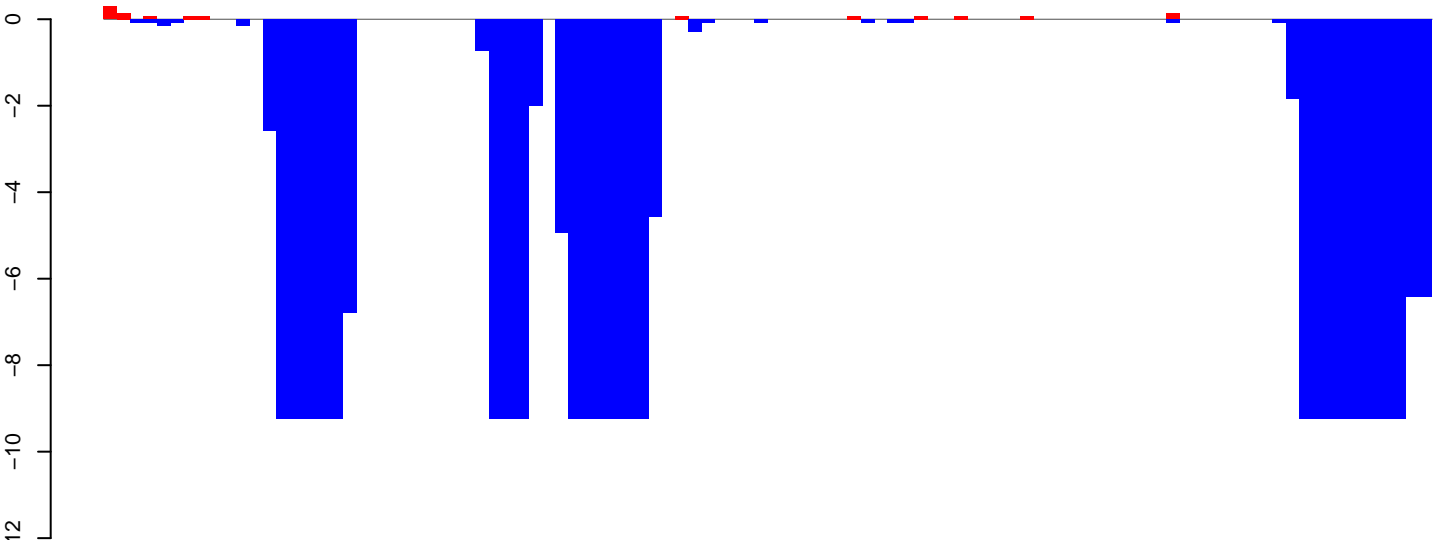


CuQuin_Testes_NL.rep



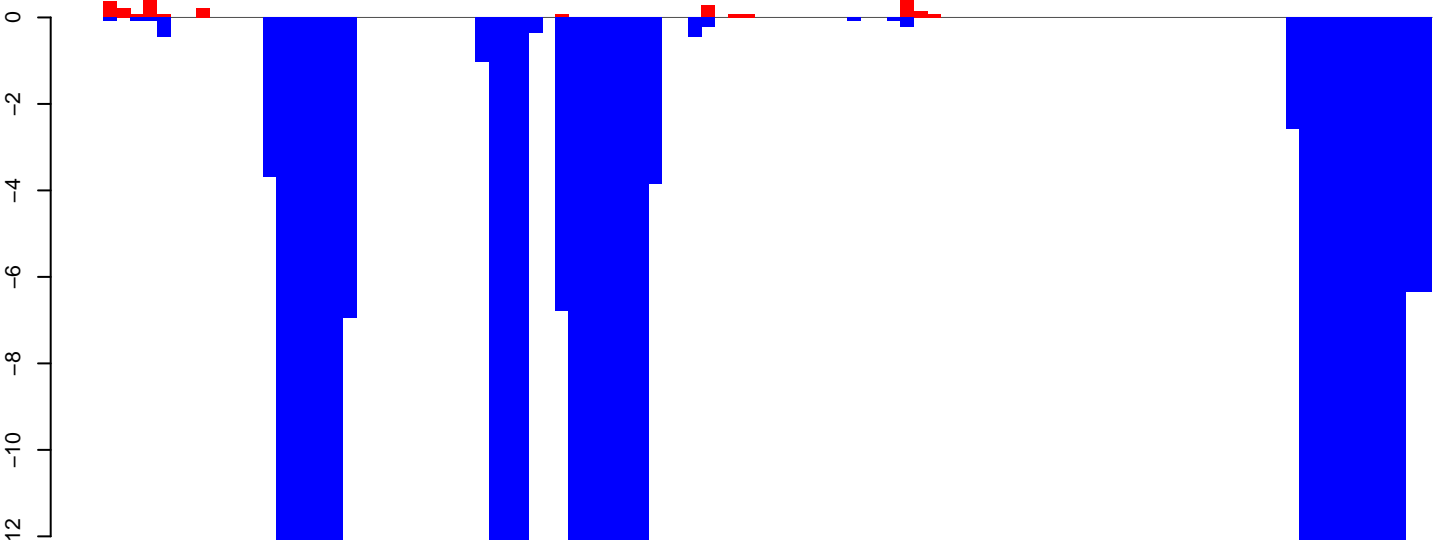
0 500 1000 1500 2000 2500

CuQuin_Testes_NL.18_23.rep



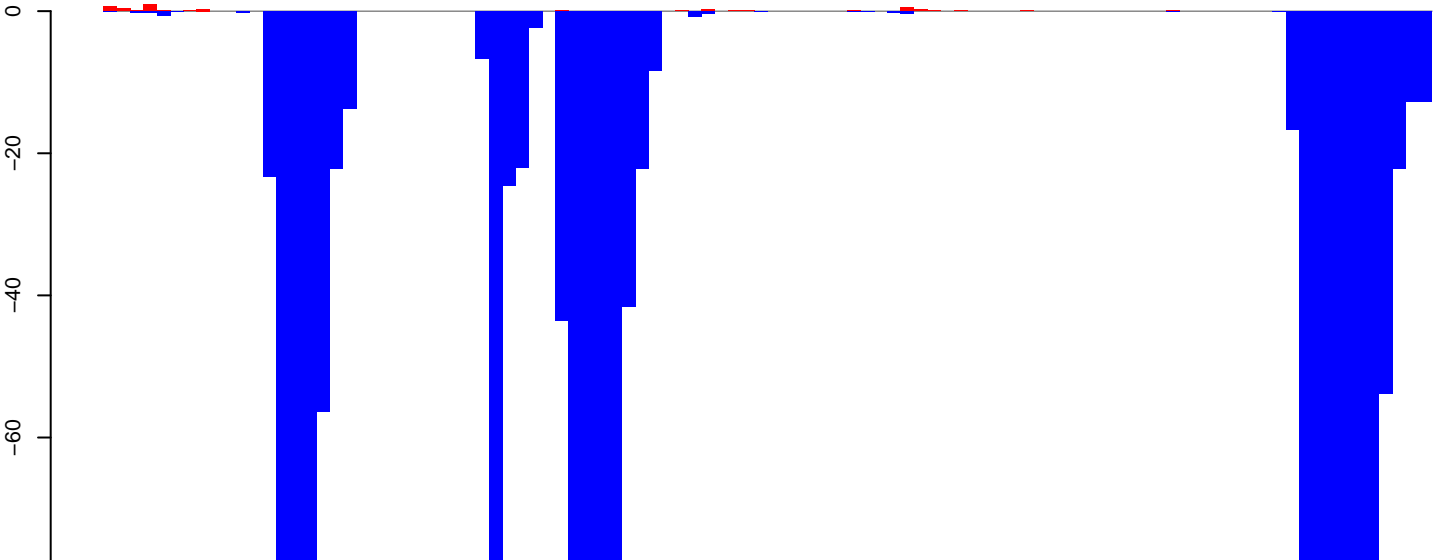
positive=1, negative=241, total=242

CuQuin_Testes_NL.24_35.rep



positive=3, negative=324, total=327

CuQuin_Testes_NL.rep

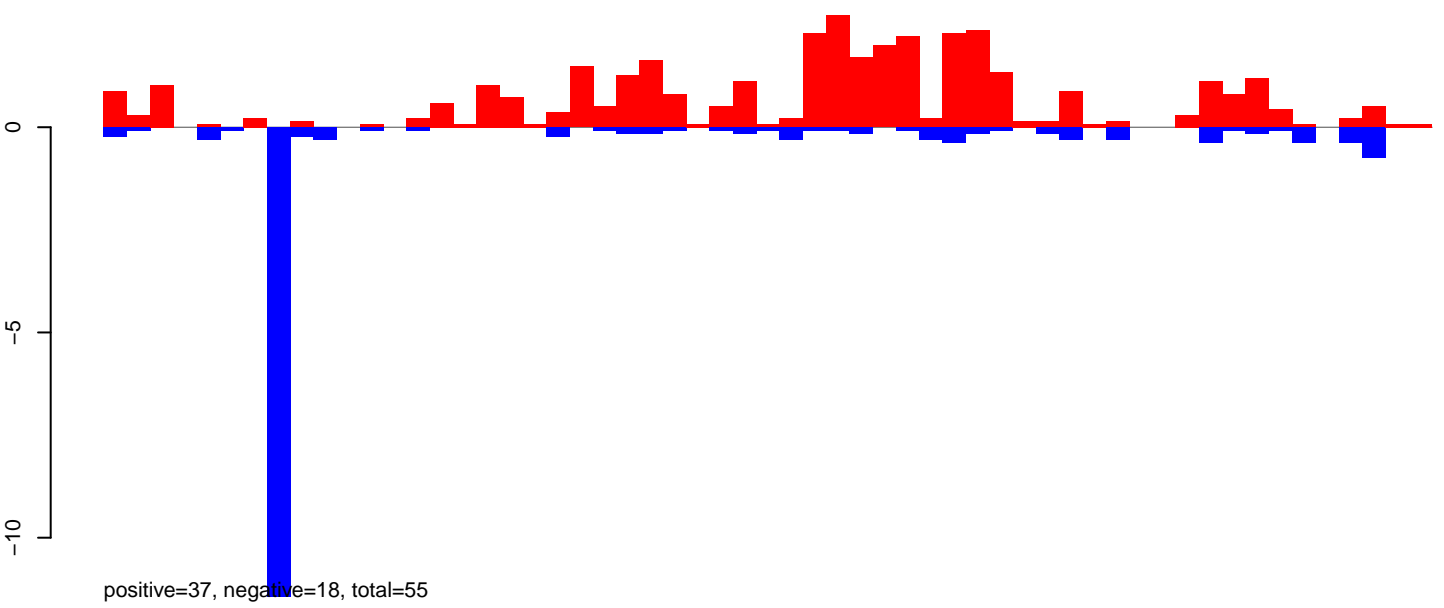


positive=4, negative=1571, total=1575

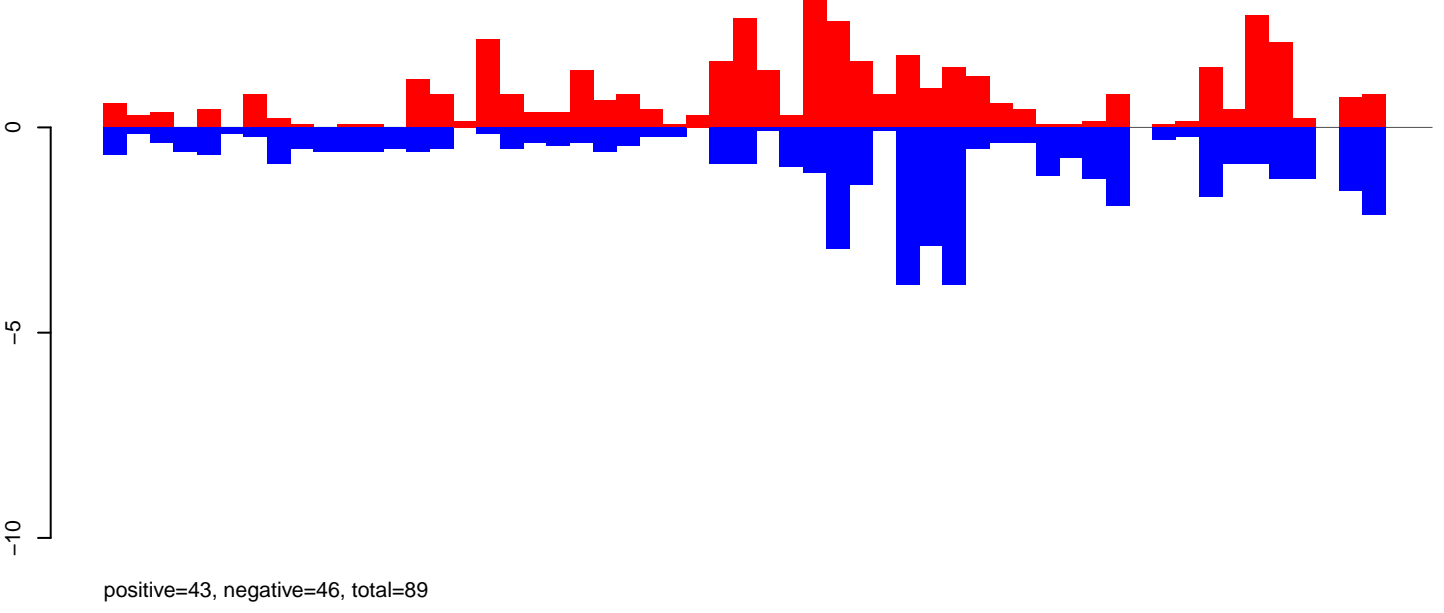
Window size=25, length=2489, TE@rnd-3_family-266@DNA_Zator@3279:1-2489

0 500 1000 1500 2000 2500

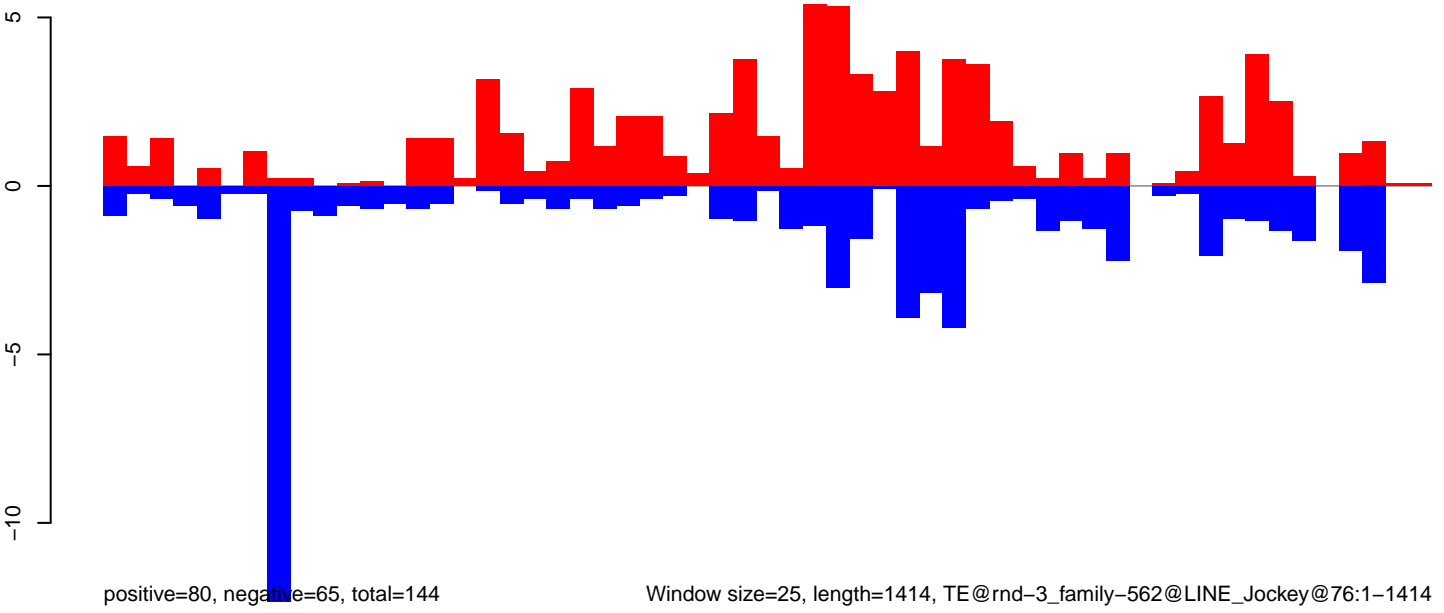
CuQuin_Testes_NL.18_23.rep



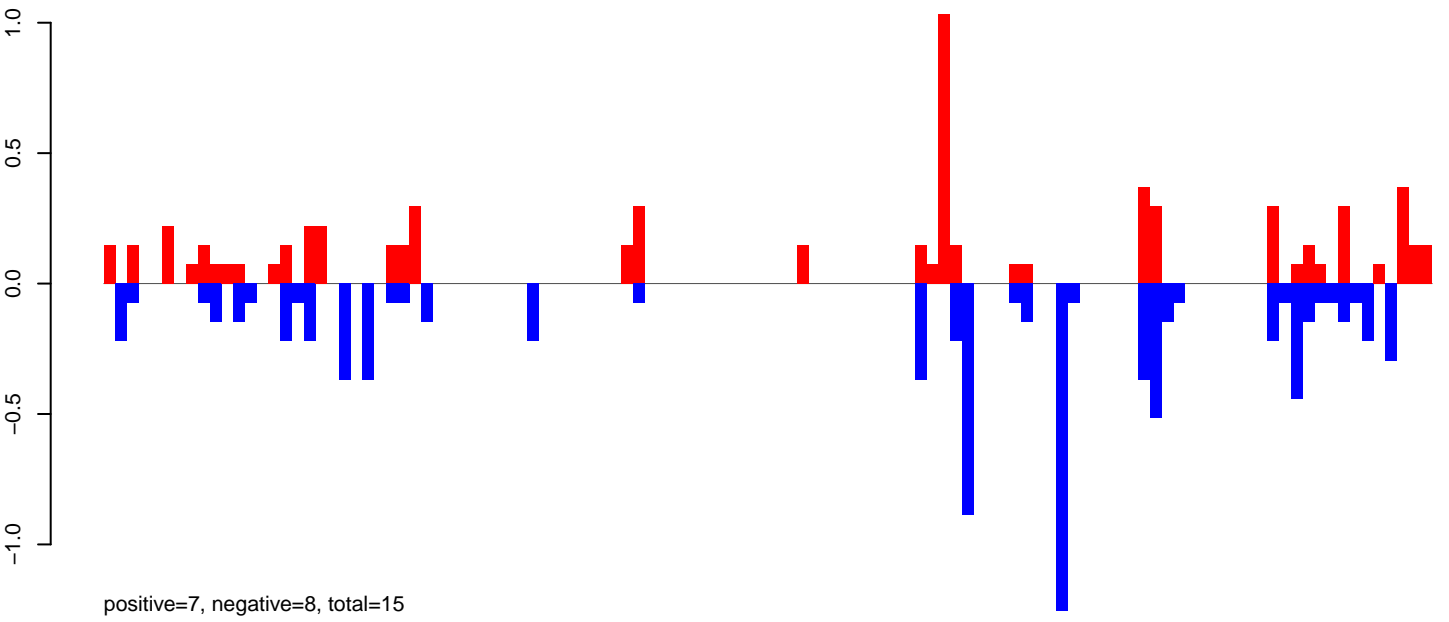
CuQuin_Testes_NL.24_35.rep



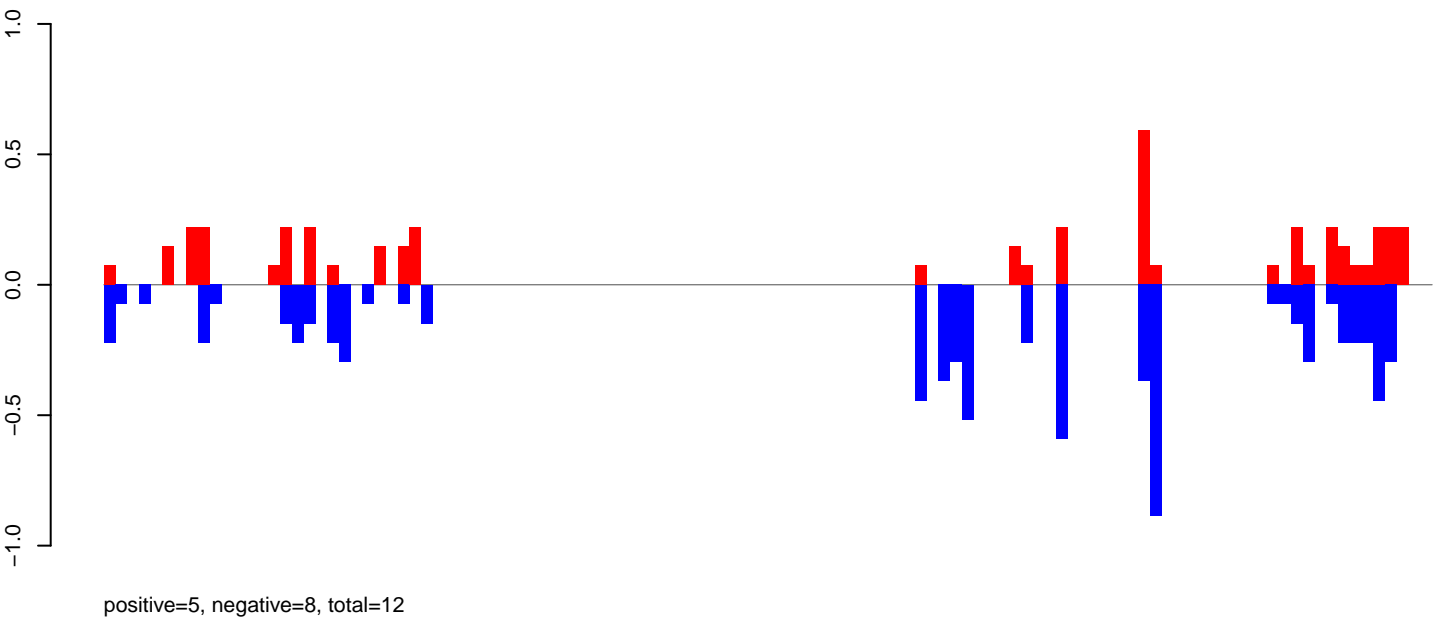
CuQuin_Testes_NL.rep



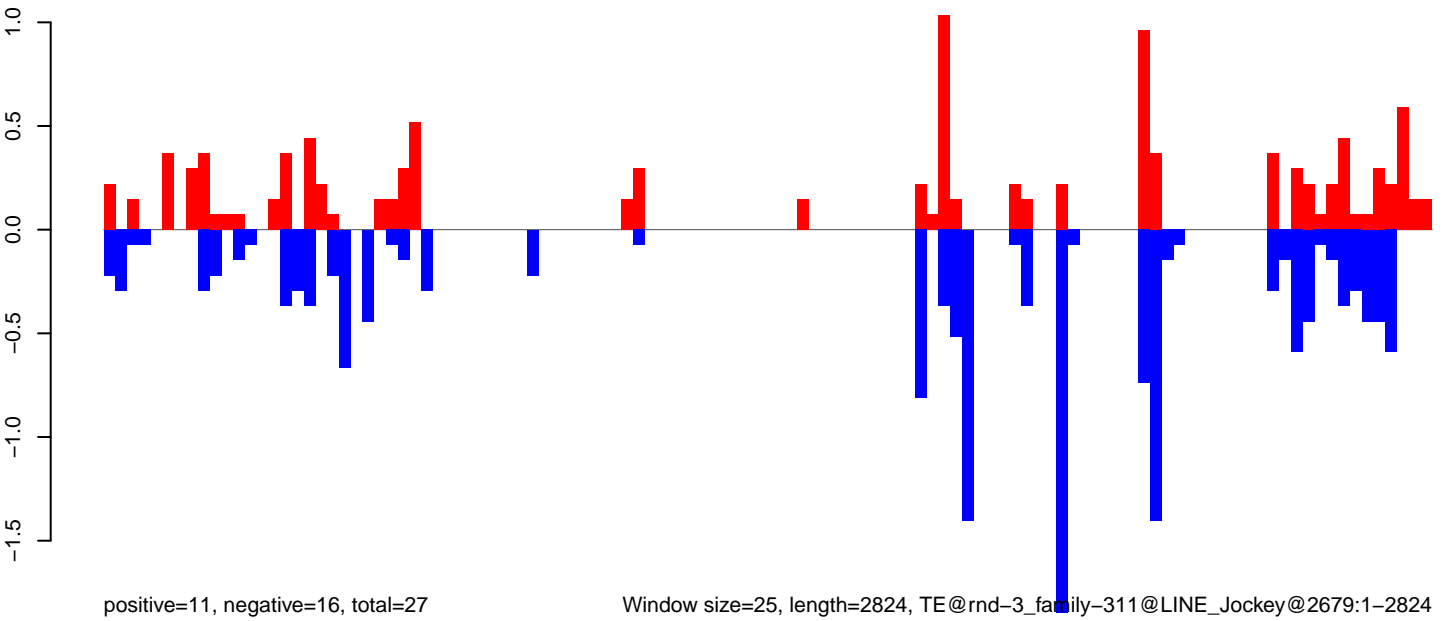
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

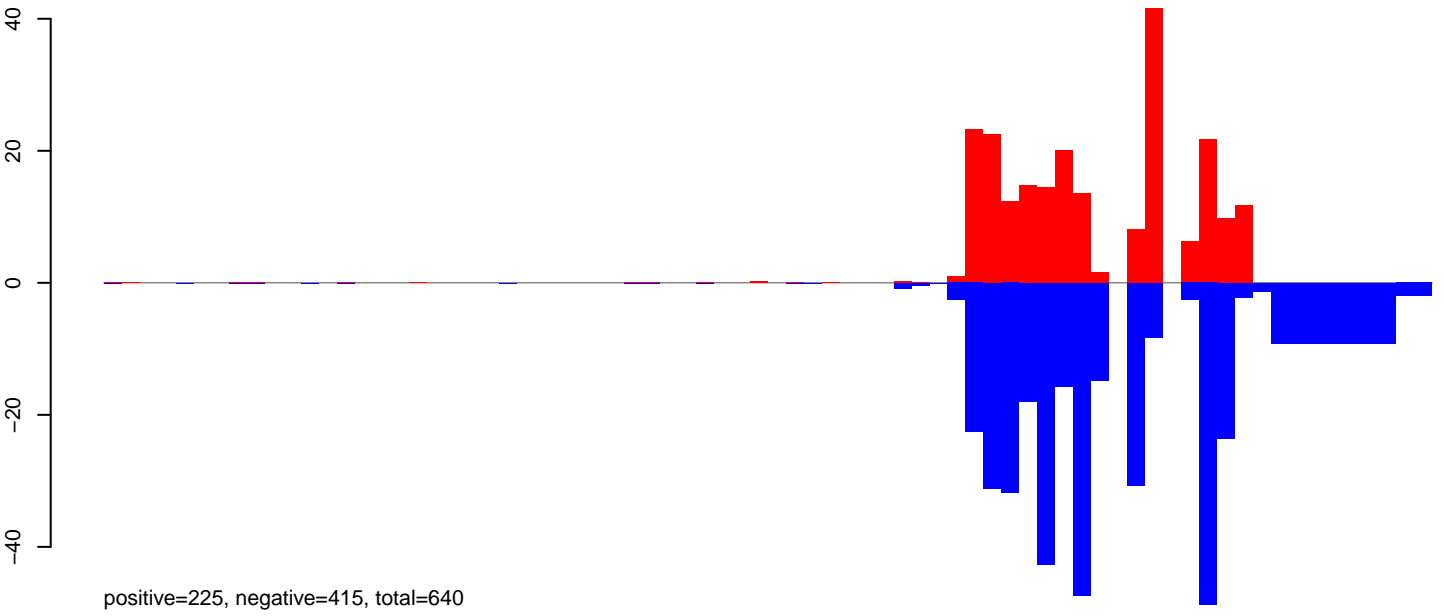


CuQuin_Testes_NL.rep

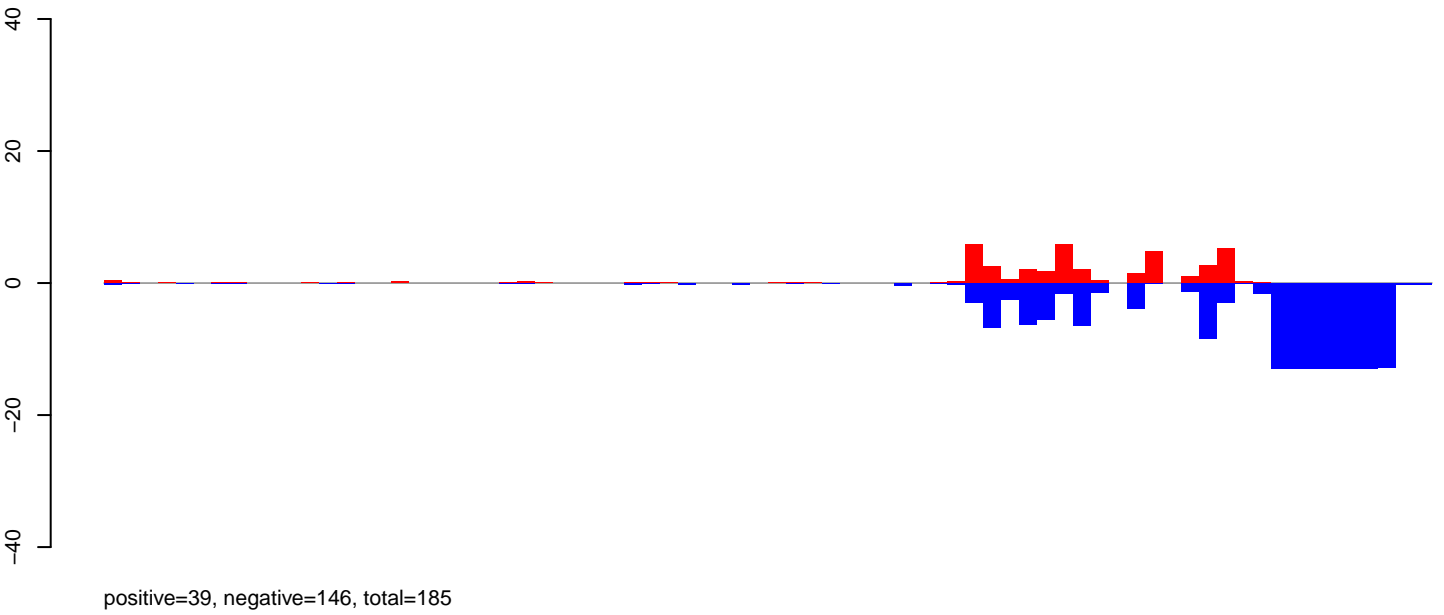


Window size=25, length=2824, TE@rnd-3_family-311@LINE_Jockey@2679:1-2824

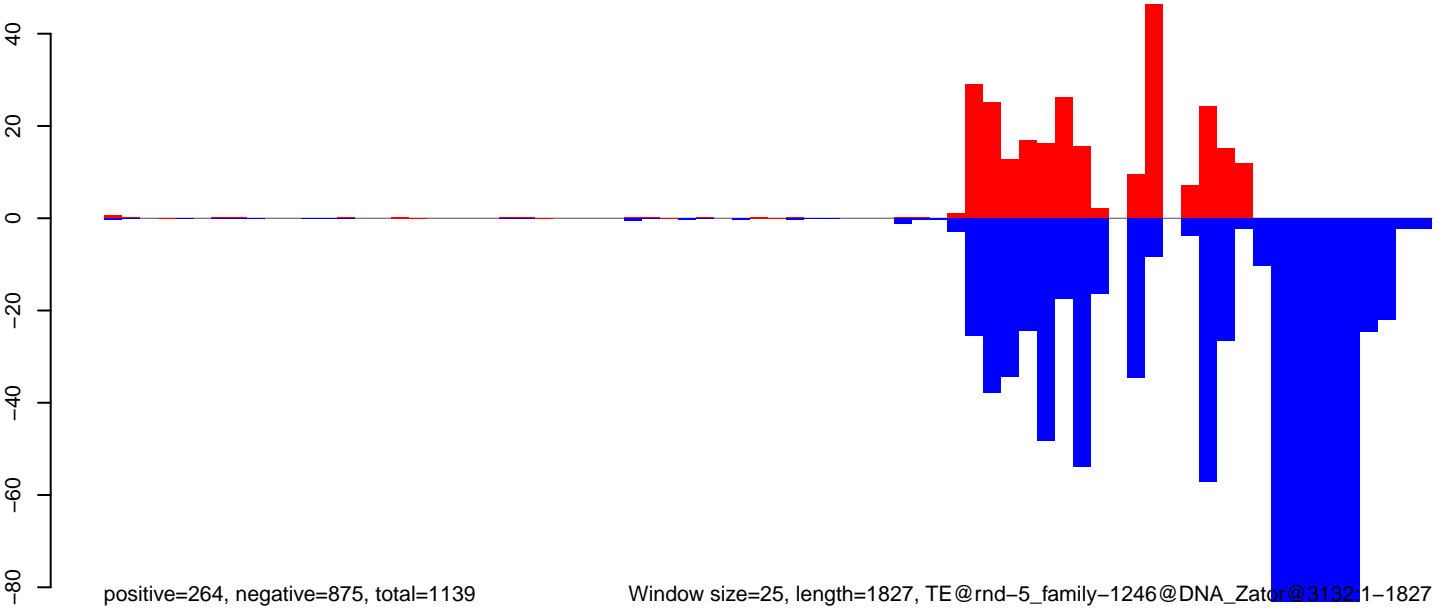
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

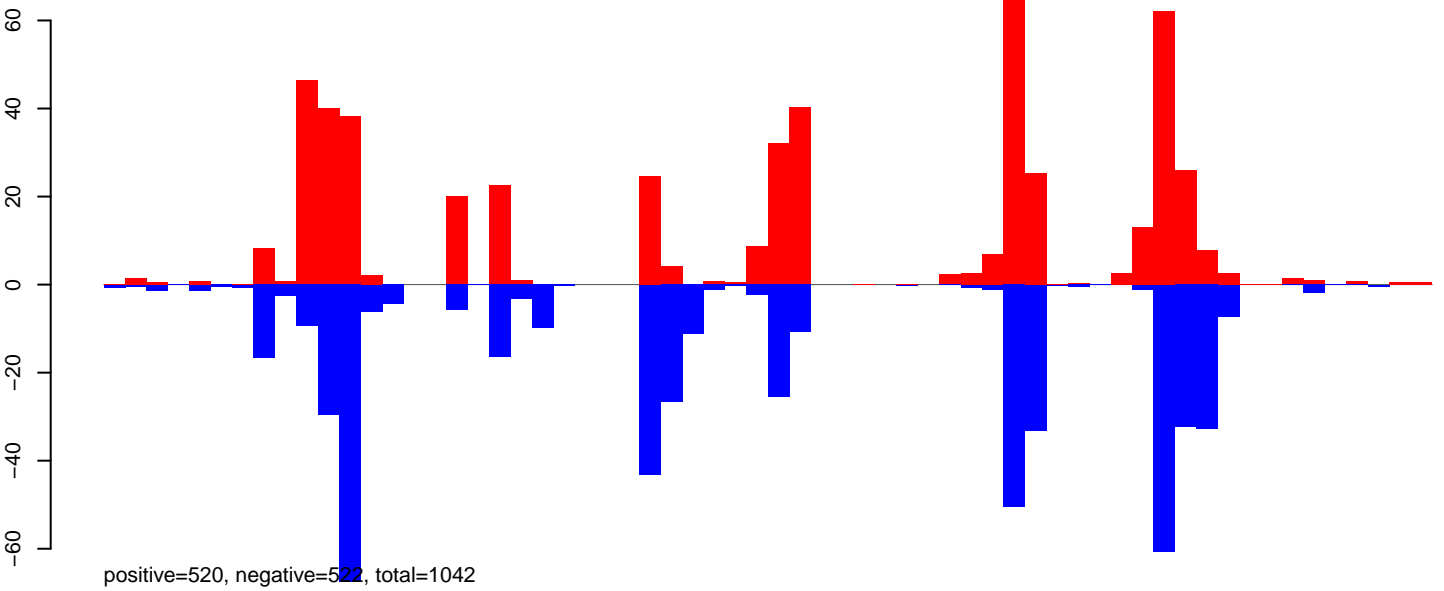


CuQuin_Testes_NL.rep

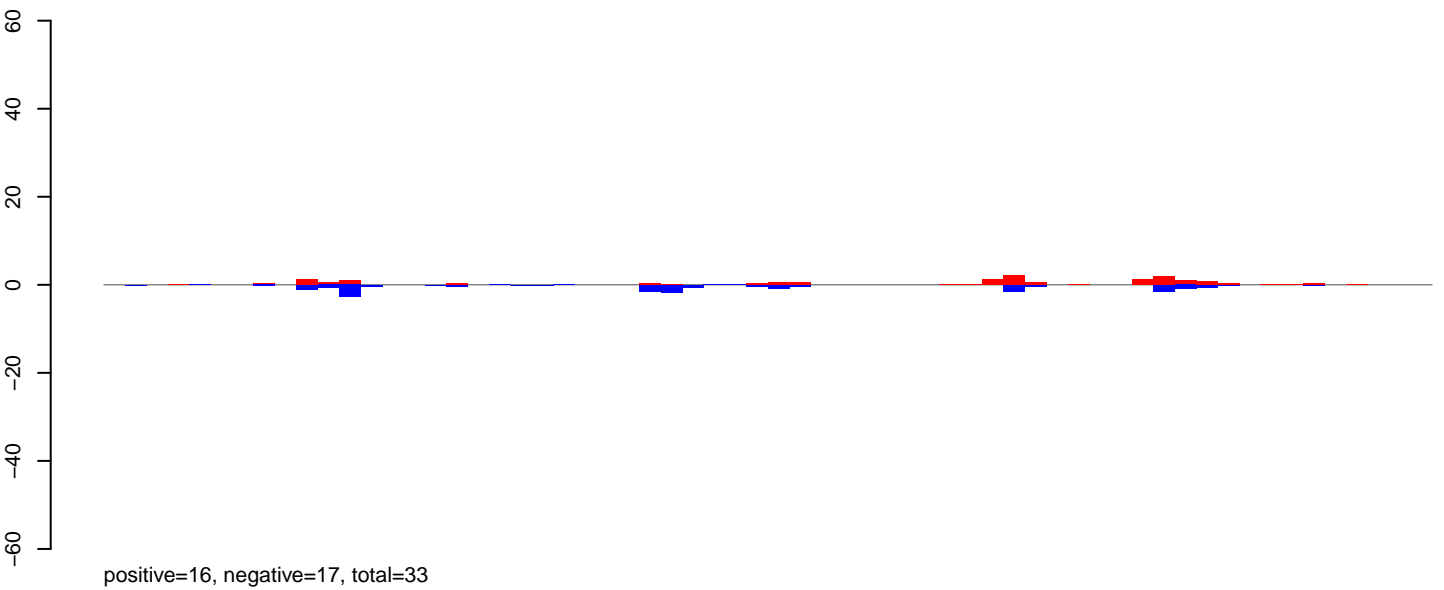


Window size=25, length=1827, TE@rnd-5_family-1246@DNA_Zator@3132:1-1827

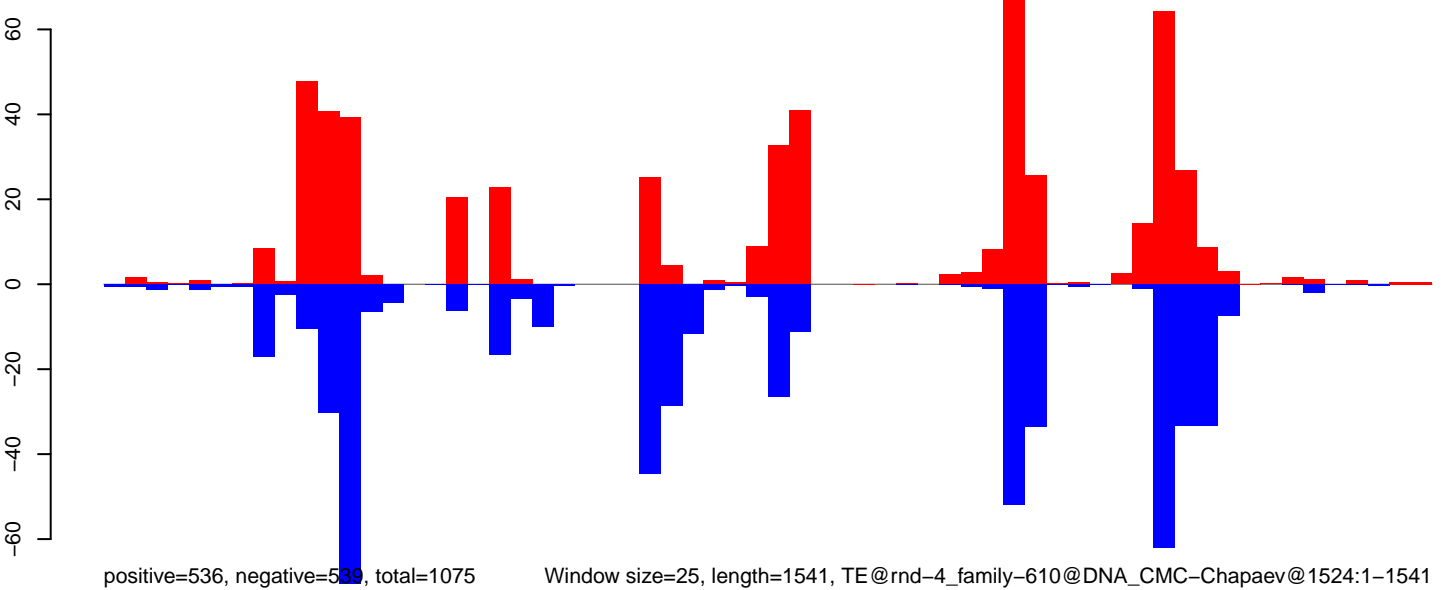
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

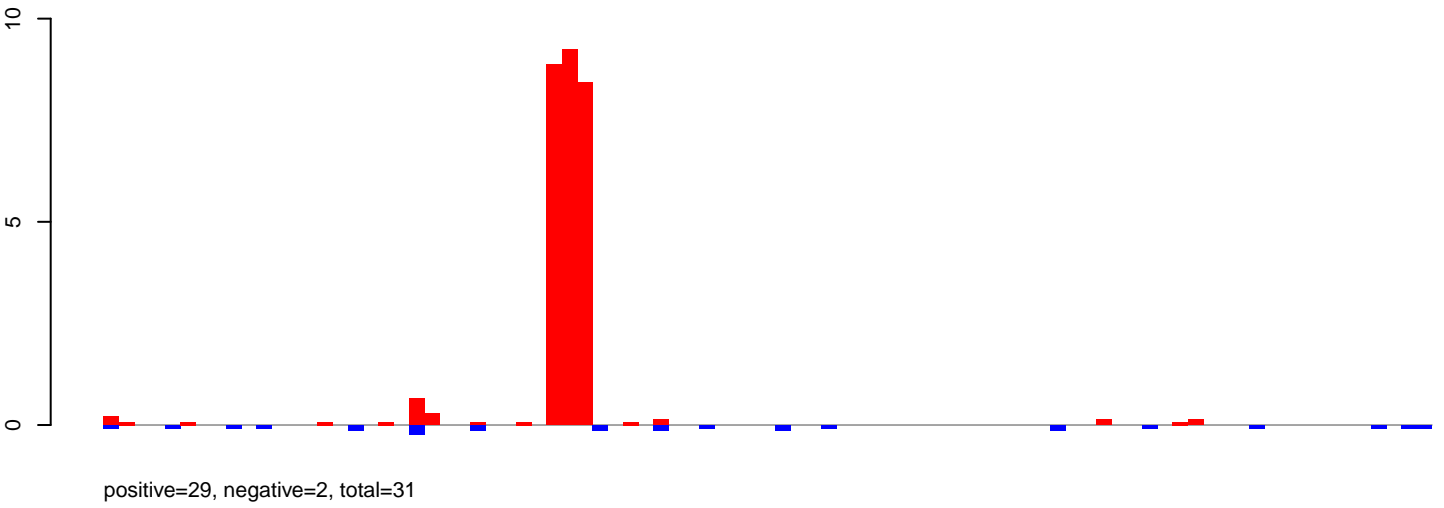


CuQuin_Testes_NL.rep

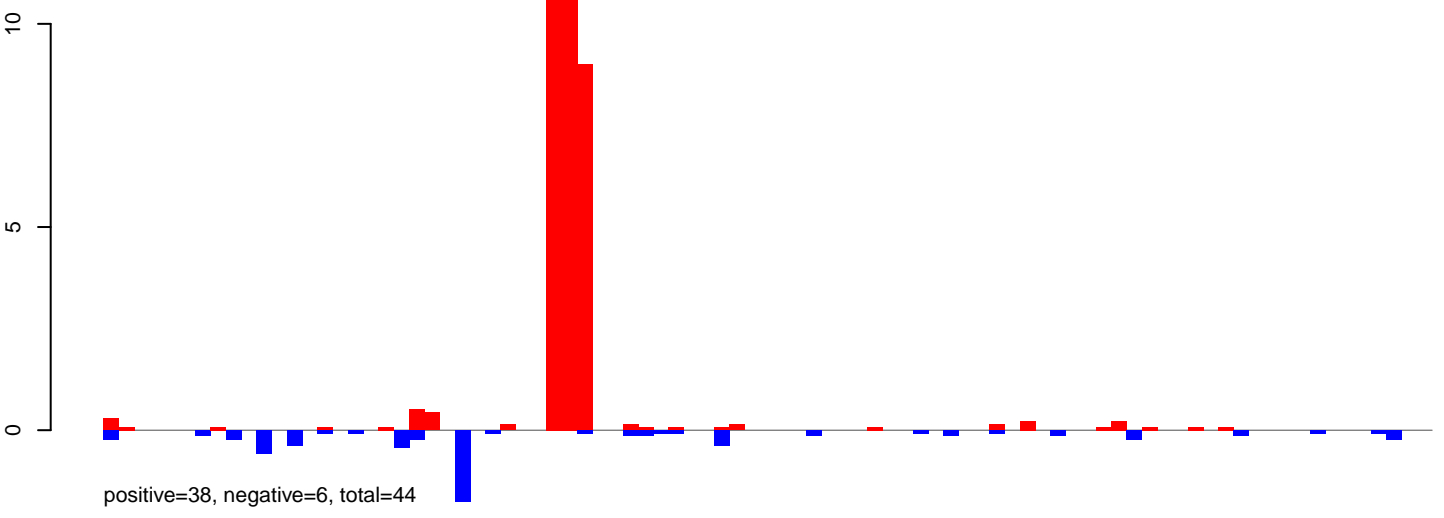


Window size=25, length=1541, TE@rnd-4_family-610@DNA_CMC-Chapaev@1524:1-1541

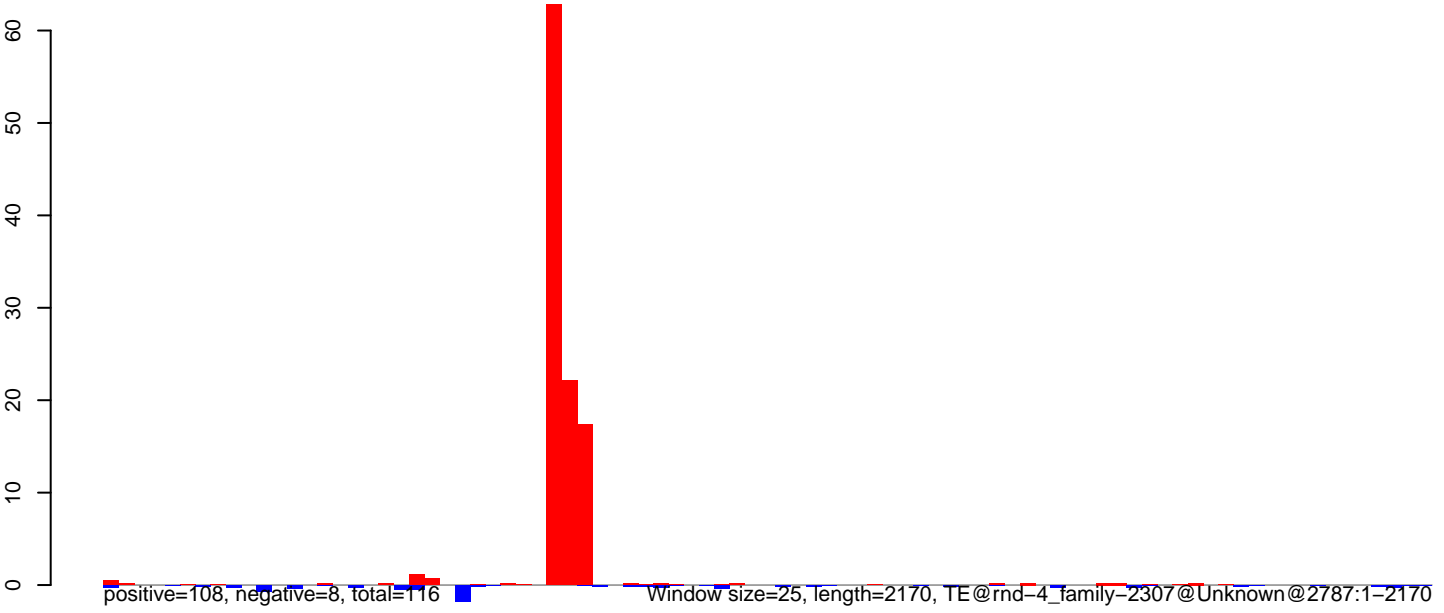
CuQuin_Testes_NL.18_23.rep



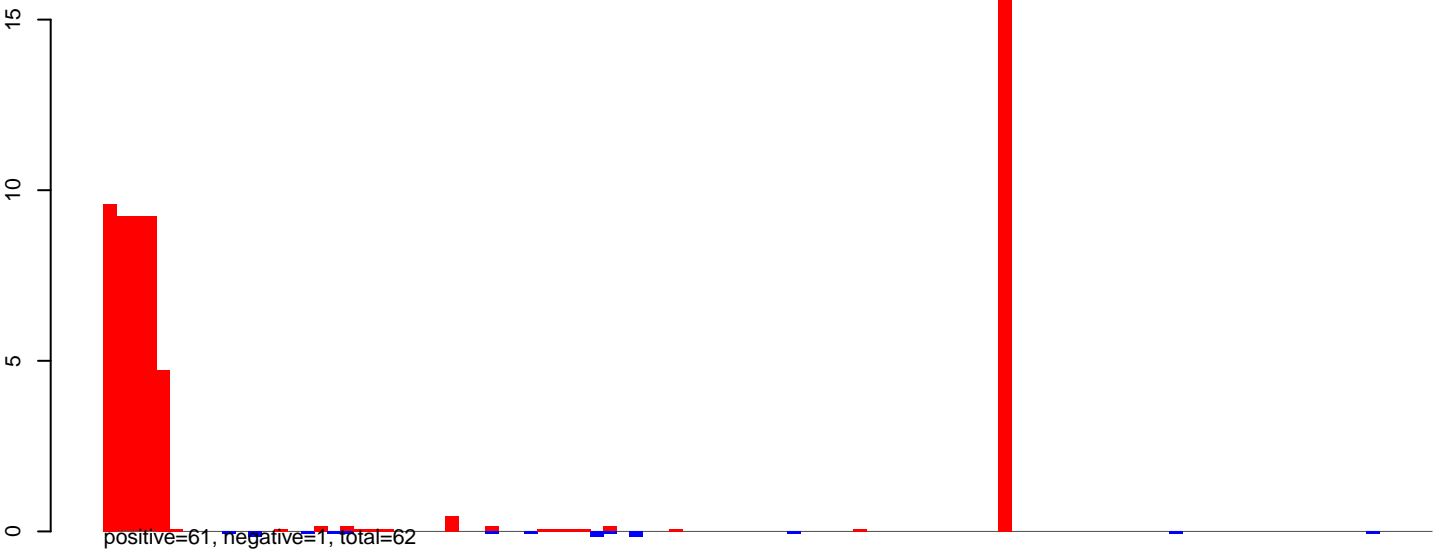
CuQuin_Testes_NL.24_35.rep



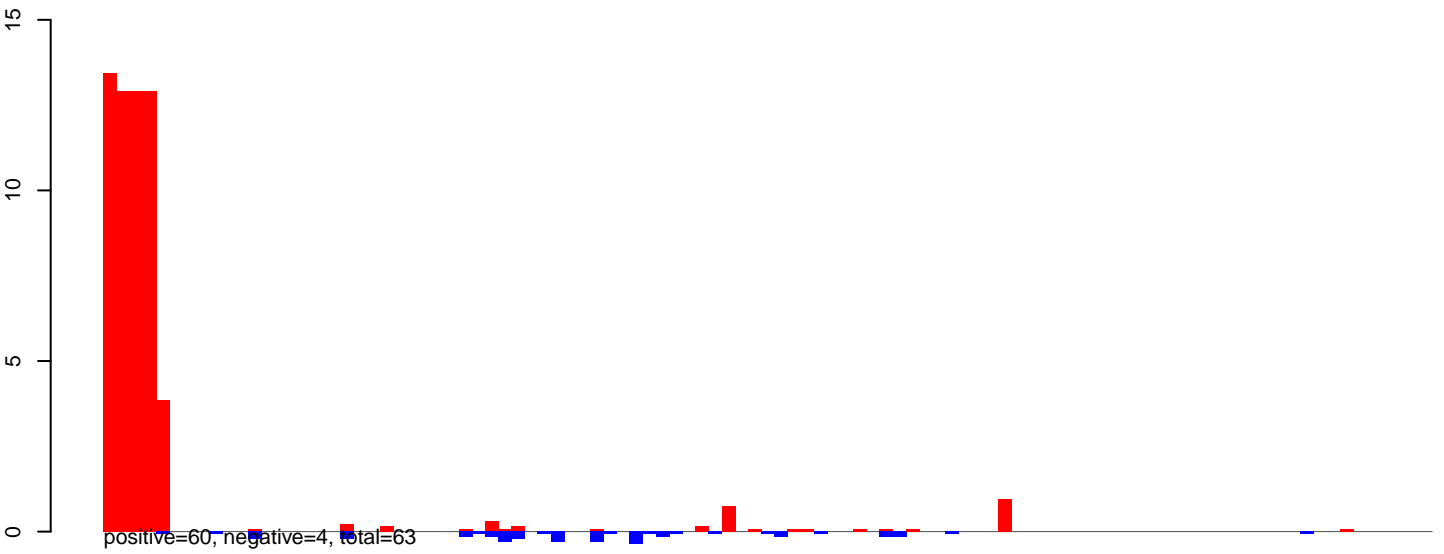
CuQuin_Testes_NL.rep



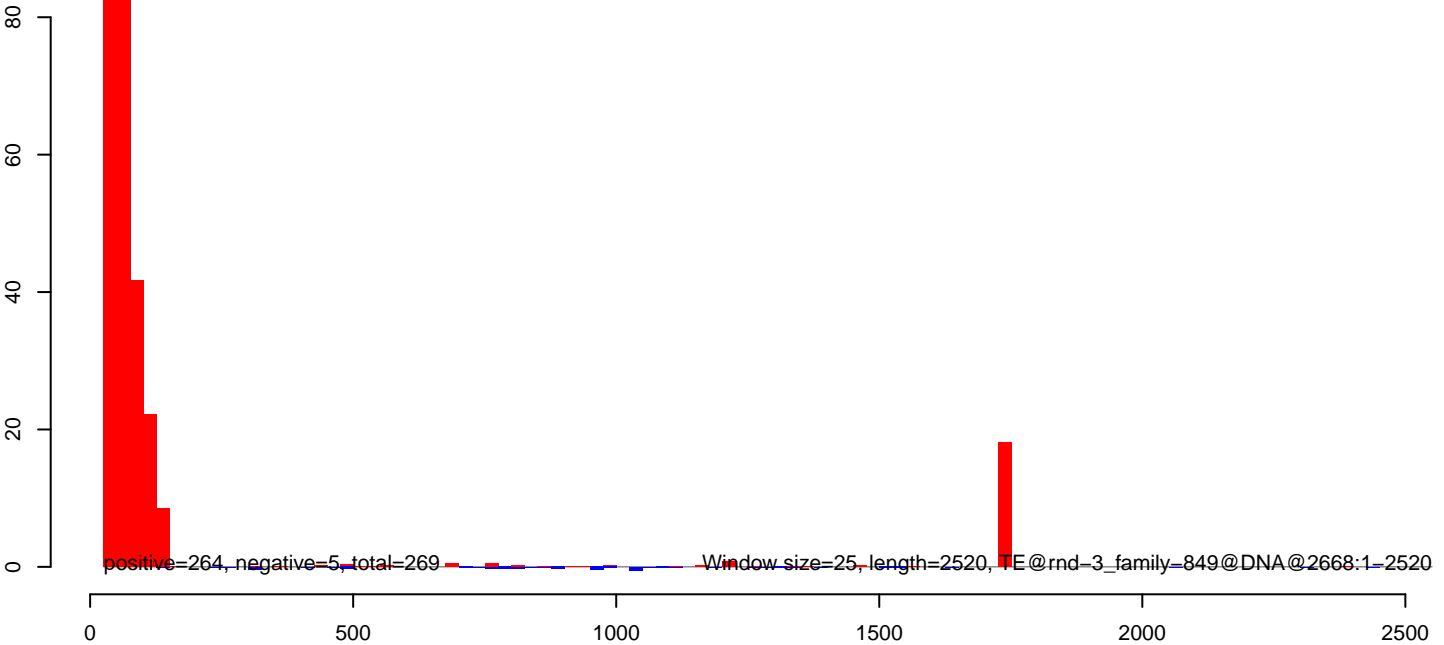
CuQuin_Testes_NL.18_23.rep



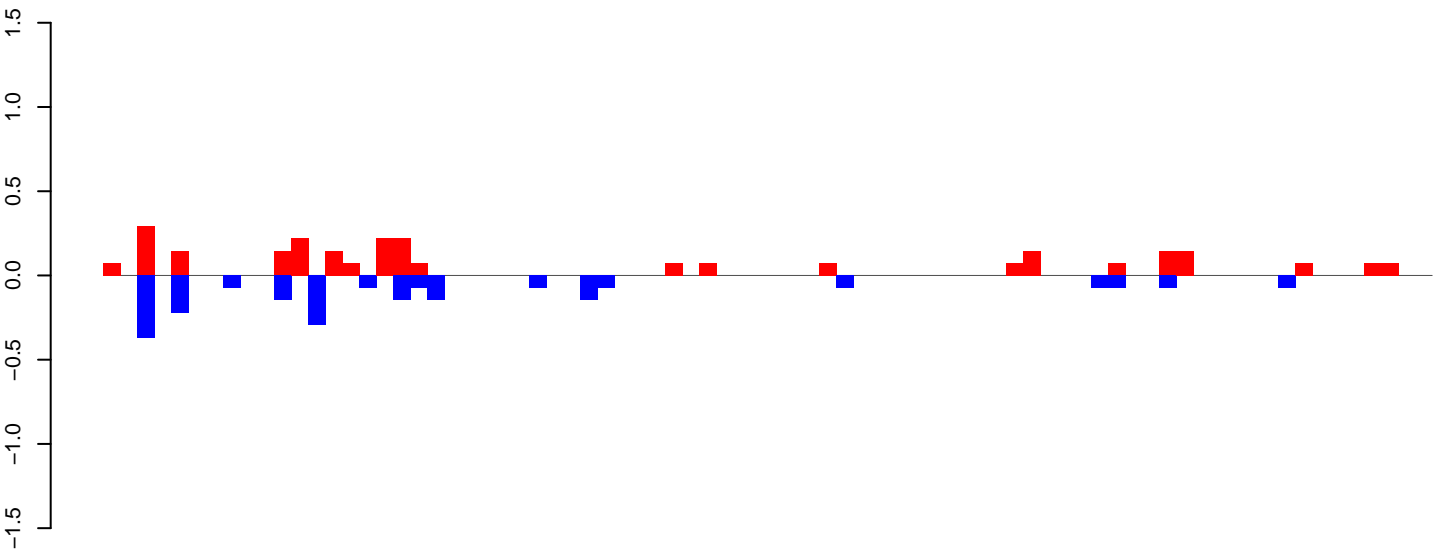
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

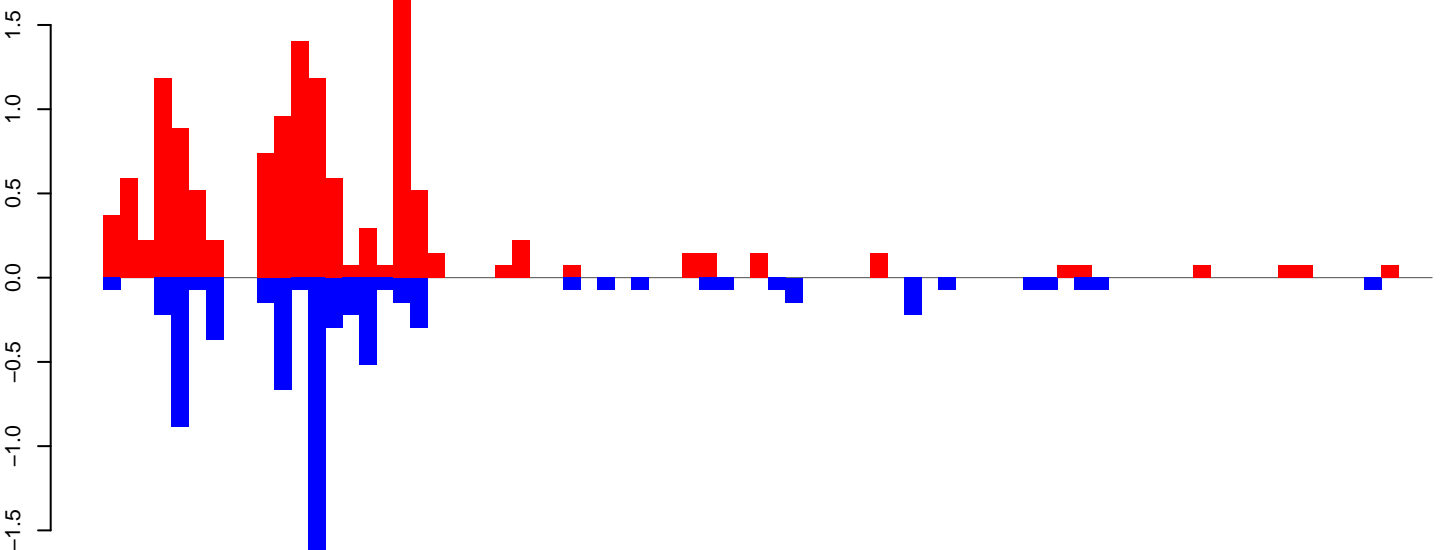


CuQuin_Testes_NL.18_23.rep



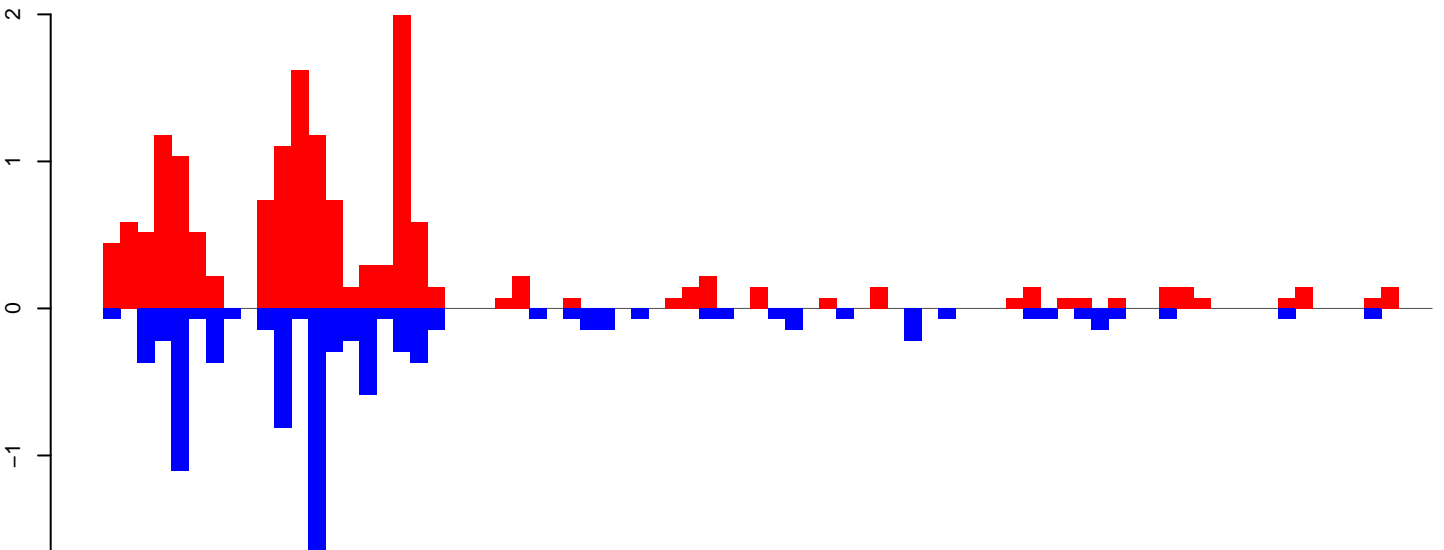
positive=3, negative=2, total=5

CuQuin_Testes_NL.24_35.rep



positive=13, negative=7, total=20

CuQuin_Testes_NL.rep

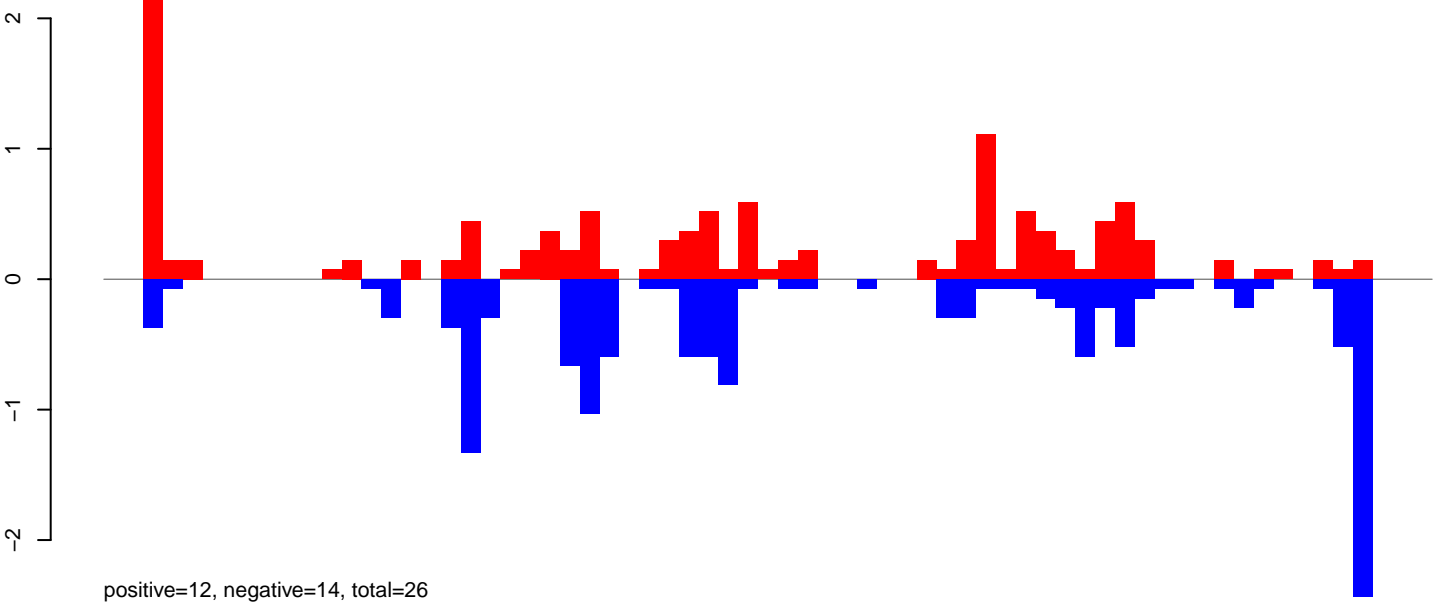


positive=16, negative=9, total=25

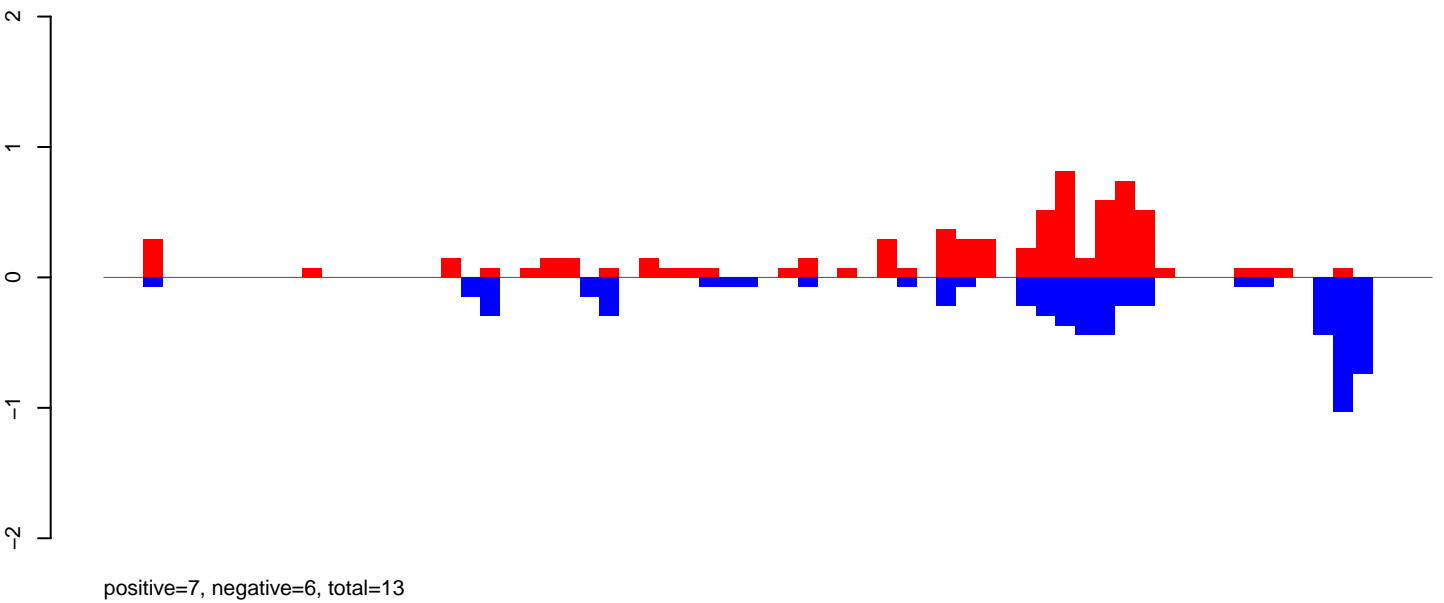
Window size=25, length=1936, TE@rnd-5_family-23@DNA@2315:1-1936

0 500 1000 1500 2000

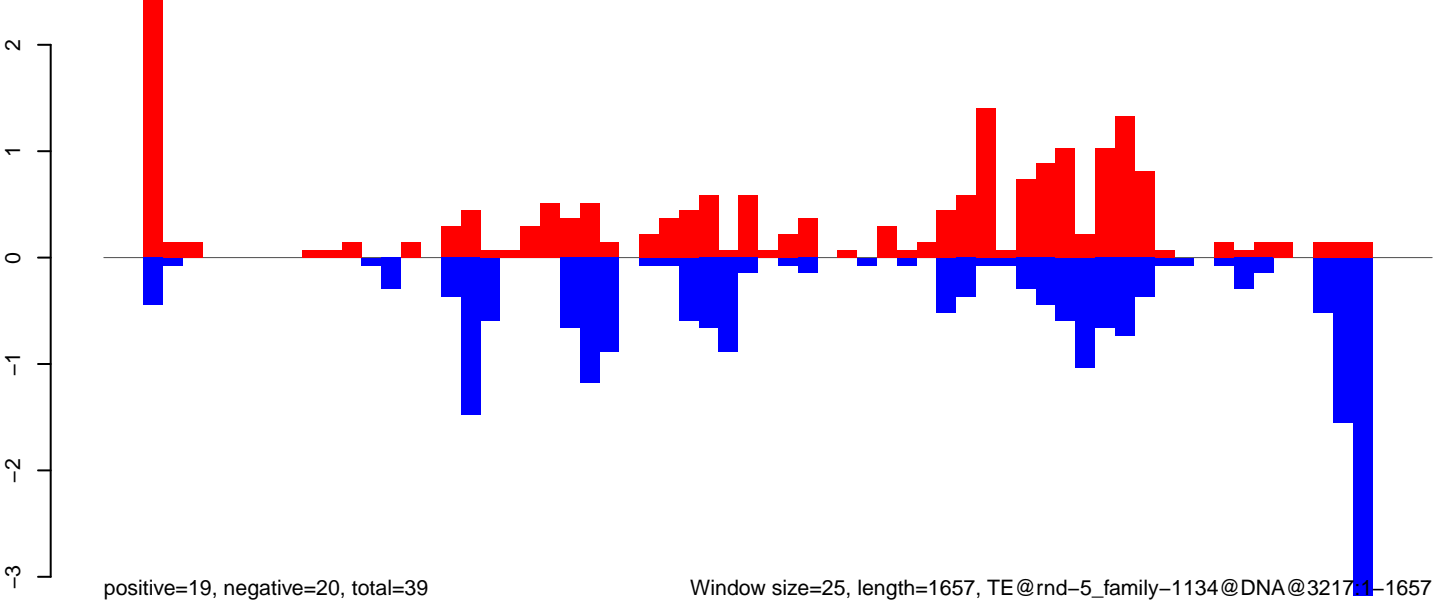
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



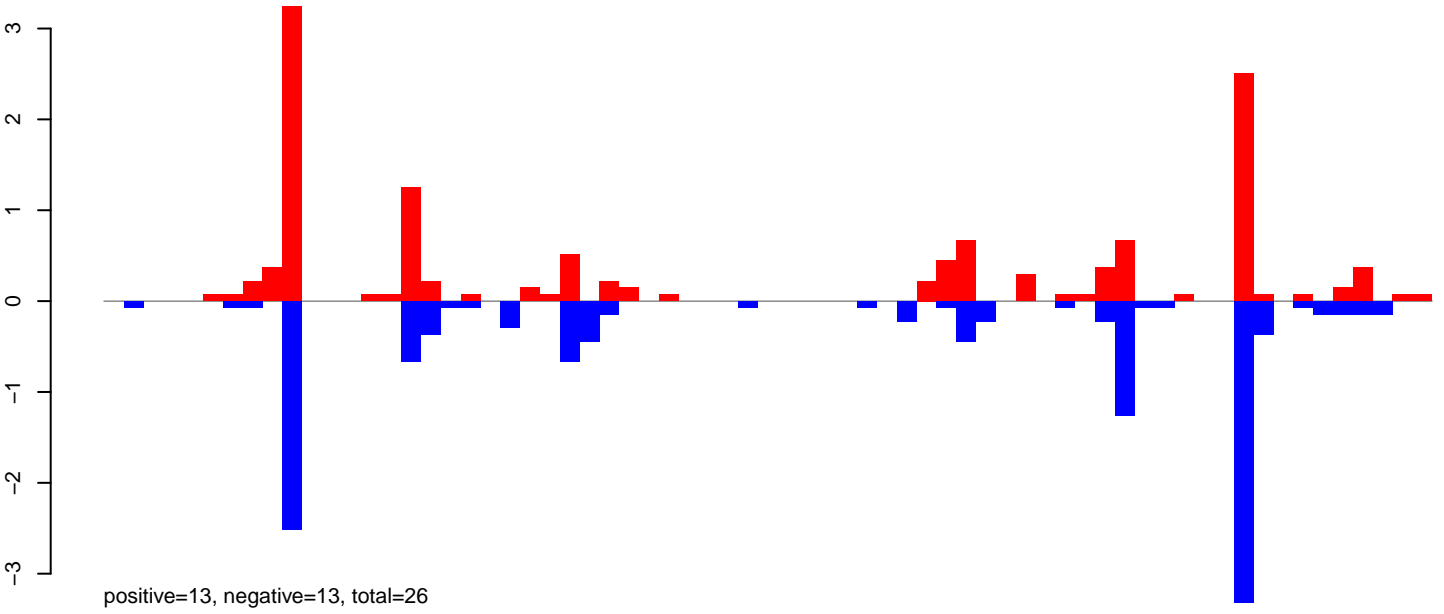
CuQuin_Testes_NL.rep



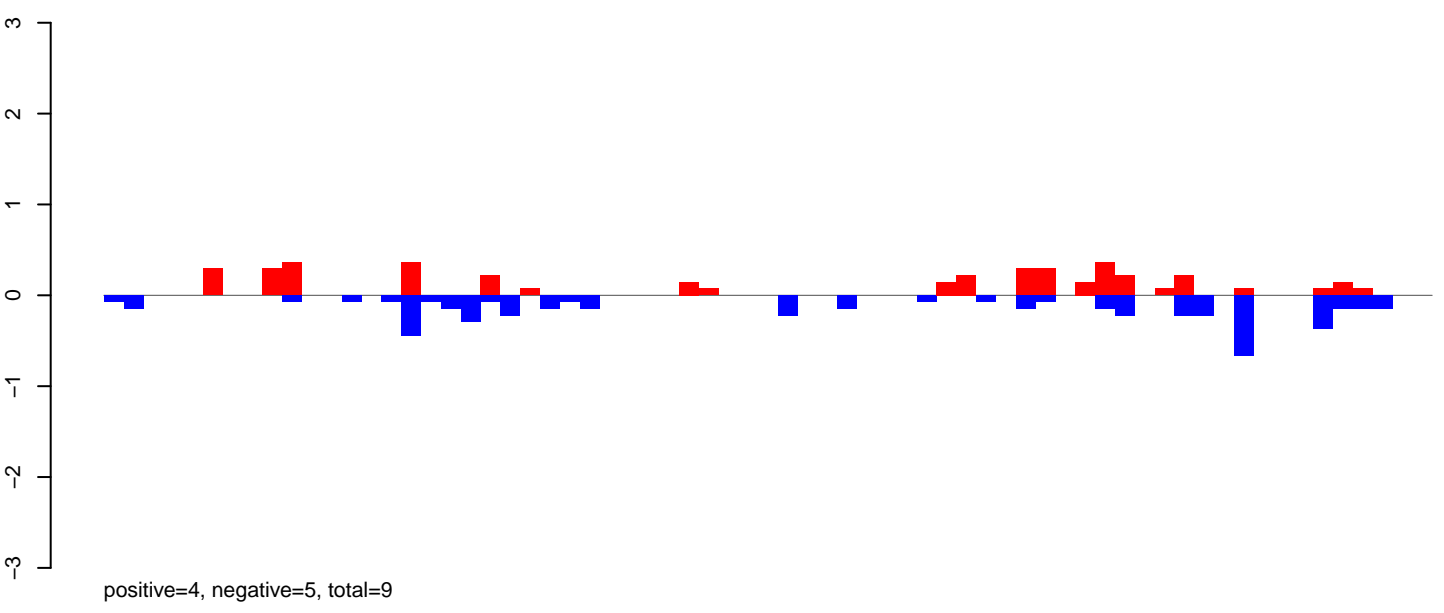
Window size=25, length=1657, TE@rnd-5_family-1134@DNA@321741-1657

0 500 1000 1500

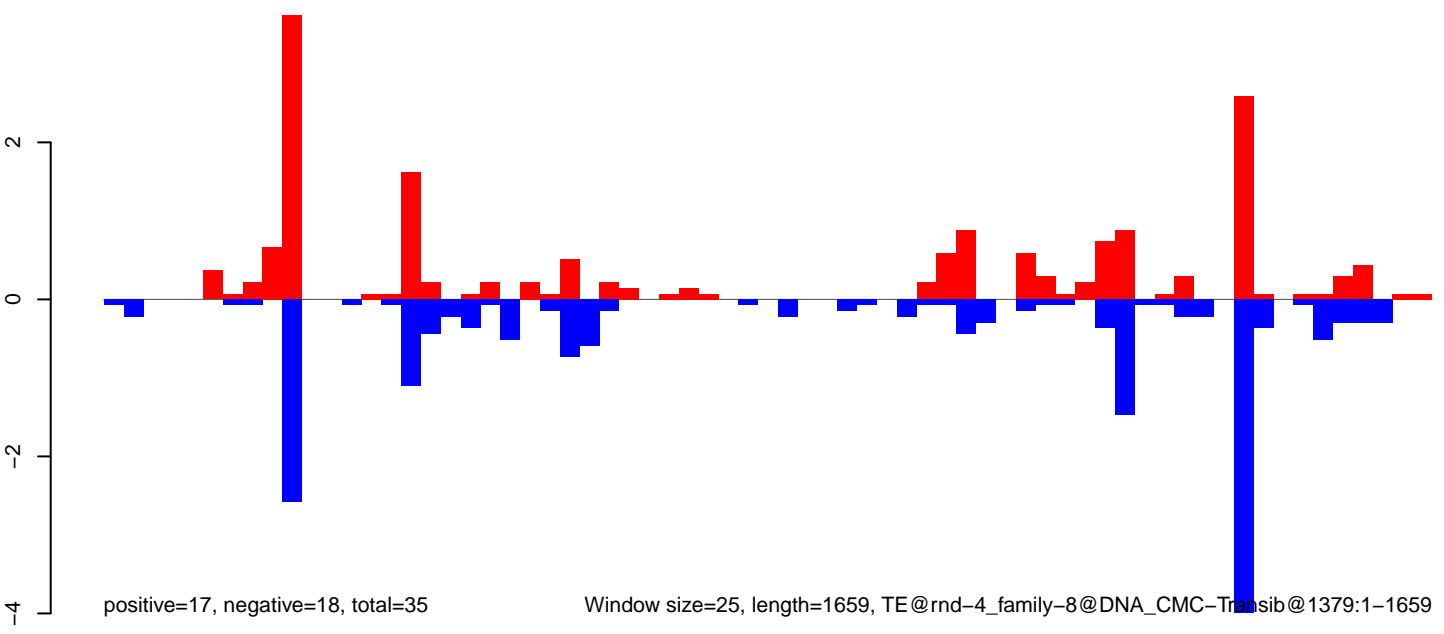
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

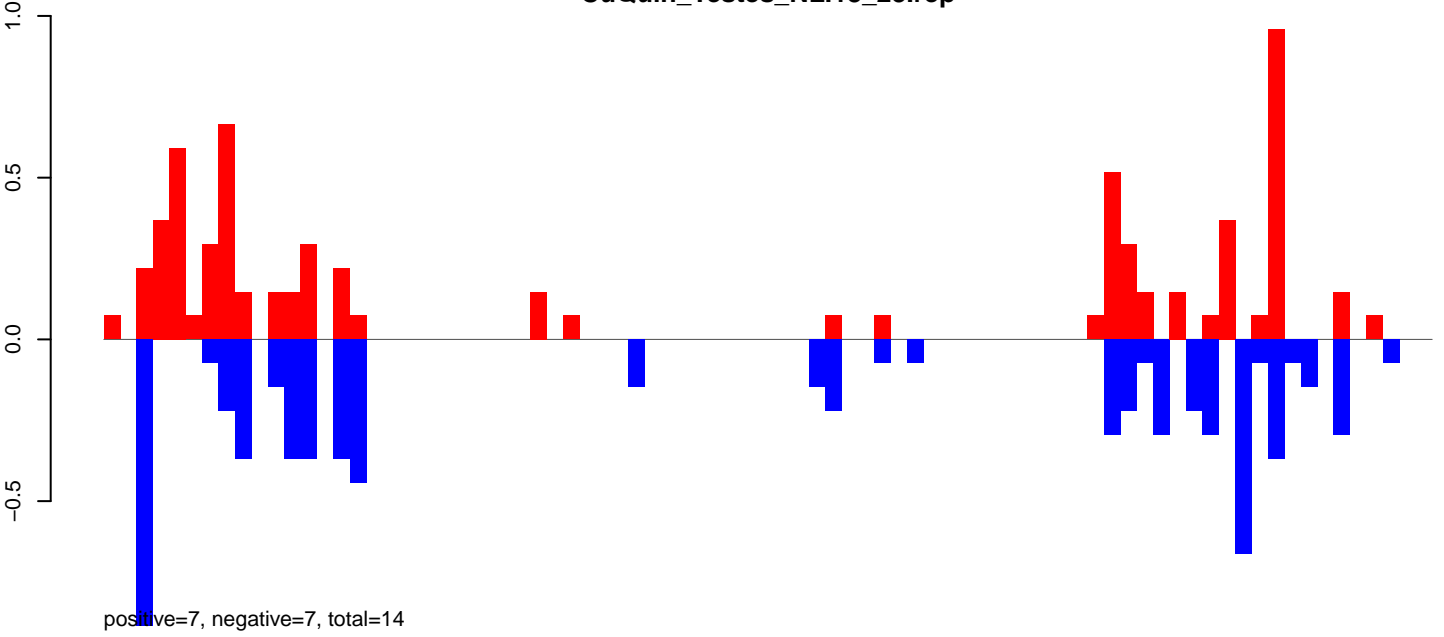


CuQuin_Testes_NL.rep

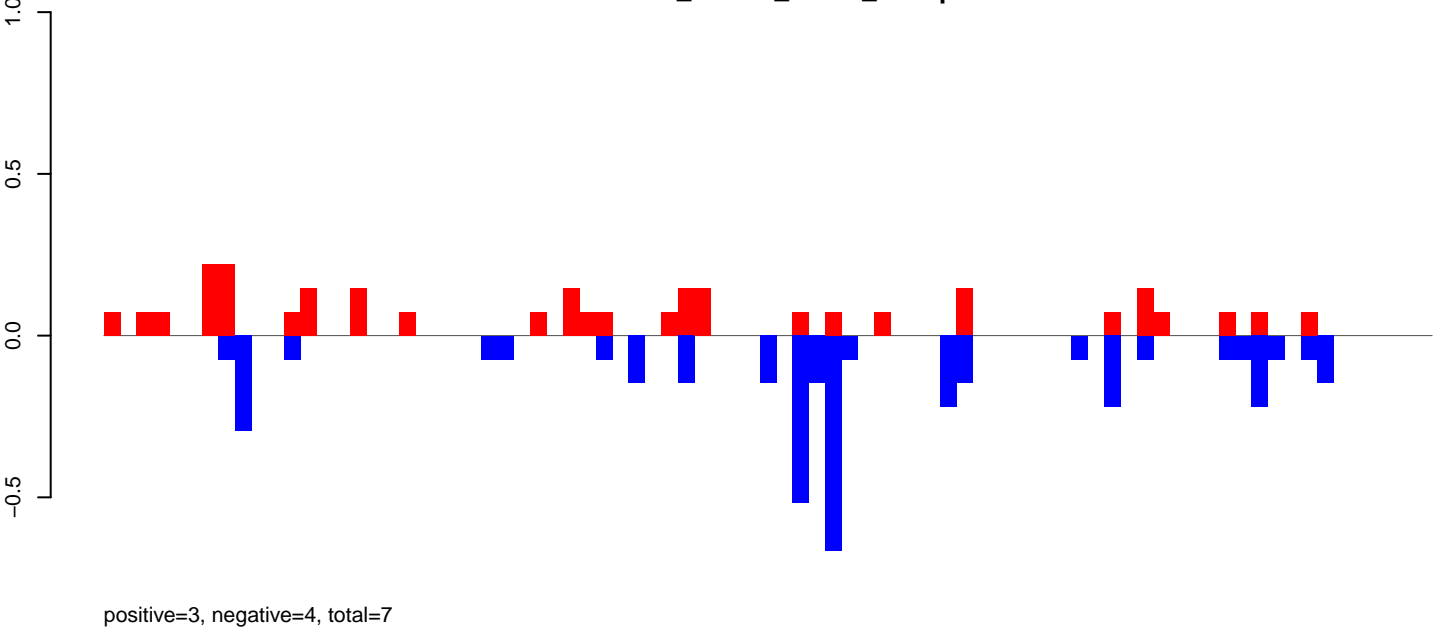


Window size=25, length=1659, TE@rnd-4_family-8@DNA_CMC-Transib@1379:1-1659

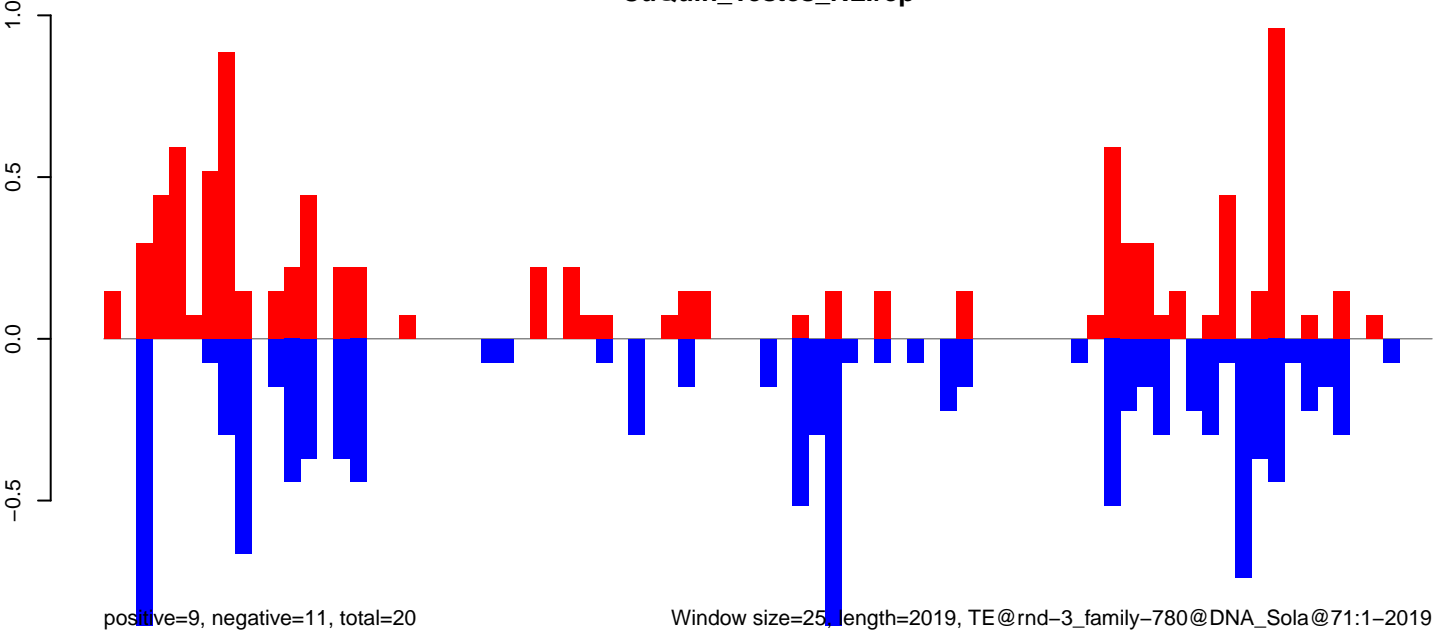
CuQuin_Testes_NL.18_23.rep



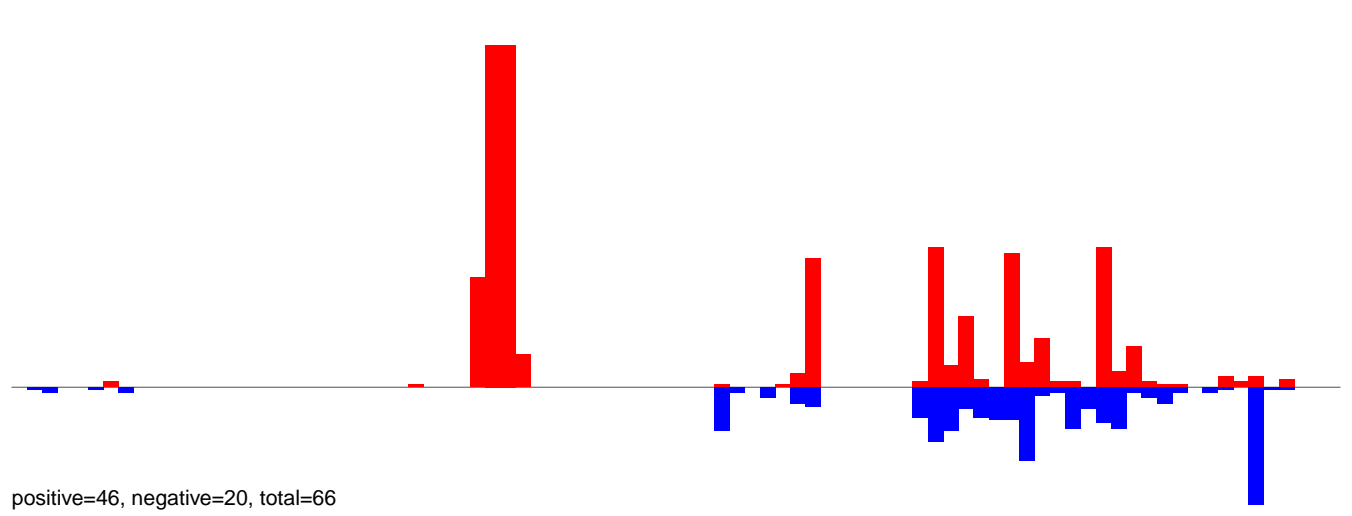
CuQuin_Testes_NL.24_35.rep



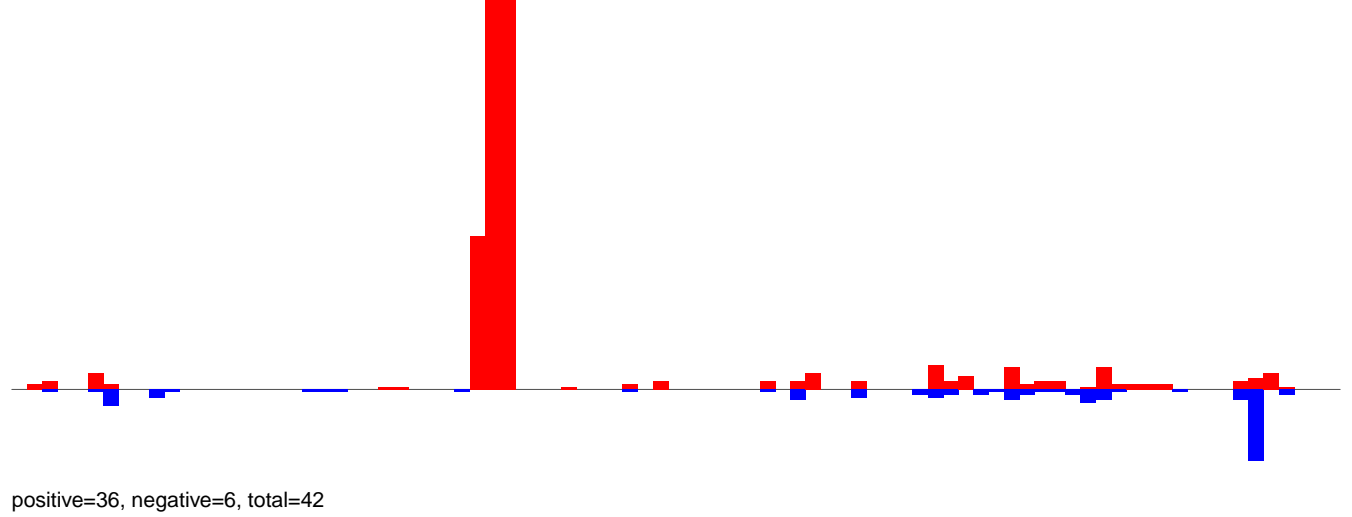
CuQuin_Testes_NL.rep



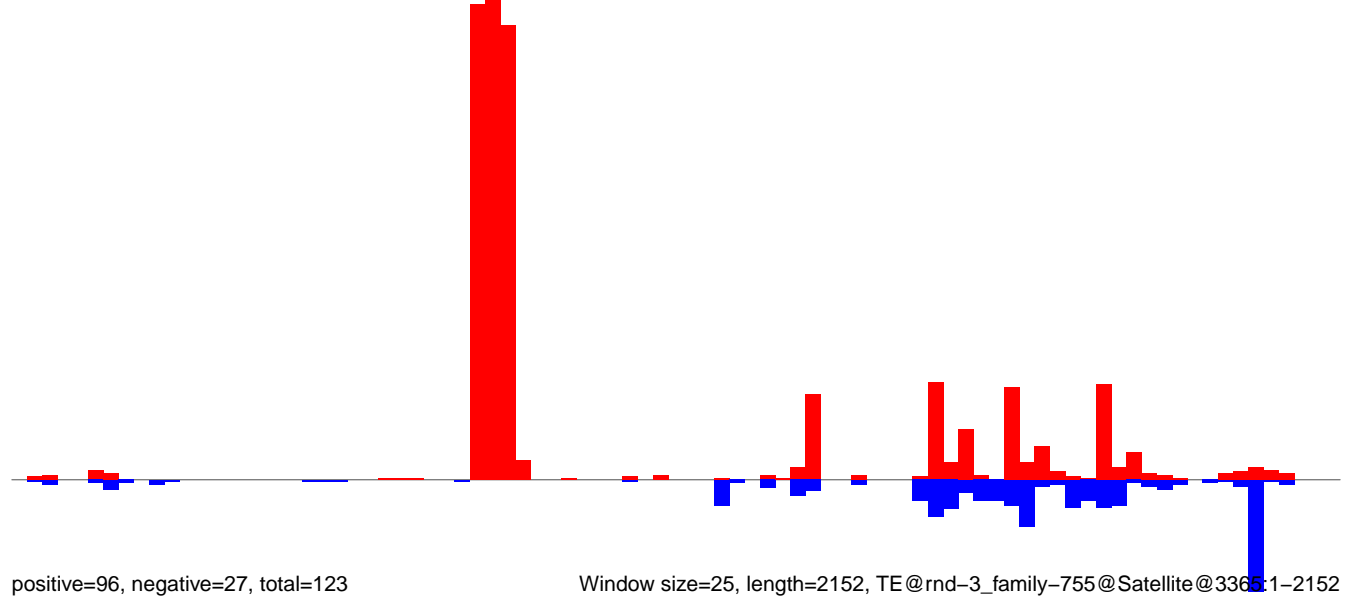
CuQuin_Testes_NL.18_23.rep



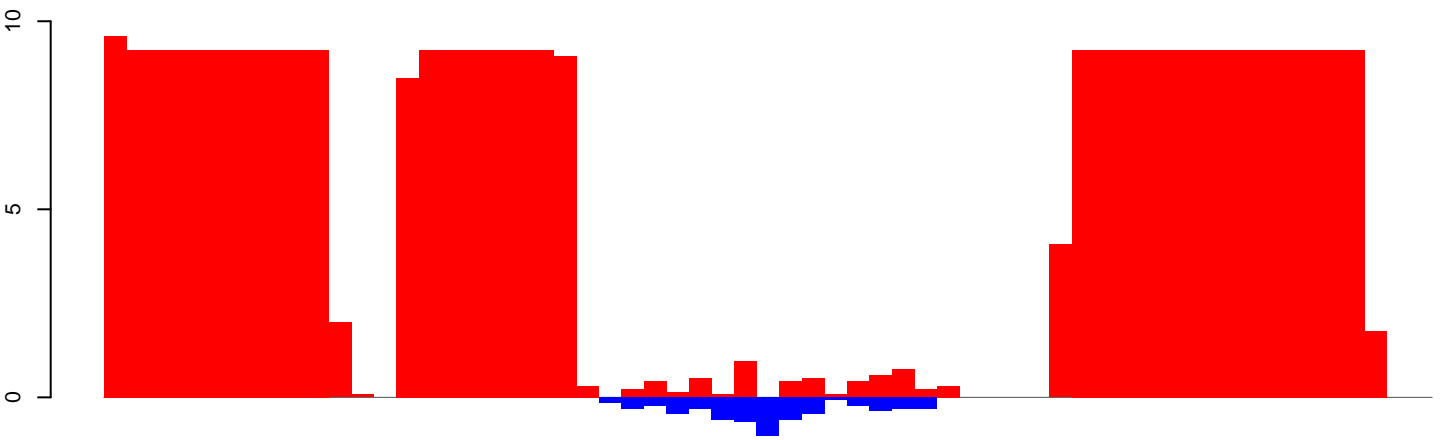
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

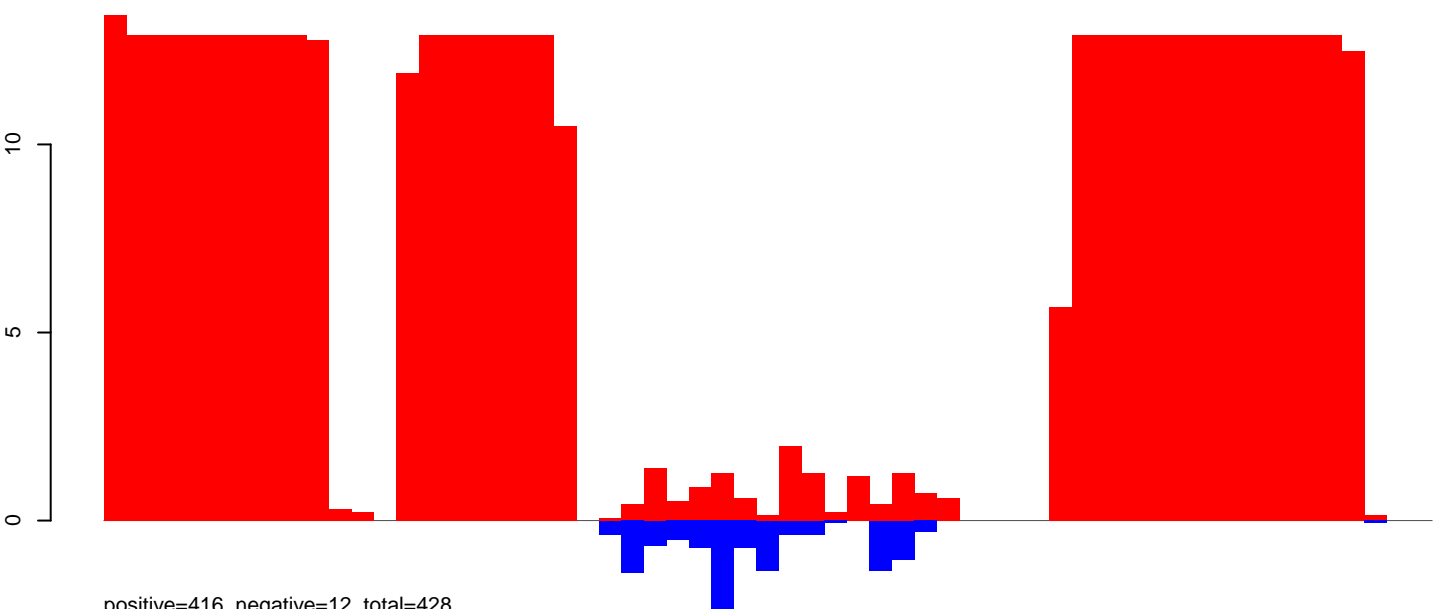


CuQuin_Testes_NL.18_23.rep



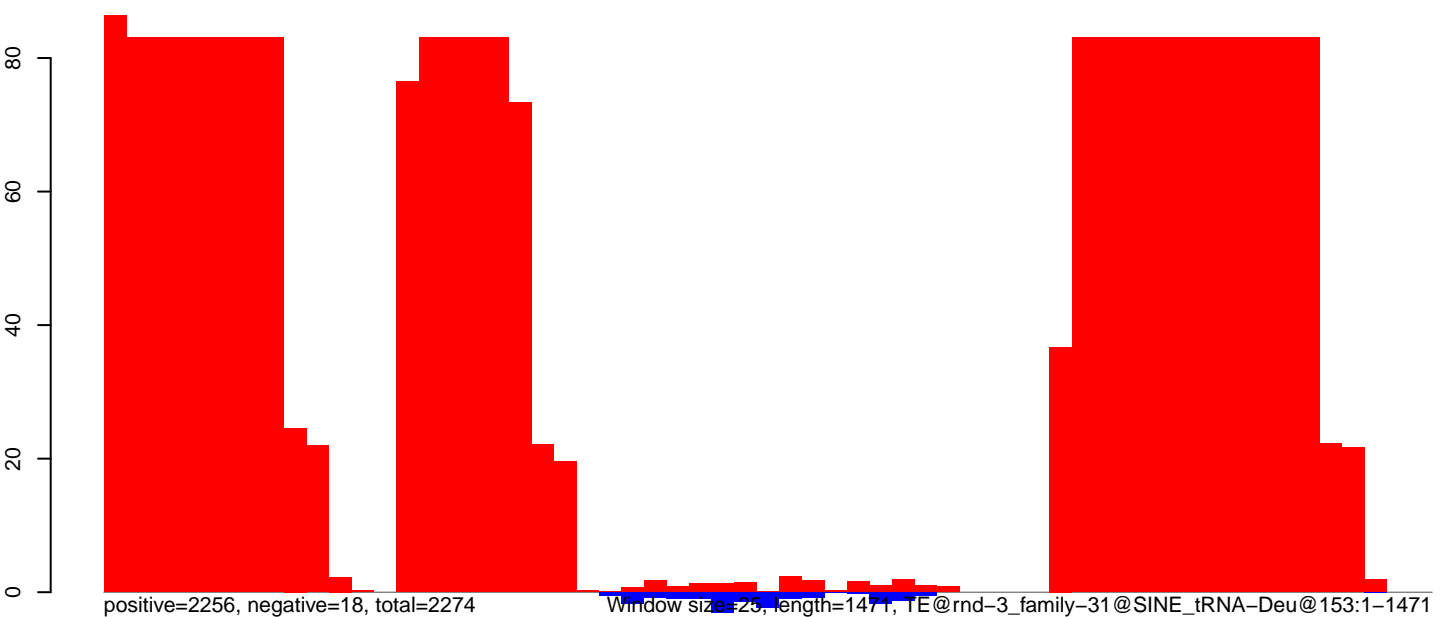
positive=299, negative=6, total=305

CuQuin_Testes_NL.24_35.rep



positive=416, negative=12, total=428

CuQuin_Testes_NL.rep

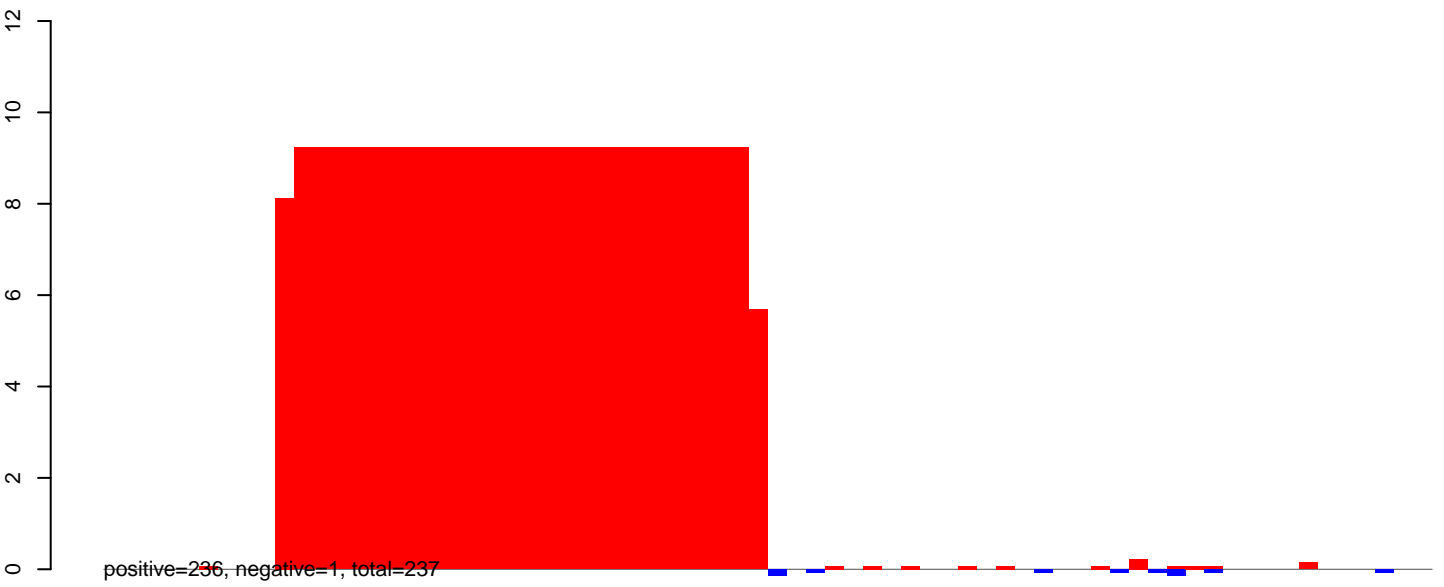


positive=2256, negative=18, total=2274

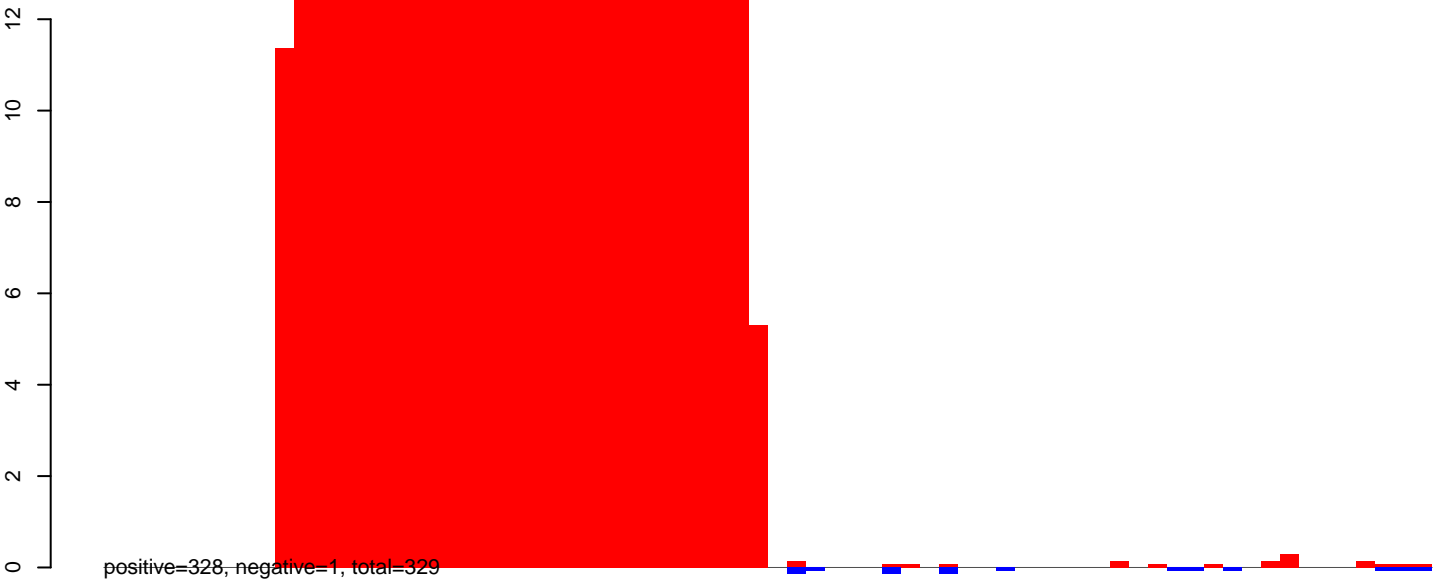
Window size=25, length=1471, TE@rnd-3_family-31@SINE_tRNA-Deu@153:1-1471

0 500 1000 1500

CuQuin_Testes_NL.18_23.rep



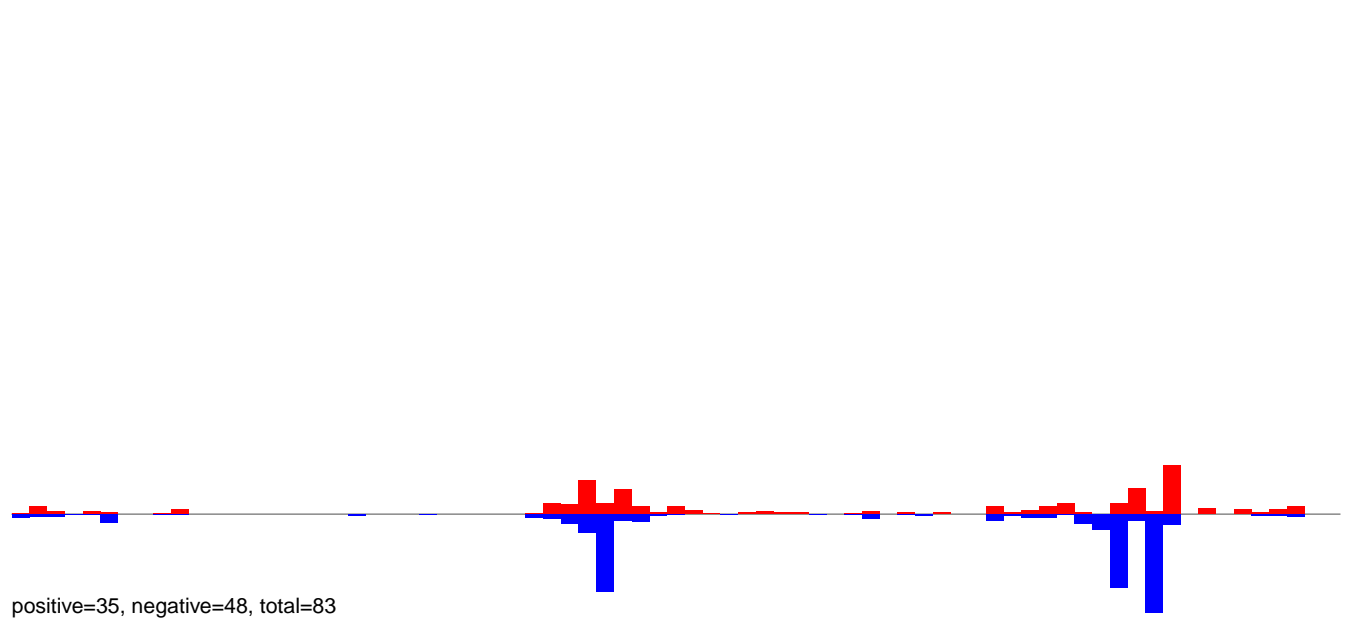
CuQuin_Testes_NL.24_35.rep



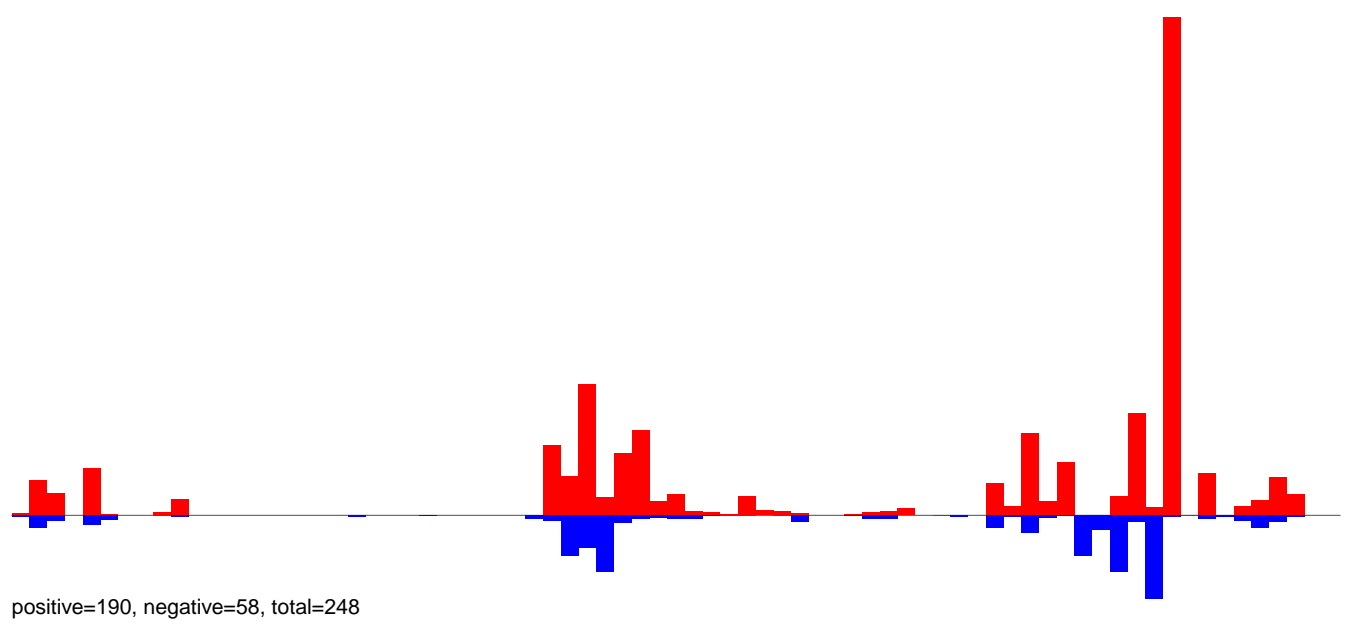
CuQuin_Testes_NL.rep



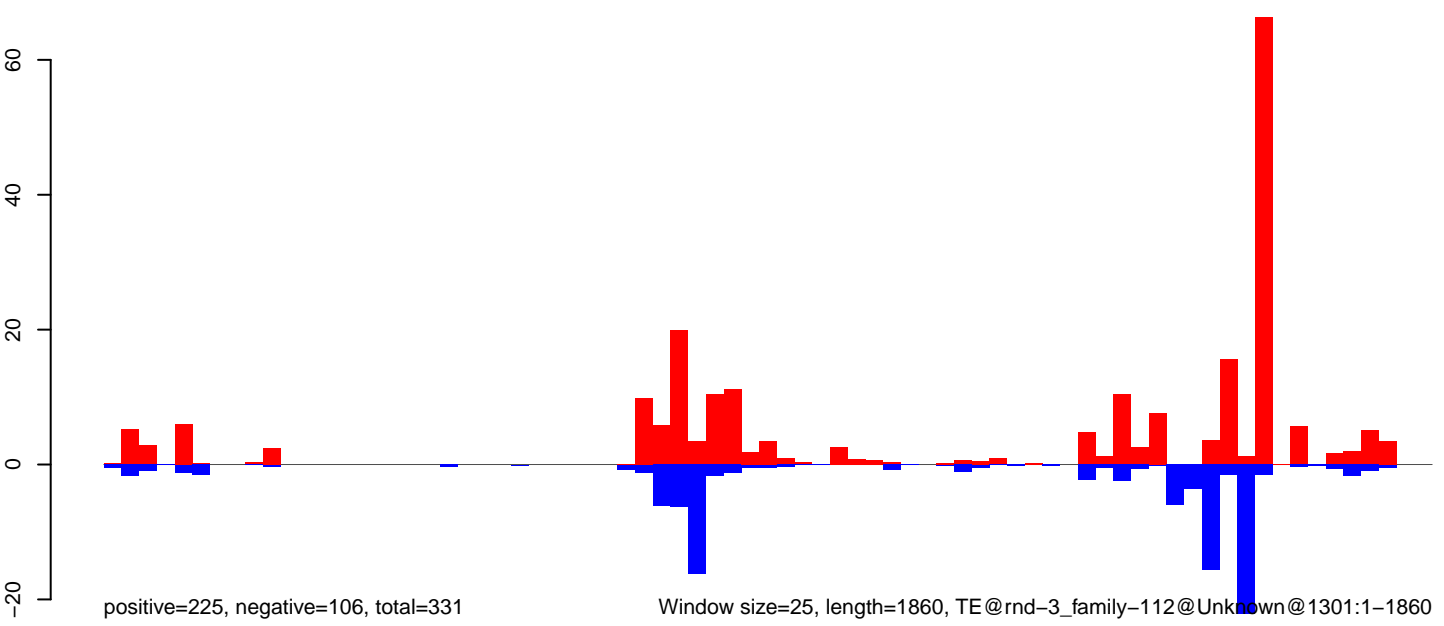
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



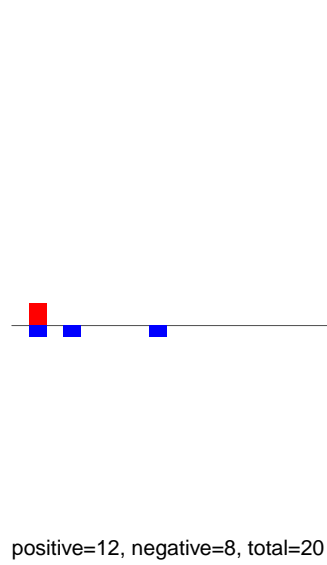
CuQuin_Testes_NL.rep



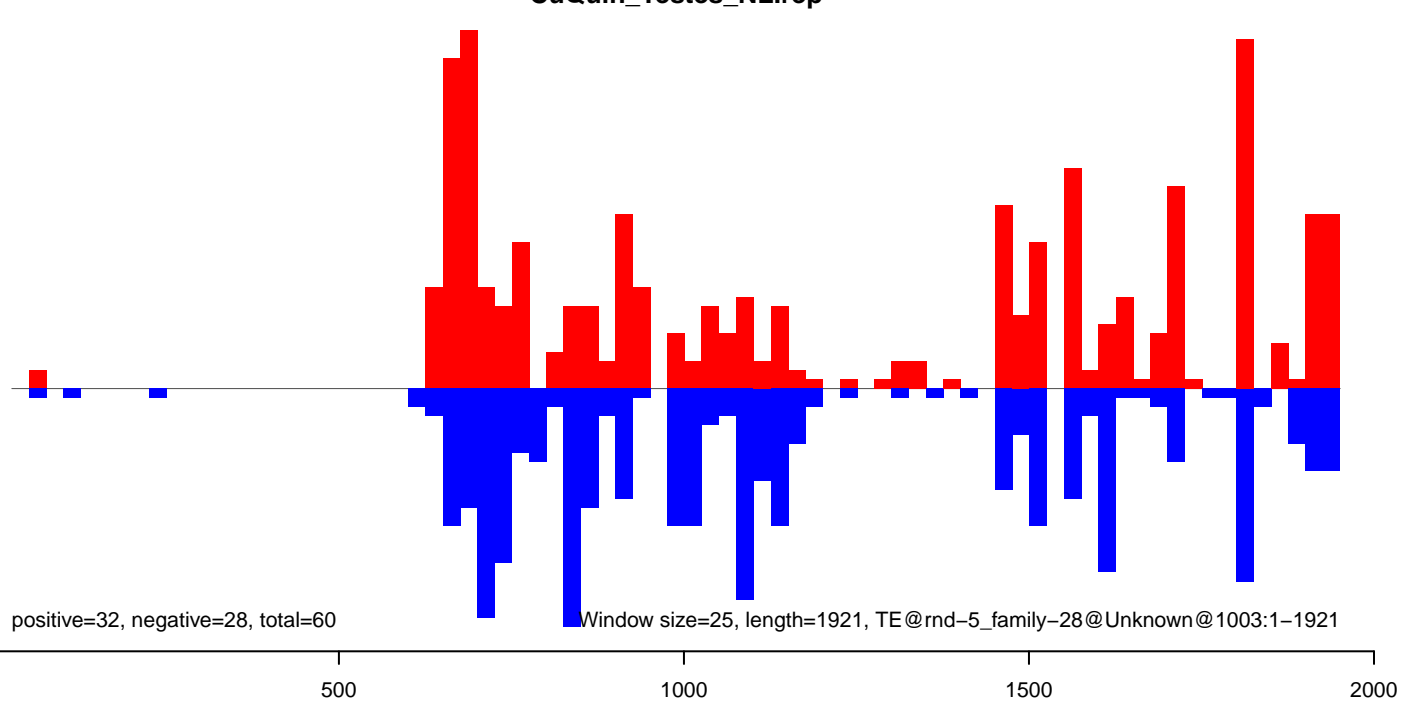
CuQuin_Testes_NL.18_23.rep



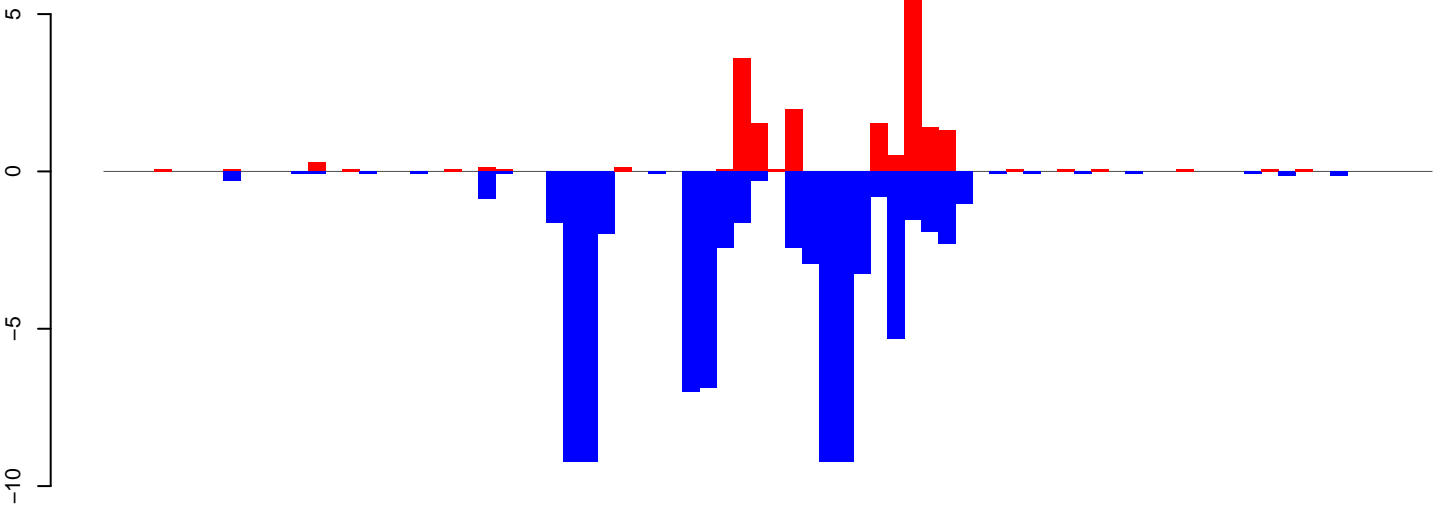
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

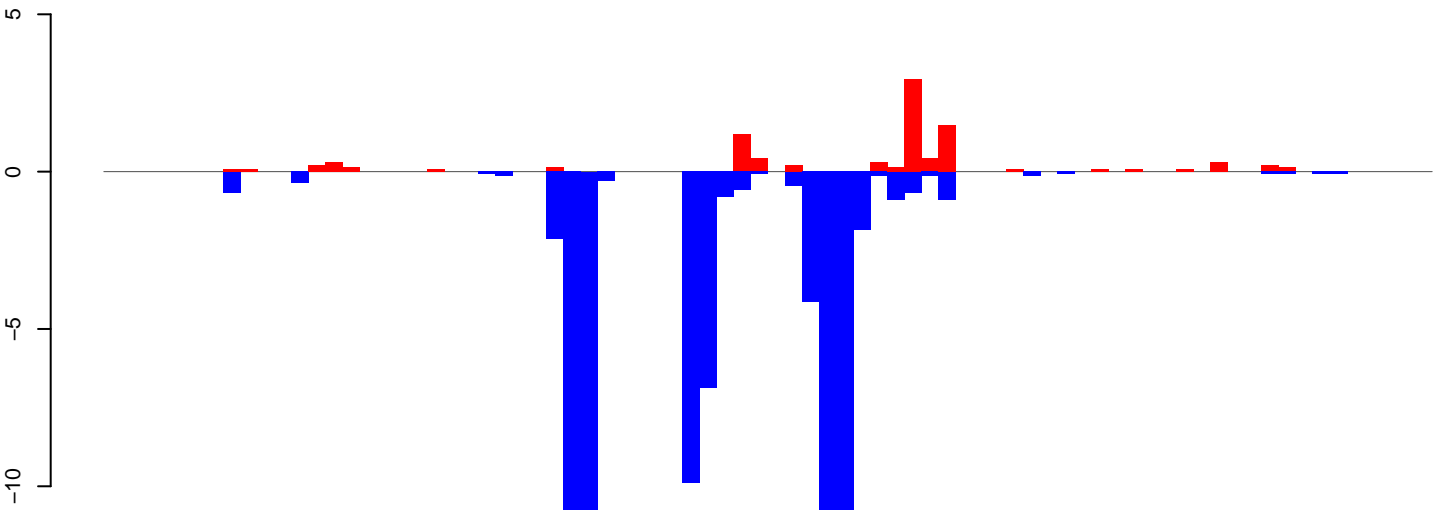


CuQuin_Testes_NL.18_23.rep



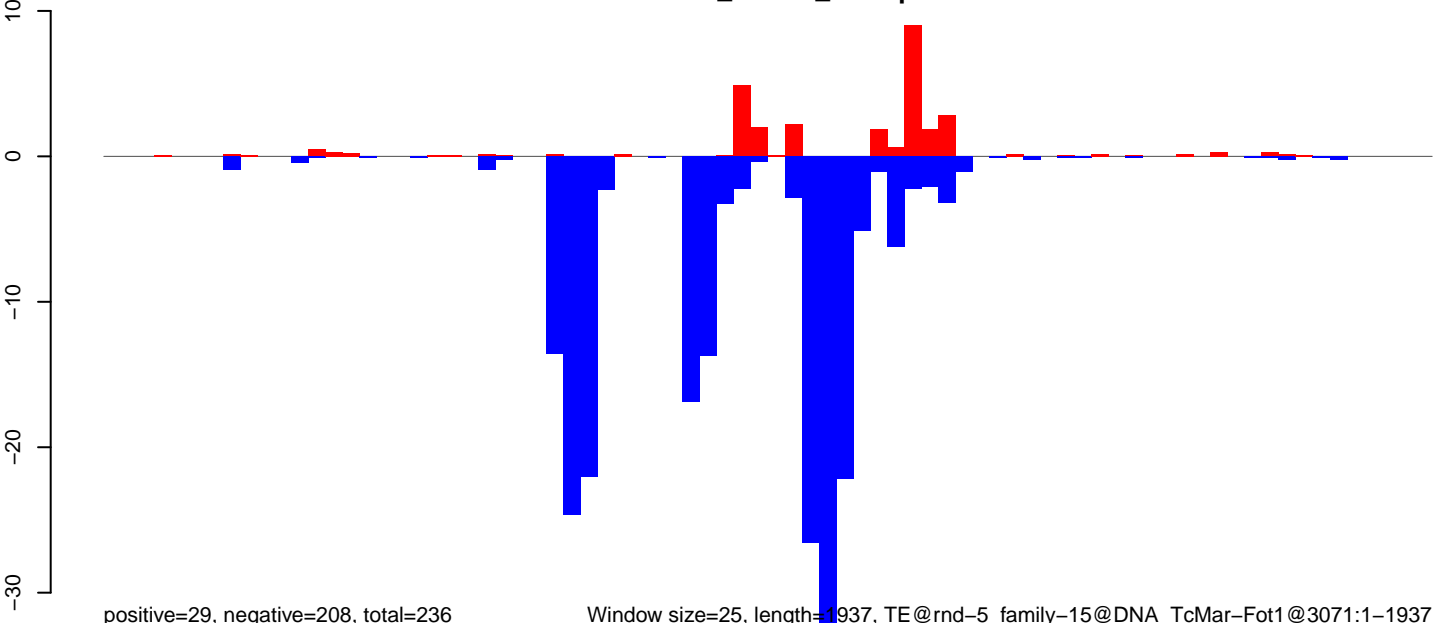
positive=20, negative=83, total=102

CuQuin_Testes_NL.24_35.rep



positive=9, negative=83, total=92

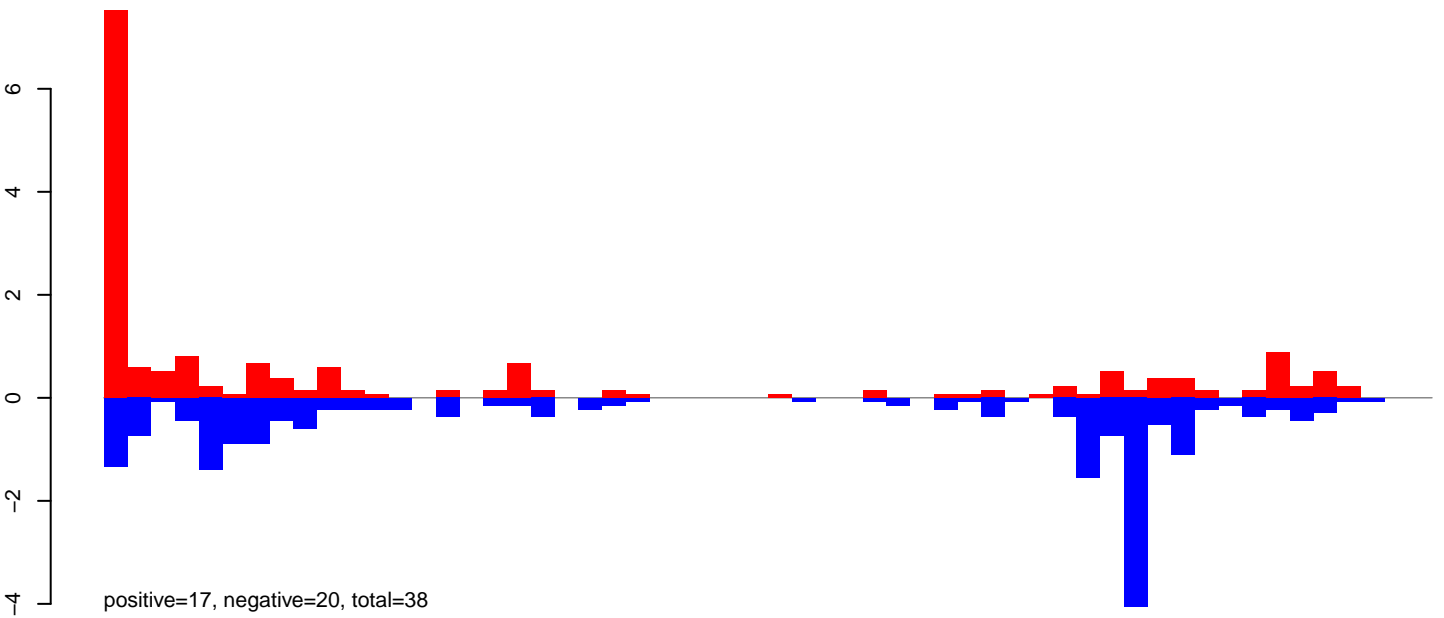
CuQuin_Testes_NL.rep



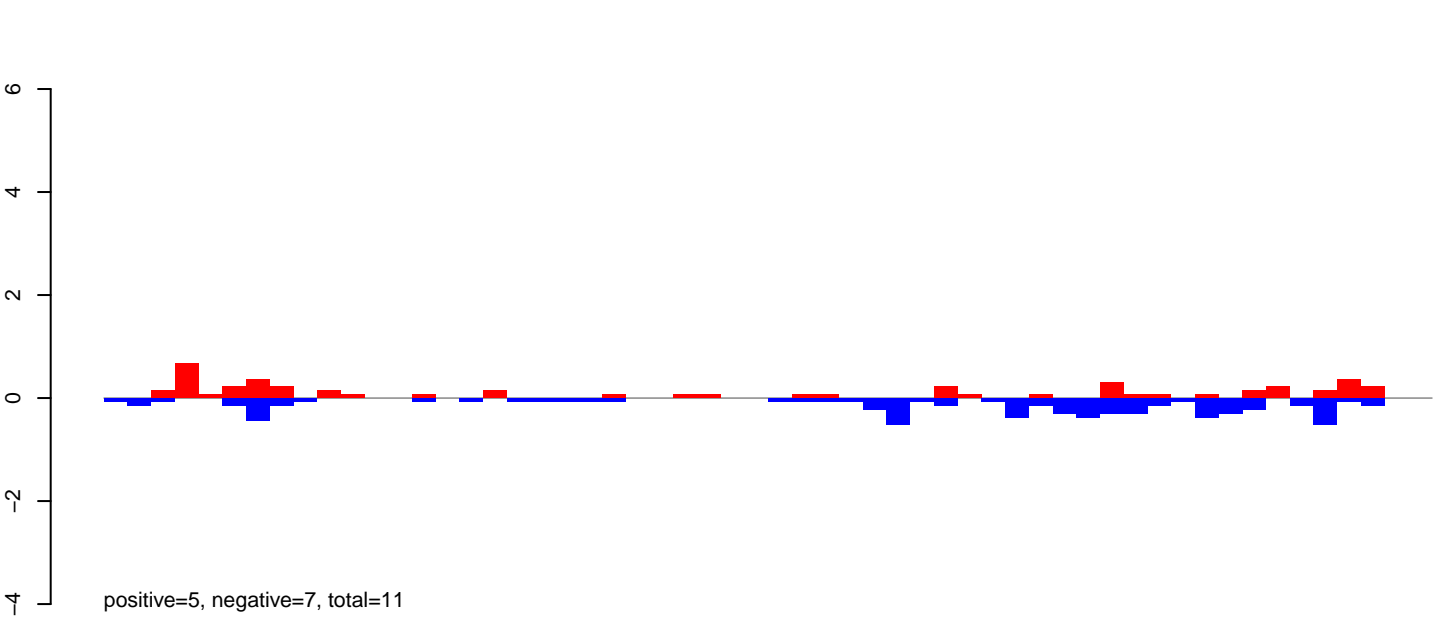
positive=29, negative=208, total=236

Window size=25, length=1937, TE@rnd-5_family-15@DNA_TcMar-Fot1@3071:1-1937

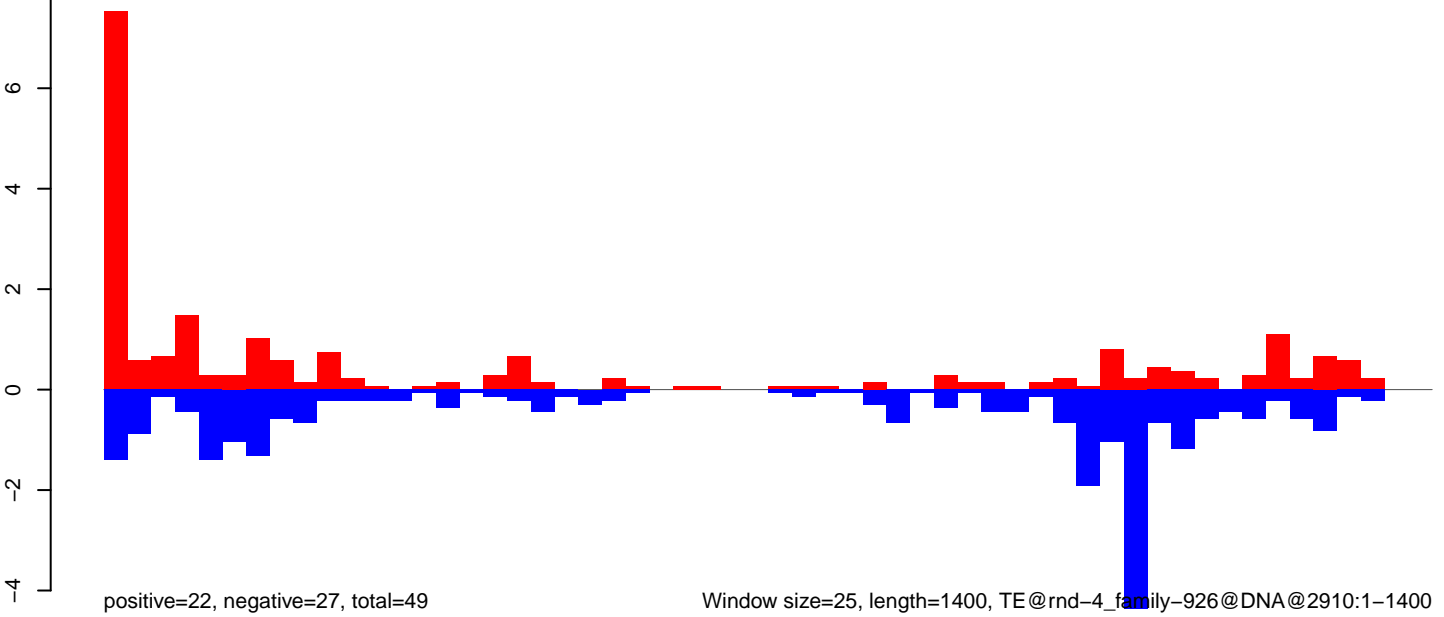
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

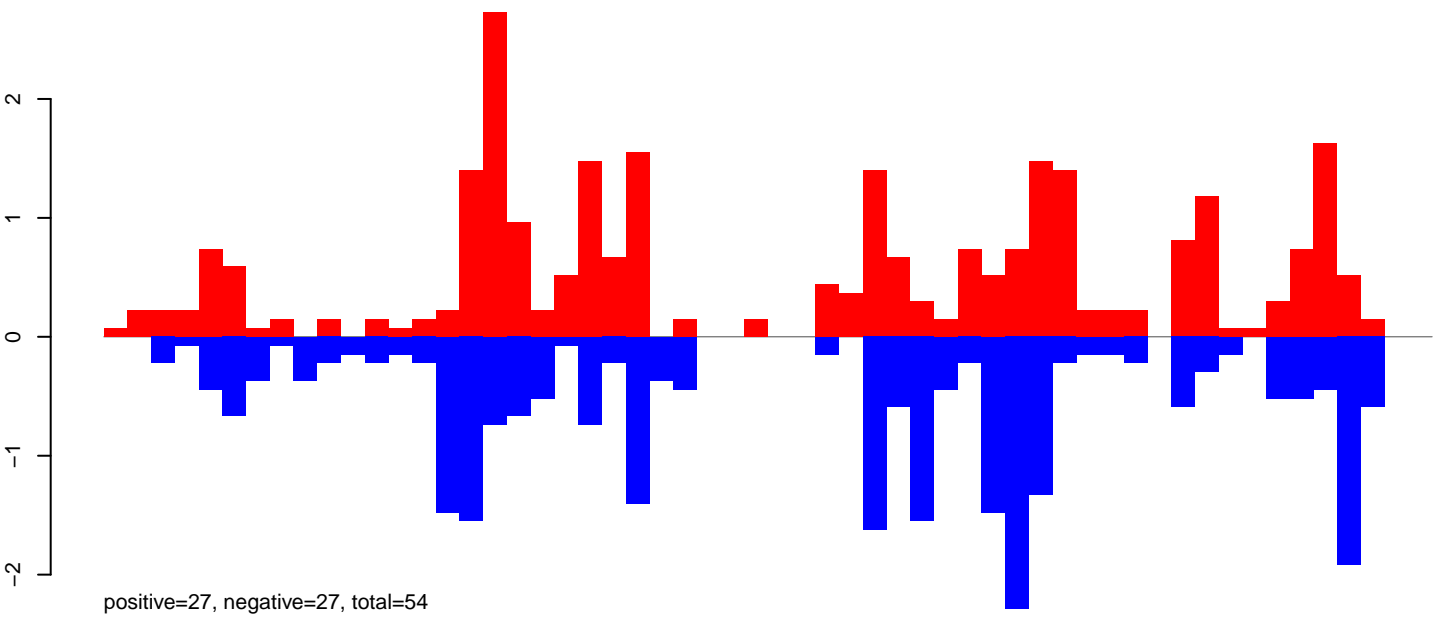


CuQuin_Testes_NL.rep

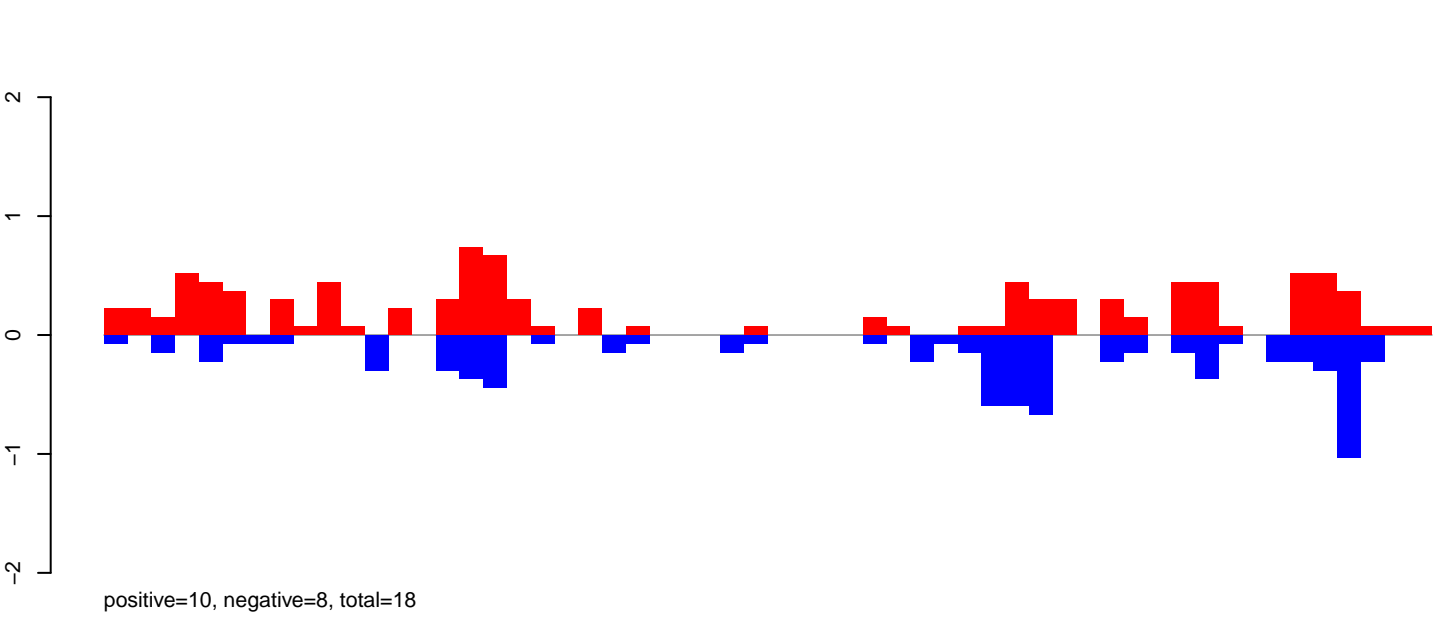


Window size=25, length=1400, TE@rnd-4_family-926@DNA@2910:1-1400

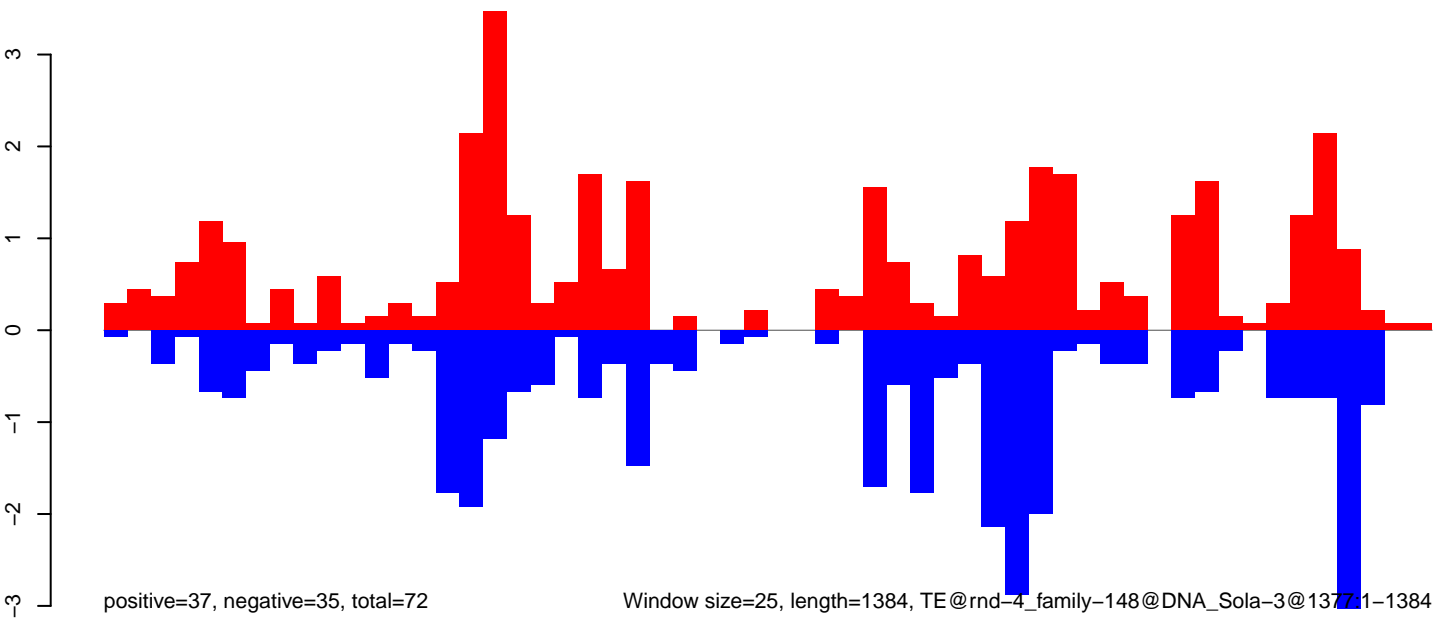
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

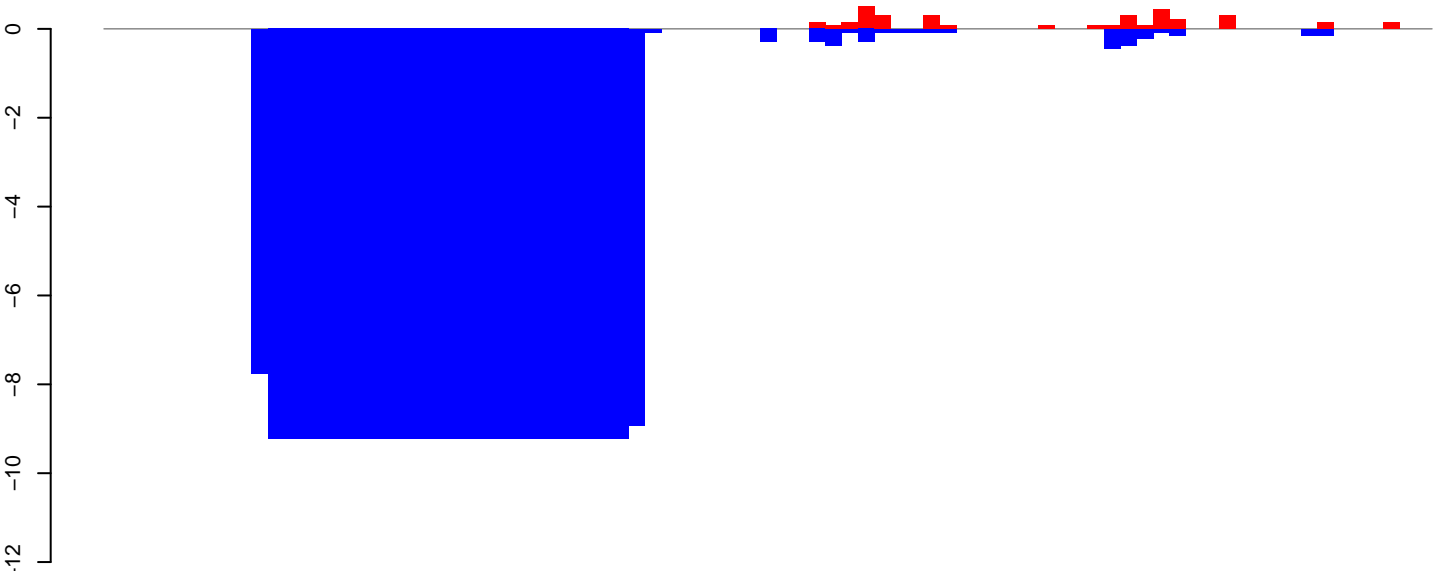


CuQuin_Testes_NL.rep



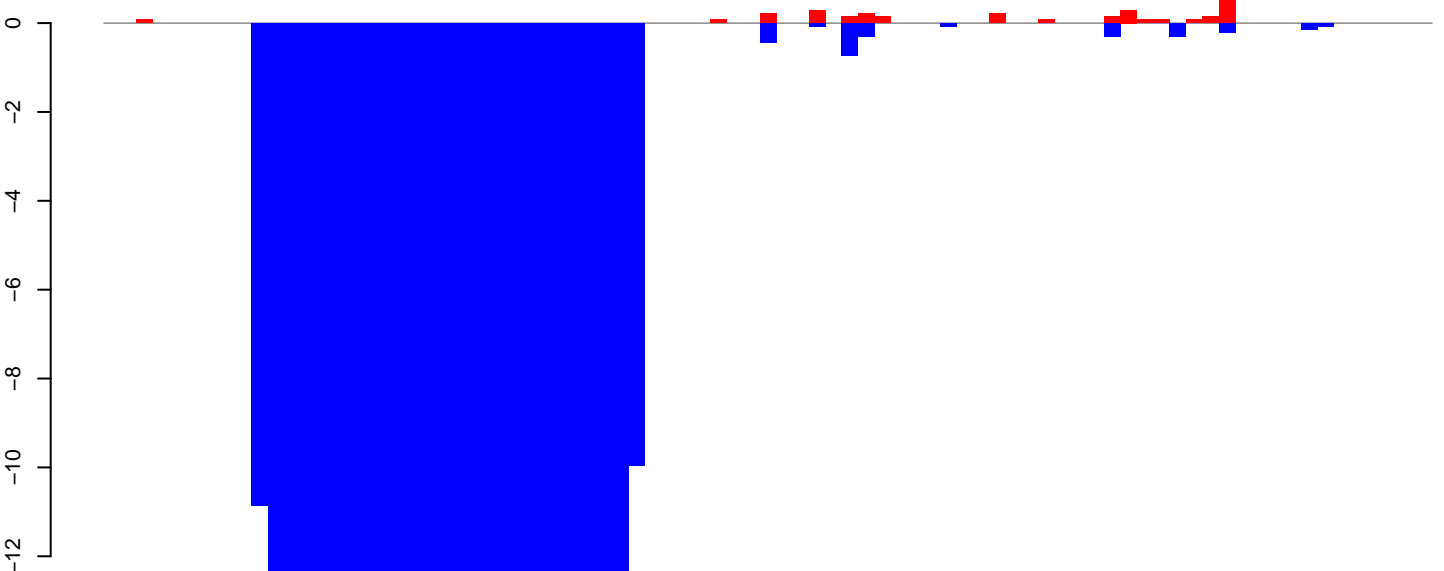
Window size=25, length=1384, TE@rnd-4_family-148@DNA_Sola-3@1377:1-1384

CuQuin_Testes_NL.18_23.rep



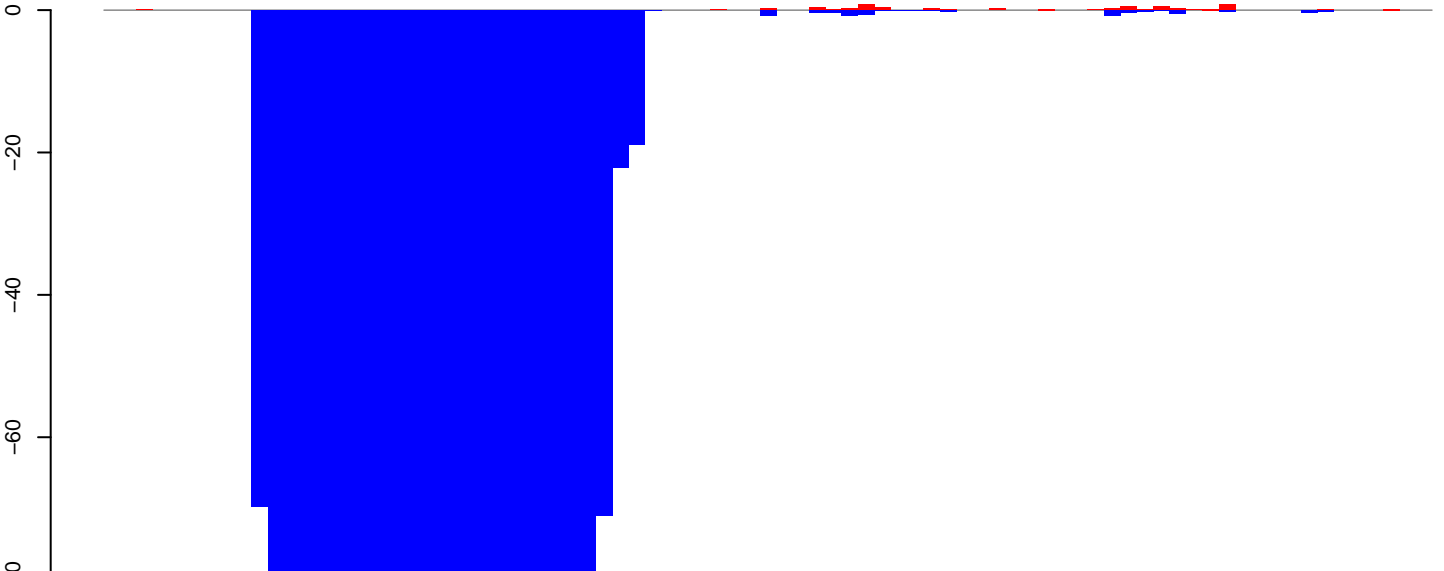
positive=3, negative=223, total=226

CuQuin_Testes_NL.24_35.rep



positive=3, negative=308, total=311

CuQuin_Testes_NL.rep

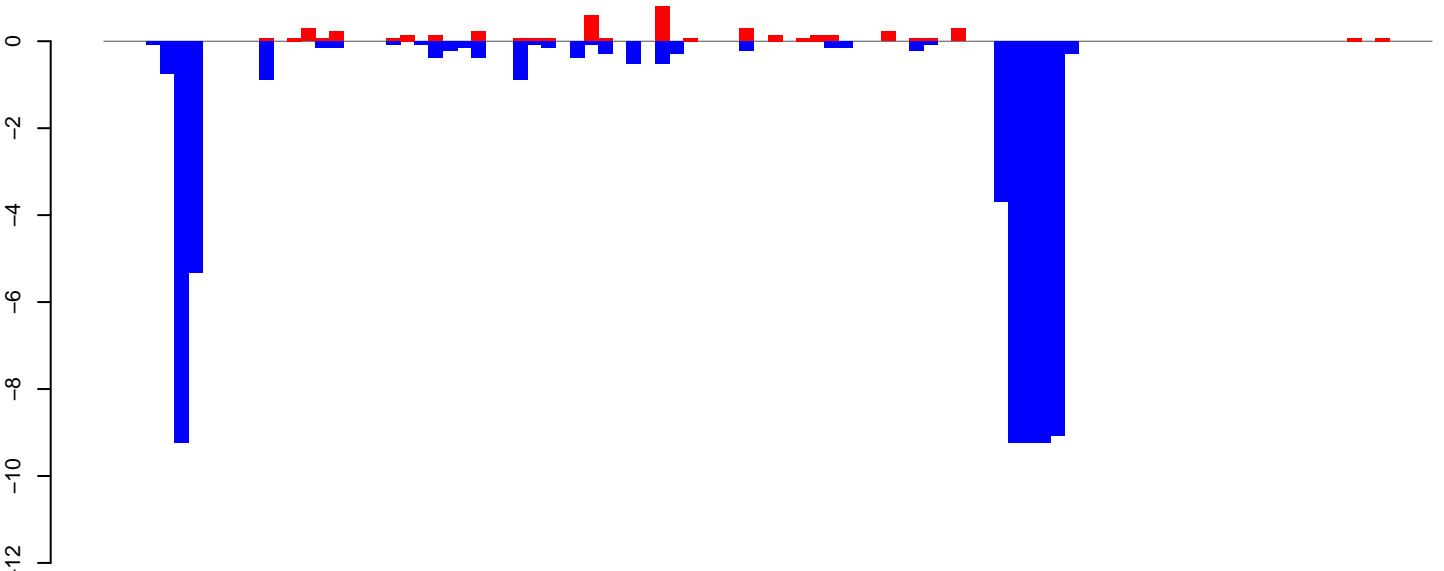


positive=6, negative=1849, total=1855

Window size=25, length=2012, TE@rnd-3_family-9@DNA@711:1-2012

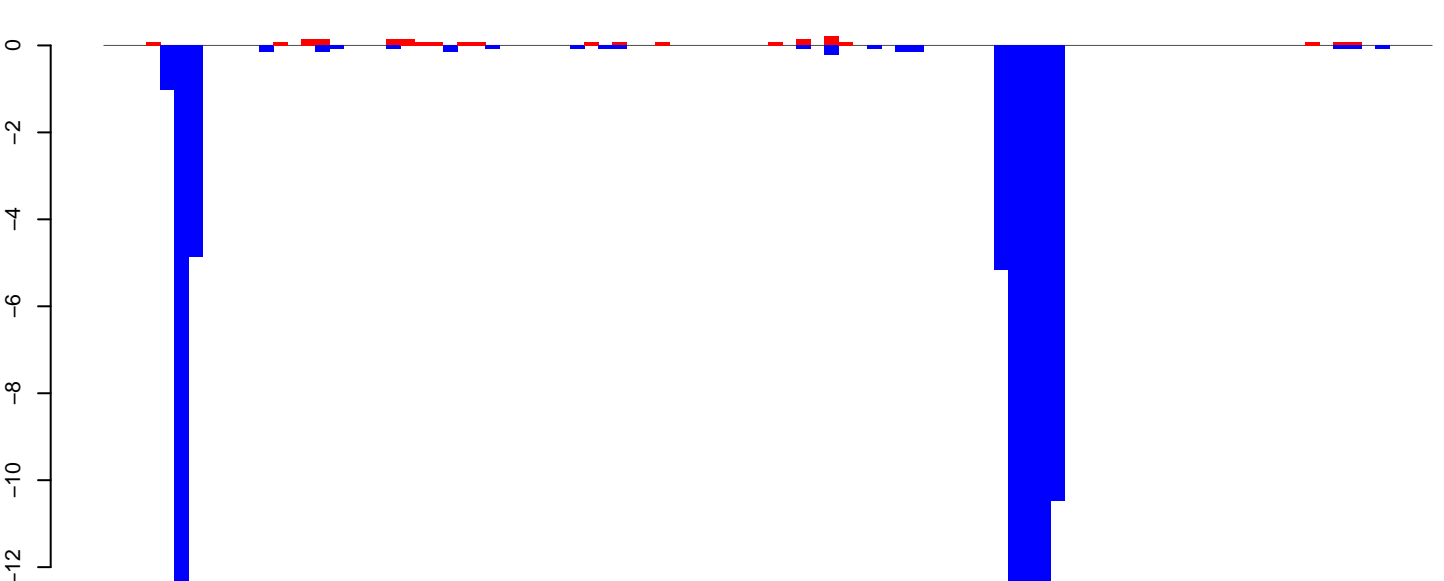
0 500 1000 1500 2000

CuQuin_Testes_NL.18_23.rep



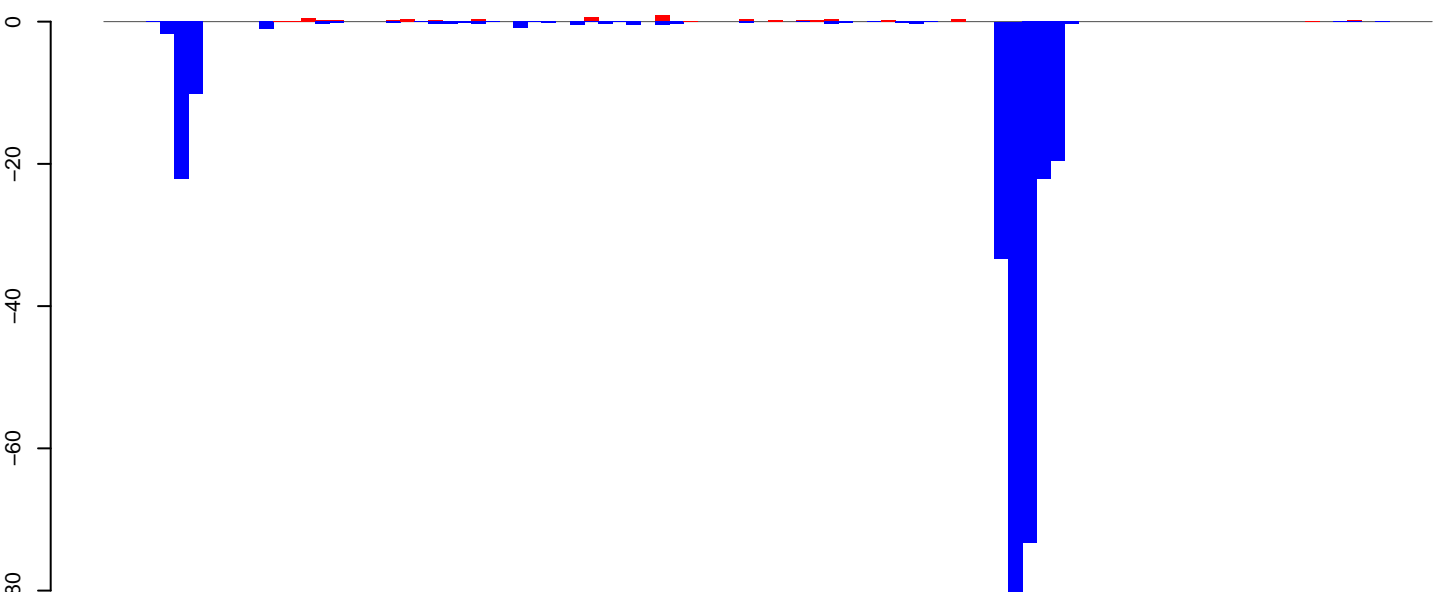
positive=5, negative=63, total=67

CuQuin_Testes_NL.24_35.rep



positive=2, negative=75, total=77

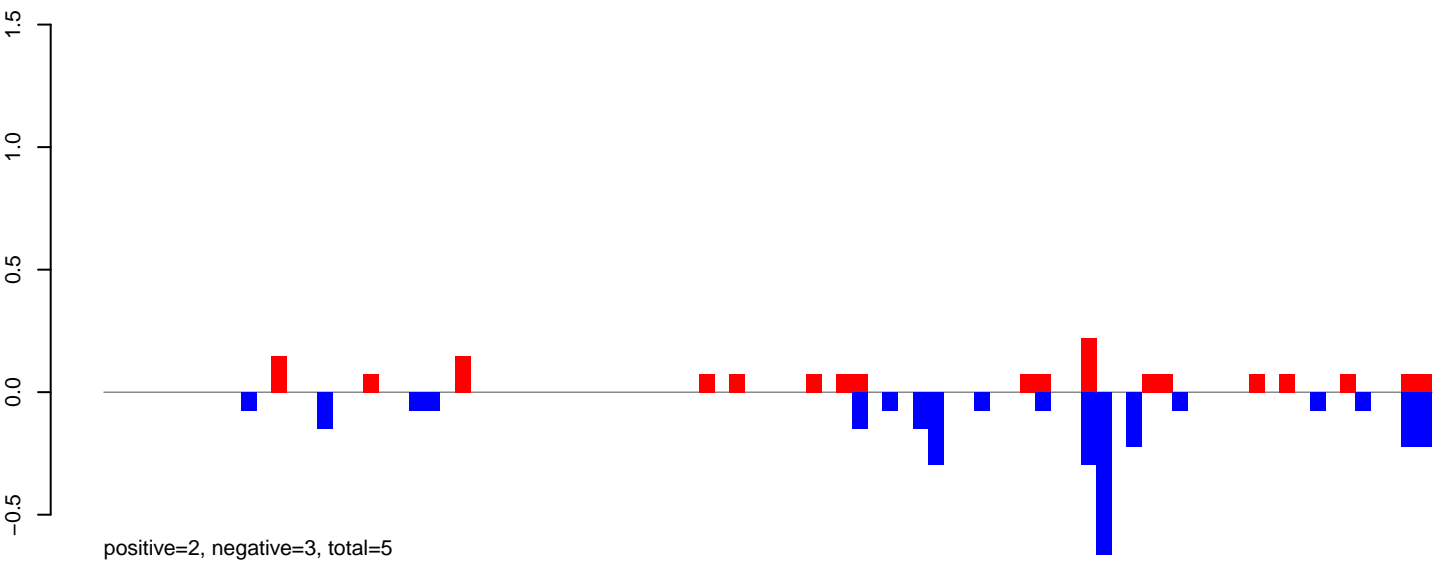
CuQuin_Testes_NL.rep



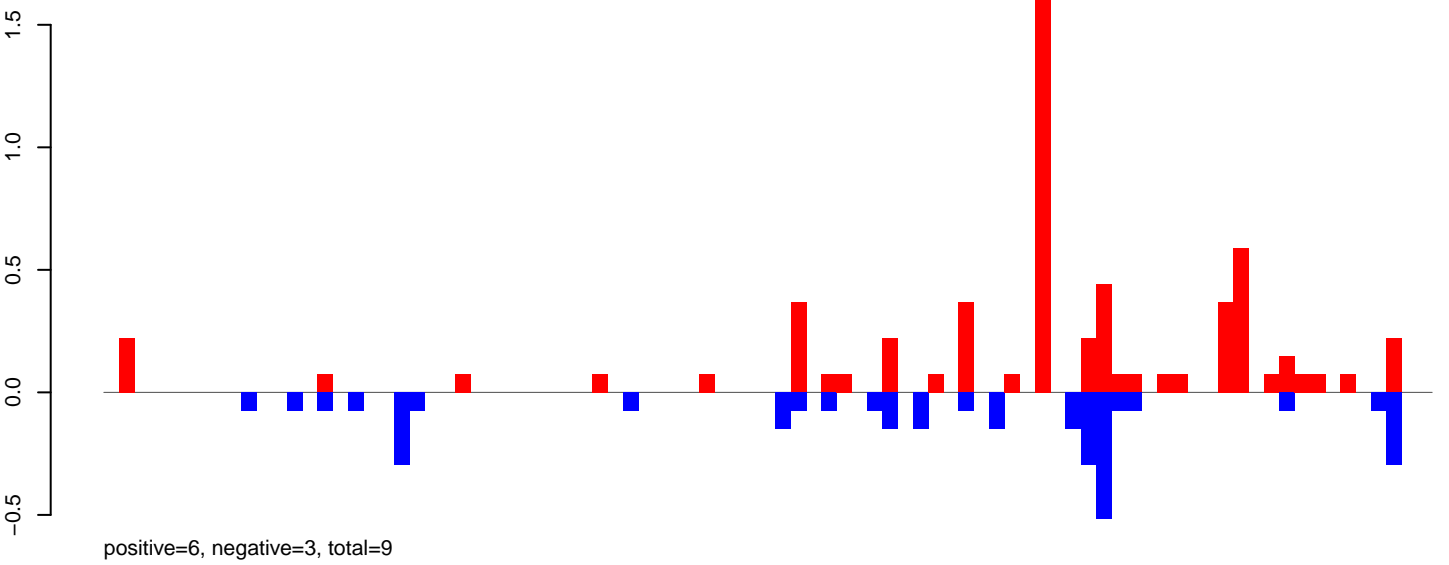
positive=7, negative=274, total=281

Window size=25, length=2336, TE@[rmd-3_family-417@DNA_hAT@1345:1-2336](#)

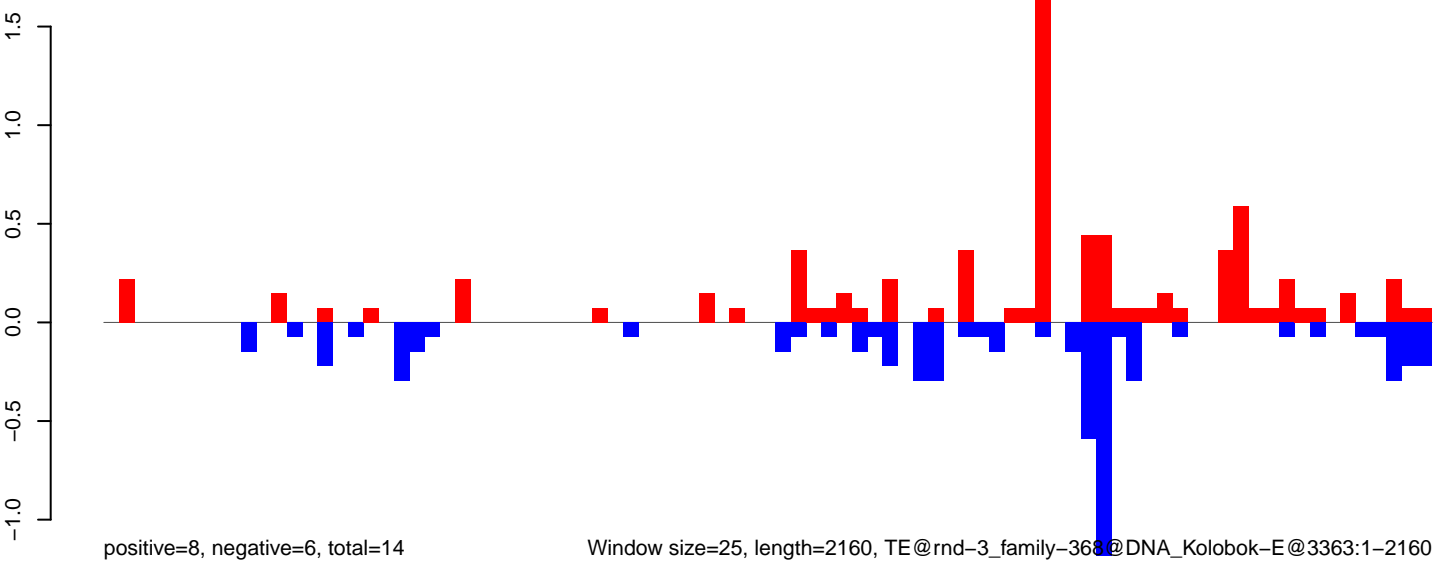
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

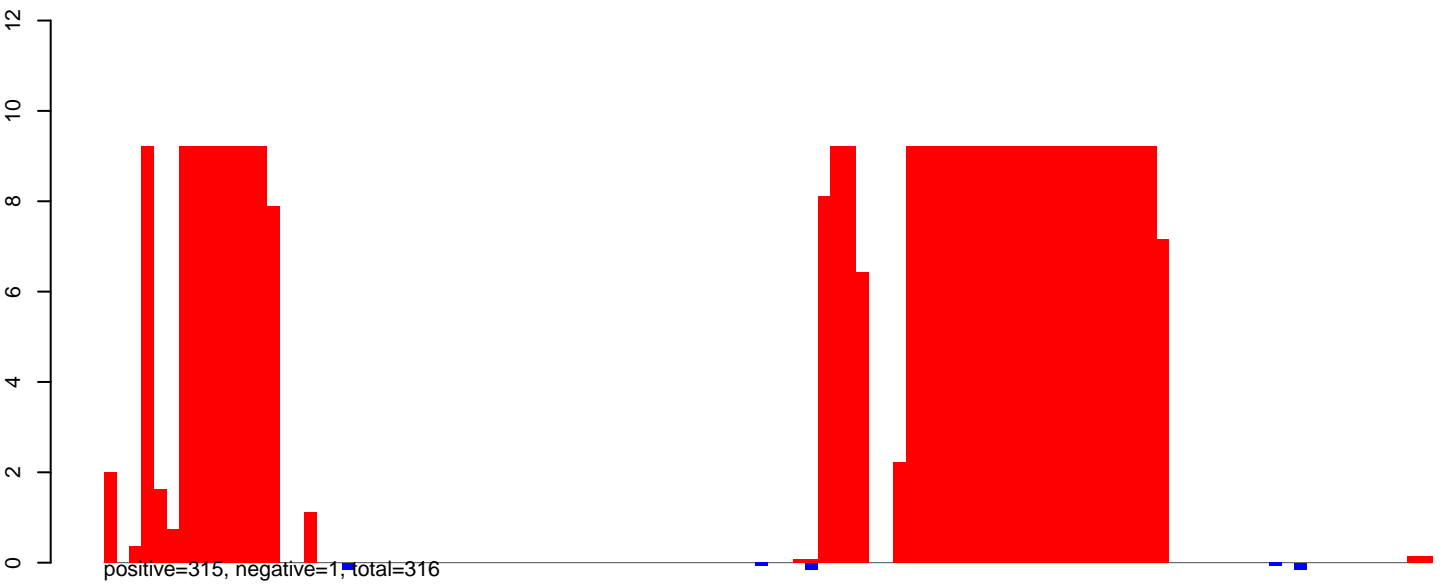


CuQuin_Testes_NL.rep

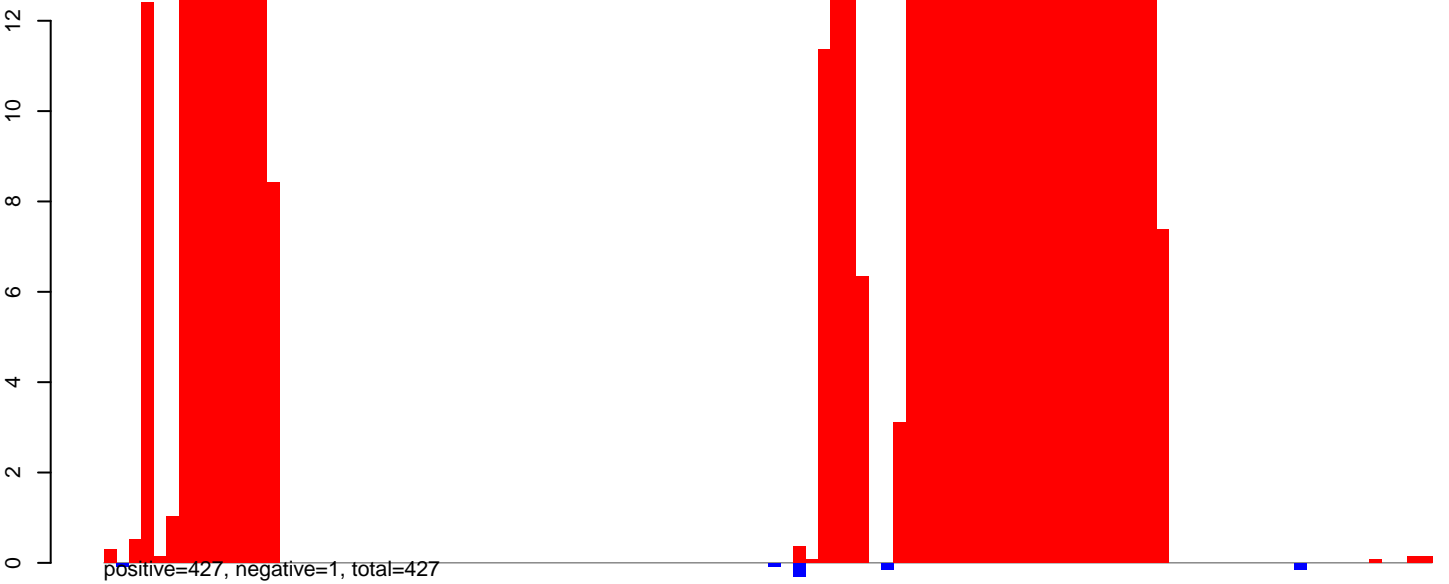


Window size=25, length=2160, TE@rnd-3_family-368@DNA_Kolobok-E@3363:1-2160

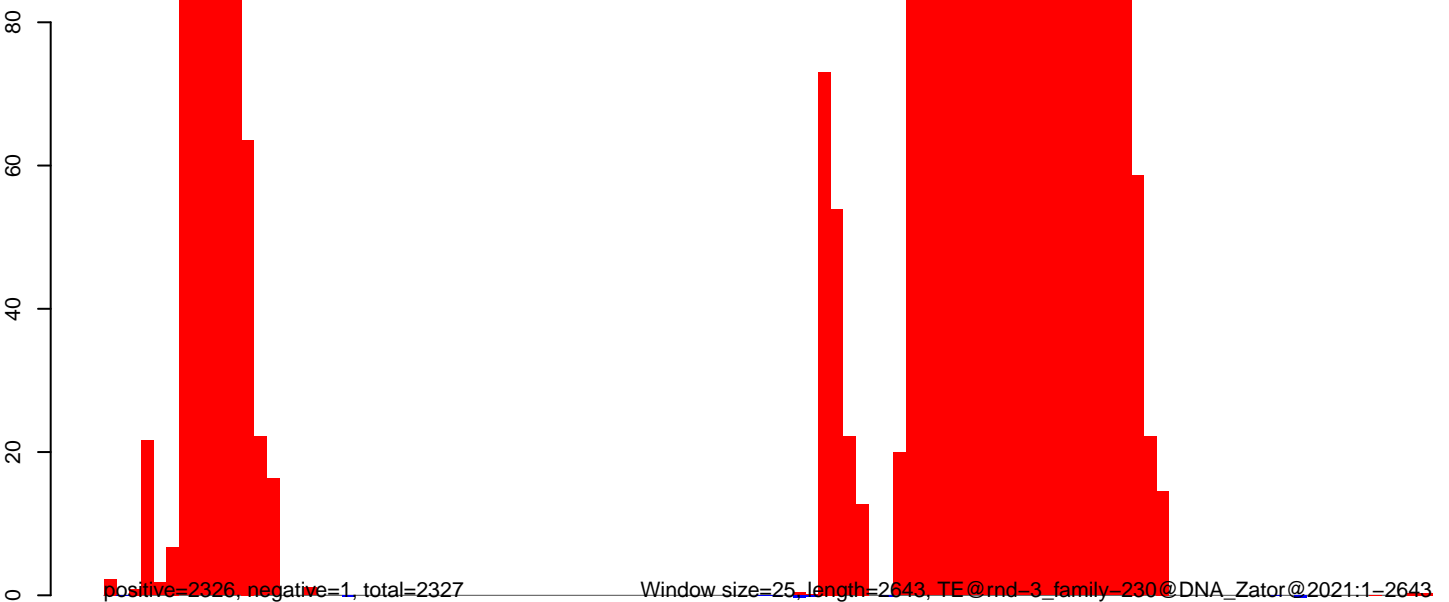
CuQuin_Testes_NL.18_23.rep



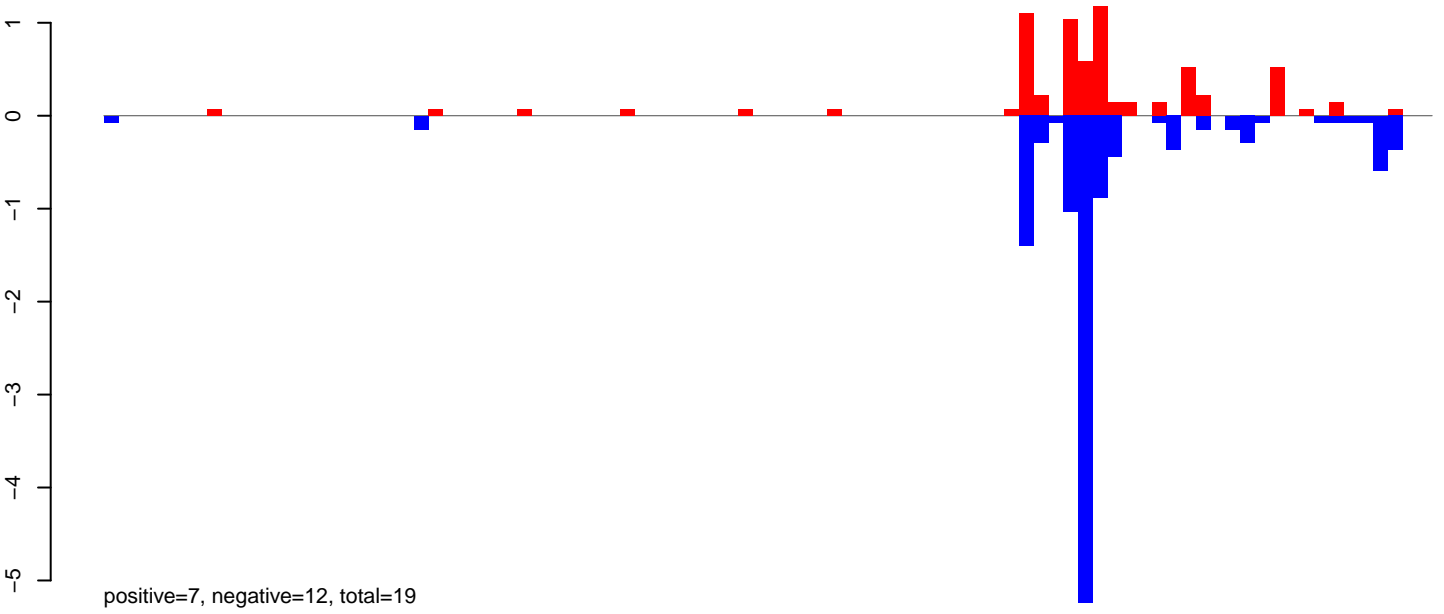
CuQuin_Testes_NL.24_35.rep



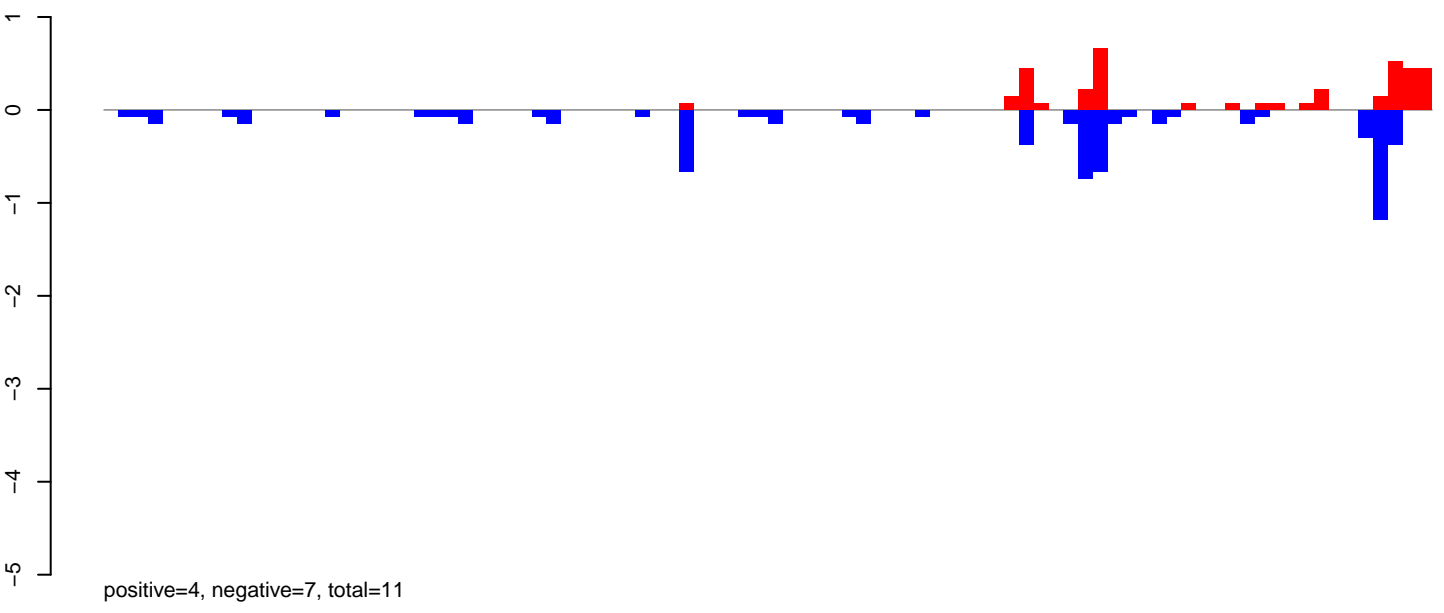
CuQuin_Testes_NL.rep



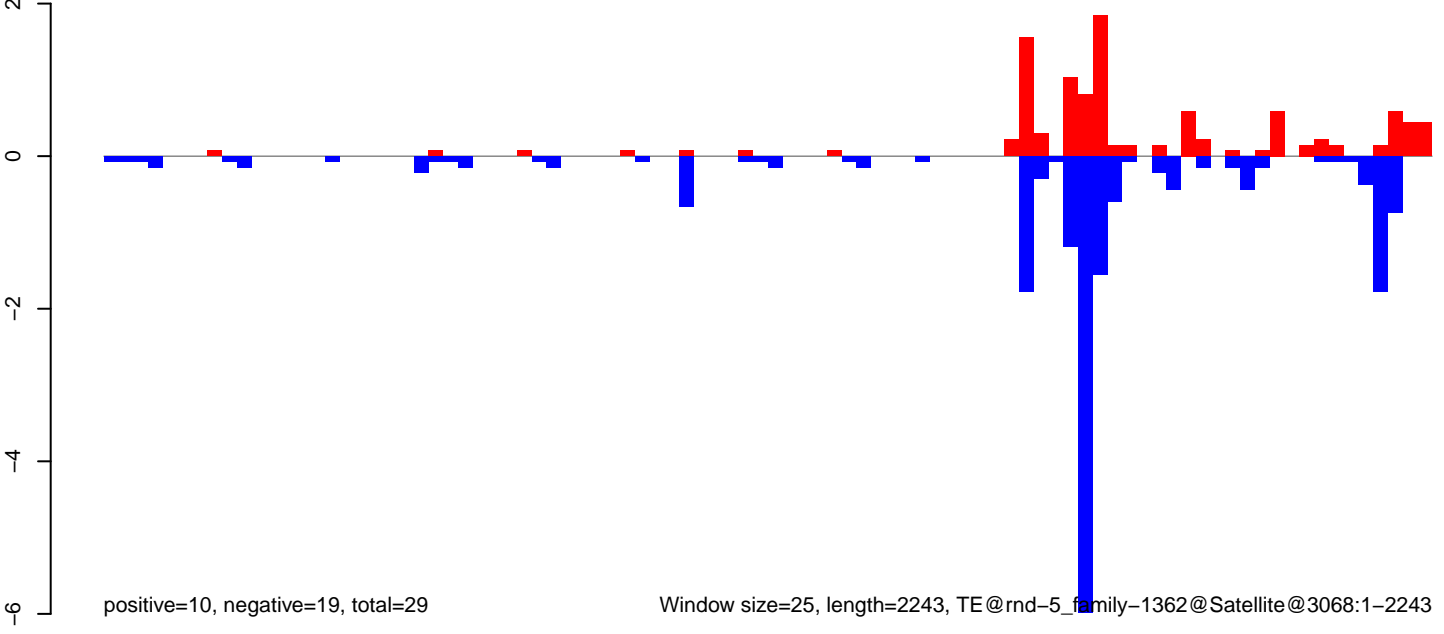
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

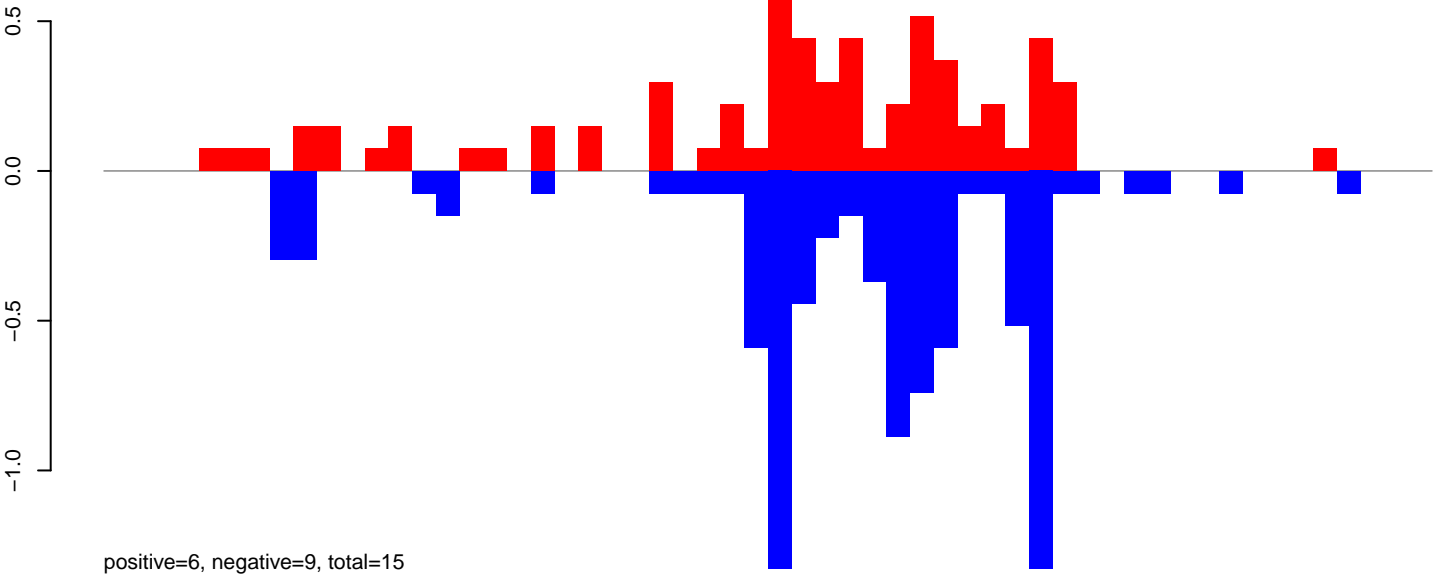


CuQuin_Testes_NL.rep

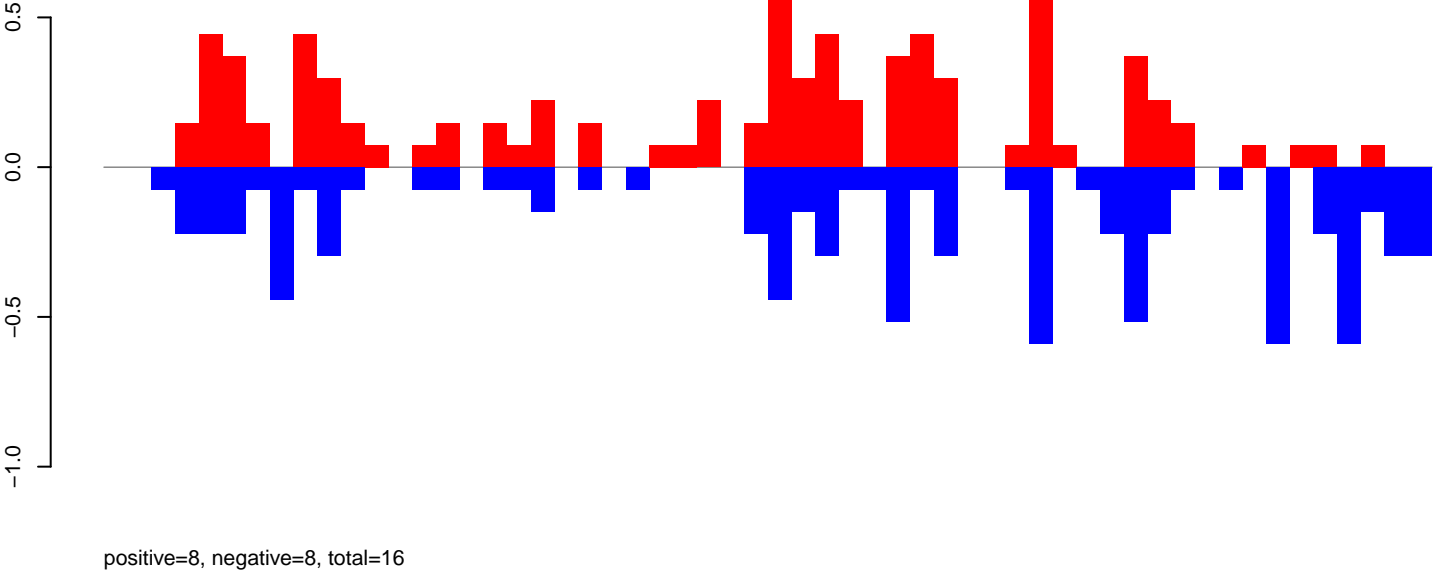


Window size=25, length=2243, TE@rnd-5_family-1362@Satellite@3068:1-2243

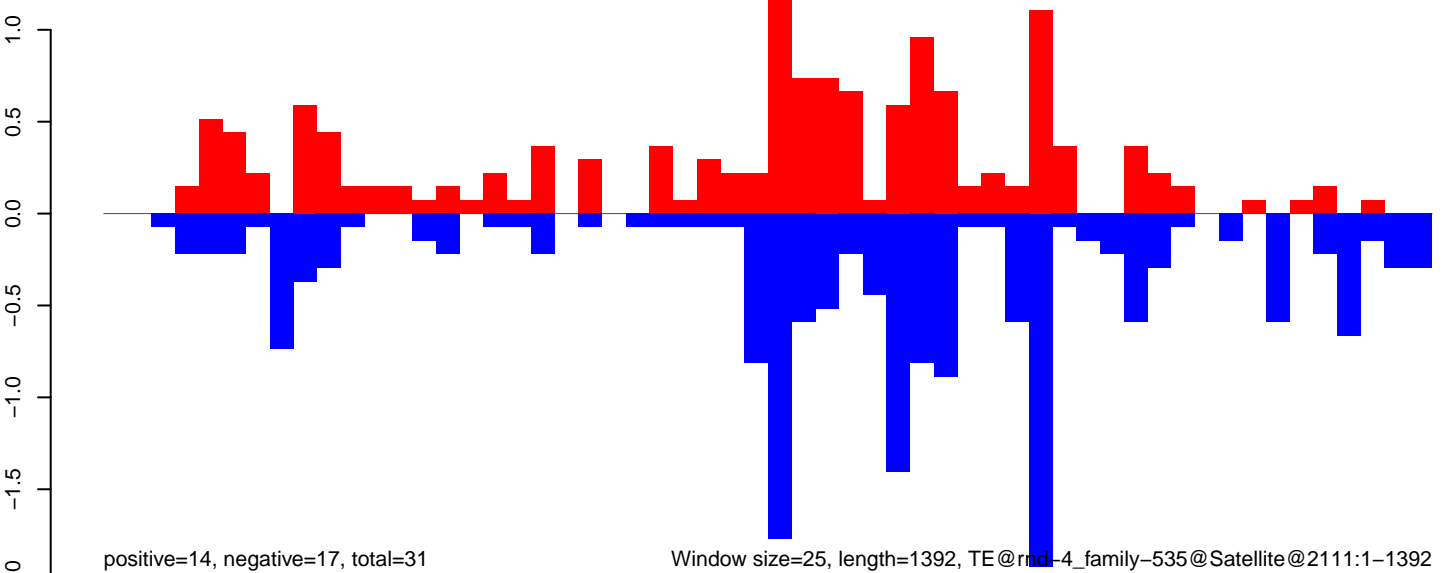
CuQuin_Testes_NL.18_23.rep



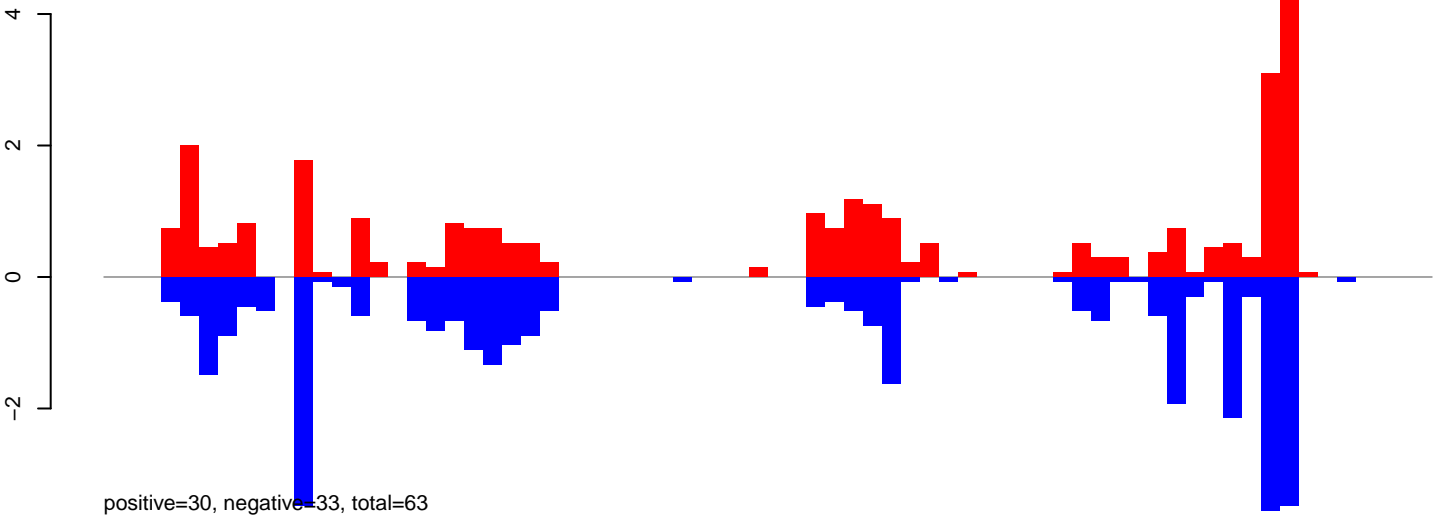
CuQuin_Testes_NL.24_35.rep



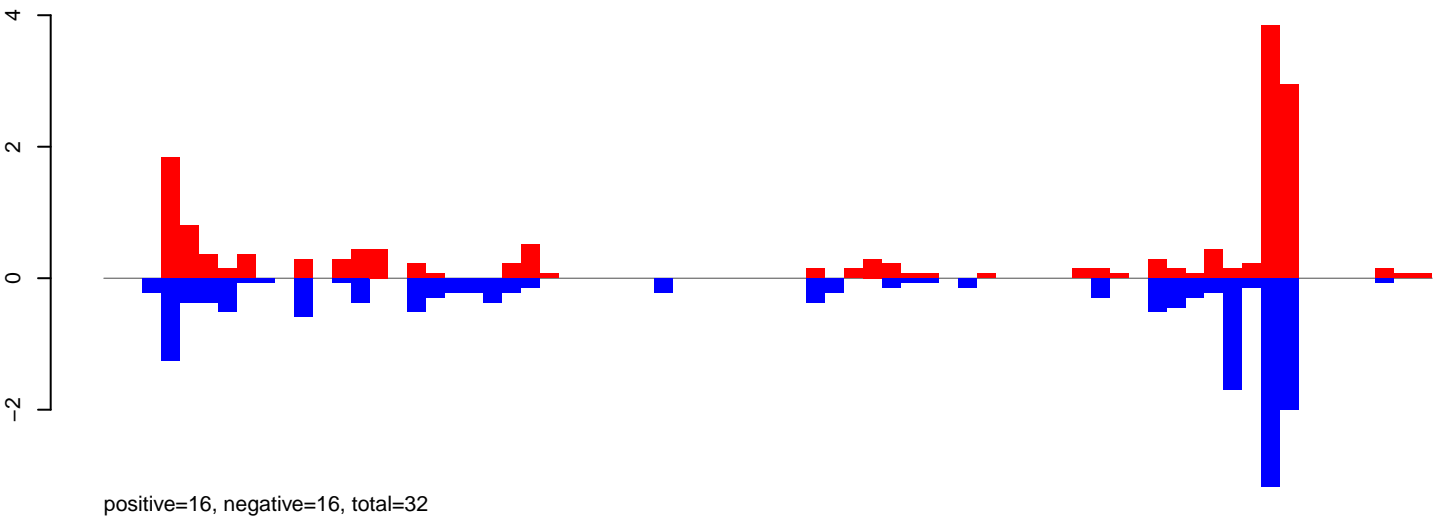
CuQuin_Testes_NL.rep



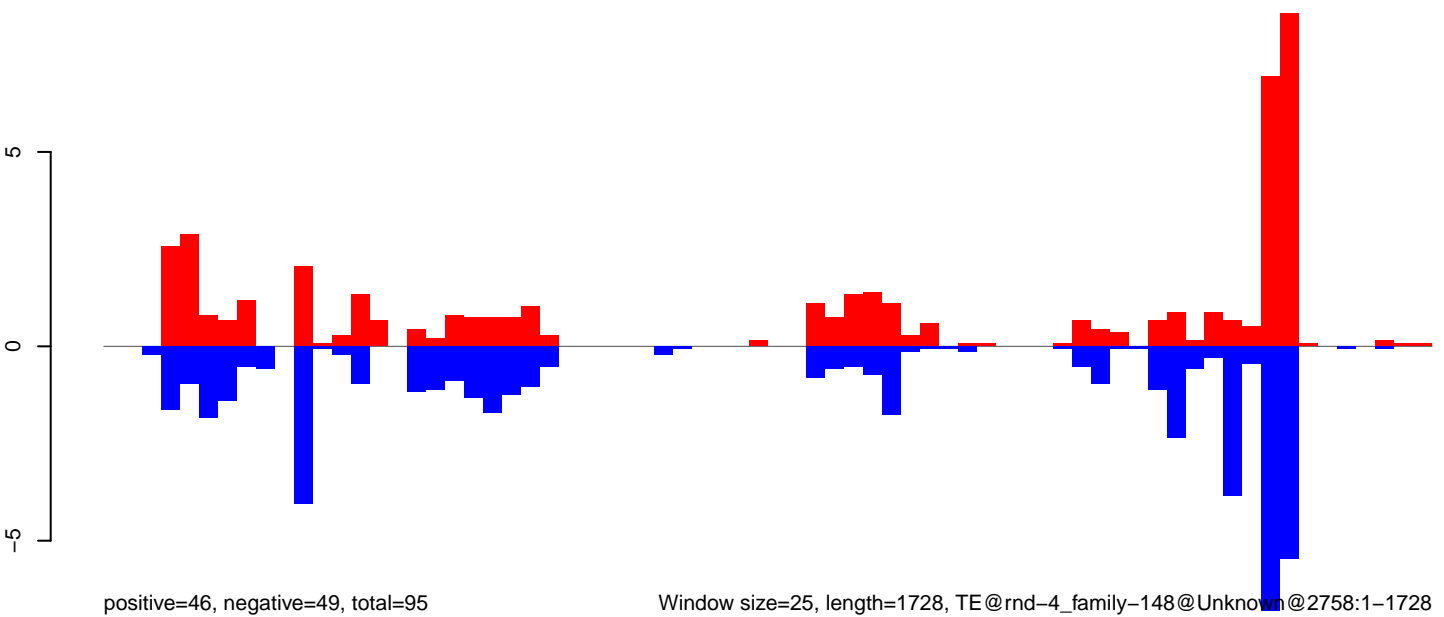
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

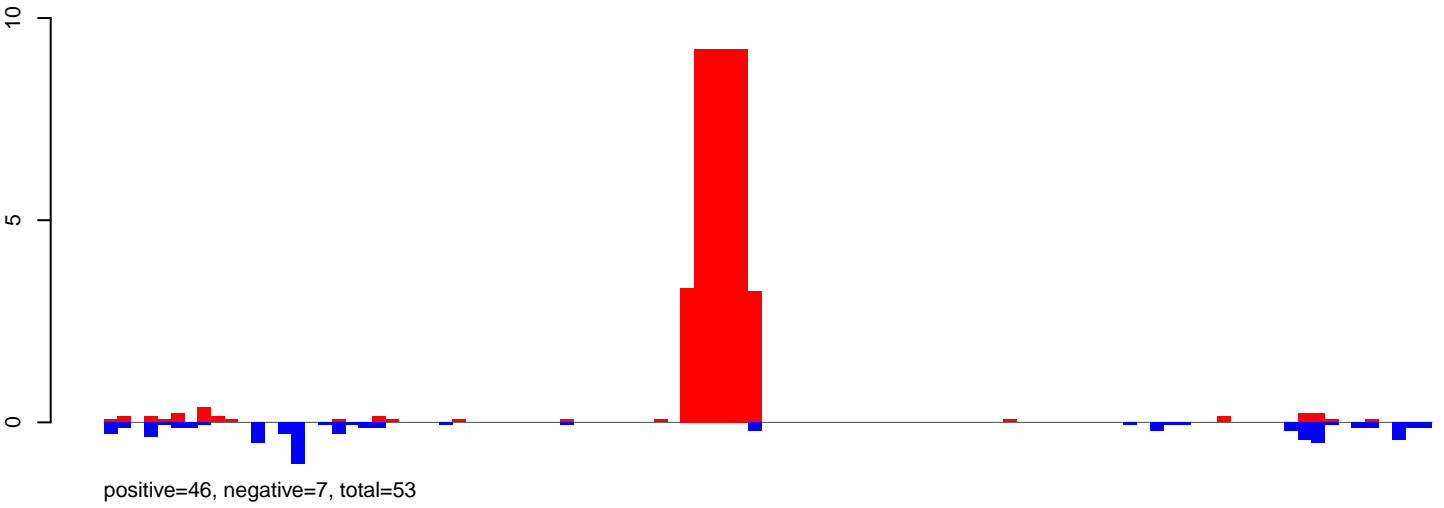


CuQuin_Testes_NL.rep

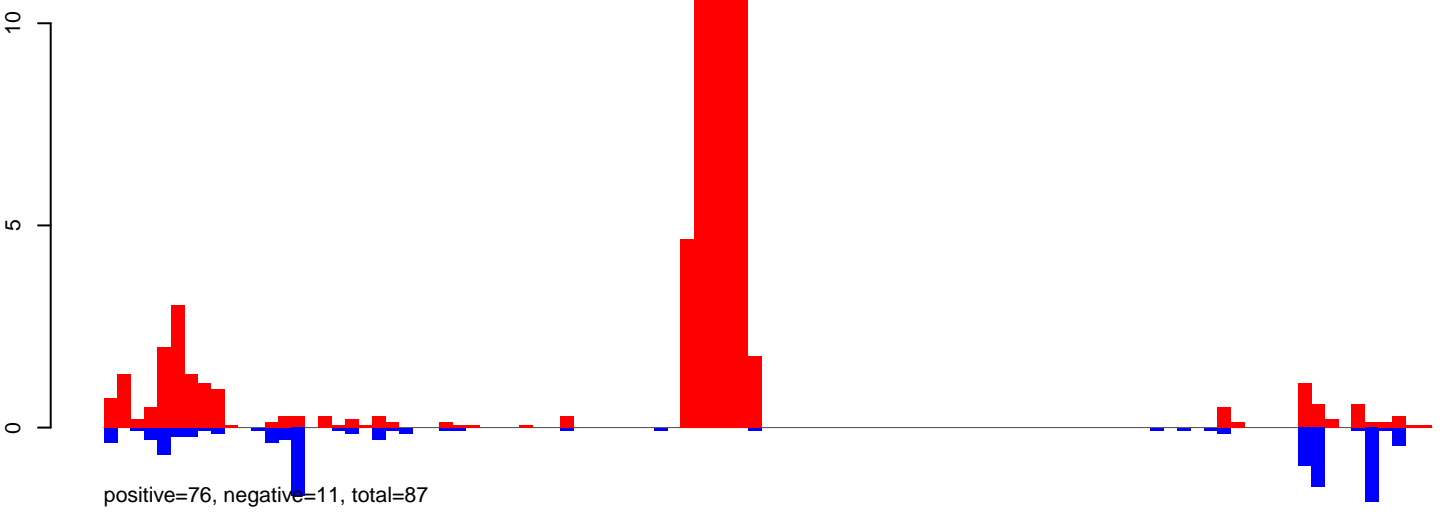


Window size=25, length=1728, TE@rnd-4_family-148@Unknown@2758:1-1728

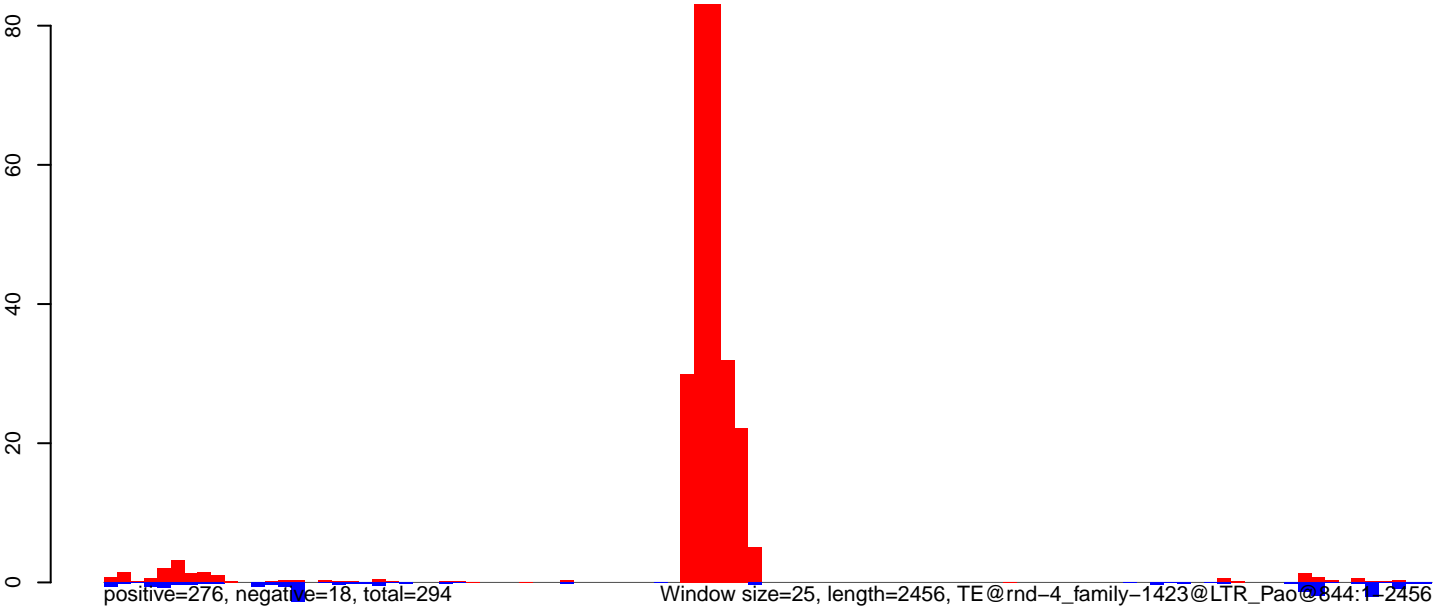
CuQuin_Testes_NL.18_23.rep



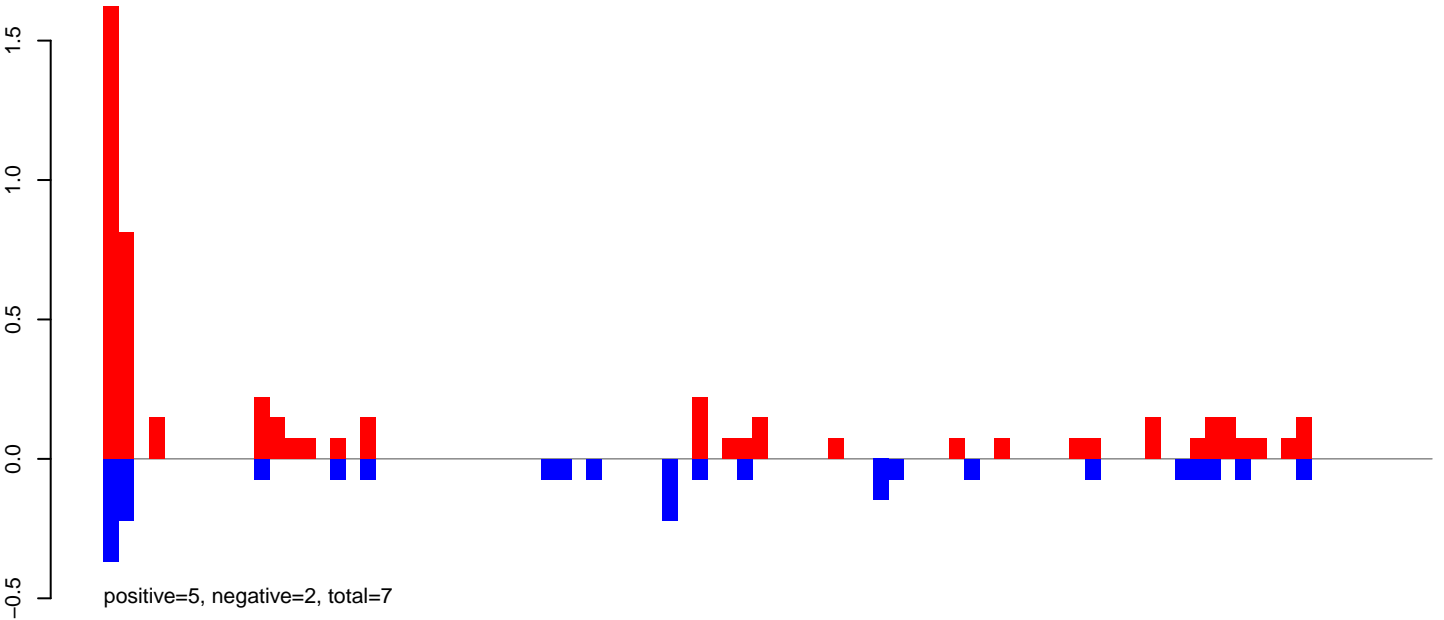
CuQuin_Testes_NL.24_35.rep



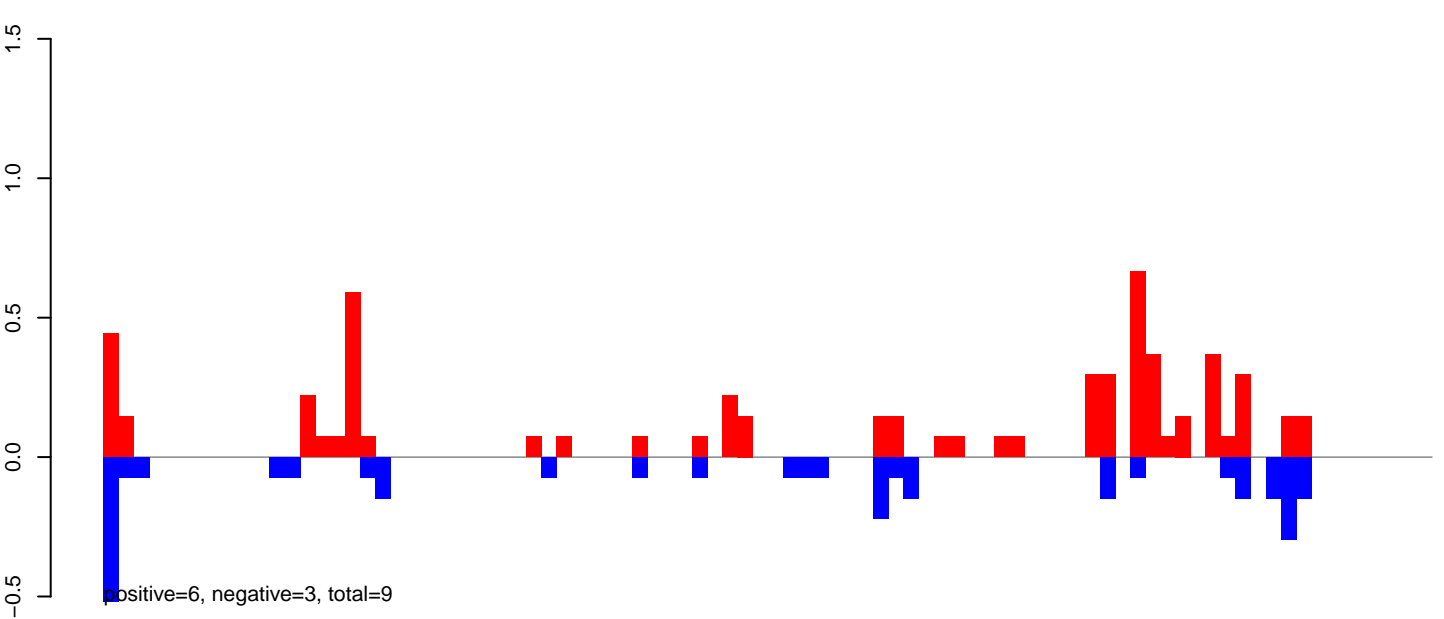
CuQuin_Testes_NL.rep



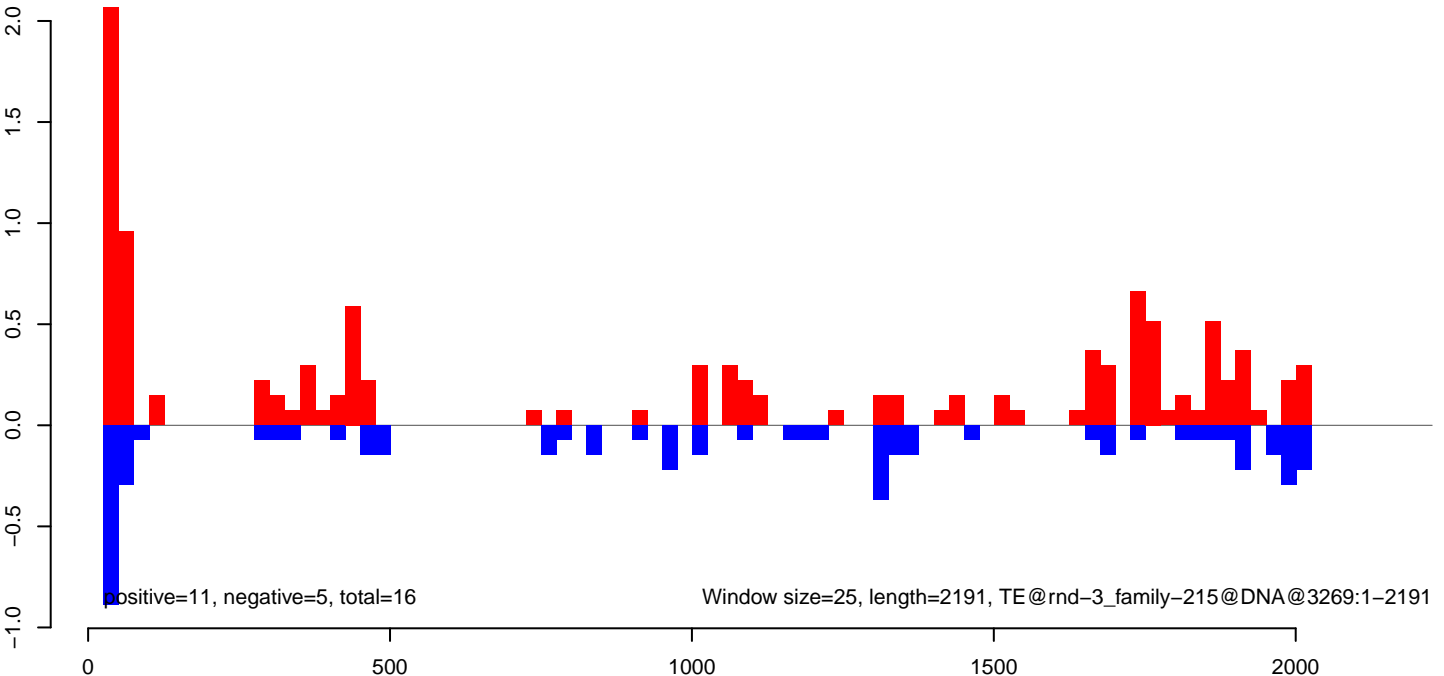
CuQuin_Testes_NL.18_23.rep



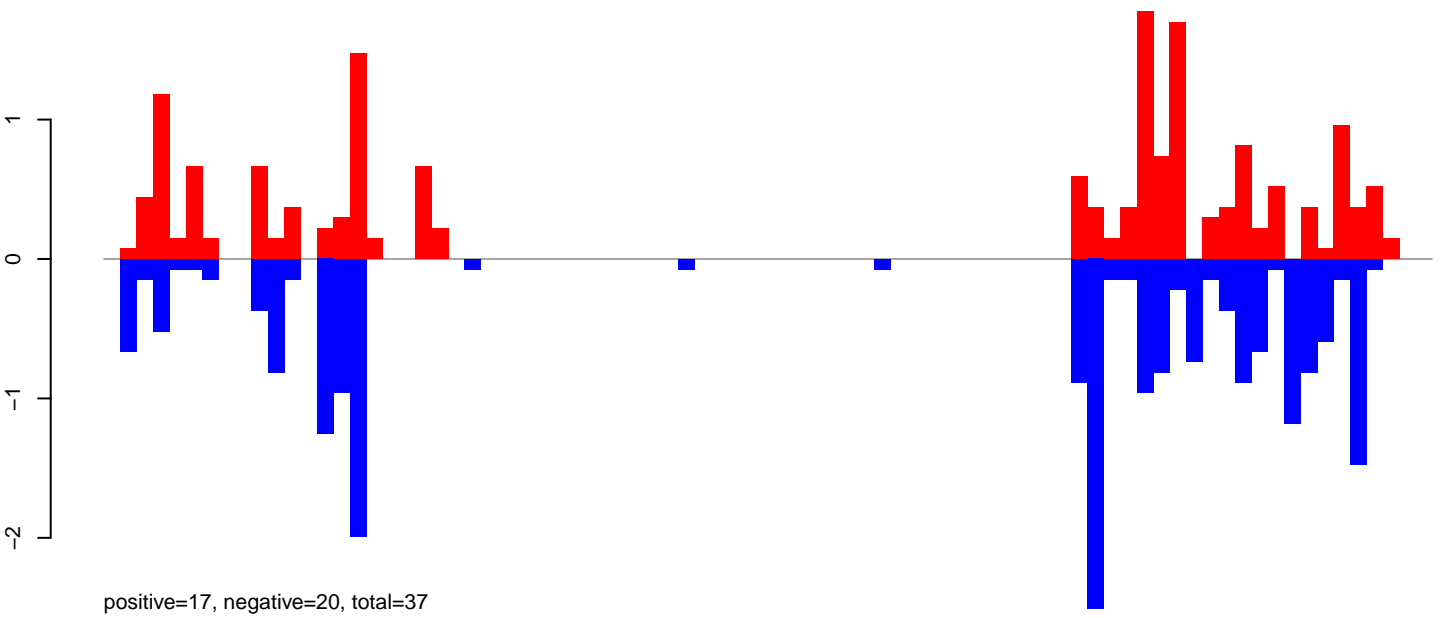
CuQuin_Testes_NL.24_35.rep



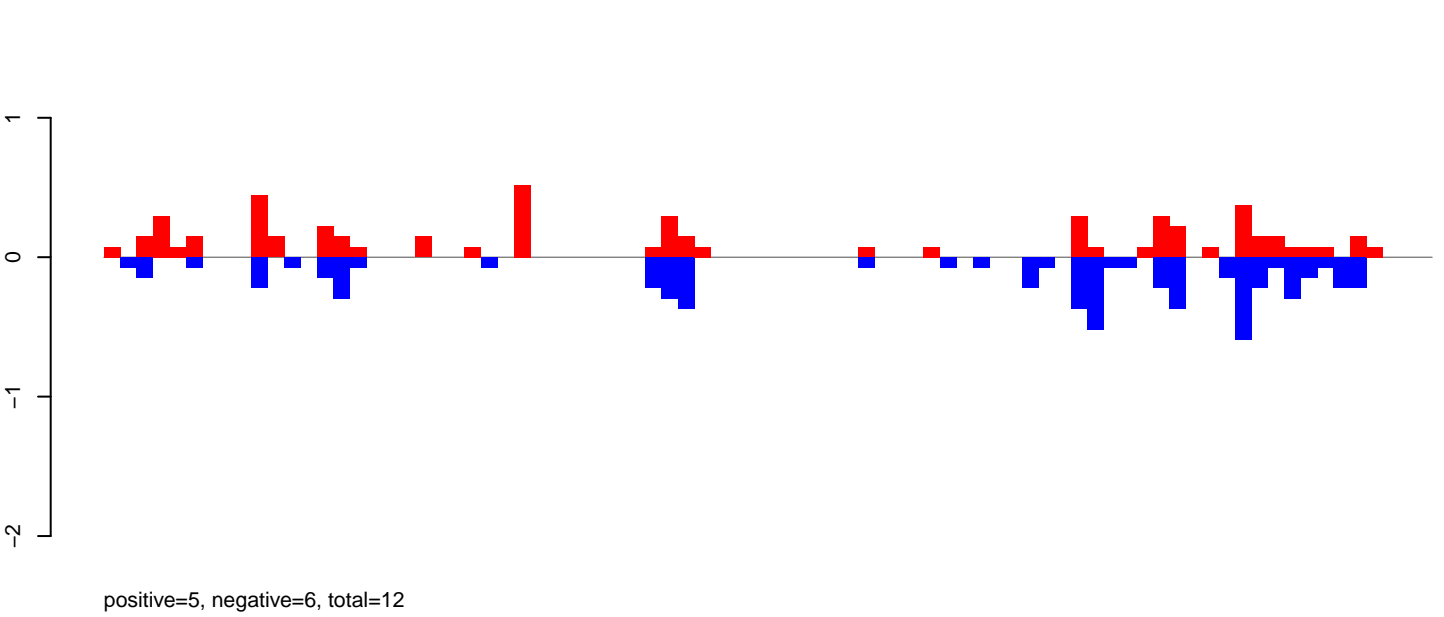
CuQuin_Testes_NL.rep



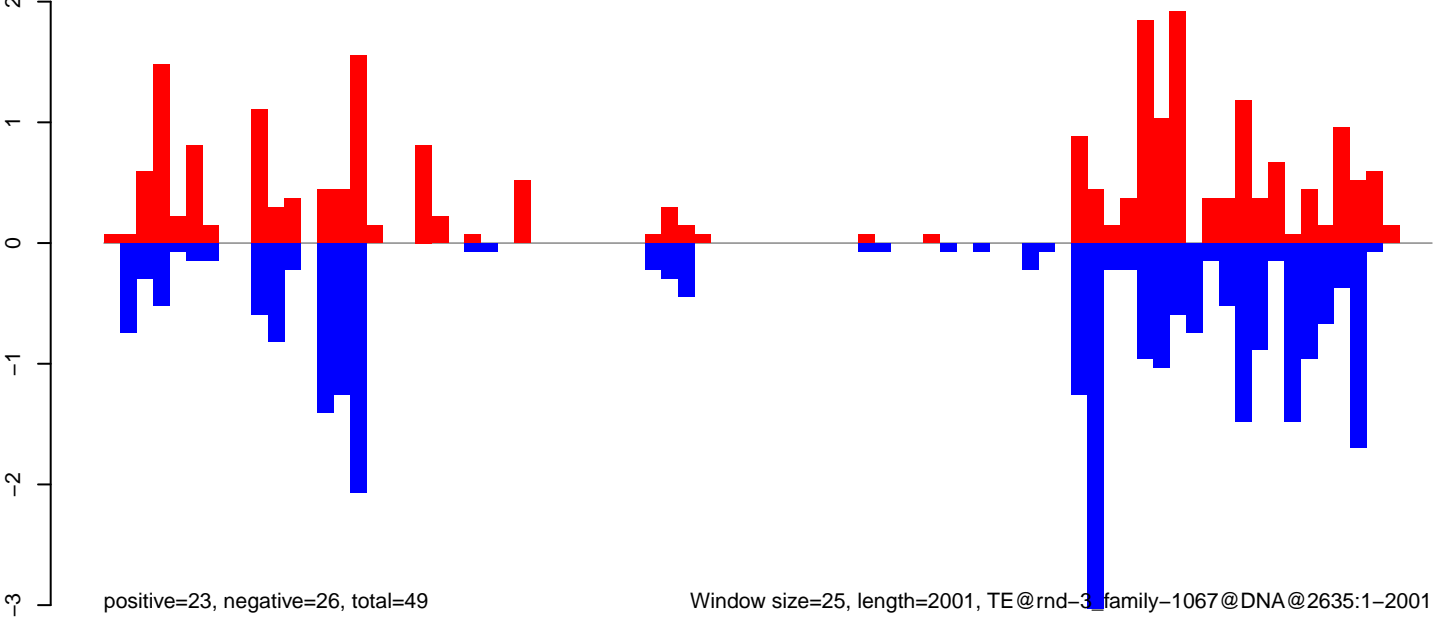
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

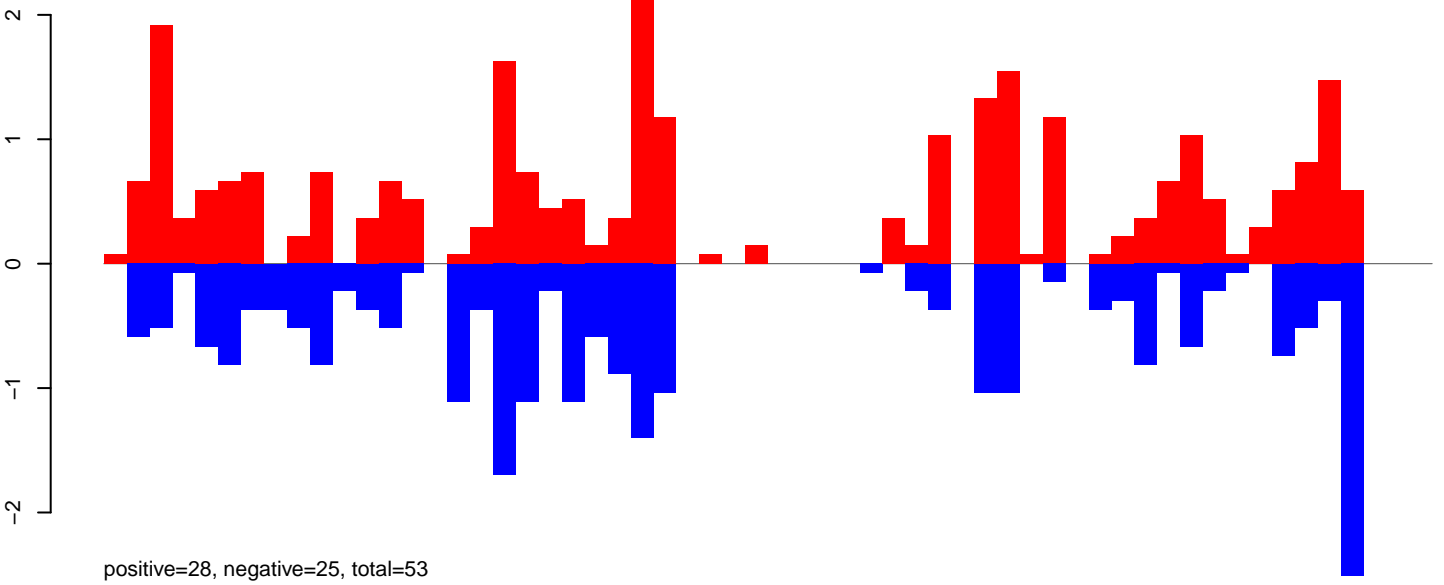


CuQuin_Testes_NL.rep

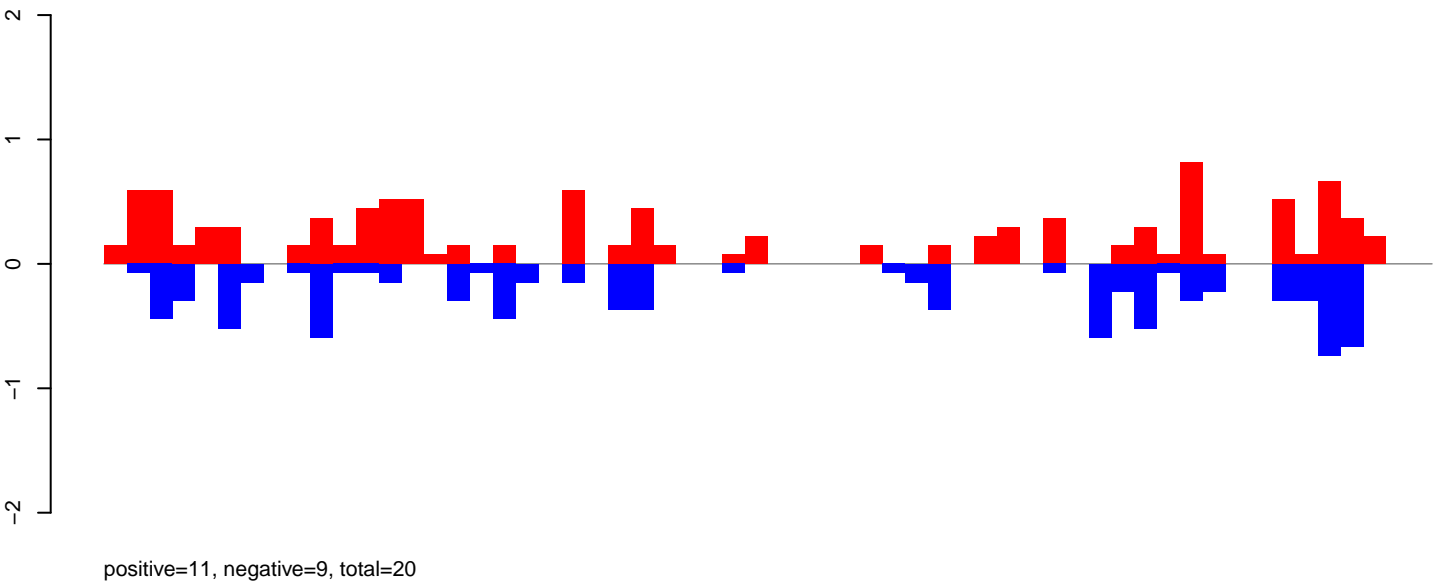


Window size=25, length=2001, TE@rnd-3 family-1067@DNA@2635:1-2001

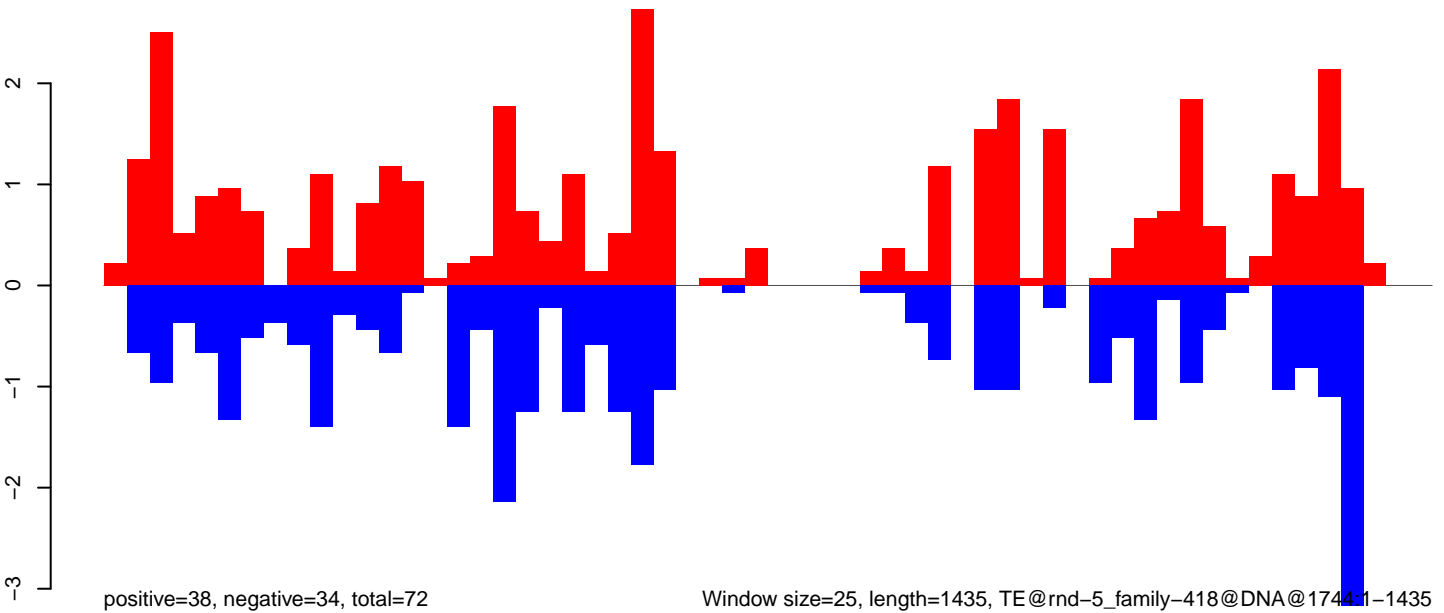
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

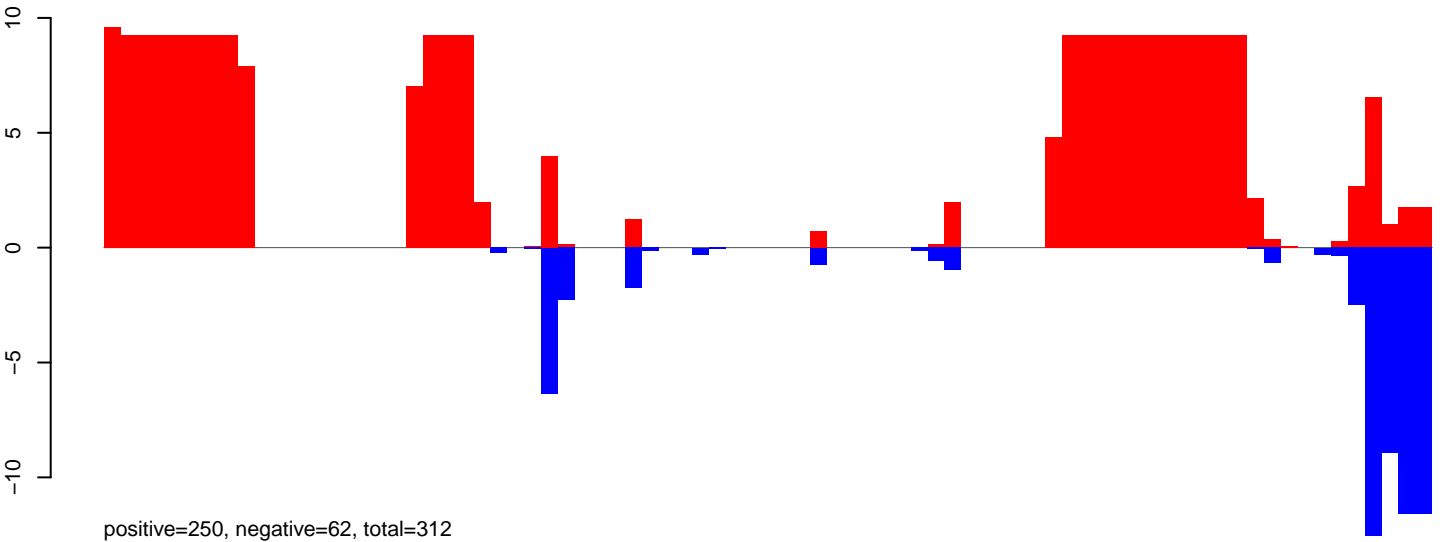


CuQuin_Testes_NL.rep

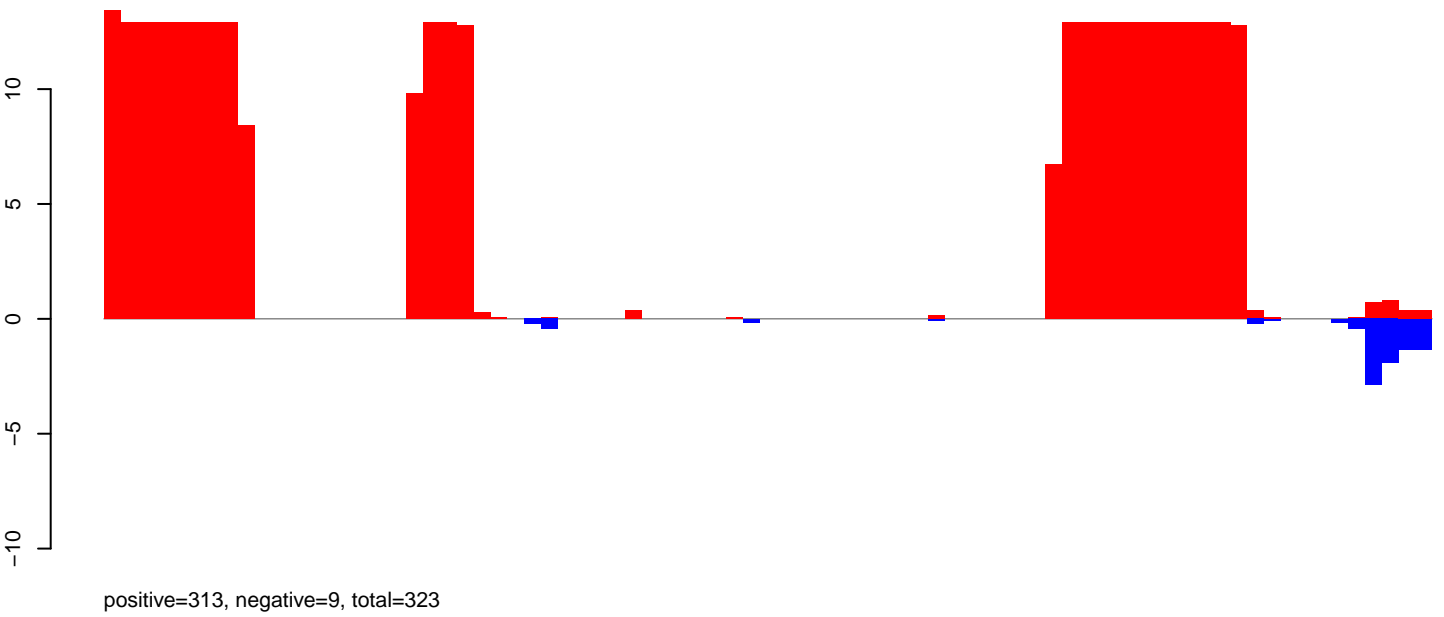


Window size=25, length=1435, TE@rnd-5_family-418@DNA@1744:1-1435

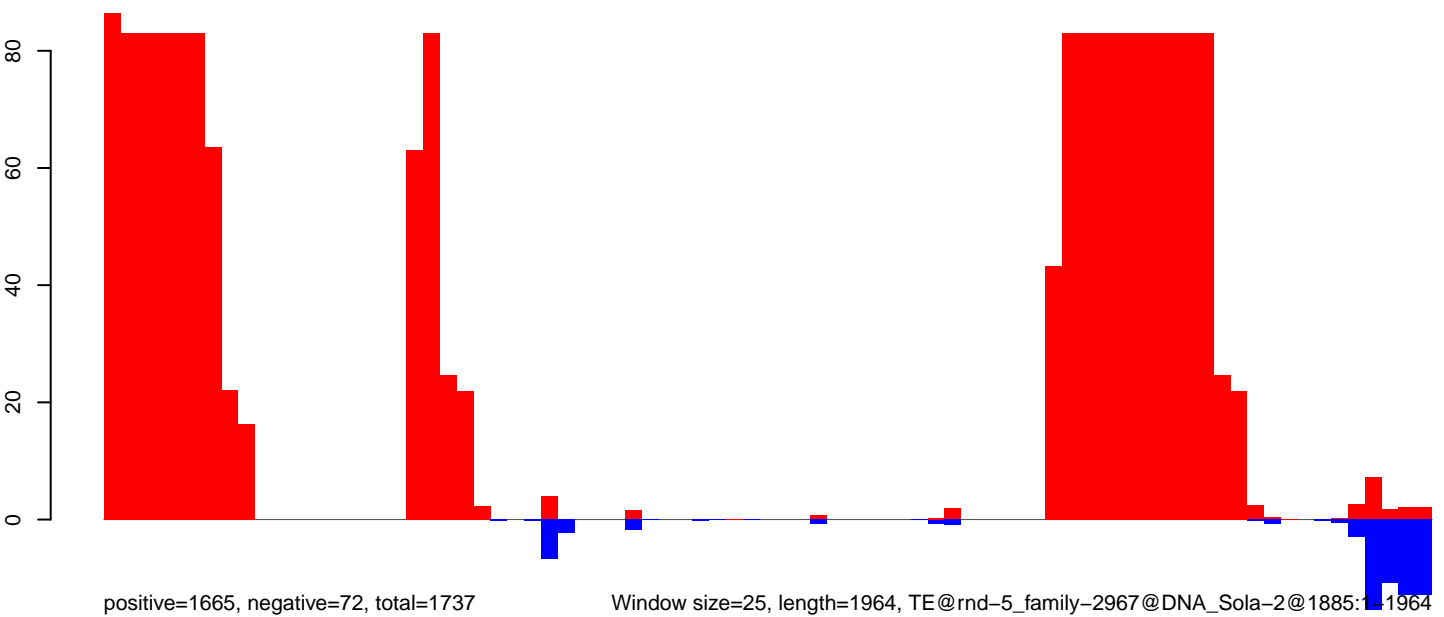
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



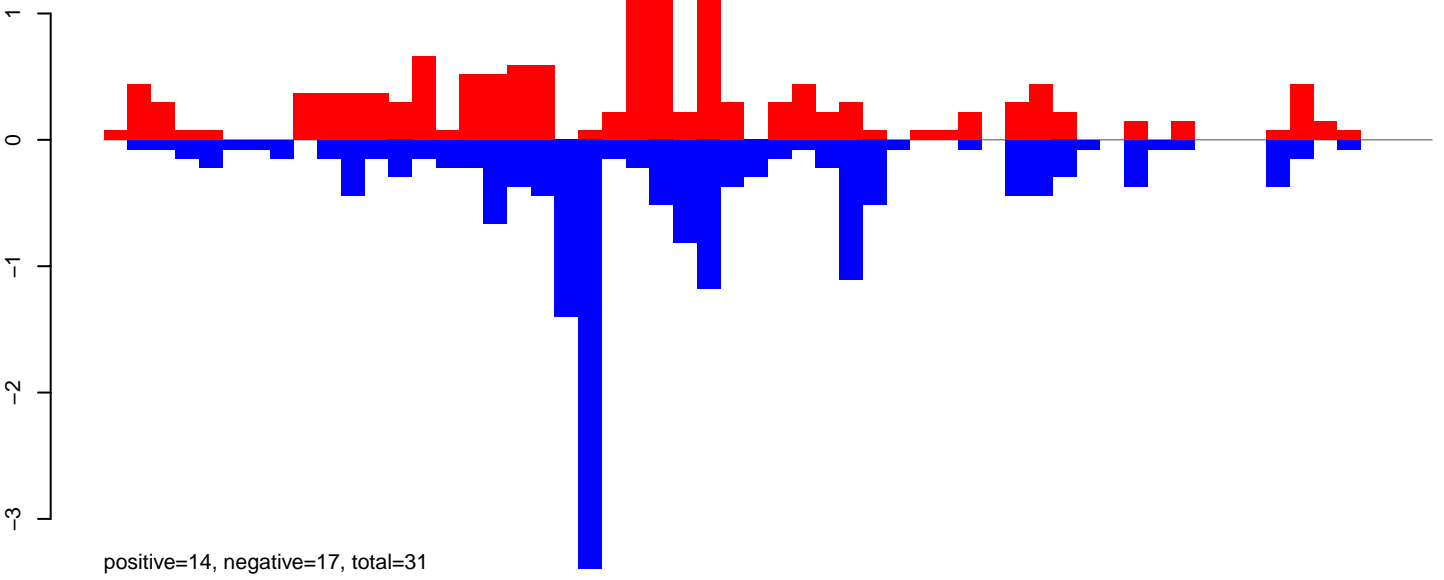
CuQuin_Testes_NL.rep



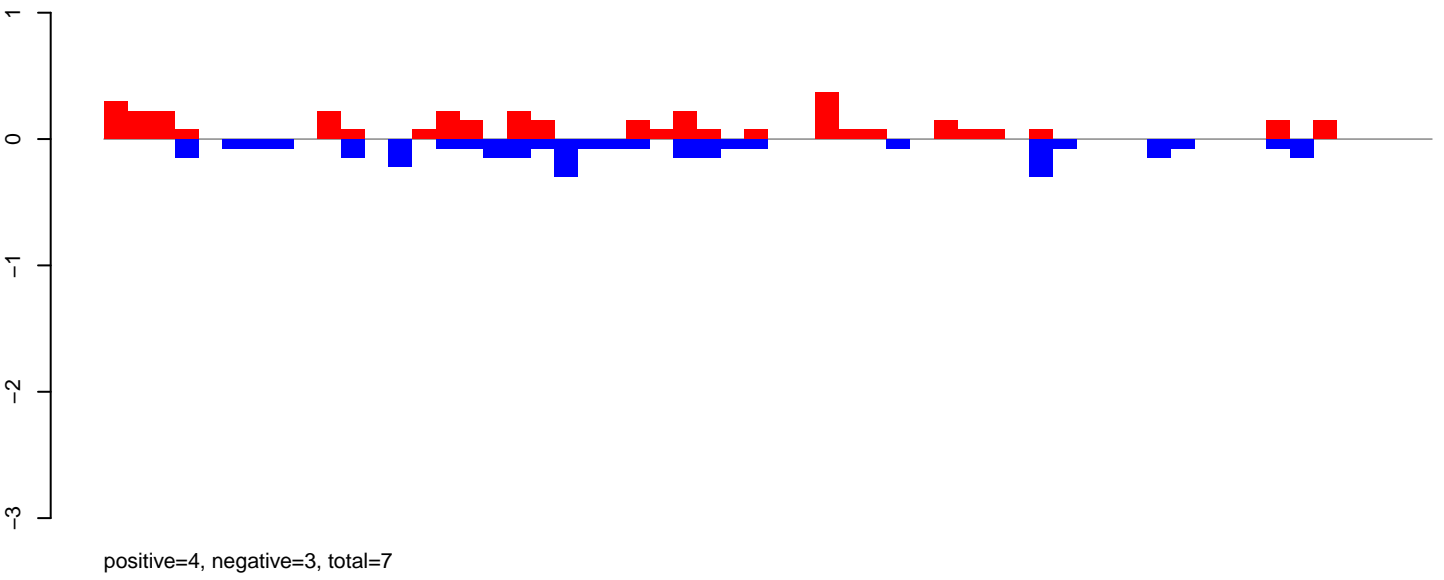
Window size=25, length=1964, TE@rnd-5_family-2967@DNA_Sola-2@1885:1-1964

0 500 1000 1500 2000

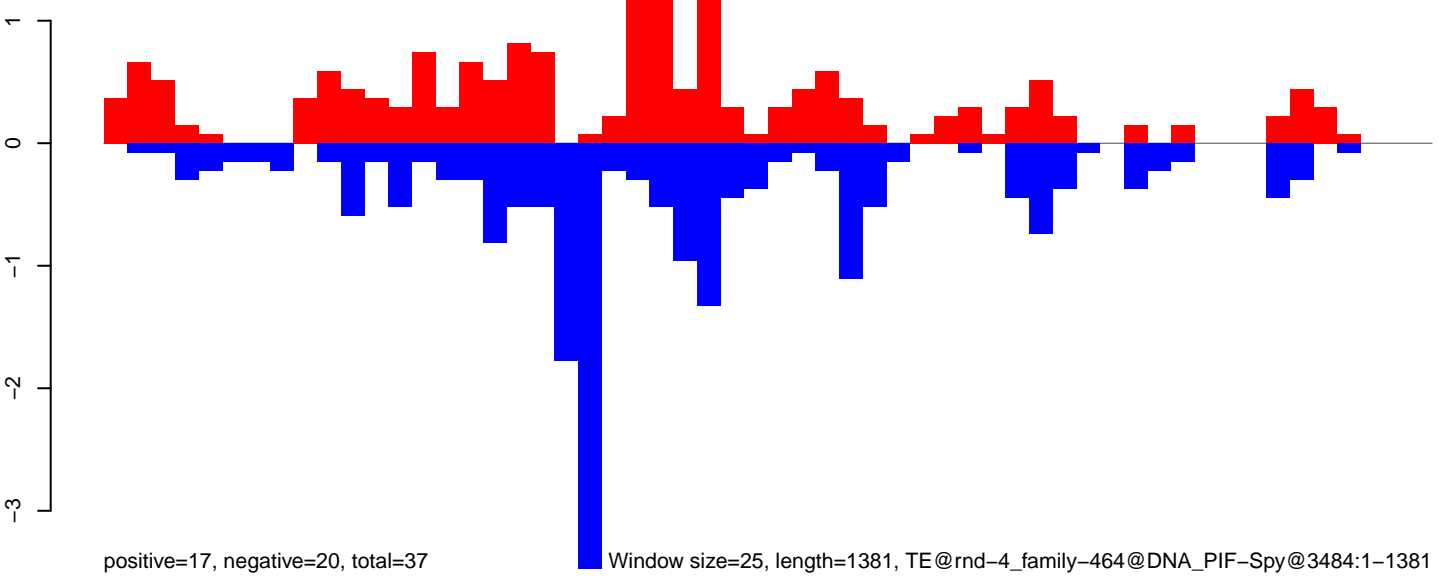
CuQuin_Testes_NL.18_23.rep



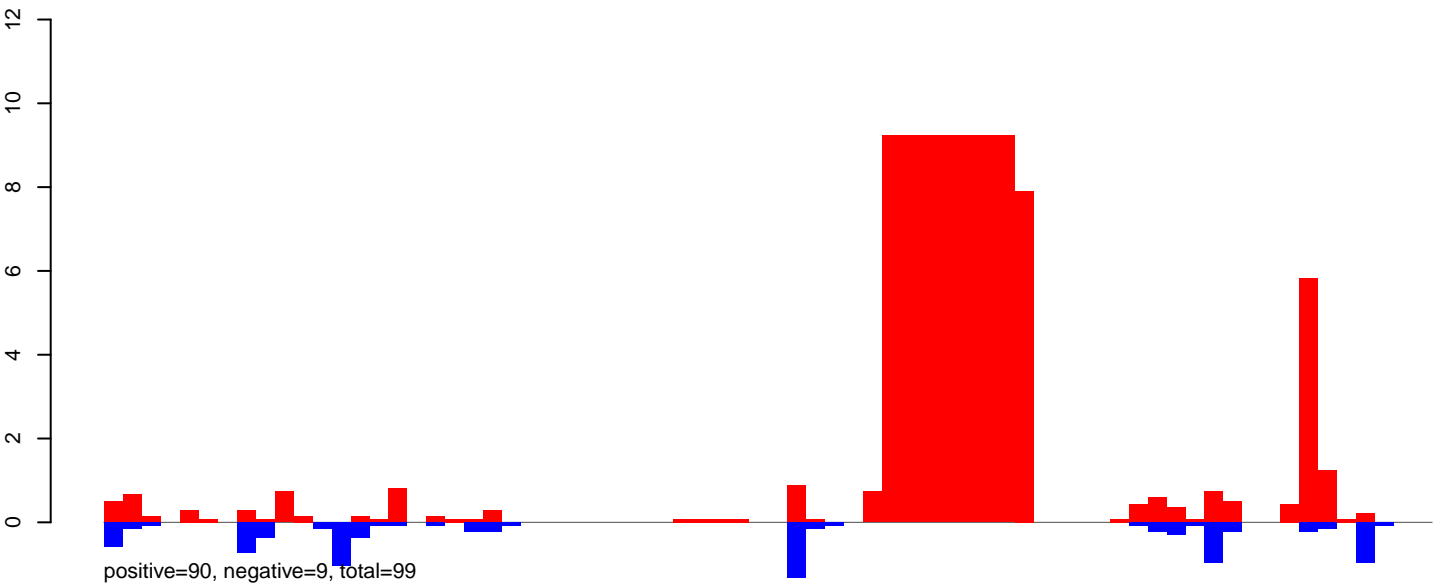
CuQuin_Testes_NL.24_35.rep



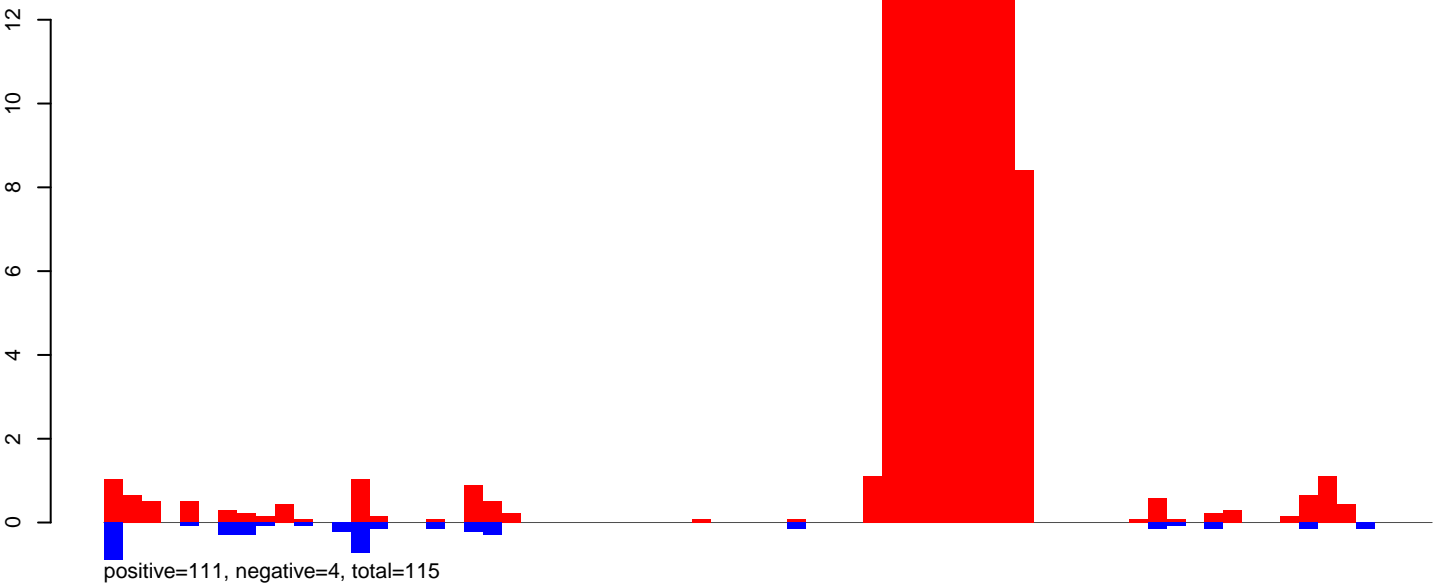
CuQuin_Testes_NL.rep



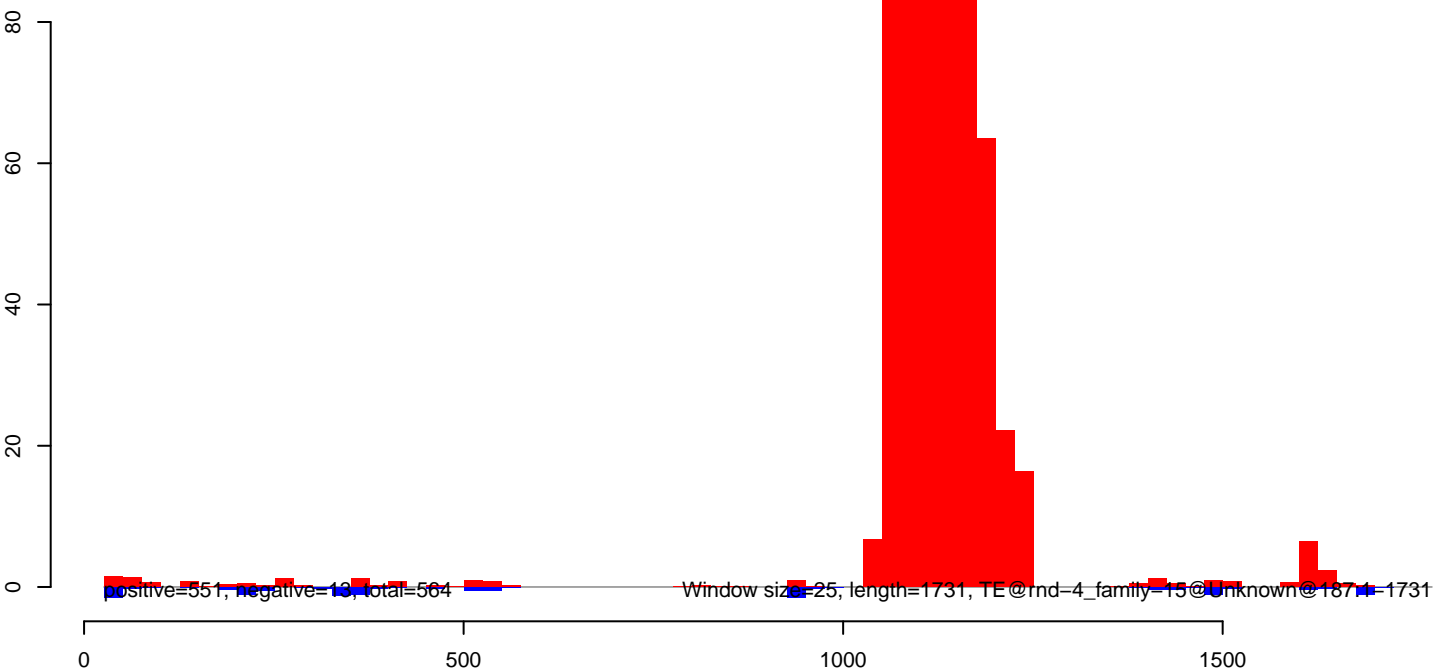
CuQuin_Testes_NL.18_23.rep



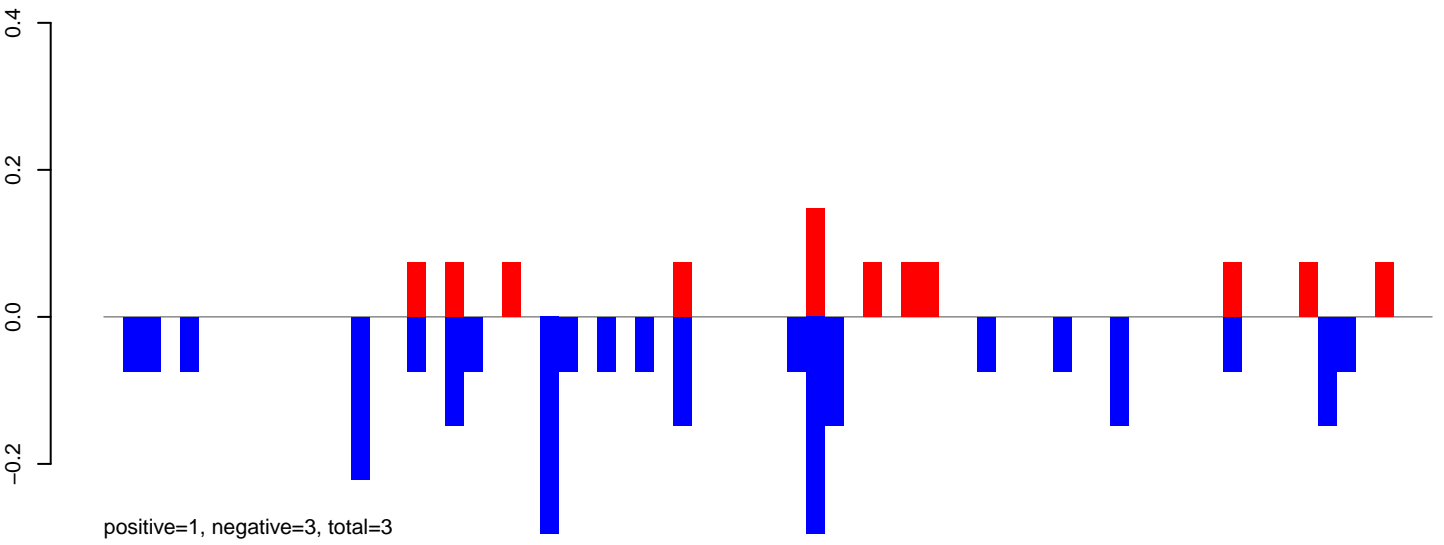
CuQuin_Testes_NL.24_35.rep



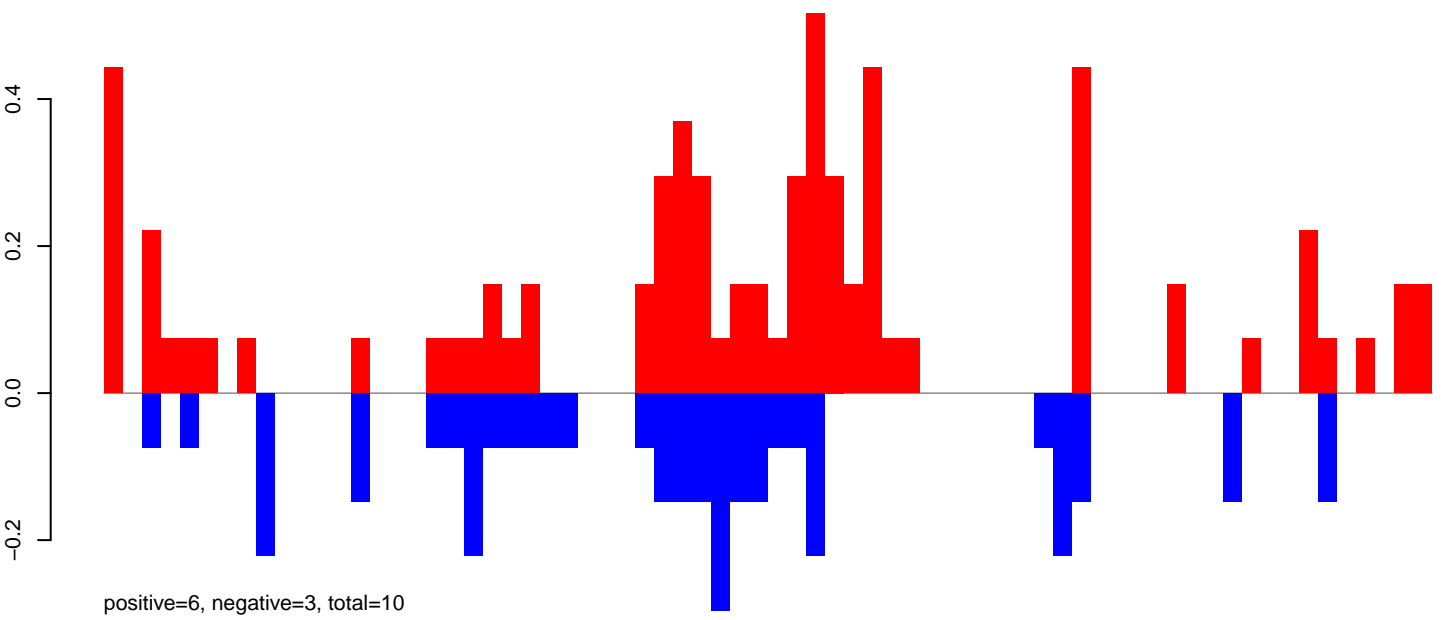
CuQuin_Testes_NL.rep



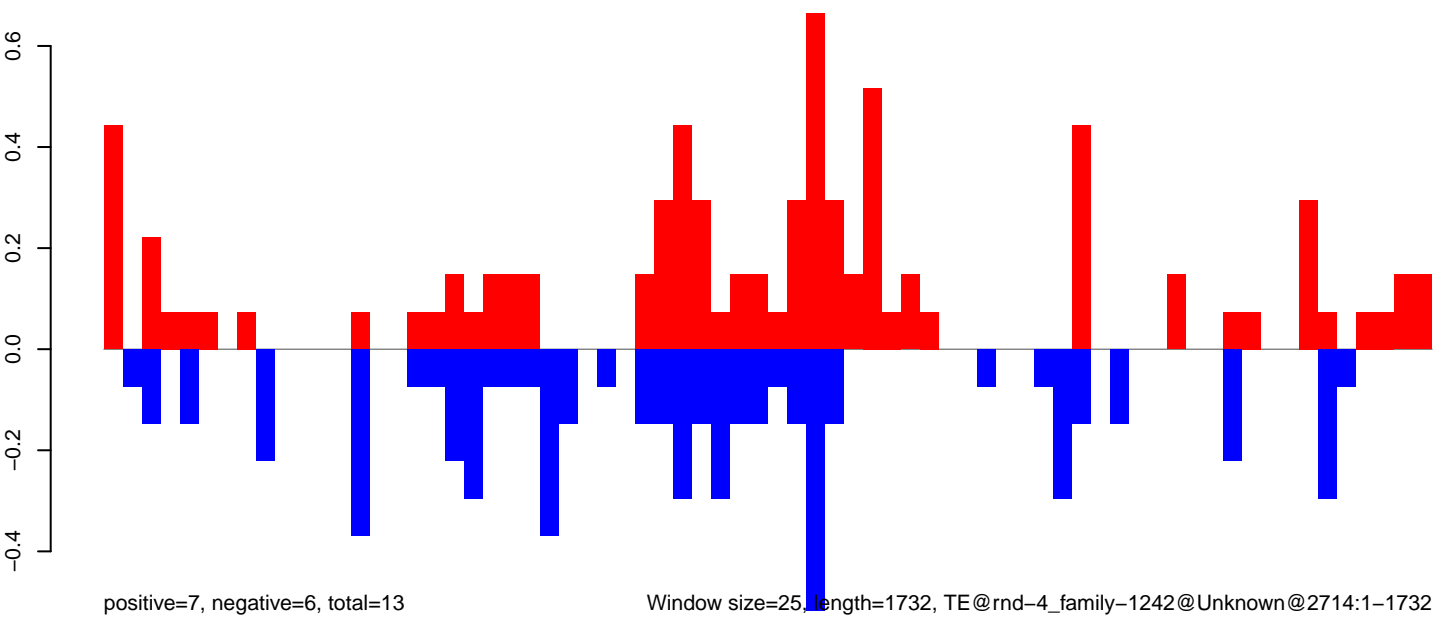
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



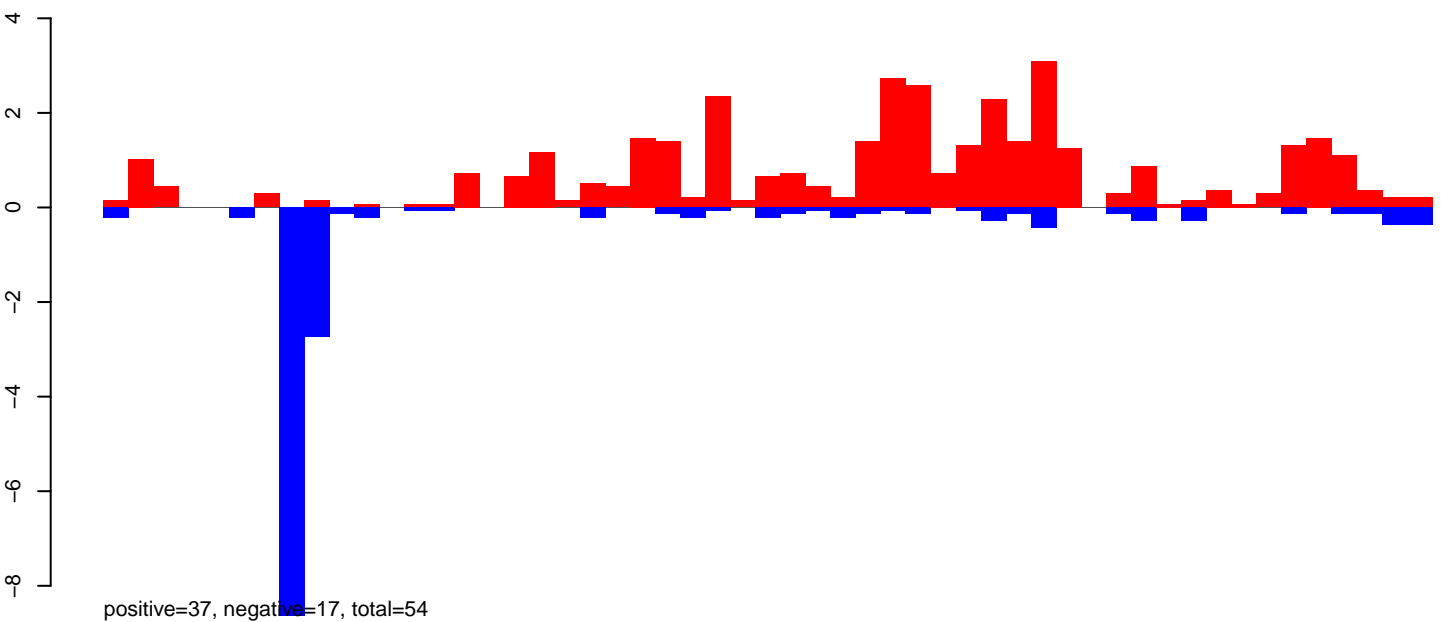
0

500

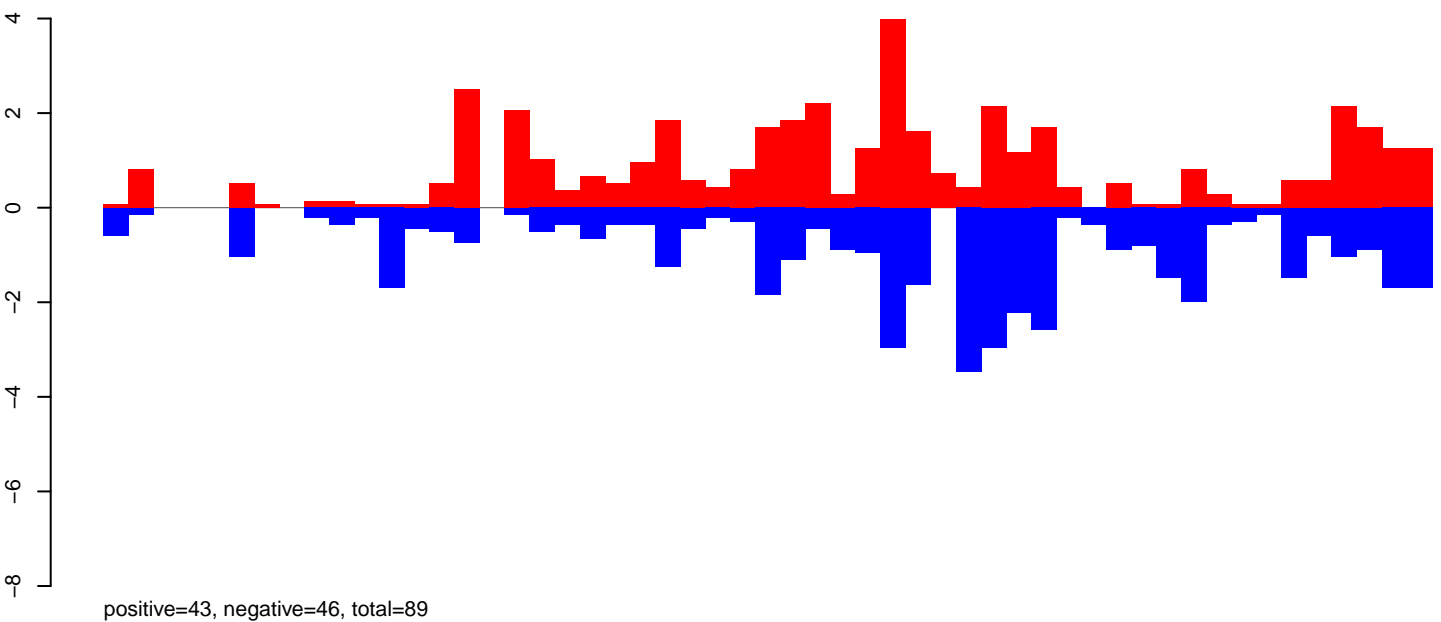
1000

1500

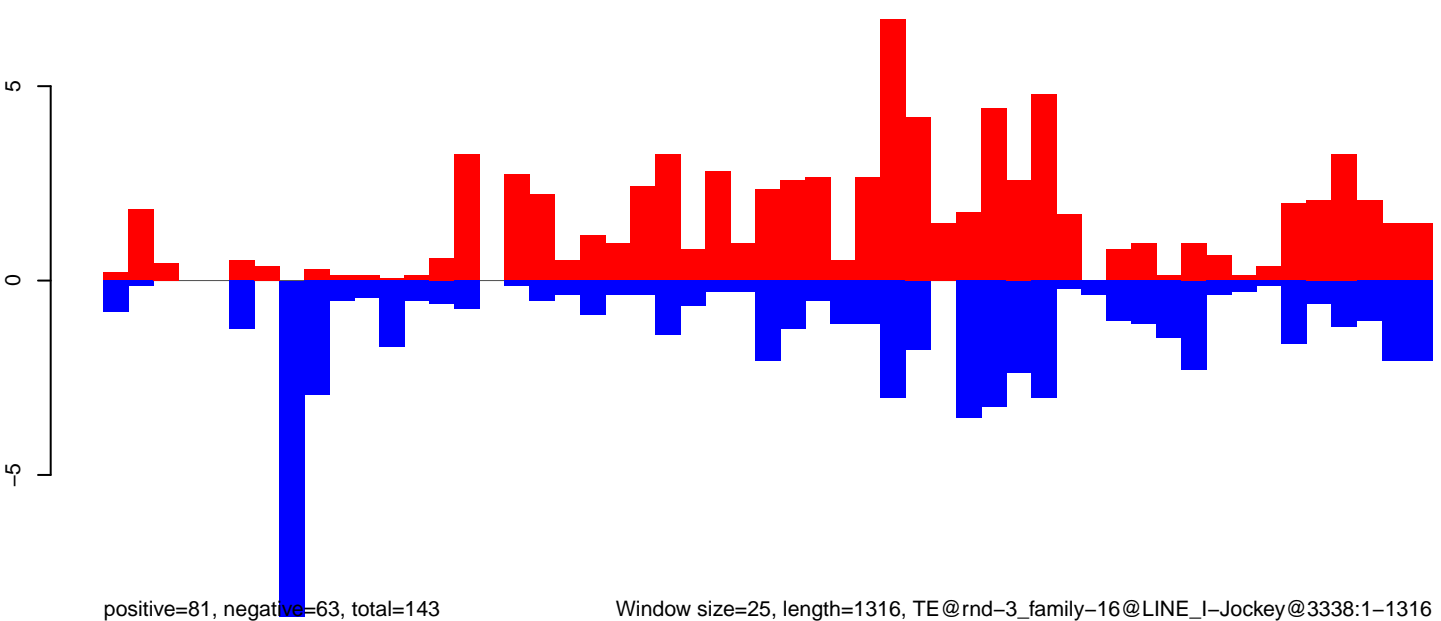
CuQuin_Testes_NL.18_23.rep



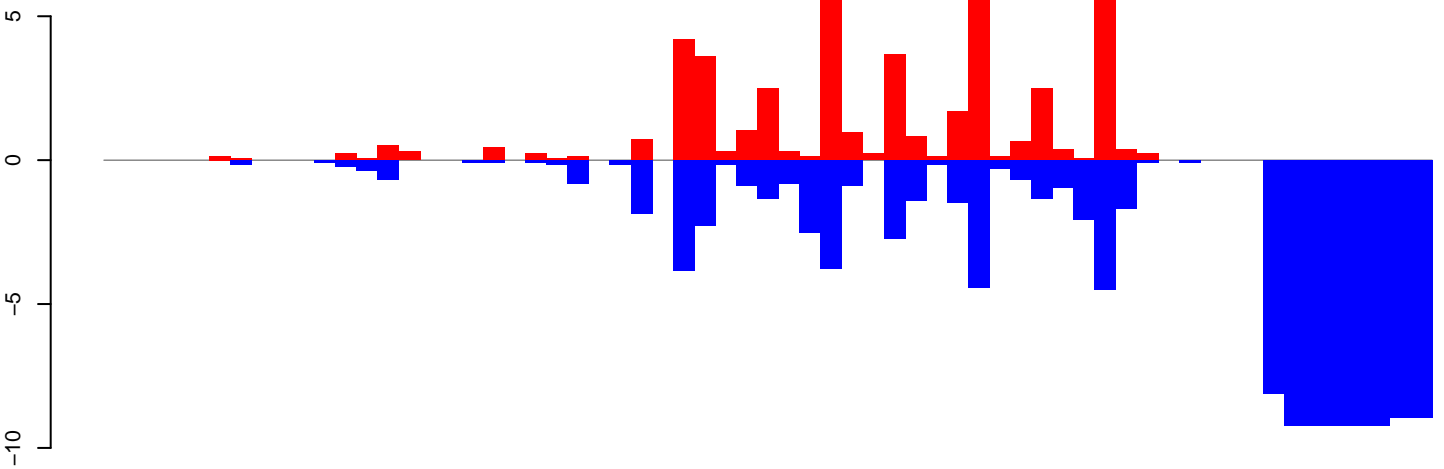
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

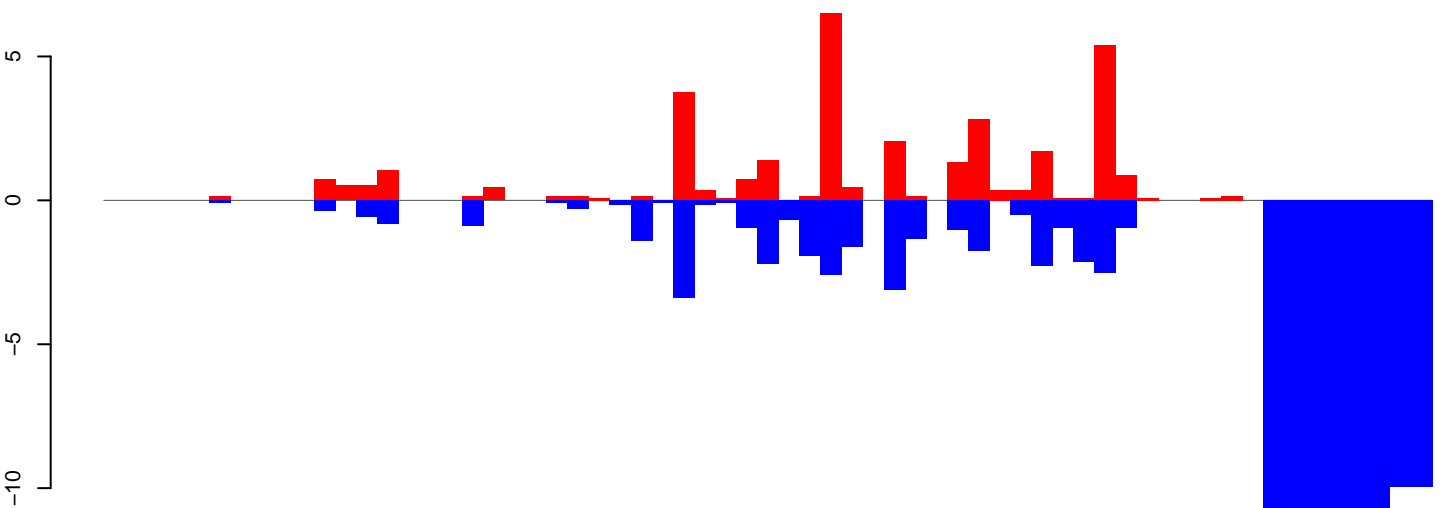


CuQuin_Testes_NL.18_23.rep



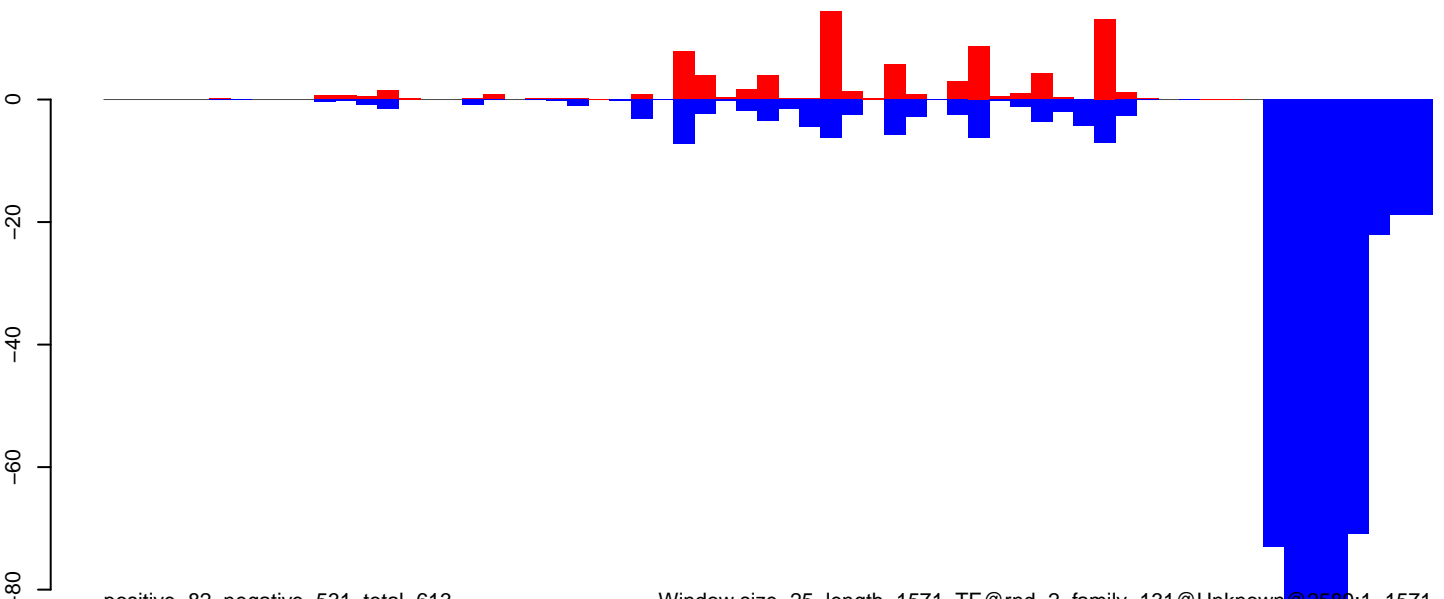
positive=49, negative=115, total=164

CuQuin_Testes_NL.24_35.rep



positive=33, negative=131, total=164

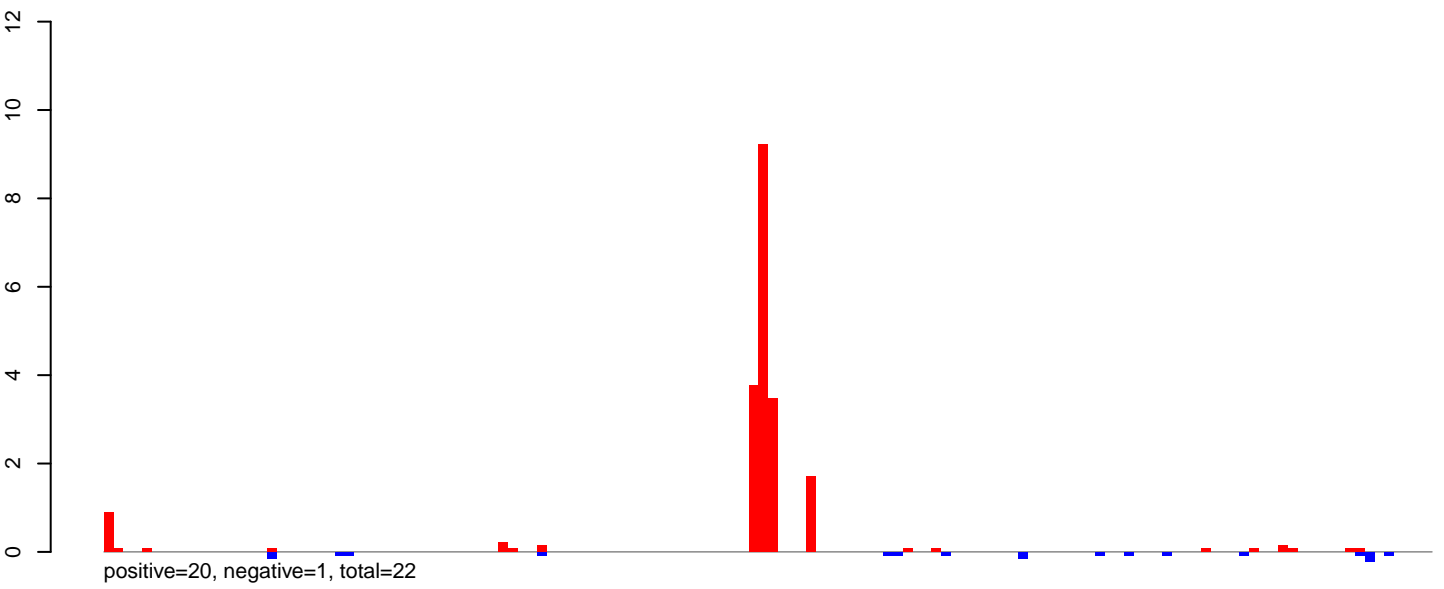
CuQuin_Testes_NL.rep



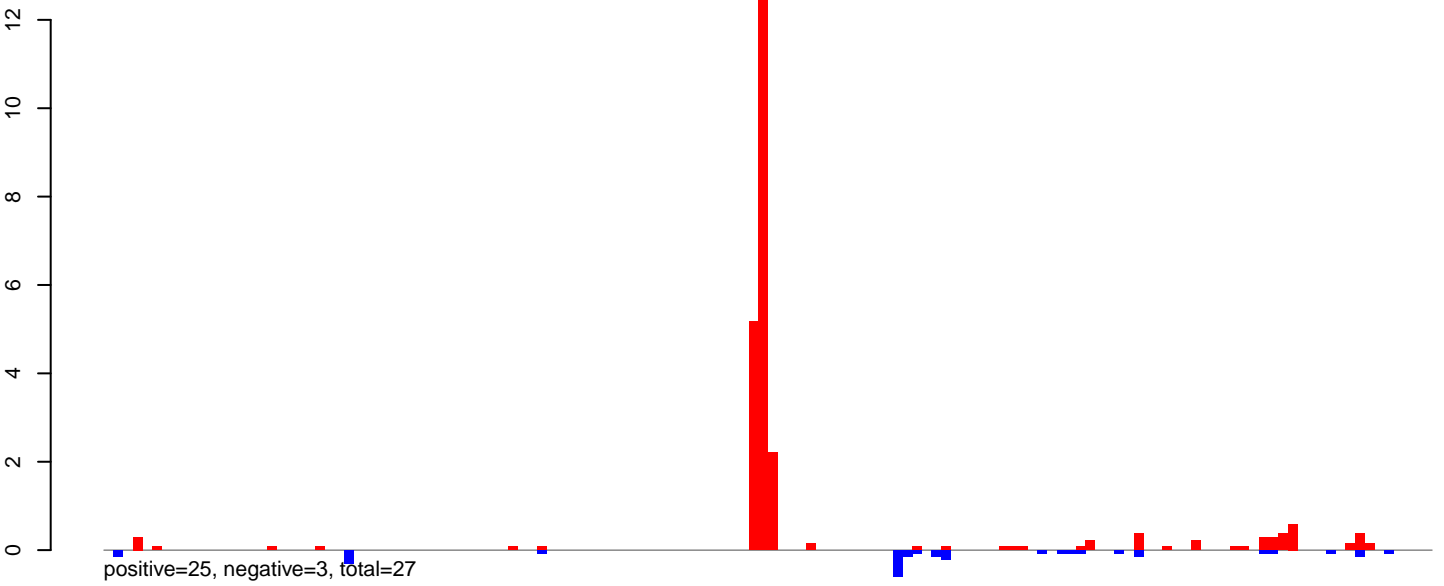
positive=82, negative=531, total=613

Window size=25, length=1571, TE@rnd-2_family-131@Unknown@2589:1-1571

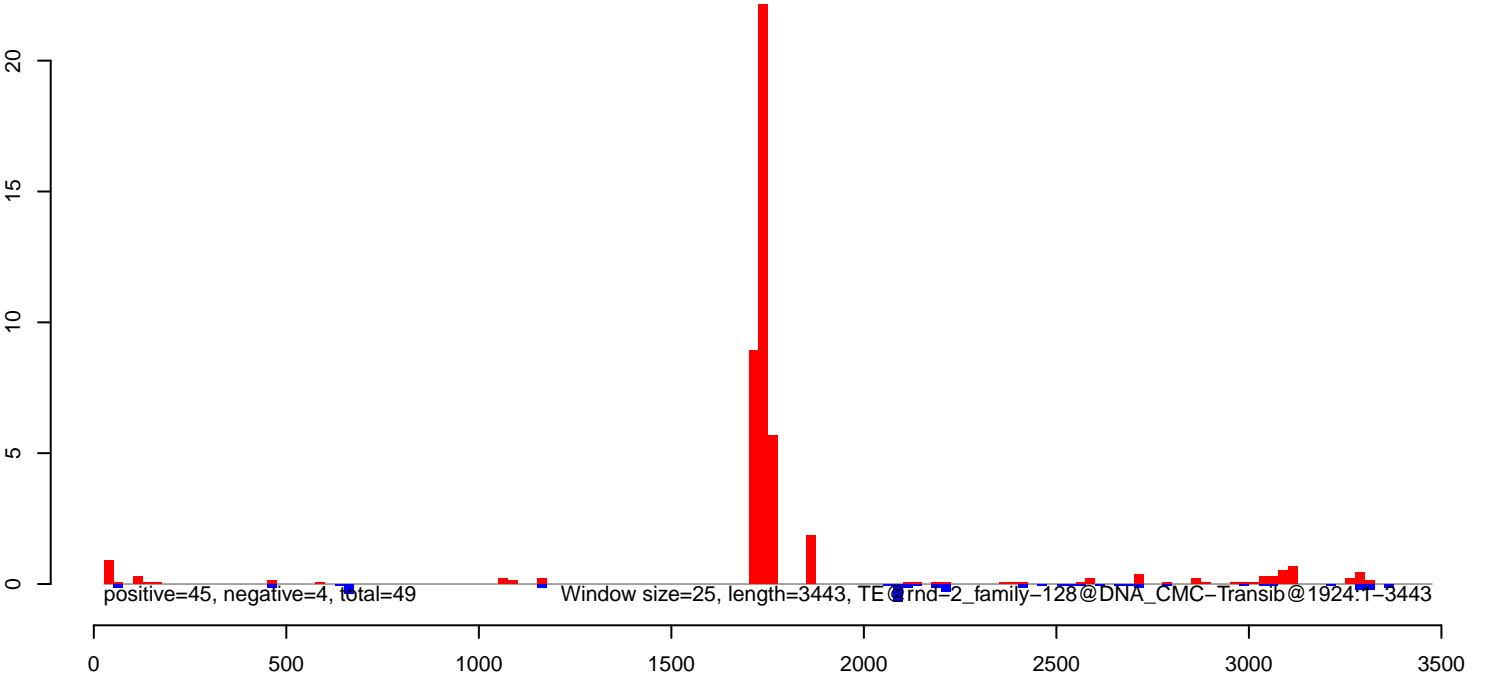
CuQuin_Testes_NL.18_23.rep



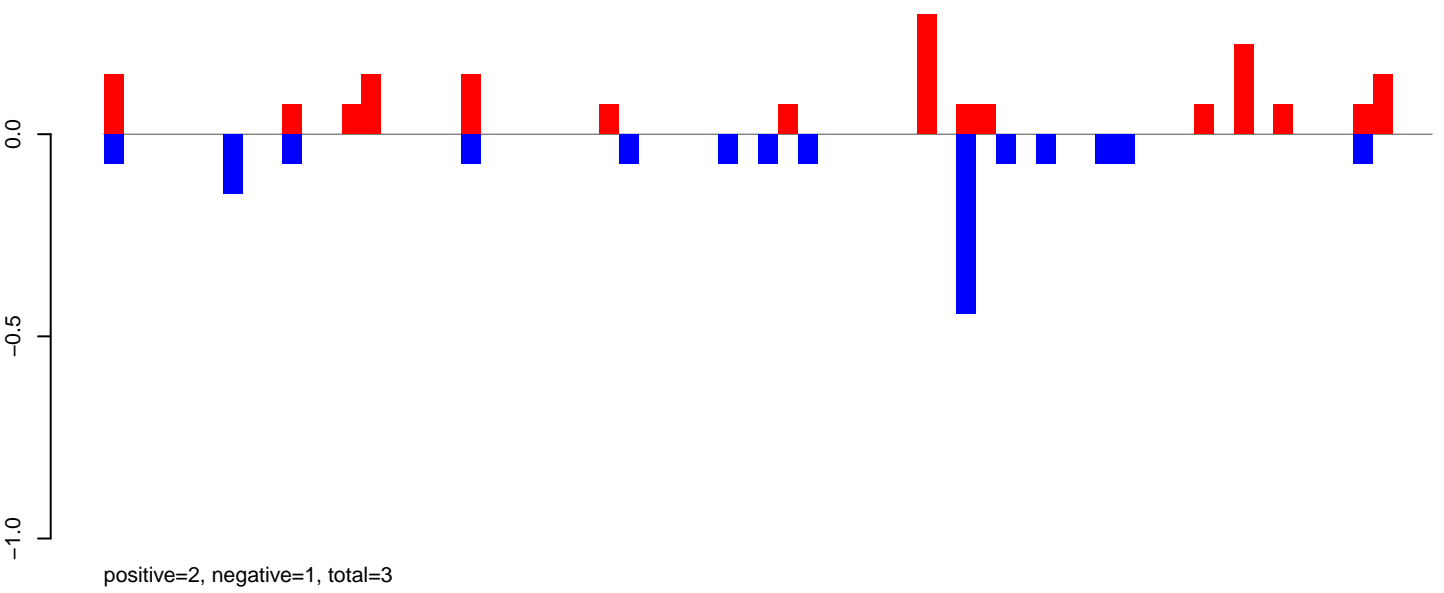
CuQuin_Testes_NL.24_35.rep



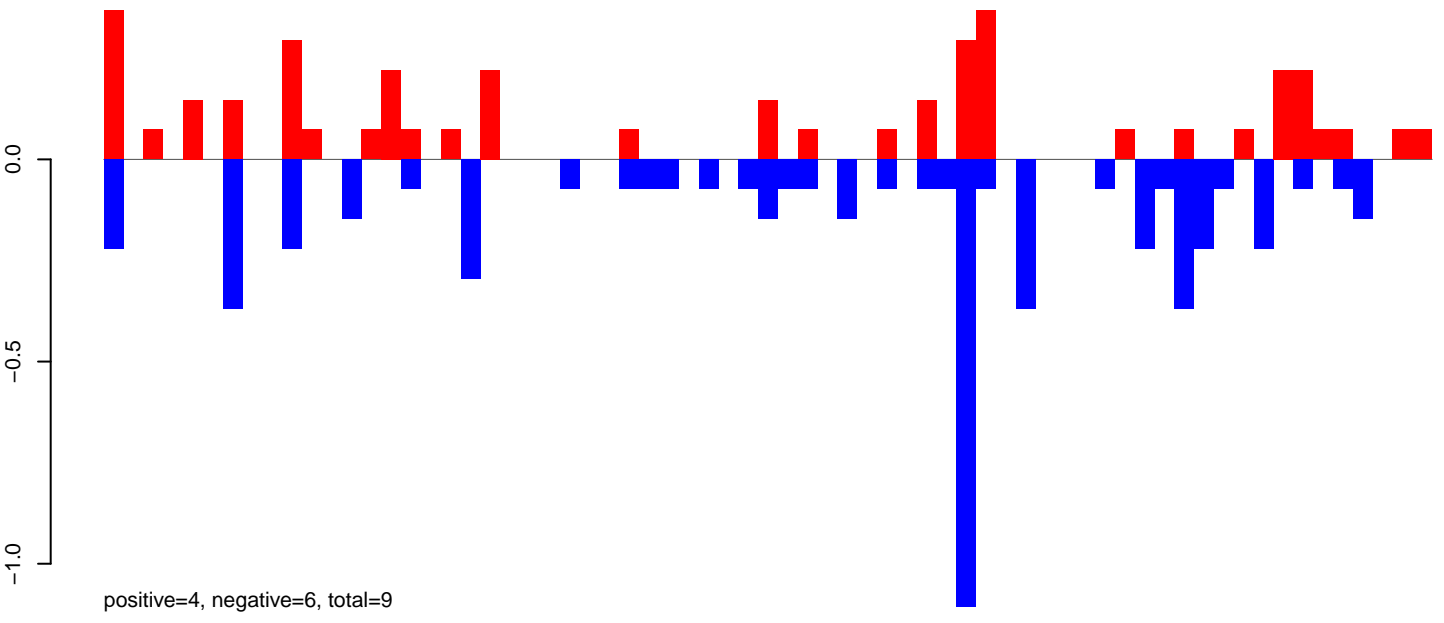
CuQuin_Testes_NL.rep



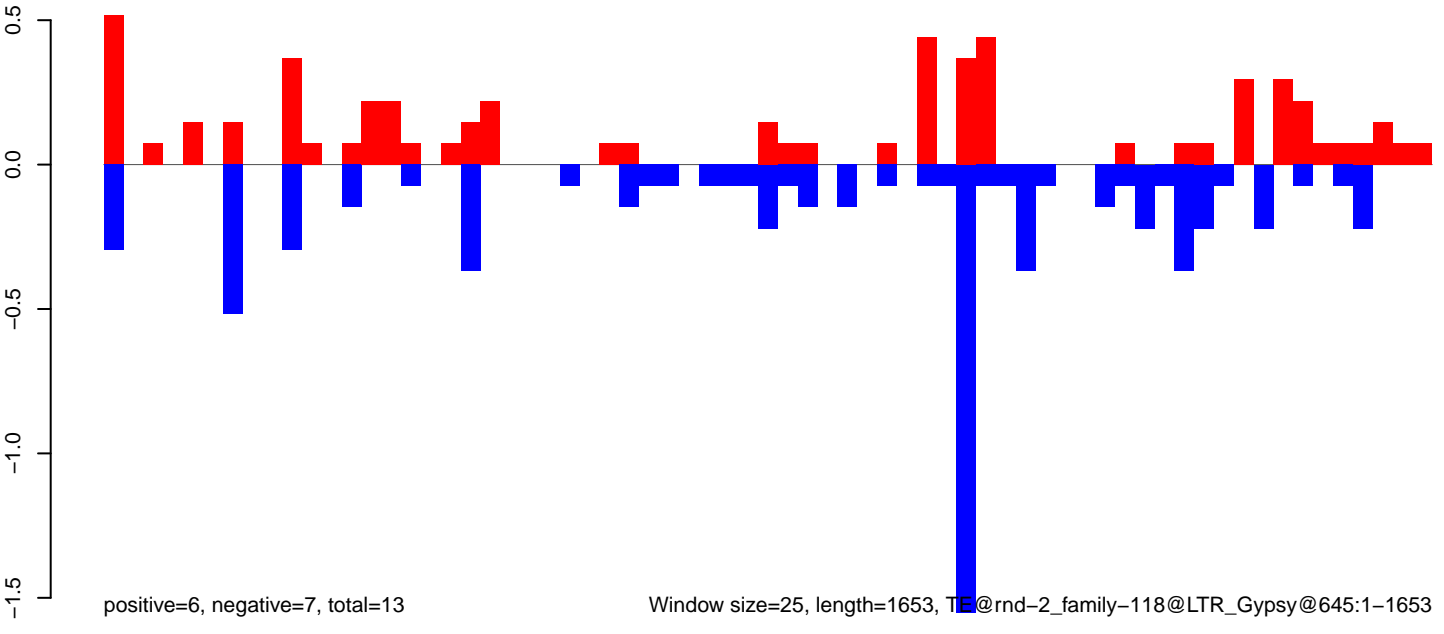
CuQuin_Testes_NL.18_23.rep



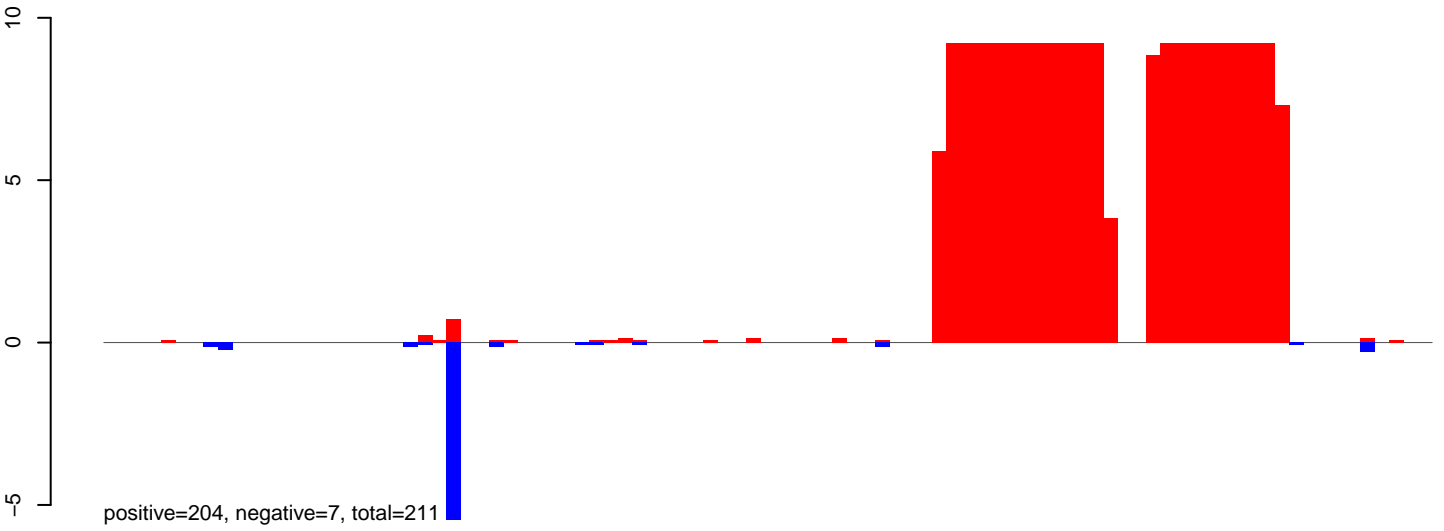
CuQuin_Testes_NL.24_35.rep



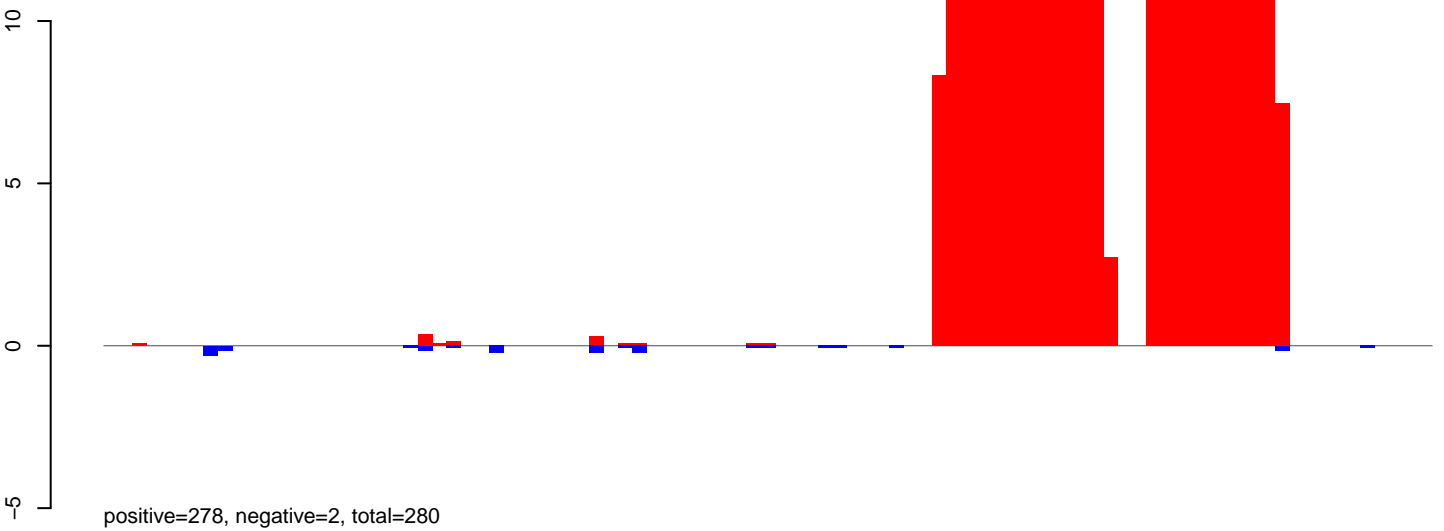
CuQuin_Testes_NL.rep



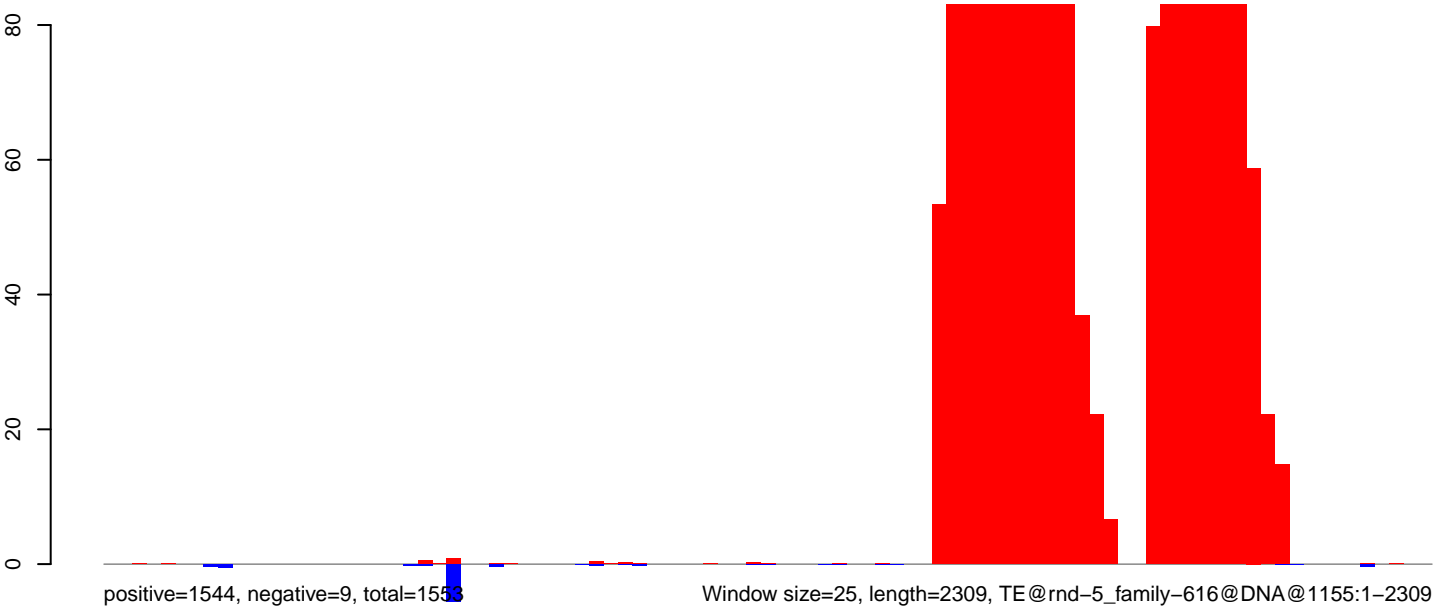
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

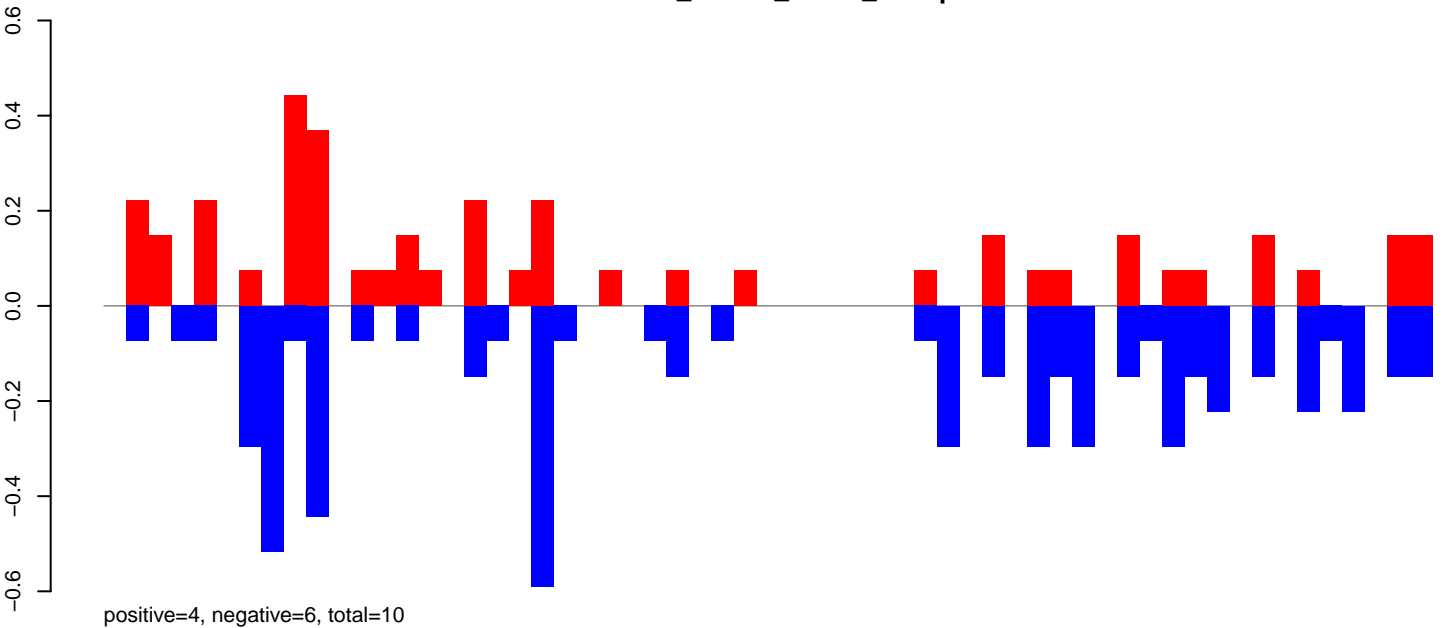


CuQuin_Testes_NL.rep

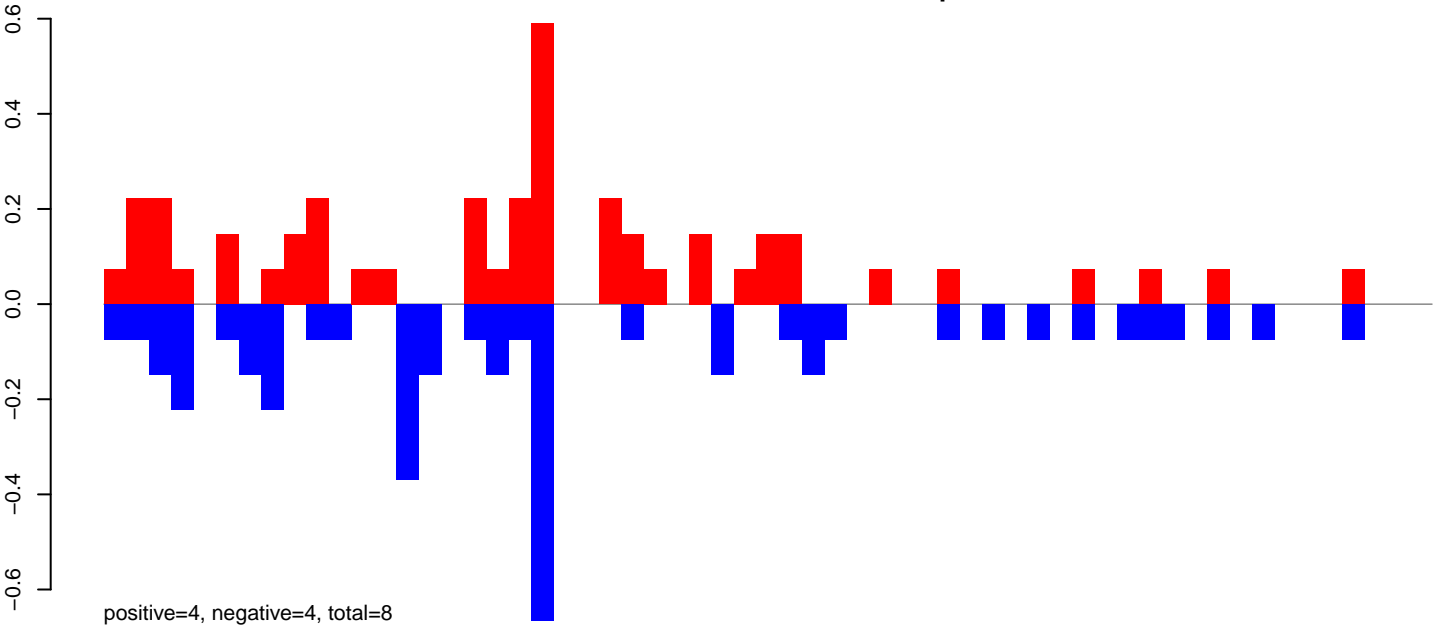


Window size=25, length=2309, TE@rnd-5_family-616@DNA@1155:1-2309

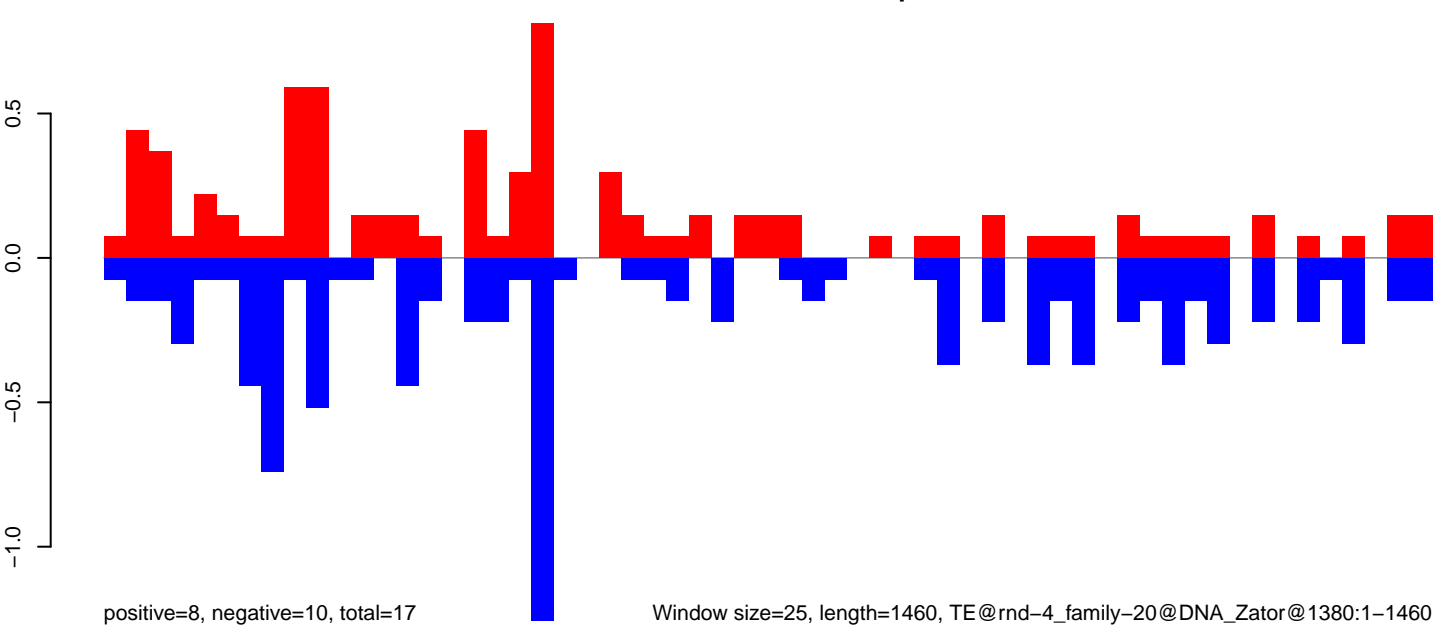
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



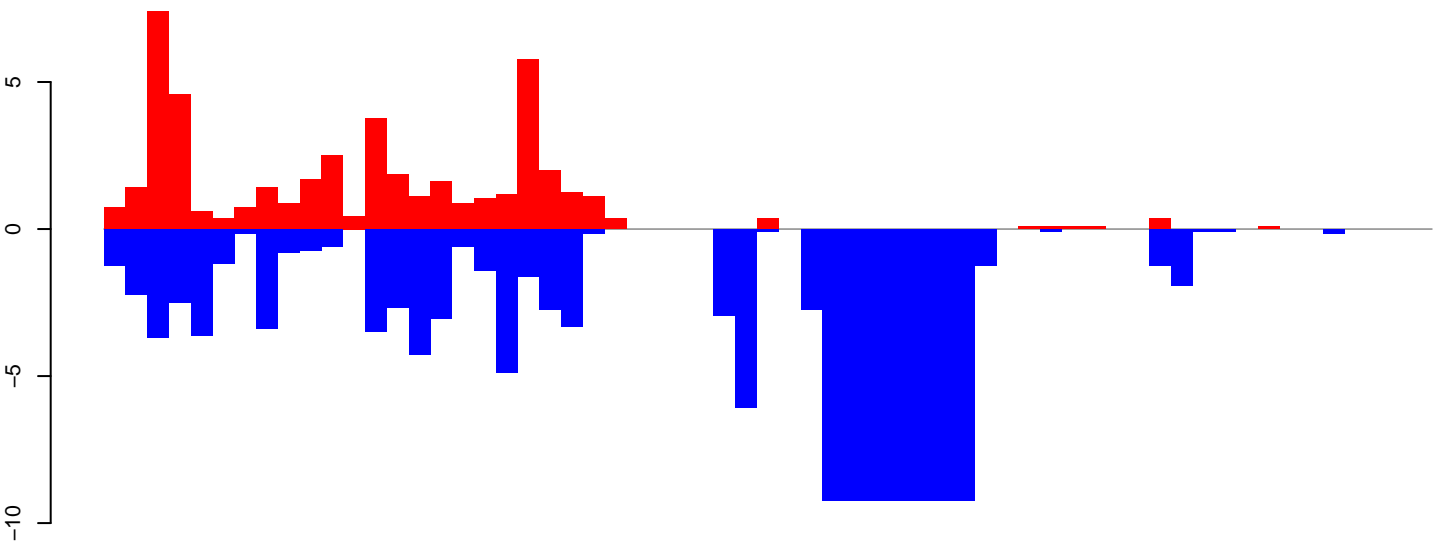
CuQuin_Testes_NL.rep



Window size=25, length=1460, TE@rnd-4_family-20@DNA_Zator@1380:1-1460

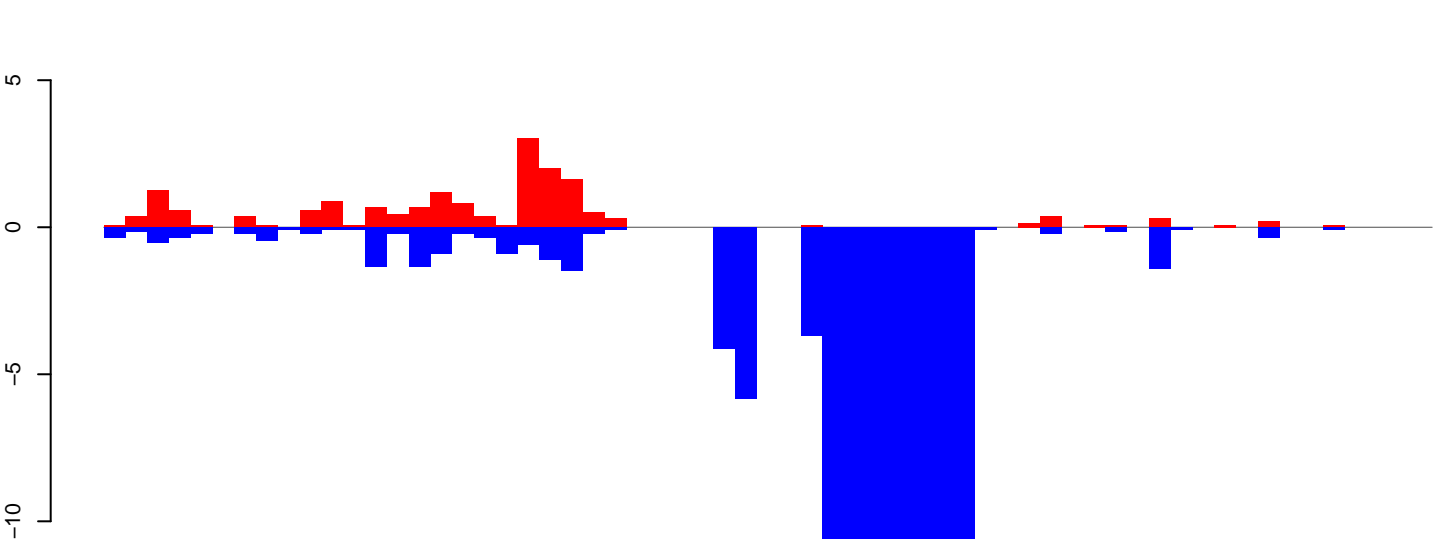
0 500 1000 1500

CuQuin_Testes_NL.18_23.rep



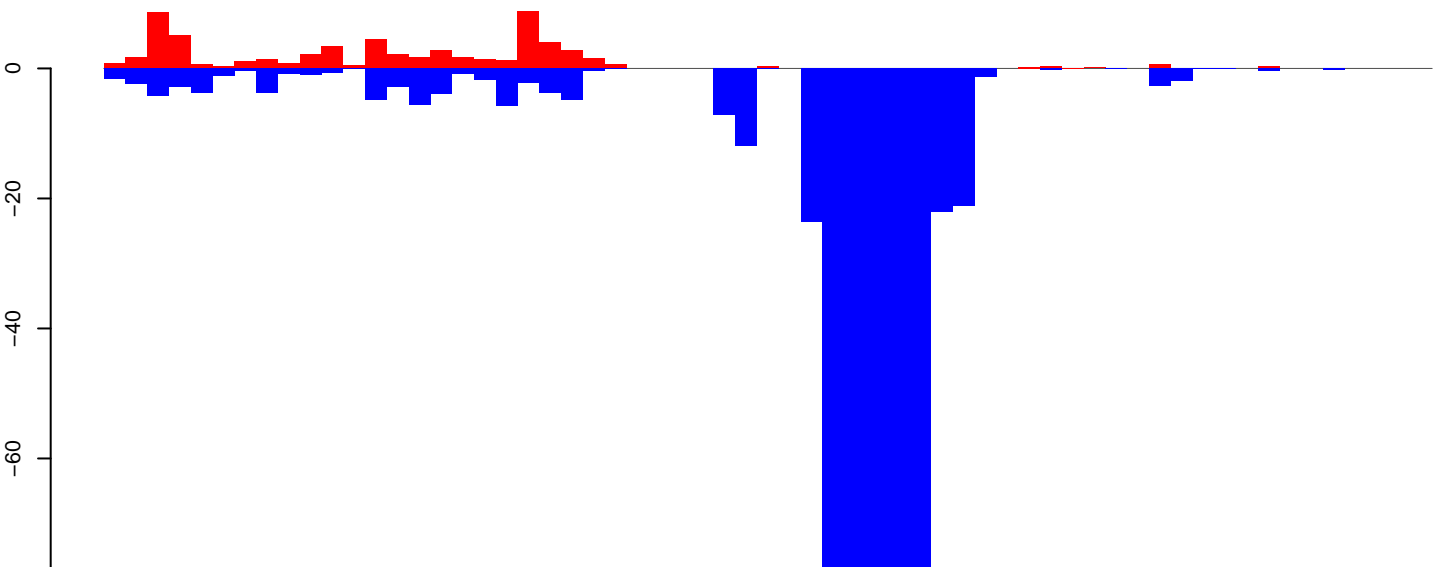
positive=46, negative=130, total=175

CuQuin_Testes_NL.24_35.rep



positive=17, negative=117, total=134

CuQuin_Testes_NL.rep

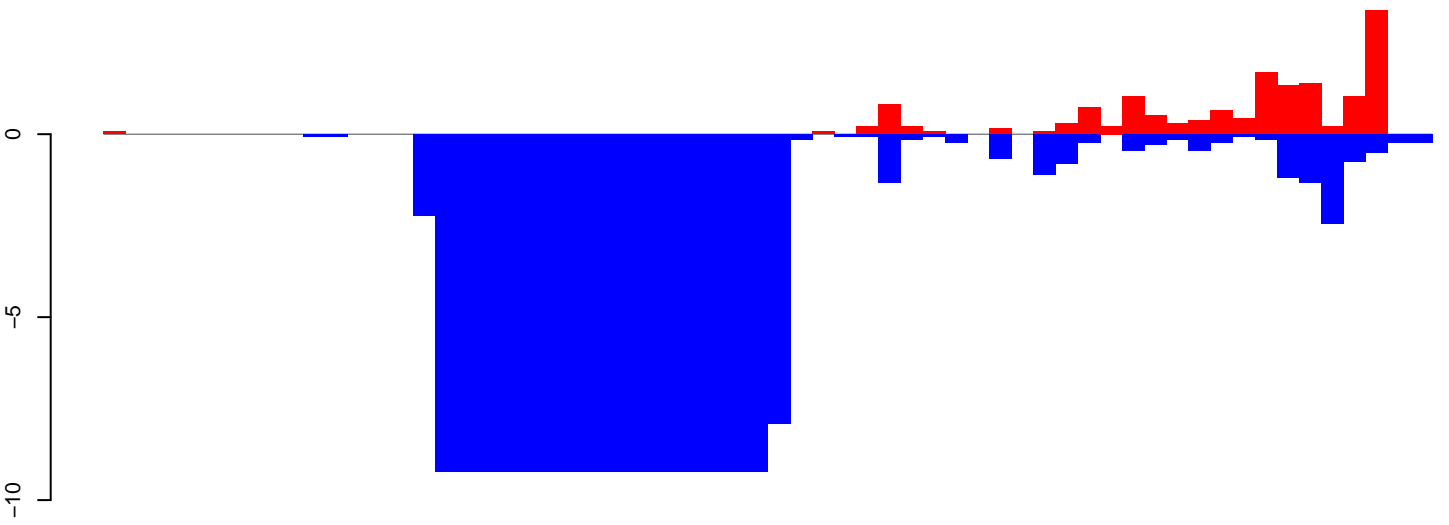


positive=63, negative=566, total=629

Window size=25, length=1524, TE@rnd-5_family-122@DNA_Sola-2@1646:1-1524

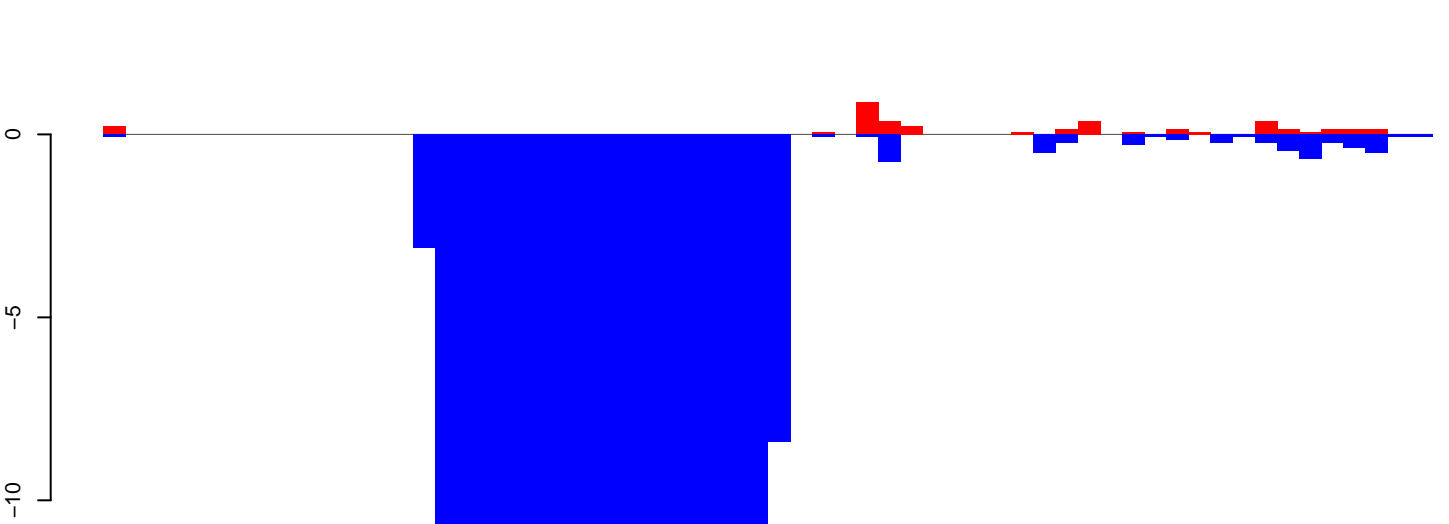
0 500 1000 1500

CuQuin_Testes_NL.18_23.rep



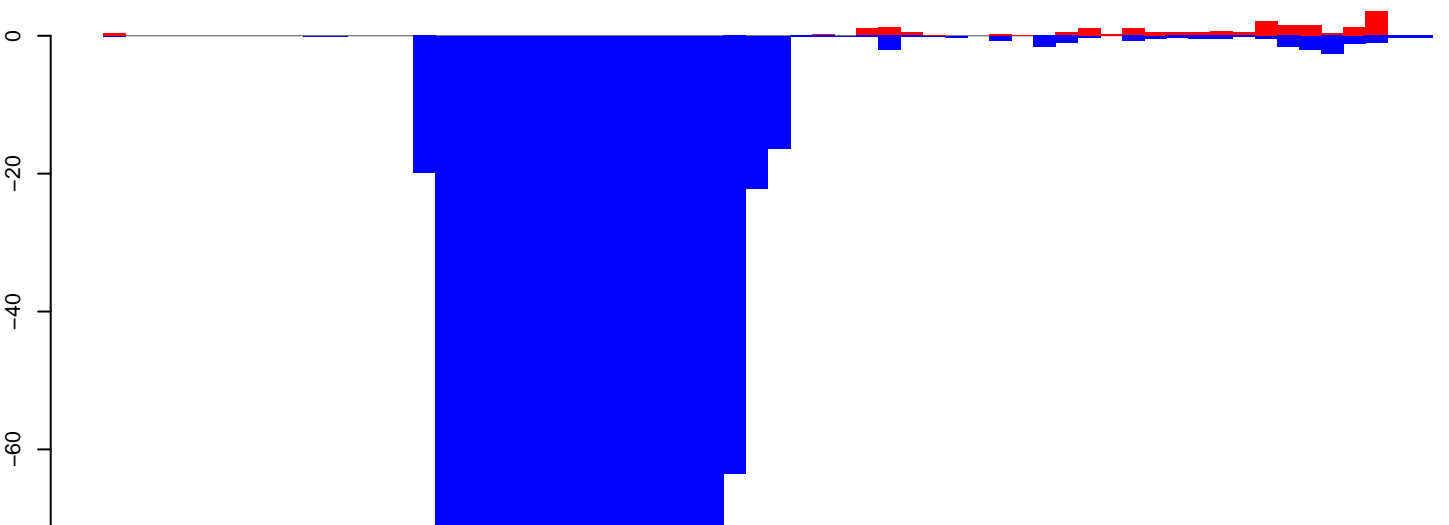
positive=15, negative=162, total=177

CuQuin_Testes_NL.24_35.rep



positive=4, negative=210, total=214

CuQuin_Testes_NL.rep

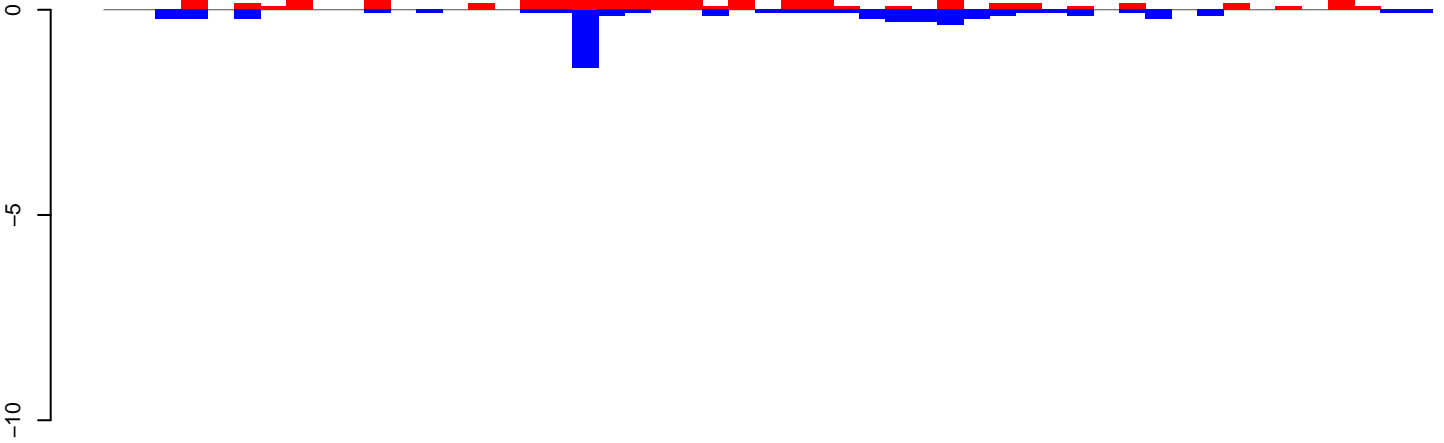


positive=19, negative=1220, total=1240

Window size=25, length=1497, TE@rnd-4_family-967@DNA_Sola-3@1564:1-1497

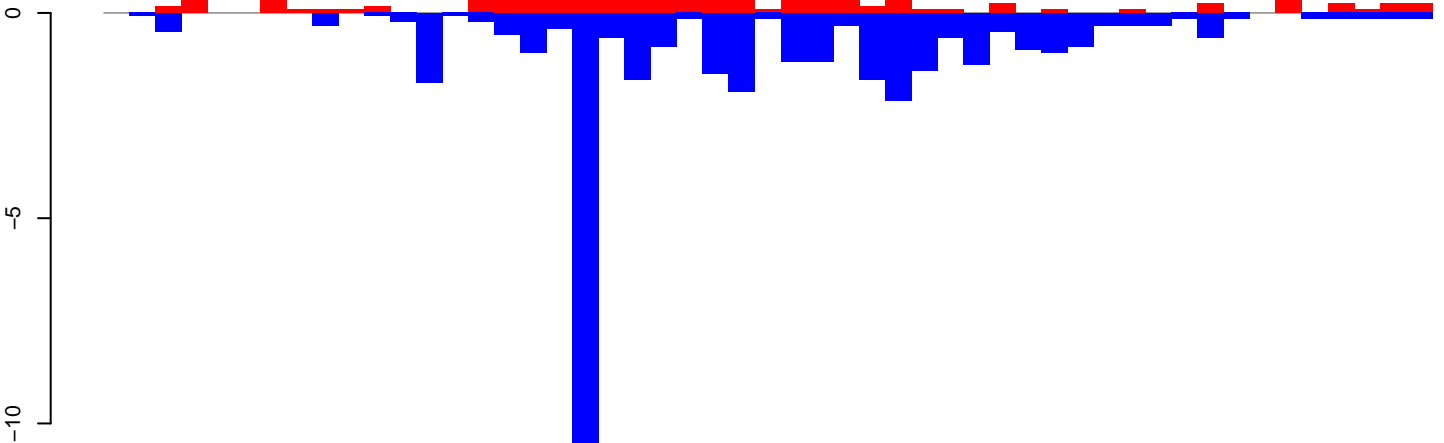
0 500 1000 1500

CuQuin_Testes_NL.18_23.rep



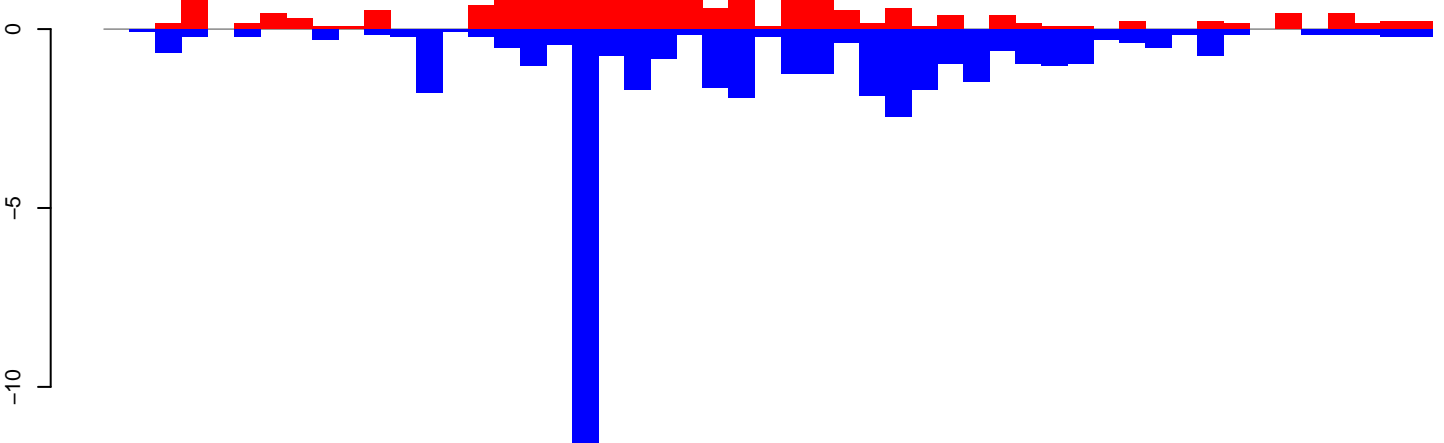
positive=11, negative=5, total=16

CuQuin_Testes_NL.24_35.rep



positive=14, negative=40, total=54

CuQuin_Testes_NL.rep

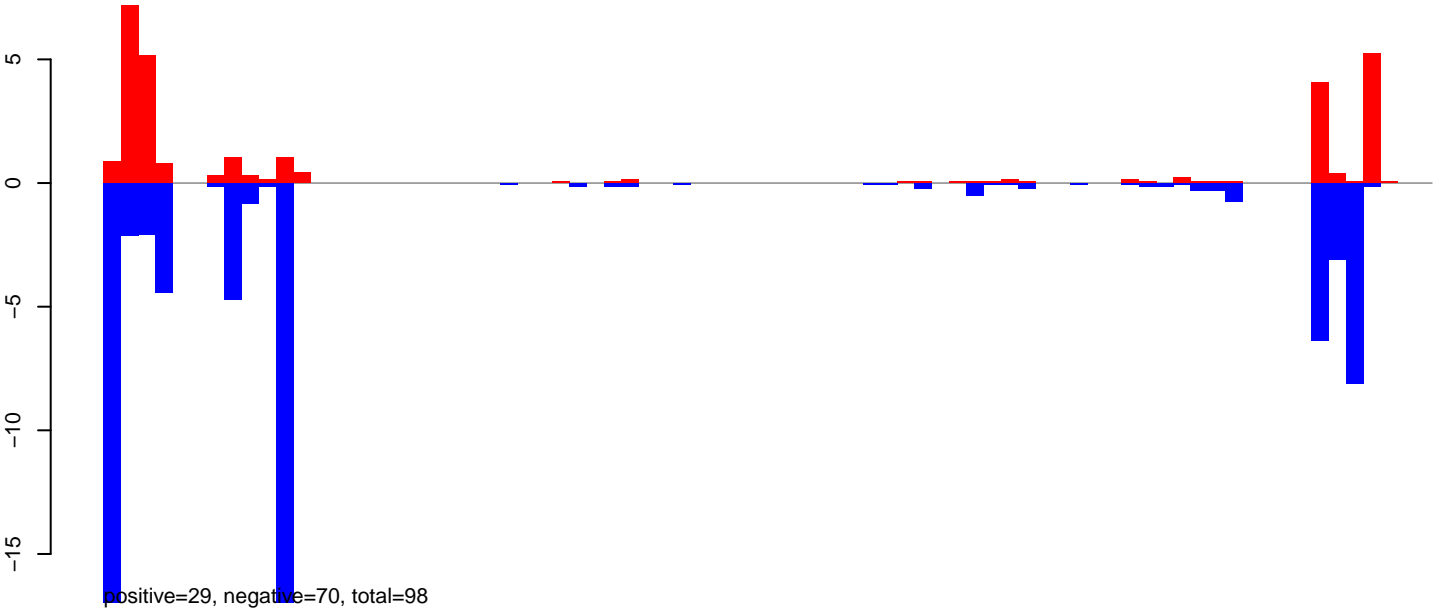


positive=25, negative=45, total=70

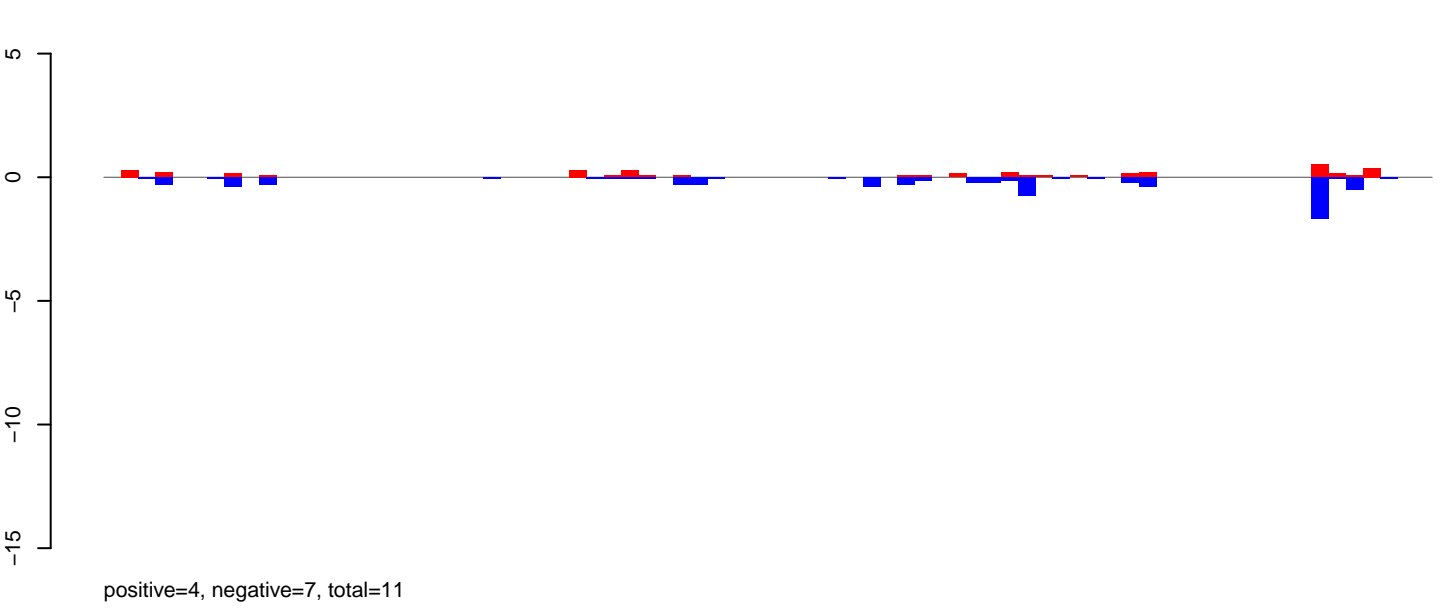
Window size=25, length=1268, TE@rnd-4_family-142@LINE_RTE-BovB@888:1-1268

0 200 400 600 800 1000 1200

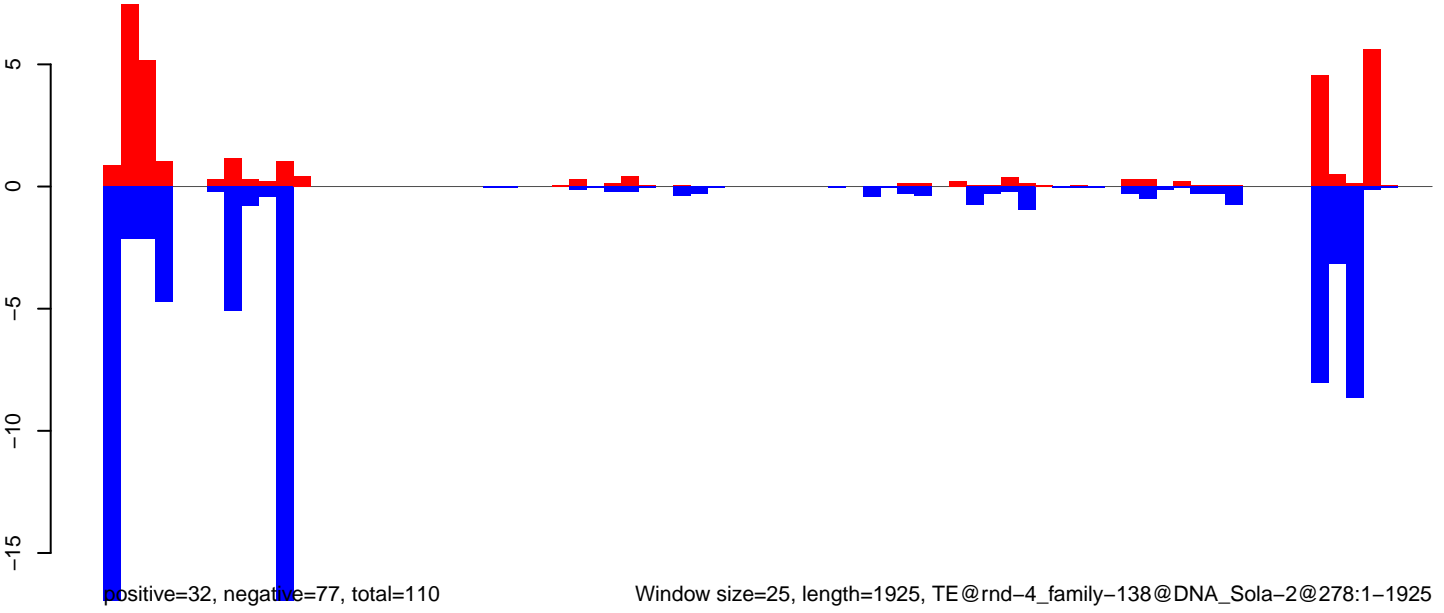
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

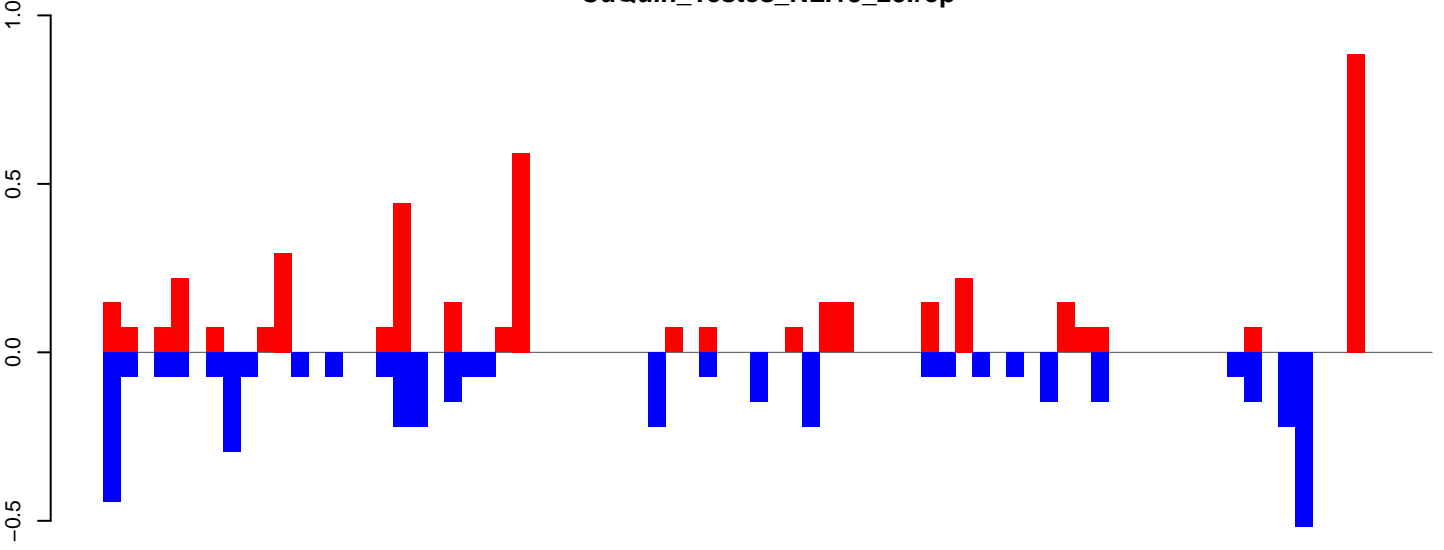


CuQuin_Testes_NL.rep



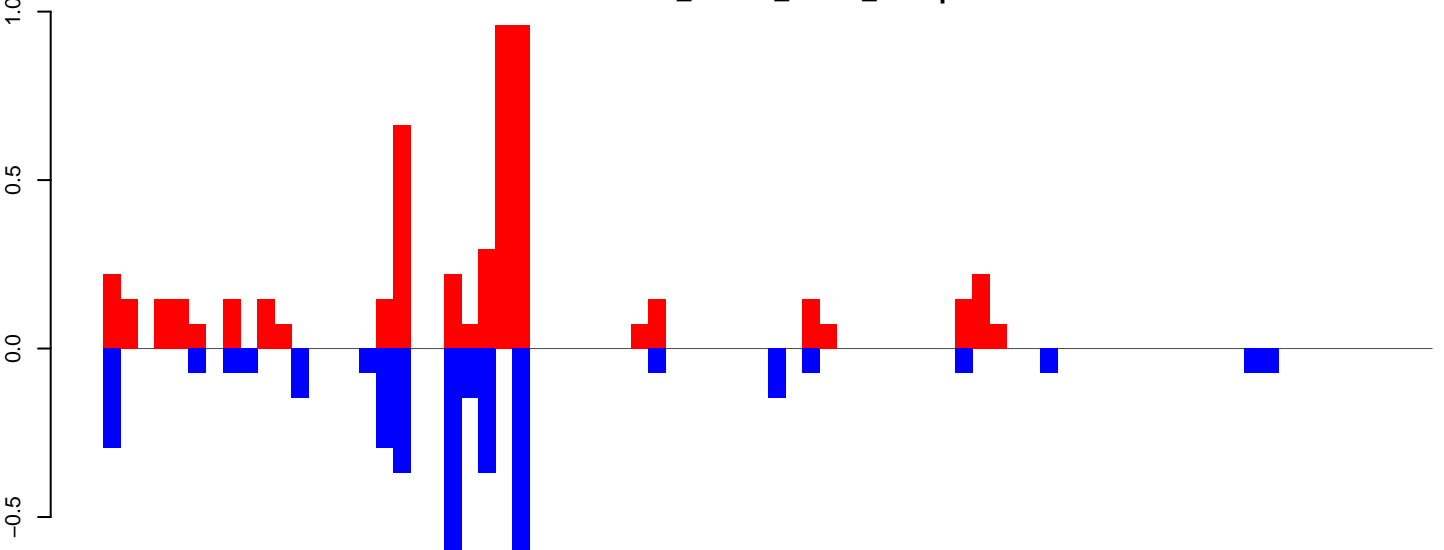
Window size=25, length=1925, TE@rnd-4_family-138@DNA_Sola-2@278:1-1925

CuQuin_Testes_NL.18_23.rep



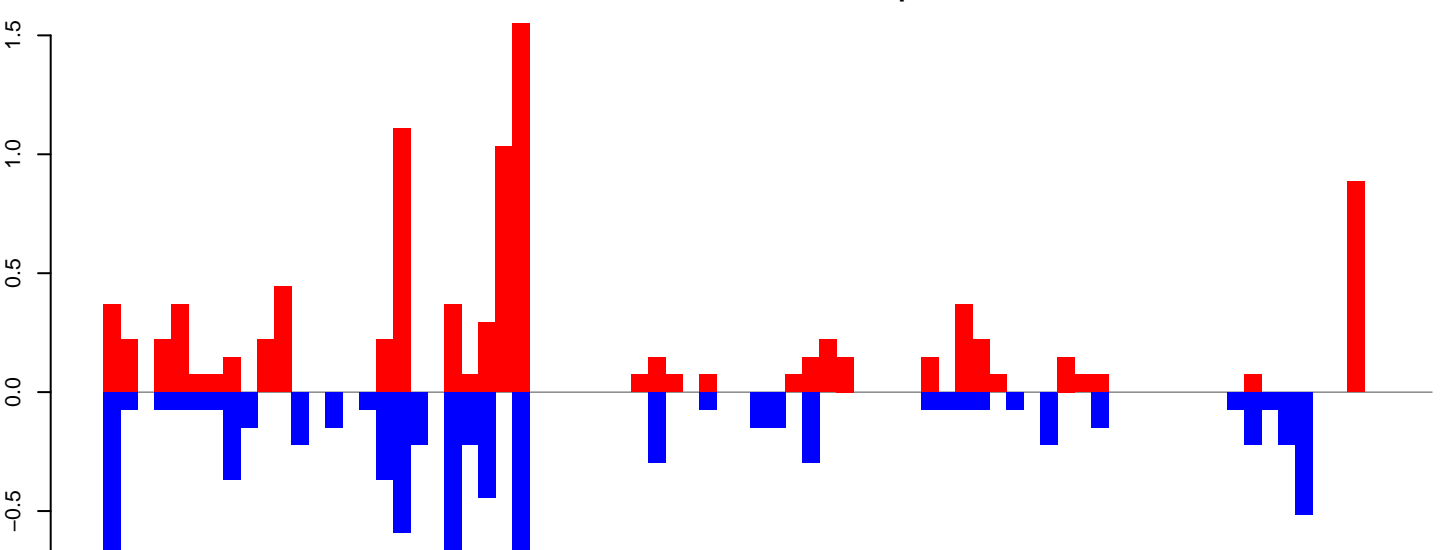
positive=4, negative=4, total=9

CuQuin_Testes_NL.24_35.rep



positive=5, negative=4, total=9

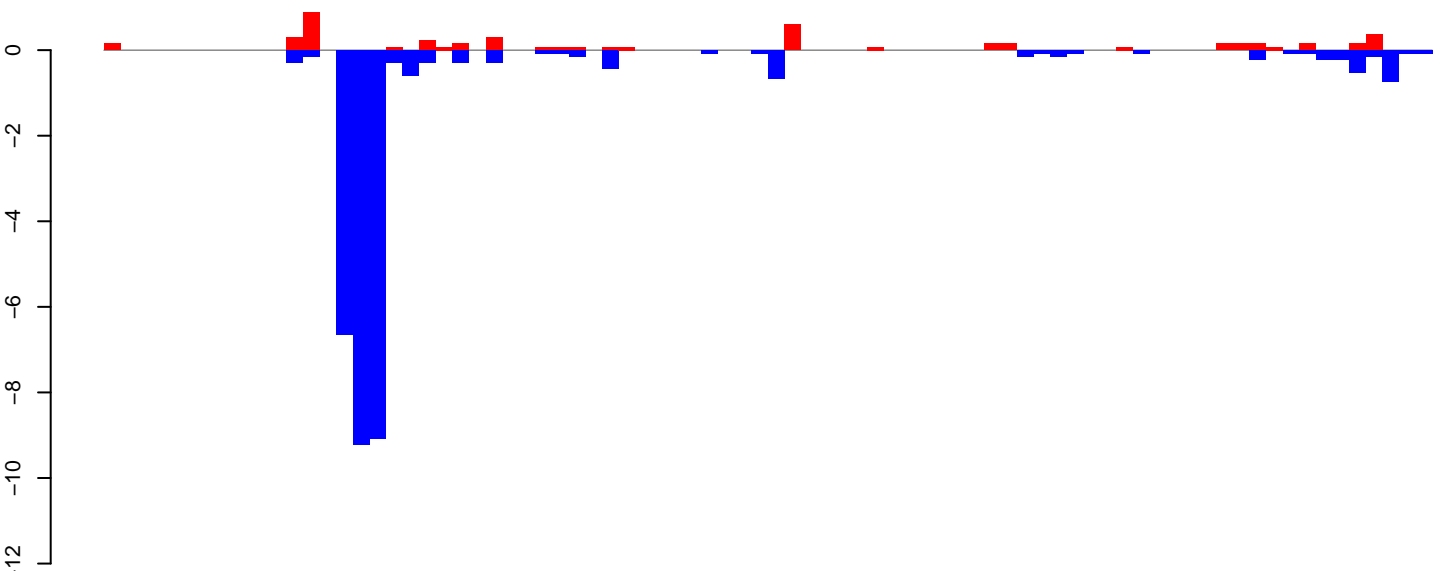
CuQuin_Testes_NL.rep



positive=10, negative=8, total=18

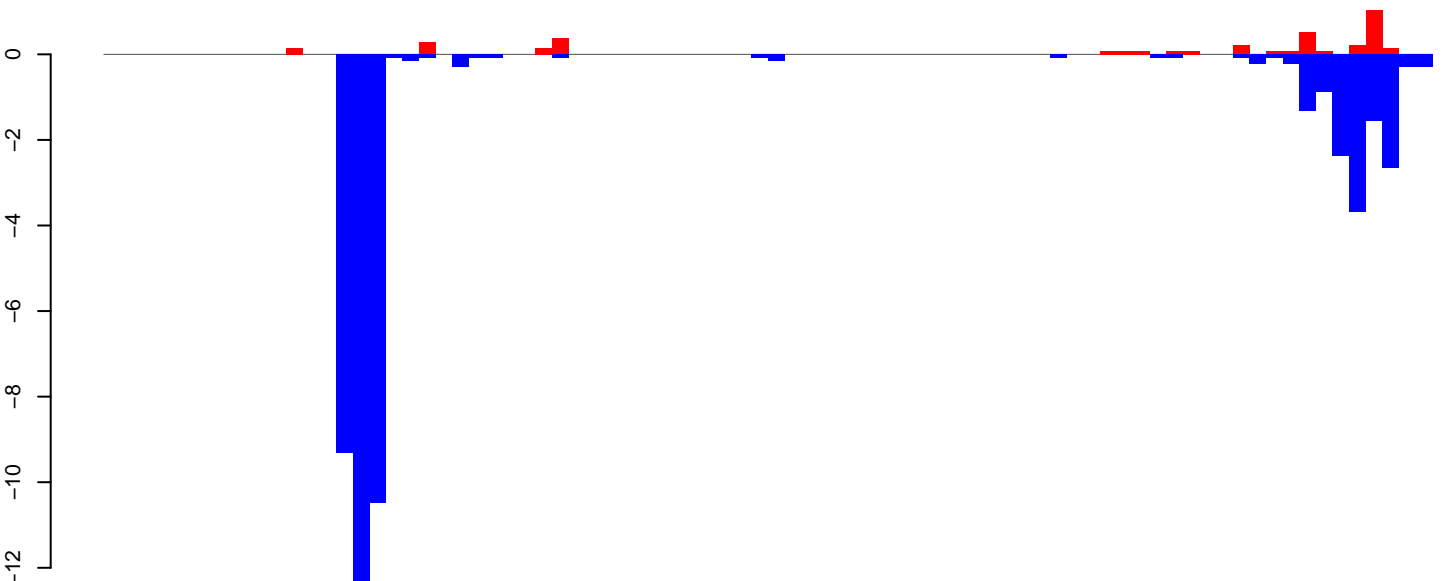
Window size=25, length=1926, TE@rnd-3_family-538@Unknown@3327:1-1926

CuQuin_Testes_NL.18_23.rep



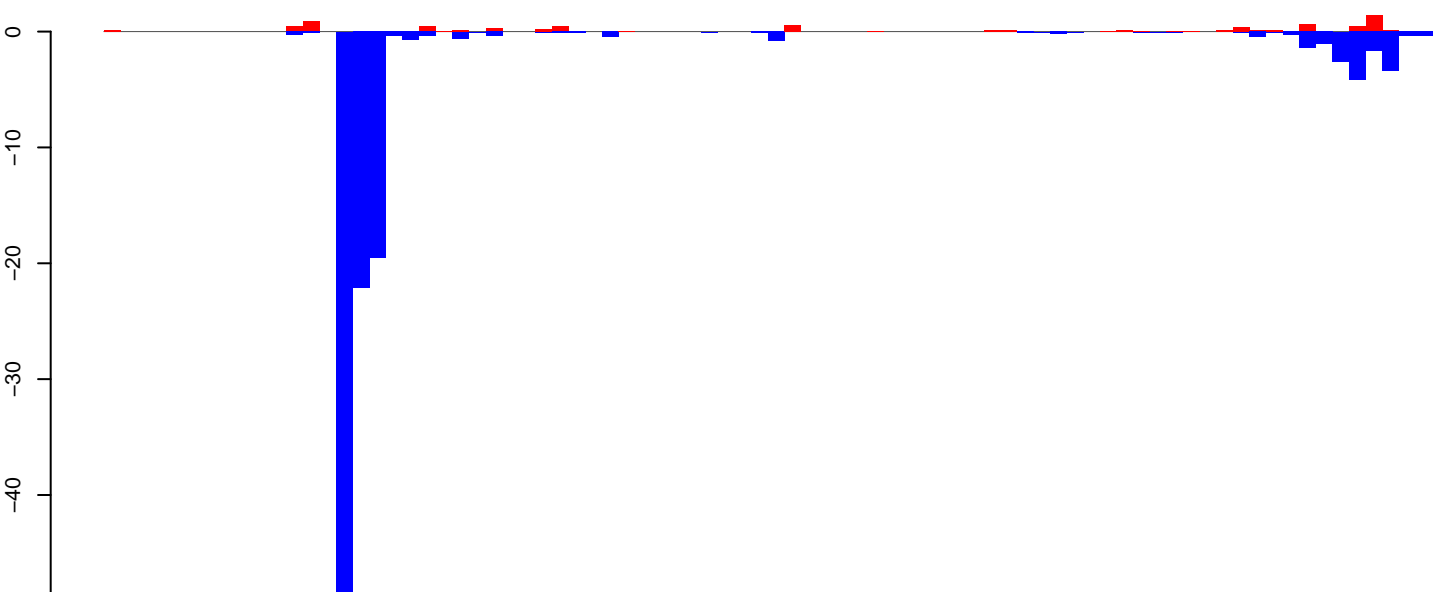
positive=5, negative=32, total=36

CuQuin_Testes_NL.24_35.rep



positive=4, negative=48, total=51

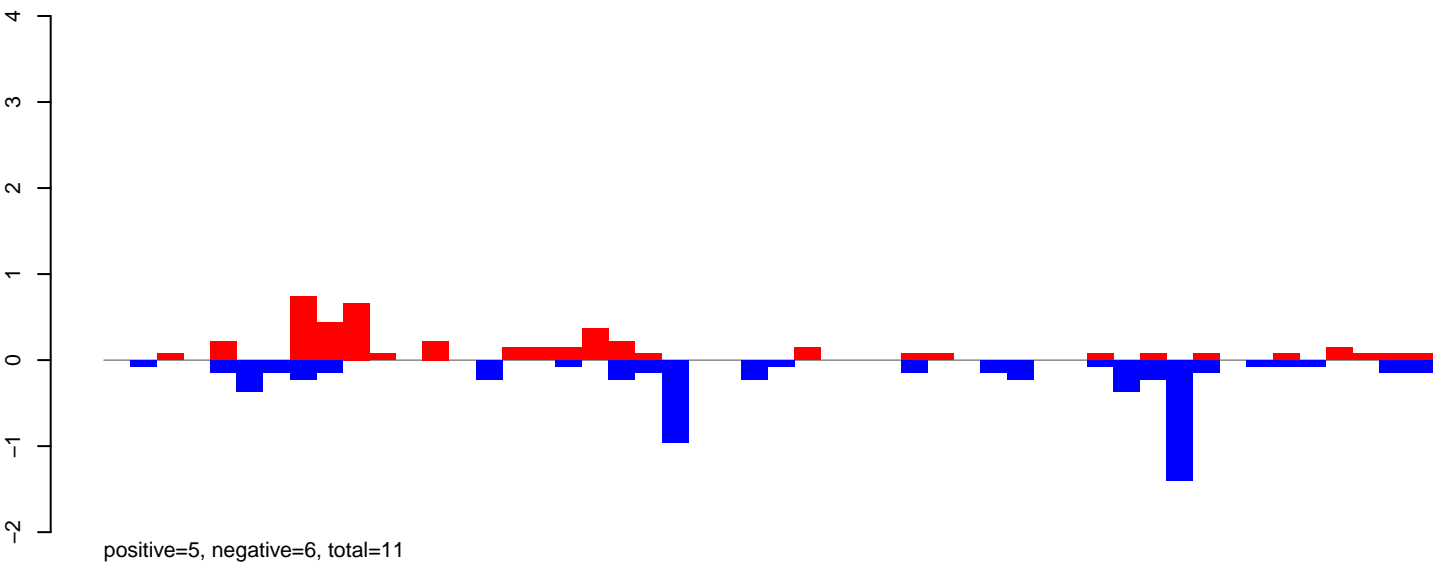
CuQuin_Testes_NL.rep



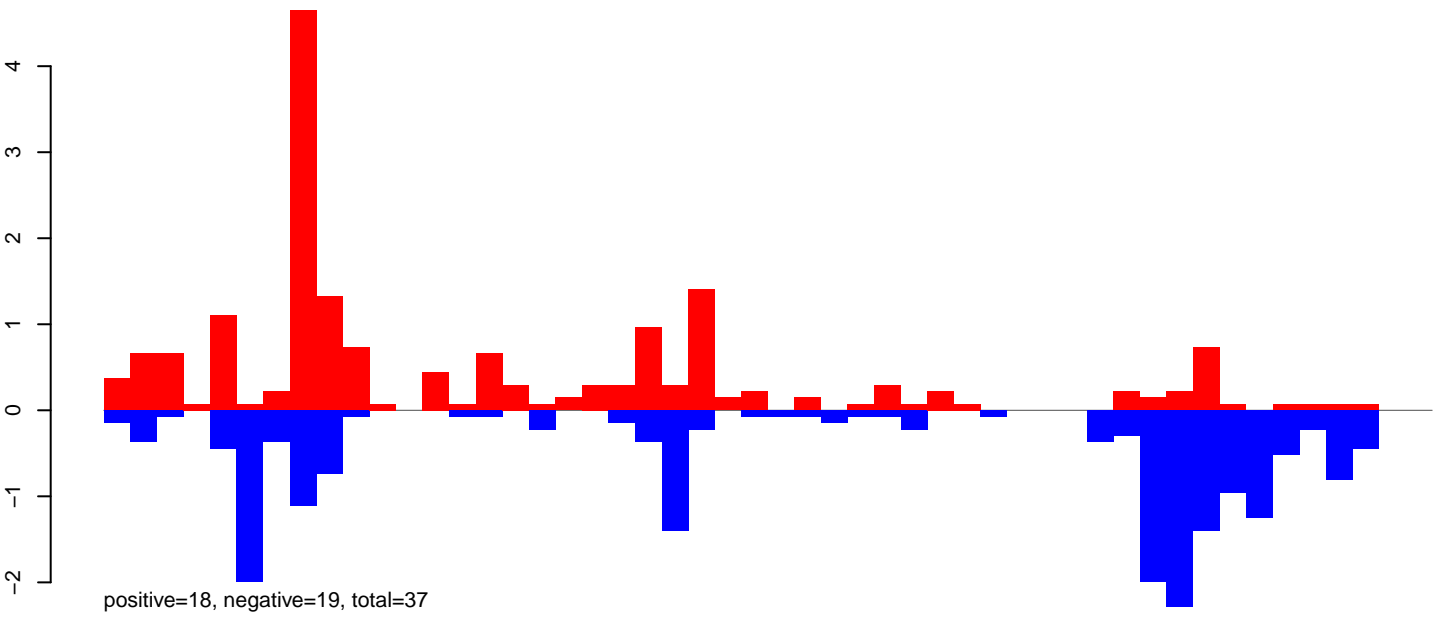
positive=8, negative=113, total=122

Window size=25, length=1991, TE@rnd-3_family-14@DNA_Sola-2@2010:1-1991

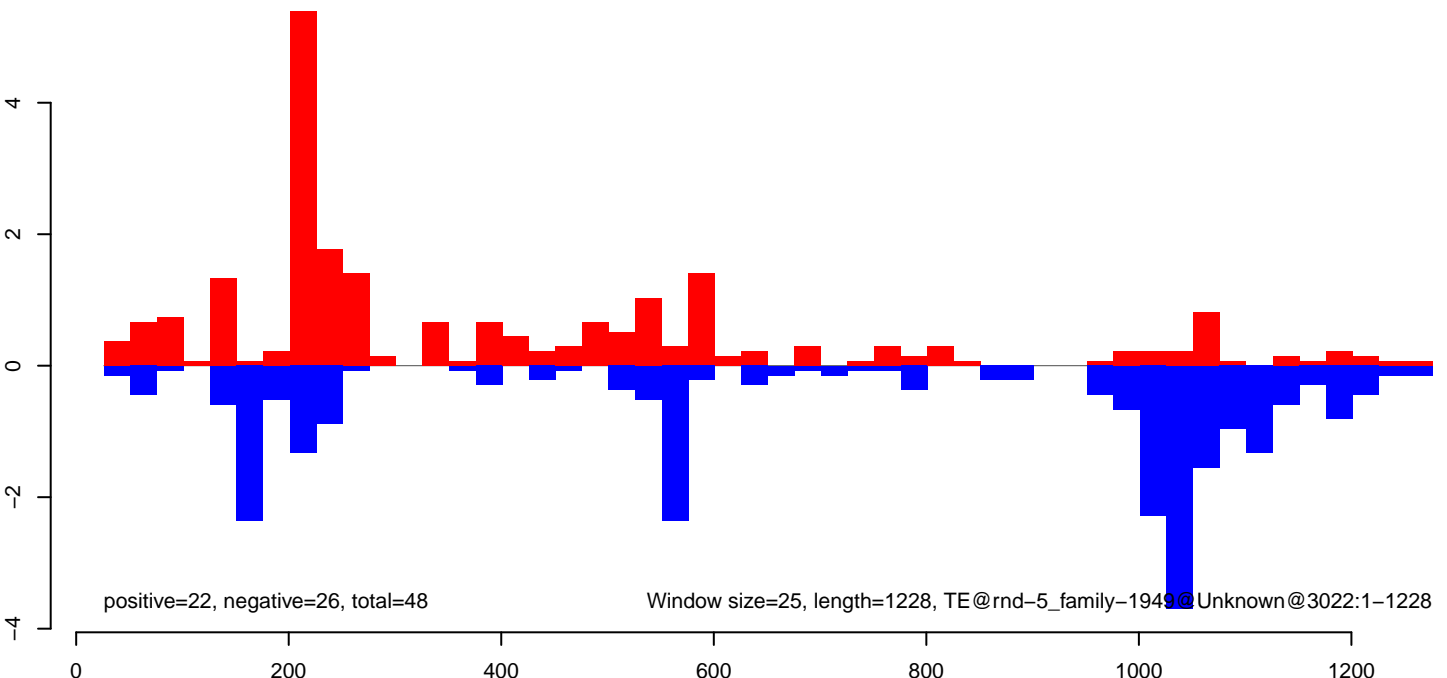
CuQuin_Testes_NL.18_23.rep



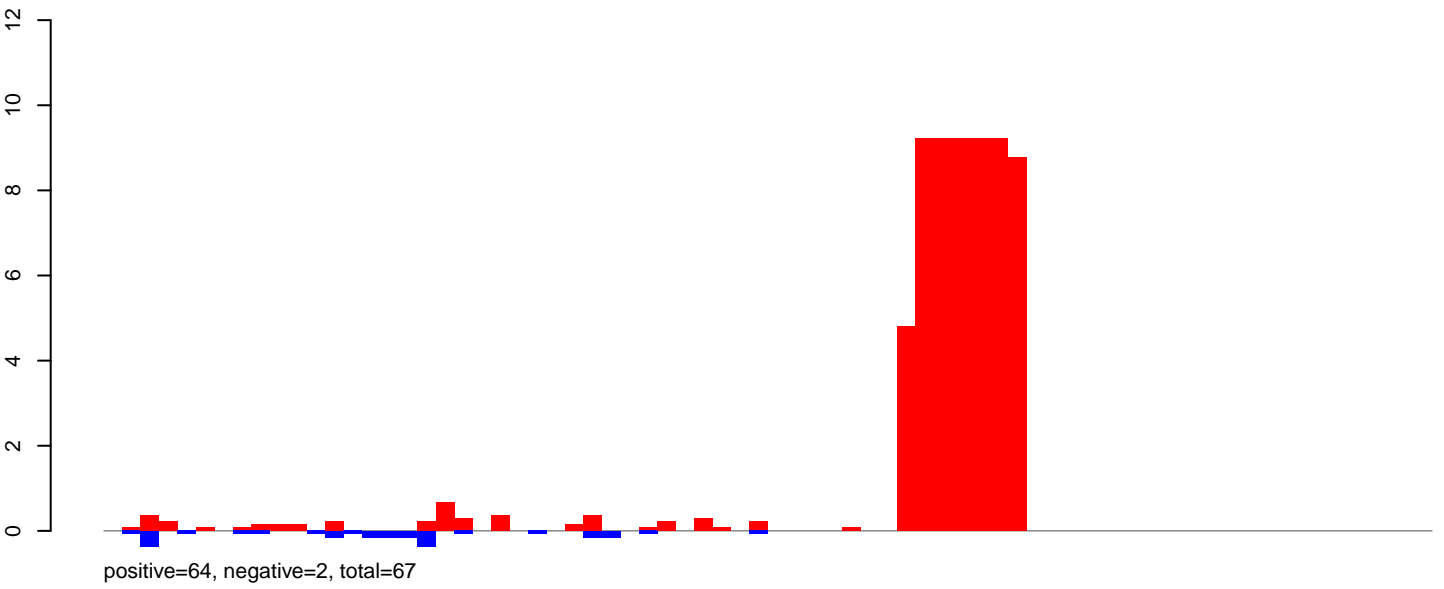
CuQuin_Testes_NL.24_35.rep



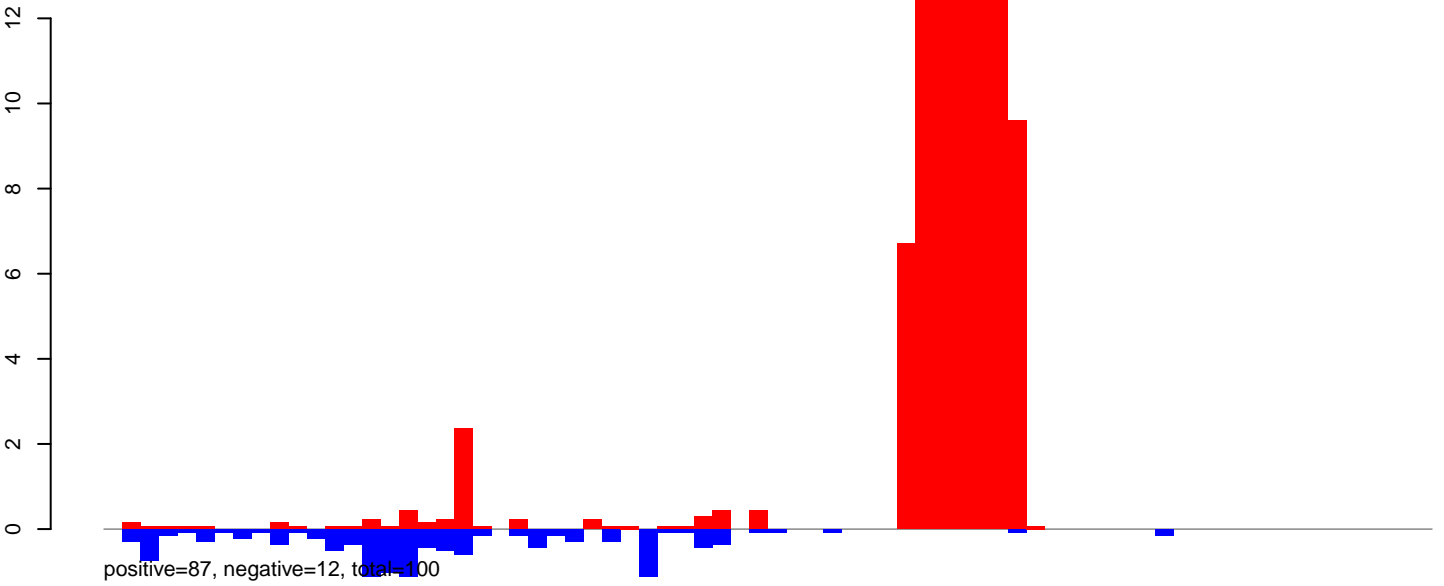
CuQuin_Testes_NL.rep



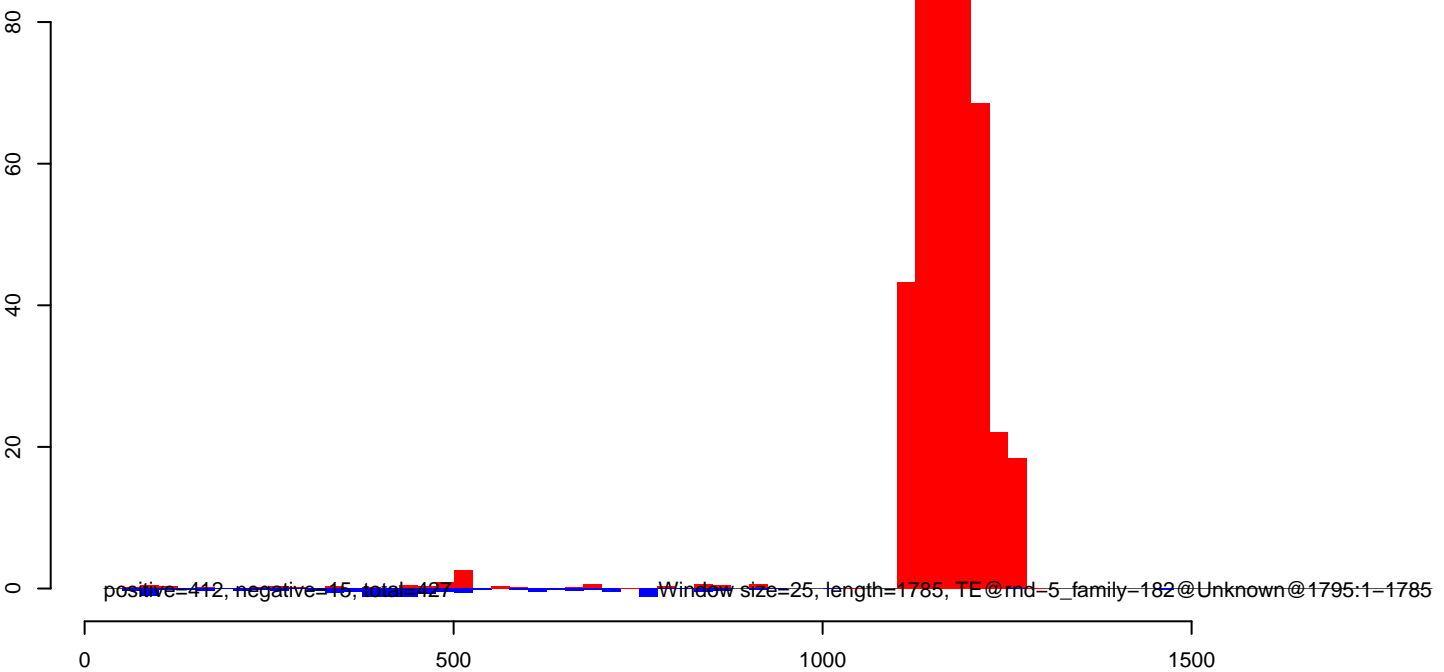
CuQuin_Testes_NL.18_23.rep



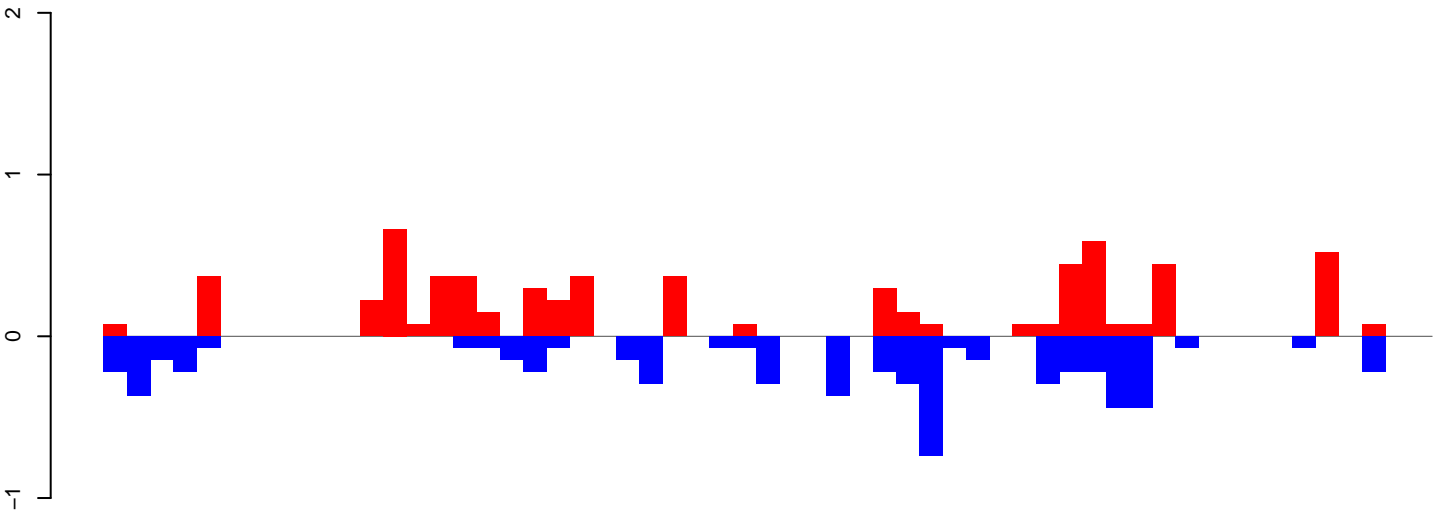
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

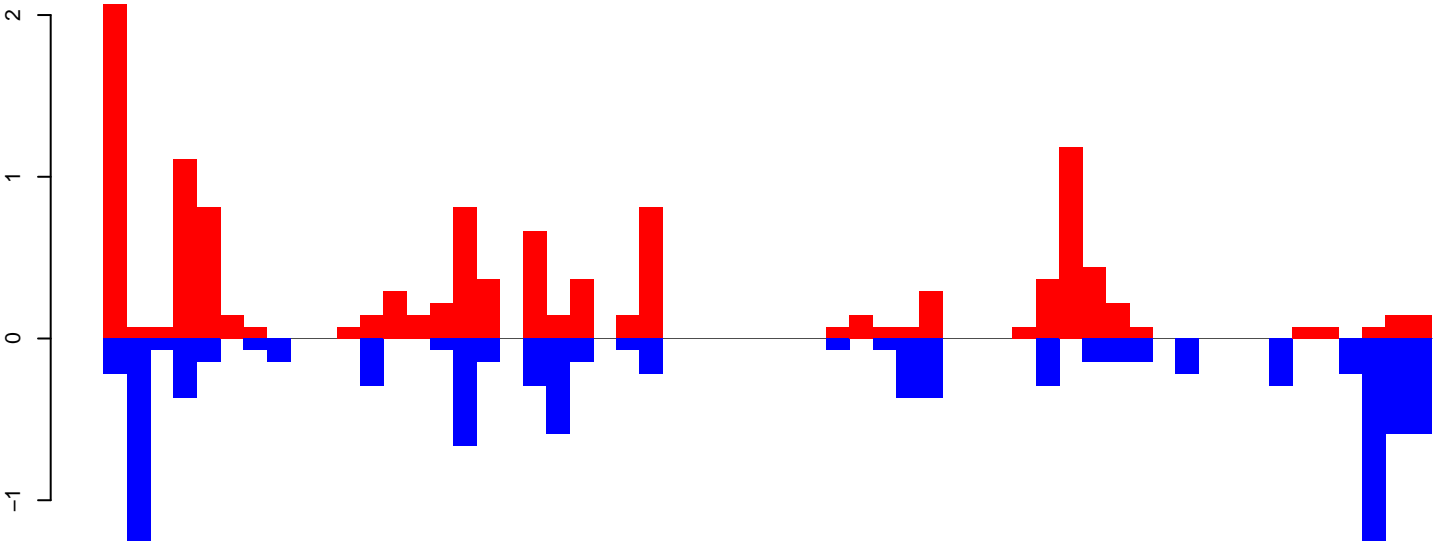


CuQuin_Testes_NL.18_23.rep



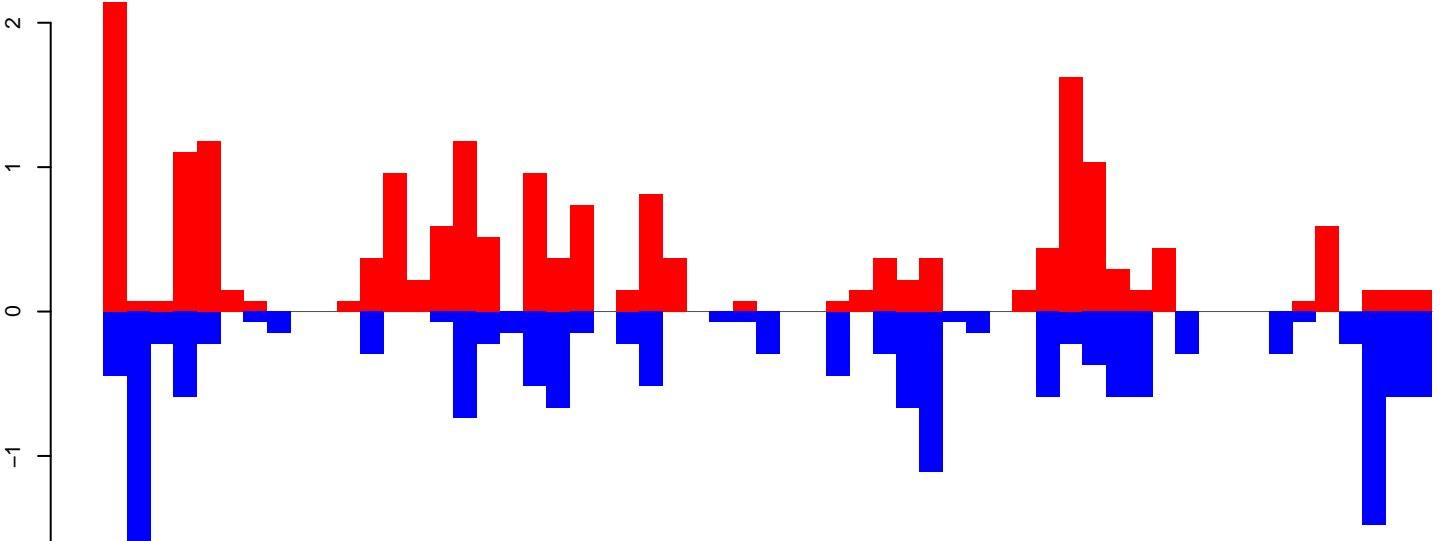
positive=6, negative=6, total=13

CuQuin_Testes_NL.24_35.rep



positive=12, negative=10, total=22

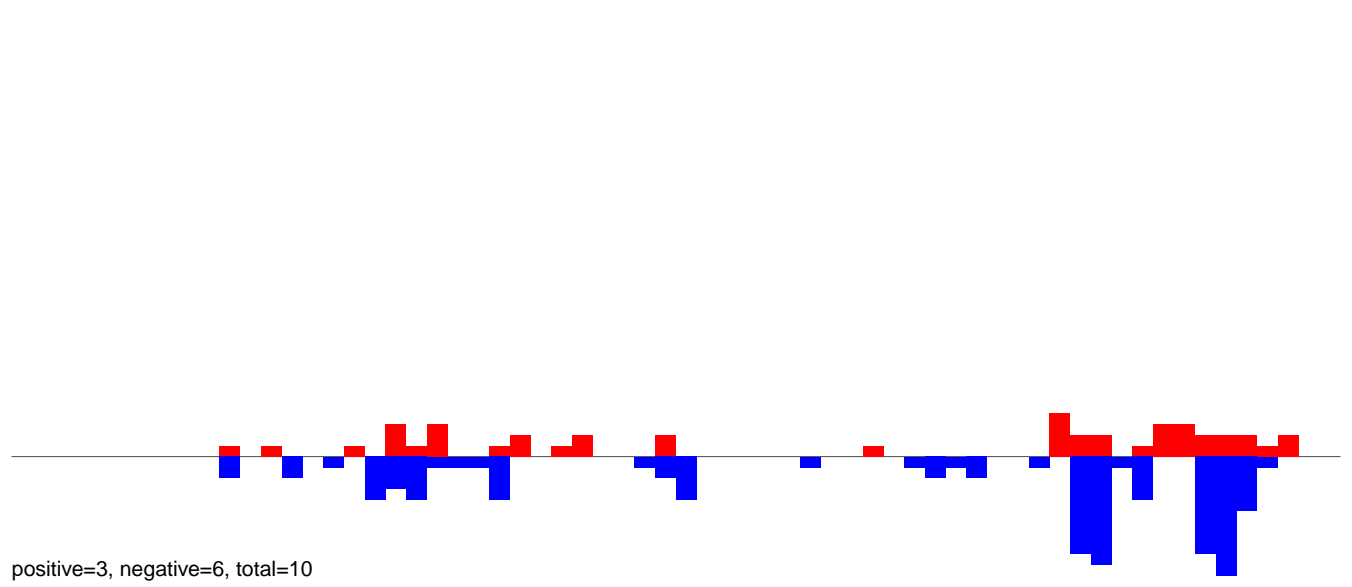
CuQuin_Testes_NL.rep



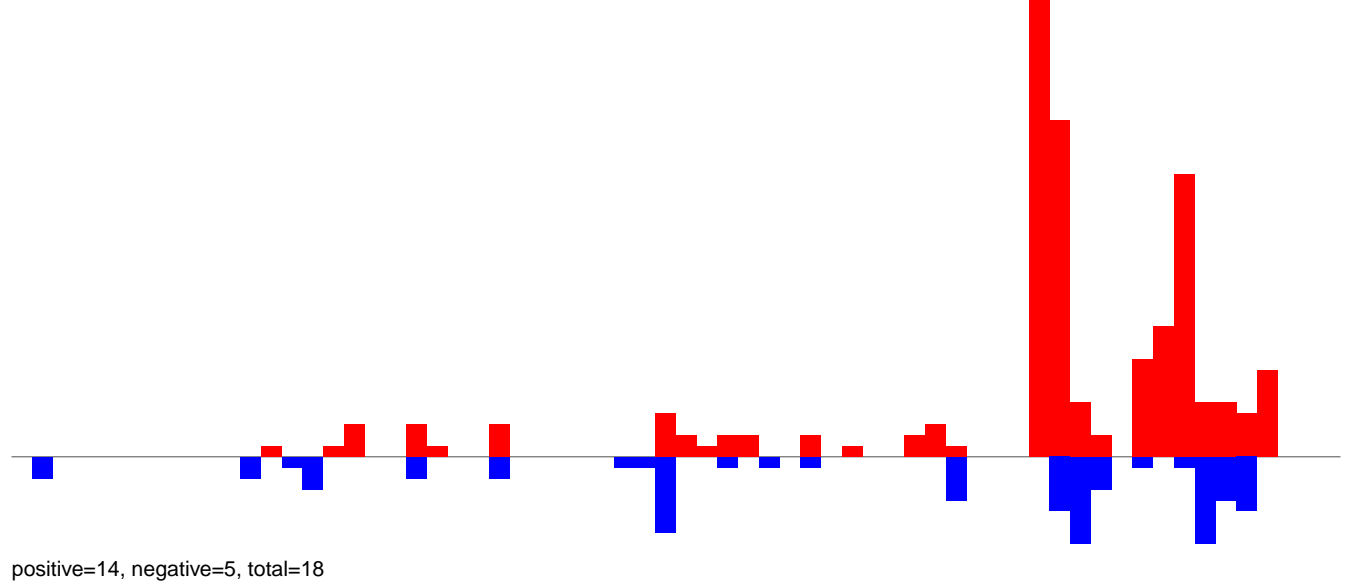
positive=19, negative=16, total=35

Window size=25, length=1422, TE@rnd-5_family-1609@Unknown@1238:1-1422

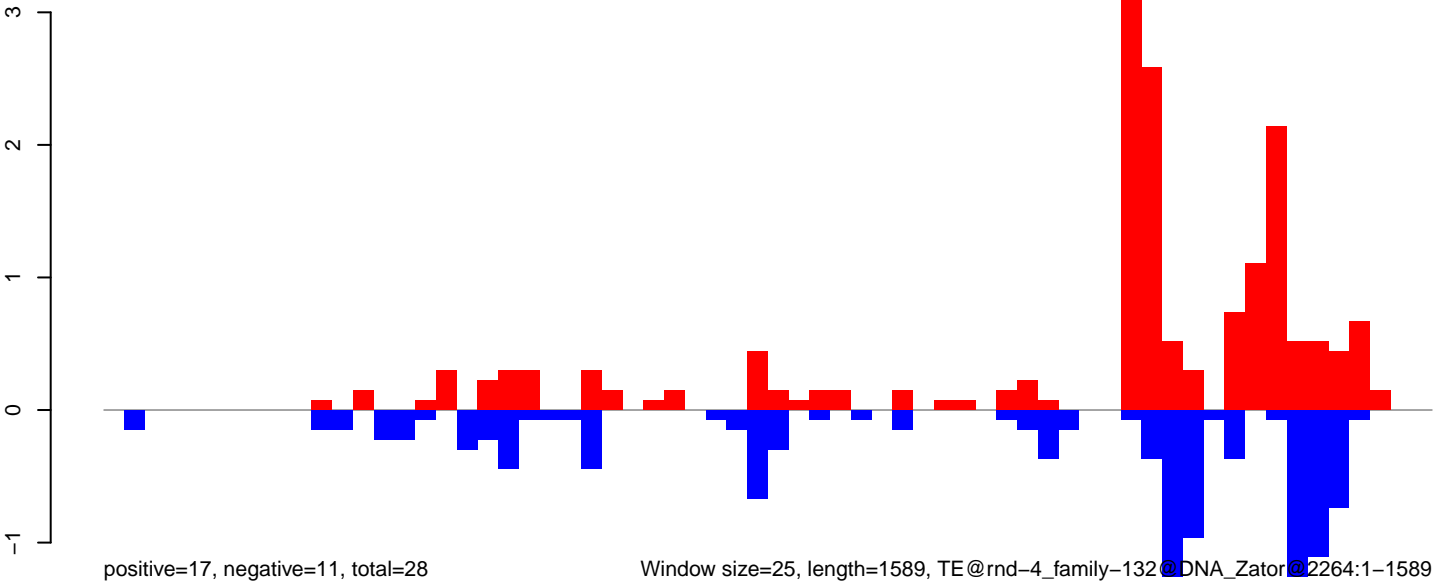
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



Window size=25, length=1589, TE@rnd-4_family-132@DNA_Zator@2264:1-1589

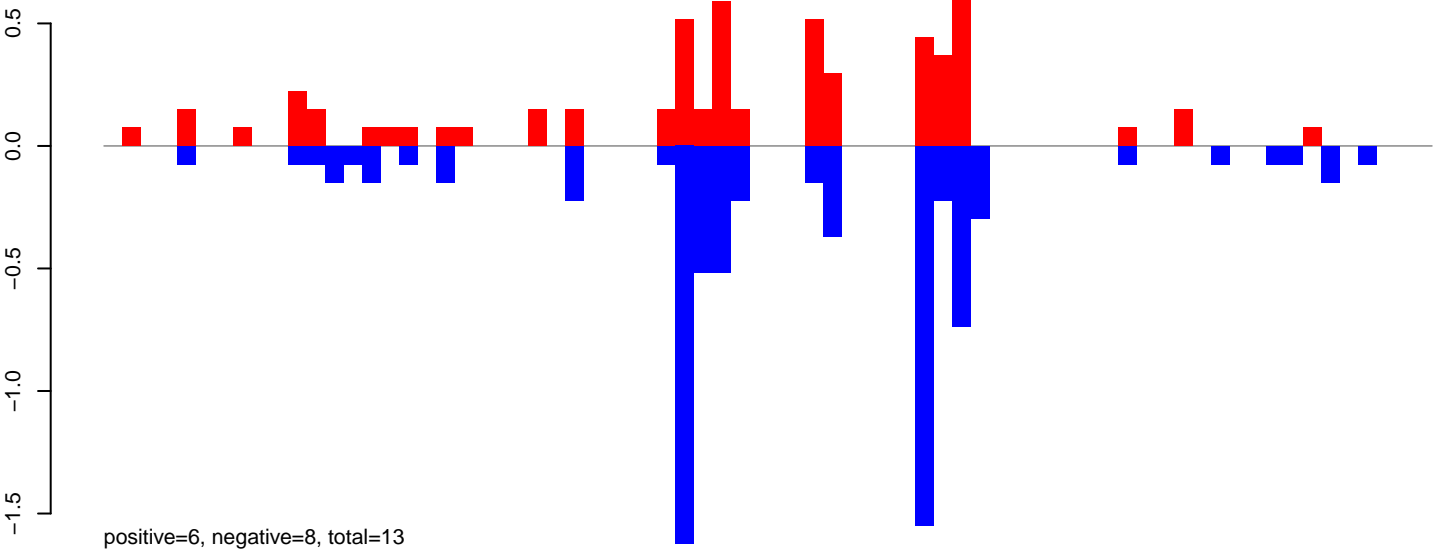
0

500

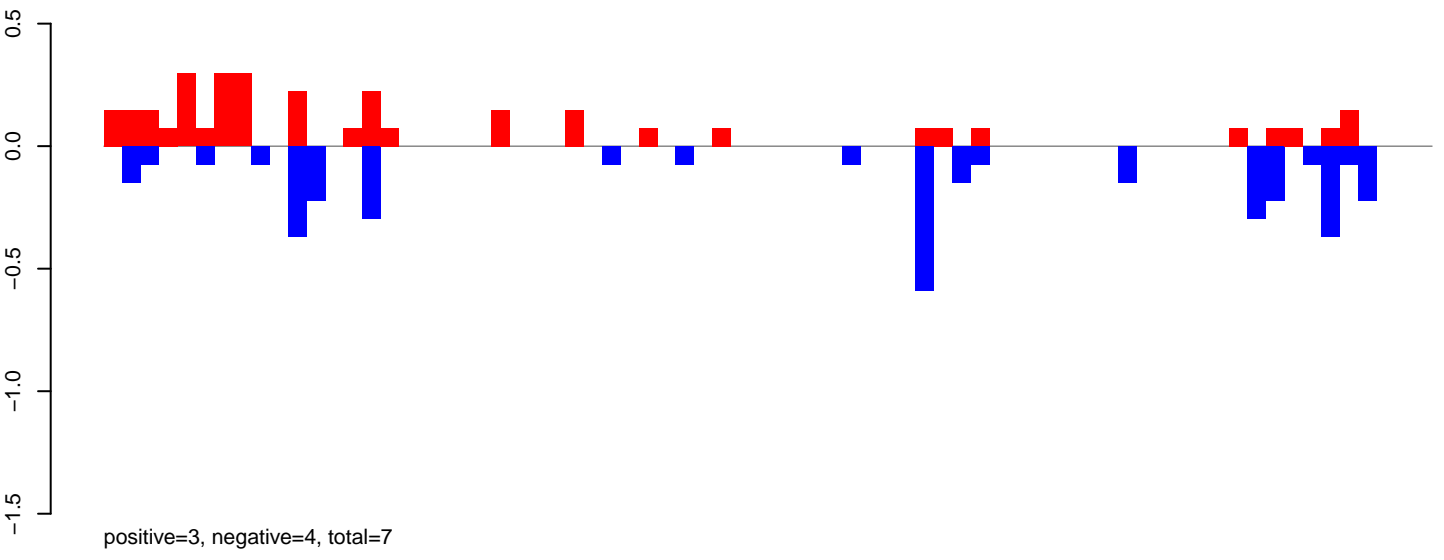
1000

1500

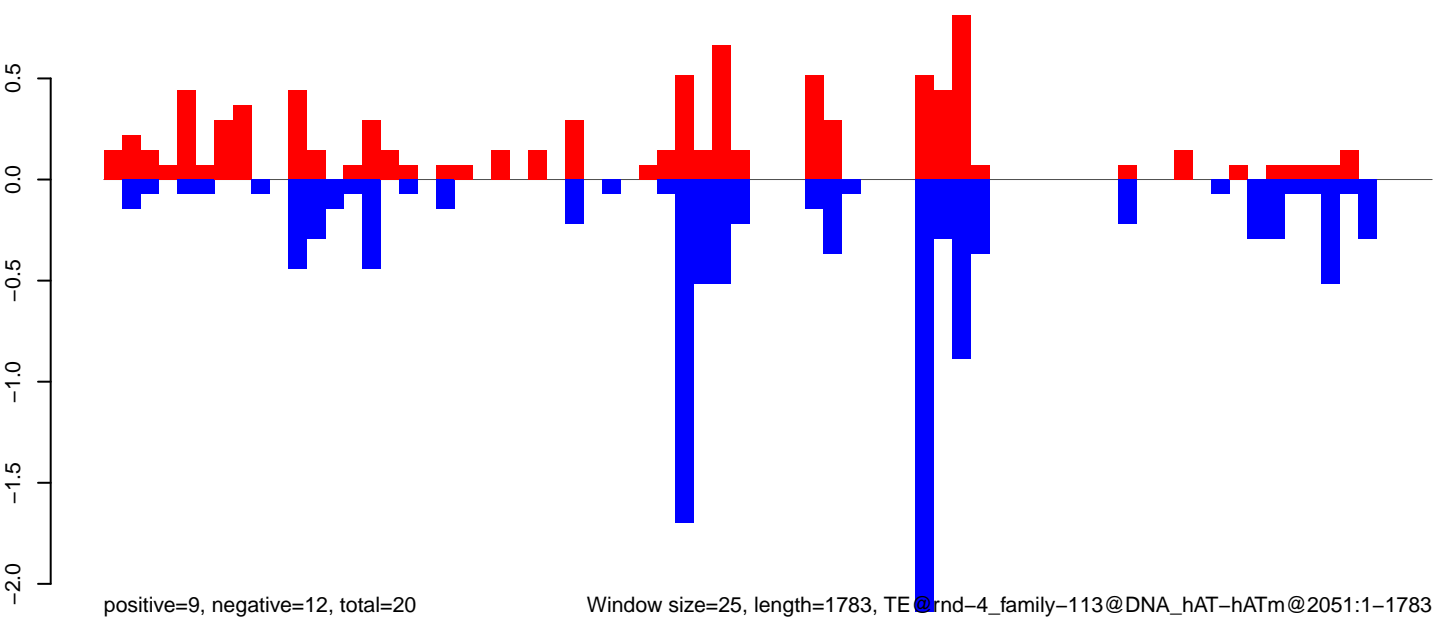
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



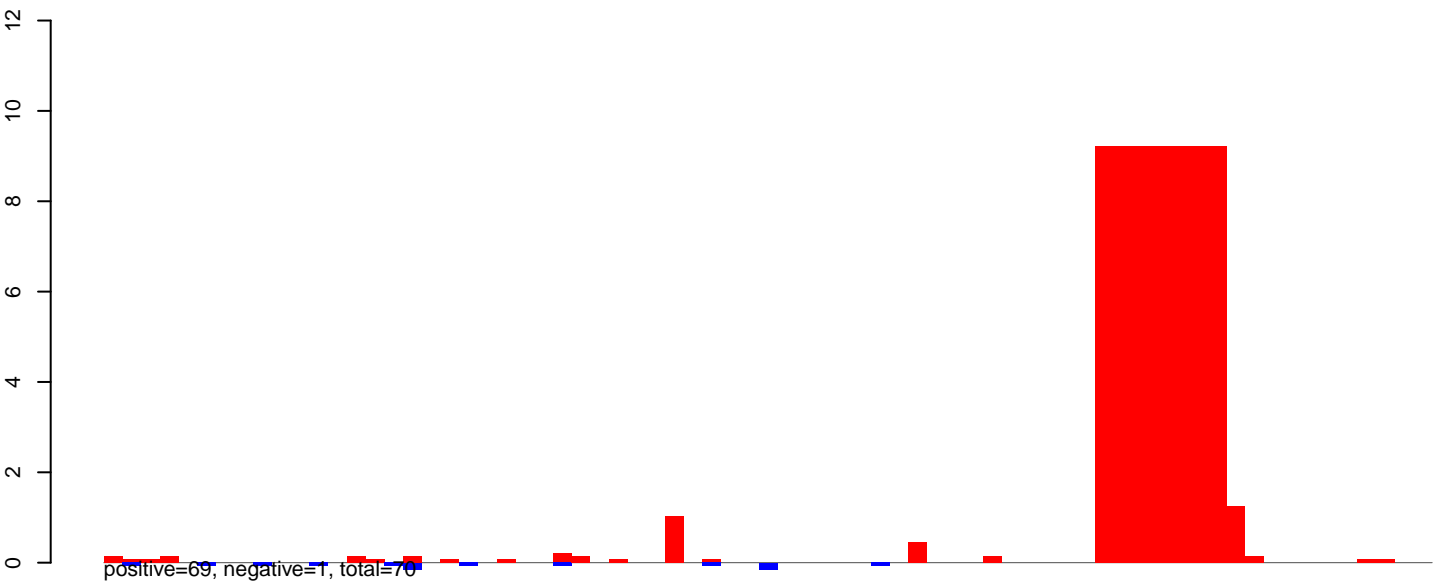
CuQuin_Testes_NL.rep



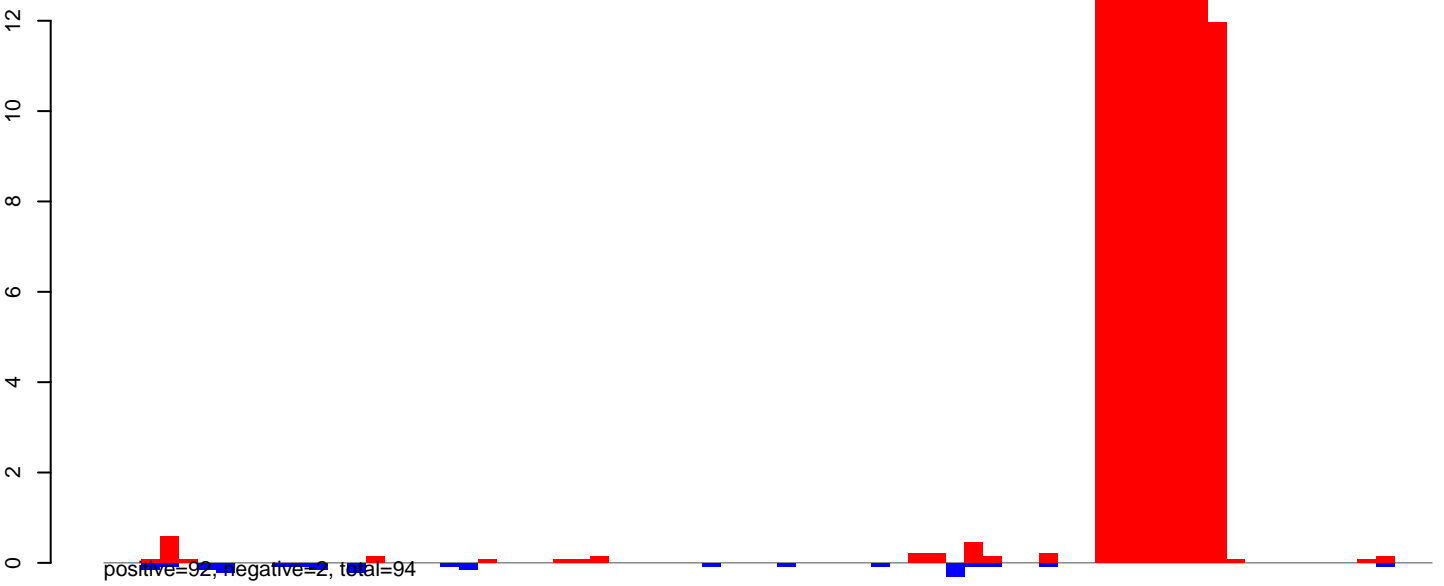
Window size=25, length=1783, TE @ rnd-4_family-113@DNA_hAT-hATm@2051:1-1783

0 500 1000 1500

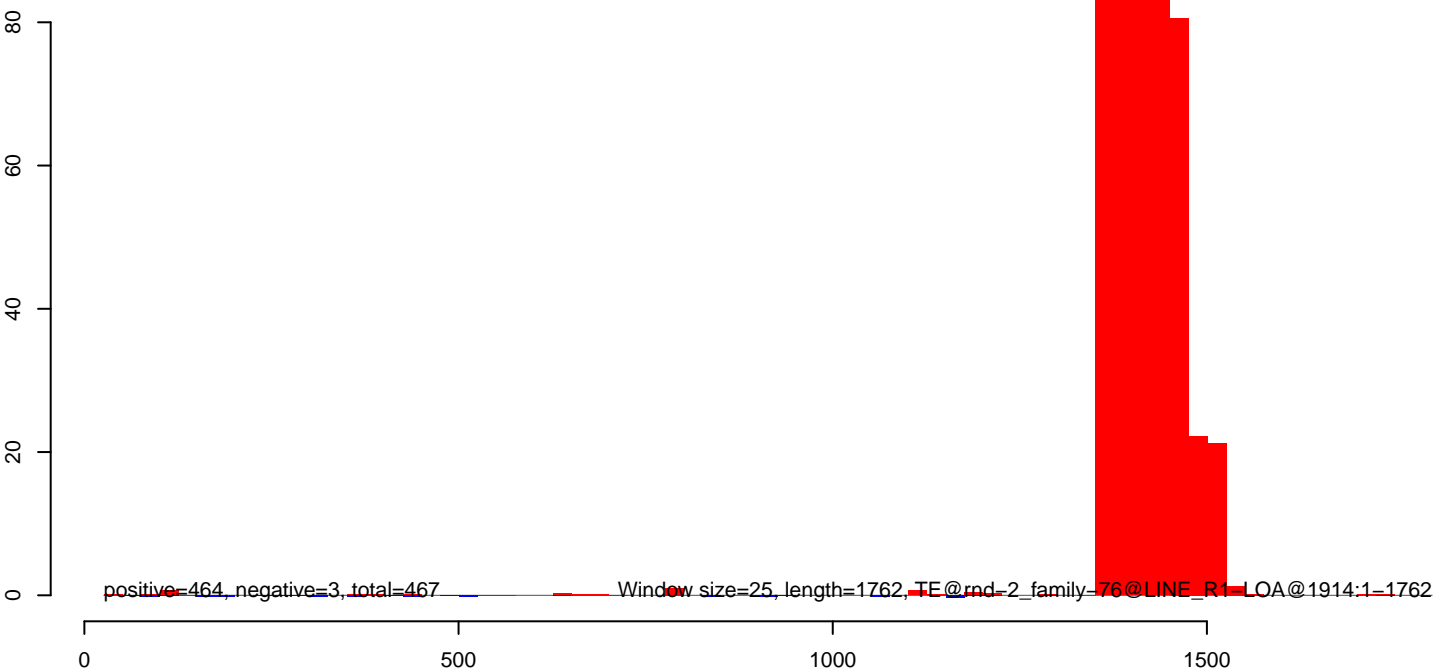
CuQuin_Testes_NL.18_23.rep



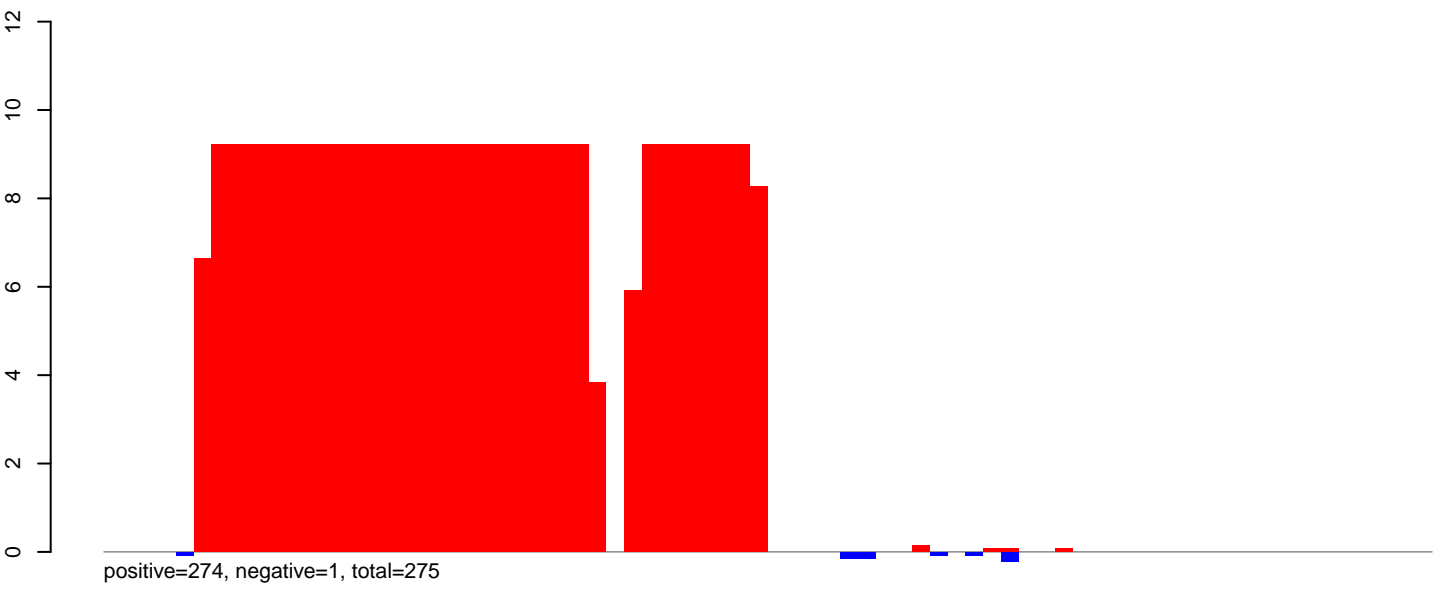
CuQuin_Testes_NL.24_35.rep



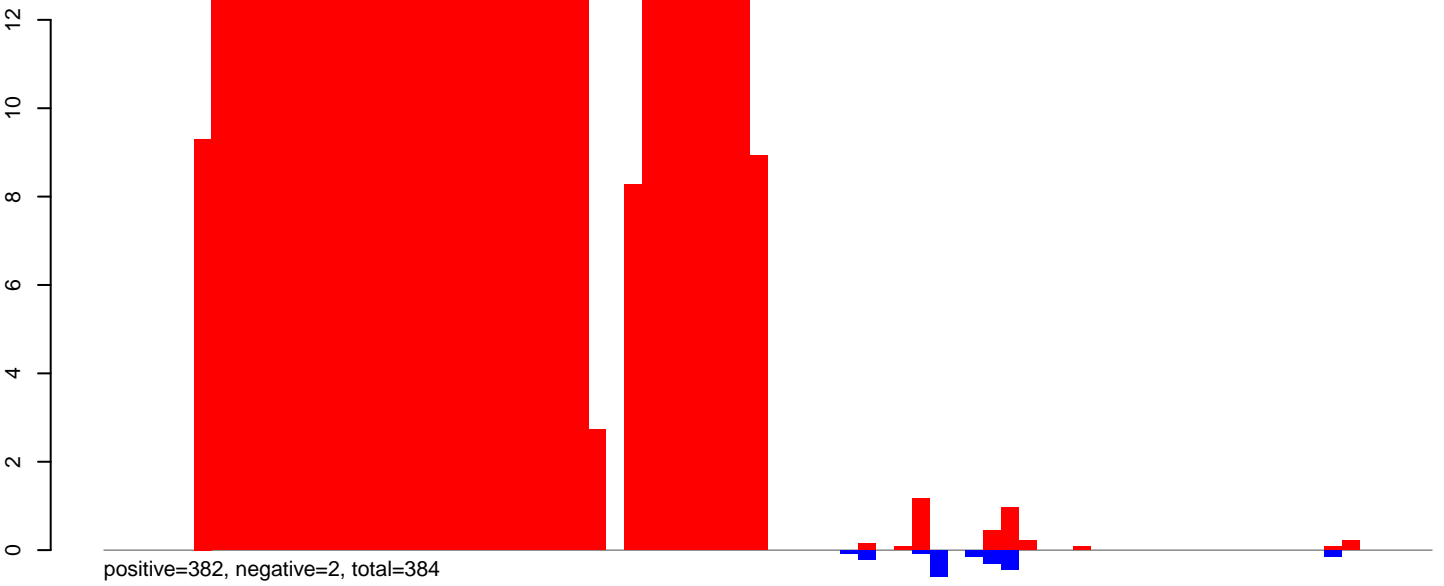
CuQuin_Testes_NL.rep



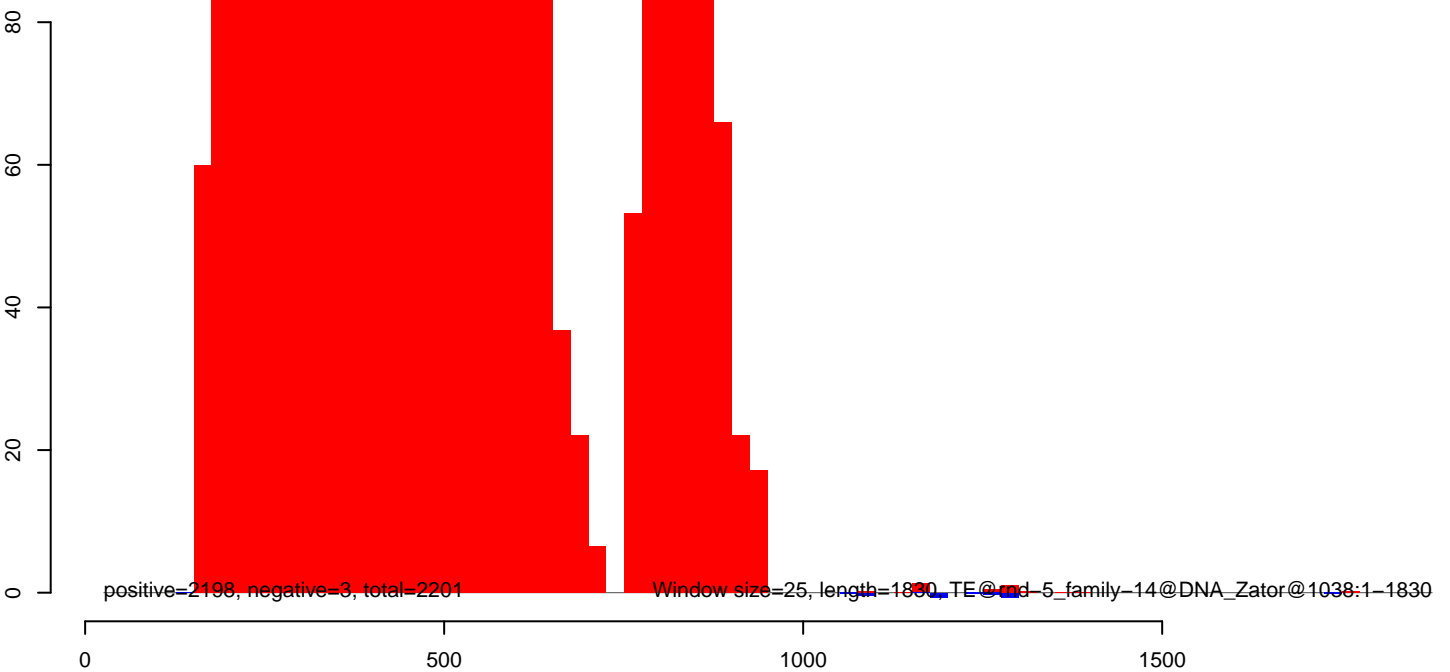
CuQuin_Testes_NL.18_23.rep



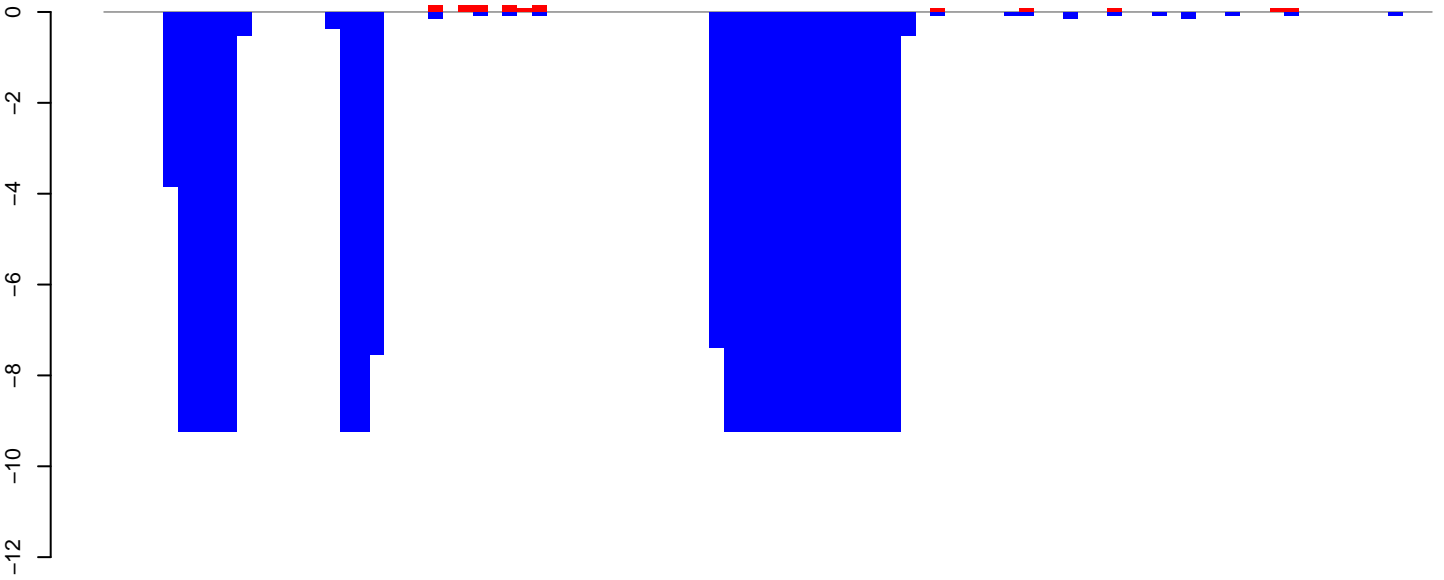
CuQuin_Testes_NL.24_35.rep



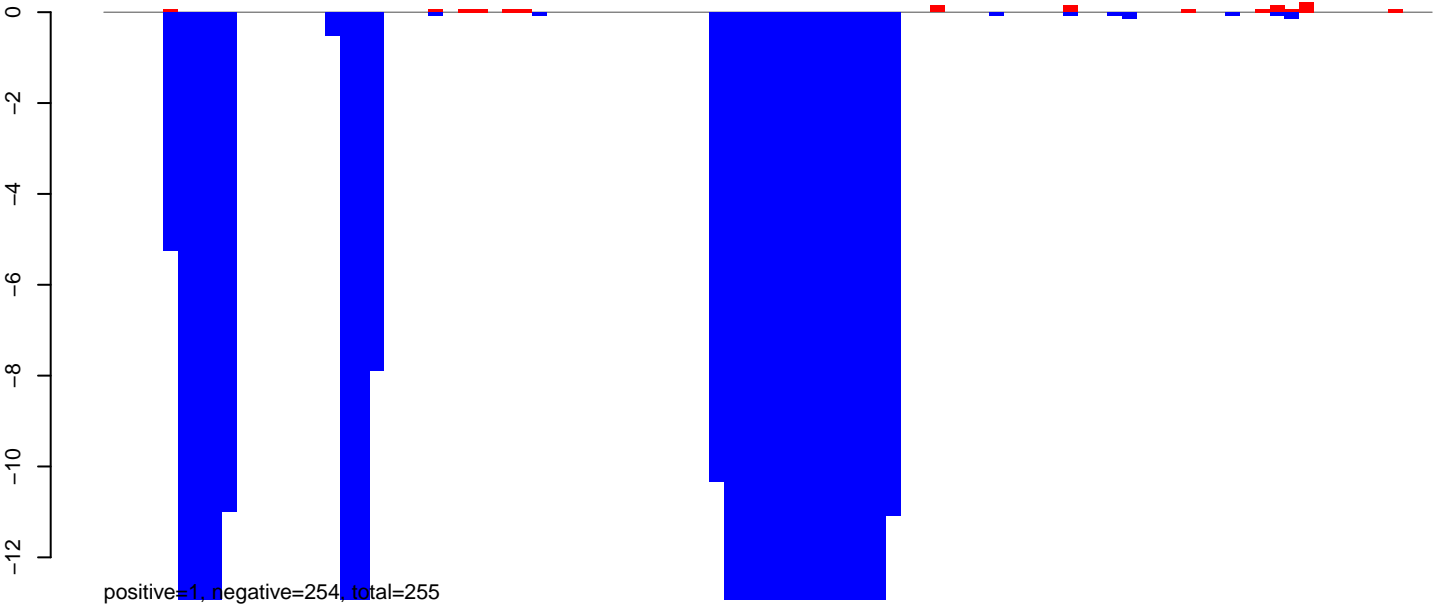
CuQuin_Testes_NL.rep



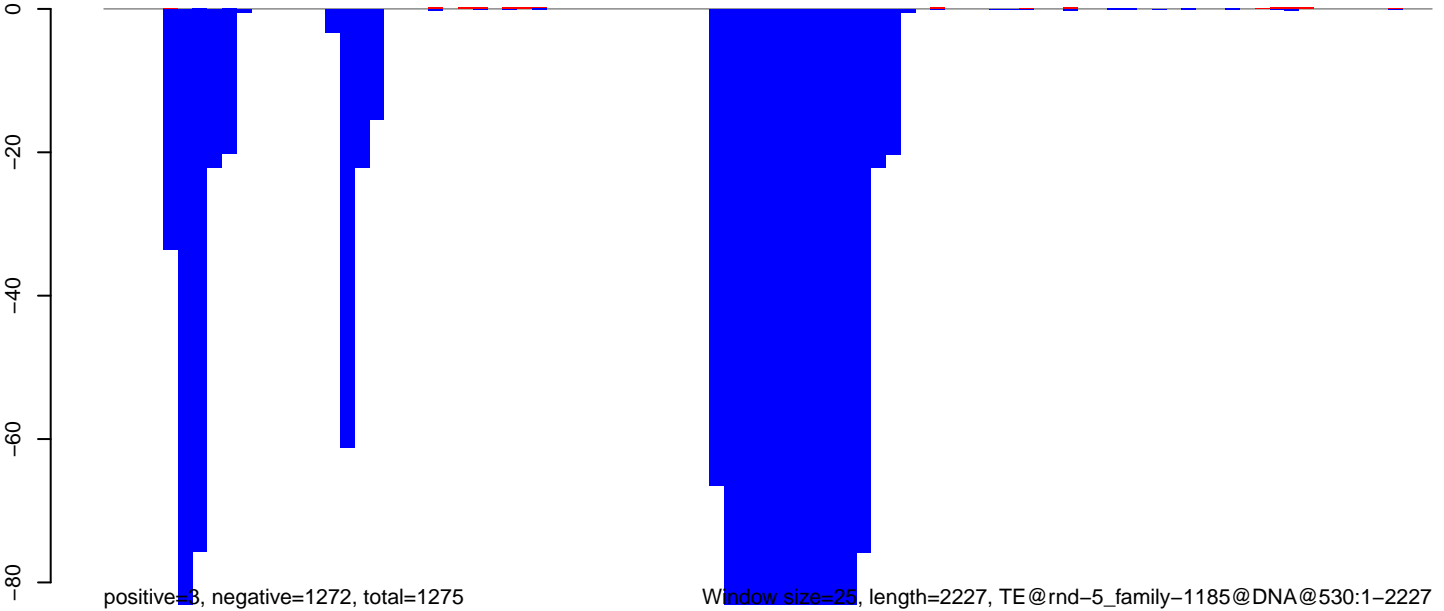
CuQuin_Testes_NL.18_23.rep



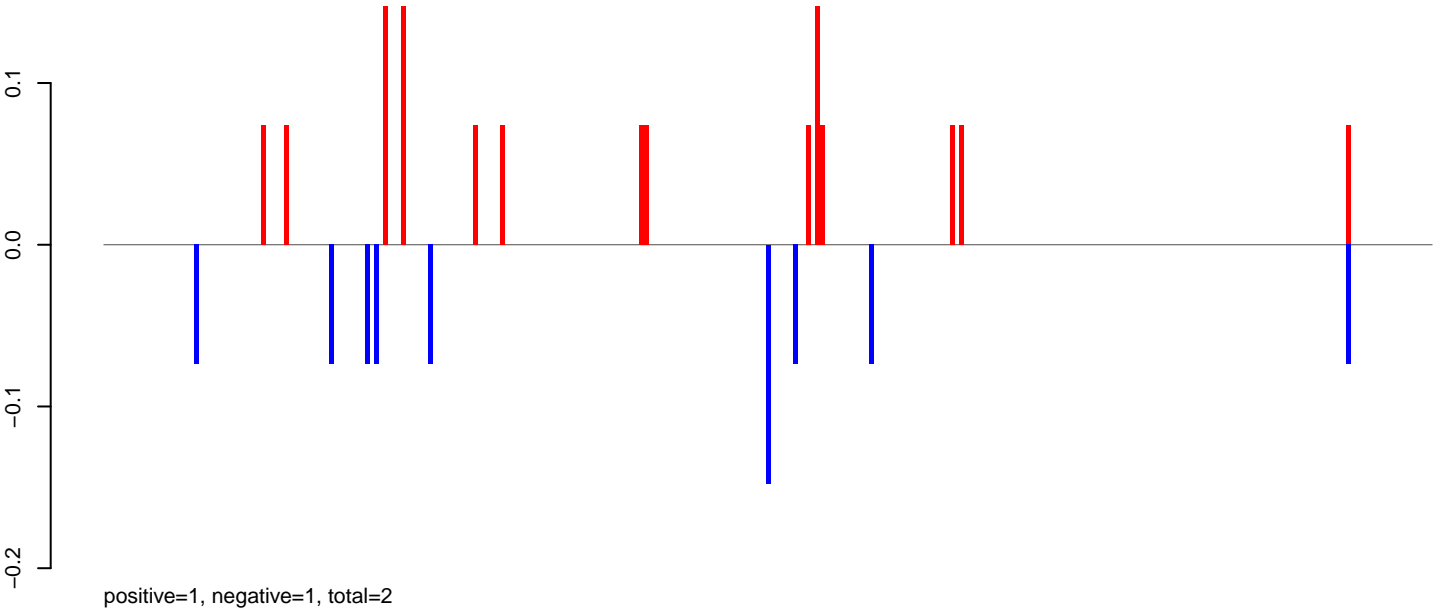
CuQuin_Testes_NL.24_35.rep



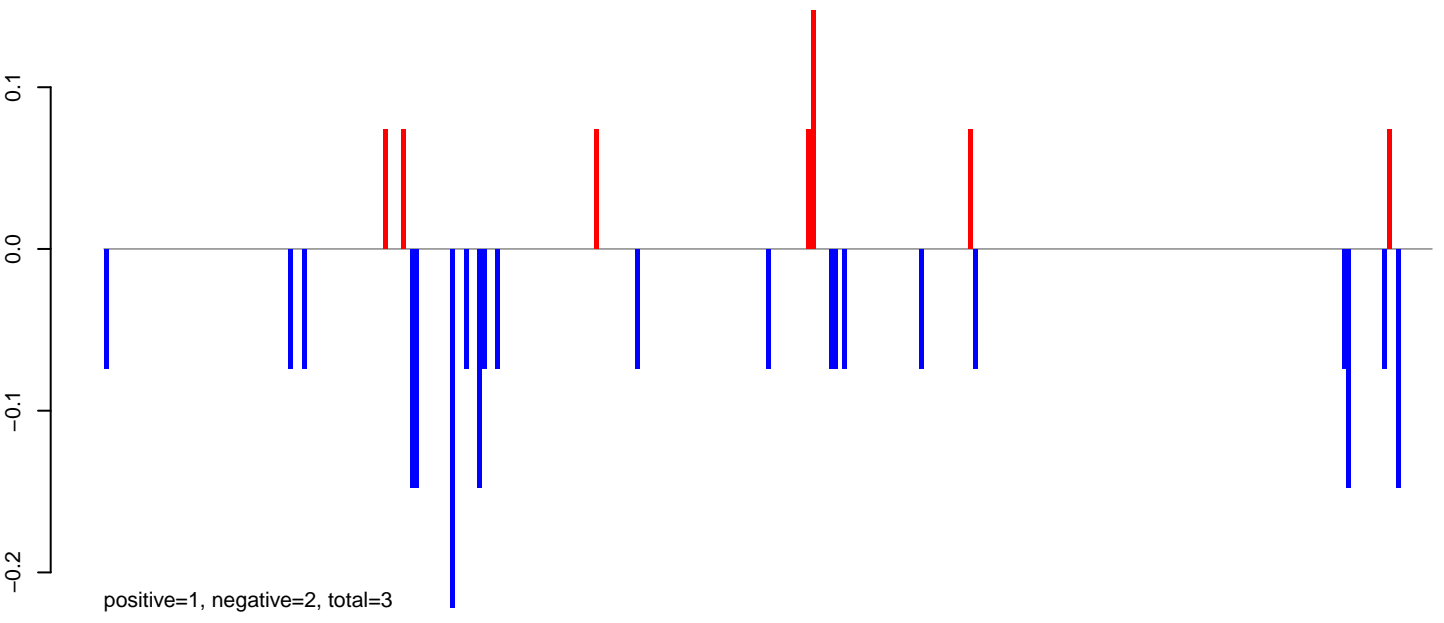
CuQuin_Testes_NL.rep



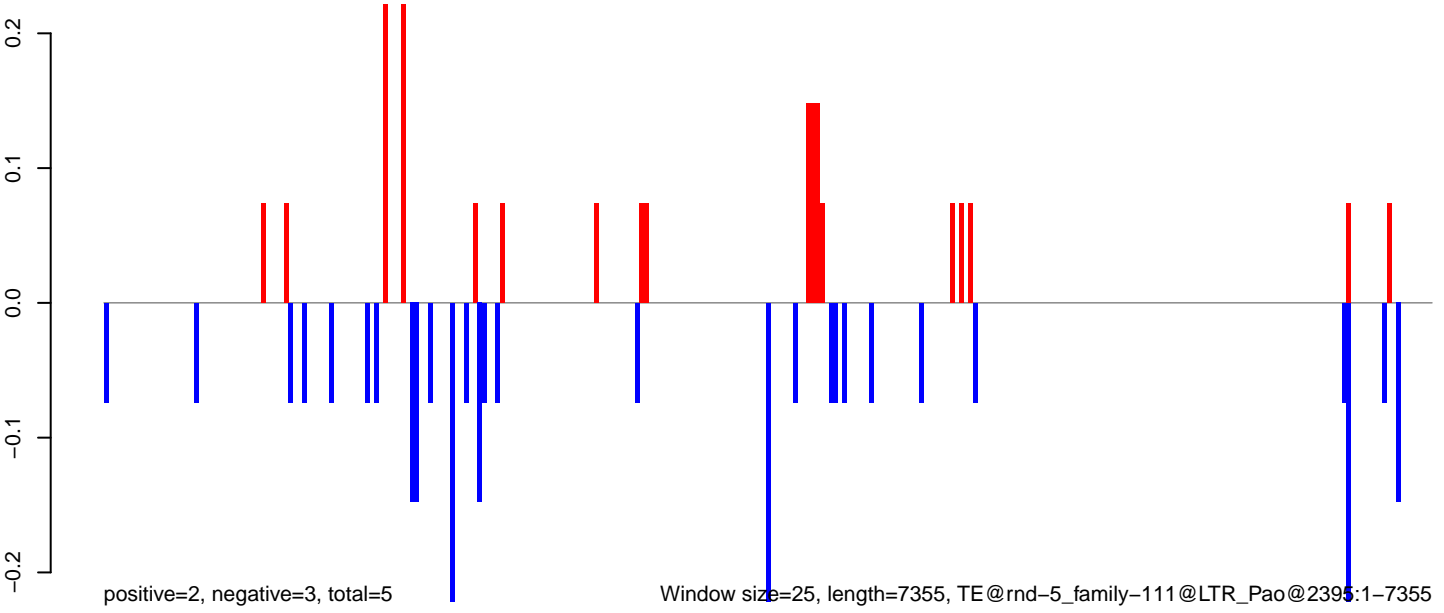
CuQuin_Testes_NL.18_23.rep



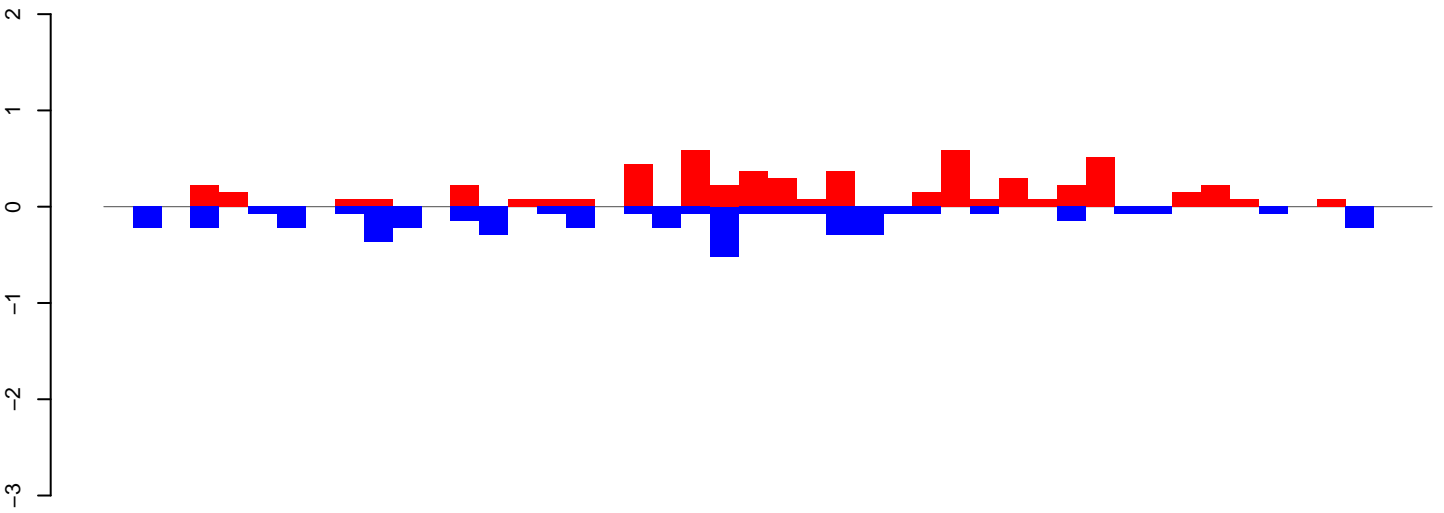
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

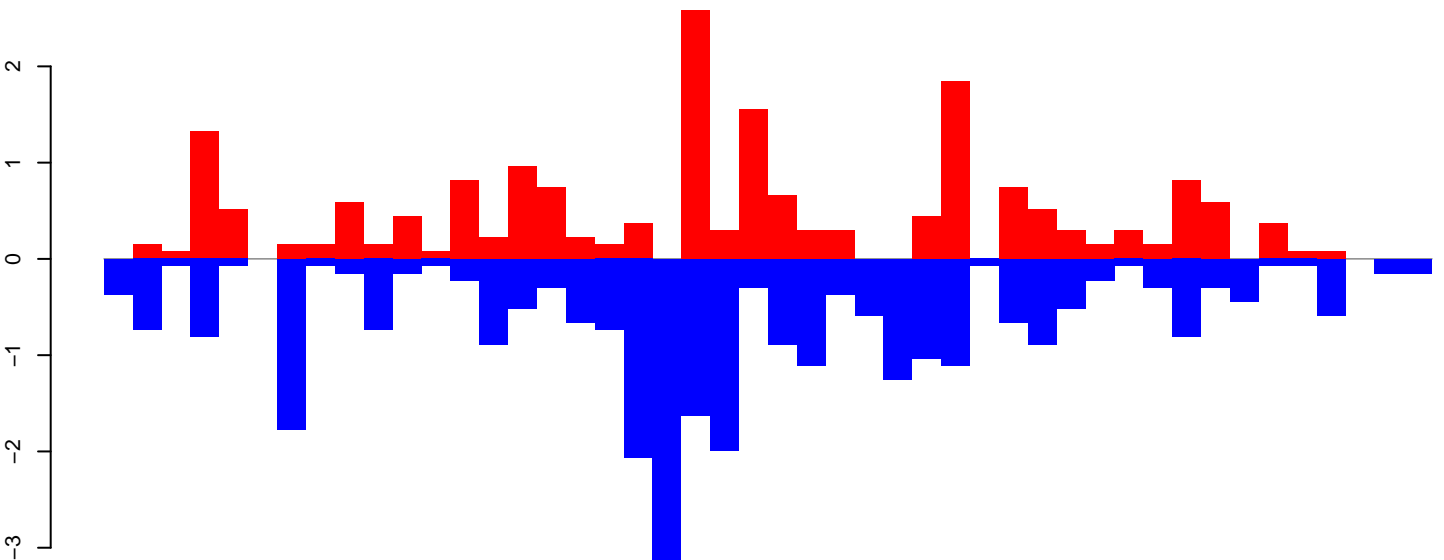


CuQuin_Testes_NL.18_23.rep



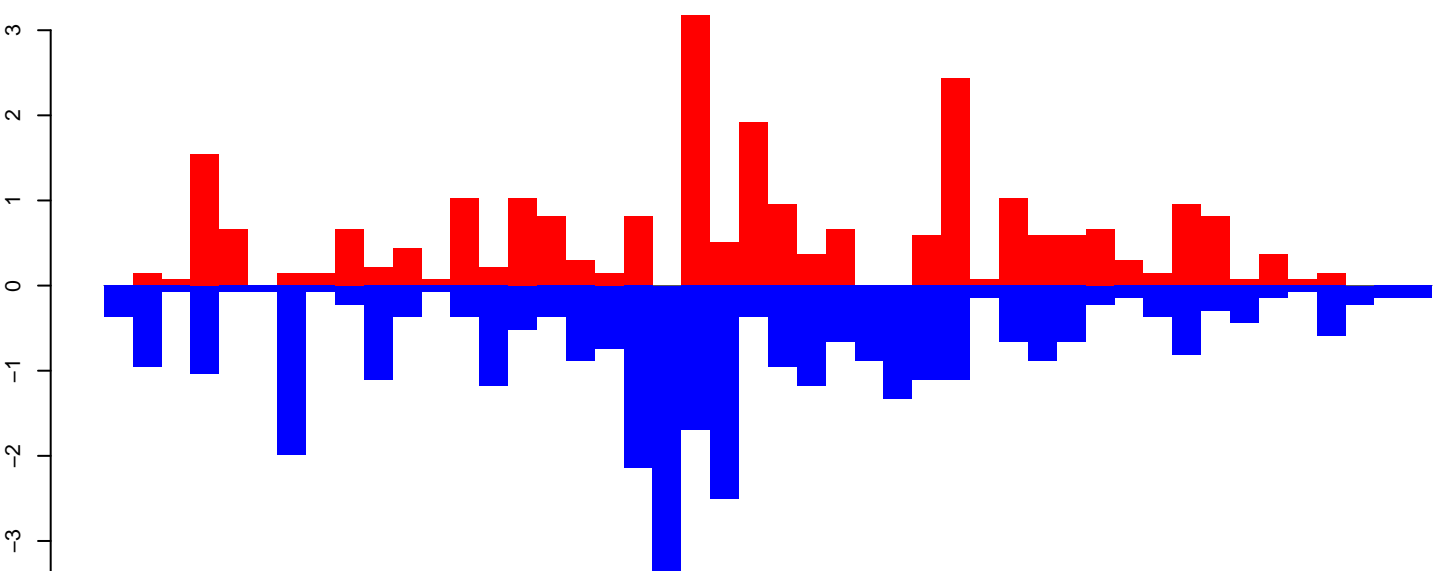
positive=6, negative=5, total=10

CuQuin_Testes_NL.24_35.rep



positive=19, negative=30, total=49

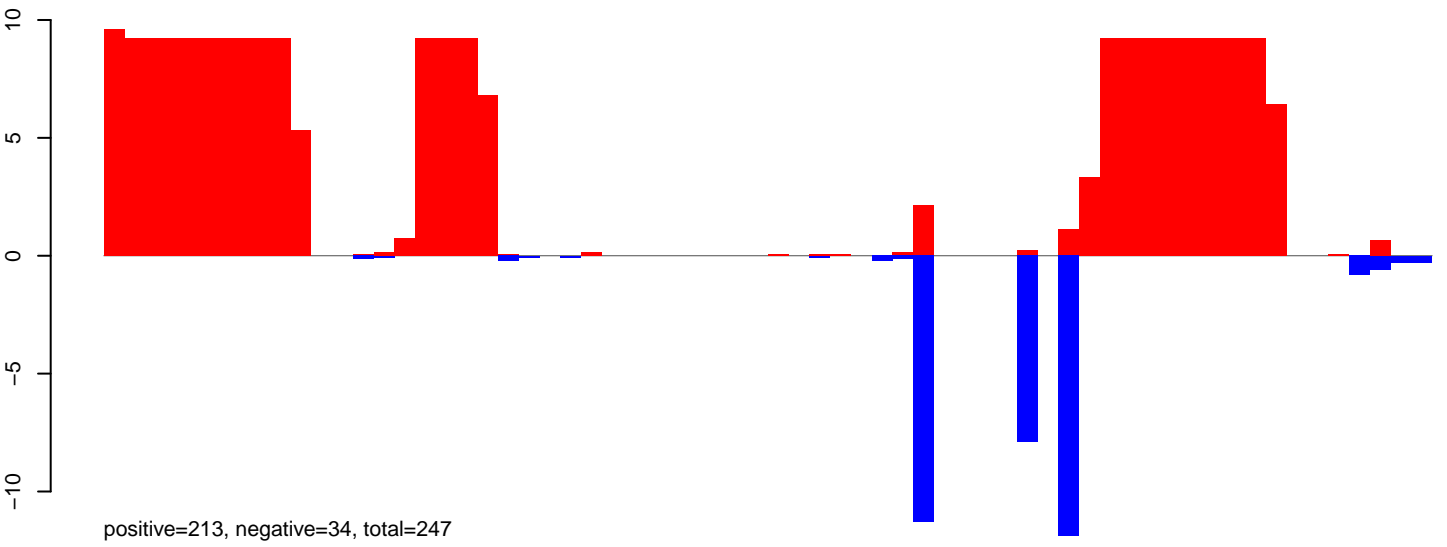
CuQuin_Testes_NL.rep



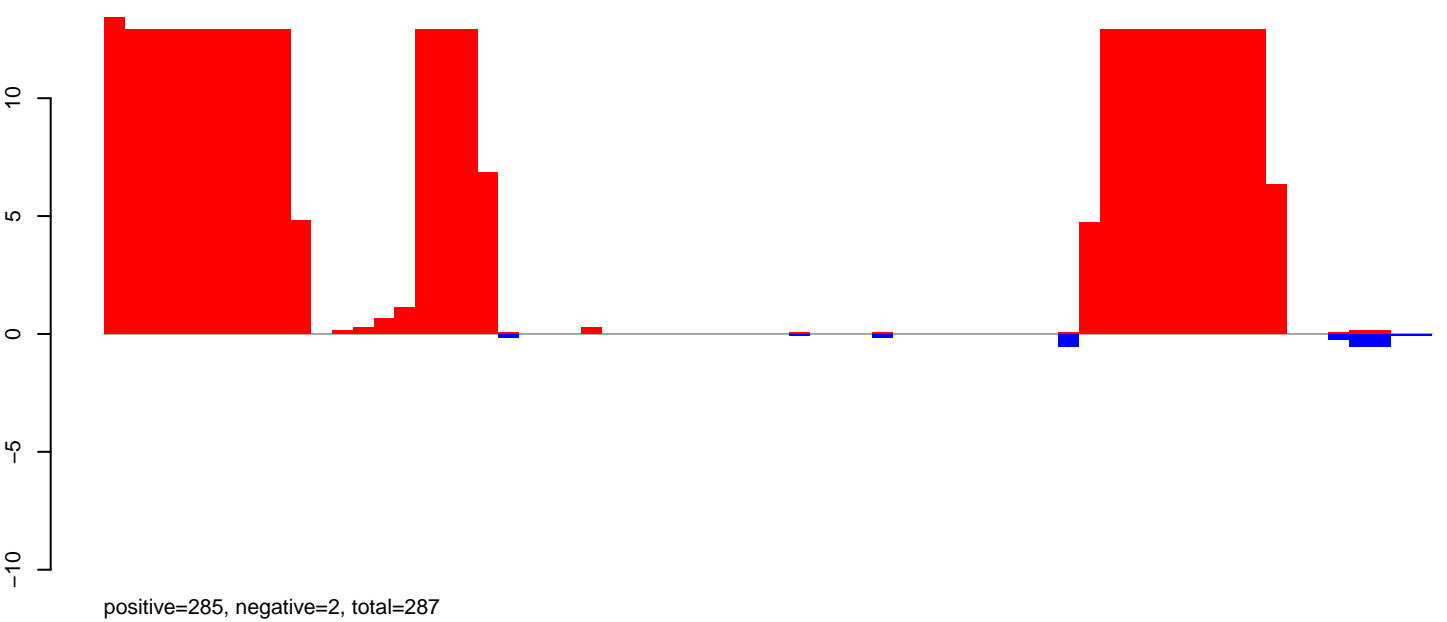
positive=25, negative=34, total=59

Window size=25, length=1149, TE@rnd-4_family-721@LINE_L1-Tx1@3542:1-1149

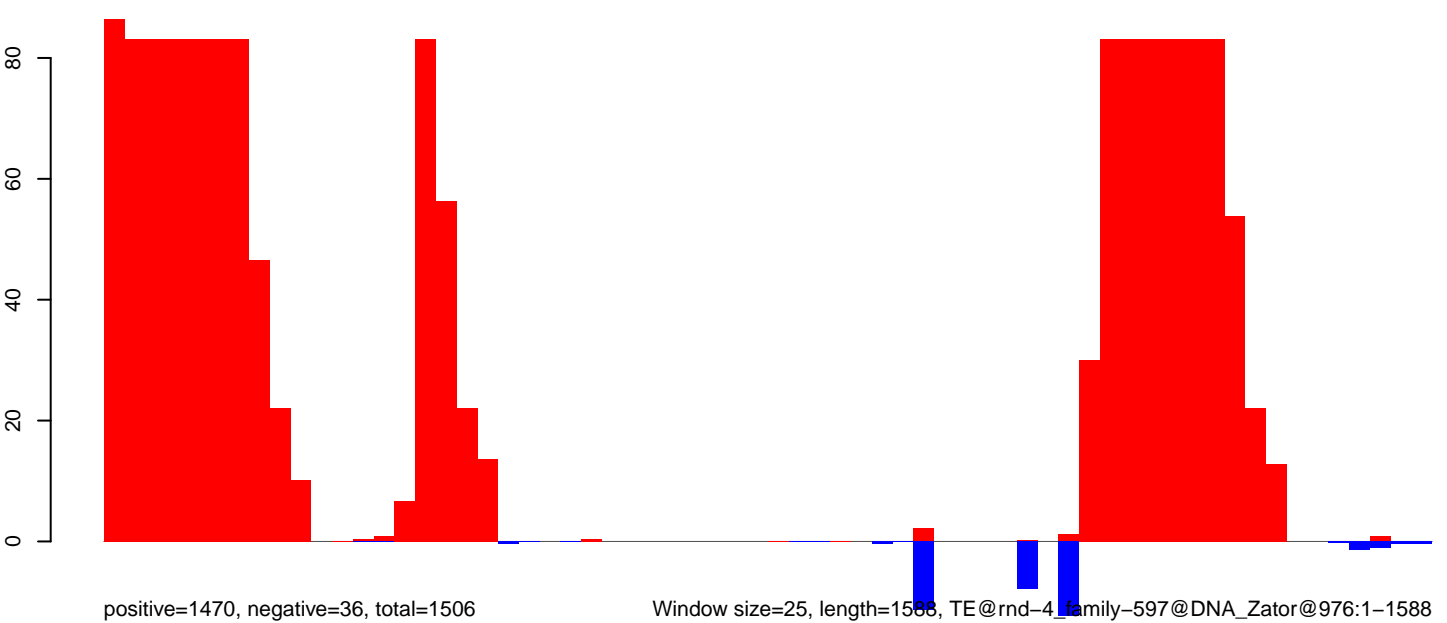
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



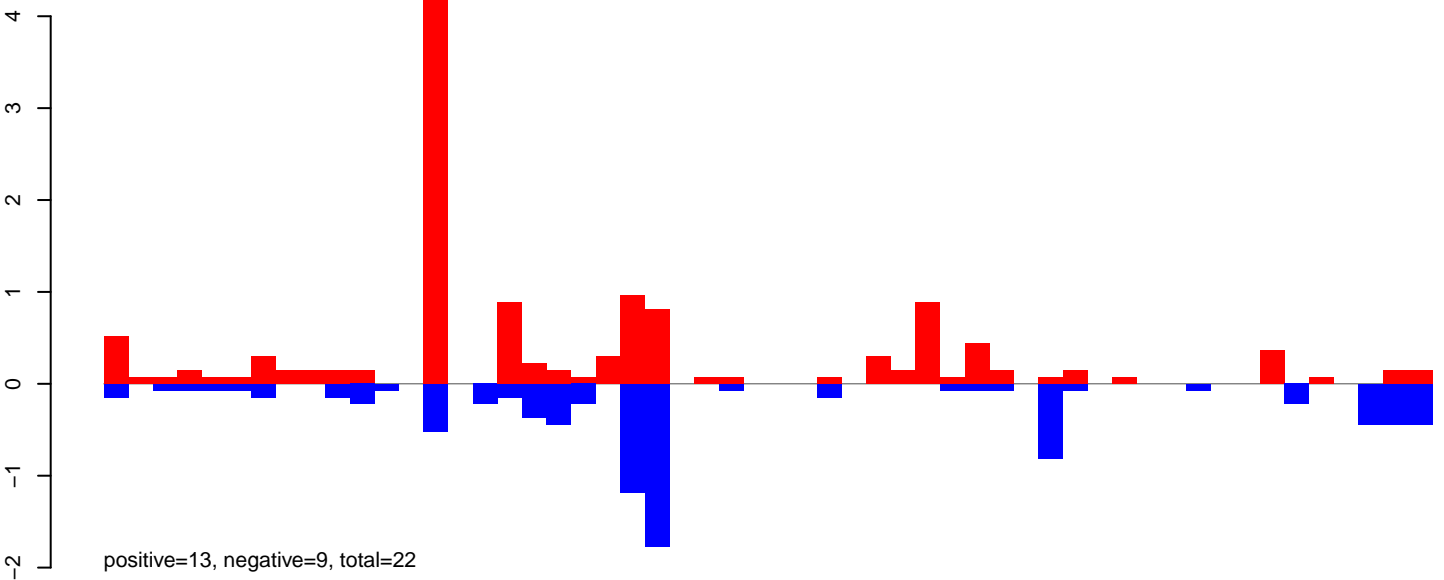
CuQuin_Testes_NL.rep



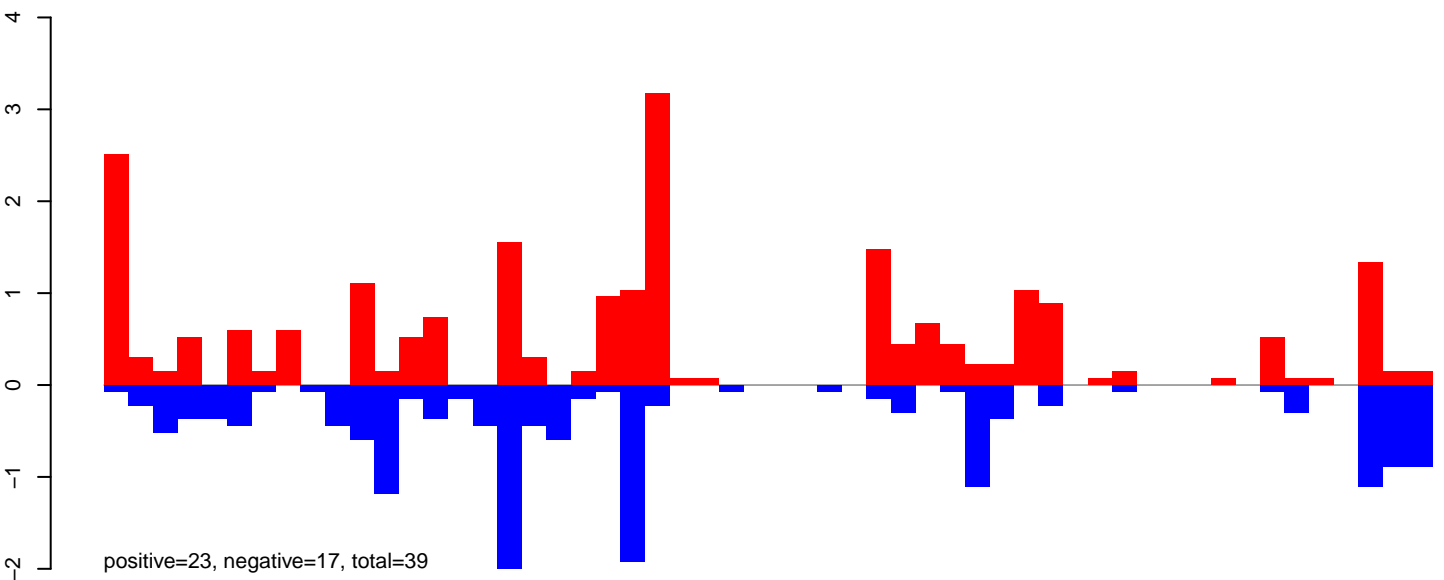
Window size=25, length=1588, TE@rnd-4_family-597@DNA_Zator@976:1-1588

0 500 1000 1500

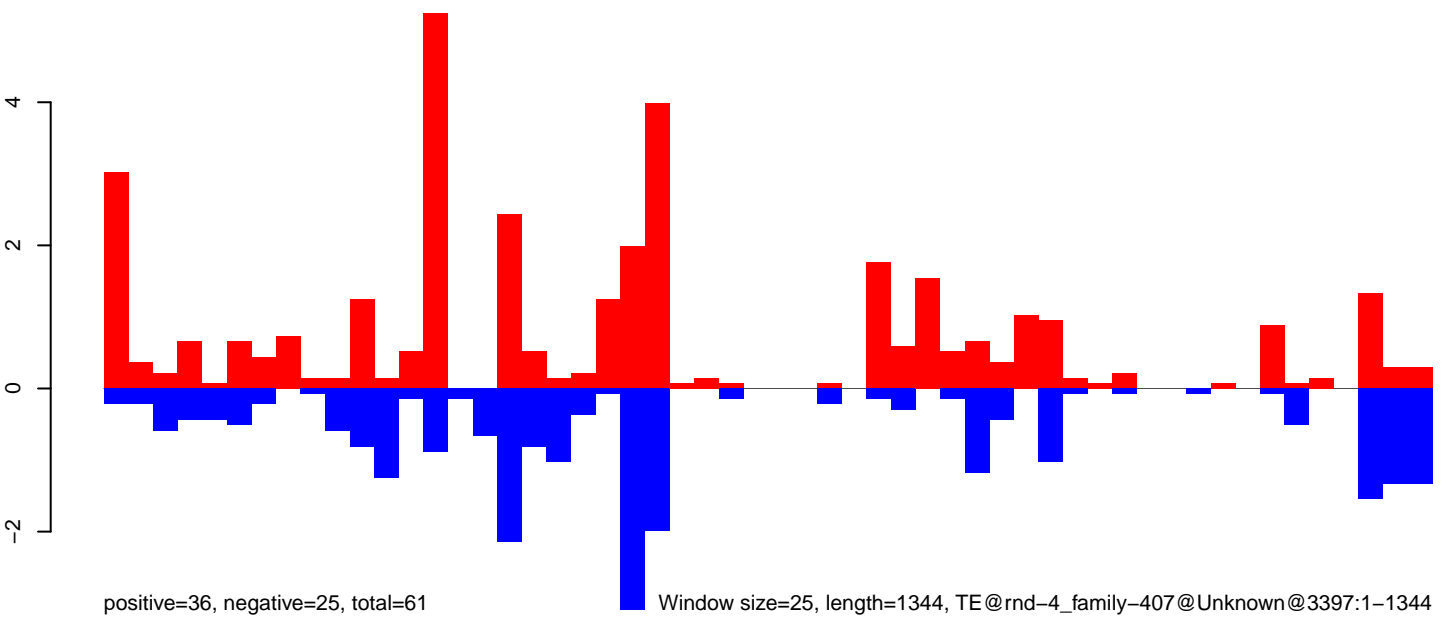
CuQuin_Testes_NL.18_23.rep



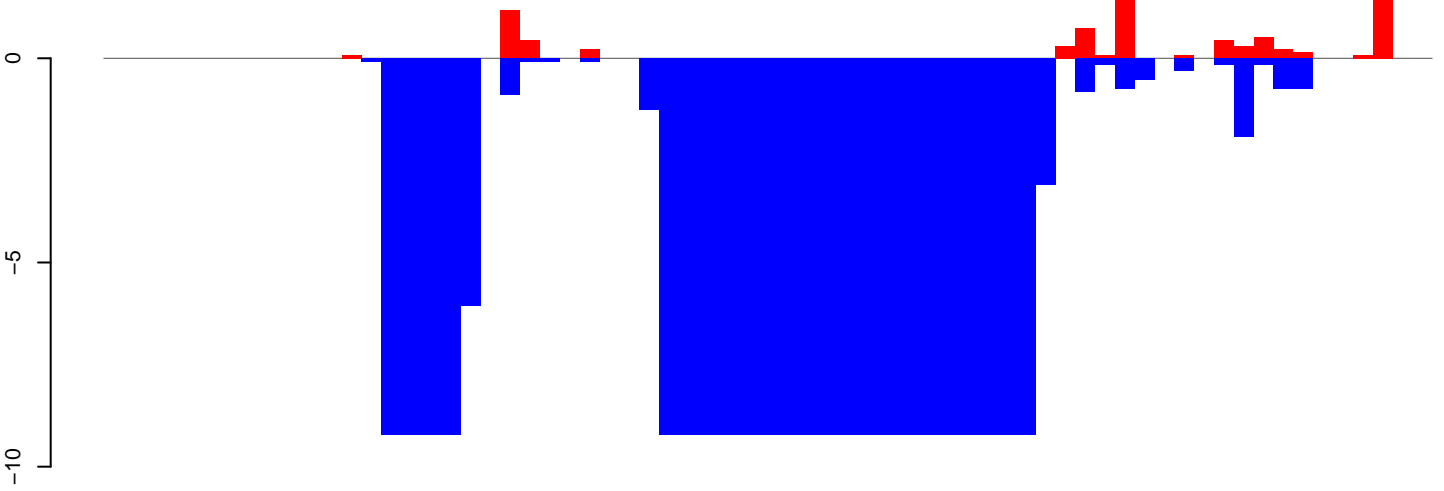
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

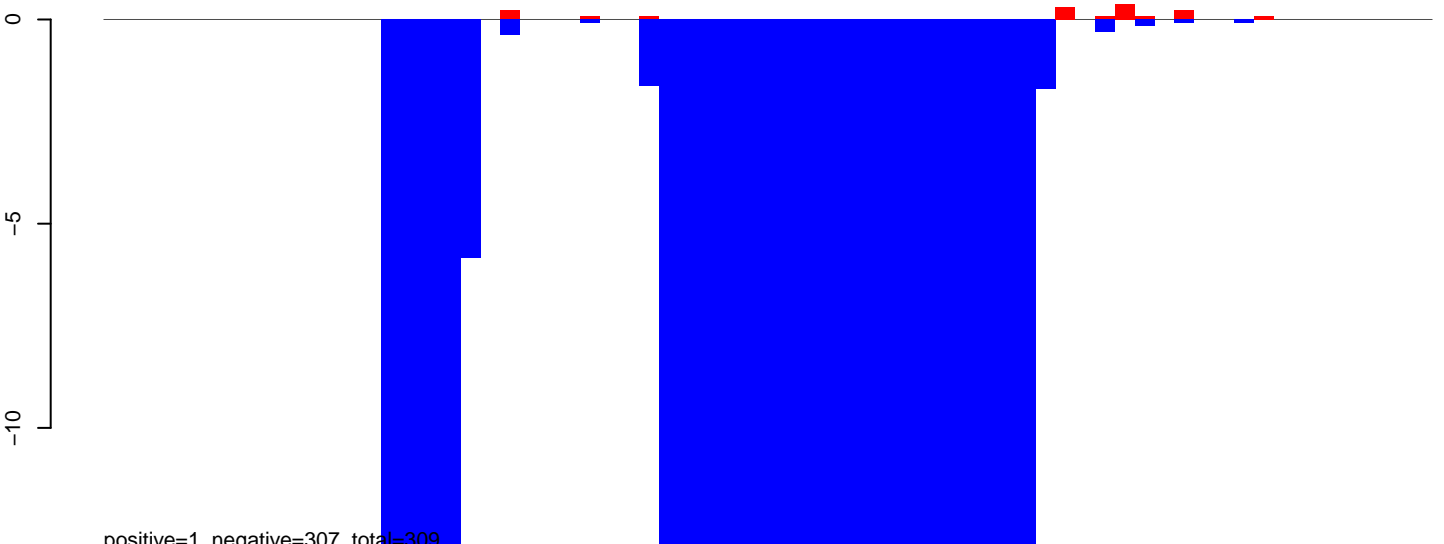


CuQuin_Testes_NL.18_23.rep



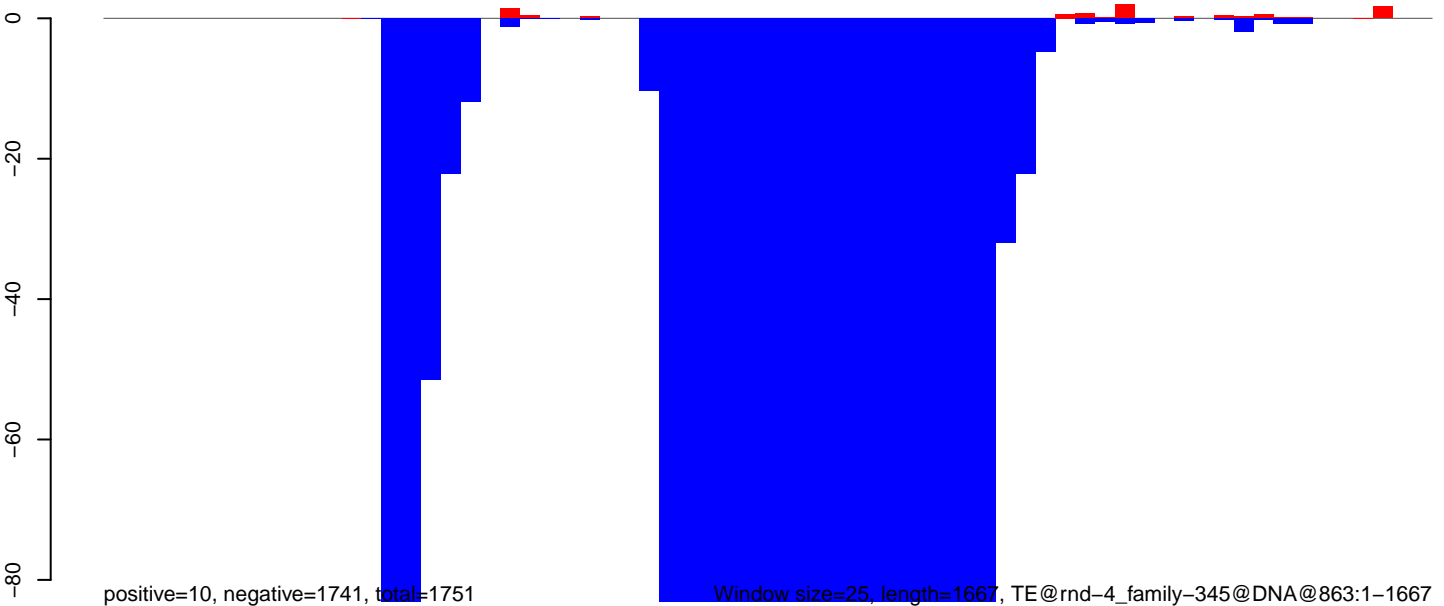
positive=8, negative=230, total=238

CuQuin_Testes_NL.24_35.rep



positive=1, negative=307, total=309

CuQuin_Testes_NL.rep

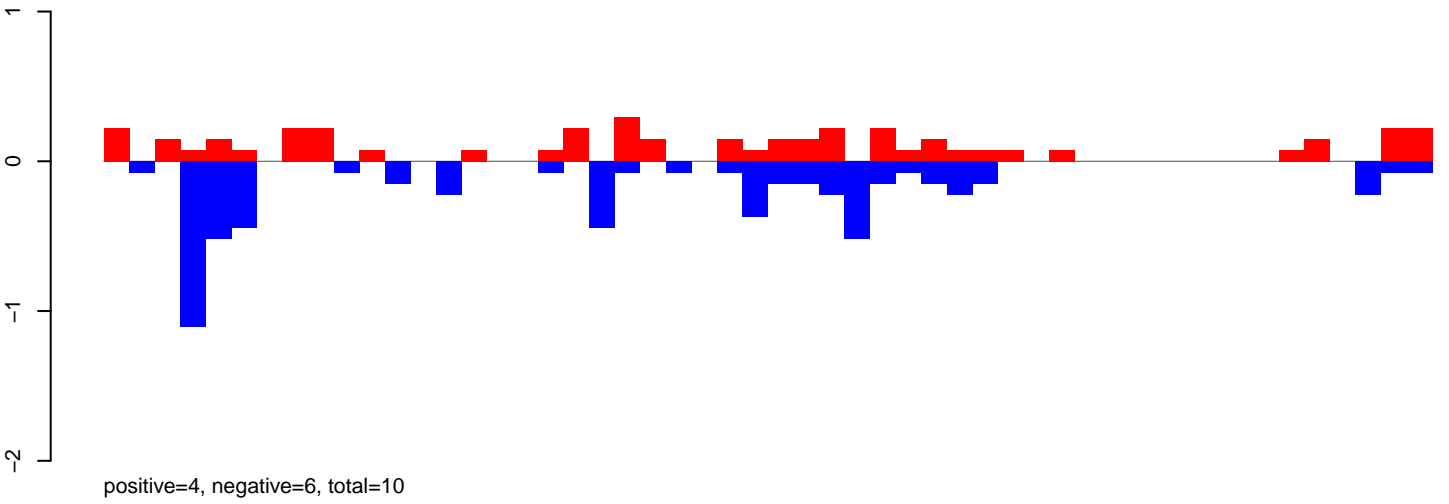


positive=10, negative=1741, total=1751

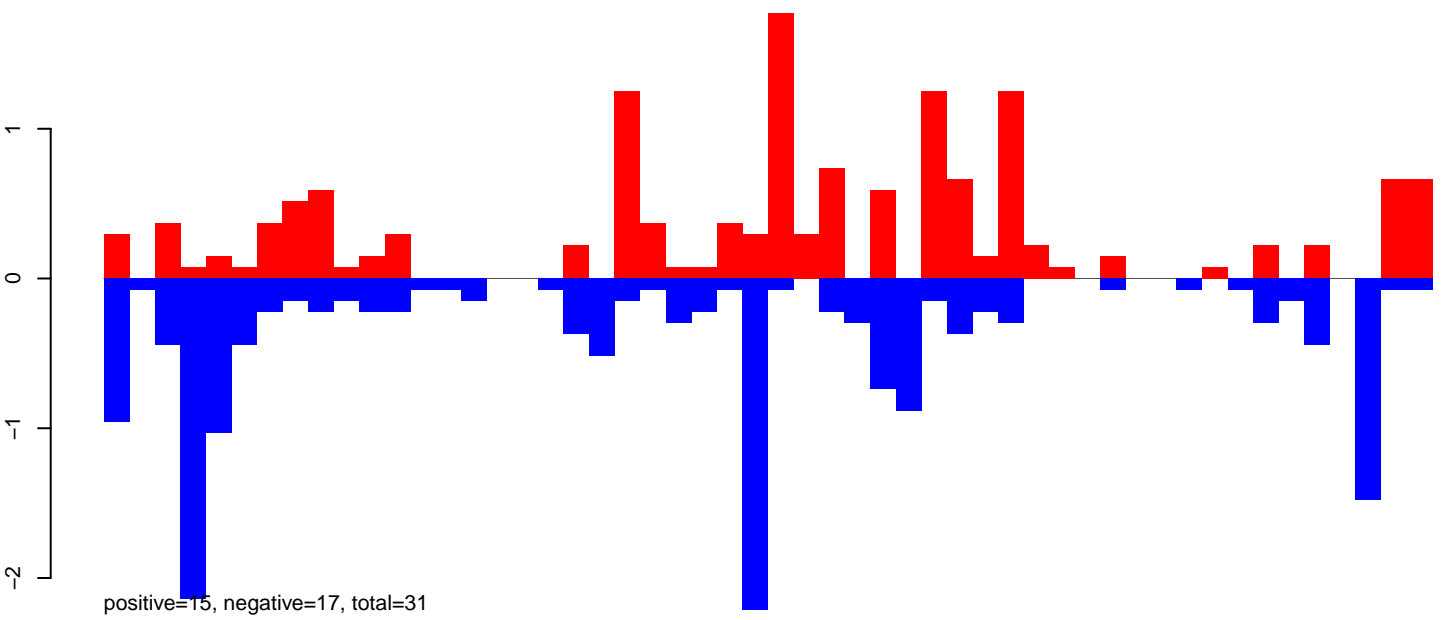
Window size=25, length=1667, TE@rnd-4_family-345@DNA@863:1-1667

0 500 1000 1500

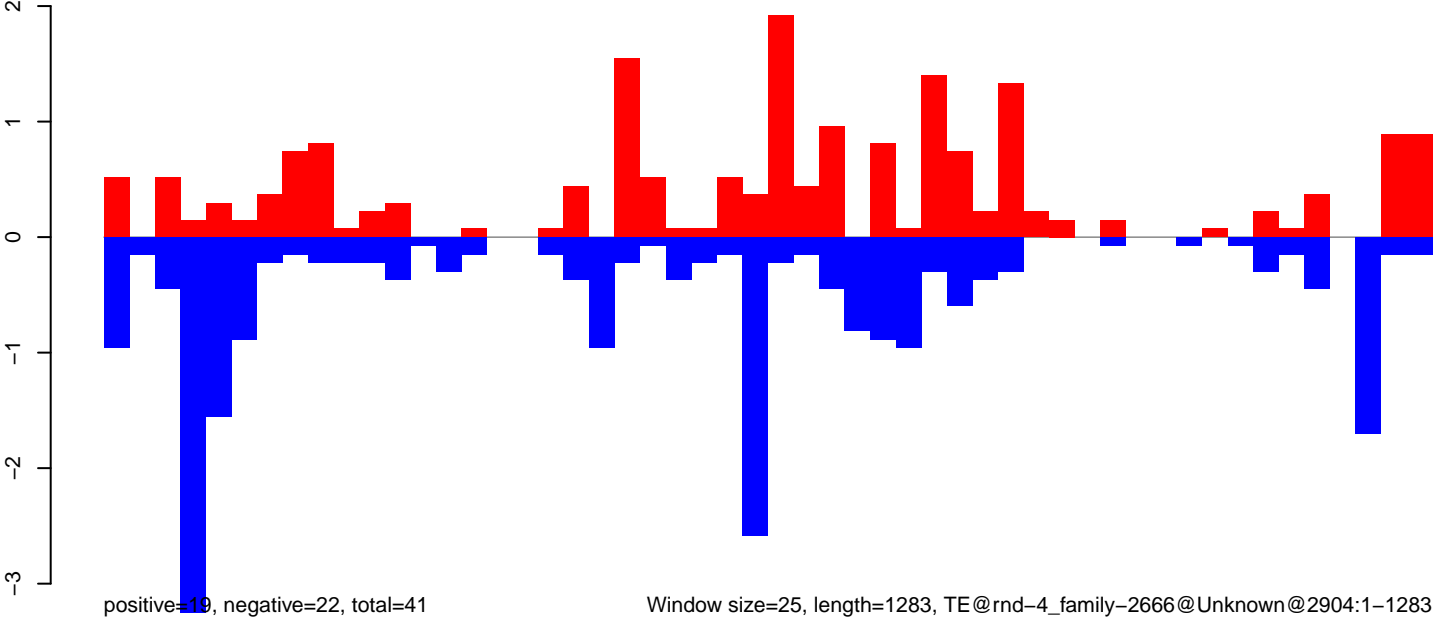
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

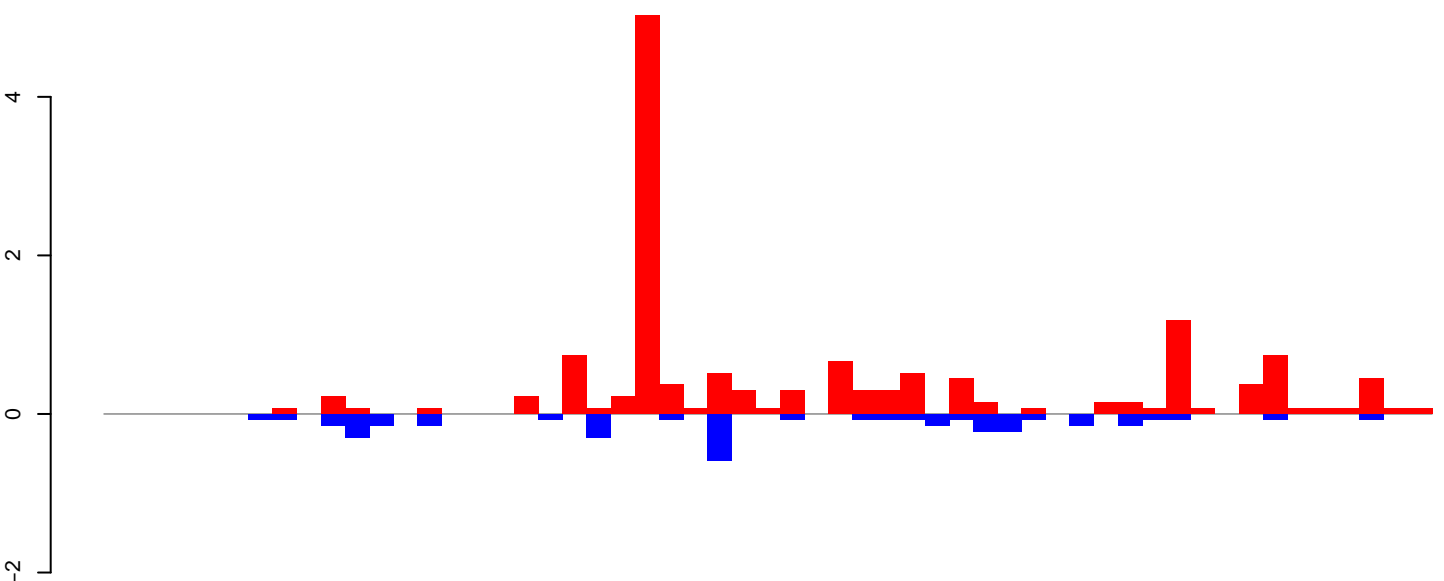


CuQuin_Testes_NL.rep



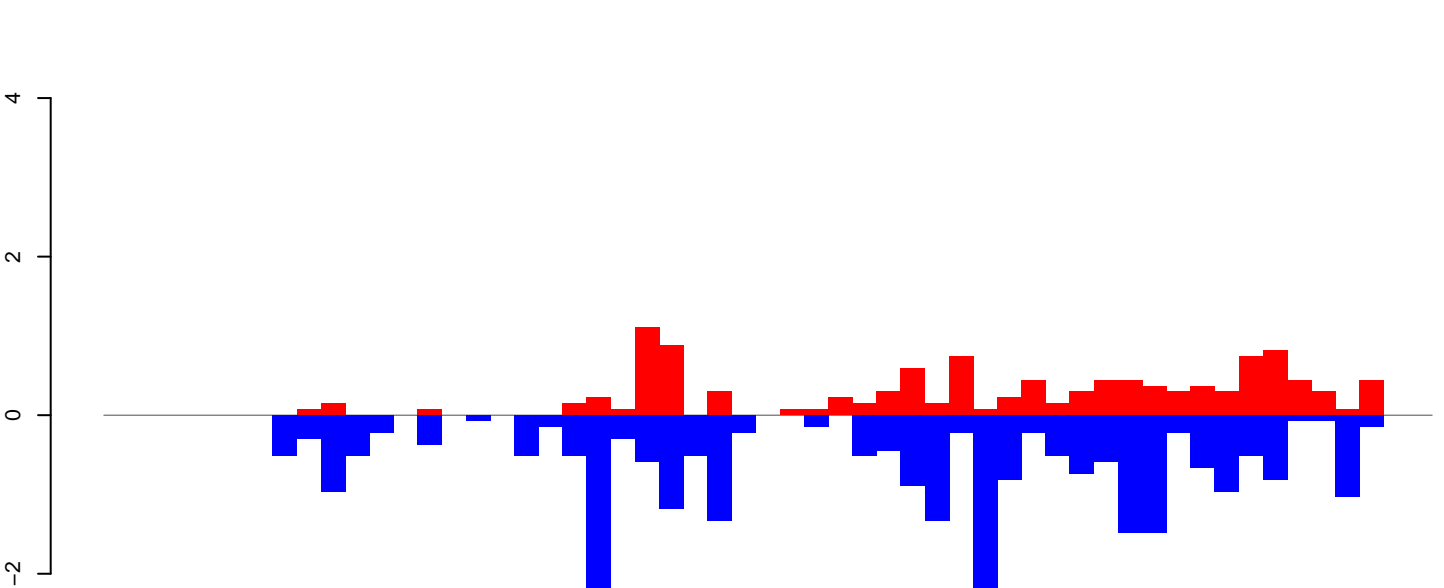
Window size=25, length=1283, TE@rnd-4_family-2666@Unknown@2904:1-1283

CuQuin_Testes_NL.18_23.rep



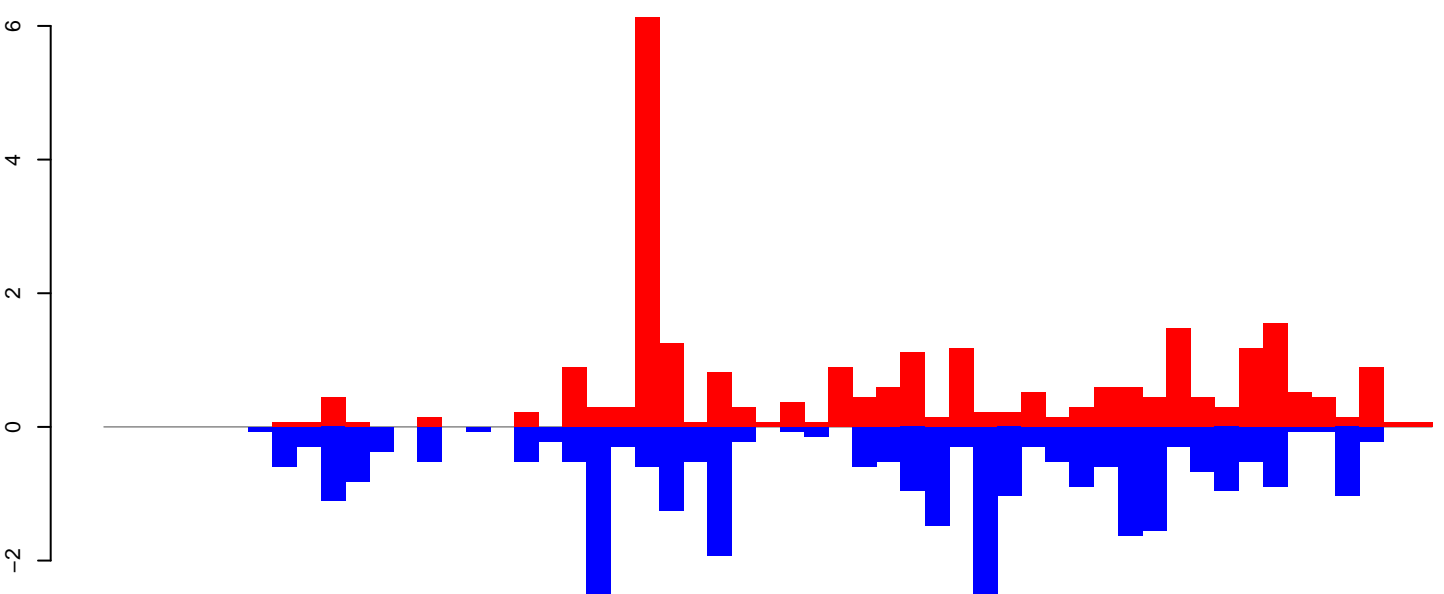
positive=14, negative=4, total=18

CuQuin_Testes_NL.24_35.rep



positive=12, negative=27, total=39

CuQuin_Testes_NL.rep

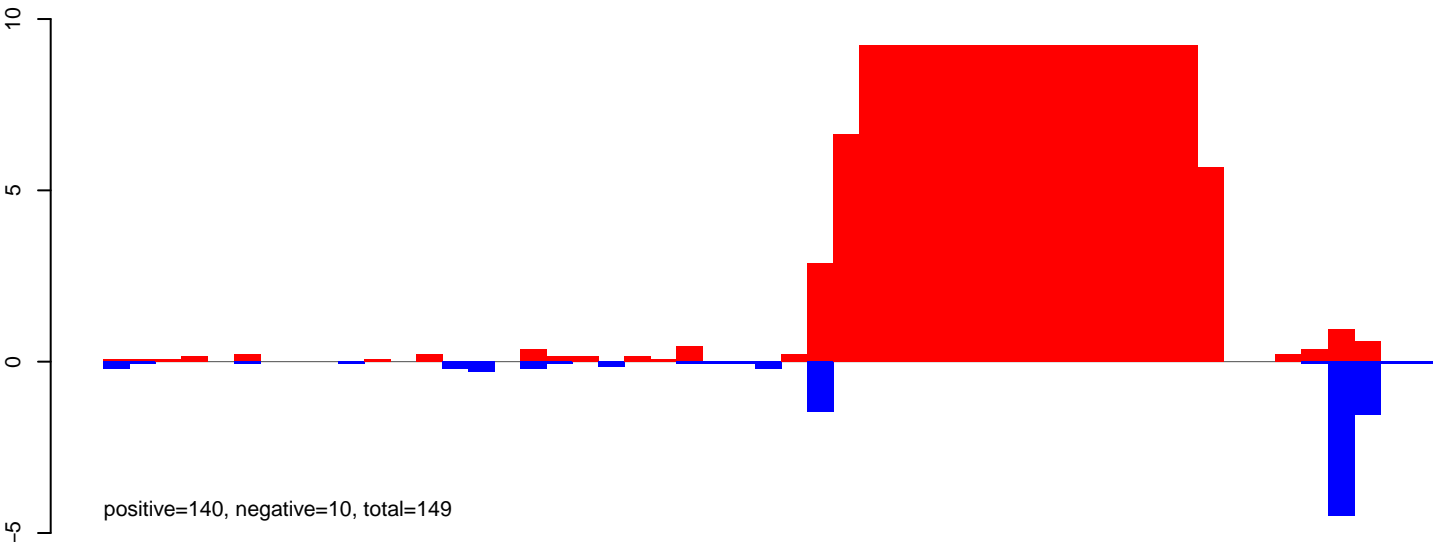


positive=26, negative=31, total=57

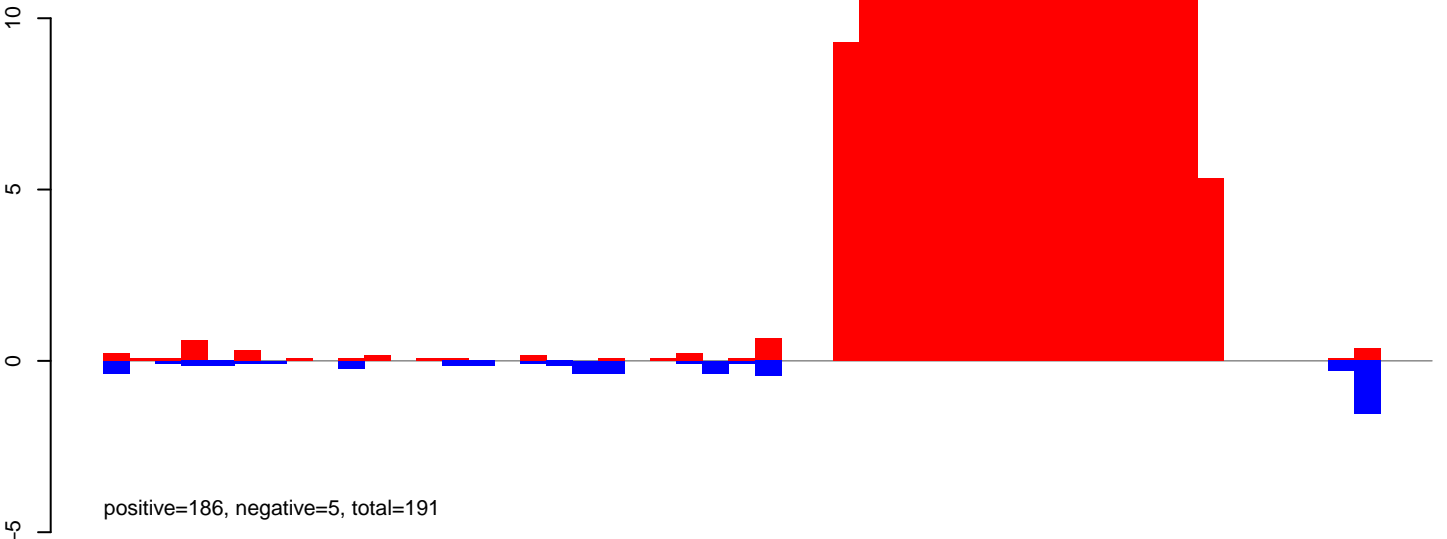
Window size=25, length=1373, TE@rnd-4_family-124@Unknown@962:1-1373

0 200 400 600 800 1000 1200 1400

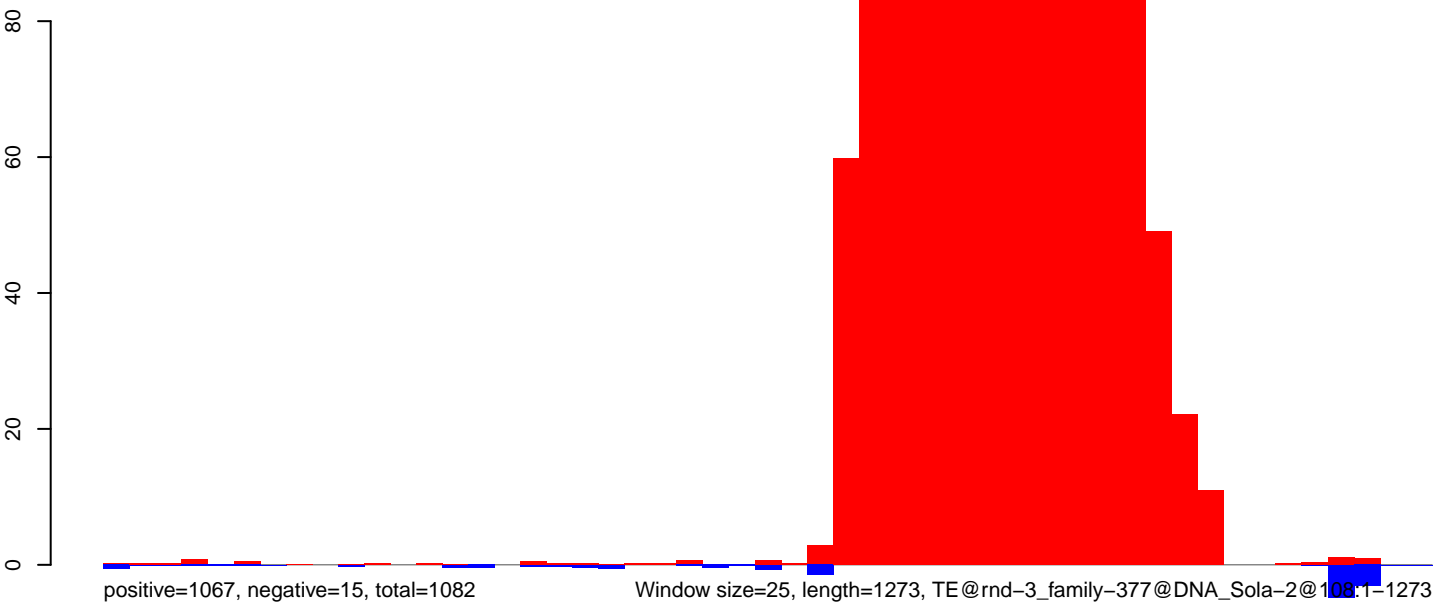
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

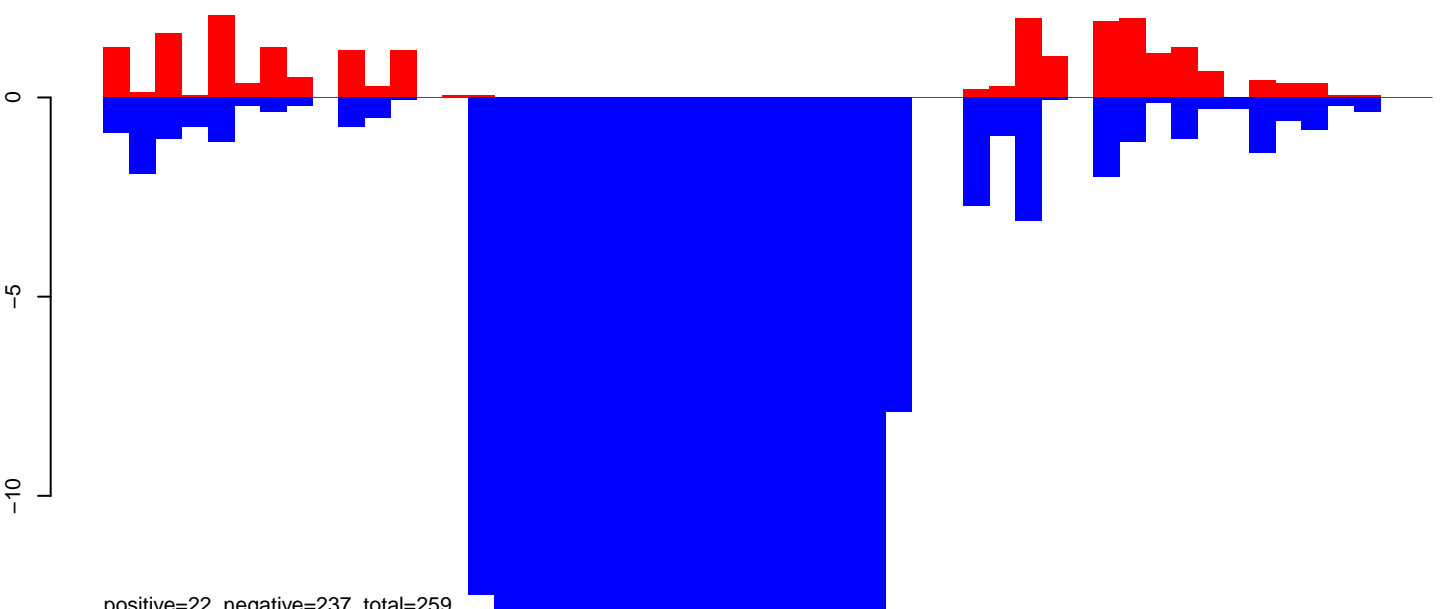


CuQuin_Testes_NL.18_23.rep



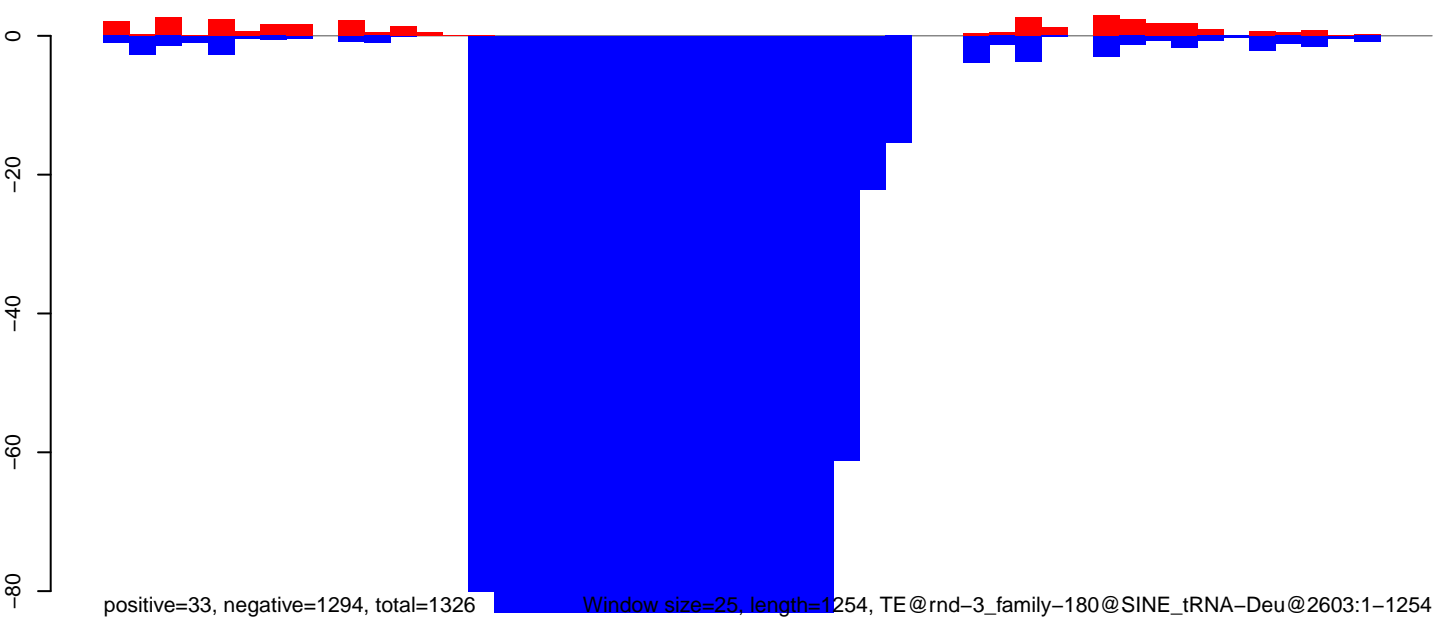
positive=11, negative=167, total=177

CuQuin_Testes_NL.24_35.rep



positive=22, negative=237, total=259

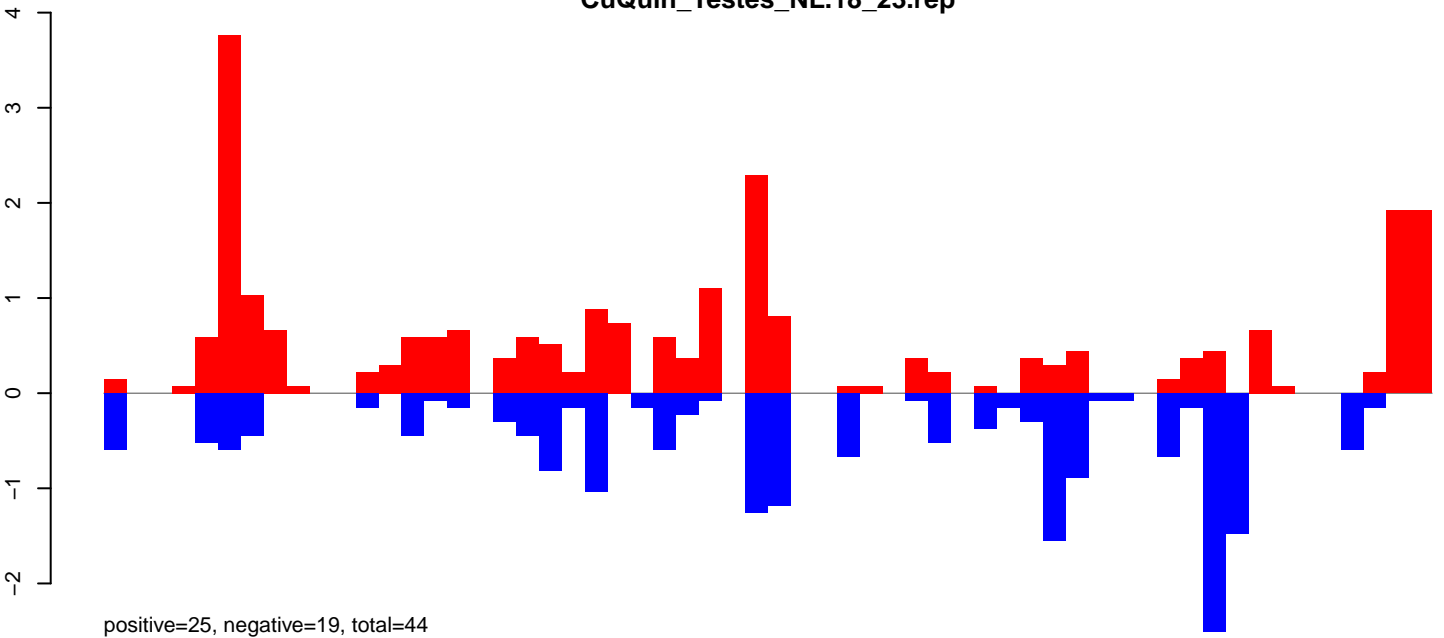
CuQuin_Testes_NL.rep



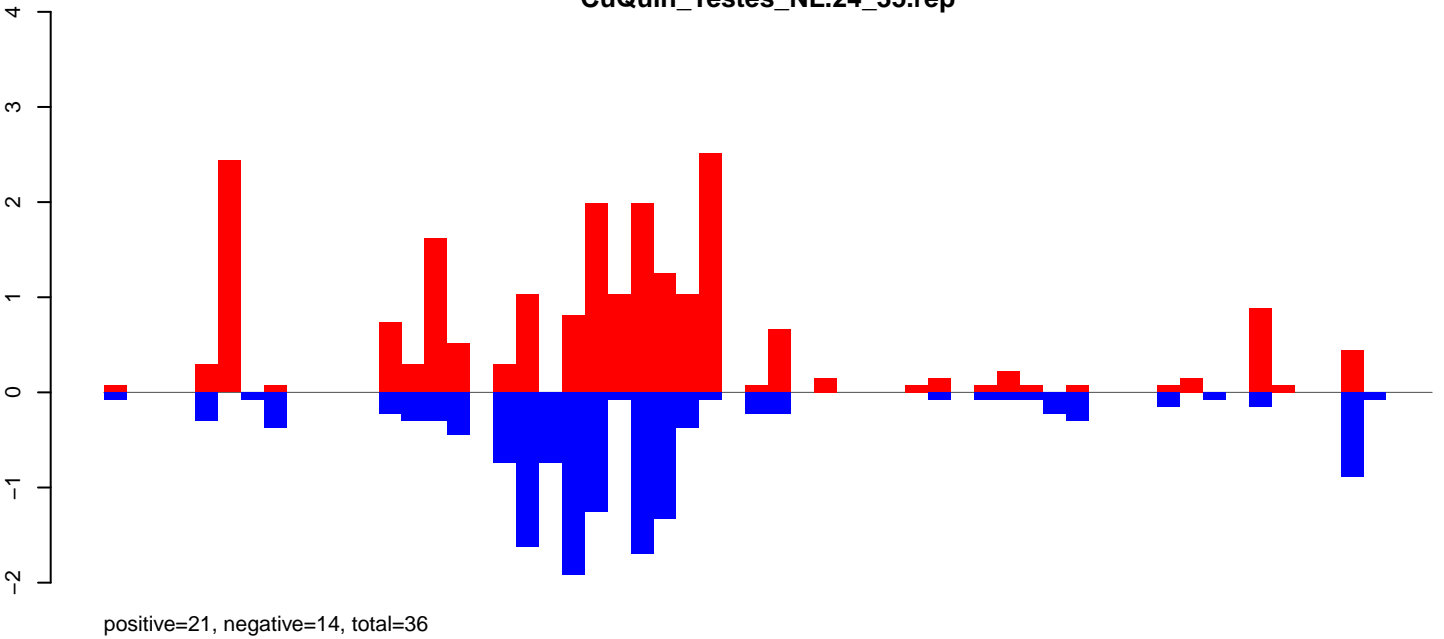
positive=33, negative=1294, total=1326

Window size=25, length=1254, TE@rnd-3_family-180@SINE_tRNA-Deu@2603:1-1254

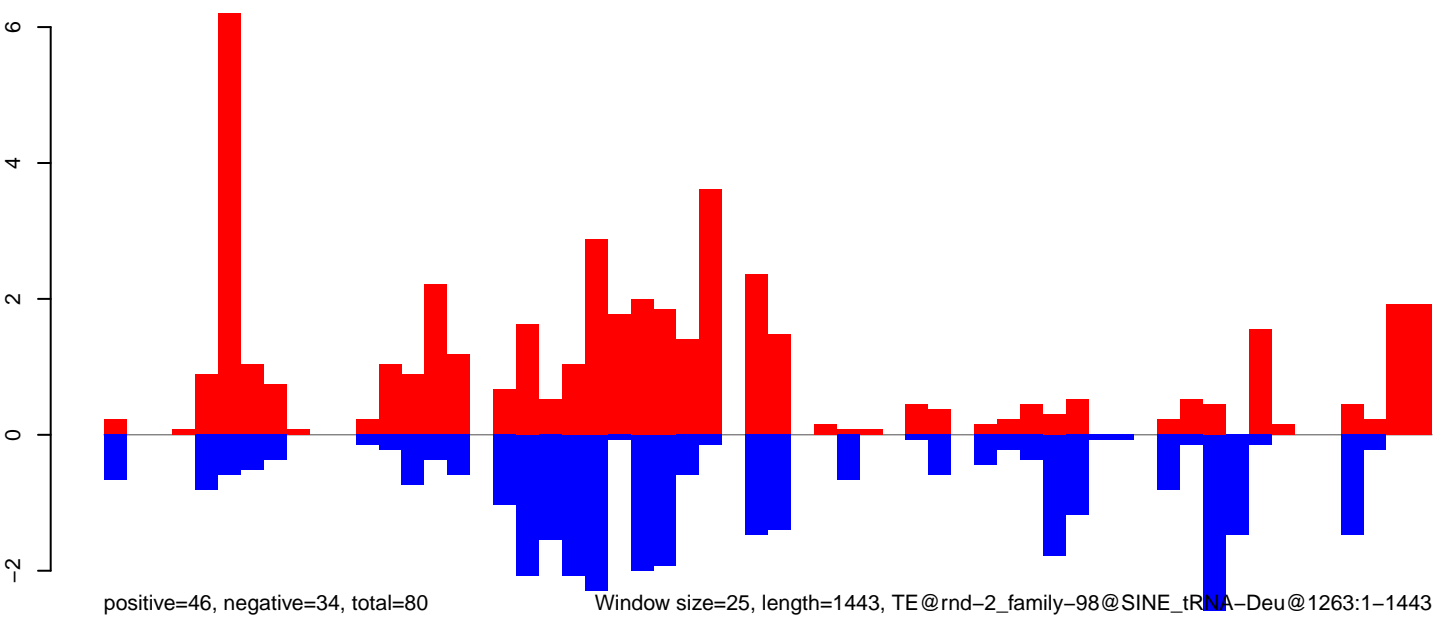
CuQuin_Testes_NL.18_23.rep



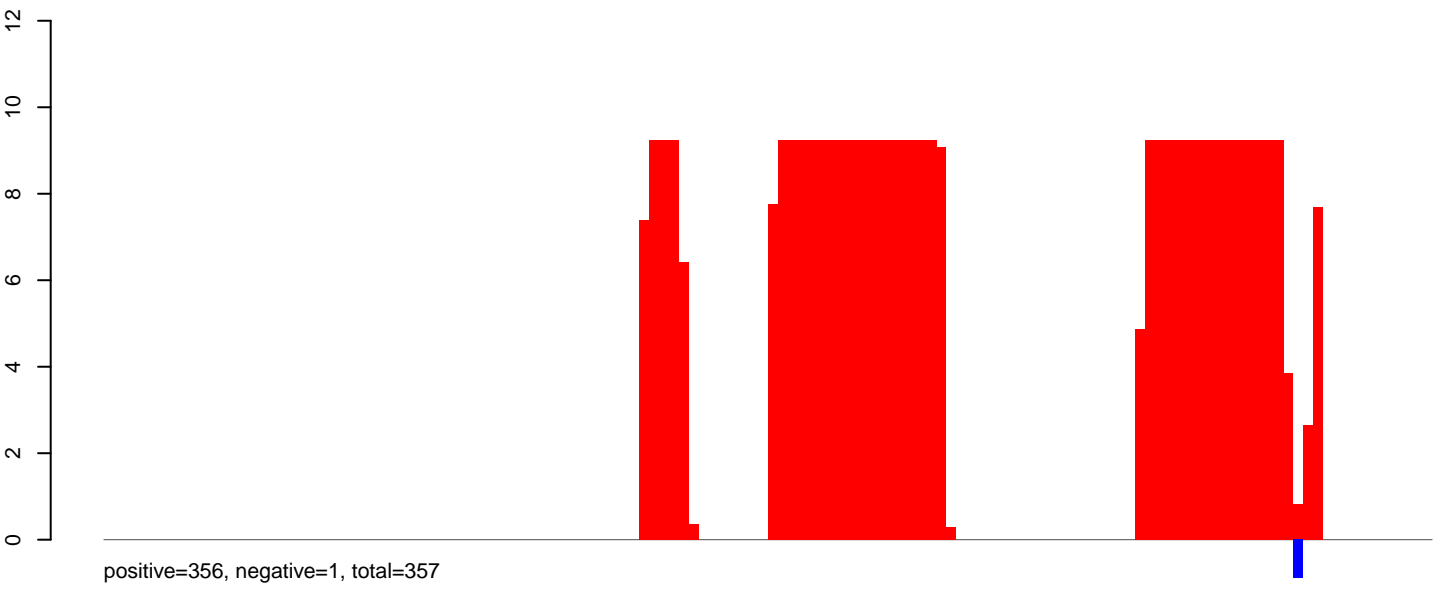
CuQuin_Testes_NL.24_35.rep



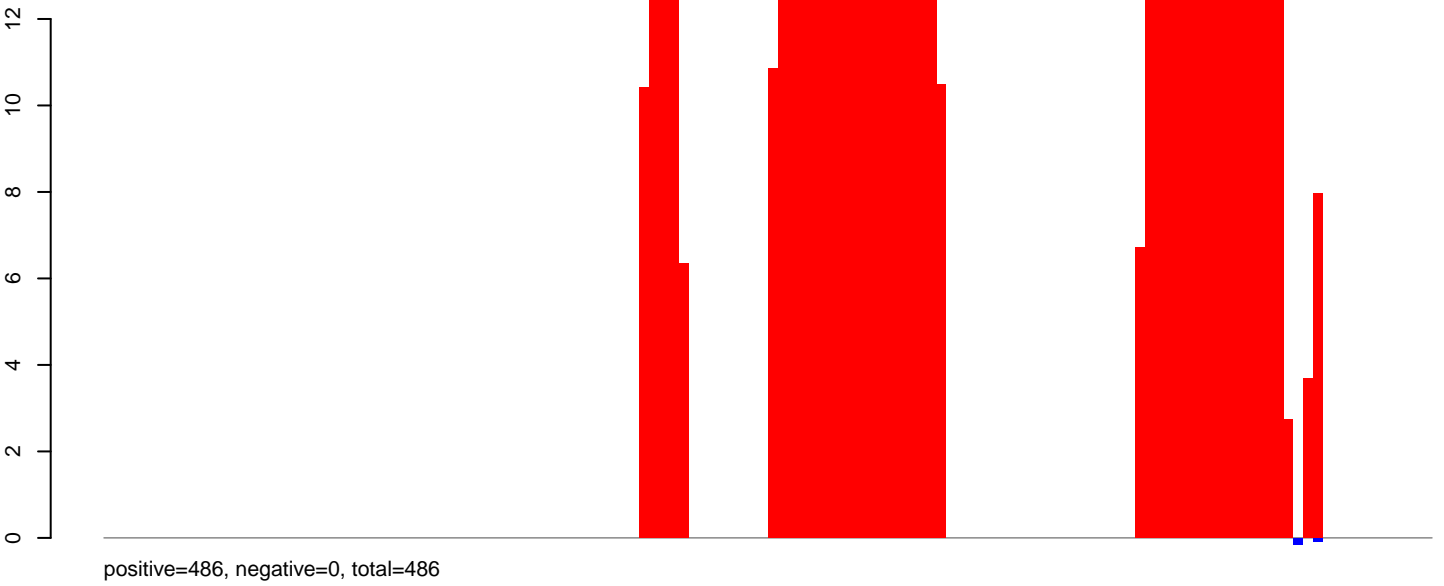
CuQuin_Testes_NL.rep



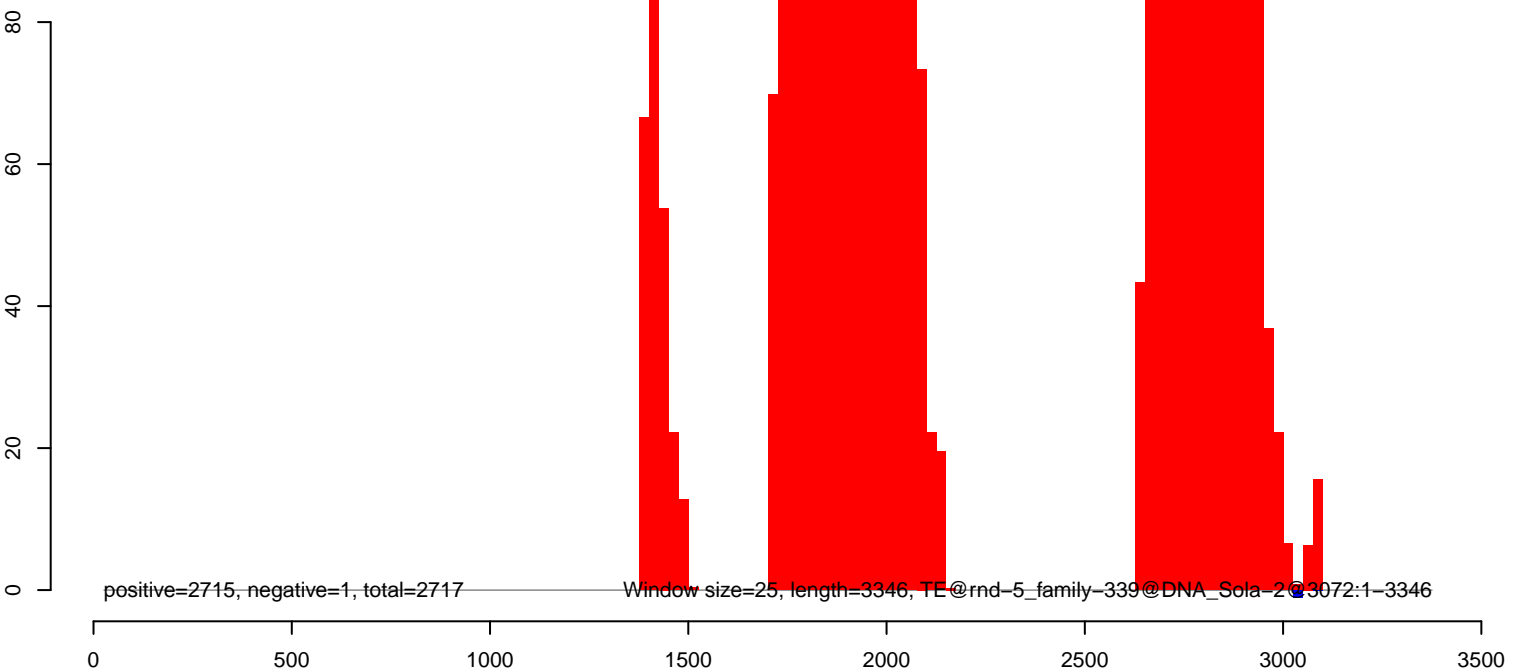
CuQuin_Testes_NL.18_23.rep



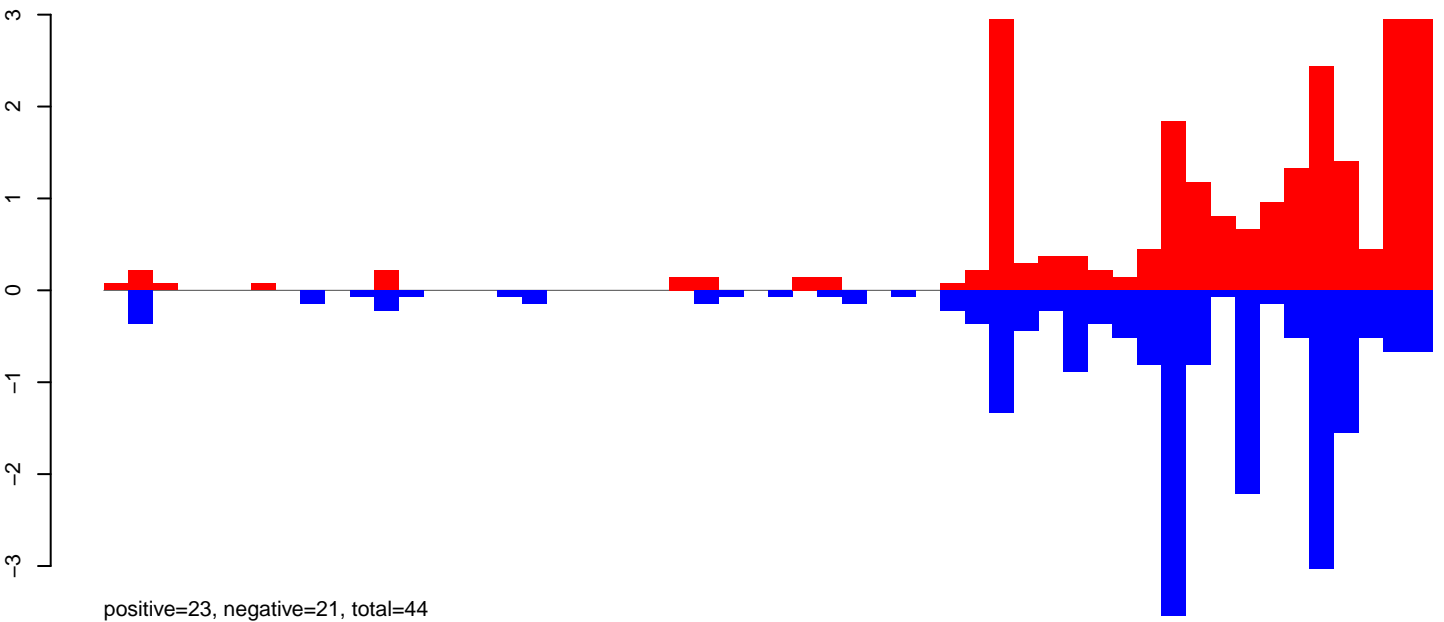
CuQuin_Testes_NL.24_35.rep



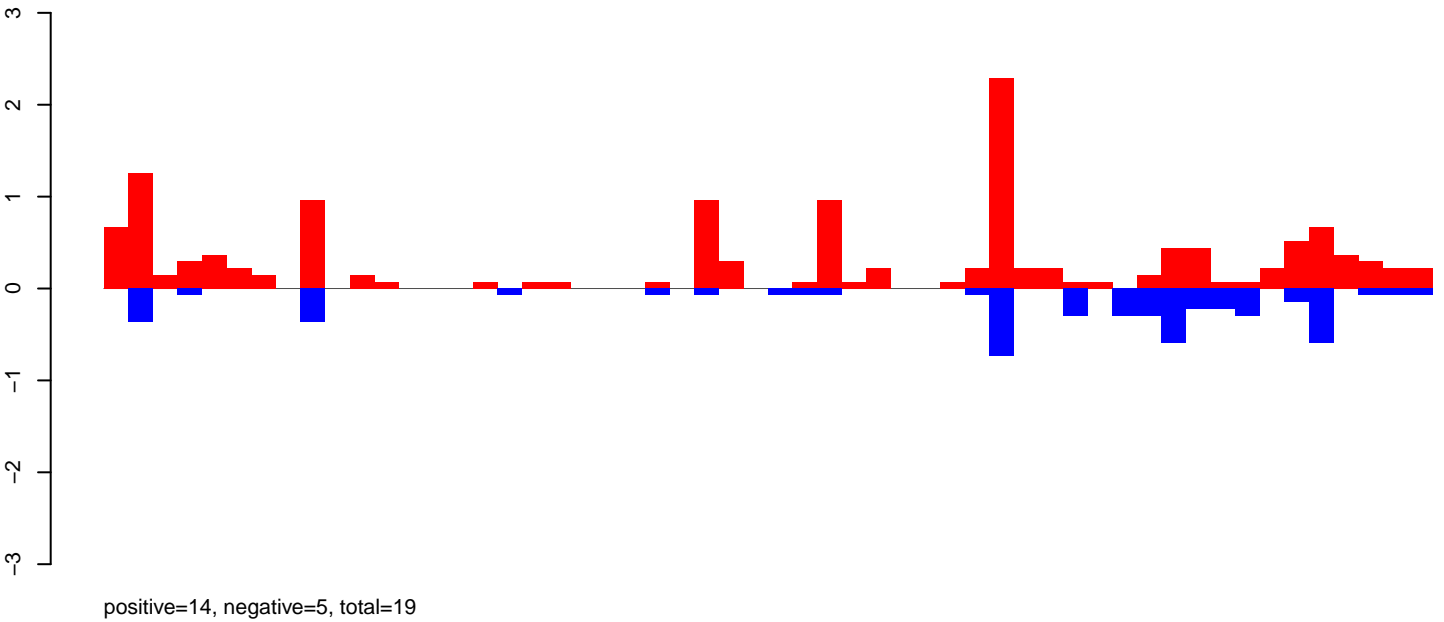
CuQuin_Testes_NL.rep



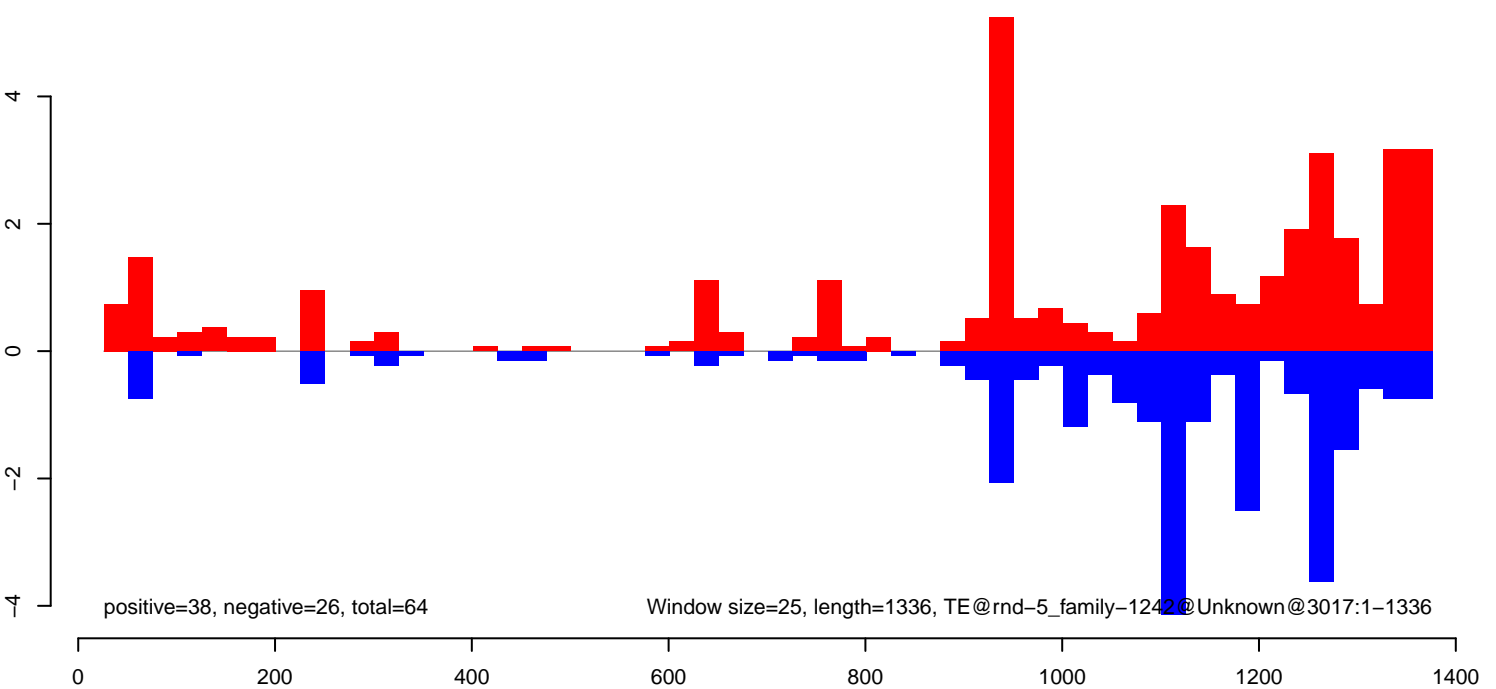
CuQuin_Testes_NL.18_23.rep



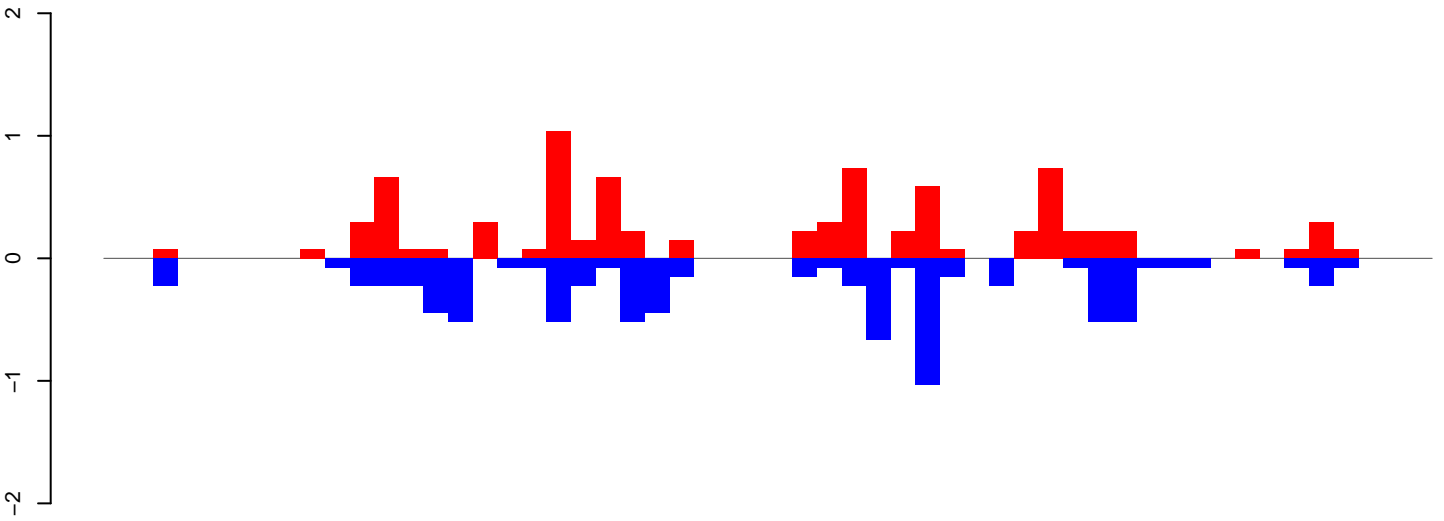
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

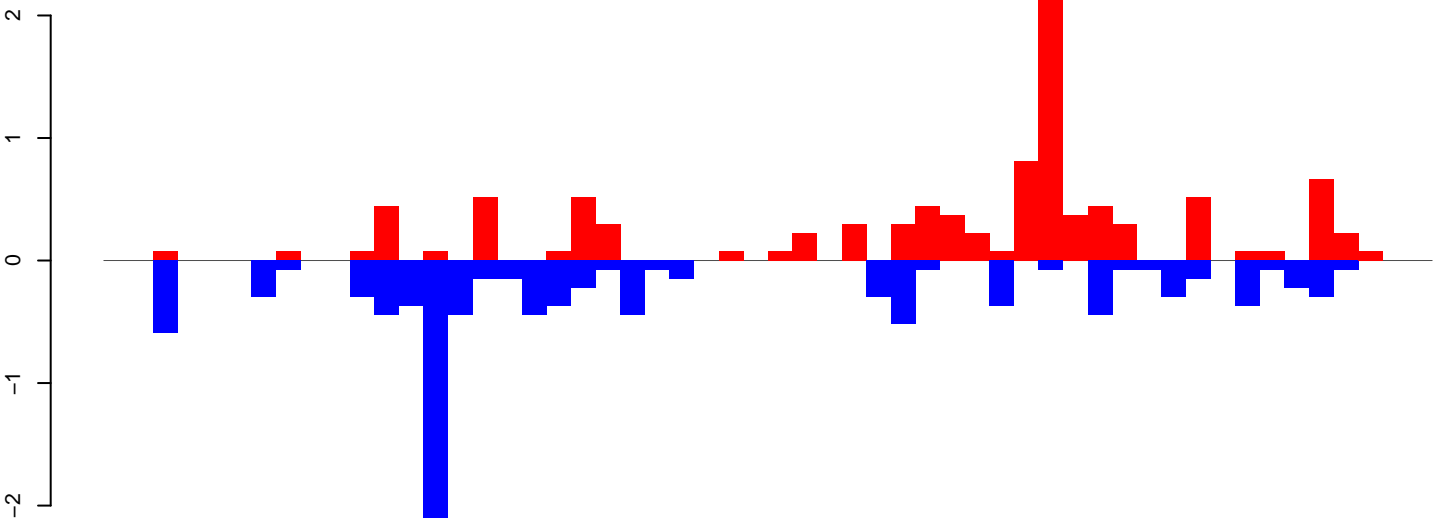


CuQuin_Testes_NL.18_23.rep



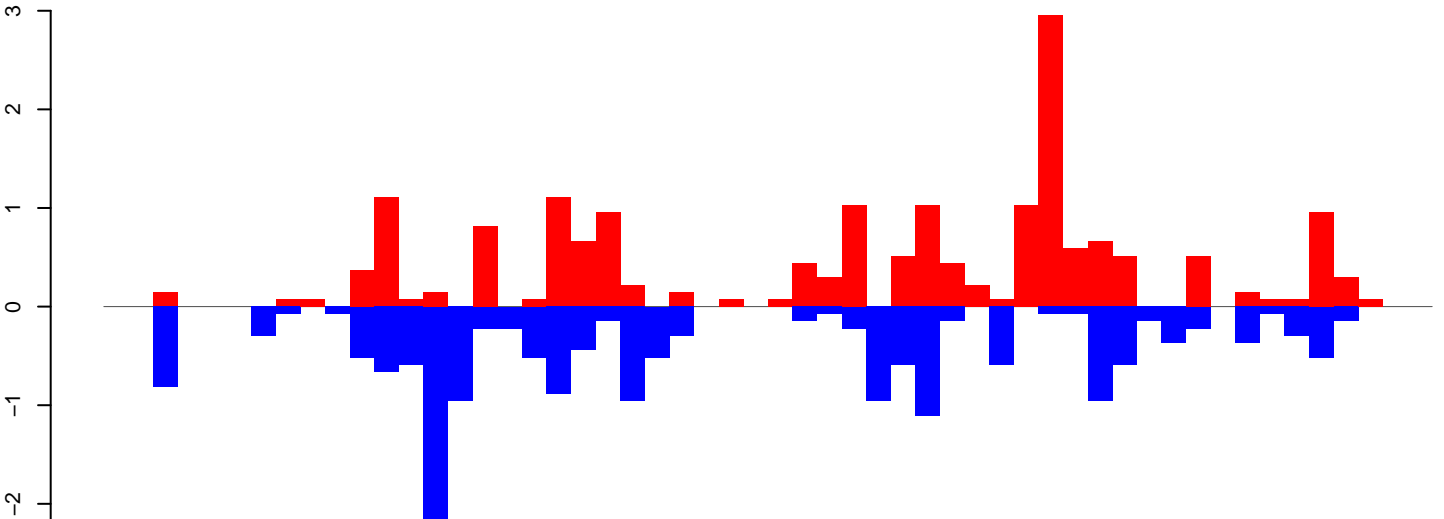
positive=8, negative=8, total=16

CuQuin_Testes_NL.24_35.rep



positive=10, negative=11, total=21

CuQuin_Testes_NL.rep

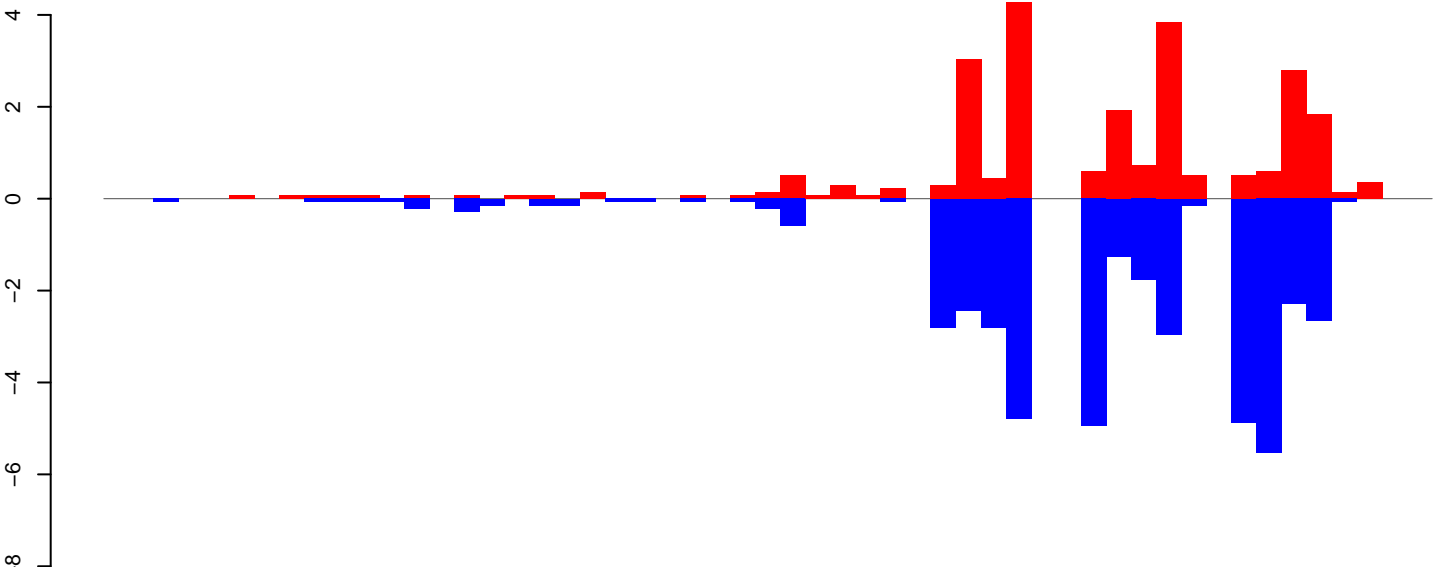


positive=18, negative=19, total=37

Window size=25, length=1327, TE@rnd-5_family-11@Unknown@2332:1-1327

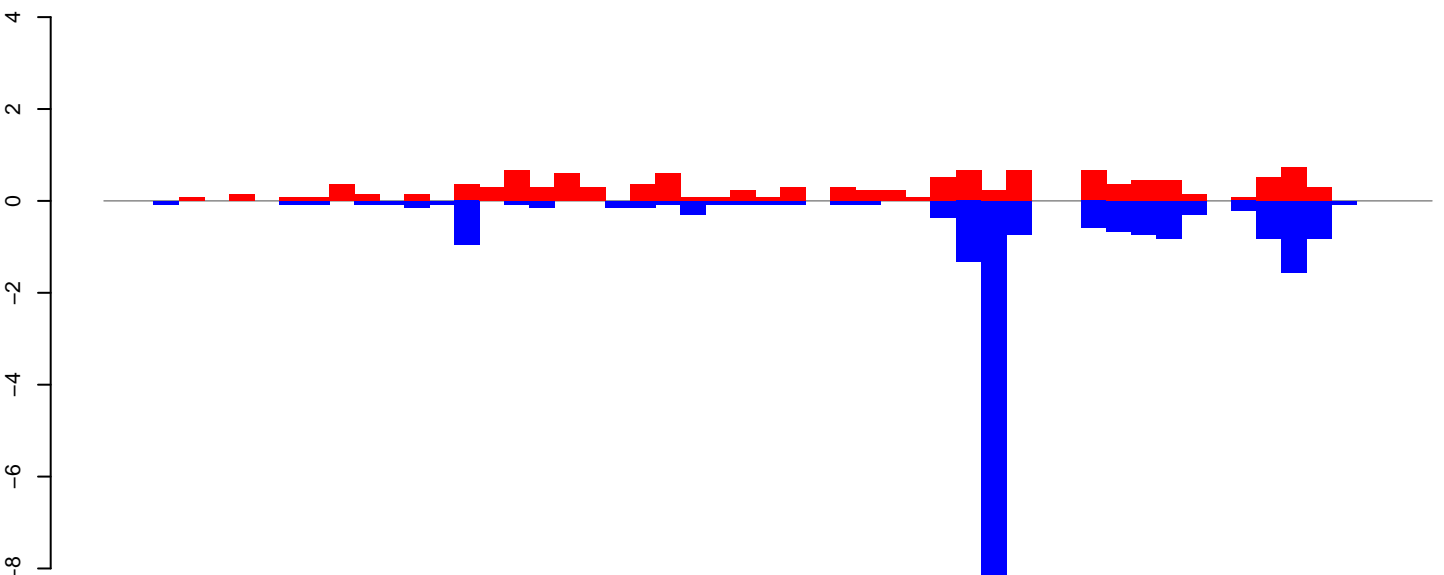
0 200 400 600 800 1000 1200 1400

CuQuin_Testes_NL.18_23.rep



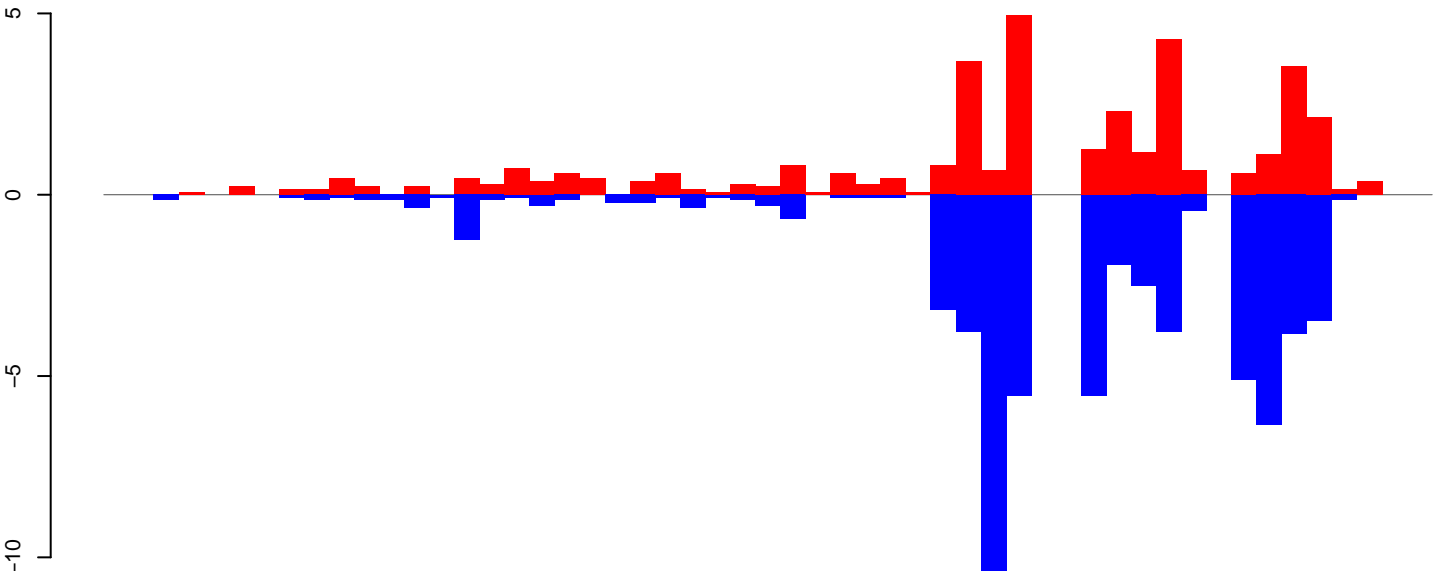
positive=24, negative=42, total=66

CuQuin_Testes_NL.24_35.rep



positive=12, negative=21, total=32

CuQuin_Testes_NL.rep

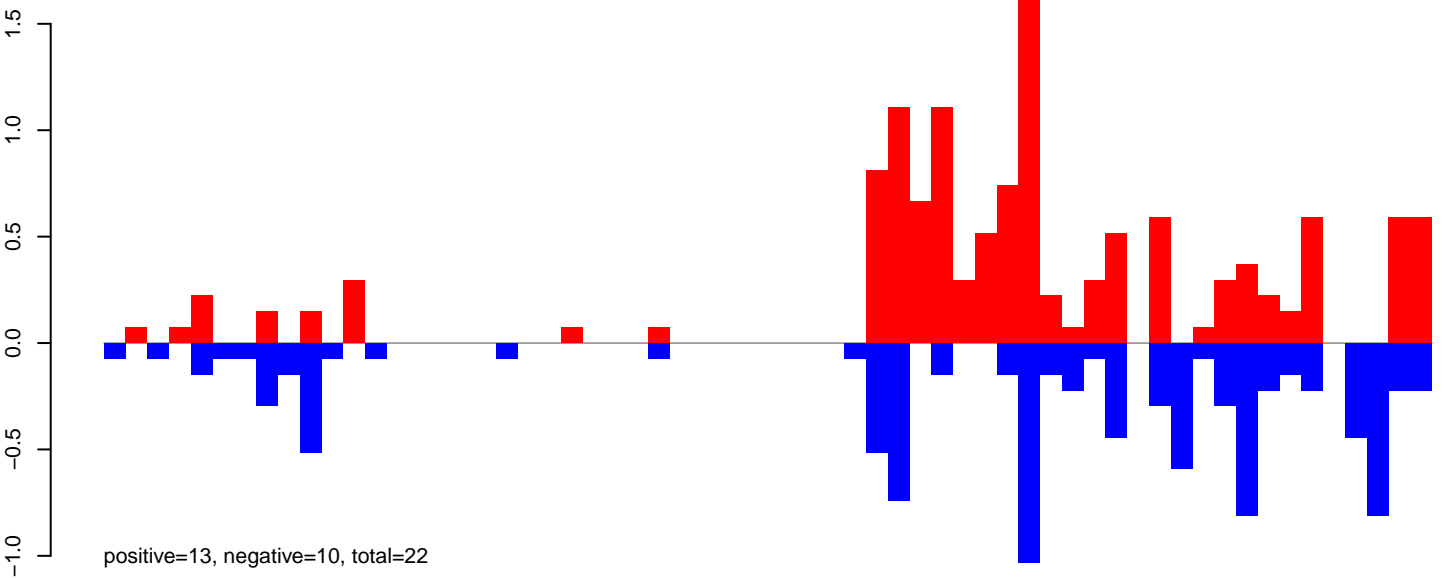


positive=36, negative=62, total=98

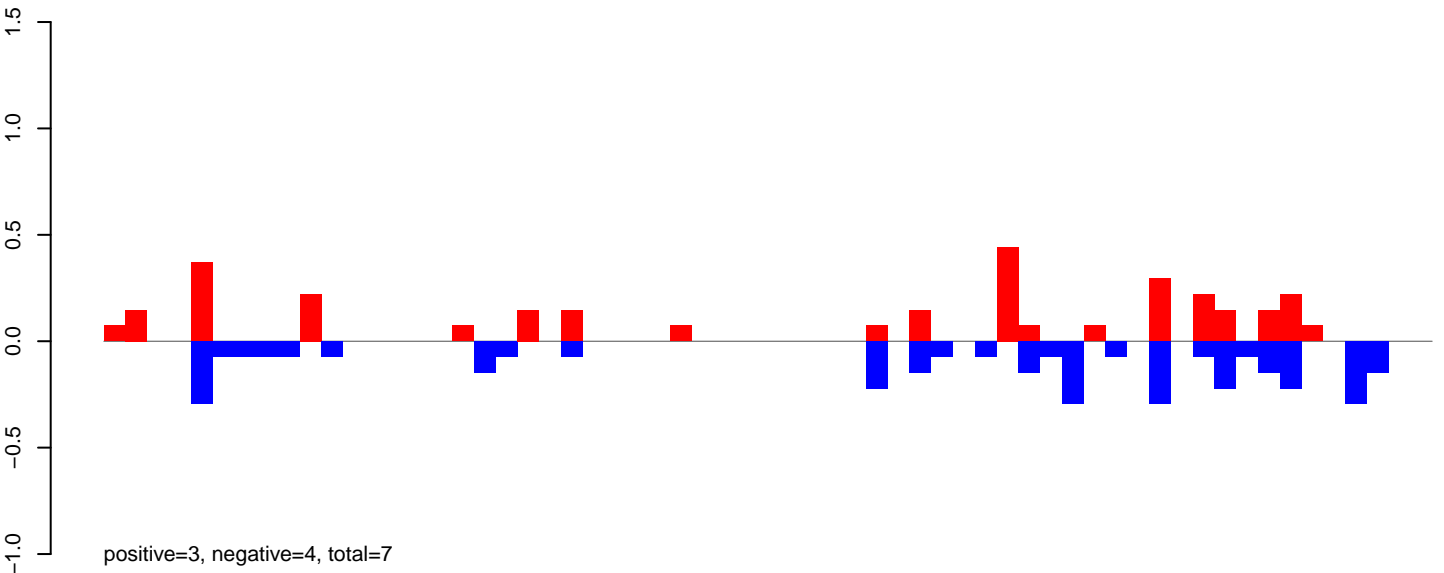
Window size=25, length=1324, TE@rnd-5_family-1002@Satellite@1811:1-1324

0 200 400 600 800 1000 1200 1400

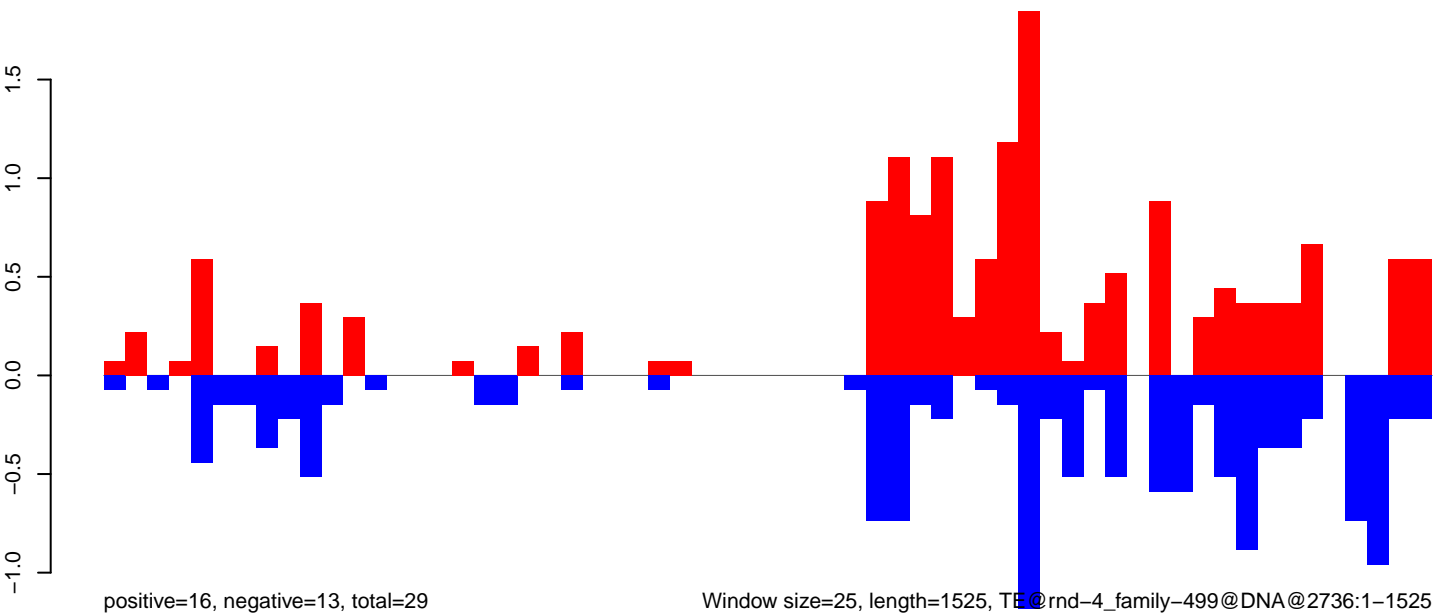
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



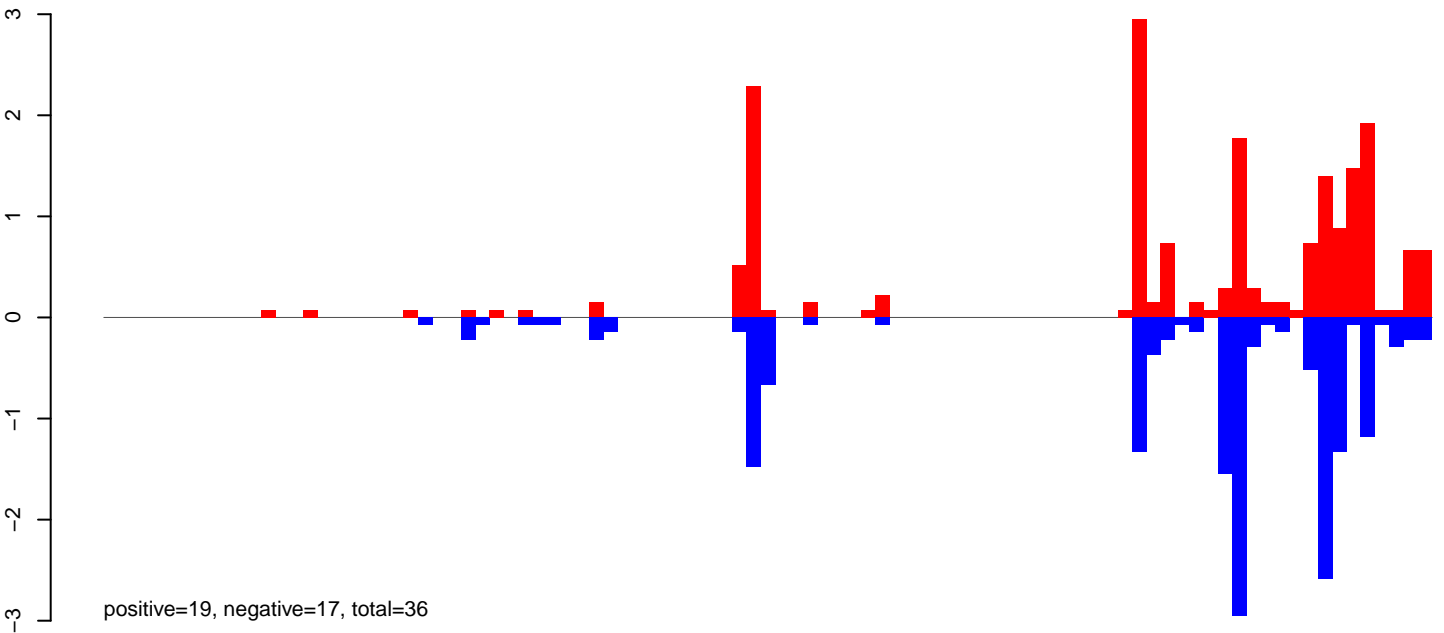
CuQuin_Testes_NL.rep



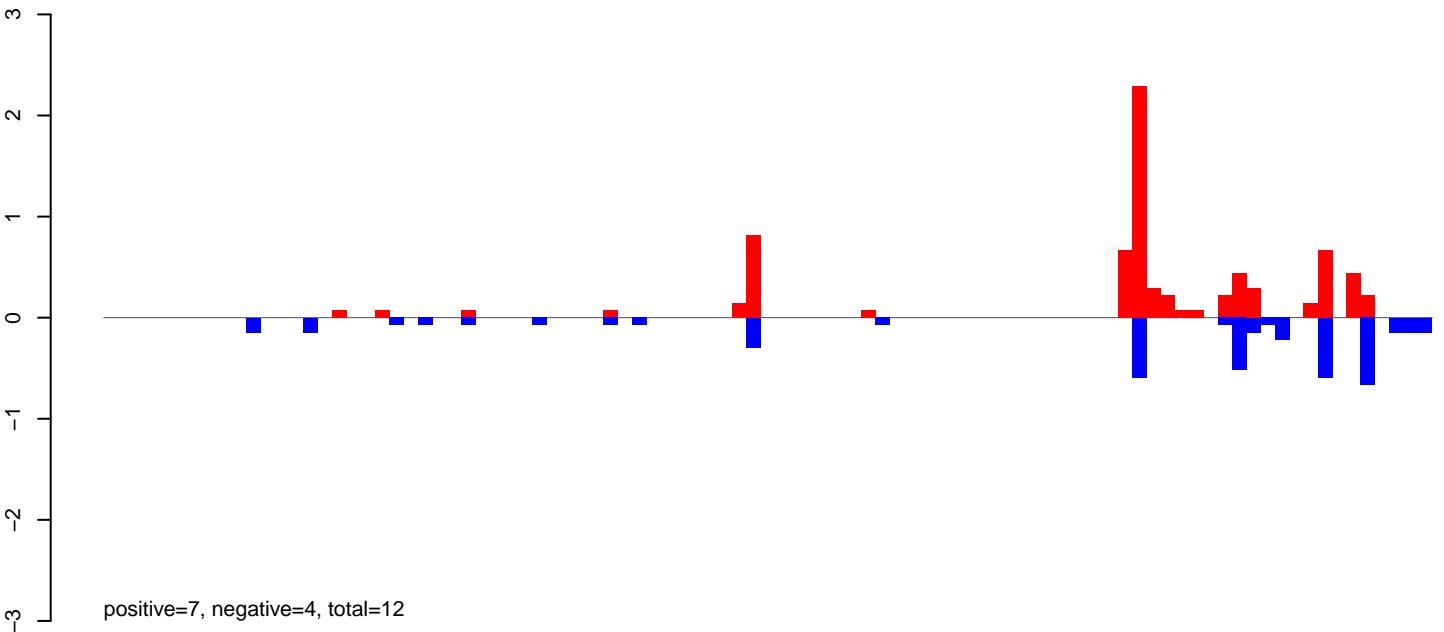
Window size=25, length=1525, TE @ rnd-4_family-499@DNA@2736:1-1525

0 500 1000 1500

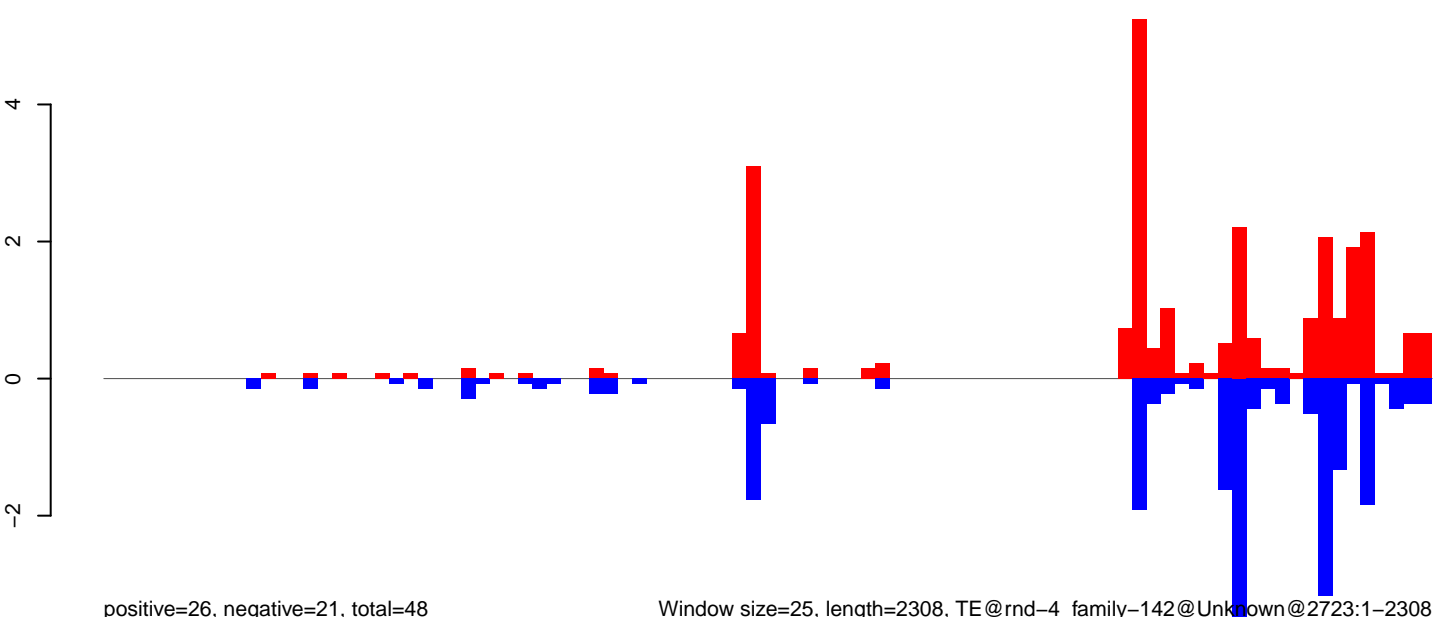
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



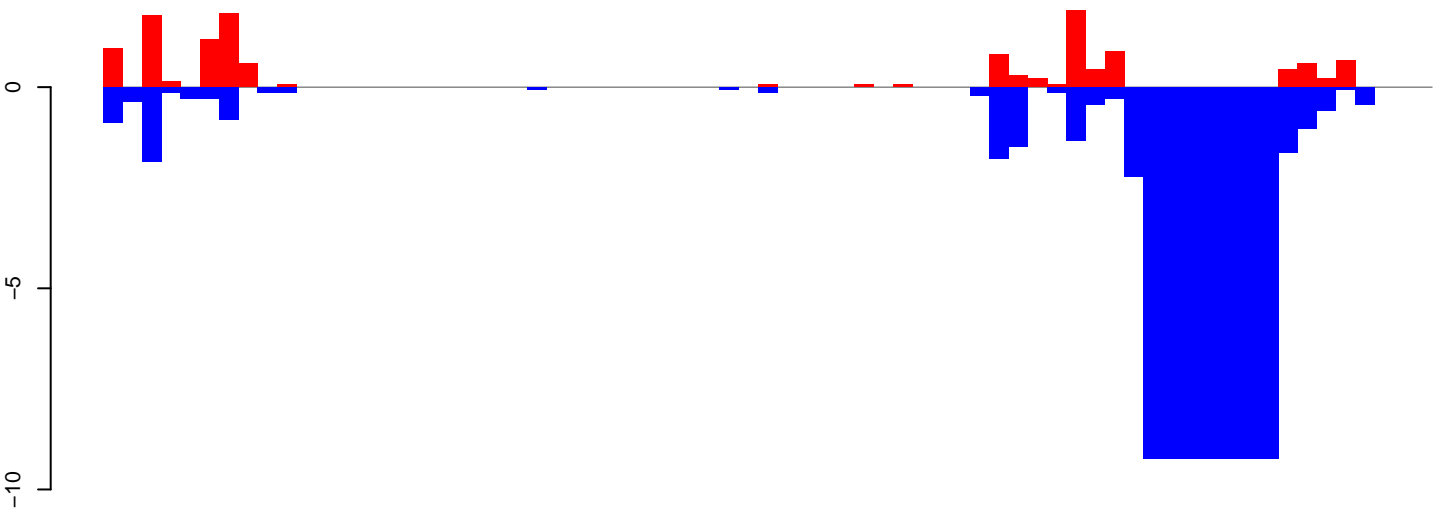
CuQuin_Testes_NL.rep



Window size=25, length=2308, TE@rnd-4_family-142@Unknown@2723:1-2308

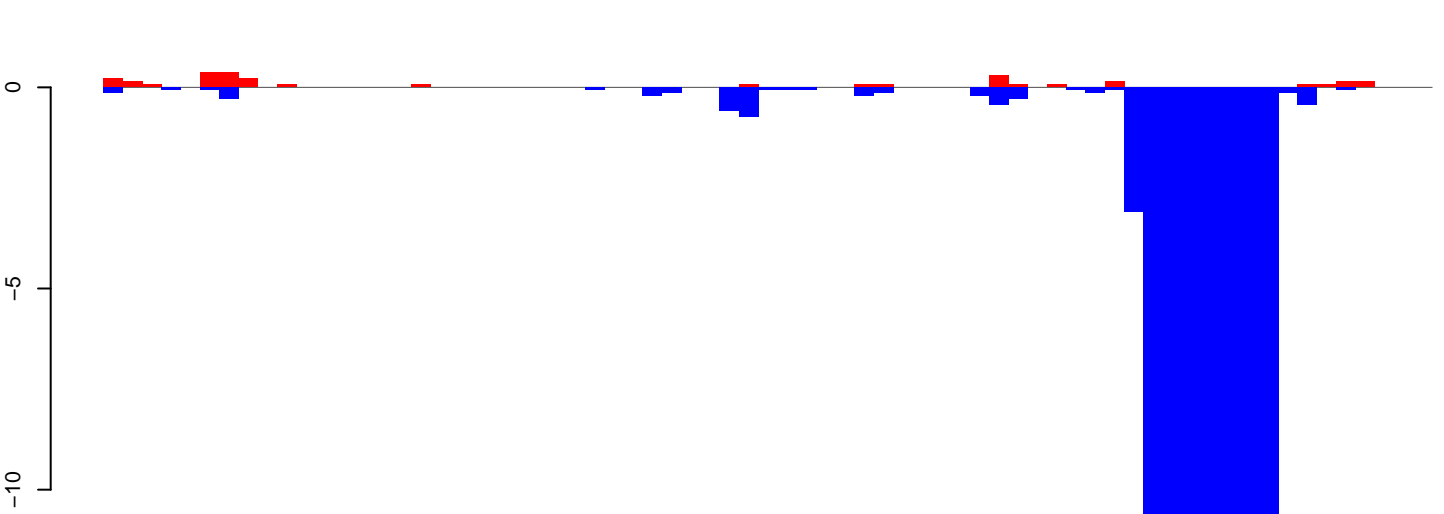
0 500 1000 1500 2000

CuQuin_Testes_NL.18_23.rep



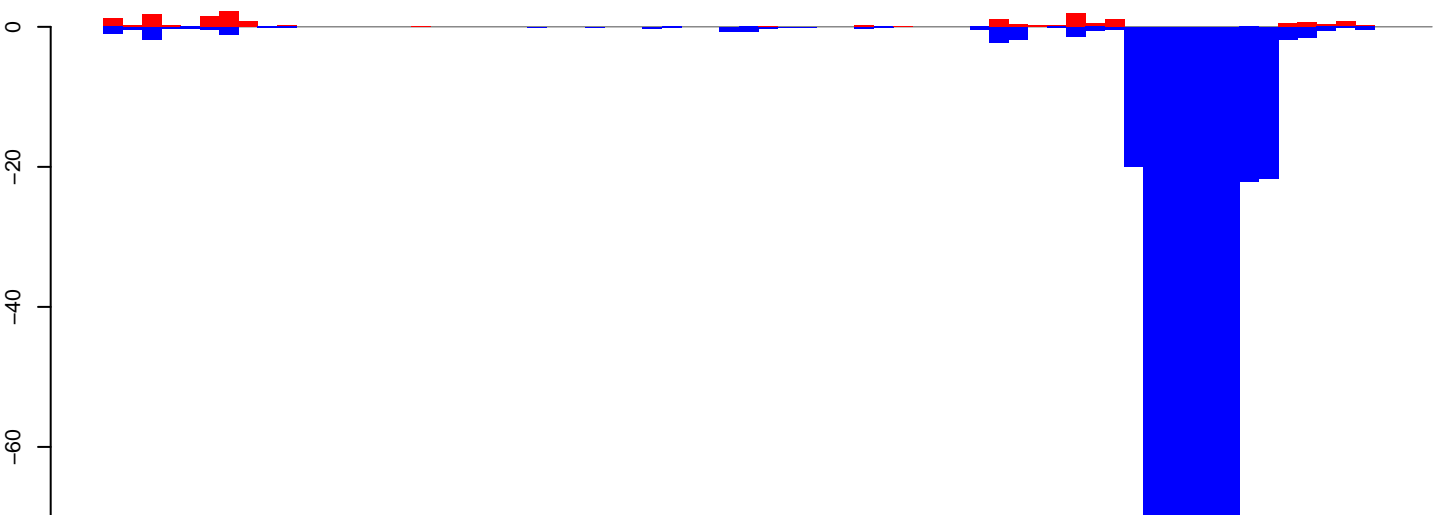
positive=13, negative=82, total=95

CuQuin_Testes_NL.24_35.rep



positive=3, negative=98, total=101

CuQuin_Testes_NL.rep

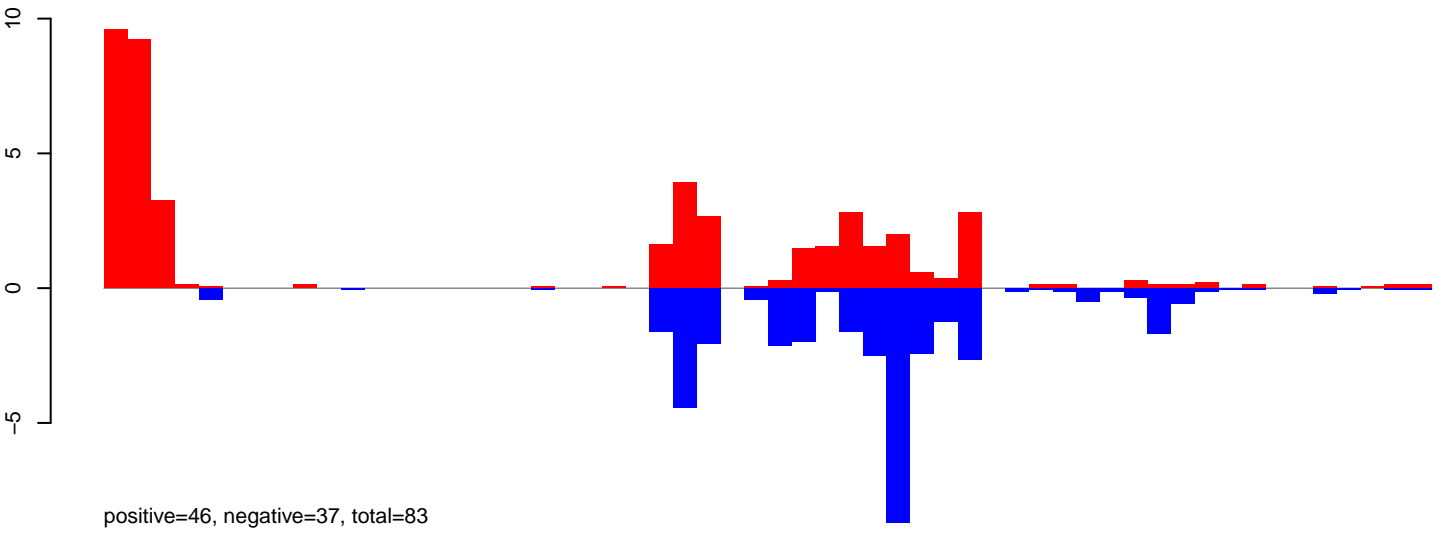


positive=16, negative=499, total=515

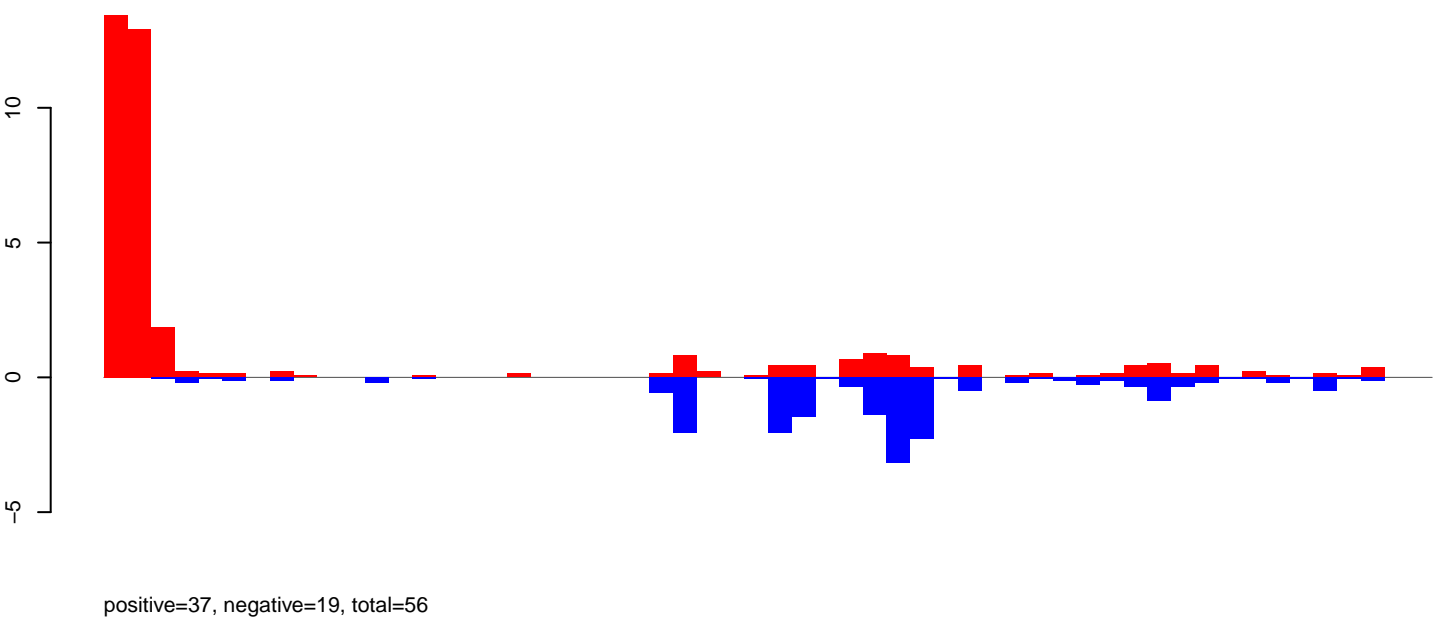
Window size=25, length=1718, TE@rnd-4_family-107@DNA@2292:1-1718

0 500 1000 1500

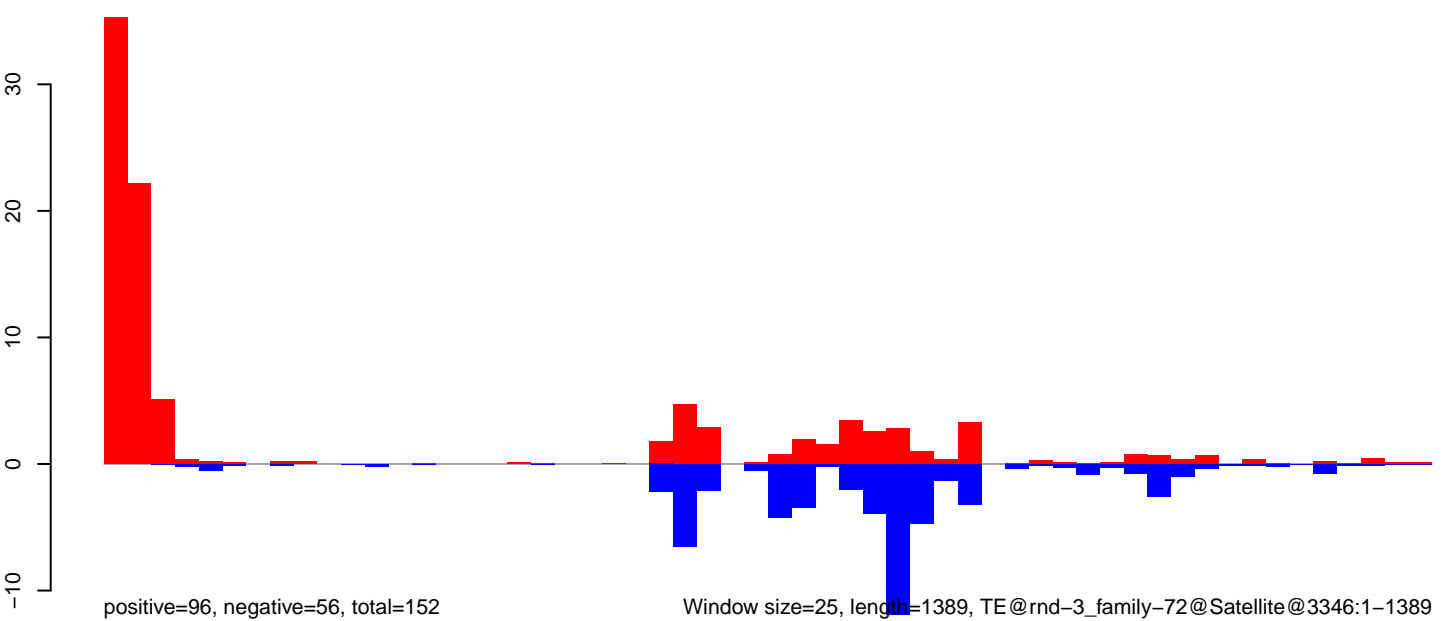
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

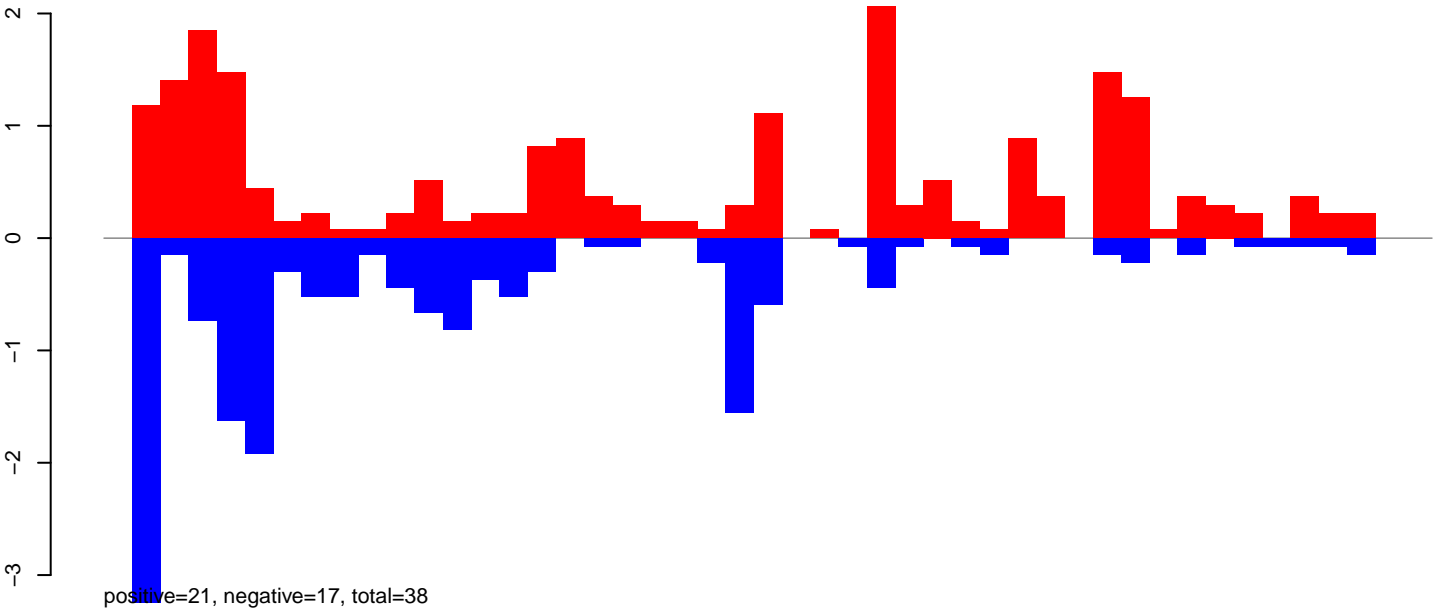


CuQuin_Testes_NL.rep

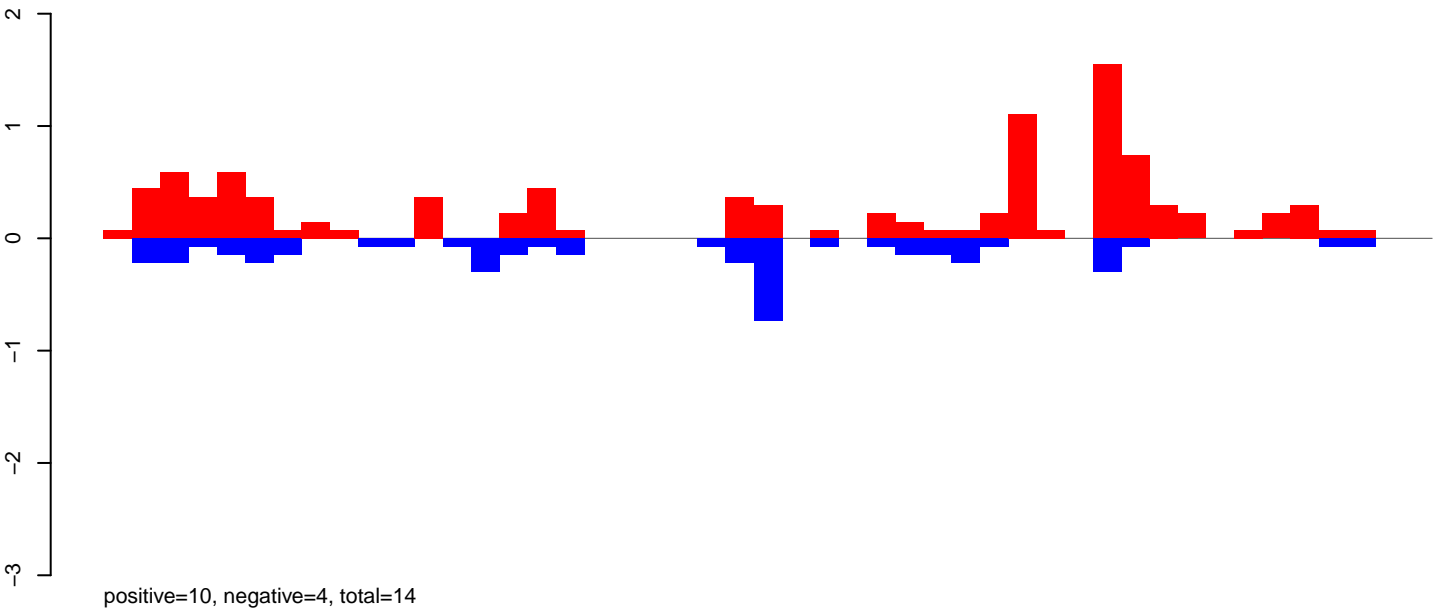


0 200 400 600 800 1000 1200 1400

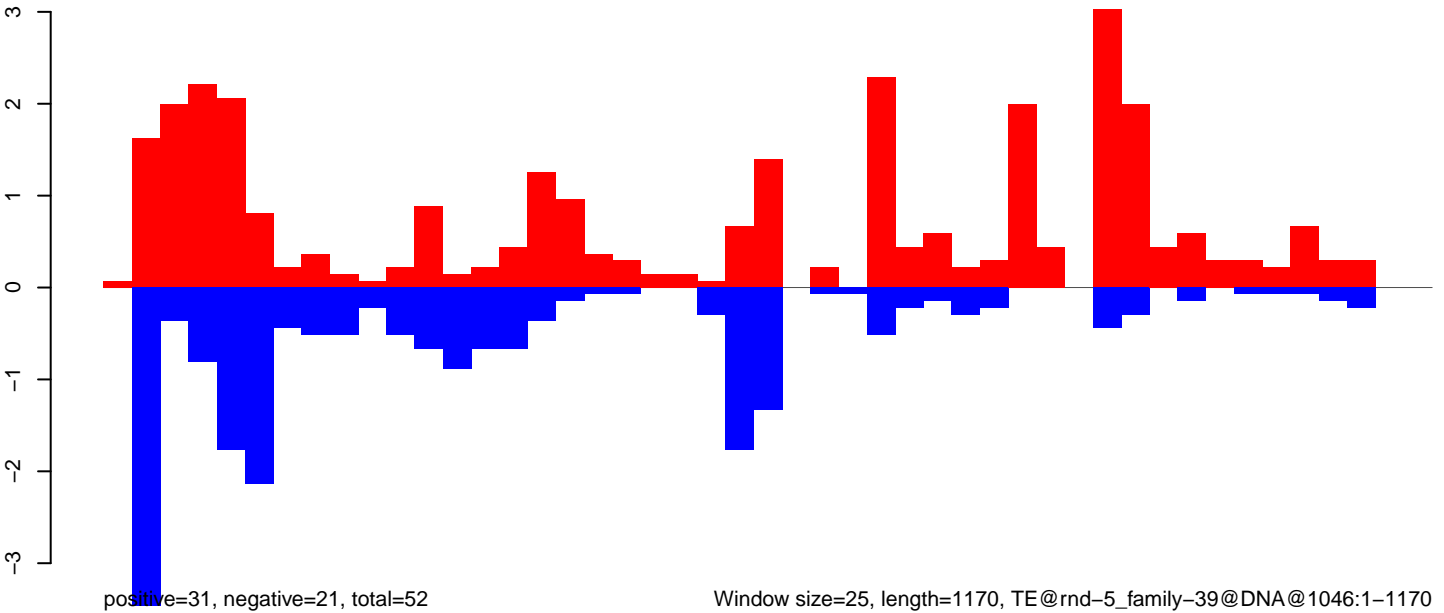
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



Window size=25, length=1170, TE@rnd-5_family-39@DNA@1046:1-1170

0 200 400 600 800 1000 1200

CuQuin_Testes_NL.18_23.rep



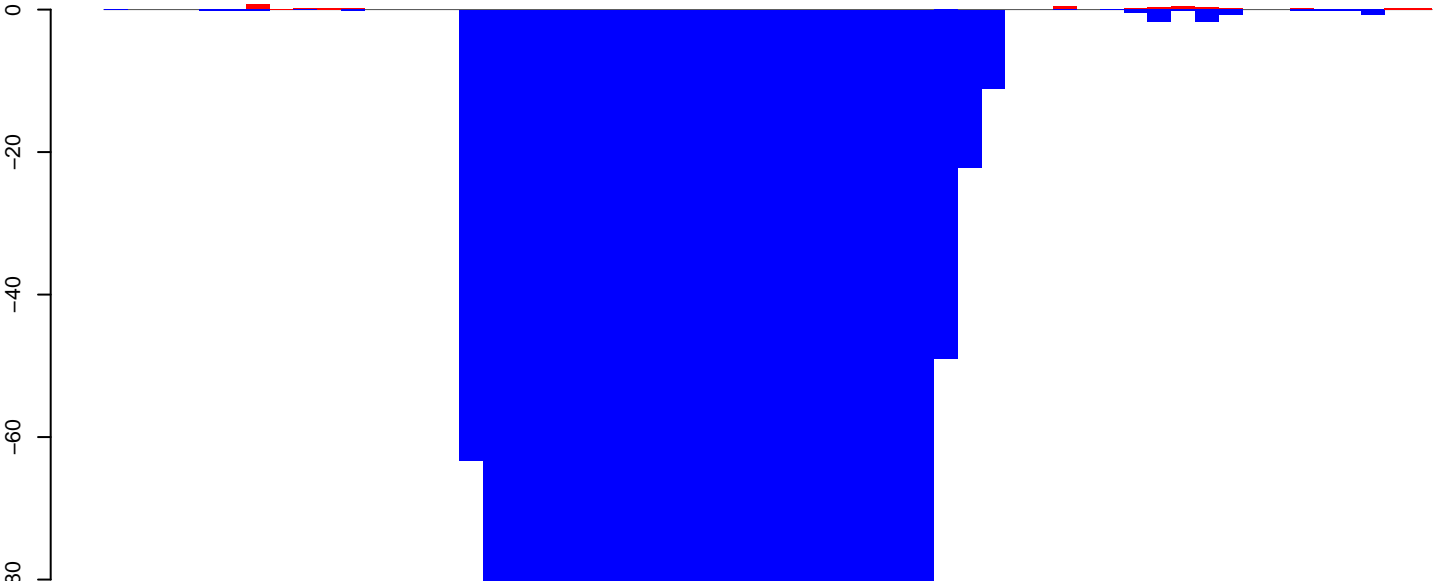
positive=2, negative=209, total=211

CuQuin_Testes_NL.24_35.rep



positive=2, negative=291, total=294

CuQuin_Testes_NL.rep

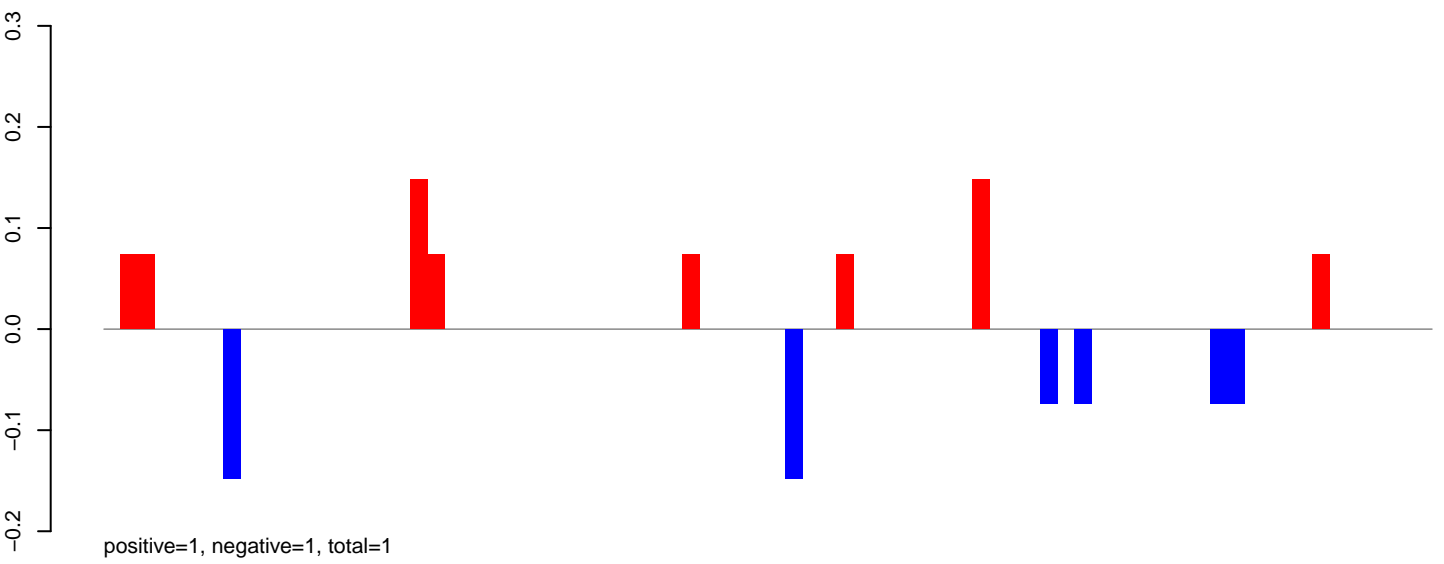


positive=4, negative=1731, total=1735

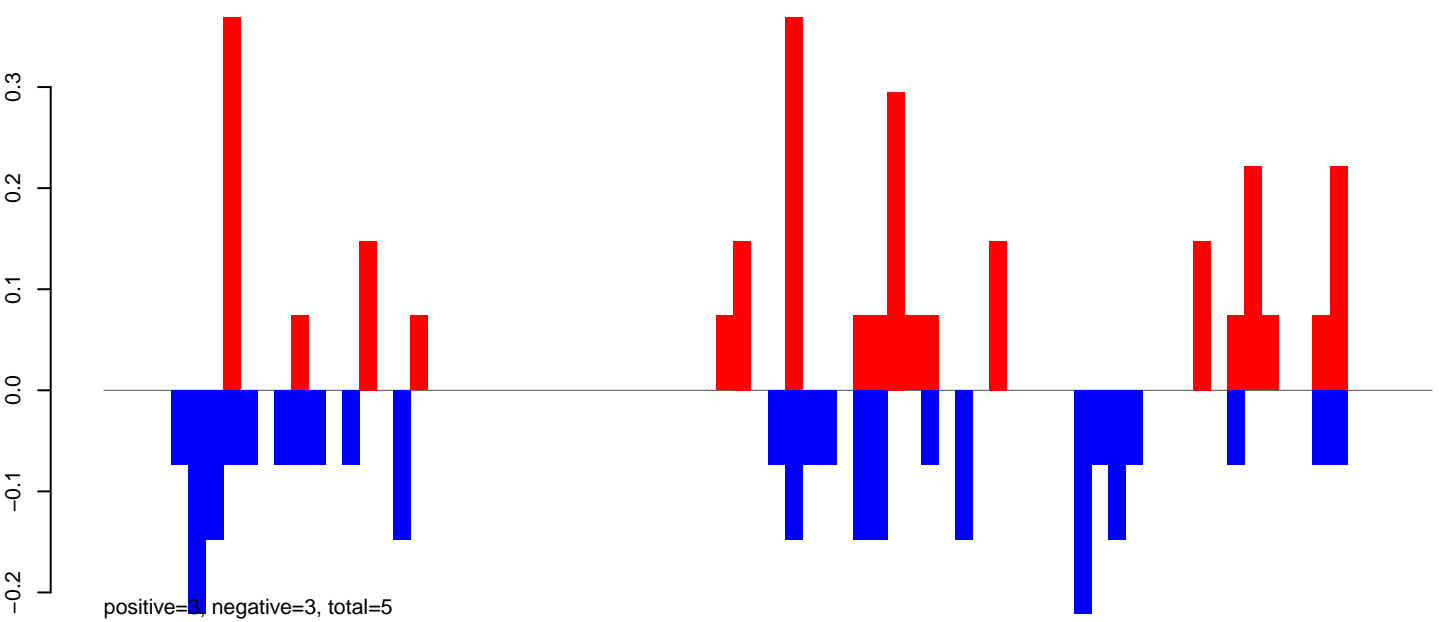
Window size=25, length=1389, TE@rnd-5_family-306@Unknown@3040:1-1389

0 200 400 600 800 1000 1200 1400

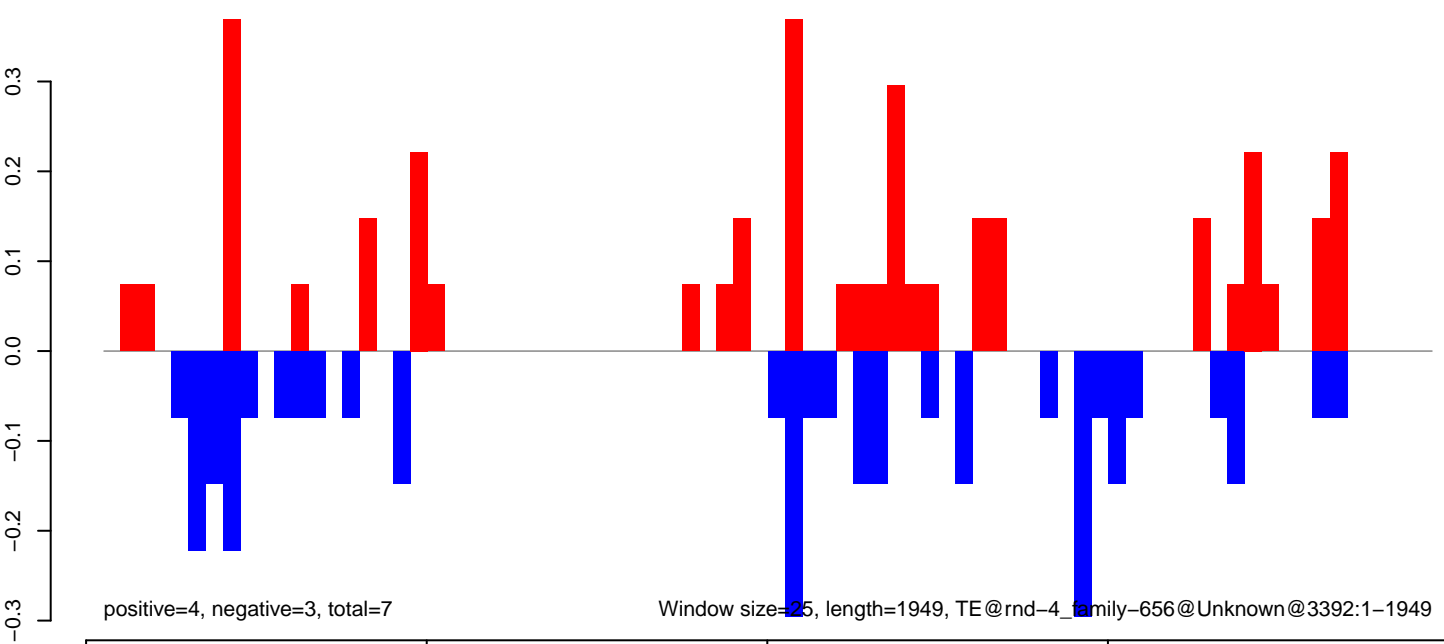
CuQuin_Testes_NL.18_23.rep



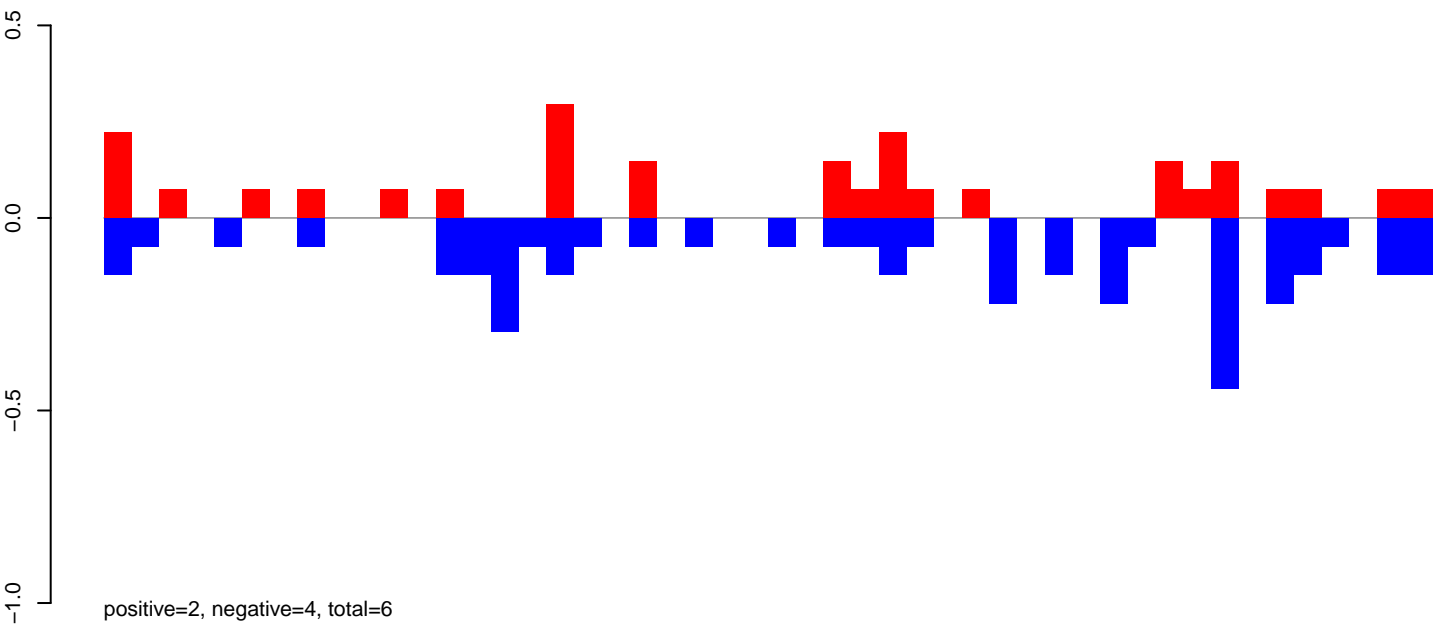
CuQuin_Testes_NL.24_35.rep



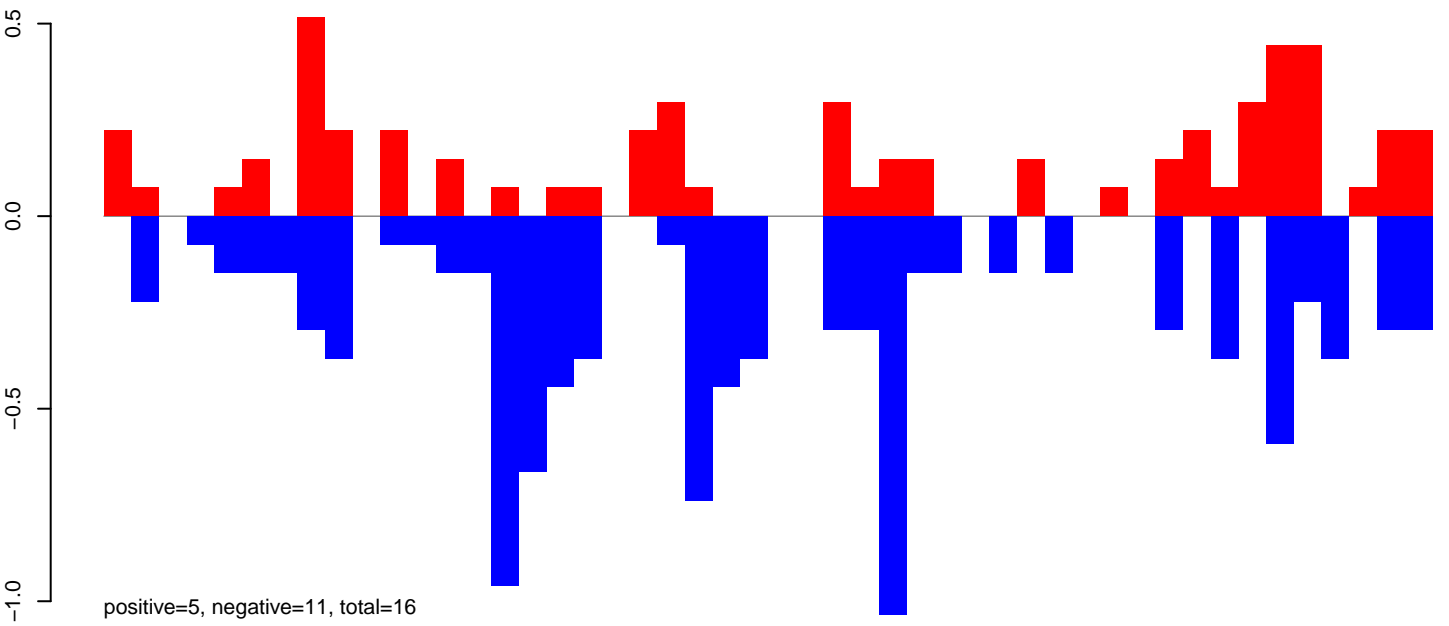
CuQuin_Testes_NL.rep



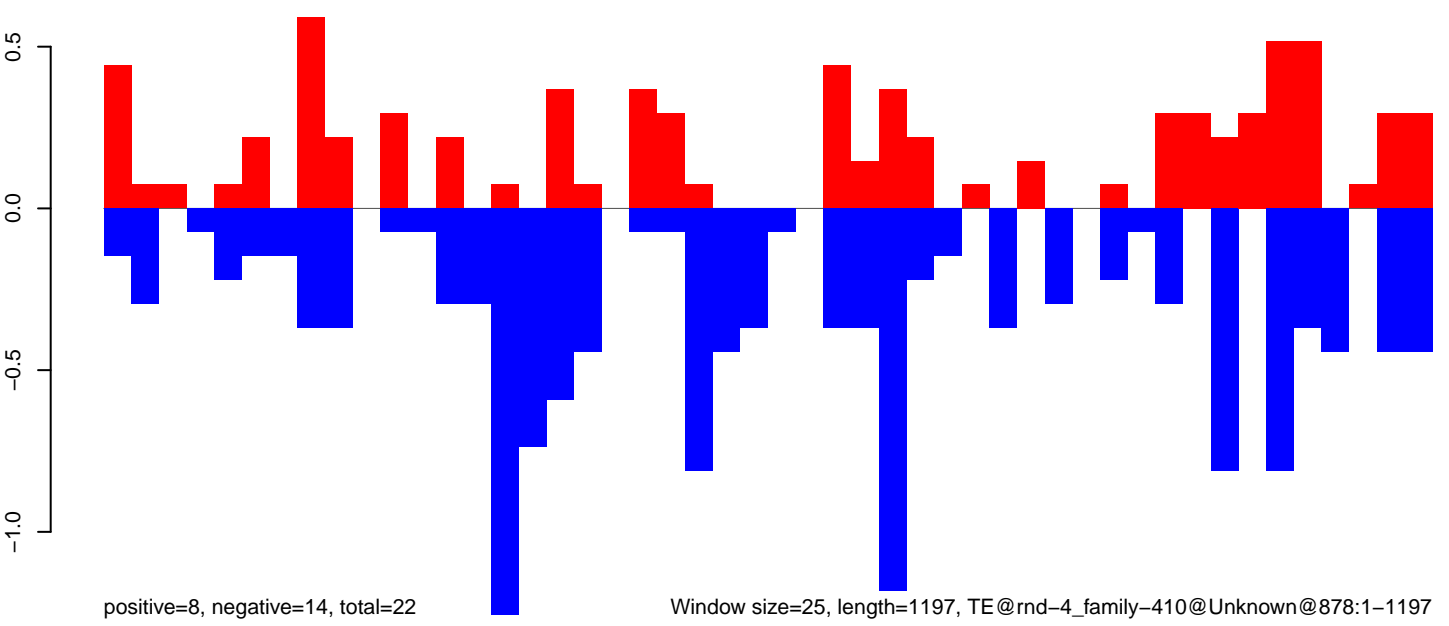
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



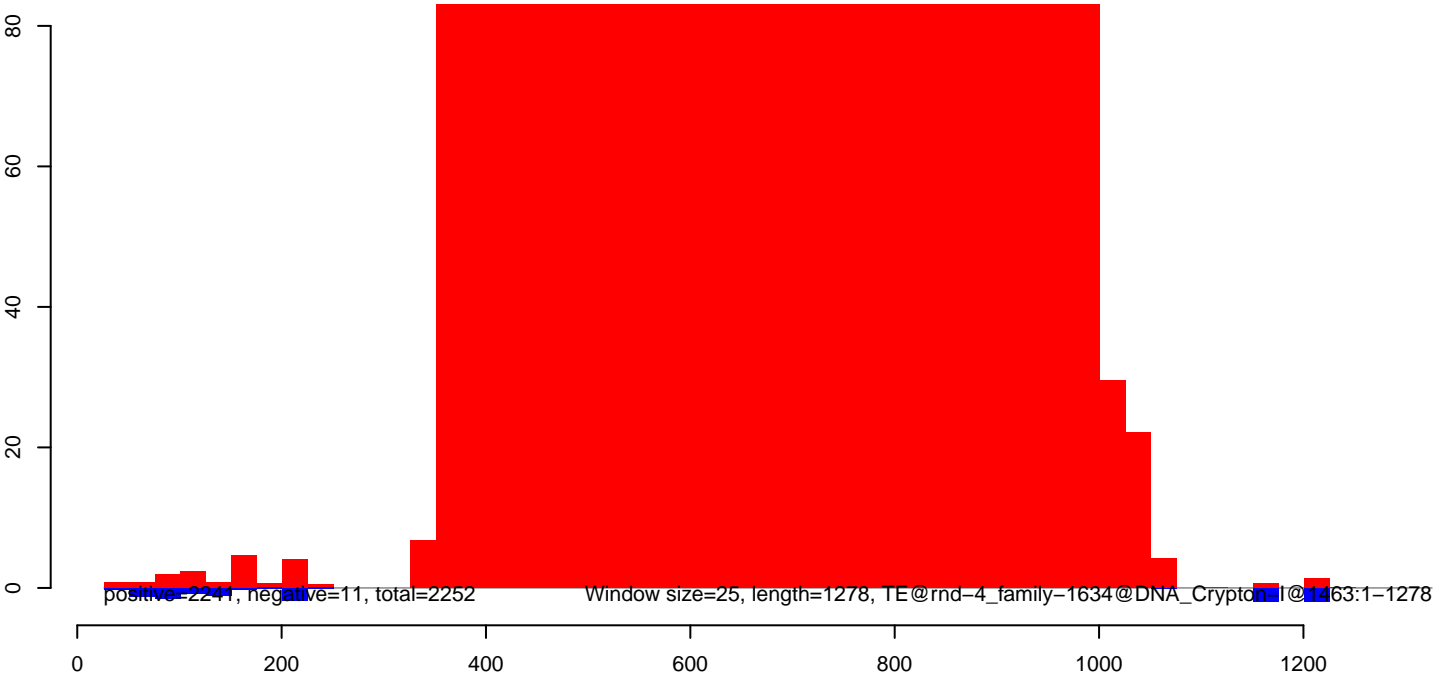
CuQuin_Testes_NL.18_23.rep



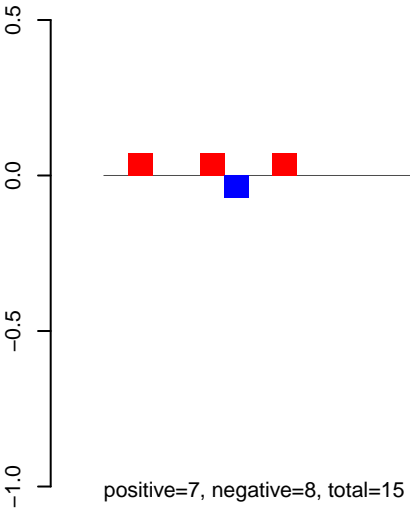
CuQuin_Testes_NL.24_35.rep



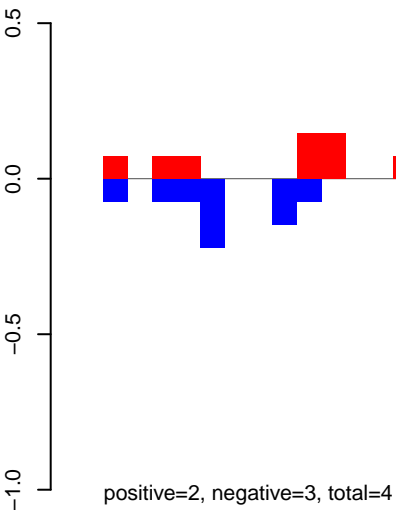
CuQuin_Testes_NL.rep



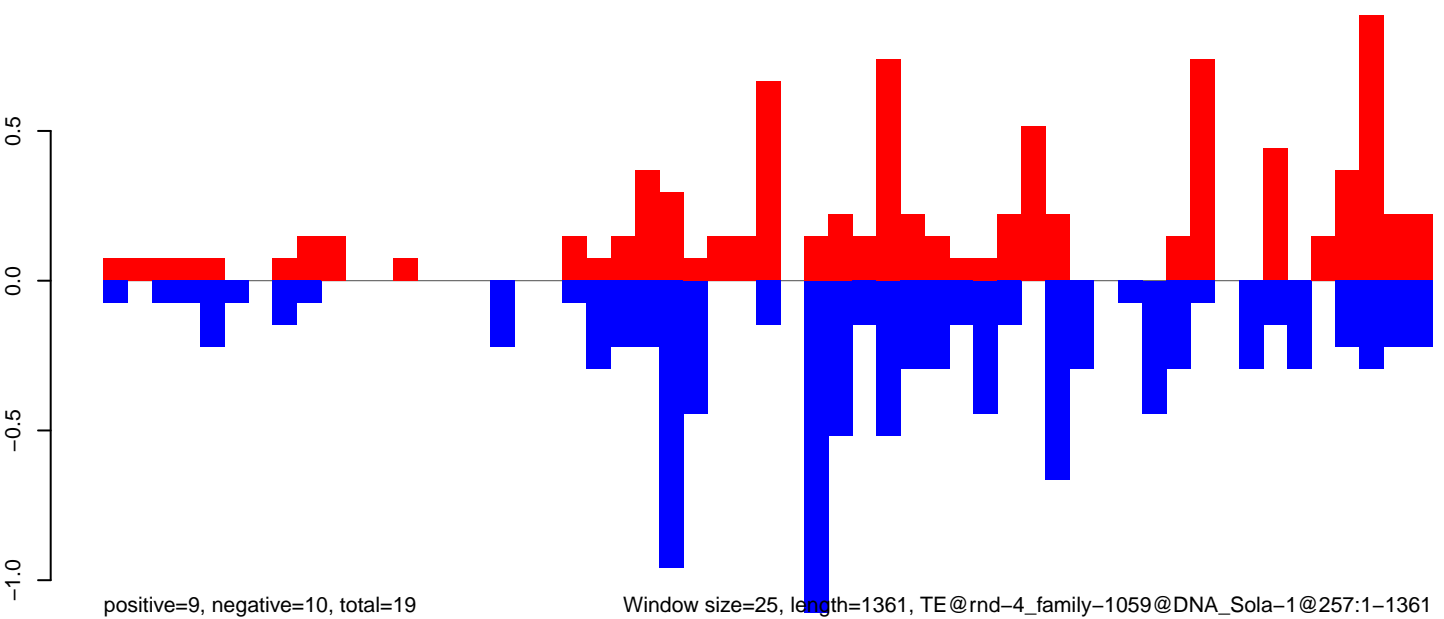
CuQuin_Testes_NL.18_23.rep



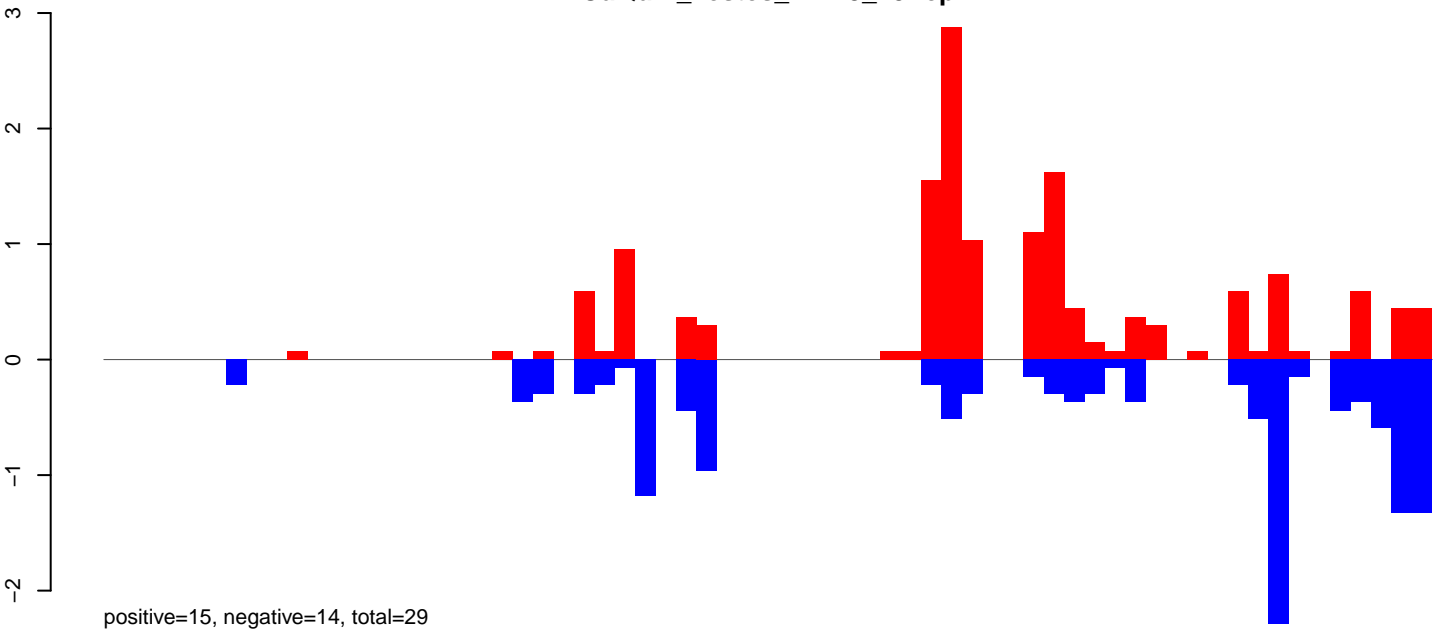
CuQuin_Testes_NL.24_35.rep



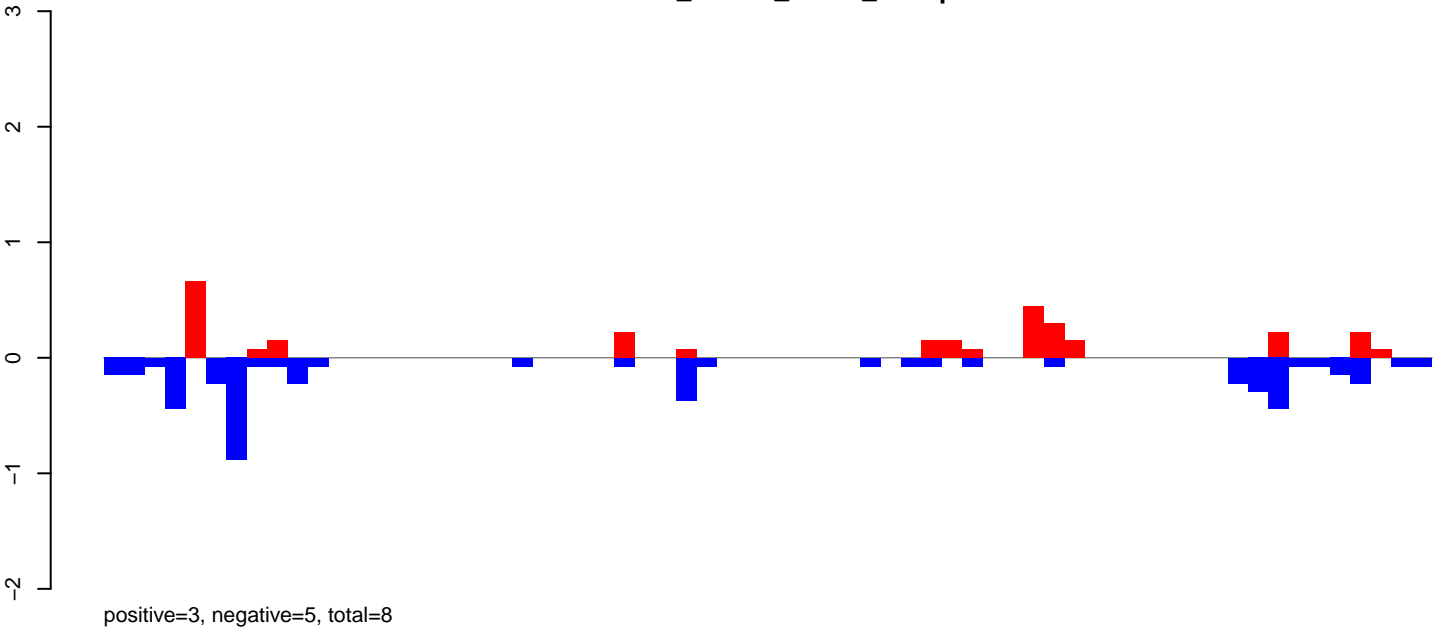
CuQuin_Testes_NL.rep



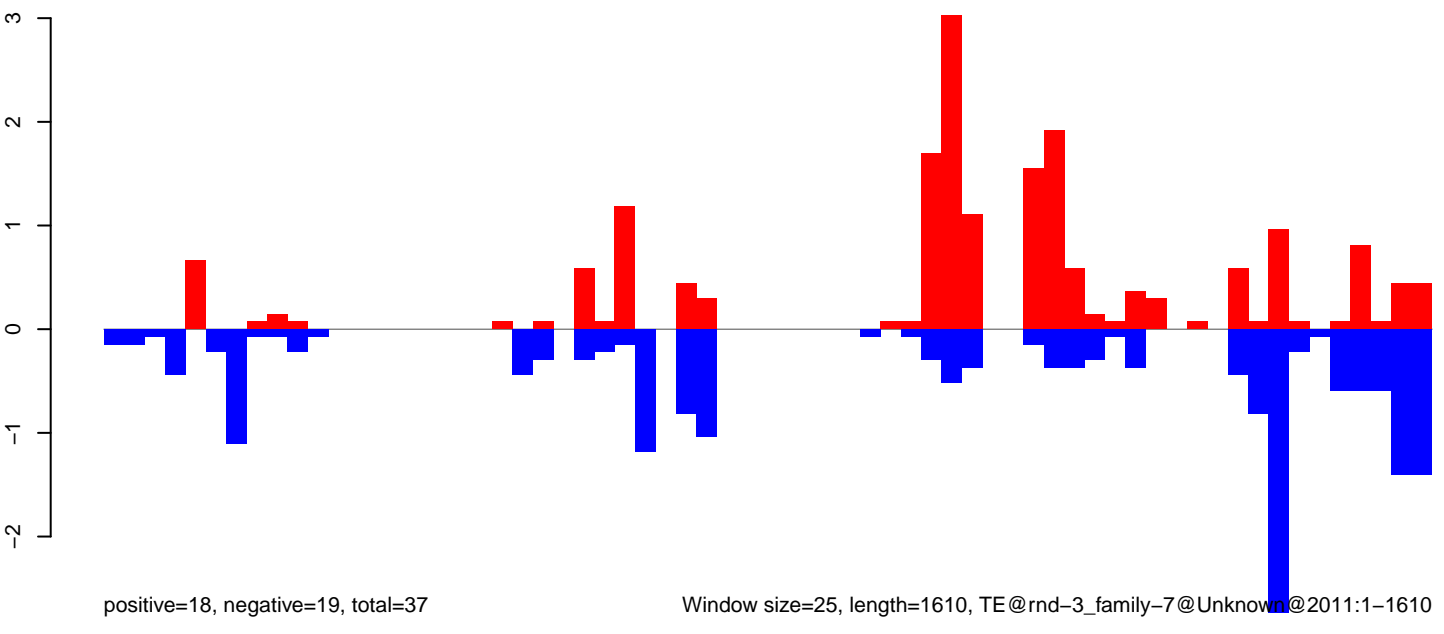
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



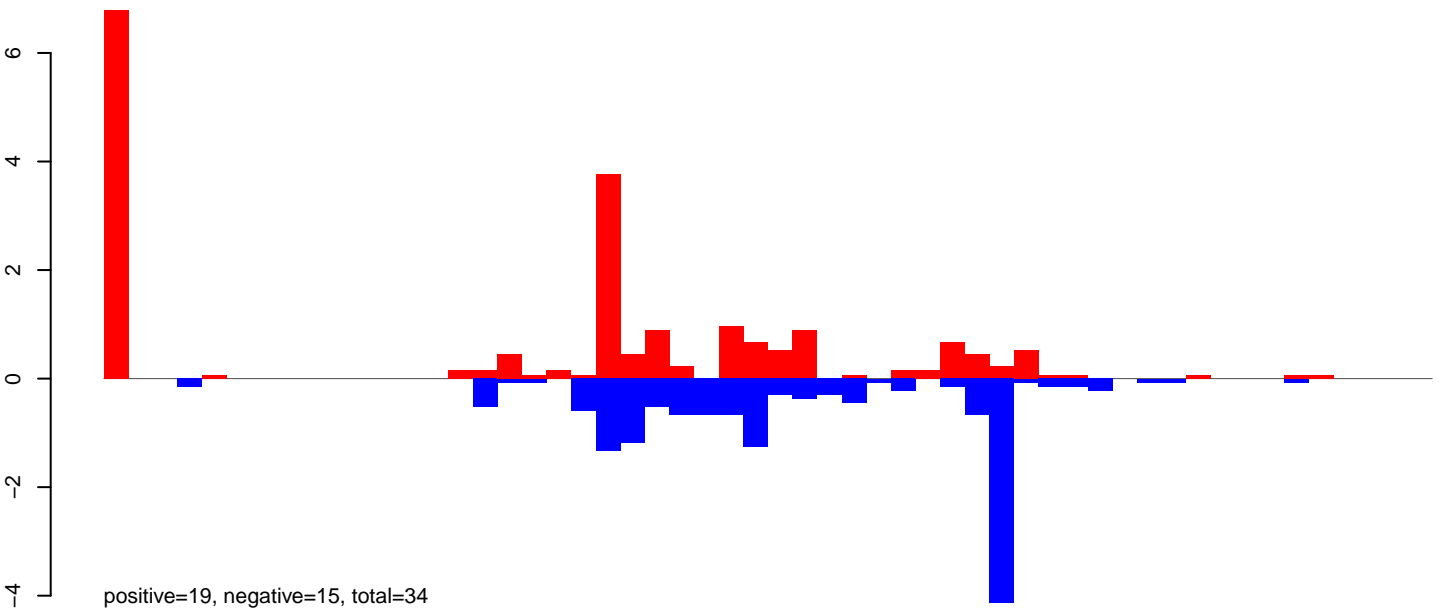
CuQuin_Testes_NL.rep



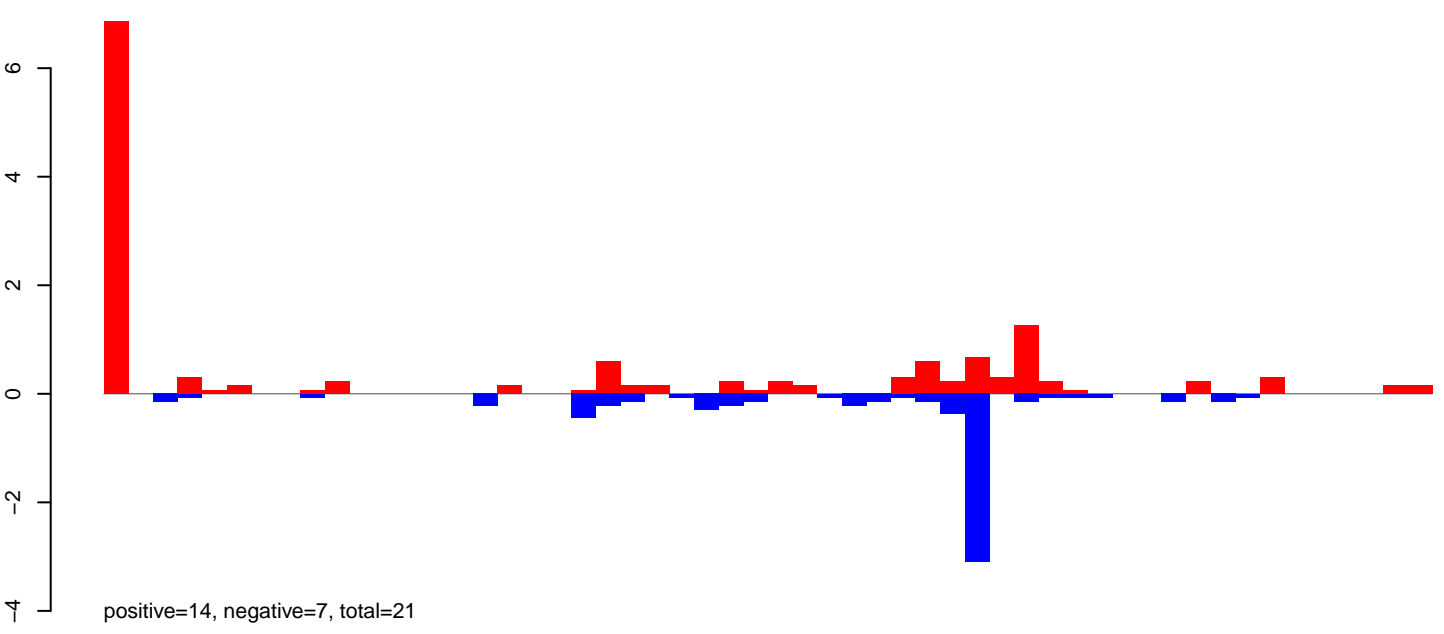
Window size=25, length=1610, TE@rnd-3_family-7@Unknown@2011:1-1610

0 500 1000 1500

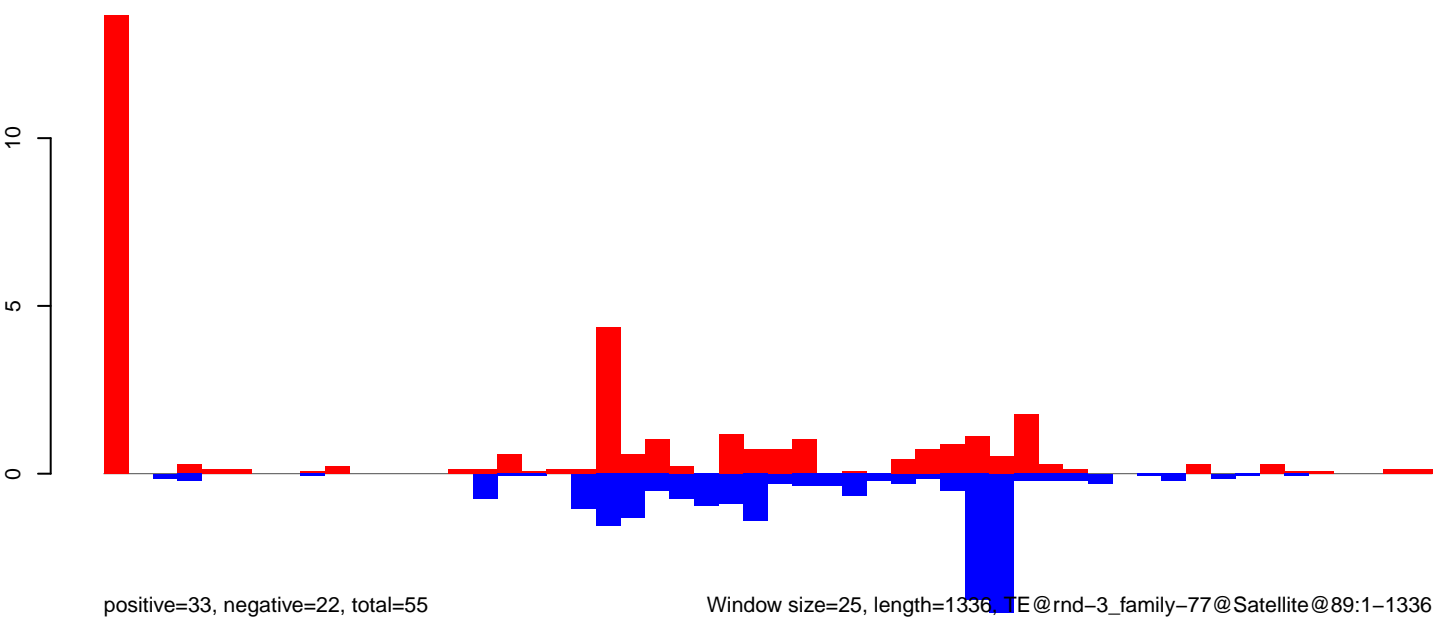
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



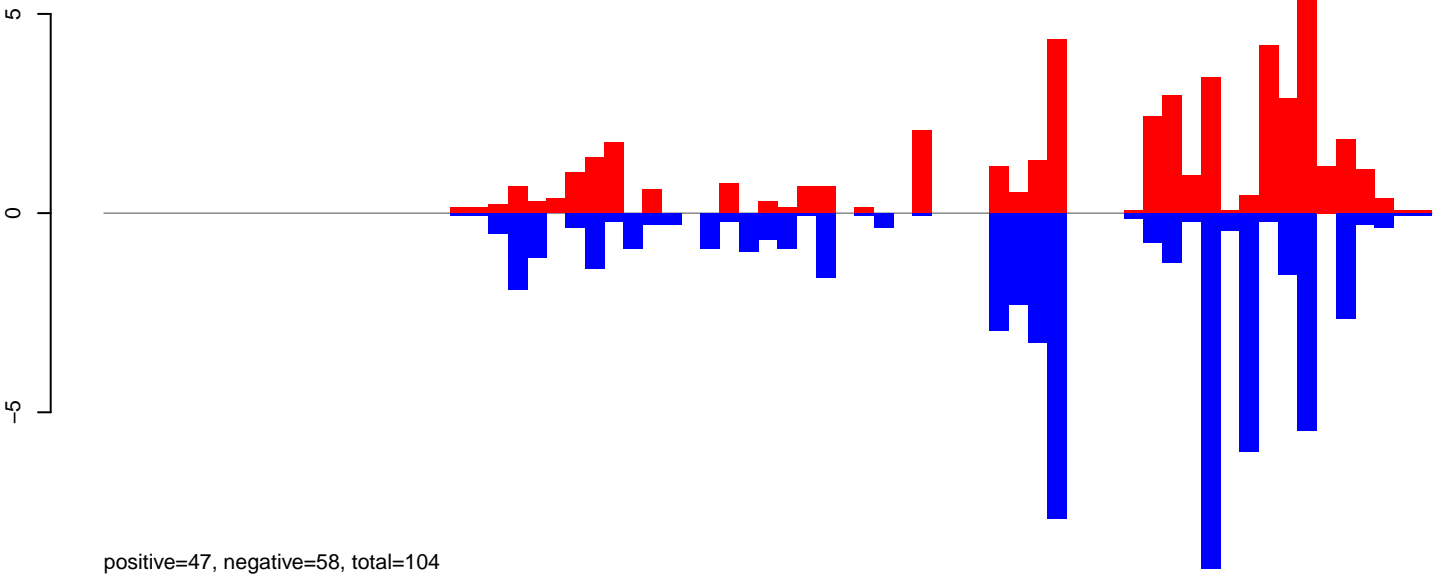
CuQuin_Testes_NL.rep



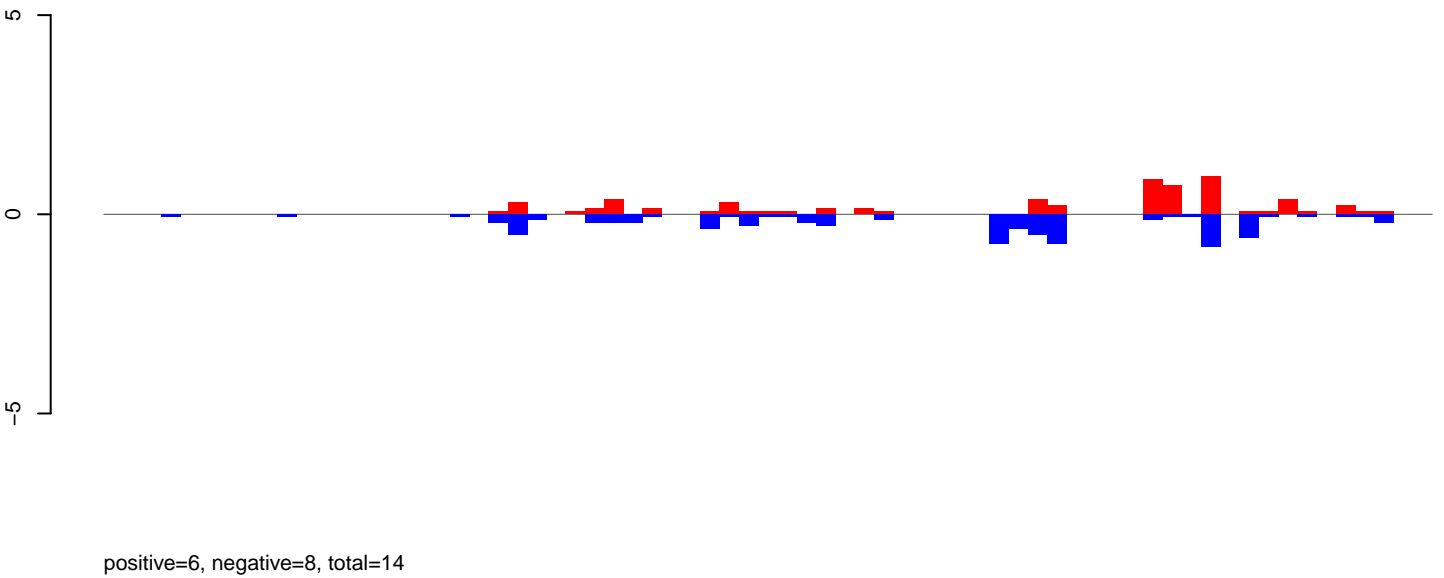
Window size=25, length=1336, TE@rnd-3_family-77@Satellite@89:1-1336

0 200 400 600 800 1000 1200 1400

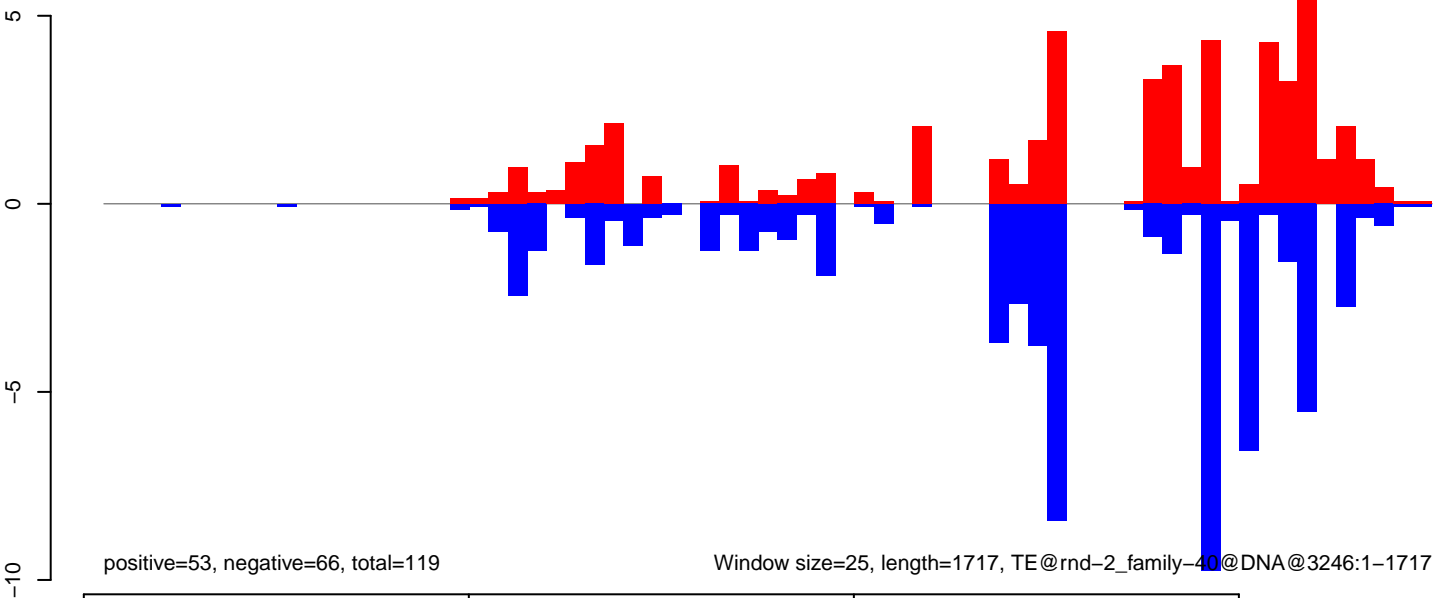
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

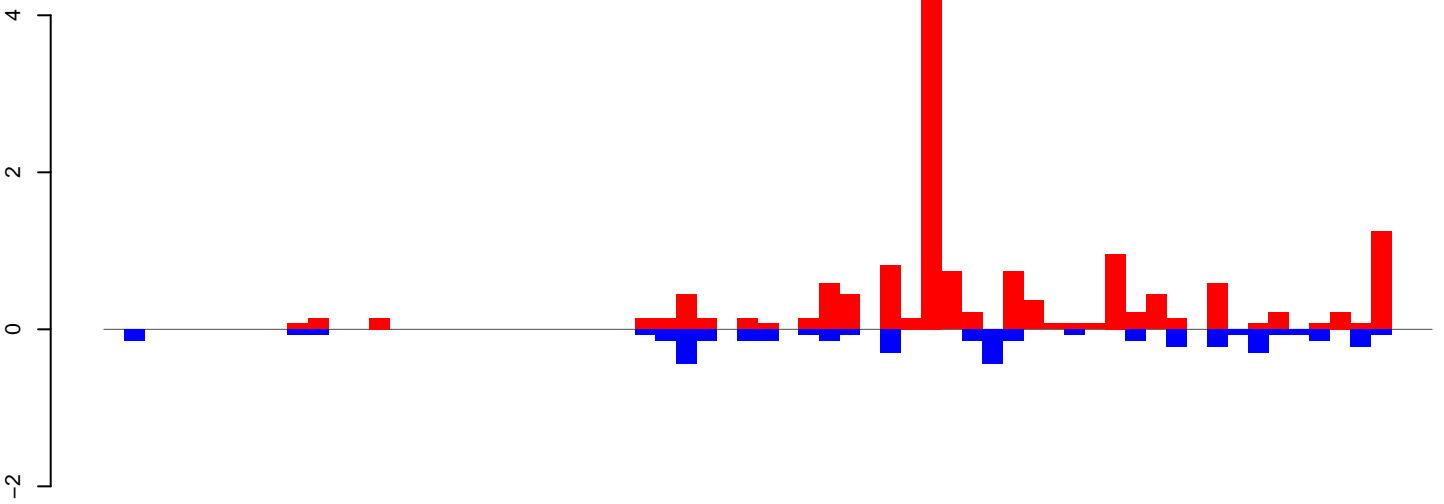


CuQuin_Testes_NL.rep



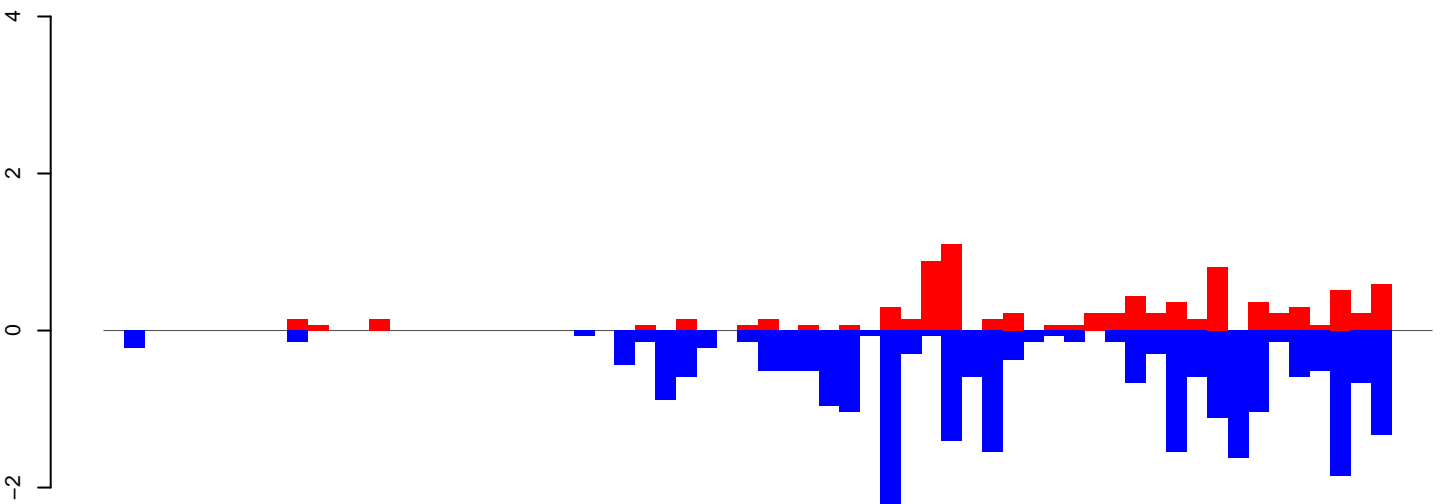
Window size=25, length=1717, TE@rnd-2_family-40@DNA@3246:1-1717

CuQuin_Testes_NL.18_23.rep



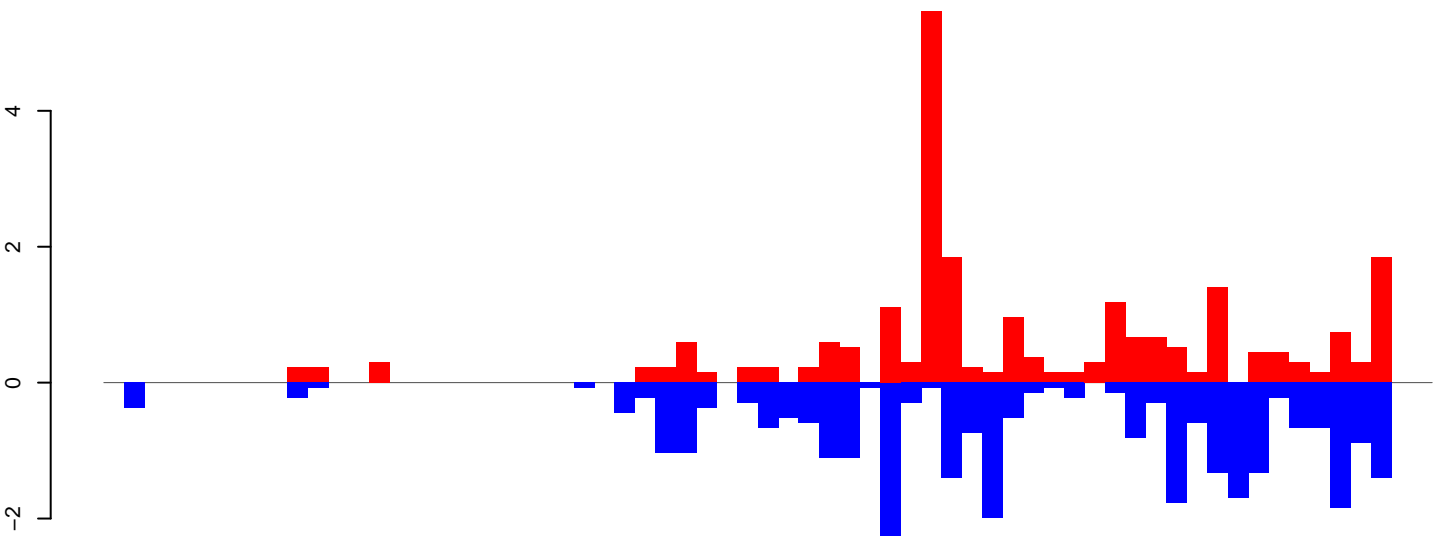
positive=15, negative=4, total=19

CuQuin_Testes_NL.24_35.rep



positive=9, negative=26, total=35

CuQuin_Testes_NL.rep

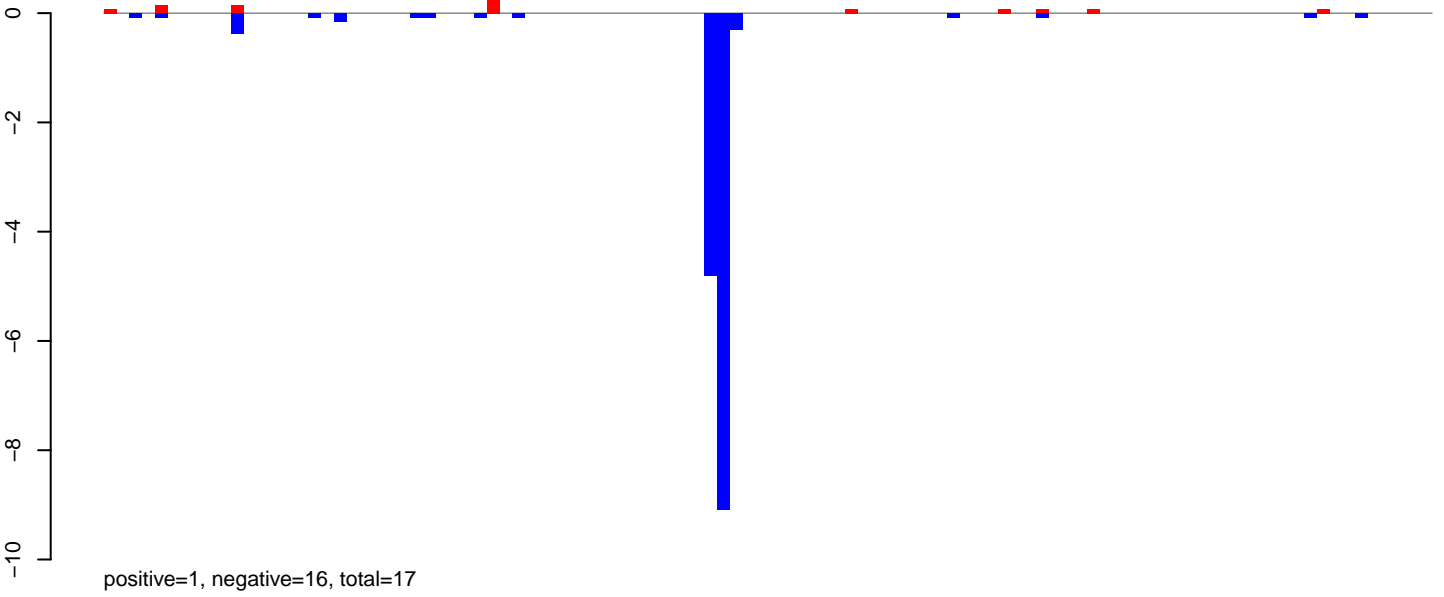


positive=23, negative=31, total=54

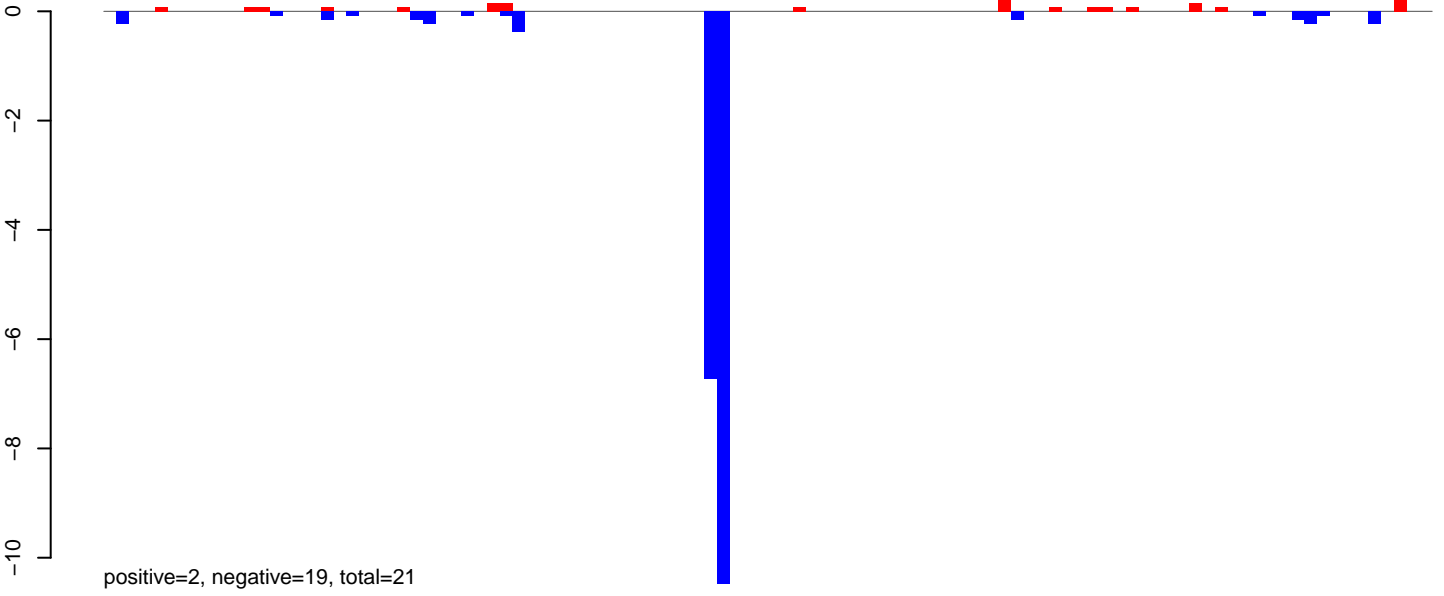
Window size=25, length=1602 TE@rnd-5_family-488@LINE RTE-BovB@409:1-1602

0 500 1000 1500

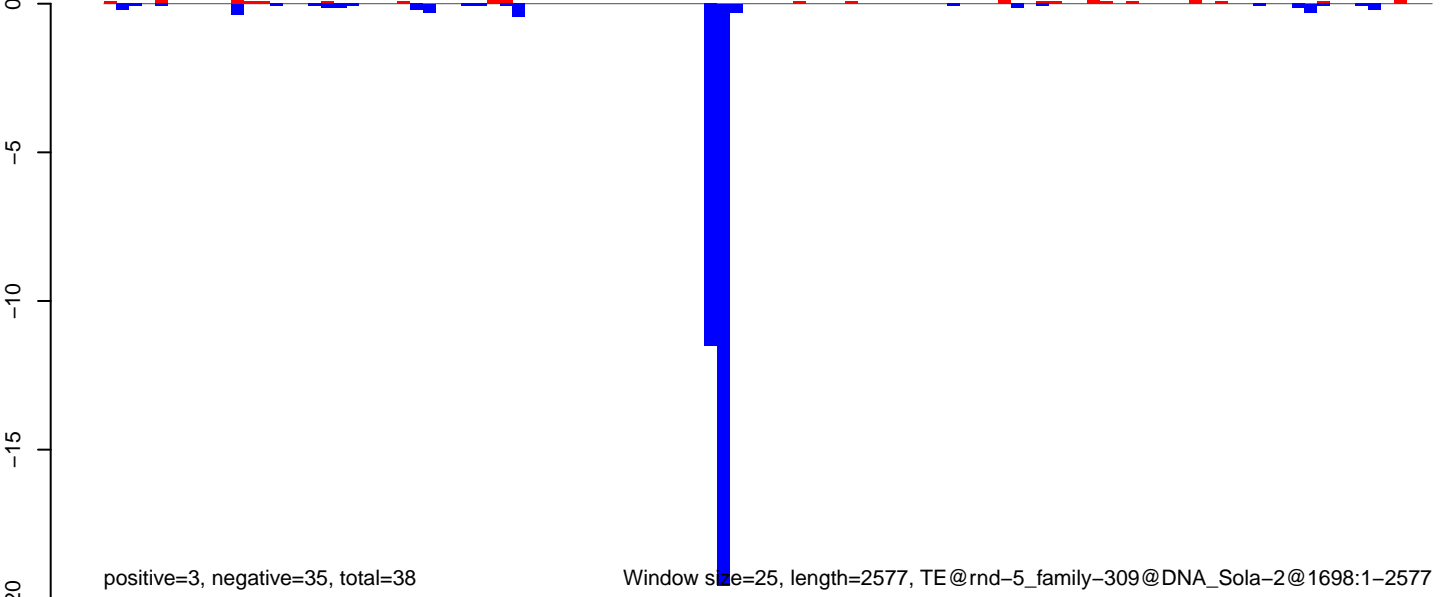
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

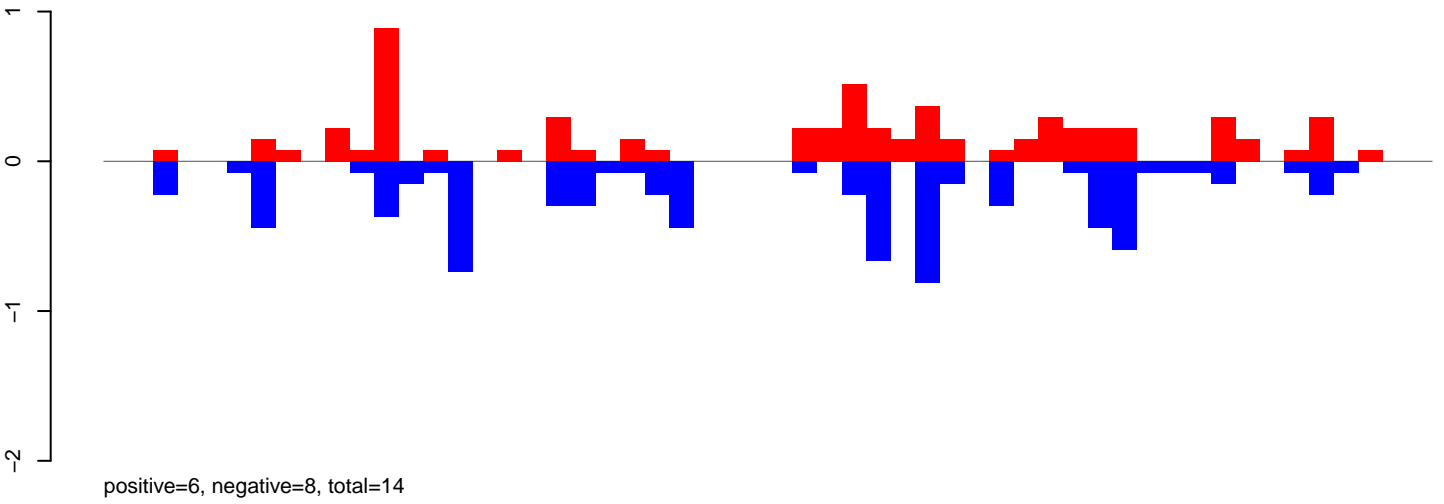


CuQuin_Testes_NL.rep

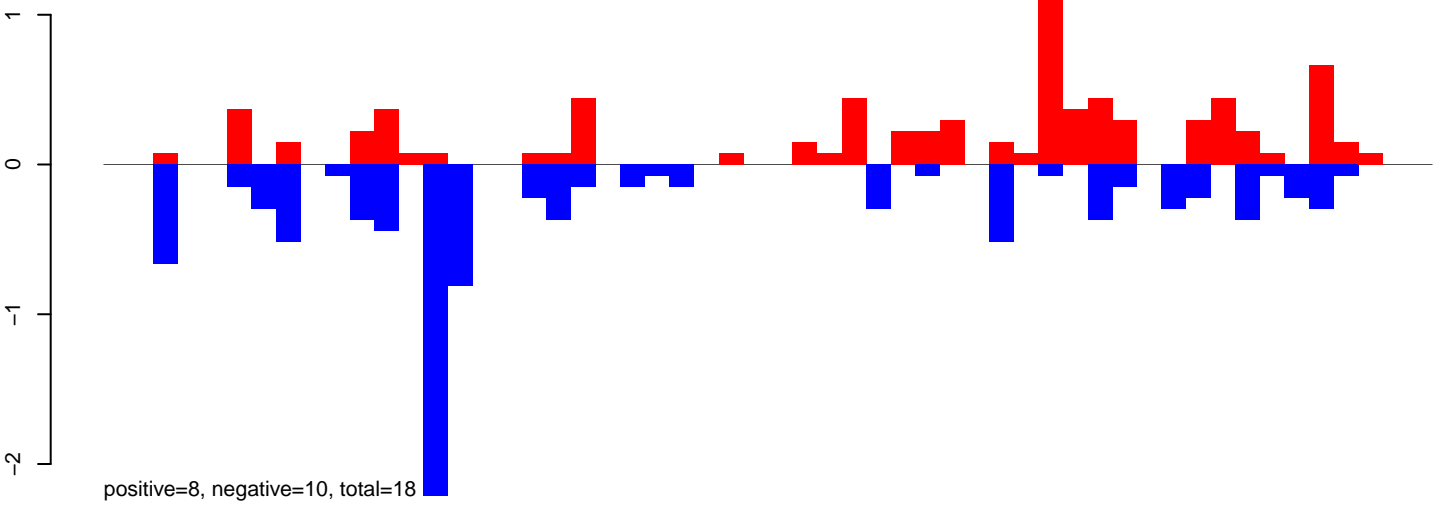


Window size=25, length=2577, TE@rnd-5_family-309@DNA_Sola-2@1698:1-2577

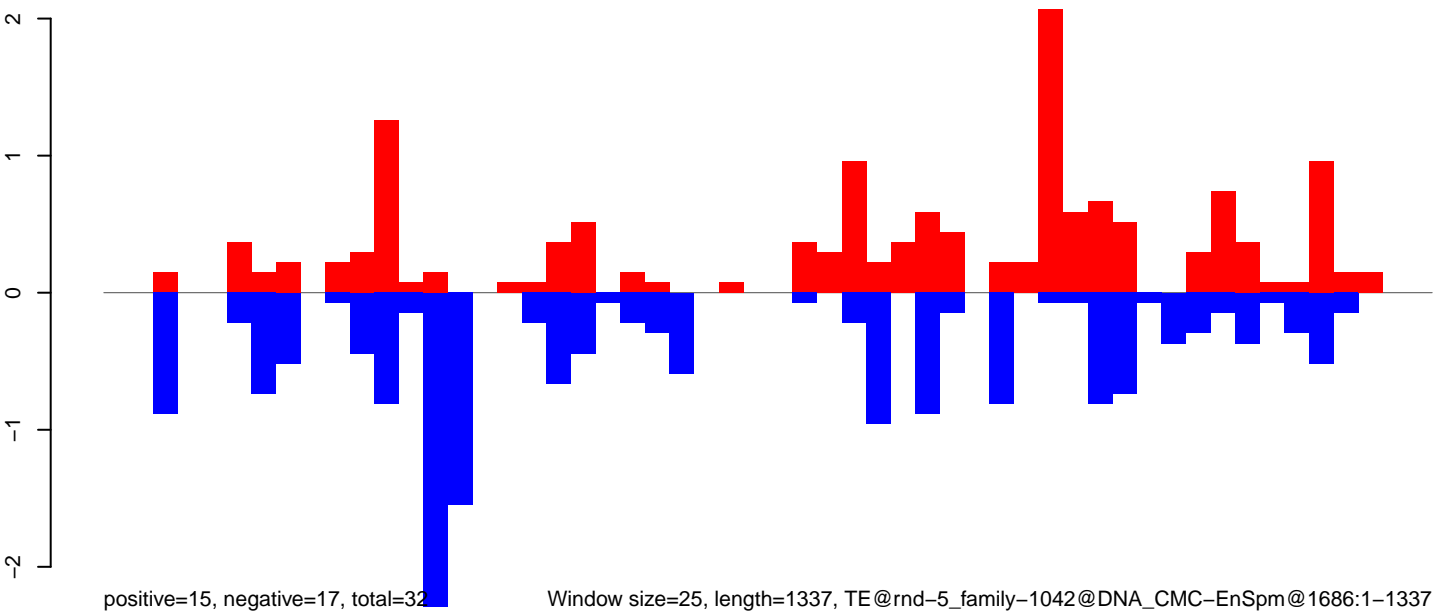
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



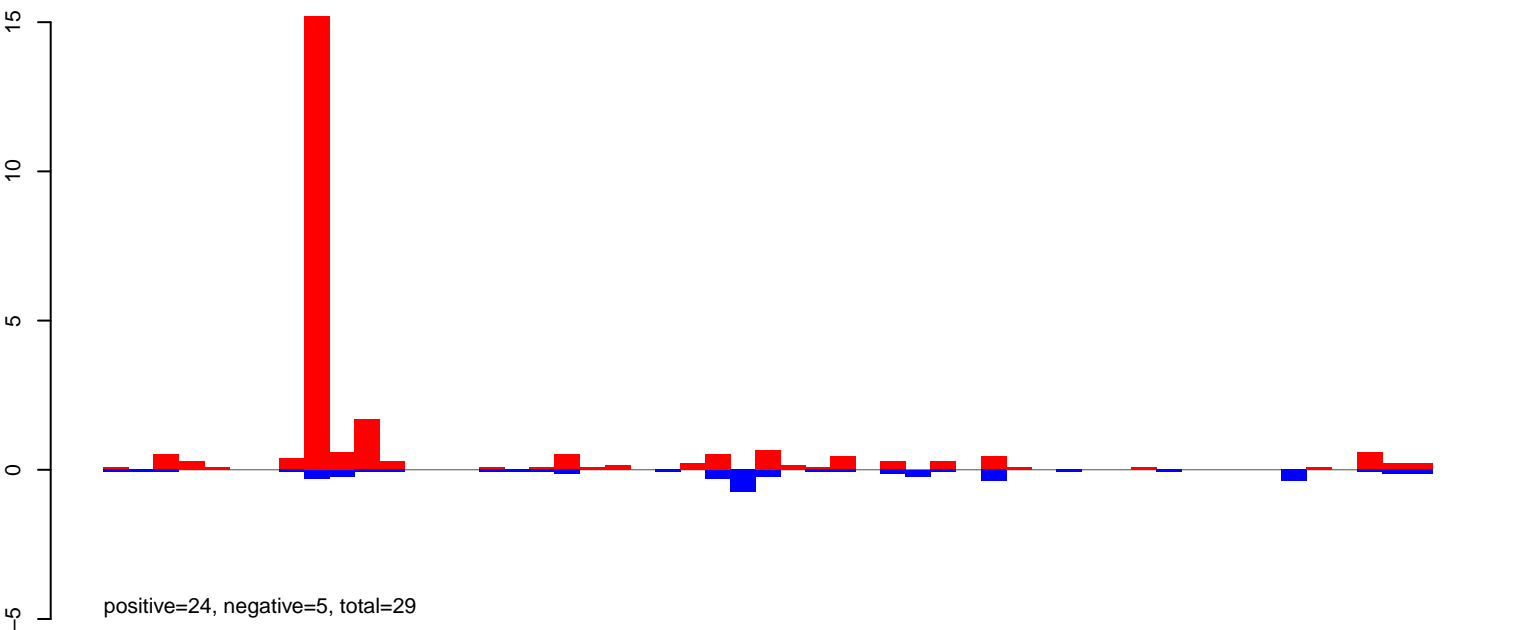
CuQuin_Testes_NL.rep



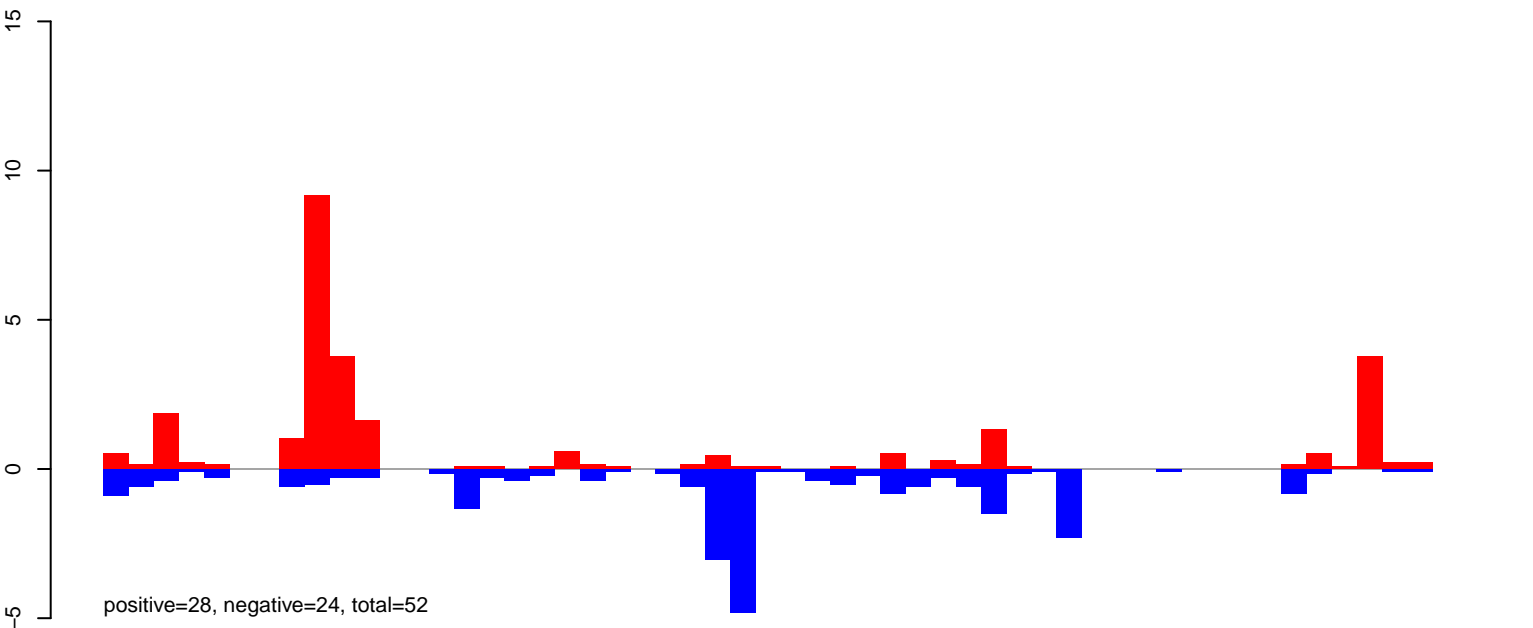
Window size=25, length=1337, TE@rnd-5_family-1042@DNA_CMC-EnSpm@1686:1-1337

0 200 400 600 800 1000 1200 1400

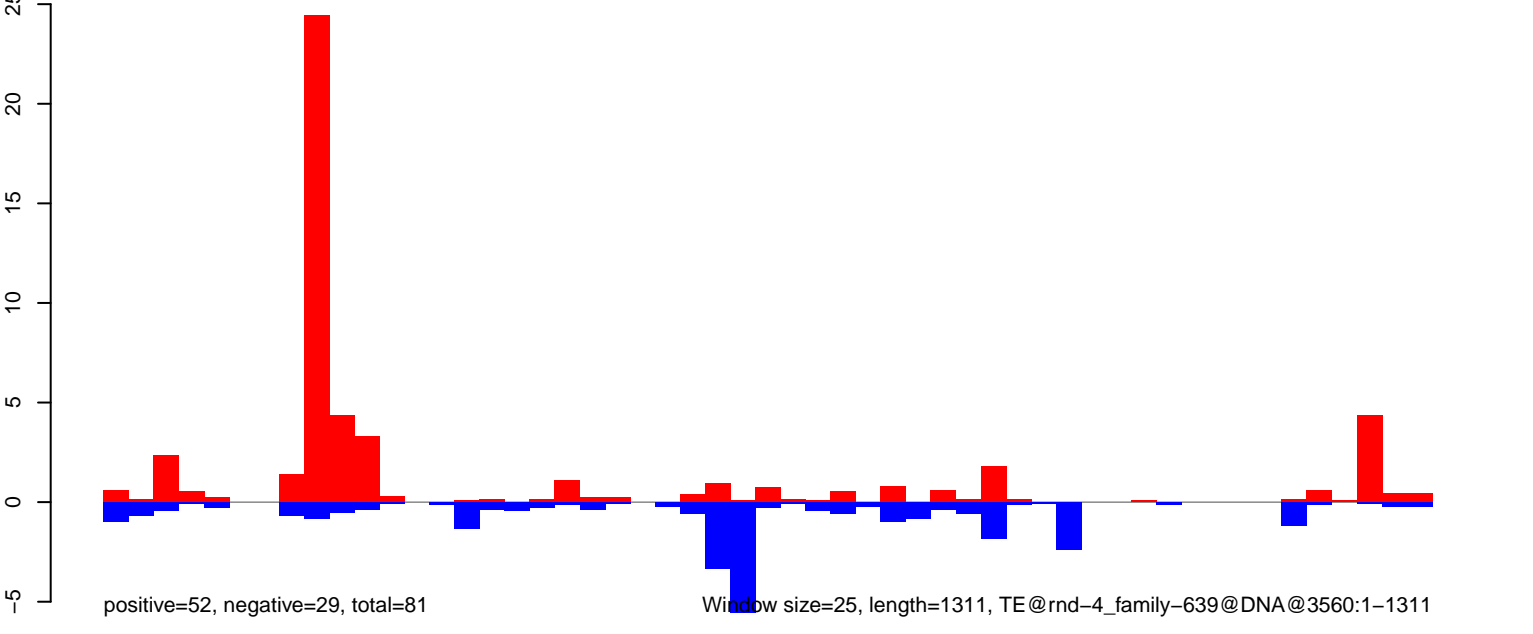
CuQuin_Testes_NL.18_23.rep



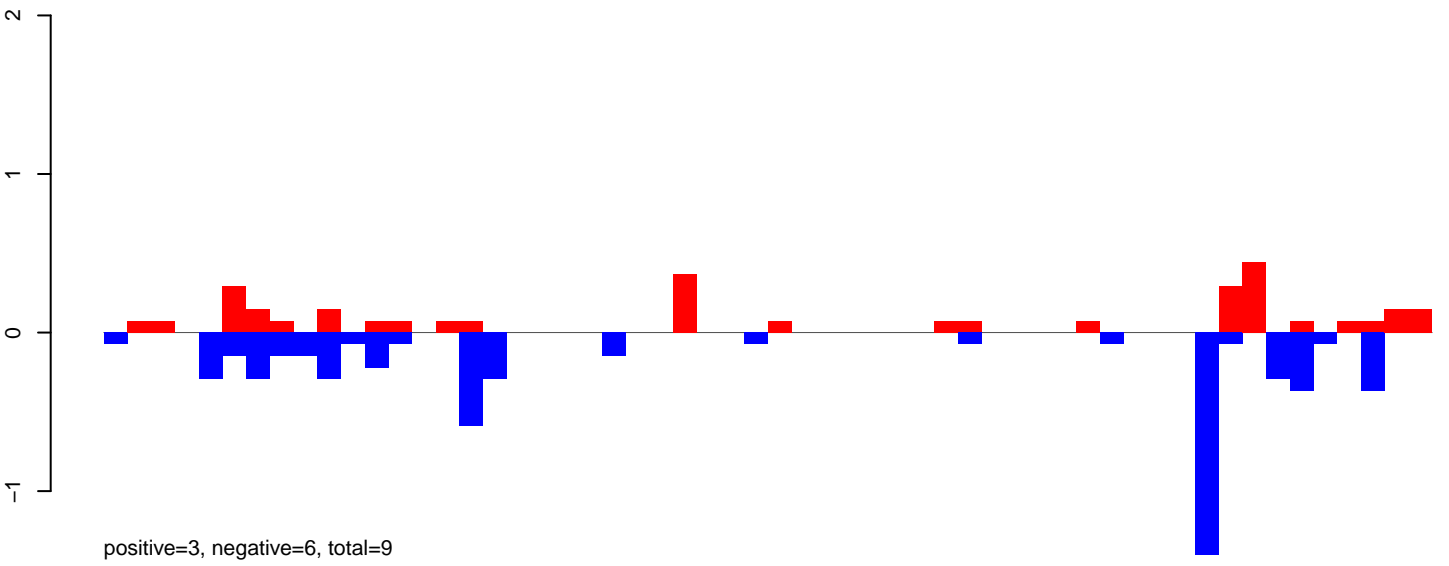
CuQuin_Testes_NL.24_35.rep



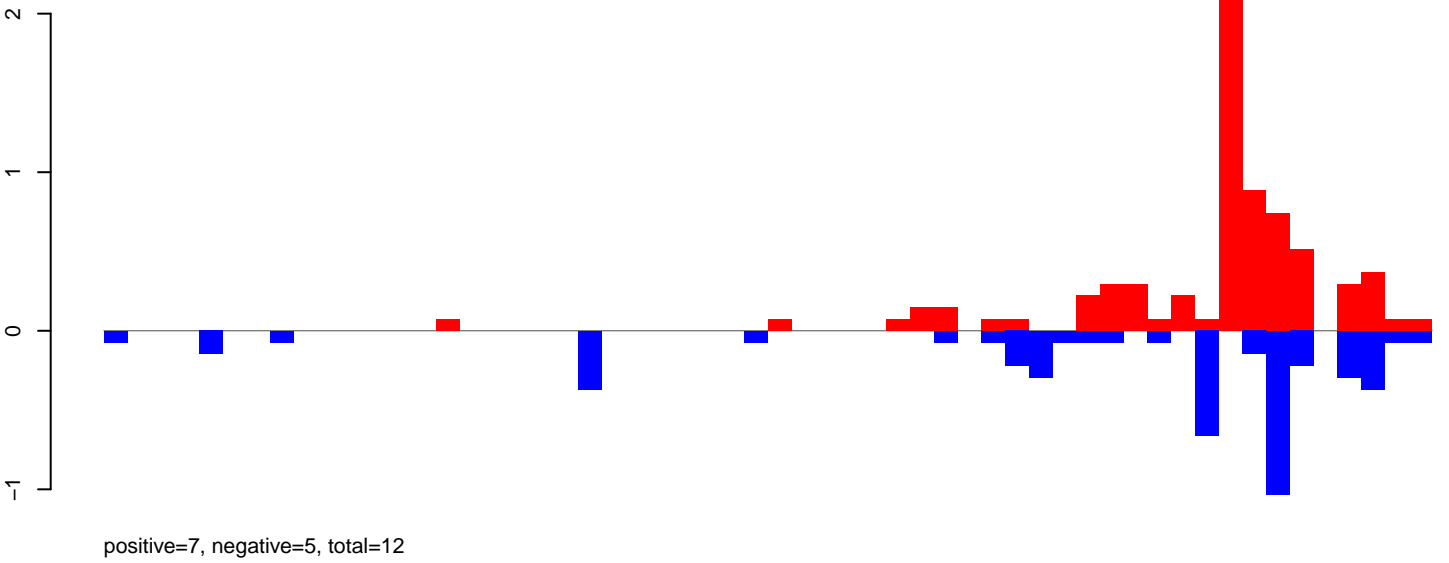
CuQuin_Testes_NL.rep



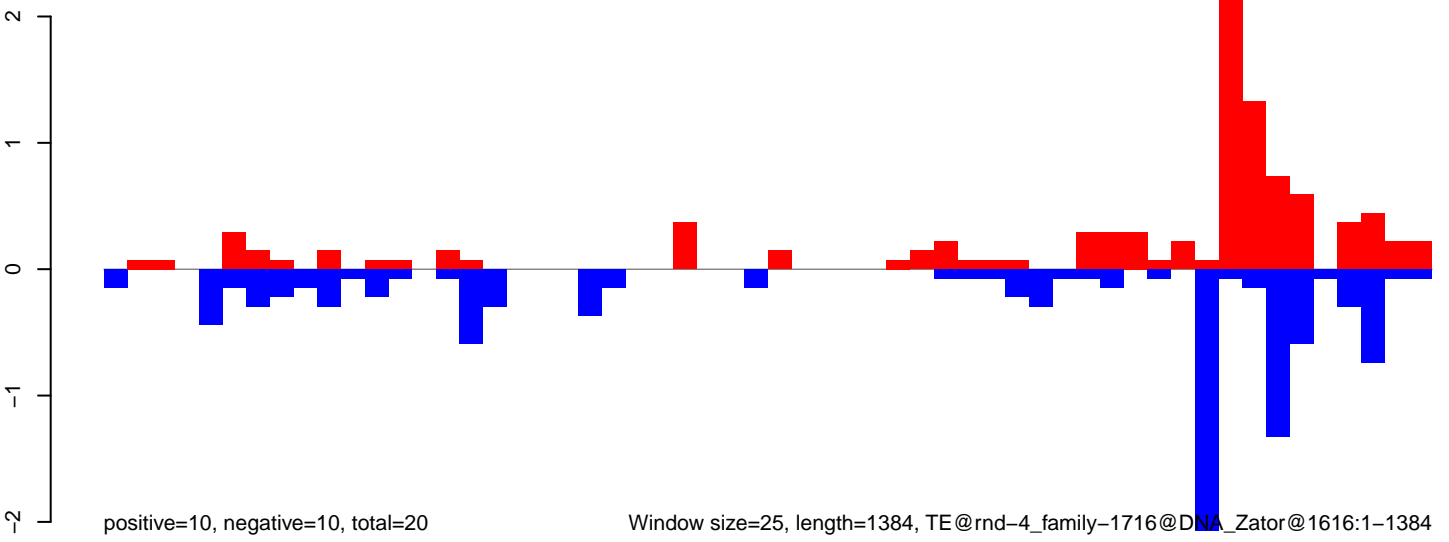
CuQuin_Testes_NL.18_23.rep



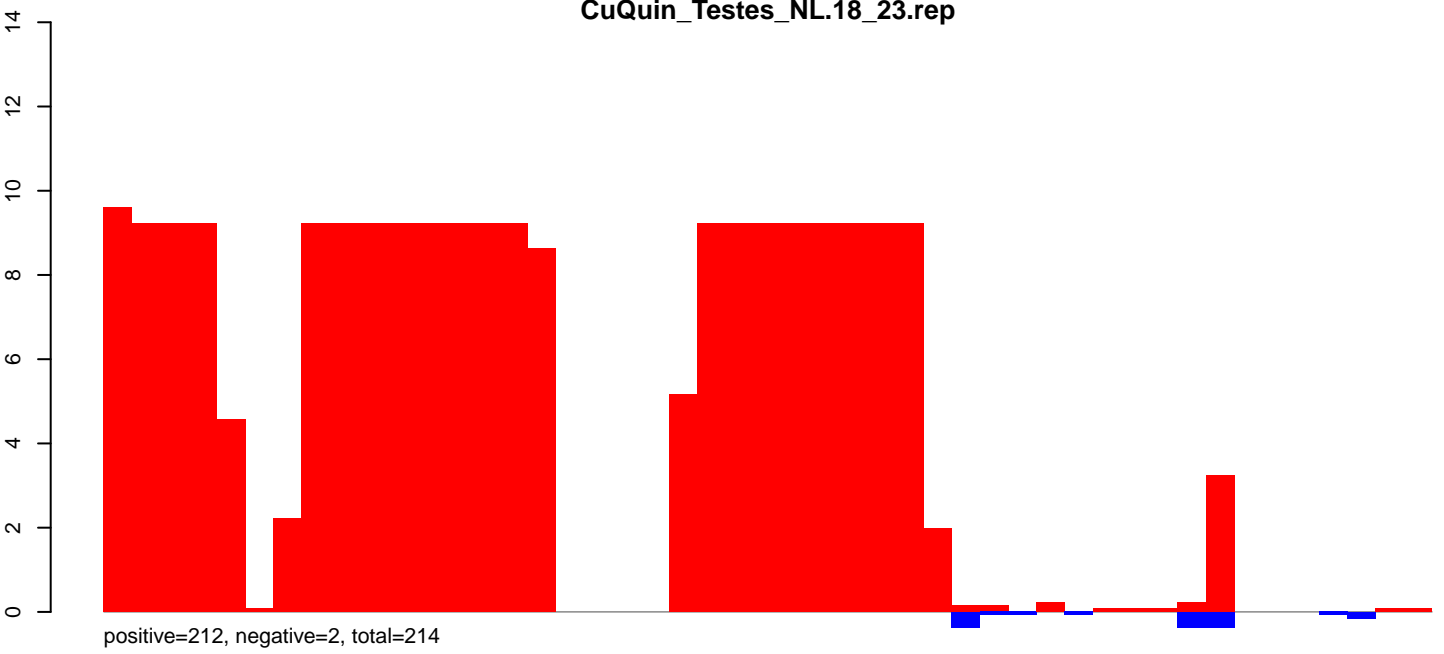
CuQuin_Testes_NL.24_35.rep



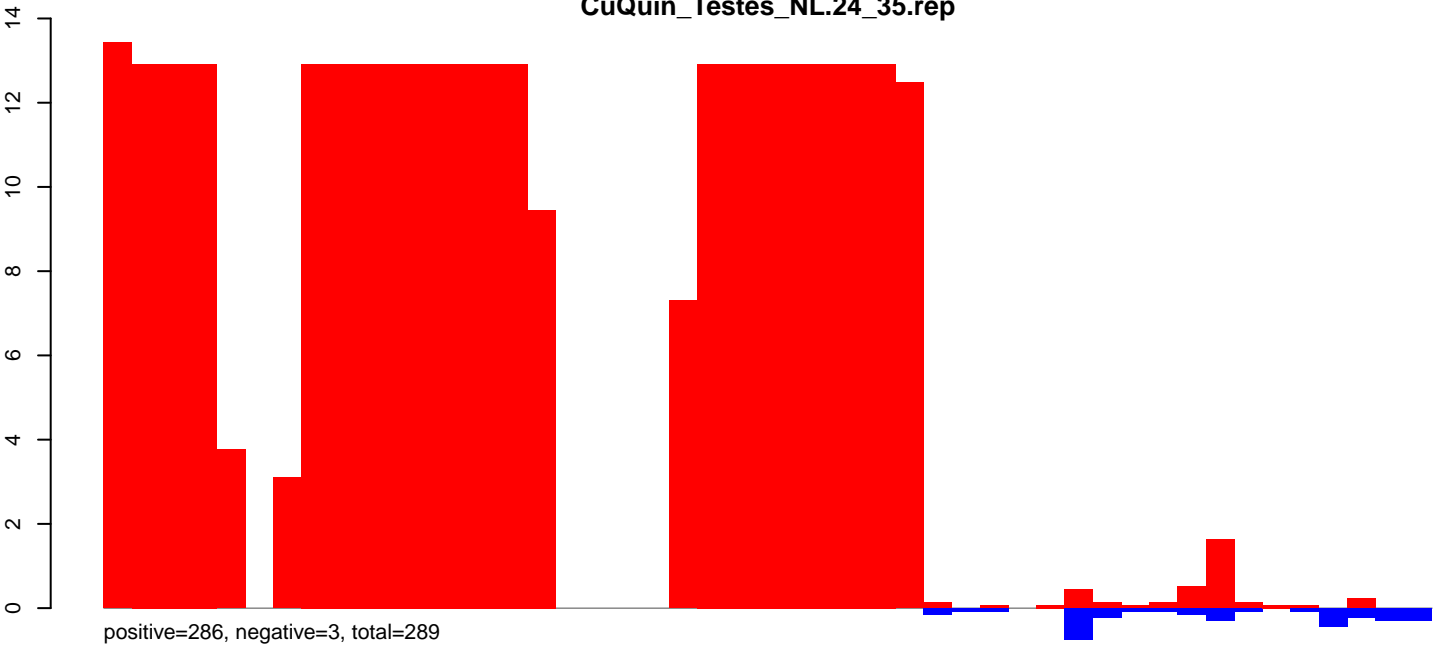
CuQuin_Testes_NL.rep



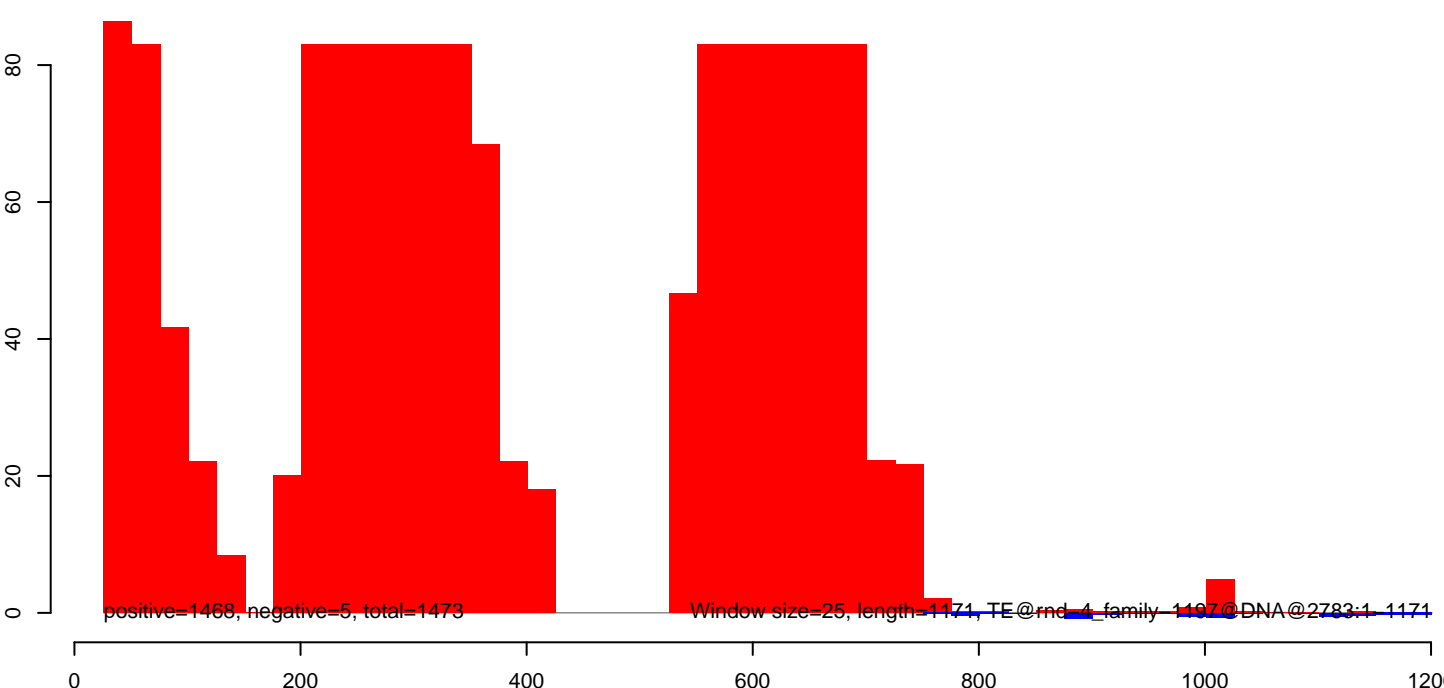
CuQuin_Testes_NL.18_23.rep



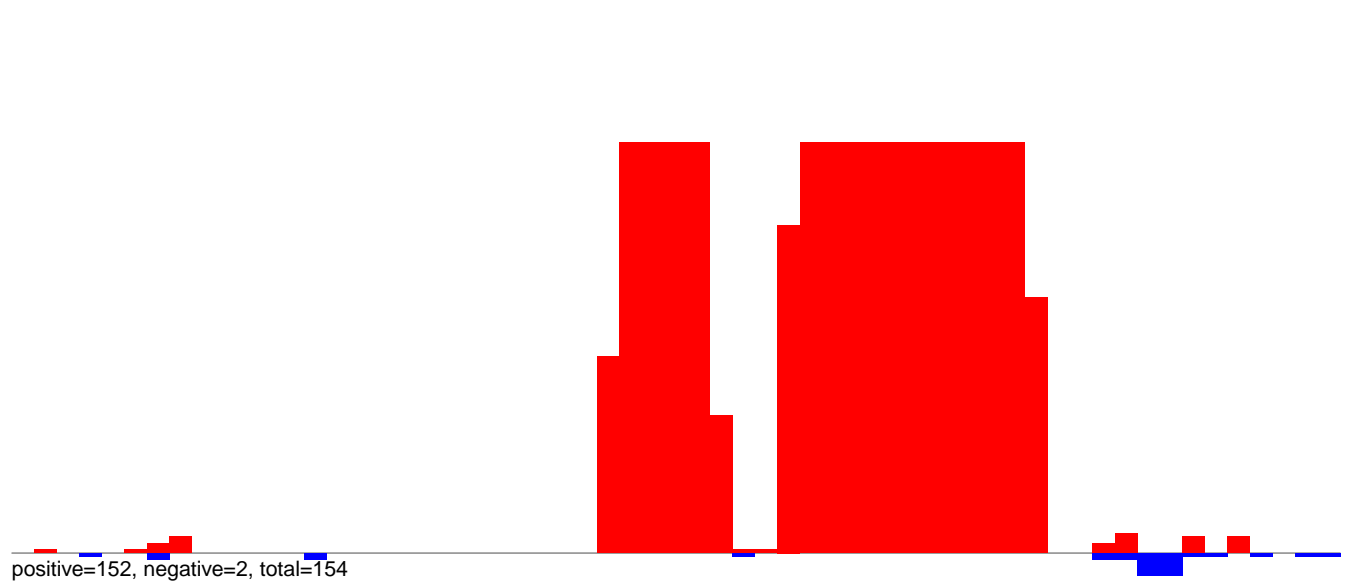
CuQuin_Testes_NL.24_35.rep



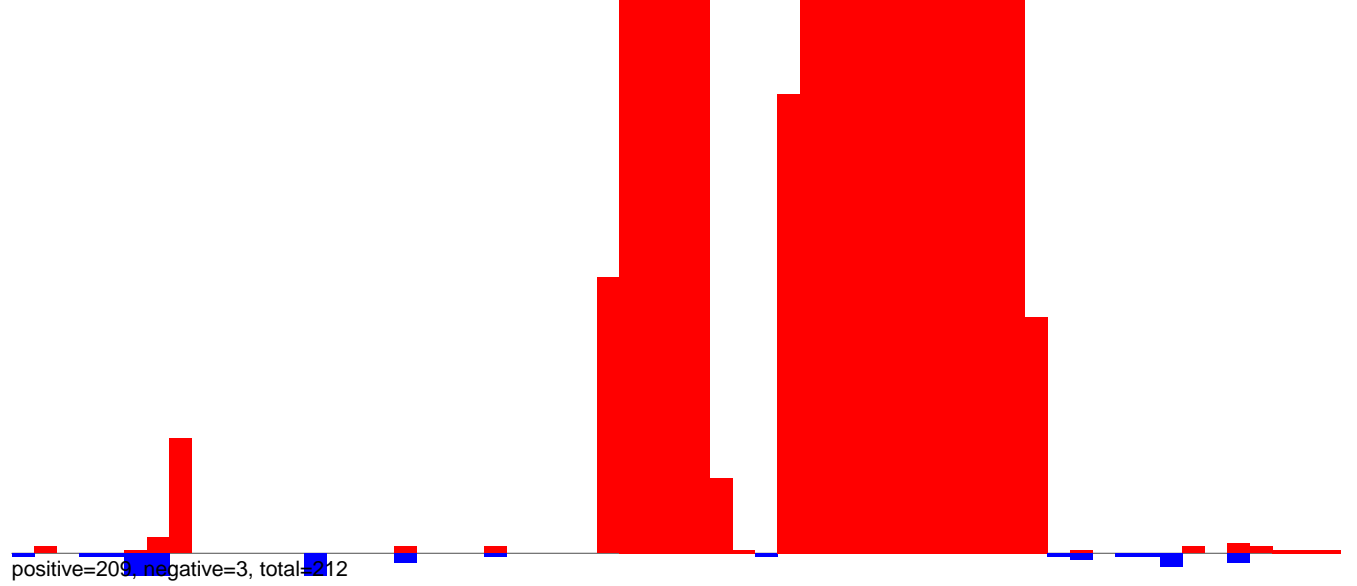
CuQuin_Testes_NL.rep



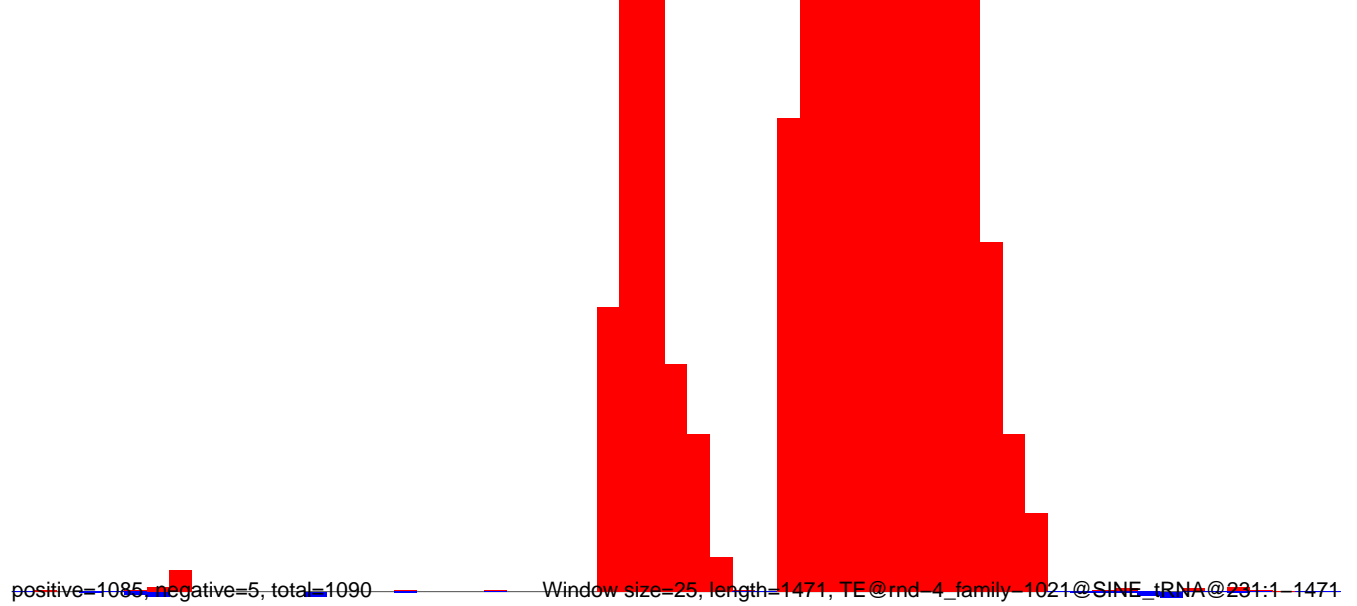
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

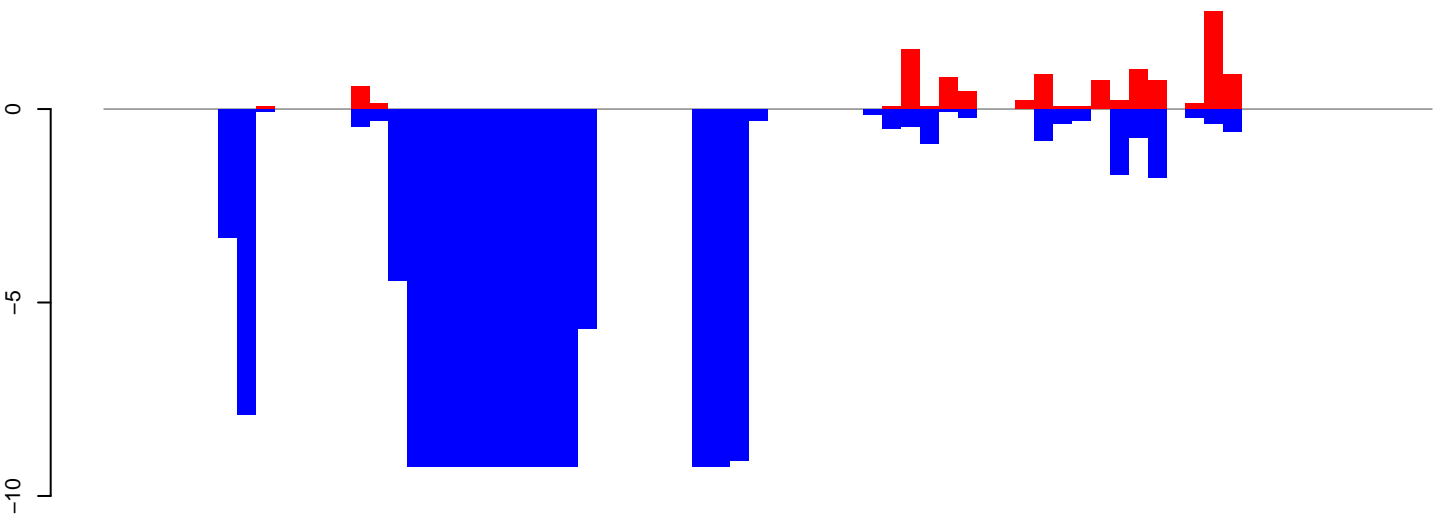


CuQuin_Testes_NL.rep



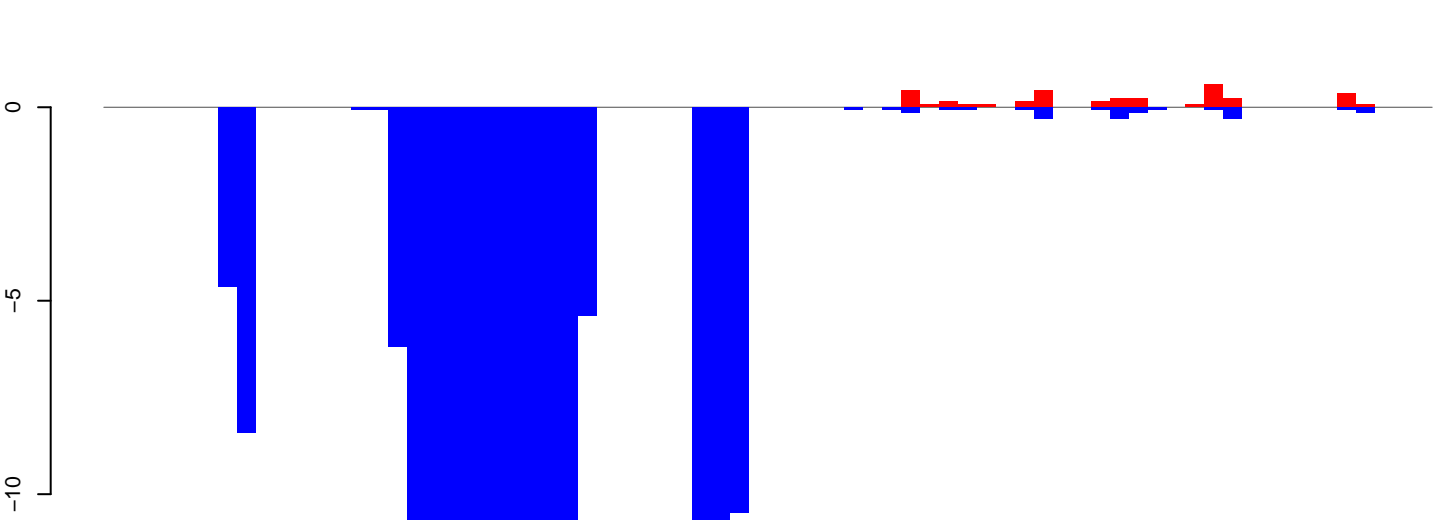
0 500 1000 1500

CuQuin_Testes_NL.18_23.rep



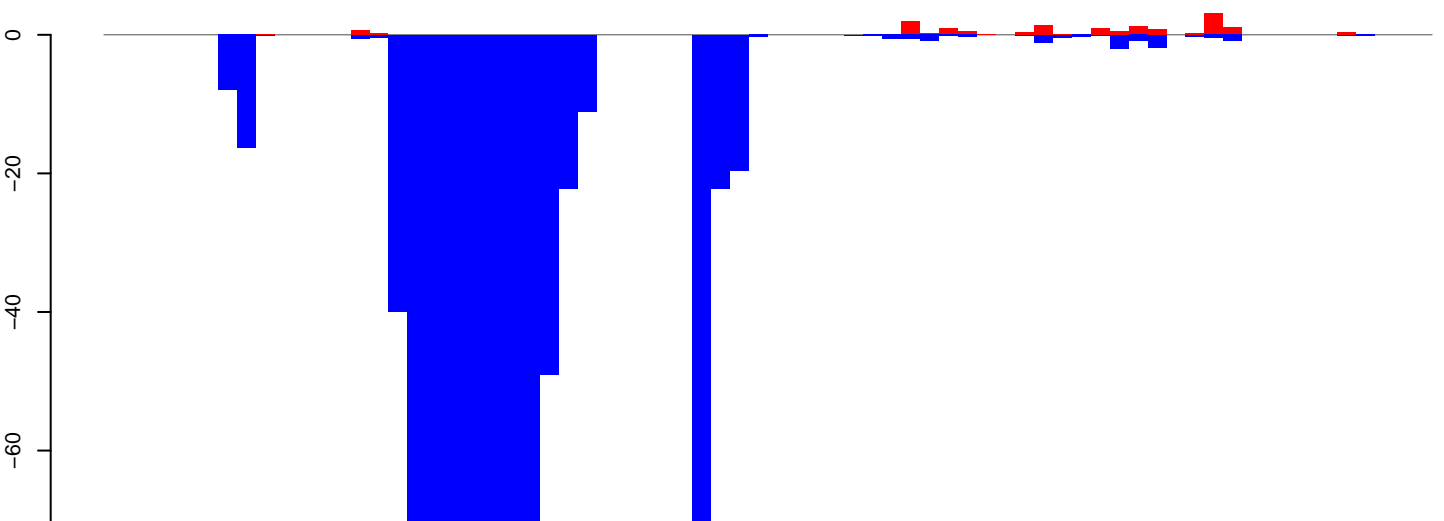
positive=11, negative=142, total=154

CuQuin_Testes_NL.24_35.rep



positive=3, negative=179, total=183

CuQuin_Testes_NL.rep

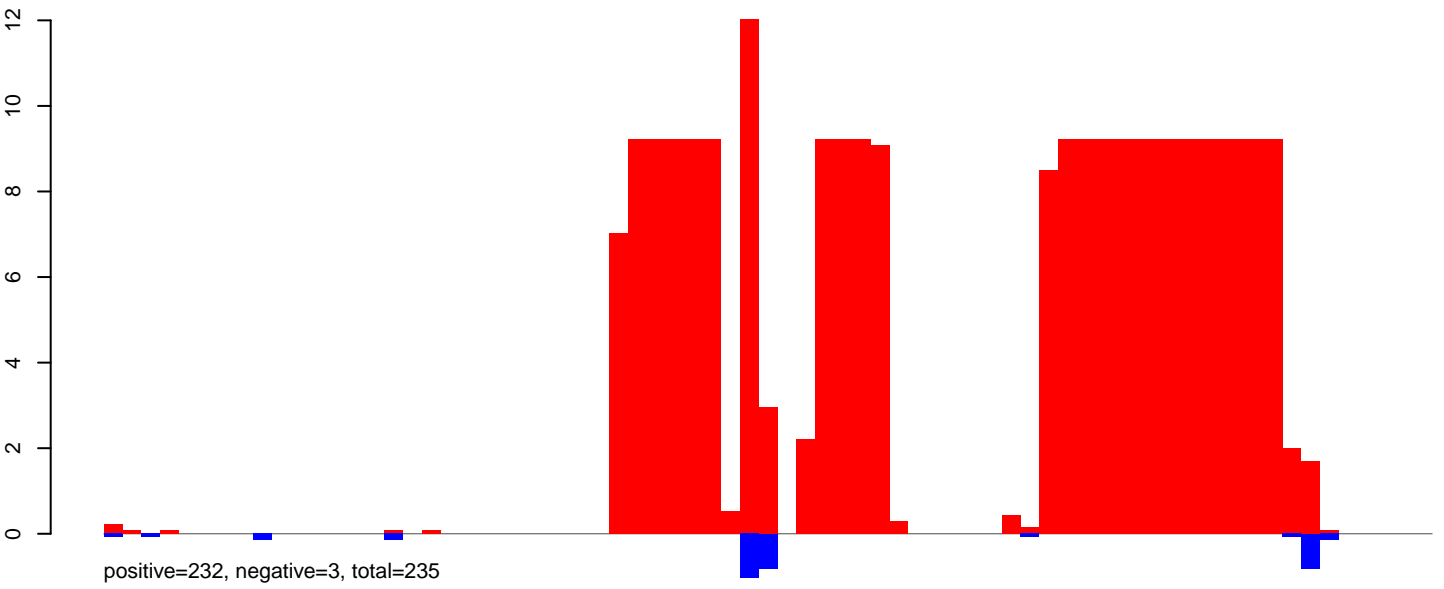


positive=15, negative=855, total=870

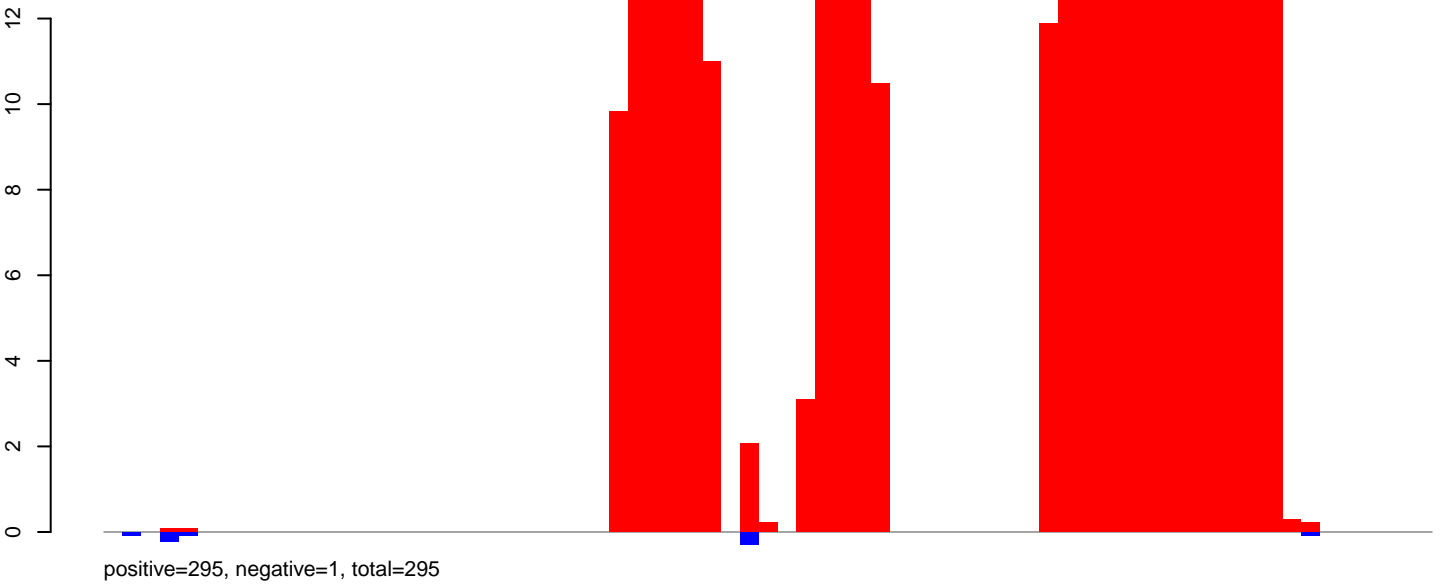
Window size=25, length=1741, TE@rnd-5_family-973@DNA@1859:1-1741

0 500 1000 1500

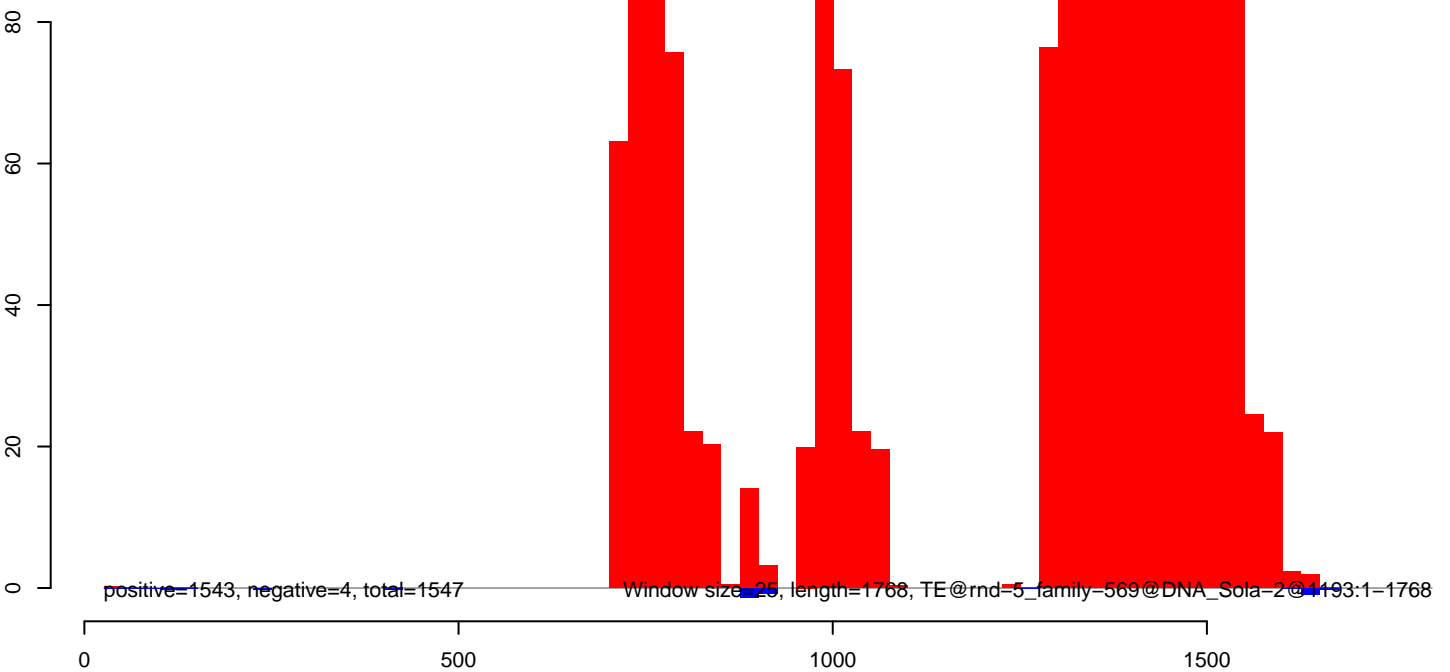
CuQuin_Testes_NL.18_23.rep



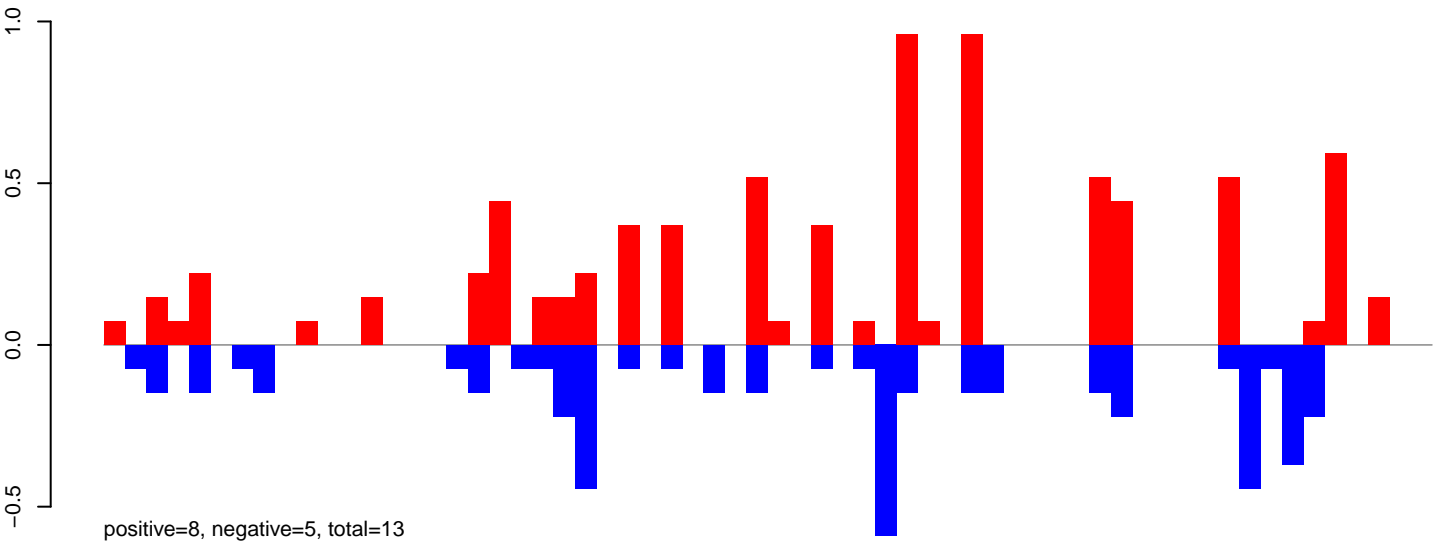
CuQuin_Testes_NL.24_35.rep



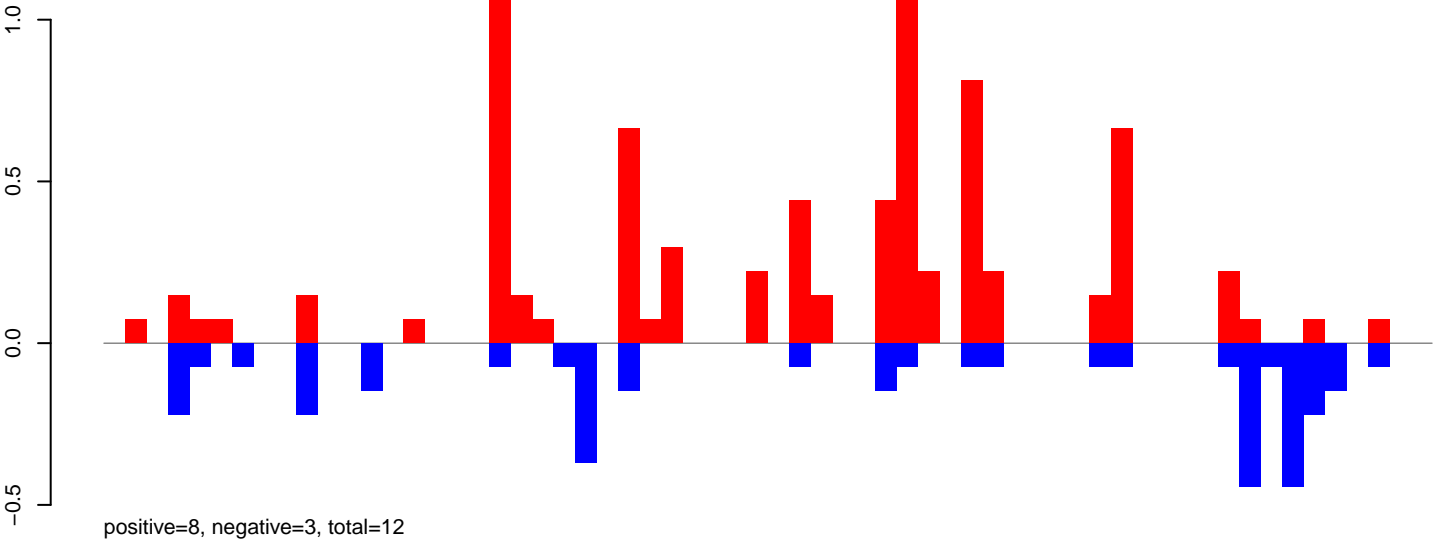
CuQuin_Testes_NL.rep



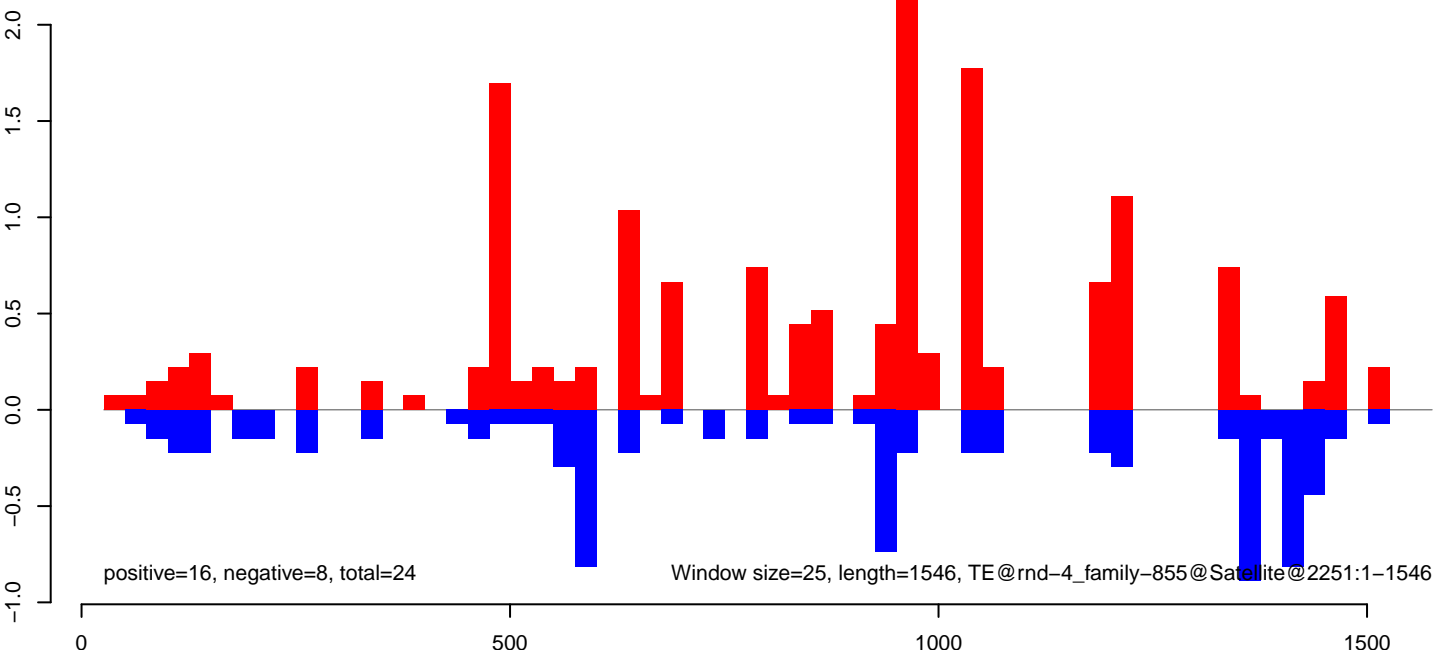
CuQuin_Testes_NL.18_23.rep



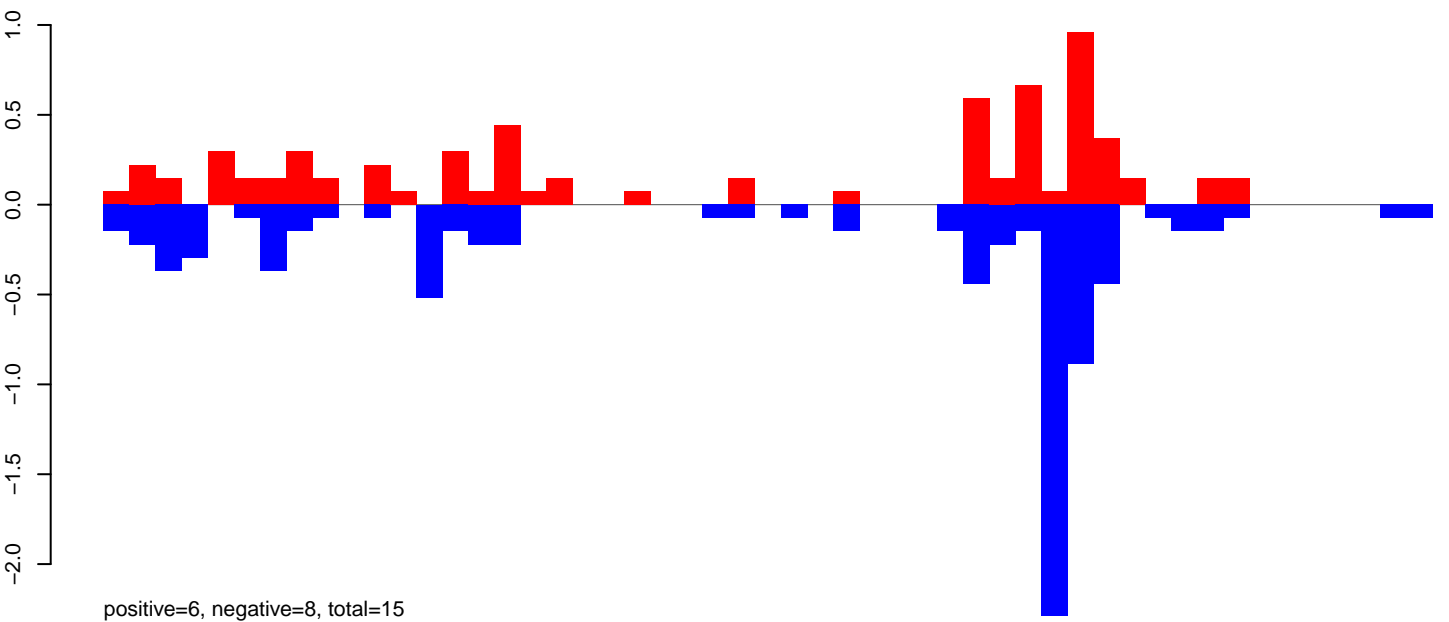
CuQuin_Testes_NL.24_35.rep



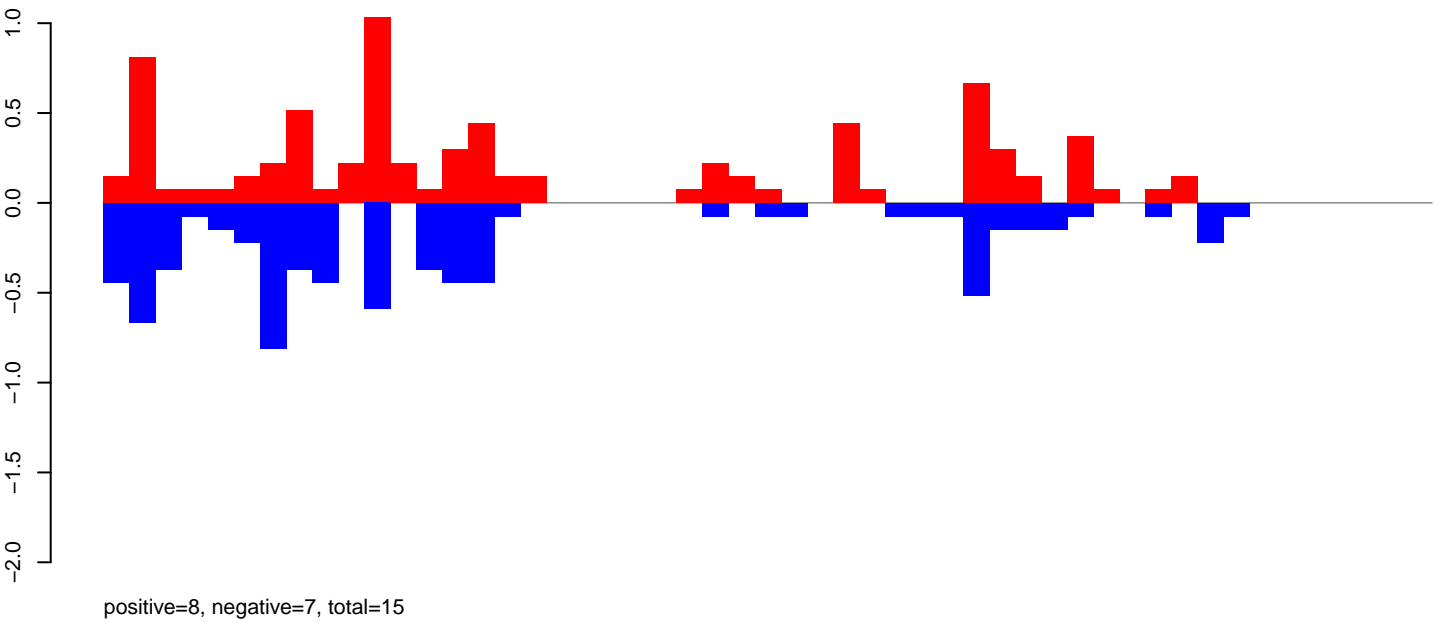
CuQuin_Testes_NL.rep



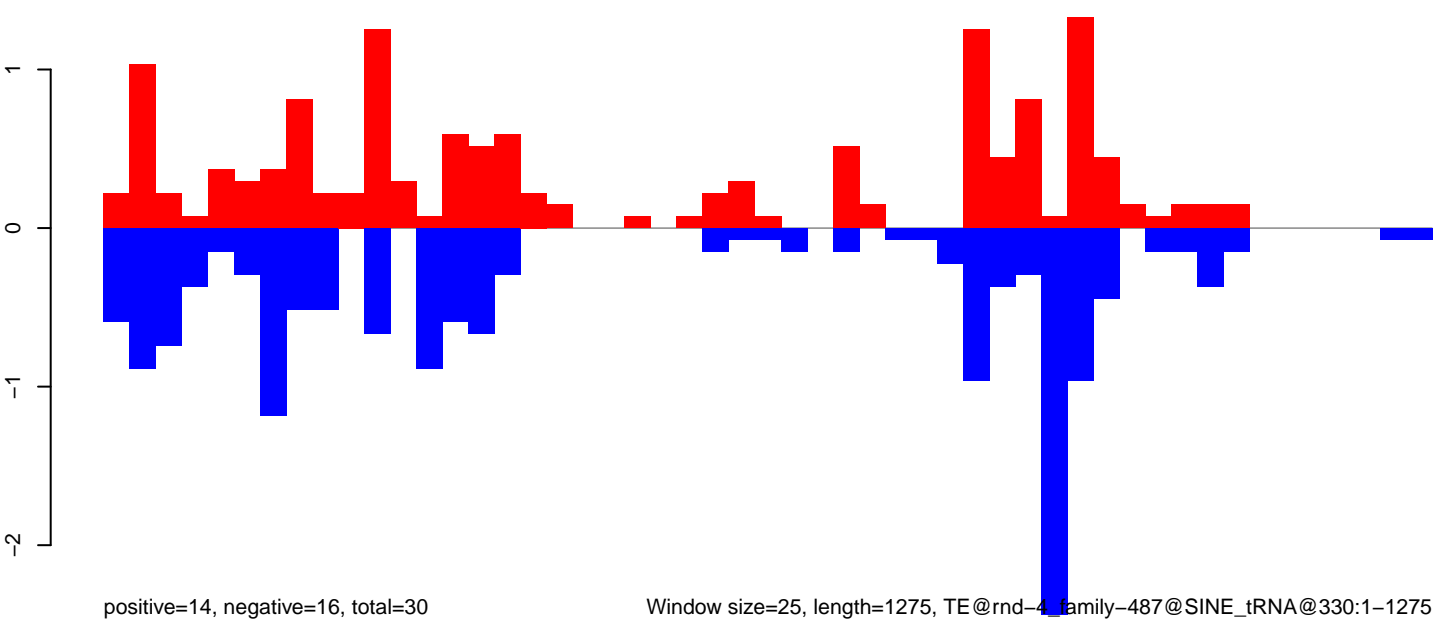
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



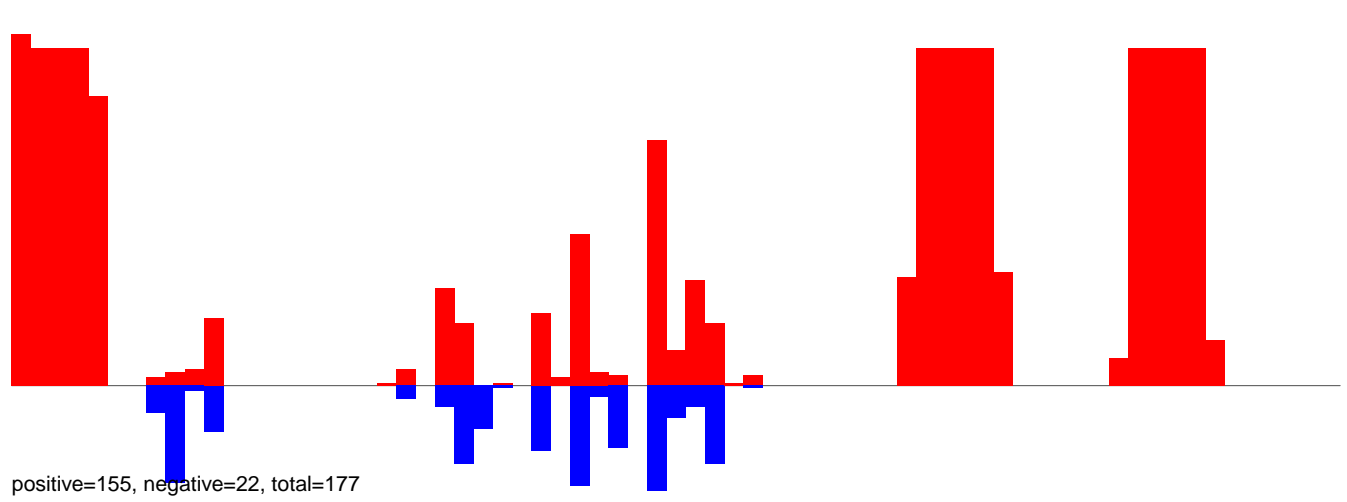
CuQuin_Testes_NL.rep



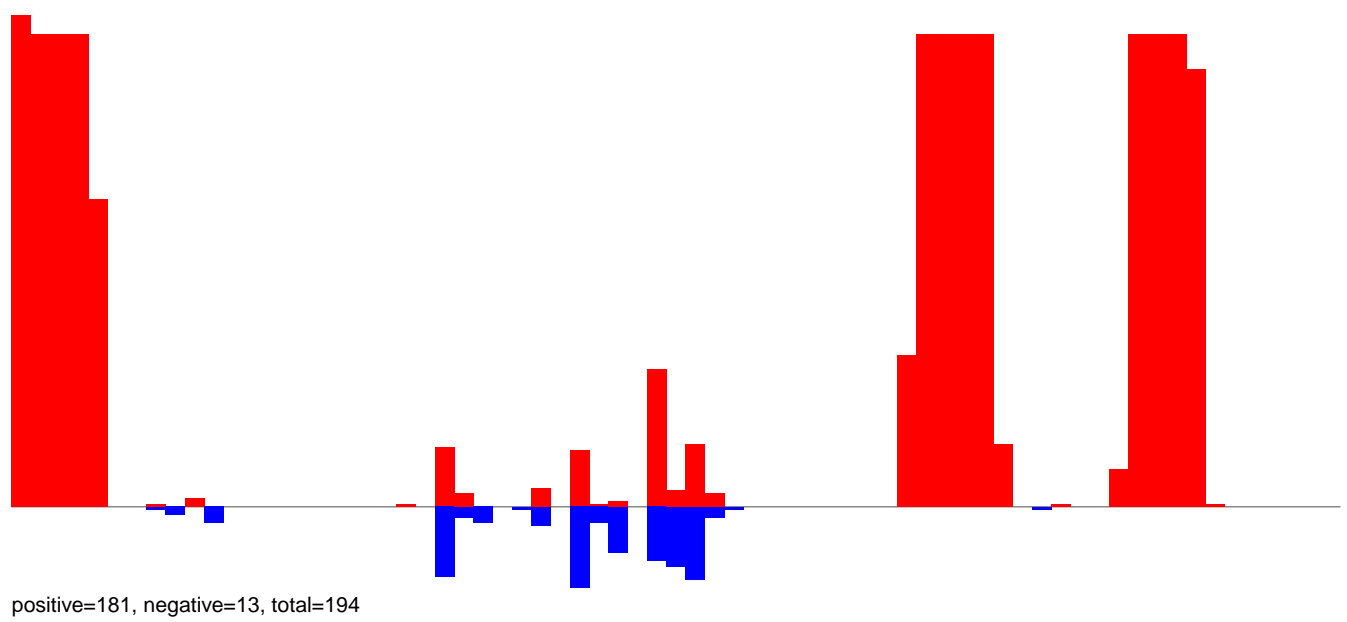
Window size=25, length=1275, TE@rnd-4, family-487@SINE_tRNA@330:1-1275

0 200 400 600 800 1000 1200

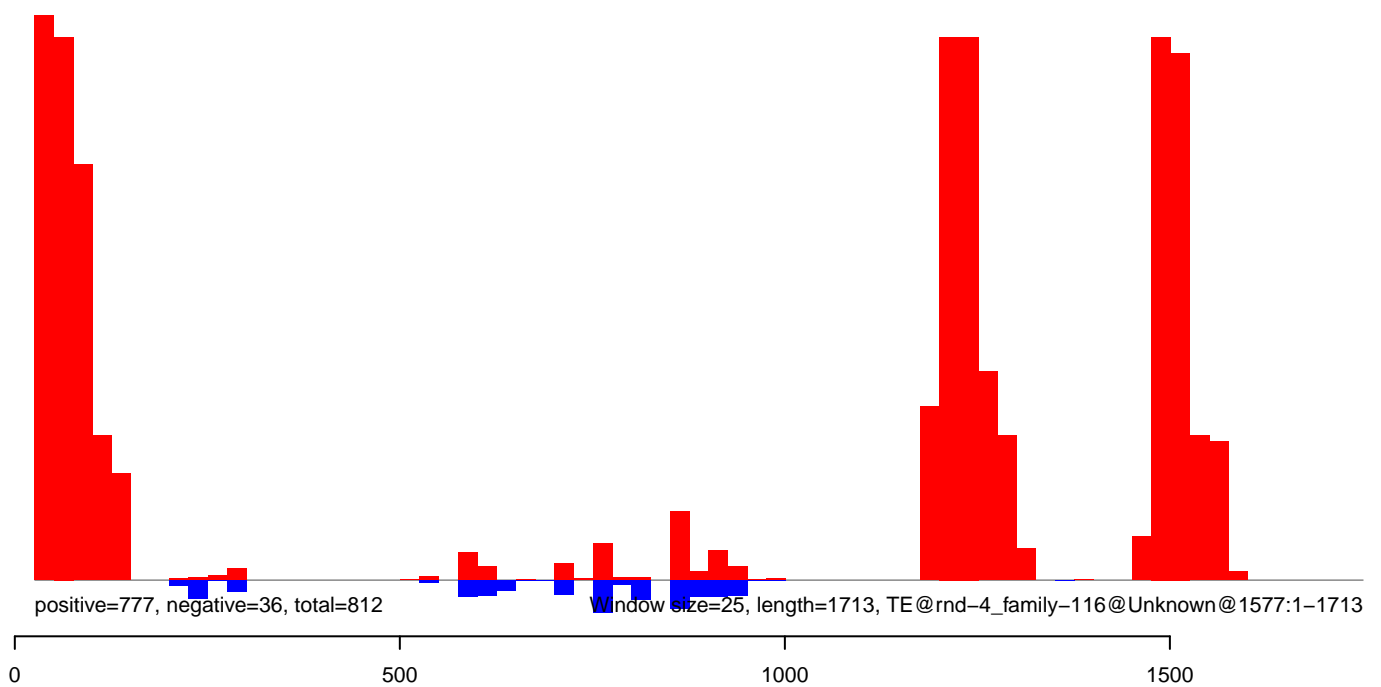
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

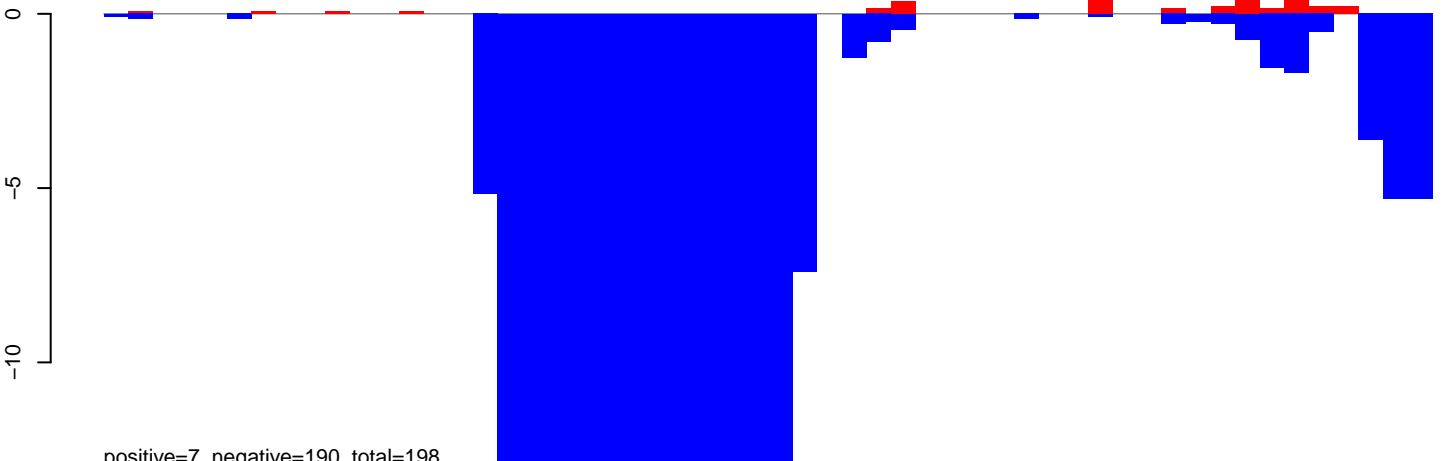


CuQuin_Testes_NL.18_23.rep



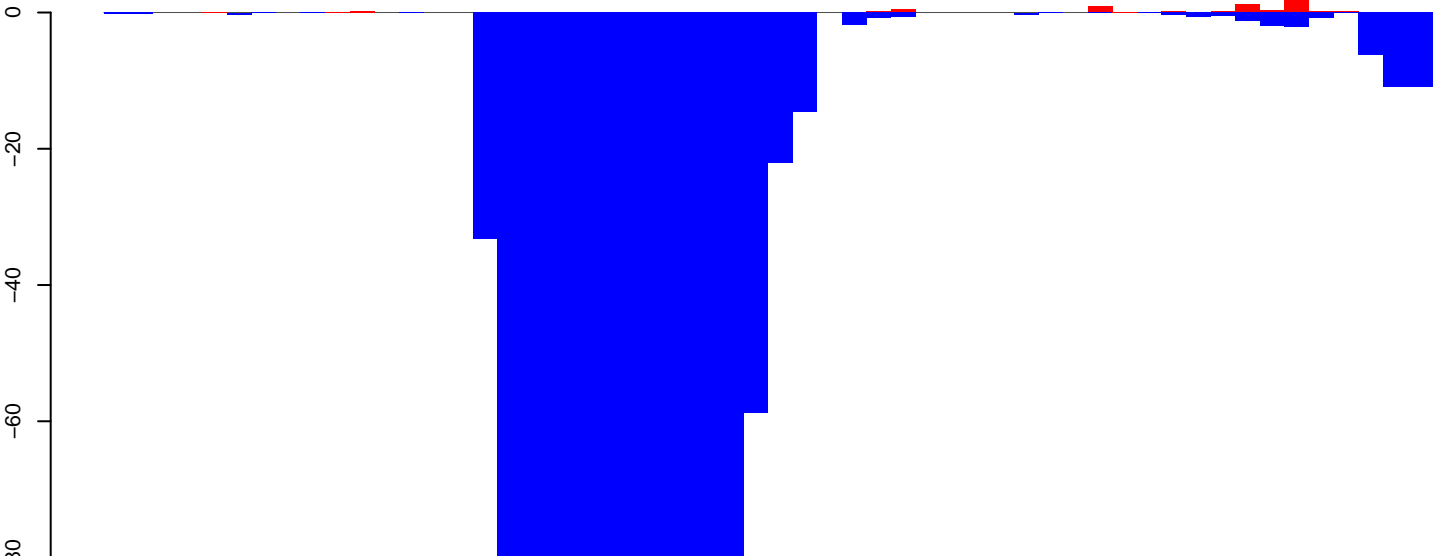
positive=2, negative=139, total=141

CuQuin_Testes_NL.24_35.rep



positive=7, negative=190, total=198

CuQuin_Testes_NL.rep

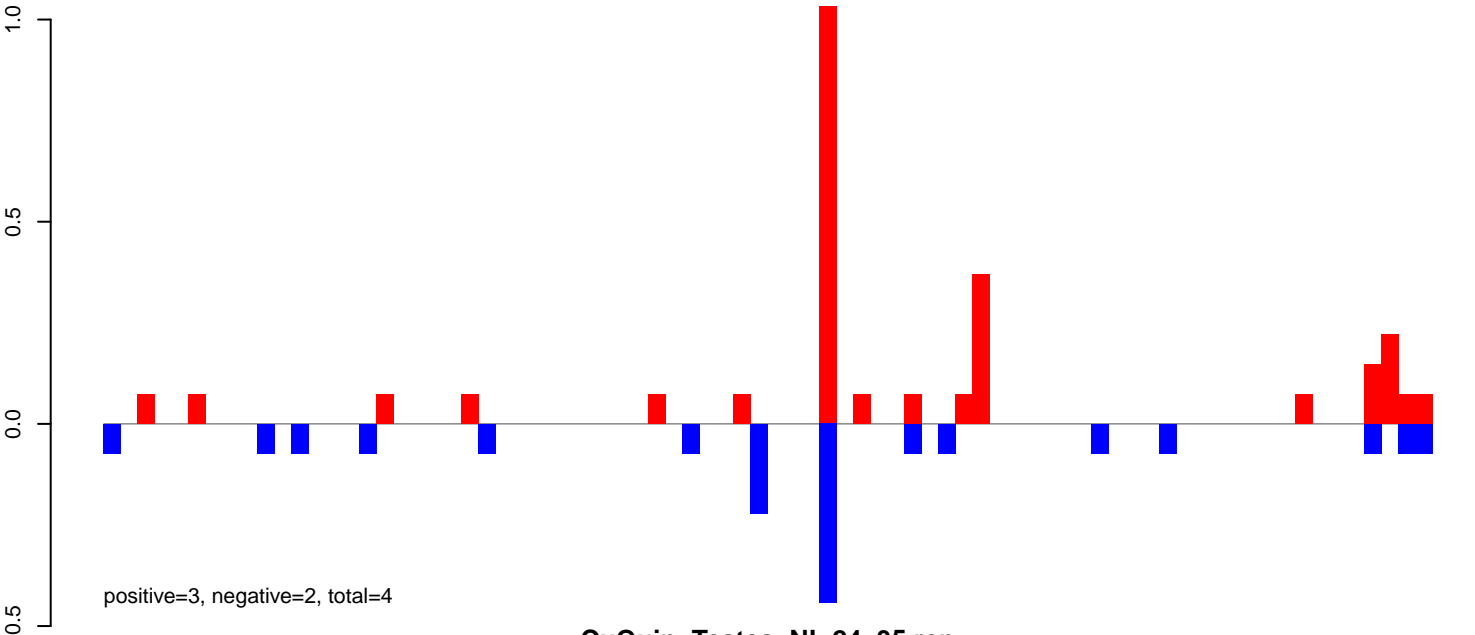


positive=9, negative=1000, total=1009

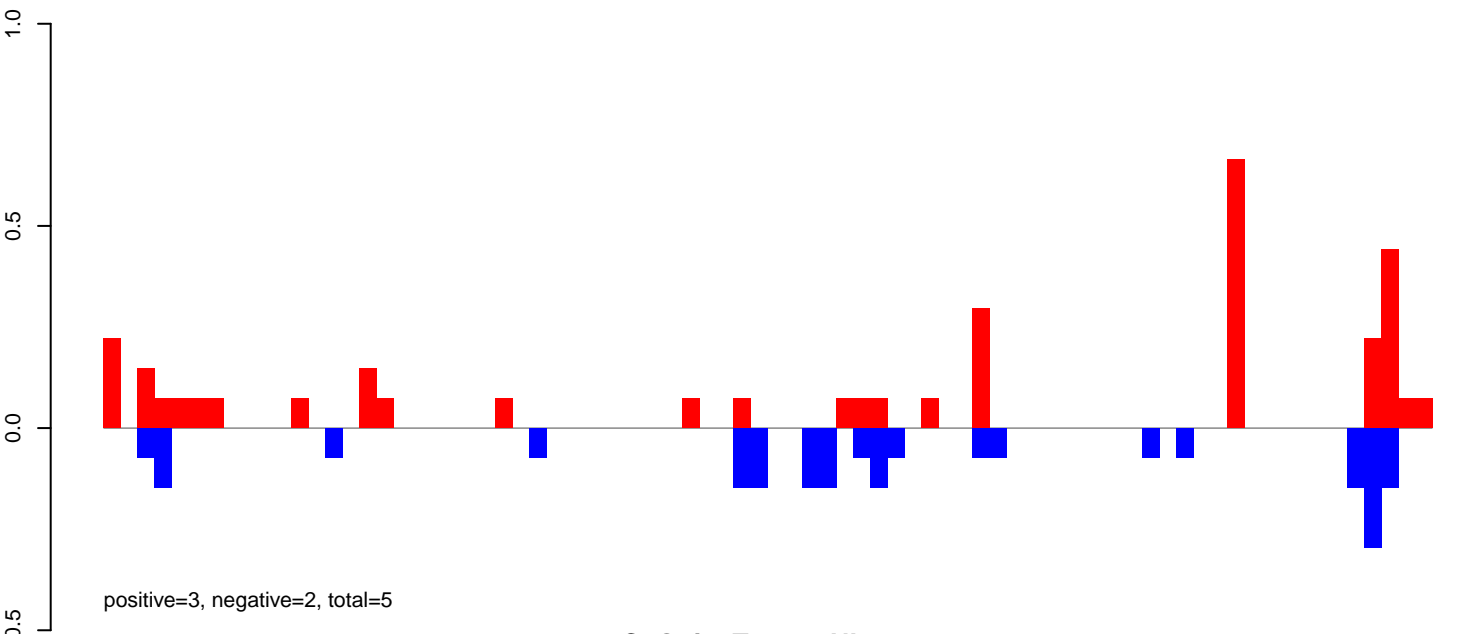
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0 200 400 600 800 1000 1200 1400

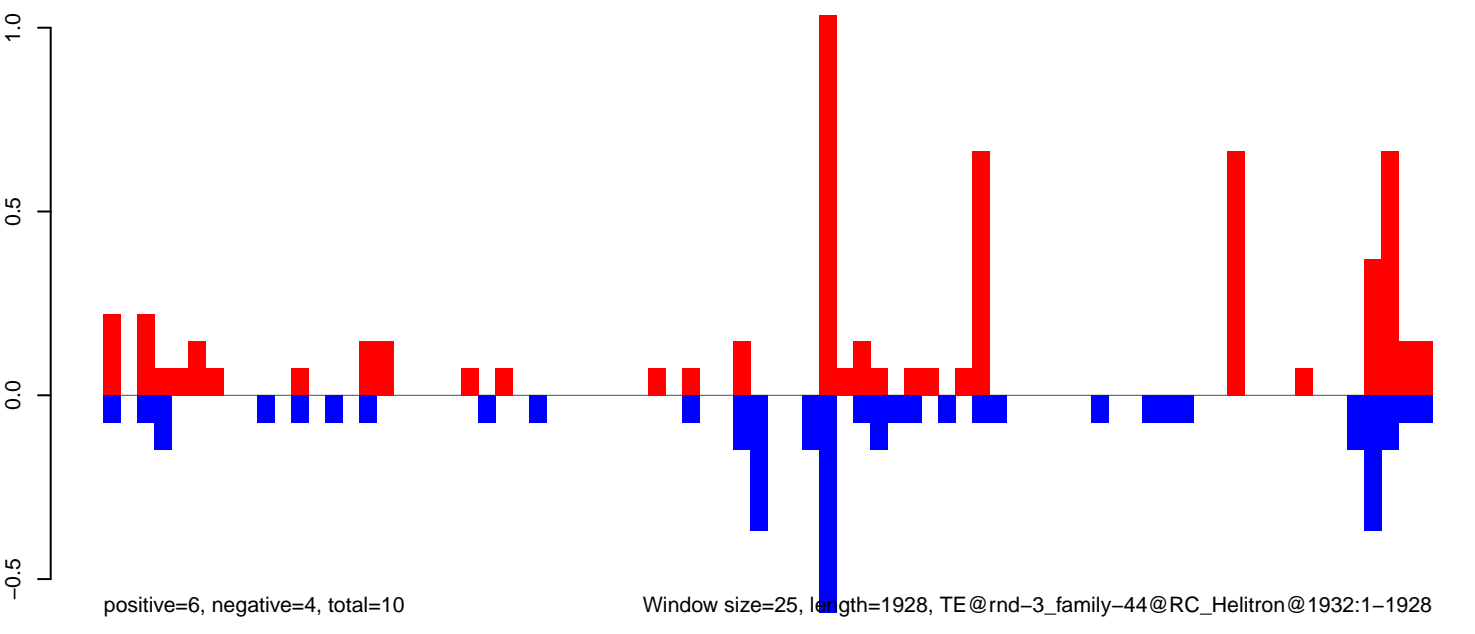
CuQuin_Testes_NL.18_23.rep



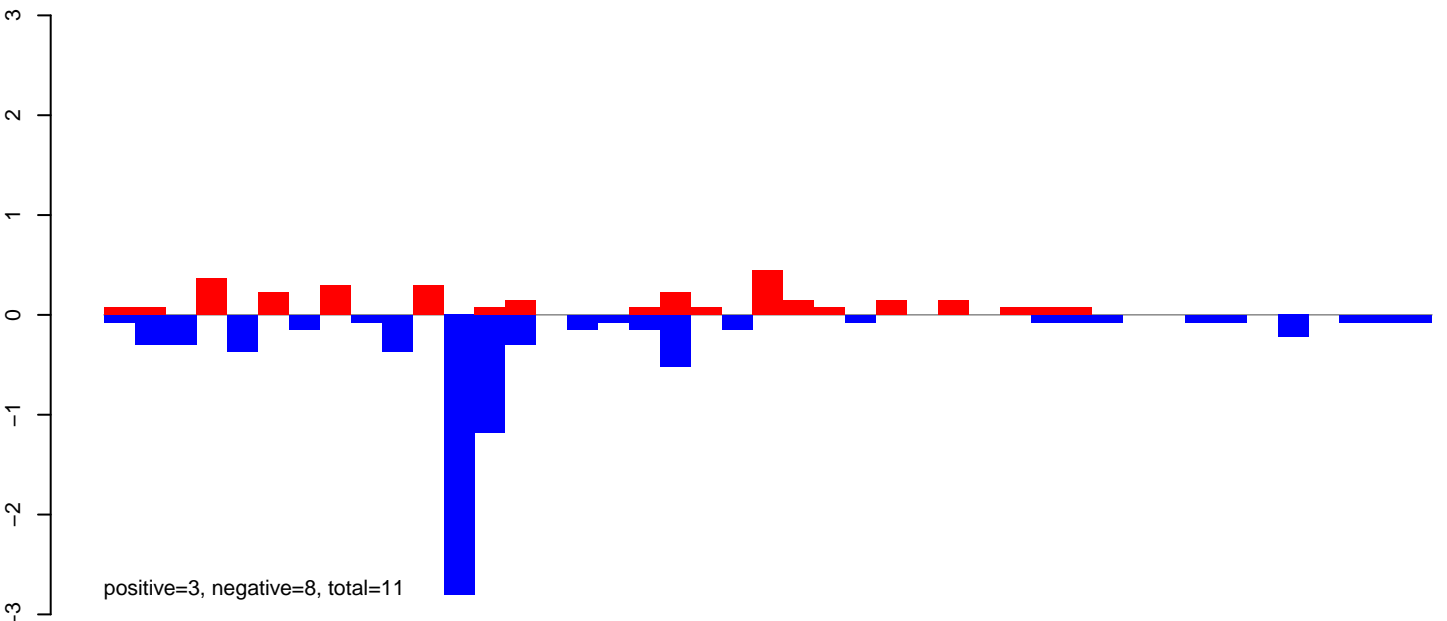
CuQuin_Testes_NL.24_35.rep



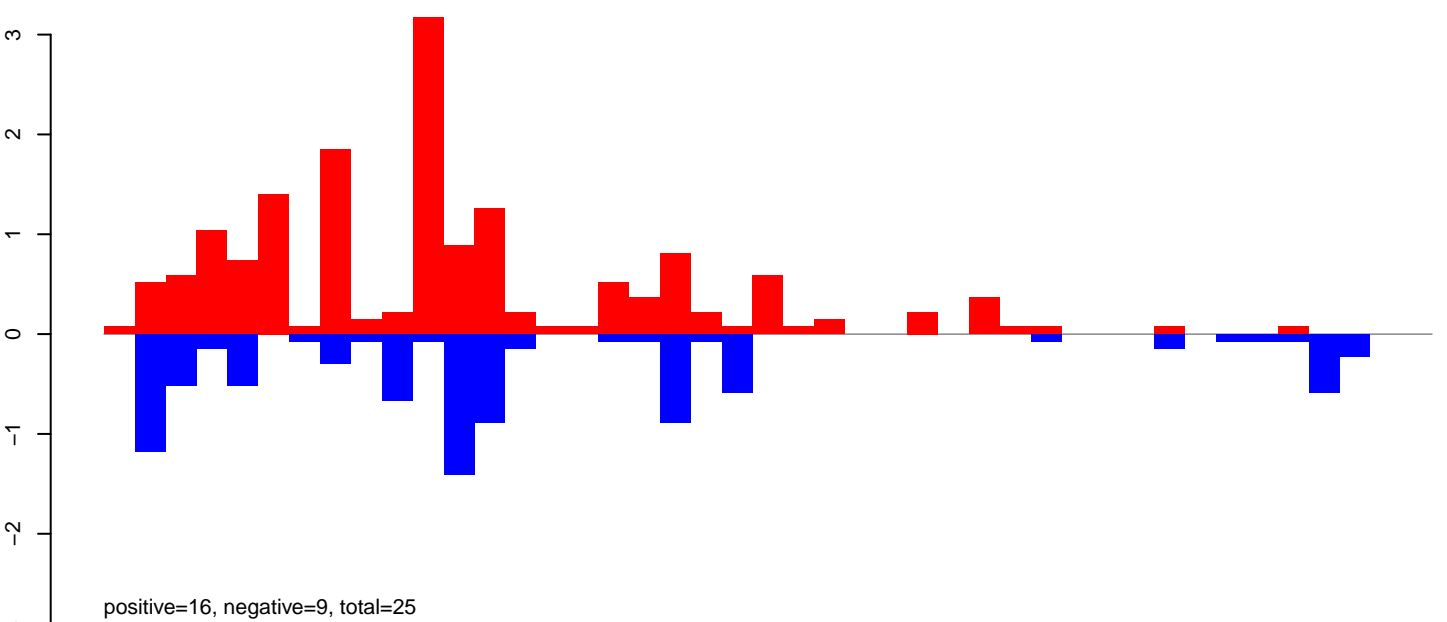
CuQuin_Testes_NL.rep



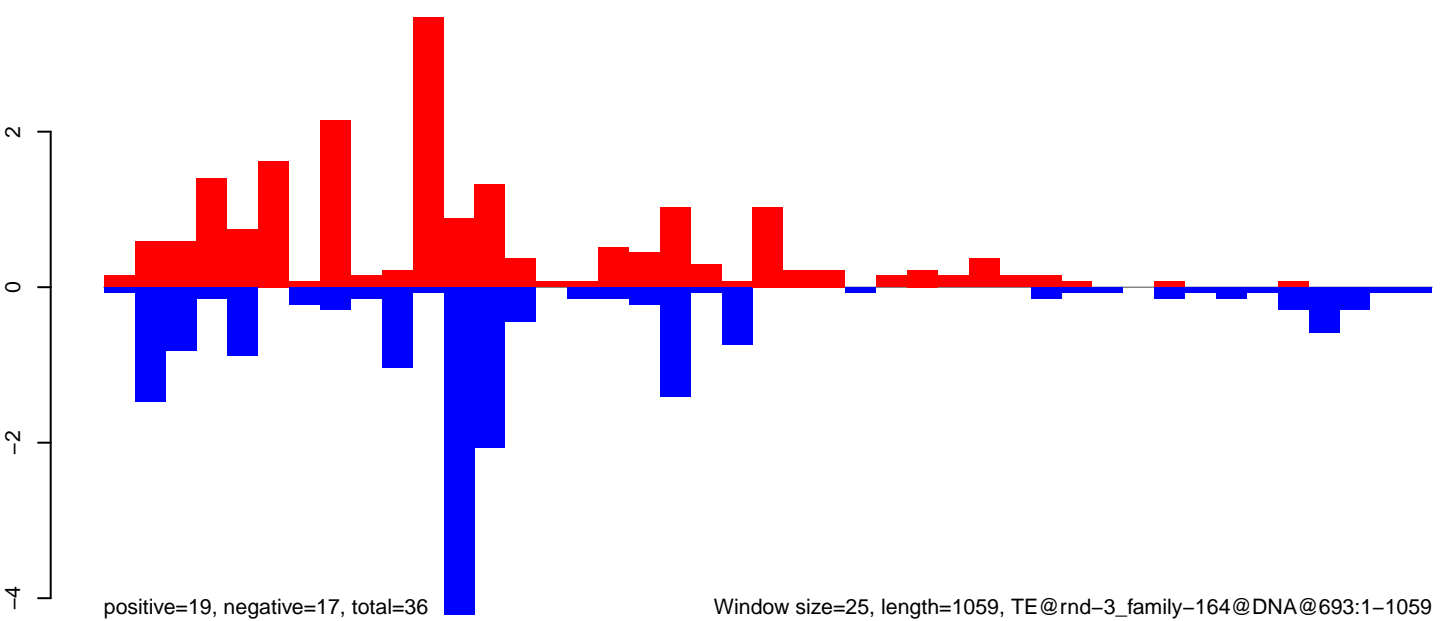
CuQuin_Testes_NL.18_23.rep



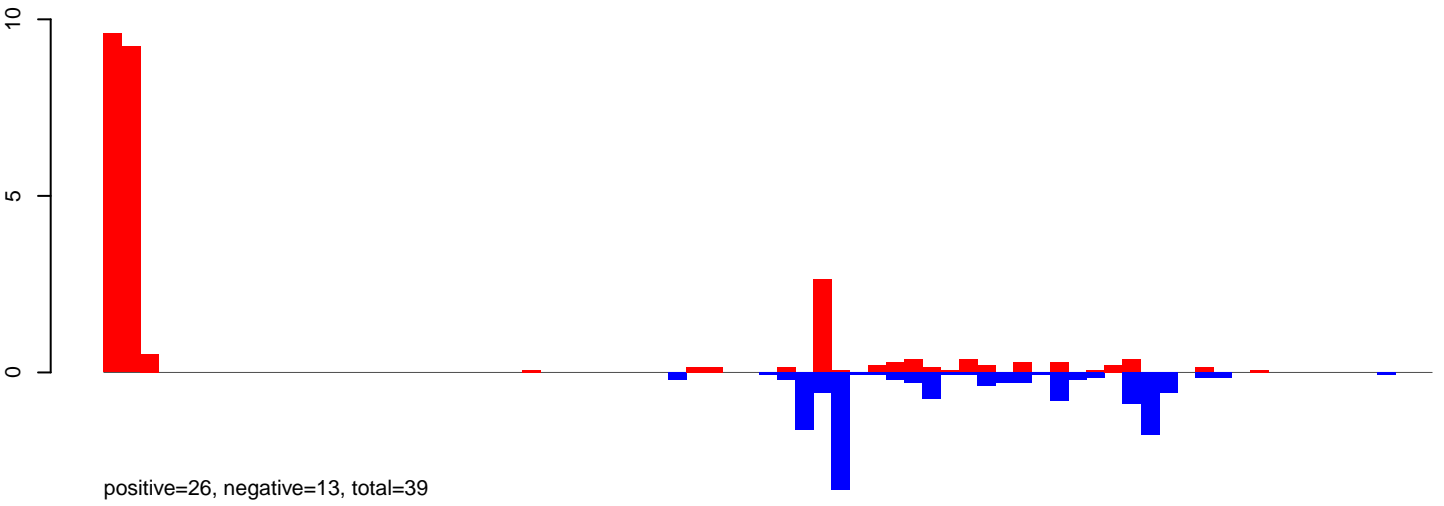
CuQuin_Testes_NL.24_35.rep



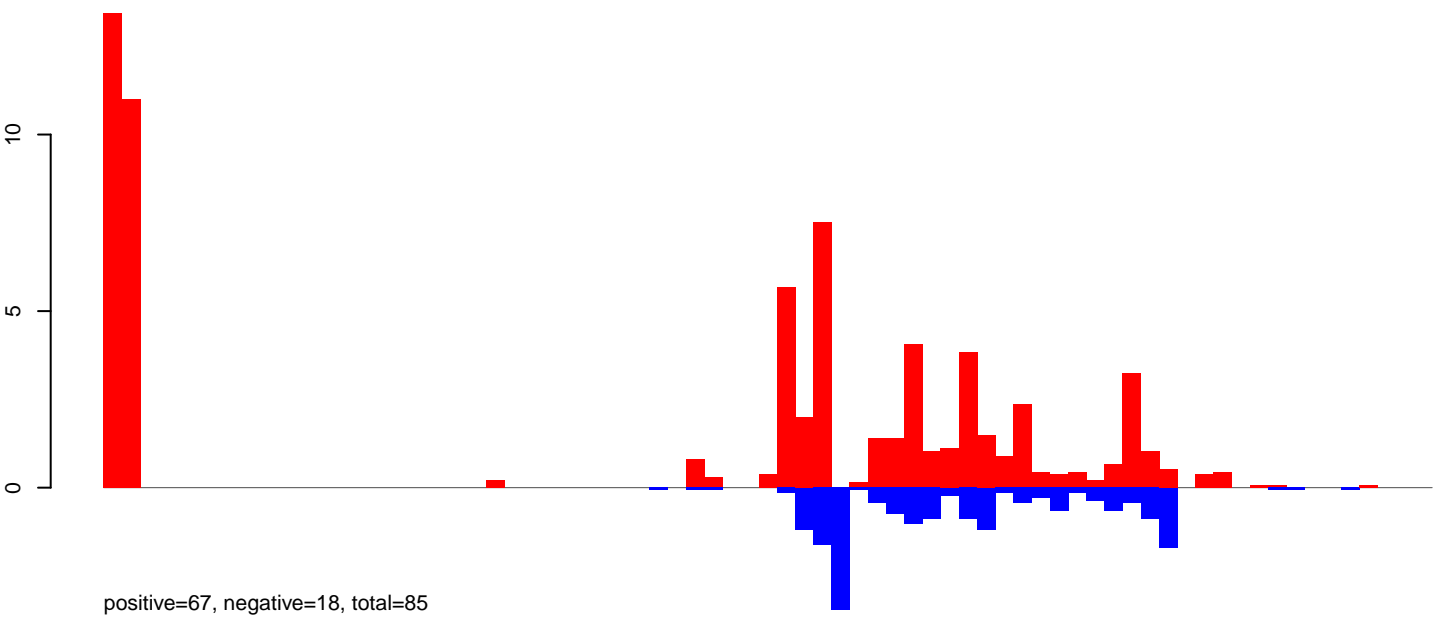
CuQuin_Testes_NL.rep



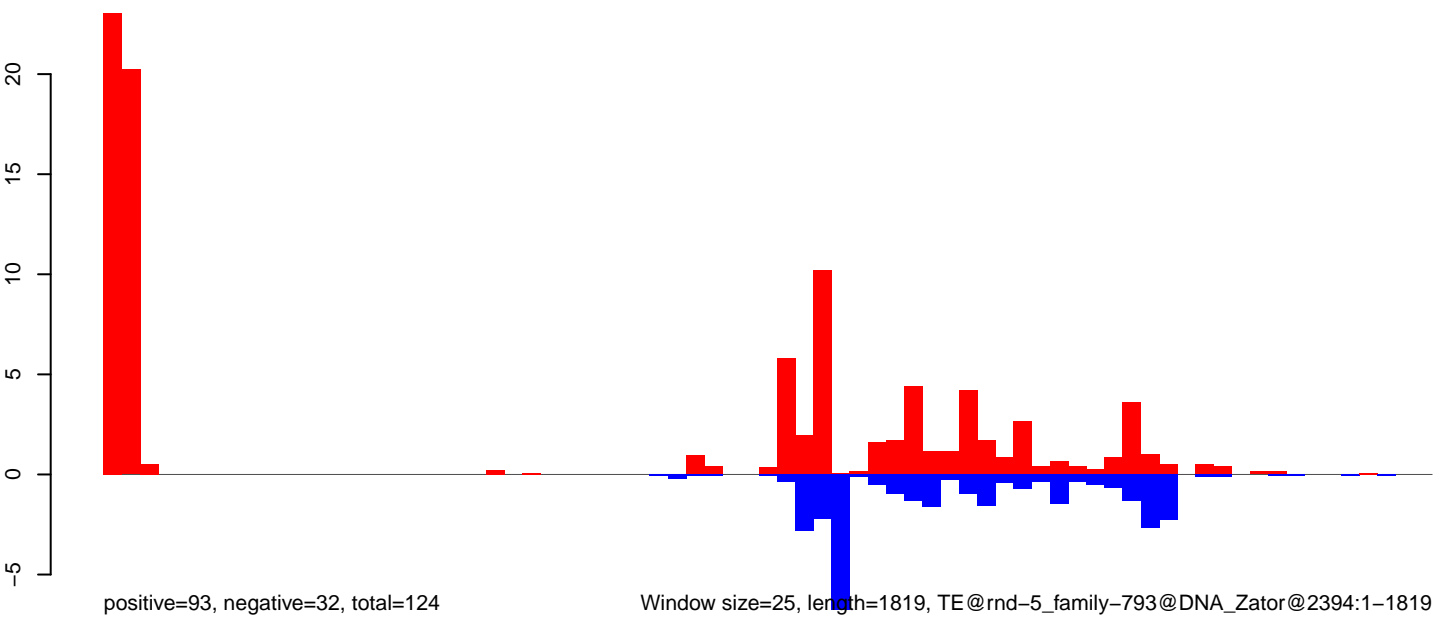
CuQuin_Testes_NL.18_23.rep



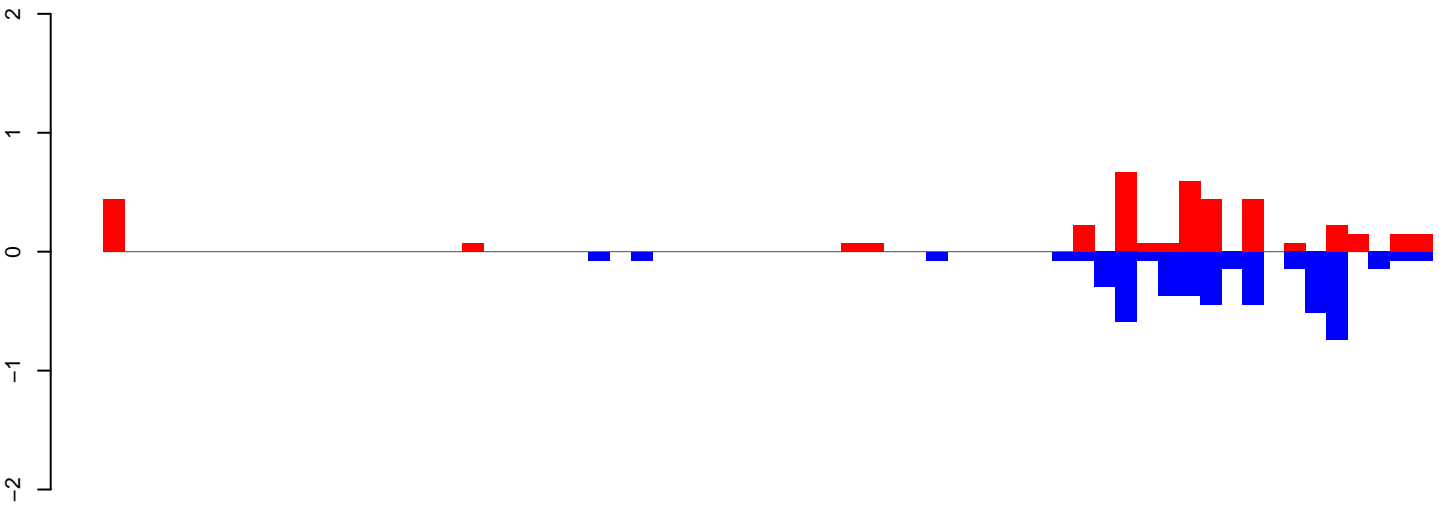
CuQuin_Testes_NL.24_35.rep



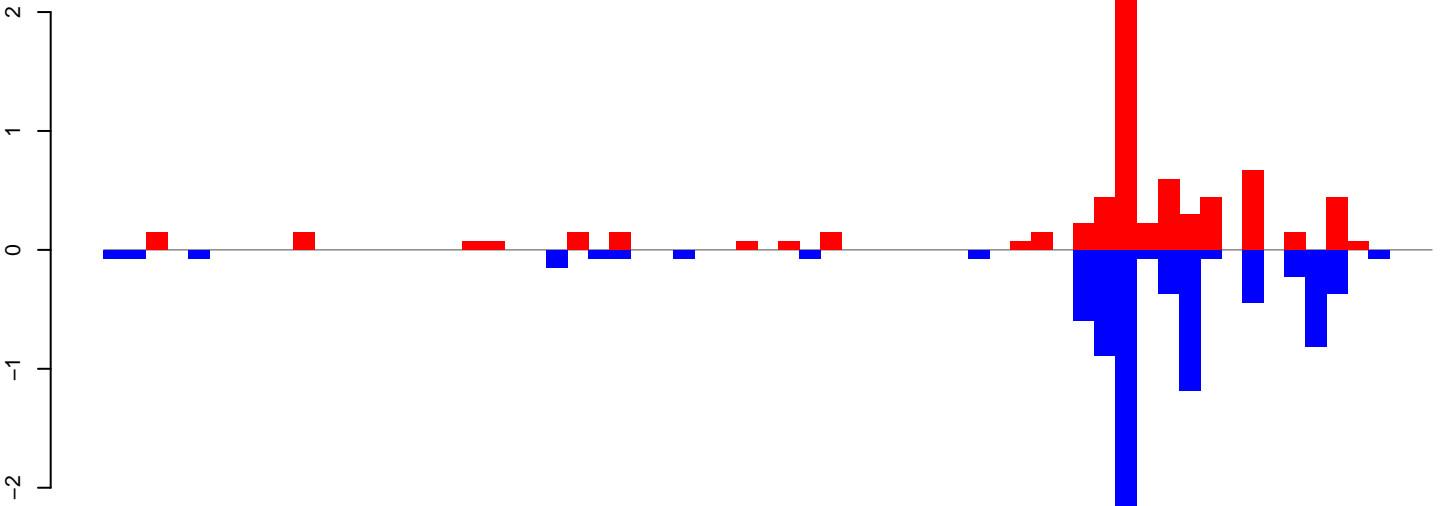
CuQuin_Testes_NL.rep



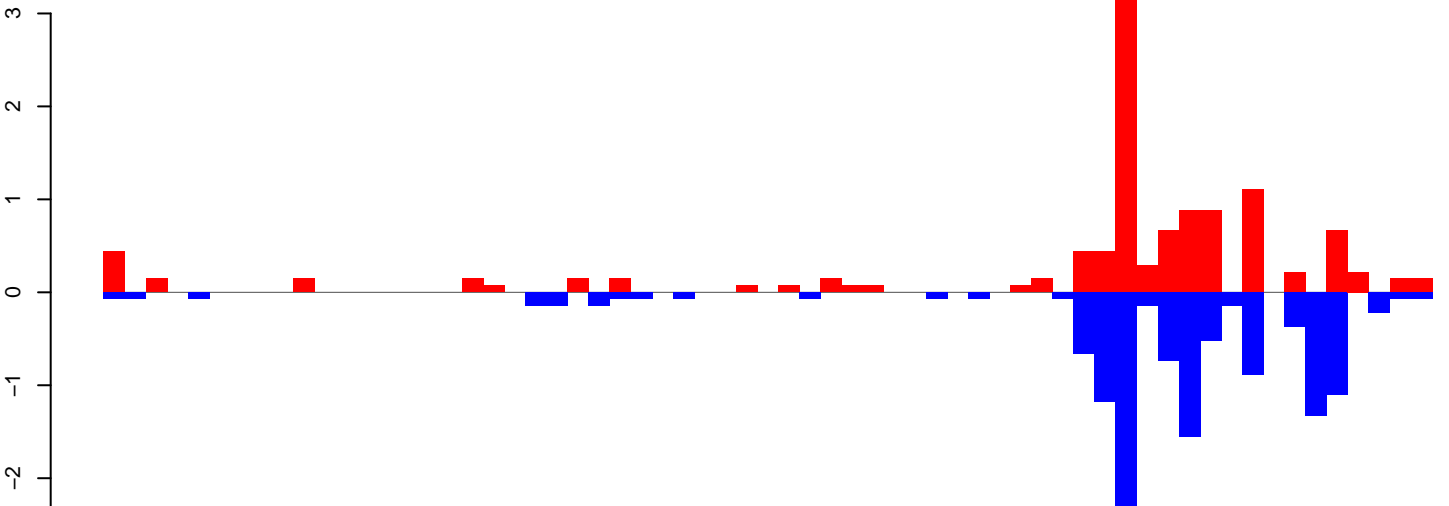
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

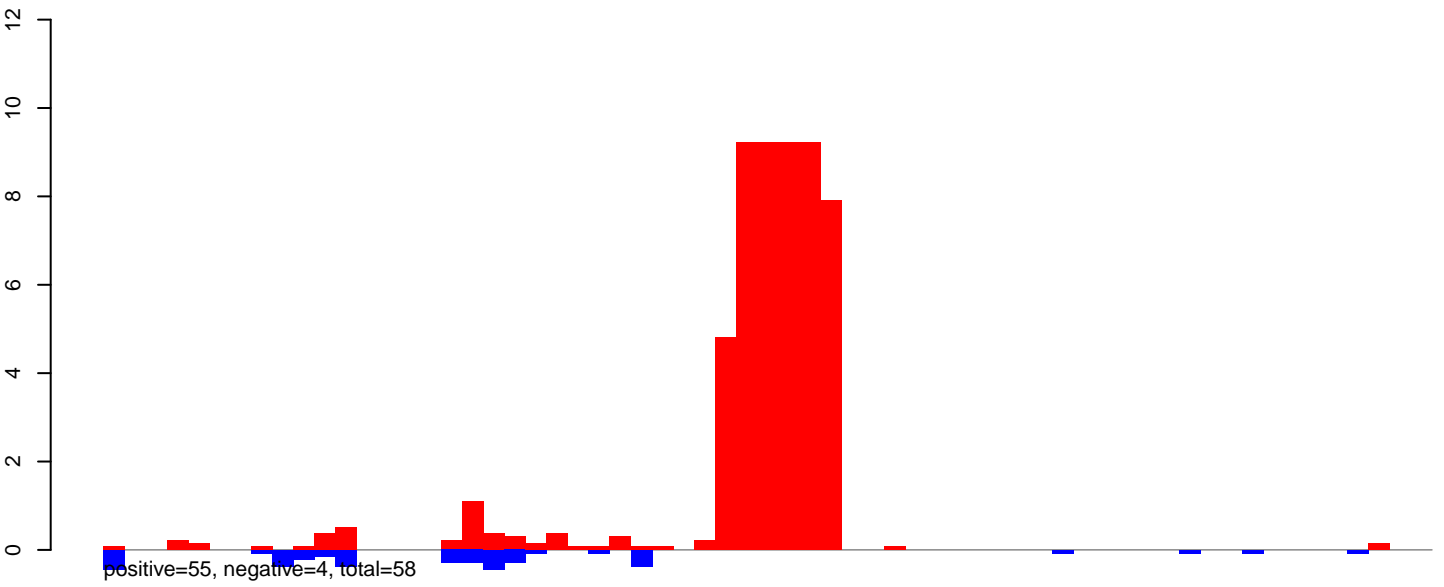


CuQuin_Testes_NL.rep

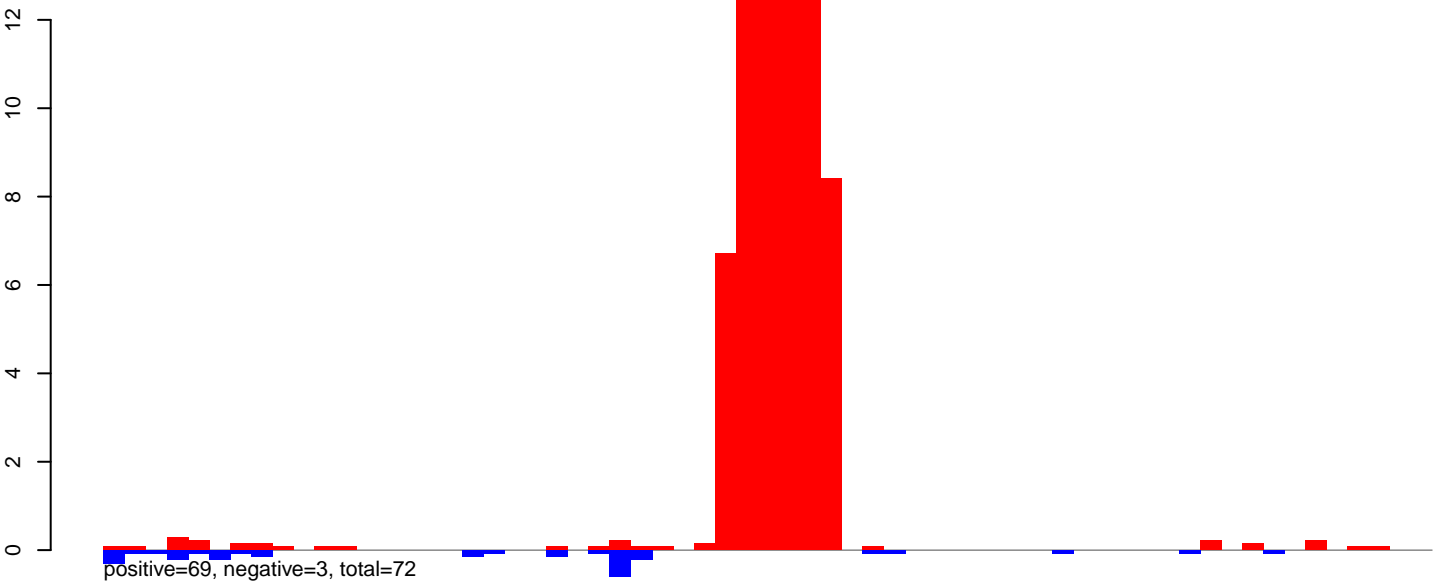


Window size=25, length=1561, TE@rnd-5_family-51@DNA_hAT@552:1-1561

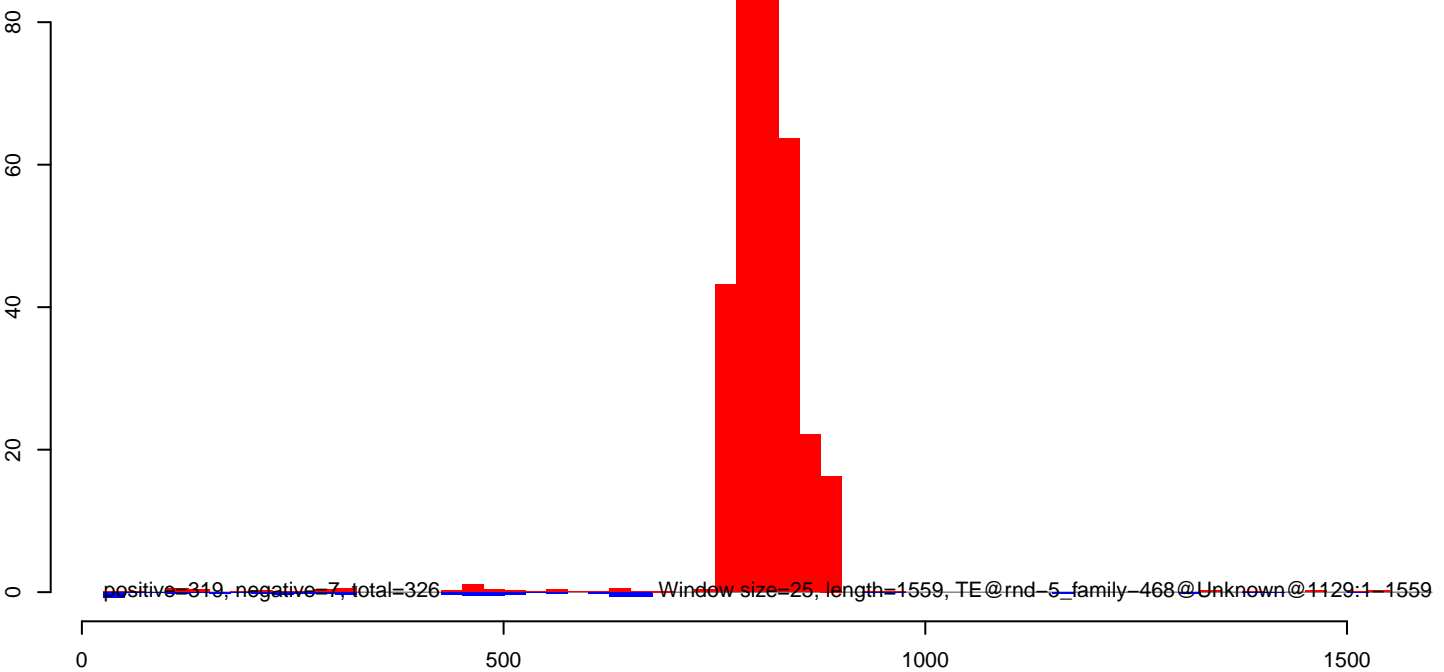
CuQuin_Testes_NL.18_23.rep



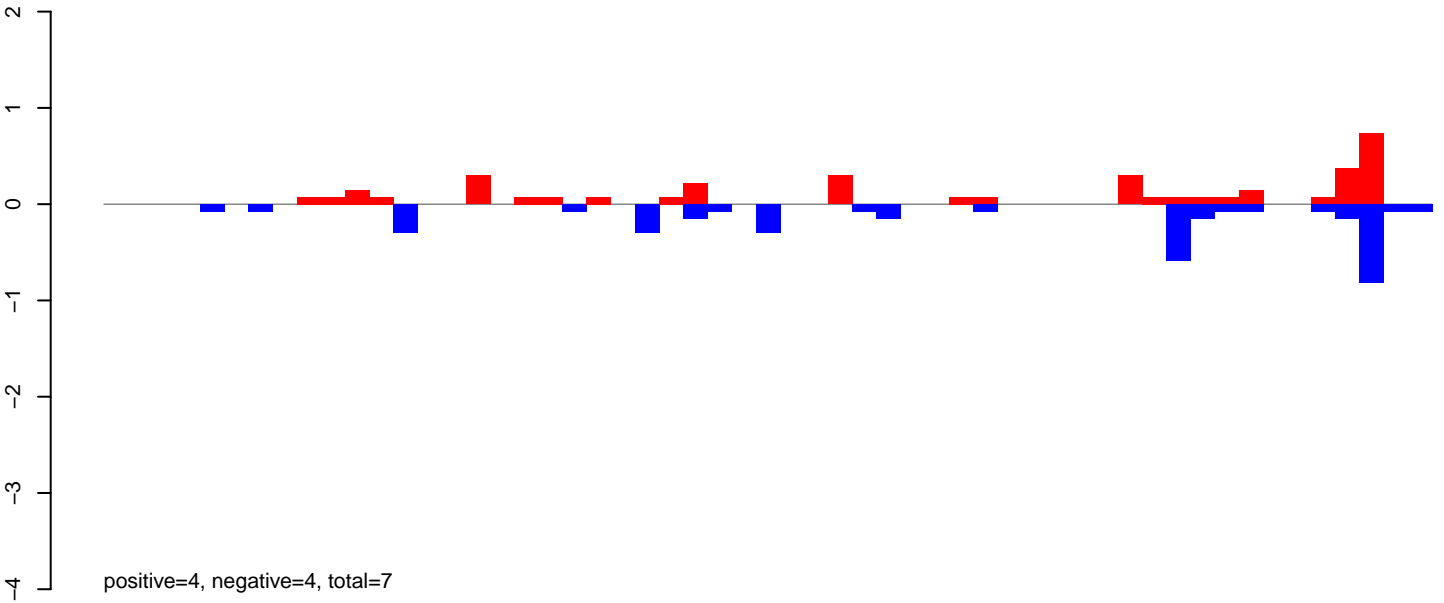
CuQuin_Testes_NL.24_35.rep



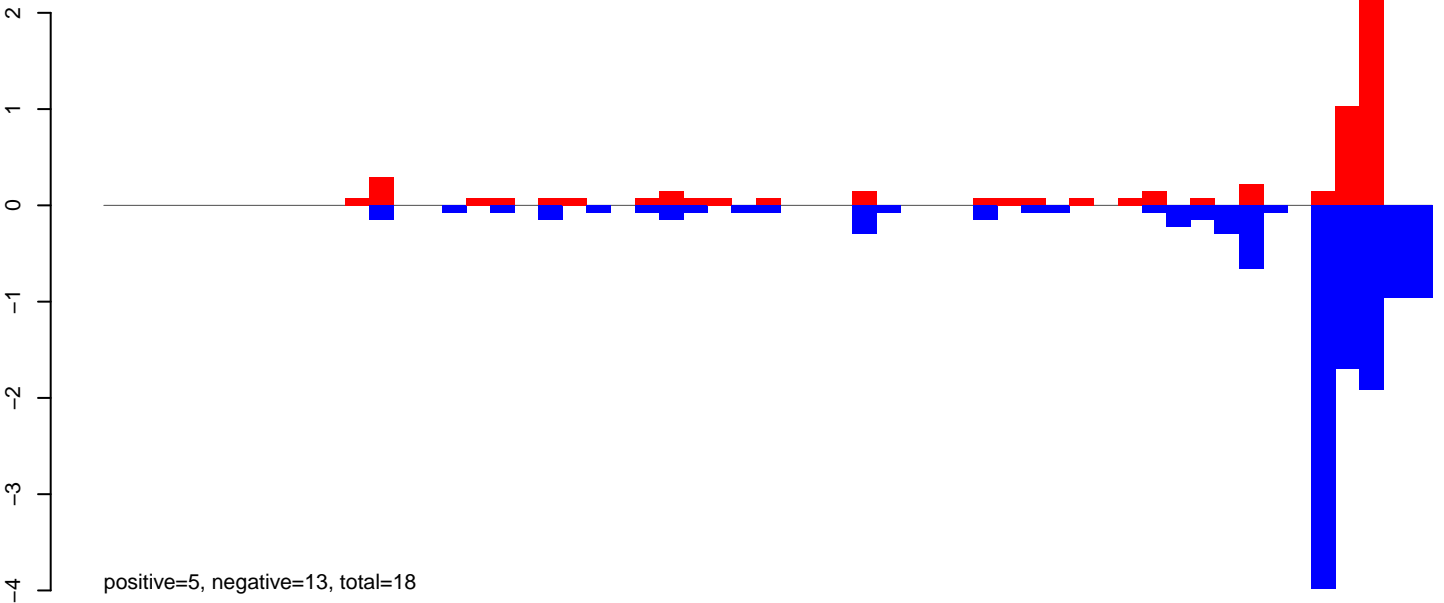
CuQuin_Testes_NL.rep



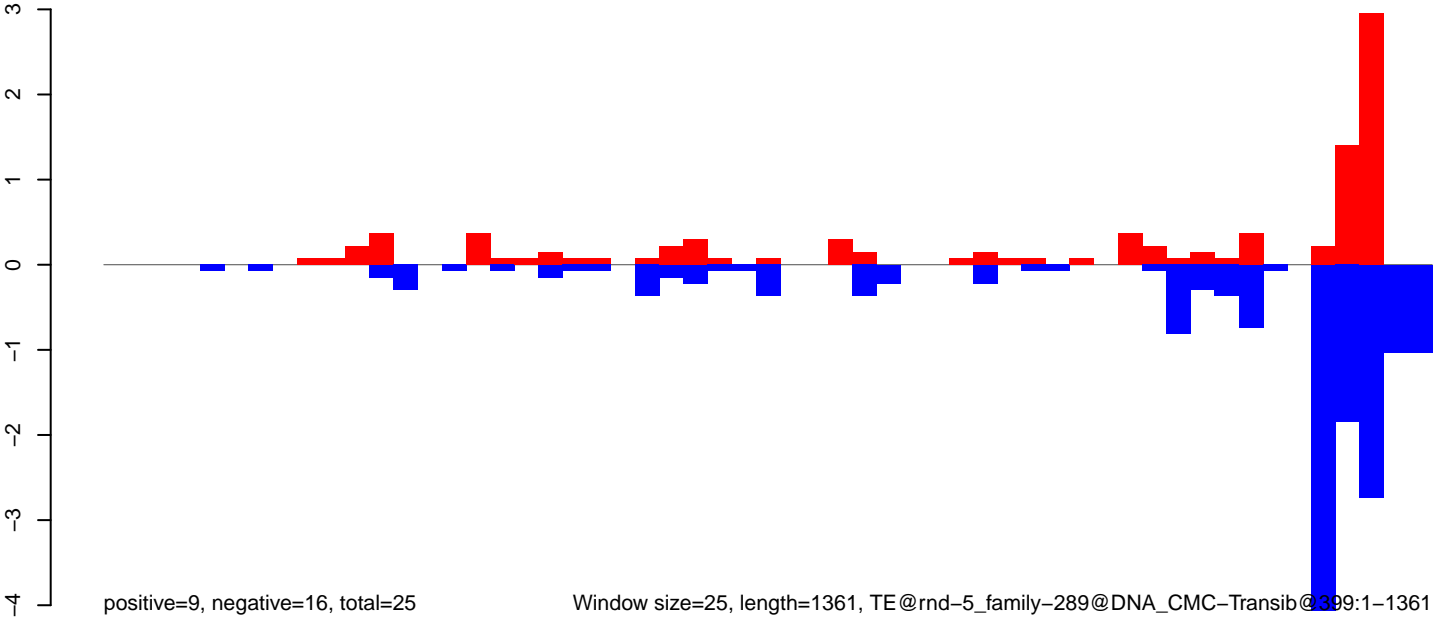
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

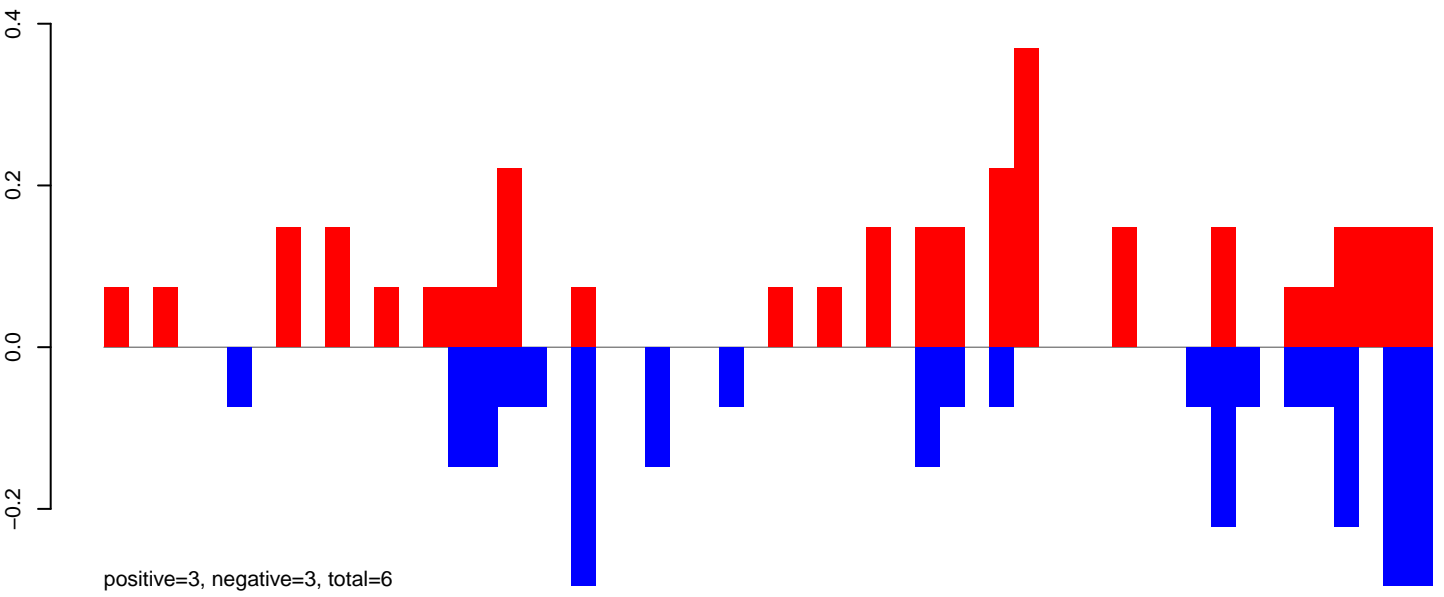


CuQuin_Testes_NL.rep

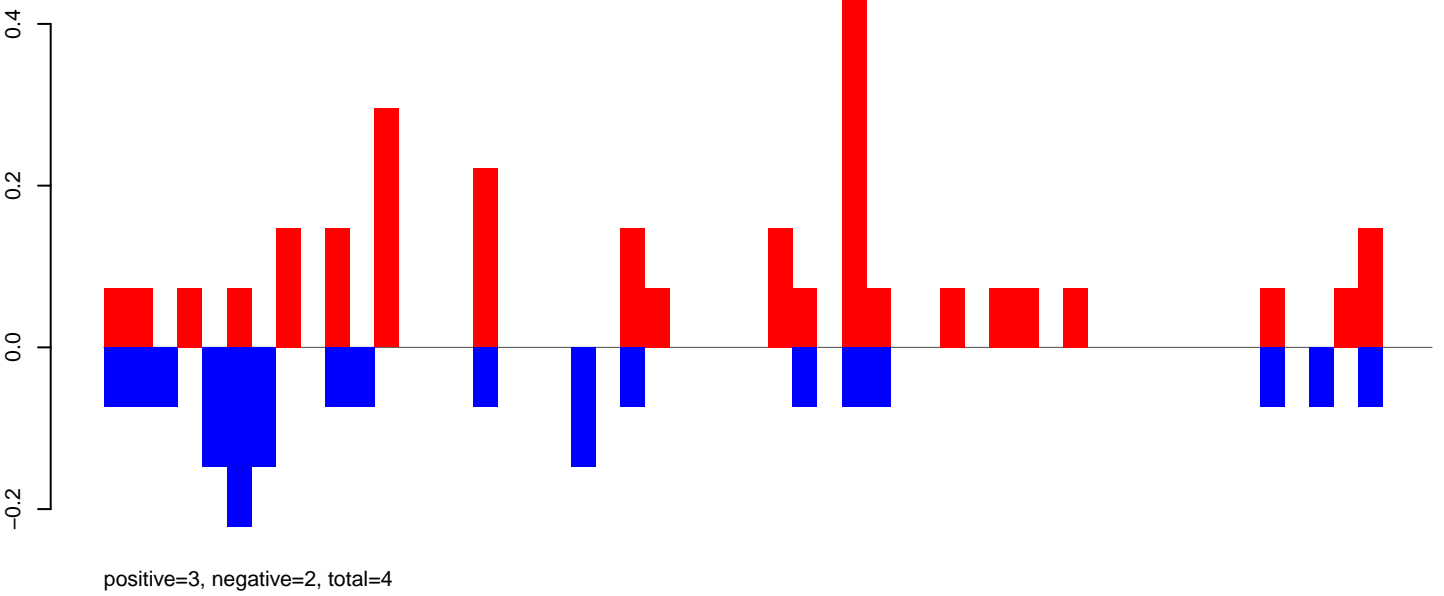


Window size=25, length=1361, TE@rnd-5_family-289@DNA_CMC-Transib@399:1-1361

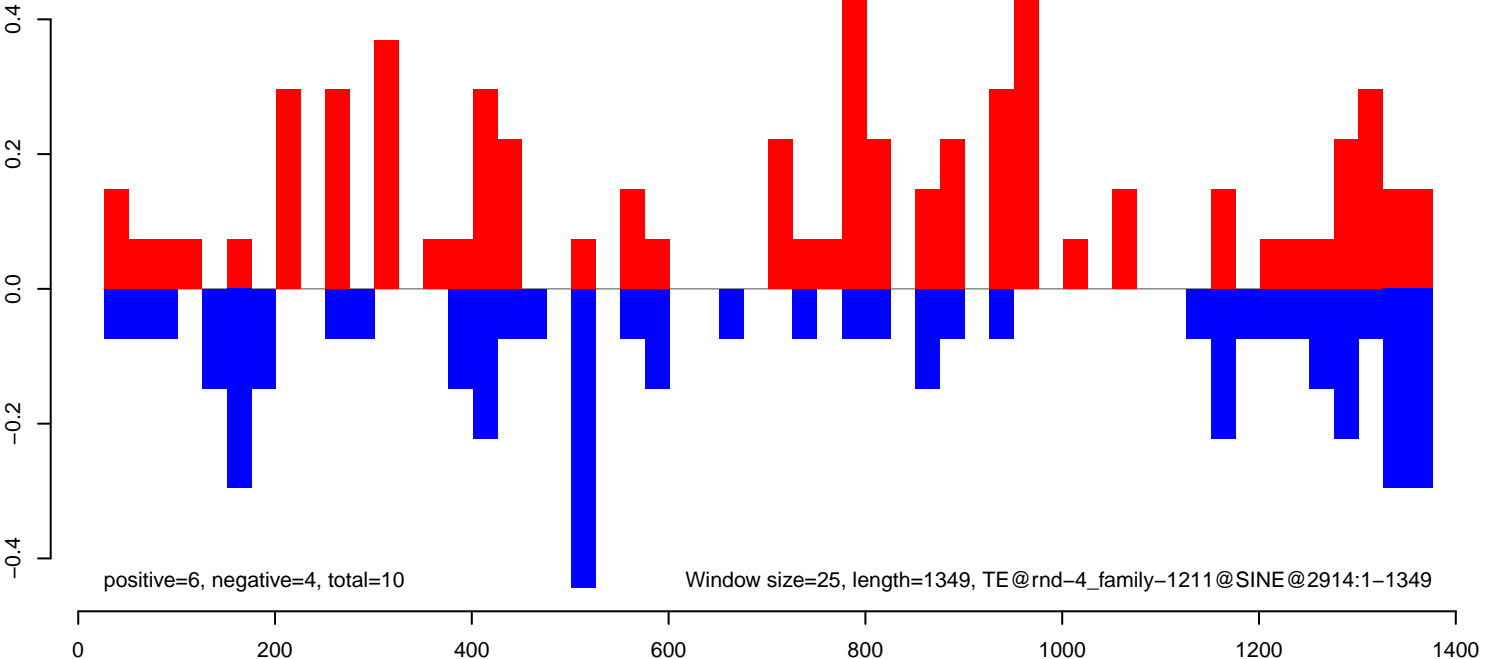
CuQuin_Testes_NL.18_23.rep



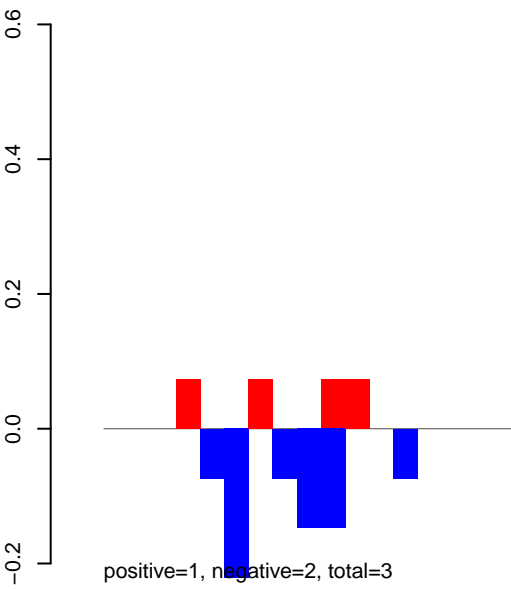
CuQuin_Testes_NL.24_35.rep



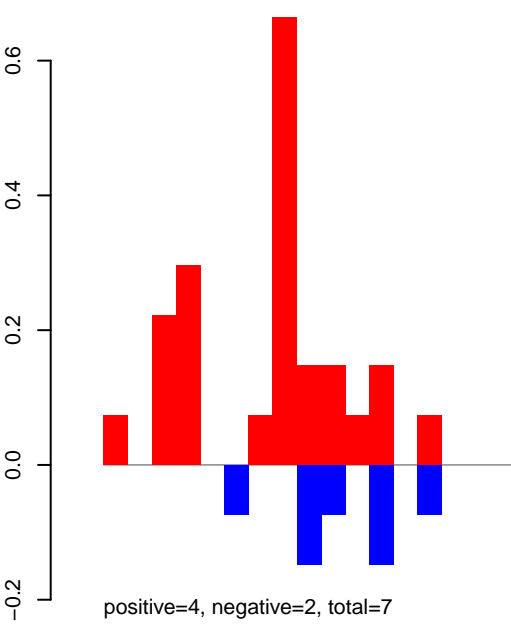
CuQuin_Testes_NL.rep



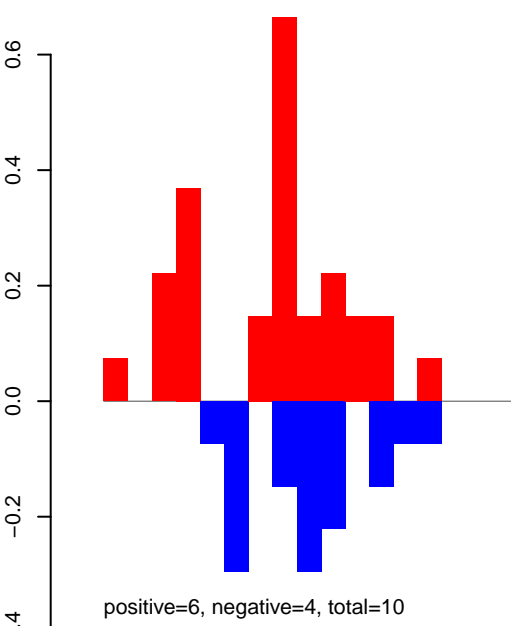
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

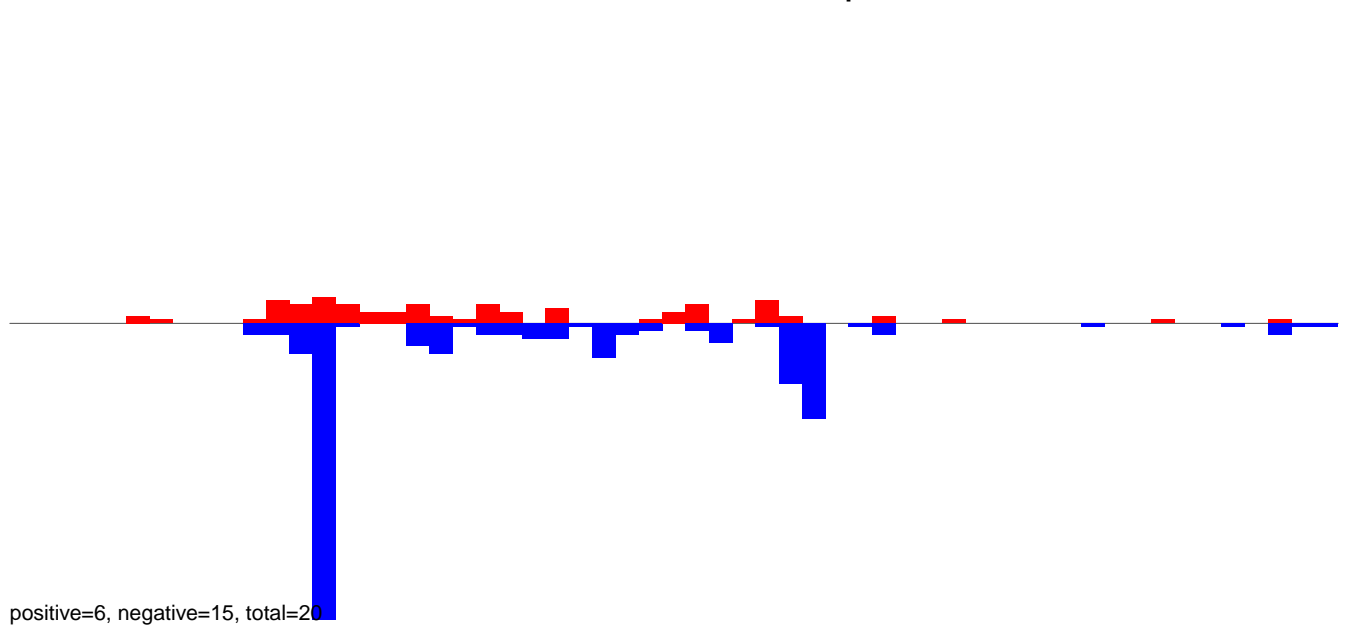


CuQuin_Testes_NL.rep

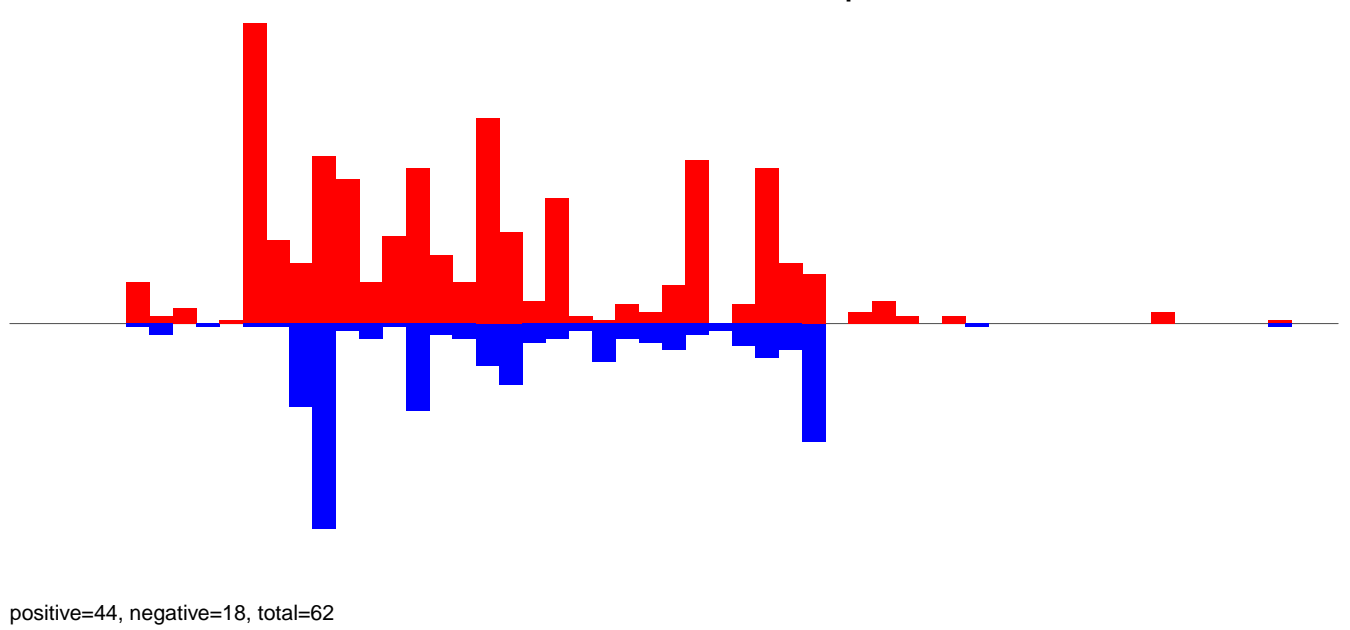


0 200 400 600 800 1000 1200 1400

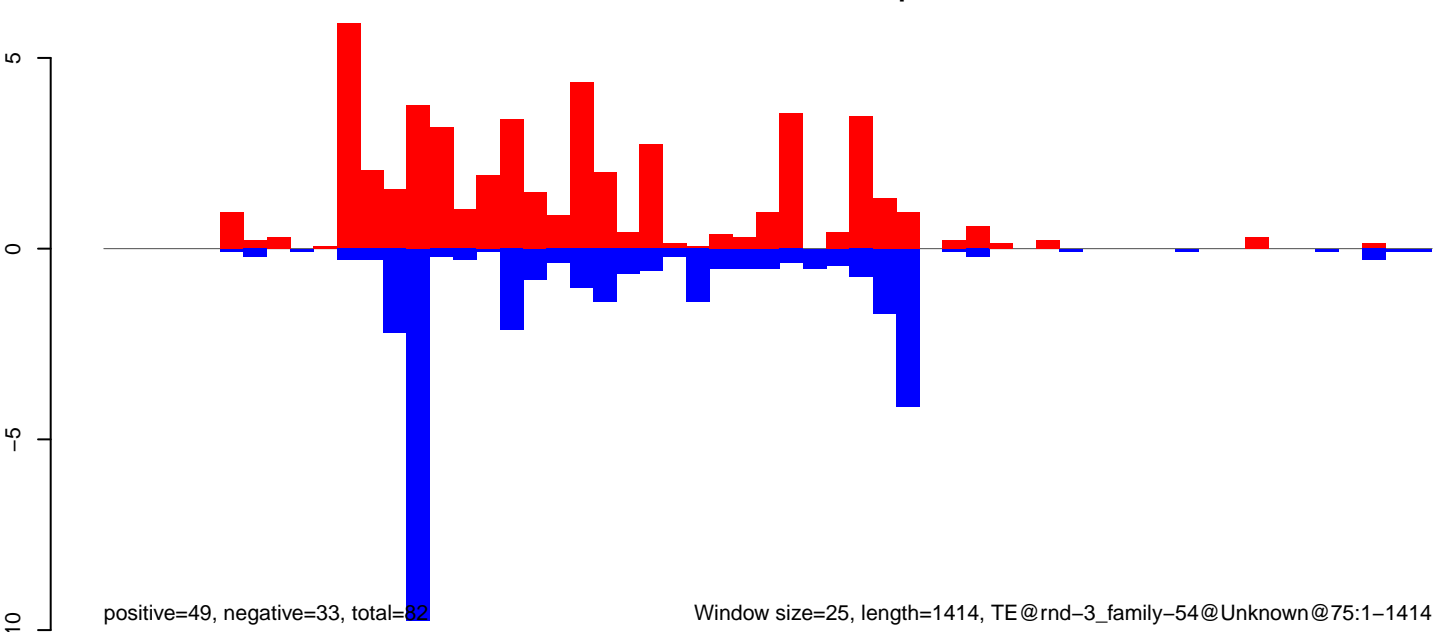
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

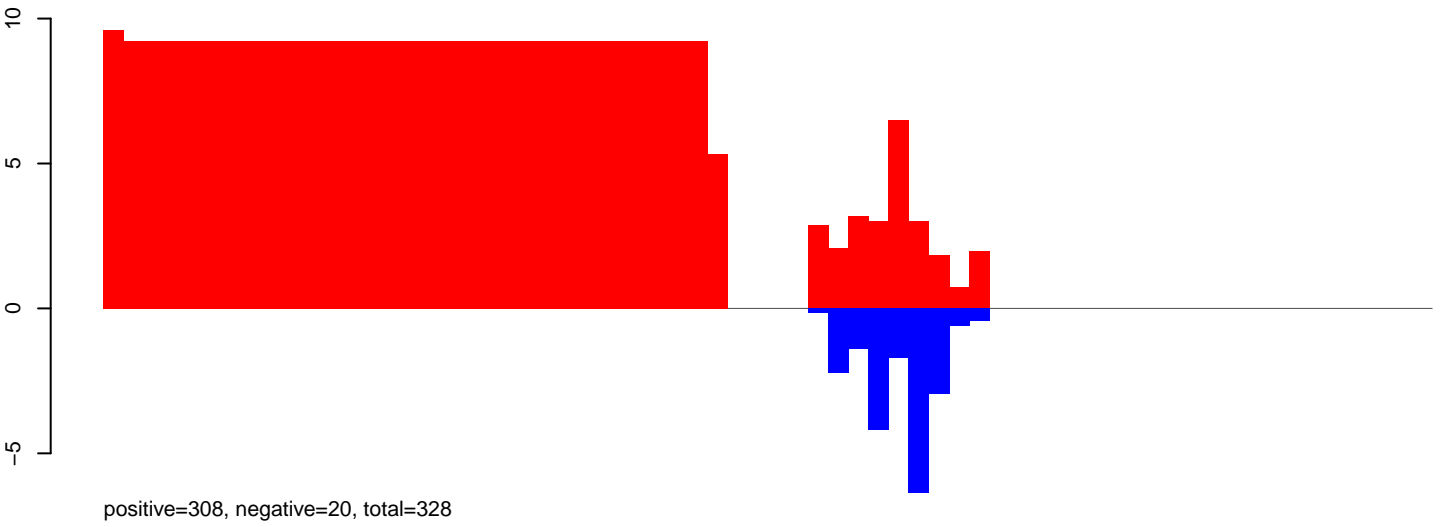


CuQuin_Testes_NL.rep

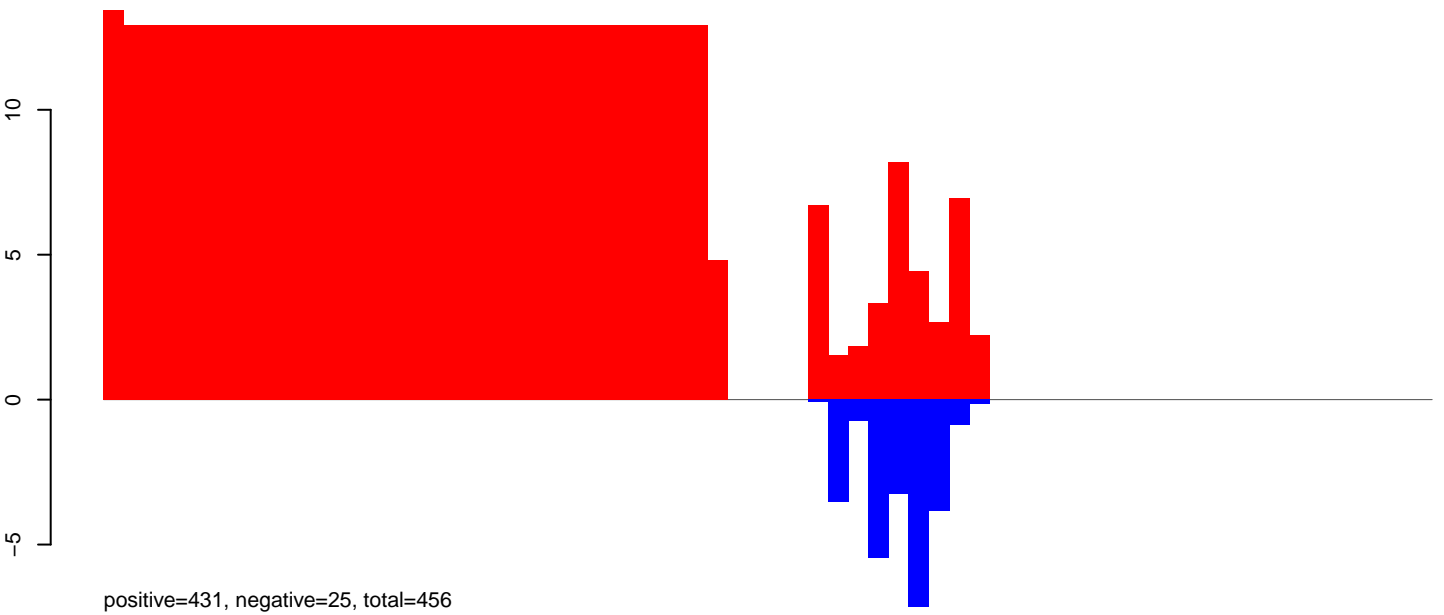


Window size=25, length=1414, TE@rnd-3_family-54@Unknown@75:1-1414

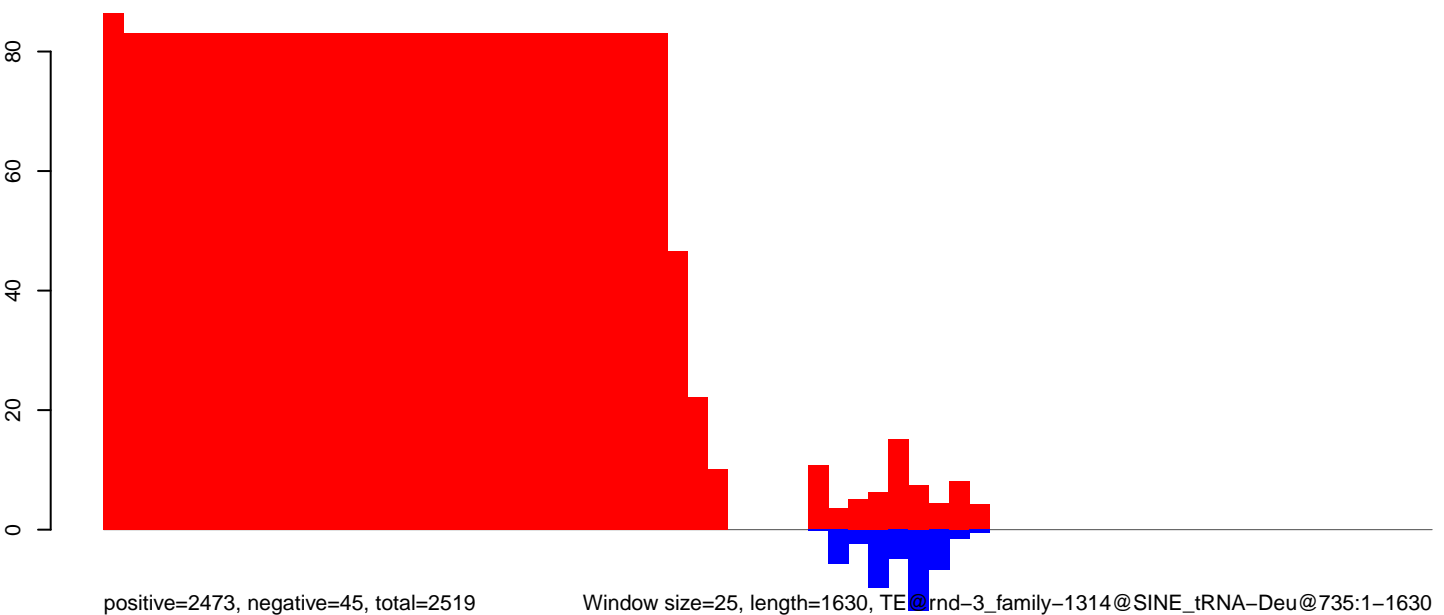
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



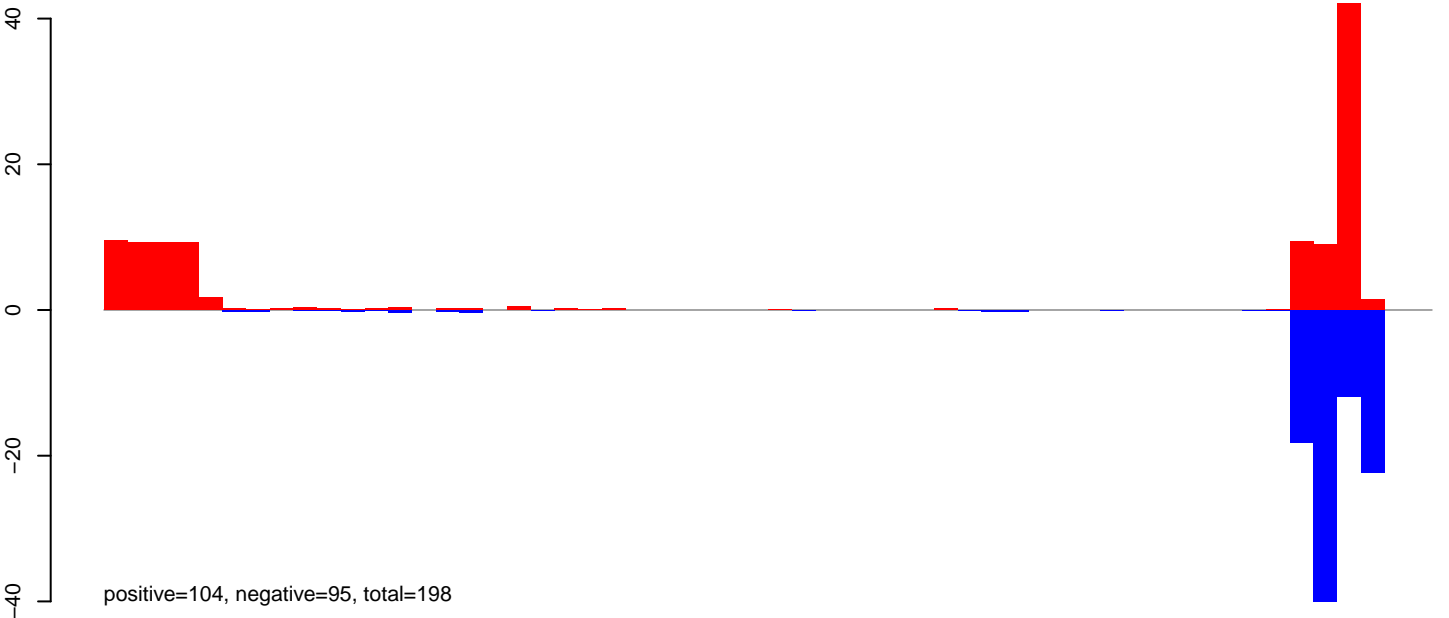
CuQuin_Testes_NL.rep



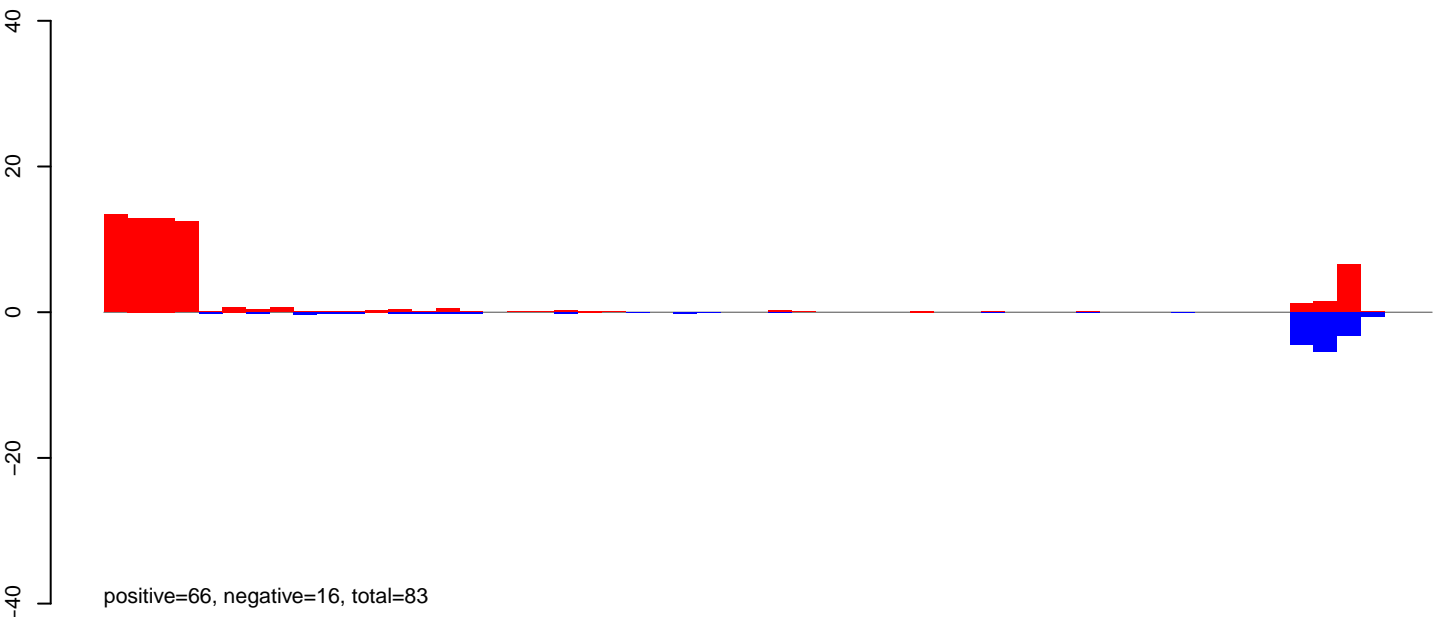
Window size=25, length=1630, TE@rnd-3_family-1314@SINE_tRNA-Deu@735:1-1630

0 500 1000 1500

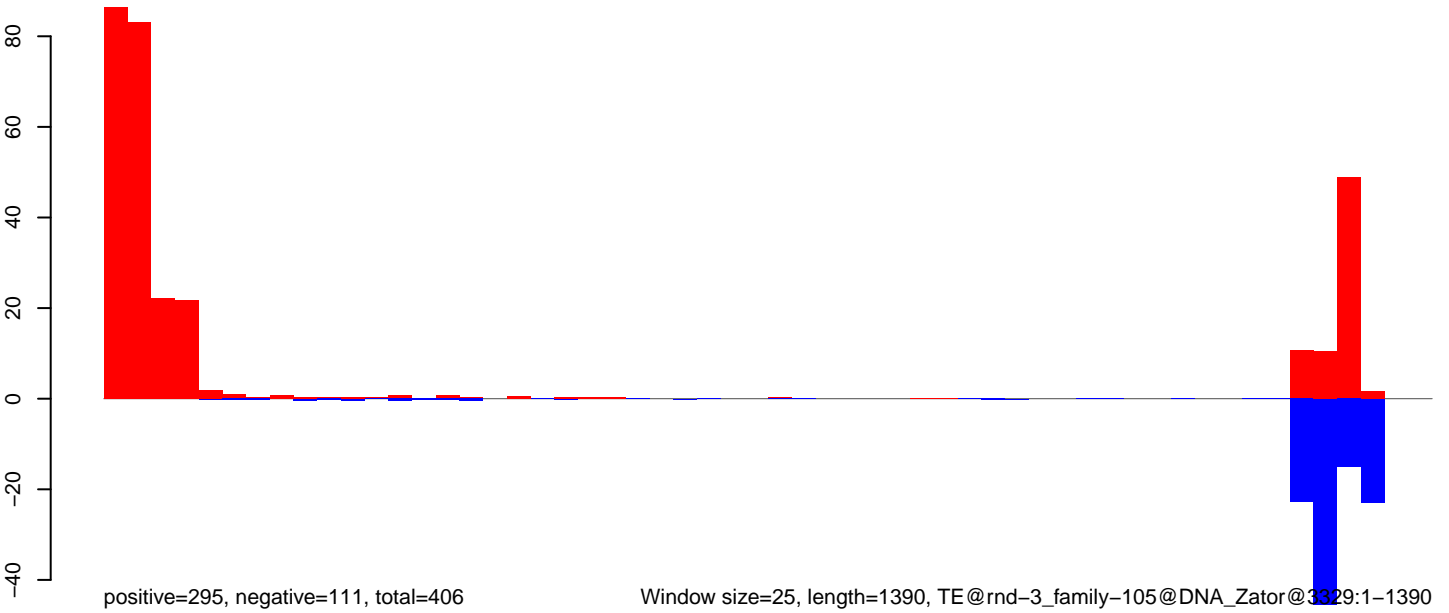
CuQuin_Testes_NL.18_23.rep



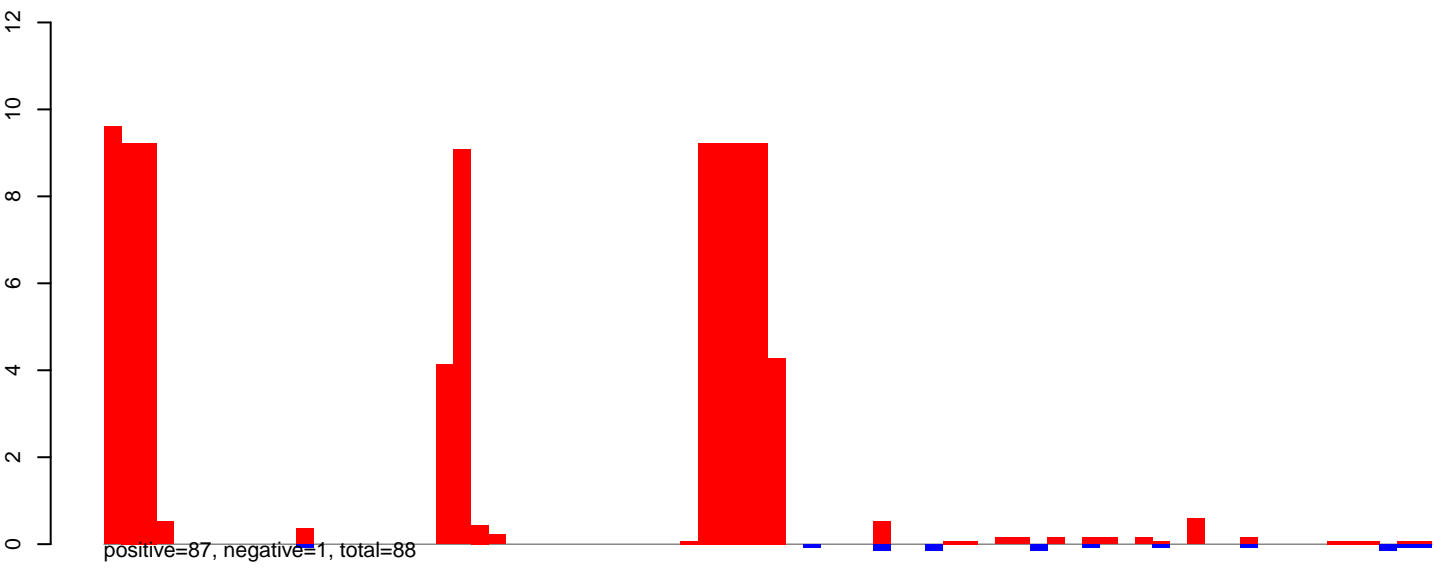
CuQuin_Testes_NL.24_35.rep



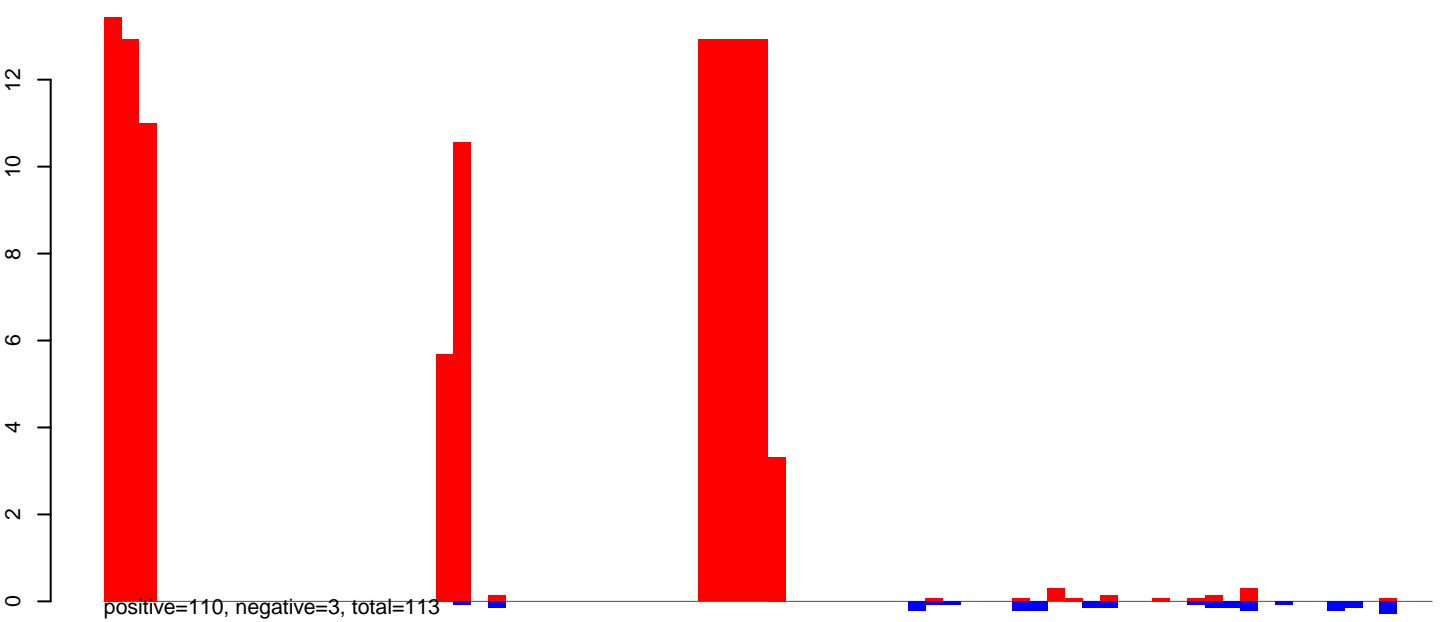
CuQuin_Testes_NL.rep



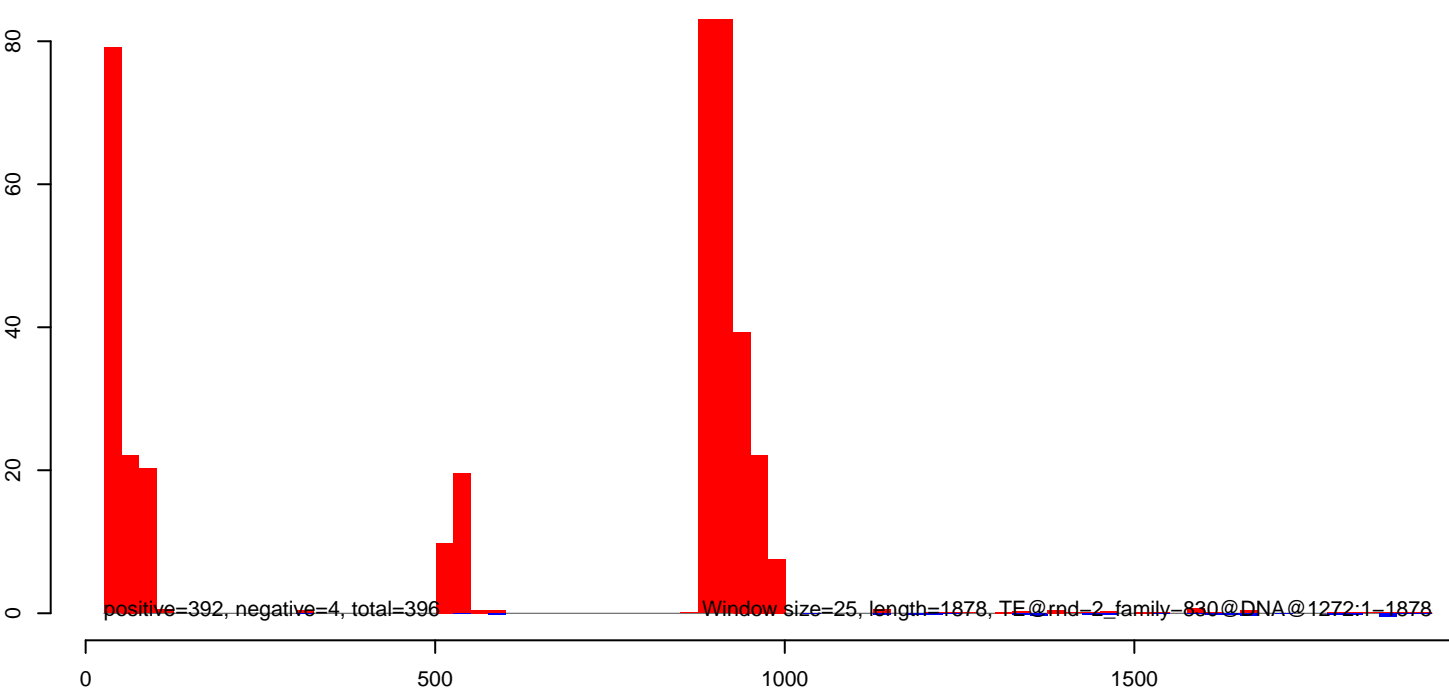
CuQuin_Testes_NL.18_23.rep



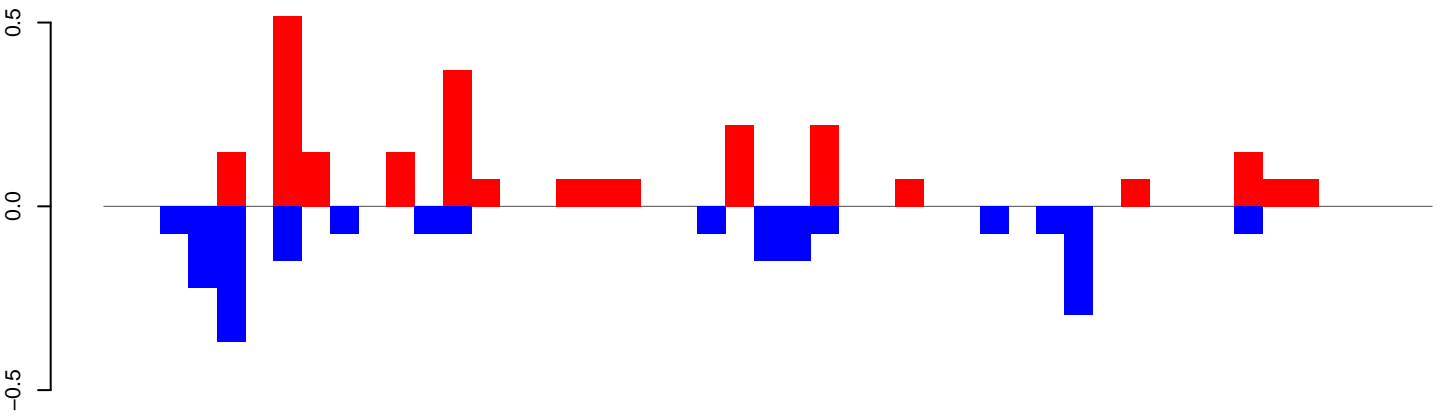
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

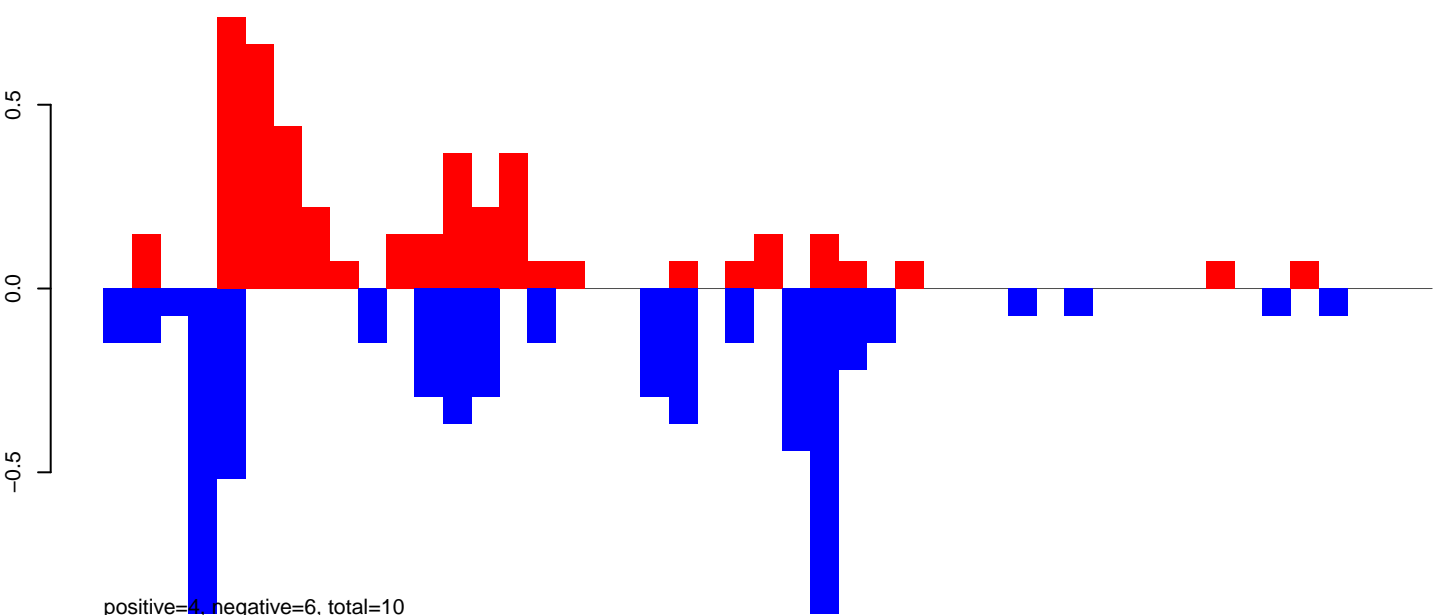


CuQuin_Testes_NL.18_23.rep



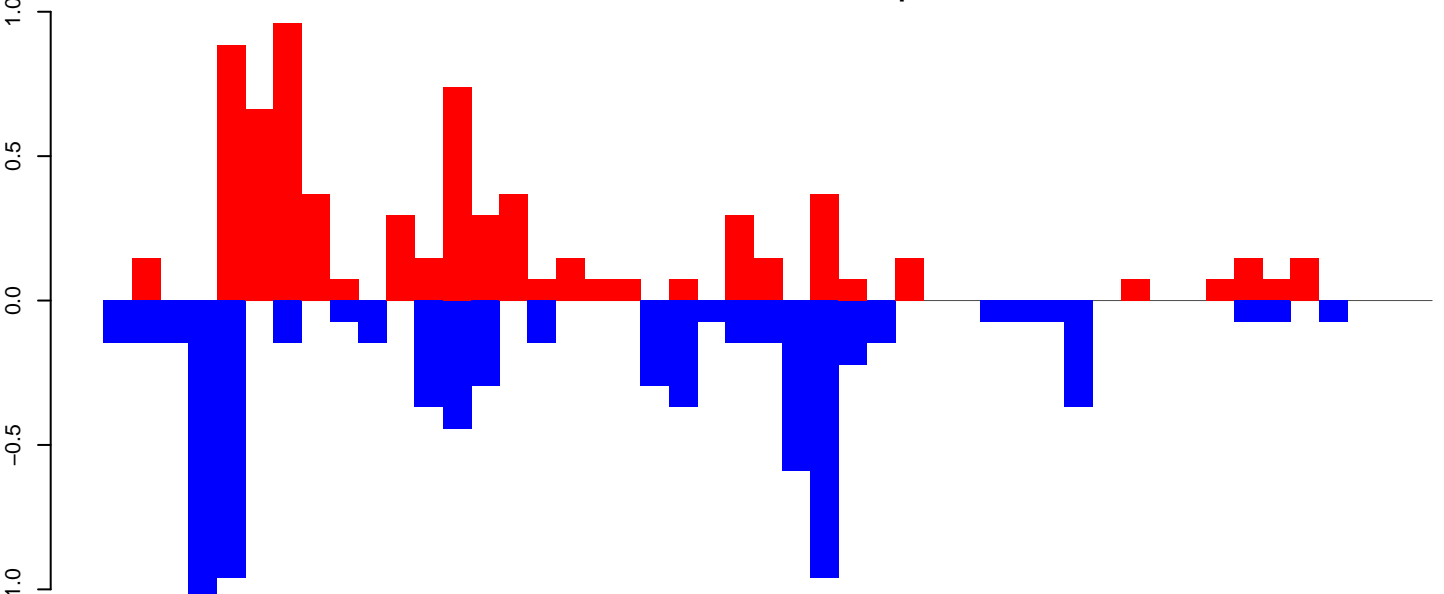
positive=3, negative=2, total=5

CuQuin_Testes_NL.24_35.rep



positive=4, negative=6, total=10

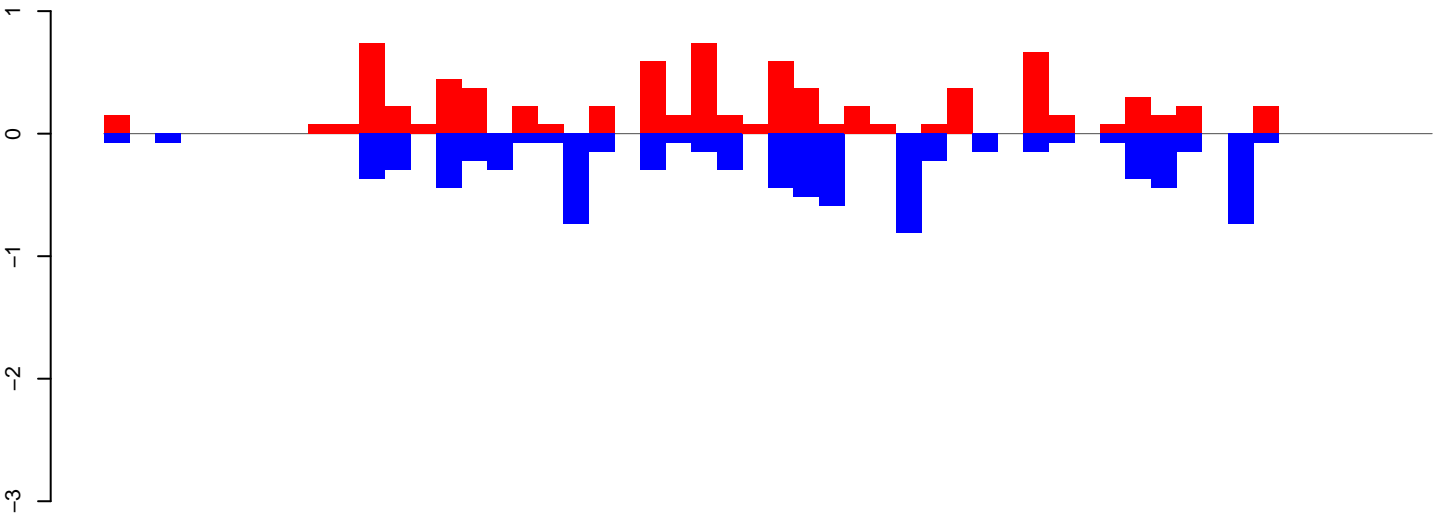
CuQuin_Testes_NL.rep



positive=7, negative=8, total=15

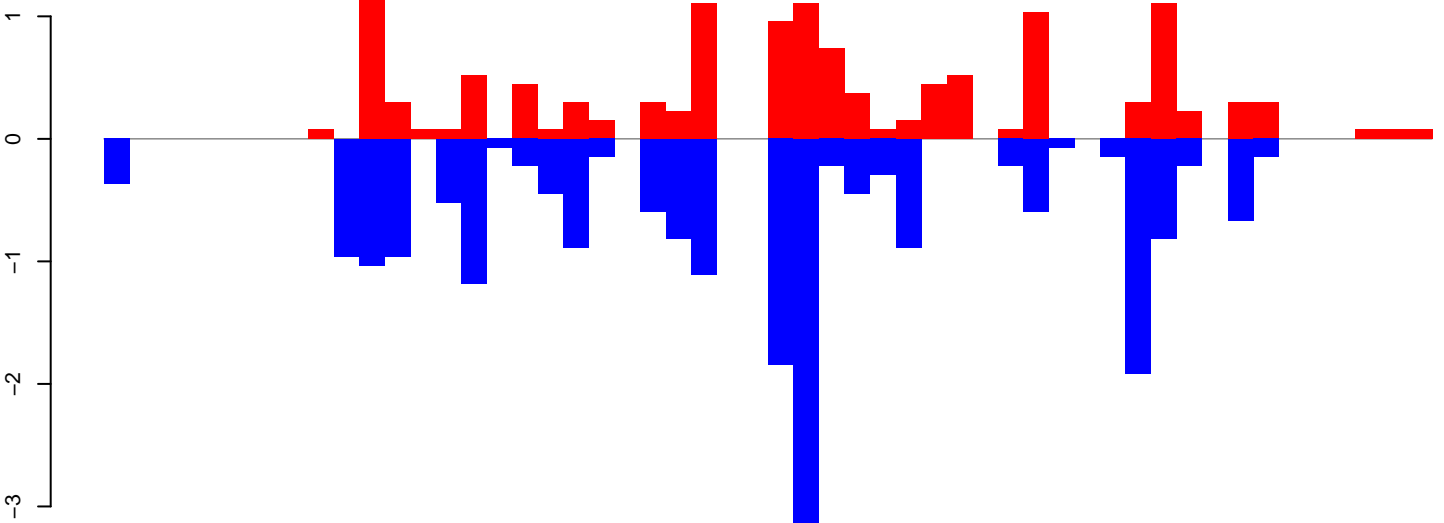
Window size=25, length=1154, TE@rnd-5_family-2356@LTR_Gypsy@2972:1-1154

CuQuin_Testes_NL.18_23.rep



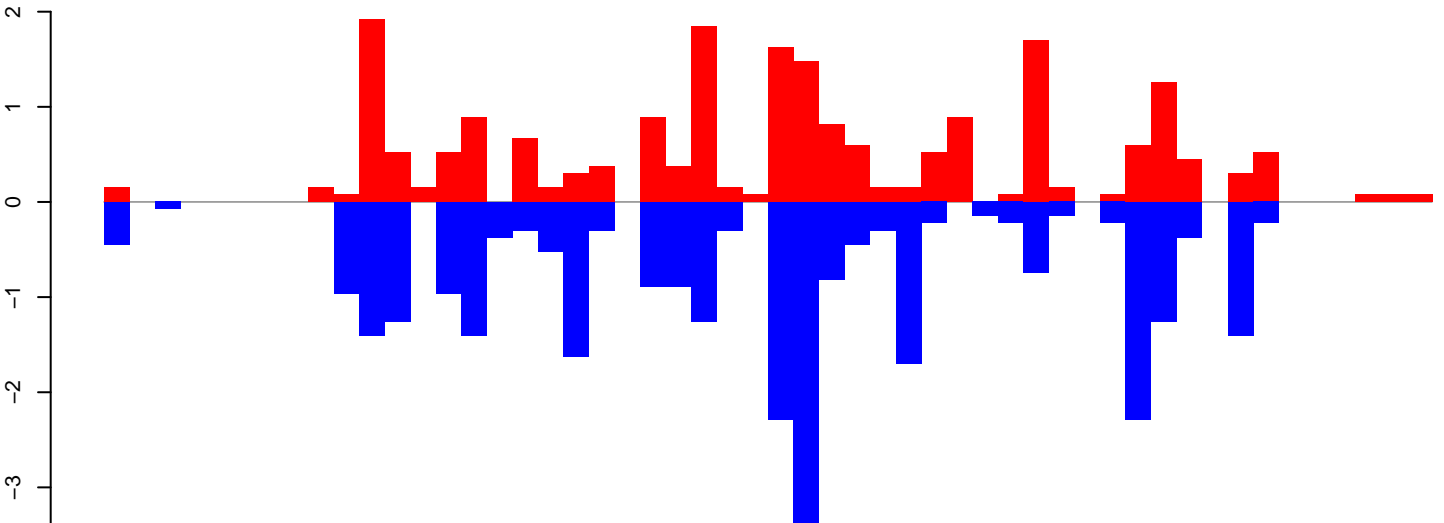
positive=8, negative=8, total=16

CuQuin_Testes_NL.24_35.rep



positive=13, negative=21, total=34

CuQuin_Testes_NL.rep

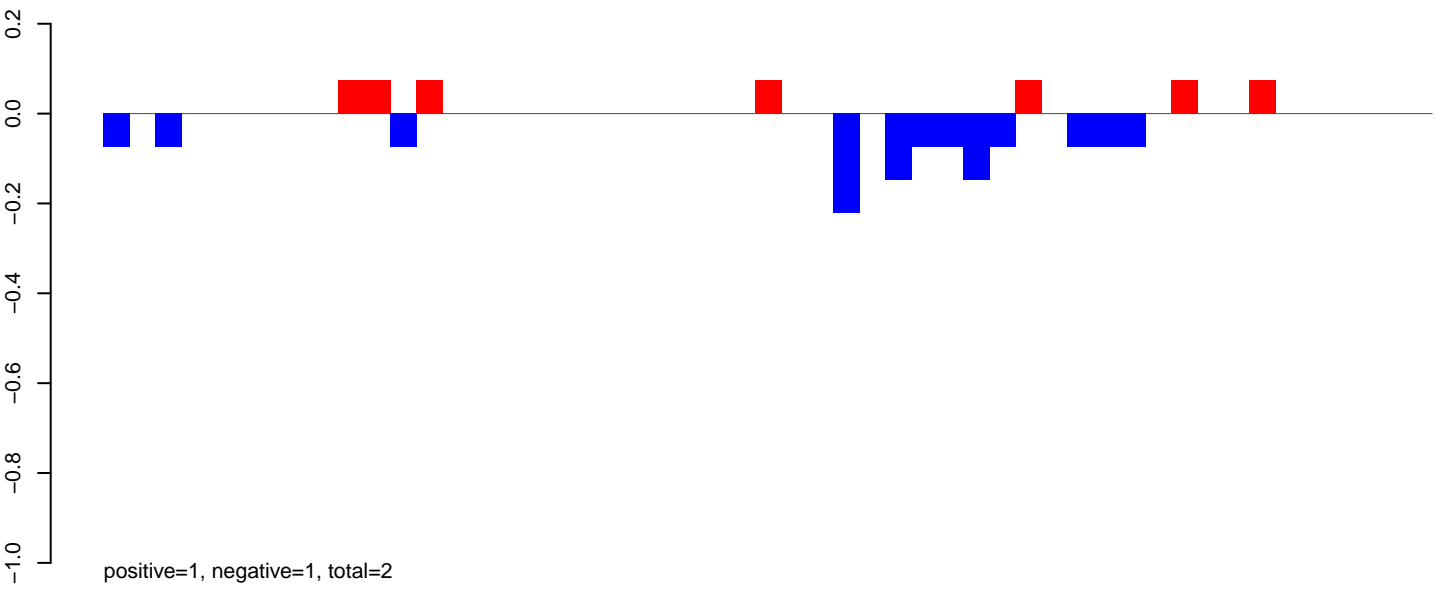


positive=21, negative=30, total=51

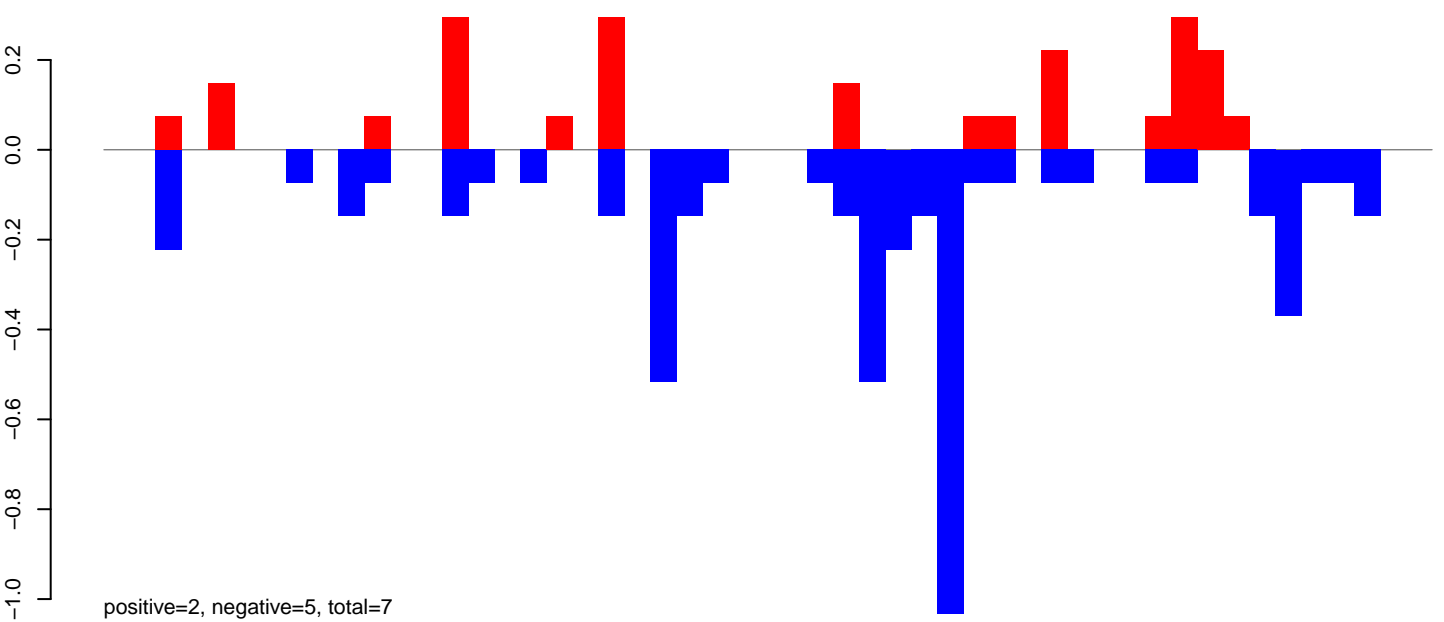
Window size=25, length=1295, TE@rnd-5_family-188@DNA@555:1-1295

0 200 400 600 800 1000 1200

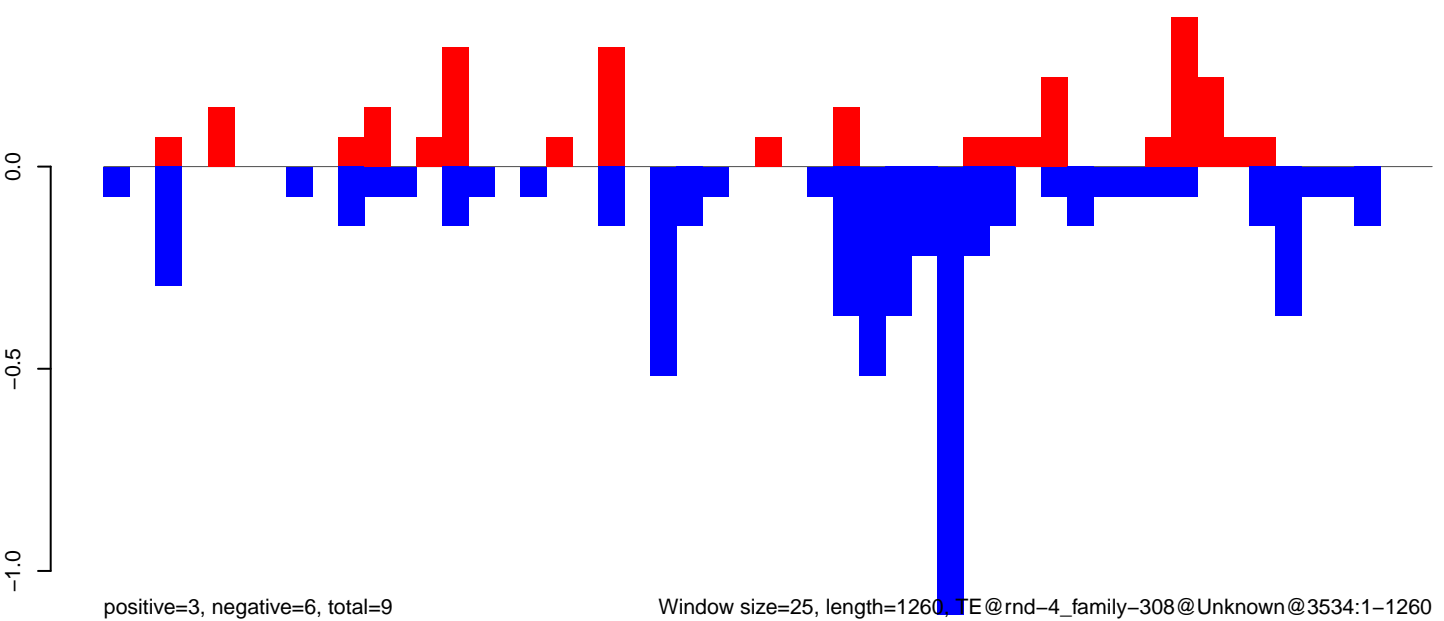
CuQuin_Testes_NL.18_23.rep



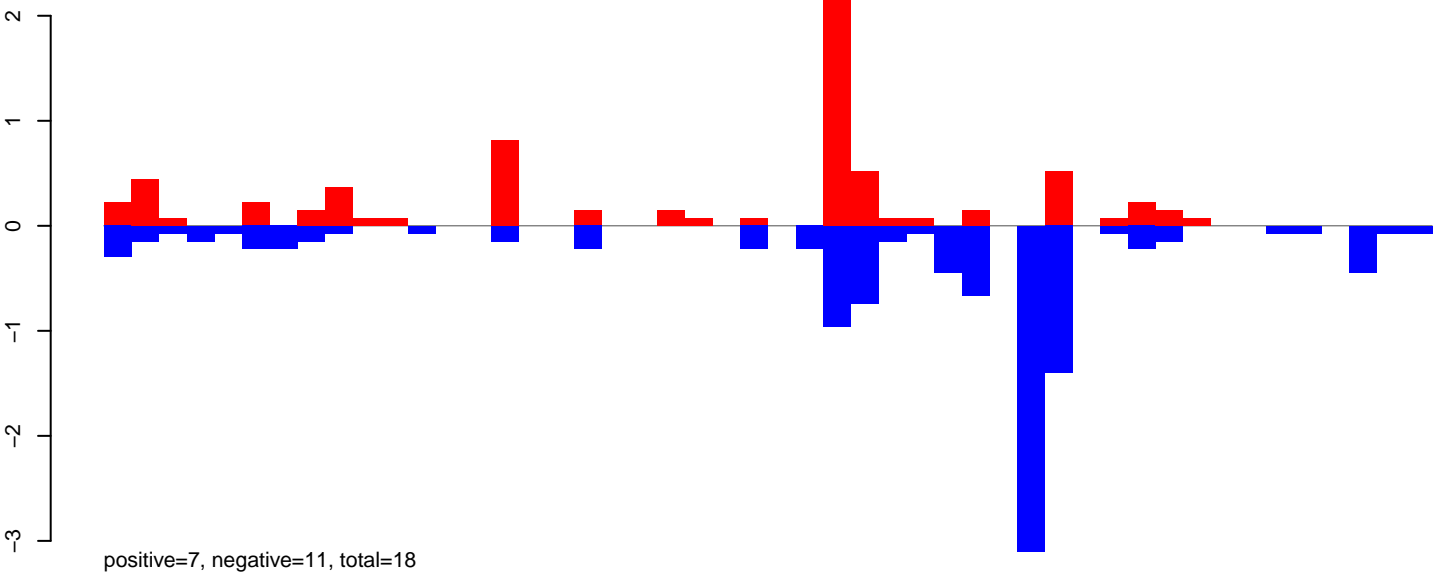
CuQuin_Testes_NL.24_35.rep



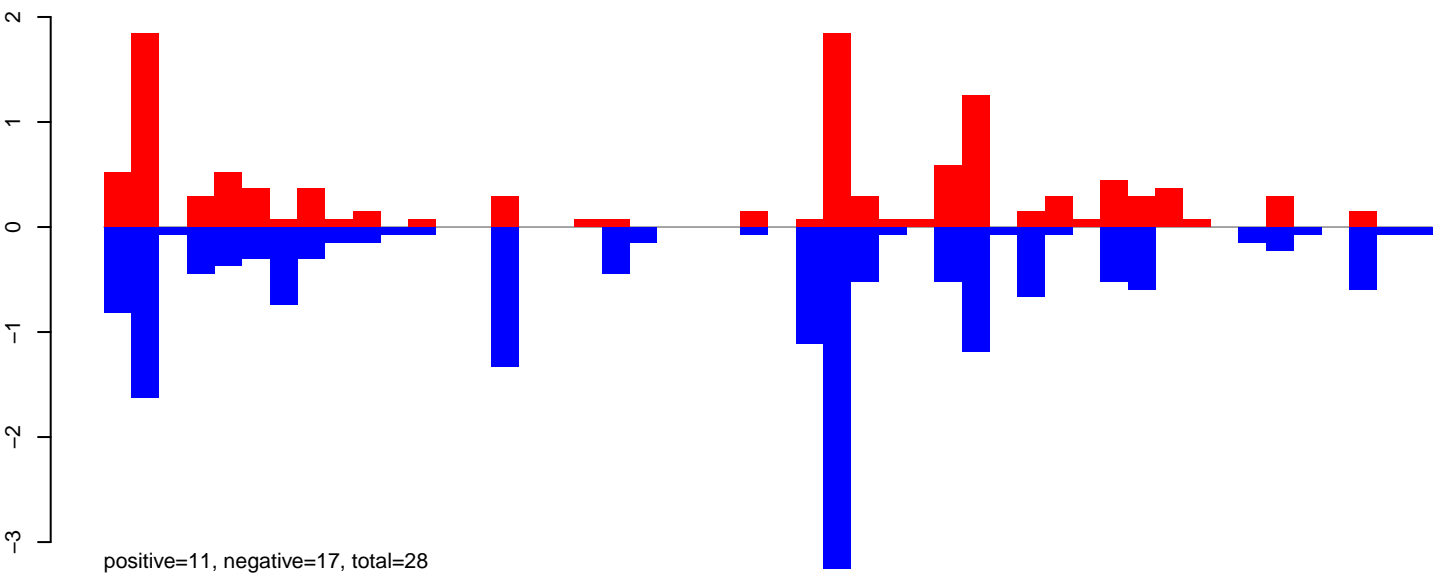
CuQuin_Testes_NL.rep



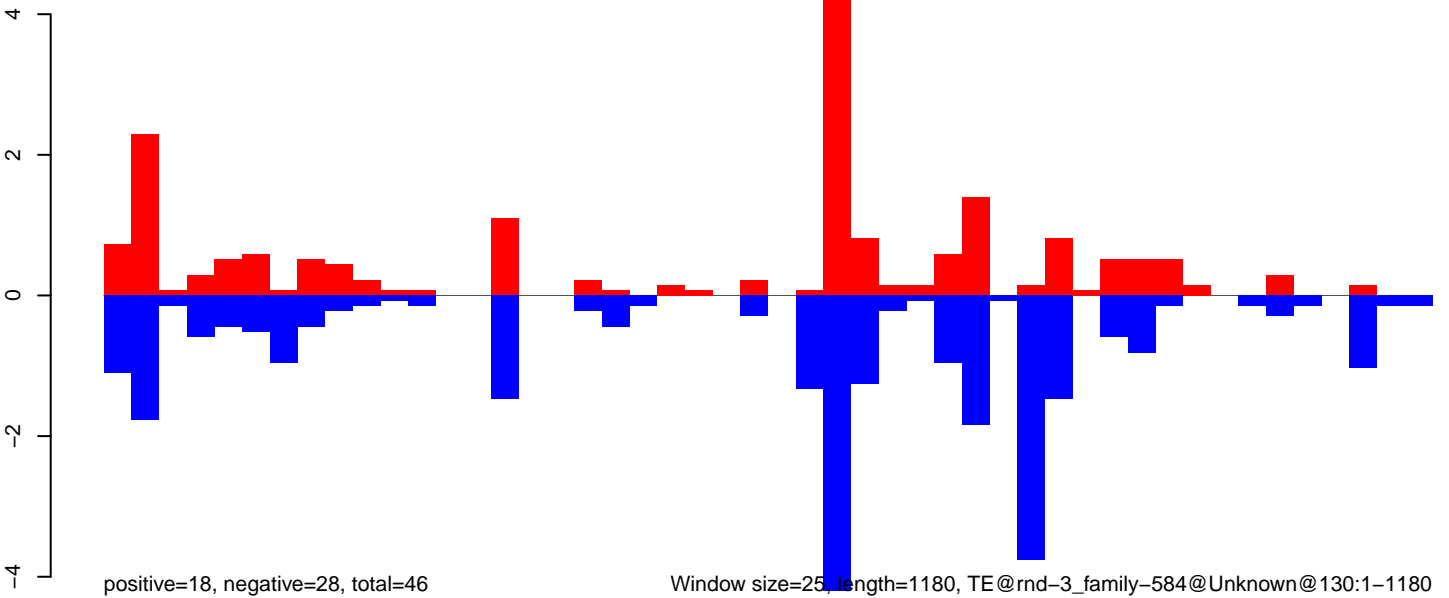
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

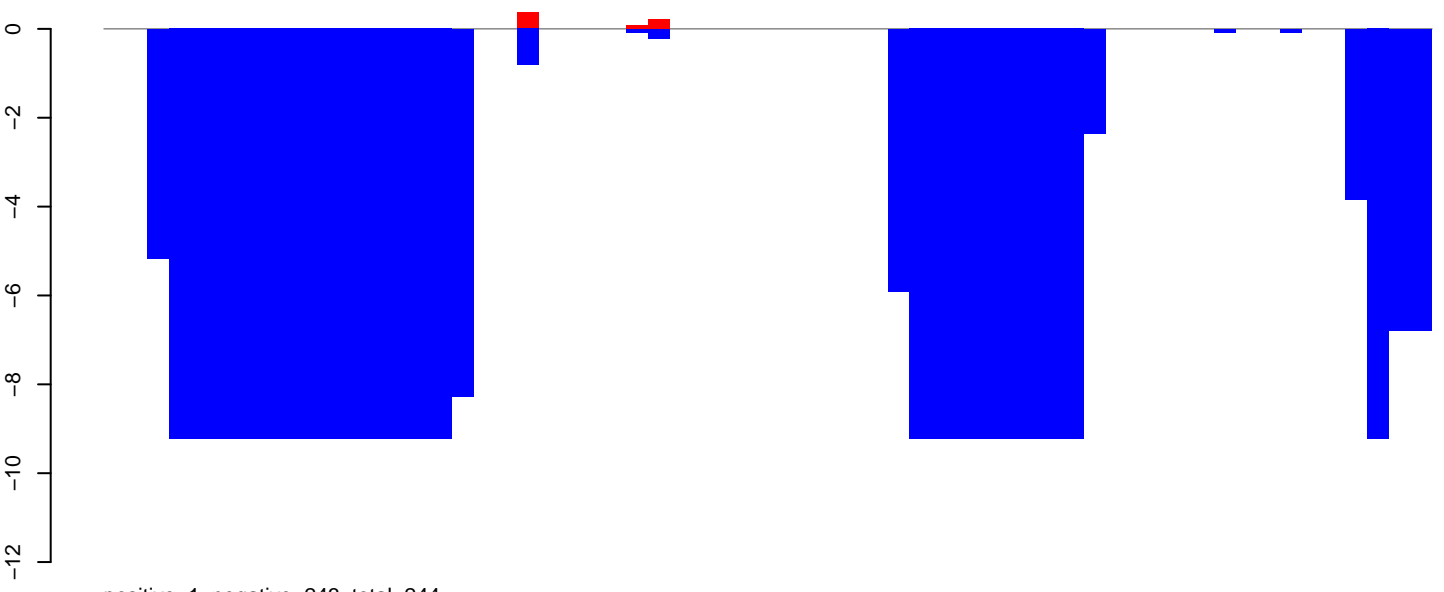


CuQuin_Testes_NL.rep



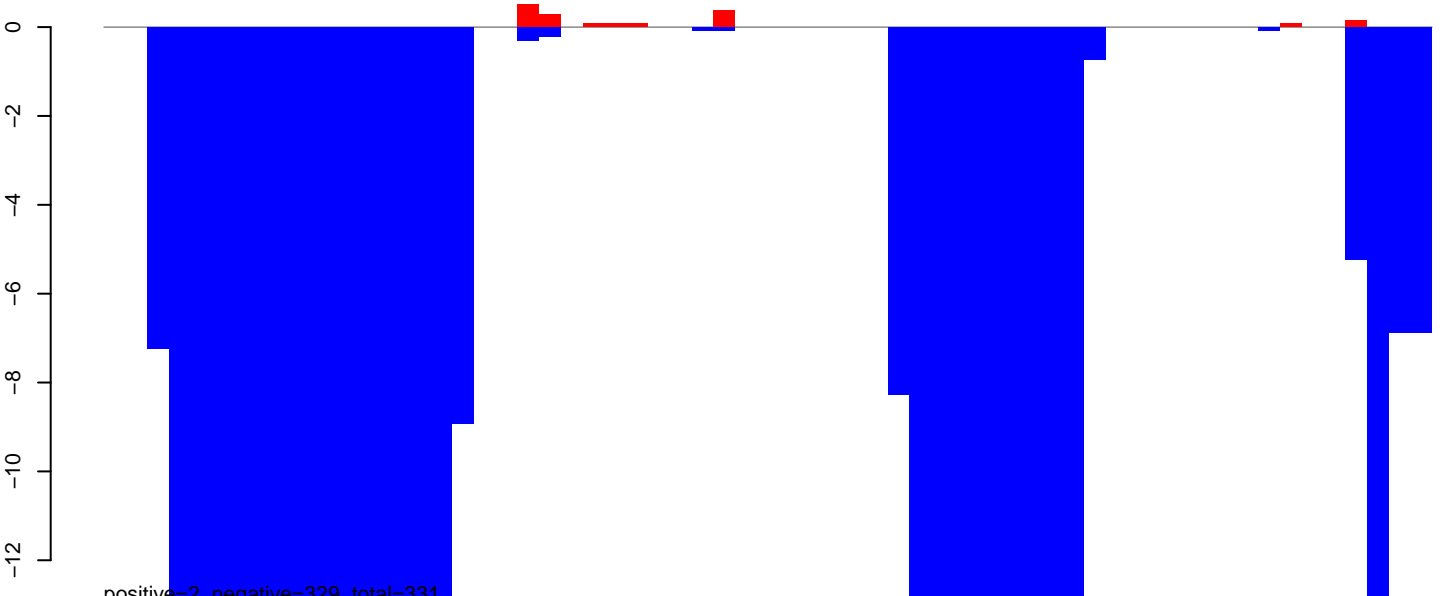
Window size=25, length=1180, TE@rnd-3_family-584@Unknown@130:1-1180

CuQuin_Testes_NL.18_23.rep



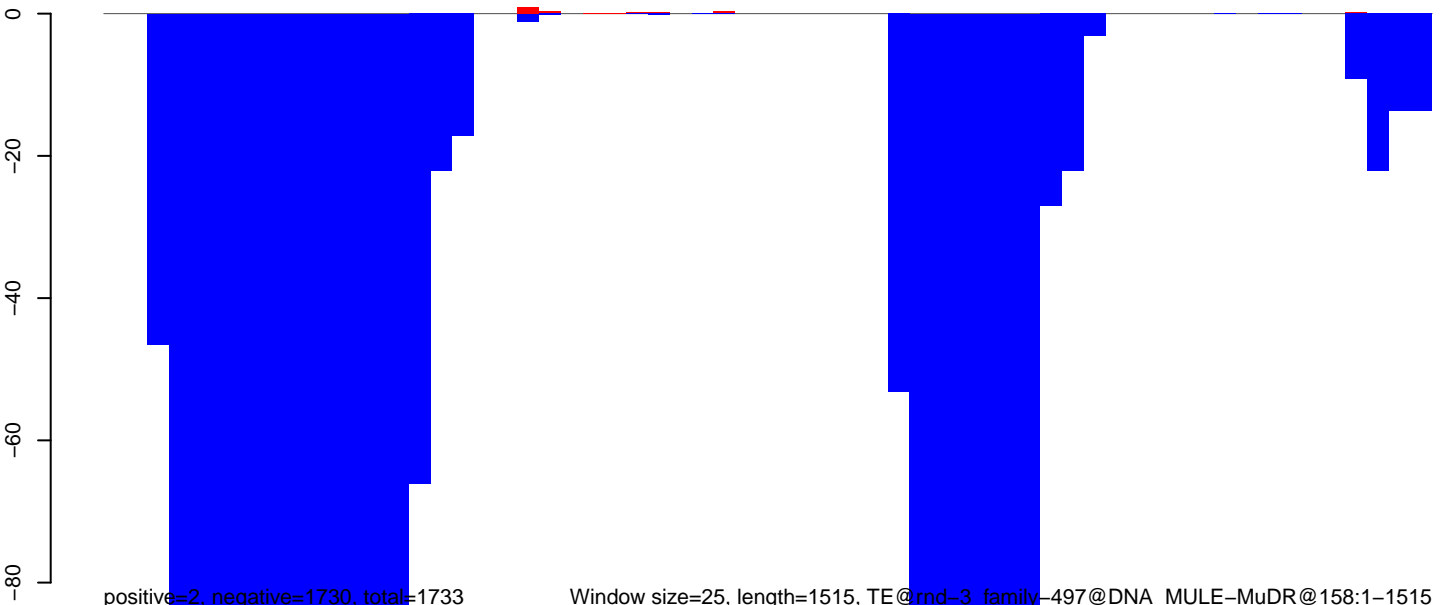
positive=1, negative=243, total=244

CuQuin_Testes_NL.24_35.rep



positive=2, negative=329, total=331

CuQuin_Testes_NL.rep

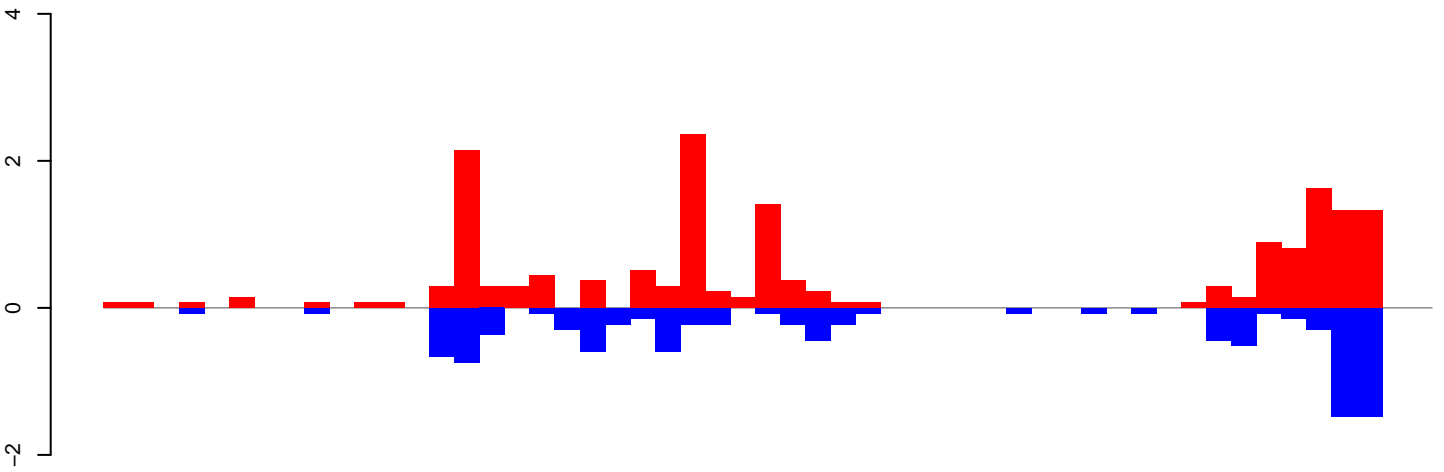


positive=2, negative=1730, total=1732

Window size=25, length=1515, TE@rnd-3_family-497@DNA_MULE-MuDR@158:1-1515

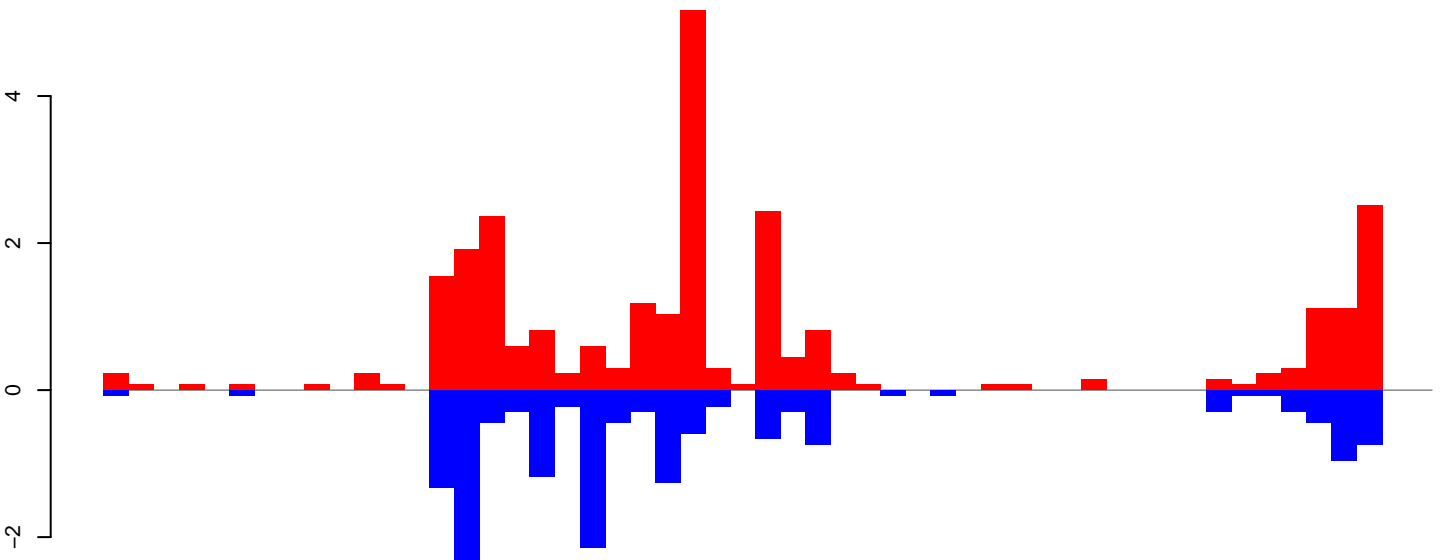
0 500 1000 1500

CuQuin_Testes_NL.18_23.rep



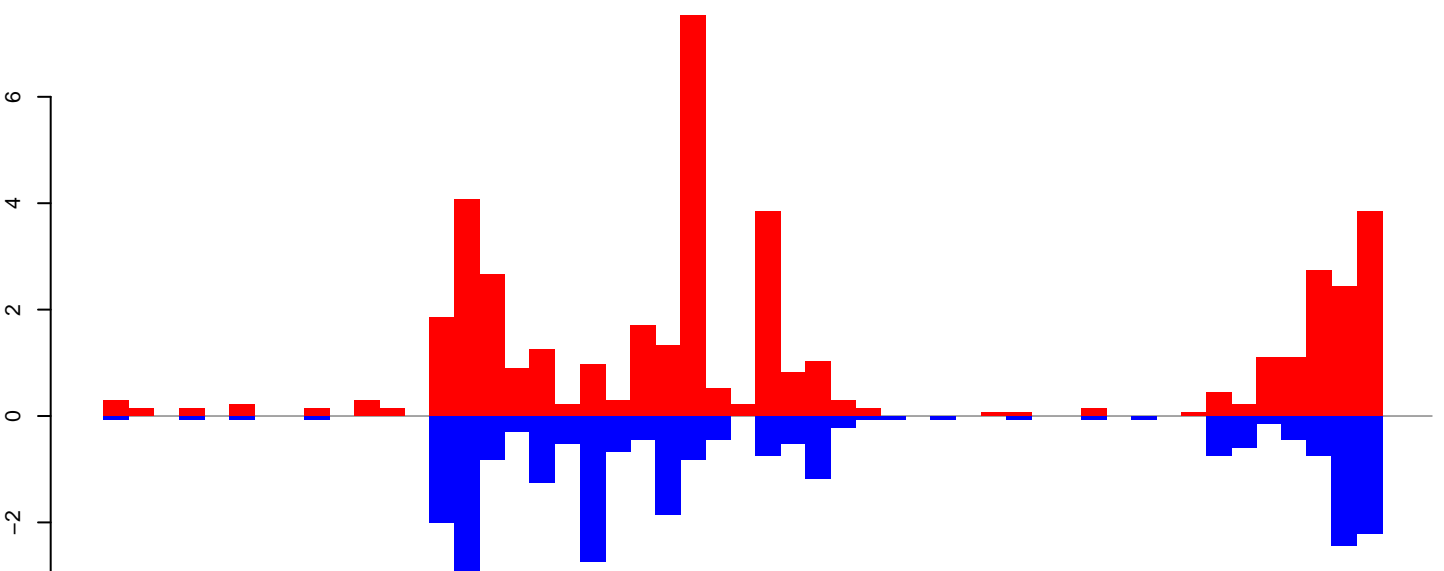
positive=17, negative=10, total=27

CuQuin_Testes_NL.24_35.rep



positive=27, negative=16, total=43

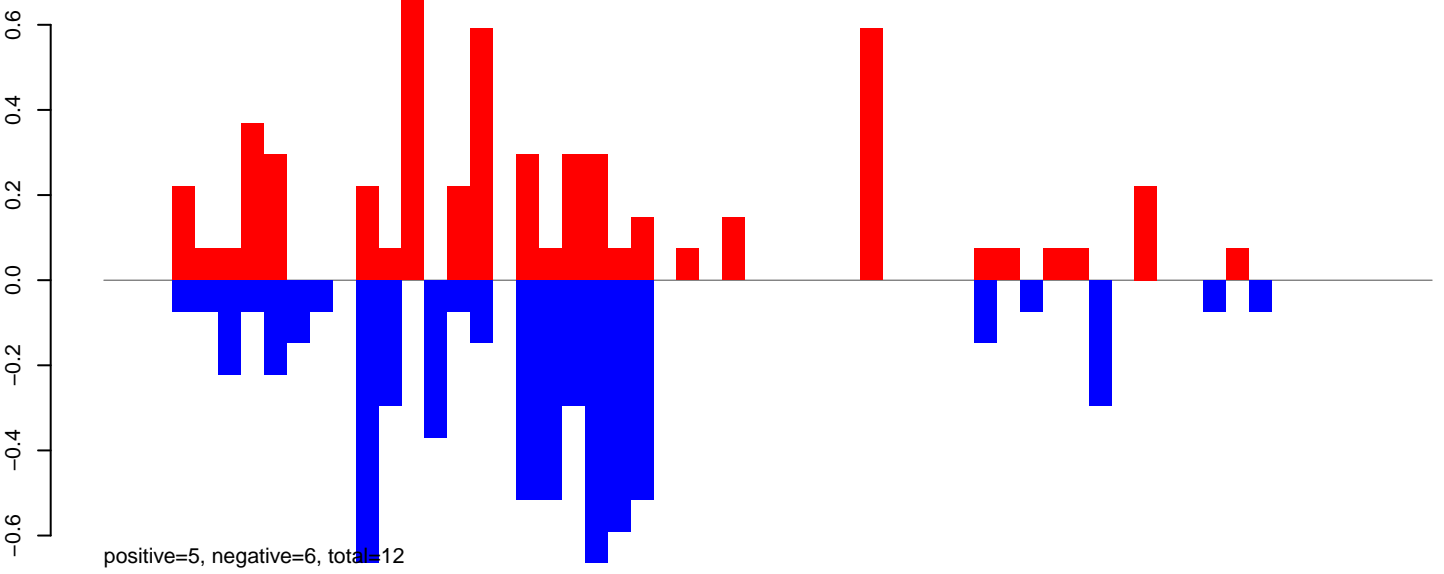
CuQuin_Testes_NL.rep



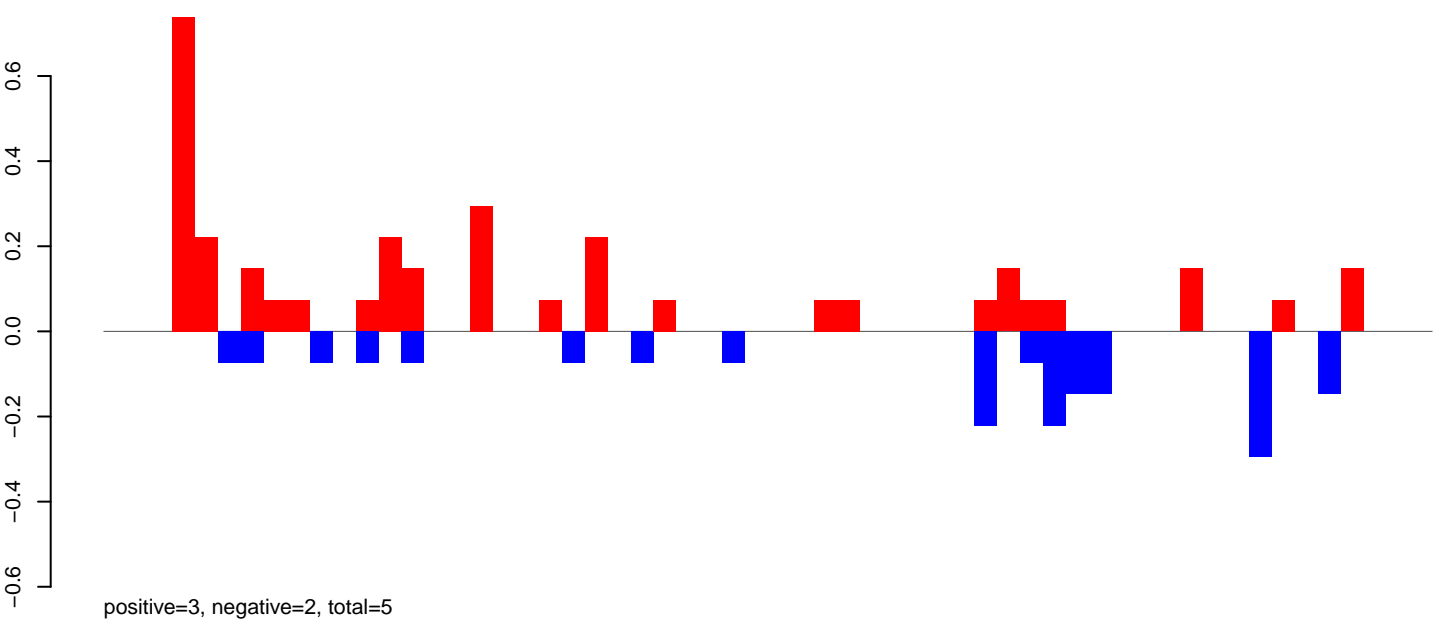
positive=43, negative=26, total=69

Window size=25, length=1314, TE@rnd-3_family-190@DNA_TcMar@3273:1-1314

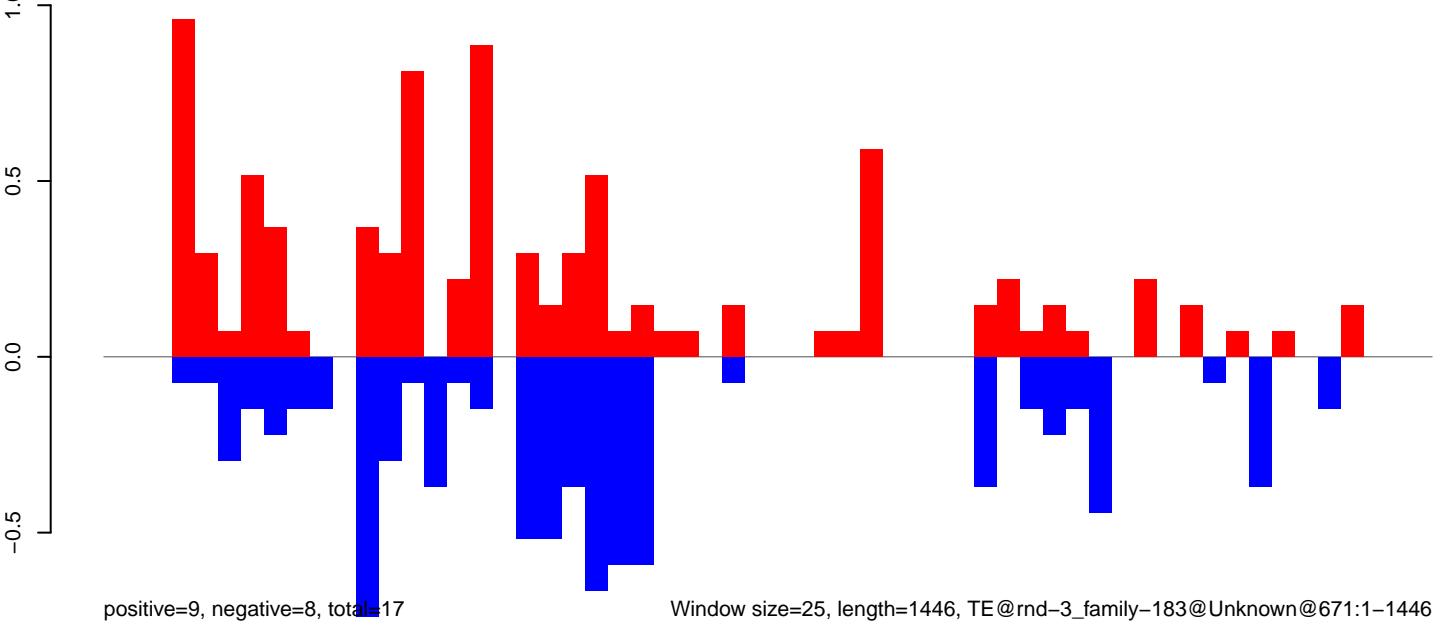
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

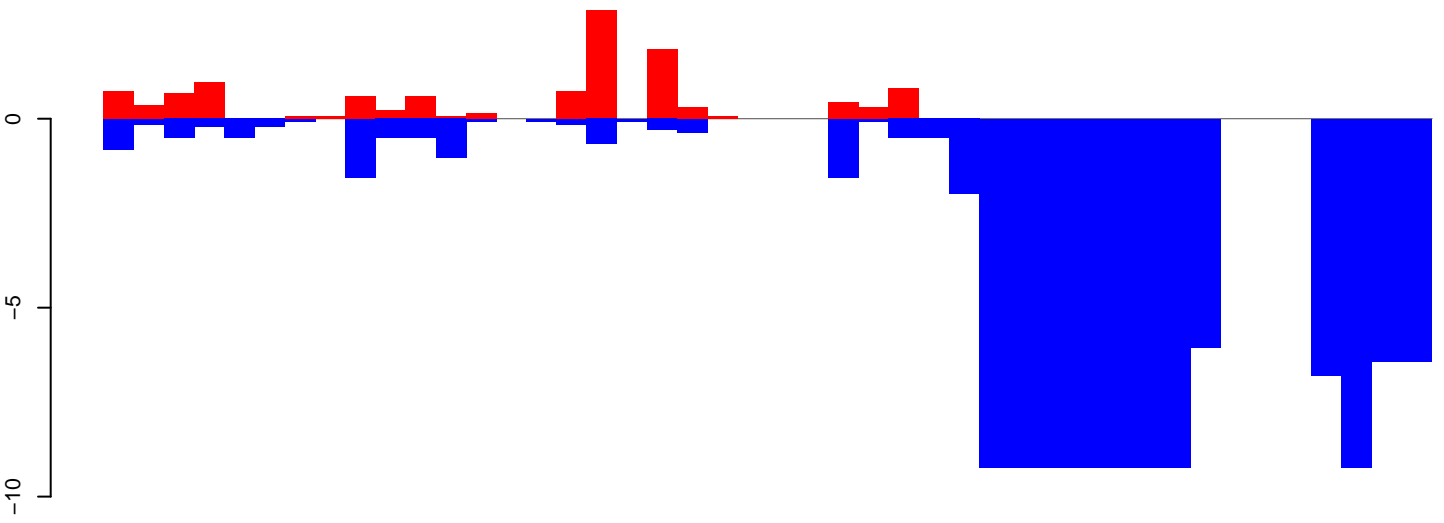


CuQuin_Testes_NL.rep



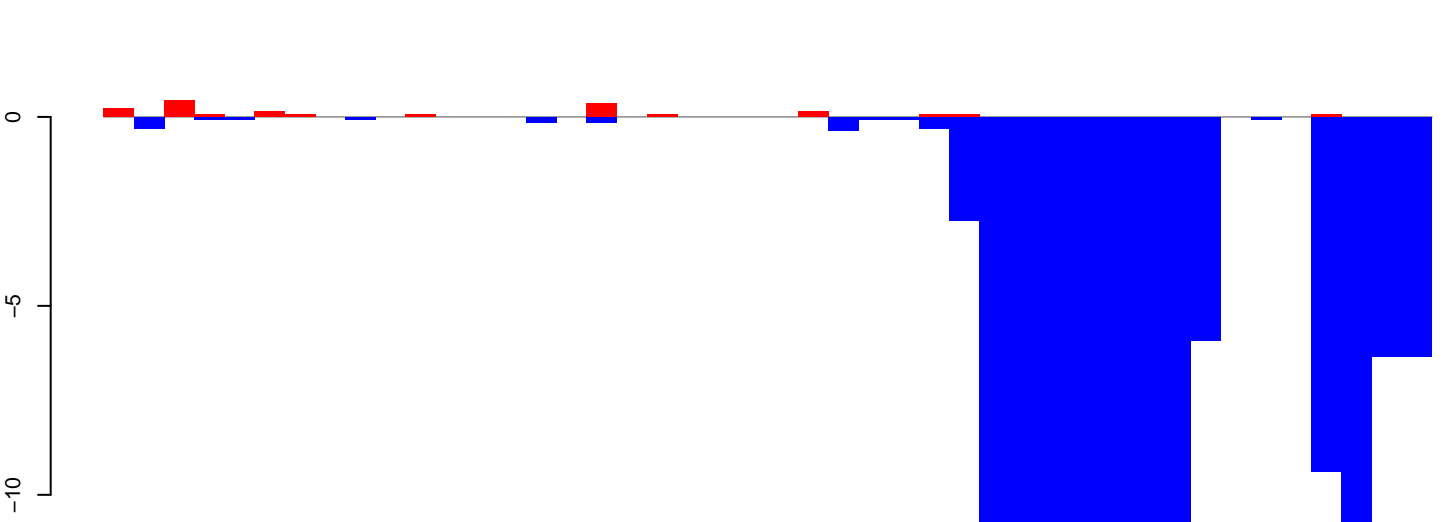
Window size=25, length=1446, TE@rnd-3_family-183@Unknown@671:1-1446

CuQuin_Testes_NL.18_23.rep



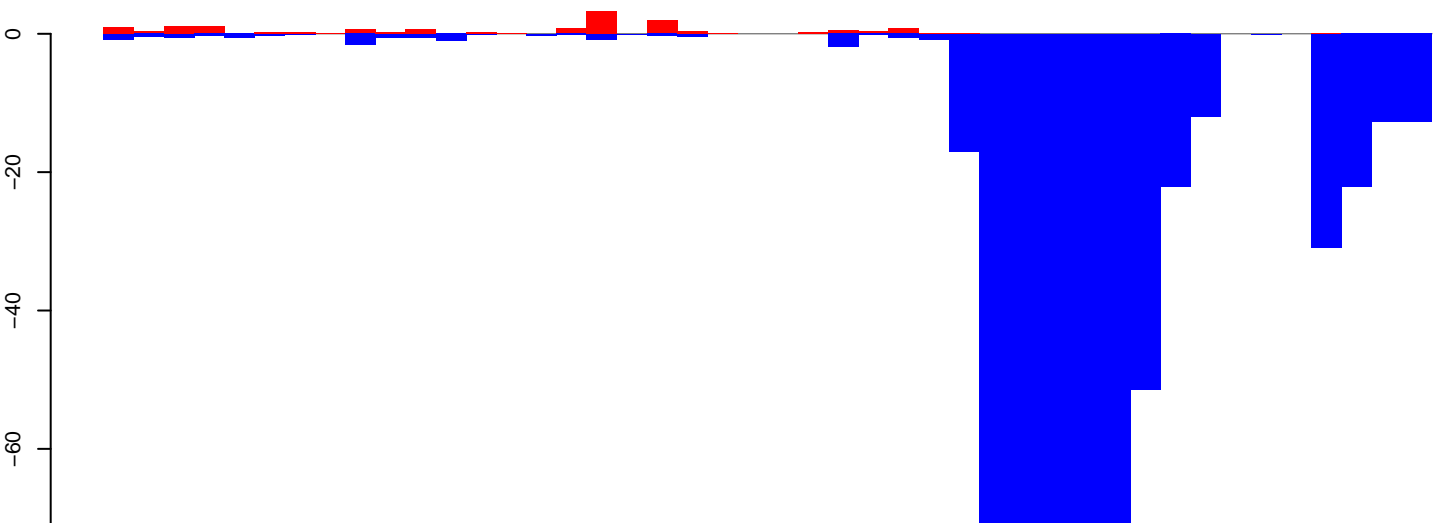
positive=12, negative=112, total=124

CuQuin_Testes_NL.24_35.rep



positive=2, negative=136, total=138

CuQuin_Testes_NL.rep

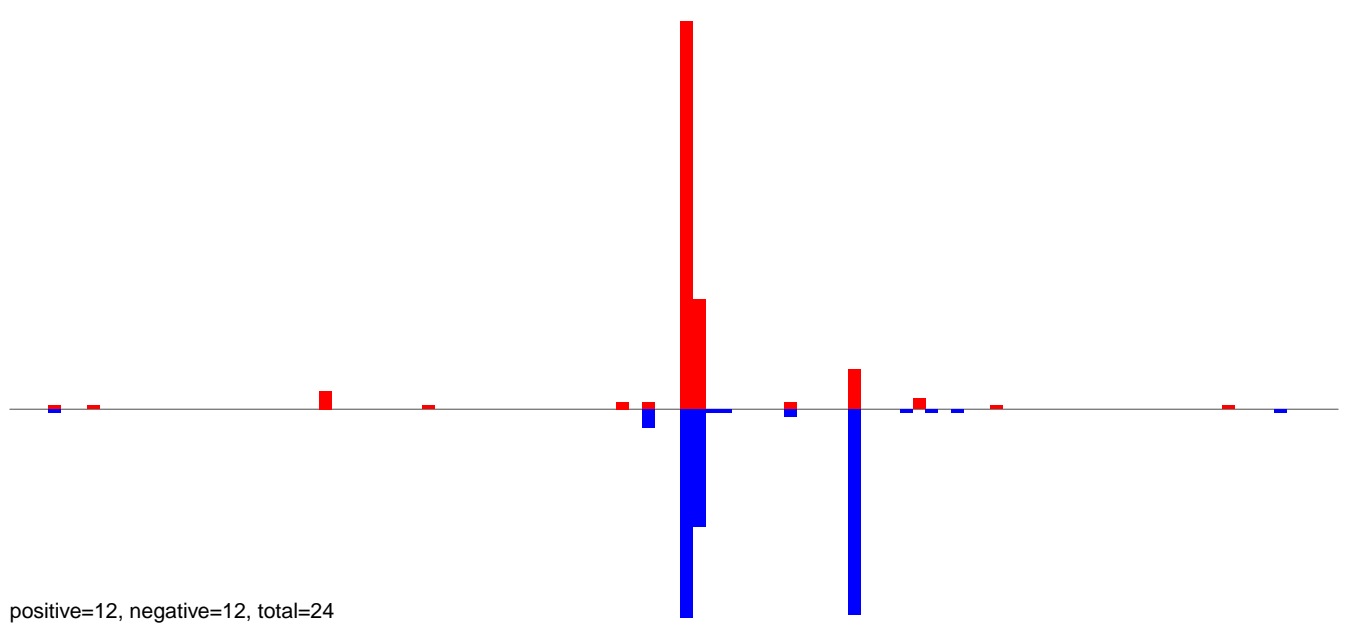


positive=14, negative=609, total=622

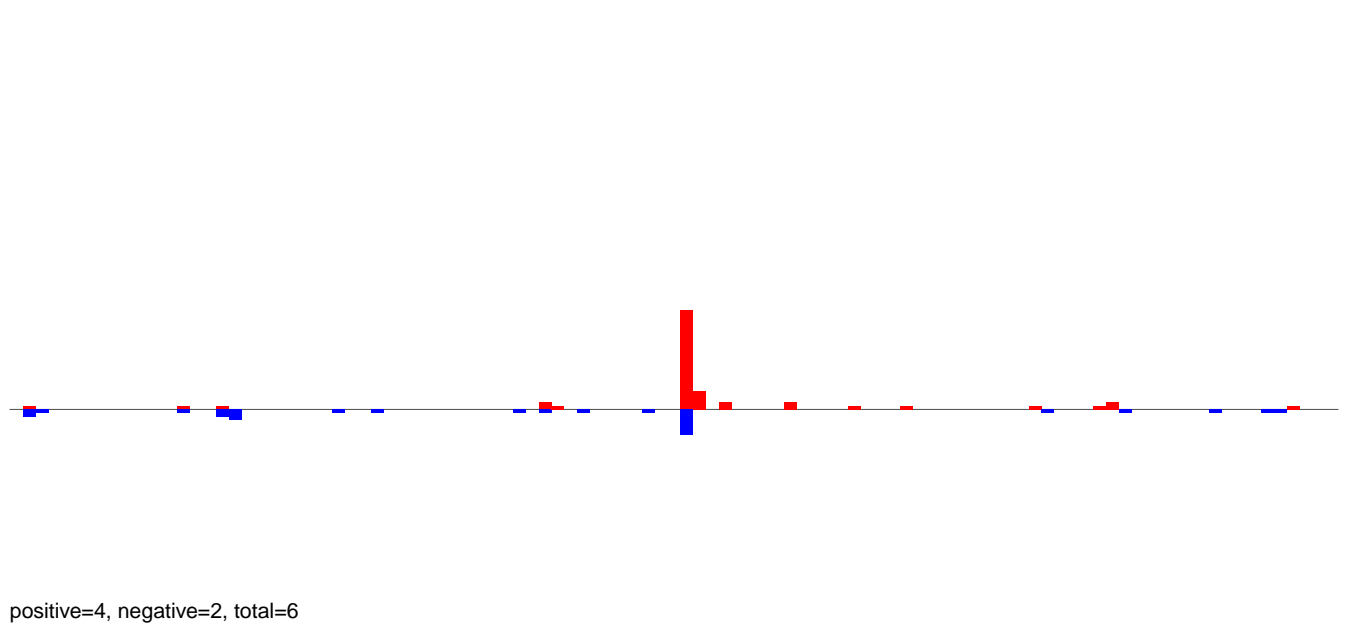
Window size=25, length=1089, TE@rnd-3_family-176@DNA@1323:1-1089

0 200 400 600 800 1000

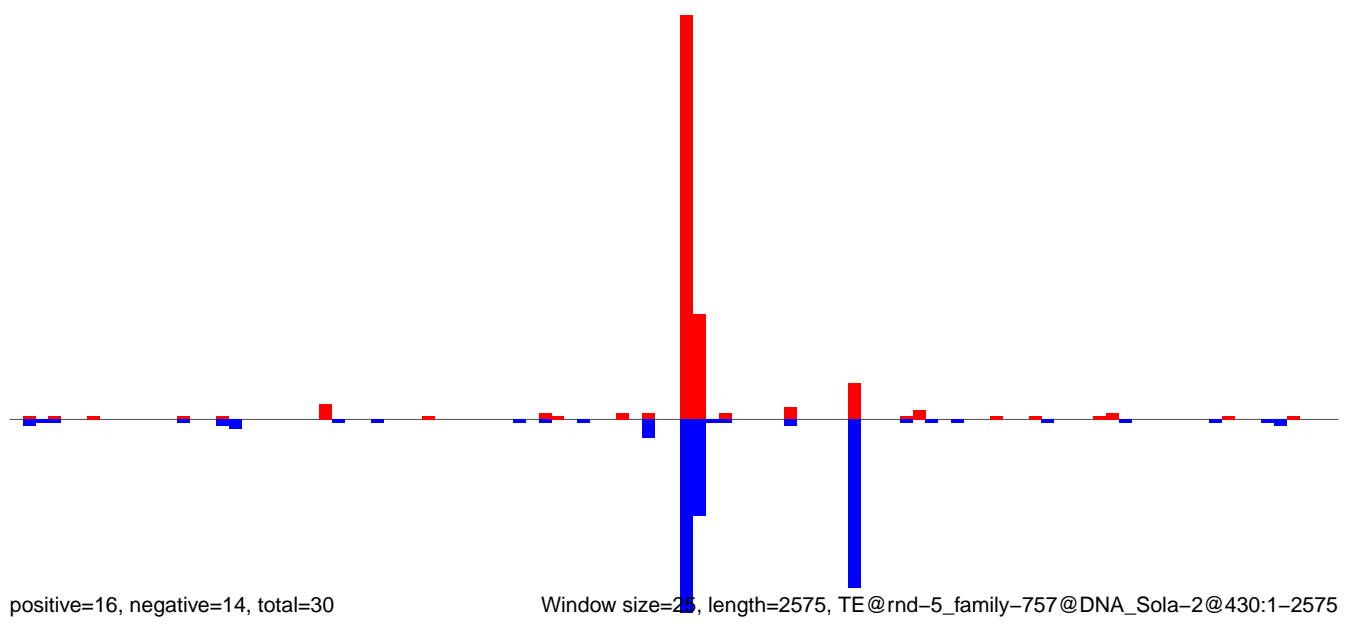
CuQuin_Testes_NL.18_23.rep



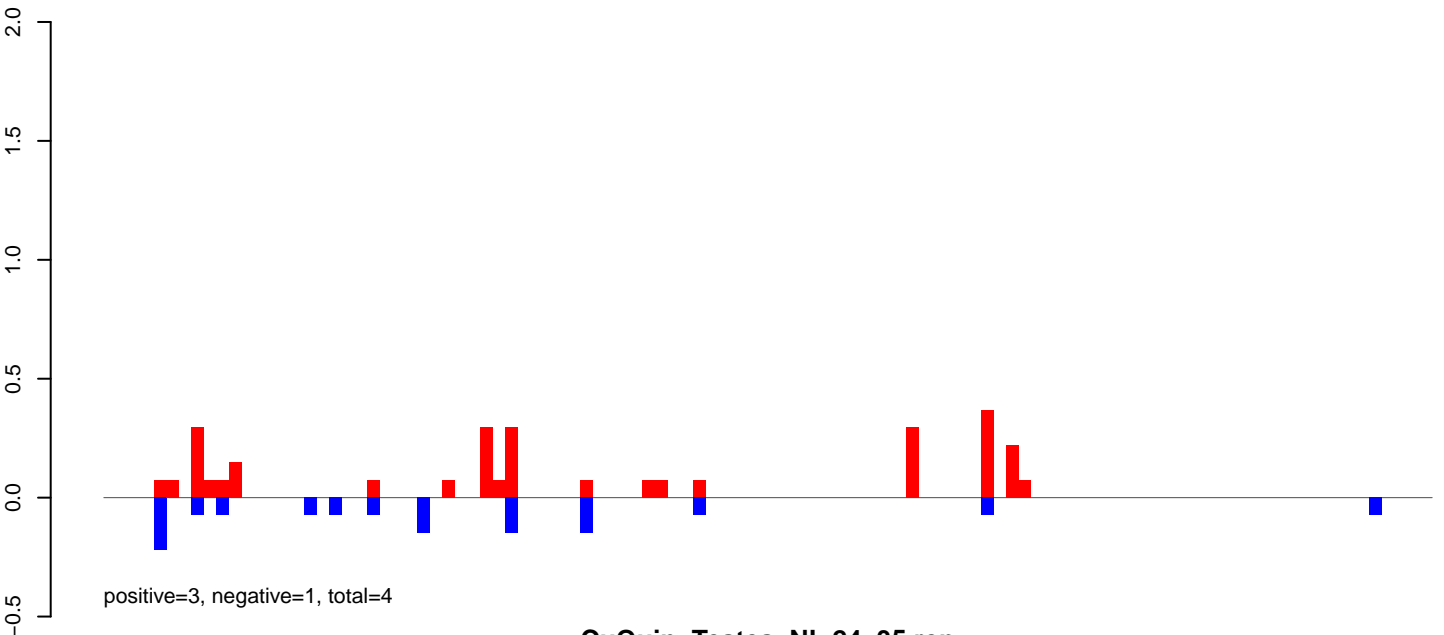
CuQuin_Testes_NL.24_35.rep



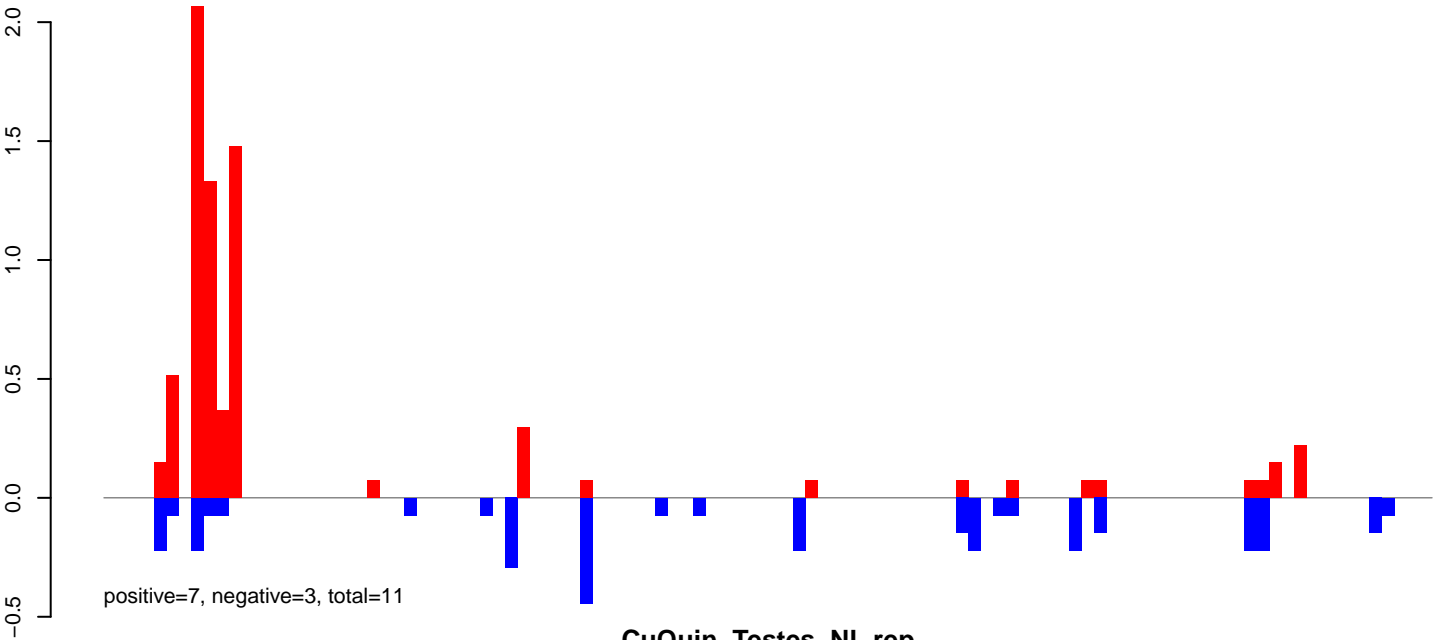
CuQuin_Testes_NL.rep



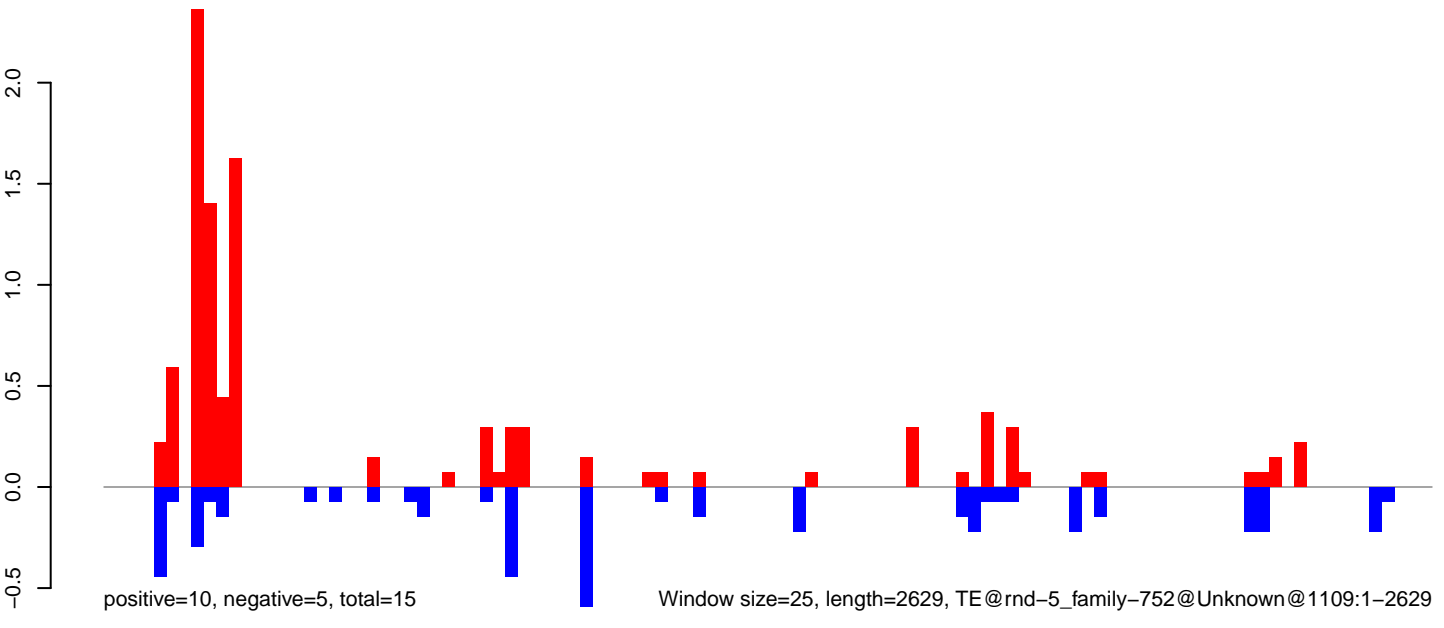
CuQuin_Testes_NL.18_23.rep



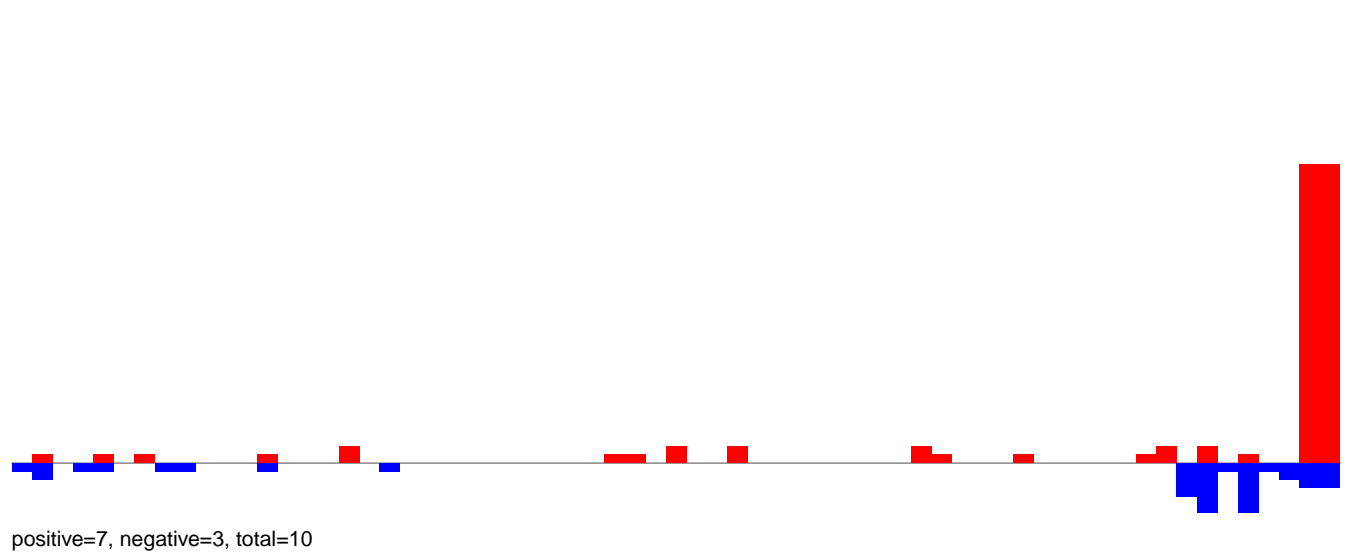
CuQuin_Testes_NL.24_35.rep



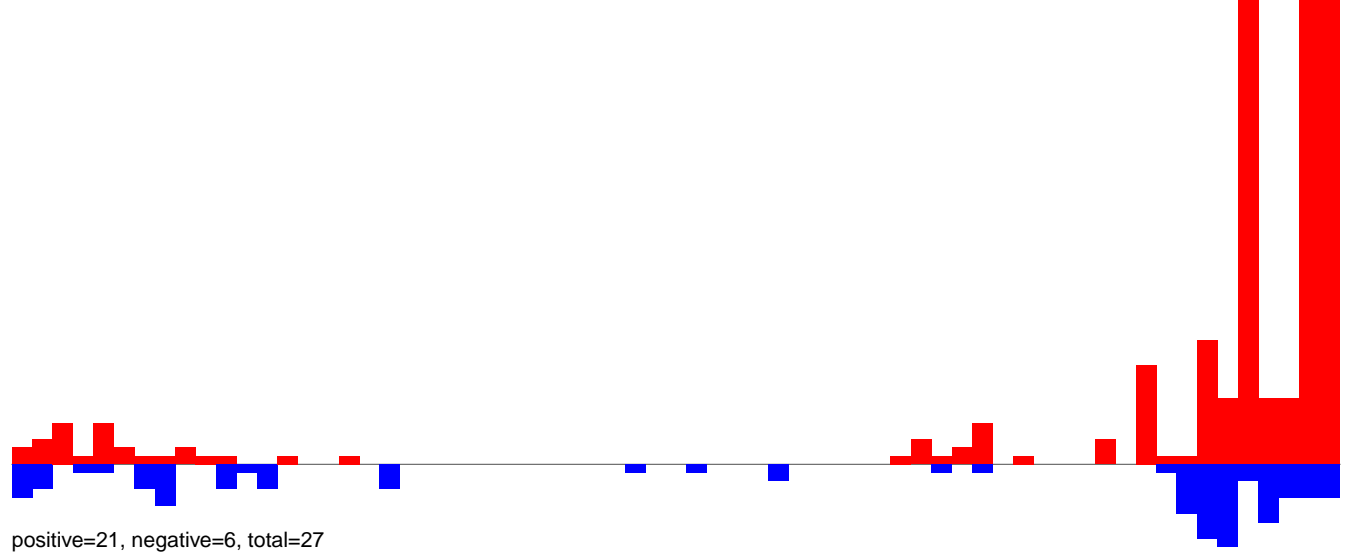
CuQuin_Testes_NL.rep



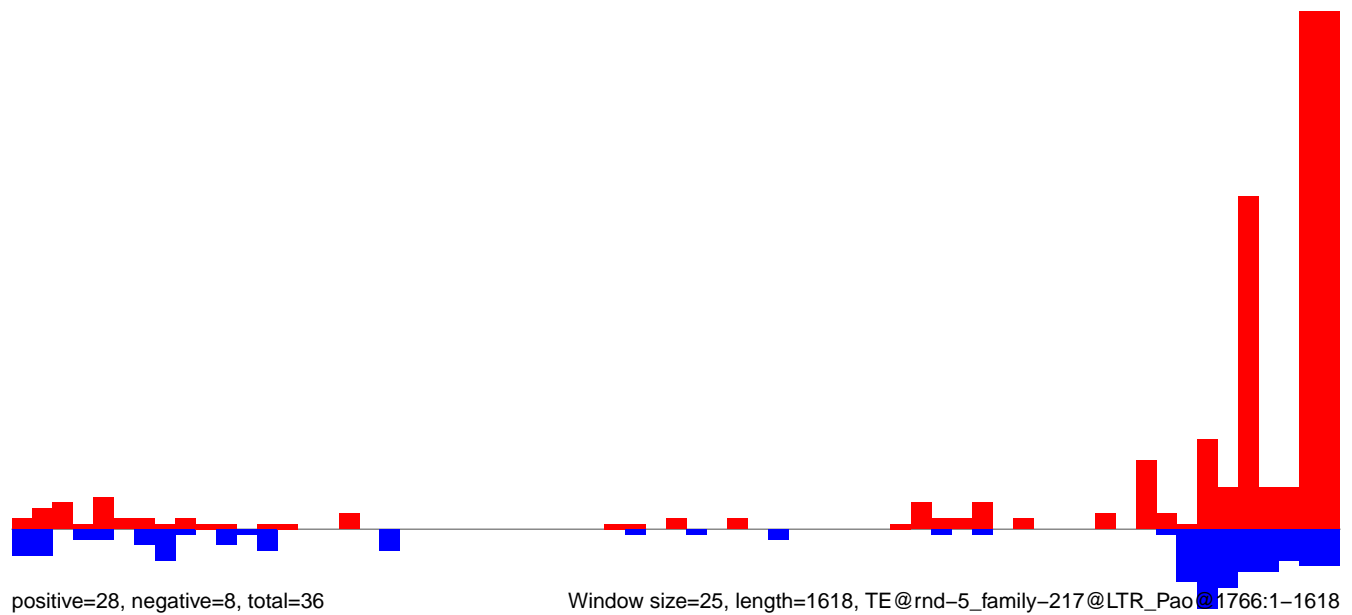
CuQuin_Testes_NL.18_23.rep



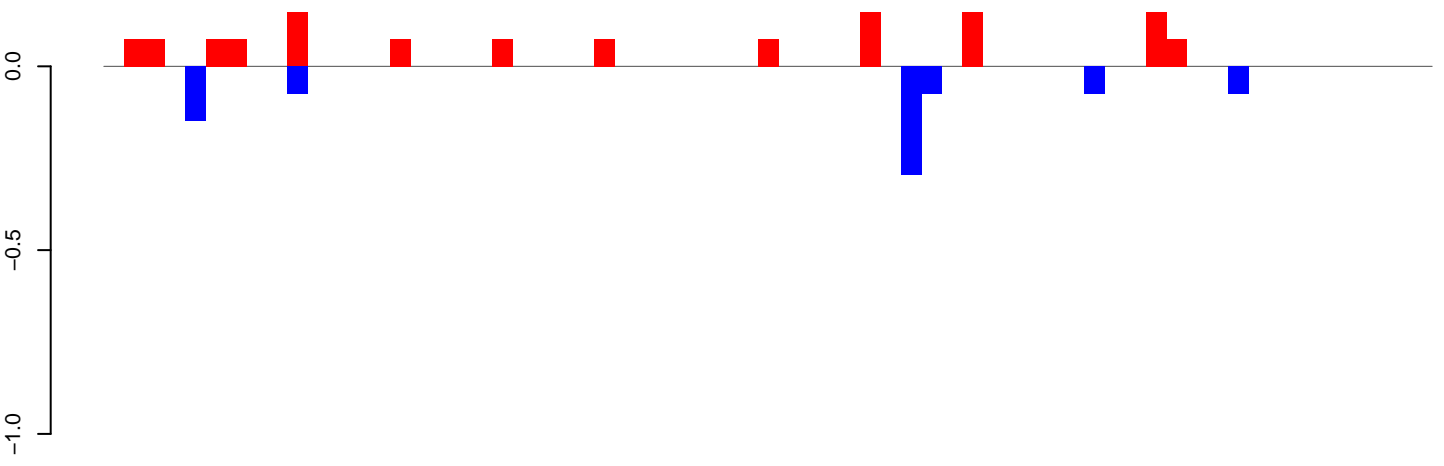
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

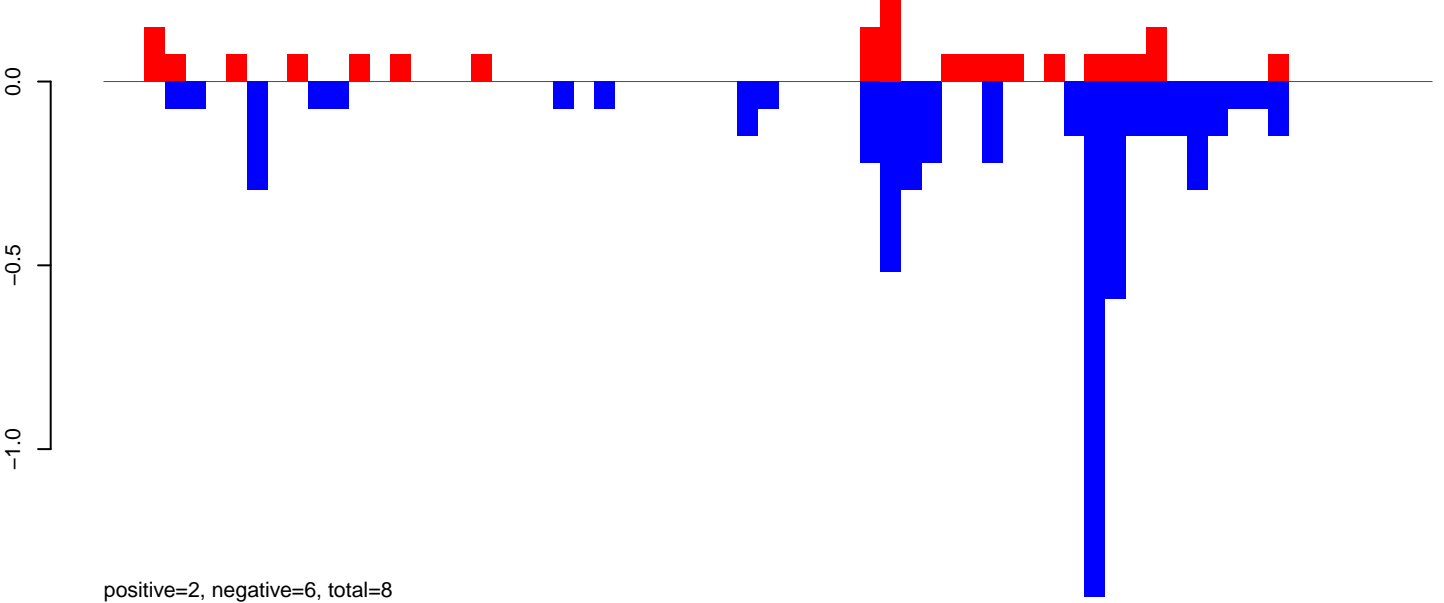


CuQuin_Testes_NL.18_23.rep



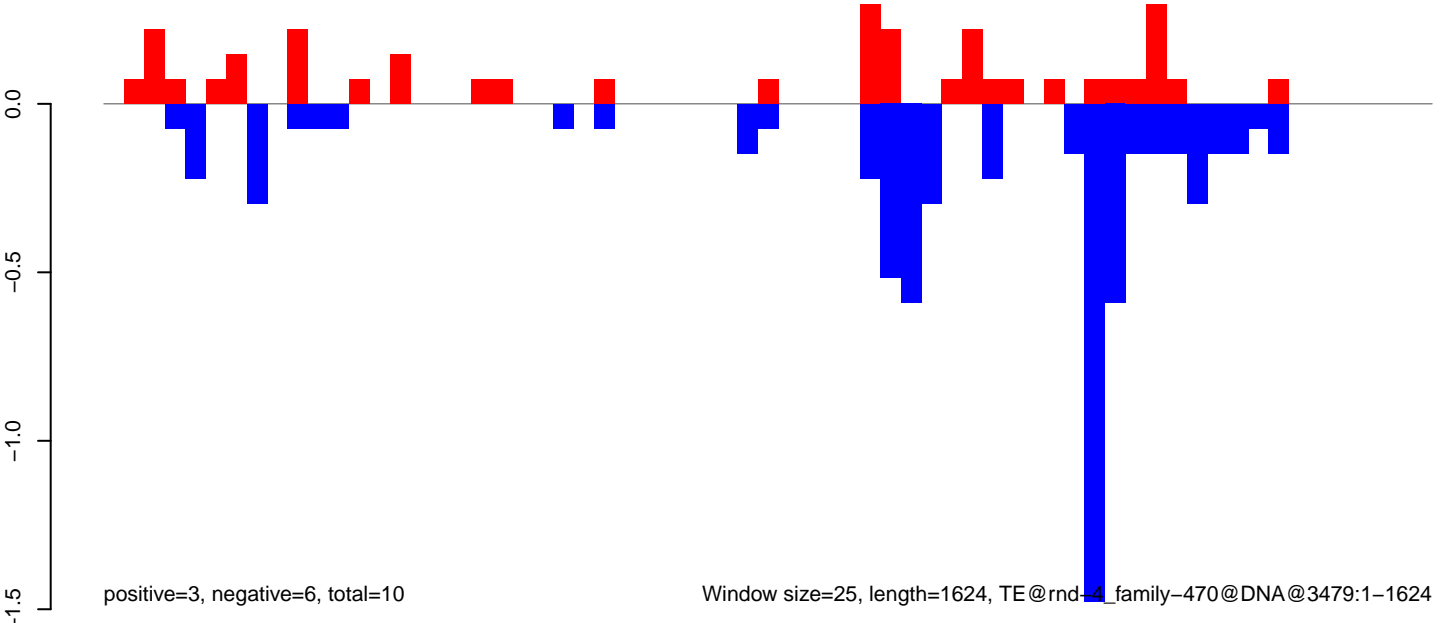
positive=1, negative=1, total=2

CuQuin_Testes_NL.24_35.rep



positive=2, negative=6, total=8

CuQuin_Testes_NL.rep

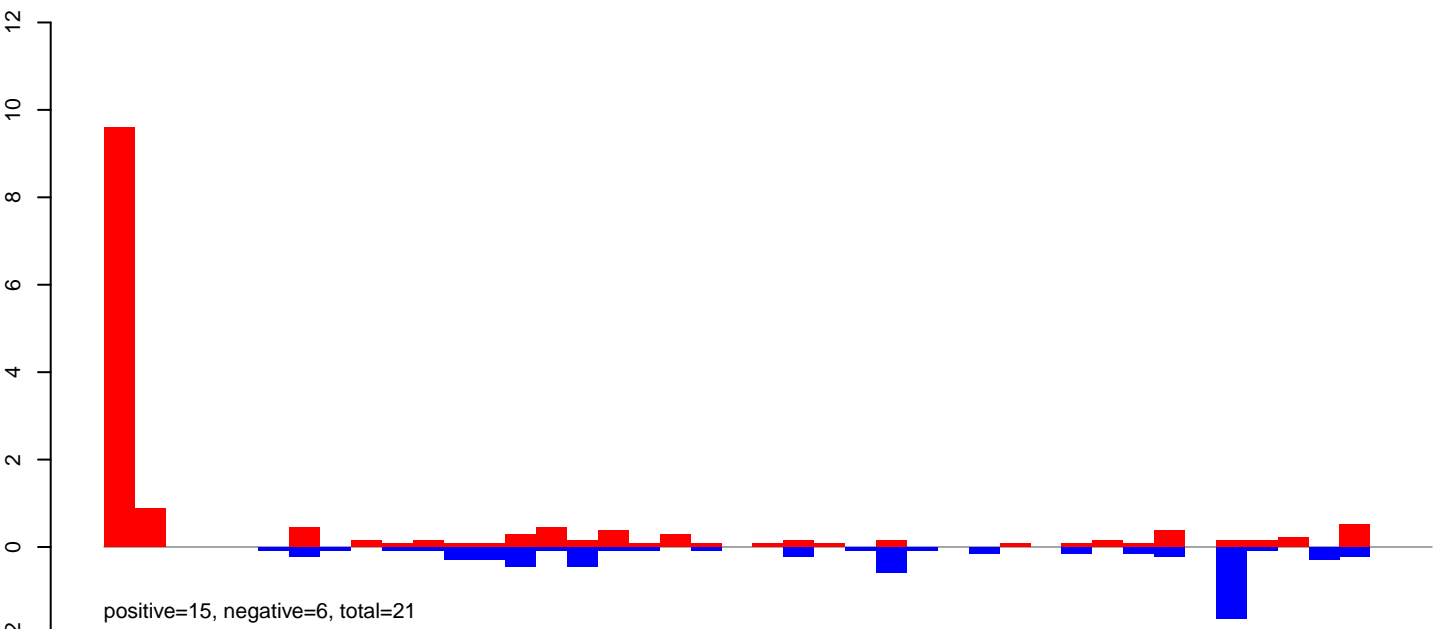


positive=3, negative=6, total=10

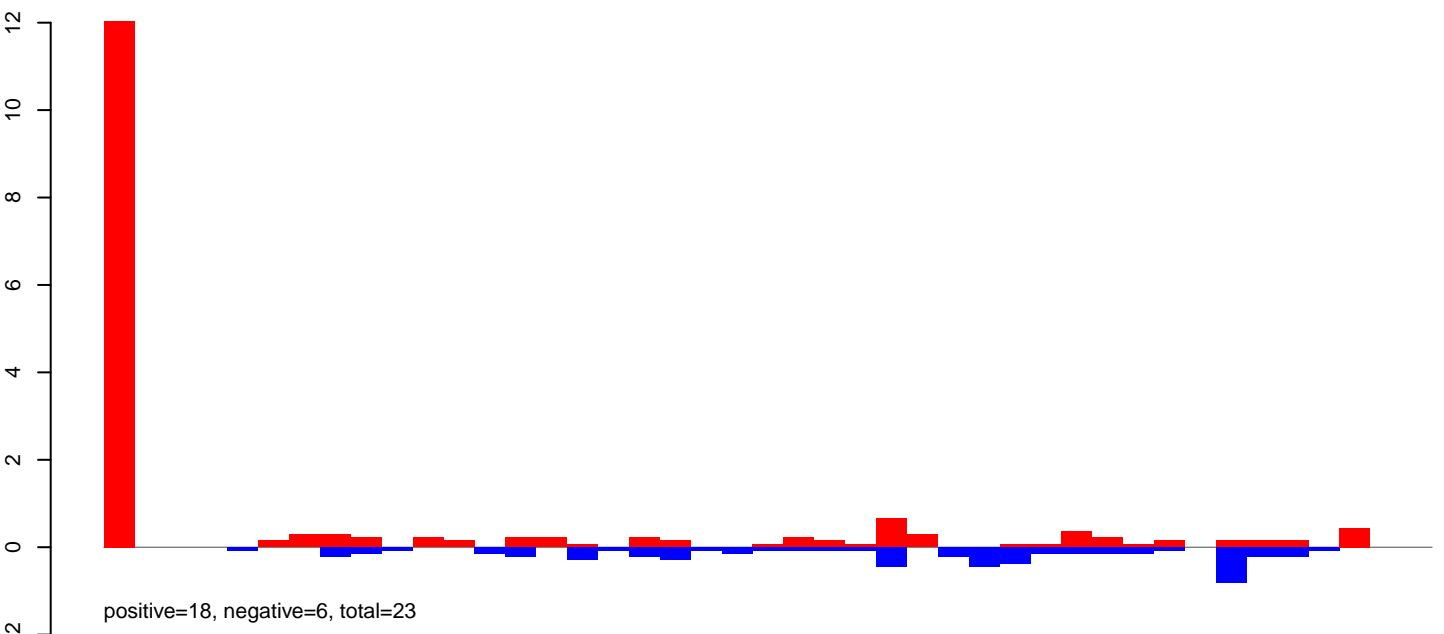
Window size=25, length=1624, TE@rnd-4_family-470@DNA@3479:1-1624

0 500 1000 1500

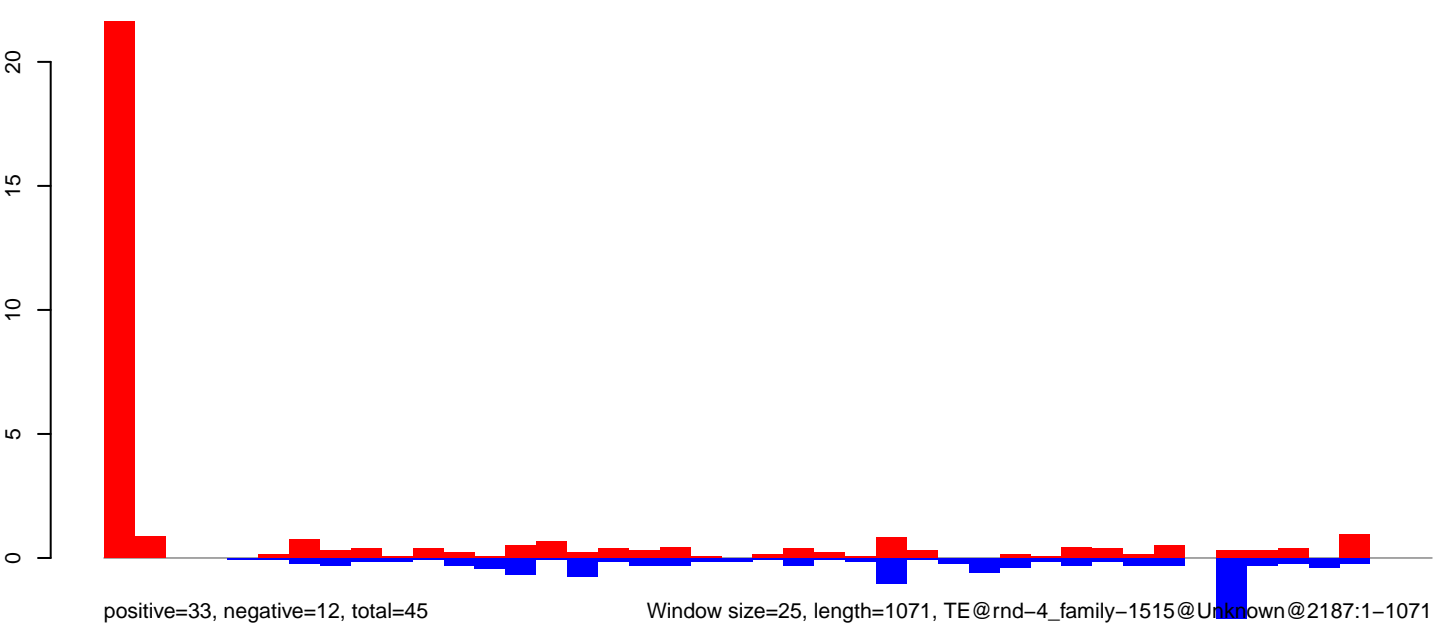
CuQuin_Testes_NL.18_23.rep



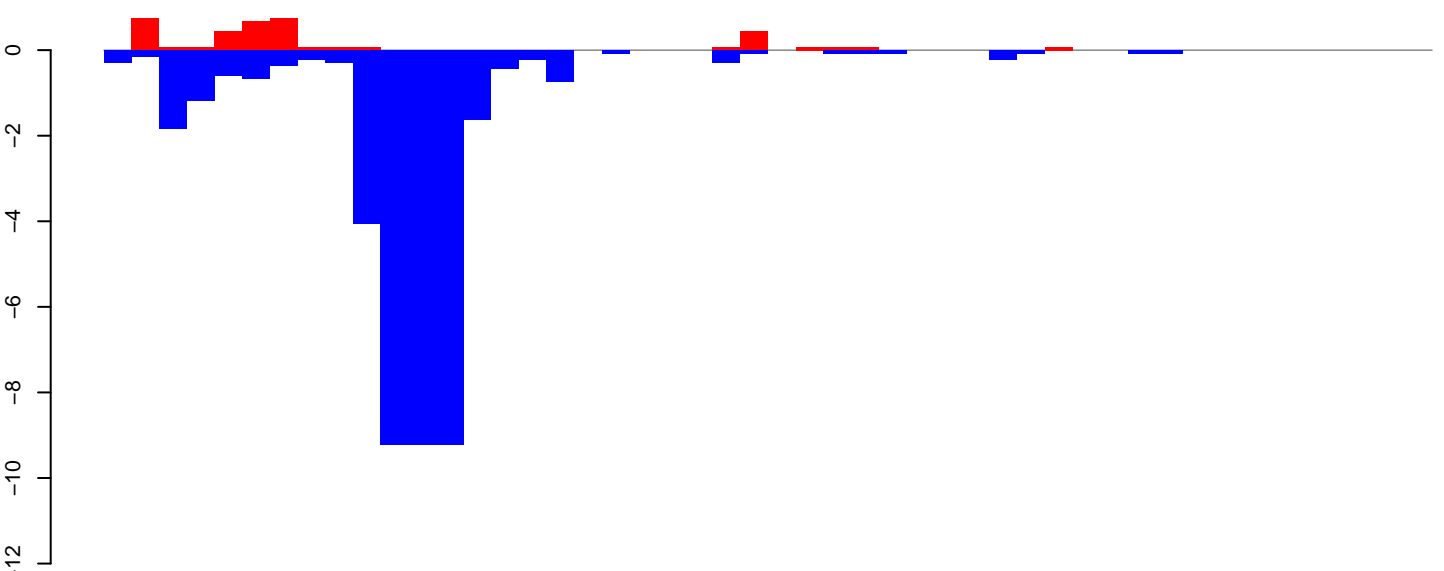
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

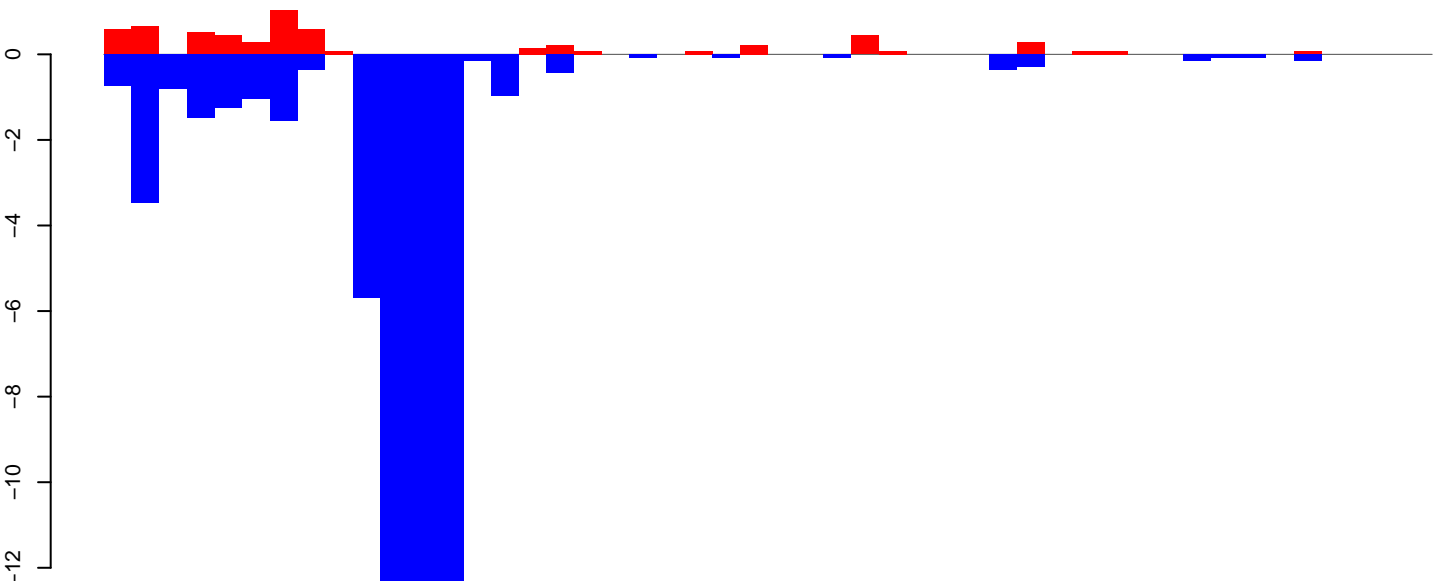


CuQuin_Testes_NL.18_23.rep



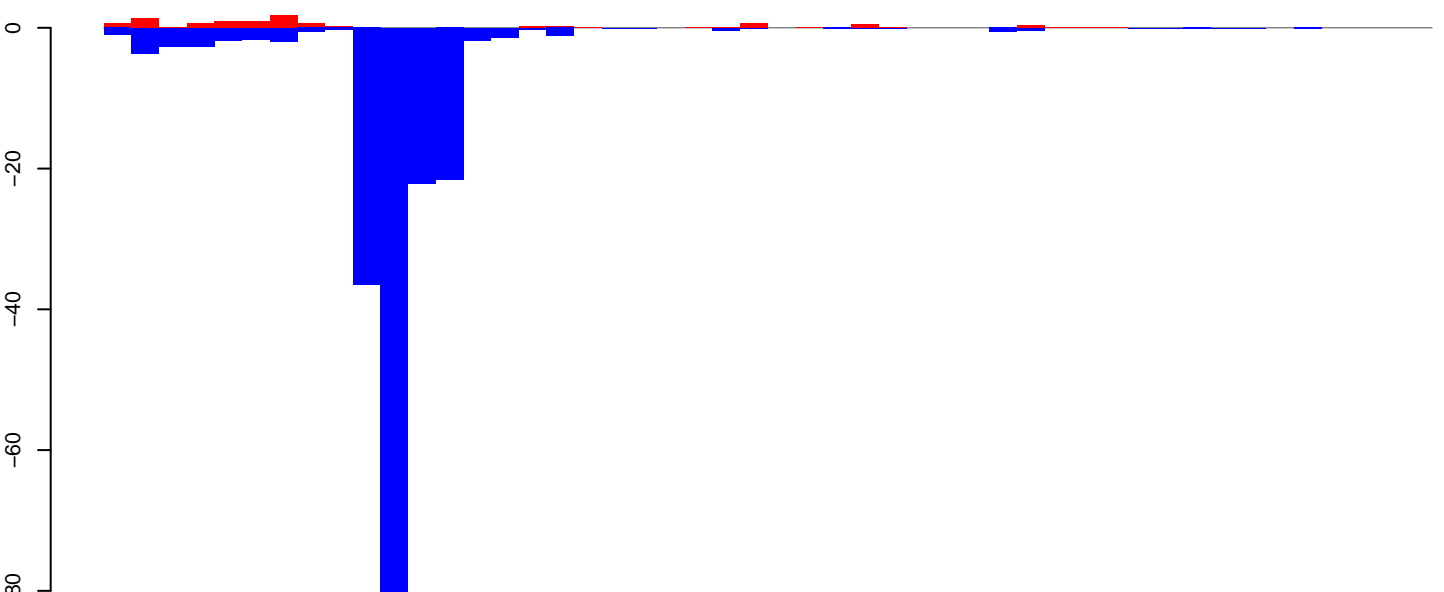
positive=4, negative=41, total=45

CuQuin_Testes_NL.24_35.rep



positive=6, negative=58, total=64

CuQuin_Testes_NL.rep

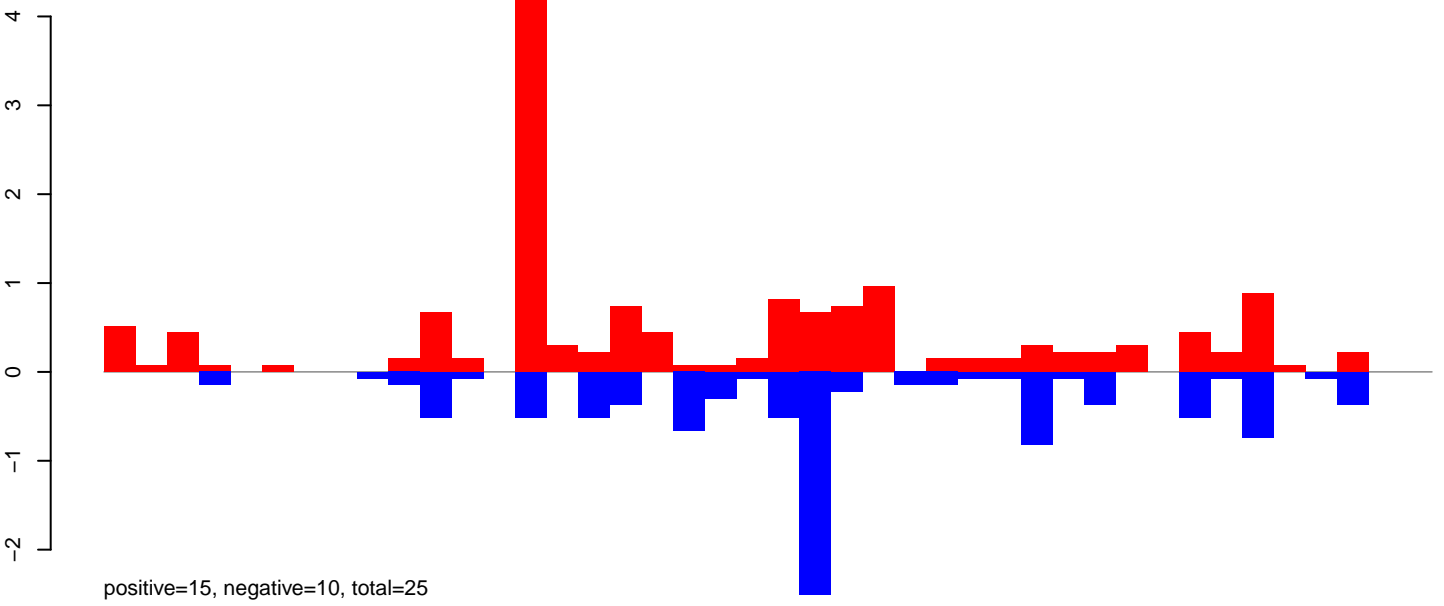


positive=10, negative=187, total=197

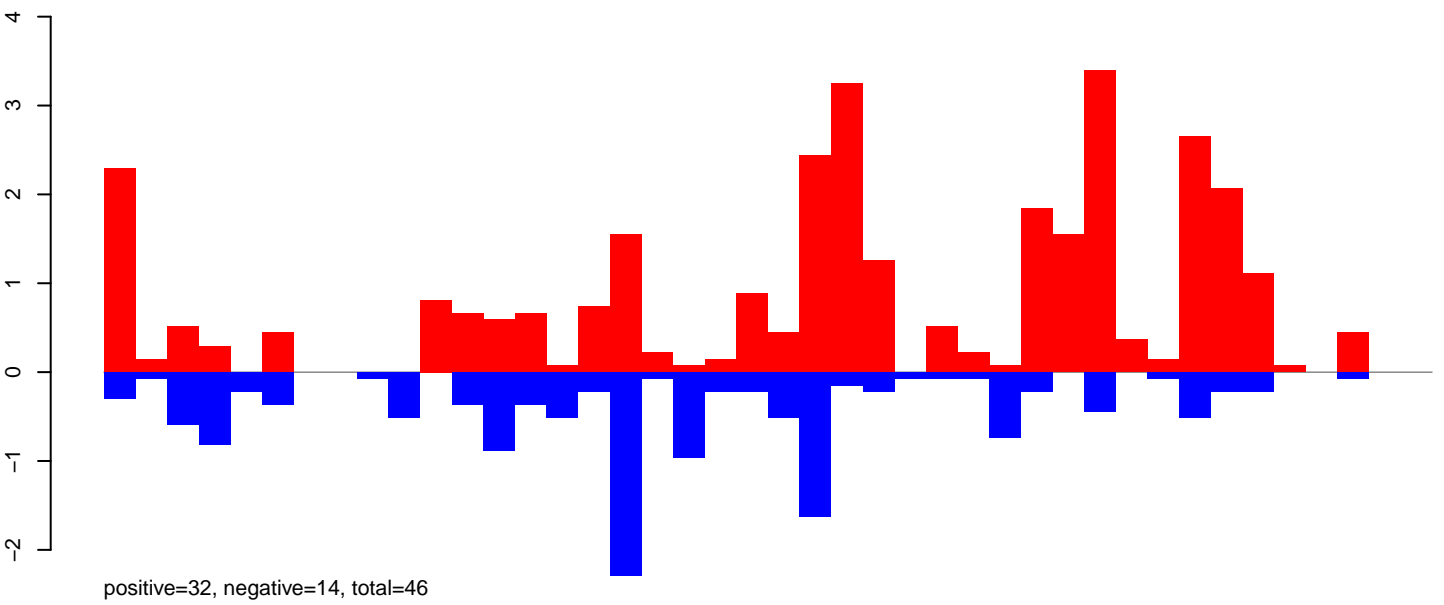
Window size=25, length=1183, TE@rnd-3_family-729@DNA_Crypton-I@670:1-1183

0 200 400 600 800 1000 1200

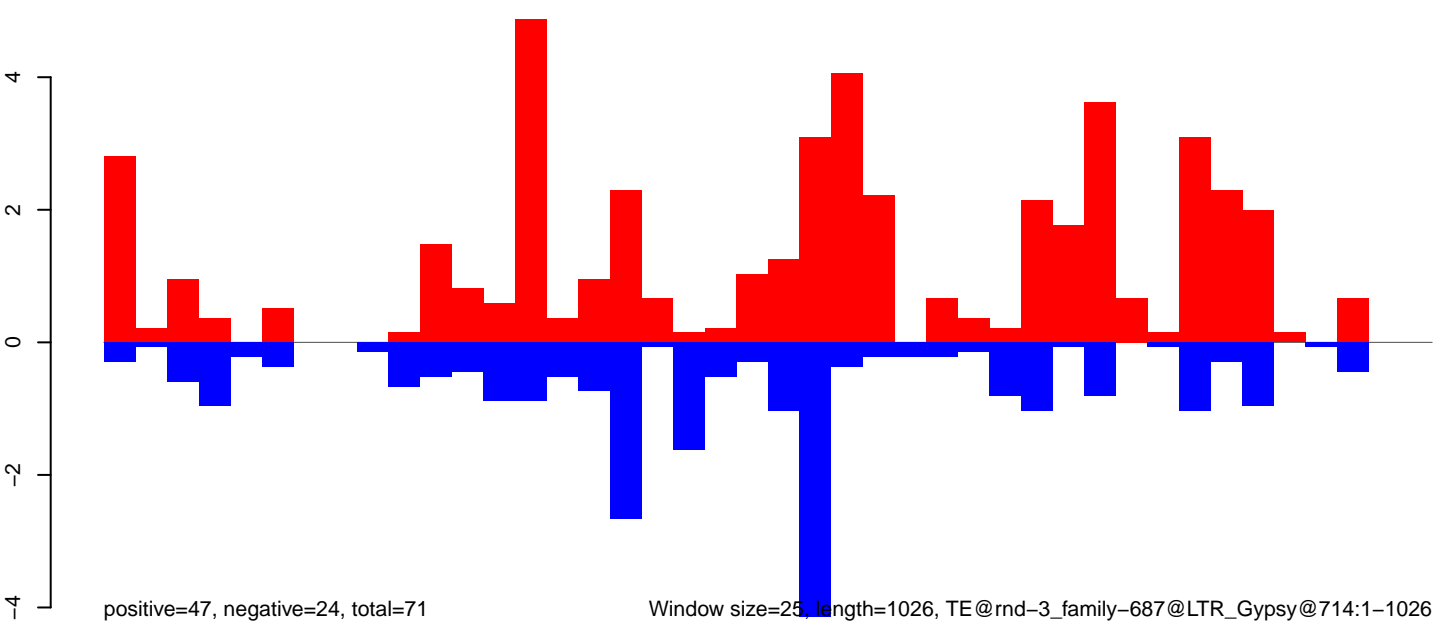
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

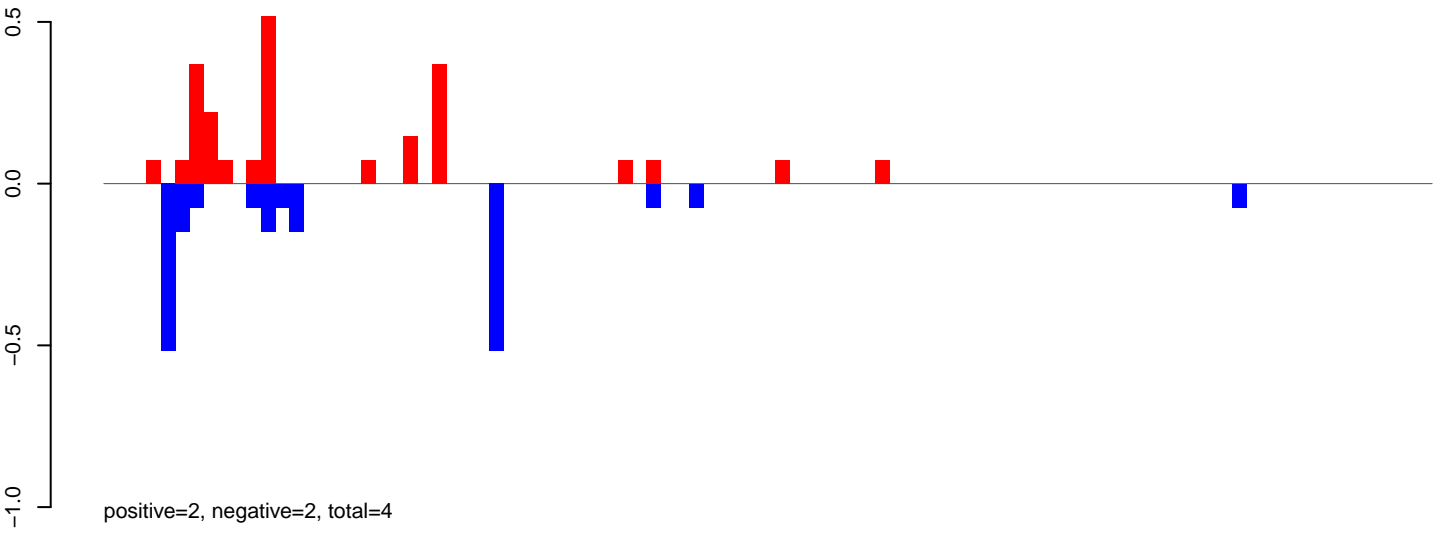


CuQuin_Testes_NL.rep

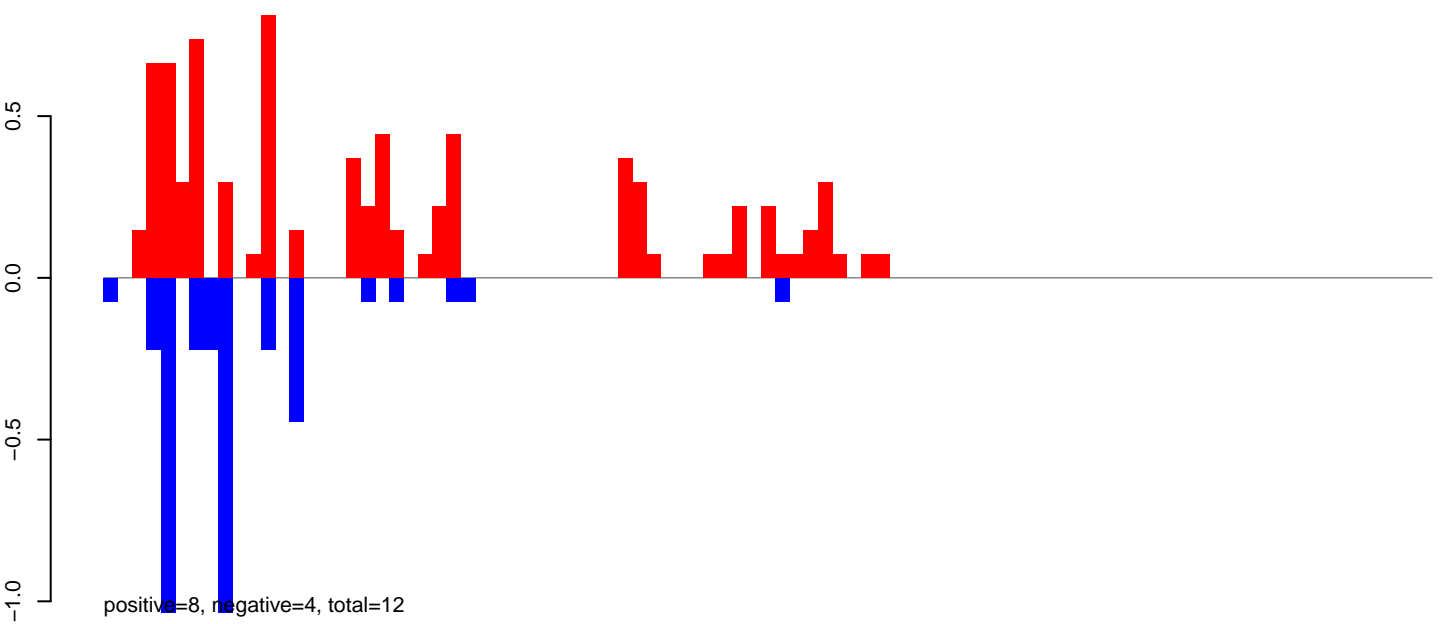


Window size=25, length=1026, TE@rnd-3_family-687@LTR_Gypsy@714:1-1026

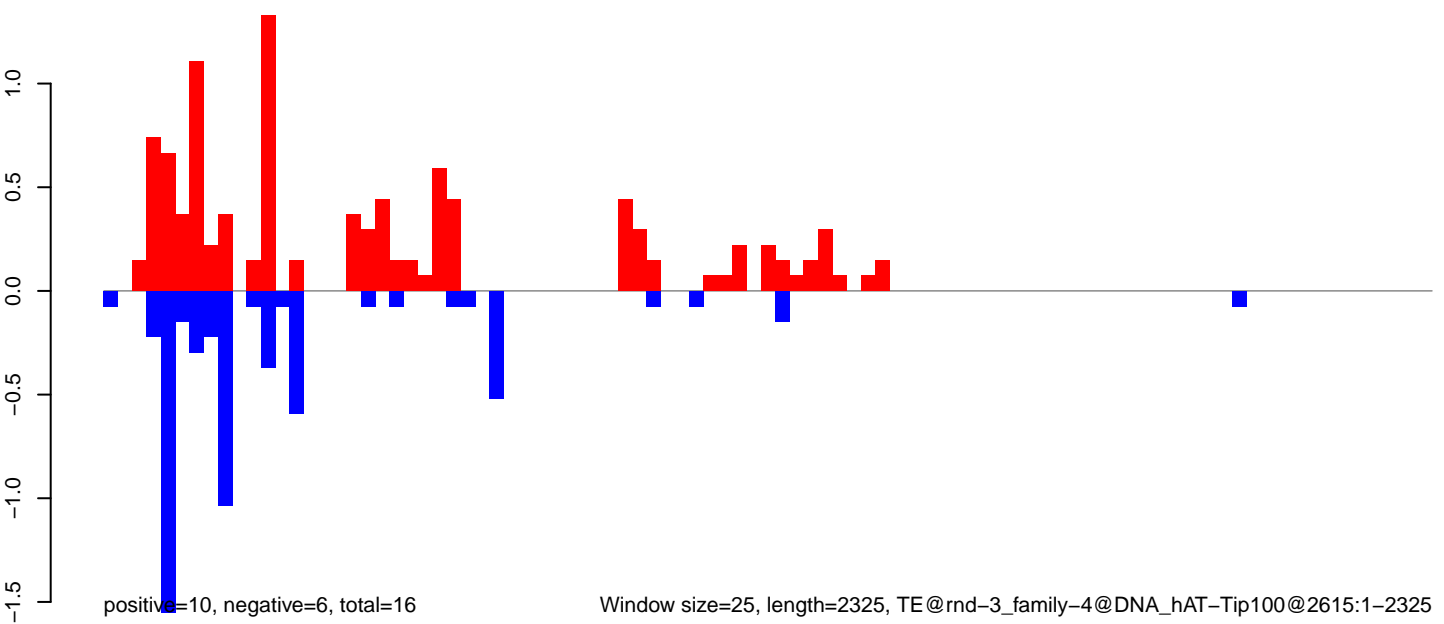
CuQuin_Testes_NL.18_23.rep



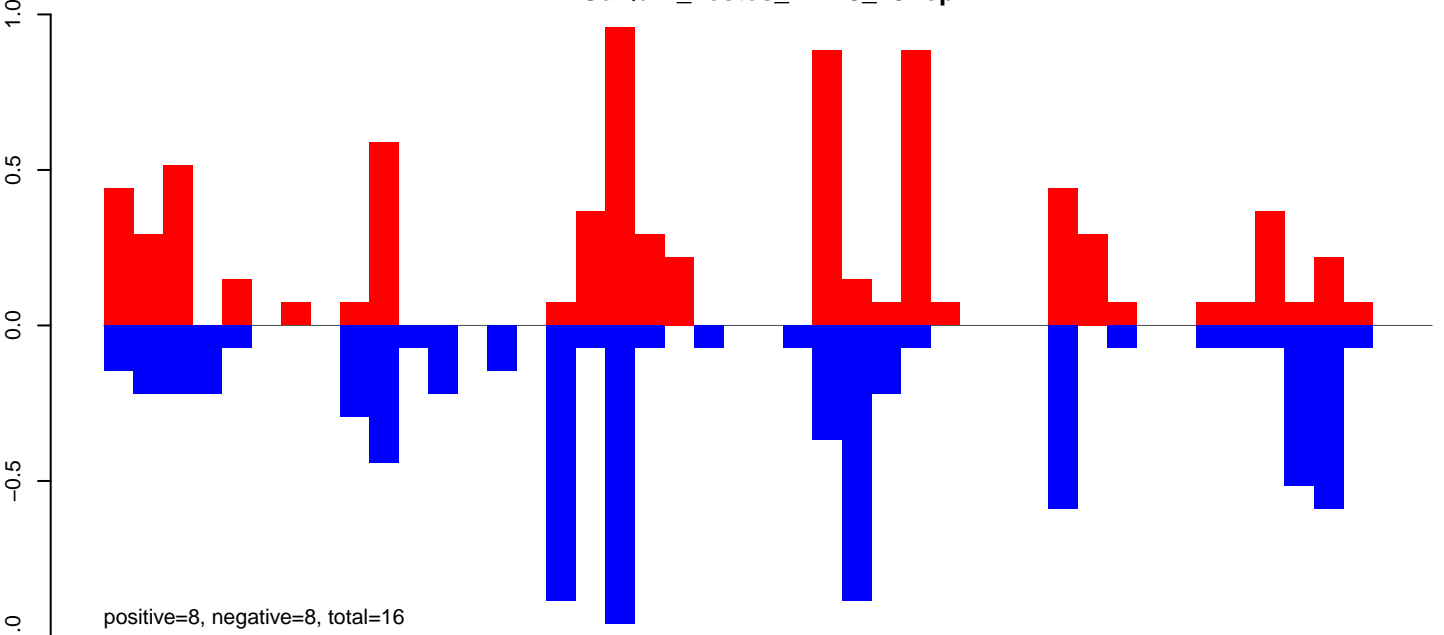
CuQuin_Testes_NL.24_35.rep



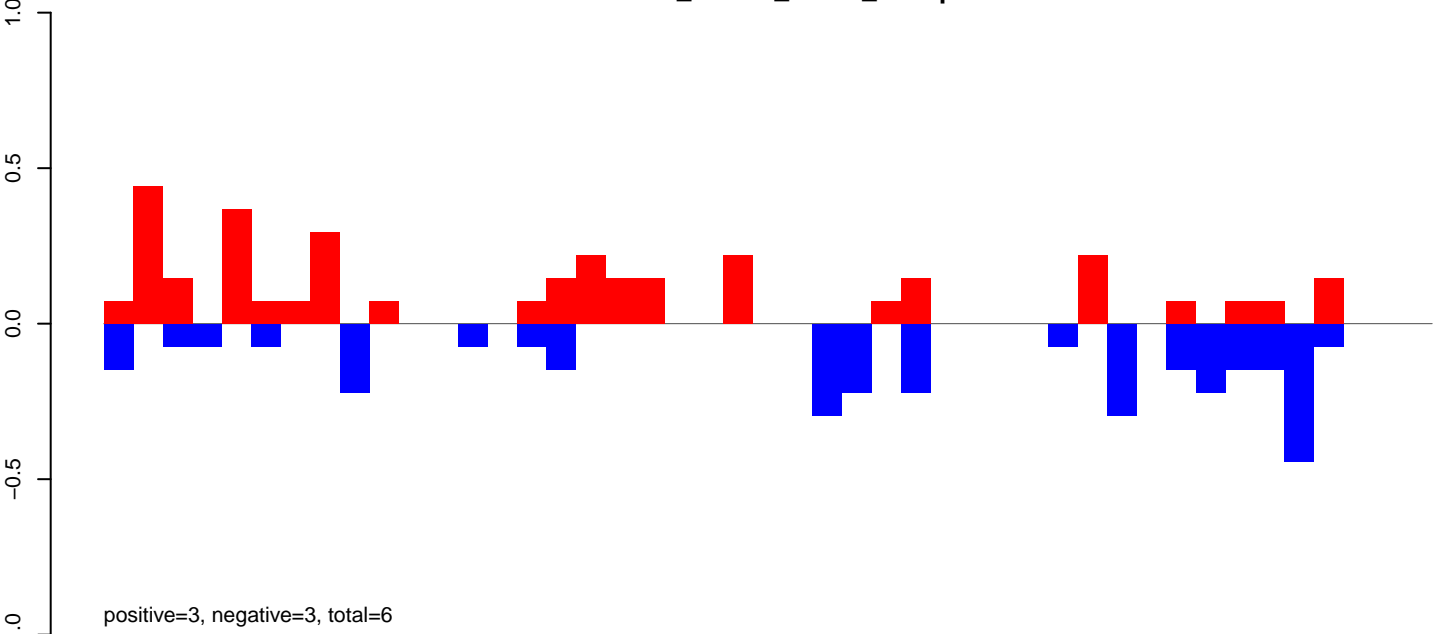
CuQuin_Testes_NL.rep



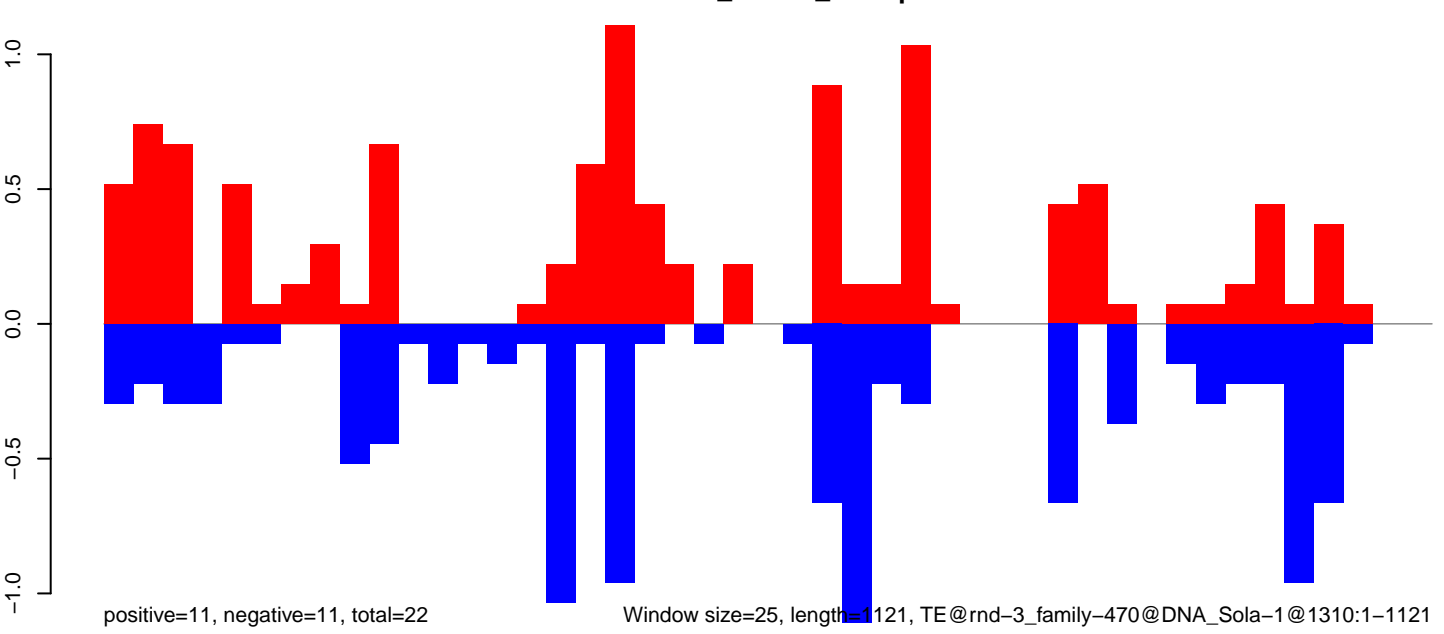
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

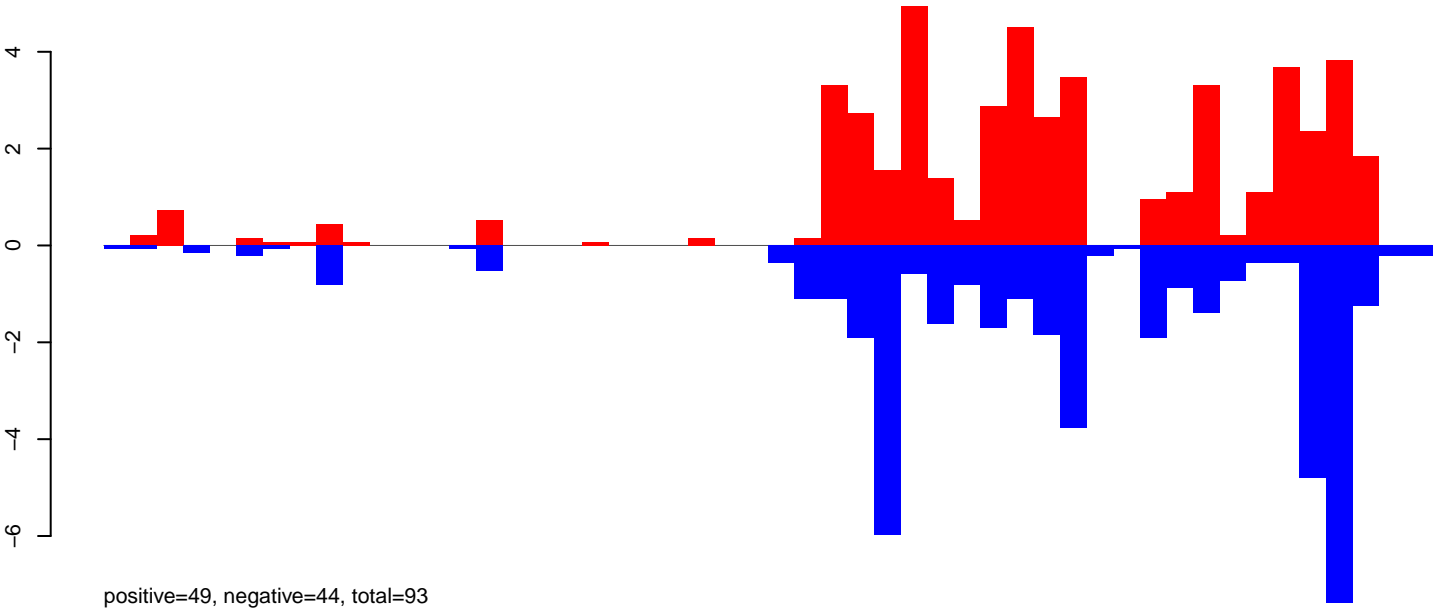


CuQuin_Testes_NL.rep

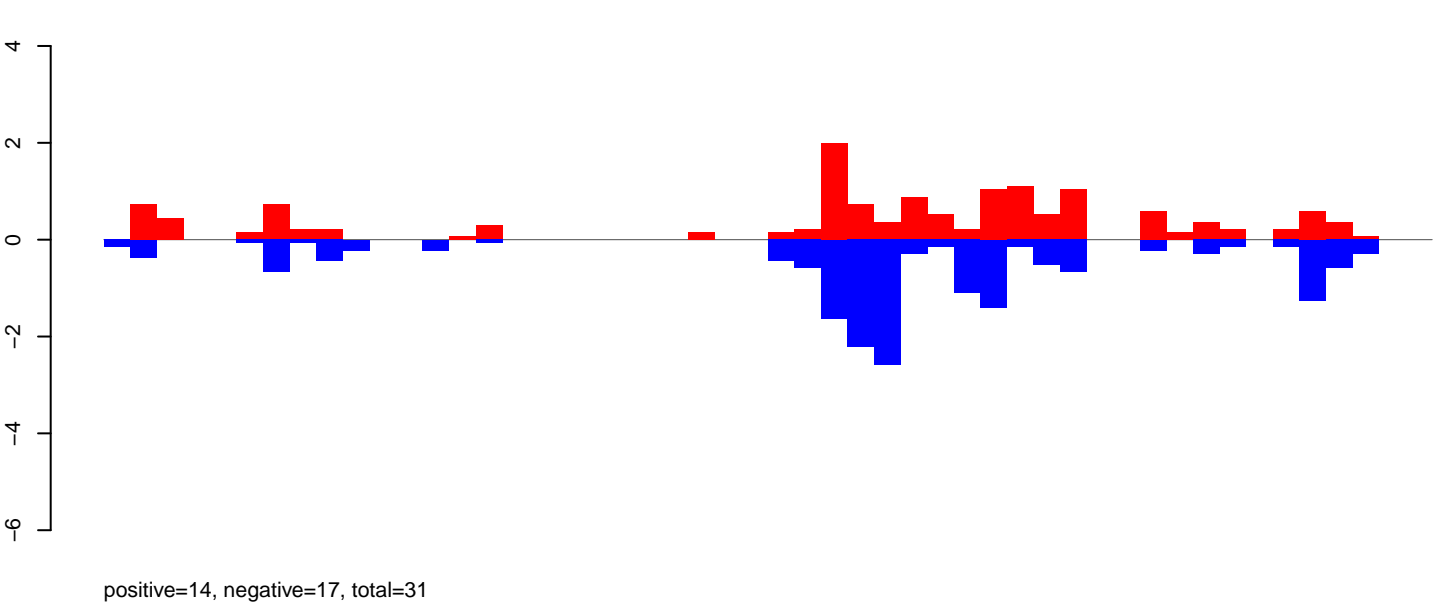


Window size=25, length=1121, TE@rnd-3_family-470@DNA_Sola-1@1310:1-1121

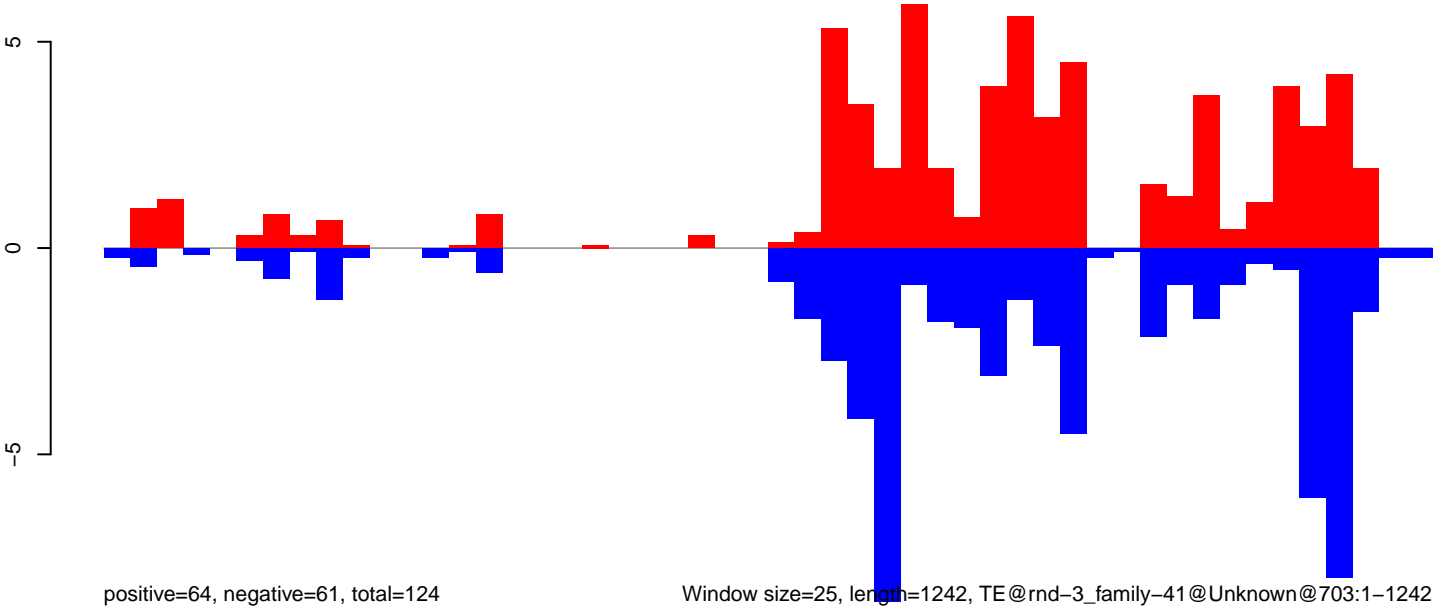
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

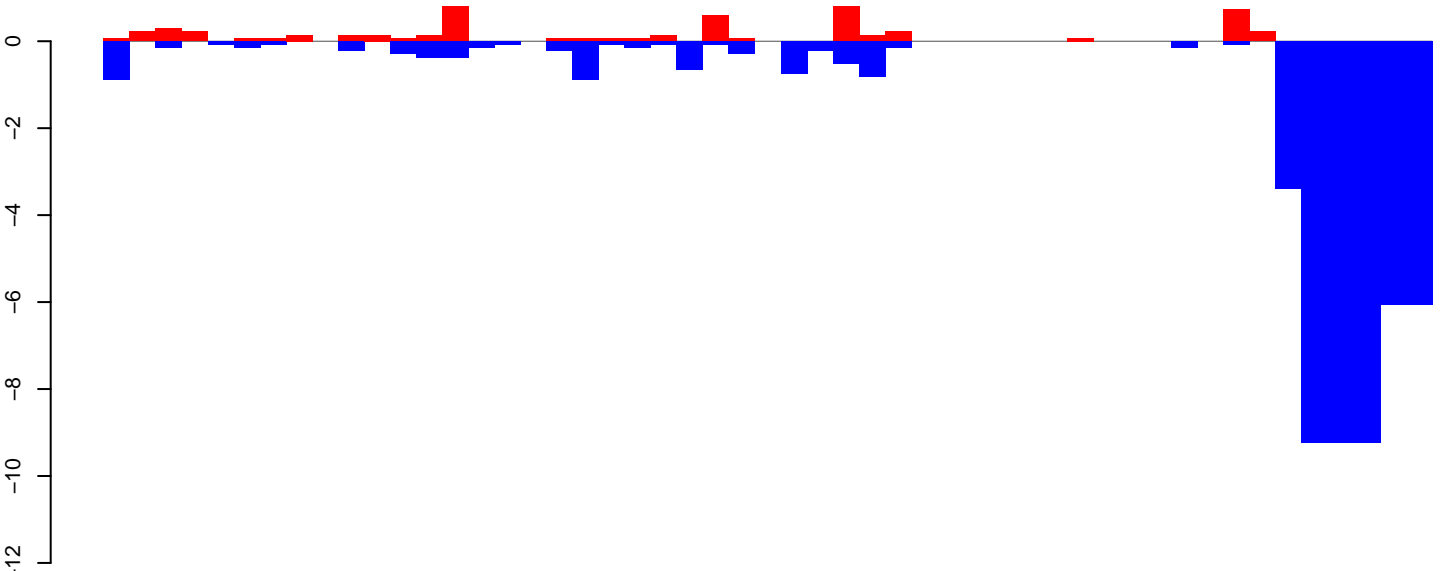


CuQuin_Testes_NL.rep



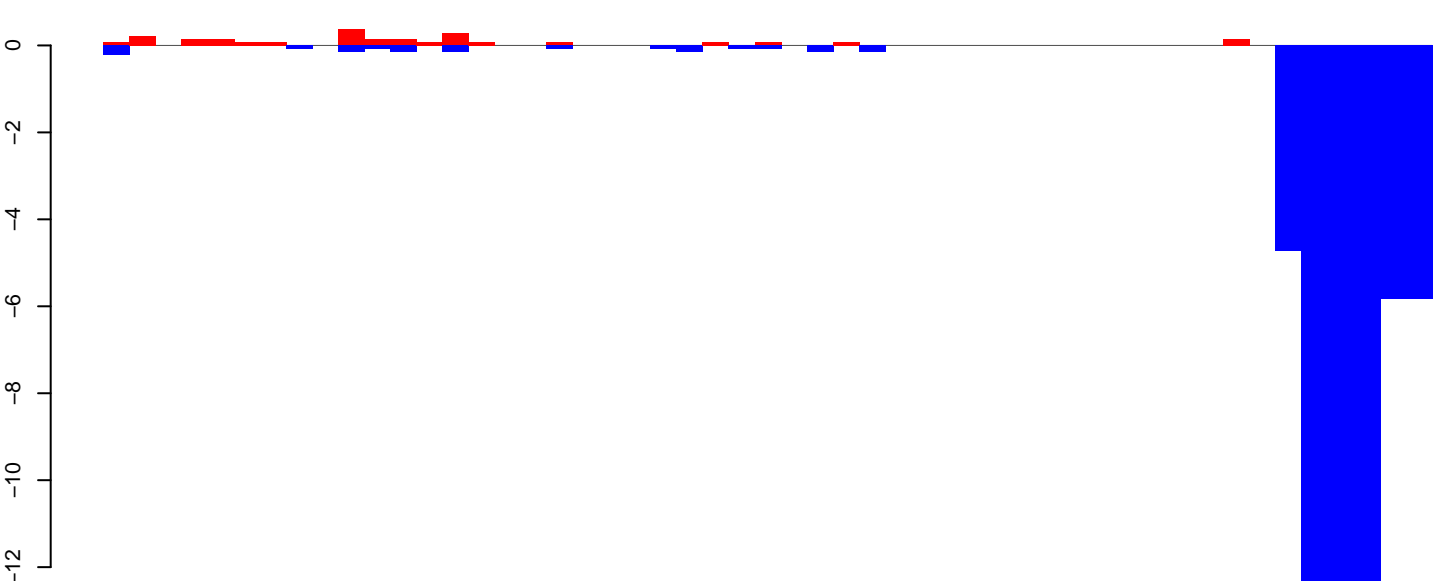
Window size=25, length=1242, TE@rnd-3_family-41@Unknown@703:1-1242

CuQuin_Testes_NL.18_23.rep



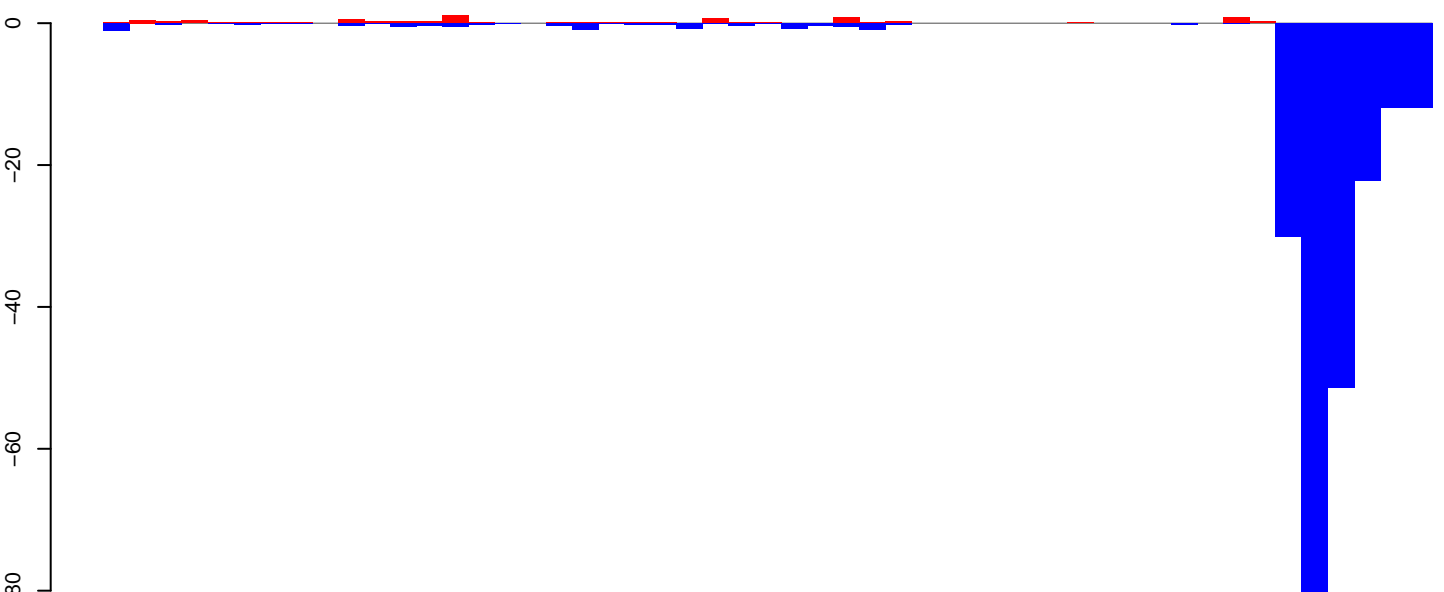
positive=6, negative=51, total=57

CuQuin_Testes_NL.24_35.rep



positive=2, negative=57, total=59

CuQuin_Testes_NL.rep

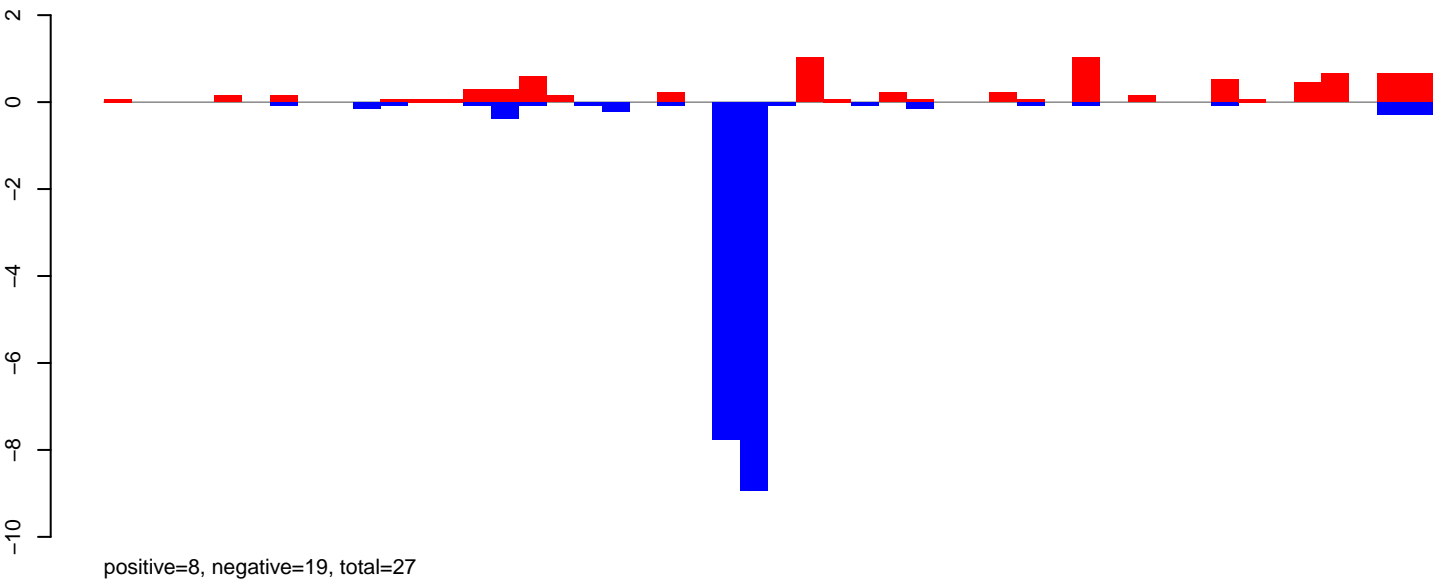


positive=8, negative=220, total=228

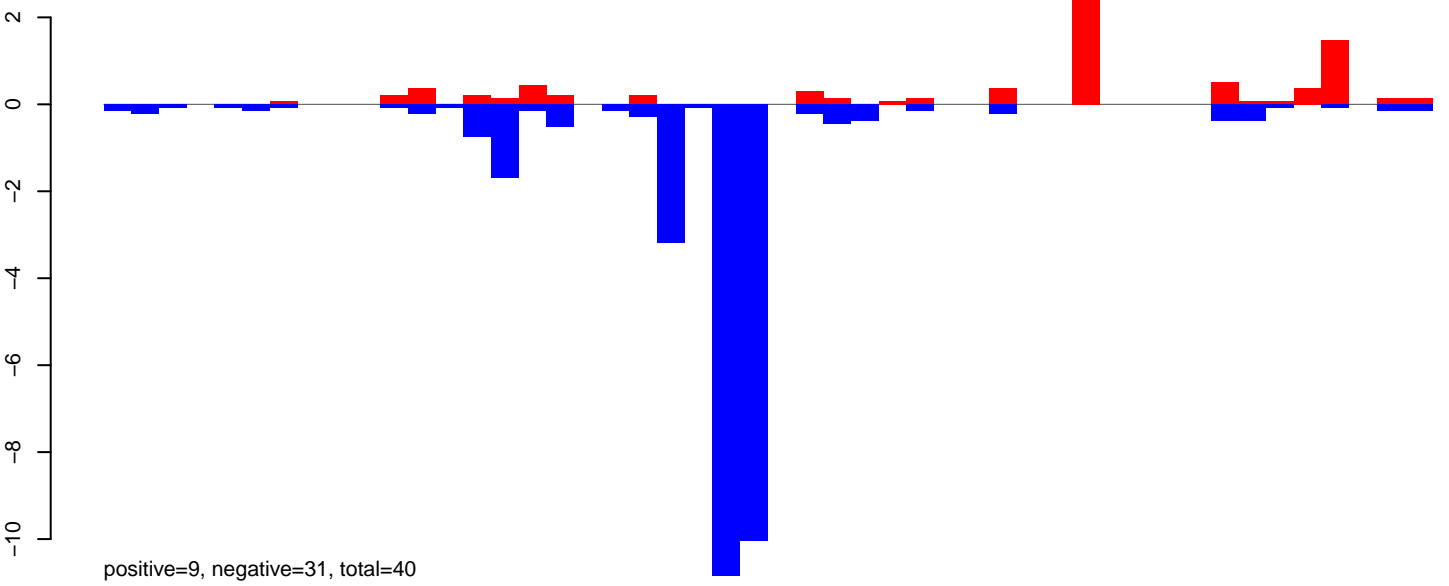
Window size=25, length=1263, TE@rnd-3_family-267@DNA_CMC-Transib@2654:1-1263

0 200 400 600 800 1000 1200

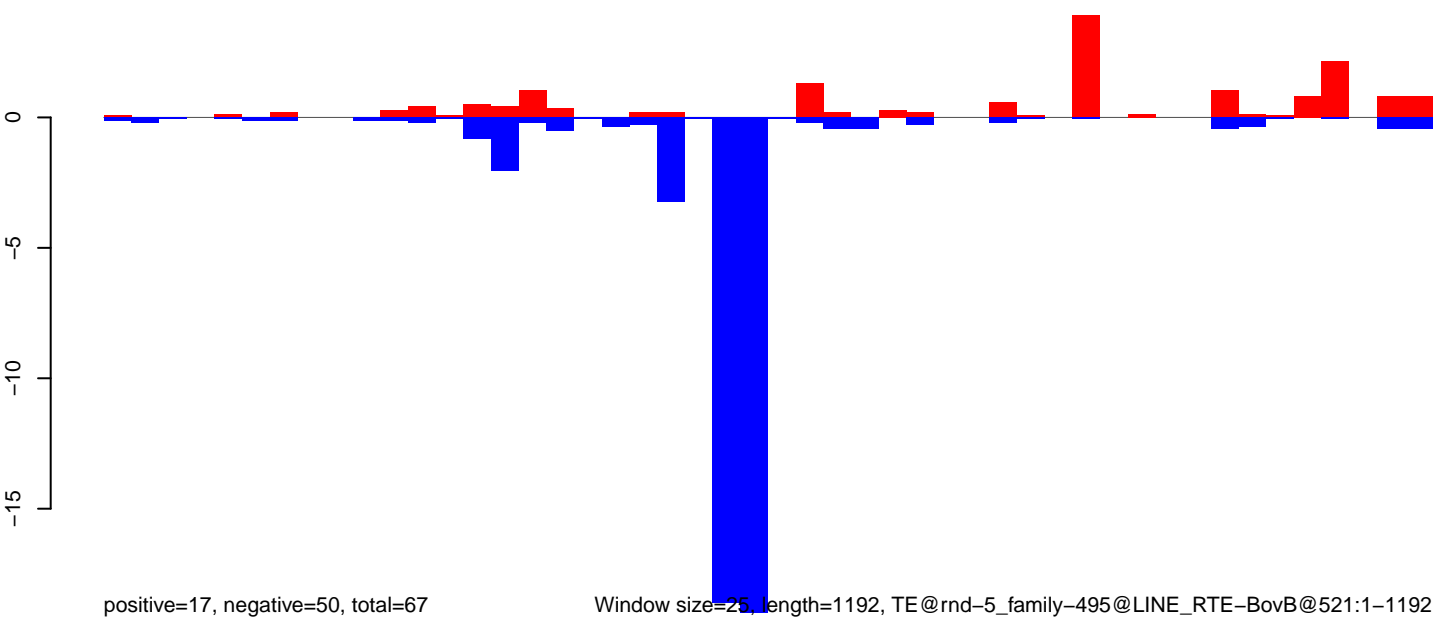
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



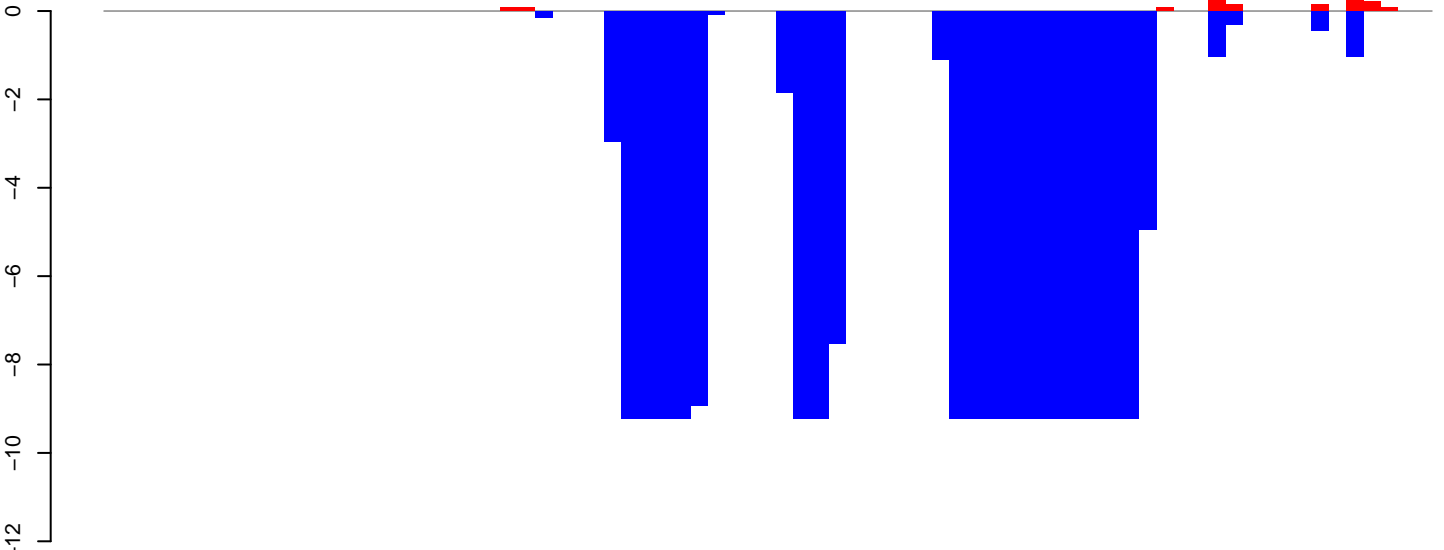
CuQuin_Testes_NL.rep



Window size=25, length=1192, TE@rnd-5_family-495@LINE_RTE-BovB@521:1-1192

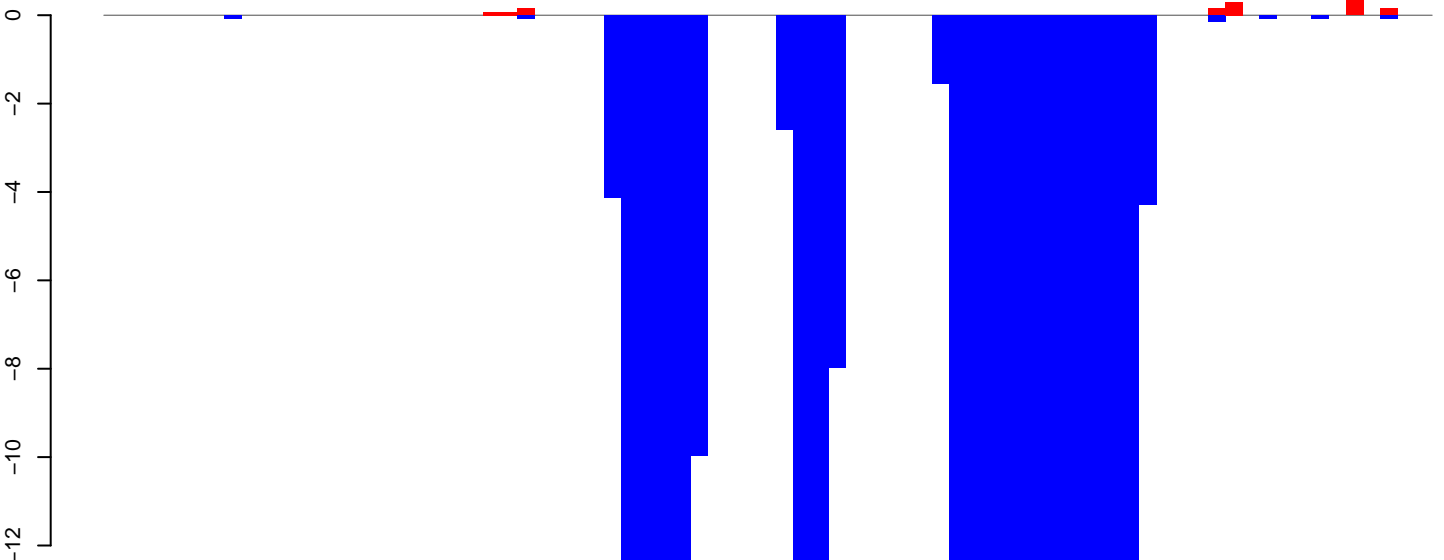
0 200 400 600 800 1000 1200

CuQuin_Testes_NL.18_23.rep



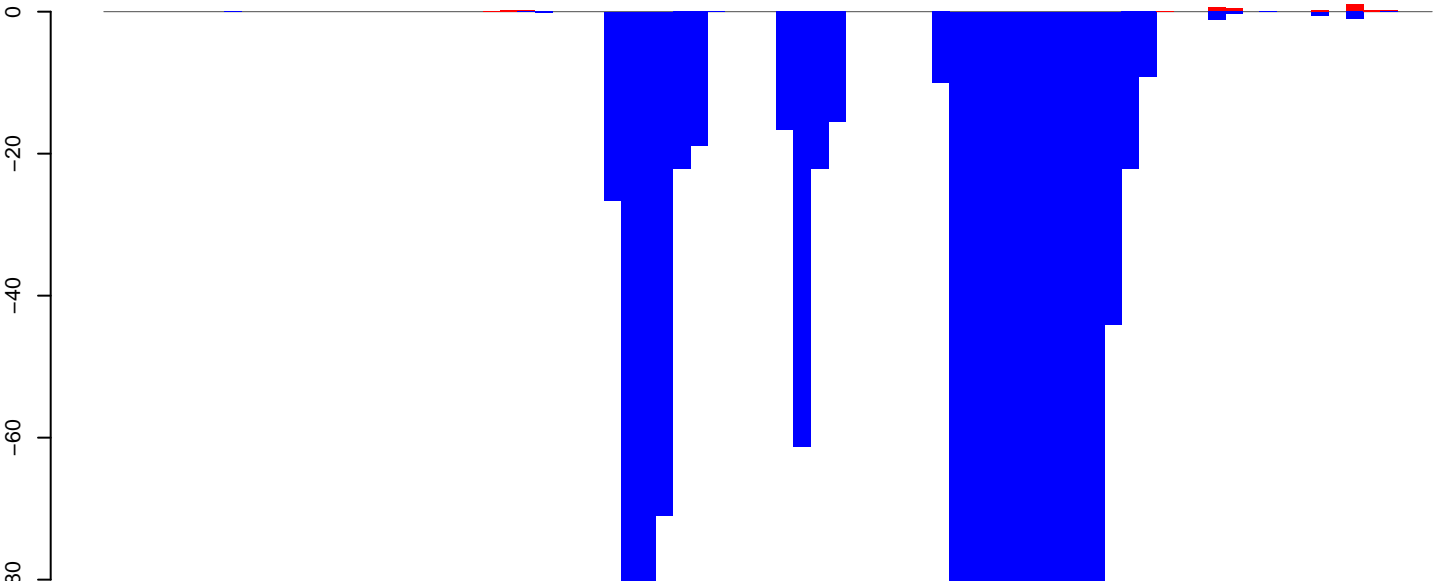
positive=2, negative=187, total=189

CuQuin_Testes_NL.24_35.rep



positive=1, negative=251, total=252

CuQuin_Testes_NL.rep

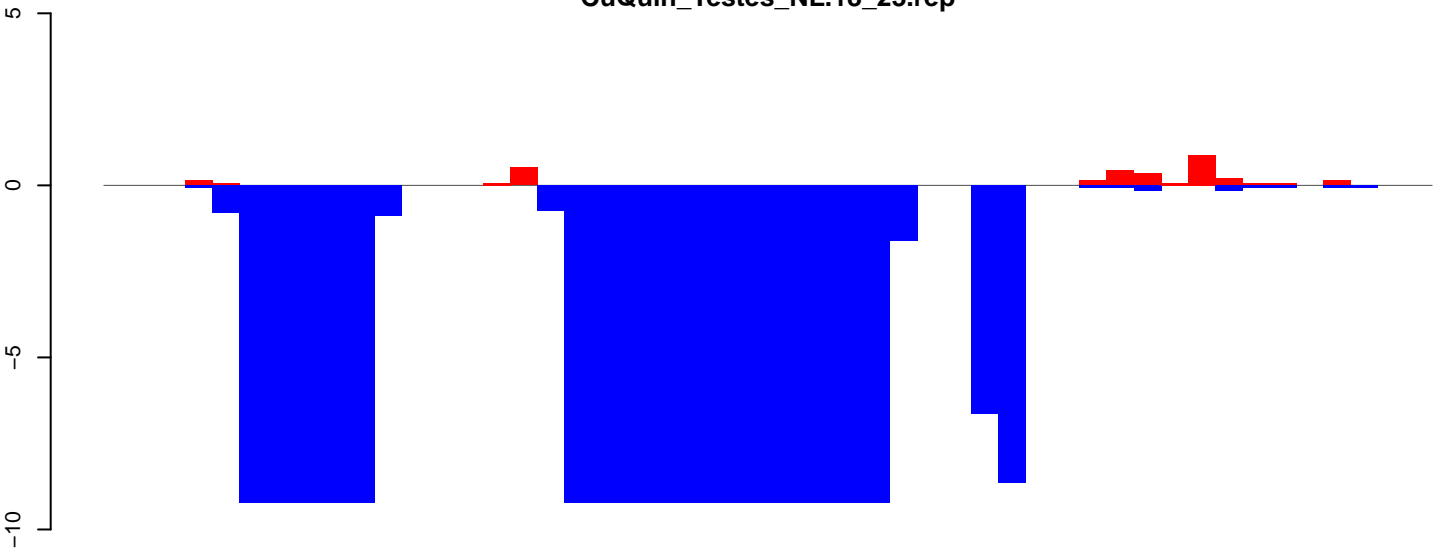


positive=3, negative=1257, total=1260

Window size=25, length=1913, TE@rnd-5_family-450@DNA_hAT@1756:1-1913

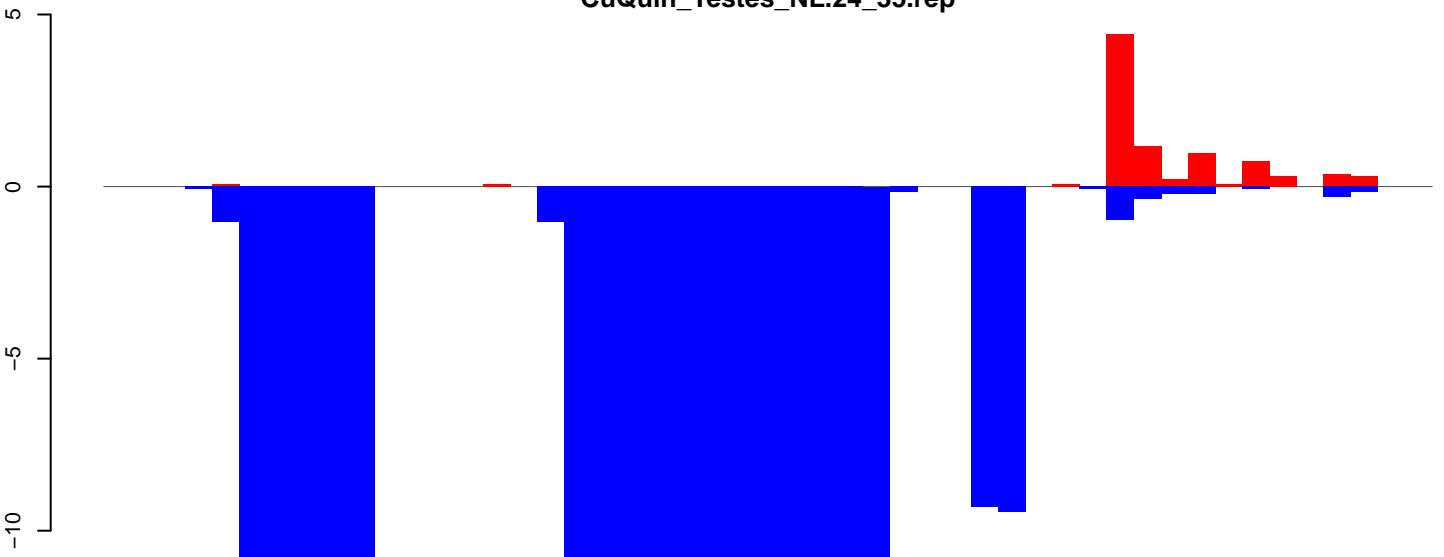
0 500 1000 1500 2000

CuQuin_Testes_NL.18_23.rep



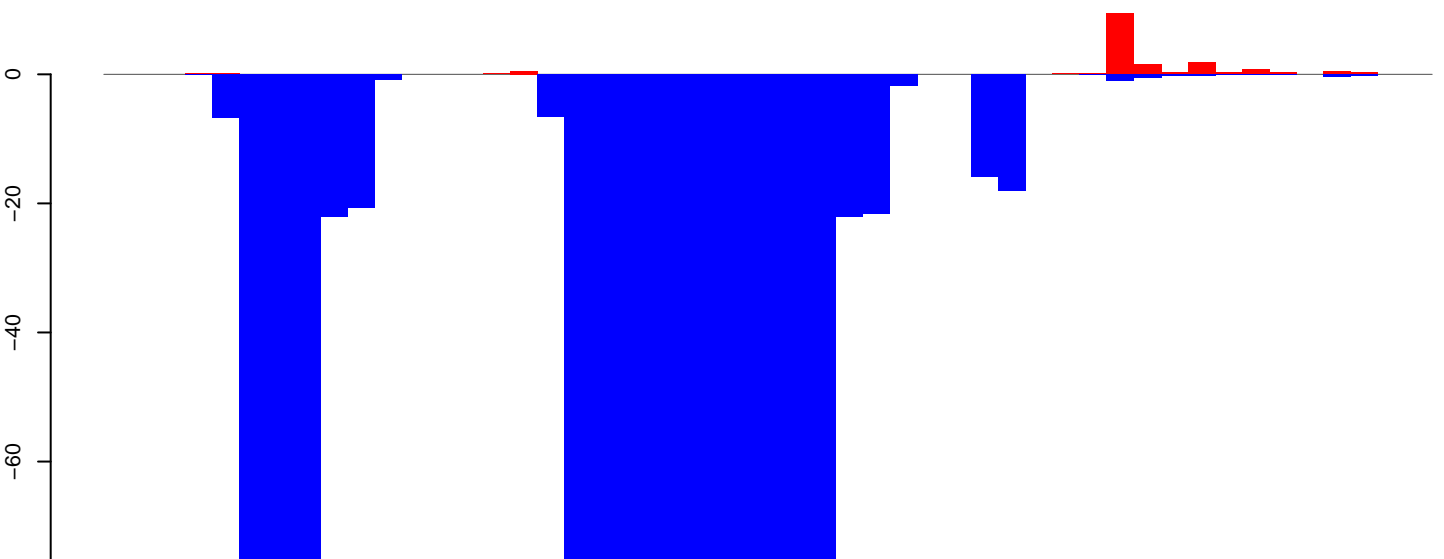
positive=3, negative=177, total=180

CuQuin_Testes_NL.24_35.rep



positive=9, negative=241, total=250

CuQuin_Testes_NL.rep

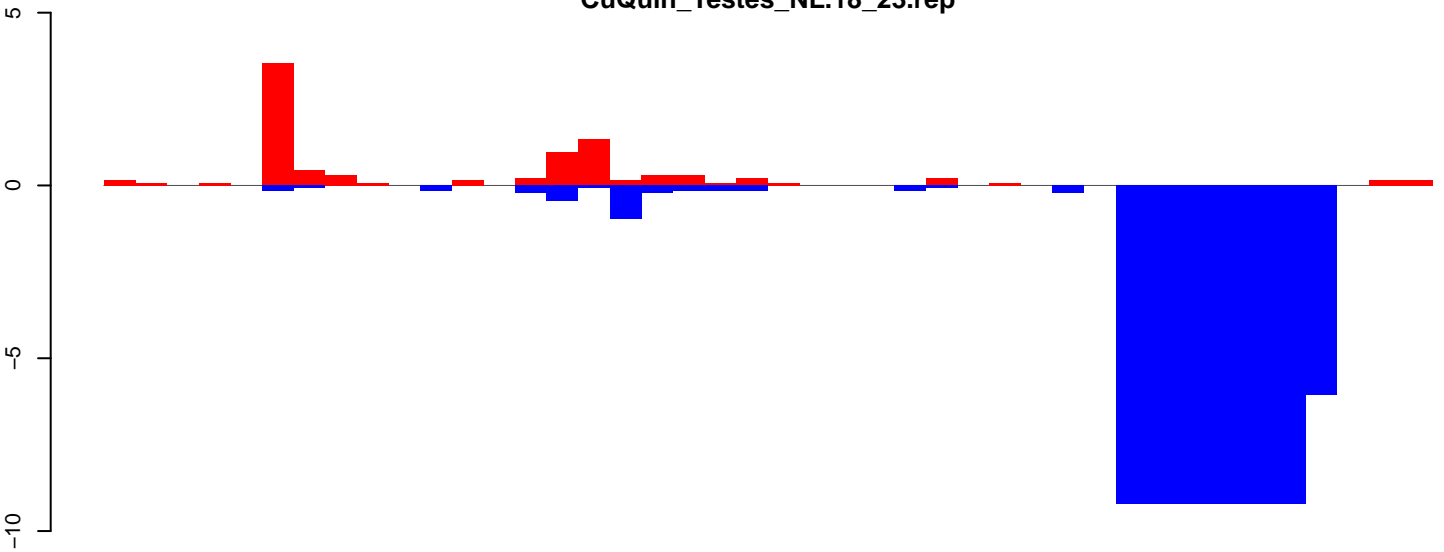


positive=17, negative=1215, total=1232

Window size=25, length=1212, TE@rnd-4_family-681@DNA_Sola-2@3581:1-1212

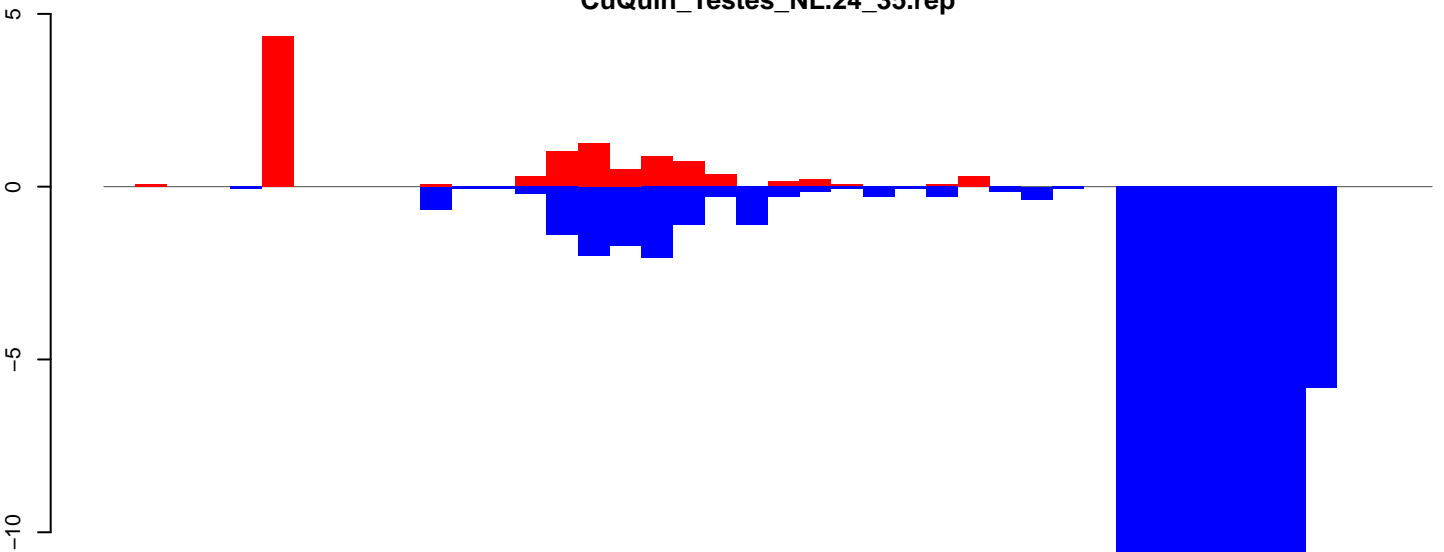
0 200 400 600 800 1000 1200

CuQuin_Testes_NL.18_23.rep



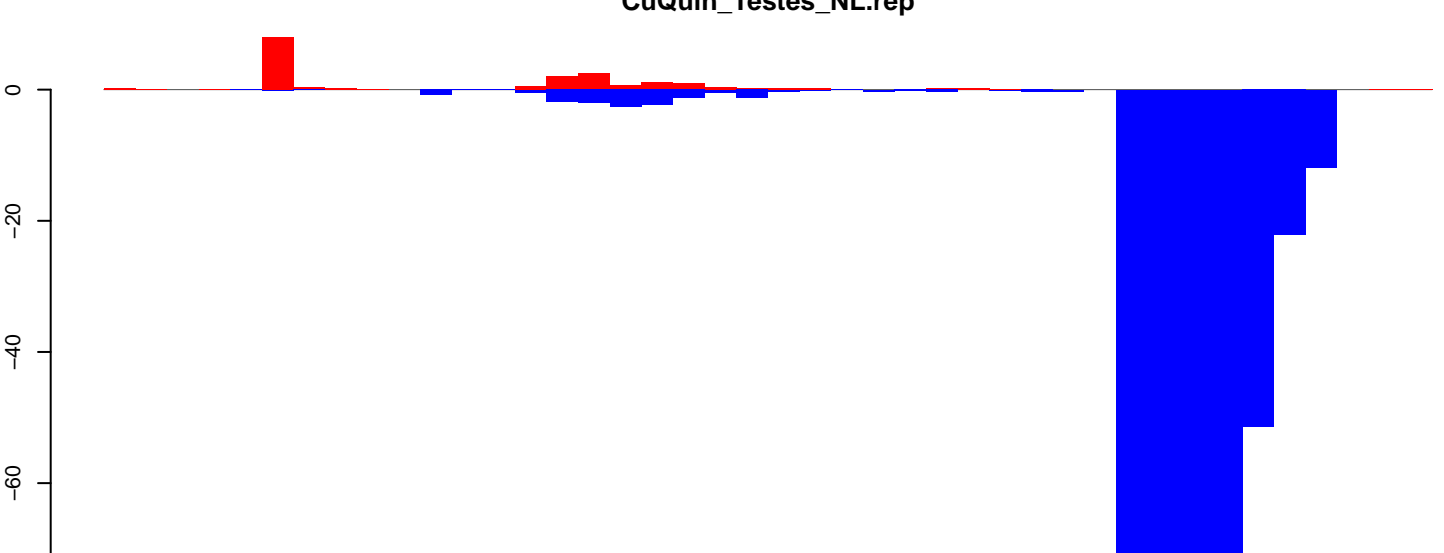
positive=9, negative=65, total=74

CuQuin_Testes_NL.24_35.rep



positive=10, negative=96, total=106

CuQuin_Testes_NL.rep

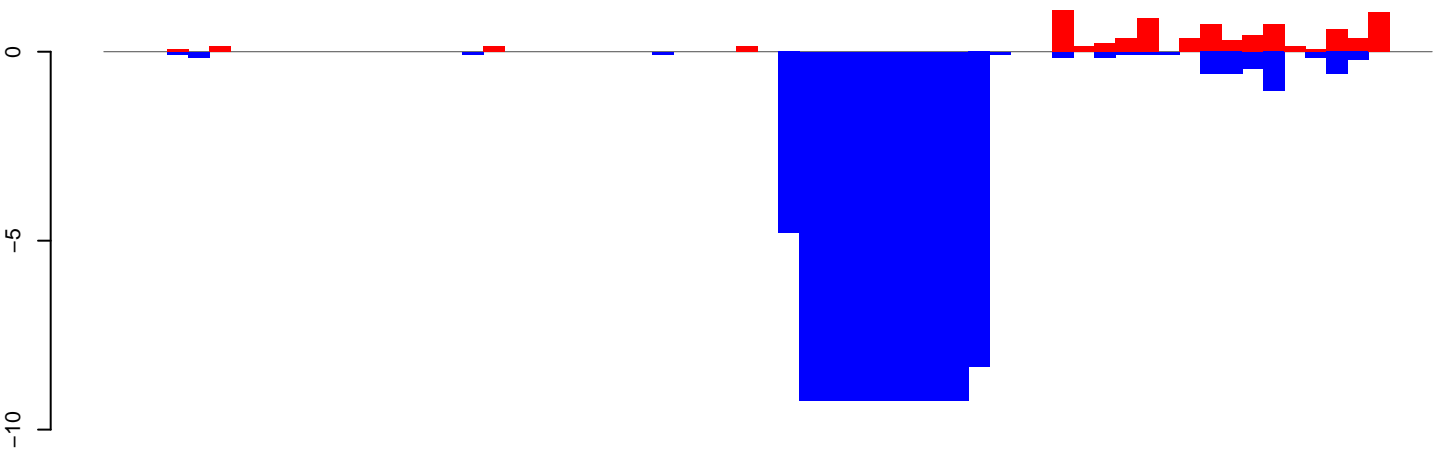


positive=20, negative=433, total=453

Window size=25, length=1033, TE@rnd-4_family-632@Satellite@3595:1-1033

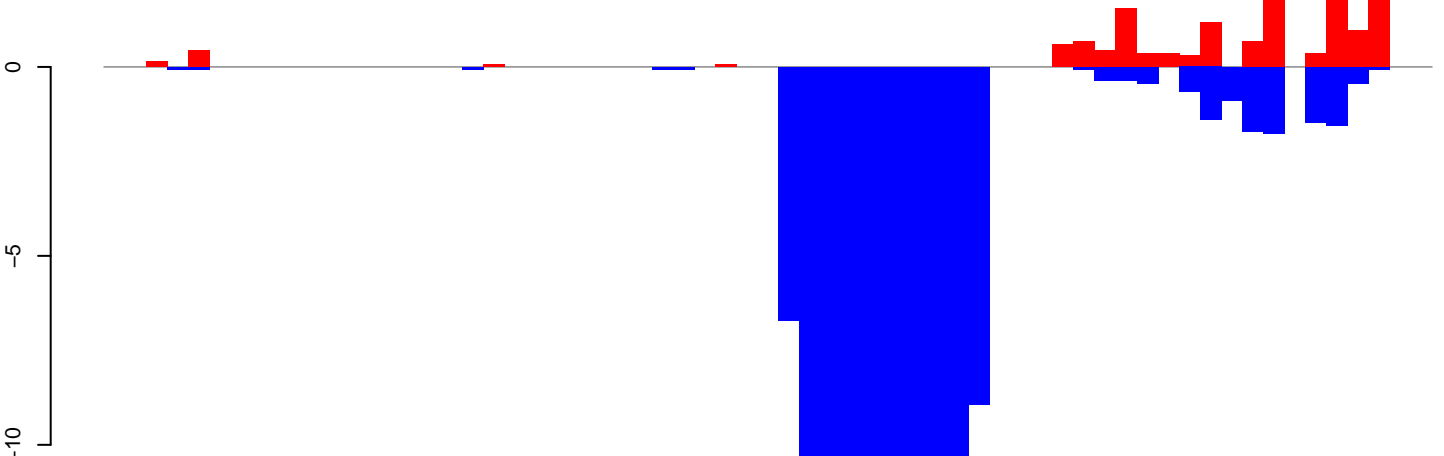
0 200 400 600 800 1000

CuQuin_Testes_NL.18_23.rep



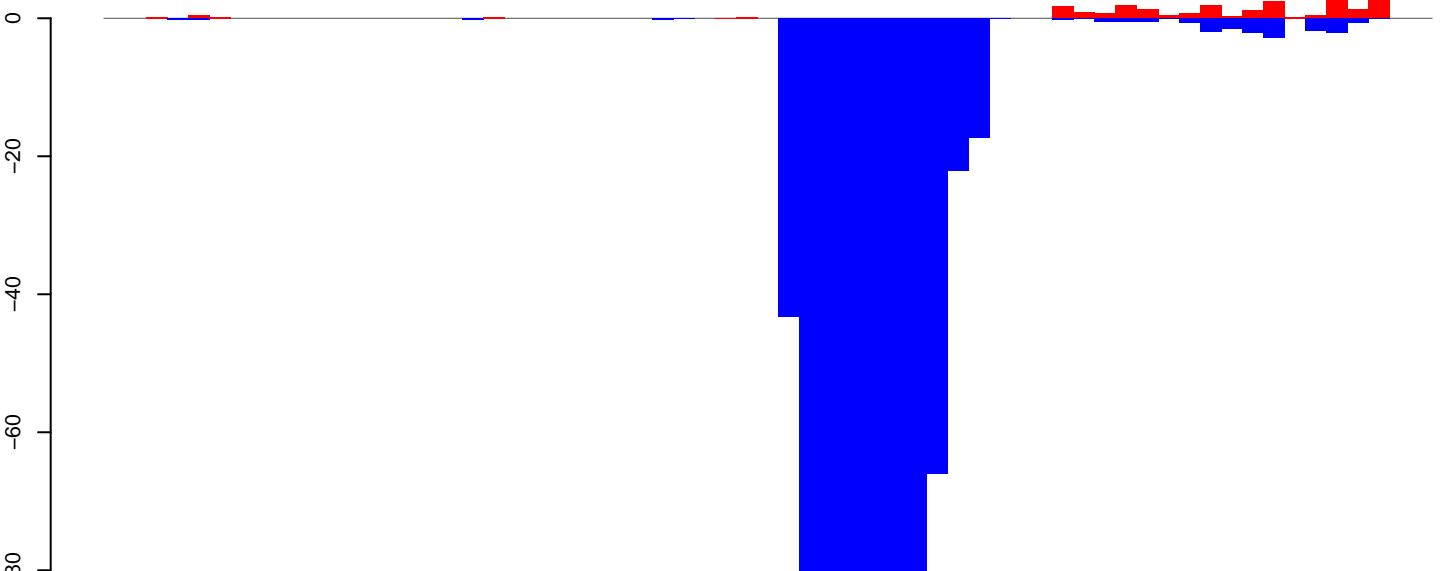
positive=8, negative=92, total=100

CuQuin_Testes_NL.24_35.rep



positive=15, negative=131, total=146

CuQuin_Testes_NL.rep

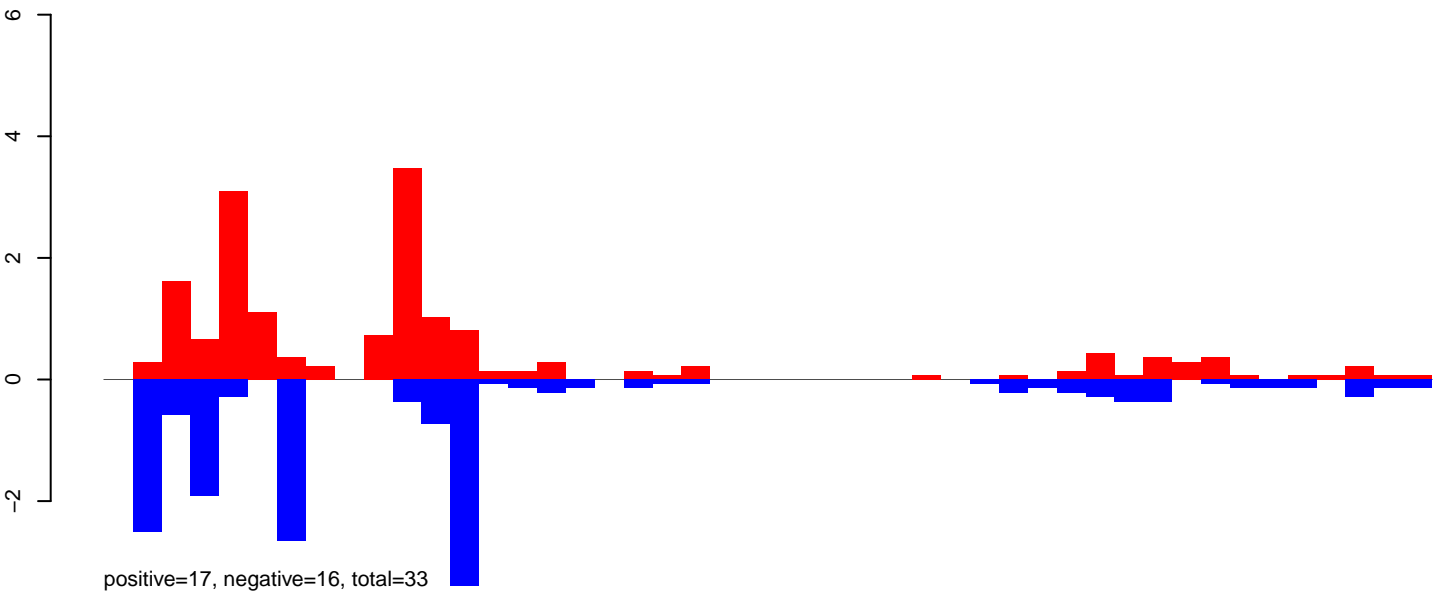


positive=23, negative=663, total=687

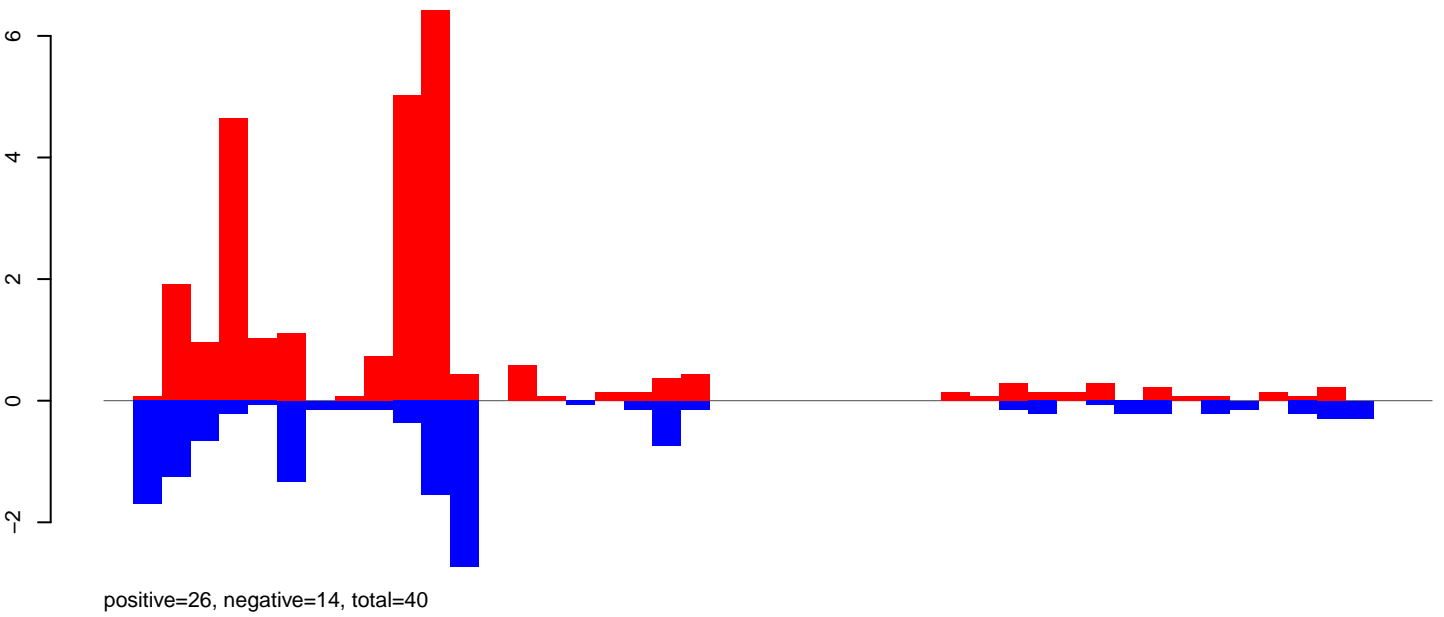
Window size=25, length=1573, TE@rnd-4_family-235@SINE_tRNA-Deu@2913:1-1573

0 500 1000 1500

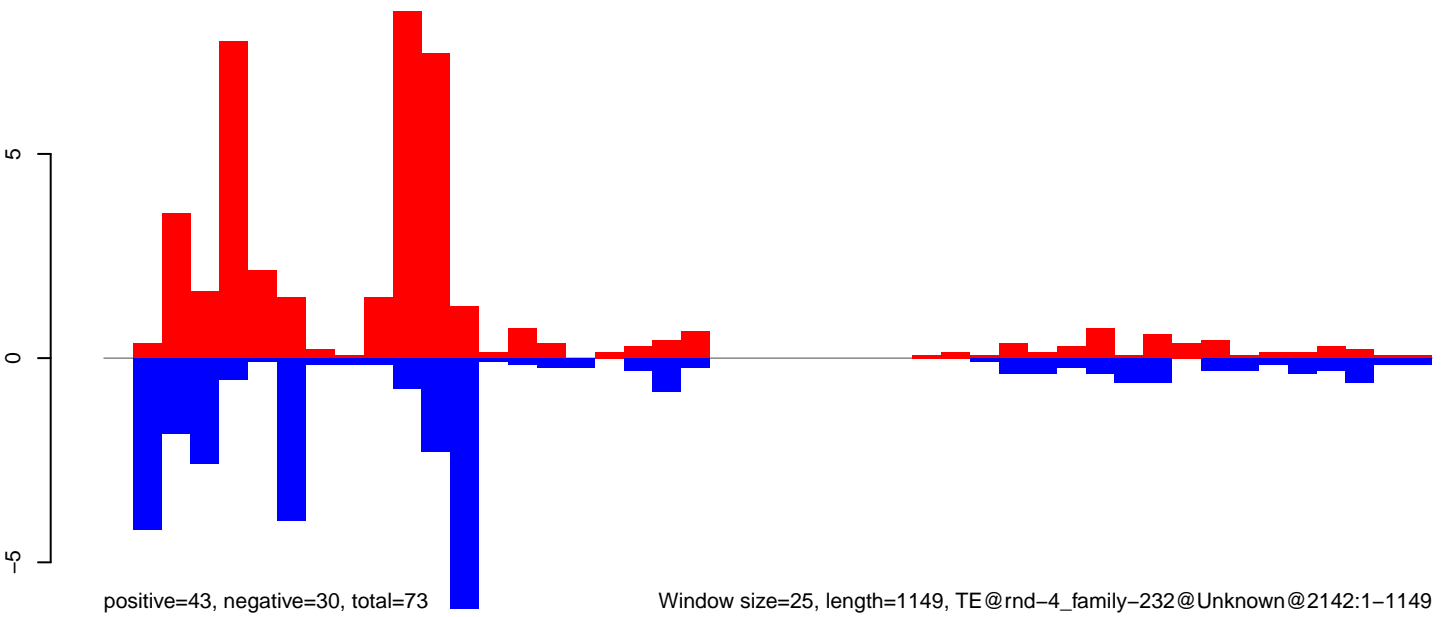
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

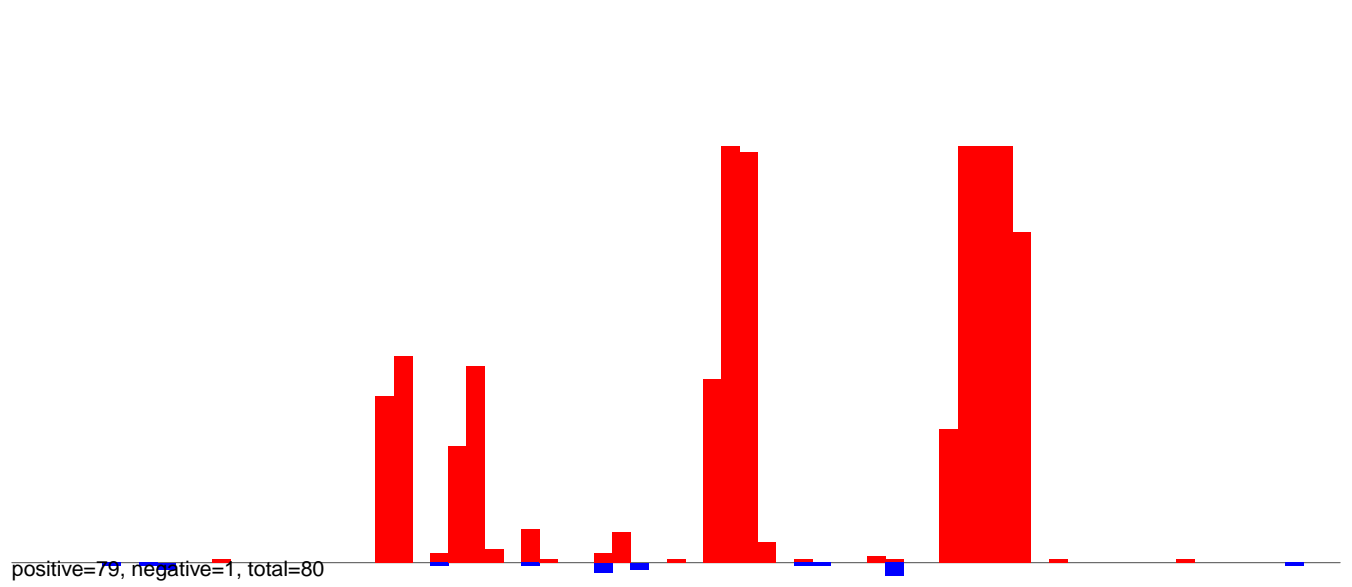


CuQuin_Testes_NL.rep

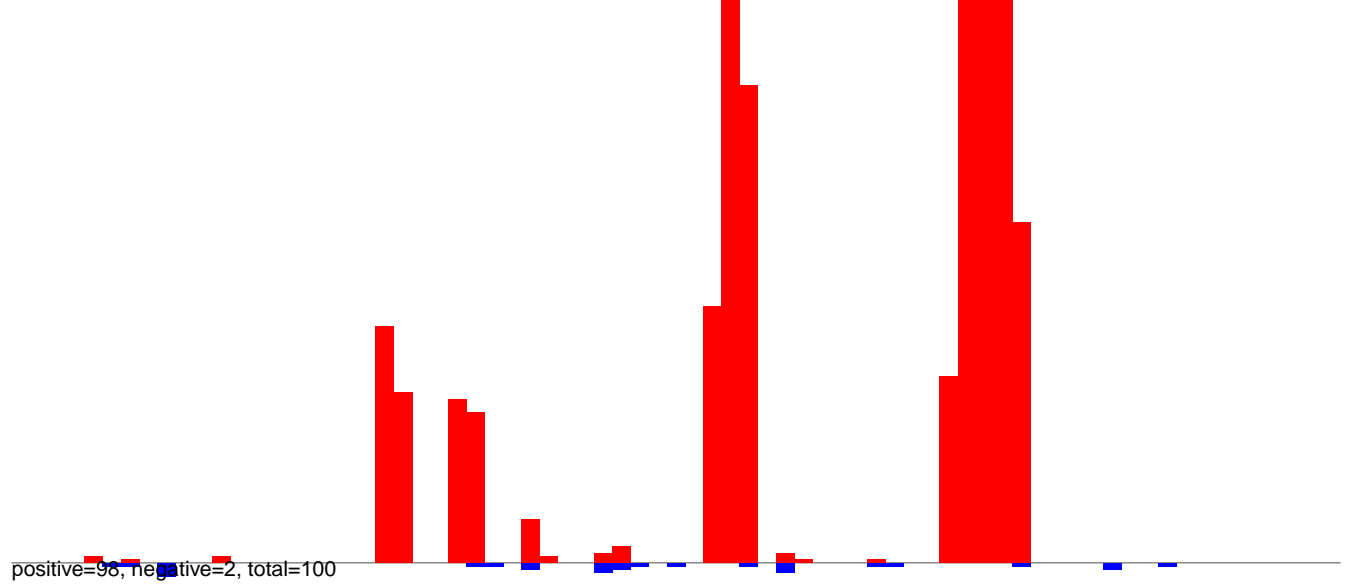


Window size=25, length=1149, TE@rnd-4_family-232@Unknown@2142:1-1149

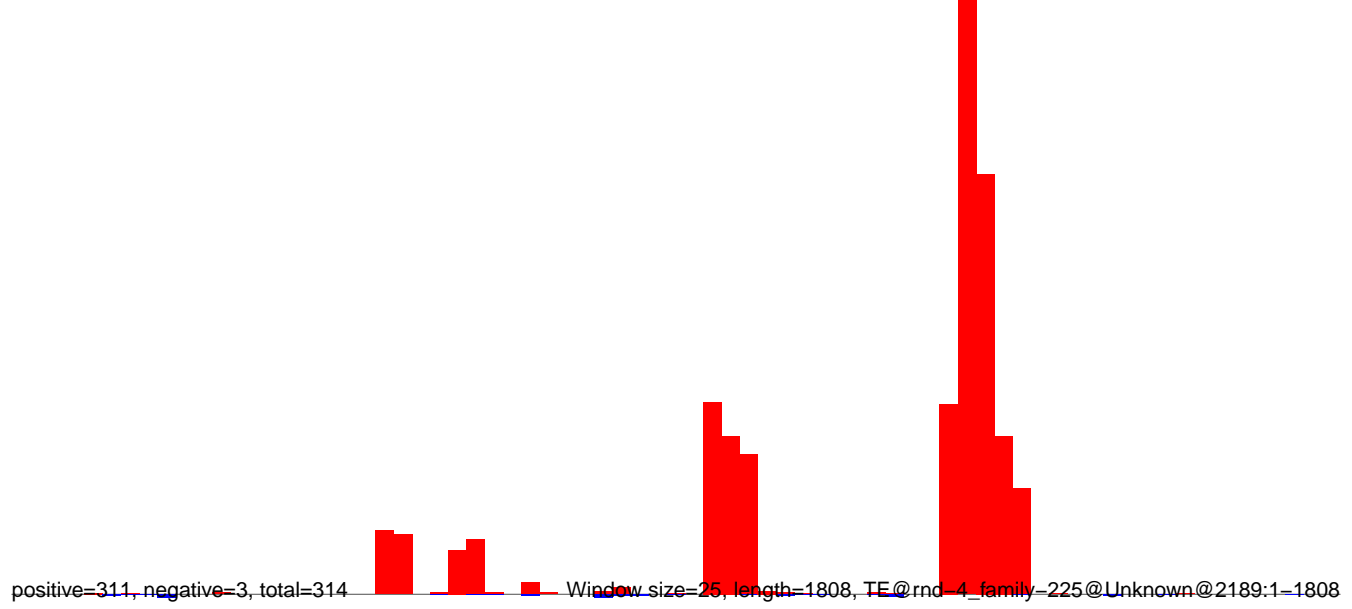
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

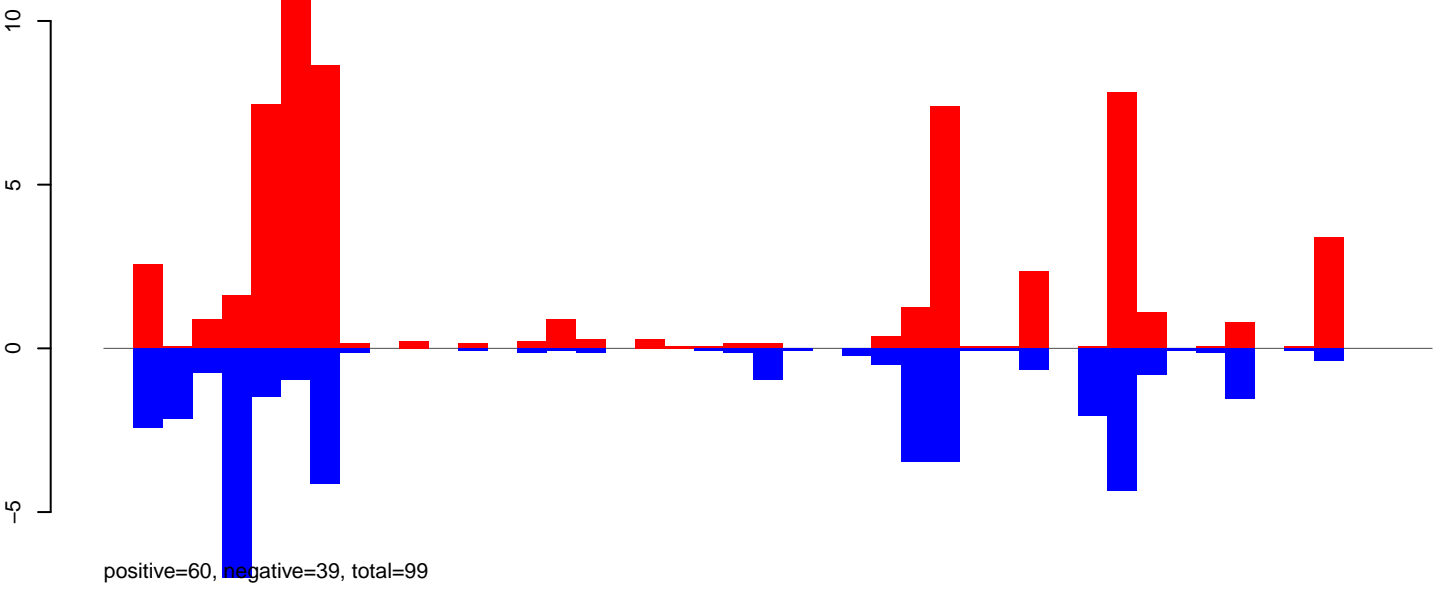


CuQuin_Testes_NL.rep

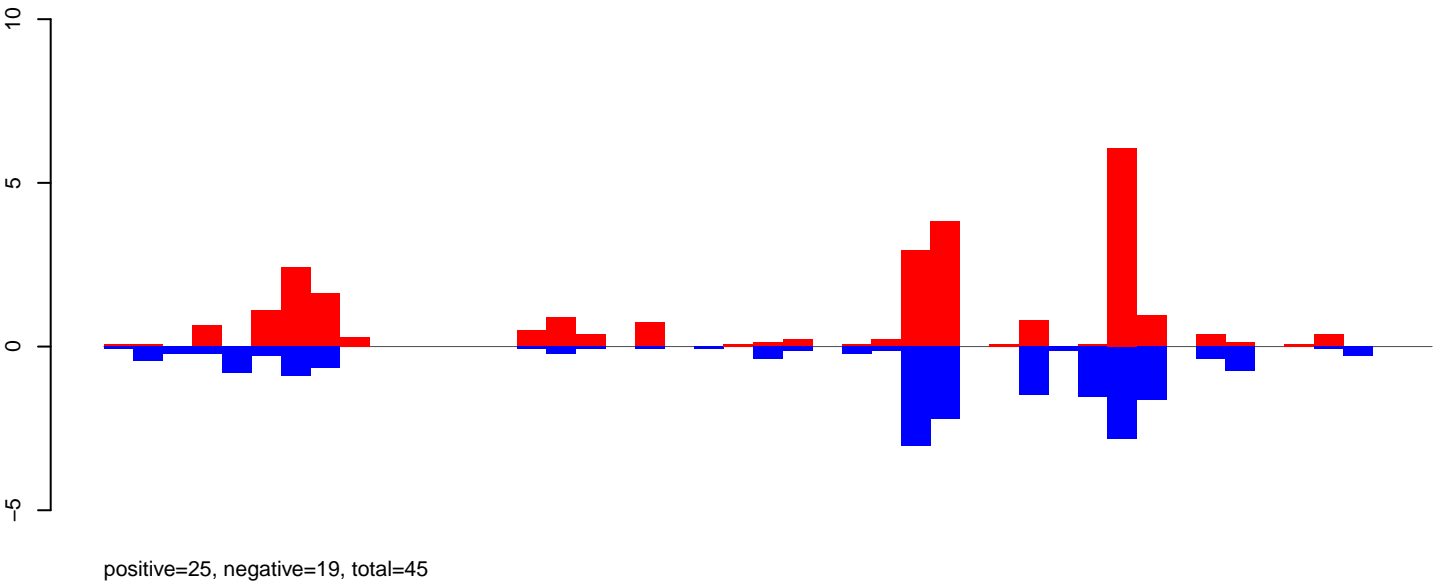


Window size=25, length=1808, TE@rnd=4, family=225@Unknown@2189:1-1808

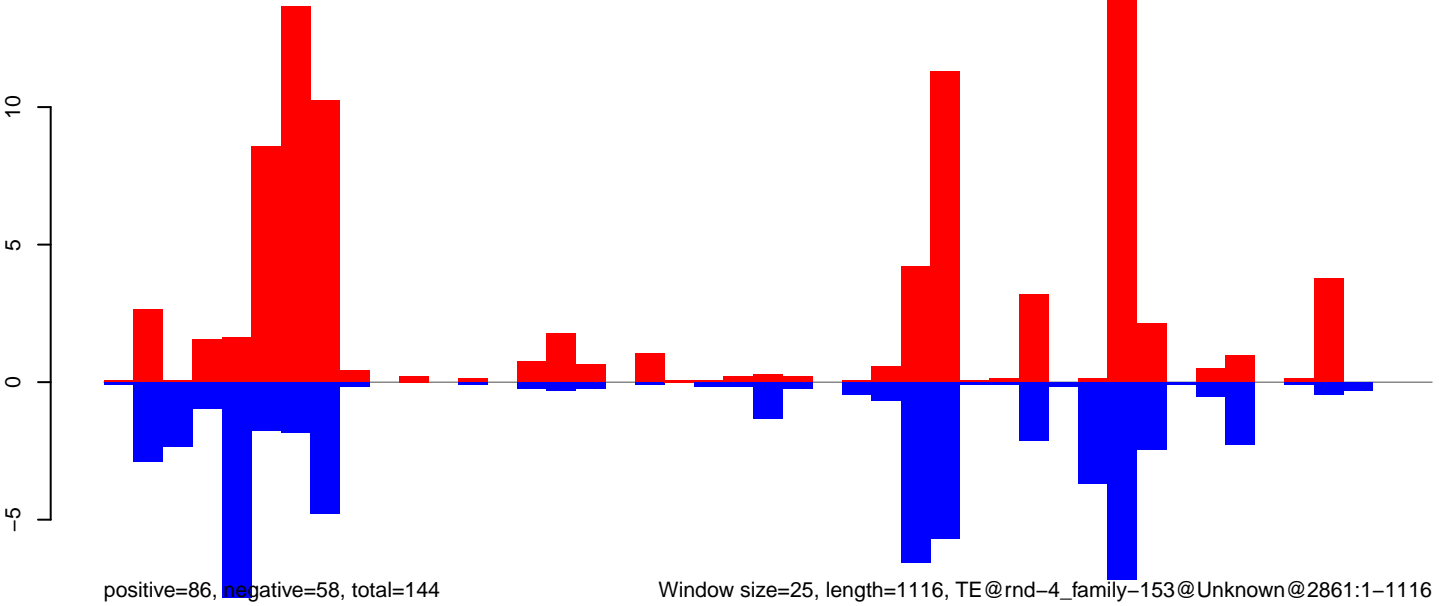
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

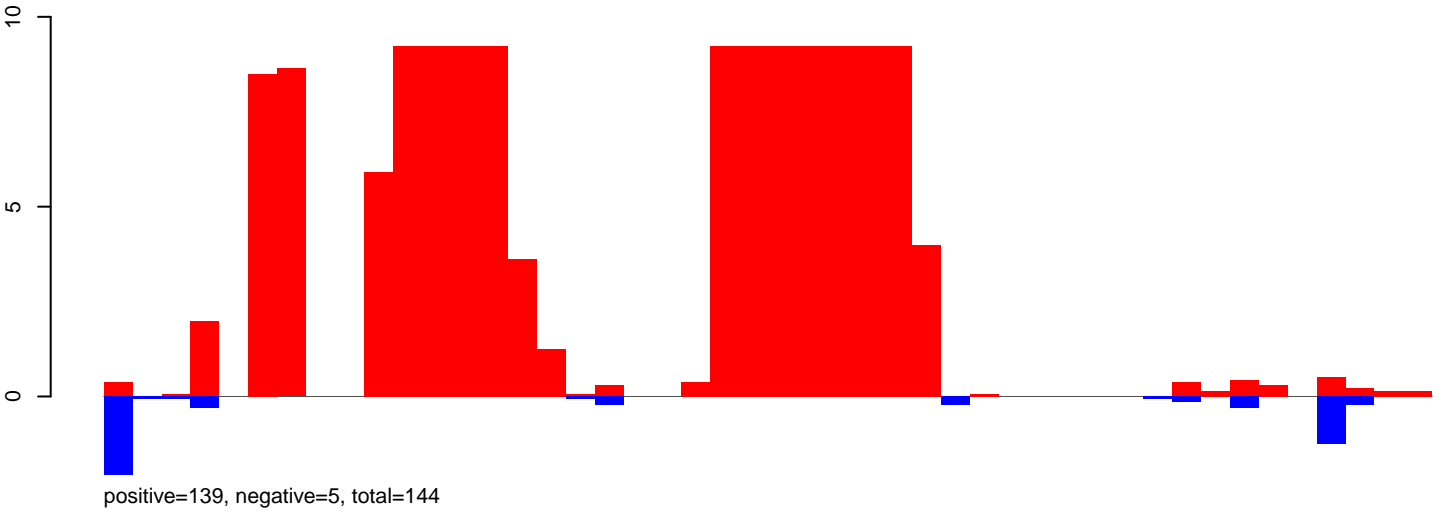


CuQuin_Testes_NL.rep

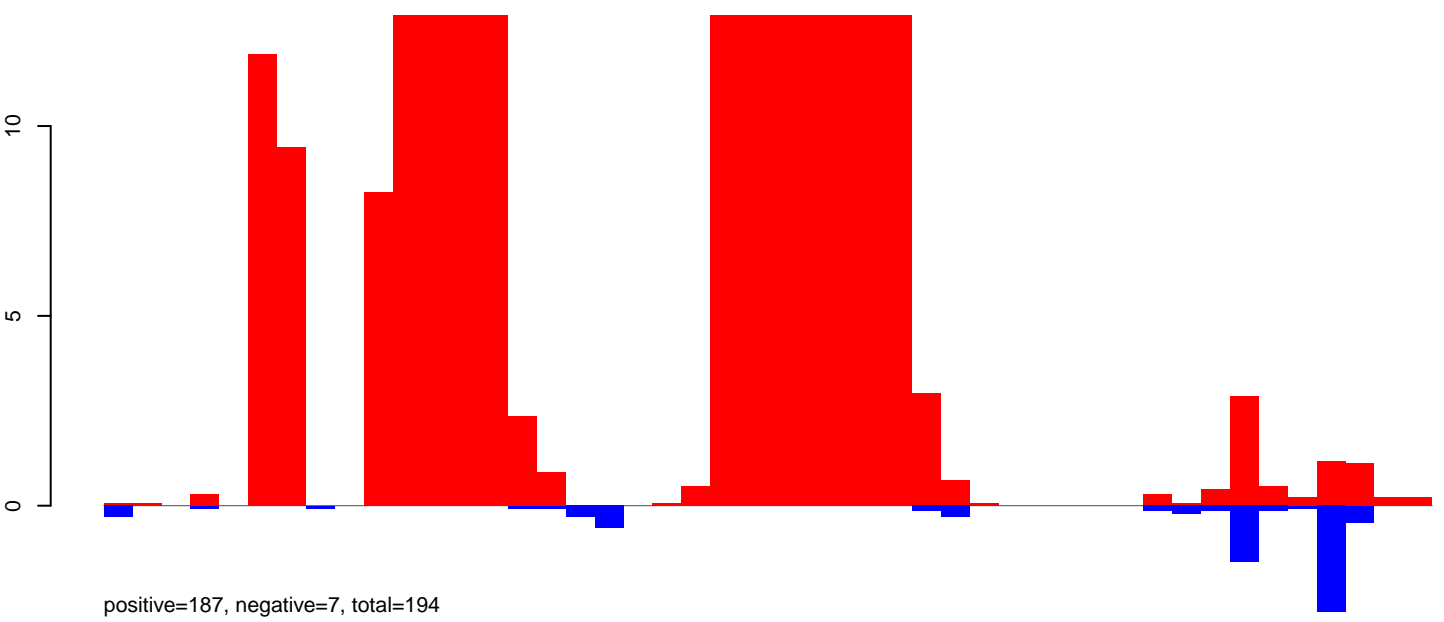


Window size=25, length=1116, TE@rnd-4_family-153@Unknown@2861:1-1116

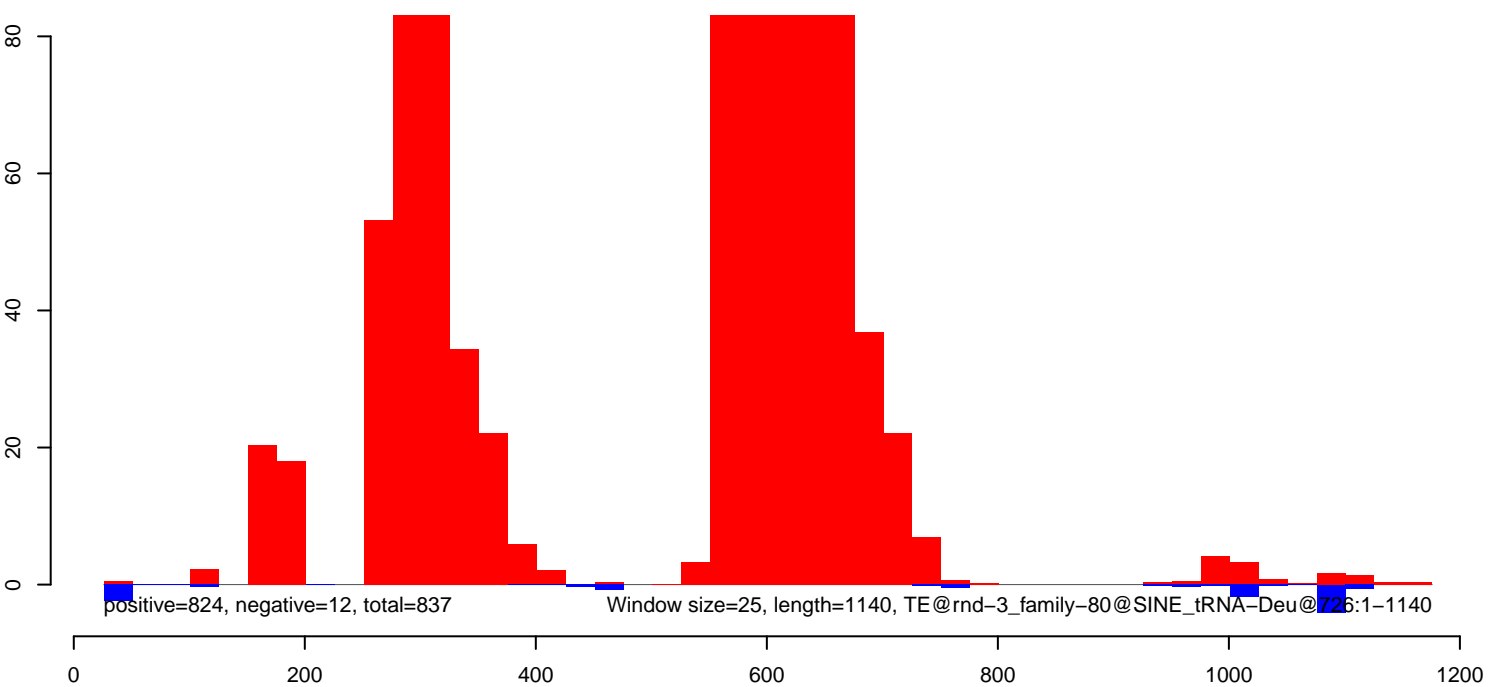
CuQuin_Testes_NL.18_23.rep



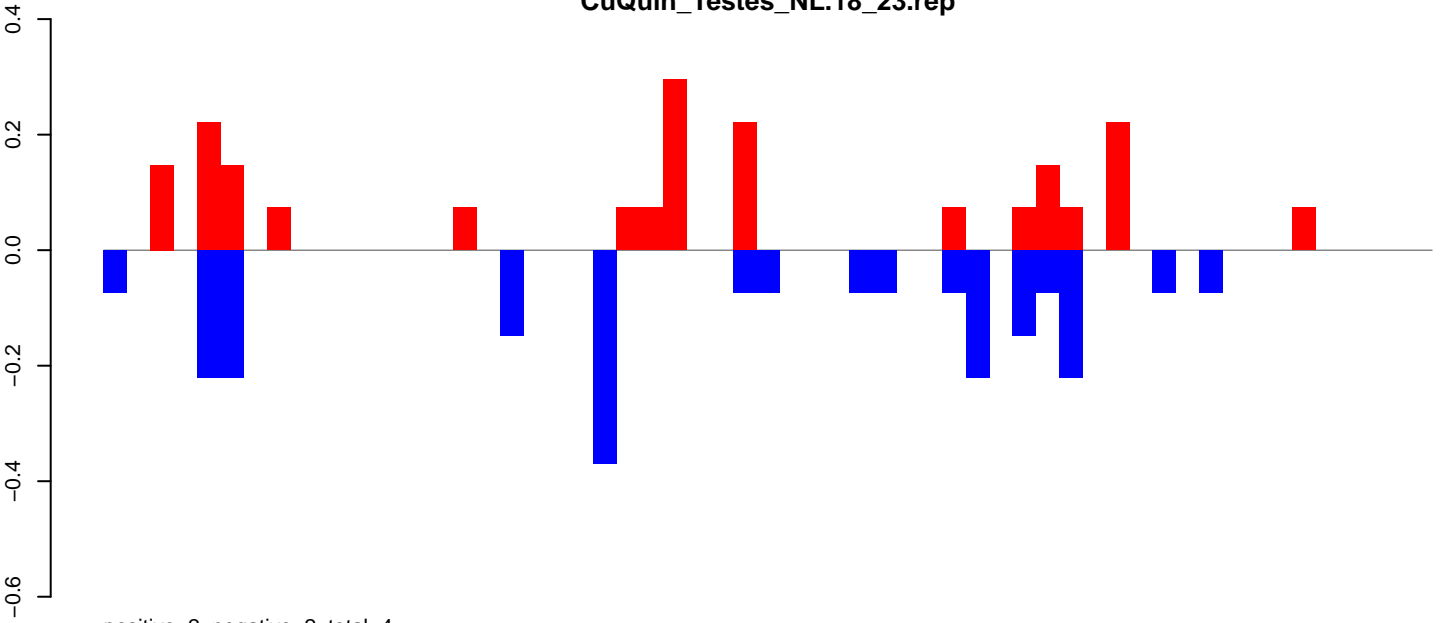
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

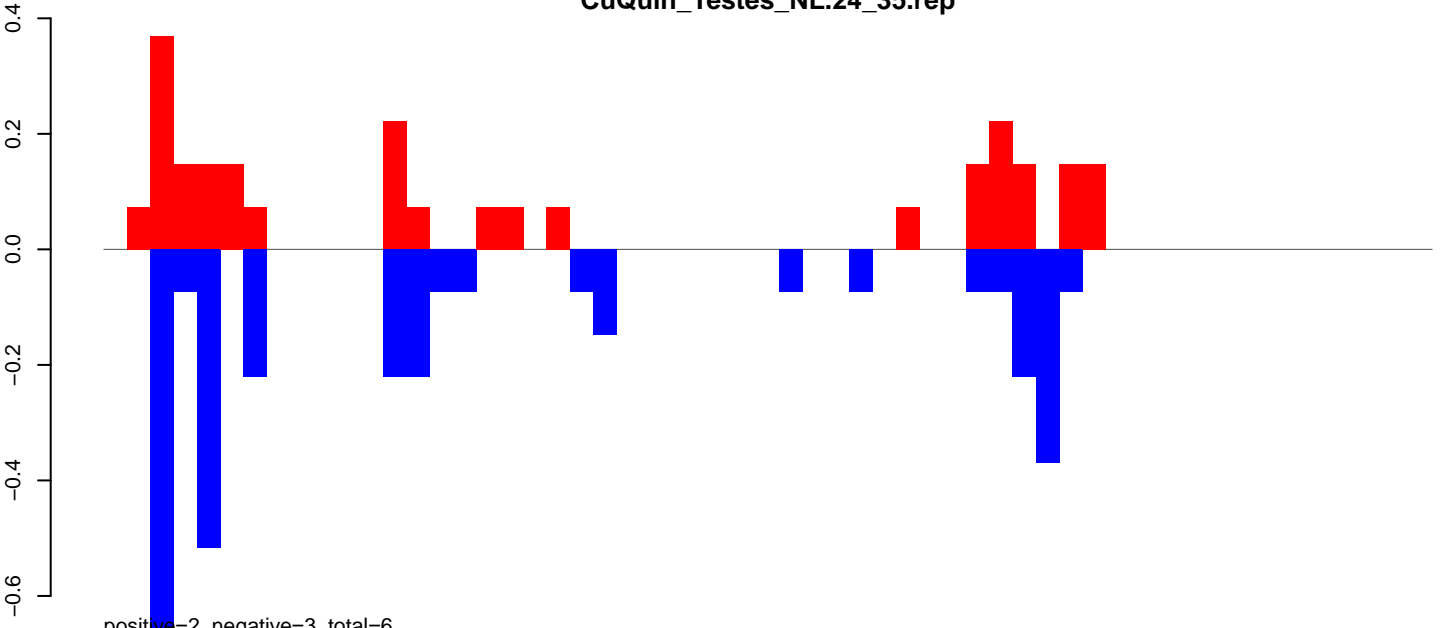


CuQuin_Testes_NL.18_23.rep



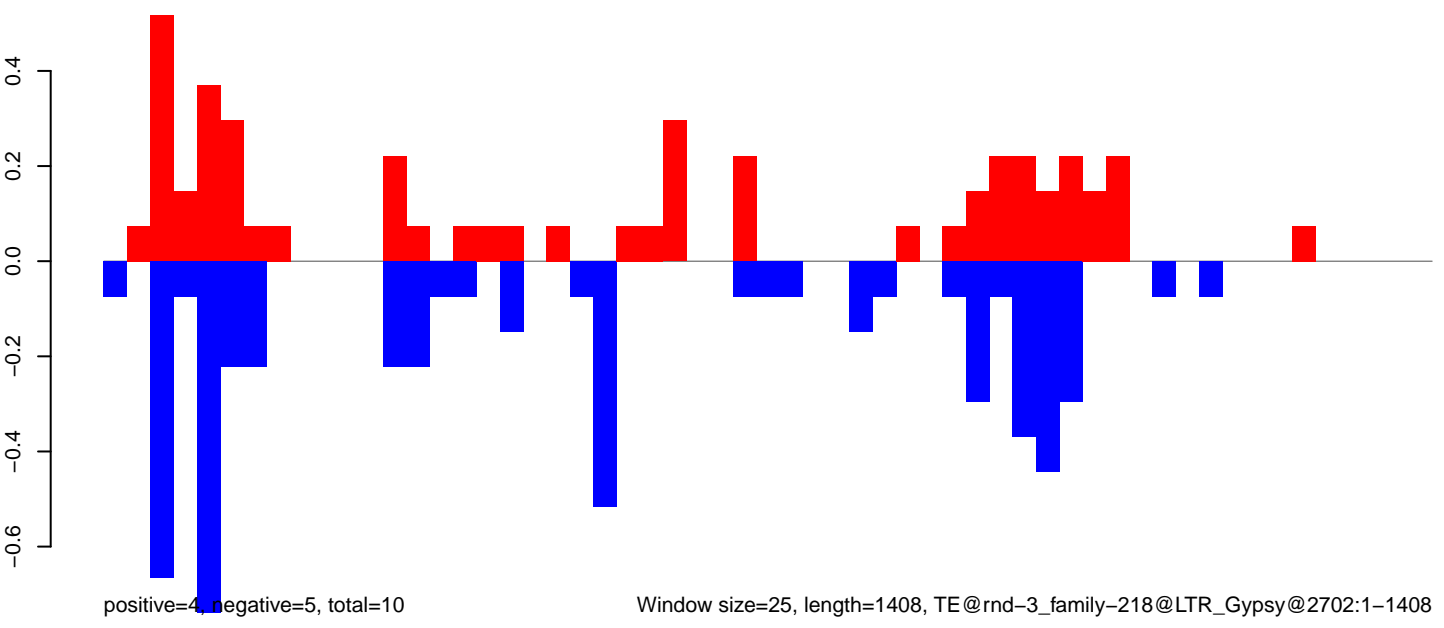
positive=2, negative=2, total=4

CuQuin_Testes_NL.24_35.rep



positive=2, negative=3, total=6

CuQuin_Testes_NL.rep

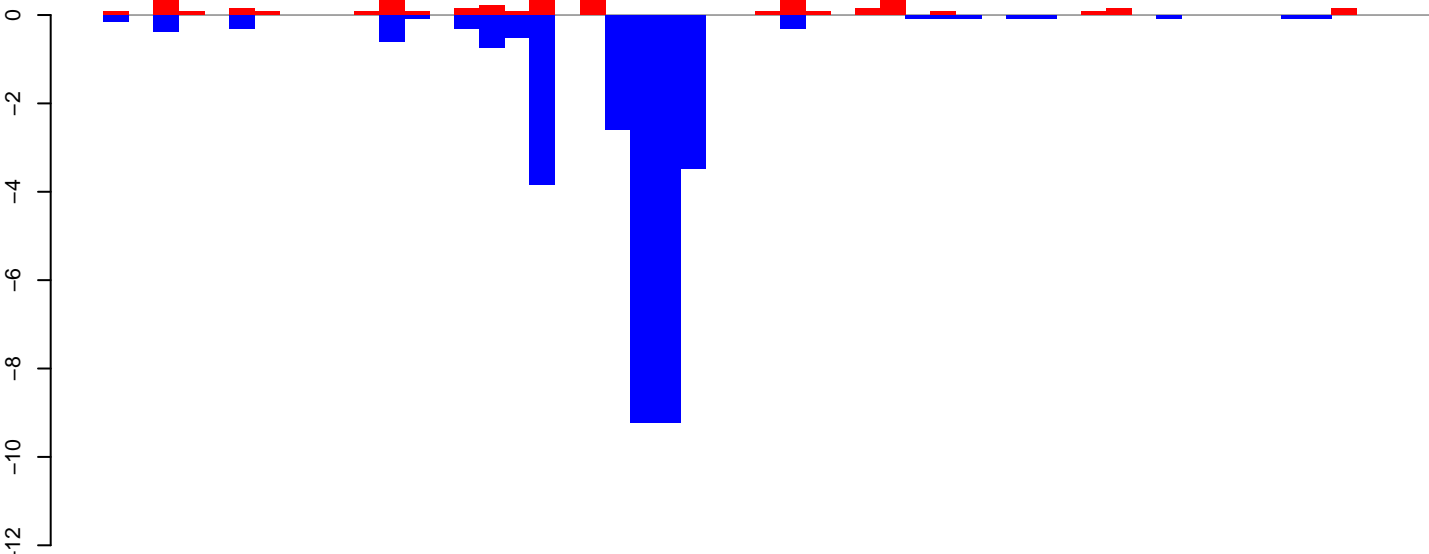


positive=4, negative=5, total=10

Window size=25, length=1408, TE@rnd-3_family-218@LTR_Gypsy@2702:1-1408

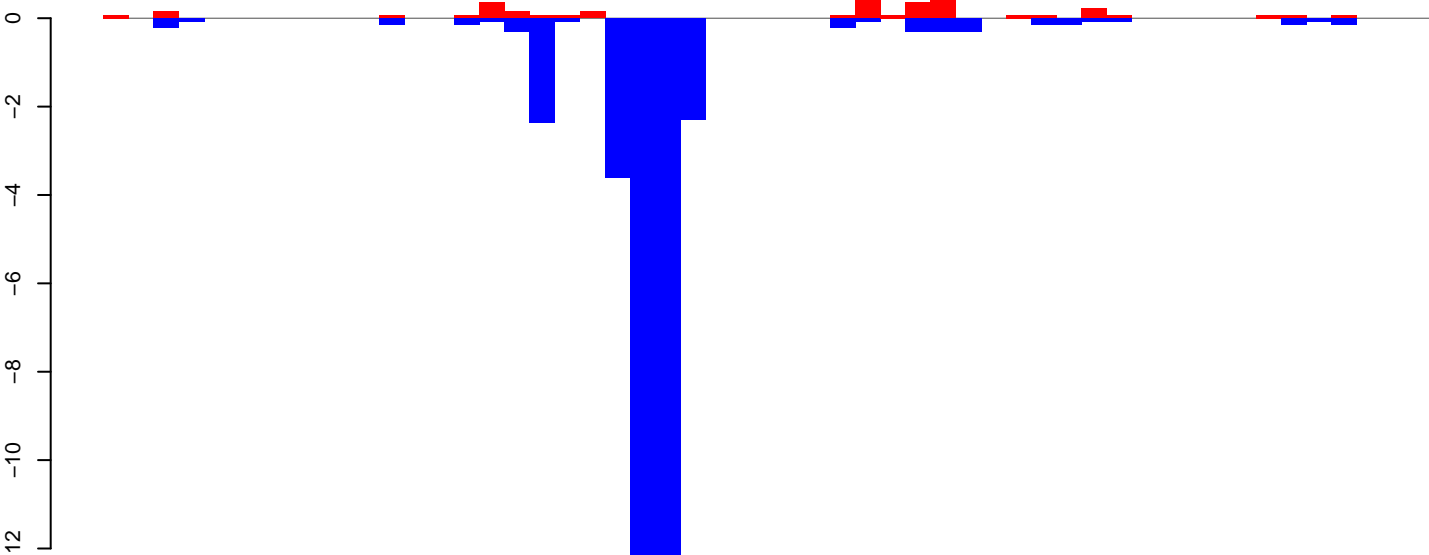
0 200 400 600 800 1000 1200 1400

CuQuin_Testes_NL.18_23.rep



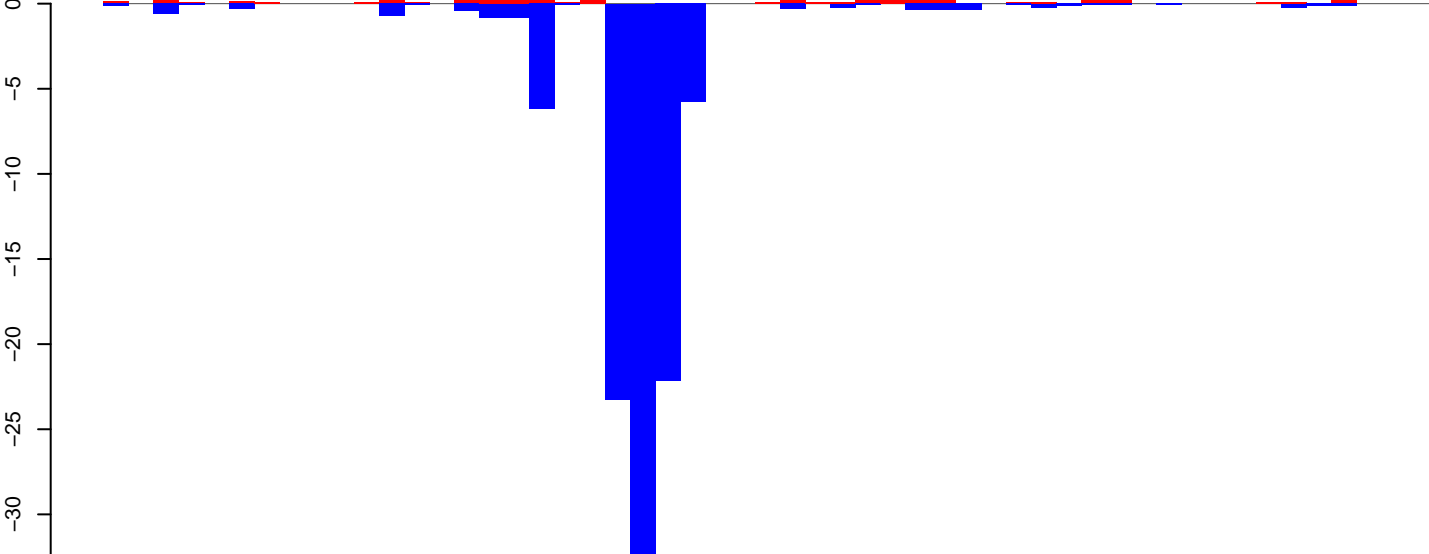
positive=4, negative=32, total=37

CuQuin_Testes_NL.24_35.rep



positive=3, negative=37, total=40

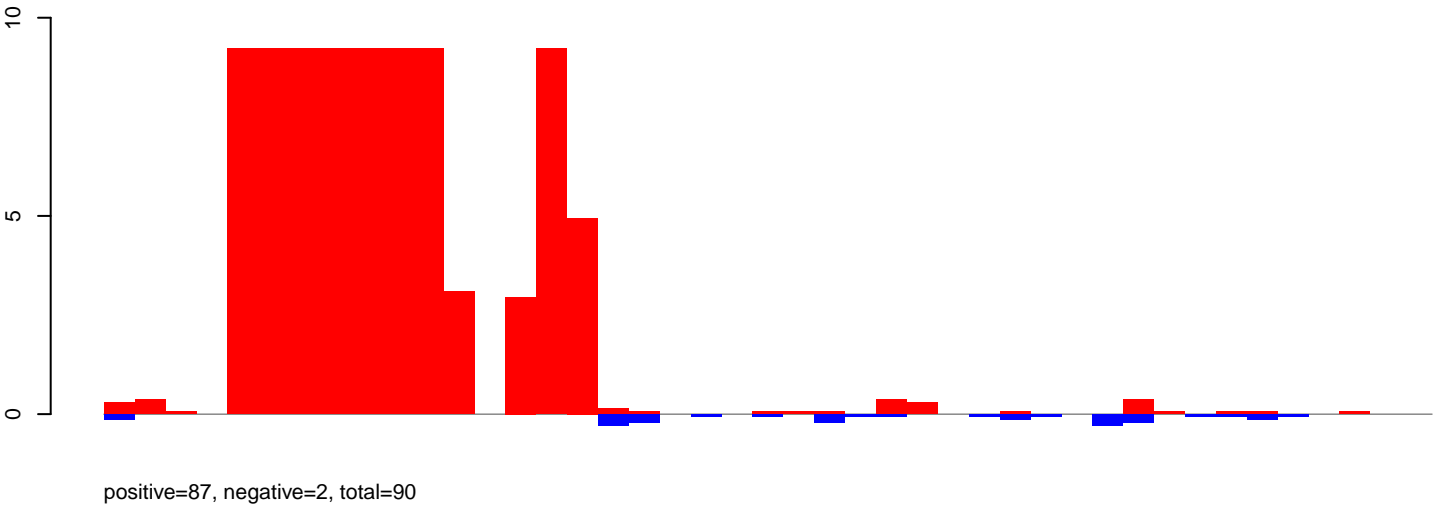
CuQuin_Testes_NL.rep



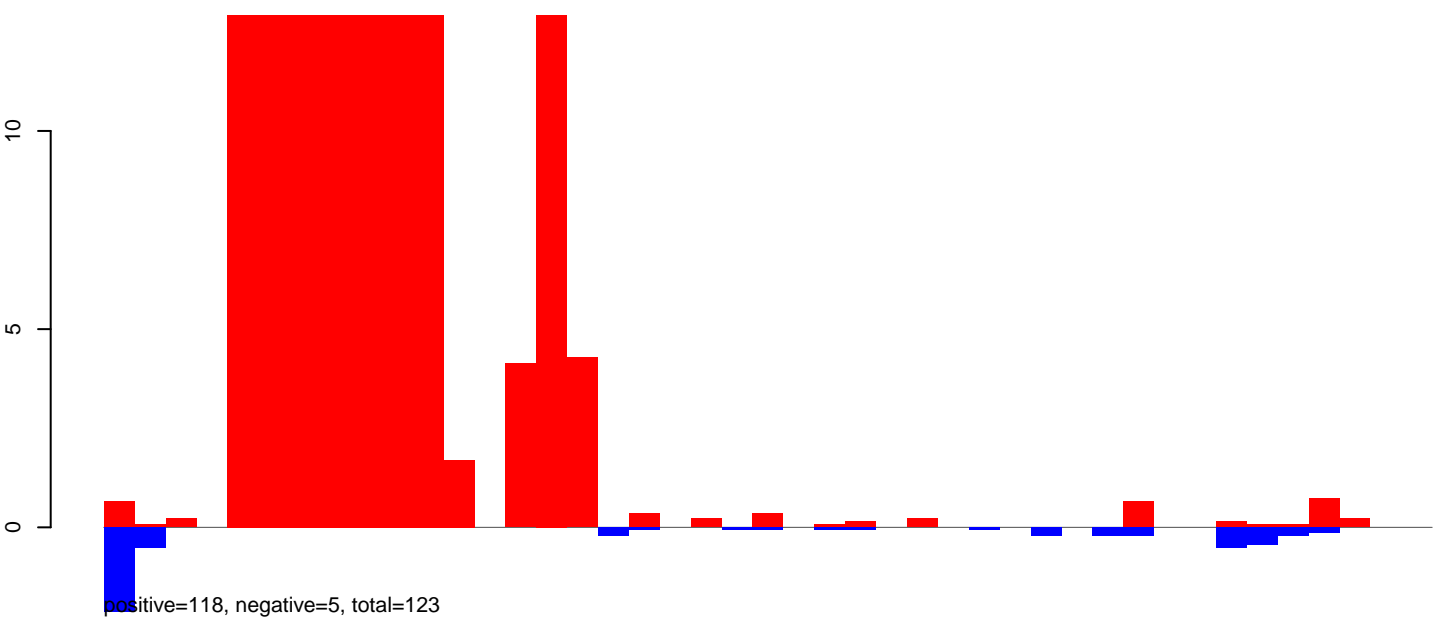
positive=8, negative=99, total=106

Window size=25, length=1303, TE@rnd-3_family-14@DNA_Zator@1370:1-1303

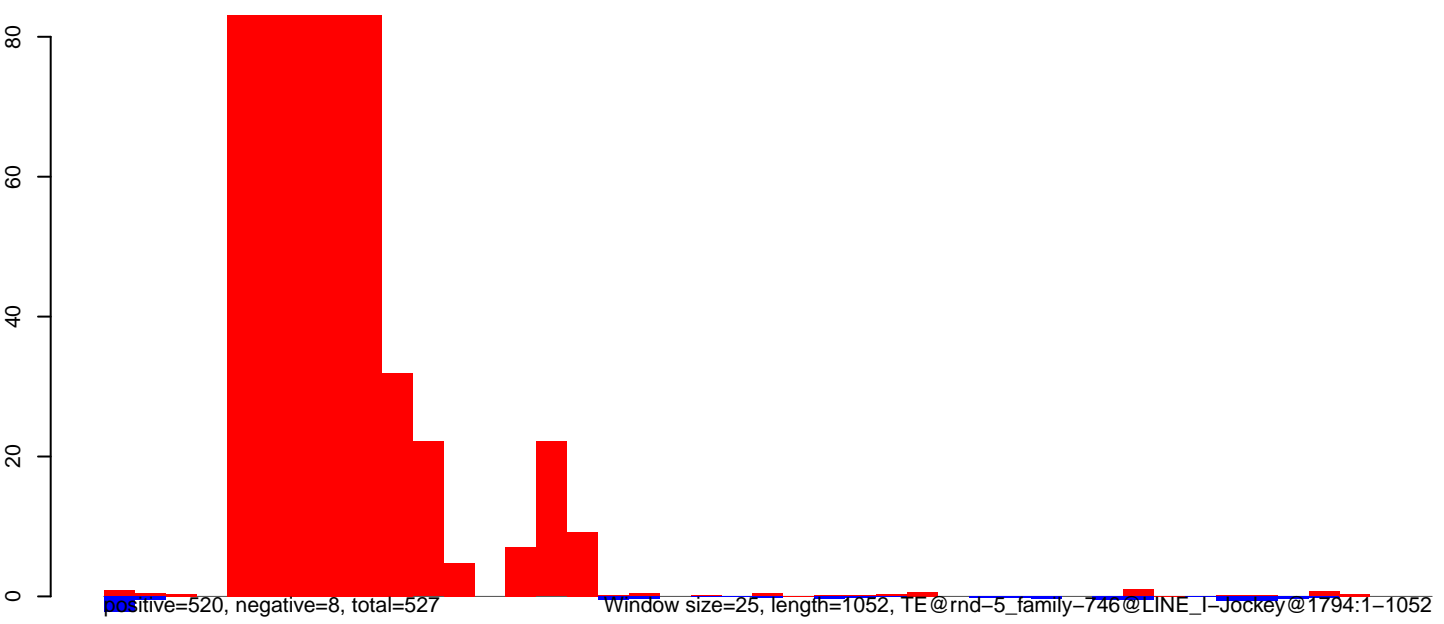
CuQuin_Testes_NL.18_23.rep



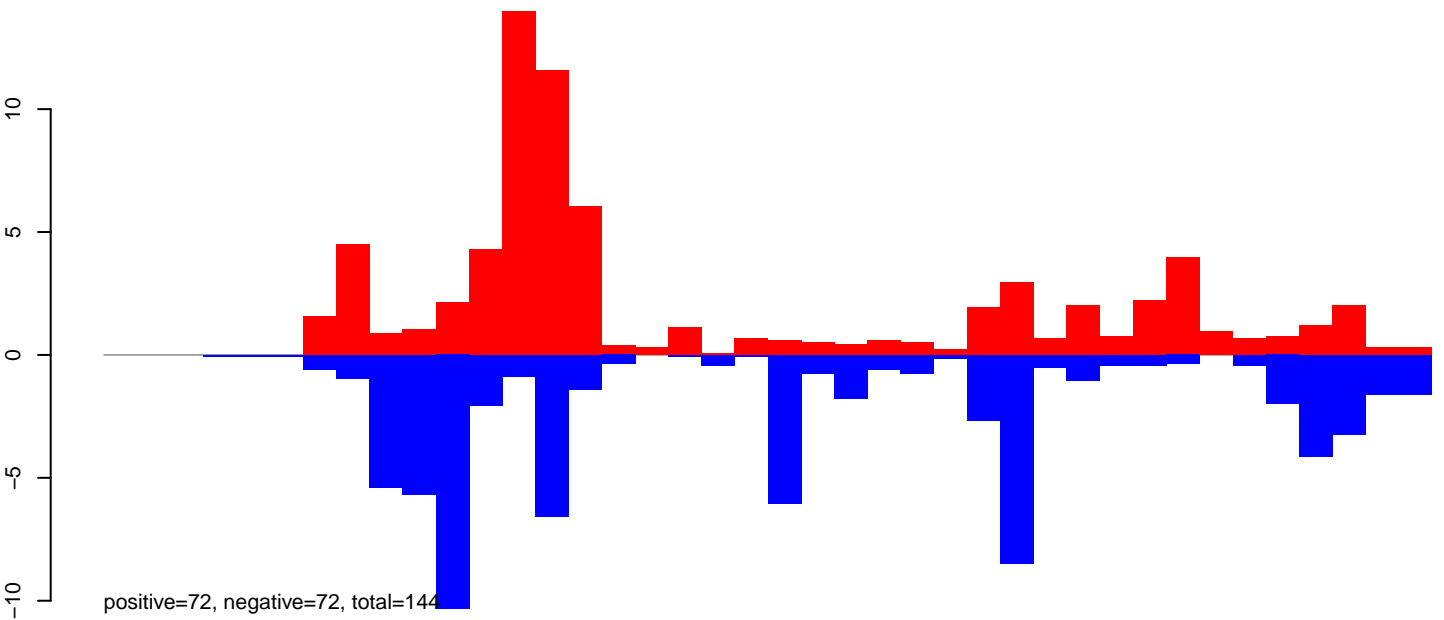
CuQuin_Testes_NL.24_35.rep



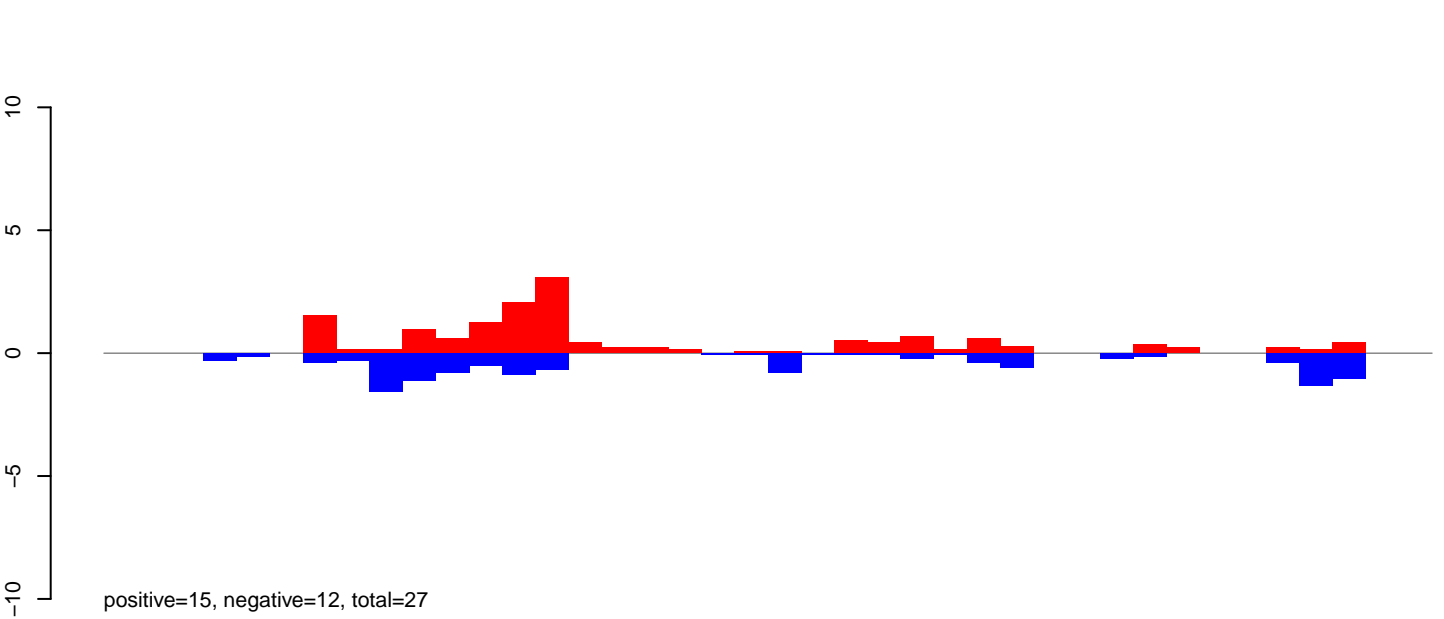
CuQuin_Testes_NL.rep



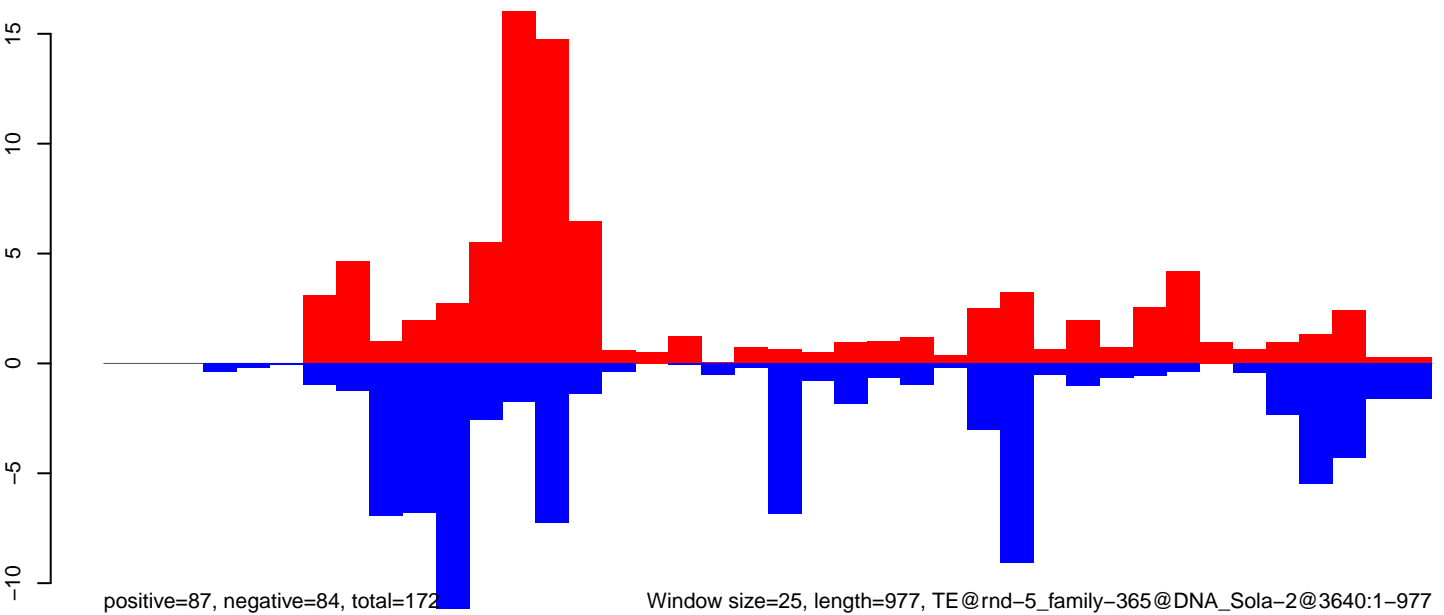
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



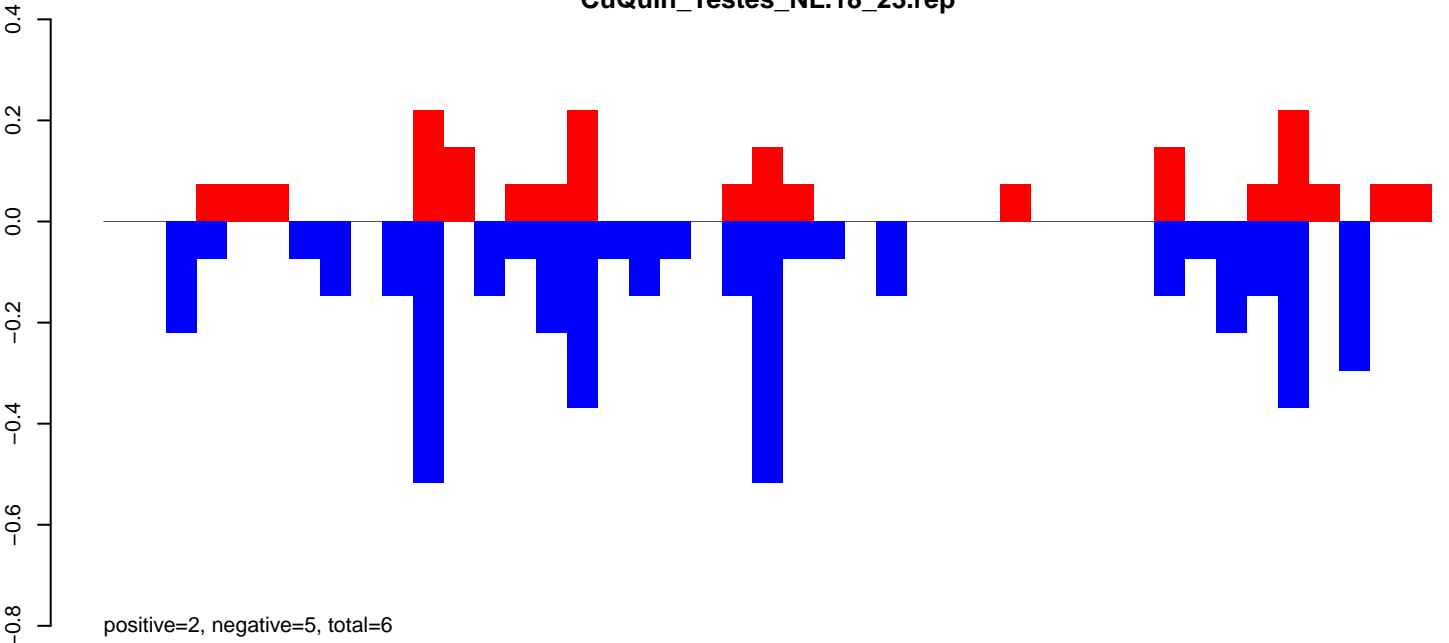
CuQuin_Testes_NL.rep



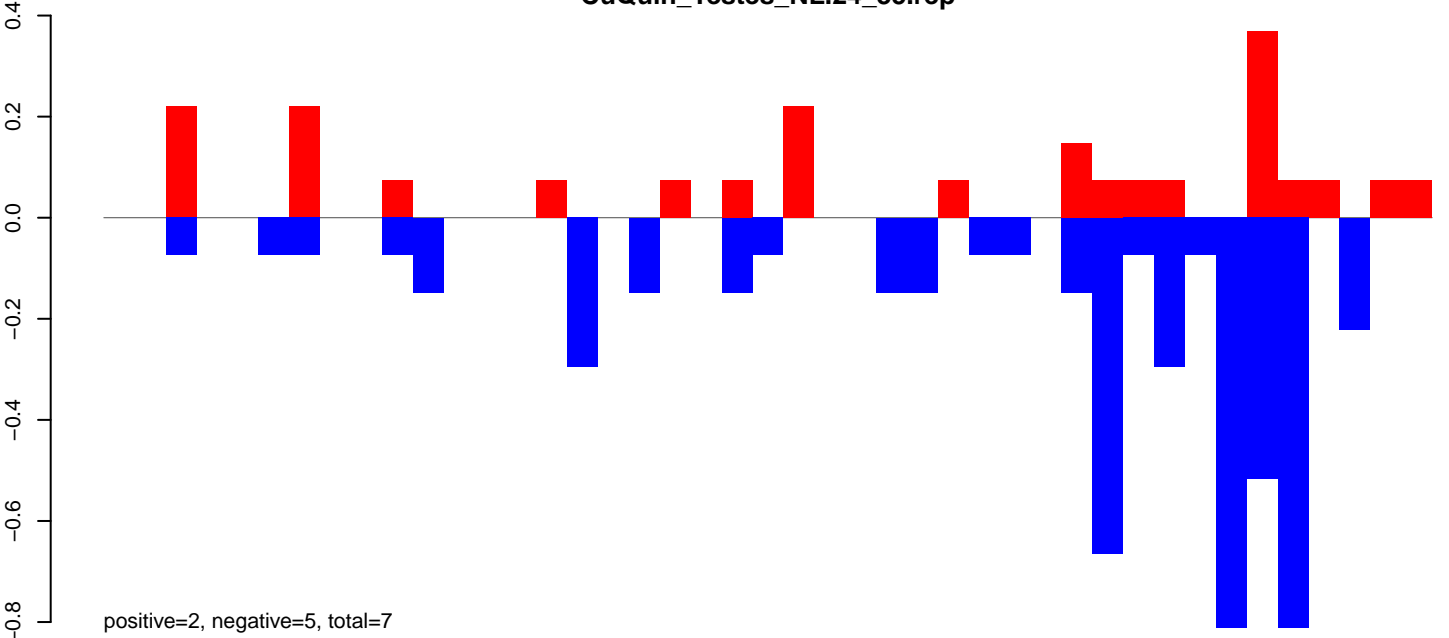
Window size=25, length=977, TE@rnd-5_family-365@DNA_Sola-2@3640:1-977

0 200 400 600 800 1000

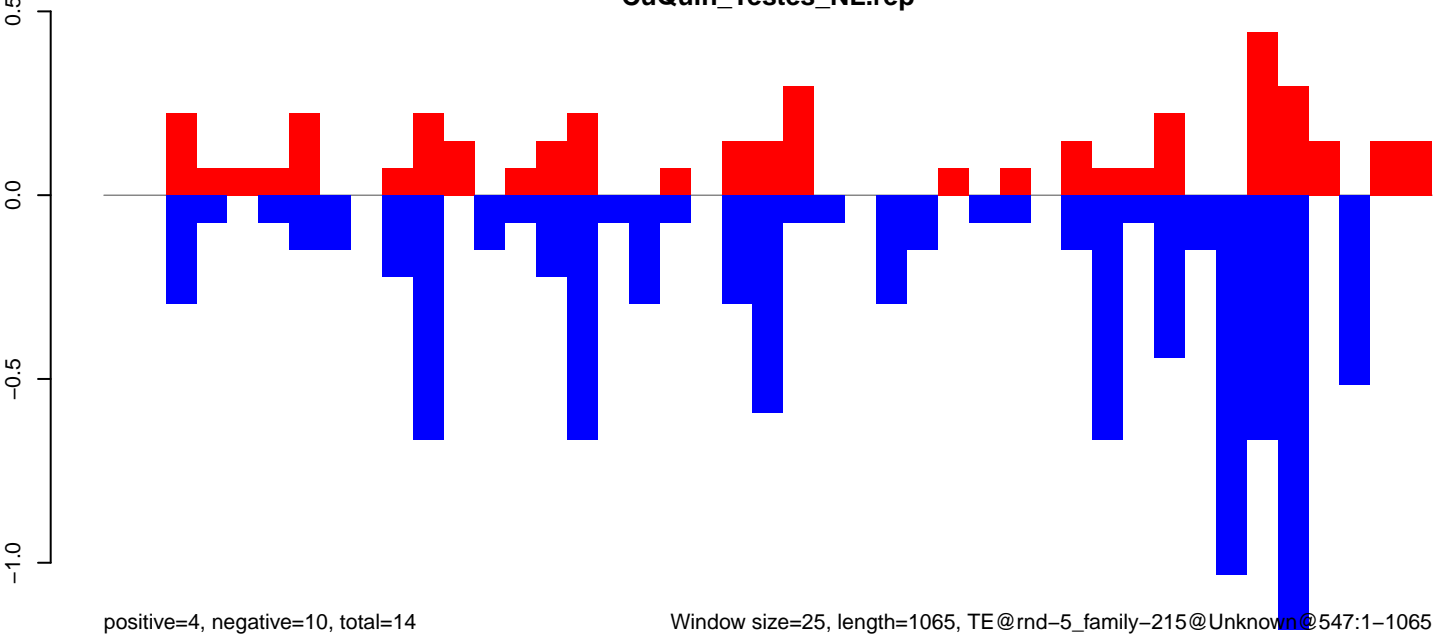
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

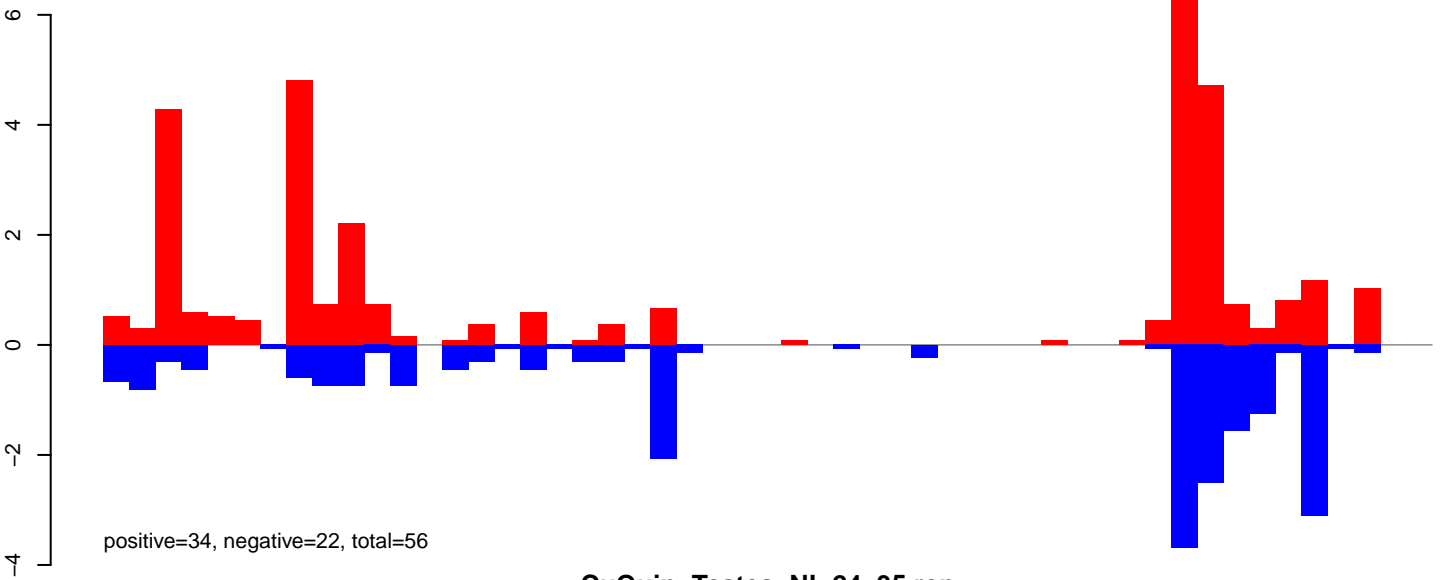


CuQuin_Testes_NL.rep

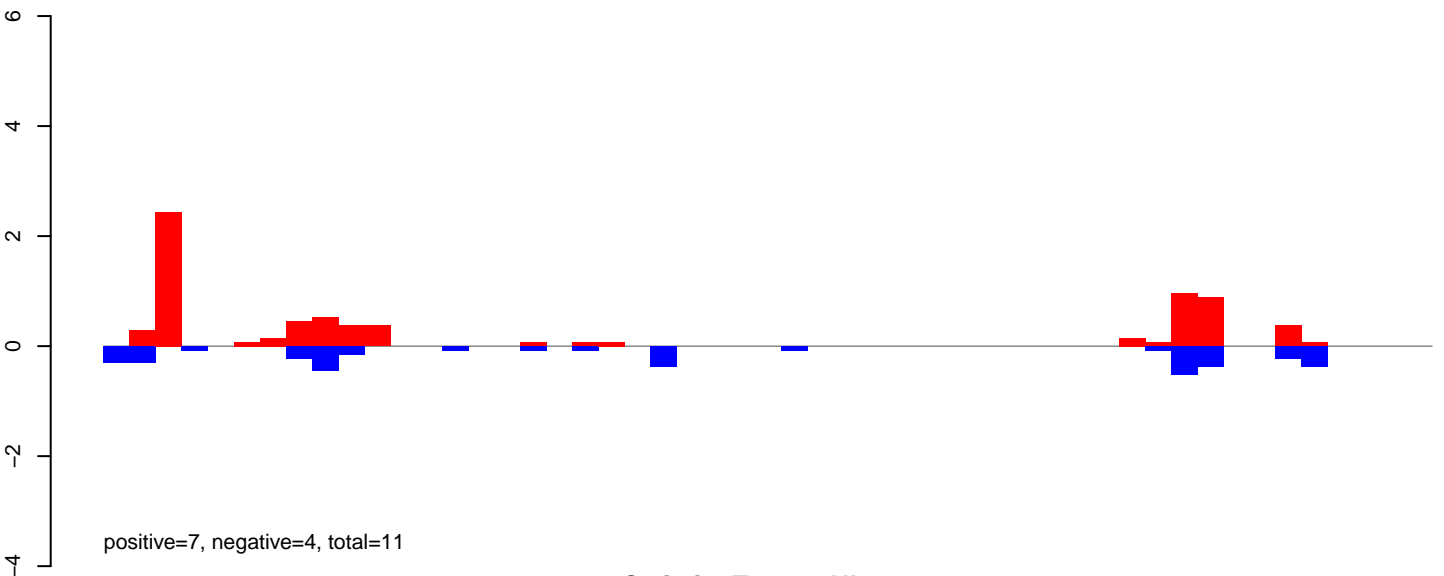


Window size=25, length=1065, TE@rnd-5_family-215@Unknown@547:1-1065

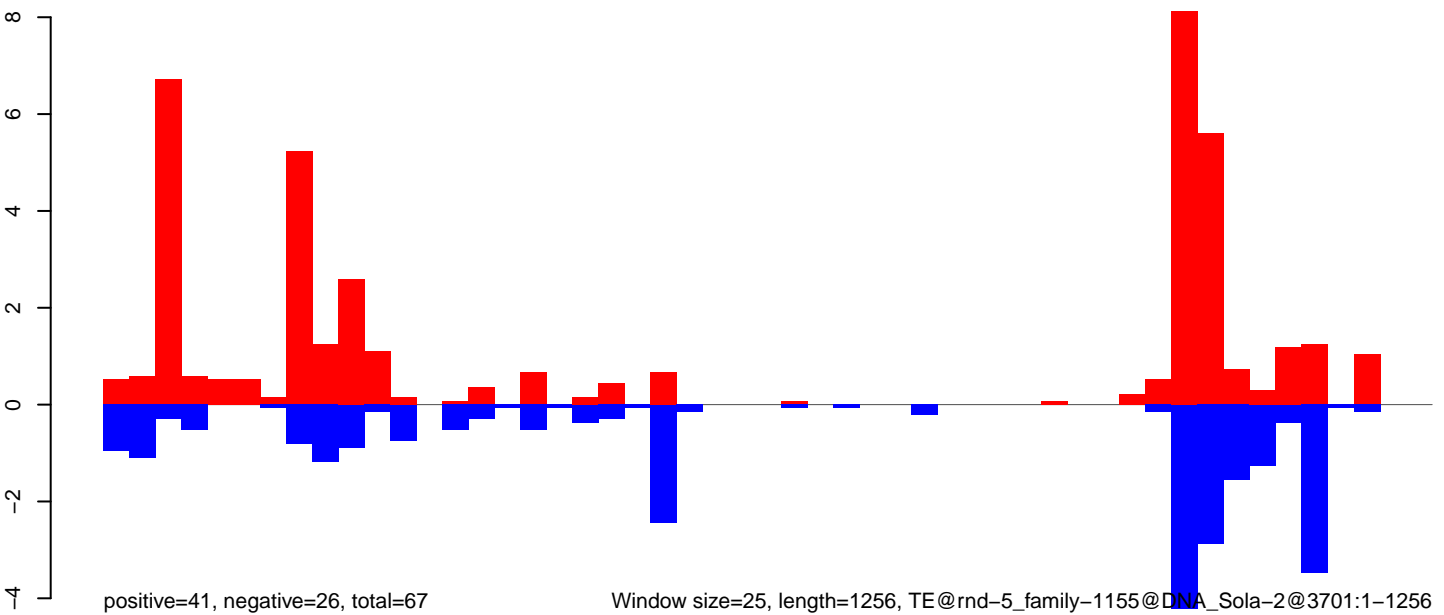
CuQuin_Testes_NL.18_23.rep



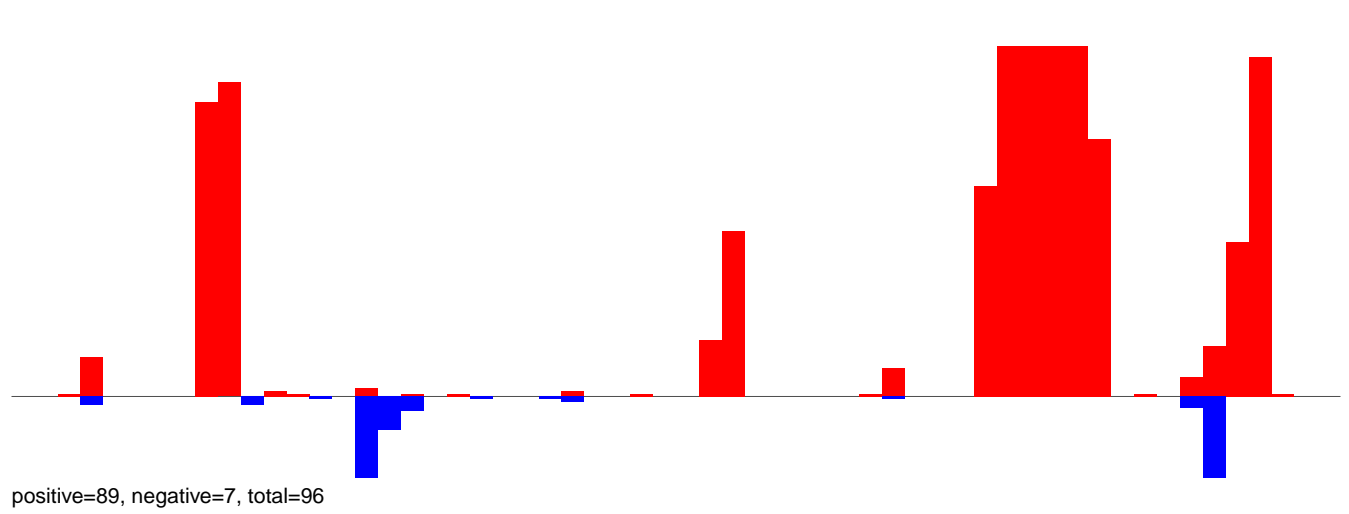
CuQuin_Testes_NL.24_35.rep



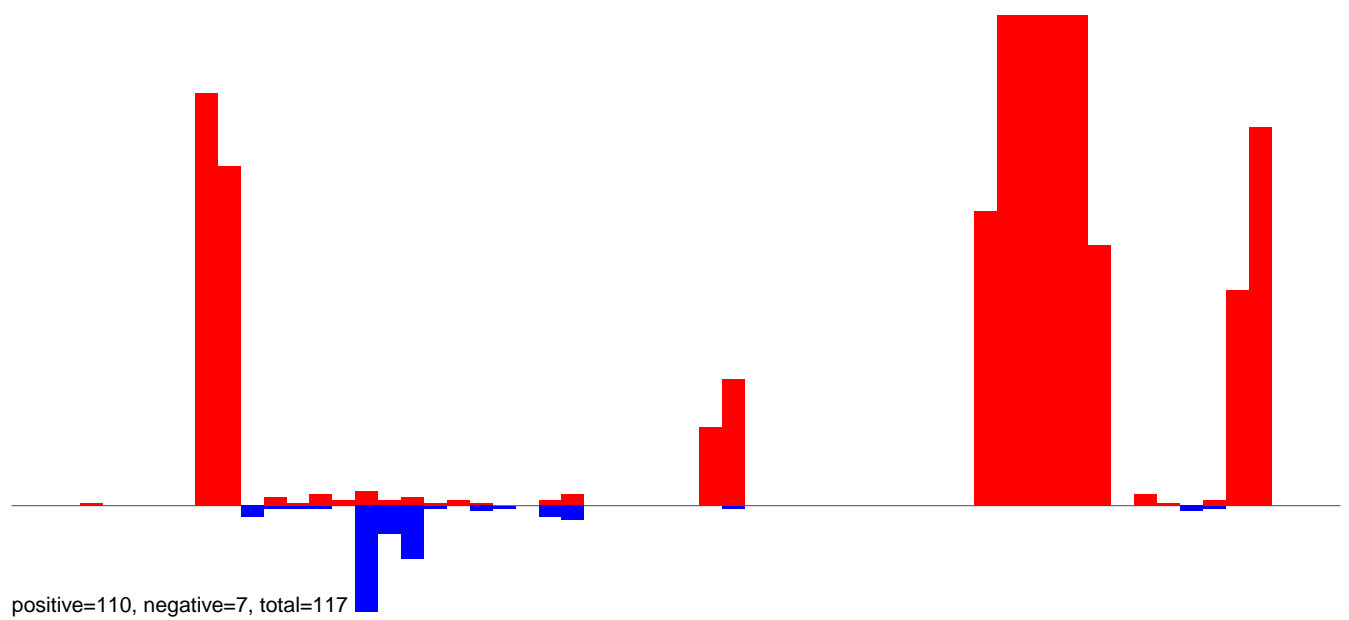
CuQuin_Testes_NL.rep



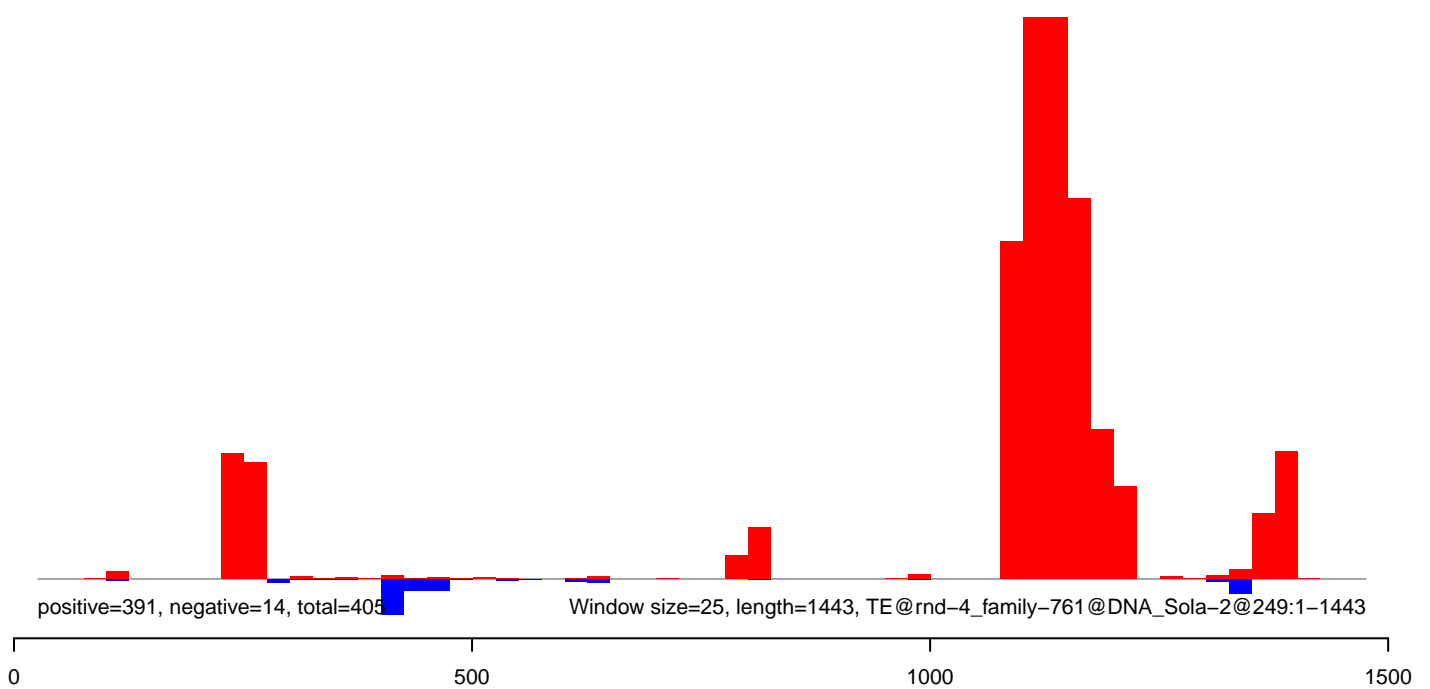
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



CuQuin_Testes_NL.18_23.rep

positive=1, negative=1, total=2

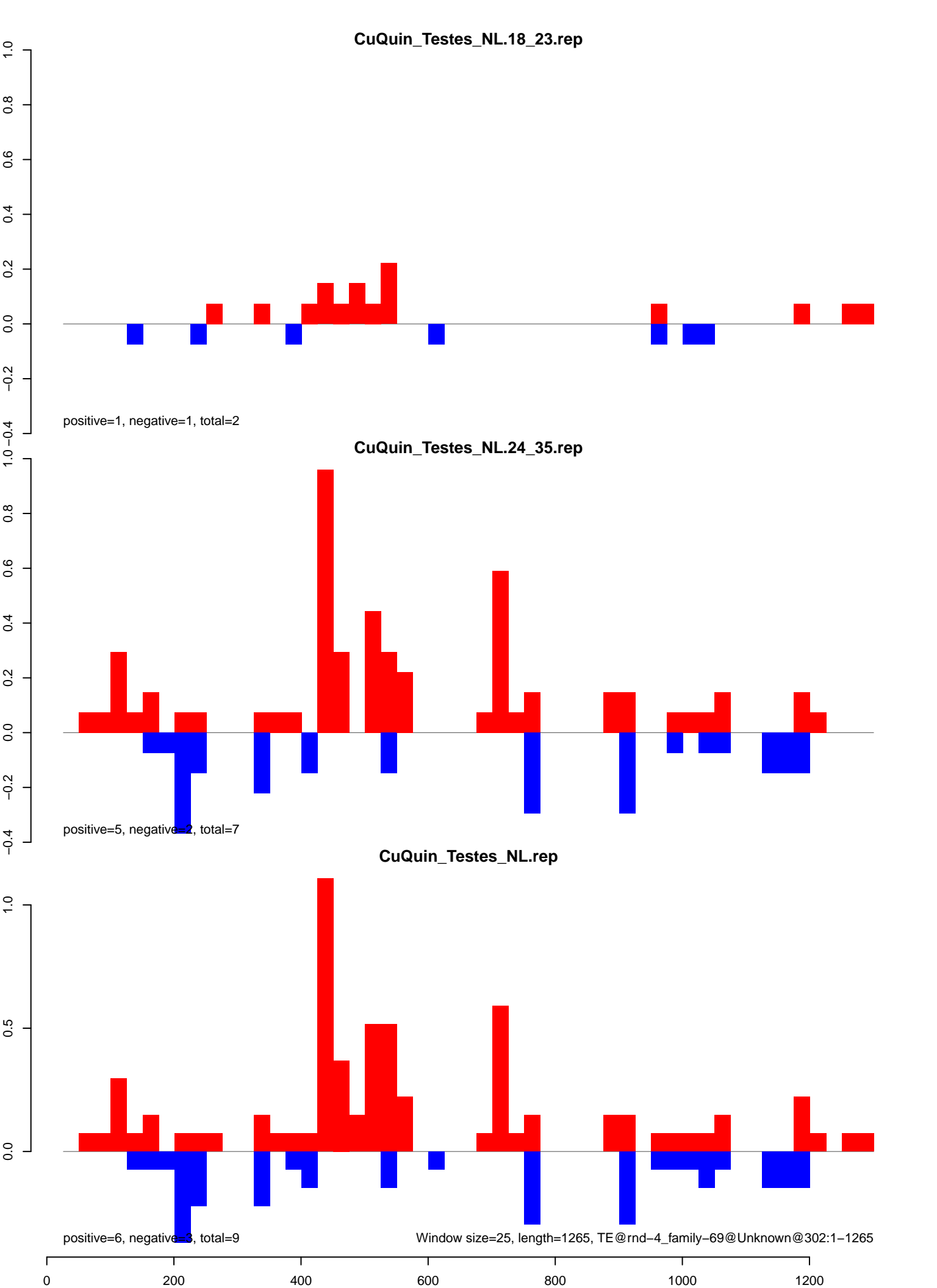
CuQuin_Testes_NL.24_35.rep

positive=5, negative=2, total=7

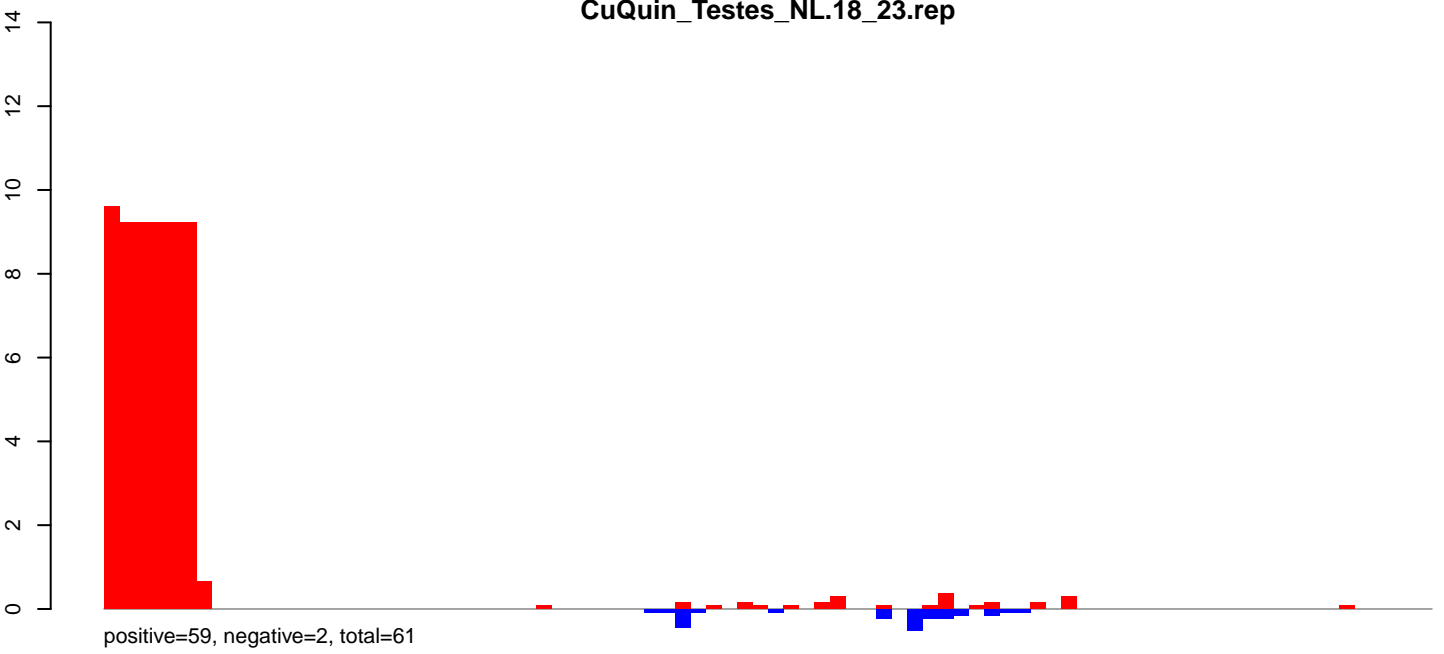
CuQuin_Testes_NL.rep

positive=6, negative=3, total=9

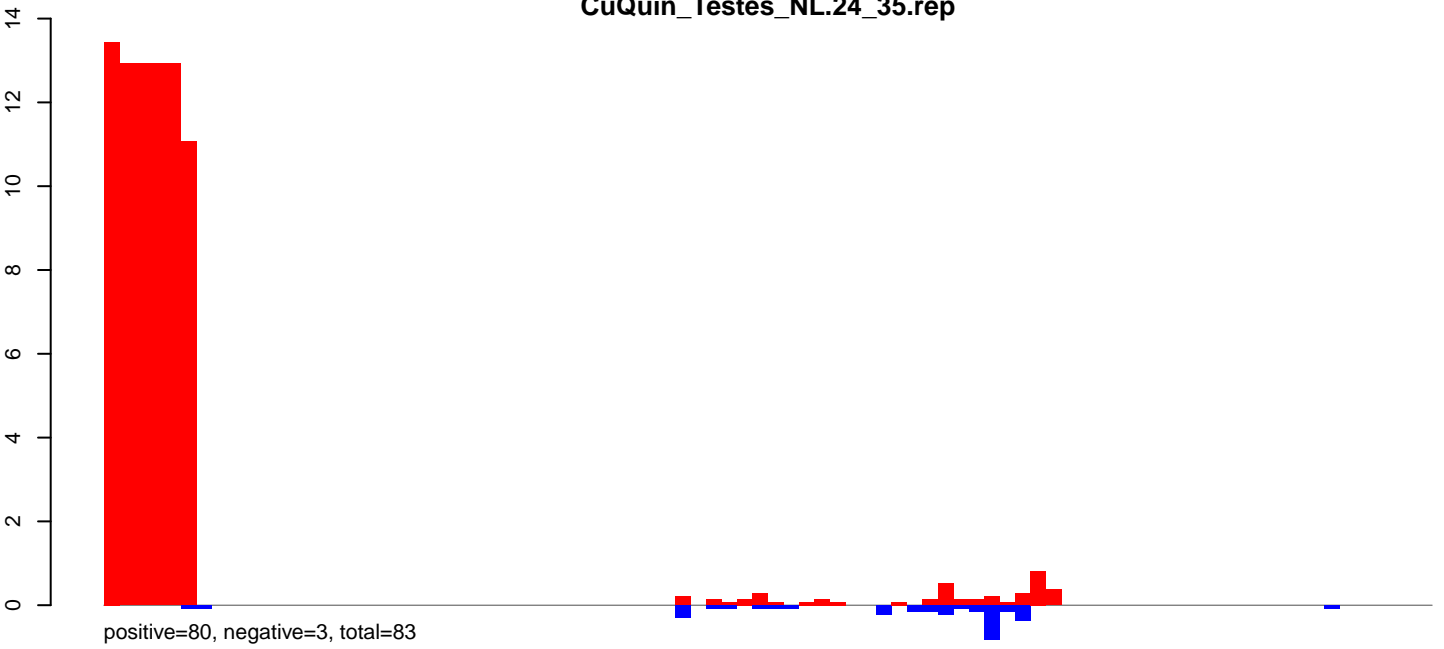
Window size=25, length=1265, TE@rnd-4_family-69@Unknown@302:1-1265



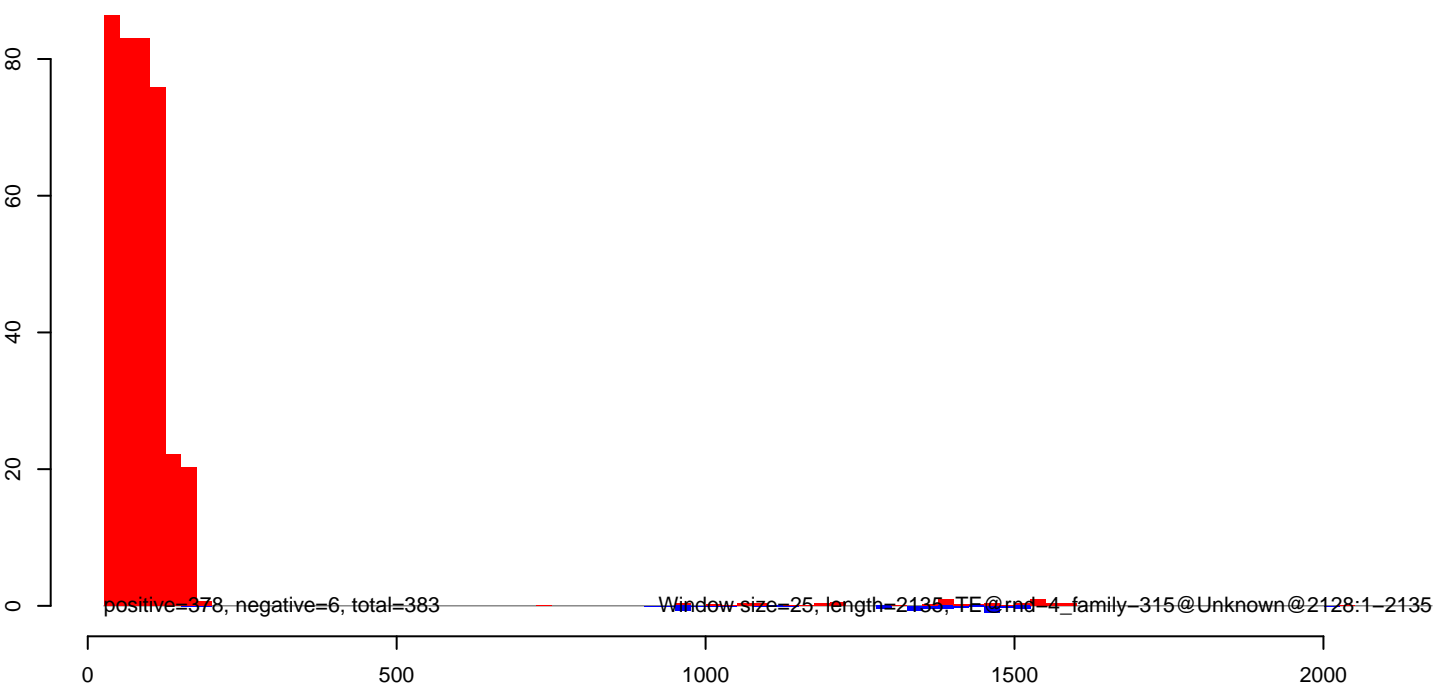
CuQuin_Testes_NL.18_23.rep



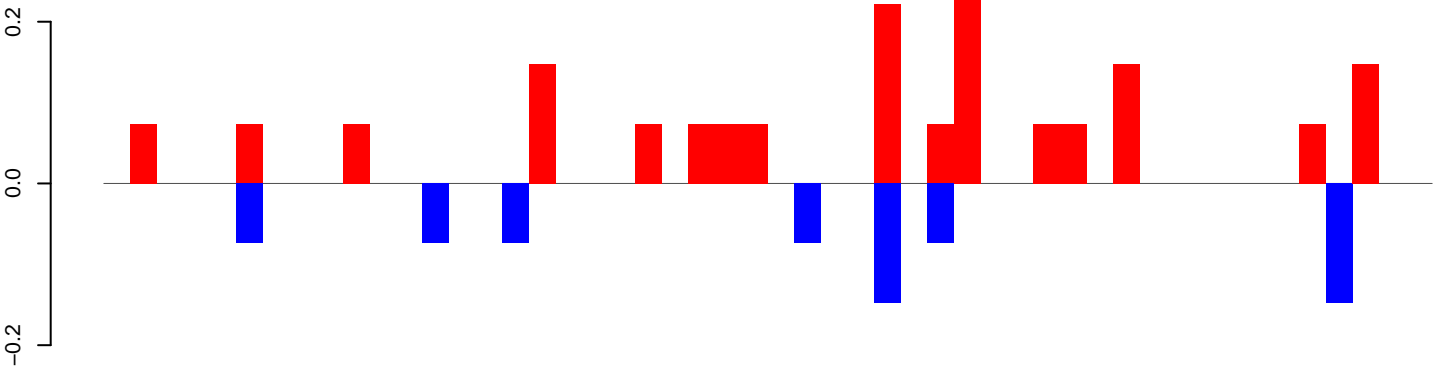
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

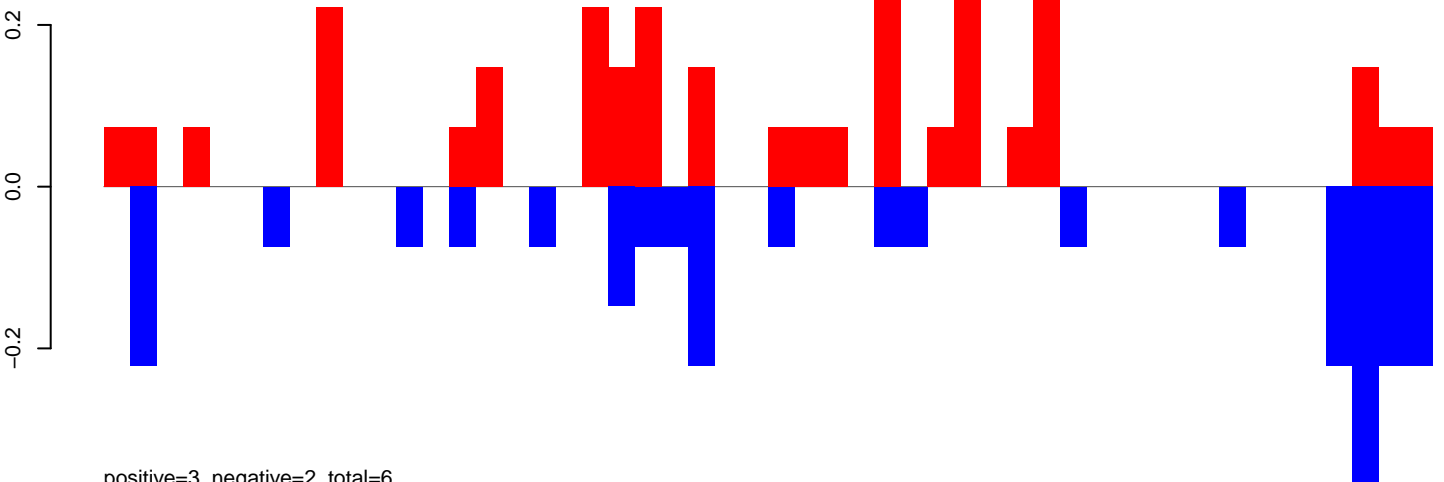


CuQuin_Testes_NL.18_23.rep



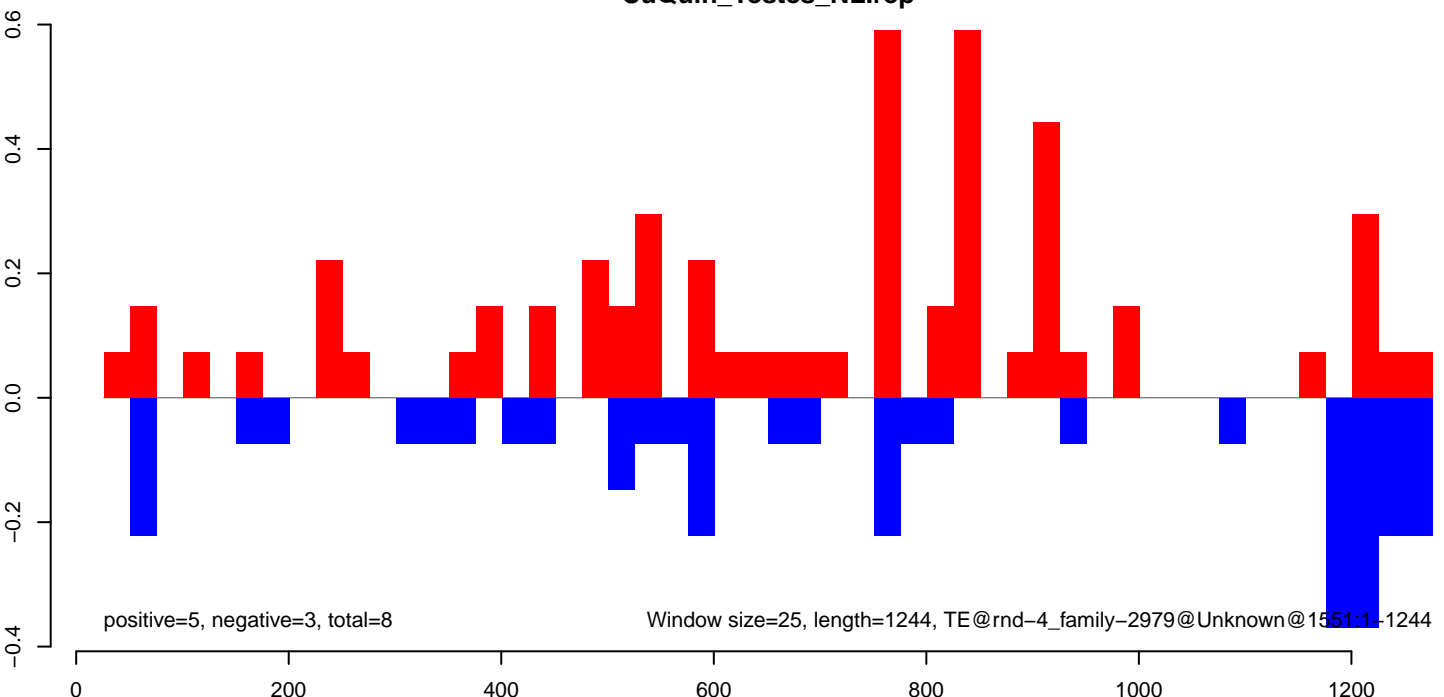
positive=2, negative=1, total=2

CuQuin_Testes_NL.24_35.rep



positive=3, negative=2, total=6

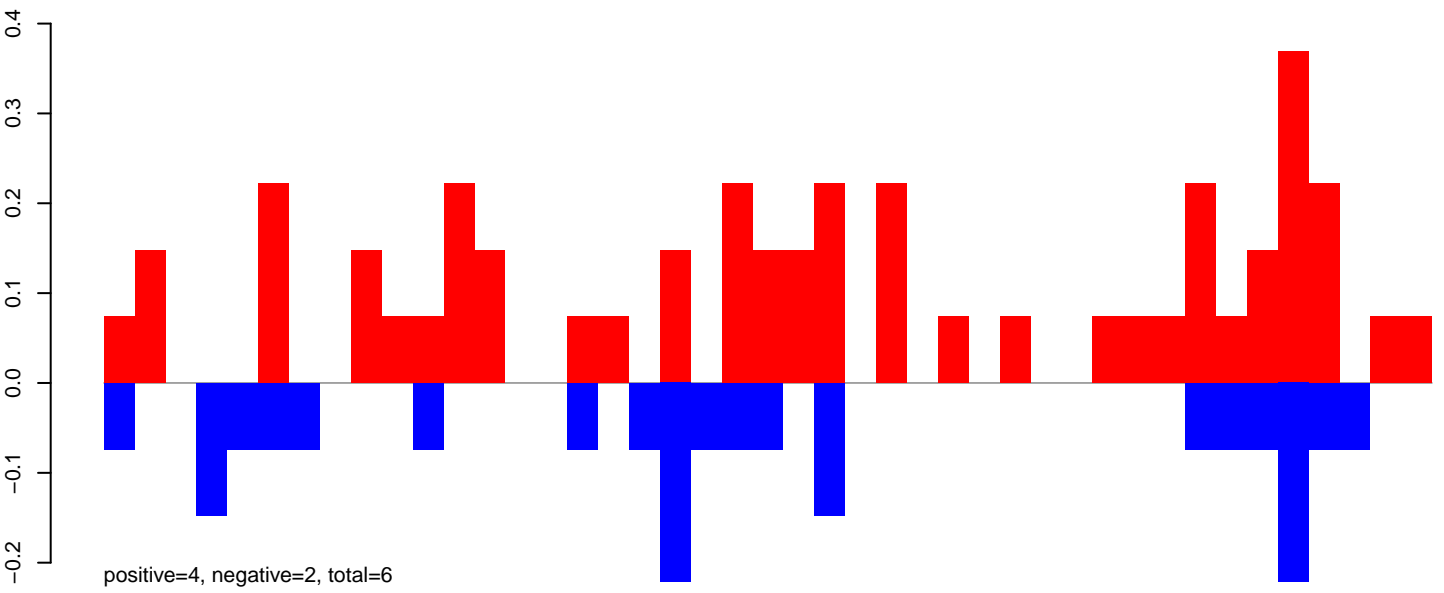
CuQuin_Testes_NL.rep



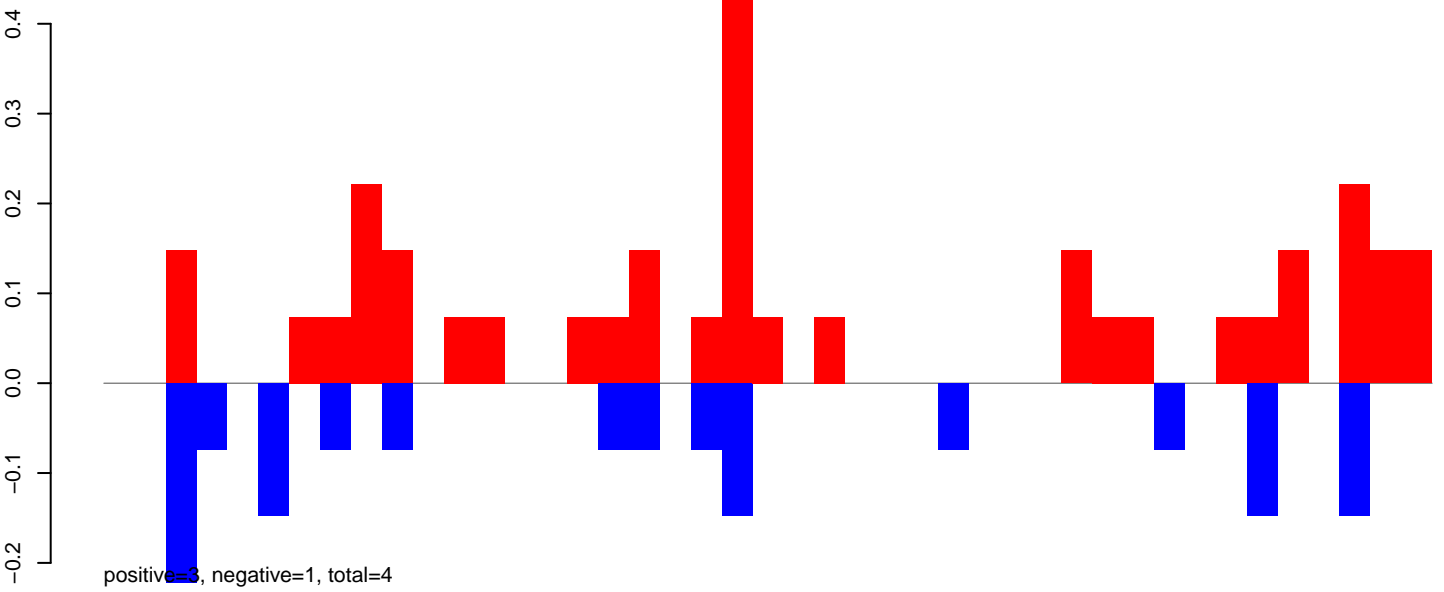
positive=5, negative=3, total=8

Window size=25, length=1244, TE@rnd-4_family-2979@Unknown@1551:1-1244

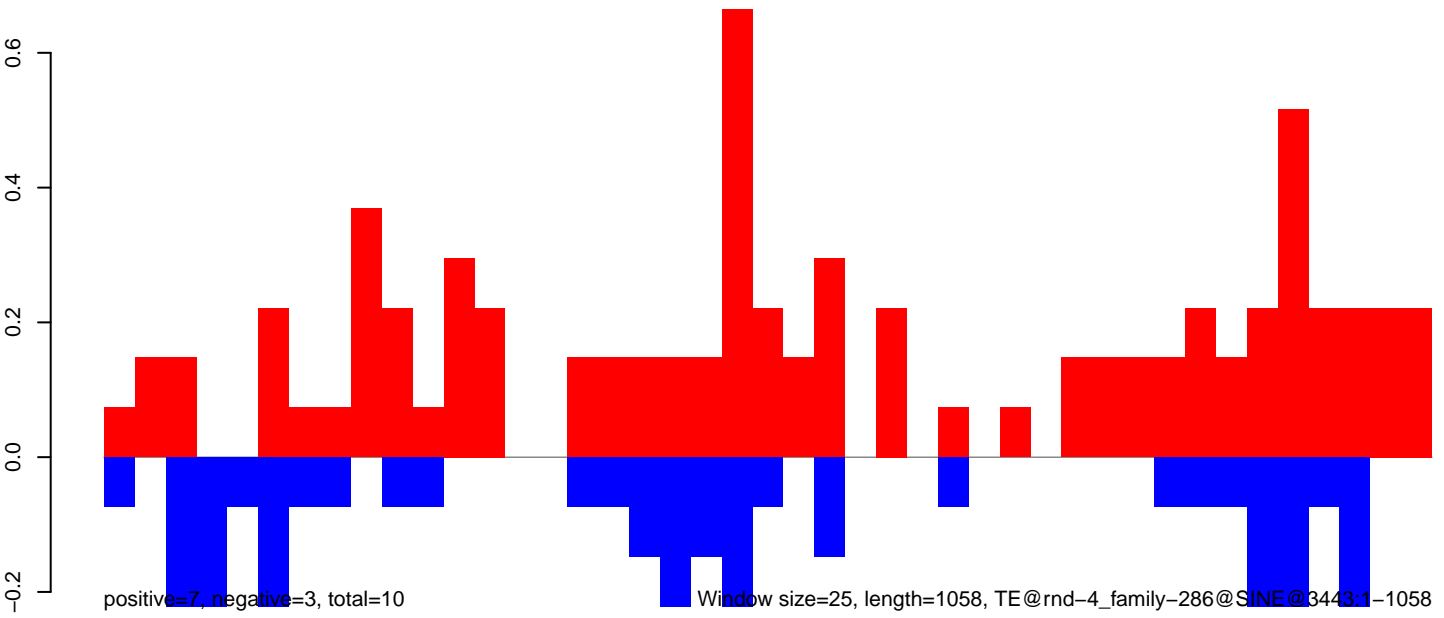
CuQuin_Testes_NL.18_23.rep



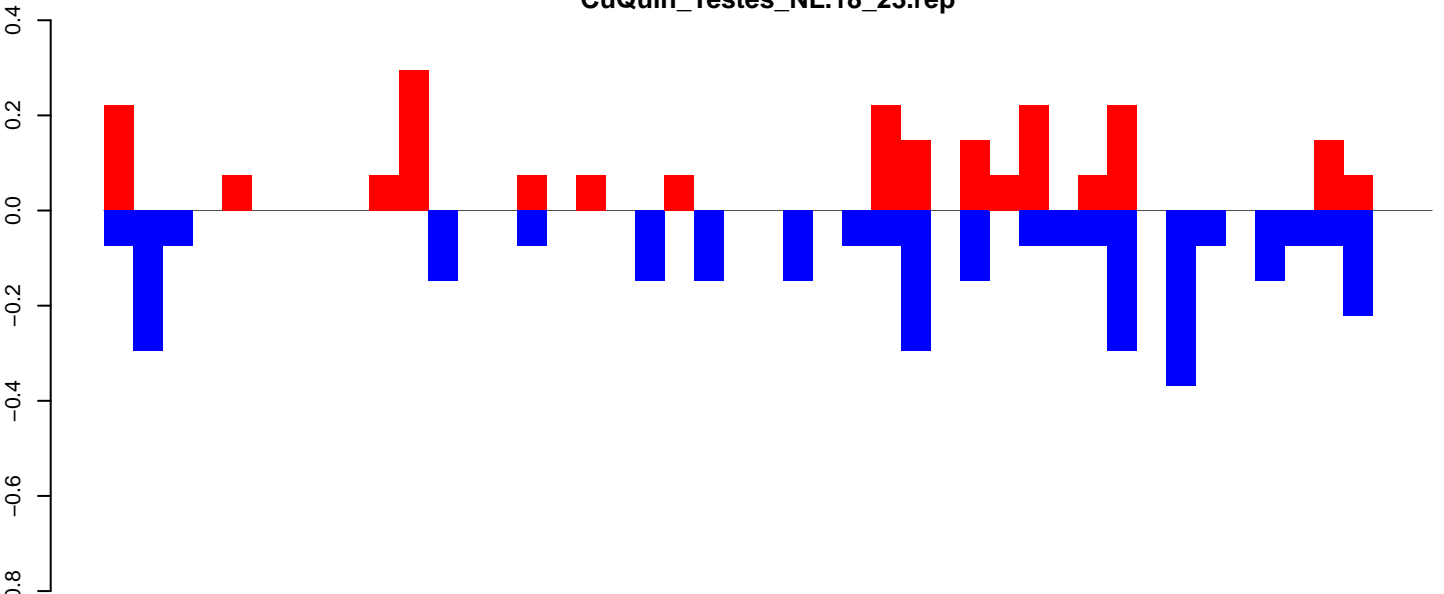
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

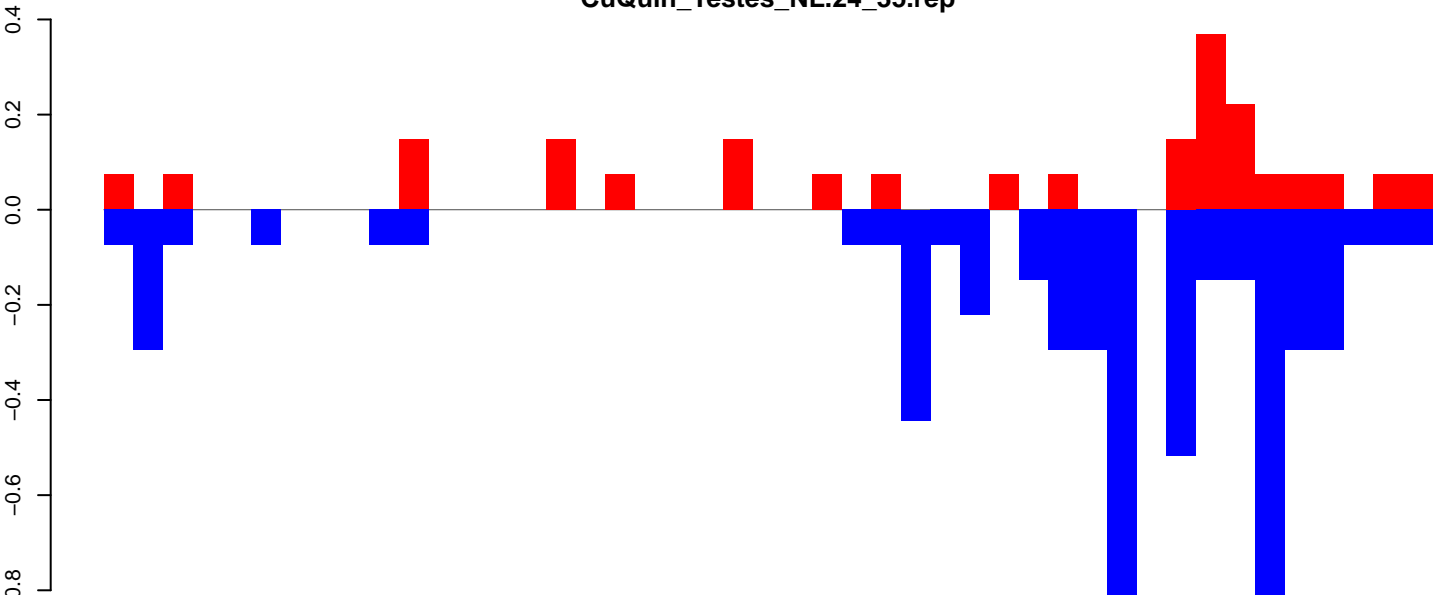


CuQuin_Testes_NL.18_23.rep



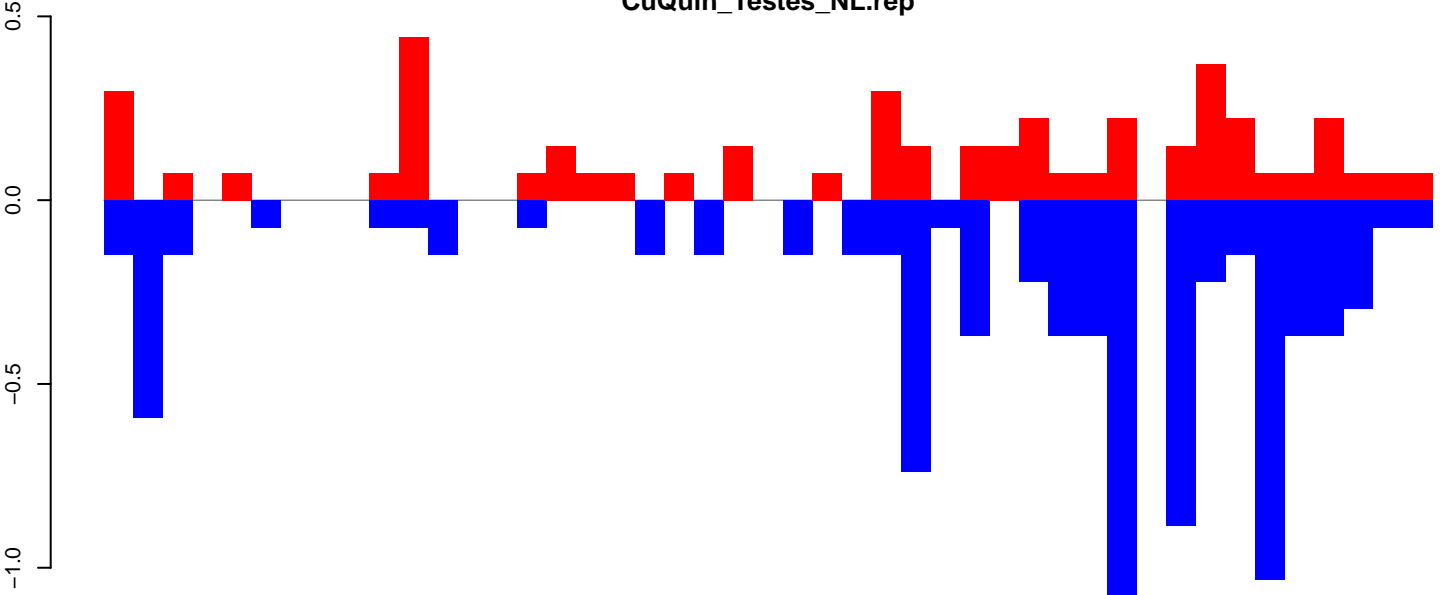
positive=2, negative=3, total=5

CuQuin_Testes_NL.24_35.rep



positive=2, negative=6, total=8

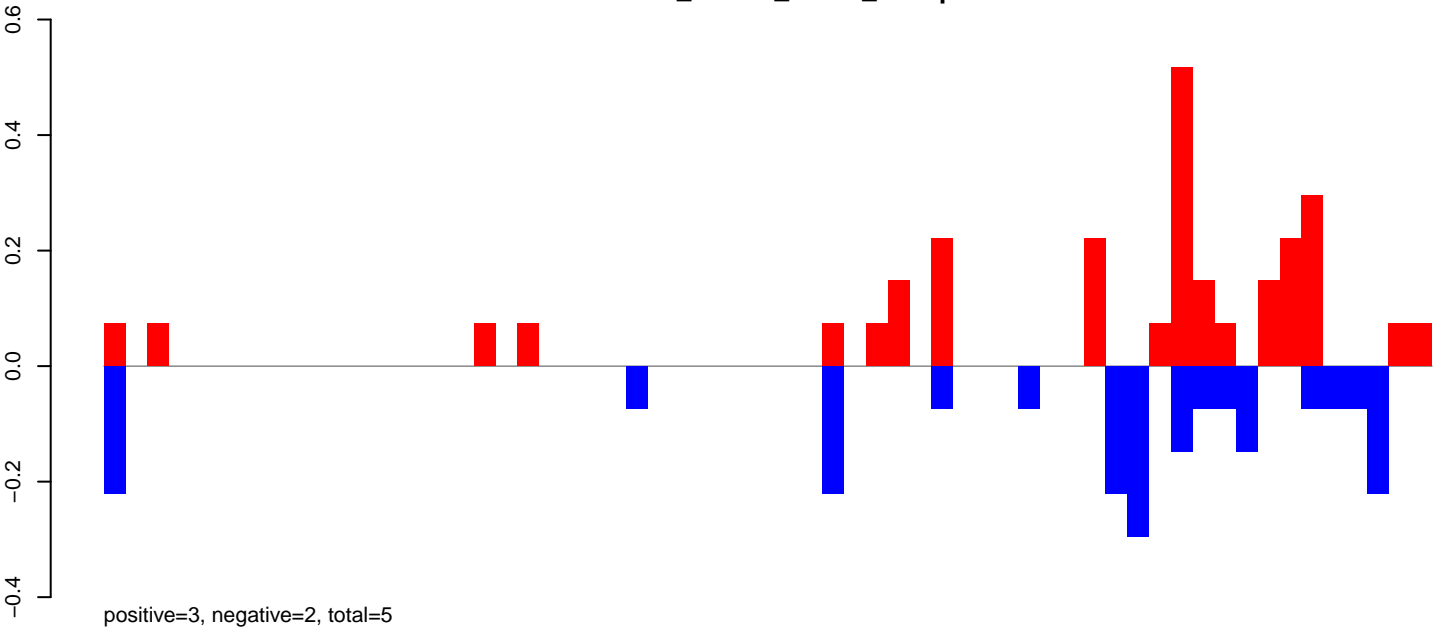
CuQuin_Testes_NL.rep



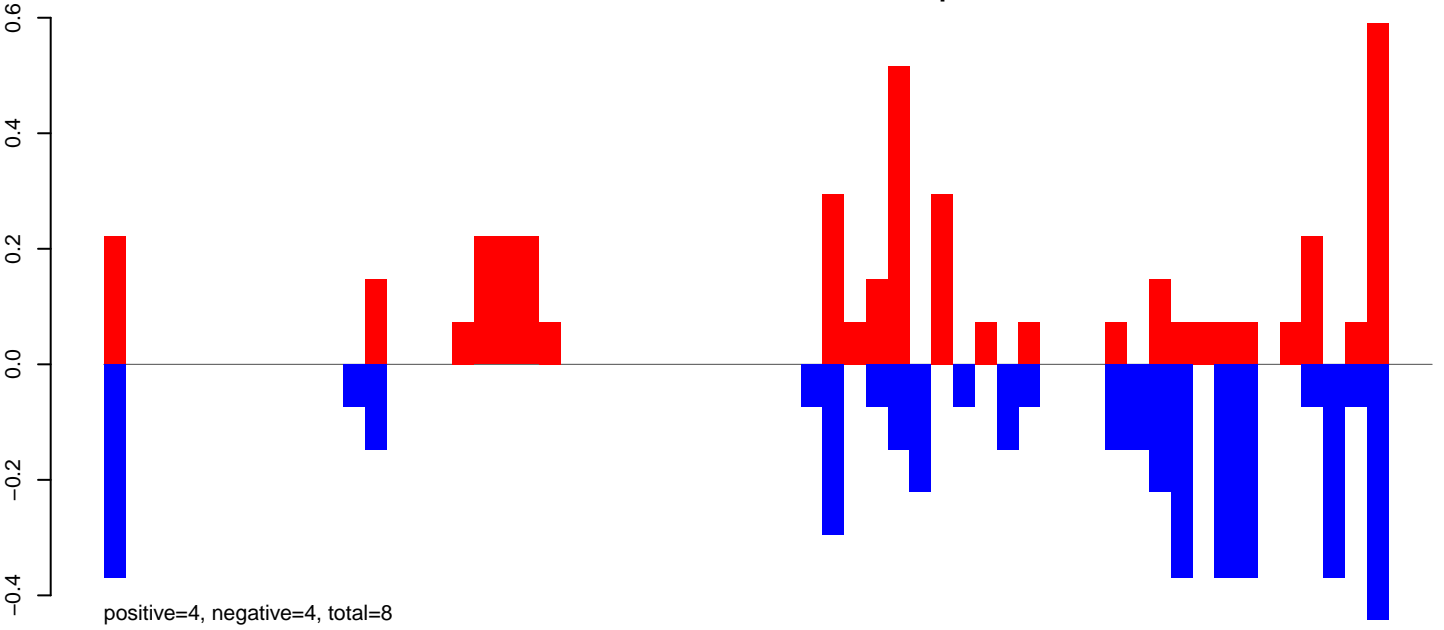
positive=4, negative=9, total=13

Window size=25, length=1109, TE@rnd-4_family-2243@DNA@2839:1-1109

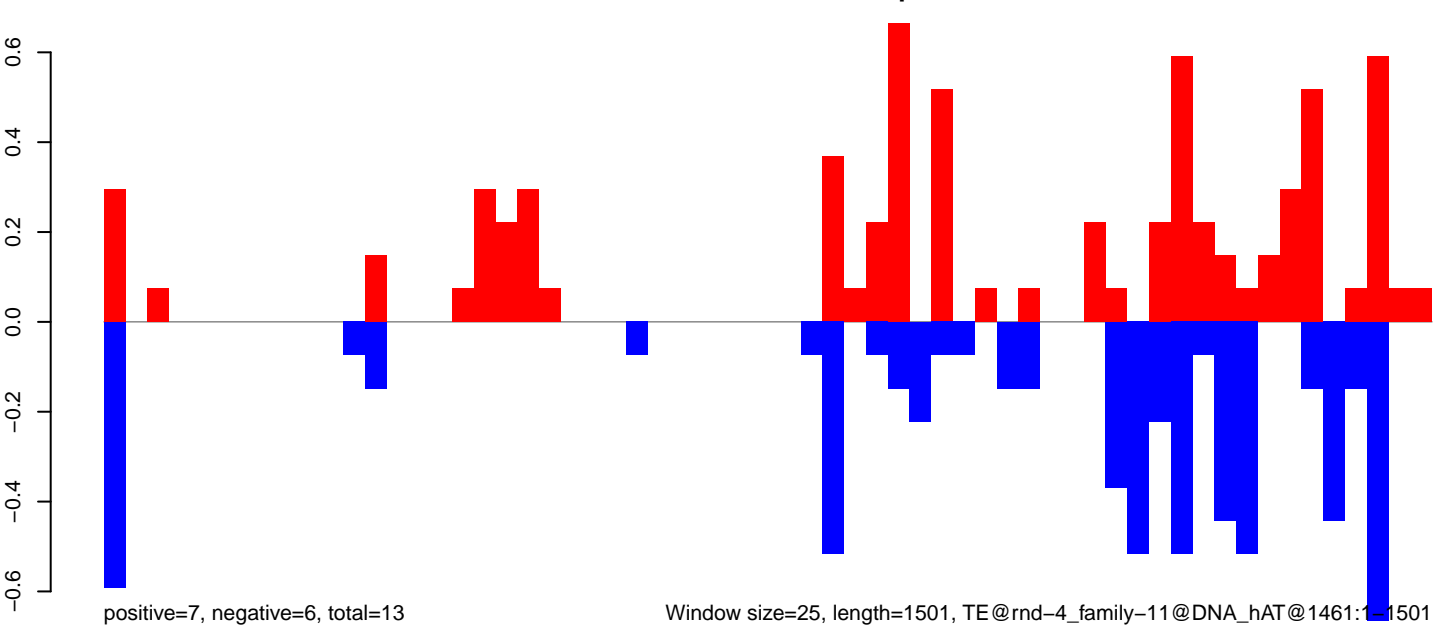
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



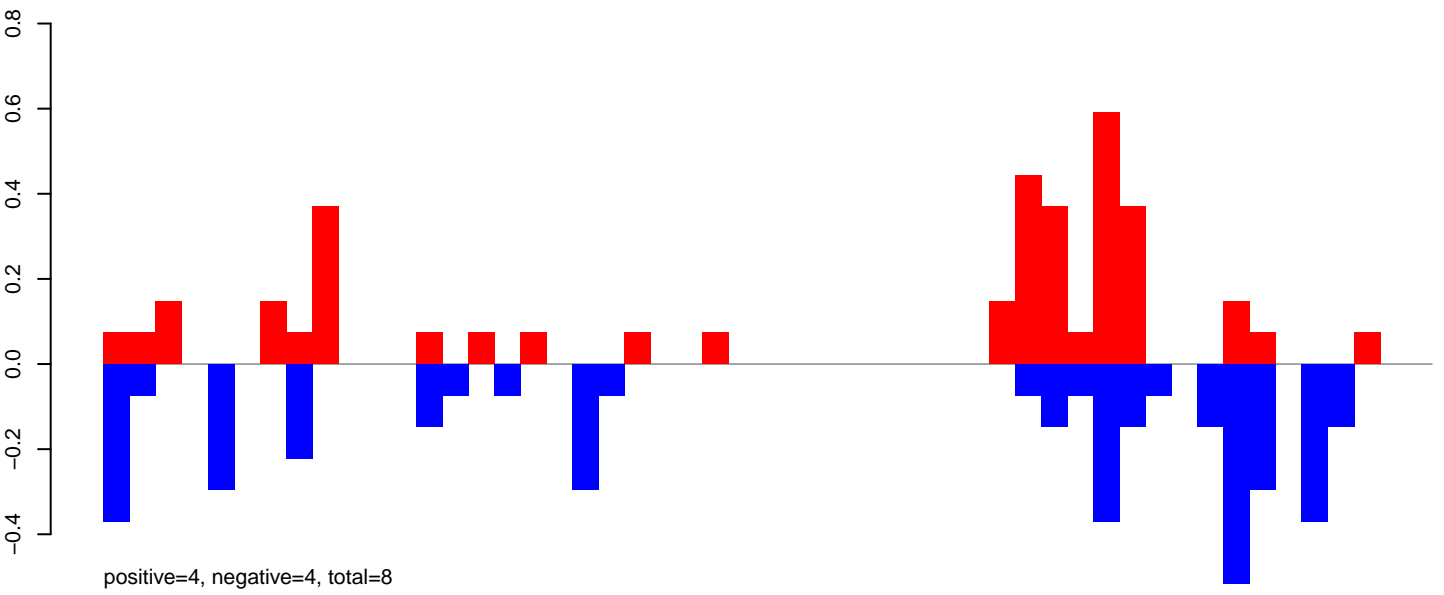
CuQuin_Testes_NL.rep



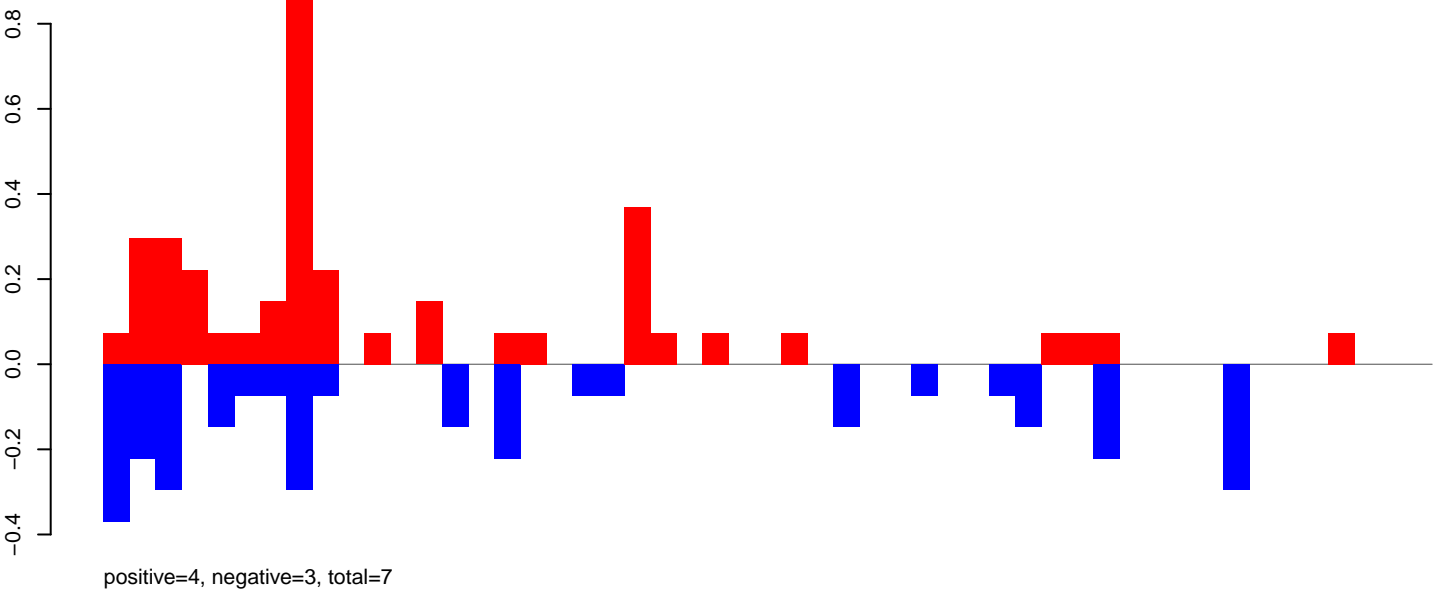
Window size=25, length=1501, TE@rnd-4_family-11@DNA_hAT@1461:1-1501

0 500 1000 1500

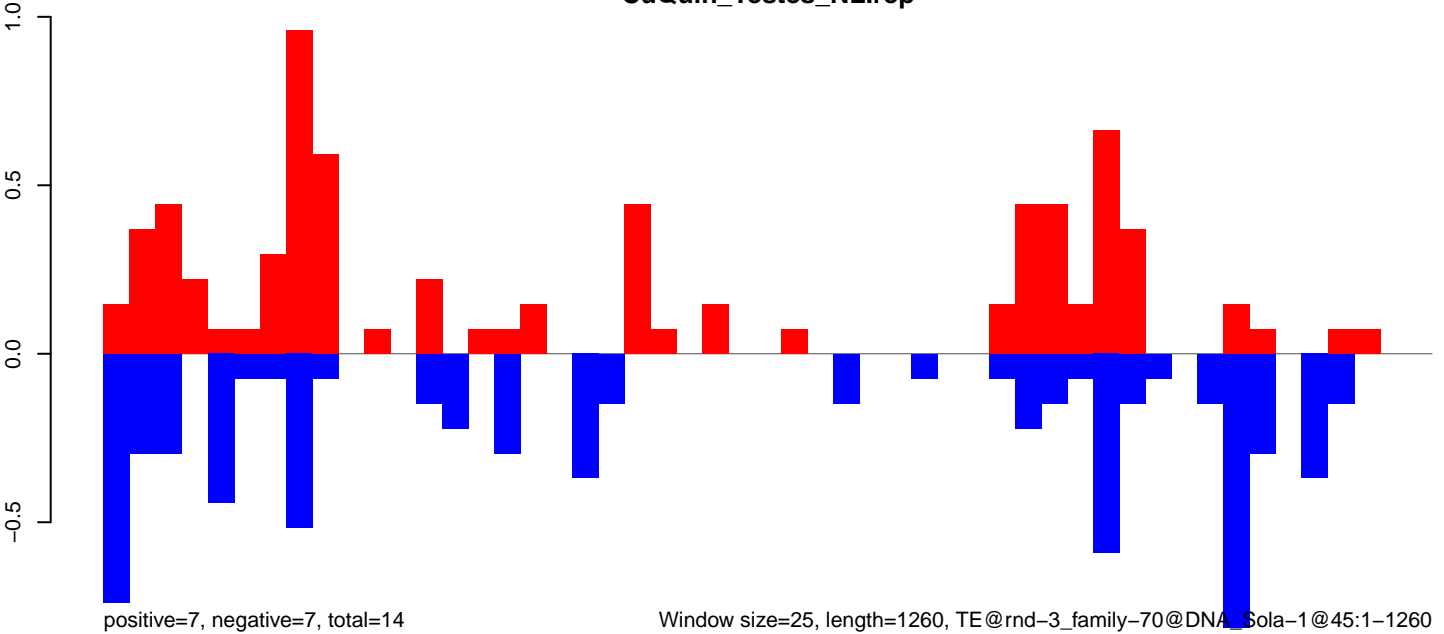
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



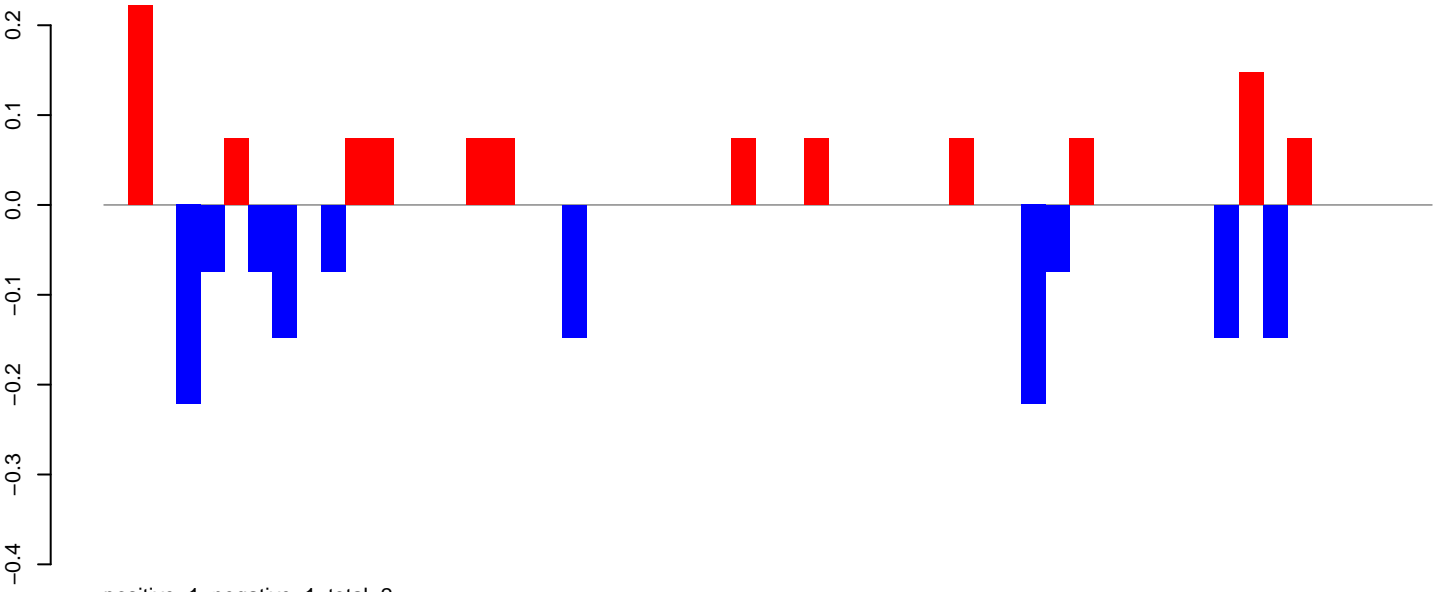
CuQuin_Testes_NL.rep



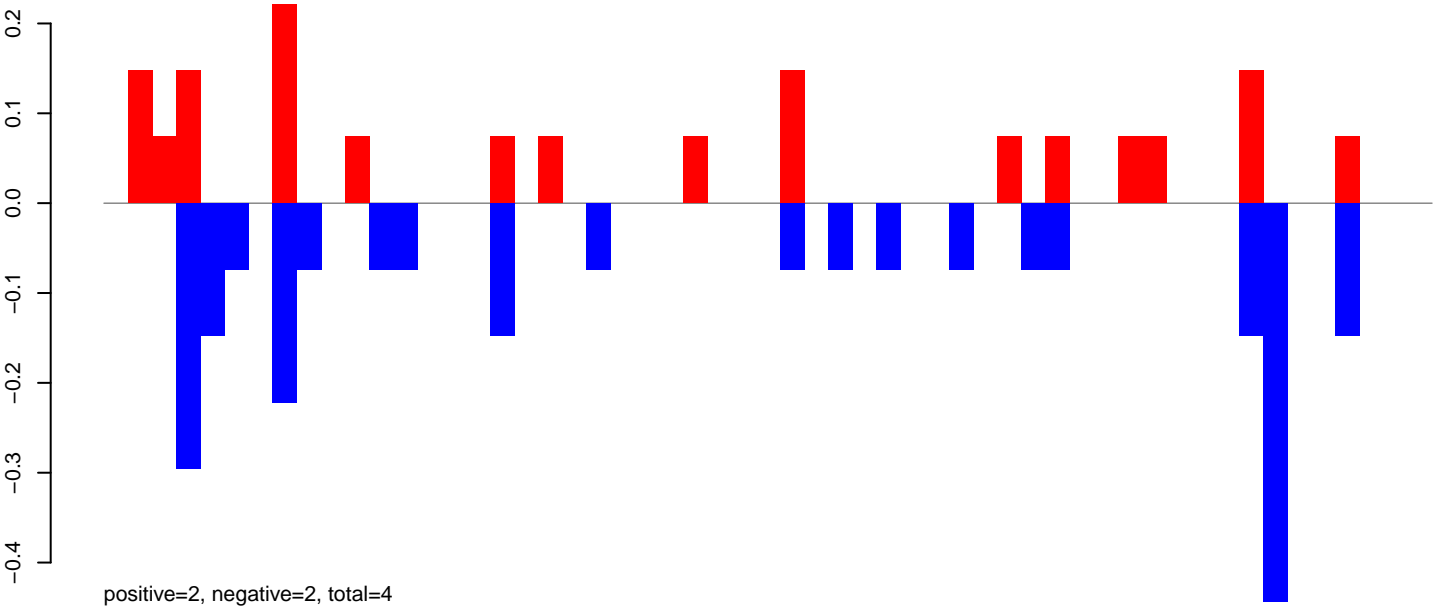
Window size=25, length=1260, TE@rnd-3_family-70@DNA_Sola-1@45:1-1260

0 200 400 600 800 1000 1200

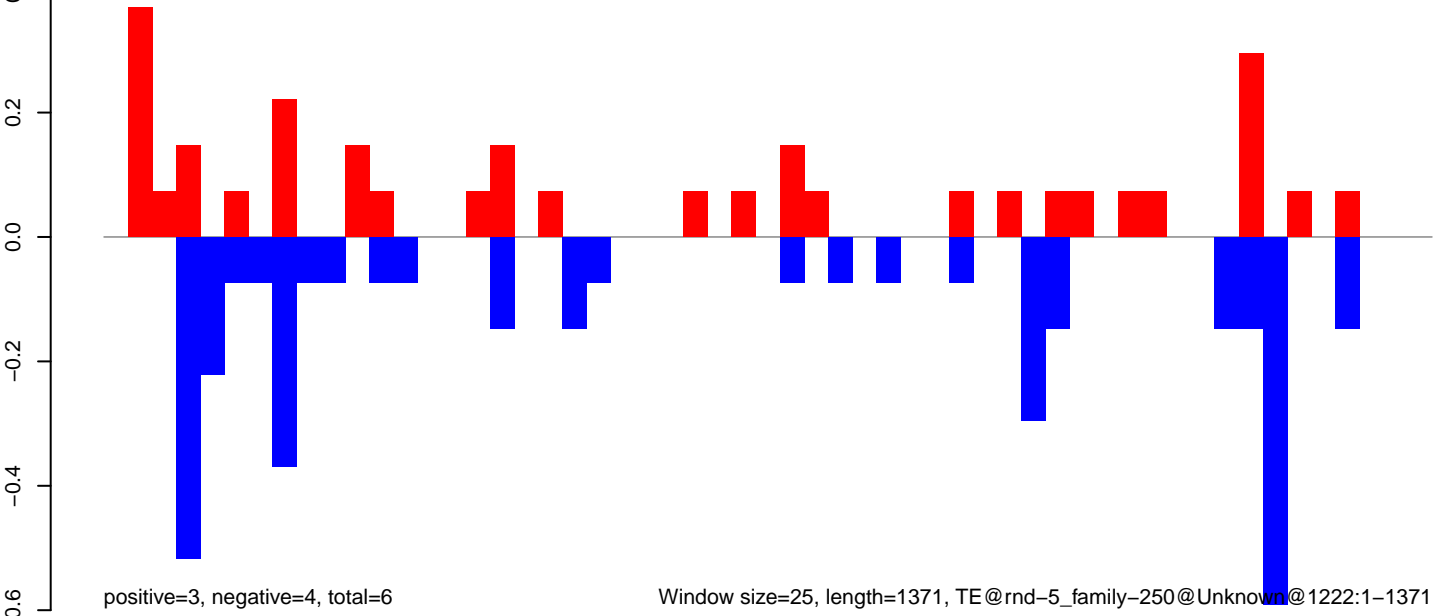
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



CuQuin_Testes_NL.18_23.rep

1.0
0.5
0.0
-0.5
-1.0

positive=4, negative=4, total=8

CuQuin_Testes_NL.24_35.rep

1.0
0.5
0.0
-0.5
-1.0

positive=9, negative=10, total=19

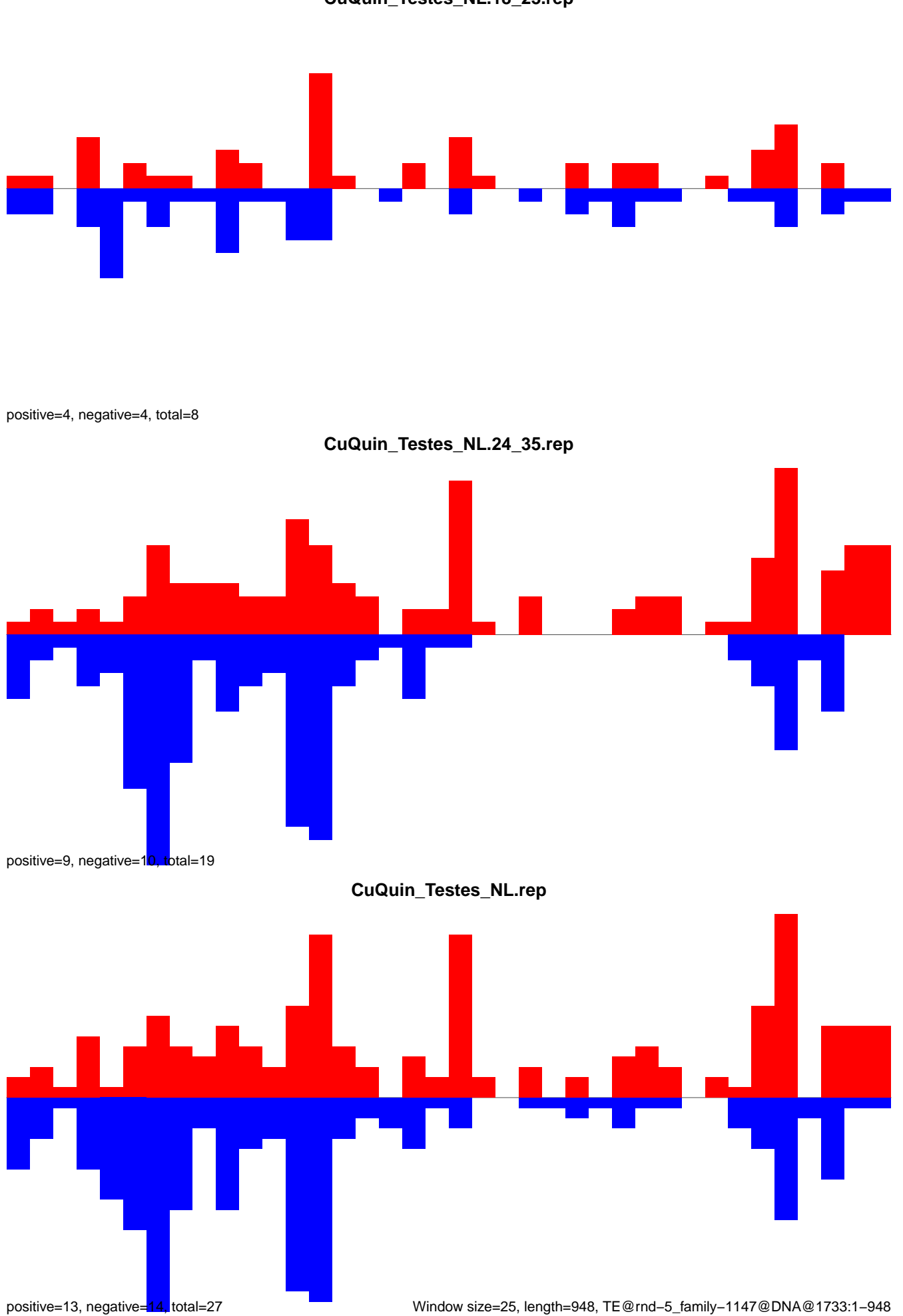
CuQuin_Testes_NL.rep

1.0
0.5
0.0
-0.5
-1.0
-1.5

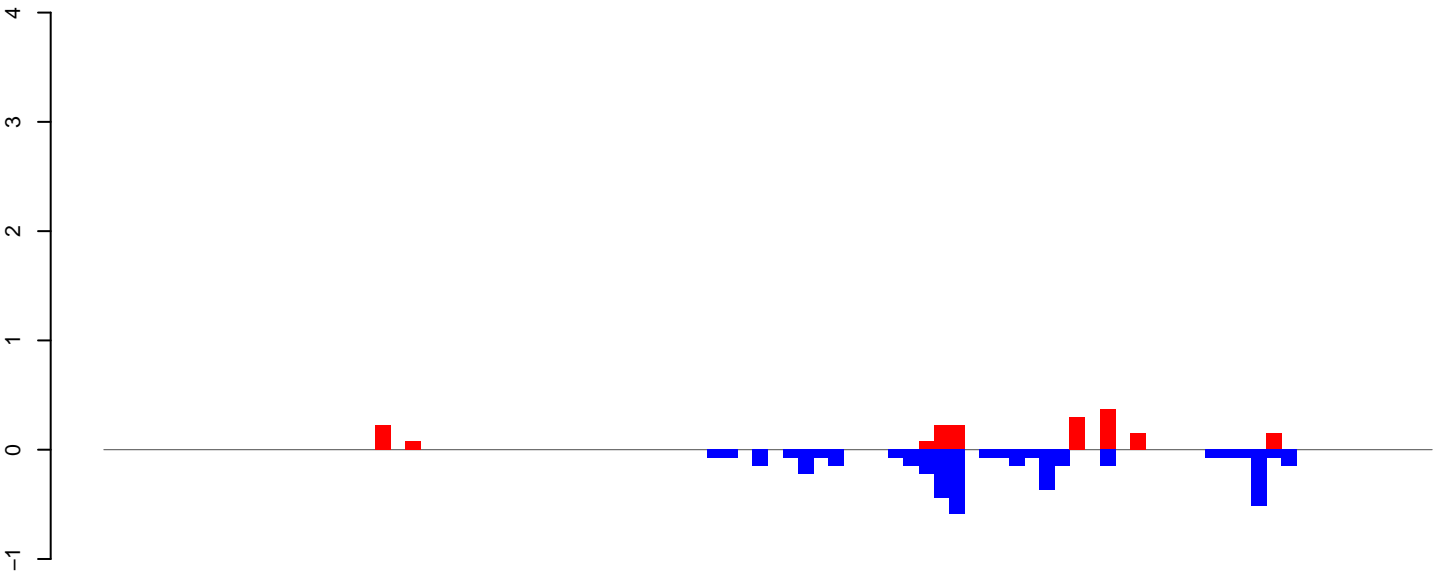
positive=13, negative=14, total=27

Window size=25, length=948, TE@rnd-5_family-1147@DNA@1733:1-948

0 200 400 600 800 1000

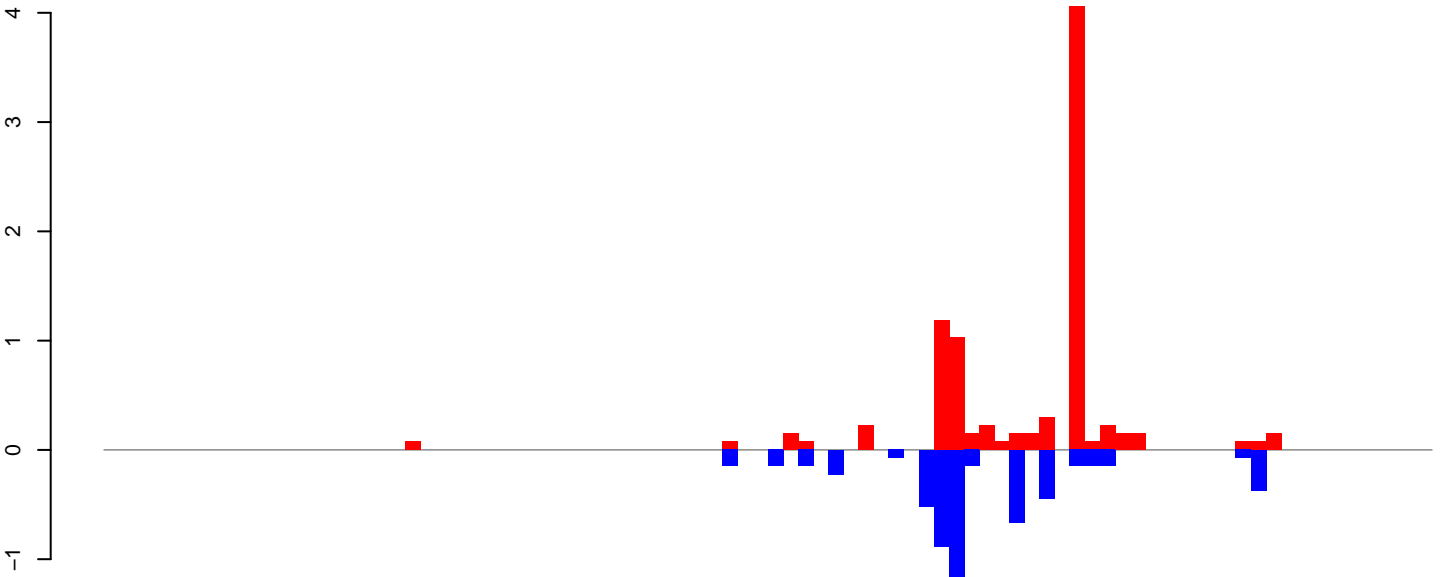


CuQuin_Testes_NL.18_23.rep



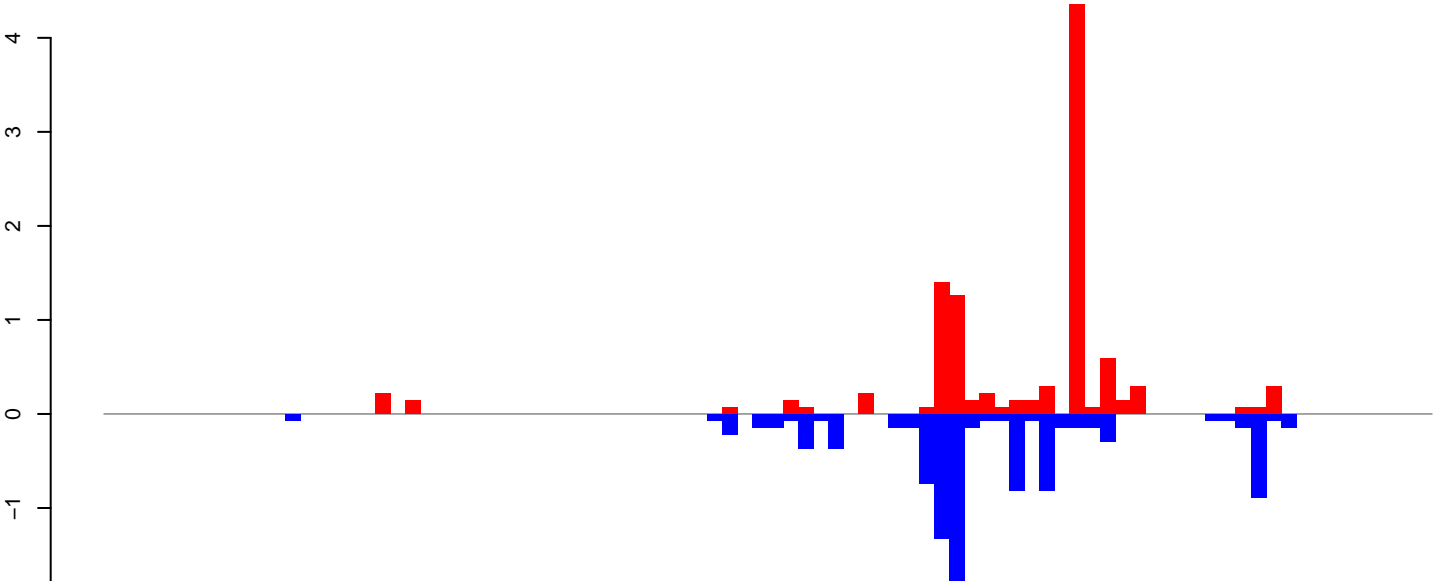
positive=2, negative=4, total=6

CuQuin_Testes_NL.24_35.rep



positive=9, negative=6, total=14

CuQuin_Testes_NL.rep

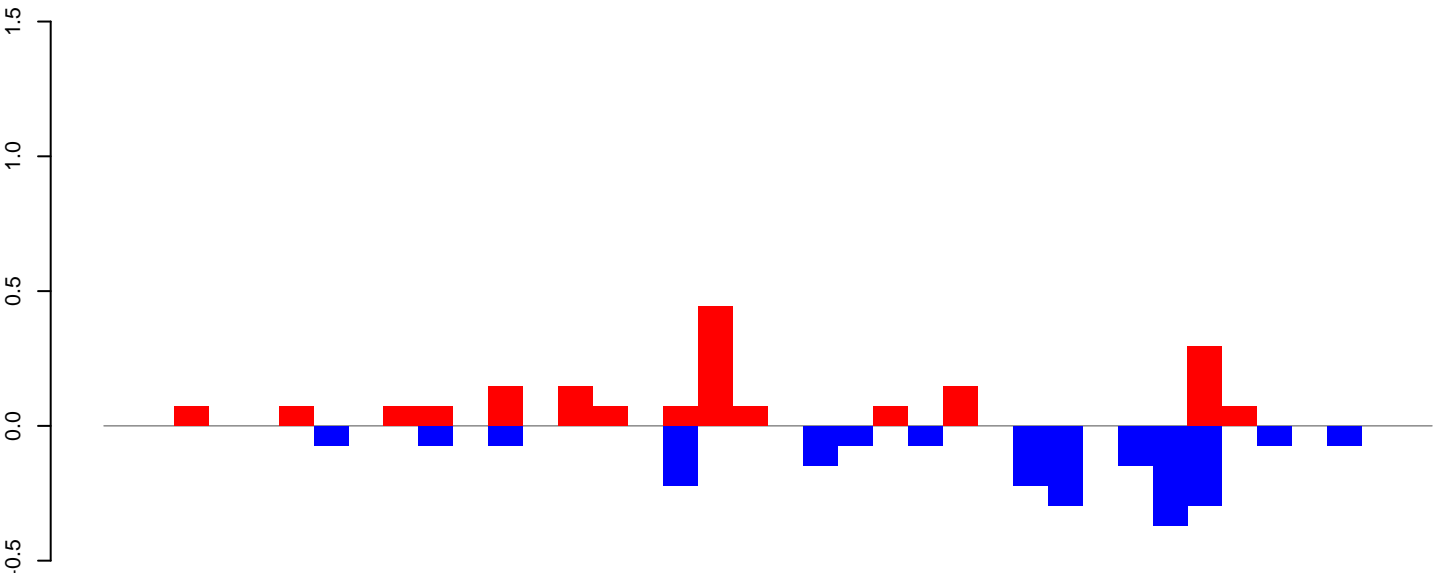


positive=11, negative=10, total=21

Window size=25, length=2198, TE@rnd-4_family-800@DNA@2308:1-2198

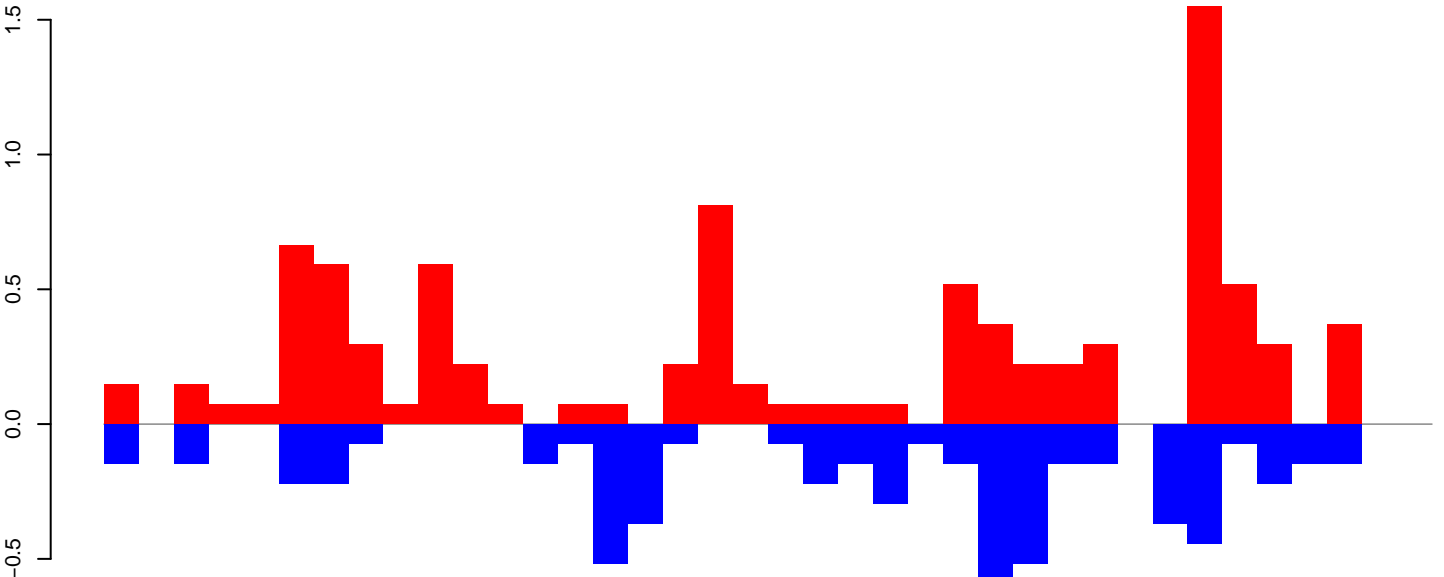
0 500 1000 1500 2000

CuQuin_Testes_NL.18_23.rep



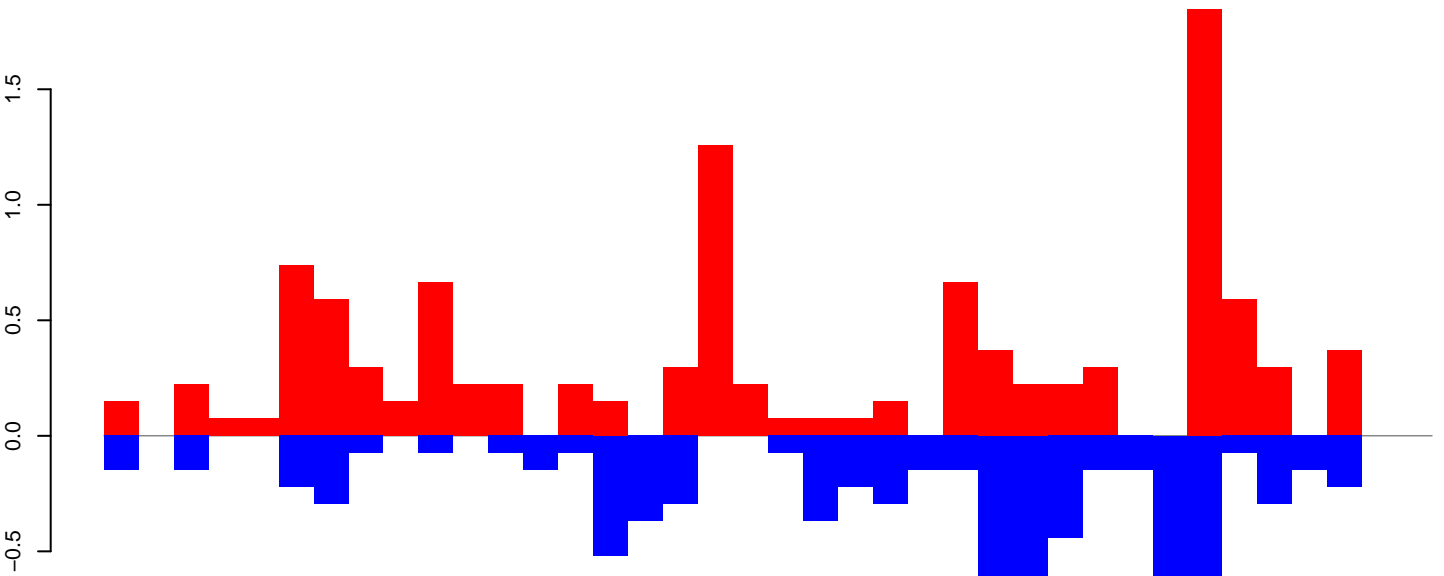
positive=2, negative=2, total=4

CuQuin_Testes_NL.24_35.rep



positive=9, negative=6, total=15

CuQuin_Testes_NL.rep

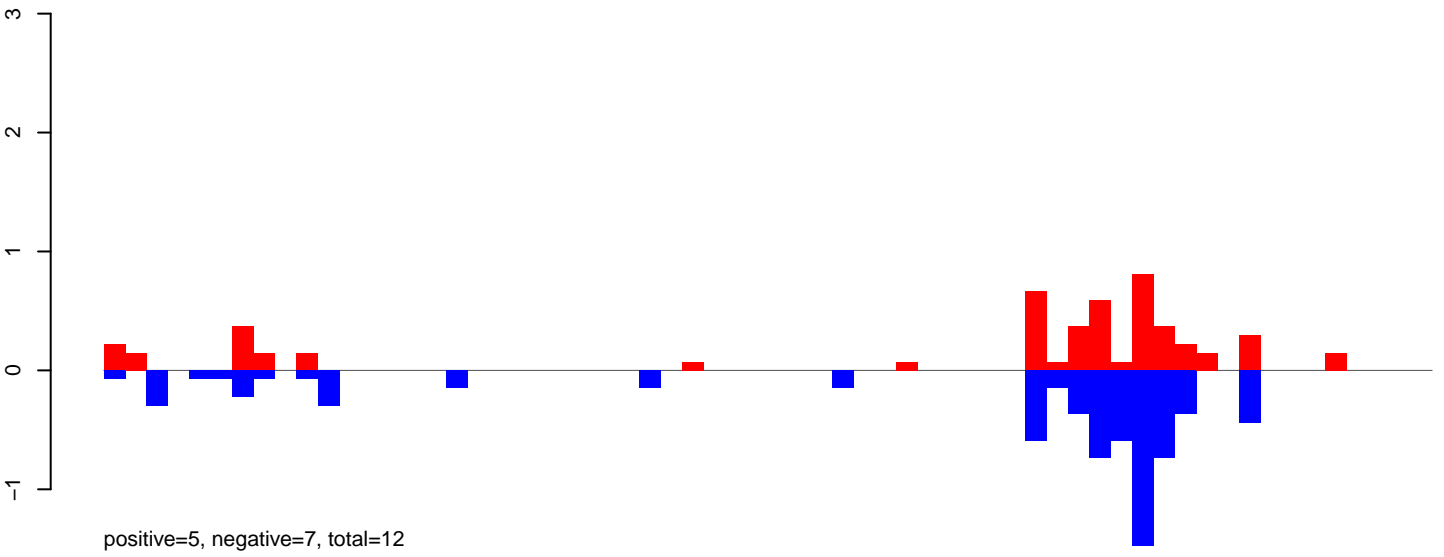


positive=11, negative=8, total=19

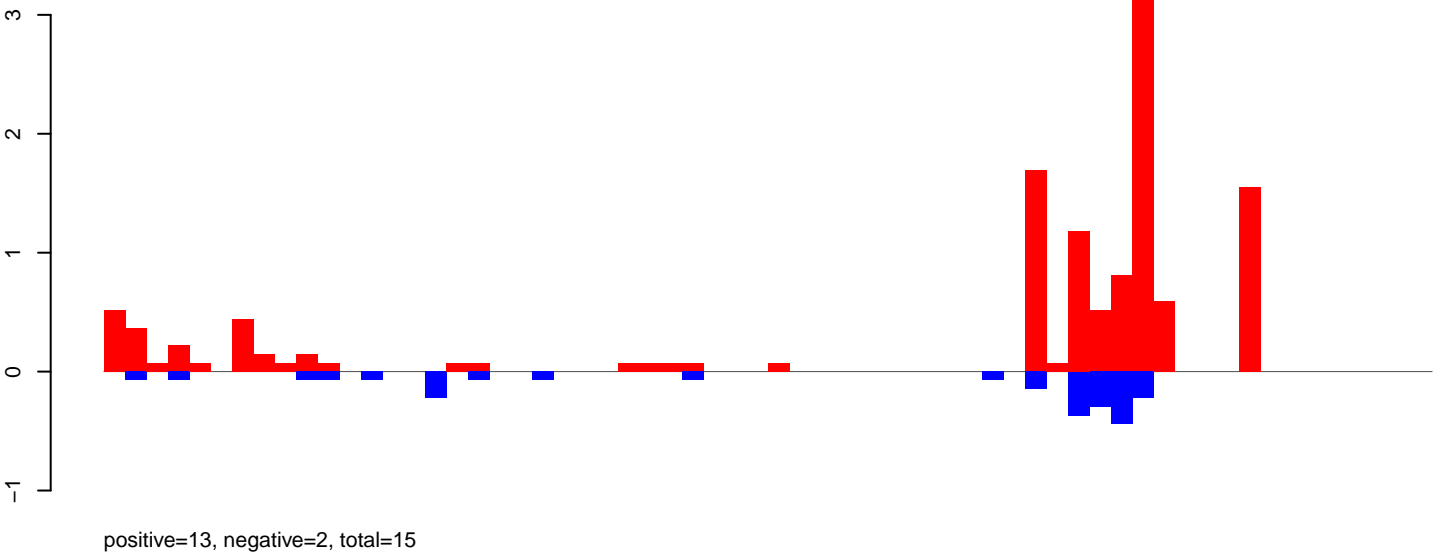
Window size=25, length=929, TE@rnc-4_family-798@LTR_Pao@2234:1-929

0 200 400 600 800 1000

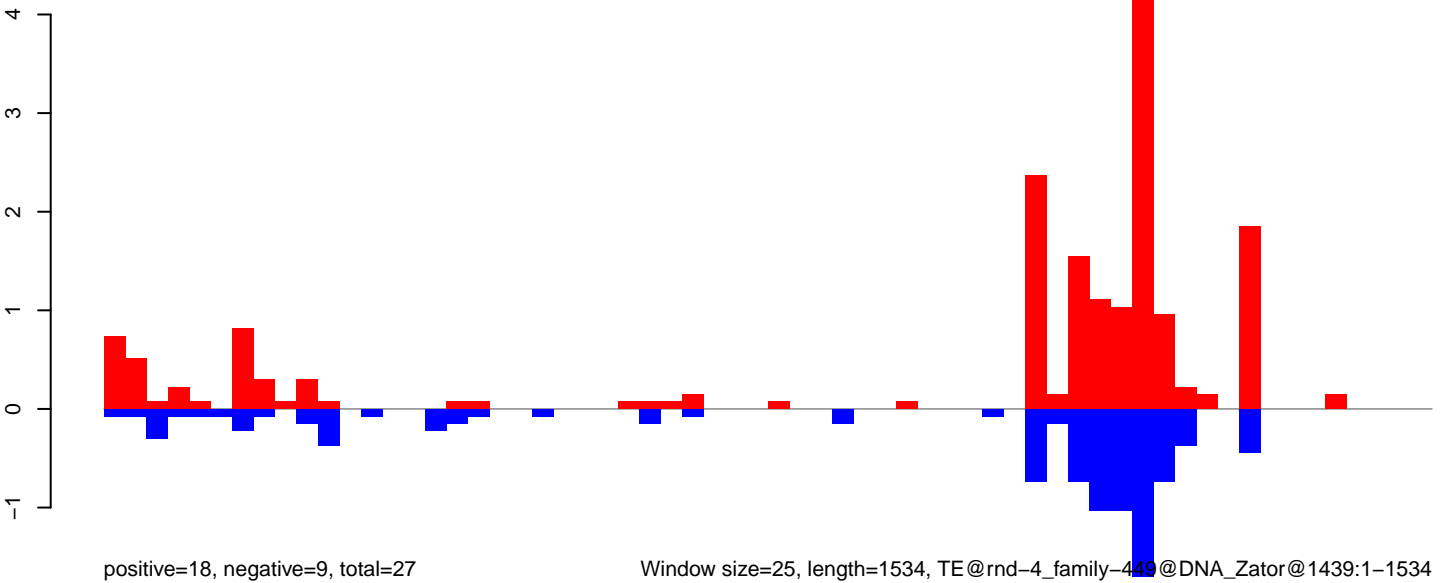
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

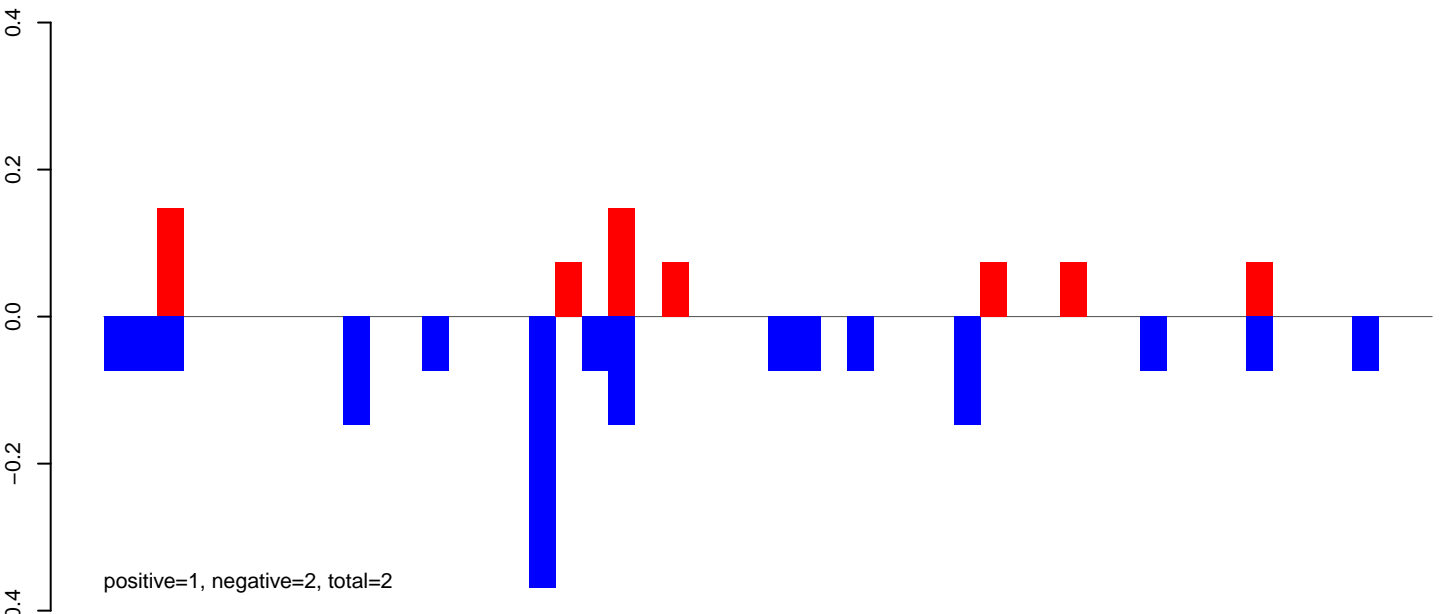


CuQuin_Testes_NL.rep

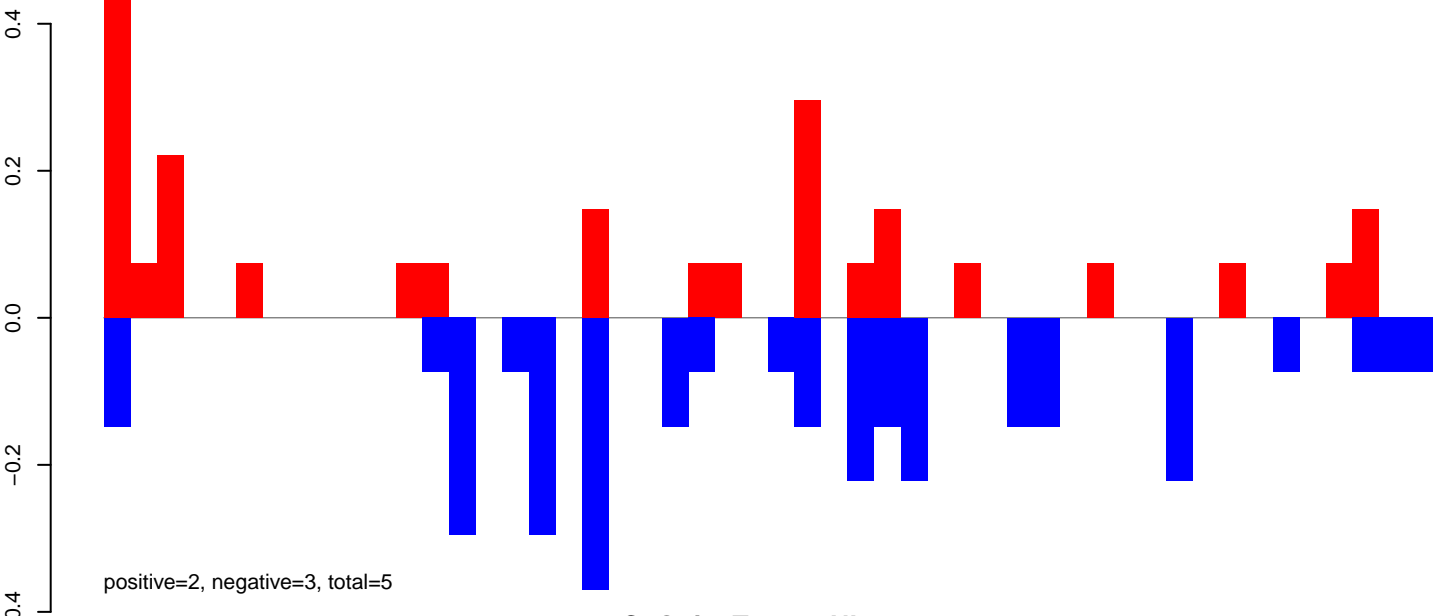


Window size=25, length=1534, TE@rnd-4_family-449@DNA_Zator@1439:1-1534

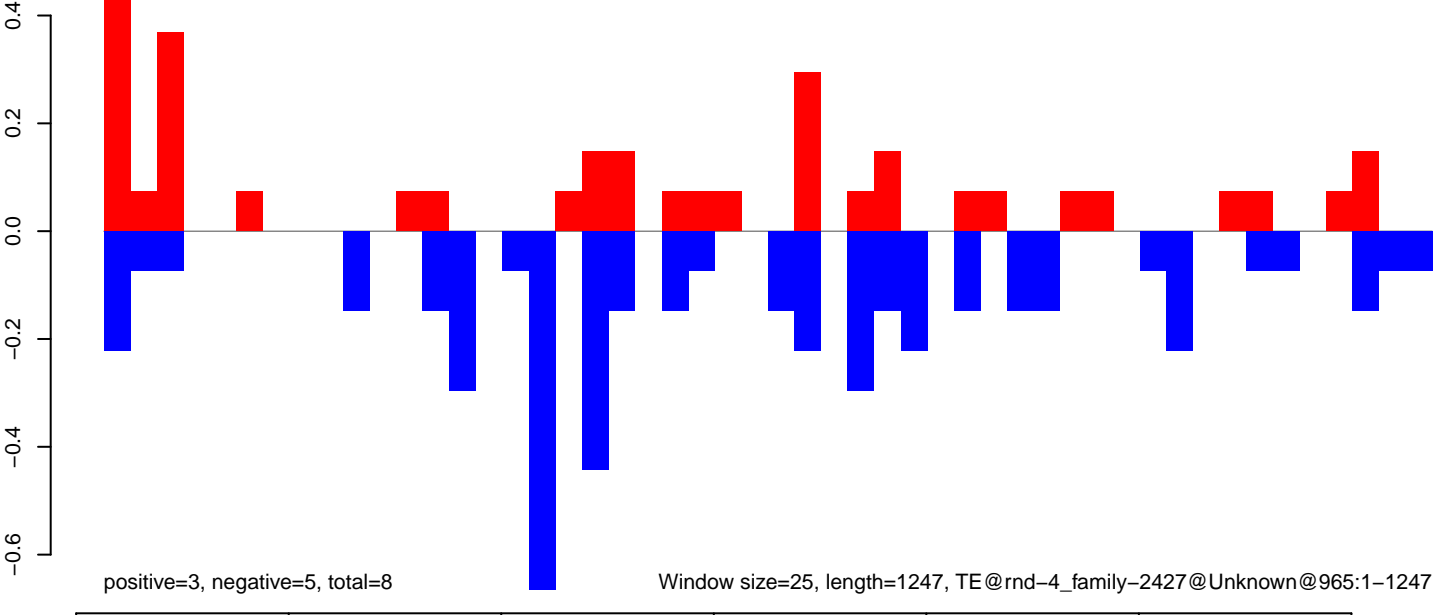
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

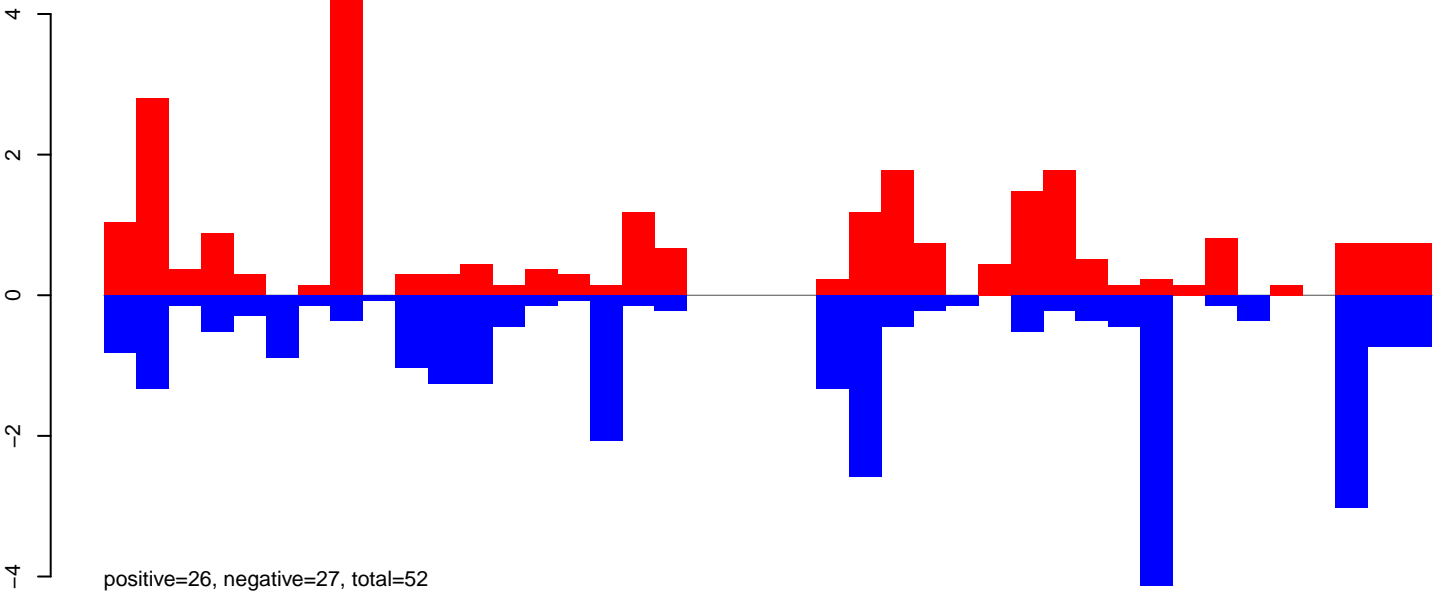


CuQuin_Testes_NL.rep

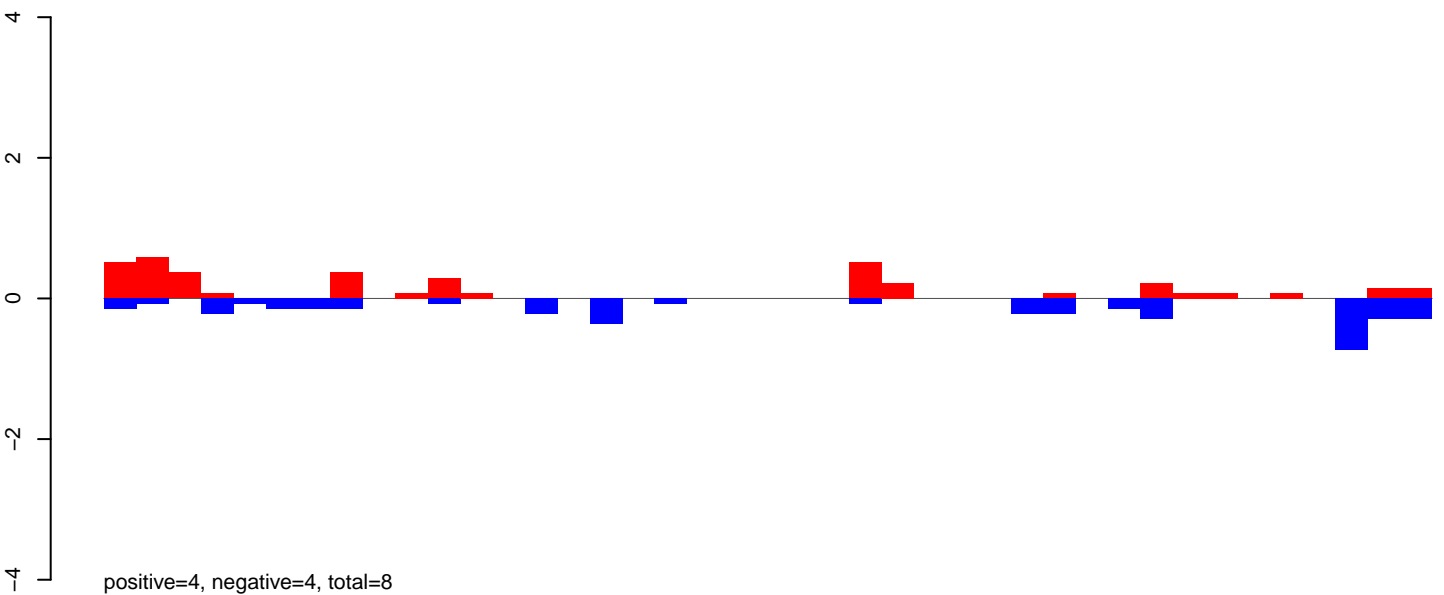


Window size=25, length=1247, TE@rnd-4_family-2427@Unknown@965:1-1247

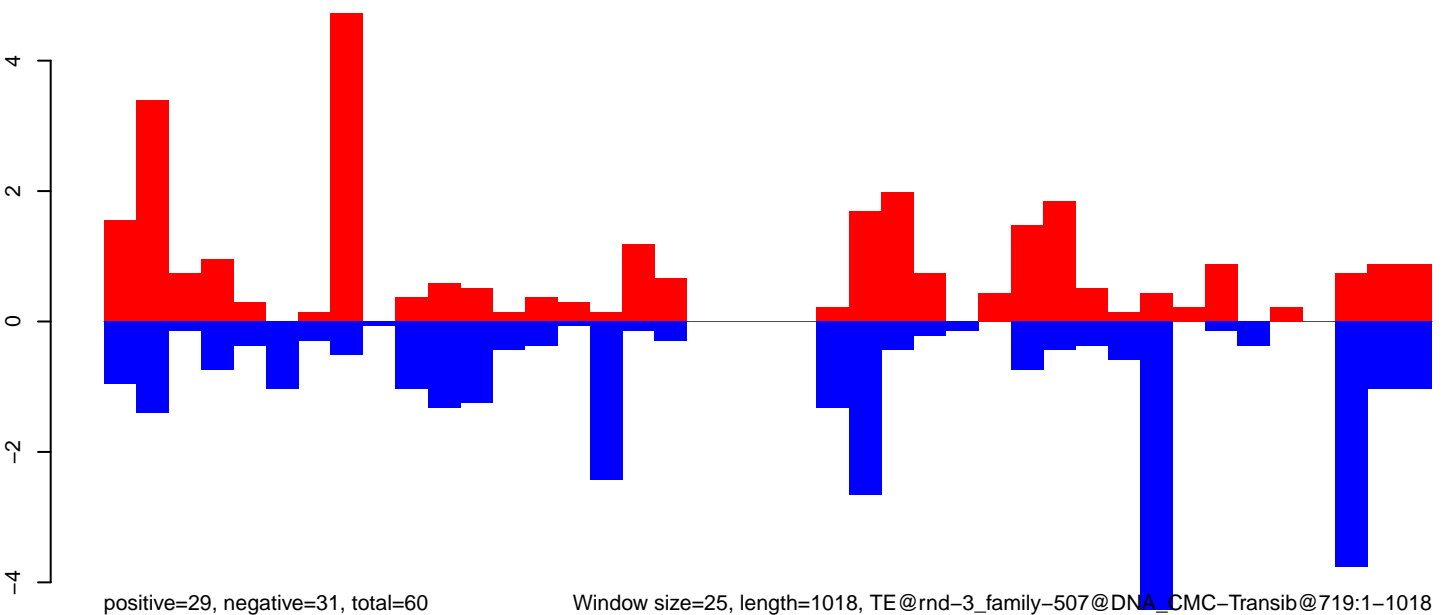
CuQuin_Testes_NL.18_23.rep



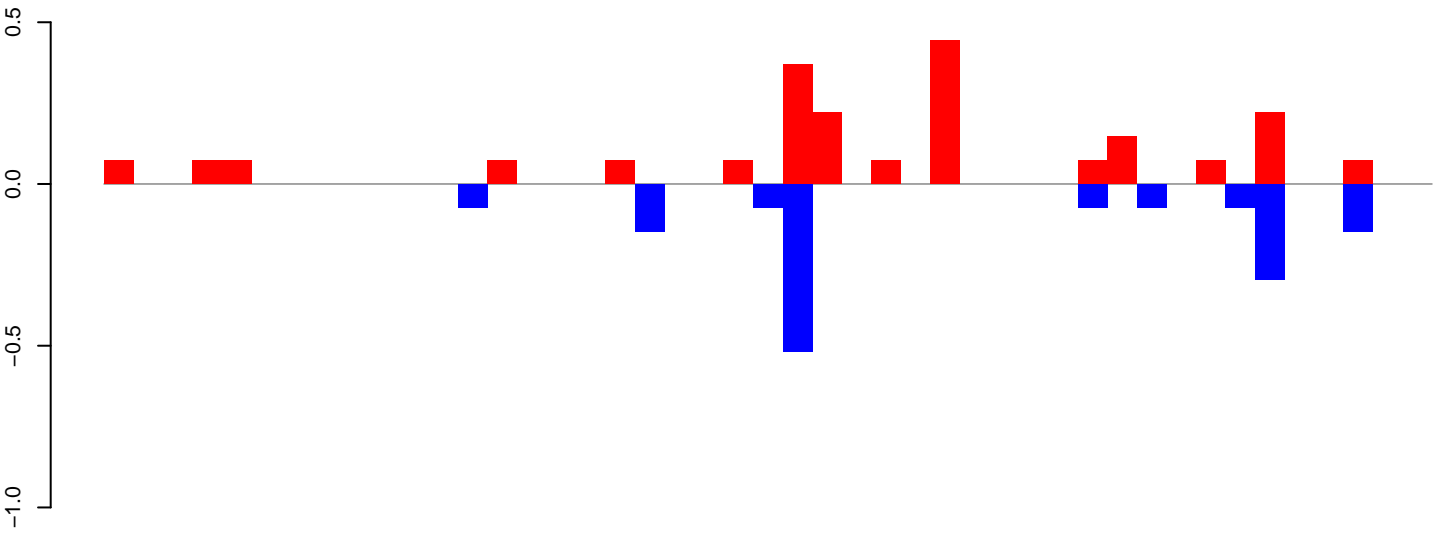
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

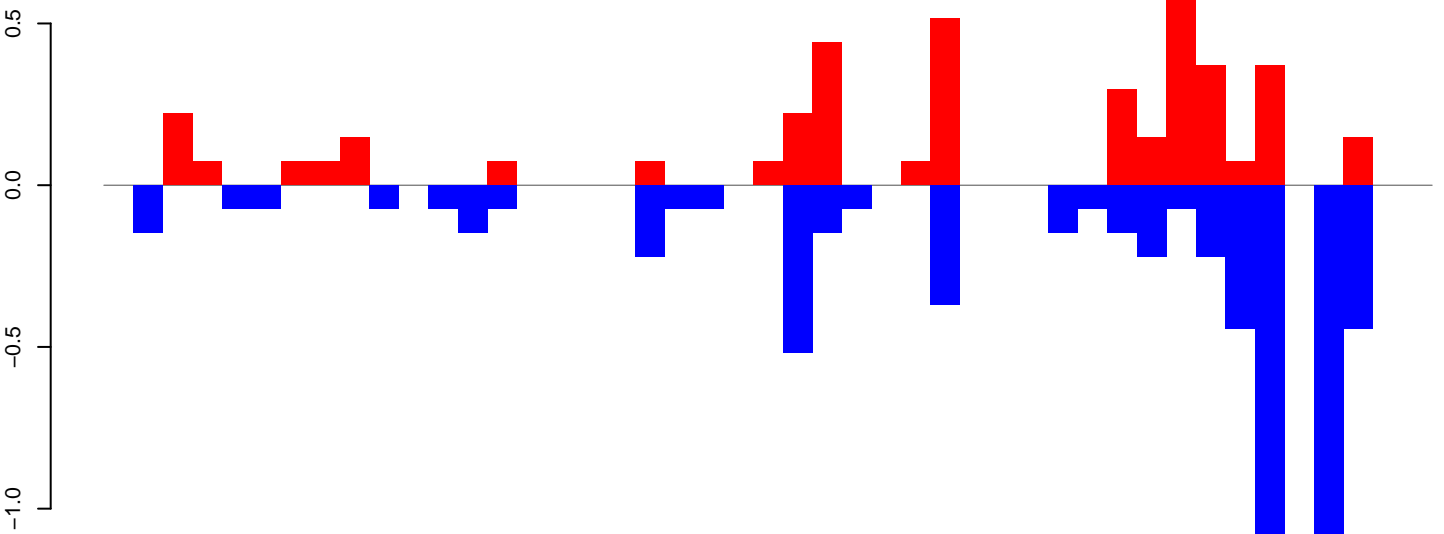


CuQuin_Testes_NL.18_23.rep



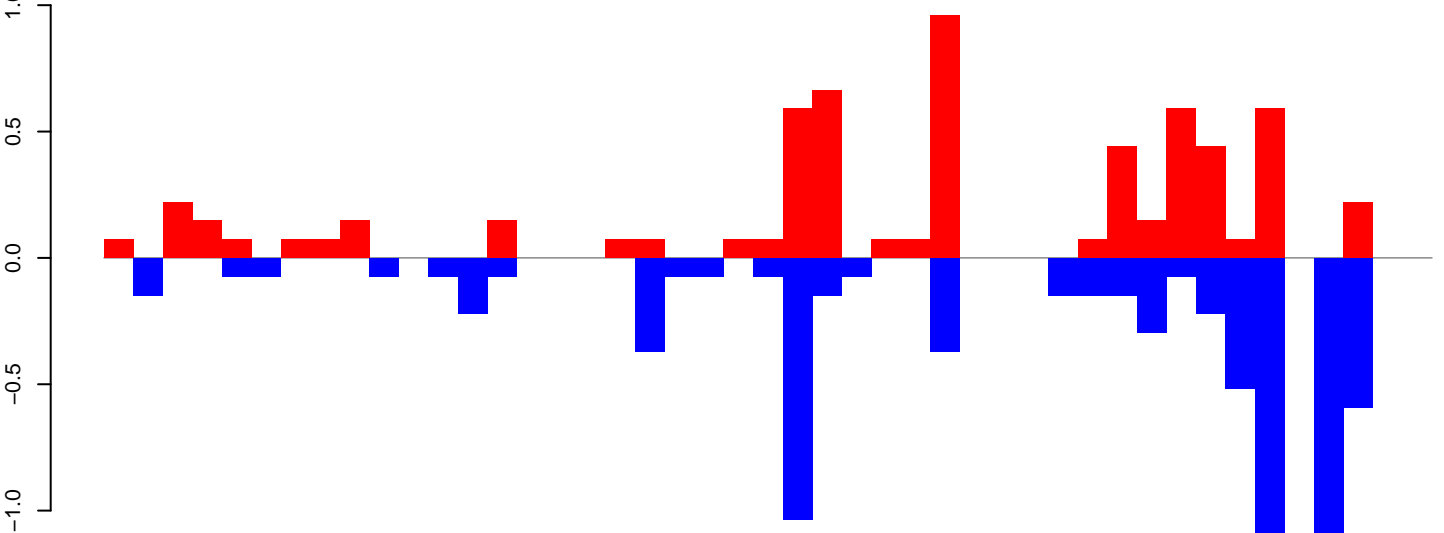
positive=2, negative=1, total=4

CuQuin_Testes_NL.24_35.rep



positive=4, negative=6, total=10

CuQuin_Testes_NL.rep

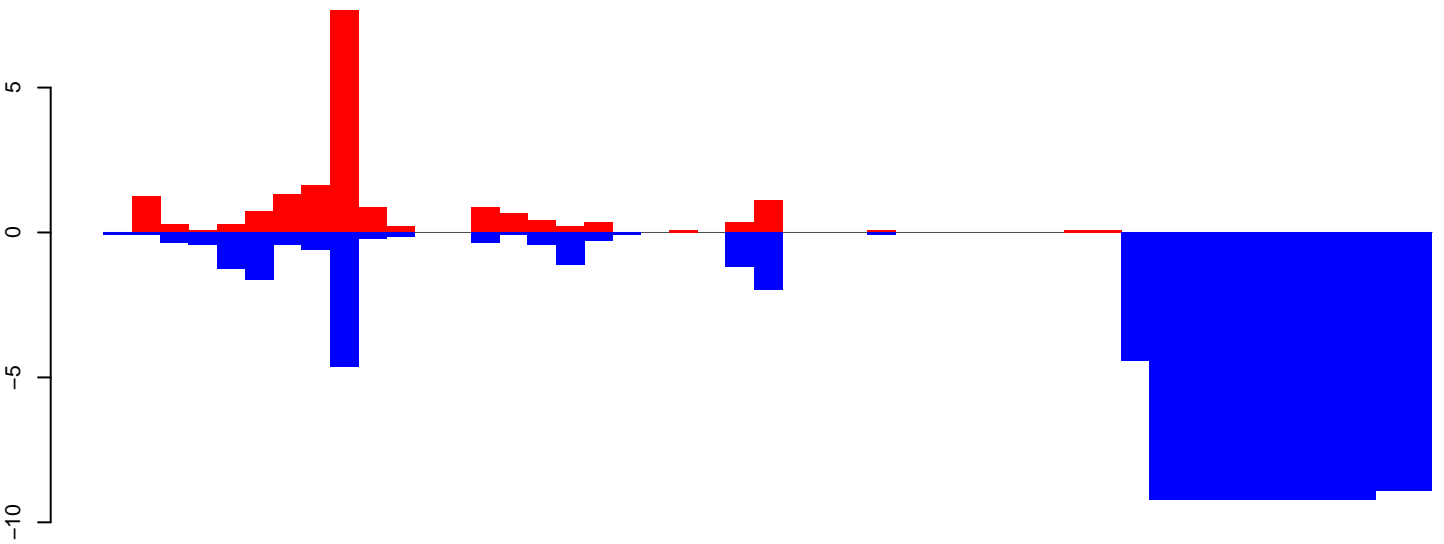


positive=6, negative=8, total=14

Window size=25, length=1118, TE@rnd-3_family-1133@DNA@725:1-1118

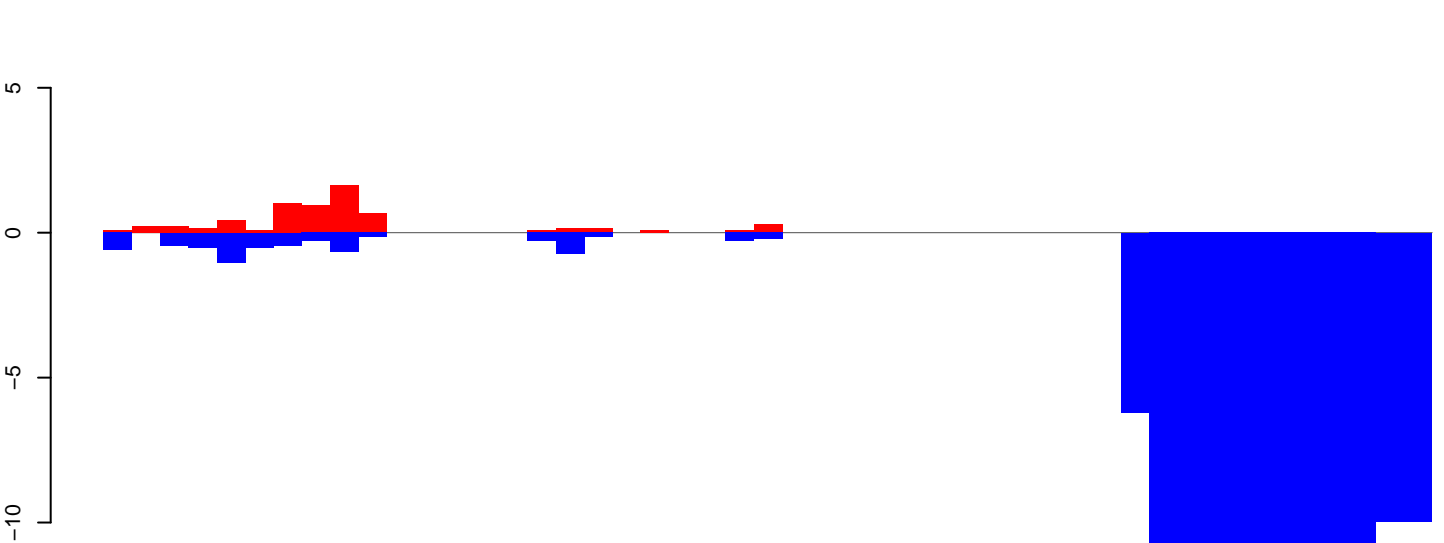
0 200 400 600 800 1000

CuQuin_Testes_NL.18_23.rep



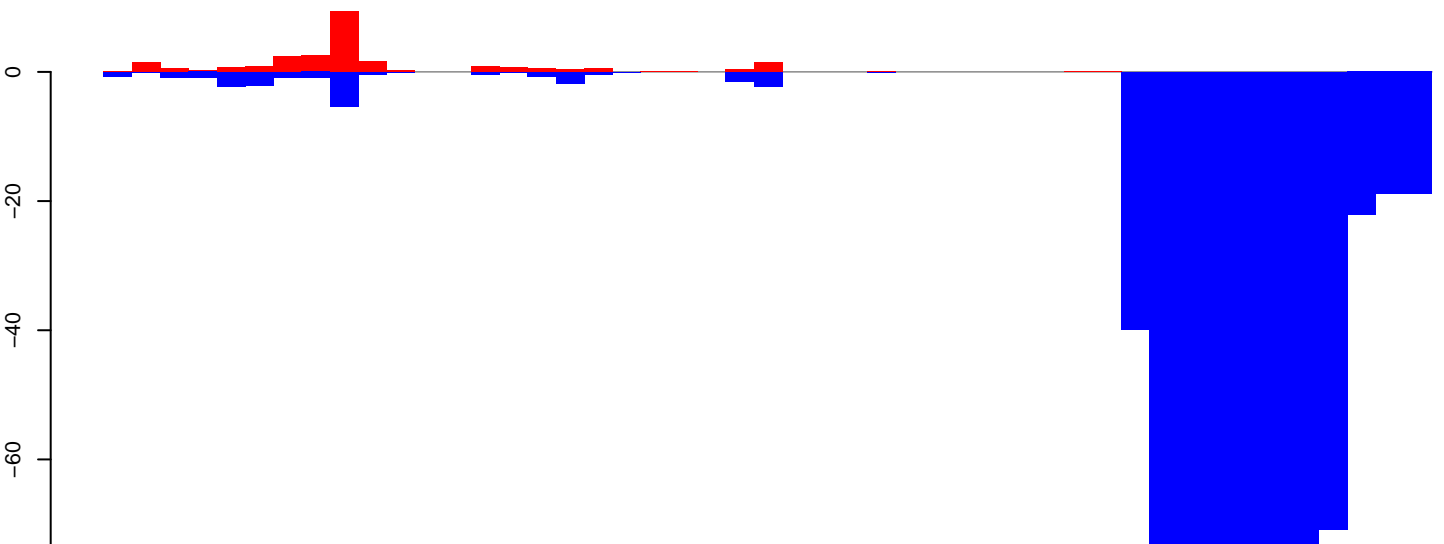
positive=19, negative=112, total=130

CuQuin_Testes_NL.24_35.rep



positive=6, negative=136, total=142

CuQuin_Testes_NL.rep

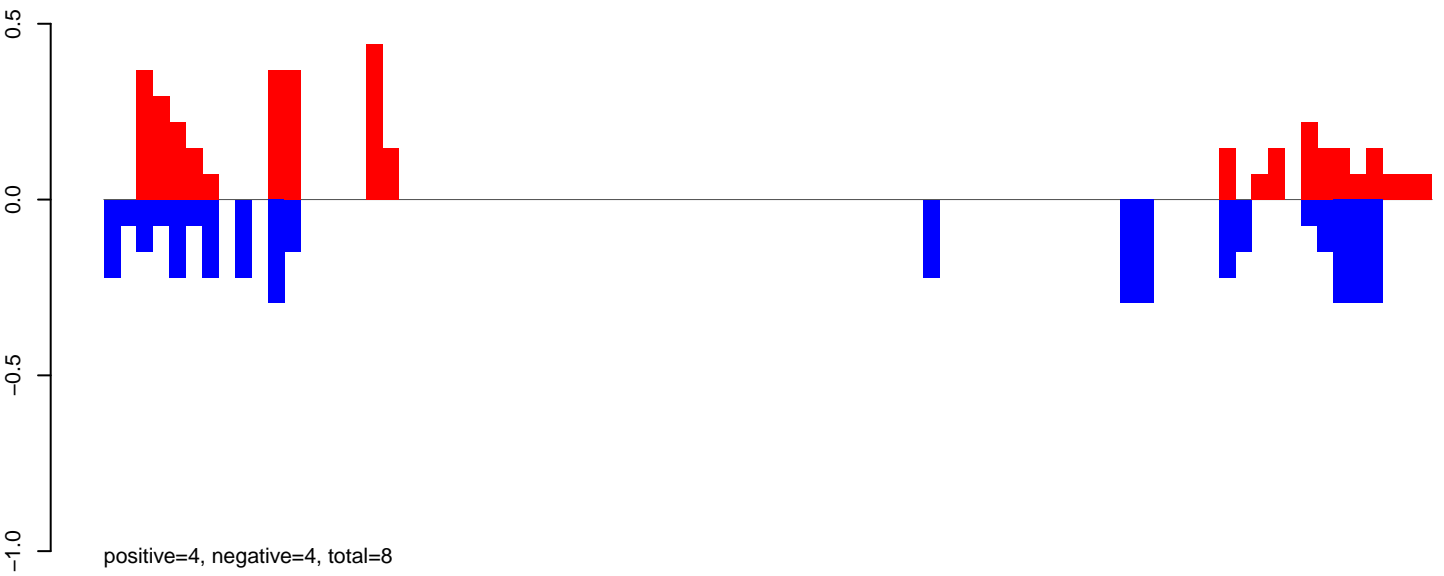


positive=25, negative=691, total=716

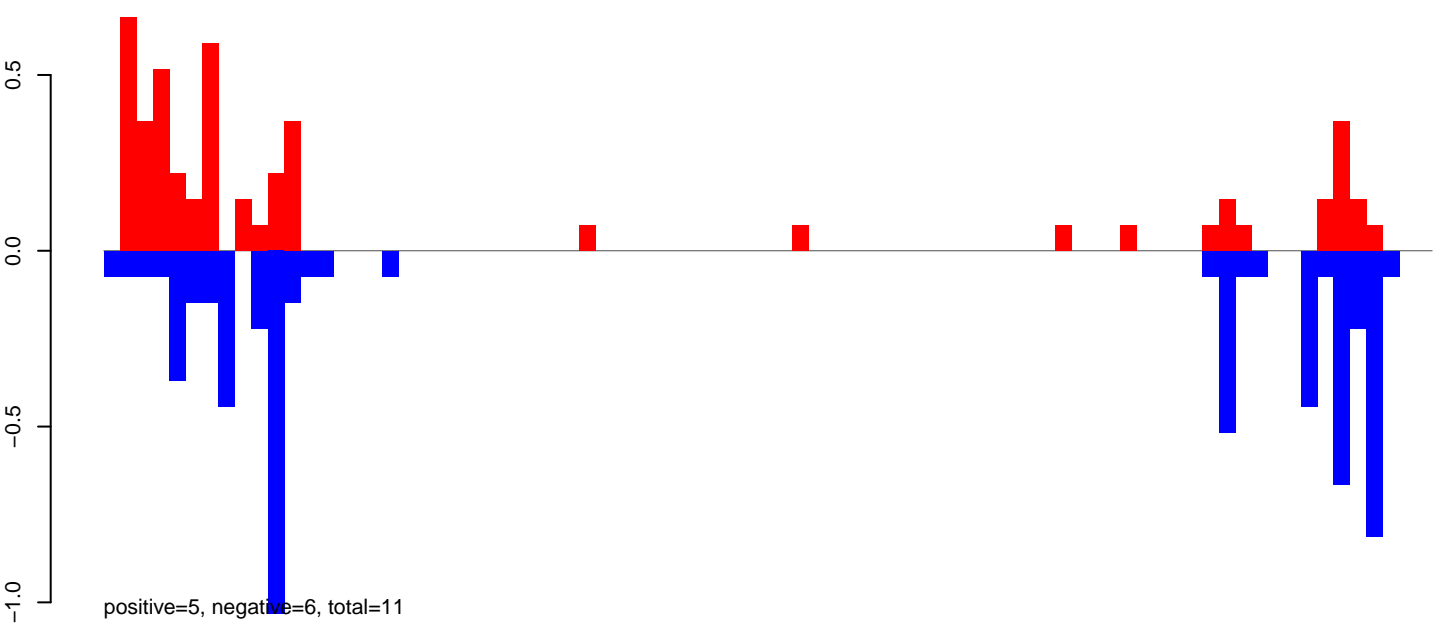
Window size=25, length=1171, TE@rnd-5_family-58@DNA_Sola-2@1074:1-1171

0 200 400 600 800 1000 1200

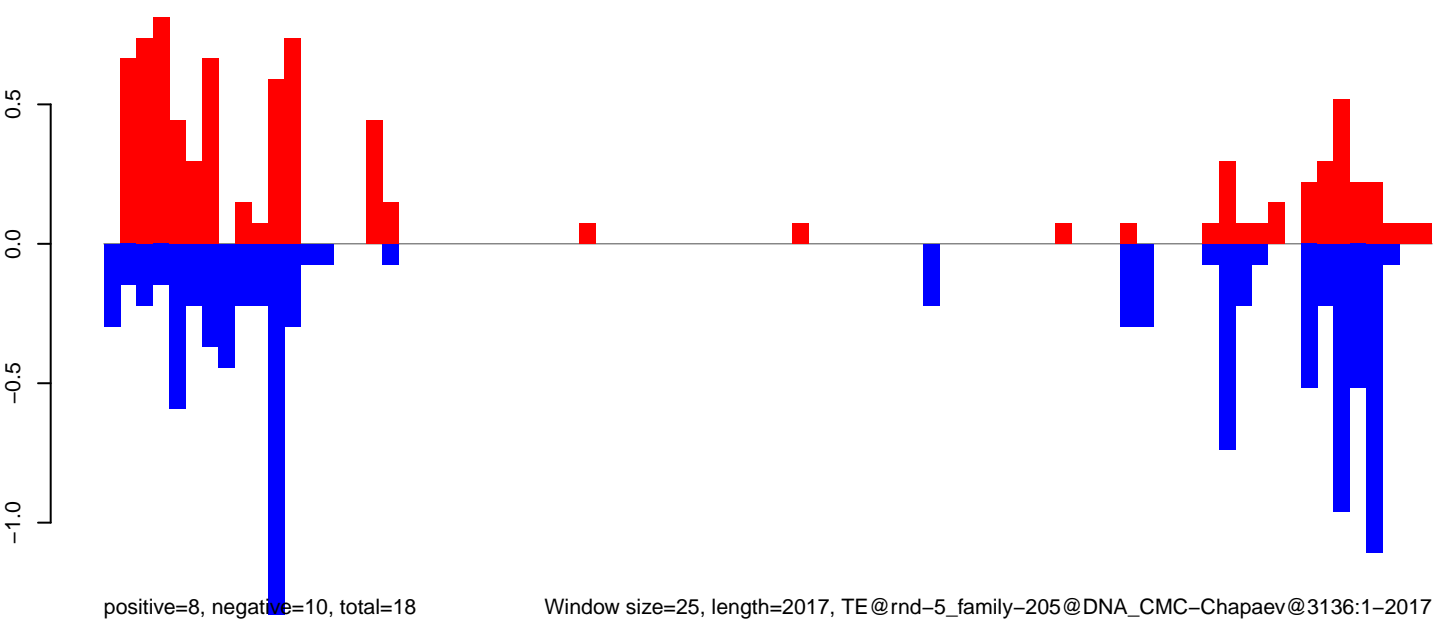
CuQuin_Testes_NL.18_23.rep



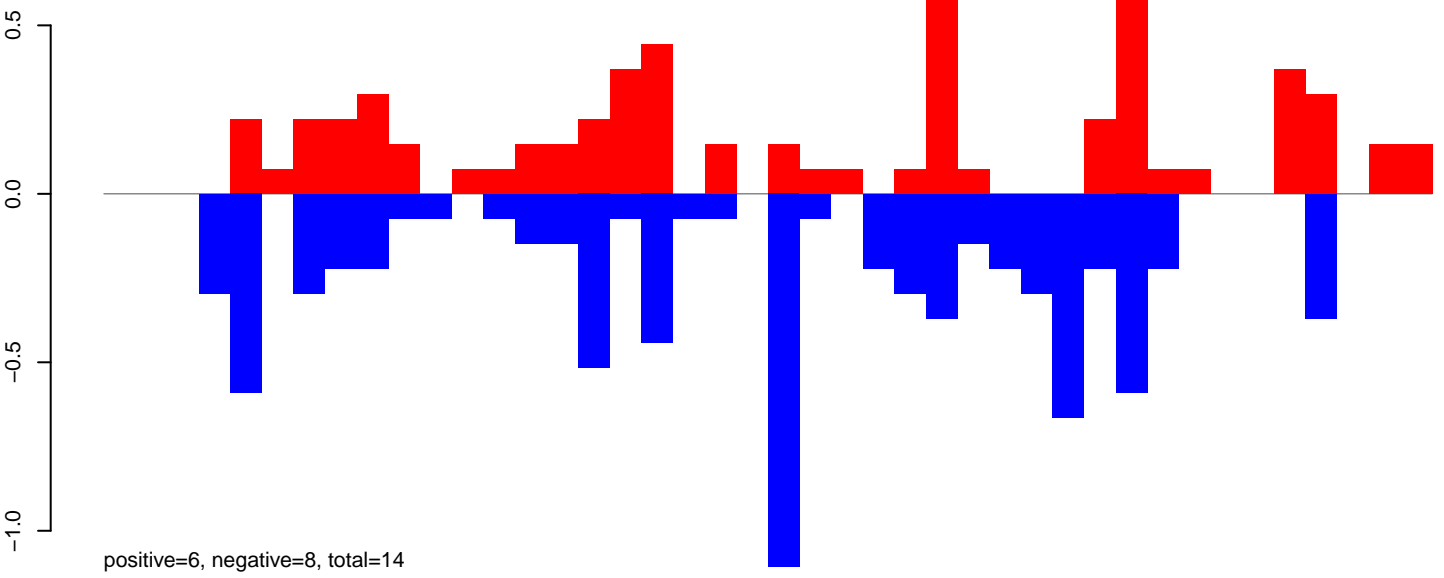
CuQuin_Testes_NL.24_35.rep



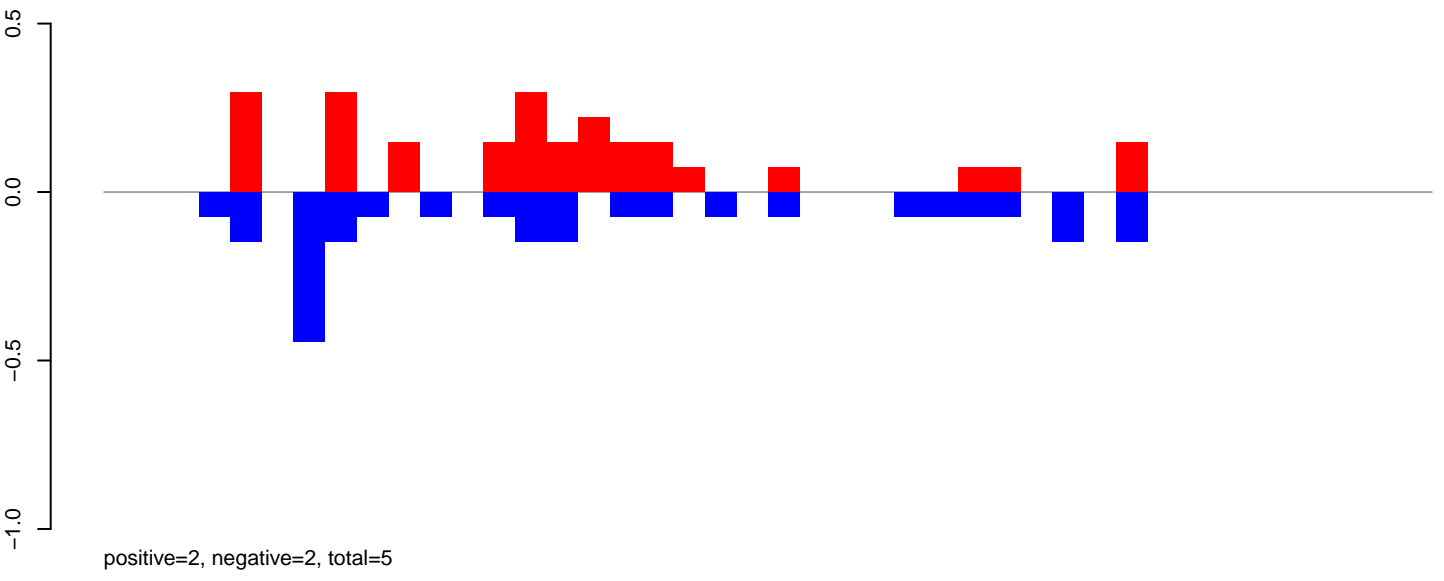
CuQuin_Testes_NL.rep



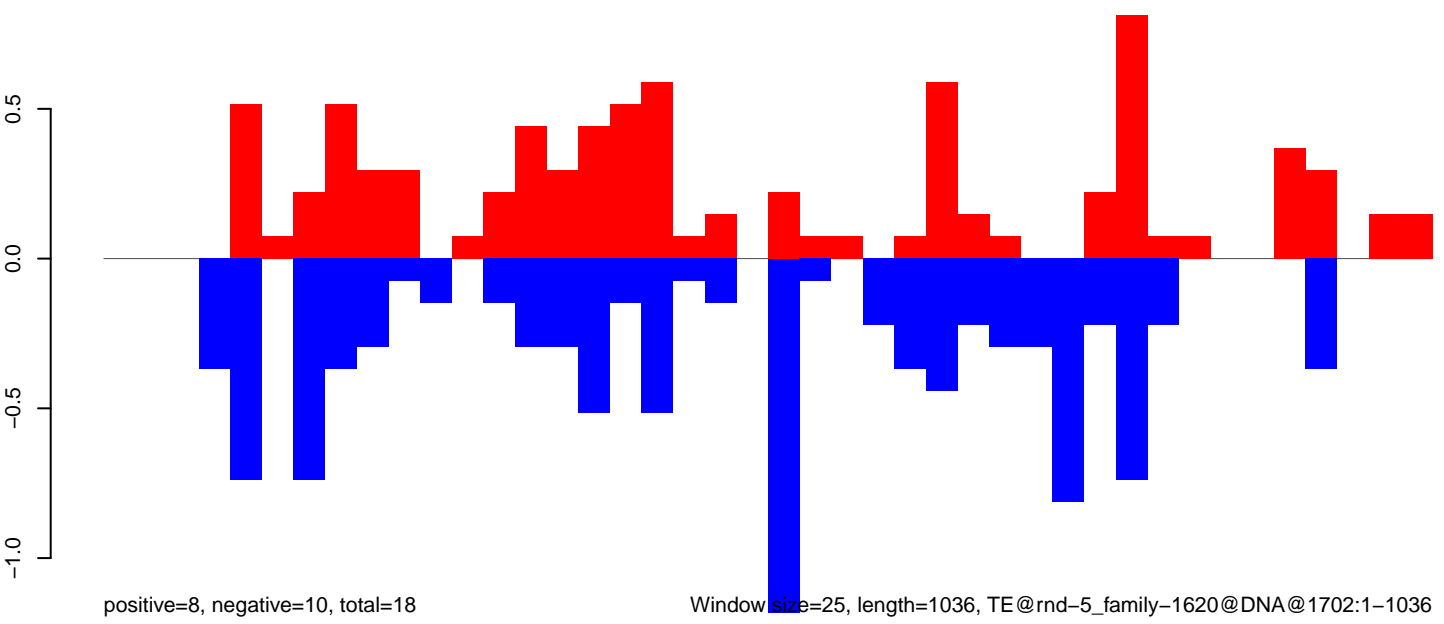
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



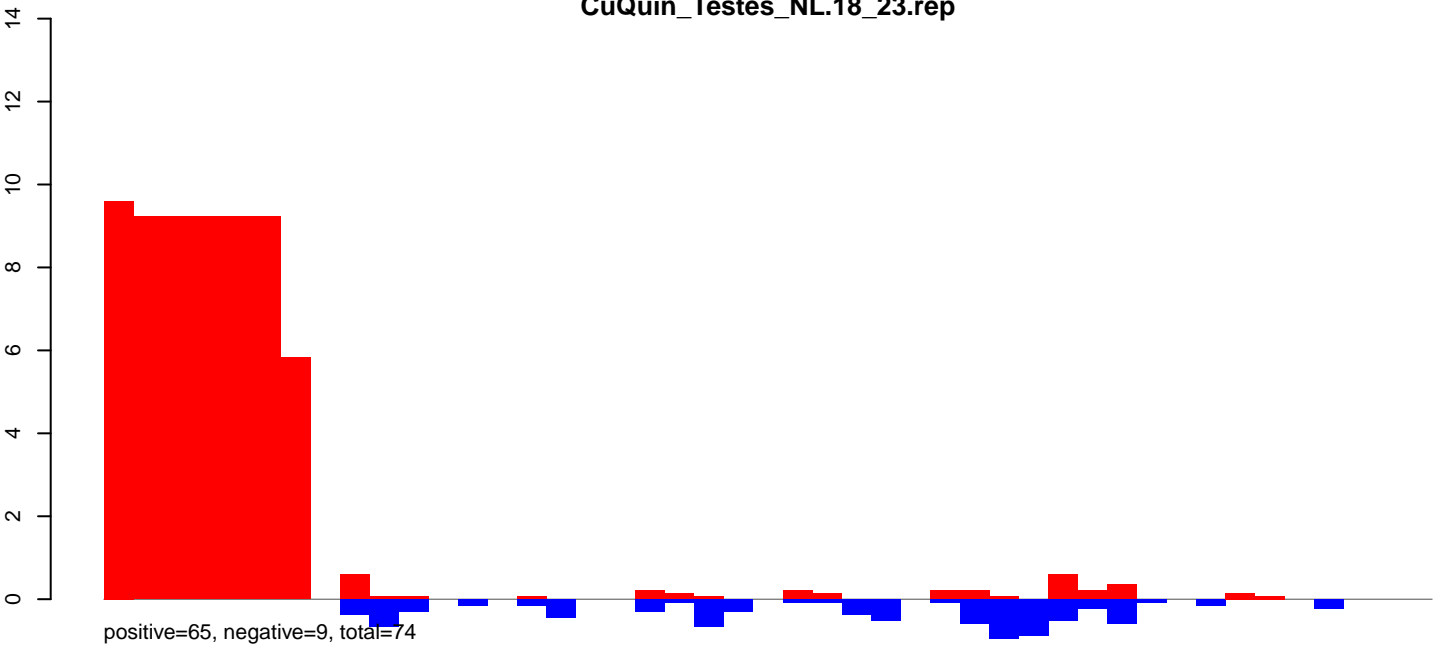
CuQuin_Testes_NL.rep



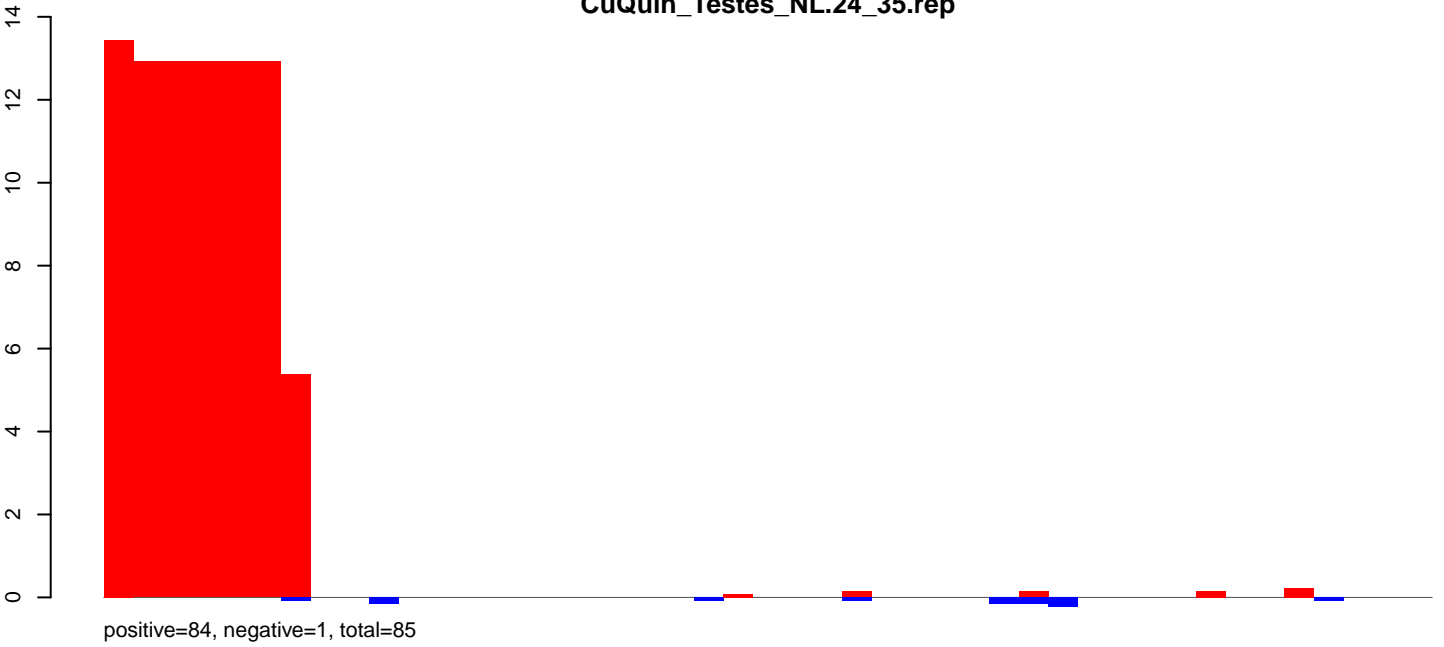
Window size=25, length=1036, TE@rnd-5_family-1620@DNA@1702:1-1036

0 200 400 600 800 1000

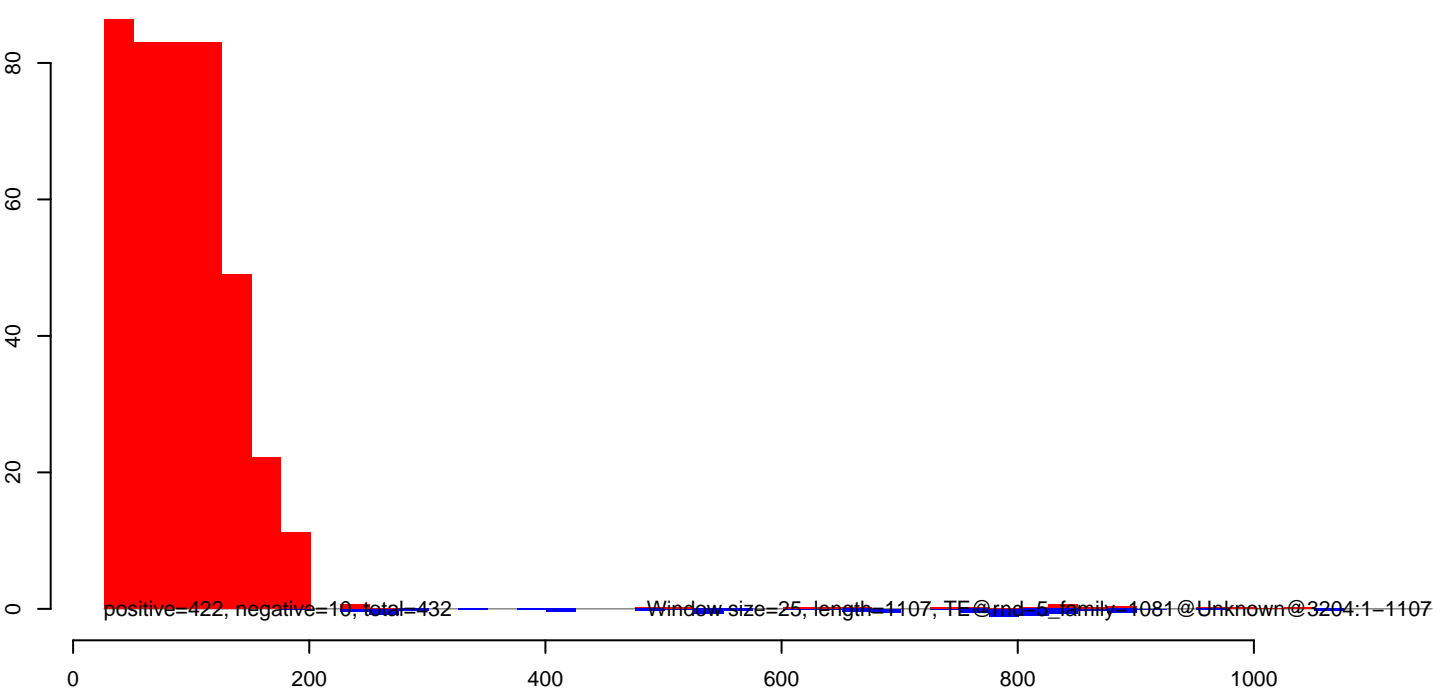
CuQuin_Testes_NL.18_23.rep



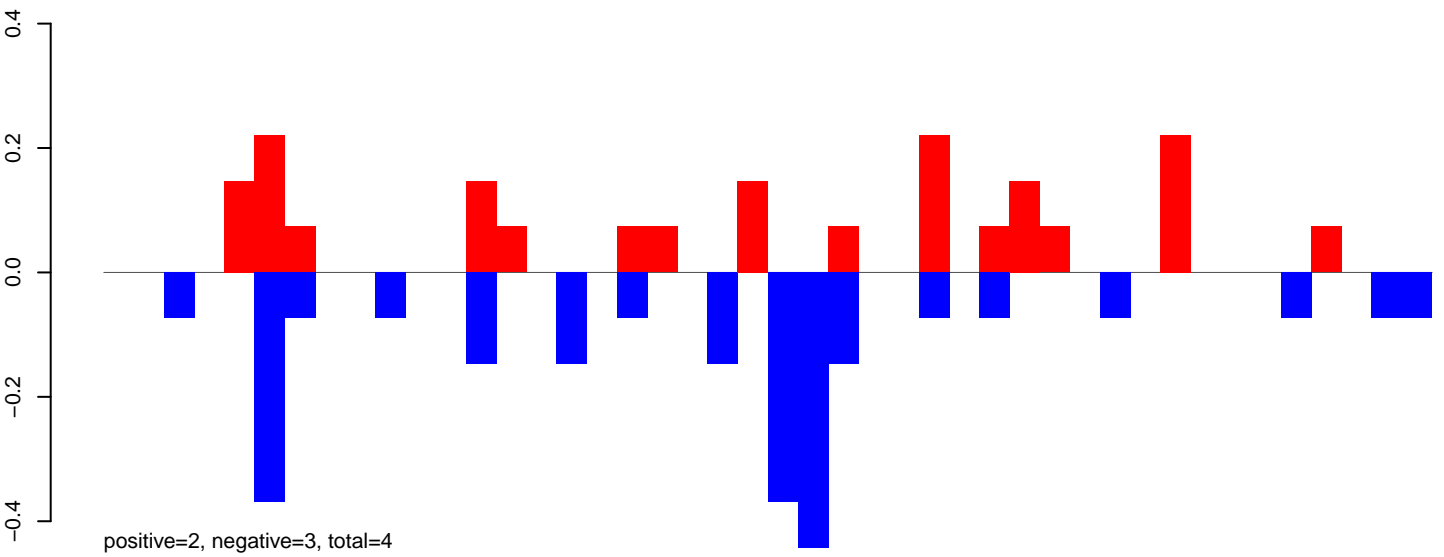
CuQuin_Testes_NL.24_35.rep



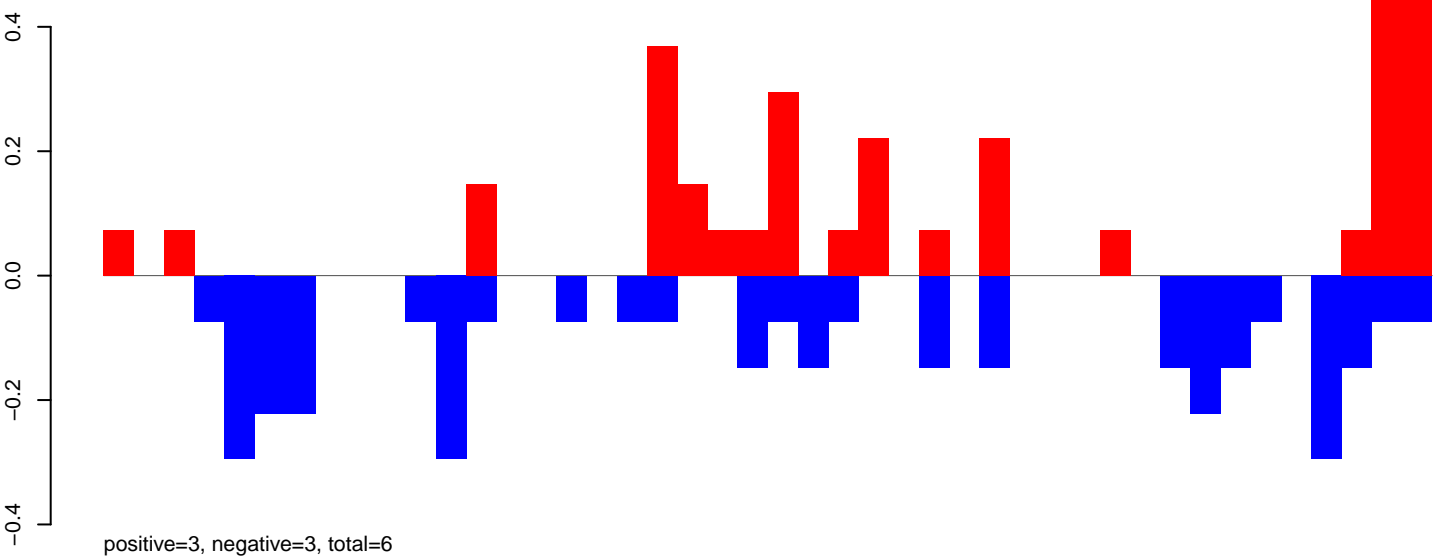
CuQuin_Testes_NL.rep



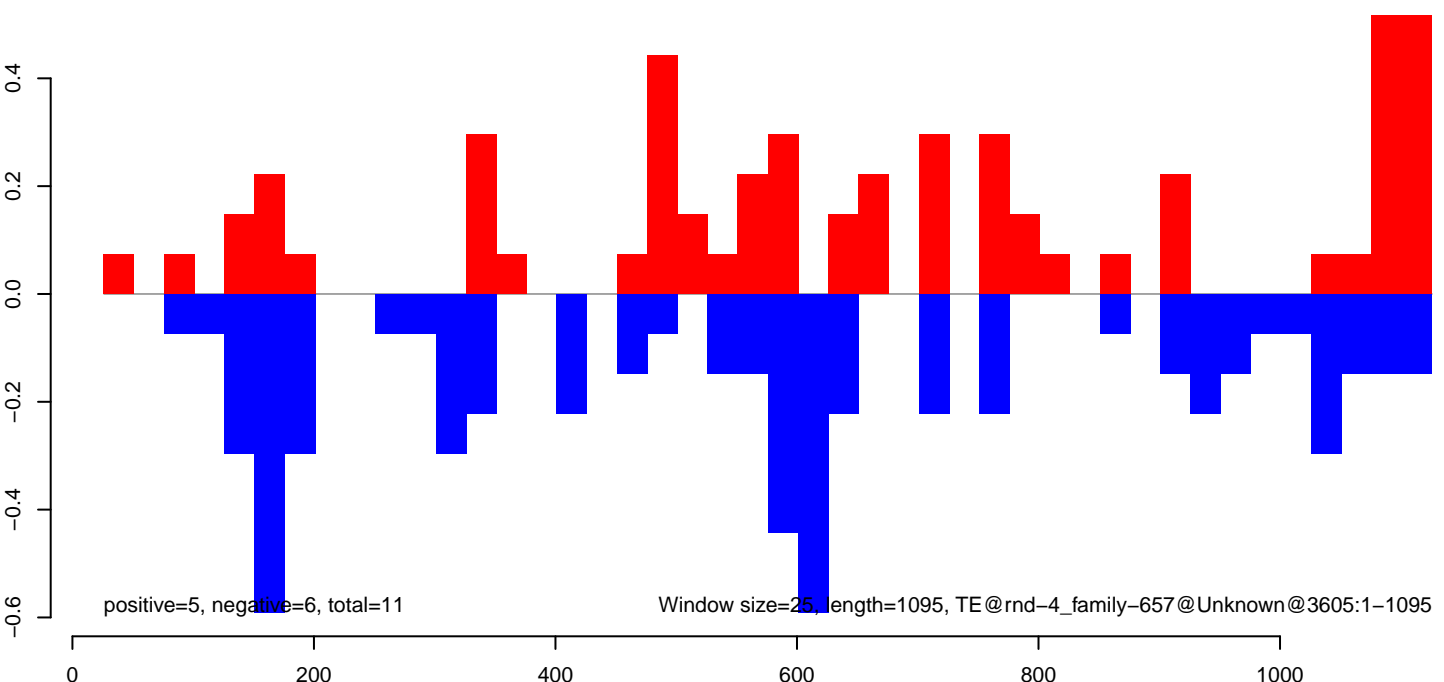
CuQuin_Testes_NL.18_23.rep



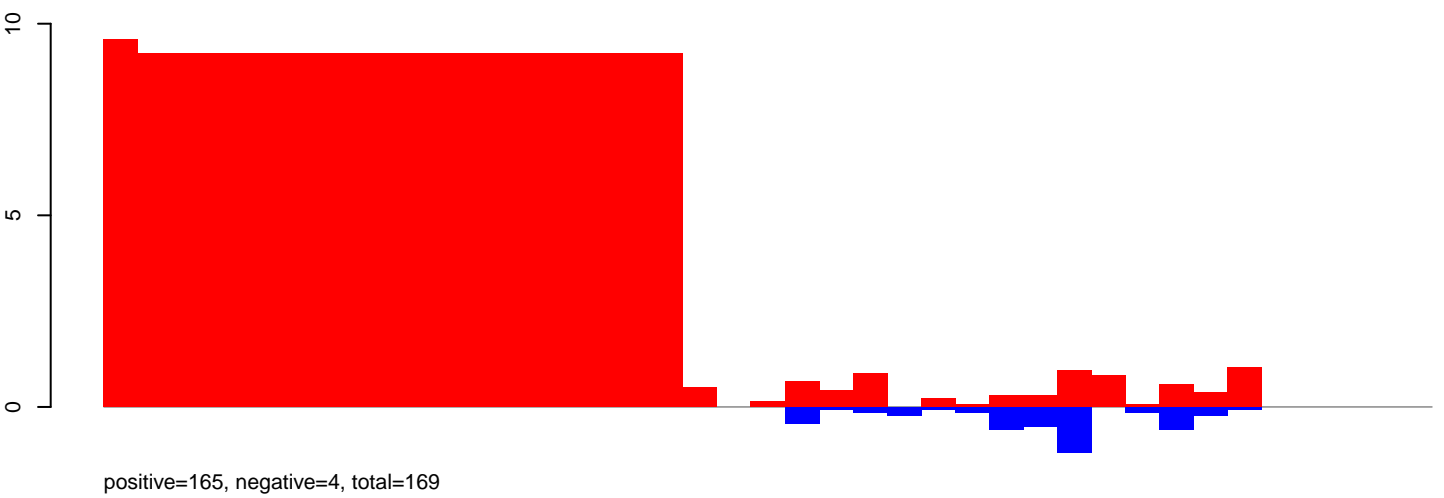
CuQuin_Testes_NL.24_35.rep



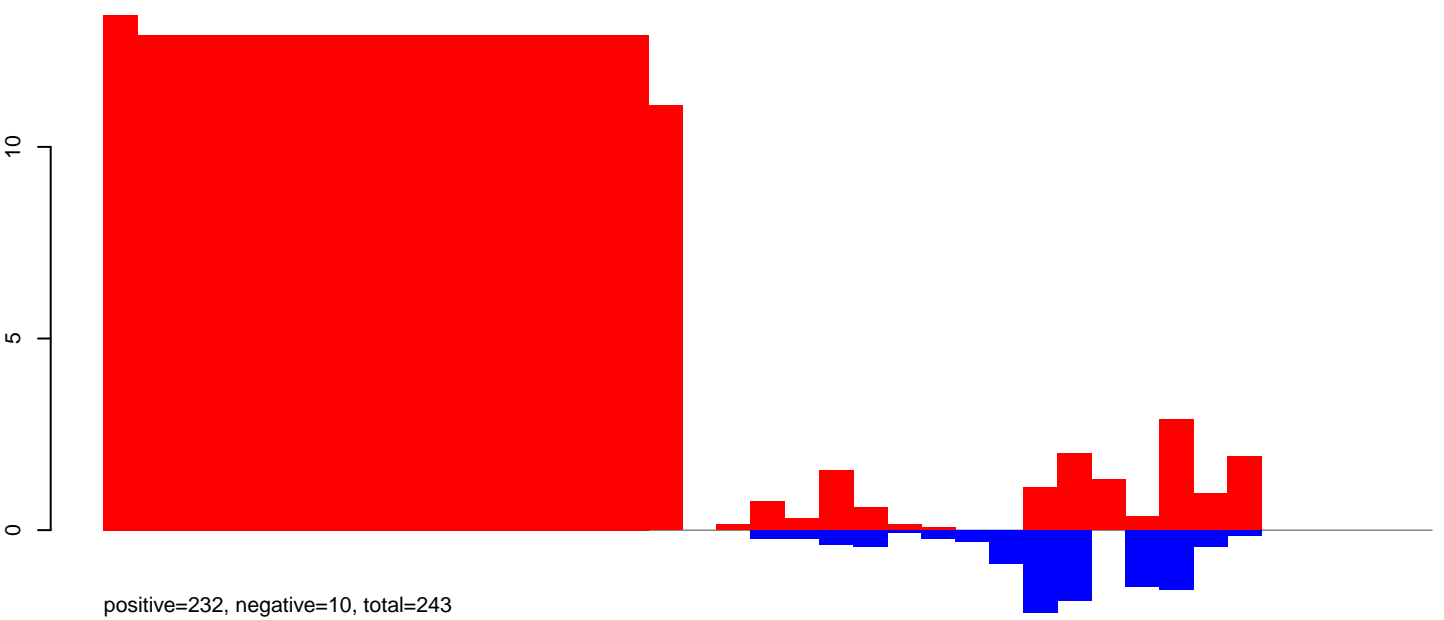
CuQuin_Testes_NL.rep



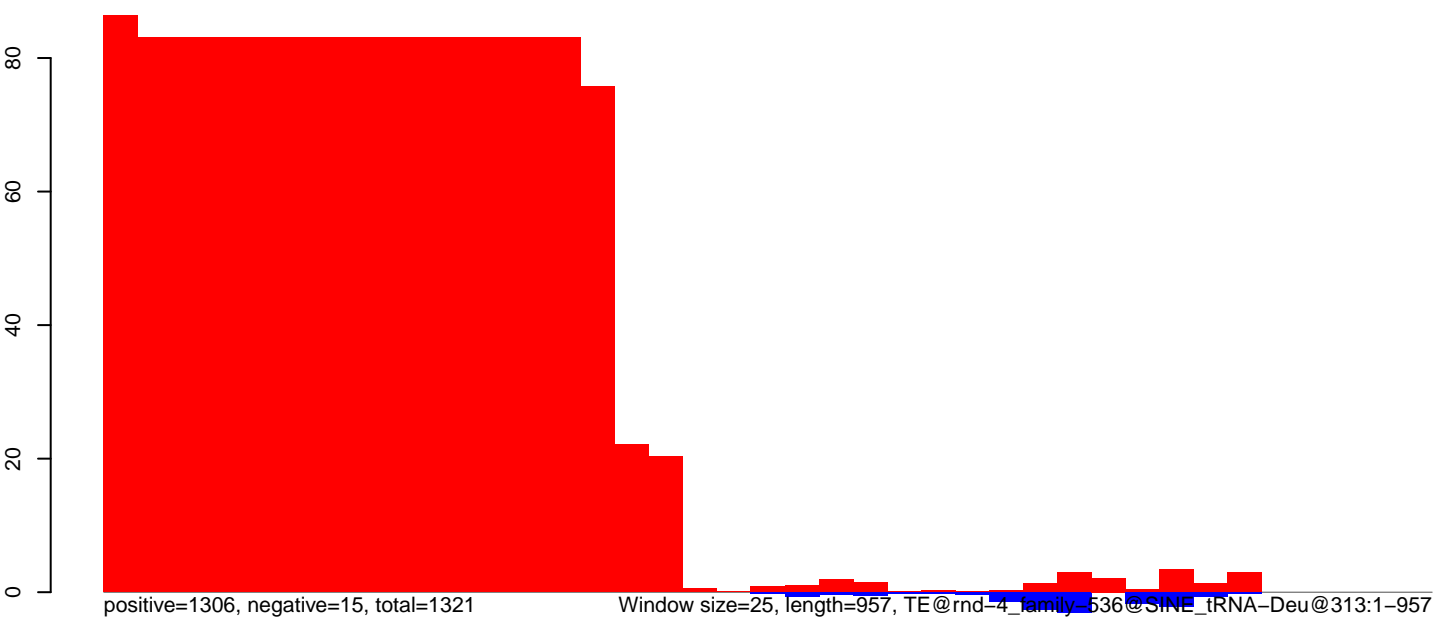
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

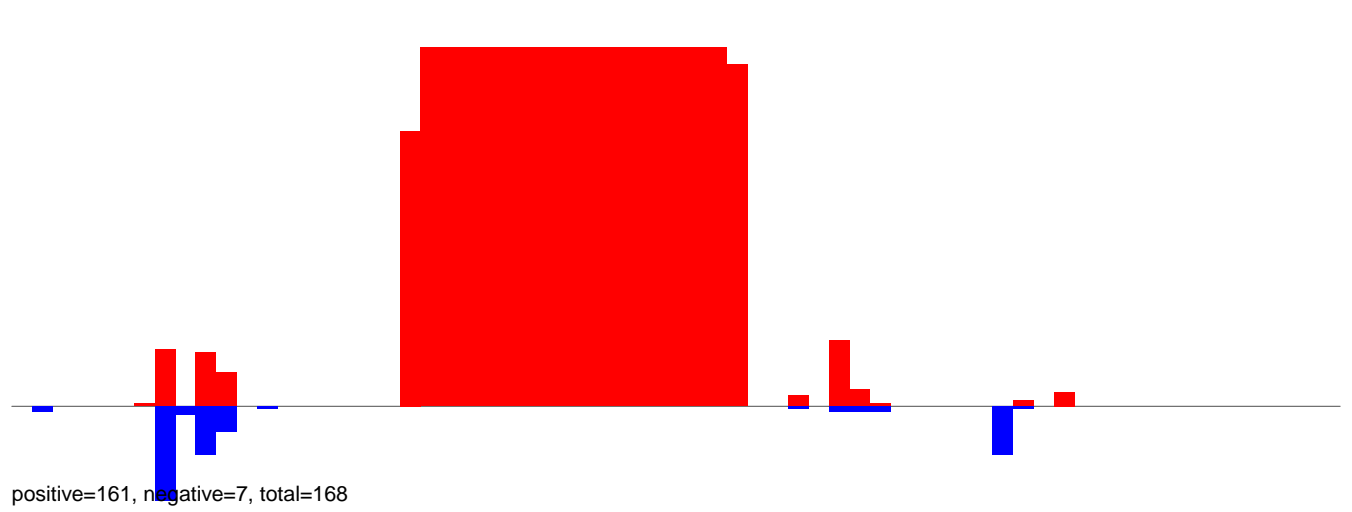


CuQuin_Testes_NL.rep



0 200 400 600 800 1000

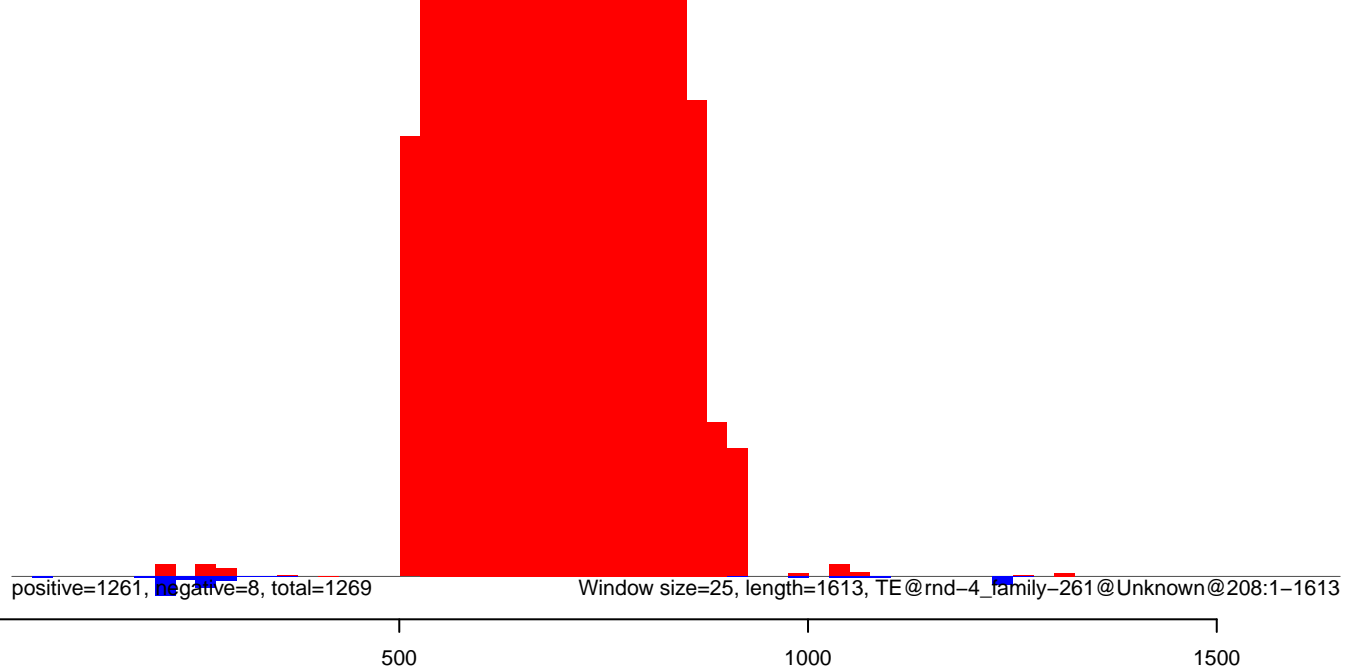
CuQuin_Testes_NL.18_23.rep



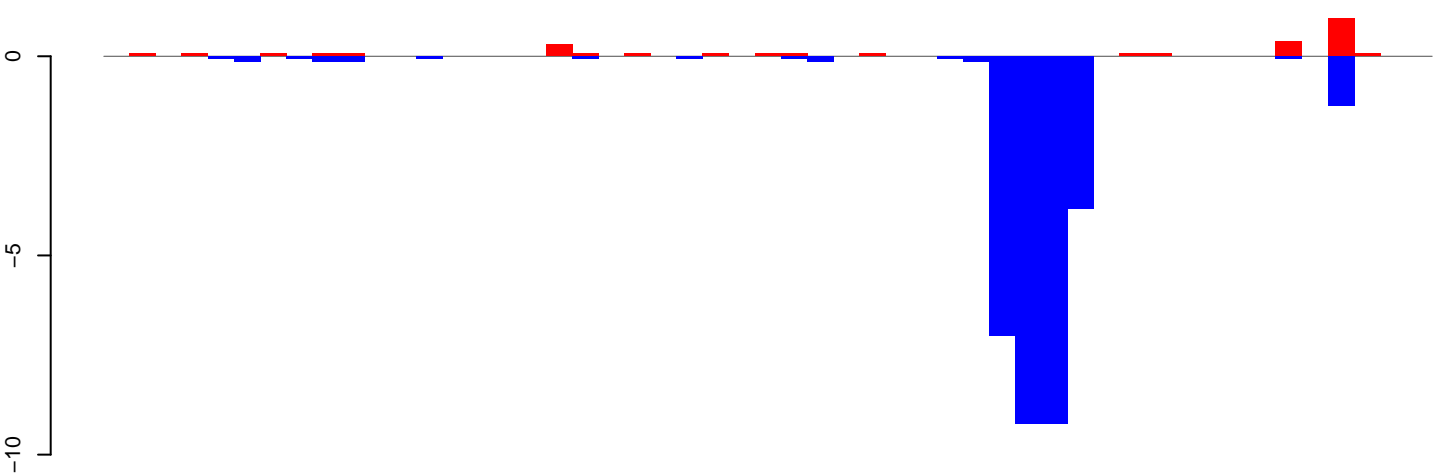
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

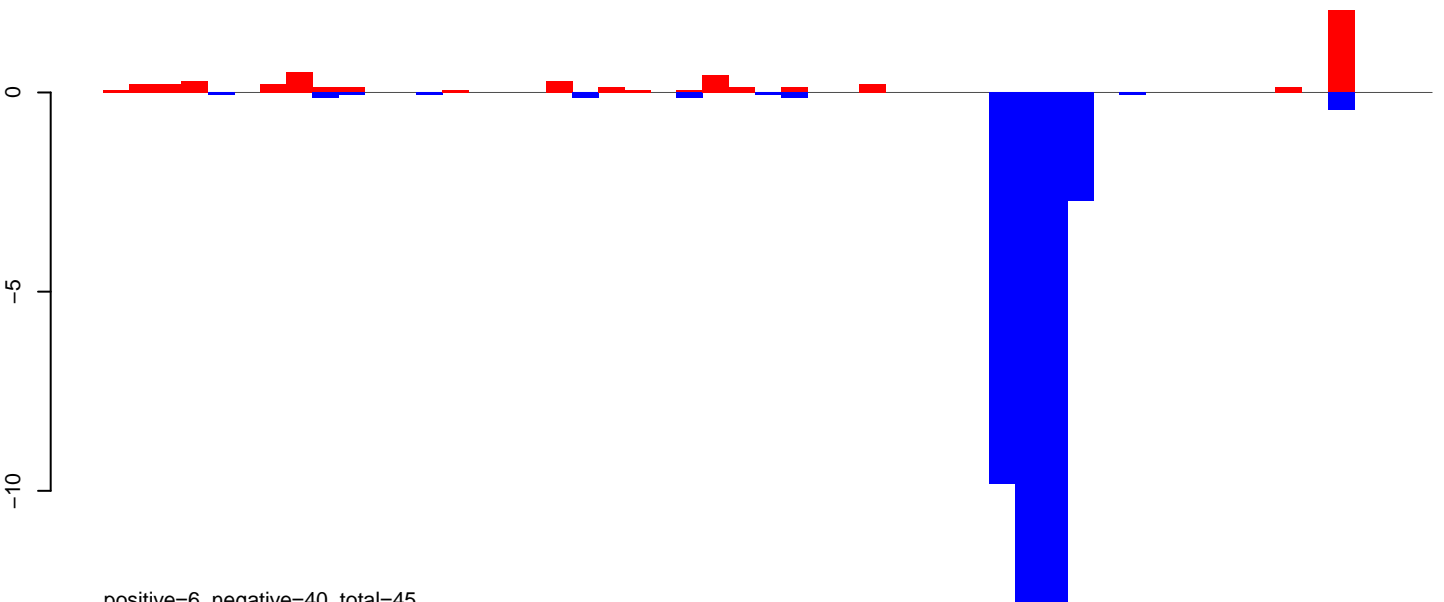


CuQuin_Testes_NL.18_23.rep



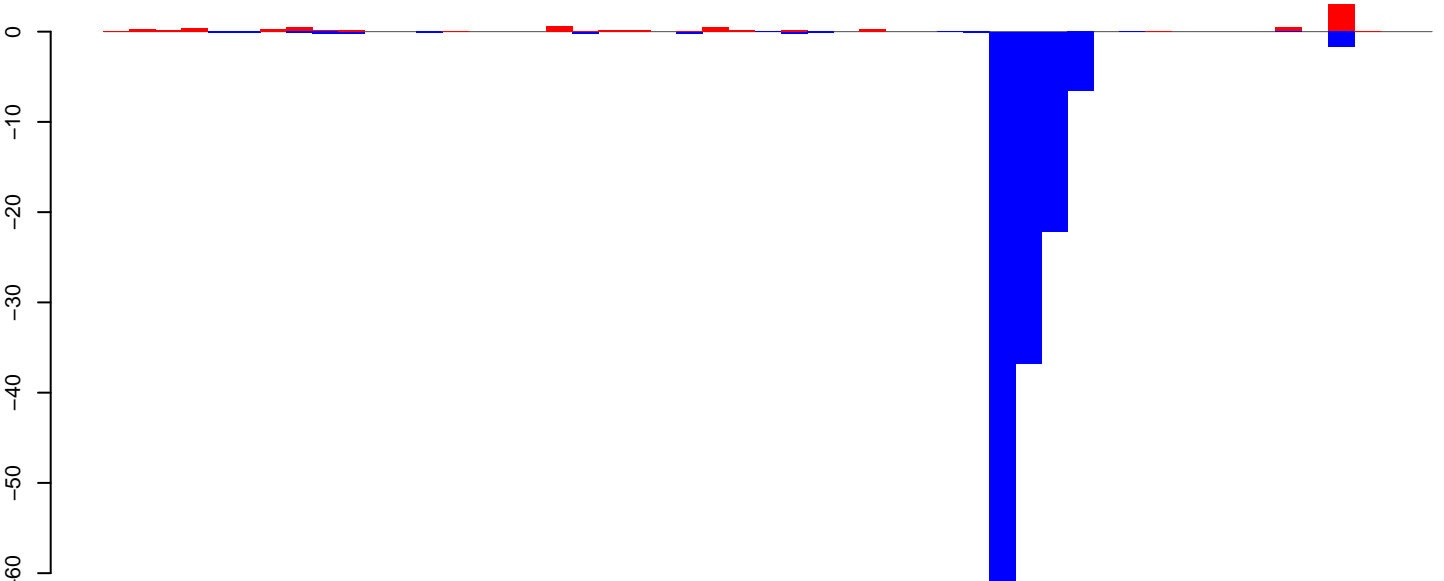
positive=3, negative=32, total=35

CuQuin_Testes_NL.24_35.rep



positive=6, negative=40, total=45

CuQuin_Testes_NL.rep

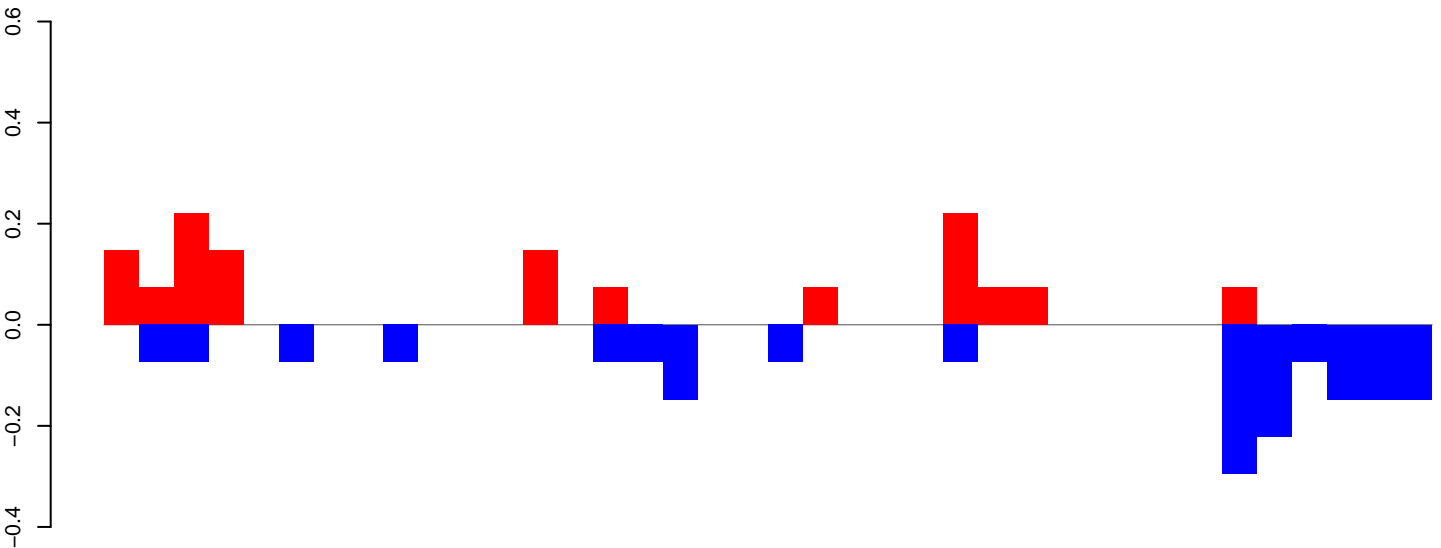


positive=8, negative=133, total=141

Window size=25, length=1269, TE@rnd-4 family-159@DNA_Crypton-I@1568:1-1269

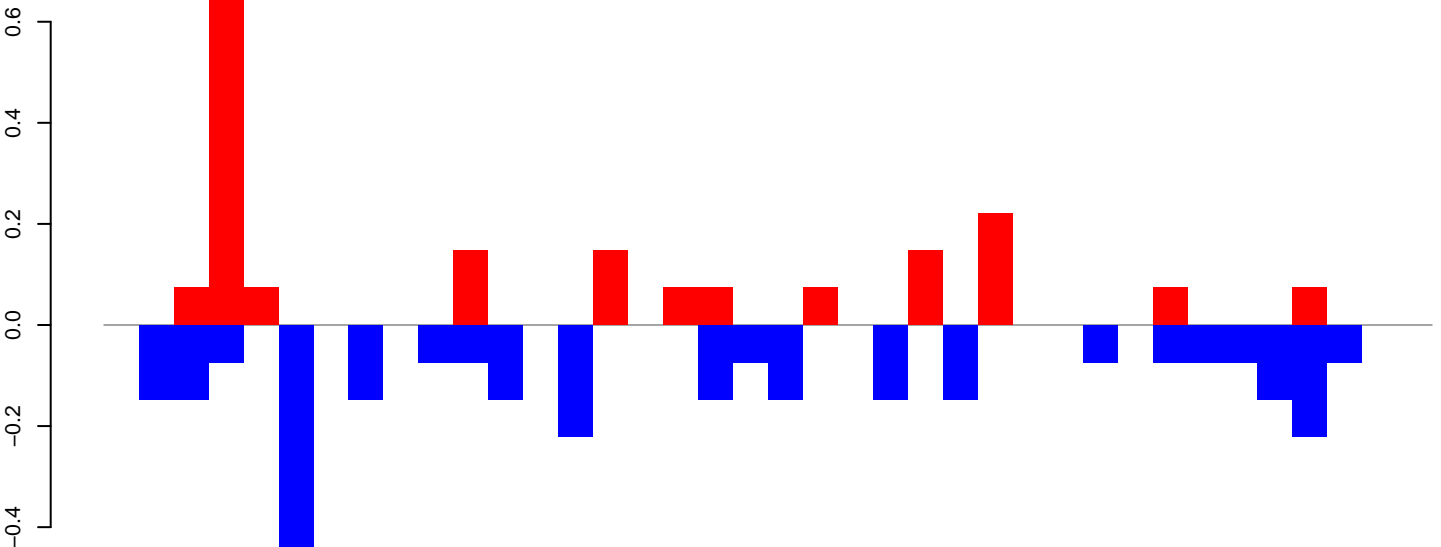
0 200 400 600 800 1000 1200

CuQuin_Testes_NL.18_23.rep



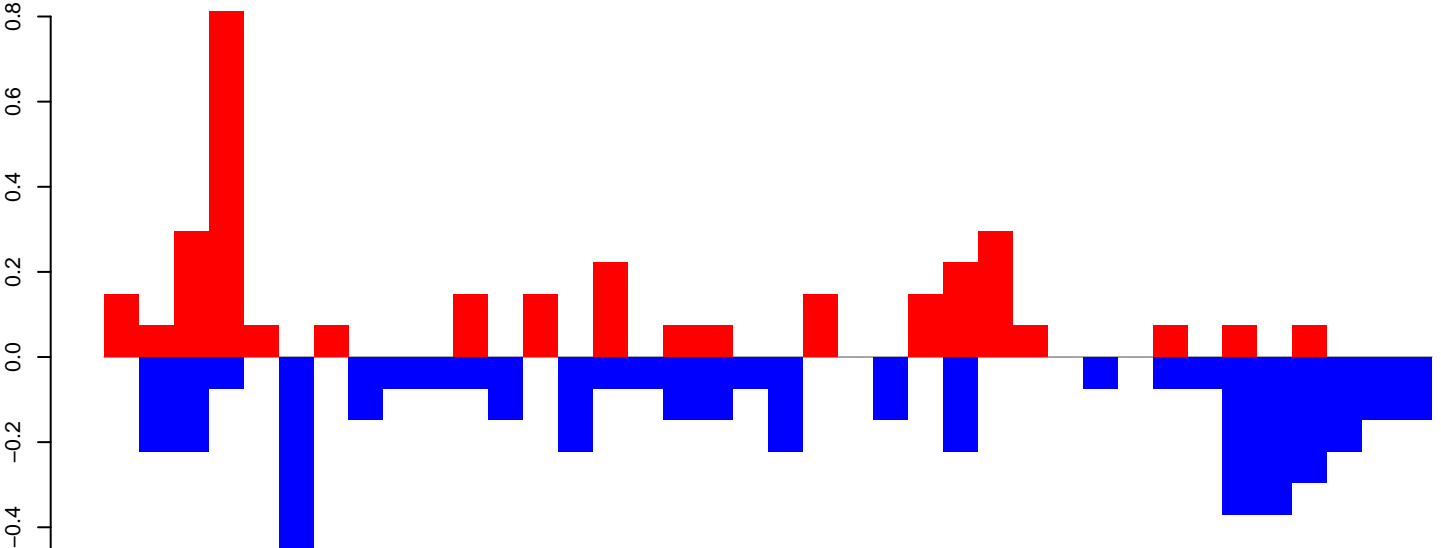
positive=1, negative=2, total=3

CuQuin_Testes_NL.24_35.rep



positive=2, negative=3, total=5

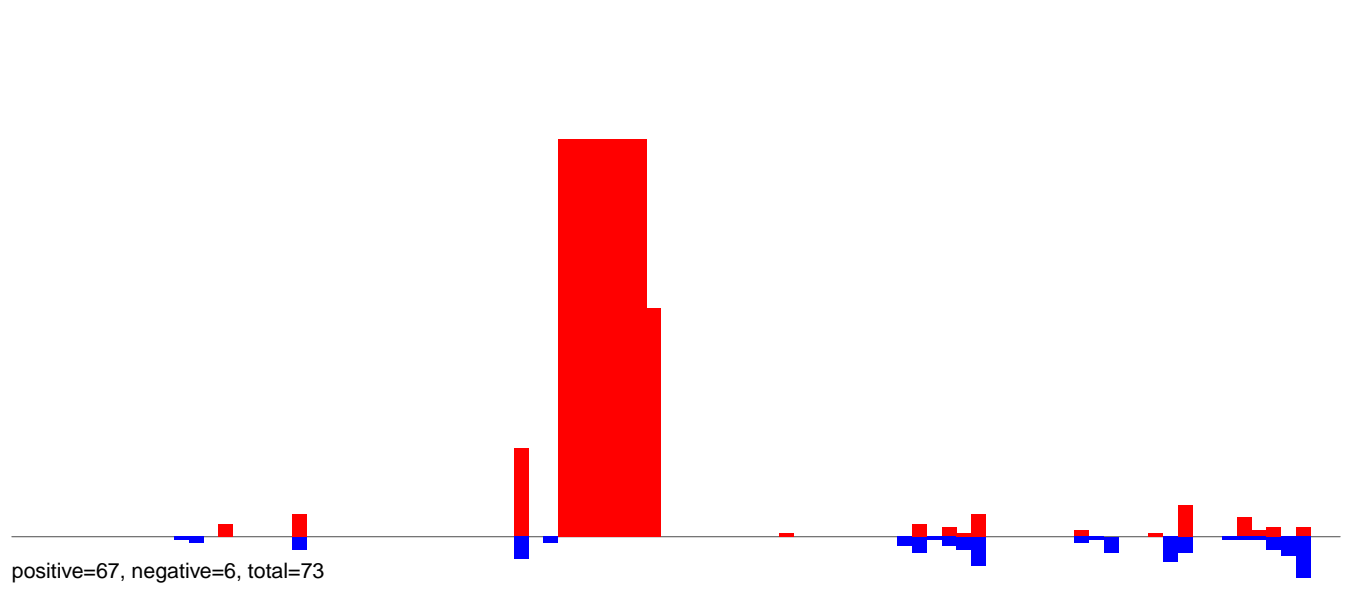
CuQuin_Testes_NL.rep



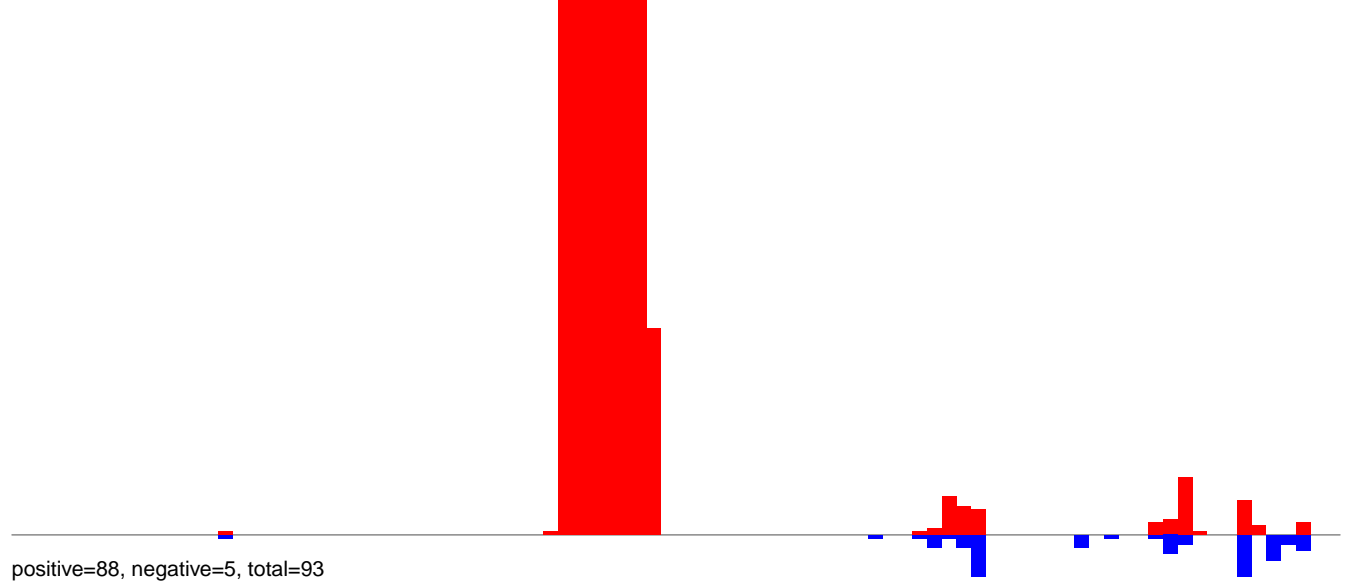
positive=3, negative=5, total=8

Window size=25, length=944, TE@rnd-3_family-794@Unknown@54:1-944

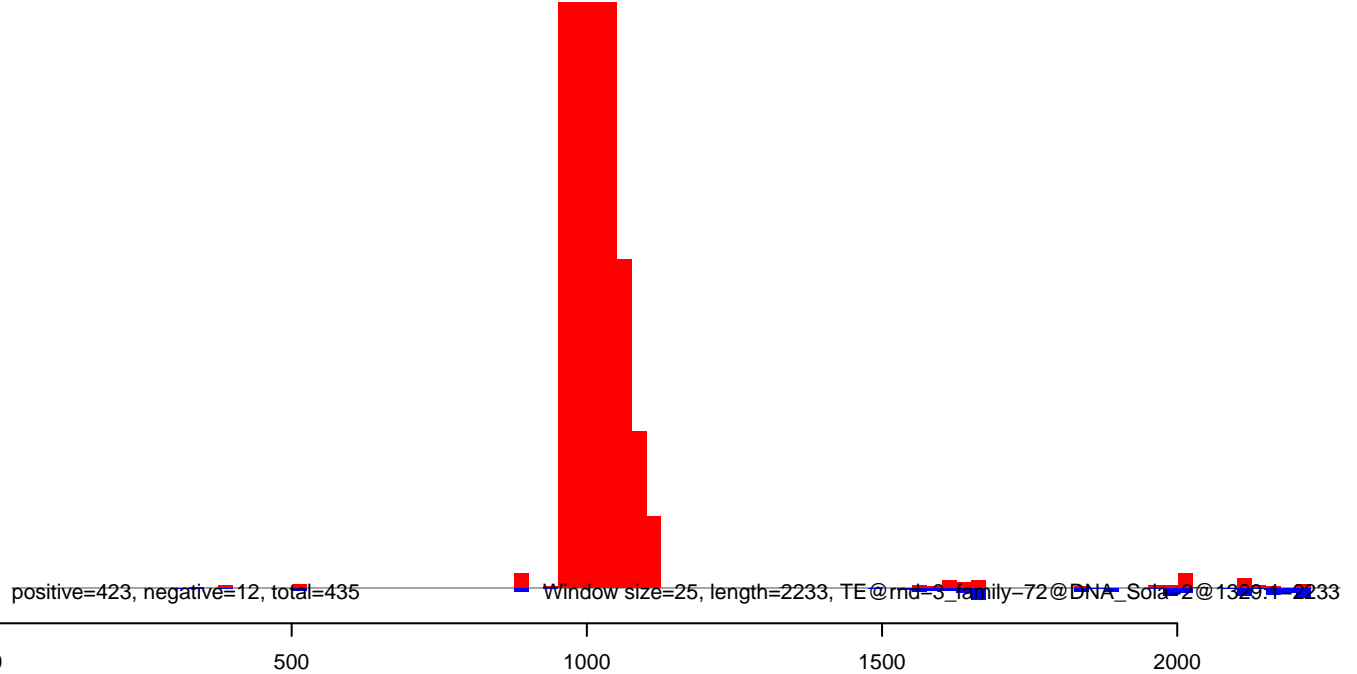
CuQuin_Testes_NL.18_23.rep



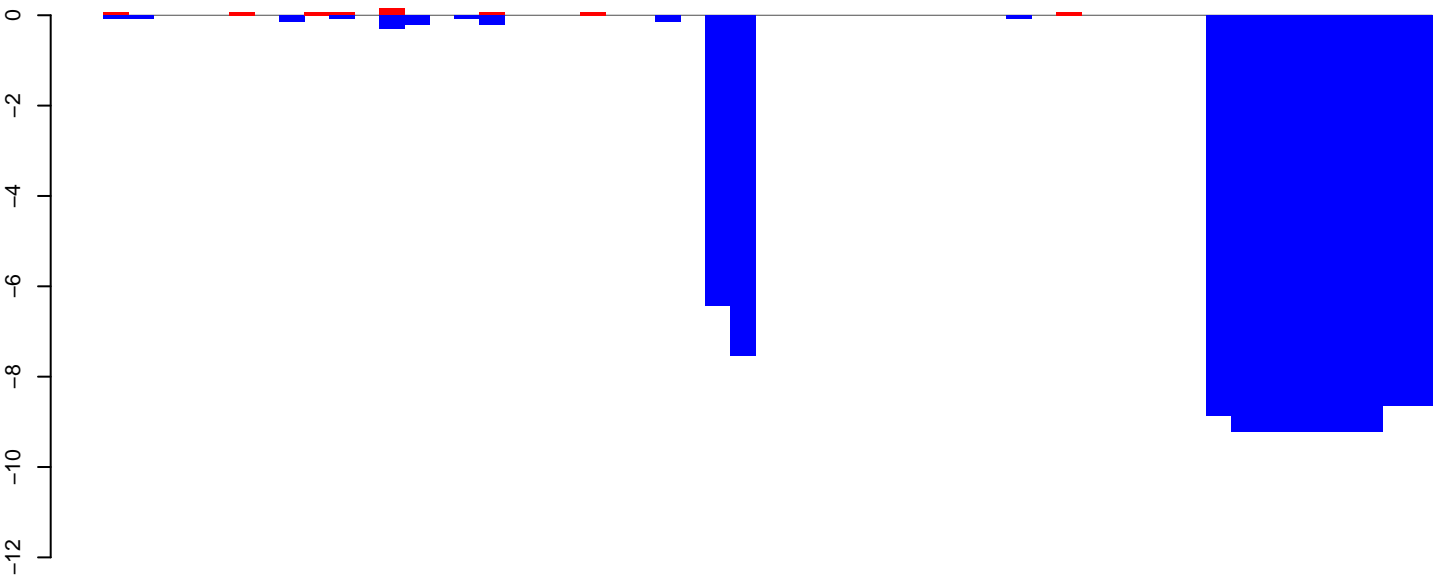
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

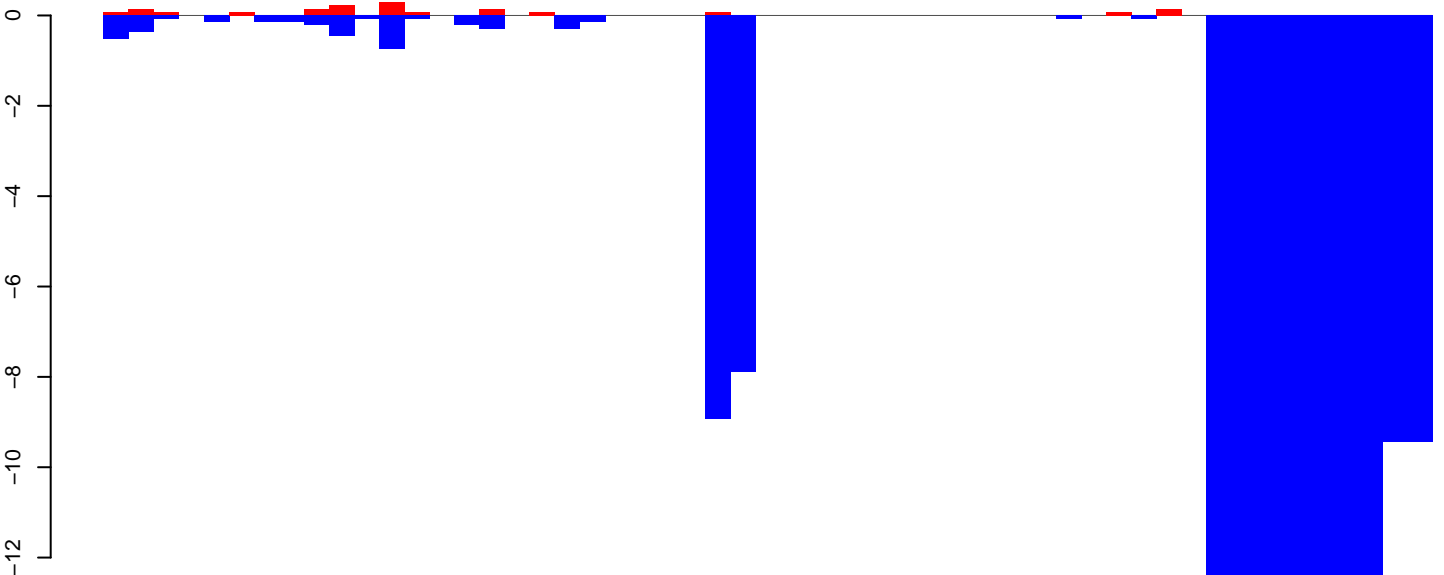


CuQuin_Testes_NL.18_23.rep



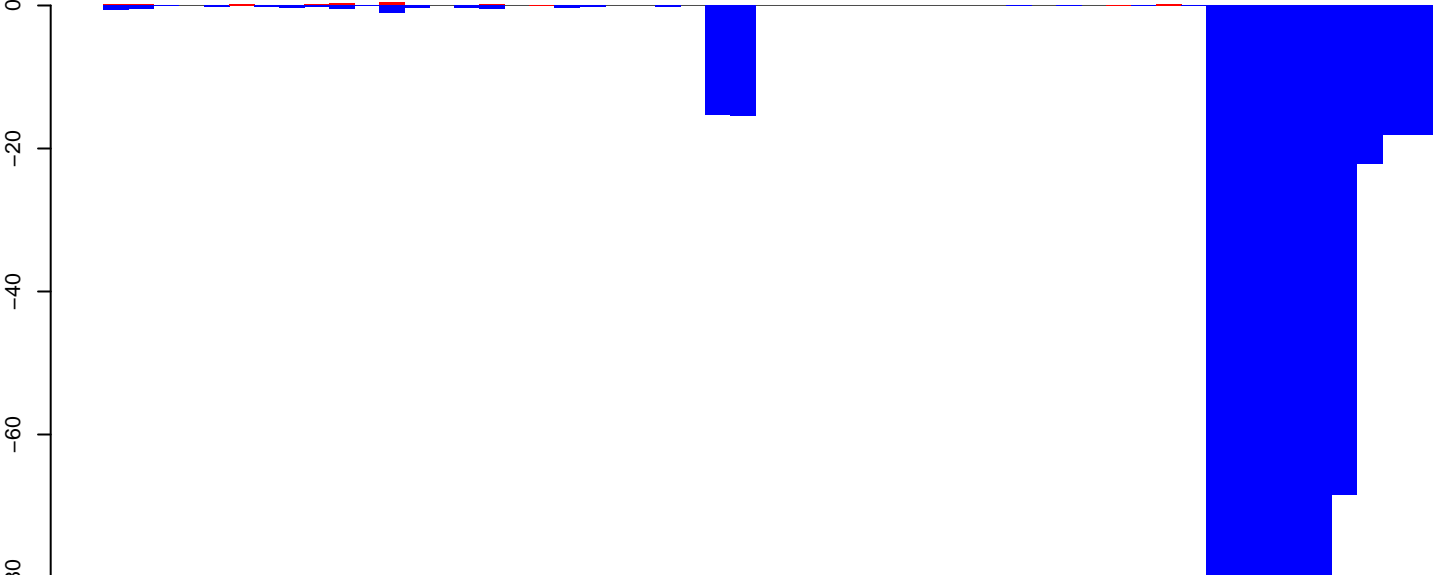
positive=1, negative=97, total=98

CuQuin_Testes_NL.24_35.rep



positive=2, negative=130, total=131

CuQuin_Testes_NL.rep

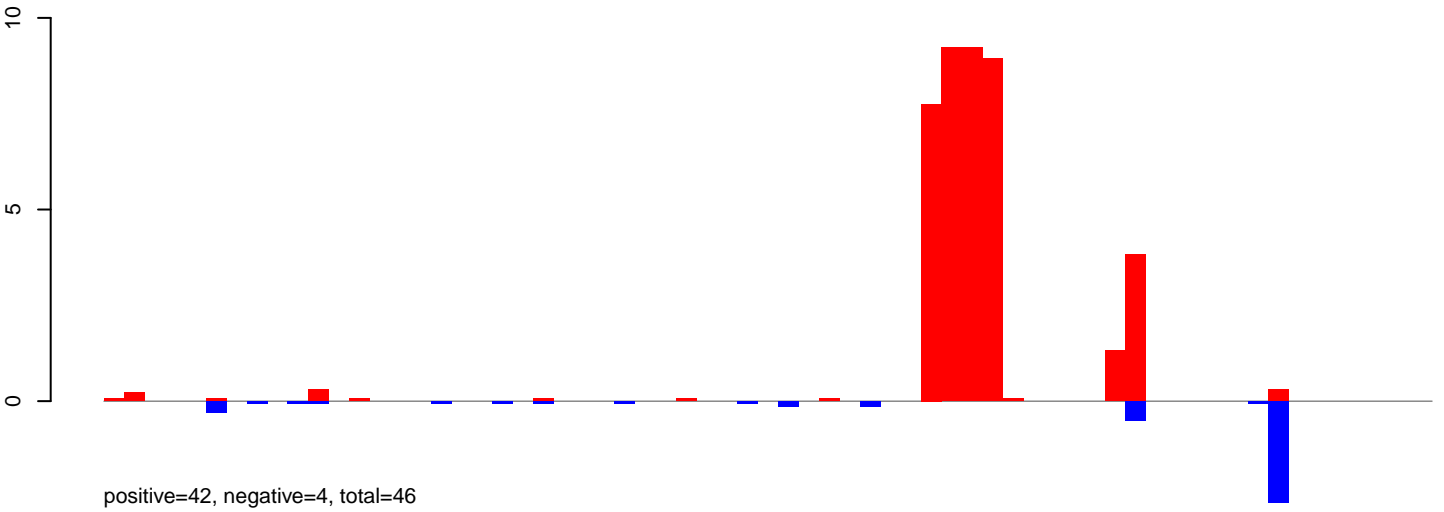


positive=2, negative=575, total=577

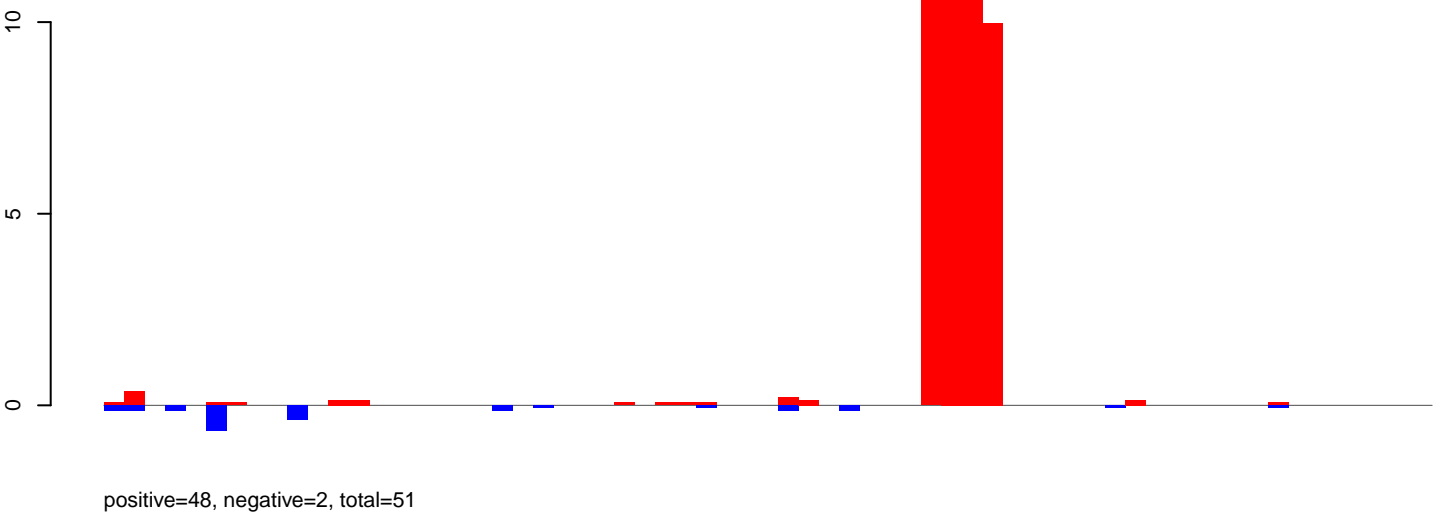
Window size=25, length=1320, TE@rnd-3_family-104@LTR_Gypsy@1335:1-1320

0 200 400 600 800 1000 1200 1400

CuQuin_Testes_NL.18_23.rep



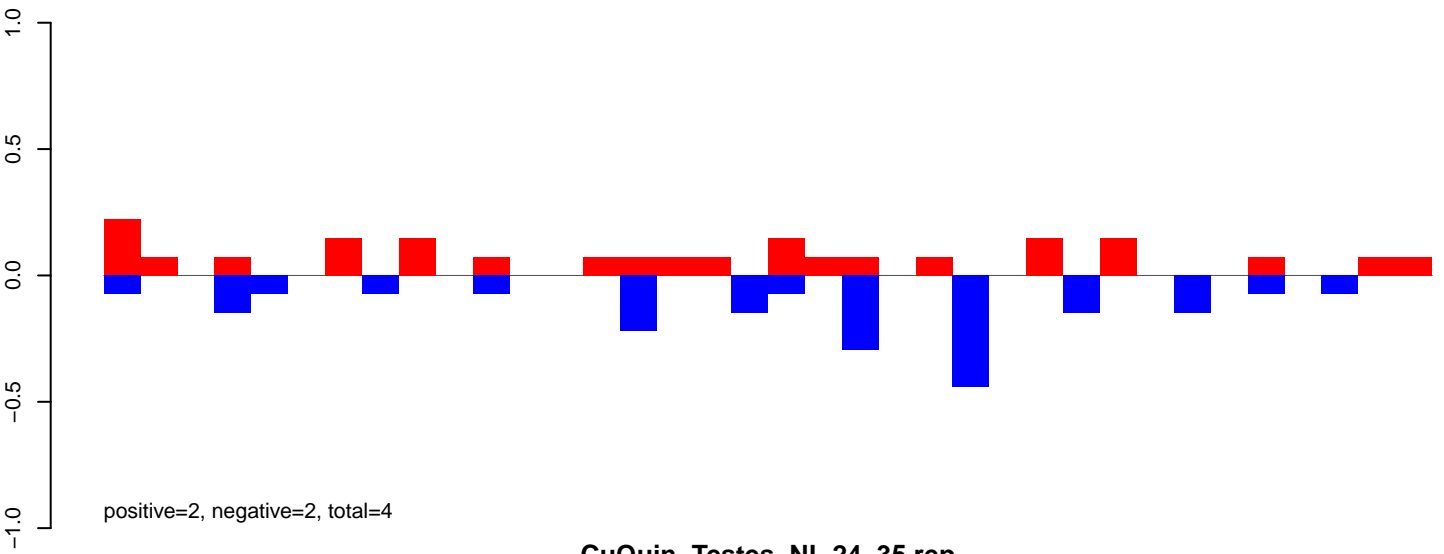
CuQuin_Testes_NL.24_35.rep



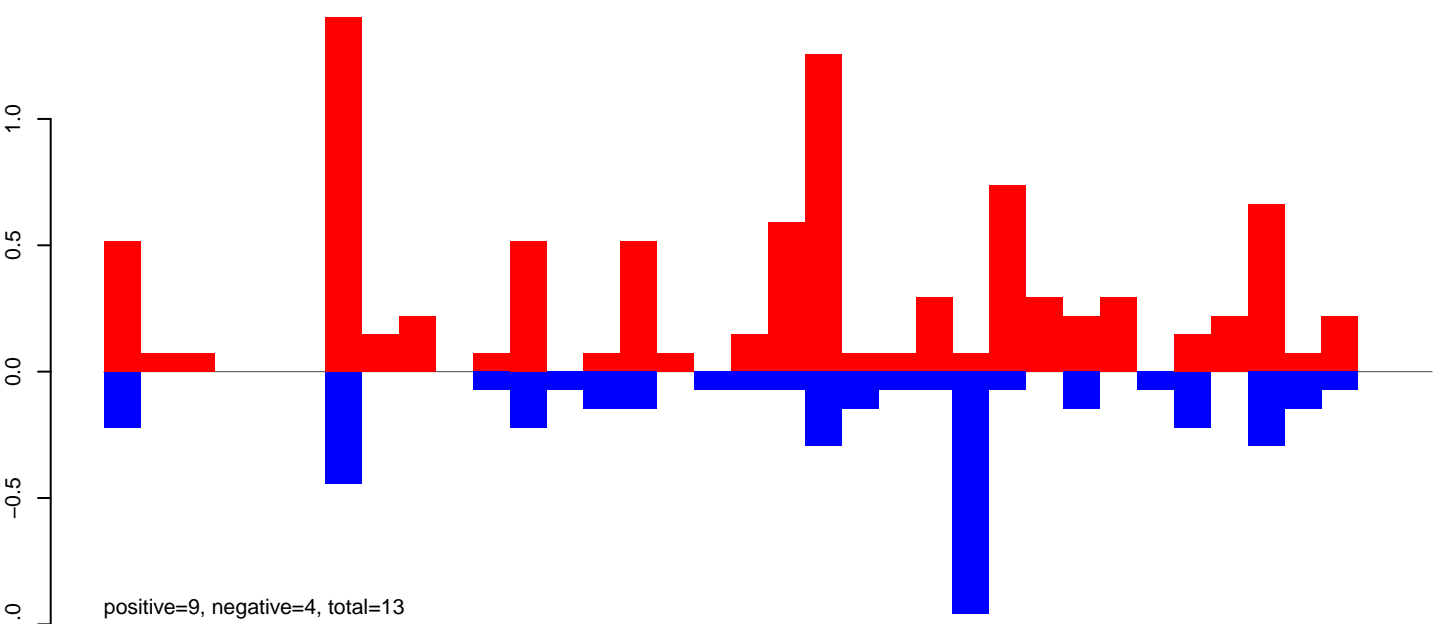
CuQuin_Testes_NL.rep



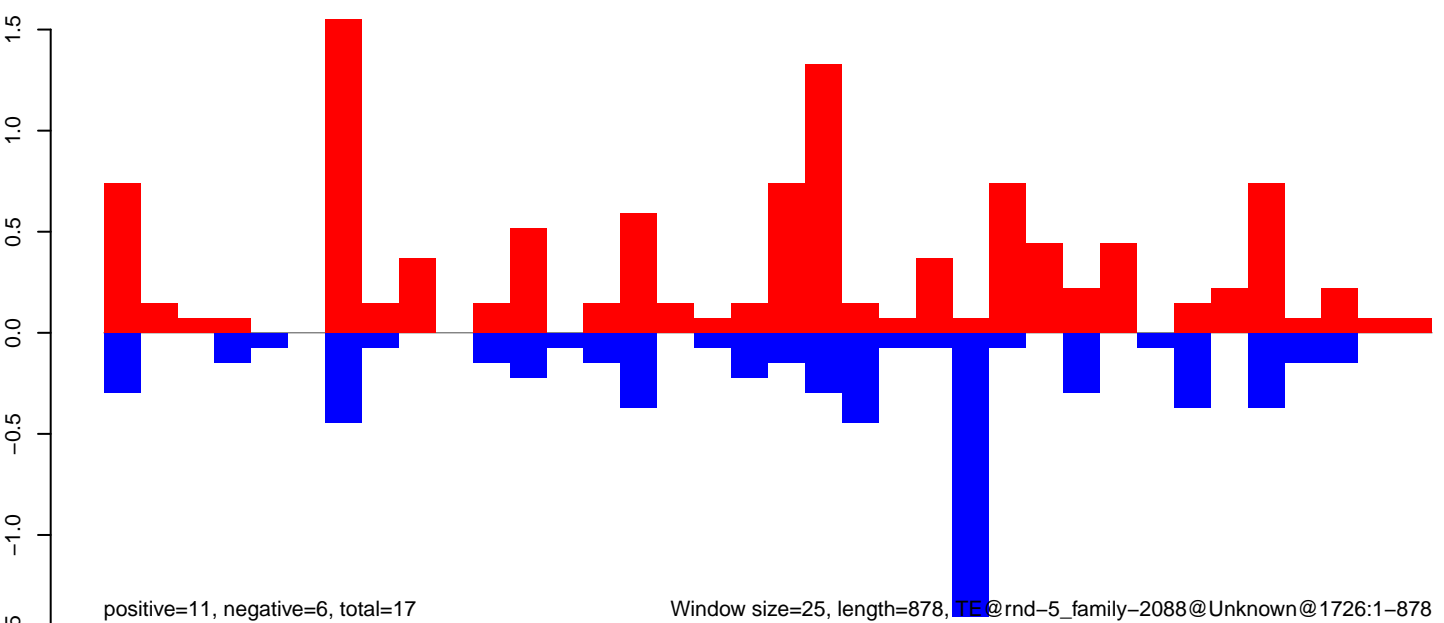
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

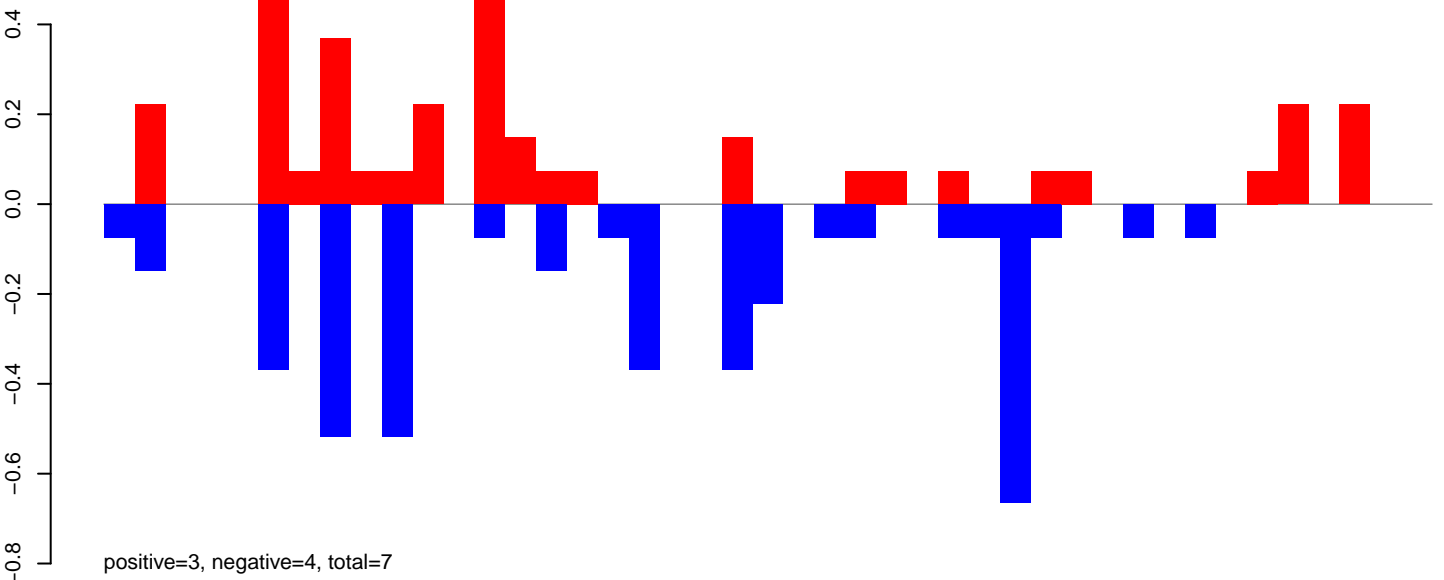


CuQuin_Testes_NL.rep

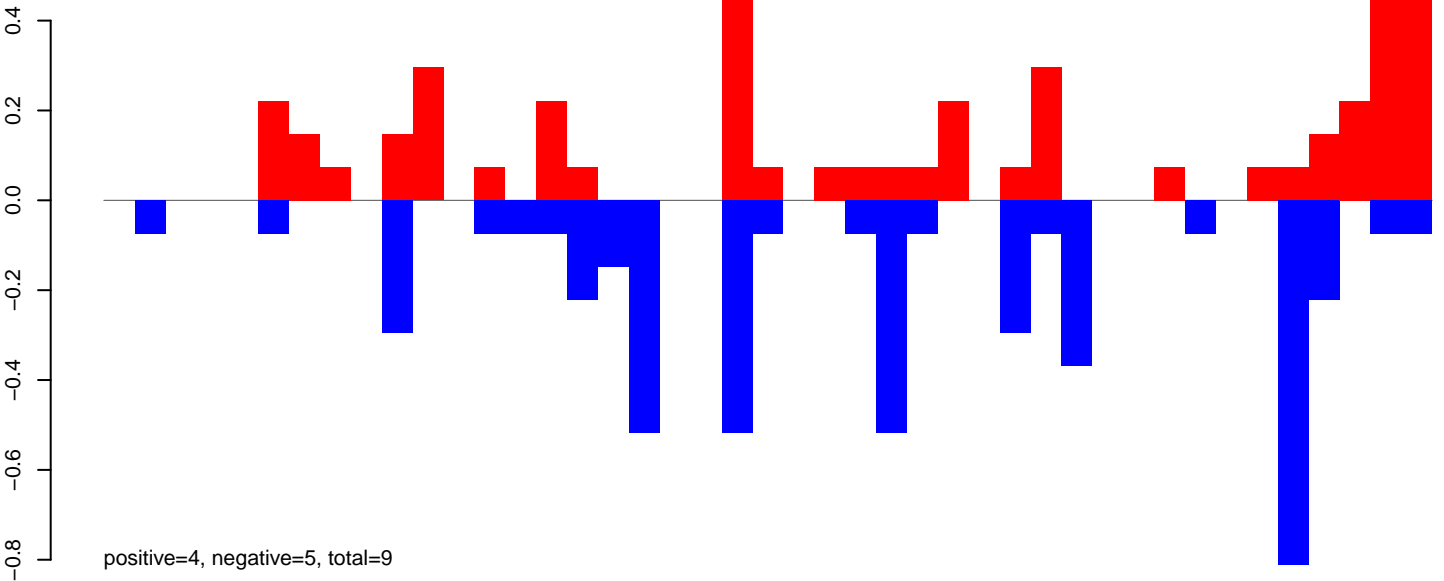


Window size=25, length=878, TE@rnd-5_family-2088@Unknown@1726:1-878

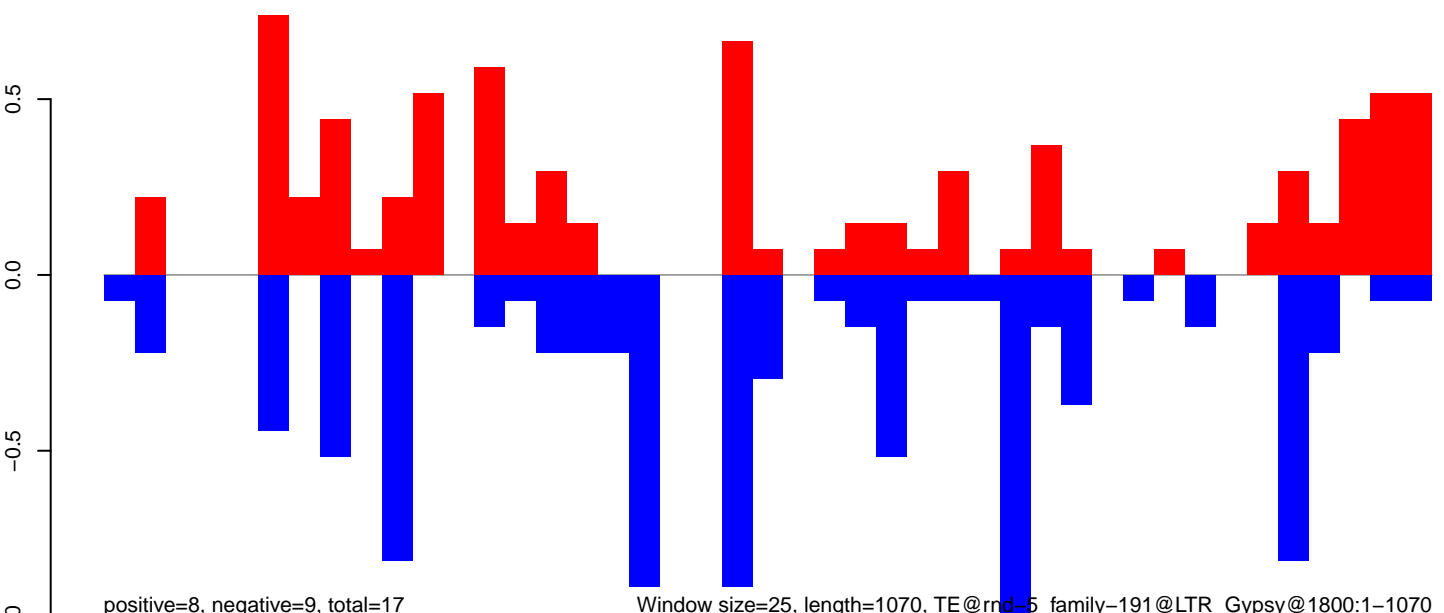
CuQuin_Testes_NL.18_23.rep



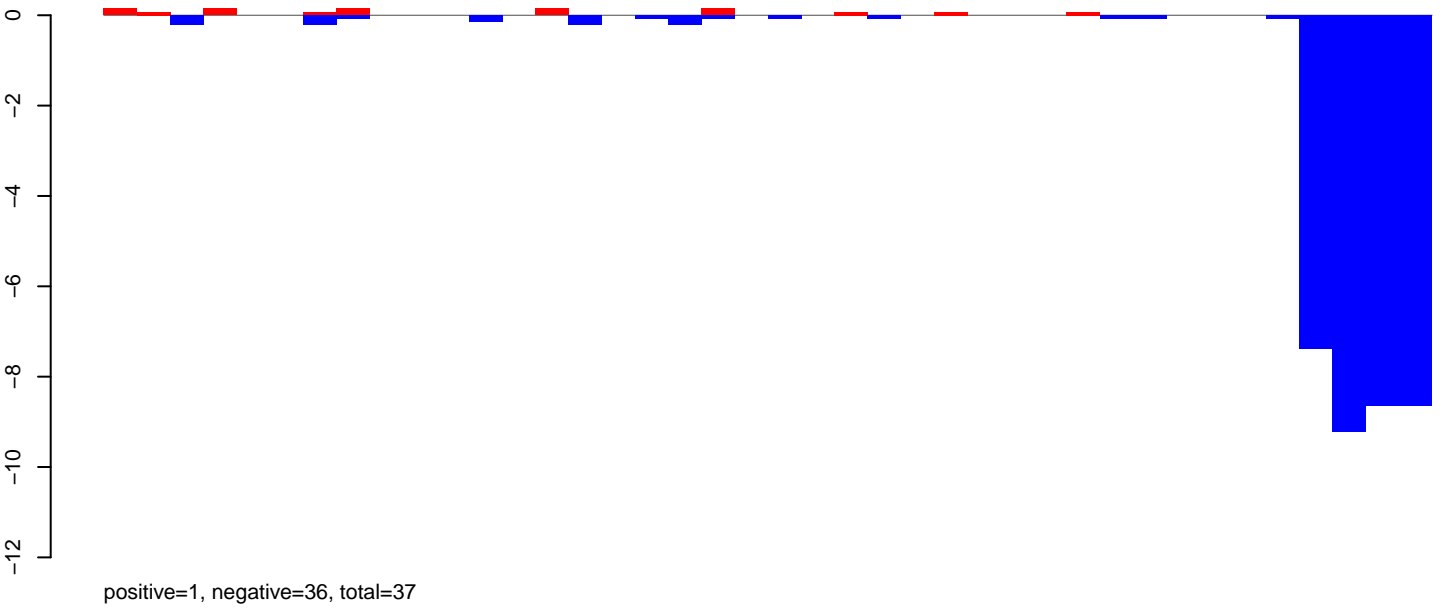
CuQuin_Testes_NL.24_35.rep



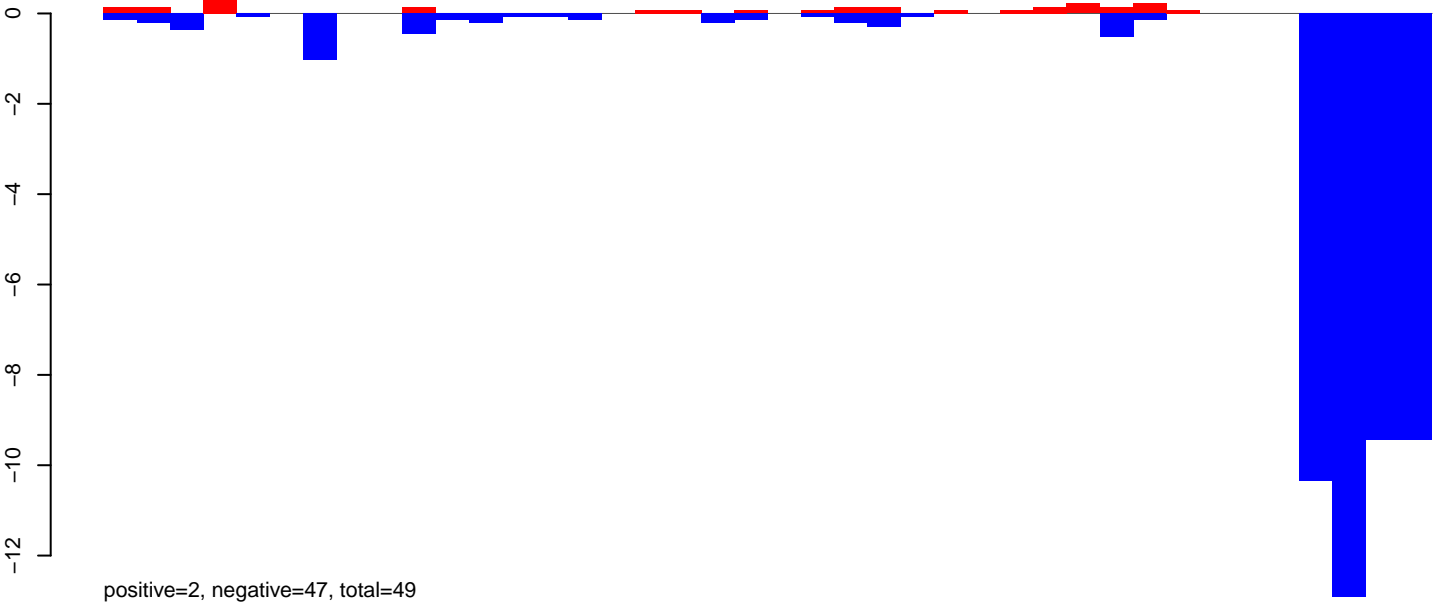
CuQuin_Testes_NL.rep



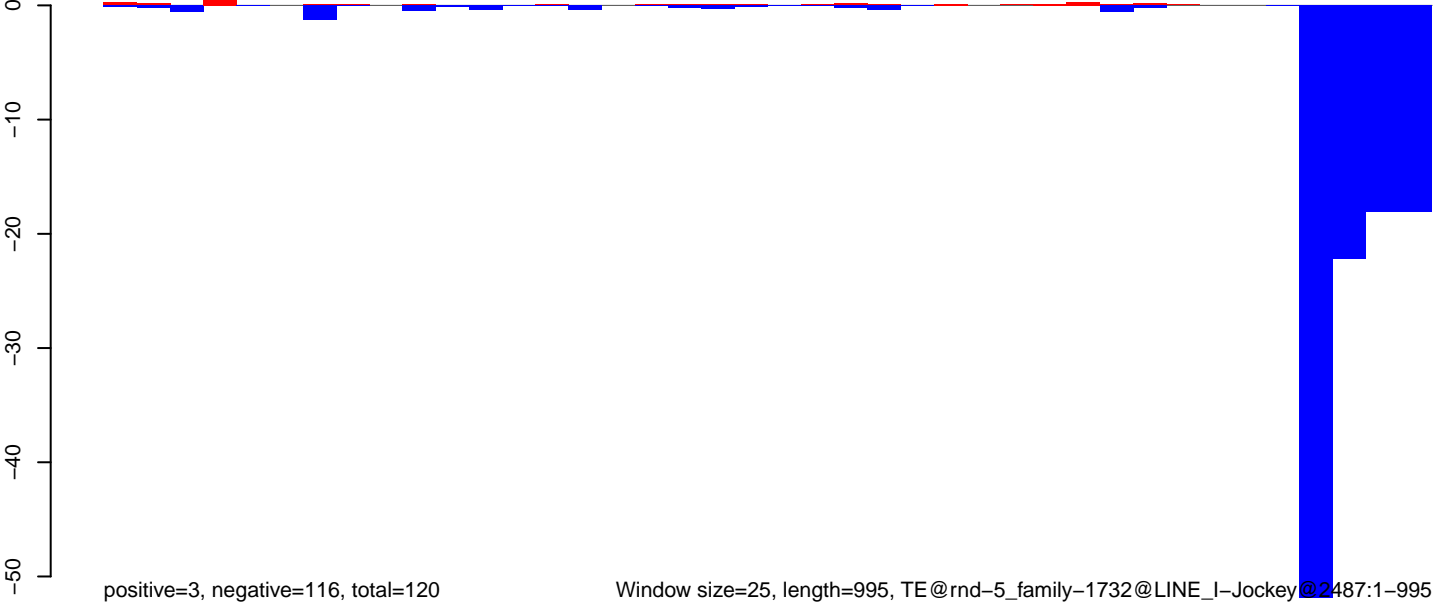
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

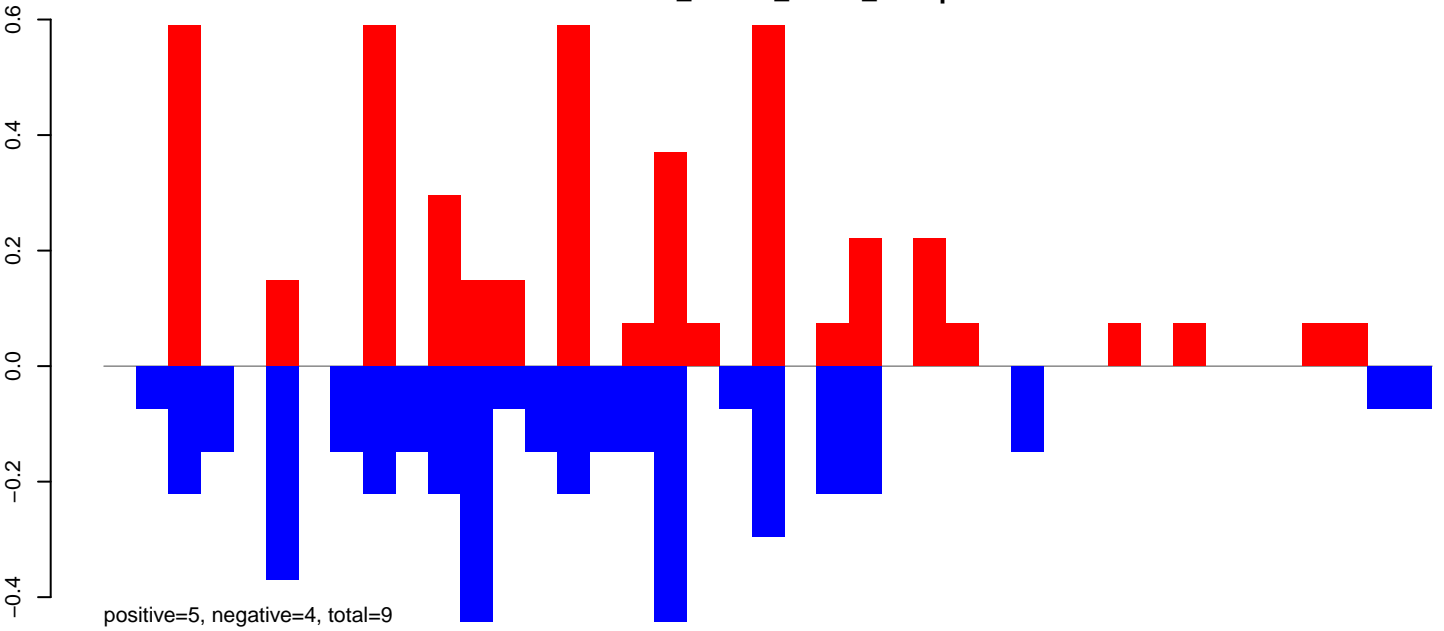


CuQuin_Testes_NL.rep

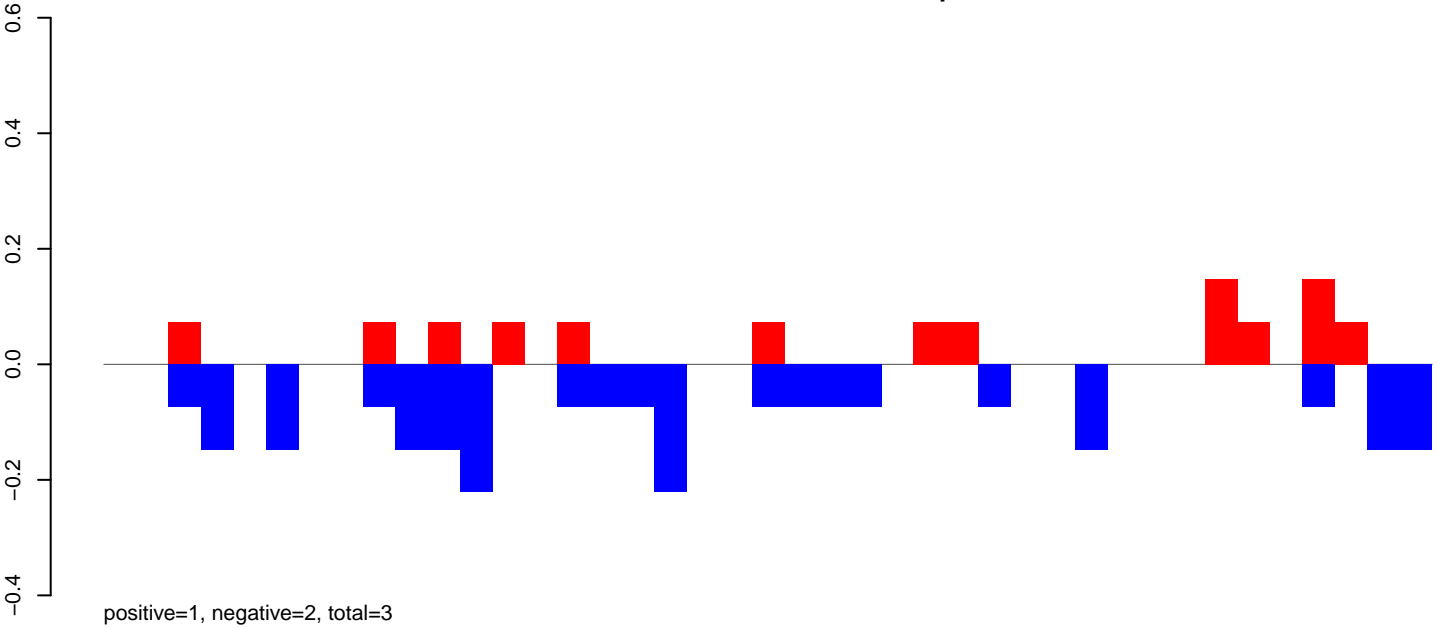


Window size=25, length=995, TE@rnd-5_family-1732@LINE_I-Jockey@2487:1-995

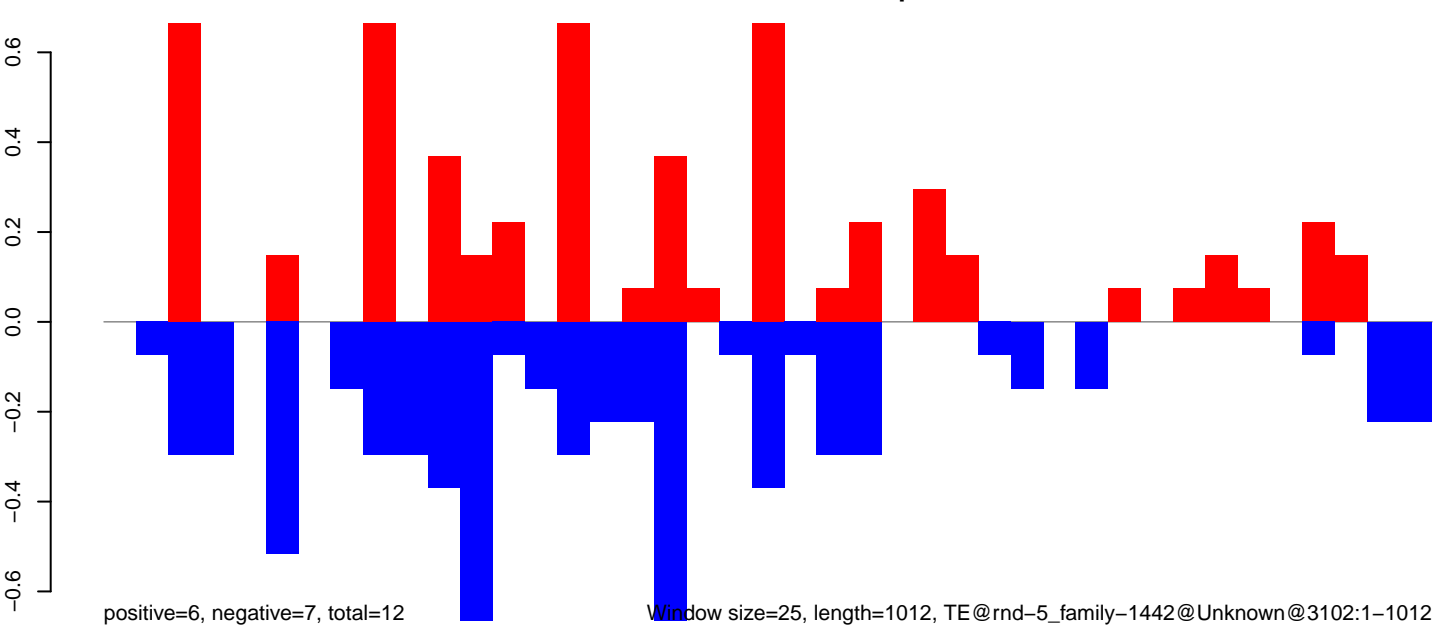
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



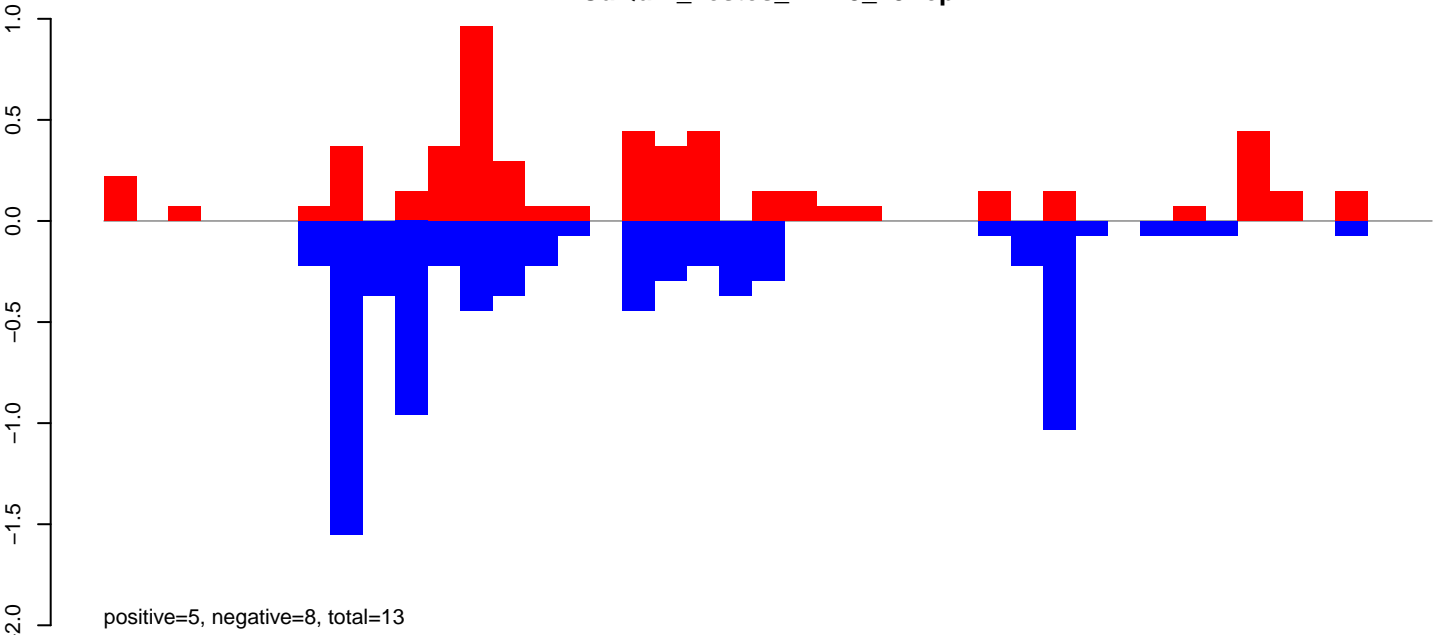
CuQuin_Testes_NL.rep



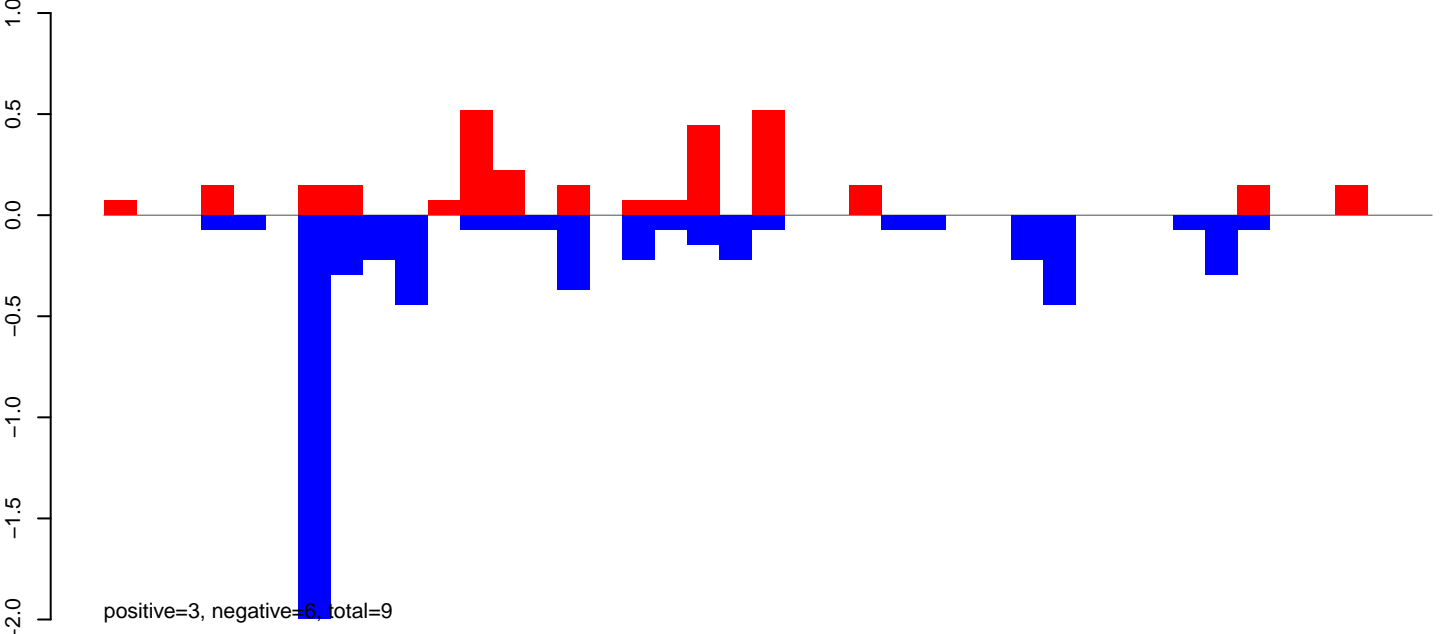
Window size=25, length=1012, TE@rnd-5_family-1442@Unknown@3102:1-1012

0 200 400 600 800 1000

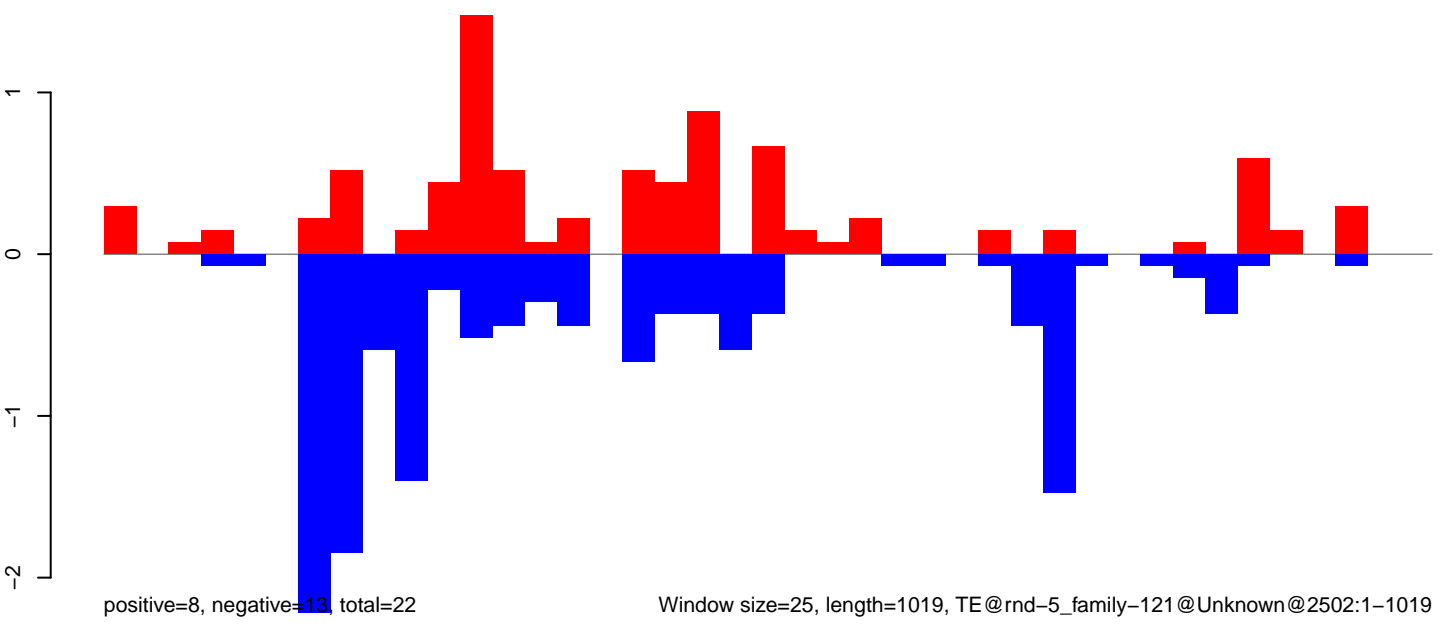
CuQuin_Testes_NL.18_23.rep



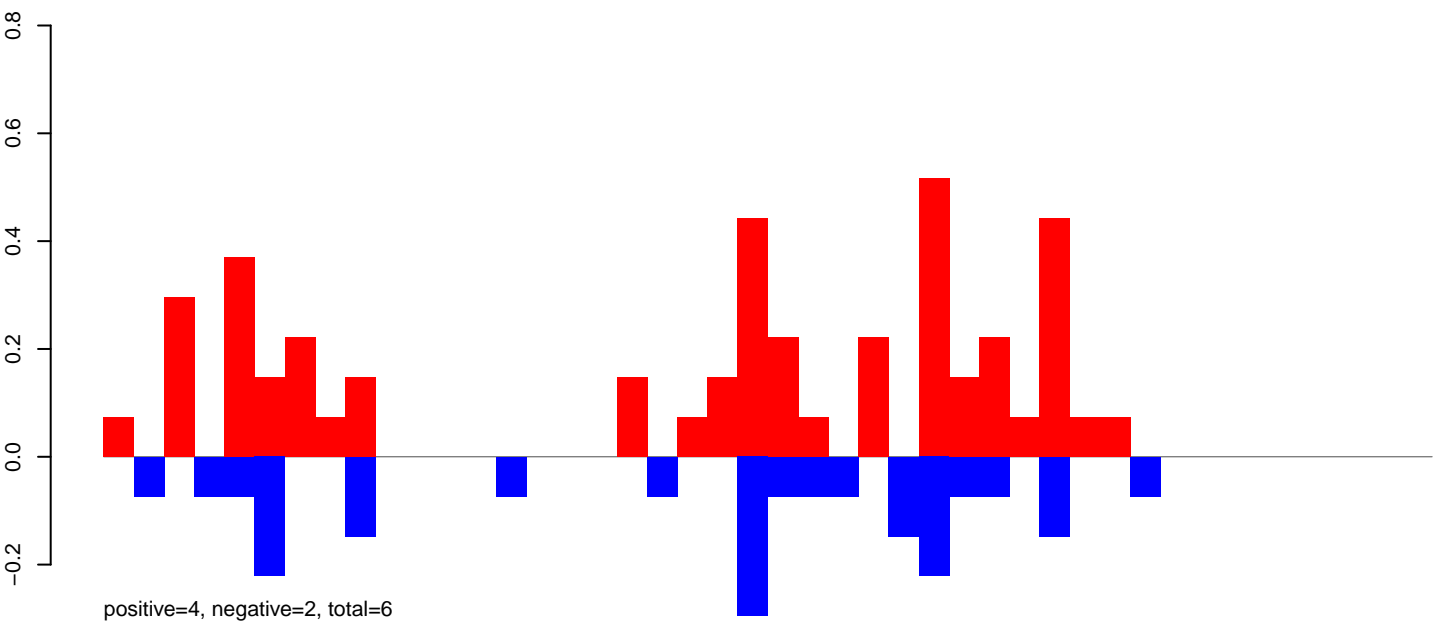
CuQuin_Testes_NL.24_35.rep



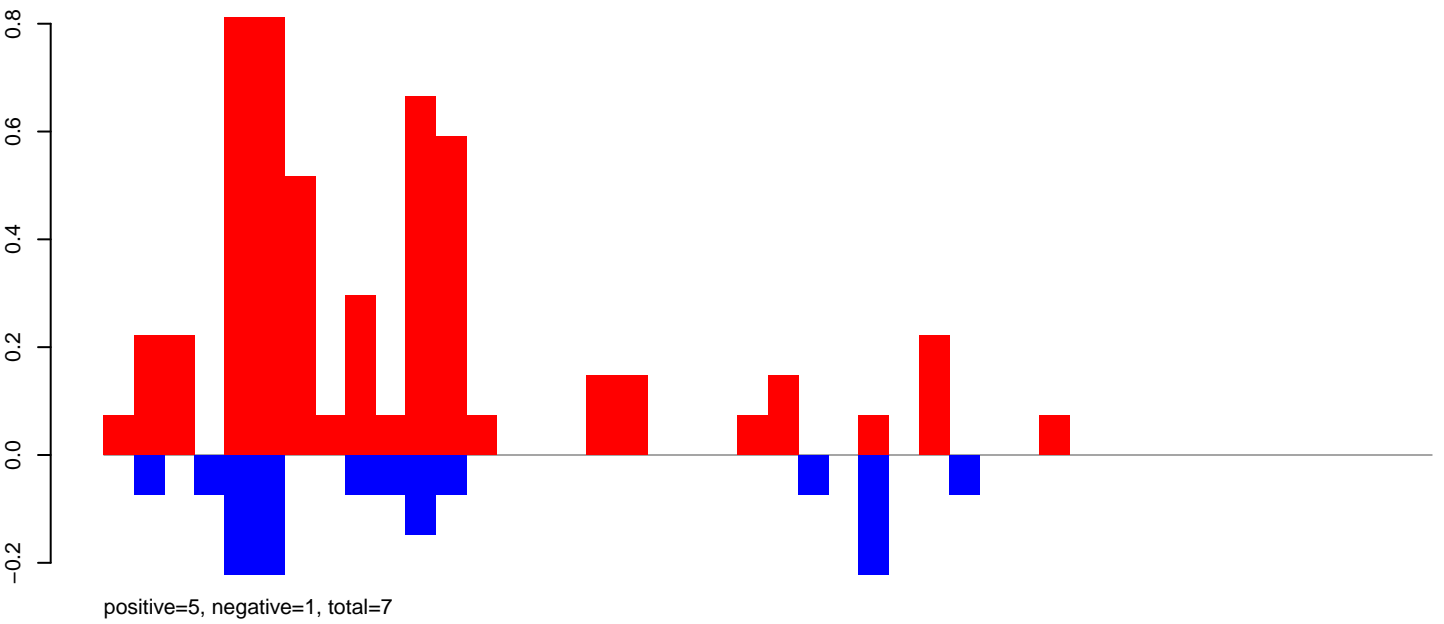
CuQuin_Testes_NL.rep



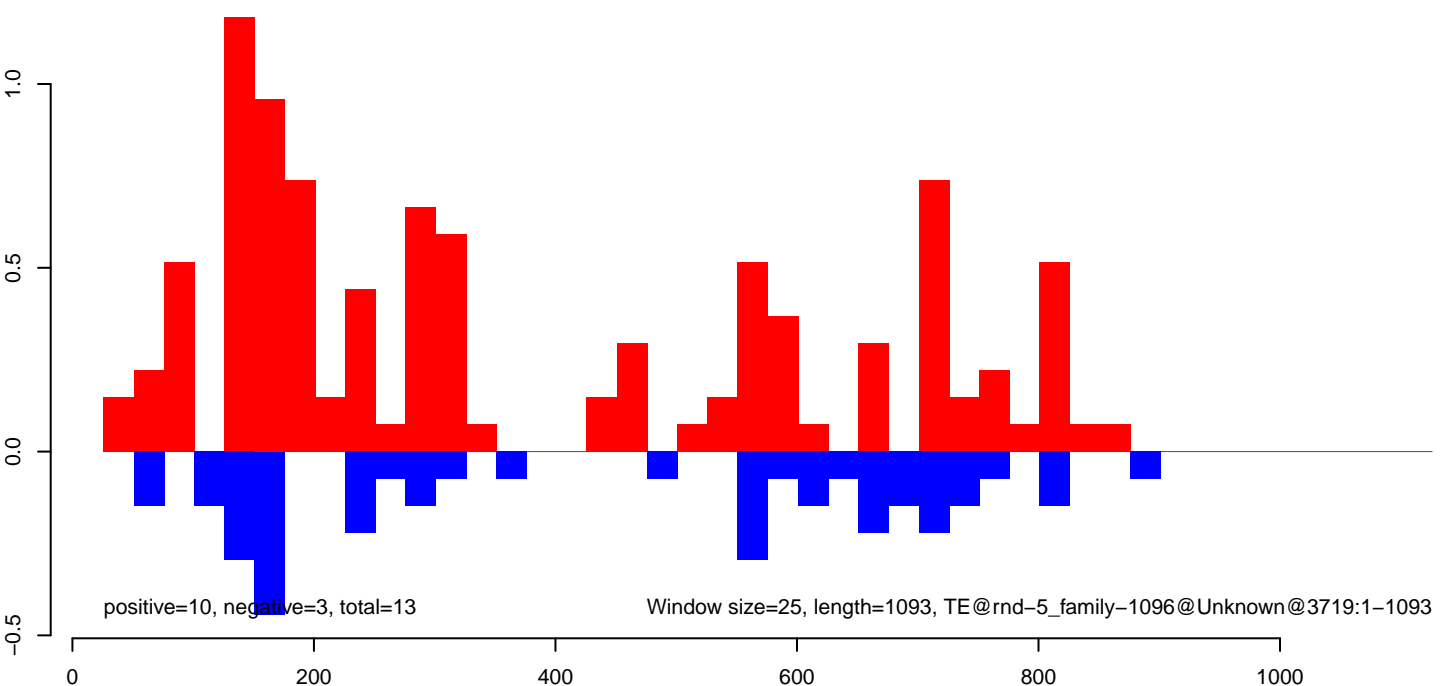
CuQuin_Testes_NL.18_23.rep



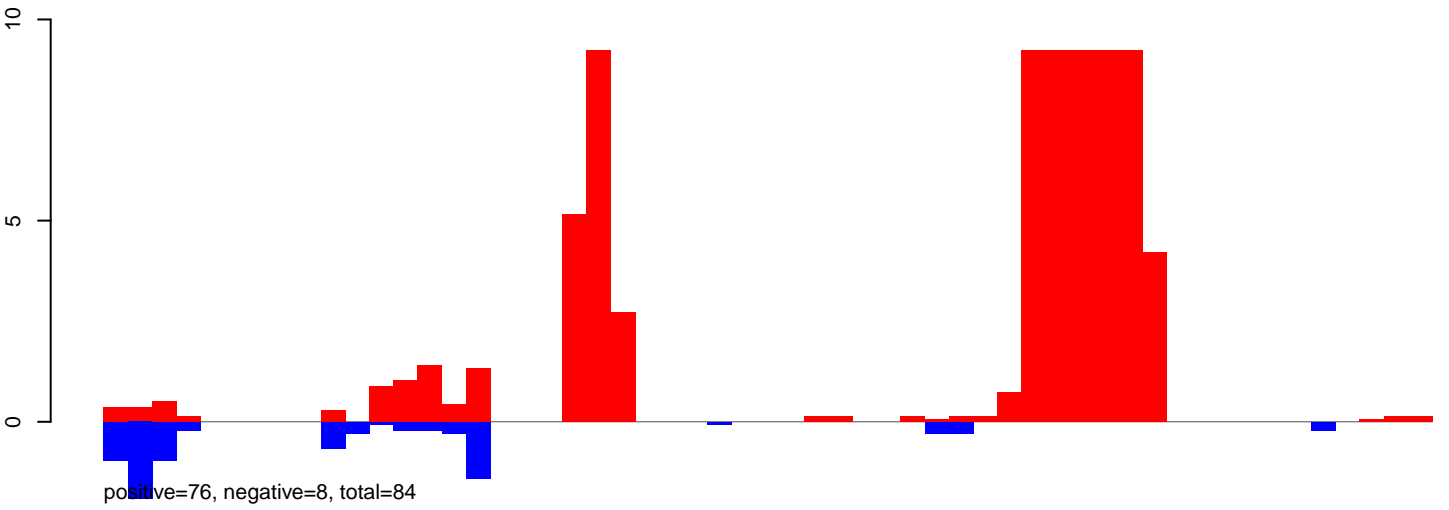
CuQuin_Testes_NL.24_35.rep



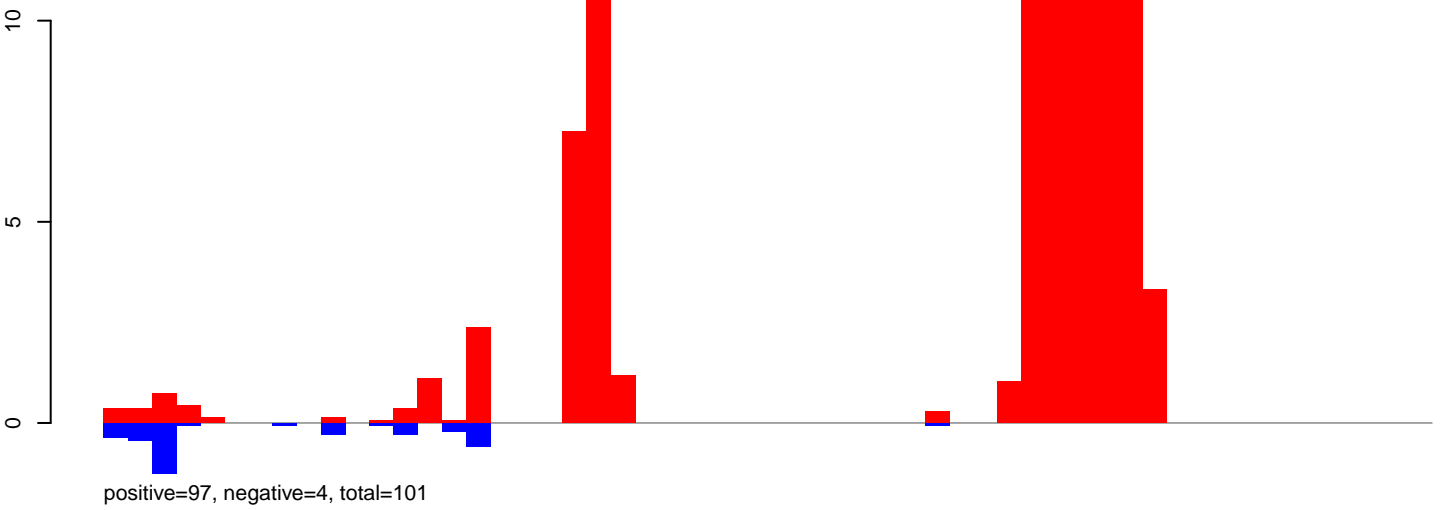
CuQuin_Testes_NL.rep



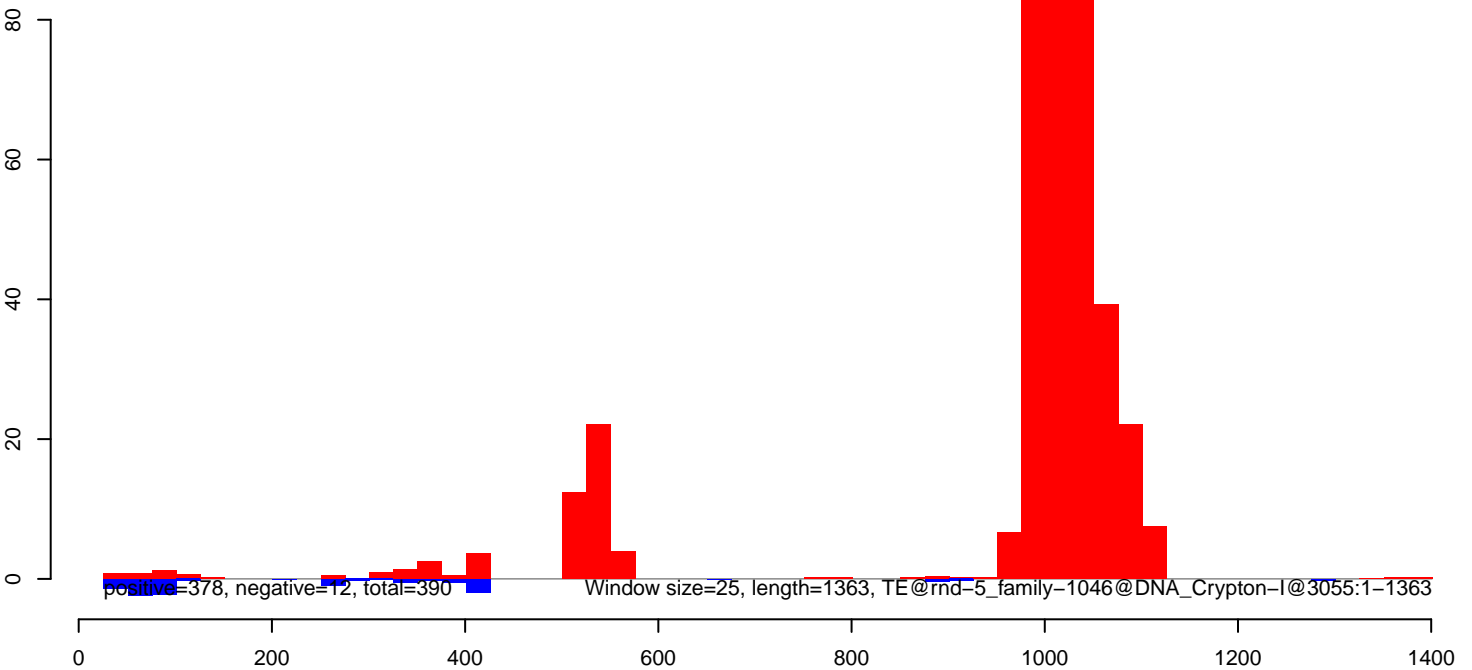
CuQuin_Testes_NL.18_23.rep



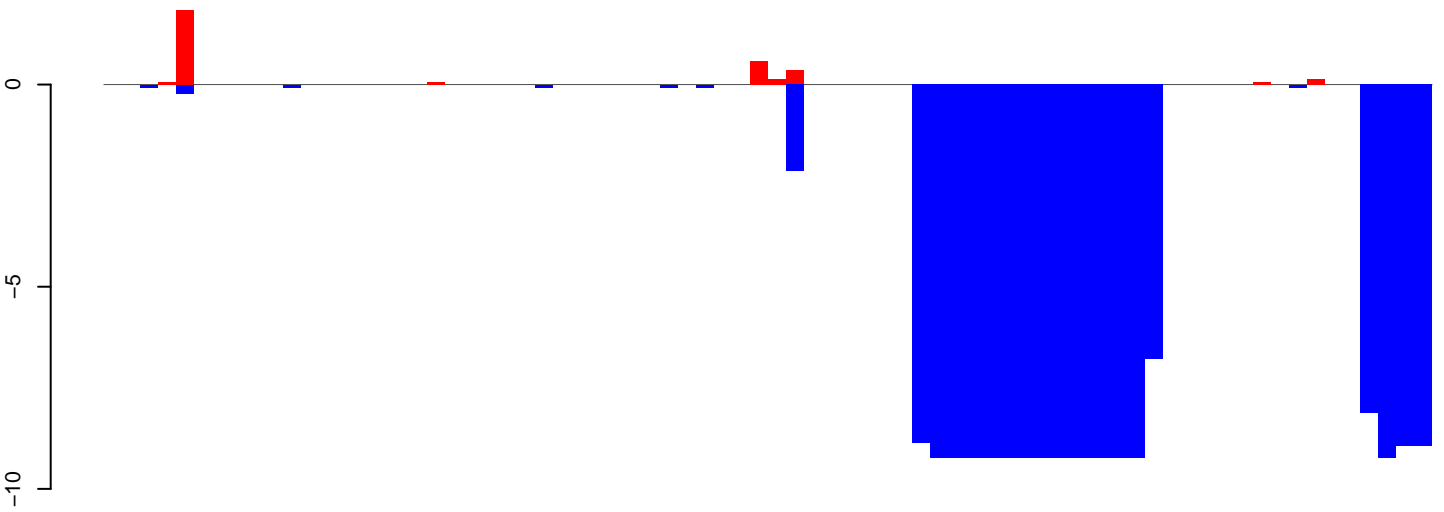
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



CuQuin_Testes_NL.18_23.rep



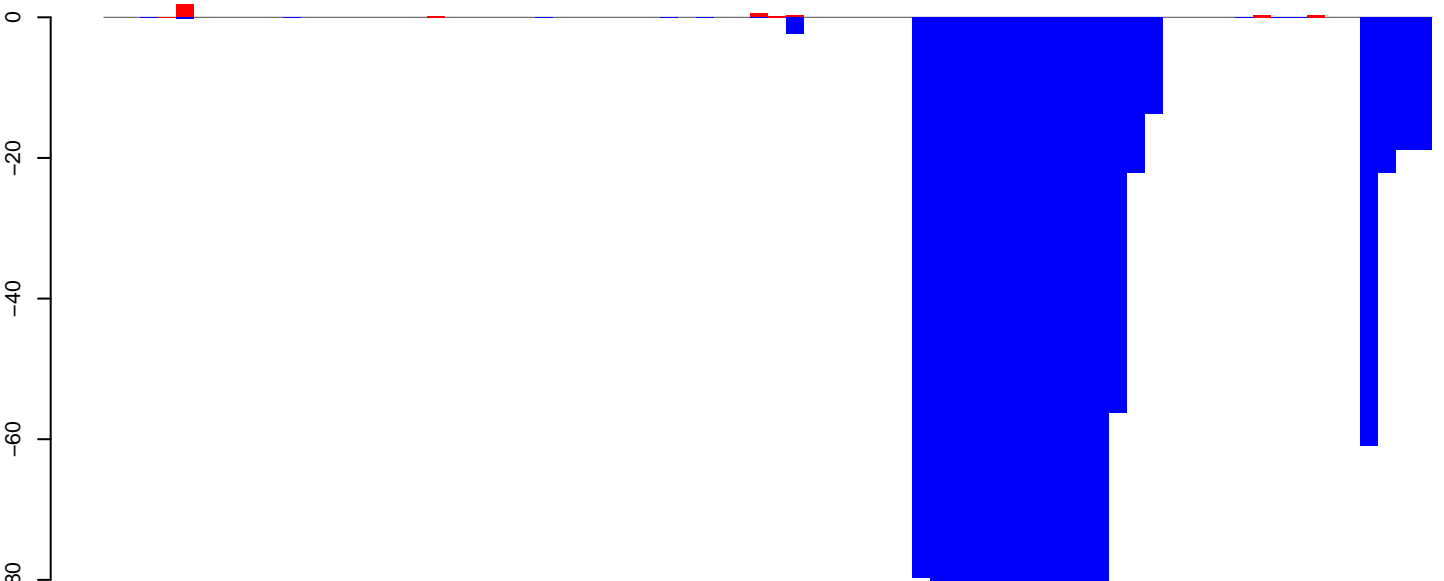
positive=3, negative=164, total=168

CuQuin_Testes_NL.24_35.rep



positive=1, negative=219, total=220

CuQuin_Testes_NL.rep

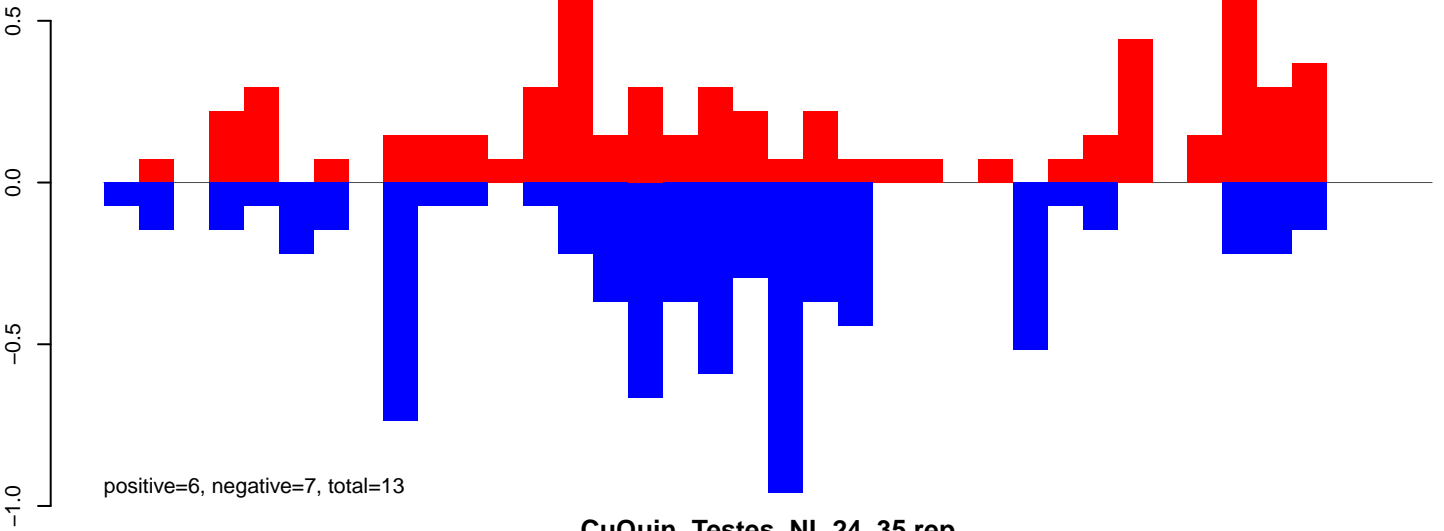


positive=4, negative=1127, total=1130

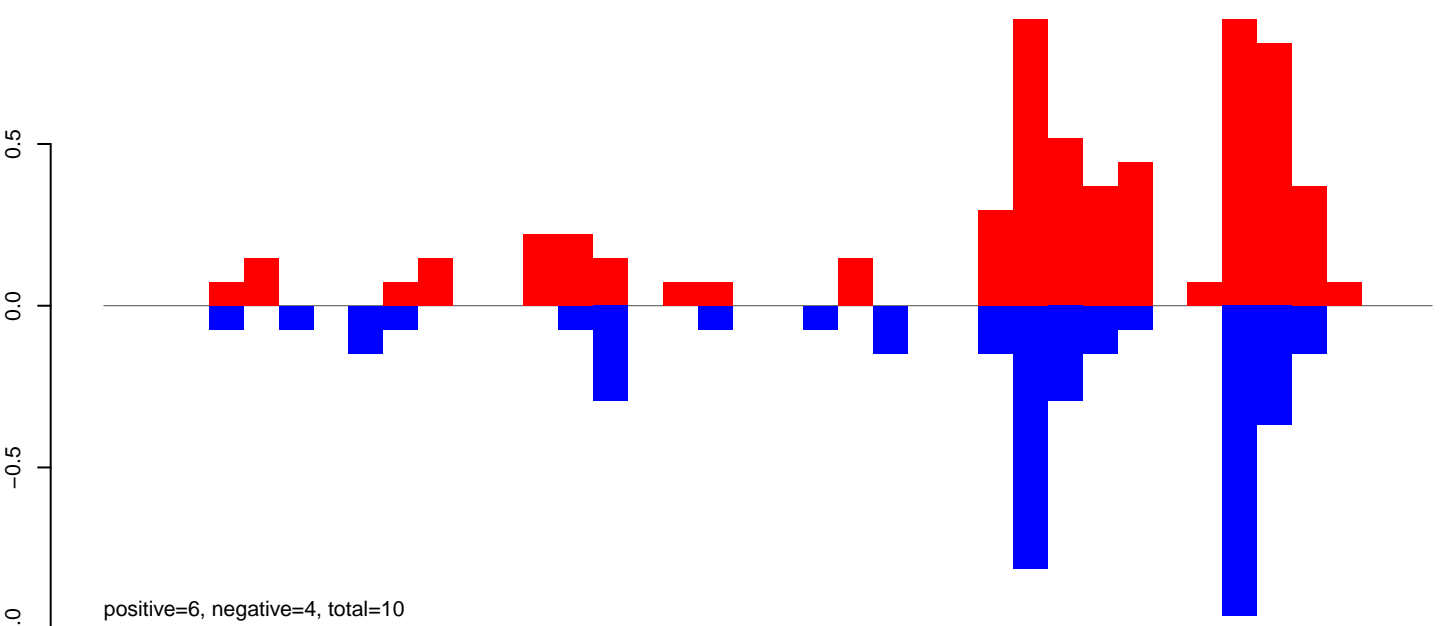
Window size=25, length=1846, TE@rmd-4_family-691@DNA_Sola-2@2738:1-1846

0 500 1000 1500

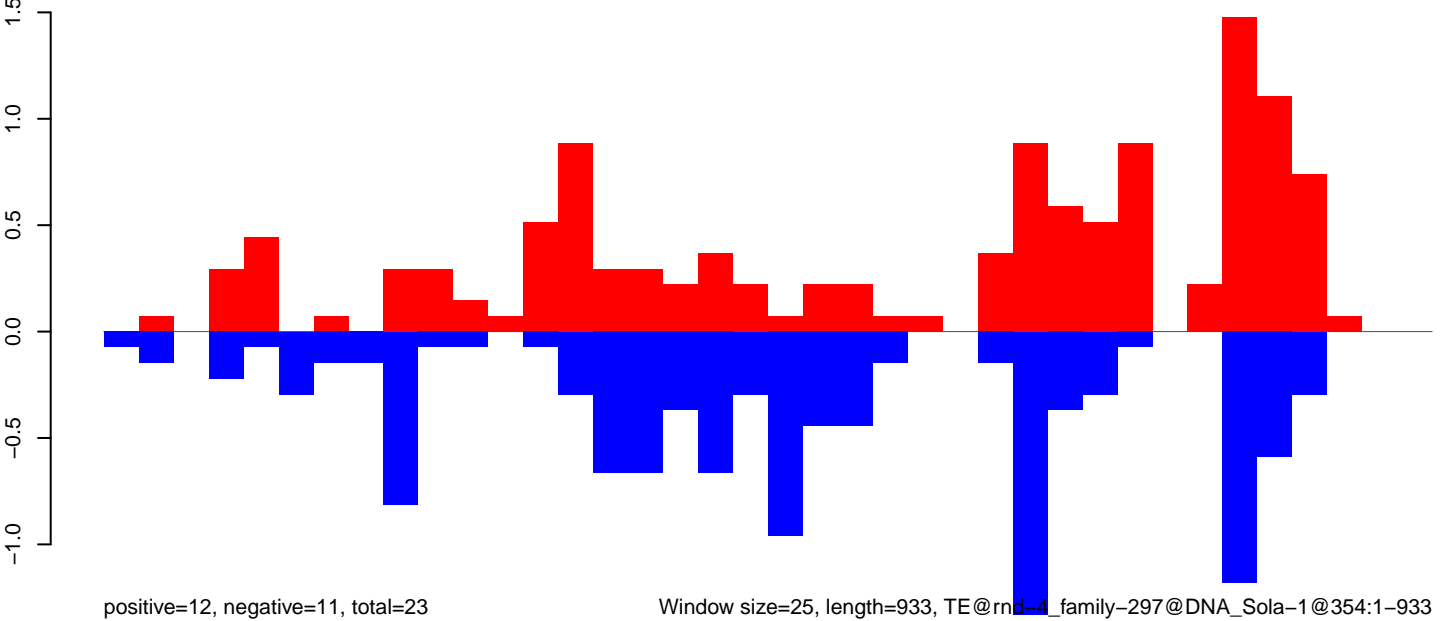
CuQuin_Testes_NL.18_23.rep



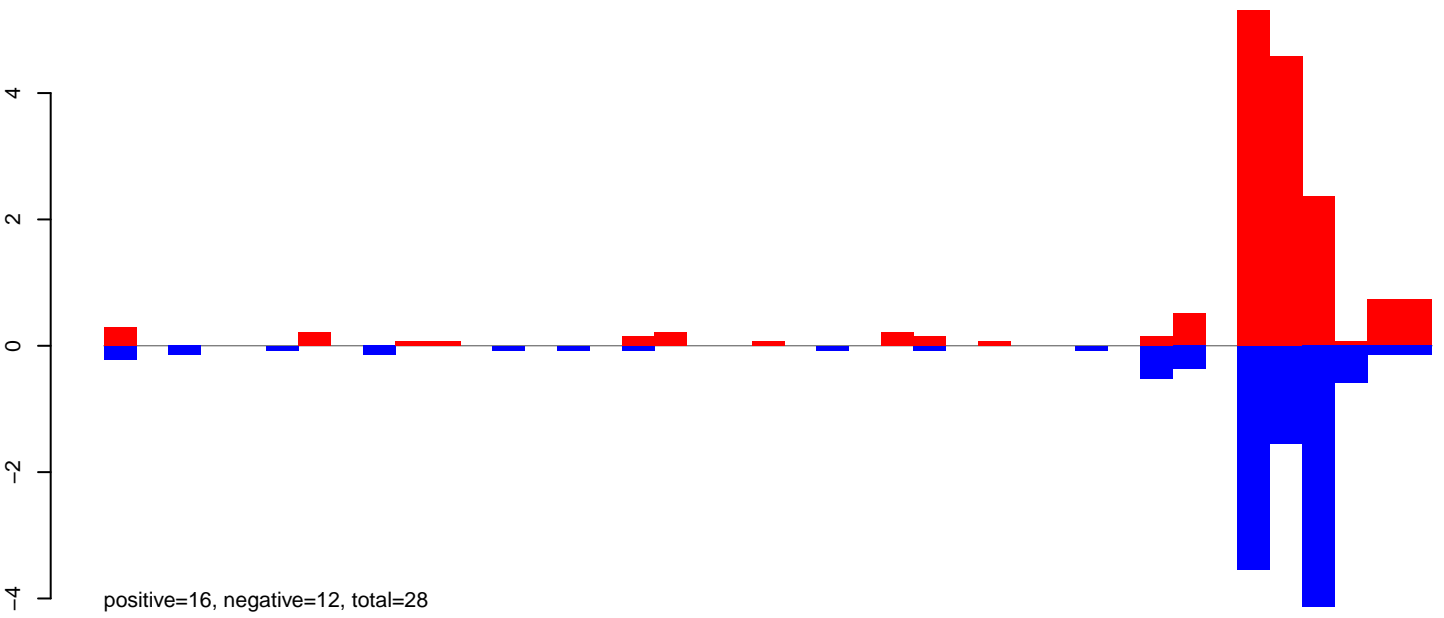
CuQuin_Testes_NL.24_35.rep



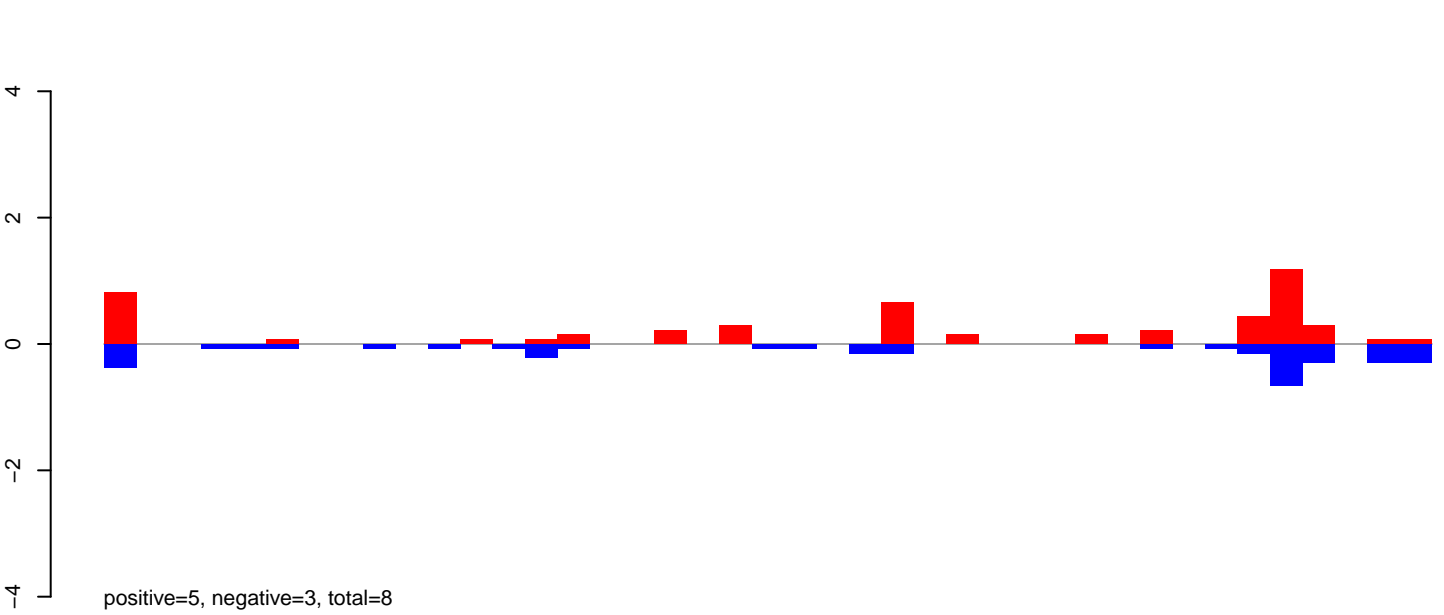
CuQuin_Testes_NL.rep



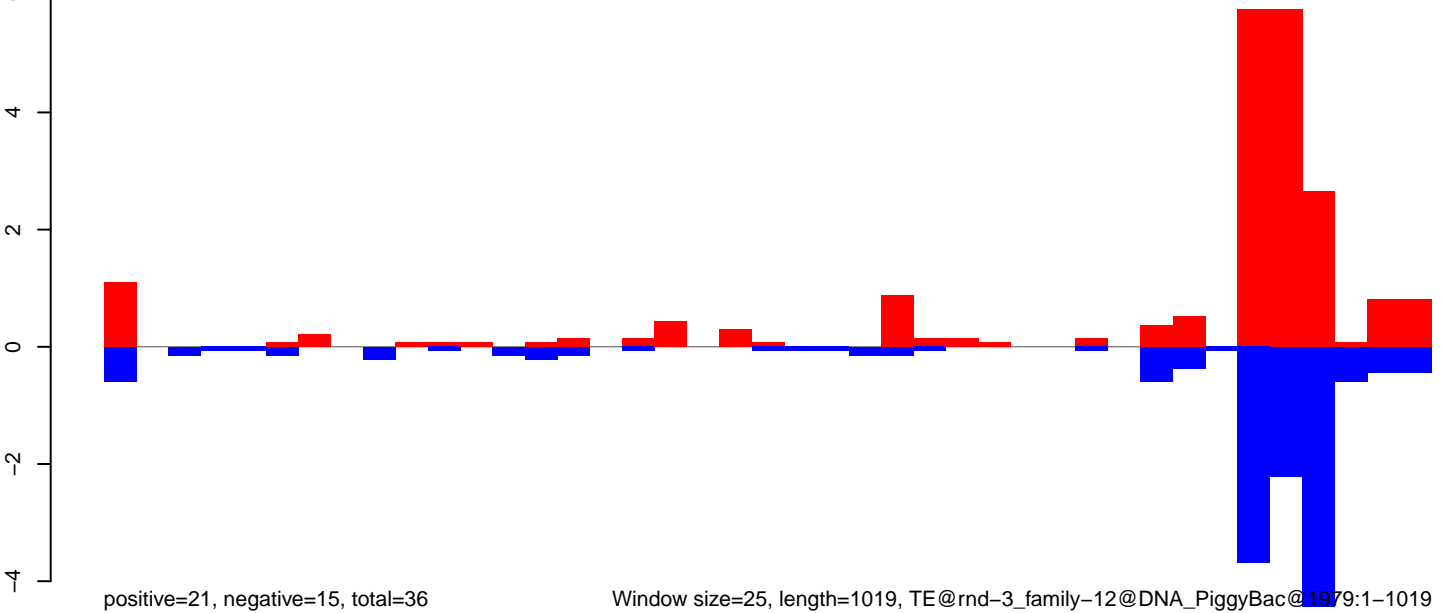
CuQuin_Testes_NL.18_23.rep



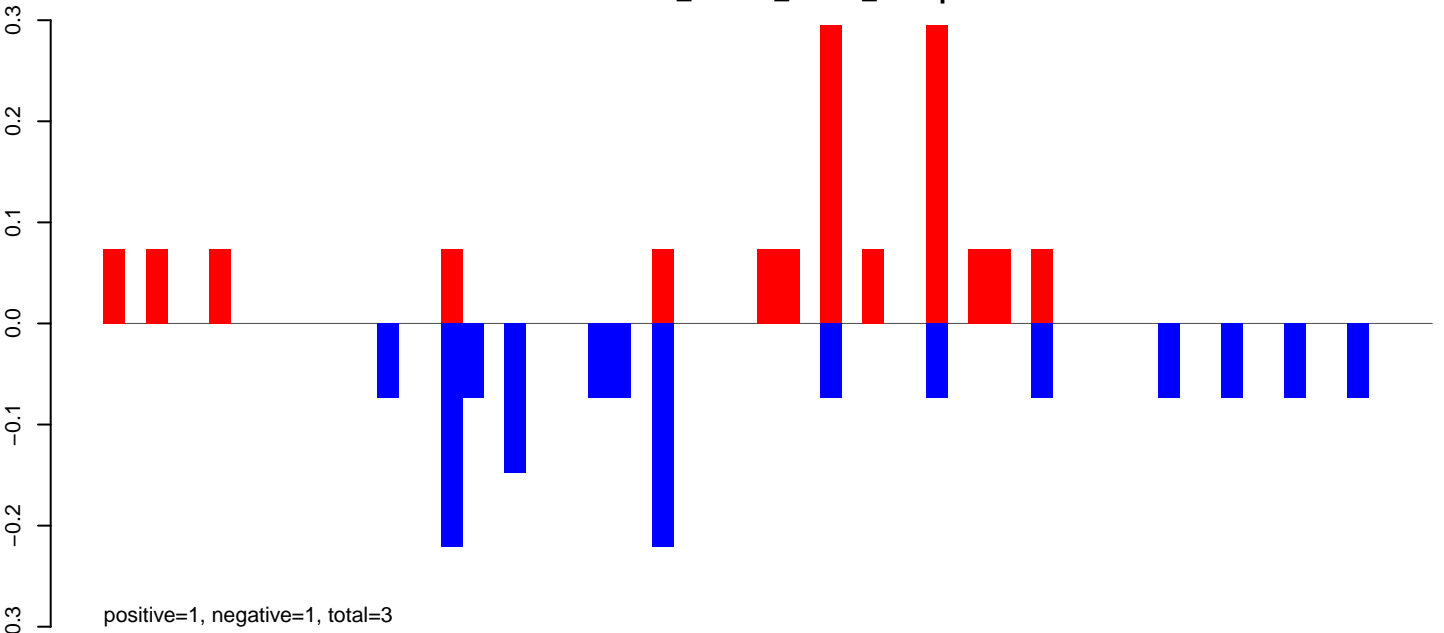
CuQuin_Testes_NL.24_35.rep



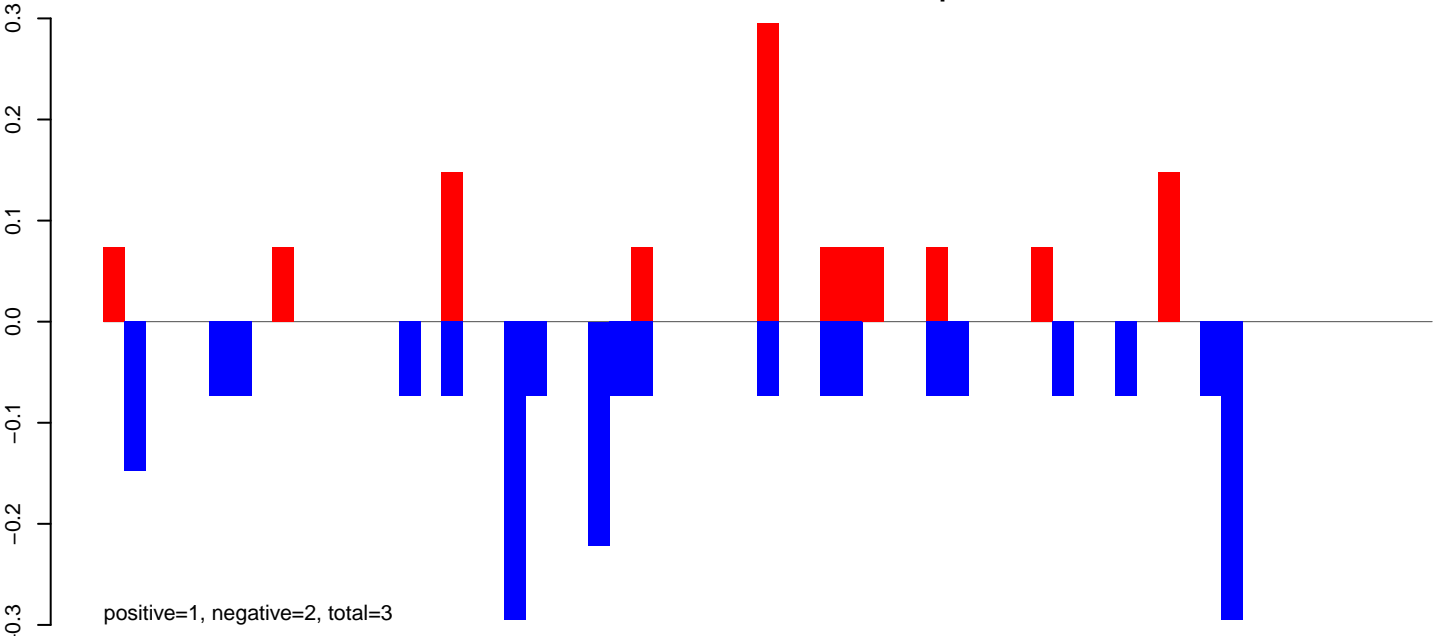
CuQuin_Testes_NL.rep



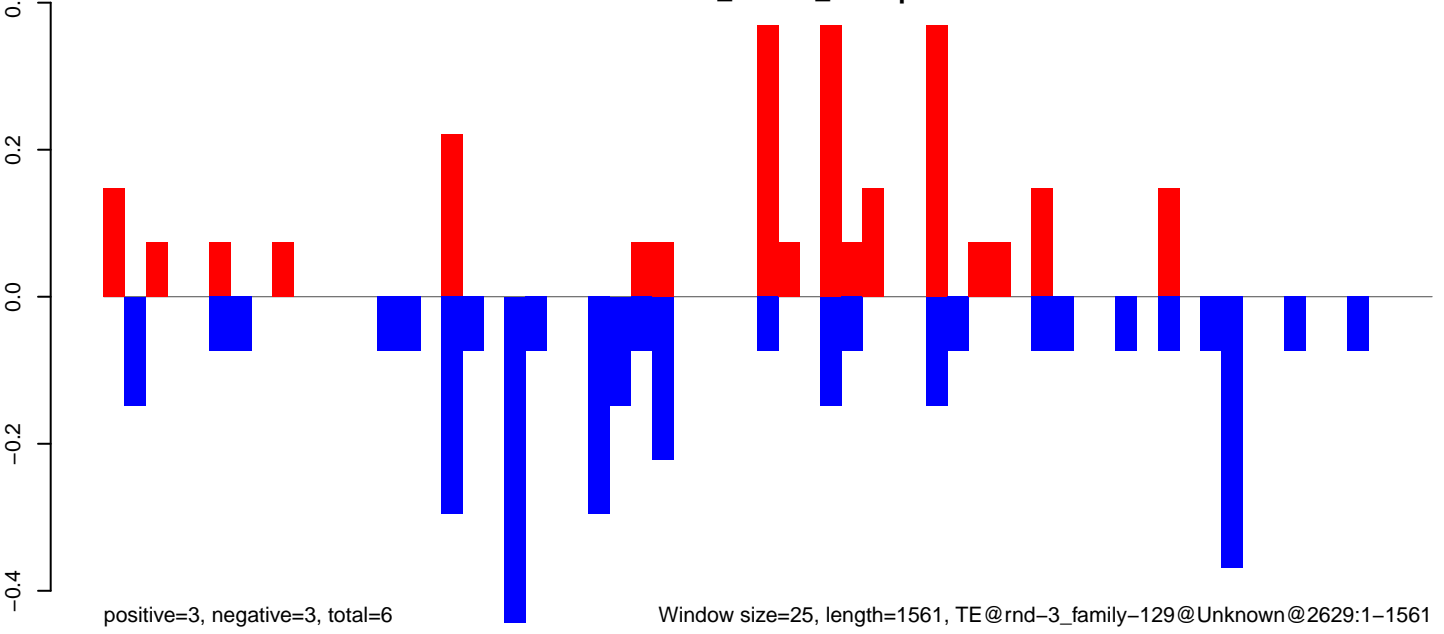
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



Window size=25, length=1561, TE@rnd-3_family-129@Unknown@2629:1-1561

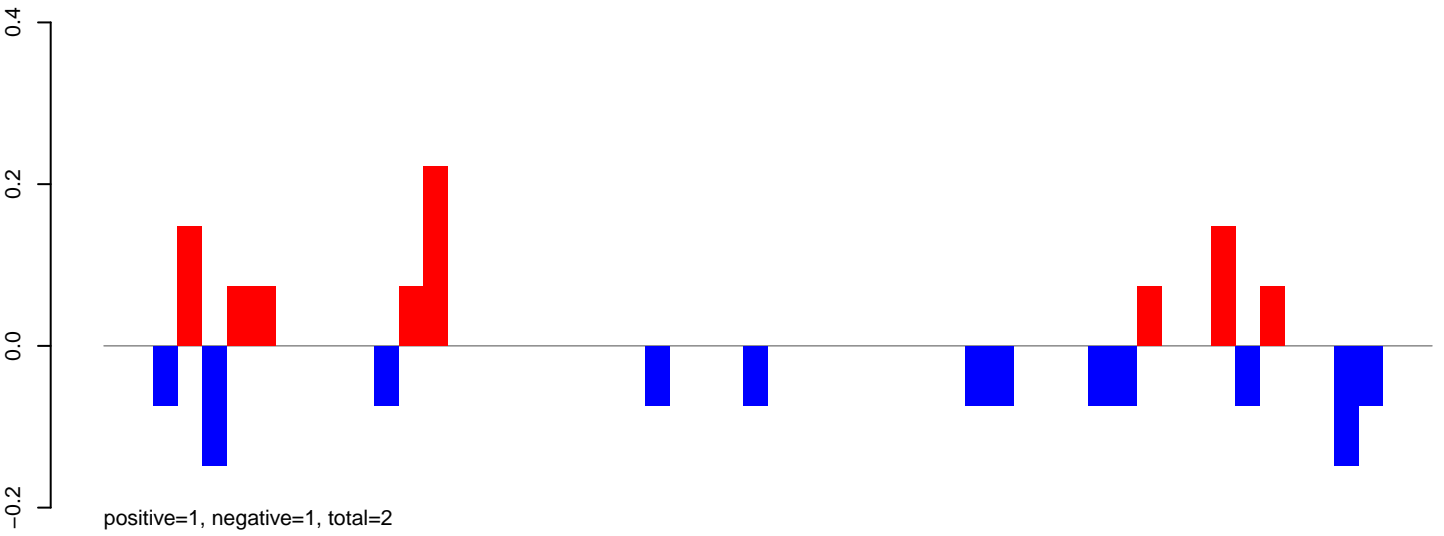
0

500

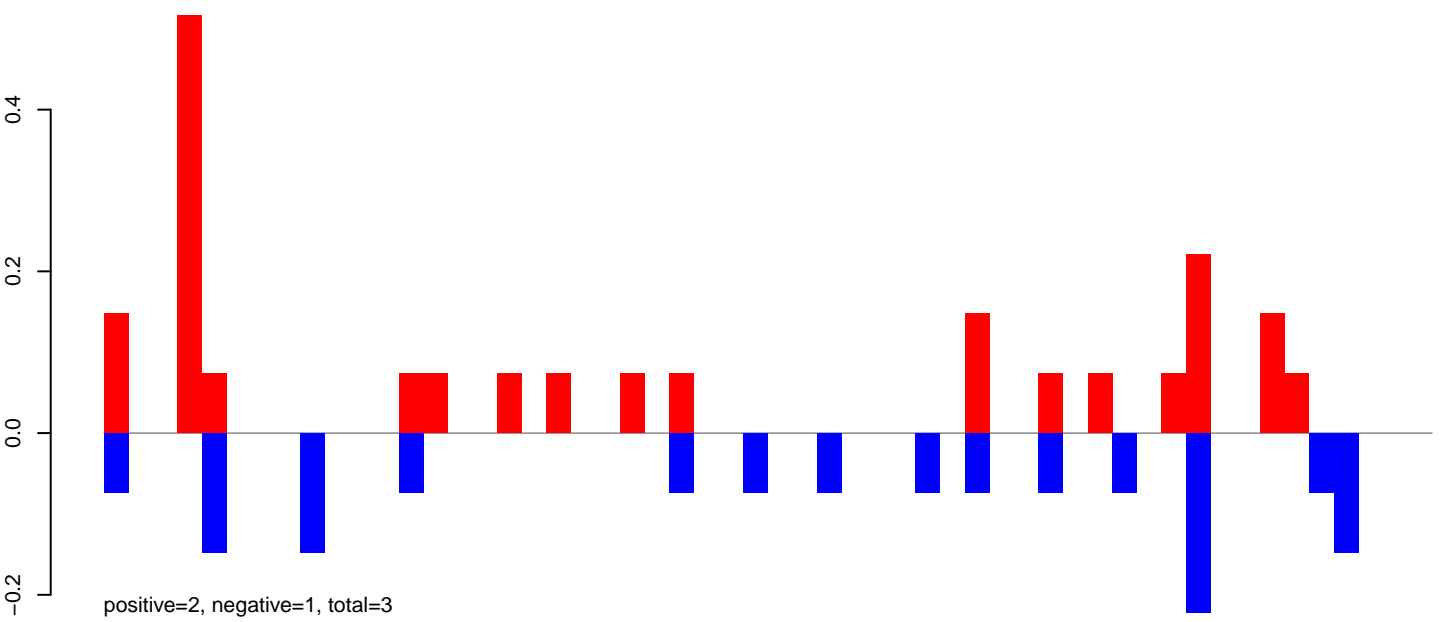
1000

1500

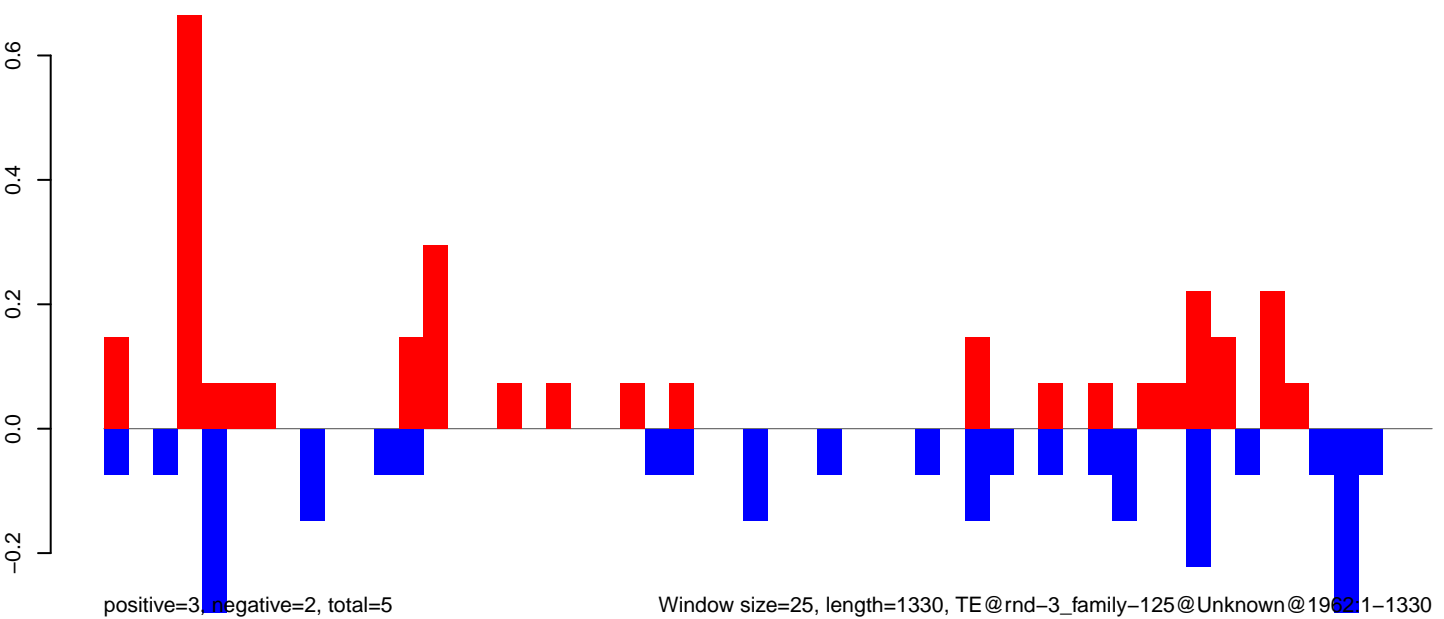
CuQuin_Testes_NL.18_23.rep



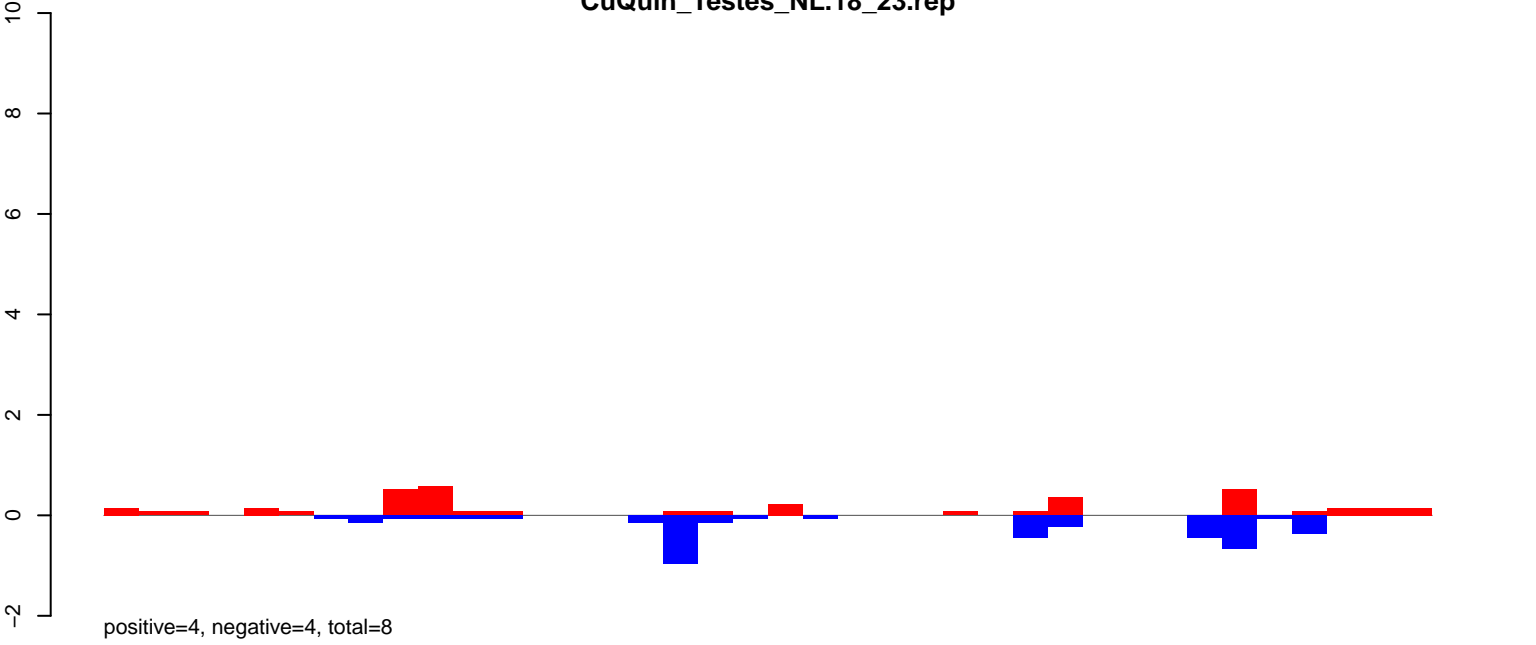
CuQuin_Testes_NL.24_35.rep



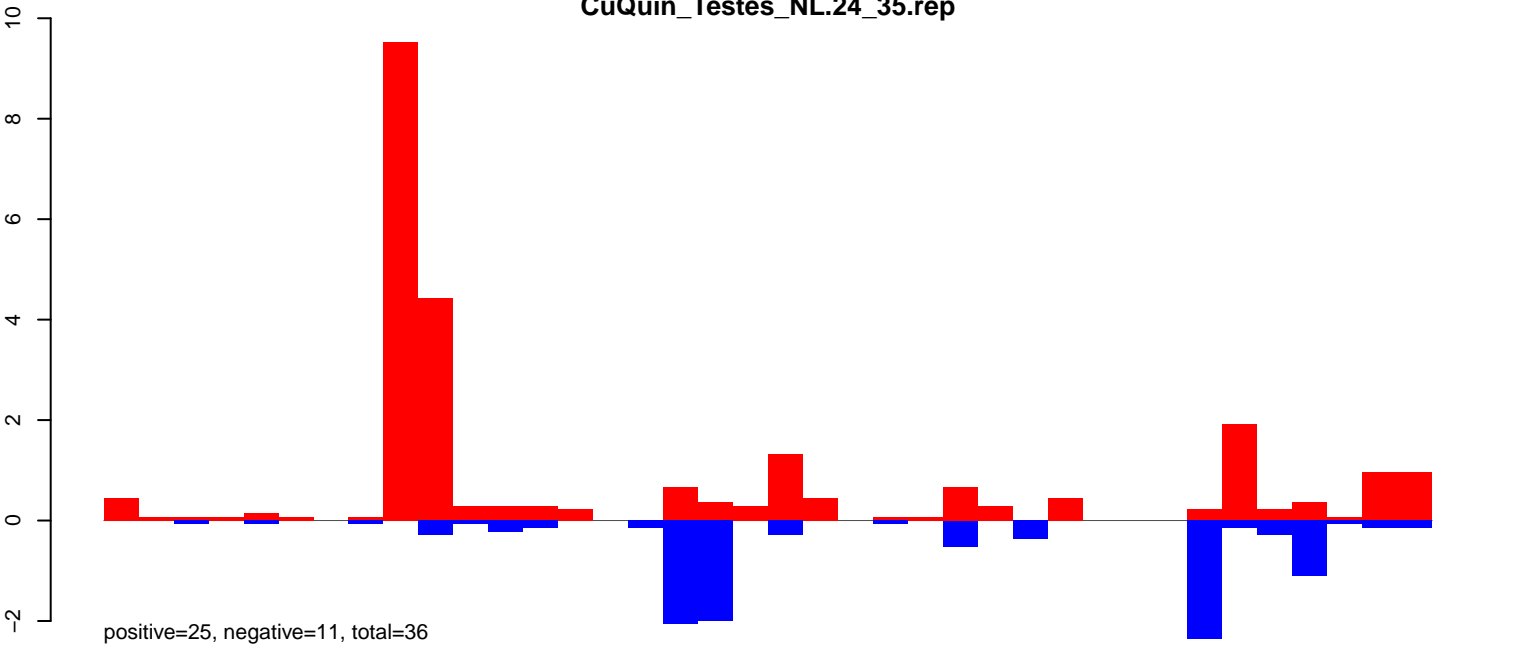
CuQuin_Testes_NL.rep



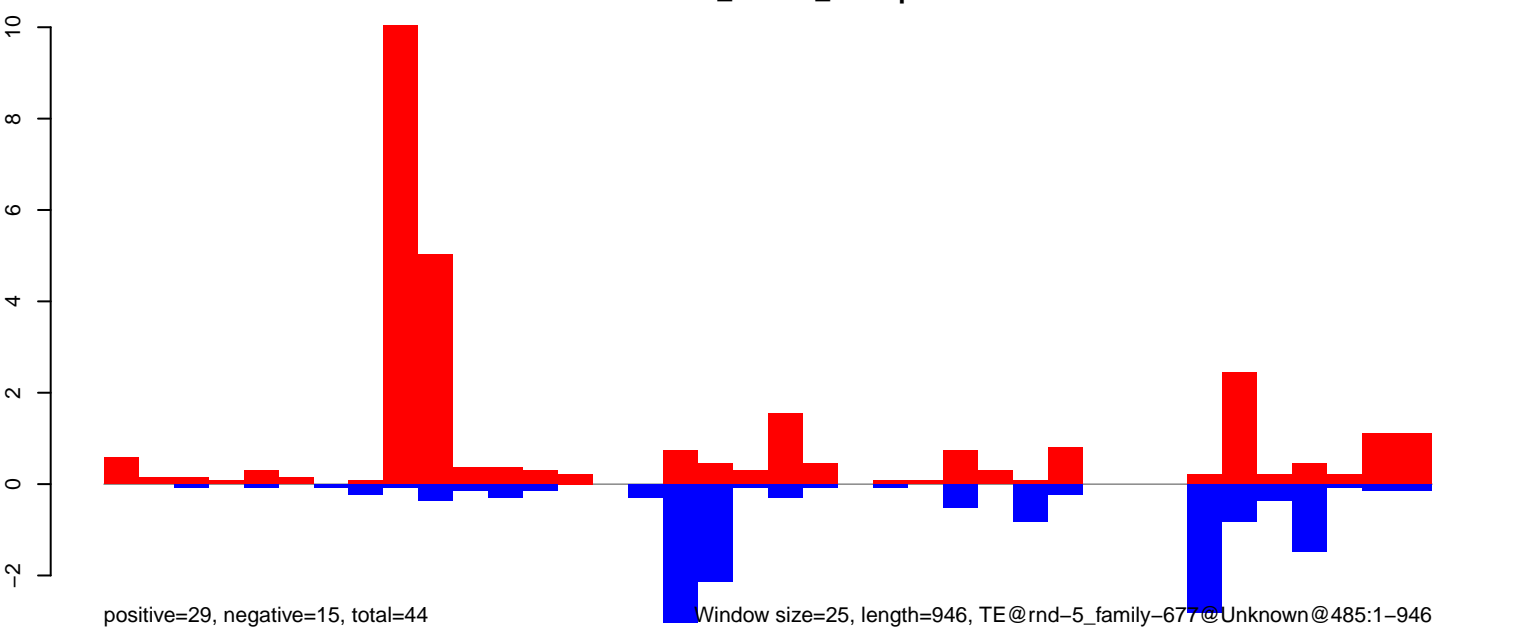
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

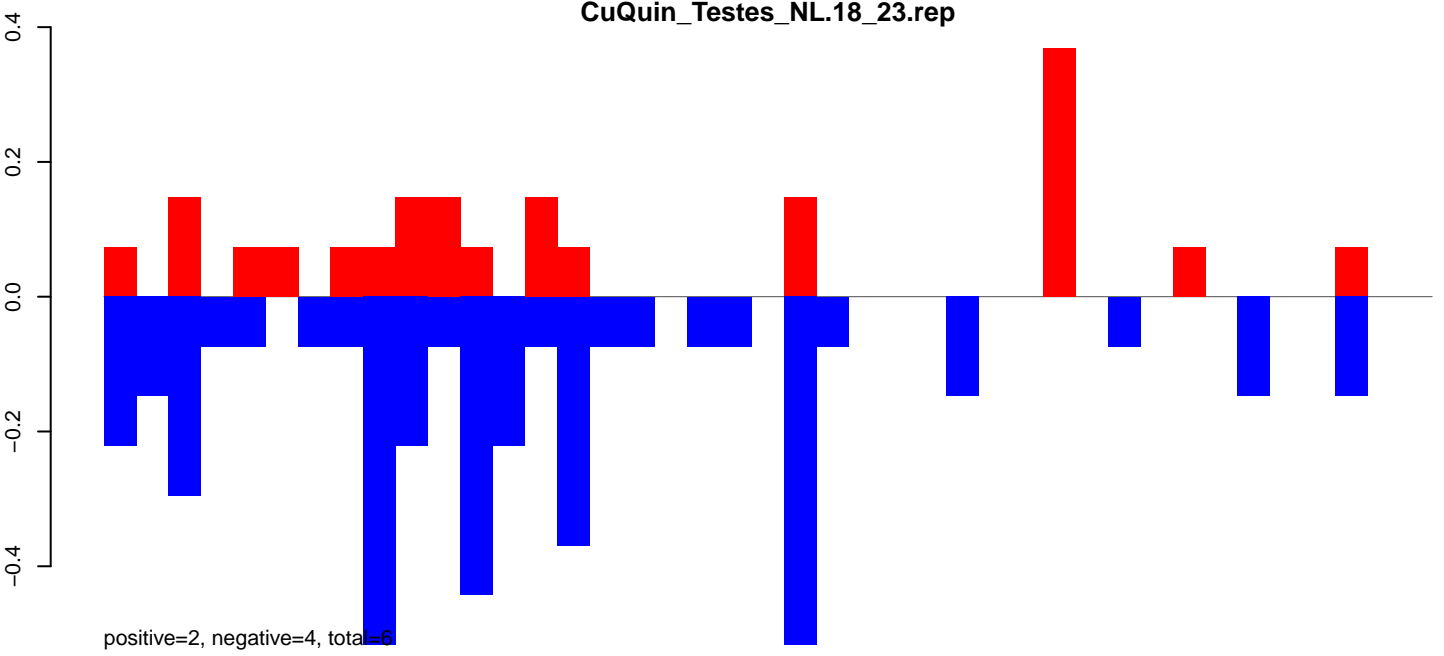


CuQuin_Testes_NL.rep

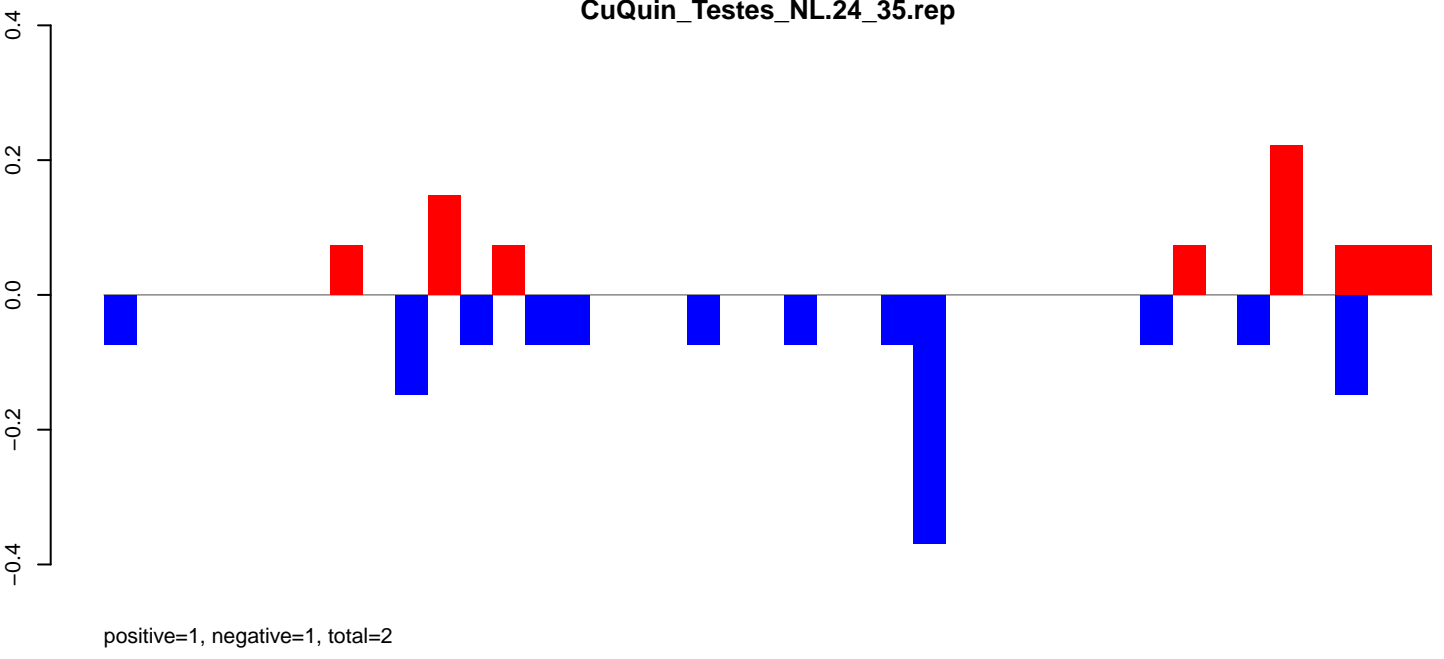


Window size=25, length=946, TE@rnd-5_family-677@Unknown@485:1-946

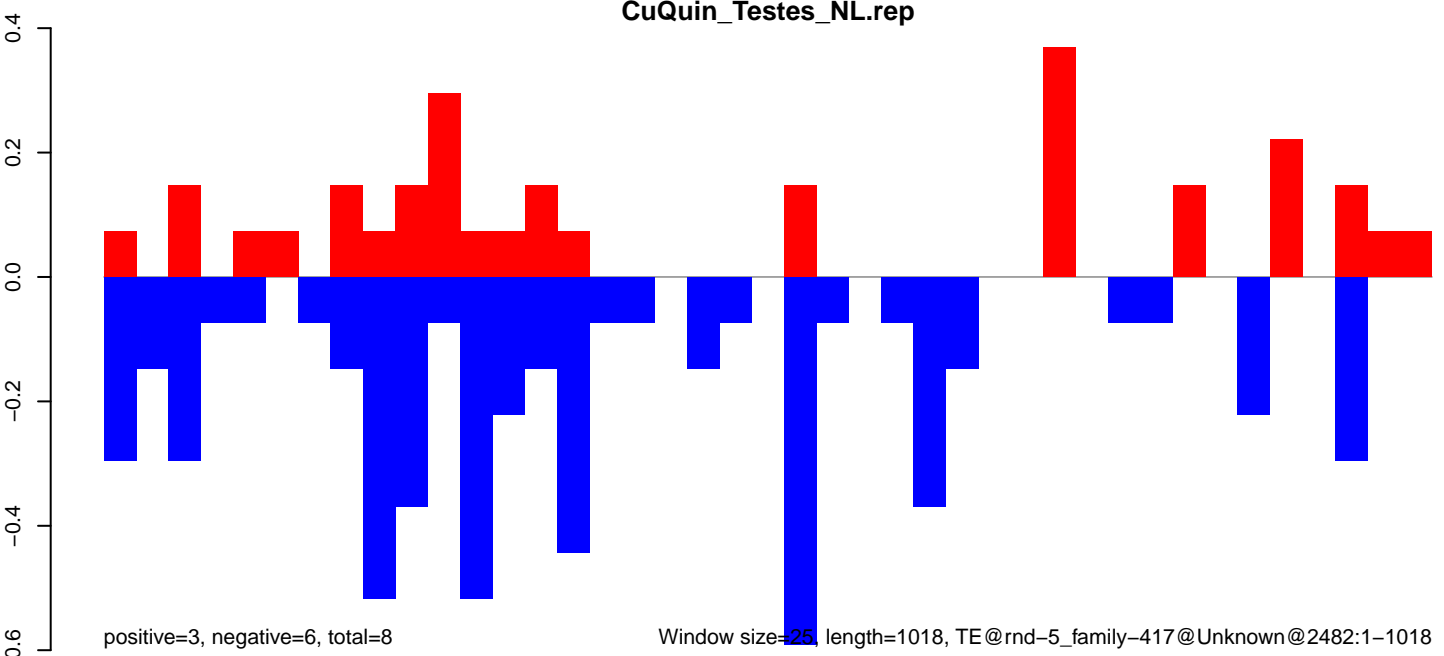
CuQuin_Testes_NL.18_23.rep



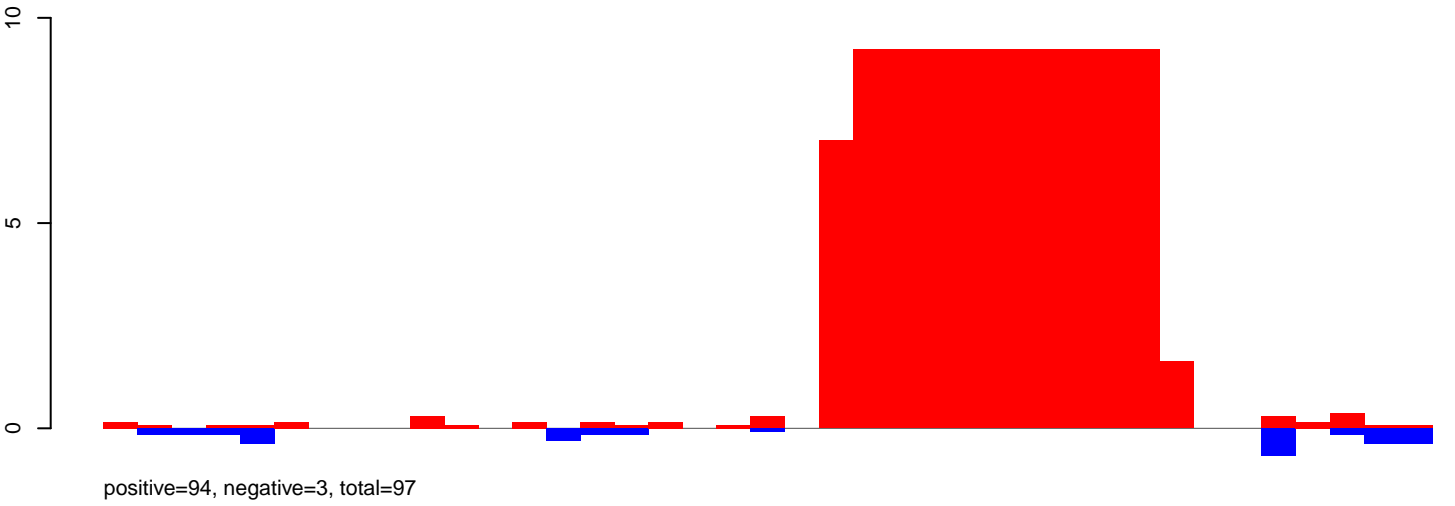
CuQuin_Testes_NL.24_35.rep



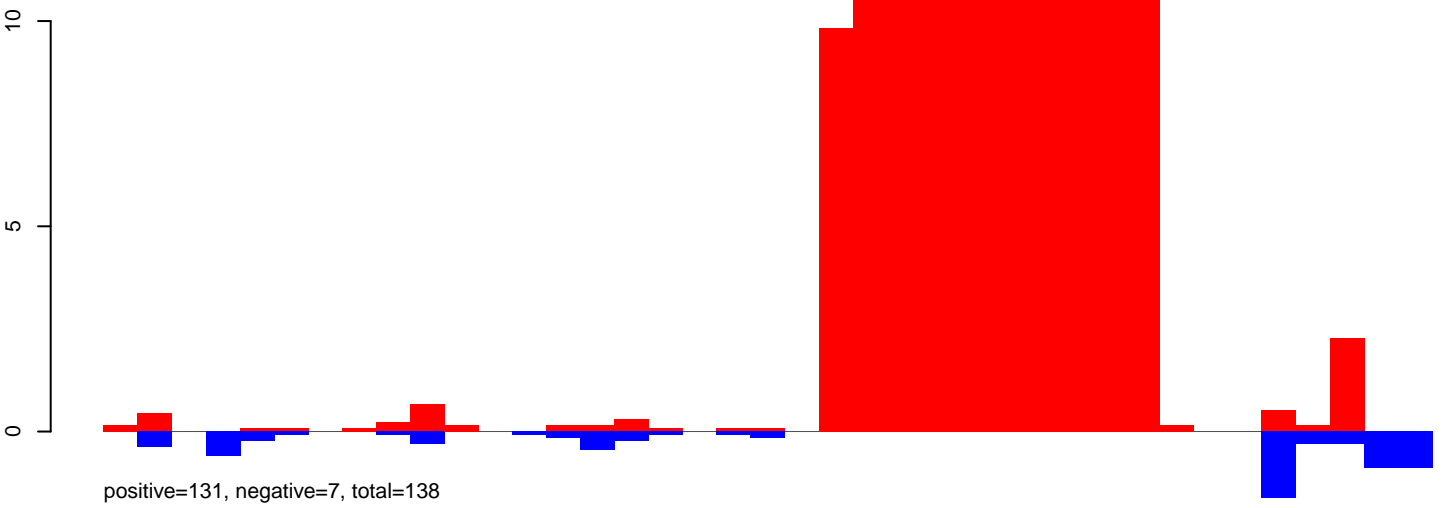
CuQuin_Testes_NL.rep



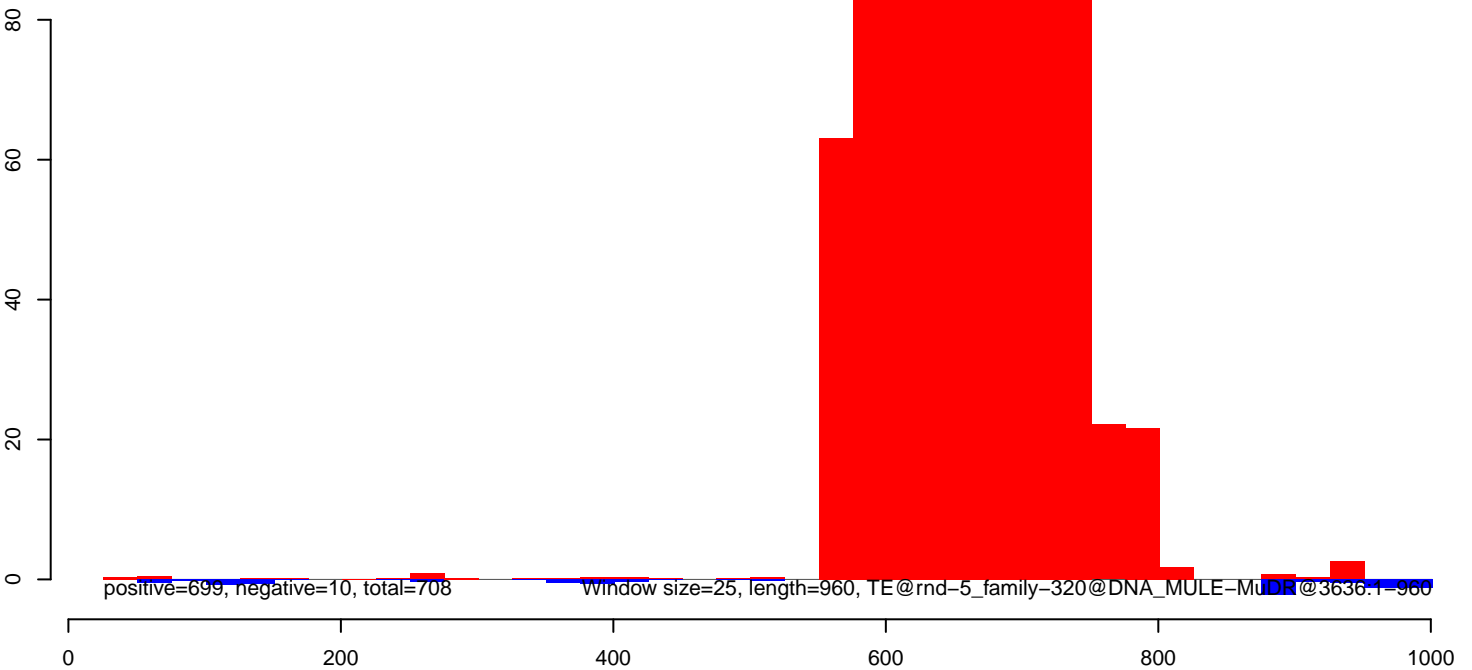
CuQuin_Testes_NL.18_23.rep



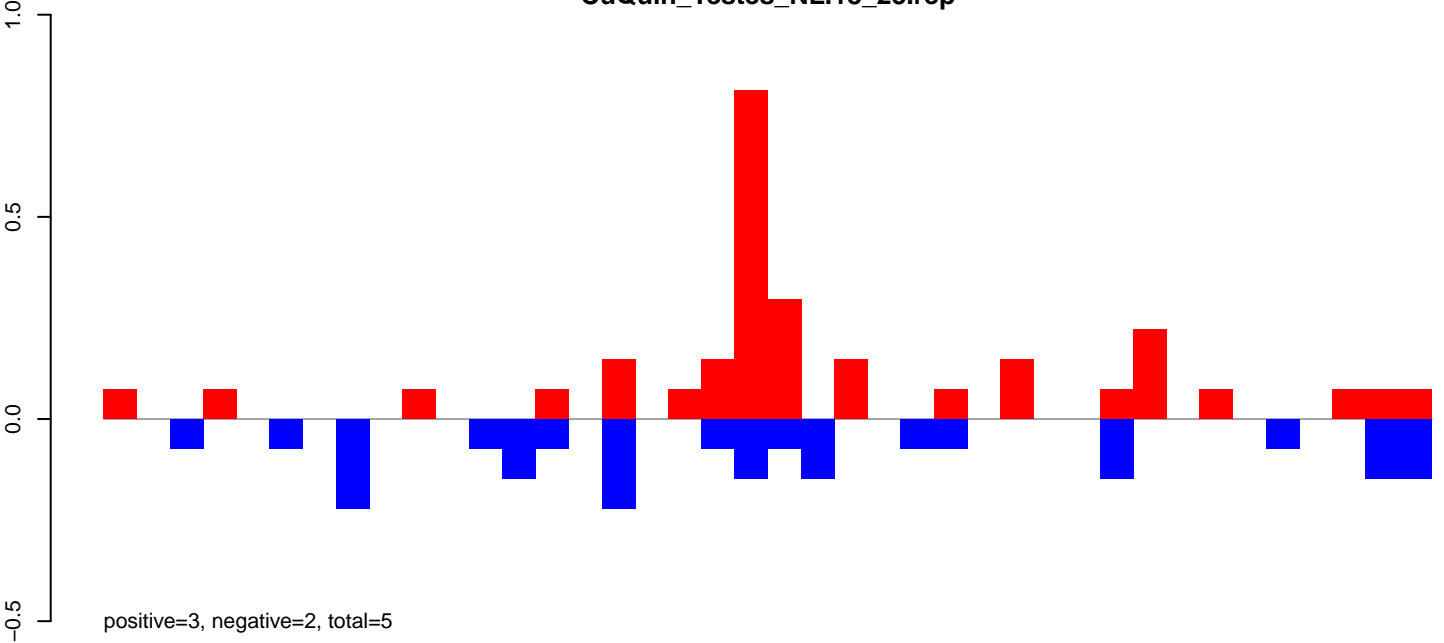
CuQuin_Testes_NL.24_35.rep



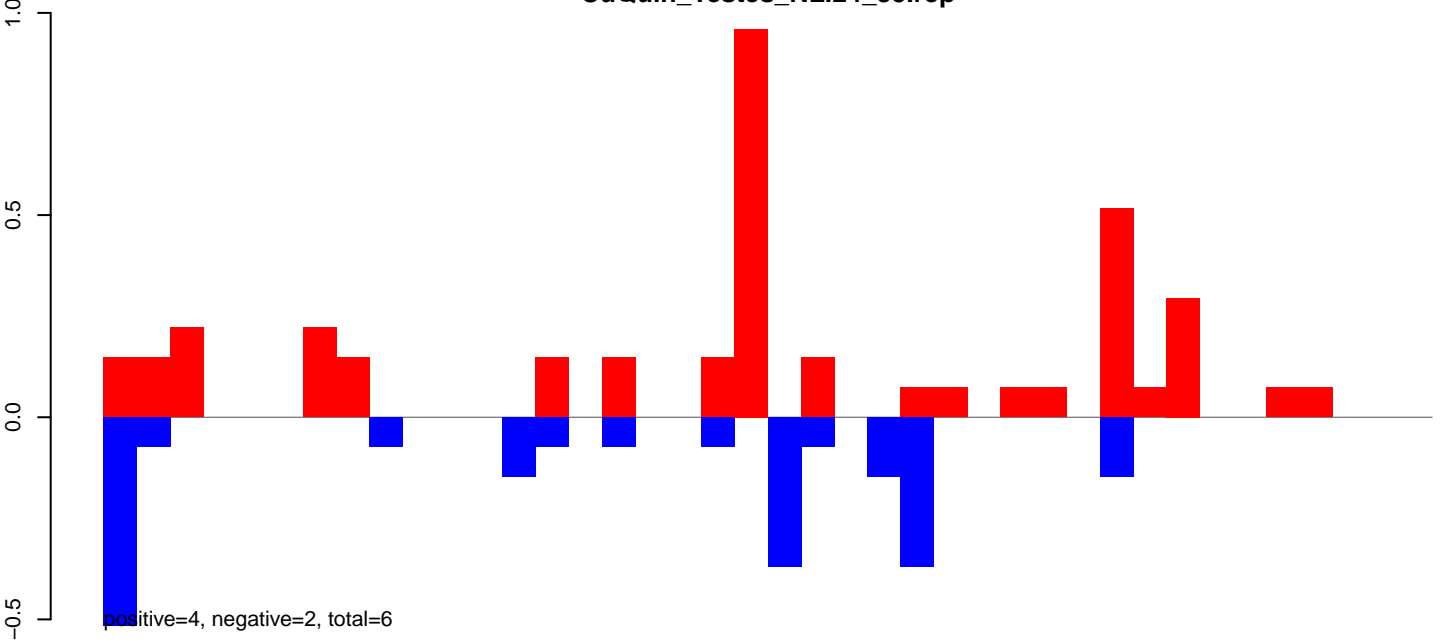
CuQuin_Testes_NL.rep



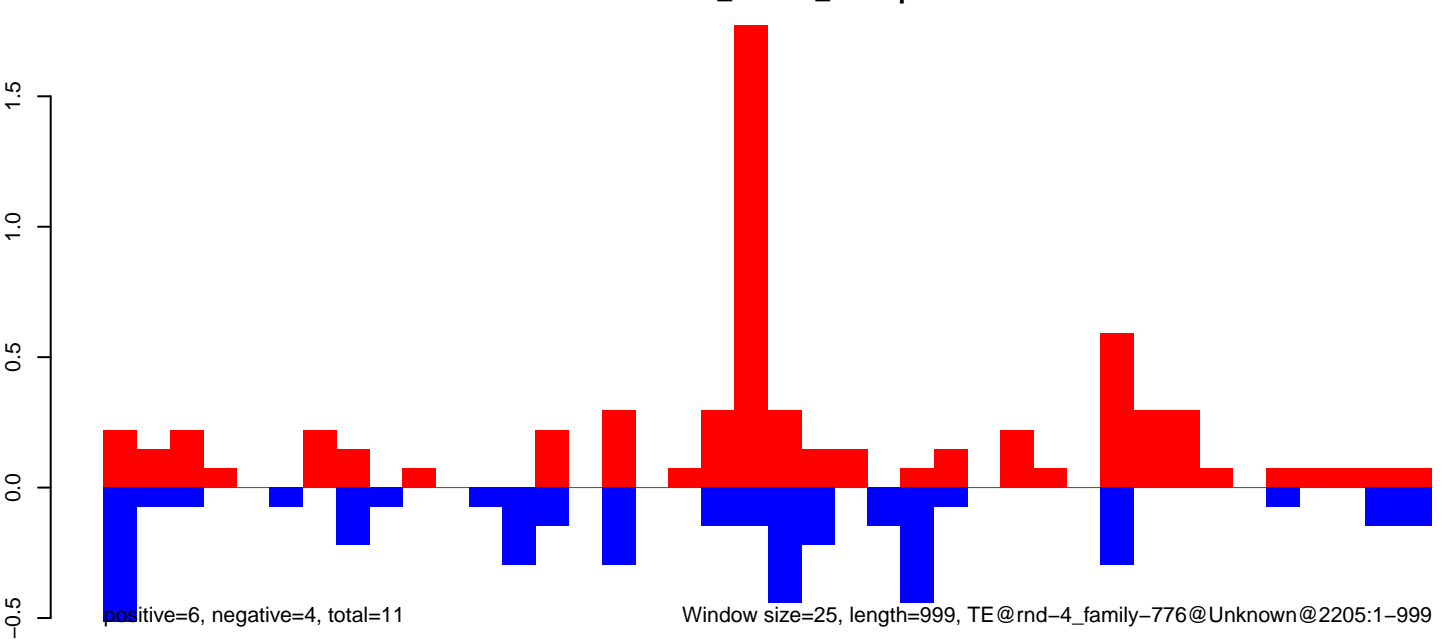
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

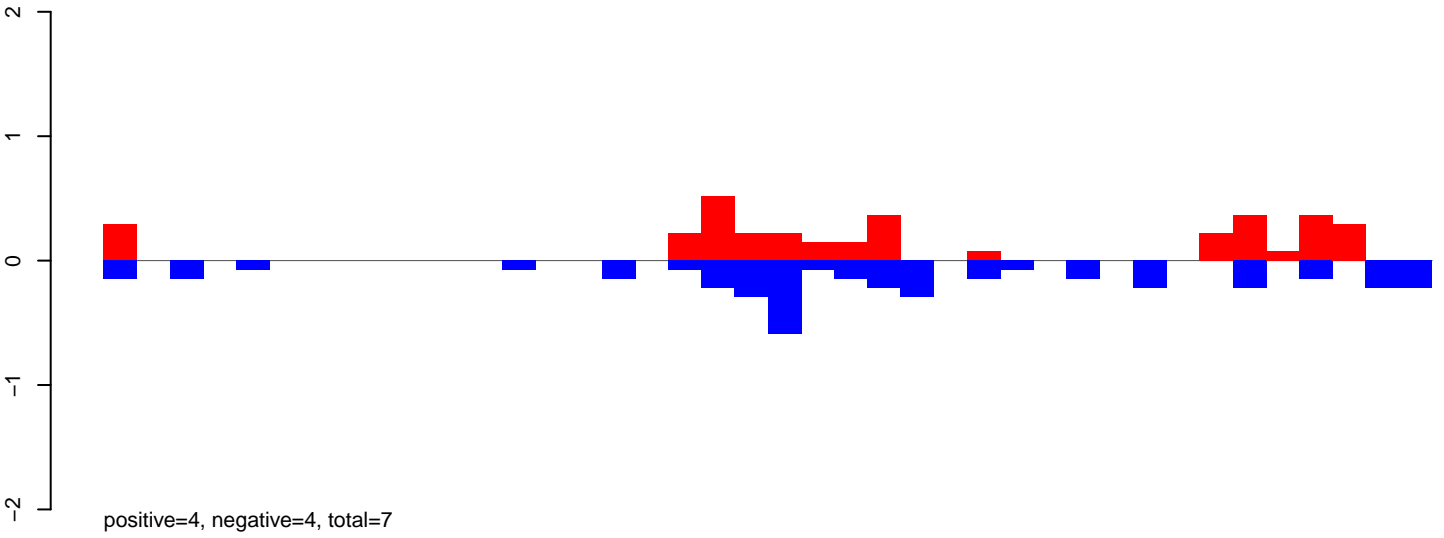


CuQuin_Testes_NL.rep

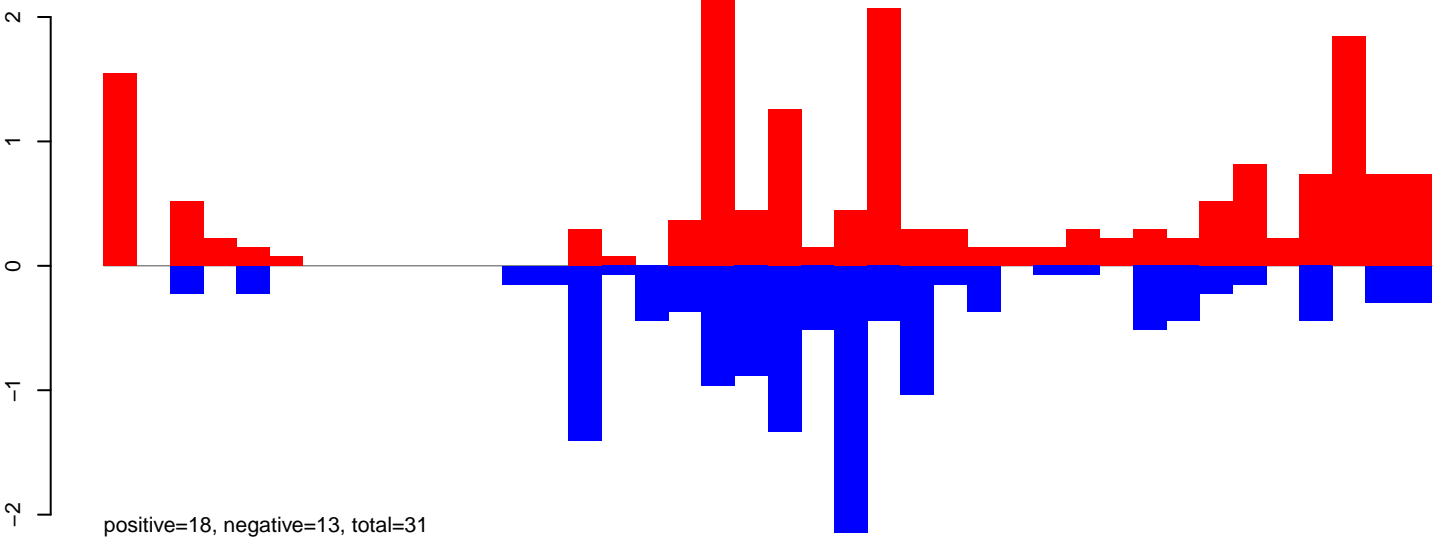


0 200 400 600 800 1000

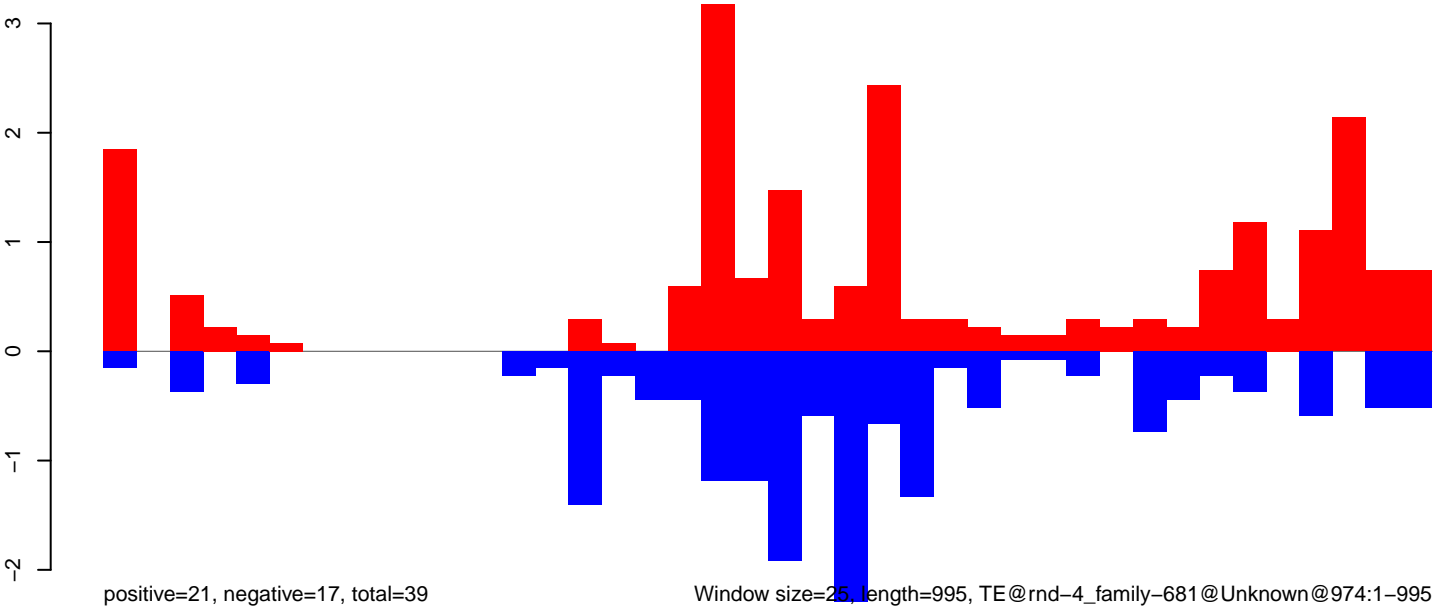
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



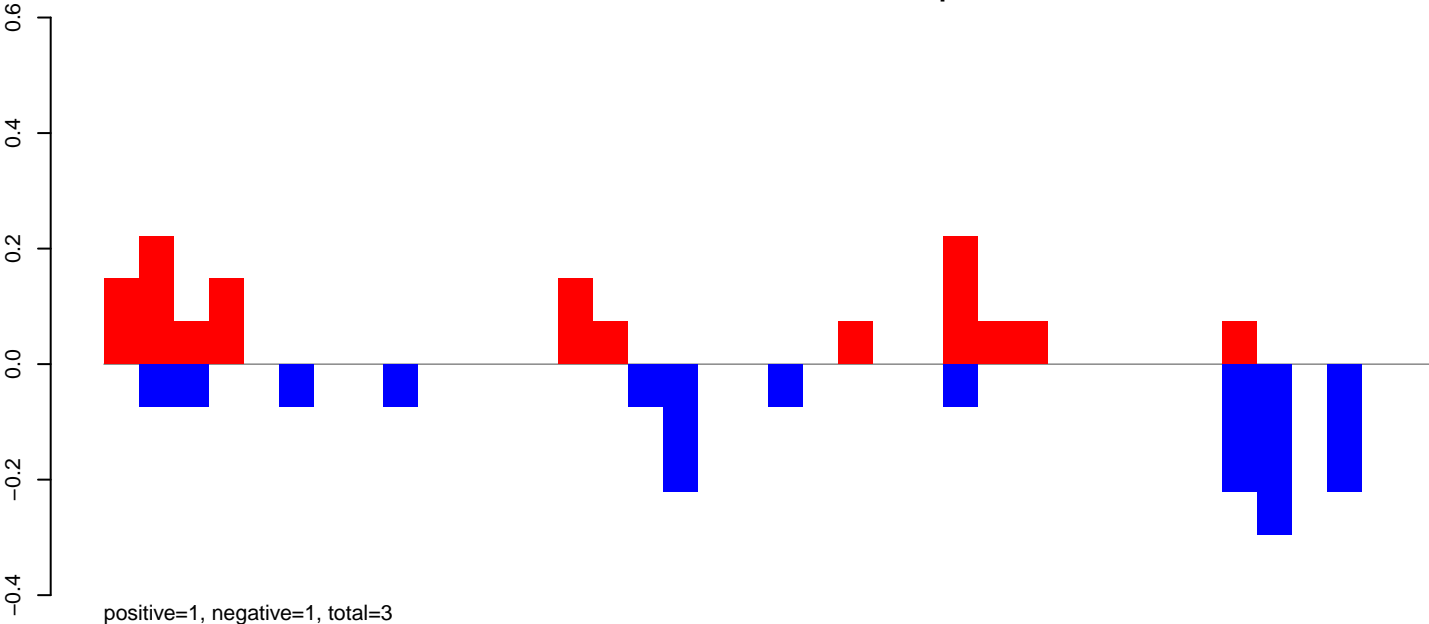
CuQuin_Testes_NL.rep



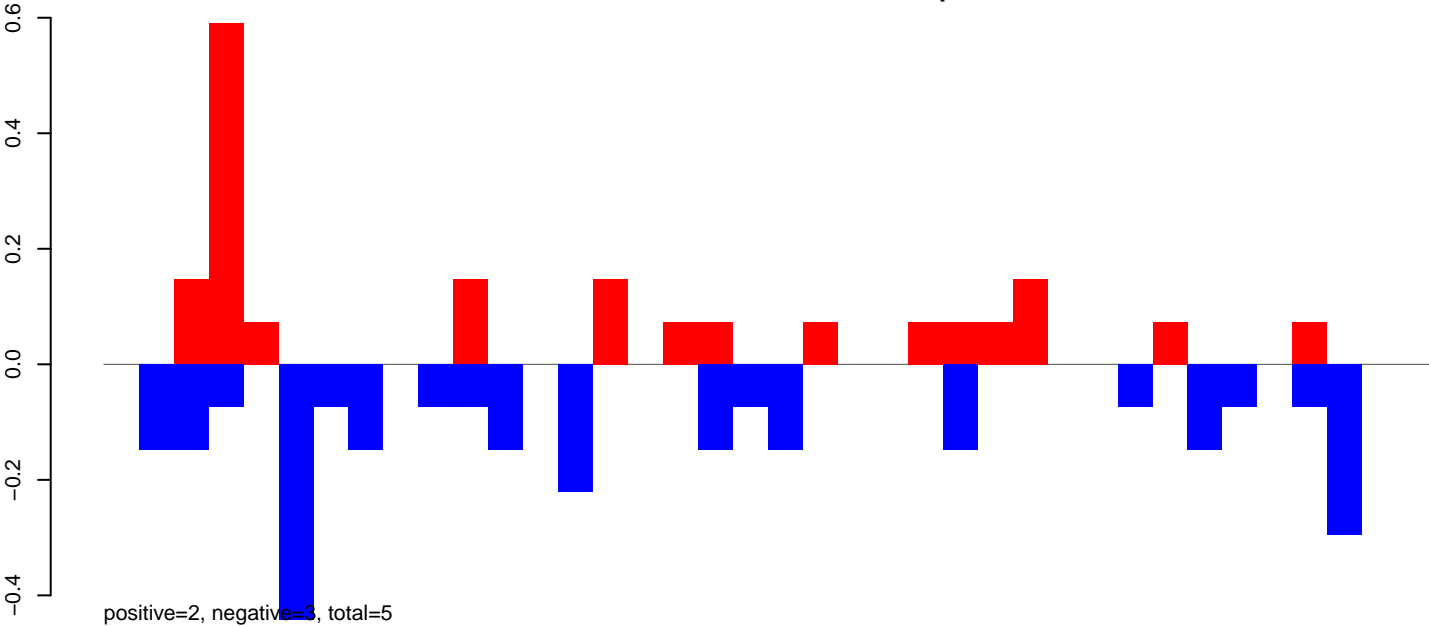
Window size=25, length=995, TE@rnd-4_family-681@Unknown@974:1-995

0 200 400 600 800 1000

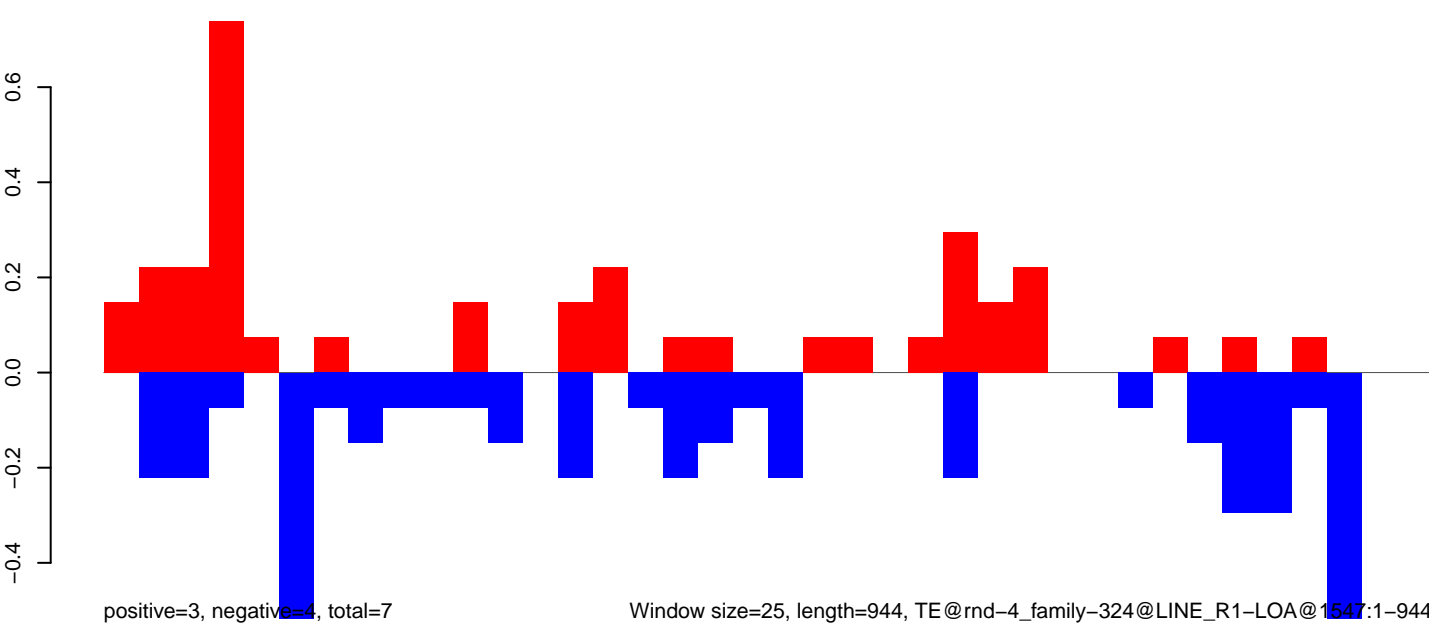
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

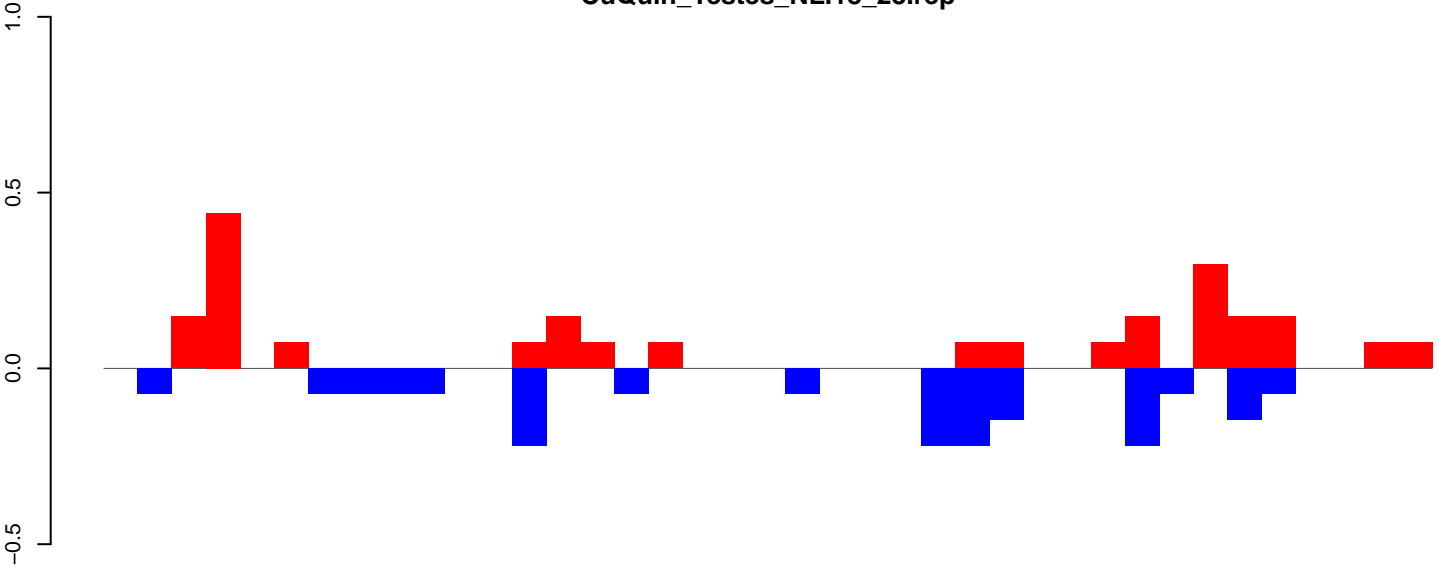


CuQuin_Testes_NL.rep



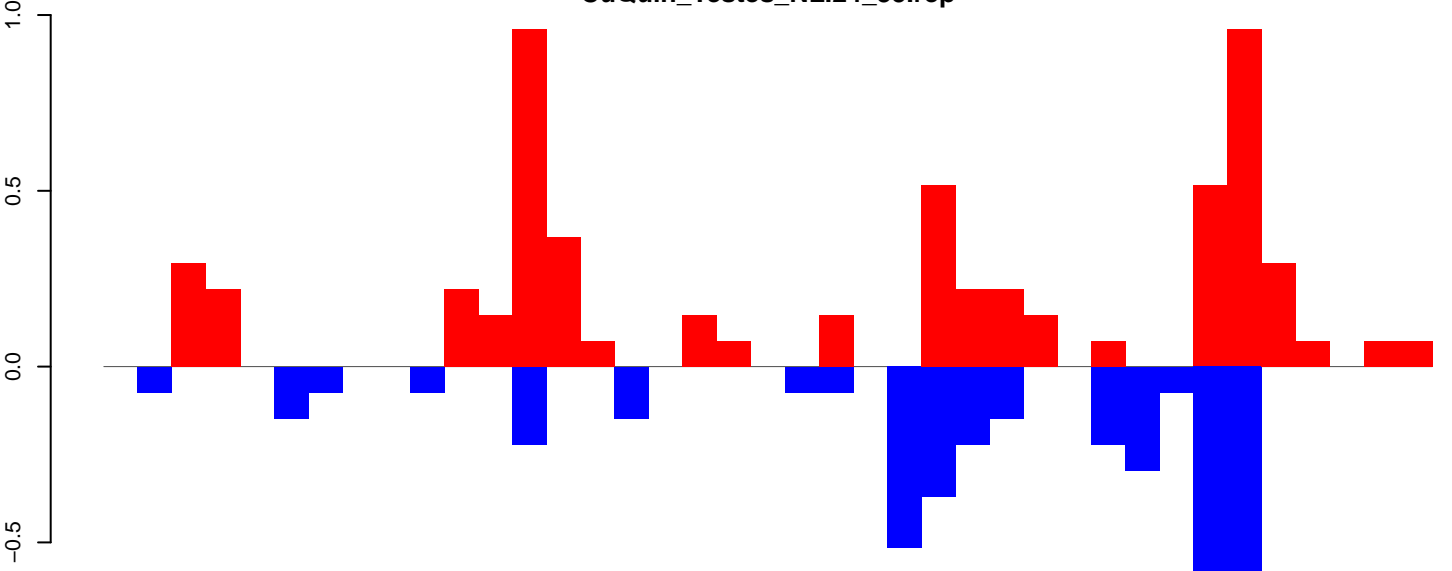
Window size=25, length=944, TE@rnd-4_family-324@LINE_R1-LOA@1547:1-944

CuQuin_Testes_NL.18_23.rep



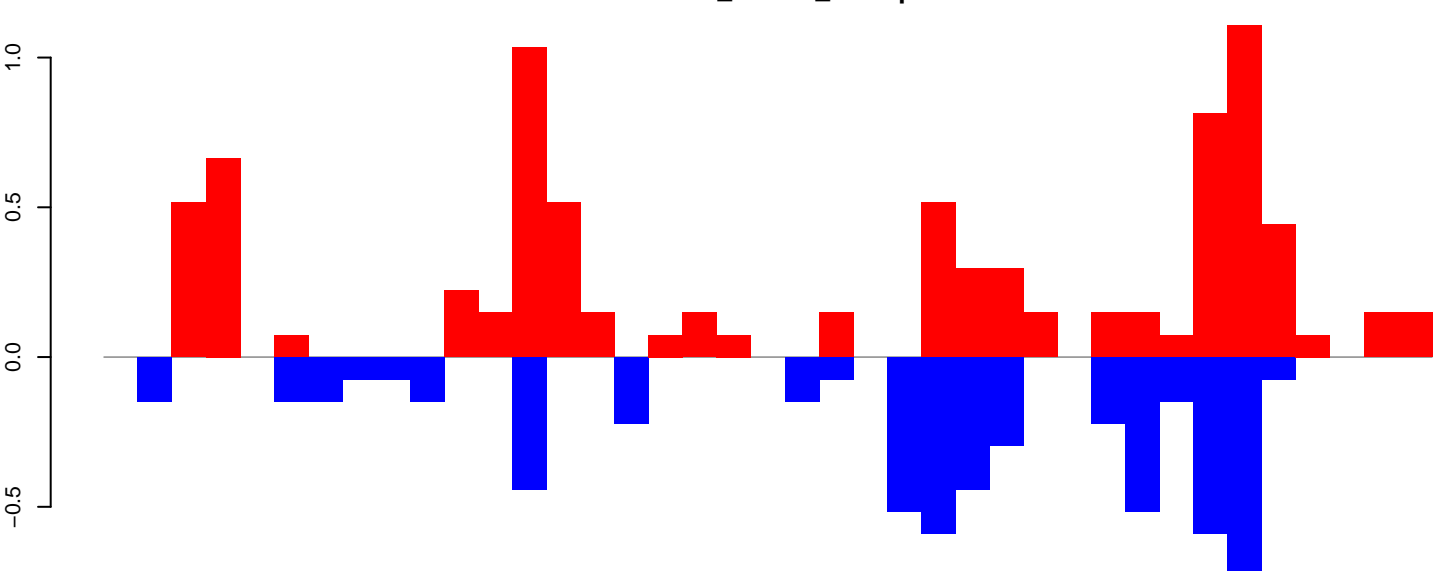
positive=2, negative=2, total=4

CuQuin_Testes_NL.24_35.rep



positive=6, negative=4, total=10

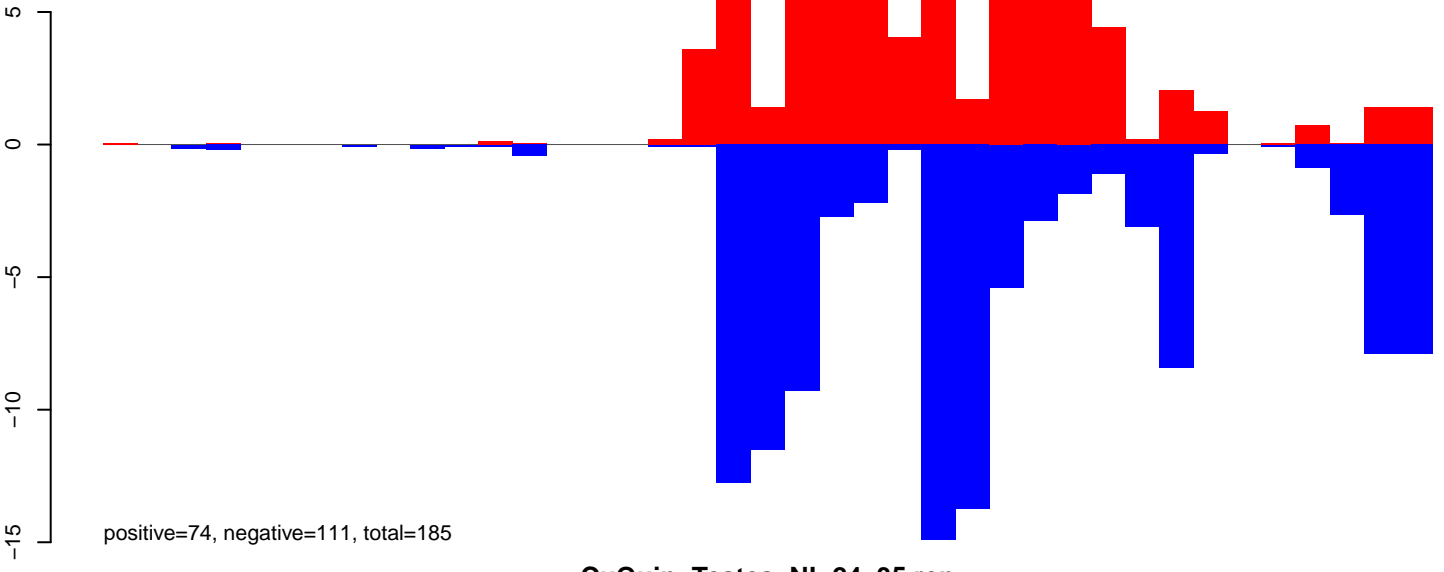
CuQuin_Testes_NL.rep



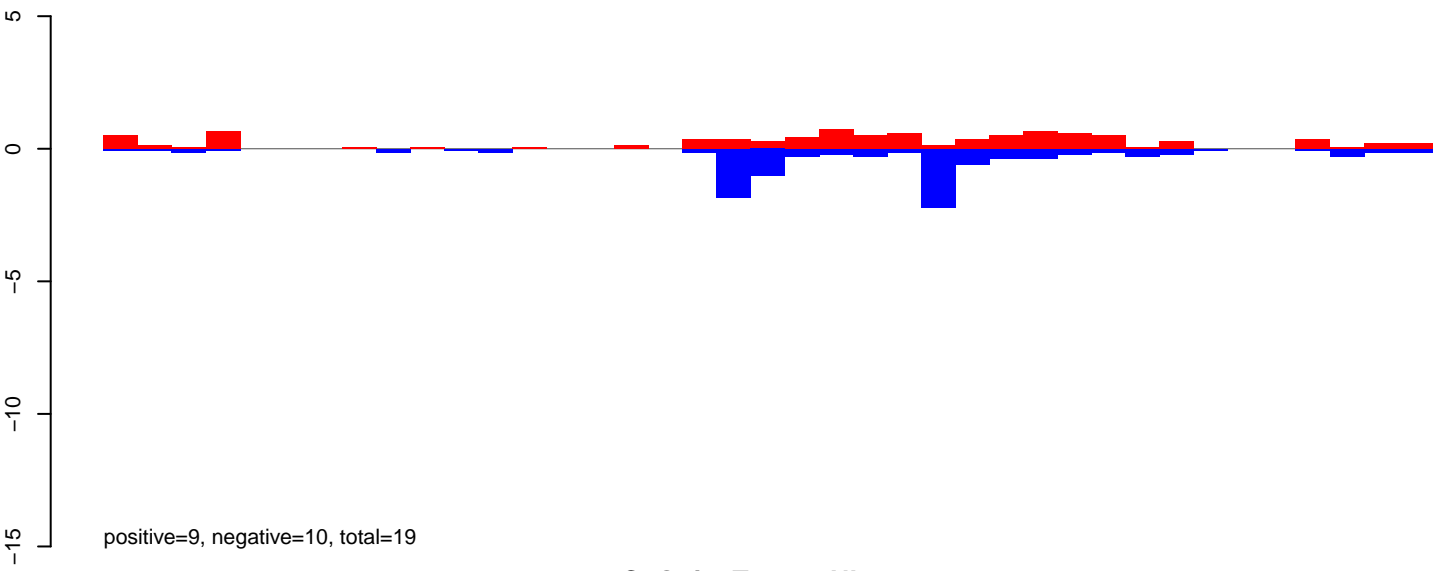
positive=8, negative=6, total=14

Window size=25, length=964, TE@rnd-4_family-220@Unknown@1567:1-964

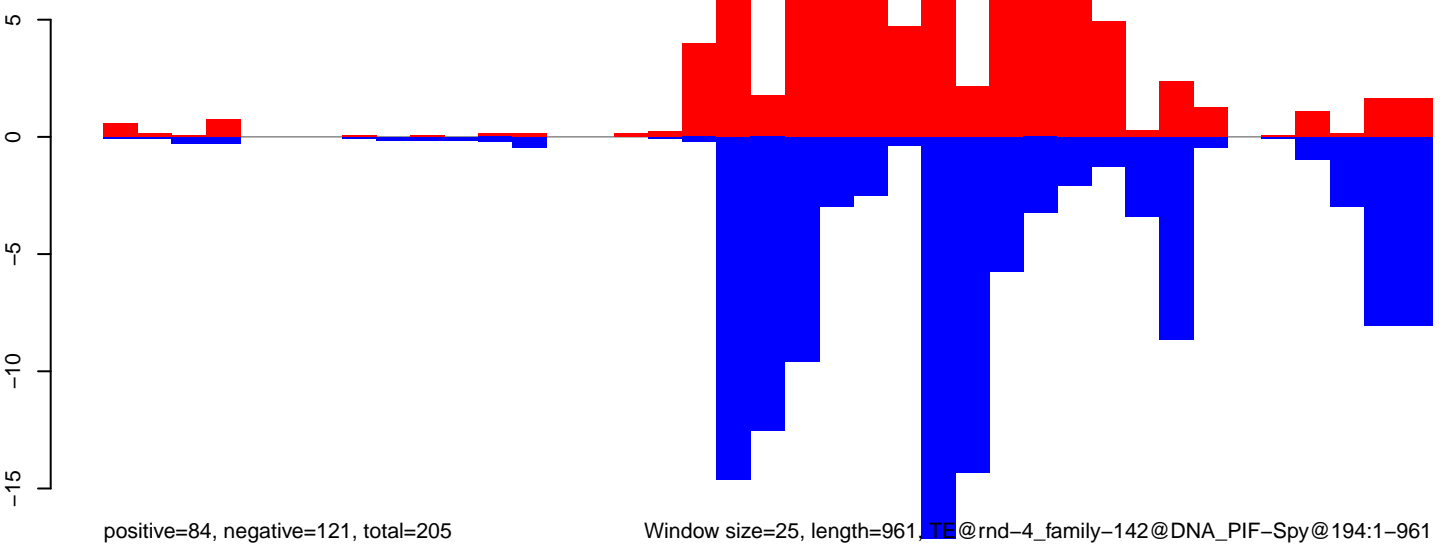
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



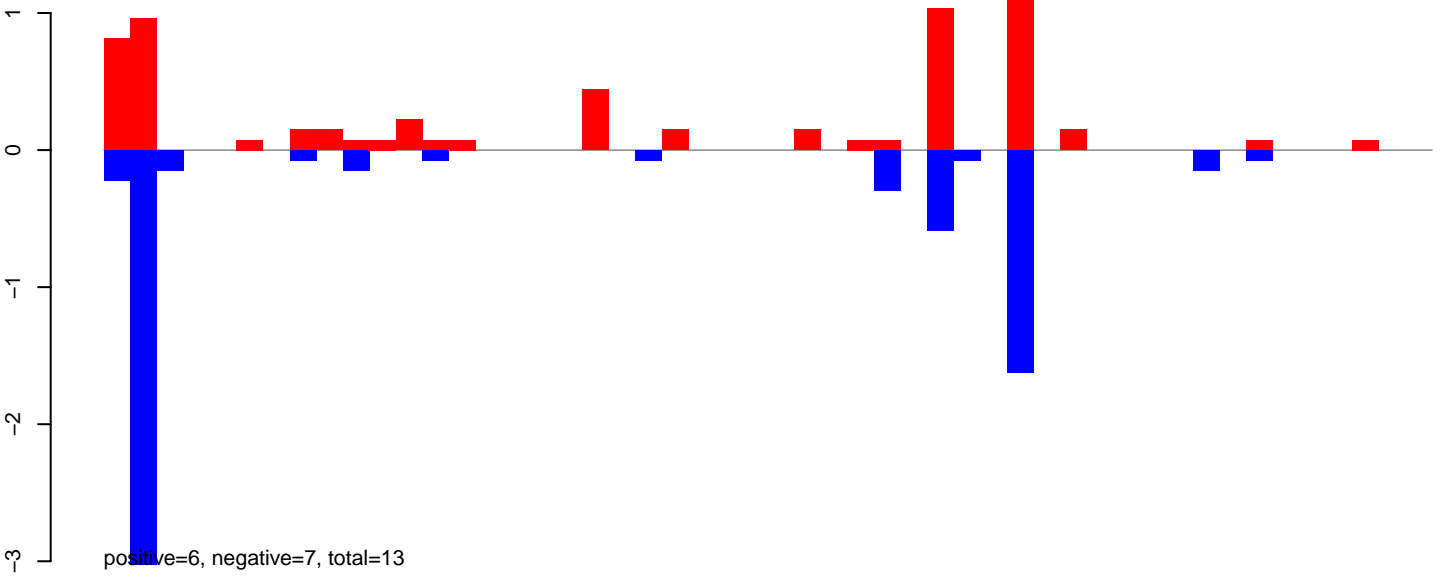
CuQuin_Testes_NL.rep



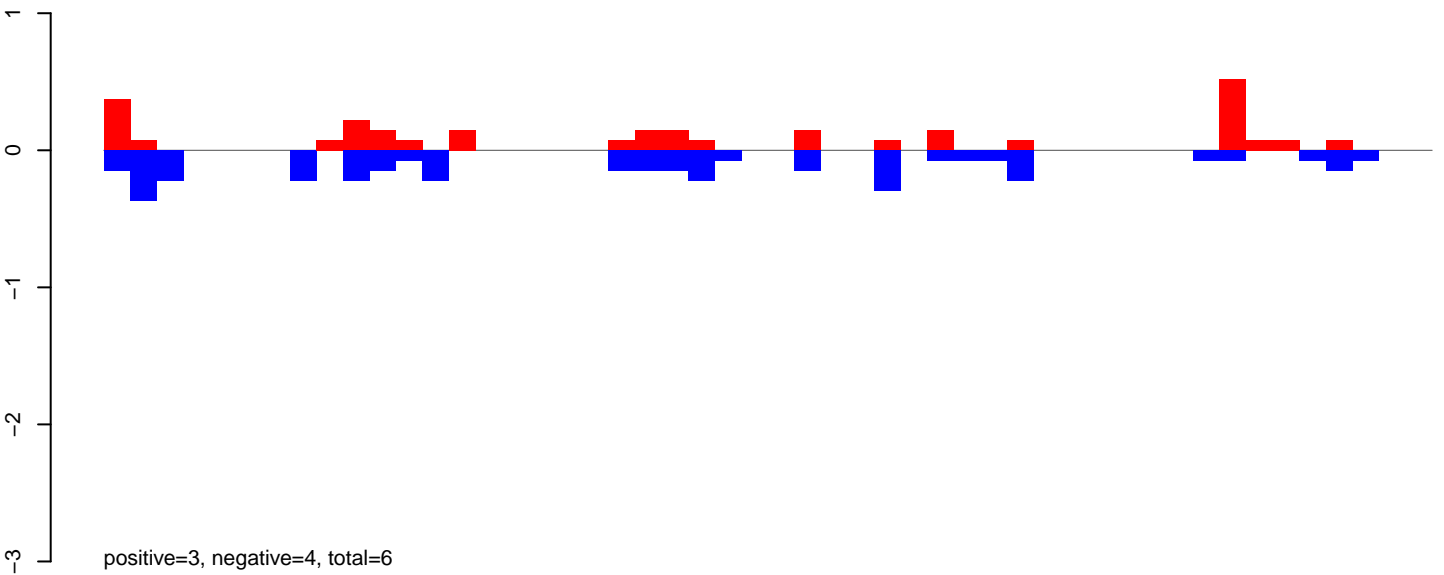
Window size=25, length=961, TE@rnd-4_family-142@DNA_PIF-Spy@194:1-961

0 200 400 600 800 1000

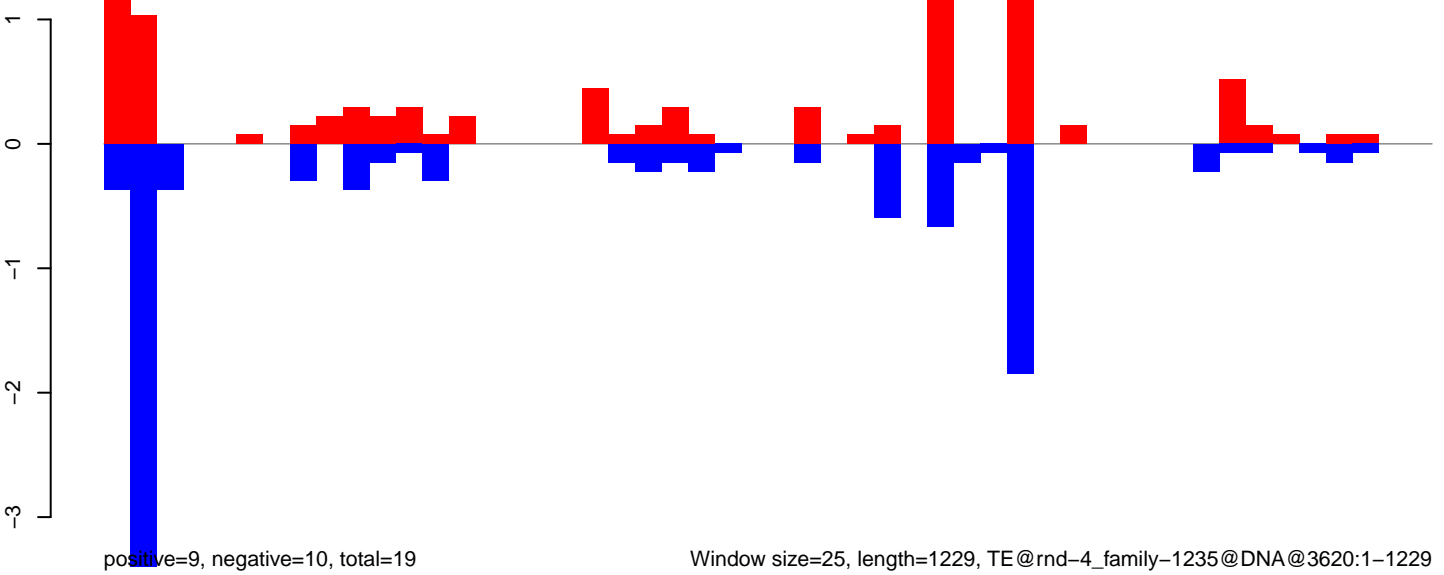
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



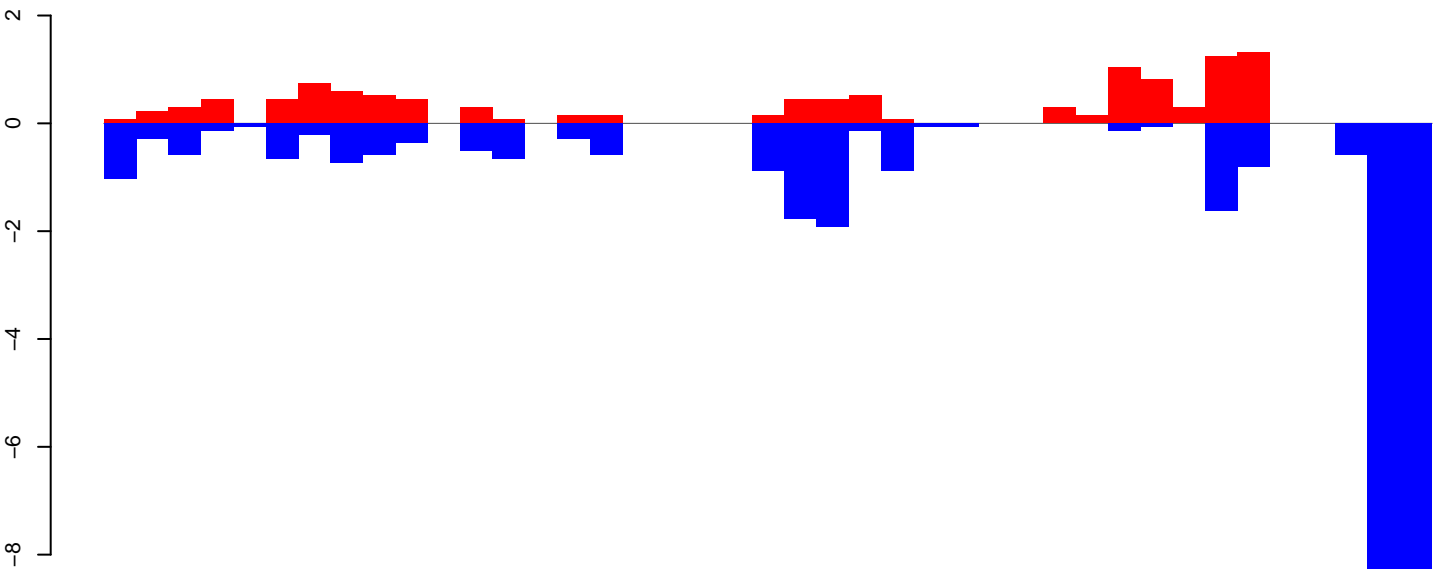
CuQuin_Testes_NL.rep



Window size=25, length=1229, TE@rnd-4_family-1235@DNA@3620:1-1229

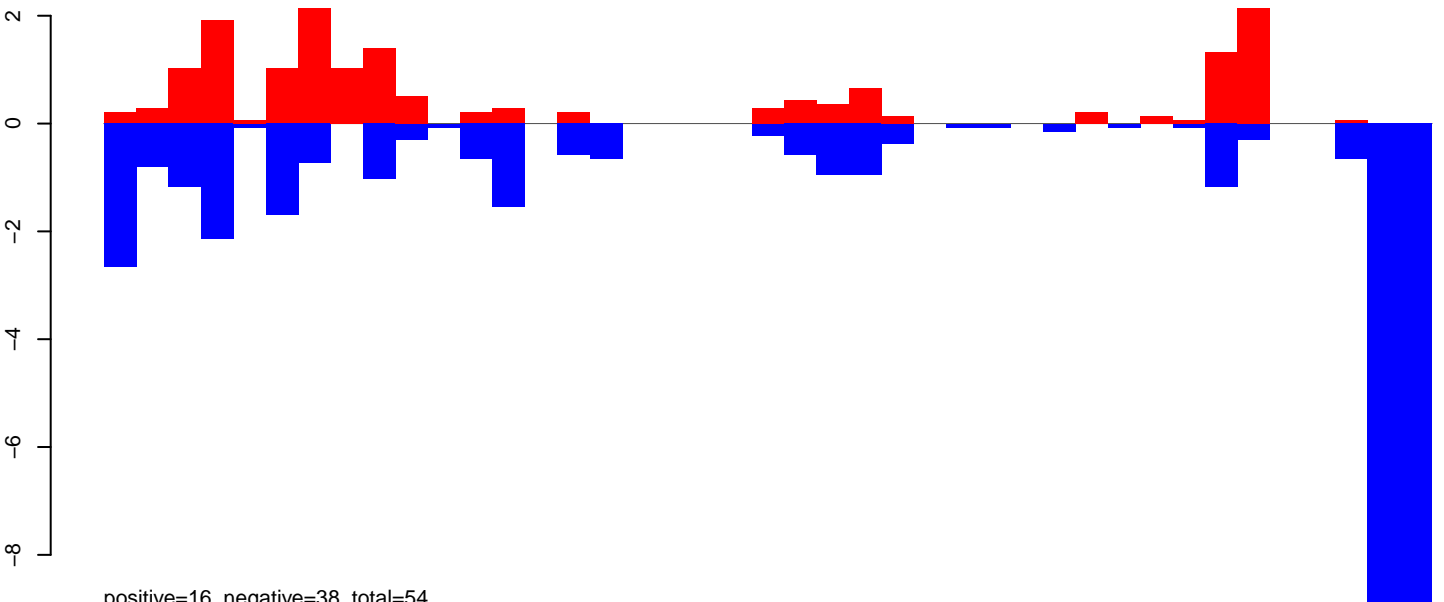
0 200 400 600 800 1000 1200

CuQuin_Testes_NL.18_23.rep



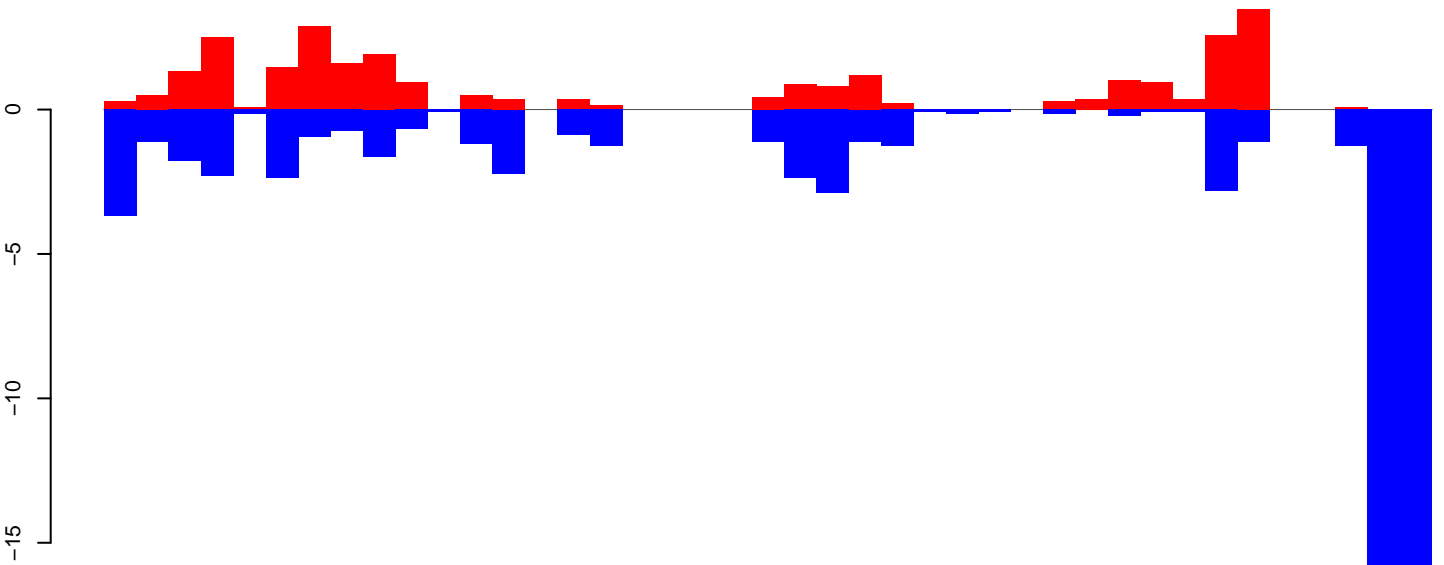
positive=11, negative=32, total=44

CuQuin_Testes_NL.24_35.rep



positive=16, negative=38, total=54

CuQuin_Testes_NL.rep

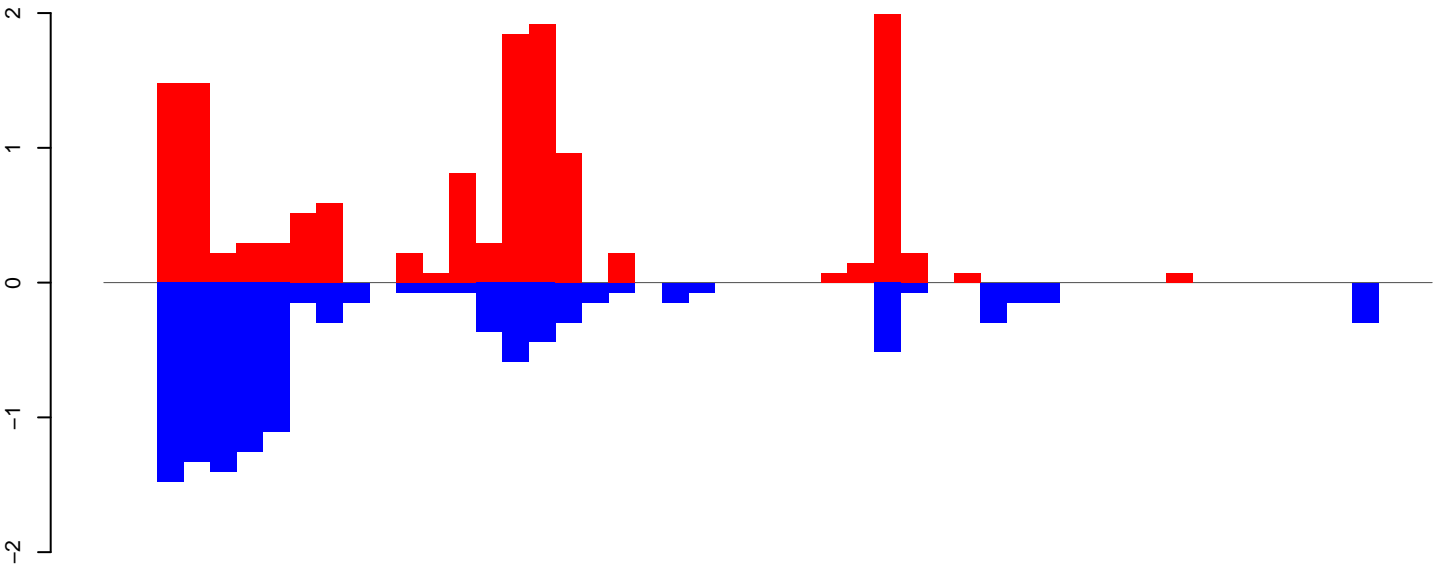


positive=28, negative=70, total=98

Window size=25, length=1019, TE@rnd-3_family-99@DNA_Crypton-I@3266:1-1019

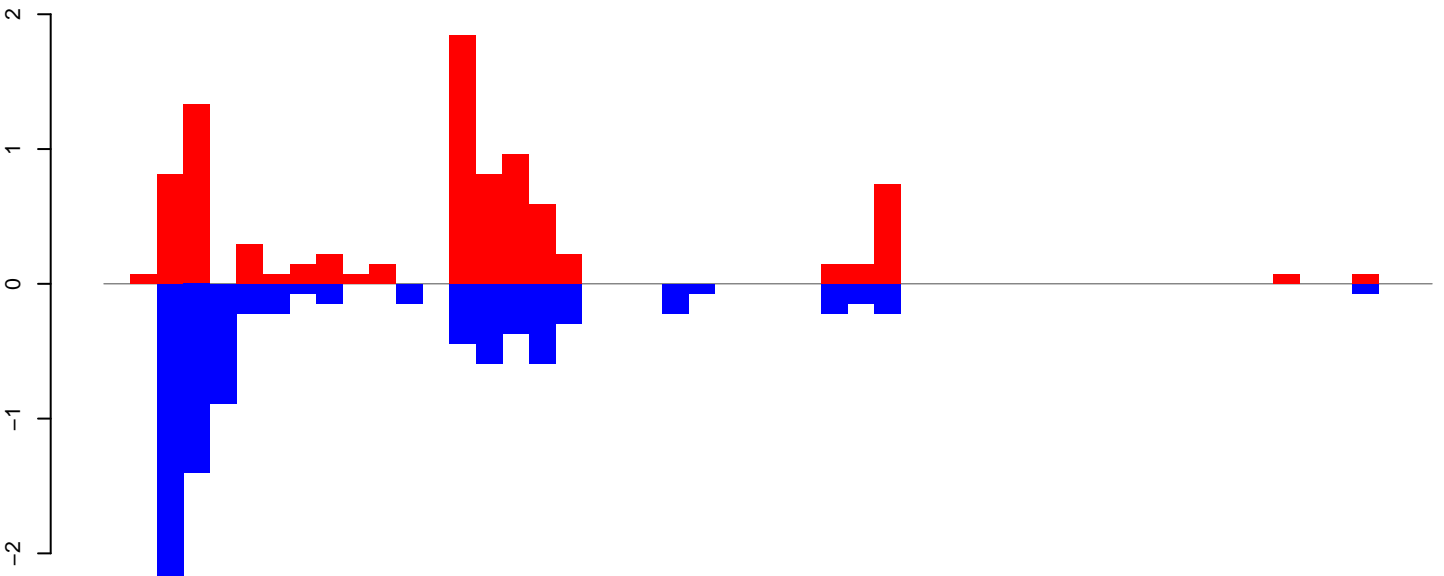
0 200 400 600 800 1000

CuQuin_Testes_NL.18_23.rep



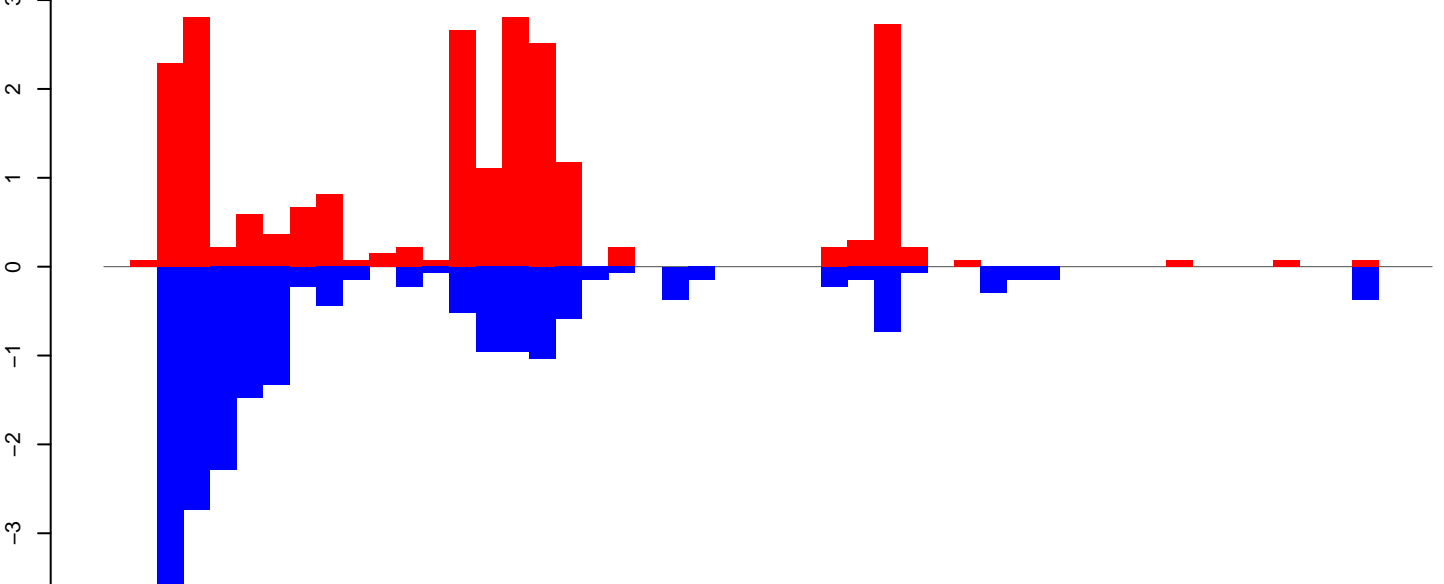
positive=14, negative=11, total=25

CuQuin_Testes_NL.24_35.rep



positive=9, negative=9, total=18

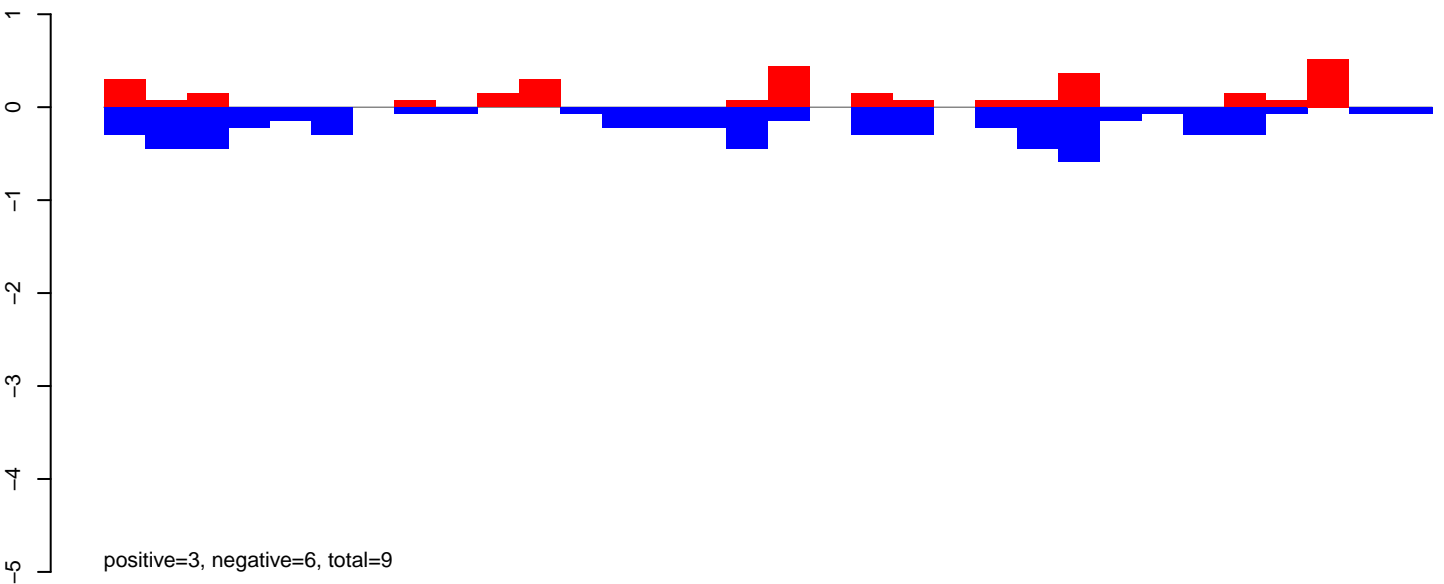
CuQuin_Testes_NL.rep



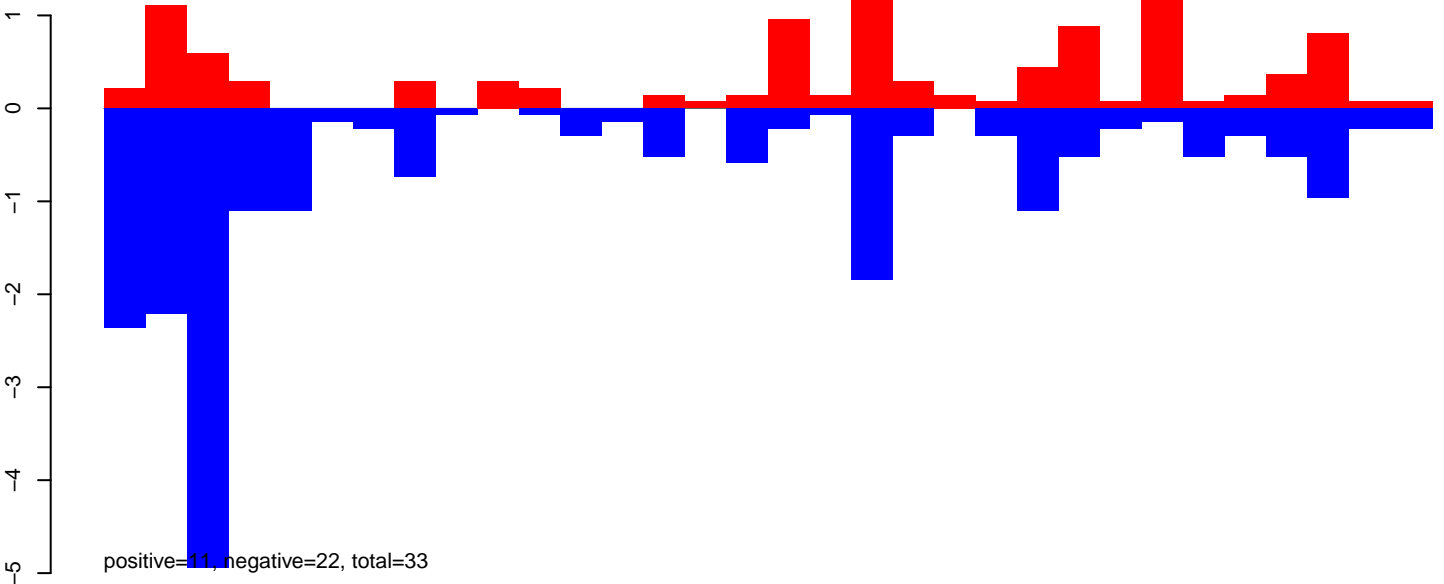
positive=23, negative=20, total=42

Window size=25, length=1230, TE@rnd-3_family-28@DNA_Zator@1289:1-1230

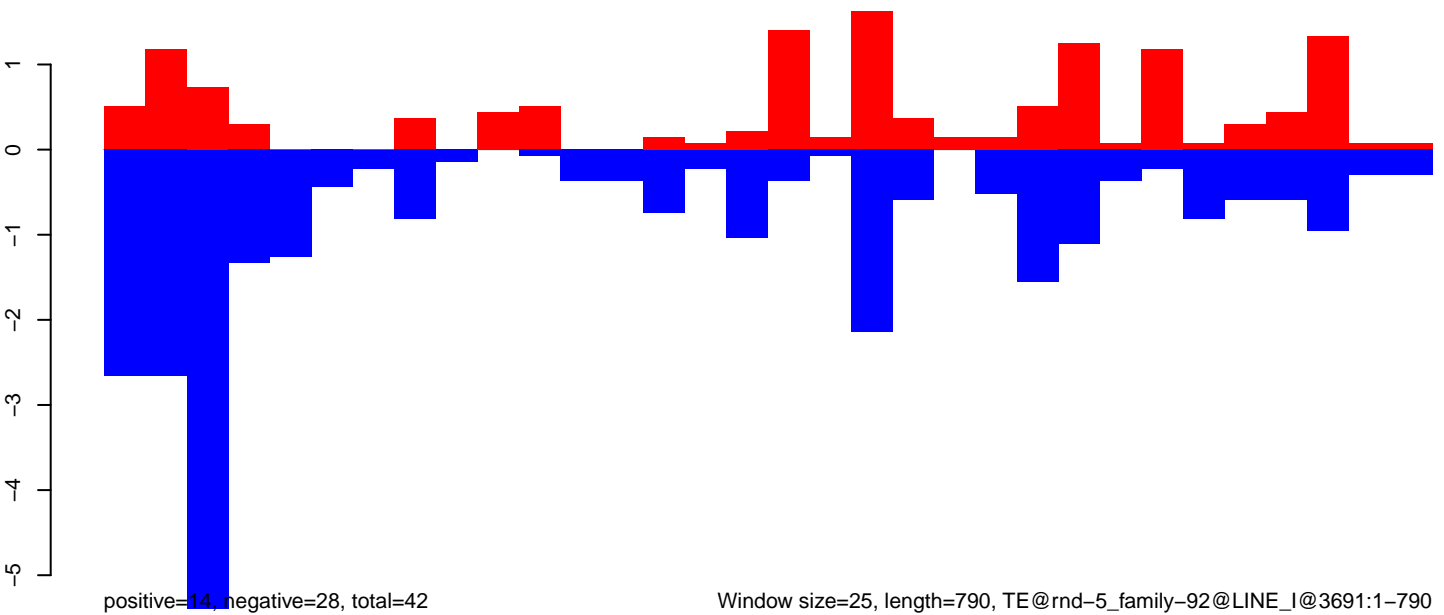
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

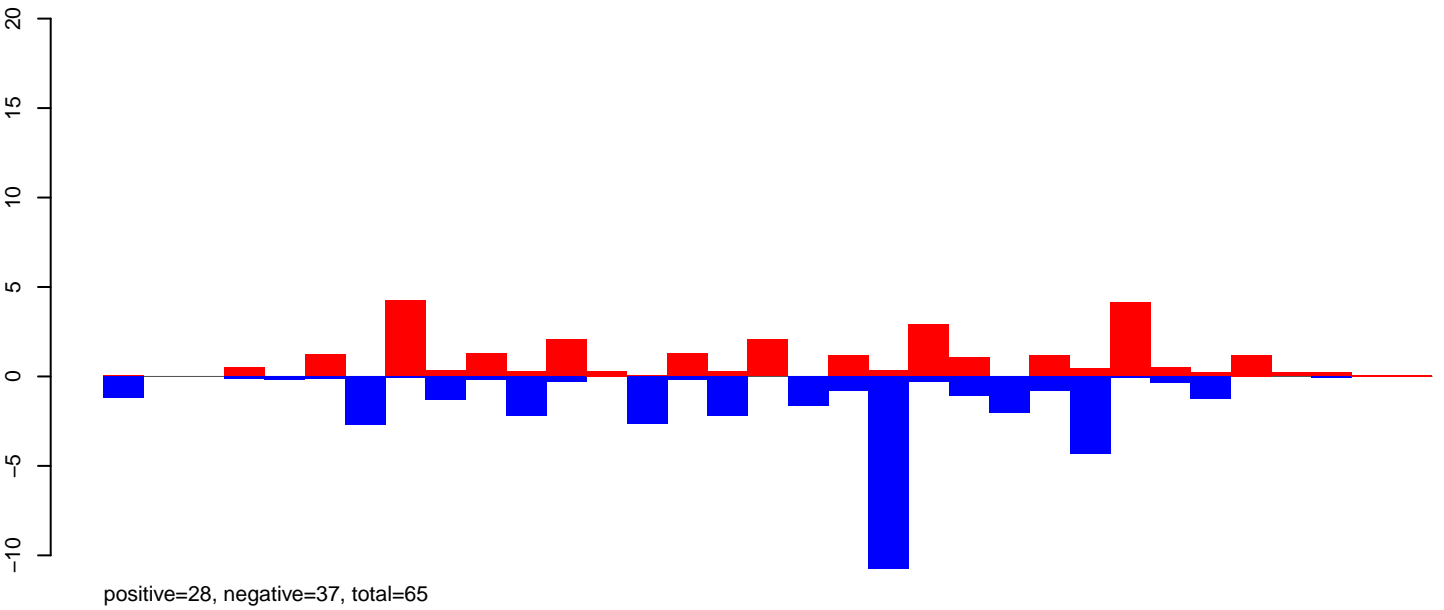


CuQuin_Testes_NL.rep

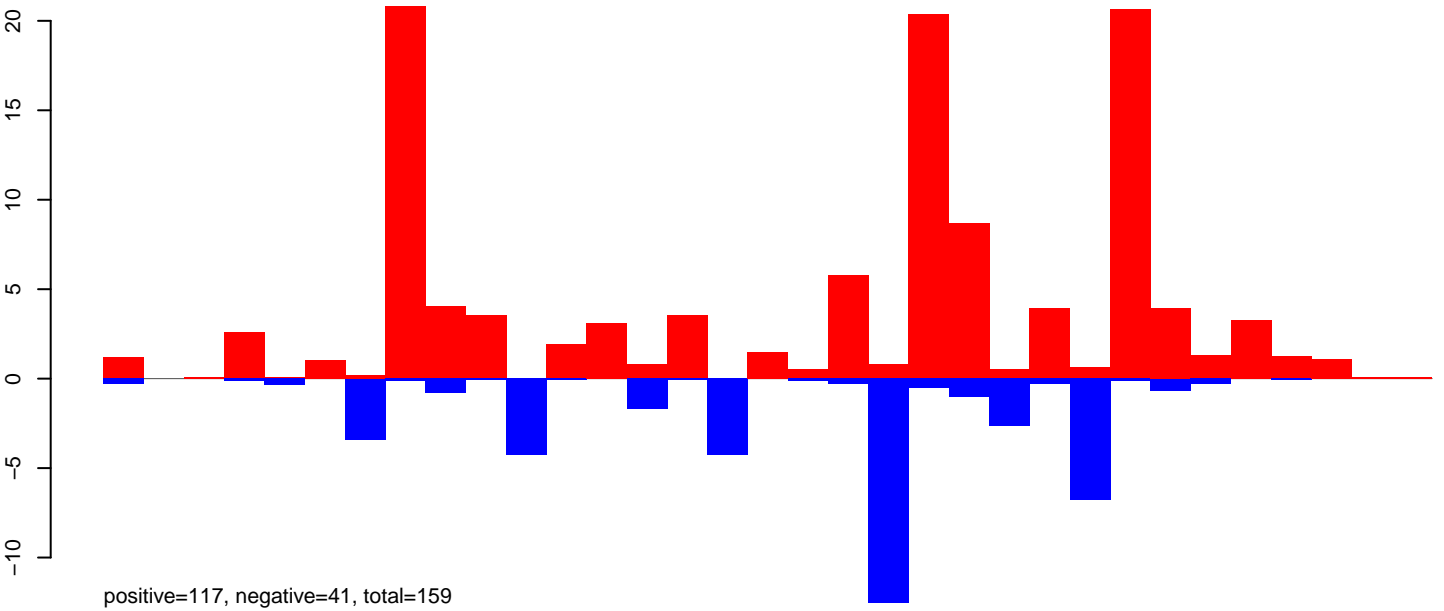


Window size=25, length=790, TE@rnd-5_family-92@LINE_I@3691:1-790

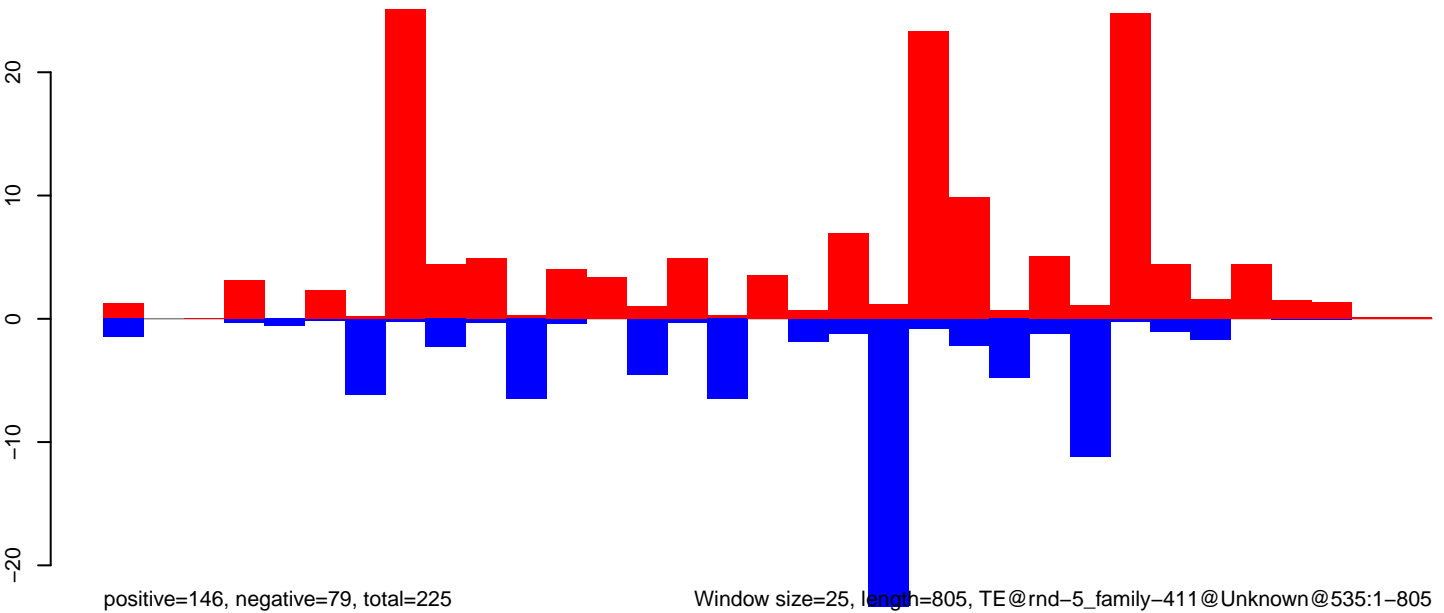
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

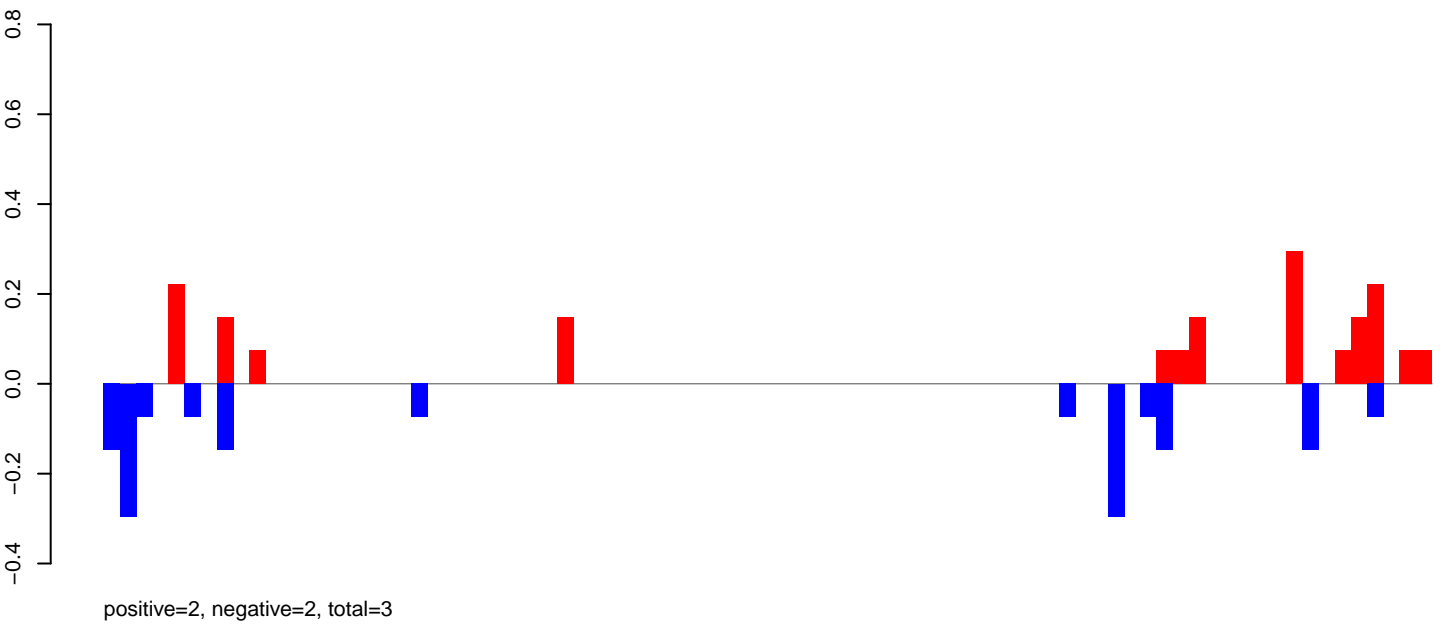


CuQuin_Testes_NL.rep

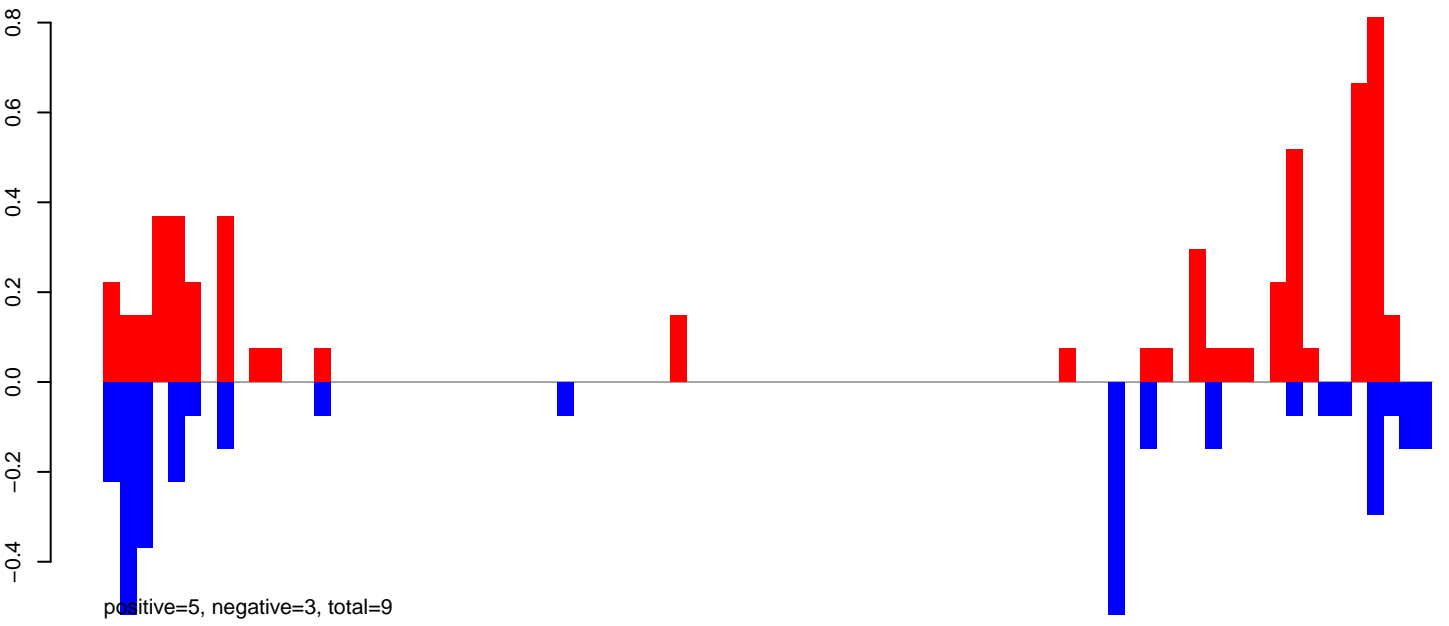


Window size=25, length=805, TE@rnd-5_family-411@Unknown@535:1-805

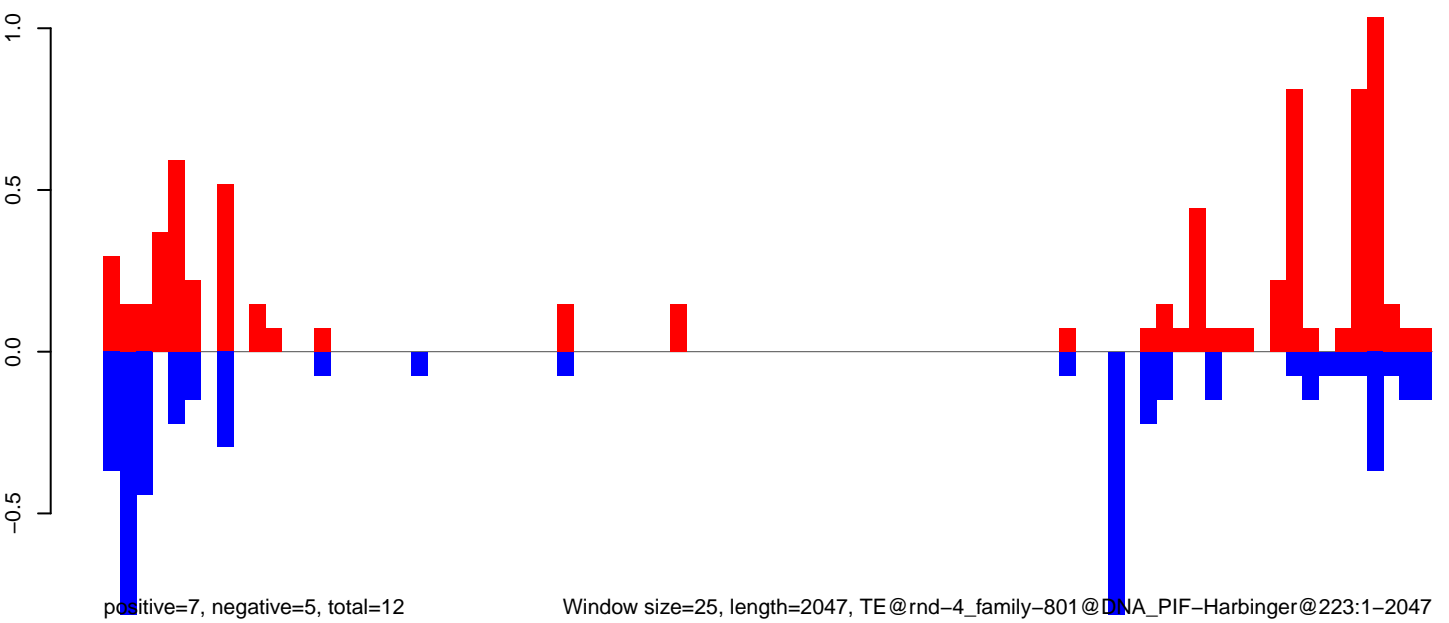
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

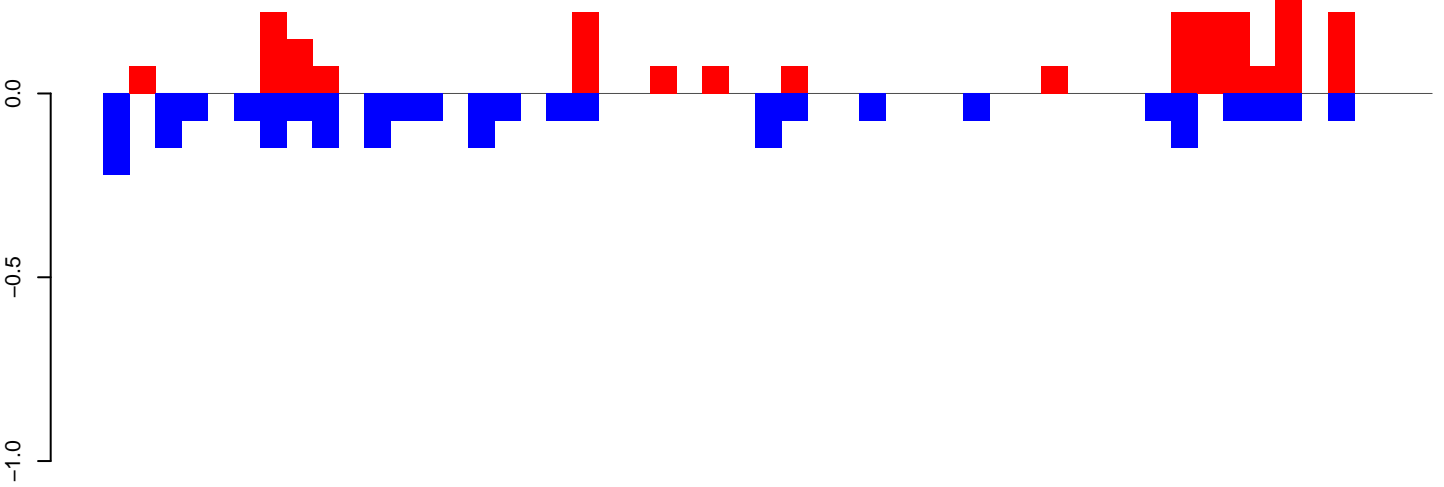


CuQuin_Testes_NL.rep



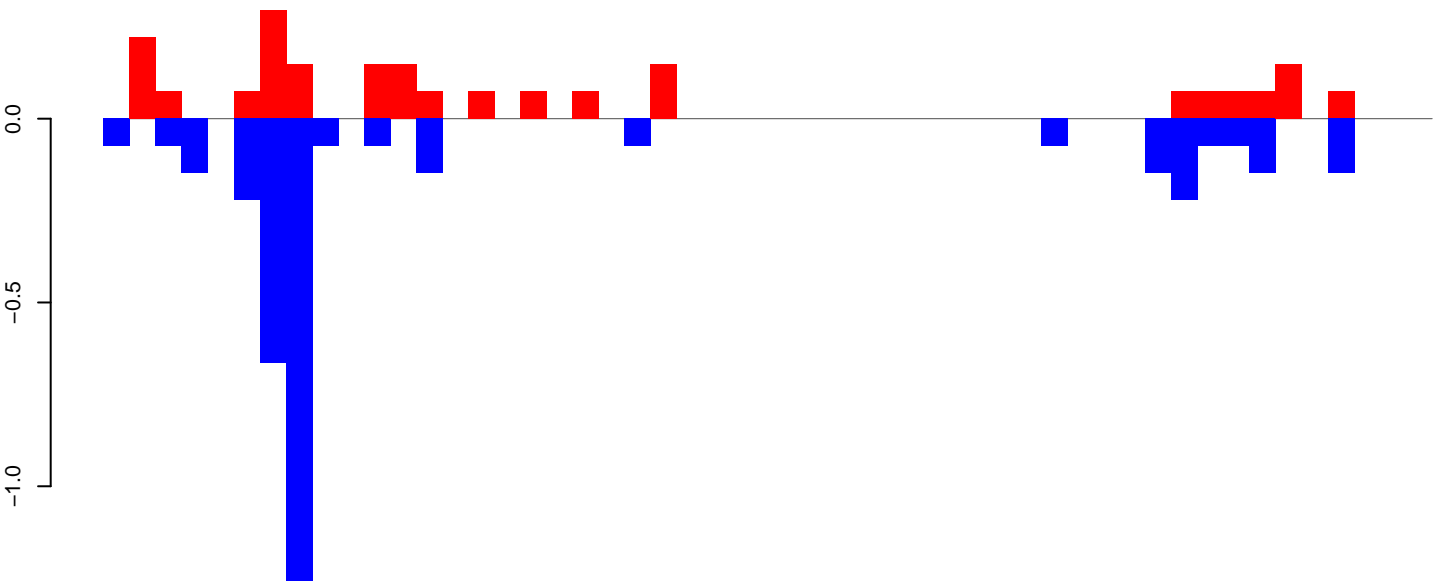
Window size=25, length=2047, TE@rnd-4_family-801@DNA_PIF-Harbinger@223:1-2047

CuQuin_Testes_NL.18_23.rep



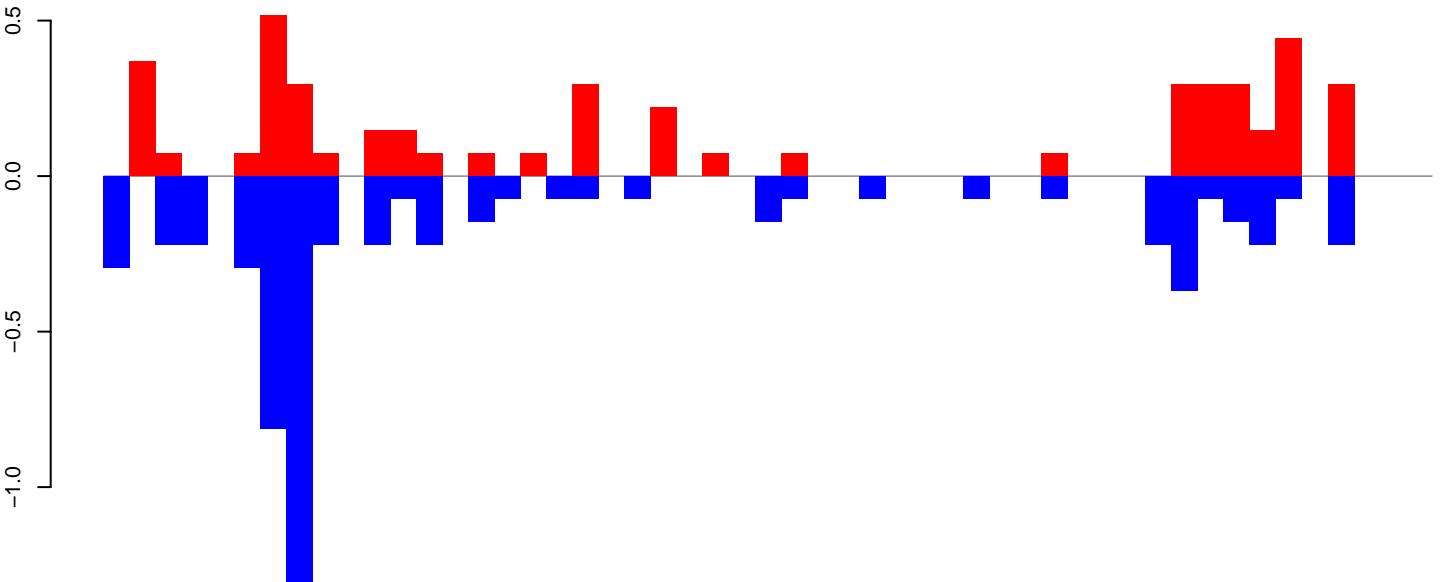
positive=2, negative=2, total=5

CuQuin_Testes_NL.24_35.rep



positive=2, negative=4, total=6

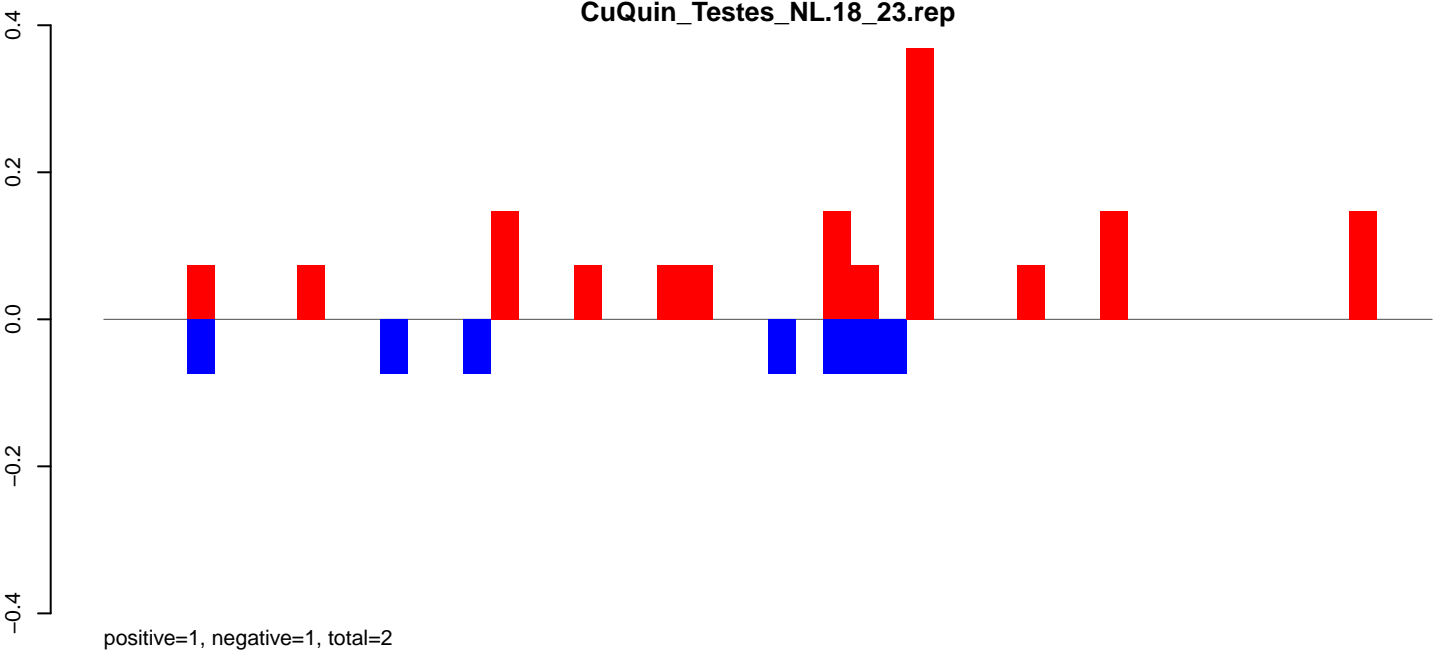
CuQuin_Testes_NL.rep



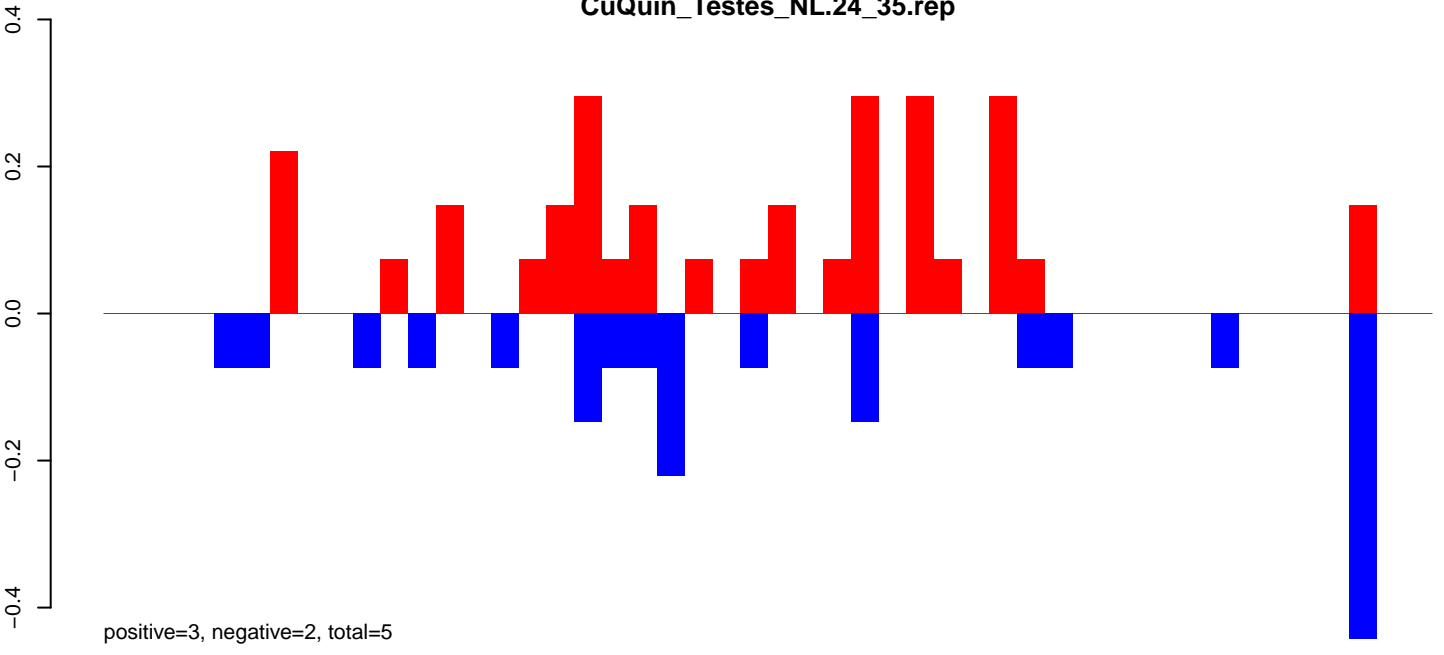
positive=4, negative=6, total=11

Window size=25, length=1252, TE@rnd-4_family-710@DNA@3450:1-1252

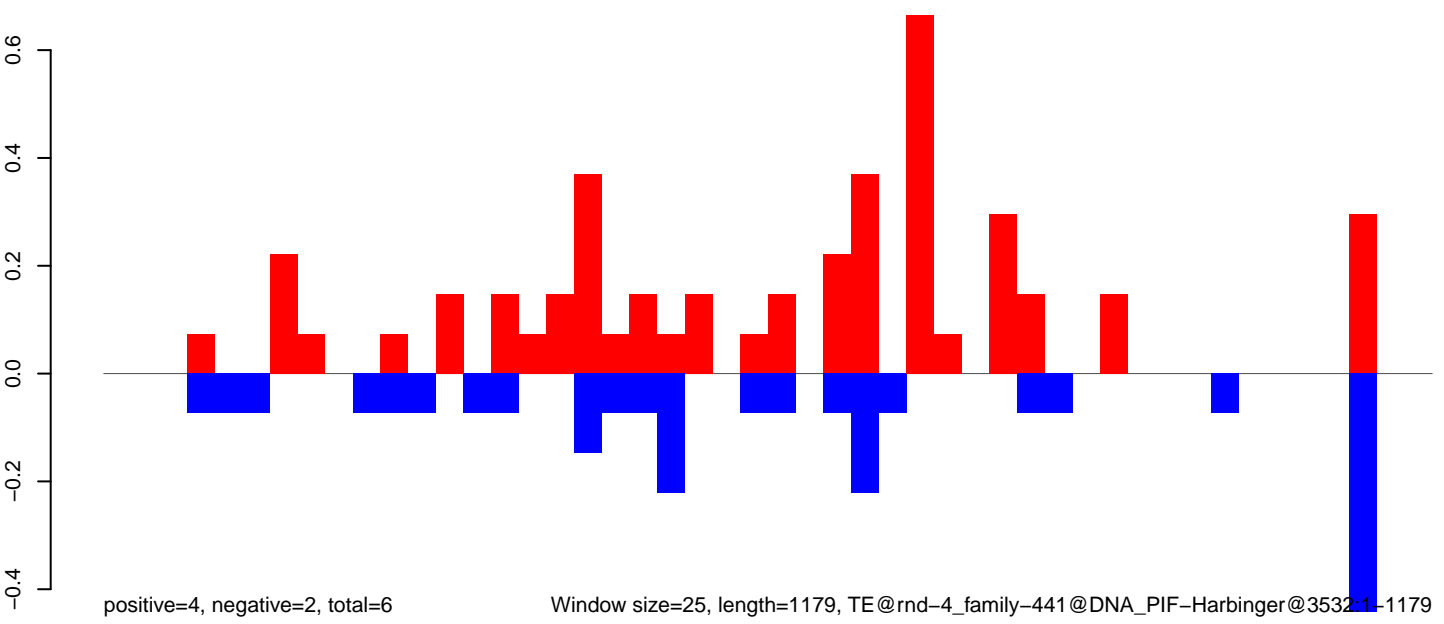
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



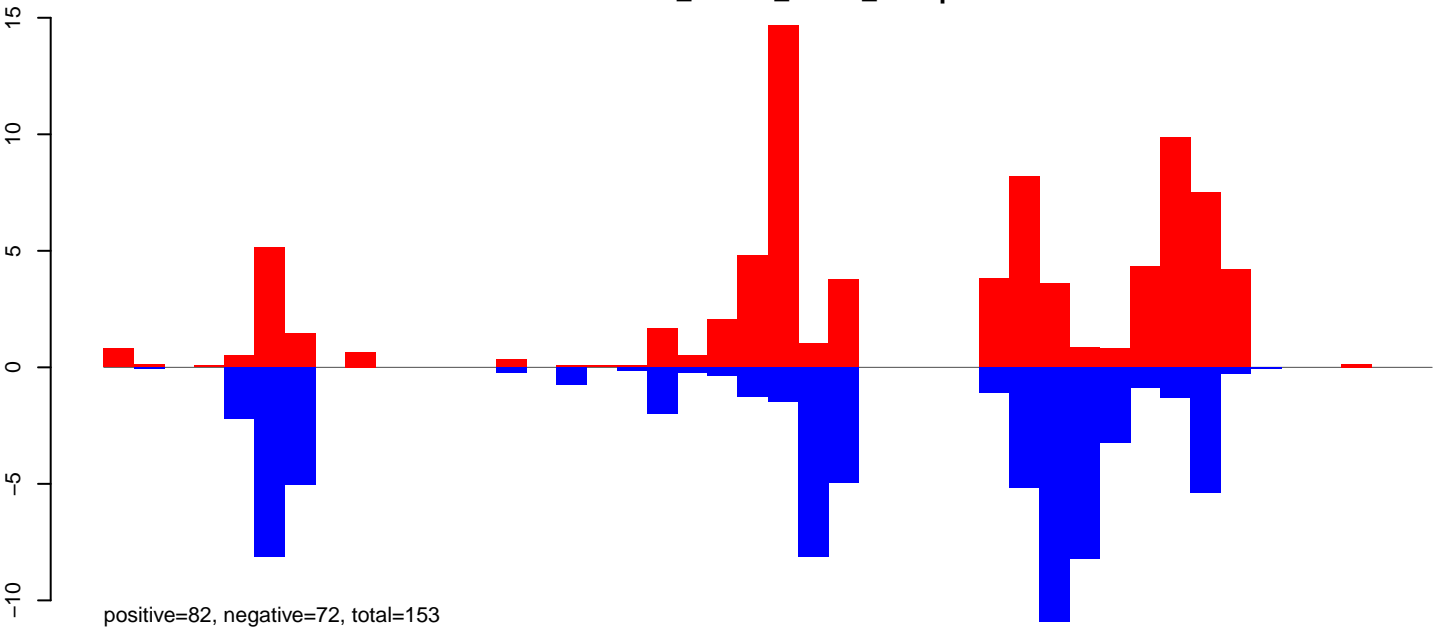
CuQuin_Testes_NL.rep



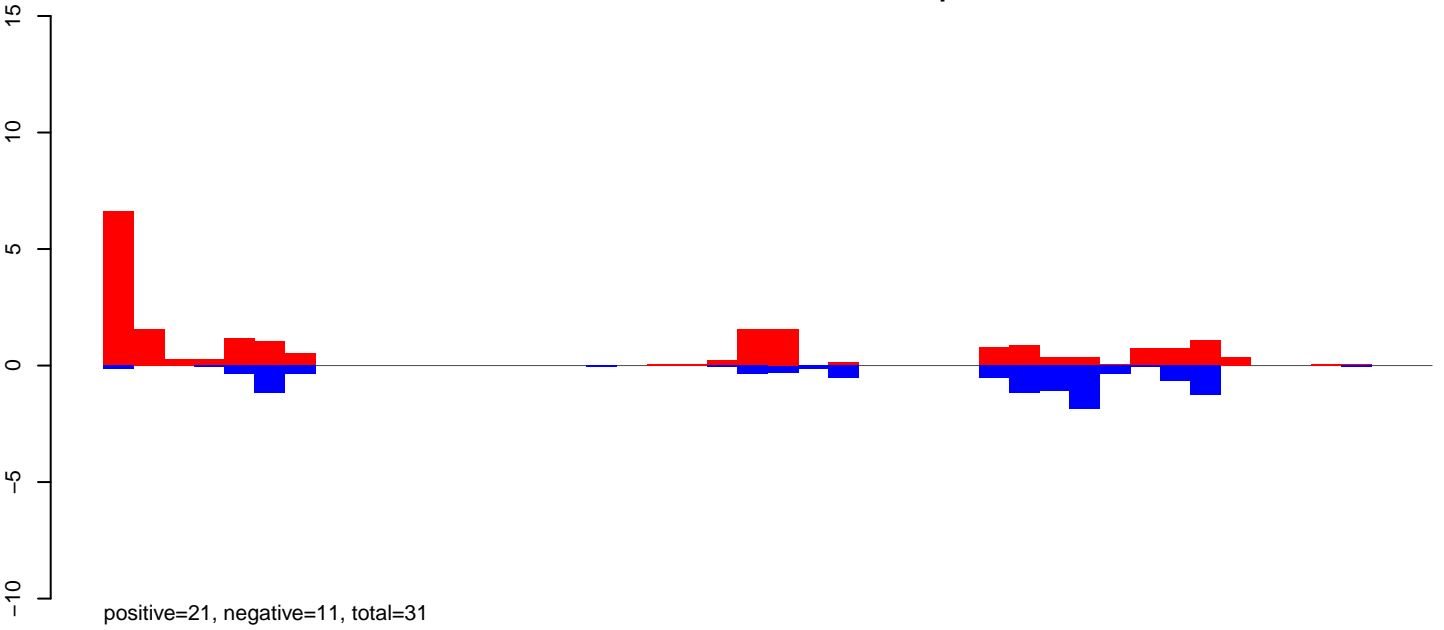
Window size=25, length=1179, TE@rnd-4_family-441@DNA_PIF-Harbinger@35321-1179

0 200 400 600 800 1000 1200

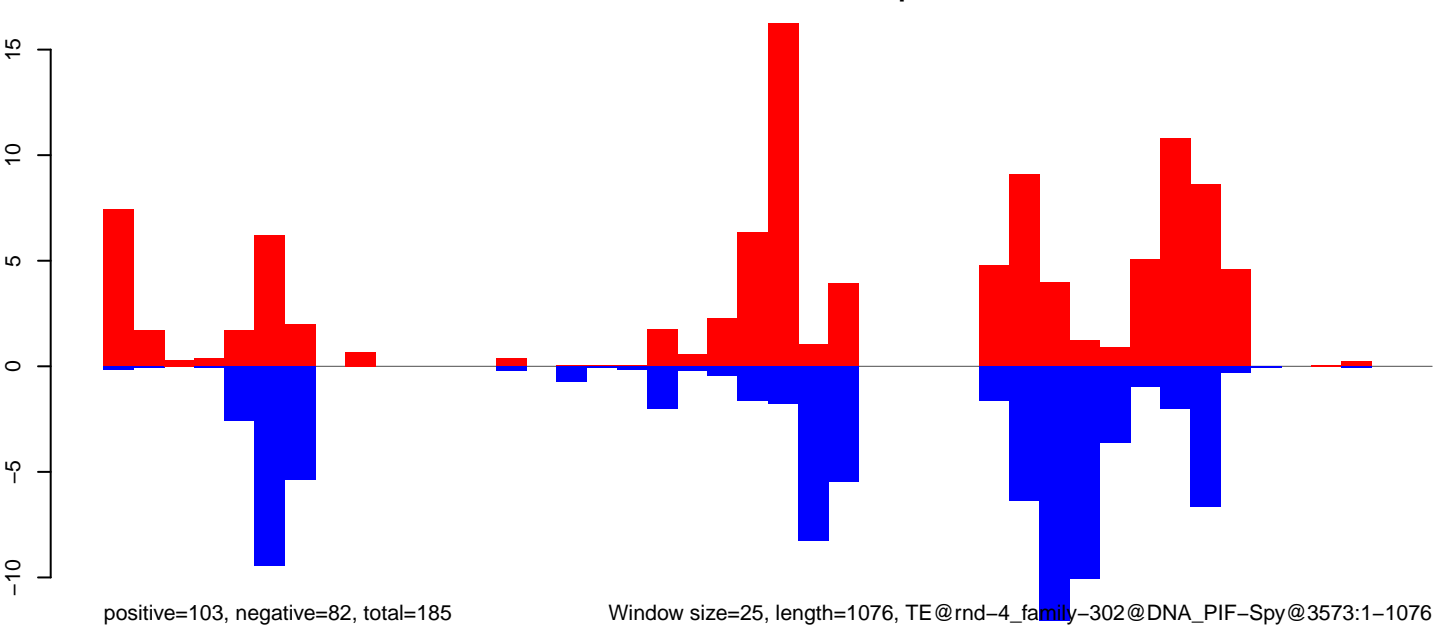
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

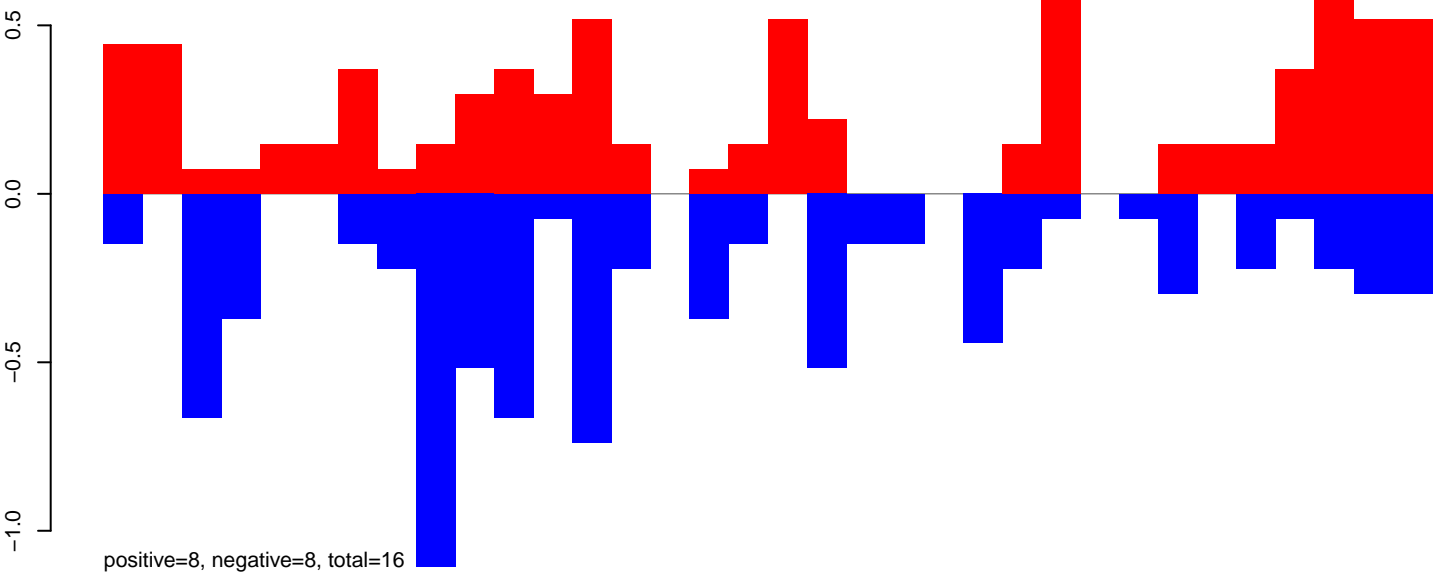


CuQuin_Testes_NL.rep

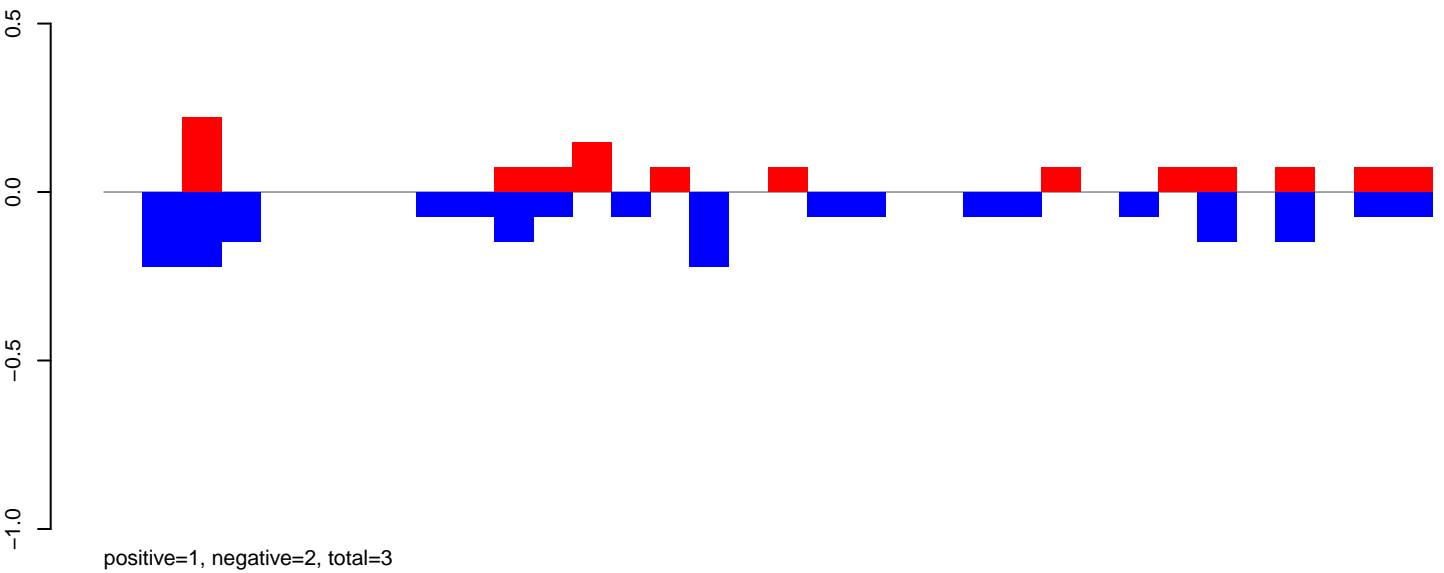


Window size=25, length=1076, TE@rnd-4_family-302@DNA_PIF-Spy@3573:1-1076

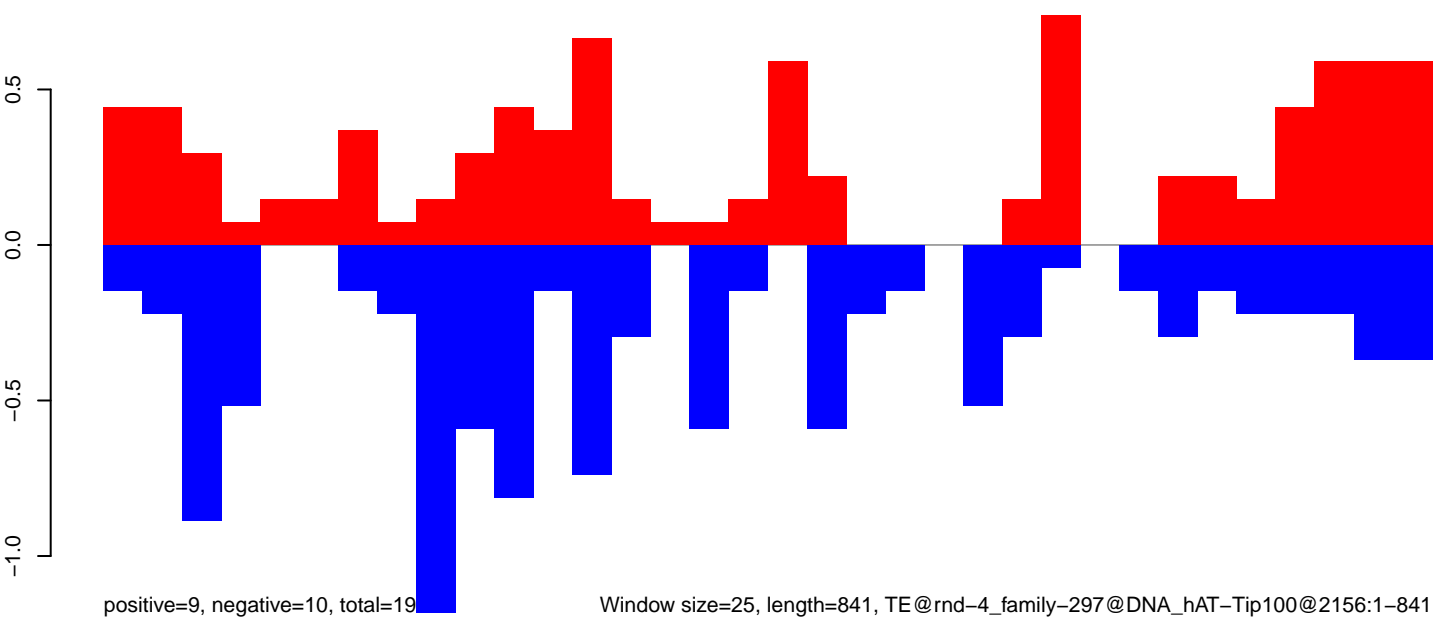
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

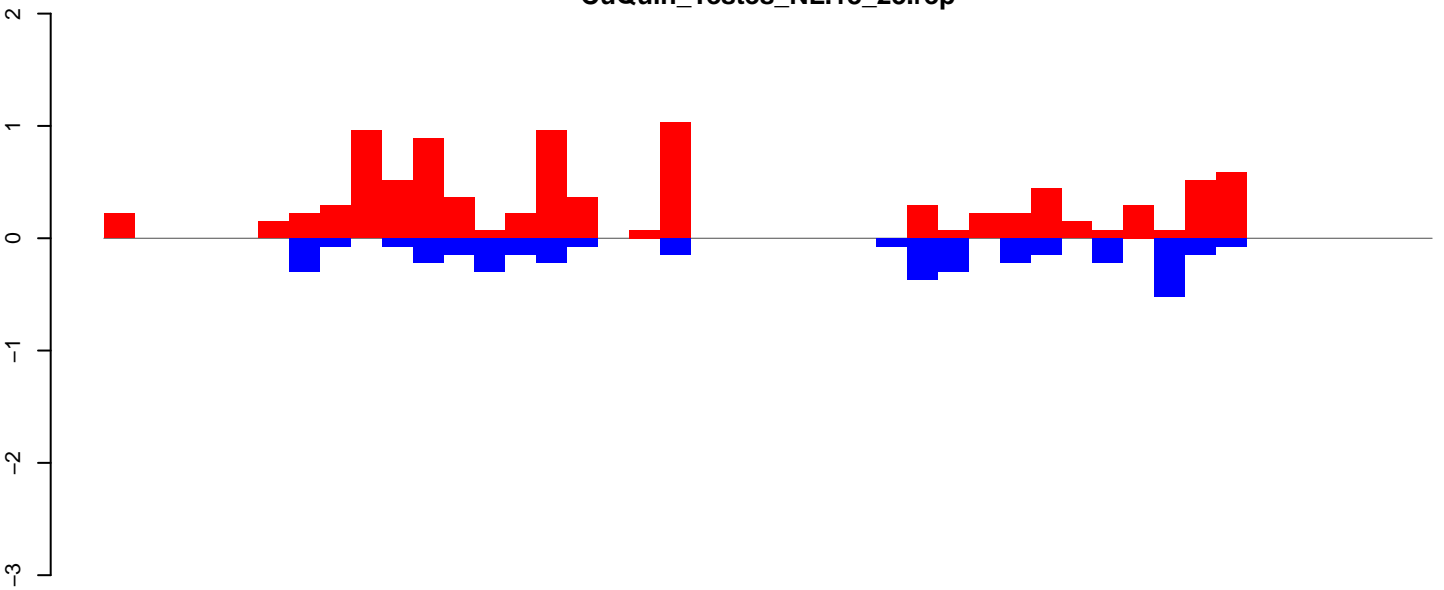


CuQuin_Testes_NL.rep



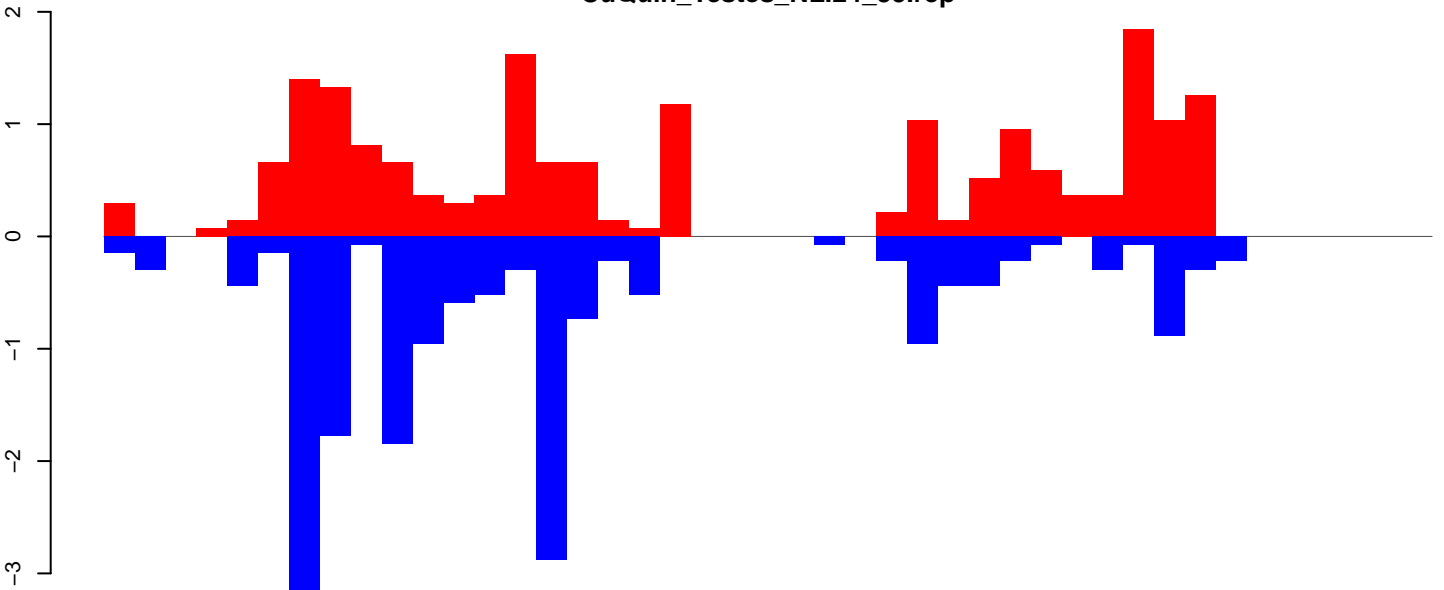
Window size=25, length=841, TE@rnd-4_family-297@DNA_hAT-Tip100@2156:1-841

CuQuin_Testes_NL.18_23.rep



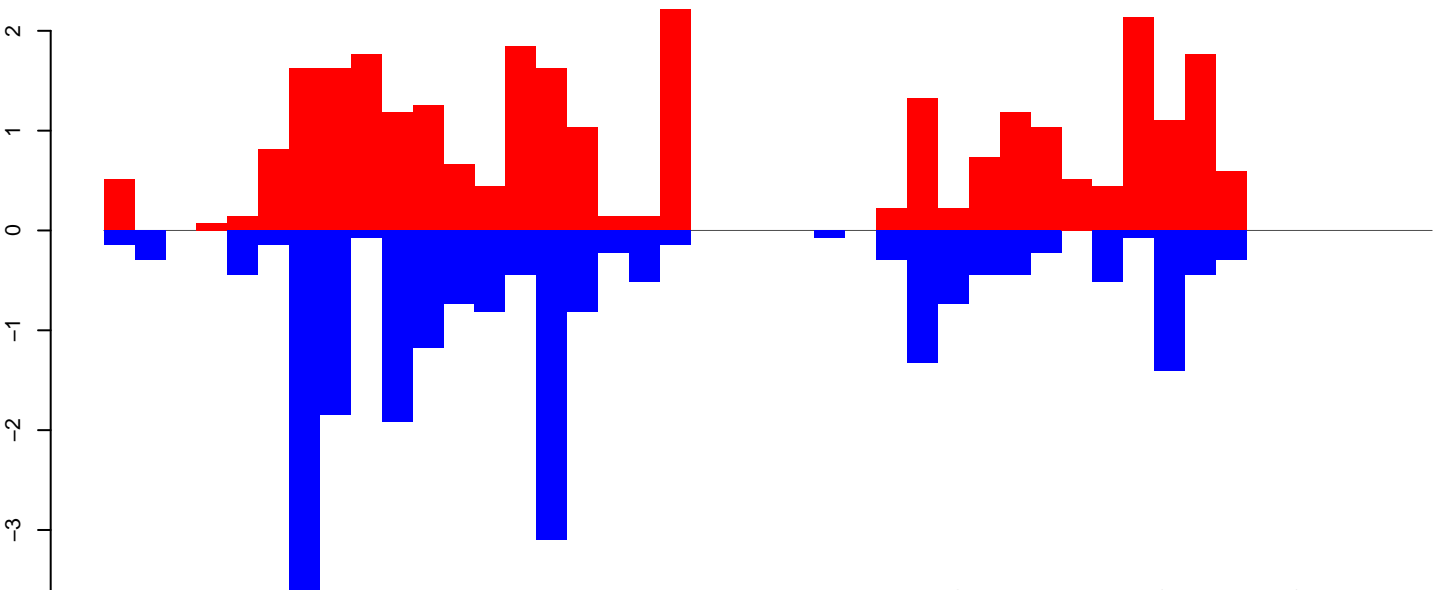
positive=9, negative=4, total=13

CuQuin_Testes_NL.24_35.rep



positive=19, negative=19, total=38

CuQuin_Testes_NL.rep



positive=28, negative=23, total=51

Window size=25, length=1062, TE@rnd-4_family-2287@SINE_tRNA@1454:1-1062

CuQuin_Testes_NL.18_23.rep



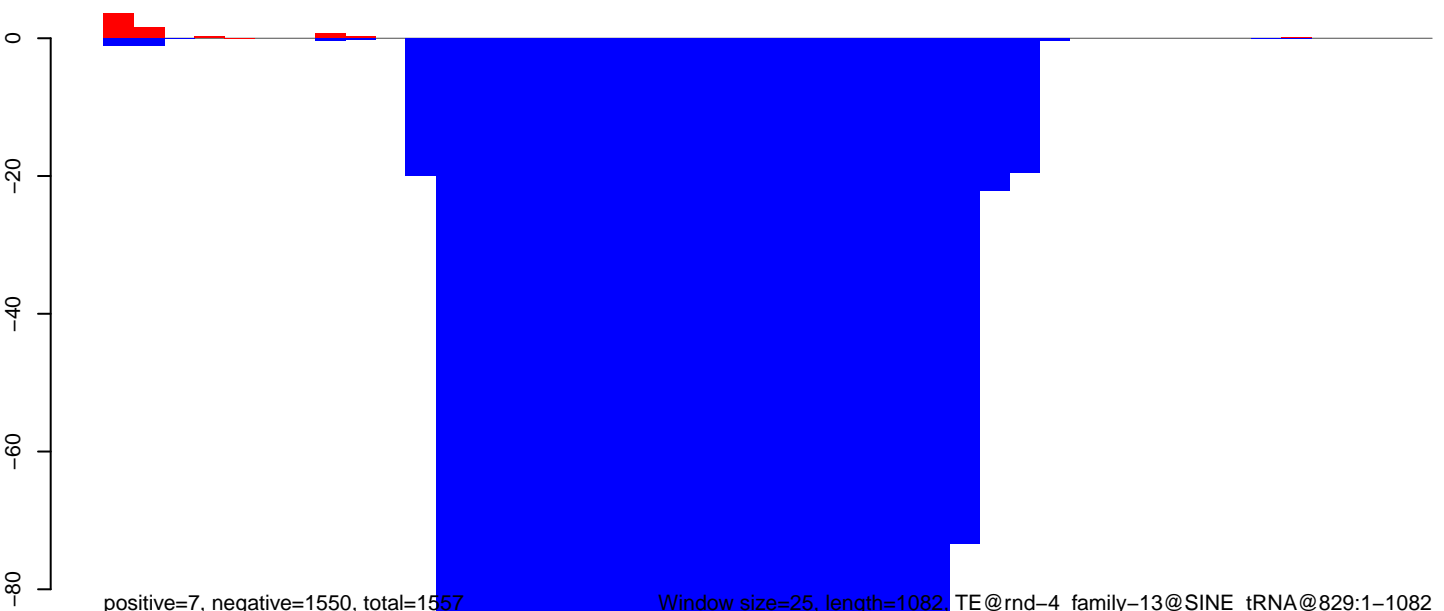
positive=1, negative=188, total=189

CuQuin_Testes_NL.24_35.rep



positive=5, negative=261, total=266

CuQuin_Testes_NL.rep

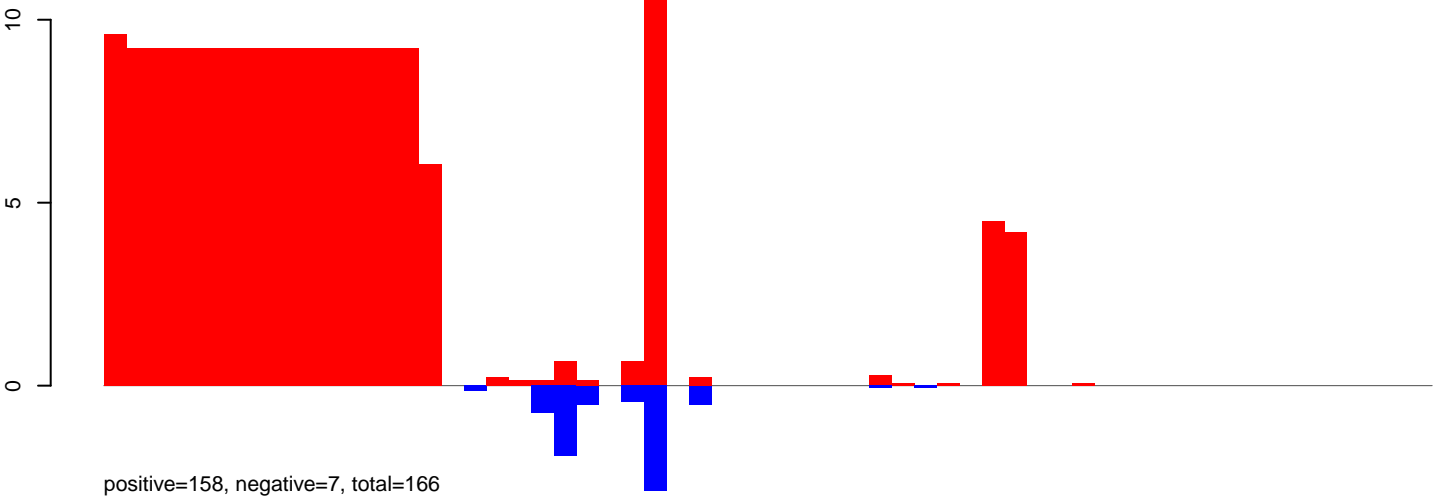


positive=7, negative=1550, total=1557

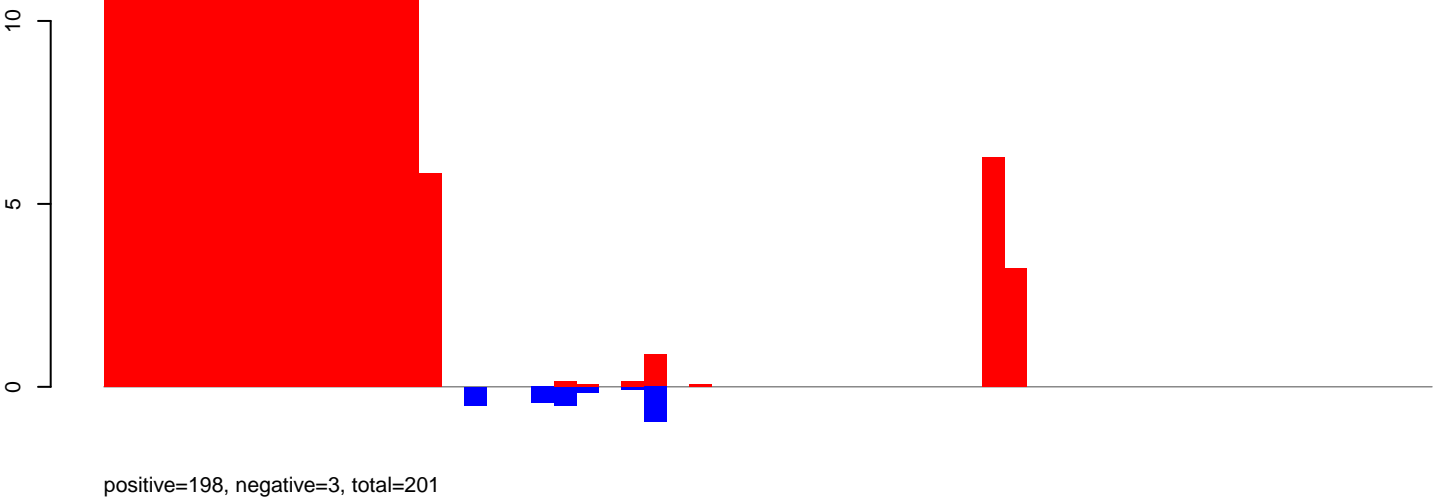
Window size=25, length=1082, TE@rnd-4_family-13@SINE_tRNA@829:1-1082

0 200 400 600 800 1000

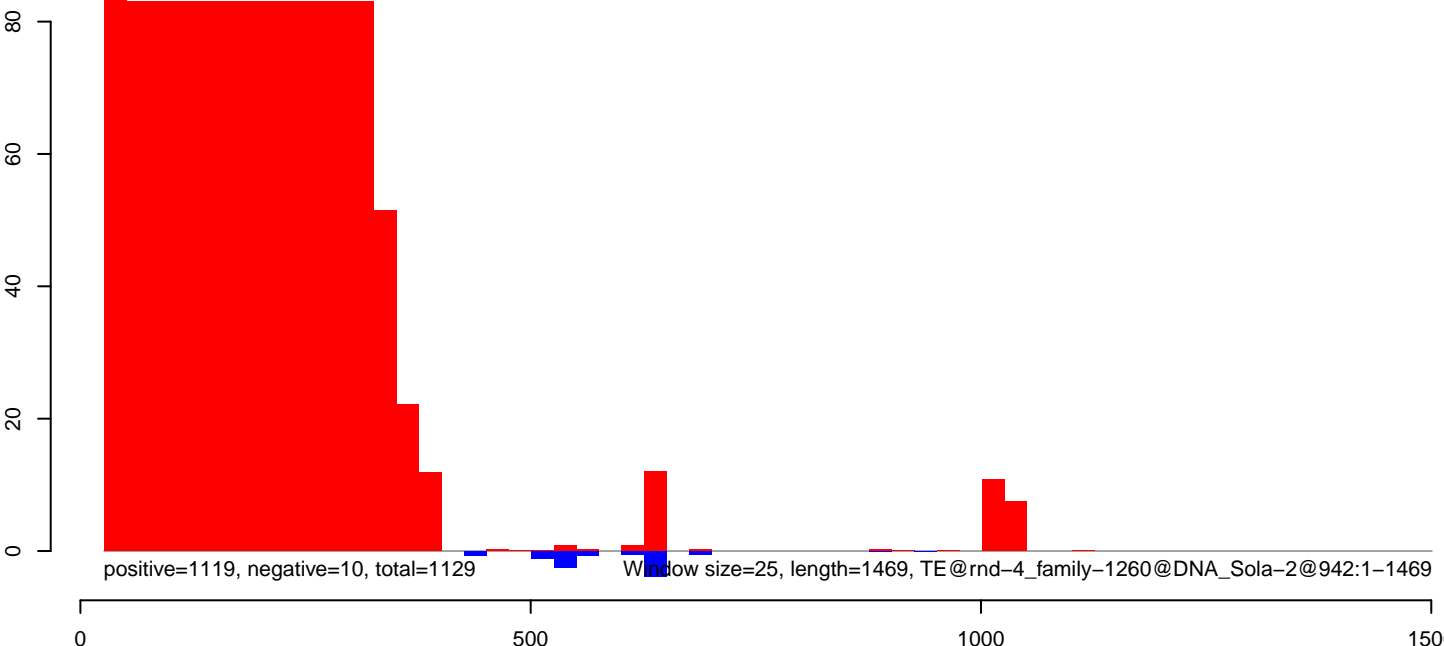
CuQuin_Testes_NL.18_23.rep



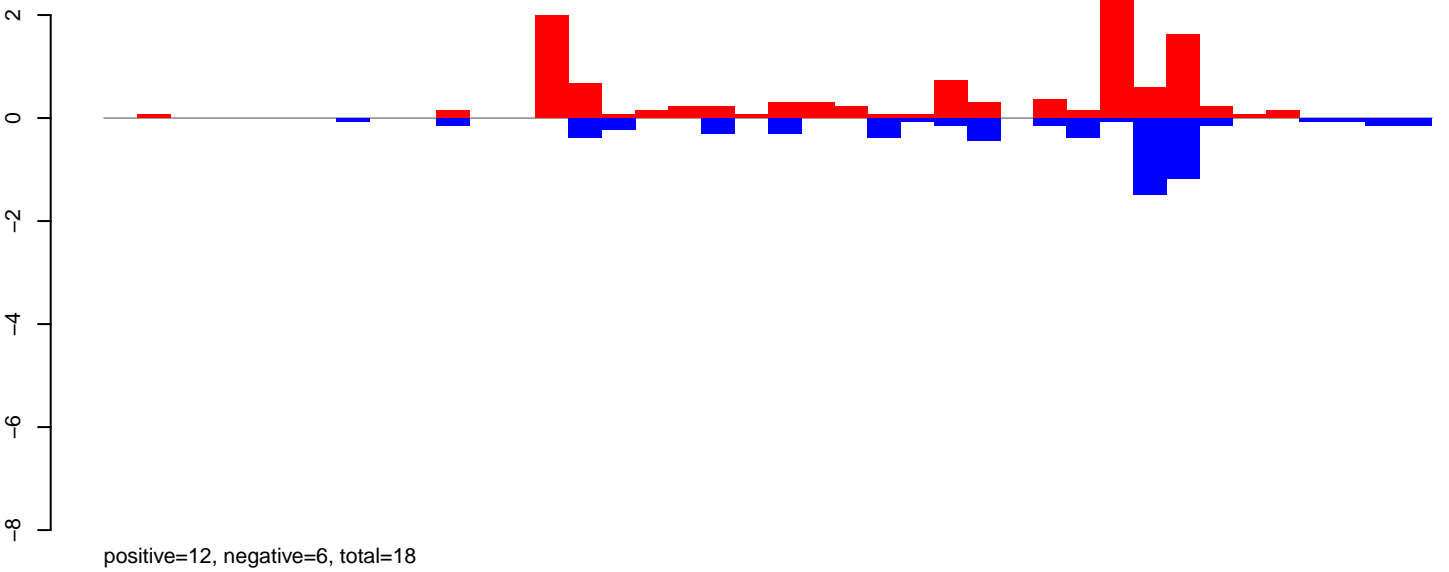
CuQuin_Testes_NL.24_35.rep



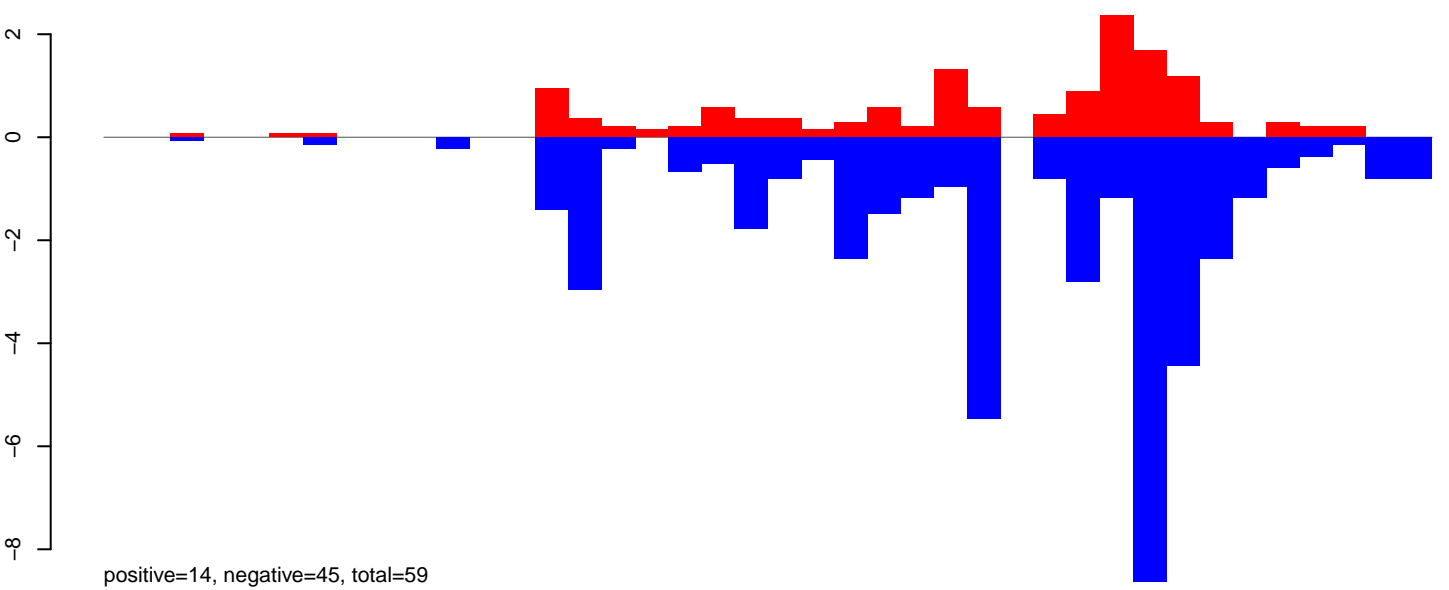
CuQuin_Testes_NL.rep



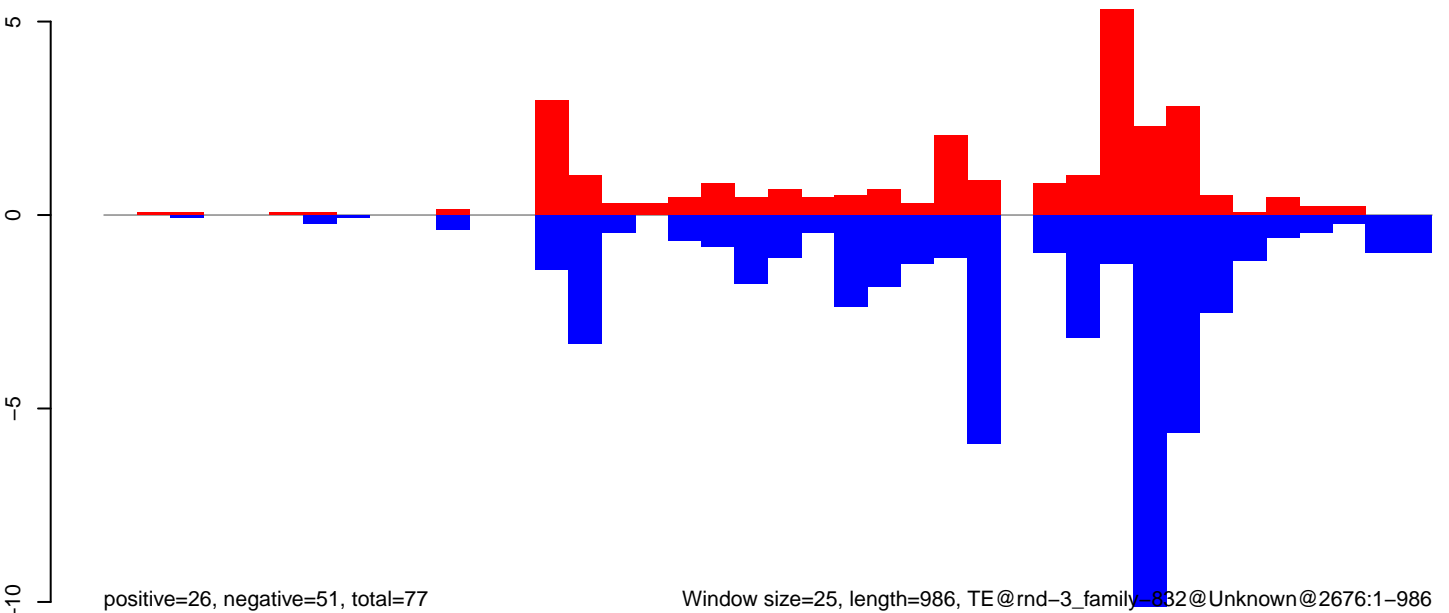
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

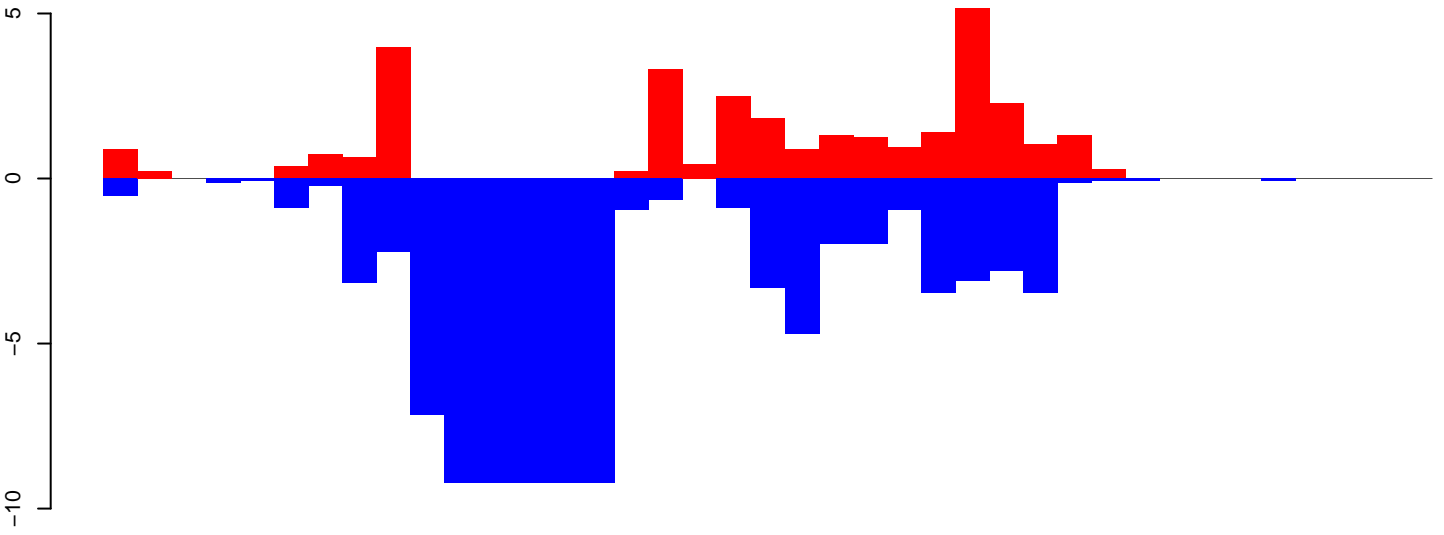


CuQuin_Testes_NL.rep



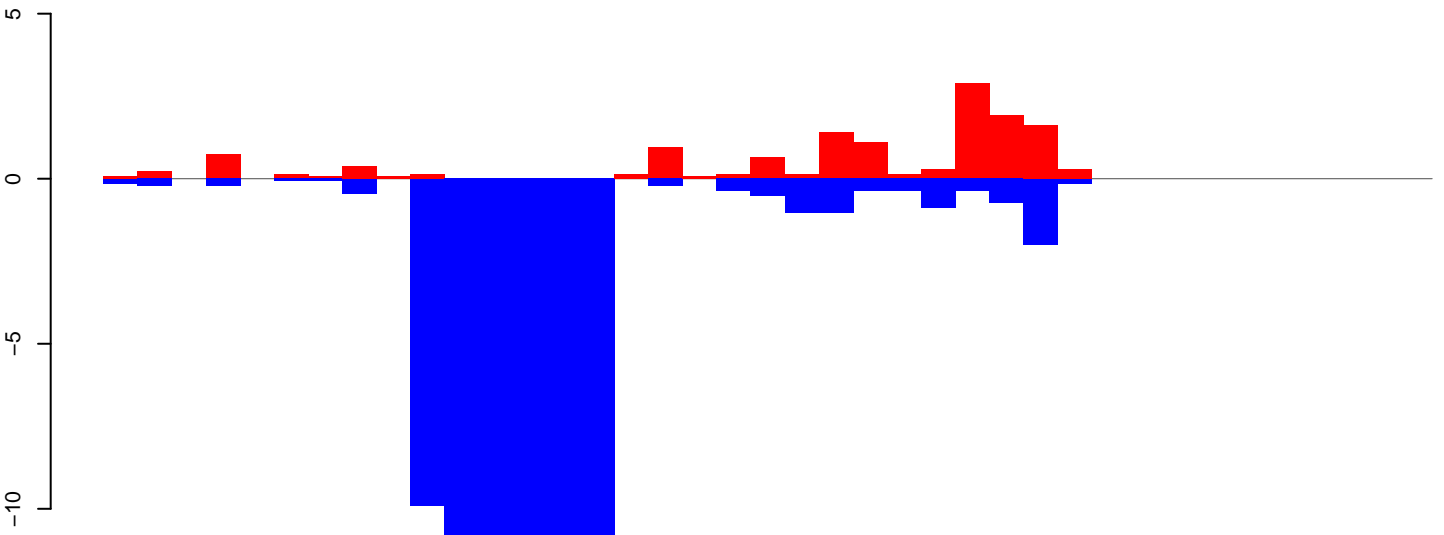
Window size=25, length=986, TE@rnd-3_family-832@Unknown@2676:1-986

CuQuin_Testes_NL.18_23.rep



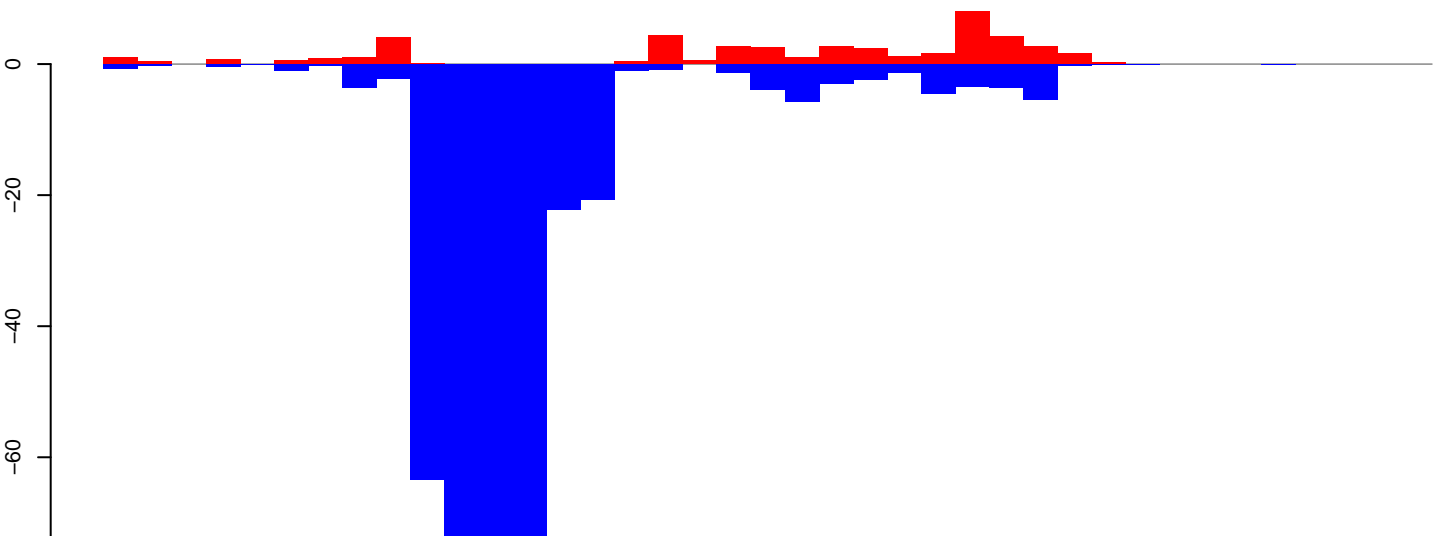
positive=31, negative=89, total=120

CuQuin_Testes_NL.24_35.rep



positive=14, negative=82, total=96

CuQuin_Testes_NL.rep

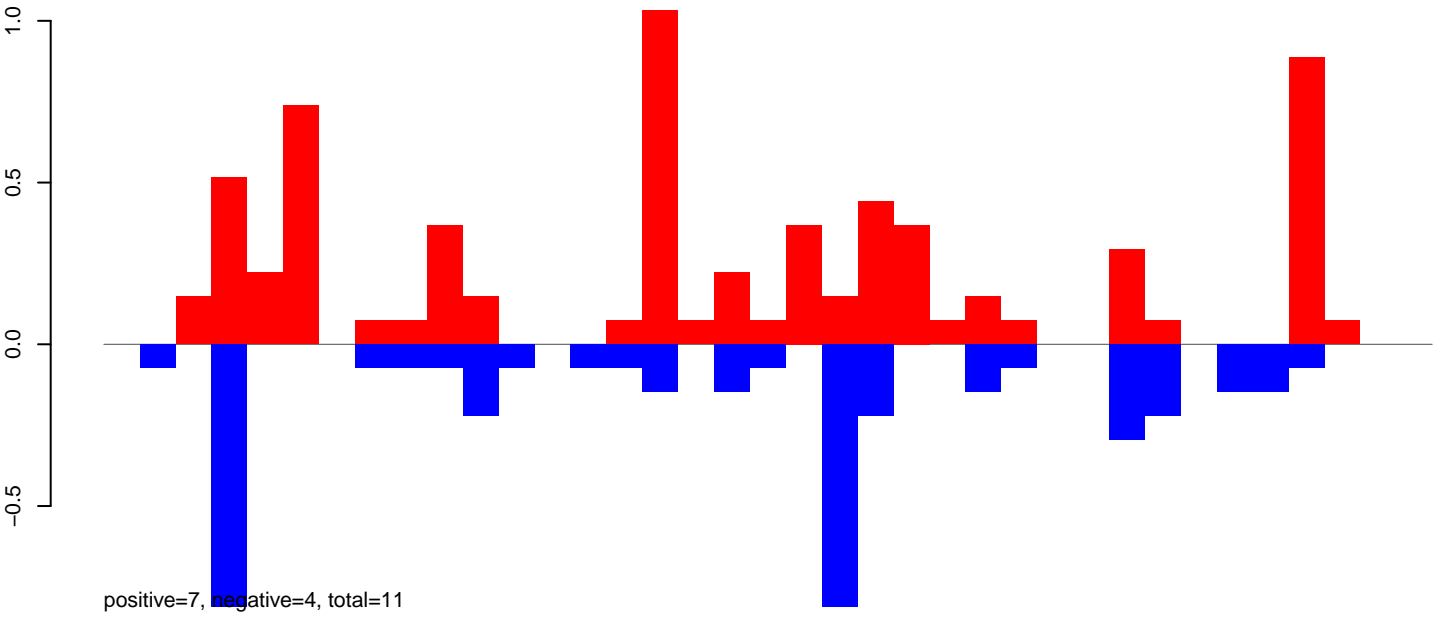


positive=45, negative=396, total=441

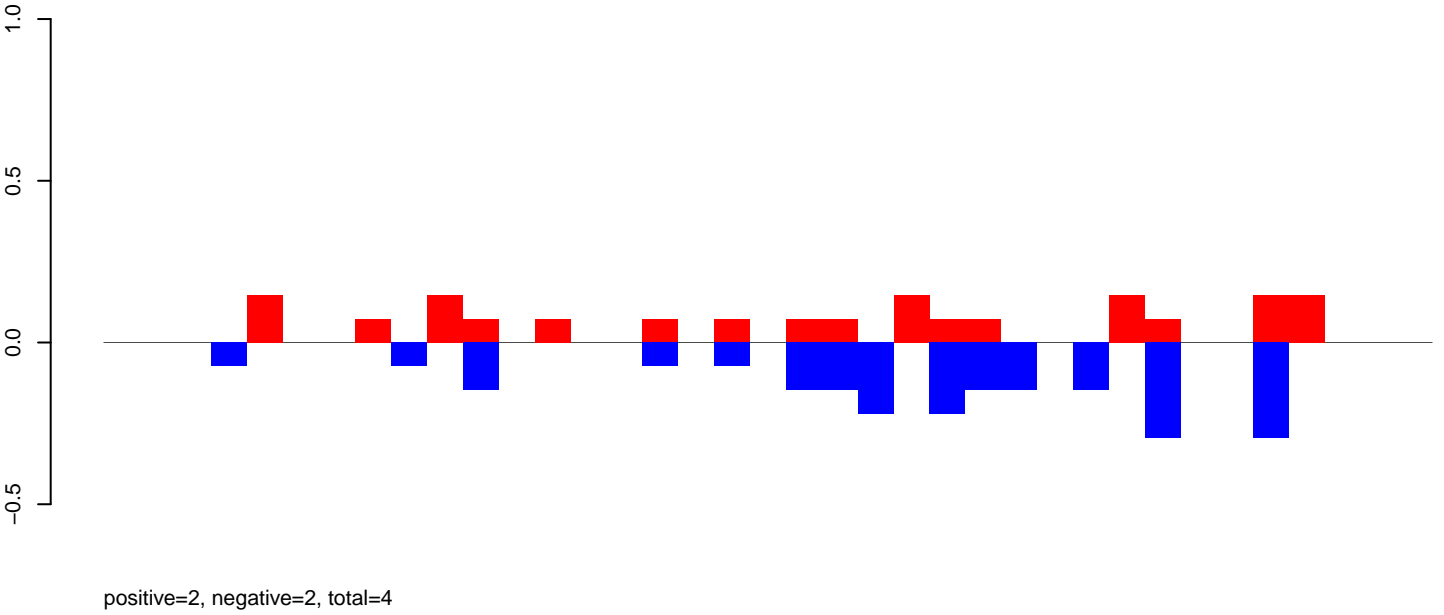
Window size=25, length=953, TE@rnd-3_family-34@DNA_Sola-2@39:1-953

0 200 400 600 800 1000

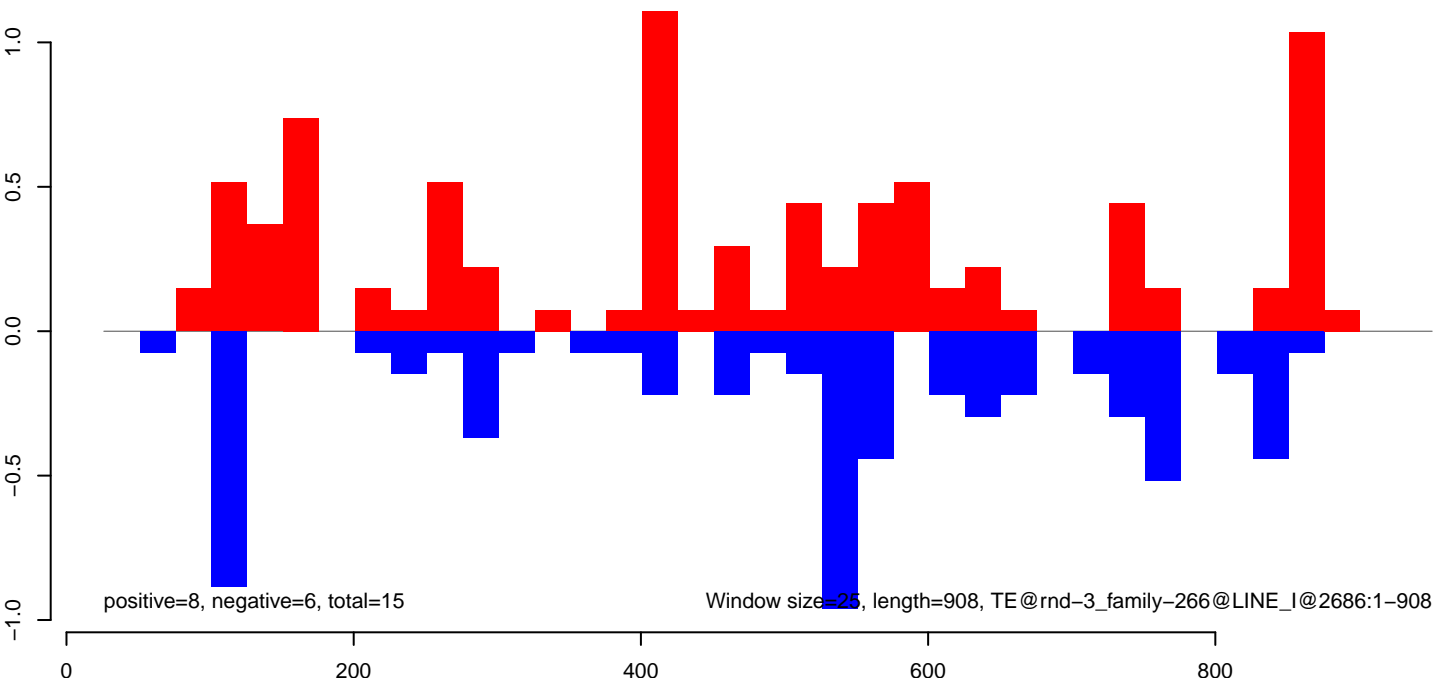
CuQuin_Testes_NL.18_23.rep



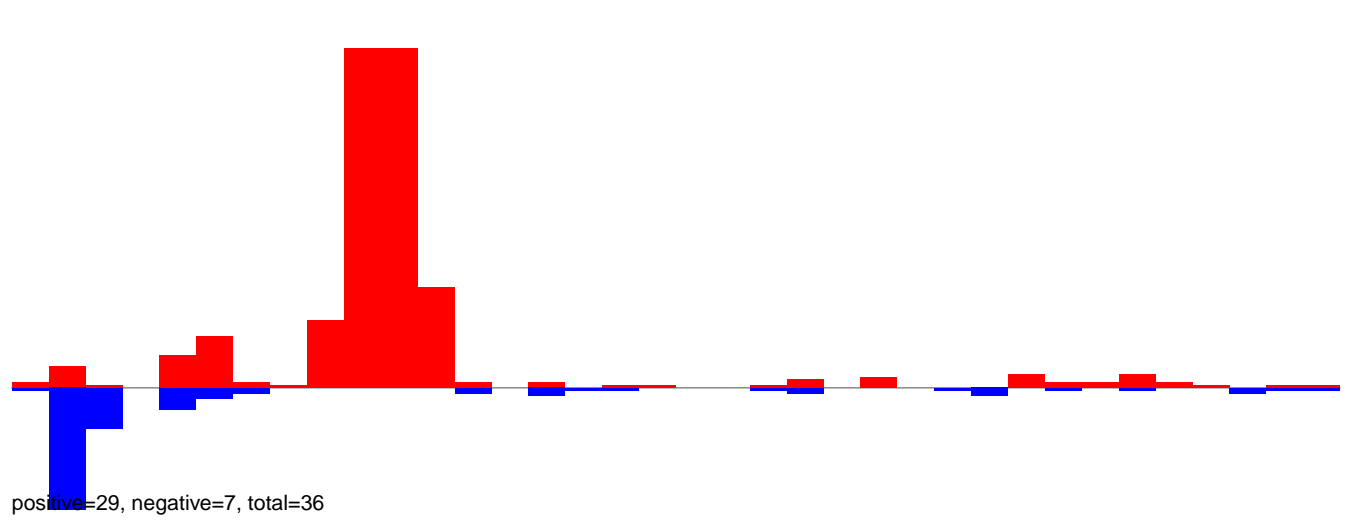
CuQuin_Testes_NL.24_35.rep



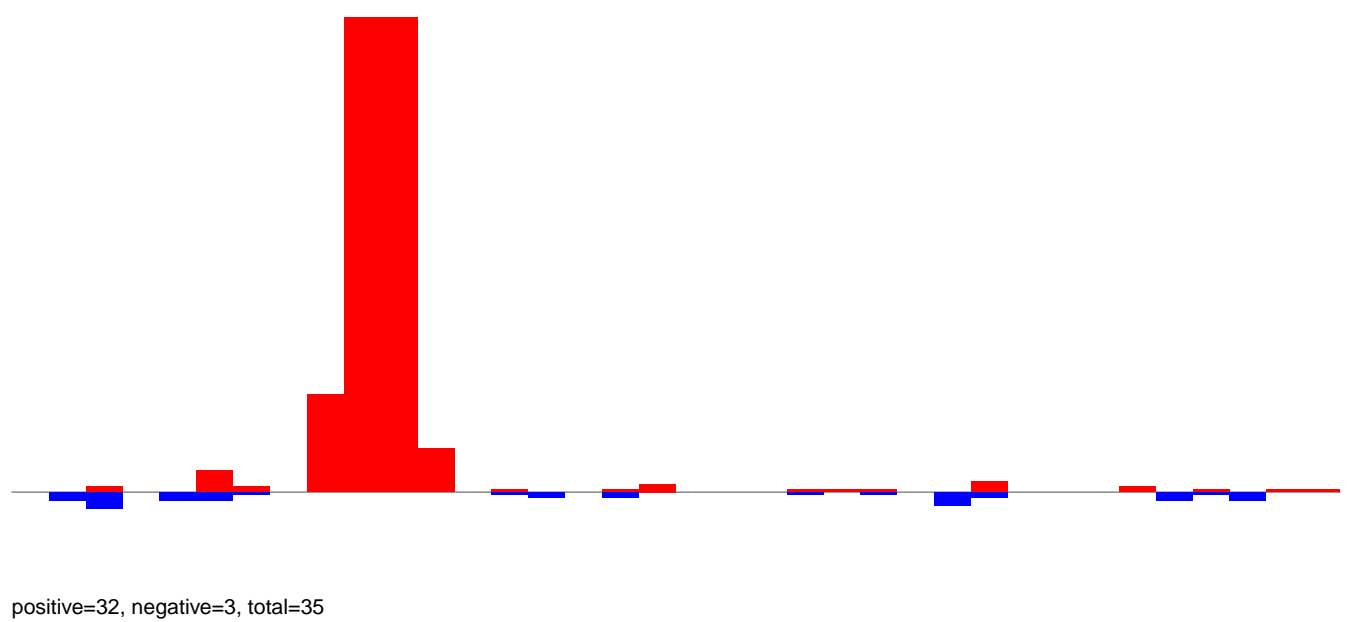
CuQuin_Testes_NL.rep



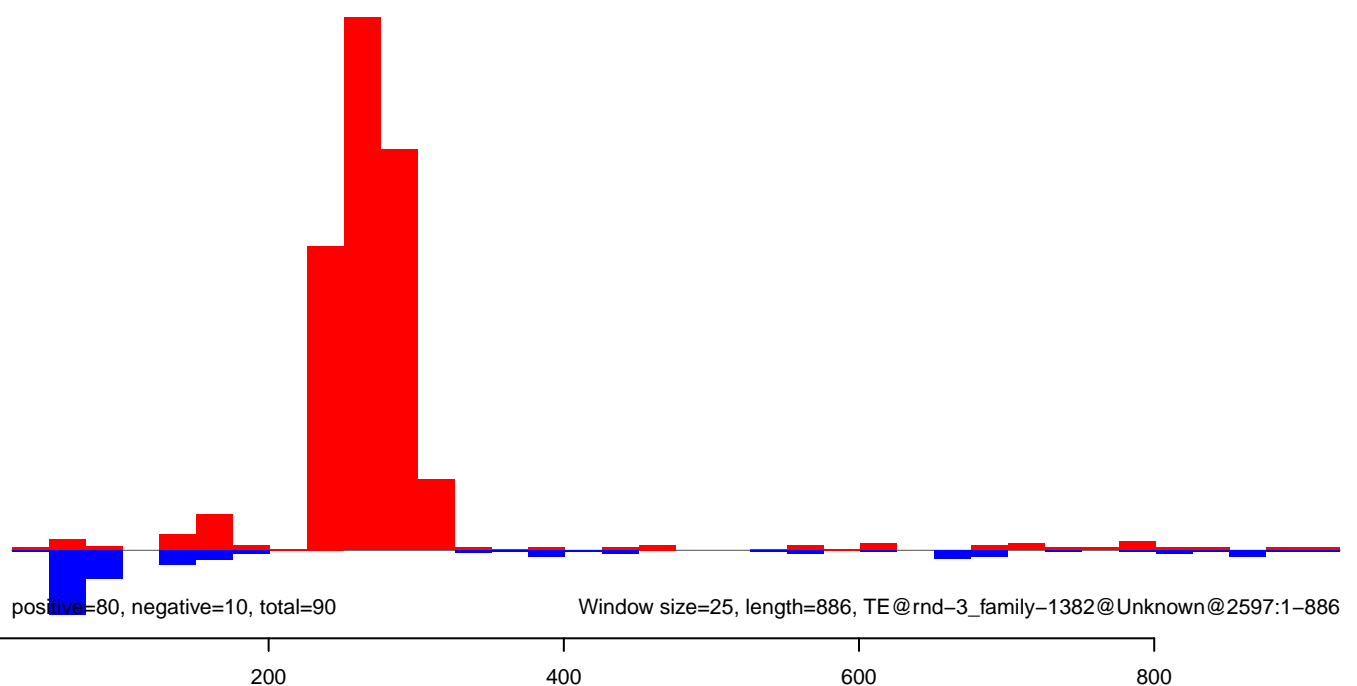
CuQuin_Testes_NL.18_23.rep



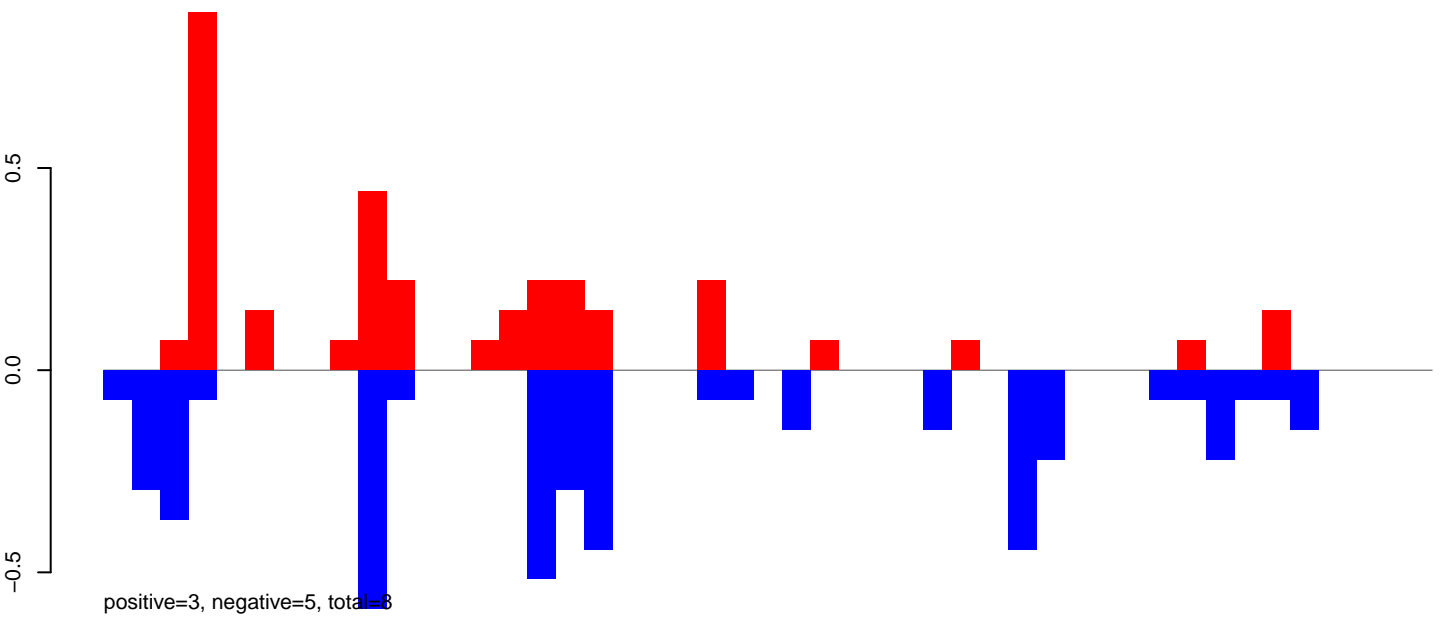
CuQuin_Testes_NL.24_35.rep



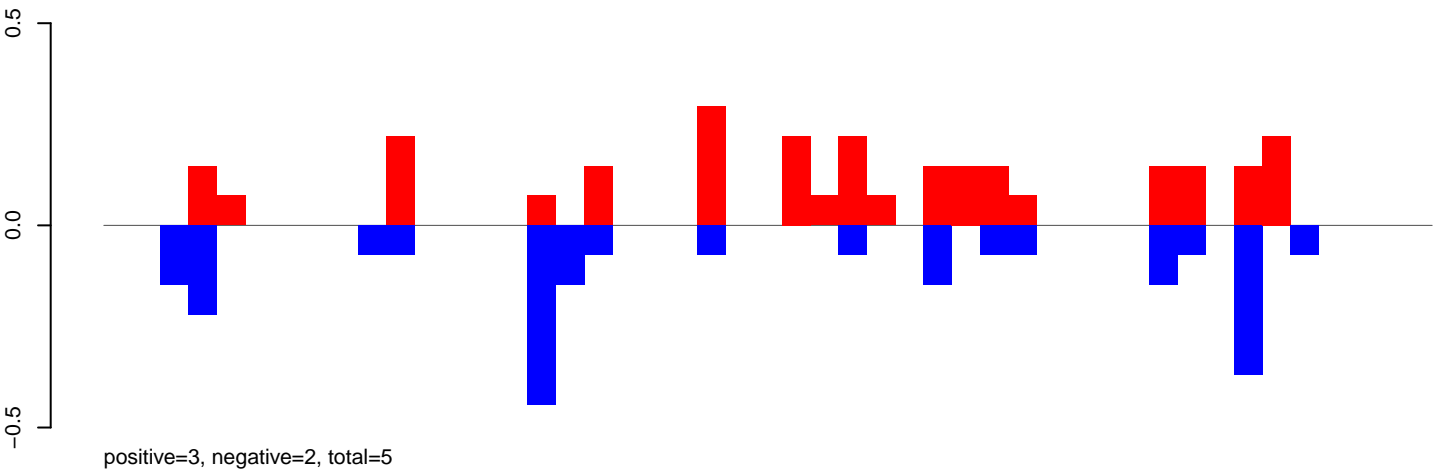
CuQuin_Testes_NL.rep



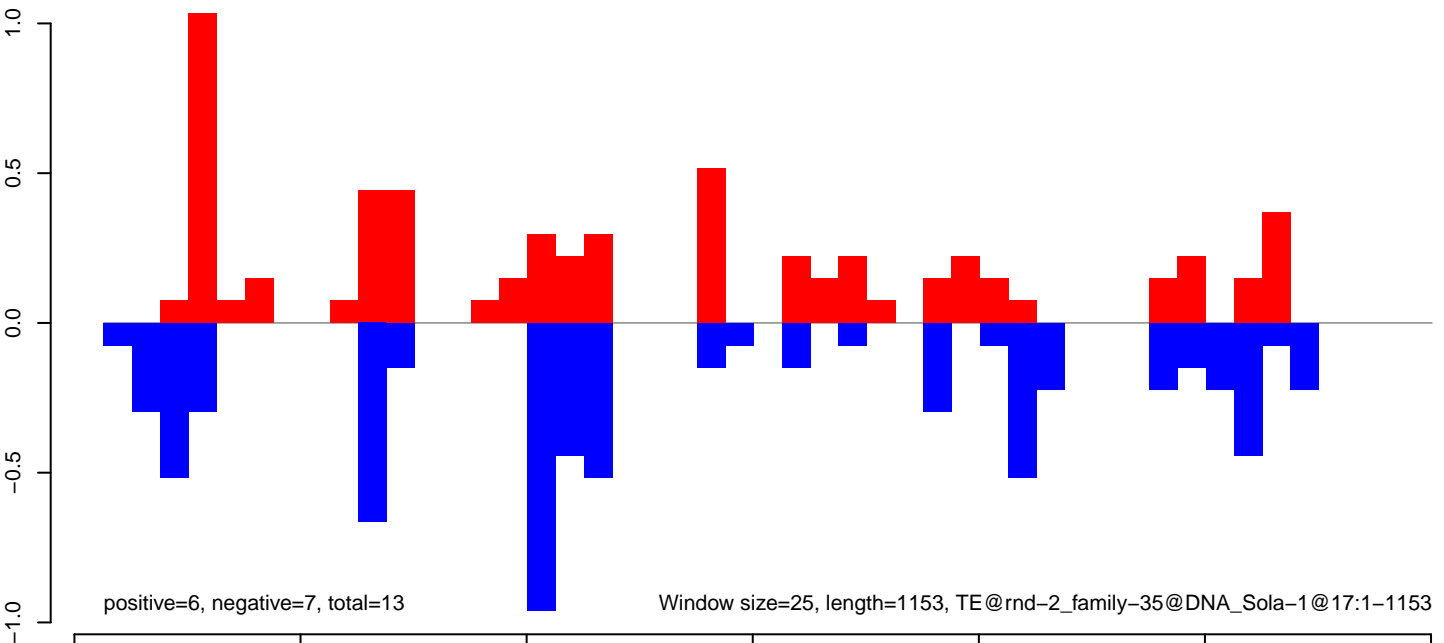
CuQuin_Testes_NL.18_23.rep



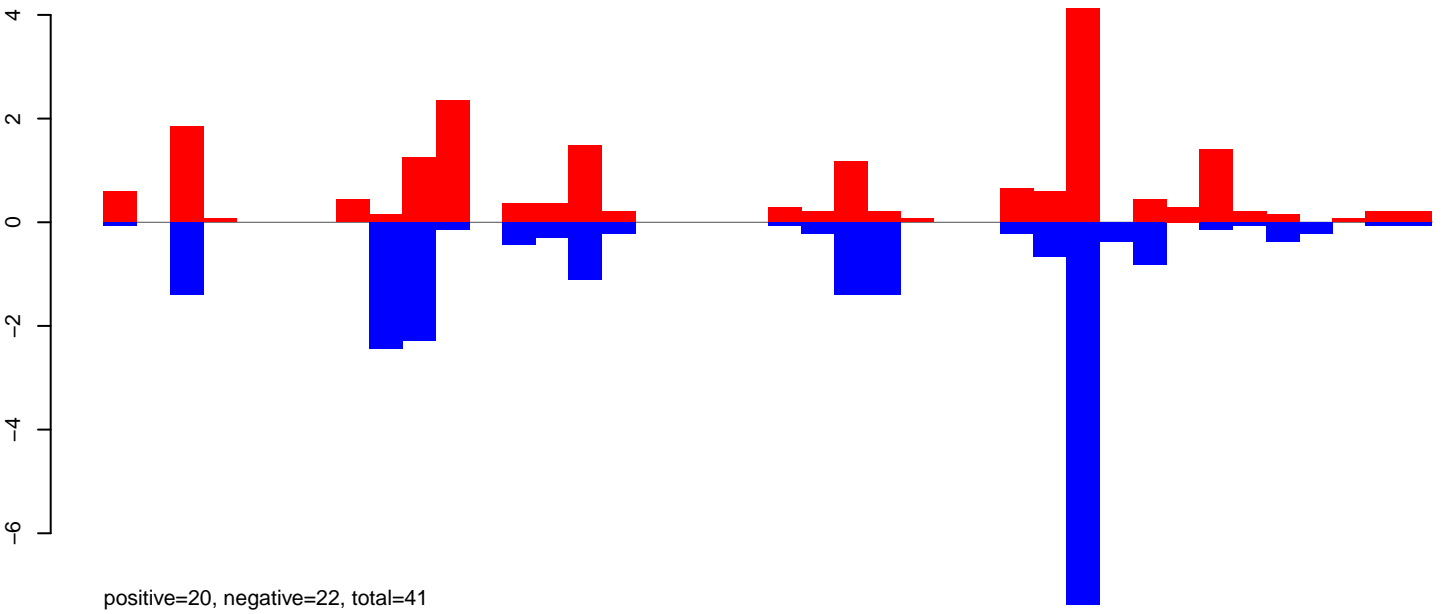
CuQuin_Testes_NL.24_35.rep



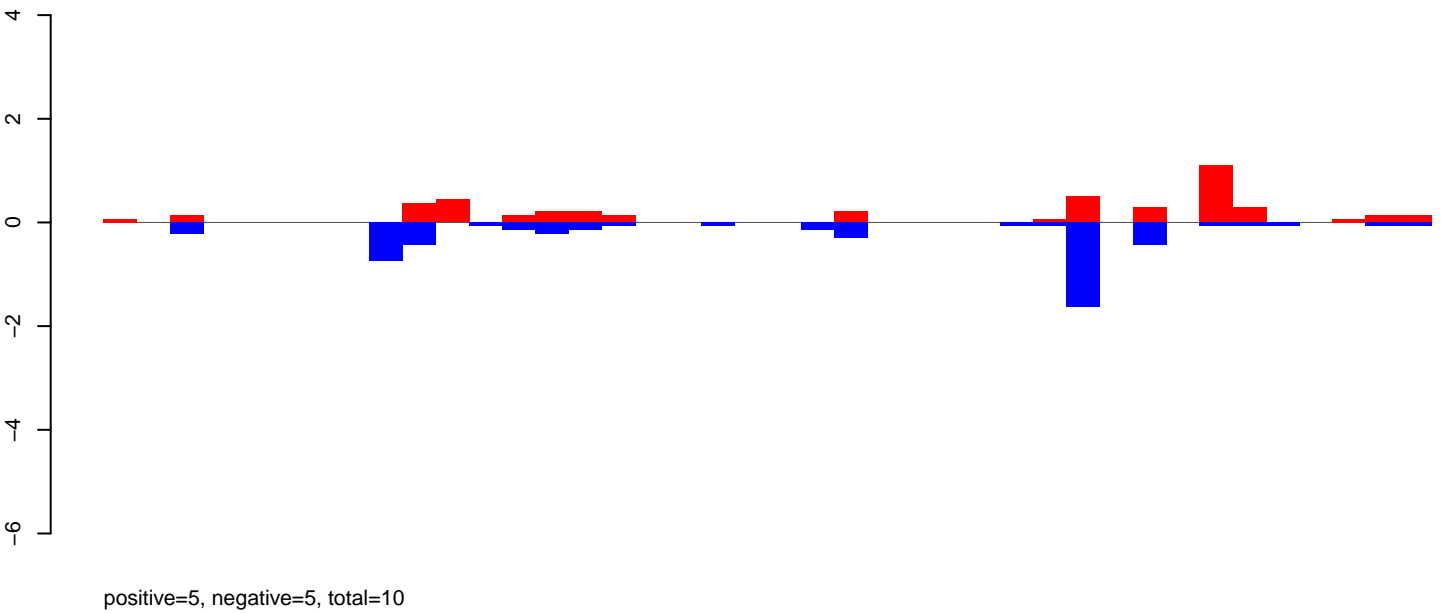
CuQuin_Testes_NL.rep



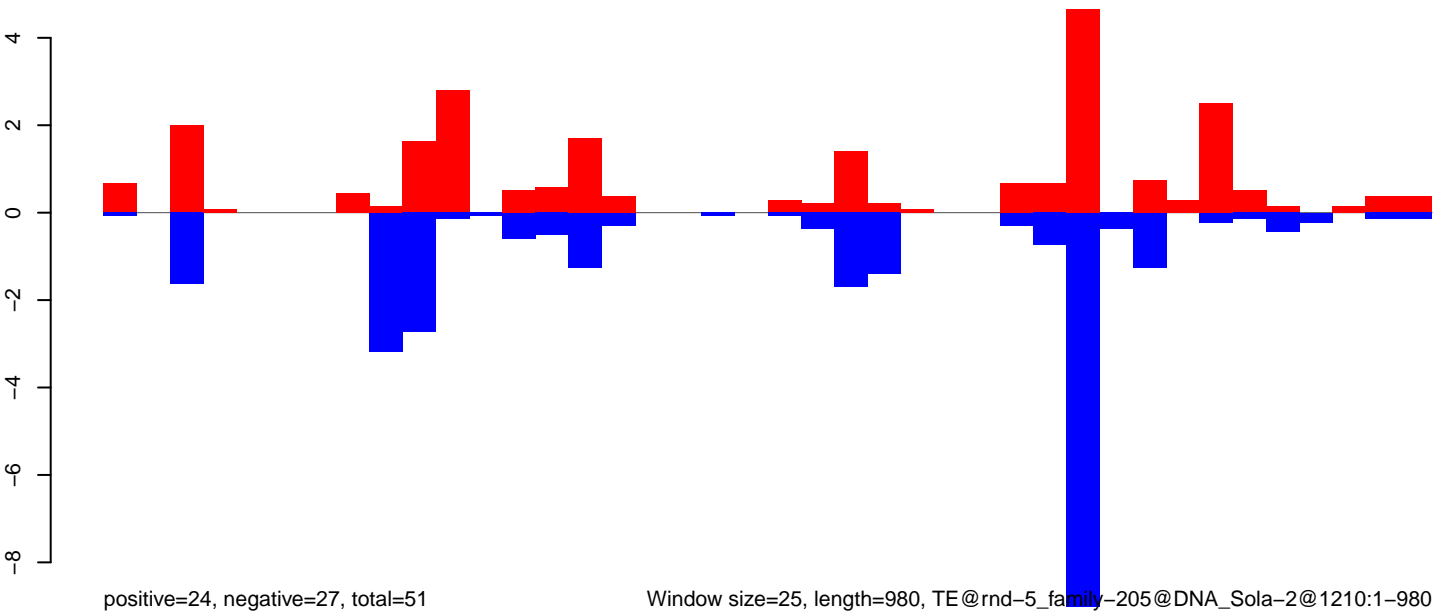
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

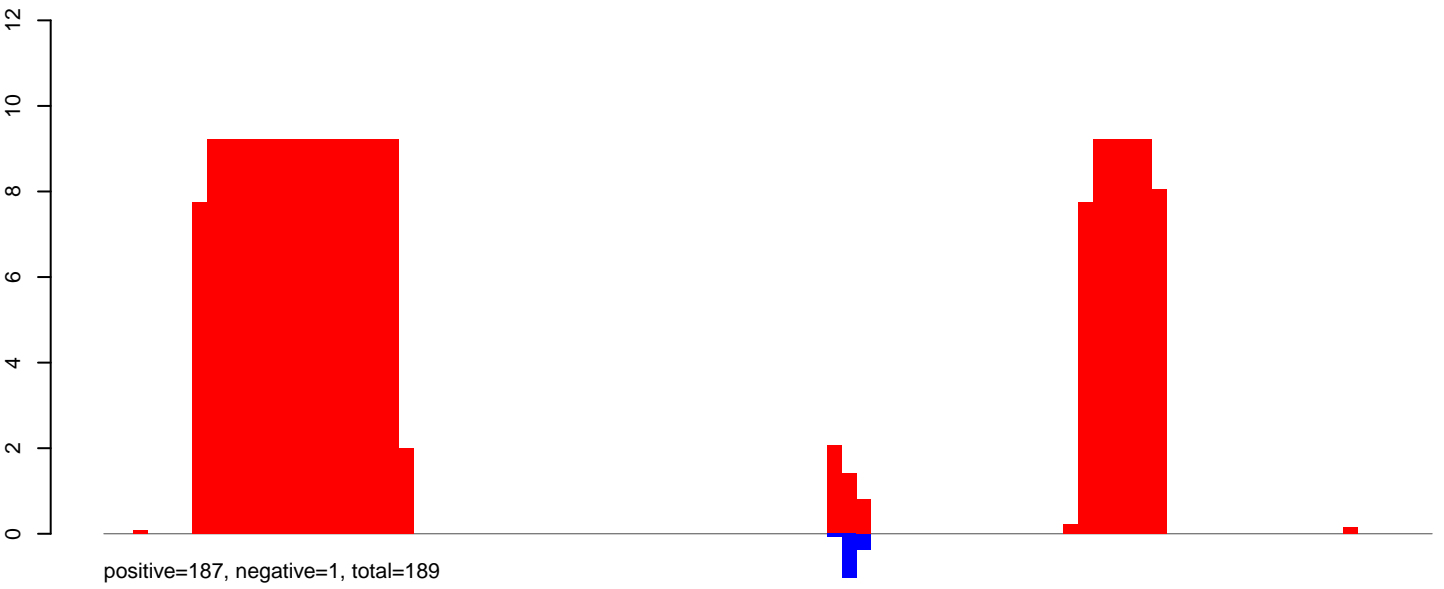


CuQuin_Testes_NL.rep

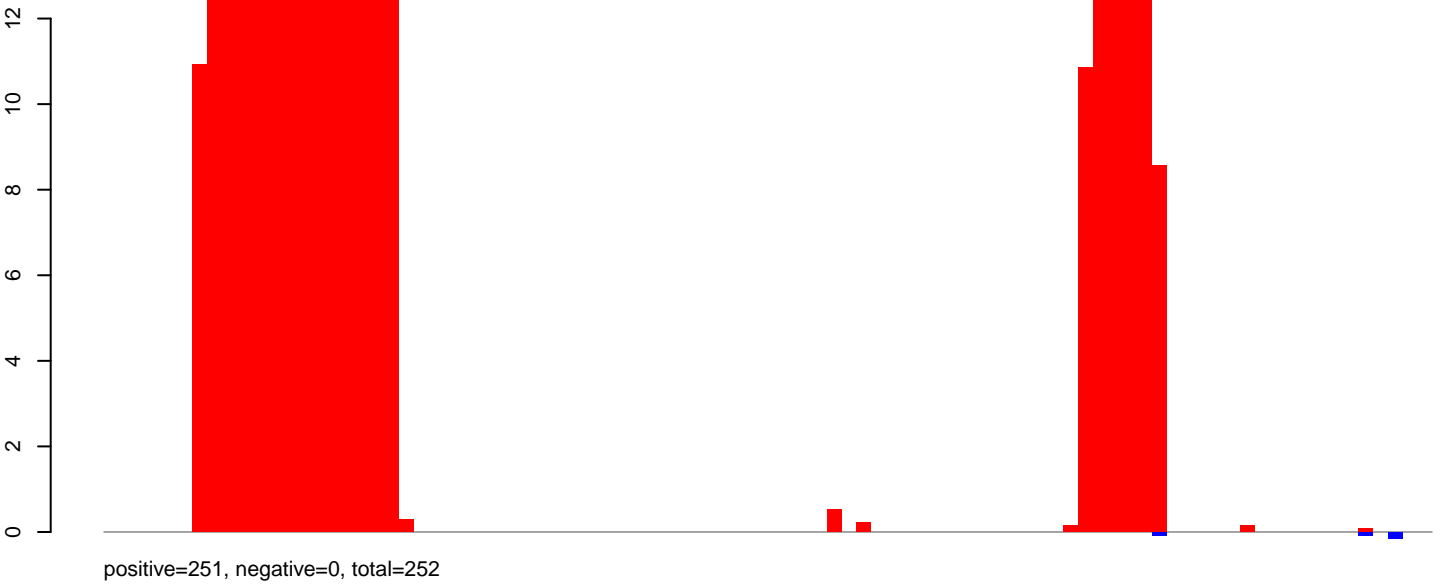


Window size=25, length=980, TE@rnd-5_family-205@DNA_Sola-2@1210:1-980

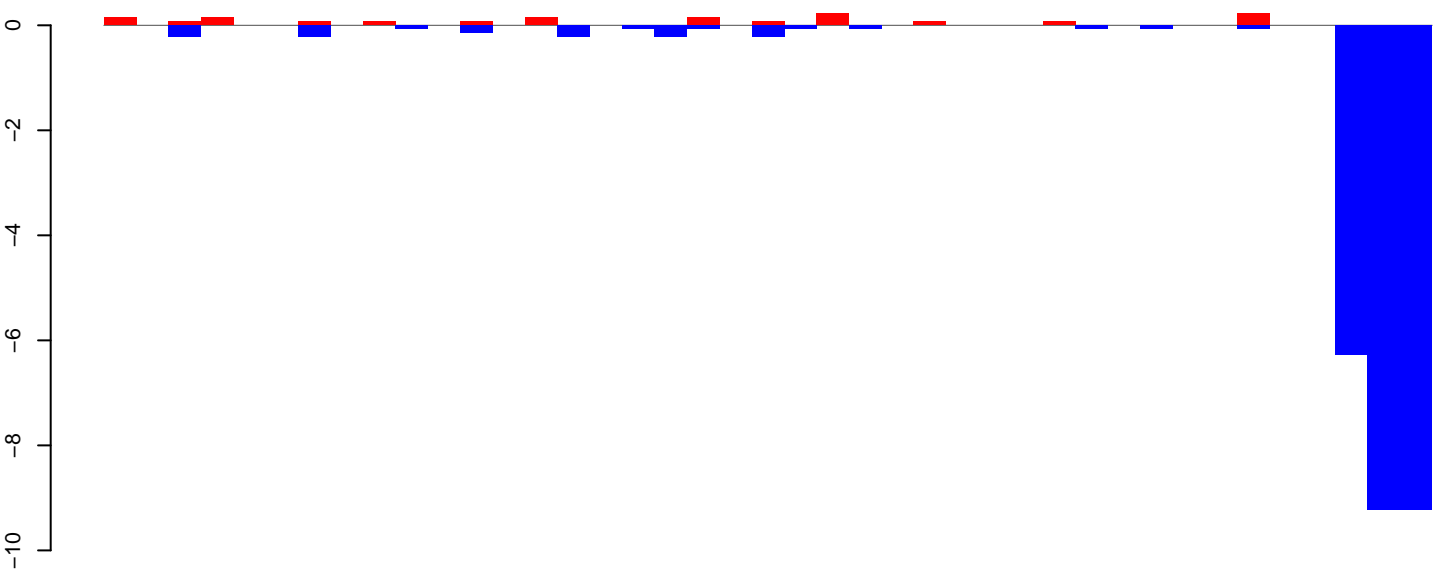
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

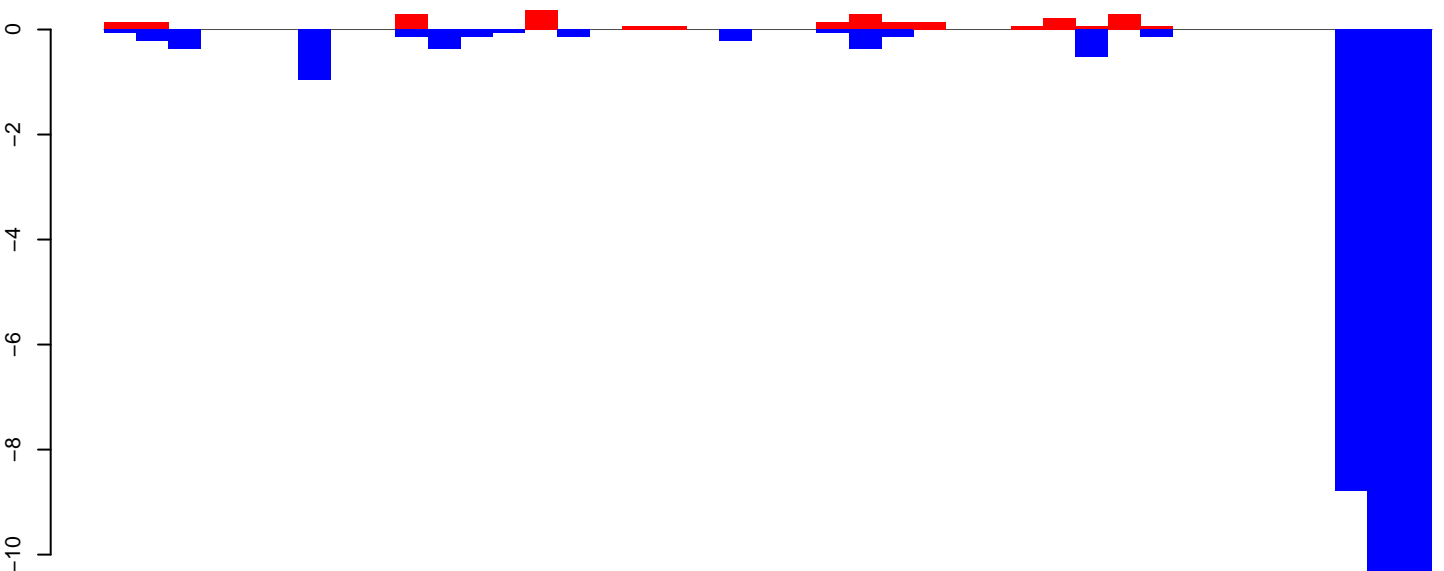


CuQuin_Testes_NL.18_23.rep



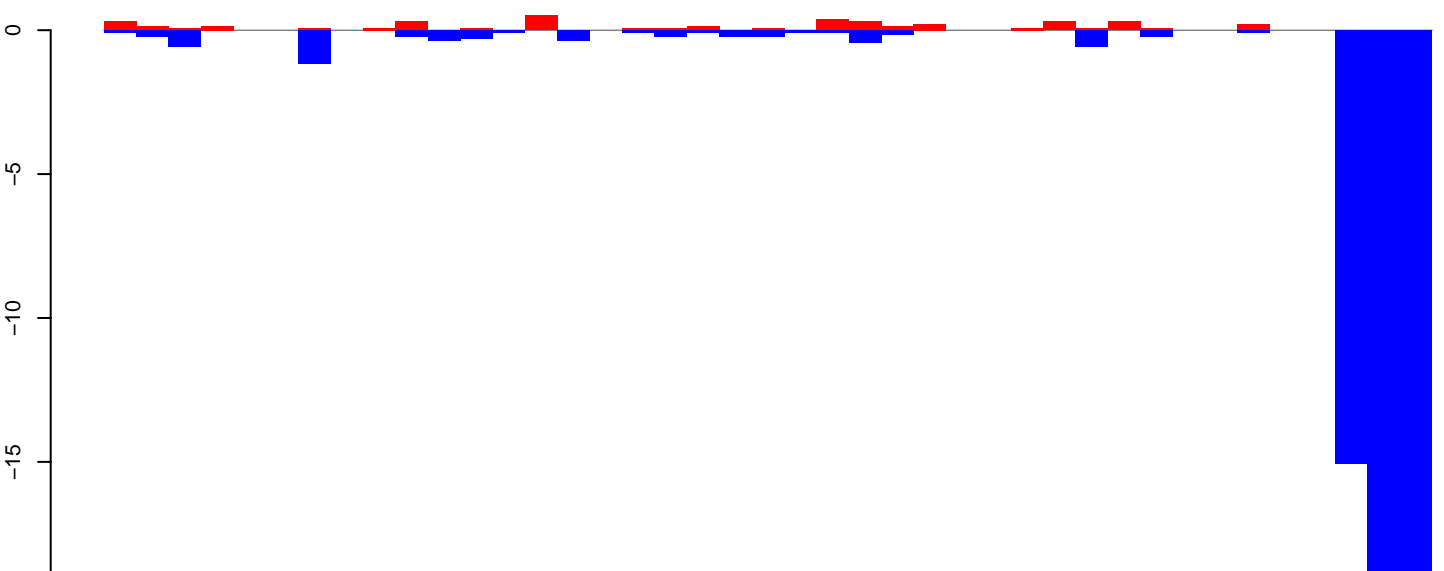
positive=2, negative=27, total=28

CuQuin_Testes_NL.24_35.rep



positive=3, negative=35, total=37

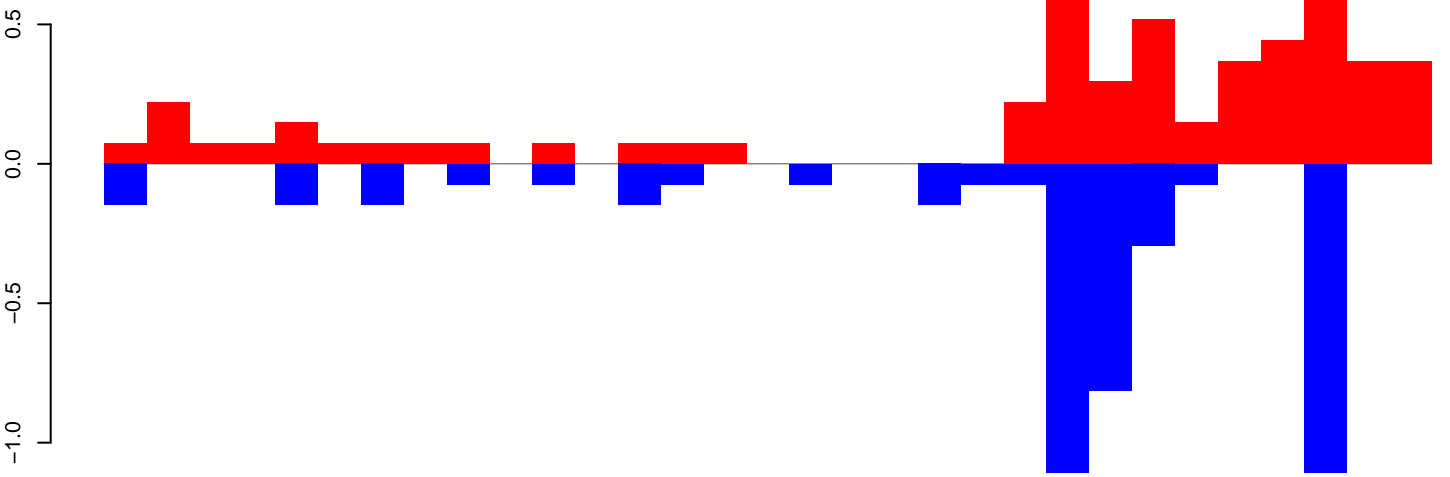
CuQuin_Testes_NL.rep



positive=4, negative=61, total=65

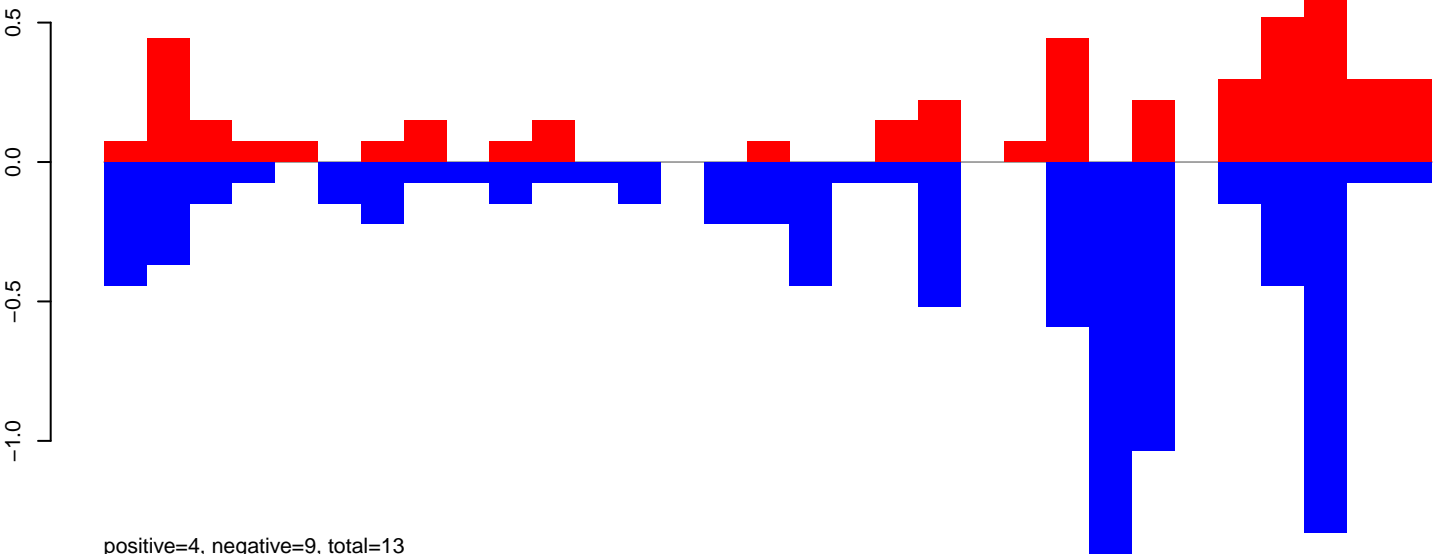
Window size=25, length=1023, TE@rnd-4_family-508@LINE_Jockey@2765:1-1023

CuQuin_Testes_NL.18_23.rep



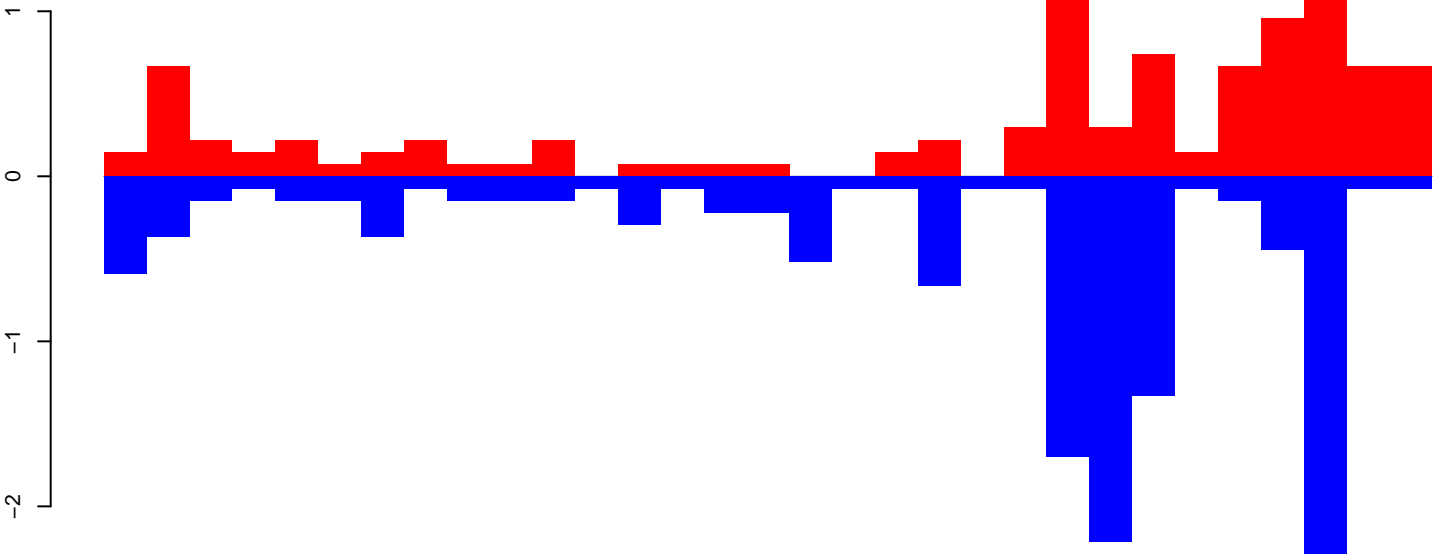
positive=5, negative=5, total=10

CuQuin_Testes_NL.24_35.rep



positive=4, negative=9, total=13

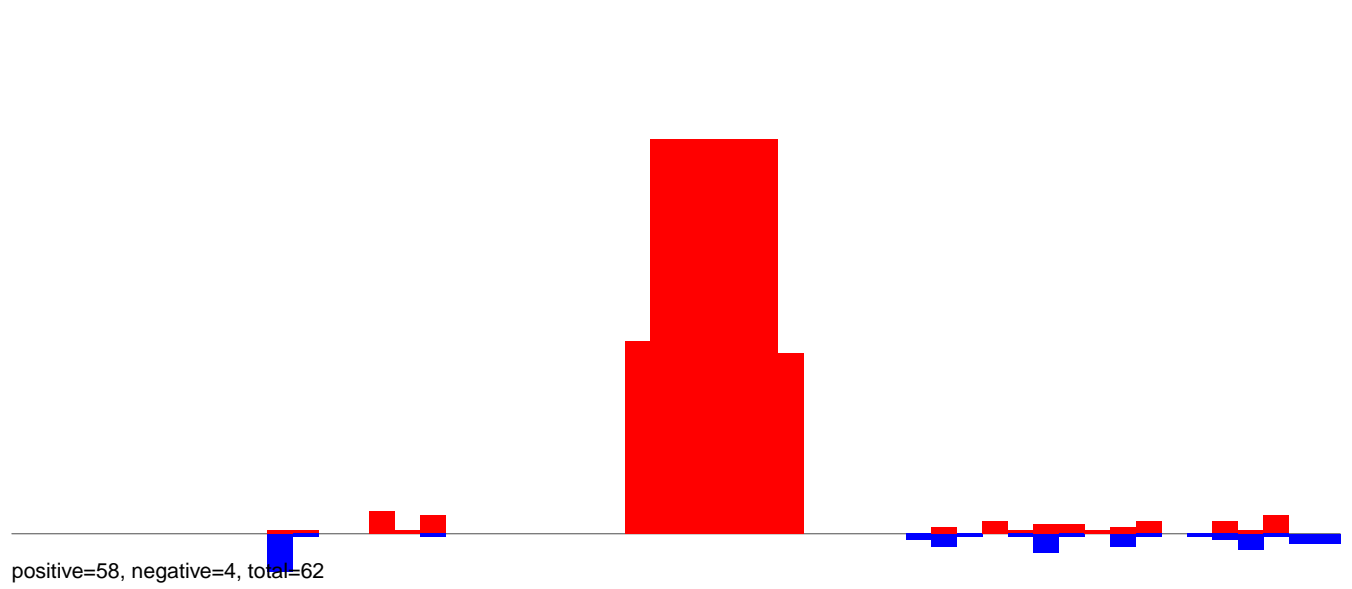
CuQuin_Testes_NL.rep



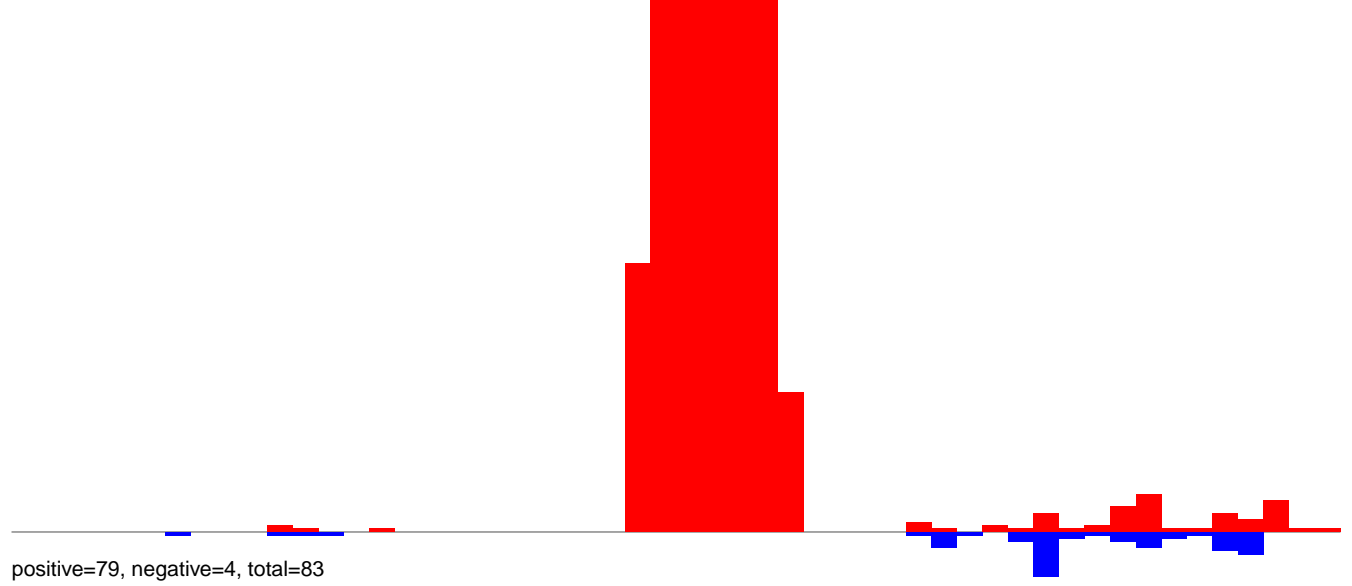
positive=10, negative=13, total=23

Window size=25, length=755, TE@rnd-4_family-4@Unknown@176:1-755

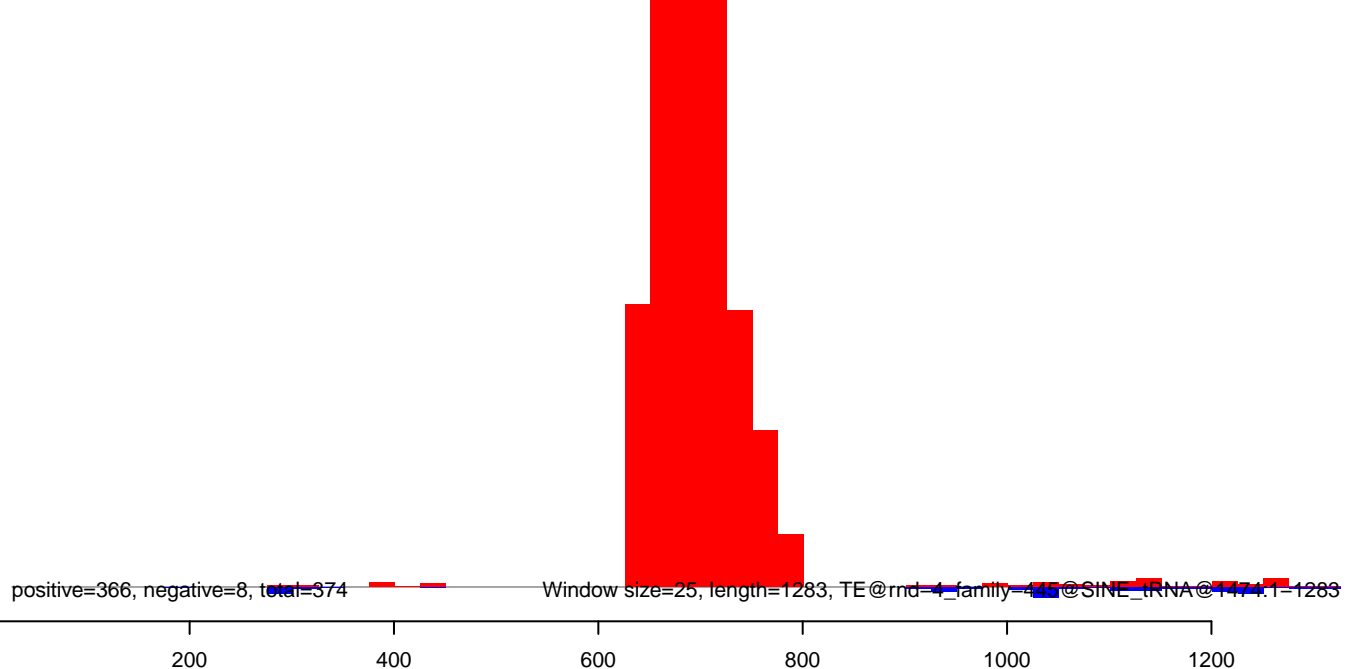
CuQuin_Testes_NL.18_23.rep



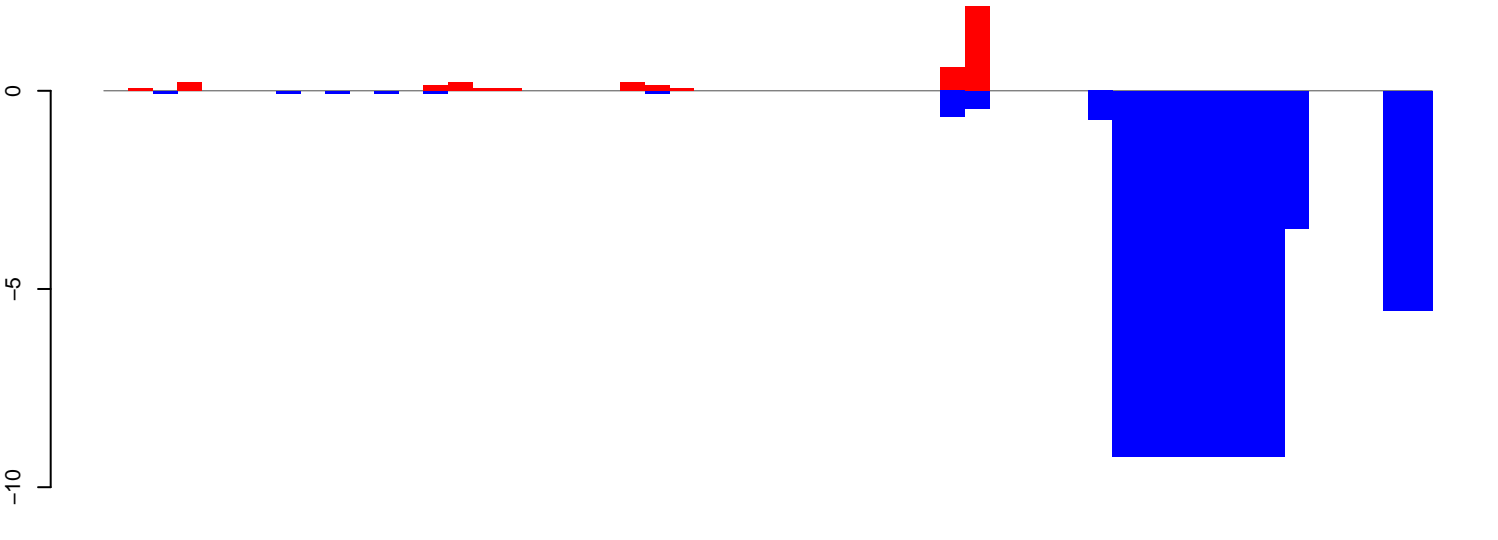
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

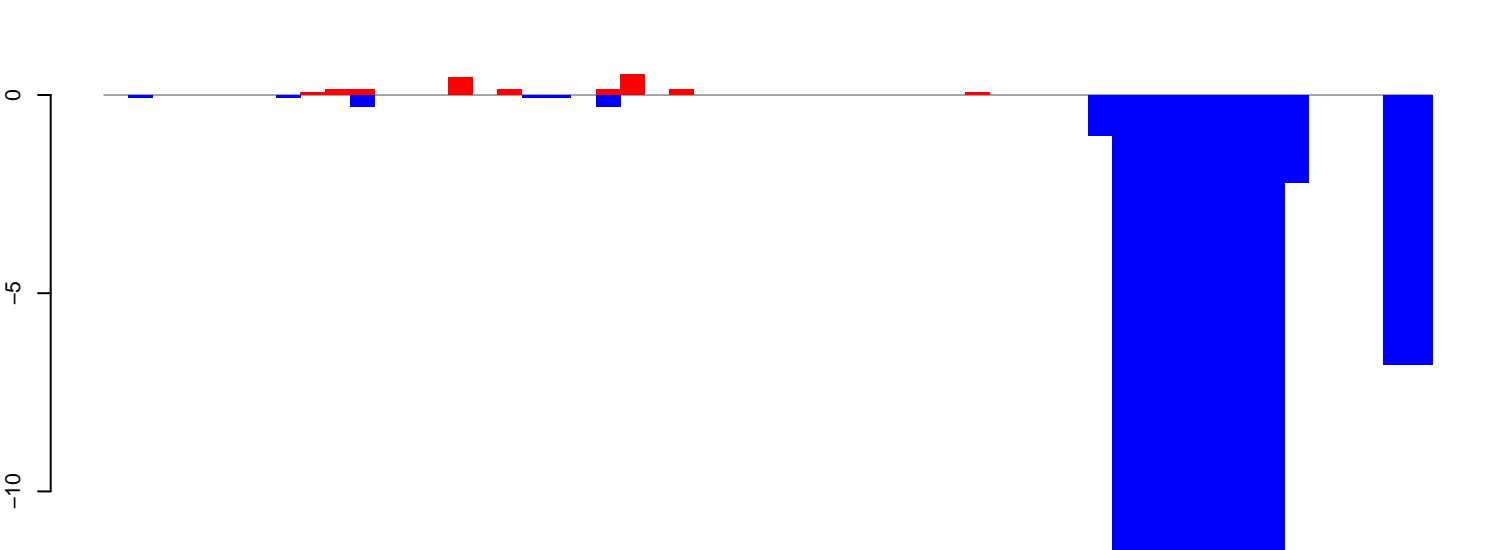


CuQuin_Testes_NL.18_23.rep



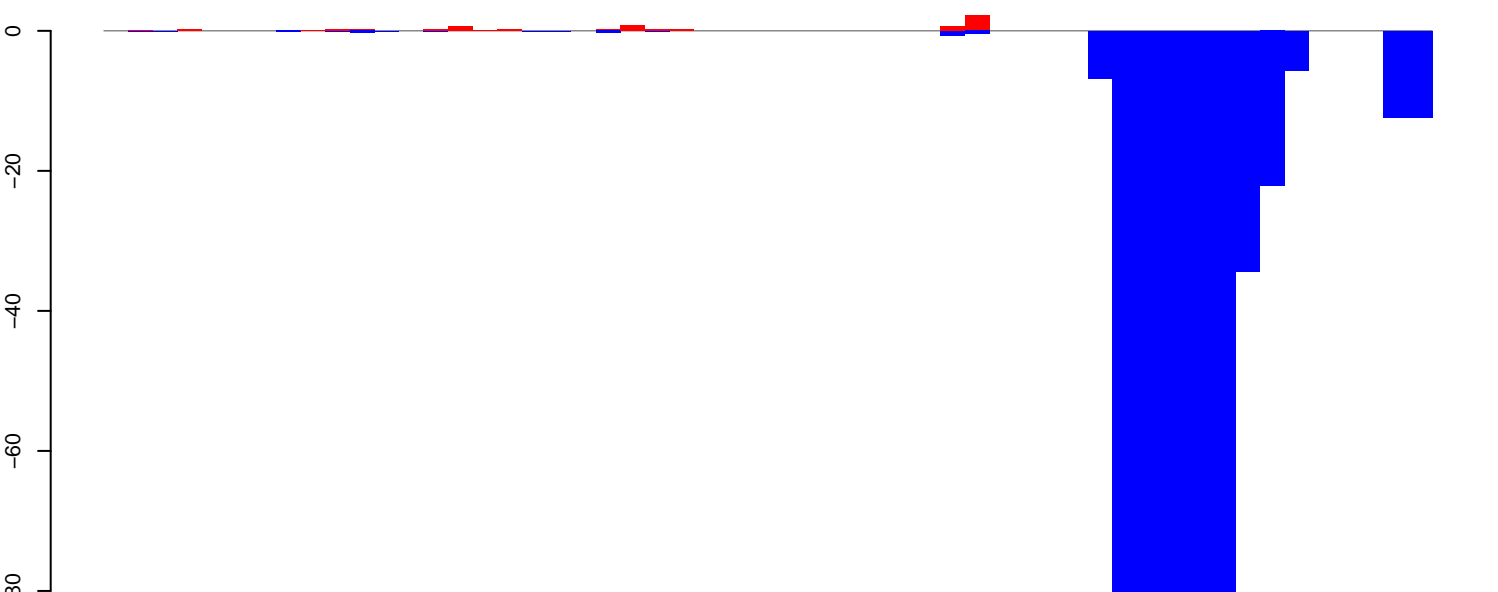
positive=4, negative=81, total=85

CuQuin_Testes_NL.24_35.rep



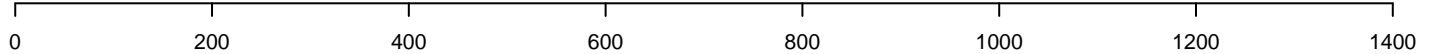
positive=2, negative=108, total=110

CuQuin_Testes_NL.rep

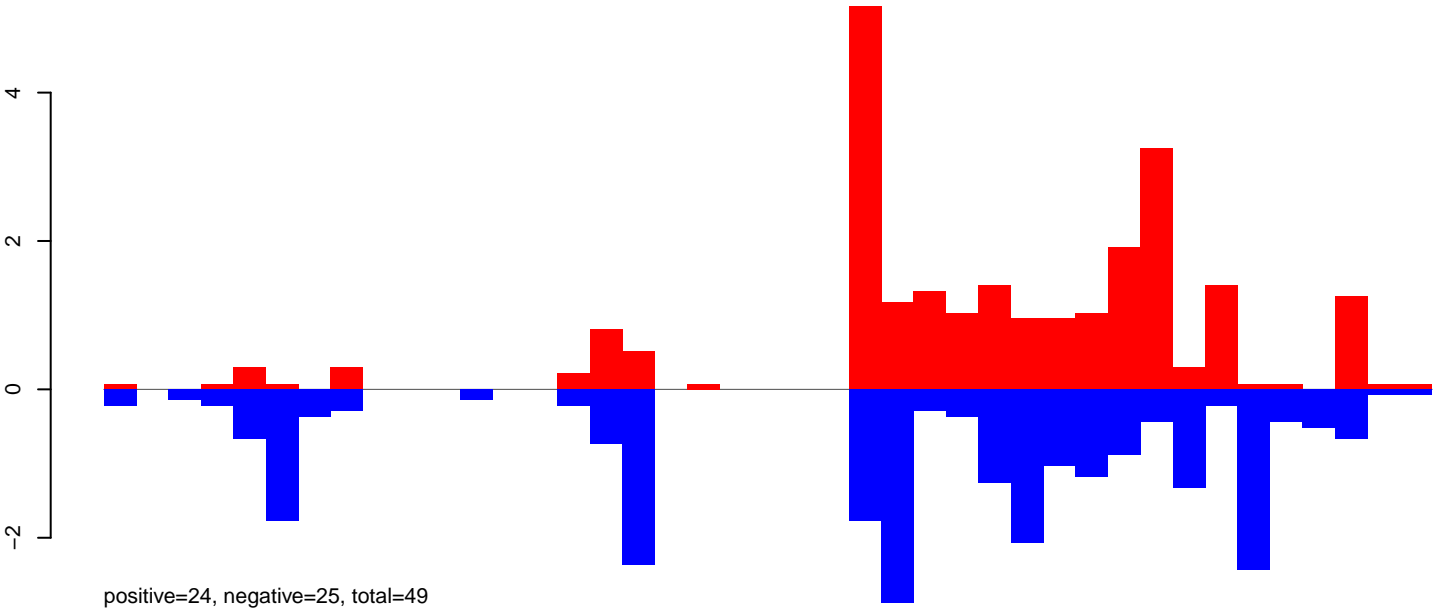


positive=6, negative=511, total=517

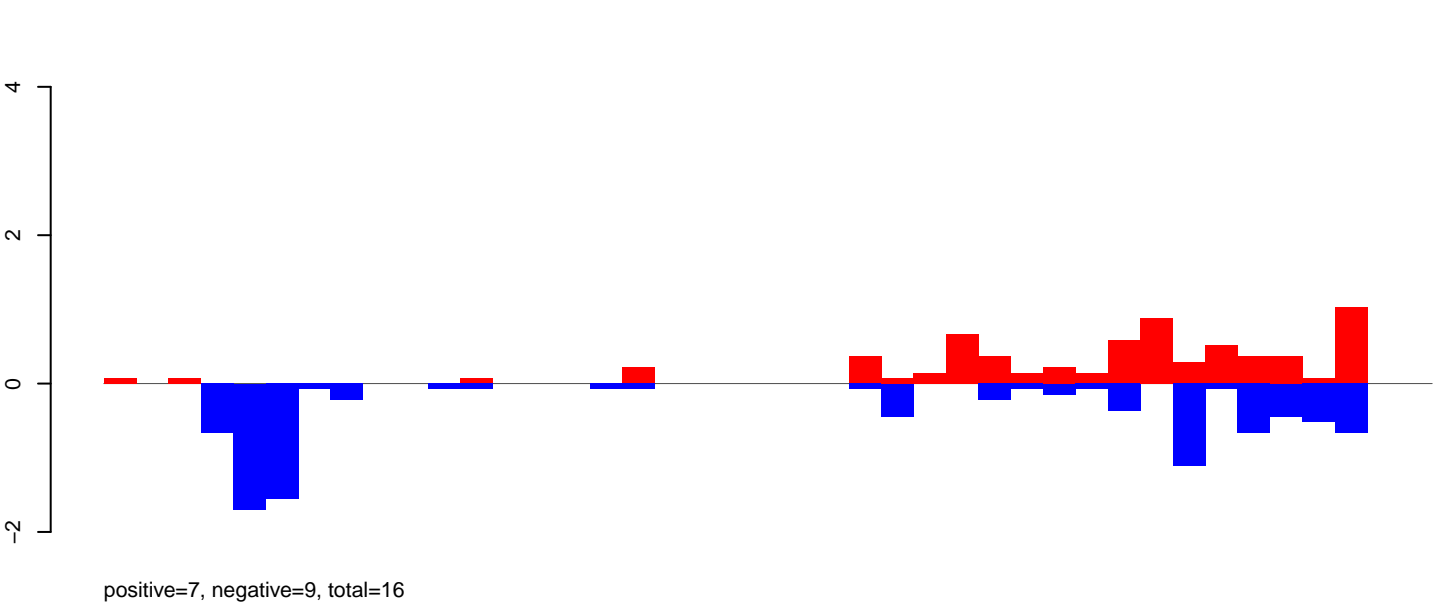
Window size=25, length=1350, TE@rnd-4_family-17@DNA_PiggyBac@2733:1-1350



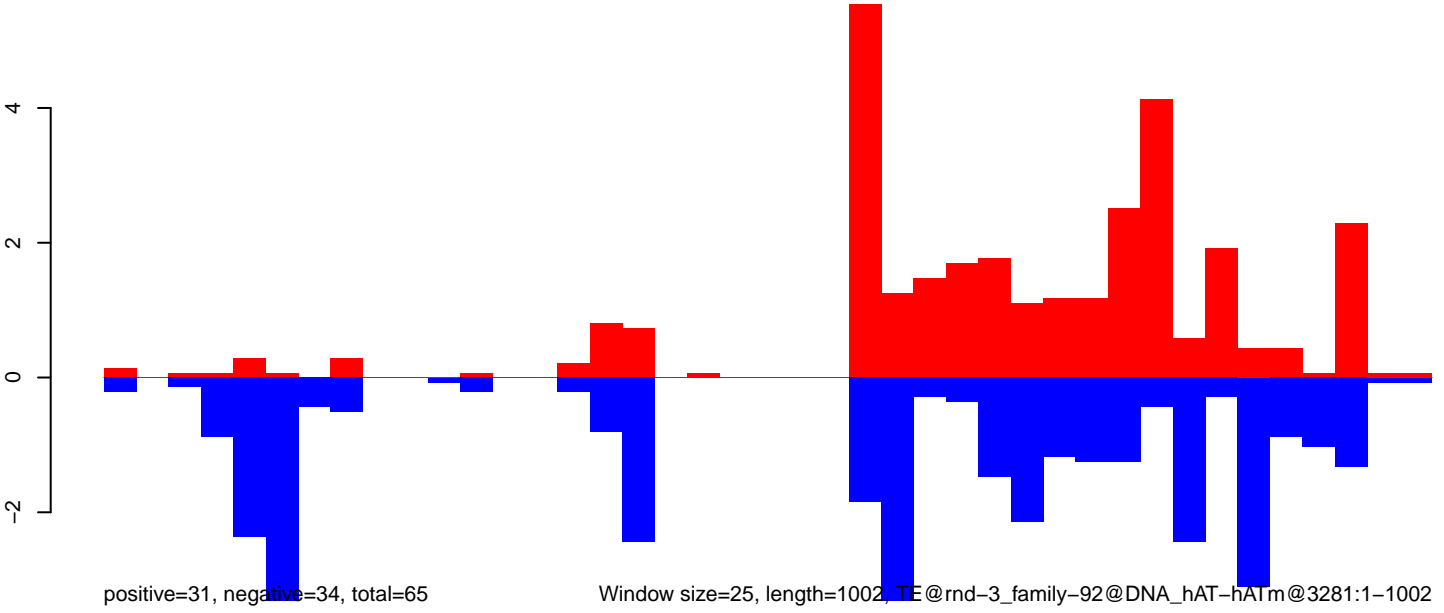
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



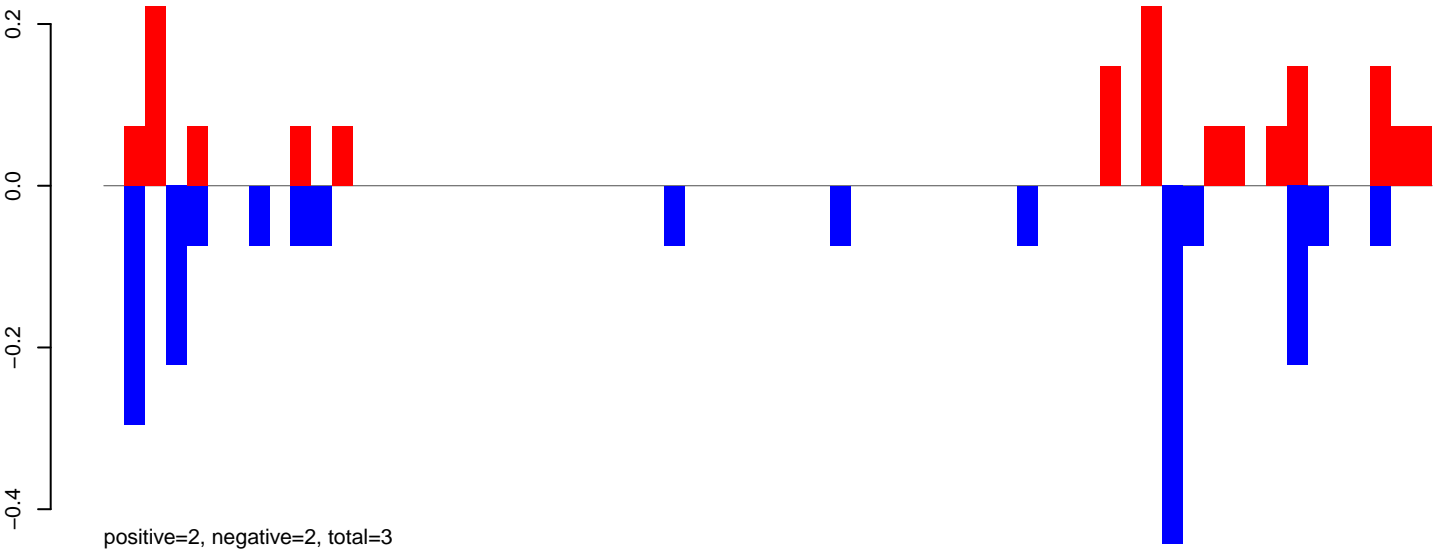
CuQuin_Testes_NL.rep



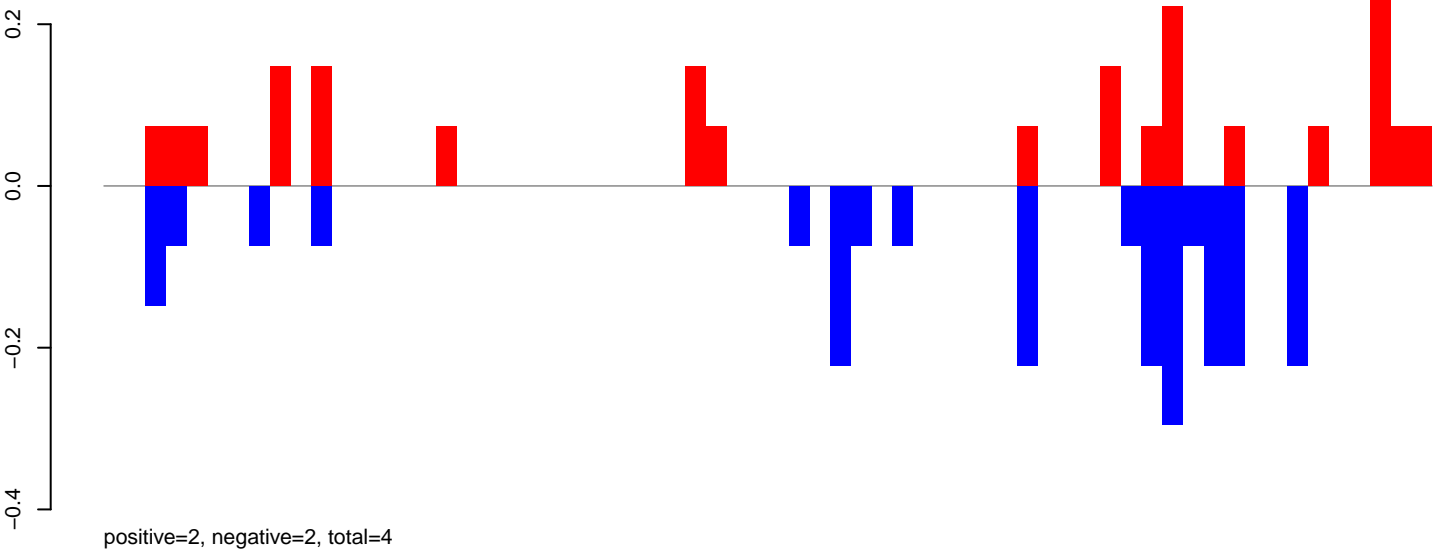
Window size=25, length=1002, TE@rnd-3_family-92@DNA_hAT-hATm@3281:1-1002

0 200 400 600 800 1000

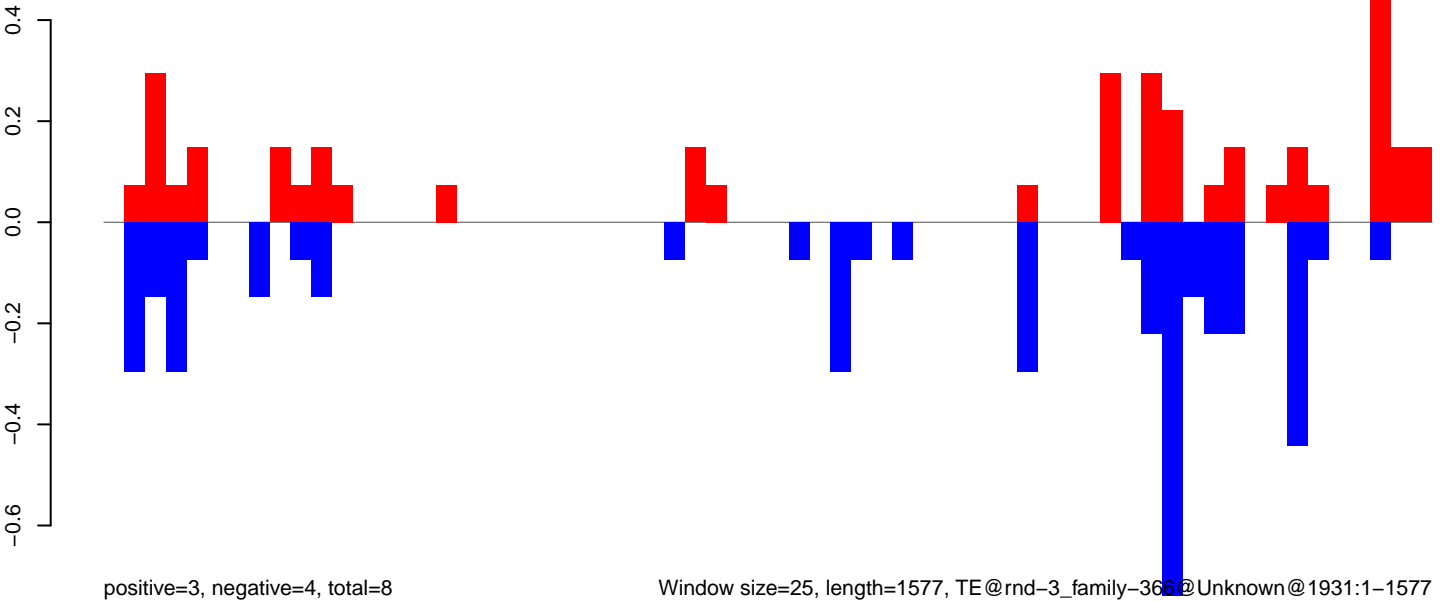
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

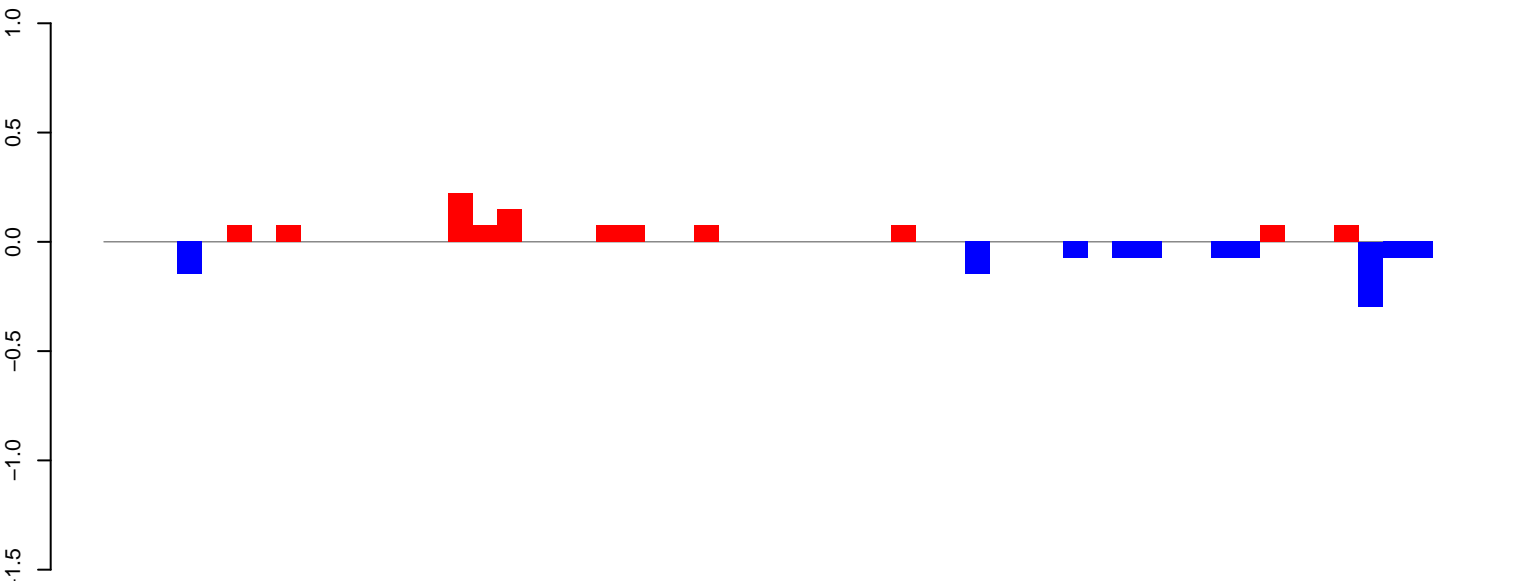


CuQuin_Testes_NL.rep



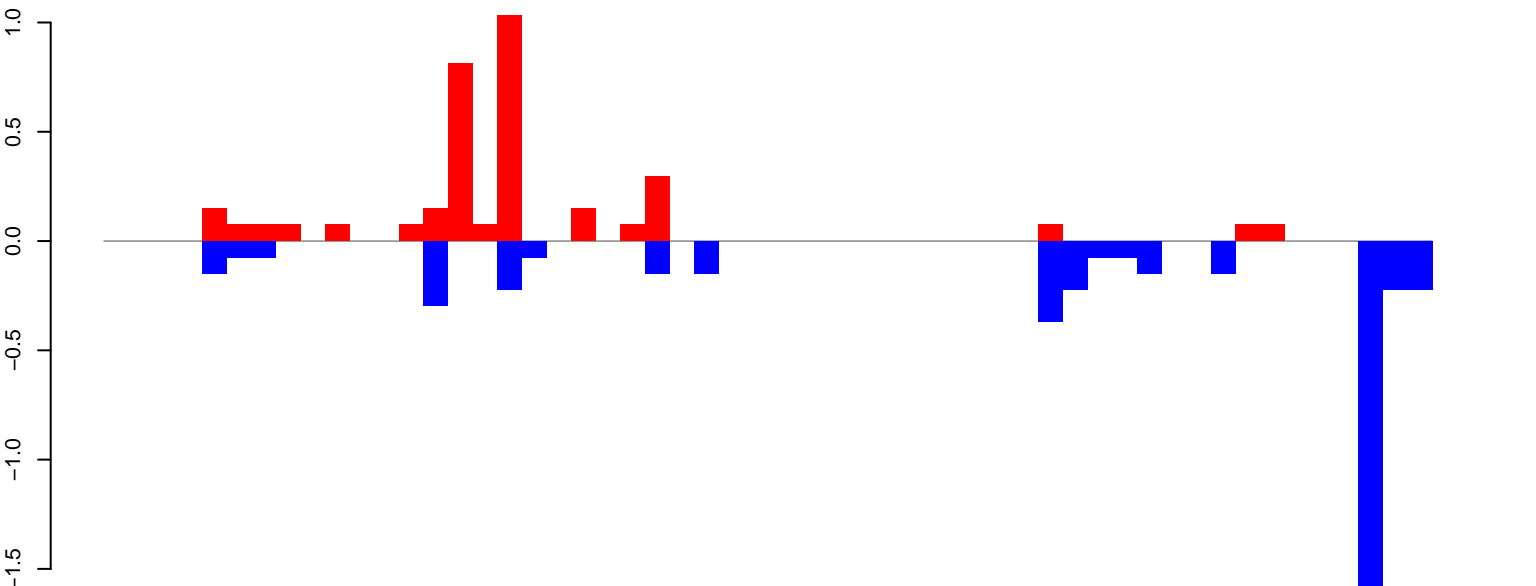
Window size=25, length=1577, TE@rnd-3_family-366@Unknown@1931:1-1577

CuQuin_Testes_NL.18_23.rep



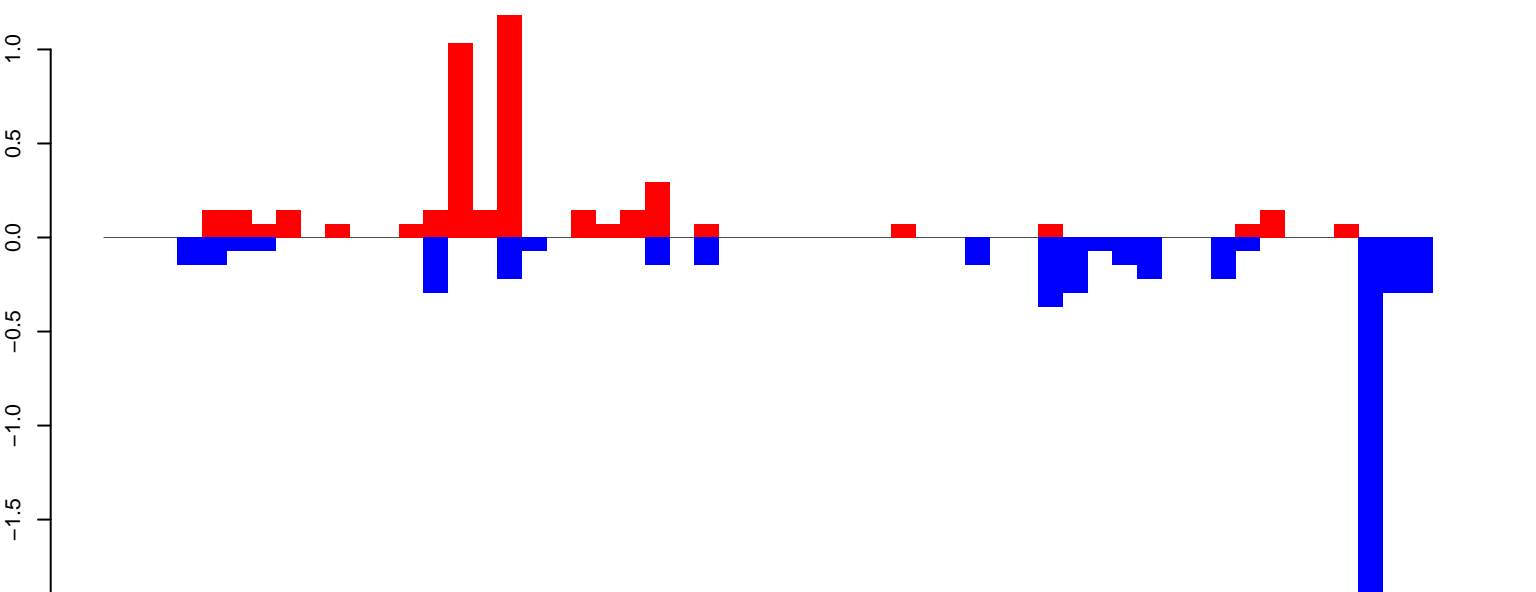
positive=1, negative=1, total=2

CuQuin_Testes_NL.24_35.rep



positive=3, negative=4, total=8

CuQuin_Testes_NL.rep

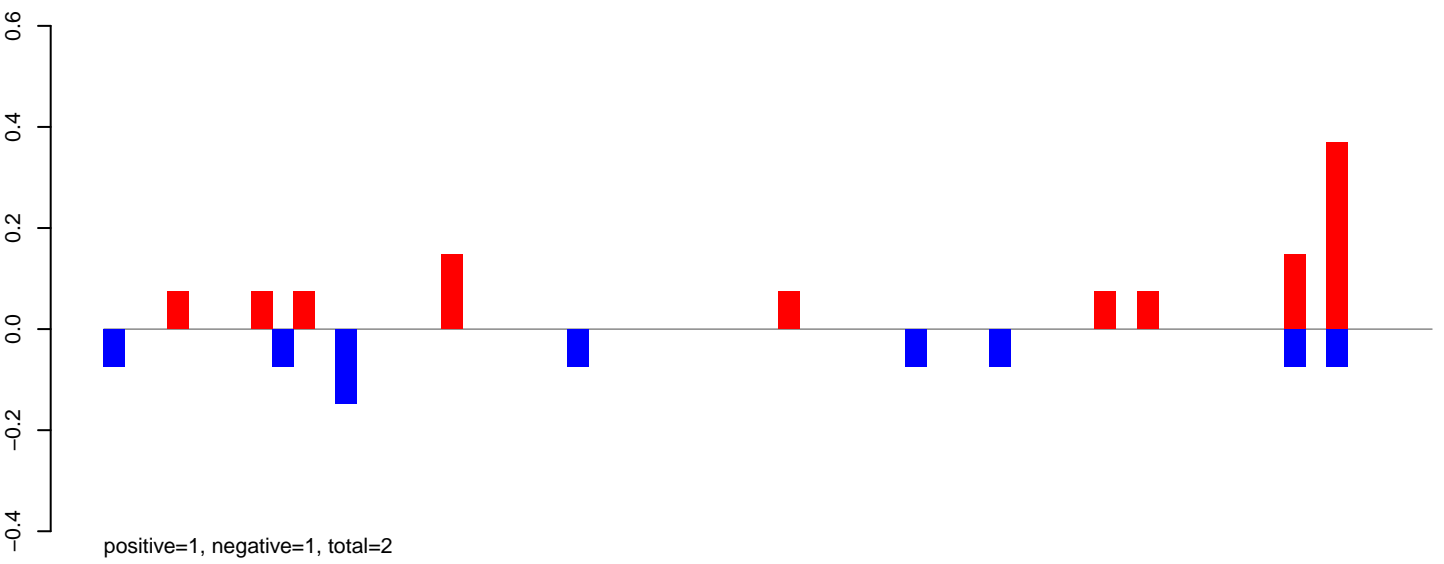


positive=4, negative=5, total=10

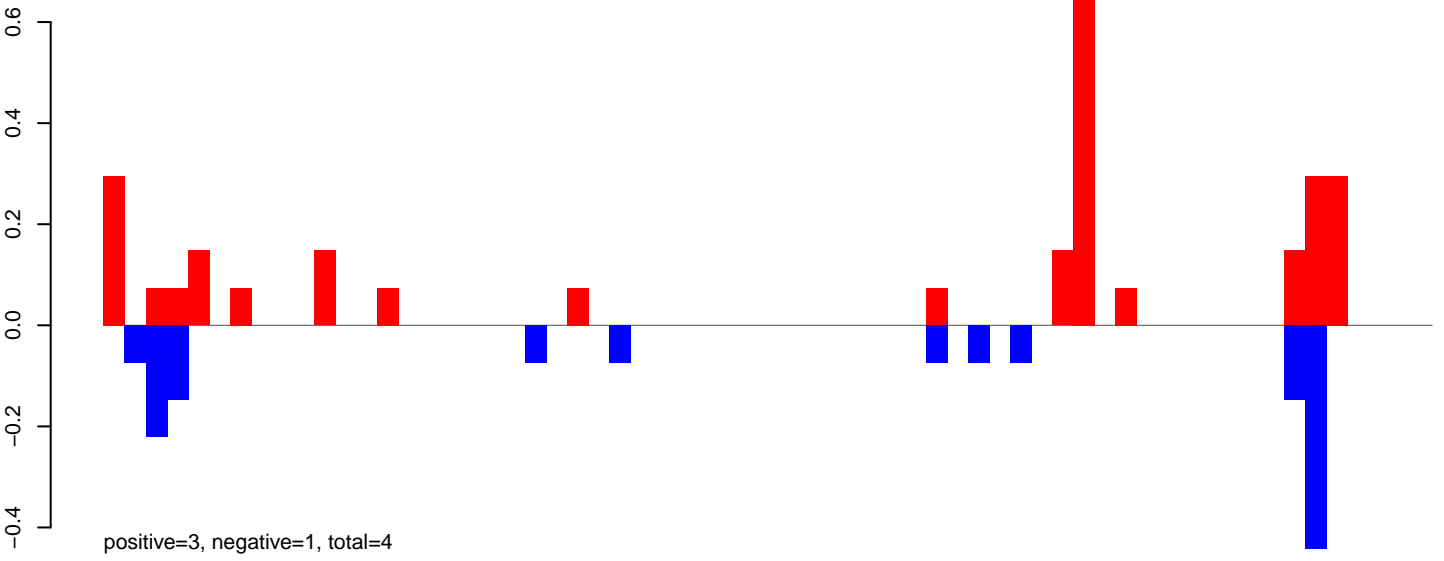
Window size=25, length=1347, TE@rnd-3_family-197@DNA_hAT-Charlie@941-1347

0 200 400 600 800 1000 1200 1400

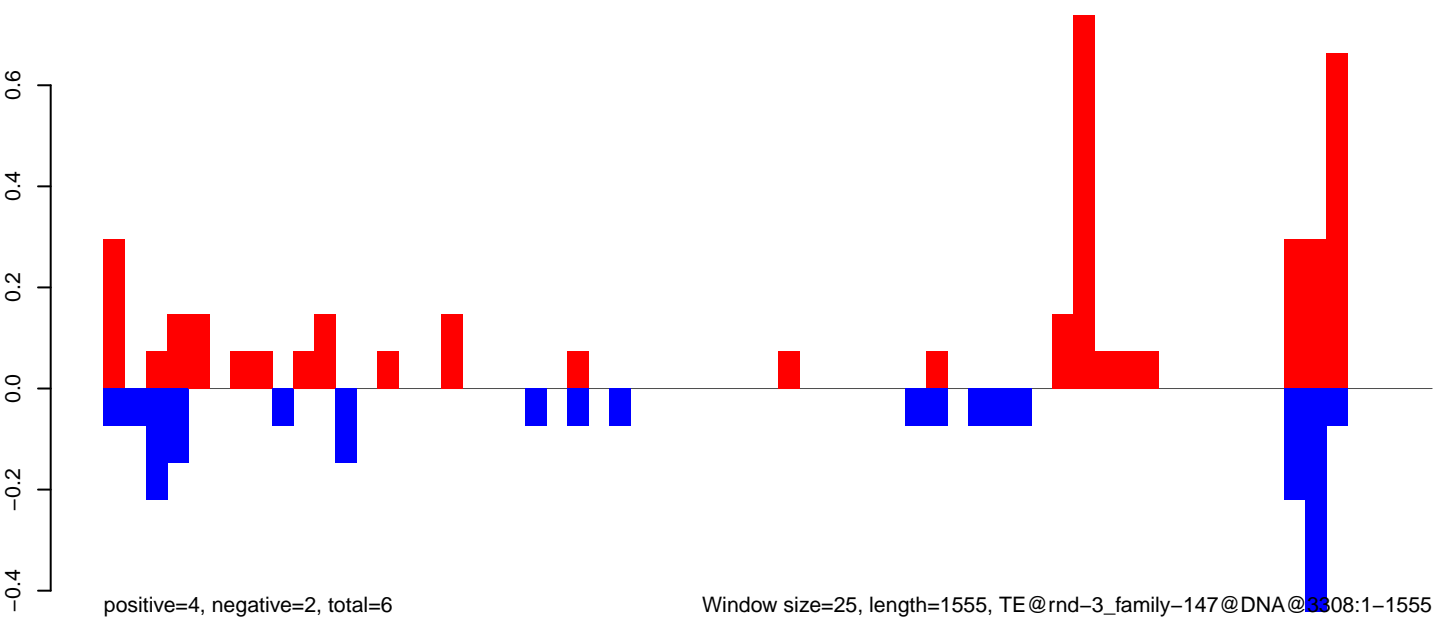
CuQuin_Testes_NL.18_23.rep



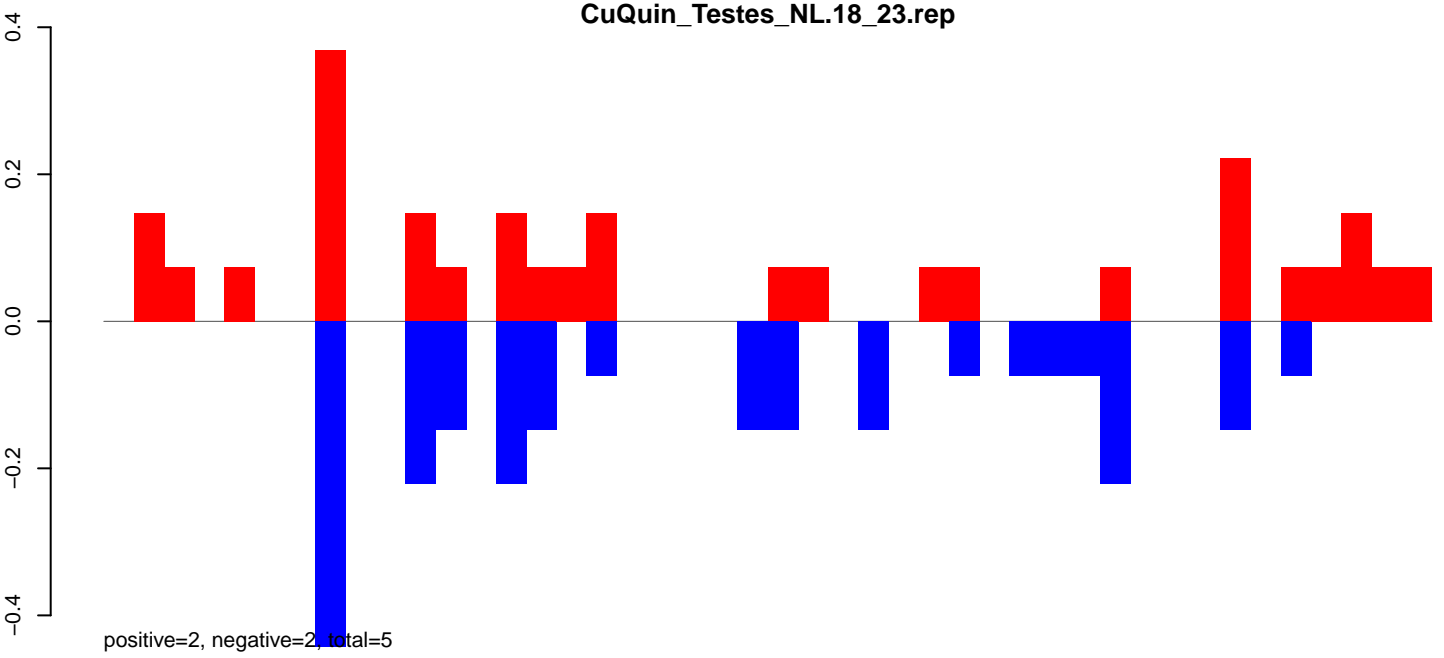
CuQuin_Testes_NL.24_35.rep



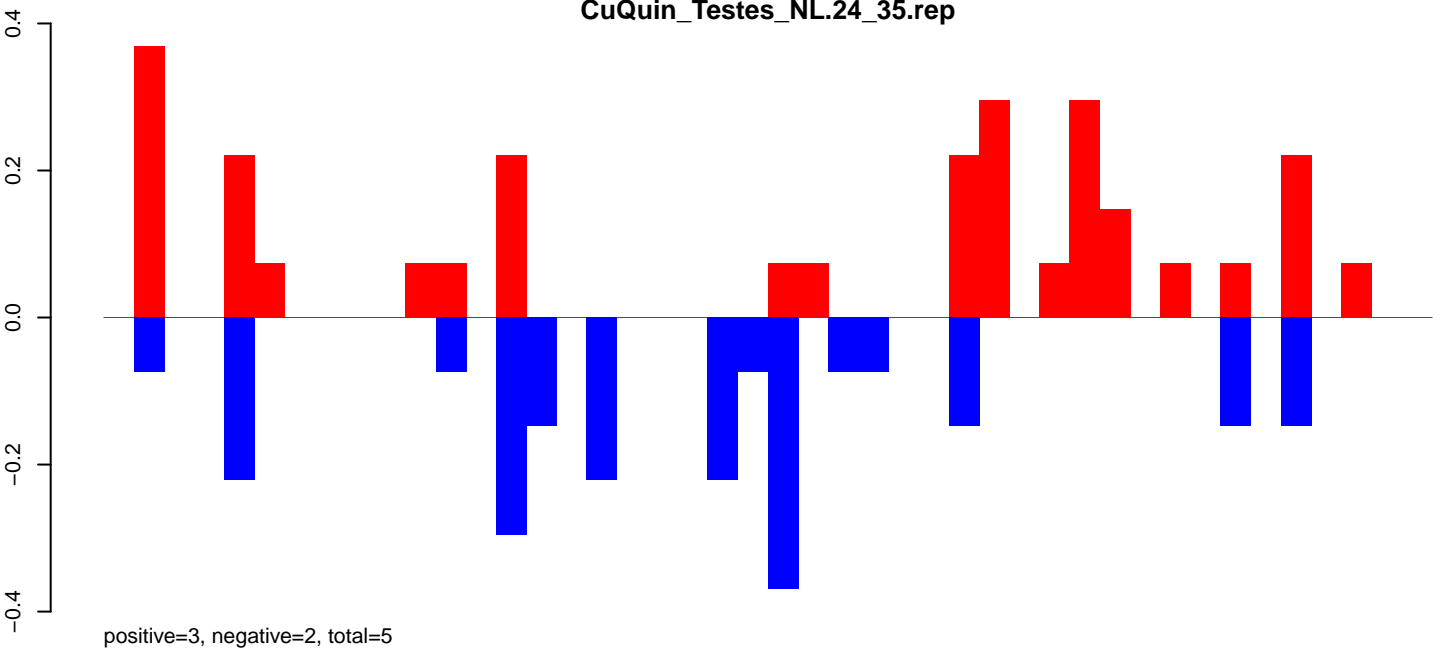
CuQuin_Testes_NL.rep



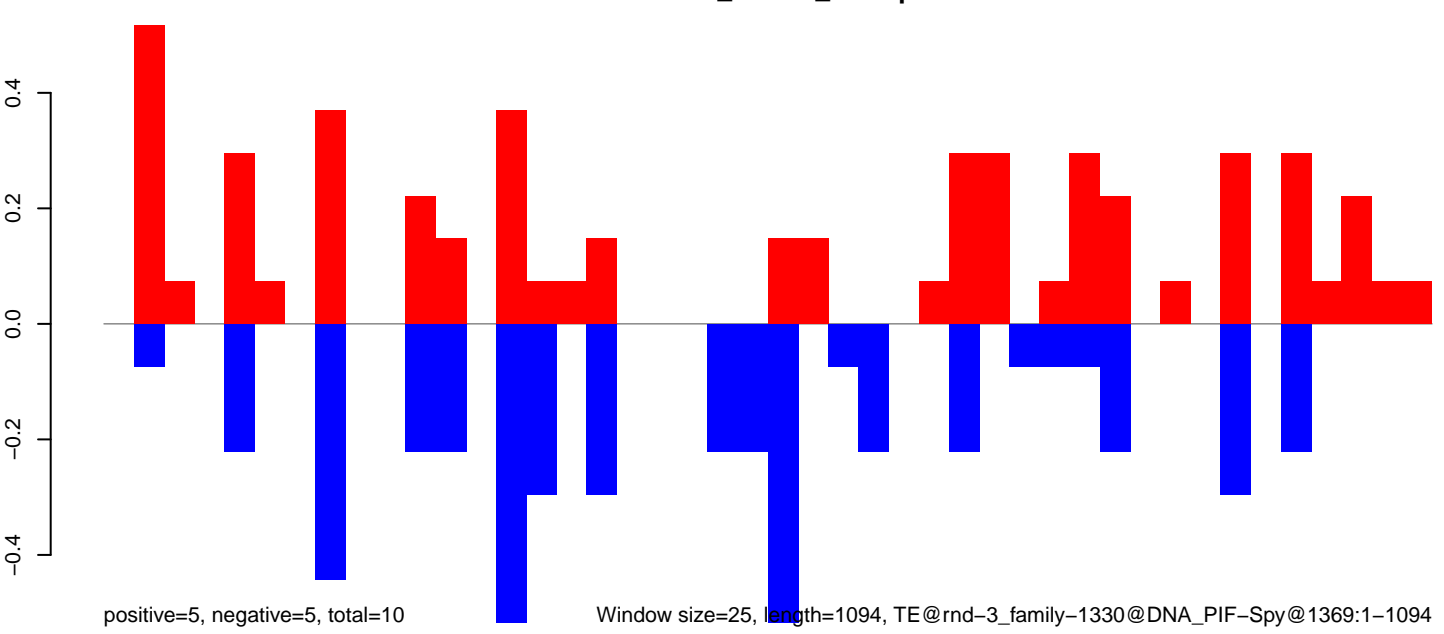
CuQuin_Testes_NL.18_23.rep



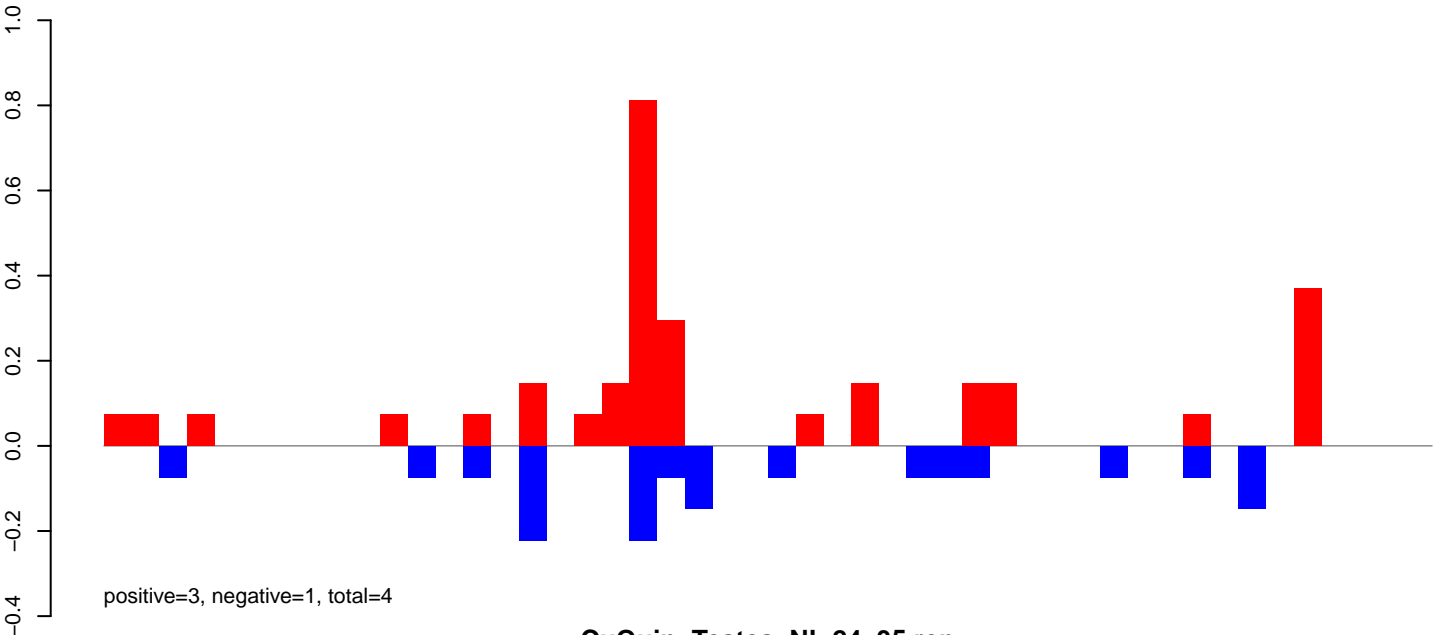
CuQuin_Testes_NL.24_35.rep



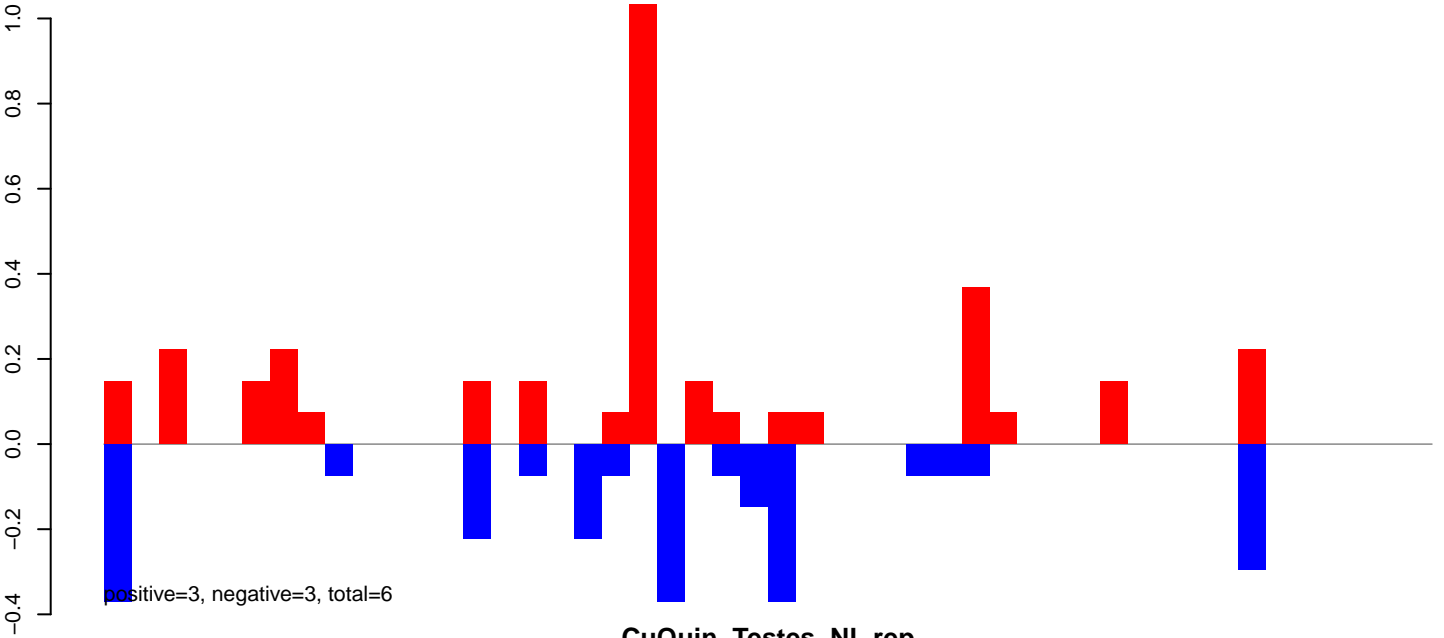
CuQuin_Testes_NL.rep



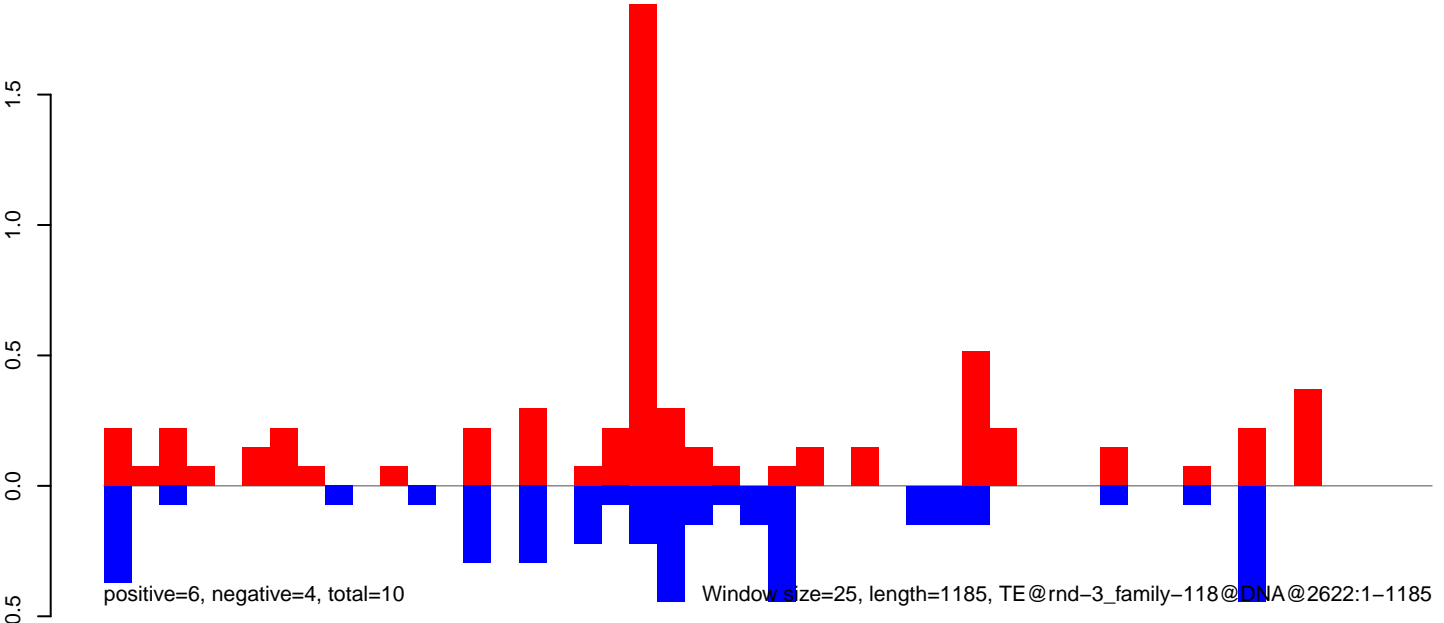
CuQuin_Testes_NL.18_23.rep



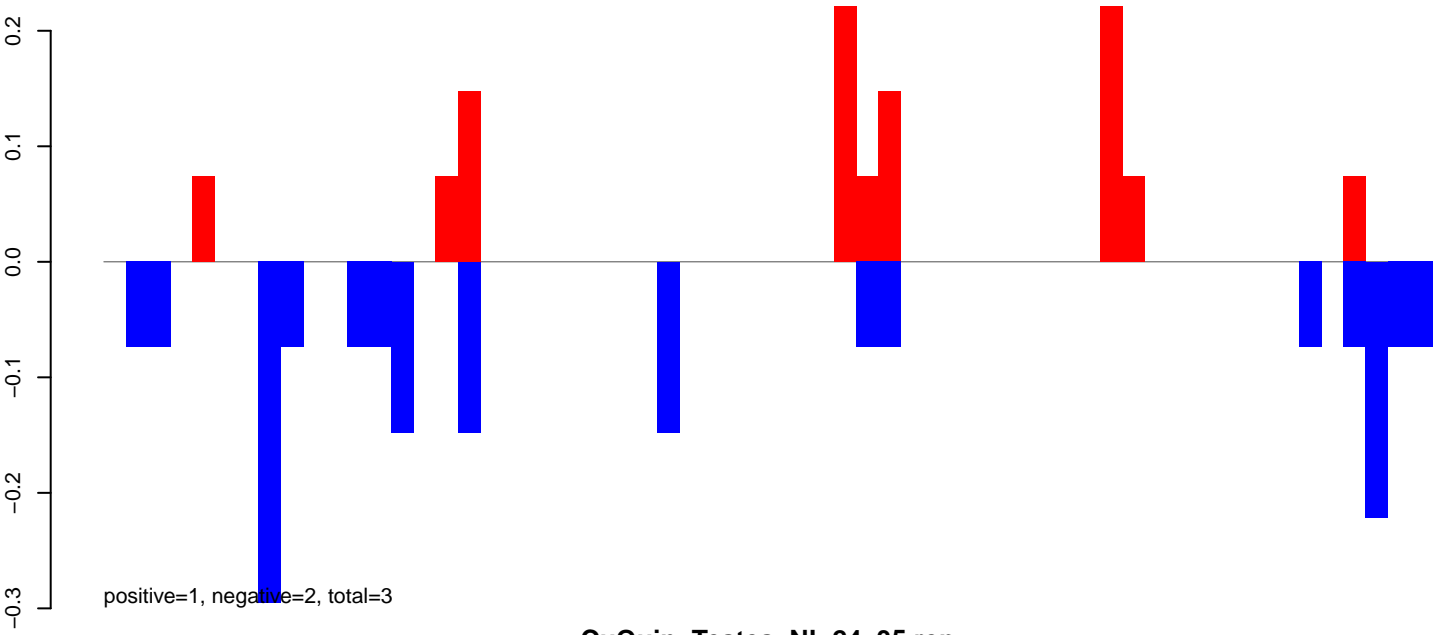
CuQuin_Testes_NL.24_35.rep



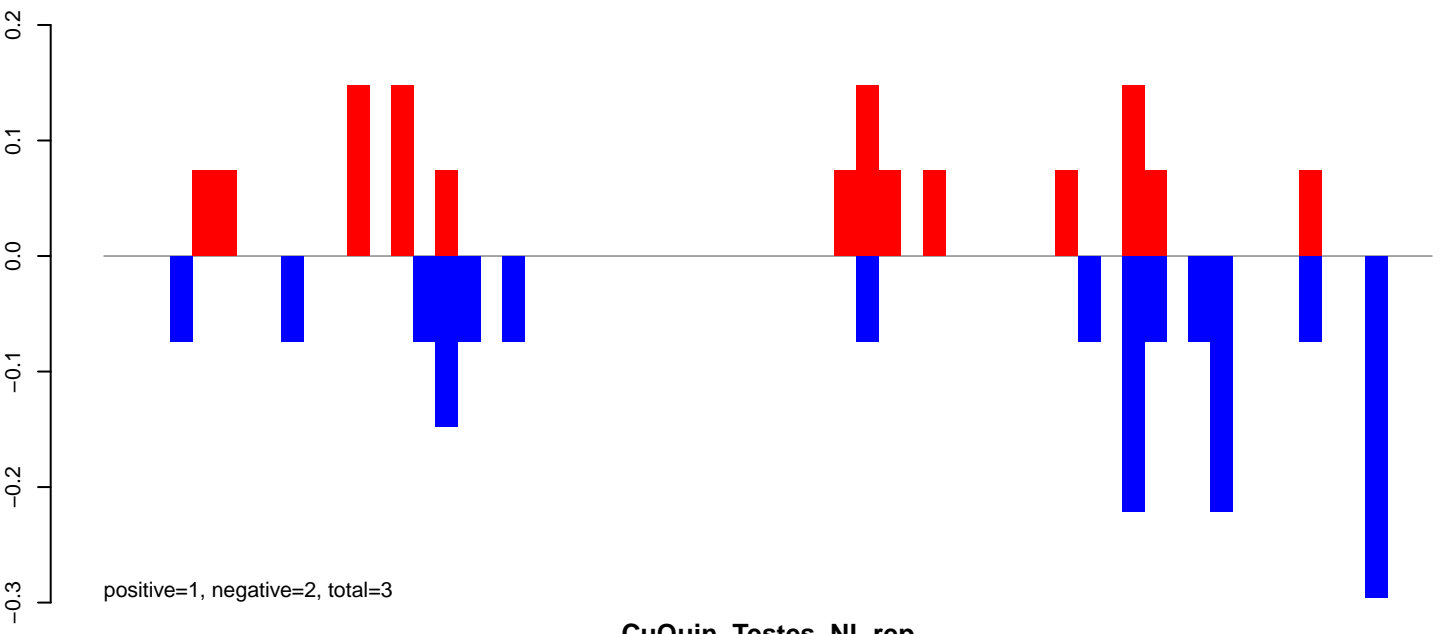
CuQuin_Testes_NL.rep



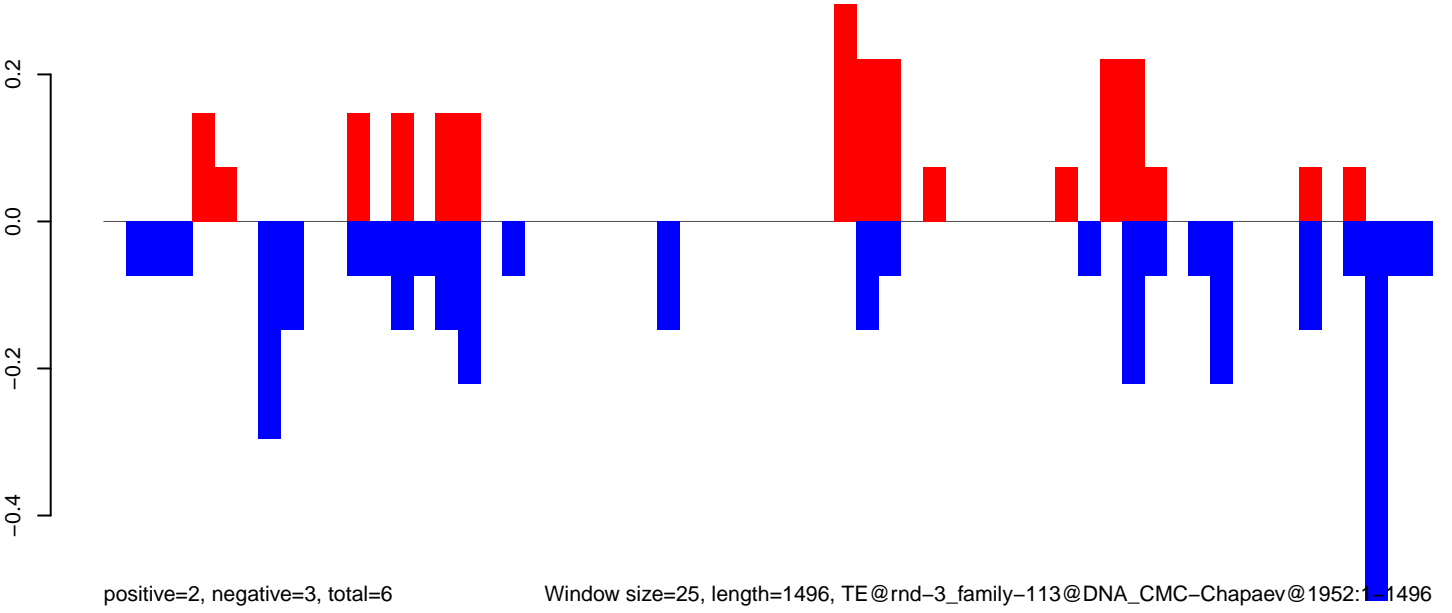
CuQuin_Testes_NL.18_23.rep



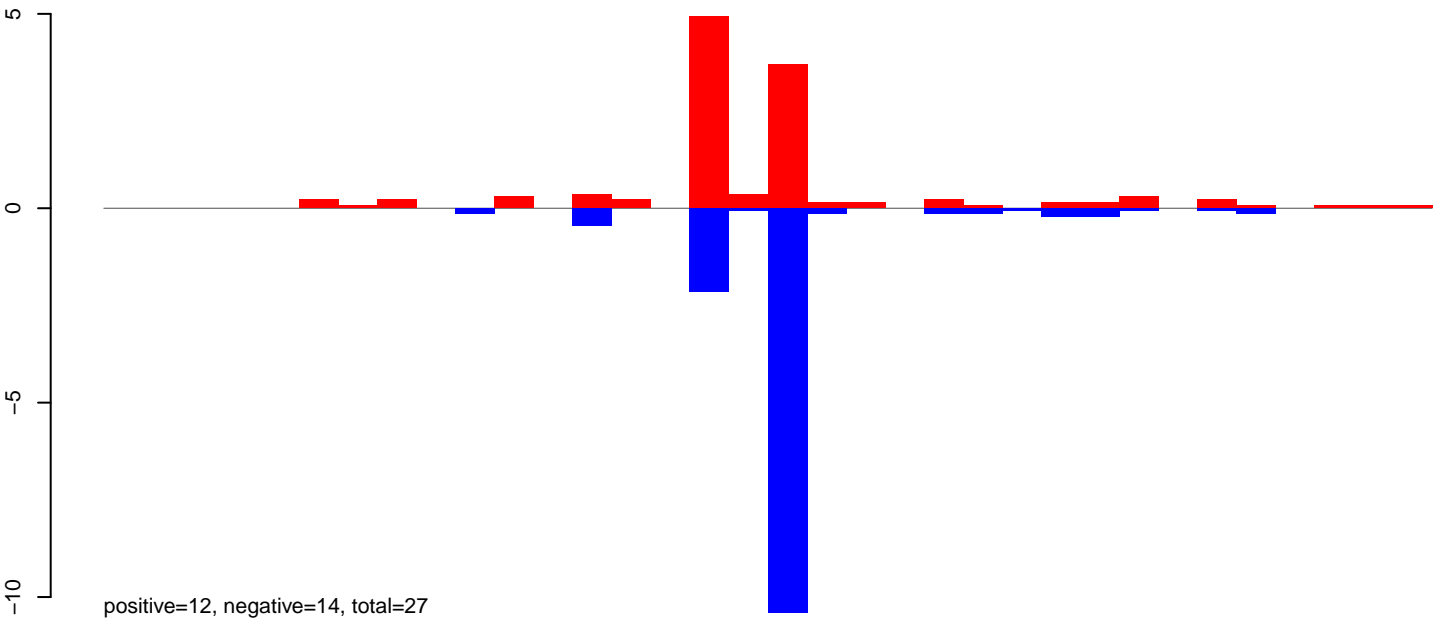
CuQuin_Testes_NL.24_35.rep



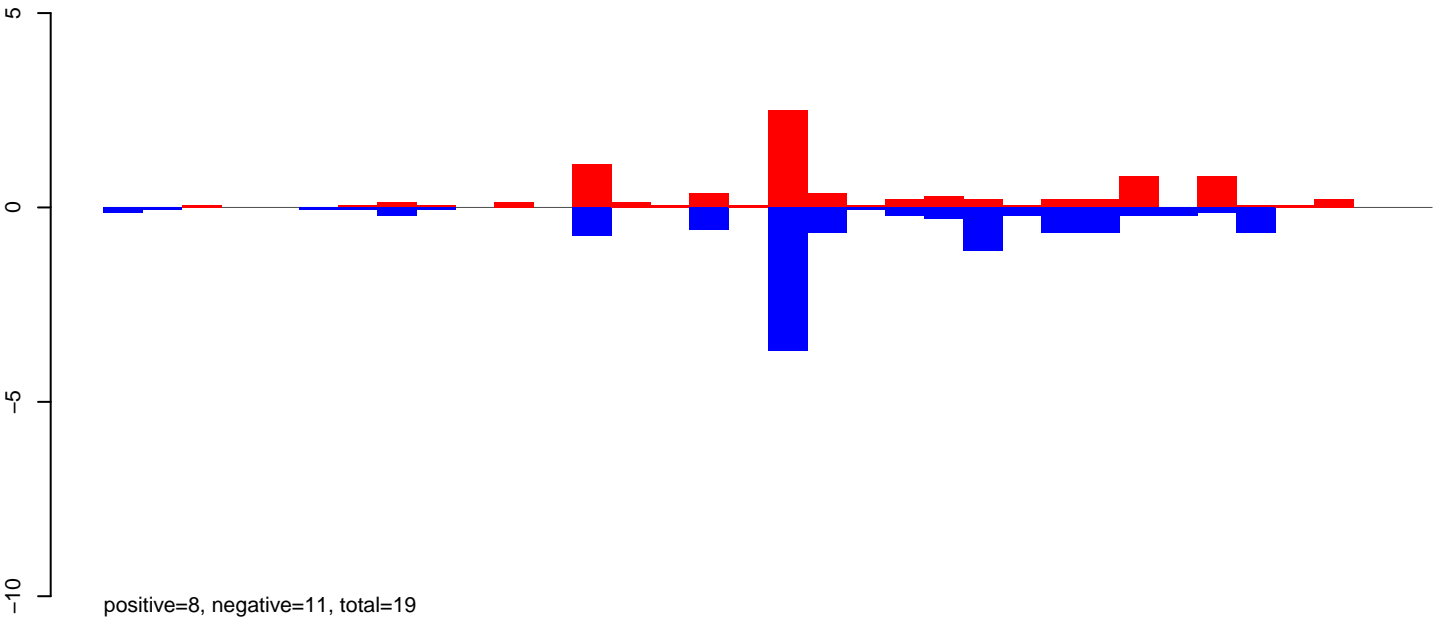
CuQuin_Testes_NL.rep



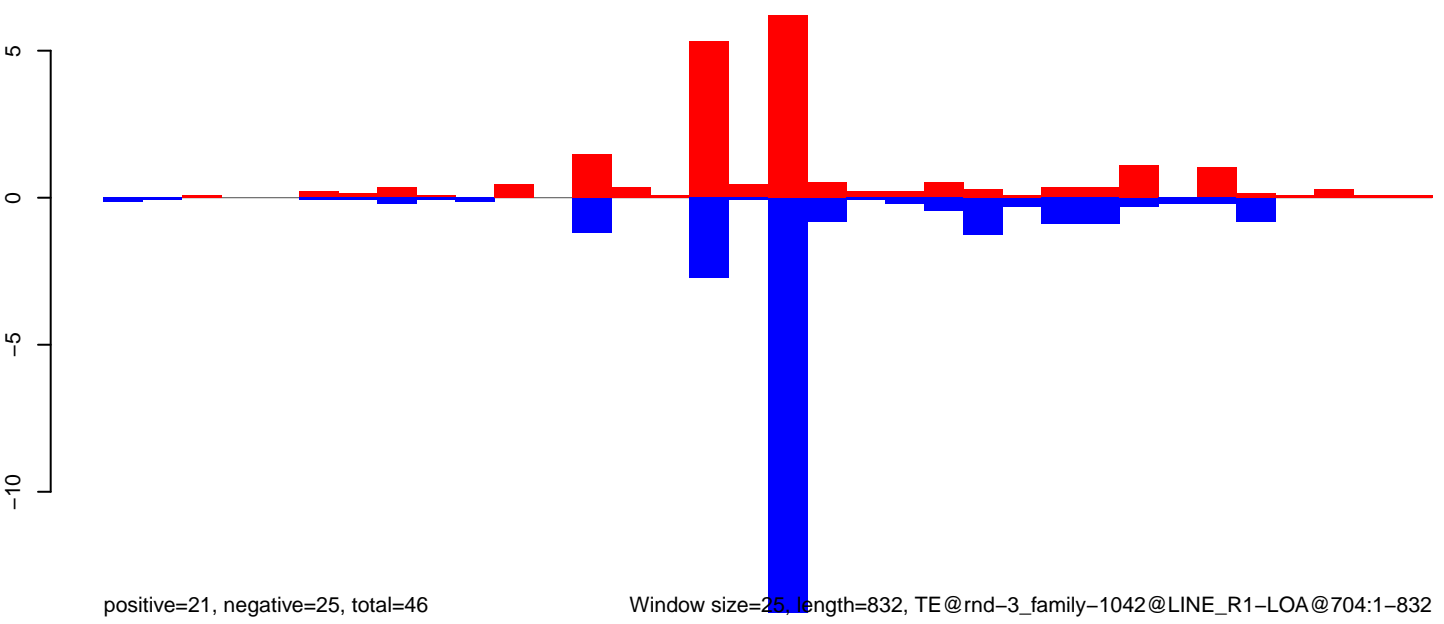
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



Window size=25, length=832, TE@rnd-3_family-1042@LINE_R1-LOA@704:1-832

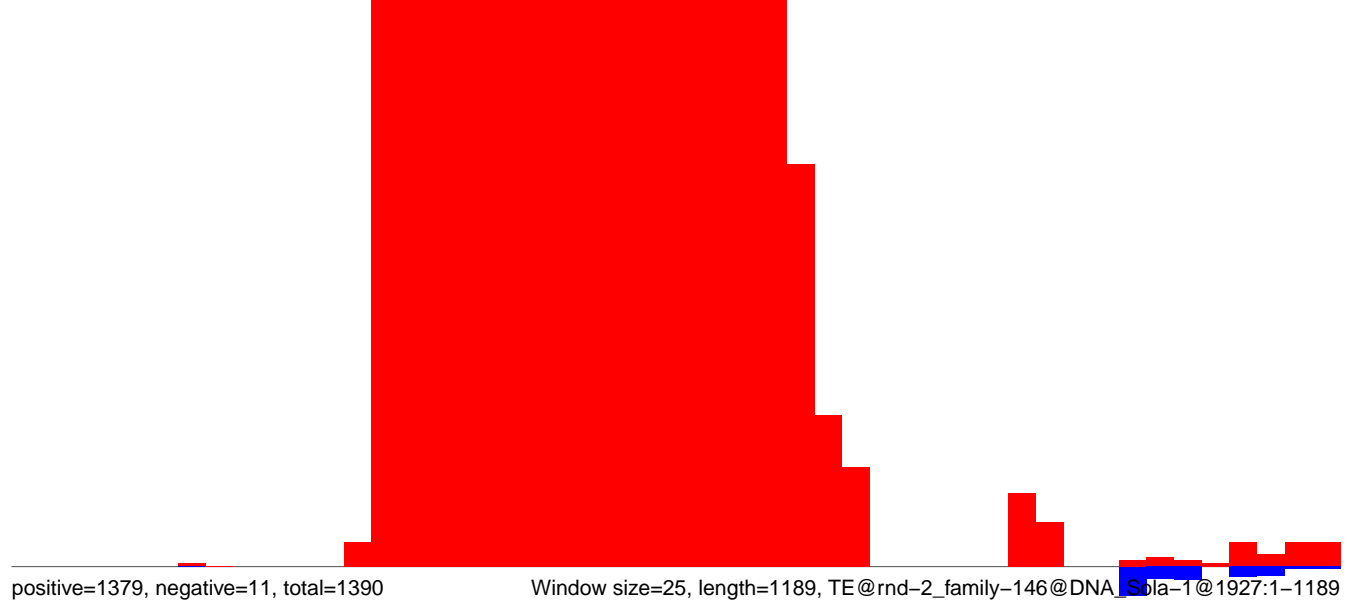
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



CuQuin_Testes_NL.18_23.rep

1.0
0.5
0.0
-0.5

positive=1, negative=2, total=4

CuQuin_Testes_NL.24_35.rep

1.0
0.5
0.0
-0.5

positive=7, negative=3, total=10

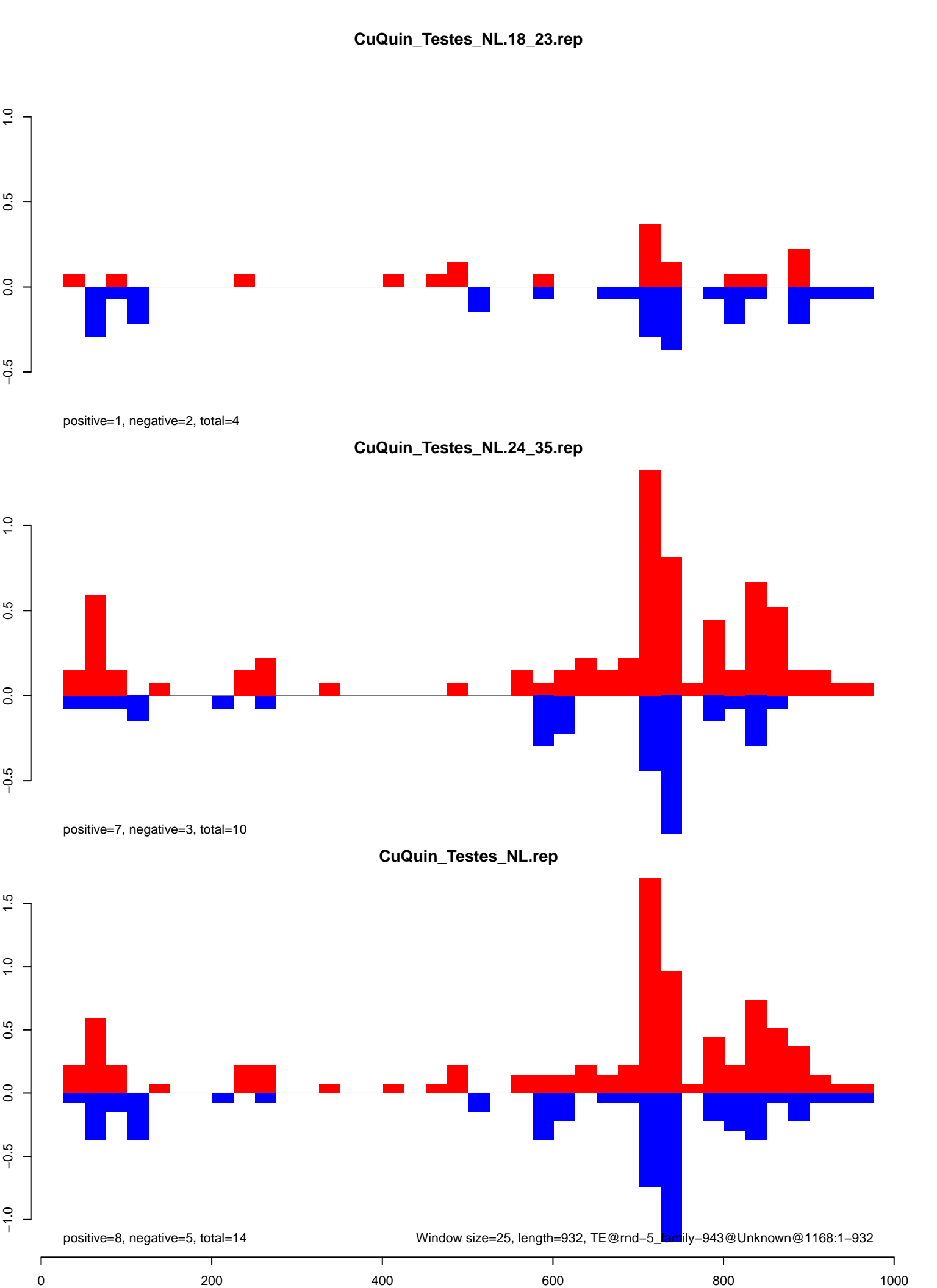
CuQuin_Testes_NL.rep

1.5
1.0
0.5
0.0
-0.5
-1.0

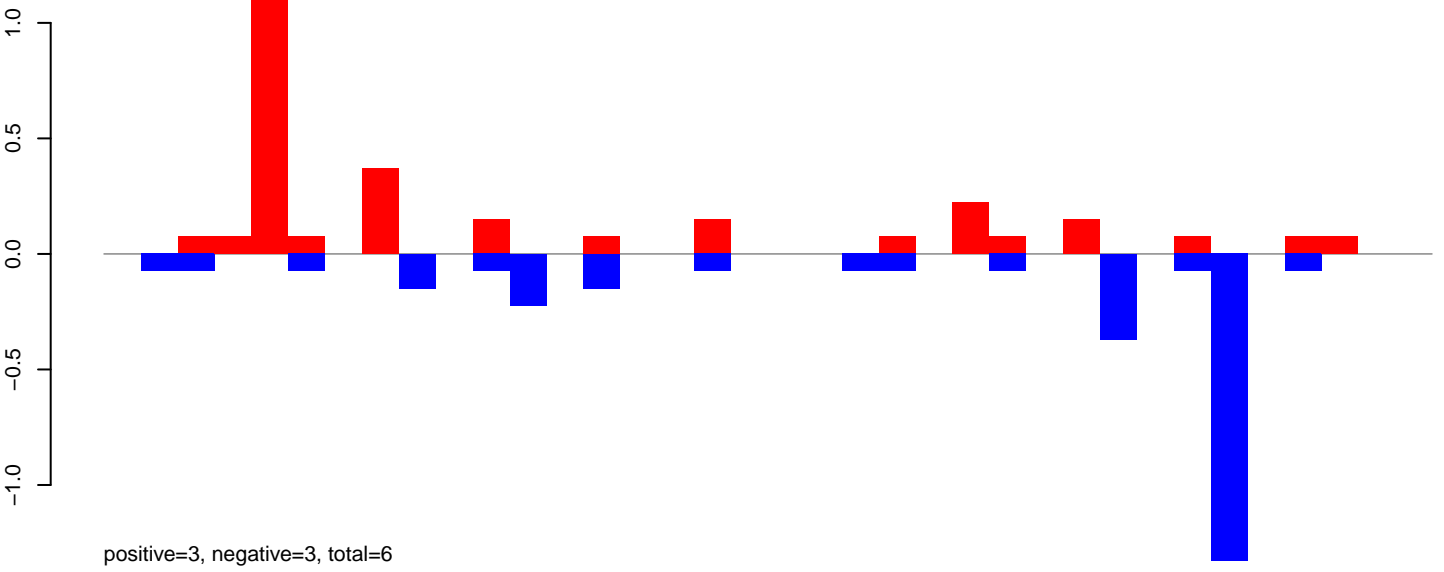
positive=8, negative=5, total=14

Window size=25, length=932, TE@rnd-5_family-943@Unknown@1168:1-932

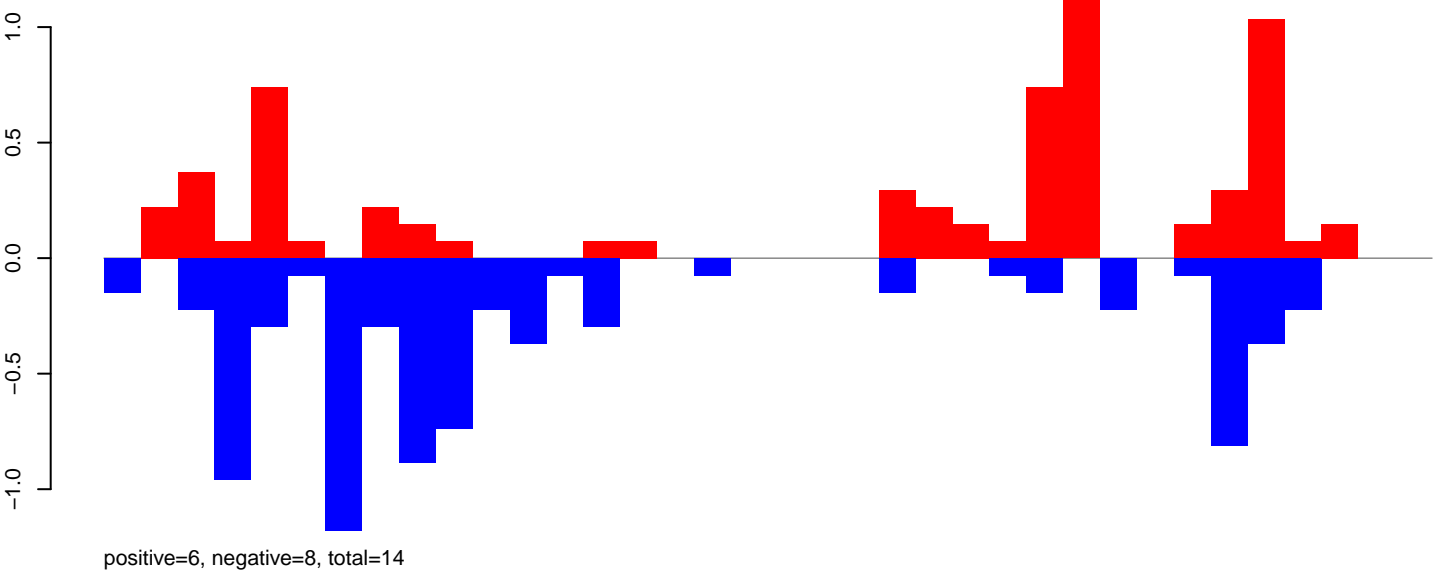
0 200 400 600 800 1000



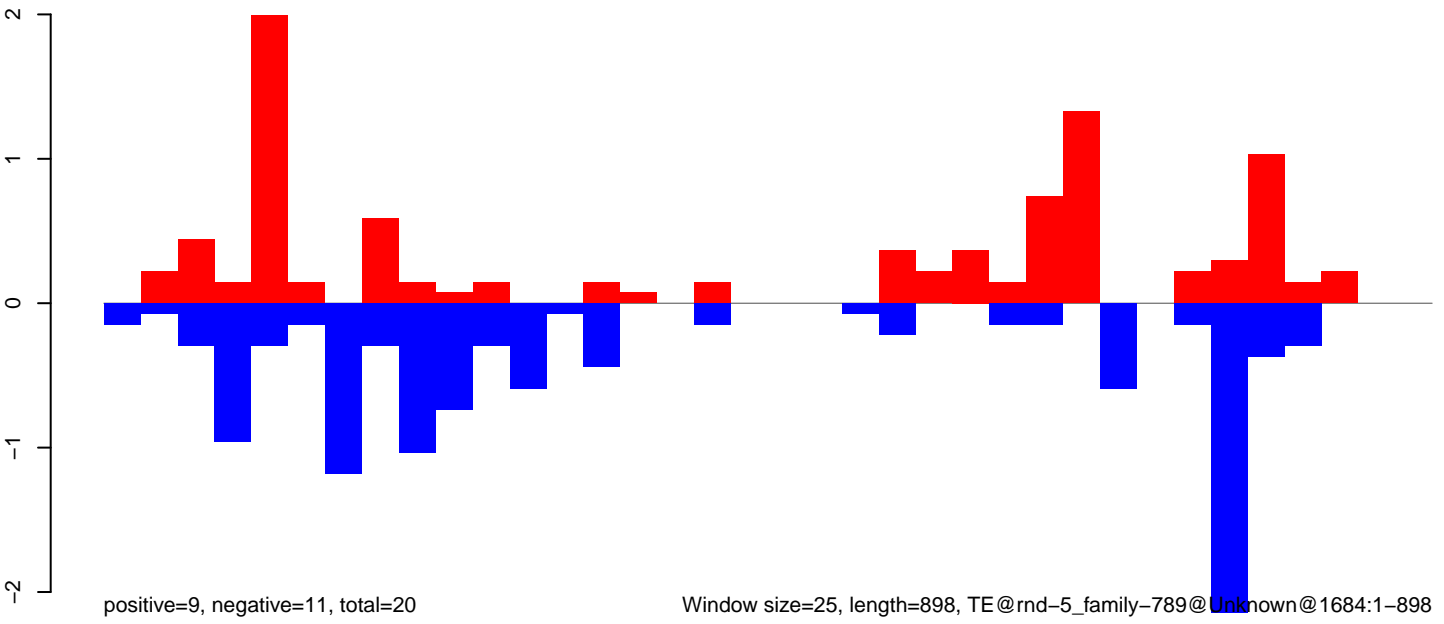
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



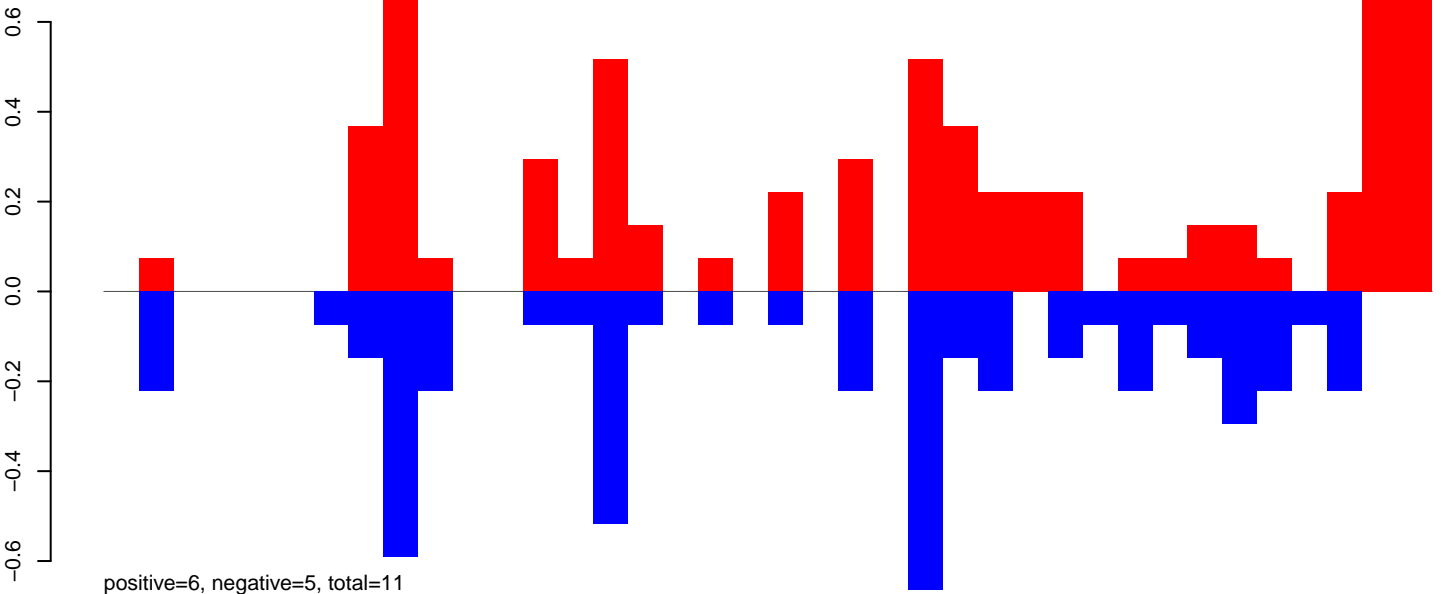
CuQuin_Testes_NL.rep



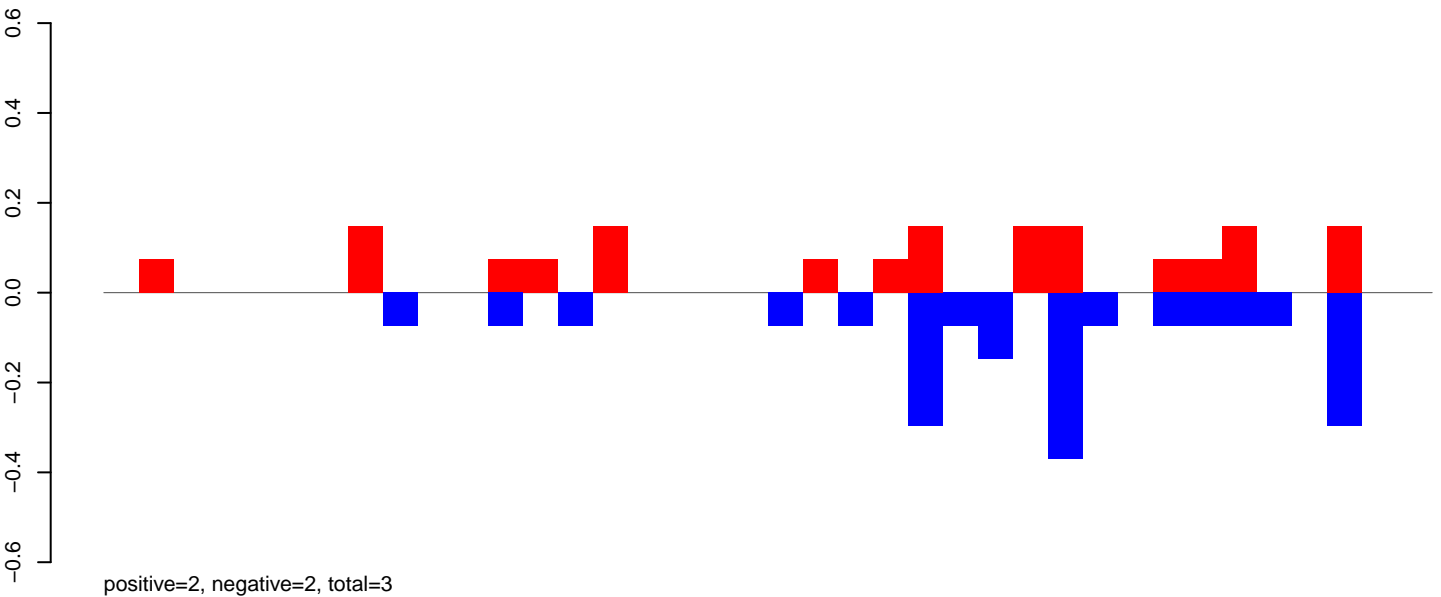
Window size=25, length=898, TE@rnd-5_family-789@Unknown@1684:1-898

0 200 400 600 800

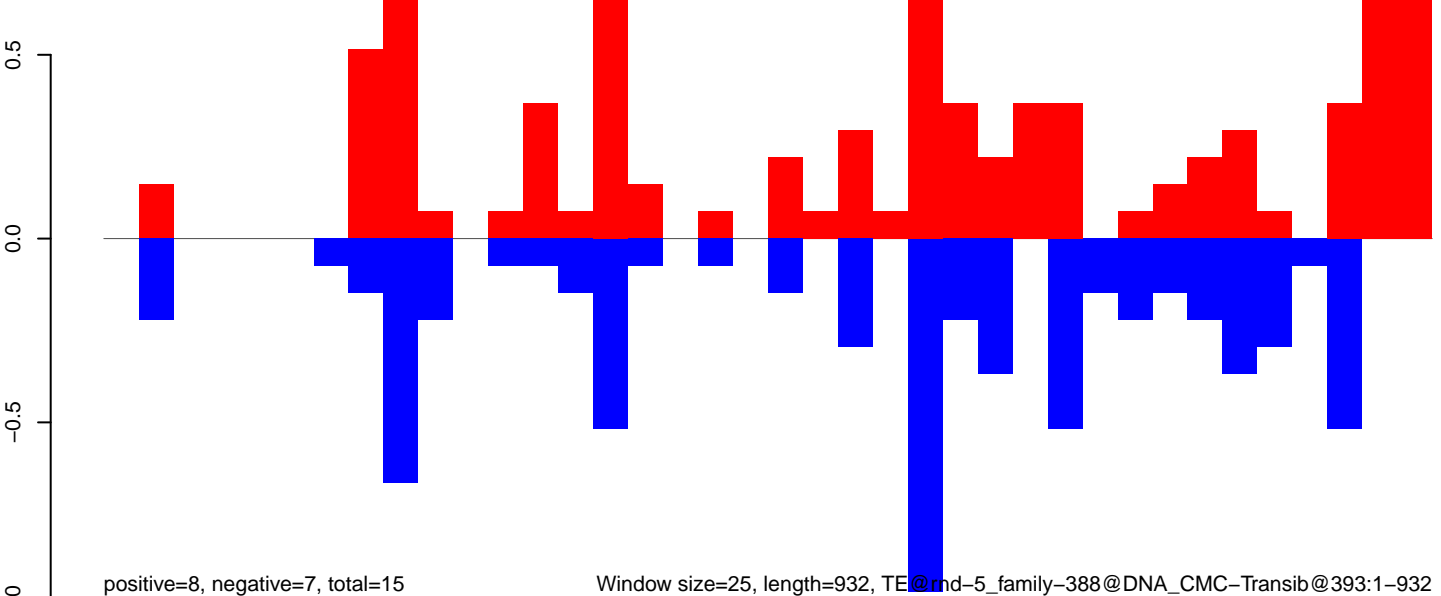
CuQuin_Testes_NL.18_23.rep



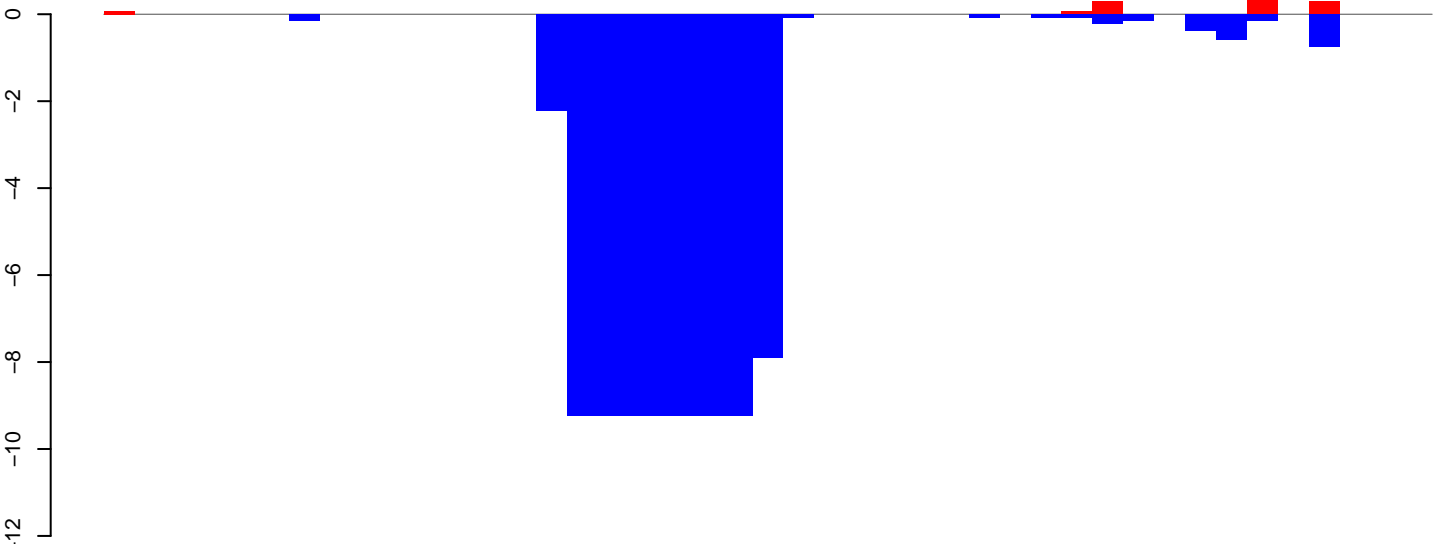
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

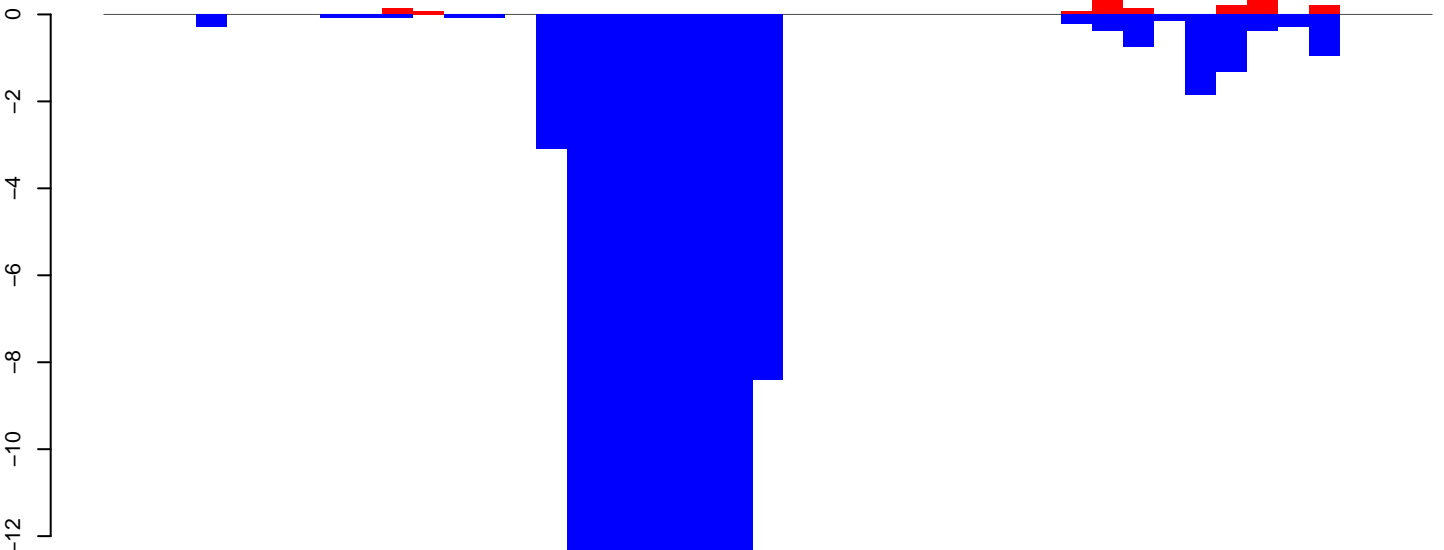


CuQuin_Testes_NL.18_23.rep



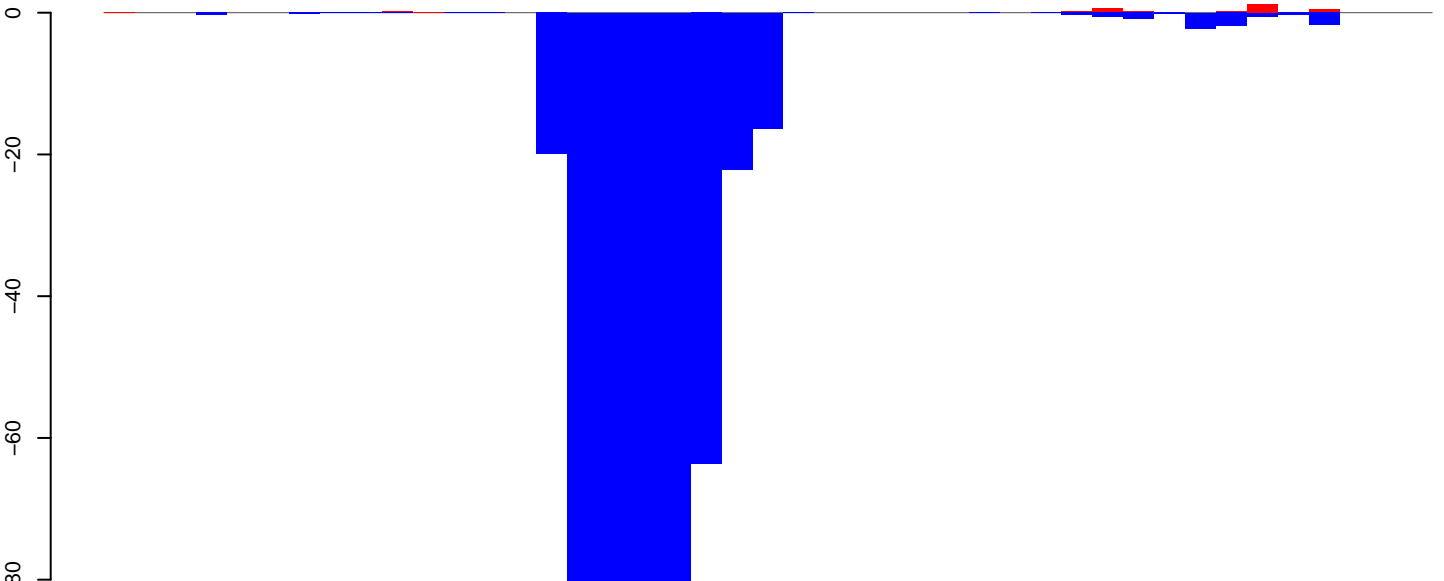
positive=2, negative=68, total=70

CuQuin_Testes_NL.24_35.rep



positive=2, negative=96, total=98

CuQuin_Testes_NL.rep

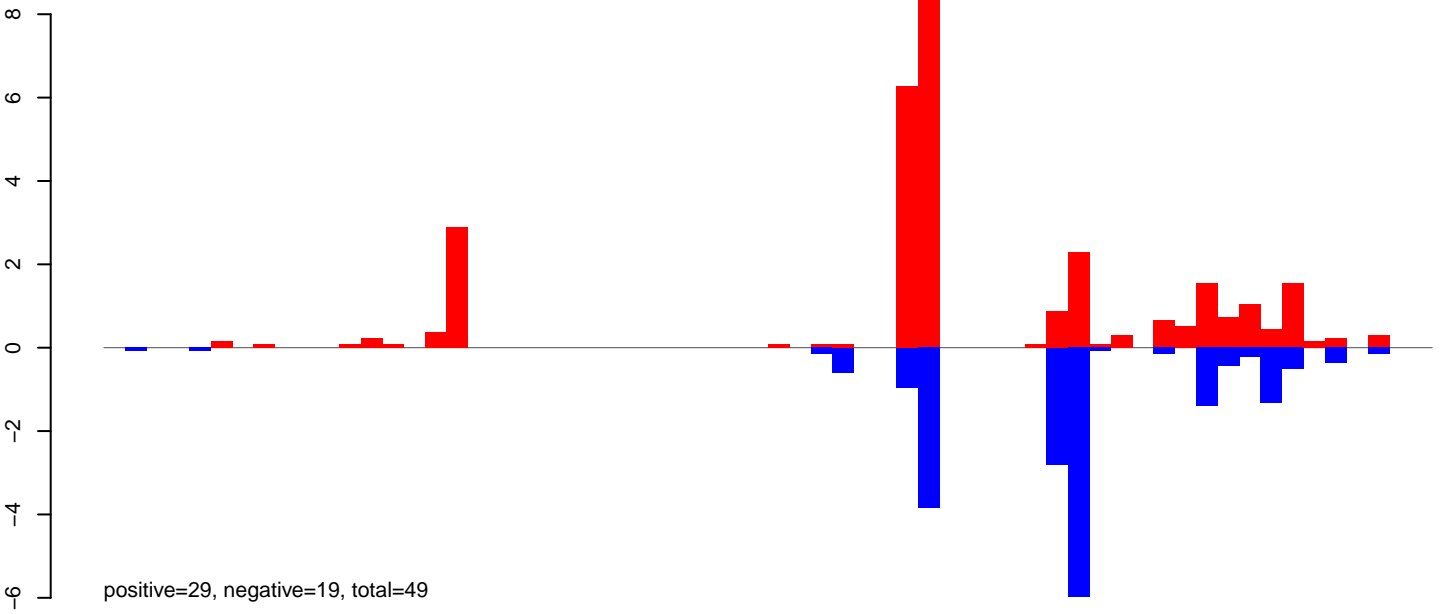


positive=3, negative=464, total=467

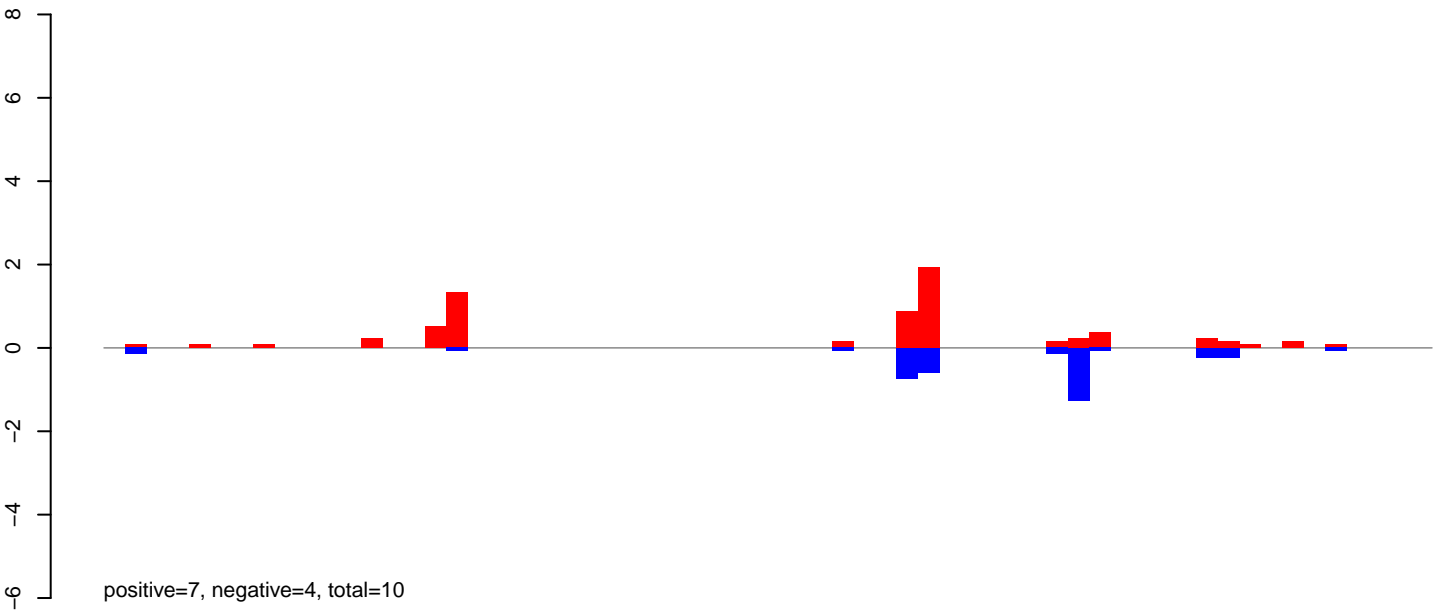
Window size=25, length=1063, TE@rnd-5_family-209@Unknown@3676:1-1063

0 200 400 600 800 1000

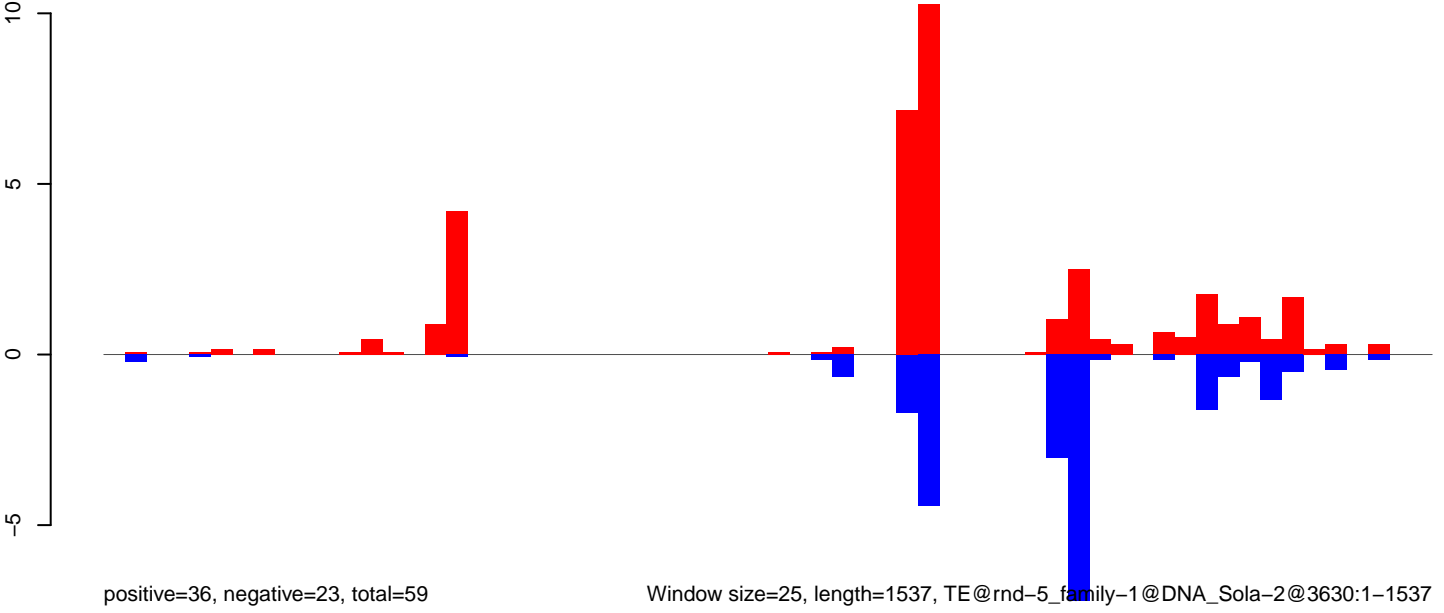
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

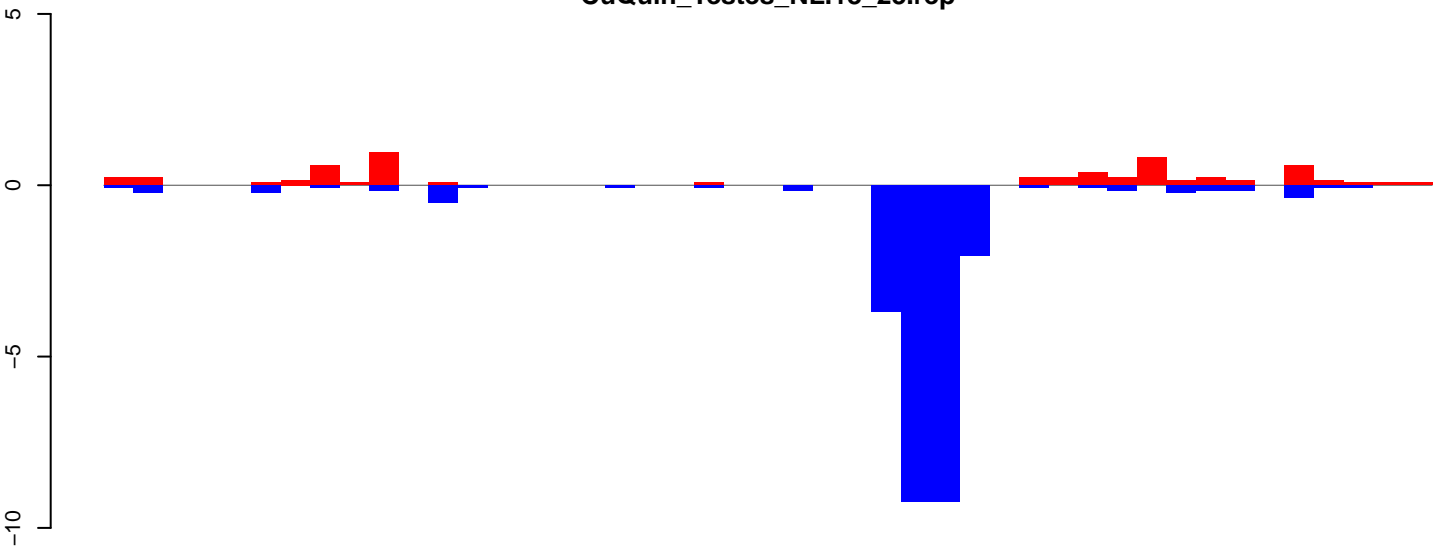


CuQuin_Testes_NL.rep



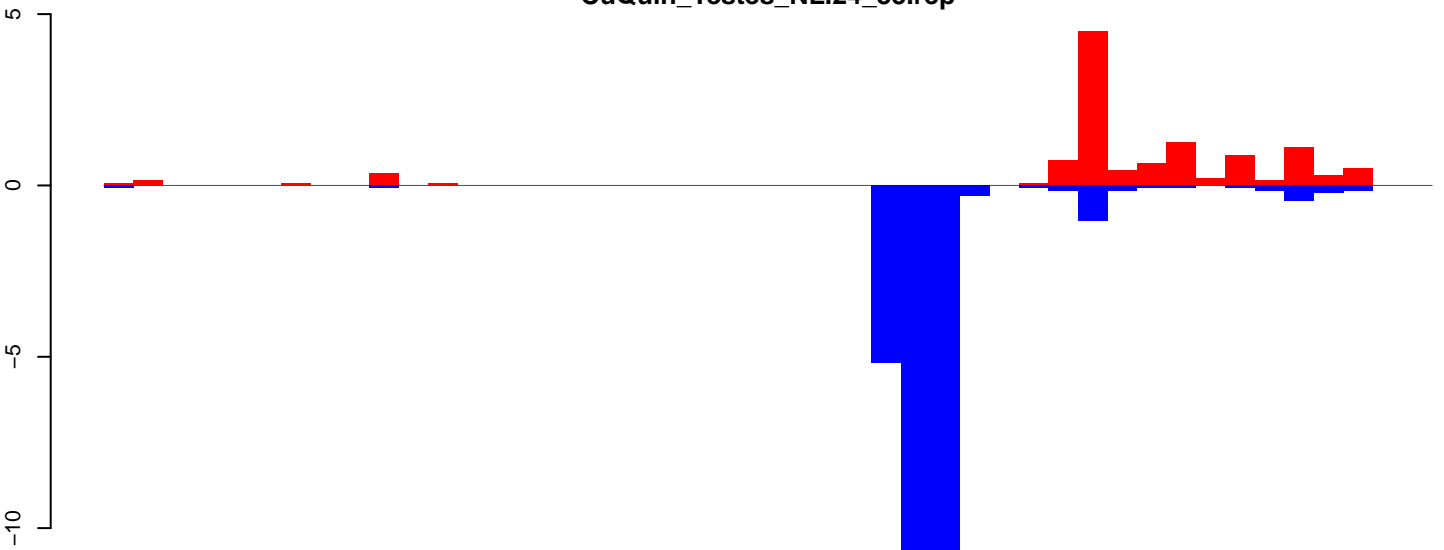
Window size=25, length=1537, TE@rnd-5_family-1@DNA_Sola-2@3630:1-1537

CuQuin_Testes_NL.18_23.rep



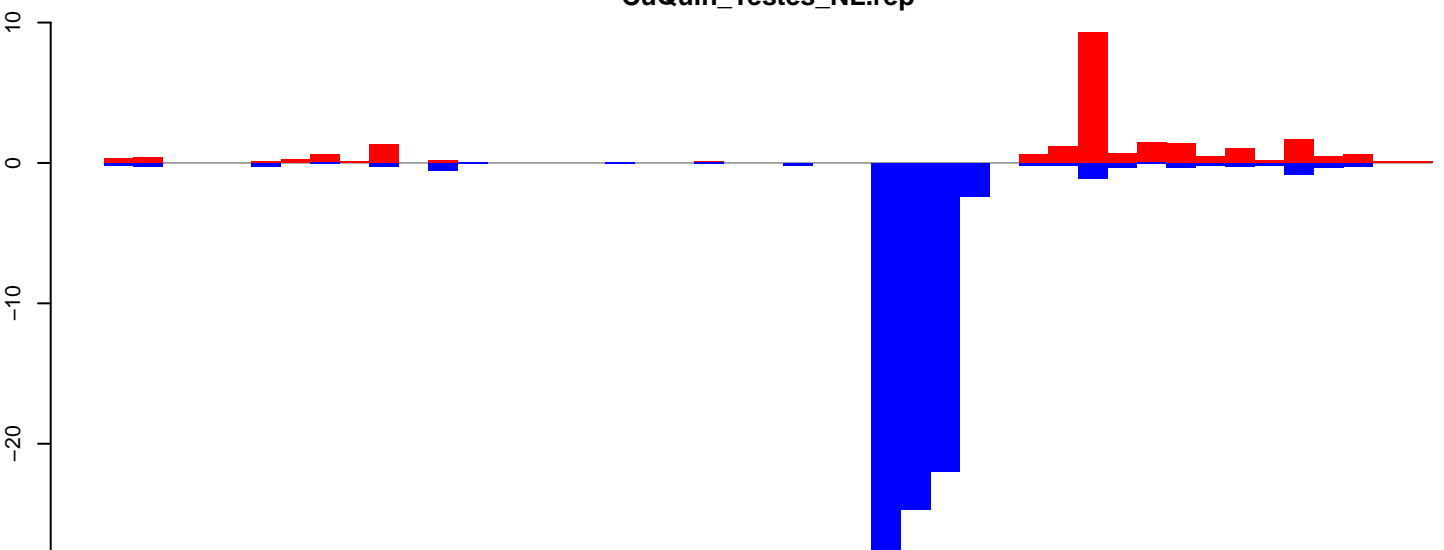
positive=6, negative=27, total=33

CuQuin_Testes_NL.24_35.rep



positive=12, negative=34, total=45

CuQuin_Testes_NL.rep

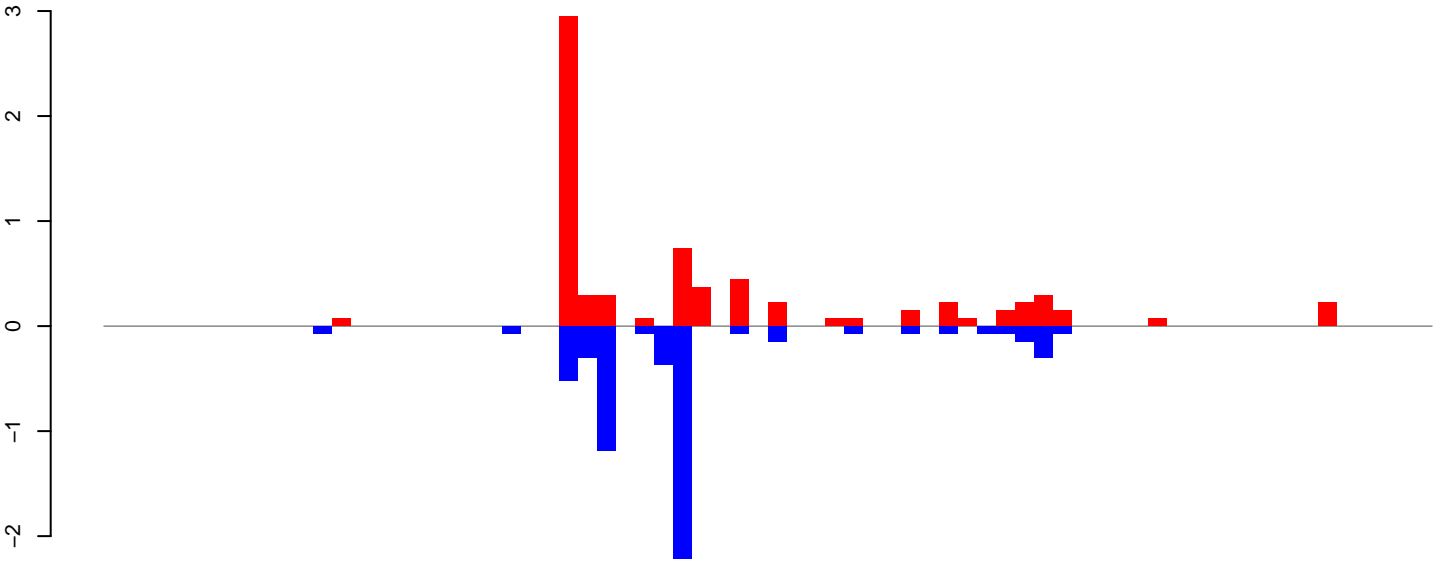


positive=22, negative=88, total=110

Window size=25, length=1103, TE@rnd-5_family-1779@DNA_Crypton-I@1198:1-1103

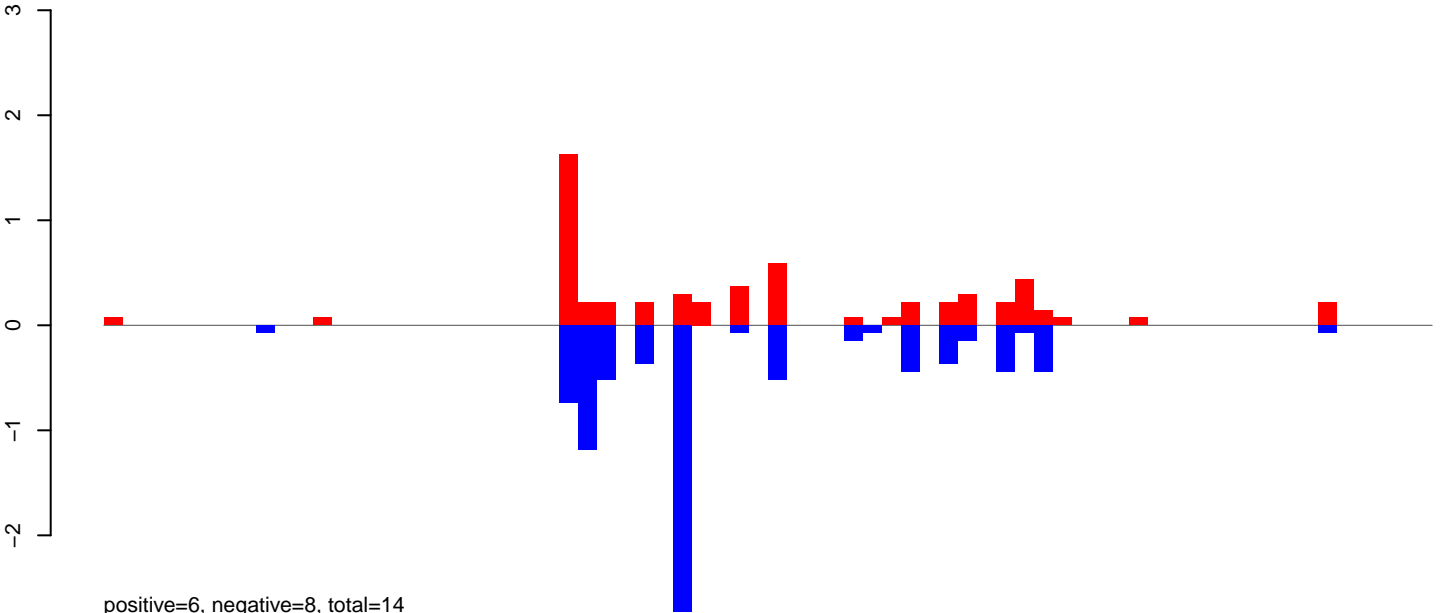
0 200 400 600 800 1000

CuQuin_Testes_NL.18_23.rep



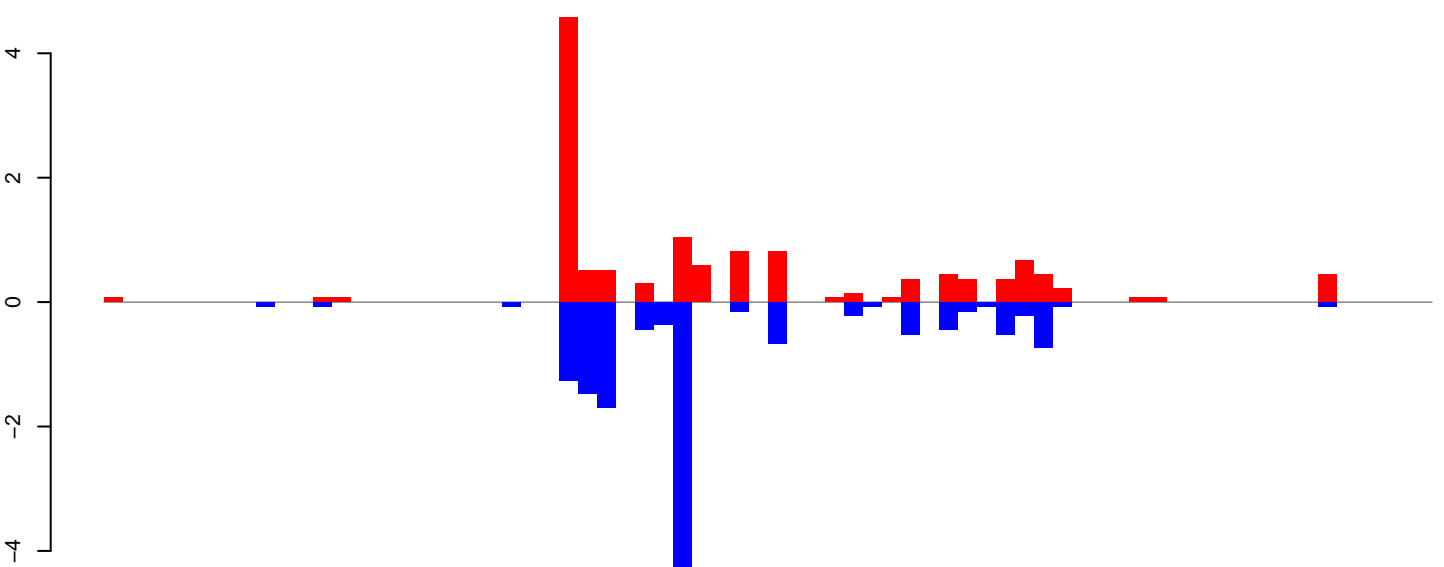
positive=7, negative=6, total=13

CuQuin_Testes_NL.24_35.rep



positive=6, negative=8, total=14

CuQuin_Testes_NL.rep

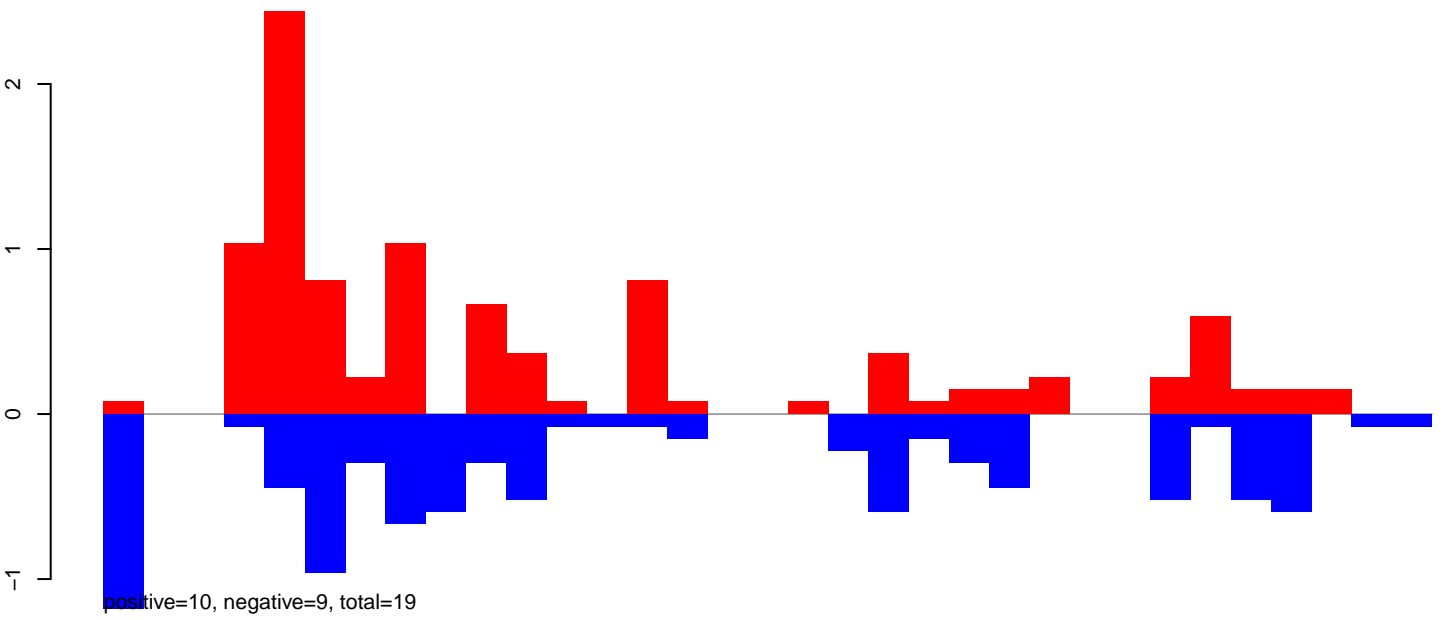


positive=13, negative=14, total=28

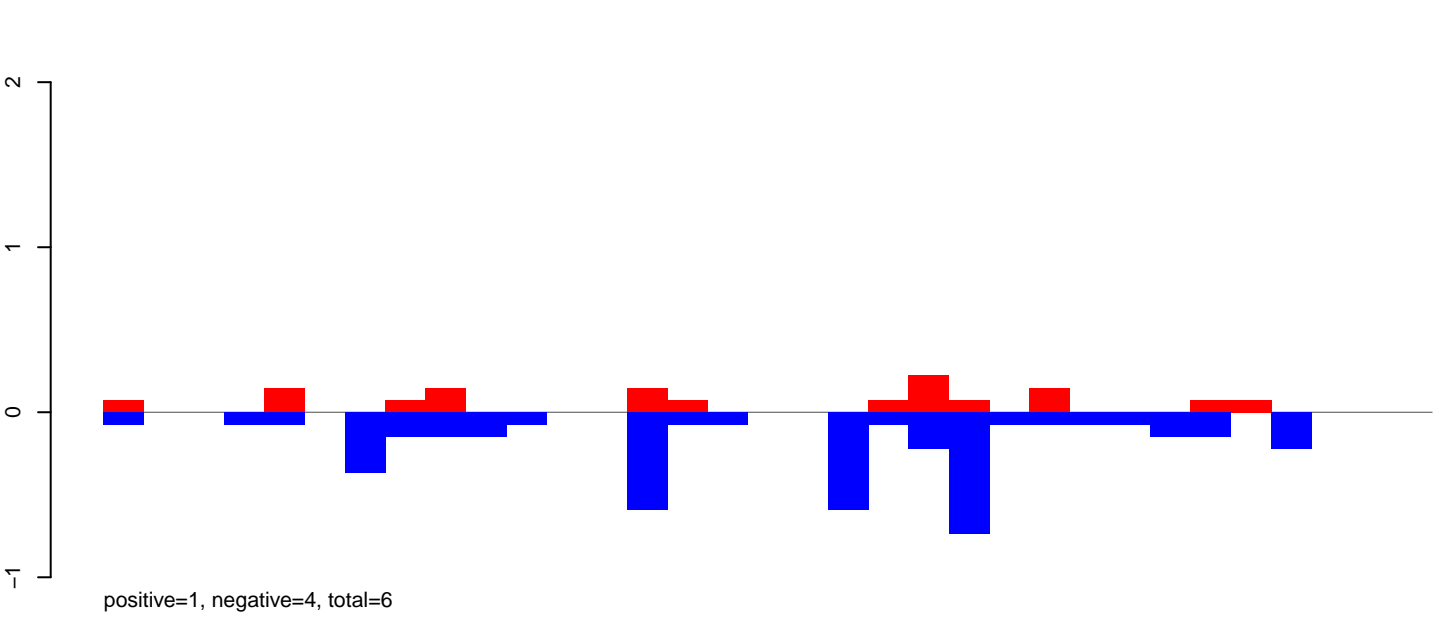
Window size=25, length=1747, TE@rnd-4_family-61@SINE_tRNA@2227:1-1747

0 500 1000 1500

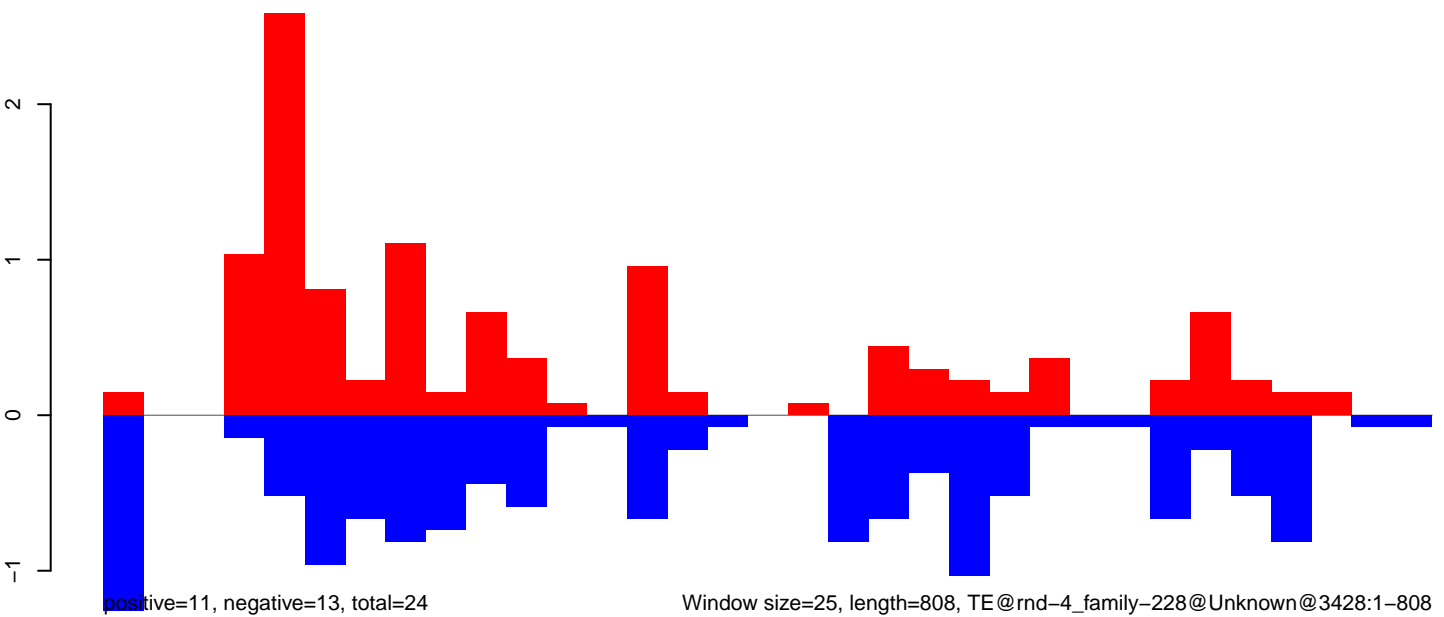
CuQuin_Testes_NL.18_23.rep



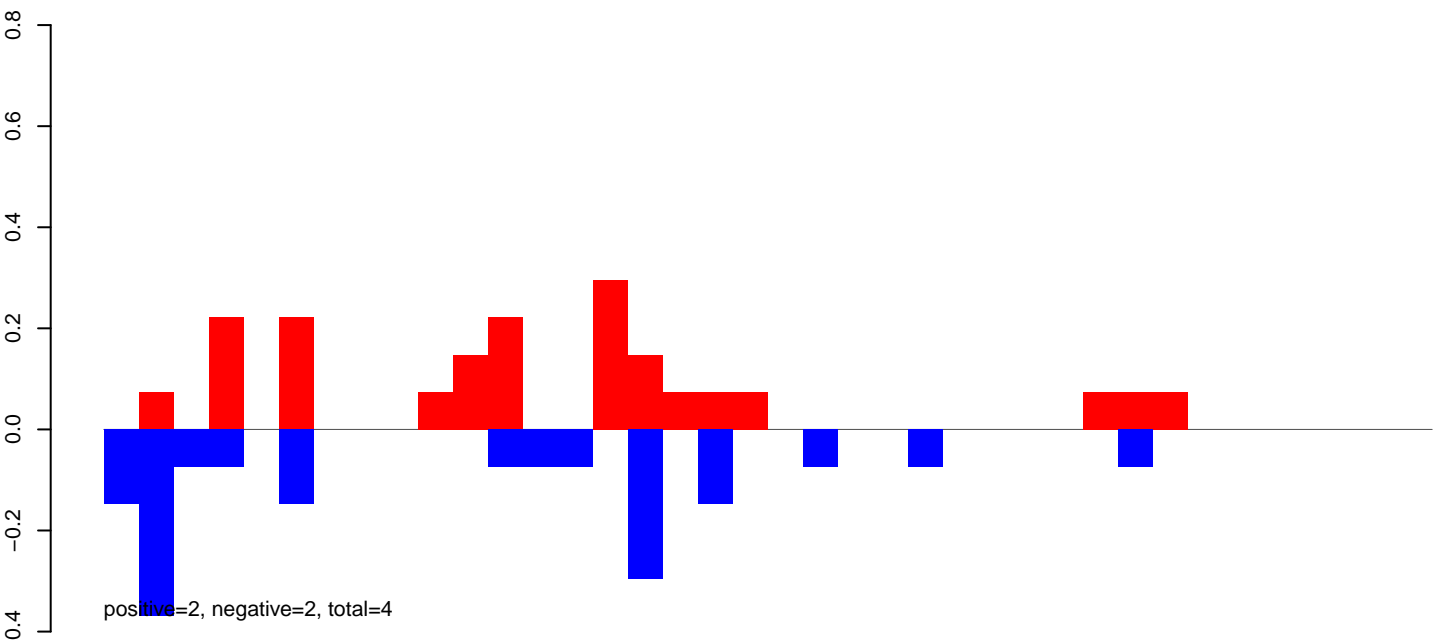
CuQuin_Testes_NL.24_35.rep



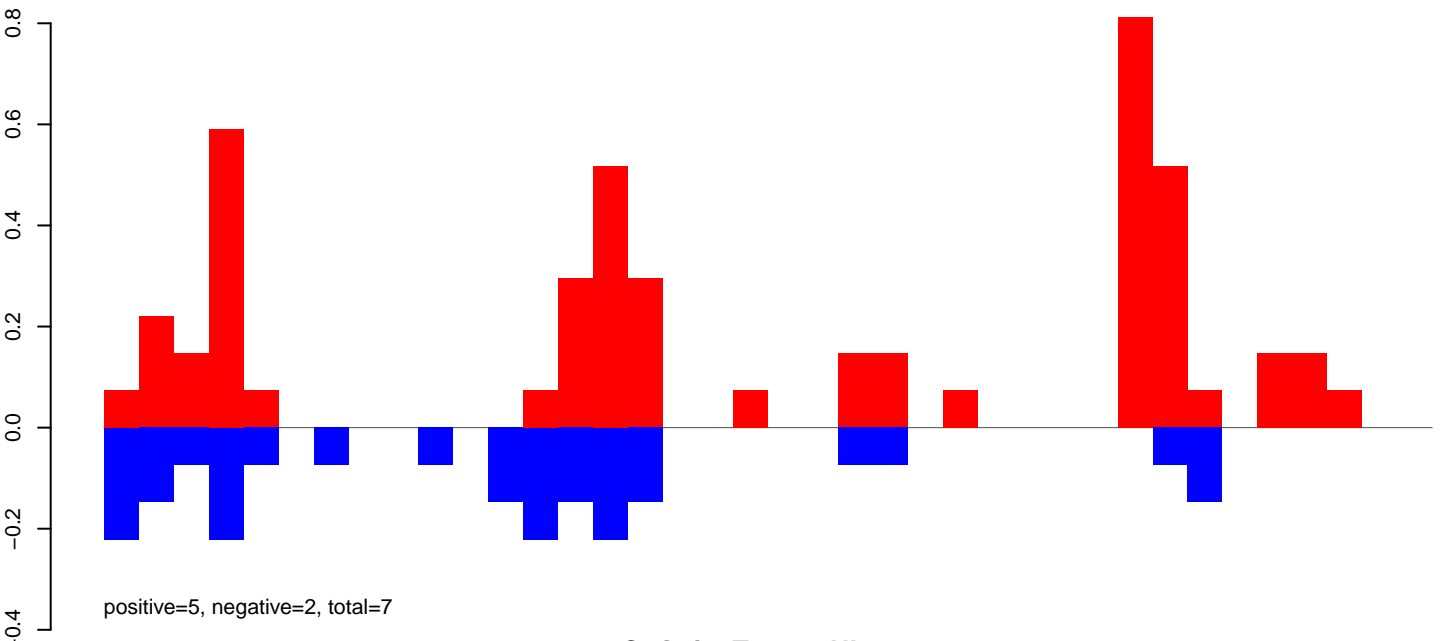
CuQuin_Testes_NL.rep



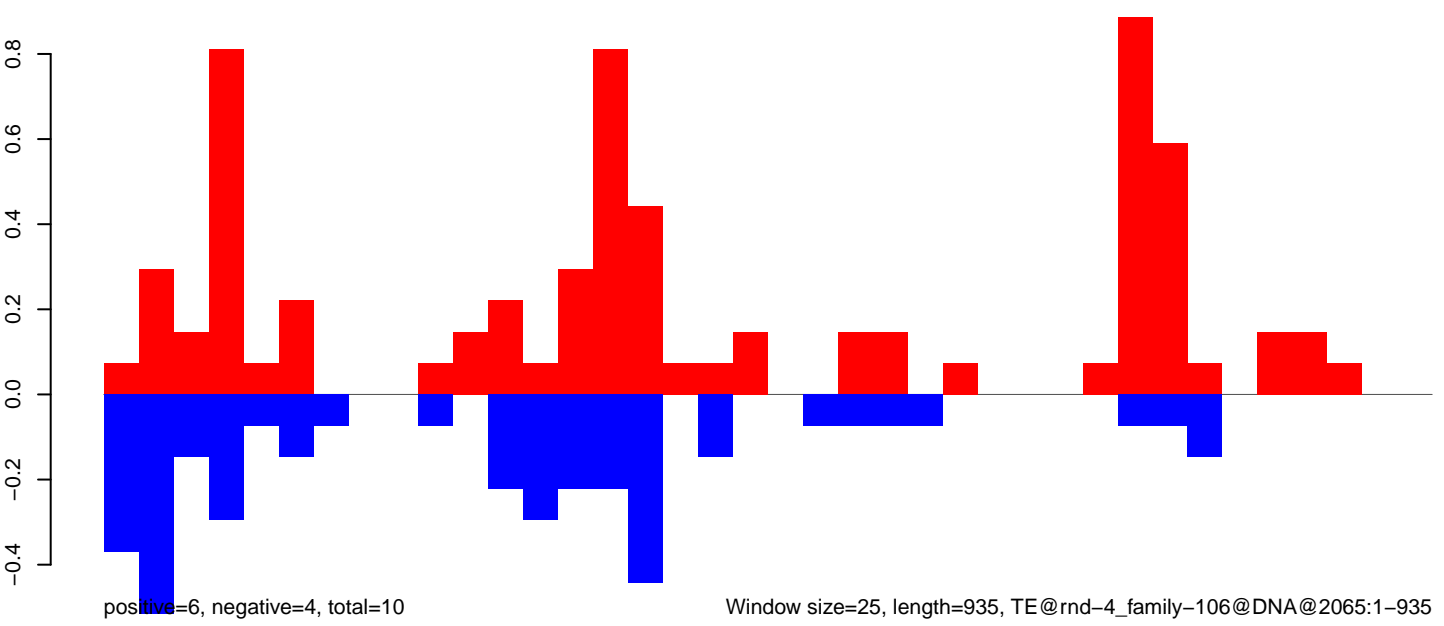
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

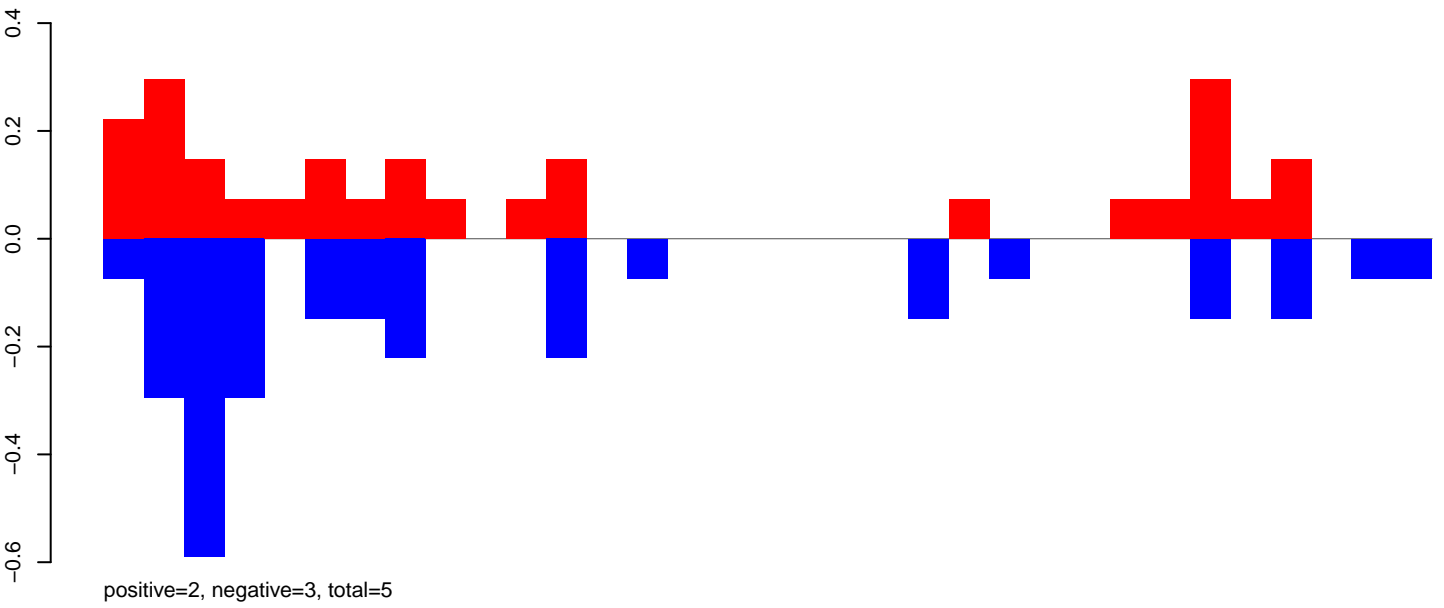


CuQuin_Testes_NL.rep

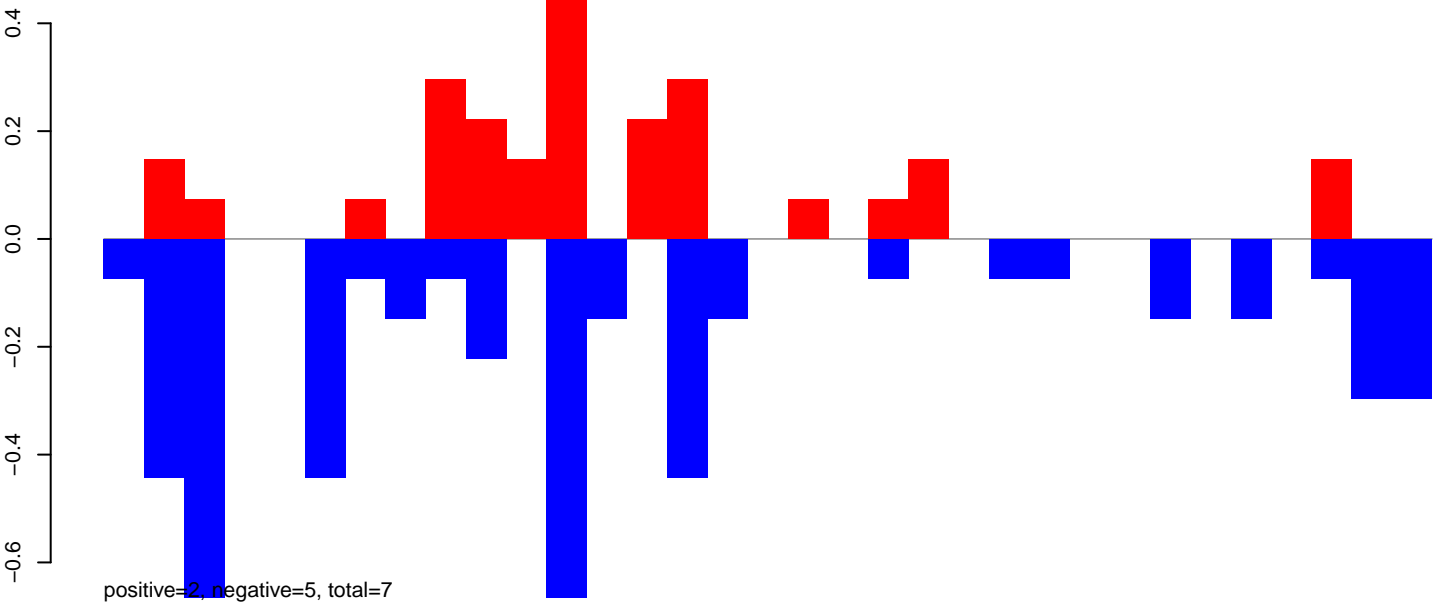


Window size=25, length=935, TE@rnd-4_family-106@DNA@2065:1-935

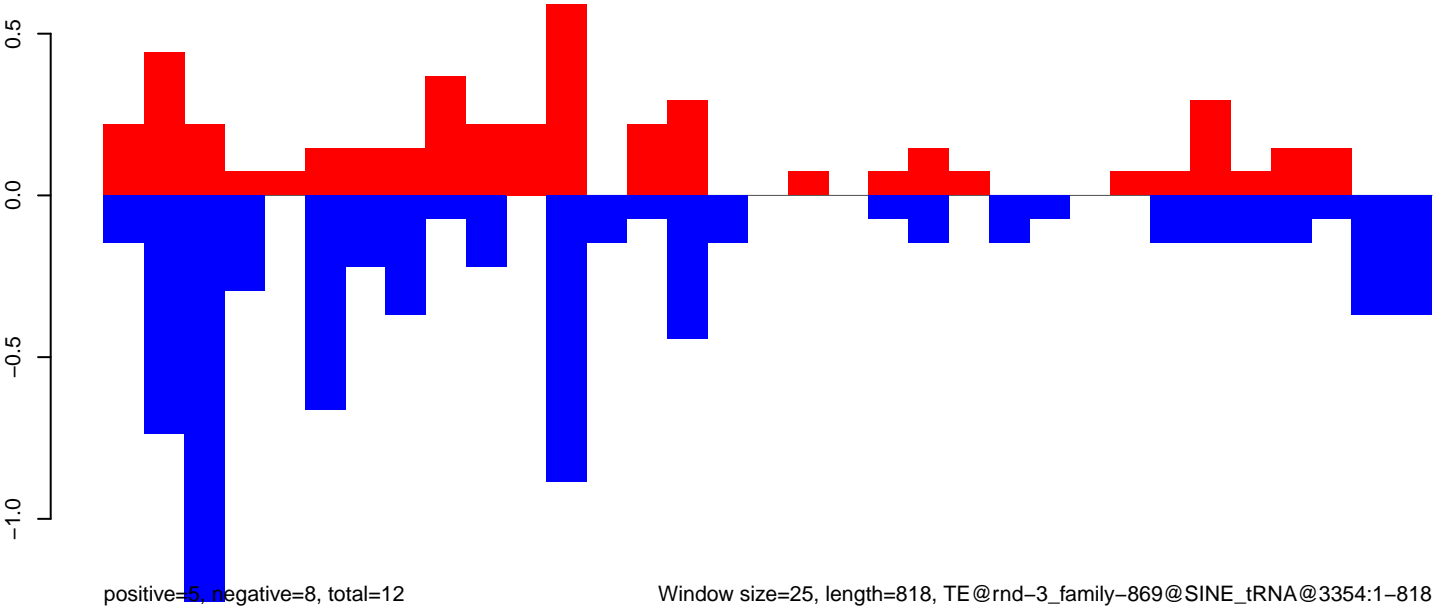
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



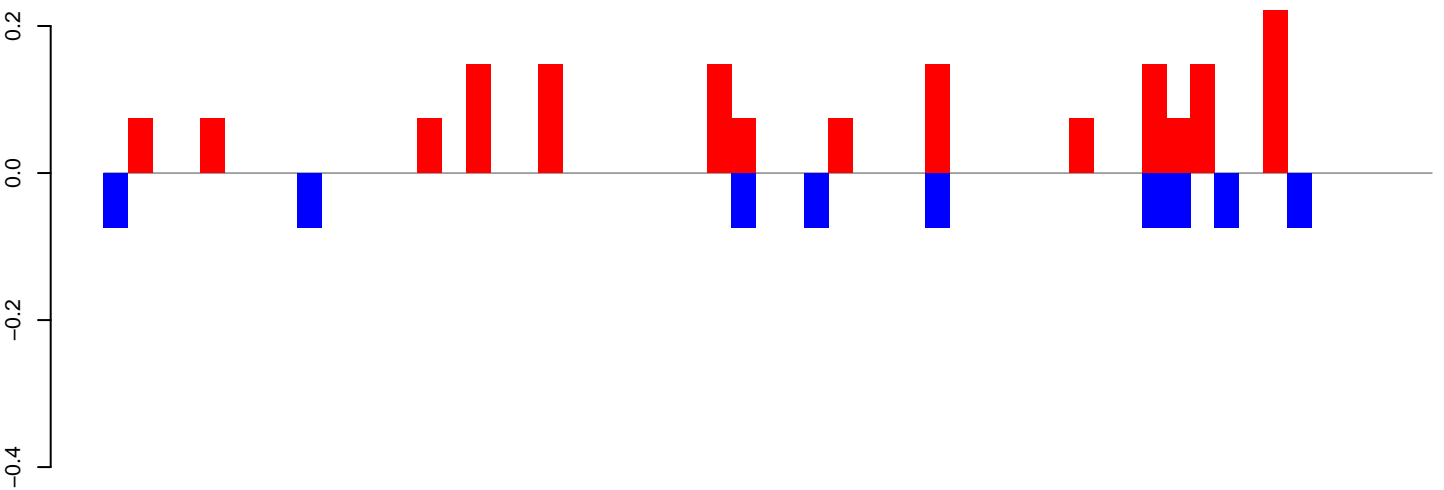
CuQuin_Testes_NL.rep



Window size=25, length=818, TE@rnd-3_family-869@SINE_tRNA@3354:1-818

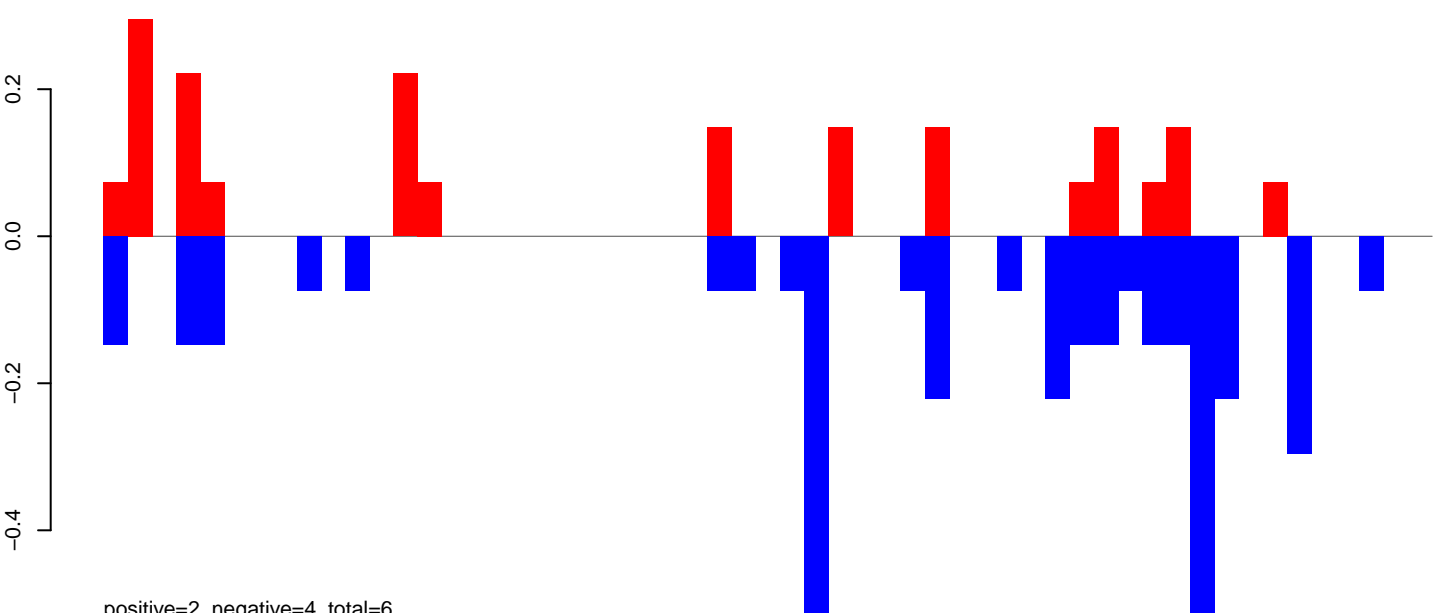
0 200 400 600 800

CuQuin_Testes_NL.18_23.rep



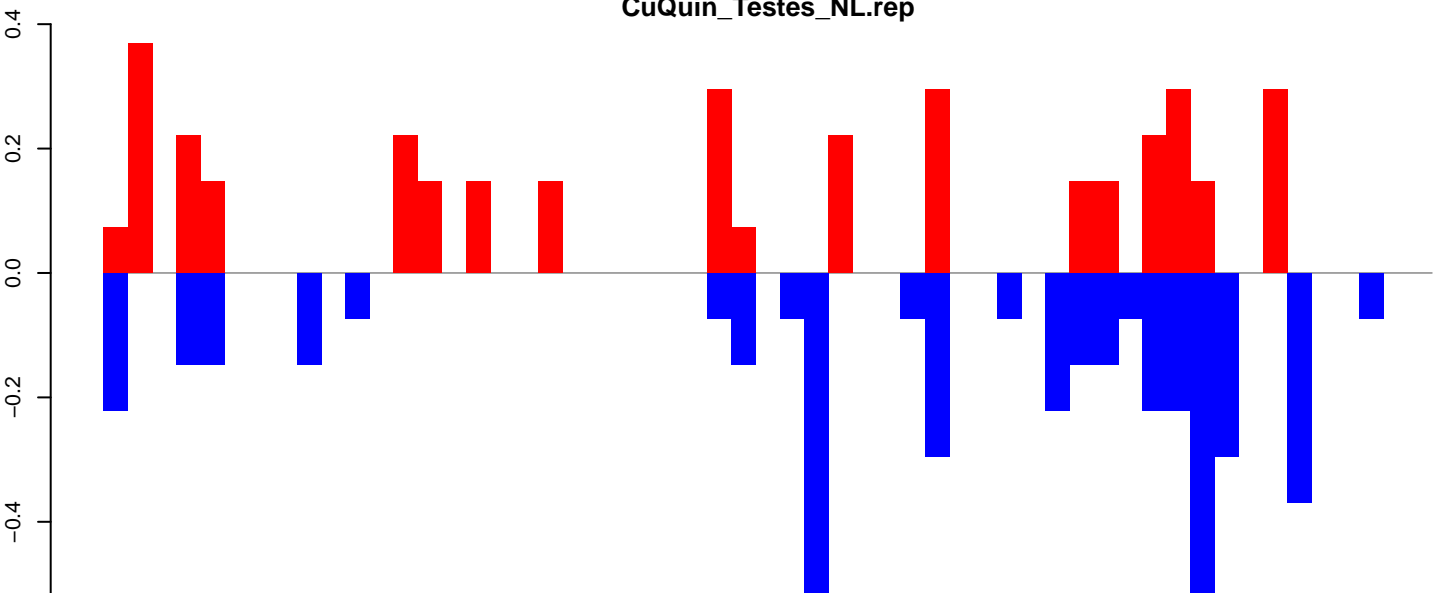
positive=2, negative=1, total=2

CuQuin_Testes_NL.24_35.rep



positive=2, negative=4, total=6

CuQuin_Testes_NL.rep

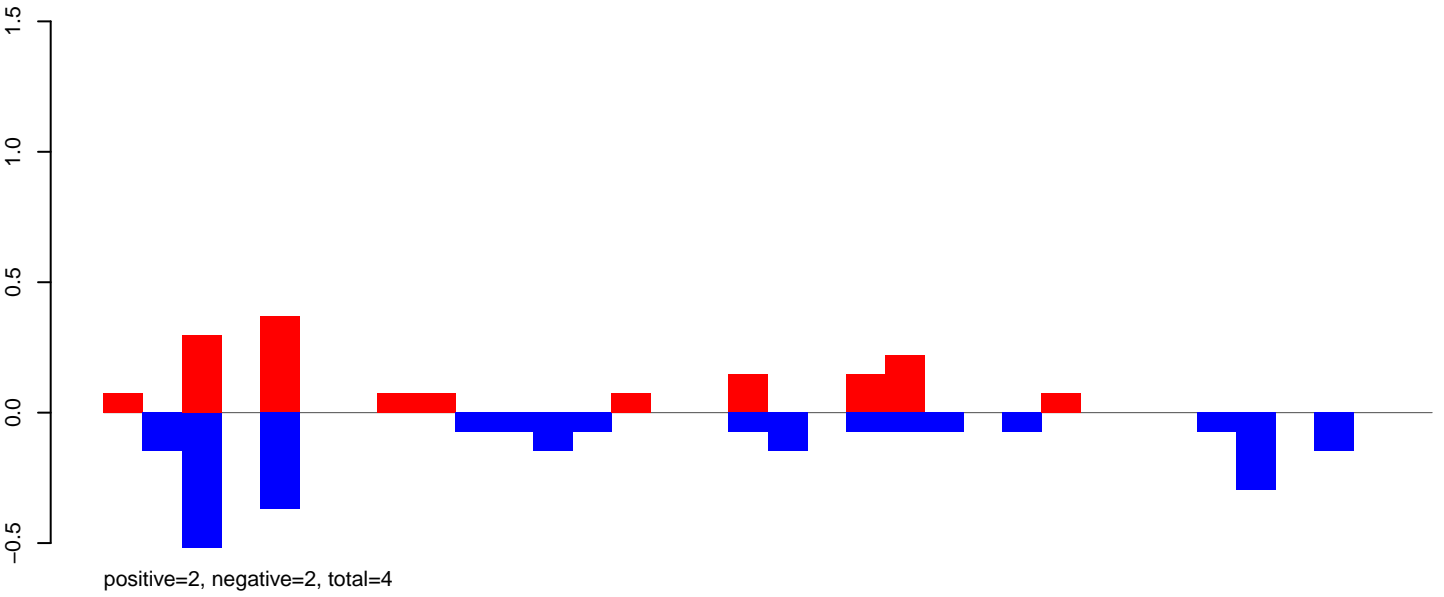


positive=4, negative=4, total=8

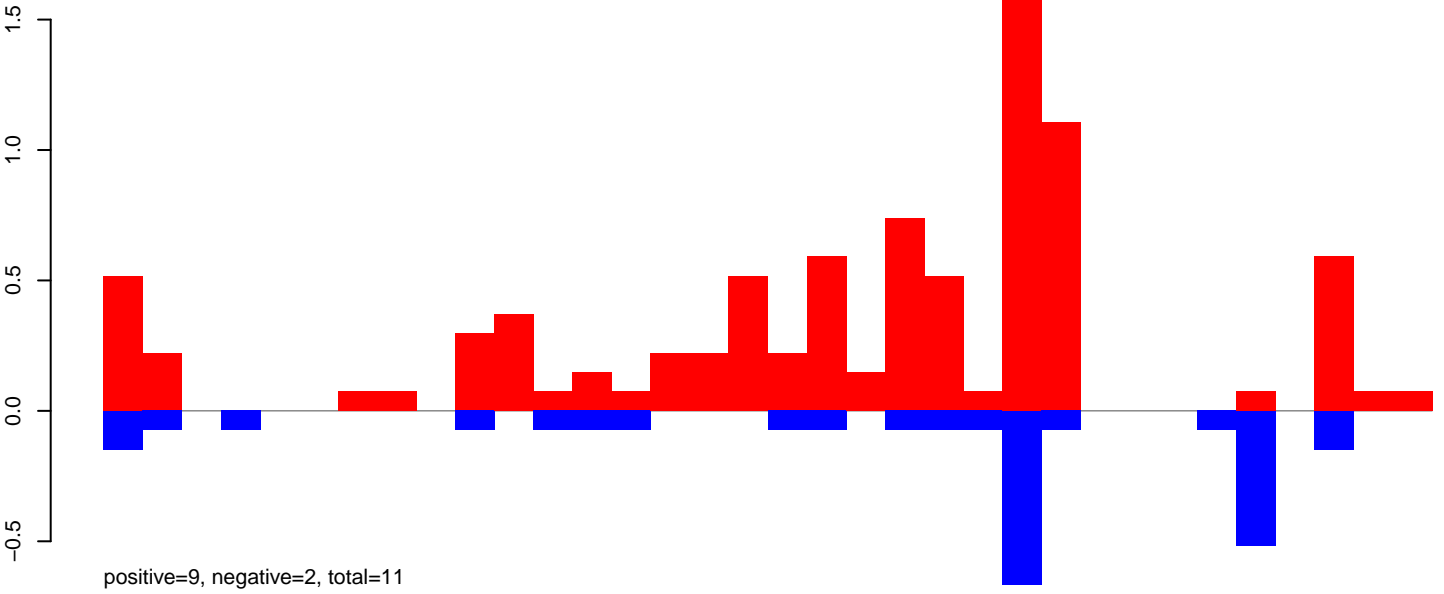
Window size=25, length=1356, TE@rnd-3_family-830@Unknown@140:1-1356

0 200 400 600 800 1000 1200 1400

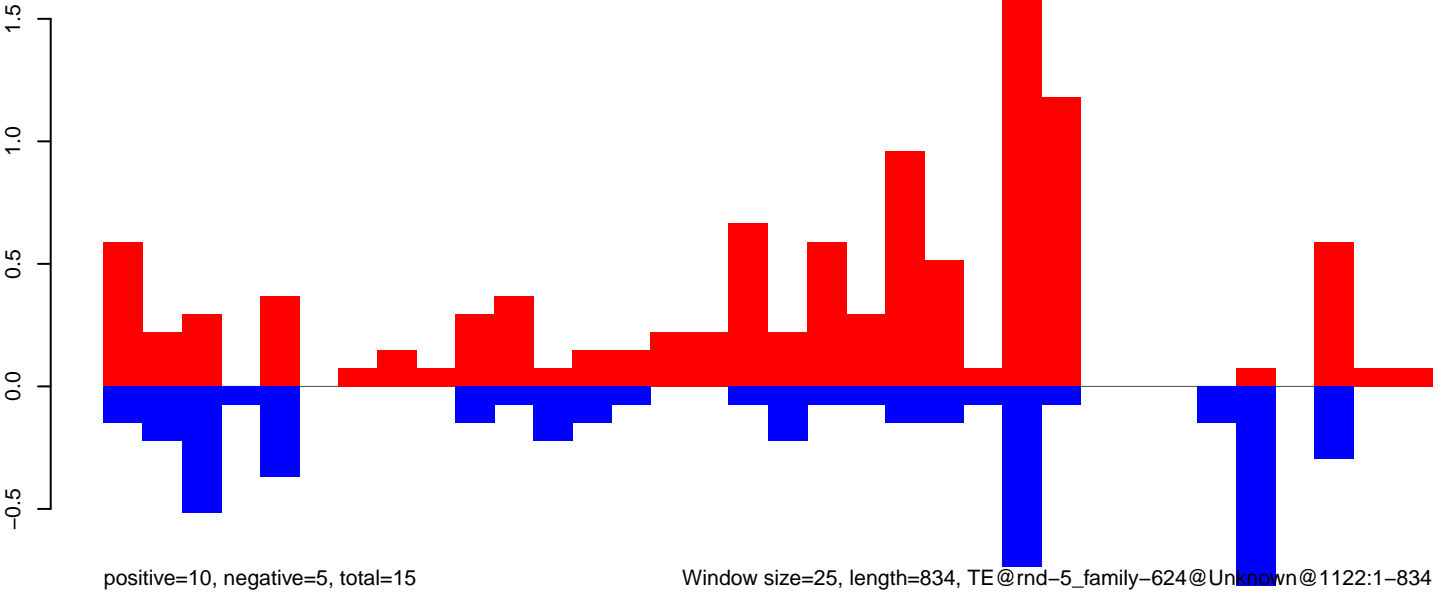
CuQuin_Testes_NL.18_23.rep



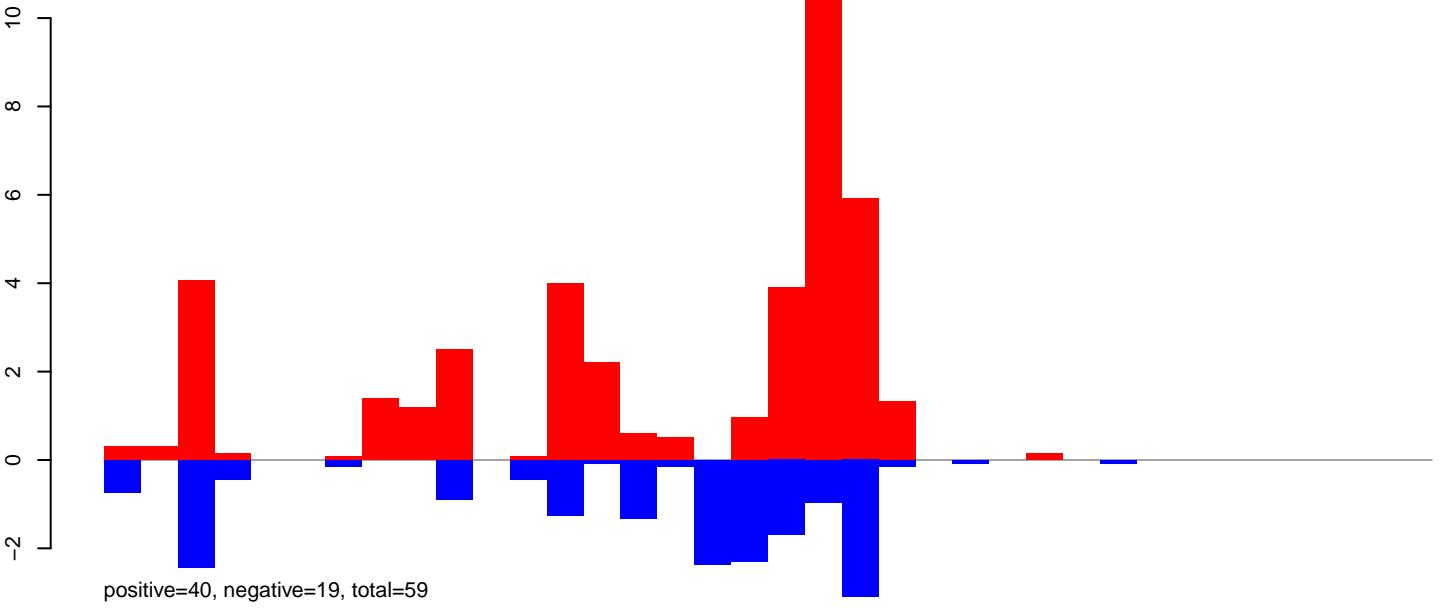
CuQuin_Testes_NL.24_35.rep



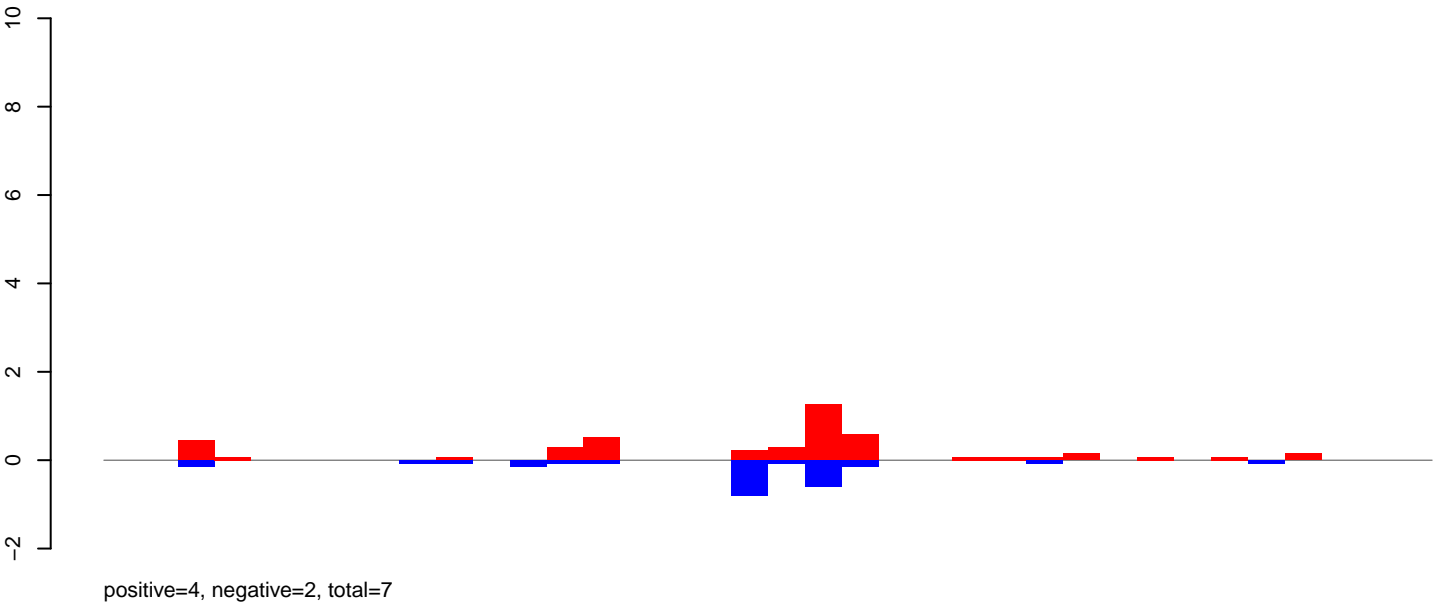
CuQuin_Testes_NL.rep



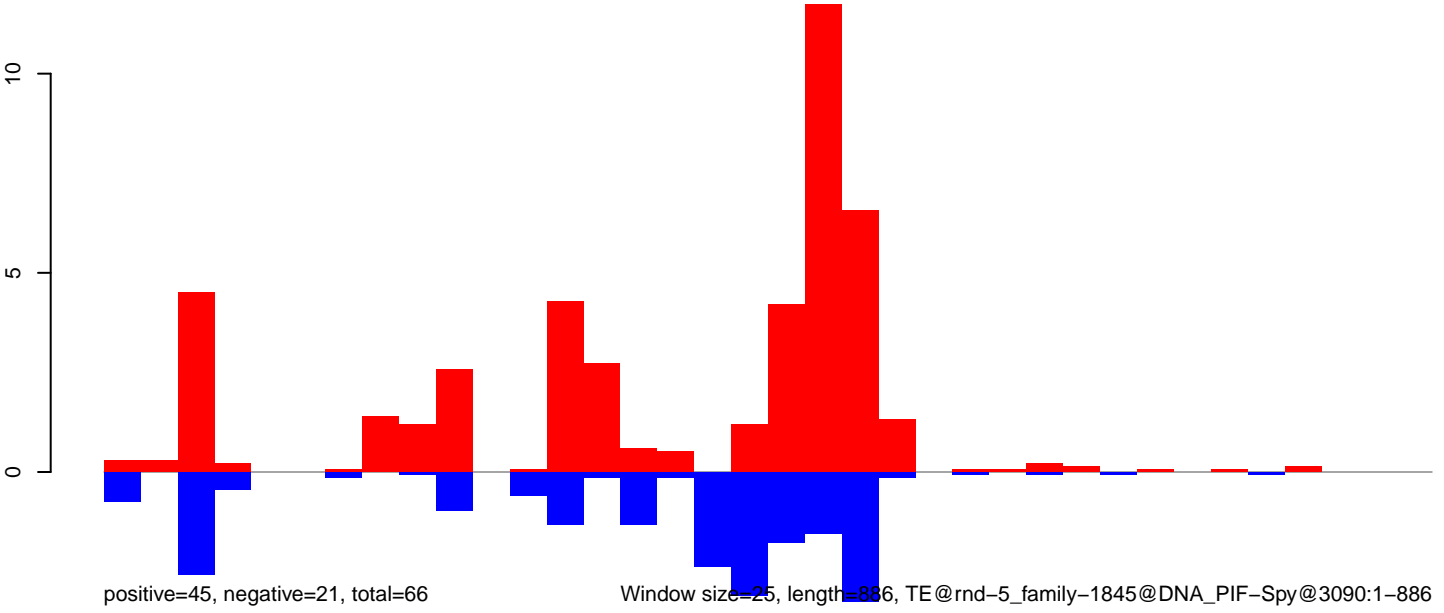
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

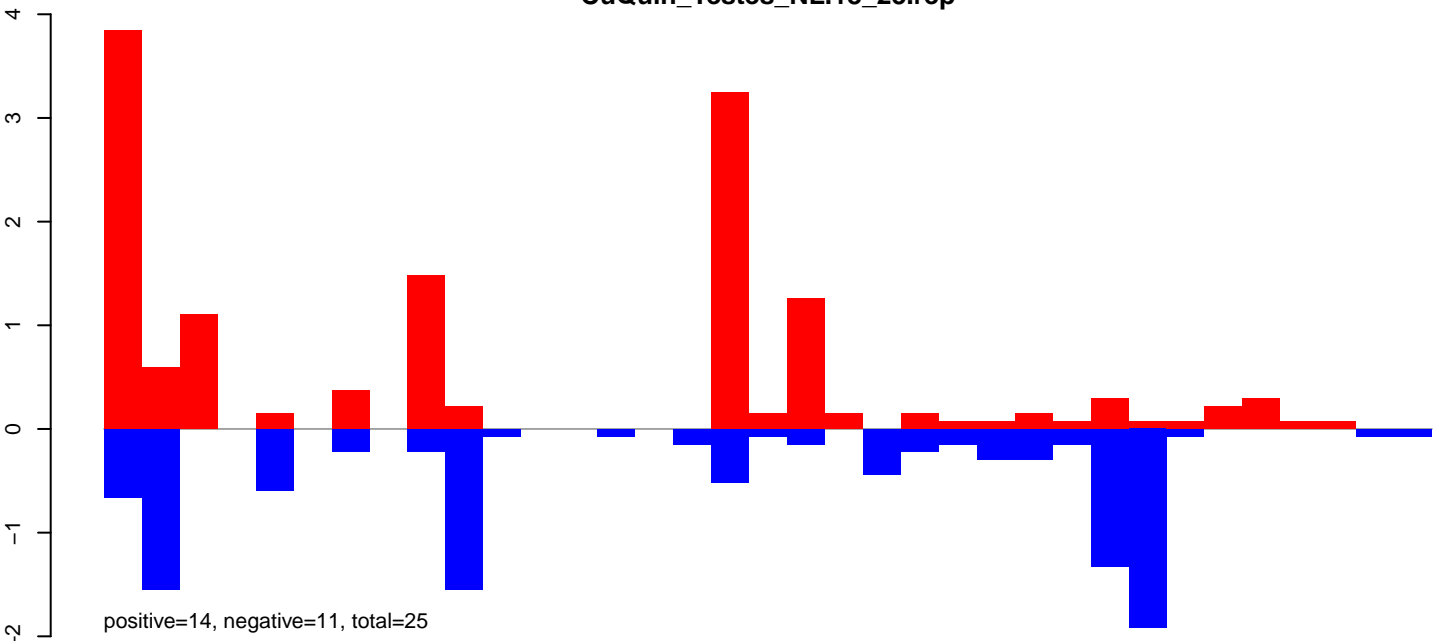


CuQuin_Testes_NL.rep

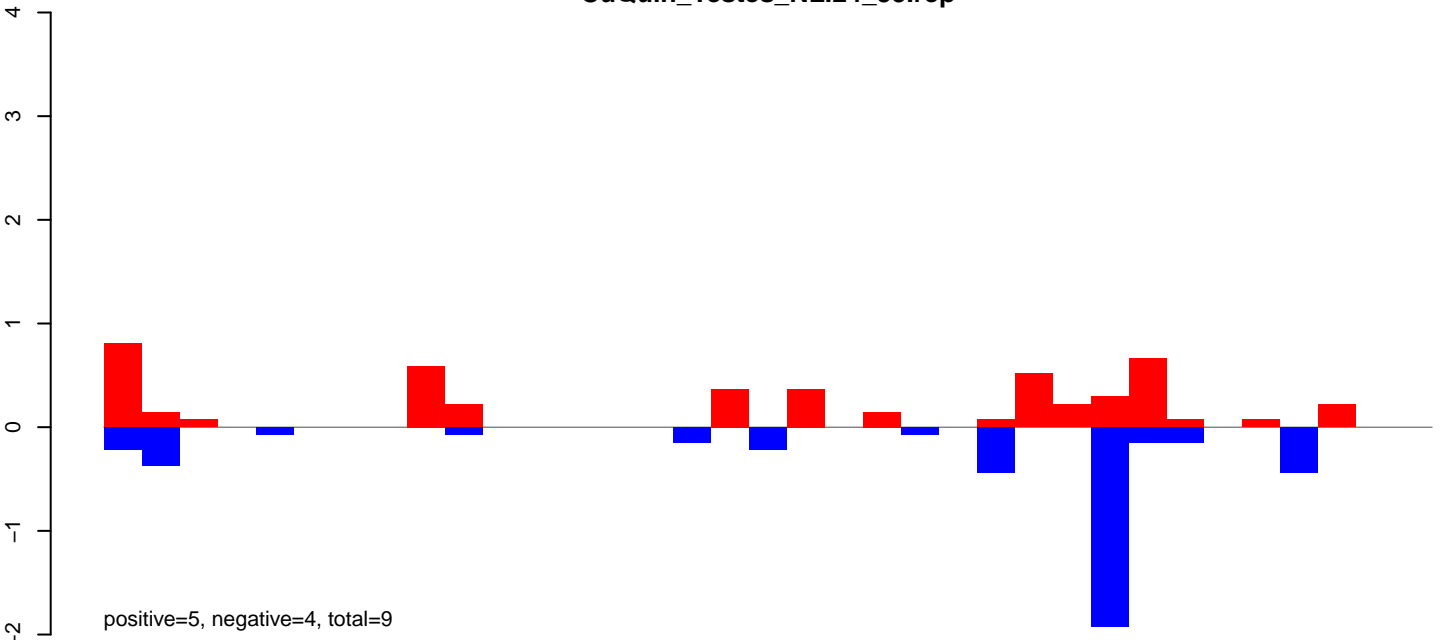


Window size=25, length=886, TE@rnd-5_family-1845@DNA_PIF-Spy@3090:1-886

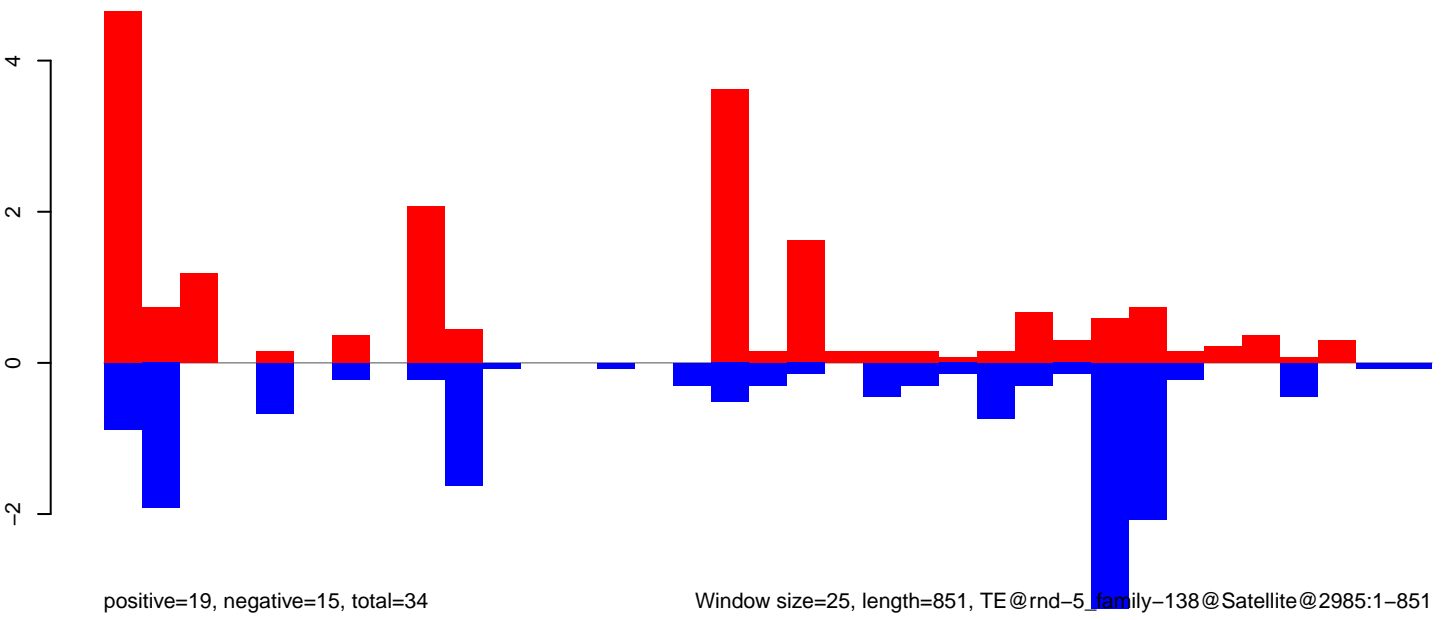
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

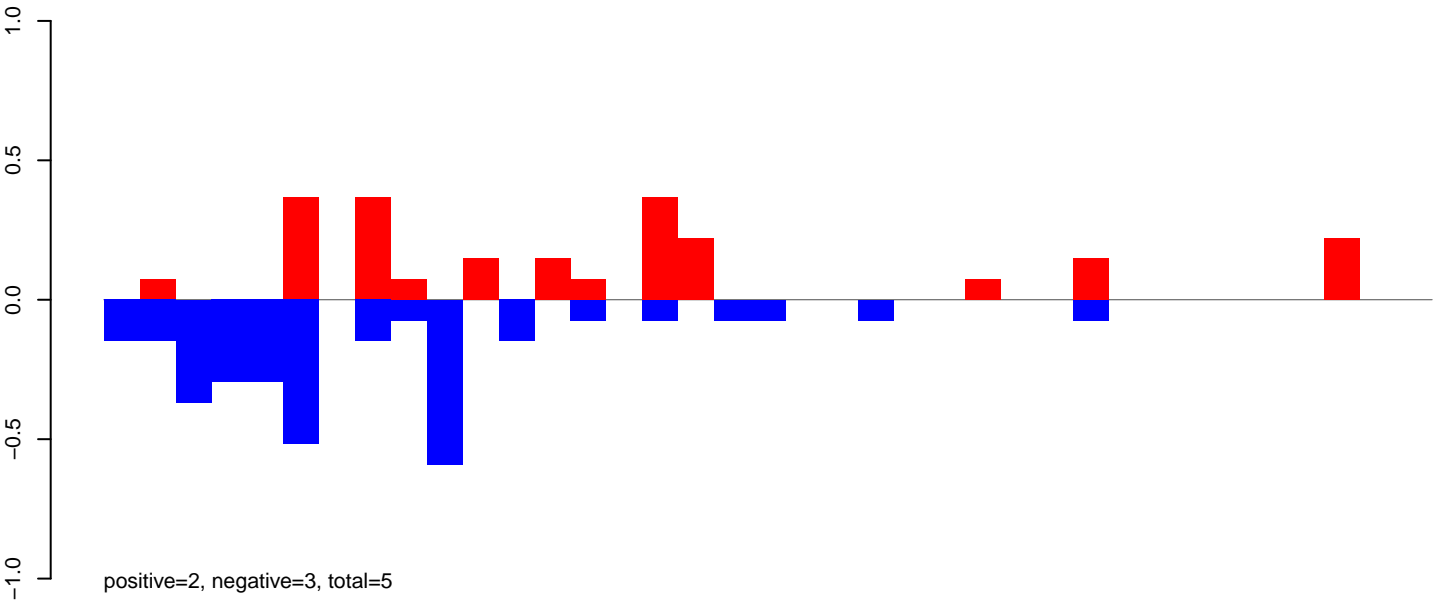


CuQuin_Testes_NL.rep

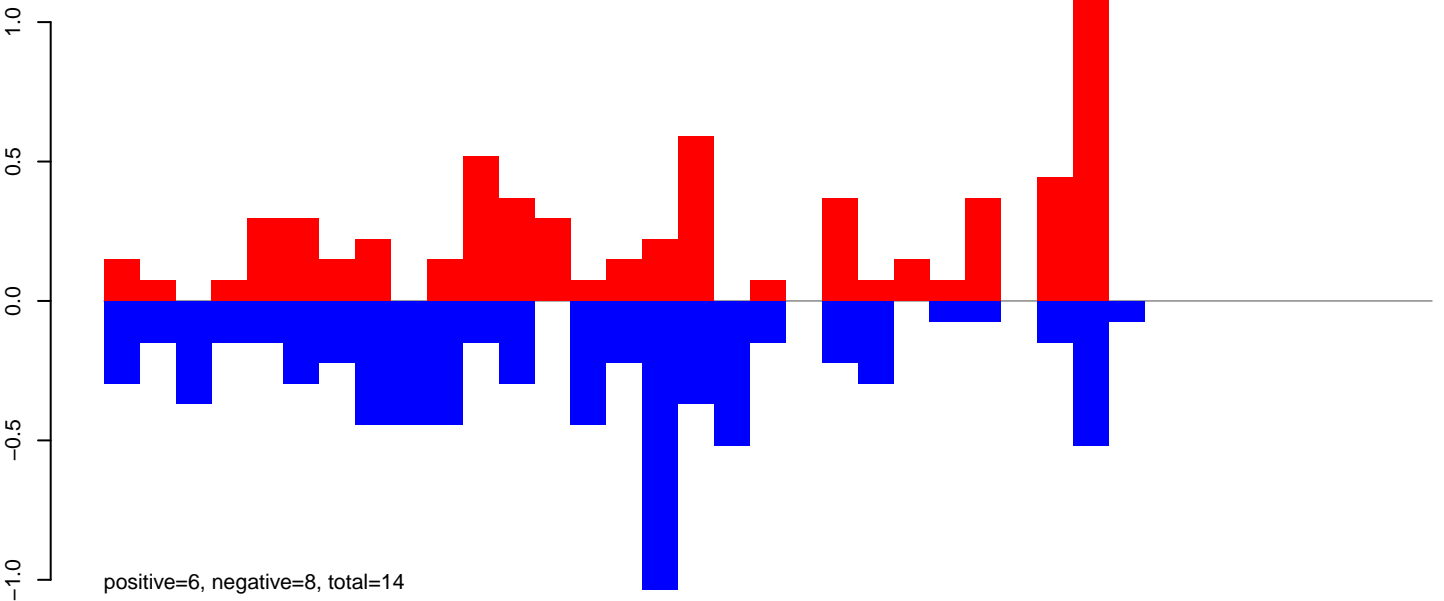


Window size=25, length=851, TE@rnd-5_family-138@Satellite@2985:1-851

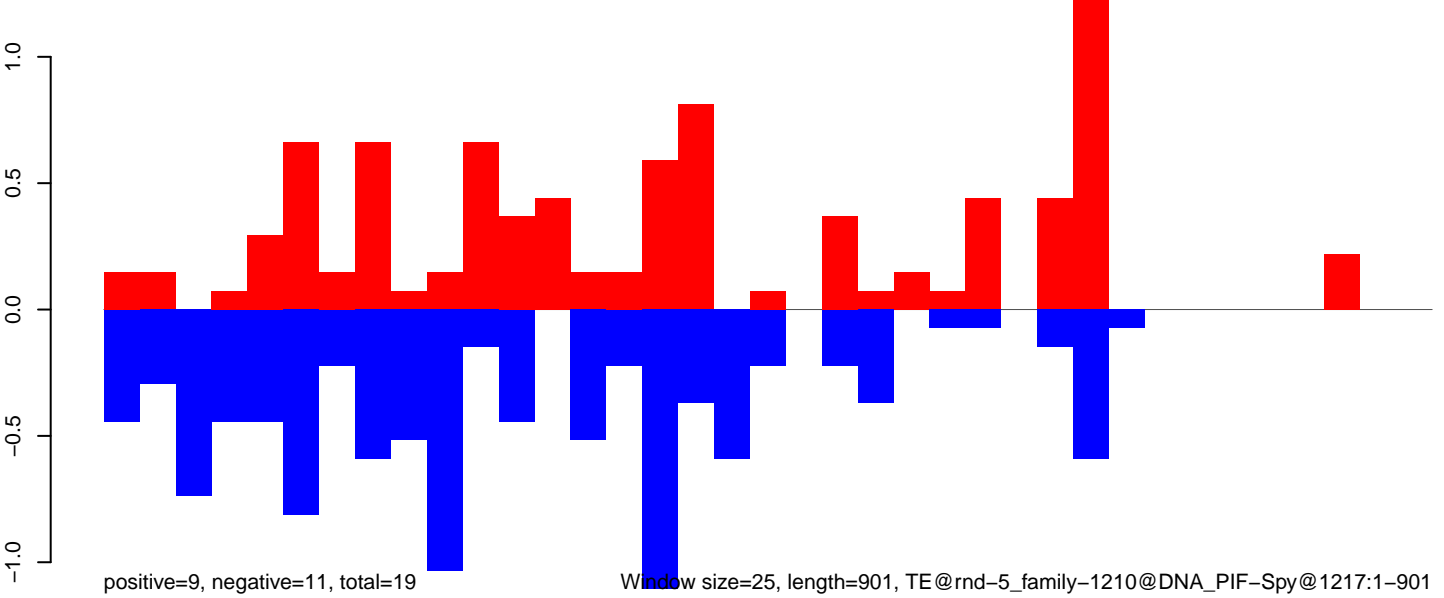
CuQuin_Testes_NL.18_23.rep



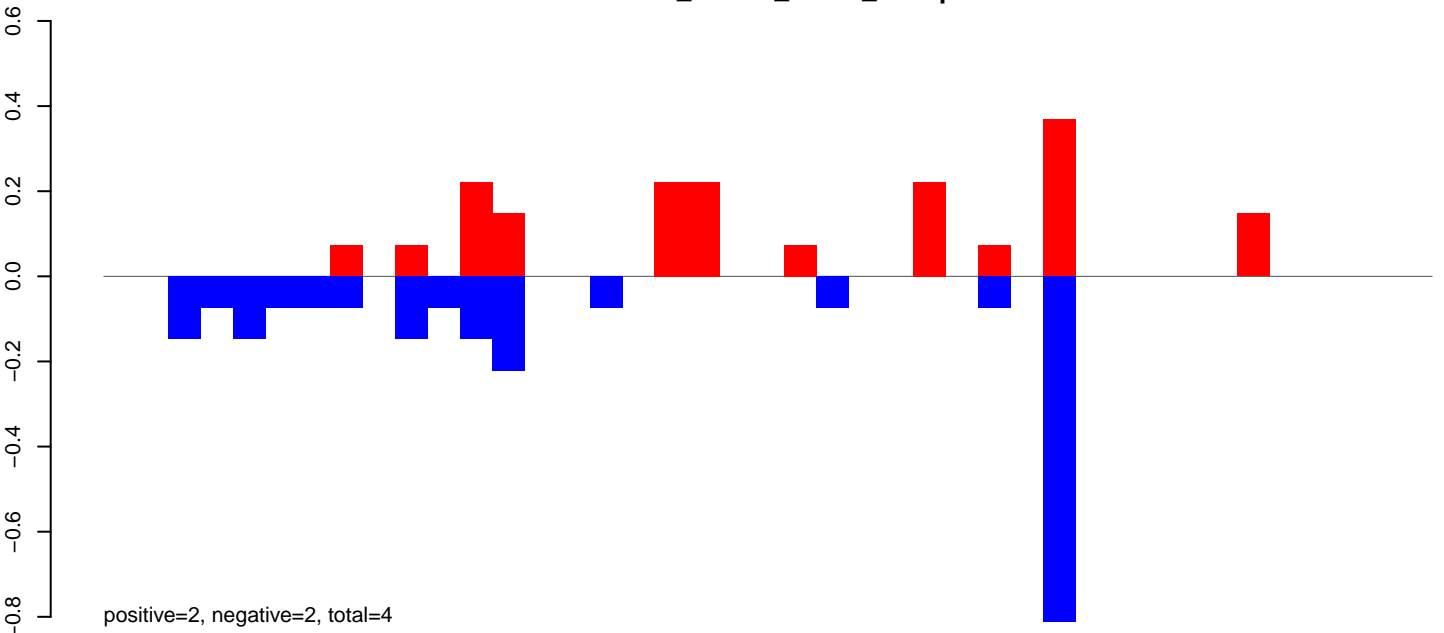
CuQuin_Testes_NL.24_35.rep



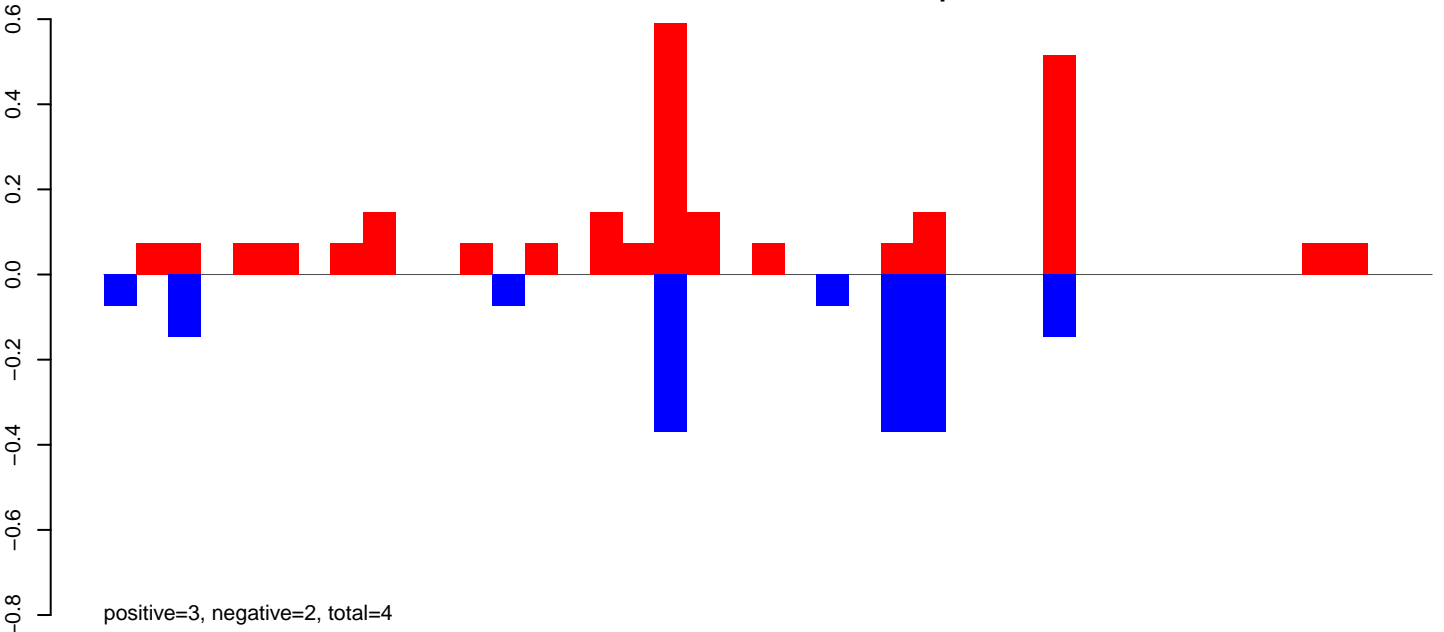
CuQuin_Testes_NL.rep



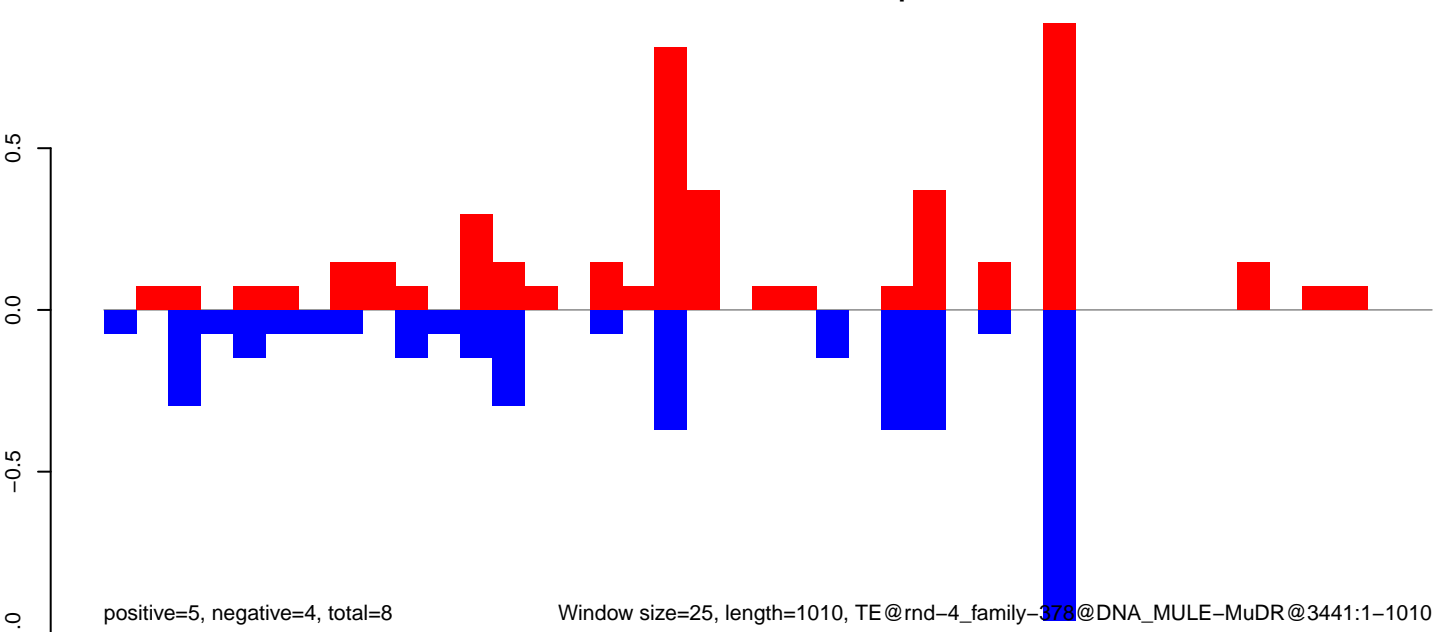
CuQuin_Testes_NL.18_23.rep



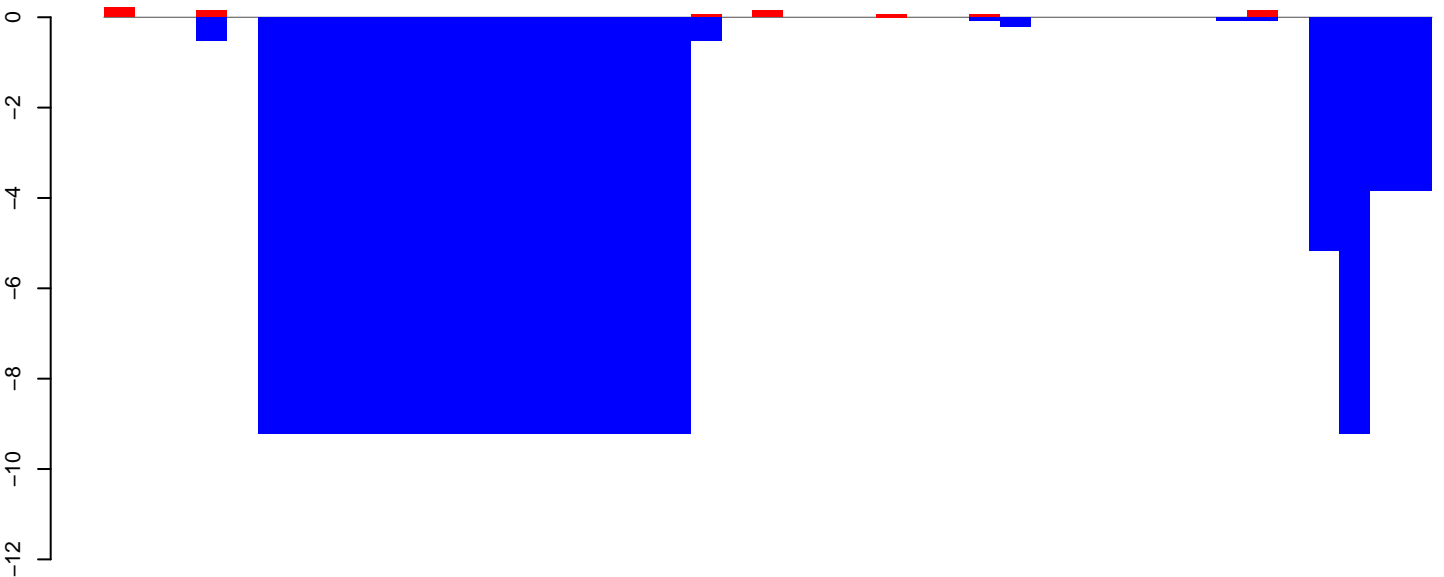
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

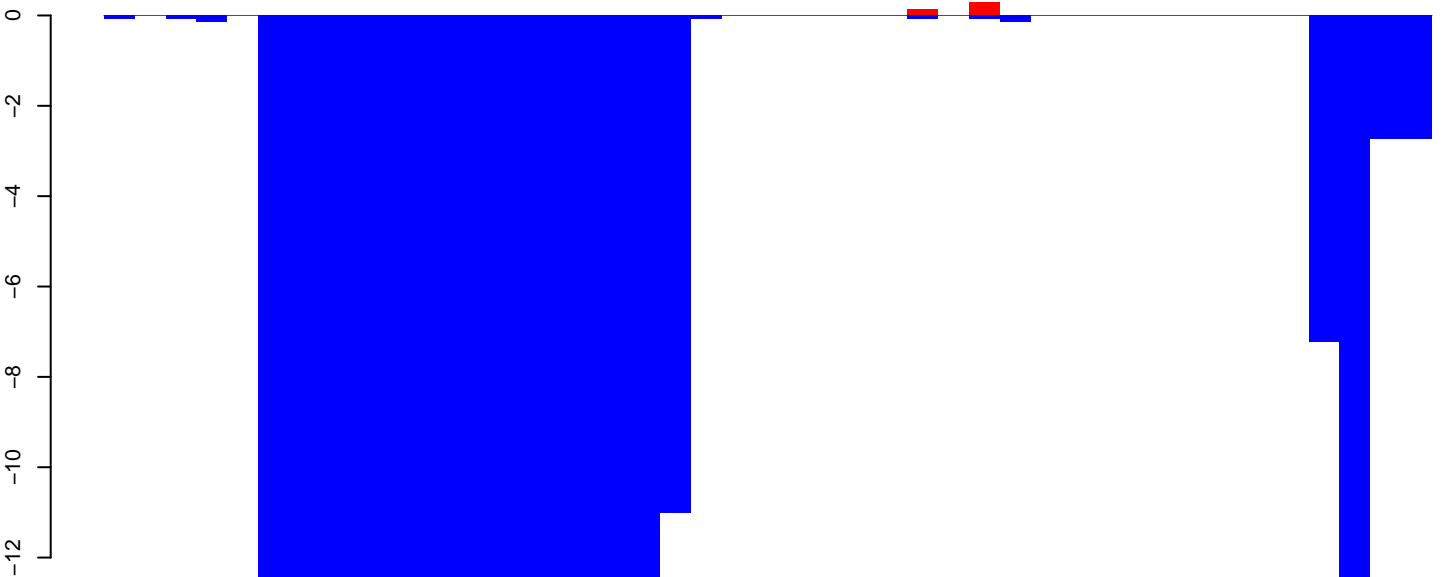


CuQuin_Testes_NL.18_23.rep



positive=1, negative=153, total=154

CuQuin_Testes_NL.24_35.rep



positive=0, negative=205, total=206

CuQuin_Testes_NL.rep

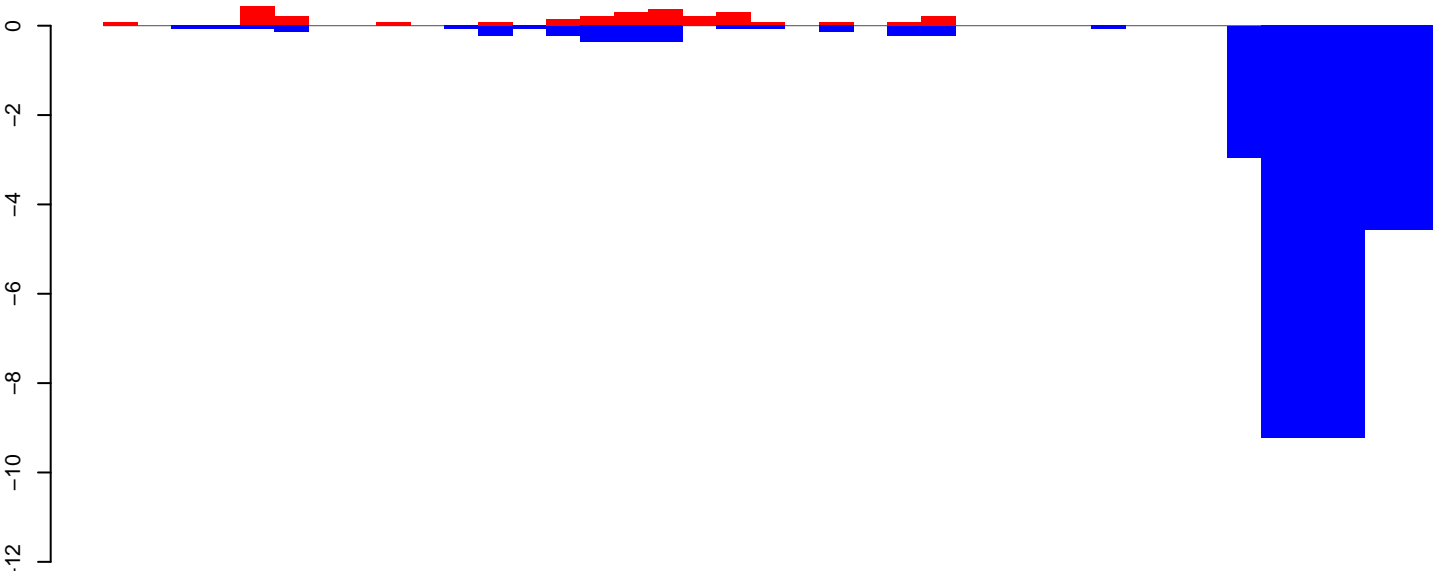


positive=1, negative=1082, total=1083

Window size=25, length=1057, TE@rnd-4_family-2761@DNA_Sola-1@1588:1-1057

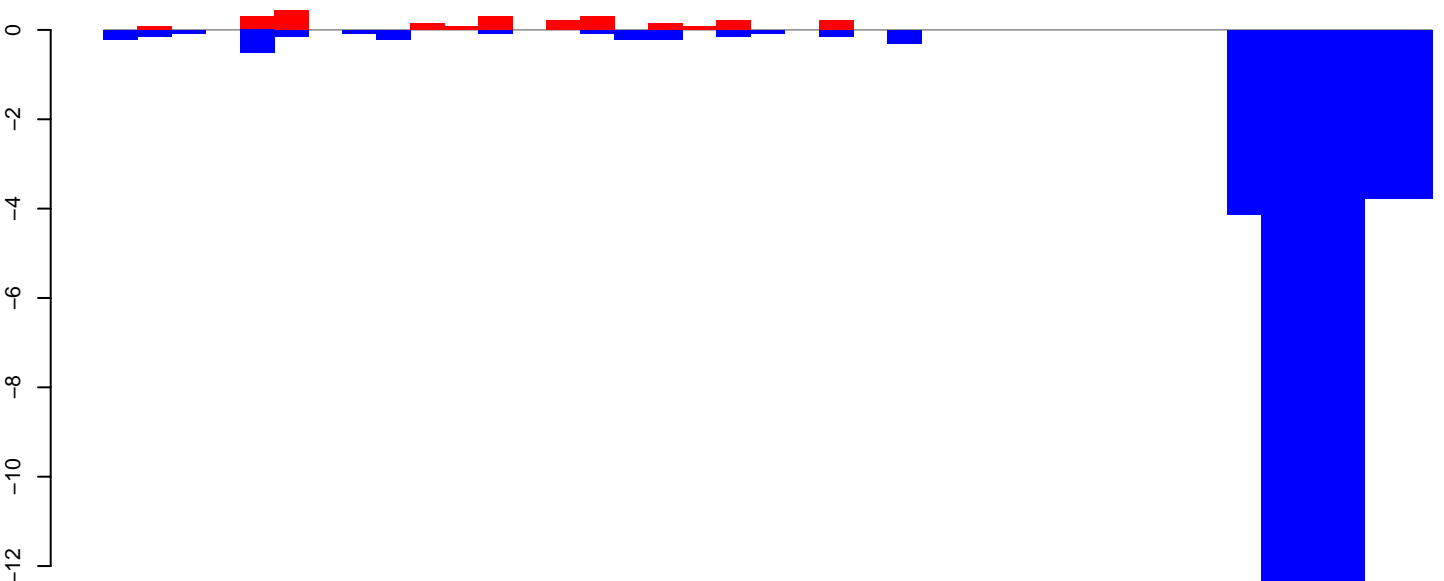
0 200 400 600 800 1000

CuQuin_Testes_NL.18_23.rep



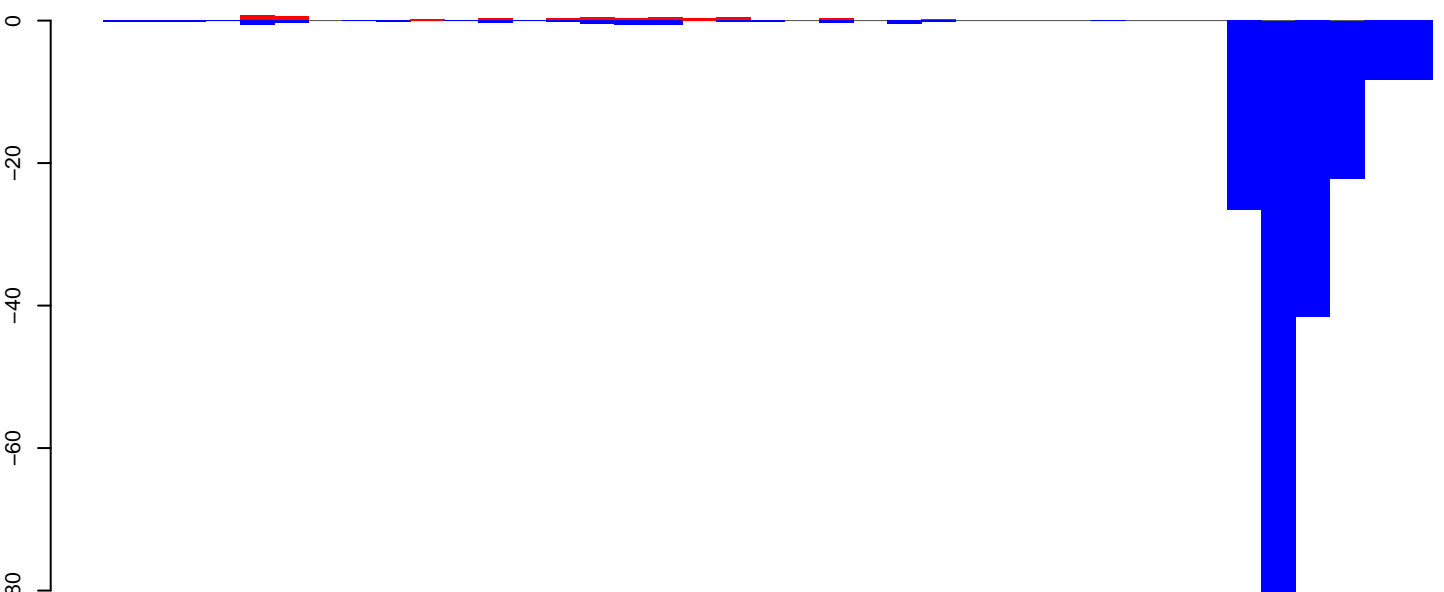
positive=3, negative=43, total=46

CuQuin_Testes_NL.24_35.rep



positive=3, negative=53, total=56

CuQuin_Testes_NL.rep

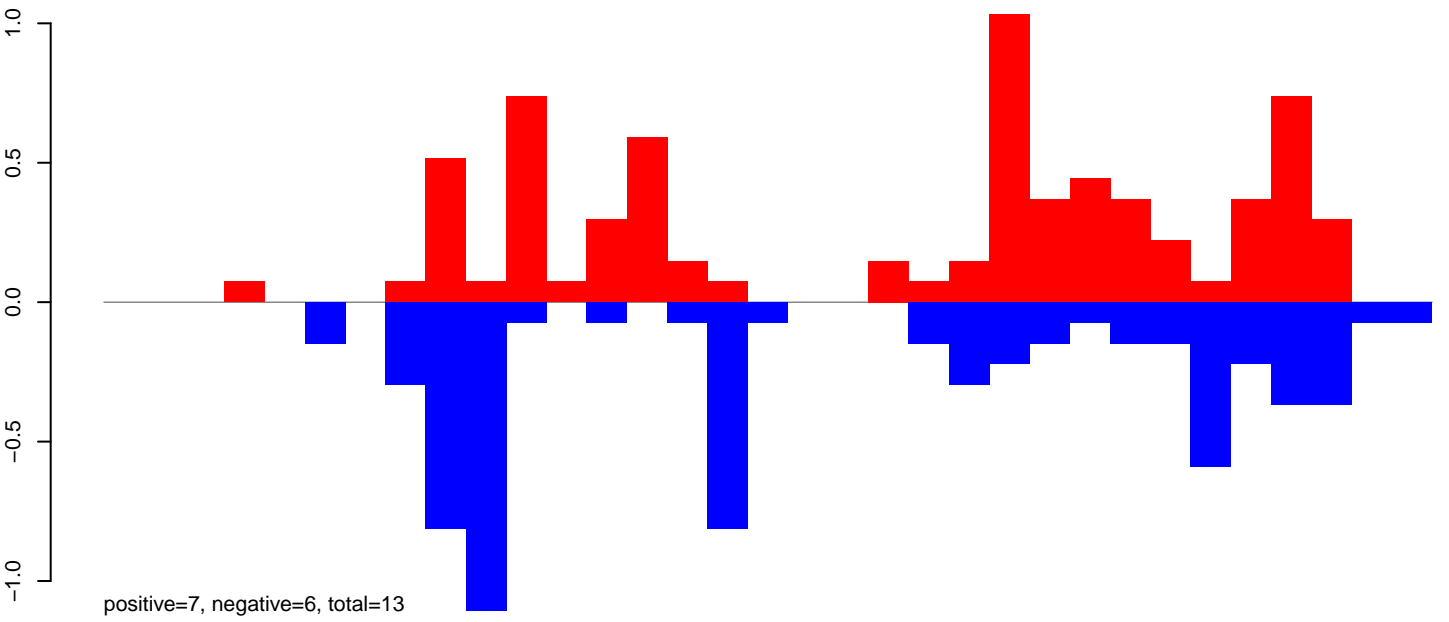


positive=5, negative=196, total=201

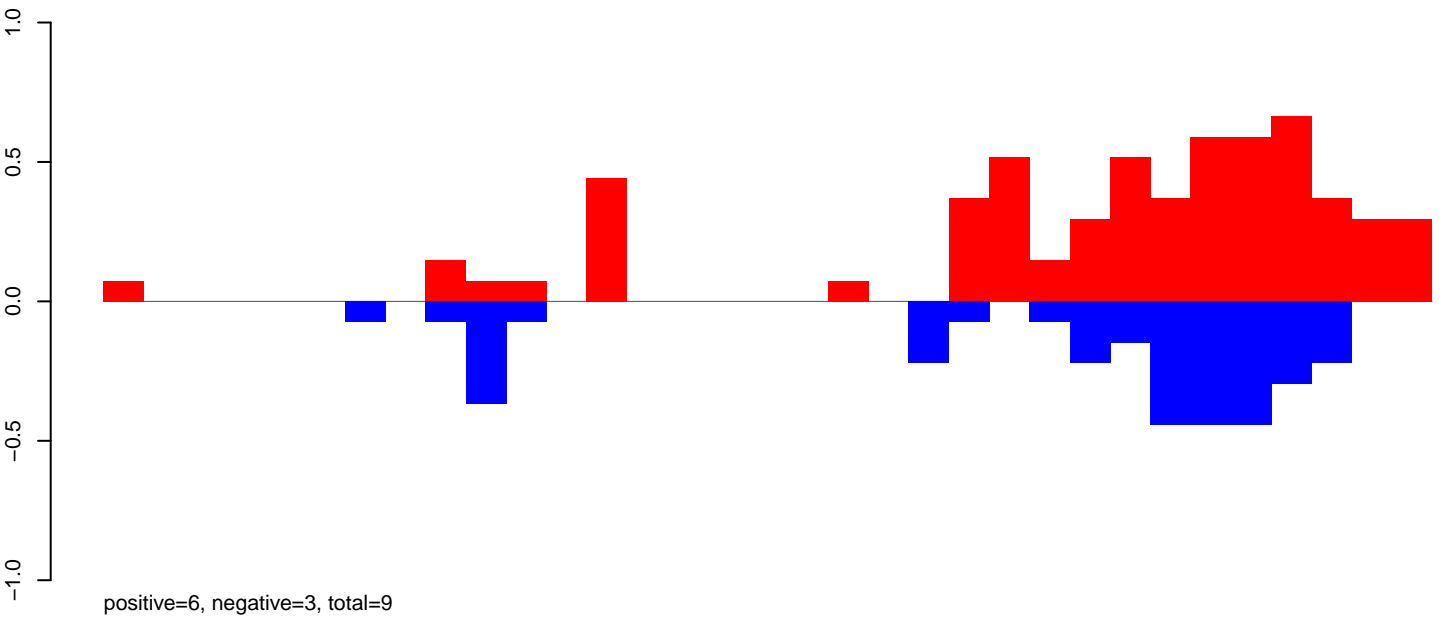
Window size=25, length=959, TE@rnd-4_family-237@DNA_Zator@227:1-959

0 200 400 600 800 1000

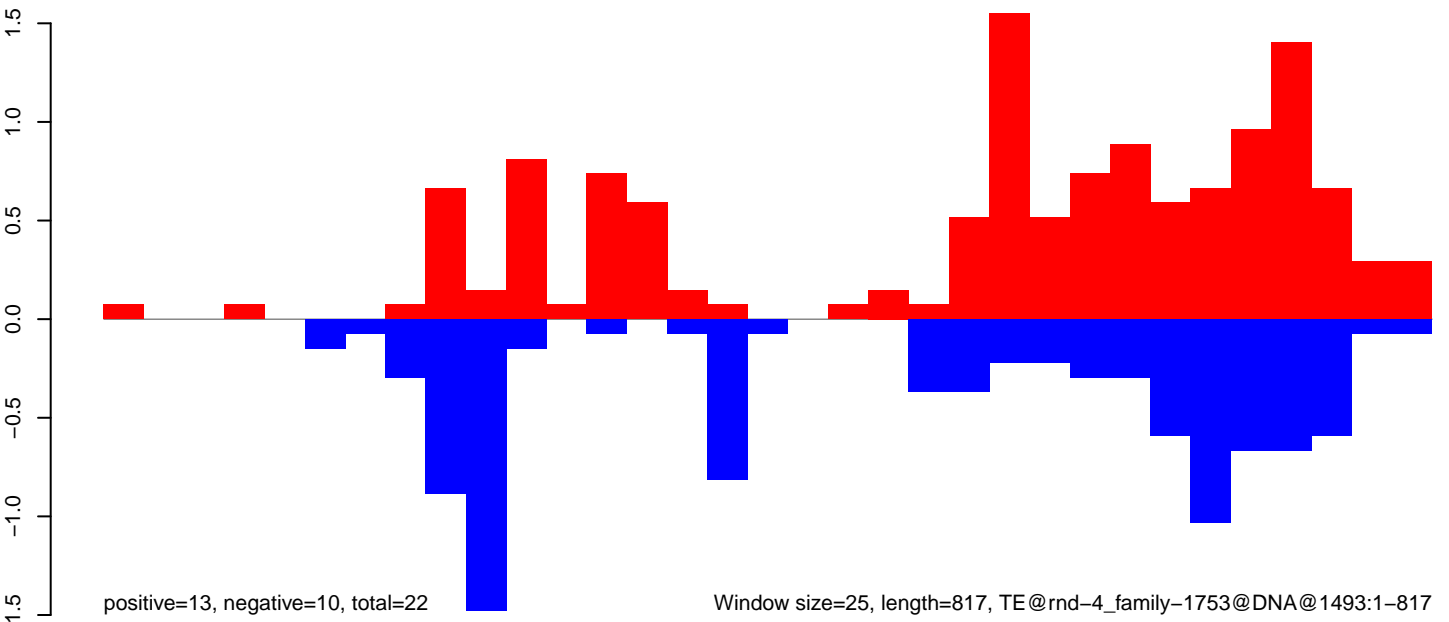
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

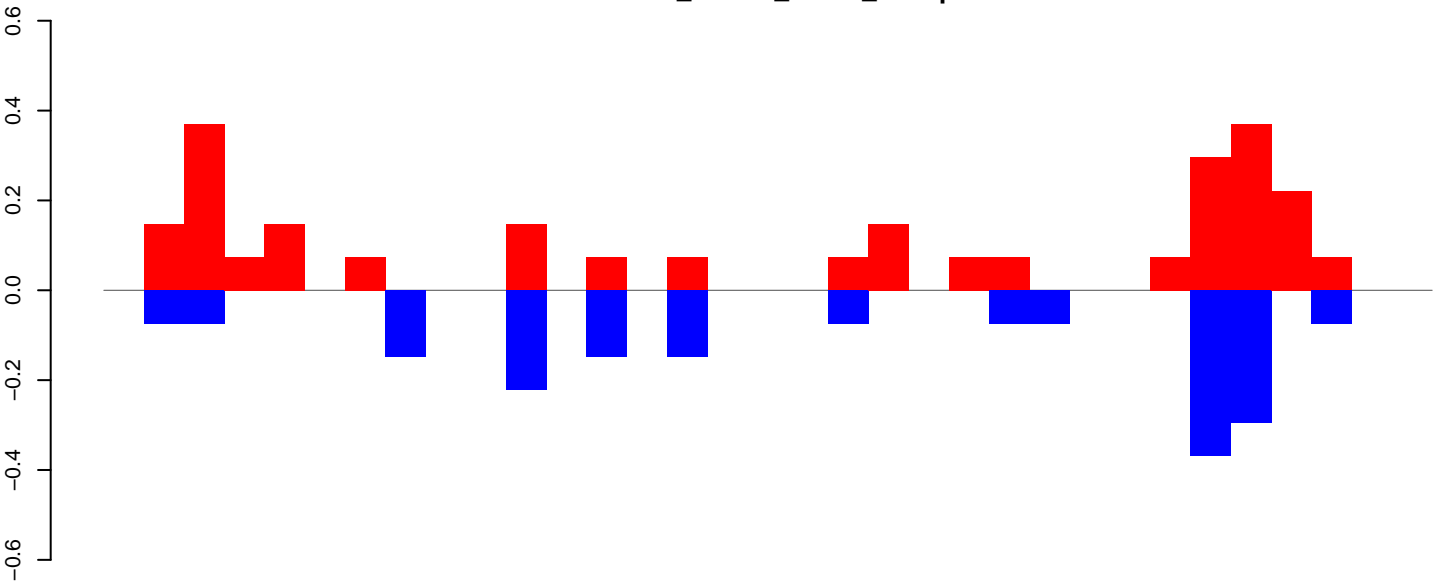


CuQuin_Testes_NL.rep



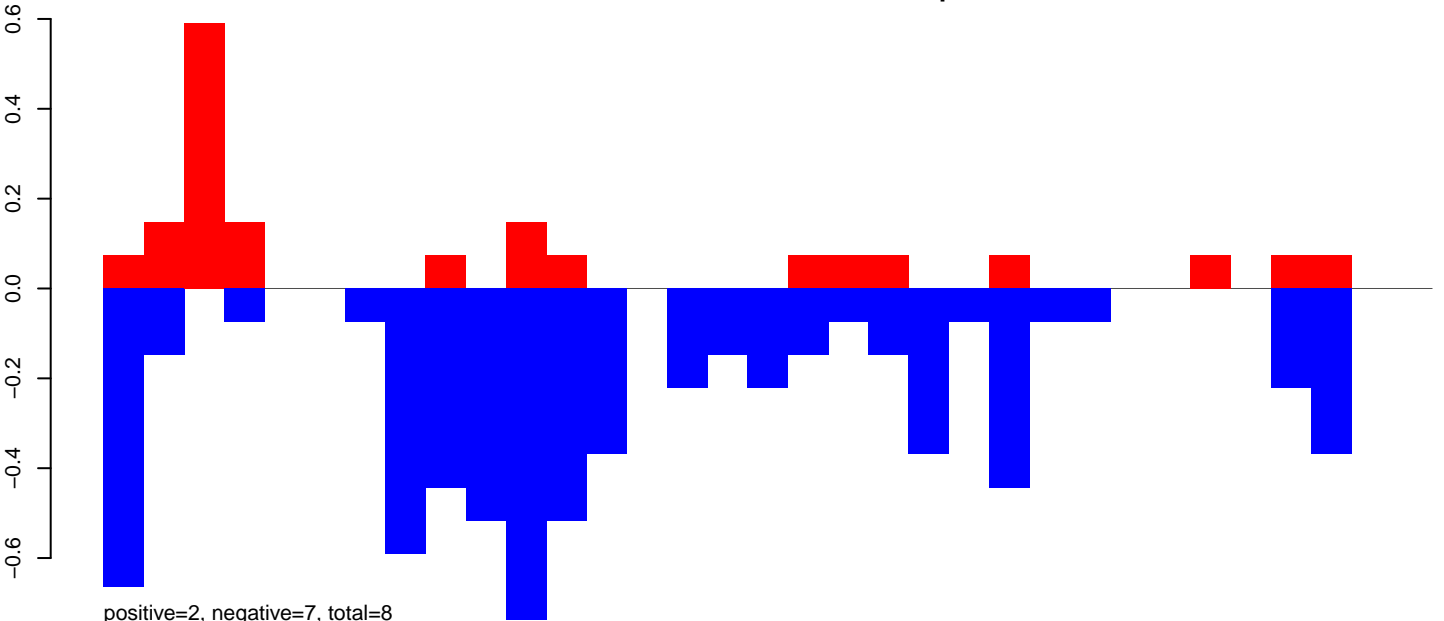
Window size=25, length=817, TE@rnd-4_family-1753@DNA@1493:1-817

CuQuin_Testes_NL.18_23.rep



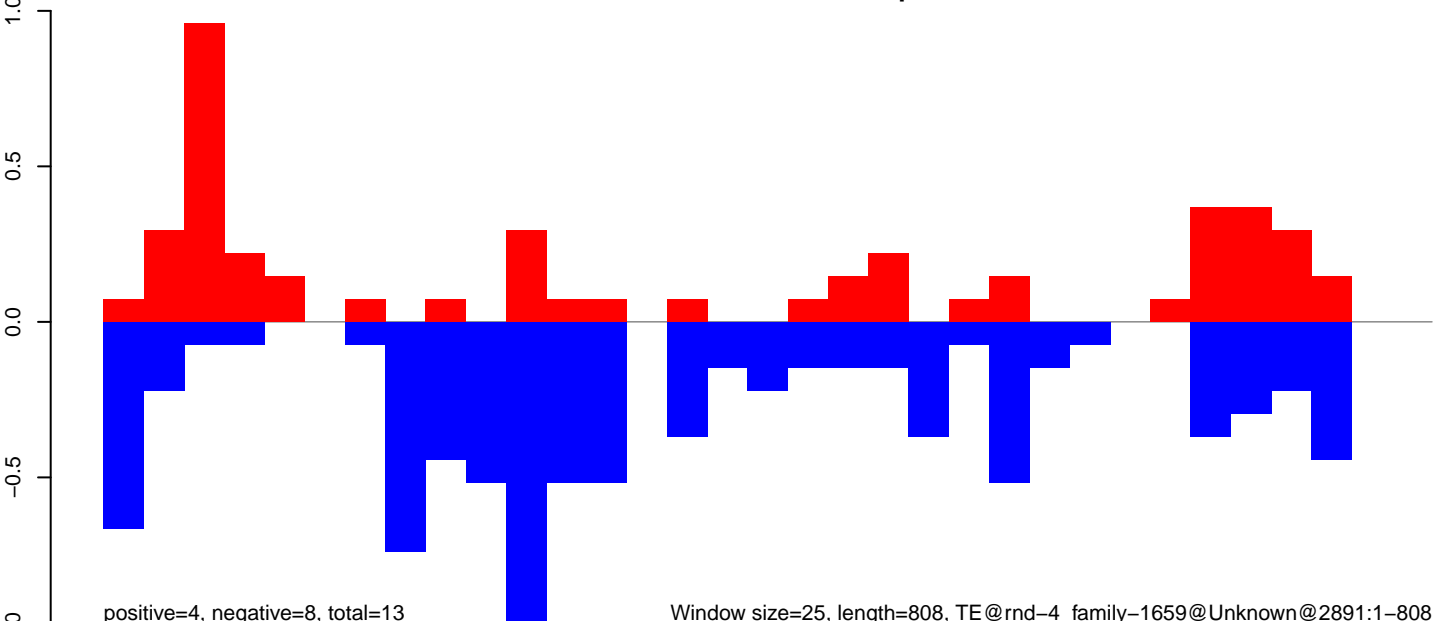
positive=3, negative=2, total=4

CuQuin_Testes_NL.24_35.rep



positive=2, negative=7, total=8

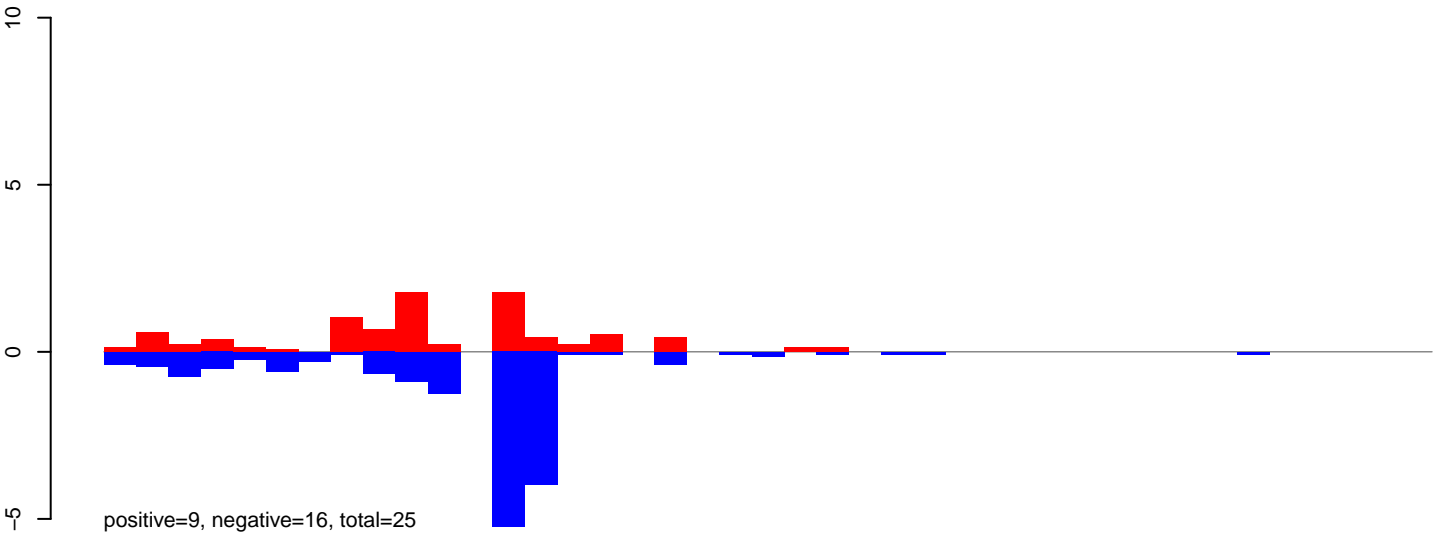
CuQuin_Testes_NL.rep



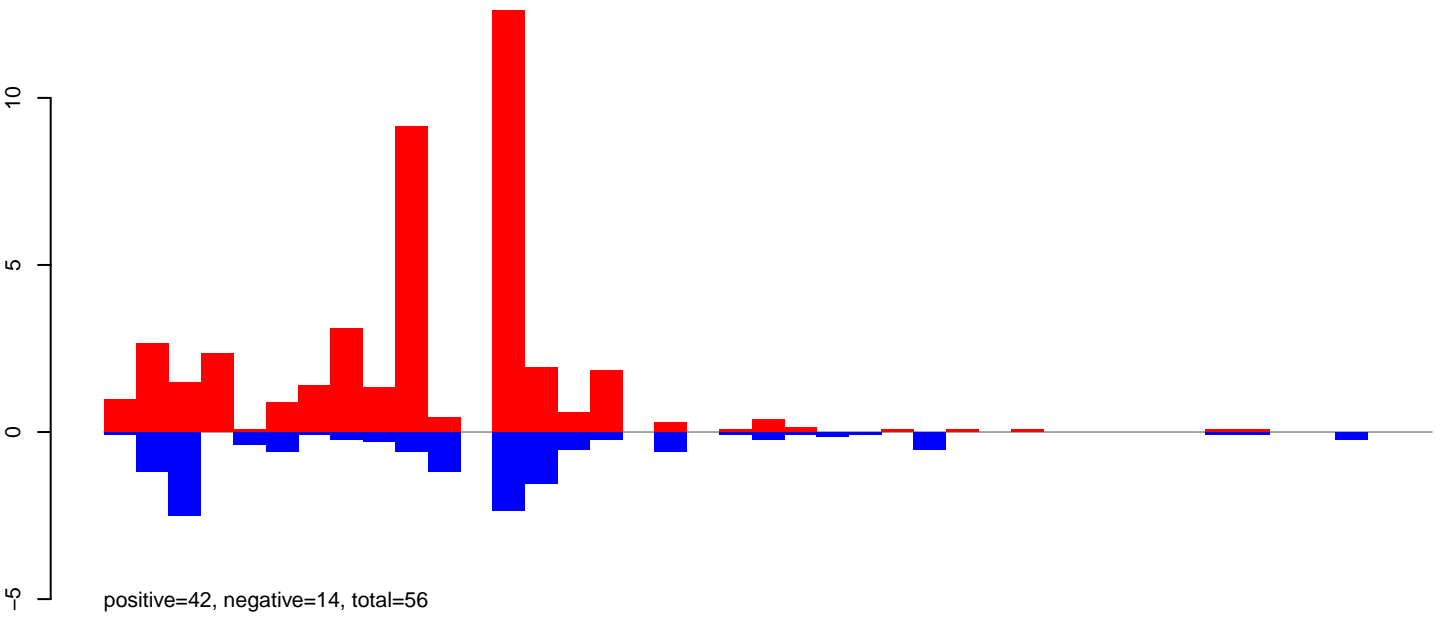
positive=4, negative=8, total=13

Window size=25, length=808, TE@rnd-4_family-1659@Unknown@2891:1-808

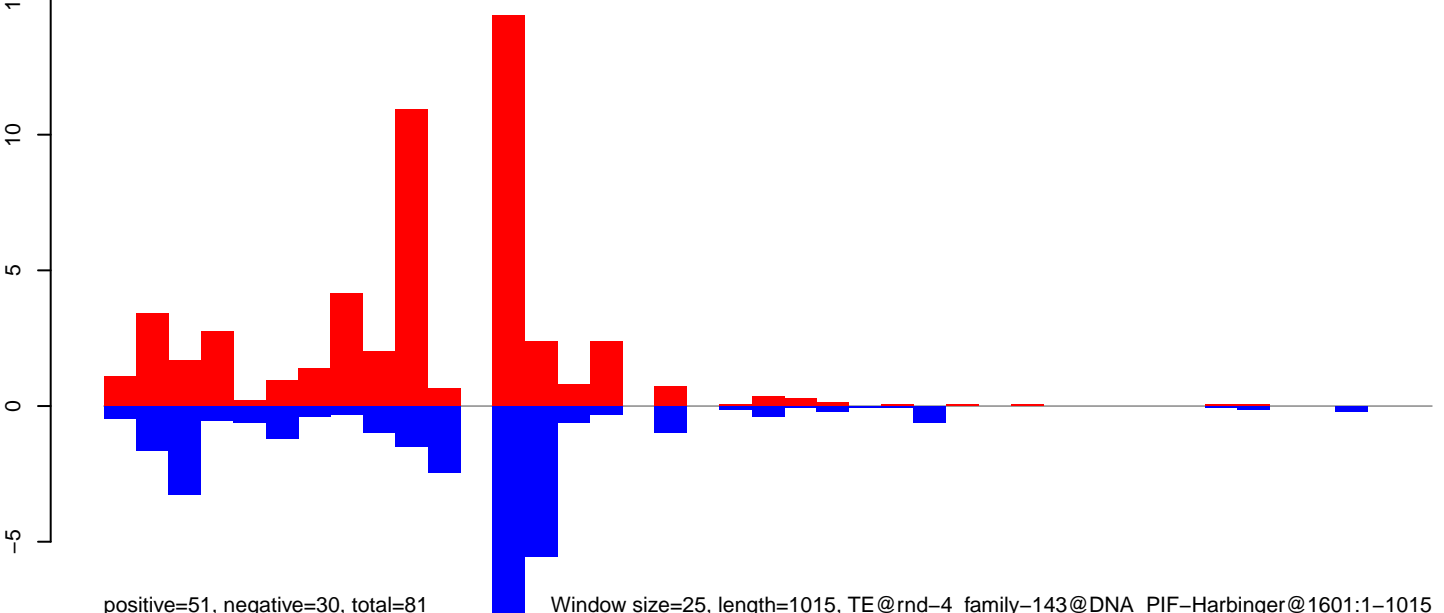
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

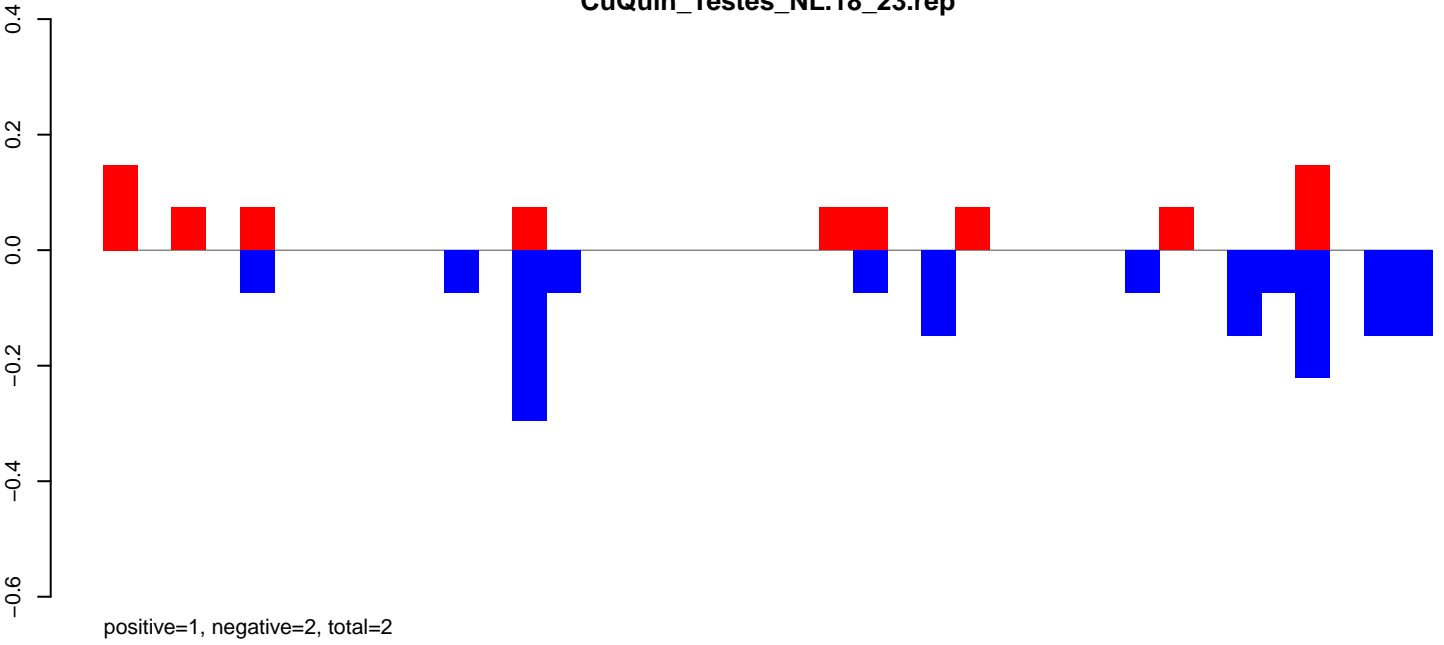


CuQuin_Testes_NL.rep

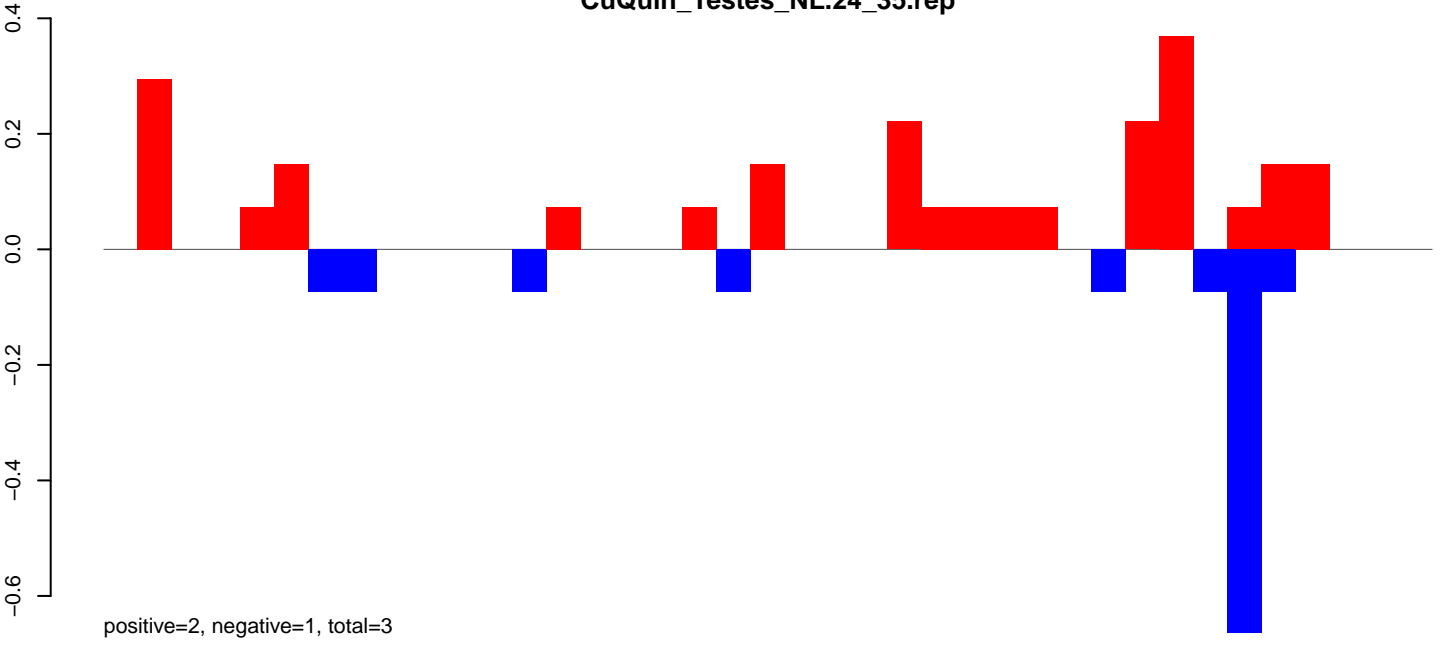


Window size=25, length=1015, TE@rnd-4_family-143@DNA_PIF-Harbinger@1601:1-1015

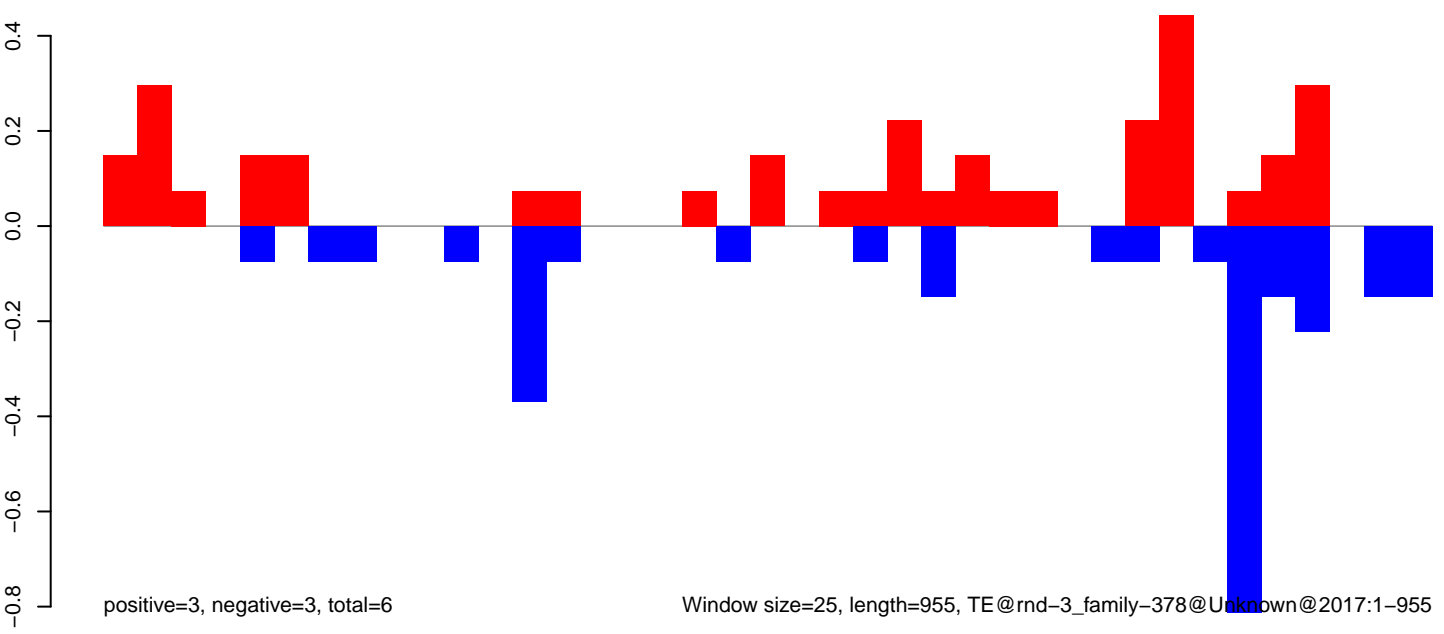
CuQuin_Testes_NL.18_23.rep



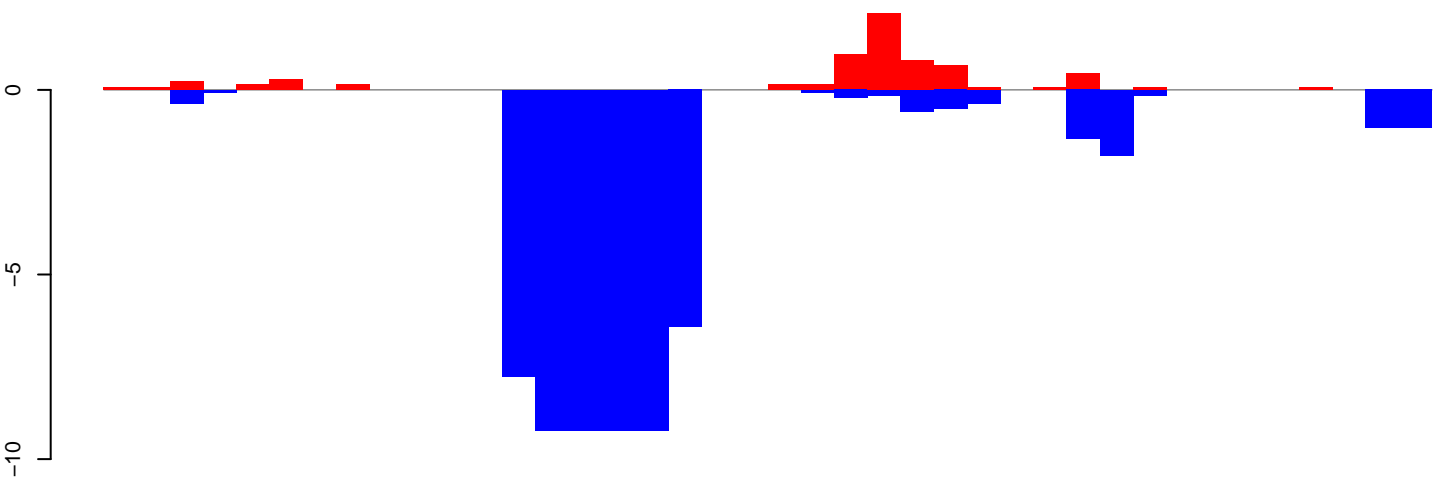
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

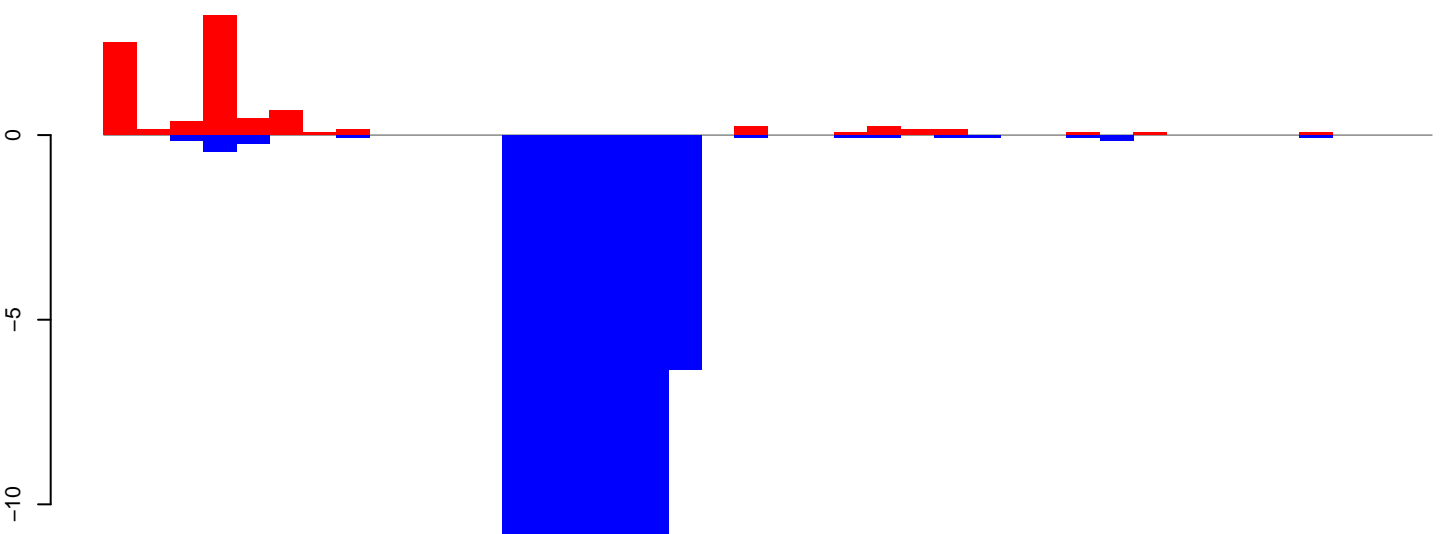


CuQuin_Testes_NL.18_23.rep



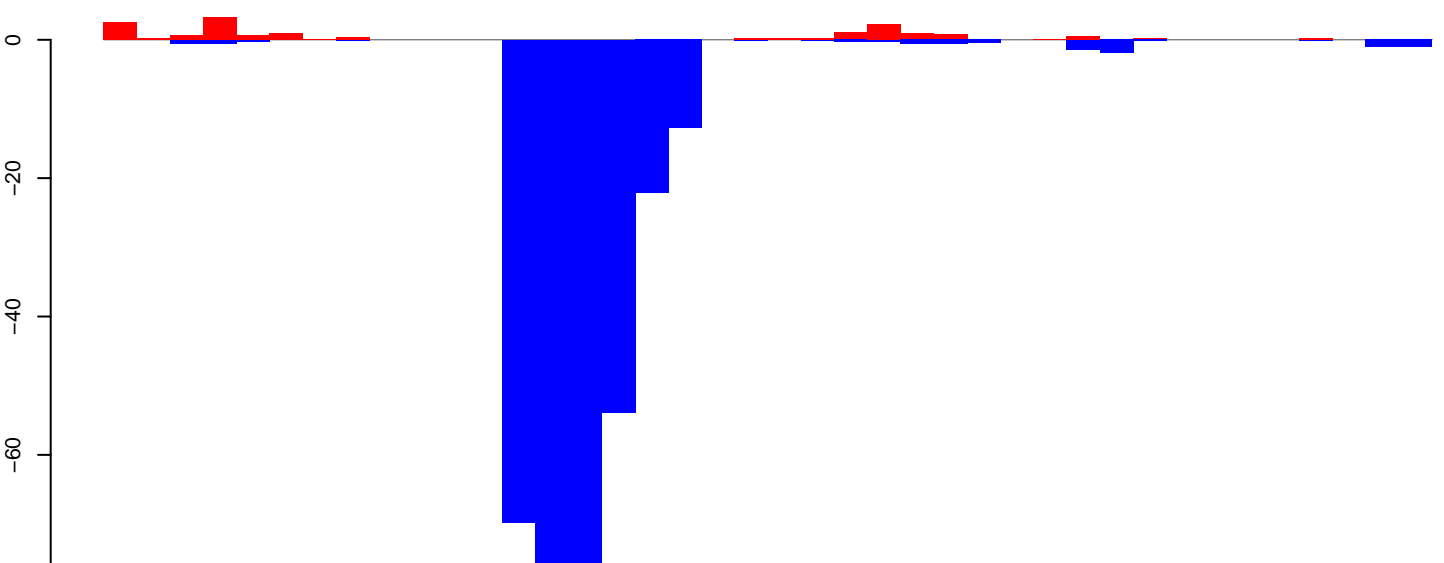
positive=6, negative=59, total=65

CuQuin_Testes_NL.24_35.rep



positive=9, negative=70, total=79

CuQuin_Testes_NL.rep

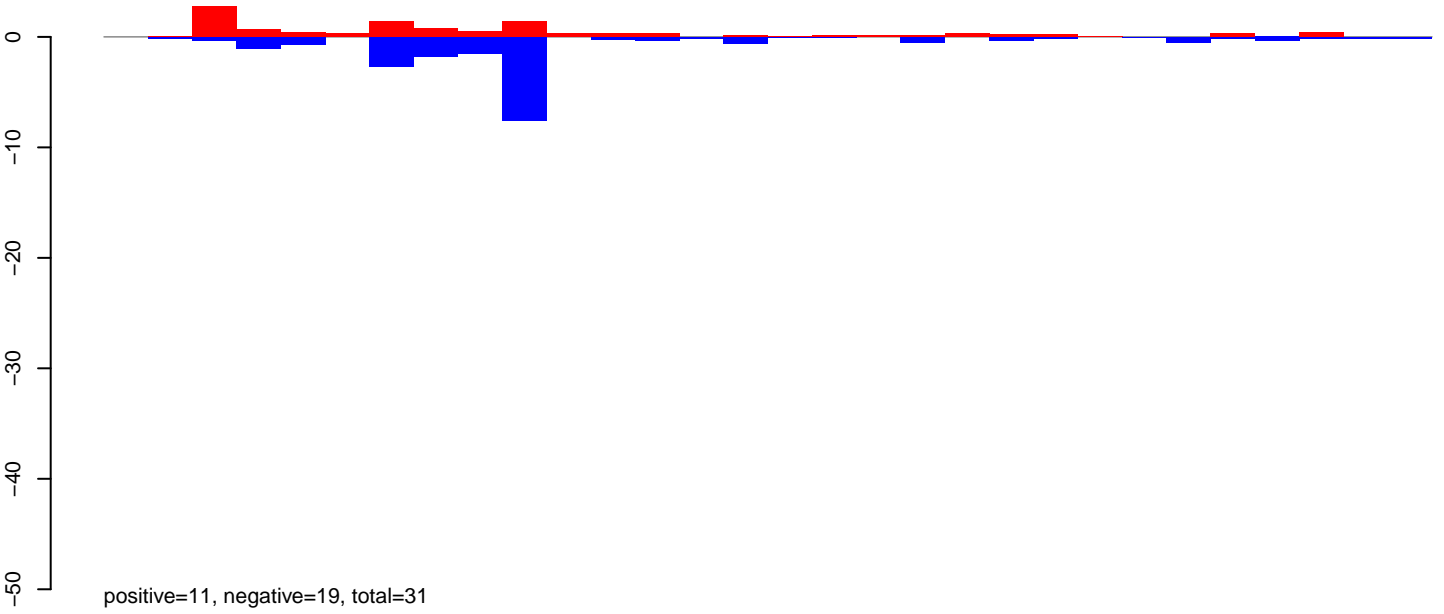


positive=15, negative=334, total=349

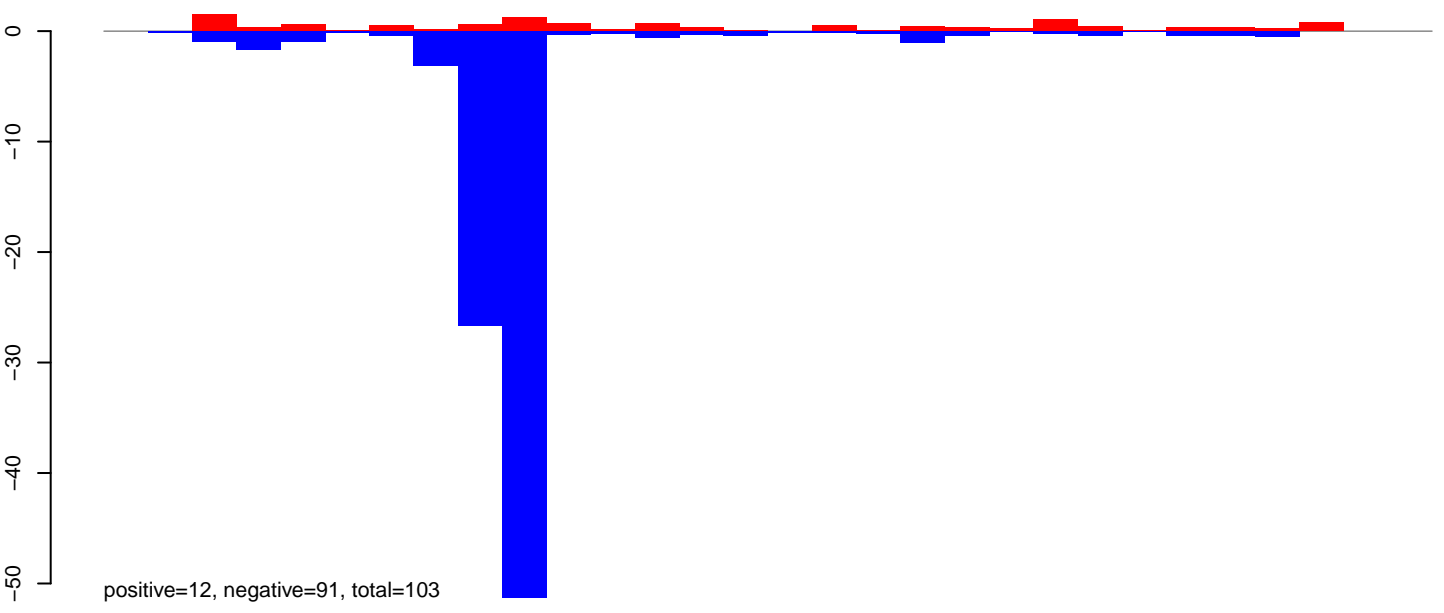
Window size=25, length=984, TE@rnd-3_family-234@DNA_hAT-Ac@2018:1-984

0 200 400 600 800 1000

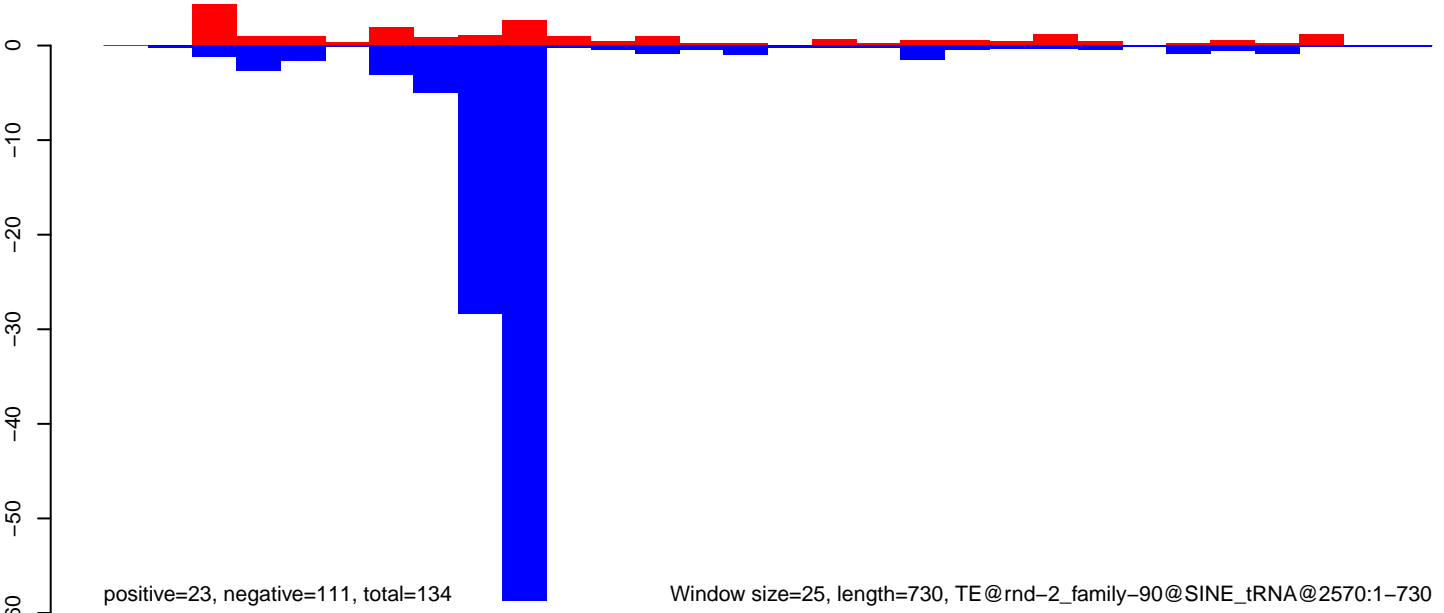
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

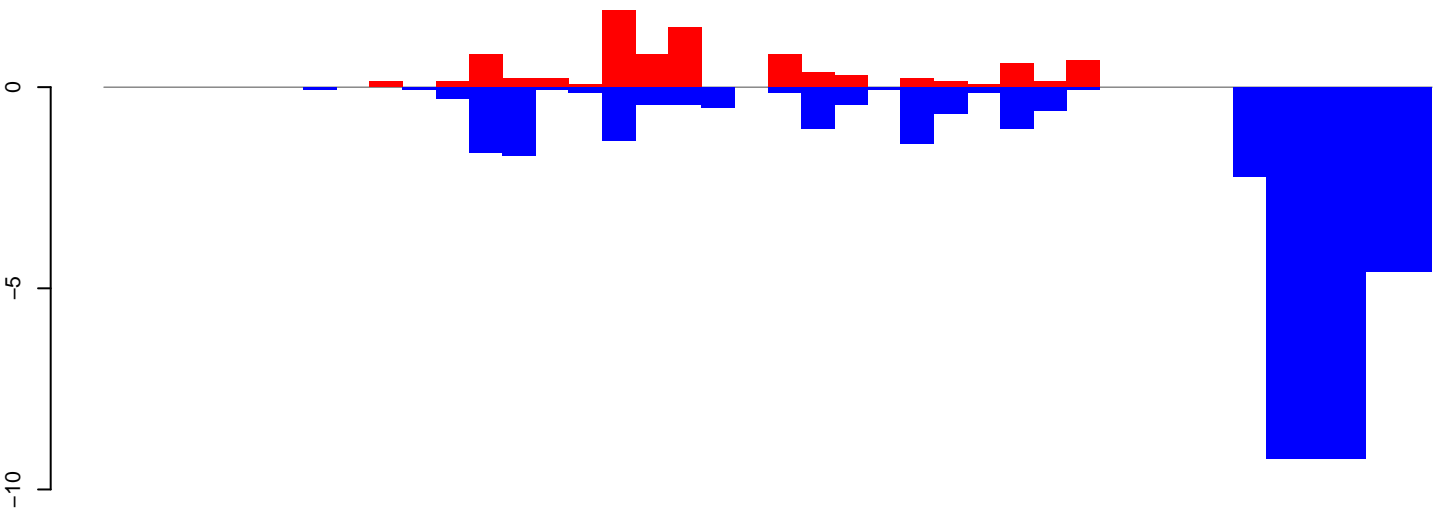


CuQuin_Testes_NL.rep



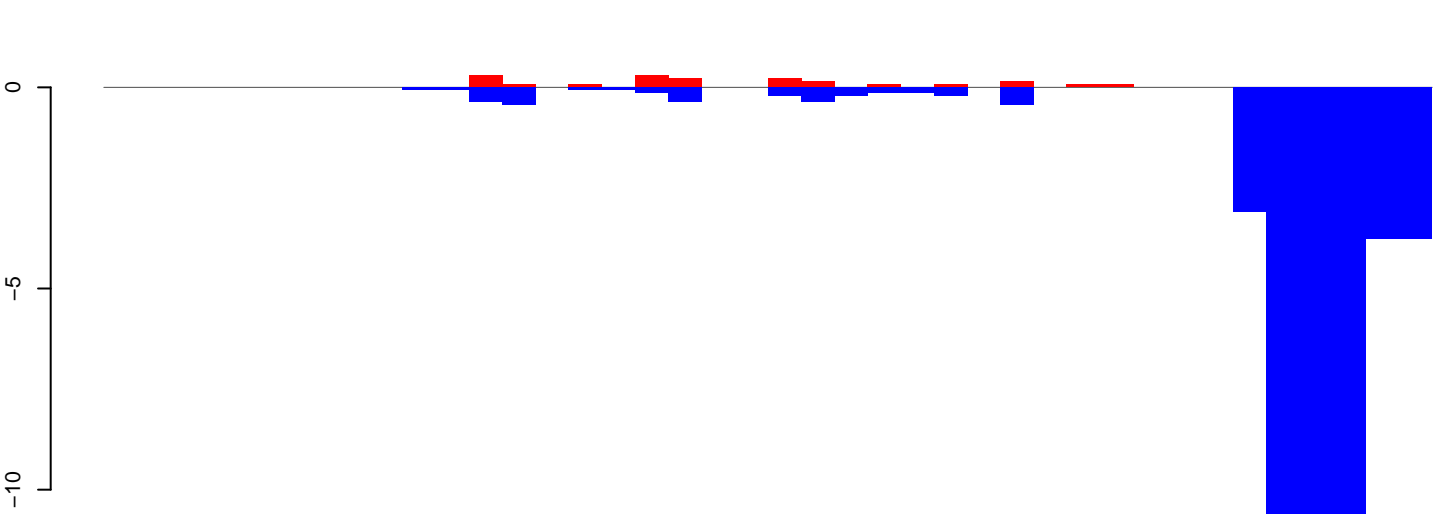
Window size=25, length=730, TE@rnd-2_family-90@SINE_tRNA@2570:1-730

CuQuin_Testes_NL.18_23.rep



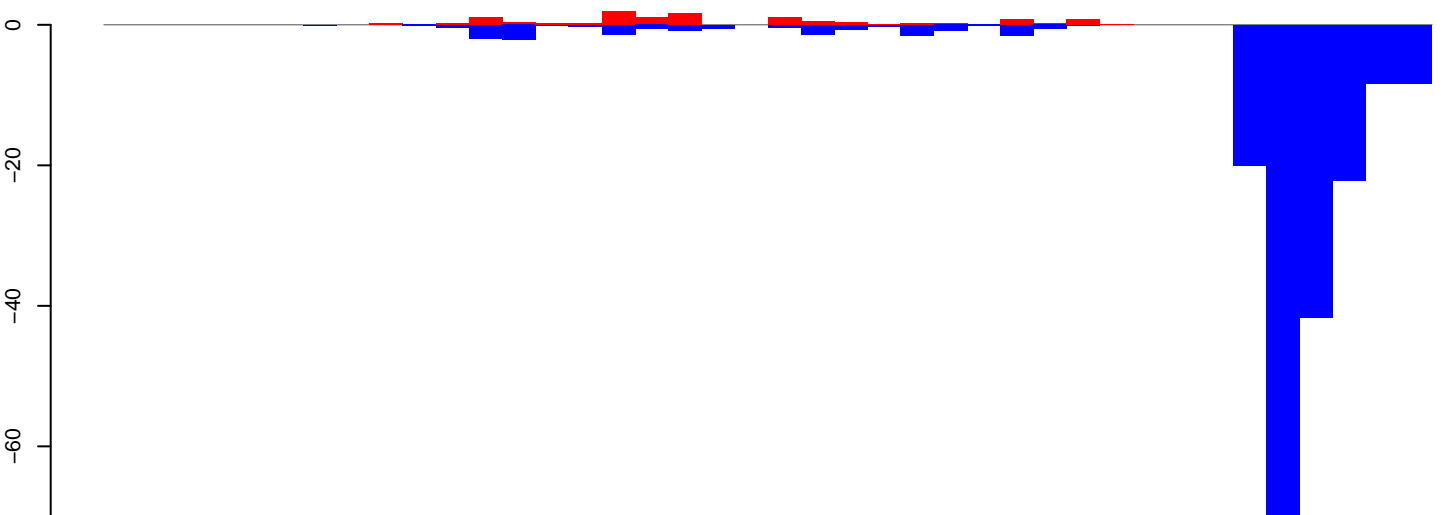
positive=9, negative=51, total=61

CuQuin_Testes_NL.24_35.rep



positive=2, negative=53, total=55

CuQuin_Testes_NL.rep

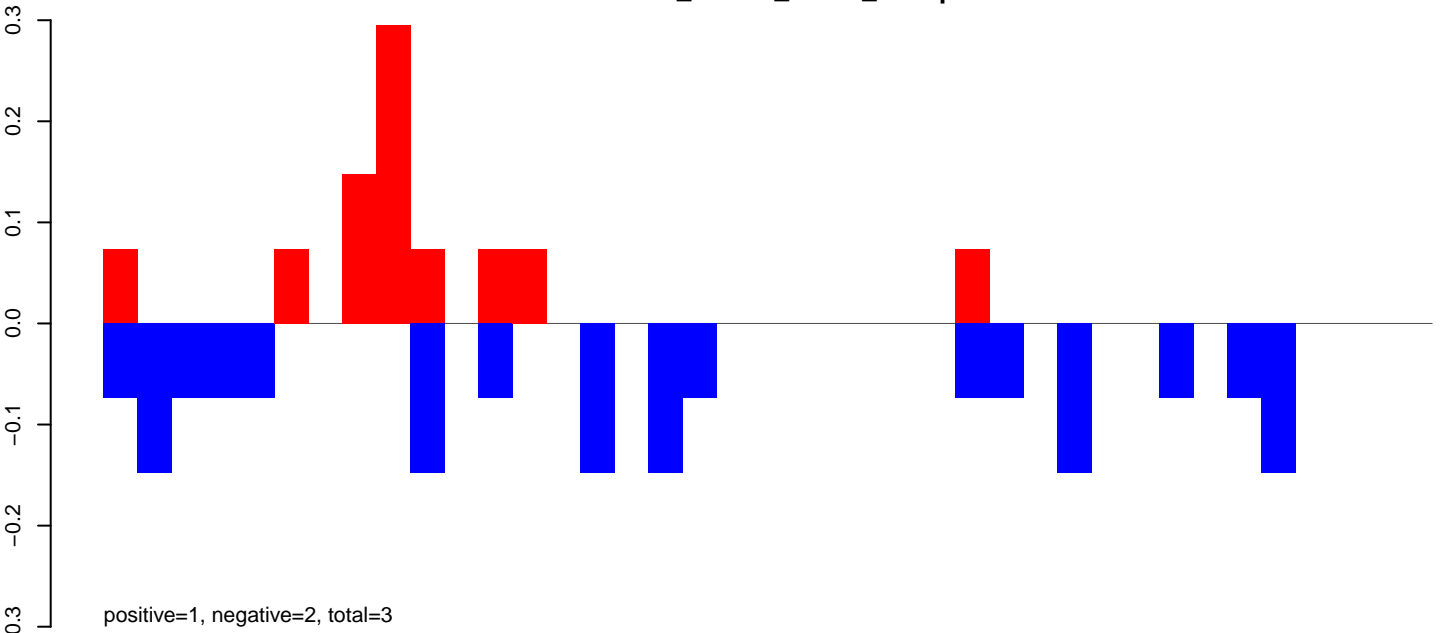


positive=11, negative=199, total=210

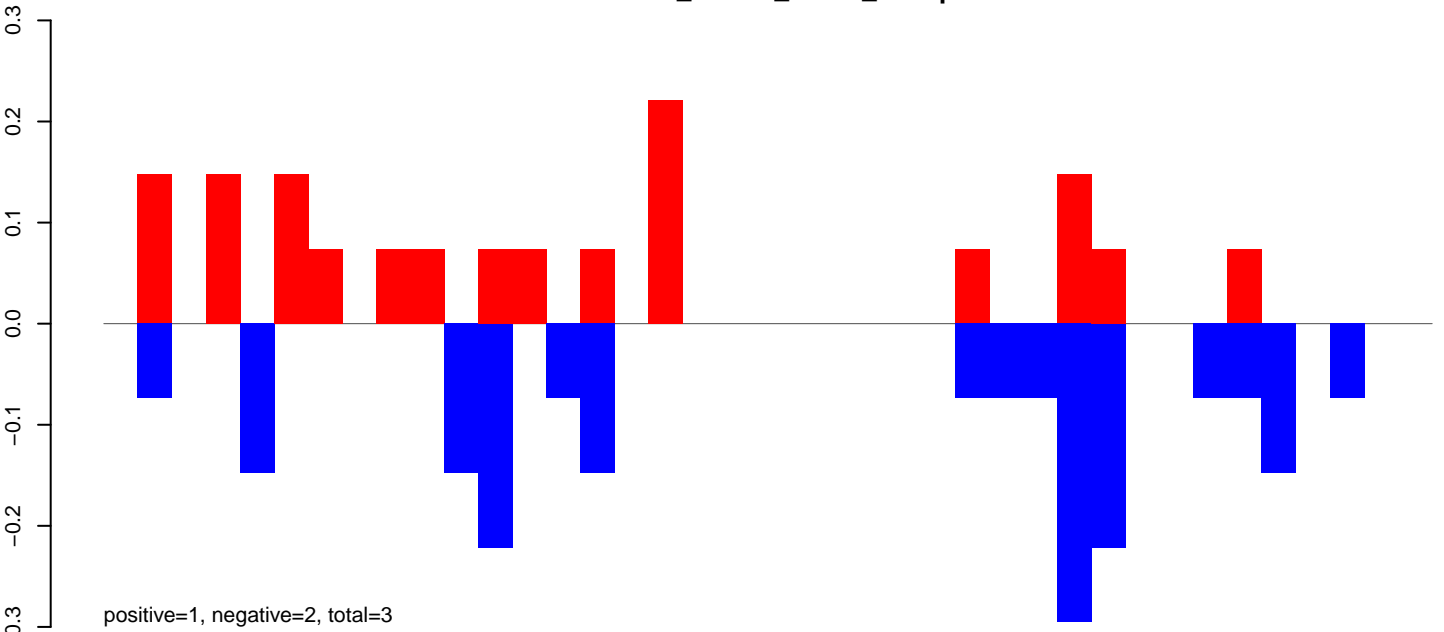
Window size=25, length=984, TE@rnd-2_family-538@DNA@647:1-984

0 200 400 600 800 1000

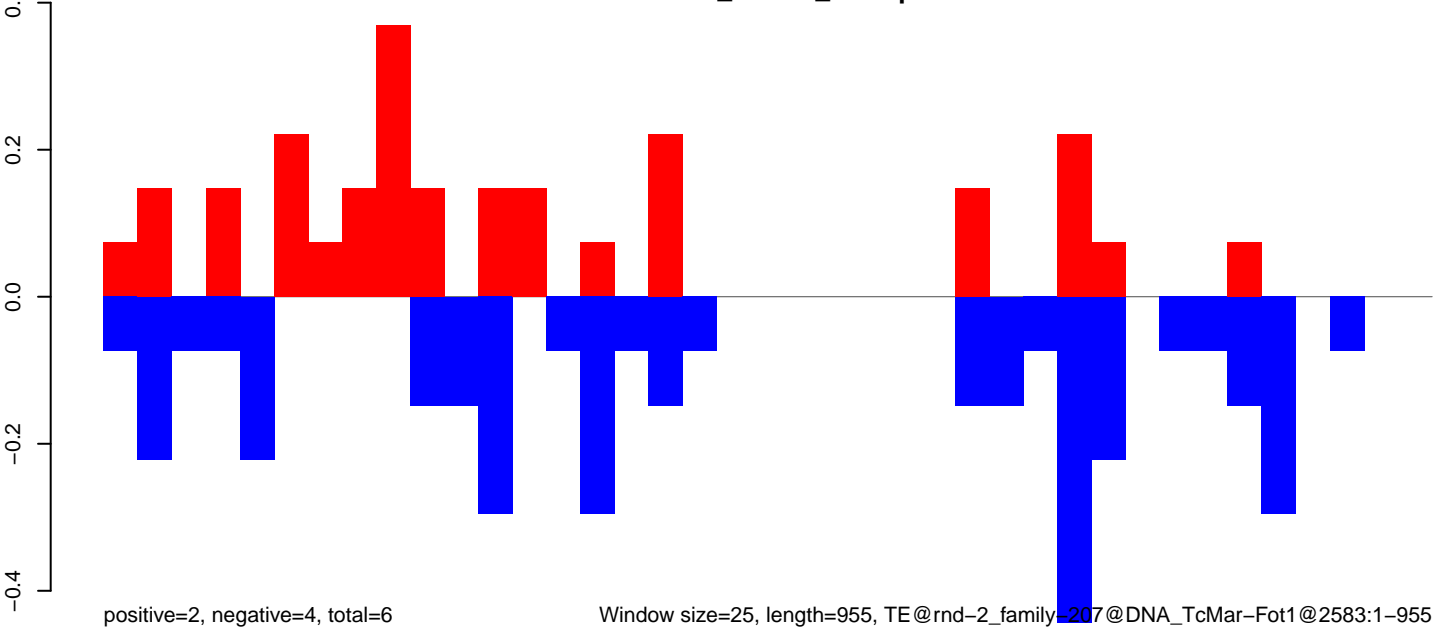
CuQuin_Testes_NL.18_23.rep



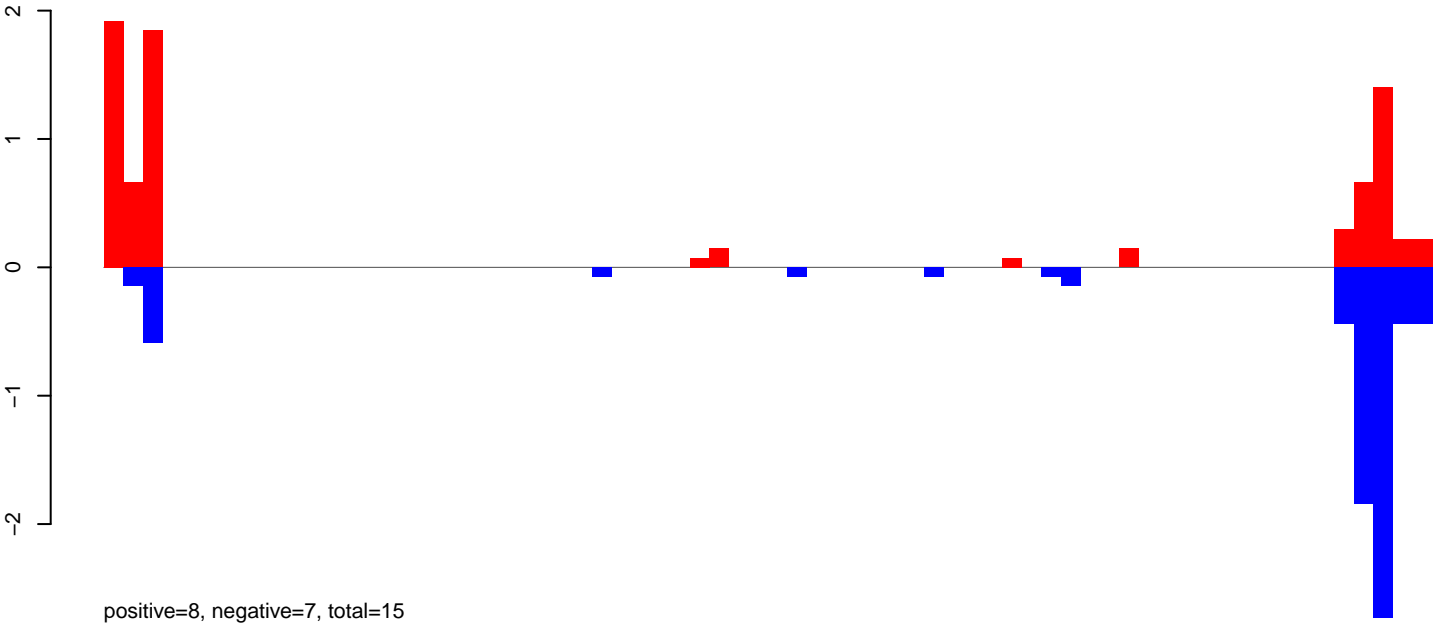
CuQuin_Testes_NL.24_35.rep



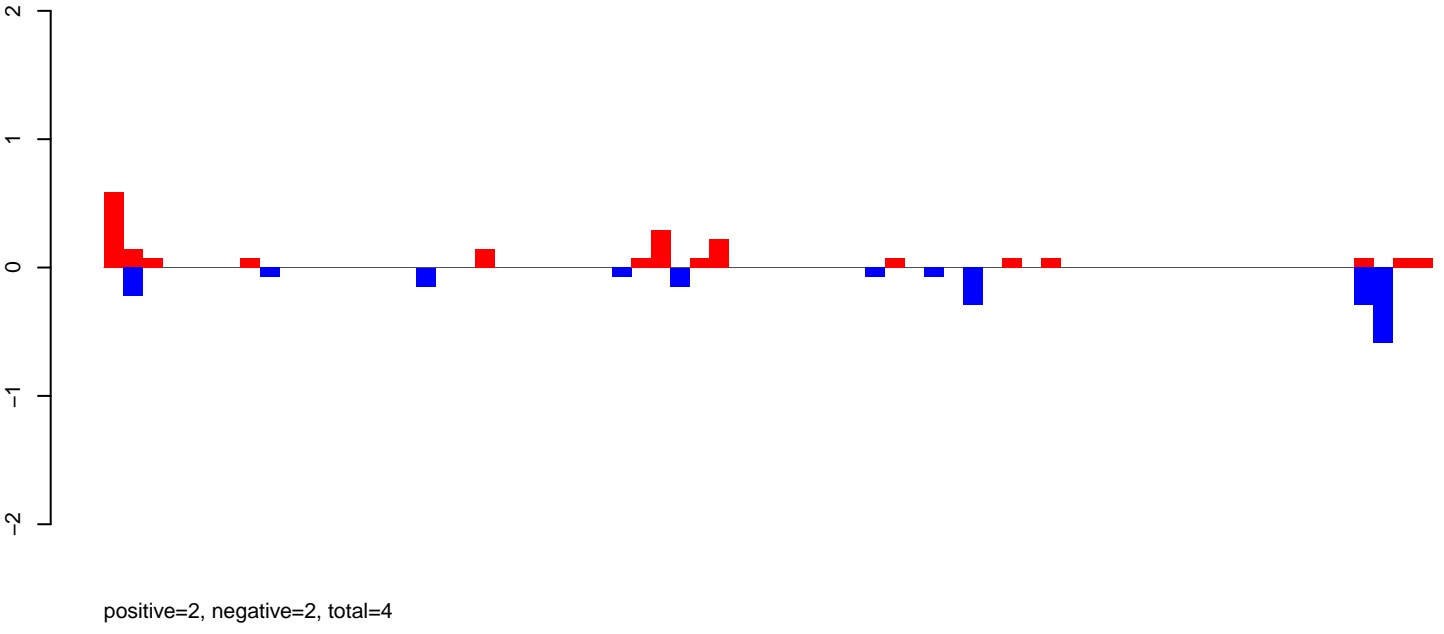
CuQuin_Testes_NL.rep



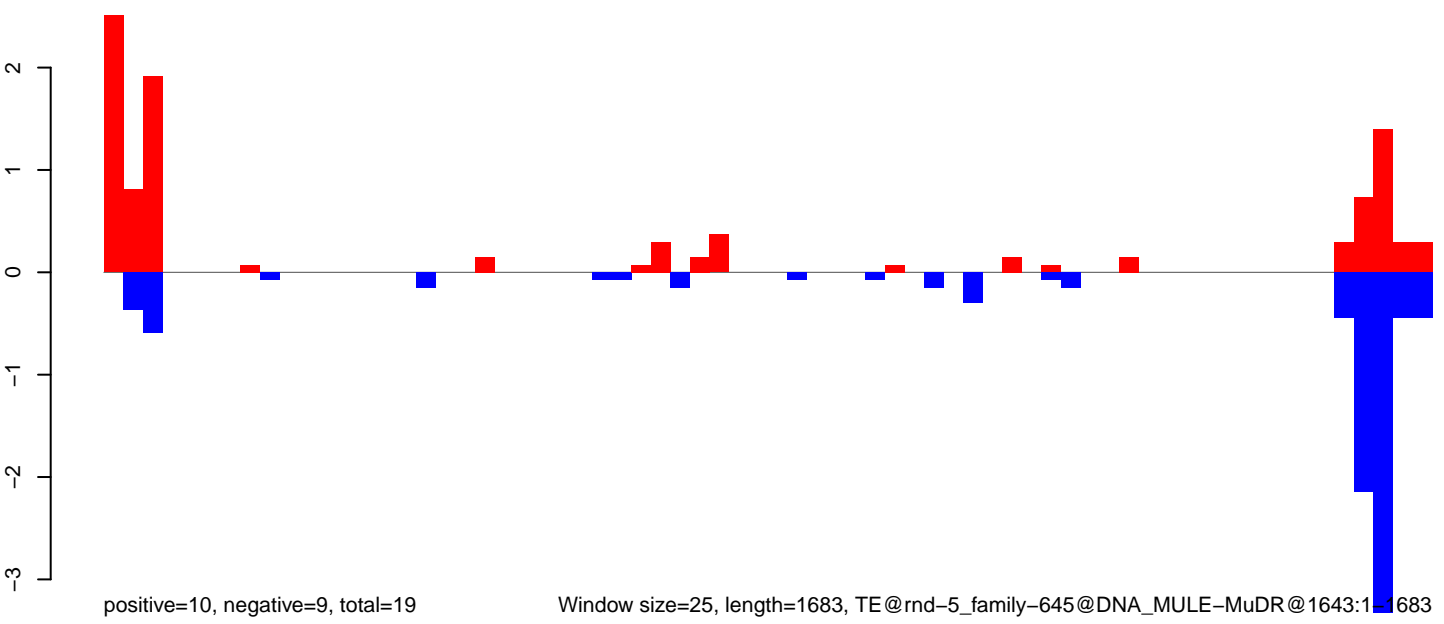
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



CuQuin_Testes_NL.18_23.rep



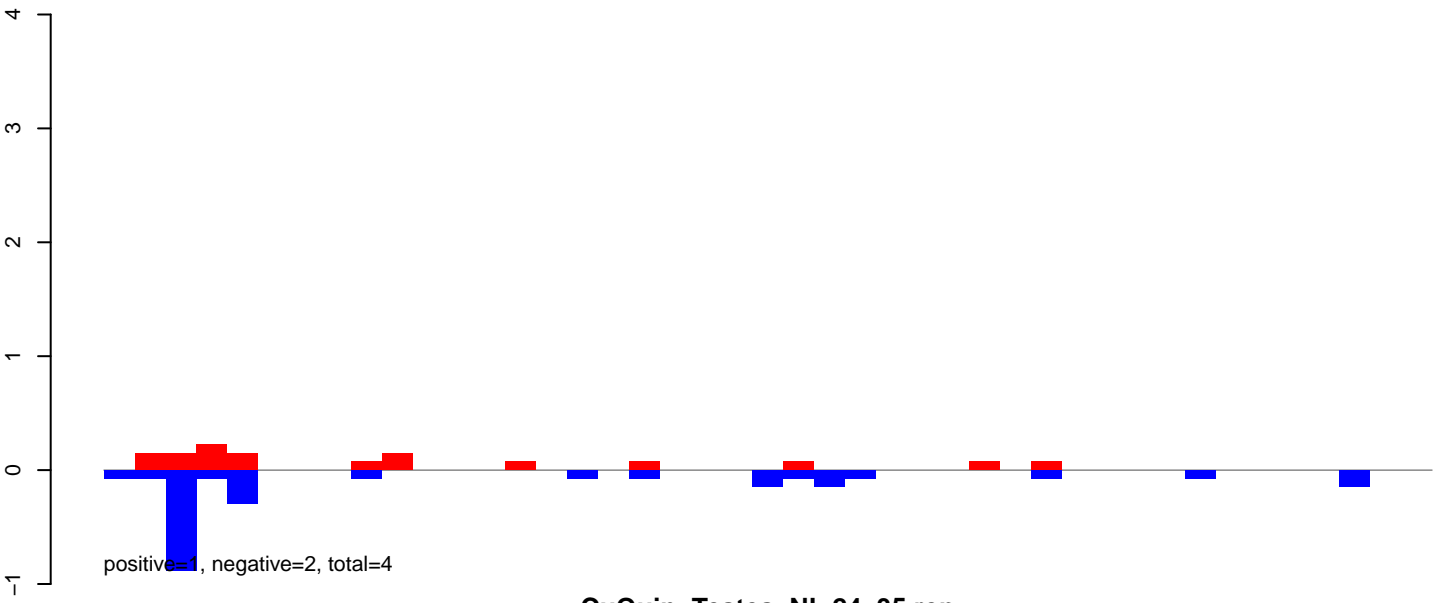
CuQuin_Testes_NL.24_35.rep



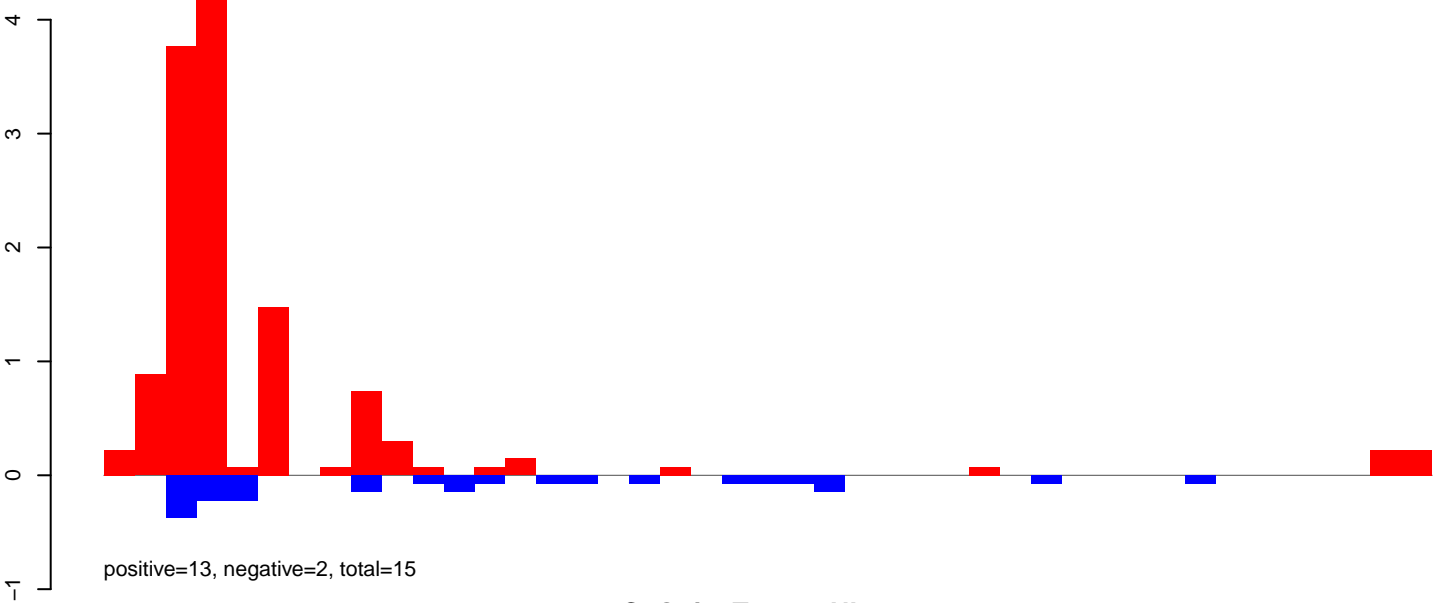
CuQuin_Testes_NL.rep



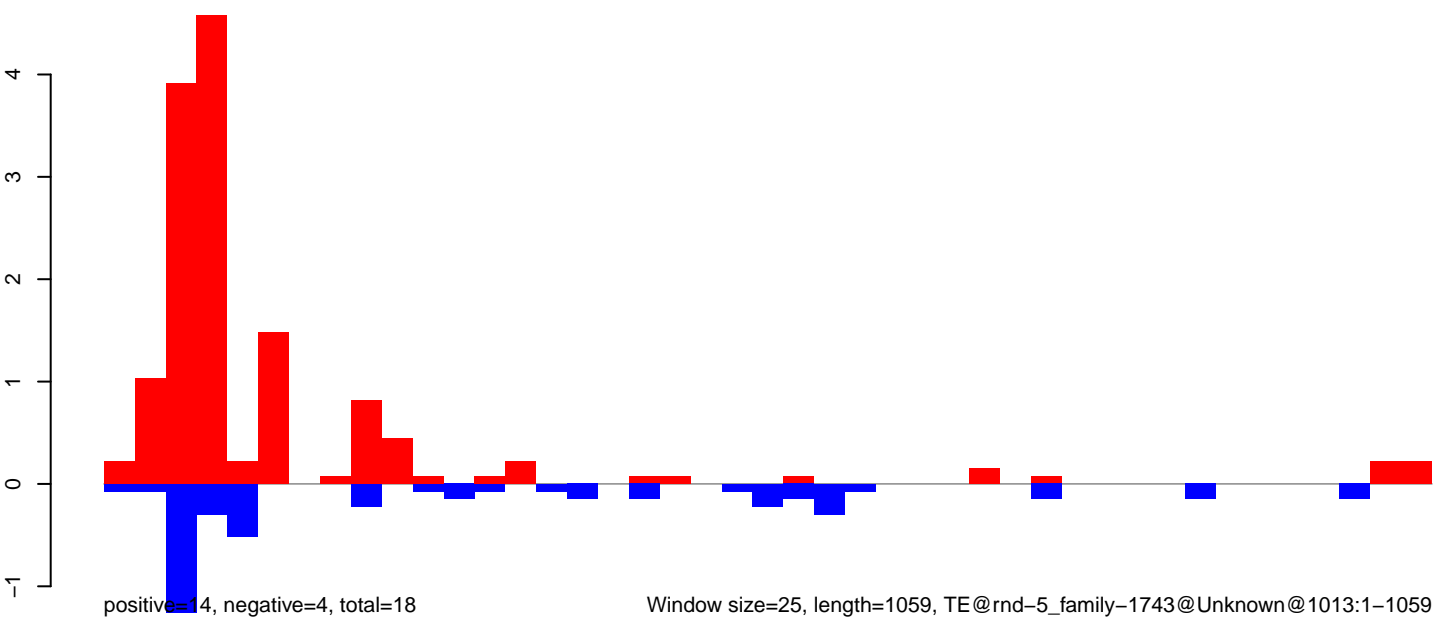
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

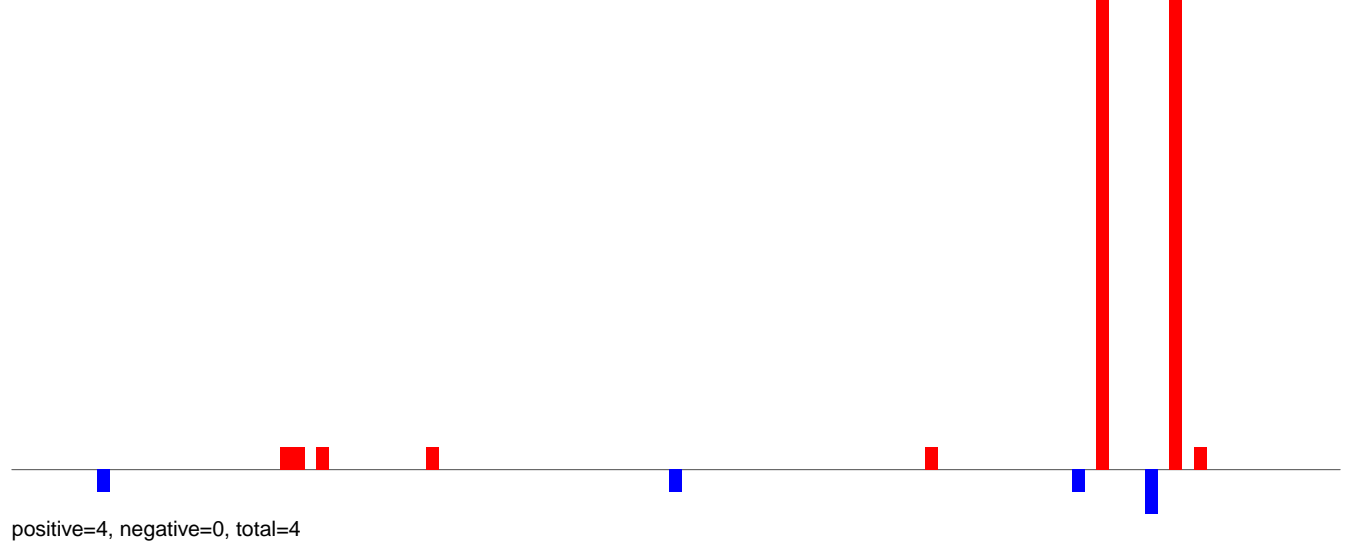


CuQuin_Testes_NL.rep

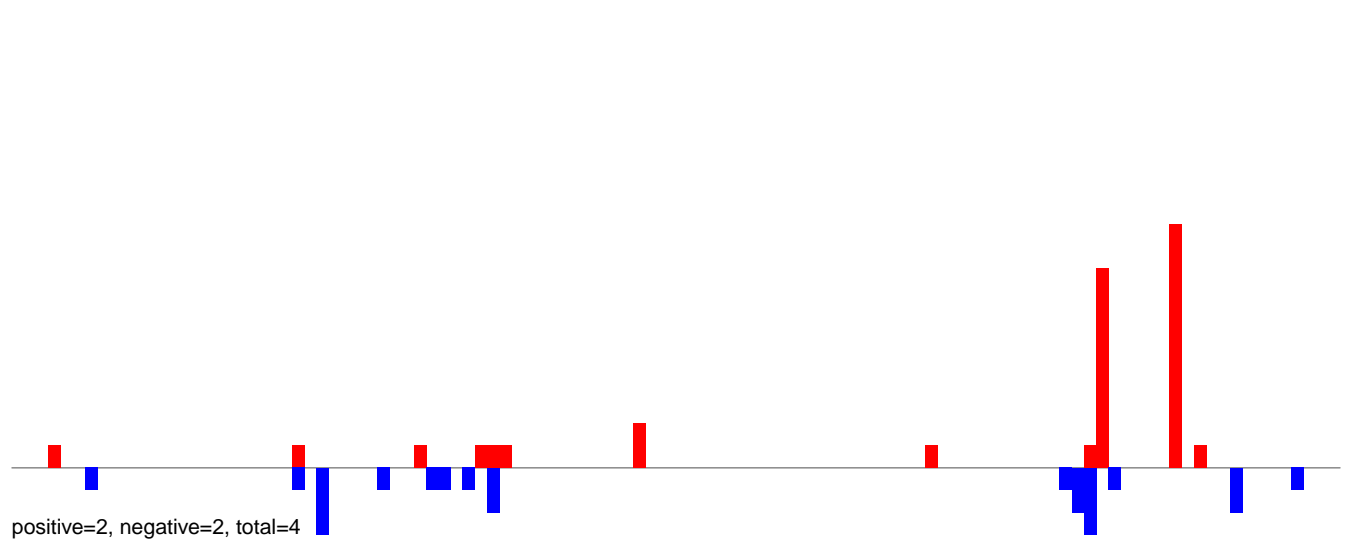


Window size=25, length=1059, TE@rnd-5_family-1743@Unknown@1013:1-1059

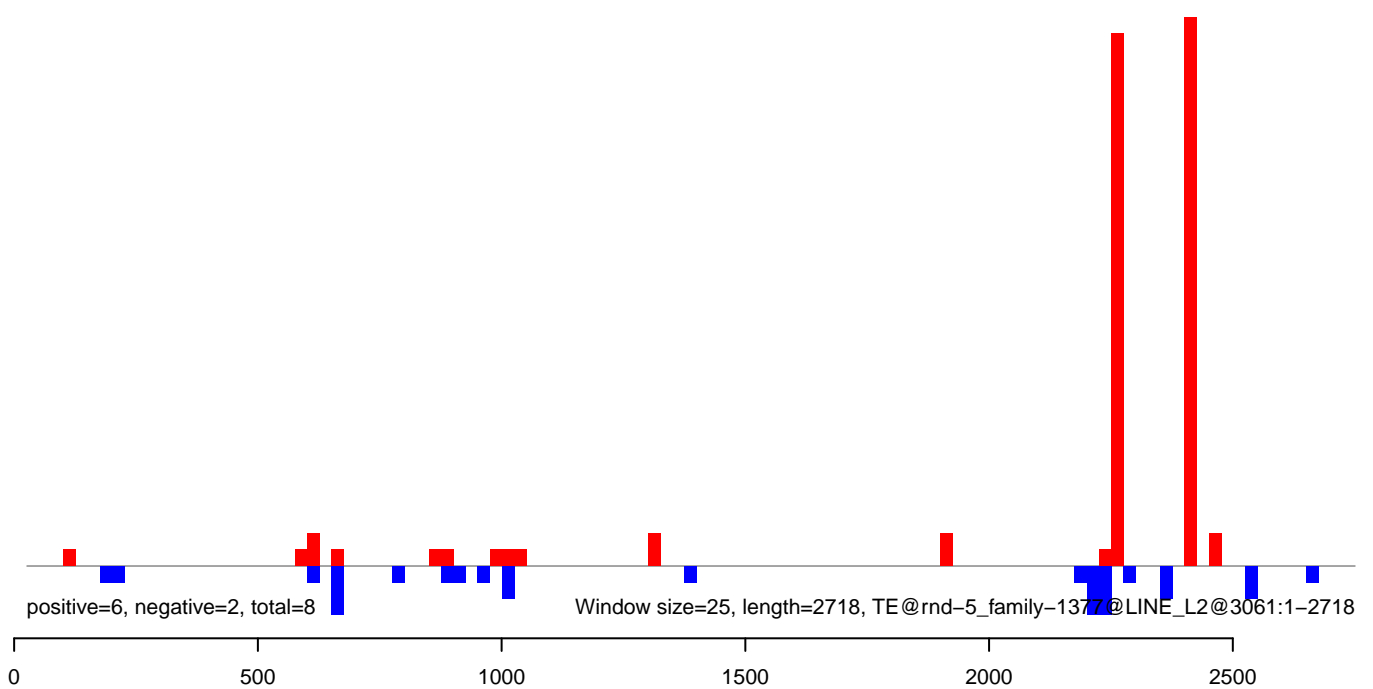
CuQuin_Testes_NL.18_23.rep



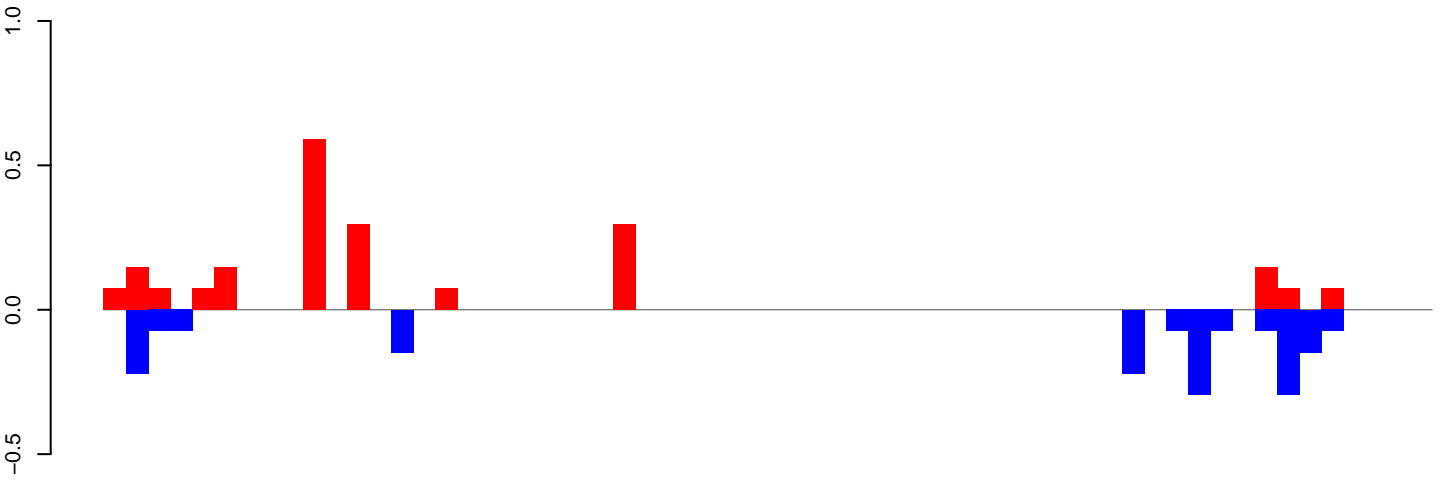
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

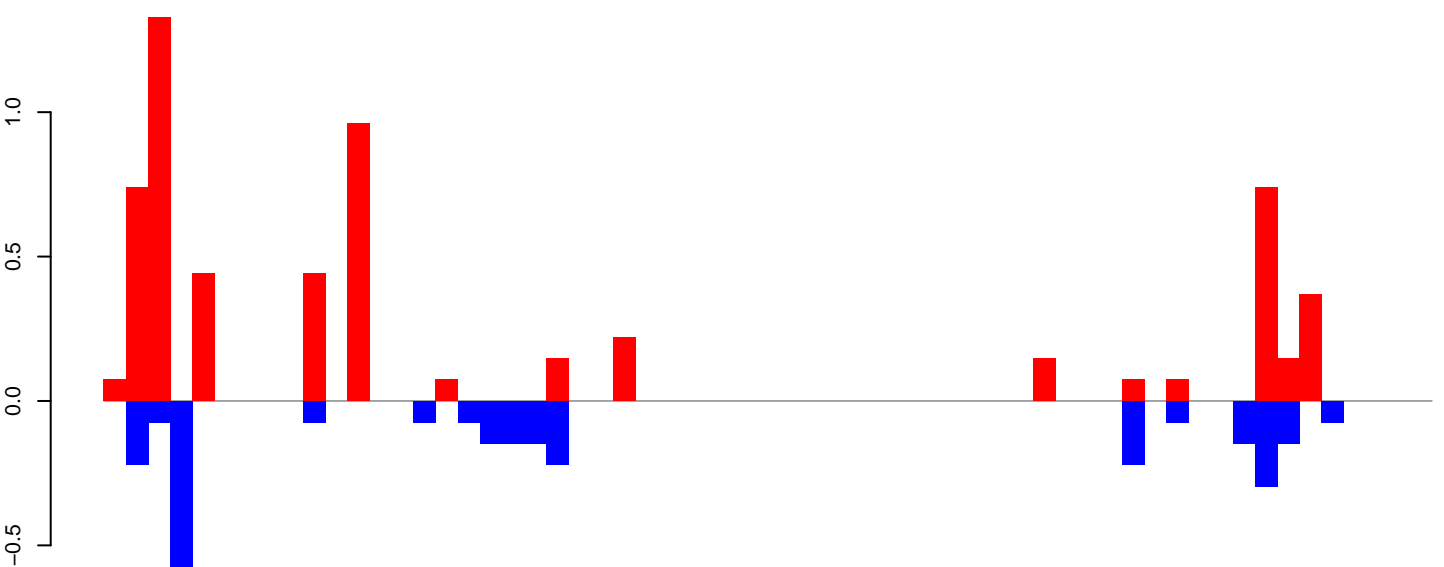


CuQuin_Testes_NL.18_23.rep



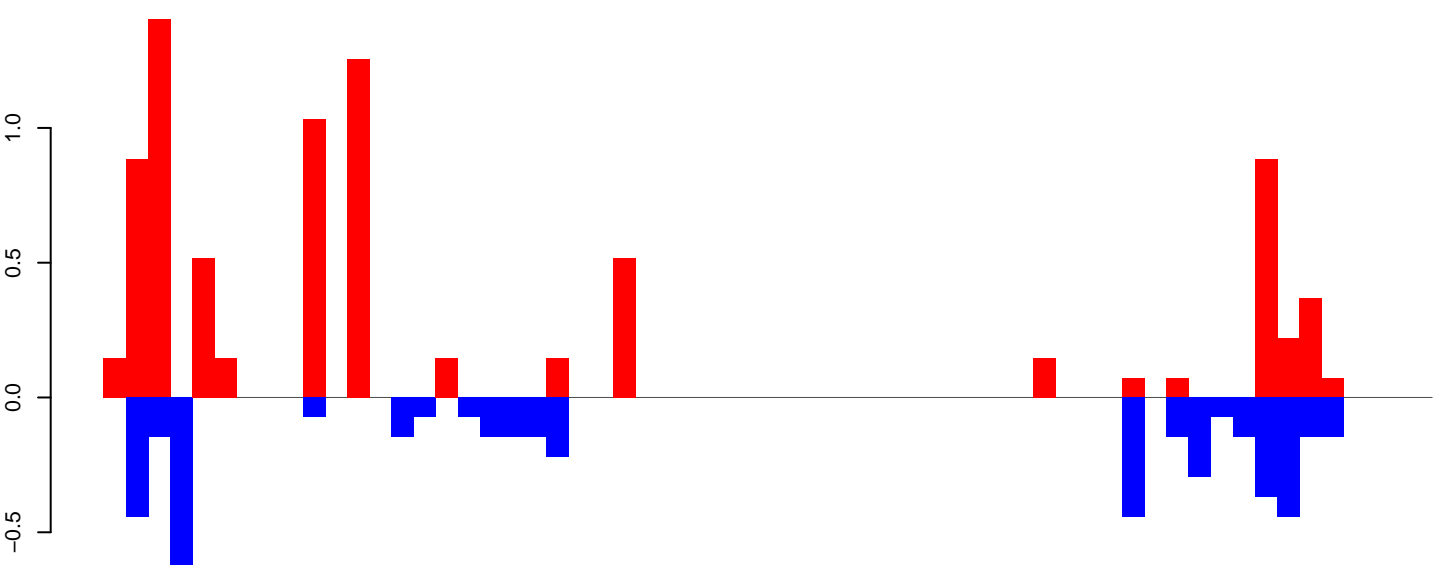
positive=2, negative=2, total=4

CuQuin_Testes_NL.24_35.rep



positive=6, negative=3, total=9

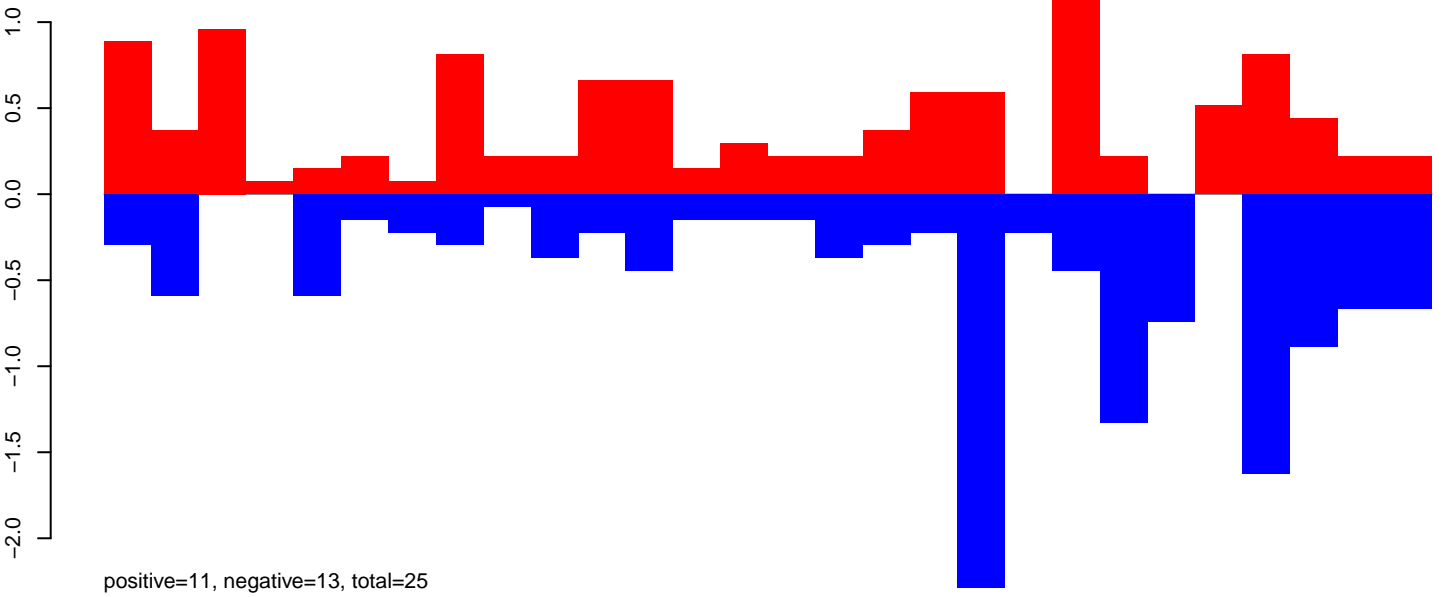
CuQuin_Testes_NL.rep



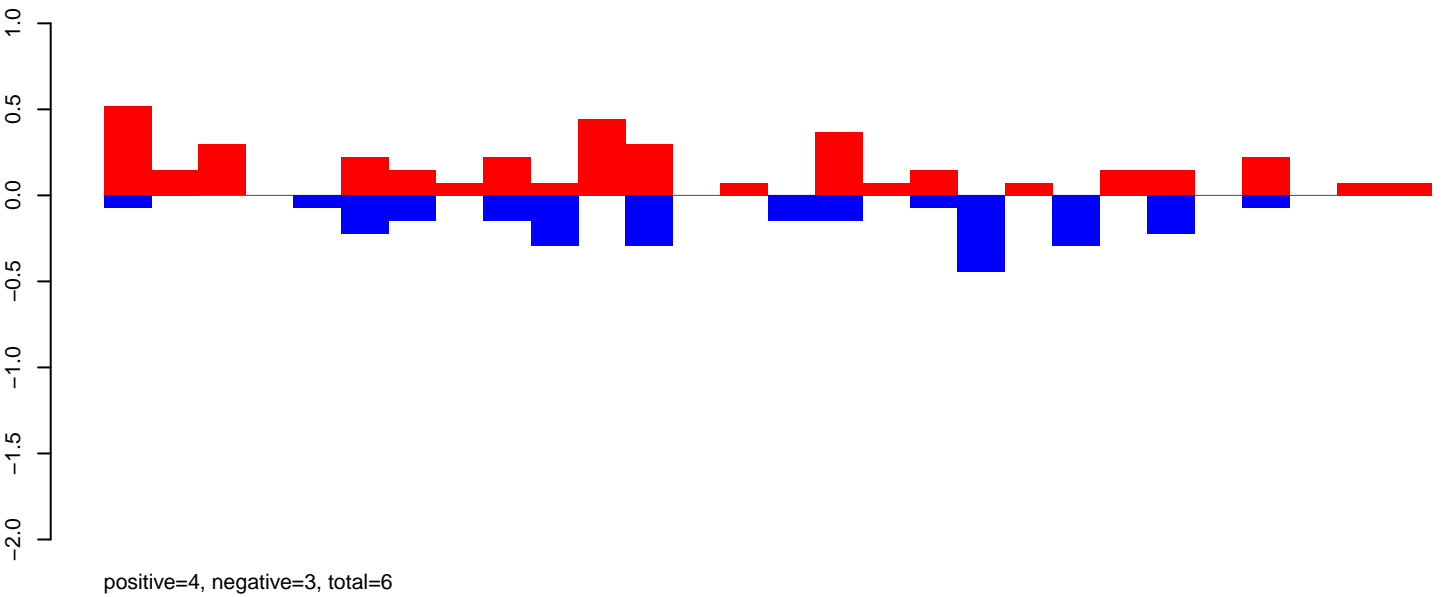
positive=8, negative=5, total=13

Window size=25, length=1479, TE@rnd-5_family-1224@Unknown@3210:1-1479

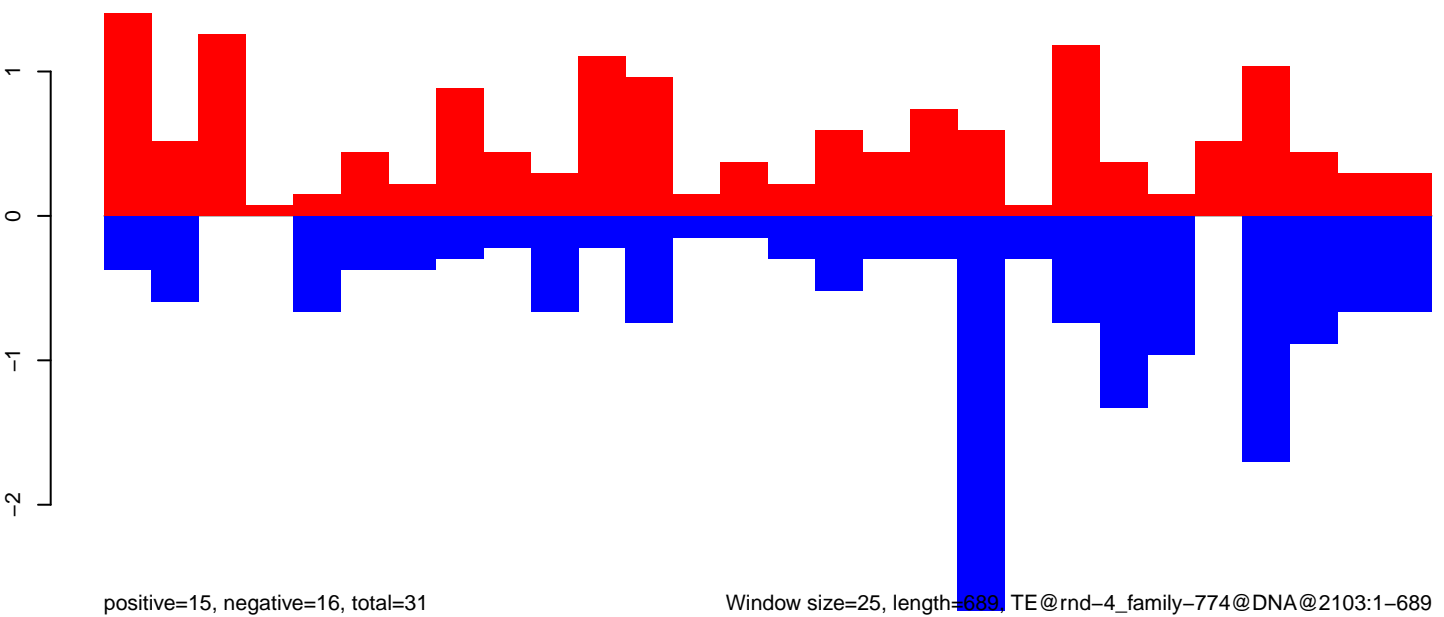
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



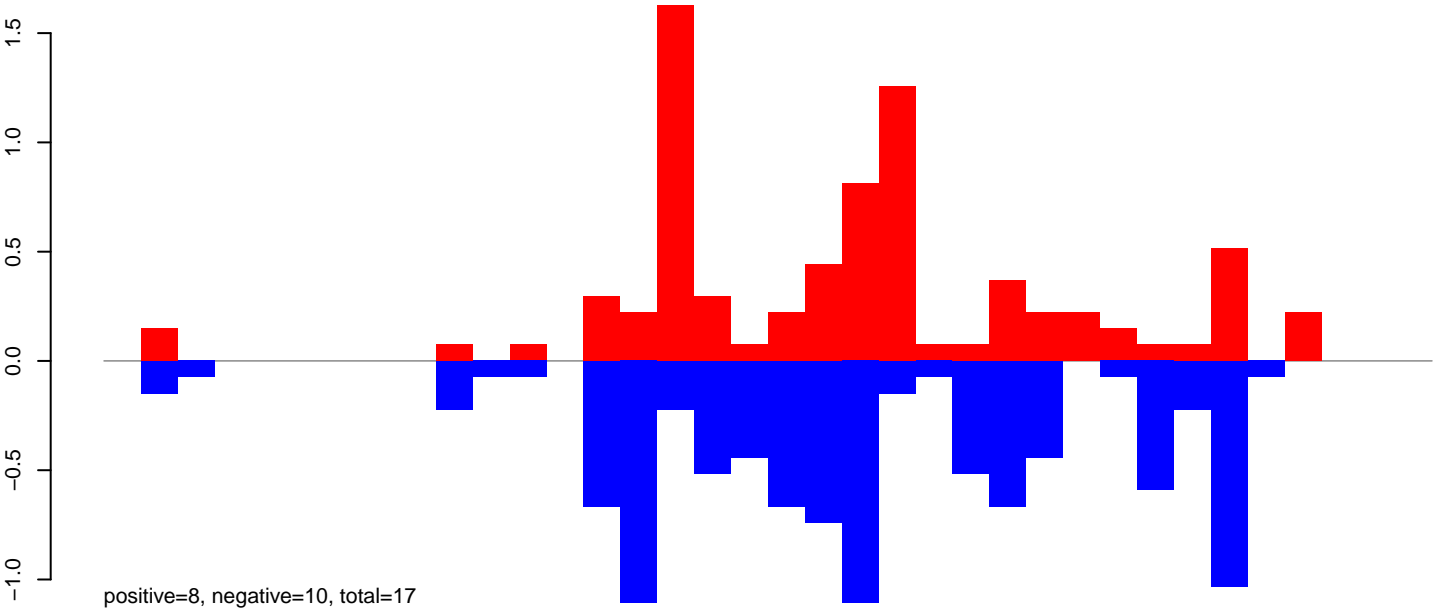
CuQuin_Testes_NL.rep



Window size=25, length=689, TE@rnd-4_family-774@DNA@2103:1-689

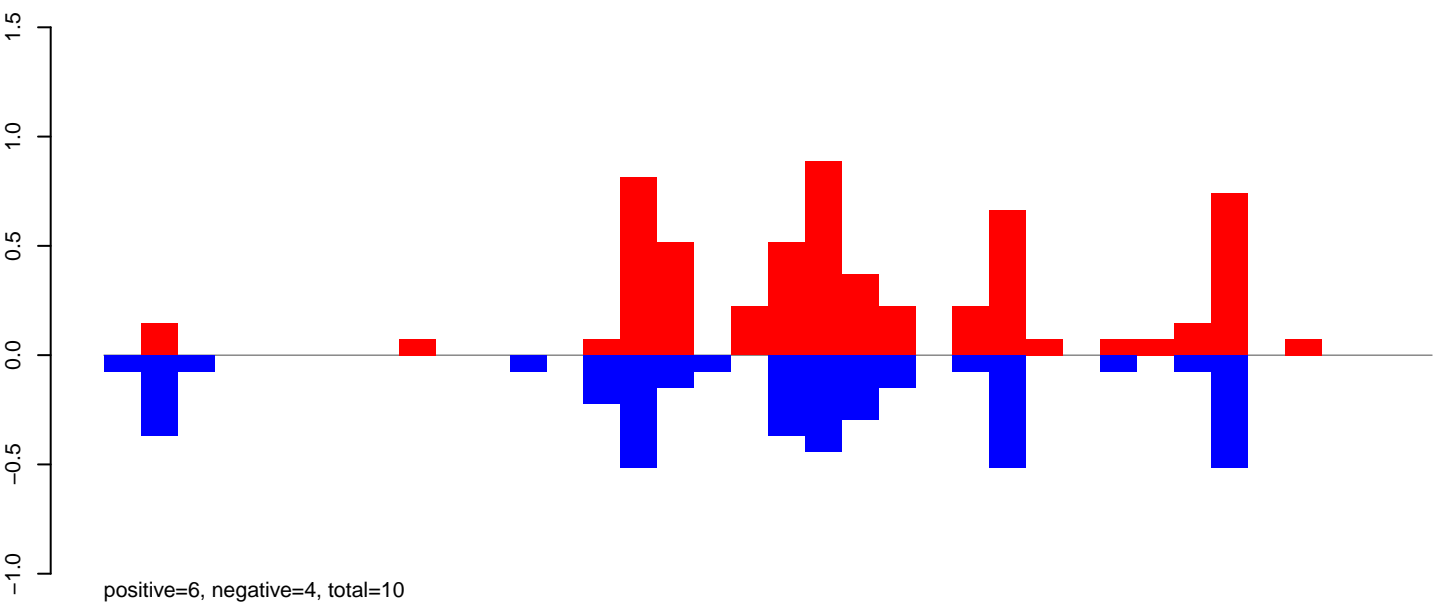
0 100 200 300 400 500 600 700

CuQuin_Testes_NL.18_23.rep



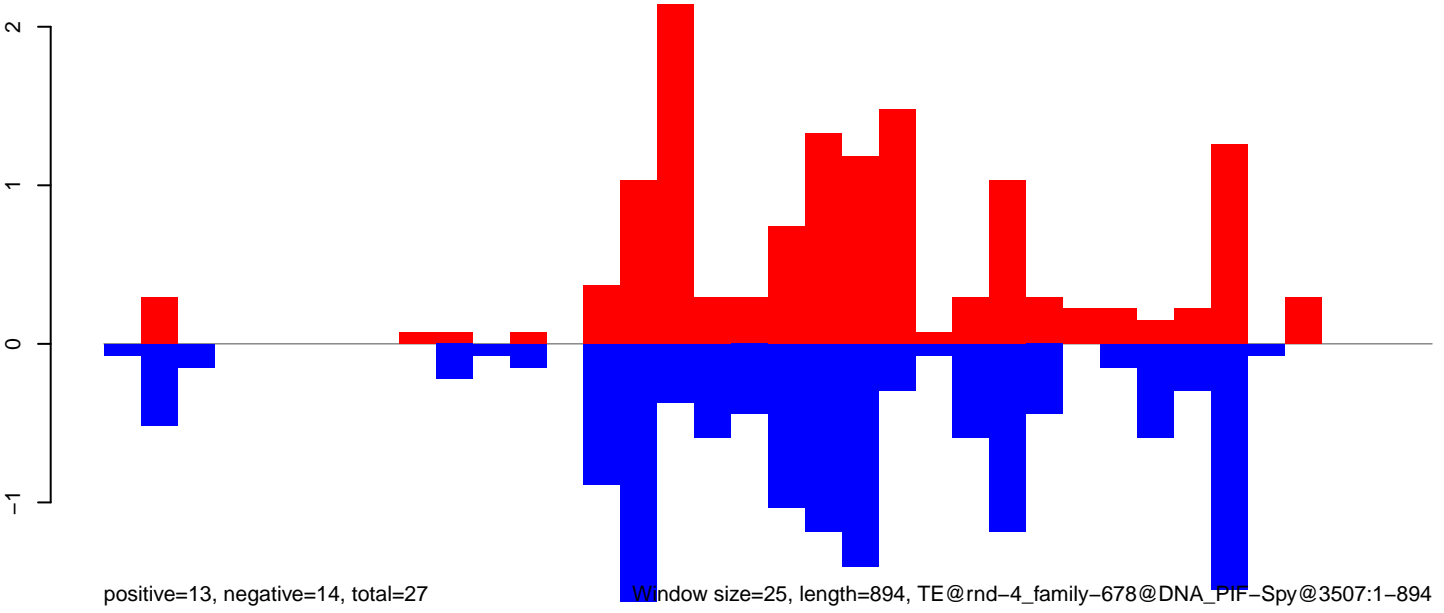
positive=8, negative=10, total=17

CuQuin_Testes_NL.24_35.rep



positive=6, negative=4, total=10

CuQuin_Testes_NL.rep

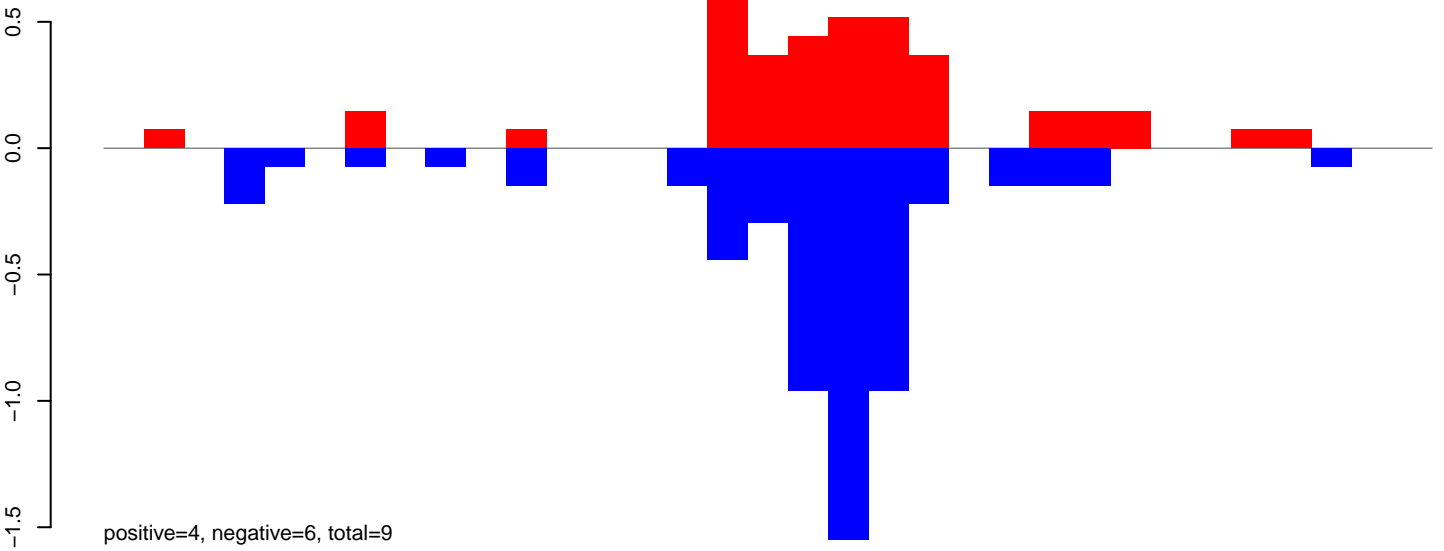


positive=13, negative=14, total=27

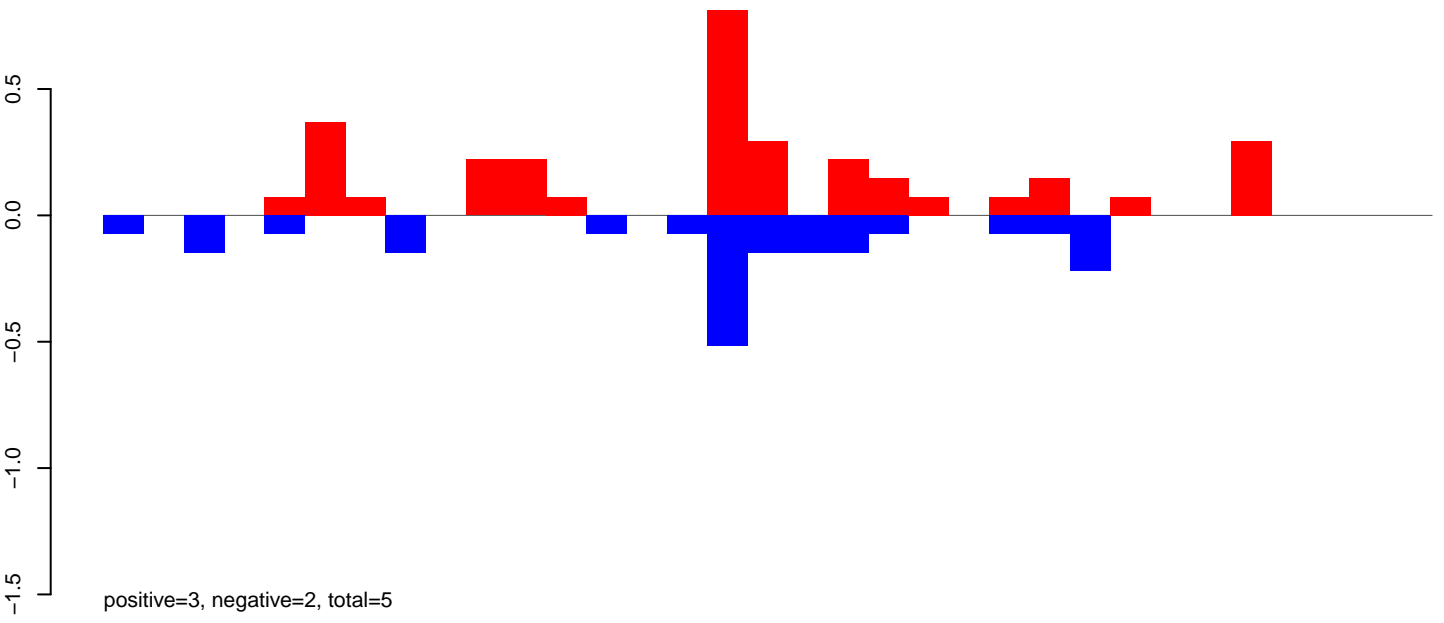
Window size=25, length=894, TE@rnd-4_family-678@DNA_PIF-Spy@3507:1-894

0 200 400 600 800

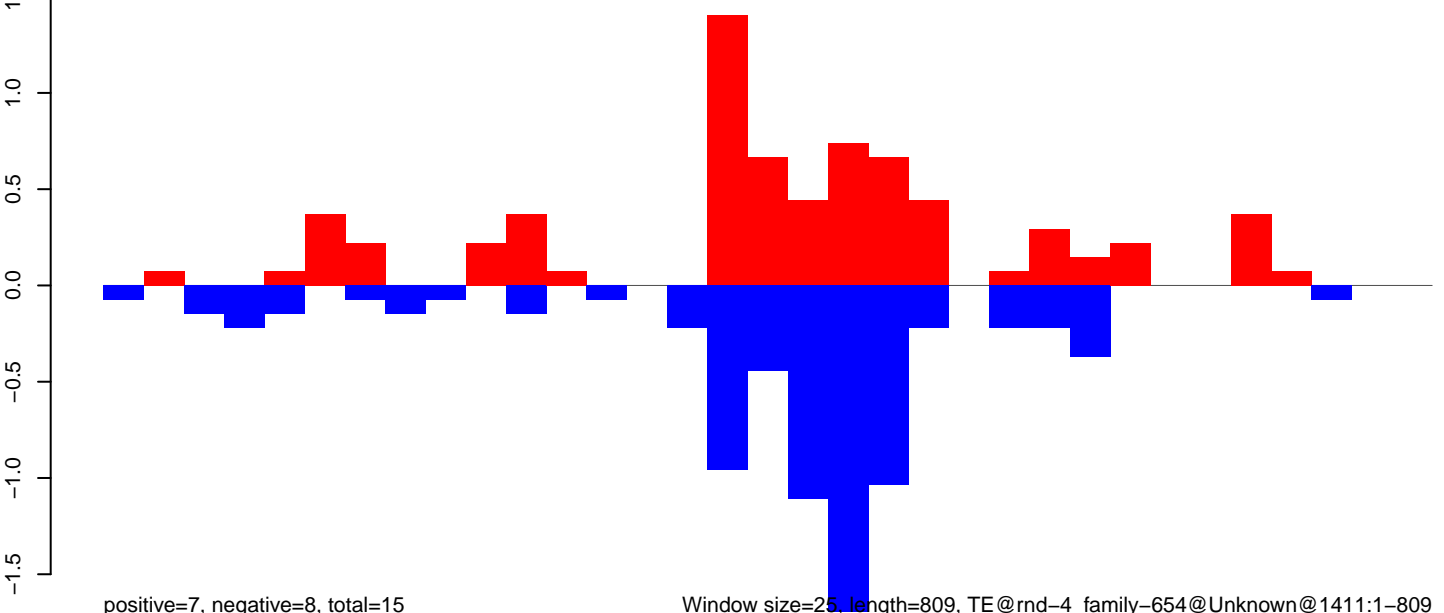
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

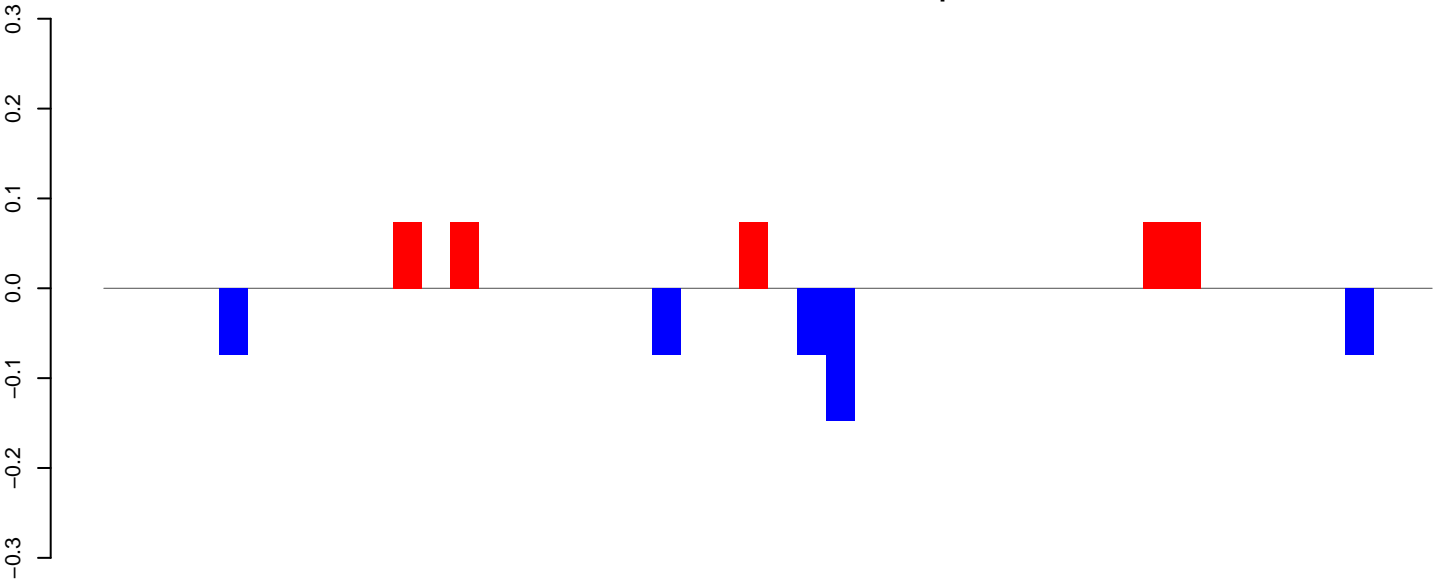


CuQuin_Testes_NL.rep



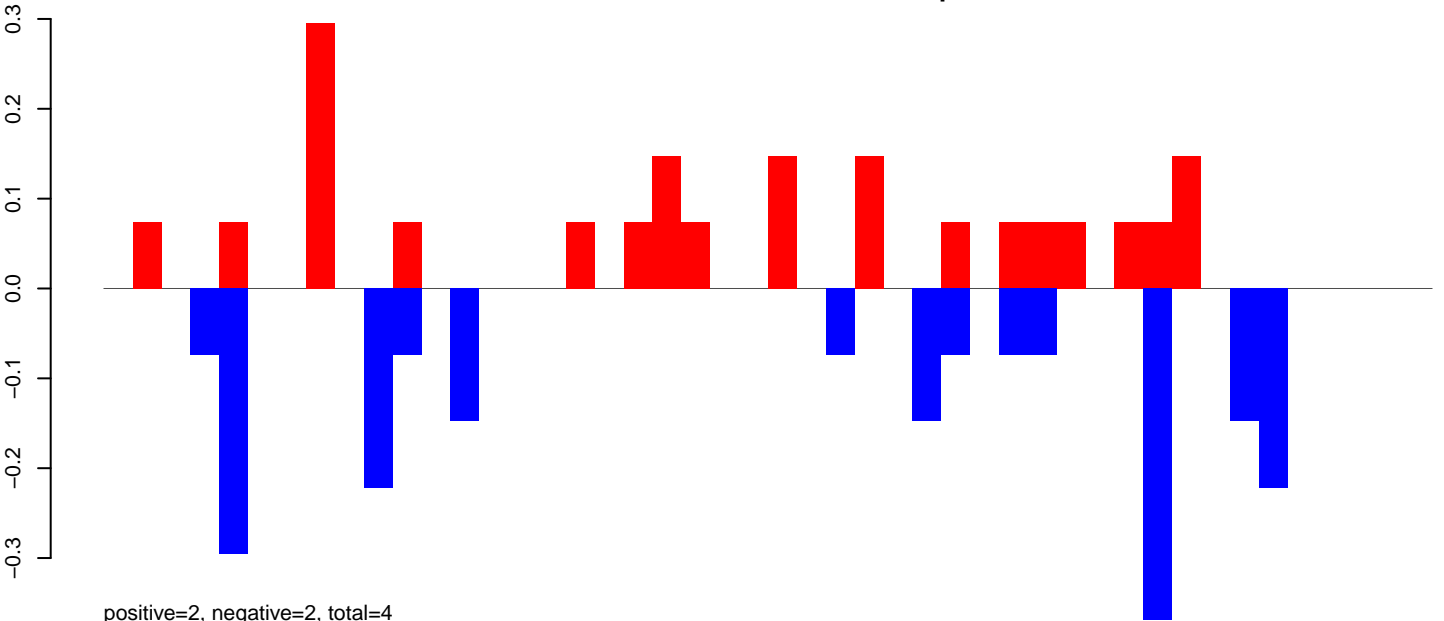
Window size=25, length=809, TE@rnd-4_family-654@Unknown@1411:1-809

CuQuin_Testes_NL.18_23.rep



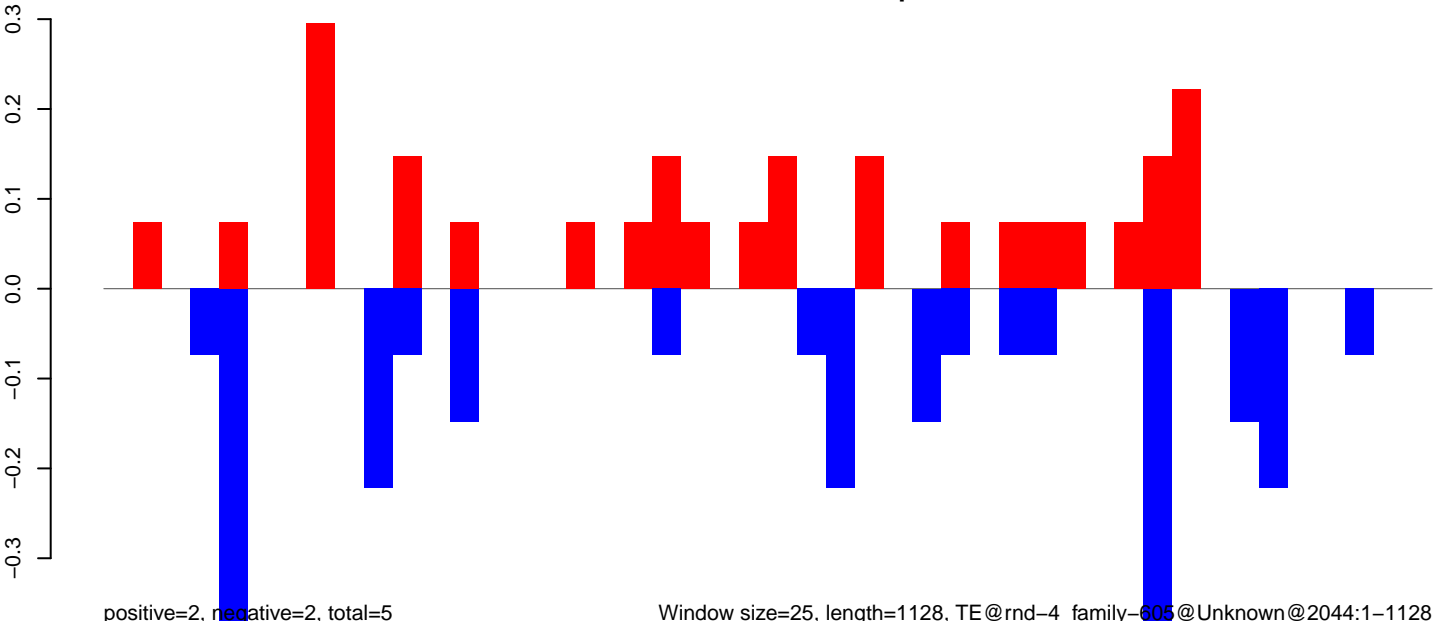
positive=0, negative=0, total=1

CuQuin_Testes_NL.24_35.rep



positive=2, negative=2, total=4

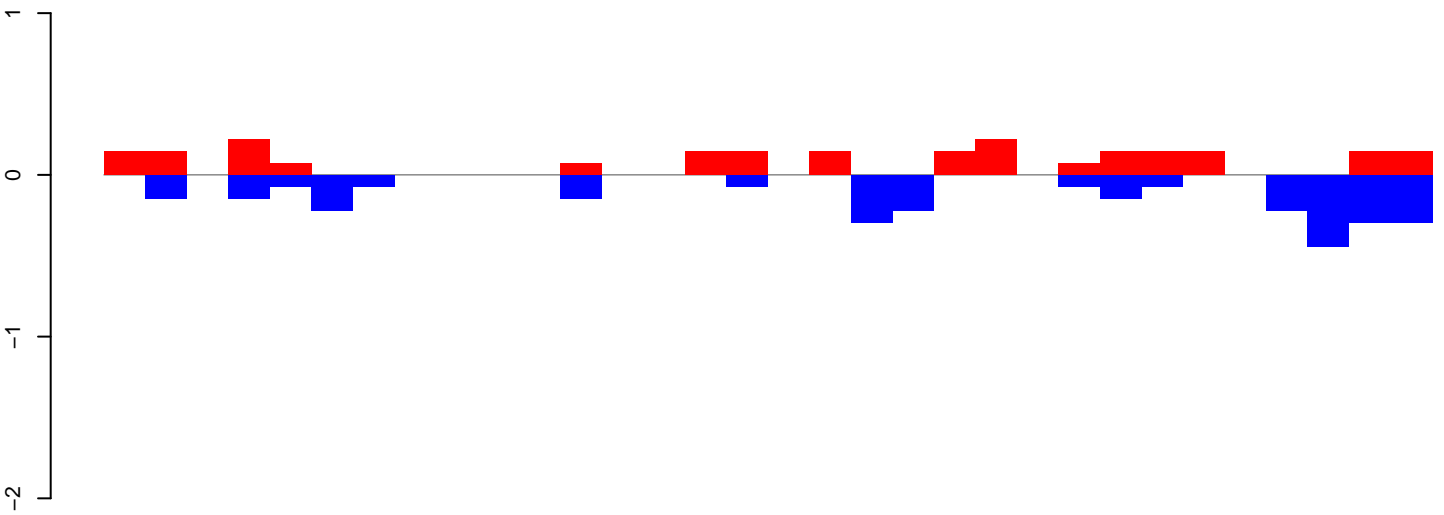
CuQuin_Testes_NL.rep



positive=2, negative=2, total=5

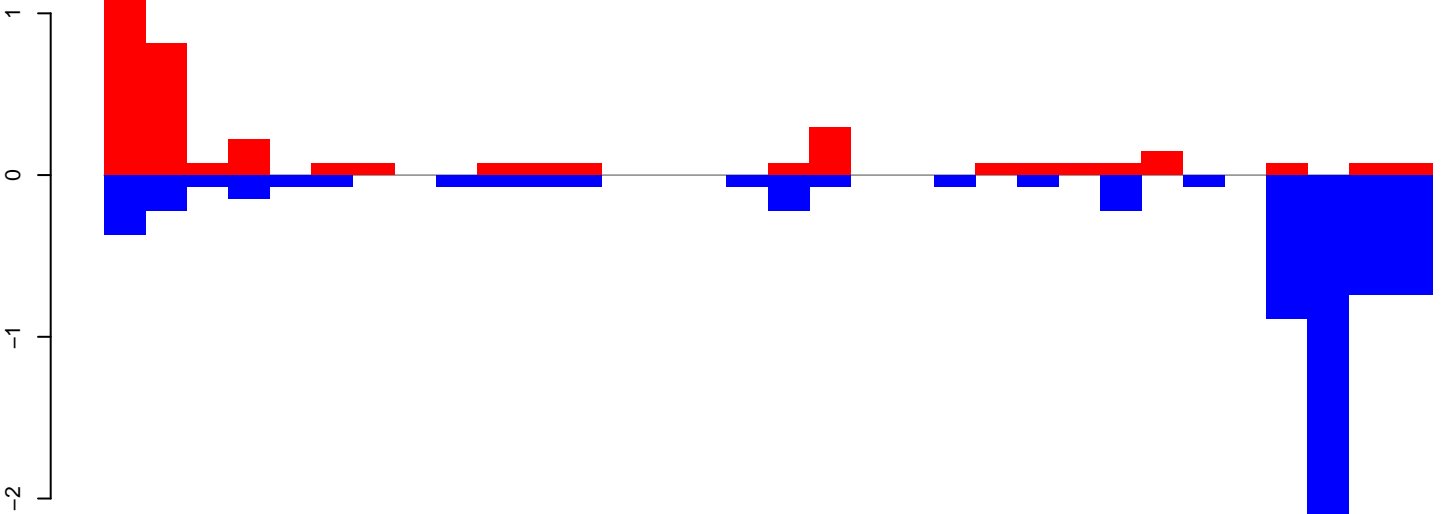
Window size=25, length=1128, TE@rnd-4_family-605@Unknown@2044:1-1128

CuQuin_Testes_NL.18_23.rep



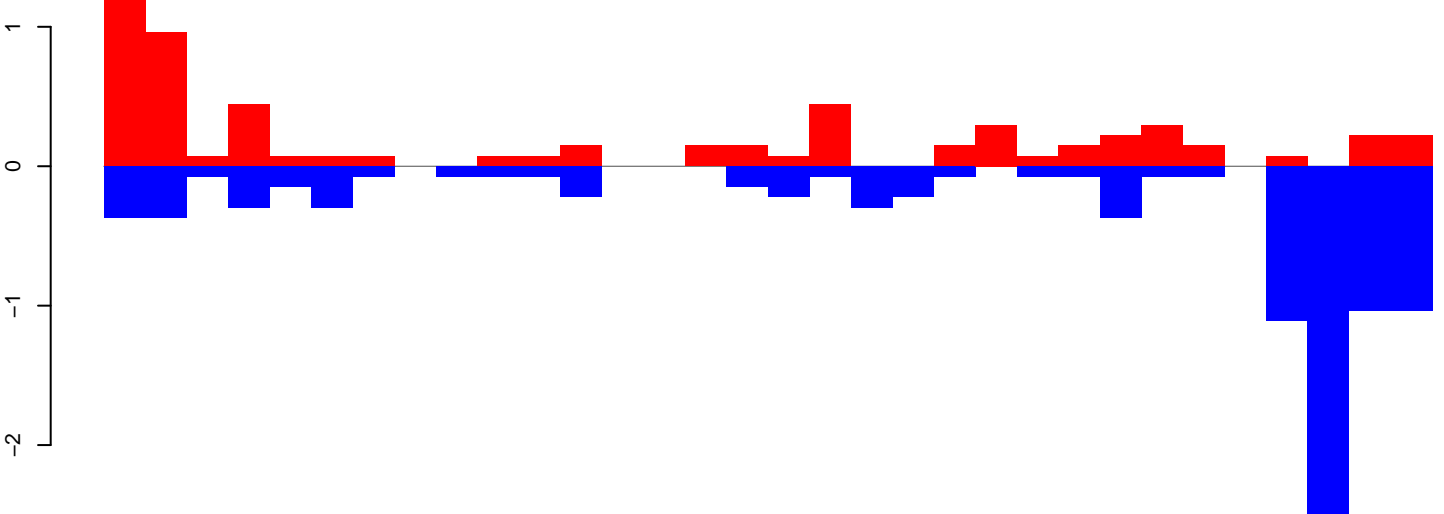
positive=2, negative=3, total=5

CuQuin_Testes_NL.24_35.rep



positive=4, negative=7, total=11

CuQuin_Testes_NL.rep

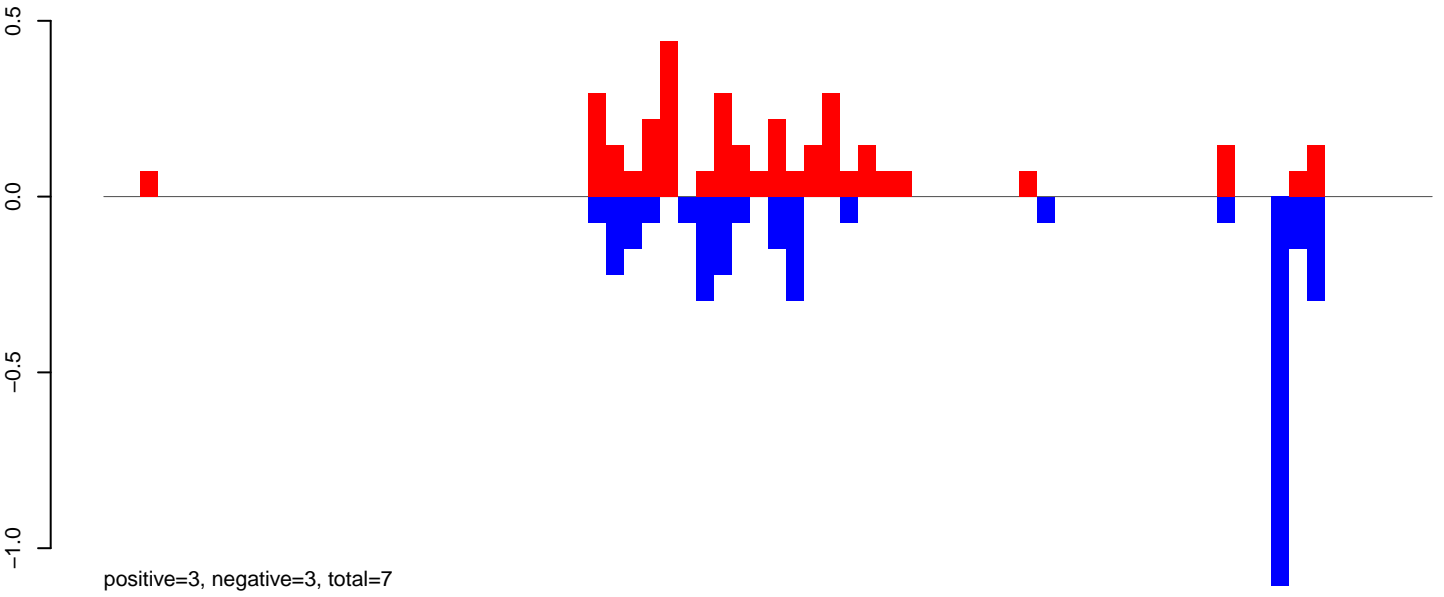


positive=6, negative=10, total=16

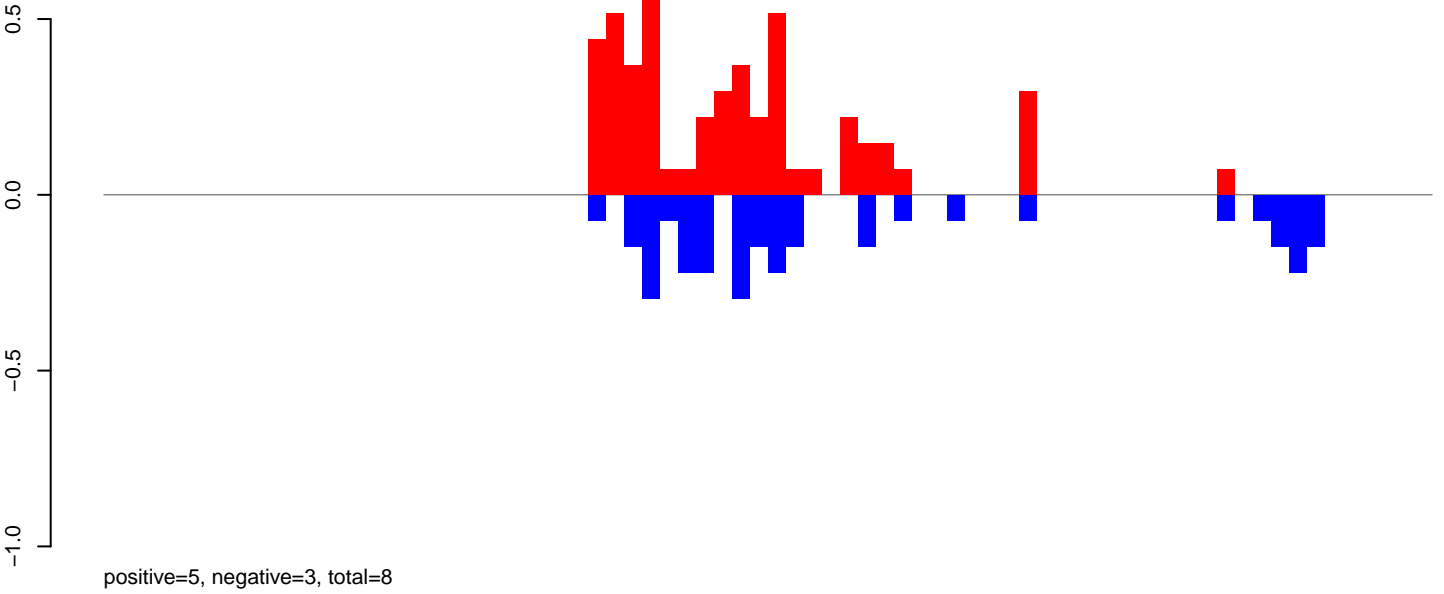
Window size=25, length=790, TE@rnd-4_family-481@Unknown@785:1-790

0 200 400 600 800

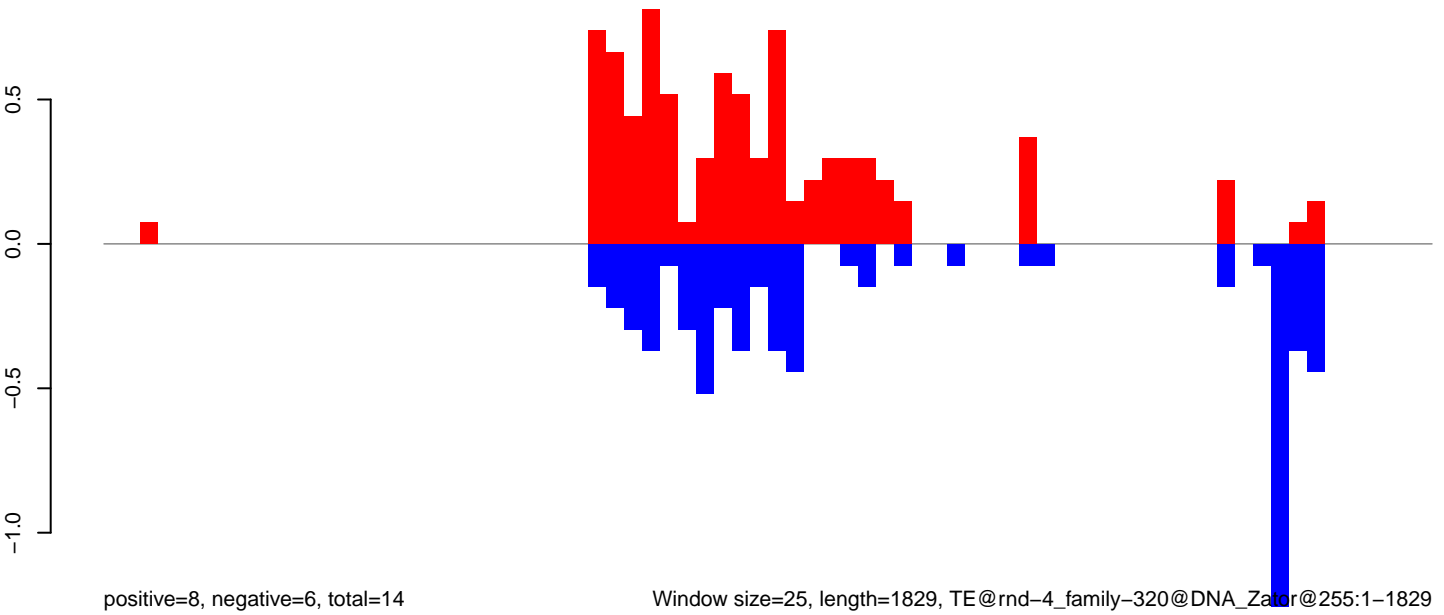
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



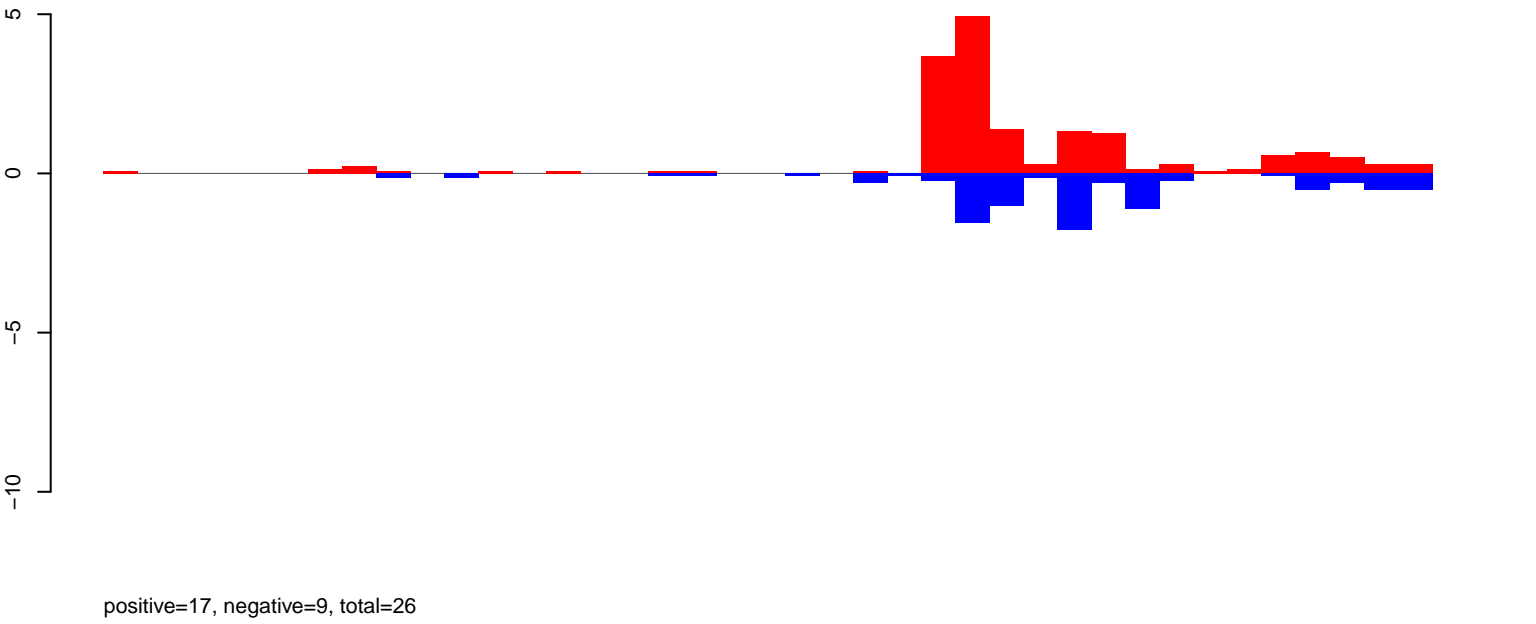
CuQuin_Testes_NL.rep



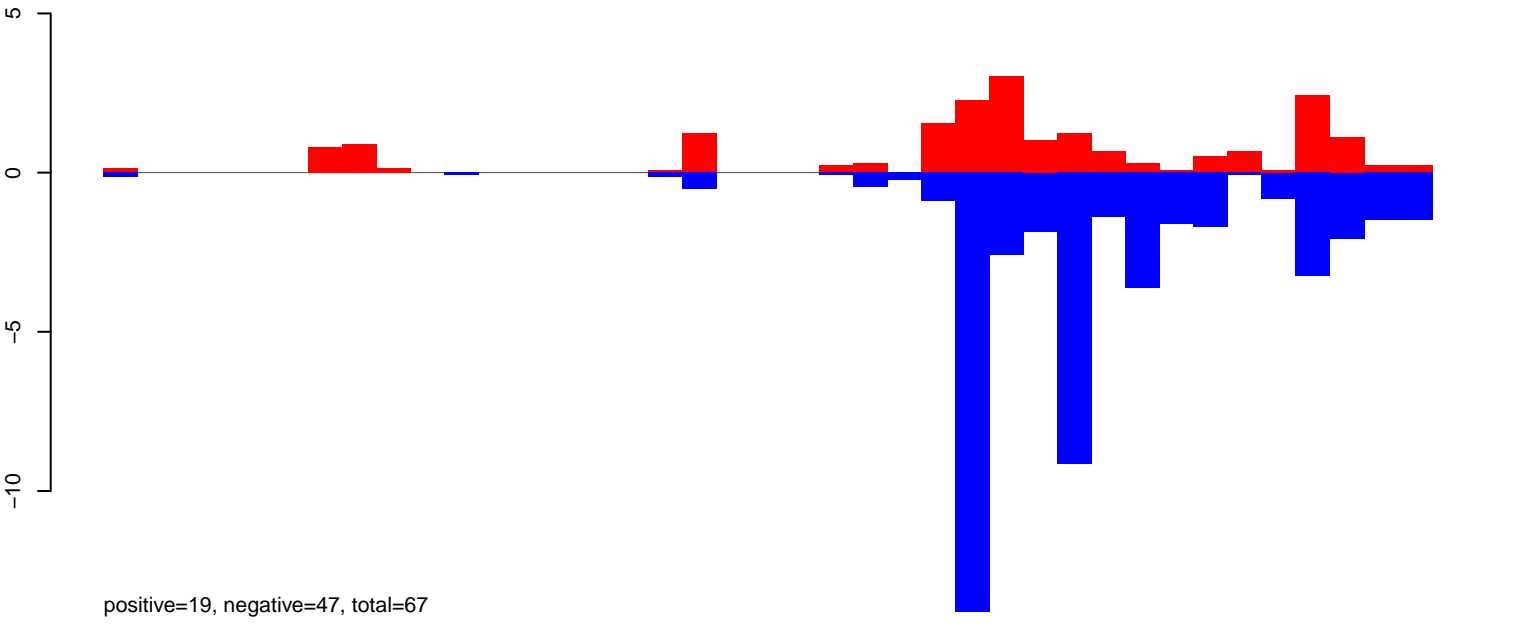
Window size=25, length=1829, TE@rnd-4_family-320@DNA_Zair@255:1-1829

0 500 1000 1500

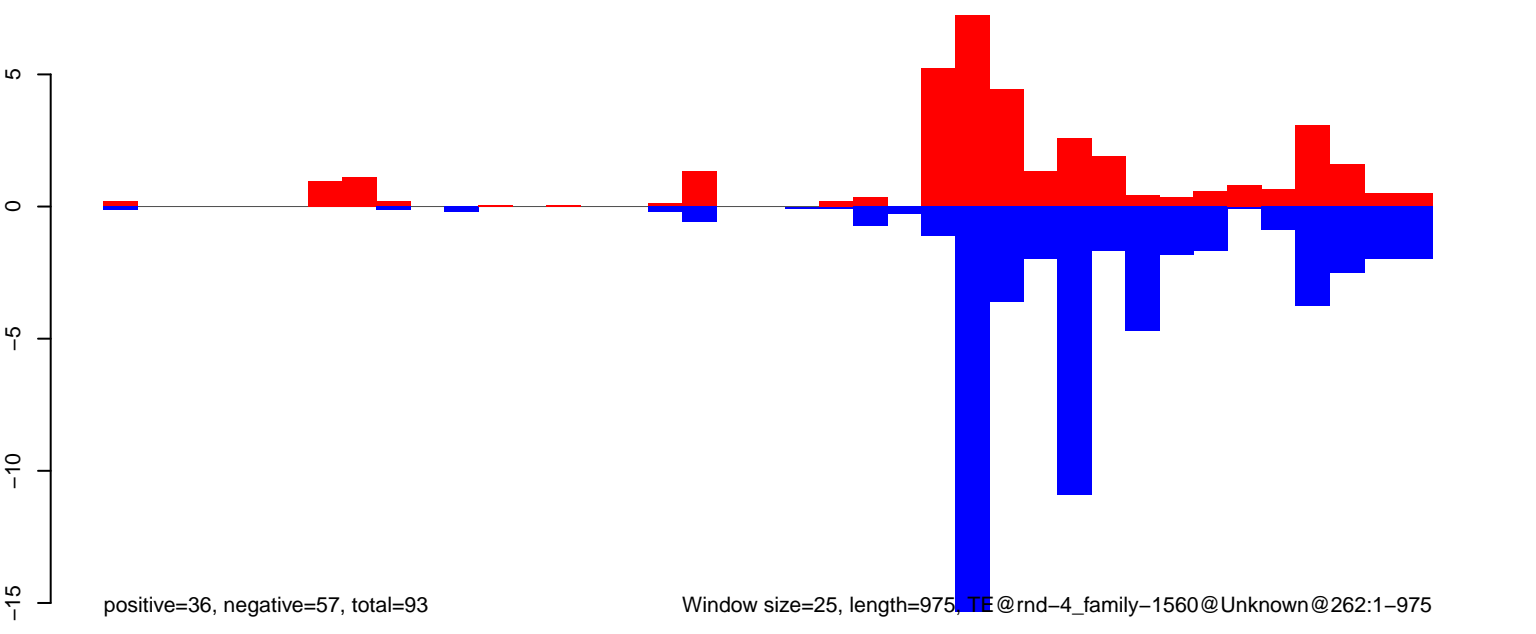
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

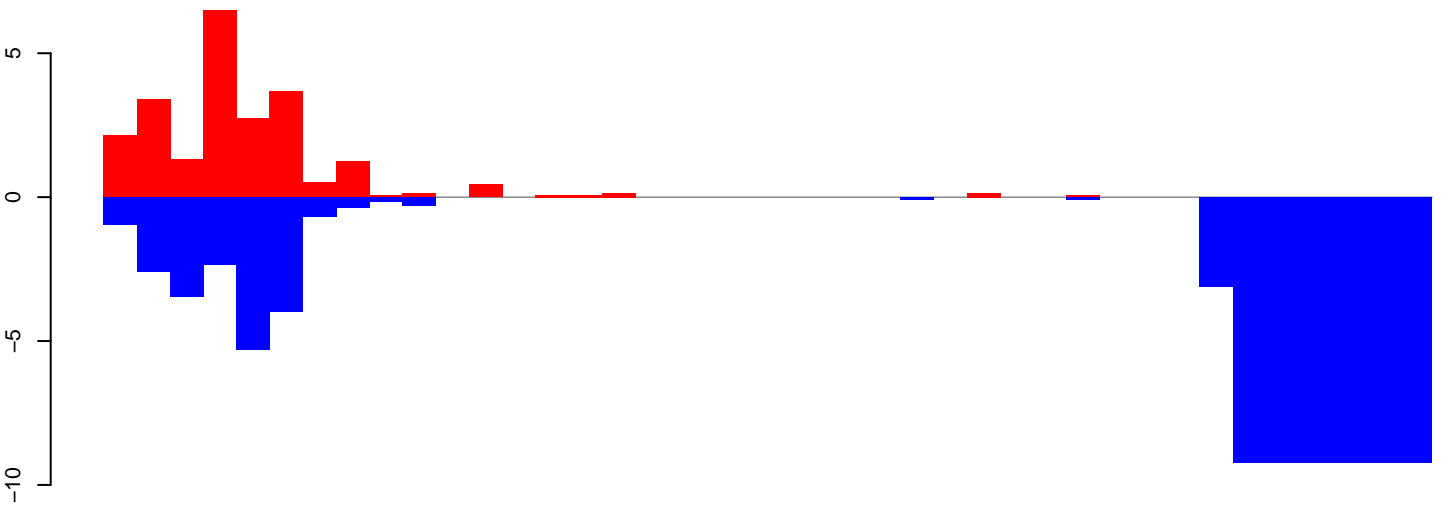


CuQuin_Testes_NL.rep



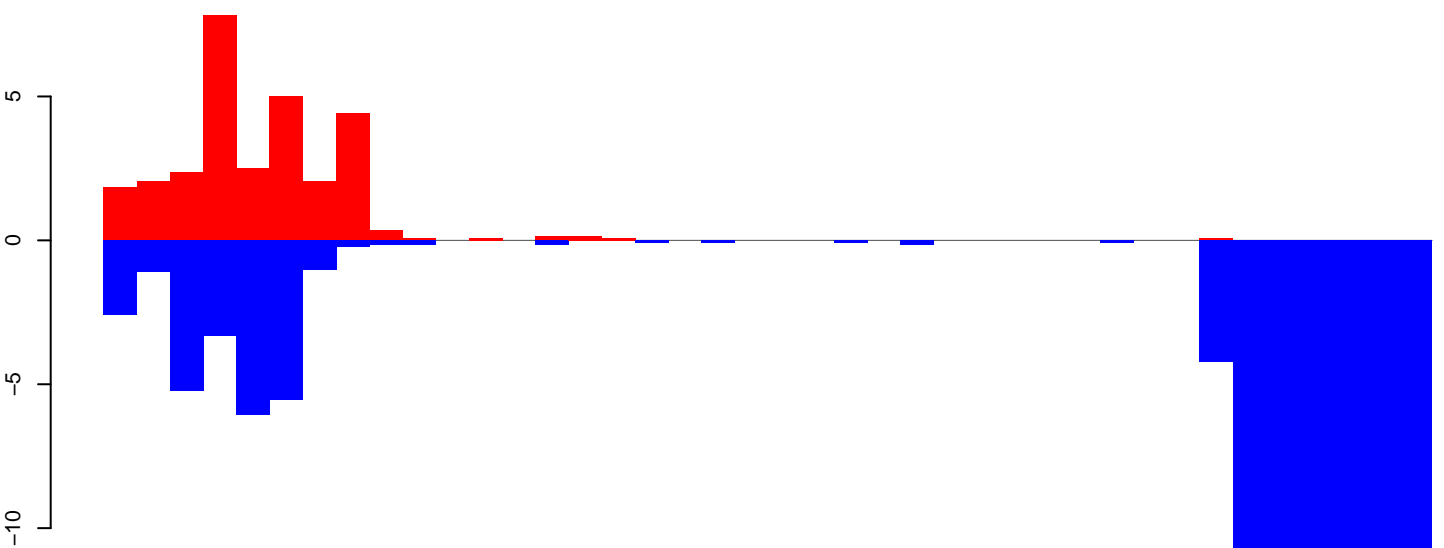
Window size=25, length=975, TE@rnd_4_family-1560@Unknown@262:1-975

CuQuin_Testes_NL.18_23.rep



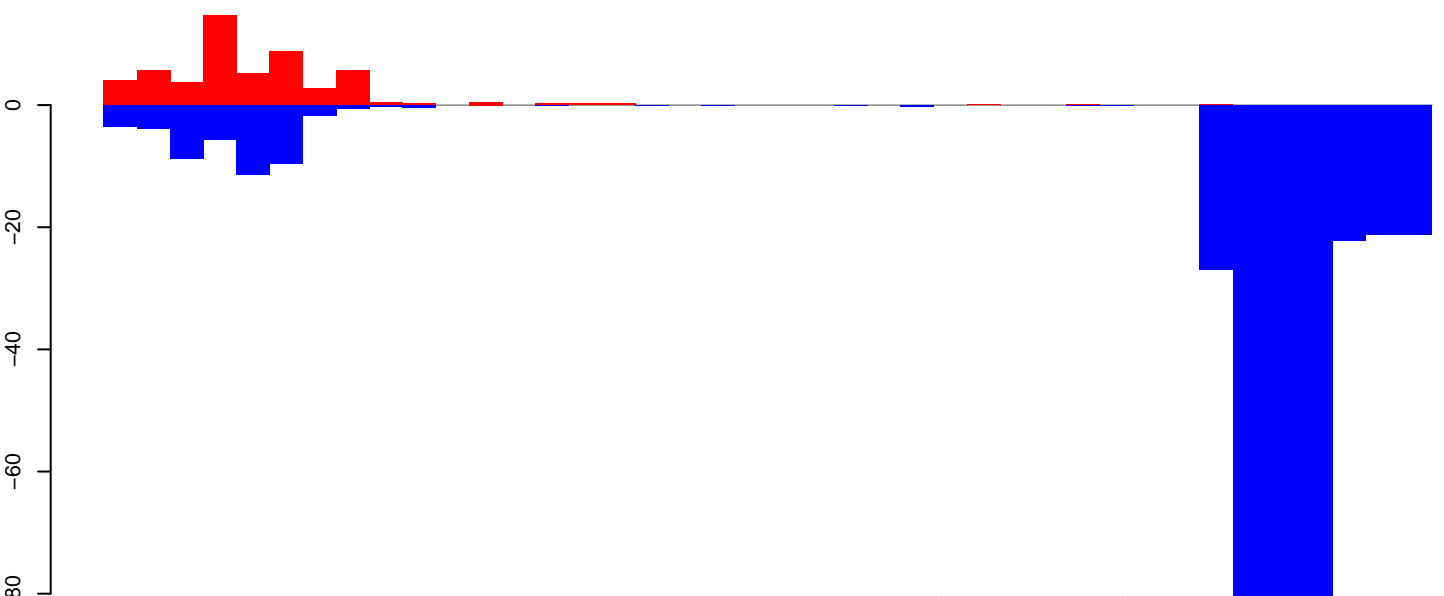
positive=23, negative=79, total=102

CuQuin_Testes_NL.24_35.rep



positive=29, negative=106, total=135

CuQuin_Testes_NL.rep

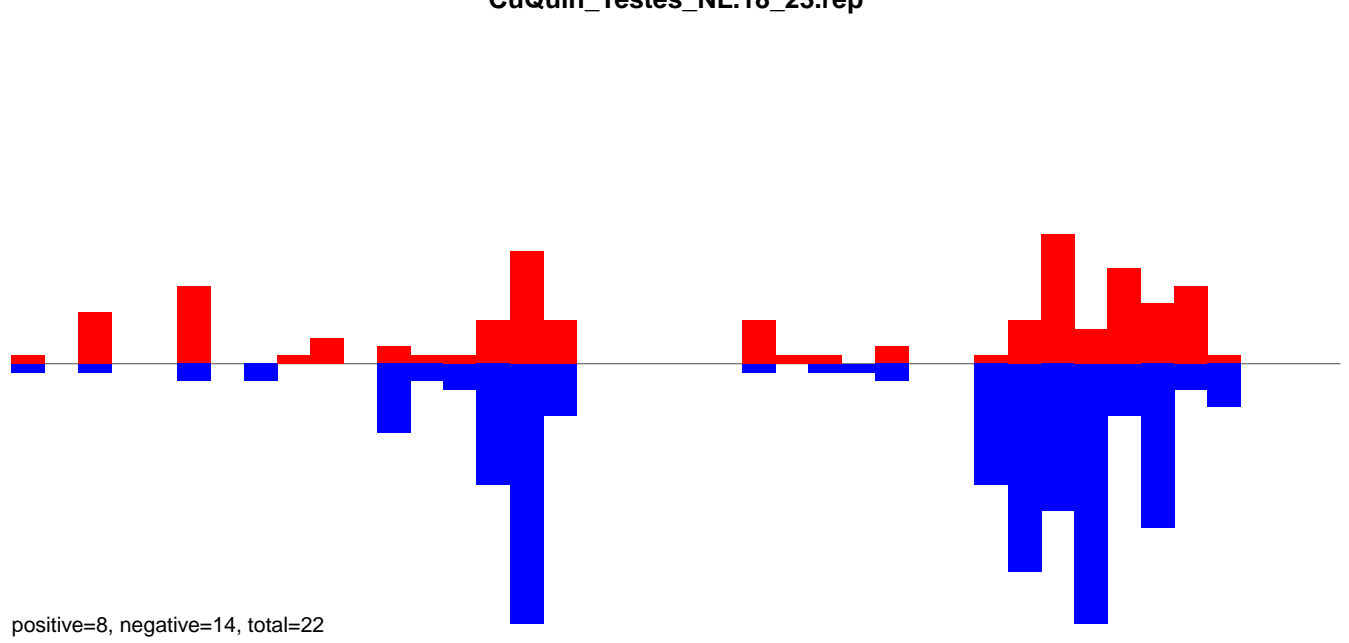


positive=52, negative=385, total=437

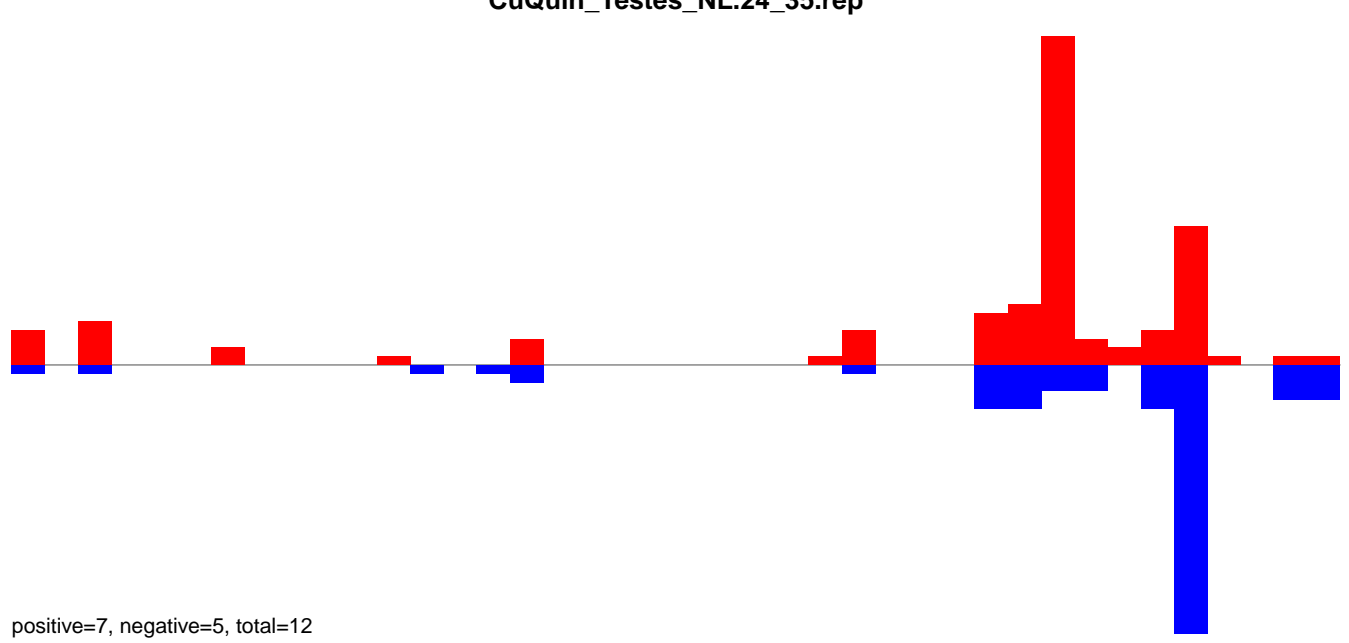
Window size=25, length=1000, TE@rnd-3_family-36@SINE_tRNA-Deu@122:1-1000

0 200 400 600 800 1000

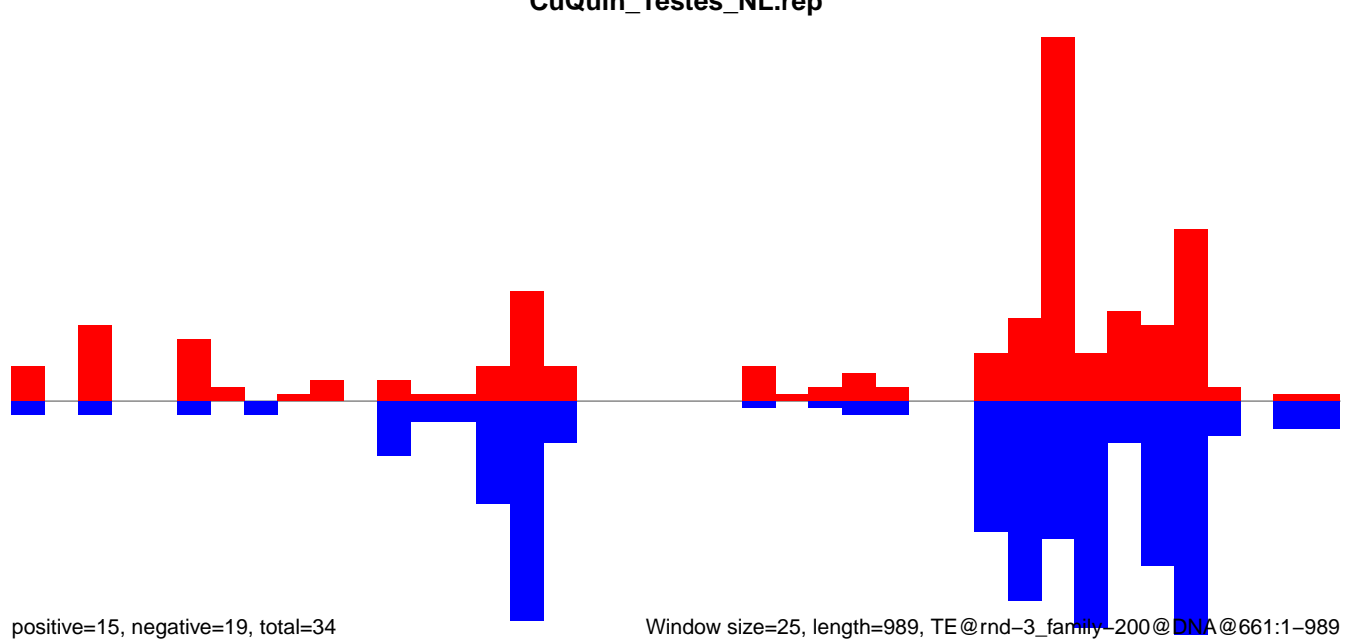
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

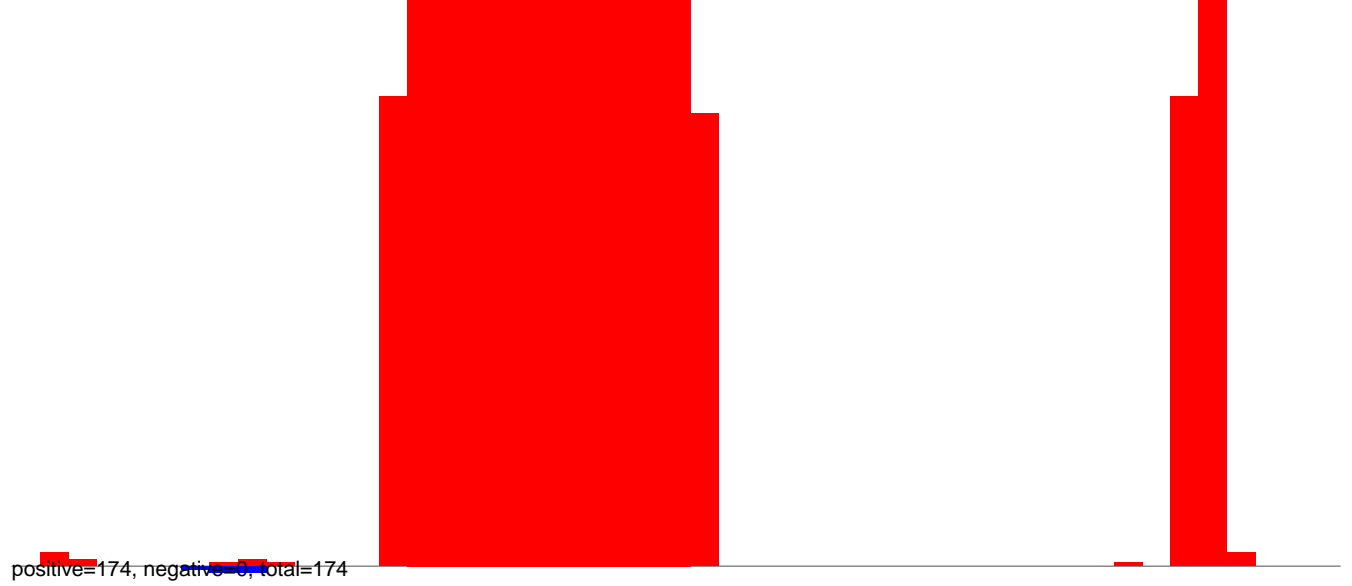


Window size=25, length=989, TE@rnd-3_family-200@DNA@661:1-989

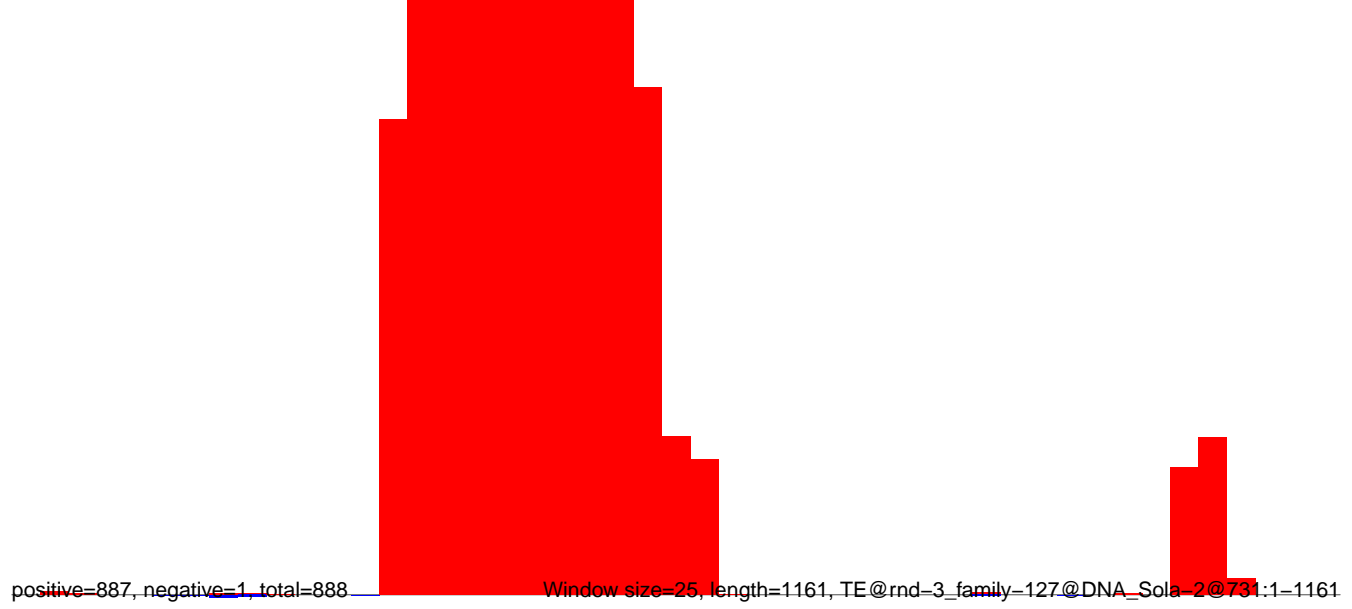
CuQuin_Testes_NL.18_23.rep



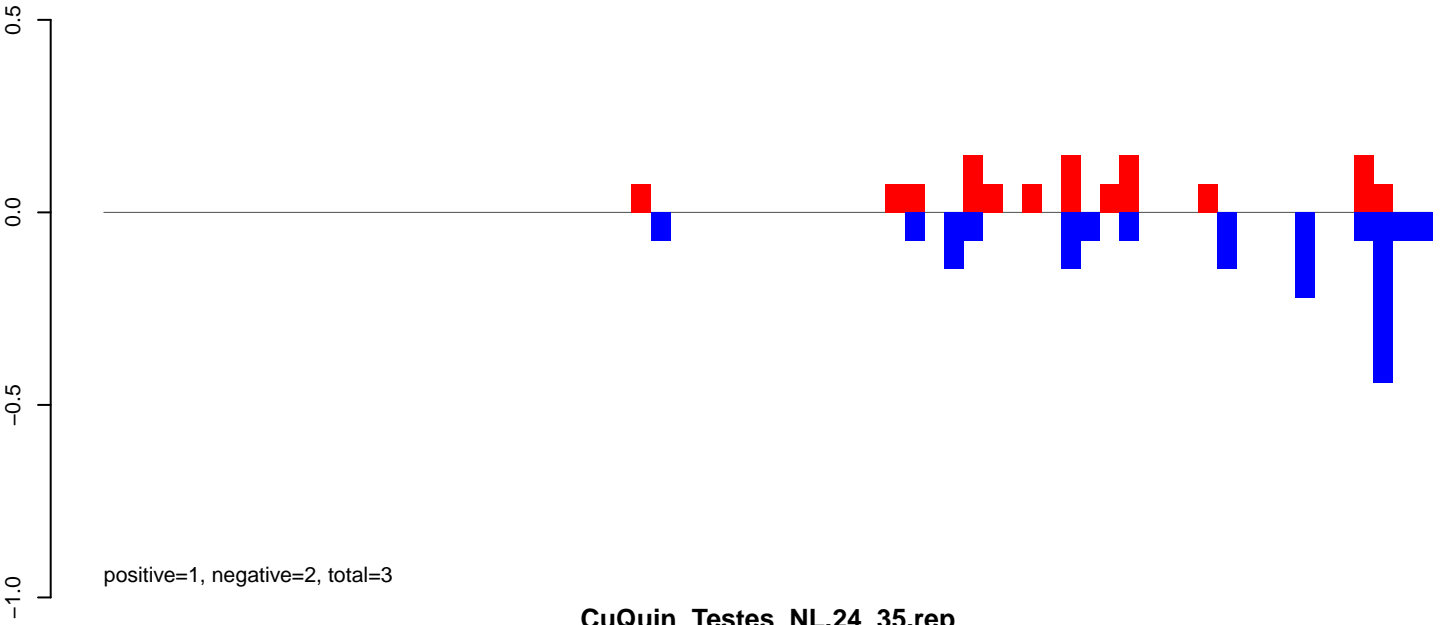
CuQuin_Testes_NL.24_35.rep



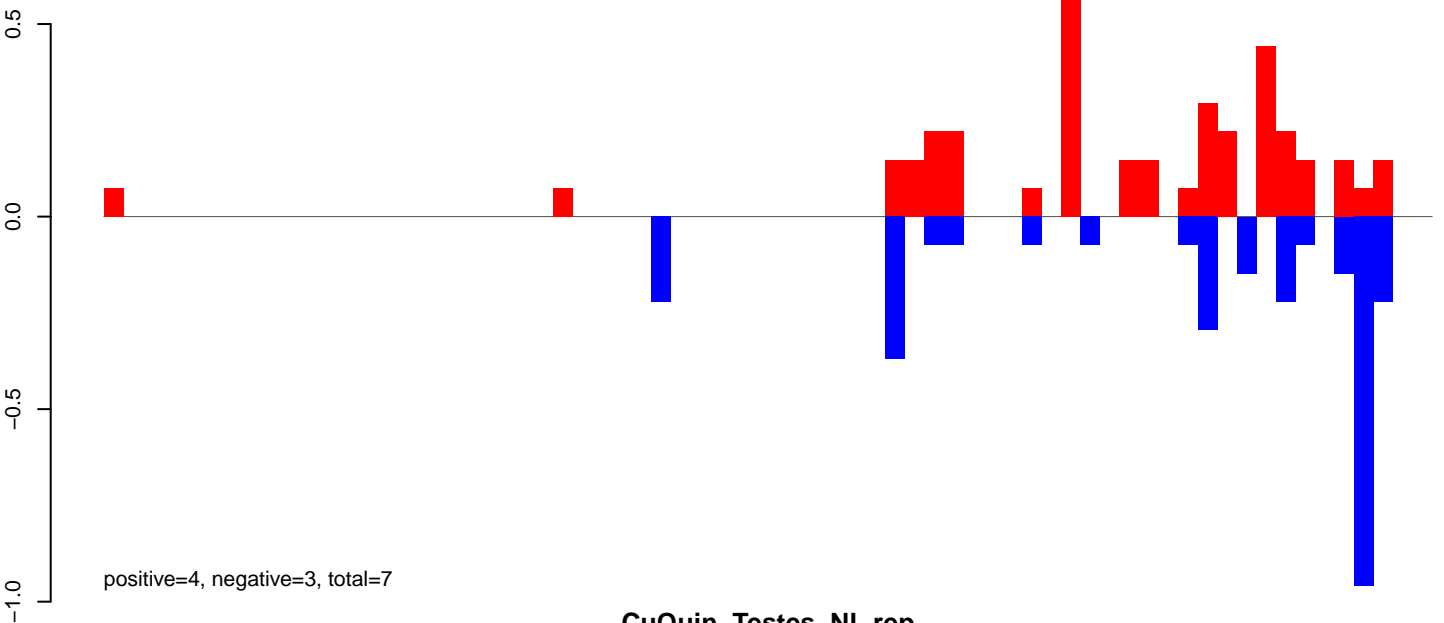
CuQuin_Testes_NL.rep



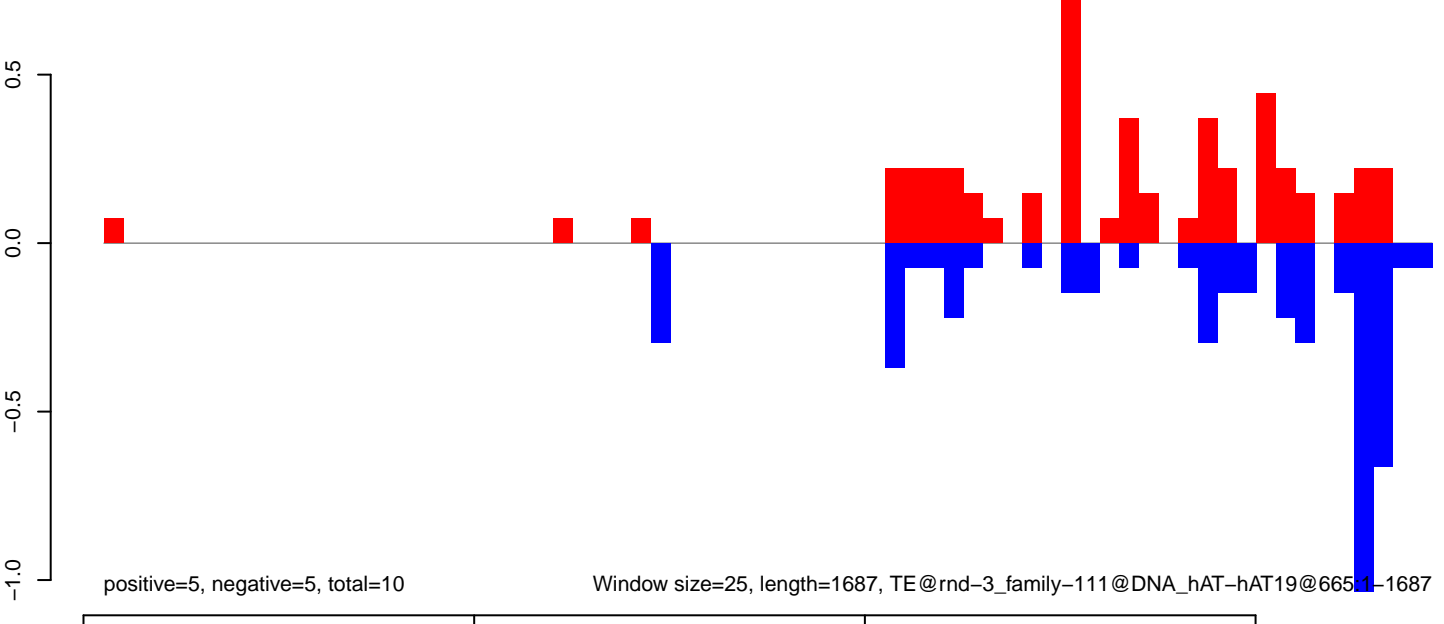
CuQuin_Testes_NL.18_23.rep



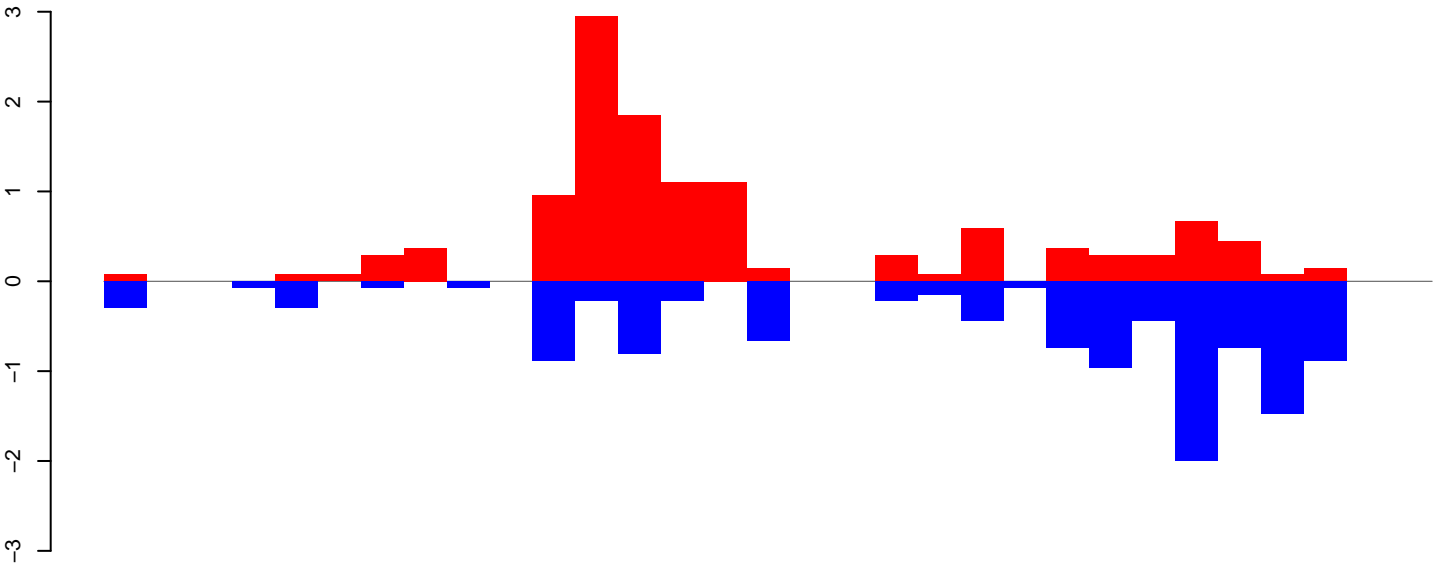
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

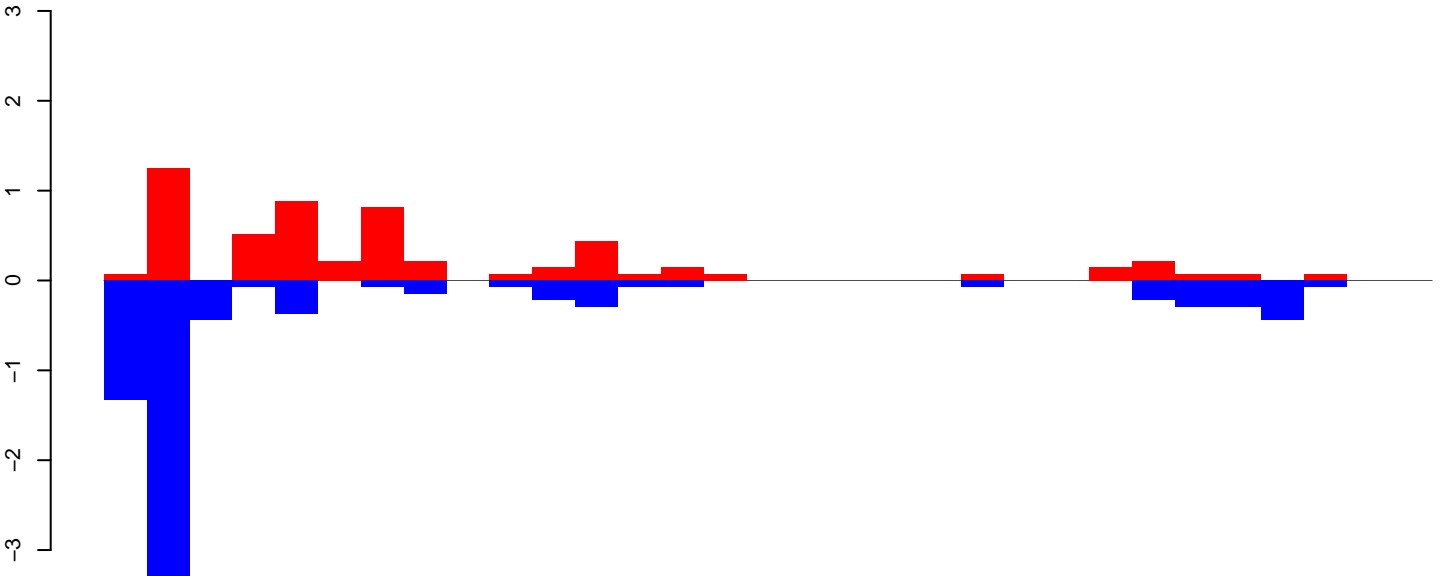


CuQuin_Testes_NL.18_23.rep



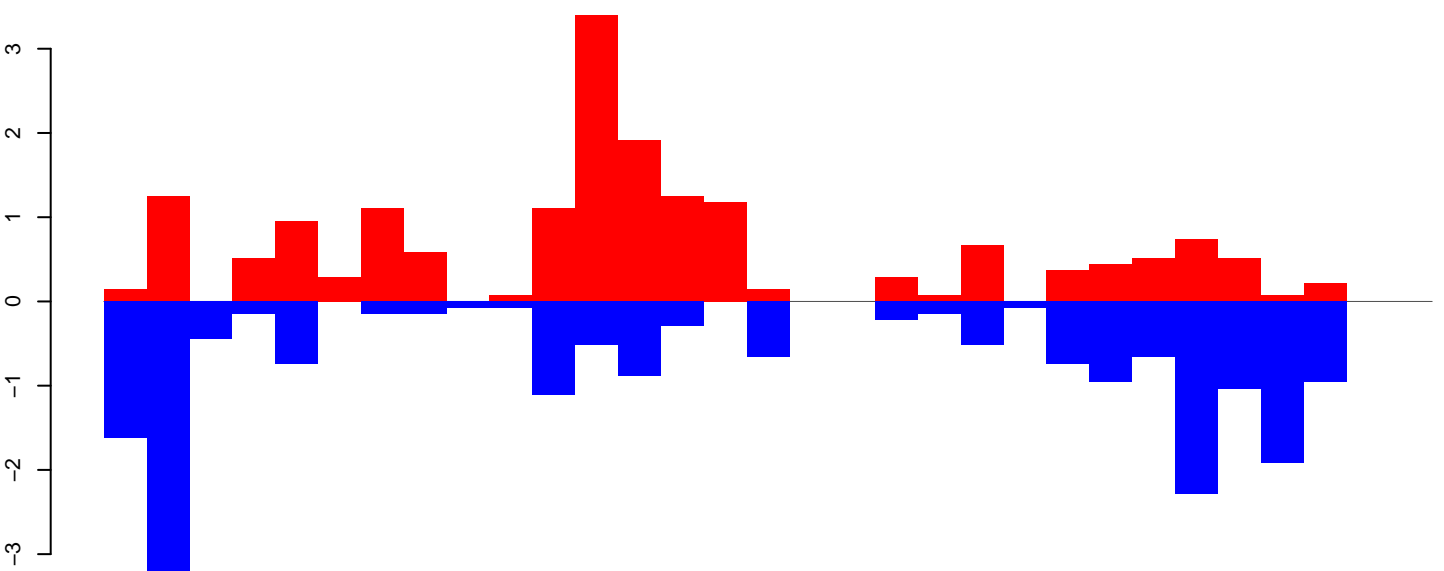
positive=12, negative=12, total=24

CuQuin_Testes_NL.24_35.rep



positive=6, negative=8, total=14

CuQuin_Testes_NL.rep

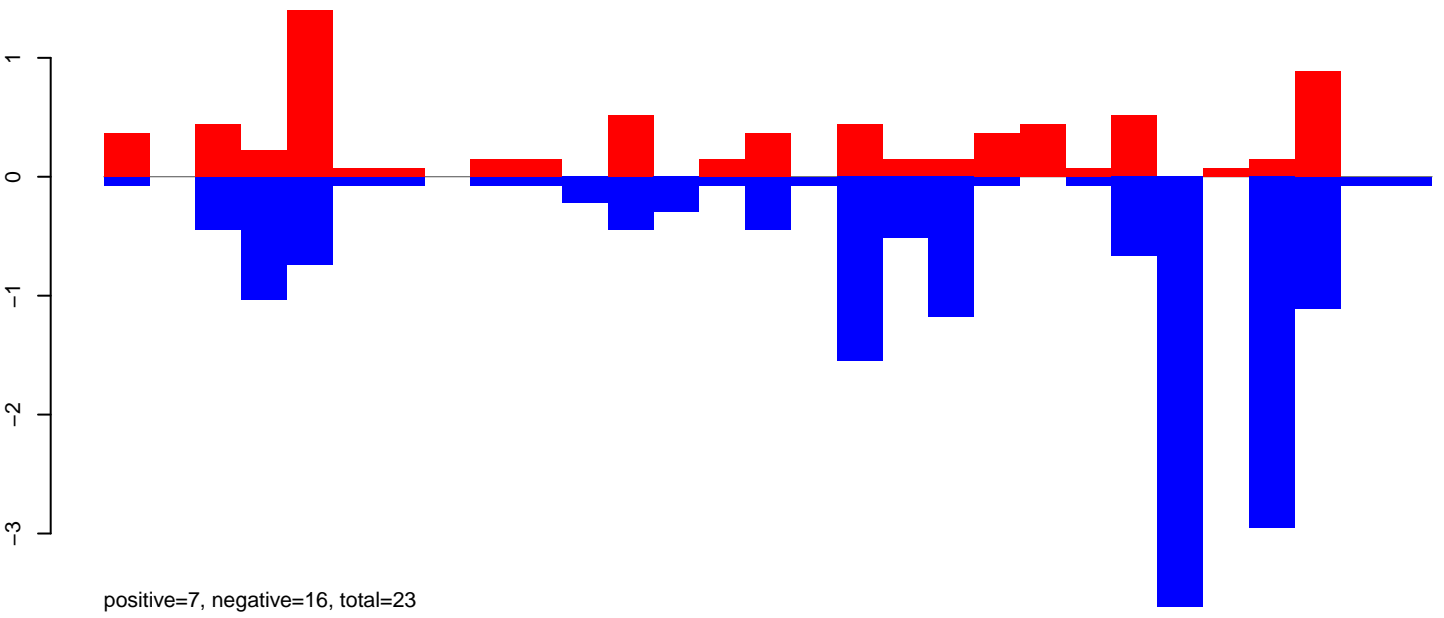


positive=18, negative=20, total=38

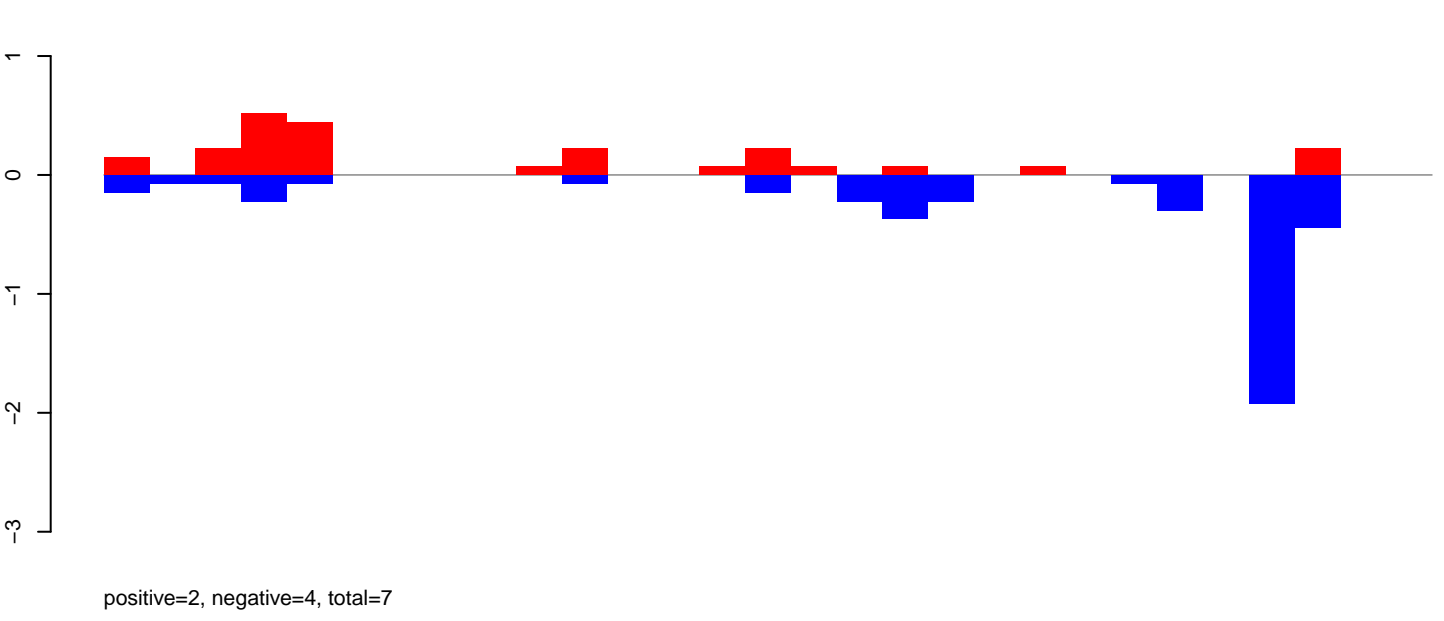
Window size=25, length=752, TE@rnd-2_family-99@Unknown@2572:1-752

0 200 400 600 800

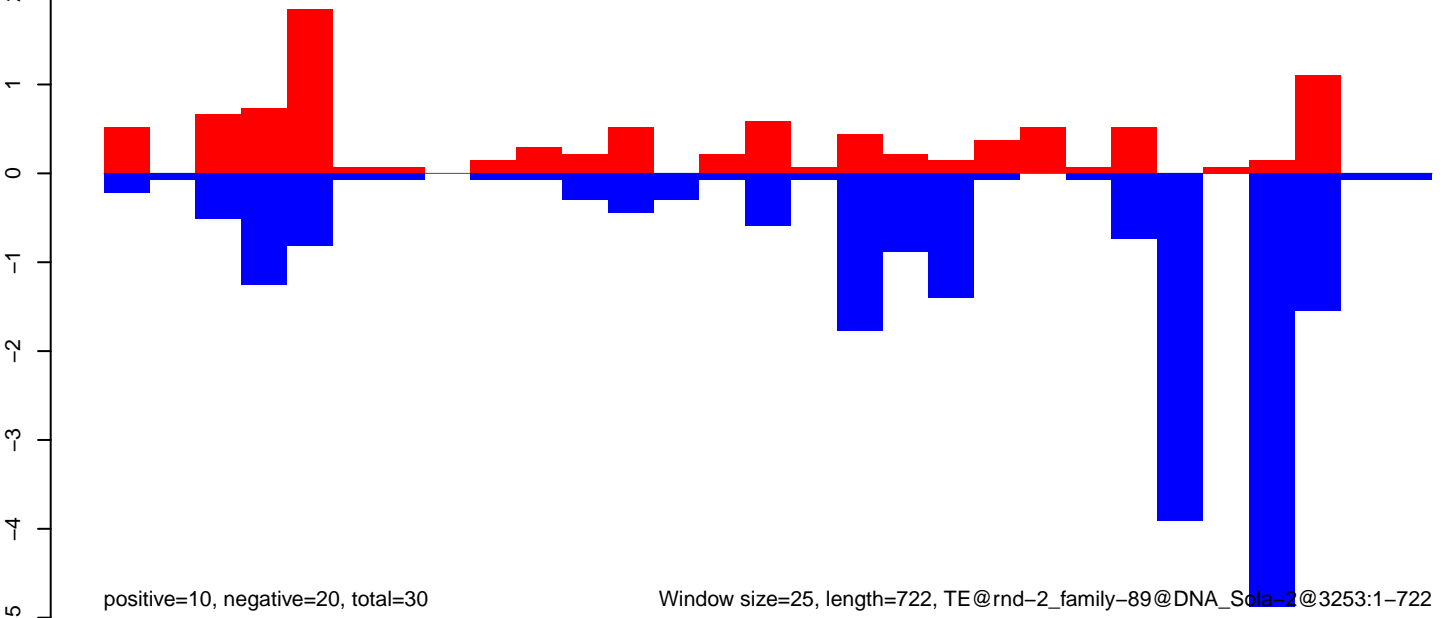
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

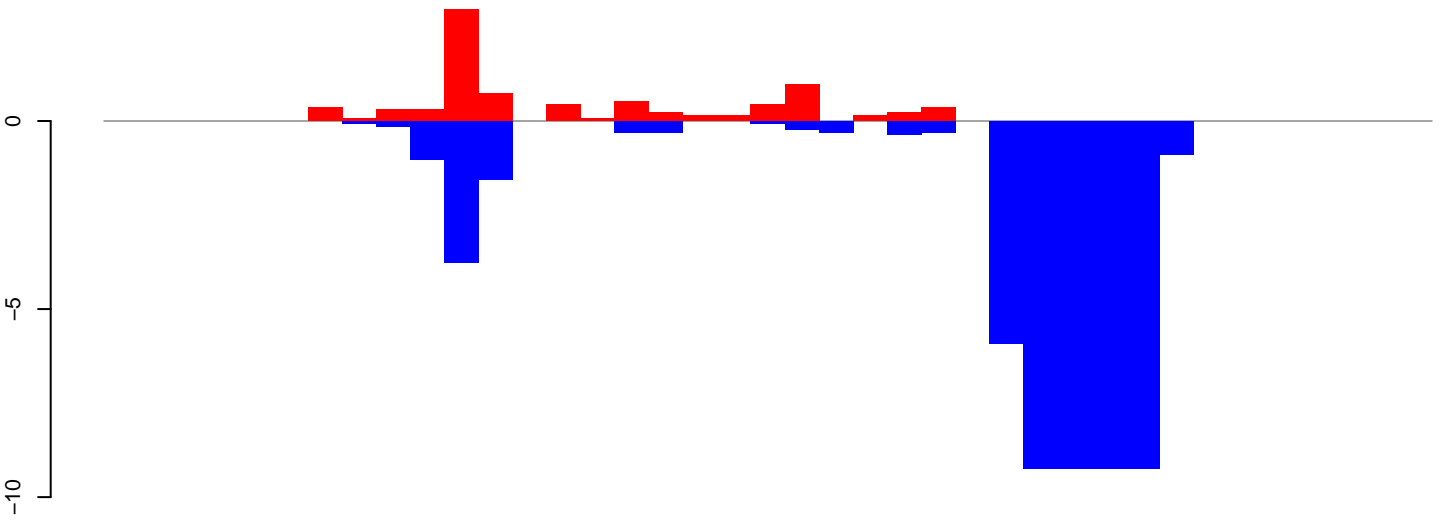


CuQuin_Testes_NL.rep



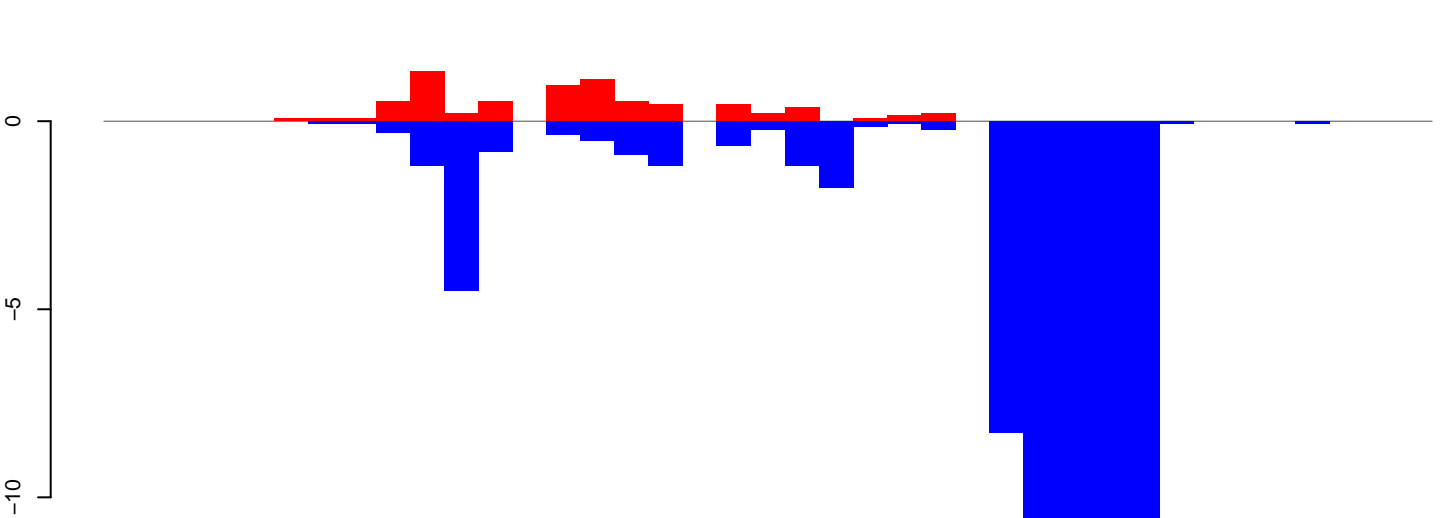
Window size=25, length=722, TE@rnd-2_family-89@DNA_Sola-2@3253:1-722

CuQuin_Testes_NL.18_23.rep



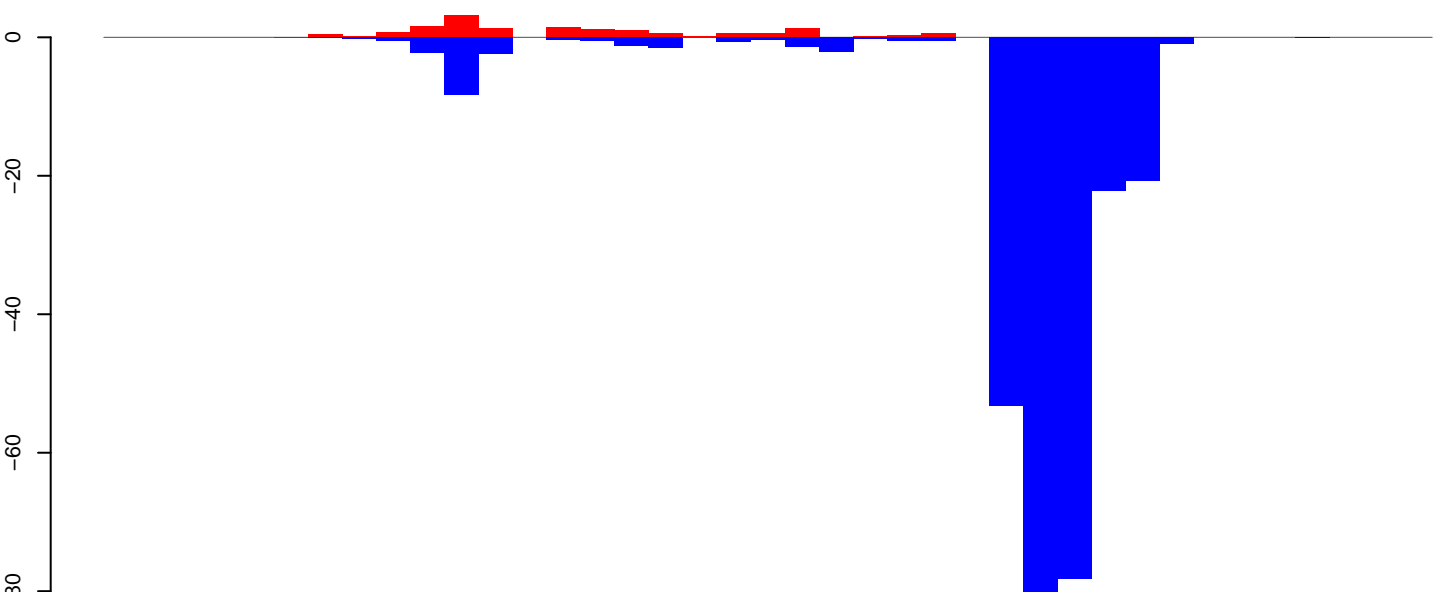
positive=8, negative=52, total=61

CuQuin_Testes_NL.24_35.rep



positive=7, negative=73, total=80

CuQuin_Testes_NL.rep

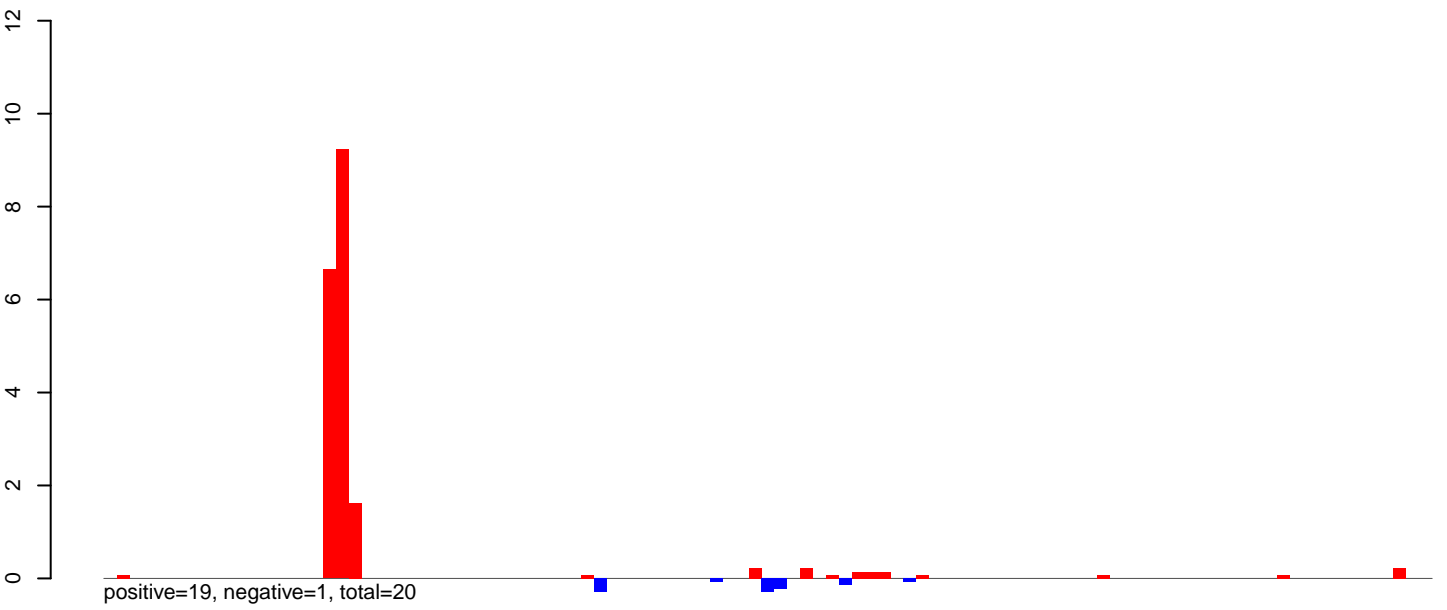


positive=16, negative=281, total=297

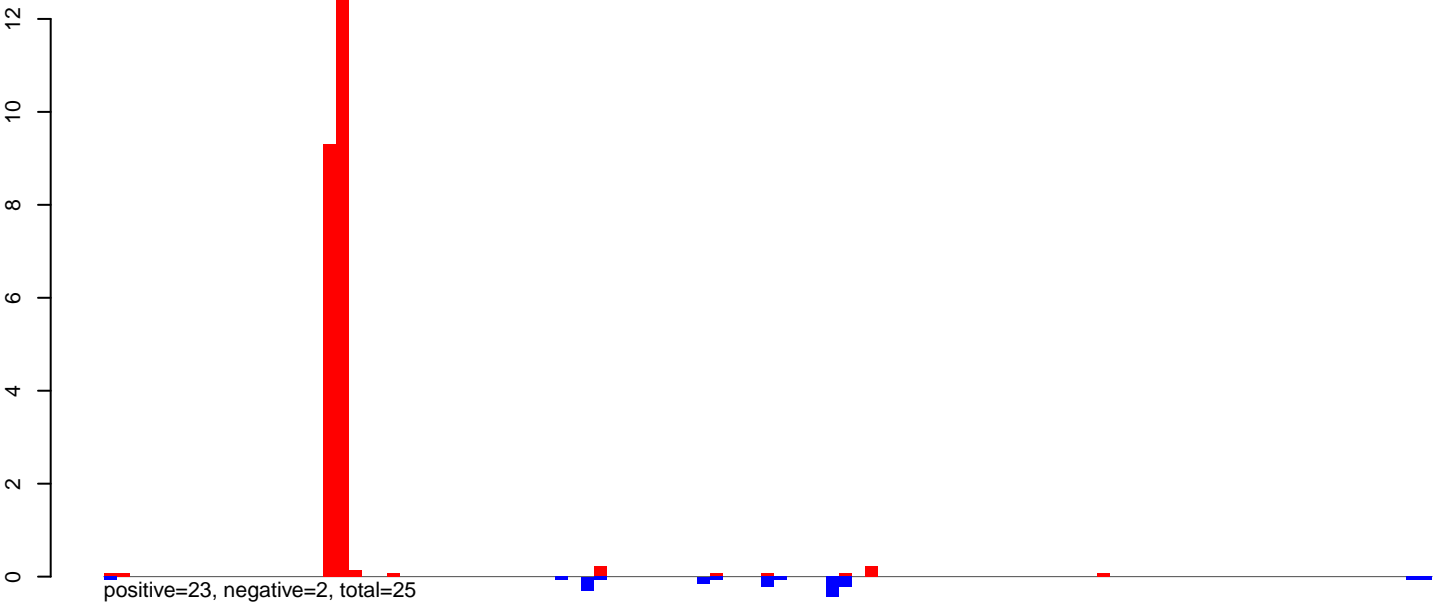
Window size=25, length=962, TE@rnd-5, family-881@SINE_tRNA@1731:1-962

0 200 400 600 800 1000

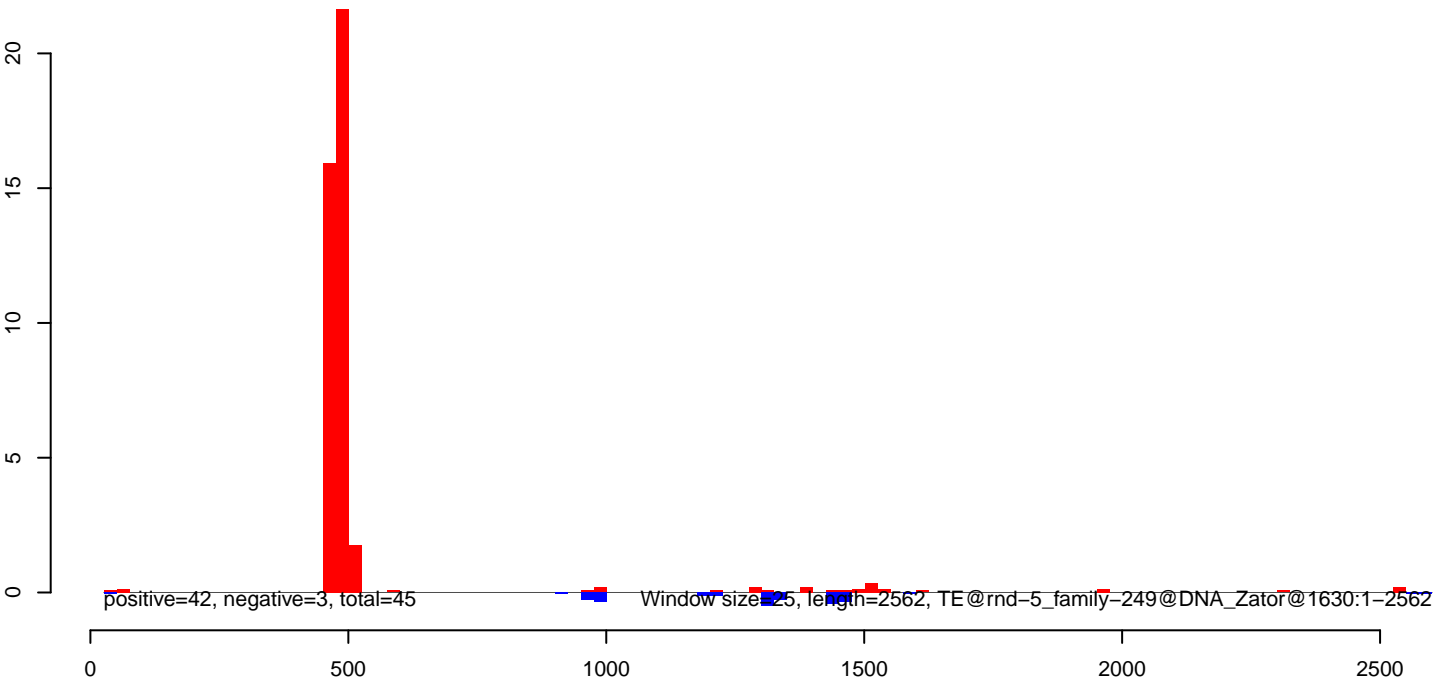
CuQuin_Testes_NL.18_23.rep



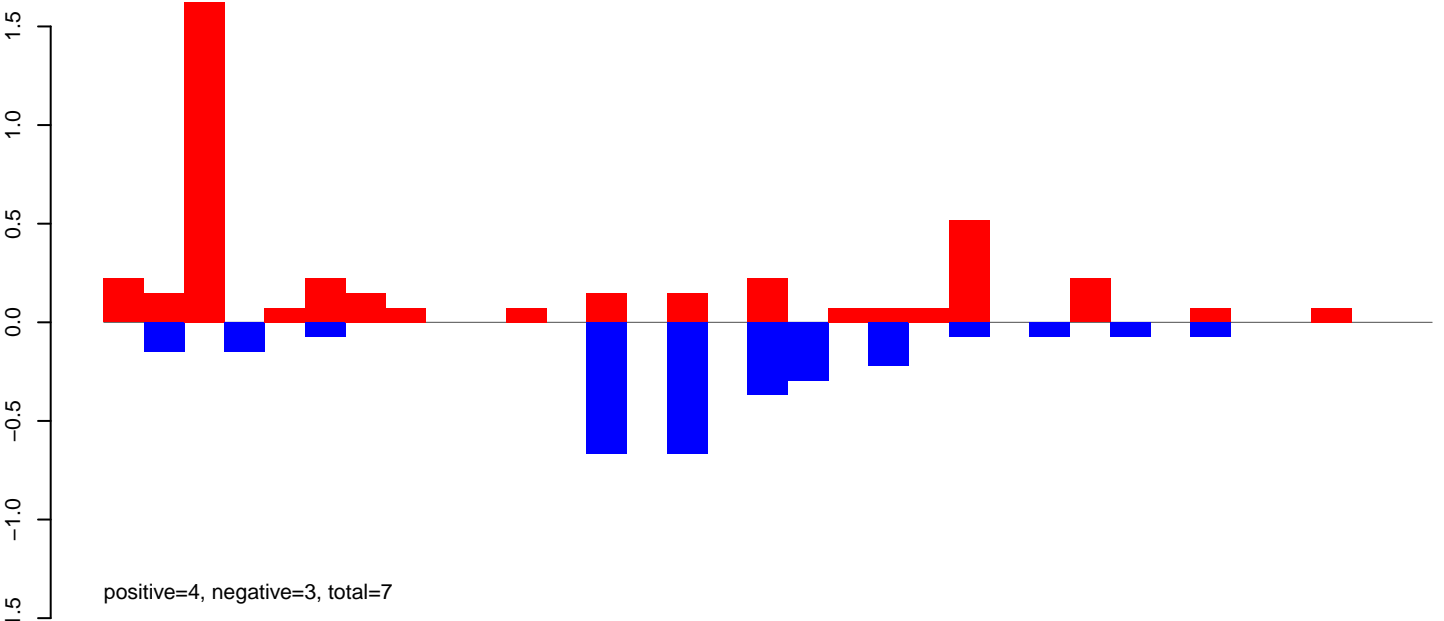
CuQuin_Testes_NL.24_35.rep



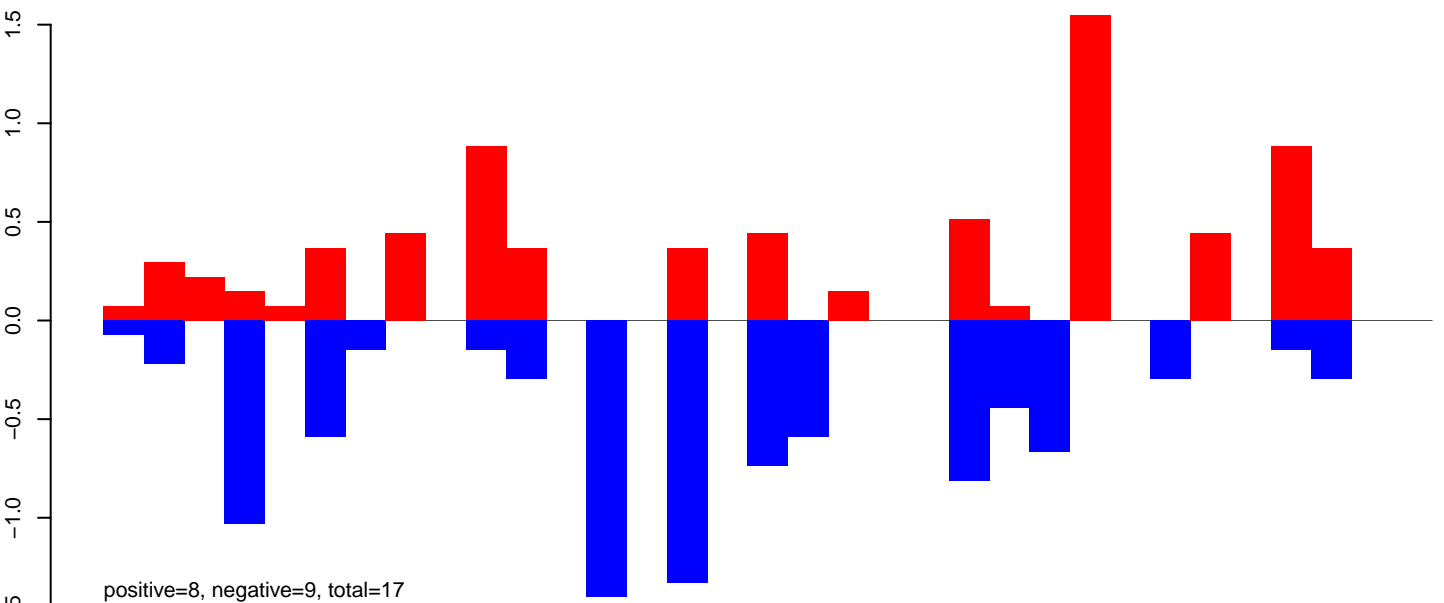
CuQuin_Testes_NL.rep



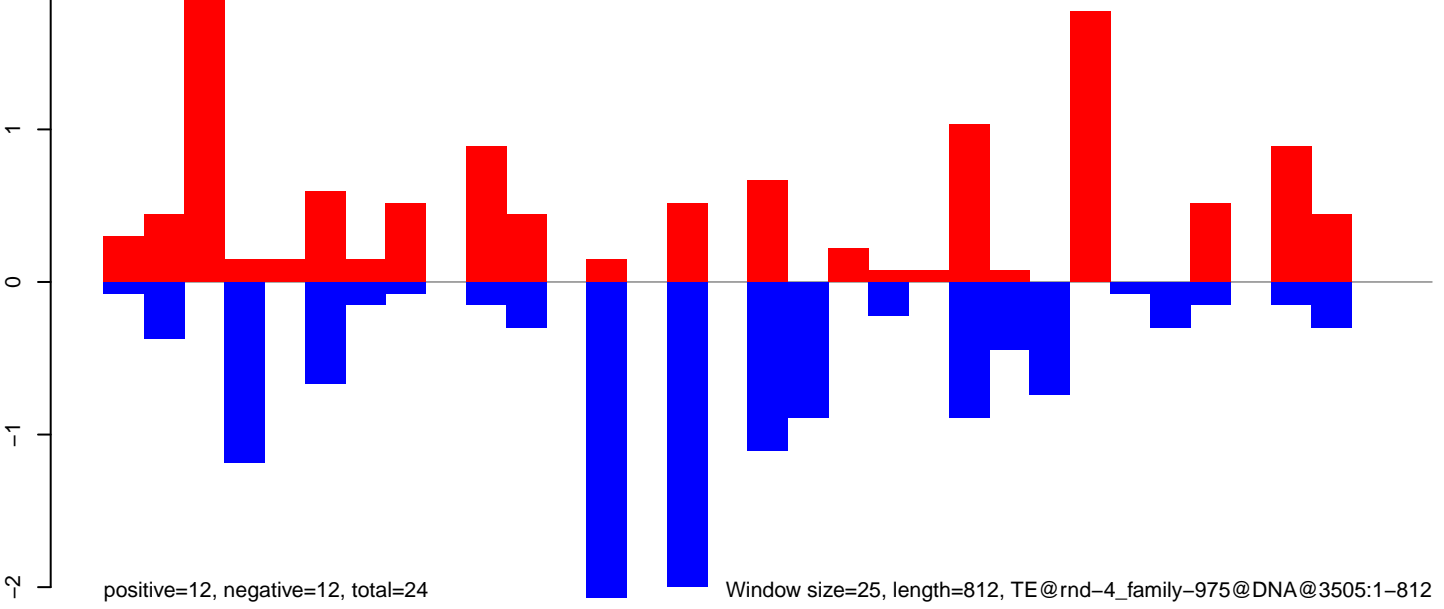
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

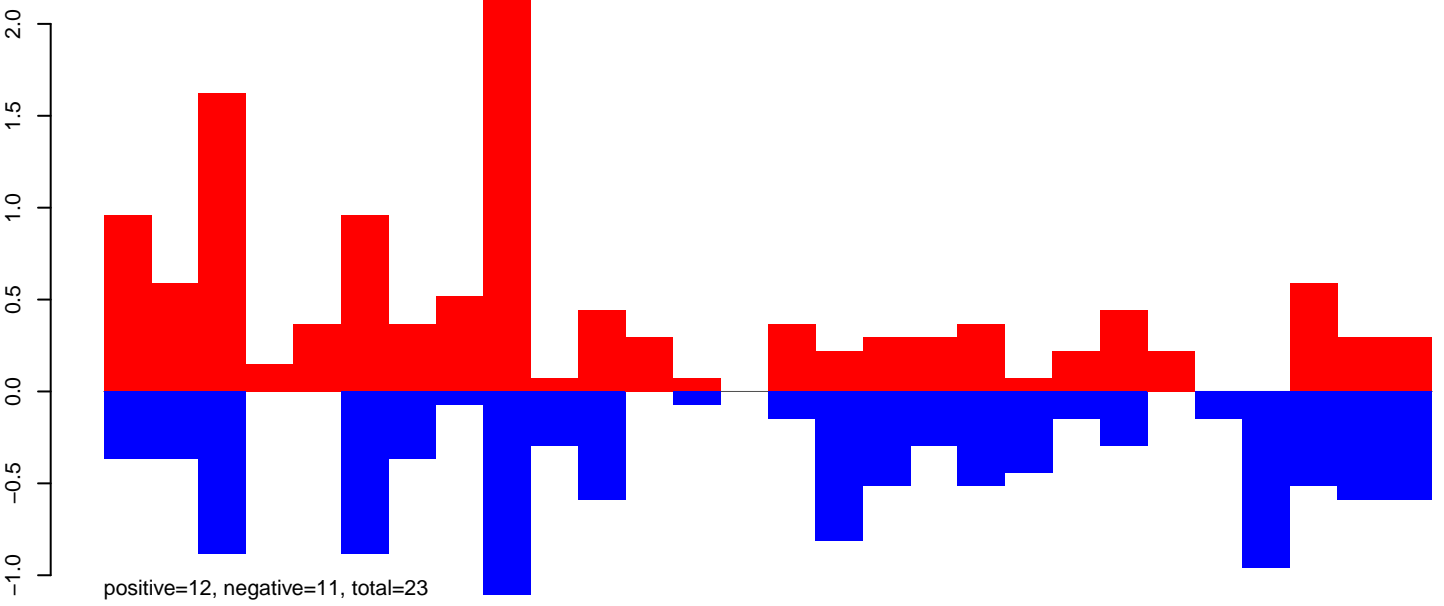


CuQuin_Testes_NL.rep

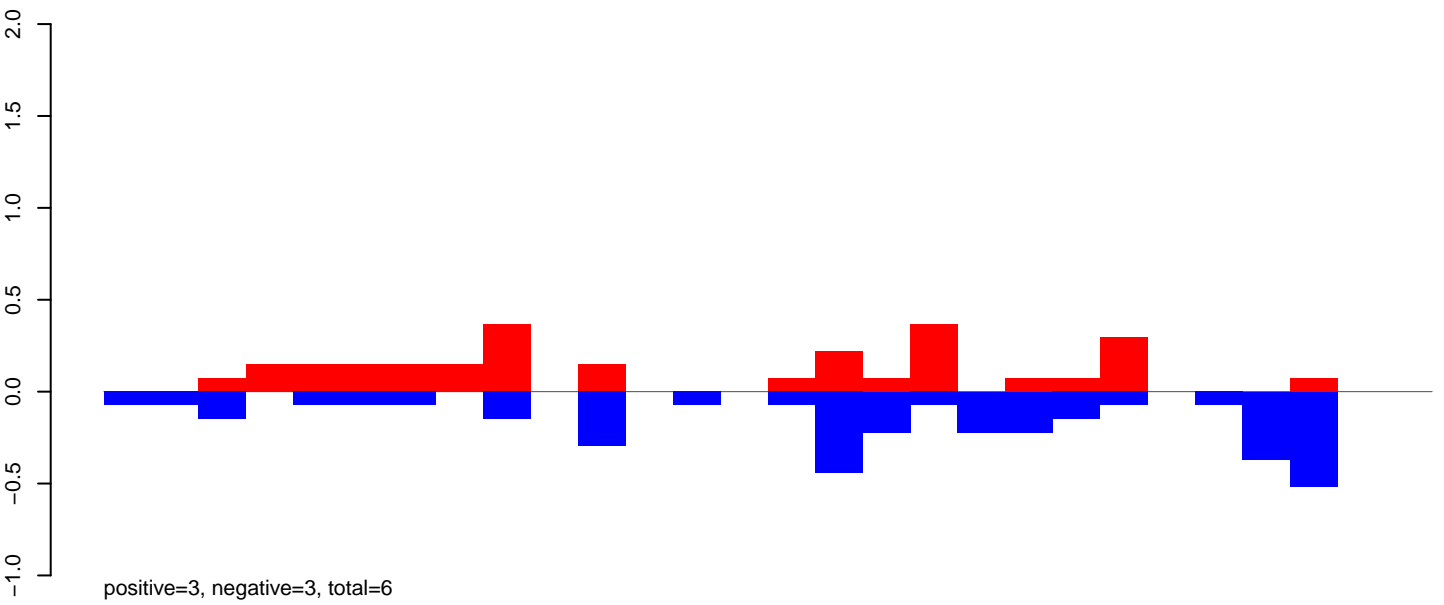


Window size=25, length=812, TE@rnd-4_family-975@DNA@3505:1-812

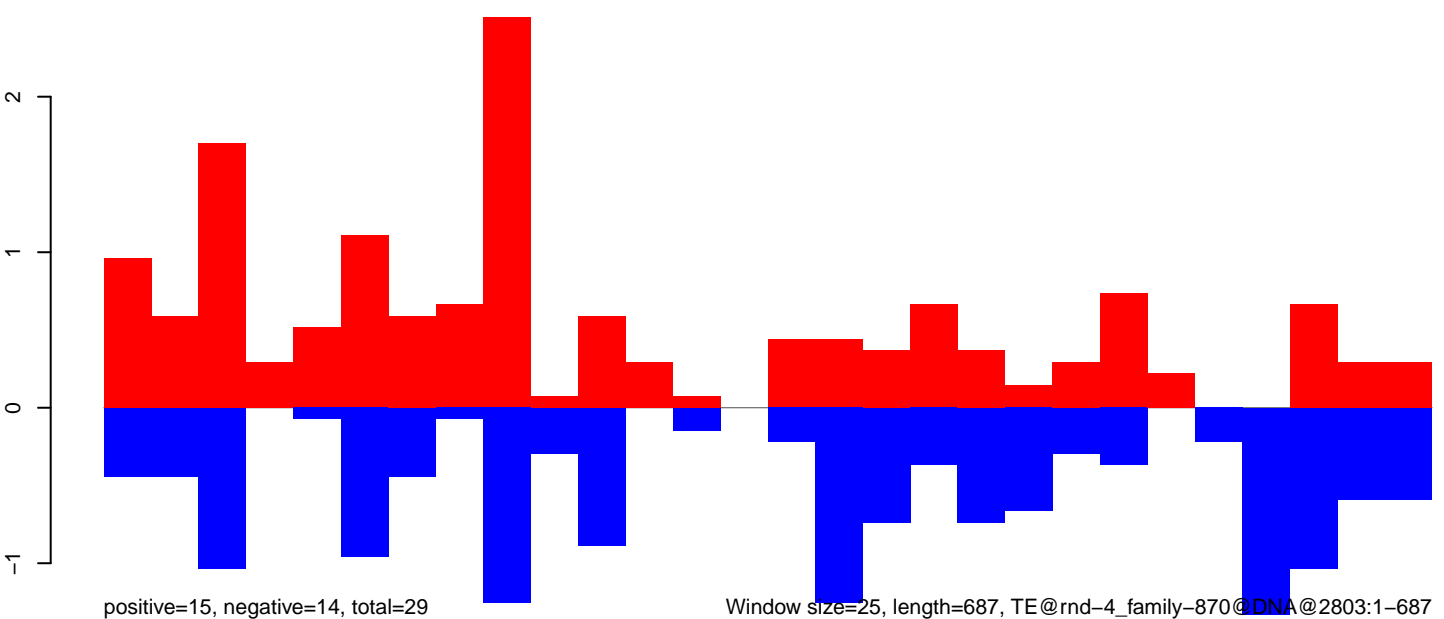
CuQuin_Testes_NL.18_23.rep



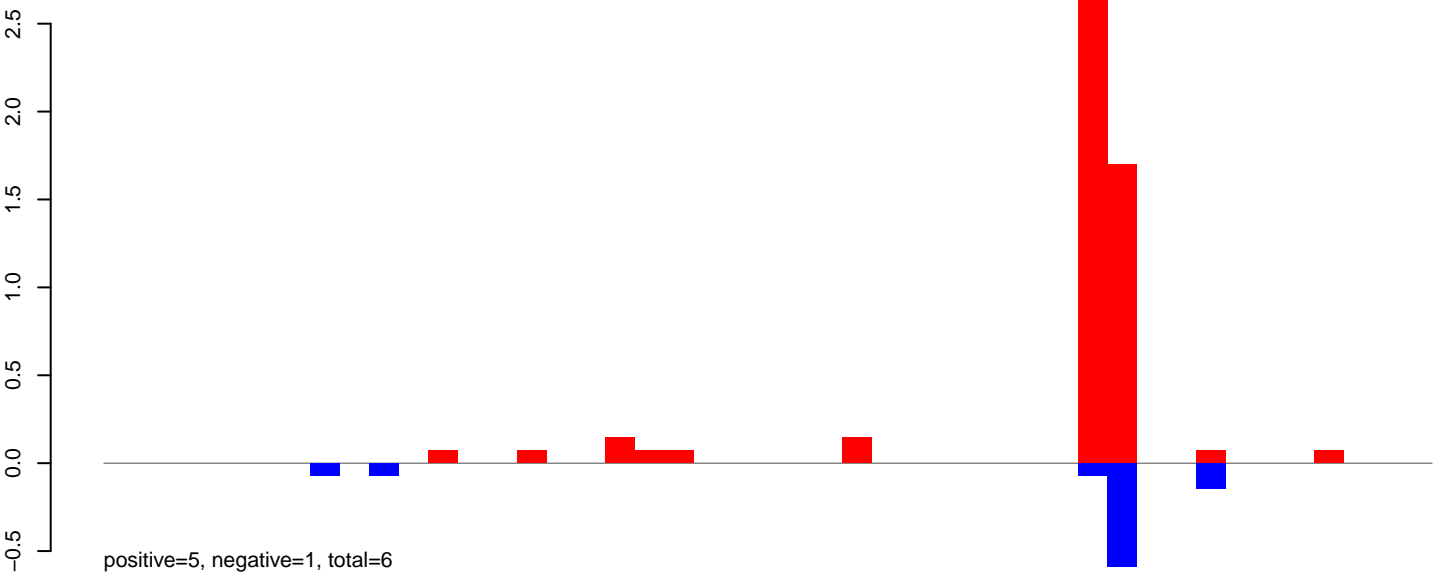
CuQuin_Testes_NL.24_35.rep



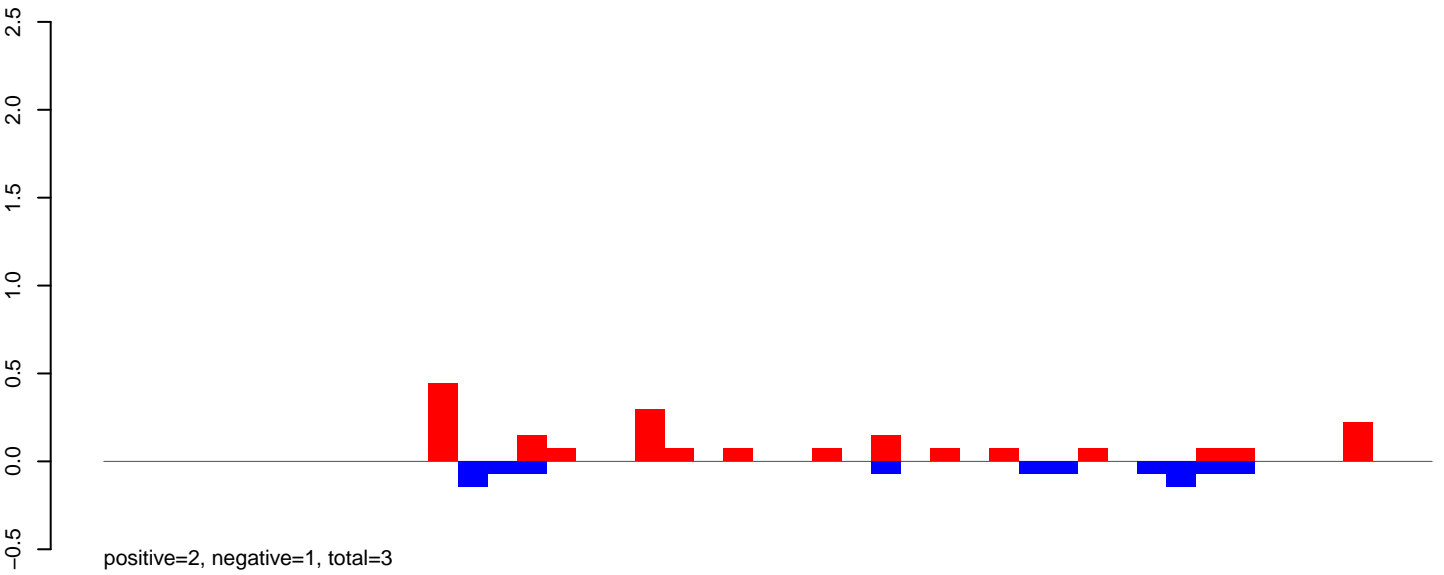
CuQuin_Testes_NL.rep



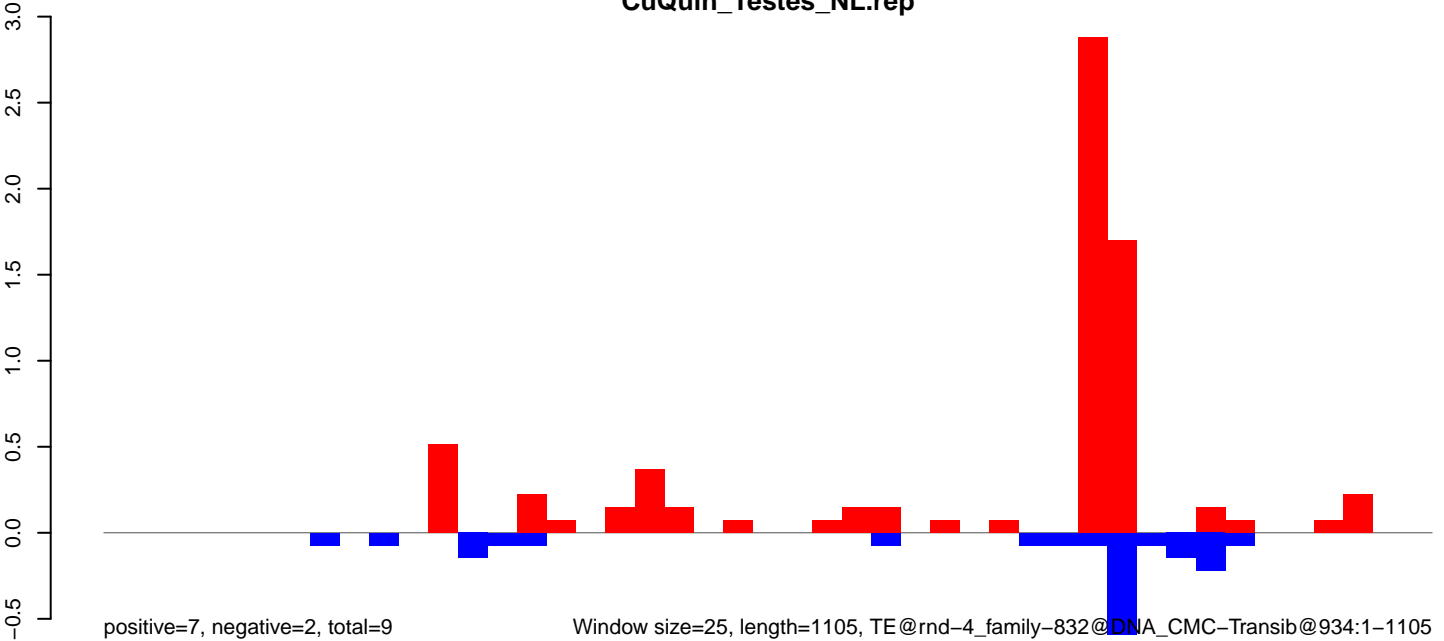
CuQuin_Testes_NL.18_23.rep



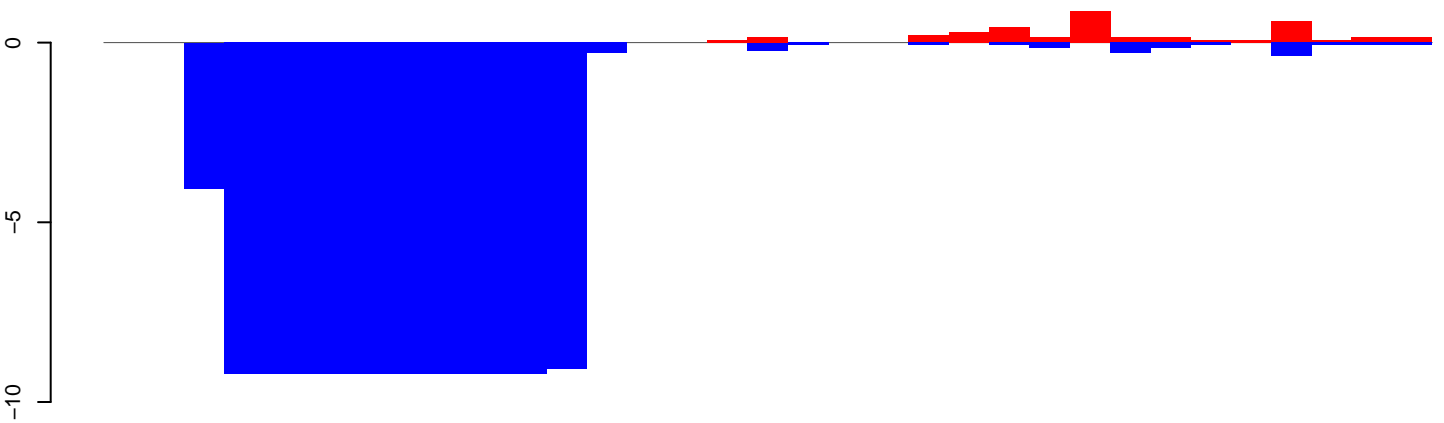
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

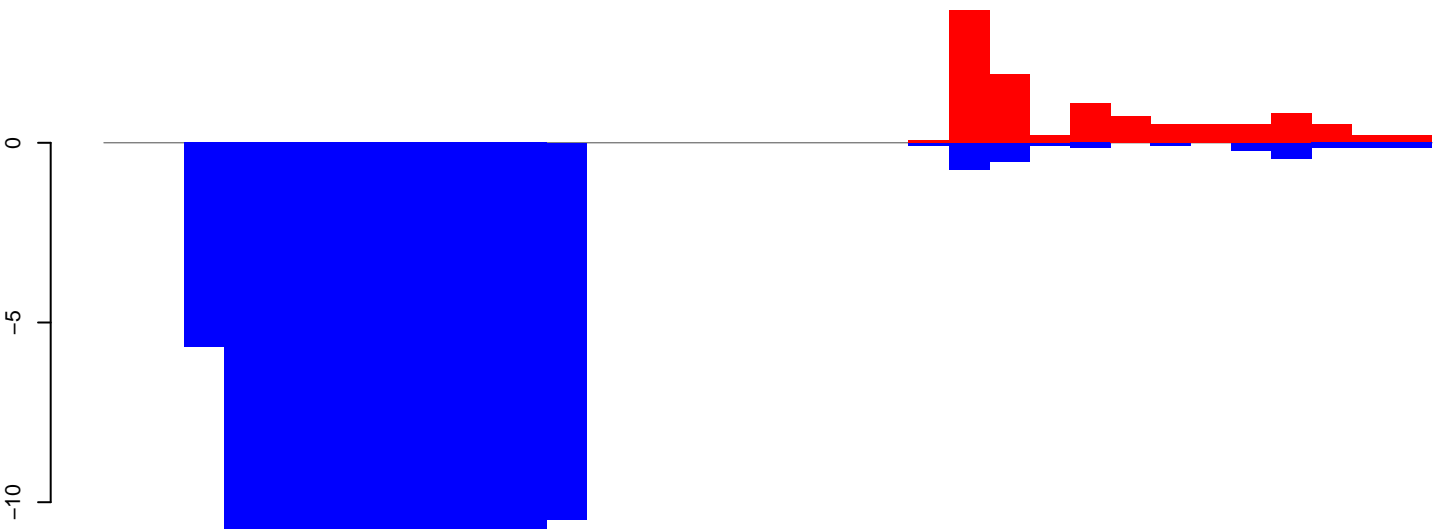


CuQuin_Testes_NL.18_23.rep



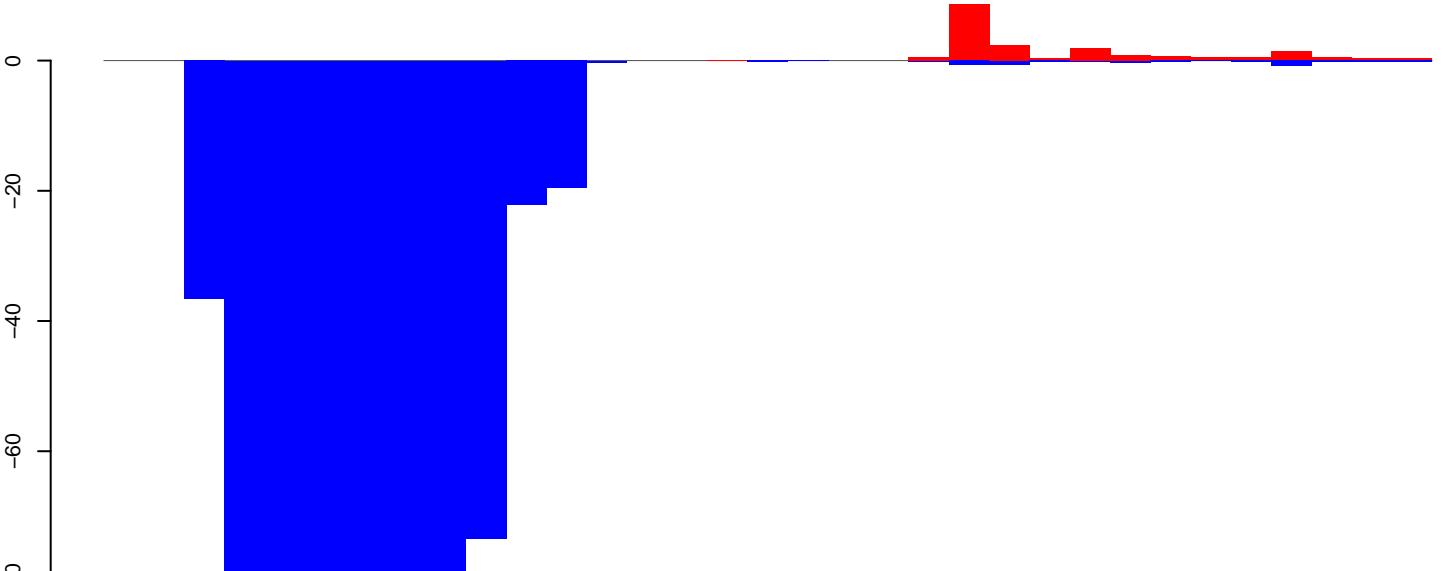
positive=4, negative=89, total=93

CuQuin_Testes_NL.24_35.rep



positive=11, negative=122, total=133

CuQuin_Testes_NL.rep

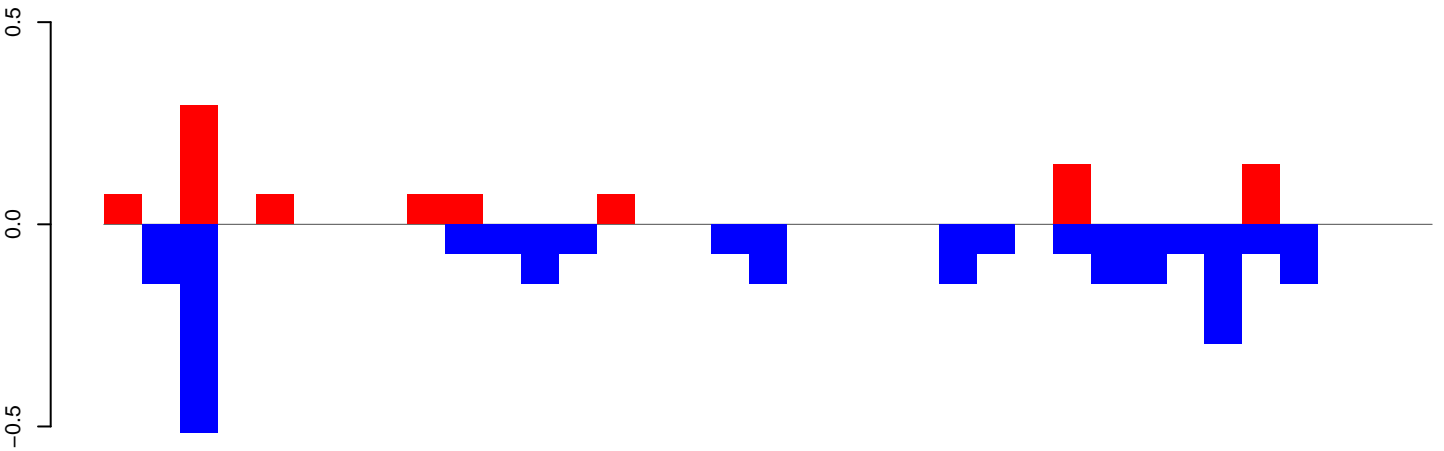


positive=20, negative=655, total=675

Window size=25, length=818, TE@rnd-4_family-316@LTR_Gypsy@1481:1-818

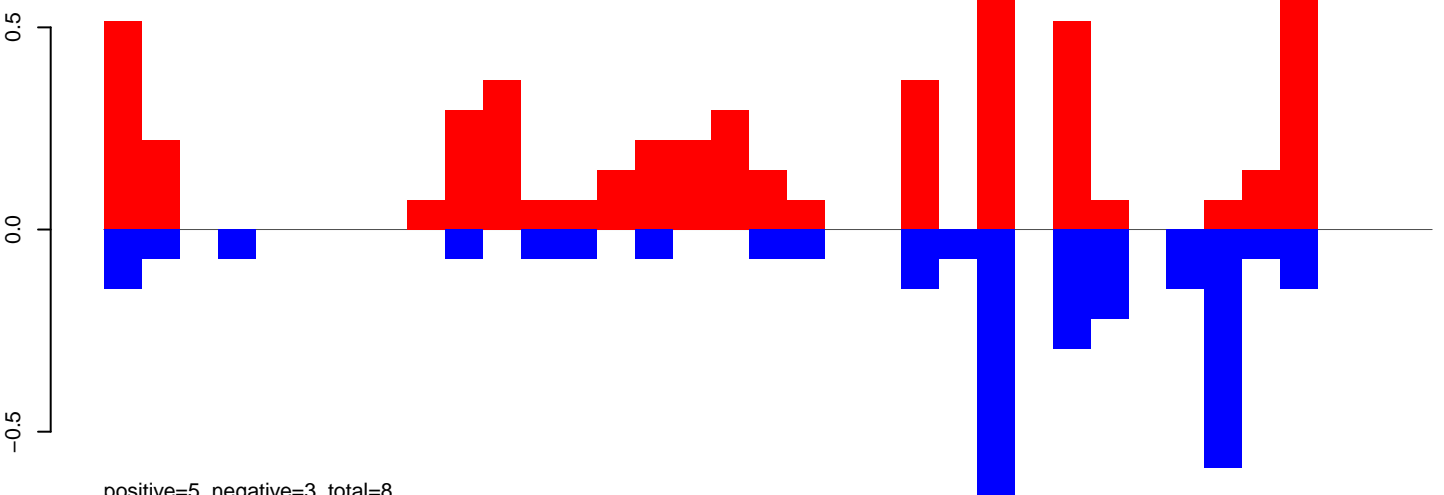
0 200 400 600 800

CuQuin_Testes_NL.18_23.rep



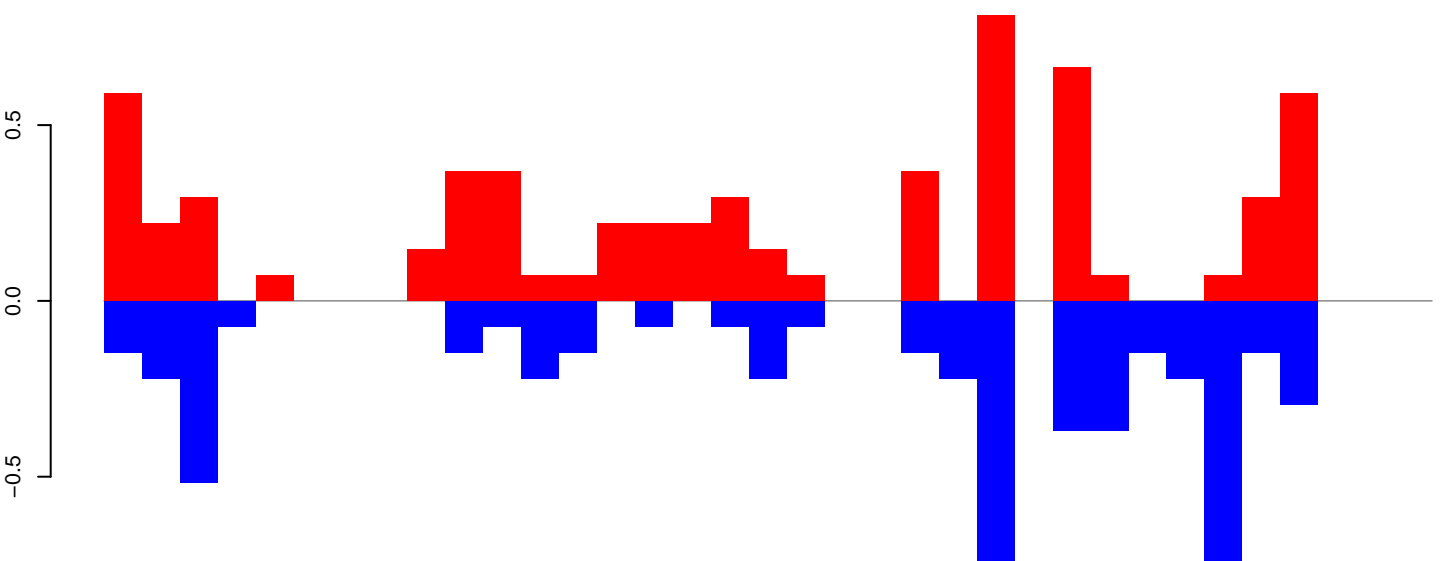
positive=1, negative=2, total=3

CuQuin_Testes_NL.24_35.rep



positive=5, negative=3, total=8

CuQuin_Testes_NL.rep

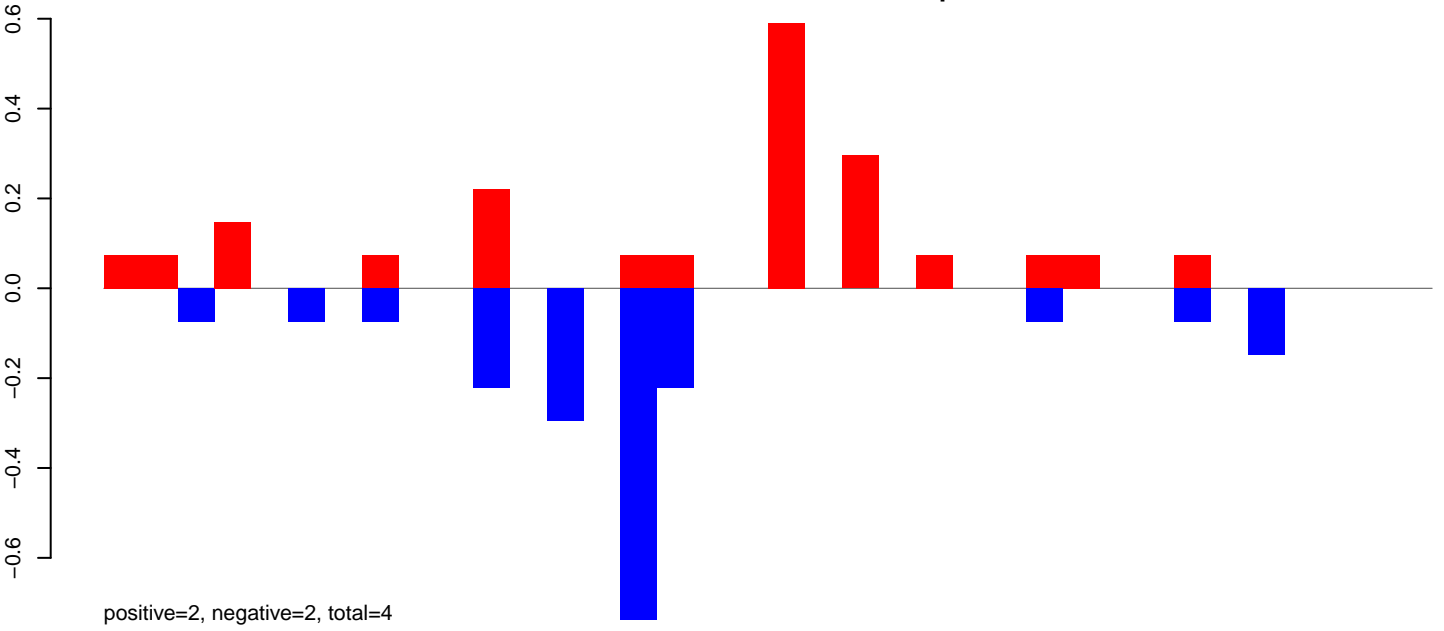


positive=6, negative=6, total=12

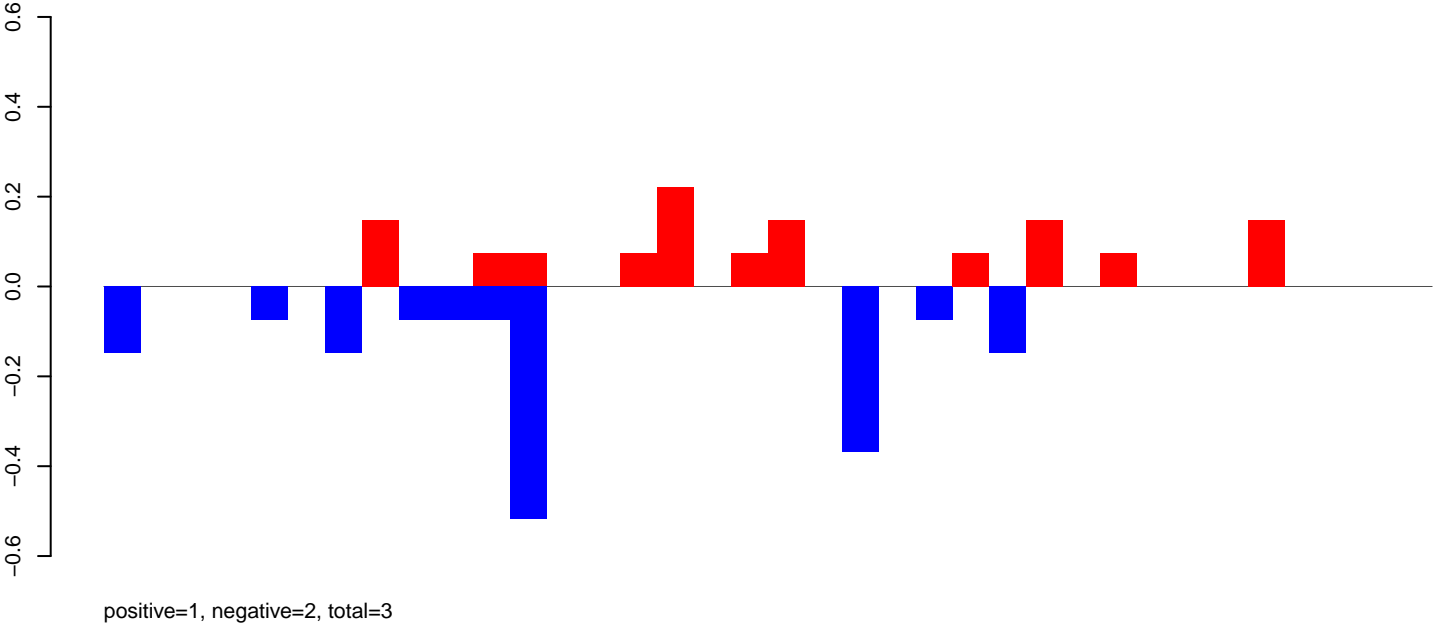
Window size=25, length=869, TE@rnd-4_family-2244@LTR_Gypsy@2241:1-869

0 200 400 600 800

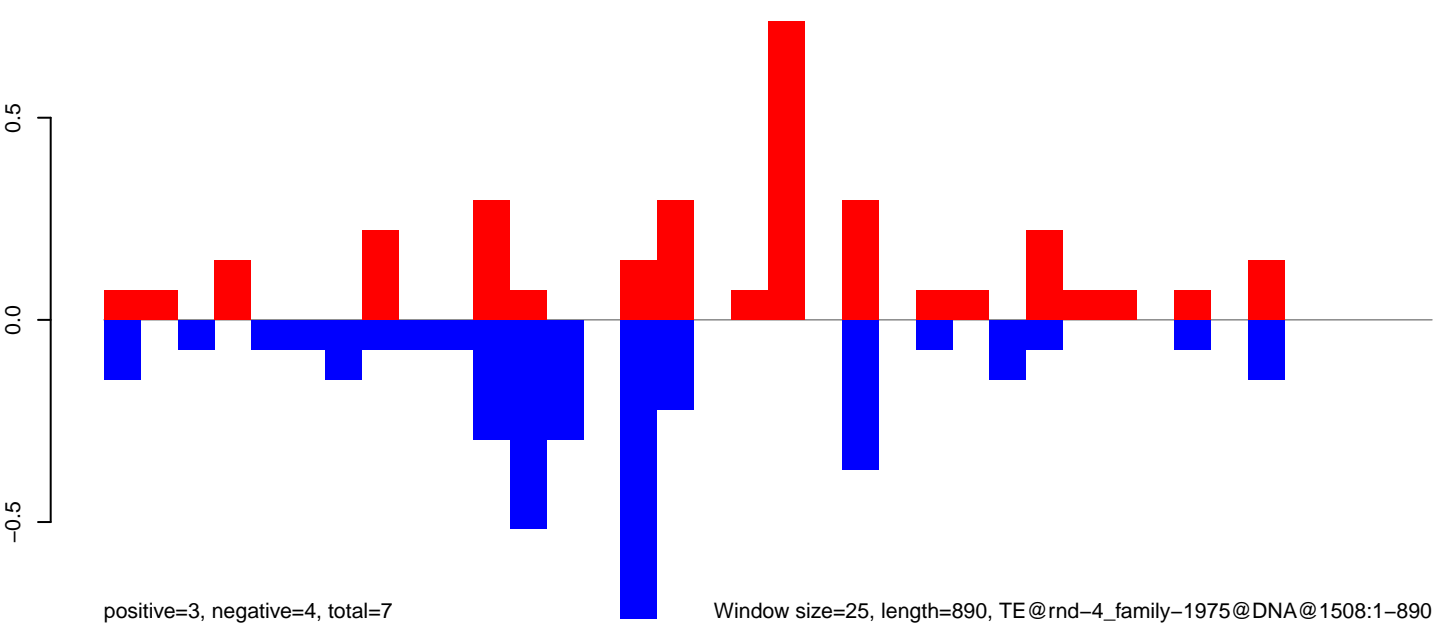
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

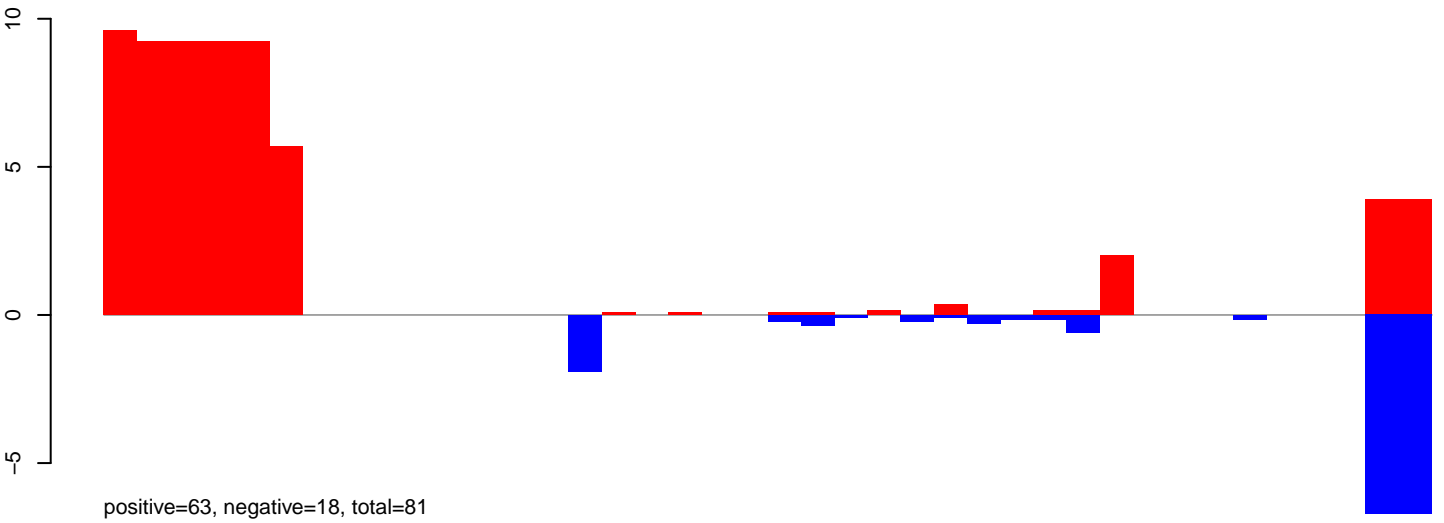


CuQuin_Testes_NL.rep

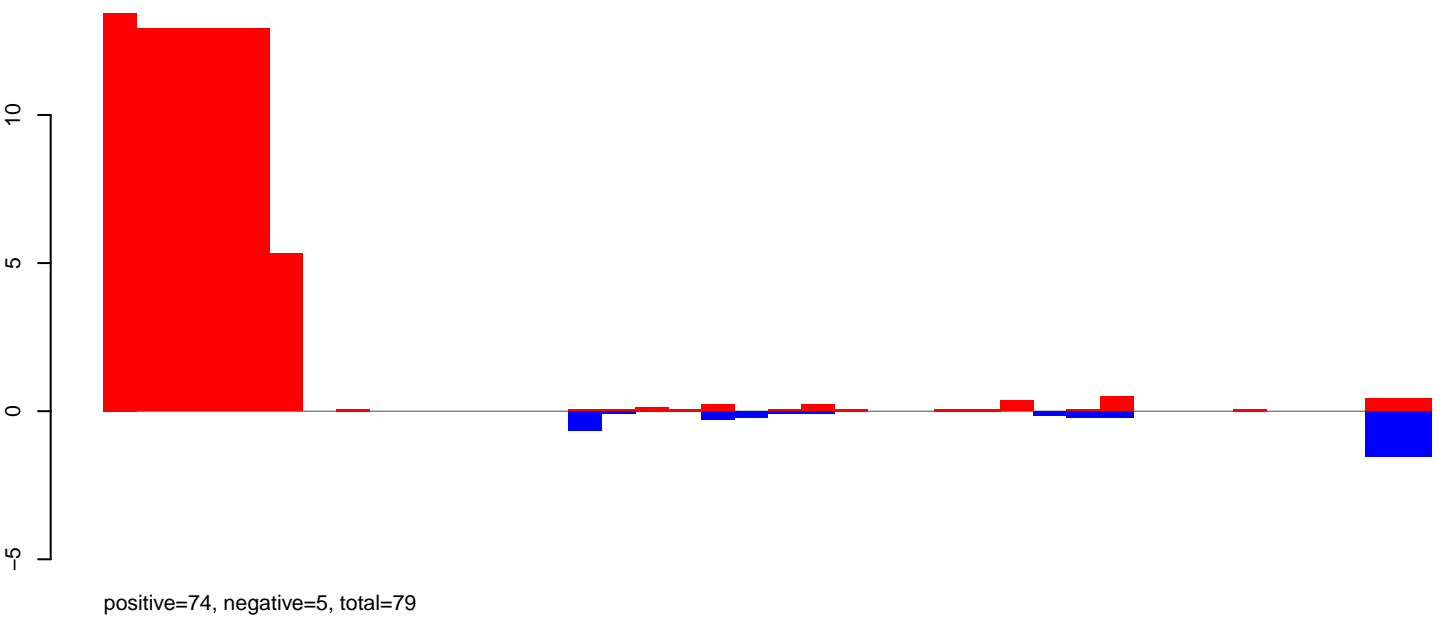


Window size=25, length=890, TE@rnd-4_family-1975@DNA@1508:1-890

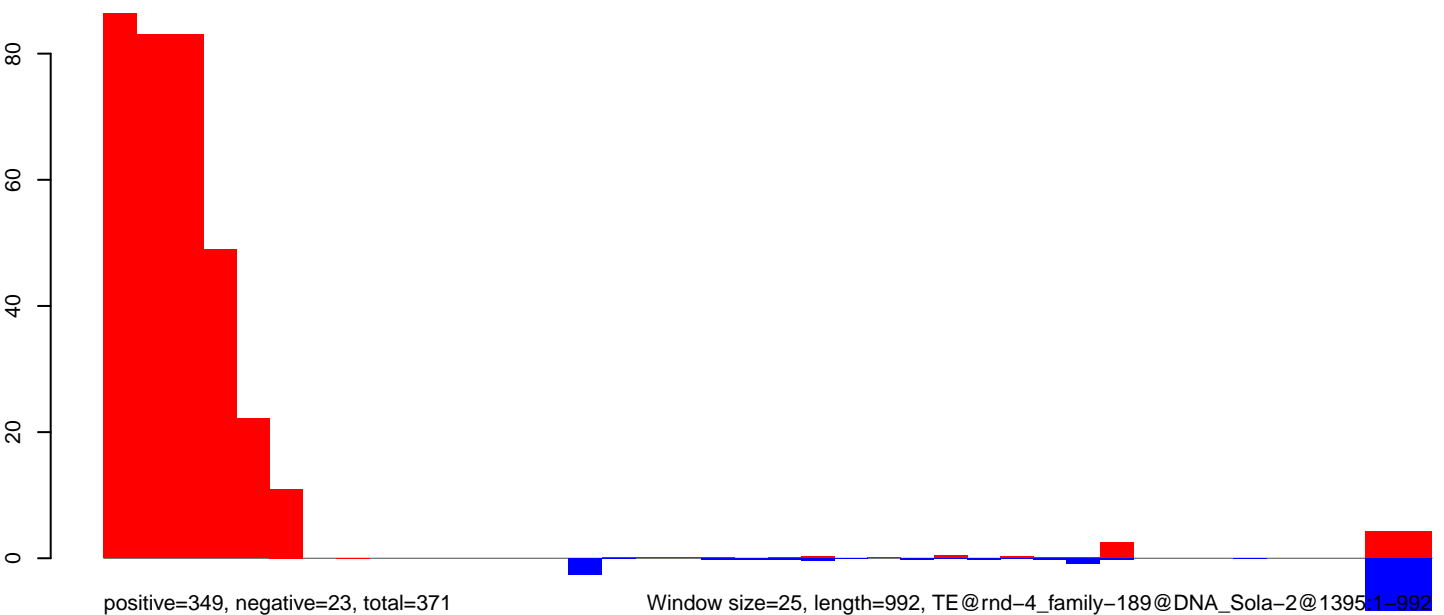
CuQuin_Testes_NL.18_23.rep



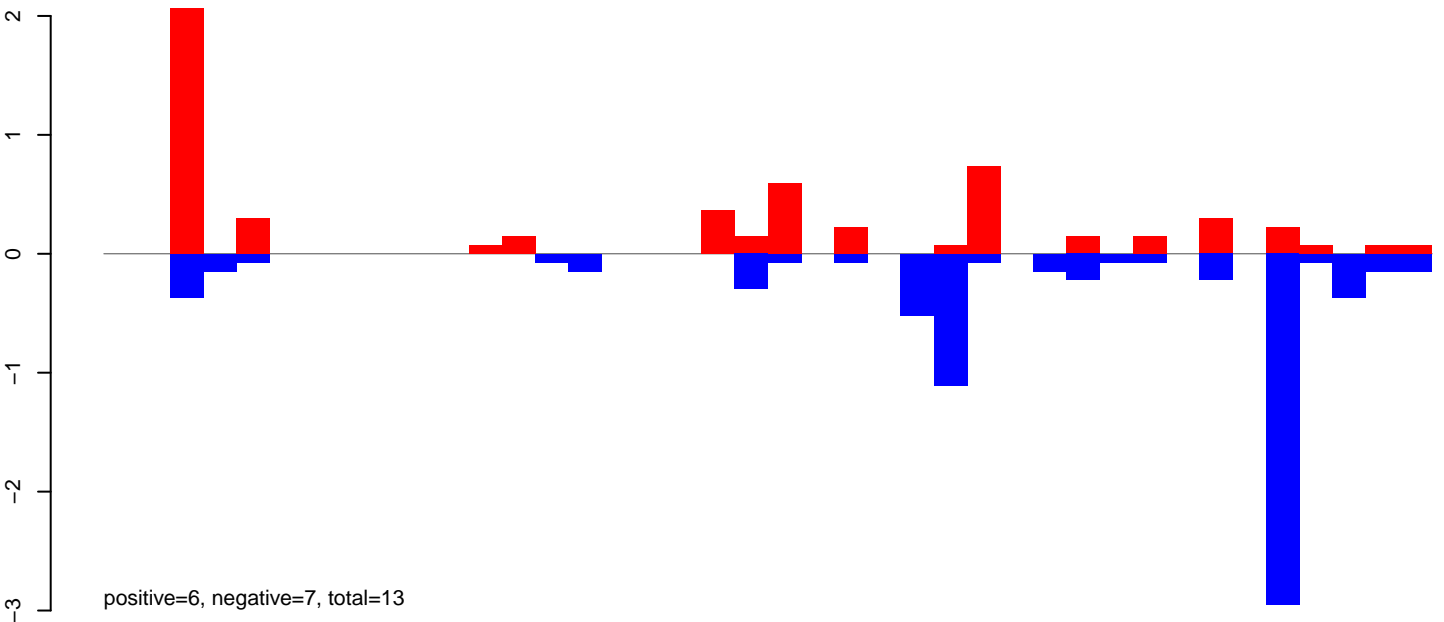
CuQuin_Testes_NL.24_35.rep



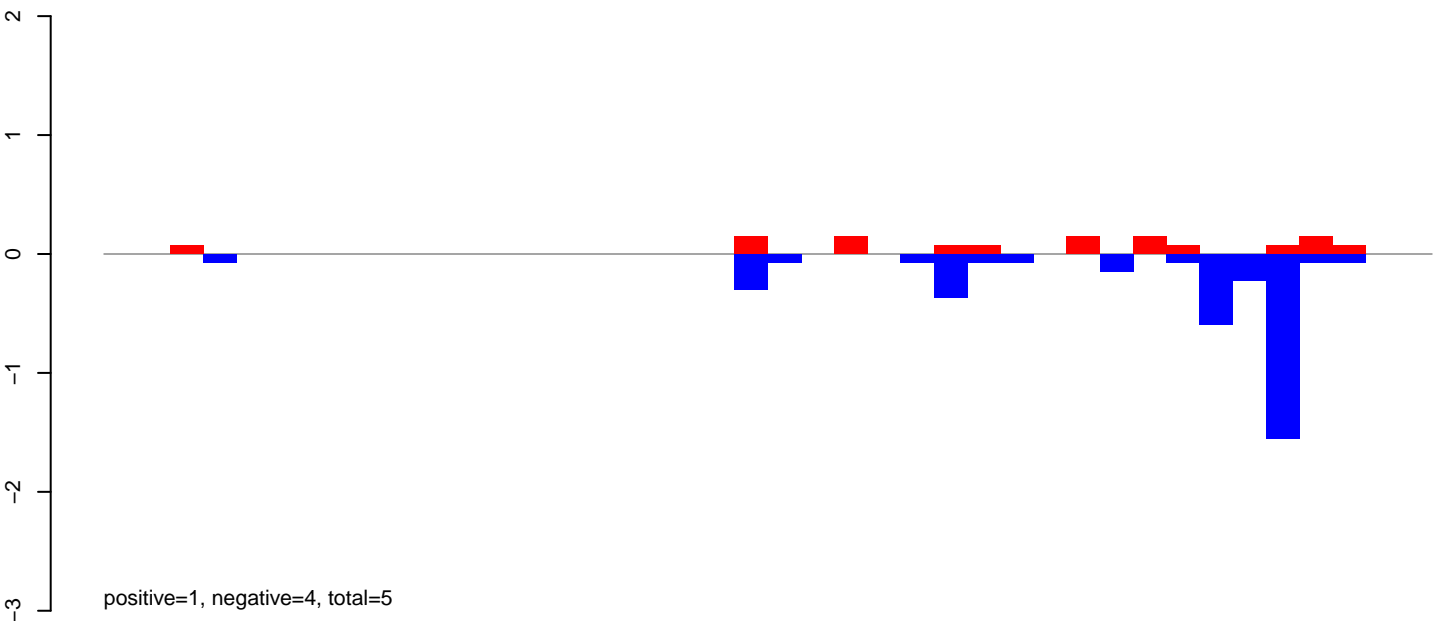
CuQuin_Testes_NL.rep



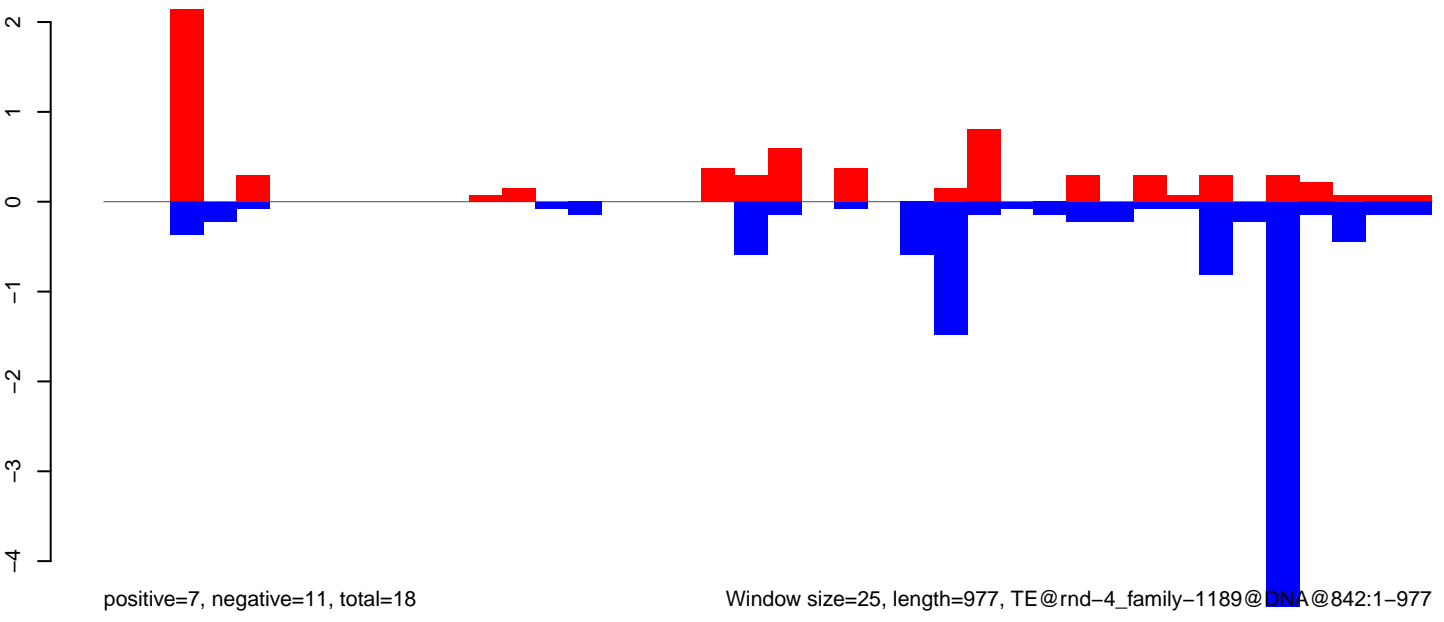
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

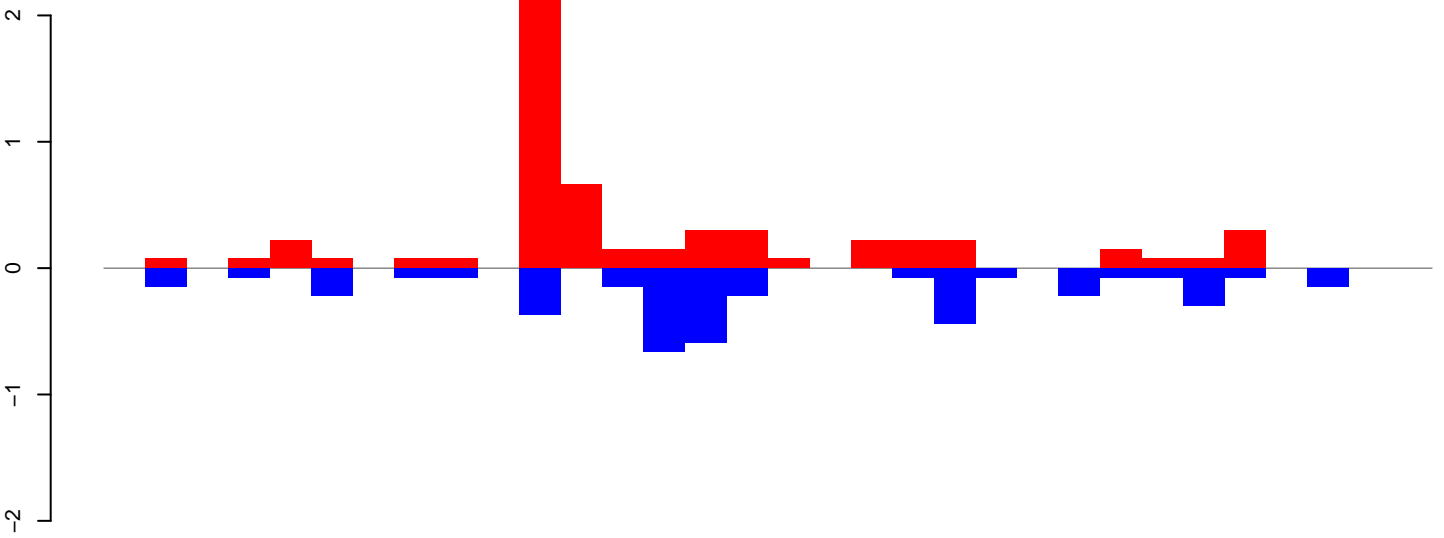


CuQuin_Testes_NL.rep



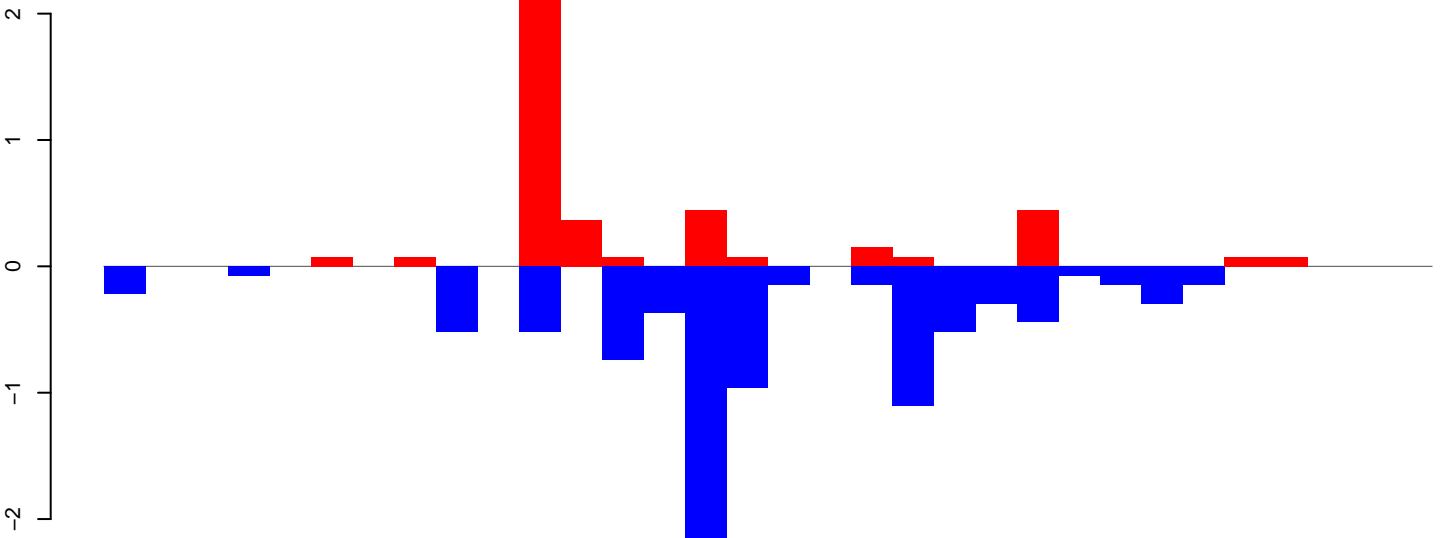
Window size=25, length=977, TE@rnd-4_family-1189@DNA@842:1-977

CuQuin_Testes_NL.18_23.rep



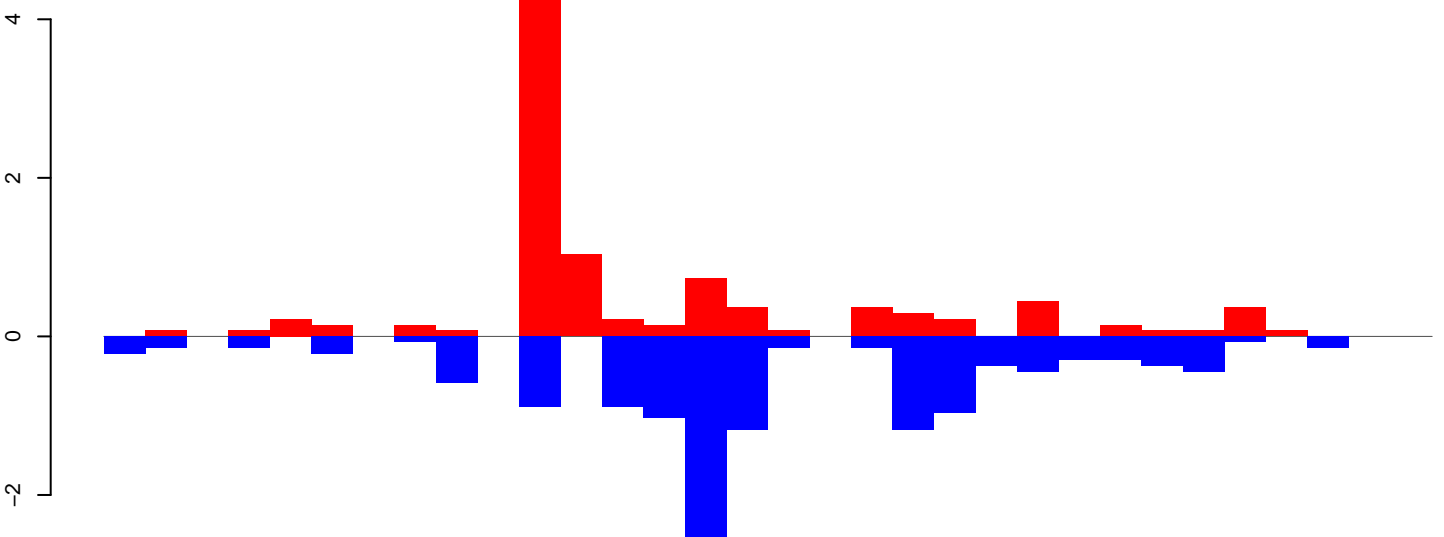
positive=6, negative=4, total=10

CuQuin_Testes_NL.24_35.rep



positive=4, negative=9, total=13

CuQuin_Testes_NL.rep

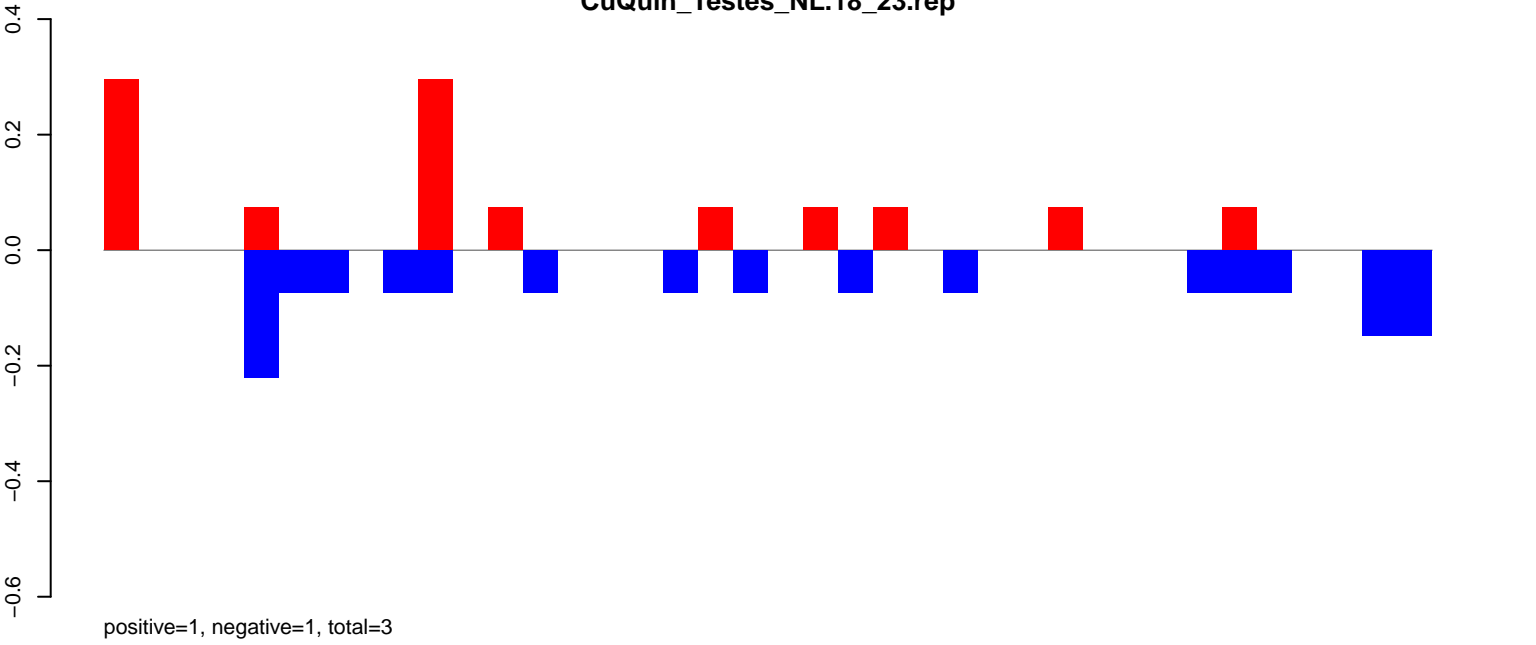


positive=10, negative=13, total=23

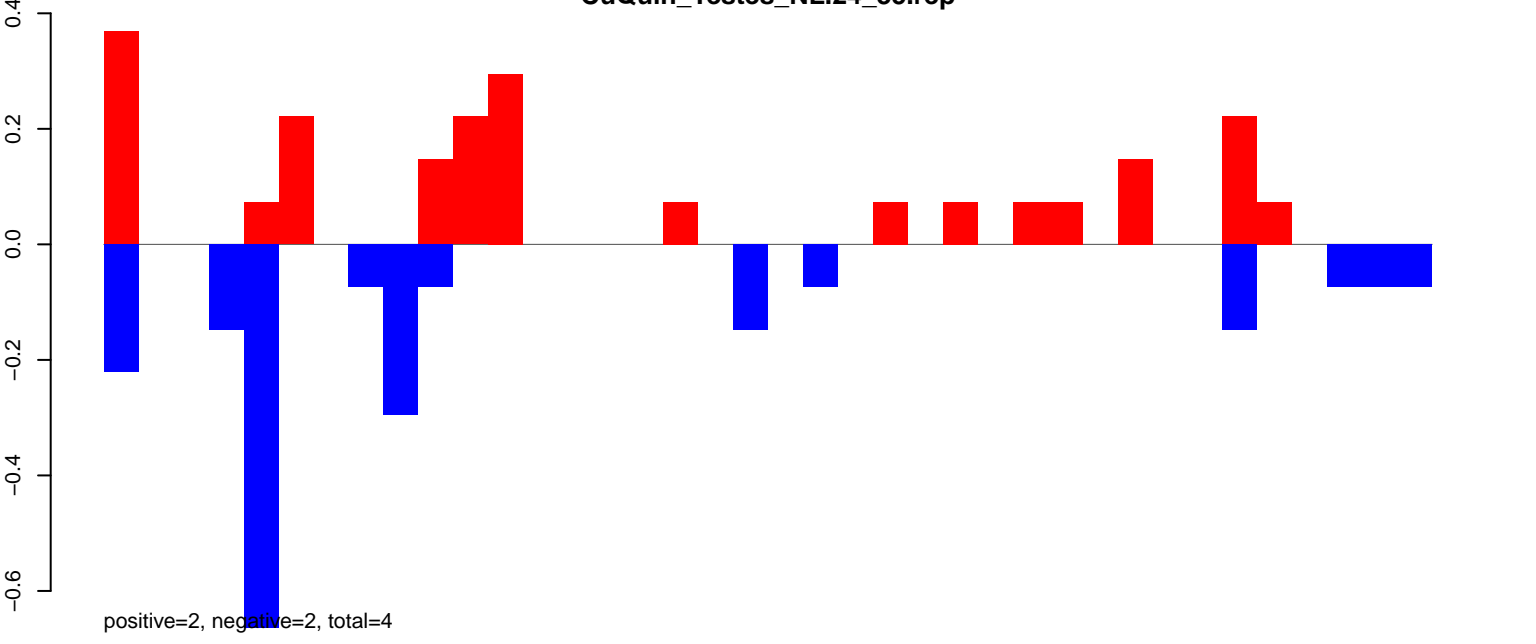
Window size=25, length=781, TE@rnd-4_family-1030@DNA@1572:1-781

0 200 400 600 800

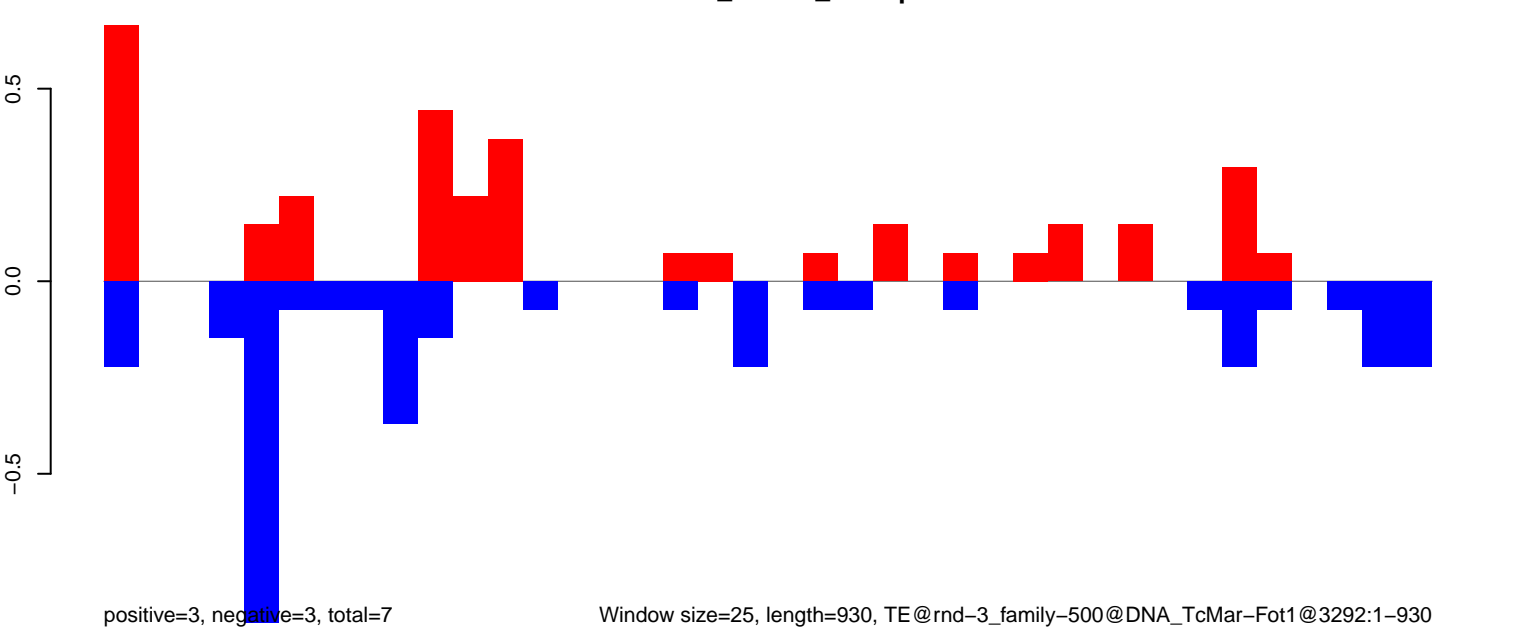
CuQuin_Testes_NL.18_23.rep



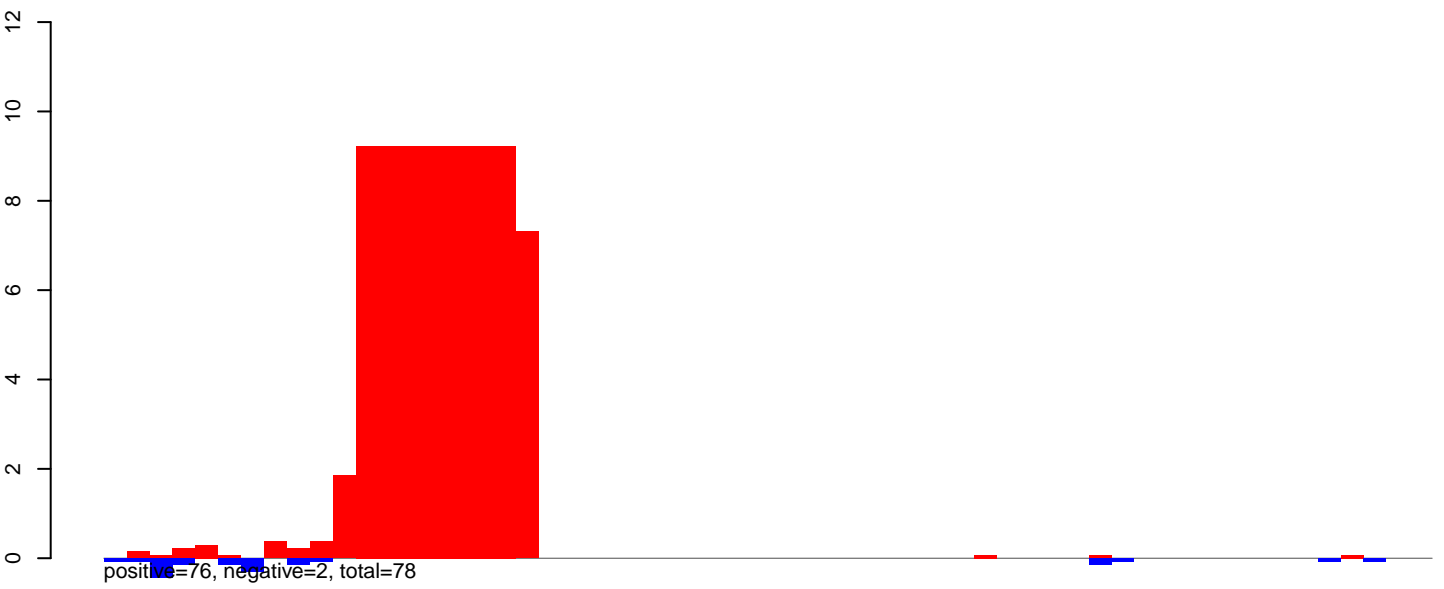
CuQuin_Testes_NL.24_35.rep



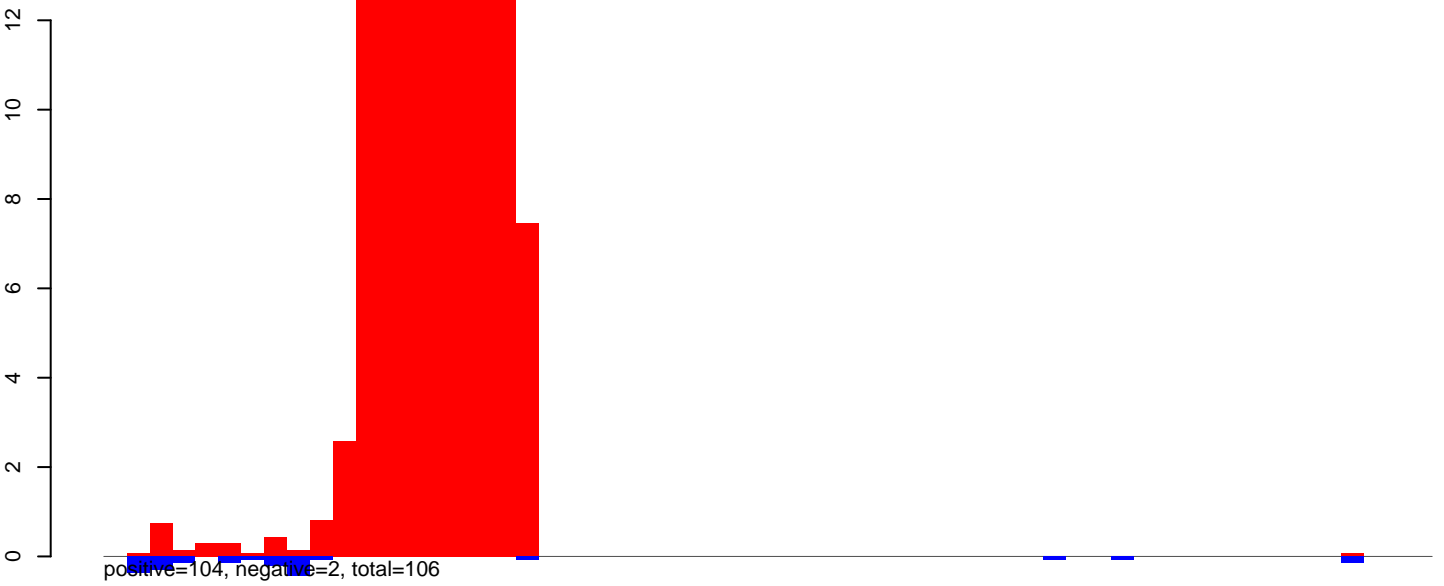
CuQuin_Testes_NL.rep



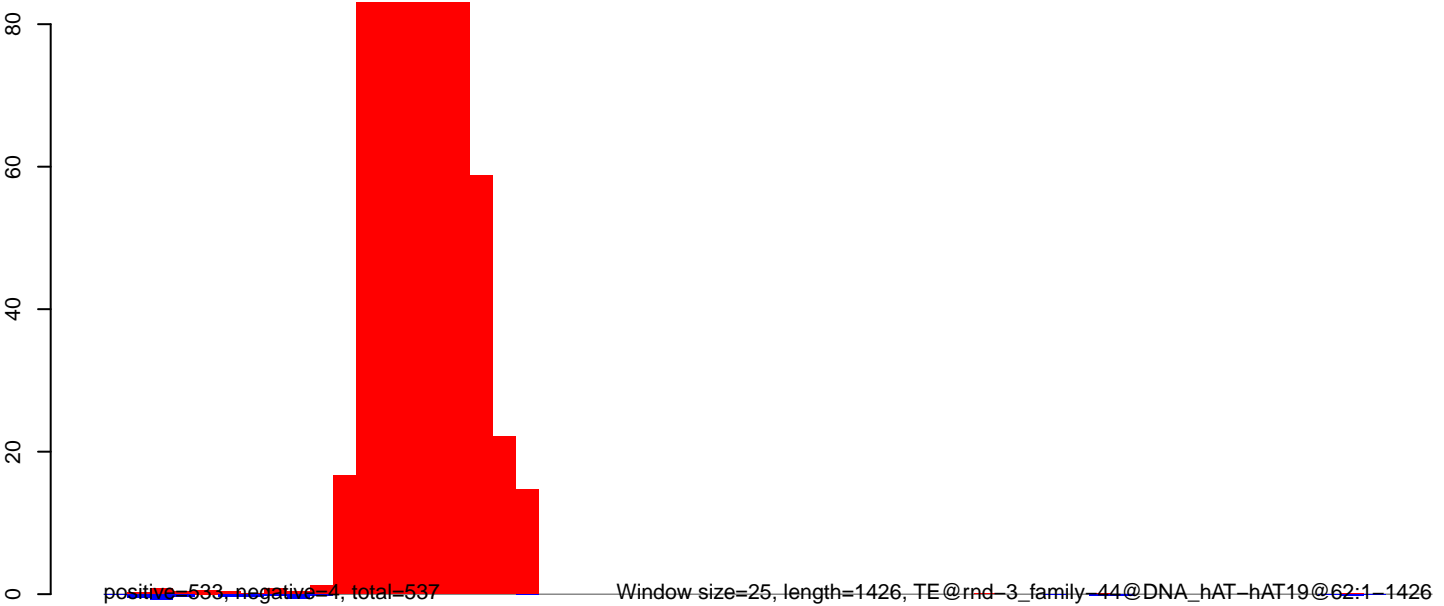
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

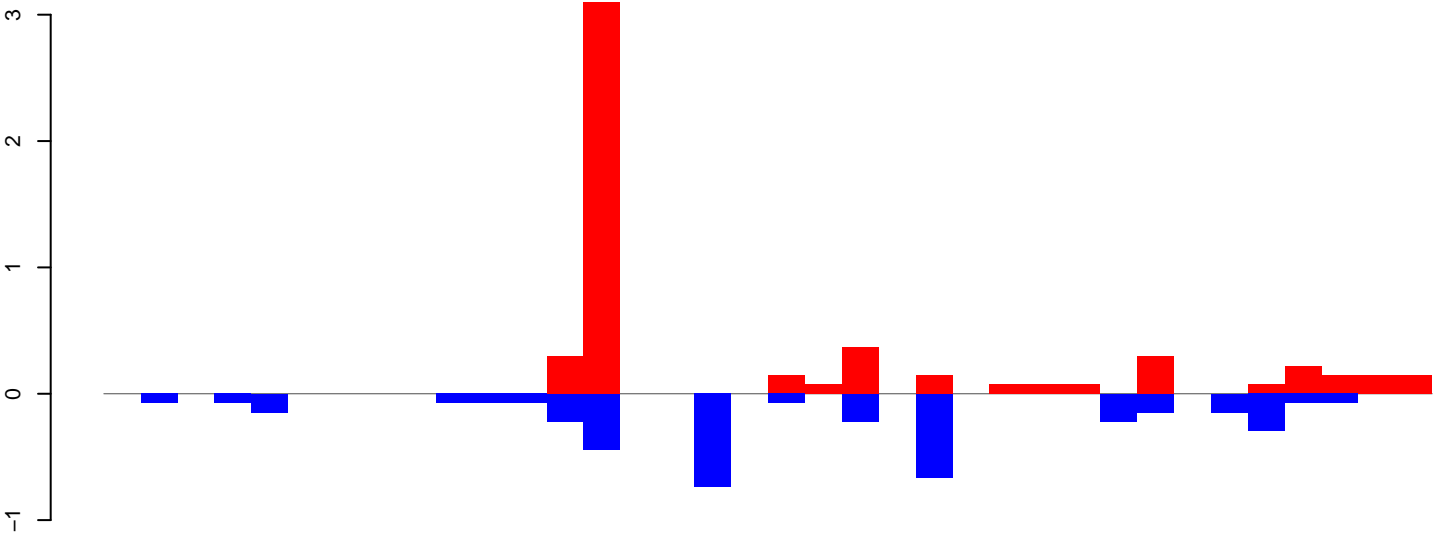


CuQuin_Testes_NL.rep



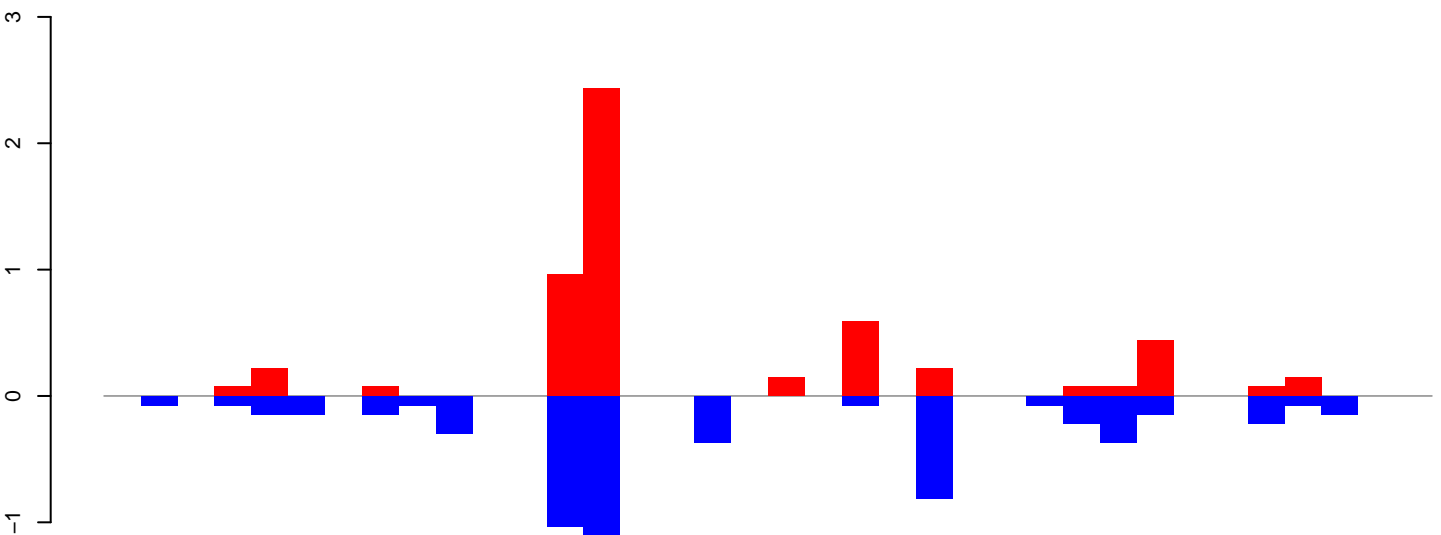
Window size=25, length=1426, TE@rnd-3_family-44@DNA_hAT-hAT19@62-1-1426

CuQuin_Testes_NL.18_23.rep



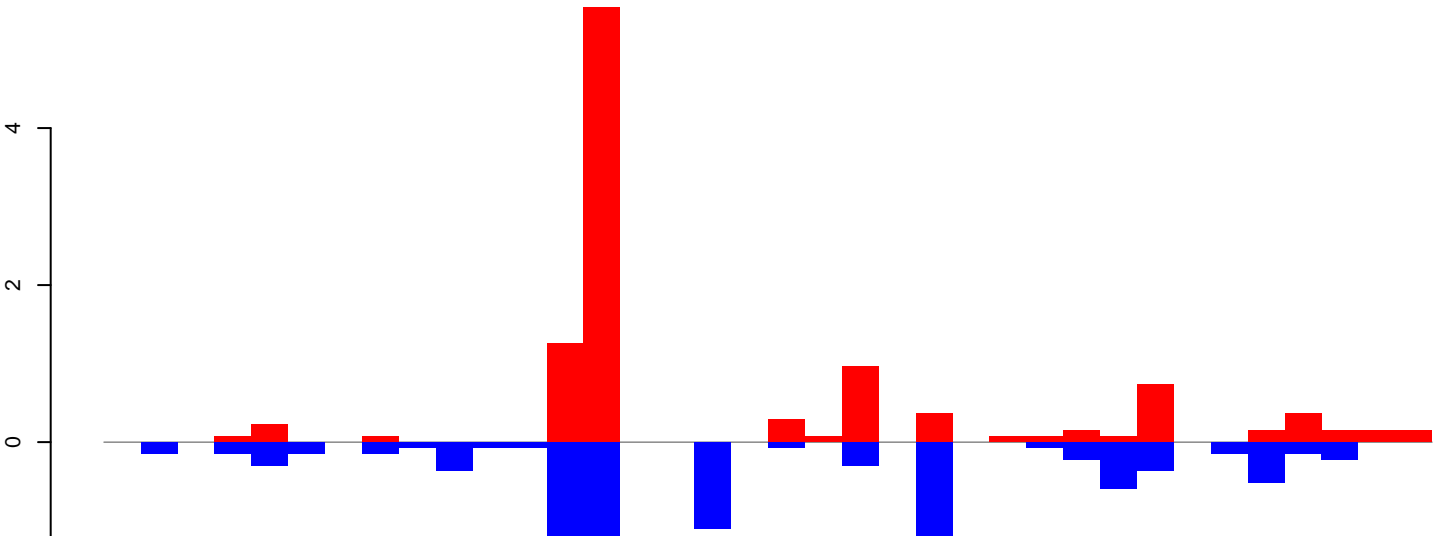
positive=5, negative=4, total=9

CuQuin_Testes_NL.24_35.rep



positive=6, negative=6, total=12

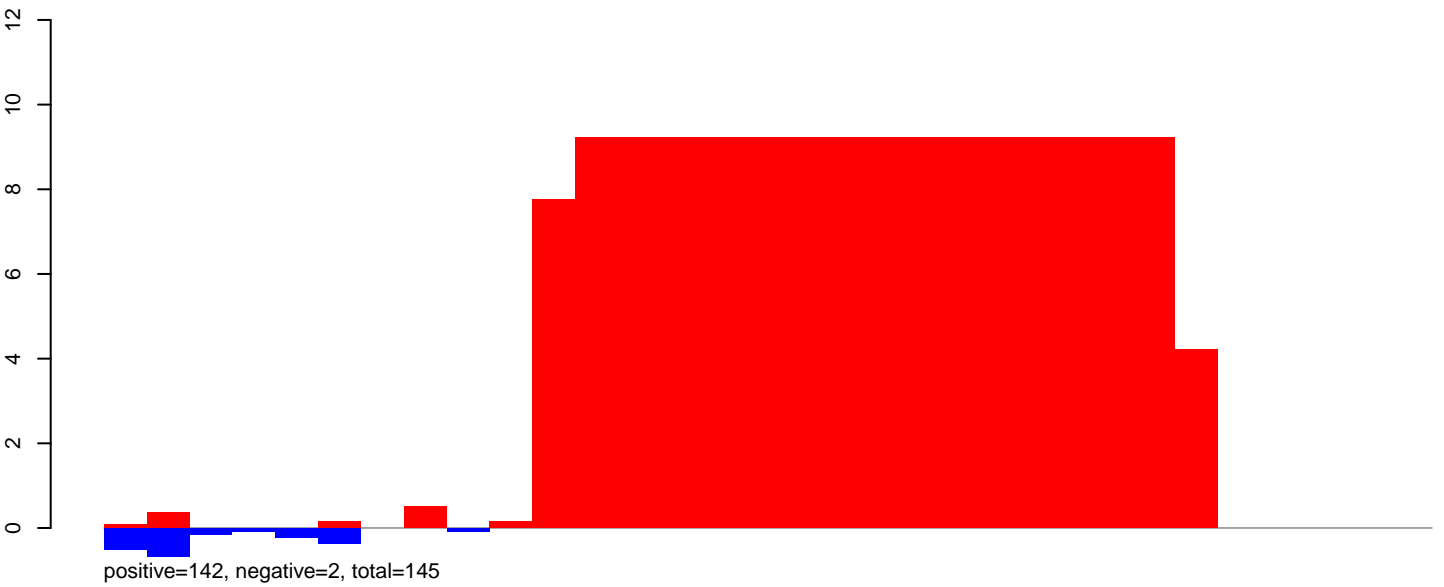
CuQuin_Testes_NL.rep



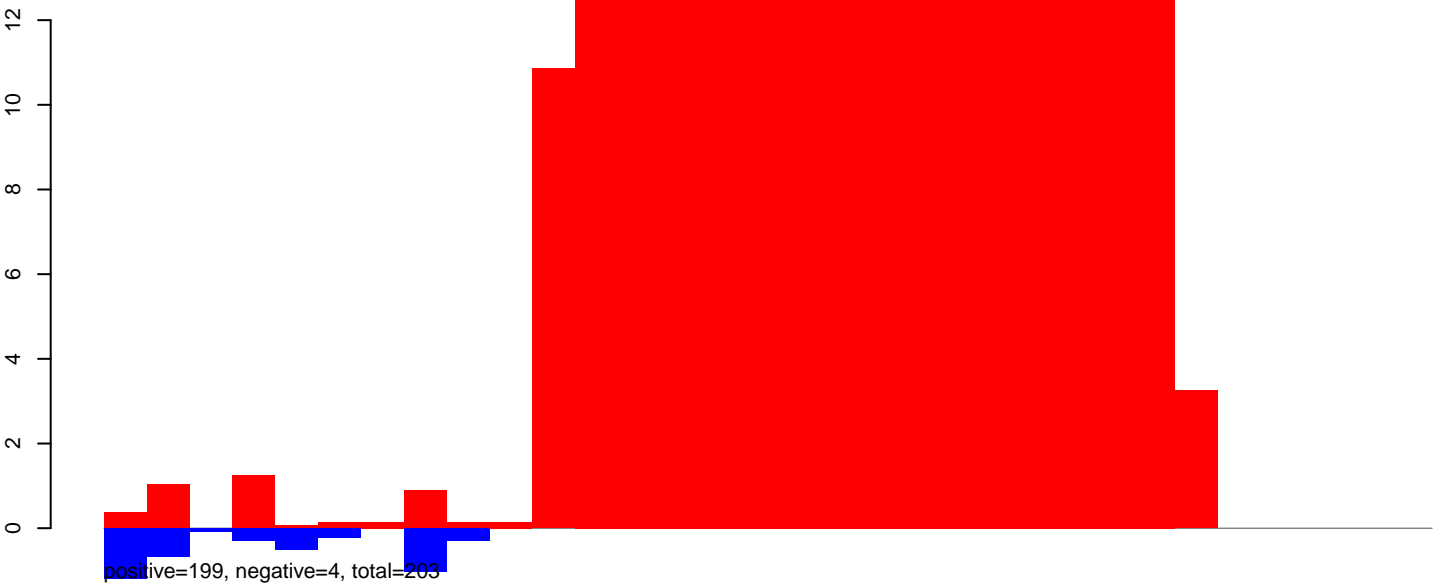
positive=11, negative=10, total=21

Window size=25, length=881, TE@rnd-3_family-287@Unknown@1284:1-881

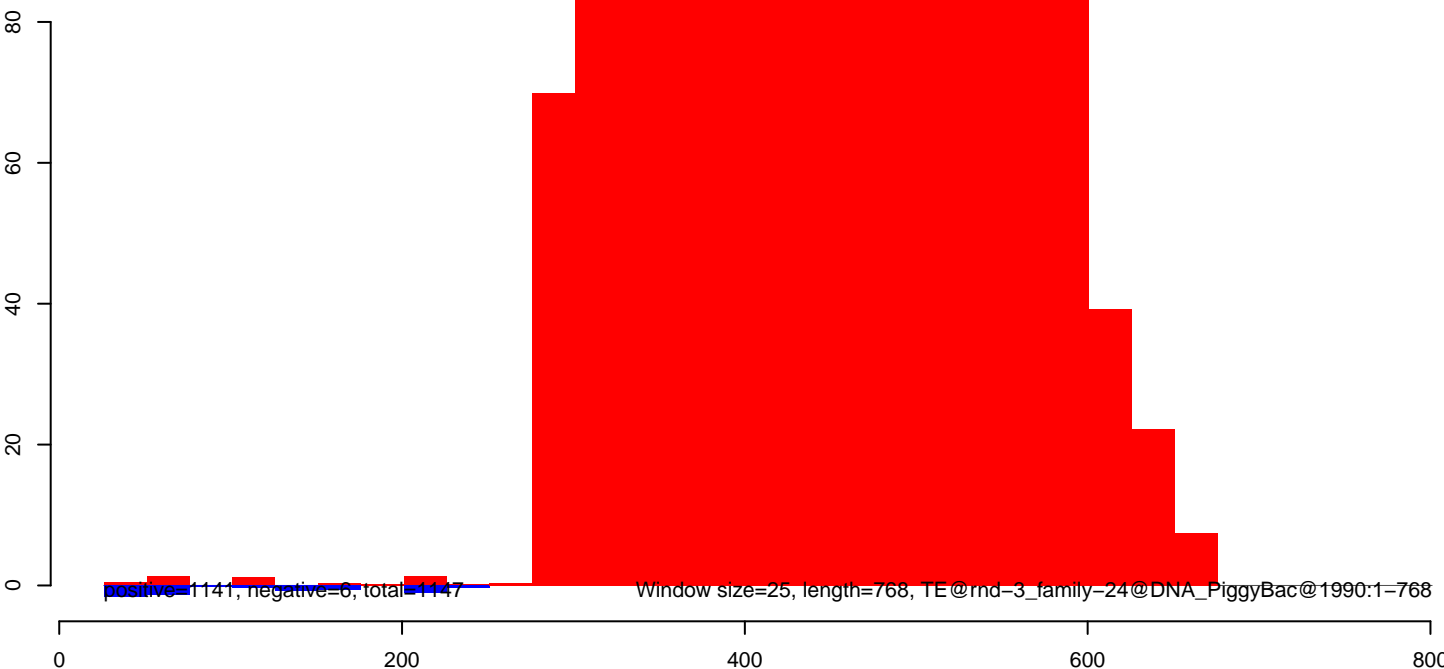
CuQuin_Testes_NL.18_23.rep



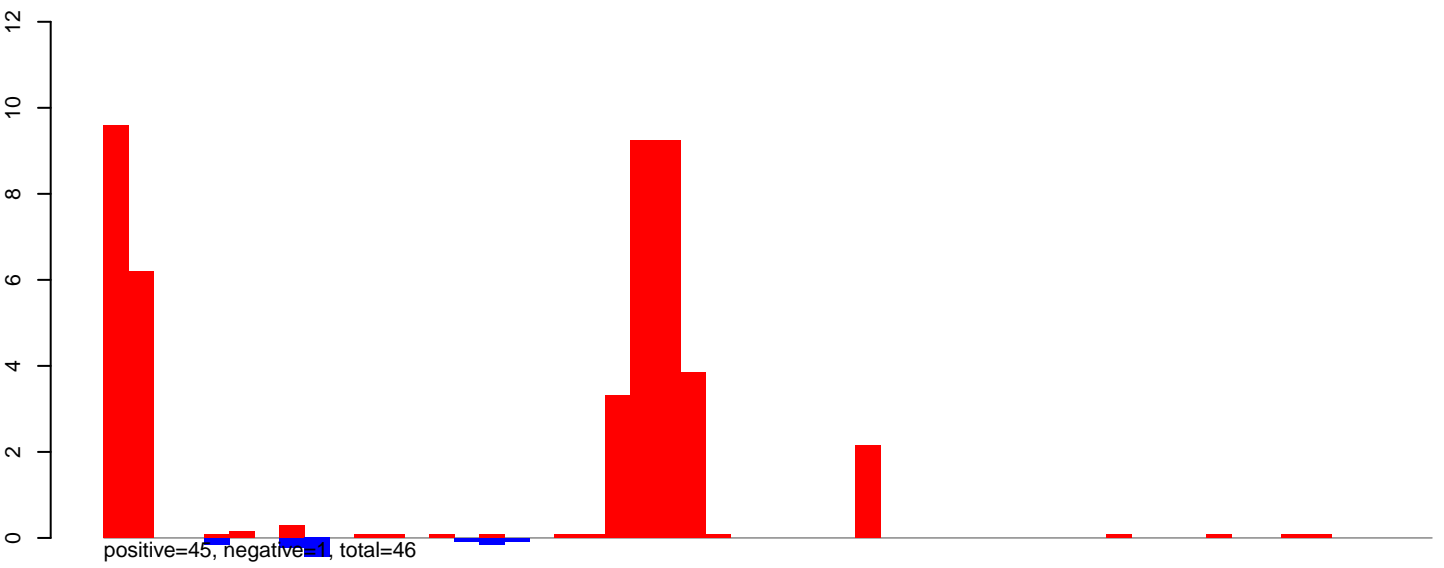
CuQuin_Testes_NL.24_35.rep



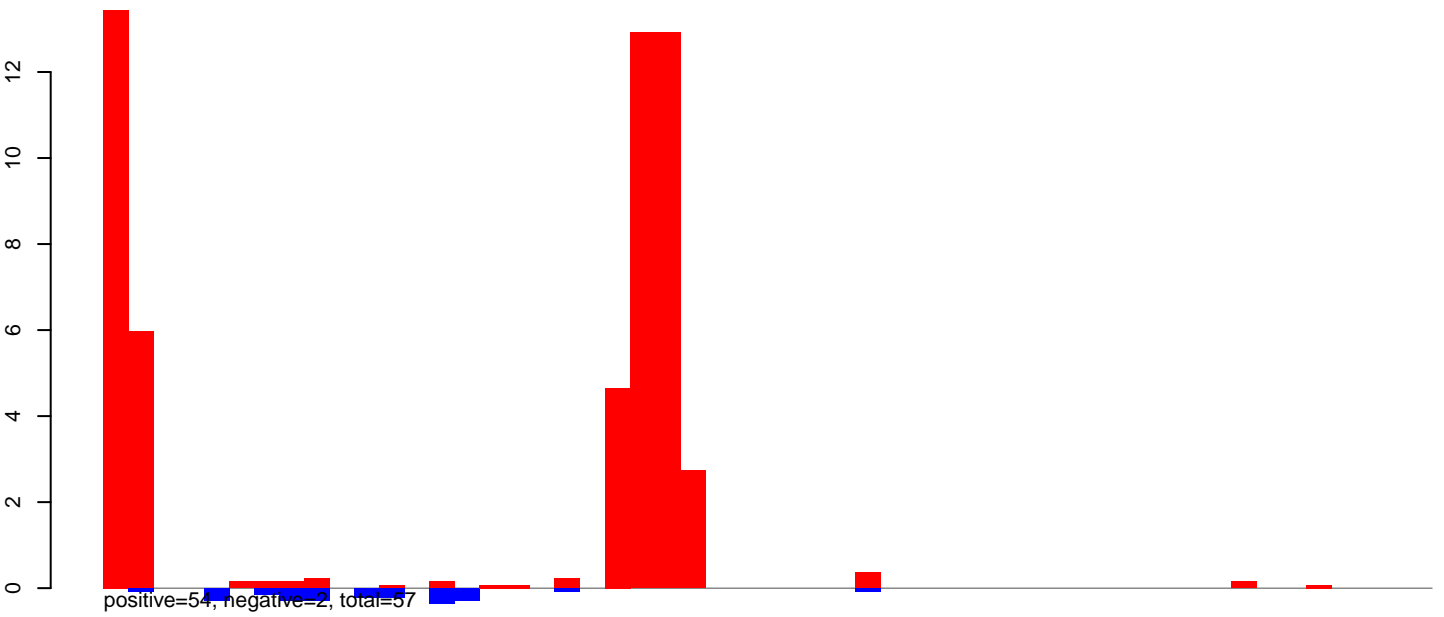
CuQuin_Testes_NL.rep



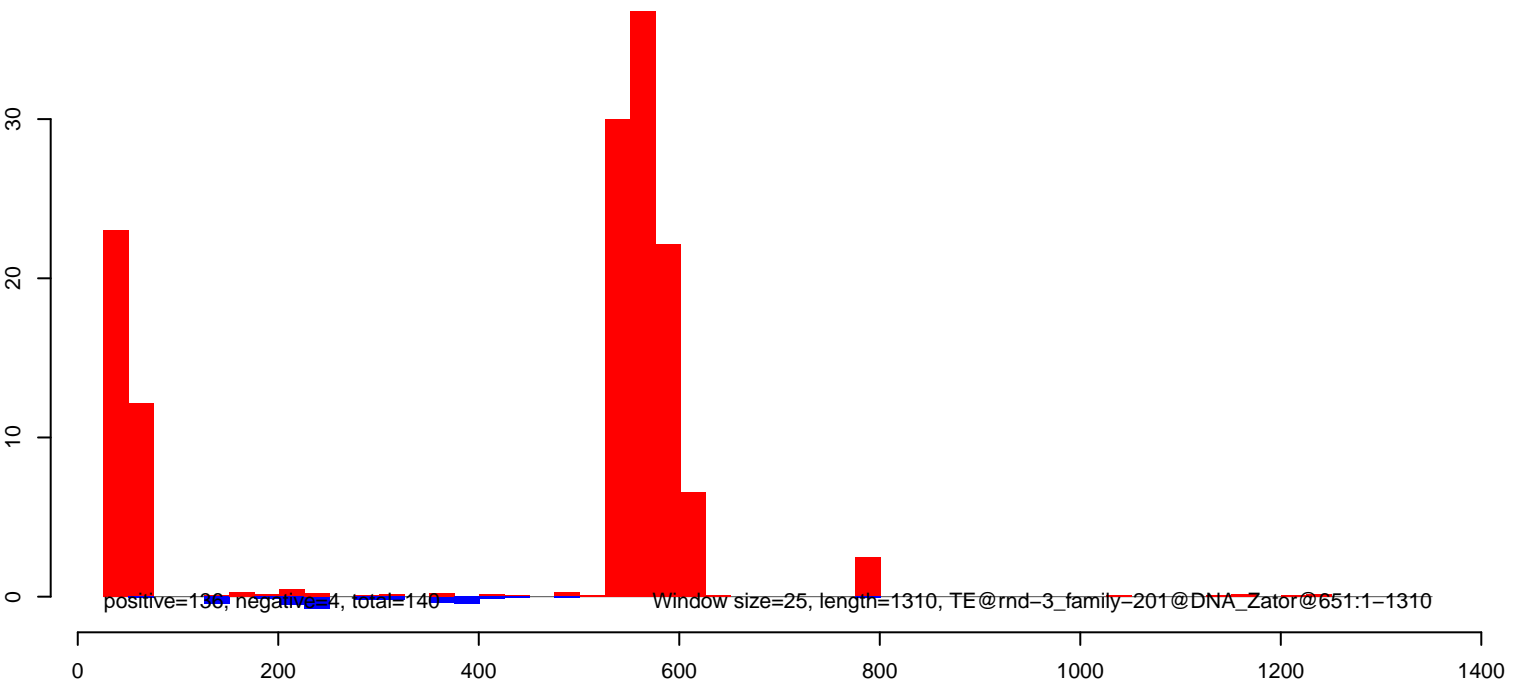
CuQuin_Testes_NL.18_23.rep



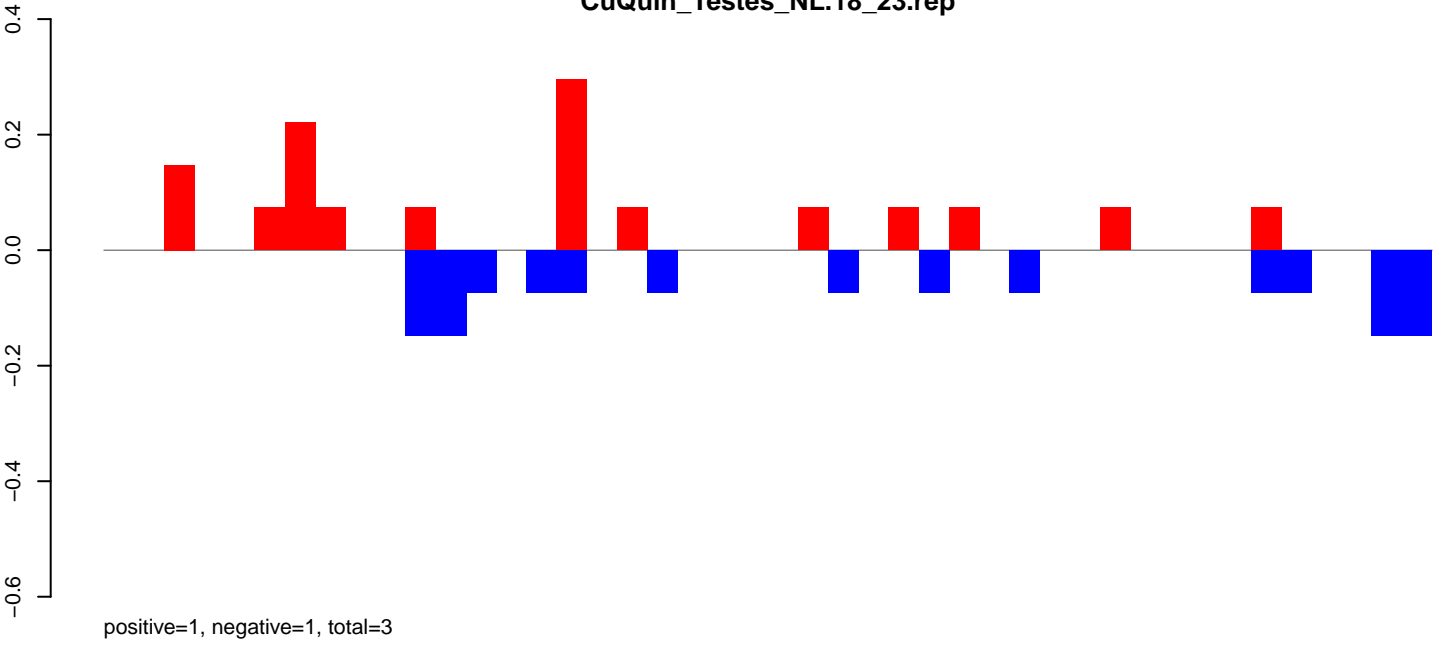
CuQuin_Testes_NL.24_35.rep



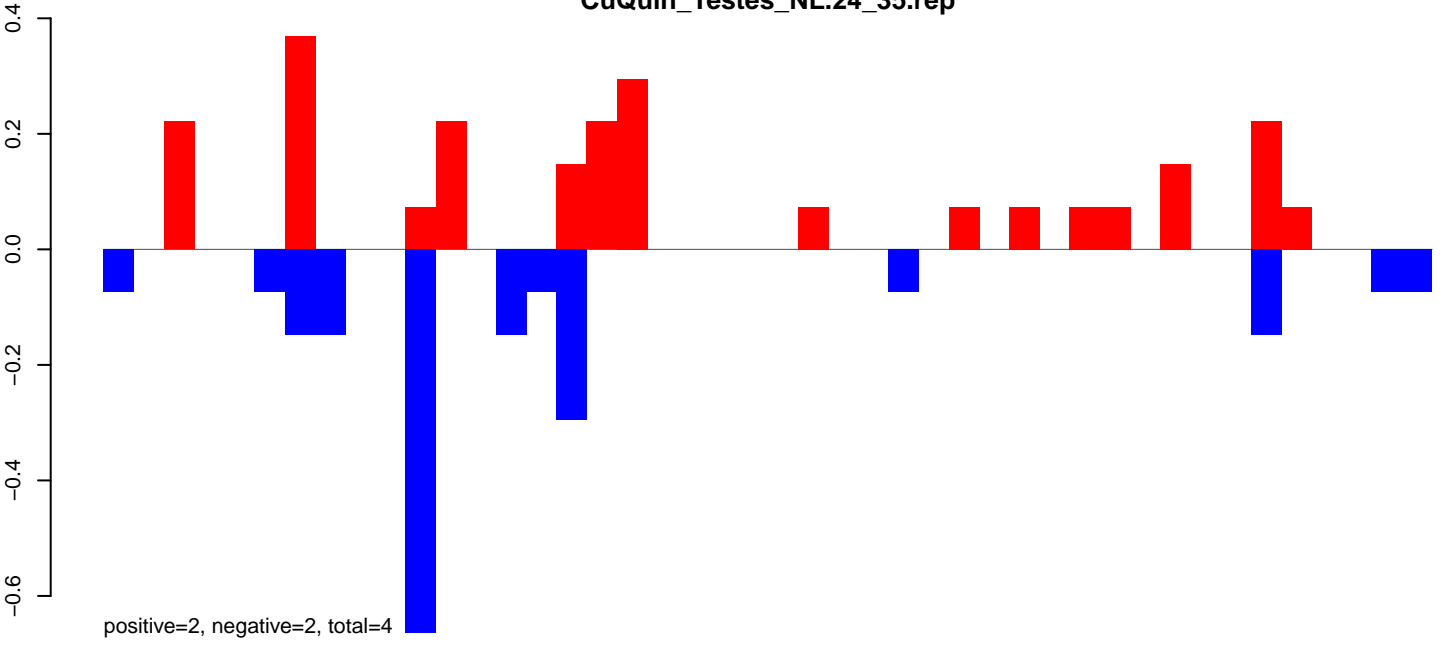
CuQuin_Testes_NL.rep



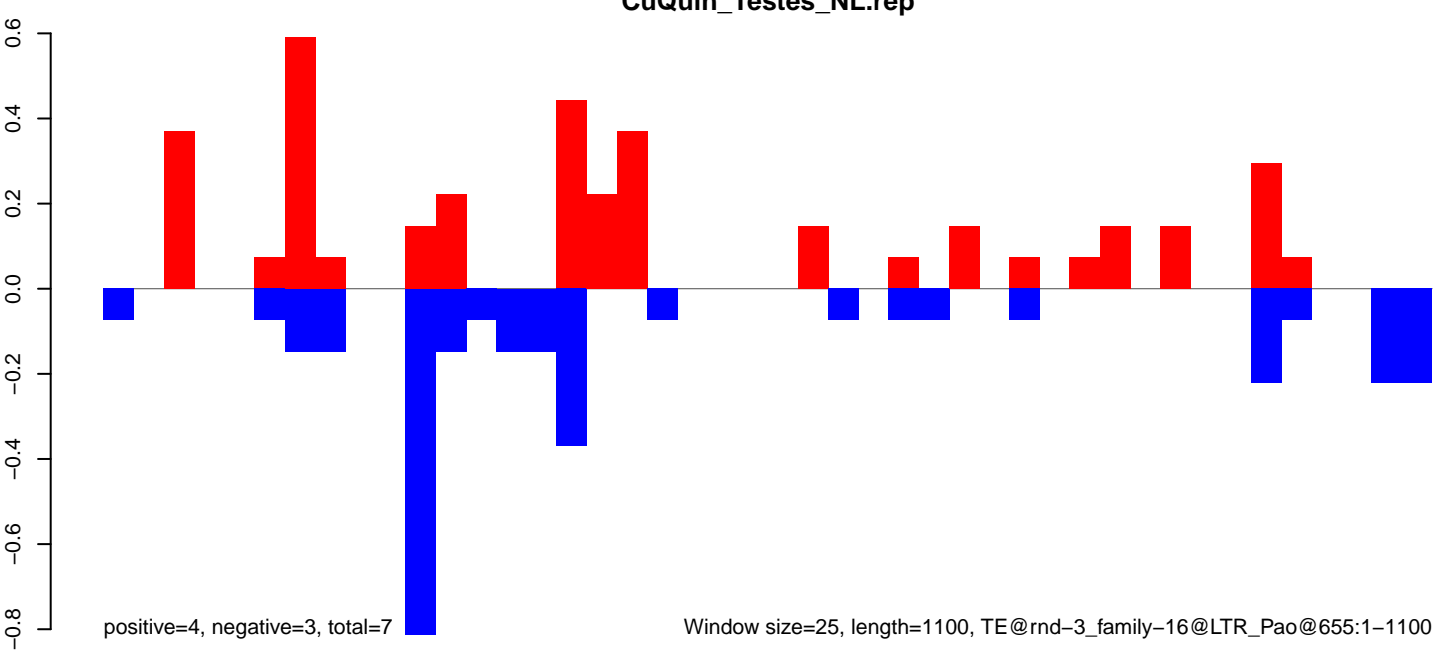
CuQuin_Testes_NL.18_23.rep



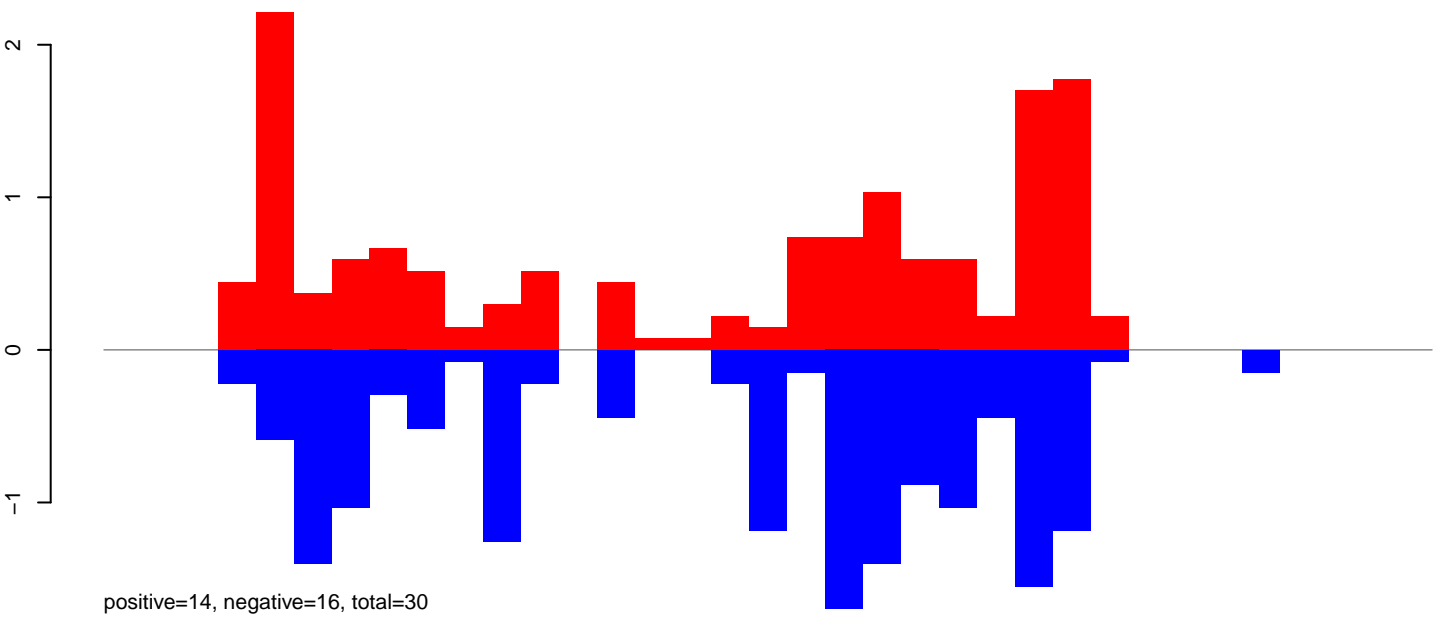
CuQuin_Testes_NL.24_35.rep



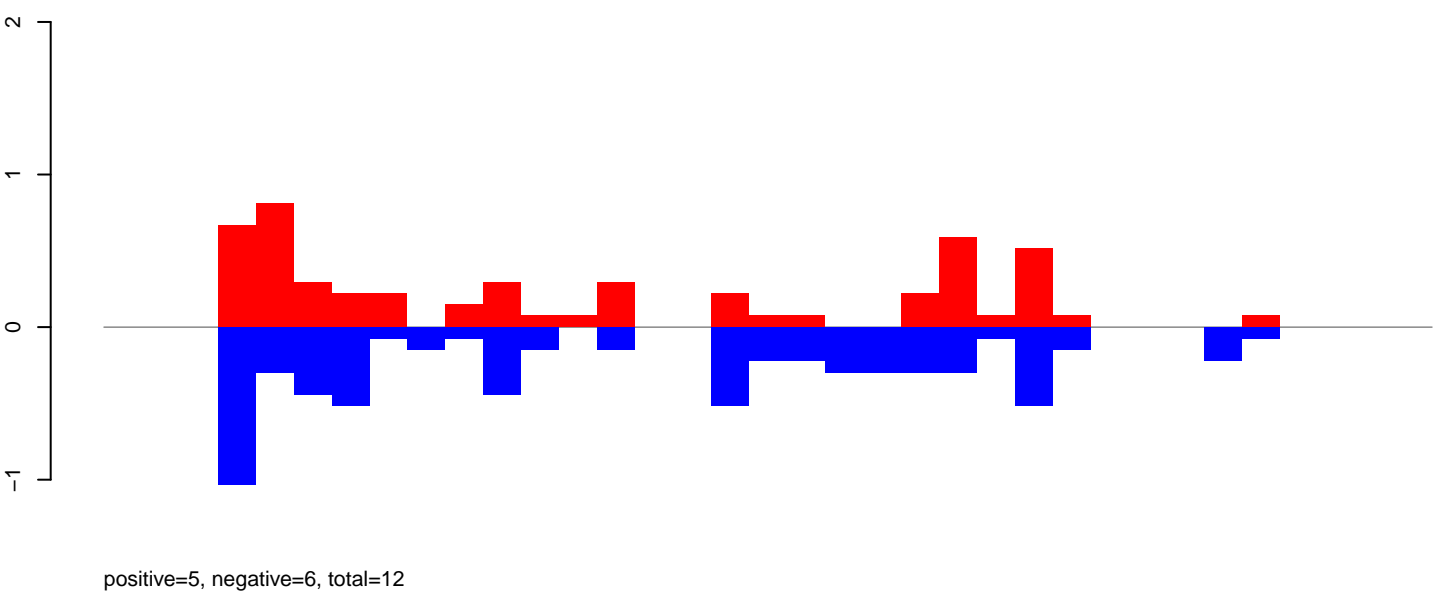
CuQuin_Testes_NL.rep



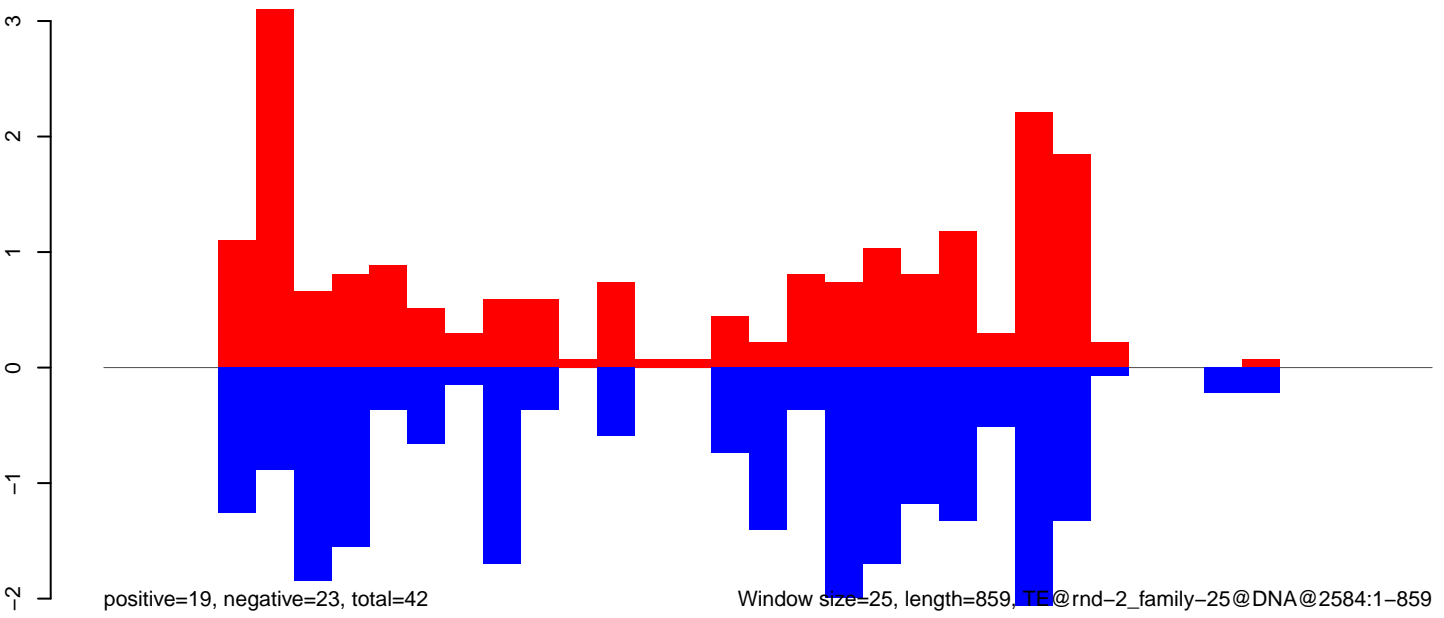
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



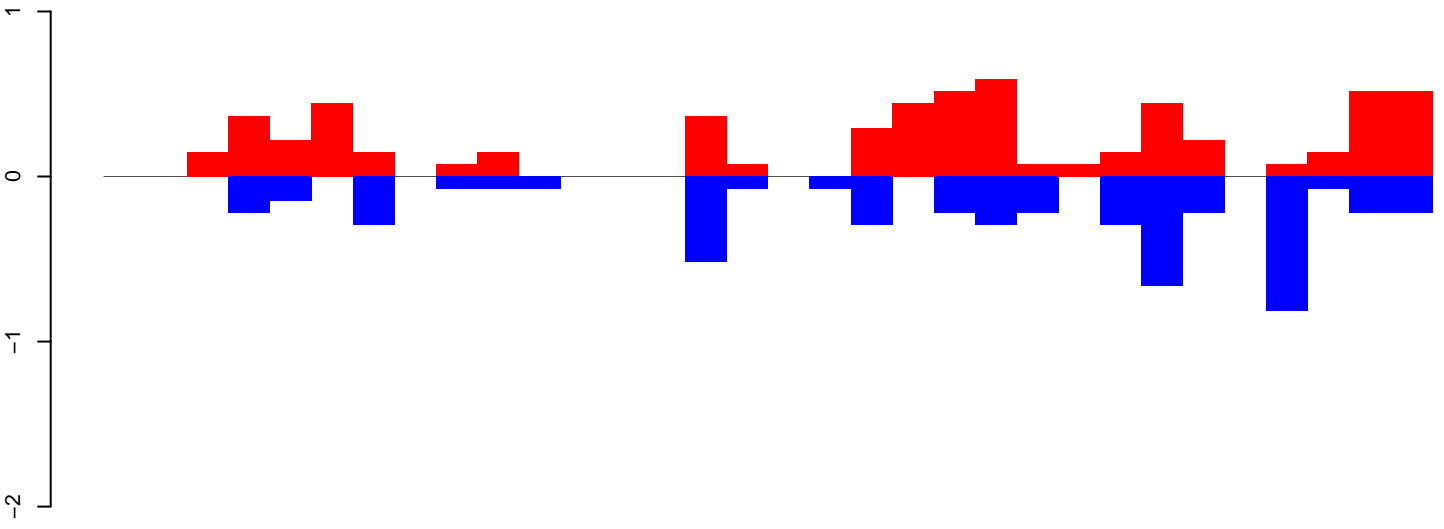
CuQuin_Testes_NL.rep



Window size=25, length=859, TE@rnd-2_family-25@DNA@2584:1-859

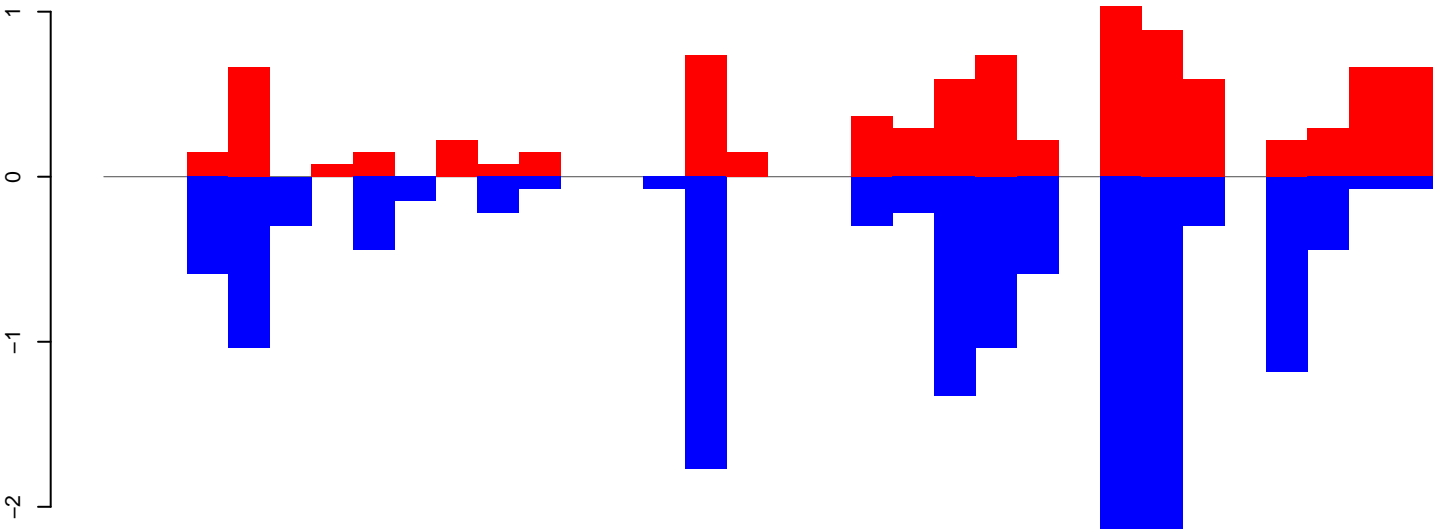
0 200 400 600 800

CuQuin_Testes_NL.18_23.rep



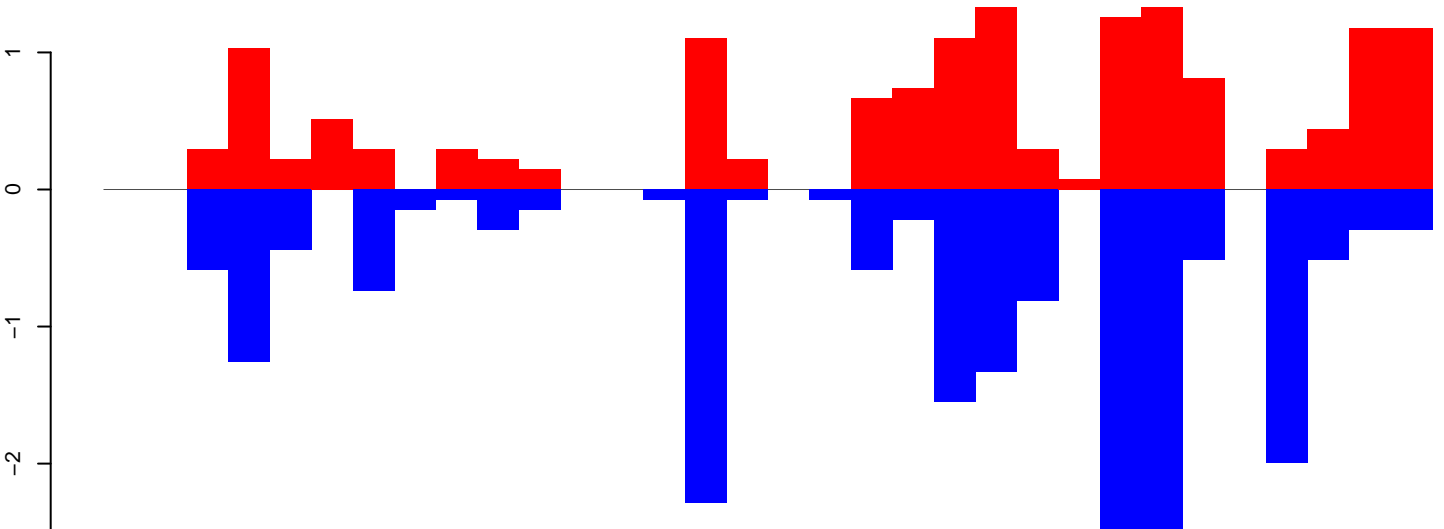
positive=6, negative=5, total=11

CuQuin_Testes_NL.24_35.rep



positive=9, negative=15, total=24

CuQuin_Testes_NL.rep

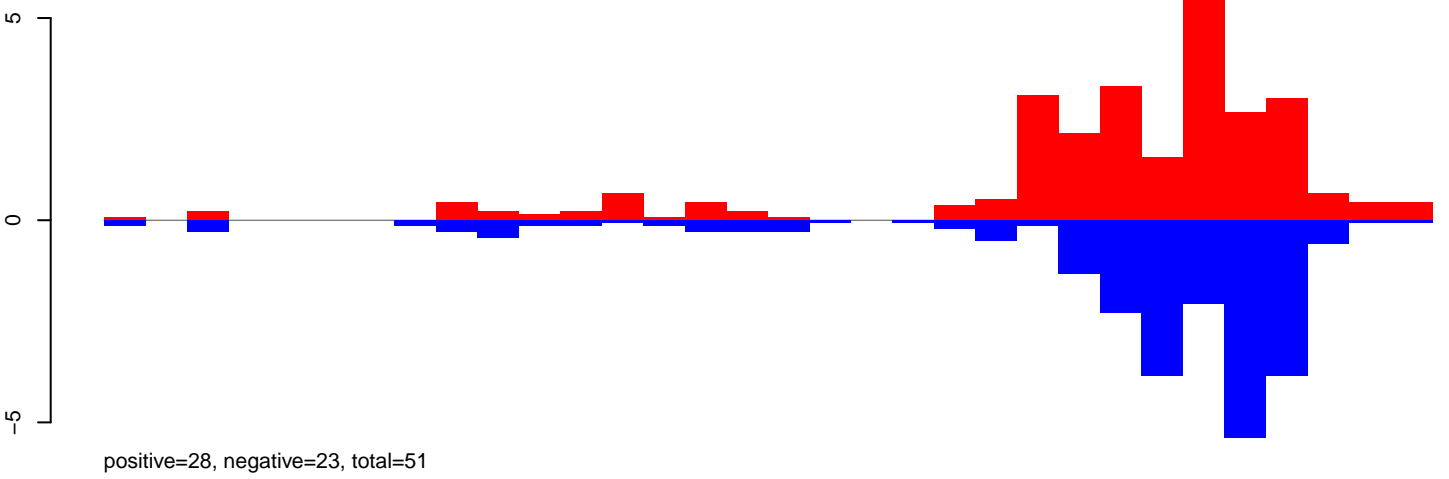


positive=15, negative=20, total=35

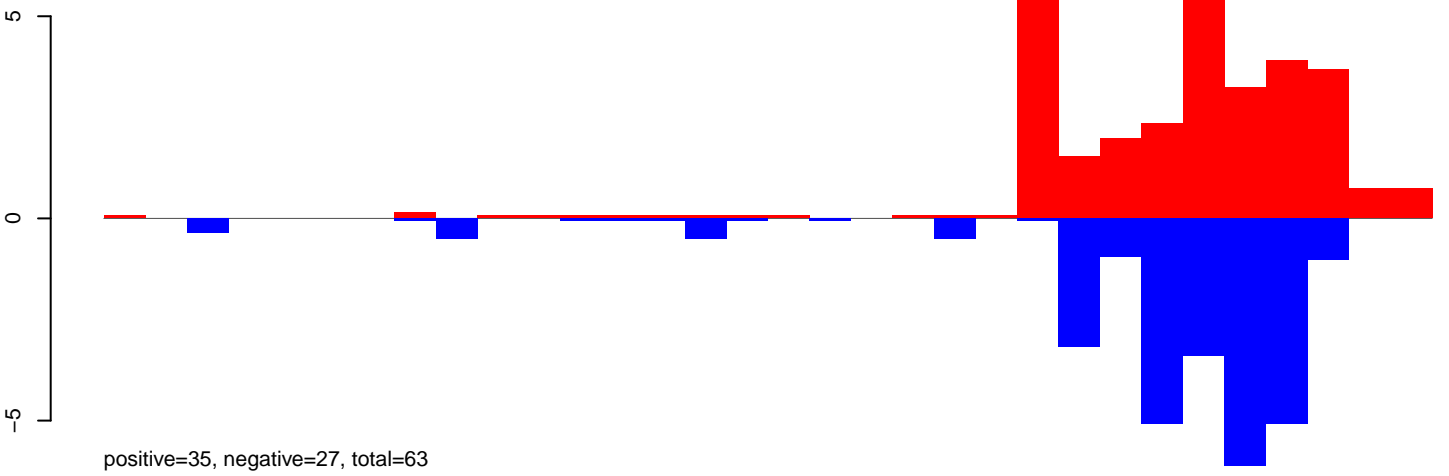
Window size=25, length=795, TE@rnd-5_family-854@DNA@2971:1-795

0 200 400 600 800

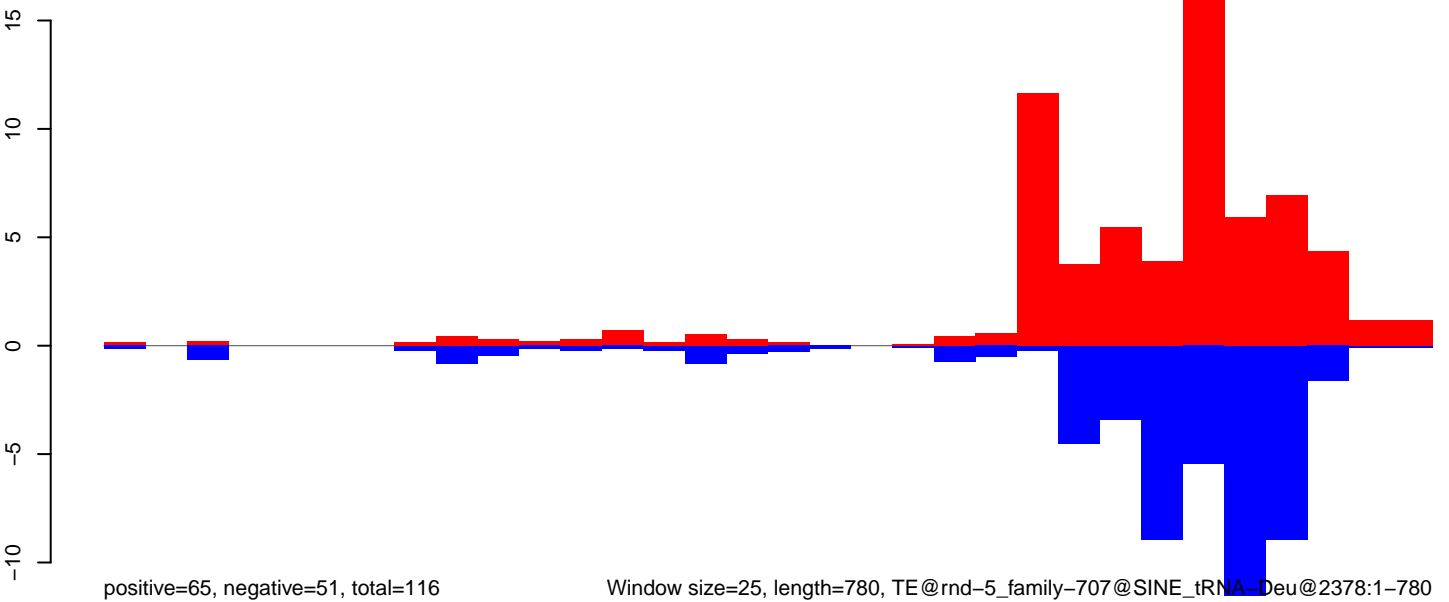
CuQuin_Testes_NL.18_23.rep



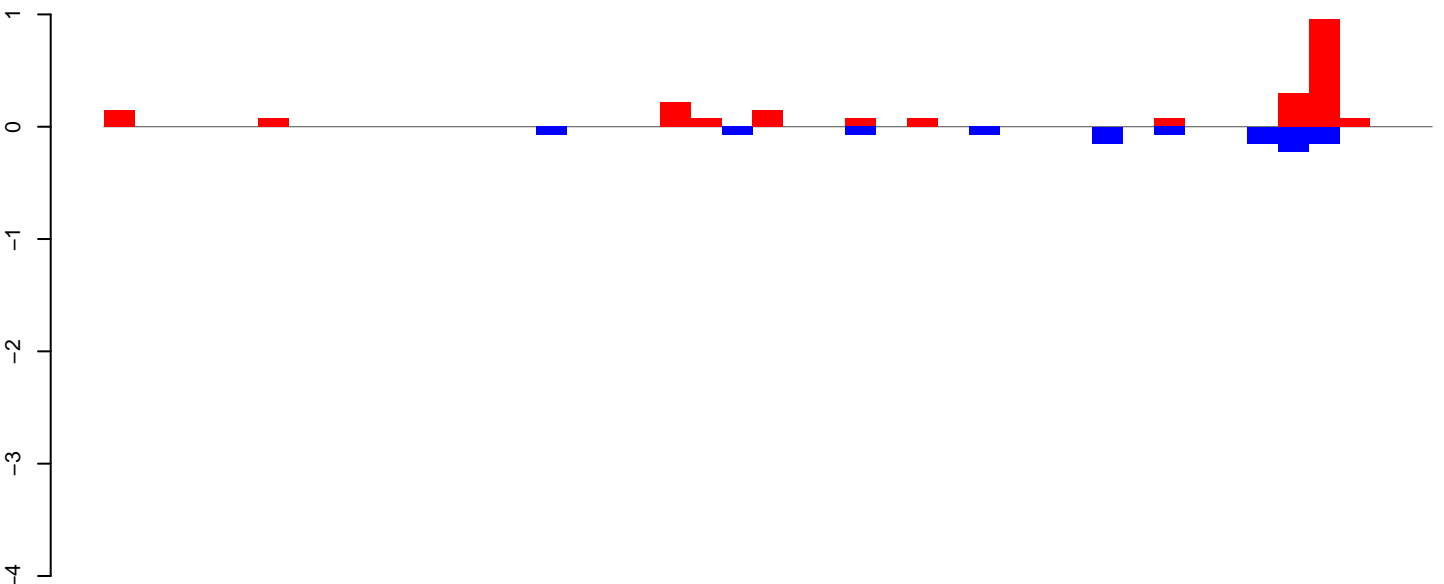
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

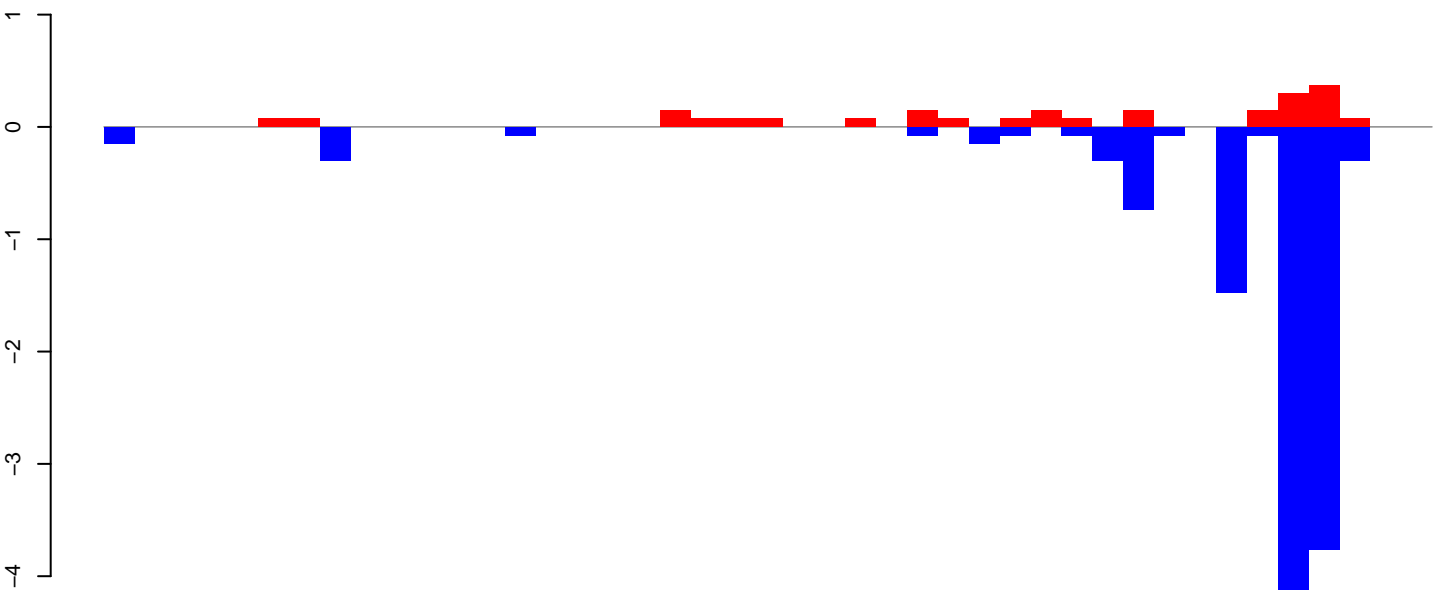


CuQuin_Testes_NL.18_23.rep



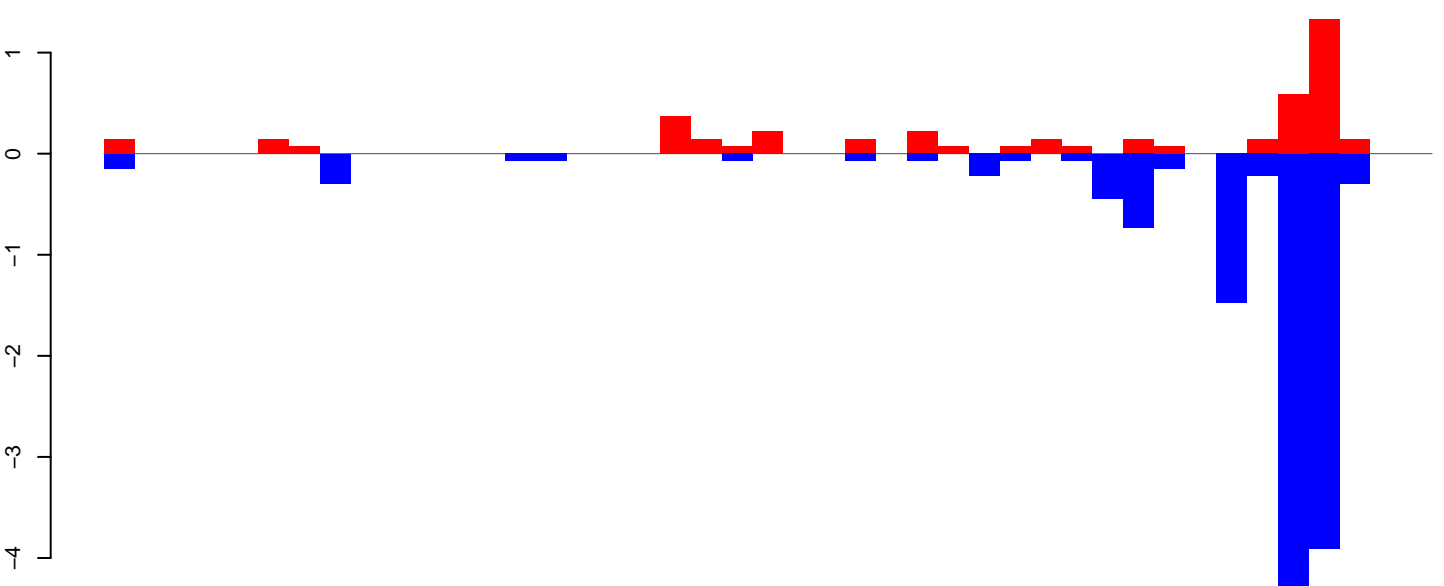
positive=2, negative=1, total=3

CuQuin_Testes_NL.24_35.rep



positive=2, negative=12, total=14

CuQuin_Testes_NL.rep

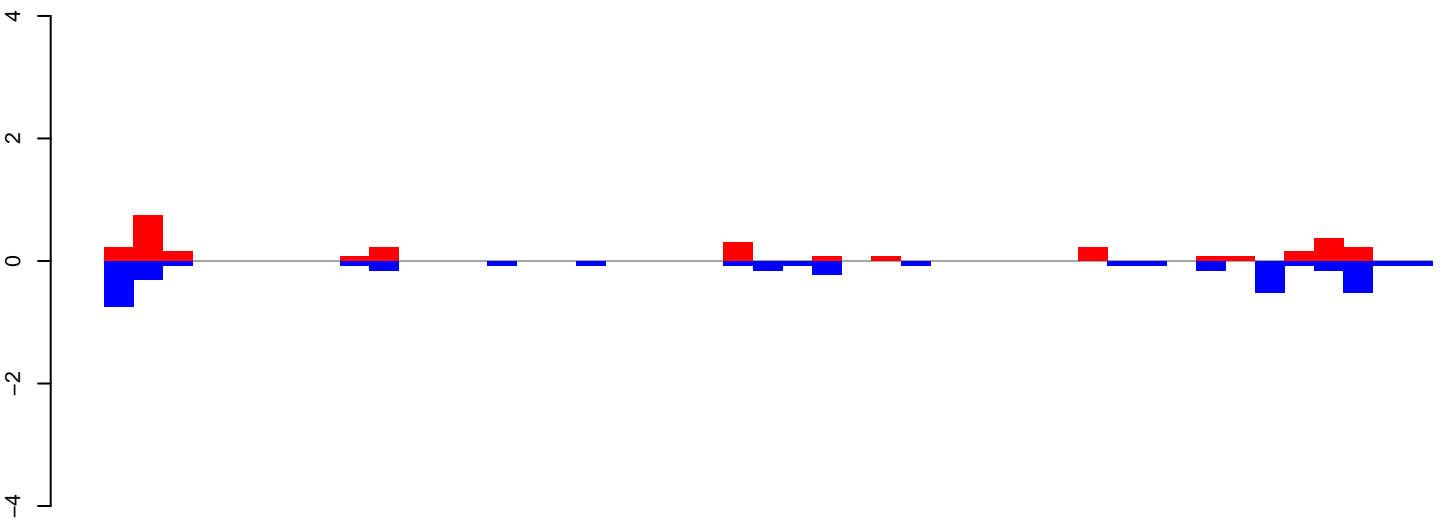


positive=4, negative=13, total=17

Window size=25, length=1067, TE@rnd-5_family-3503@Unknown@3050:1-1067

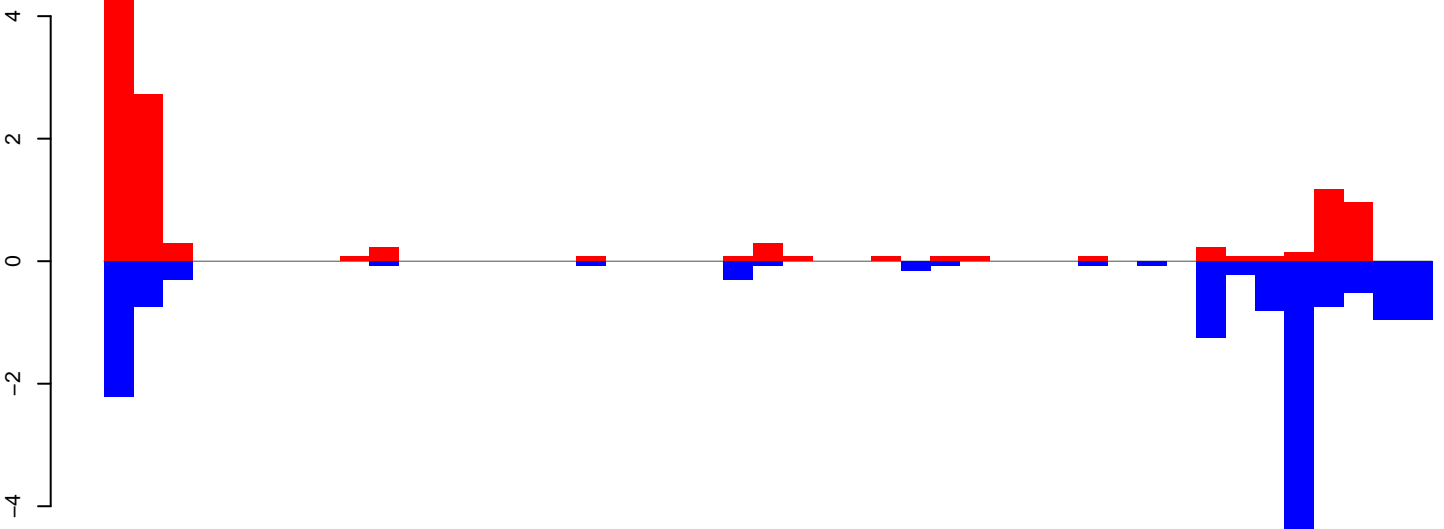
0 200 400 600 800 1000

CuQuin_Testes_NL.18_23.rep



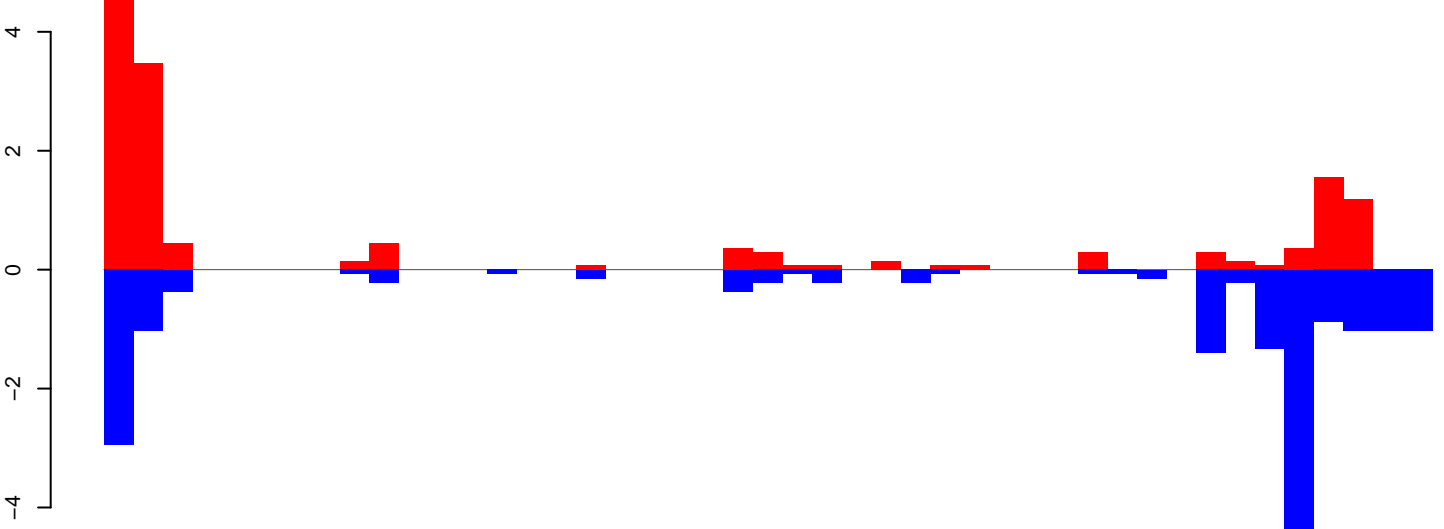
positive=3, negative=4, total=7

CuQuin_Testes_NL.24_35.rep



positive=12, negative=14, total=26

CuQuin_Testes_NL.rep

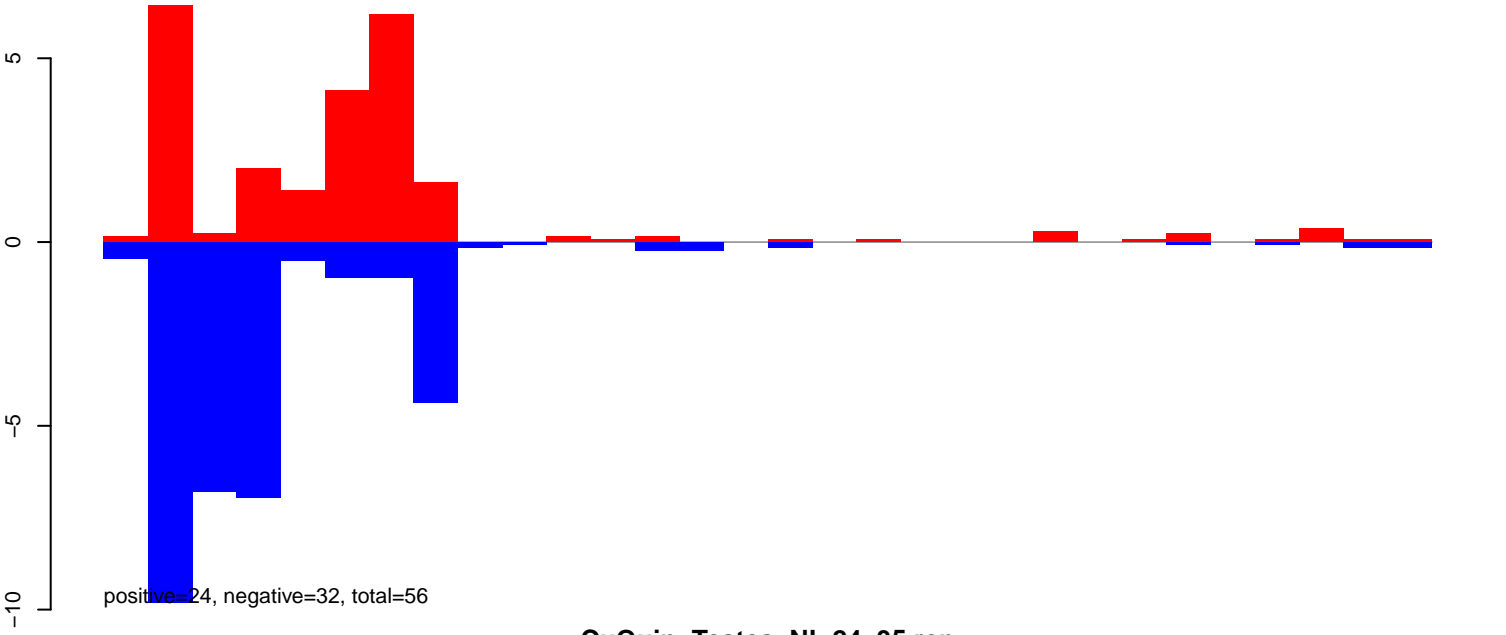


positive=15, negative=18, total=33

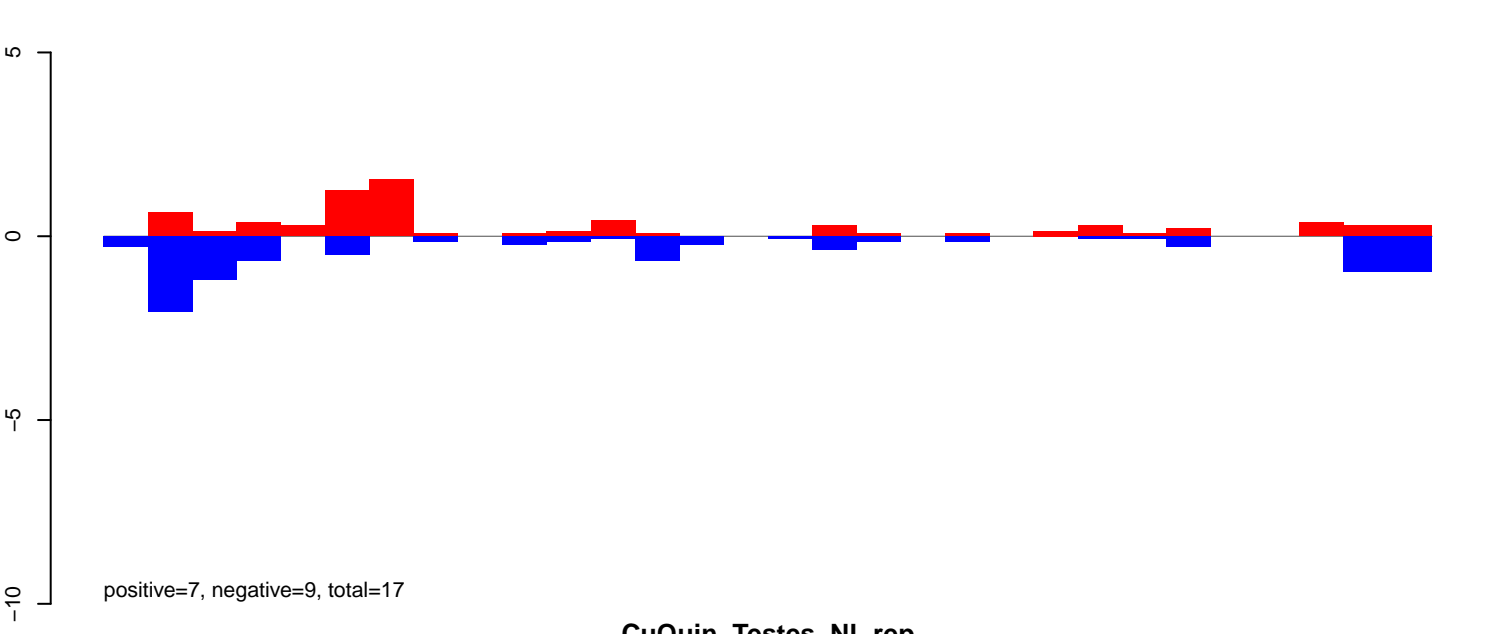
Window size=25, length=1110, TE@rnd-5_family-2842@DNA_CMC-Transib@1108:1-1110

0 200 400 600 800 1000

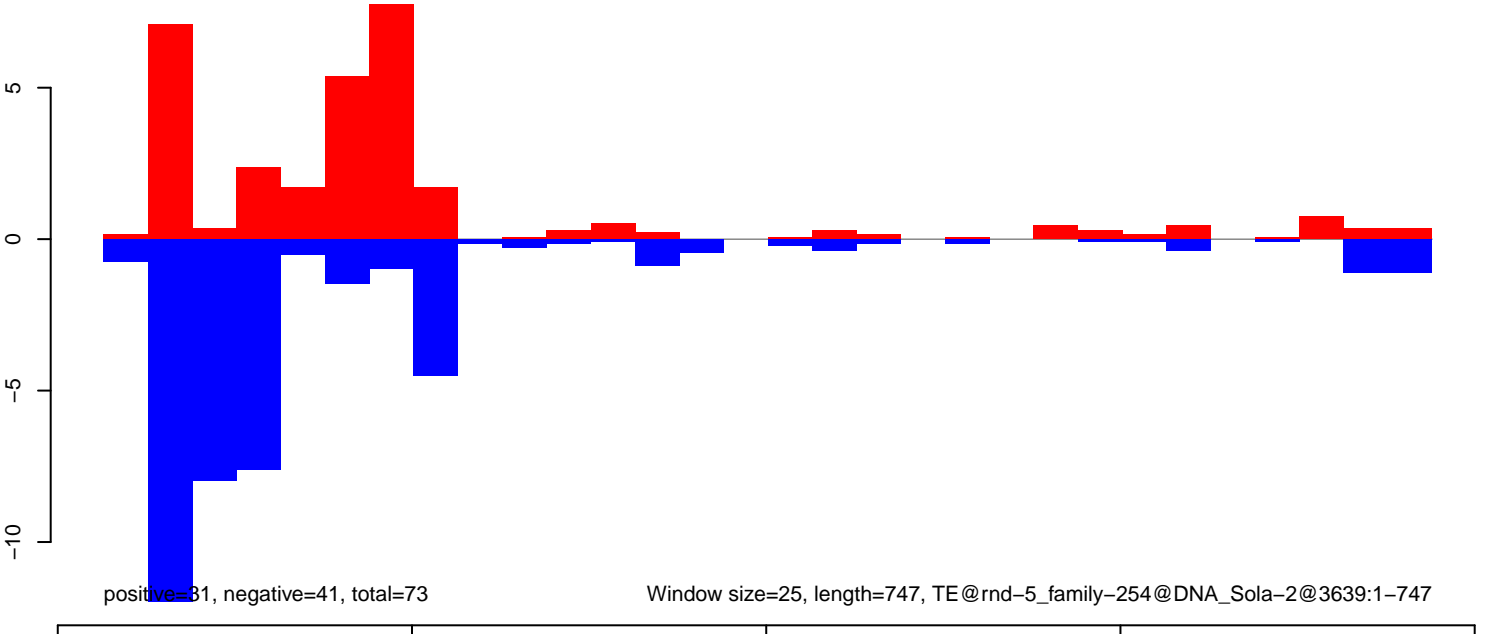
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

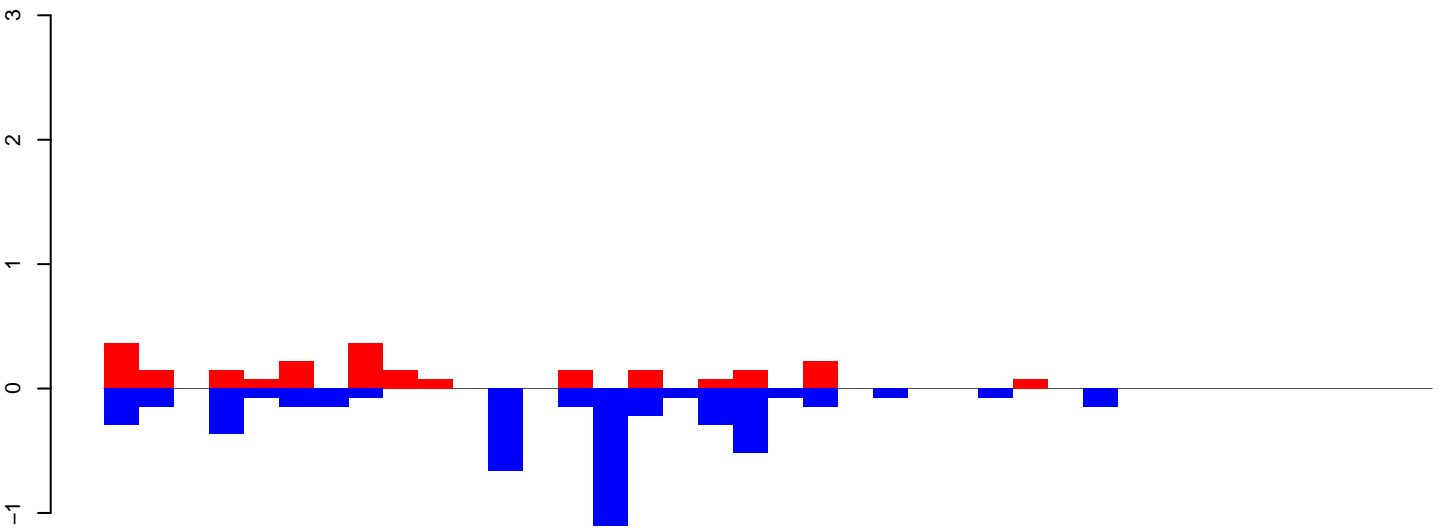


CuQuin_Testes_NL.rep



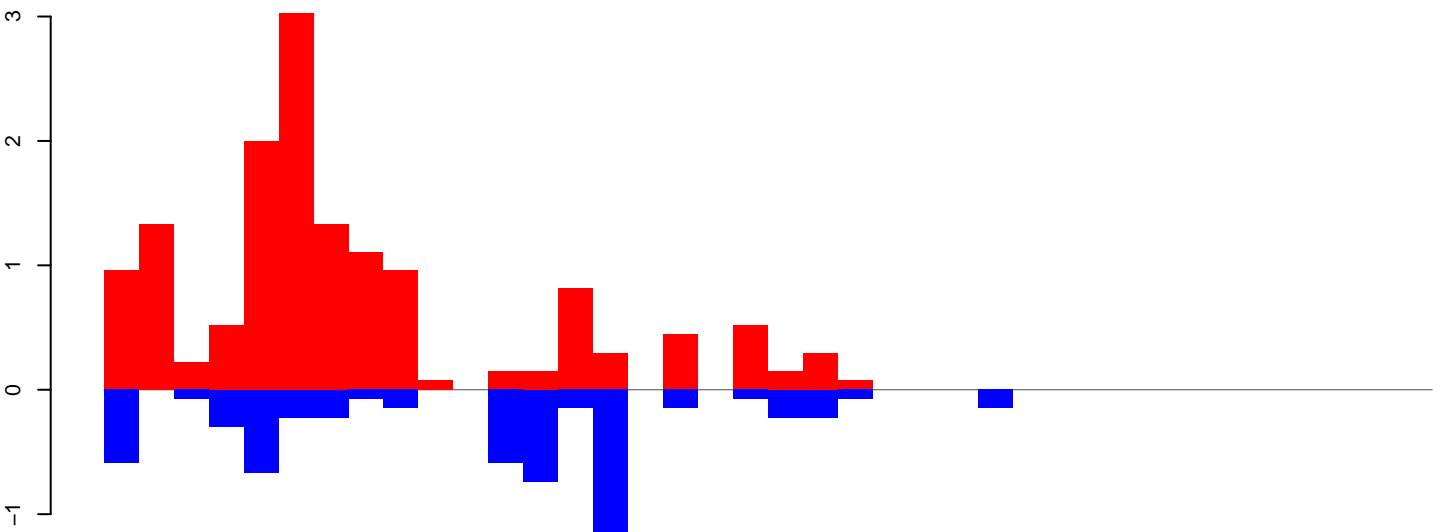
Window size=25, length=747, TE@rnd-5_family-254@DNA_Sola-2@3639:1-747

CuQuin_Testes_NL.18_23.rep



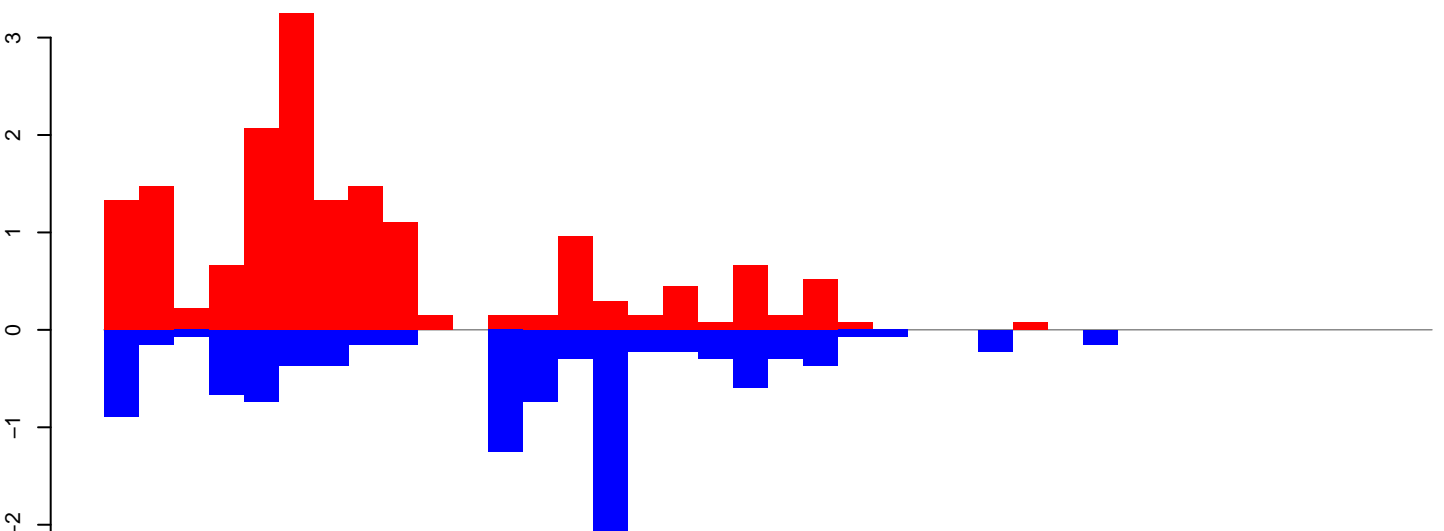
positive=2, negative=5, total=7

CuQuin_Testes_NL.24_35.rep



positive=14, negative=6, total=21

CuQuin_Testes_NL.rep

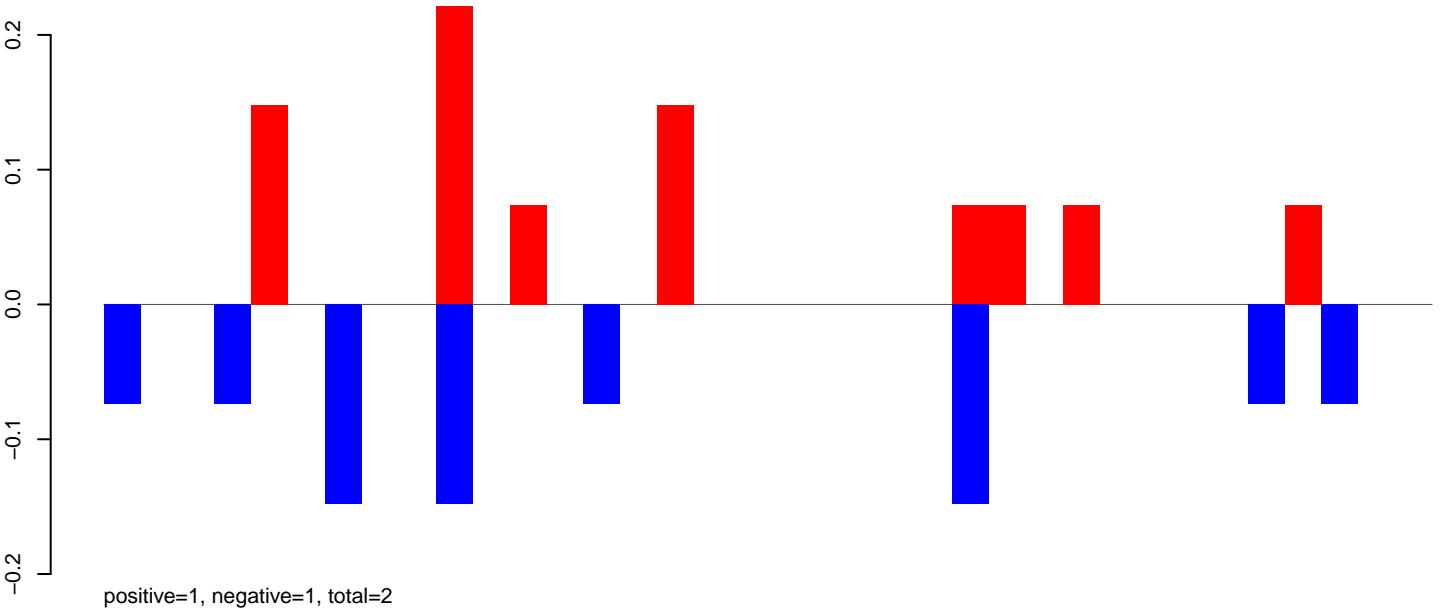


positive=17, negative=11, total=28

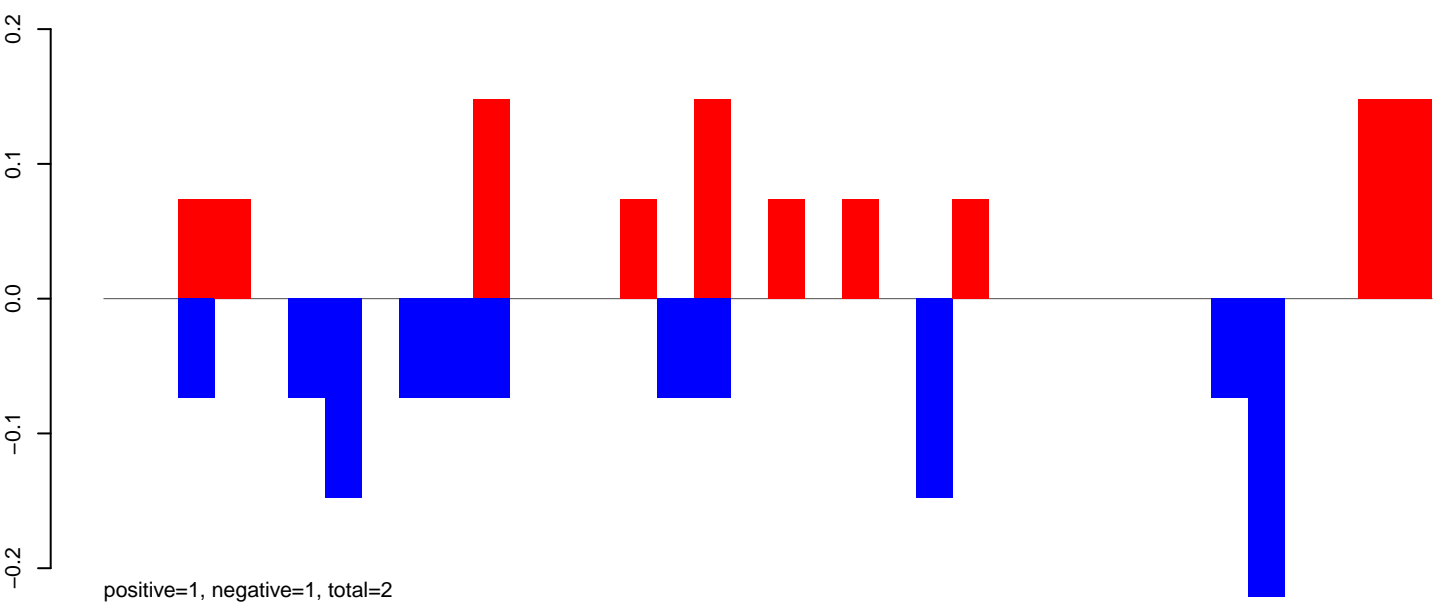
Window size=25, length=946, TE@rnd-4_family-692@LTR_Pao@3429:1-946

0 200 400 600 800 1000

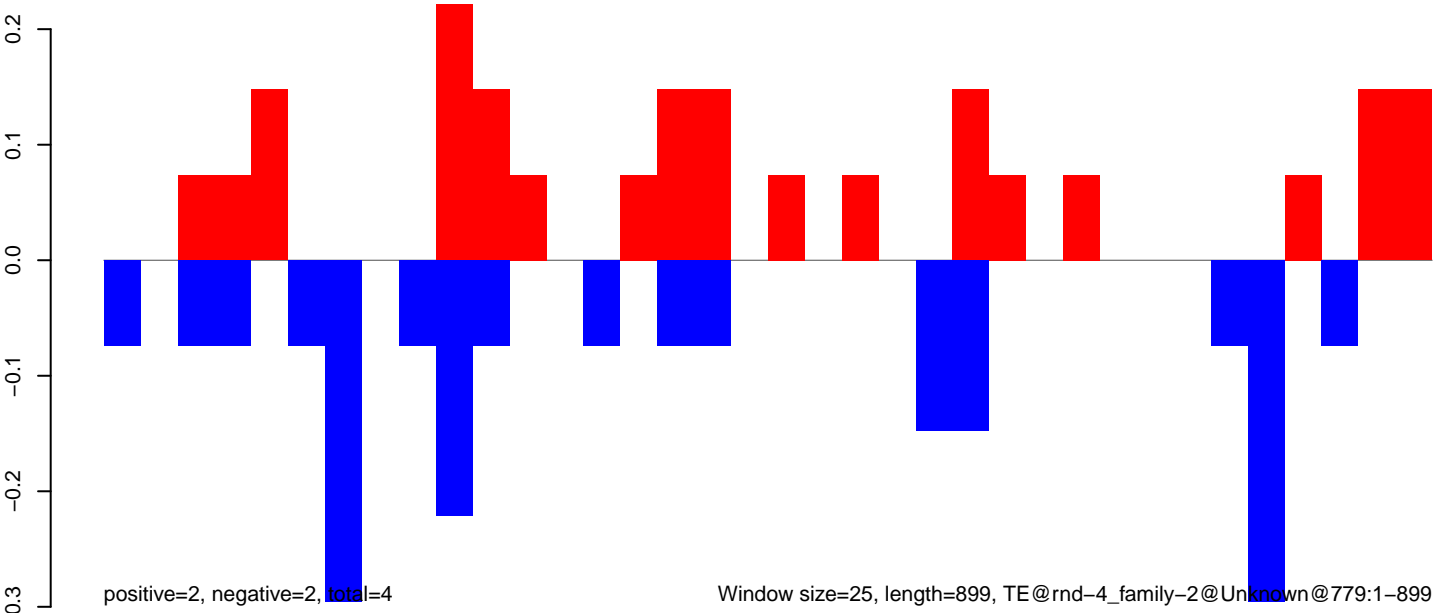
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

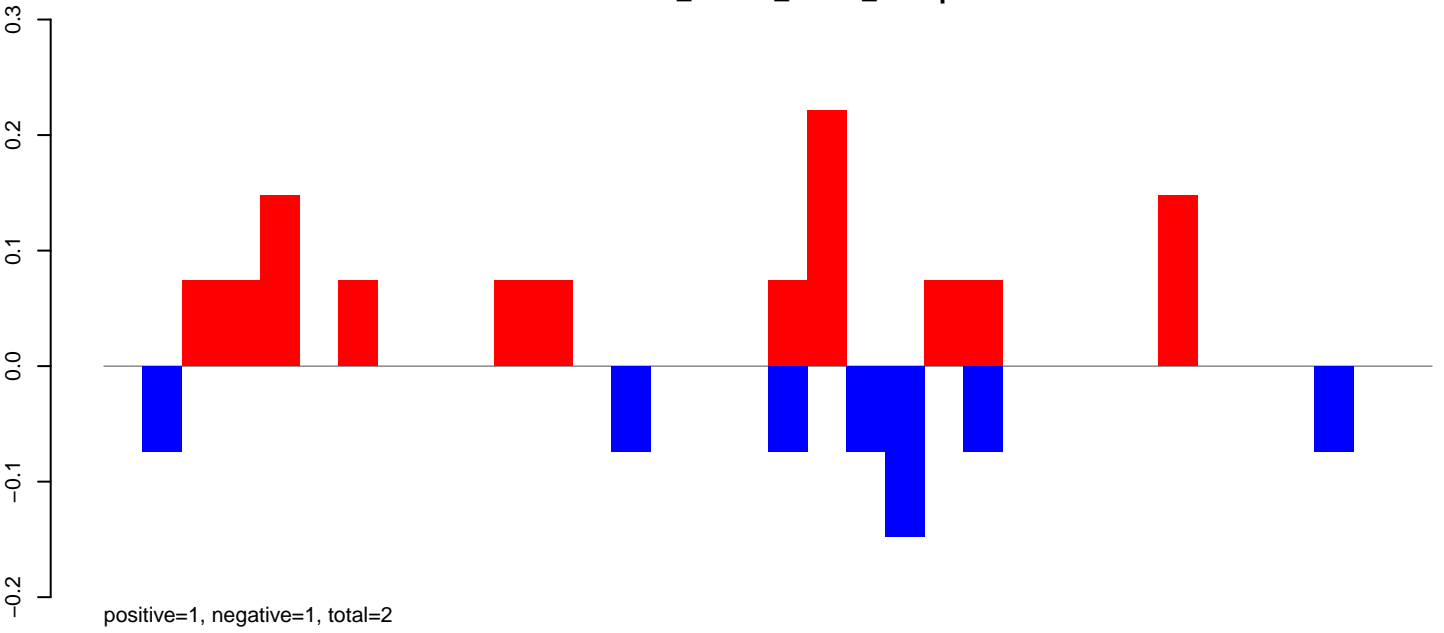


CuQuin_Testes_NL.rep

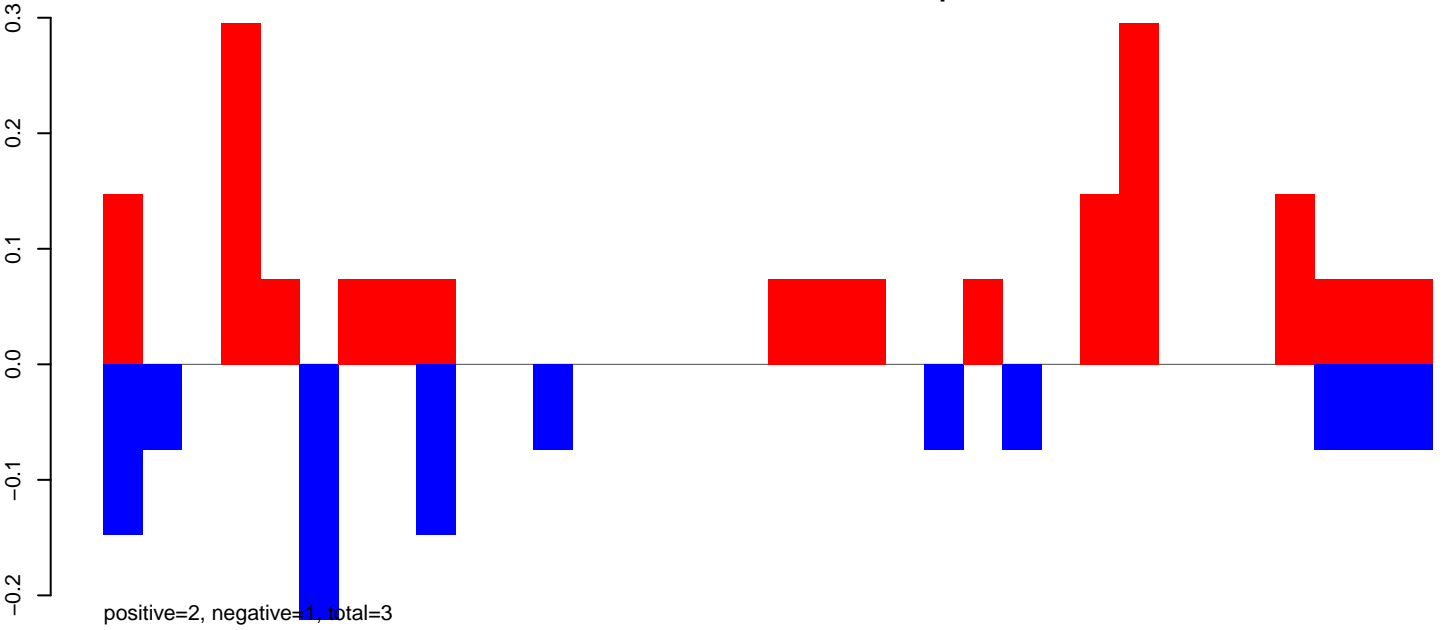


Window size=25, length=899, TE@rnd-4_family-2@Unknown@779:1-899

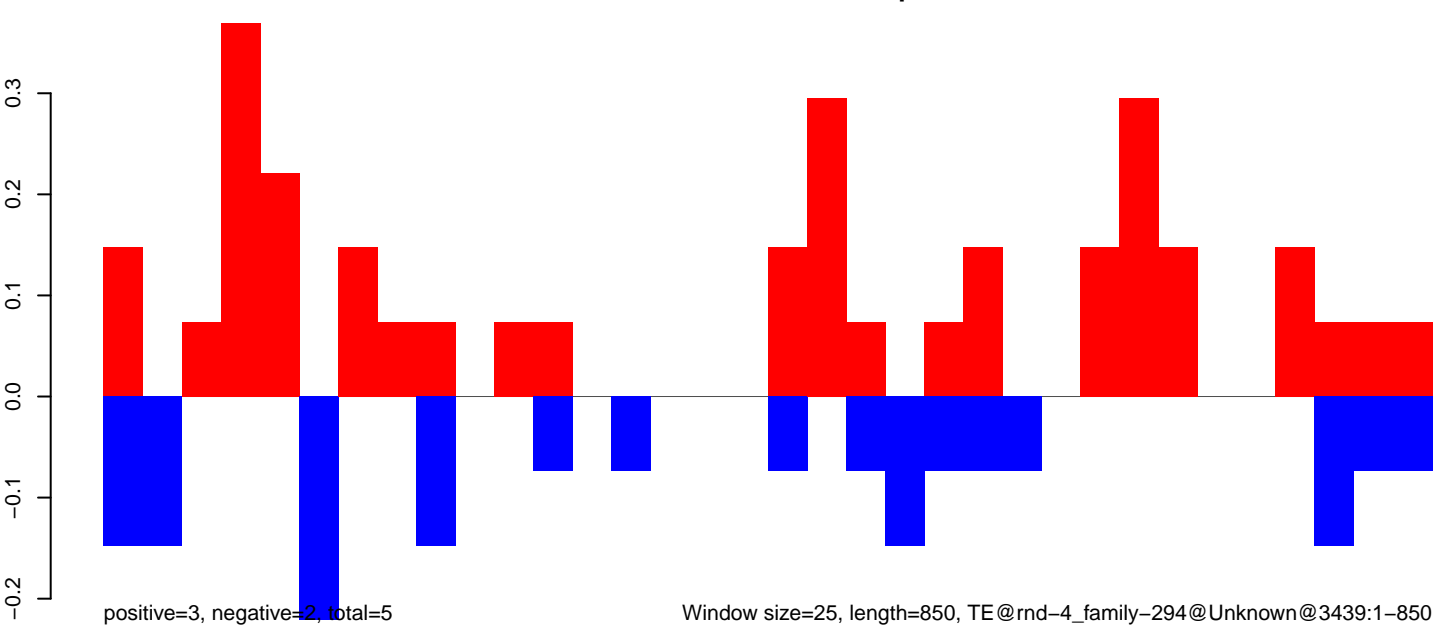
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

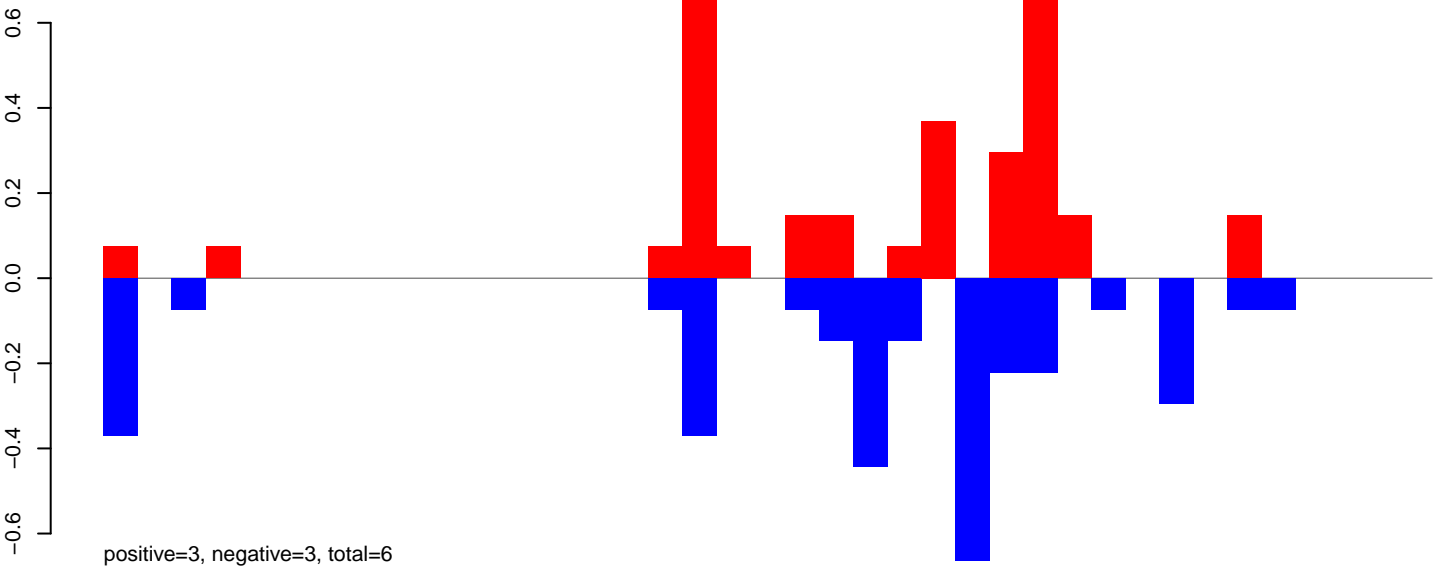


CuQuin_Testes_NL.rep

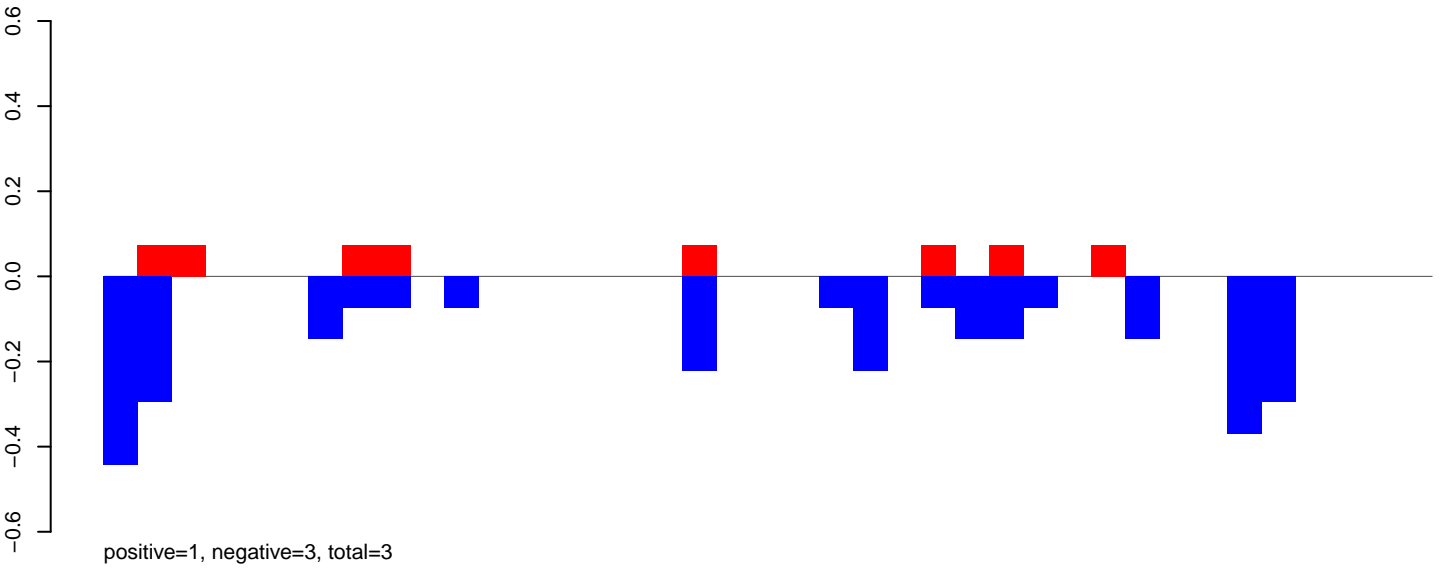


Window size=25, length=850, TE@rnd-4_family-294@Unknown@3439:1-850

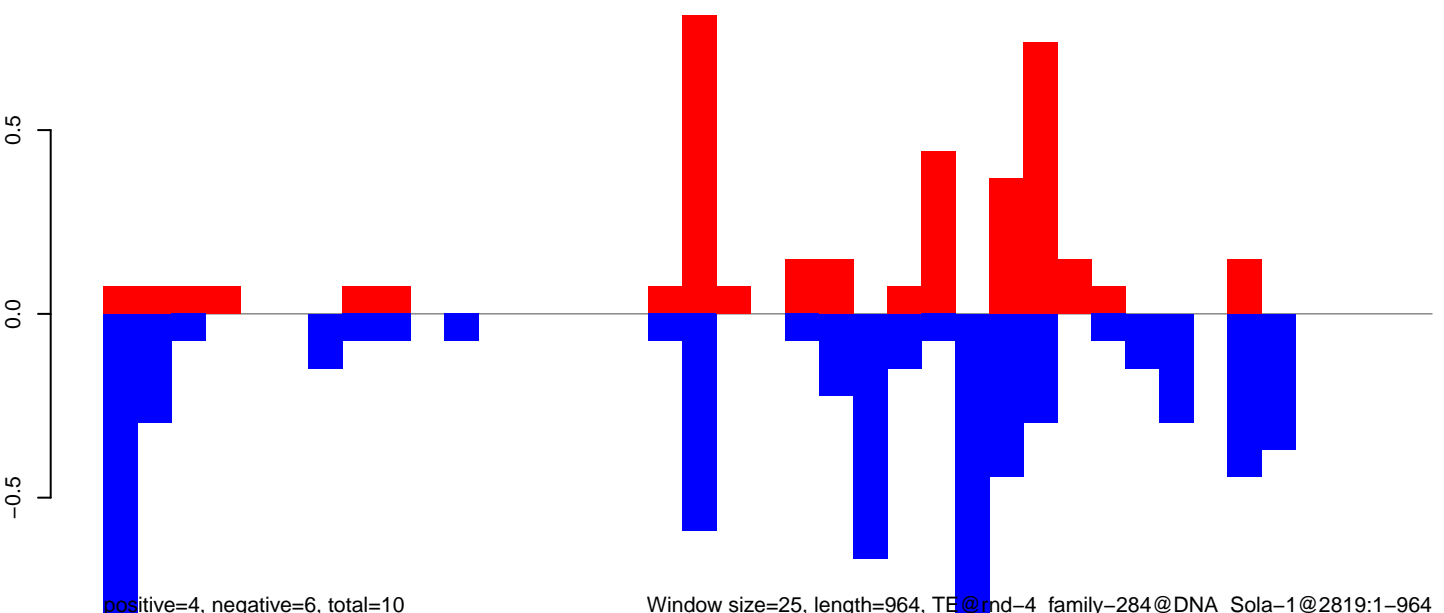
CuQuin_Testes_NL.18_23.rep



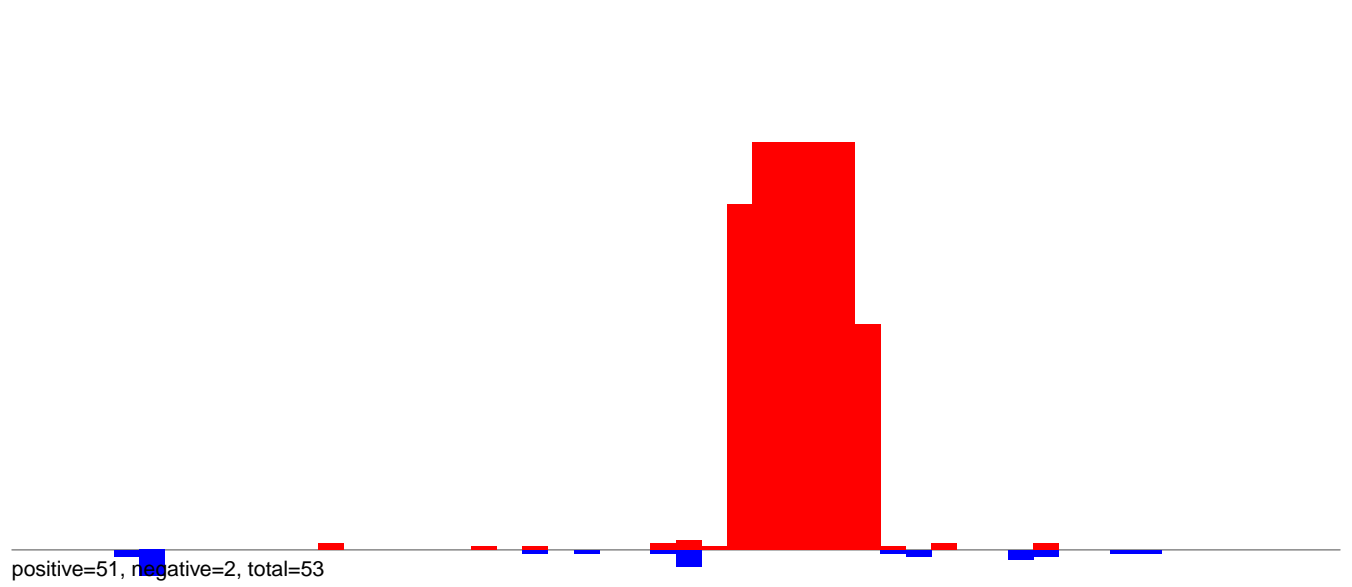
CuQuin_Testes_NL.24_35.rep



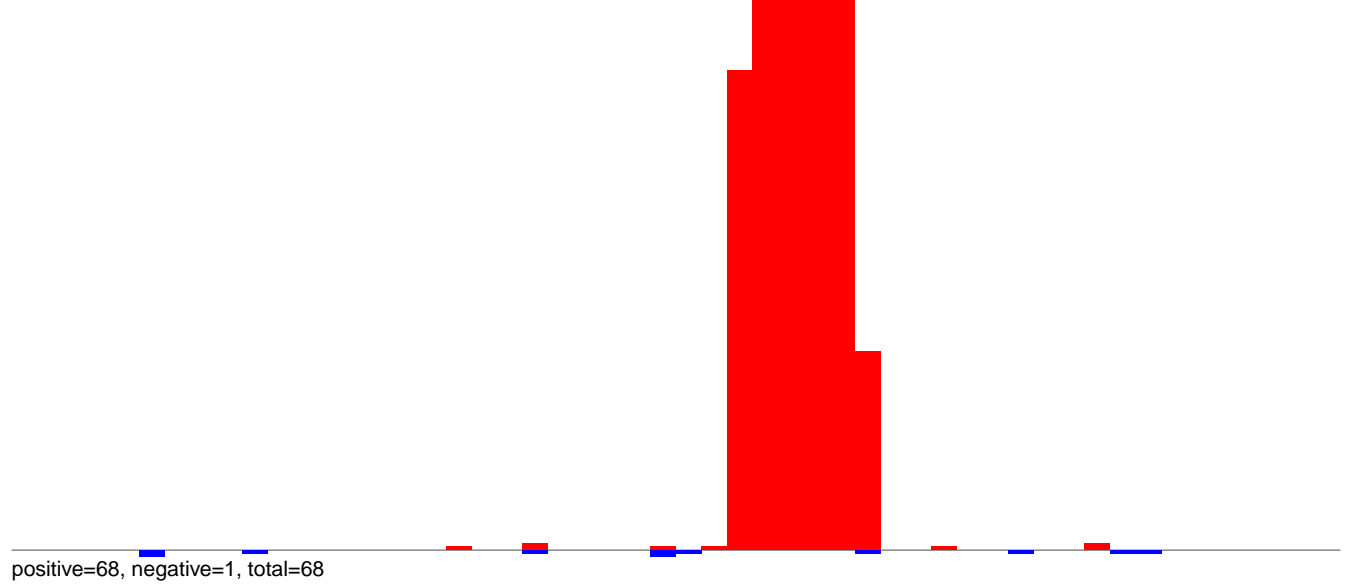
CuQuin_Testes_NL.rep



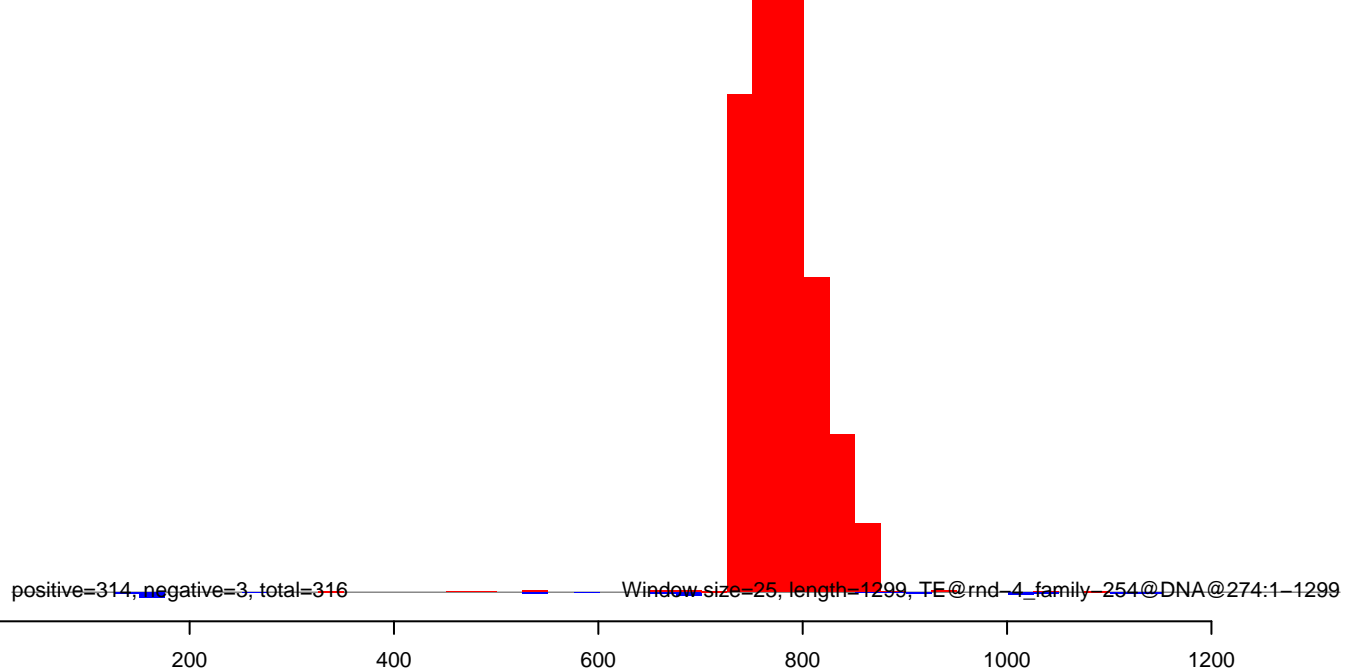
CuQuin_Testes_NL.18_23.rep



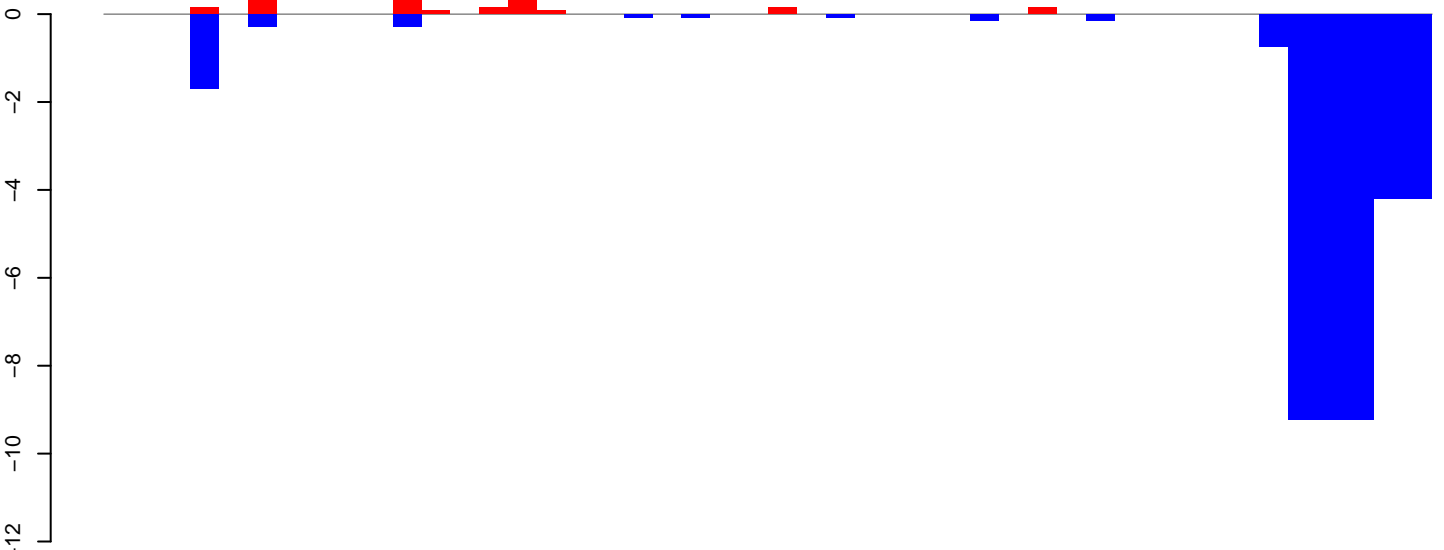
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

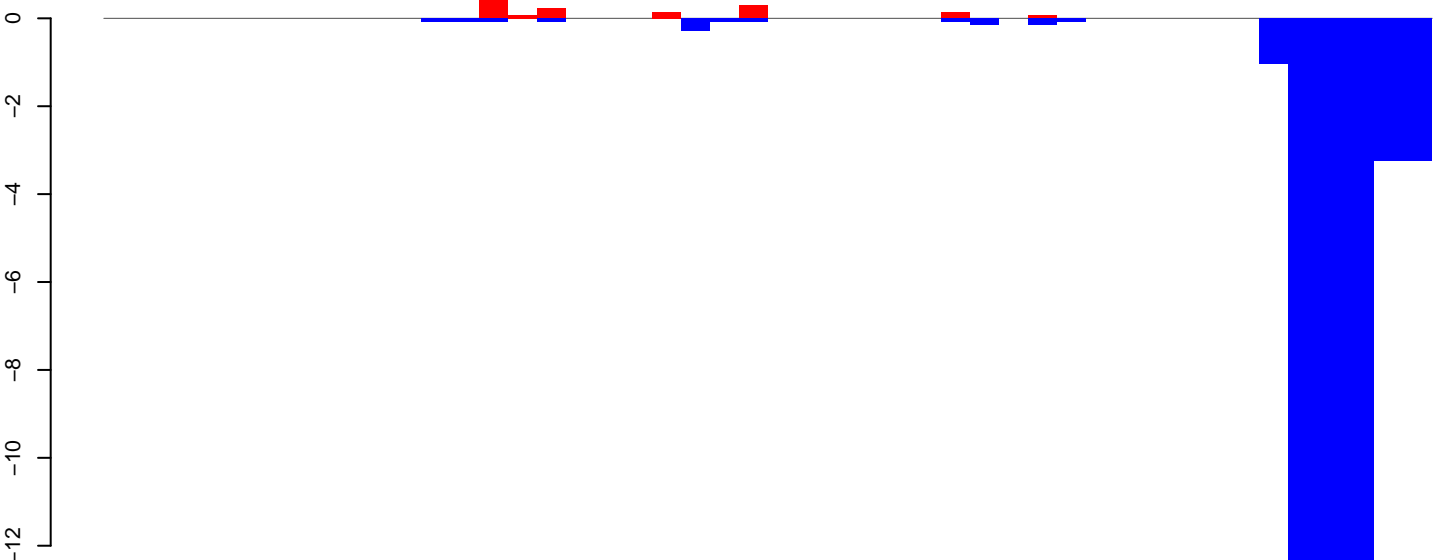


CuQuin_Testes_NL.18_23.rep



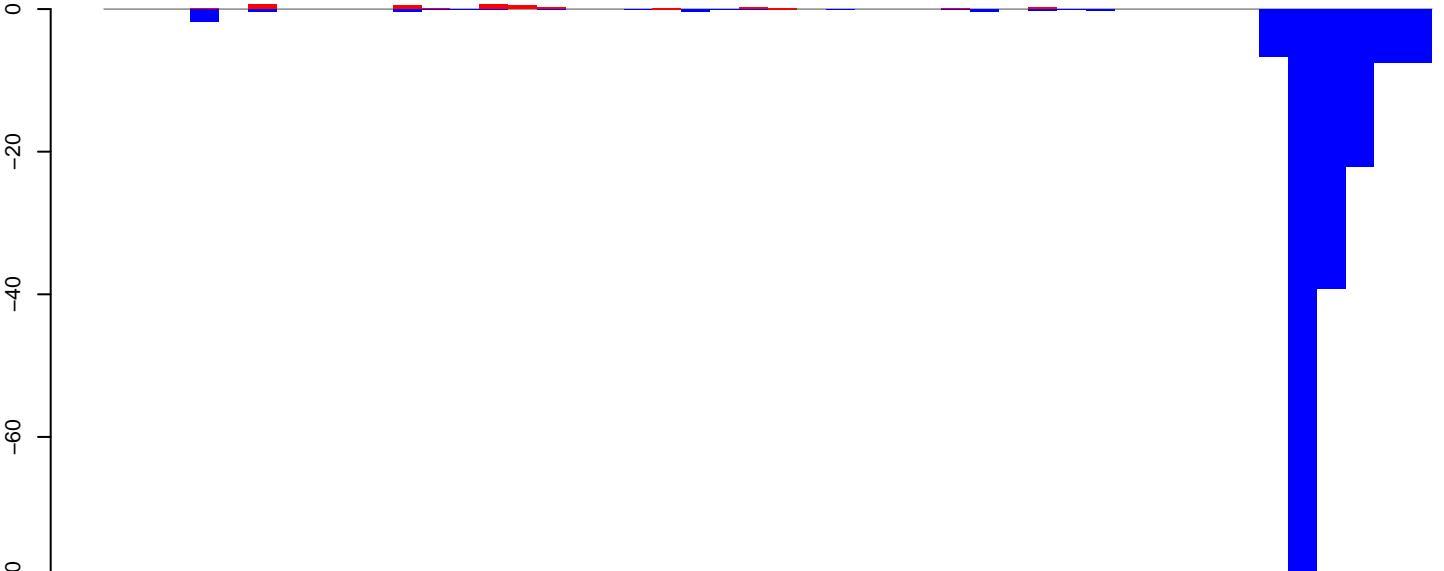
positive=2, negative=40, total=42

CuQuin_Testes_NL.24_35.rep



positive=1, negative=47, total=49

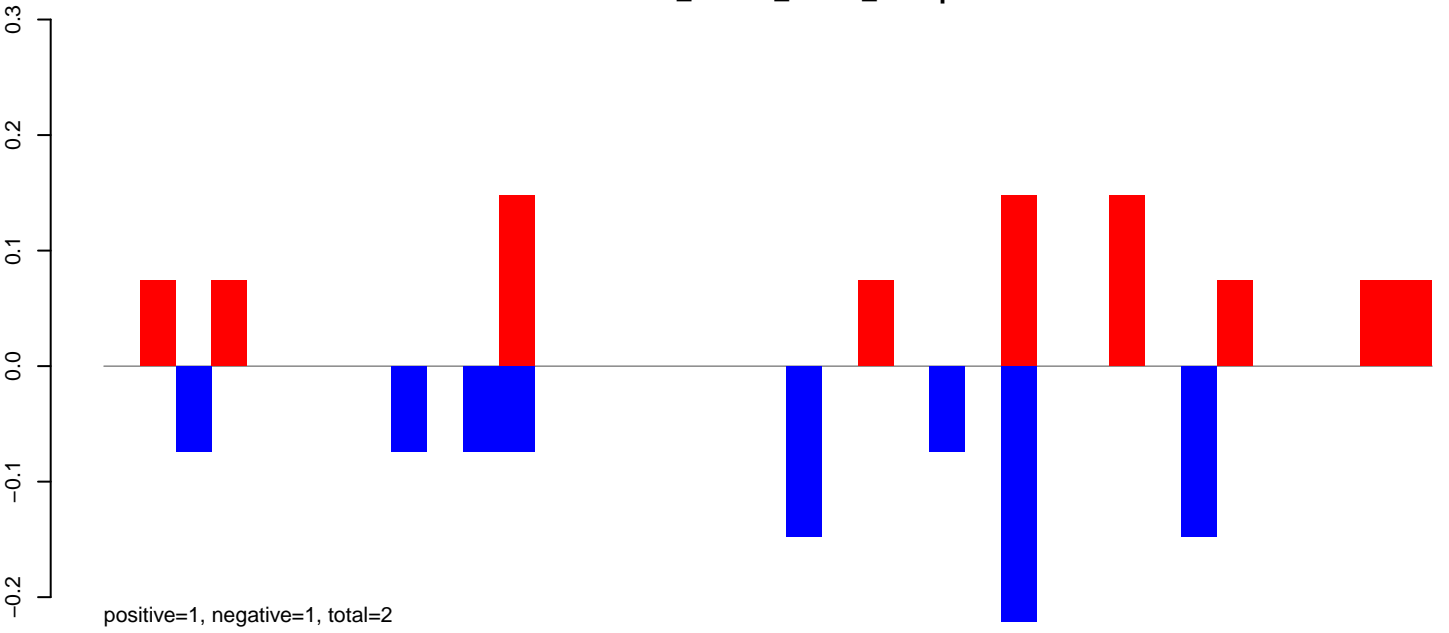
CuQuin_Testes_NL.rep



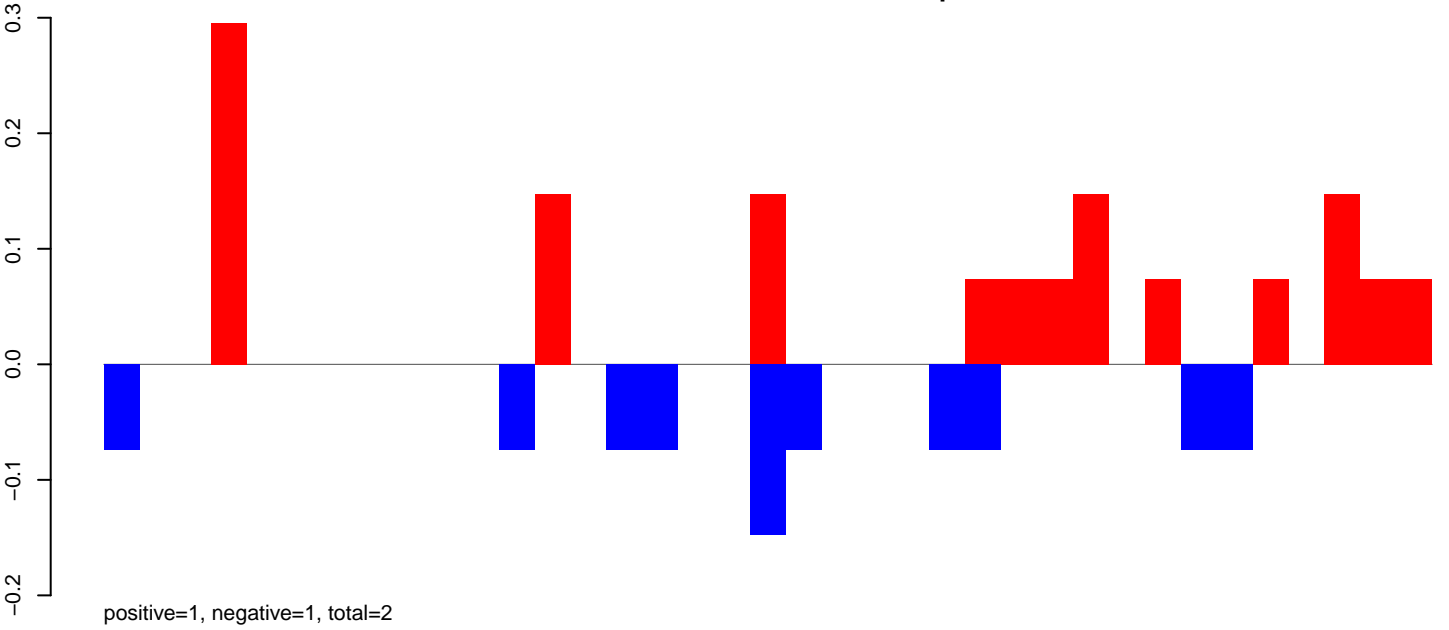
positive=4, negative=170, total=174

Window size=25, length=1133, TE@rnd-4_family-1436@DNA_Zator@905:1-1133

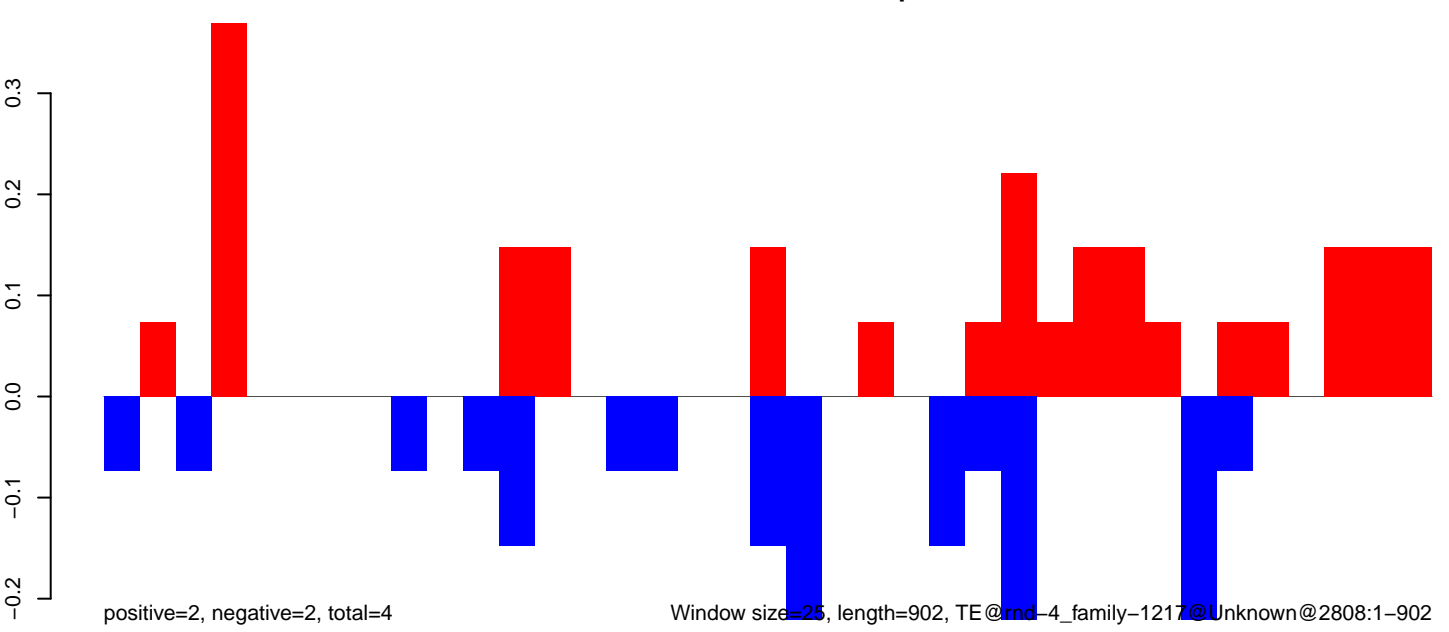
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

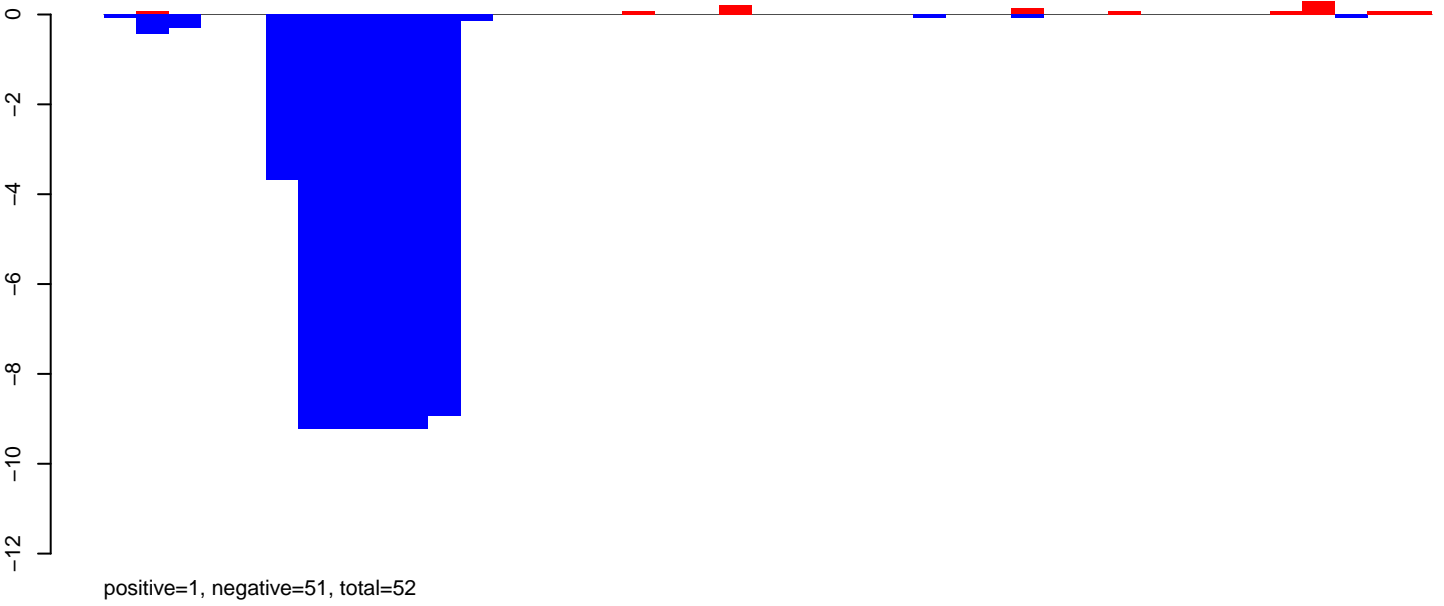


CuQuin_Testes_NL.rep

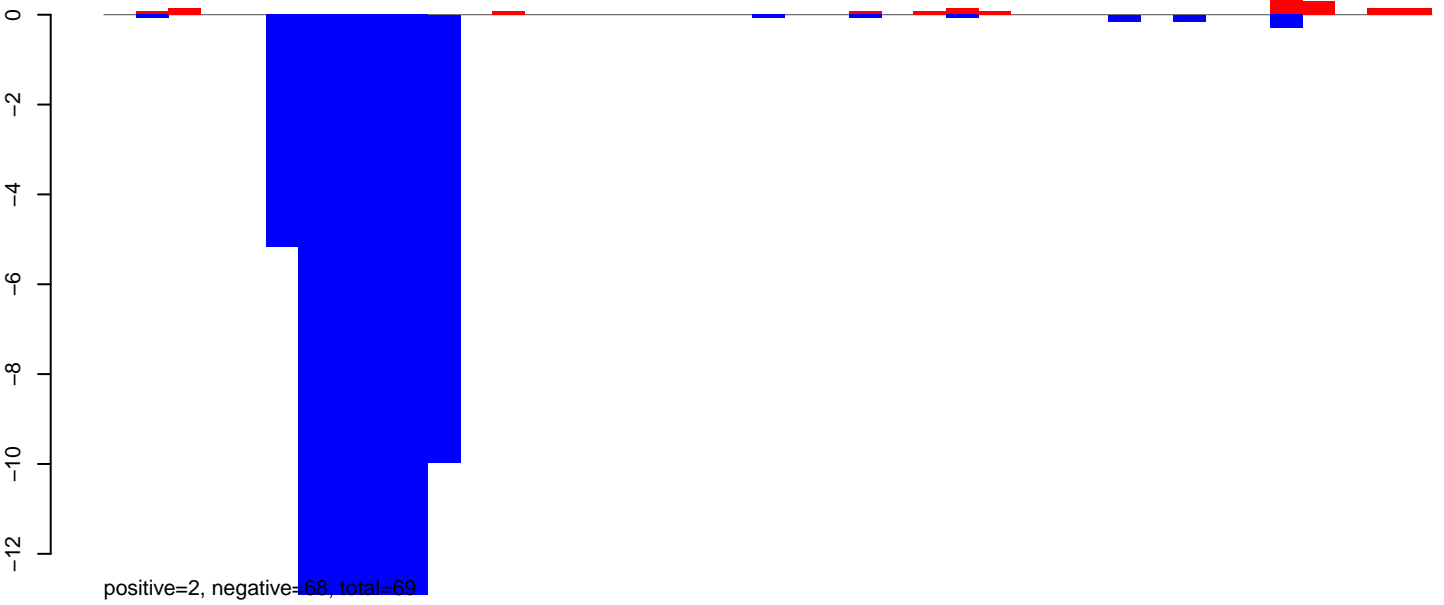


Window size=25, length=902, TE@md-4_family-1217@Unknown@2808:1-902

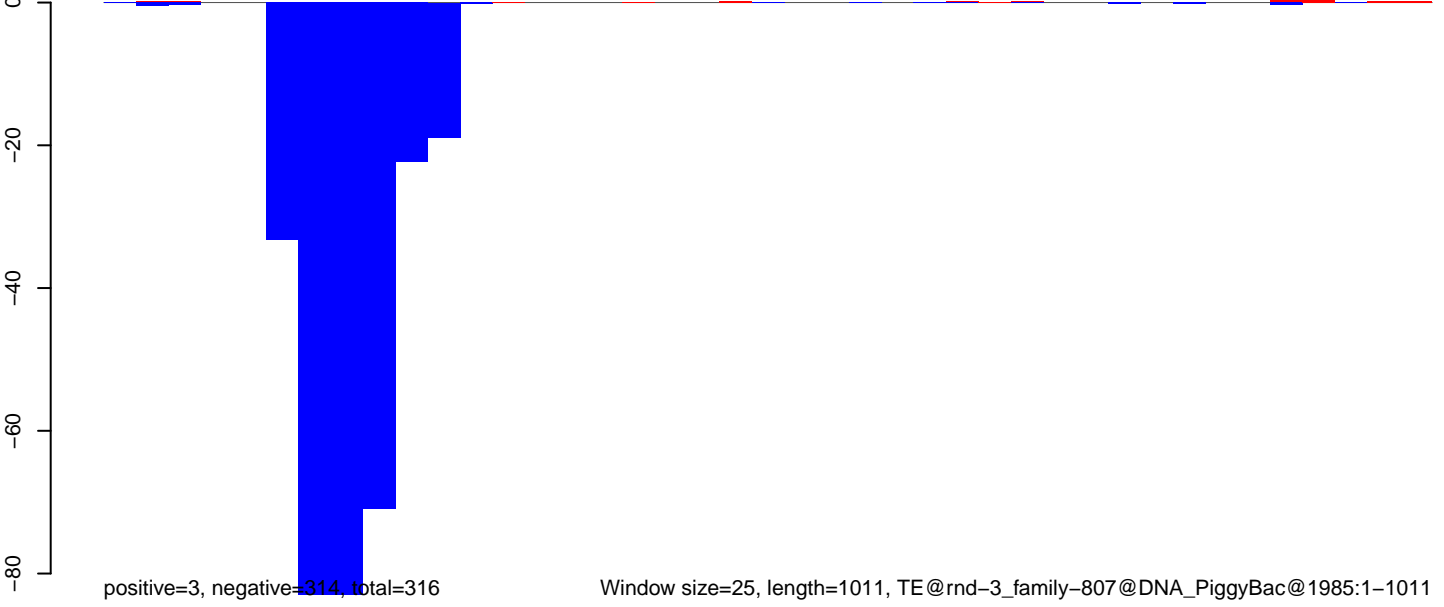
CuQuin_Testes_NL.18_23.rep



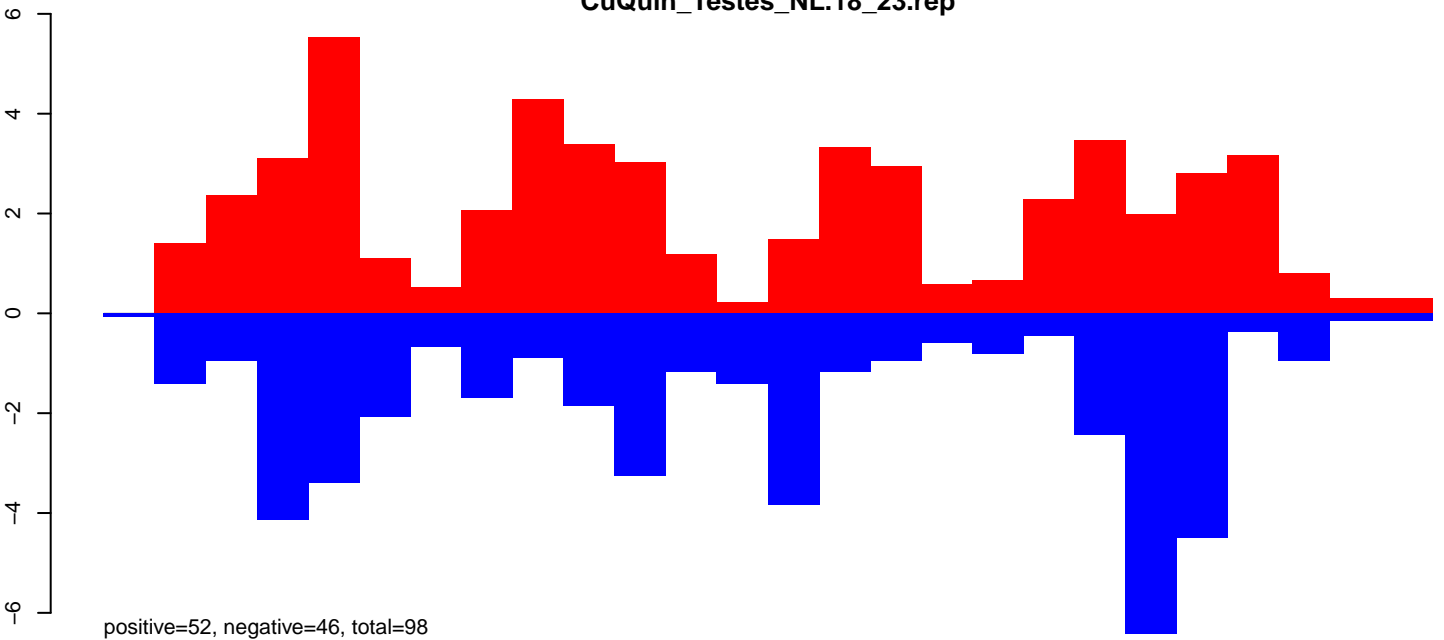
CuQuin_Testes_NL.24_35.rep



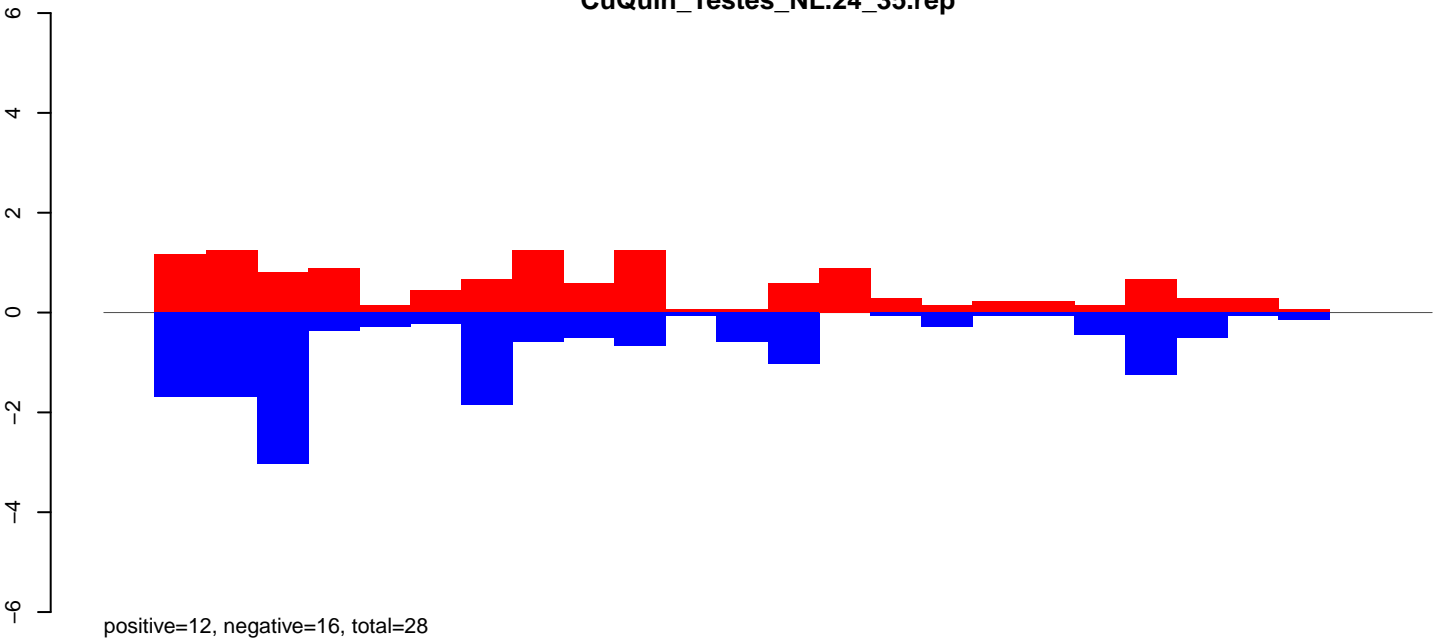
CuQuin_Testes_NL.rep



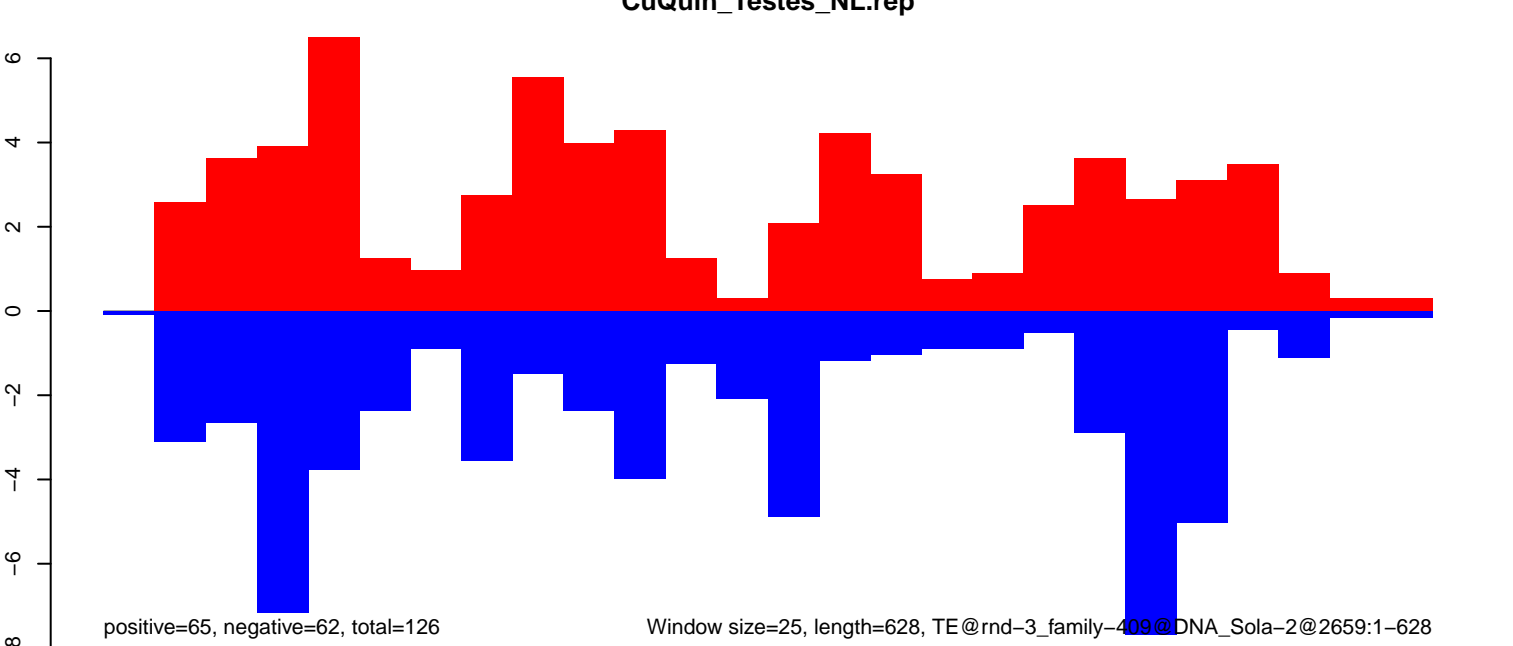
CuQuin_Testes_NL.18_23.rep



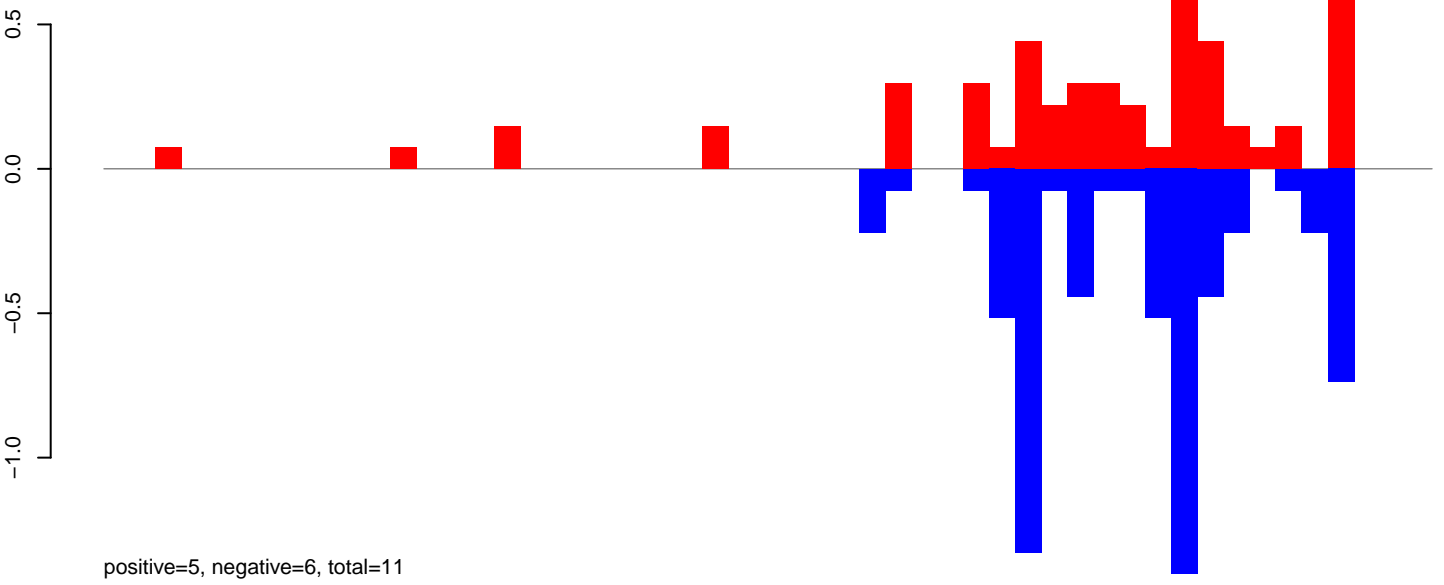
CuQuin_Testes_NL.24_35.rep



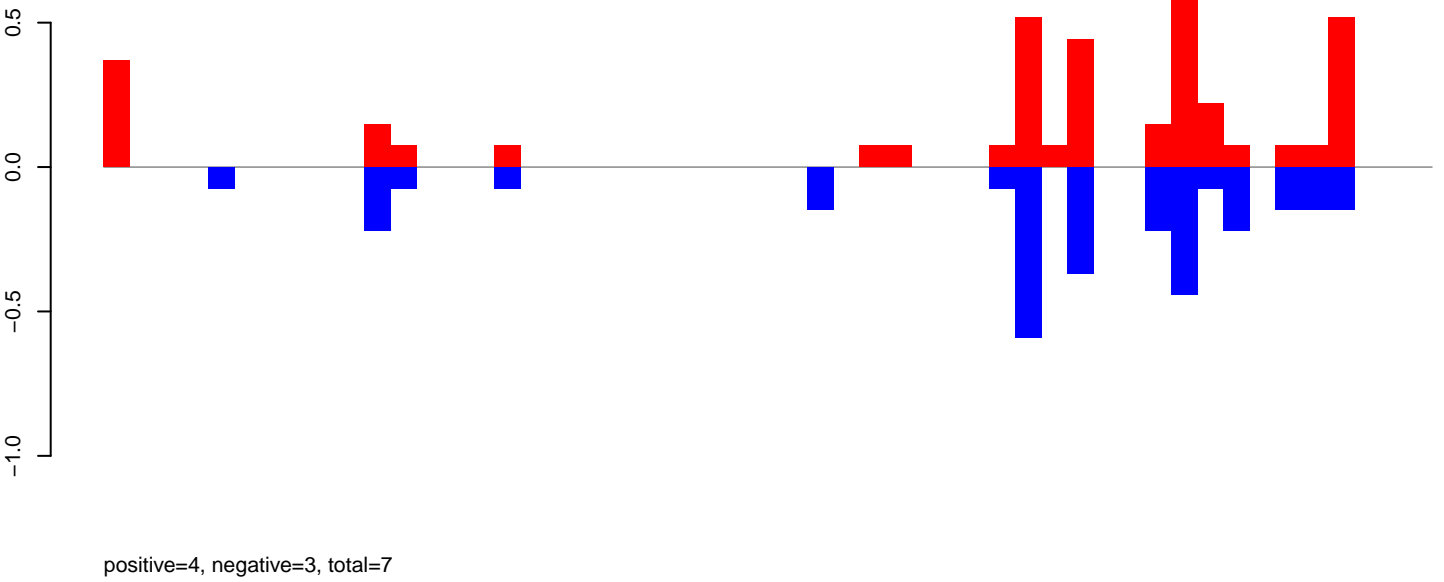
CuQuin_Testes_NL.rep



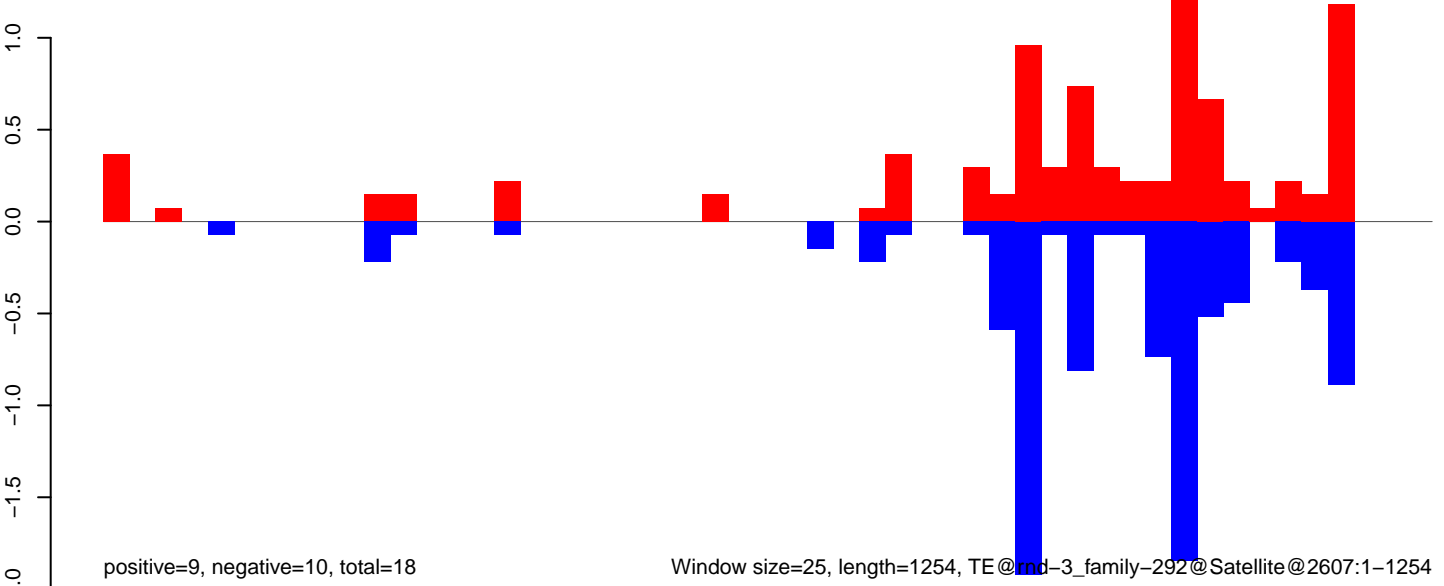
CuQuin_Testes_NL.18_23.rep



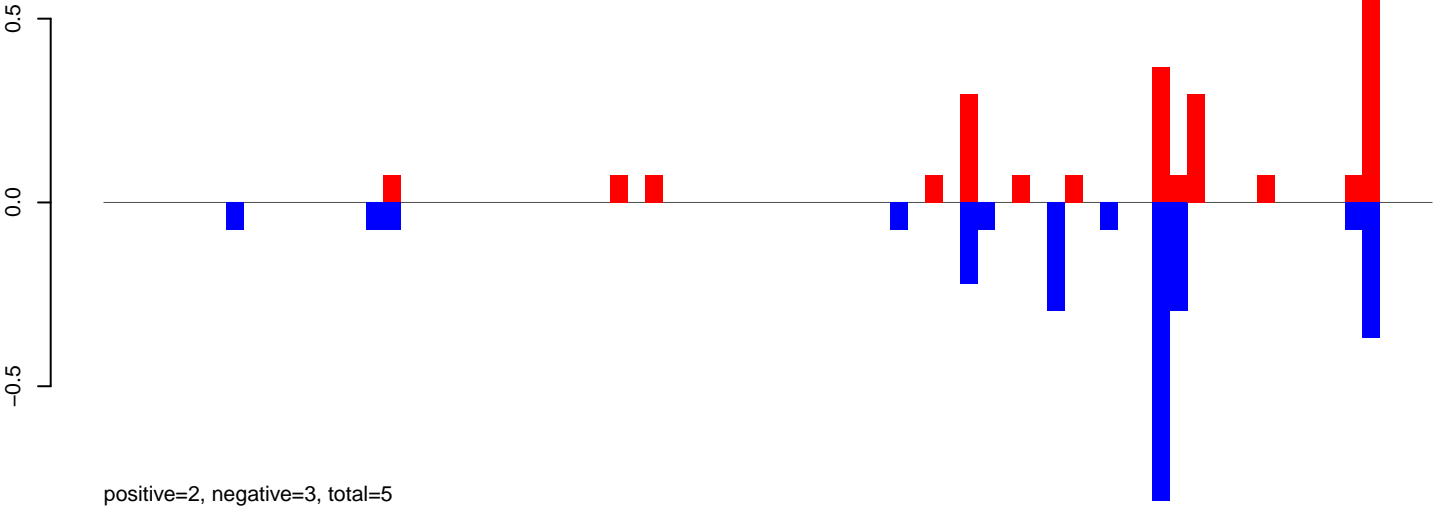
CuQuin_Testes_NL.24_35.rep



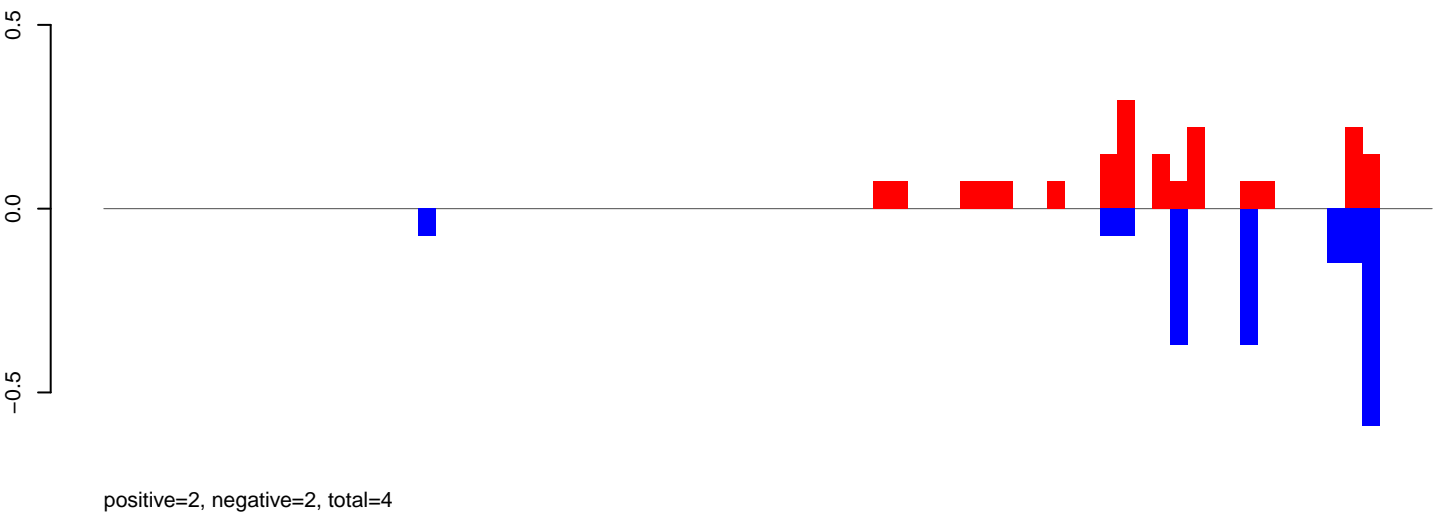
CuQuin_Testes_NL.rep



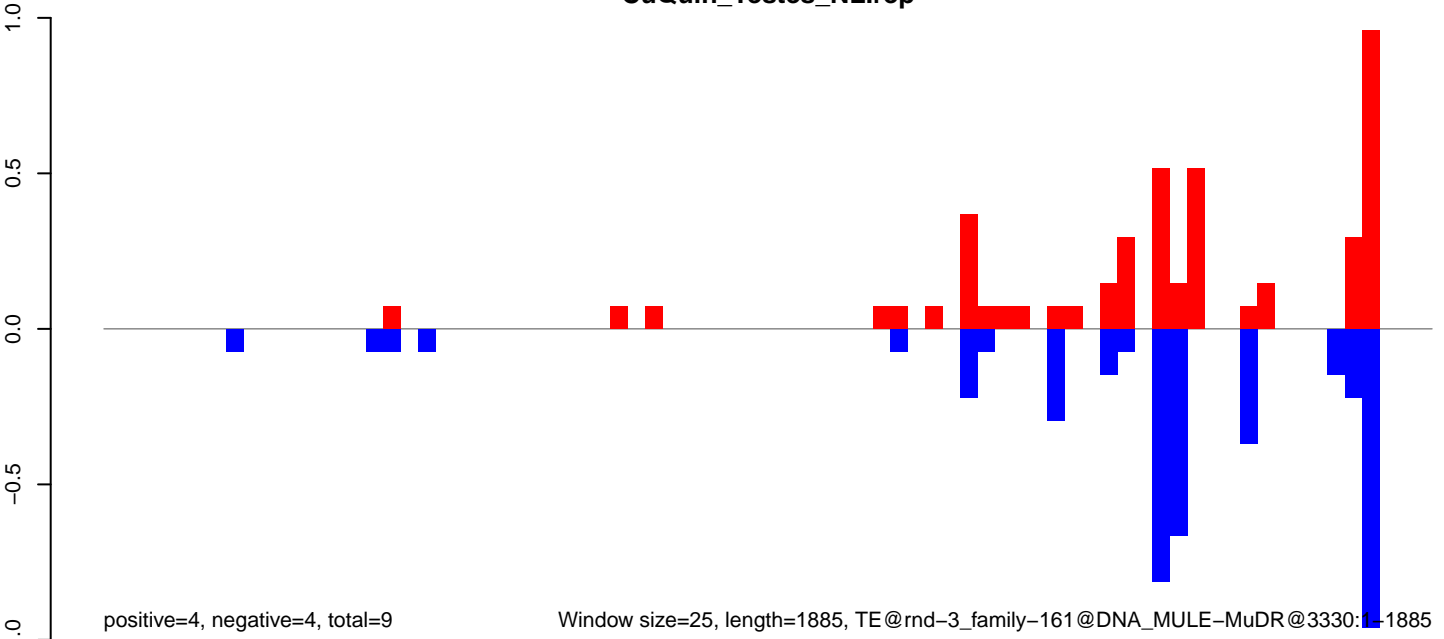
CuQuin_Testes_NL.18_23.rep



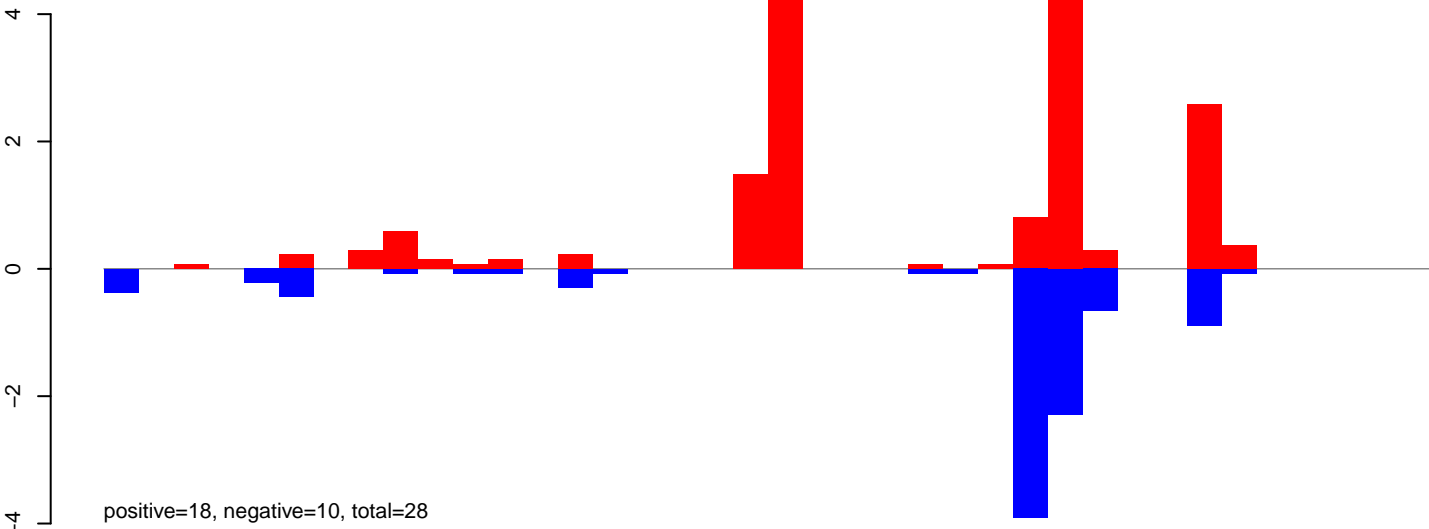
CuQuin_Testes_NL.24_35.rep



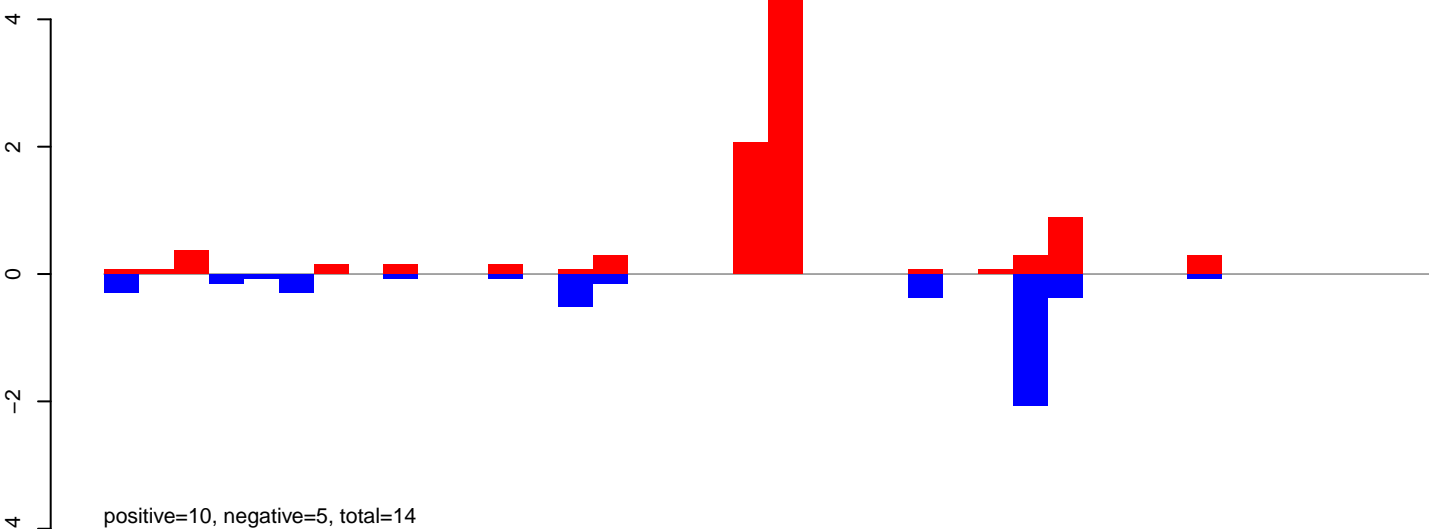
CuQuin_Testes_NL.rep



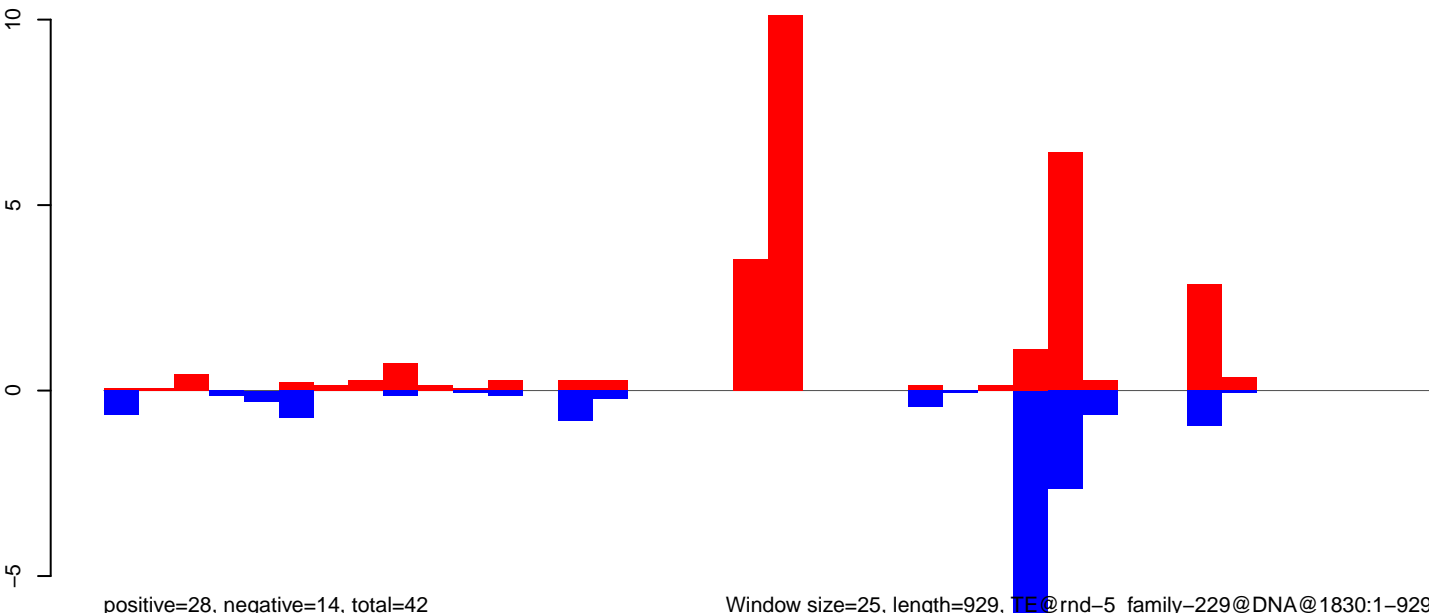
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



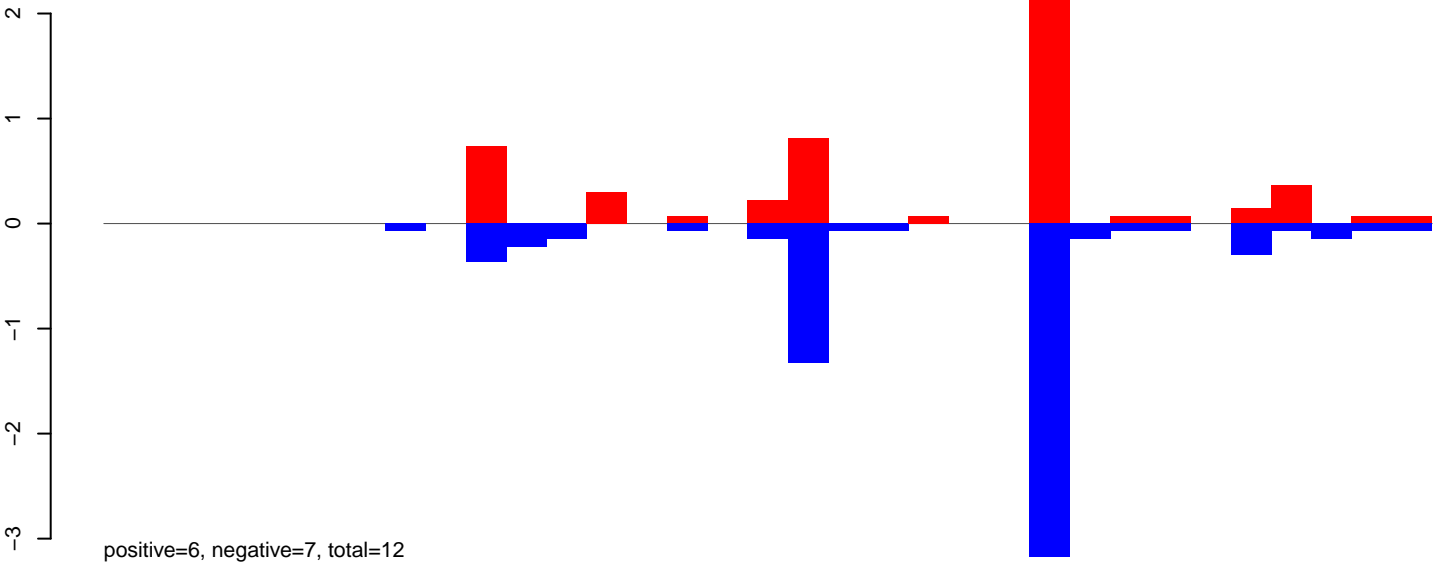
CuQuin_Testes_NL.rep



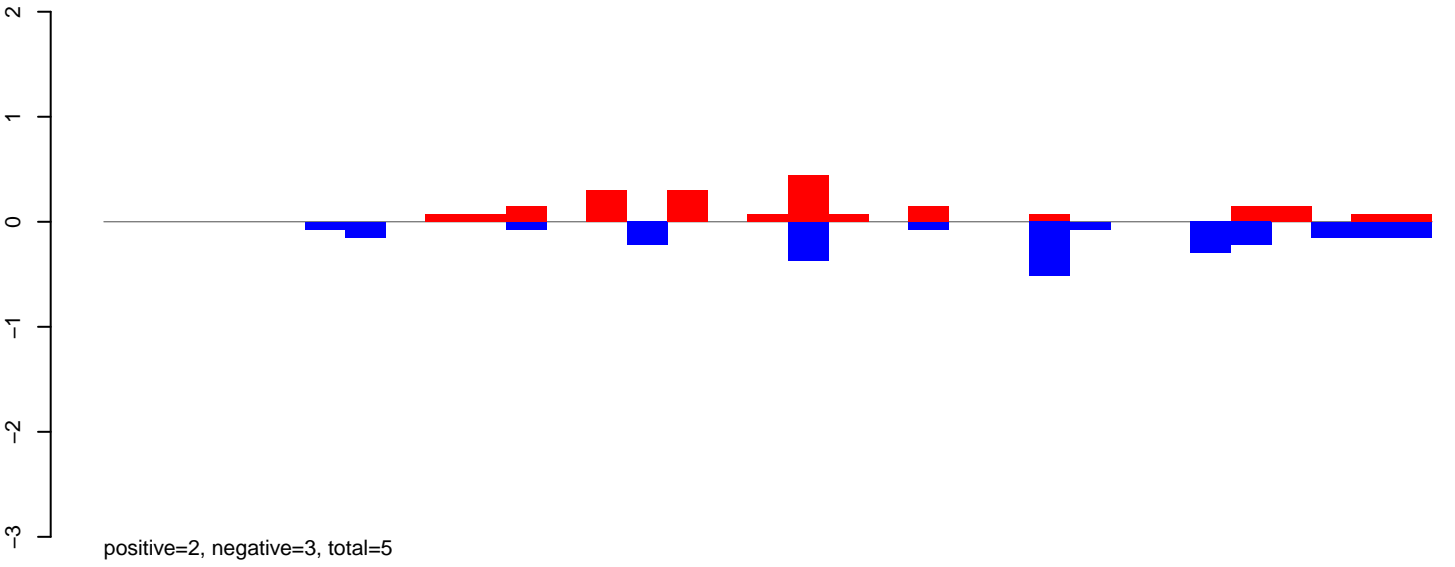
Window size=25, length=929, TE@rnd-5_family-229@DNA@1830:1-929

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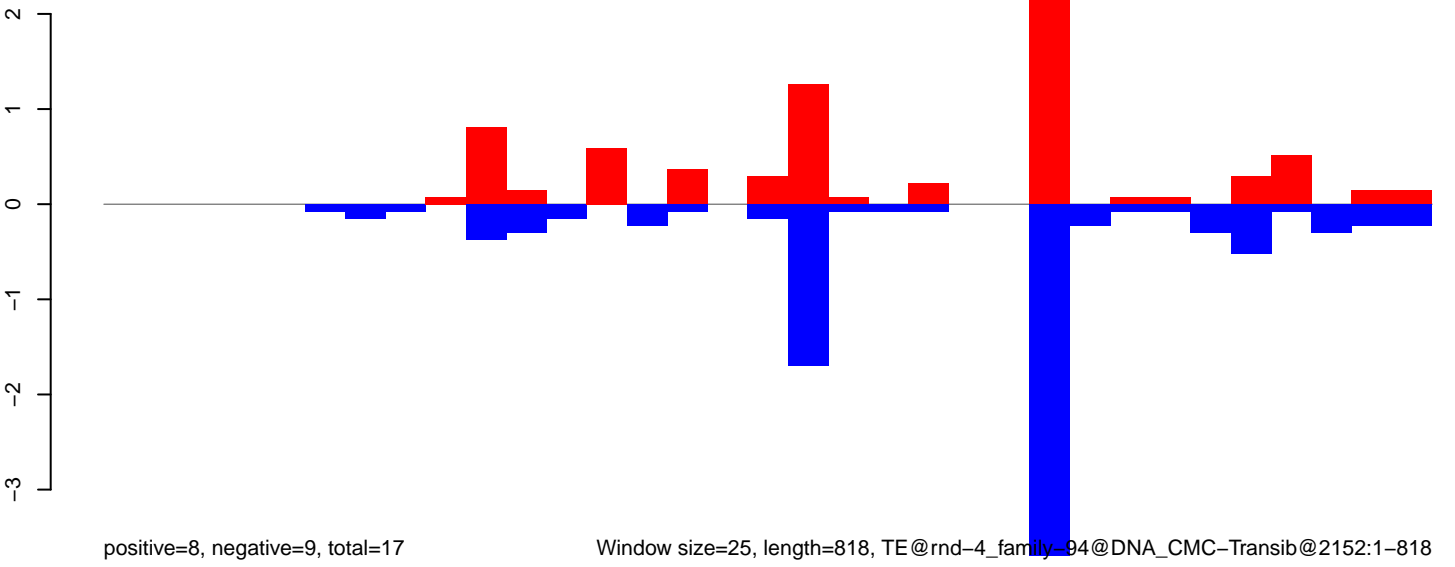
CuQuin_Testes_NL.18_23.rep



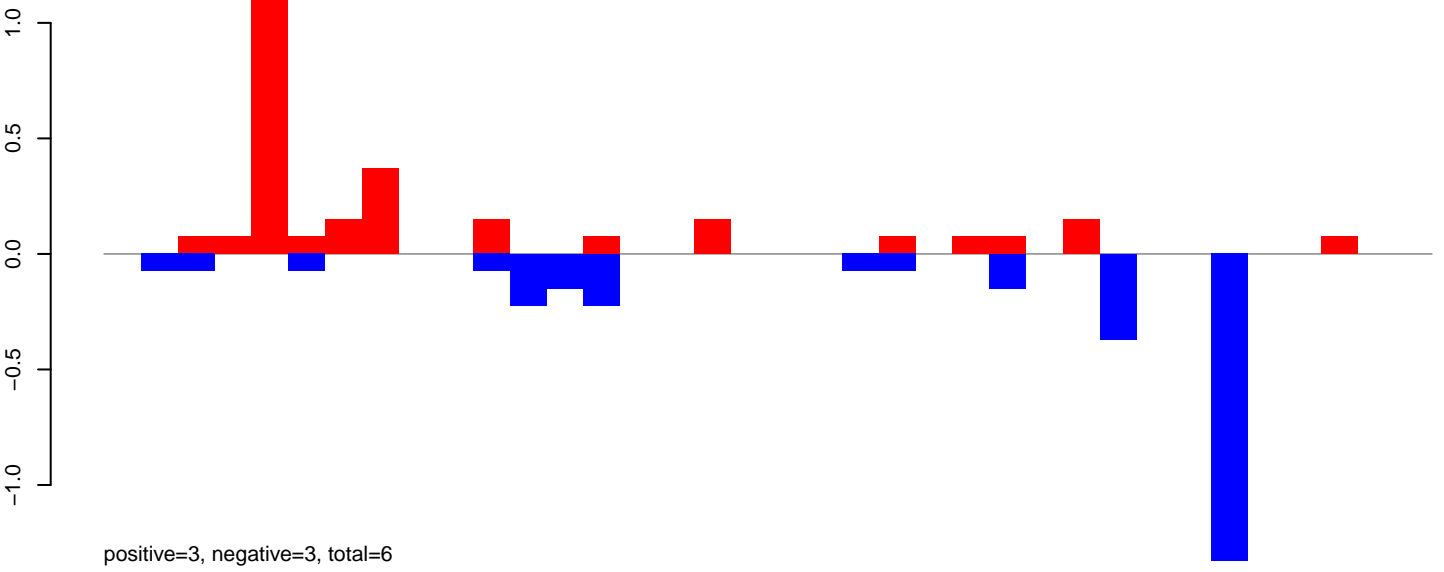
CuQuin_Testes_NL.24_35.rep



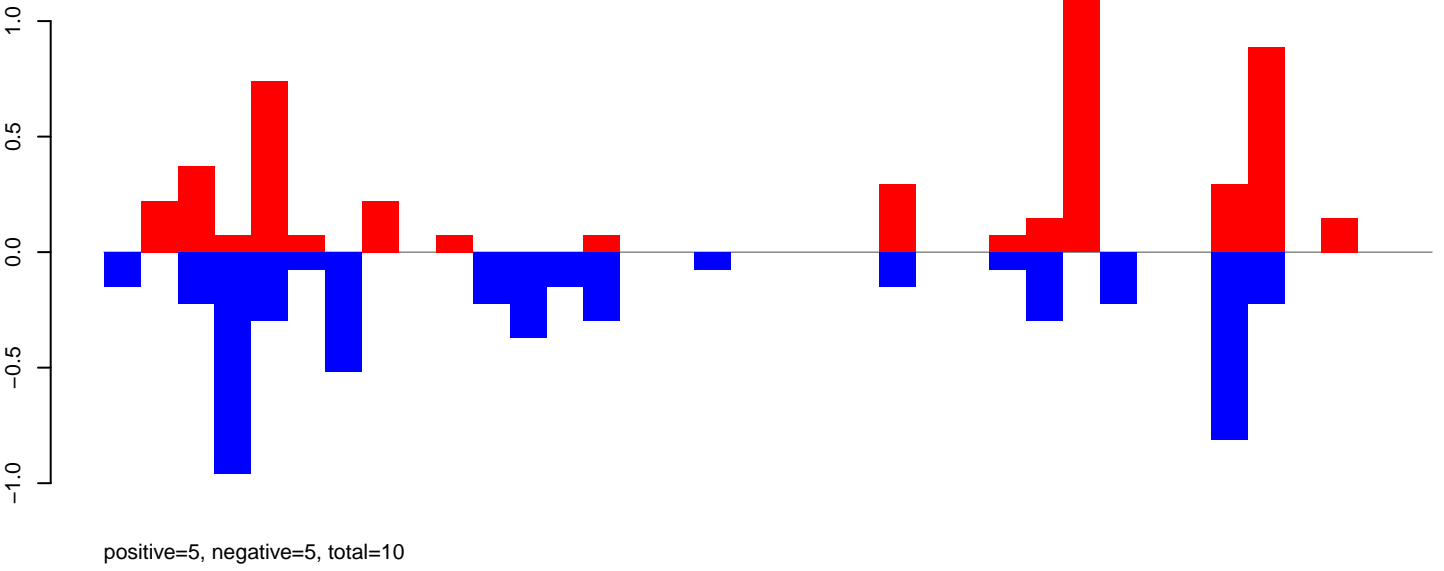
CuQuin_Testes_NL.rep



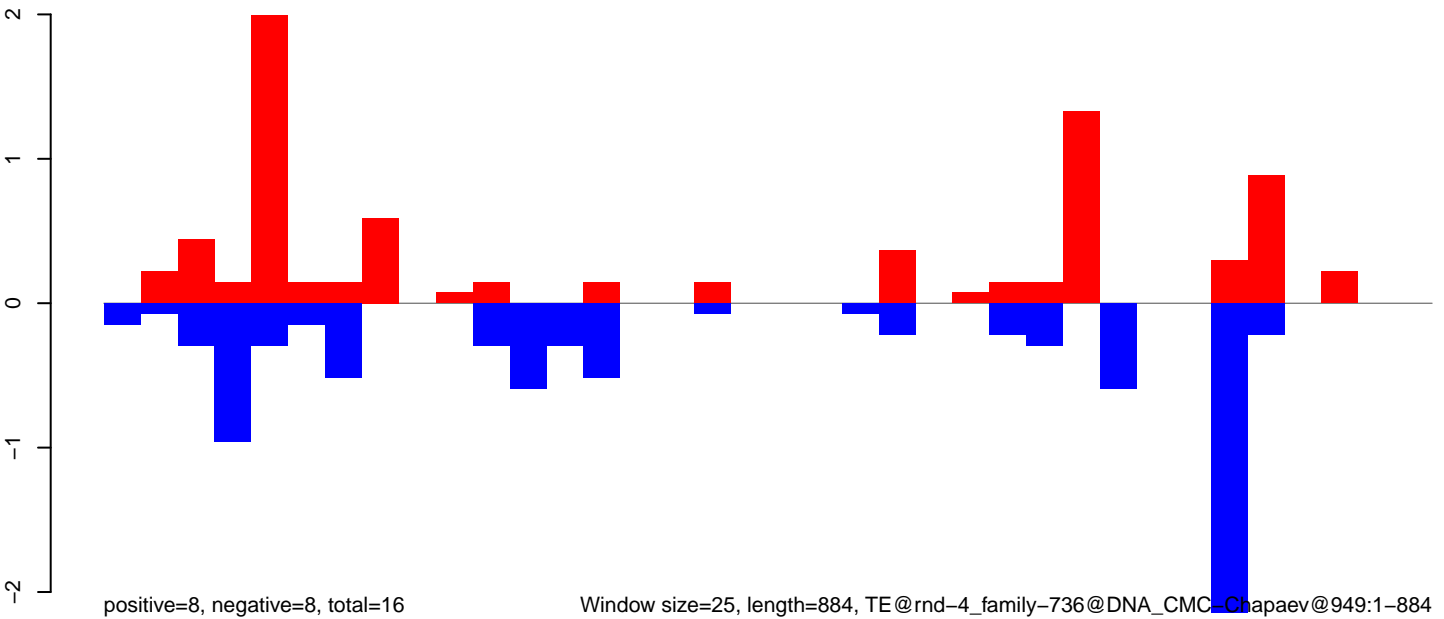
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

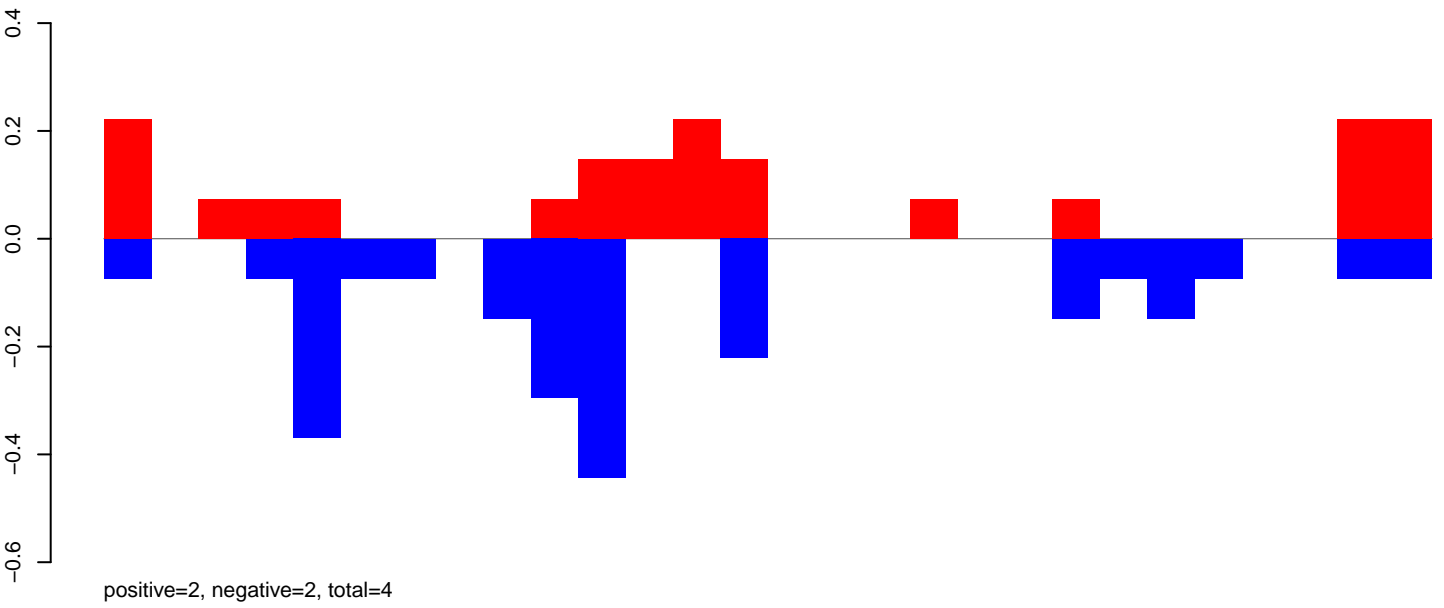


CuQuin_Testes_NL.rep

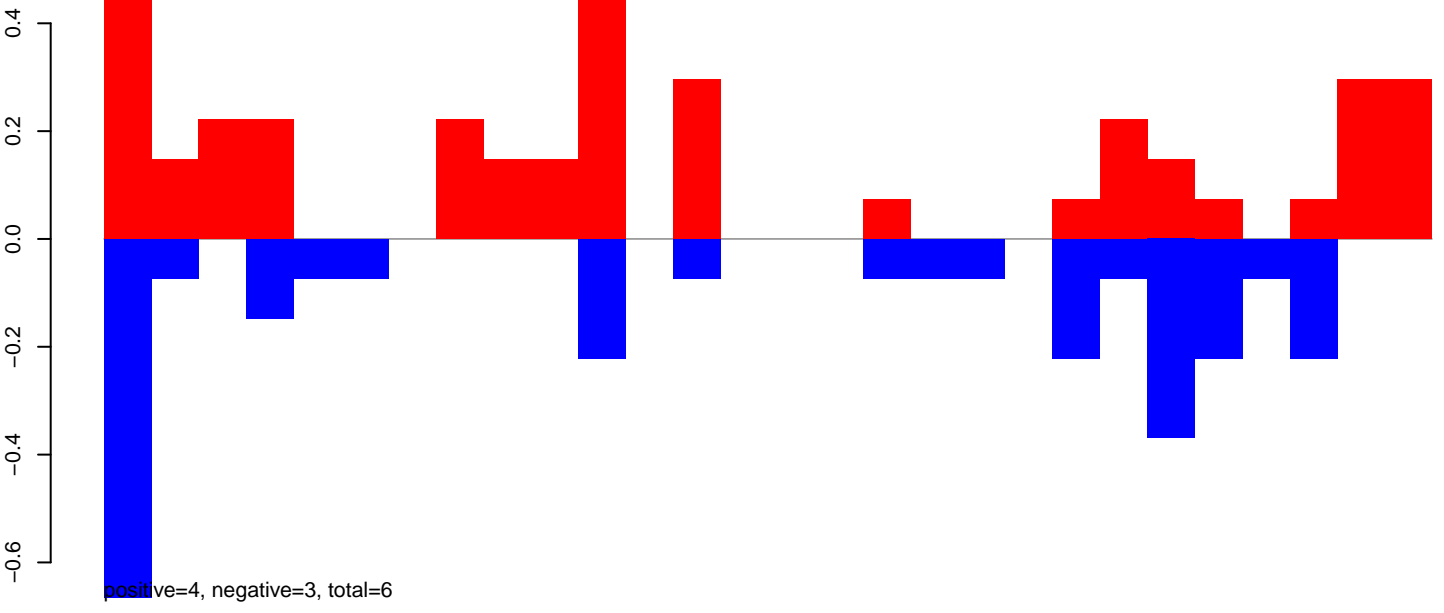


Window size=25, length=884, TE@rnd-4_family-736@DNA_CMC-Chapaev@949:1-884

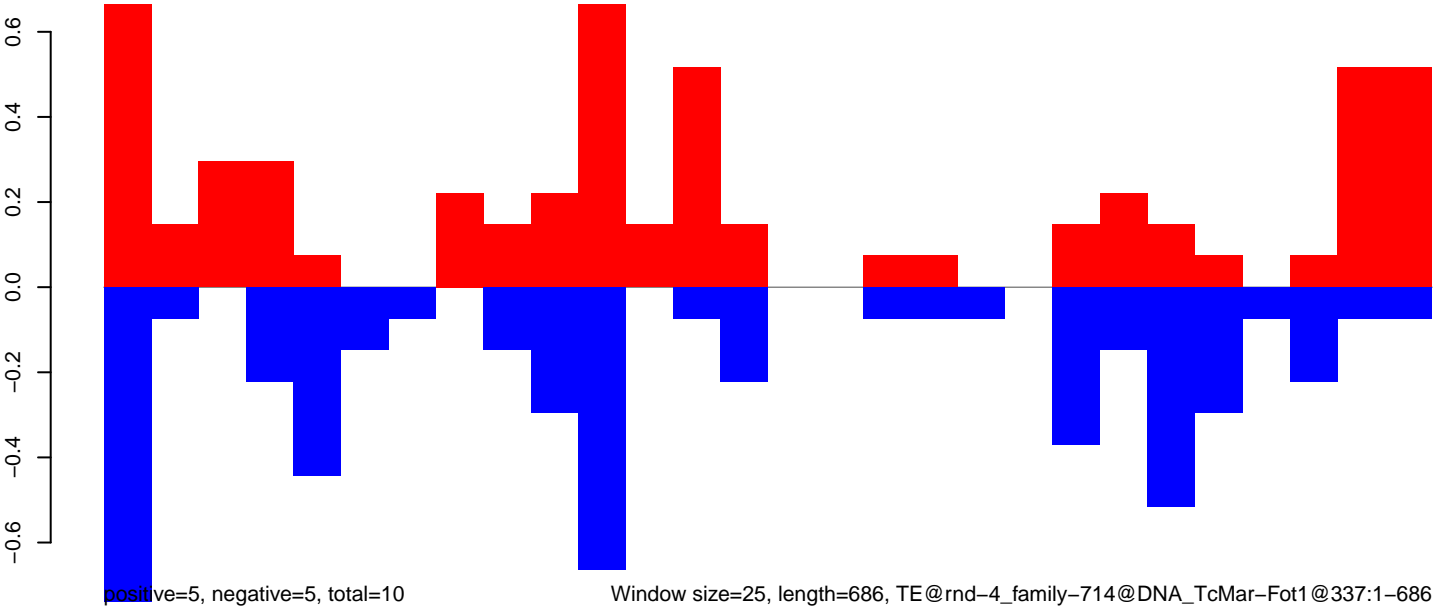
CuQuin_Testes_NL.18_23.rep



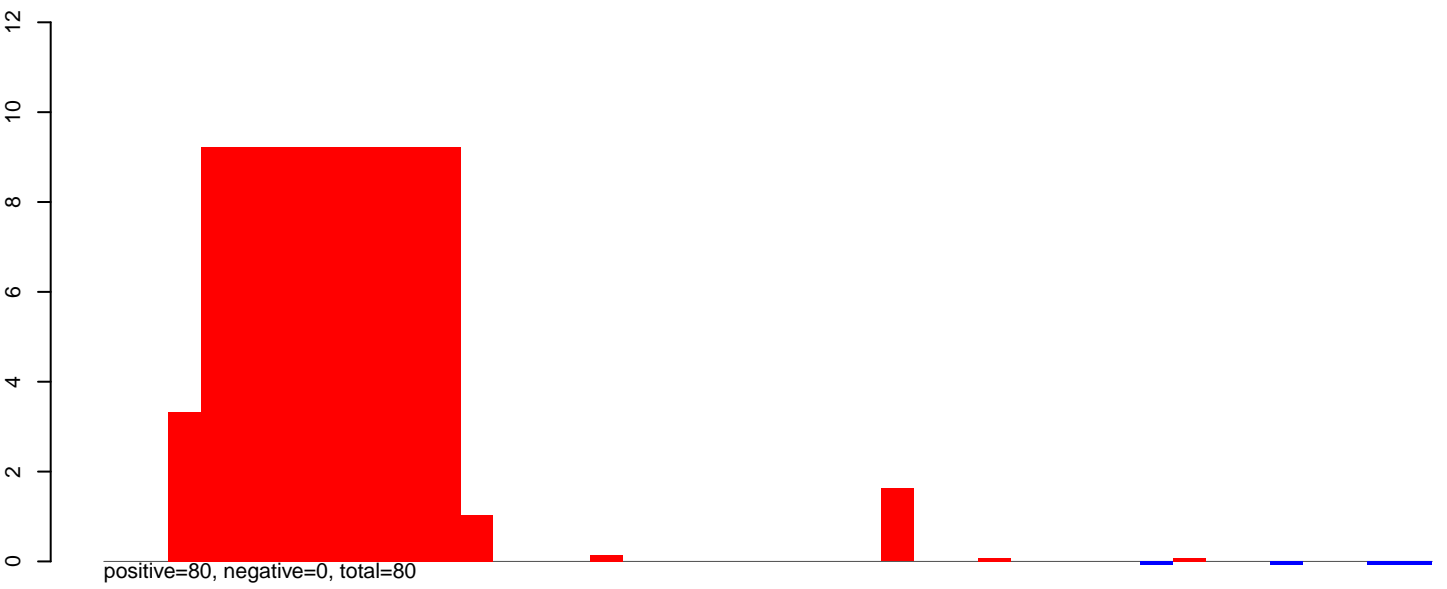
CuQuin_Testes_NL.24_35.rep



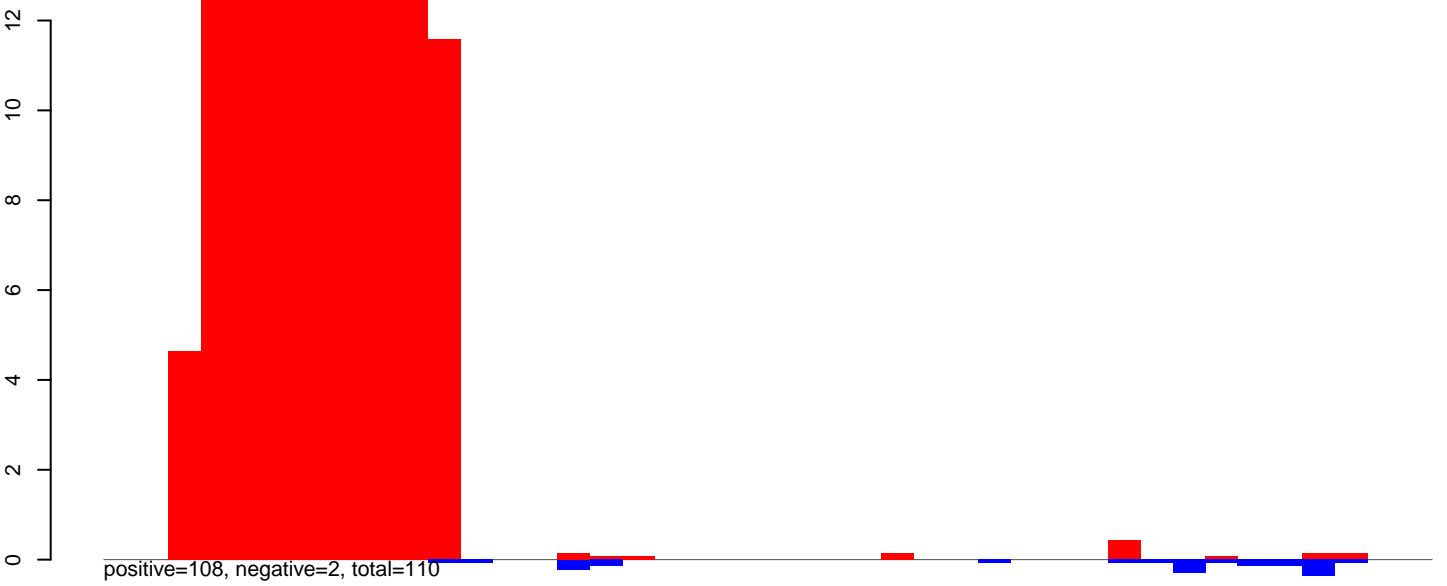
CuQuin_Testes_NL.rep



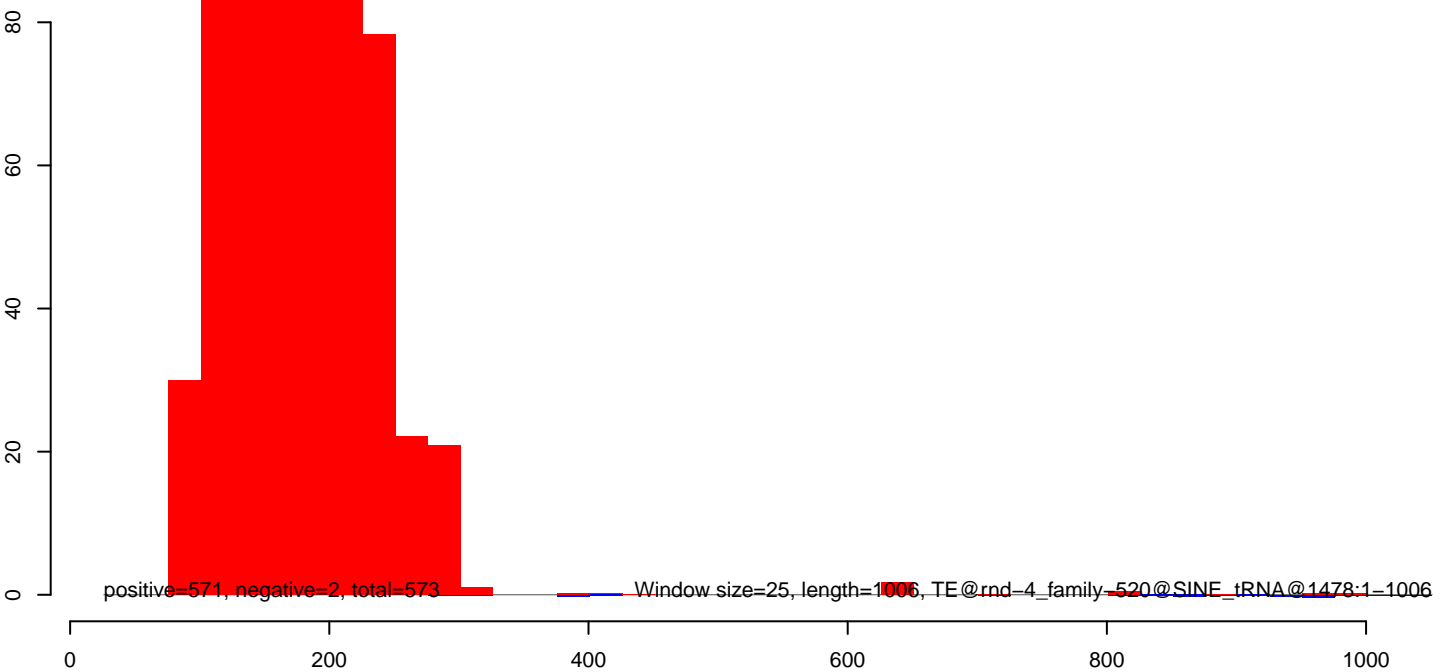
CuQuin_Testes_NL.18_23.rep



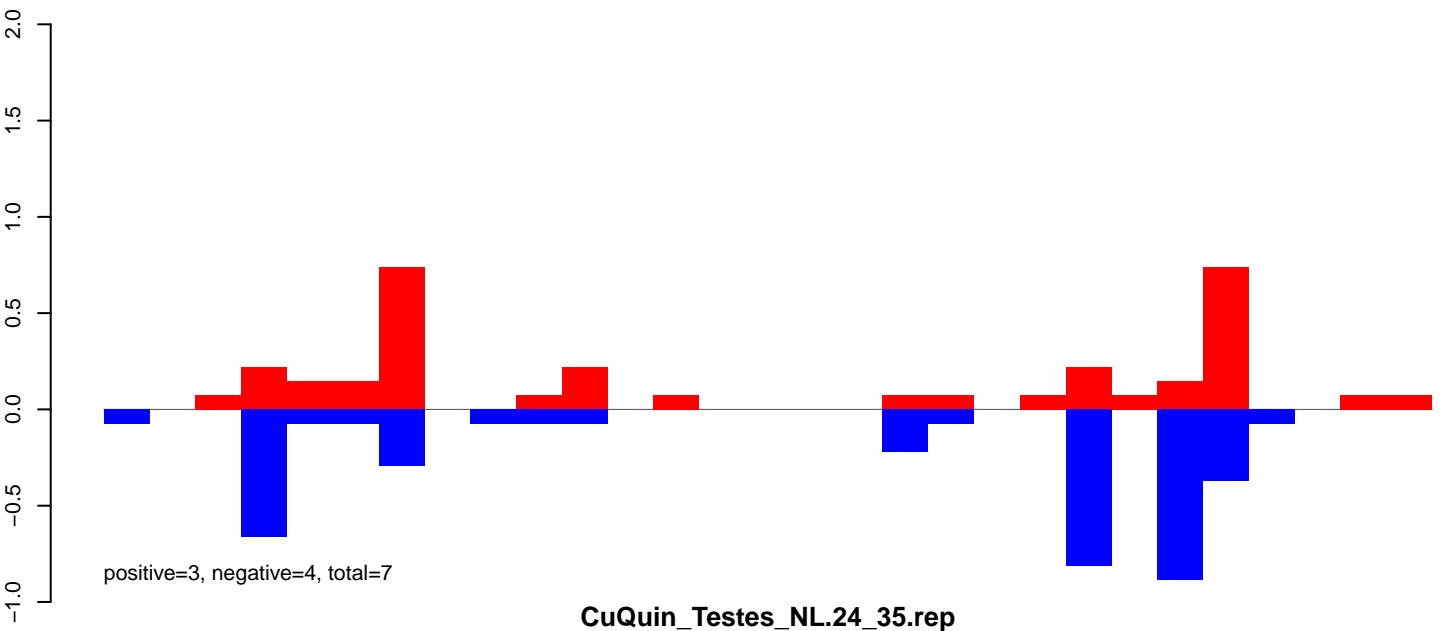
CuQuin_Testes_NL.24_35.rep



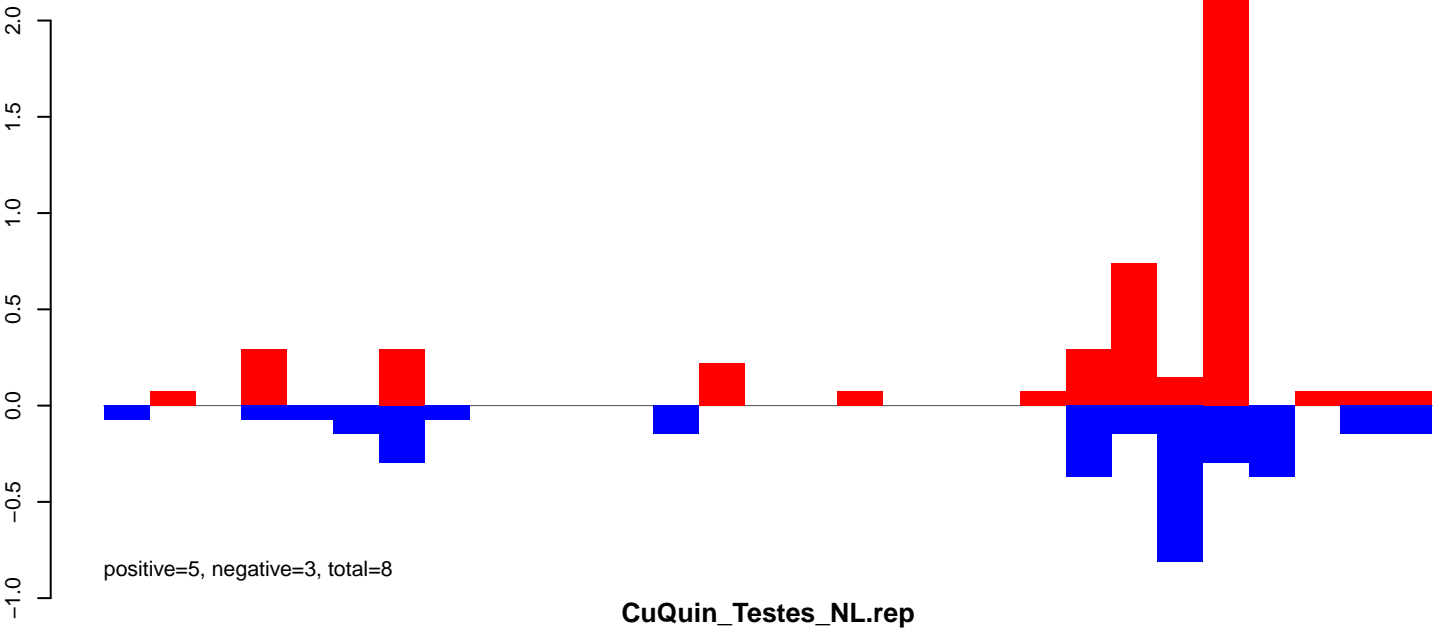
CuQuin_Testes_NL.rep



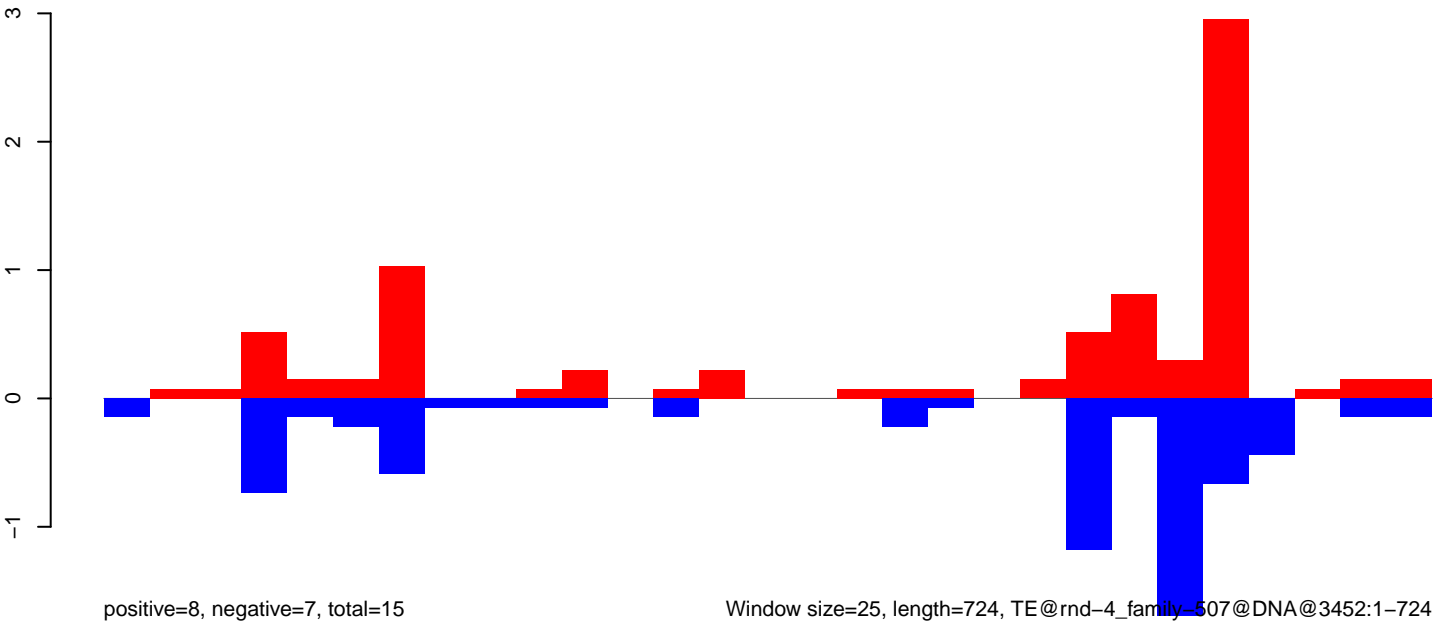
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

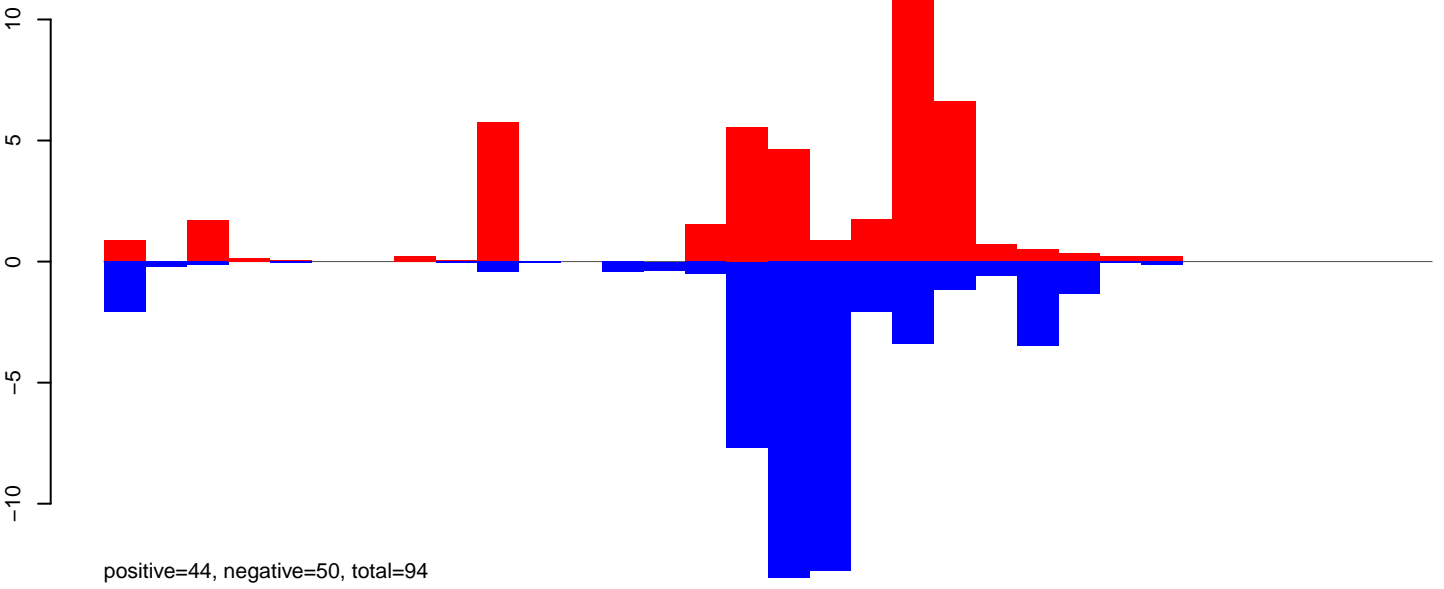


CuQuin_Testes_NL.rep

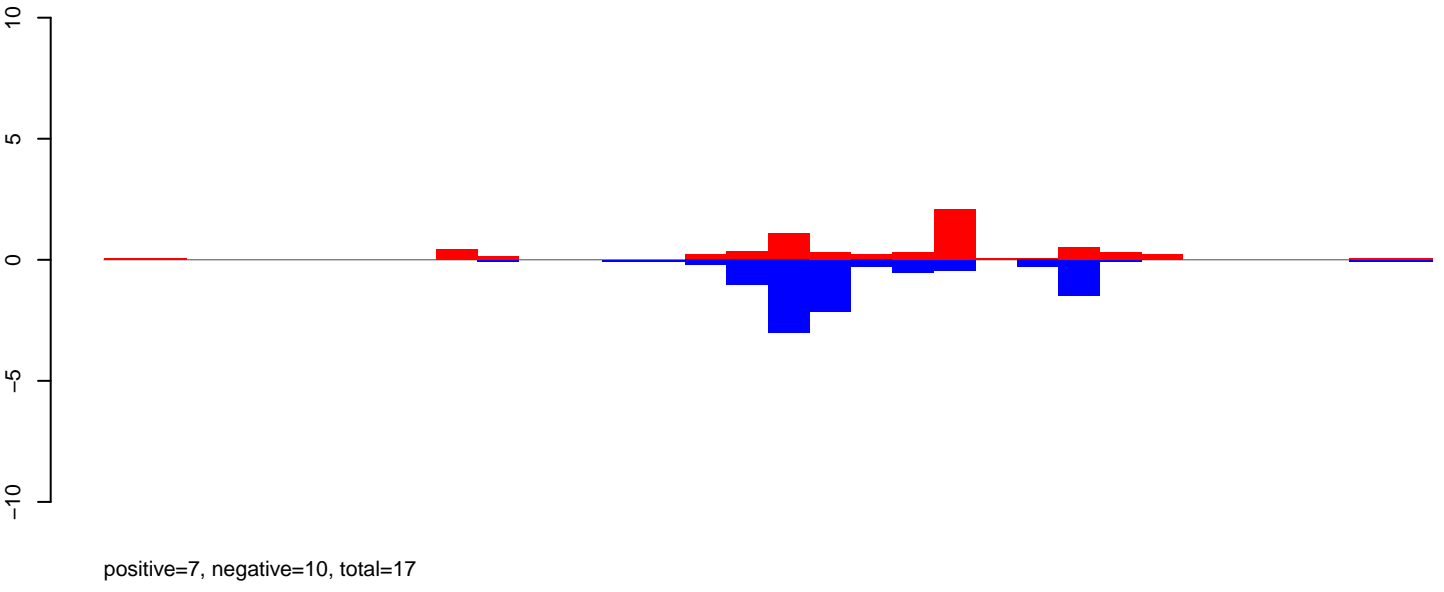


Window size=25, length=724, TE@rnd-4_family-507@DNA@3452:1-724

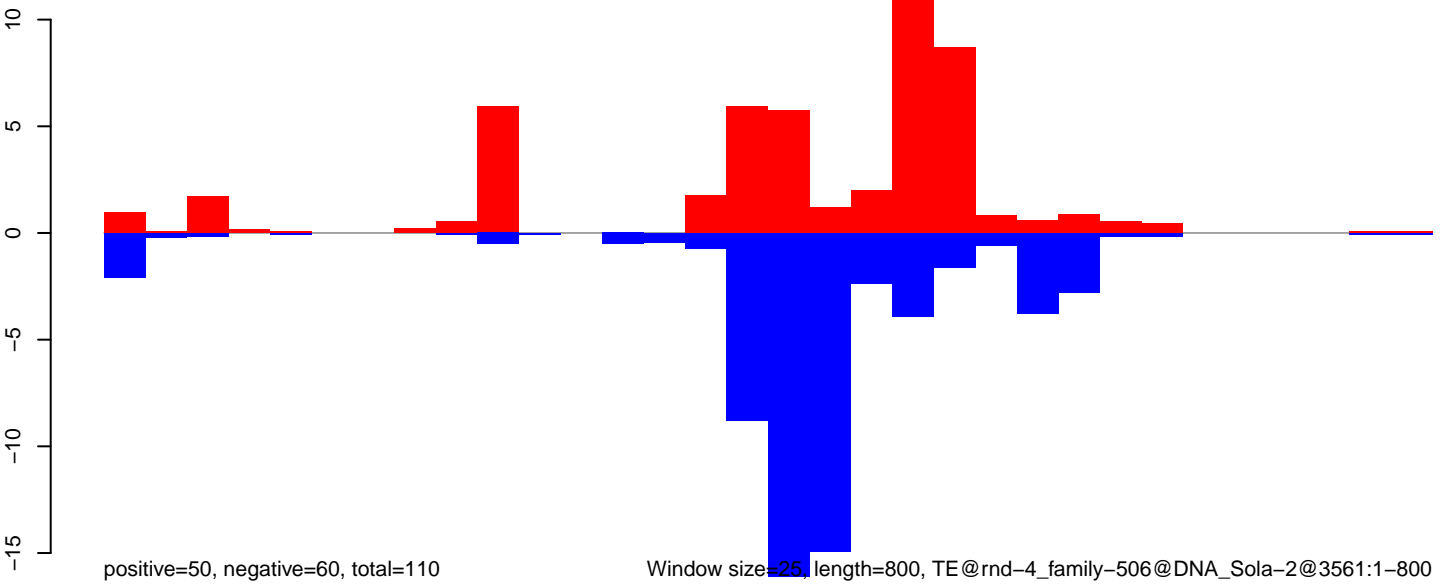
CuQuin_Testes_NL.18_23.rep



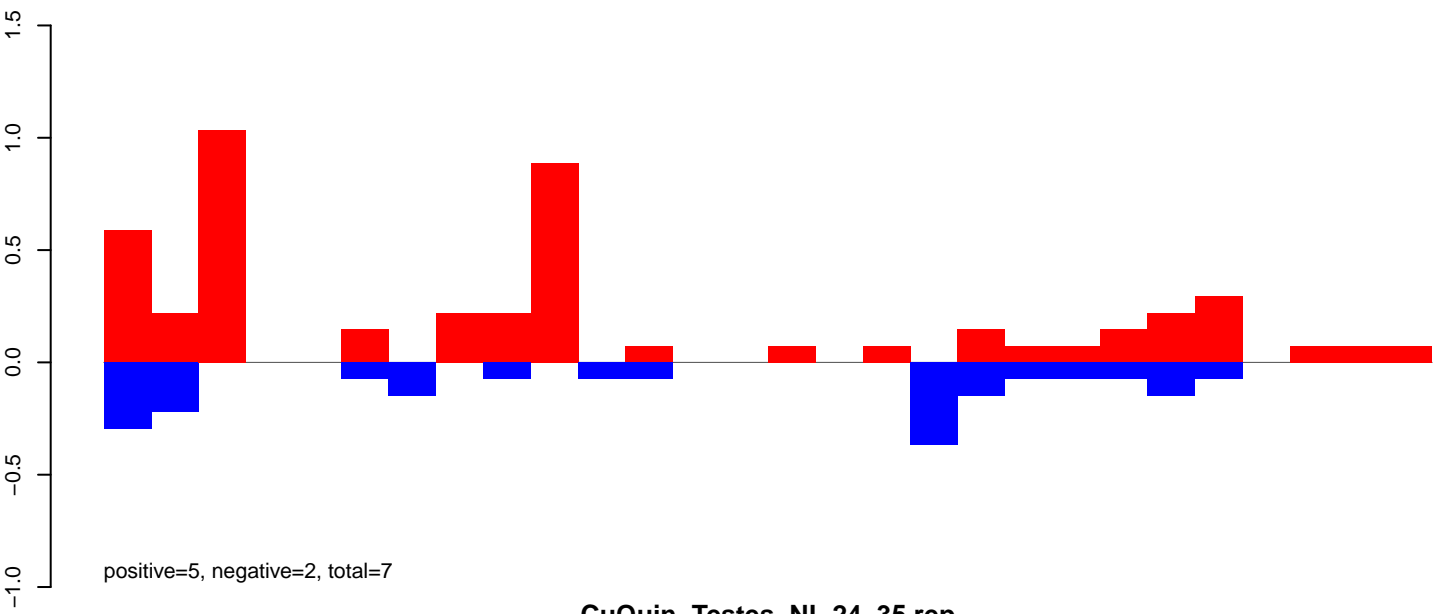
CuQuin_Testes_NL.24_35.rep



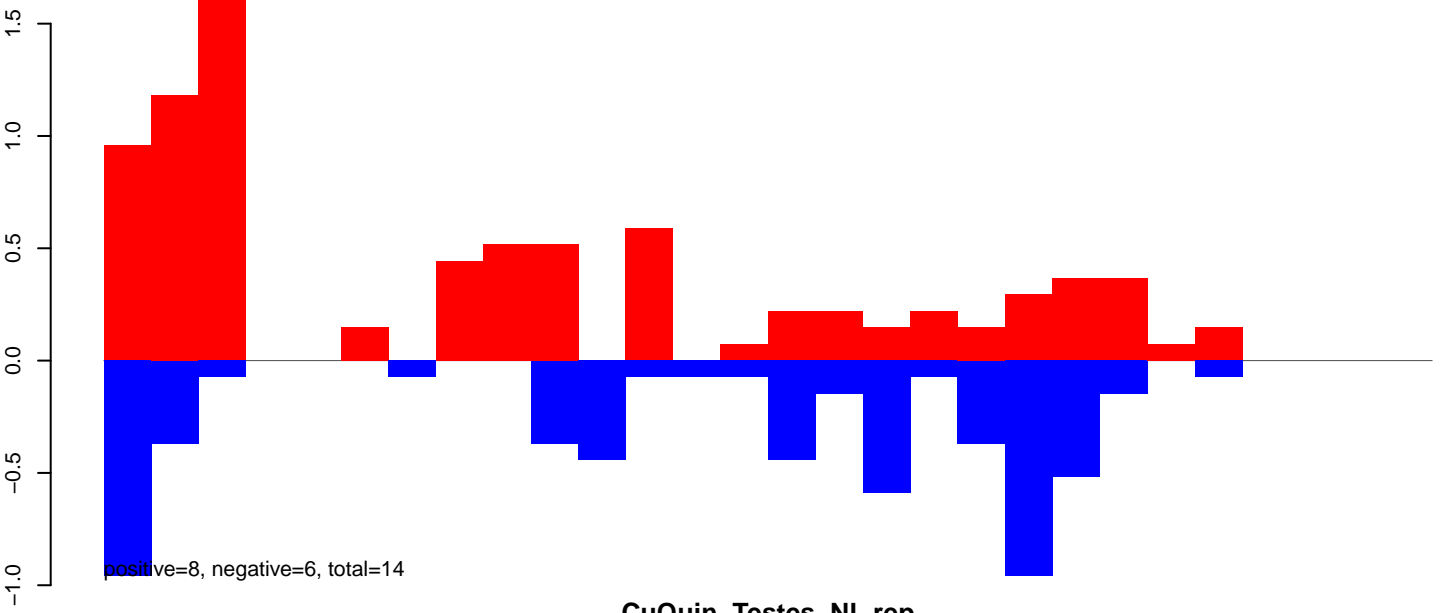
CuQuin_Testes_NL.rep



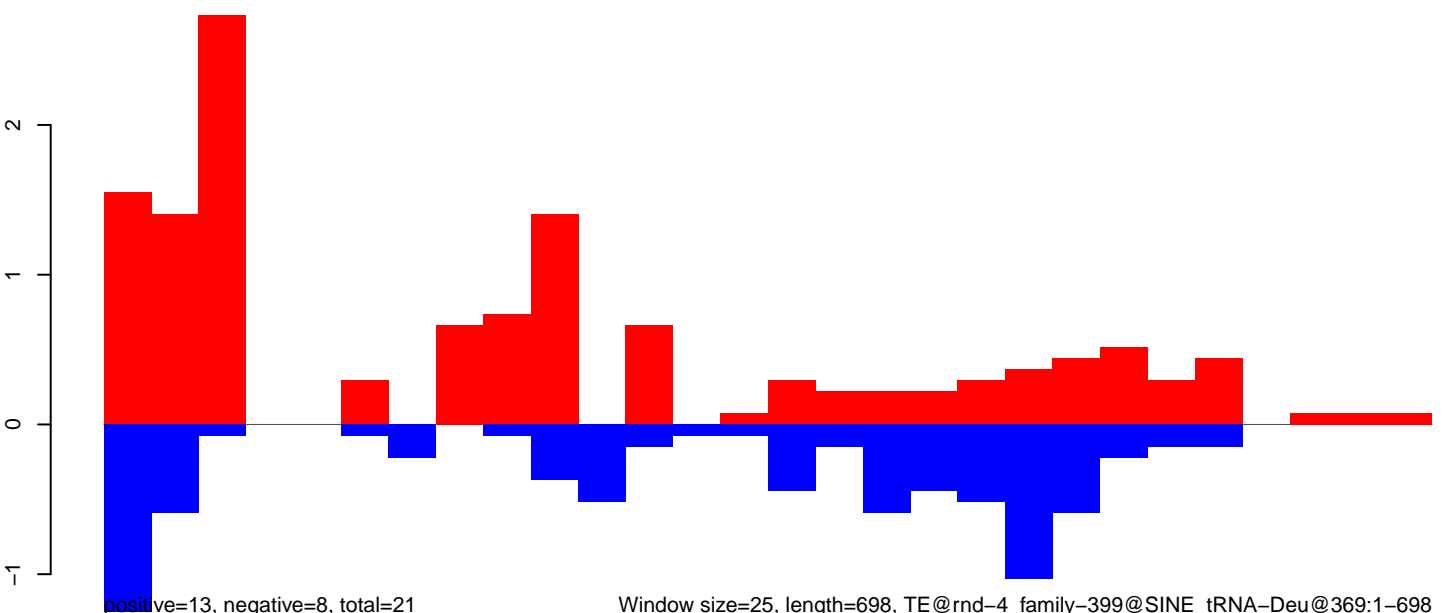
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

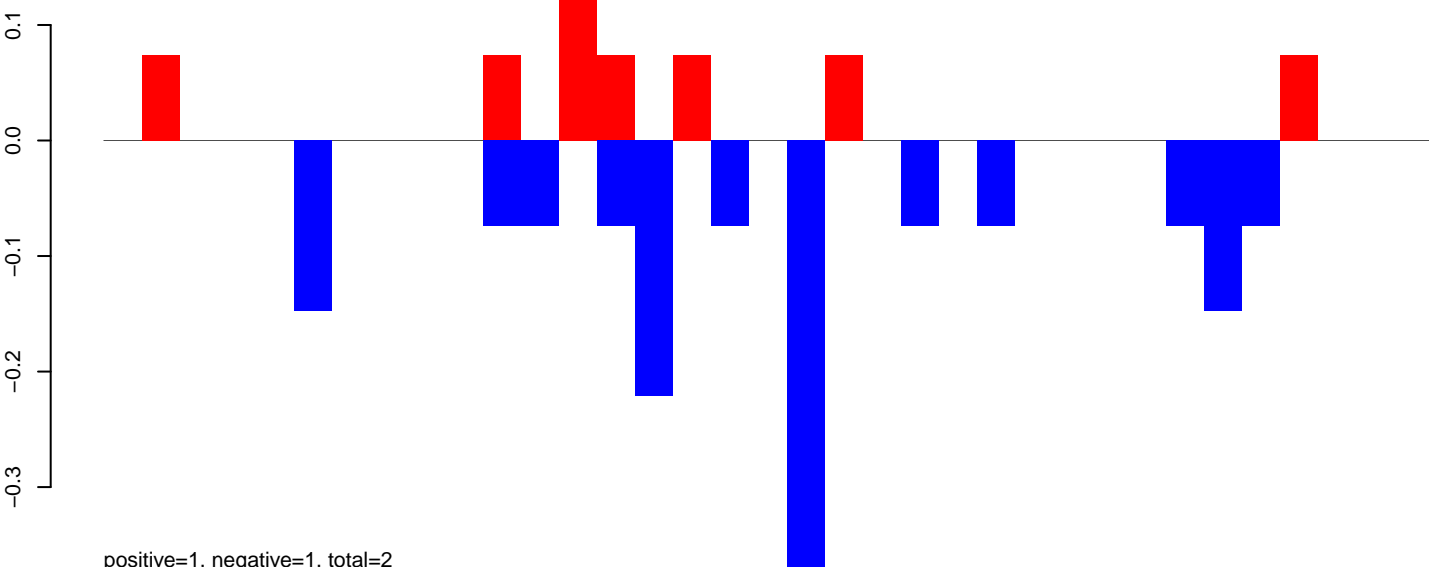


CuQuin_Testes_NL.rep

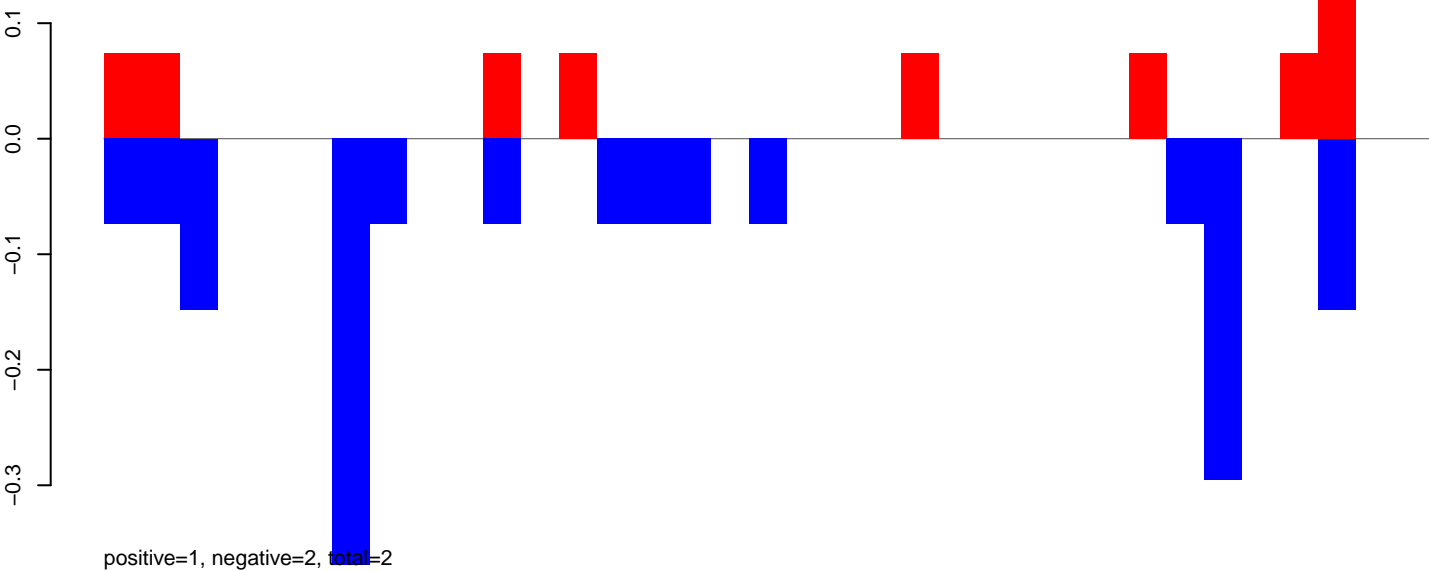


Window size=25, length=698, TE@rnd-4_family-399@SINE_tRNA-Deu@369:1-698

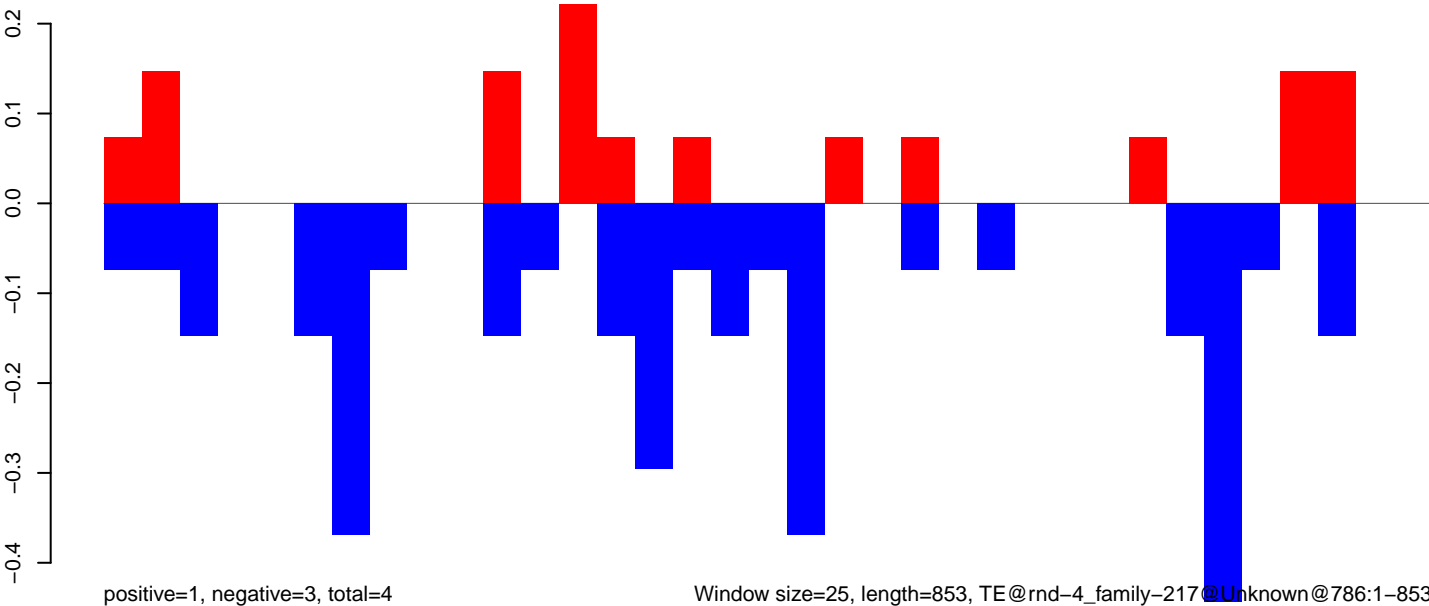
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

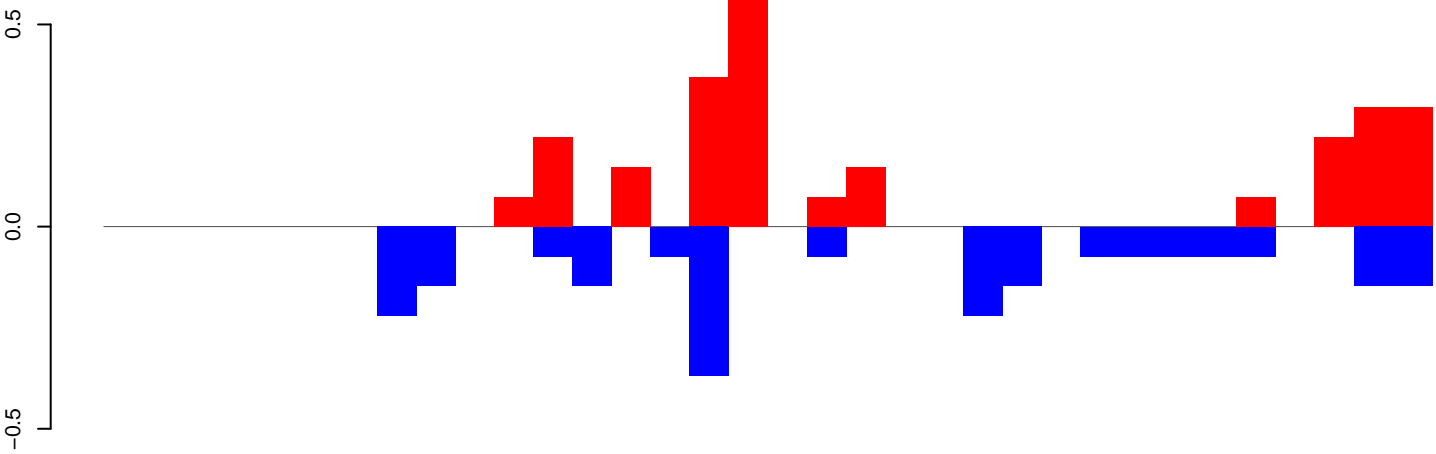


CuQuin_Testes_NL.rep



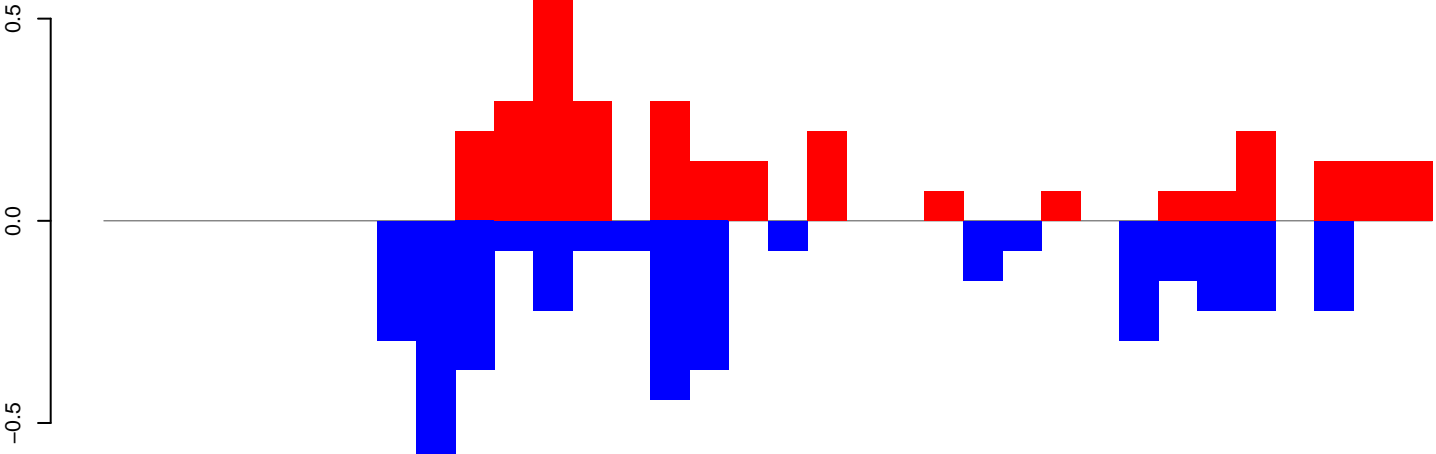
Window size=25, length=853, TE@rnd-4_family-217@Unknown@786:1-853

CuQuin_Testes_NL.18_23.rep



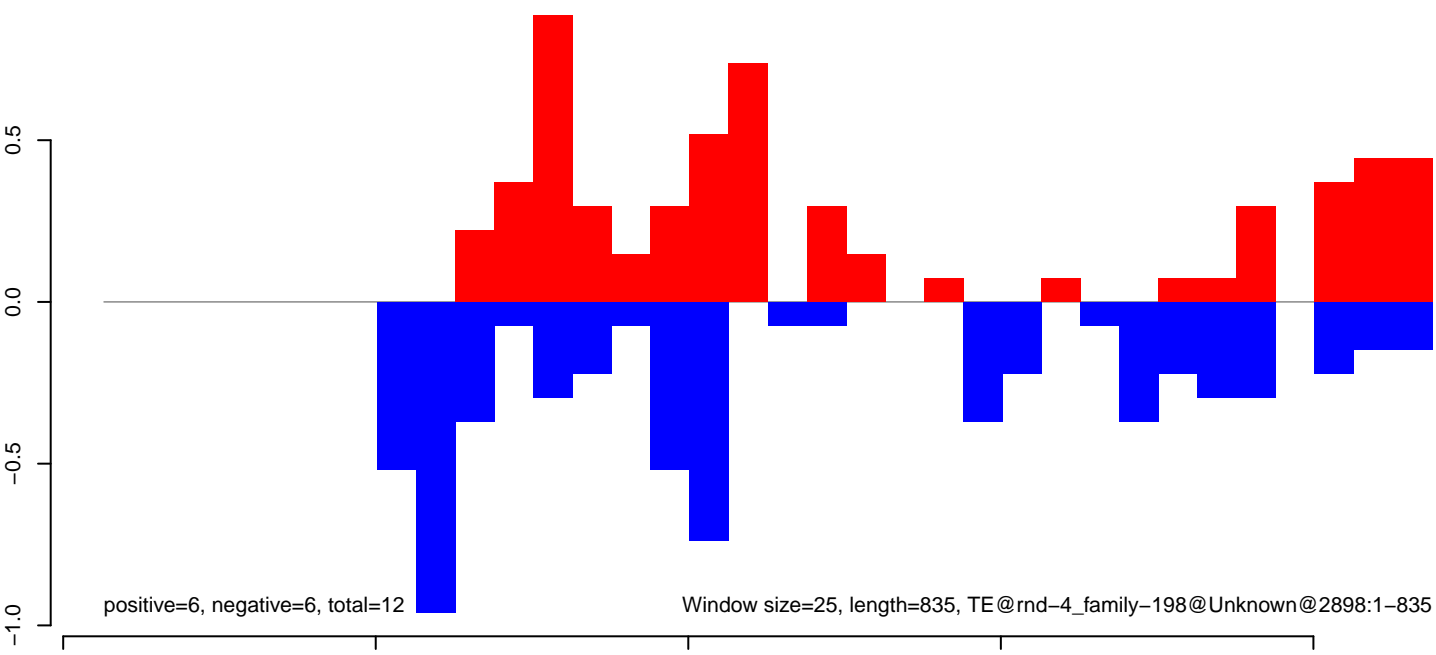
positive=3, negative=2, total=5

CuQuin_Testes_NL.24_35.rep



positive=3, negative=4, total=7

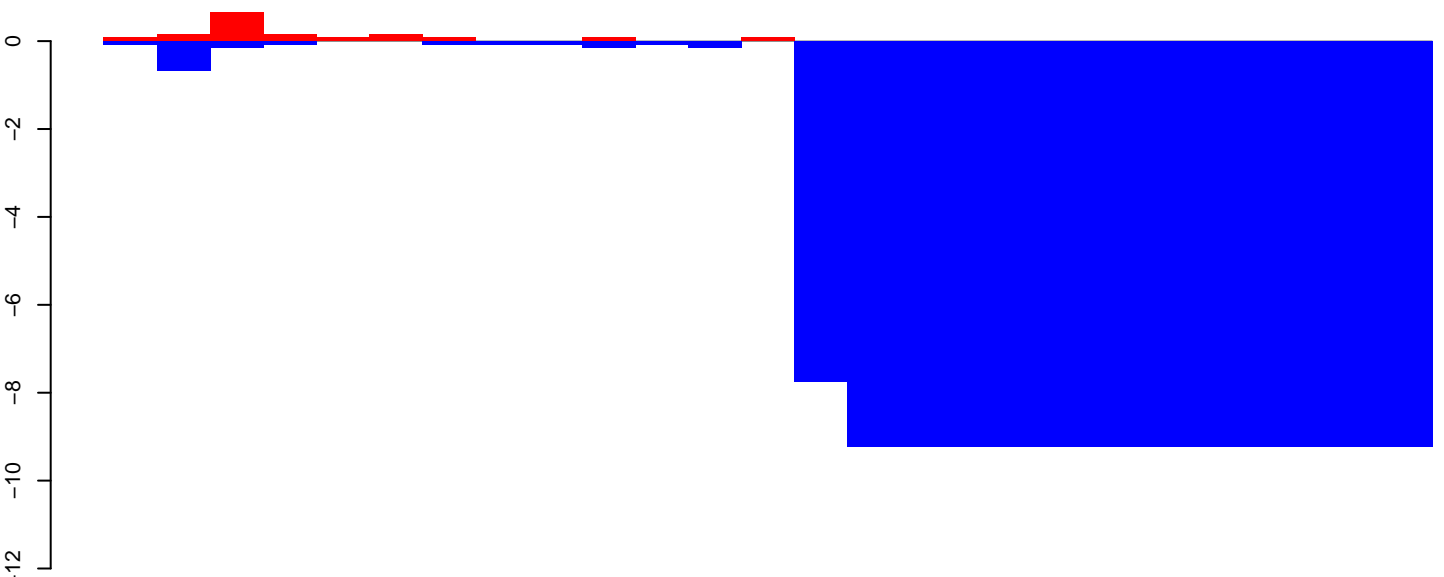
CuQuin_Testes_NL.rep



positive=6, negative=6, total=12

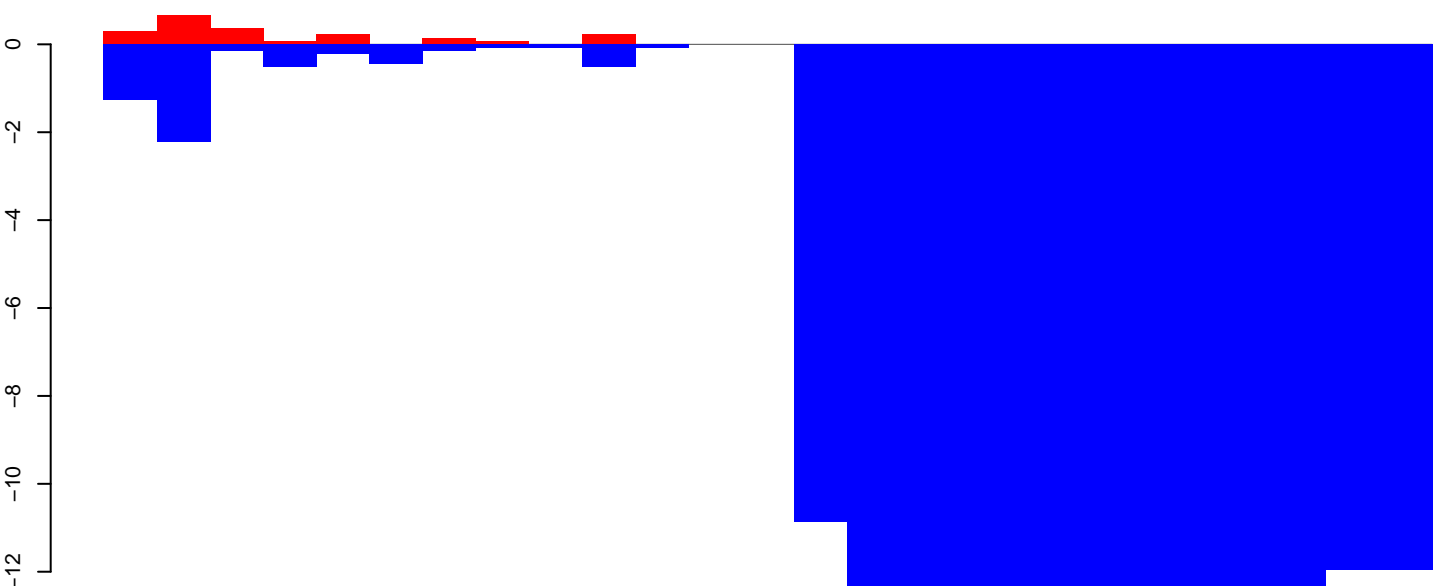
Window size=25, length=835, TE@rnd-4_family-198@Unknown@2898:1-835

CuQuin_Testes_NL.18_23.rep



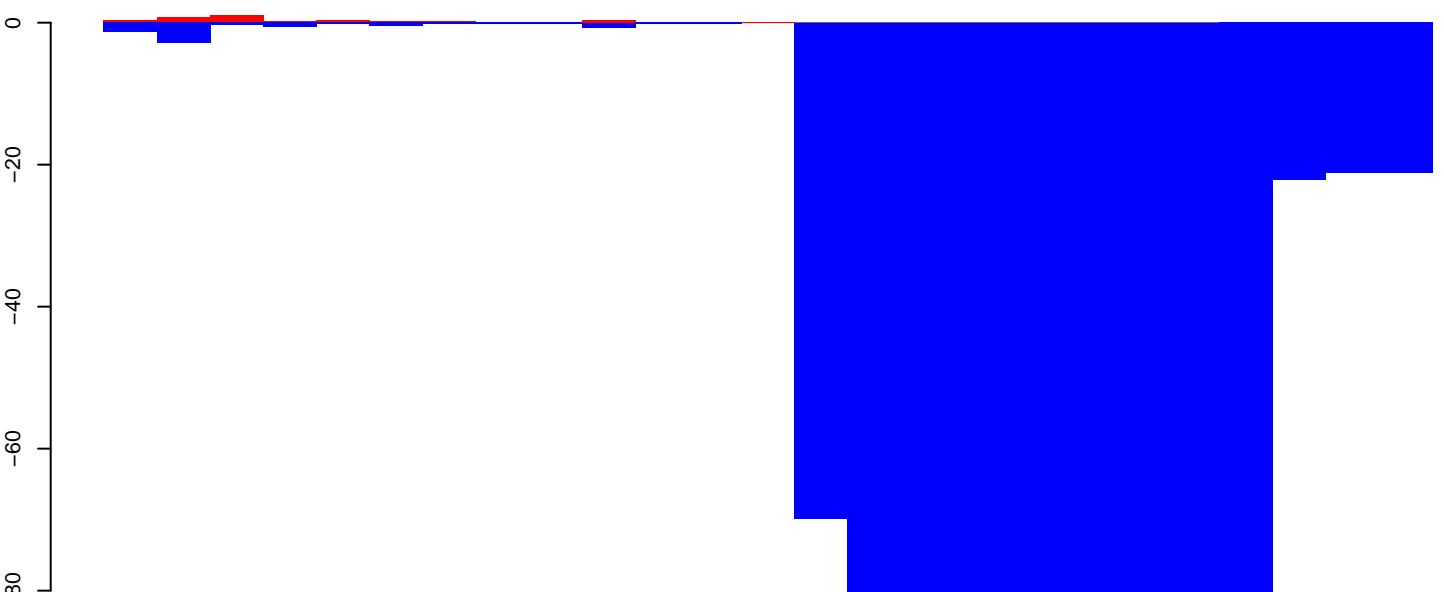
positive=1, negative=111, total=112

CuQuin_Testes_NL.24_35.rep



positive=2, negative=157, total=159

CuQuin_Testes_NL.rep

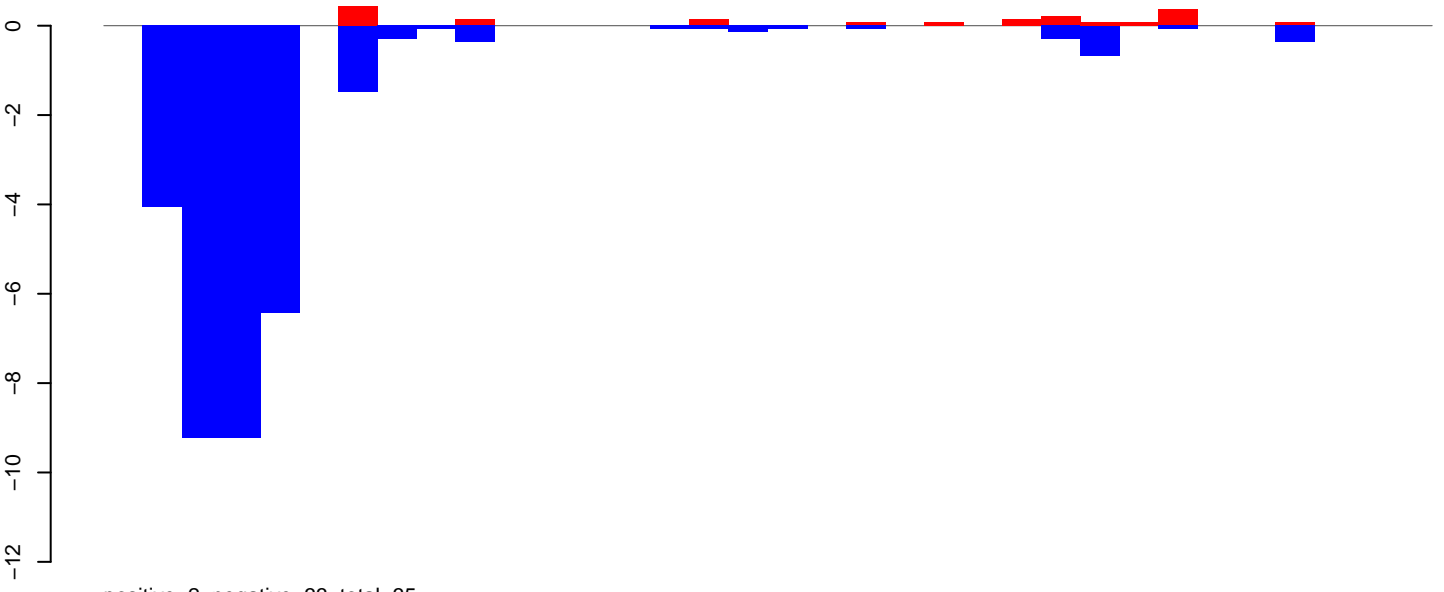


positive=4, negative=804, total=807

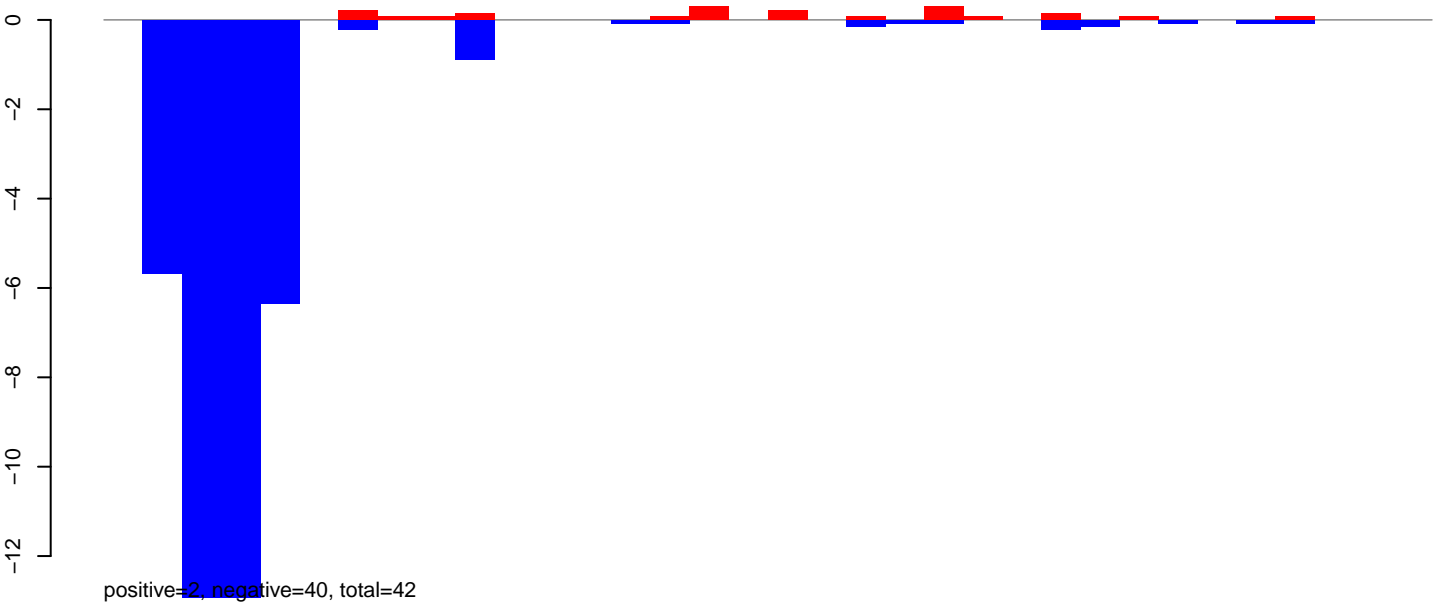
Window size=25, length=625, TE@rnd-3_family-773@LINE_1-Jockey@1981:1-625

100 200 300 400 500 600

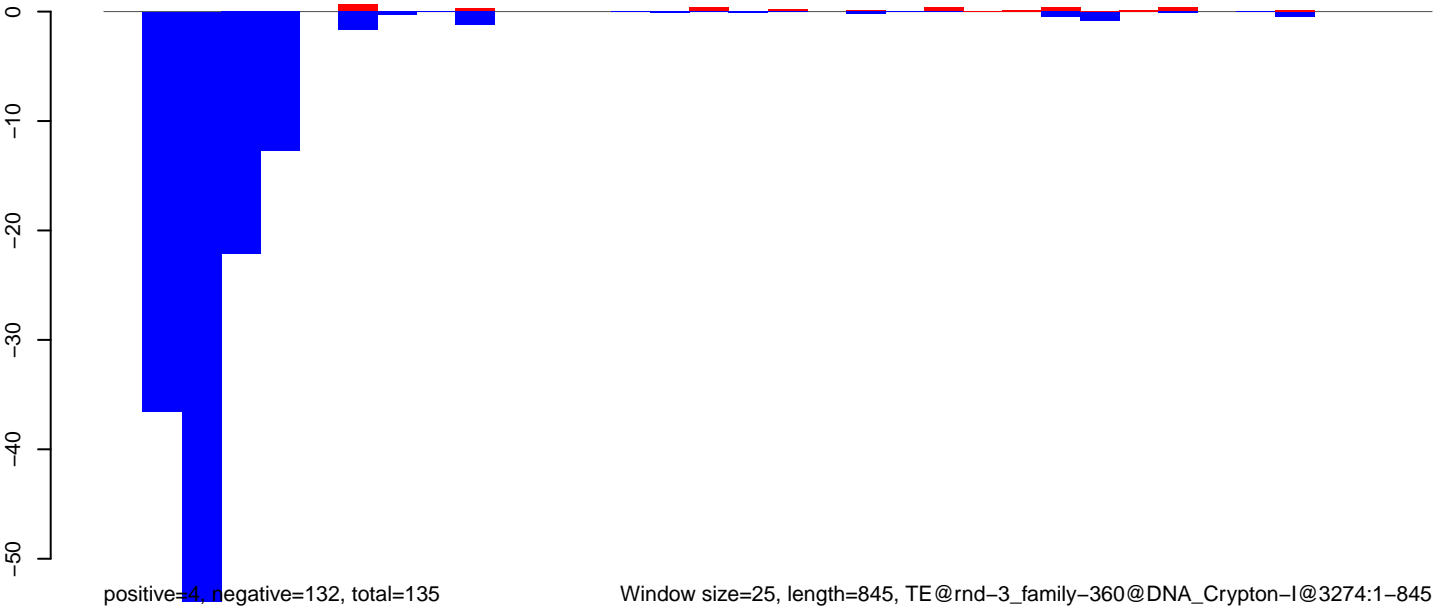
CuQuin_Testes_NL.18_23.rep



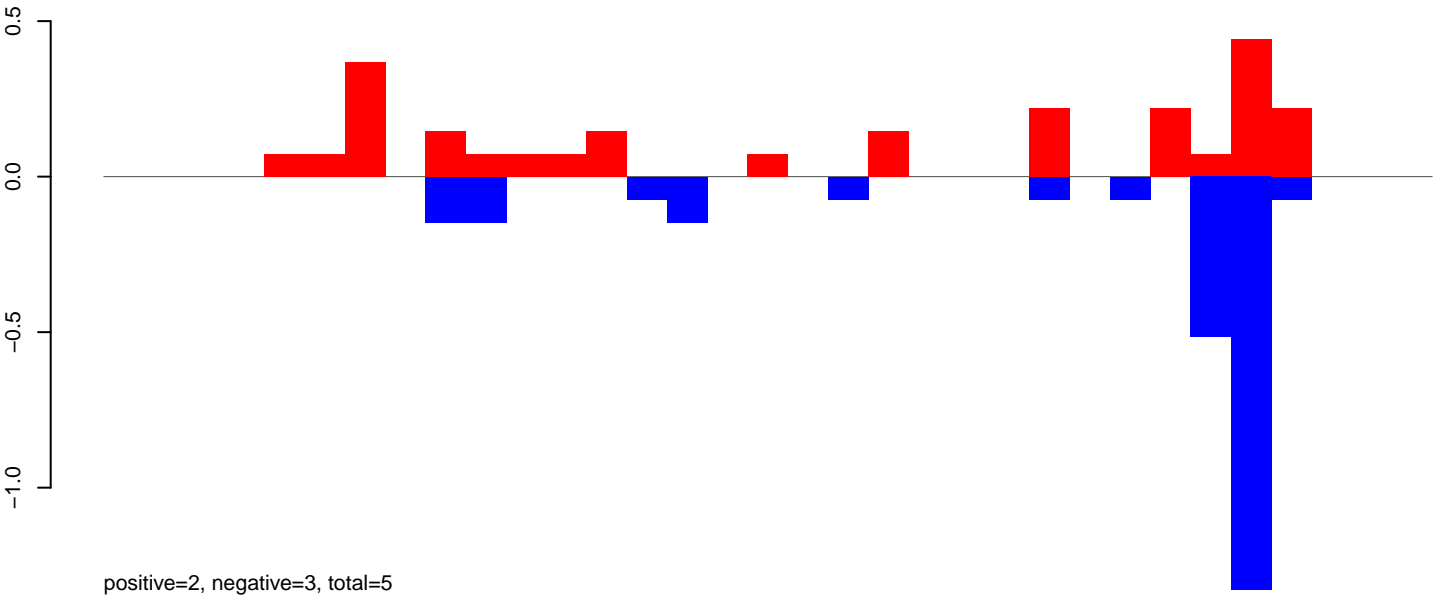
CuQuin_Testes_NL.24_35.rep



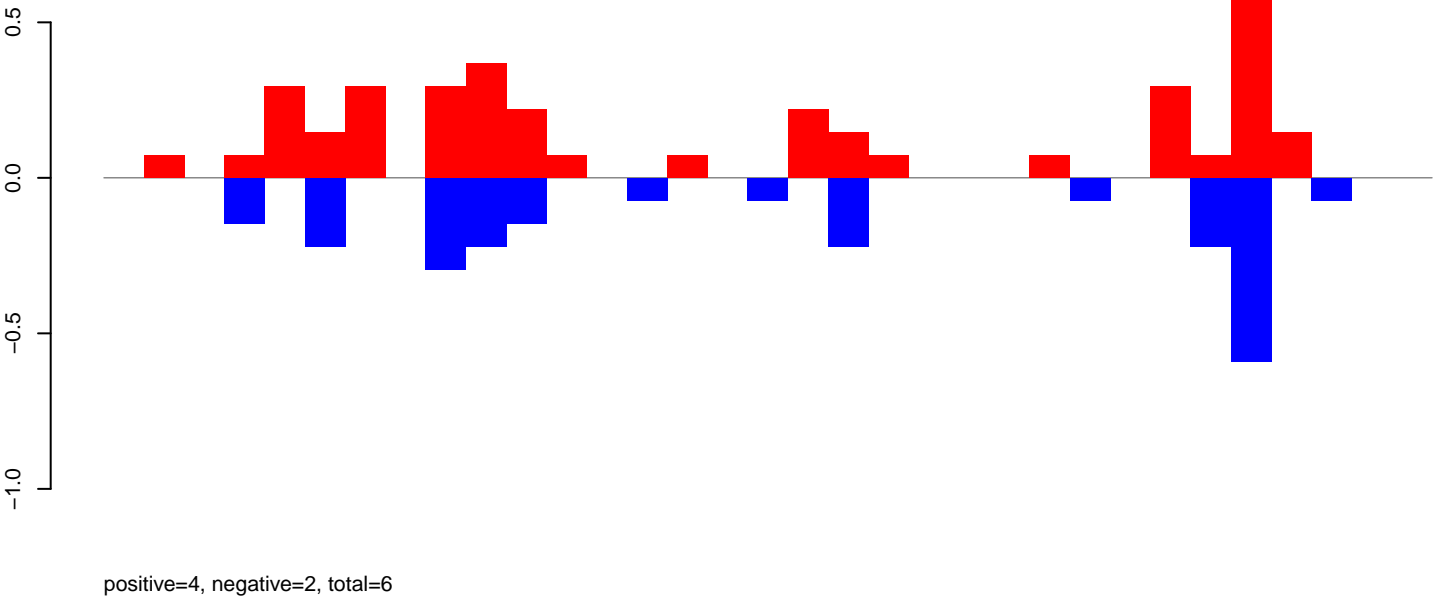
CuQuin_Testes_NL.rep



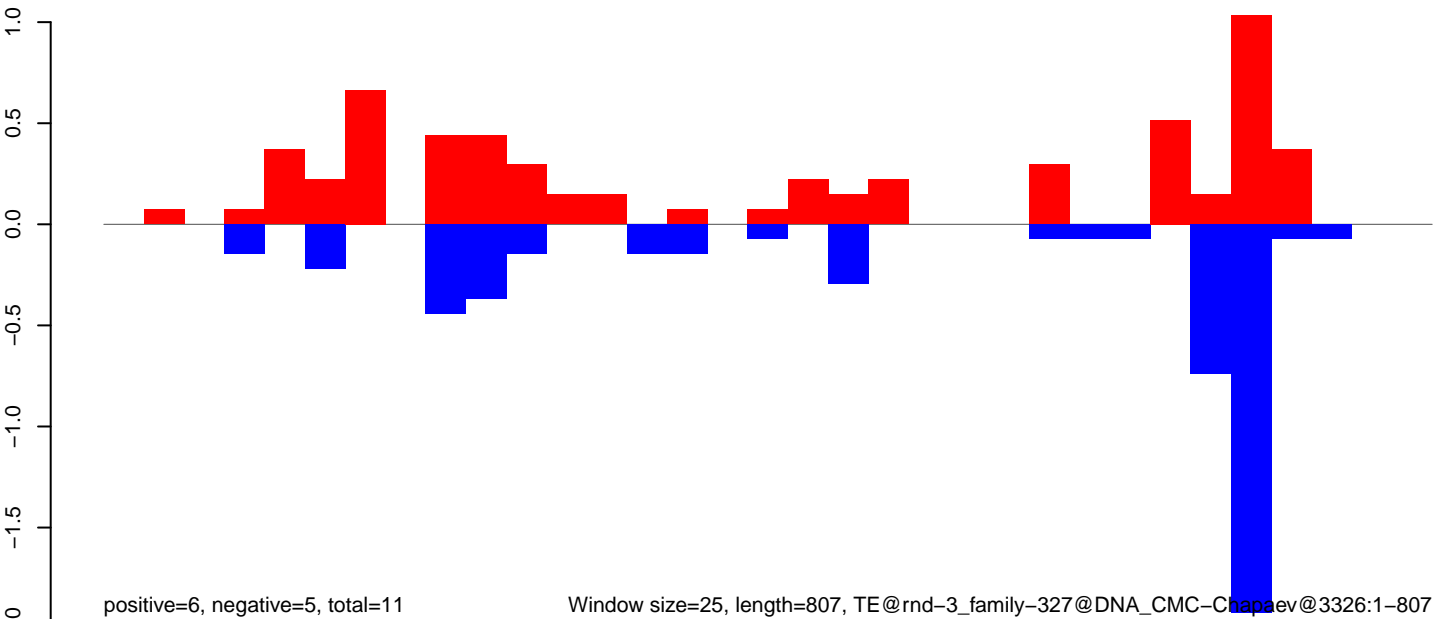
CuQuin_Testes_NL.18_23.rep



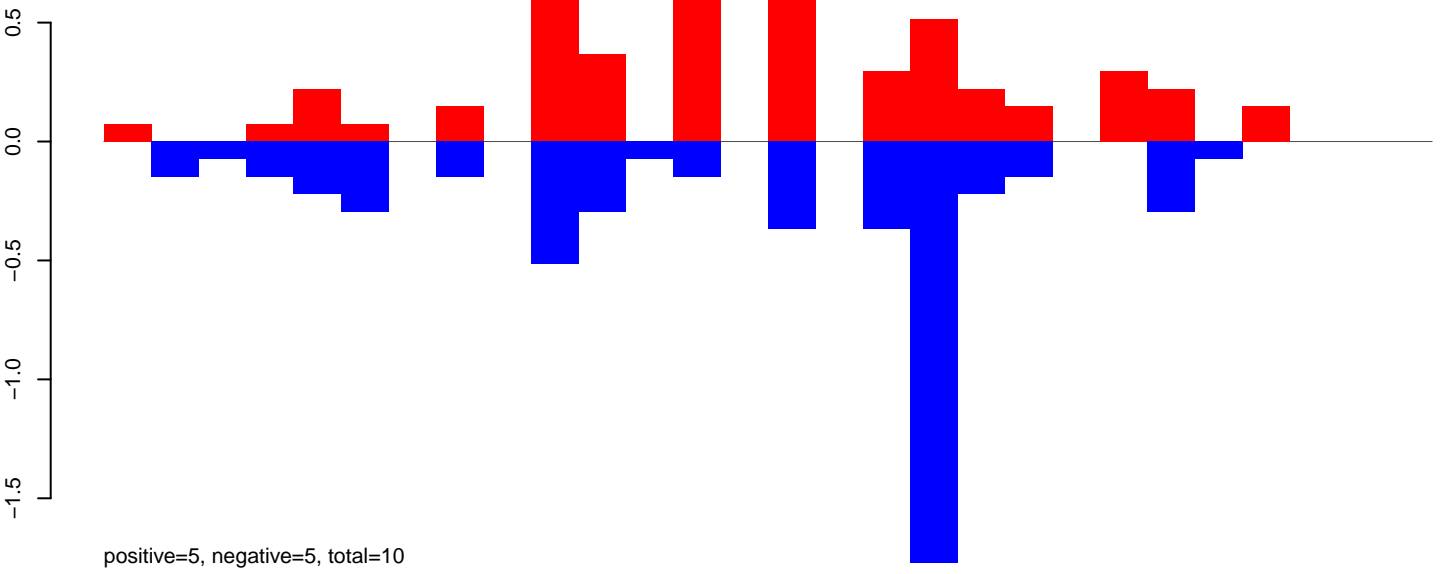
CuQuin_Testes_NL.24_35.rep



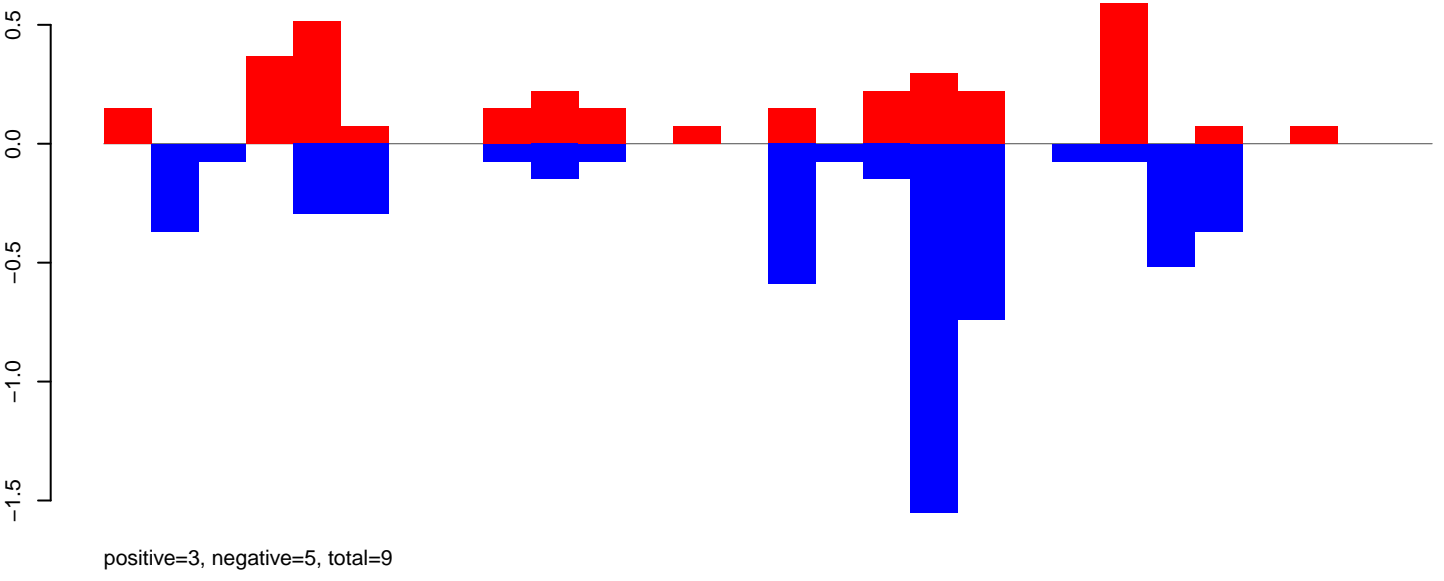
CuQuin_Testes_NL.rep



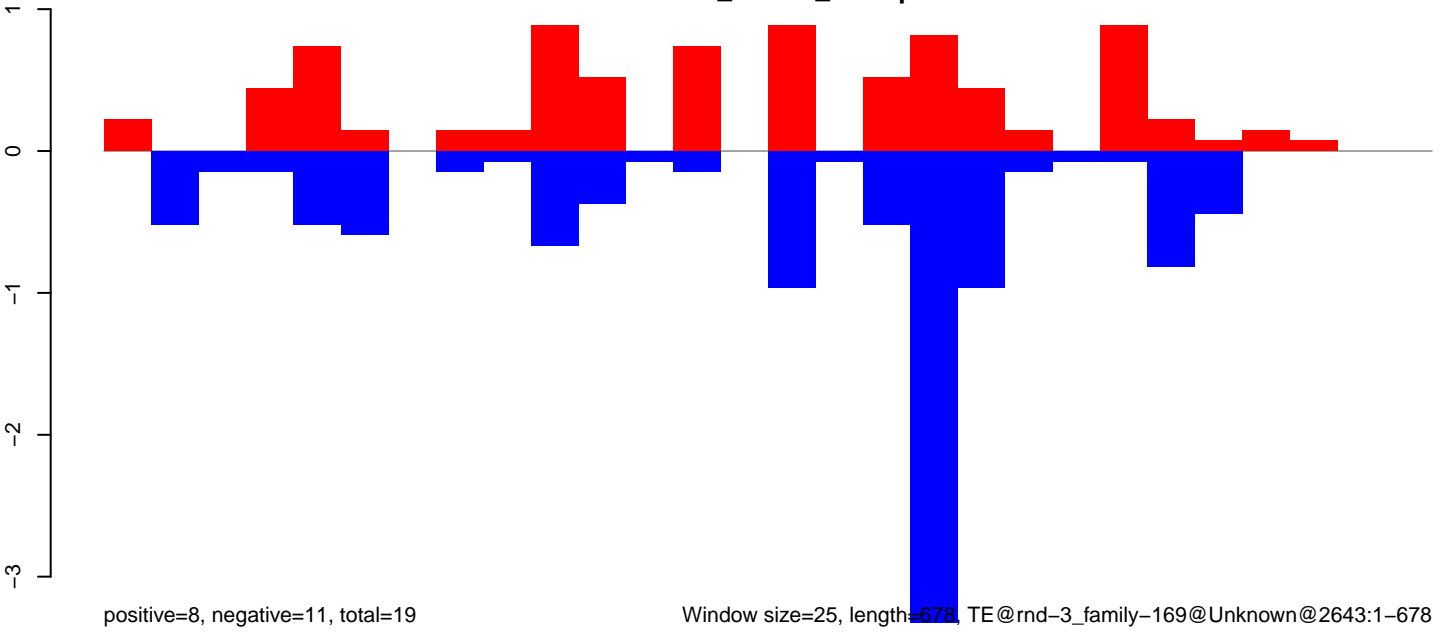
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

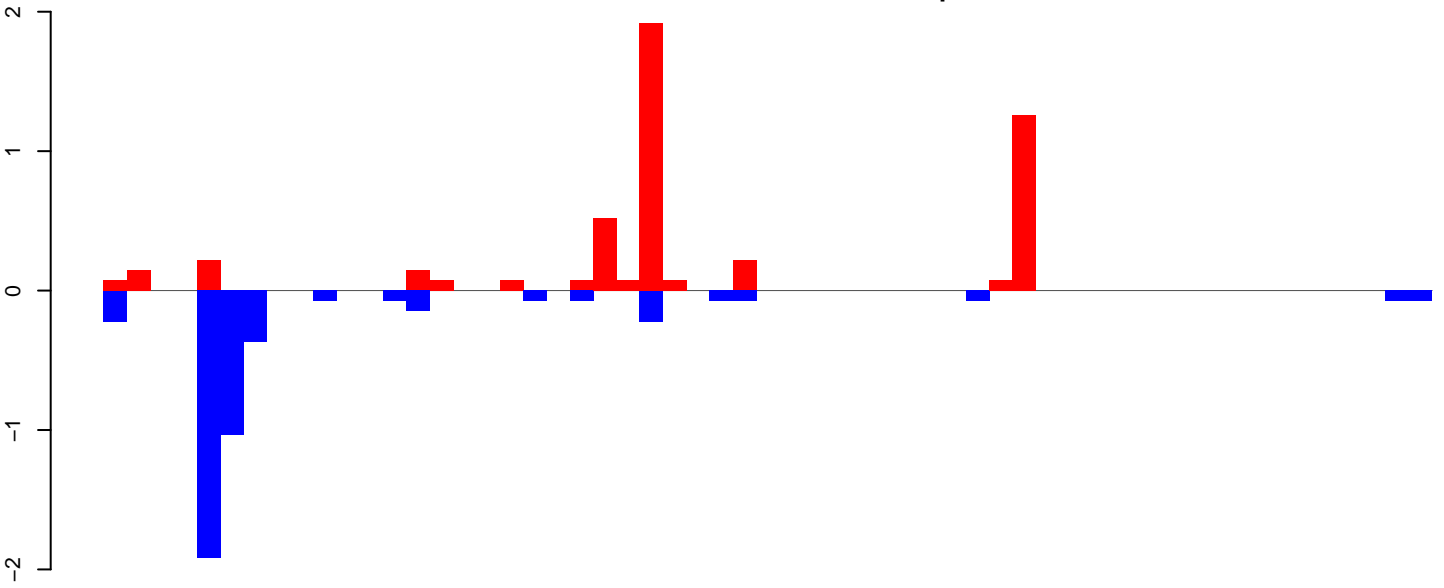


CuQuin_Testes_NL.rep



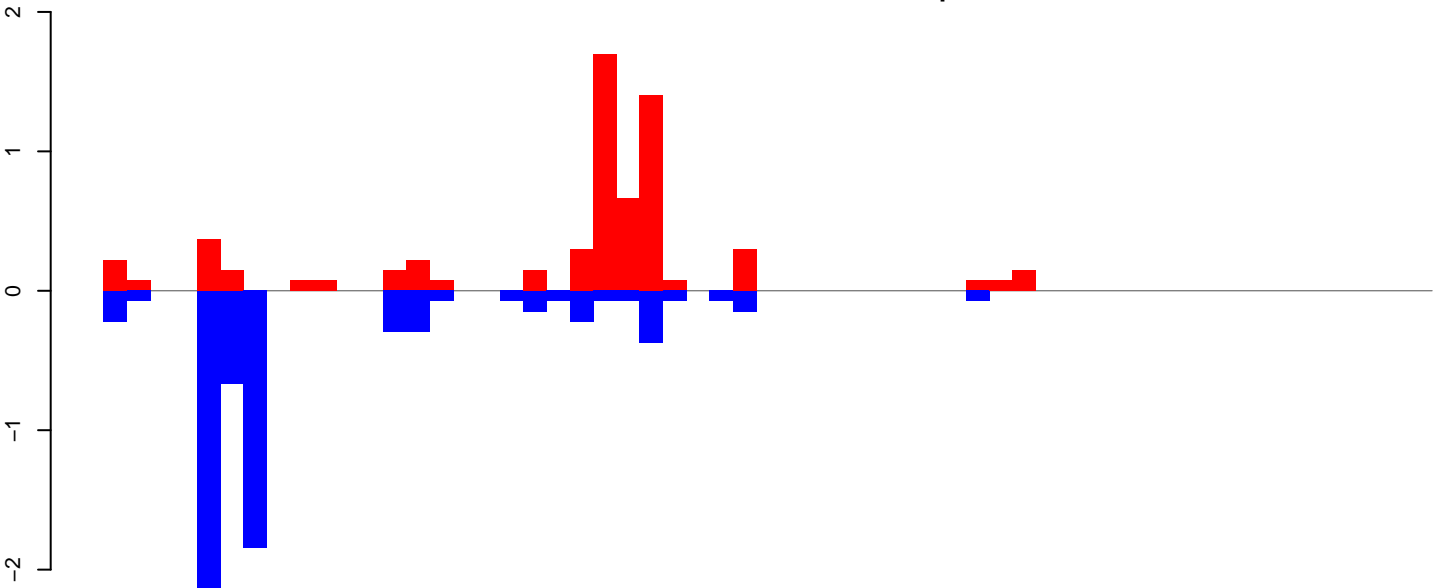
Window size=25, length=678, TE@rnd-3_family-169@Unknown@2643:1-678

CuQuin_Testes_NL.18_23.rep



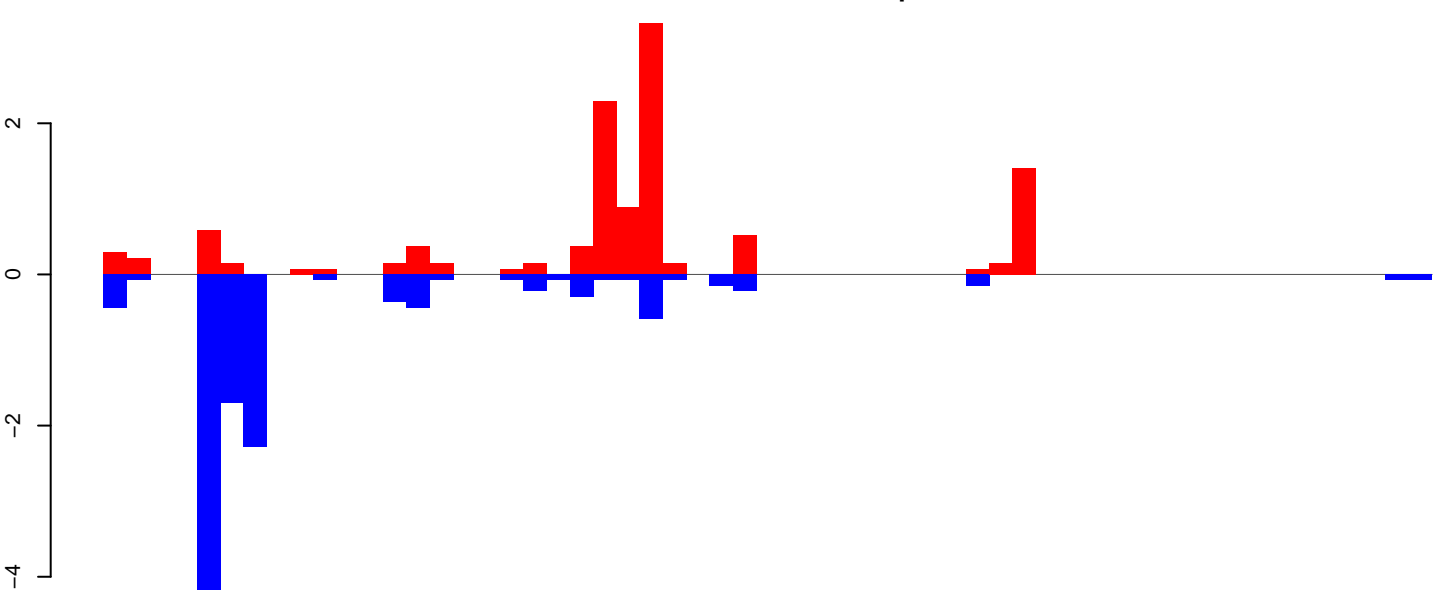
positive=5, negative=5, total=10

CuQuin_Testes_NL.24_35.rep



positive=6, negative=7, total=14

CuQuin_Testes_NL.rep

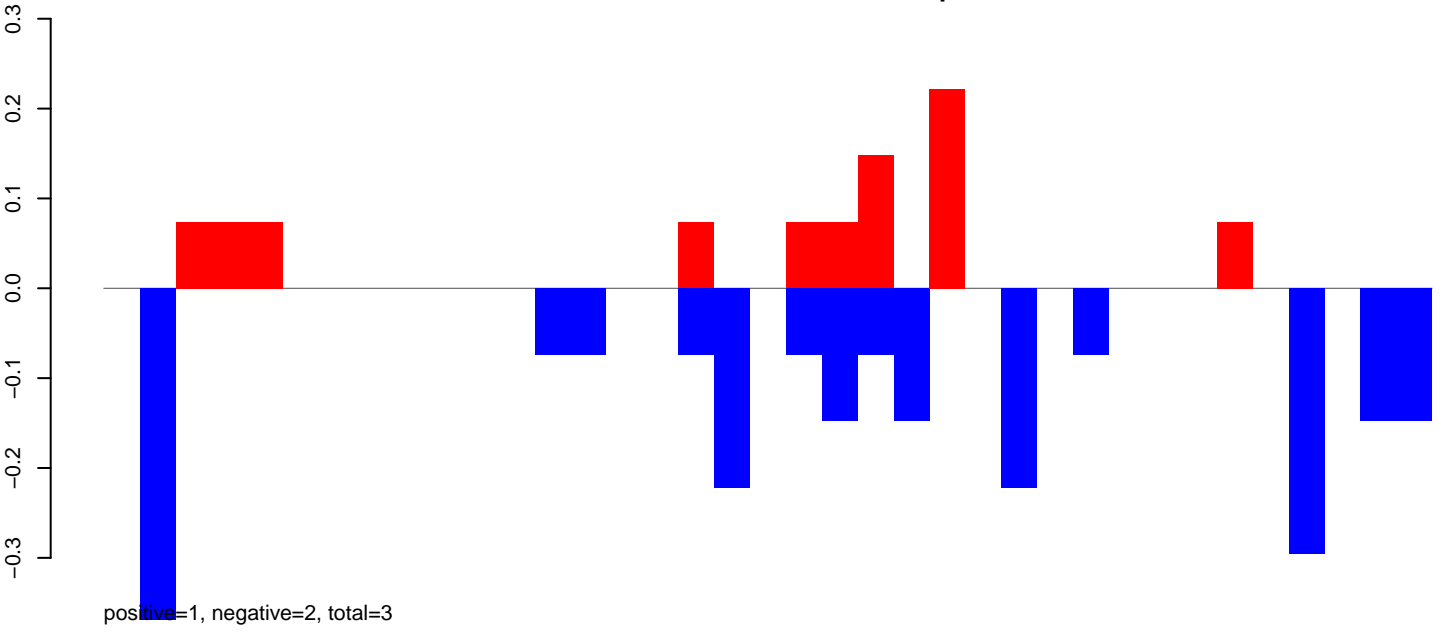


positive=11, negative=12, total=24

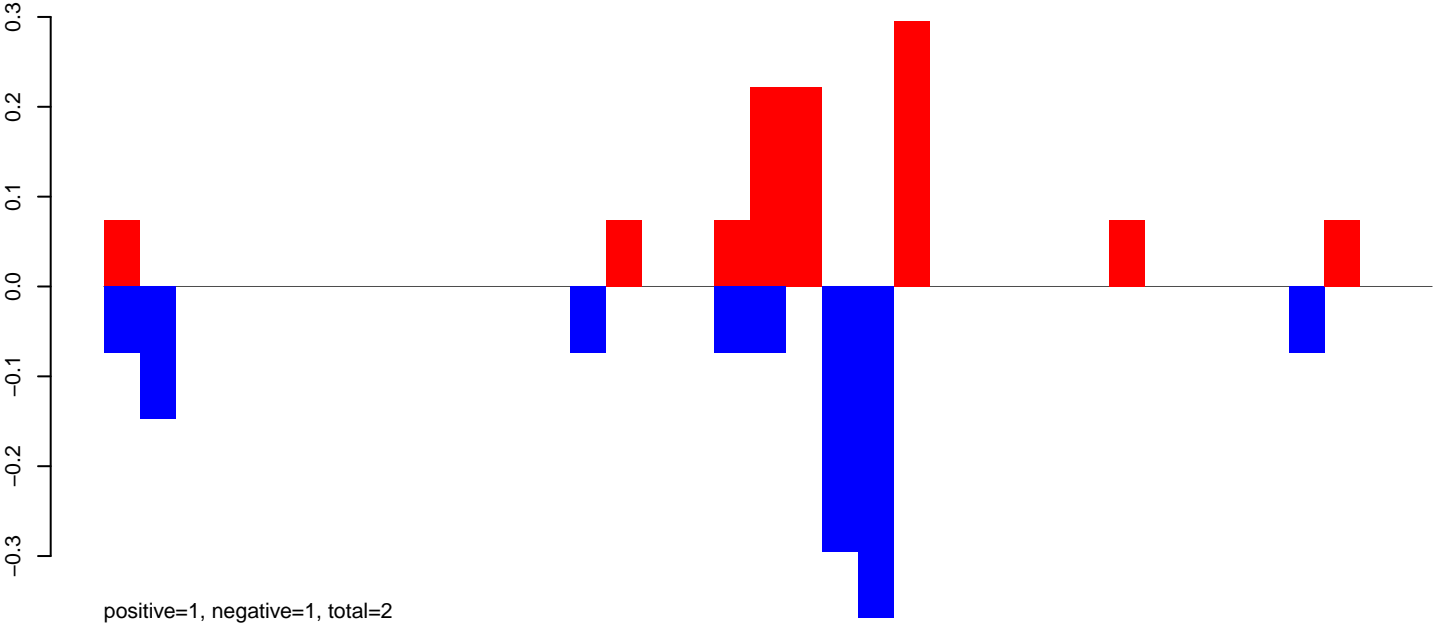
Window size=25, length=1411, TE@rnd-3_family-108@DNA_Crypton-I@1355:1-1411

0 200 400 600 800 1000 1200 1400

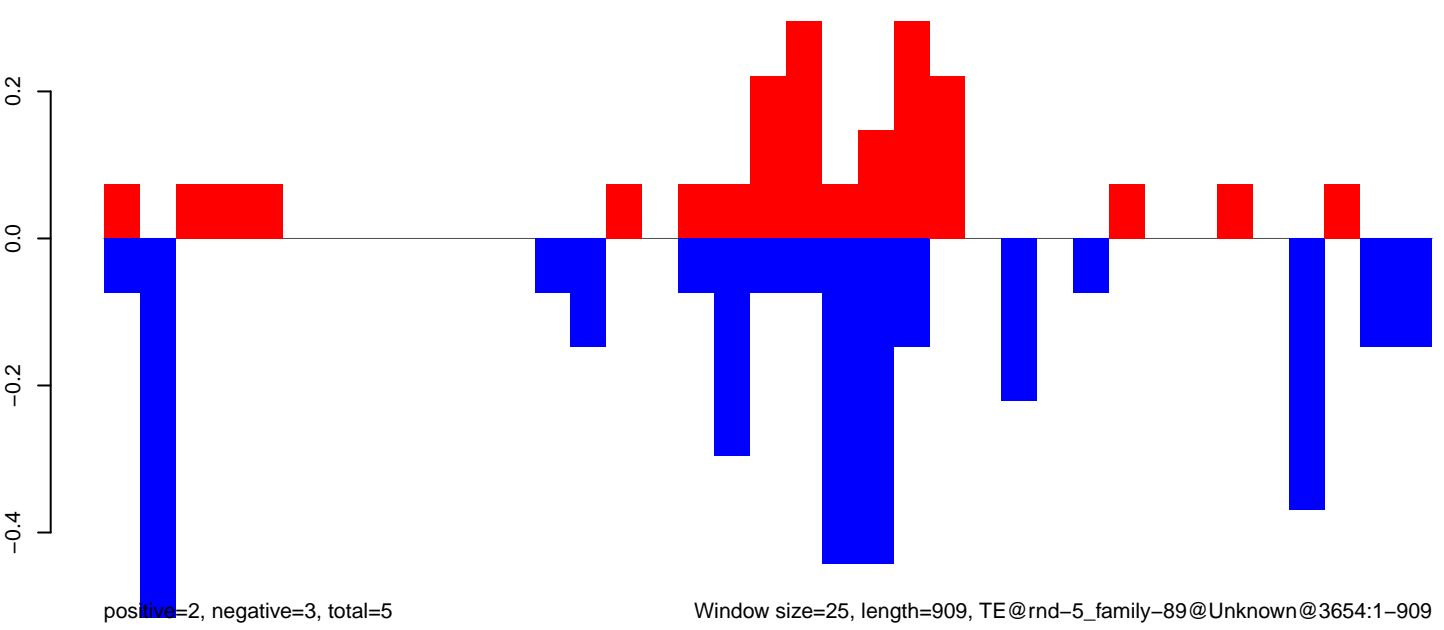
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

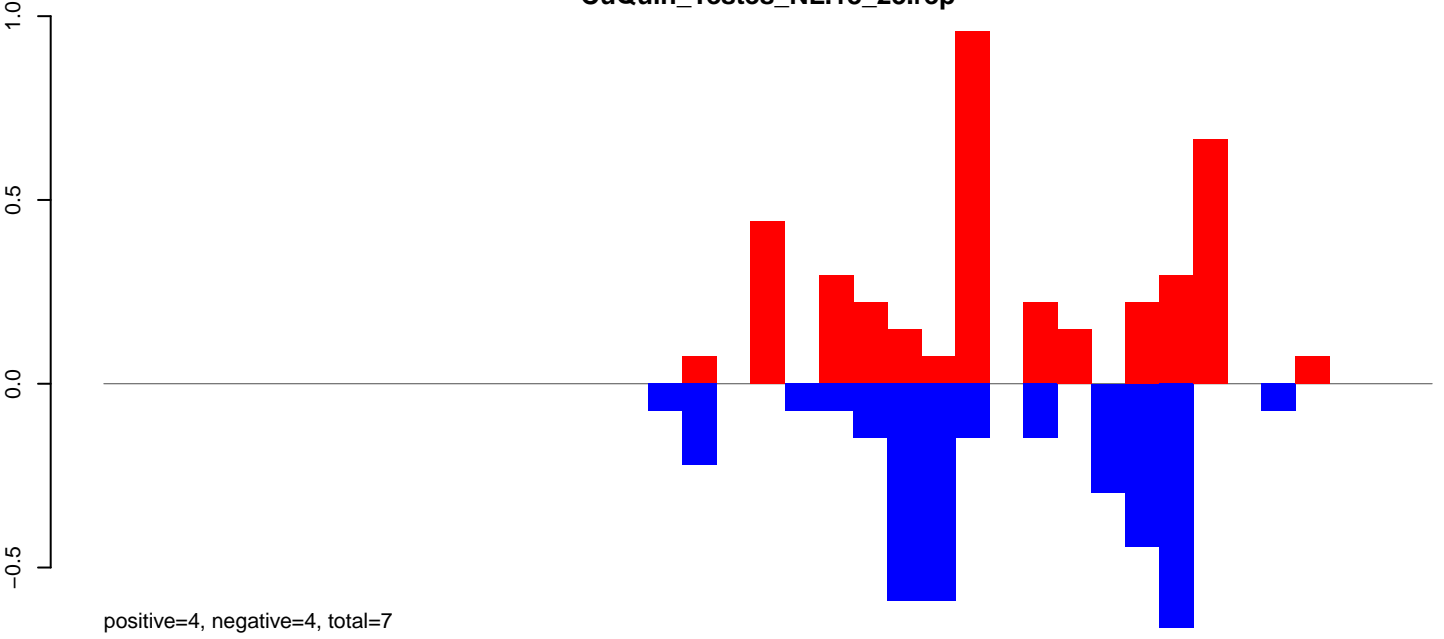


CuQuin_Testes_NL.rep

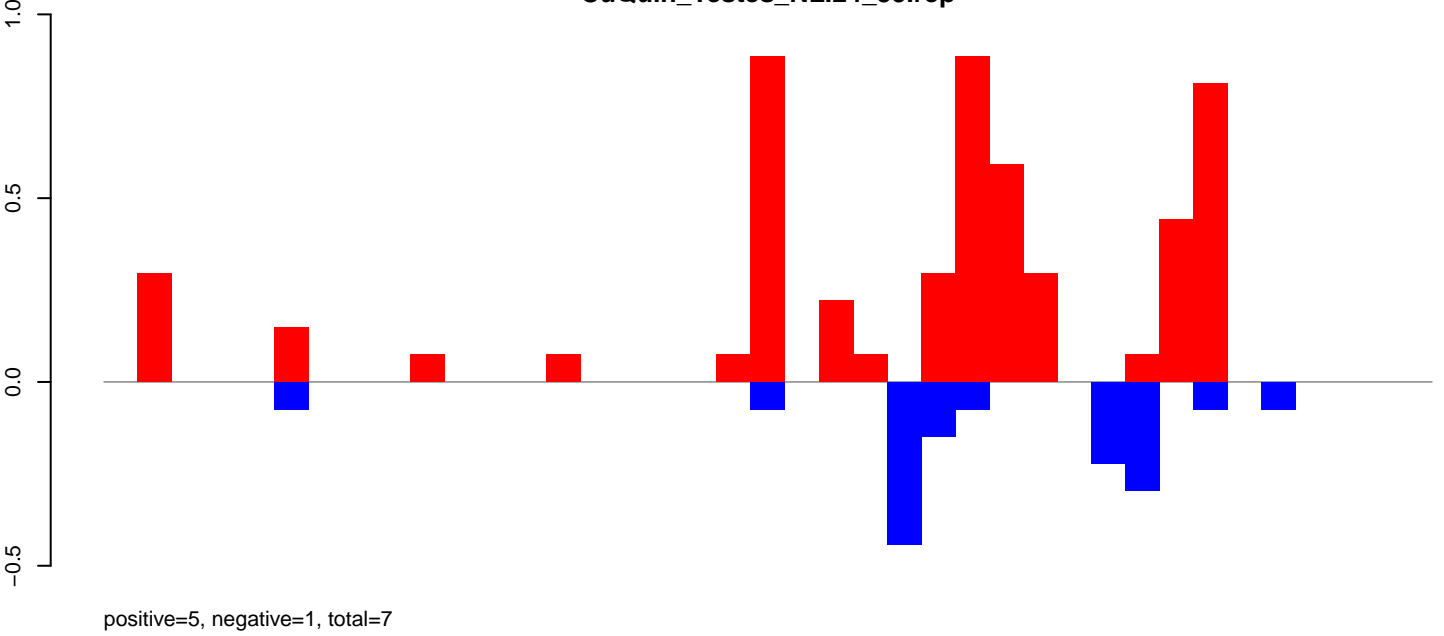


Window size=25, length=909, TE@rnd-5_family-89@Unknown@3654:1-909

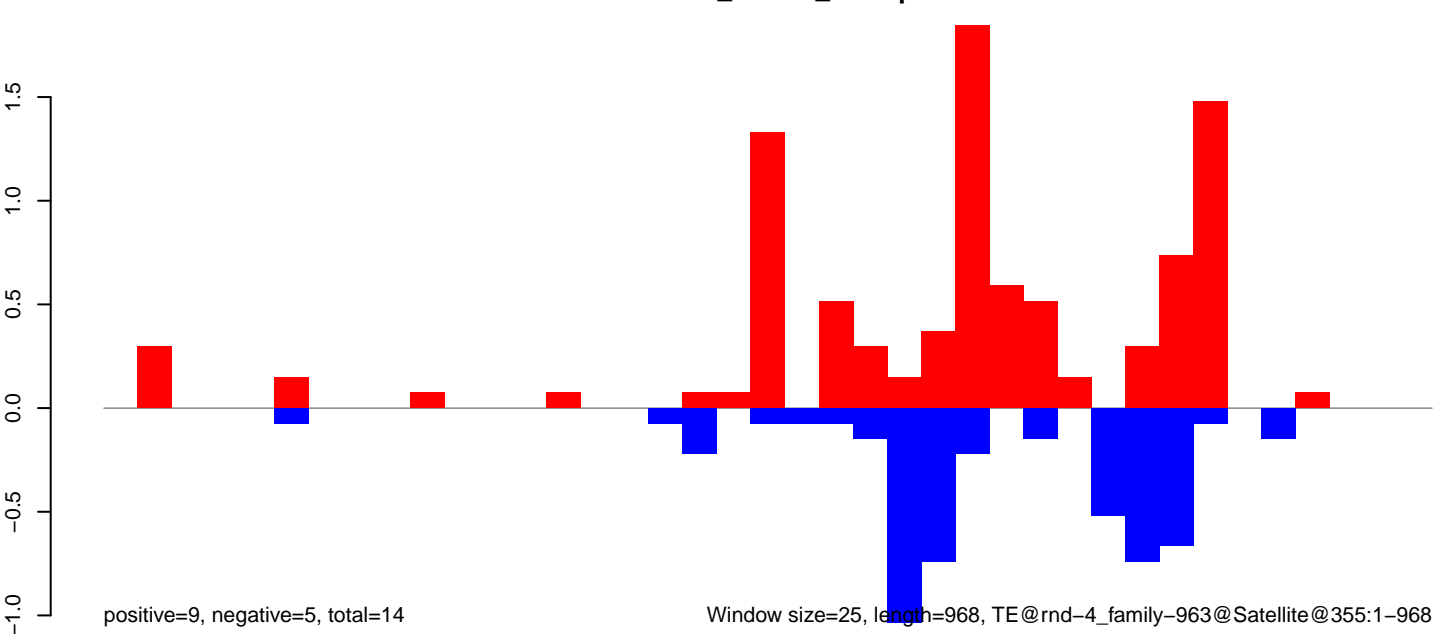
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

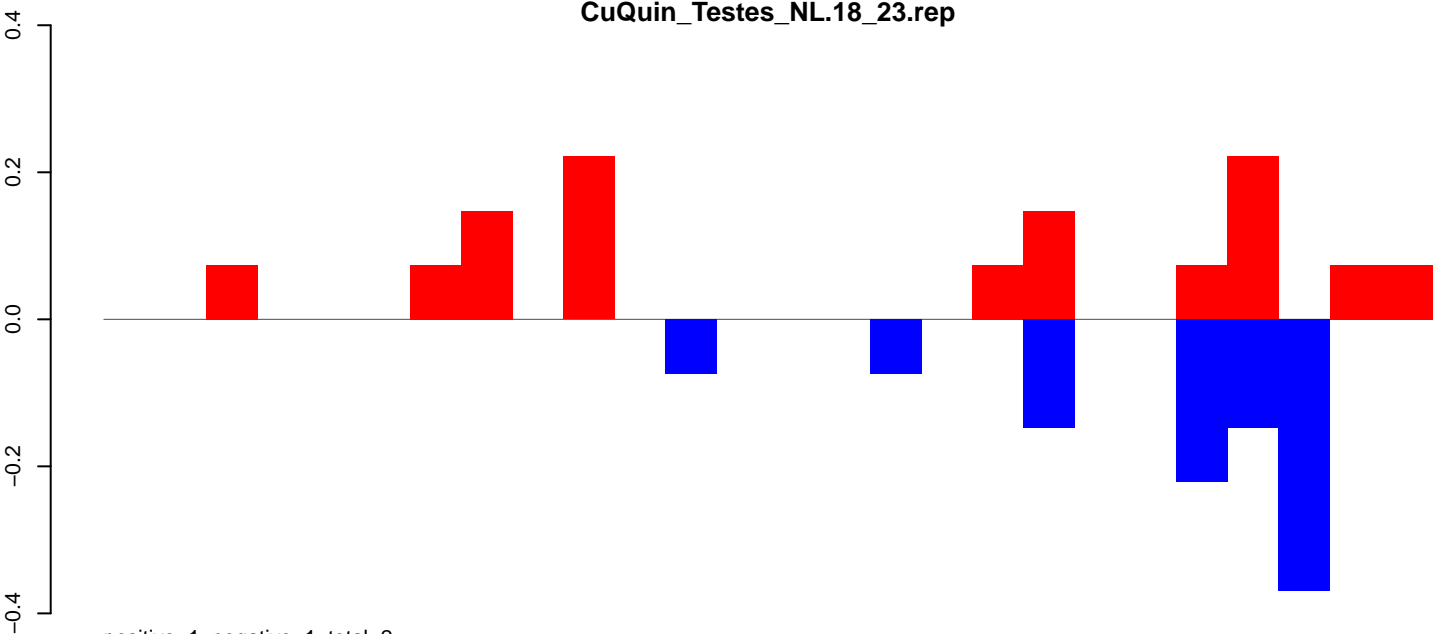


CuQuin_Testes_NL.rep



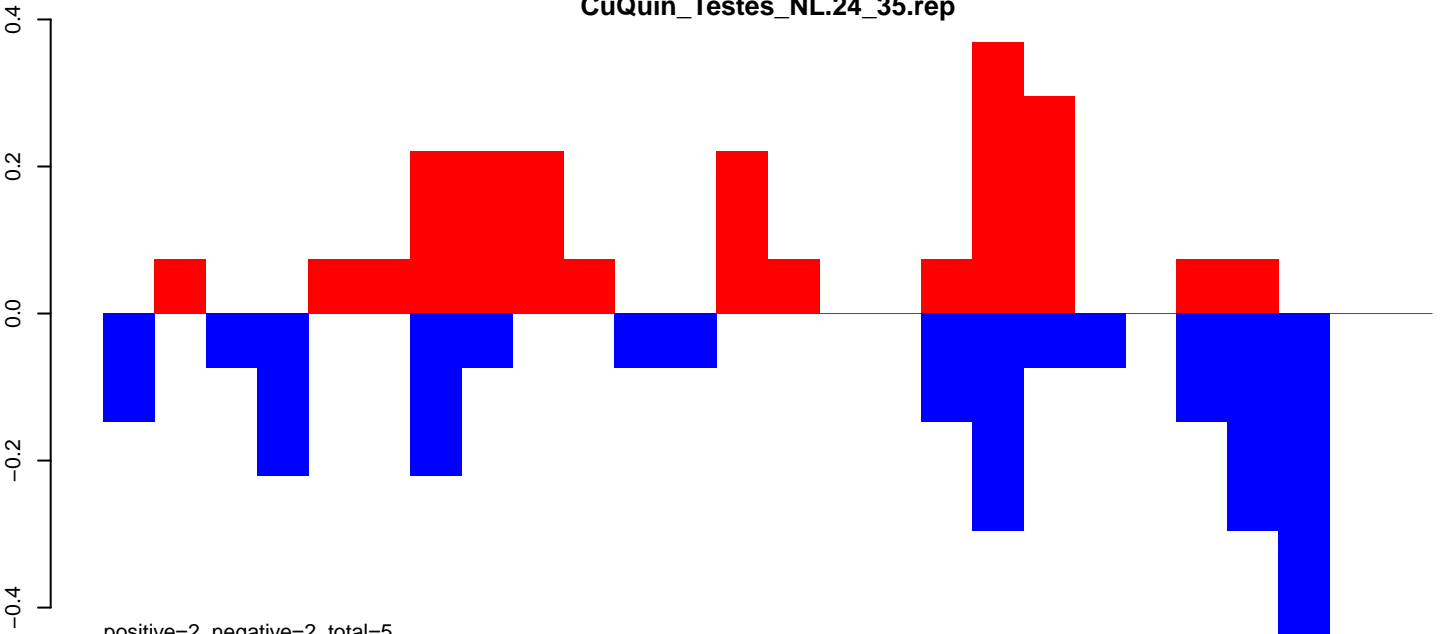
Window size=25, length=968, TE@rnd-4_family-963@Satellite@355:1-968

CuQuin_Testes_NL.18_23.rep



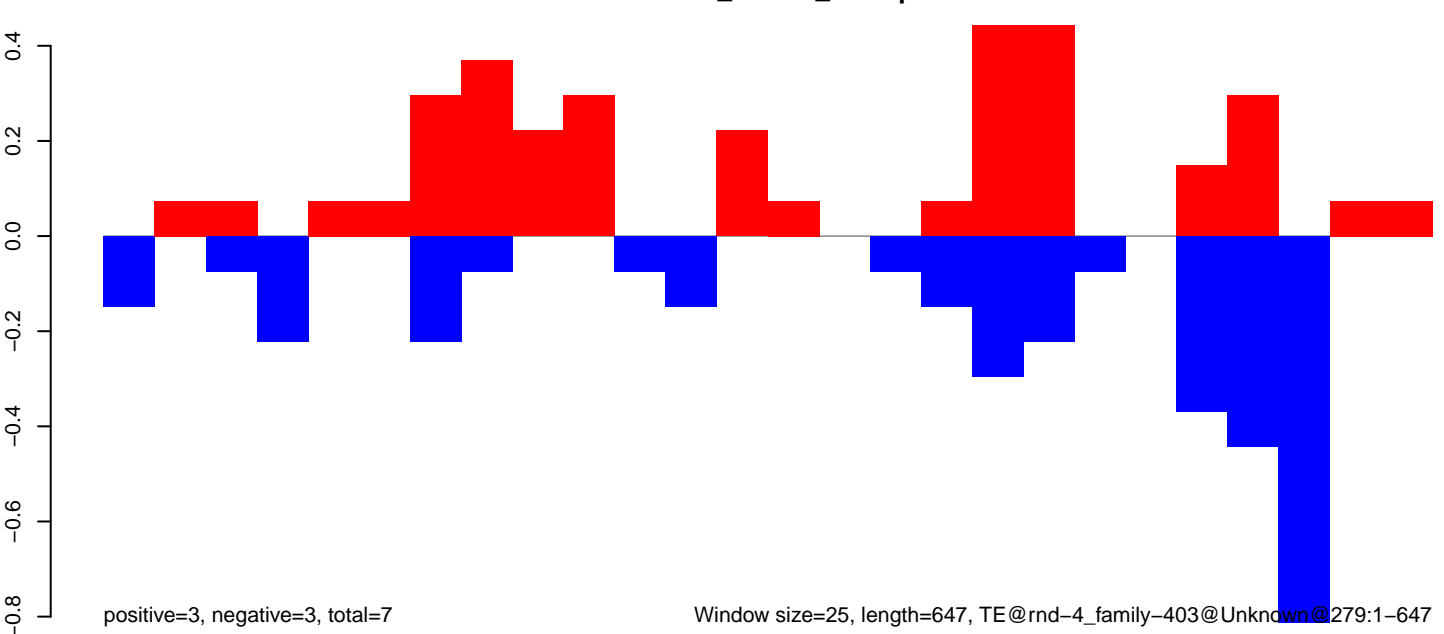
positive=1, negative=1, total=2

CuQuin_Testes_NL.24_35.rep



positive=2, negative=2, total=5

CuQuin_Testes_NL.rep

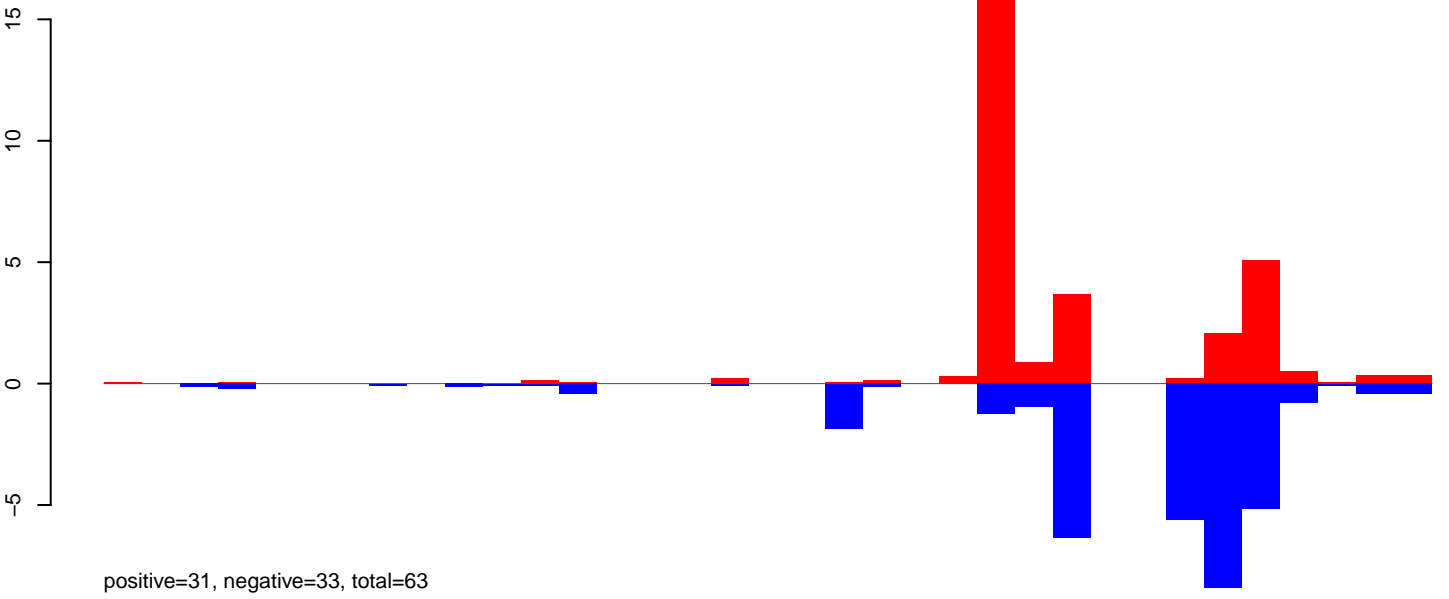


positive=3, negative=3, total=7

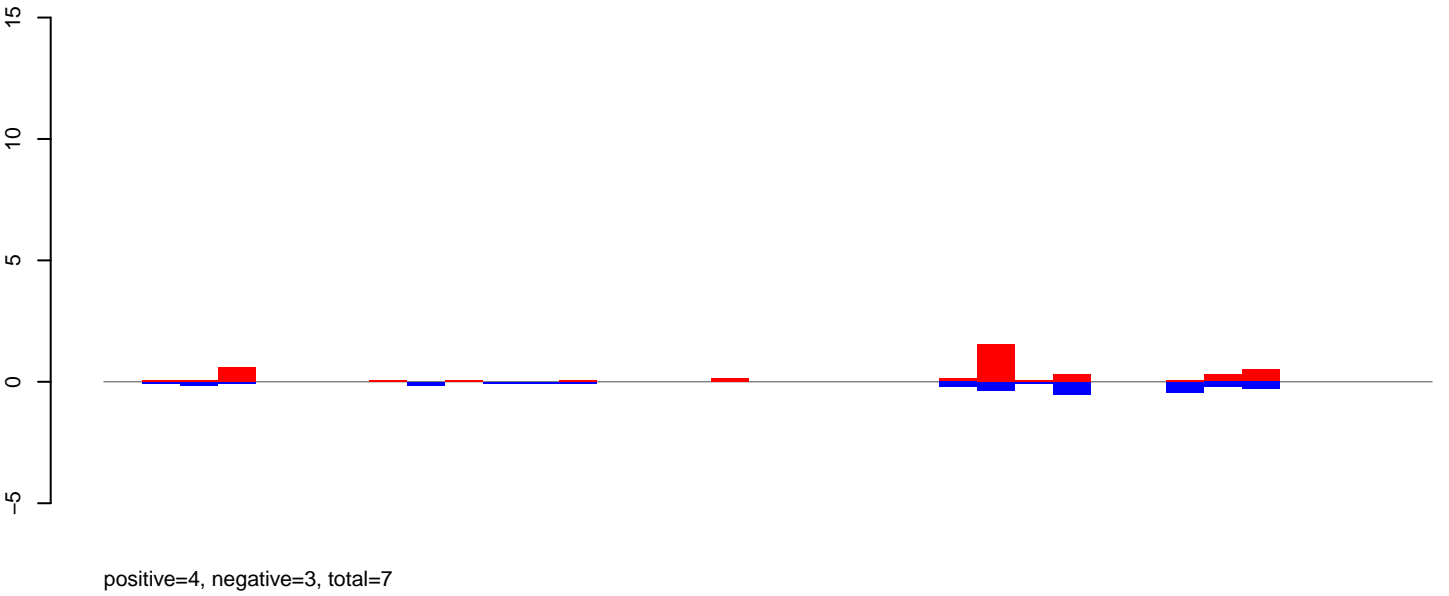
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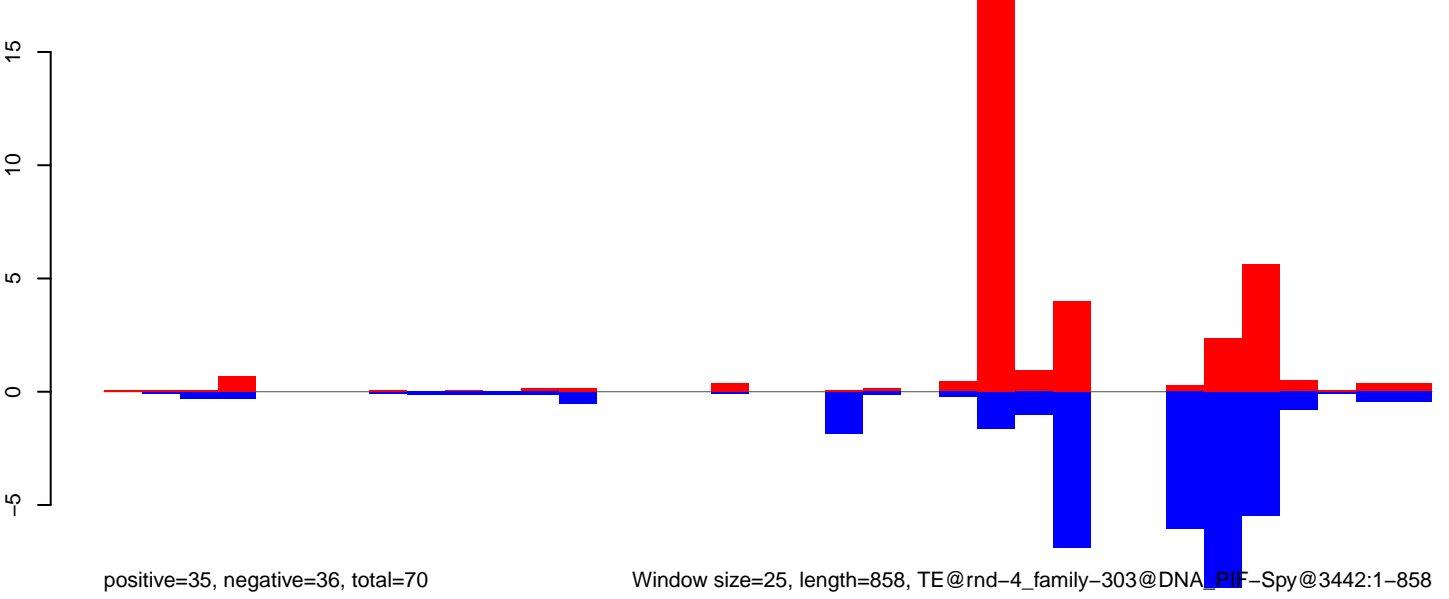
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

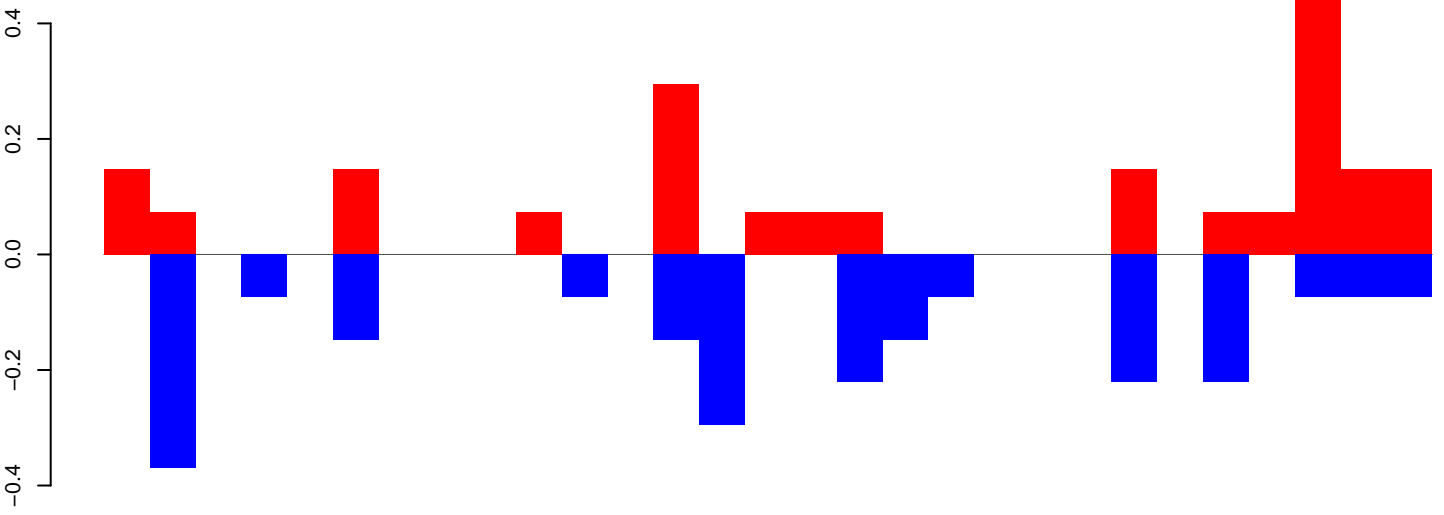


CuQuin_Testes_NL.rep



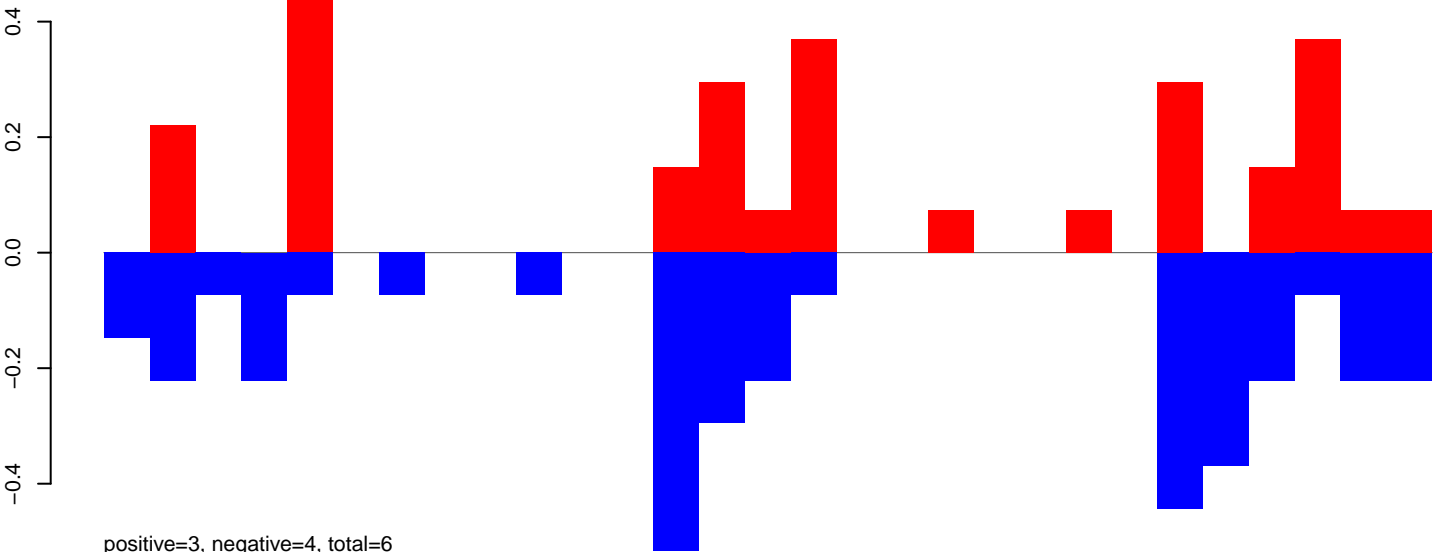
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CuQuin_Testes_NL.18_23.rep



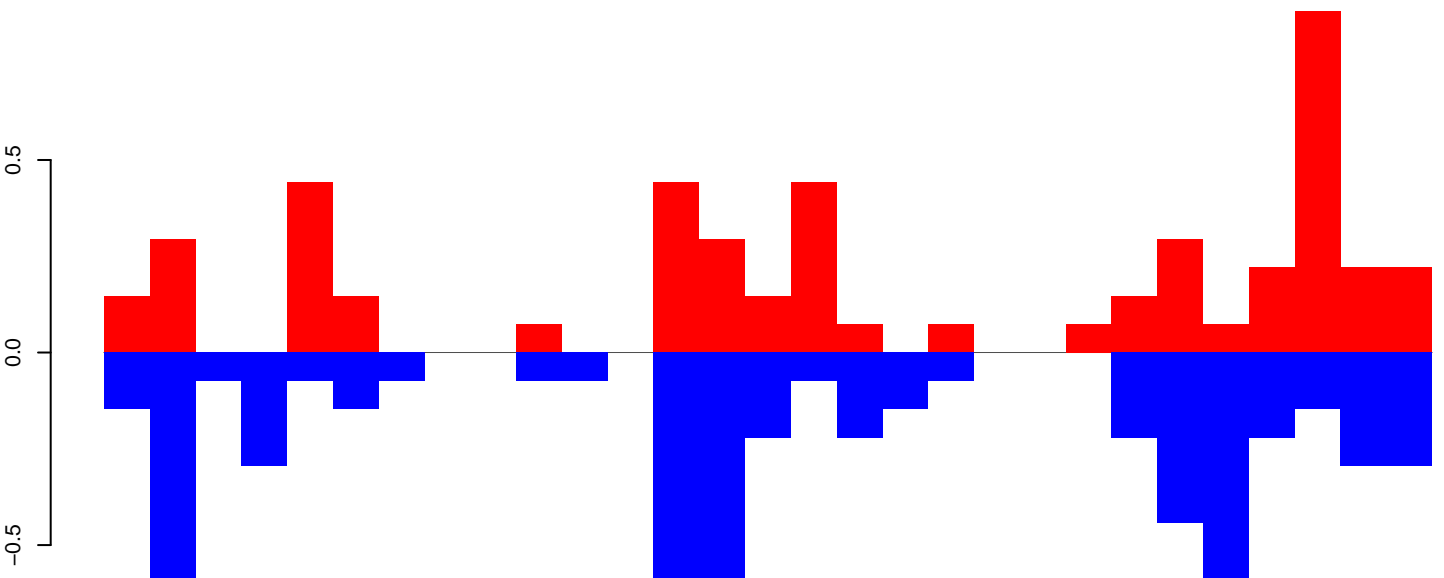
positive=2, negative=2, total=4

CuQuin_Testes_NL.24_35.rep



positive=3, negative=4, total=6

CuQuin_Testes_NL.rep

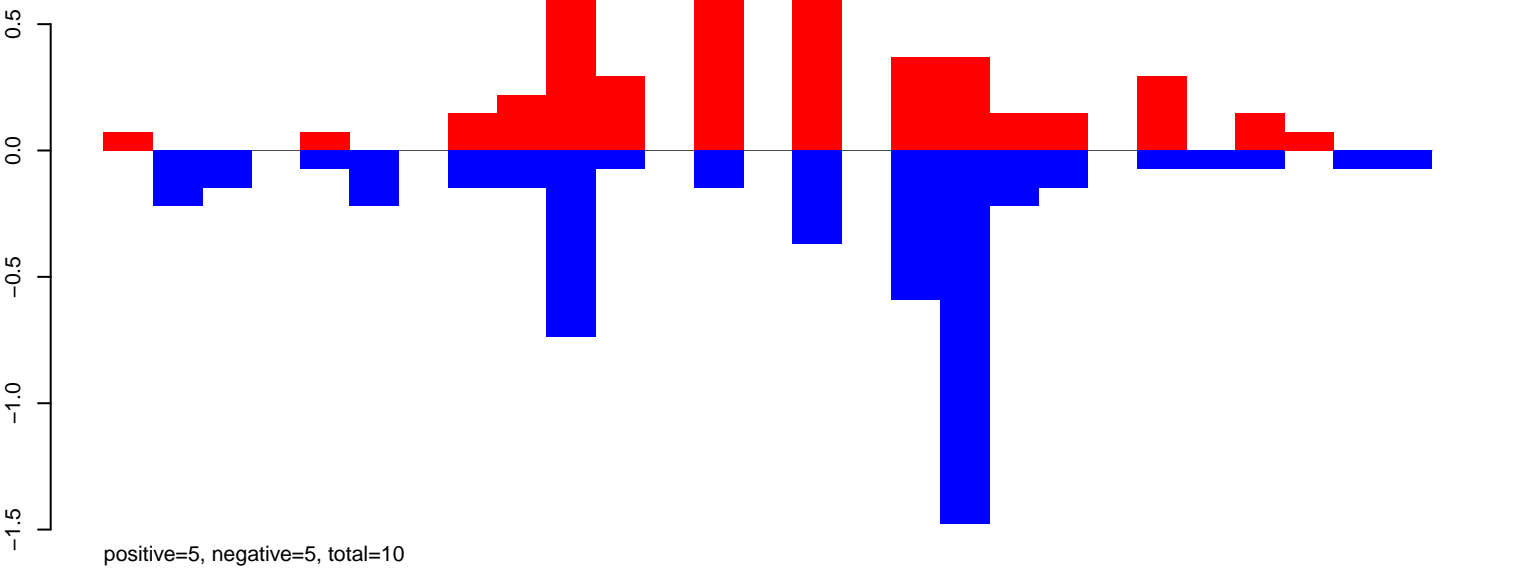


positive=5, negative=6, total=10

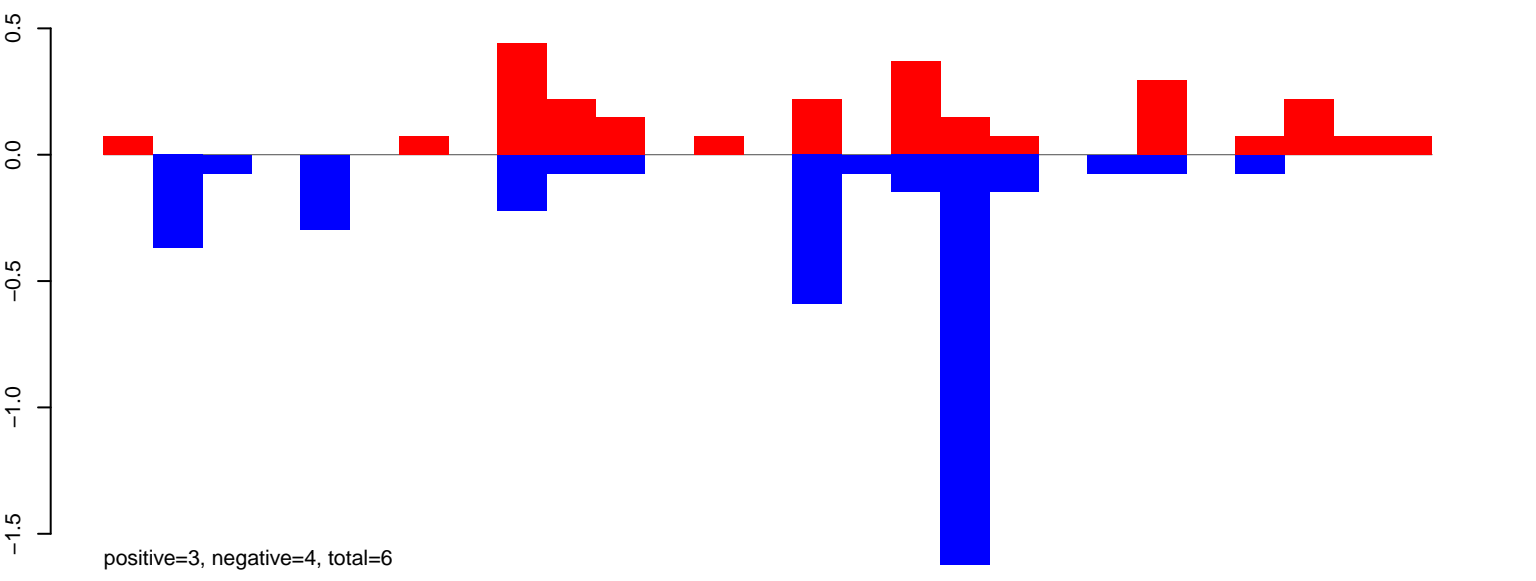
Window size=25, length=716, TE@rnd-4_family-1437@Unknown@3510:1-716

0 200 400 600

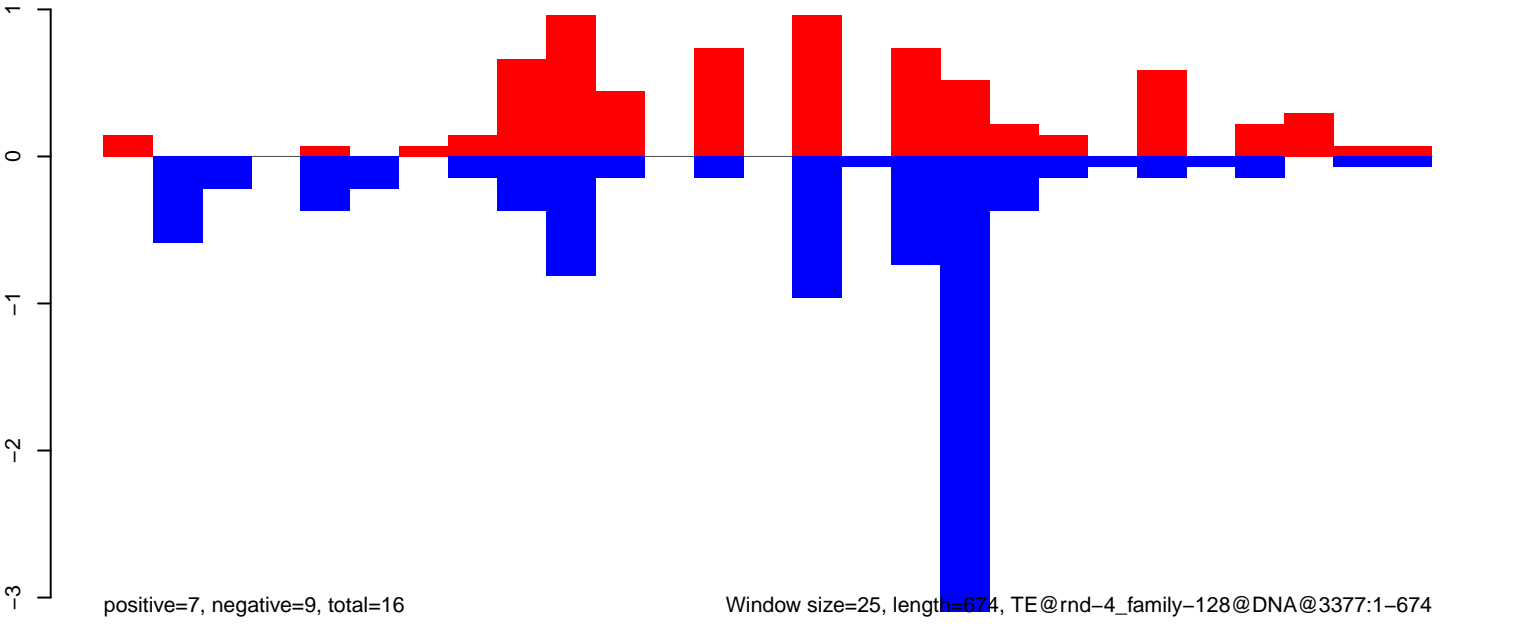
CuQuin_Testes_NL.18_23.rep



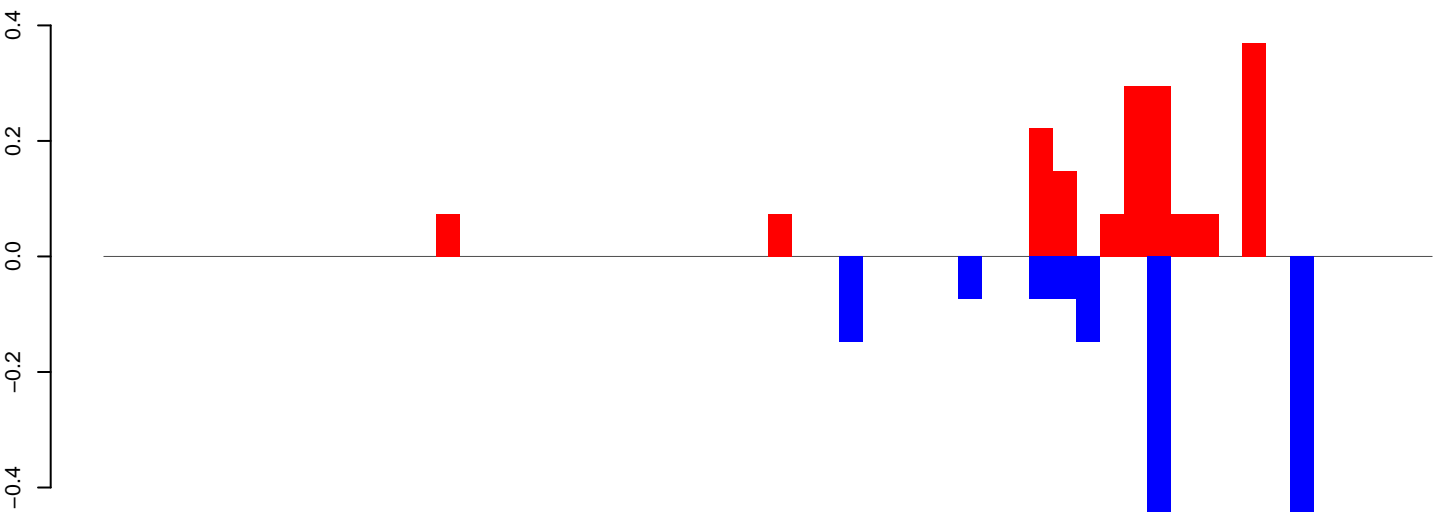
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

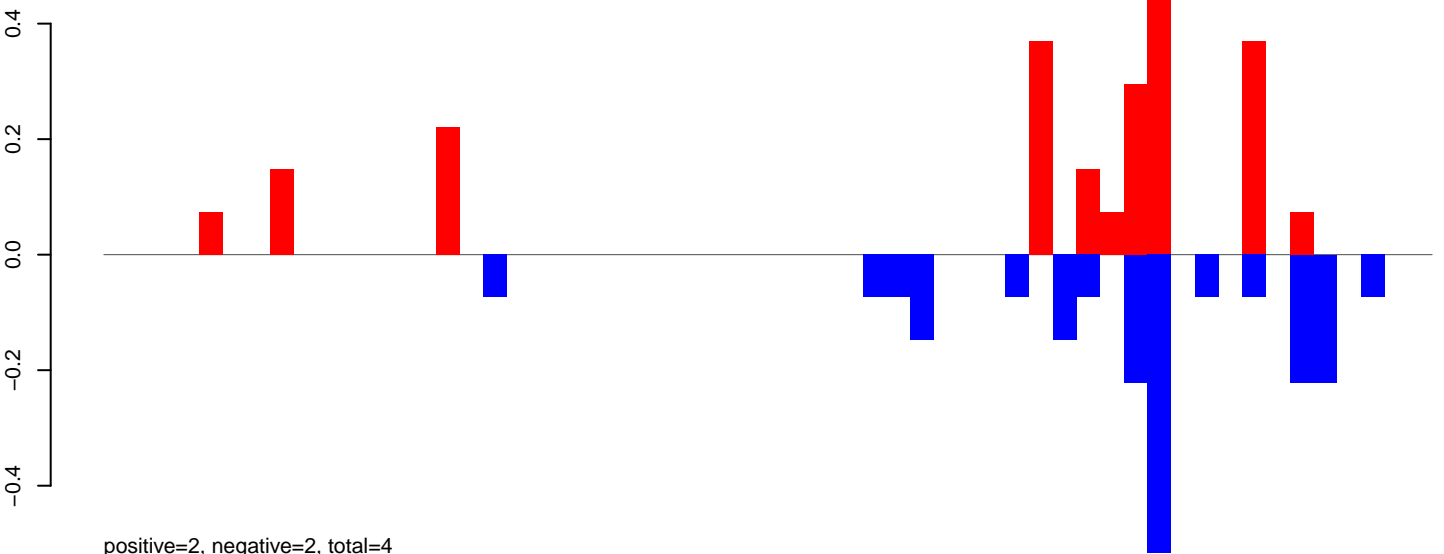


CuQuin_Testes_NL.18_23.rep



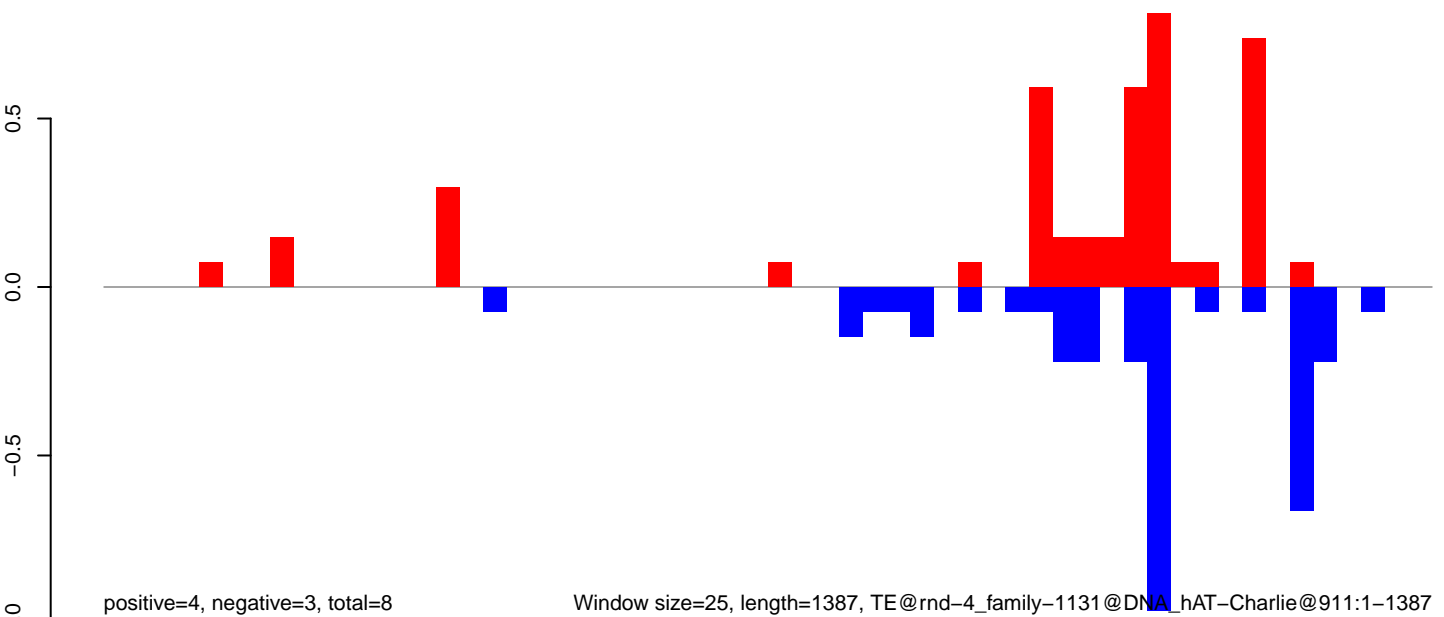
positive=2, negative=1, total=3

CuQuin_Testes_NL.24_35.rep



positive=2, negative=2, total=4

CuQuin_Testes_NL.rep

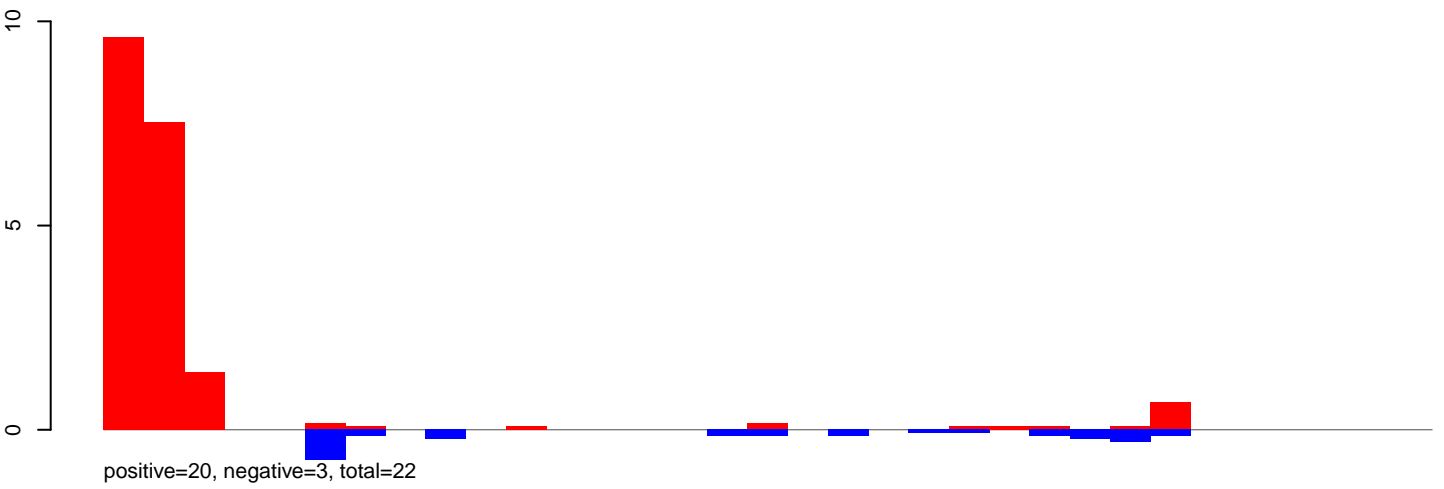


positive=4, negative=3, total=8

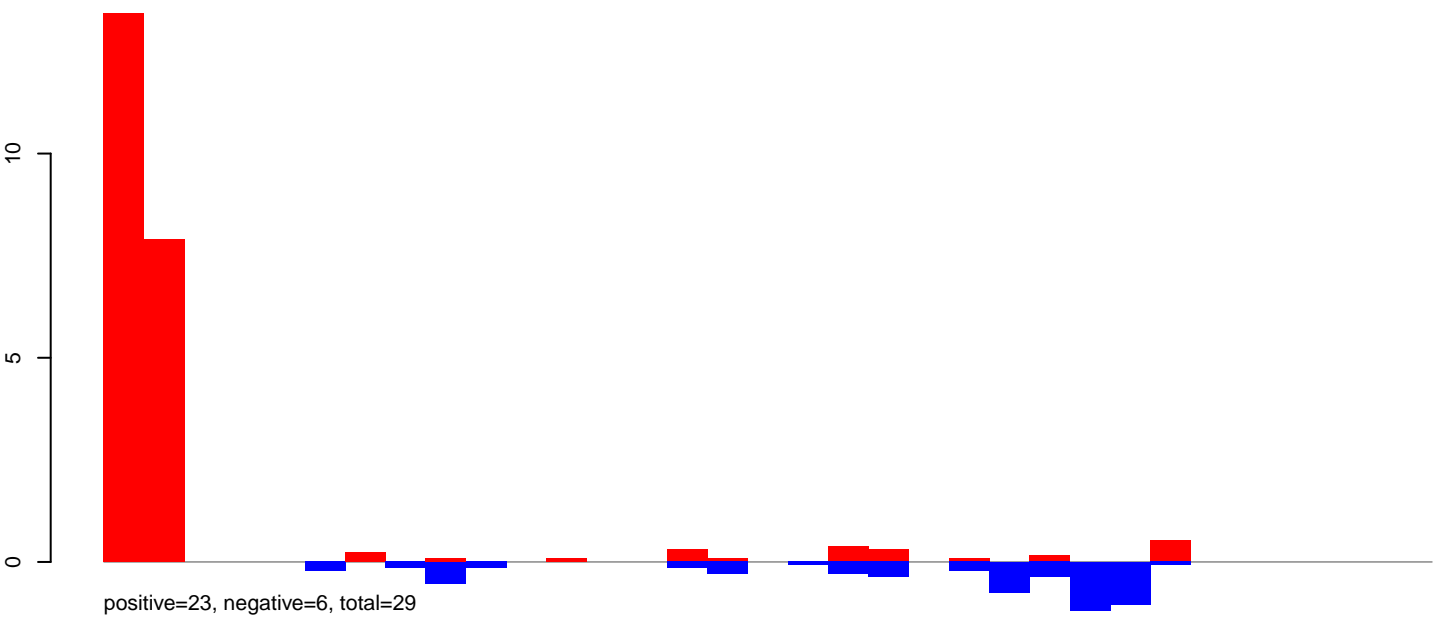
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0 200 400 600 800 1000 1200 1400

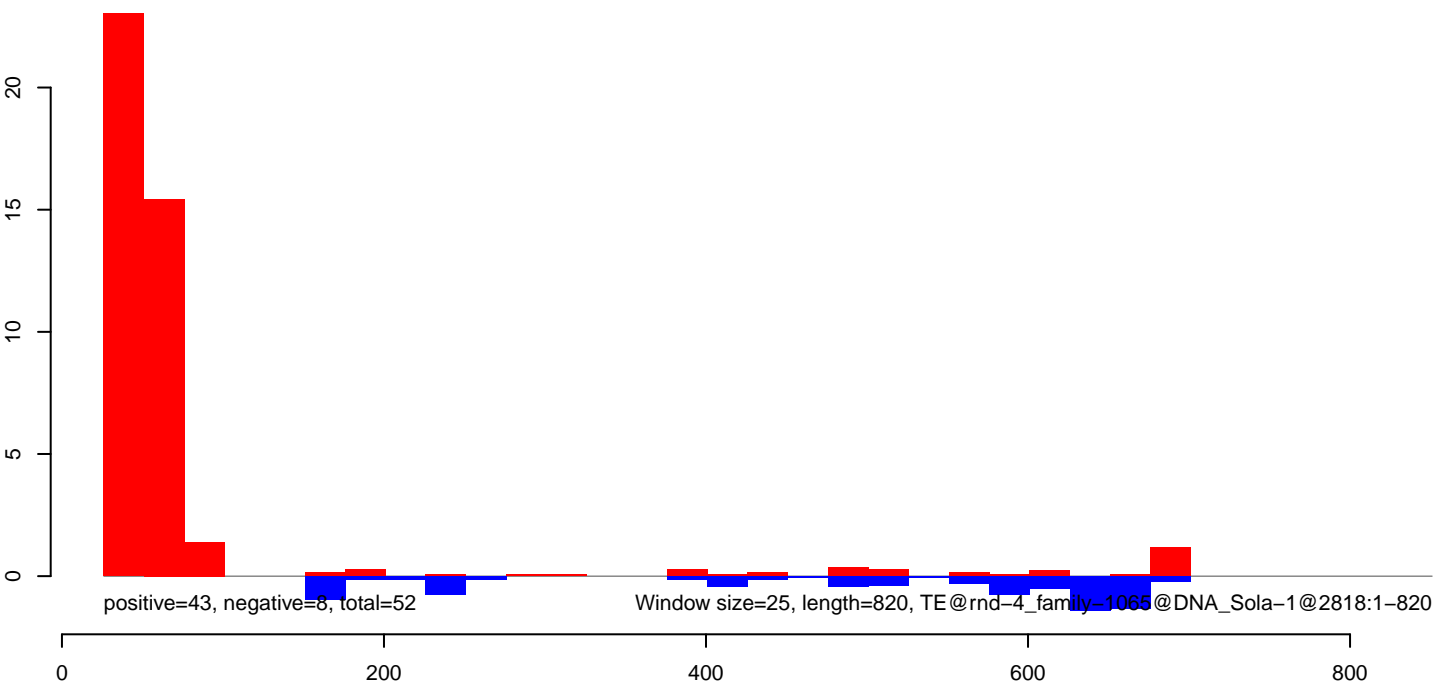
CuQuin_Testes_NL.18_23.rep



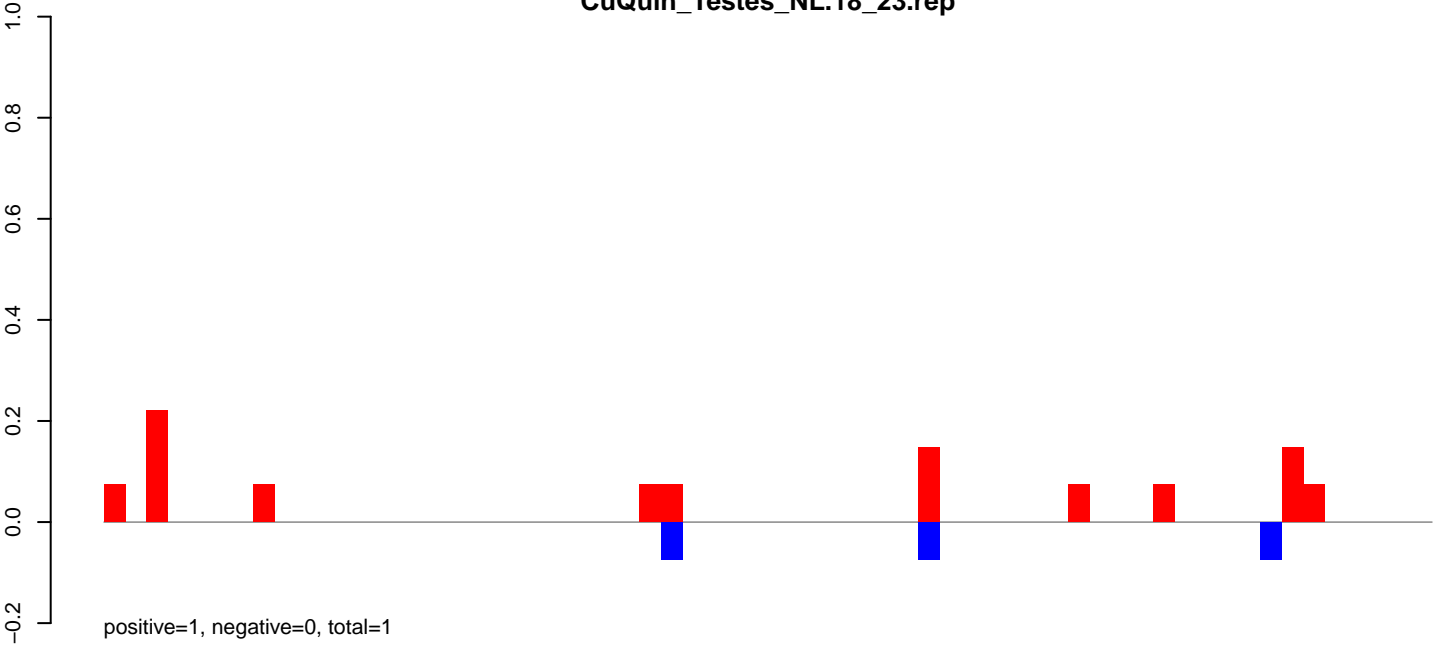
CuQuin_Testes_NL.24_35.rep



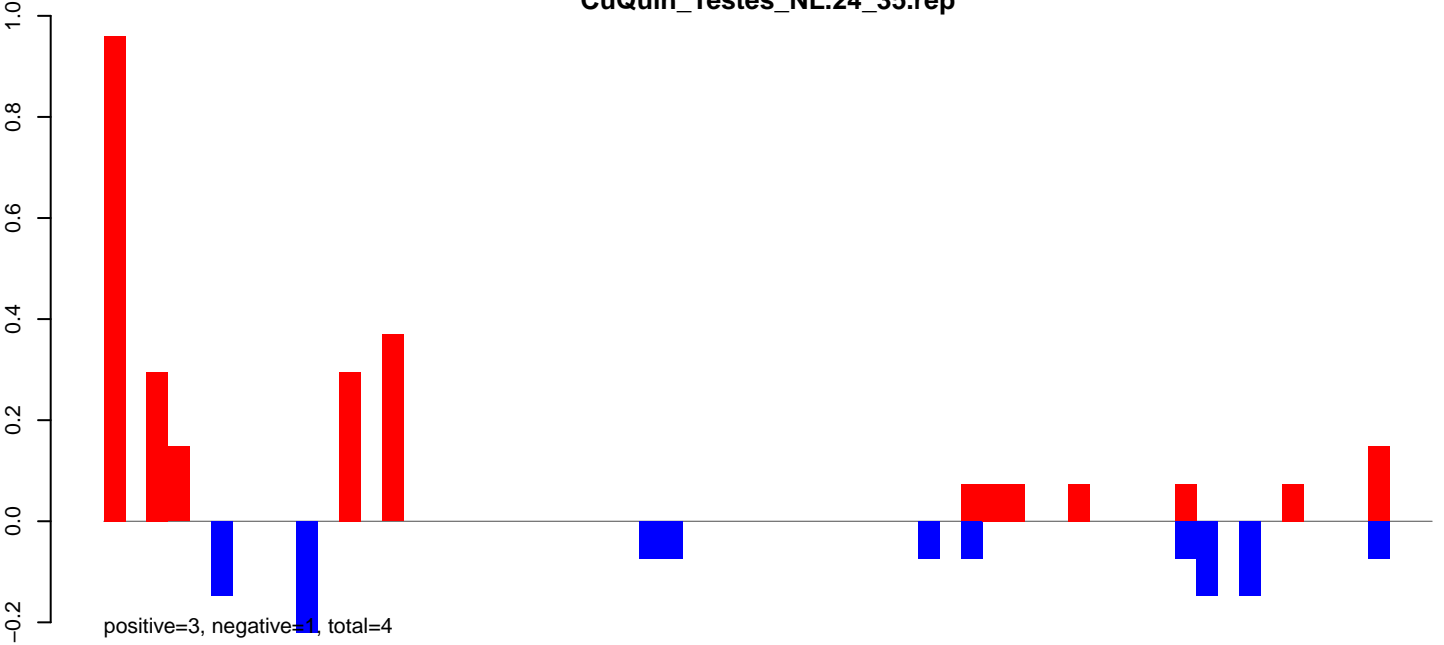
CuQuin_Testes_NL.rep



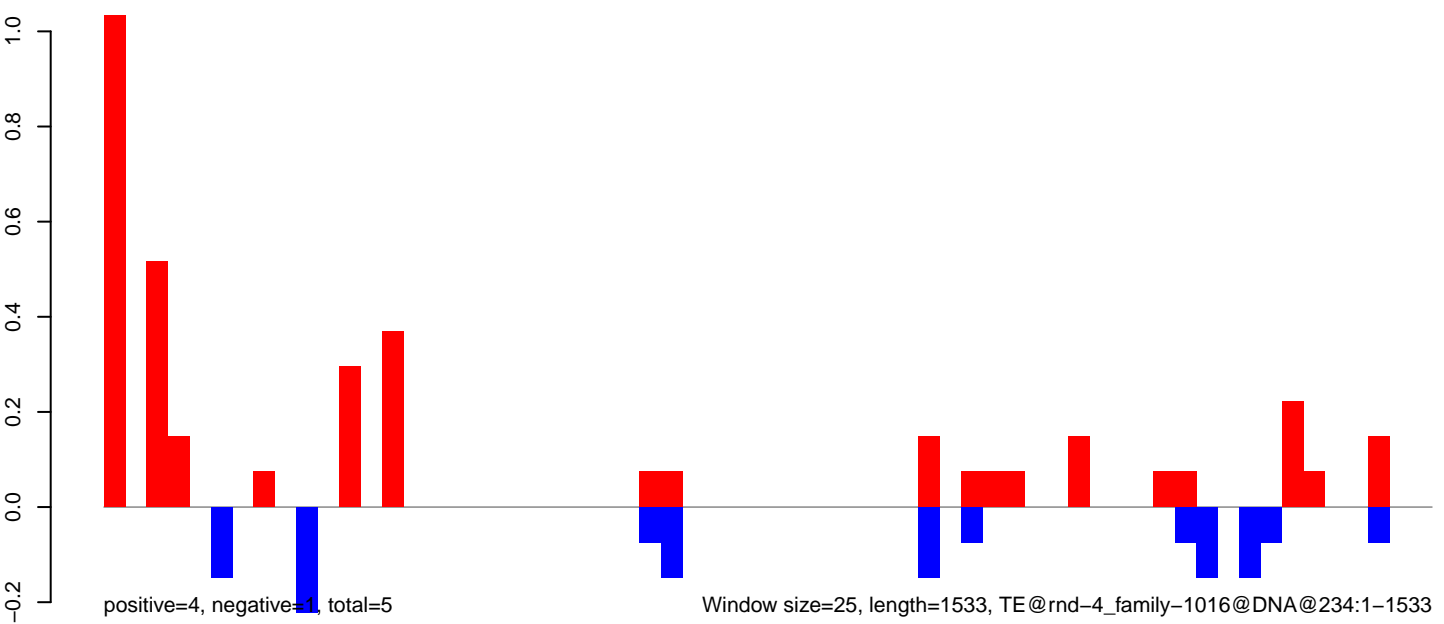
CuQuin_Testes_NL.18_23.rep



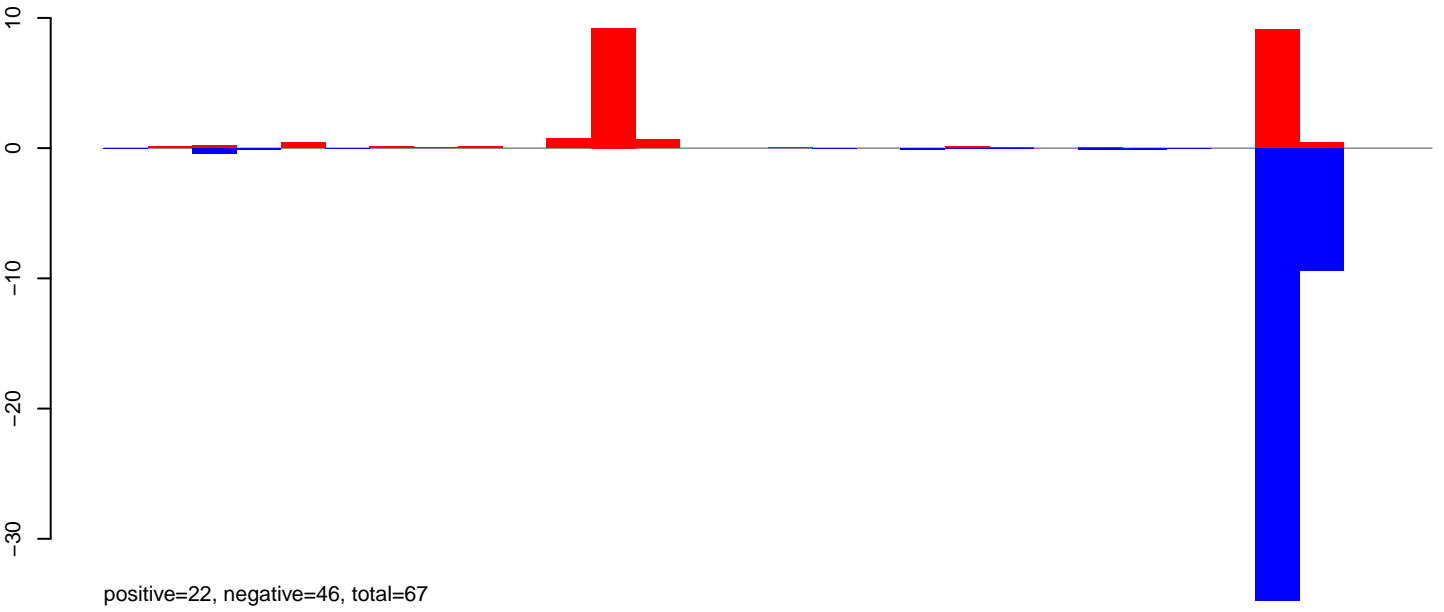
CuQuin_Testes_NL.24_35.rep



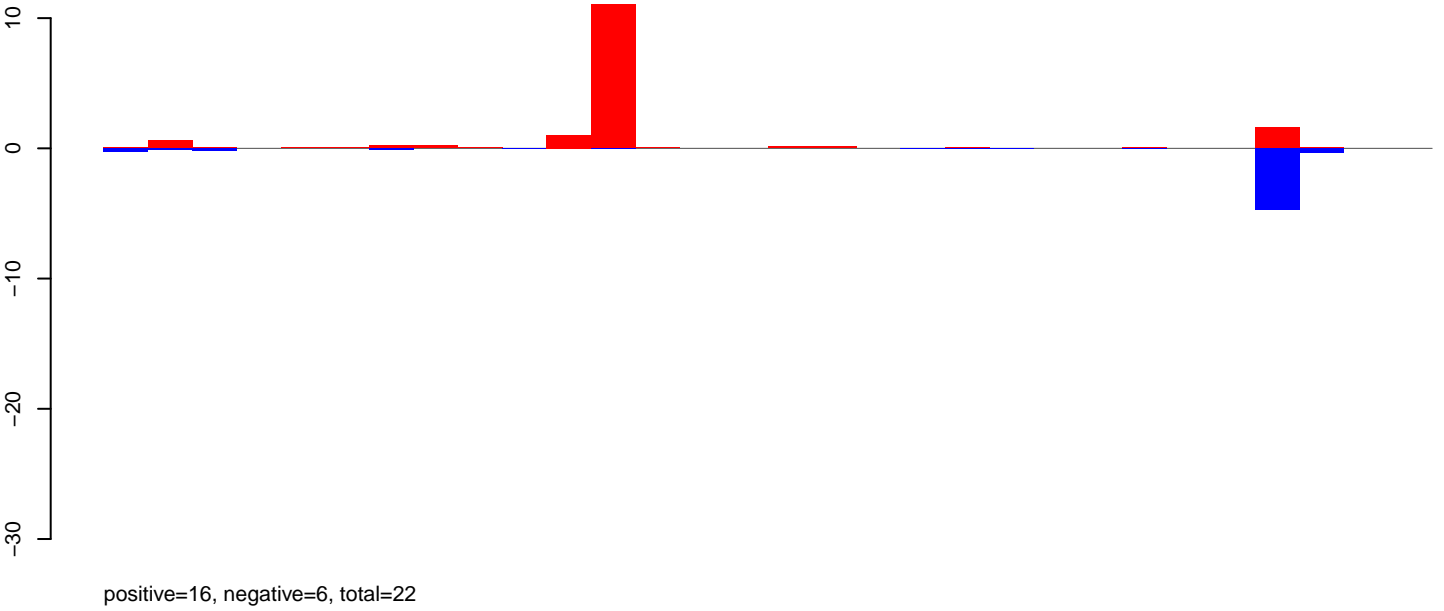
CuQuin_Testes_NL.rep



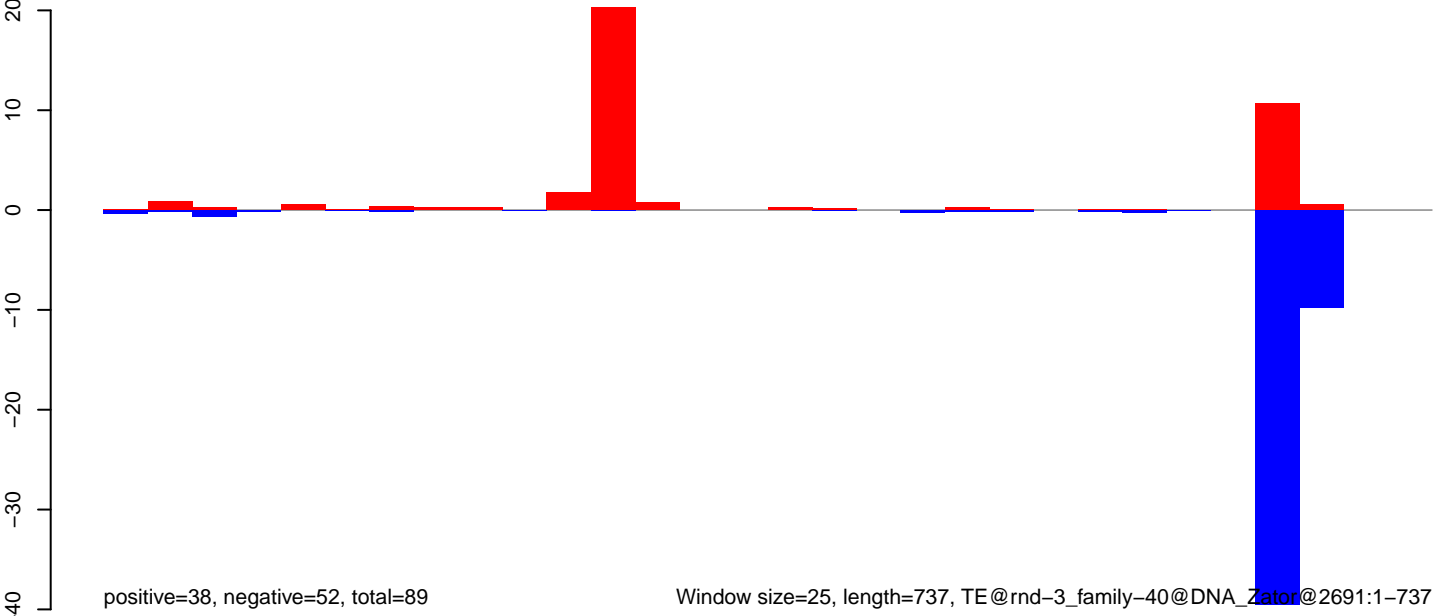
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

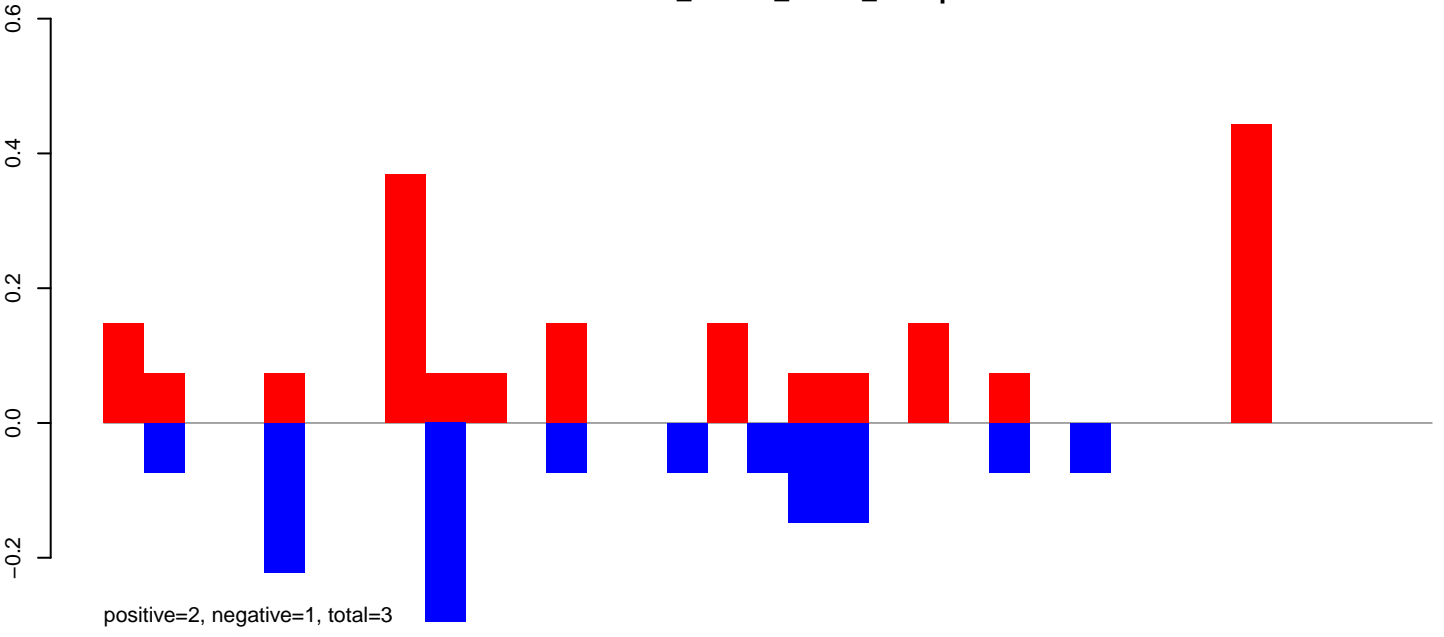


CuQuin_Testes_NL.rep

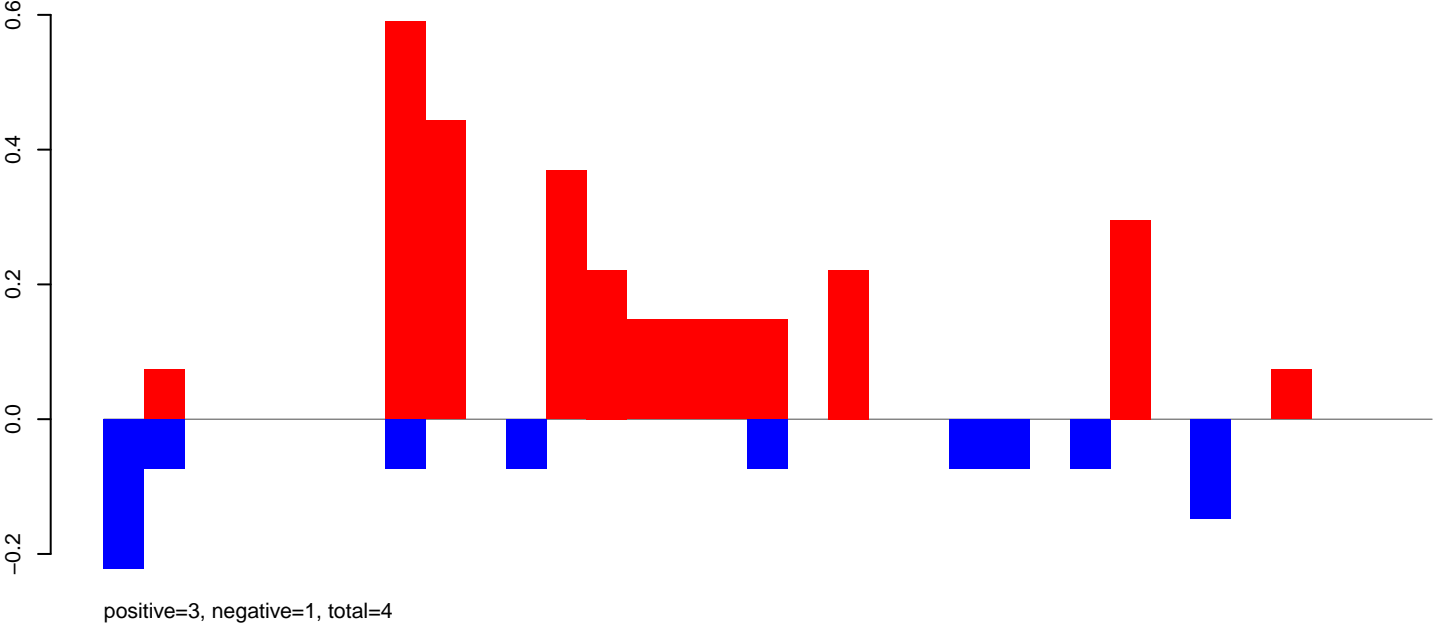


Window size=25, length=737, TE@rnd-3_family-40@DNA_Zator@2691:1-737

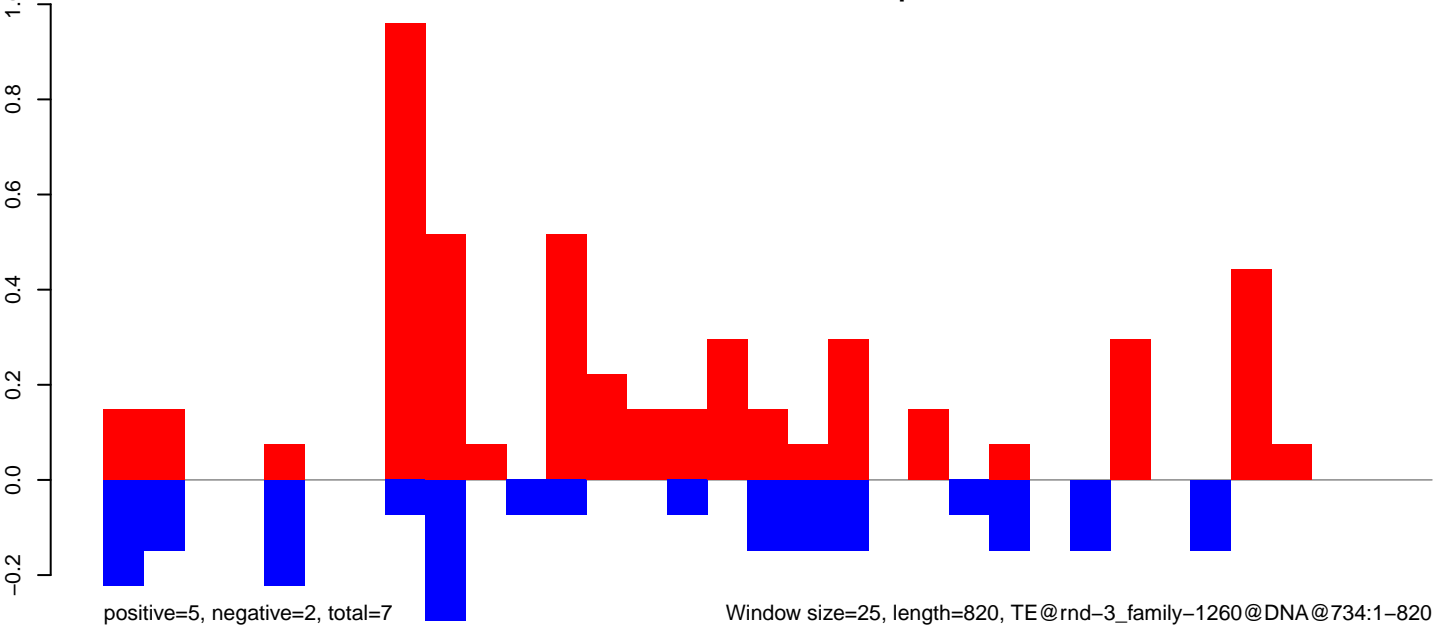
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

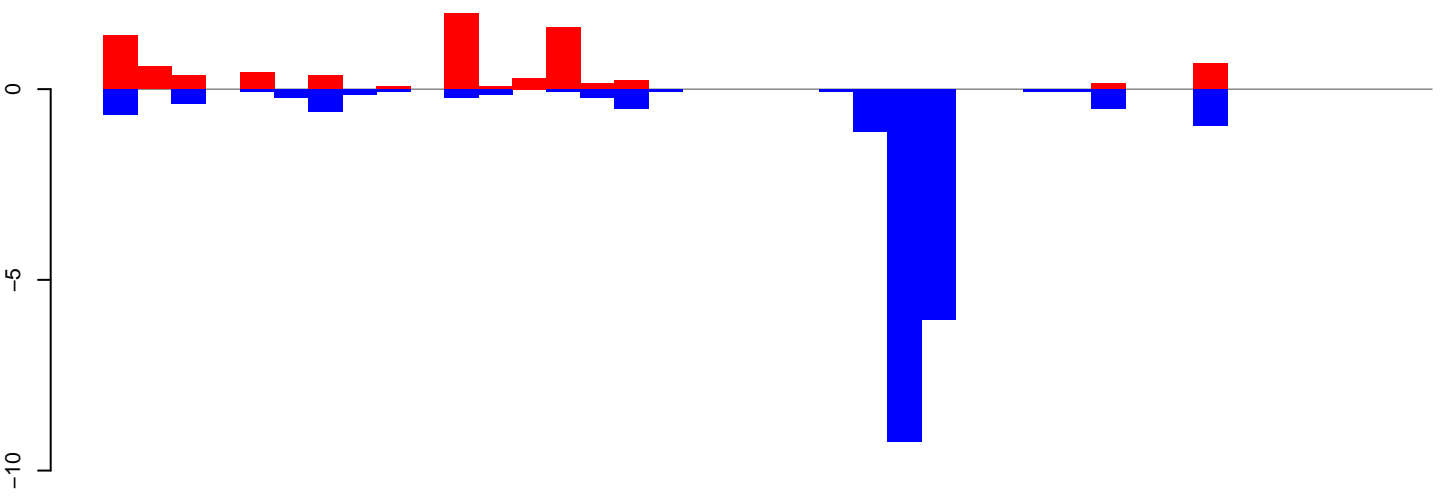


CuQuin_Testes_NL.rep



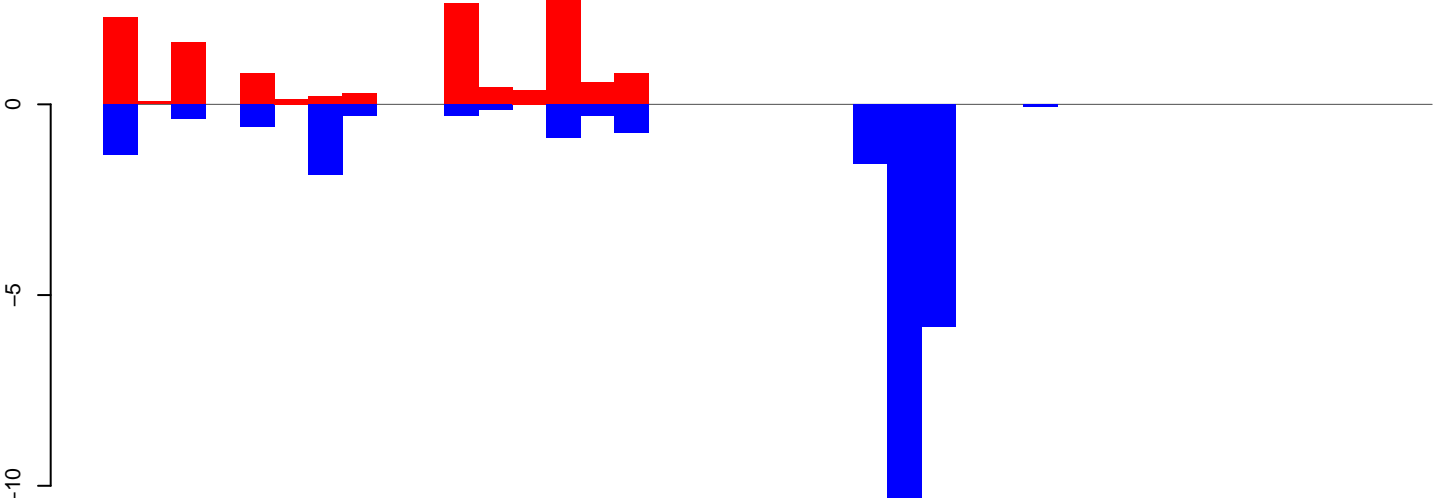
Window size=25, length=820, TE@rnd-3_family-1260@DNA@734:1-820

CuQuin_Testes_NL.18_23.rep



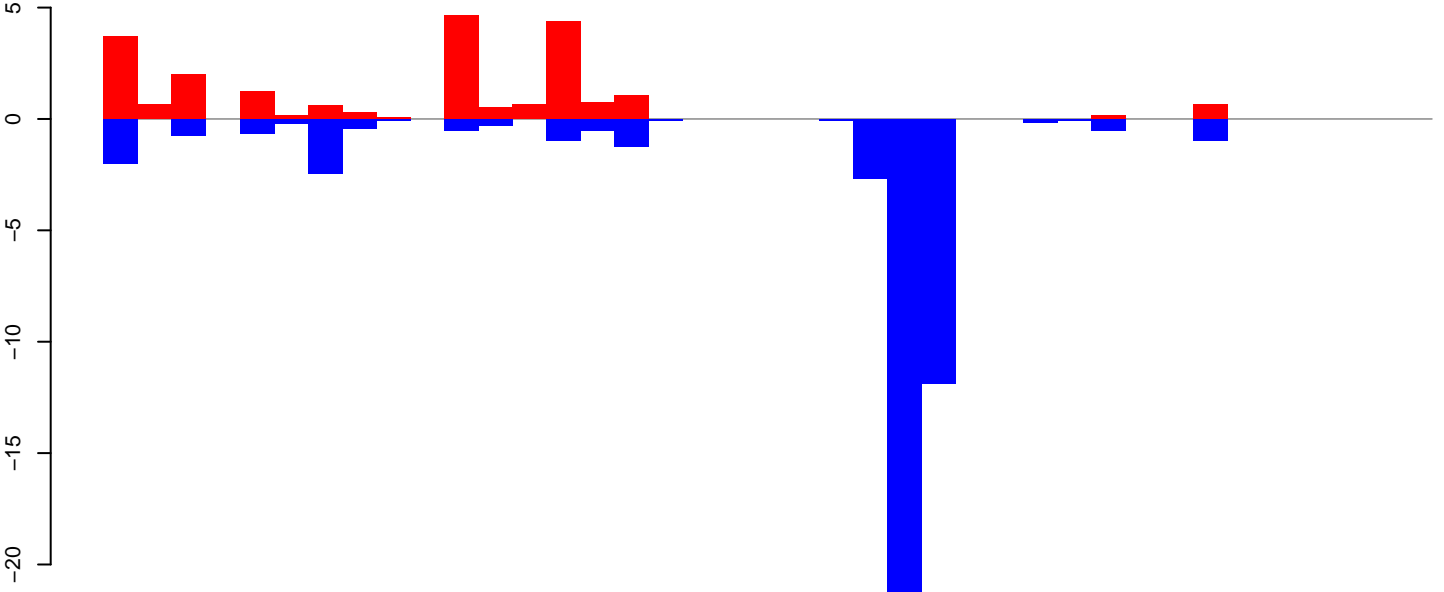
positive=8, negative=21, total=30

CuQuin_Testes_NL.24_35.rep



positive=13, negative=27, total=40

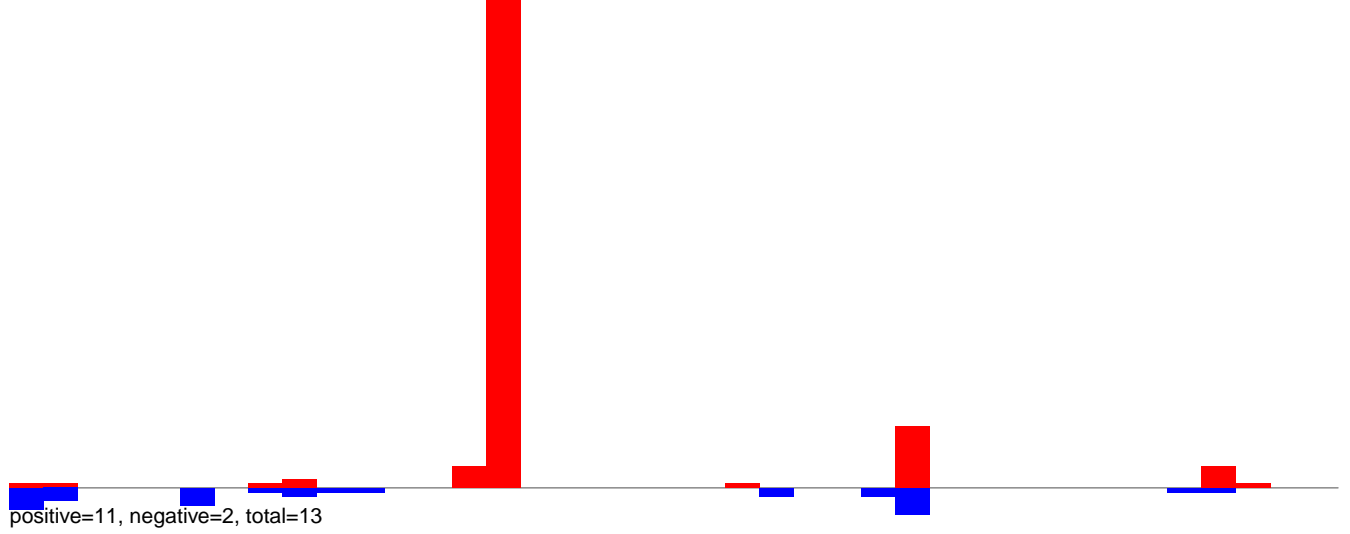
CuQuin_Testes_NL.rep



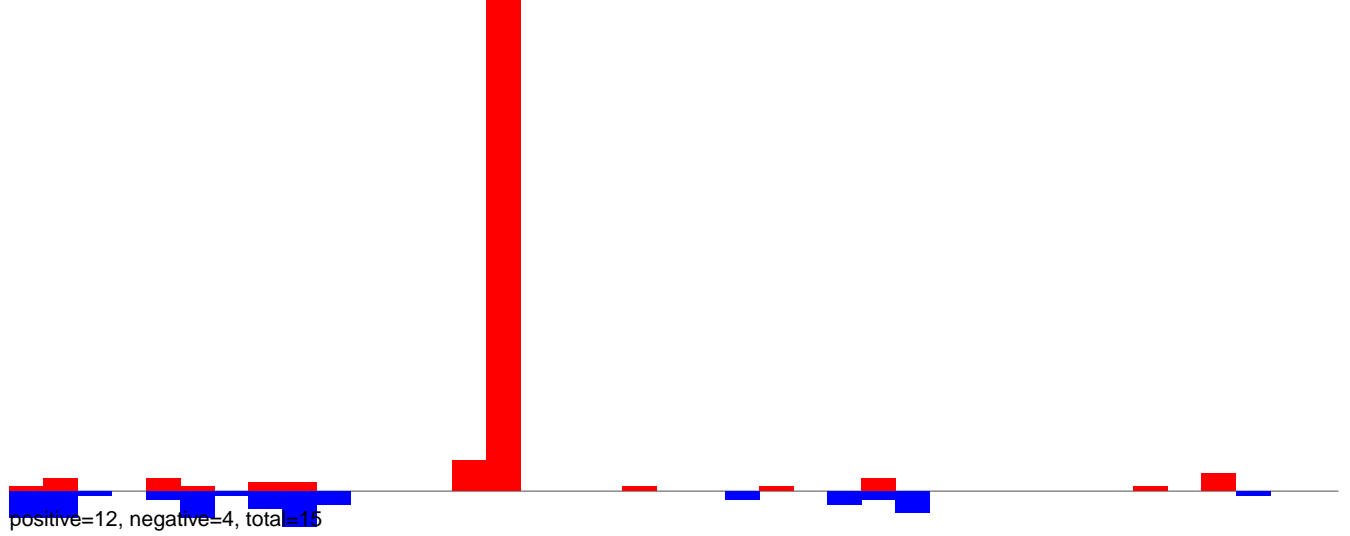
positive=21, negative=49, total=70

Window size=25, length=961, TE@rnd-2_family-669@DNA_TcMar@1281:1-961

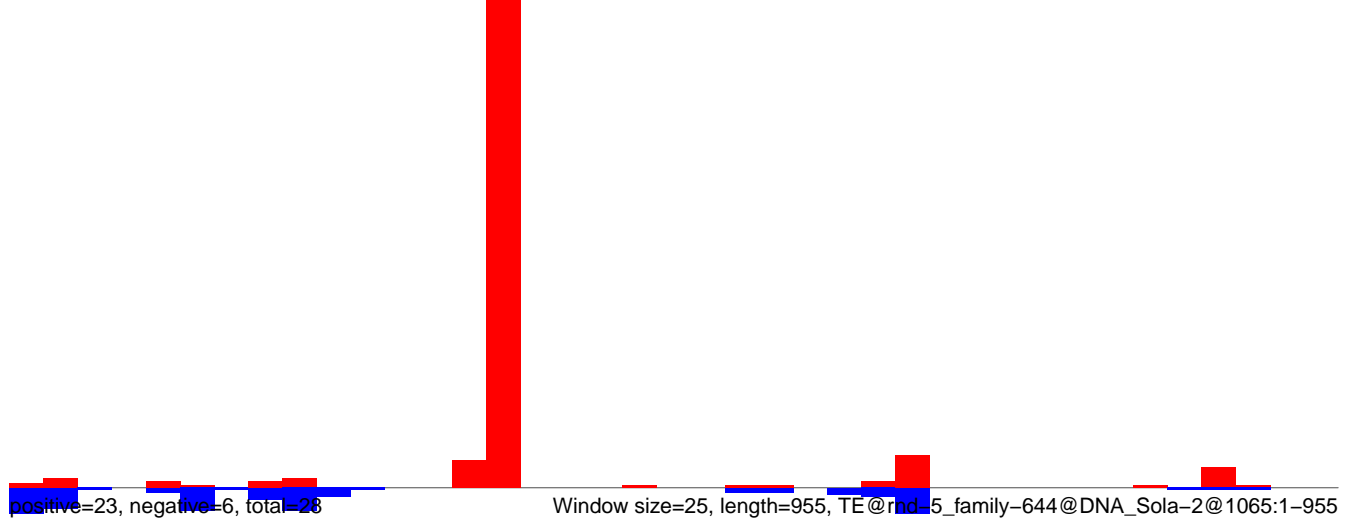
CuQuin_Testes_NL.18_23.rep



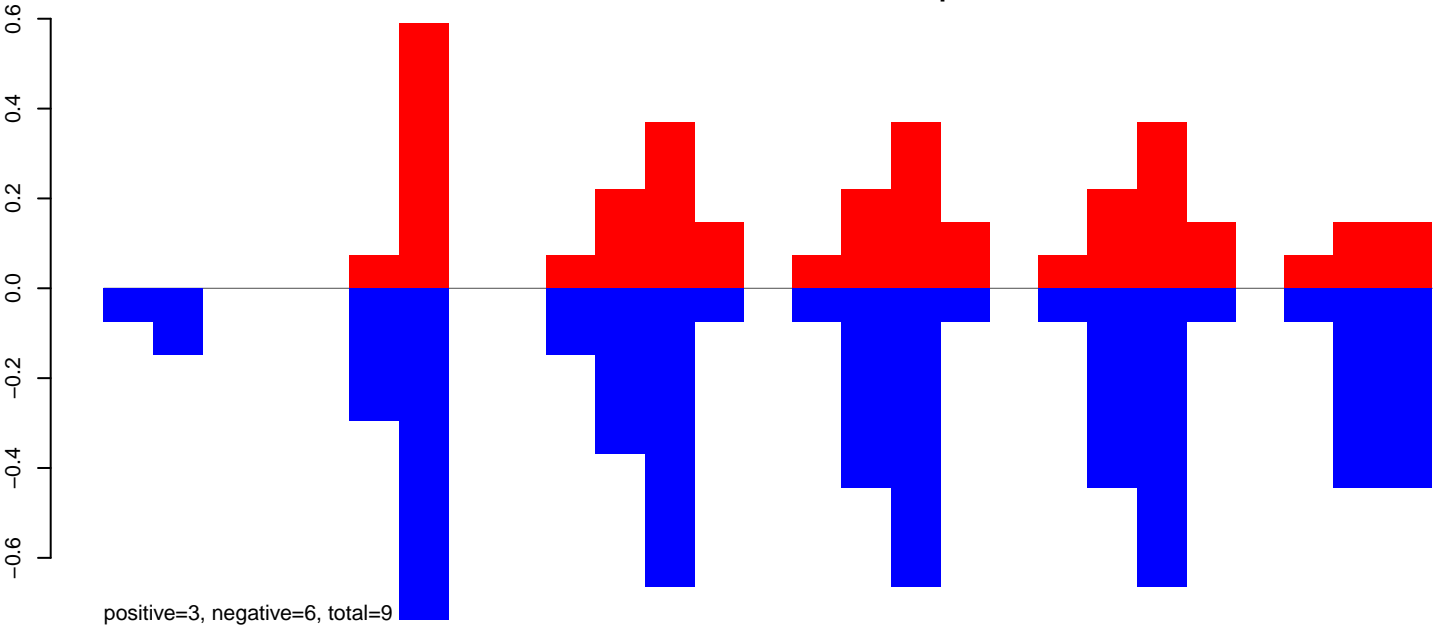
CuQuin_Testes_NL.24_35.rep



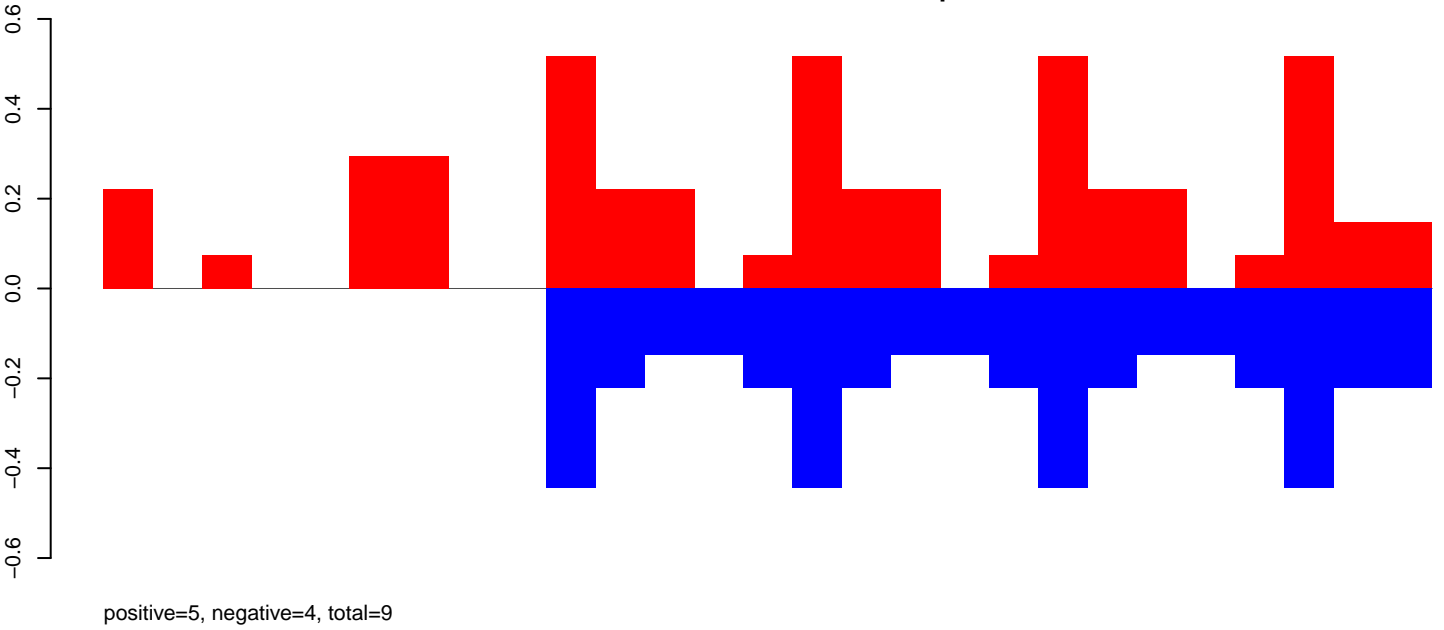
CuQuin_Testes_NL.rep



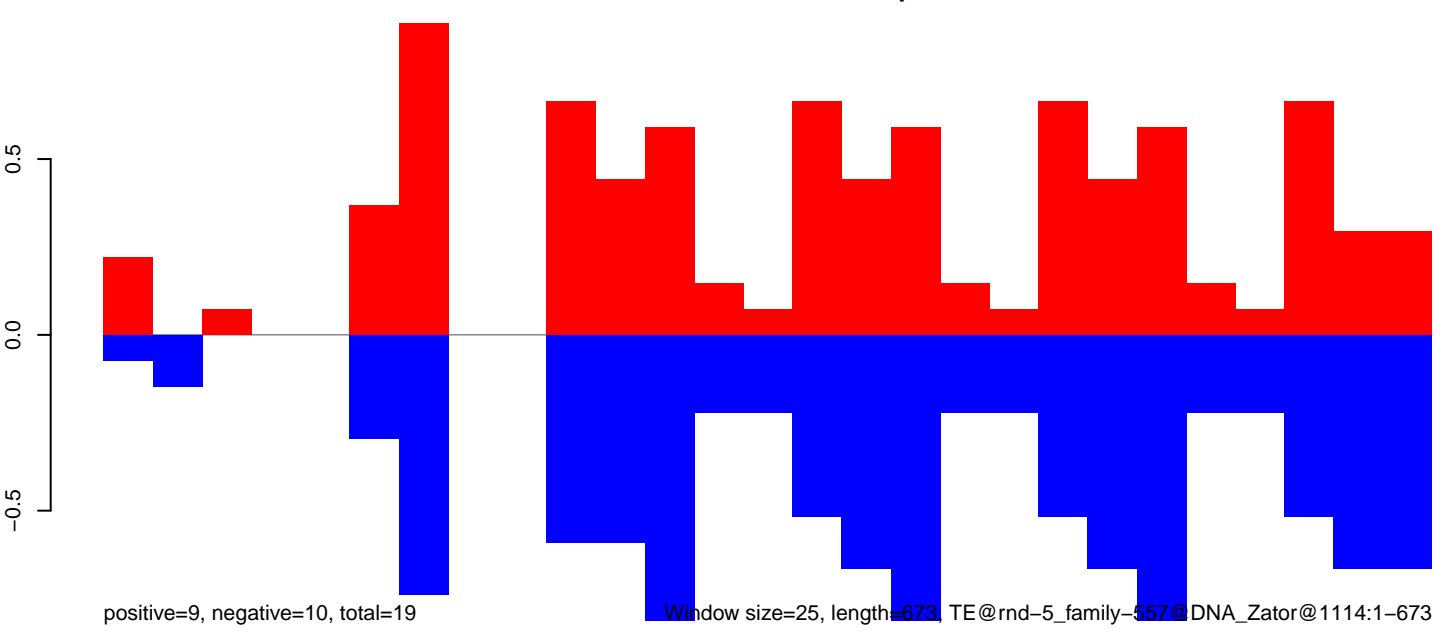
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

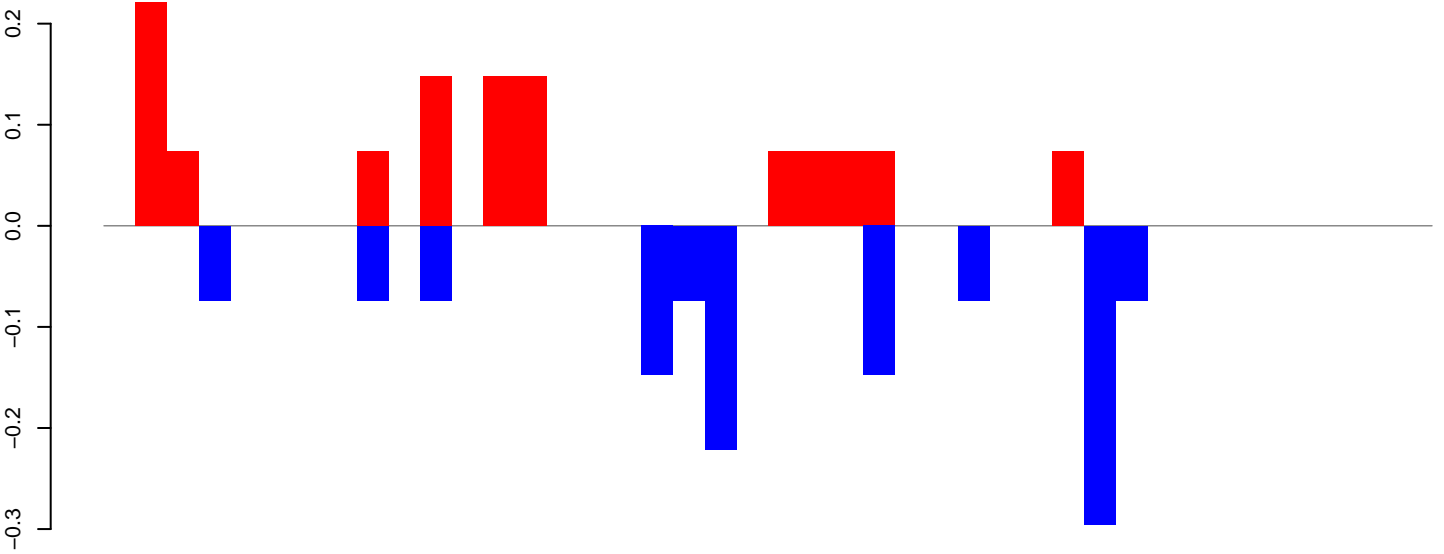


CuQuin_Testes_NL.rep



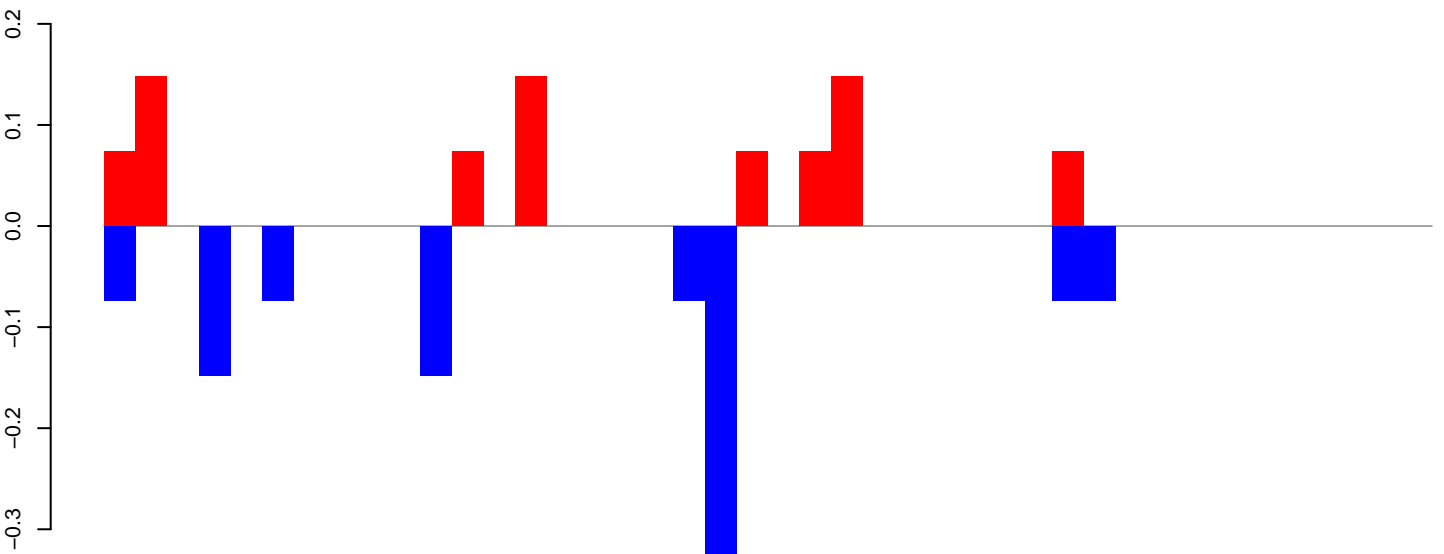
Window size=25, length=673, TE@rnd-5_family-557@DNA_Zator@1114:1-673

CuQuin_Testes_NL.18_23.rep



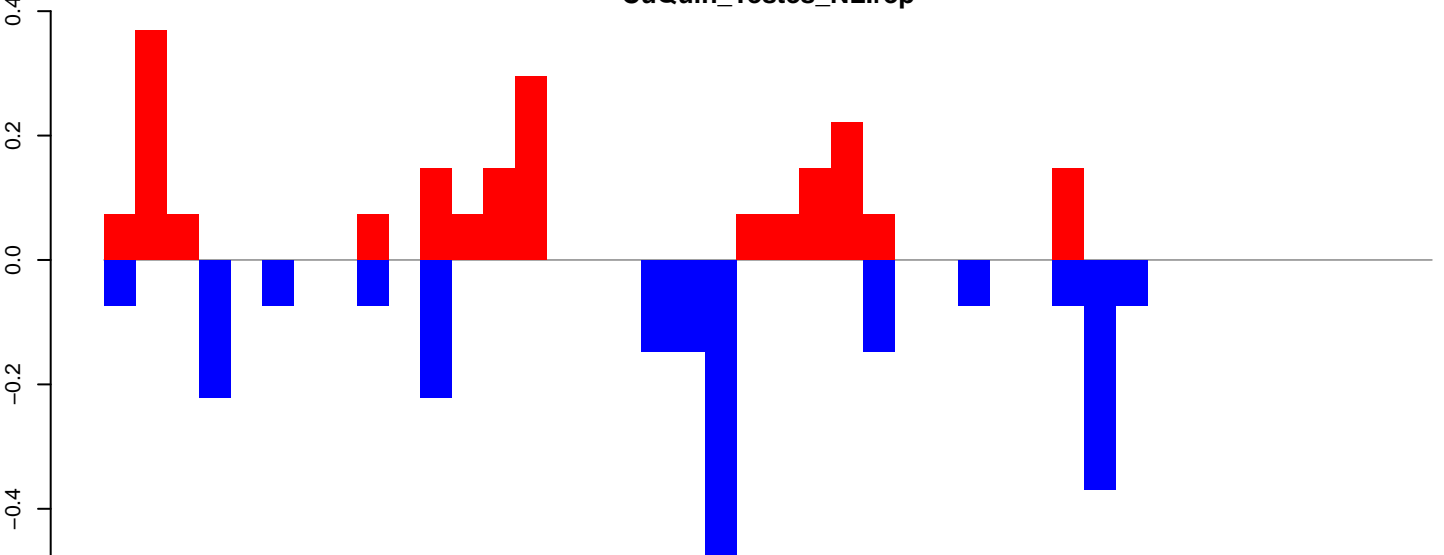
positive=1, negative=1, total=2

CuQuin_Testes_NL.24_35.rep



positive=1, negative=1, total=2

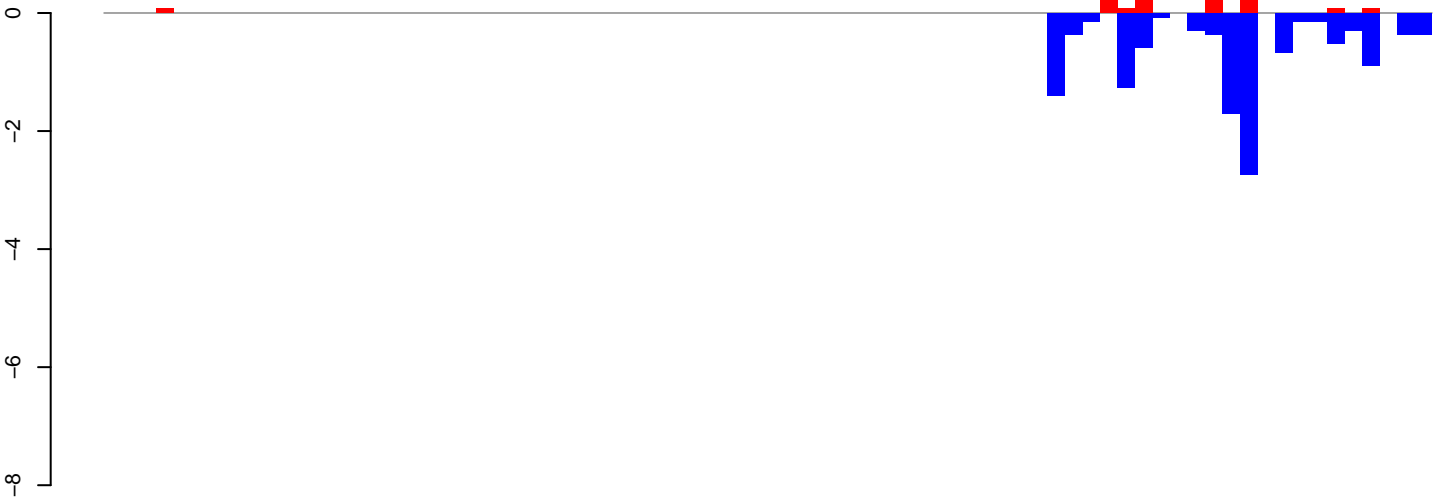
CuQuin_Testes_NL.rep



positive=2, negative=2, total=4

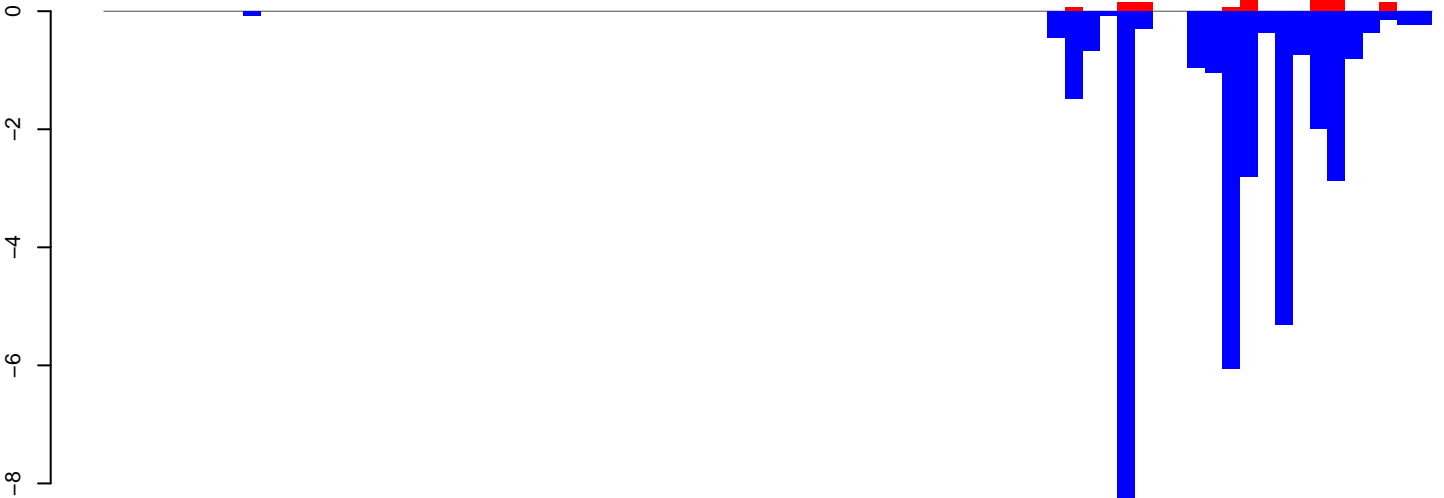
Window size=25, length=1035, TE@rnd-5_family-3929@DNA_Zator@3180:1-1035

CuQuin_Testes_NL.18_23.rep



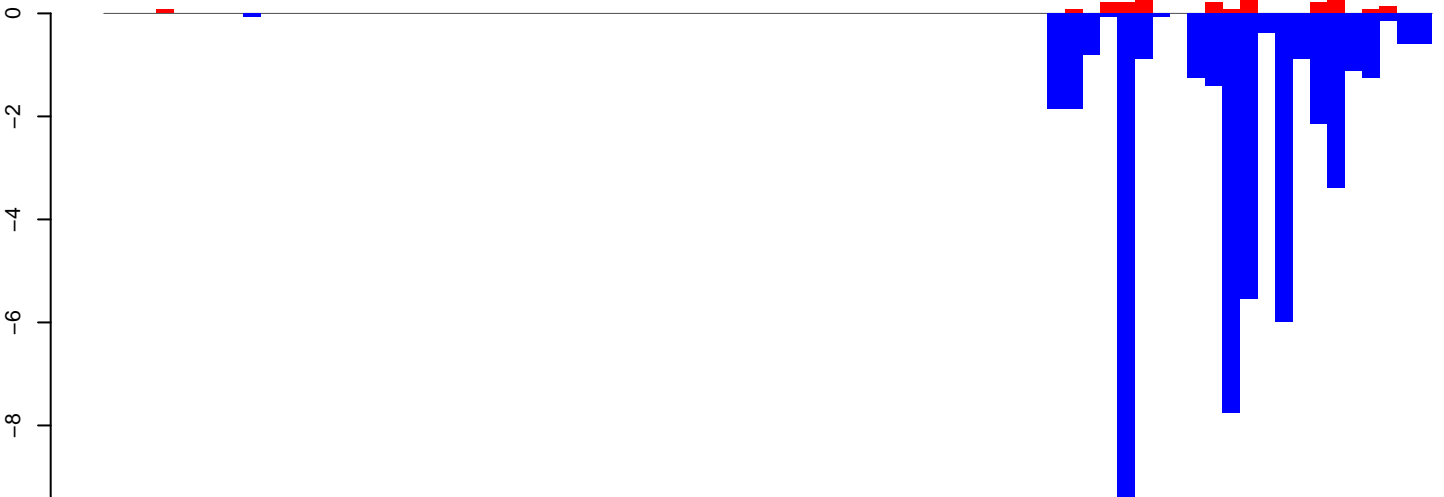
positive=1, negative=12, total=14

CuQuin_Testes_NL.24_35.rep



positive=2, negative=36, total=38

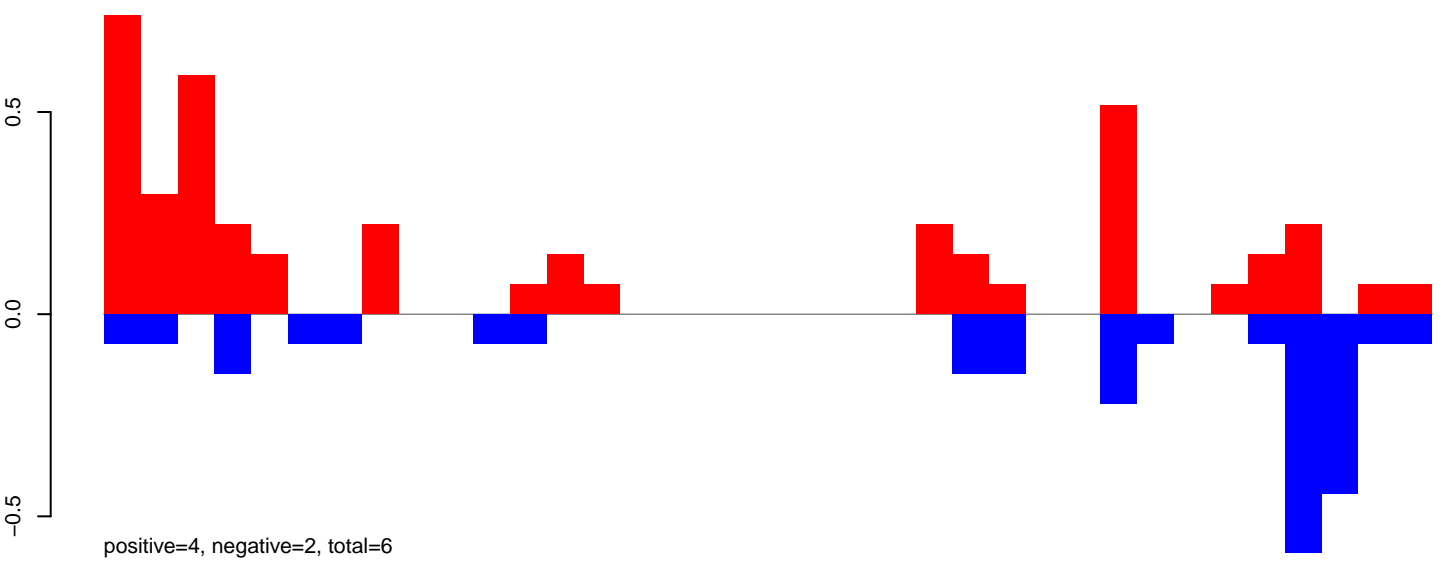
CuQuin_Testes_NL.rep



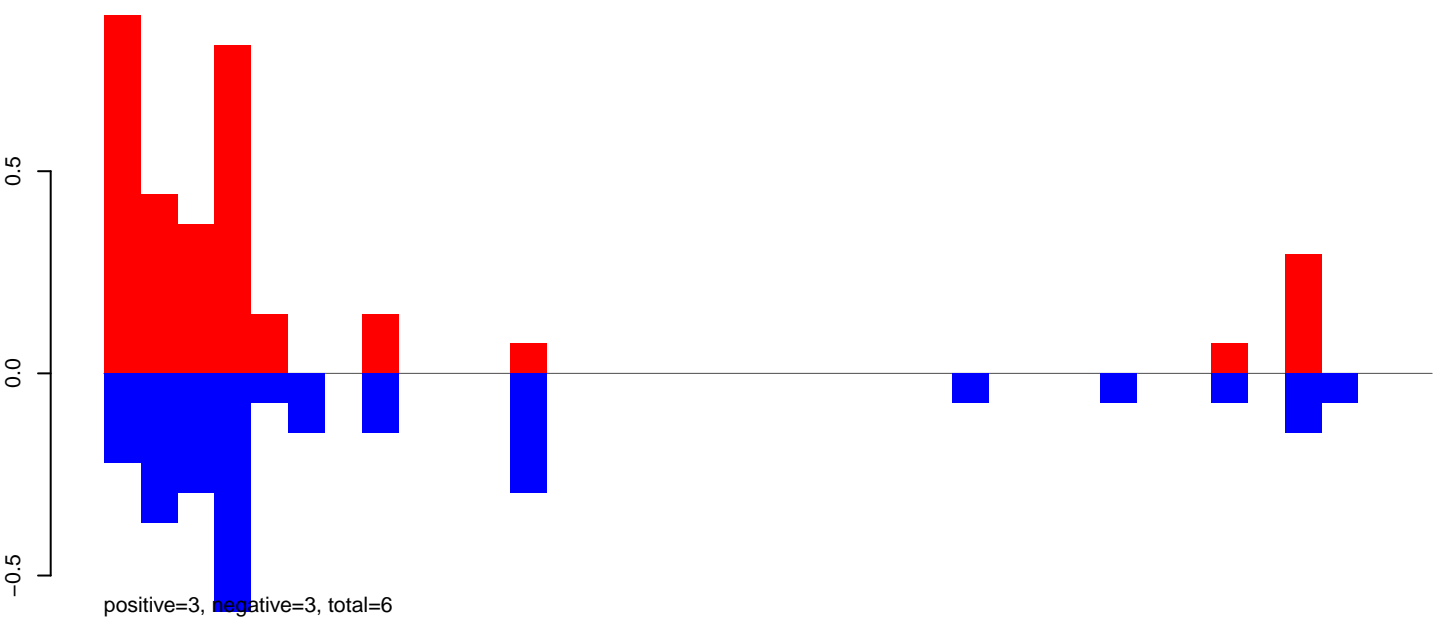
positive=4, negative=48, total=52

Window size=25, length=1900, TE@rnd-5_family-1996@LTR_Gypsy@2542:1-1900

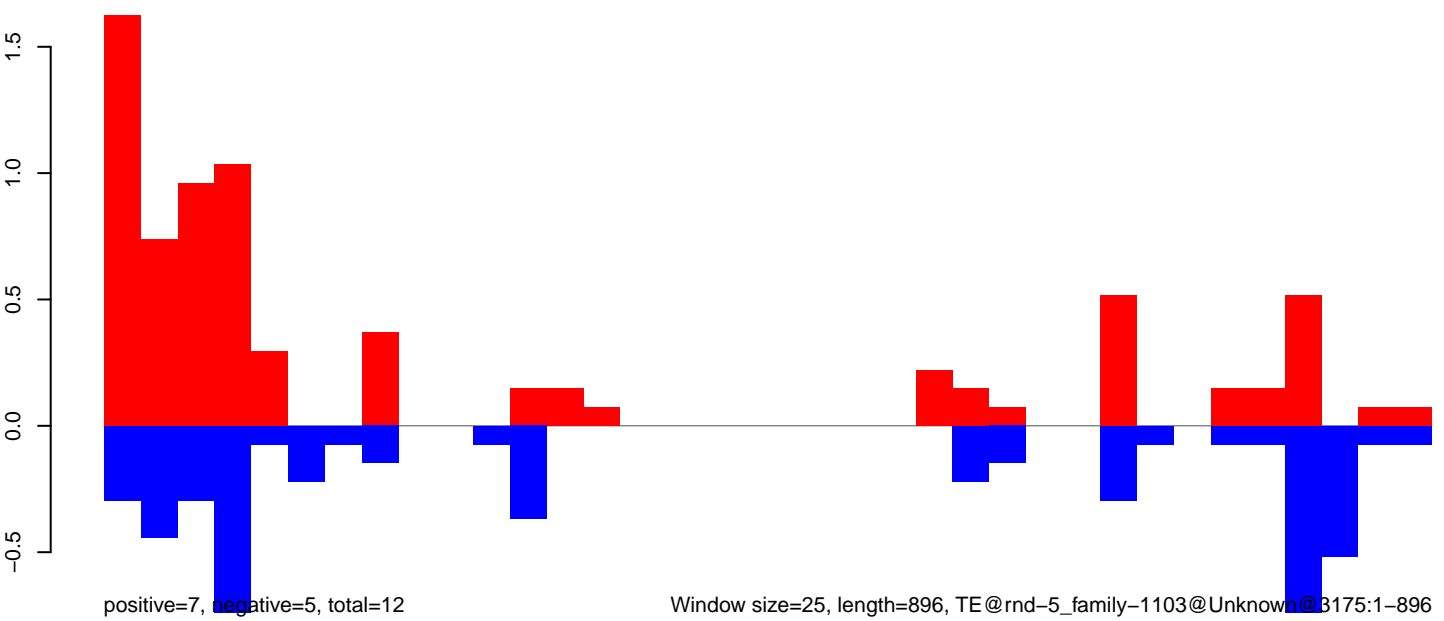
CuQuin_Testes_NL.18_23.rep



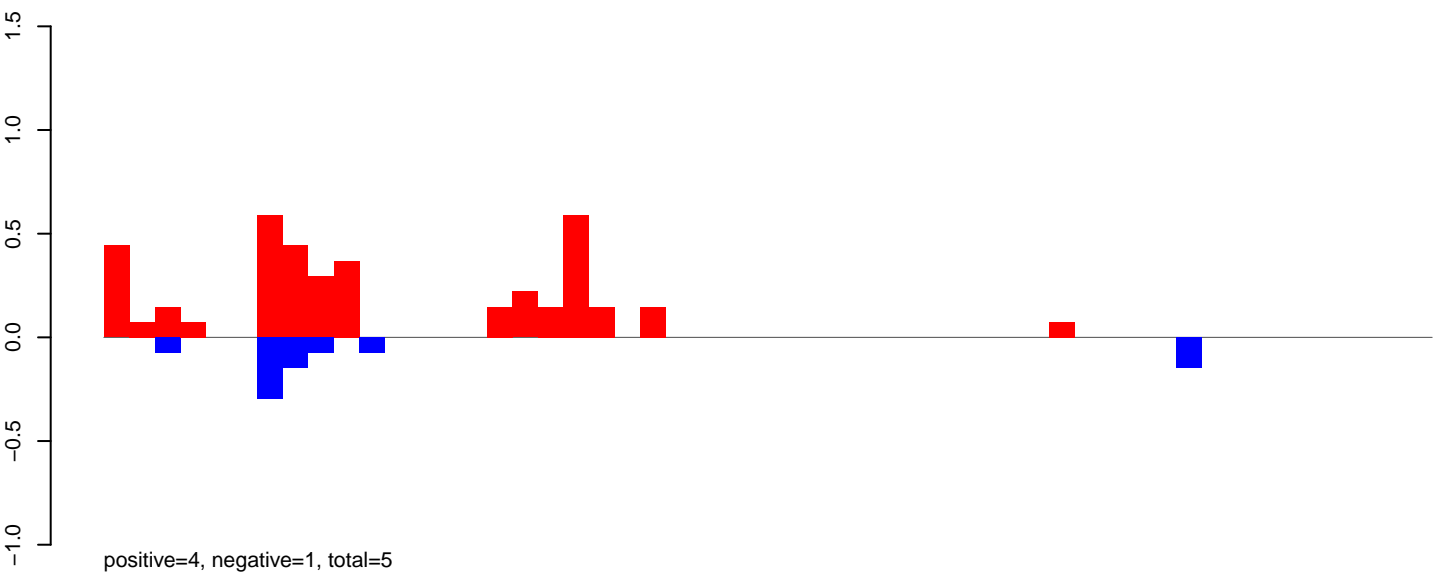
CuQuin_Testes_NL.24_35.rep



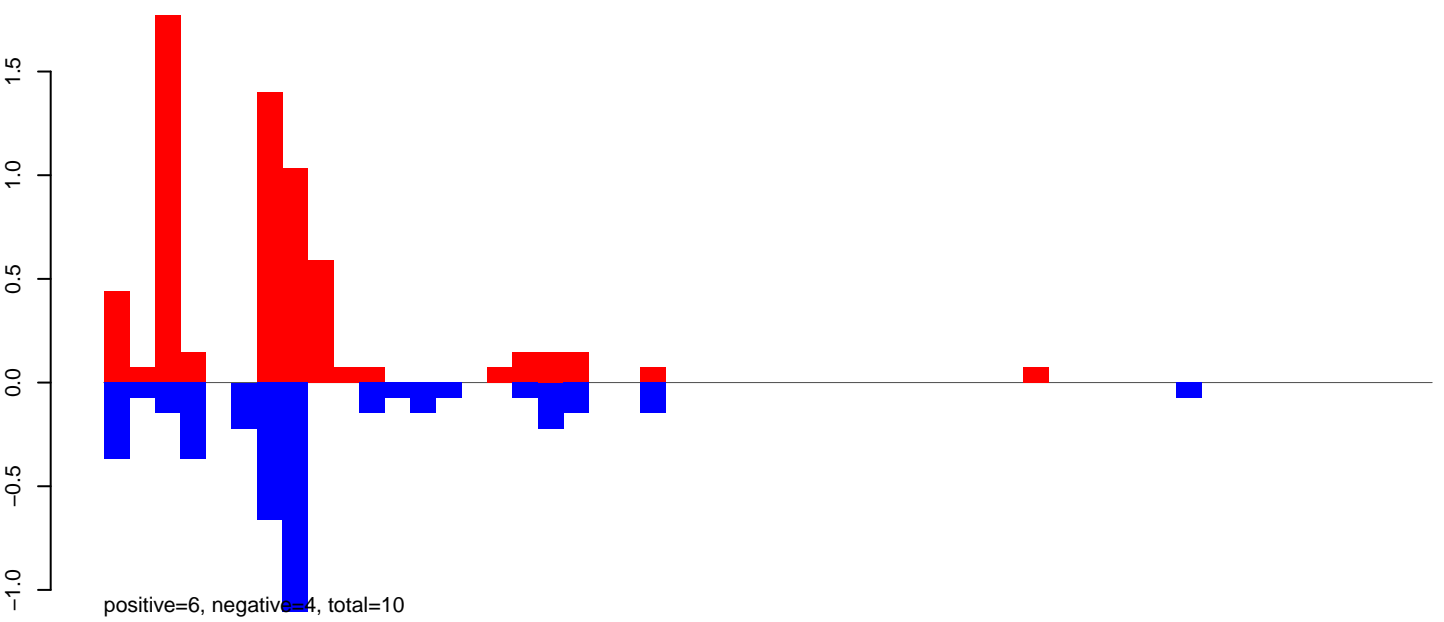
CuQuin_Testes_NL.rep



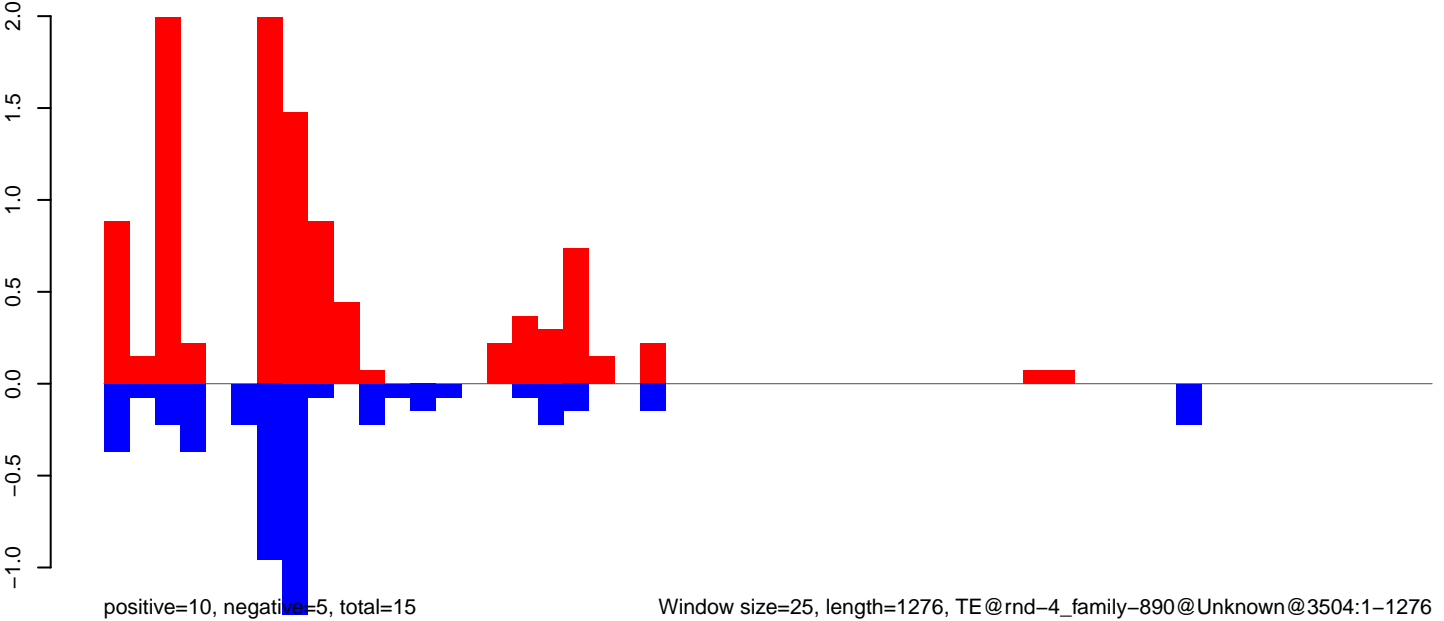
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



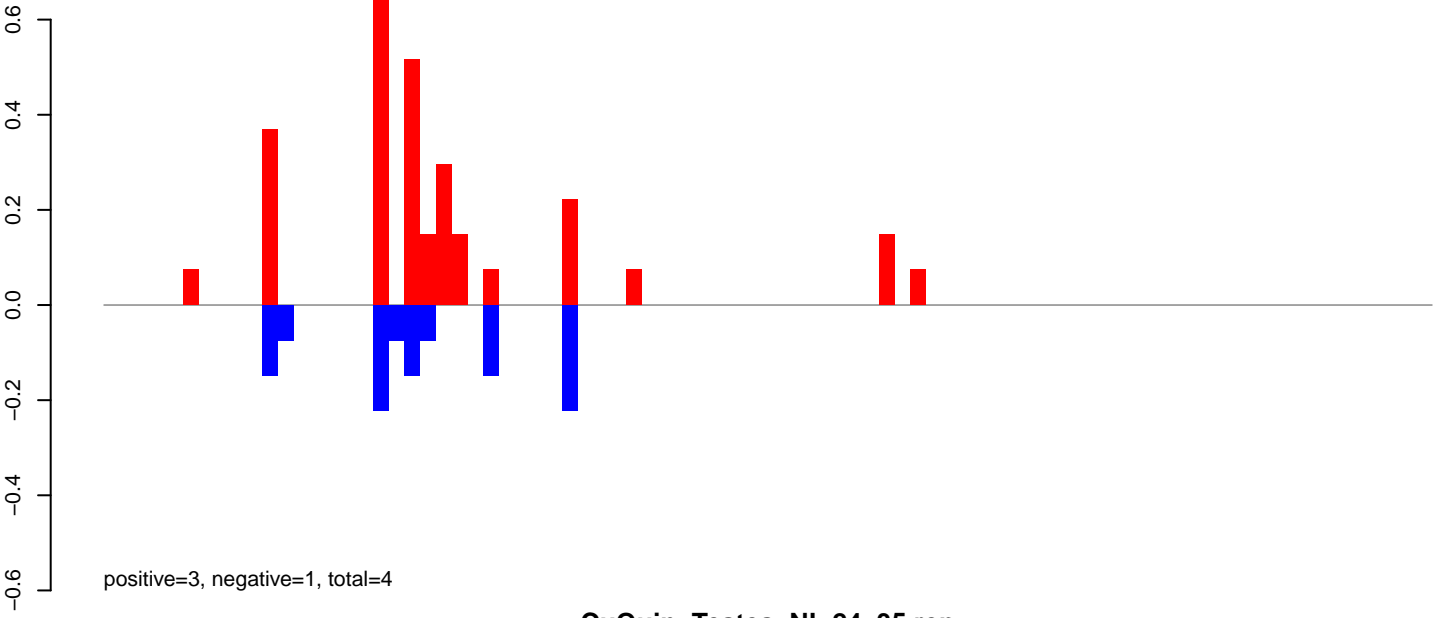
CuQuin_Testes_NL.rep



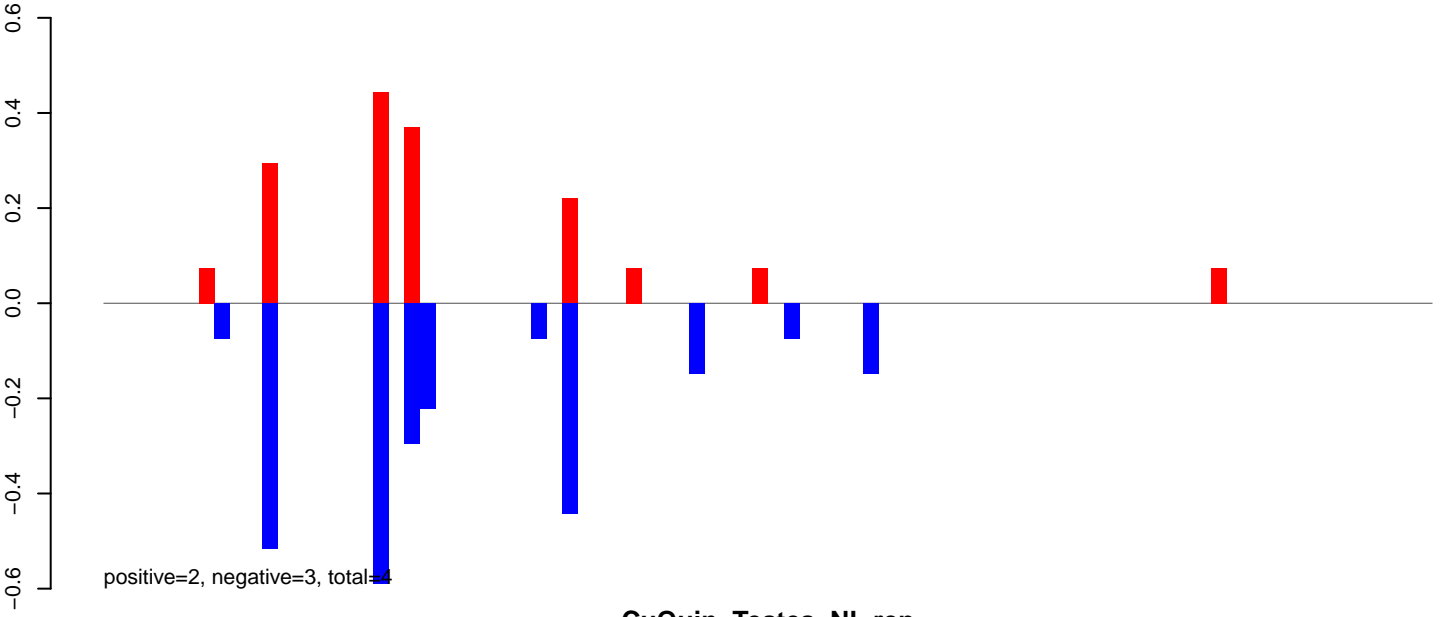
Window size=25, length=1276, TE@rnd-4_family-890@Unknown@3504:1-1276

0 200 400 600 800 1000 1200

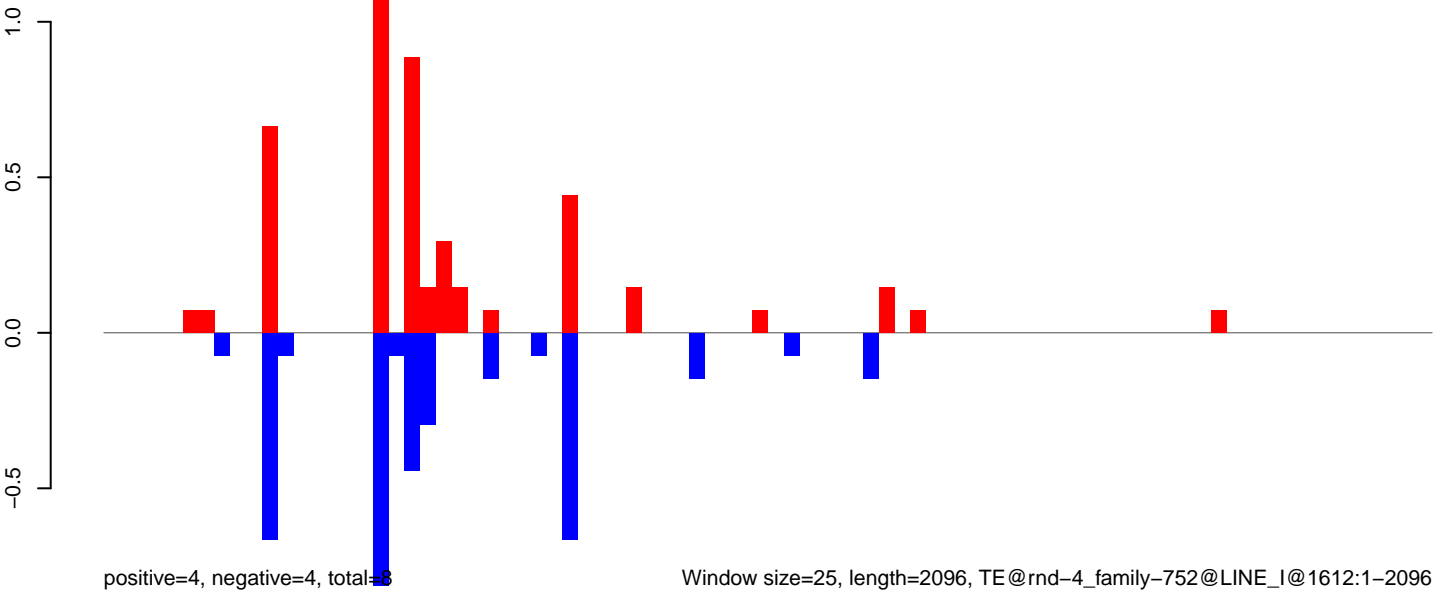
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



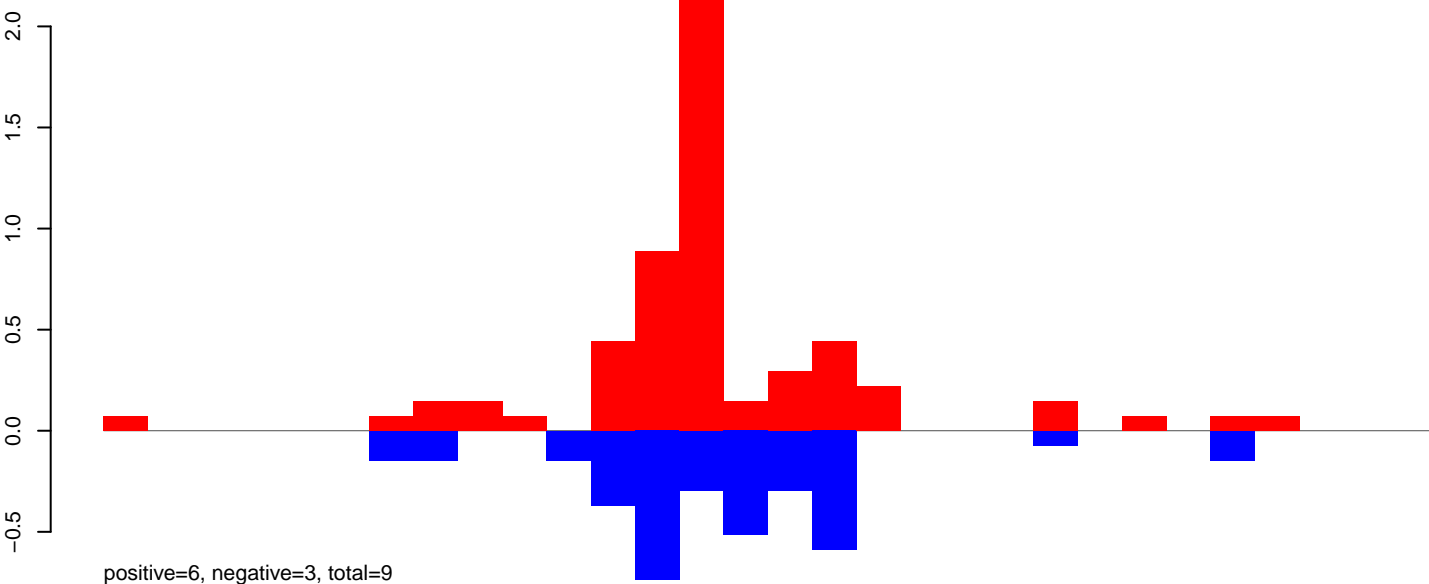
CuQuin_Testes_NL.rep



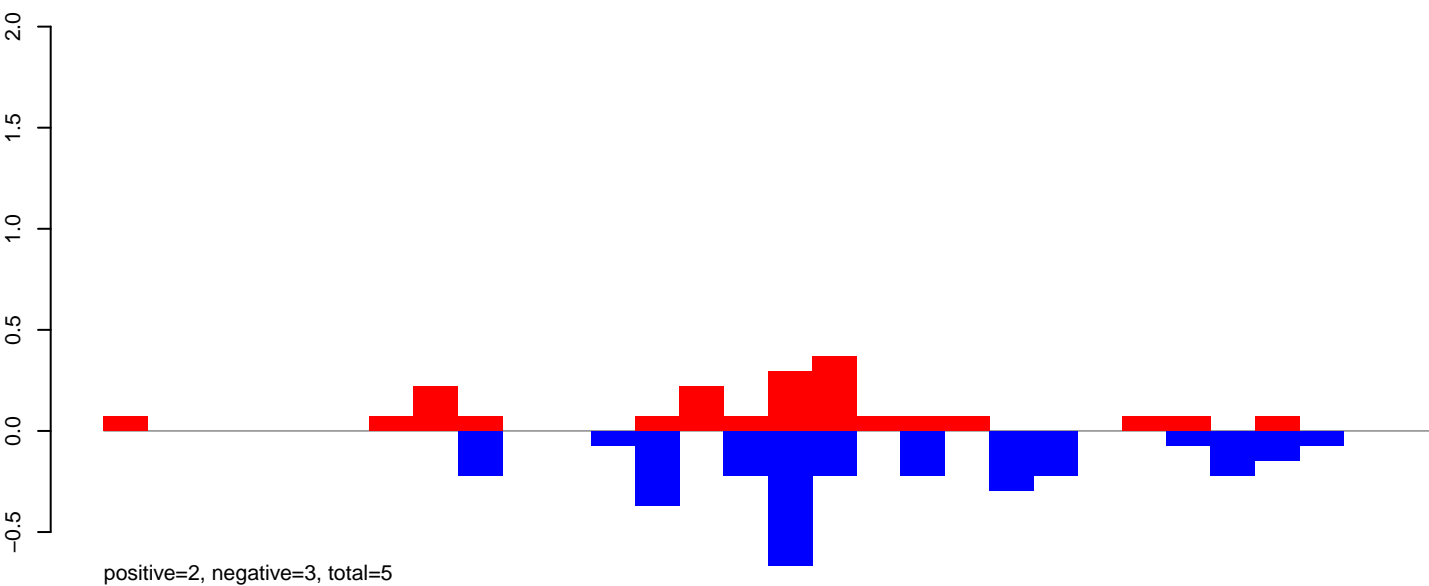
Window size=25, length=2096, TE@rnd-4_family-752@LINE_I@1612:1-2096

0 500 1000 1500 2000

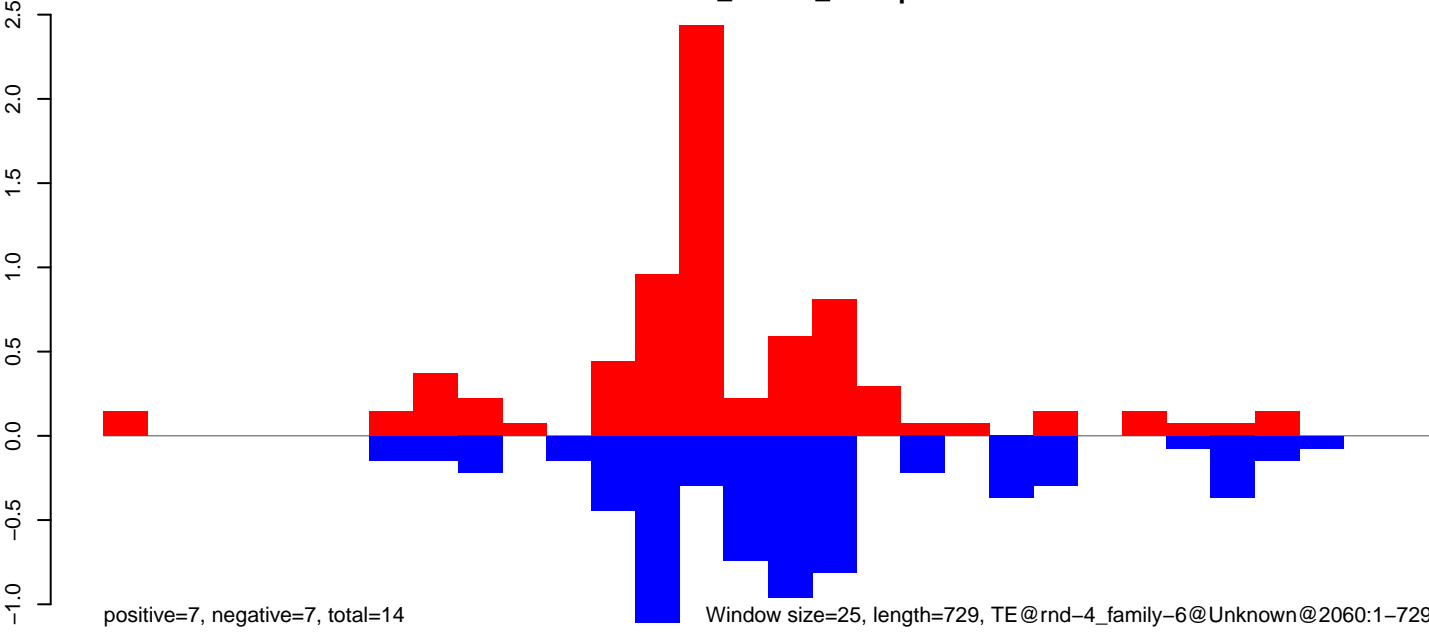
CuQuin_Testes_NL.18_23.rep



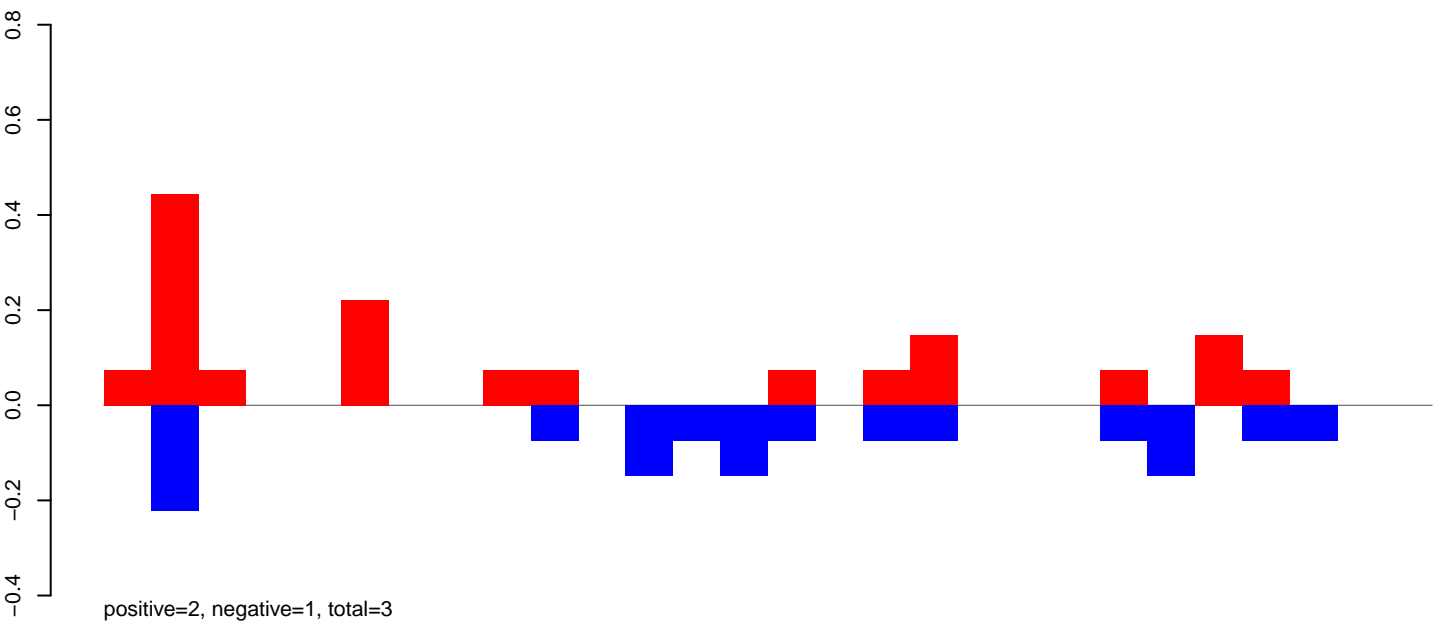
CuQuin_Testes_NL.24_35.rep



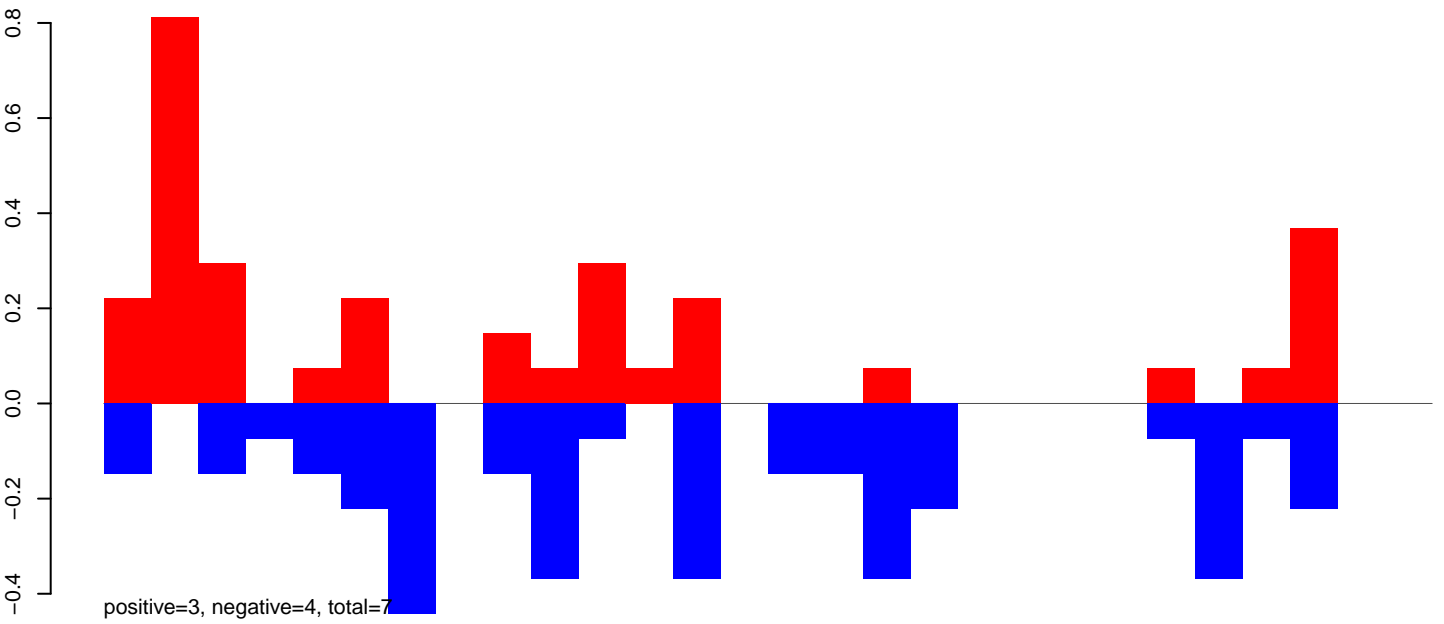
CuQuin_Testes_NL.rep



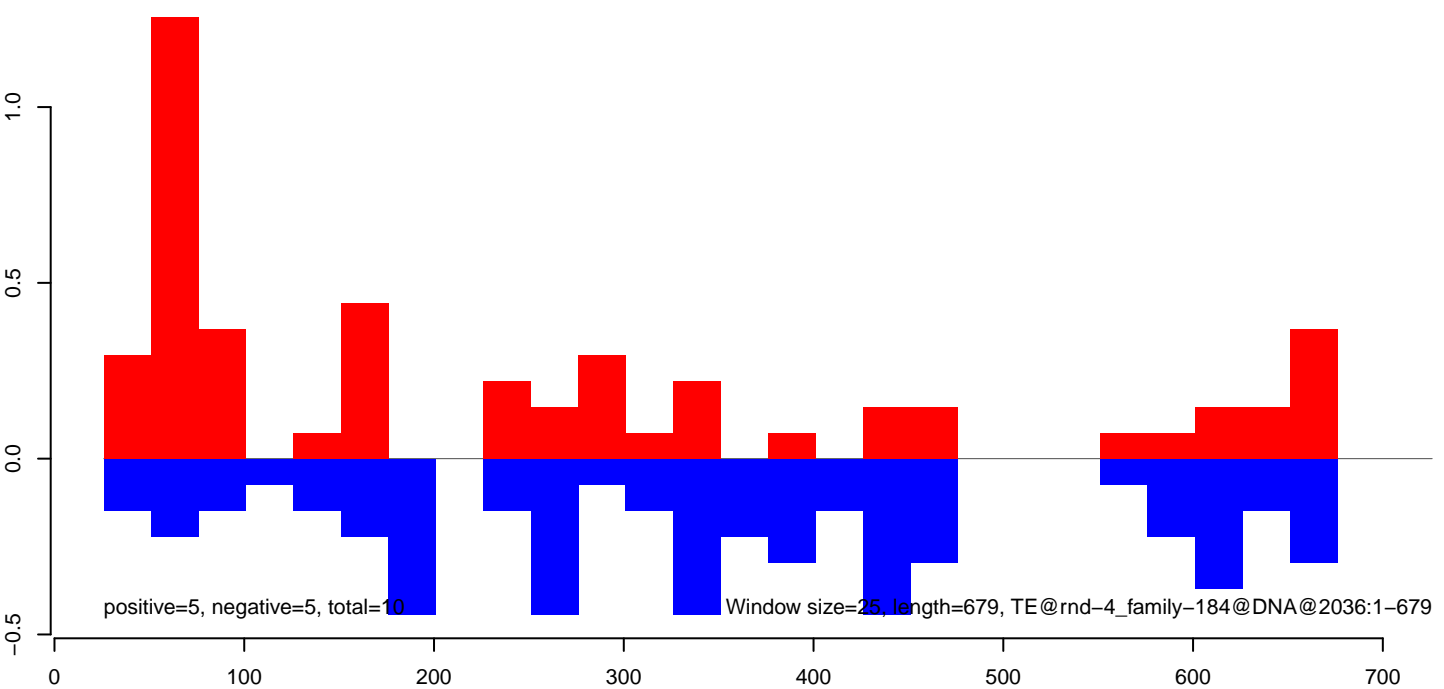
CuQuin_Testes_NL.18_23.rep



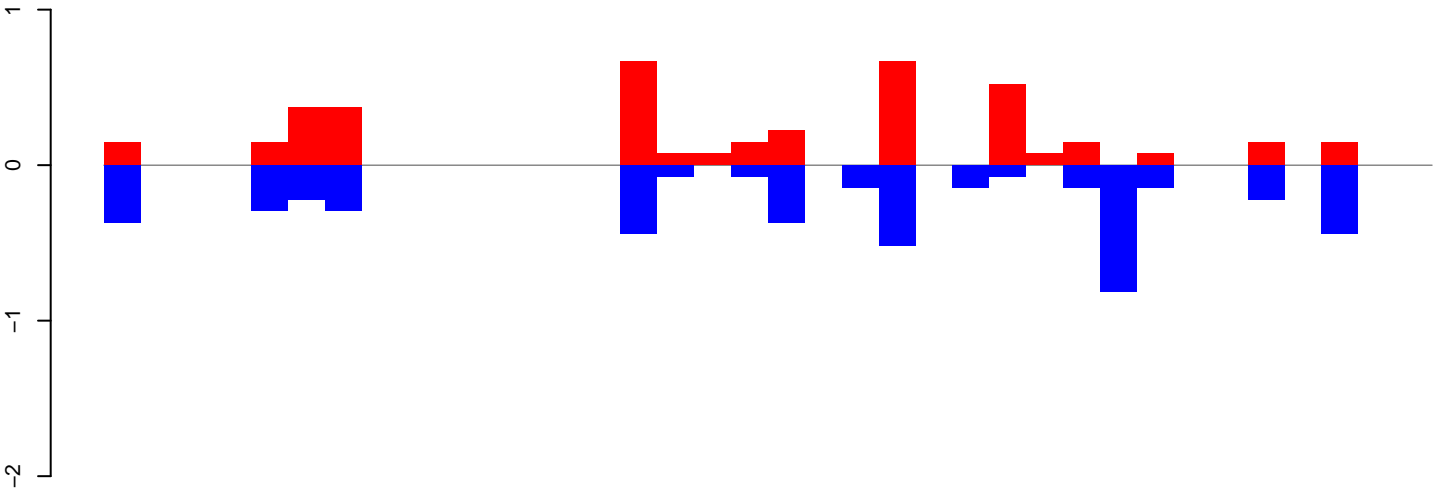
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

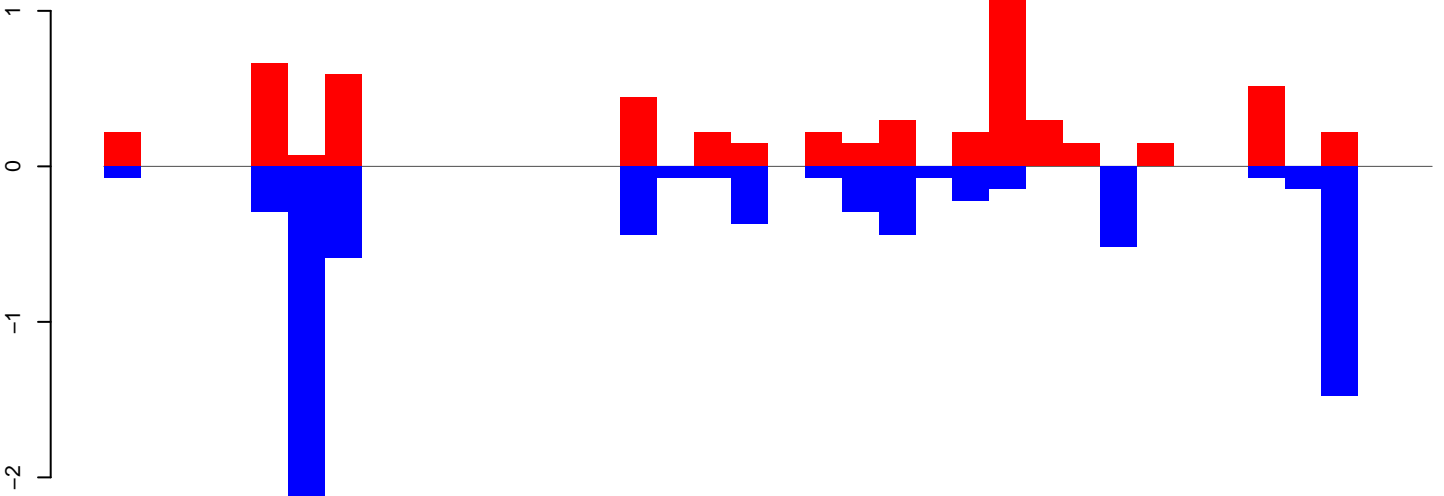


CuQuin_Testes_NL.18_23.rep



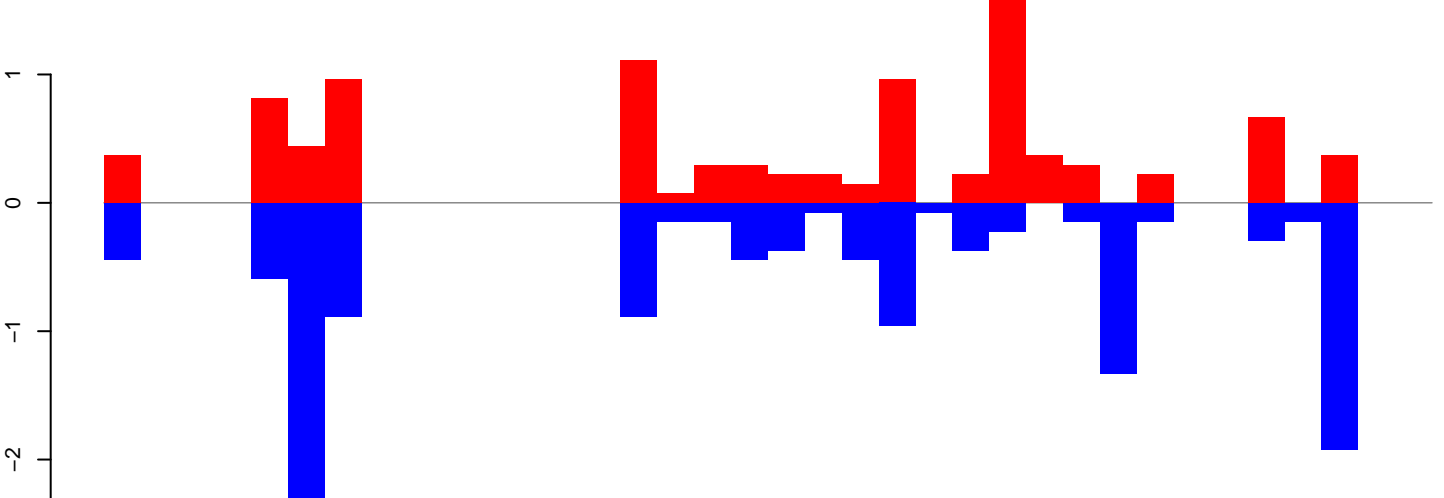
positive=4, negative=5, total=9

CuQuin_Testes_NL.24_35.rep



positive=6, negative=8, total=14

CuQuin_Testes_NL.rep



positive=10, negative=13, total=23

Window size=25, length=878, TE@rnd-4_family-151@Unknown@2786:1-878

0 200 400 600 800

CuQuin_Testes_NL.18_23.rep



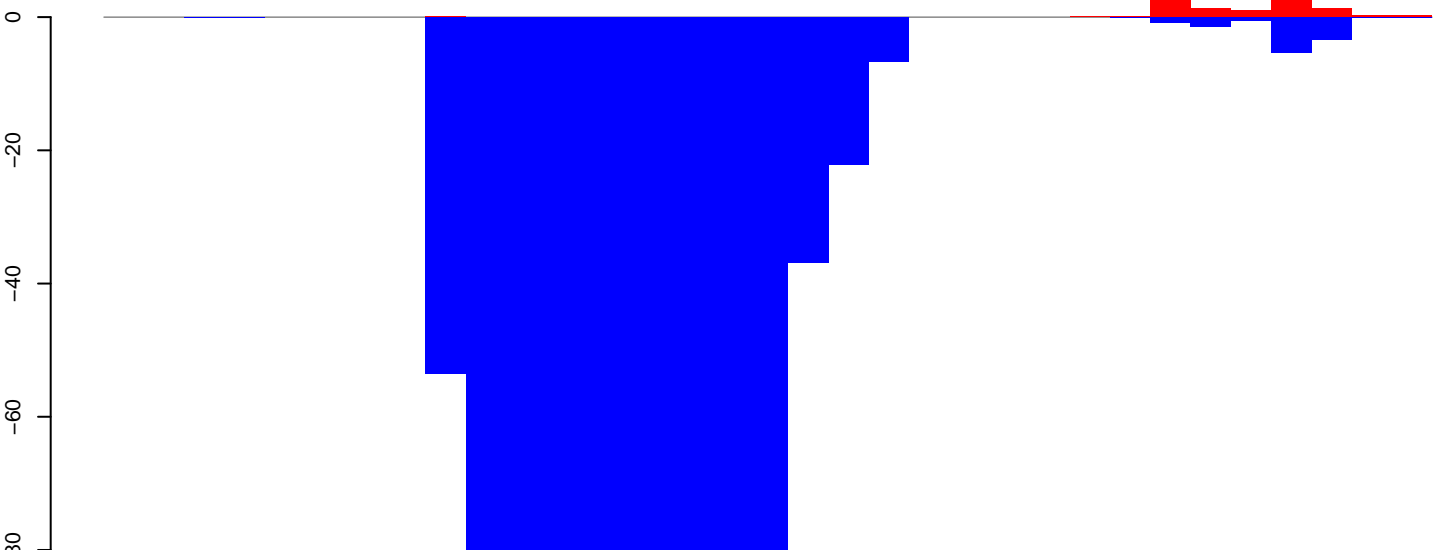
positive=2, negative=105, total=108

CuQuin_Testes_NL.24_35.rep



positive=8, negative=149, total=157

CuQuin_Testes_NL.rep

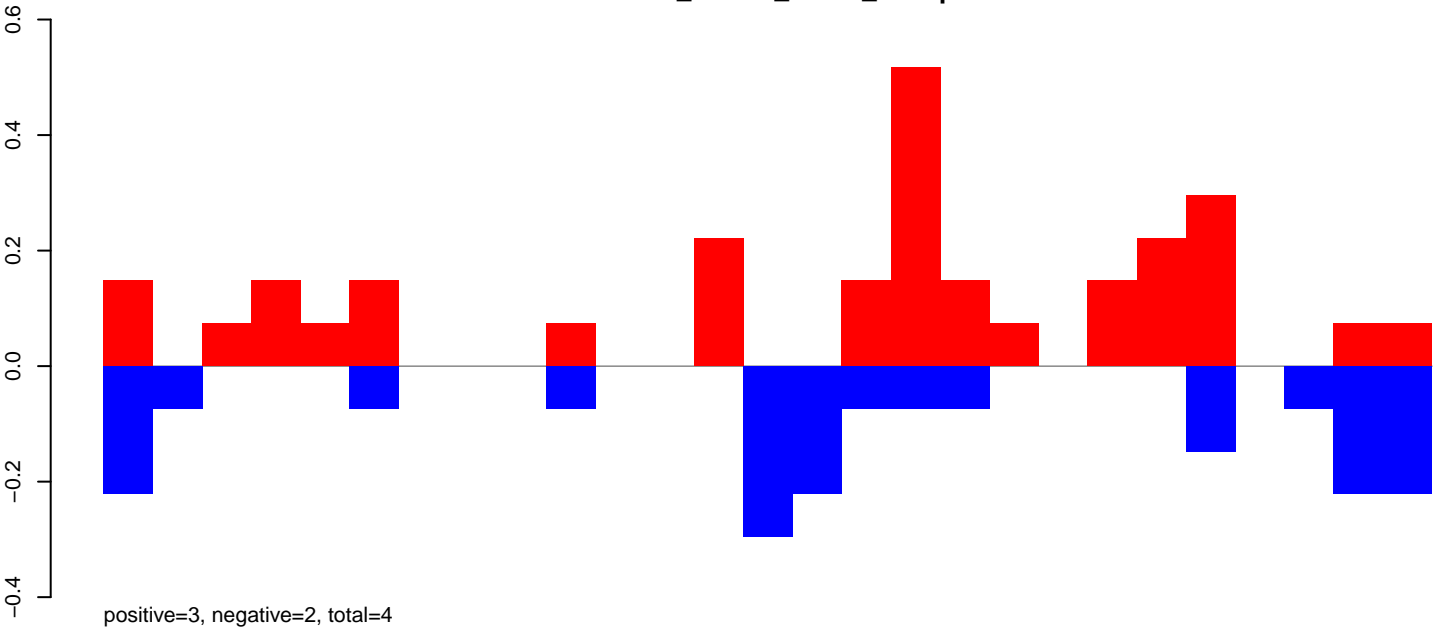


positive=14, negative=796, total=810

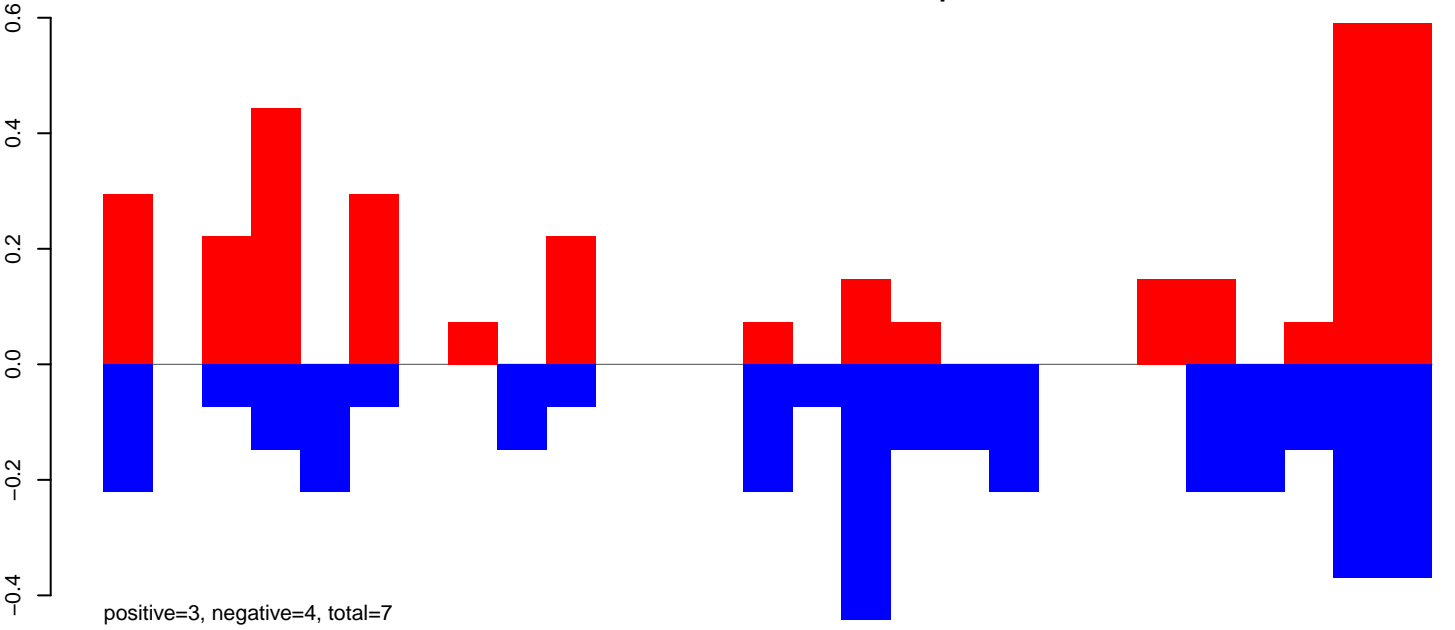
Window size=25, length=824, TE@rnd-3_family-322@Unknown@142:1-824

0 200 400 600 800

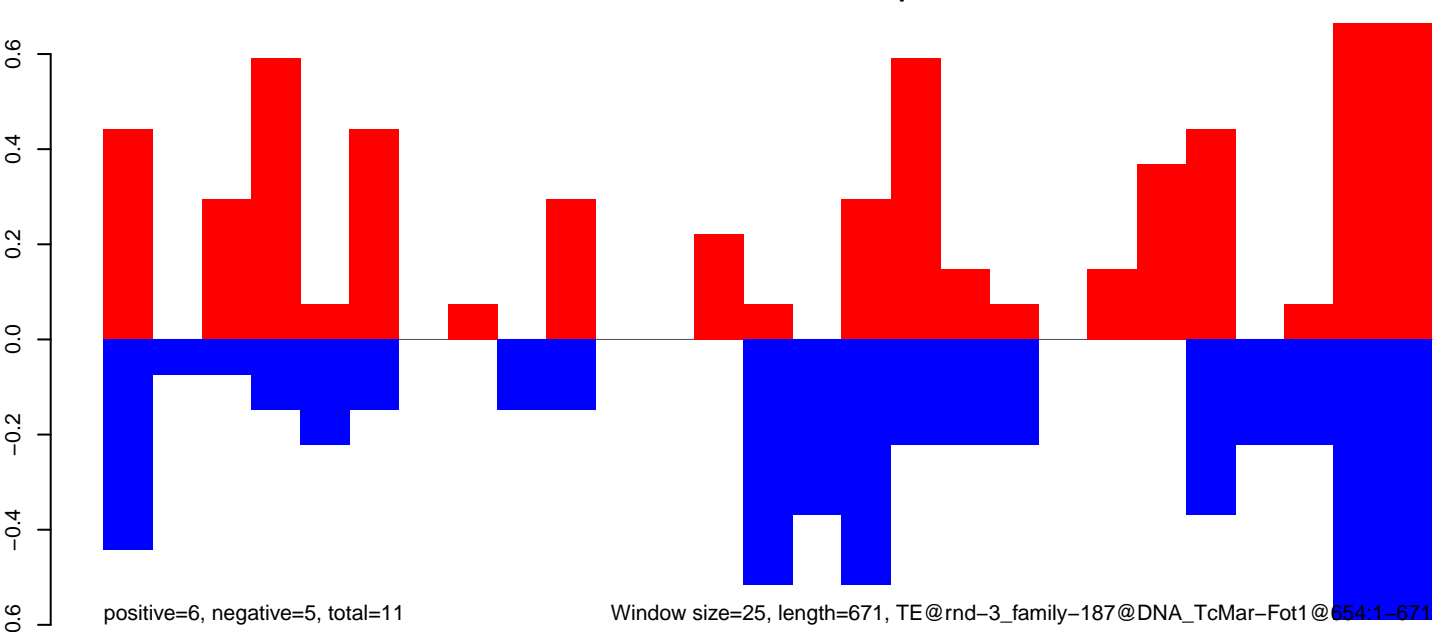
CuQuin_Testes_NL.18_23.rep



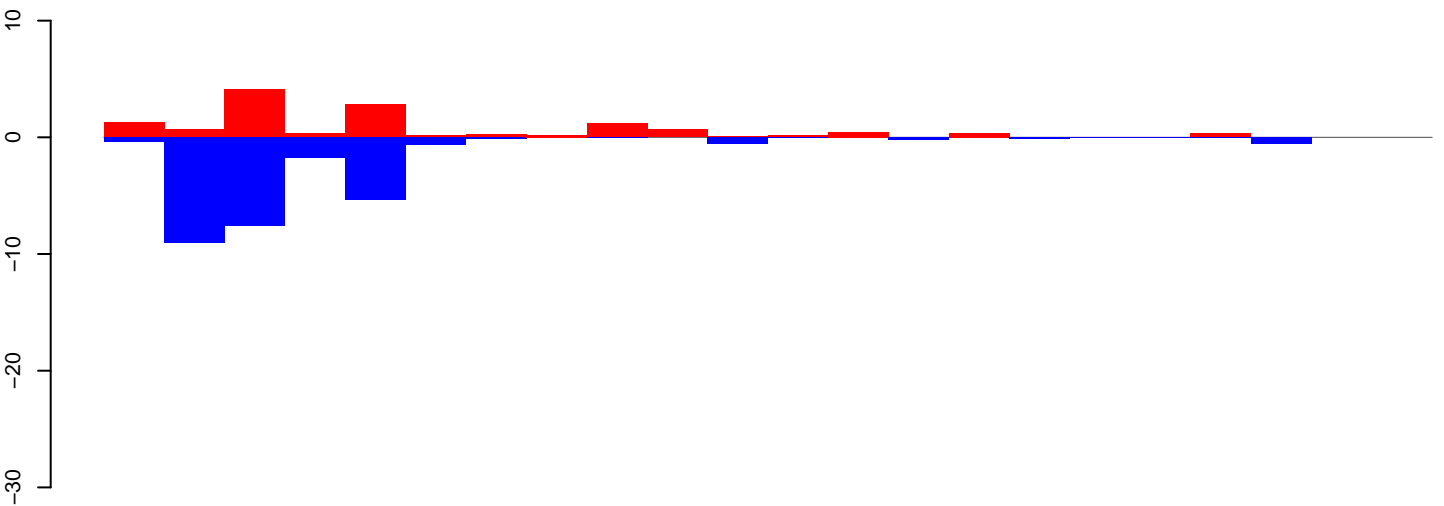
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

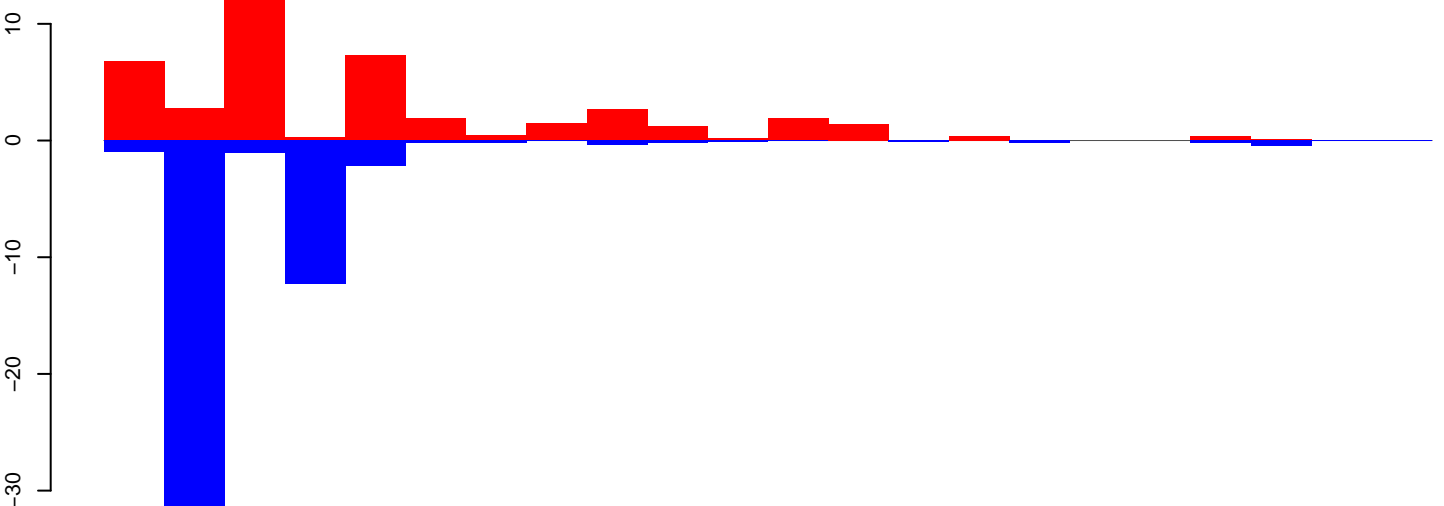


CuQuin_Testes_NL.18_23.rep



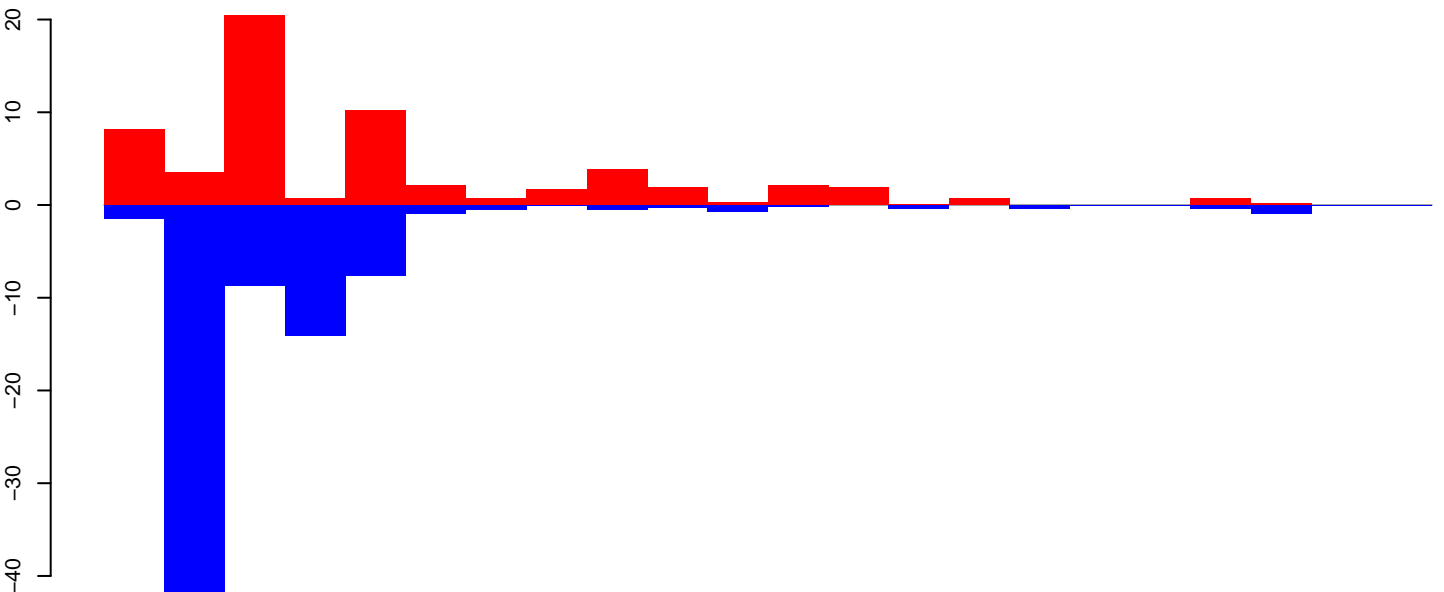
positive=13, negative=27, total=40

CuQuin_Testes_NL.24_35.rep



positive=46, negative=54, total=100

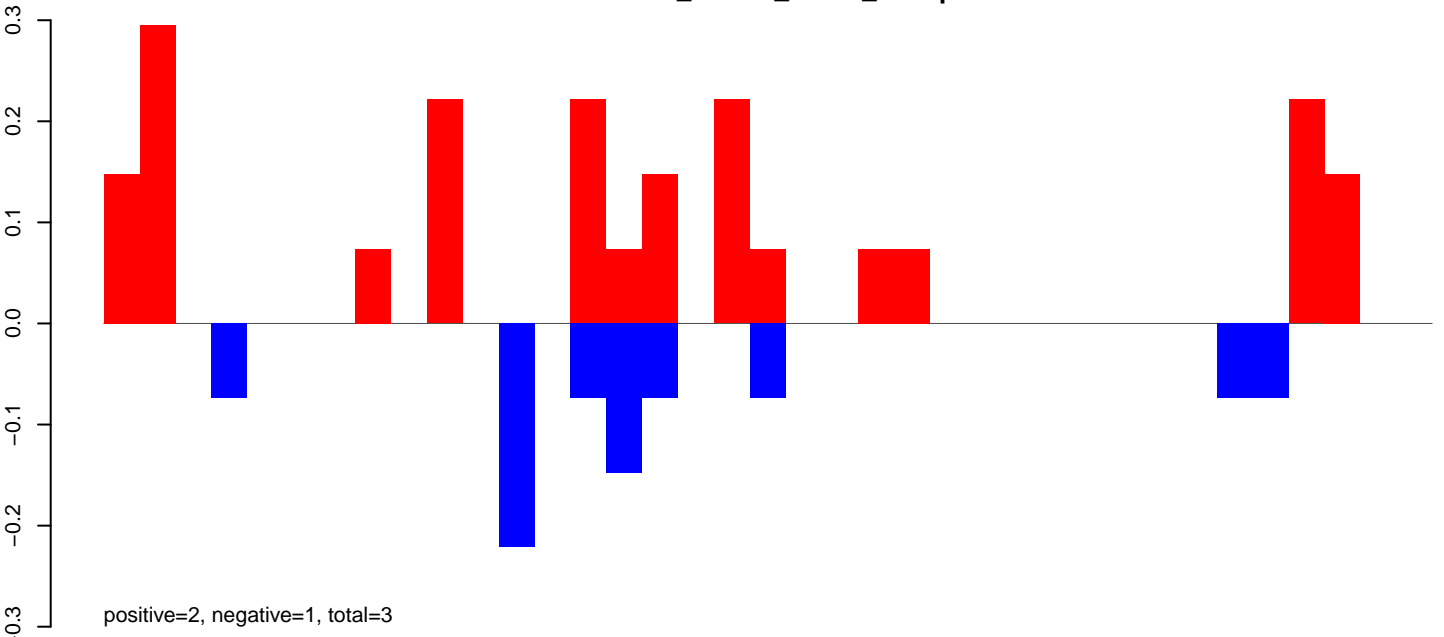
CuQuin_Testes_NL.rep



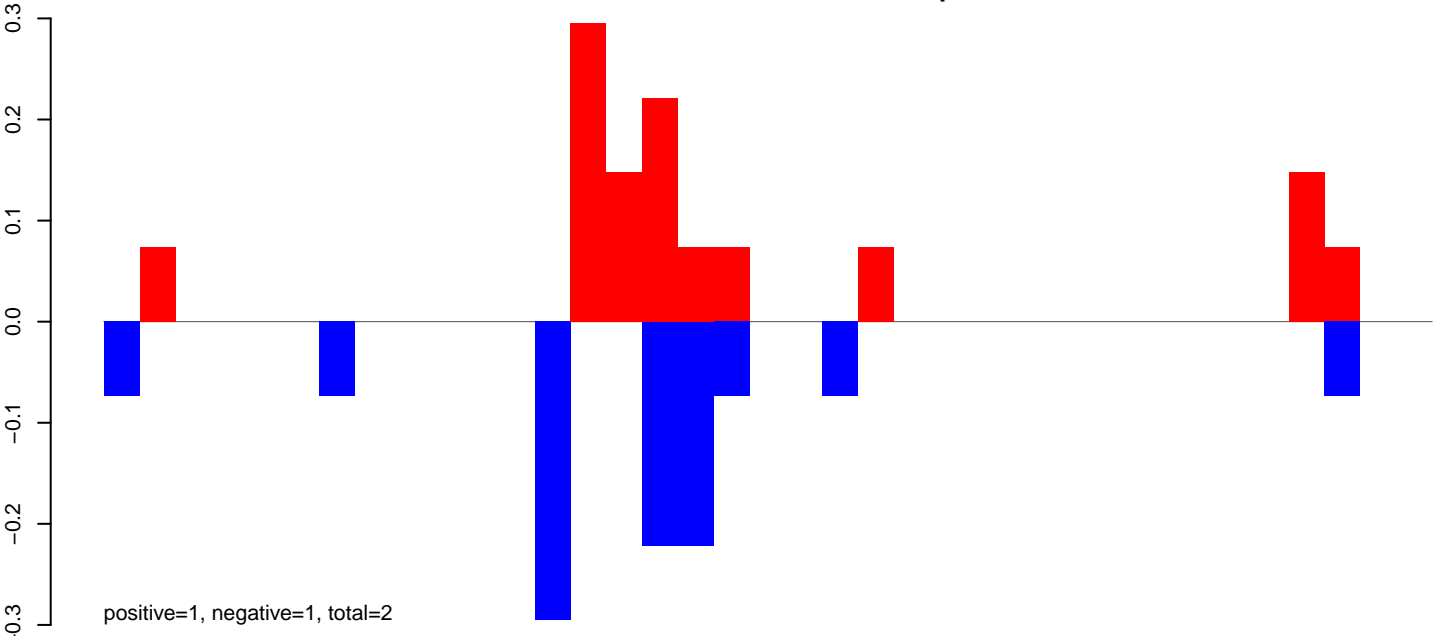
positive=59, negative=81, total=140

Window size=25, length=533, TE@rnd-5_family-2852@Unknown@3018:1-533

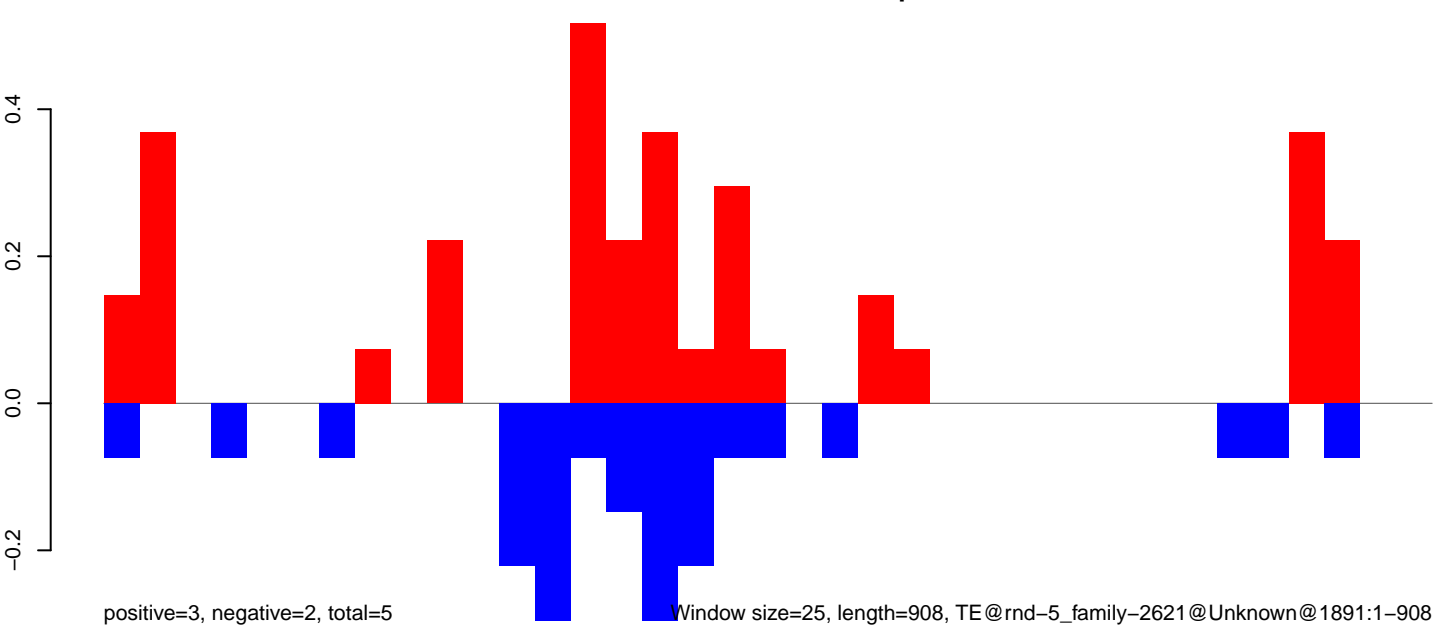
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

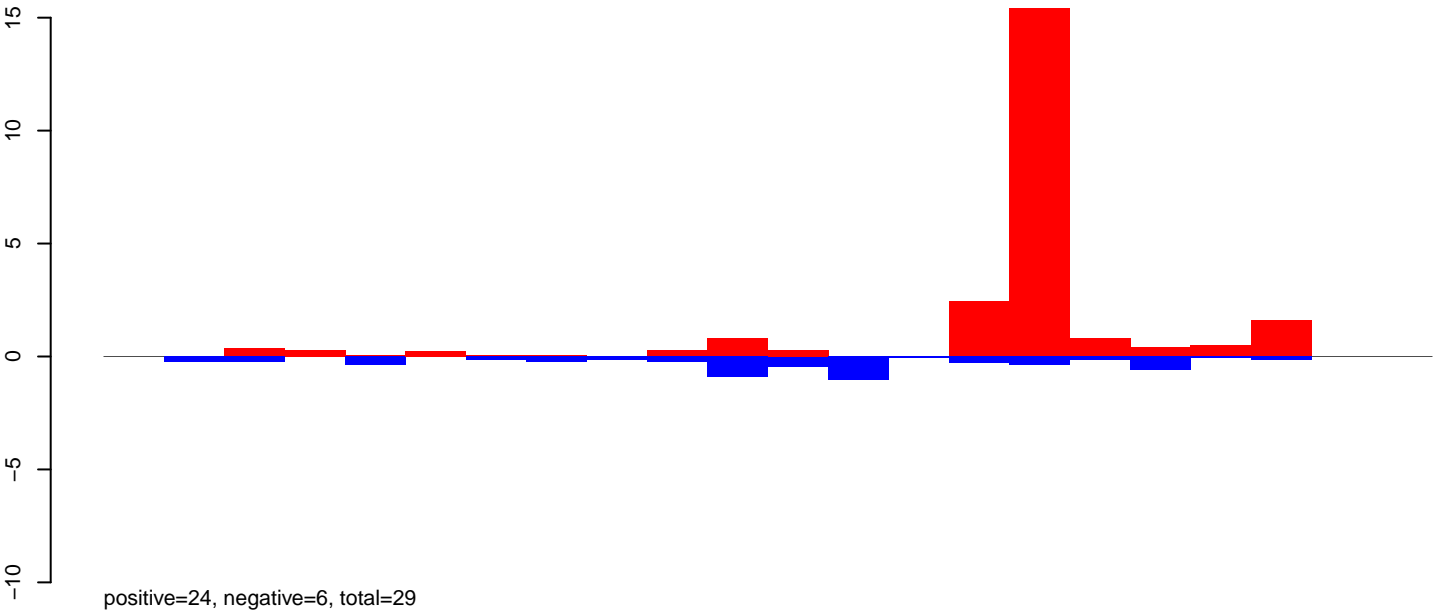


CuQuin_Testes_NL.rep

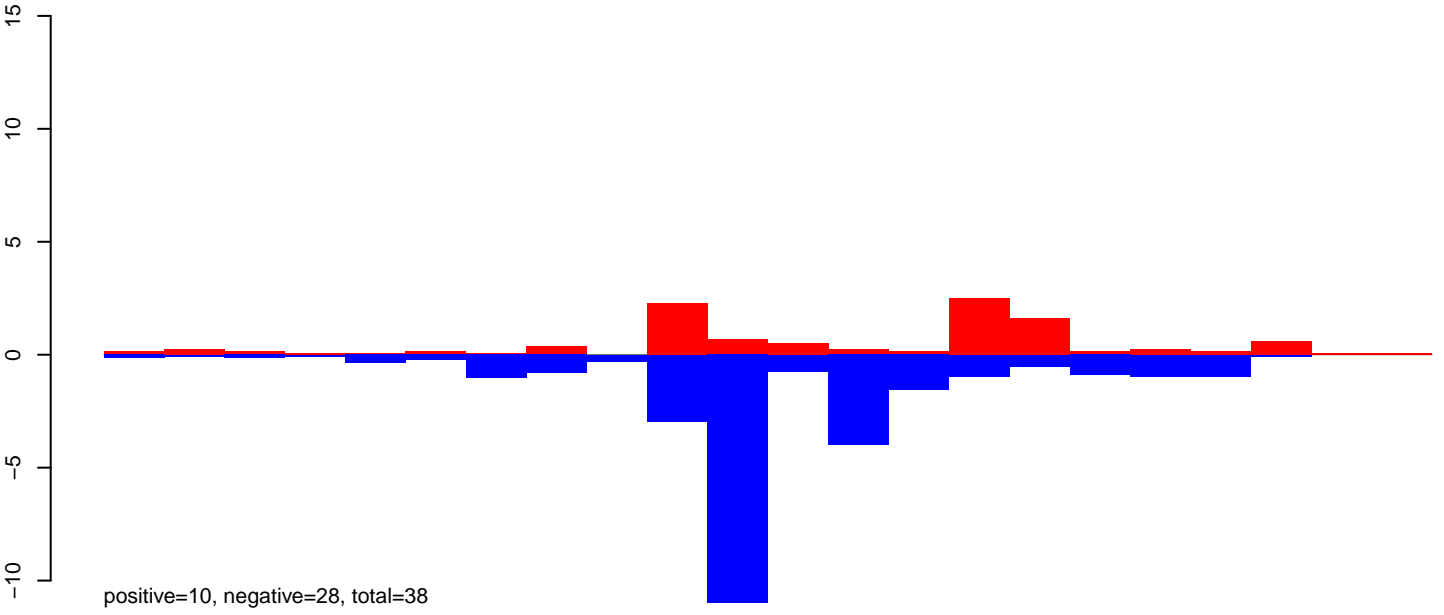


Window size=25, length=908, TE@rnd-5_family-2621@Unknown@1891:1-908

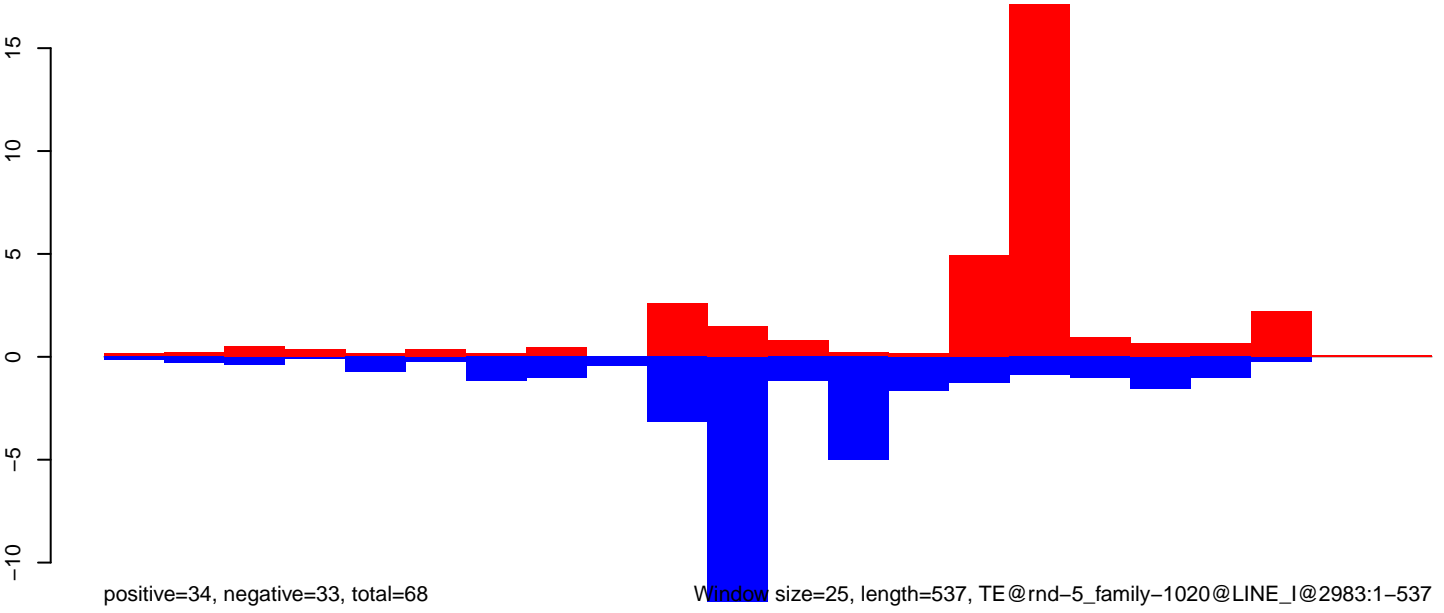
CuQuin_Testes_NL.18_23.rep



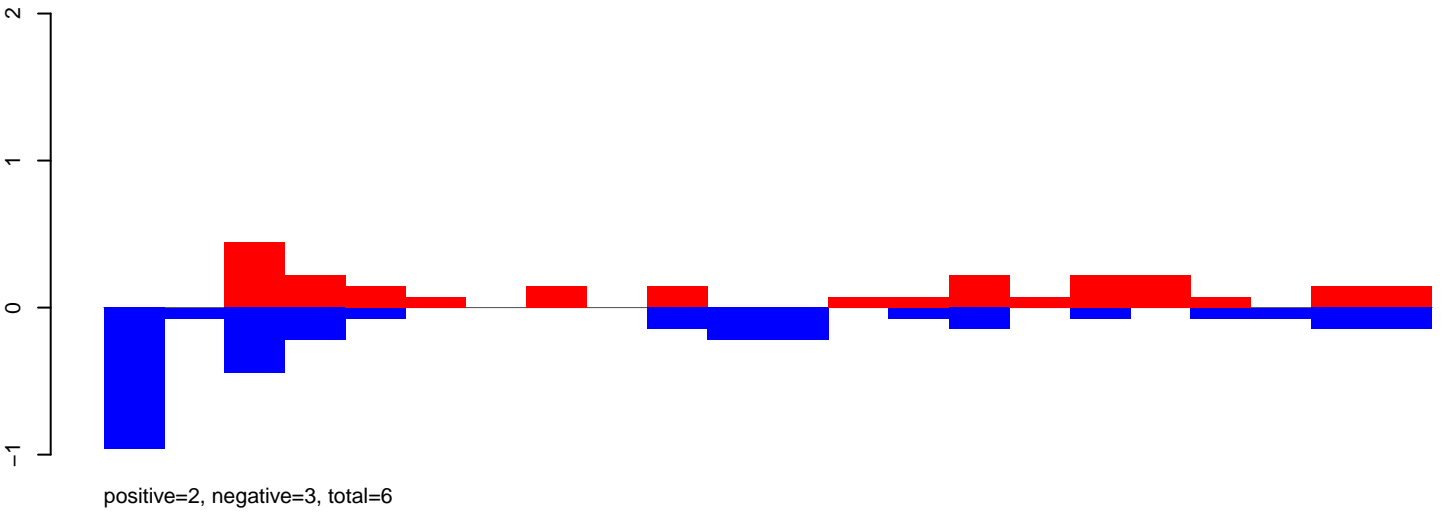
CuQuin_Testes_NL.24_35.rep



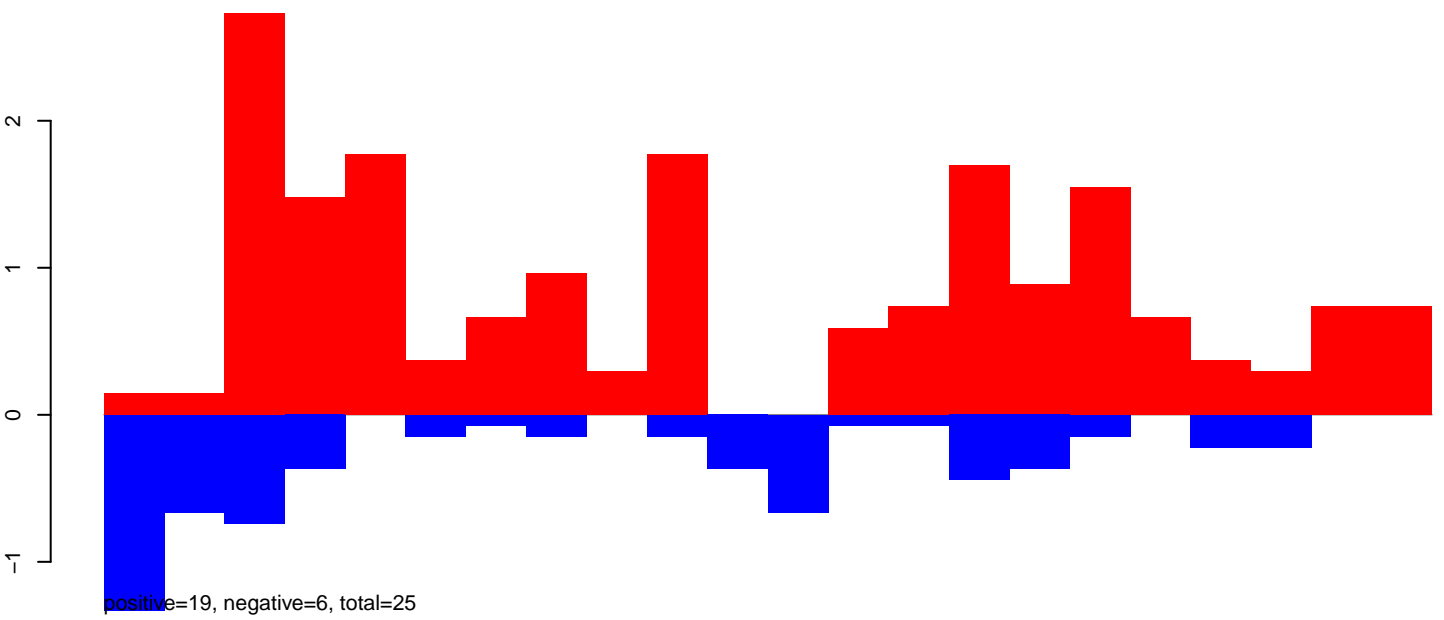
CuQuin_Testes_NL.rep



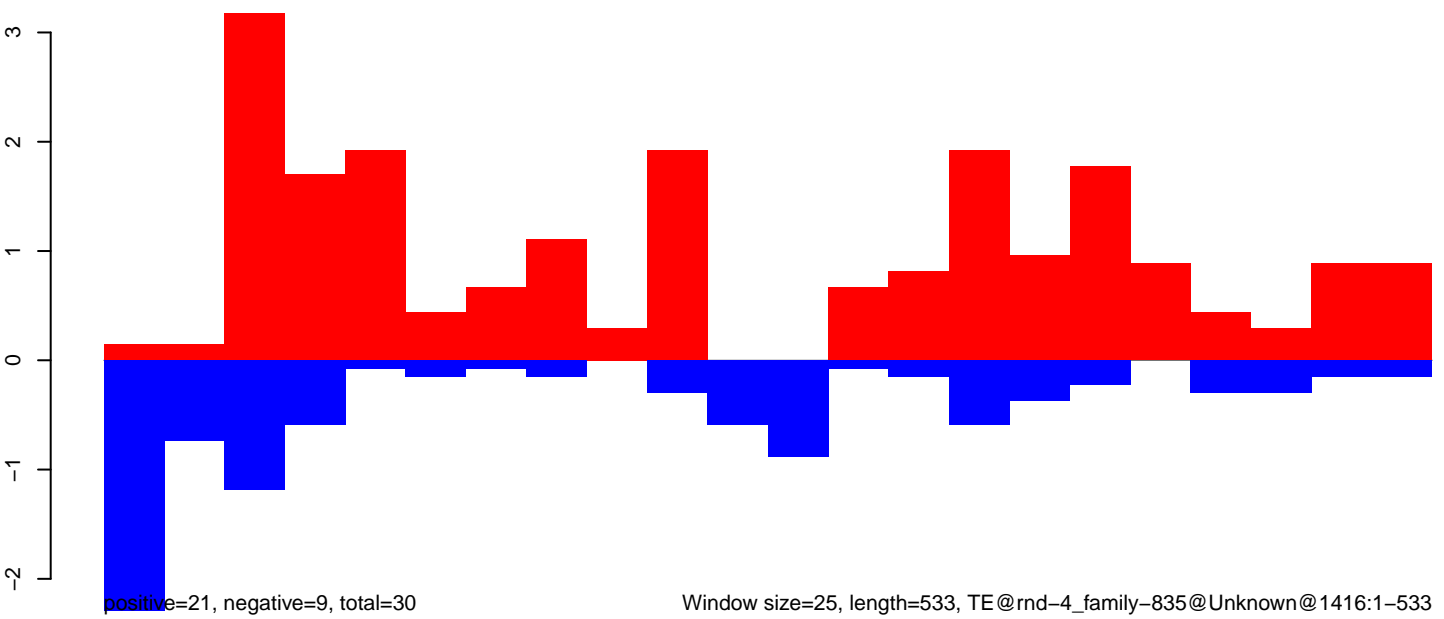
CuQuin_Testes_NL.18_23.rep



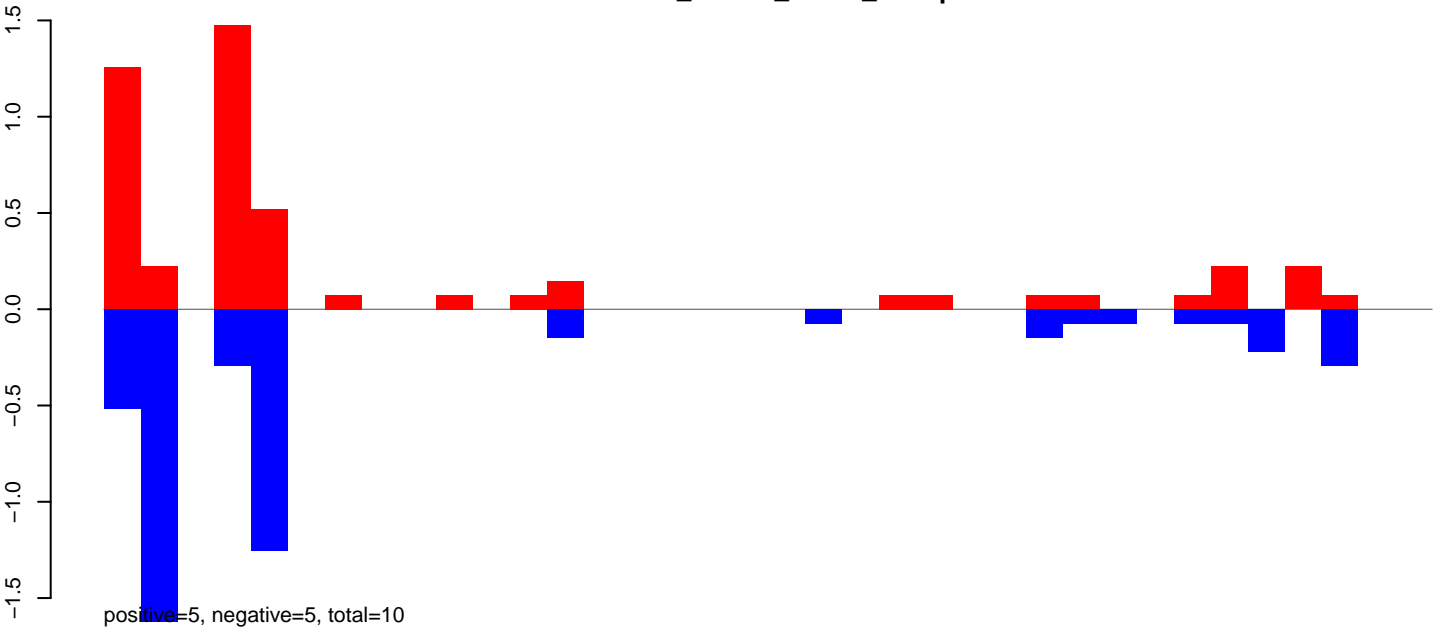
CuQuin_Testes_NL.24_35.rep



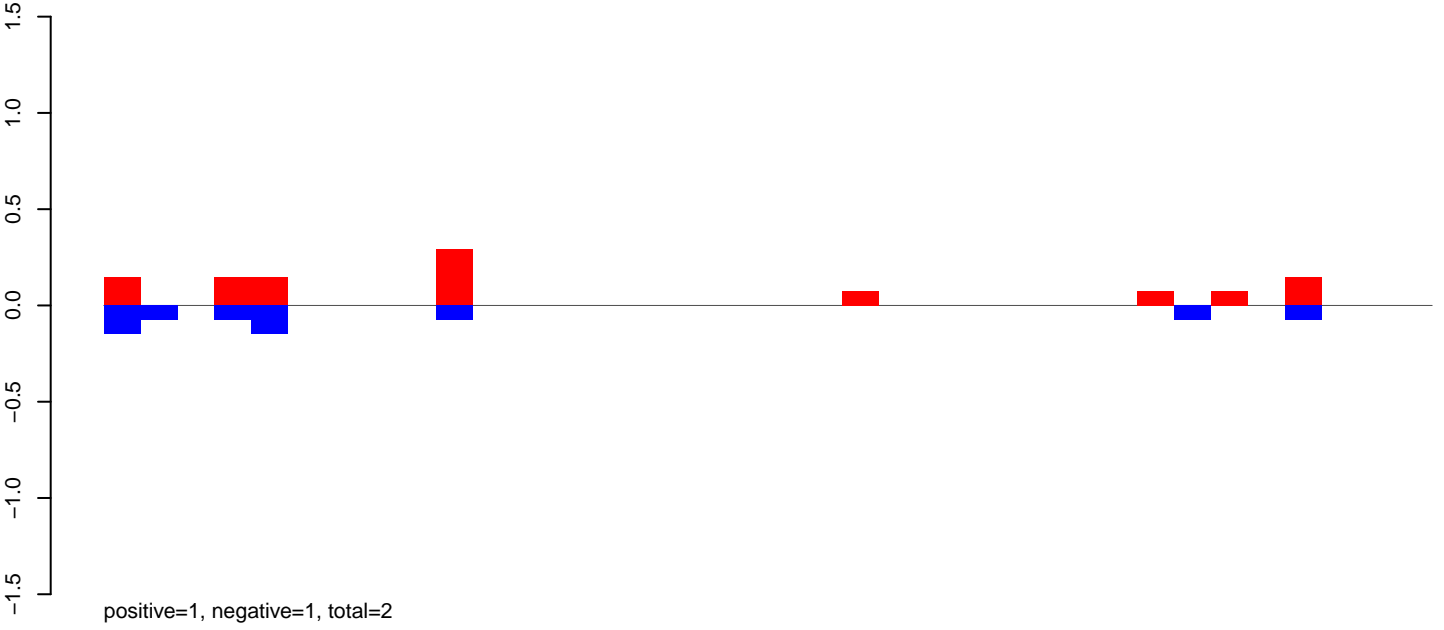
CuQuin_Testes_NL.rep



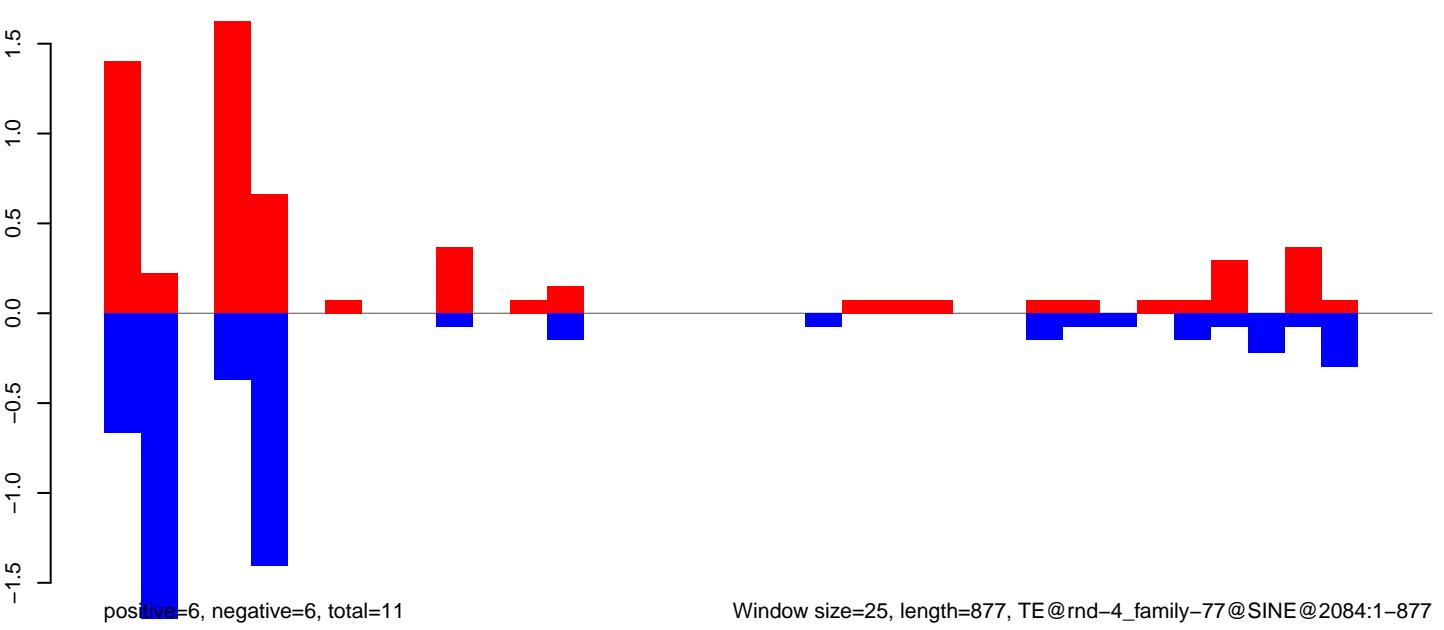
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



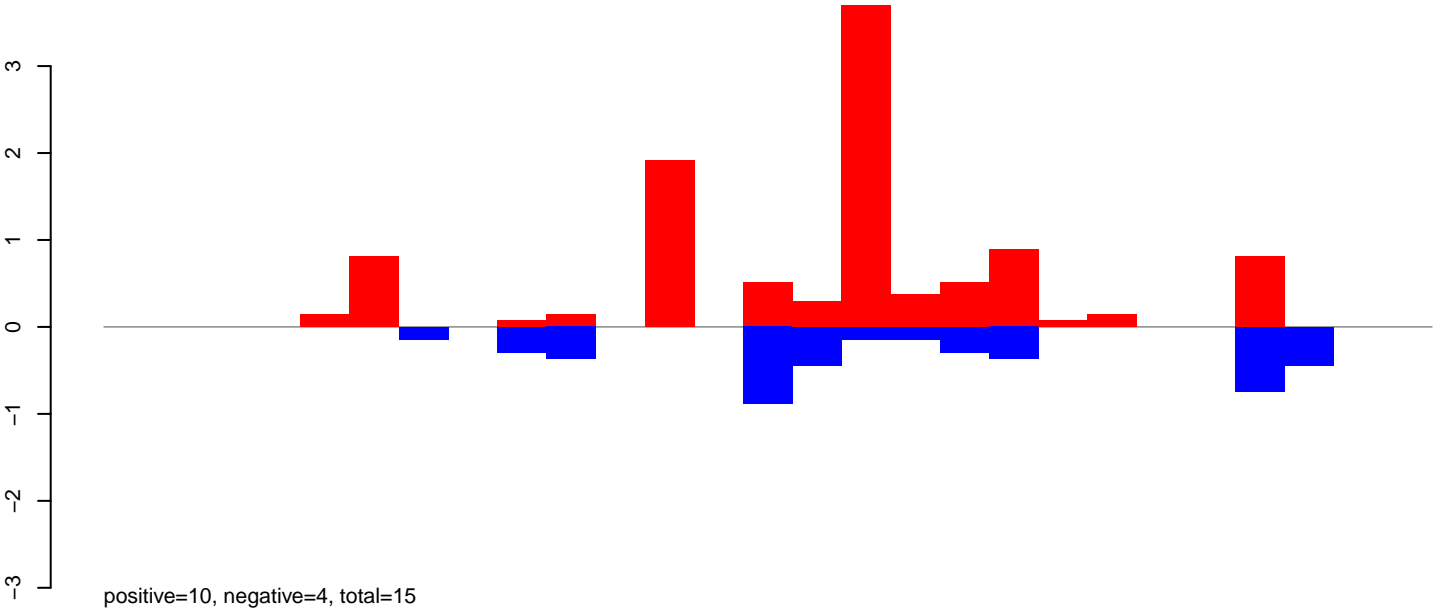
CuQuin_Testes_NL.rep



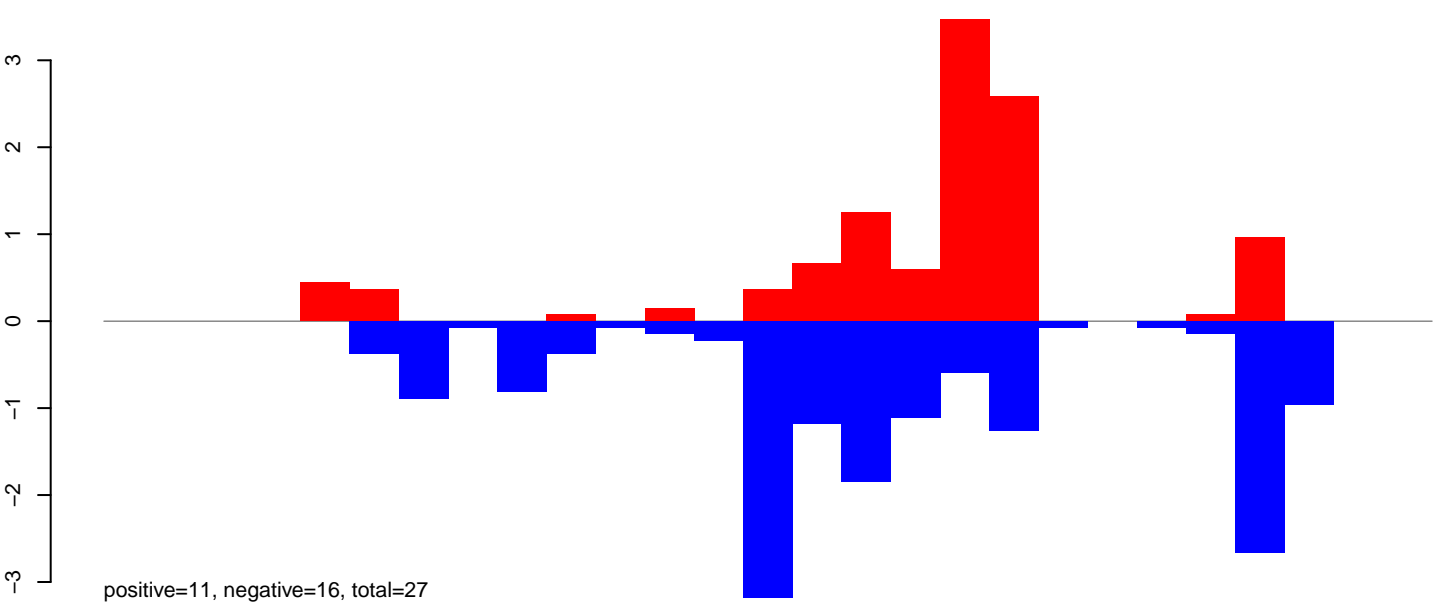
Window size=25, length=877, TE@rnd-4_family-77@SINE@2084:1-877

0 200 400 600 800

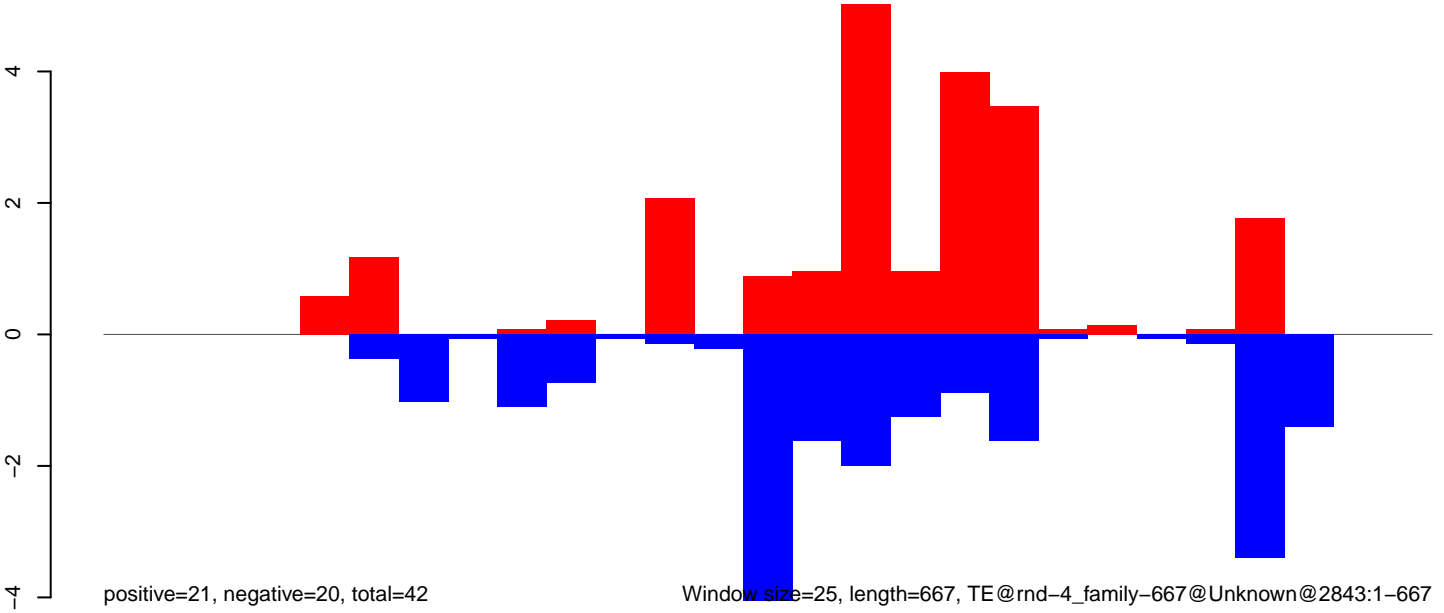
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

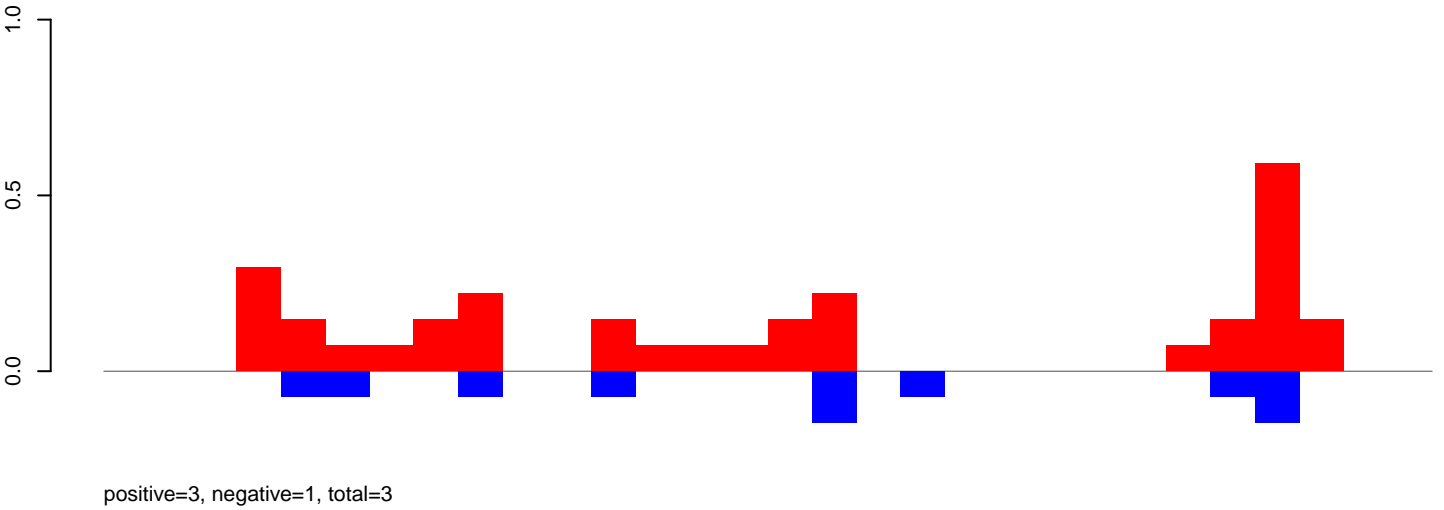


CuQuin_Testes_NL.rep

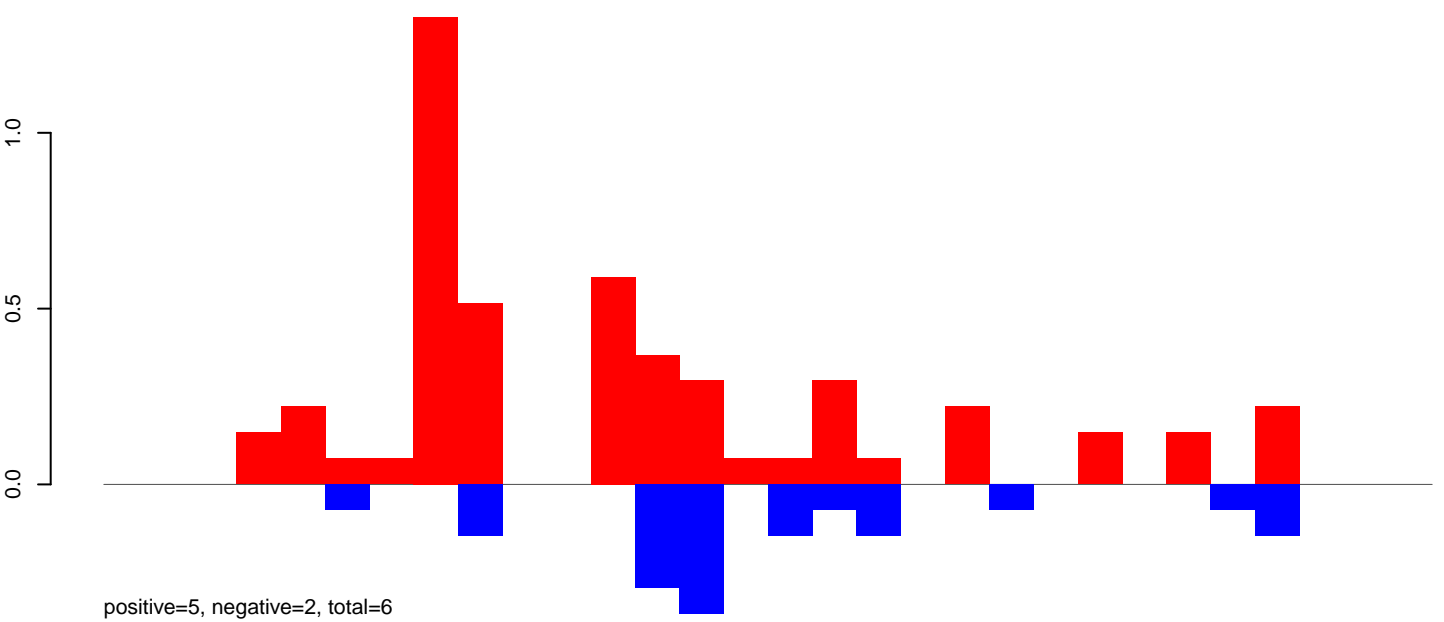


Window size=25, length=667, TE@rnd-4_family-667@Unknown@2843:1-667

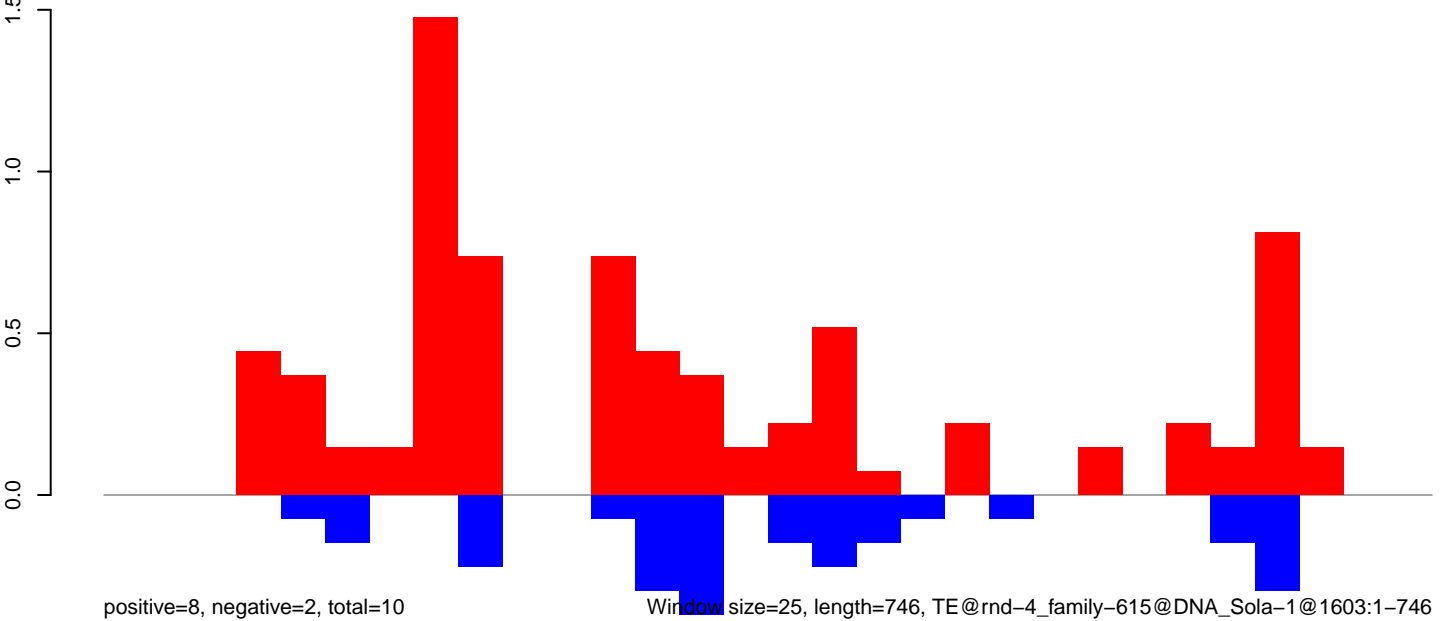
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

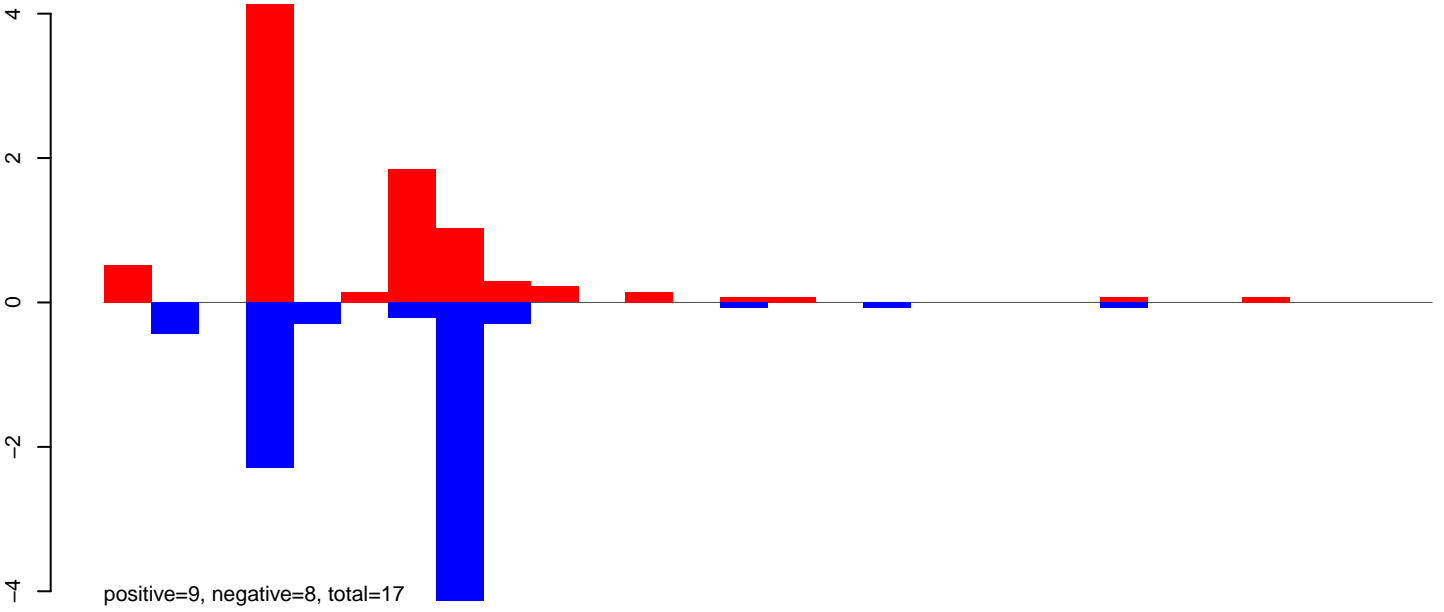


CuQuin_Testes_NL.rep

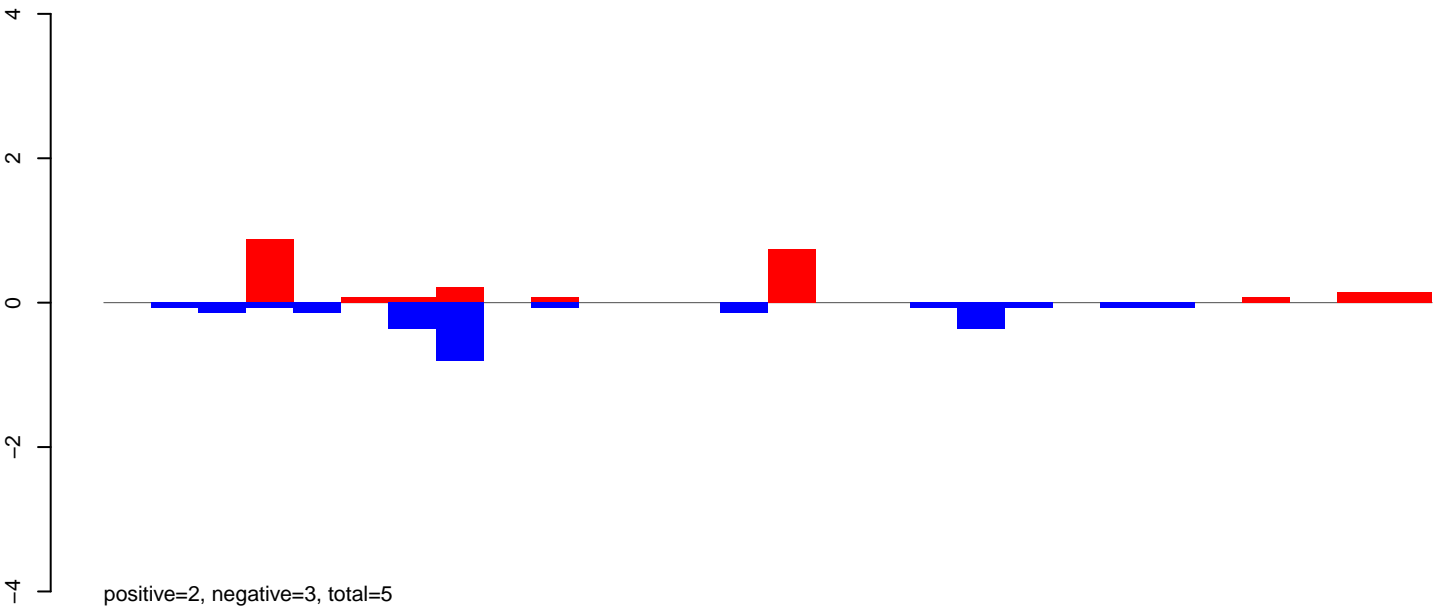


Window size=25, length=746, TE@rnd-4_family-615@DNA_Sola-1@1603:1-746

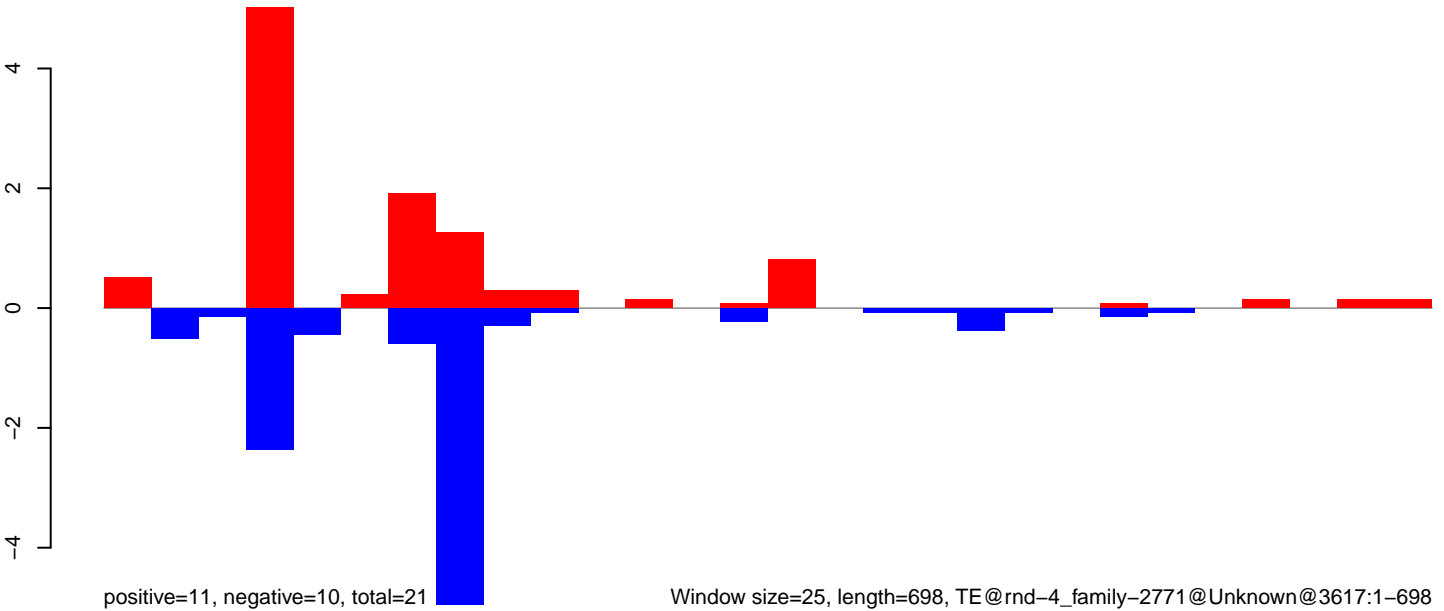
CuQuin_Testes_NL.18_23.rep



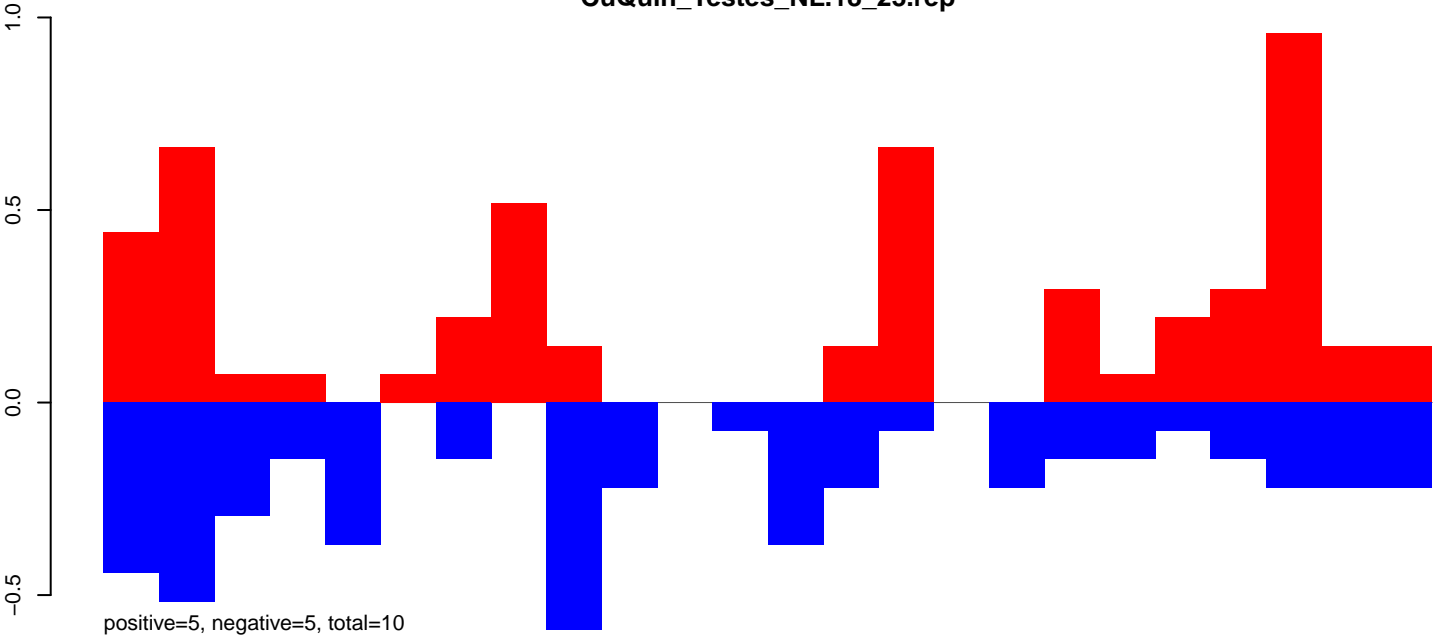
CuQuin_Testes_NL.24_35.rep



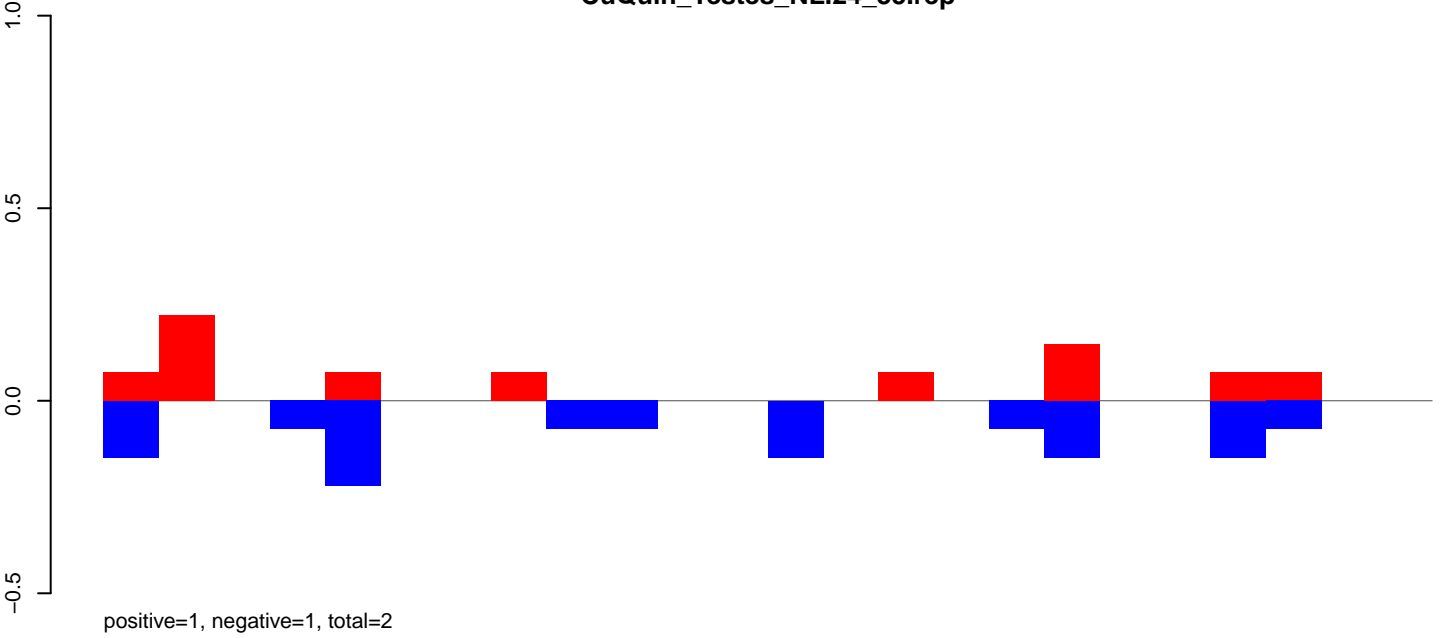
CuQuin_Testes_NL.rep



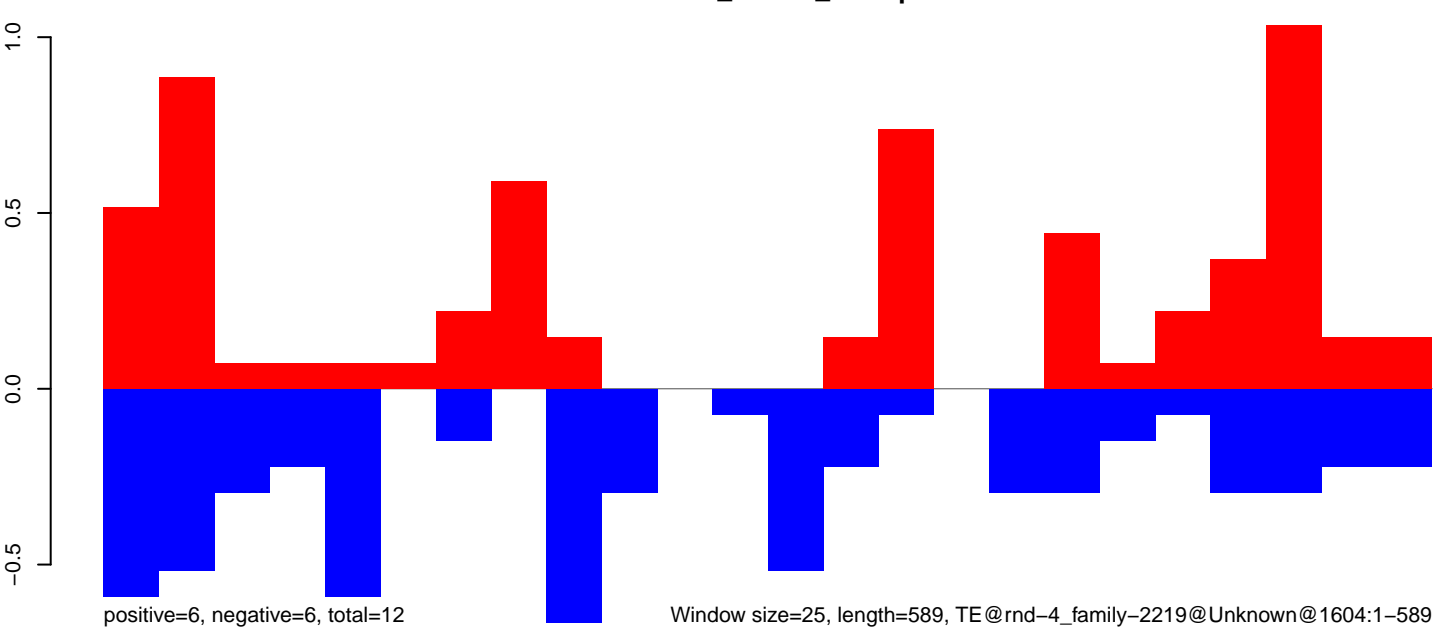
CuQuin_Testes_NL.18_23.rep



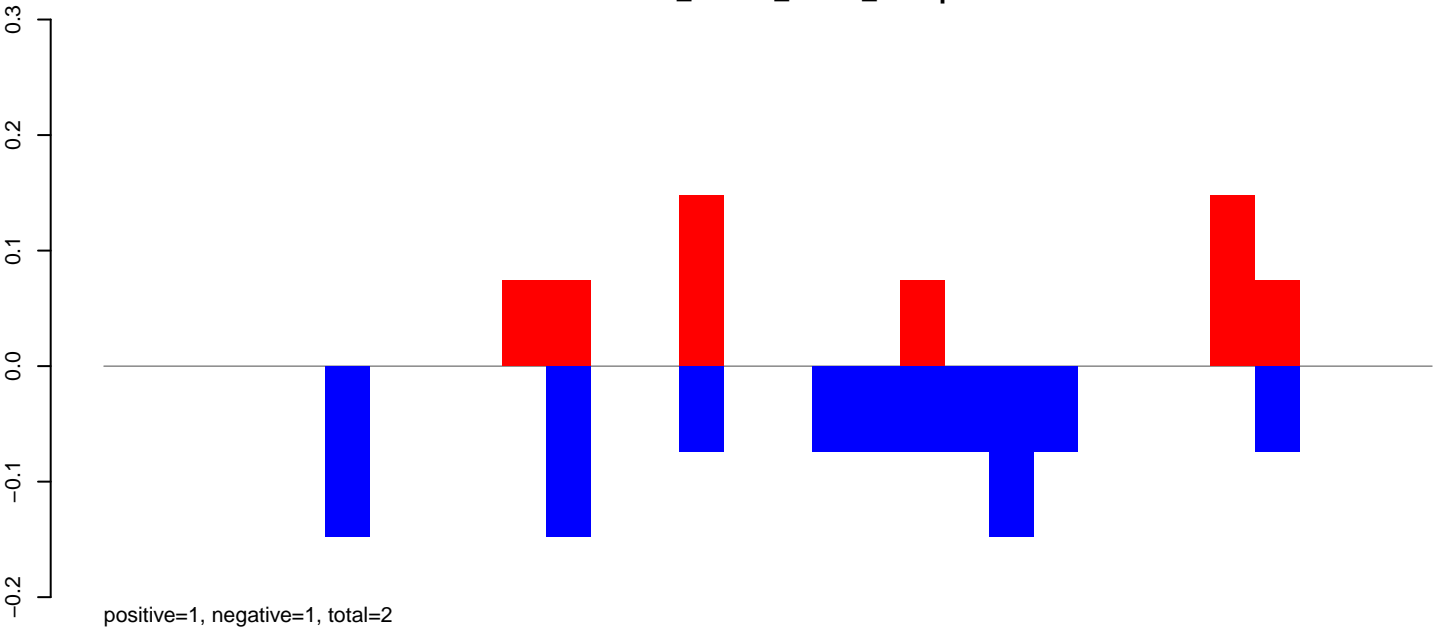
CuQuin_Testes_NL.24_35.rep



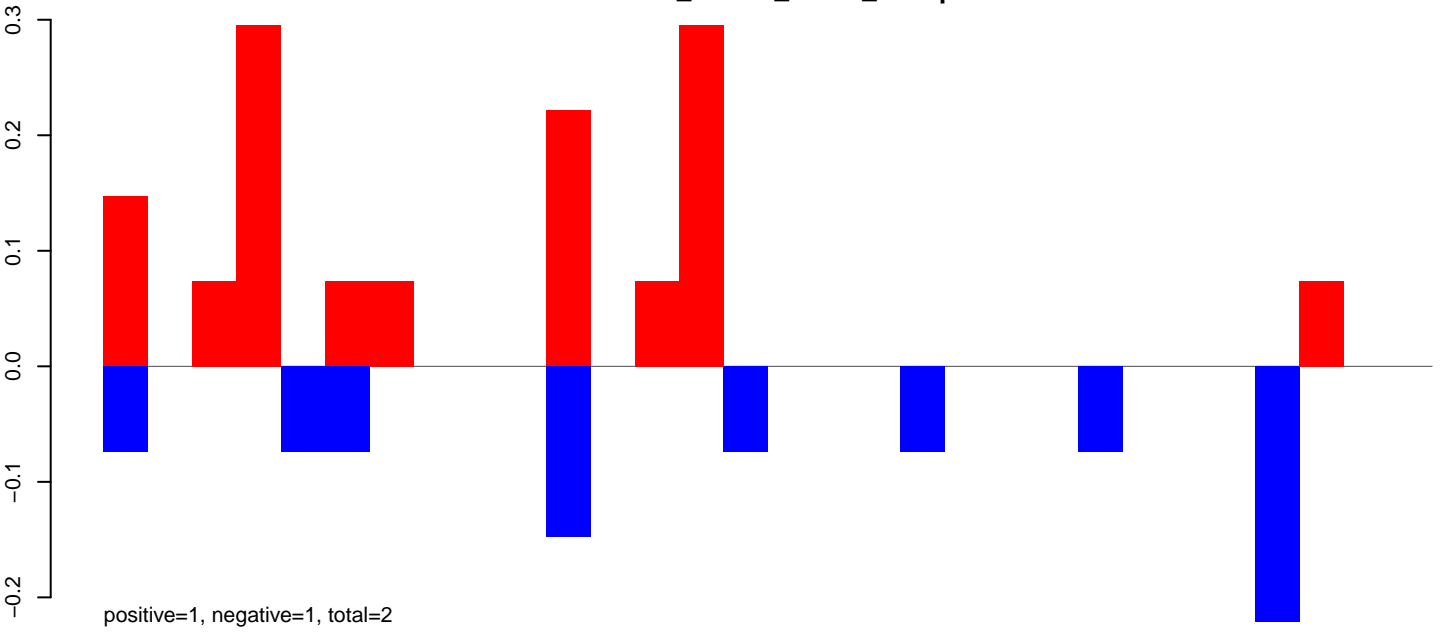
CuQuin_Testes_NL.rep



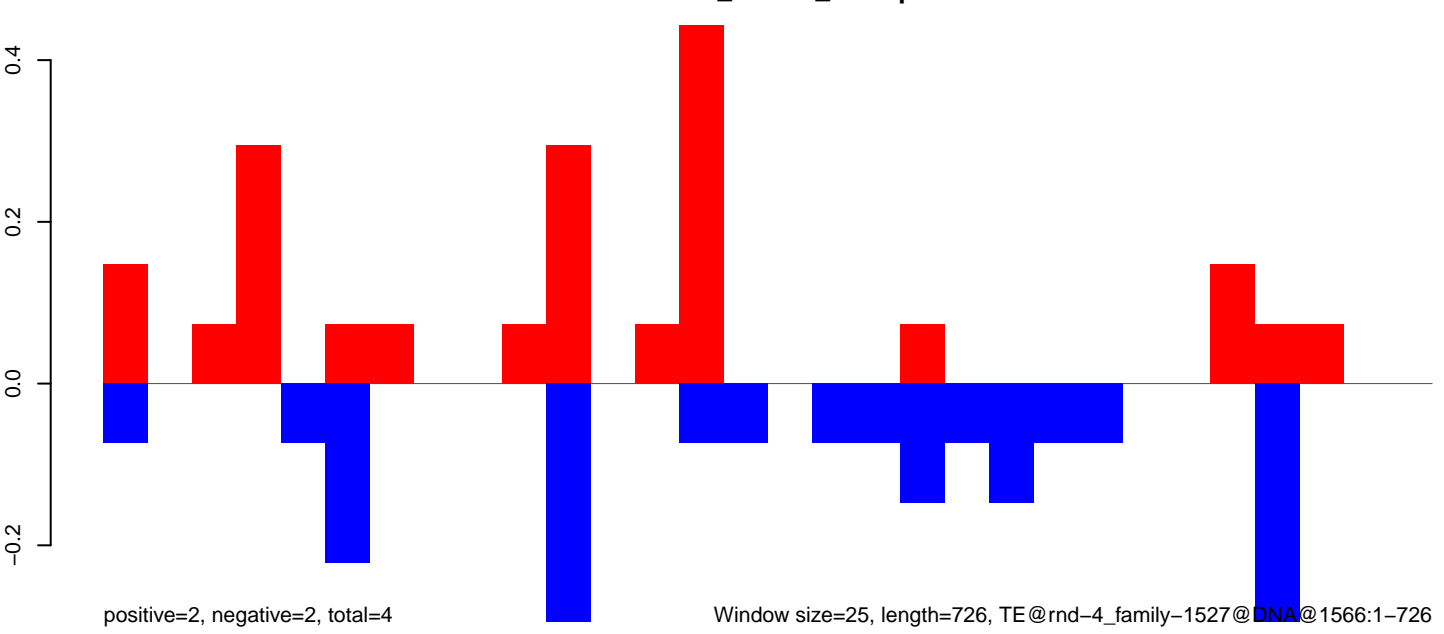
CuQuin_Testes_NL.18_23.rep



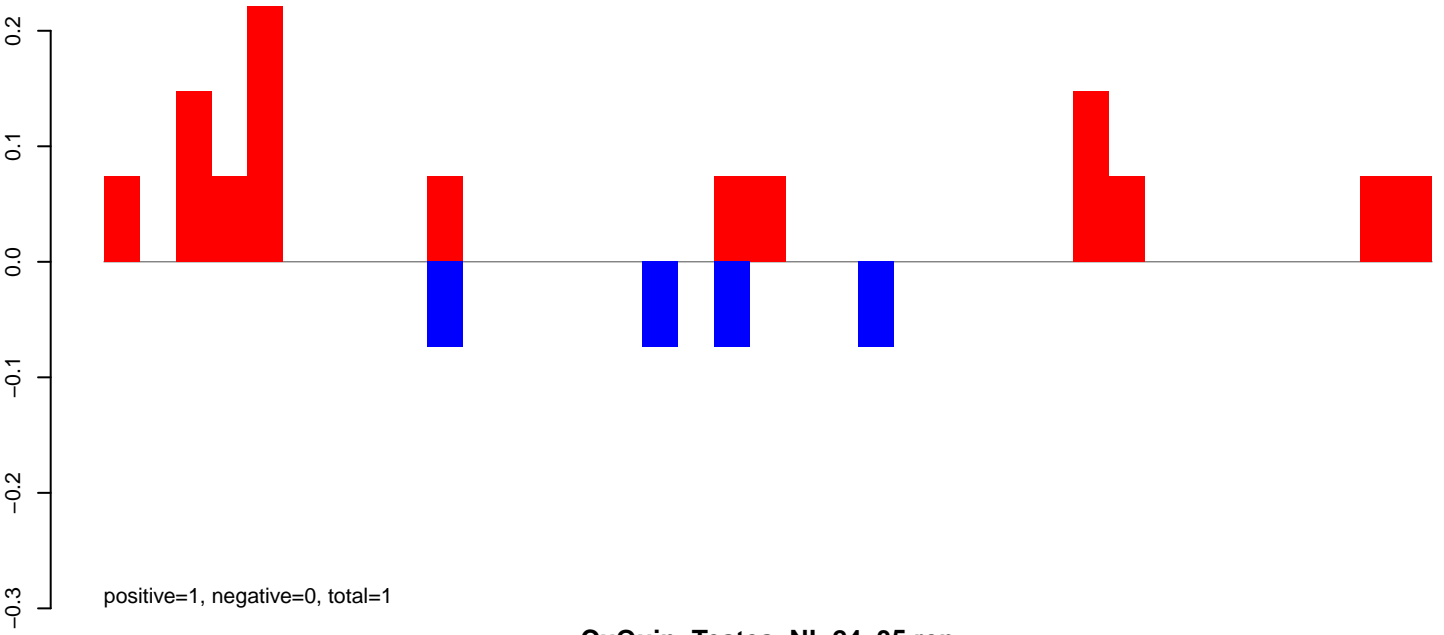
CuQuin_Testes_NL.24_35.rep



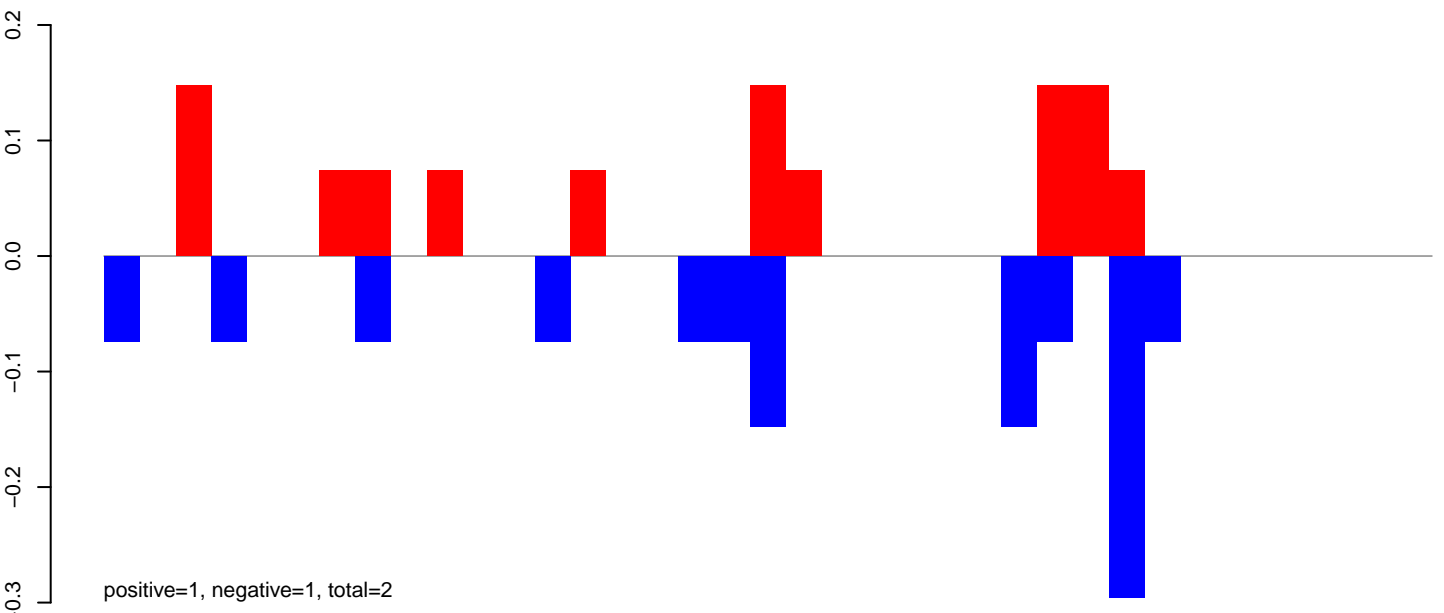
CuQuin_Testes_NL.rep



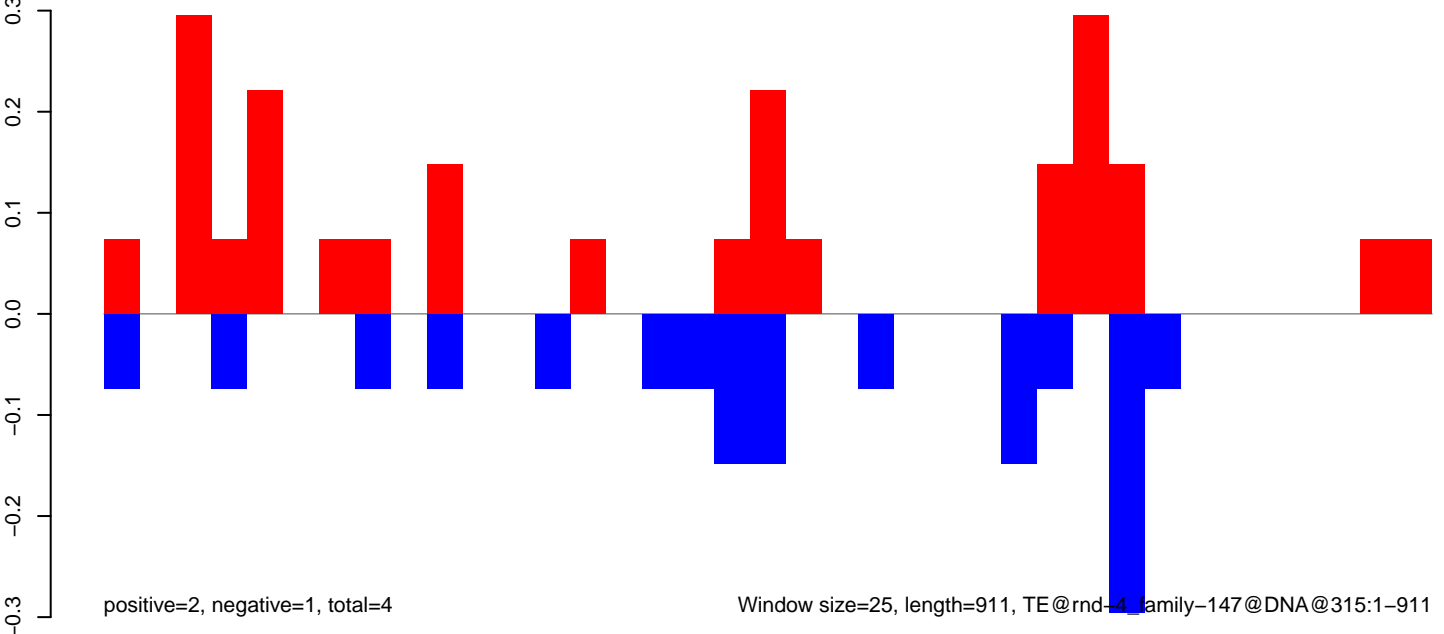
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

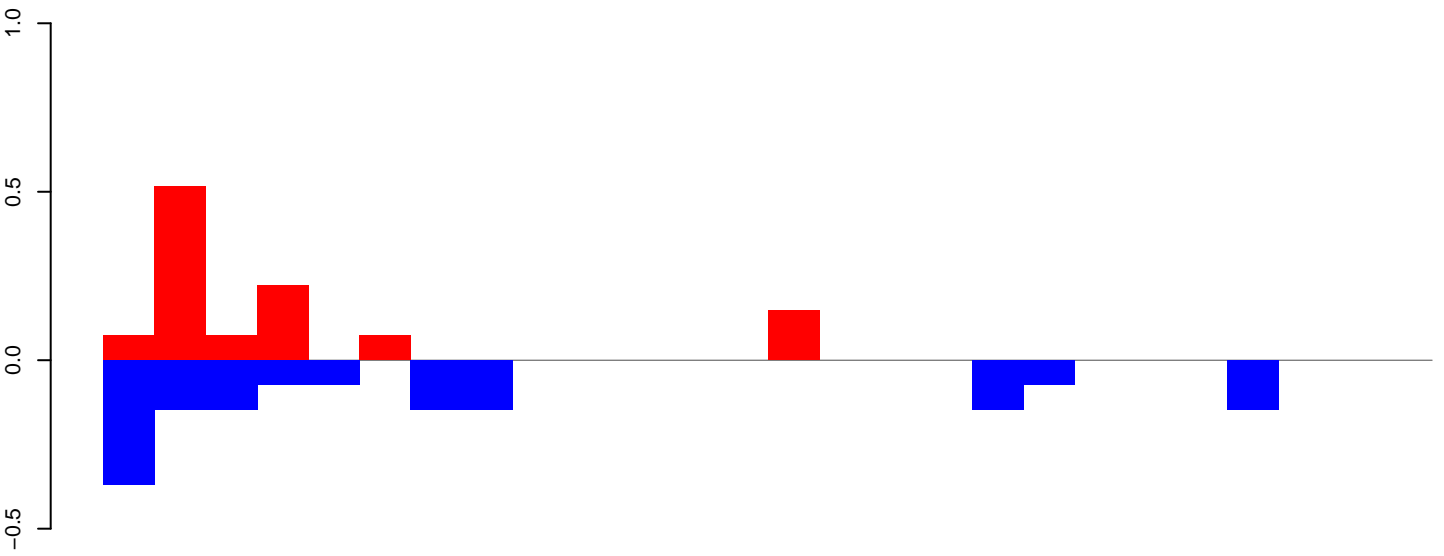


CuQuin_Testes_NL.rep



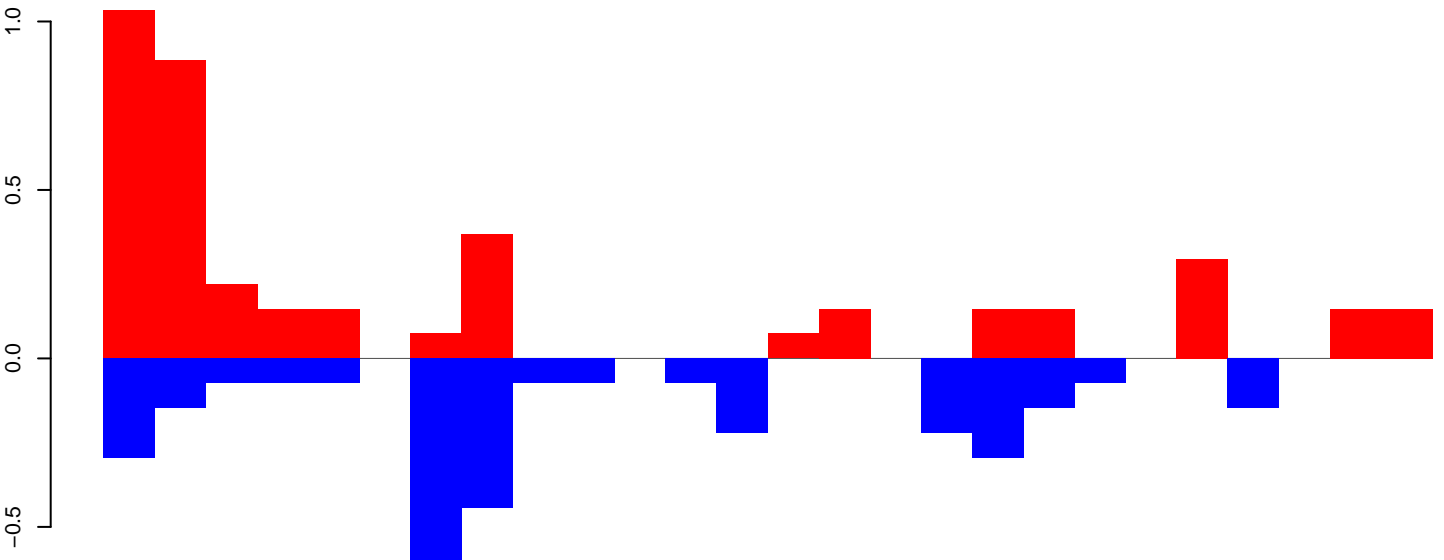
0 200 400 600 800

CuQuin_Testes_NL.18_23.rep



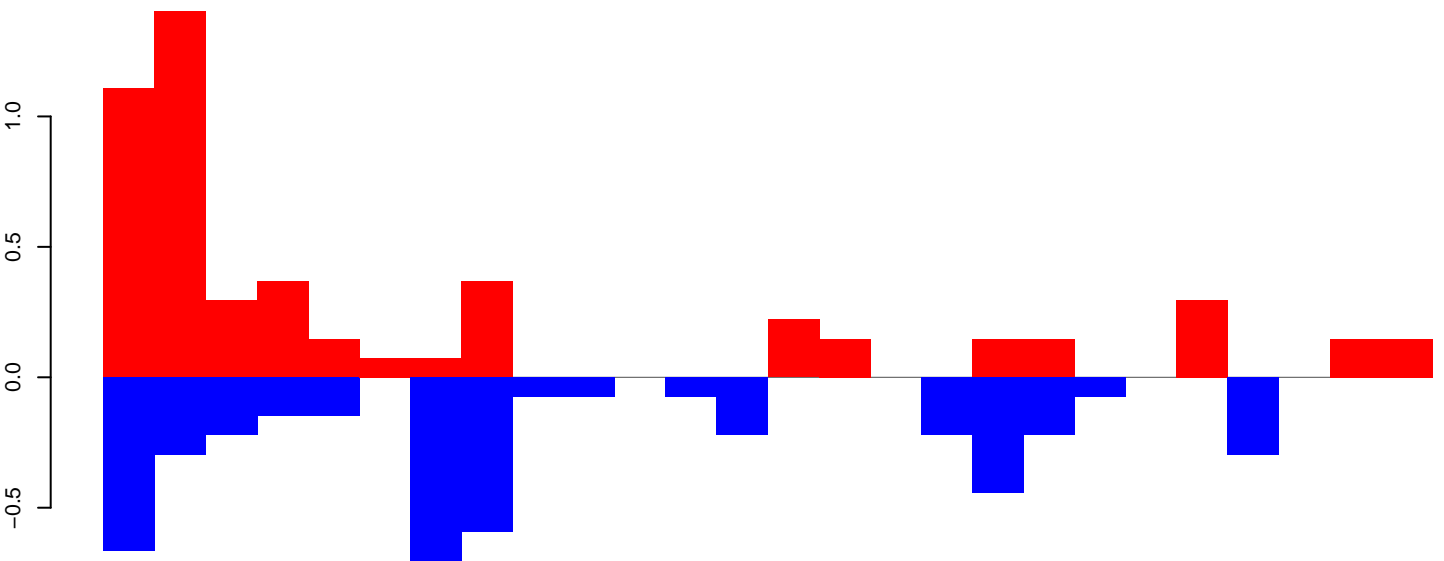
positive=1, negative=1, total=3

CuQuin_Testes_NL.24_35.rep



positive=4, negative=3, total=7

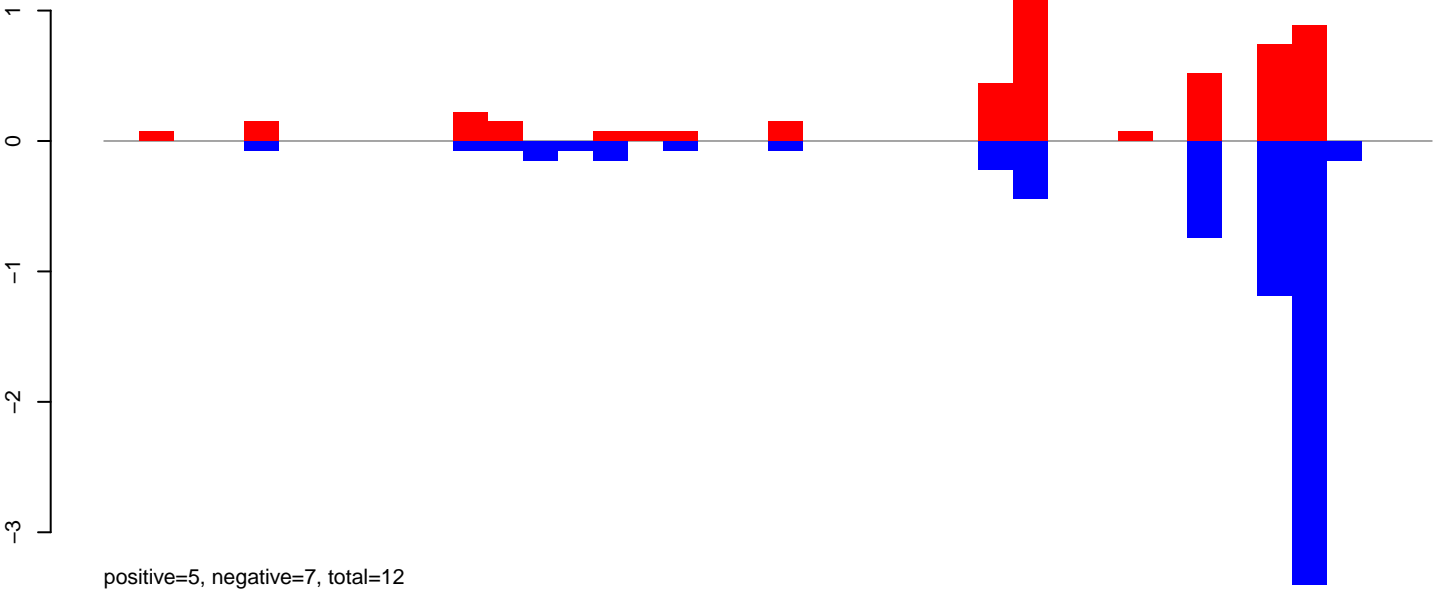
CuQuin_Testes_NL.rep



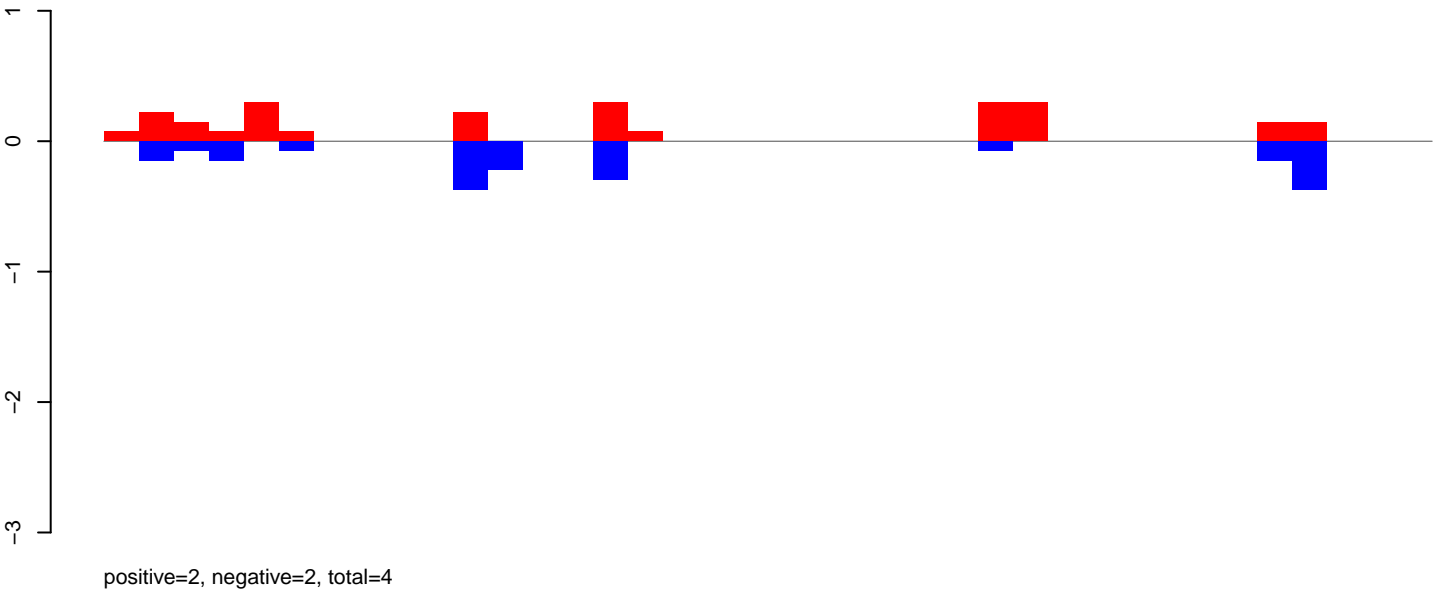
positive=5, negative=5, total=10

Window size=25, length=647, TE@rnd-4_family-1090@Unknown@3499:1-647

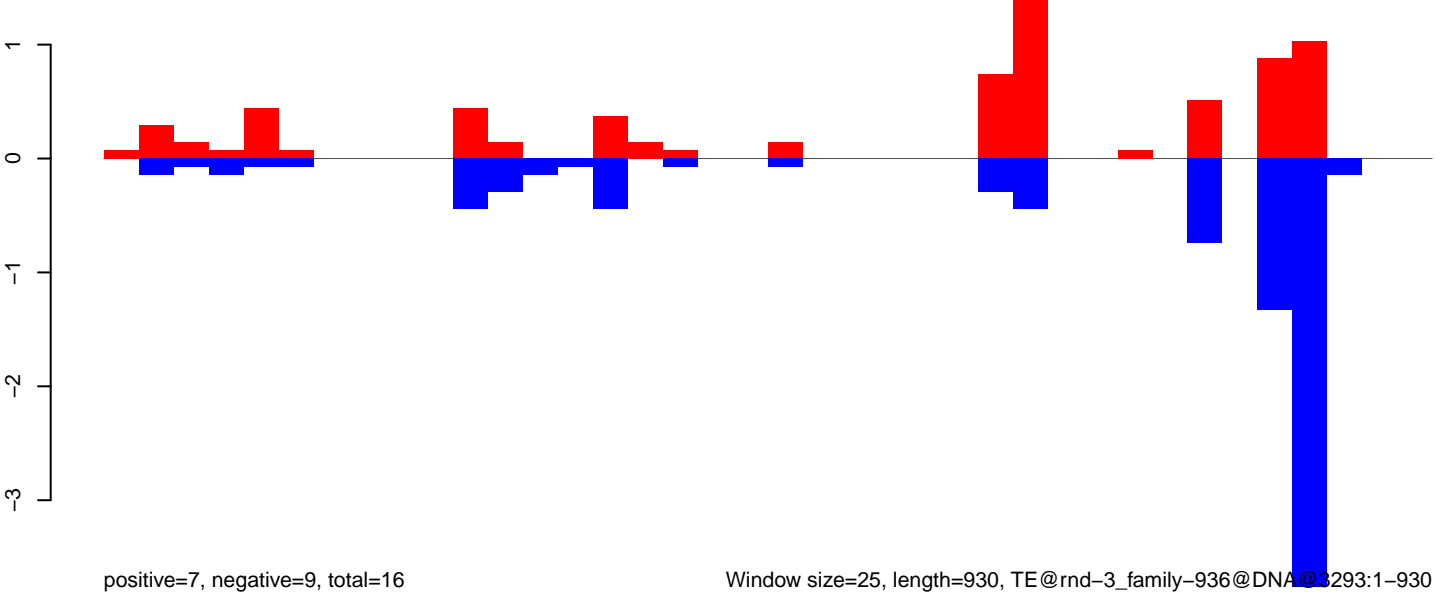
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



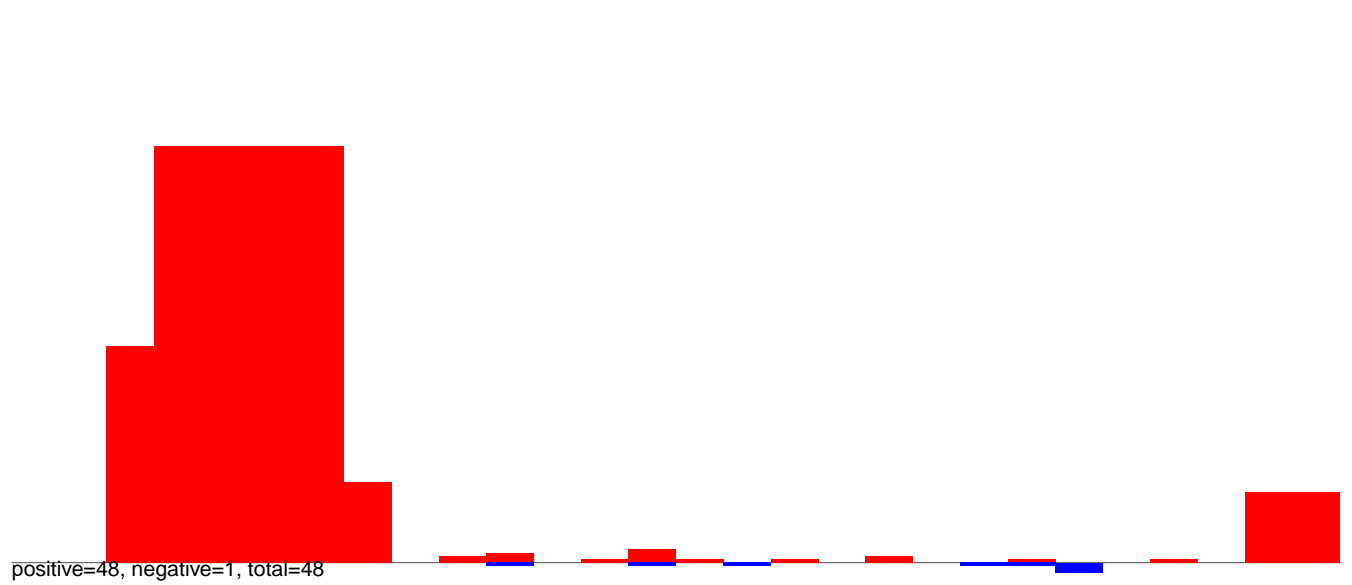
CuQuin_Testes_NL.rep



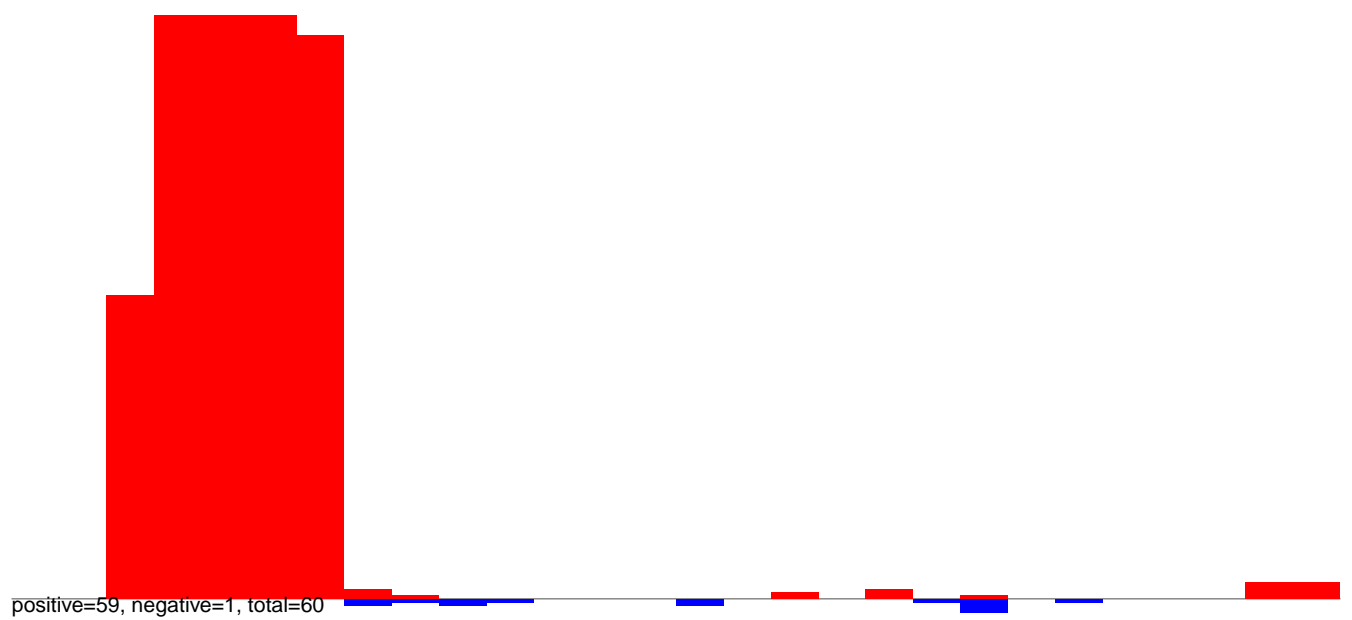
Window size=25, length=930, TE@rnd-3_family-936@DNA@3293:1-930

0 200 400 600 800 1000

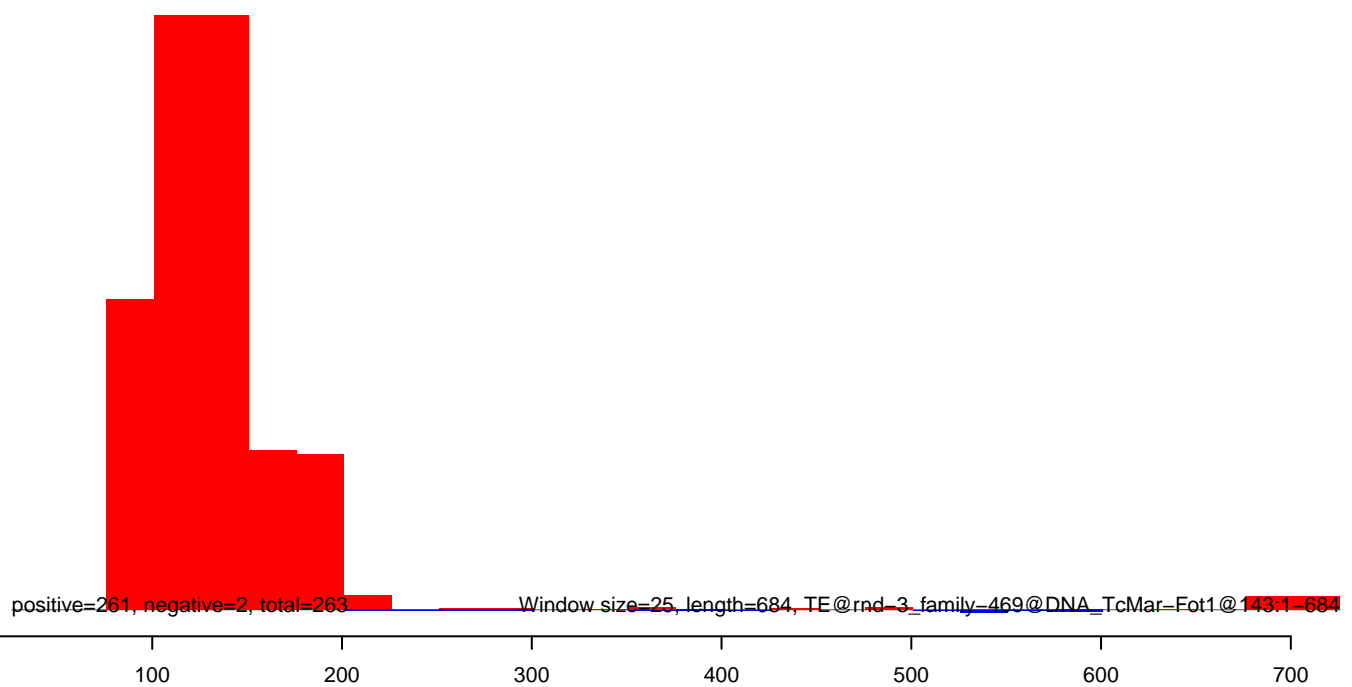
CuQuin_Testes_NL.18_23.rep



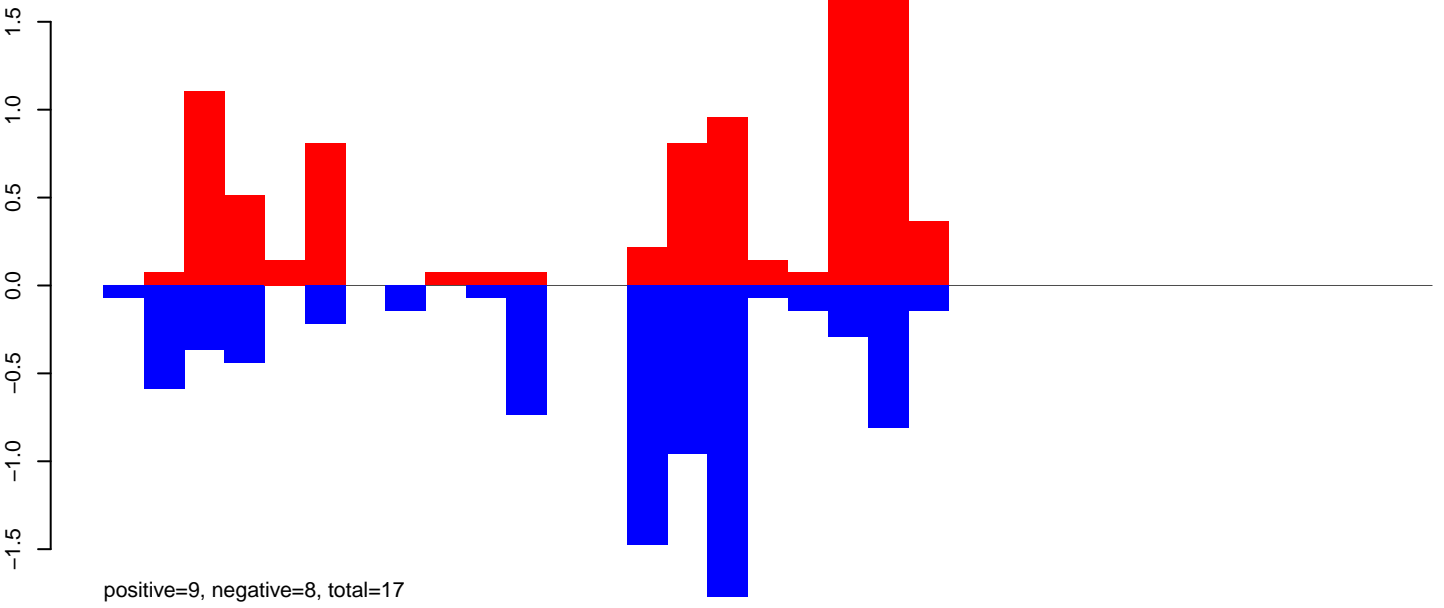
CuQuin_Testes_NL.24_35.rep



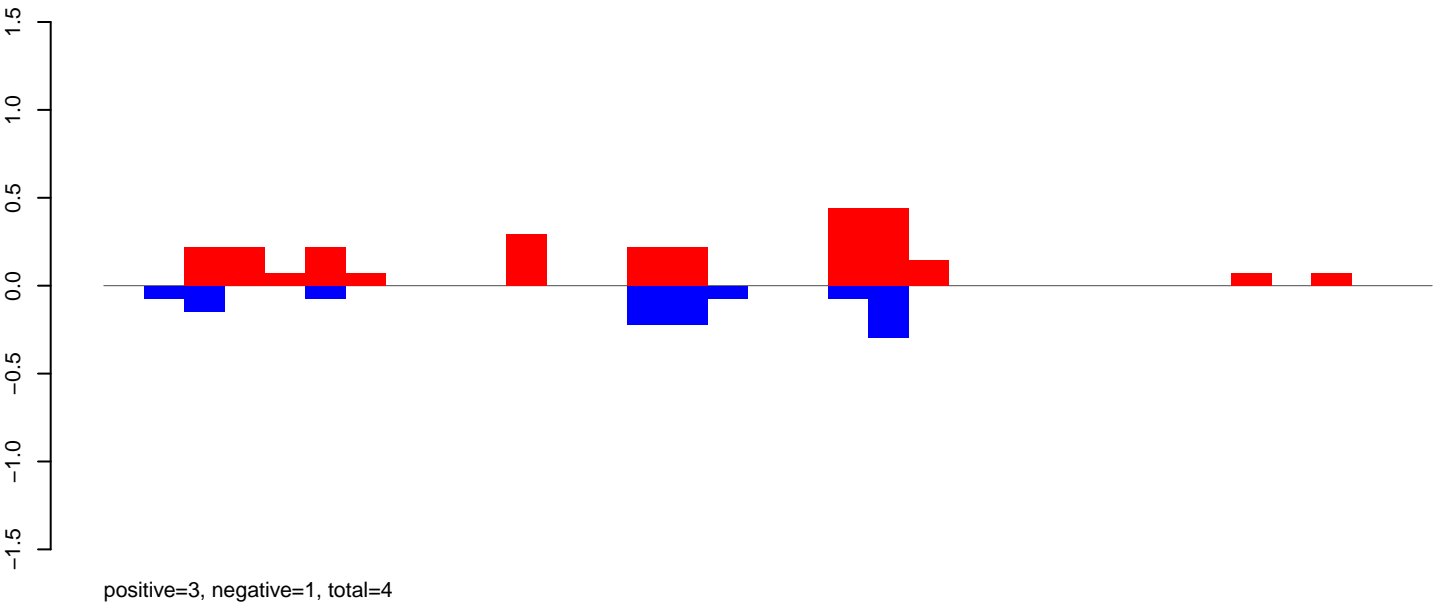
CuQuin_Testes_NL.rep



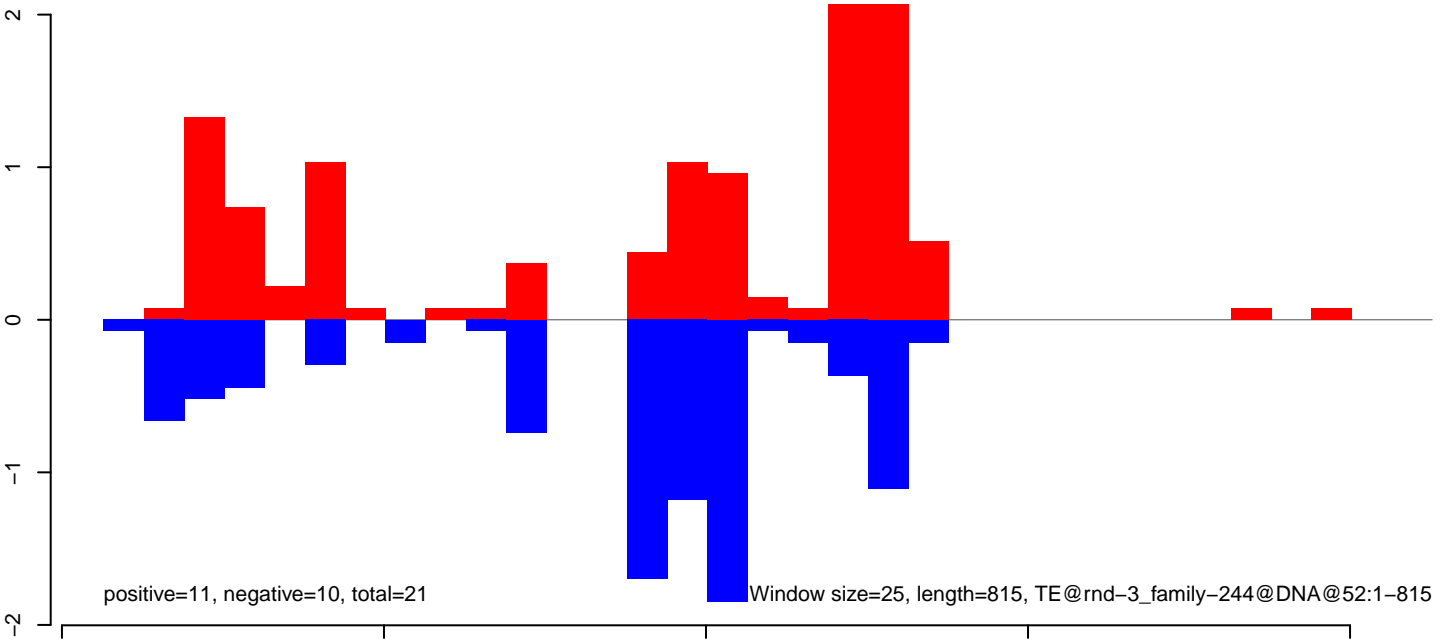
CuQuin_Testes_NL.18_23.rep



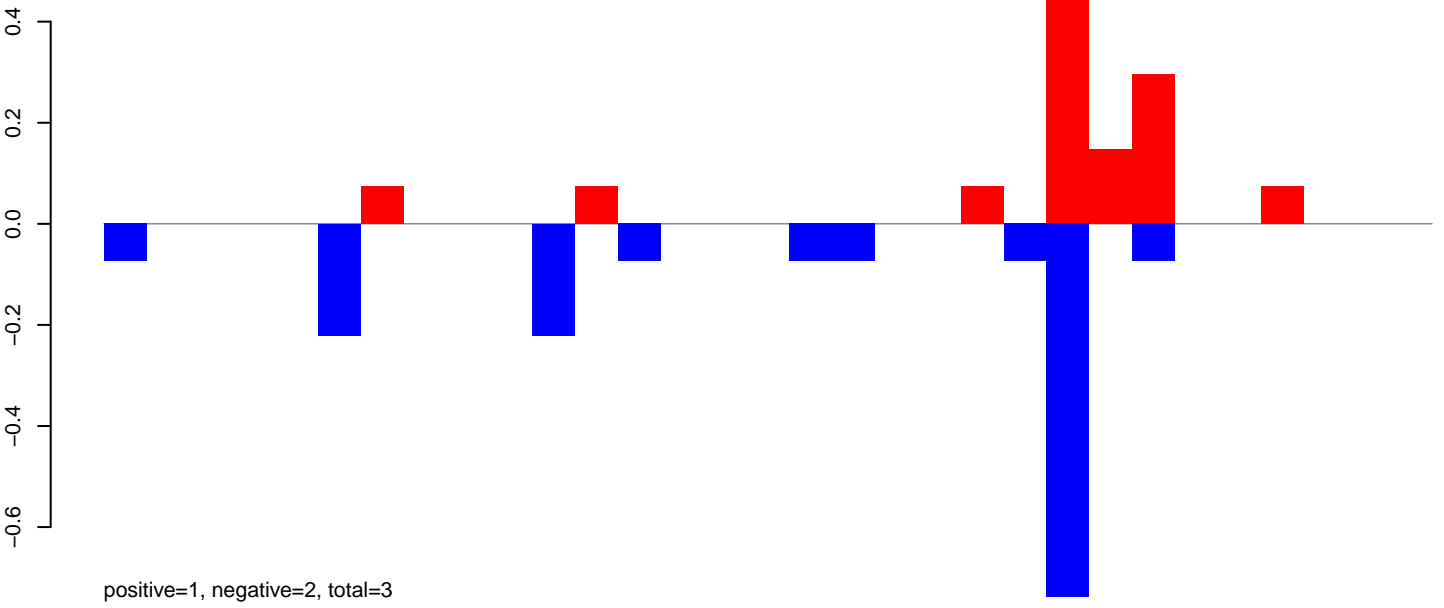
CuQuin_Testes_NL.24_35.rep



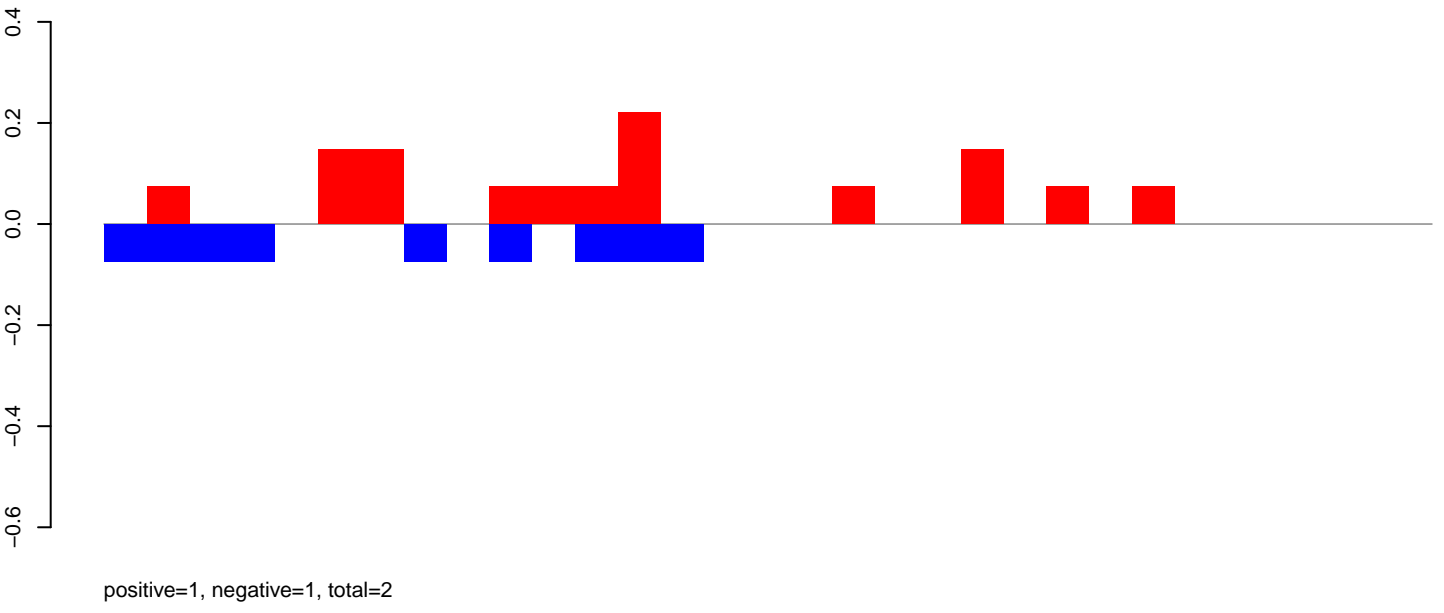
CuQuin_Testes_NL.rep



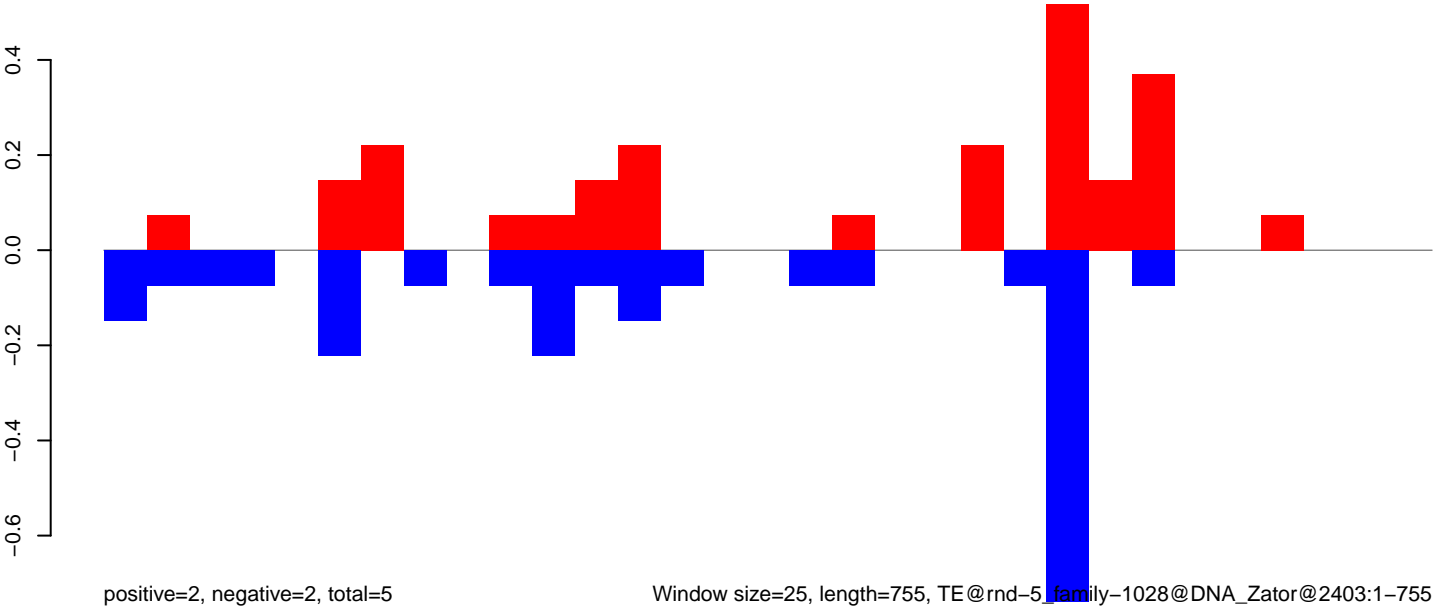
CuQuin_Testes_NL.18_23.rep



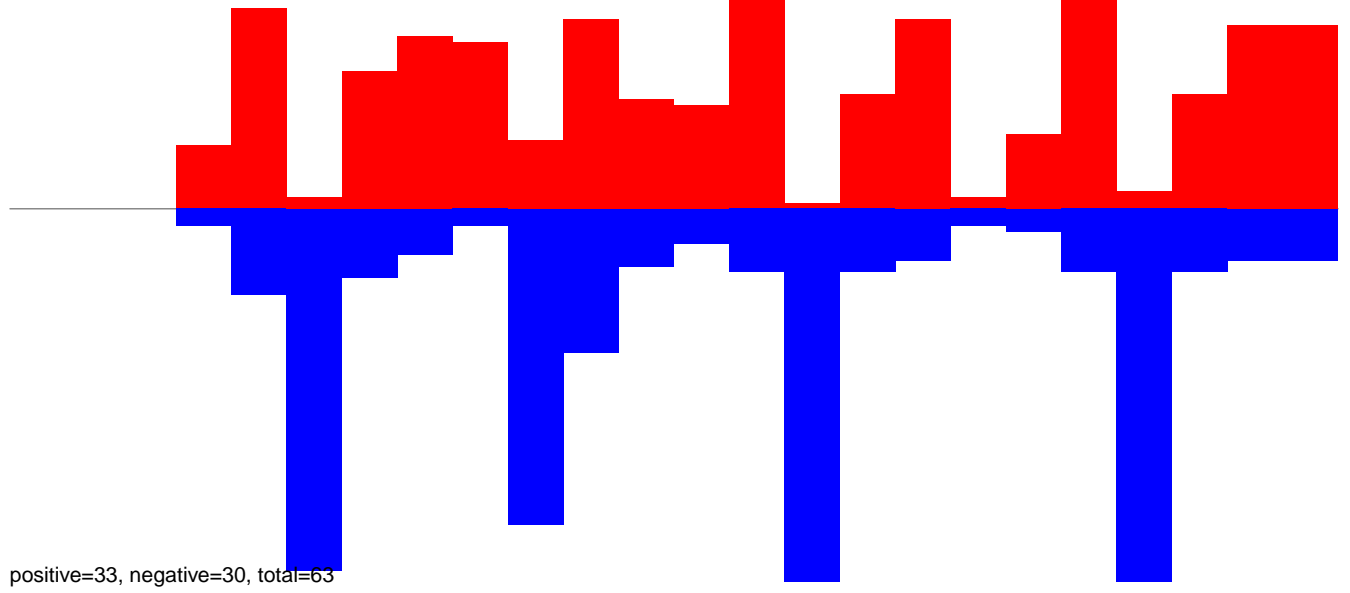
CuQuin_Testes_NL.24_35.rep



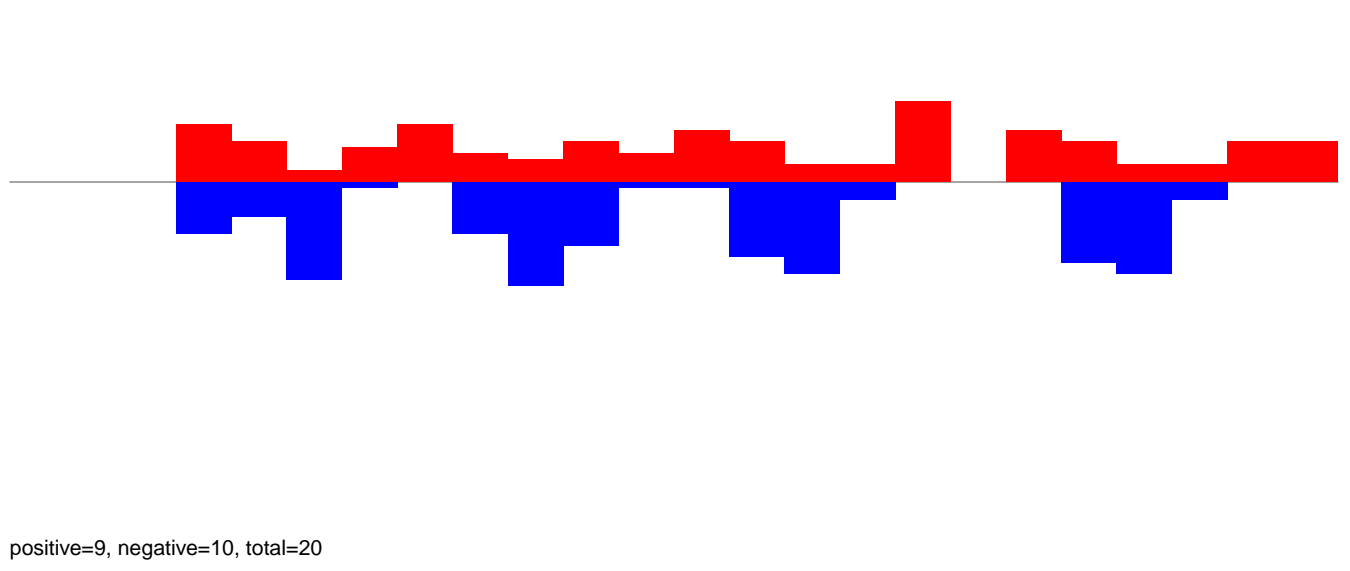
CuQuin_Testes_NL.rep



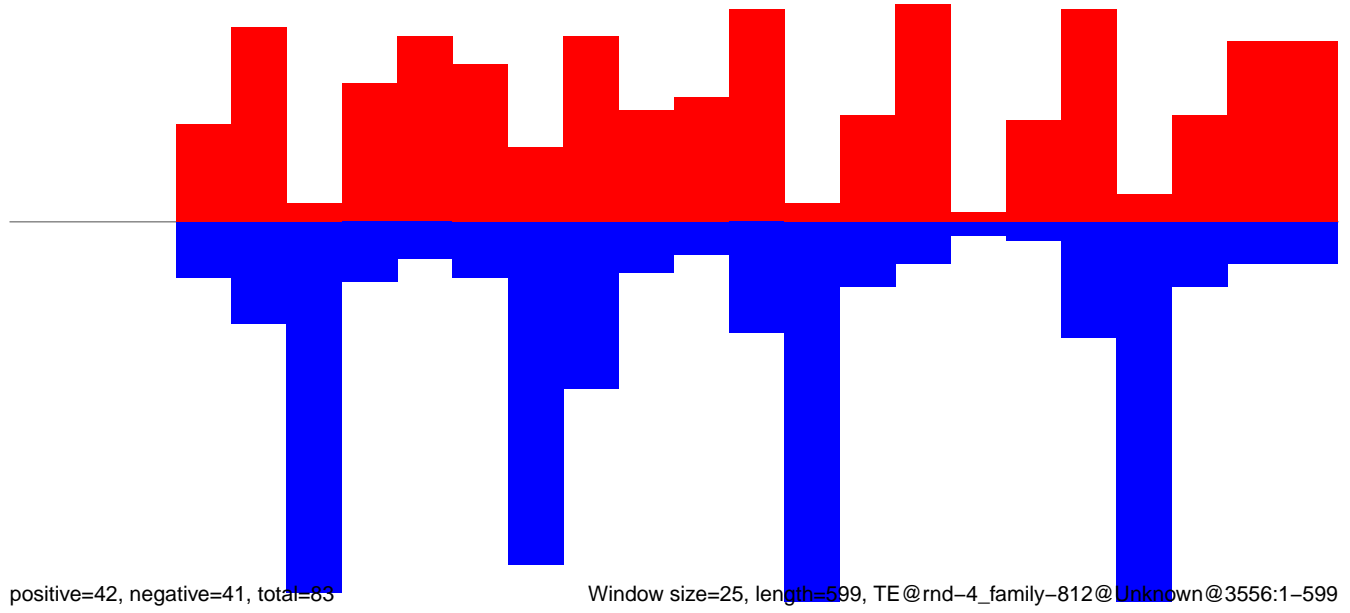
CuQuin_Testes_NL.18_23.rep



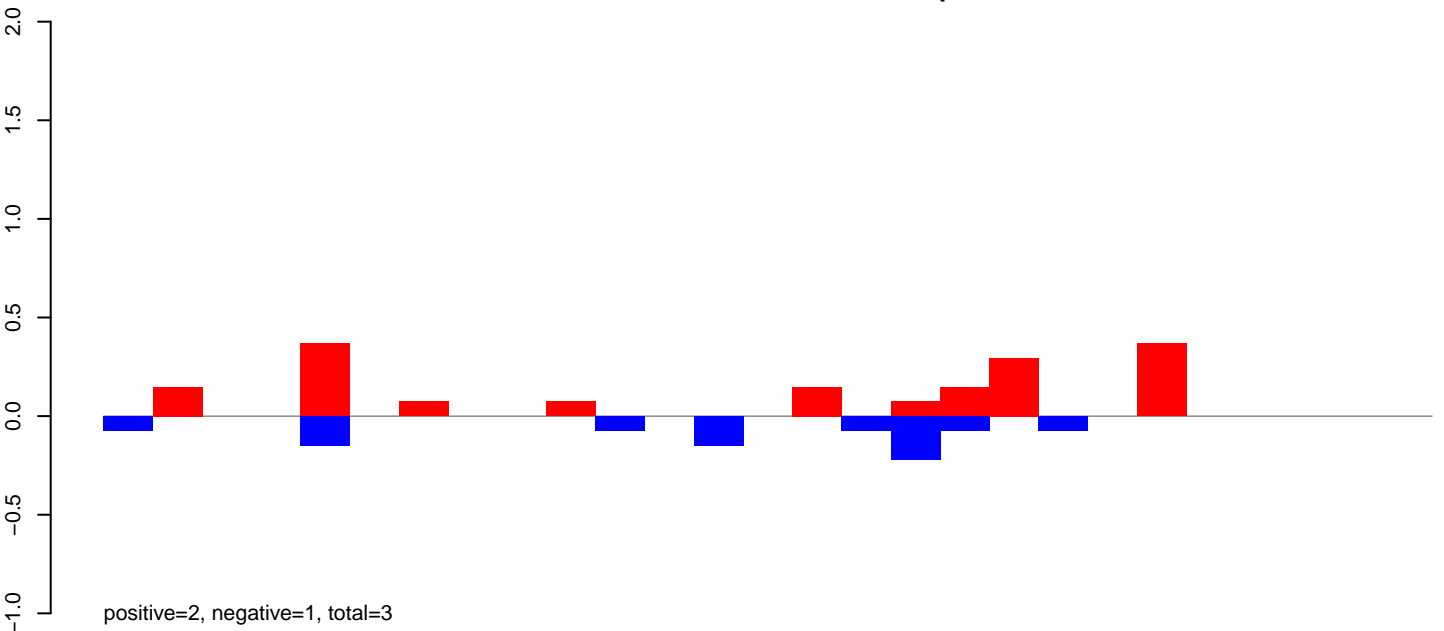
CuQuin_Testes_NL.24_35.rep



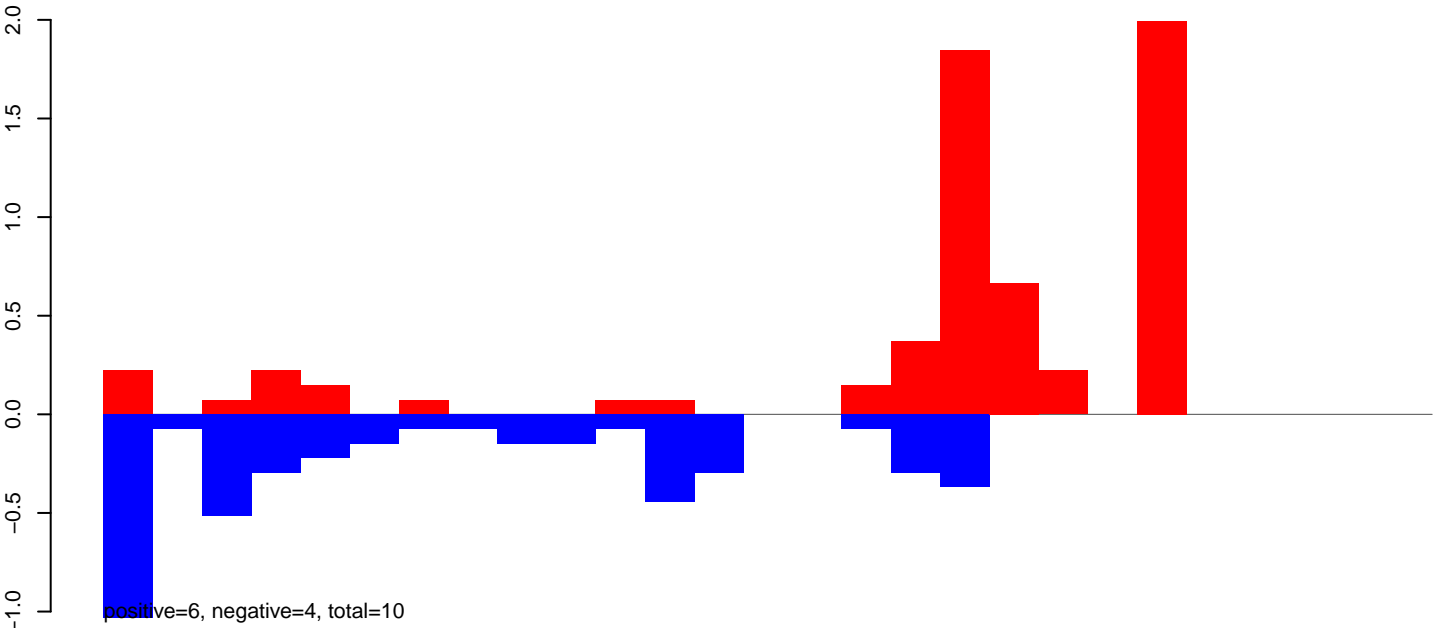
CuQuin_Testes_NL.rep



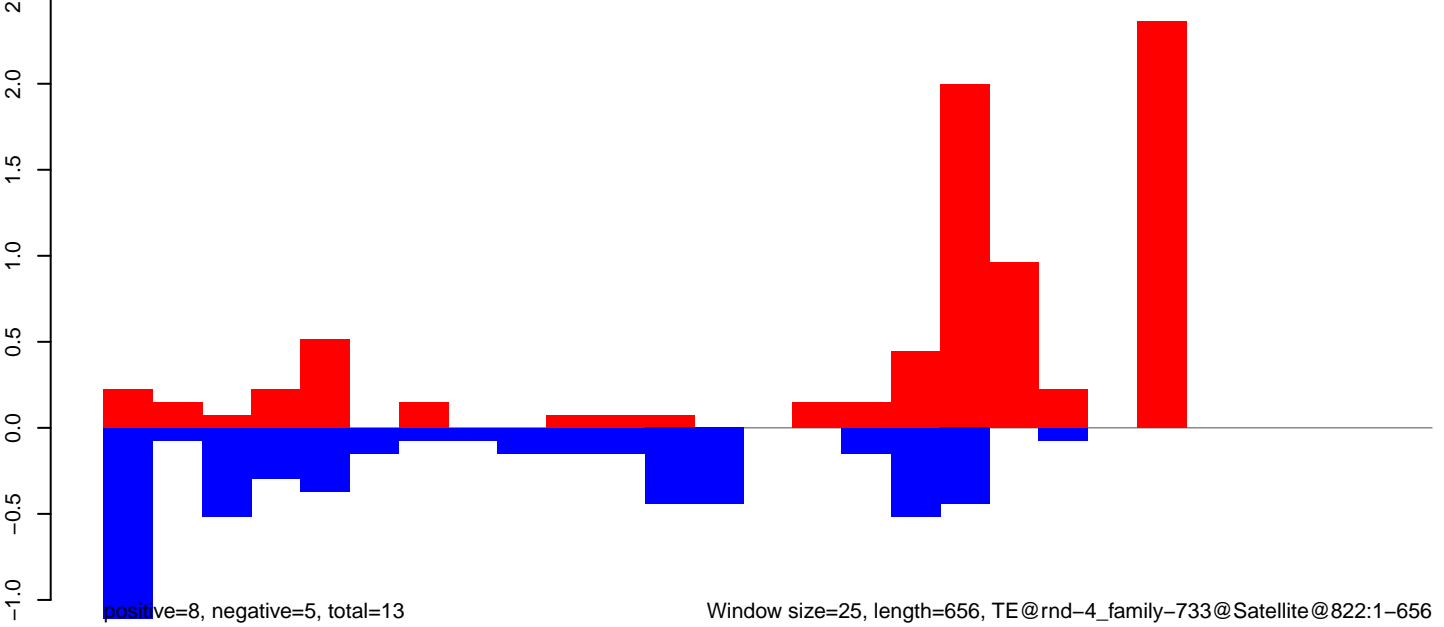
CuQuin_Testes_NL.18_23.rep



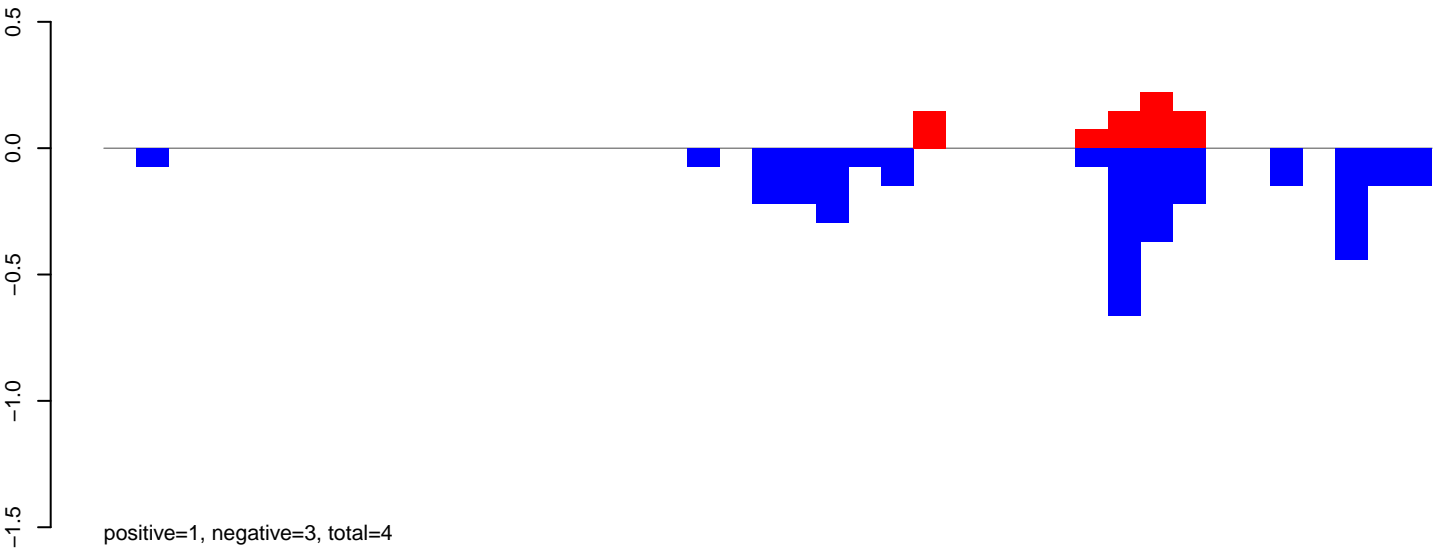
CuQuin_Testes_NL.24_35.rep



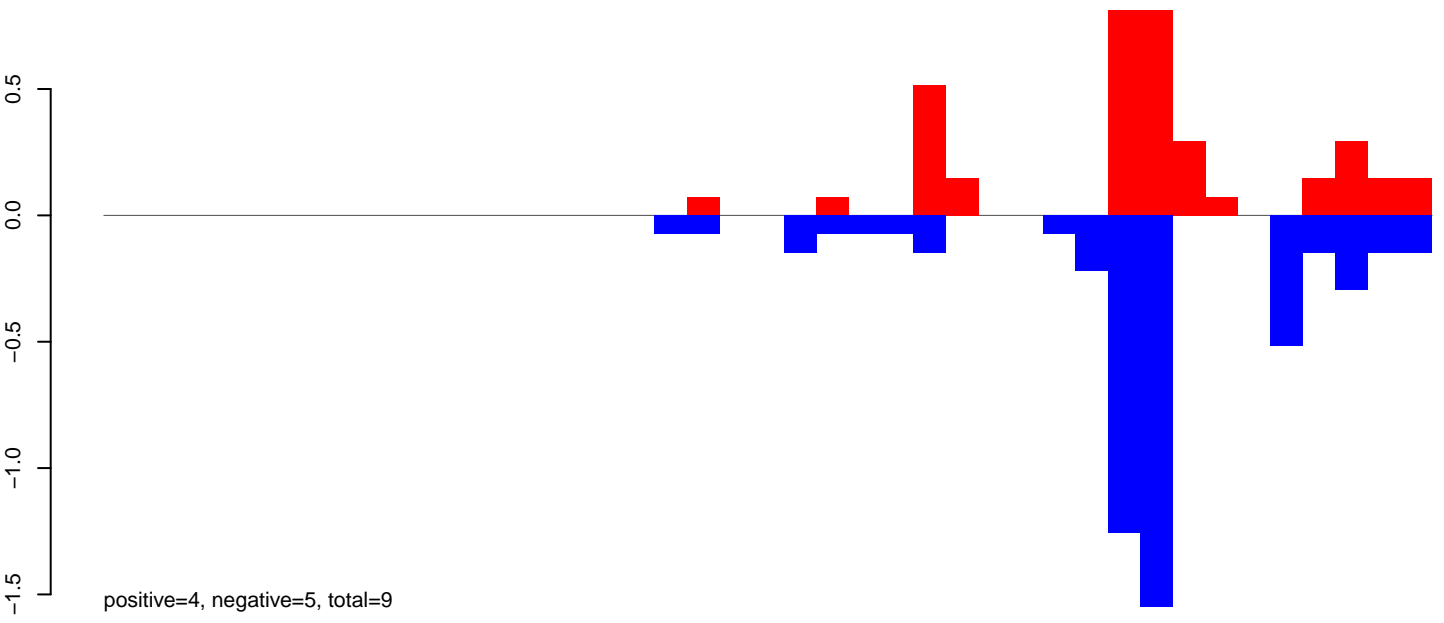
CuQuin_Testes_NL.rep



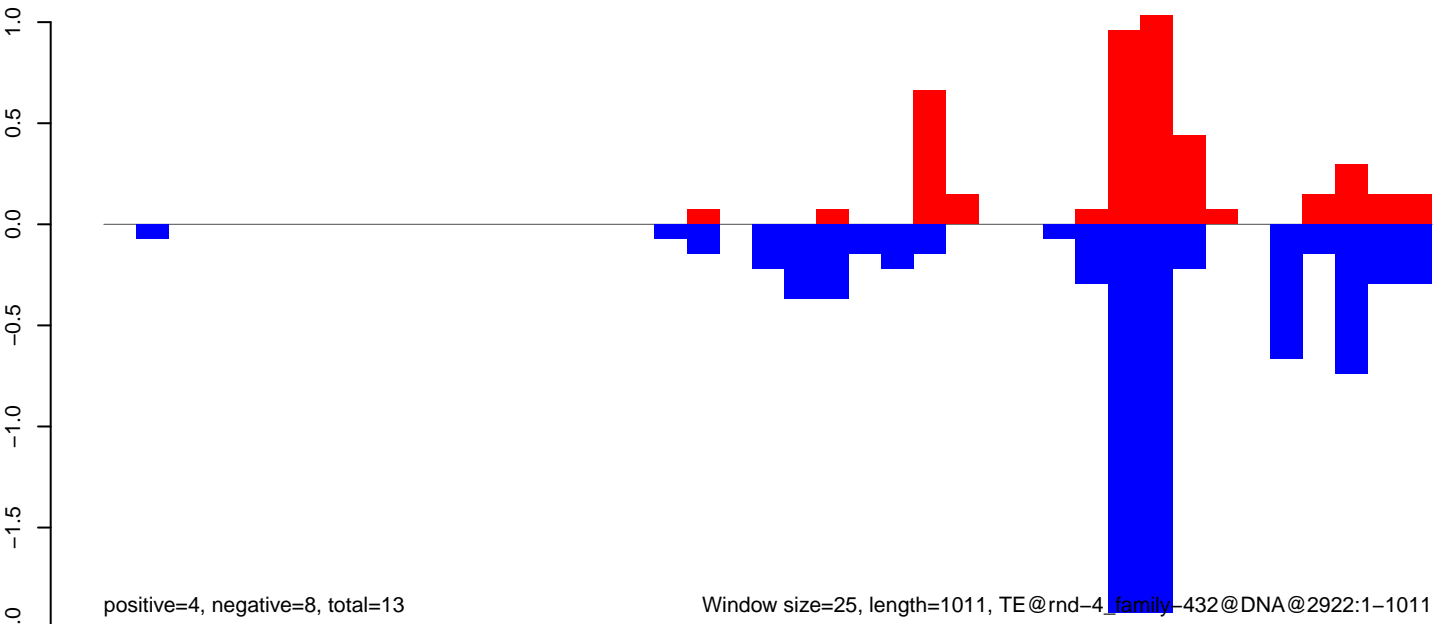
CuQuin_Testes_NL.18_23.rep



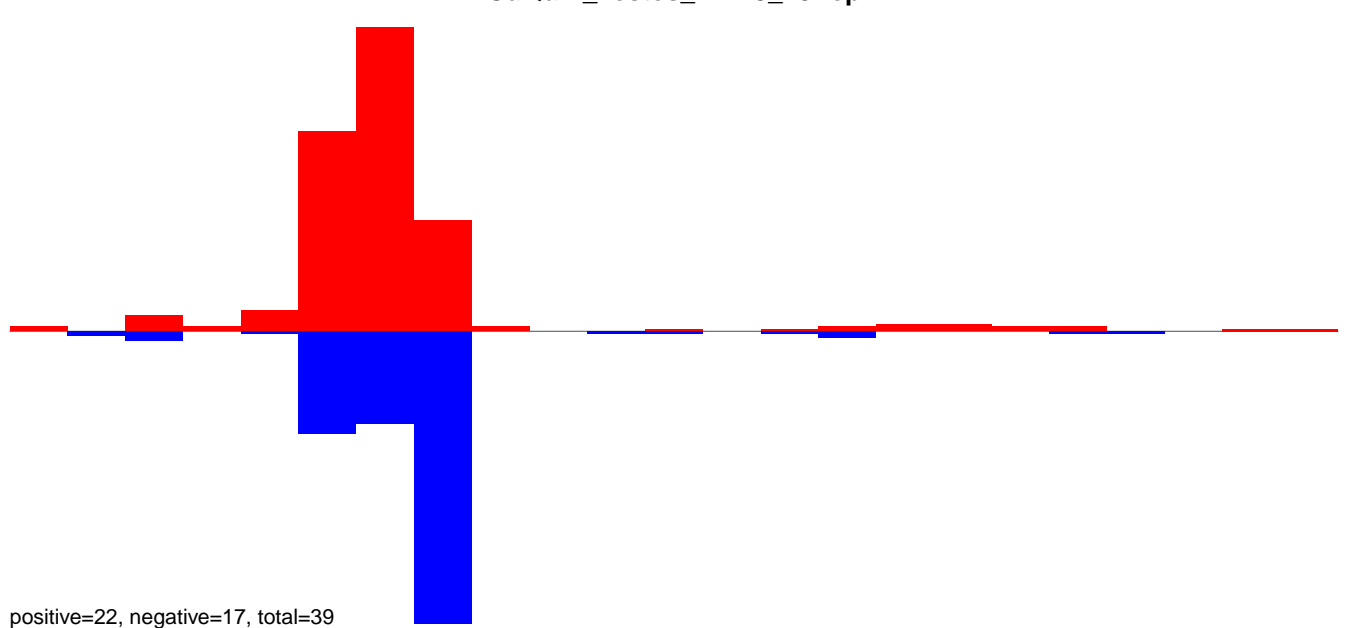
CuQuin_Testes_NL.24_35.rep



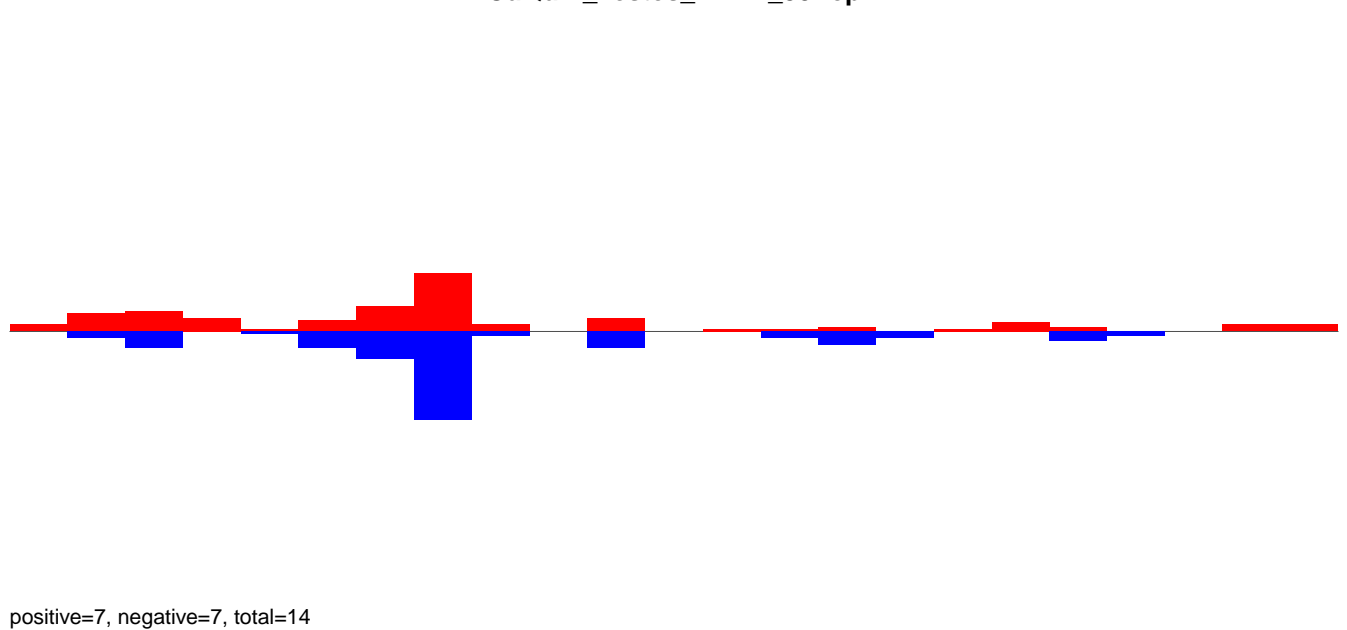
CuQuin_Testes_NL.rep



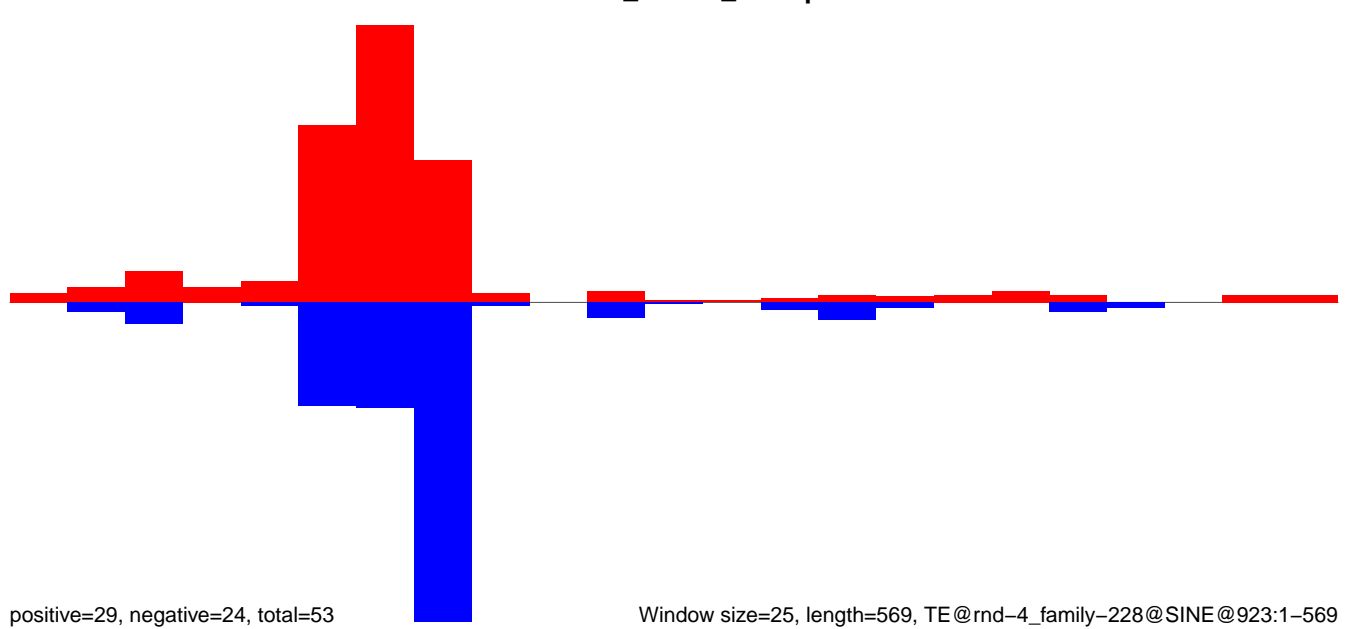
CuQuin_Testes_NL.18_23.rep



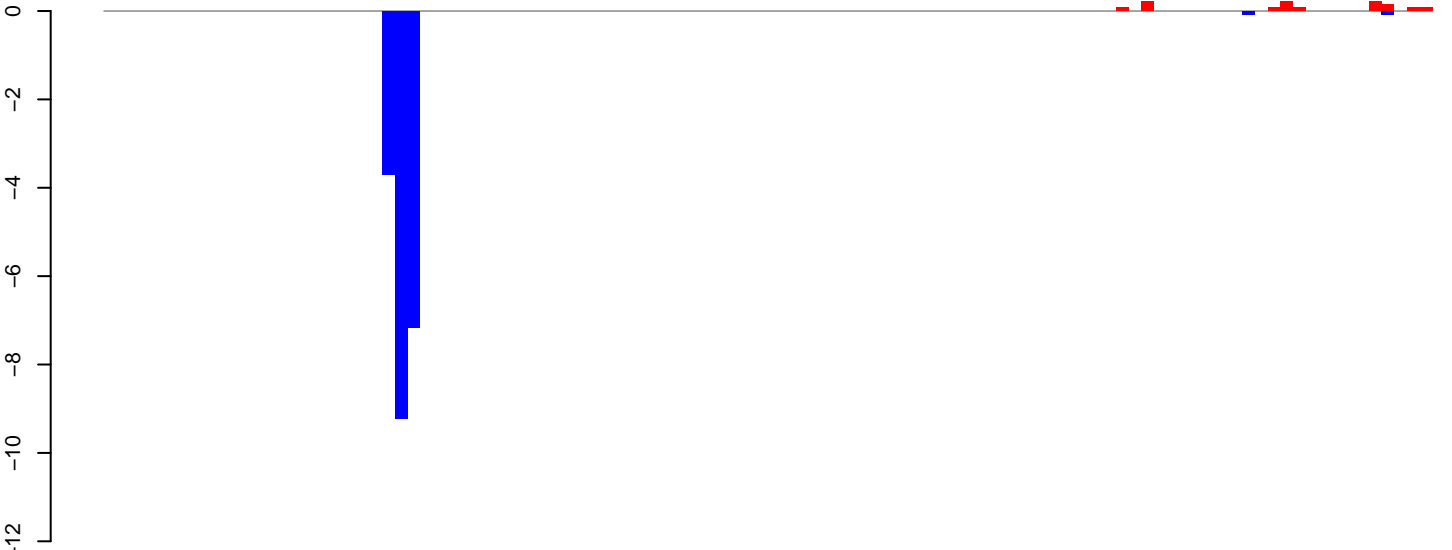
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

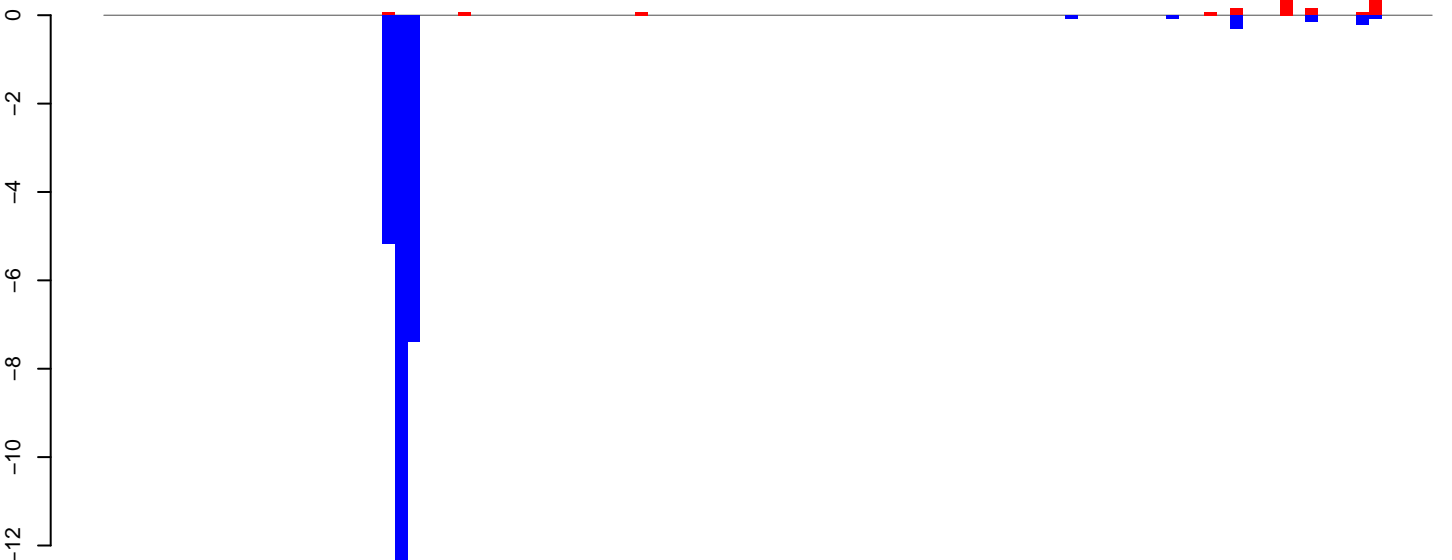


CuQuin_Testes_NL.18_23.rep



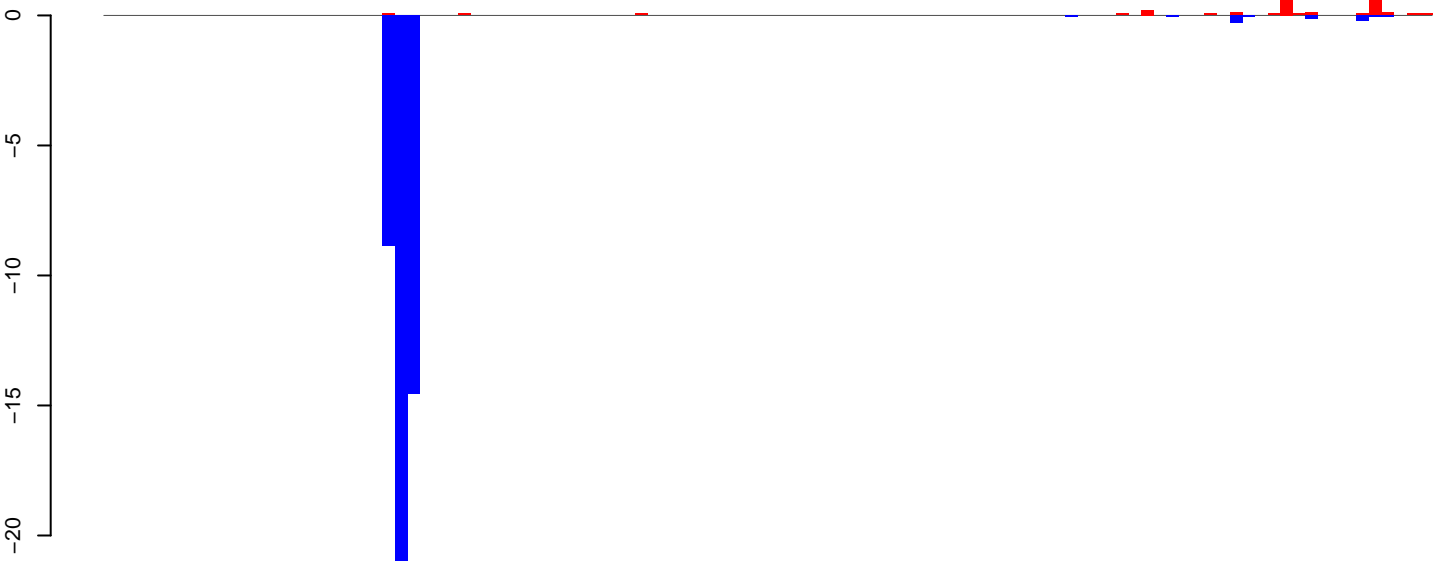
positive=1, negative=20, total=21

CuQuin_Testes_NL.24_35.rep



positive=2, negative=26, total=28

CuQuin_Testes_NL.rep

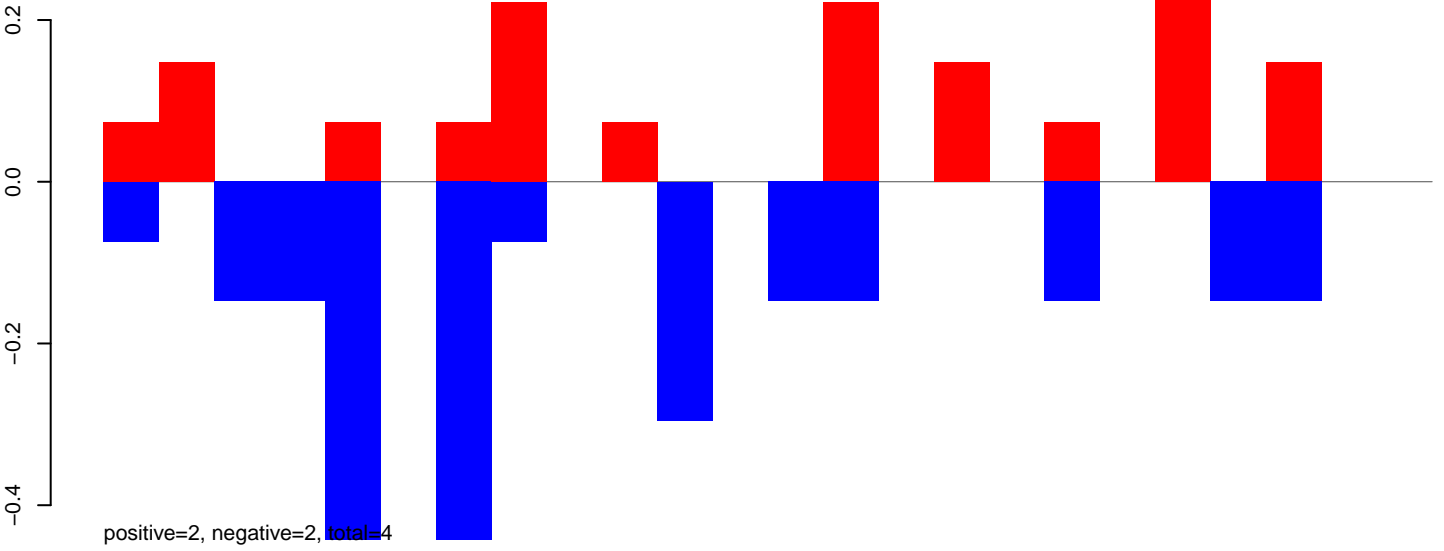


positive=3, negative=47, total=50

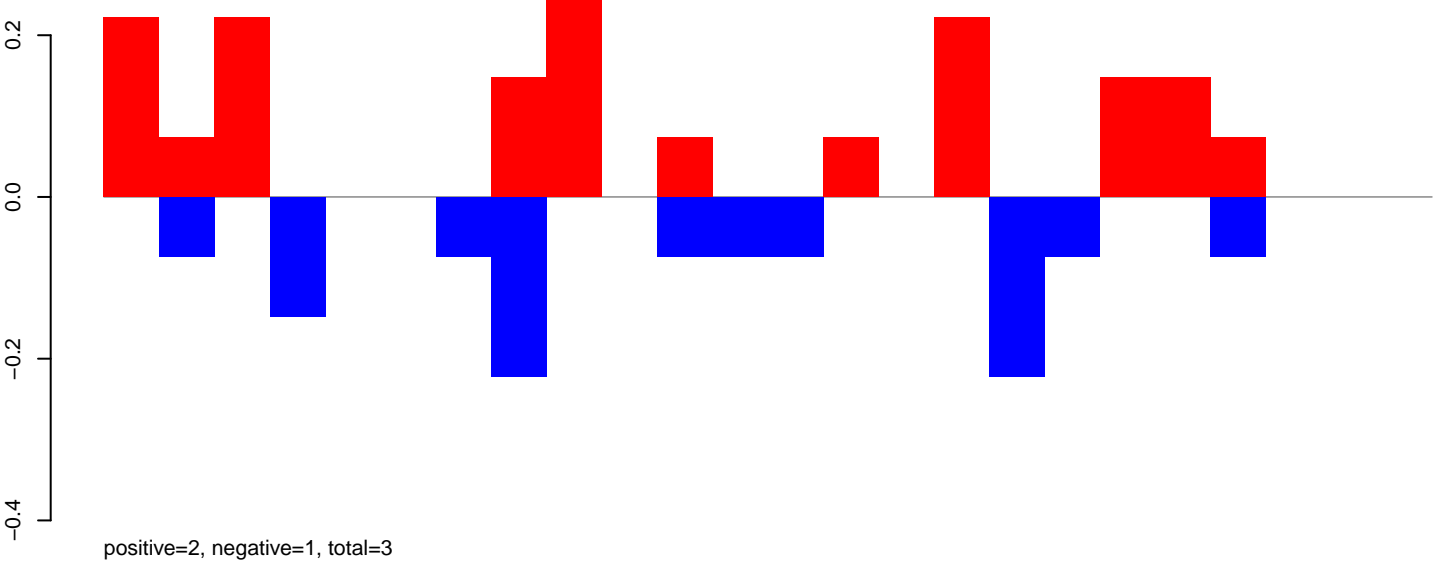
Window size=25, length=2602, TE@rnd-4_family-213@DNA_PiggyBac@2165:1-2602

0 500 1000 1500 2000 2500

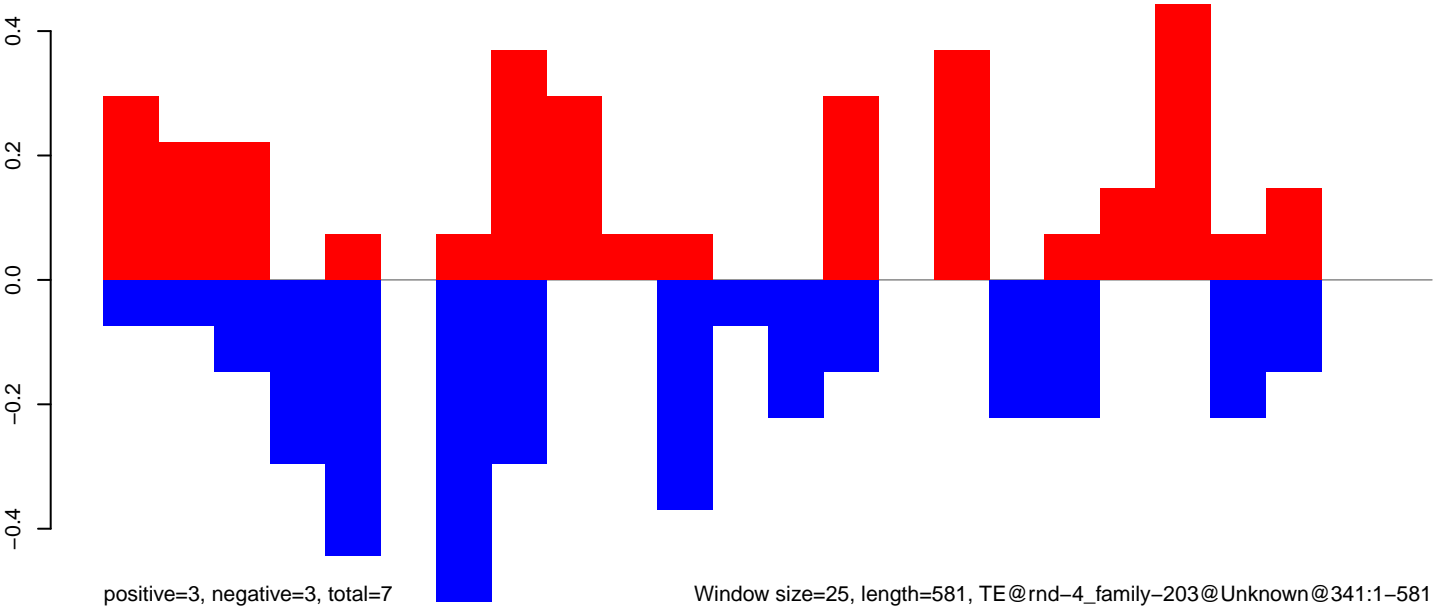
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

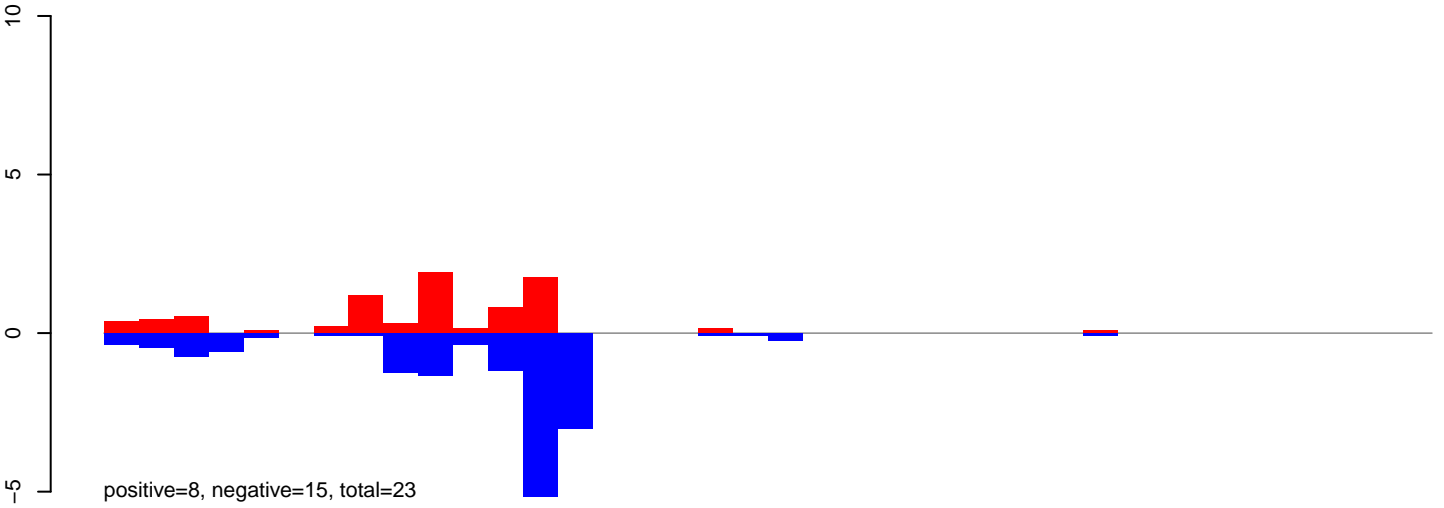


CuQuin_Testes_NL.rep

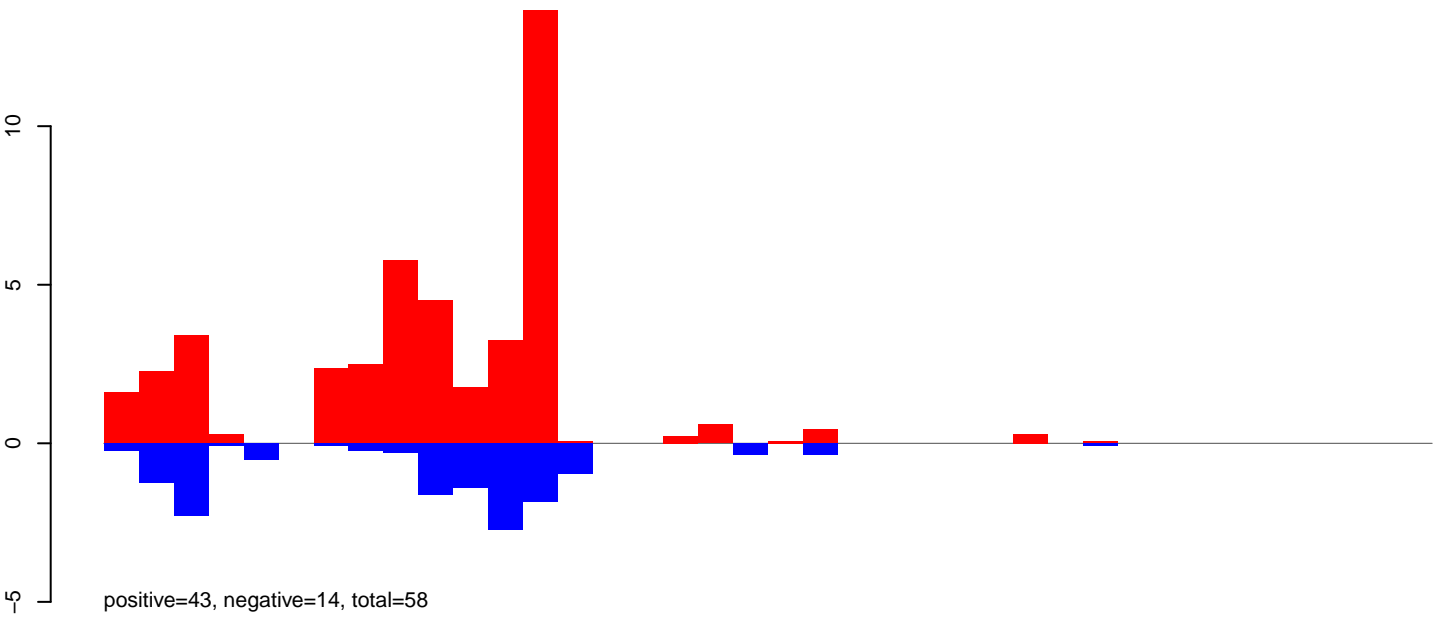


Window size=25, length=581, TE@rnd-4_family-203@Unknown@341:1-581

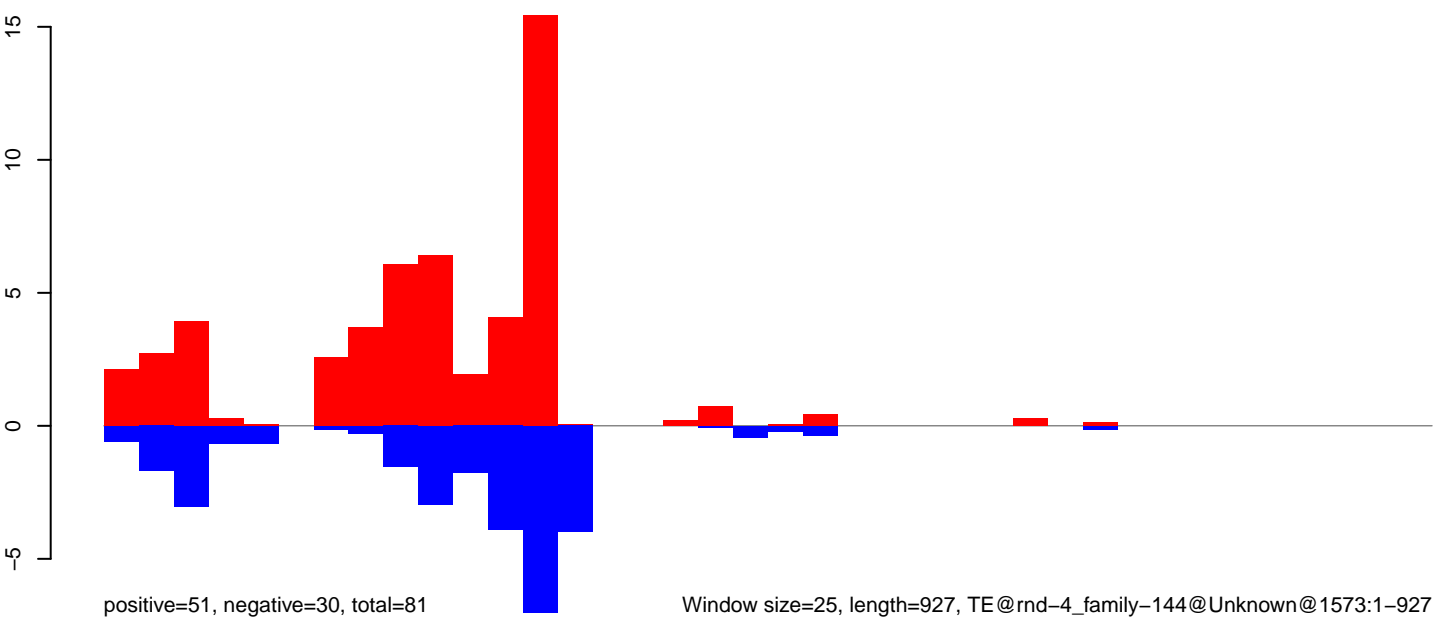
CuQuin_Testes_NL.18_23.rep



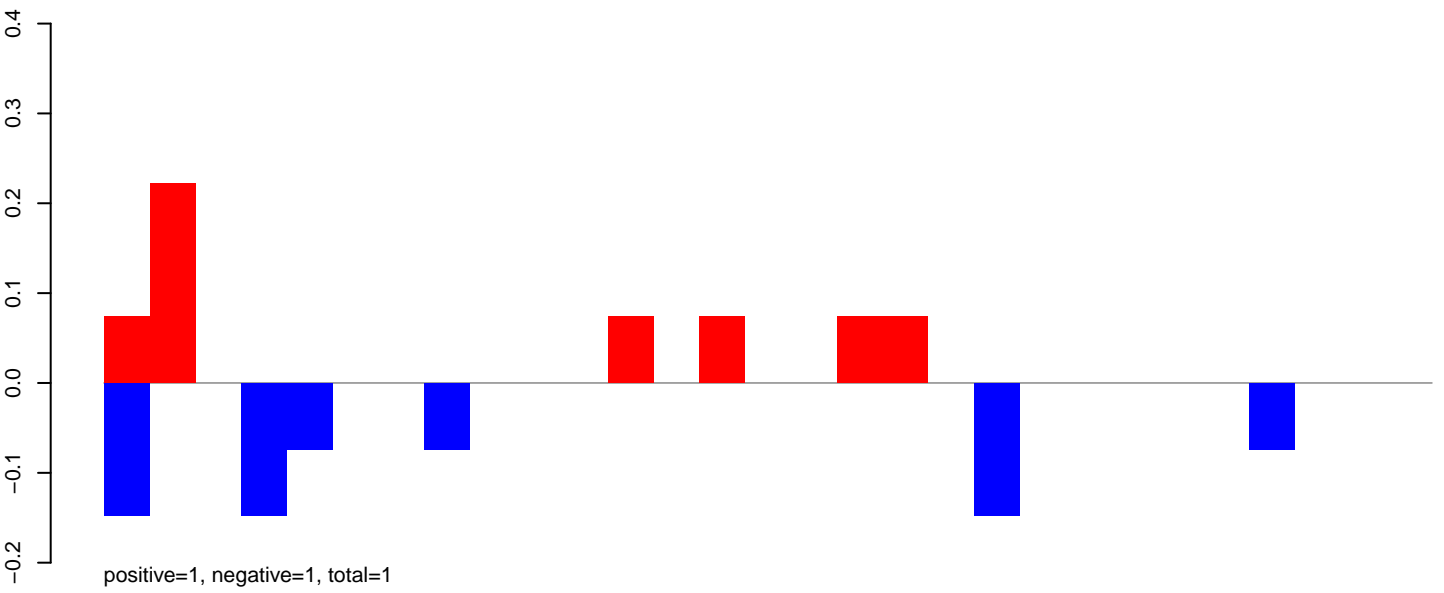
CuQuin_Testes_NL.24_35.rep



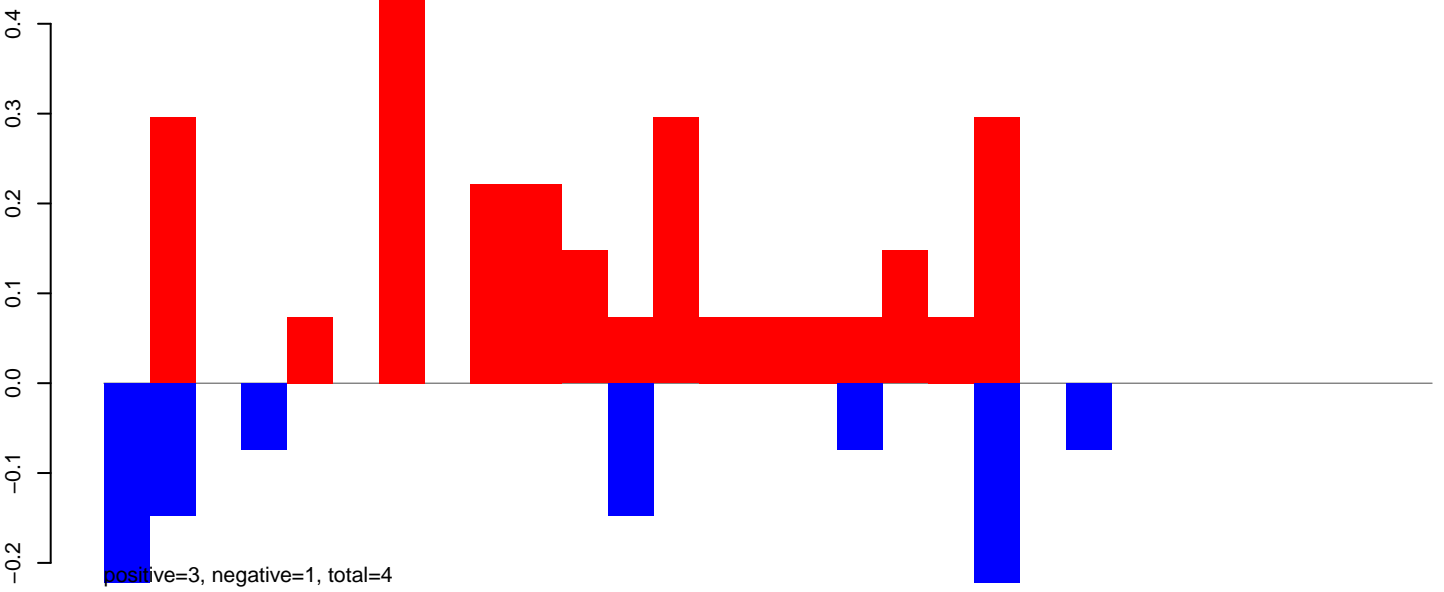
CuQuin_Testes_NL.rep



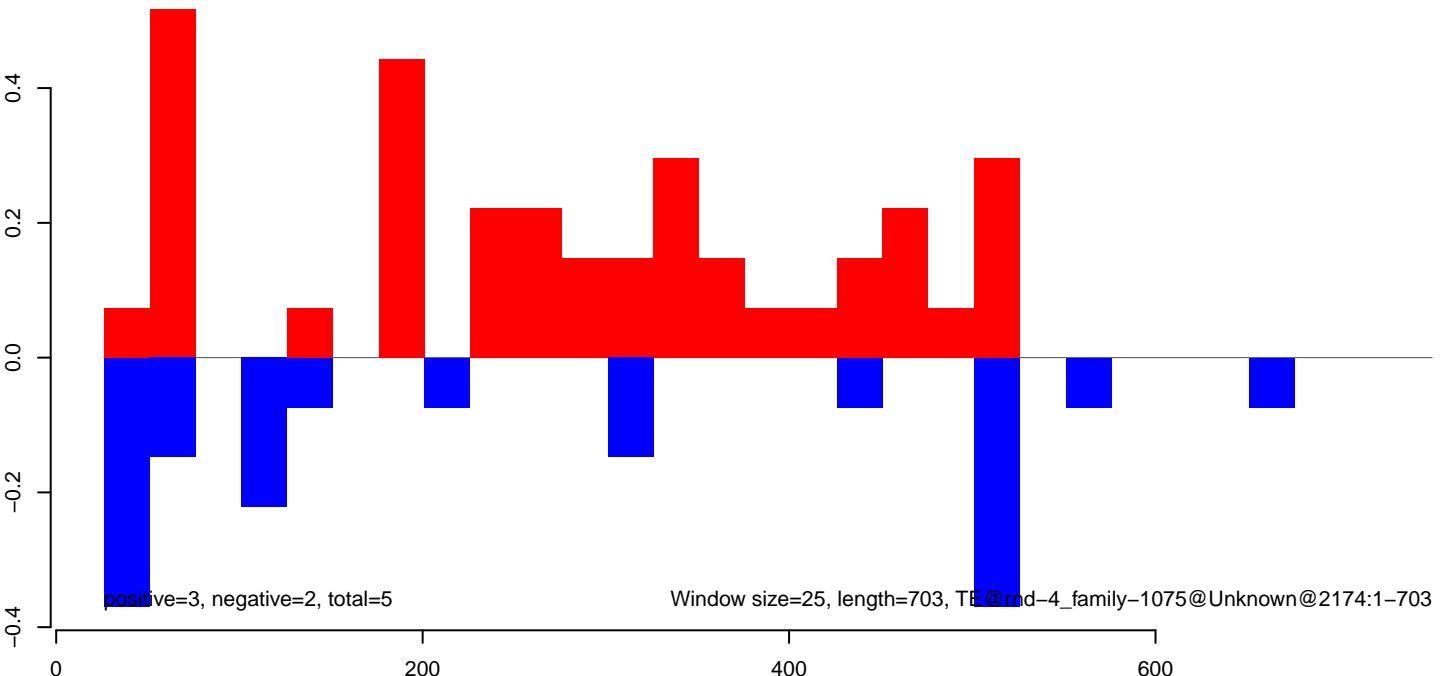
CuQuin_Testes_NL.18_23.rep



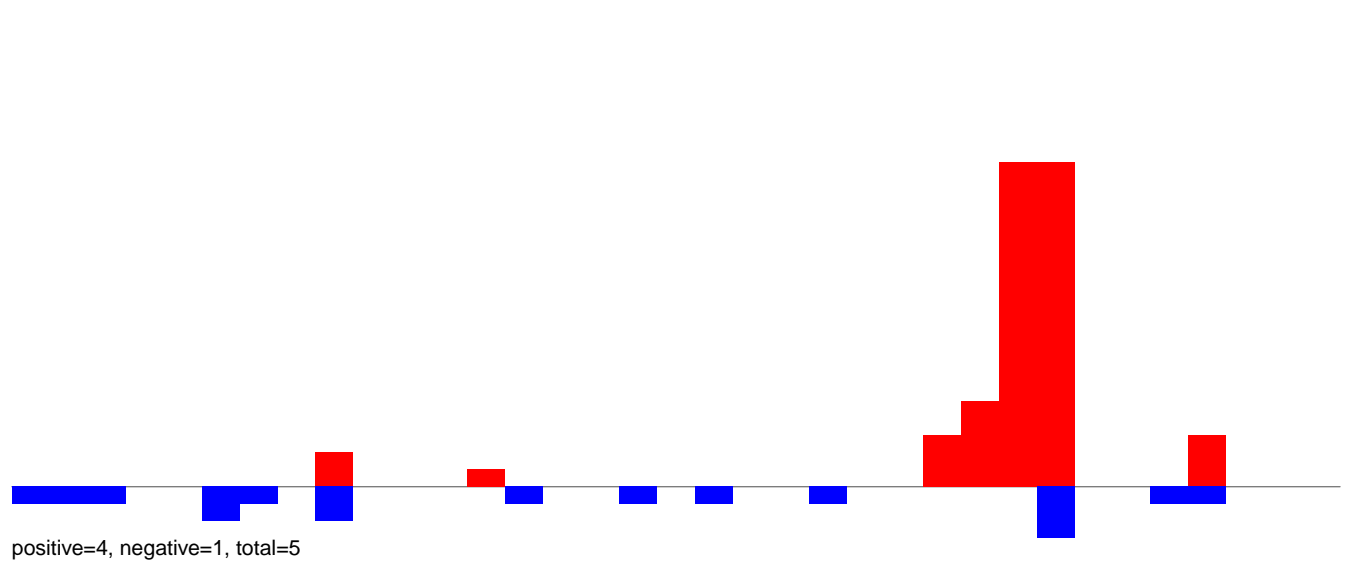
CuQuin_Testes_NL.24_35.rep



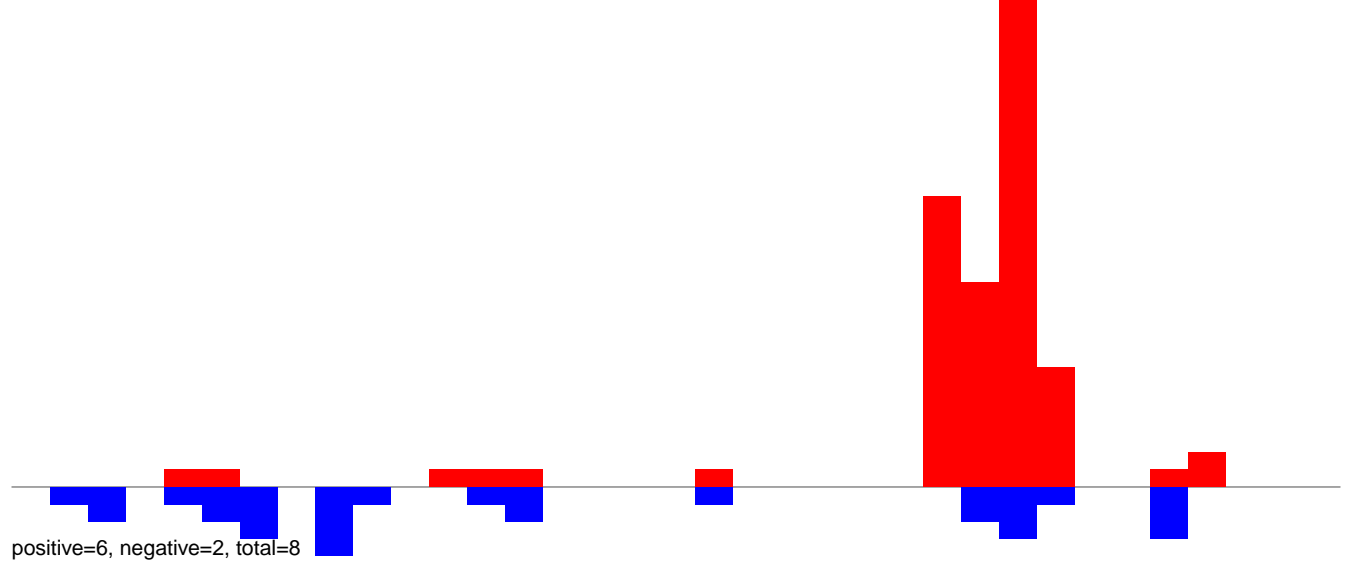
CuQuin_Testes_NL.rep



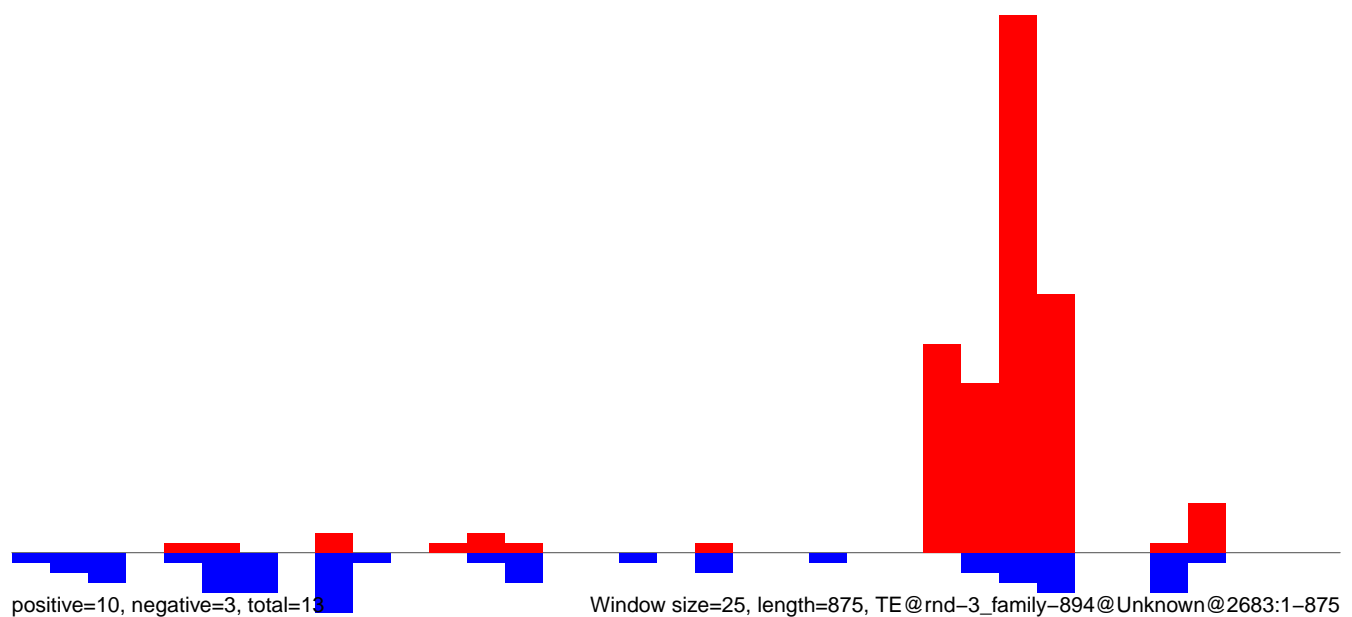
CuQuin_Testes_NL.18_23.rep



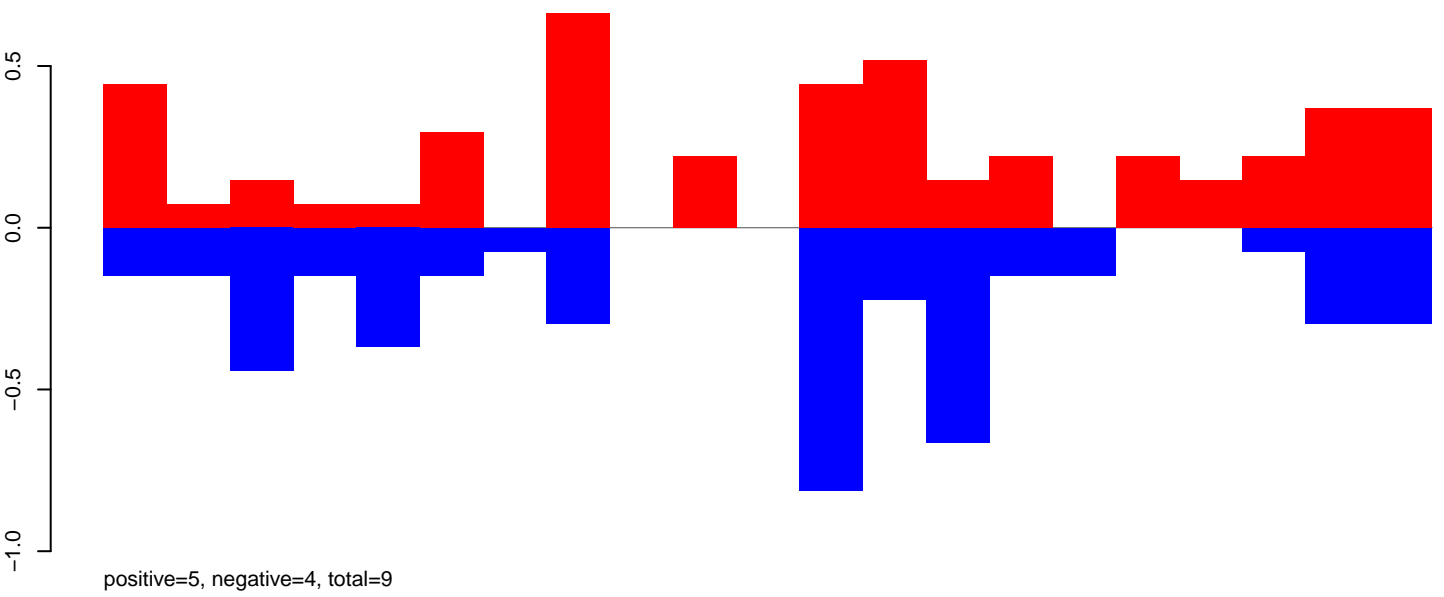
CuQuin_Testes_NL.24_35.rep



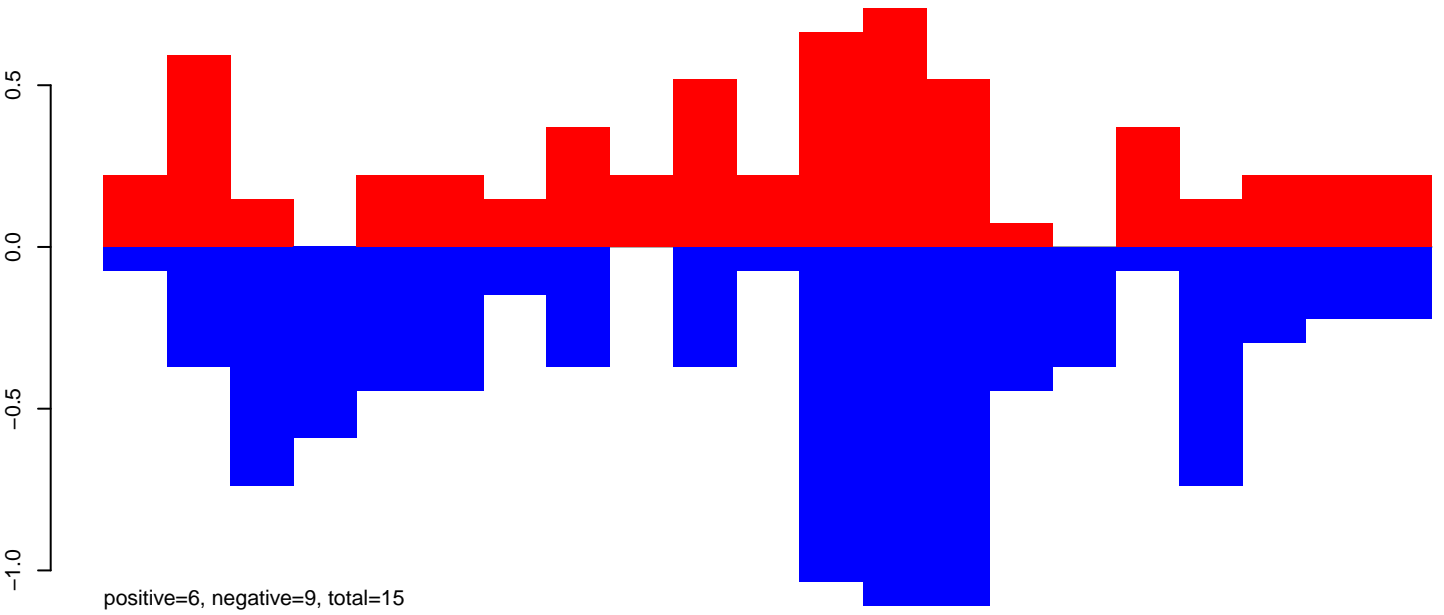
CuQuin_Testes_NL.rep



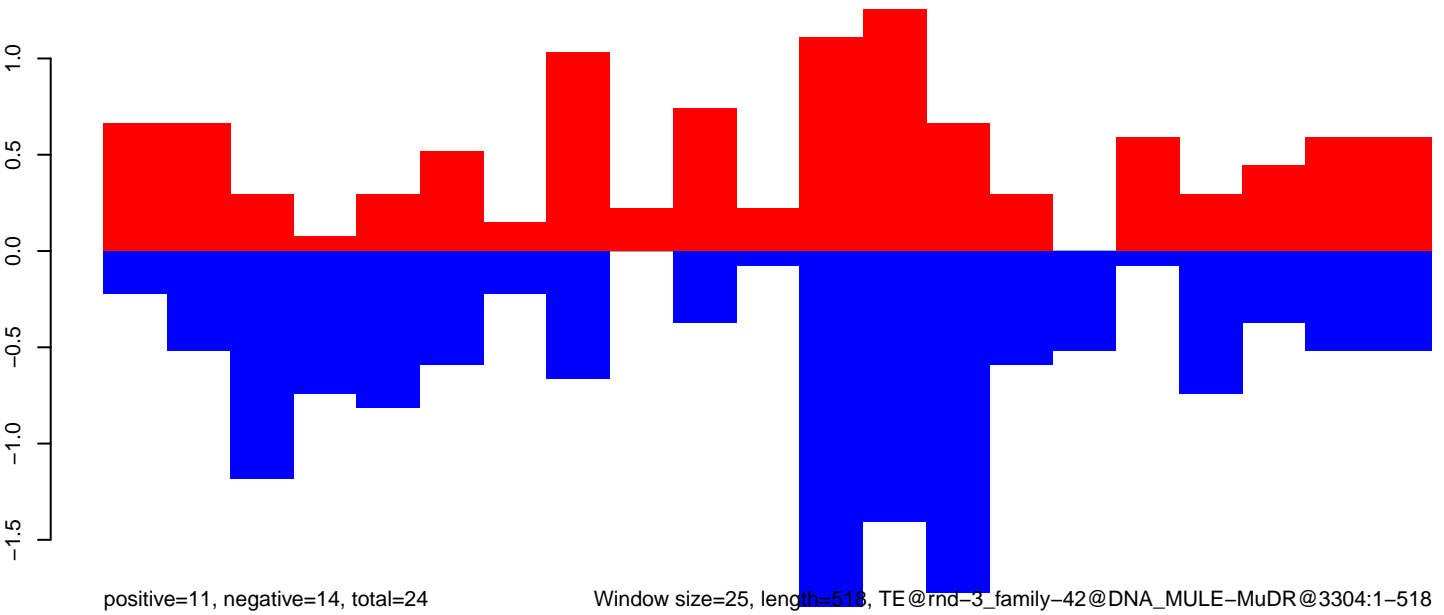
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

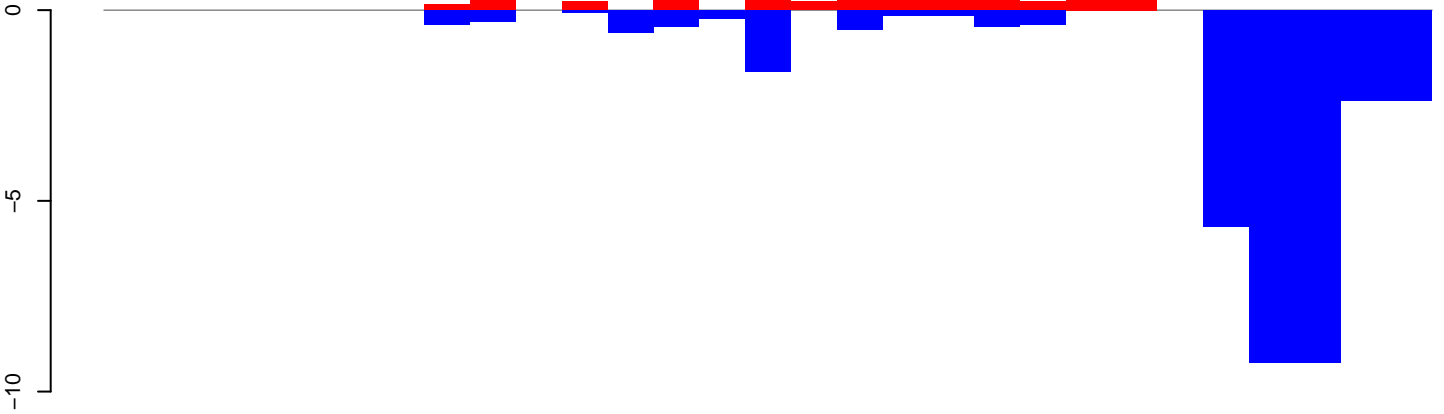


CuQuin_Testes_NL.rep



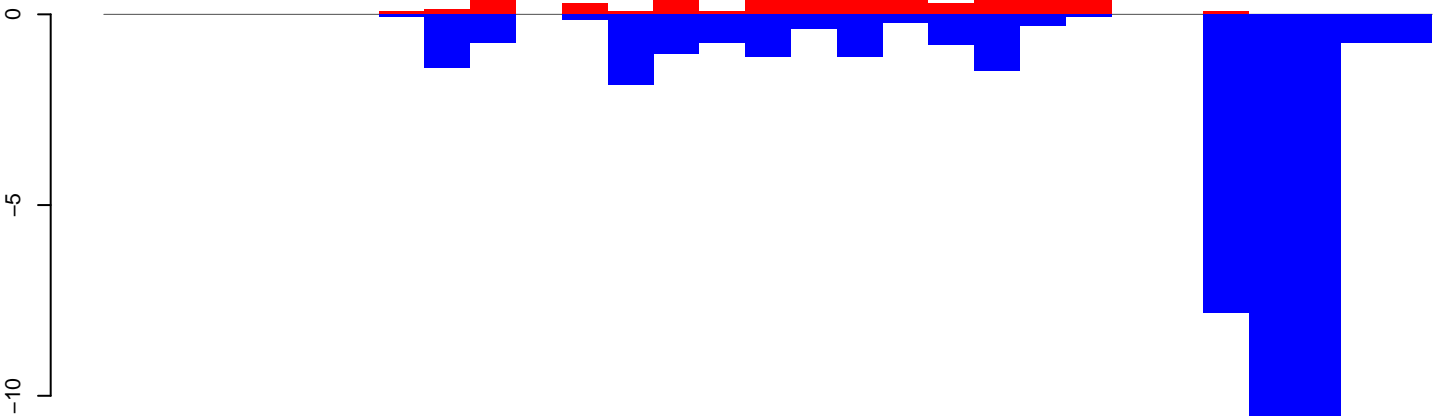
Window size=25, length=518, TE@rnd-3_family-42@DNA_MULE-MuDR@3304:1-518

CuQuin_Testes_NL.18_23.rep



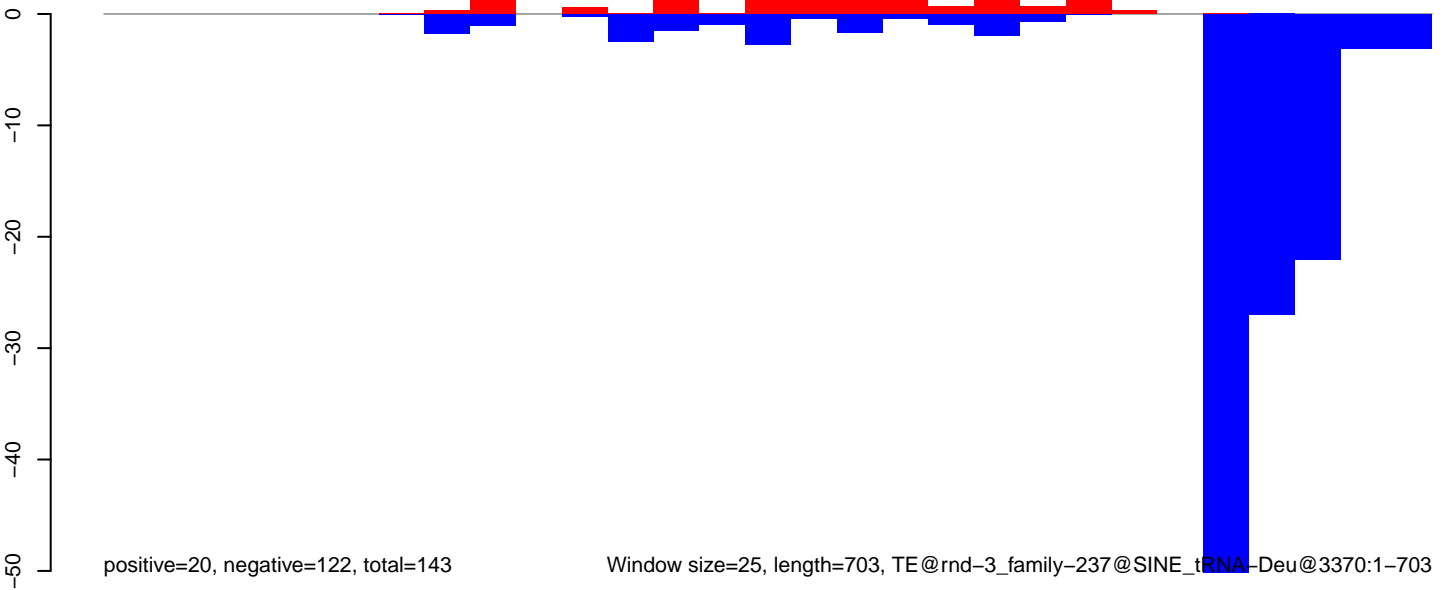
positive=7, negative=34, total=41

CuQuin_Testes_NL.24_35.rep



positive=14, negative=47, total=60

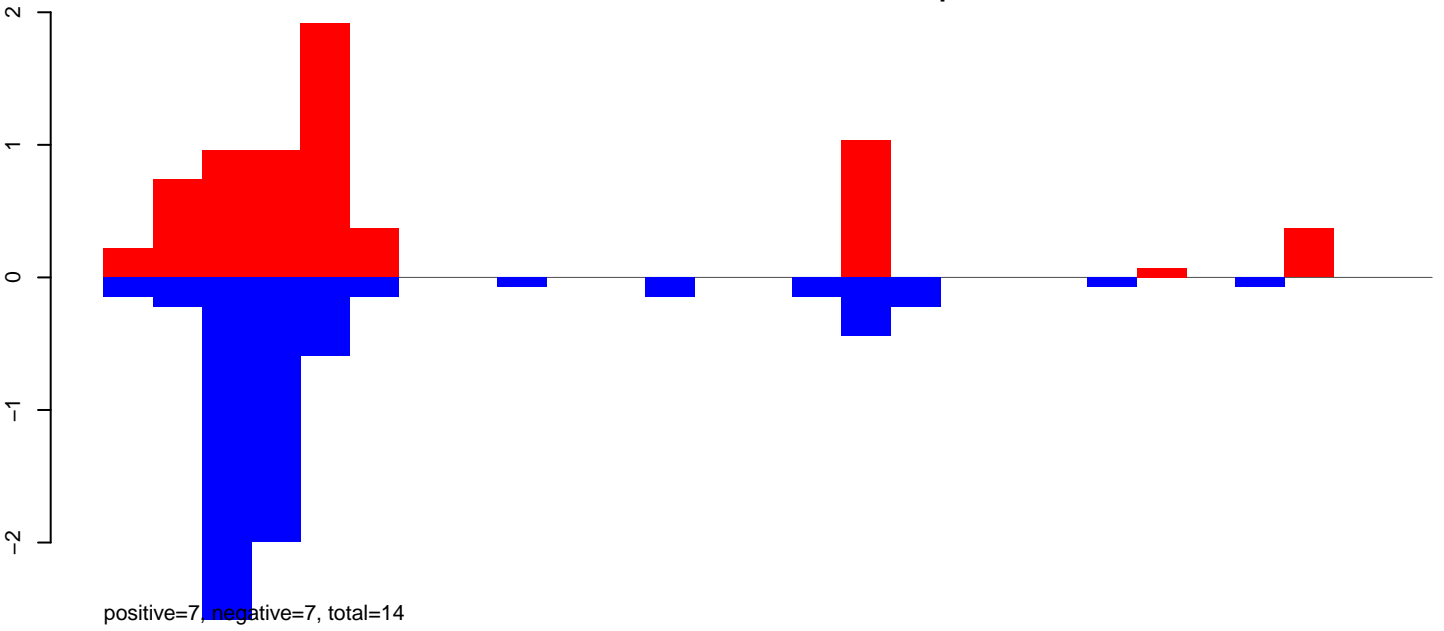
CuQuin_Testes_NL.rep



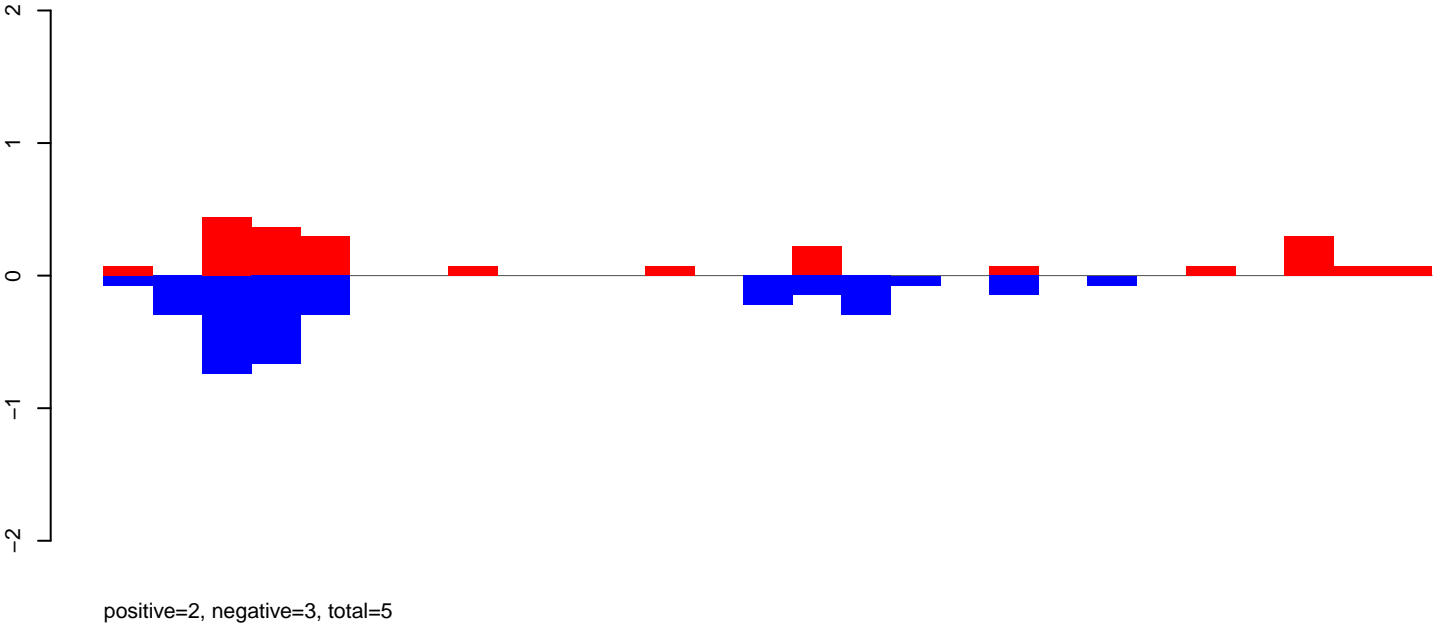
positive=20, negative=122, total=143

Window size=25, length=703, TE@rnd-3_family-237@SINE_tRNA-Deu@3370:1-703

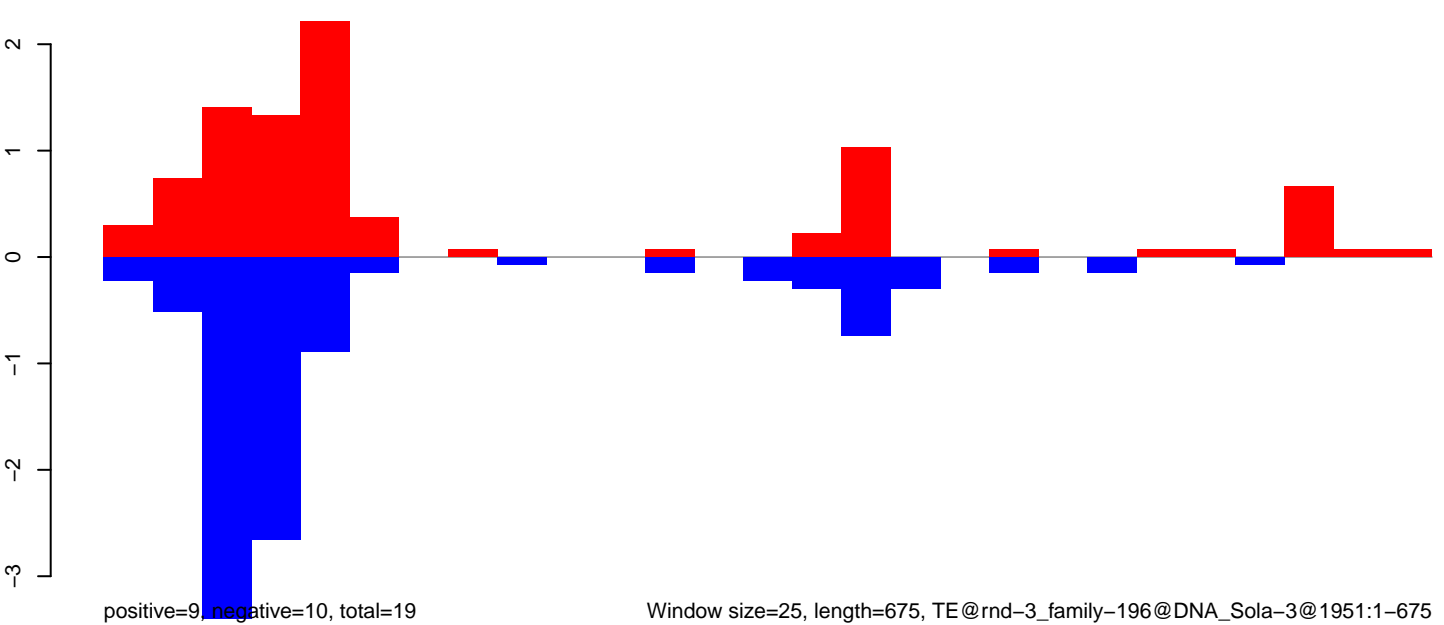
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

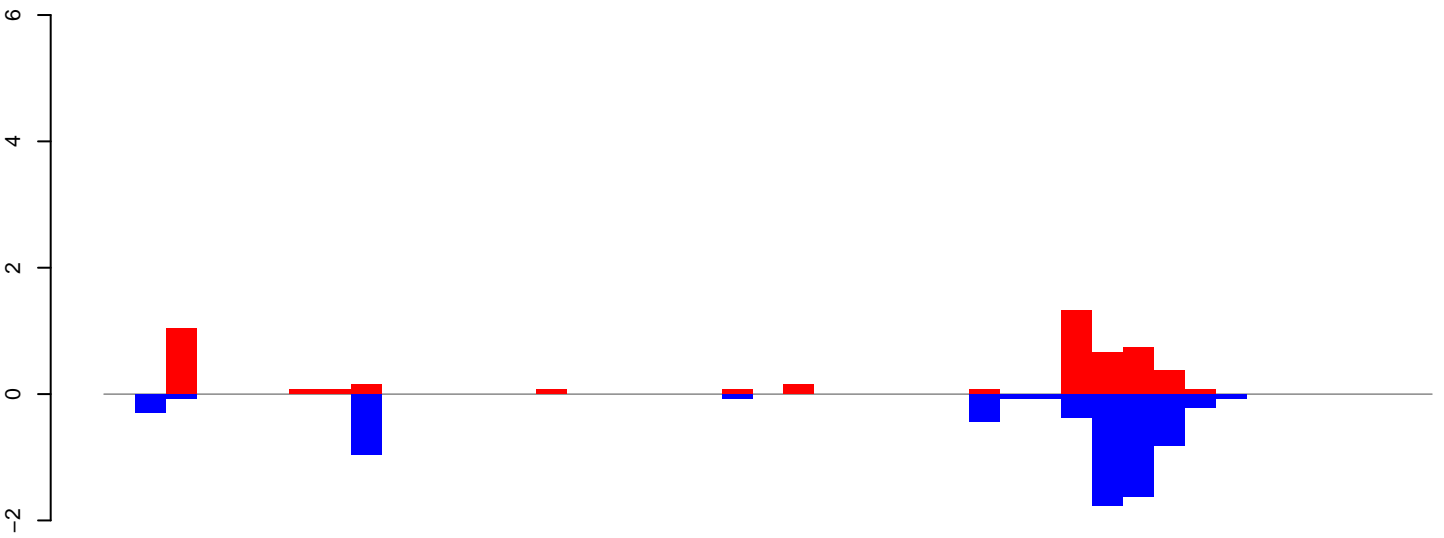


CuQuin_Testes_NL.rep



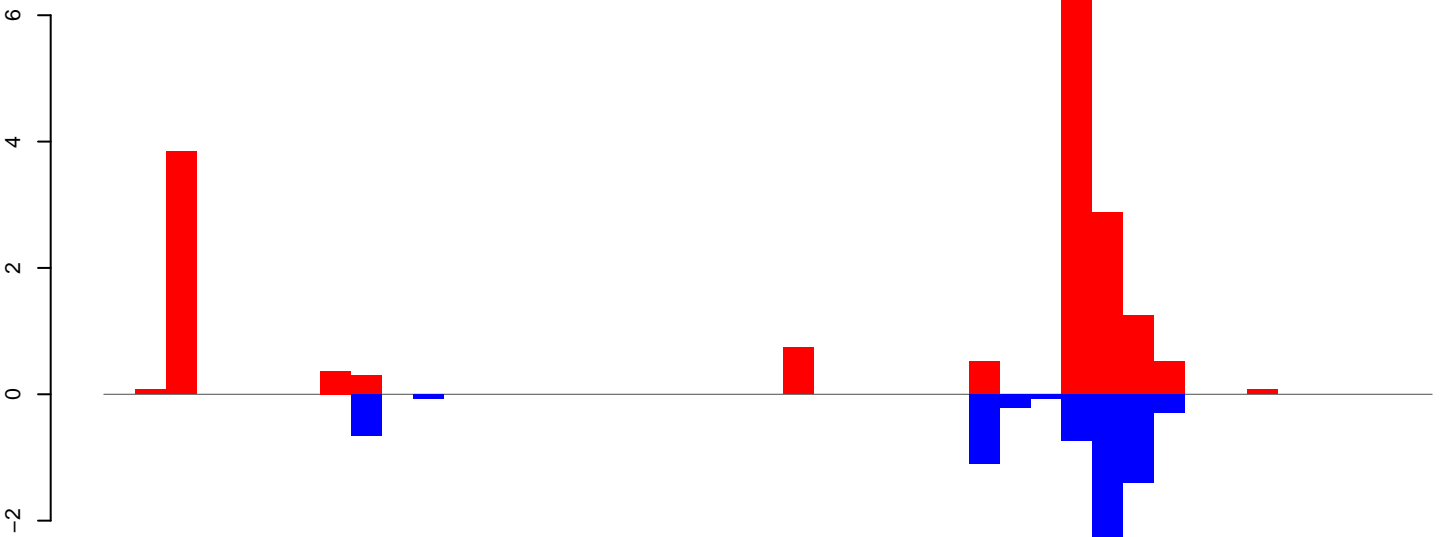
Window size=25, length=675, TE@rnd-3_family-196@DNA_Sola-3@1951:1-675

CuQuin_Testes_NL.18_23.rep



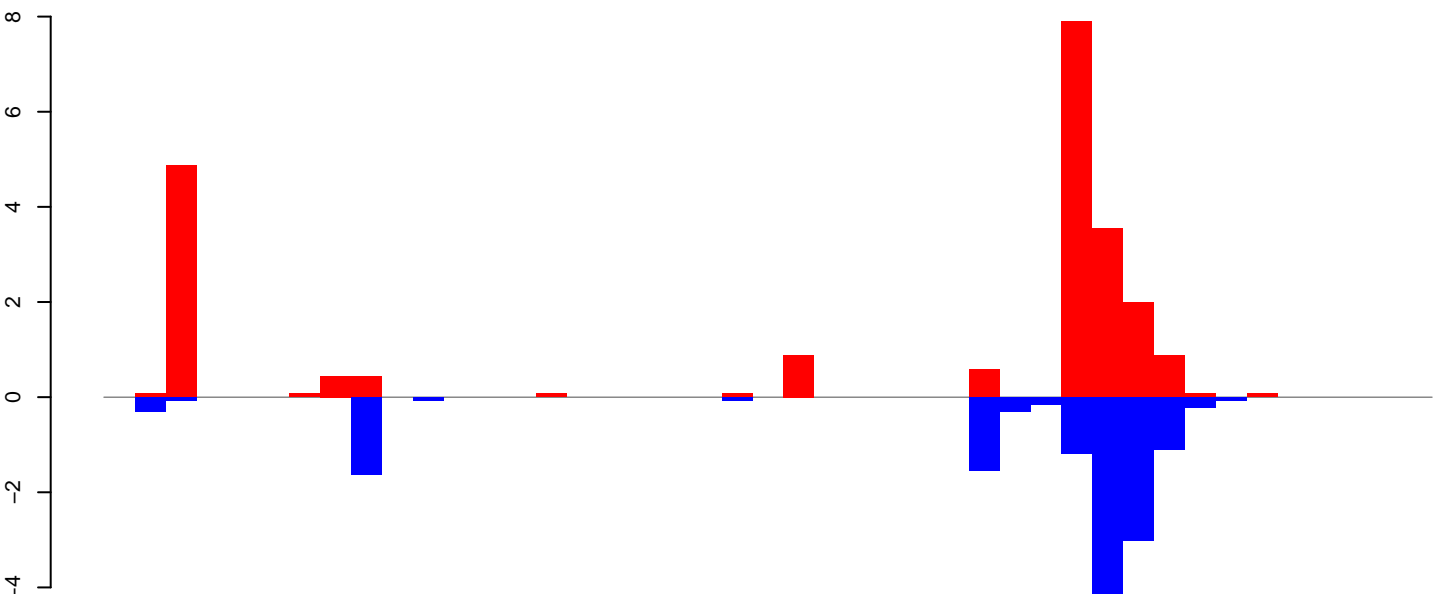
positive=5, negative=7, total=12

CuQuin_Testes_NL.24_35.rep



positive=17, negative=7, total=25

CuQuin_Testes_NL.rep

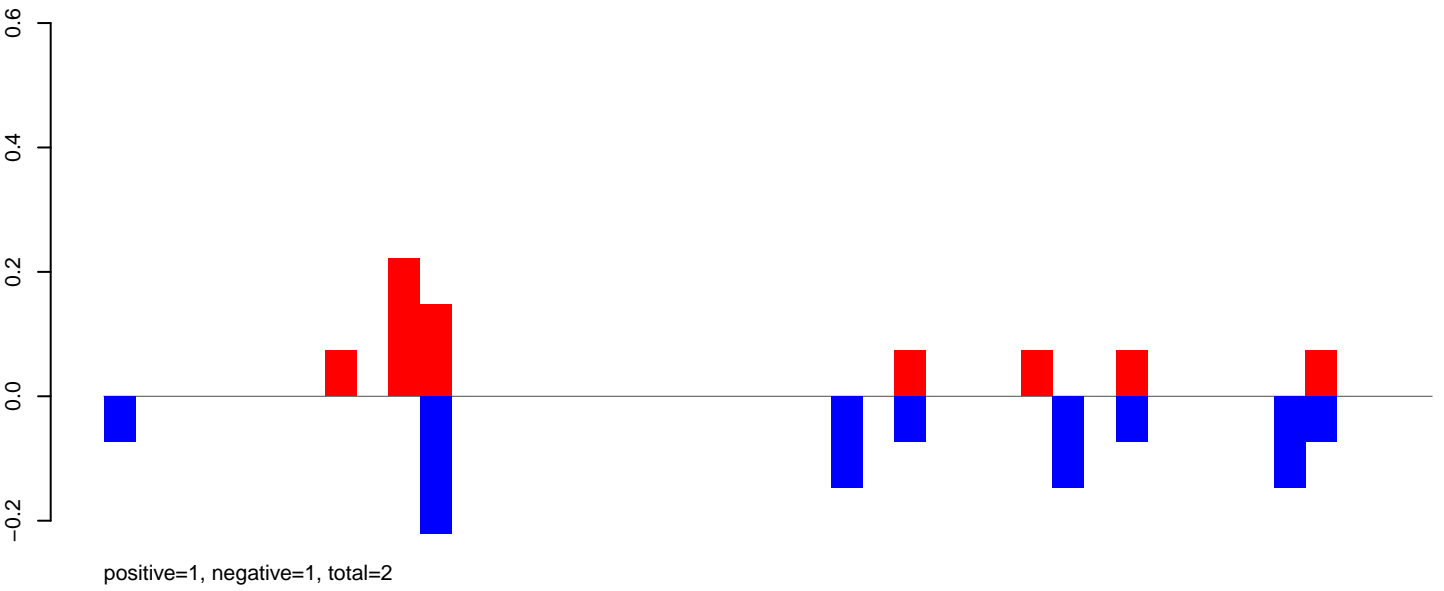


positive=22, negative=14, total=36

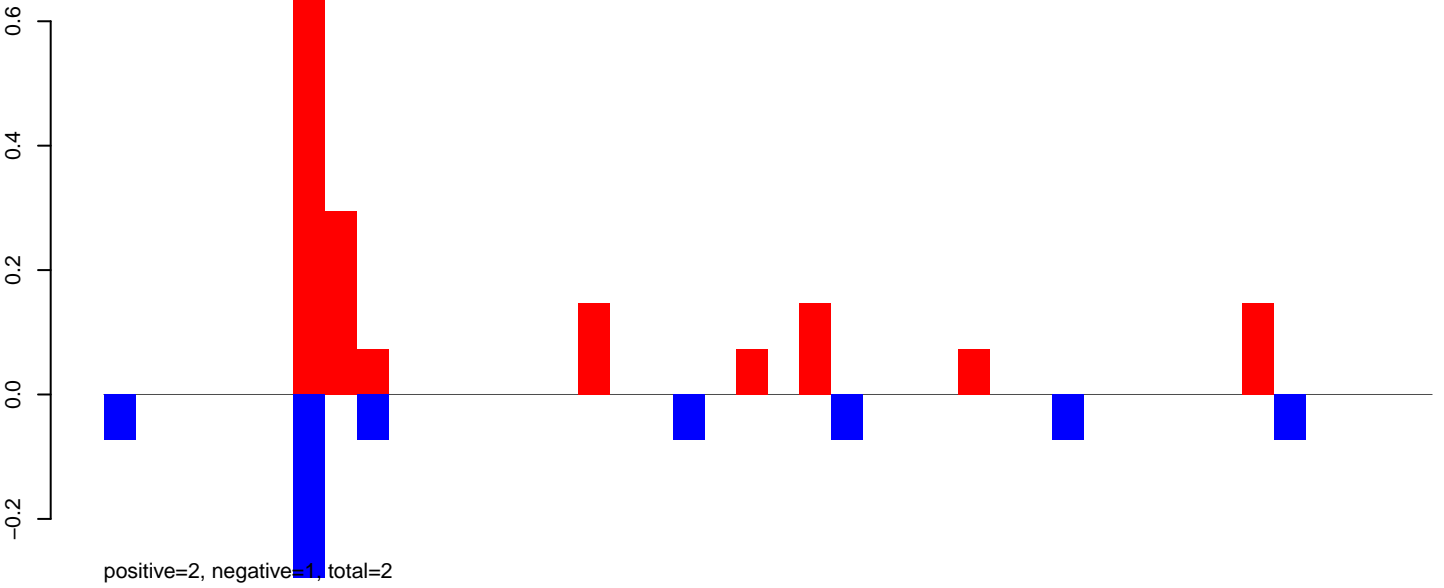
Window size=25, length=1067, TE@rnd-5_family-98@Unknown@1144:1-1067

0 200 400 600 800 1000

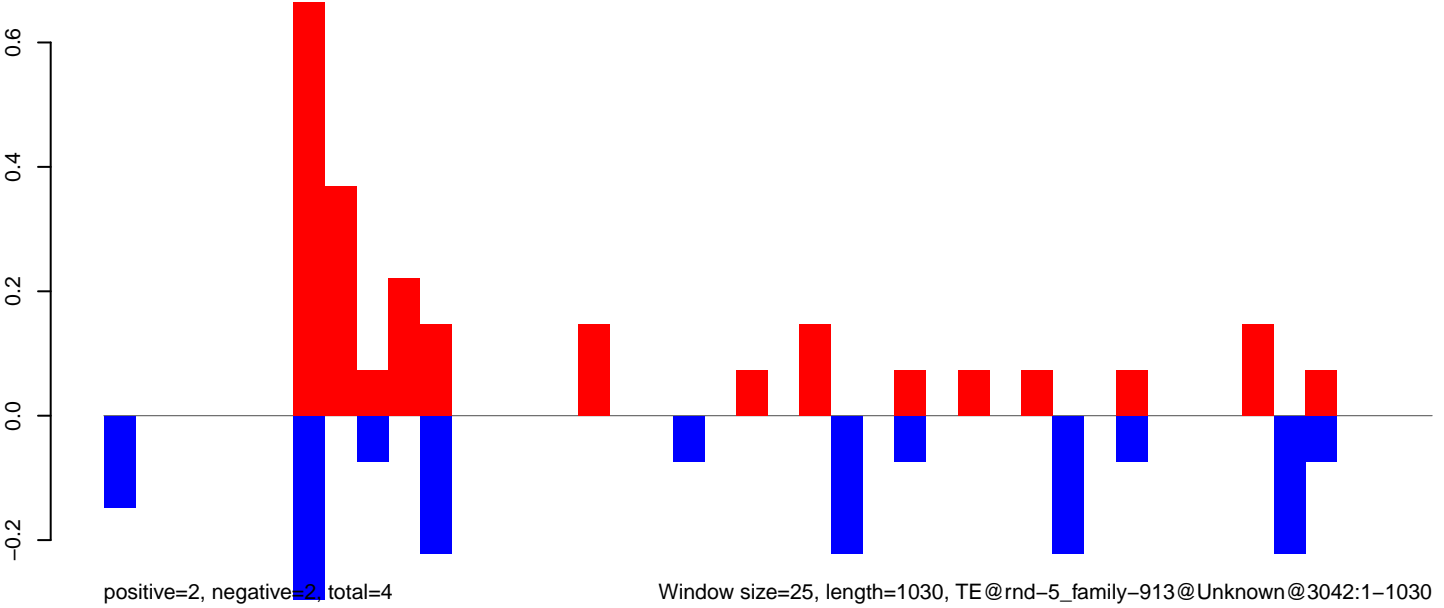
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



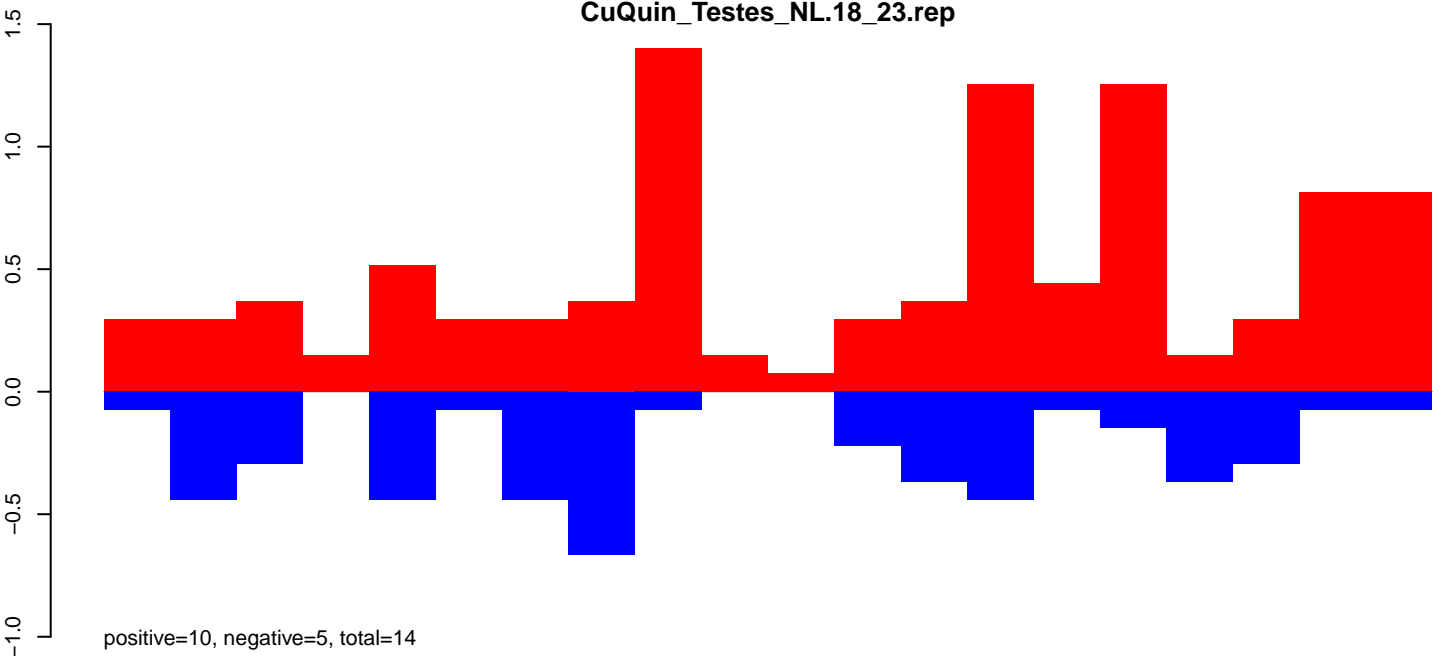
CuQuin_Testes_NL.rep



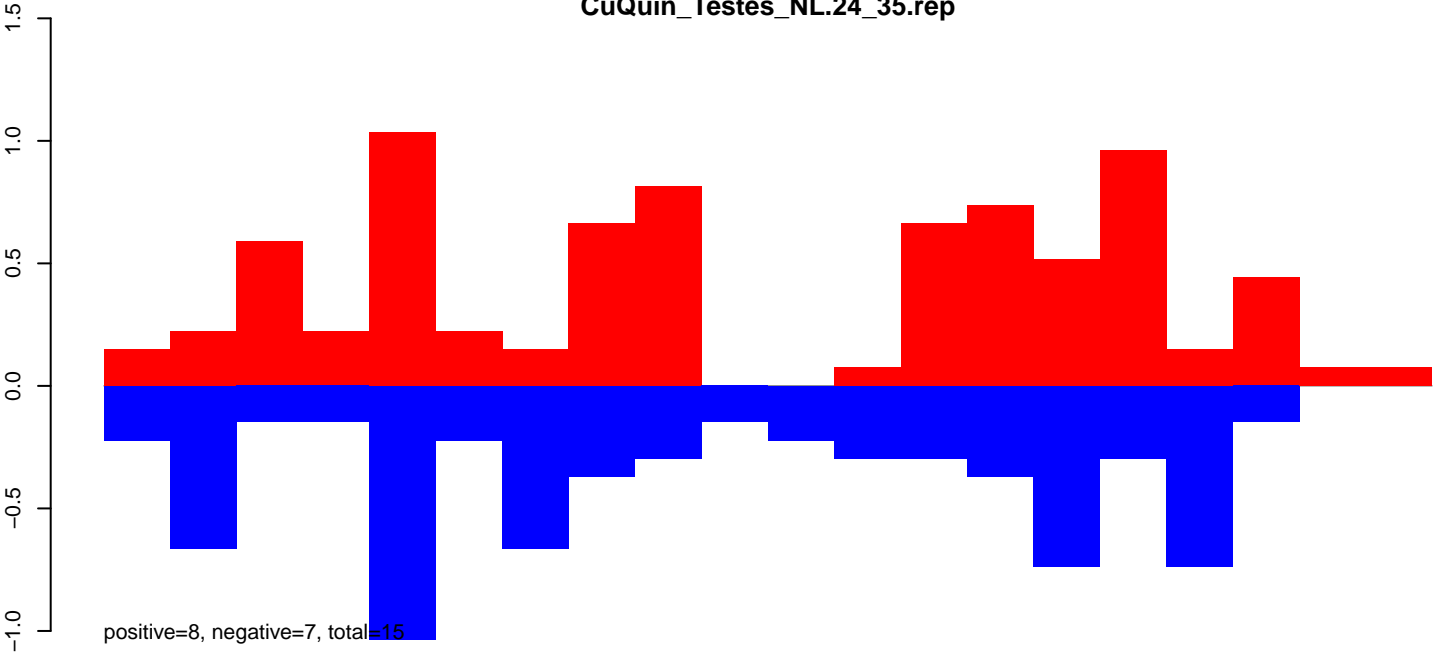
Window size=25, length=1030, TE@rnd-5_family-913@Unknown@3042:1-1030

0 200 400 600 800 1000

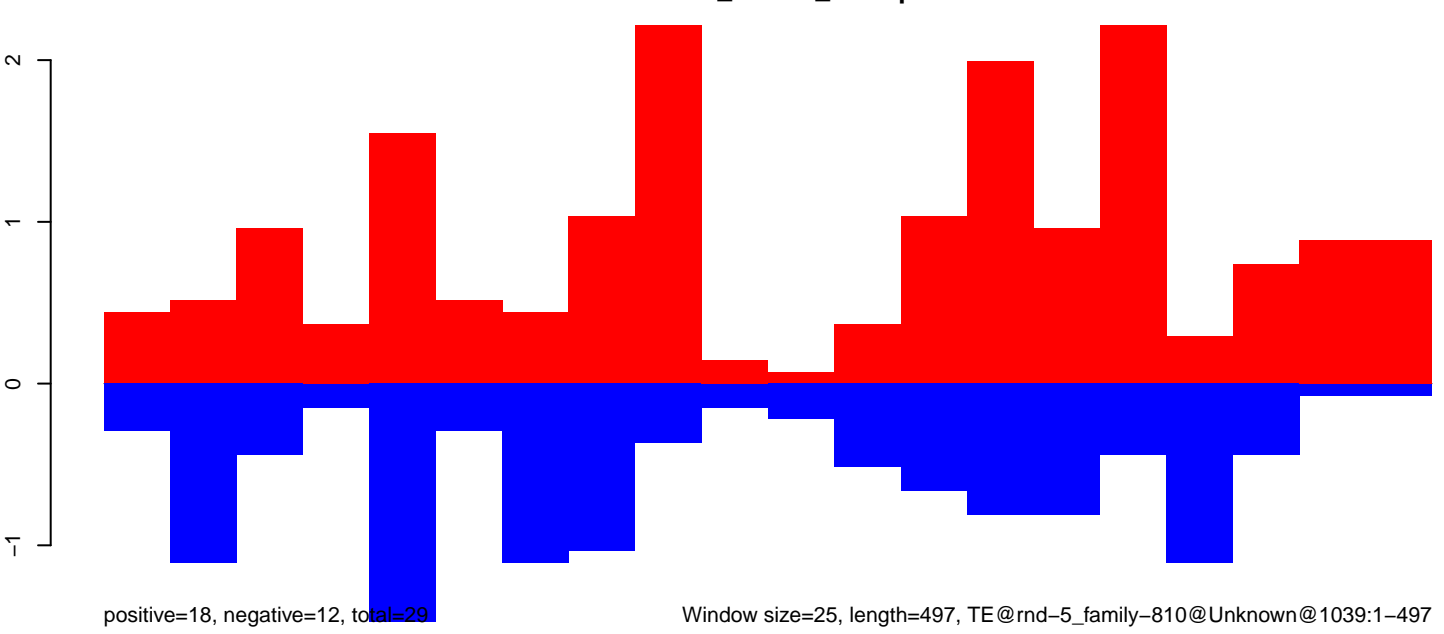
CuQuin_Testes_NL.18_23.rep



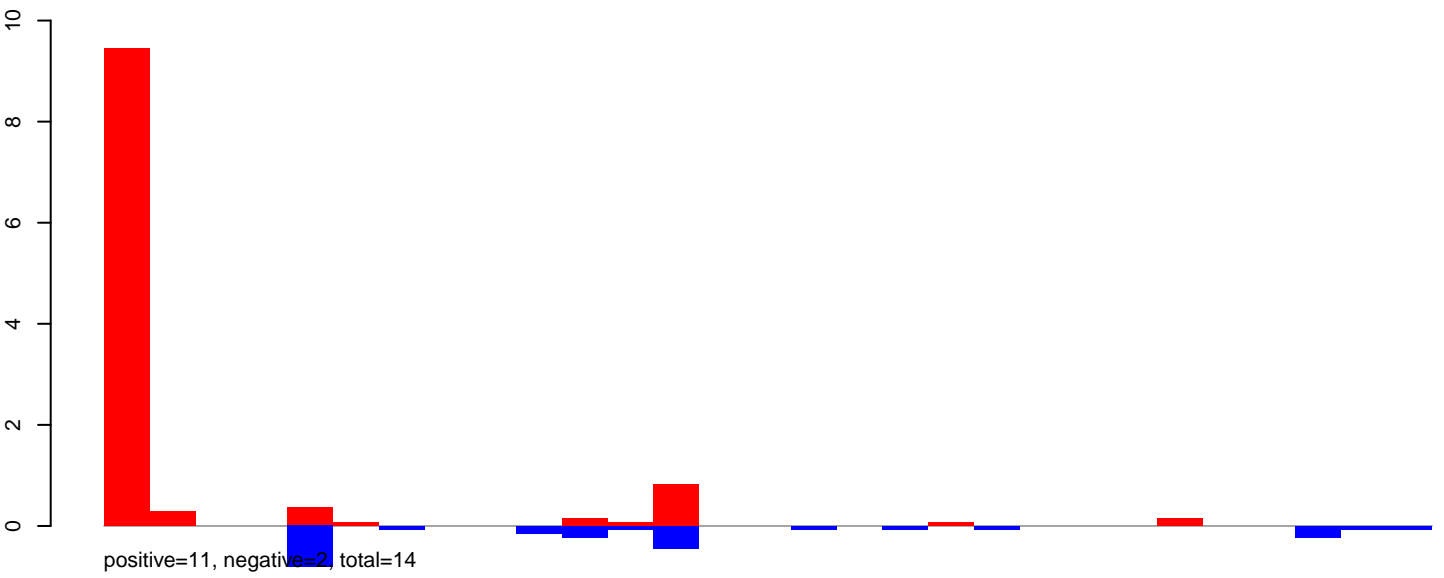
CuQuin_Testes_NL.24_35.rep



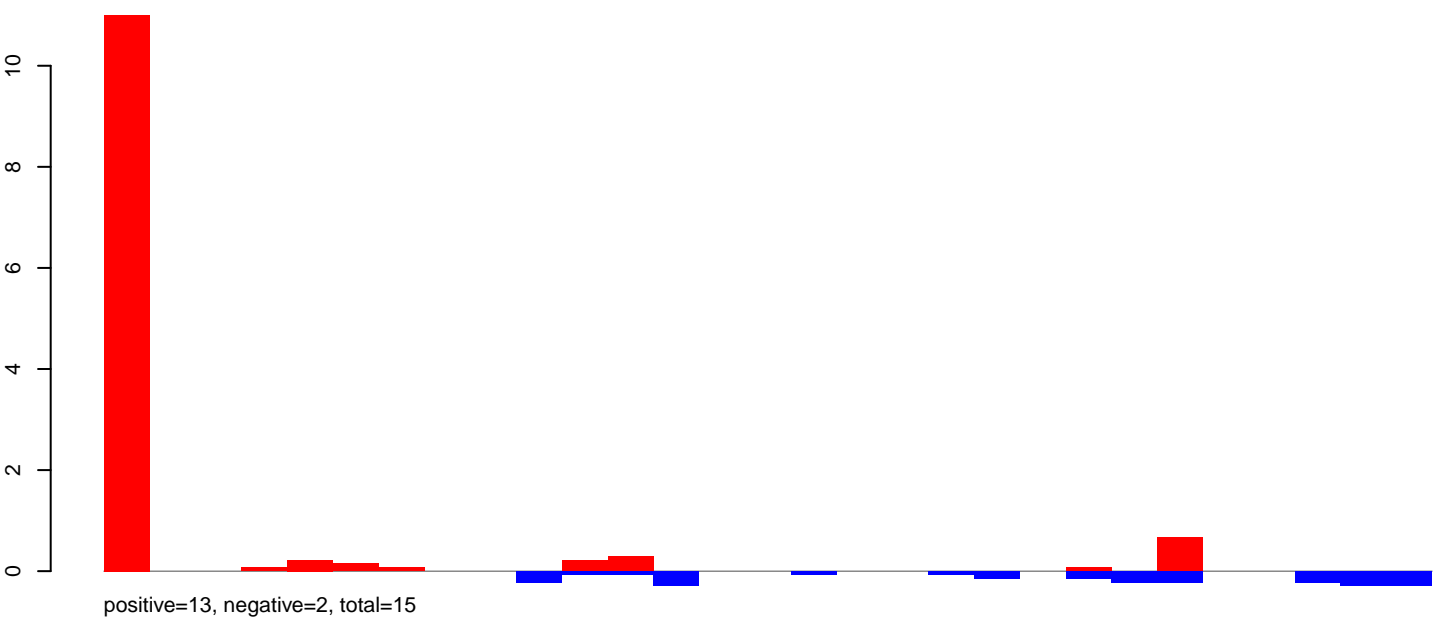
CuQuin_Testes_NL.rep



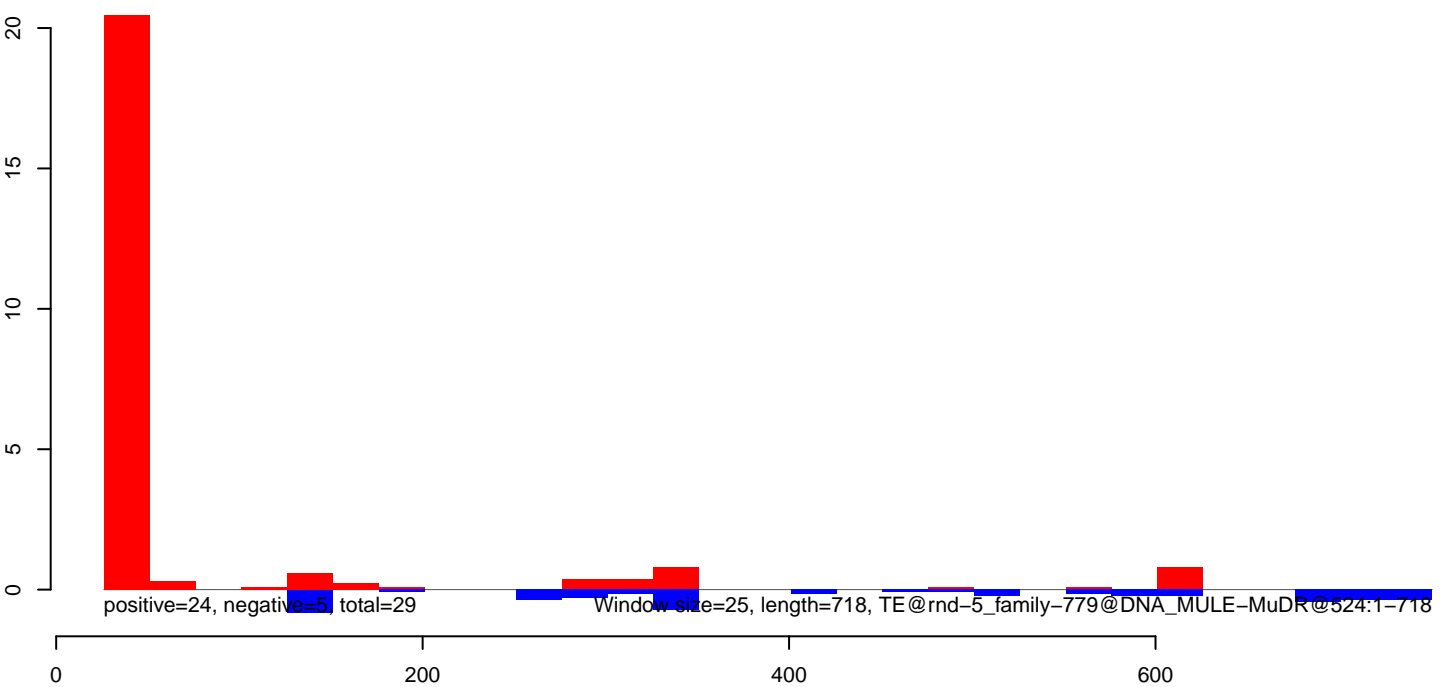
CuQuin_Testes_NL.18_23.rep



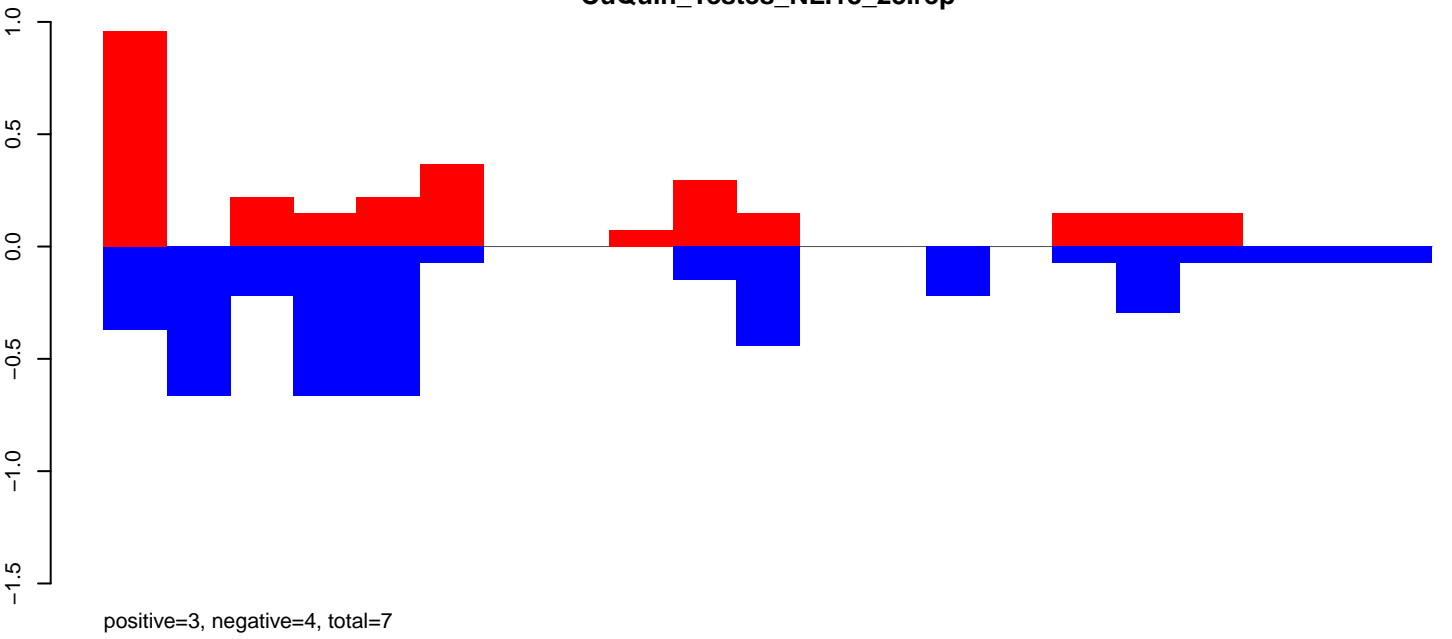
CuQuin_Testes_NL.24_35.rep



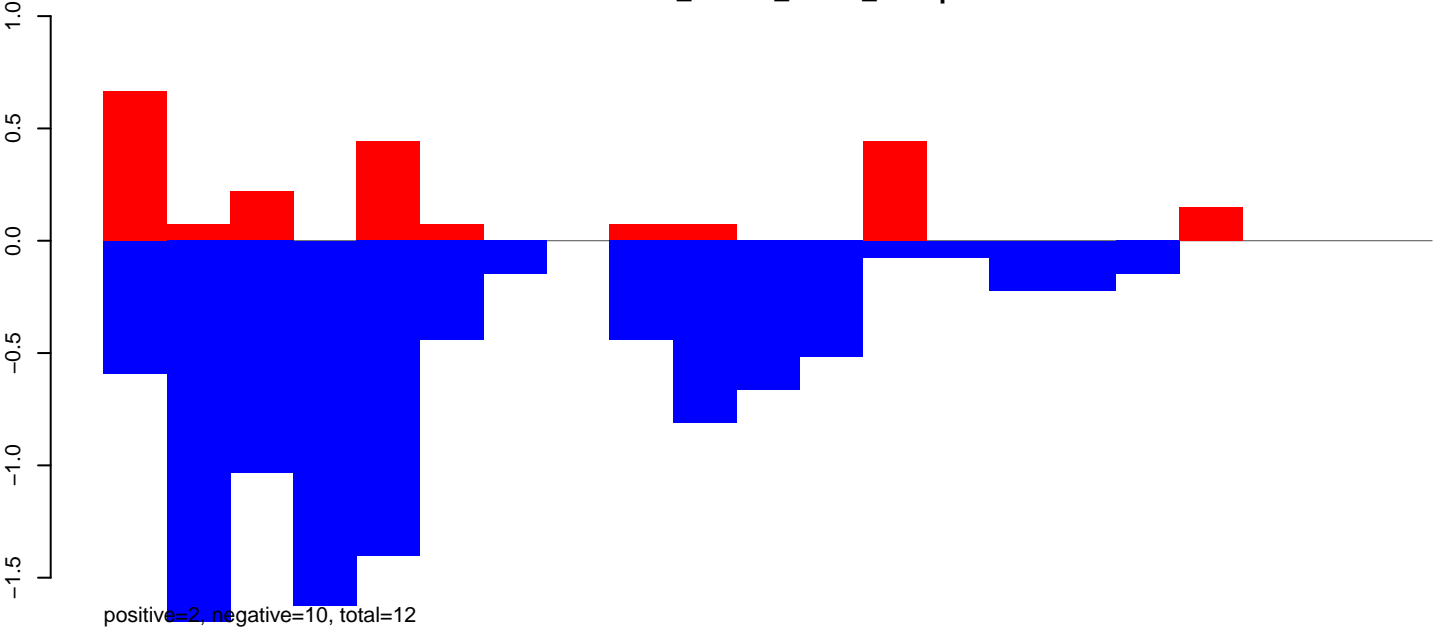
CuQuin_Testes_NL.rep



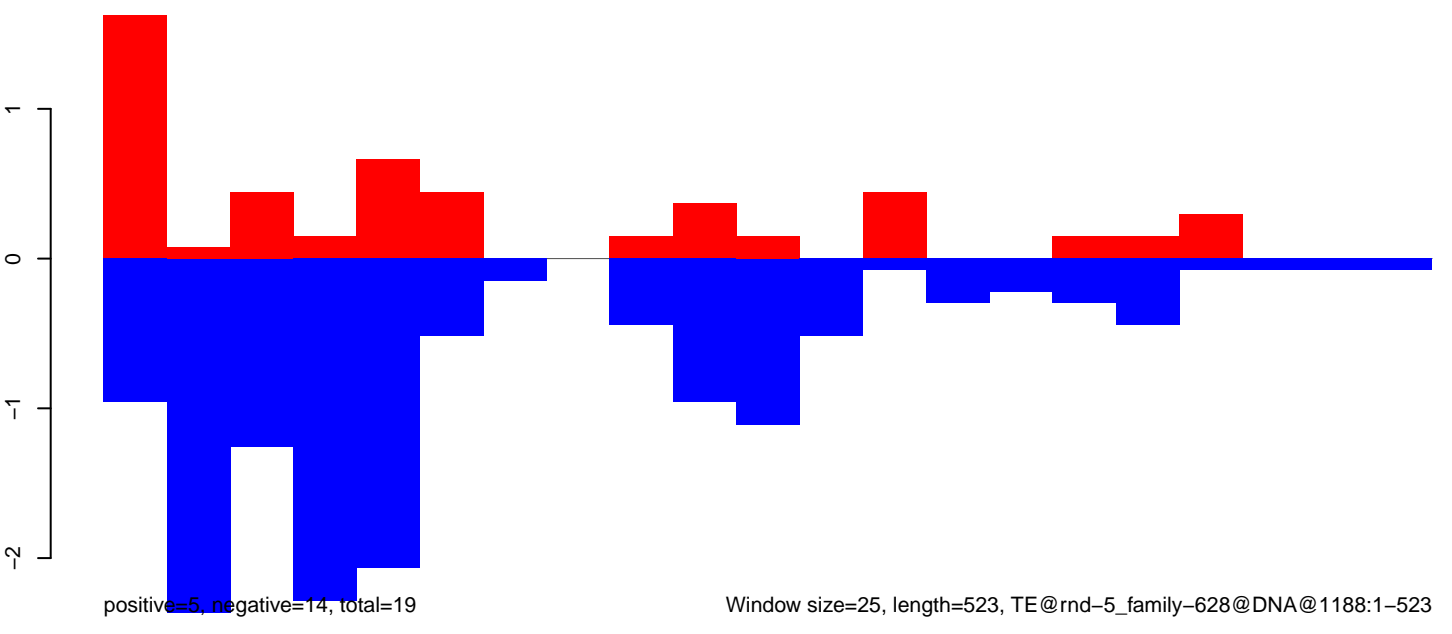
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

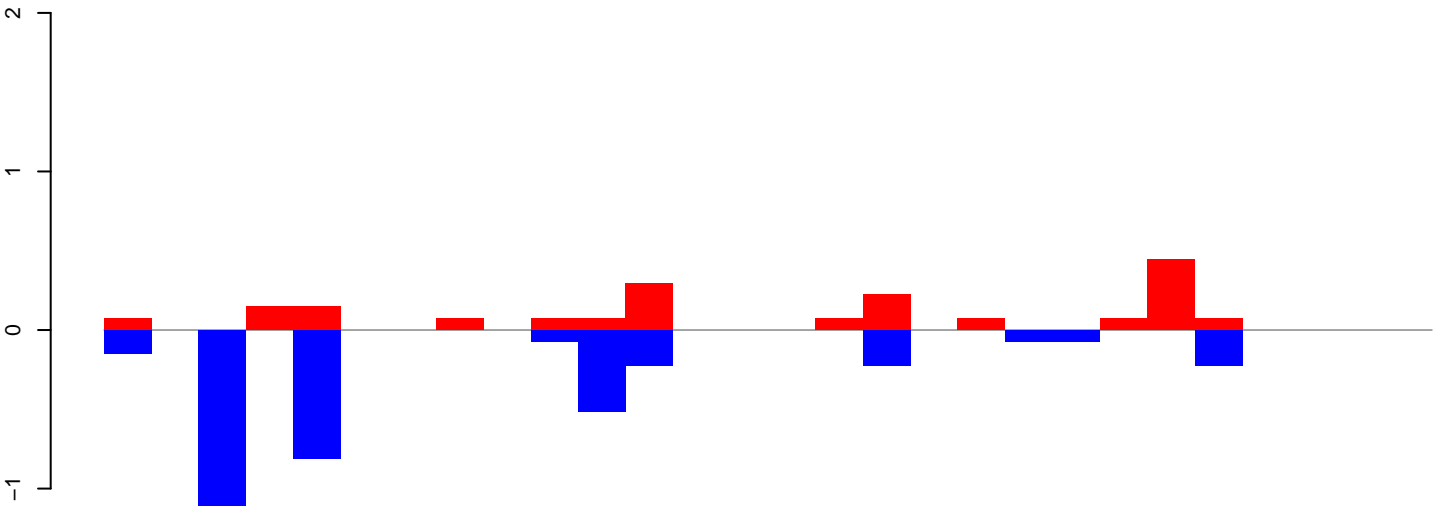


CuQuin_Testes_NL.rep



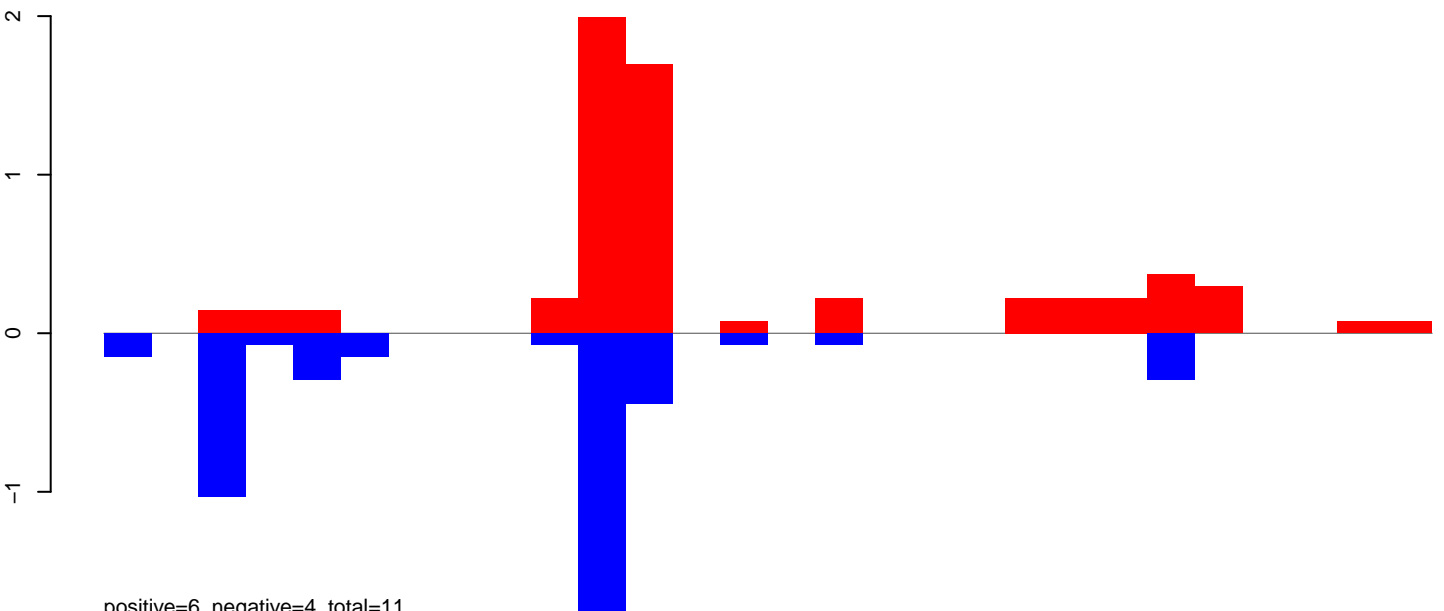
Window size=25, length=523, TE@rnd-5_family-628@DNA@1188:1-523

CuQuin_Testes_NL.18_23.rep



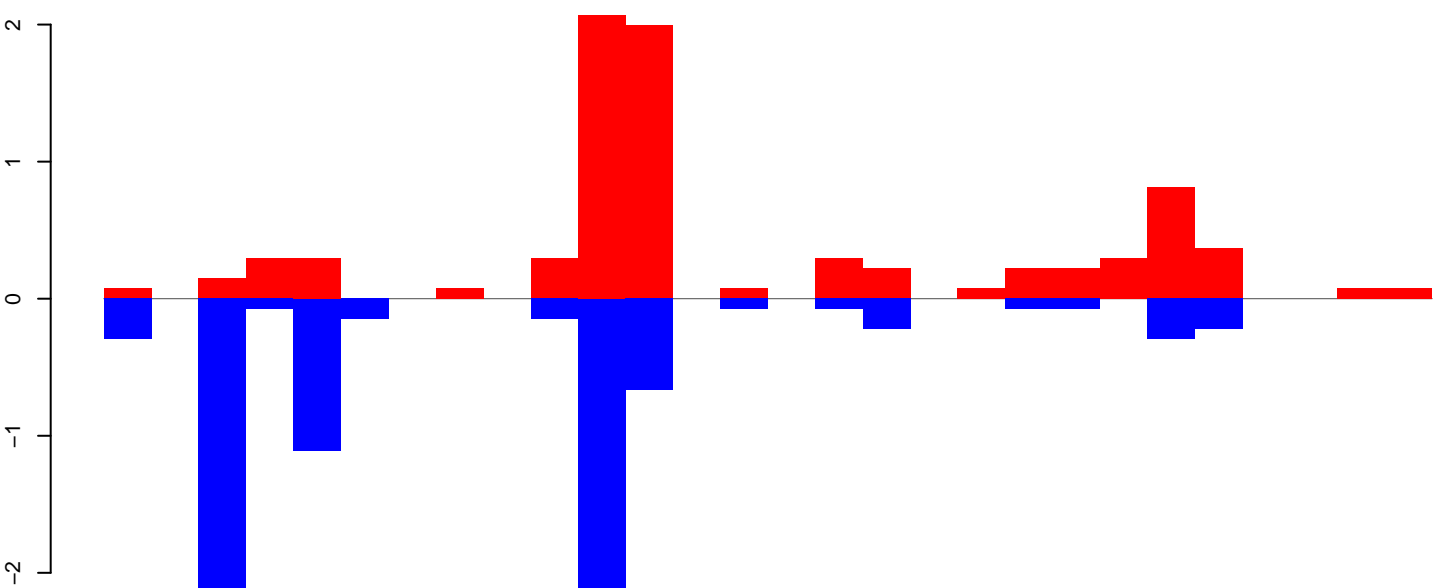
positive=2, negative=3, total=5

CuQuin_Testes_NL.24_35.rep



positive=6, negative=4, total=11

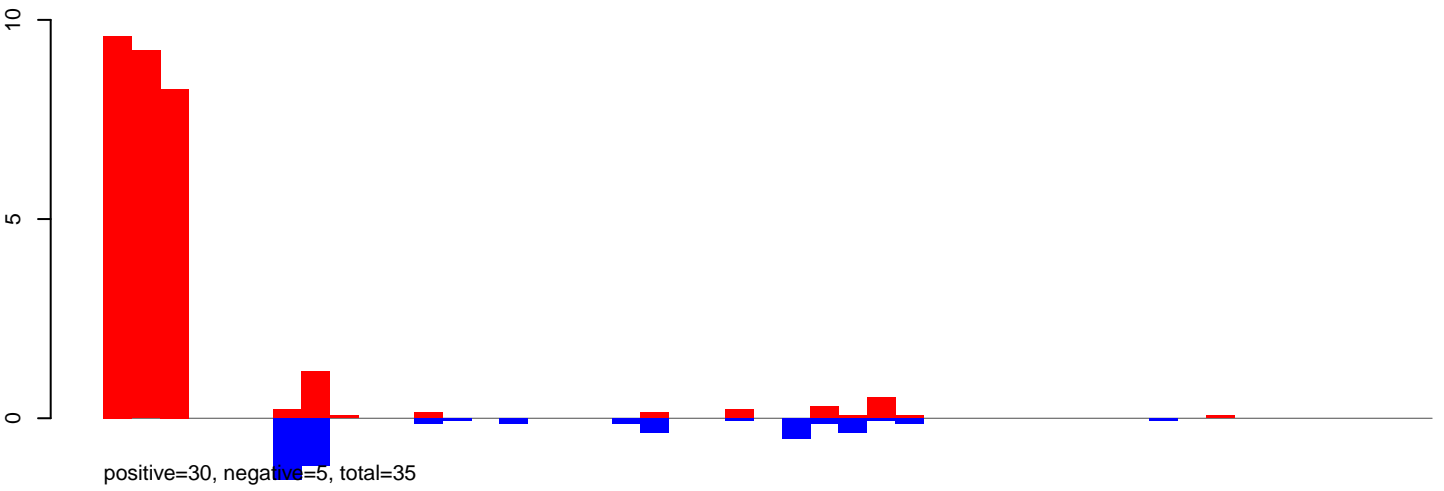
CuQuin_Testes_NL.rep



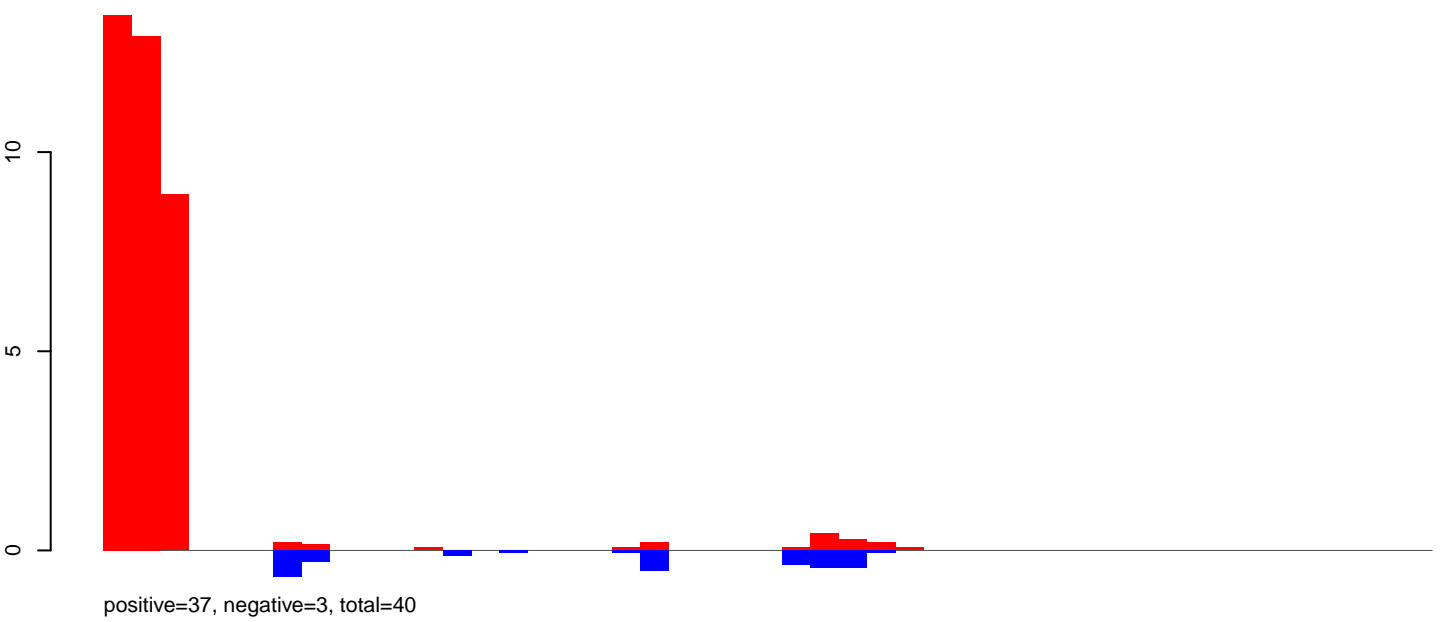
positive=8, negative=8, total=16

Window size=25, length=688, TE@rnd-5_family-471@Unknown@2355:1-688

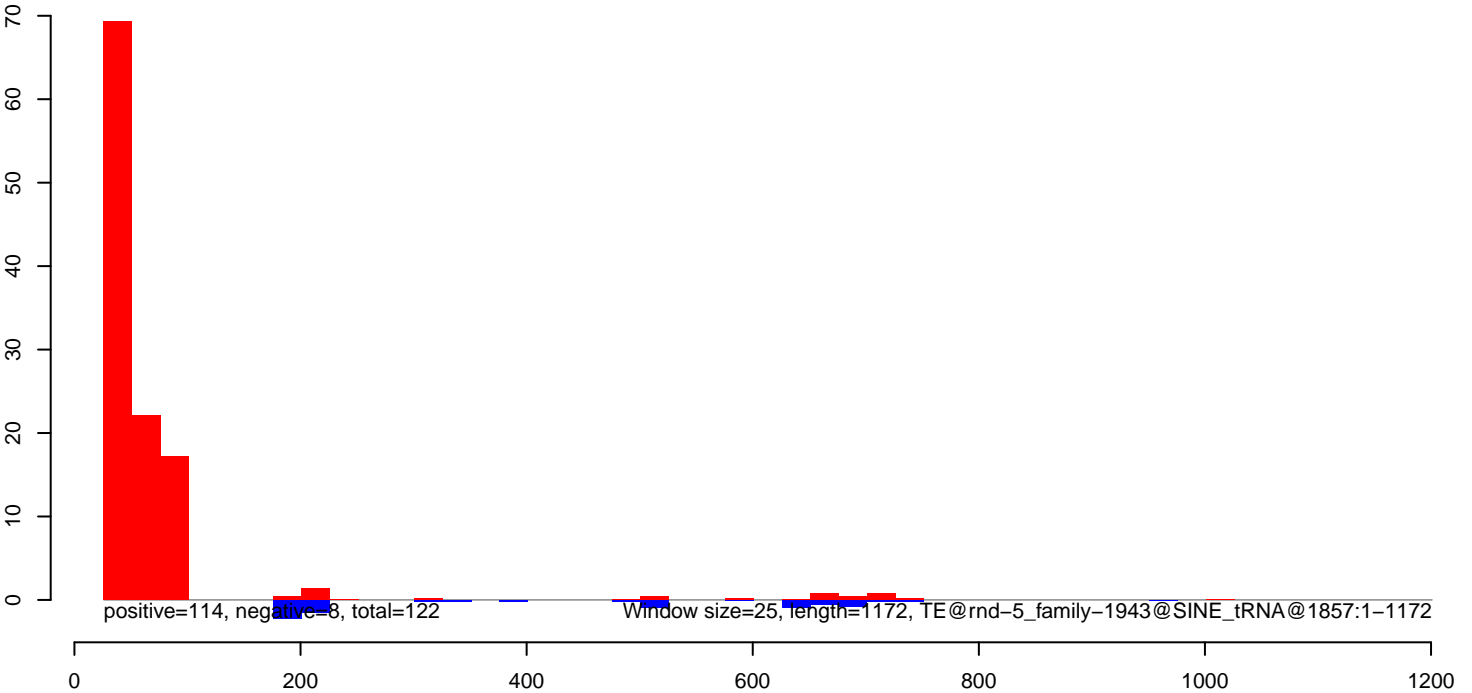
CuQuin_Testes_NL.18_23.rep



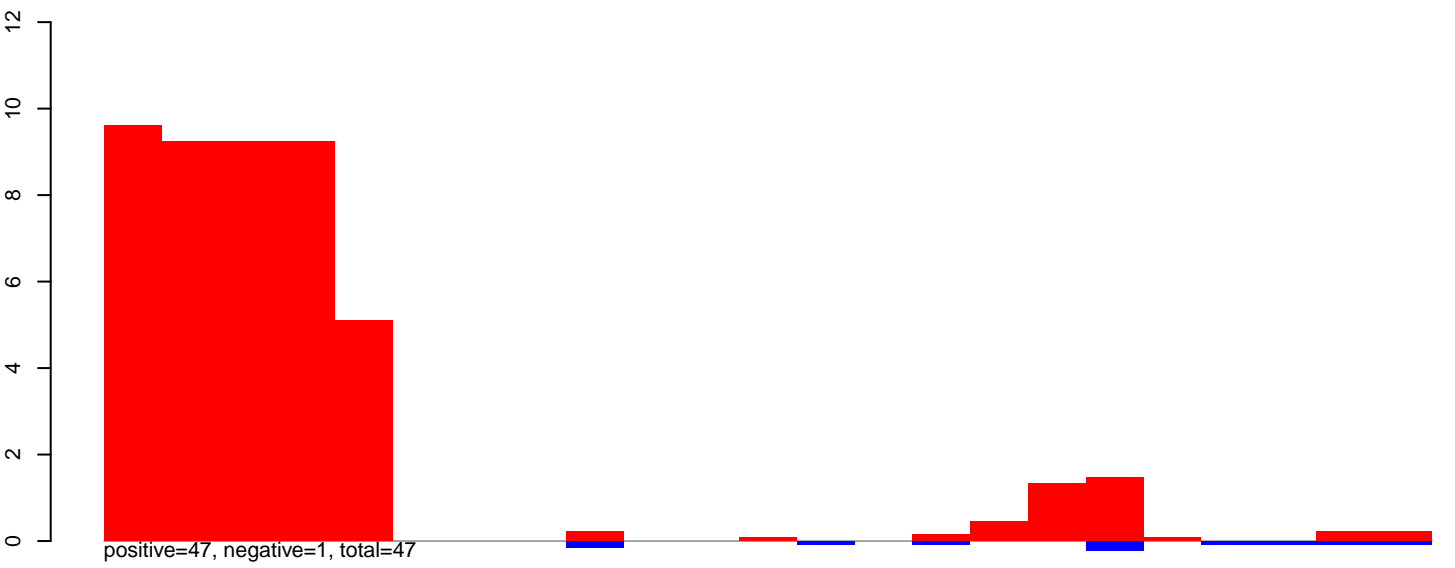
CuQuin_Testes_NL.24_35.rep



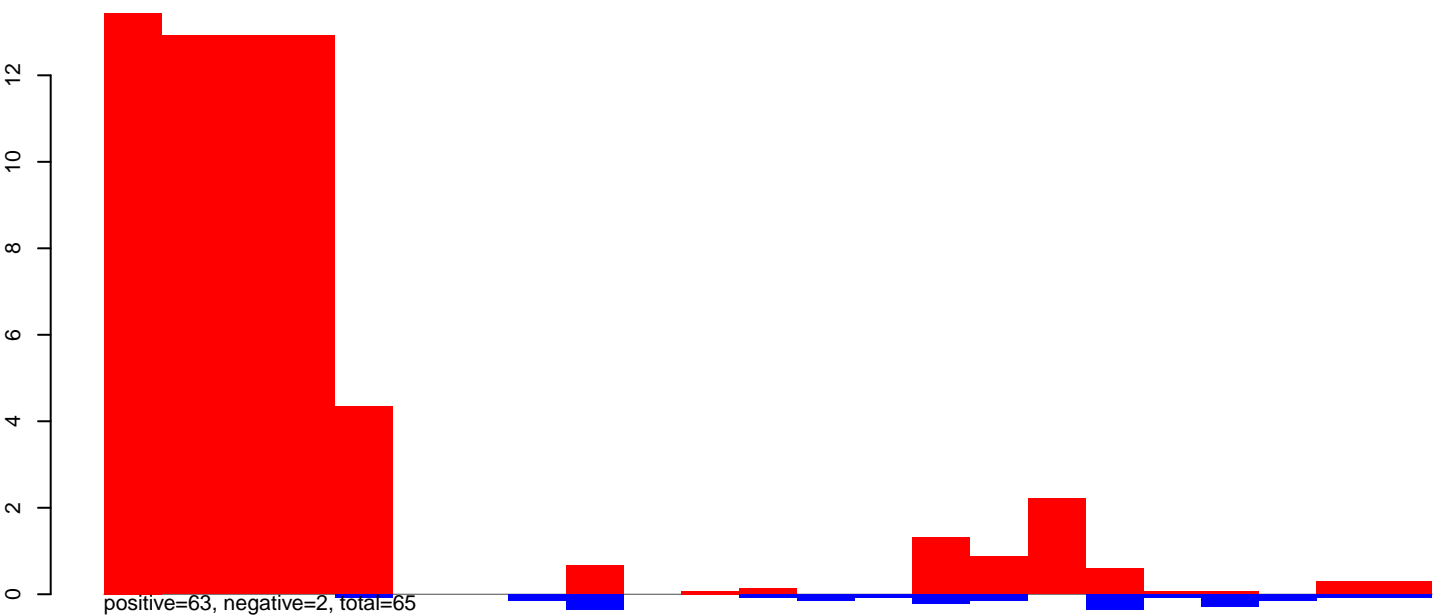
CuQuin_Testes_NL.rep



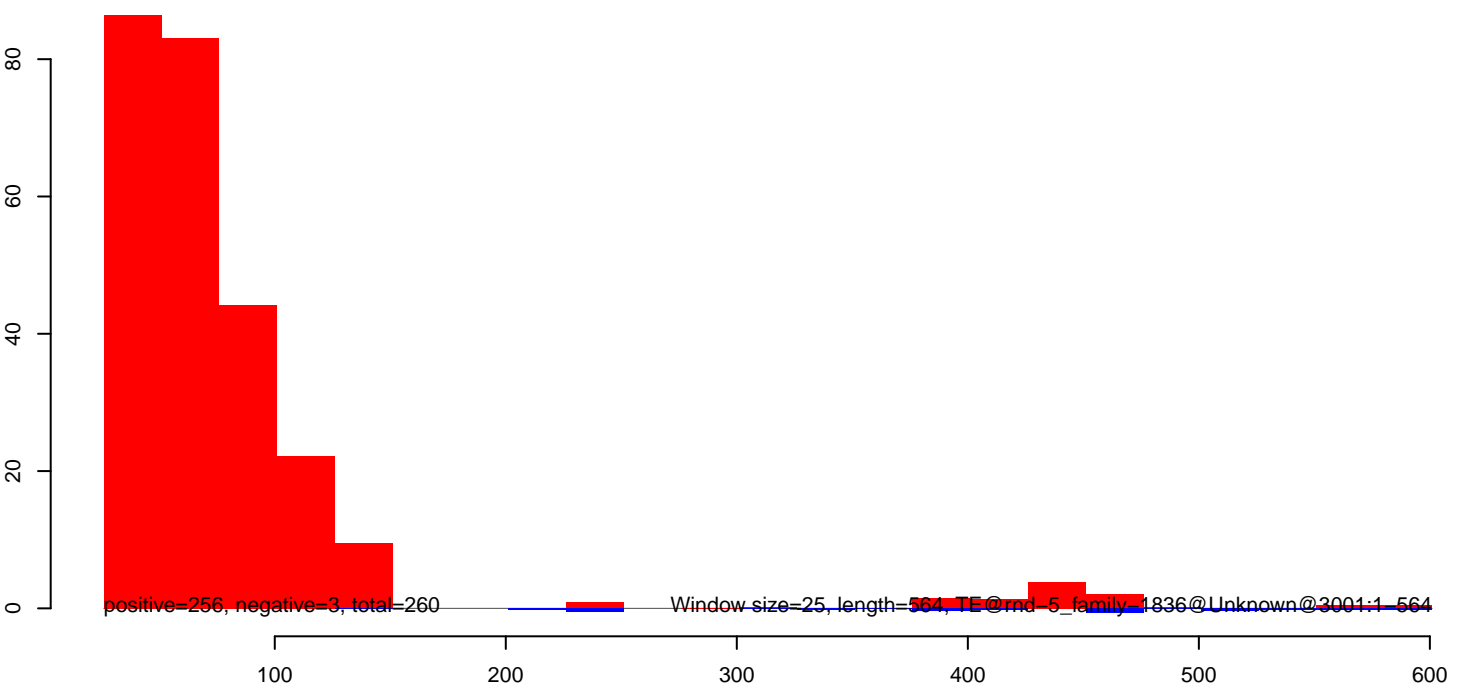
CuQuin_Testes_NL.18_23.rep



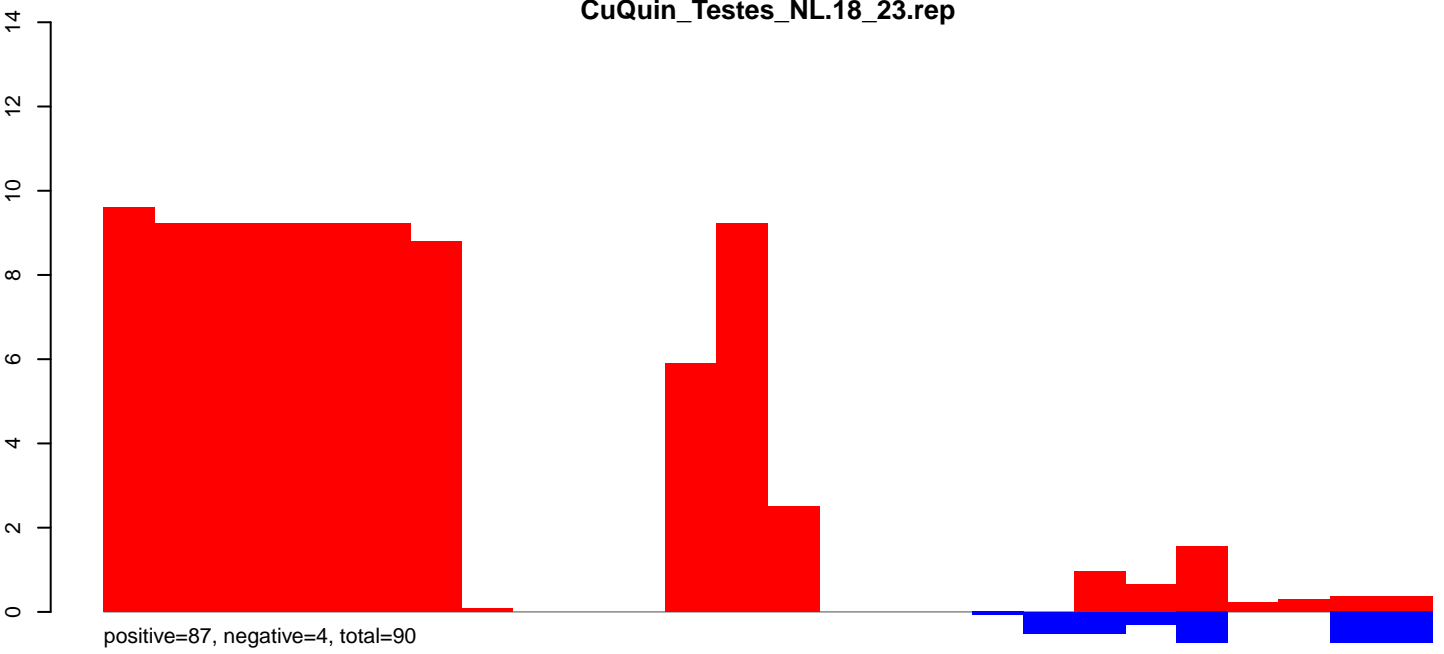
CuQuin_Testes_NL.24_35.rep



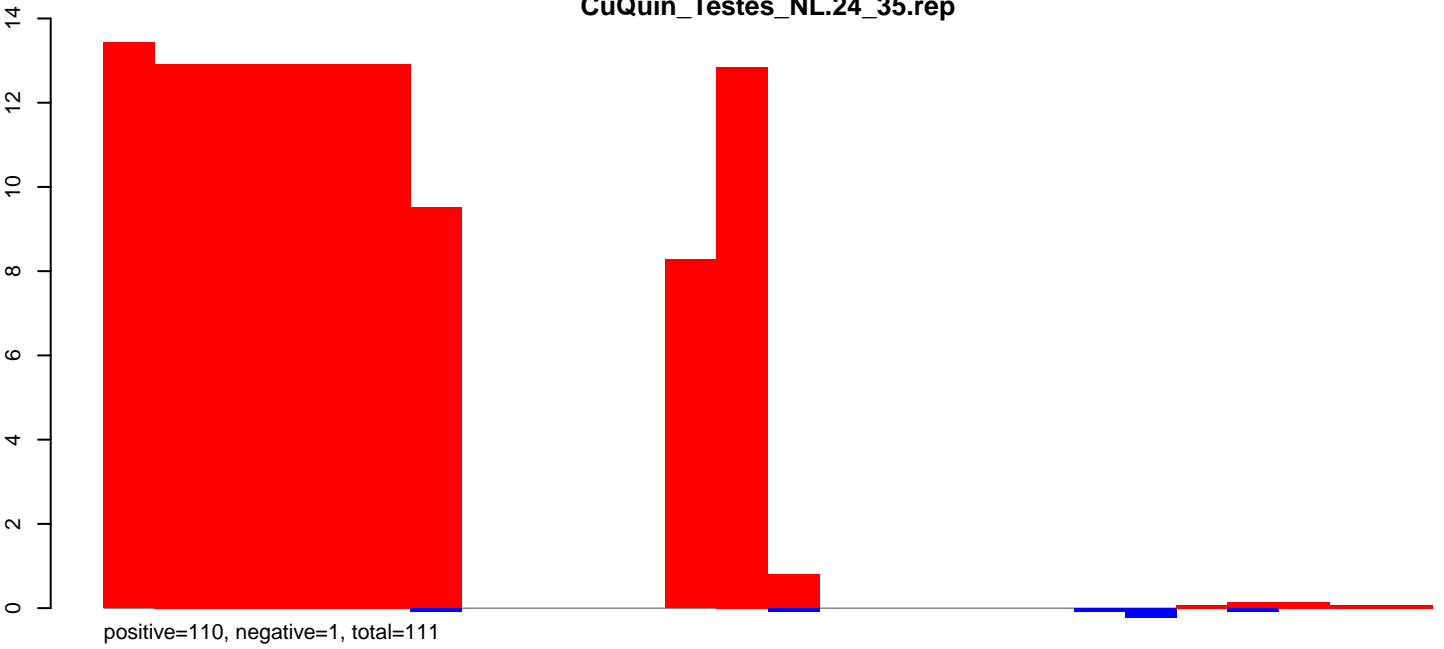
CuQuin_Testes_NL.rep



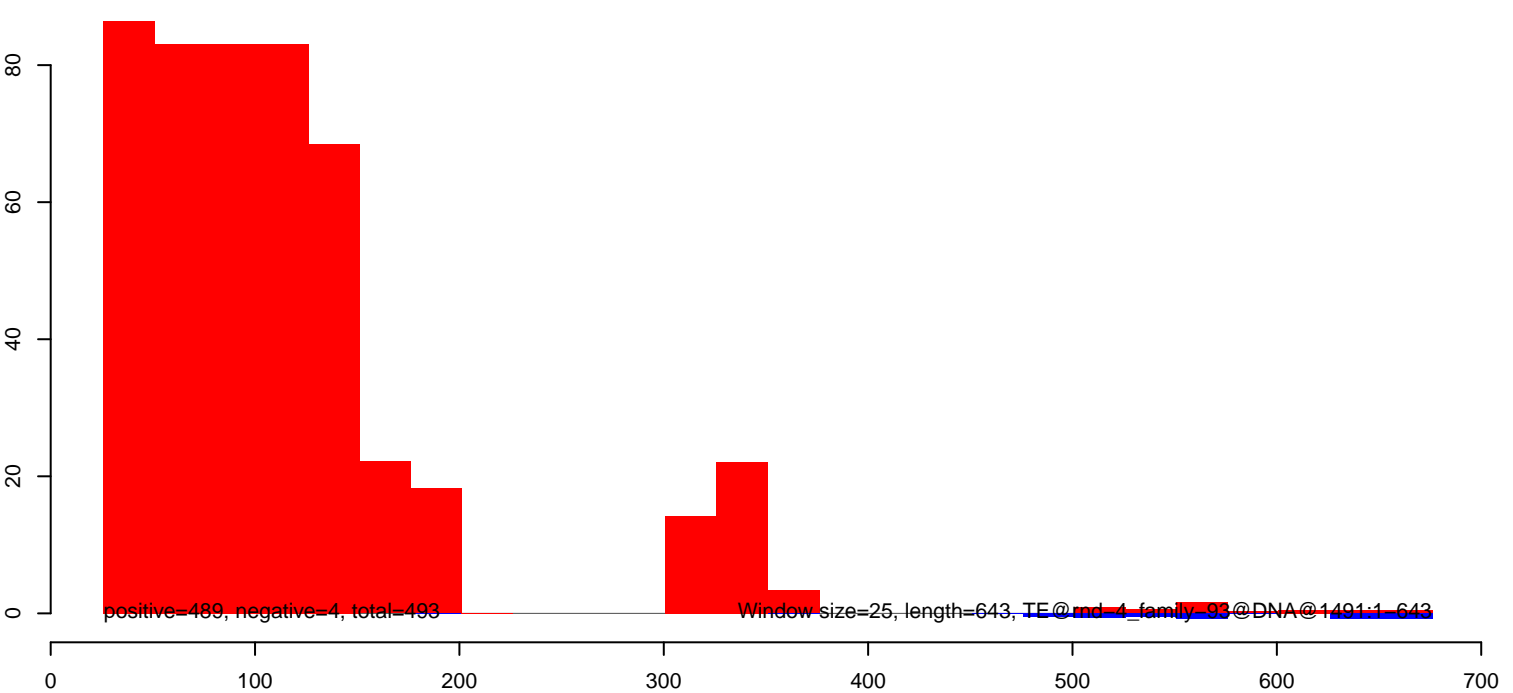
CuQuin_Testes_NL.18_23.rep



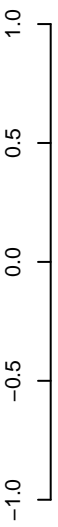
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

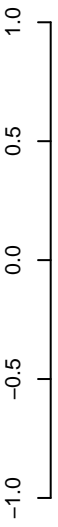


CuQuin_Testes_NL.18_23.rep



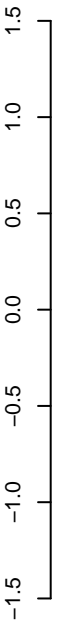
positive=2, negative=2, total=4

CuQuin_Testes_NL.24_35.rep



positive=6, negative=5, total=11

CuQuin_Testes_NL.rep

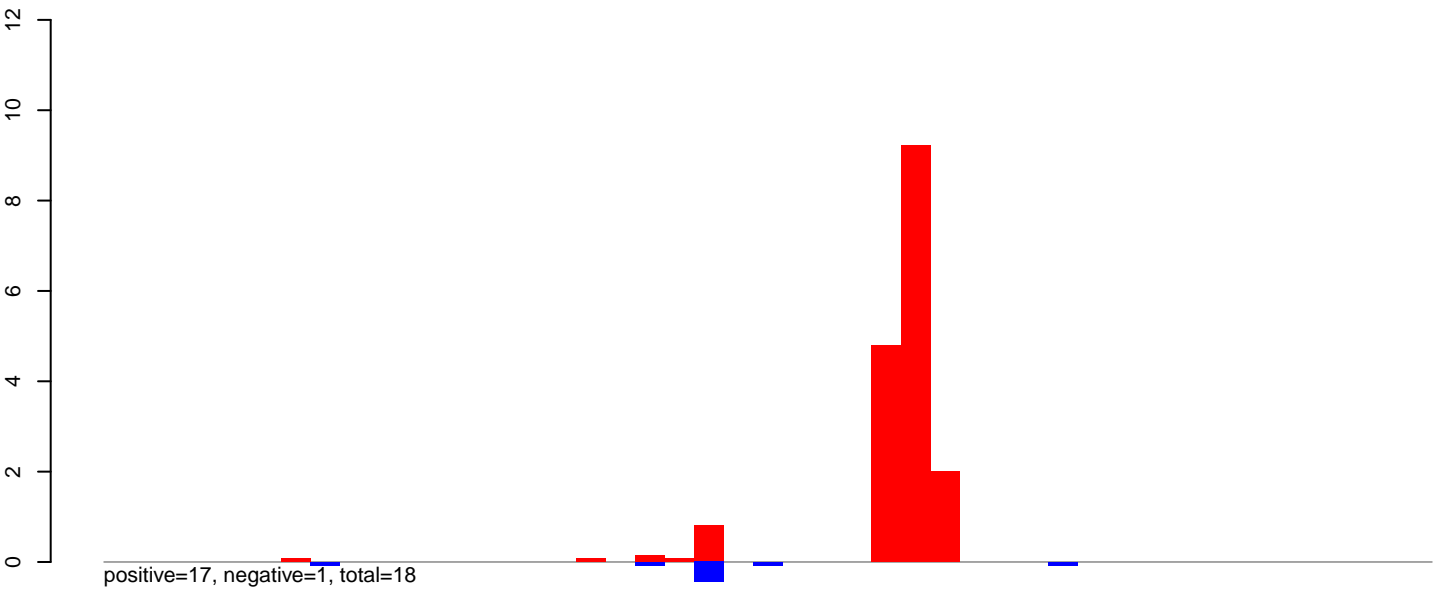


positive=8, negative=7, total=14

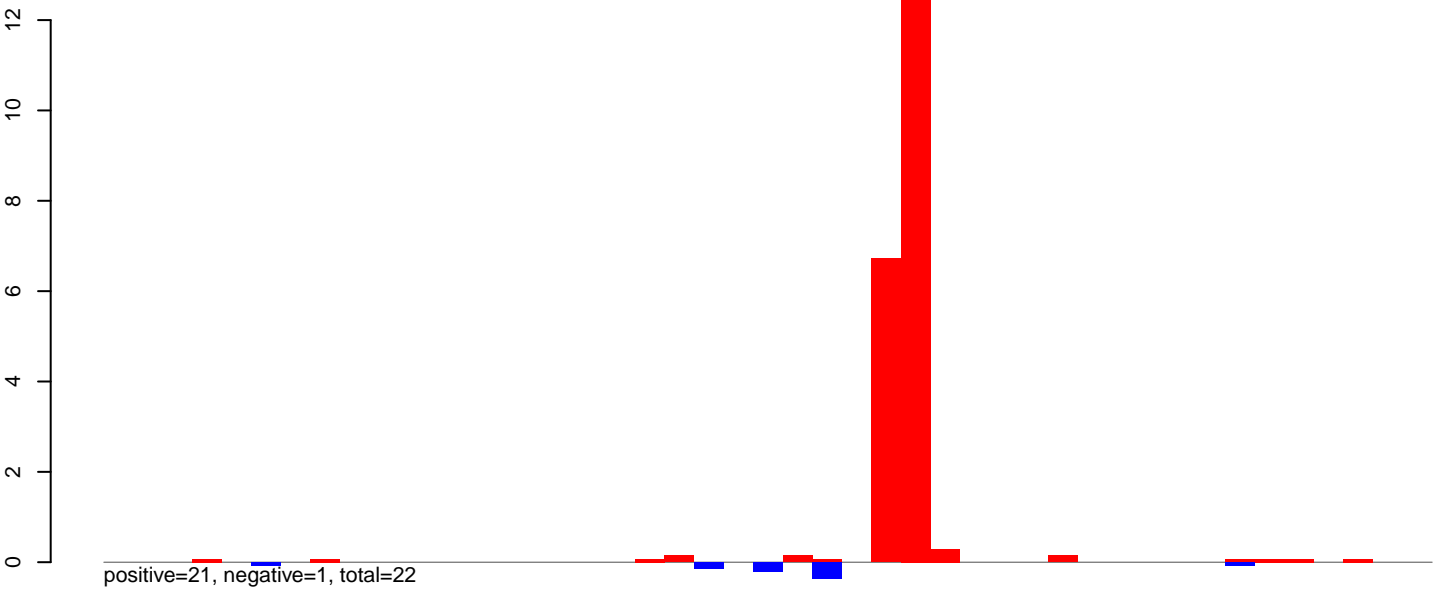
Window size=25, length=669, TE@rnd-4_family-696@DNA@2157:1-669



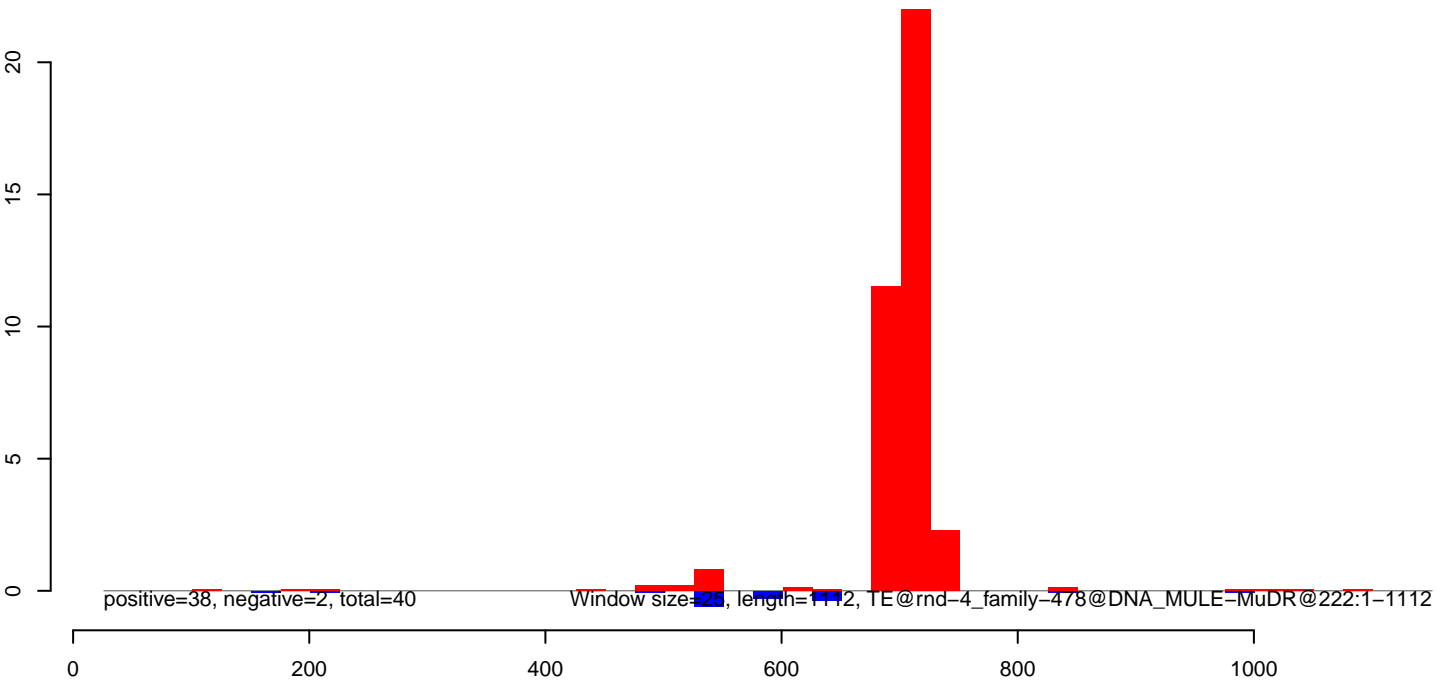
CuQuin_Testes_NL.18_23.rep



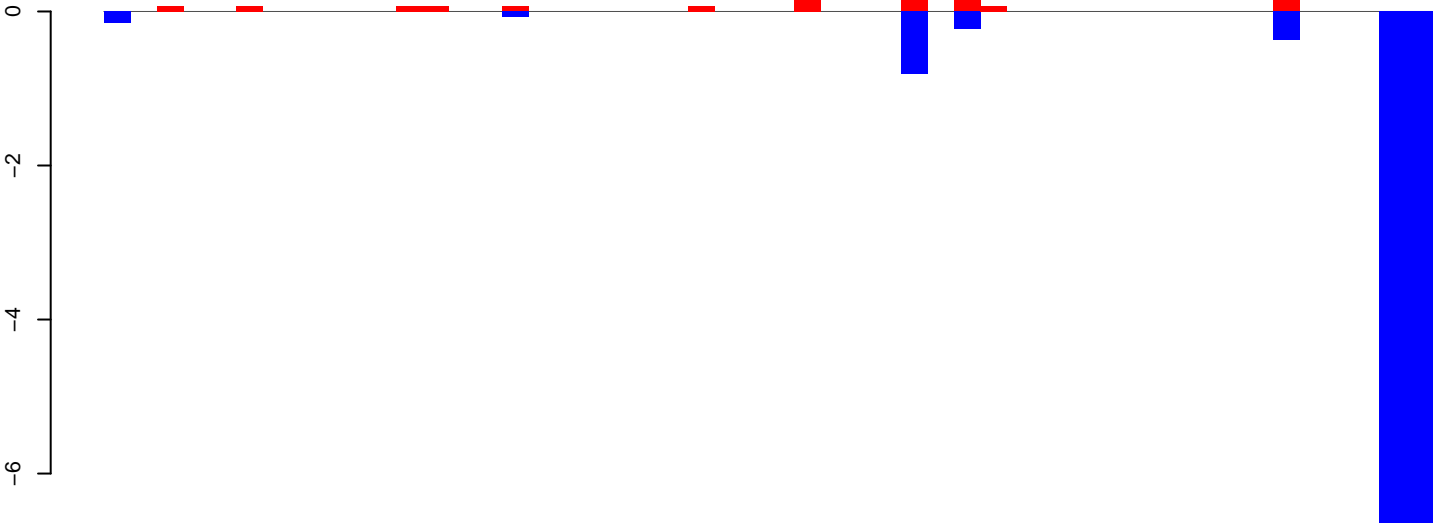
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

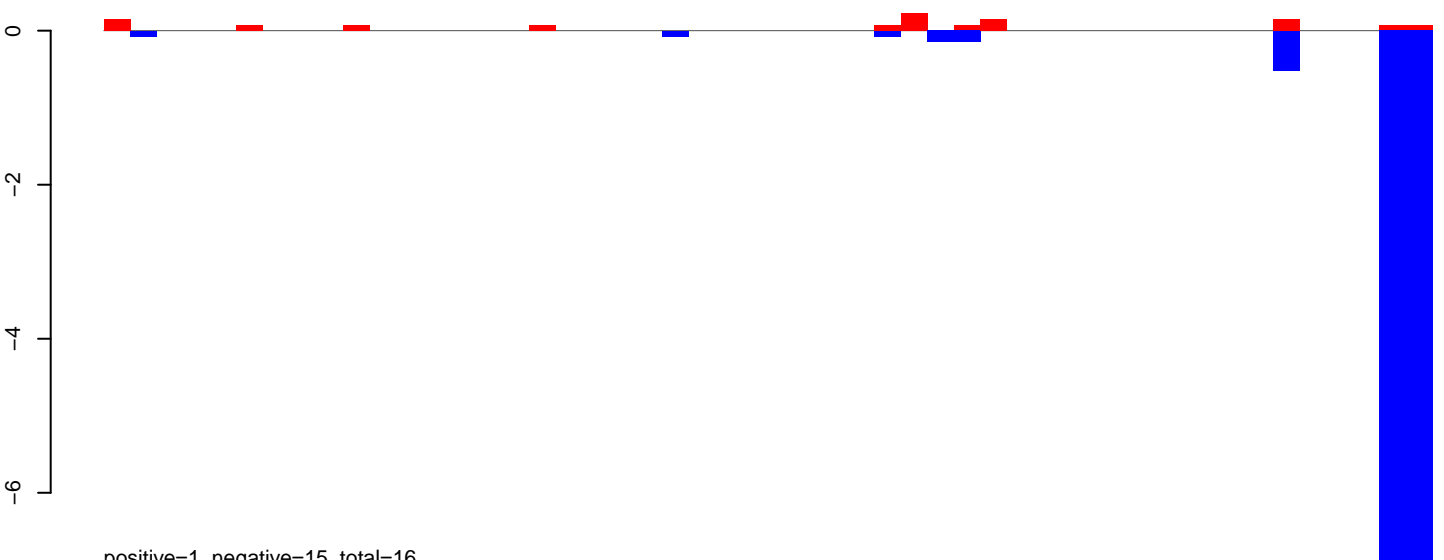


CuQuin_Testes_NL.18_23.rep



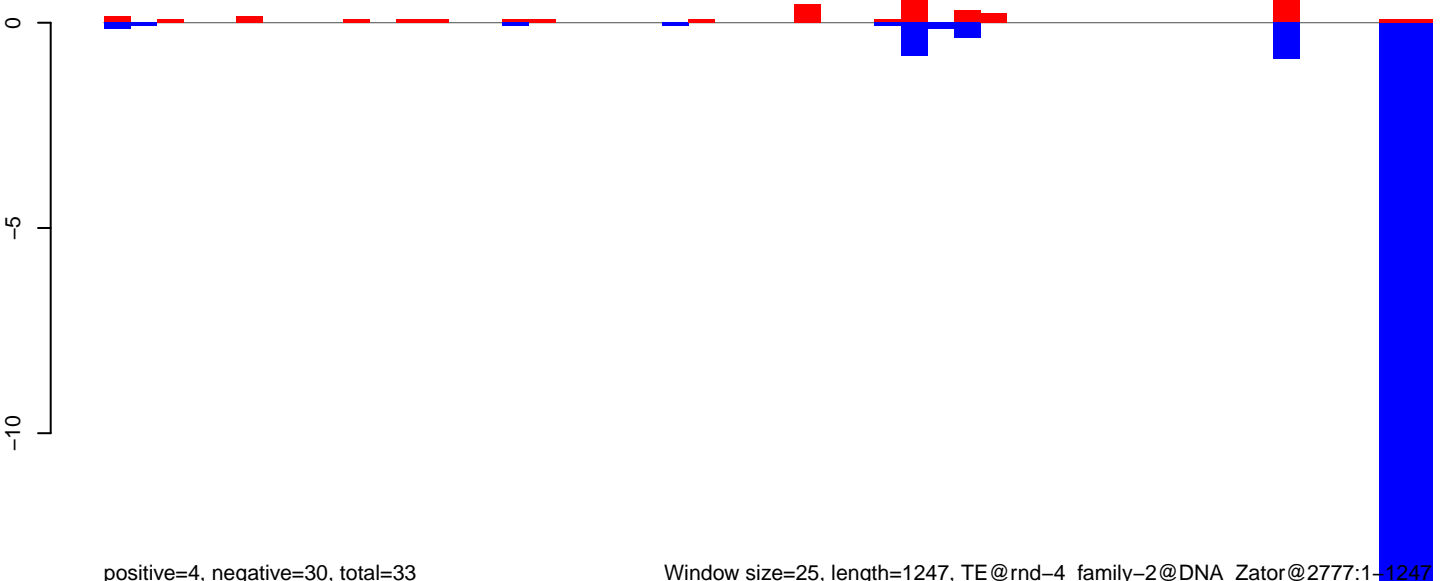
positive=2, negative=15, total=17

CuQuin_Testes_NL.24_35.rep



positive=1, negative=15, total=16

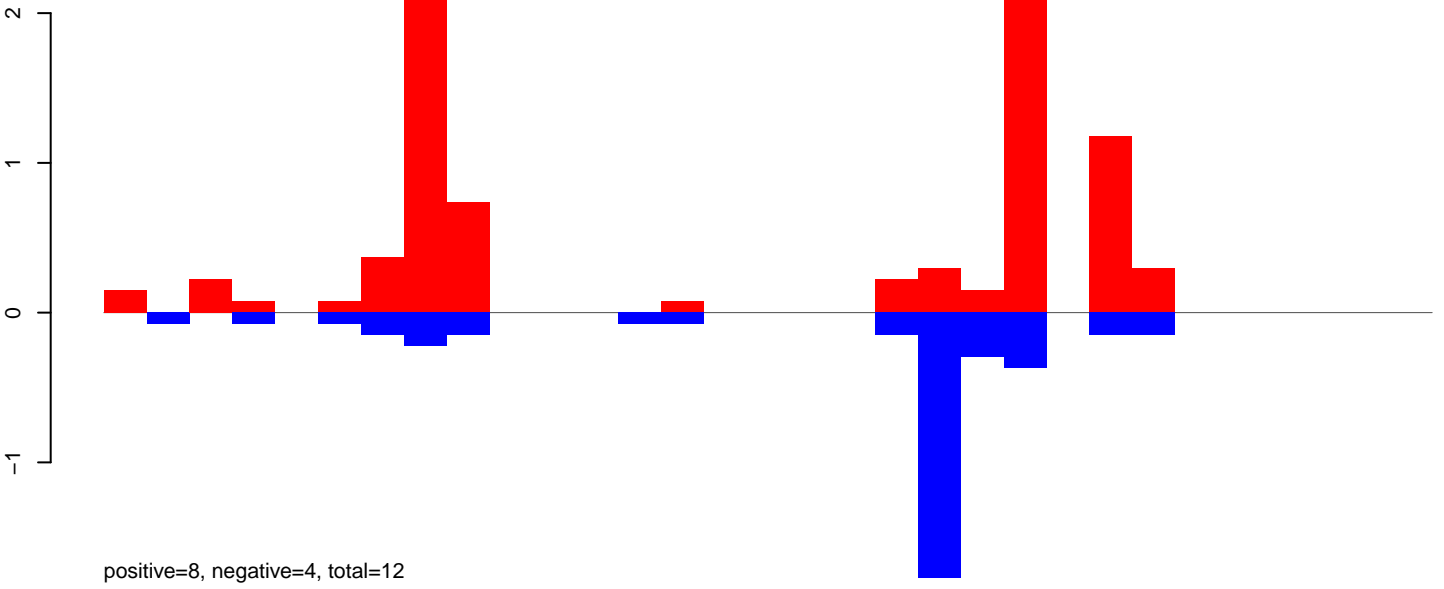
CuQuin_Testes_NL.rep



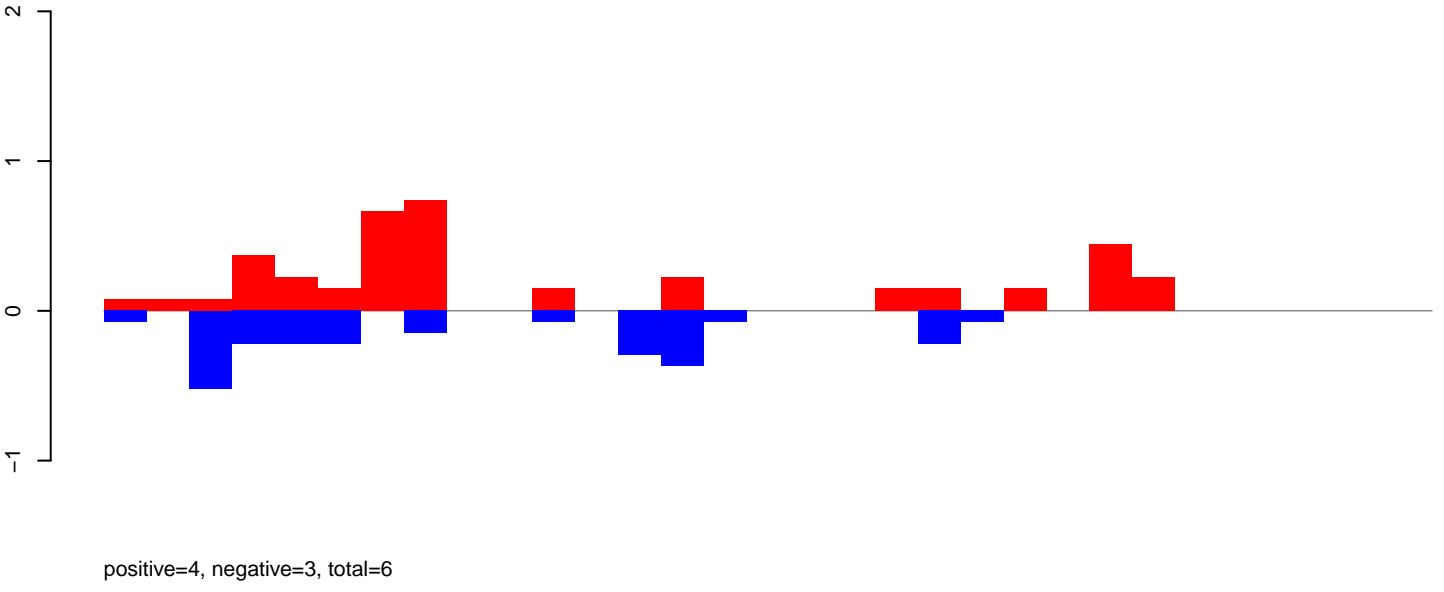
positive=4, negative=30, total=33

Window size=25, length=1247, TE@rnd-4_family-2@DNA_Zator@2777:1-1247

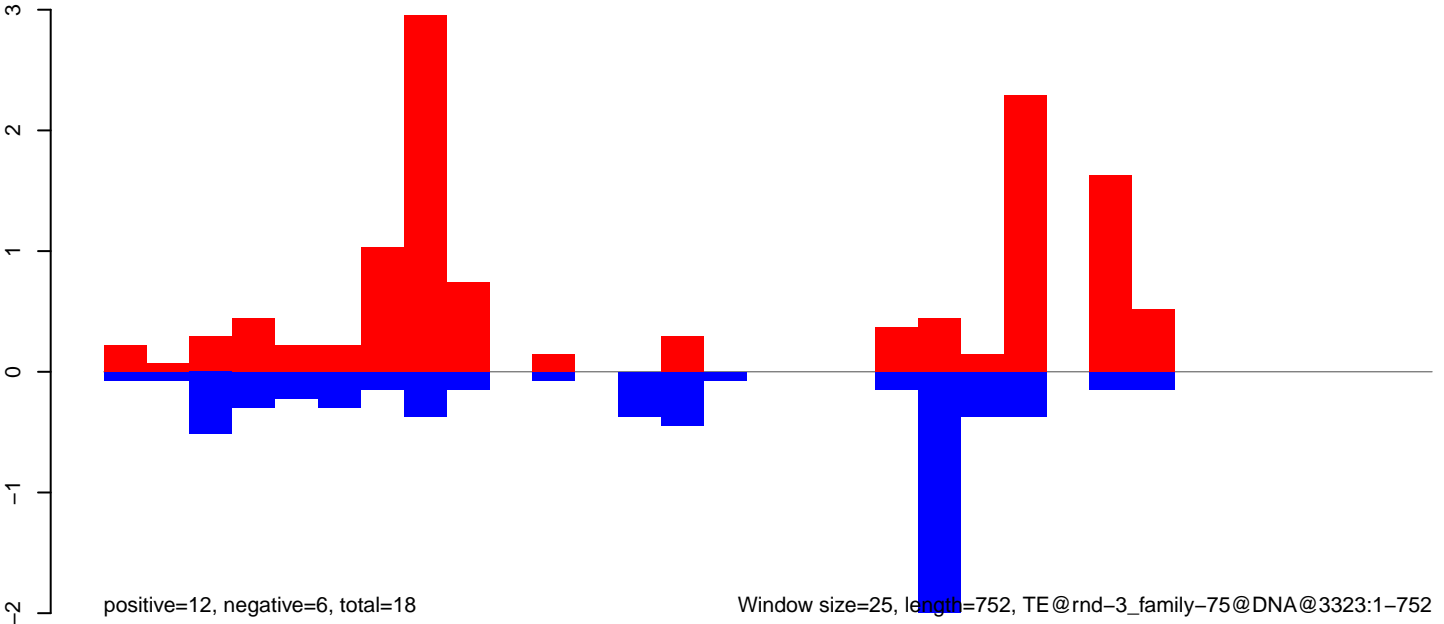
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

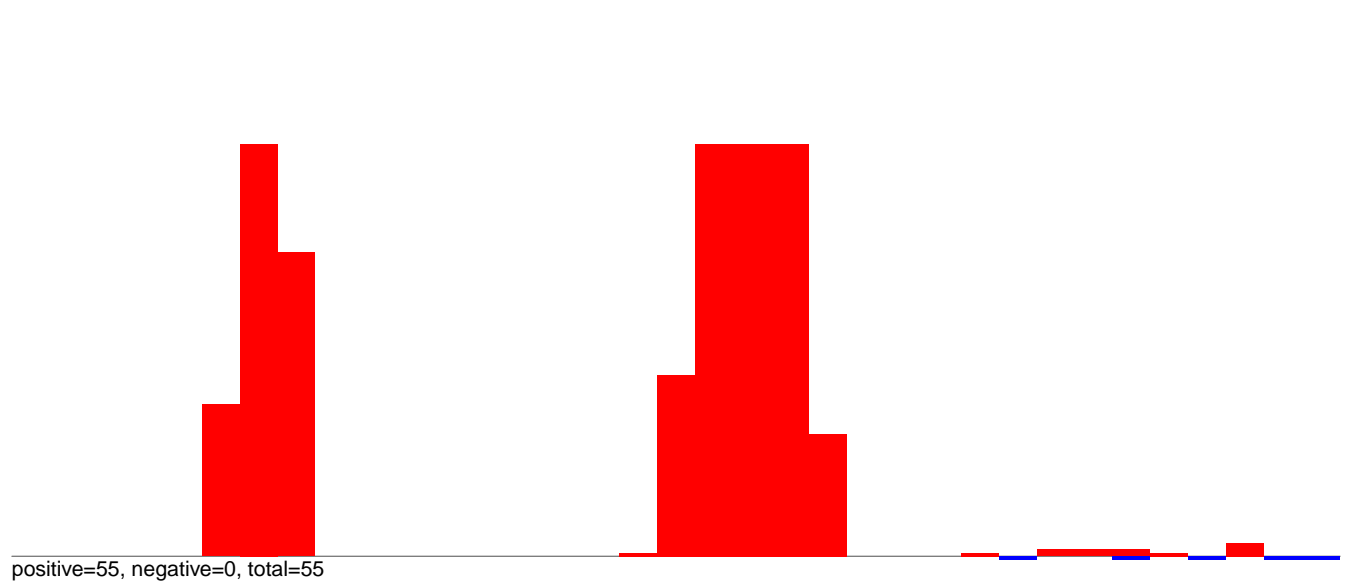


CuQuin_Testes_NL.rep

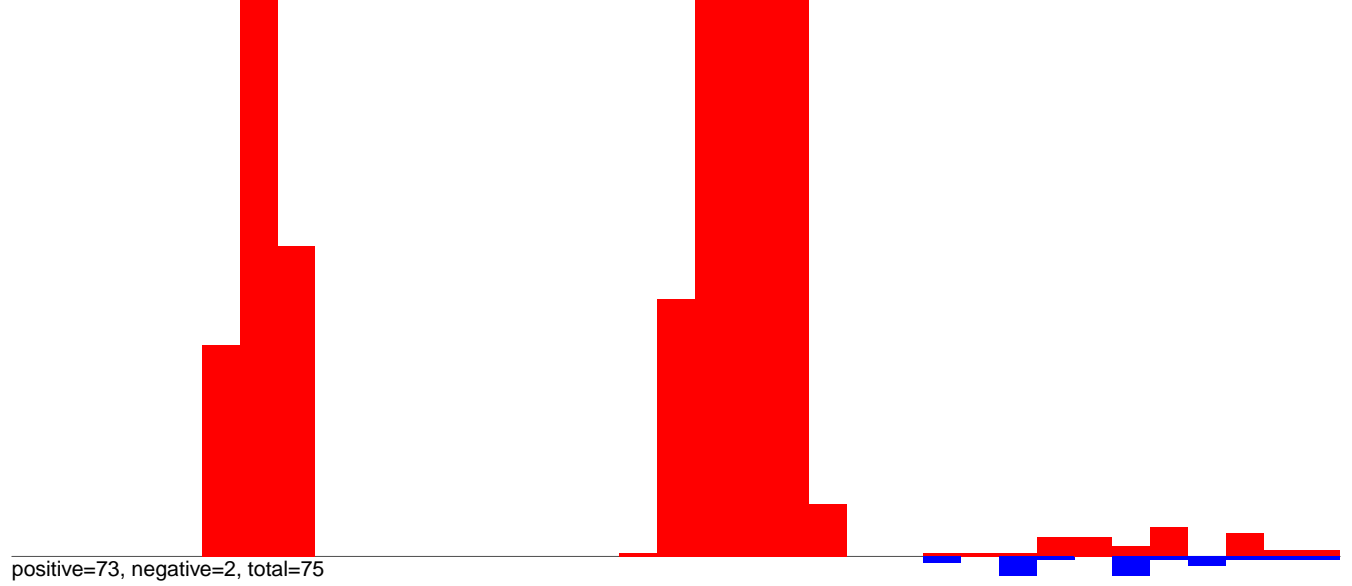


Window size=25, length=752, TE@rnd-3_family-75@DNA@3323:1-752

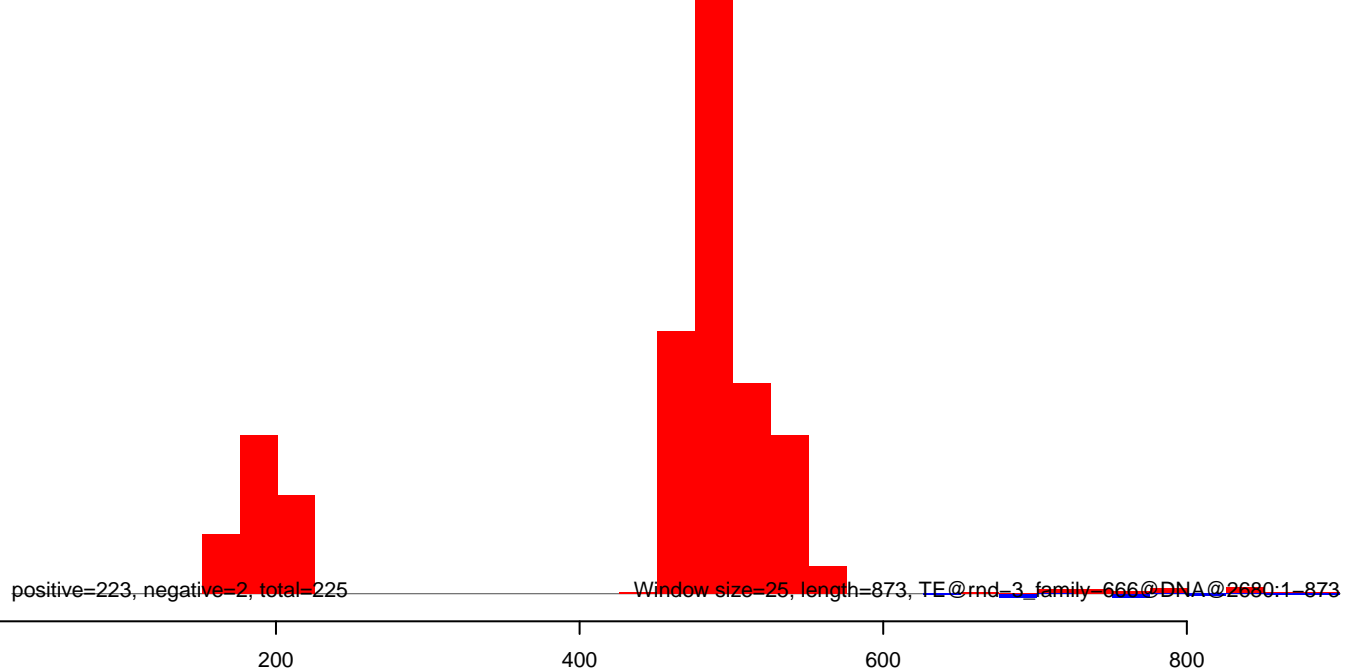
CuQuin_Testes_NL.18_23.rep



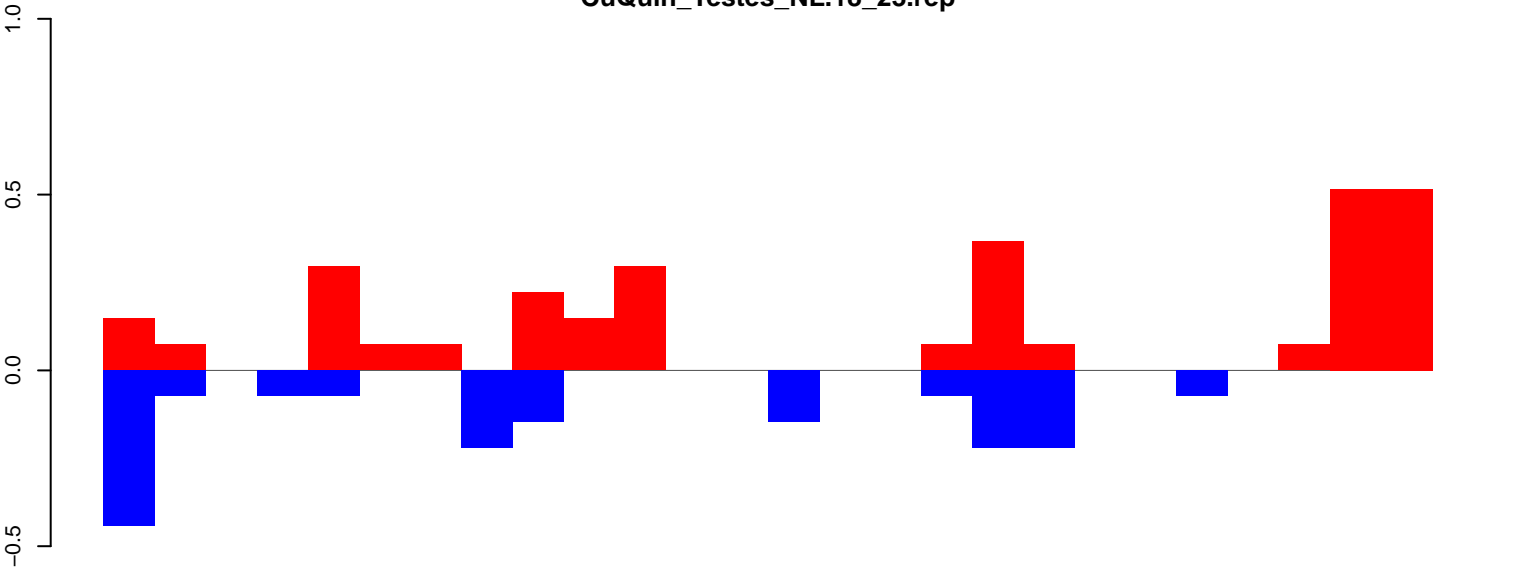
CuQuin_Testes_NL.24_35.rep



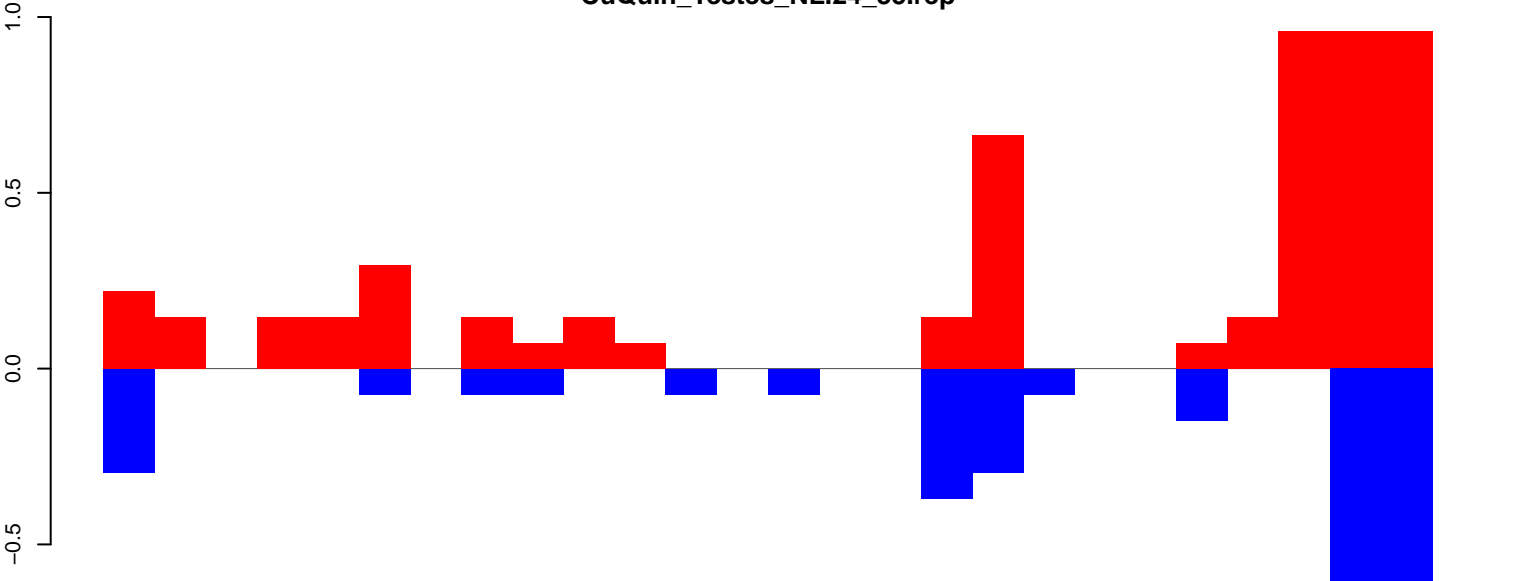
CuQuin_Testes_NL.rep



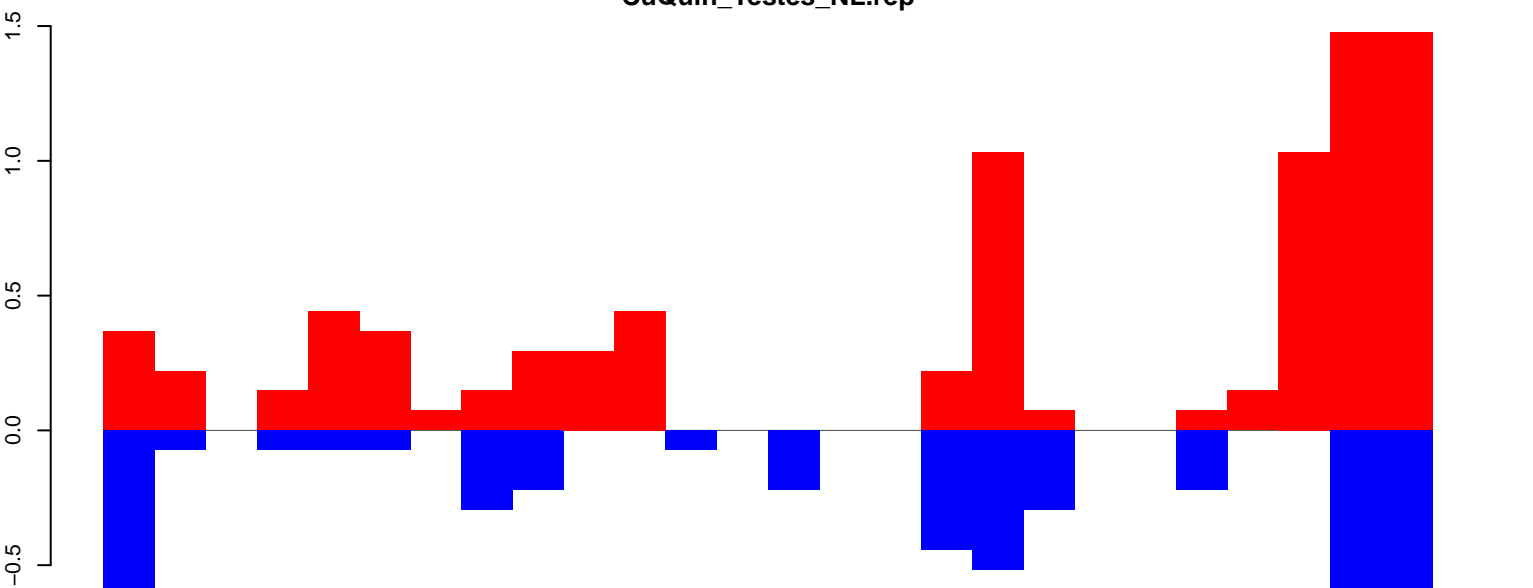
CuQuin_Testes_NL.18_23.rep



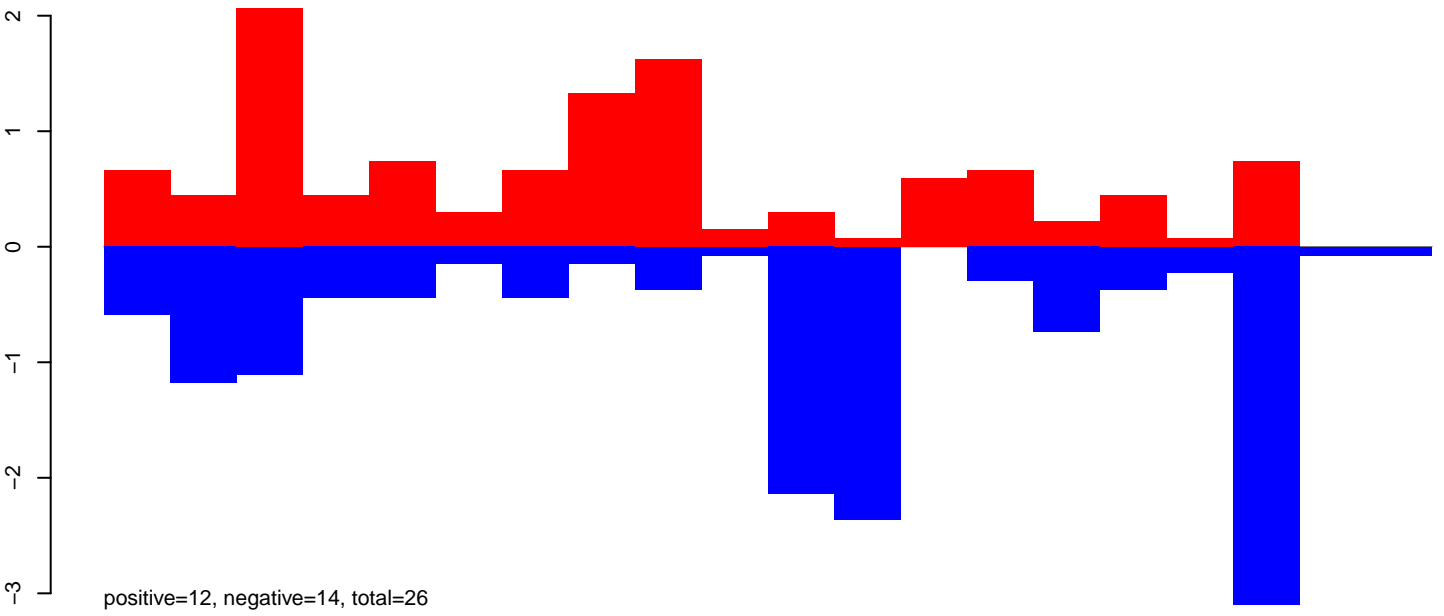
CuQuin_Testes_NL.24_35.rep



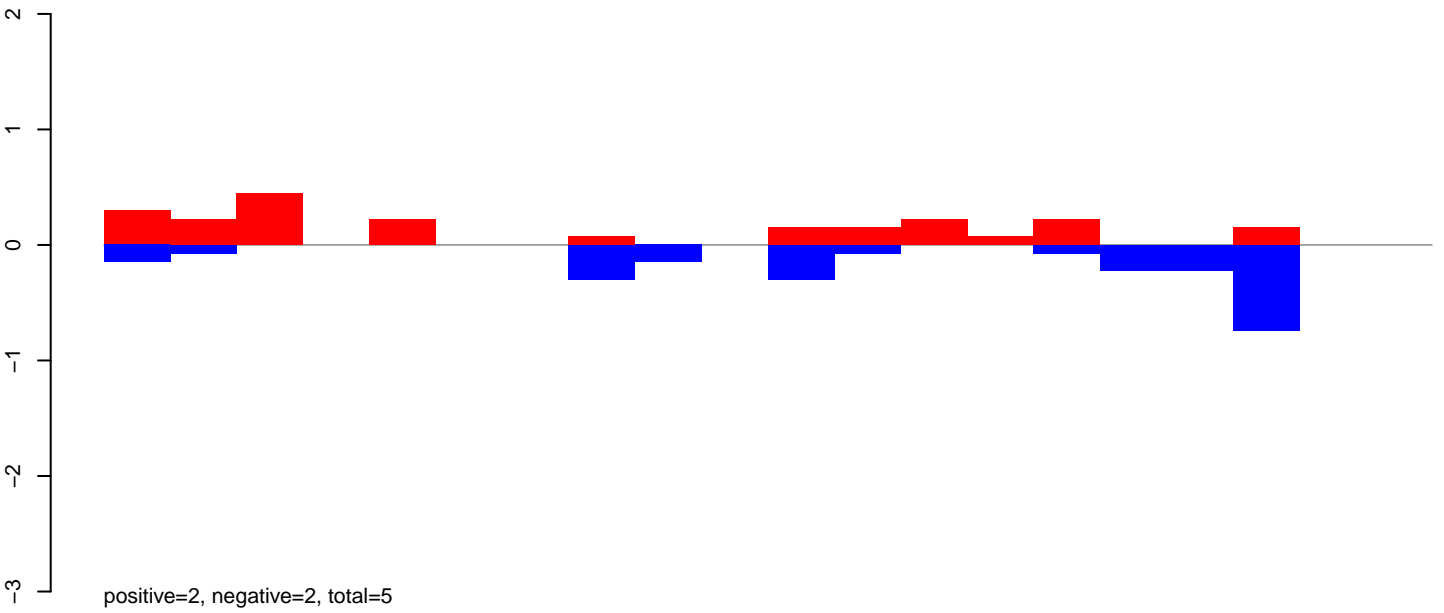
CuQuin_Testes_NL.rep



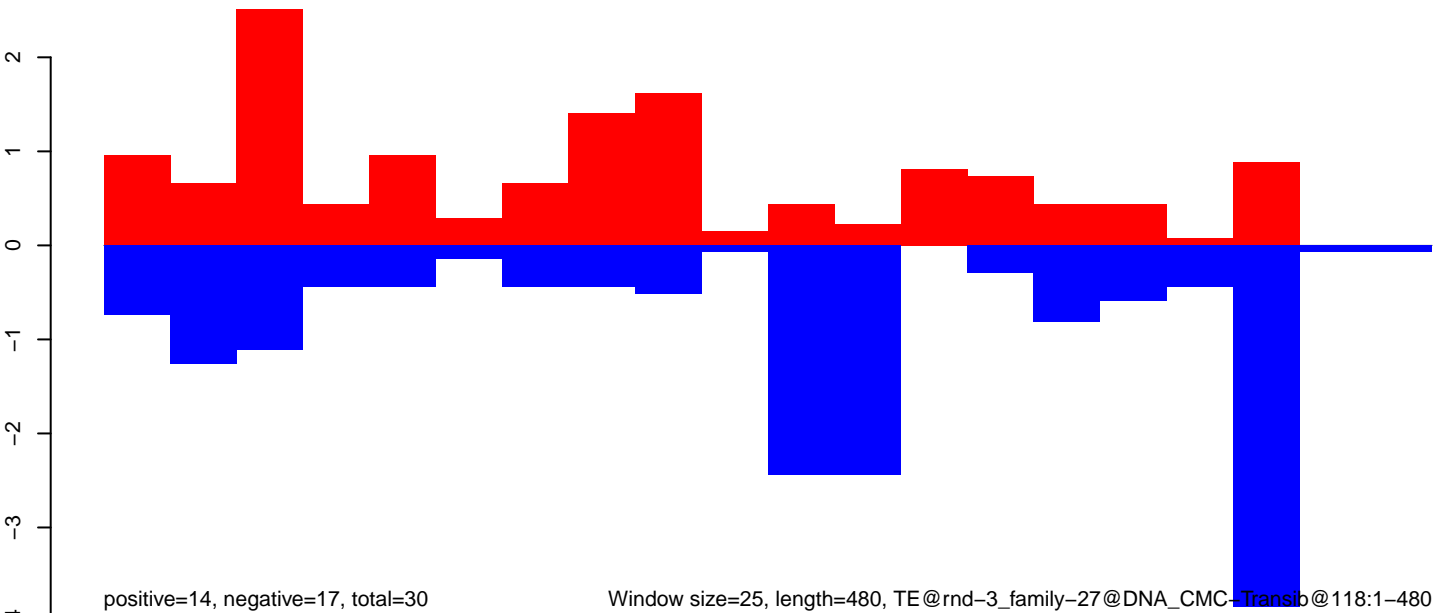
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

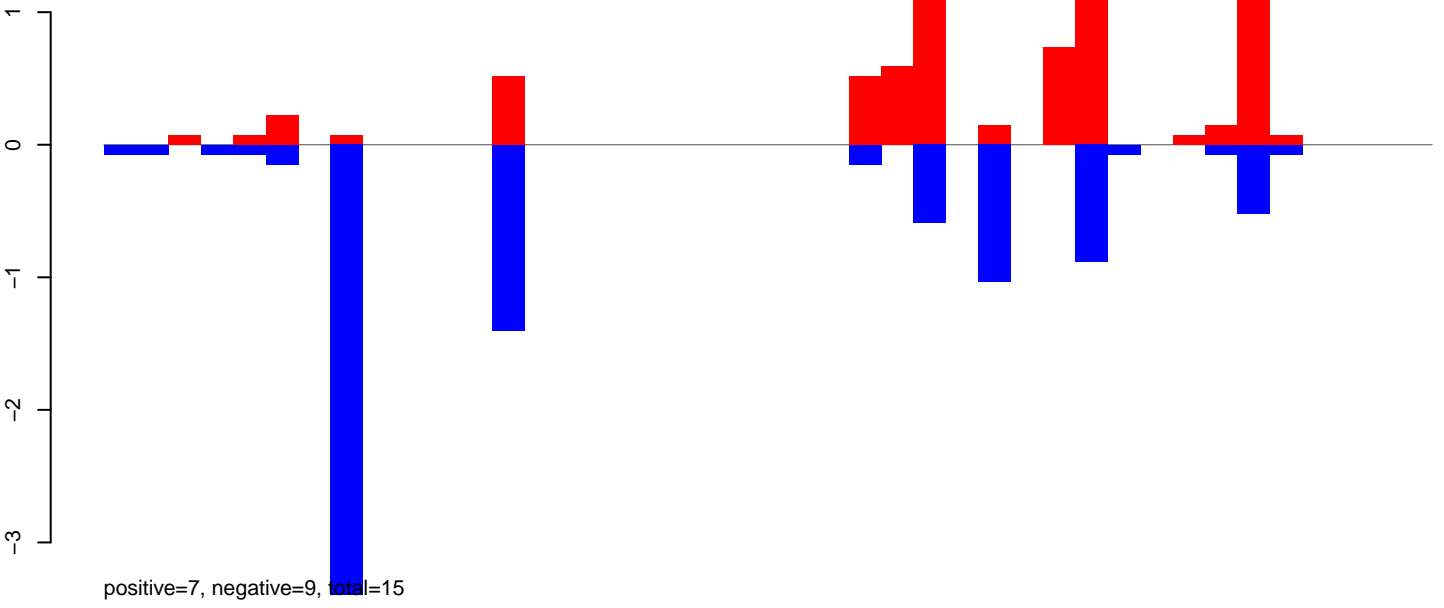


CuQuin_Testes_NL.rep

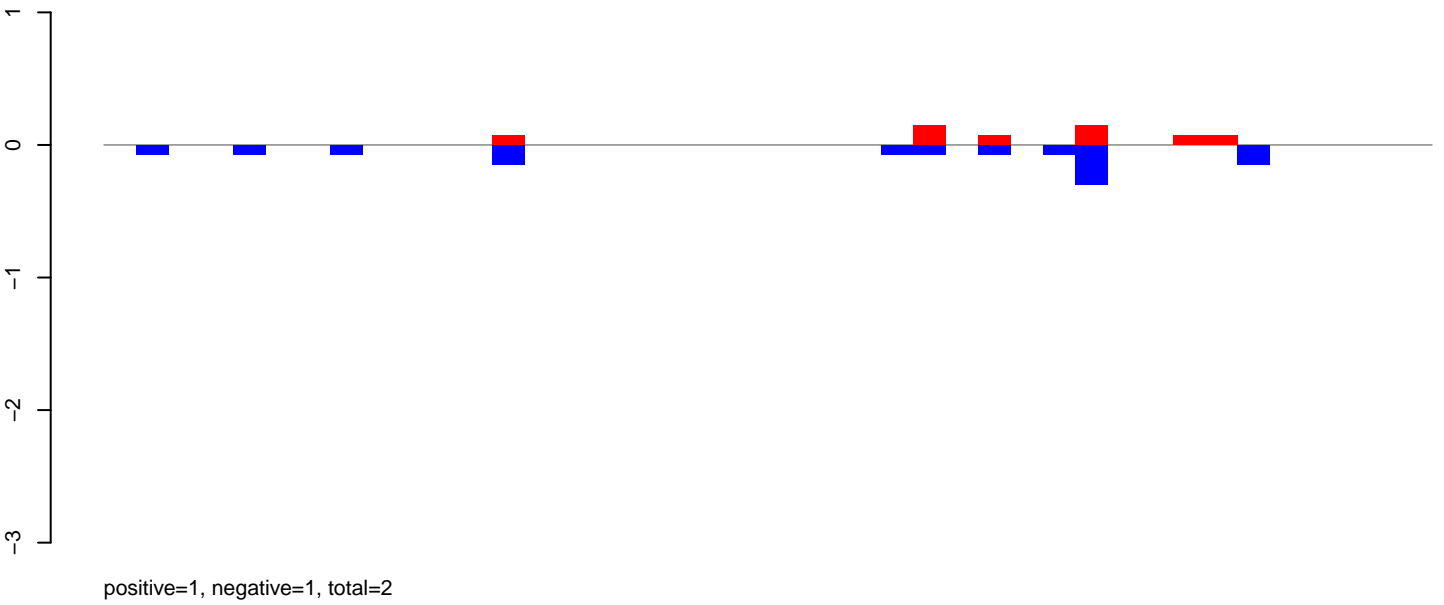


Window size=25, length=480, TE@rnd-3_family-27@DNA_CMC-Transib@118:1-480

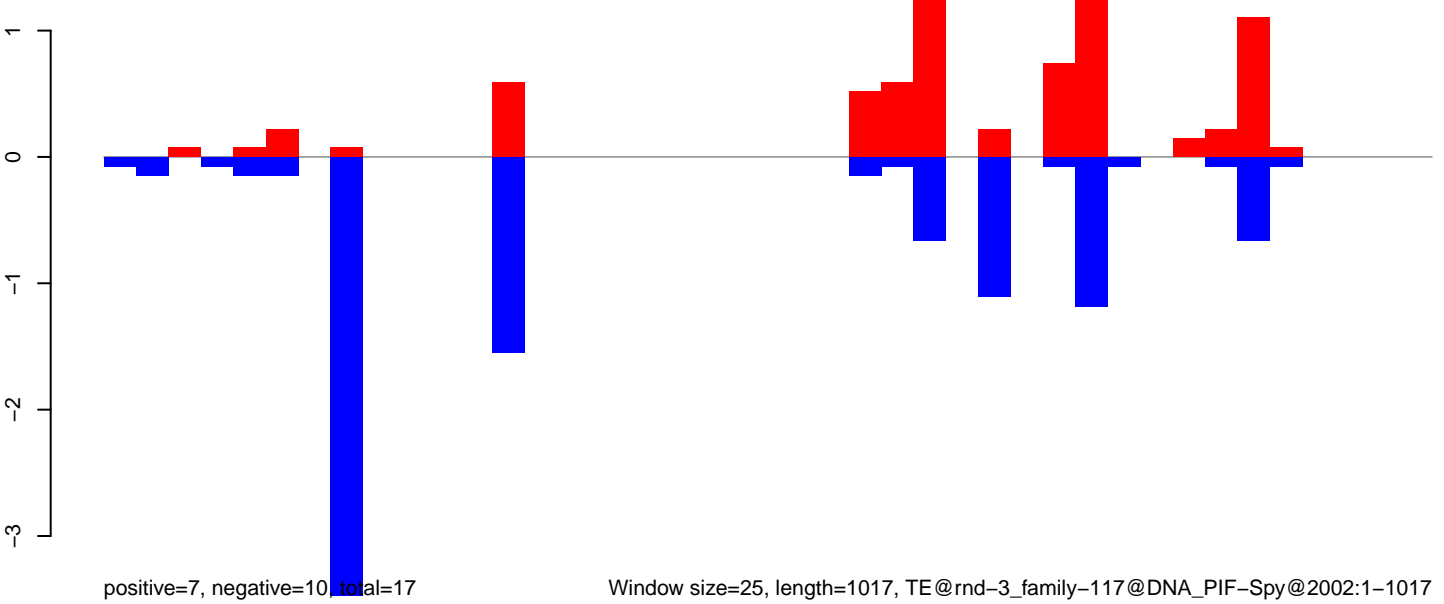
CuQuin_Testes_NL.18_23.rep



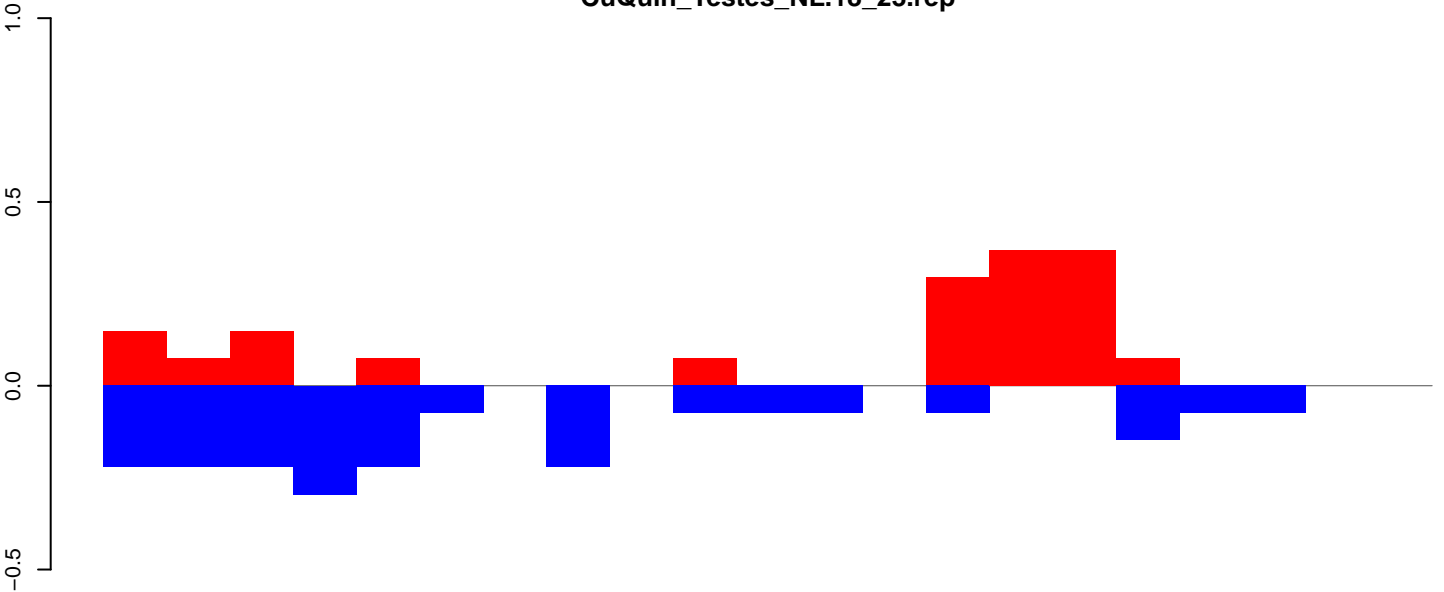
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

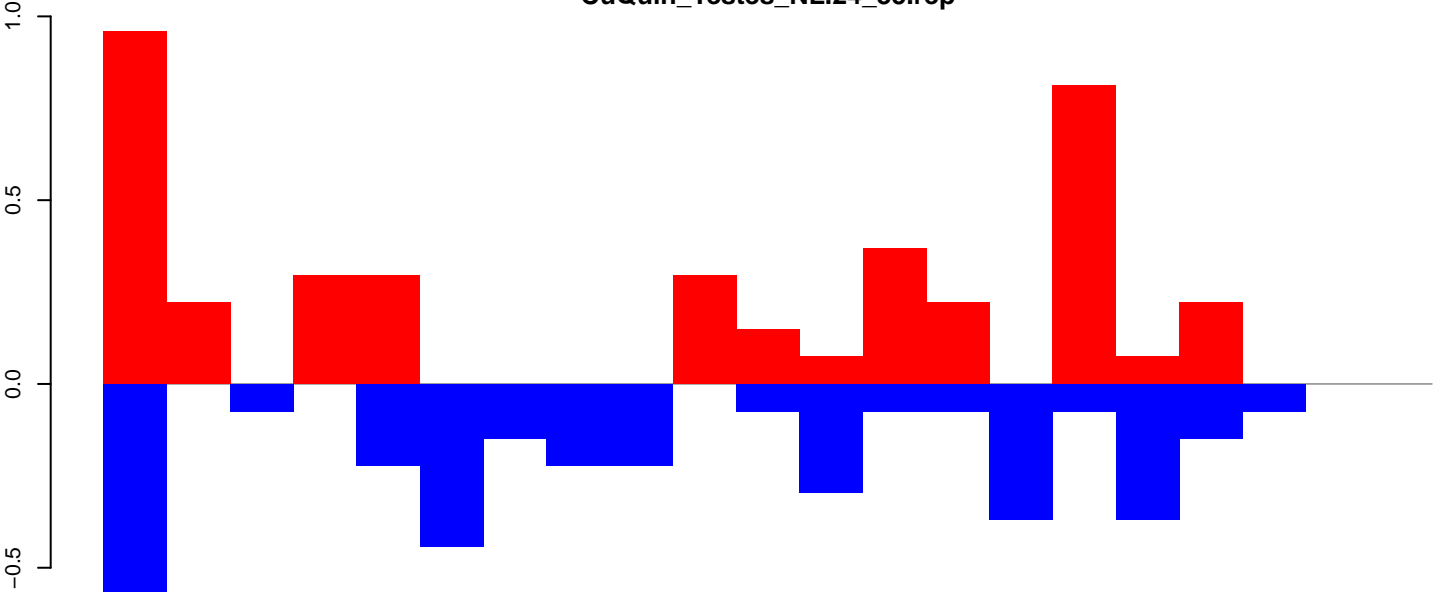


CuQuin_Testes_NL.18_23.rep



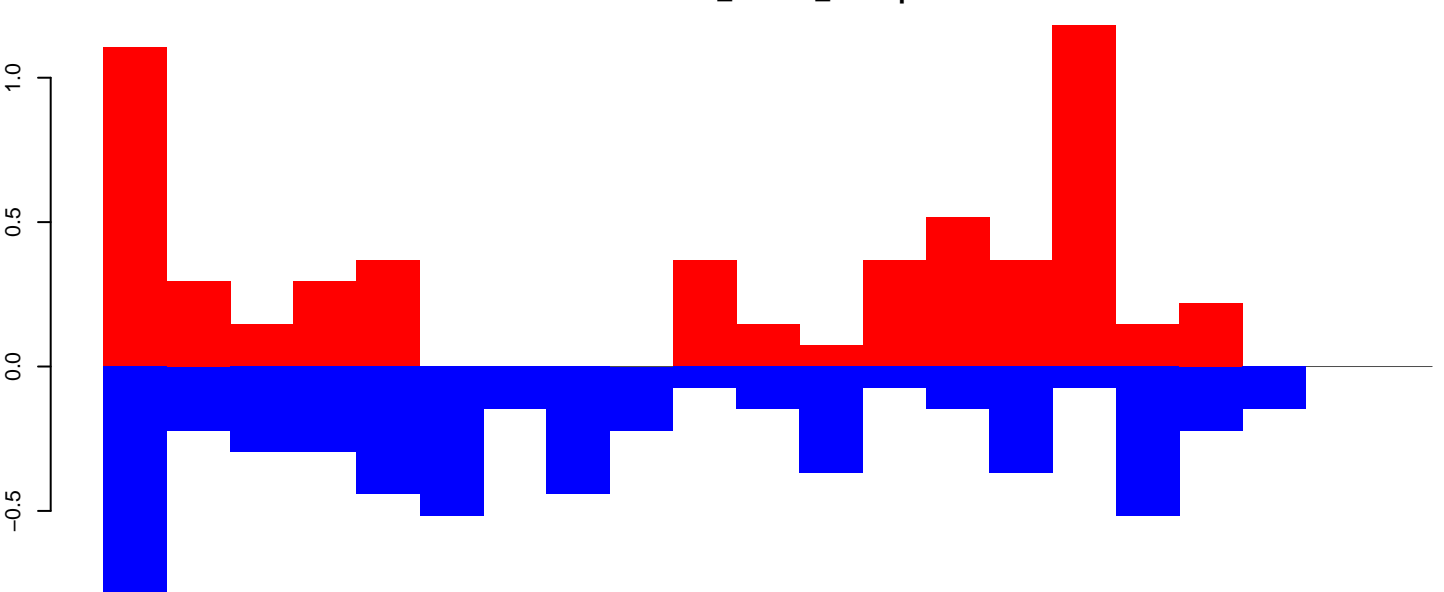
positive=2, negative=2, total=4

CuQuin_Testes_NL.24_35.rep



positive=4, negative=4, total=8

CuQuin_Testes_NL.rep



positive=6, negative=6, total=11

Window size=25, length=515, TE@rnd-3_family-101@Unknown@2687:1-515

100

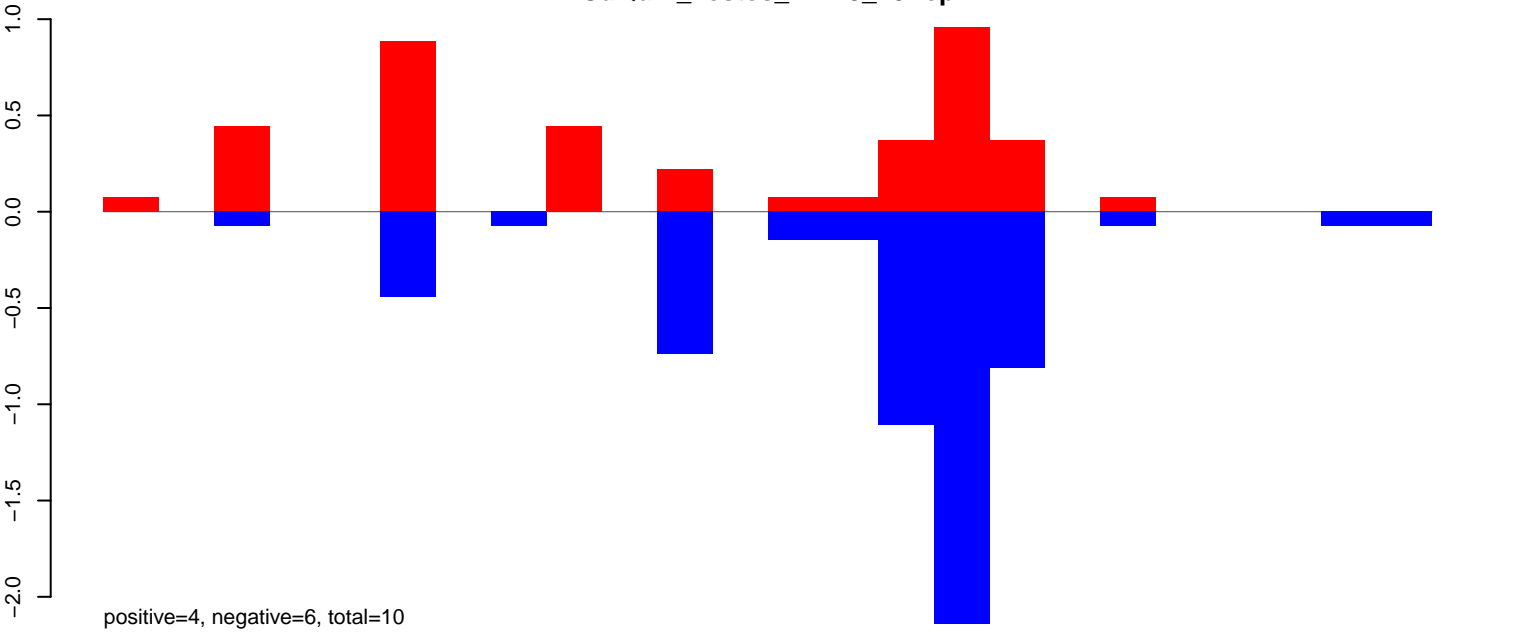
200

300

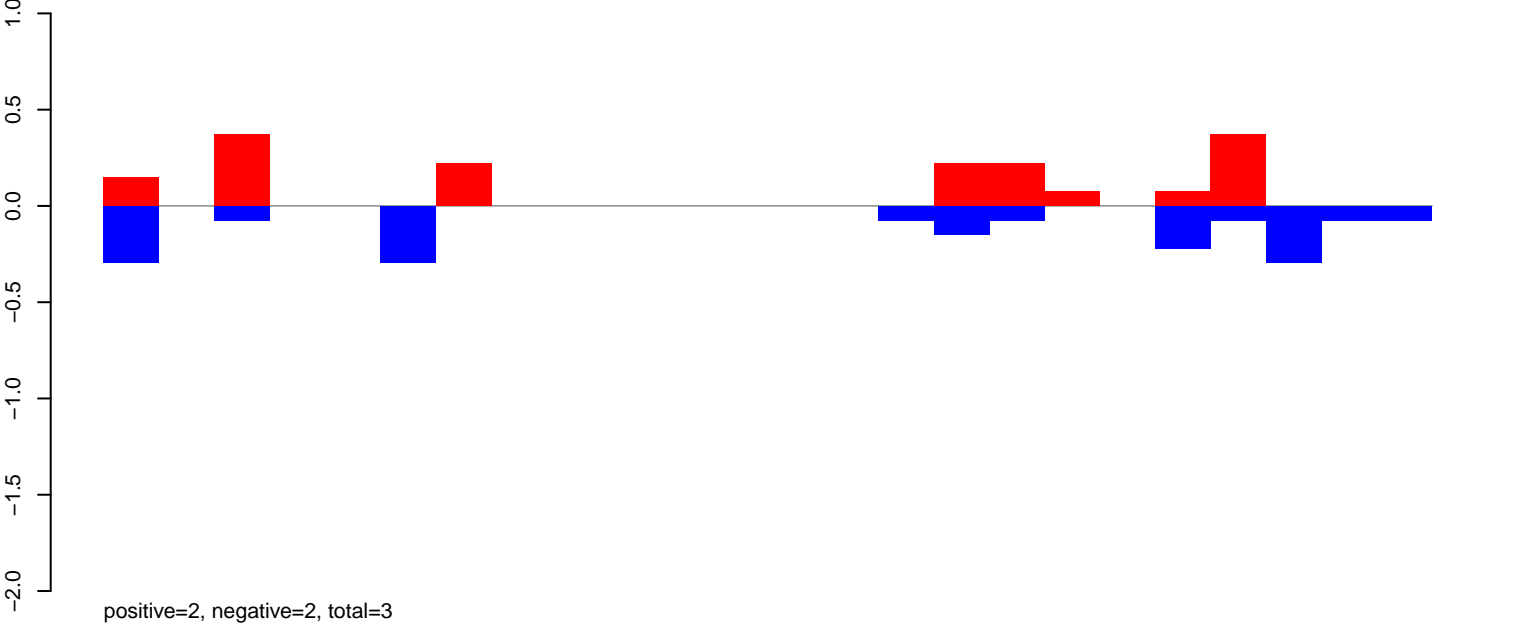
400

500

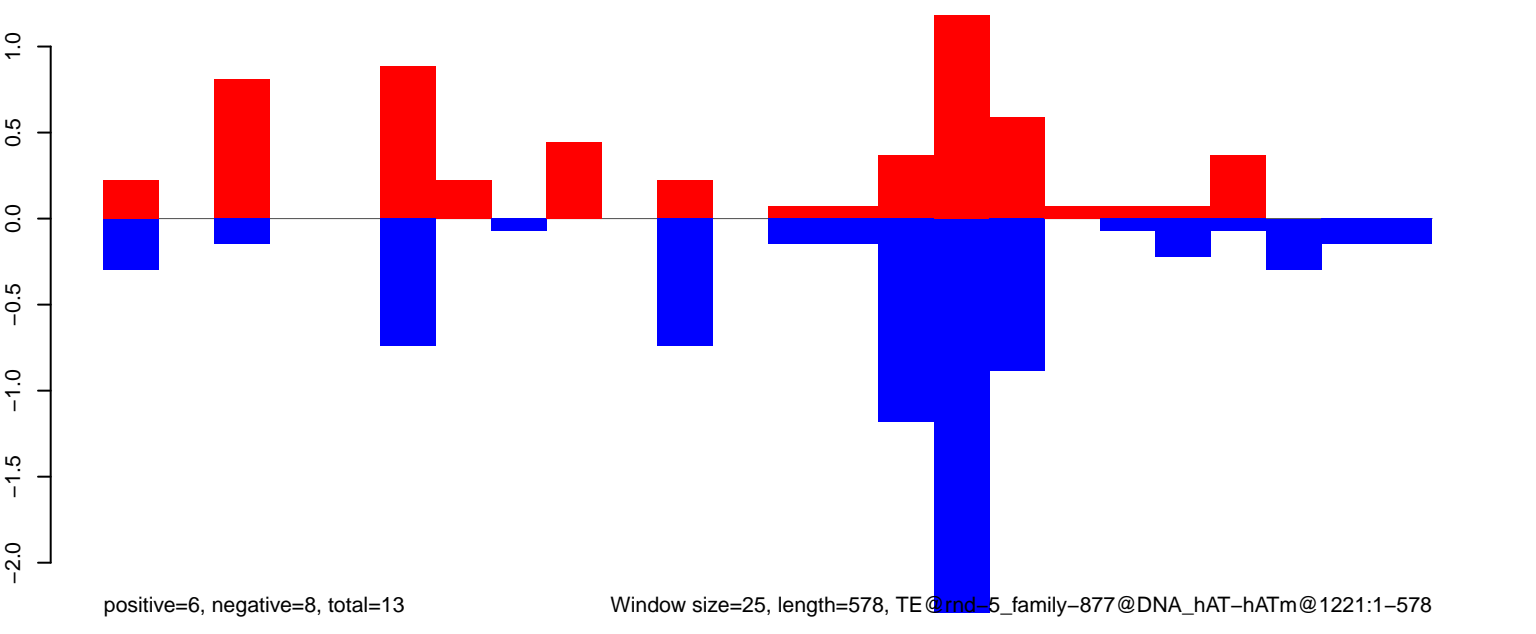
CuQuin_Testes_NL.18_23.rep



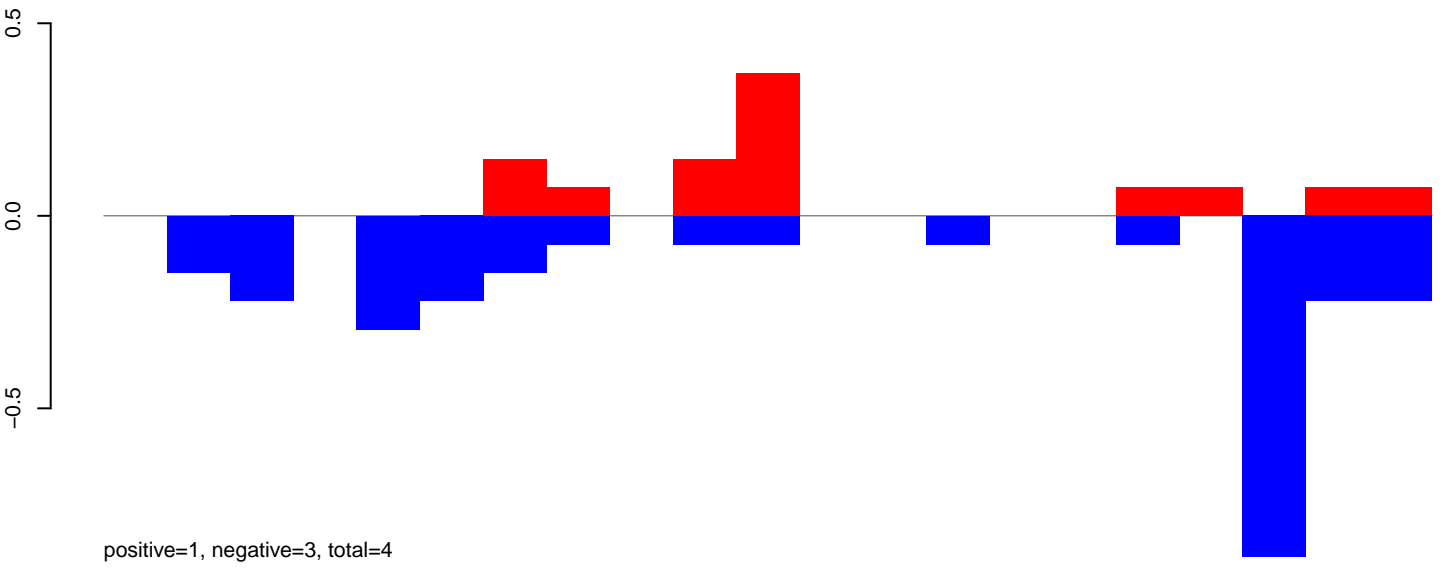
CuQuin_Testes_NL.24_35.rep



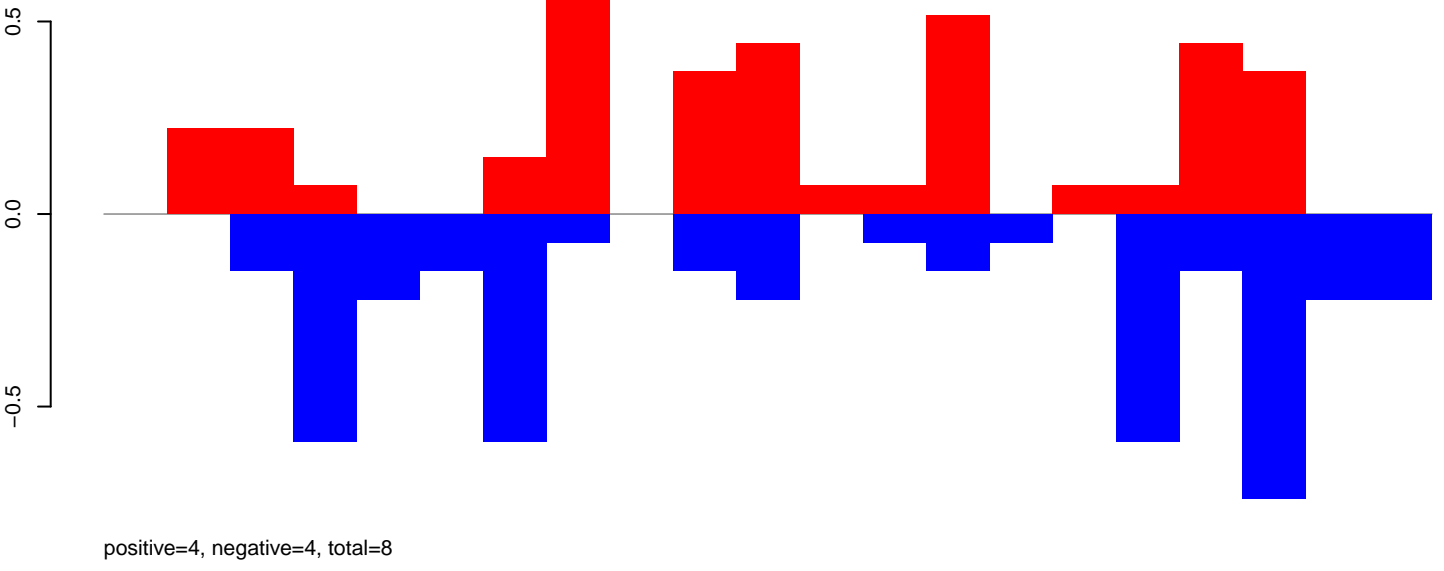
CuQuin_Testes_NL.rep



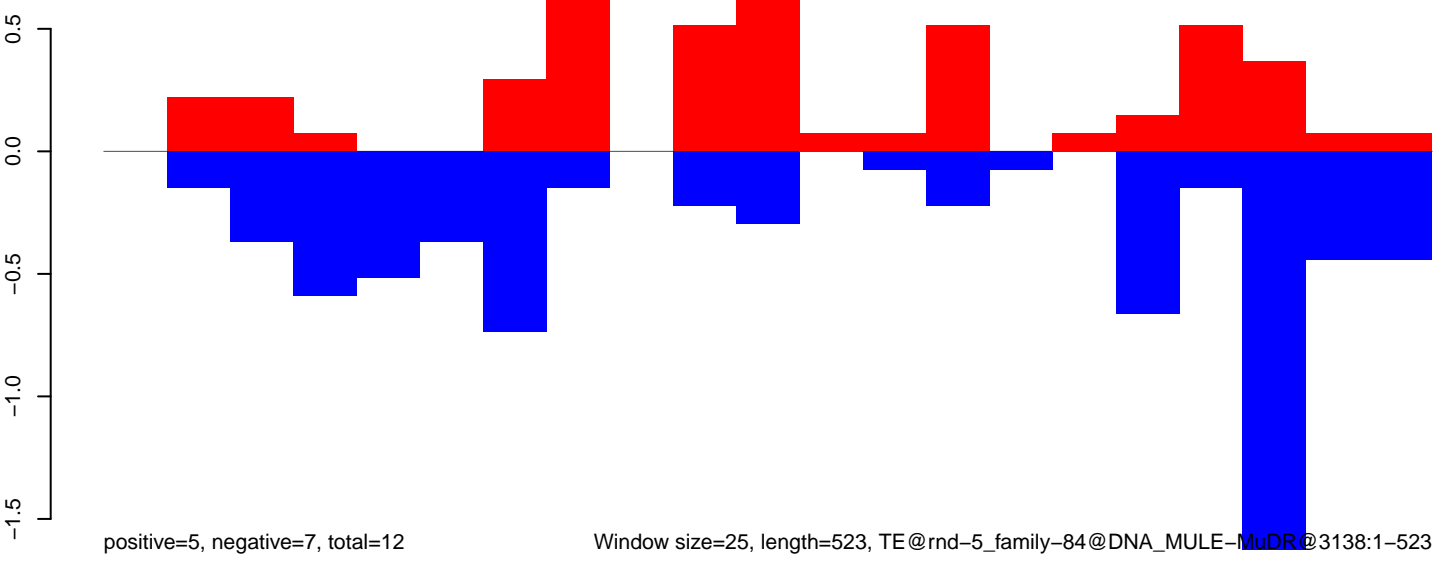
CuQuin_Testes_NL.18_23.rep



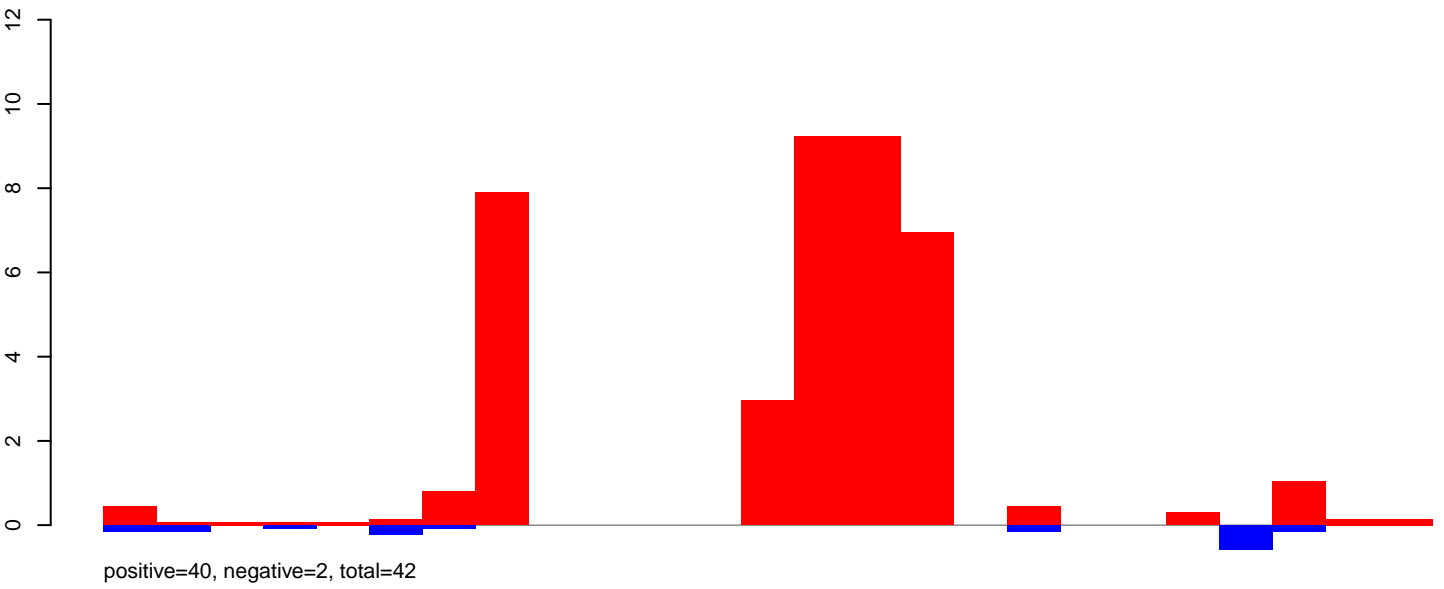
CuQuin_Testes_NL.24_35.rep



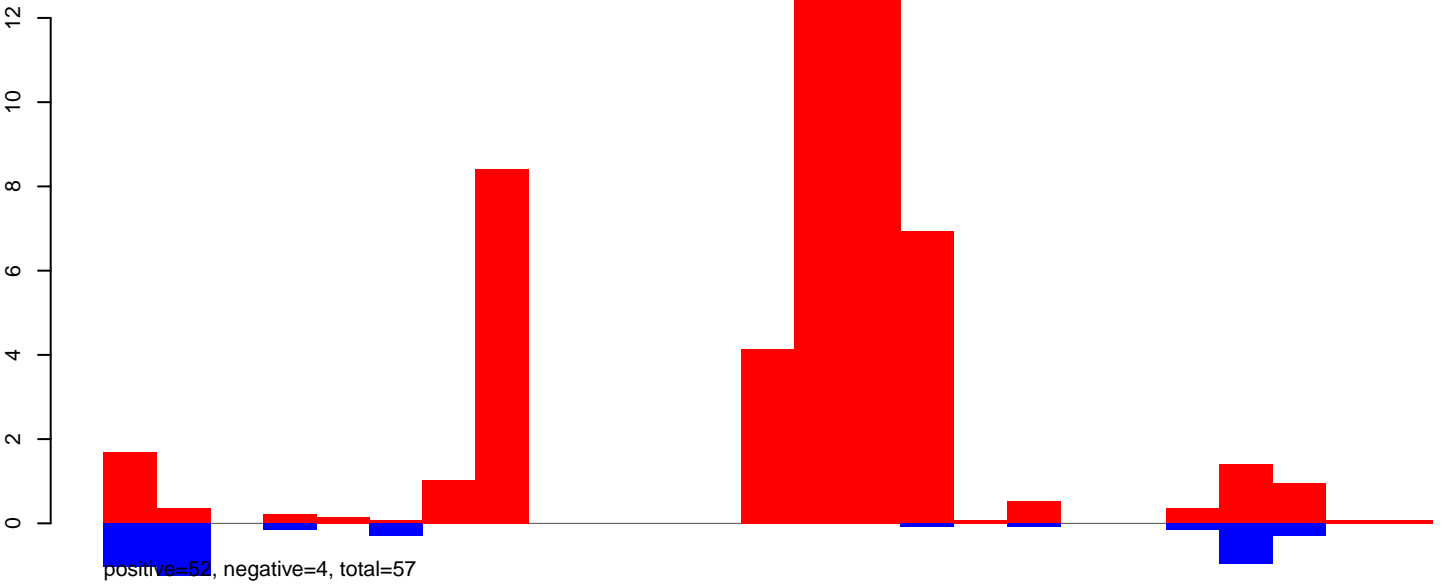
CuQuin_Testes_NL.rep



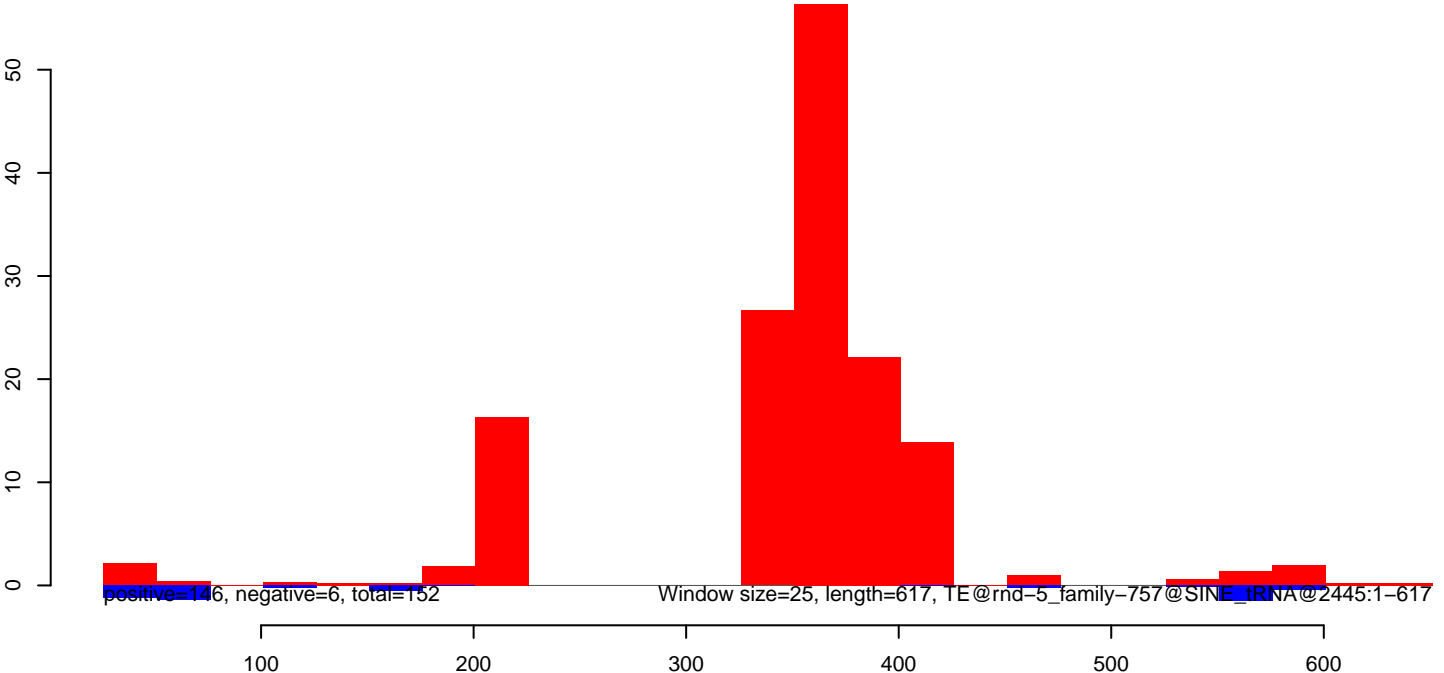
CuQuin_Testes_NL.18_23.rep



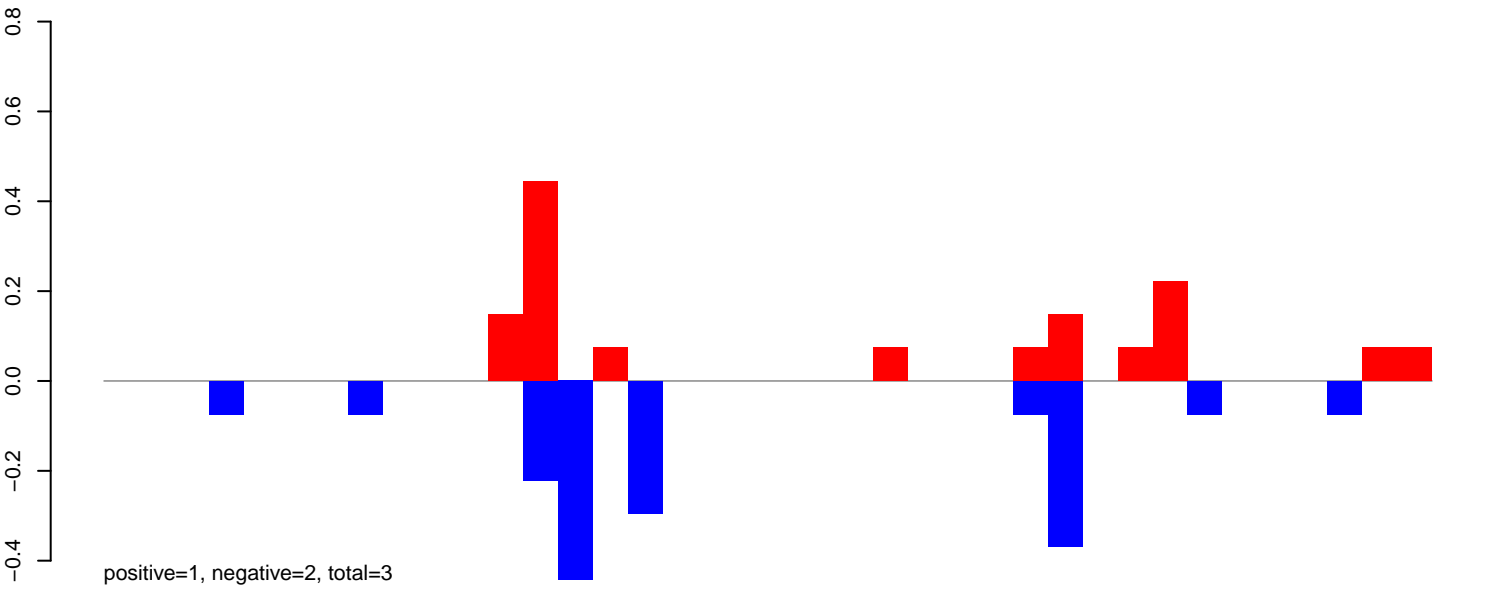
CuQuin_Testes_NL.24_35.rep



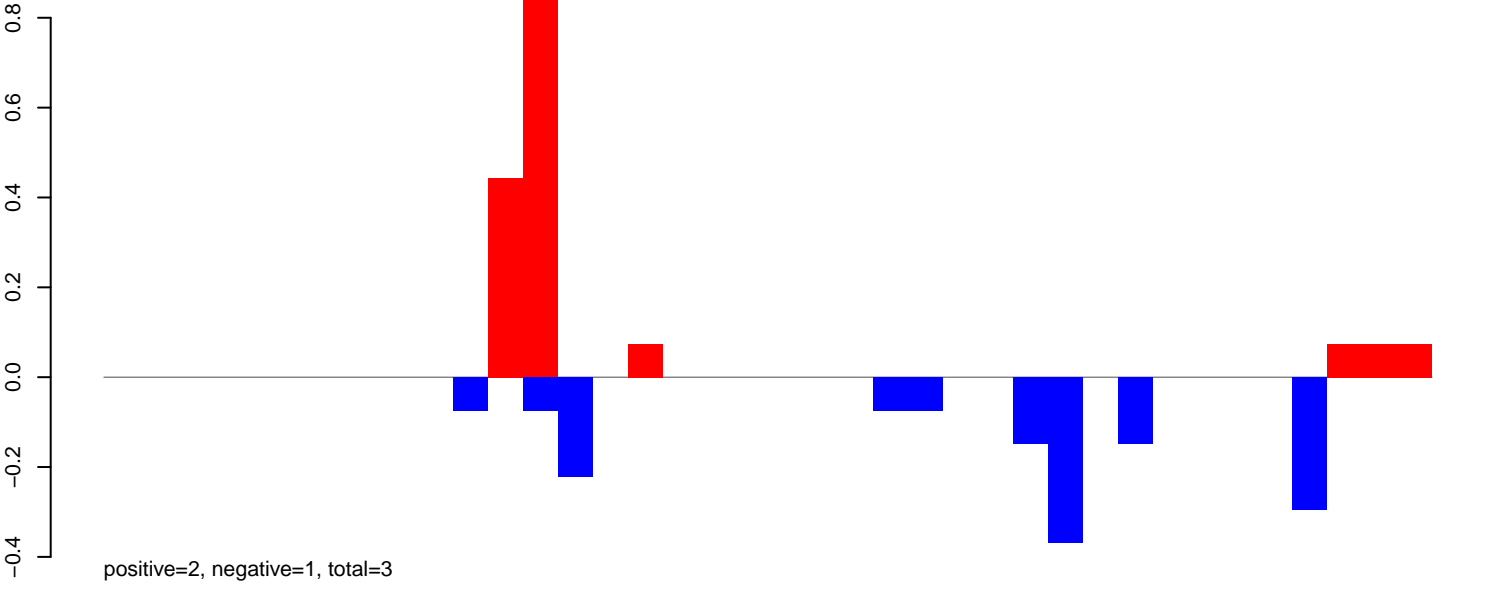
CuQuin_Testes_NL.rep



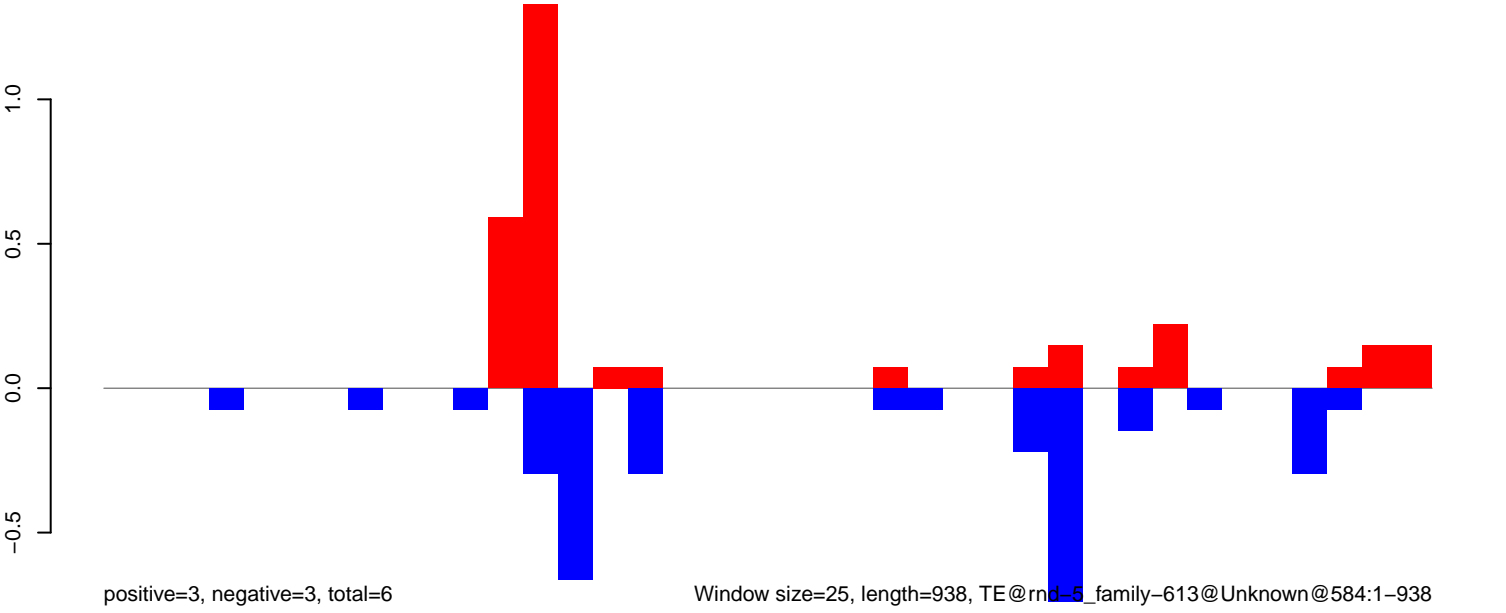
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

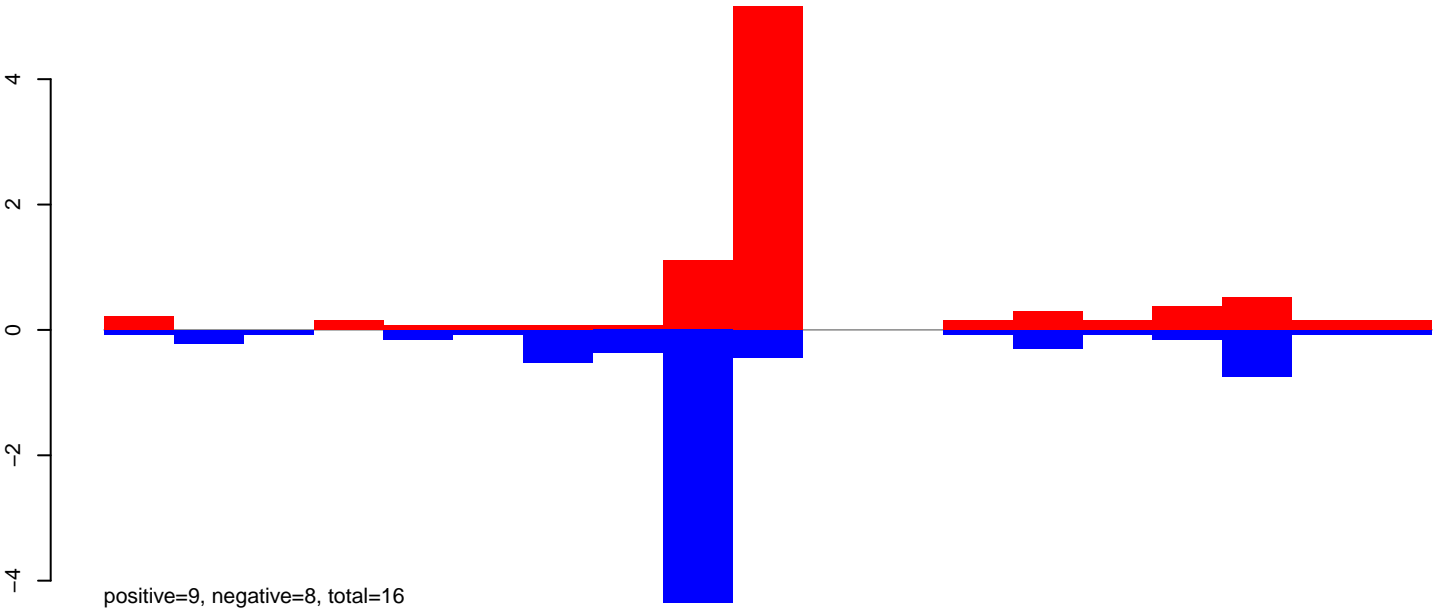


CuQuin_Testes_NL.rep

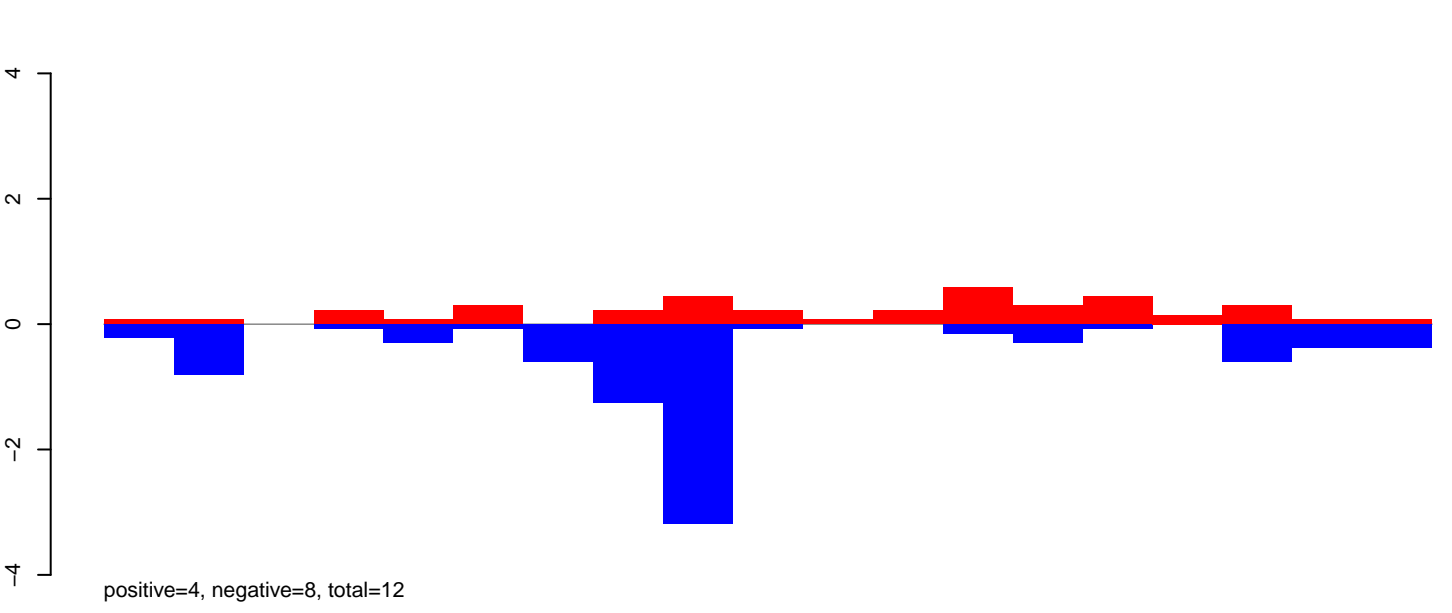


Window size=25, length=938, TE@rnl-5_family-613@Unknown@584:1-938

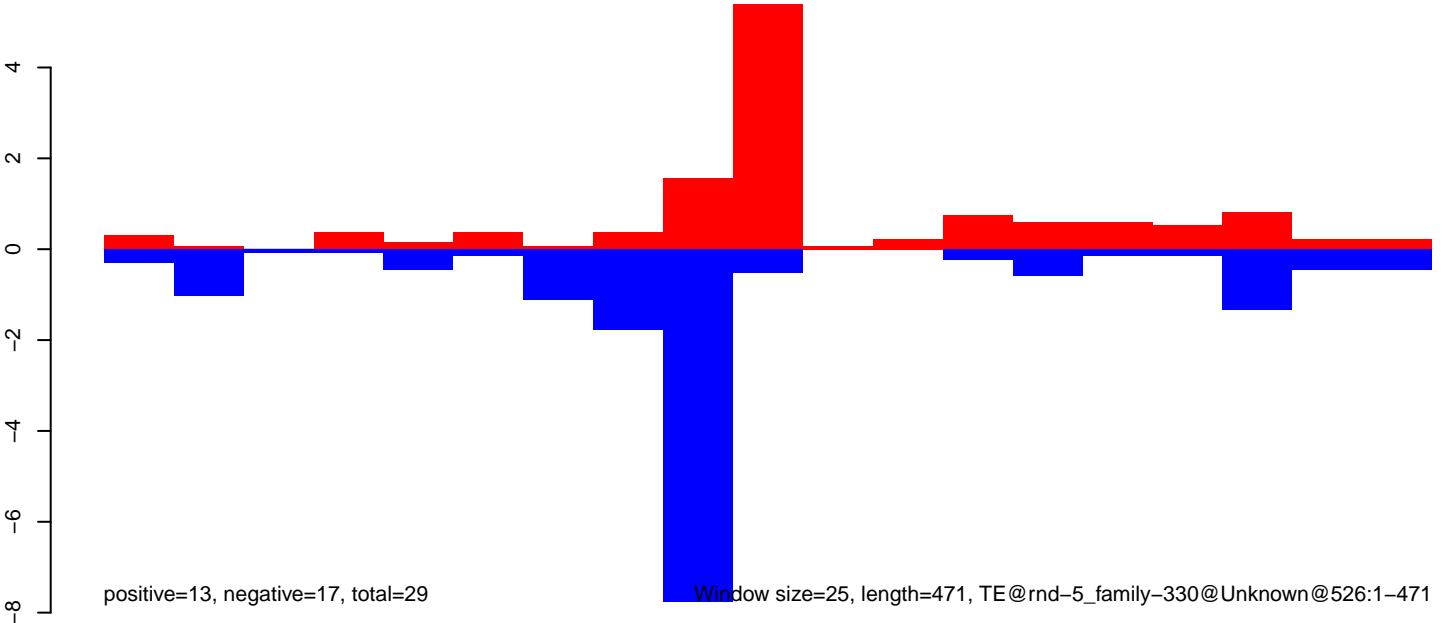
CuQuin_Testes_NL.18_23.rep



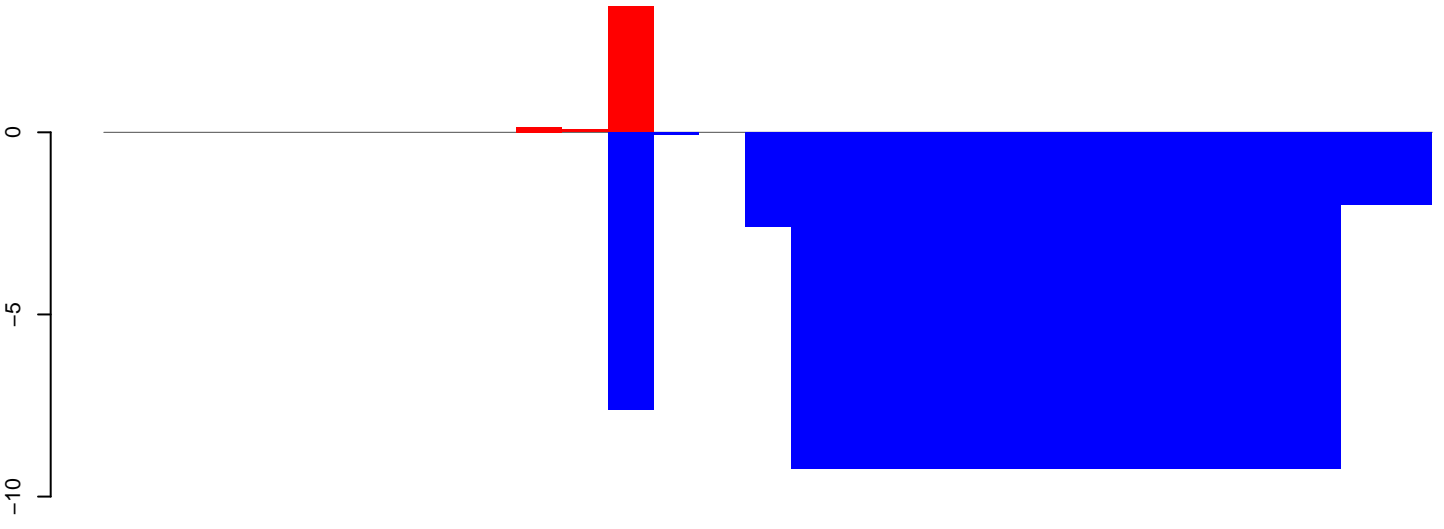
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

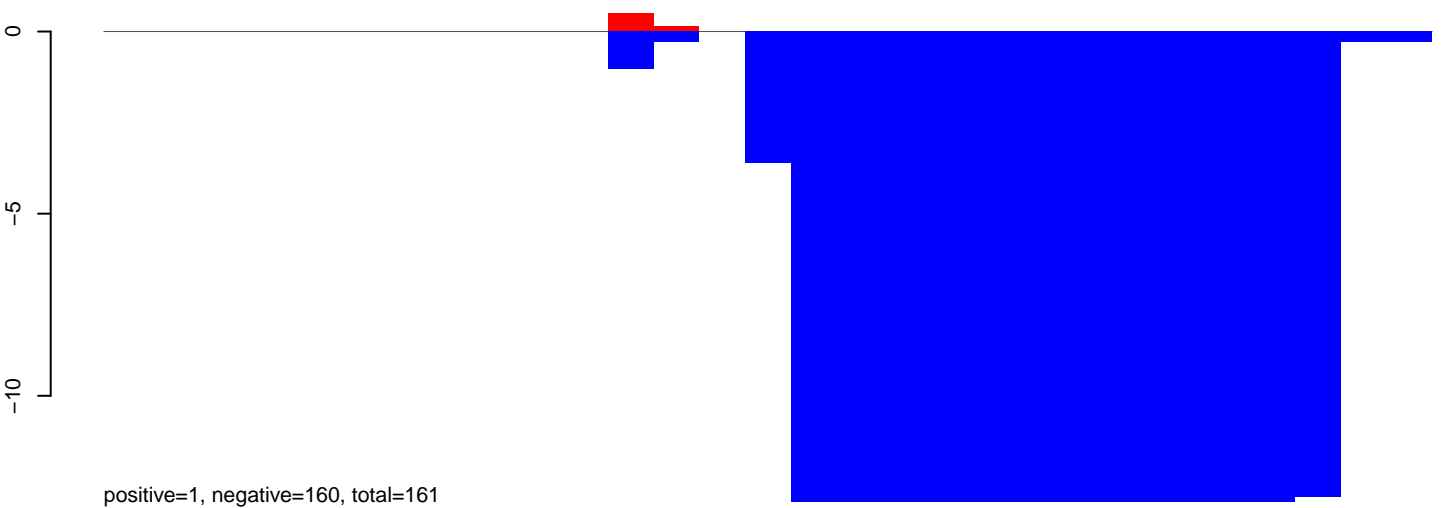


CuQuin_Testes_NL.18_23.rep



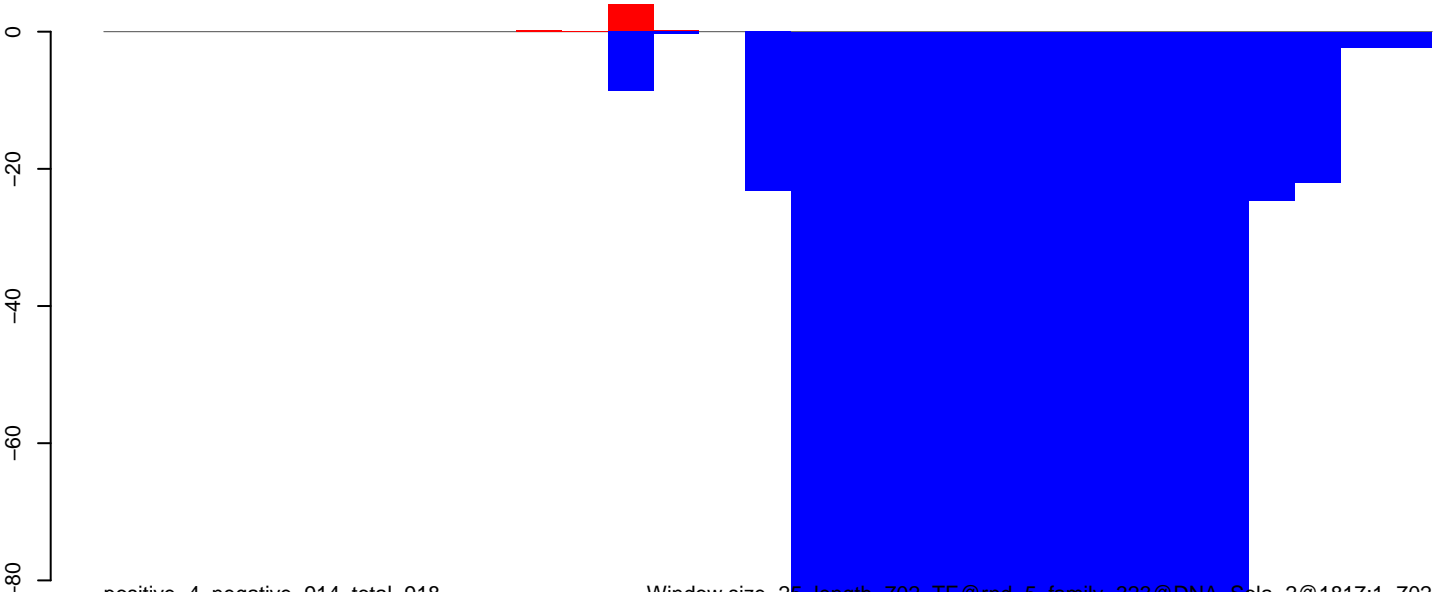
positive=4, negative=125, total=129

CuQuin_Testes_NL.24_35.rep



positive=1, negative=160, total=161

CuQuin_Testes_NL.rep

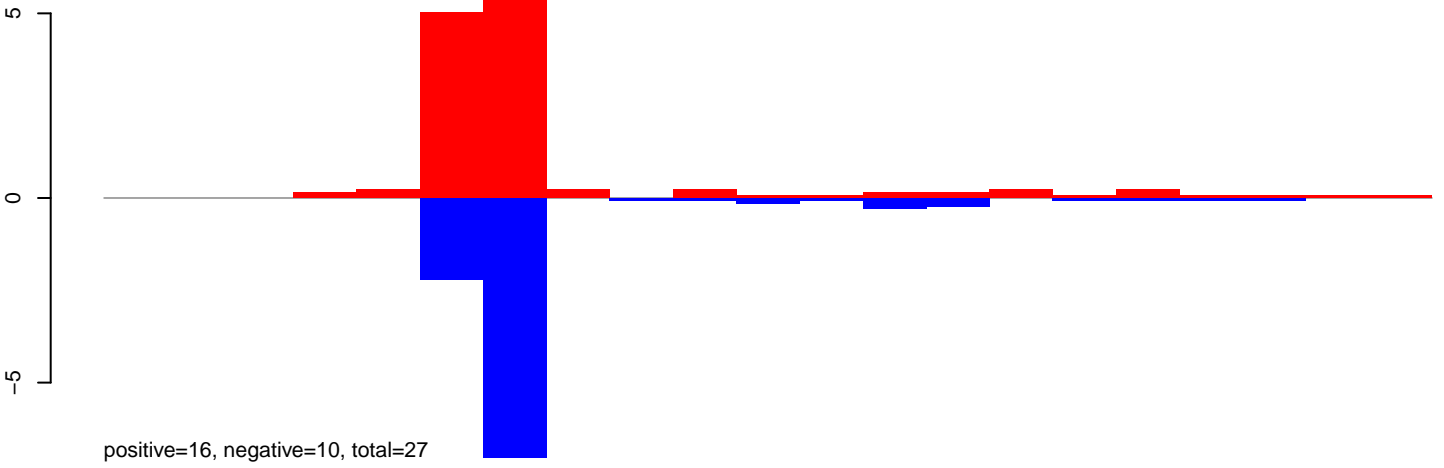


positive=4, negative=914, total=918

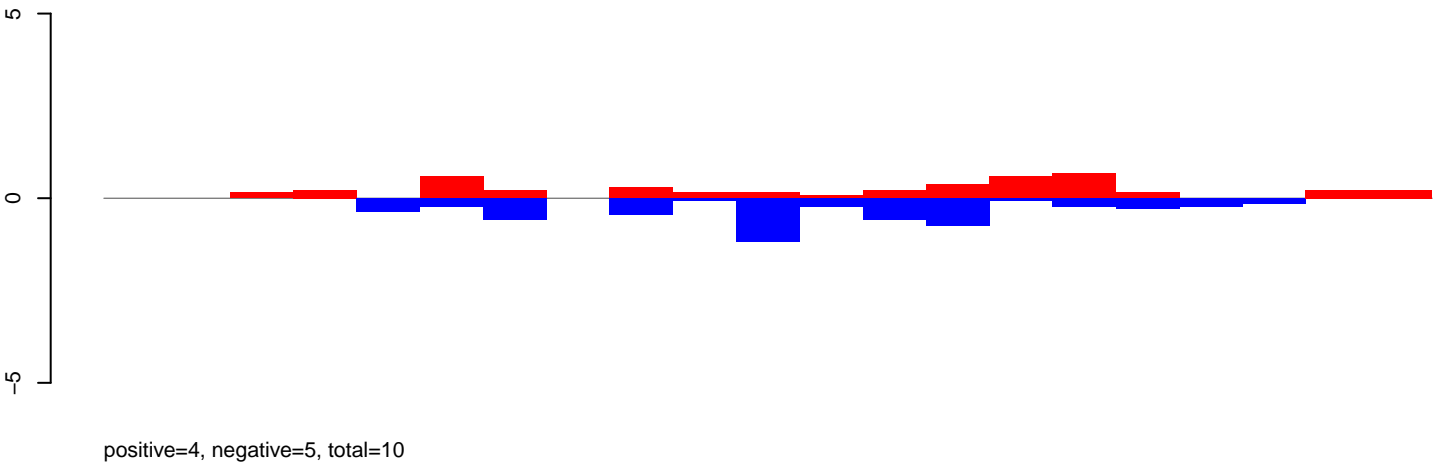
Window size=25, length=702, TE@rnd-5_family-323@DNA_Sola-2@1817:1-702

0 200 400 600

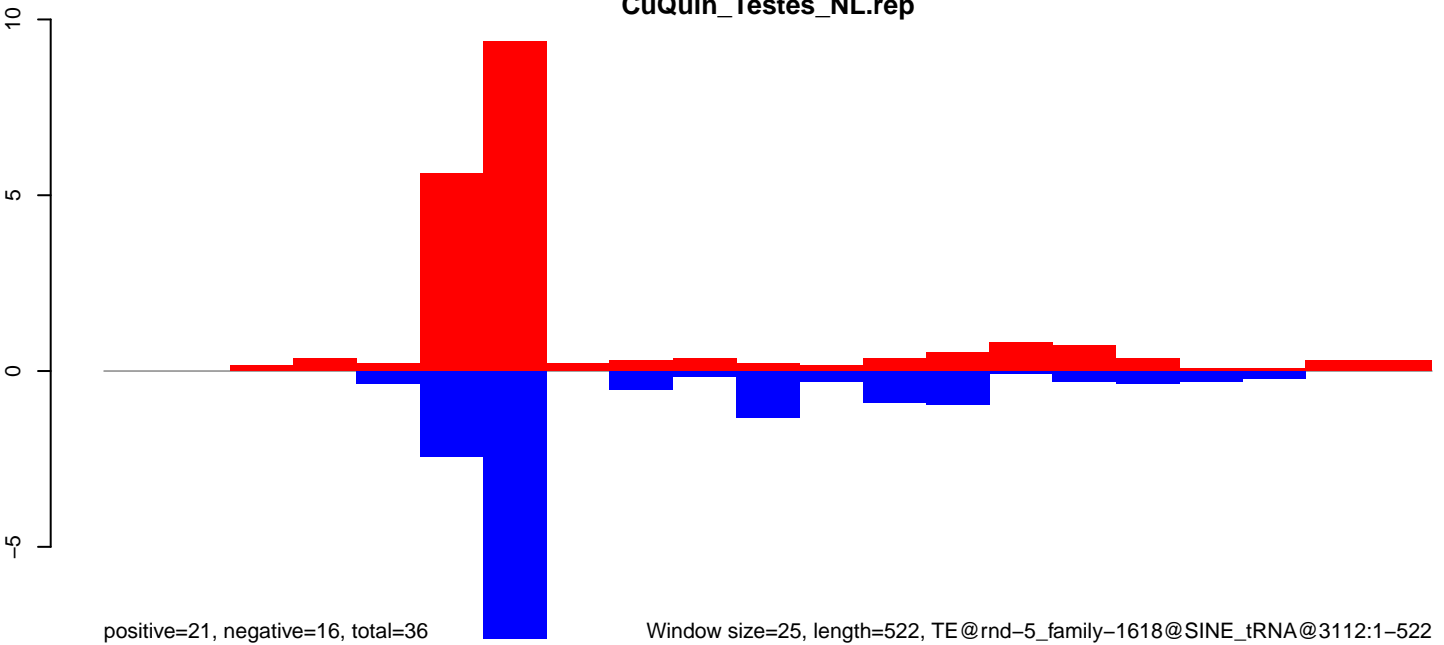
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

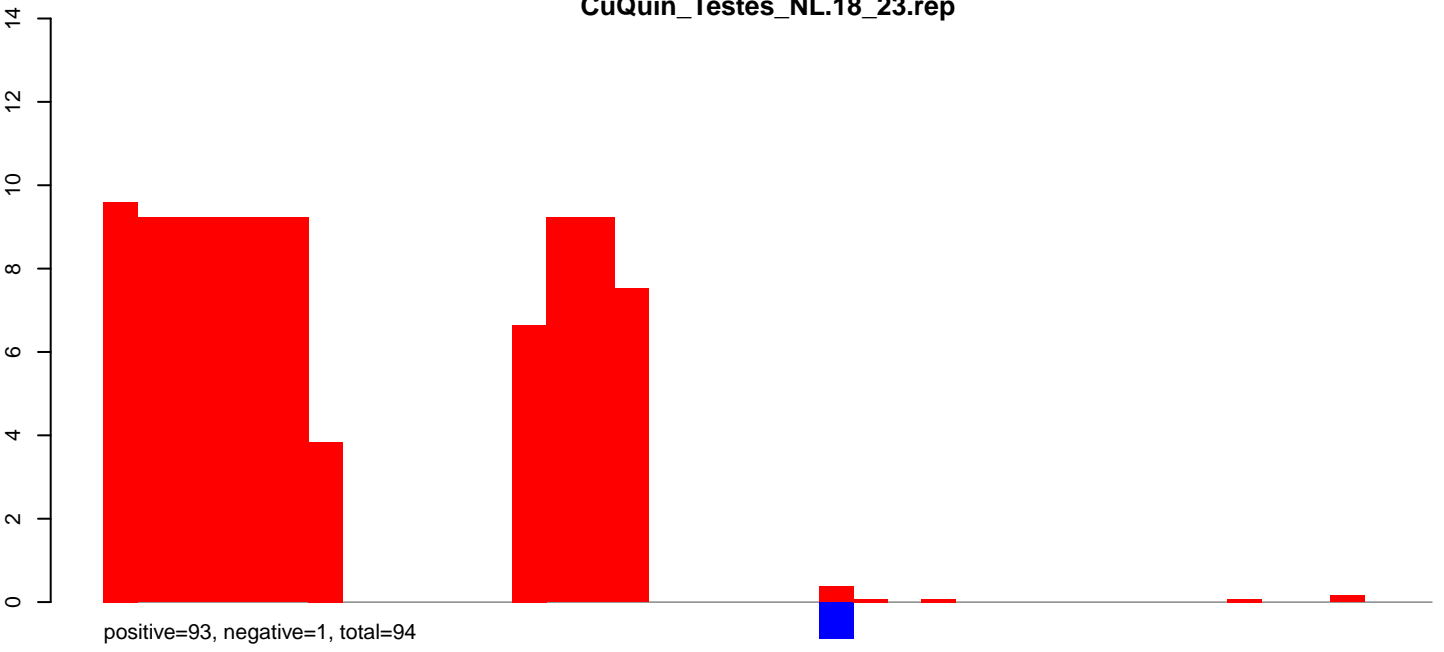


CuQuin_Testes_NL.rep

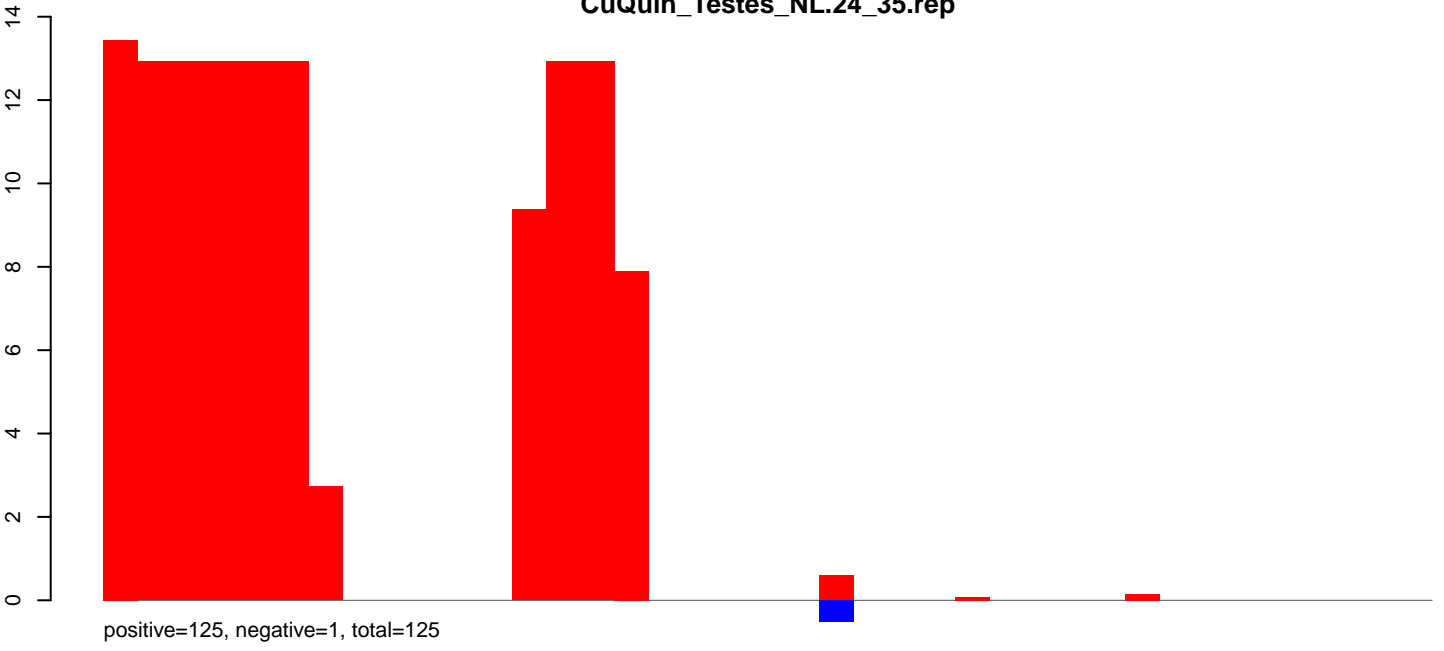


Window size=25, length=522, TE@rnd-5_family-1618@SINE_tRNA@3112:1-522

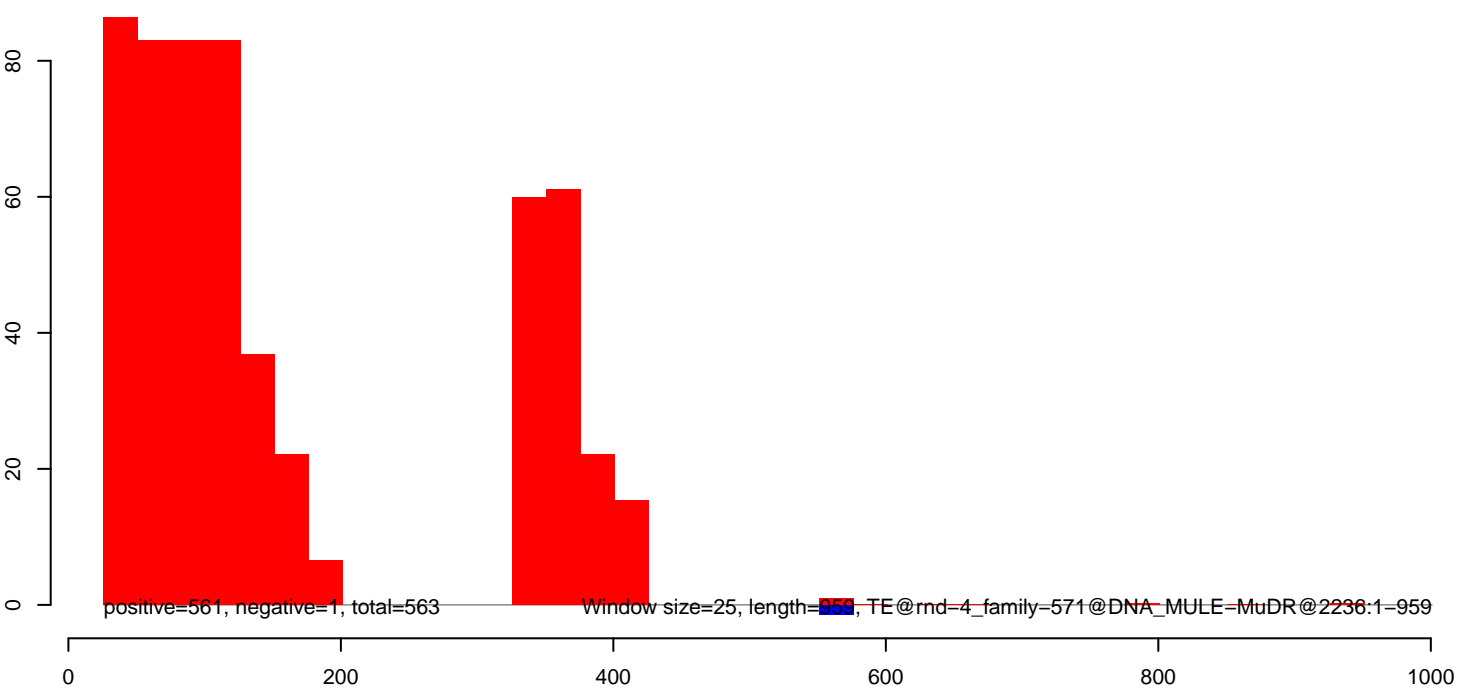
CuQuin_Testes_NL.18_23.rep



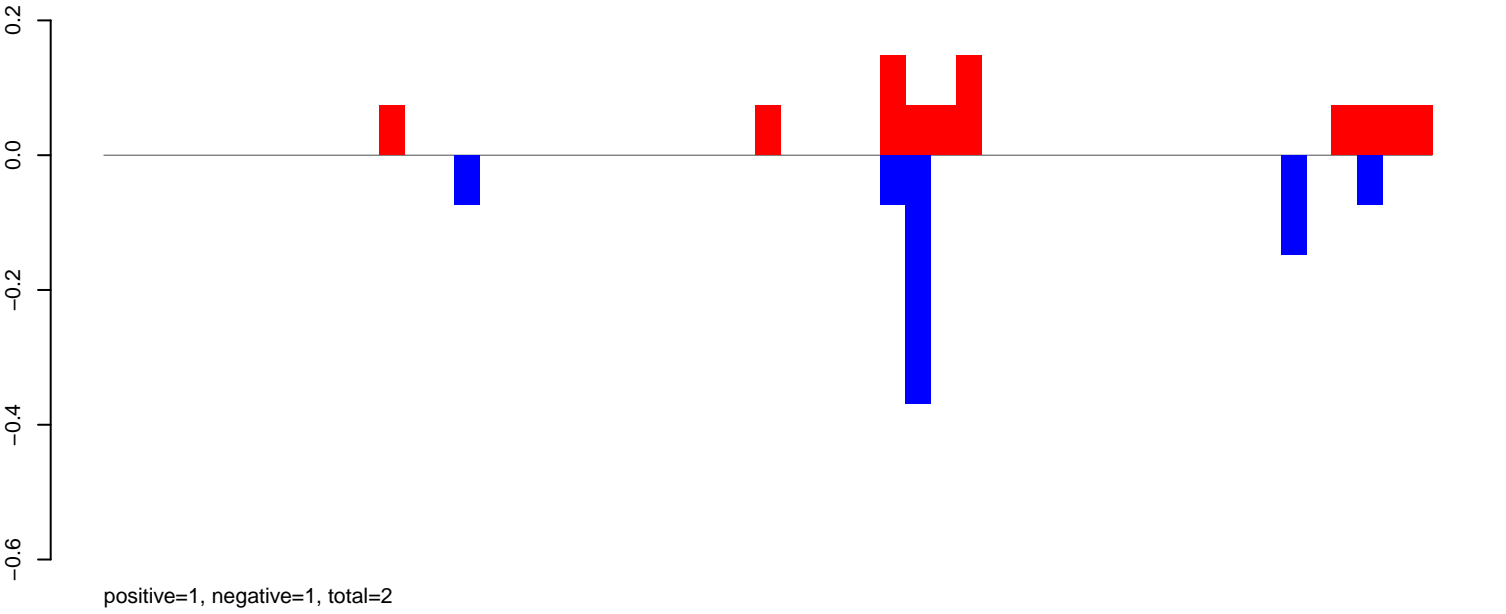
CuQuin_Testes_NL.24_35.rep



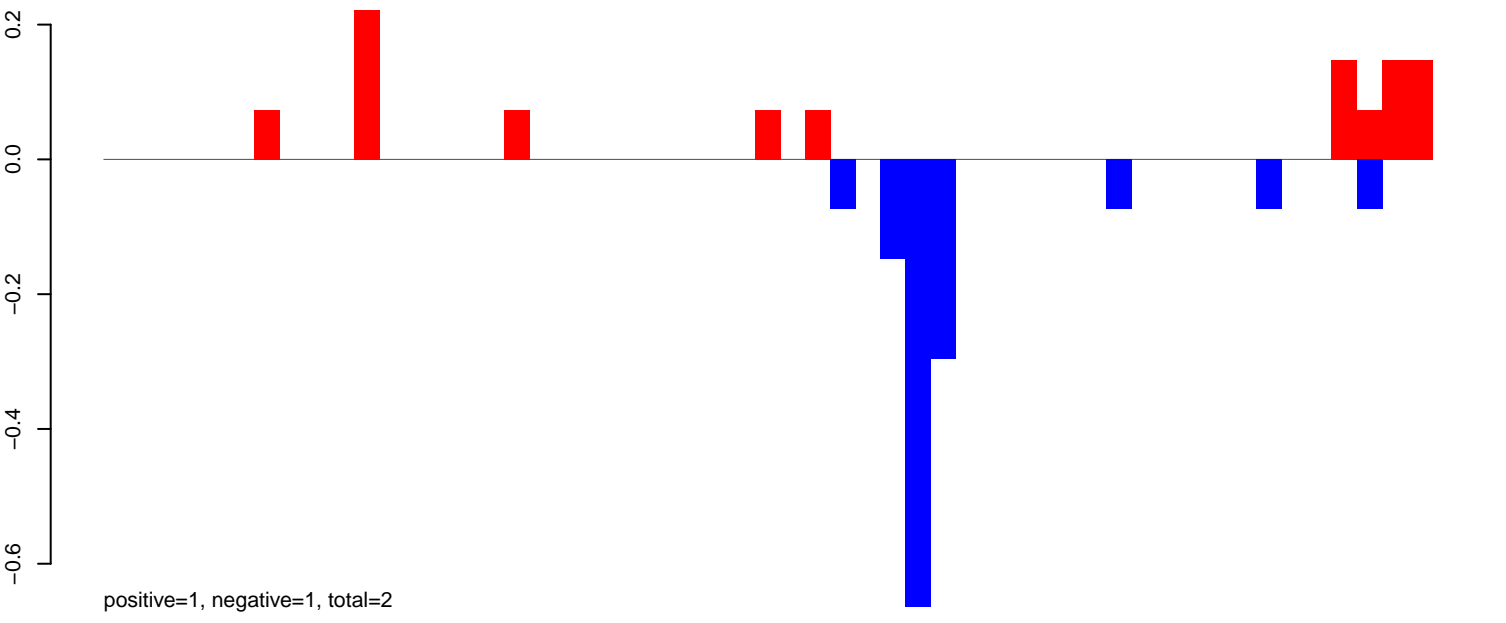
CuQuin_Testes_NL.rep



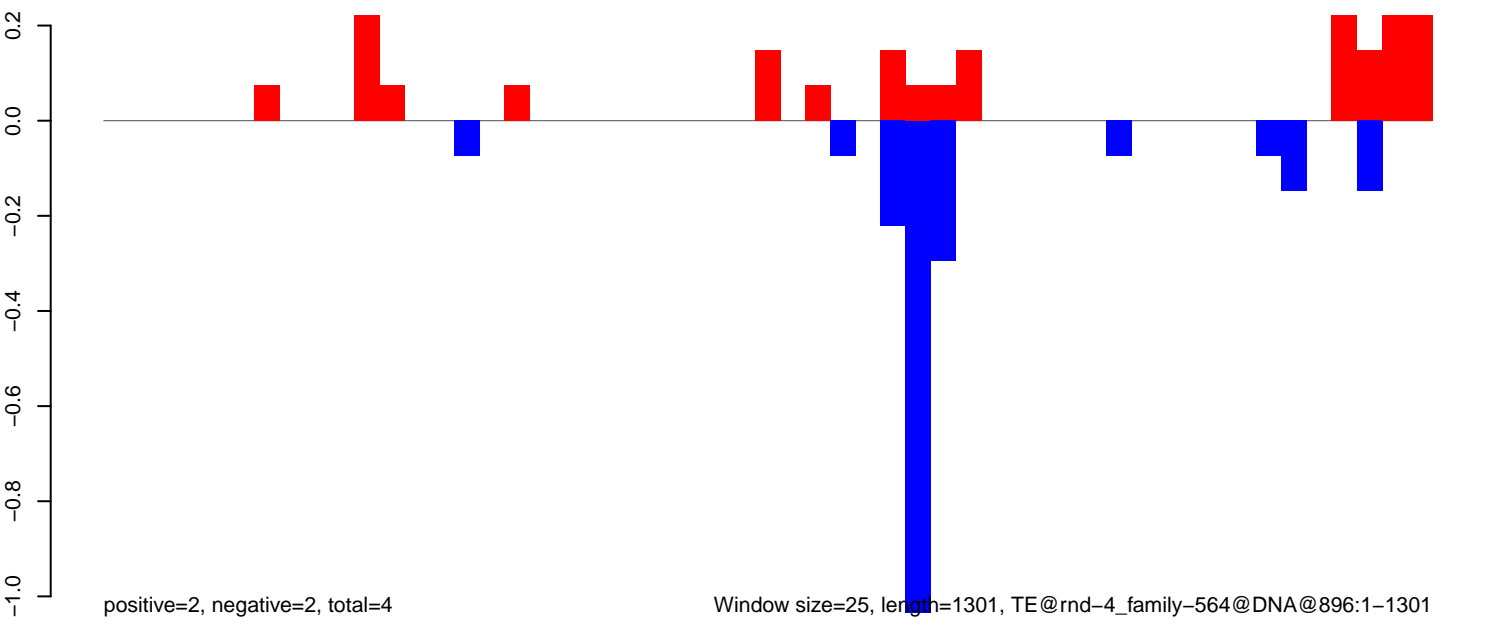
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

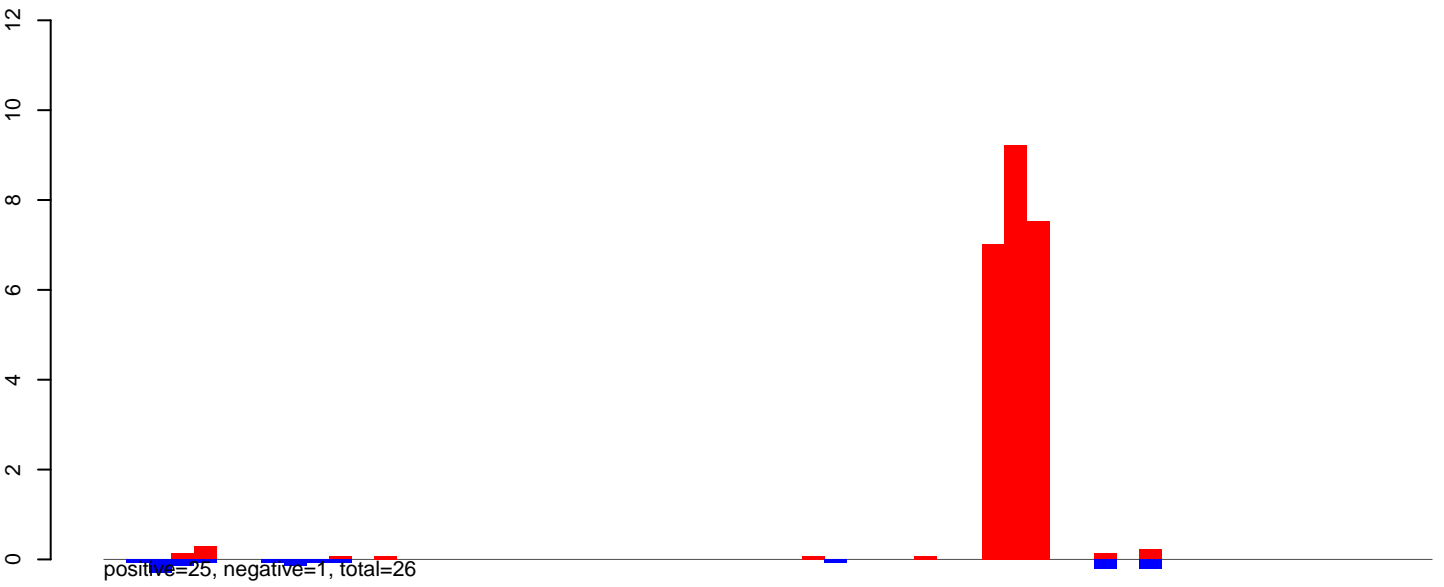


CuQuin_Testes_NL.rep

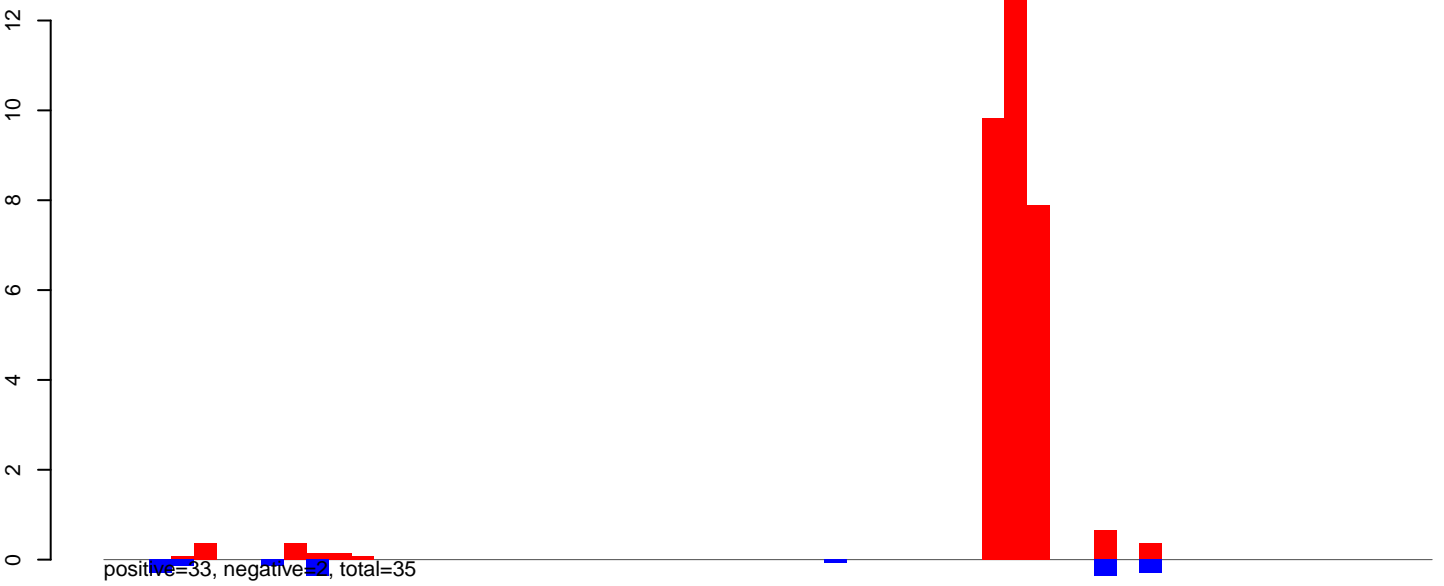


Window size=25, length=1301, TE@rnd-4_family-564@DNA@896:1-1301

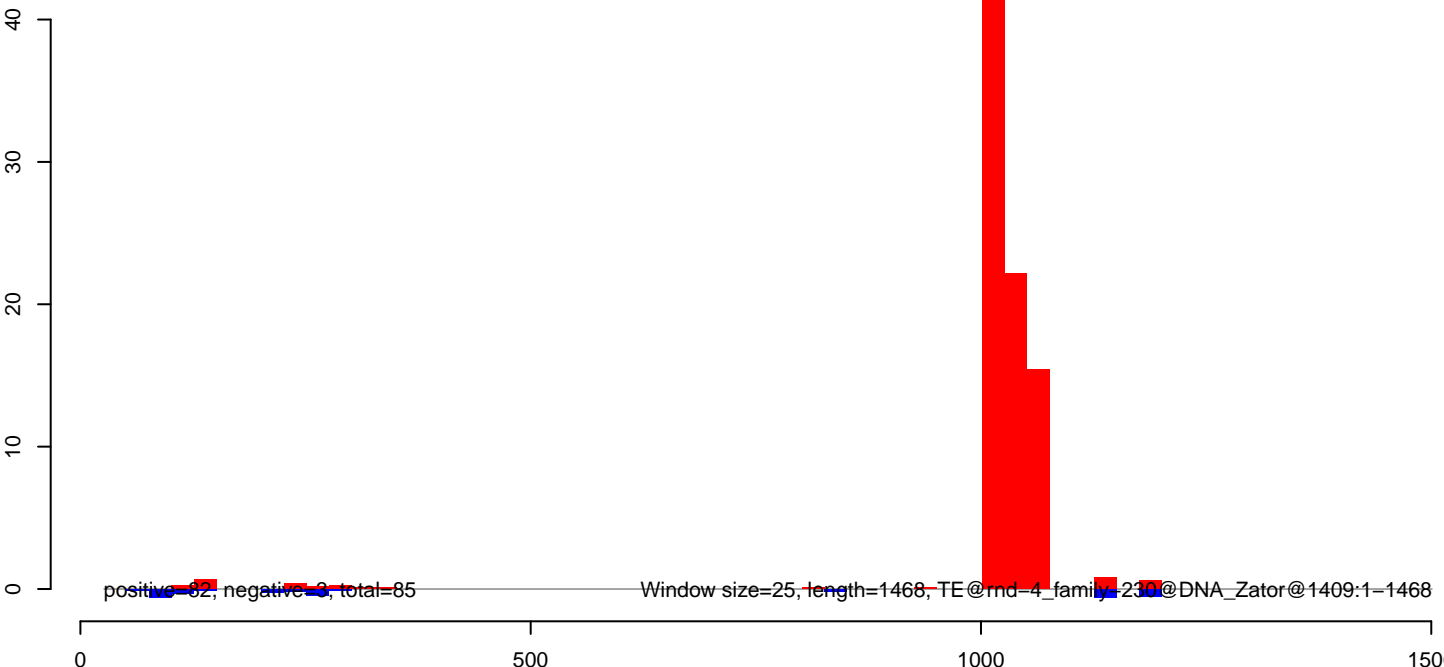
CuQuin_Testes_NL.18_23.rep



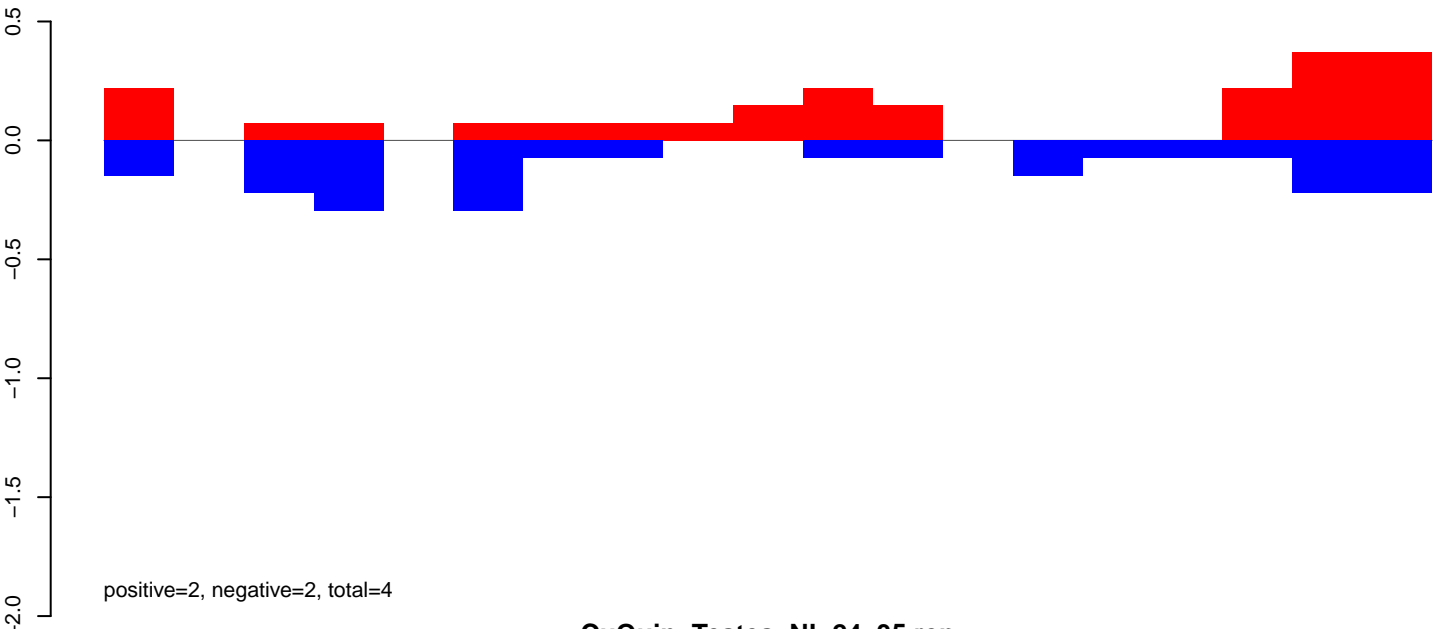
CuQuin_Testes_NL.24_35.rep



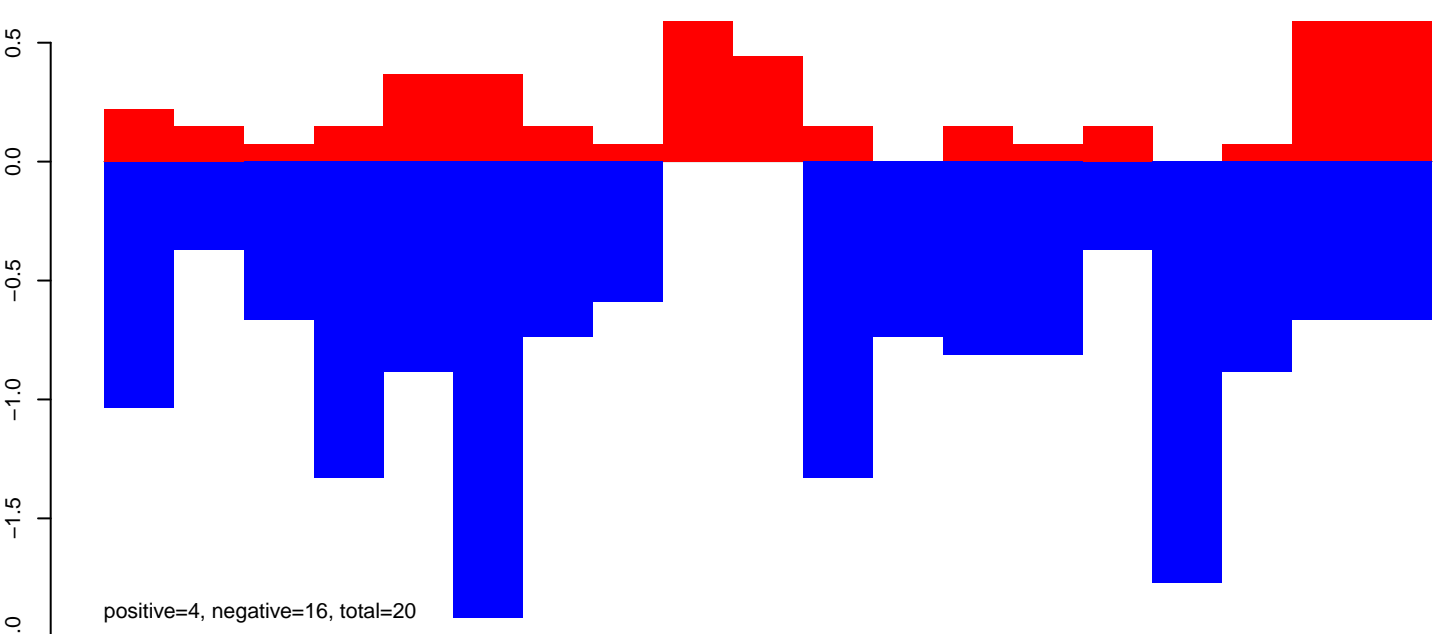
CuQuin_Testes_NL.rep



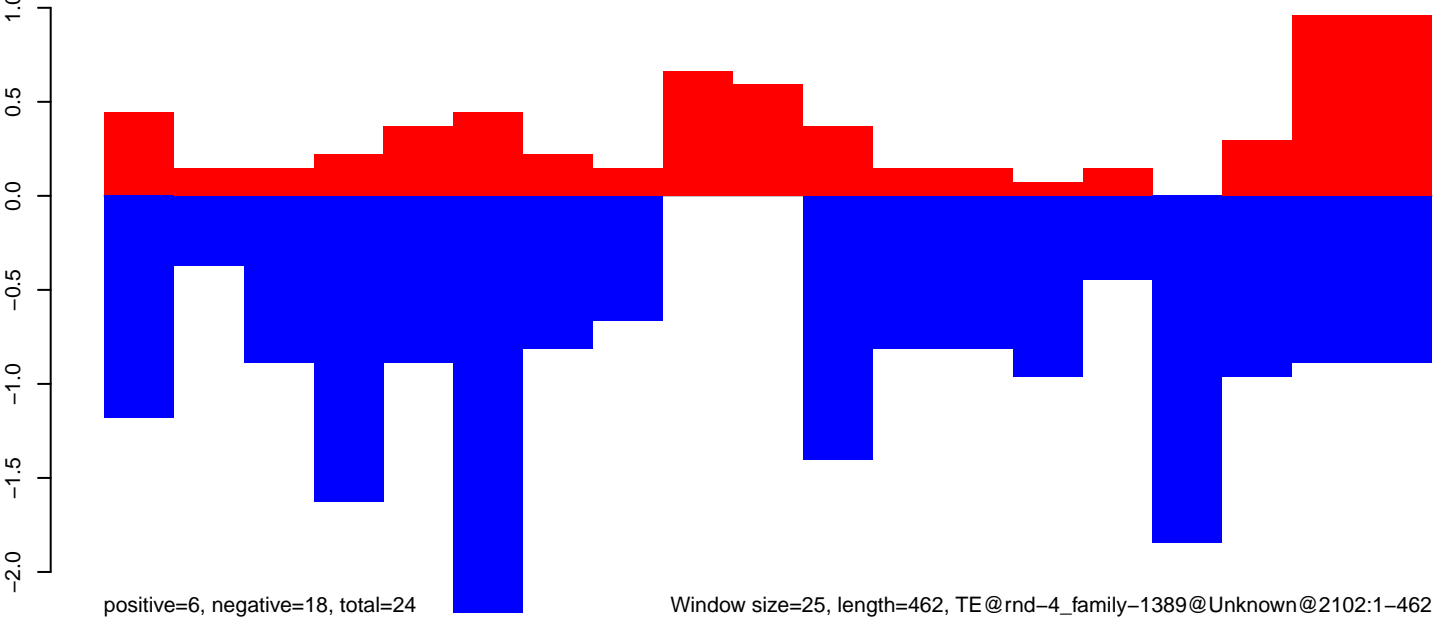
CuQuin_Testes_NL.18_23.rep



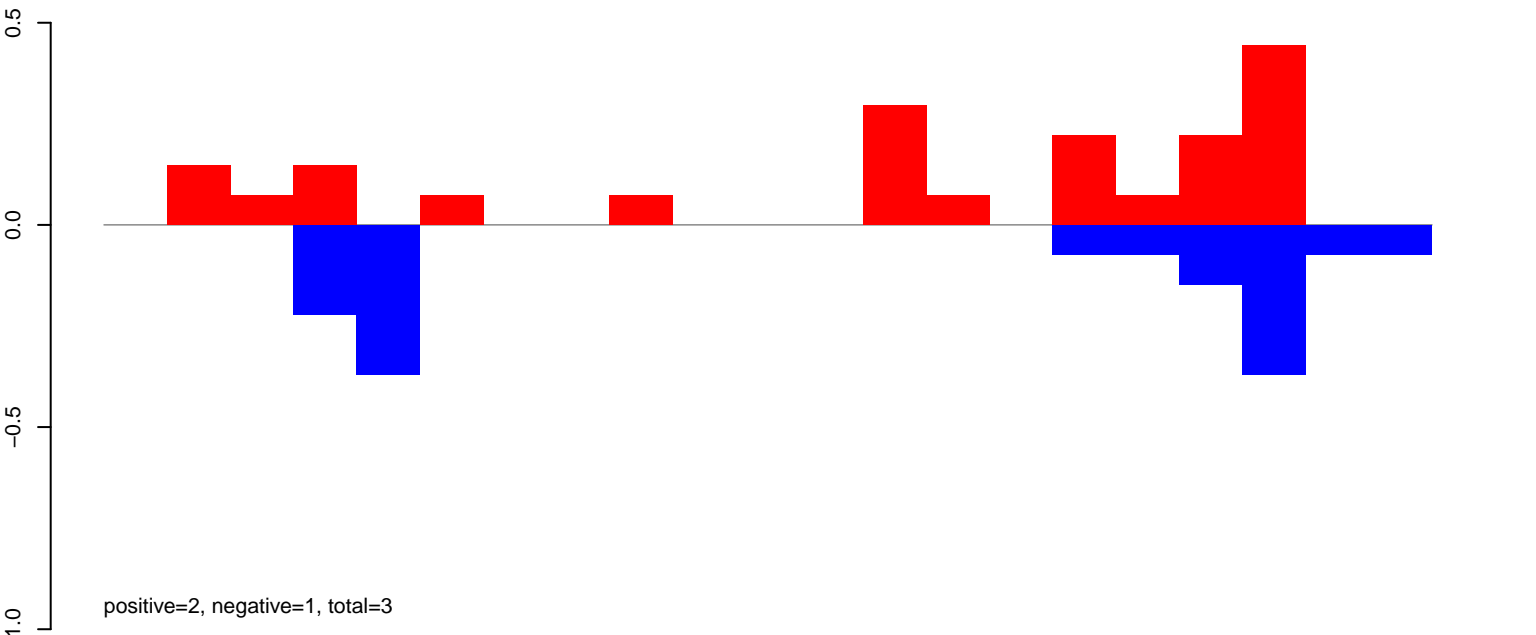
CuQuin_Testes_NL.24_35.rep



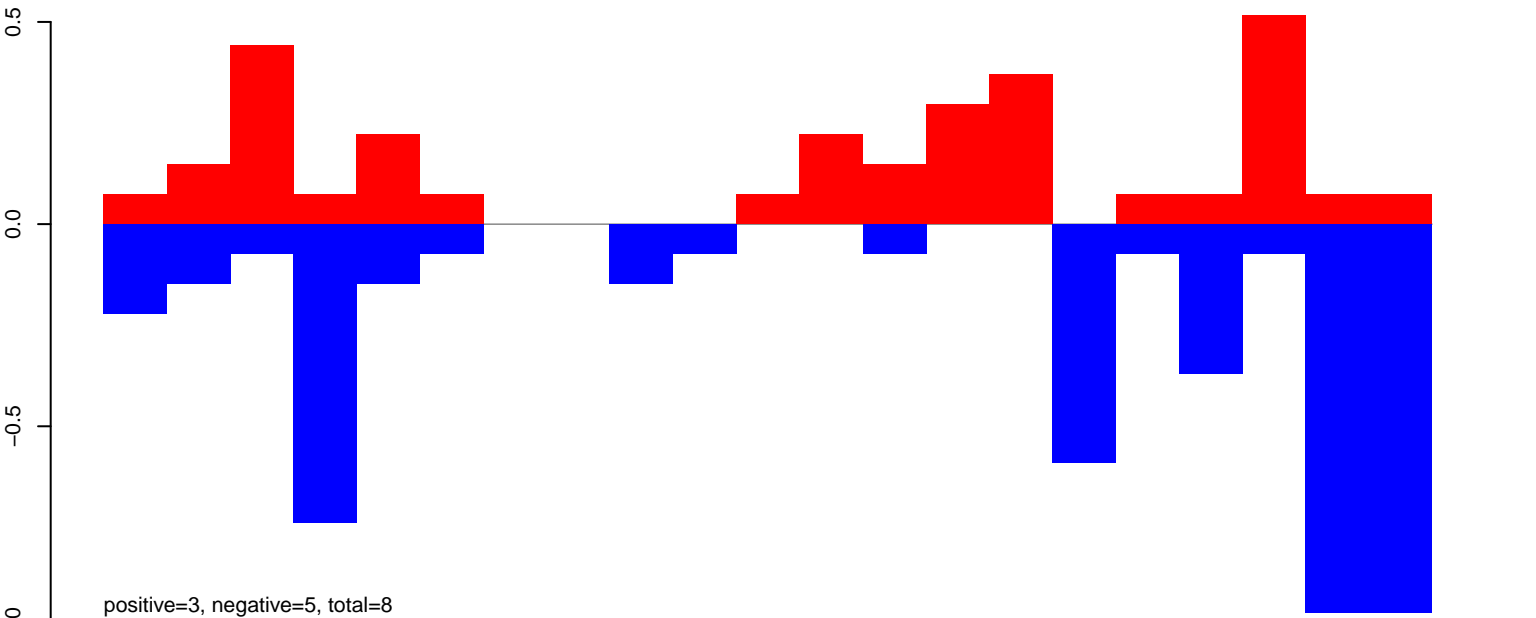
CuQuin_Testes_NL.rep



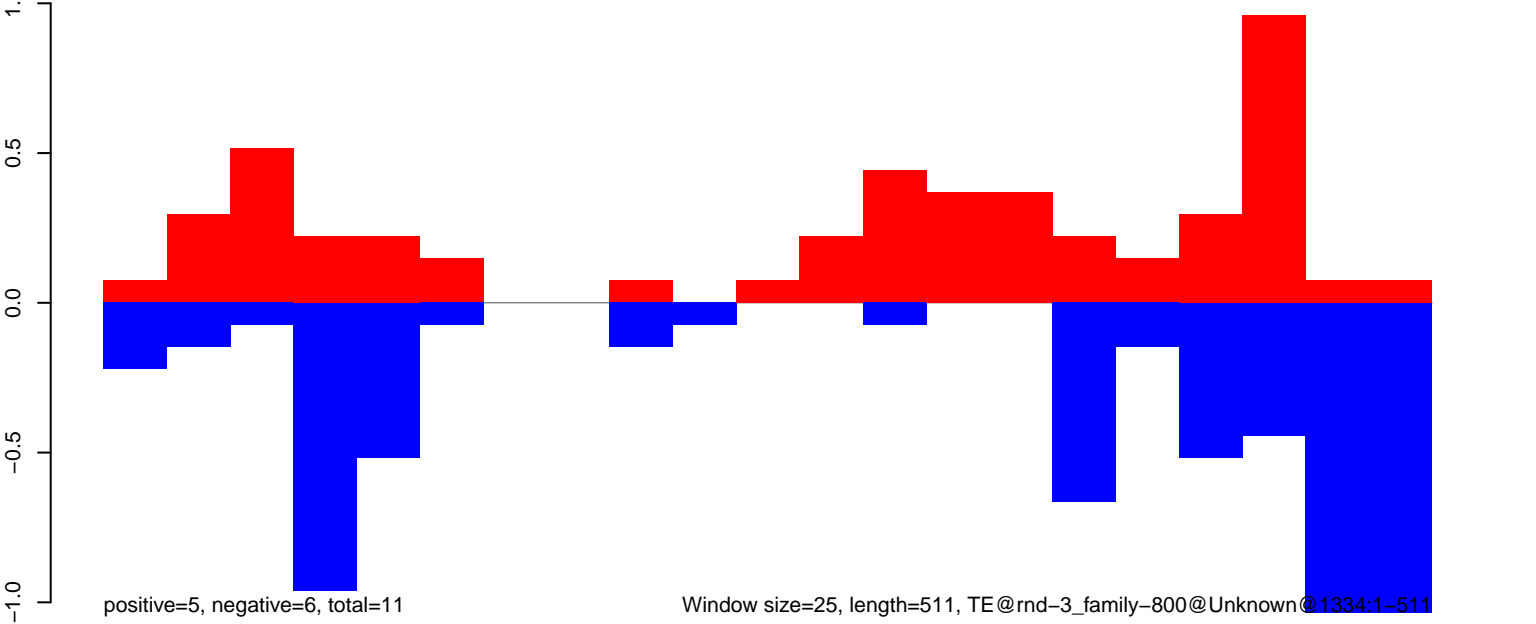
CuQuin_Testes_NL.18_23.rep



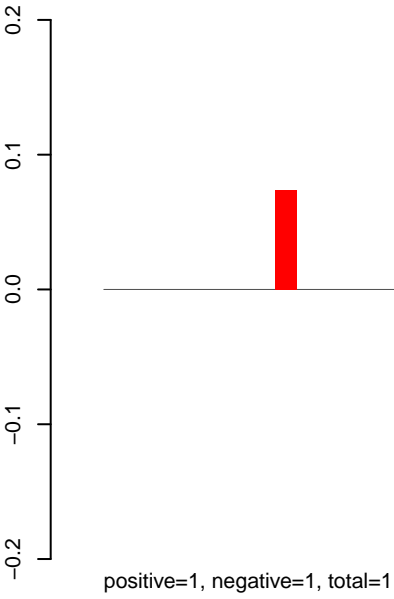
CuQuin_Testes_NL.24_35.rep



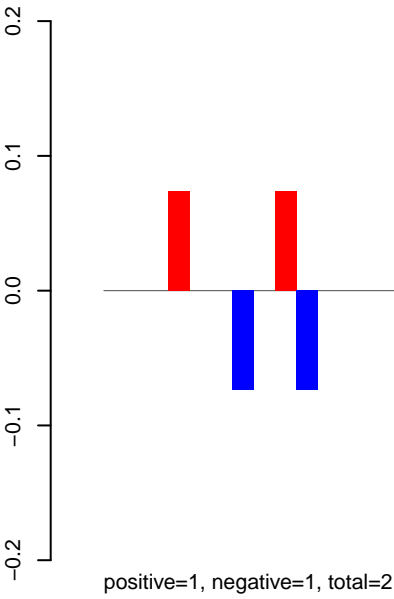
CuQuin_Testes_NL.rep



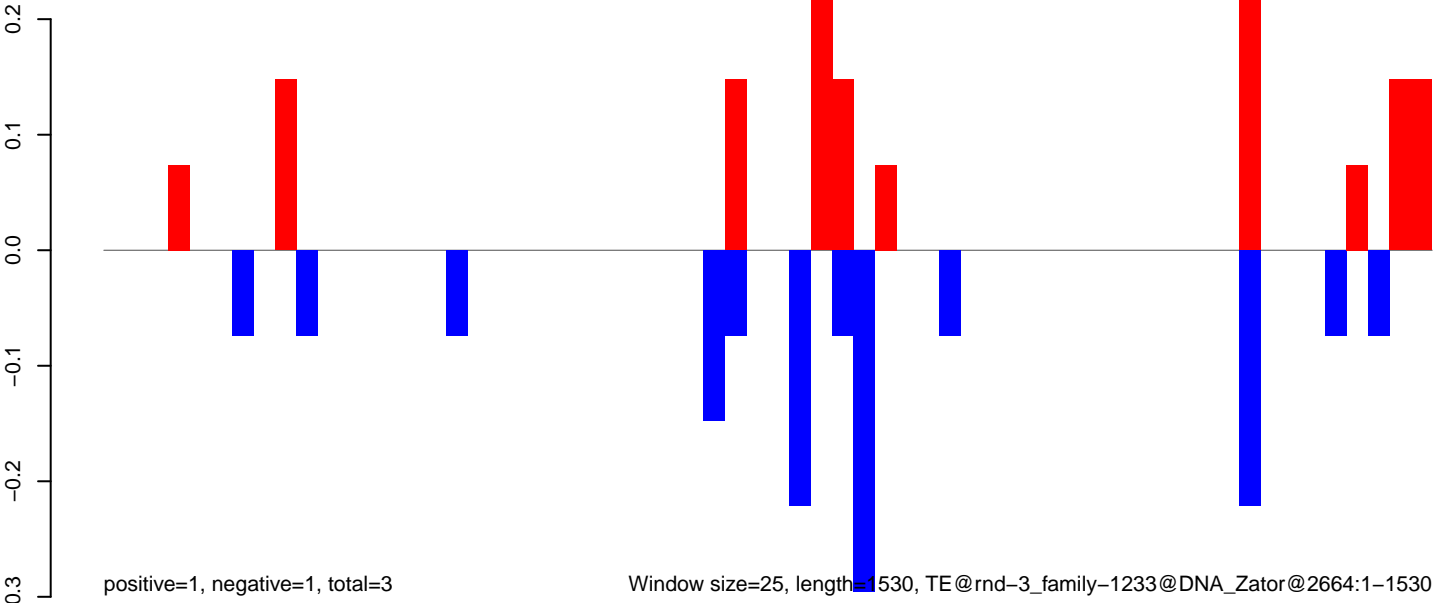
CuQuin_Testes_NL.18_23.rep



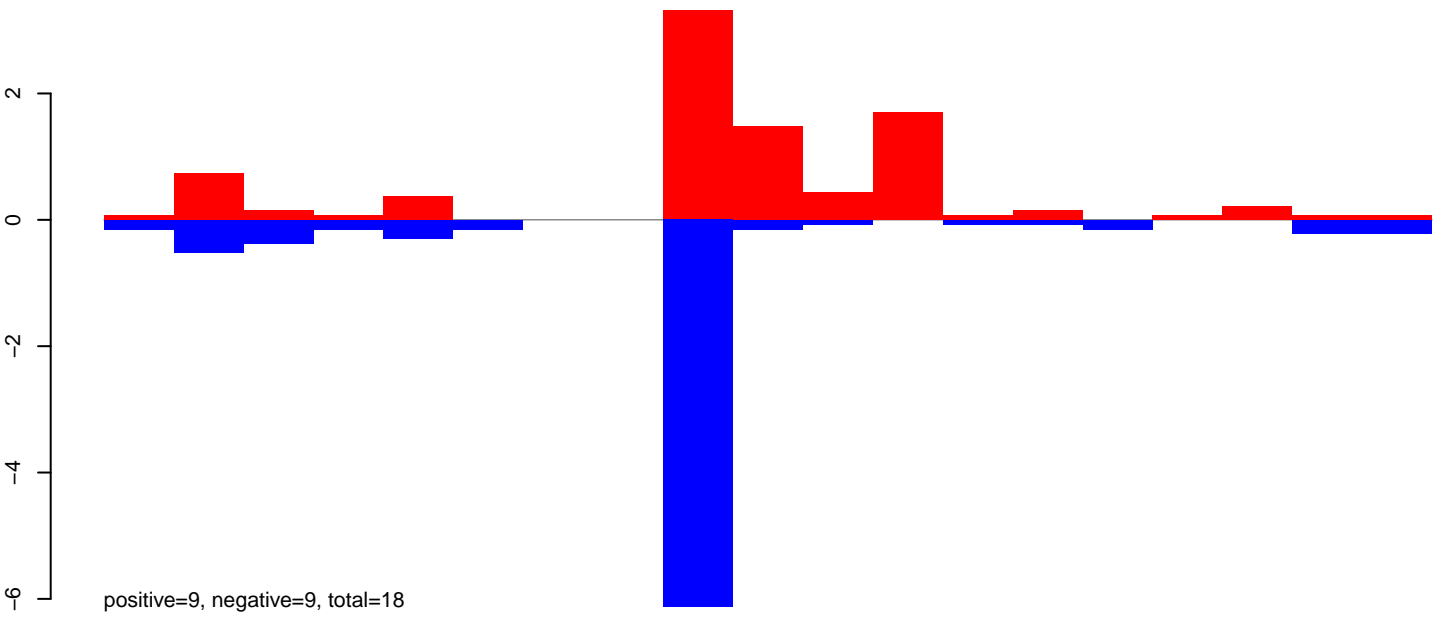
CuQuin_Testes_NL.24_35.rep



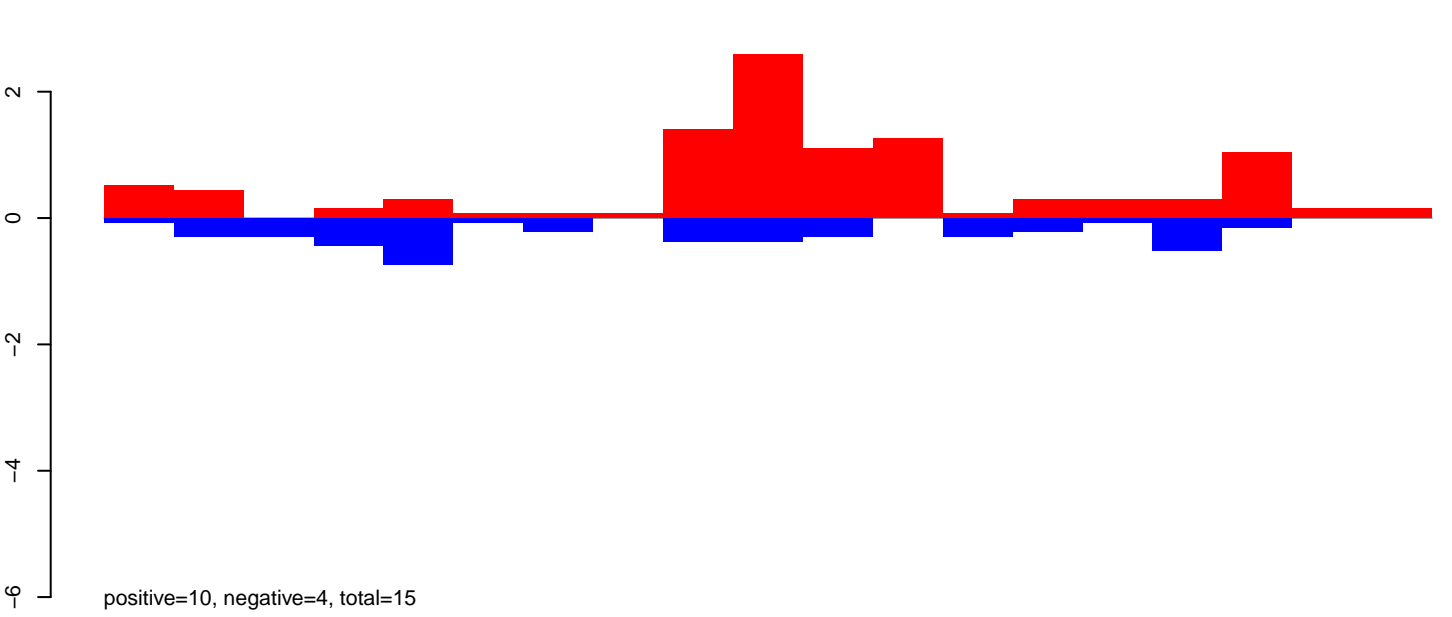
CuQuin_Testes_NL.rep



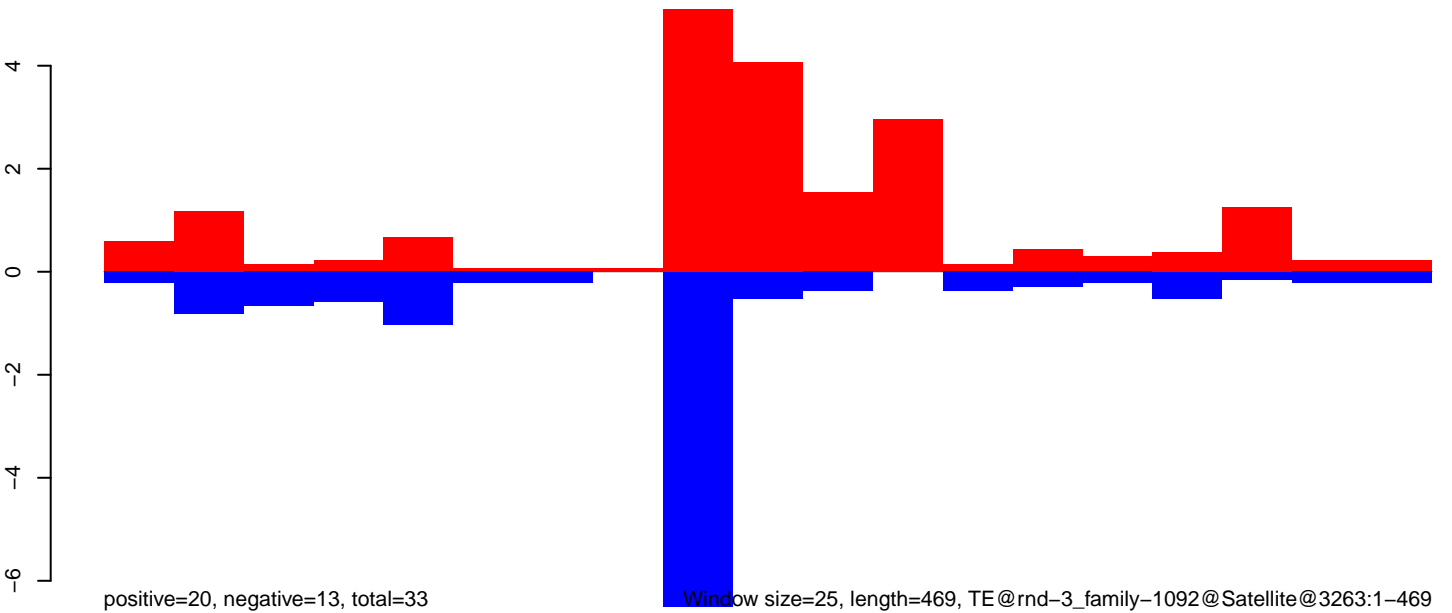
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

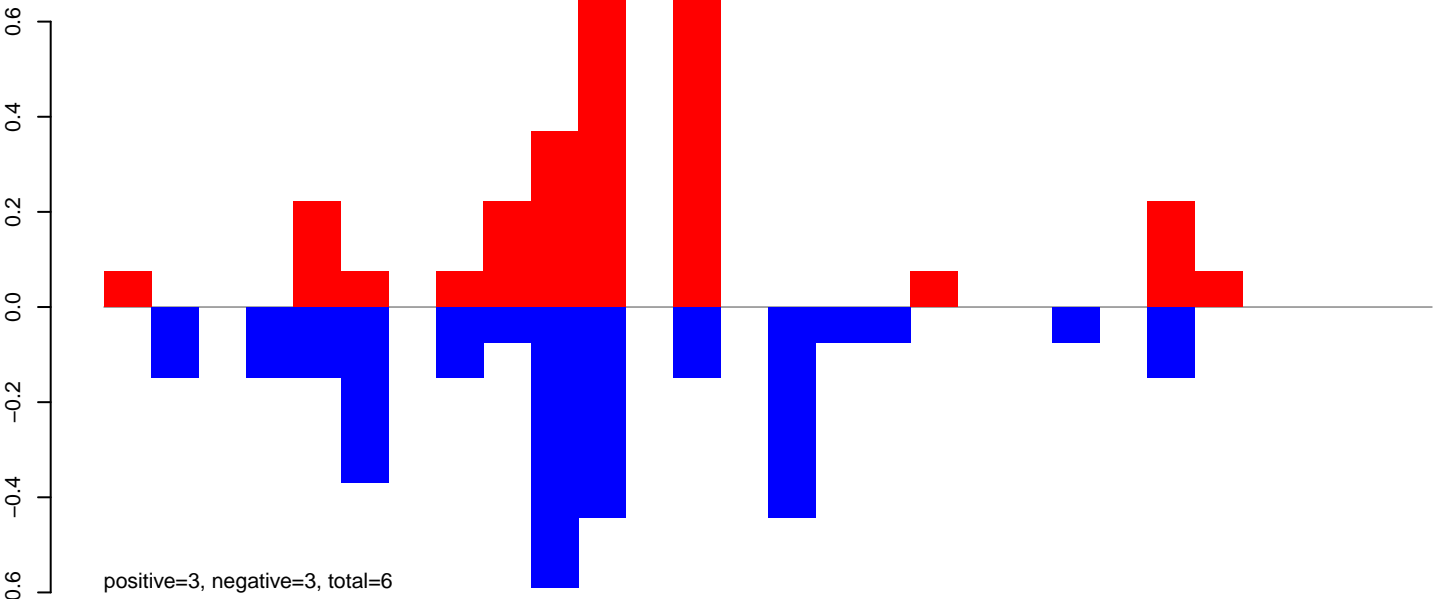


CuQuin_Testes_NL.rep

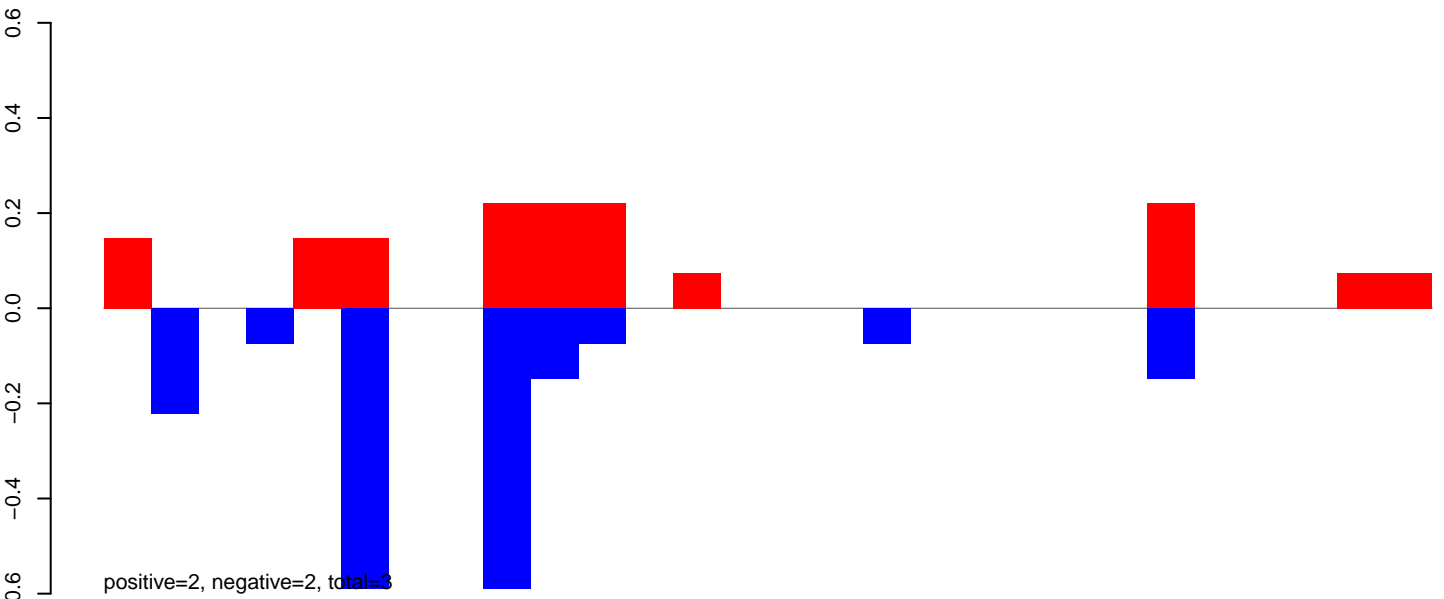


Window size=25, length=469, TE@rnd-3_family-1092@Satellite@3263:1-469

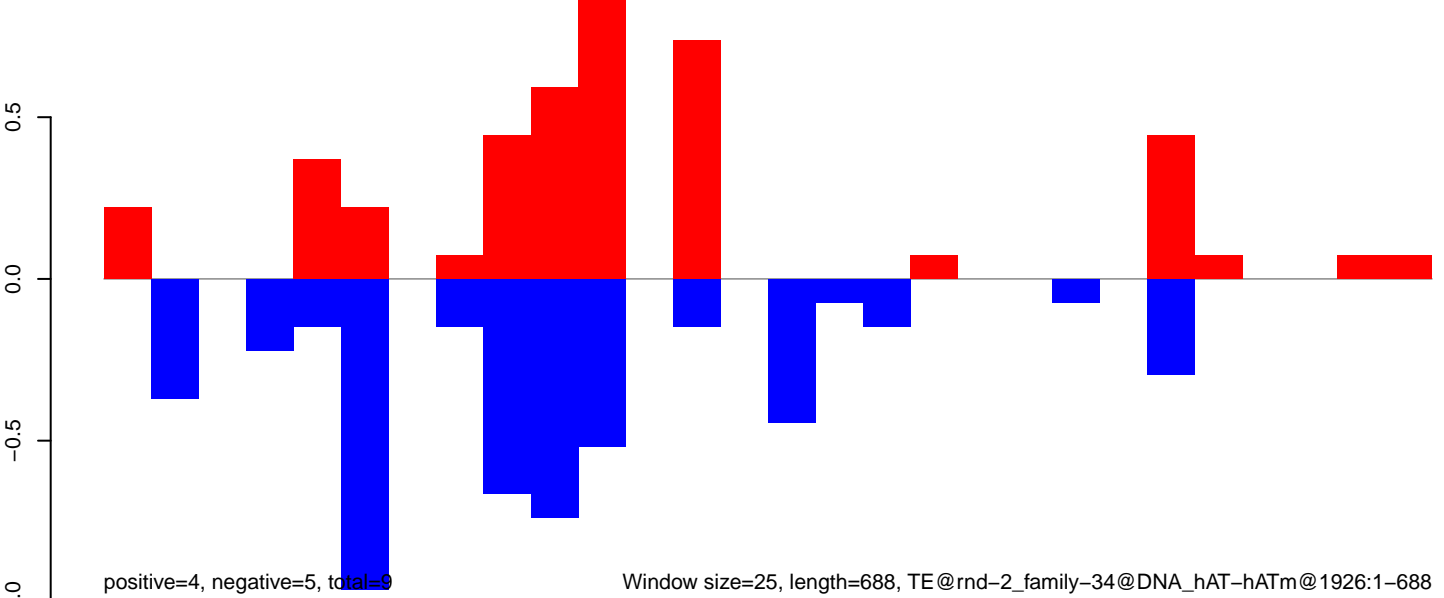
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

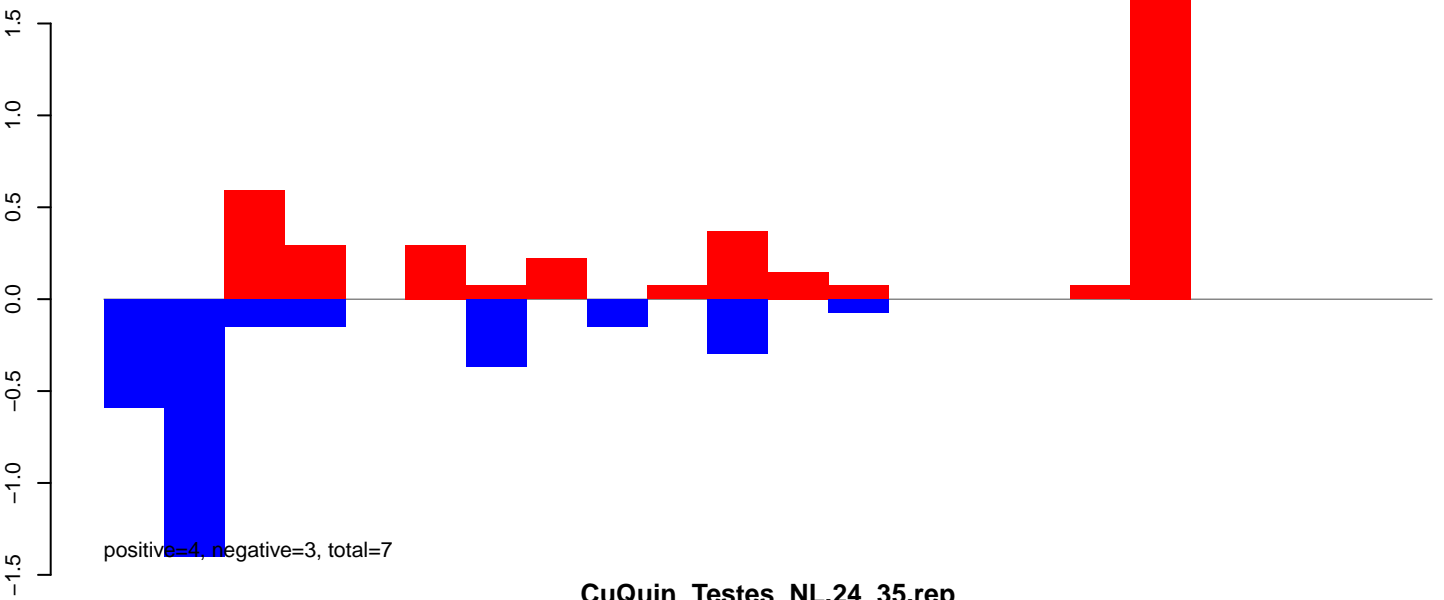


CuQuin_Testes_NL.rep

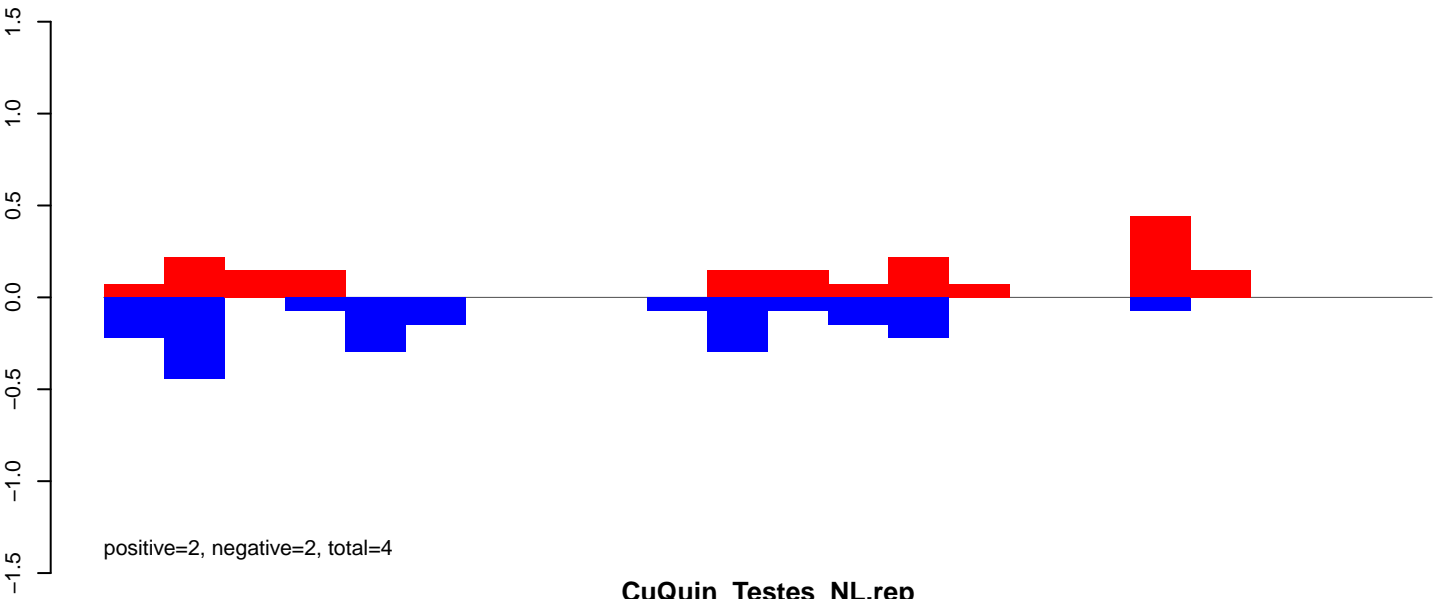


Window size=25, length=688, TE@rnd-2_family-34@DNA_hAT-hATm@1926:1-688

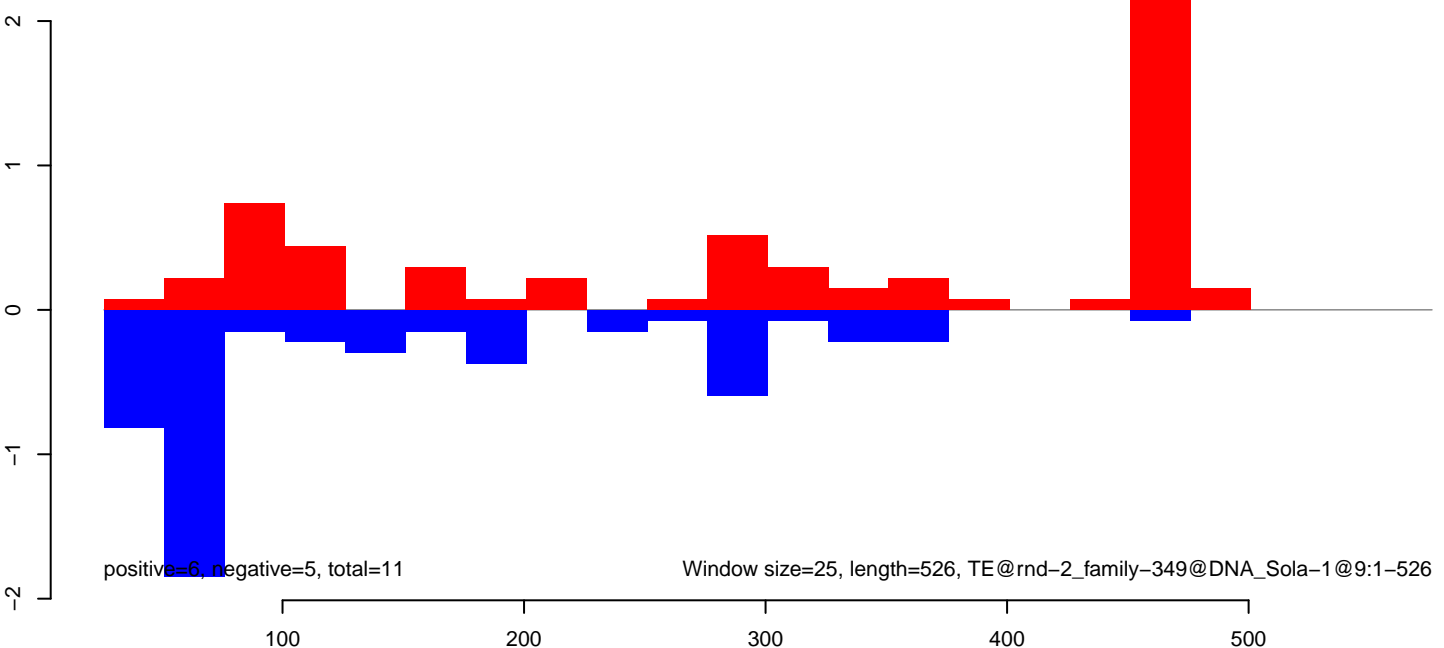
CuQuin_Testes_NL.18_23.rep



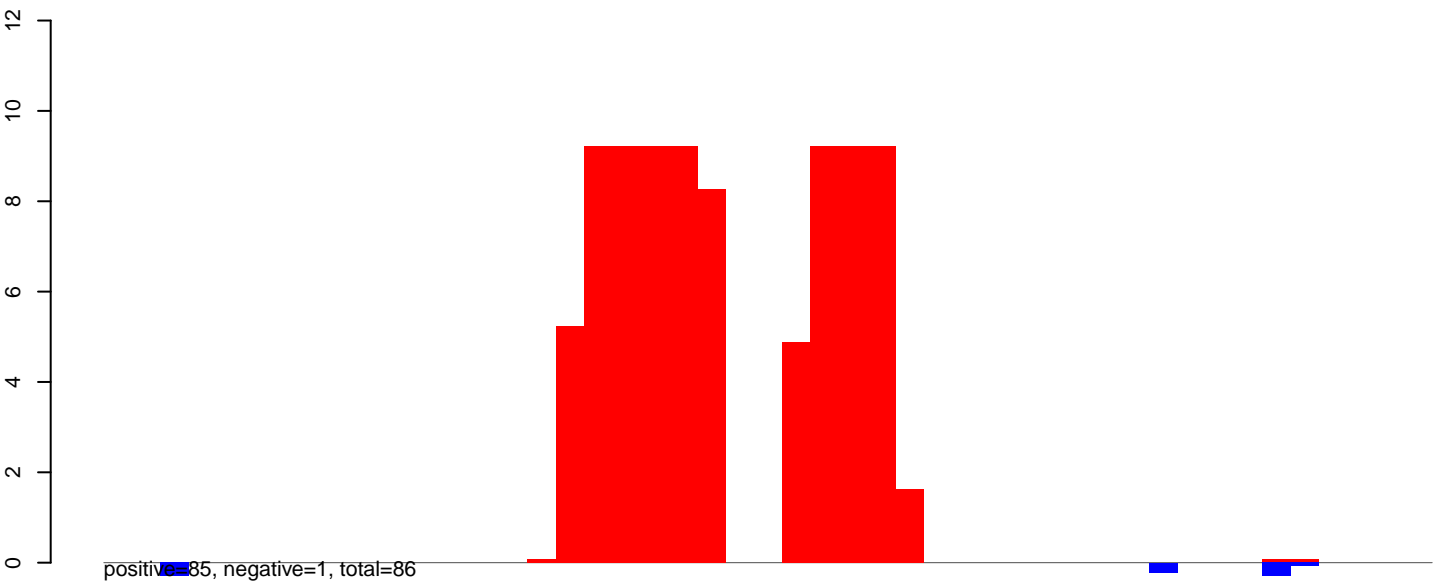
CuQuin_Testes_NL.24_35.rep



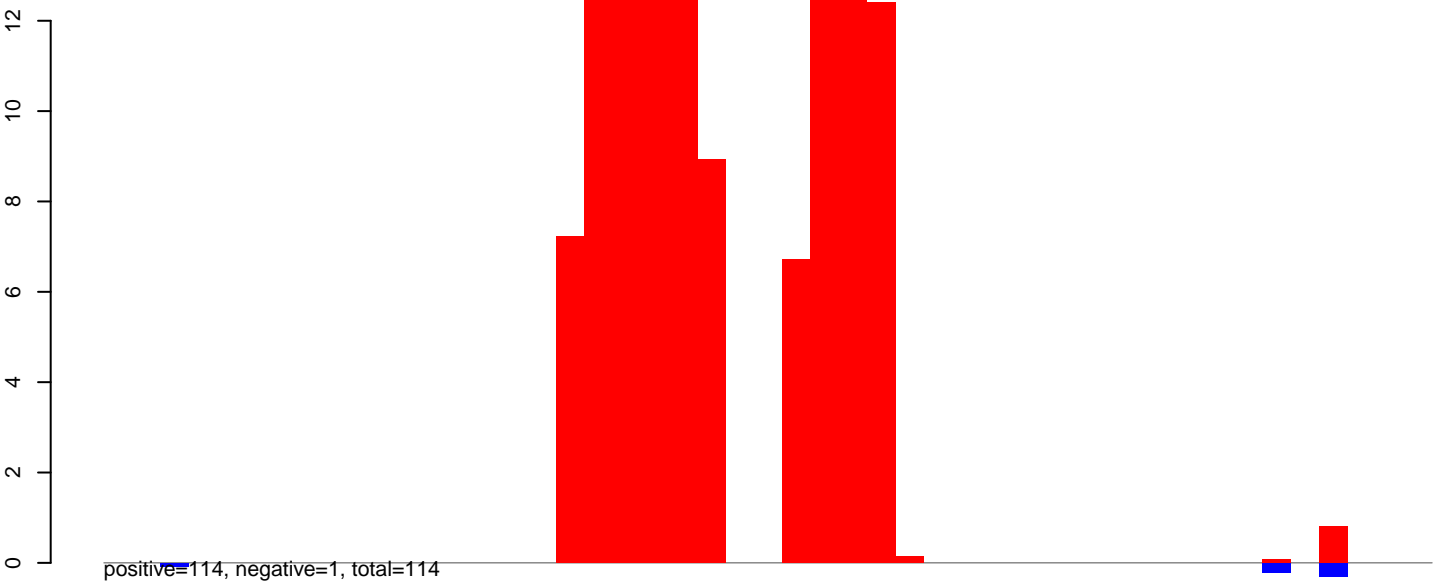
CuQuin_Testes_NL.rep



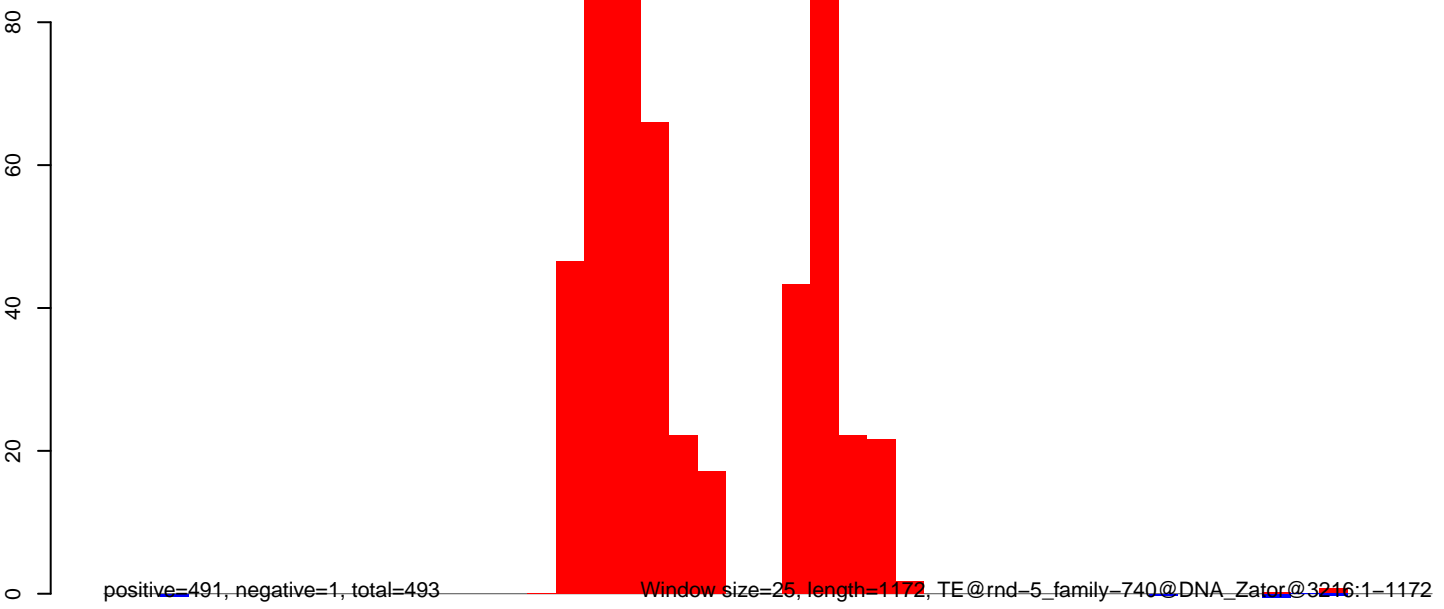
CuQuin_Testes_NL.18_23.rep



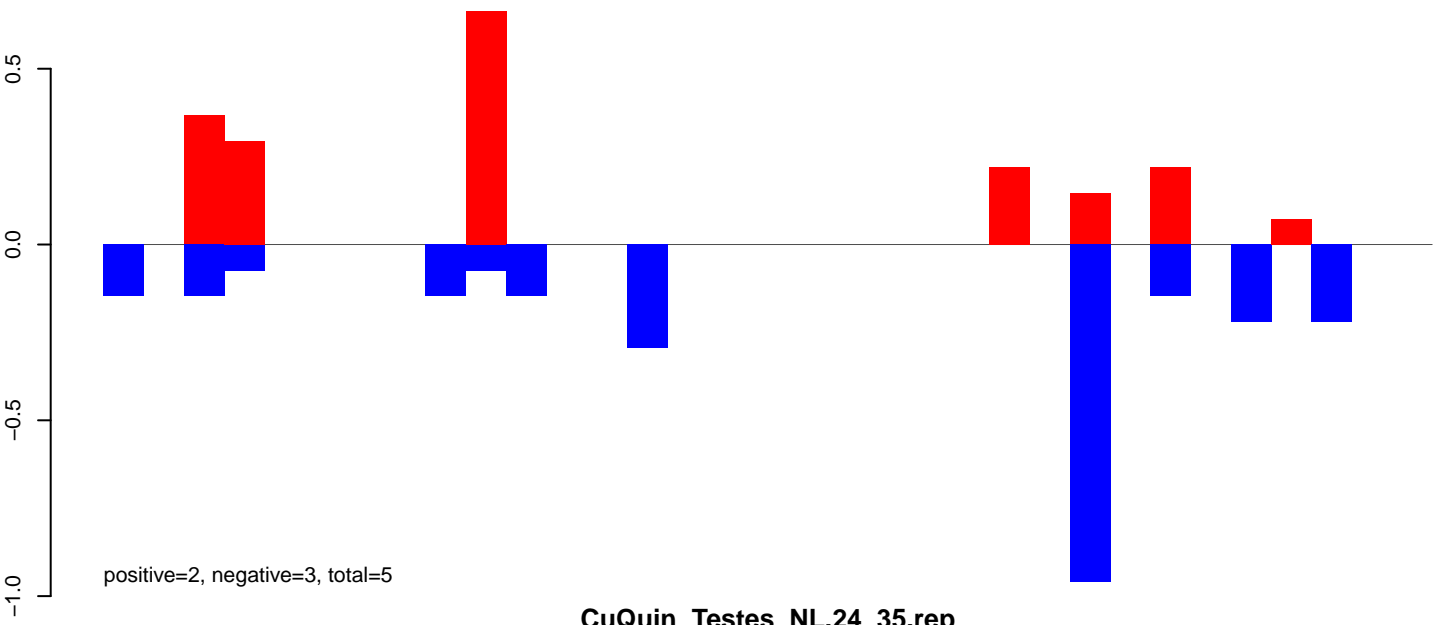
CuQuin_Testes_NL.24_35.rep



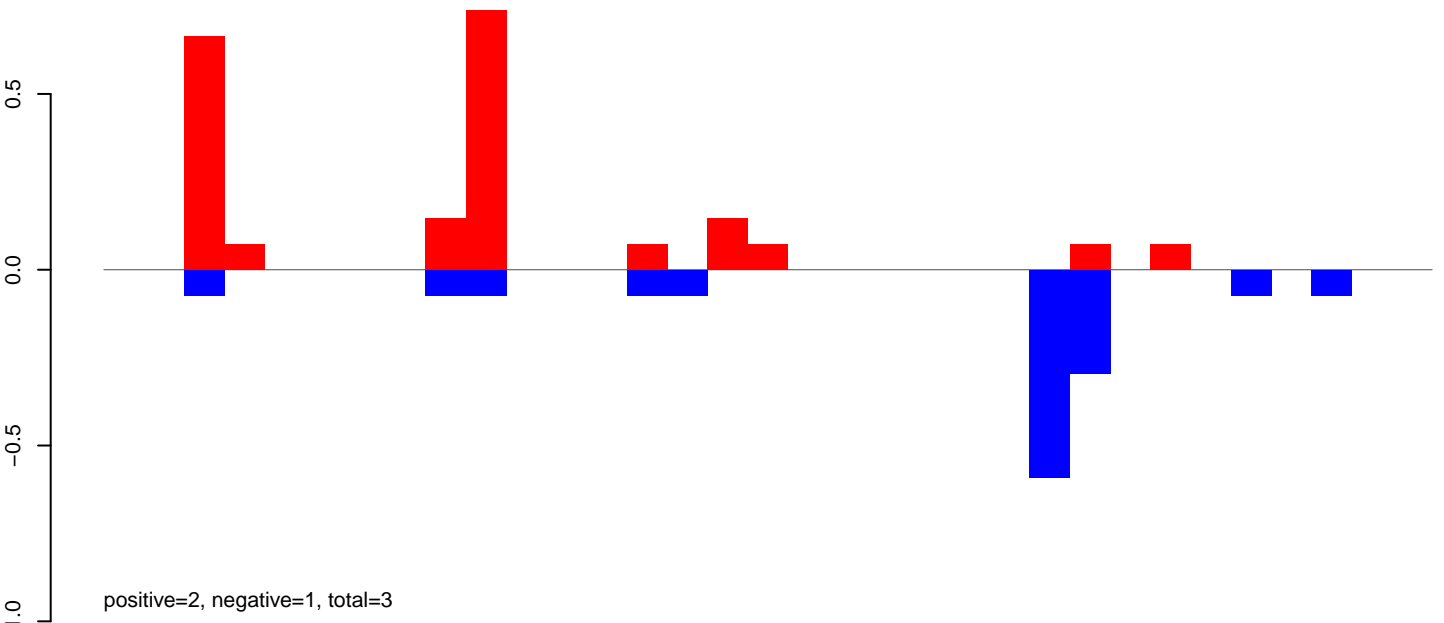
CuQuin_Testes_NL.rep



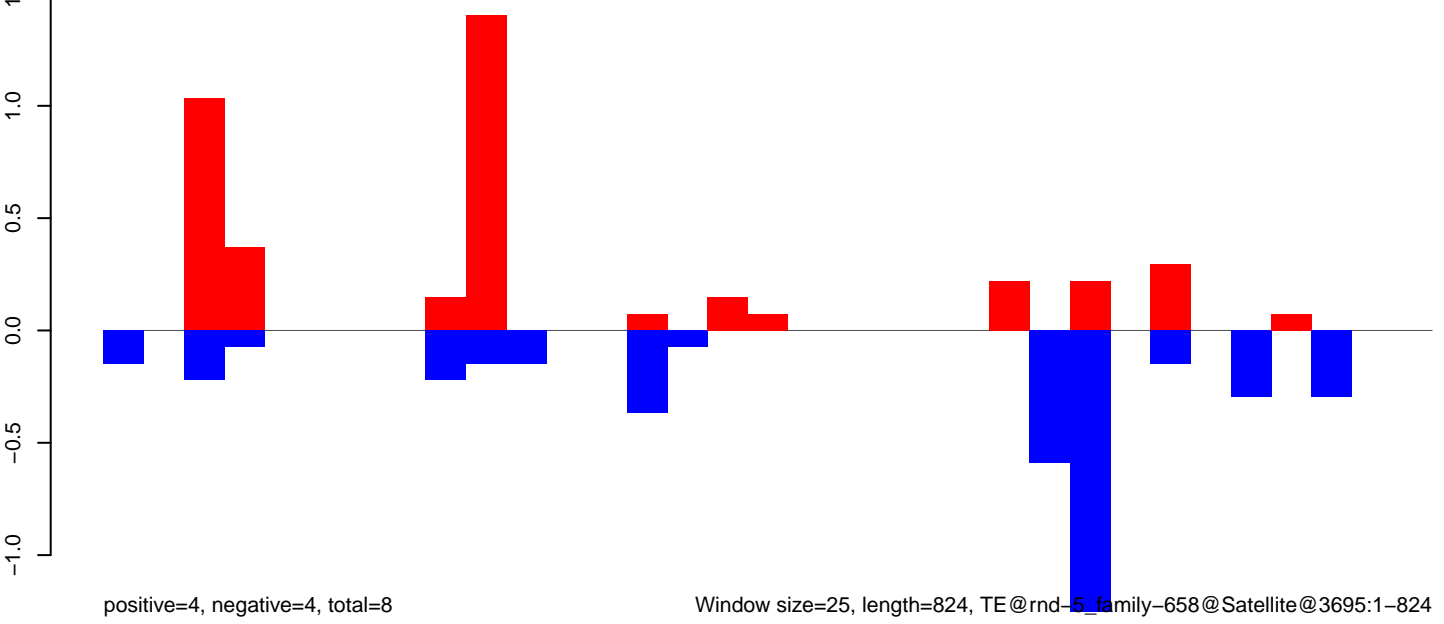
CuQuin_Testes_NL.18_23.rep



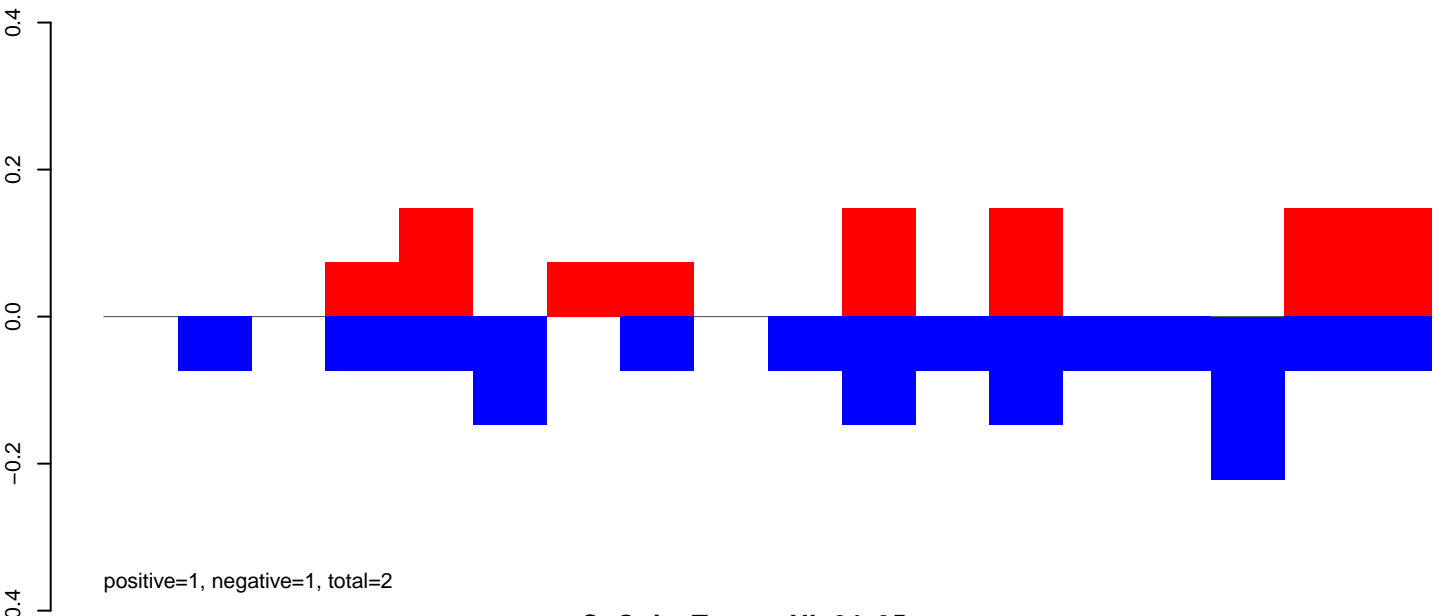
CuQuin_Testes_NL.24_35.rep



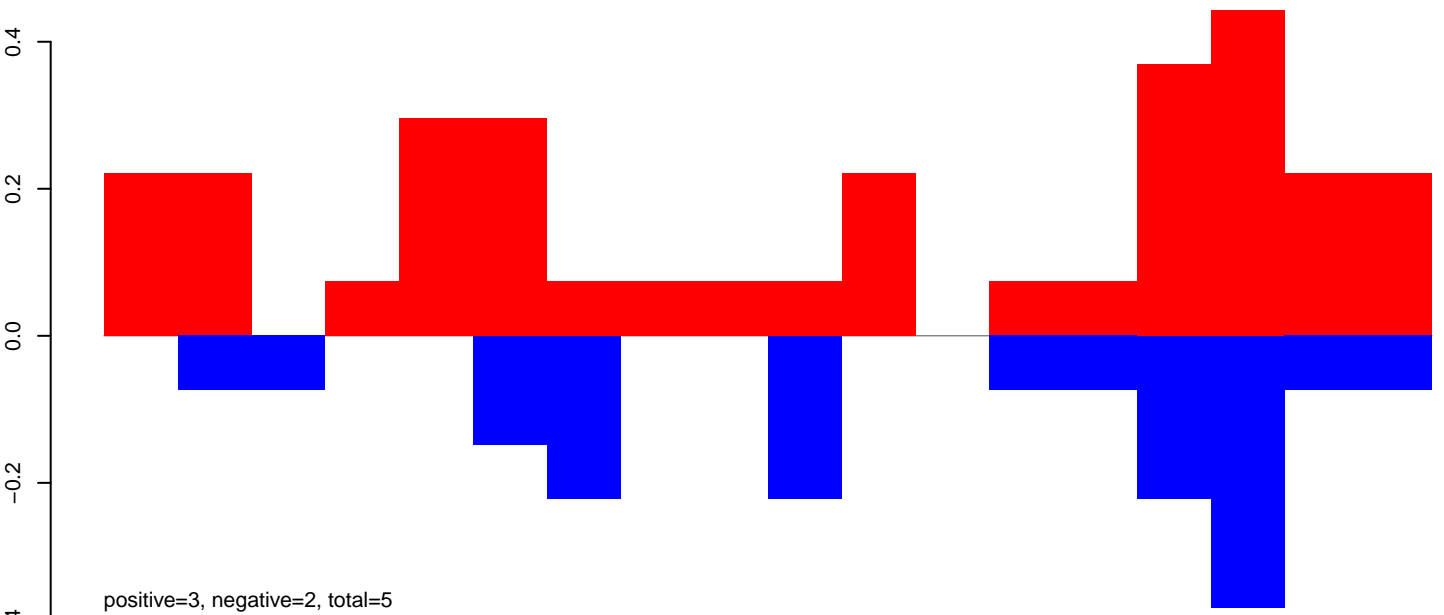
CuQuin_Testes_NL.rep



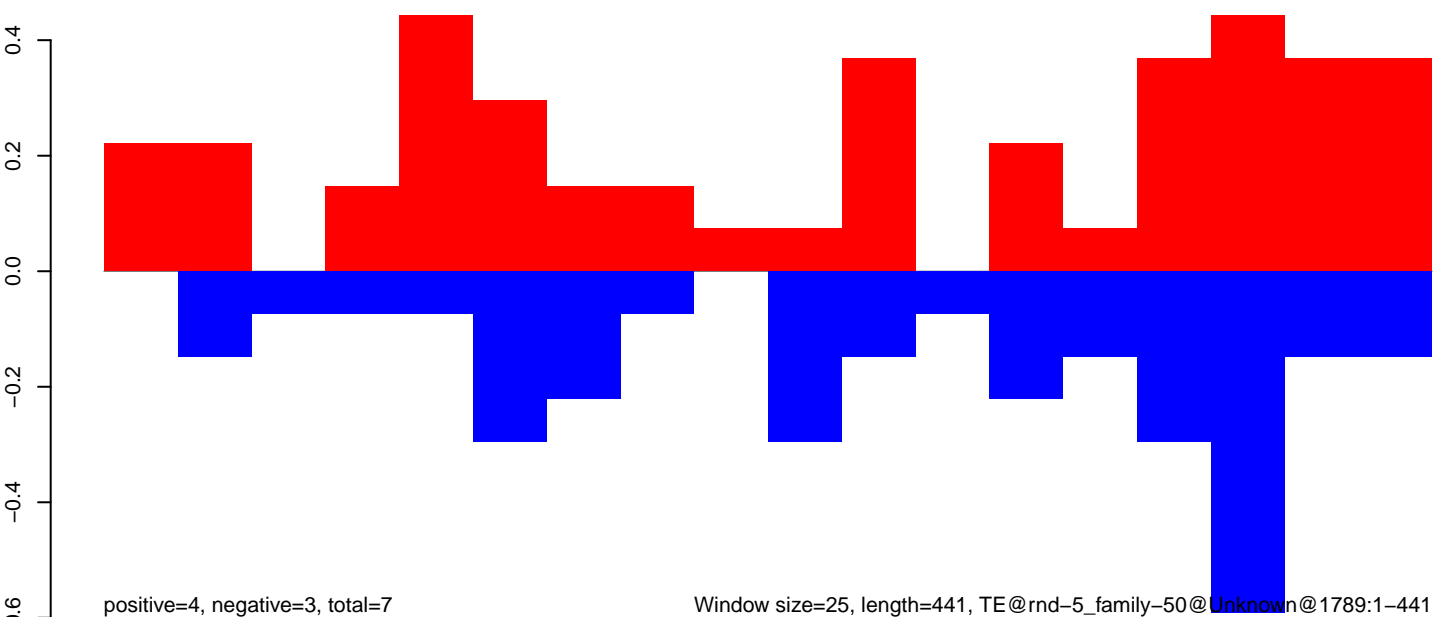
CuQuin_Testes_NL.18_23.rep



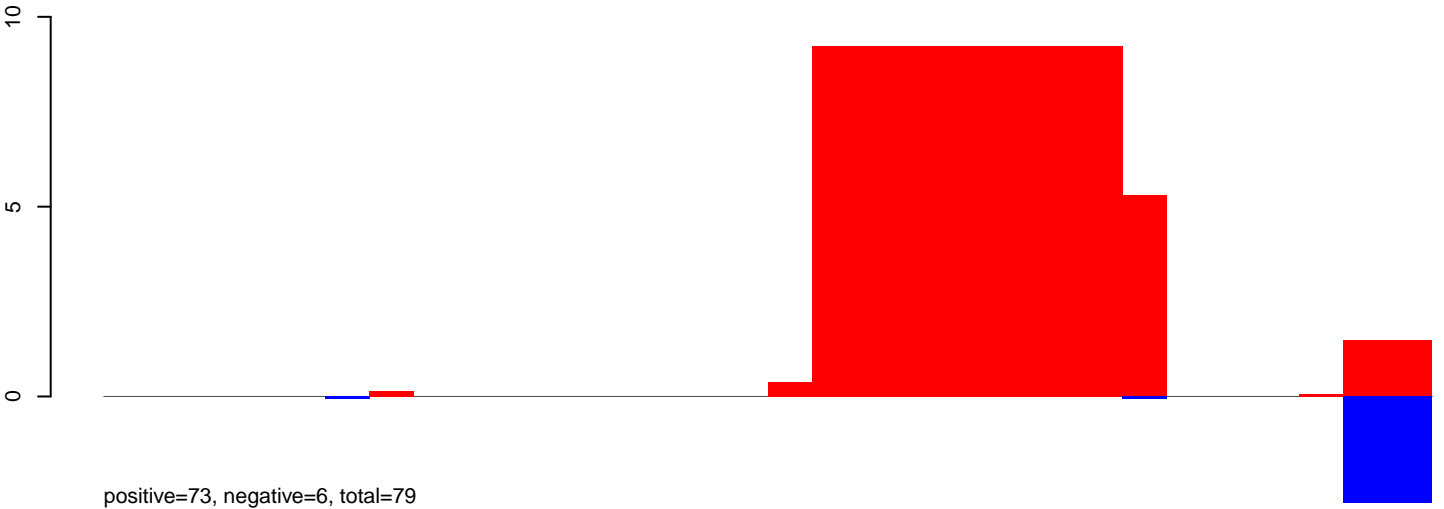
CuQuin_Testes_NL.24_35.rep



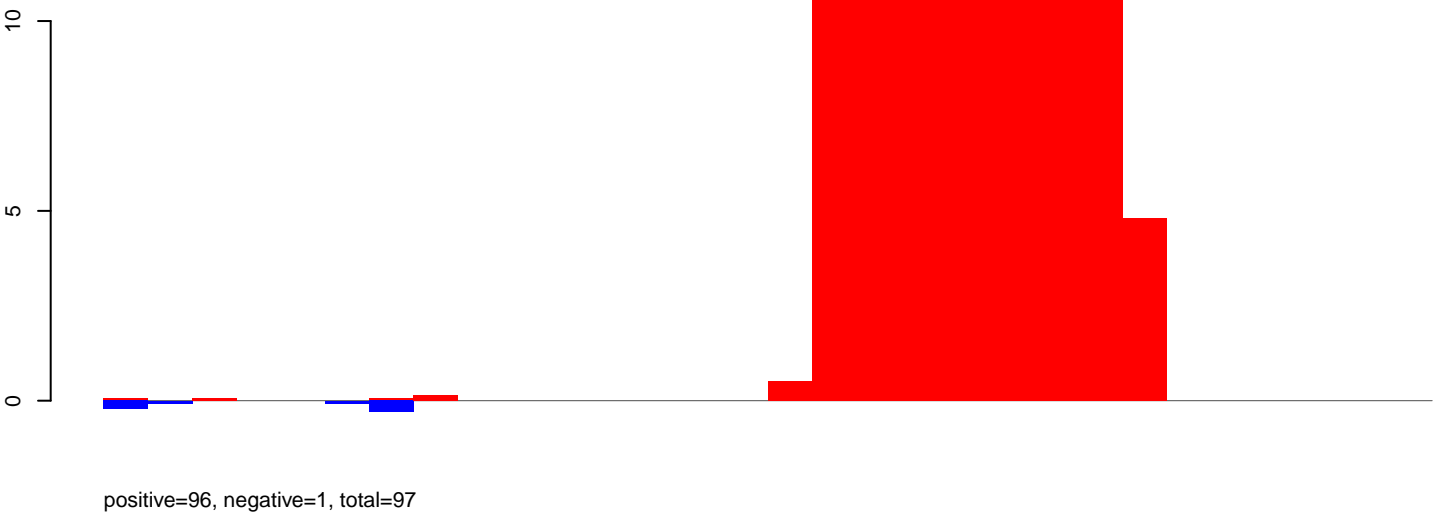
CuQuin_Testes_NL.rep



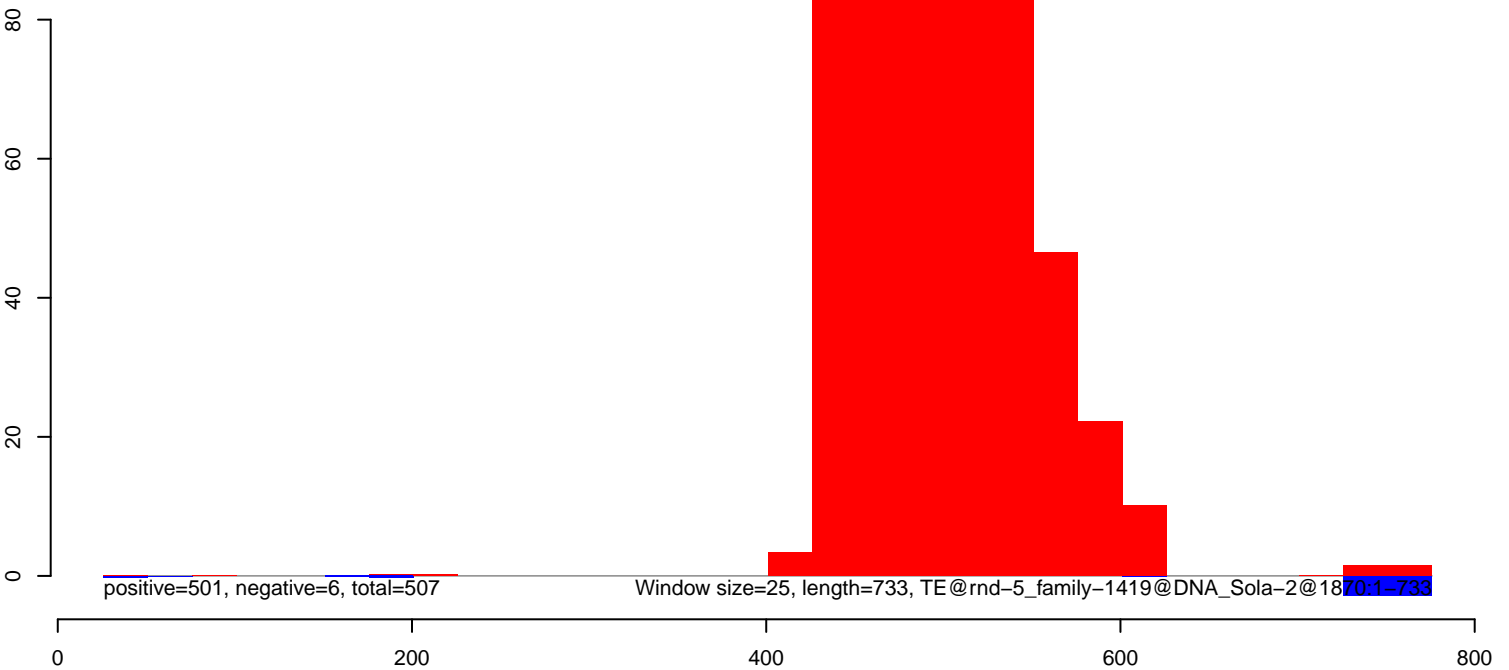
CuQuin_Testes_NL.18_23.rep



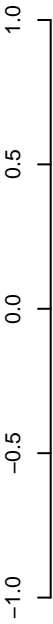
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

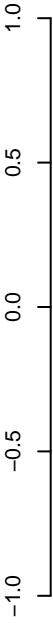


CuQuin_Testes_NL.18_23.rep



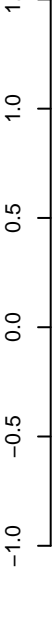
positive=3, negative=1, total=4

CuQuin_Testes_NL.24_35.rep



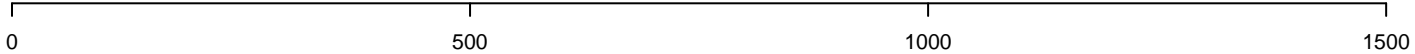
positive=3, negative=3, total=5

CuQuin_Testes_NL.rep

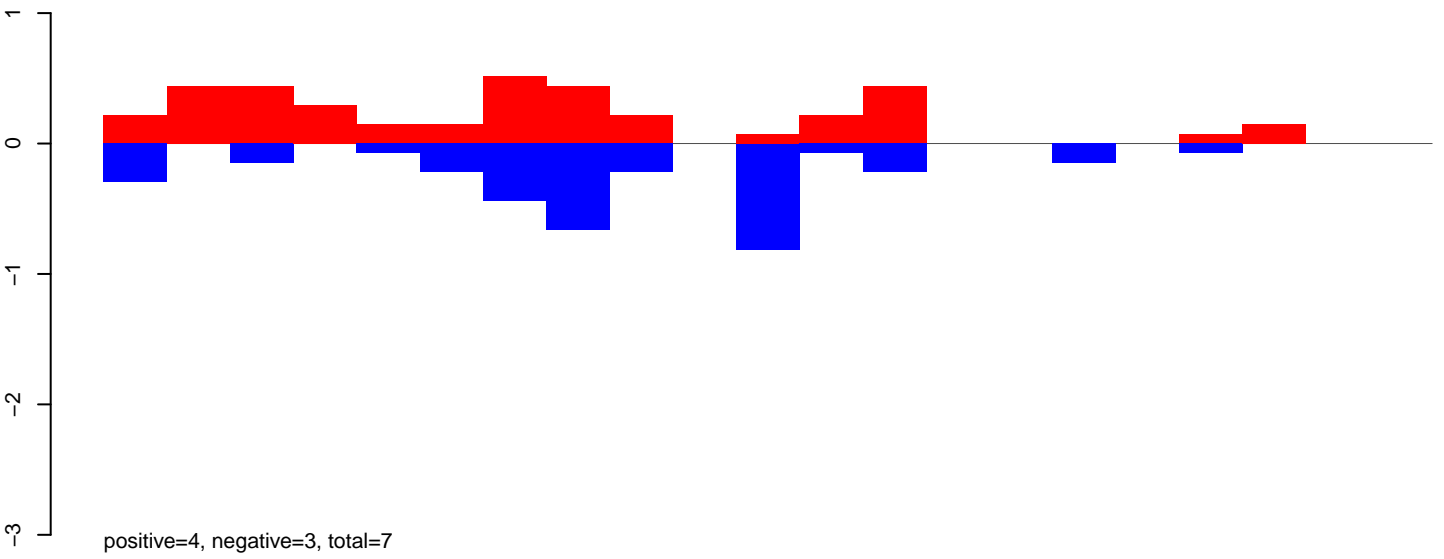


positive=5, negative=4, total=9

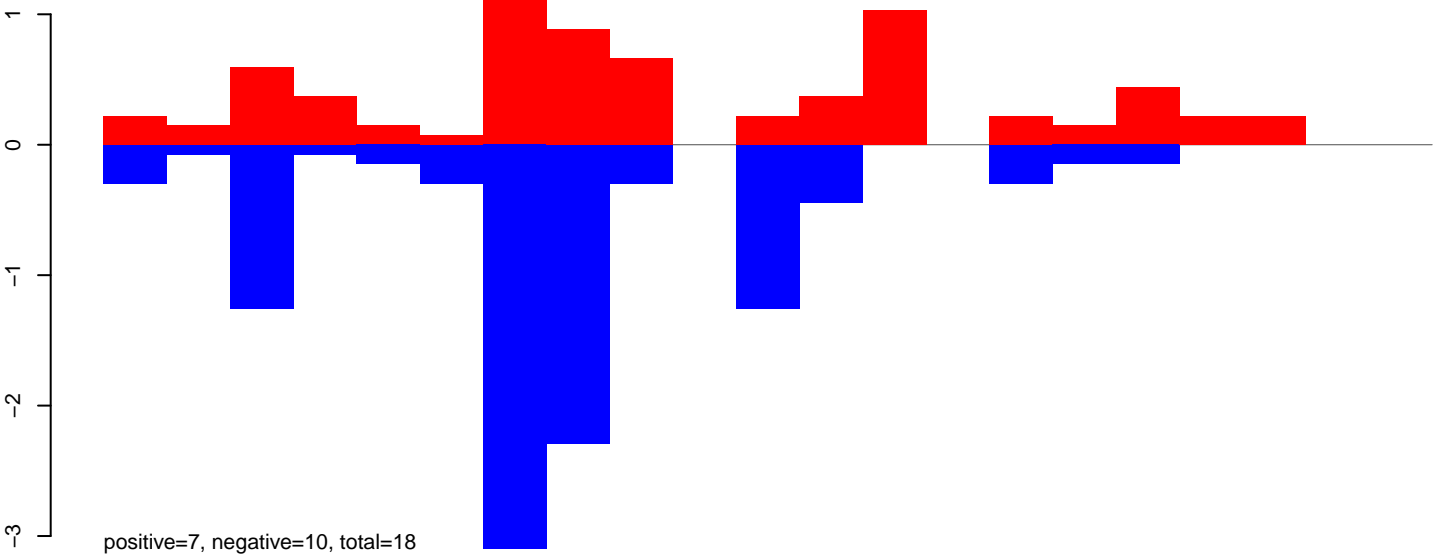
Window size=25, length=1444, TE@rnd-5_family-127@DNA_Ginger@1115:1-1444



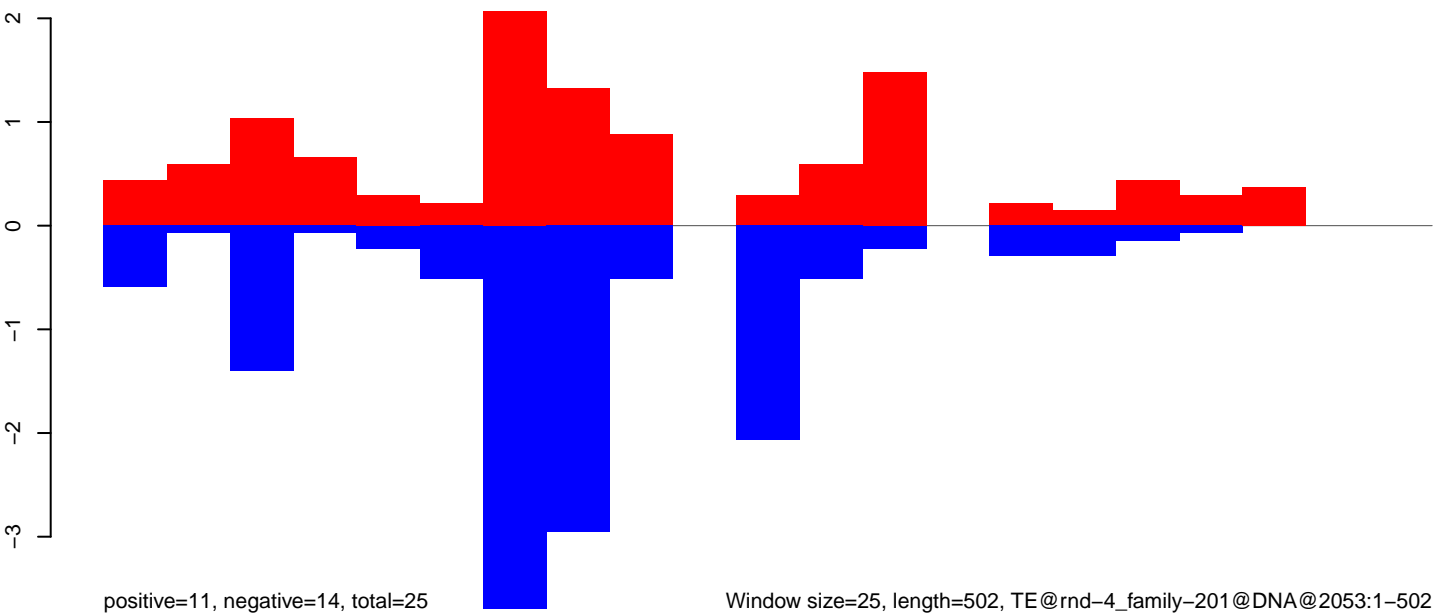
CuQuin_Testes_NL.18_23.rep



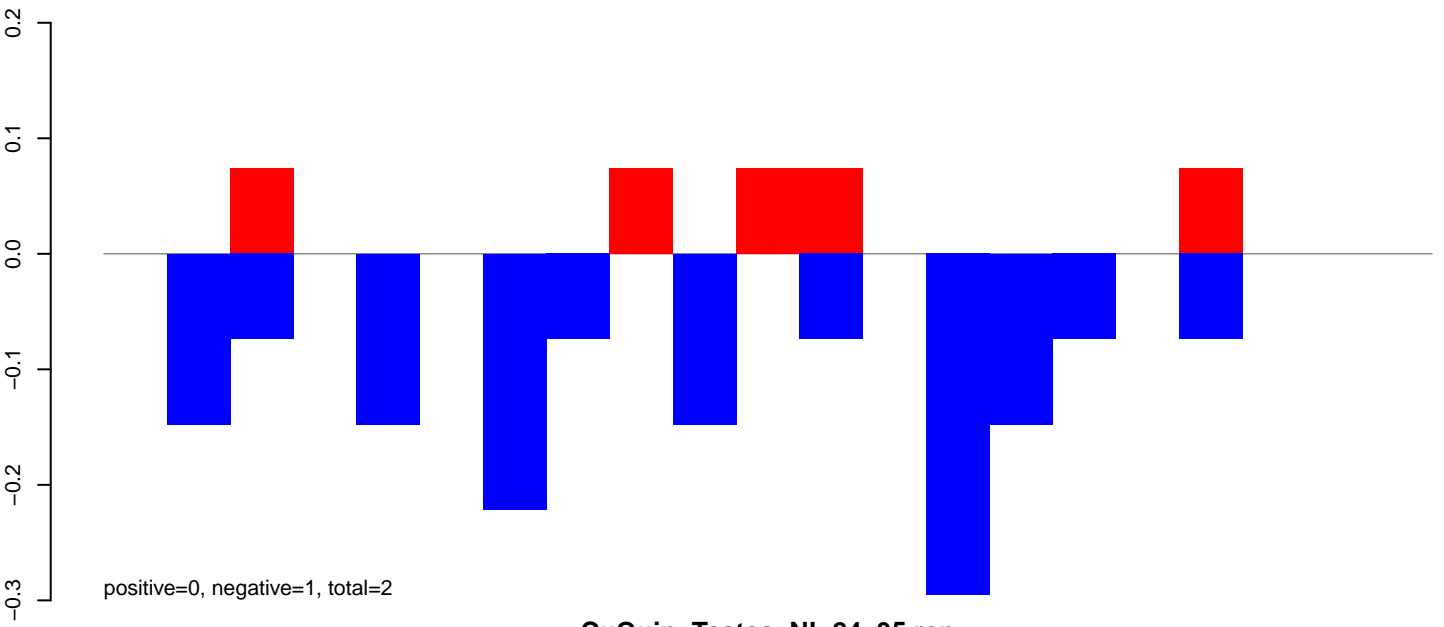
CuQuin_Testes_NL.24_35.rep



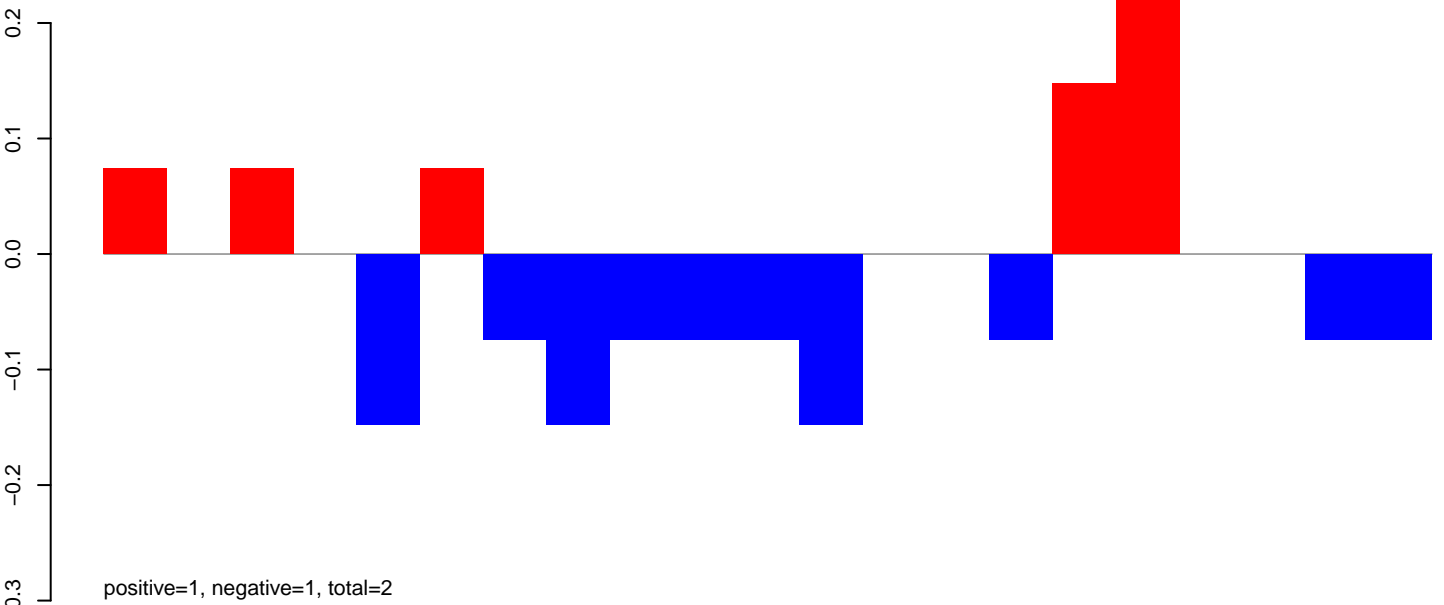
CuQuin_Testes_NL.rep



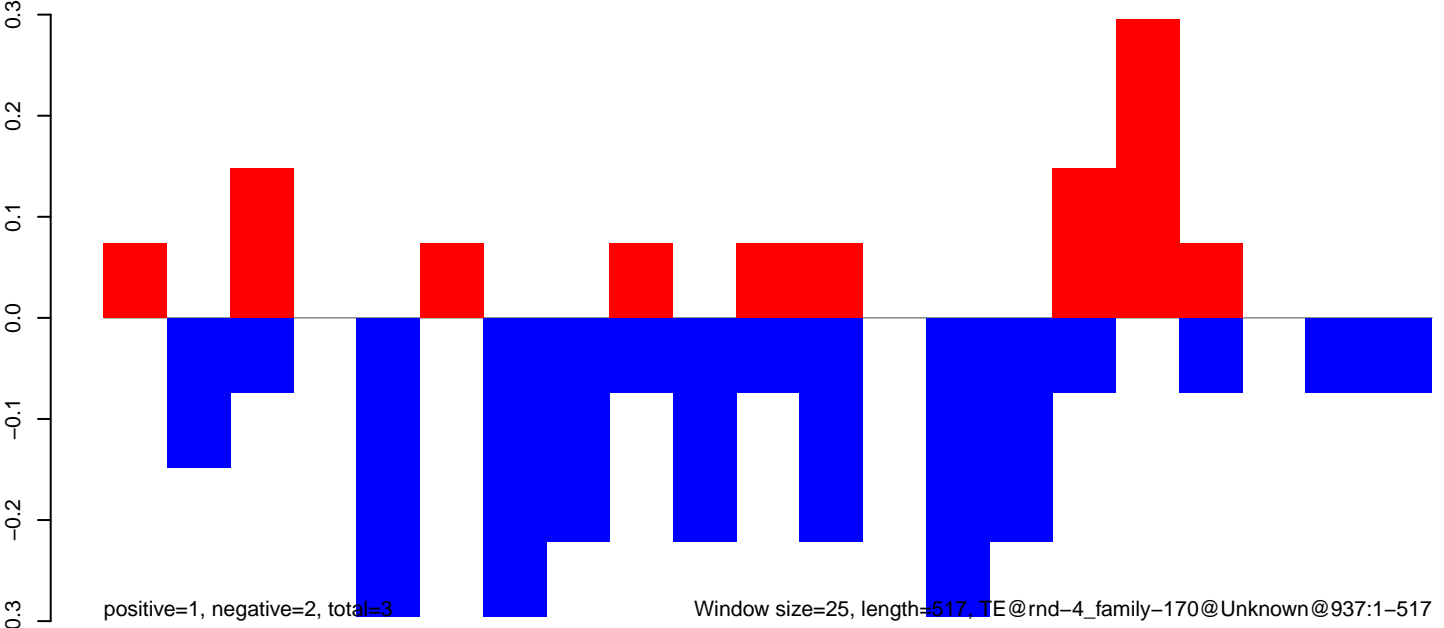
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

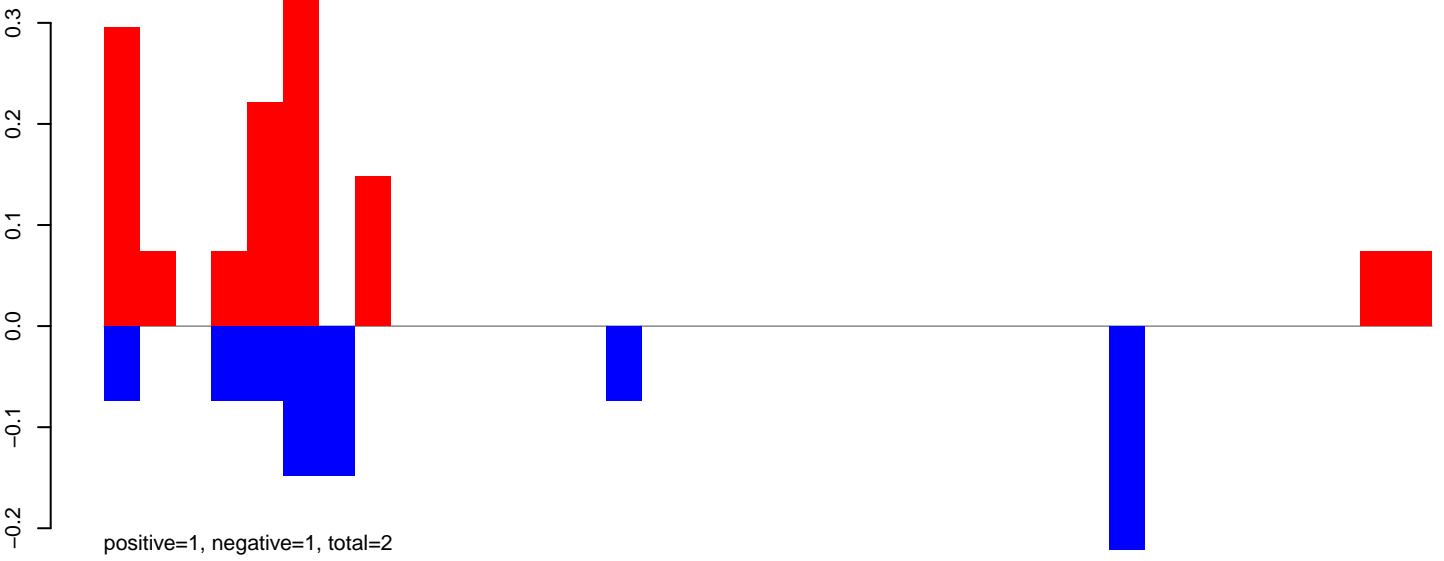


CuQuin_Testes_NL.rep

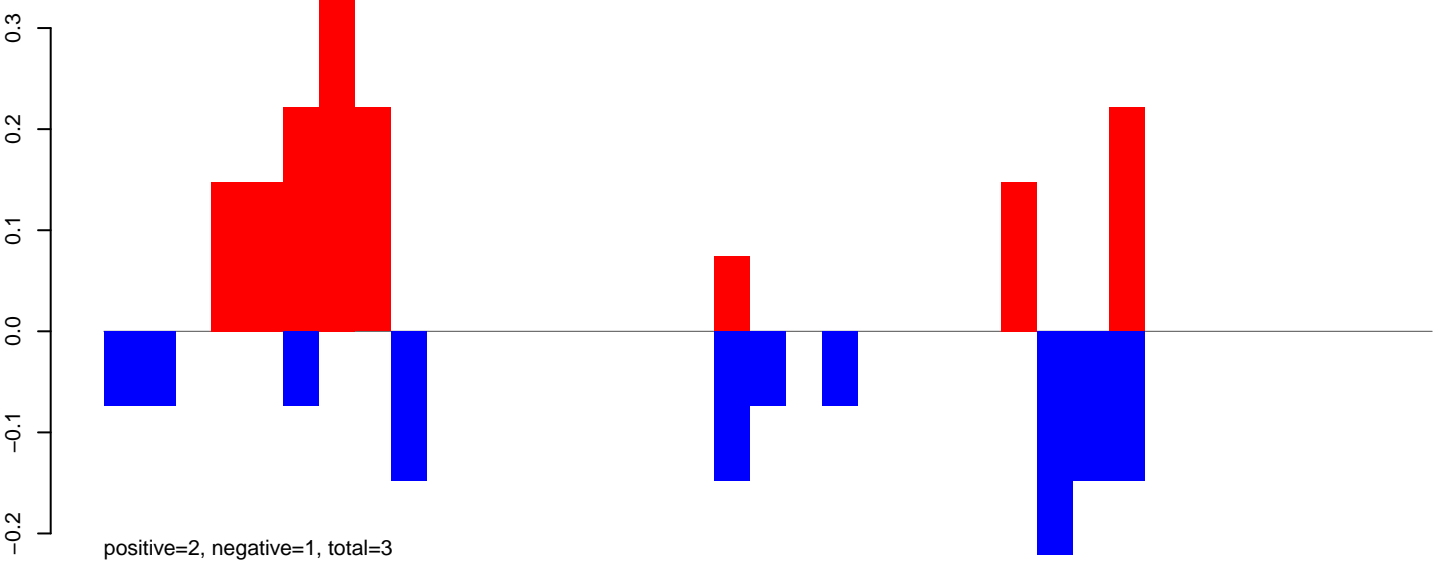


Window size=25, length=517, TE@rnd-4_family-170@Unknown@937:1-517

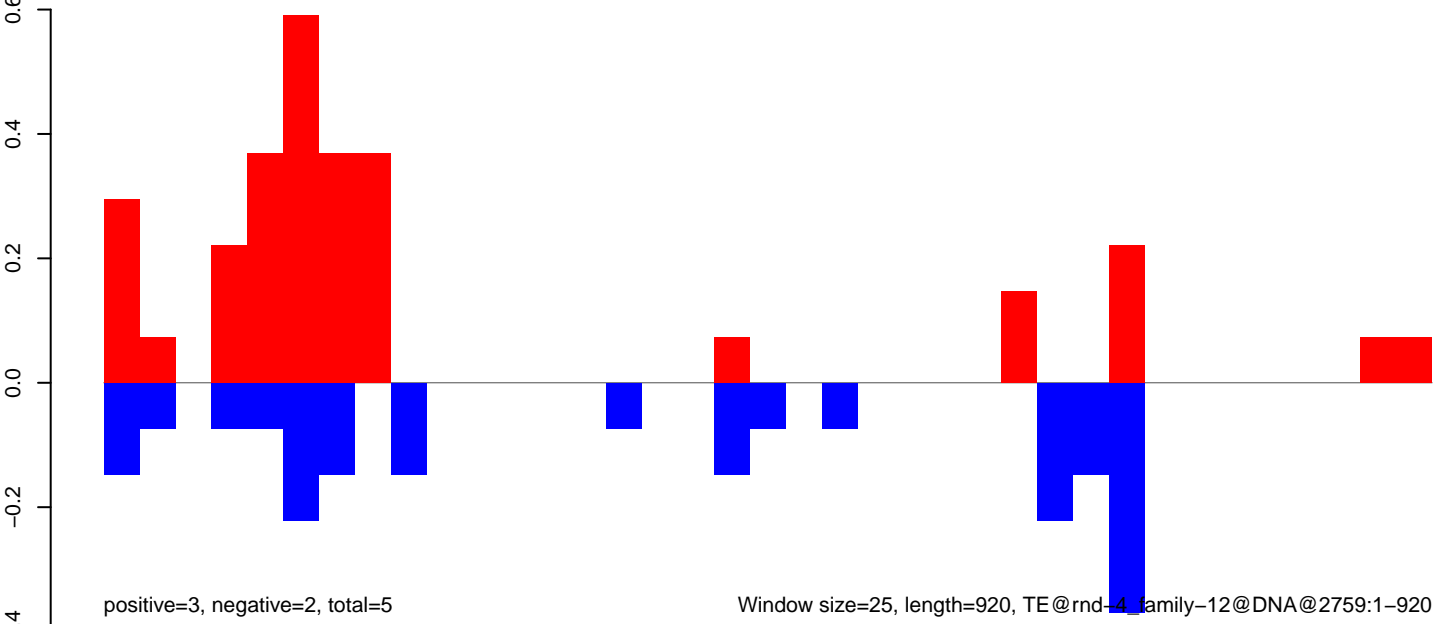
CuQuin_Testes_NL.18_23.rep



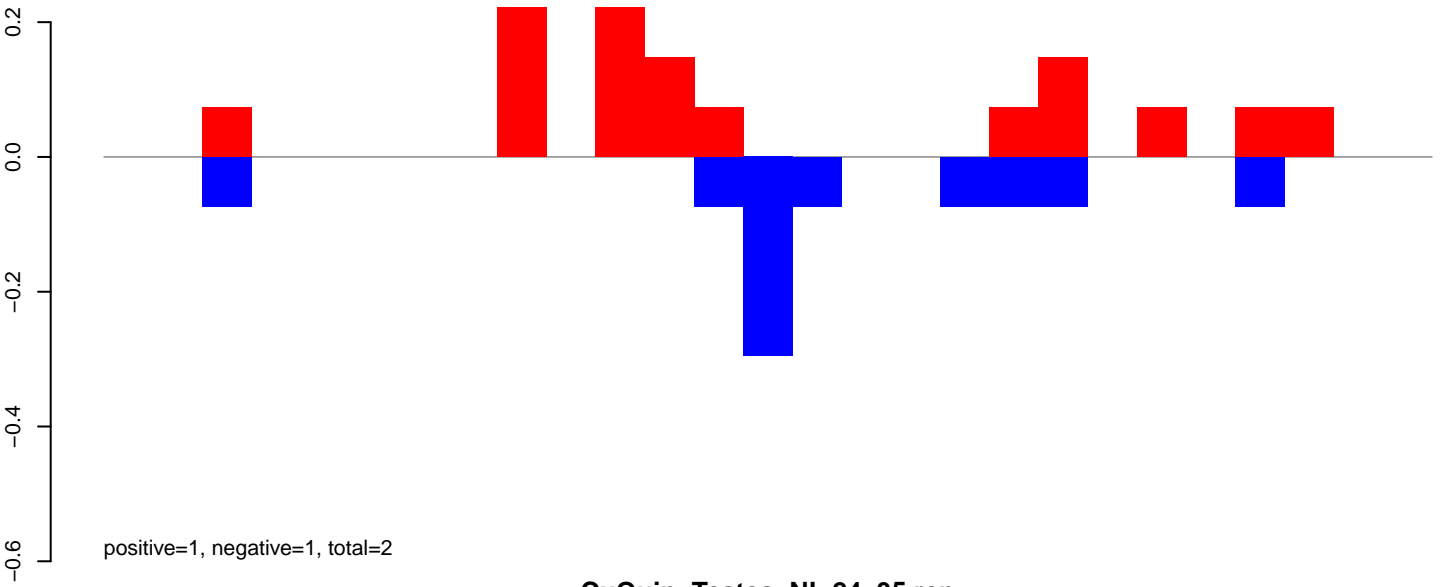
CuQuin_Testes_NL.24_35.rep



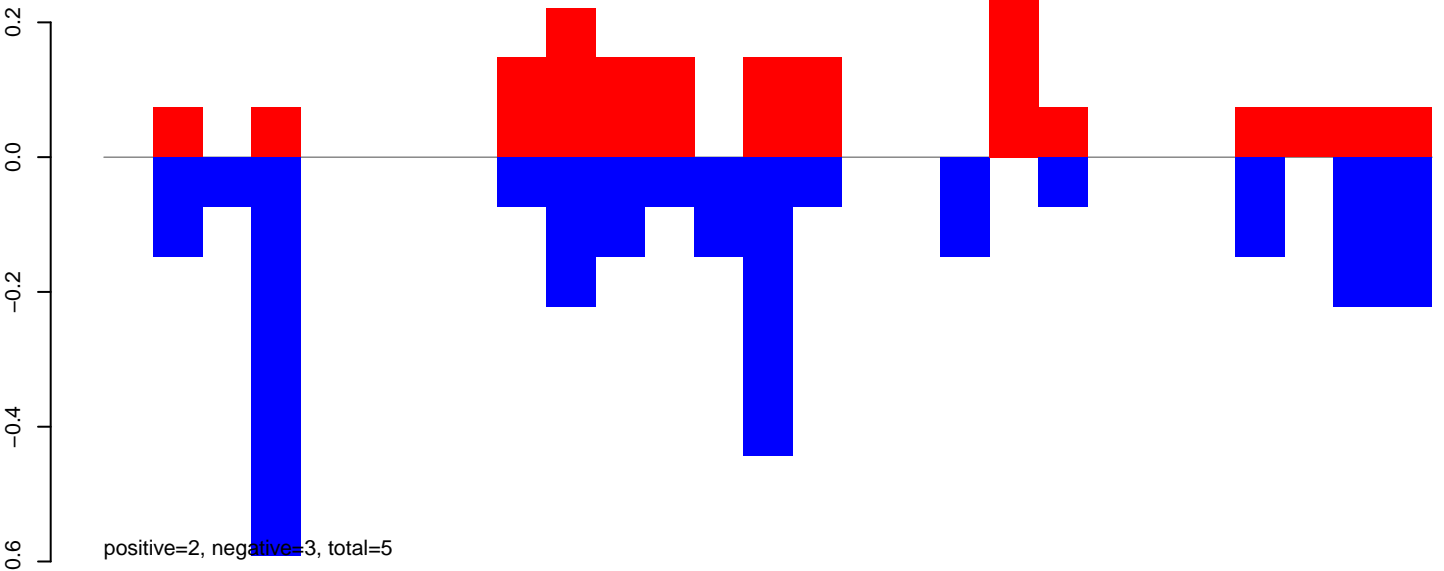
CuQuin_Testes_NL.rep



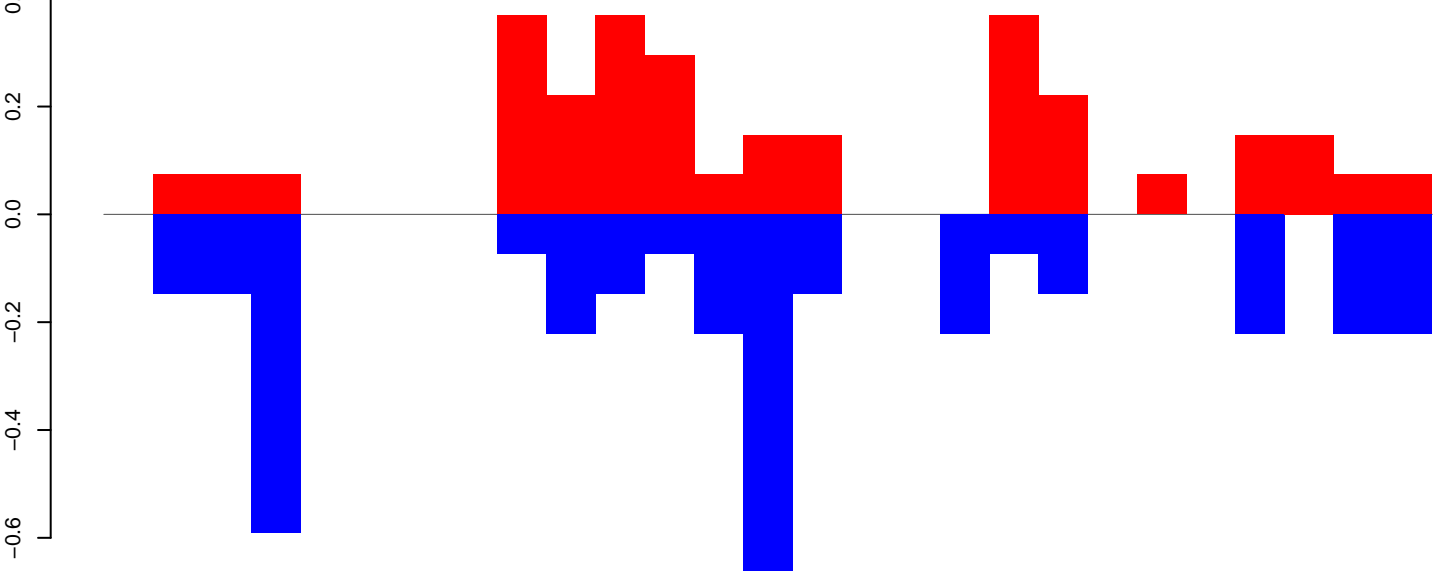
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



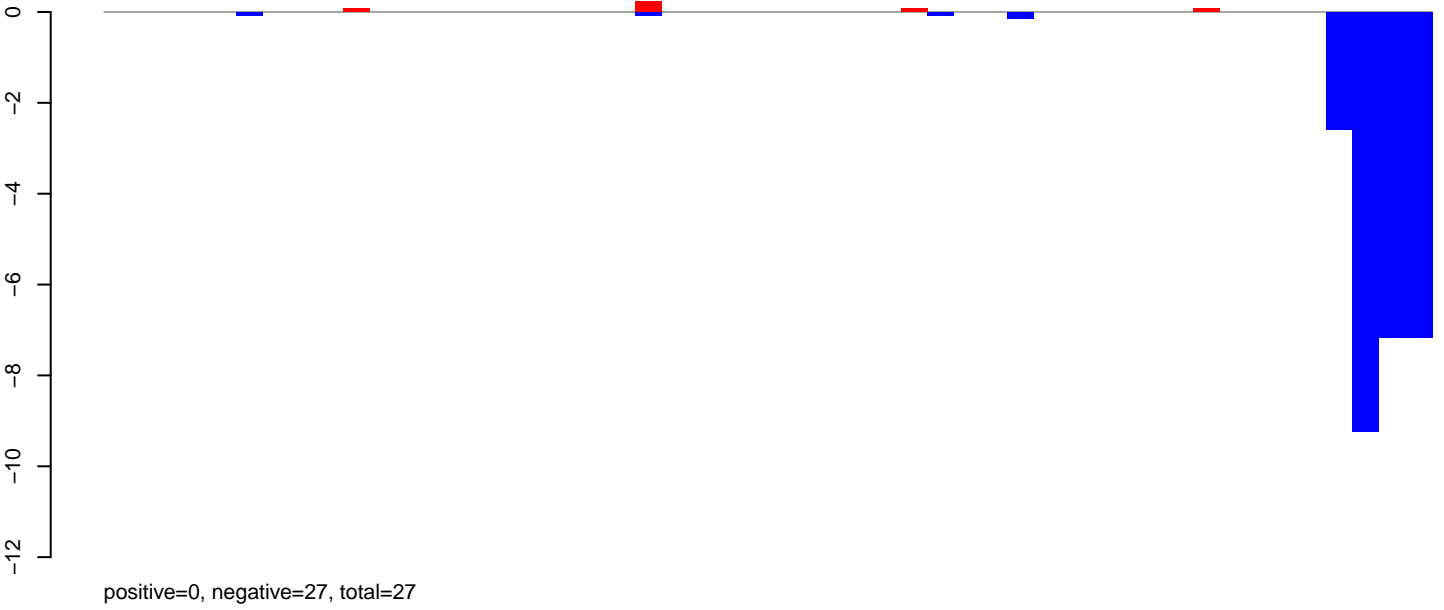
CuQuin_Testes_NL.rep



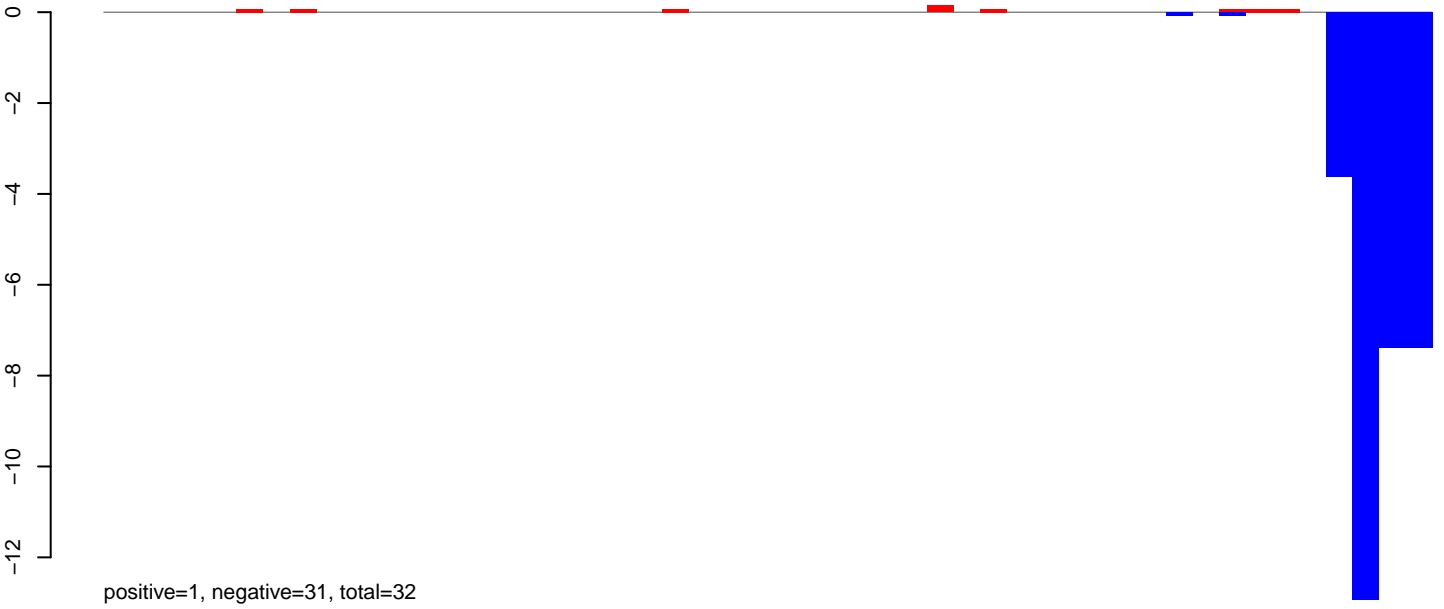
Window size=25, length=671, TE@rnd-2_family-601@LTR_Gypsy@2588:1-671

0 100 200 300 400 500 600 700

CuQuin_Testes_NL.18_23.rep



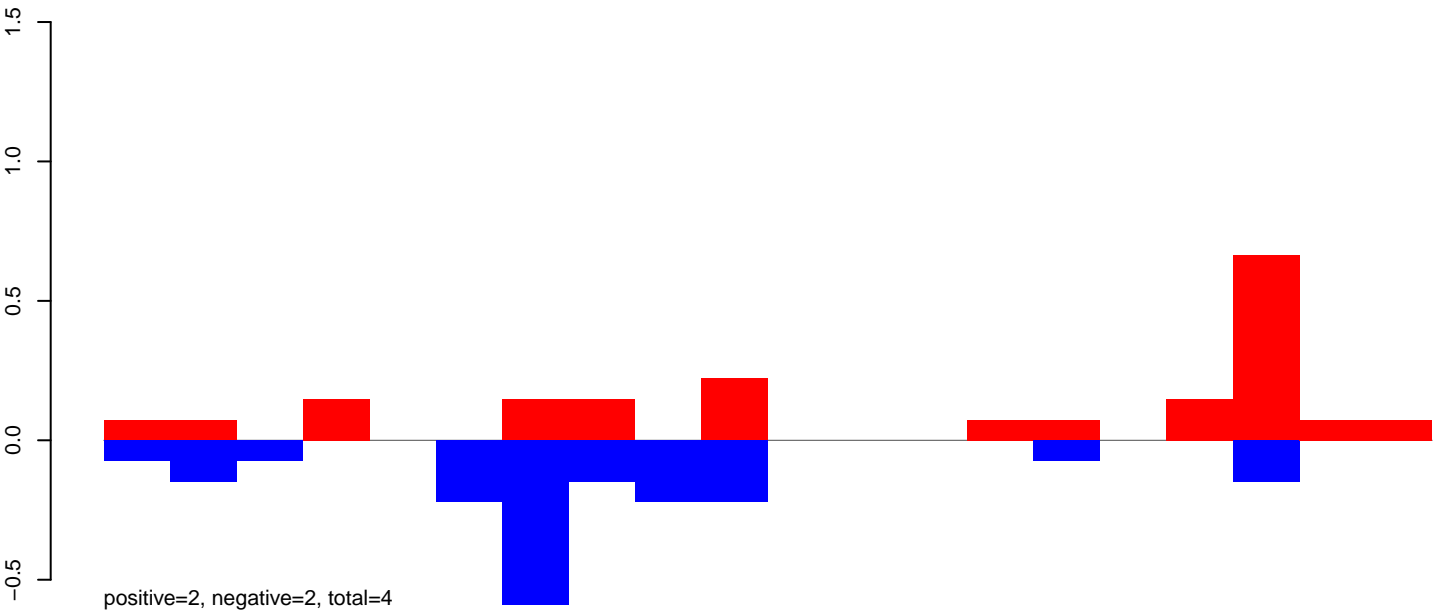
CuQuin_Testes_NL.24_35.rep



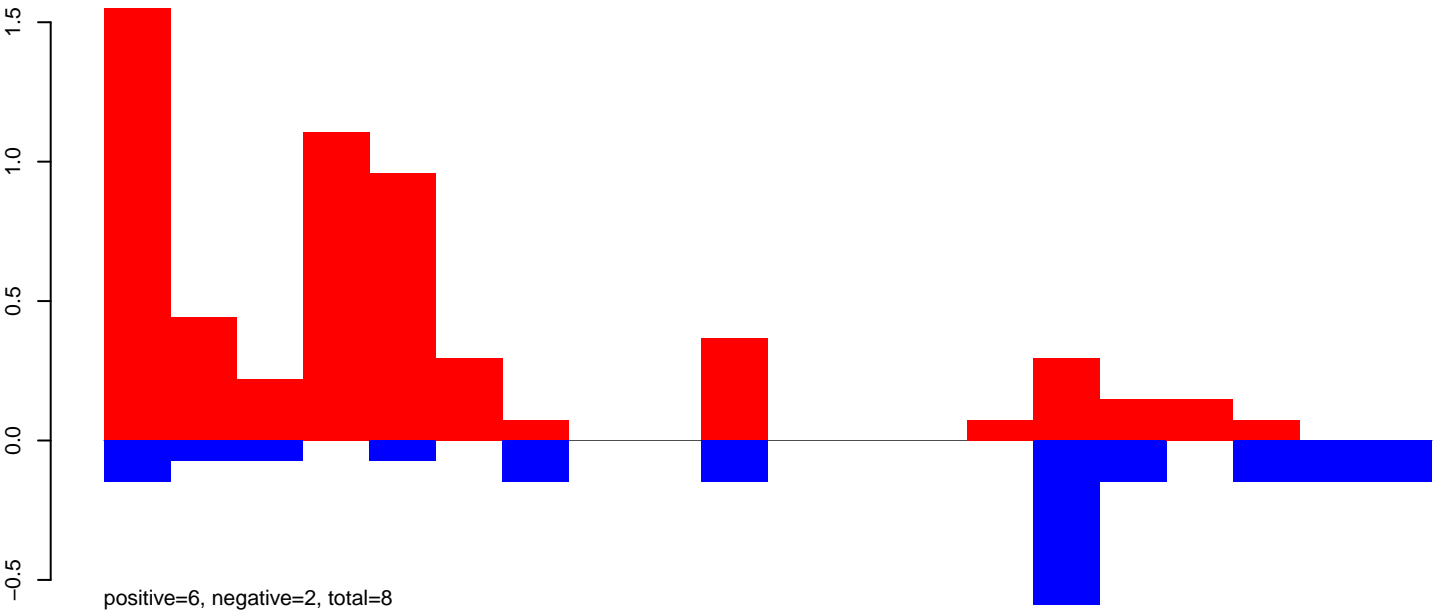
CuQuin_Testes_NL.rep



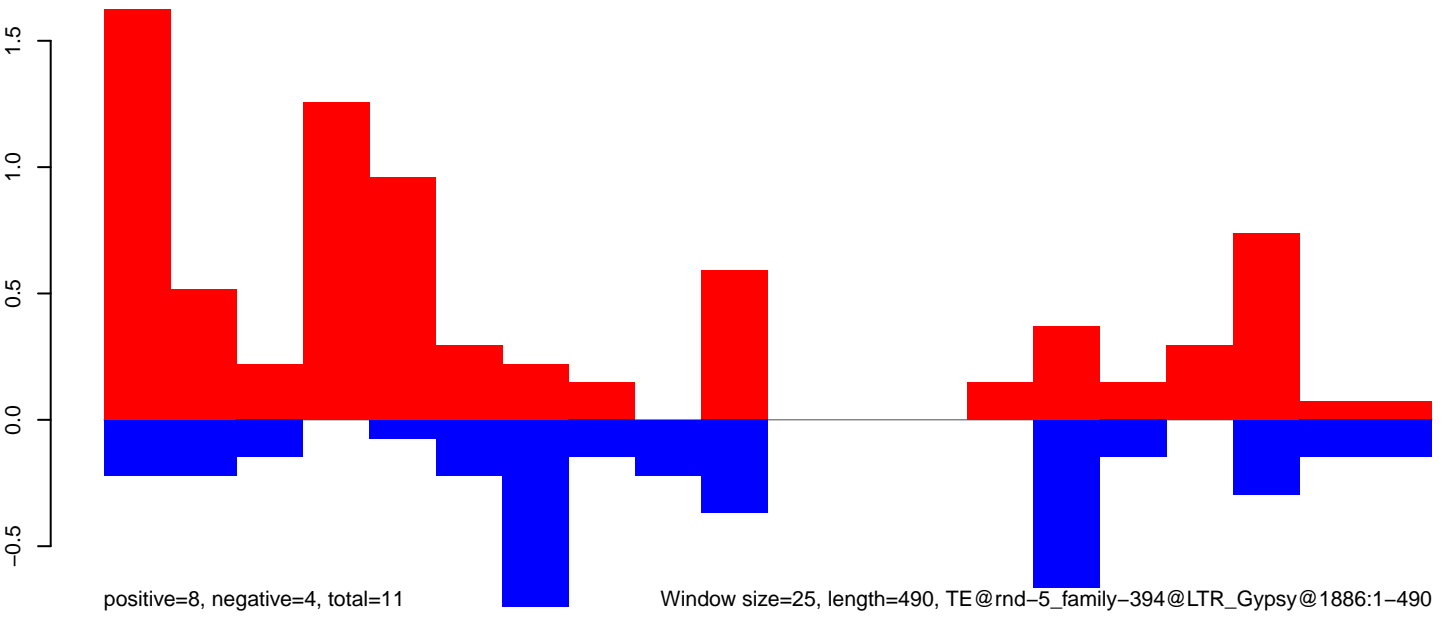
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

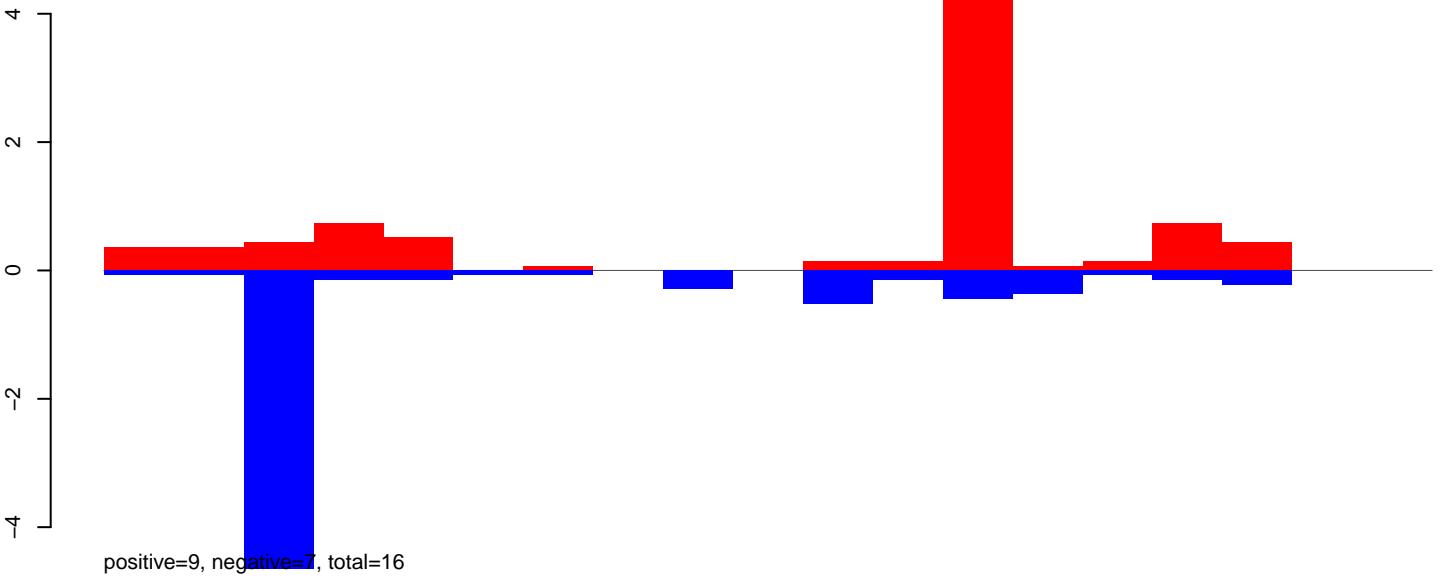


CuQuin_Testes_NL.rep

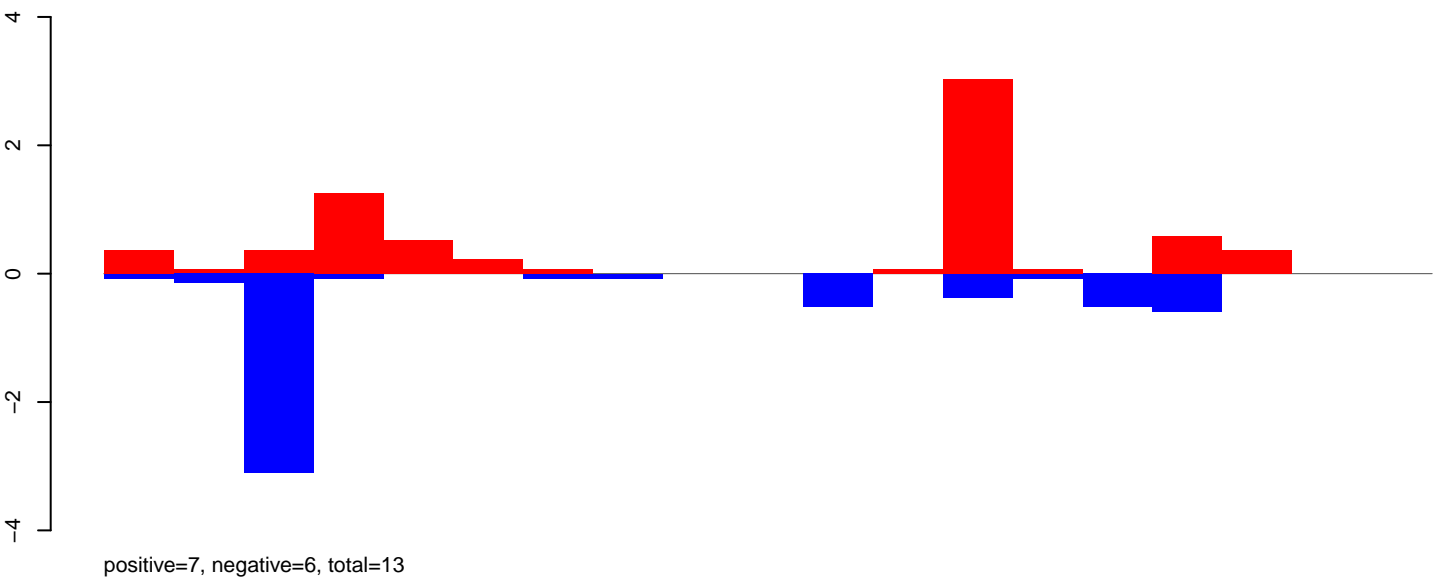


100 200 300 400 500

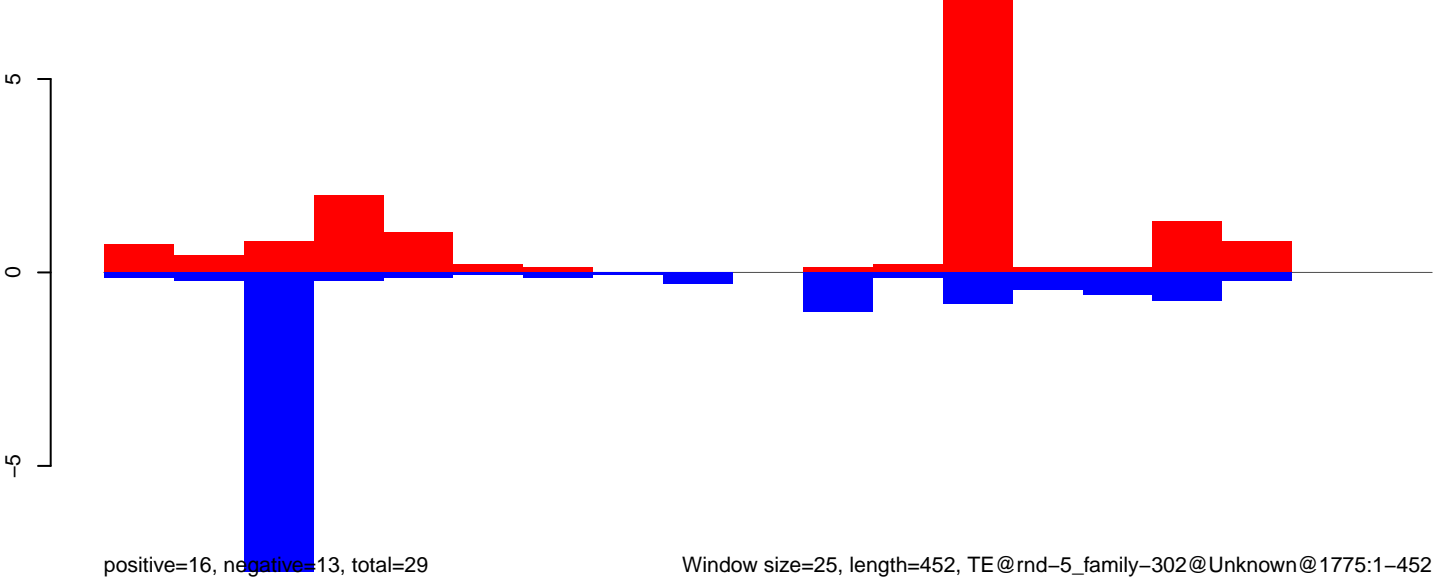
CuQuin_Testes_NL.18_23.rep



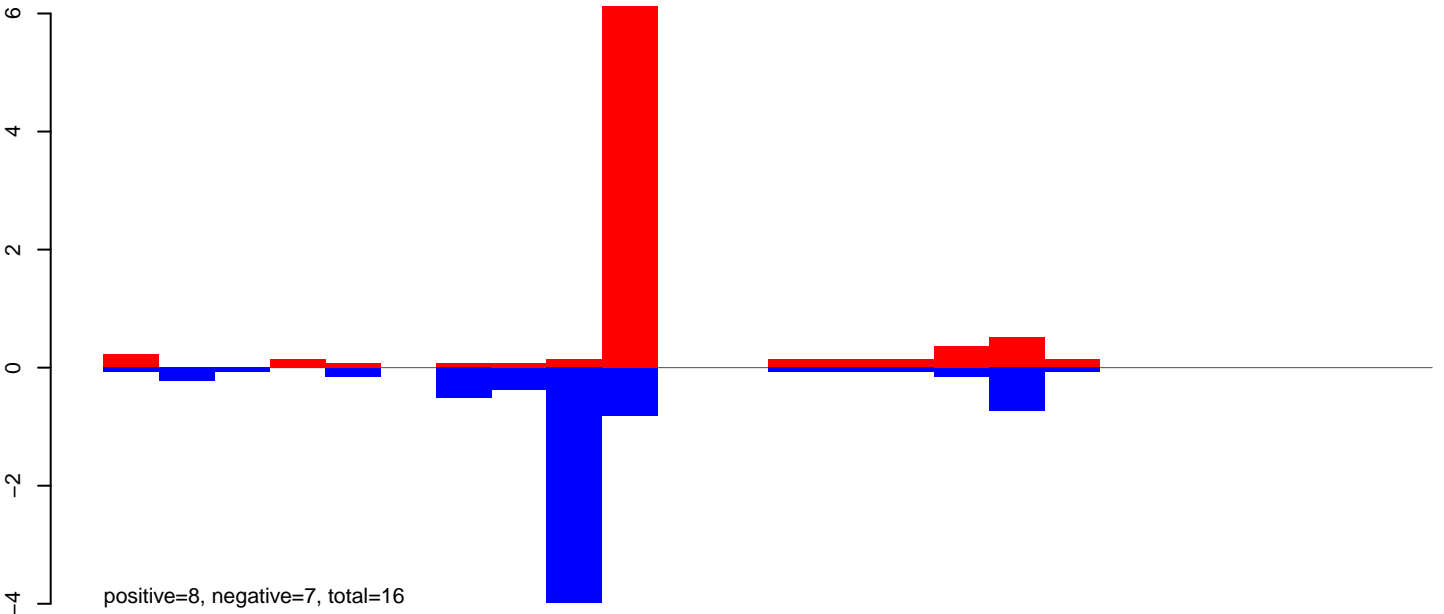
CuQuin_Testes_NL.24_35.rep



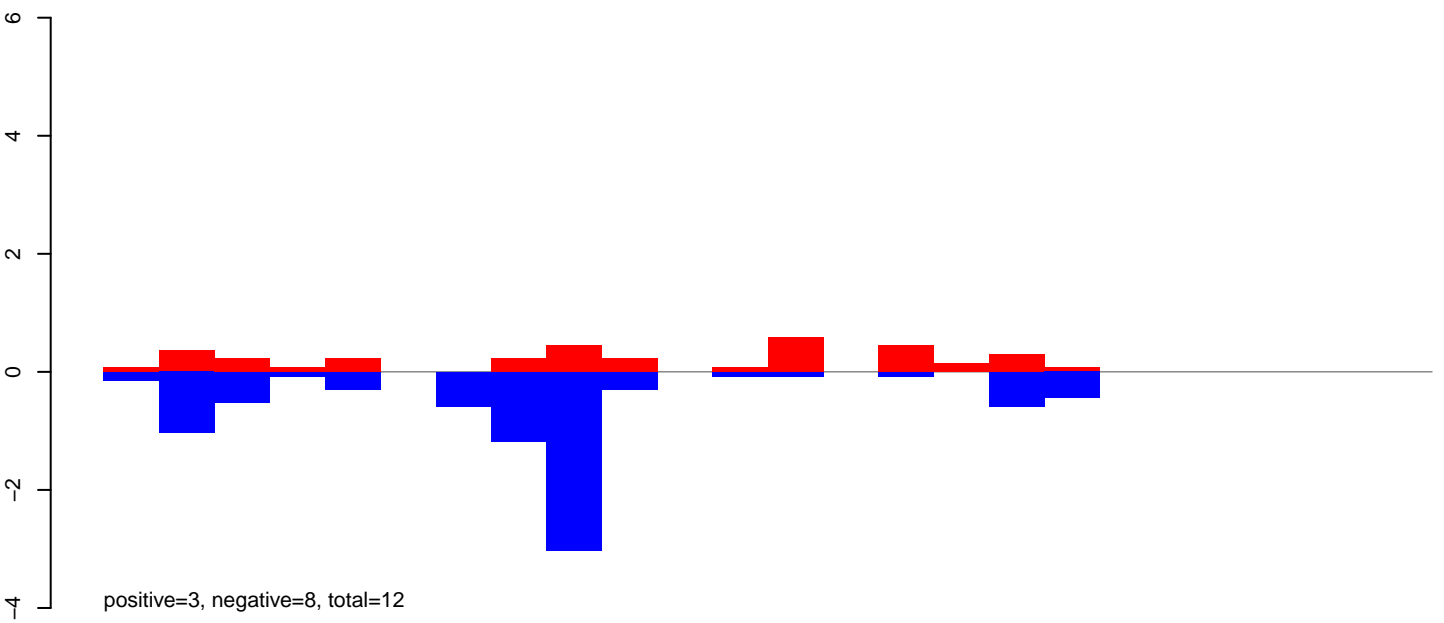
CuQuin_Testes_NL.rep



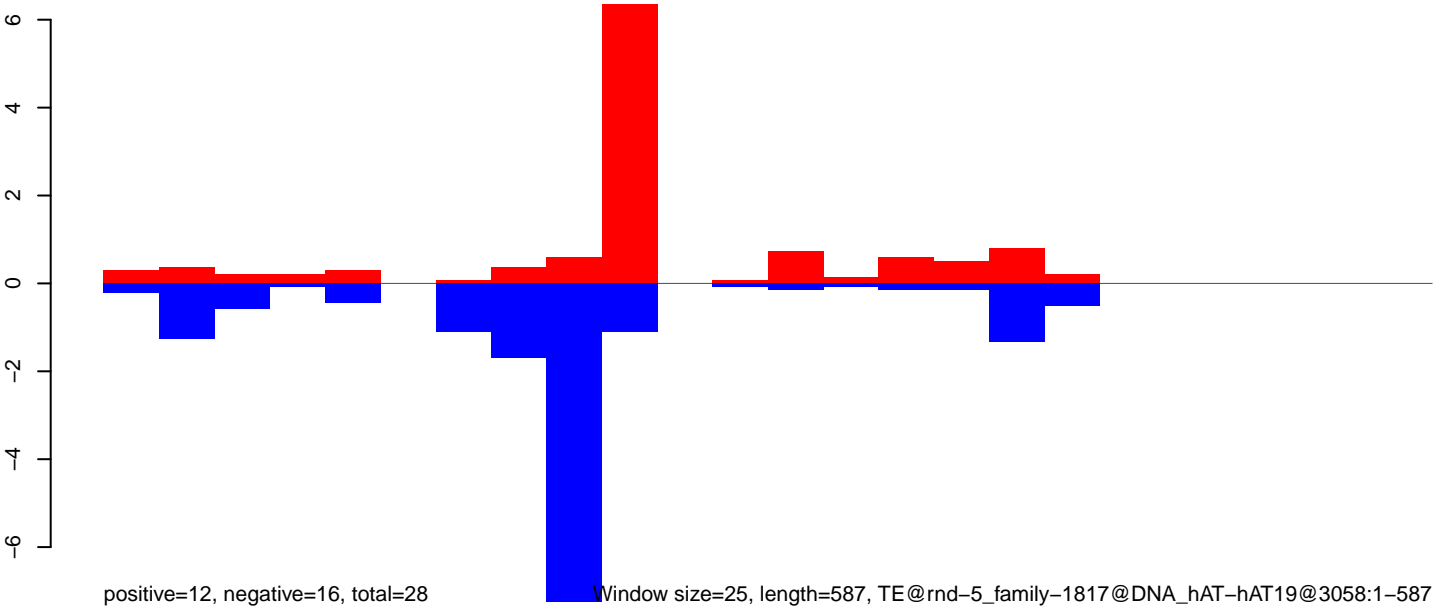
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

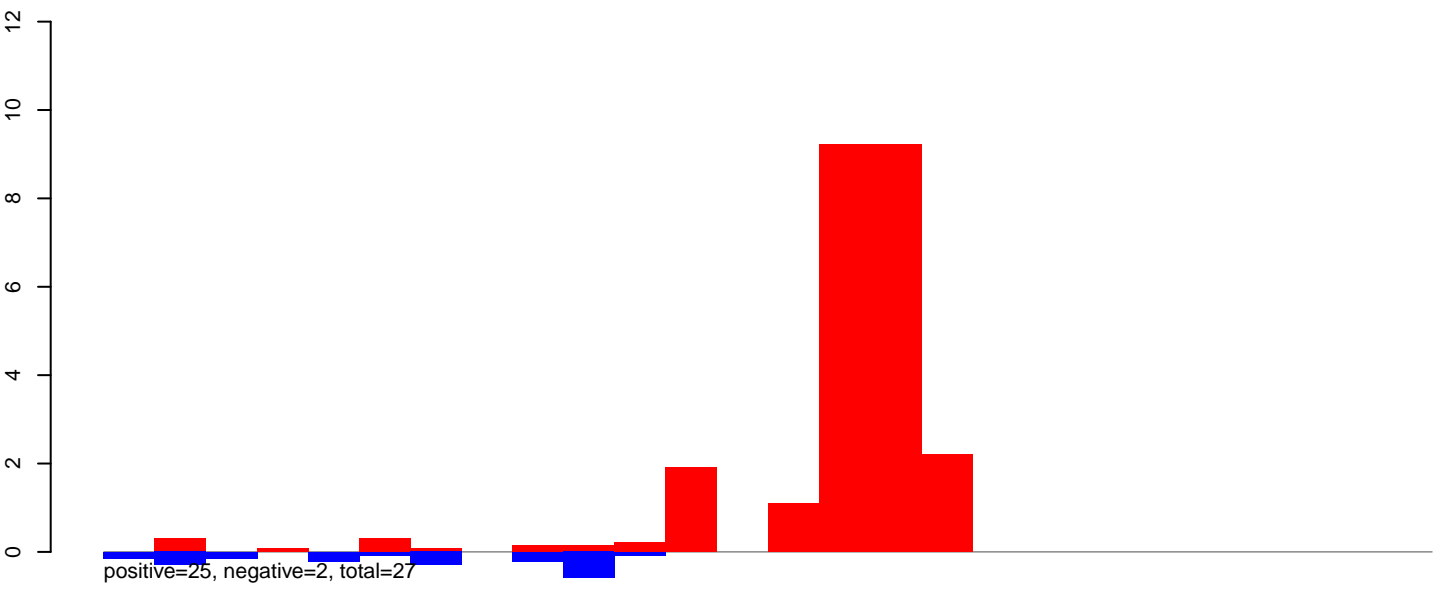


CuQuin_Testes_NL.rep

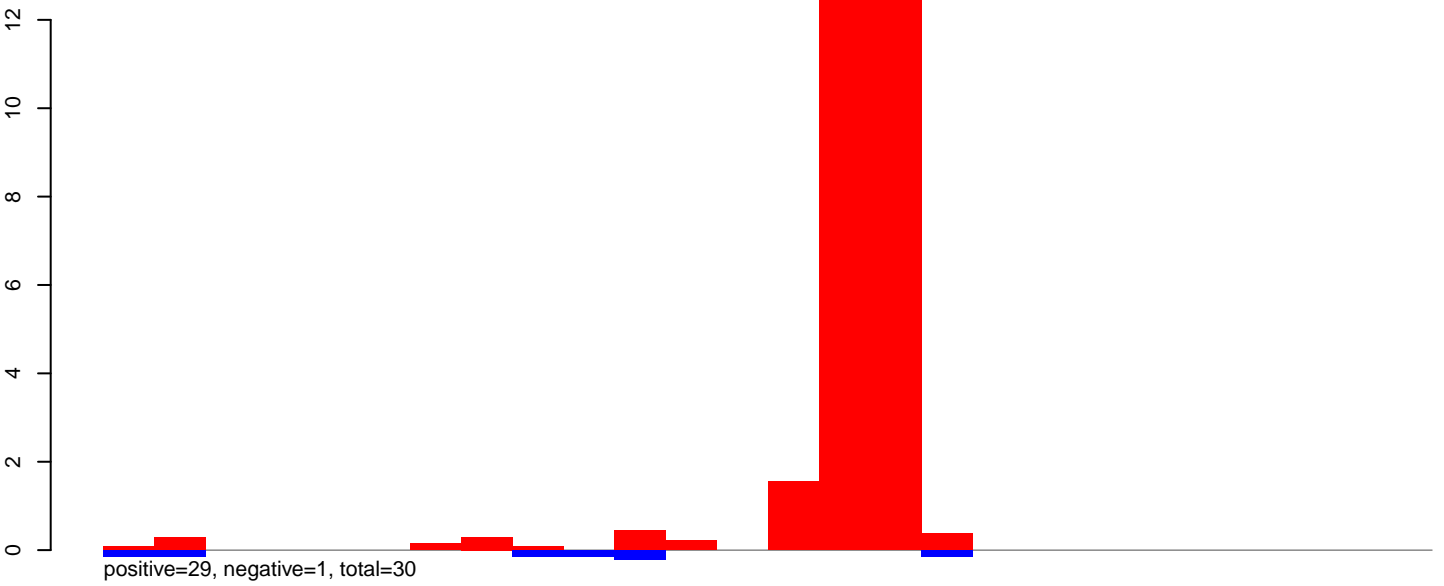


Window size=25, length=587, TE@rnd-5_family-1817@DNA_hAT-hAT19@3058:1-587

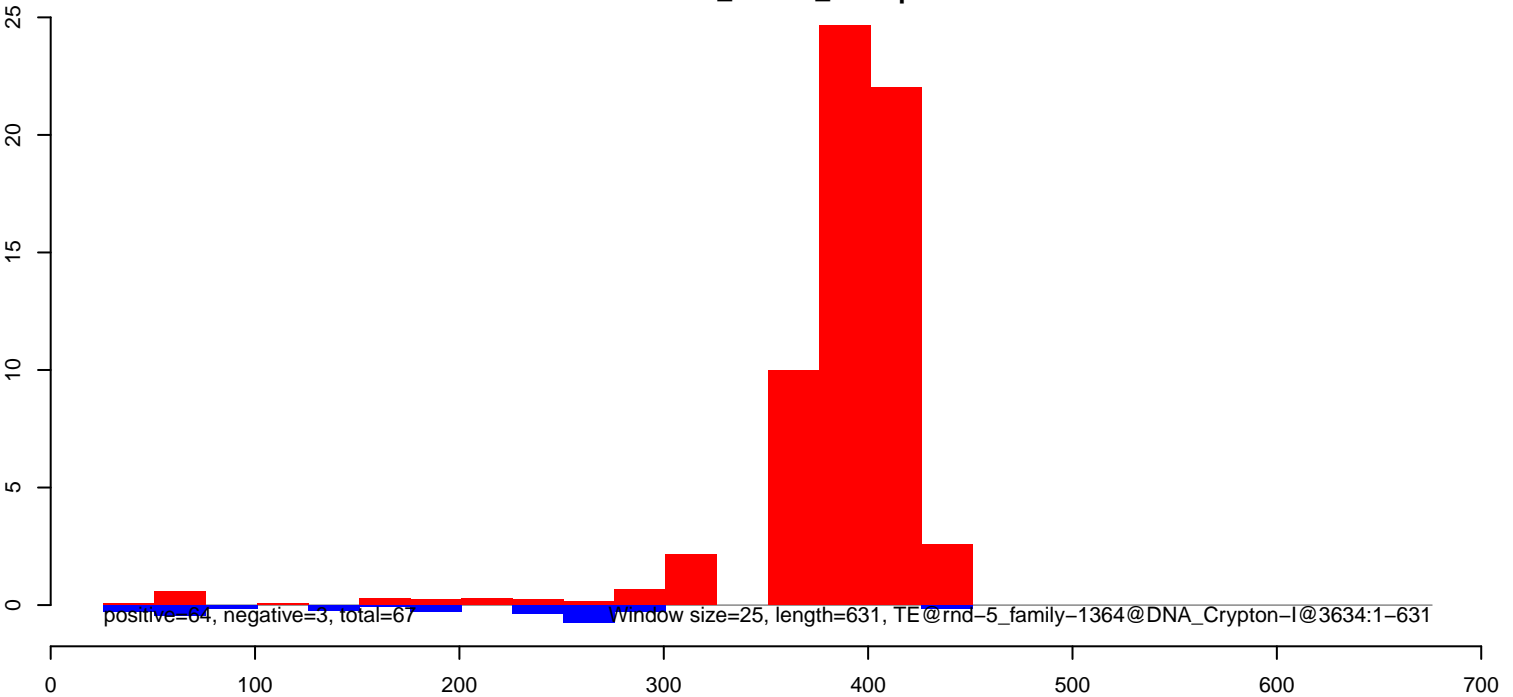
CuQuin_Testes_NL.18_23.rep



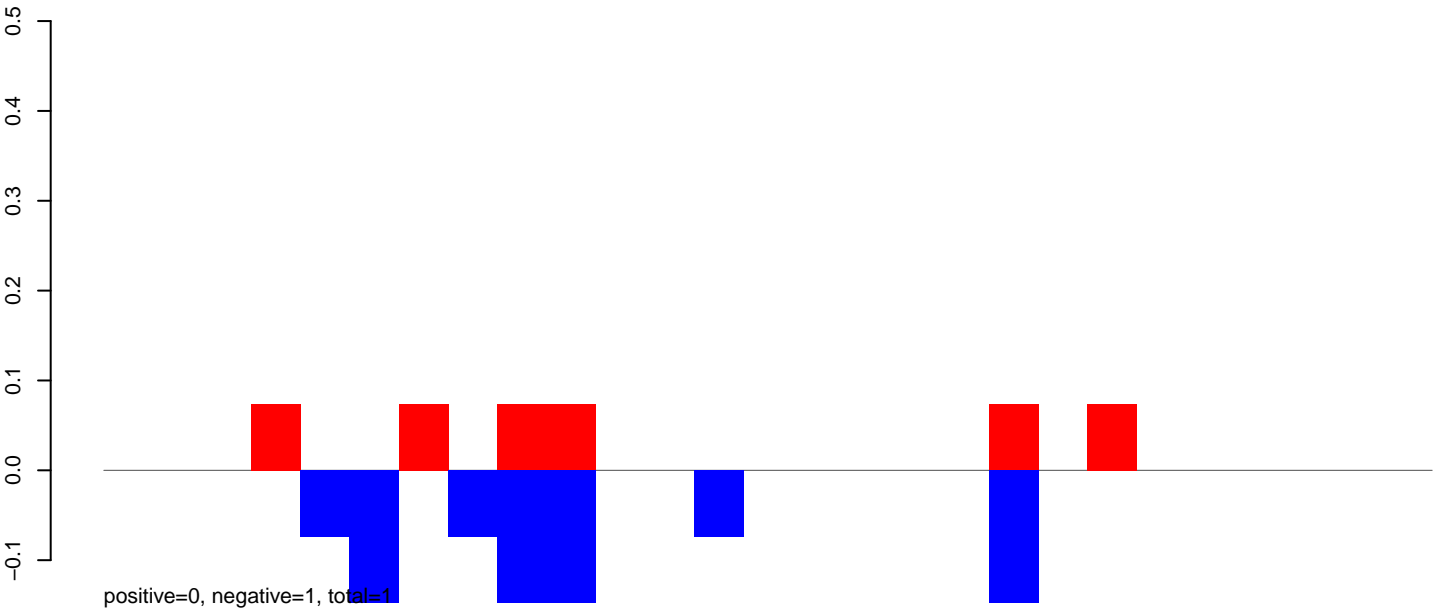
CuQuin_Testes_NL.24_35.rep



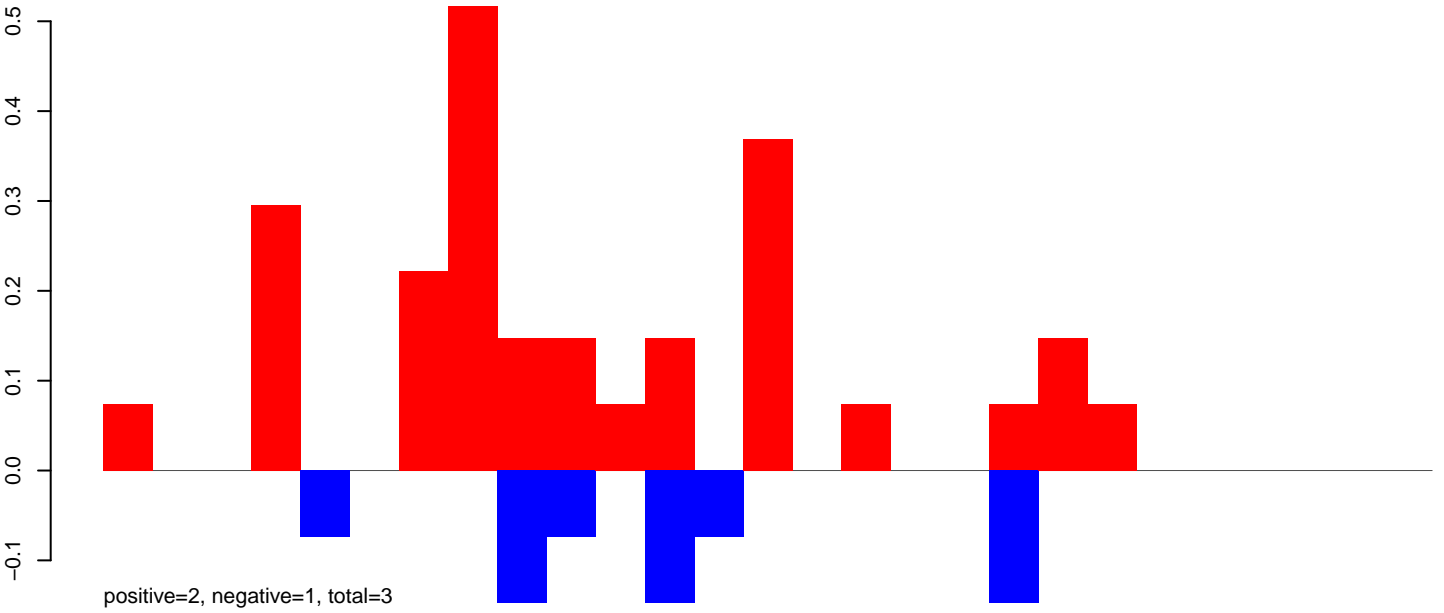
CuQuin_Testes_NL.rep



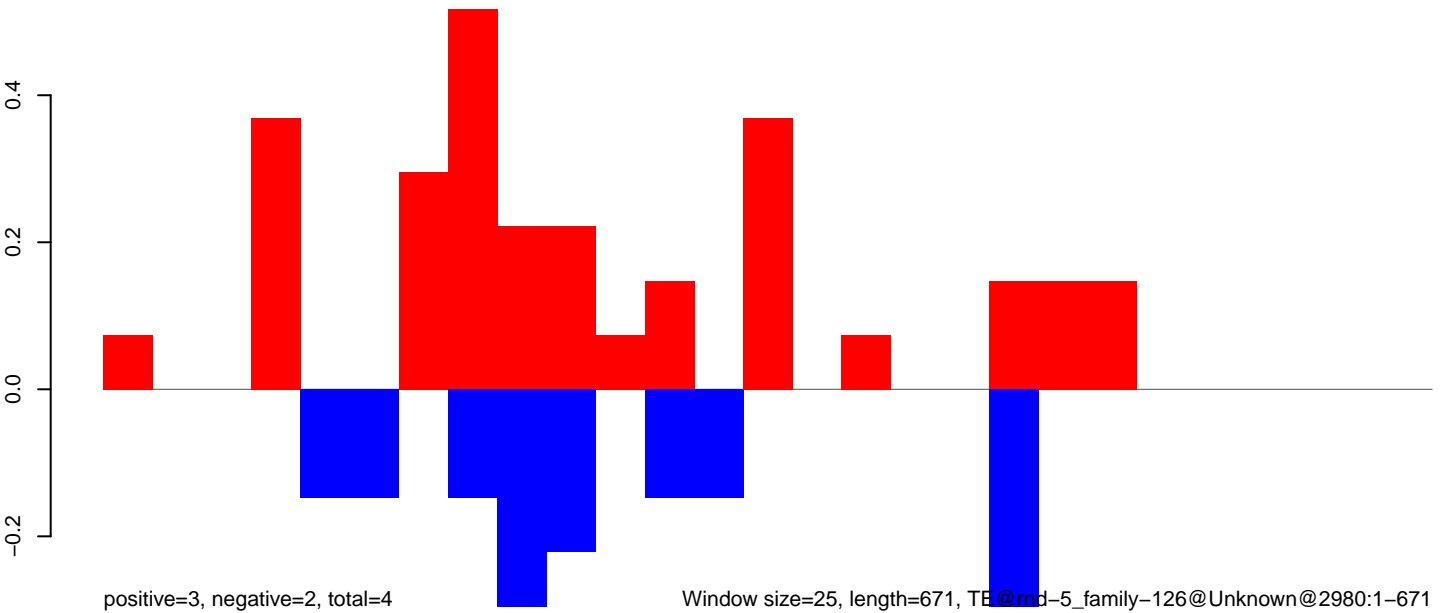
CuQuin_Testes_NL.18_23.rep



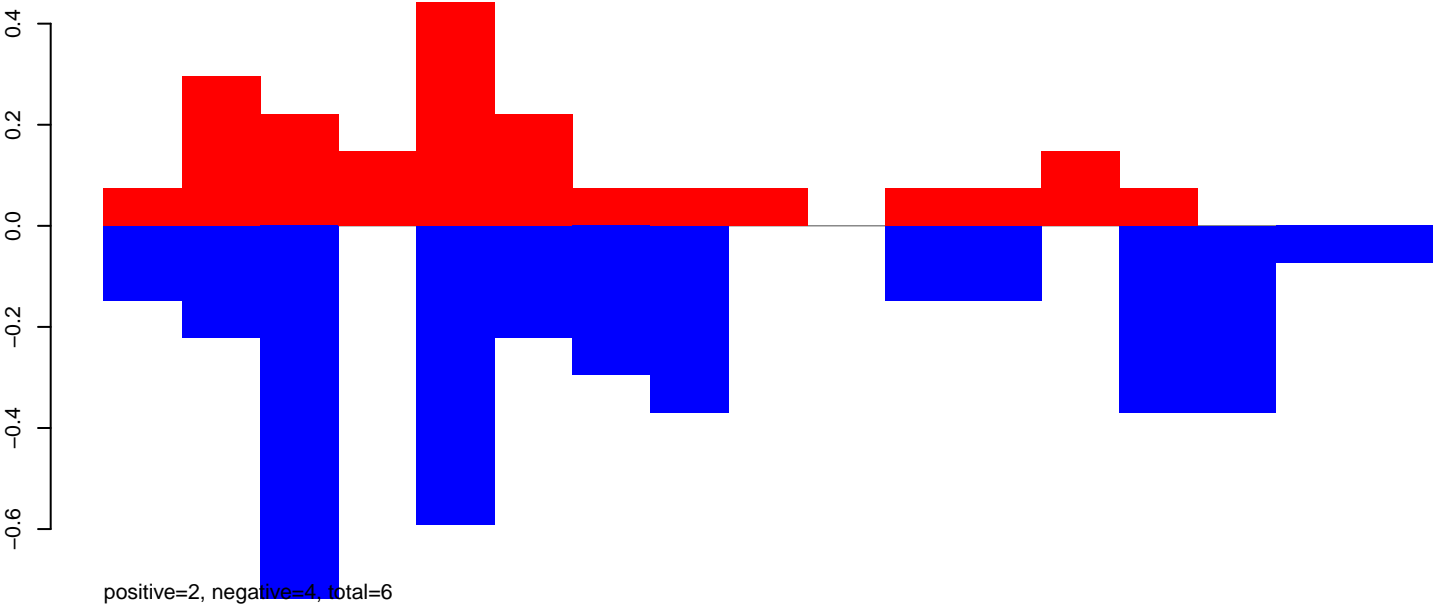
CuQuin_Testes_NL.24_35.rep



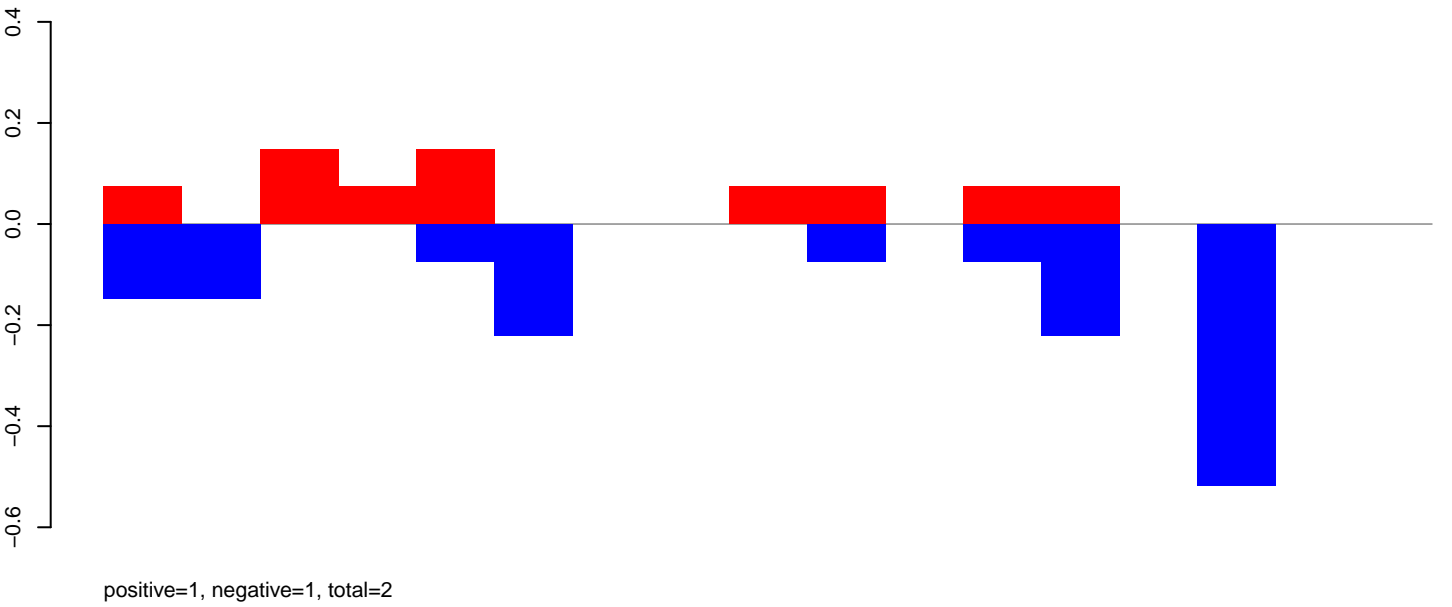
CuQuin_Testes_NL.rep



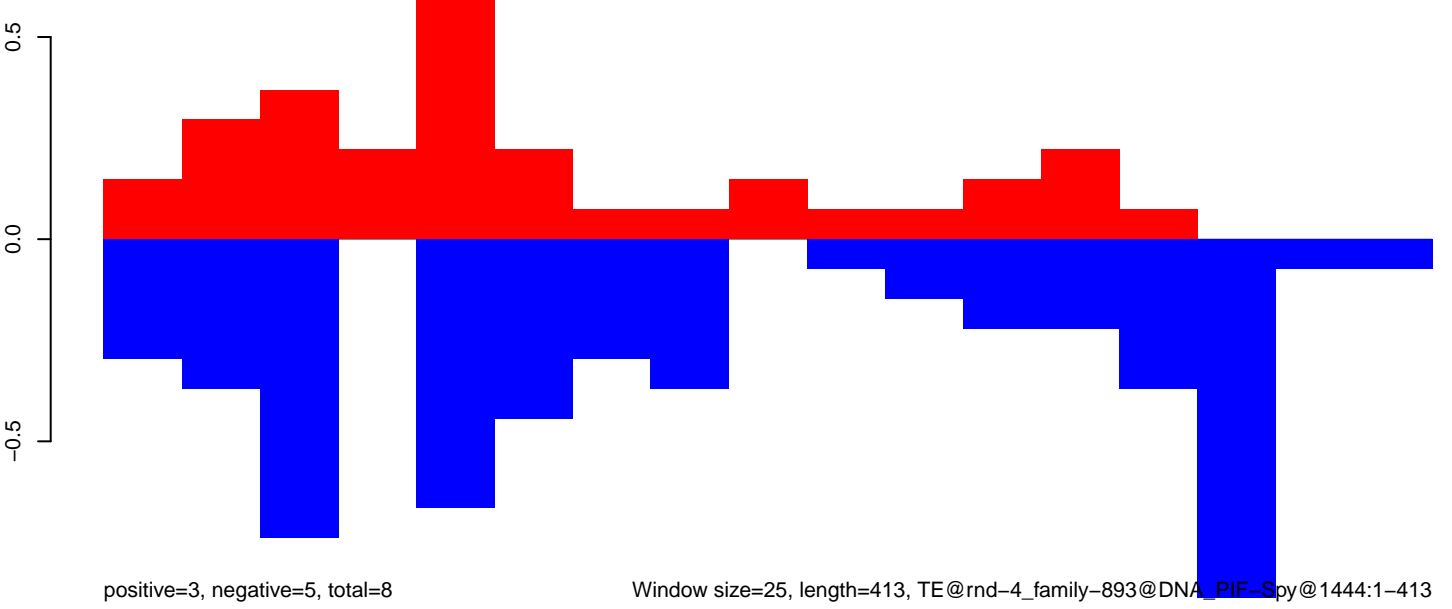
CuQuin_Testes_NL.18_23.rep



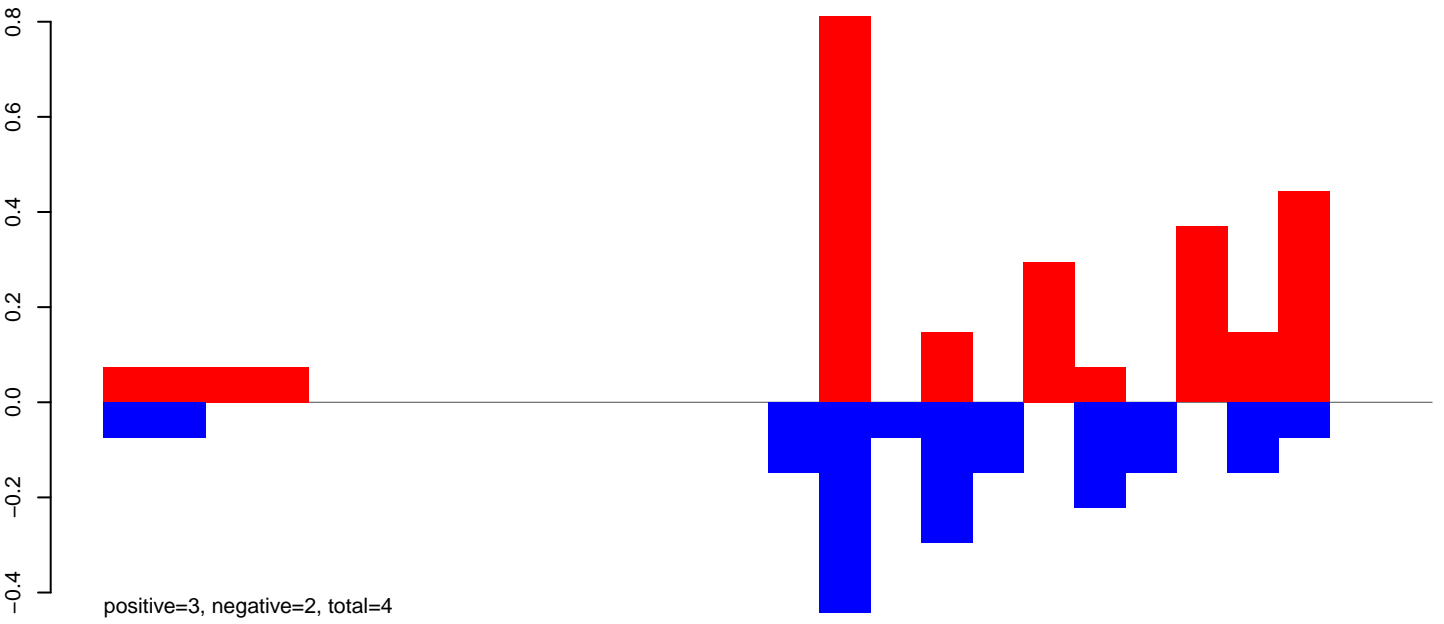
CuQuin_Testes_NL.24_35.rep



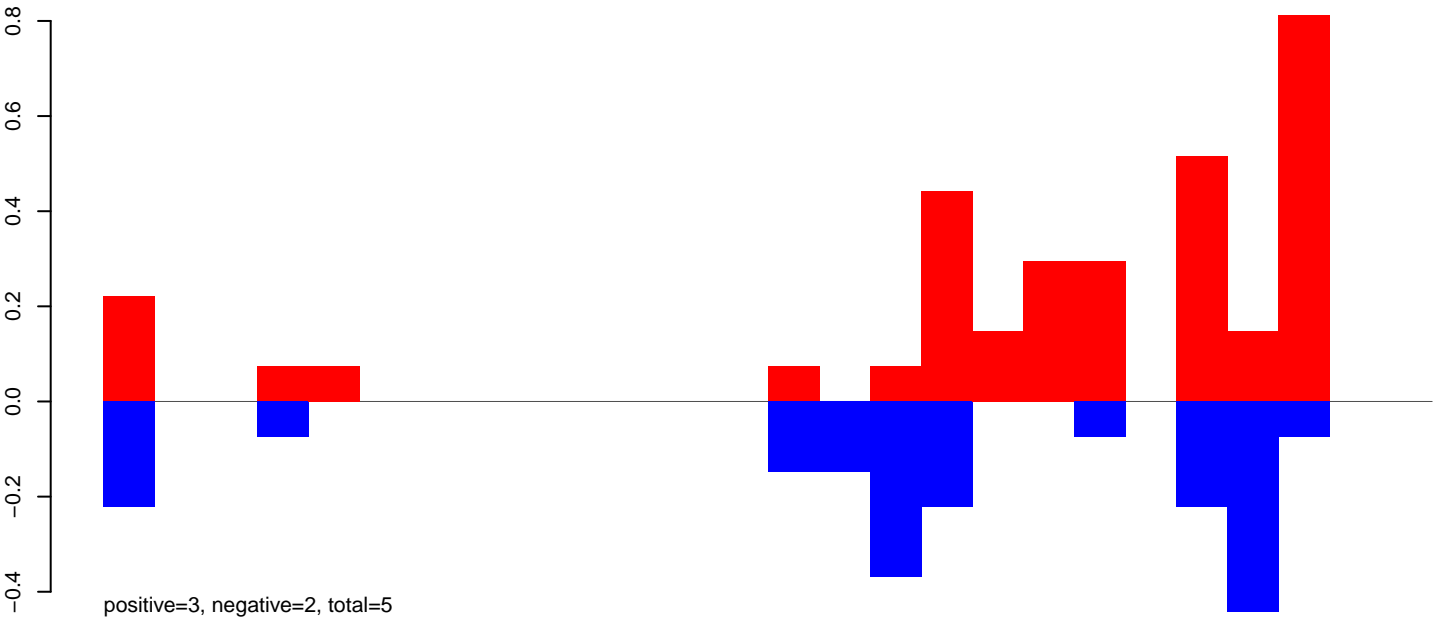
CuQuin_Testes_NL.rep



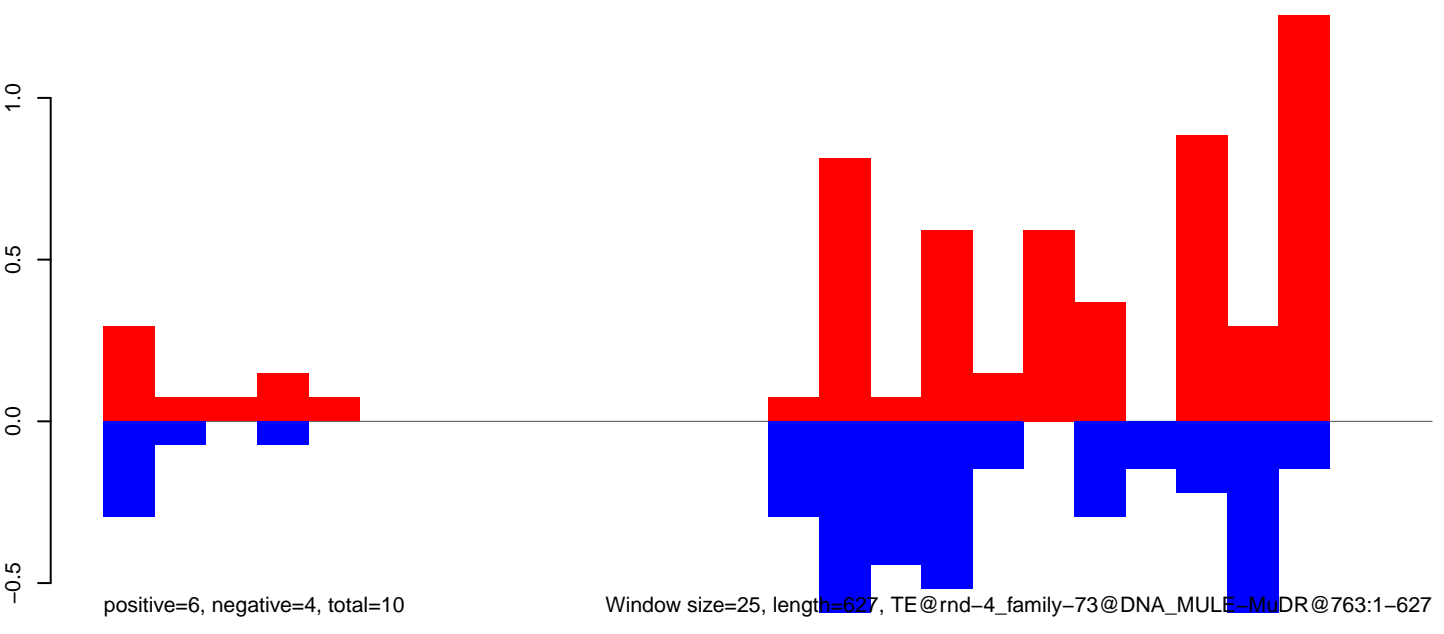
CuQuin_Testes_NL.18_23.rep



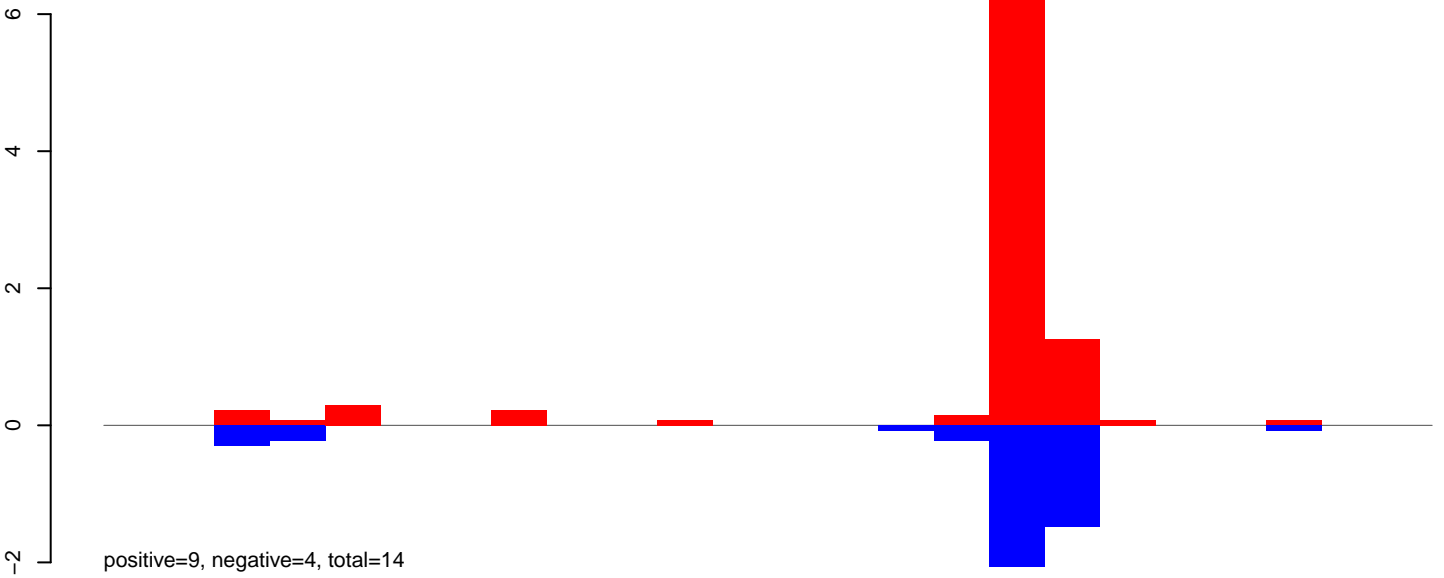
CuQuin_Testes_NL.24_35.rep



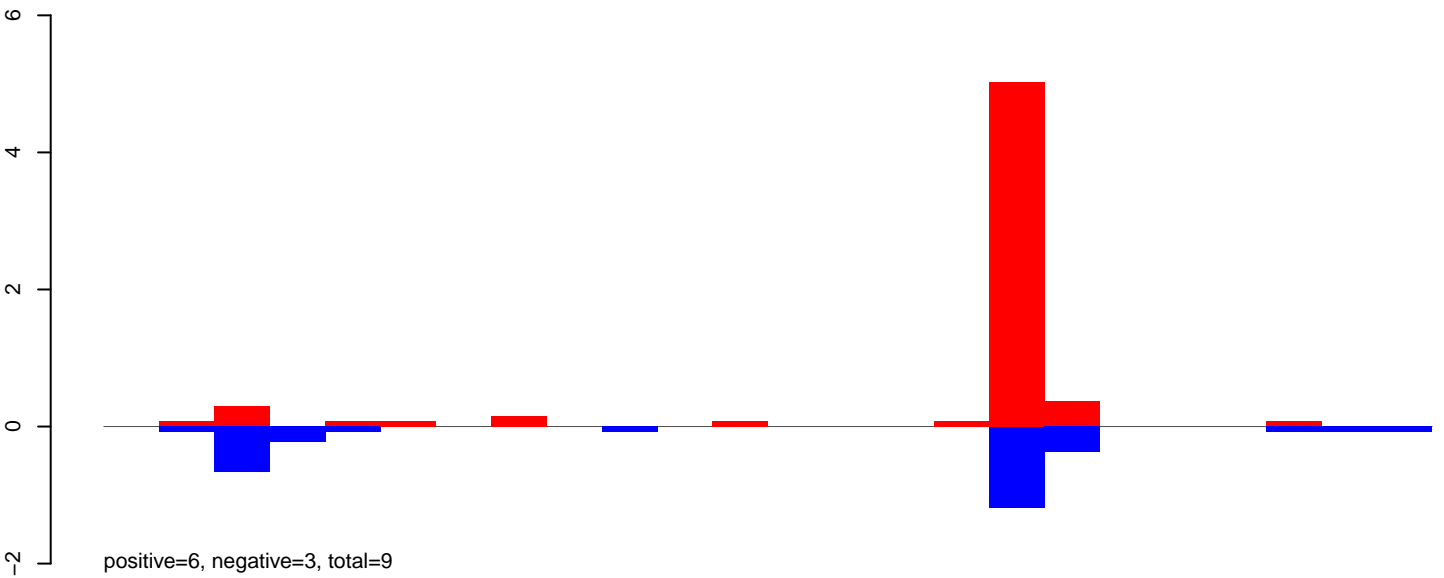
CuQuin_Testes_NL.rep



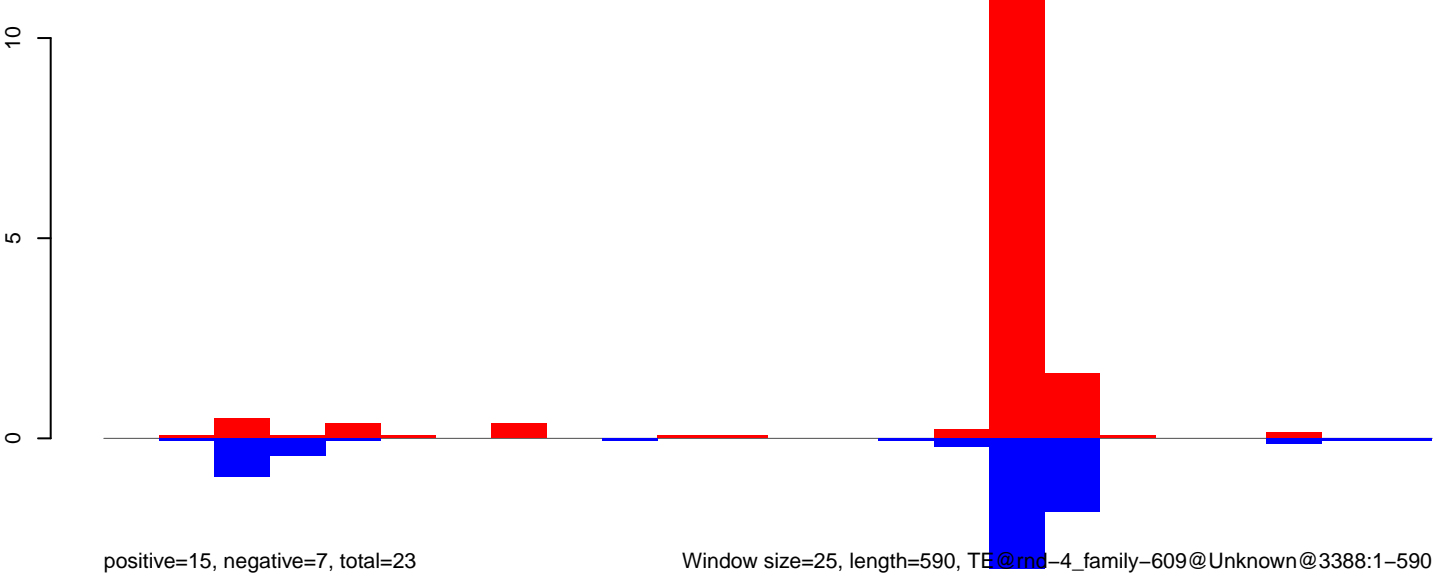
CuQuin_Testes_NL.18_23.rep



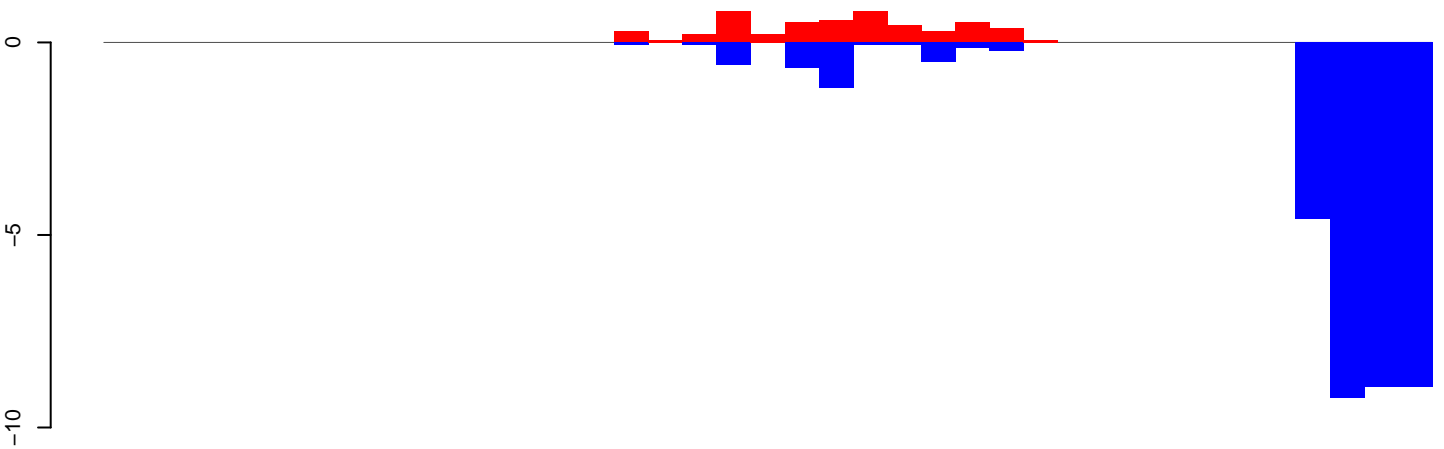
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

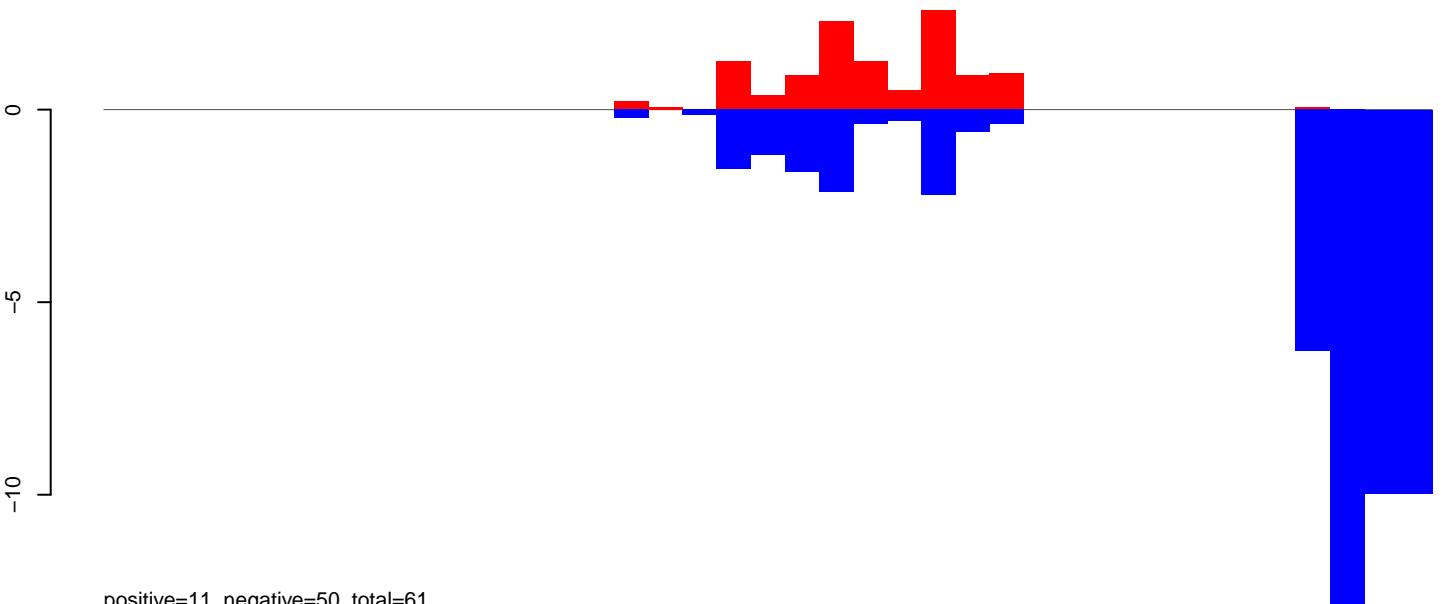


CuQuin_Testes_NL.18_23.rep



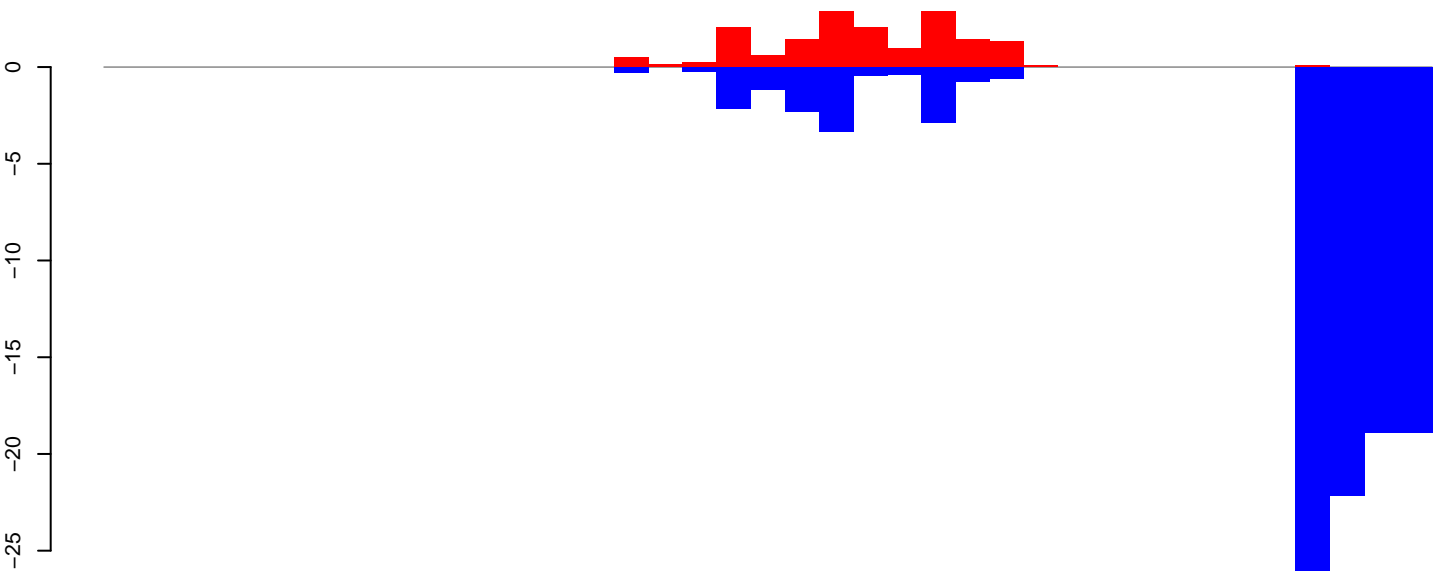
positive=5, negative=35, total=41

CuQuin_Testes_NL.24_35.rep



positive=11, negative=50, total=61

CuQuin_Testes_NL.rep

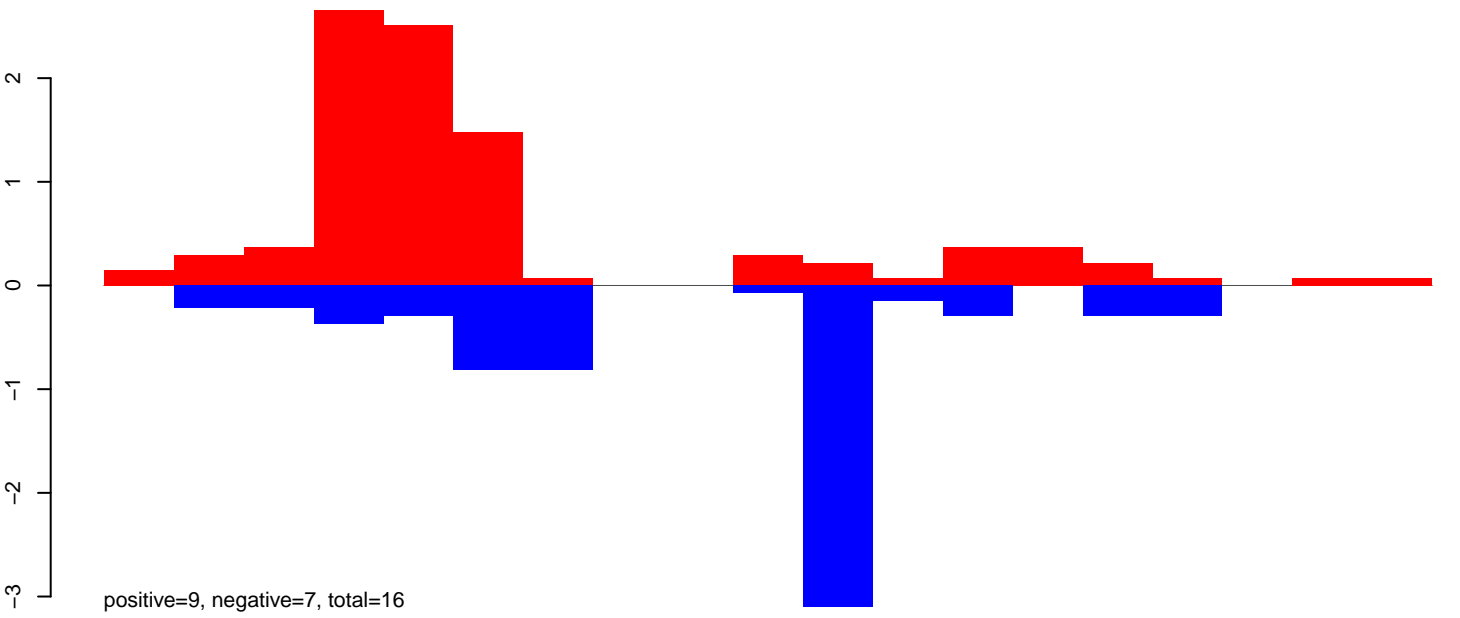


positive=17, negative=102, total=119

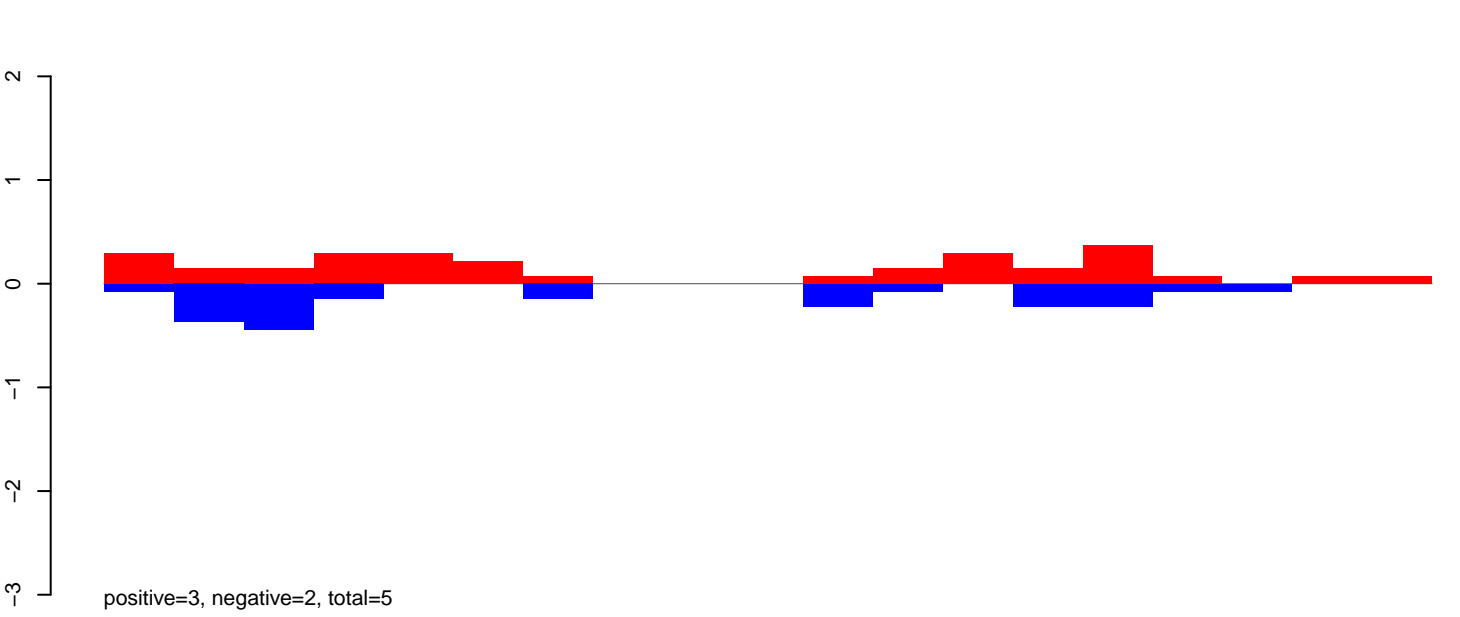
Window size=25, length=971, TE@rnd-4_family-393@SINE_tRNA-Deu@370:1-971

0 200 400 600 800 1000

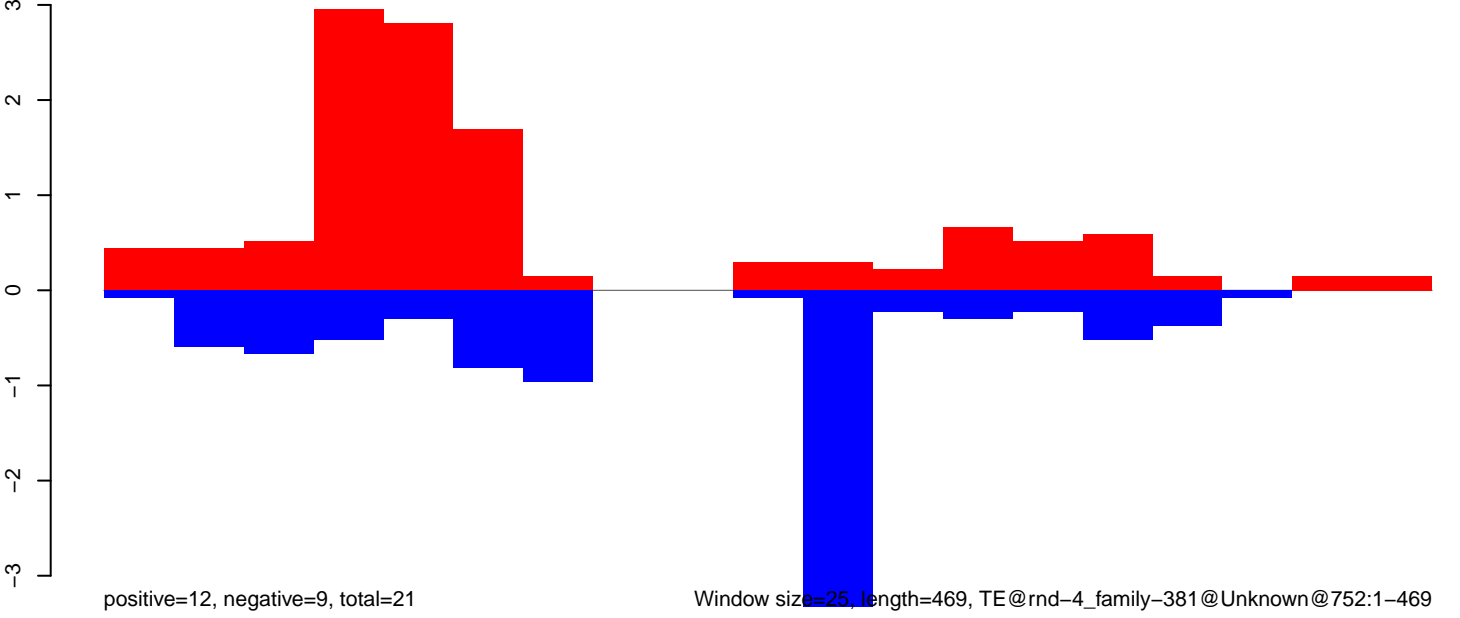
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



Window size=25, length=469, TE@rnd-4_family-381@Unknown@752:1-469

CuQuin_Testes_NL.18_23.rep

0.5
0.0
-0.5
-1.0
-1.5

positive=2, negative=2, total=3

CuQuin_Testes_NL.24_35.rep

0.5
0.0
-0.5
-1.0
-1.5

positive=1, negative=6, total=7

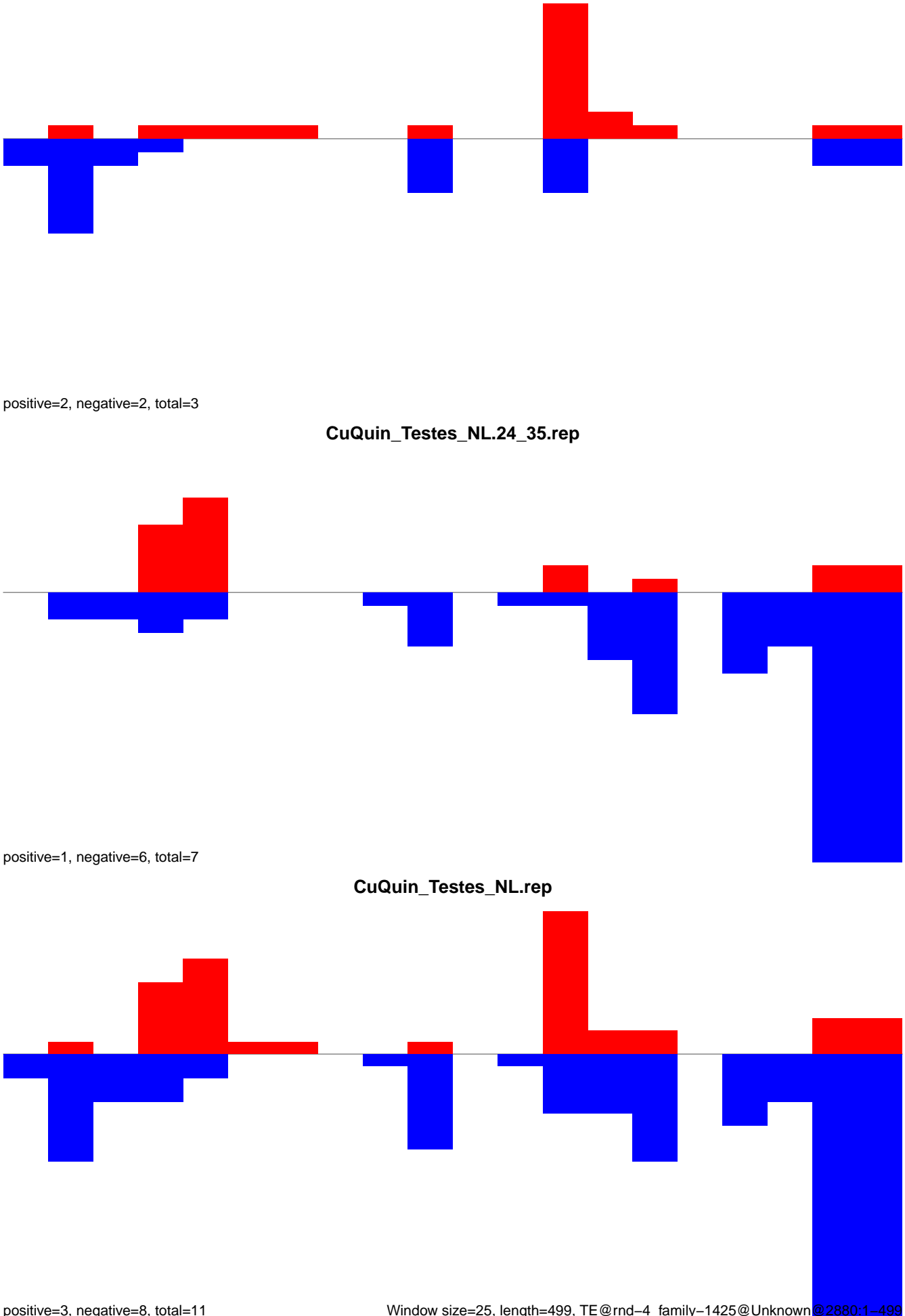
CuQuin_Testes_NL.rep

0.5
0.0
-0.5
-1.0
-1.5

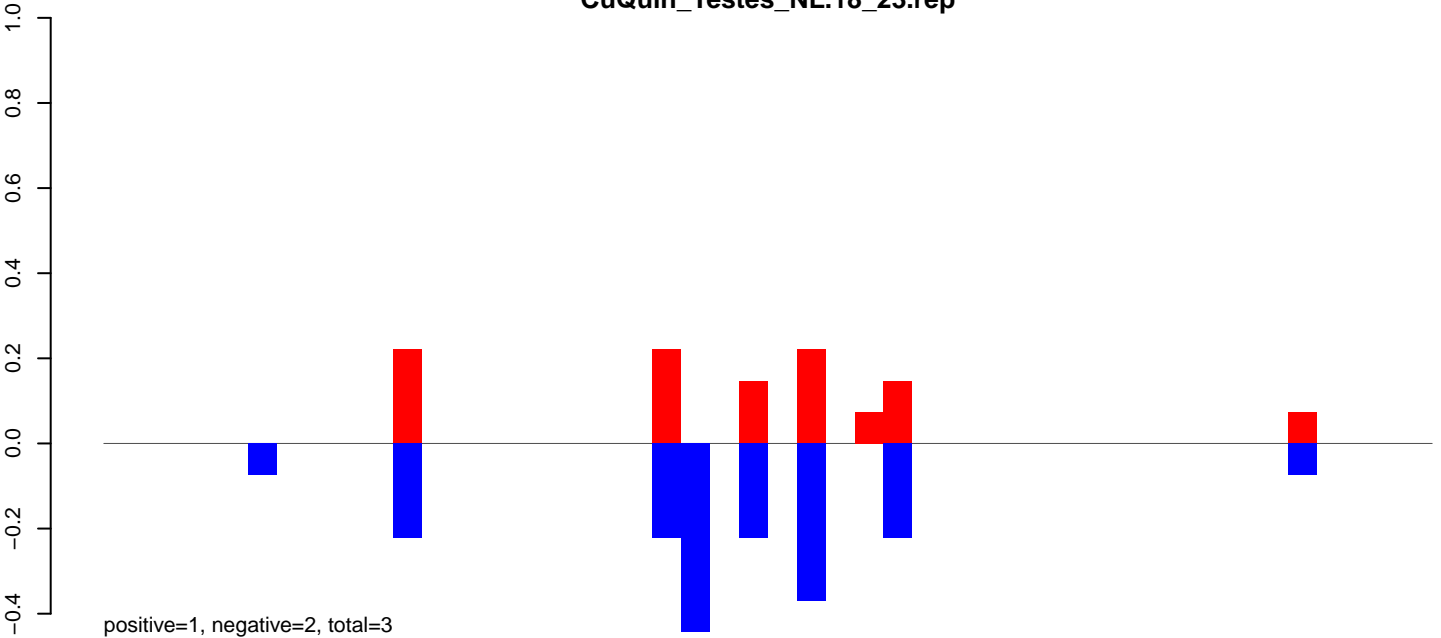
positive=3, negative=8, total=11

Window size=25, length=499, TE@rnd-4_family-1425@Unknown@2880:1-499

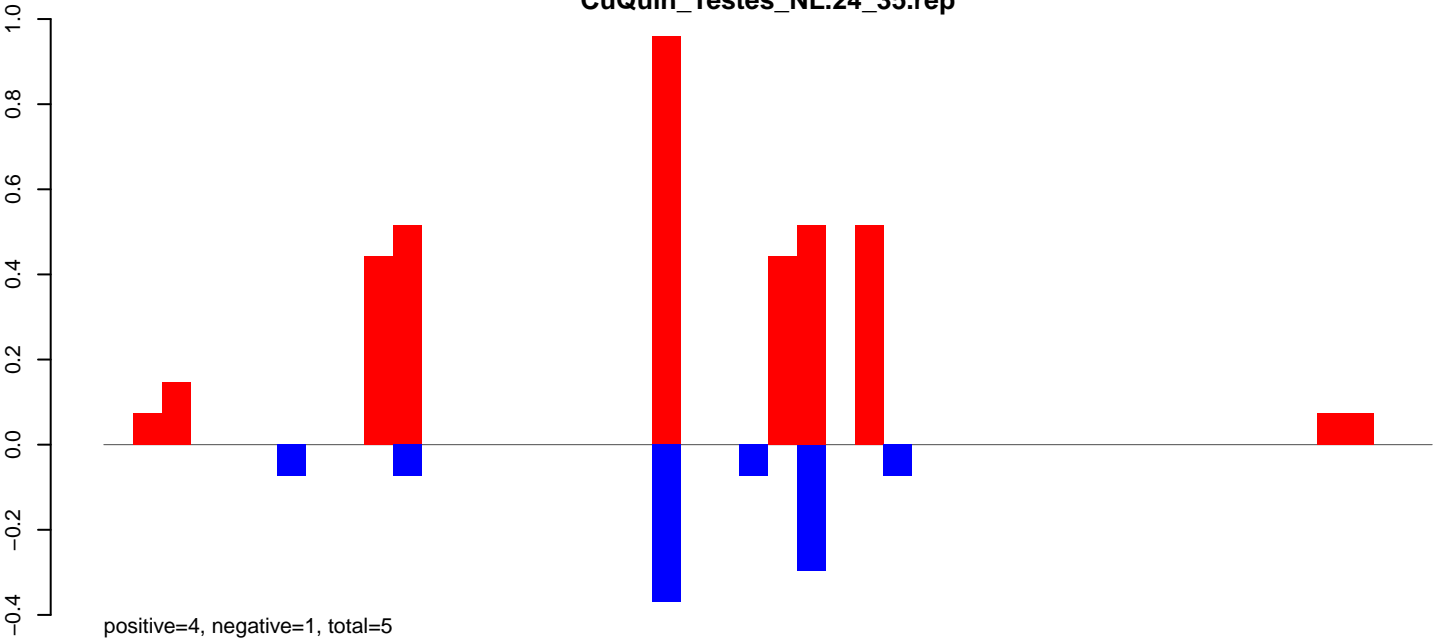
100 200 300 400 500



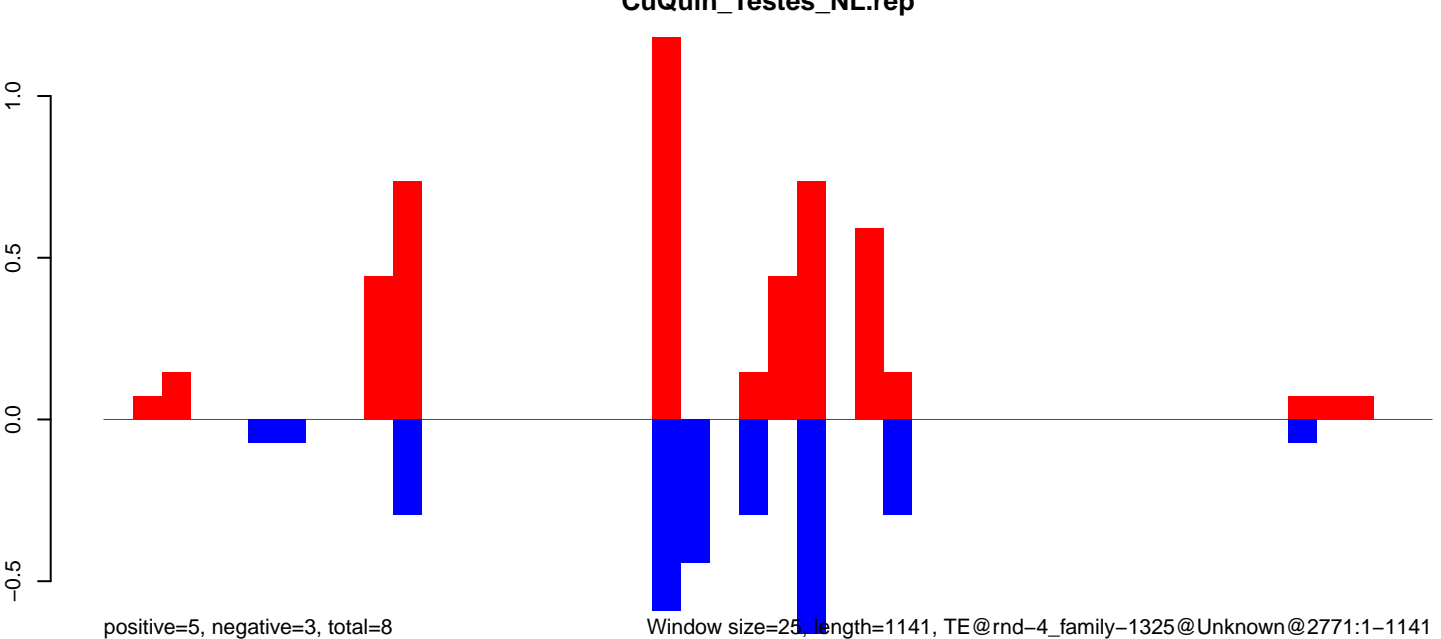
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



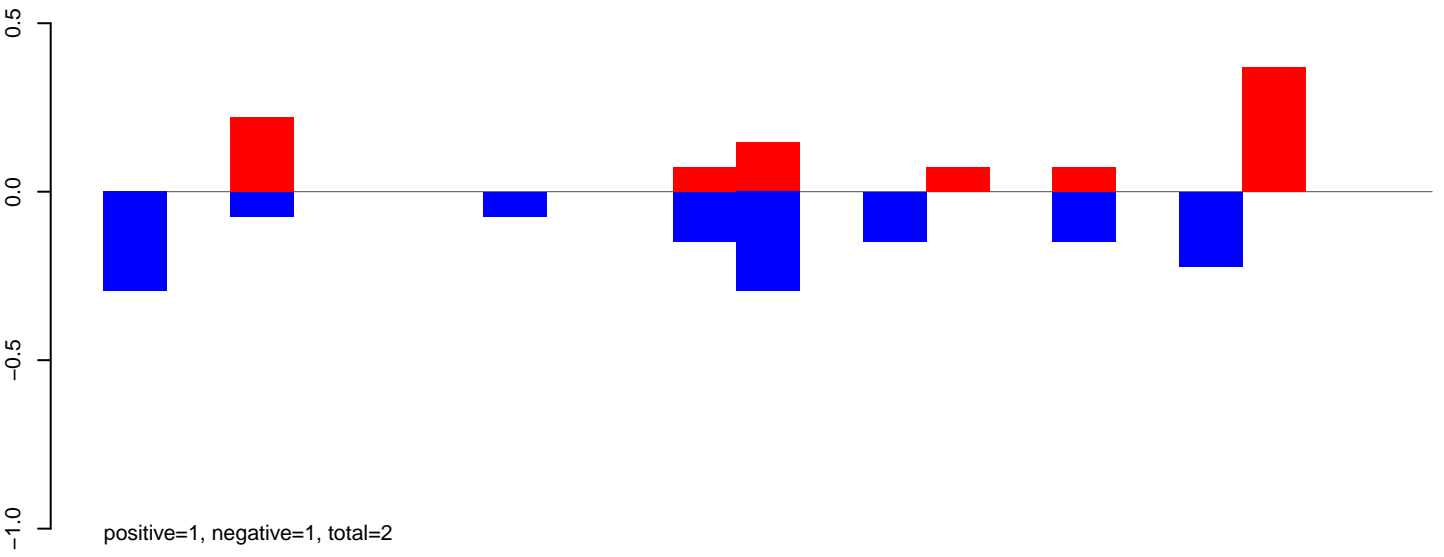
CuQuin_Testes_NL.rep



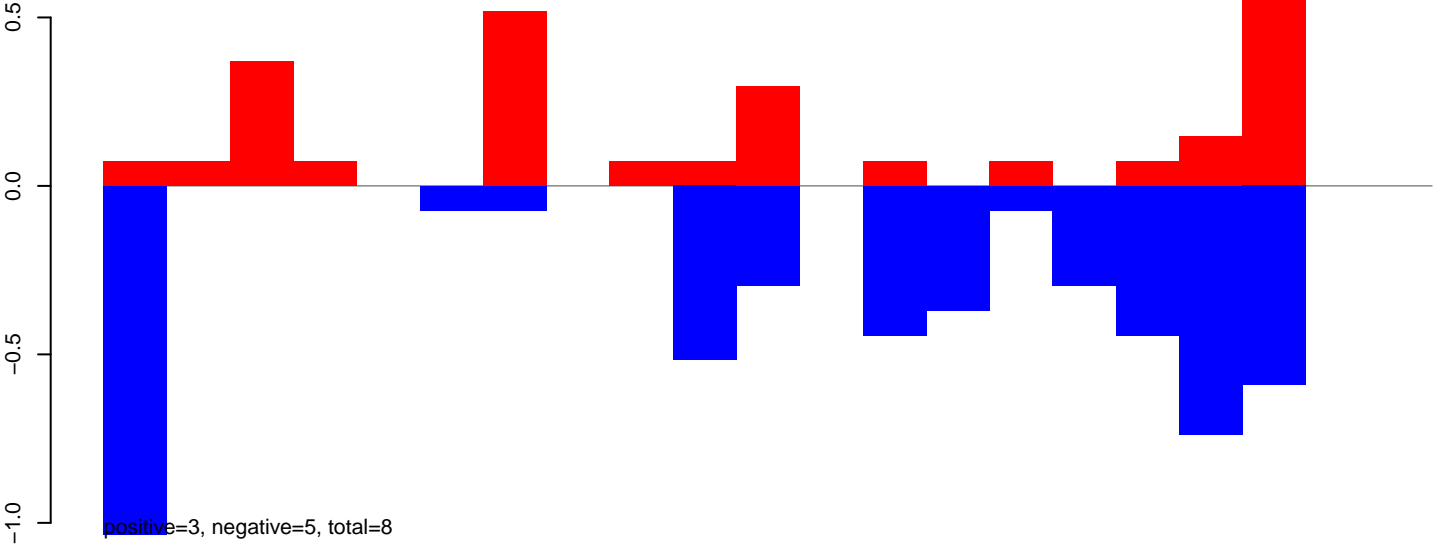
Window size=25, length=1141, TE@rnd-4_family-1325@Unknown@2771:1-1141

0 200 400 600 800 1000 1200

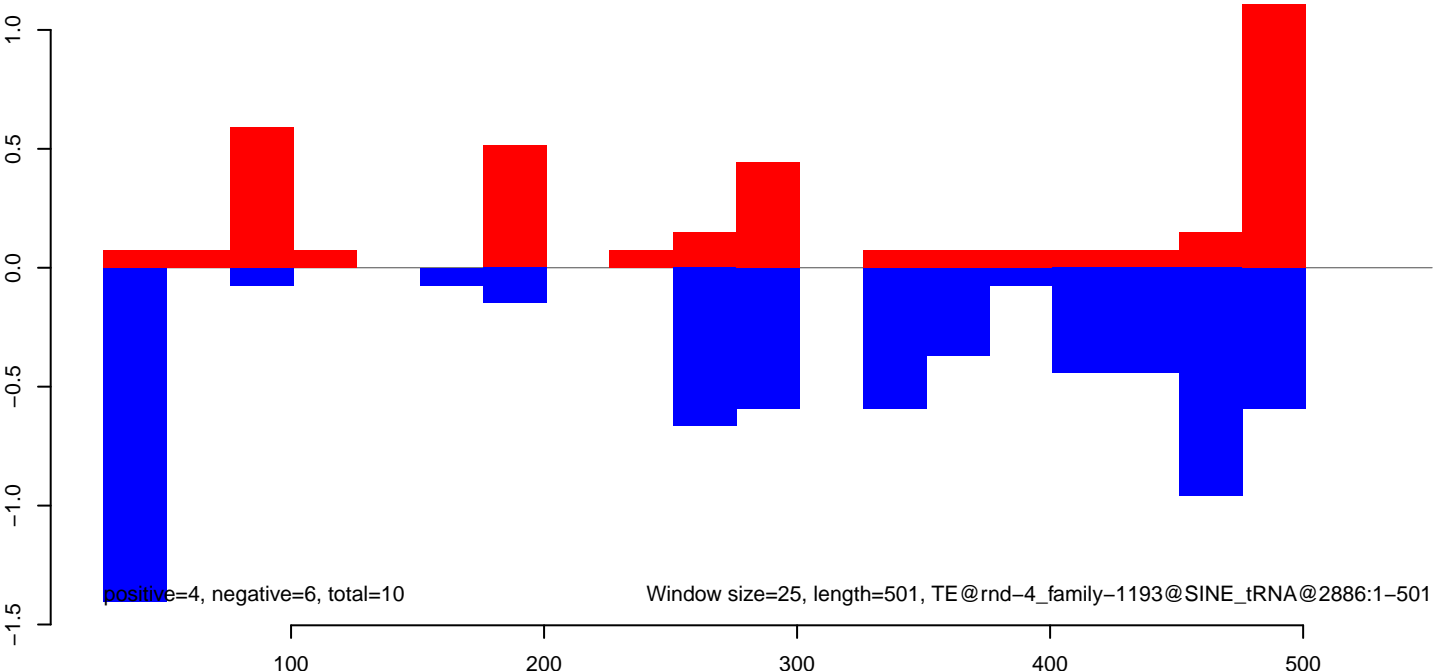
CuQuin_Testes_NL.18_23.rep



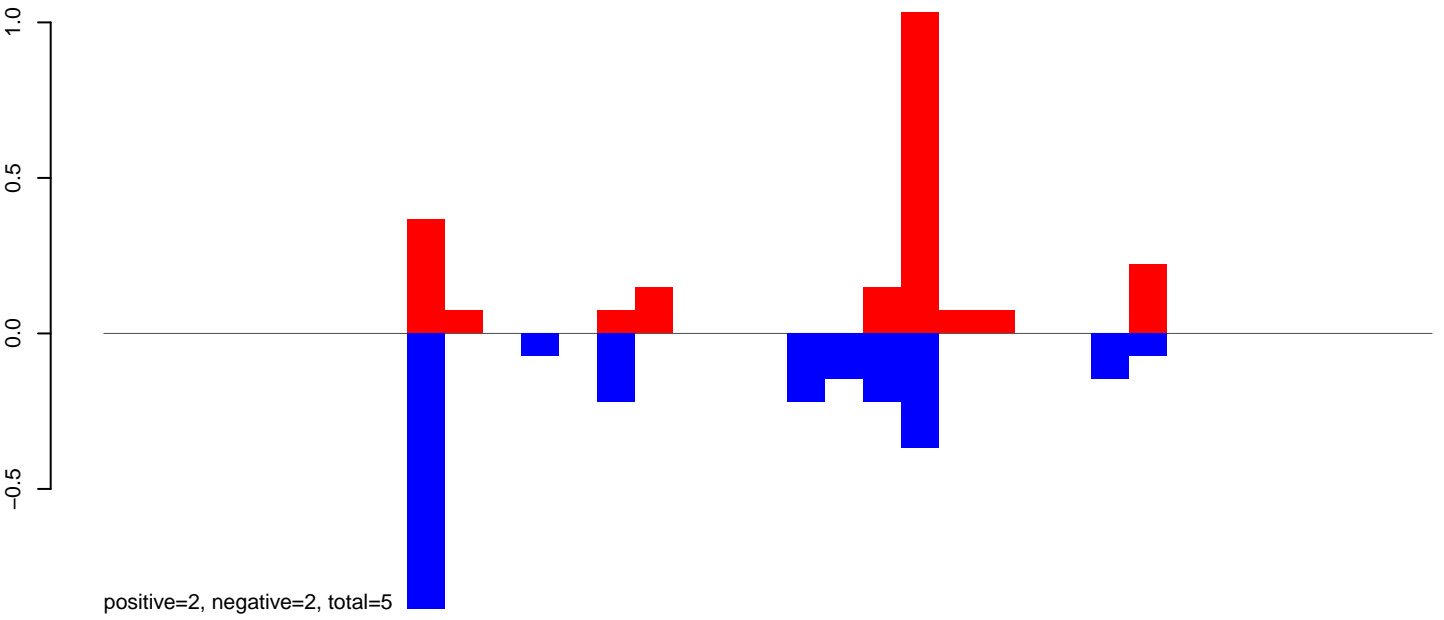
CuQuin_Testes_NL.24_35.rep



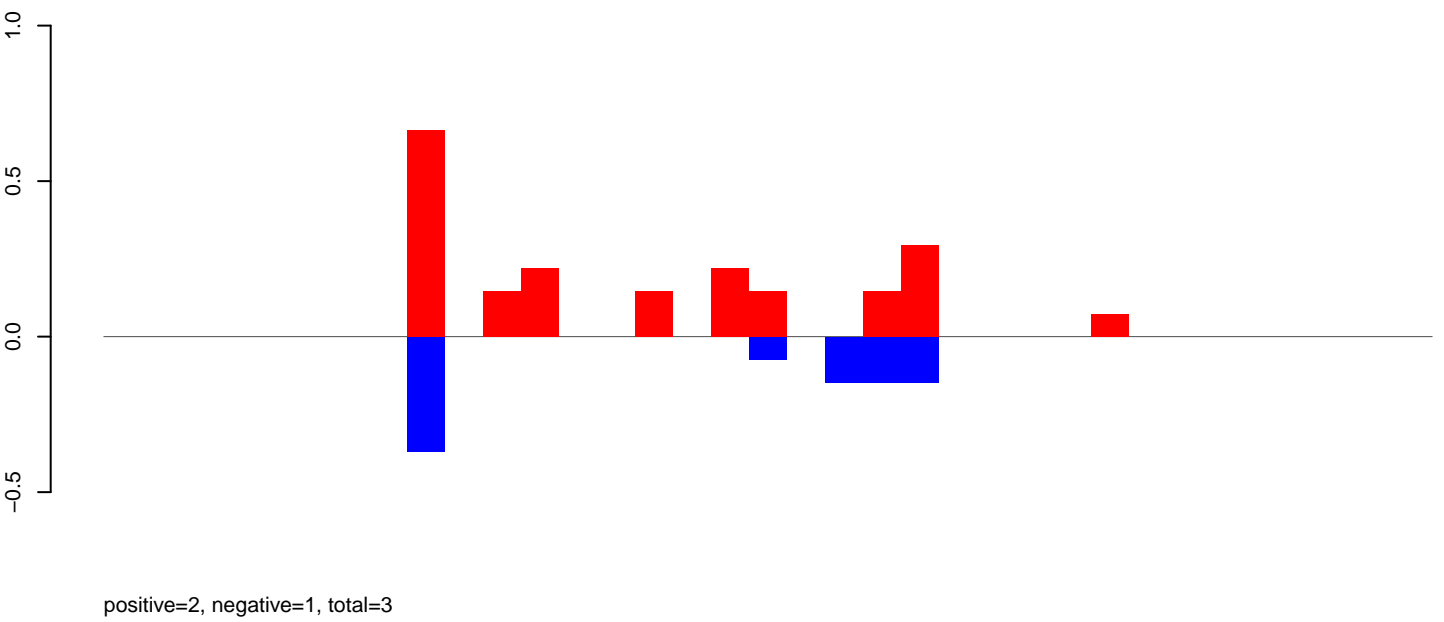
CuQuin_Testes_NL.rep



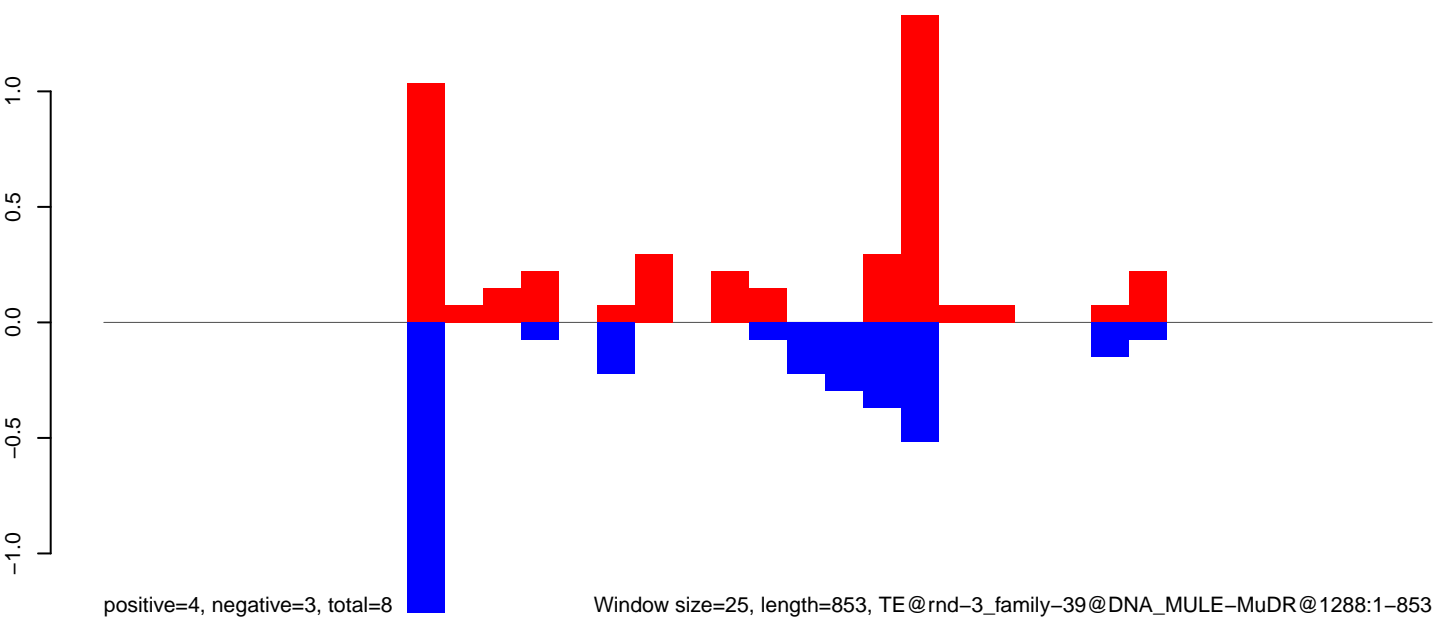
CuQuin_Testes_NL.18_23.rep



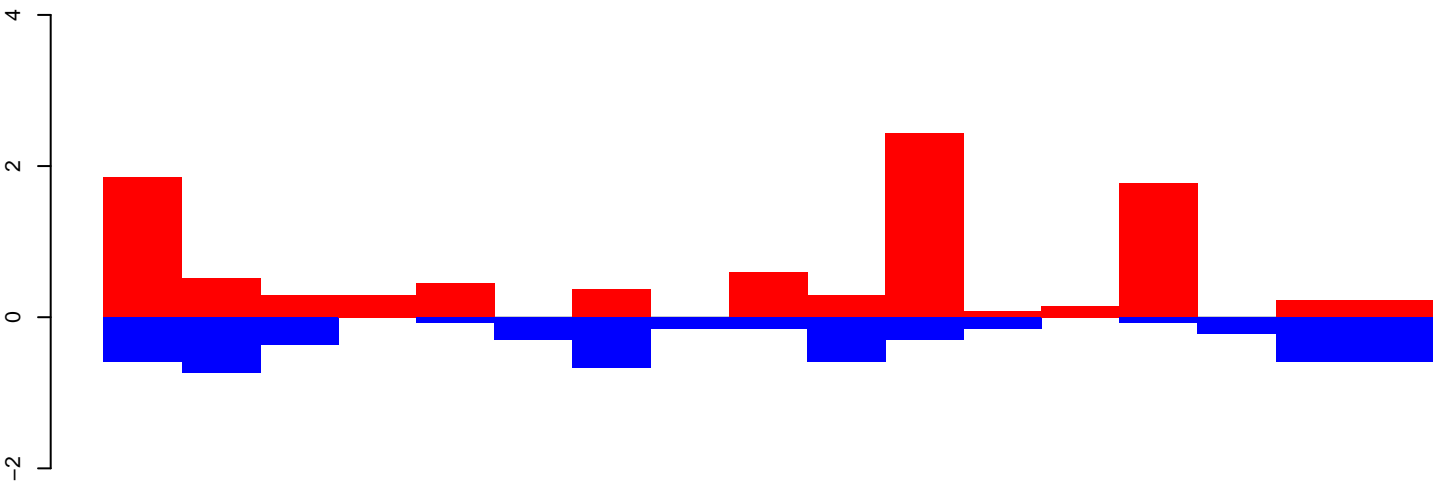
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

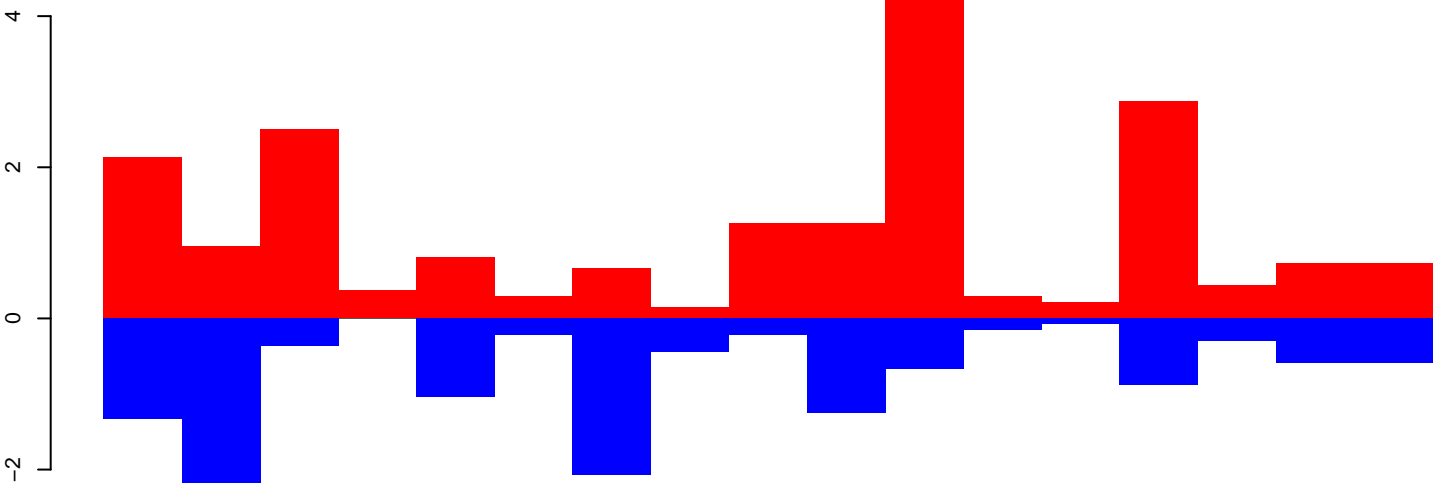


CuQuin_Testes_NL.18_23.rep



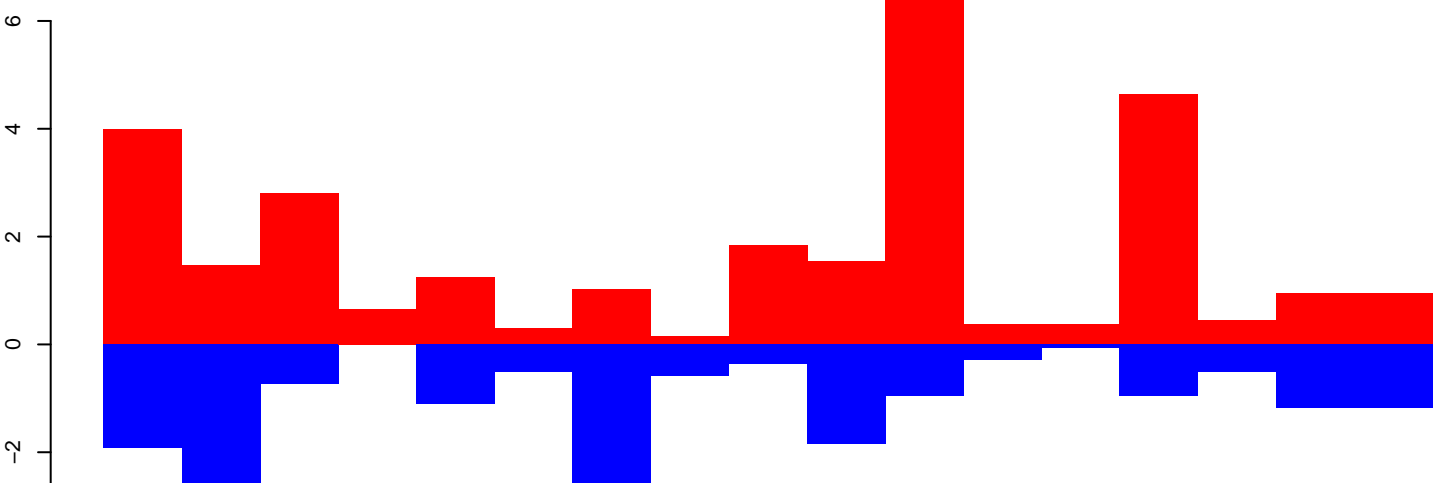
positive=10, negative=6, total=15

CuQuin_Testes_NL.24_35.rep



positive=21, negative=13, total=34

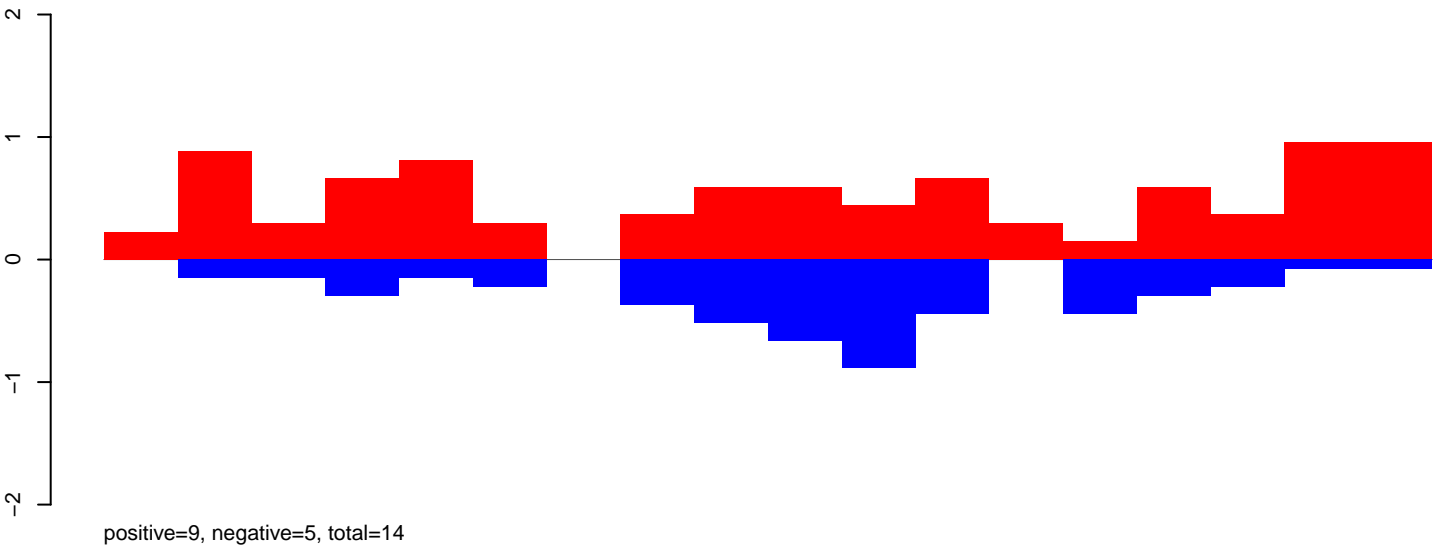
CuQuin_Testes_NL.rep



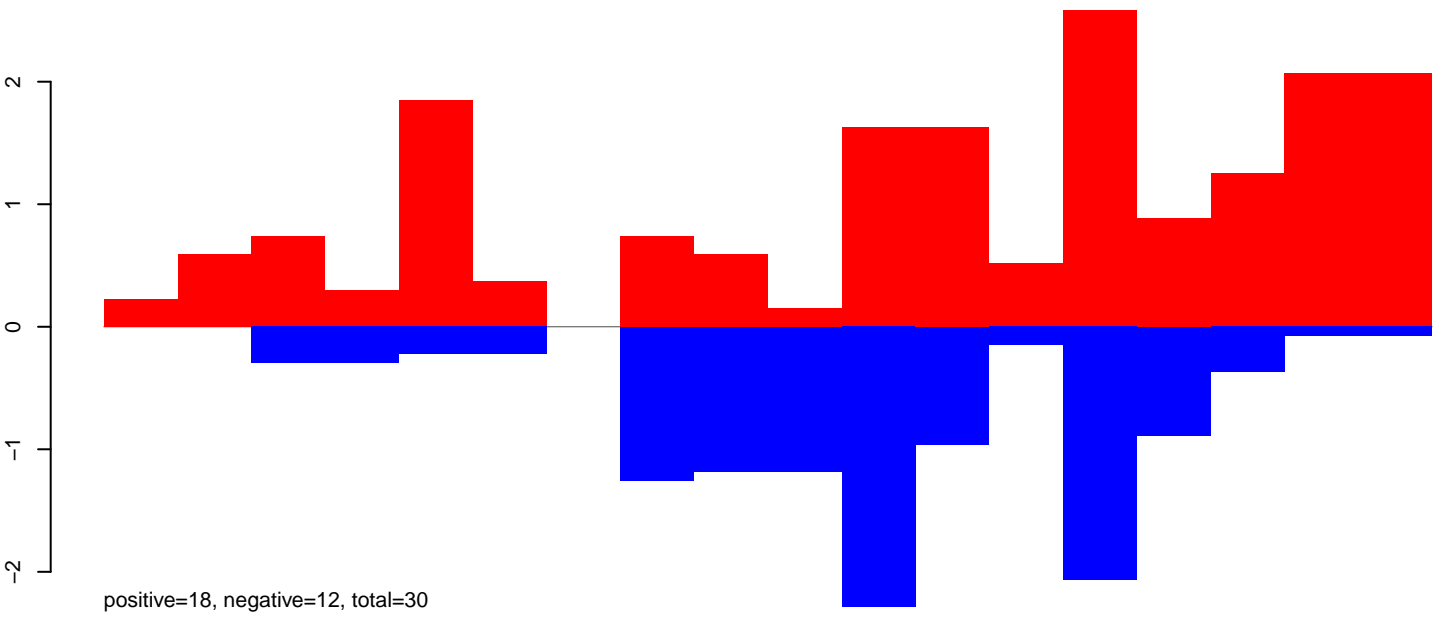
positive=30, negative=19, total=49

Window size=25, length=421, TE@rnd-3_family-326@DNA_TcMar@48:1-421

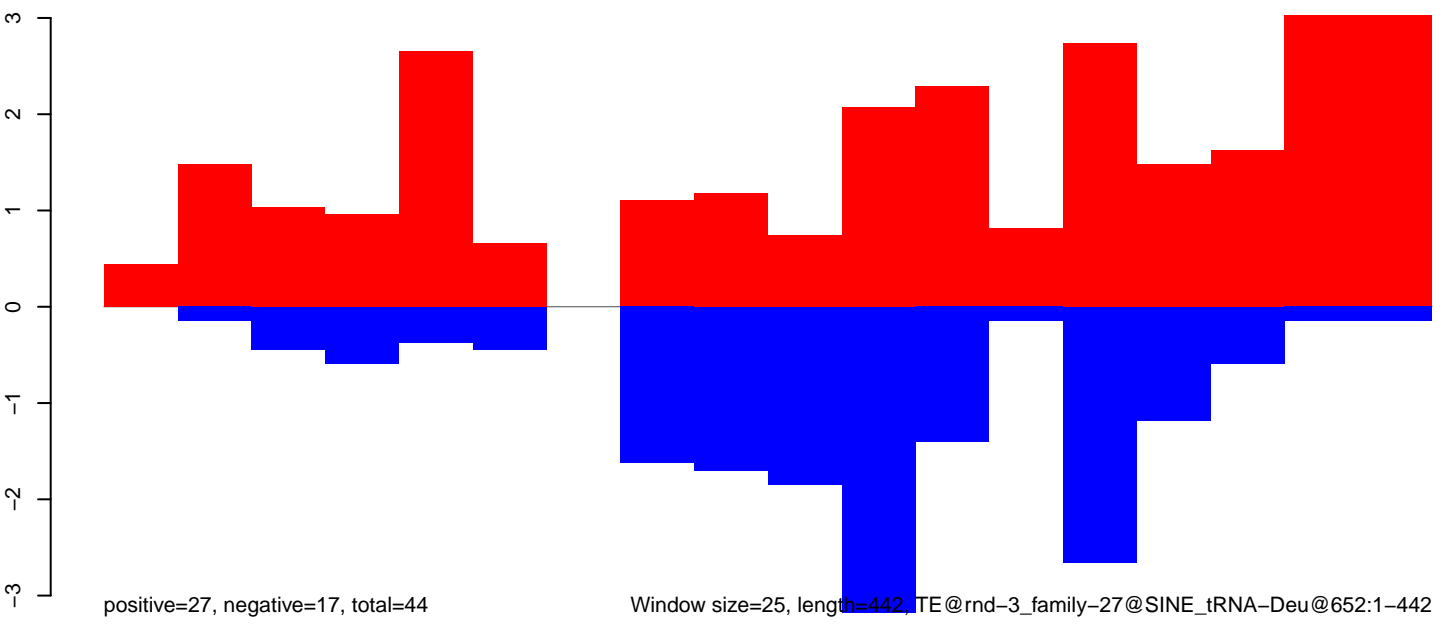
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



Window size=25, length=442, TE@rnd-3_family-27@SINE_tRNA-Deu@652:1-442

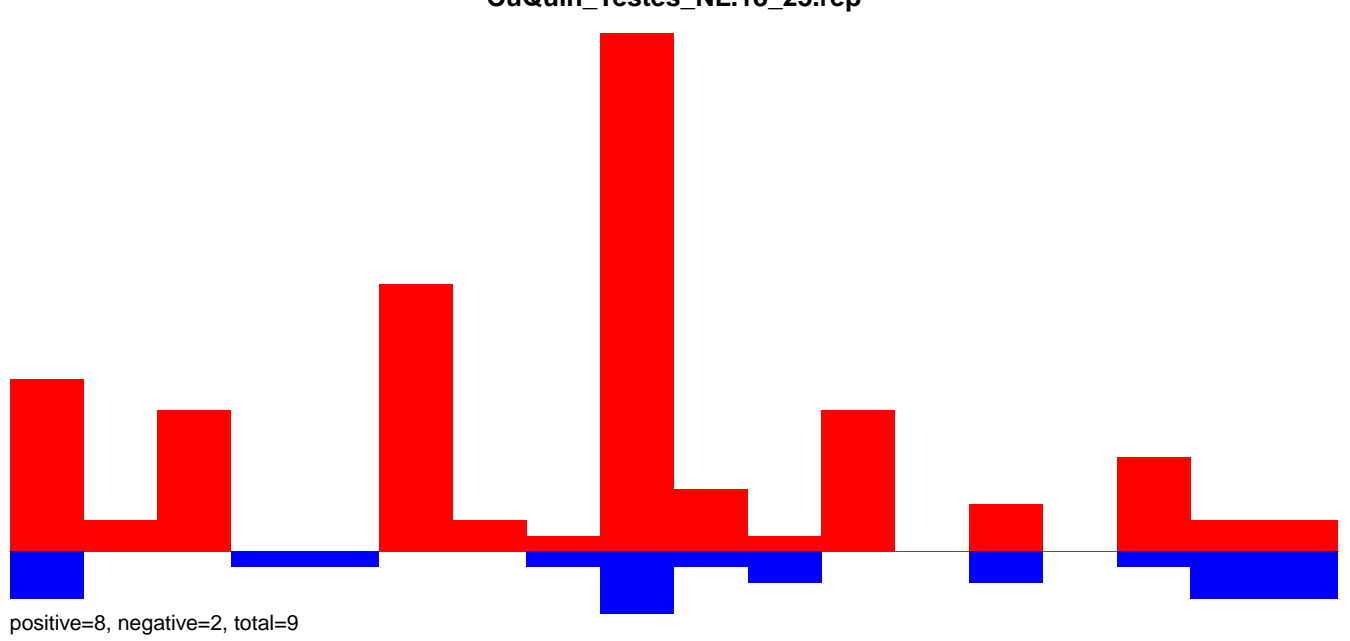
100

200

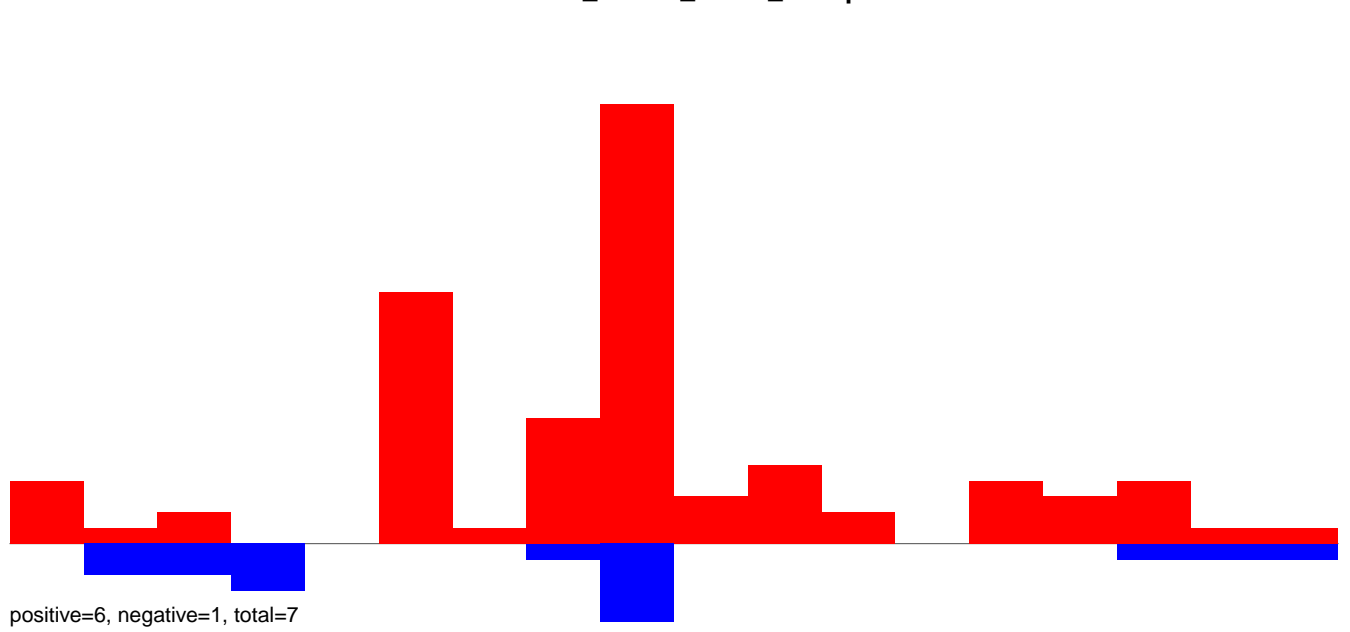
300

400

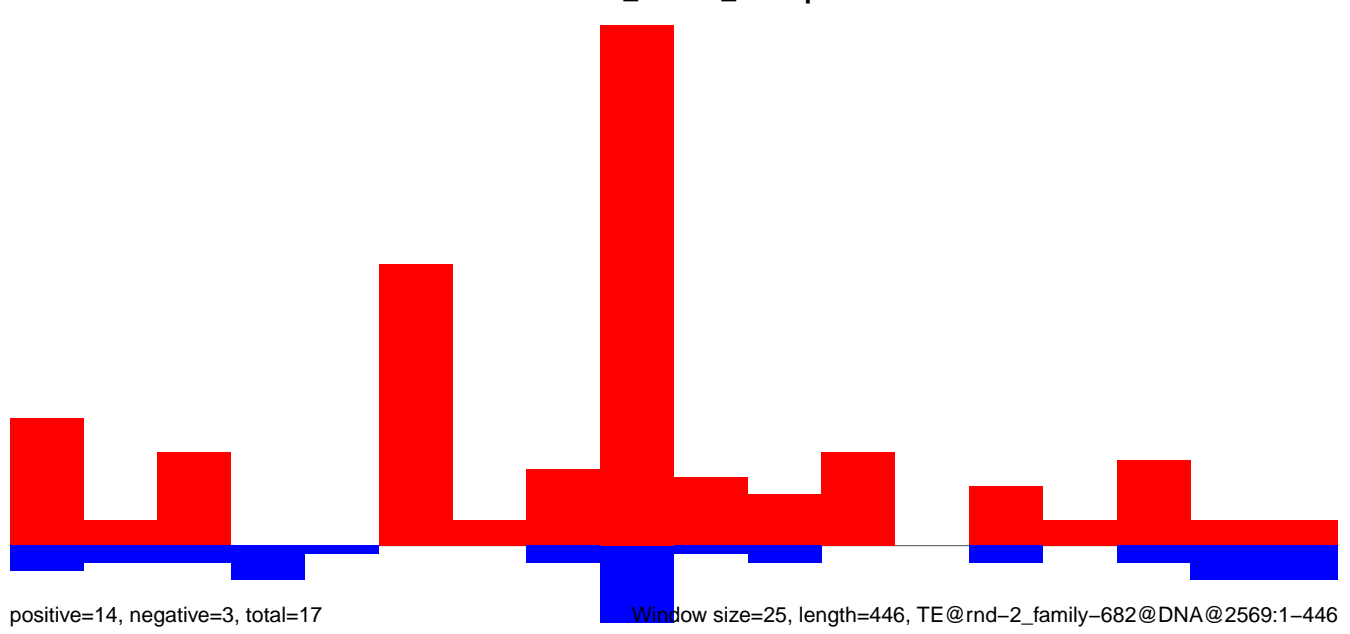
CuQuin_Testes_NL.18_23.rep



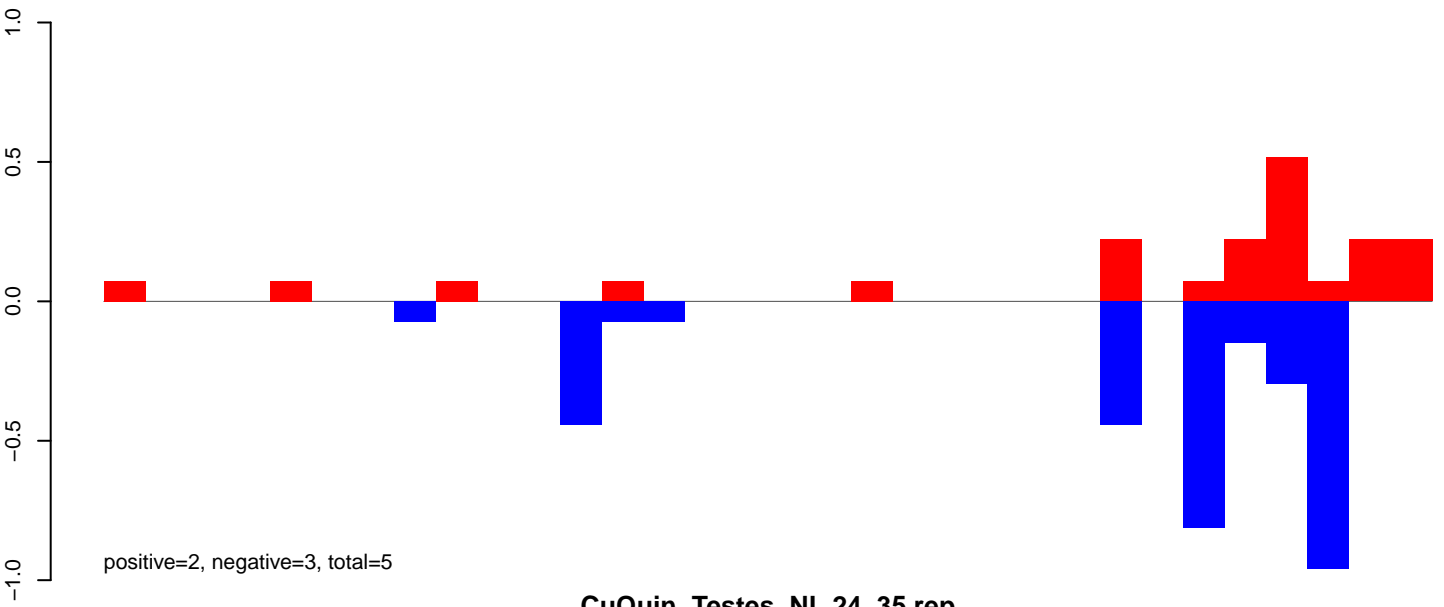
CuQuin_Testes_NL.24_35.rep



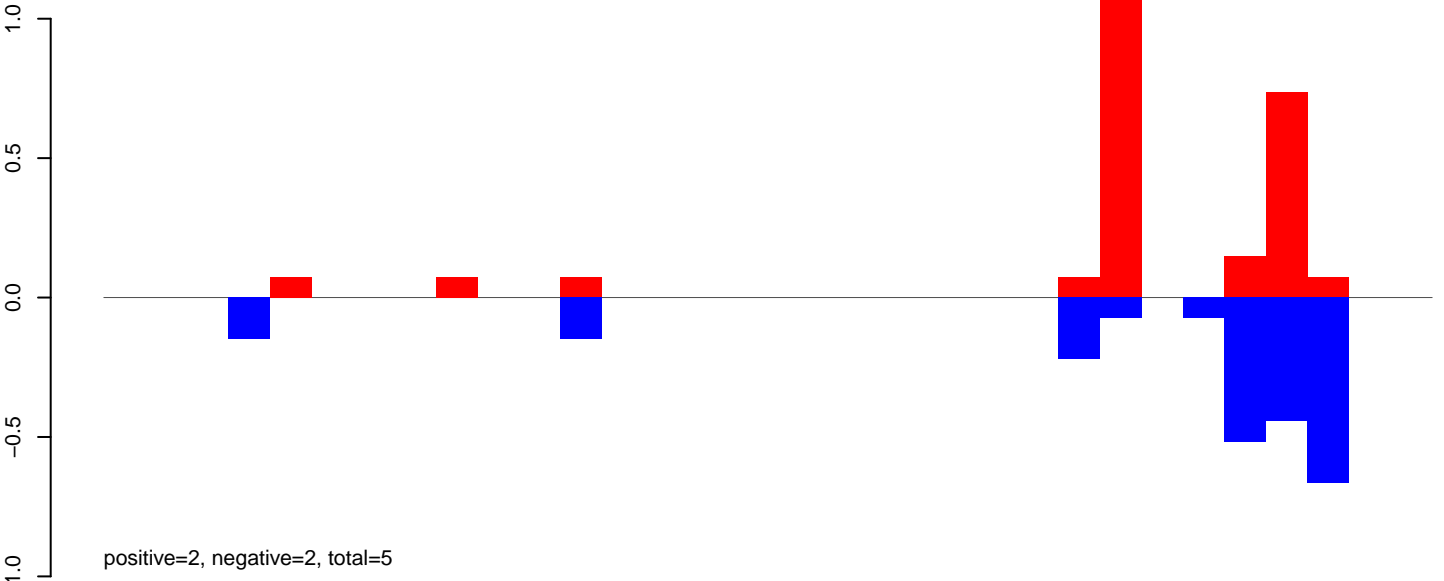
CuQuin_Testes_NL.rep



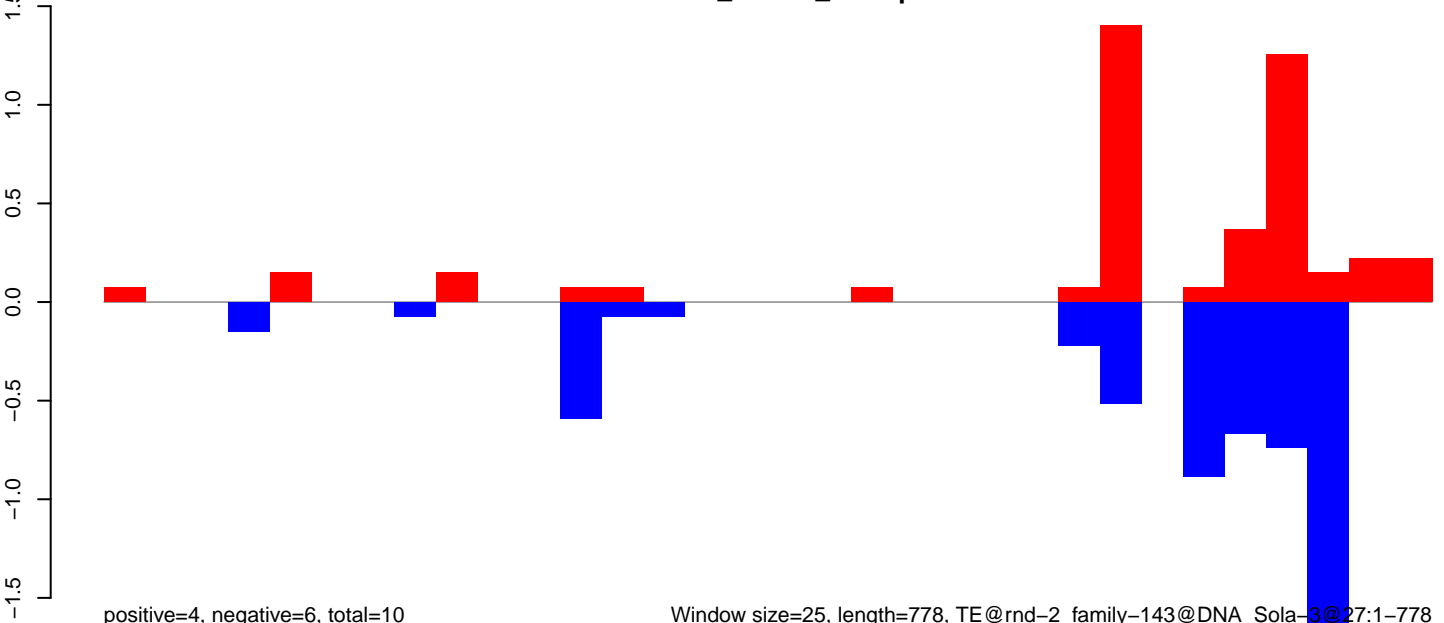
CuQuin_Testes_NL.18_23.rep



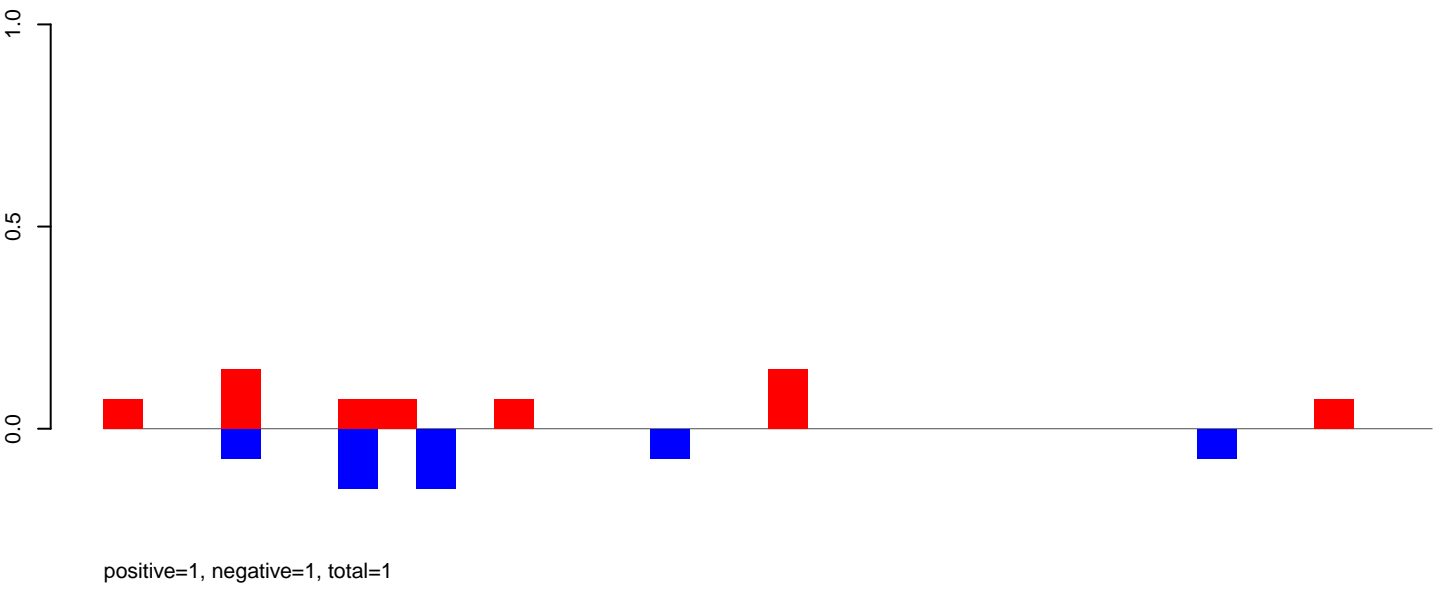
CuQuin_Testes_NL.24_35.rep



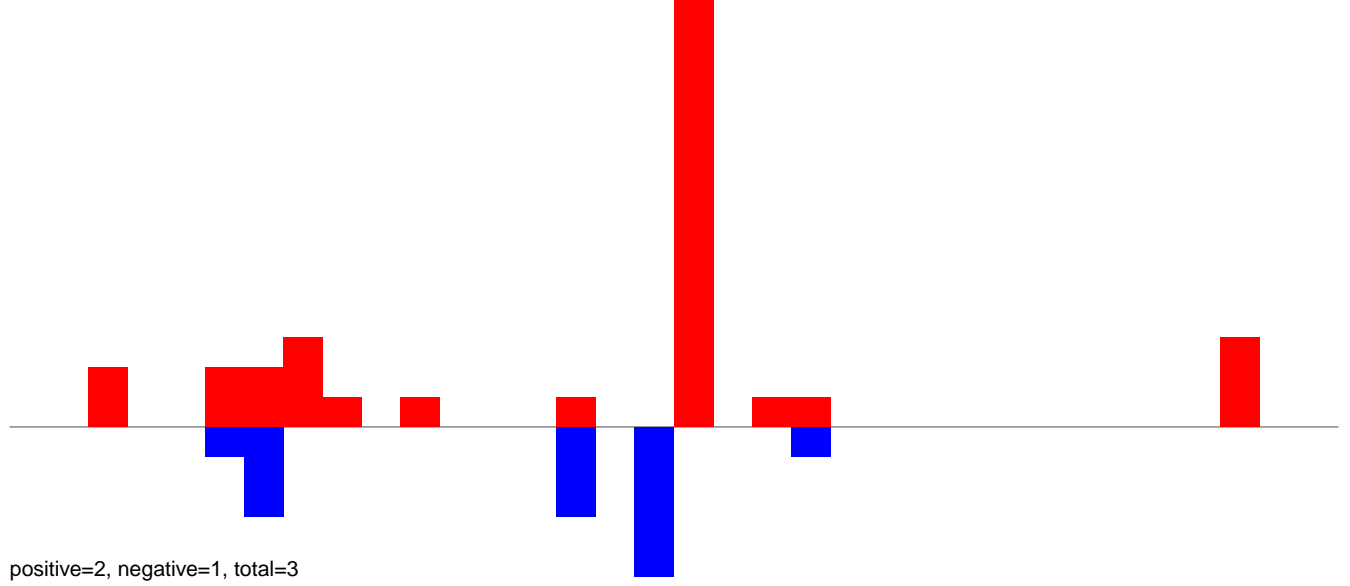
CuQuin_Testes_NL.rep



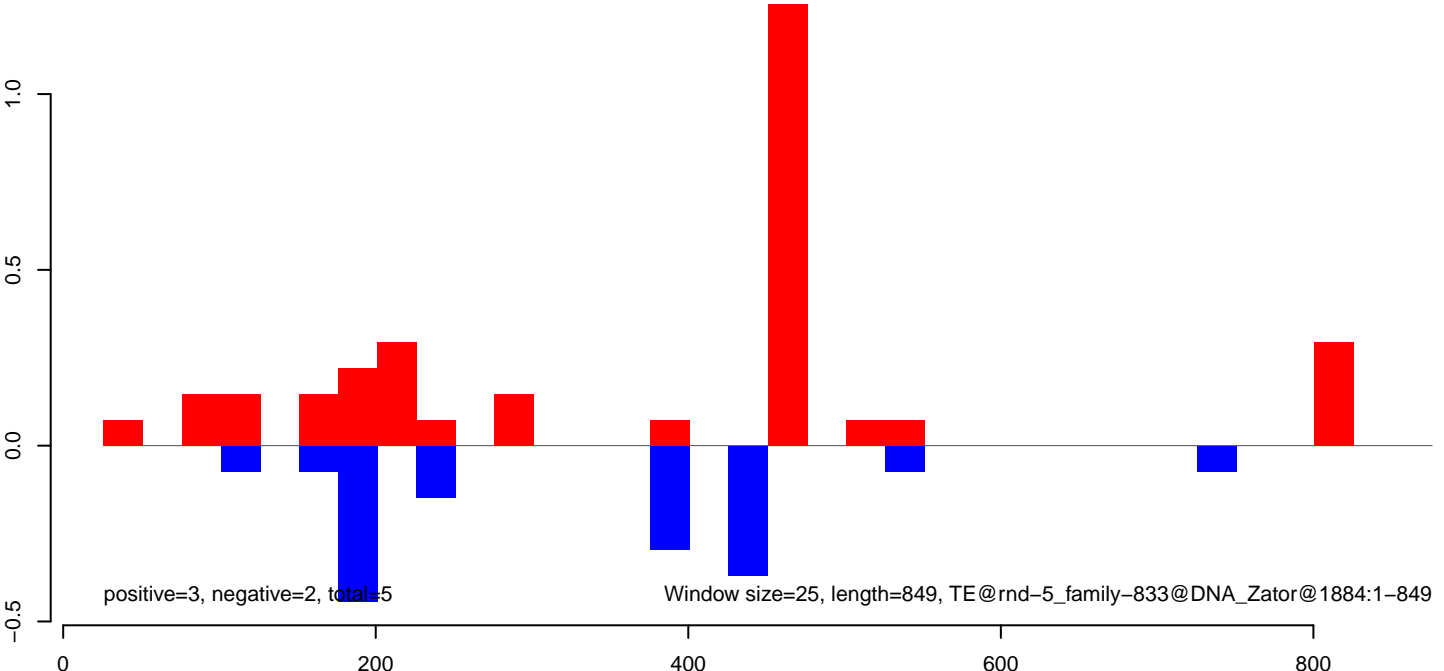
CuQuin_Testes_NL.18_23.rep



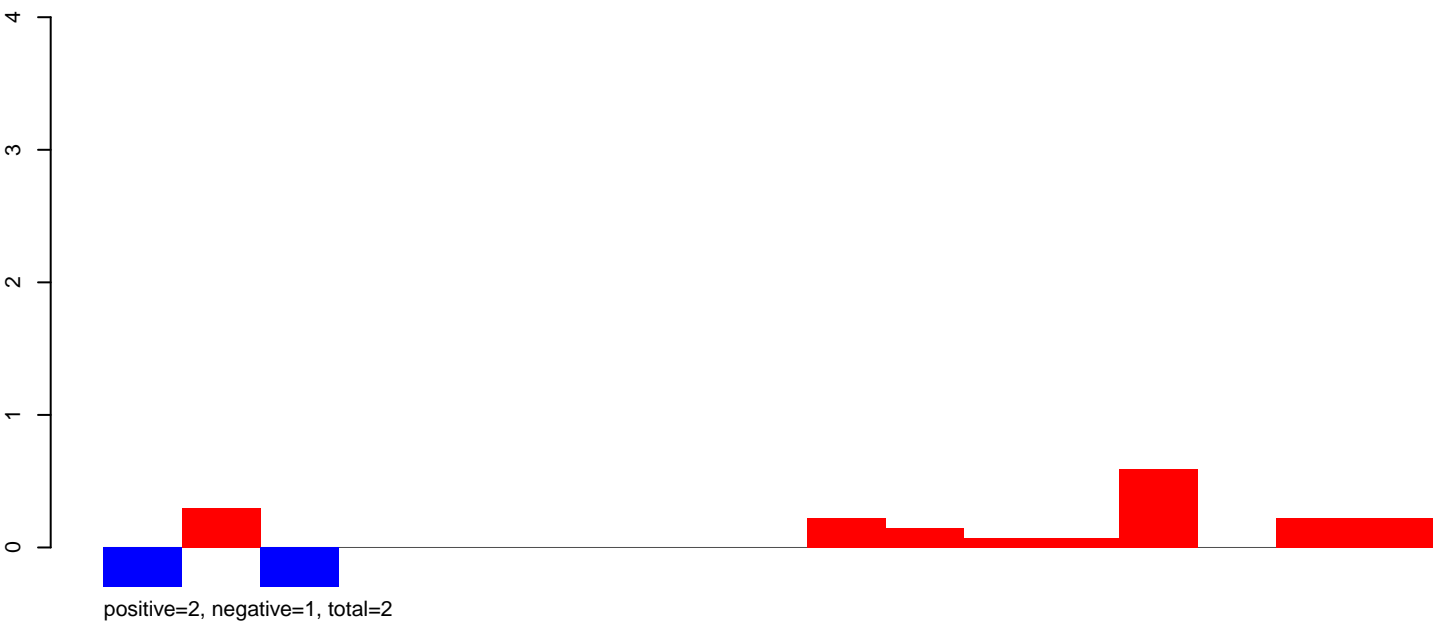
CuQuin_Testes_NL.24_35.rep



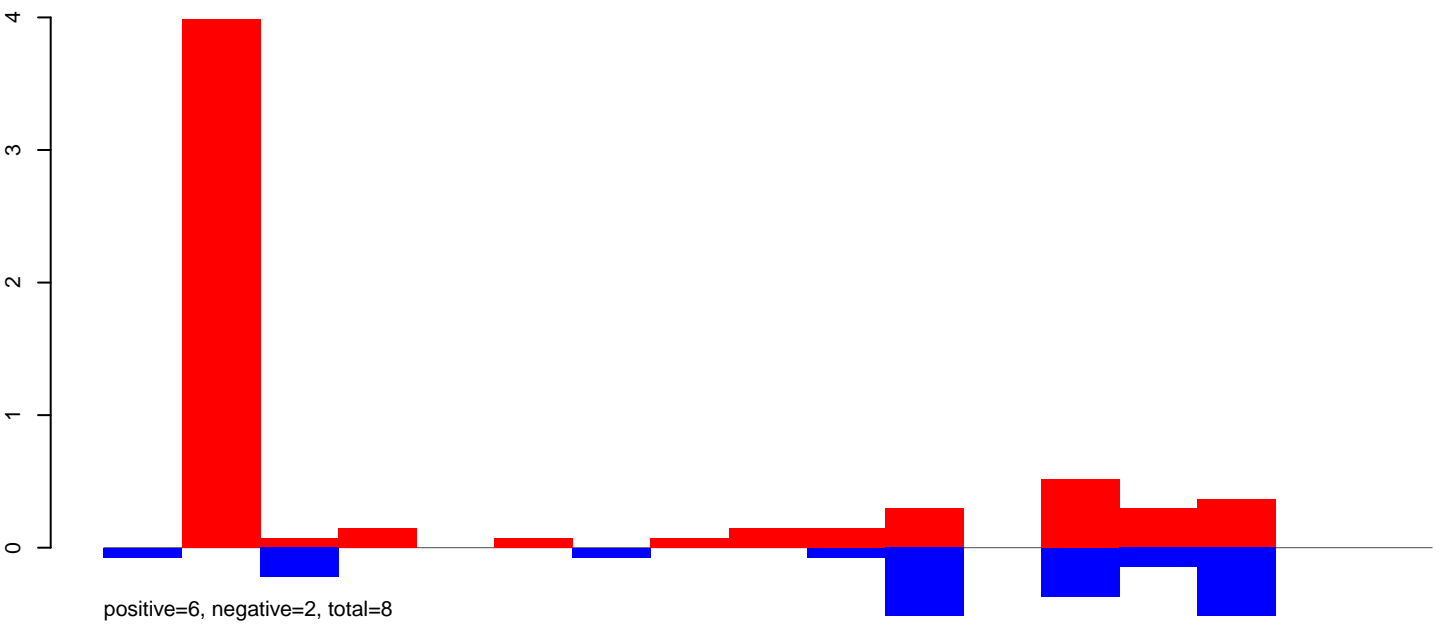
CuQuin_Testes_NL.rep



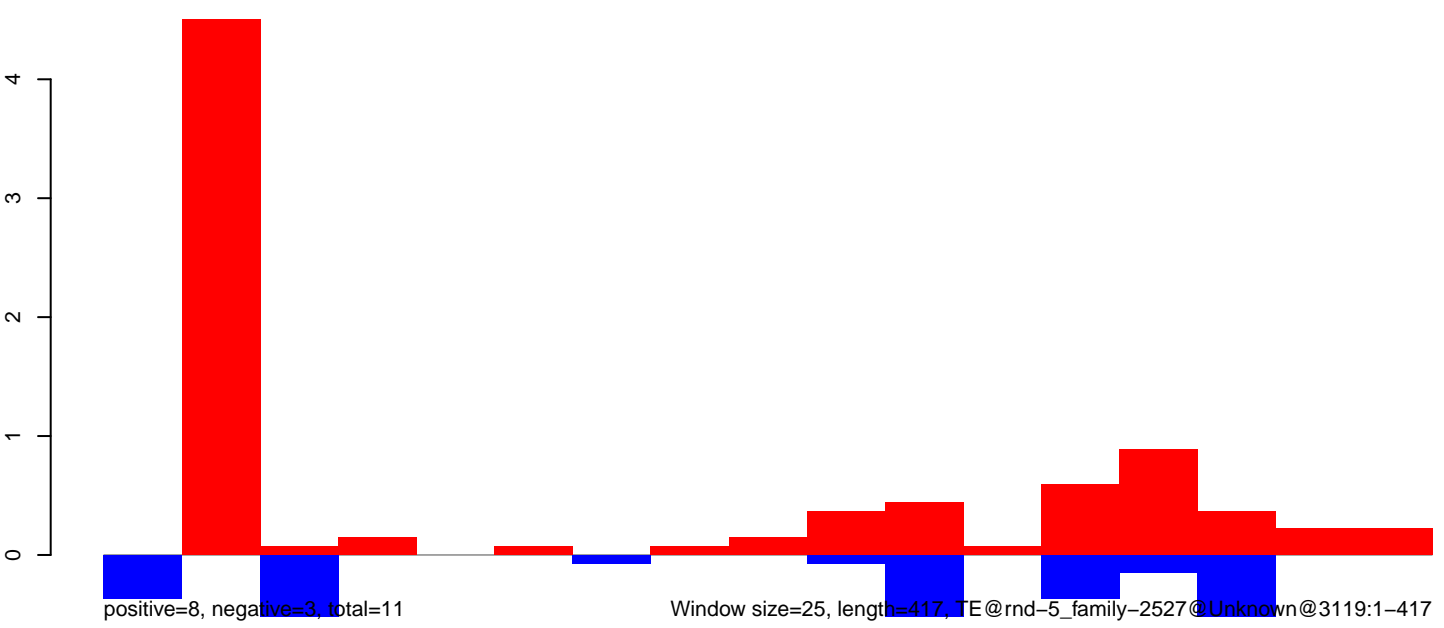
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

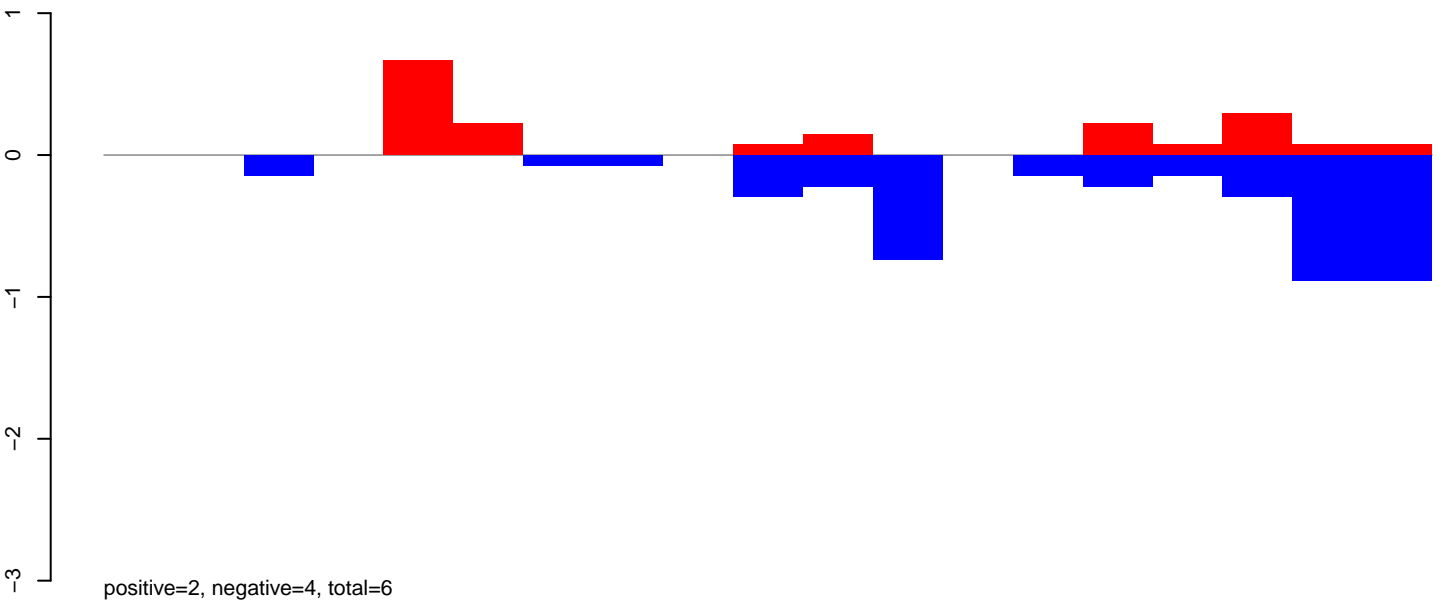


CuQuin_Testes_NL.rep

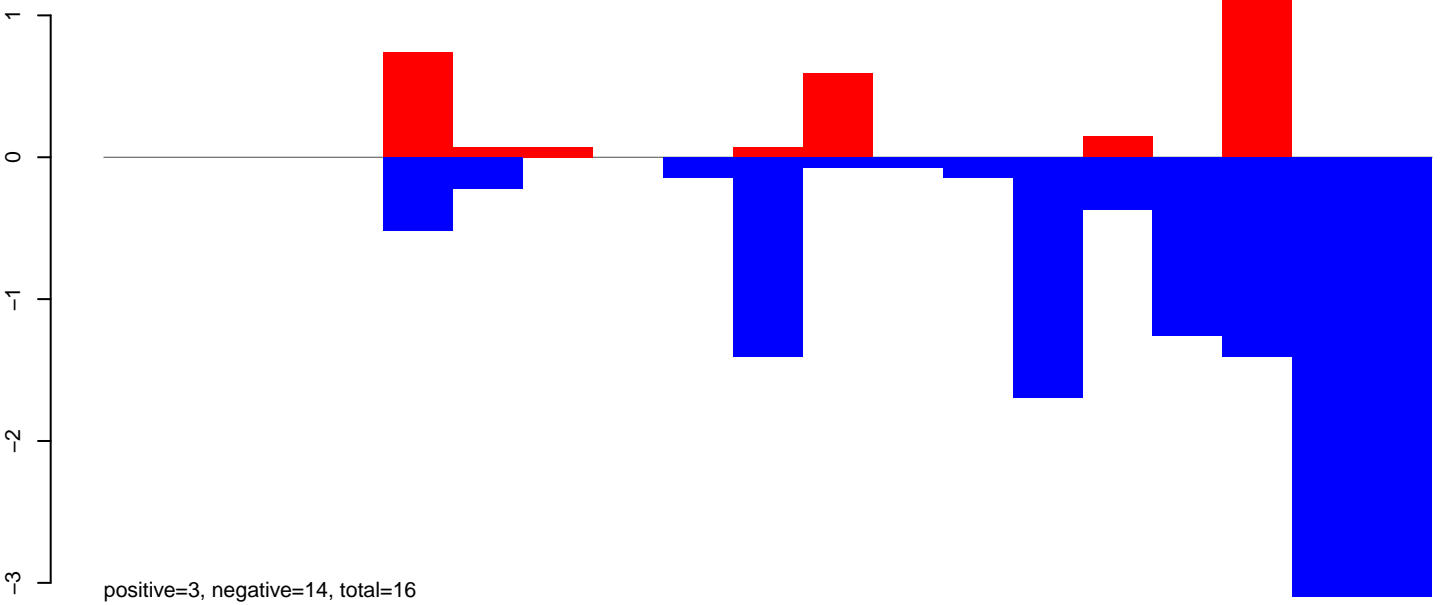


Window size=25, length=417, TE@rnd-5_family-2527@Unknown@3119:1-417

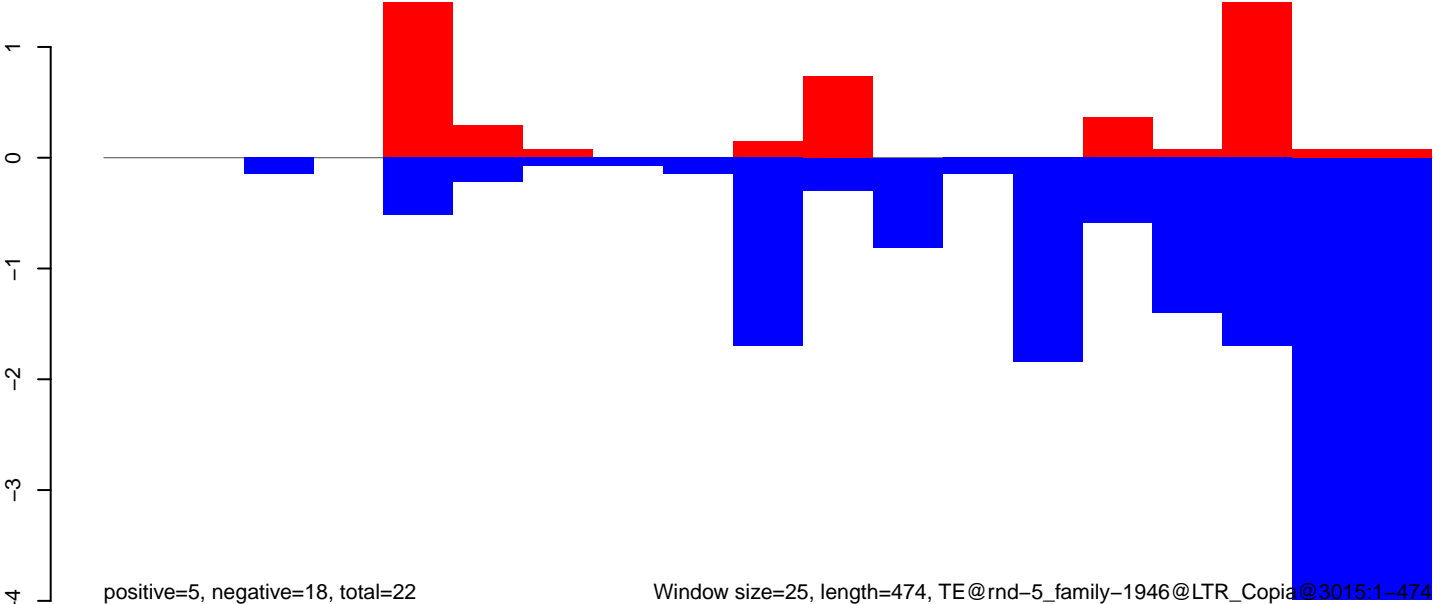
CuQuin_Testes_NL.18_23.rep



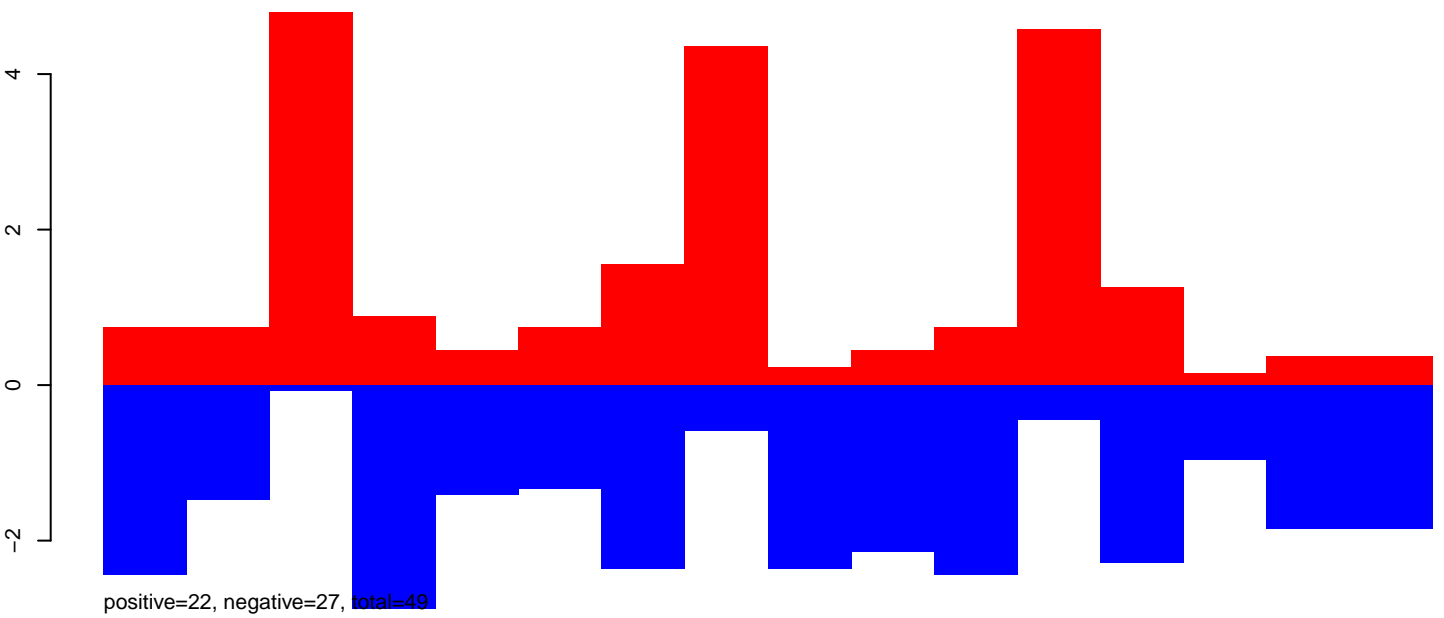
CuQuin_Testes_NL.24_35.rep



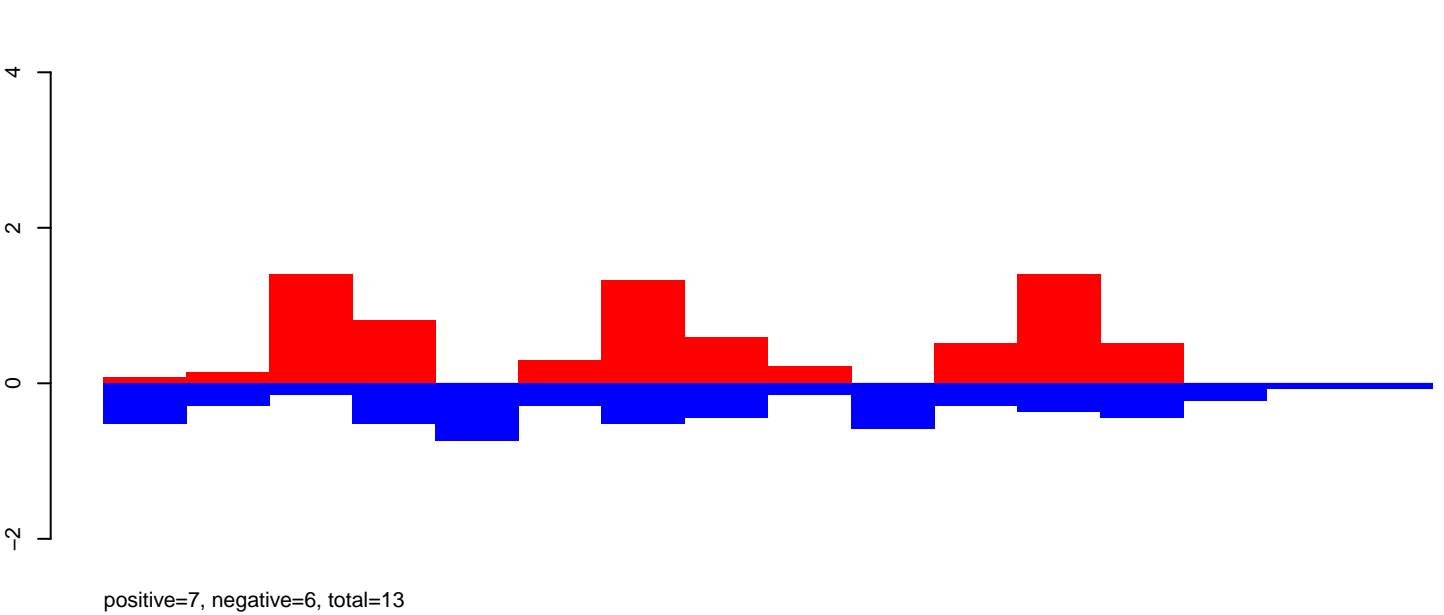
CuQuin_Testes_NL.rep



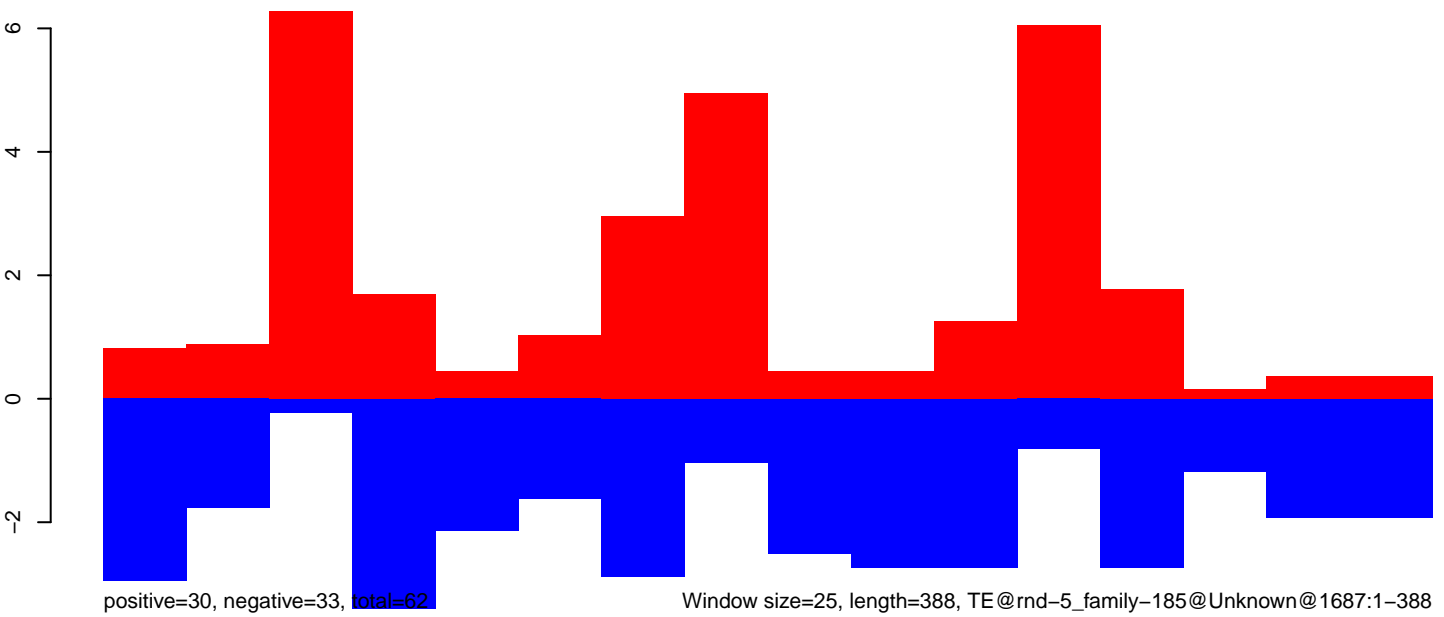
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

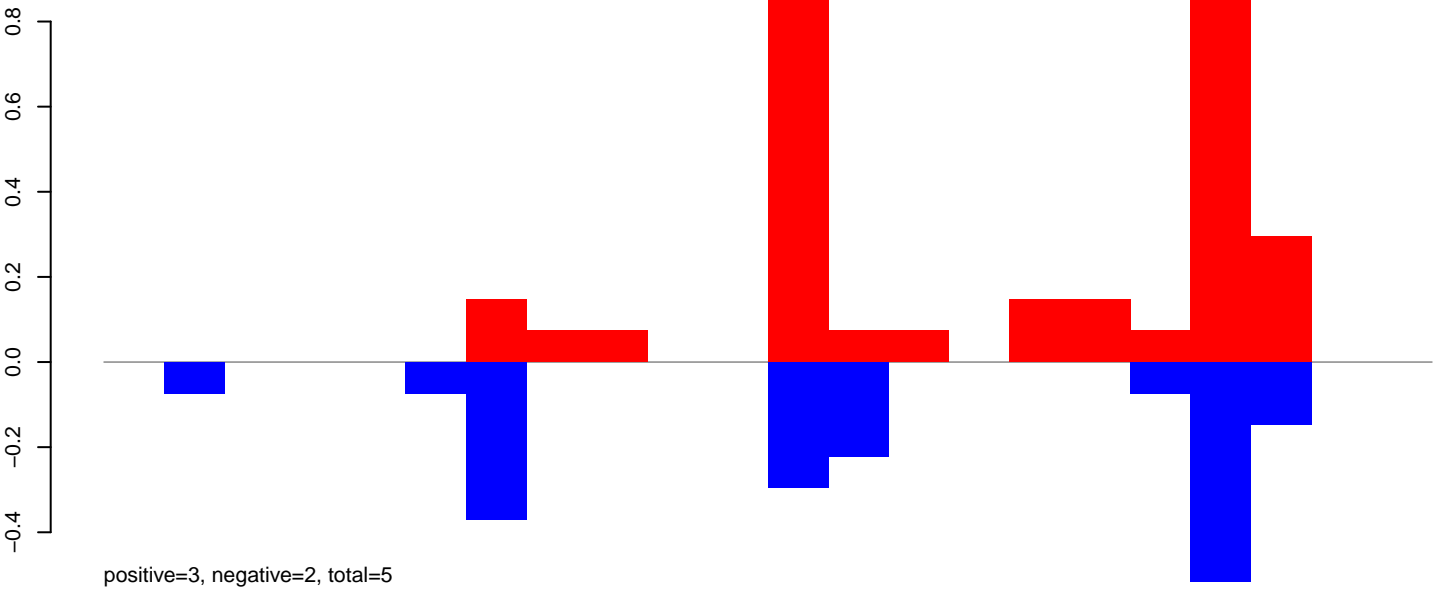


CuQuin_Testes_NL.rep

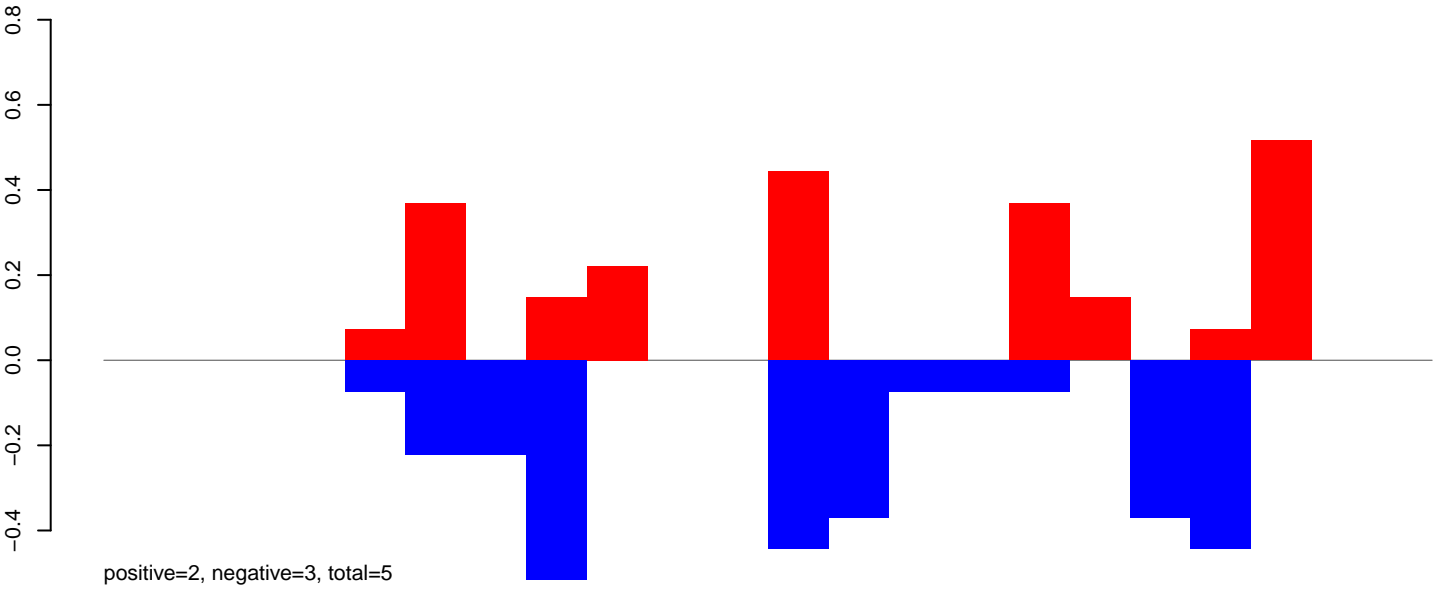


Window size=25, length=388, TE@rnd-5_family-185@Unknown@1687:1-388

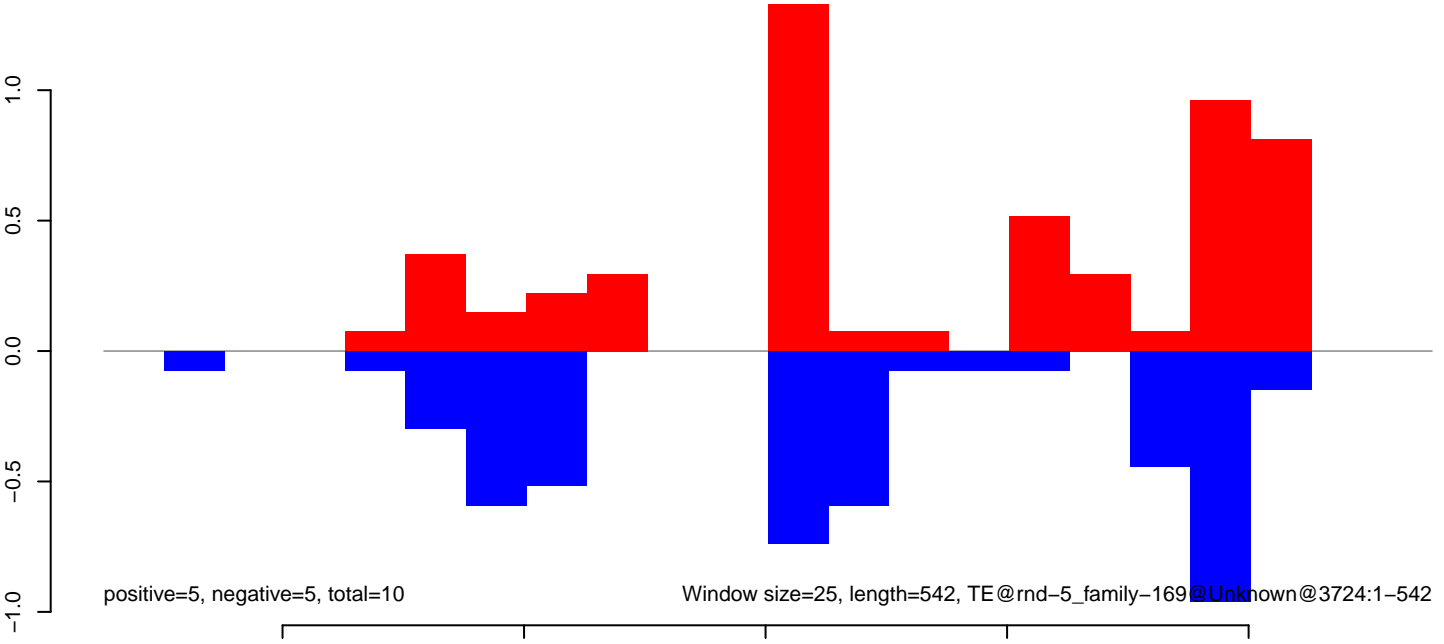
CuQuin_Testes_NL.18_23.rep



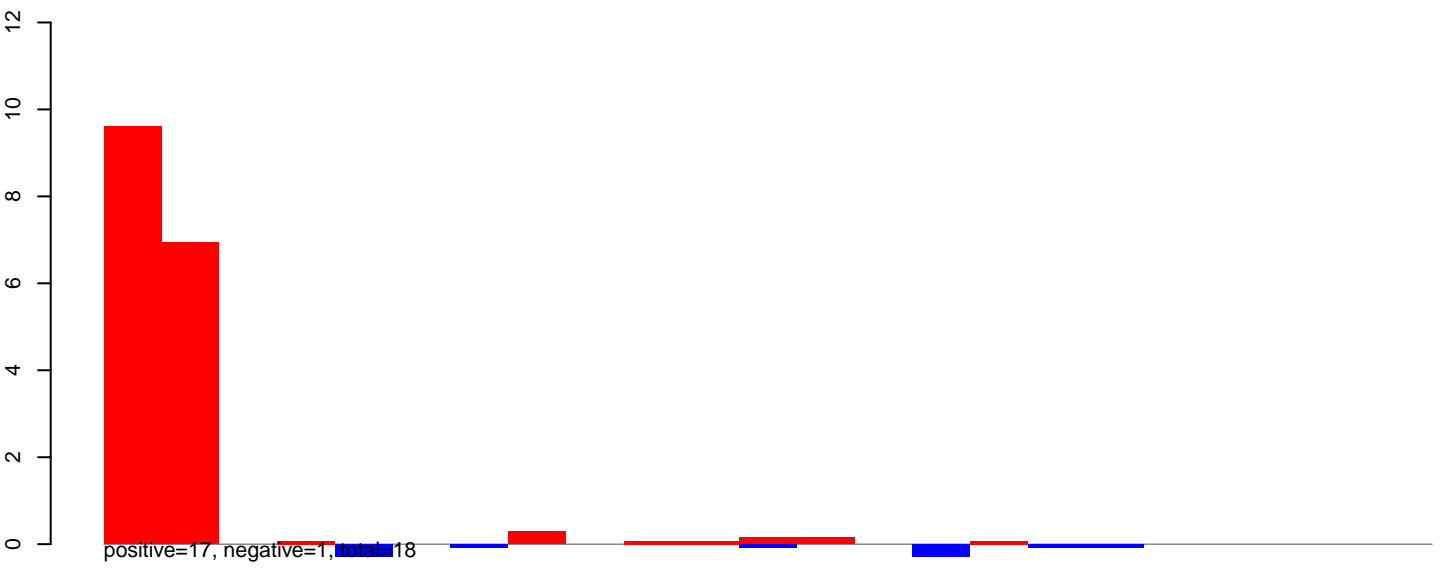
CuQuin_Testes_NL.24_35.rep



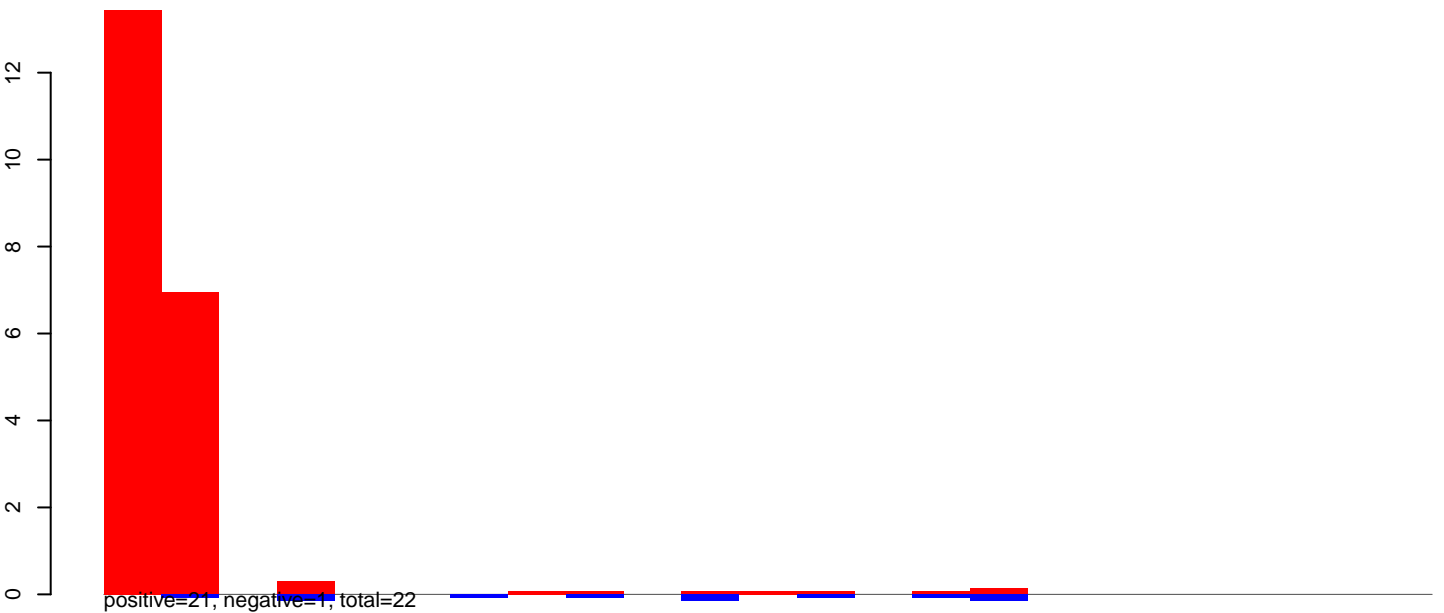
CuQuin_Testes_NL.rep



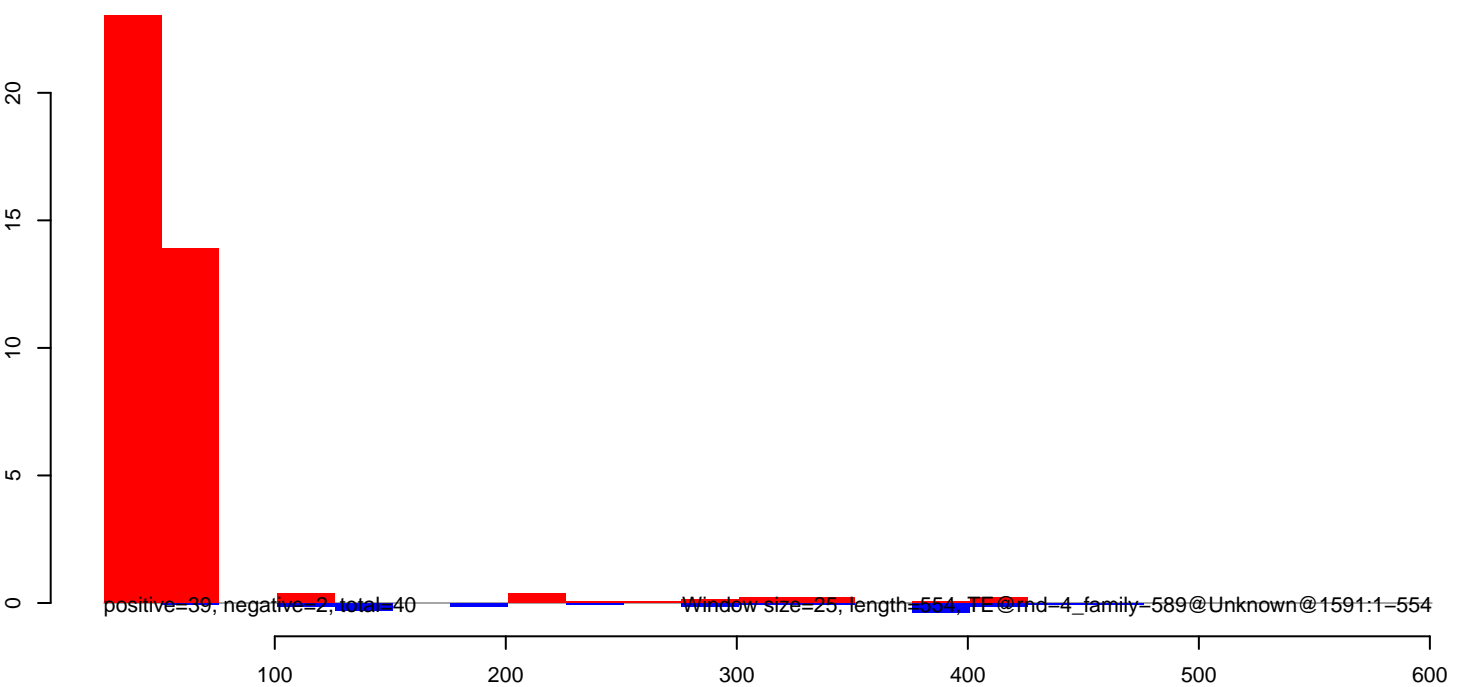
CuQuin_Testes_NL.18_23.rep



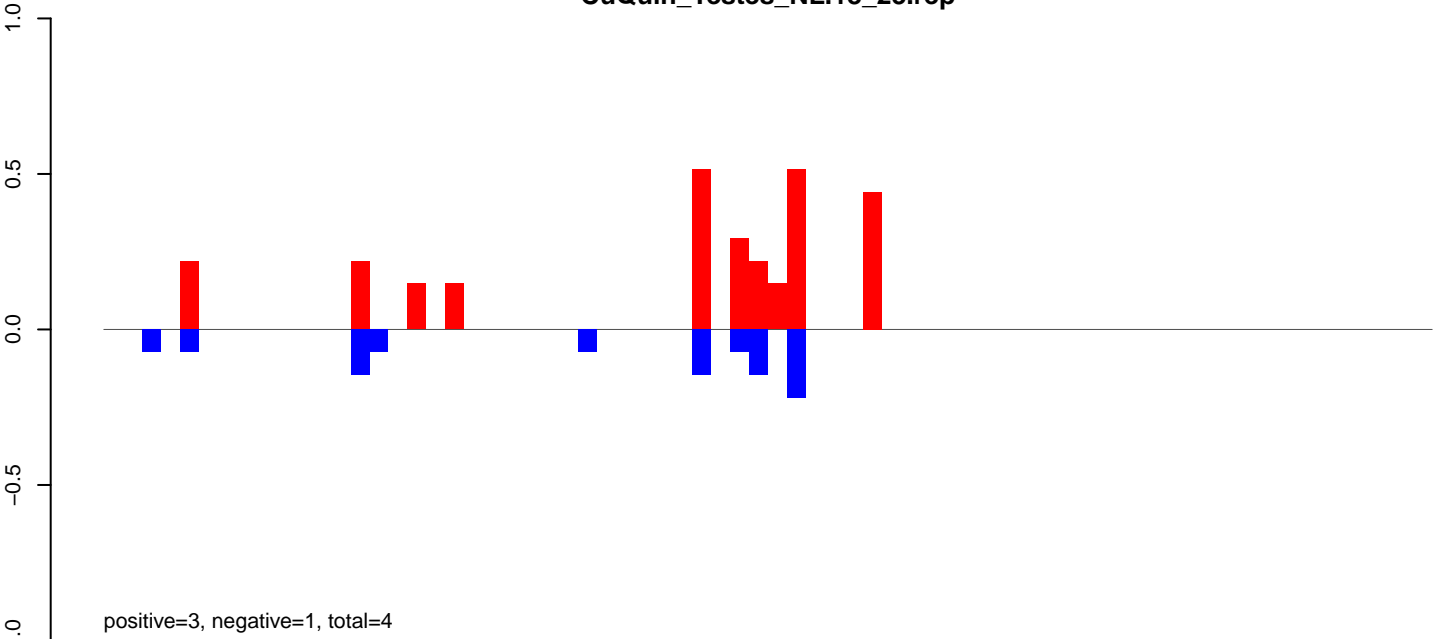
CuQuin_Testes_NL.24_35.rep



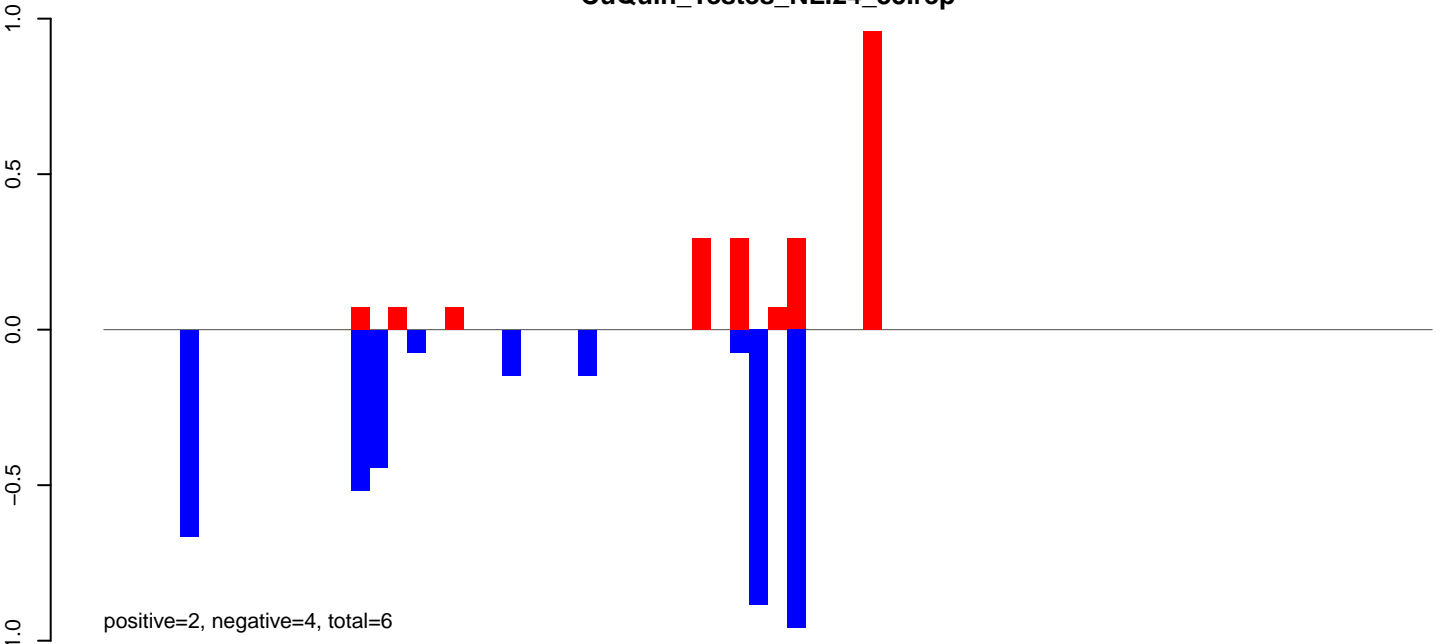
CuQuin_Testes_NL.rep



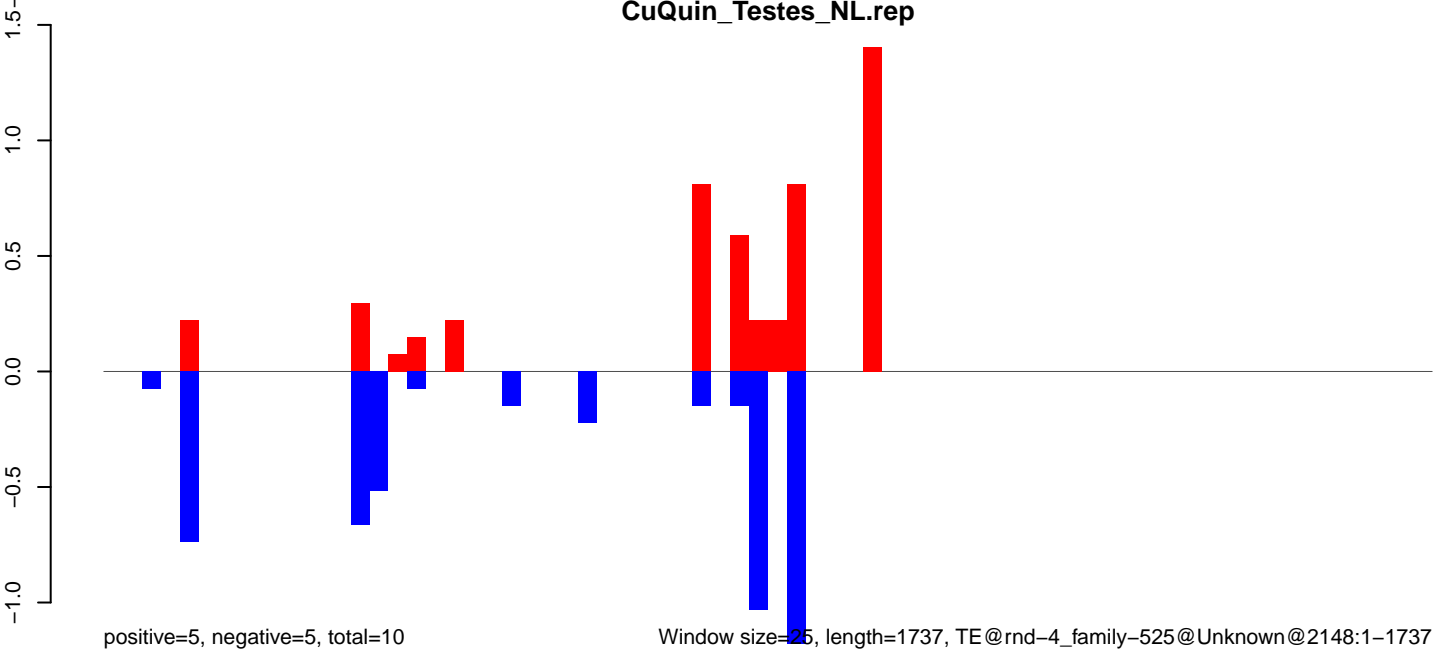
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

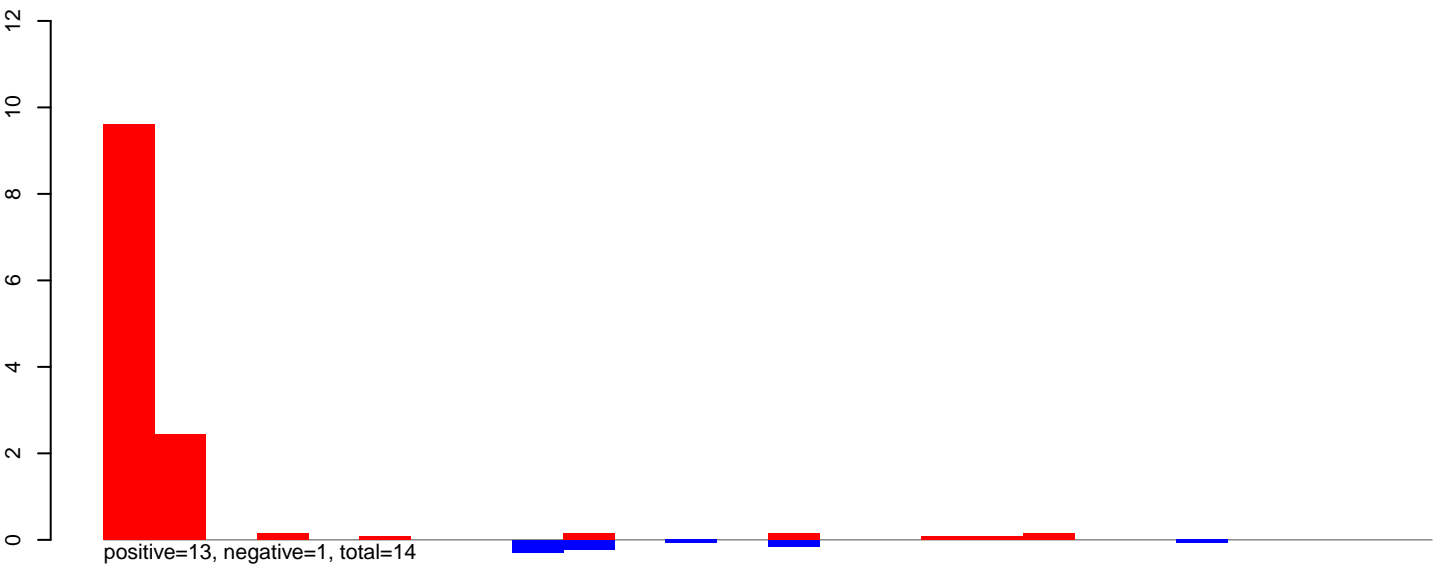


CuQuin_Testes_NL.rep

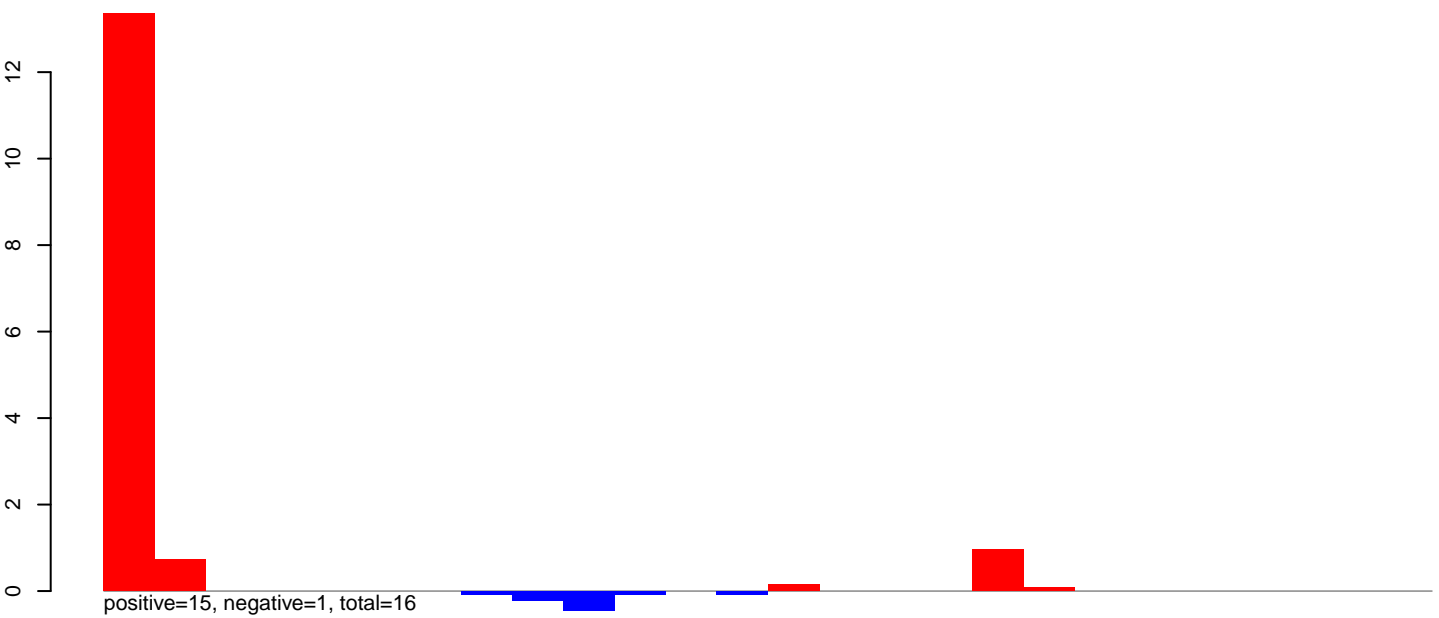


Window size=25, length=1737, TE@rnd-4_family-525@Unknown@2148:1-1737

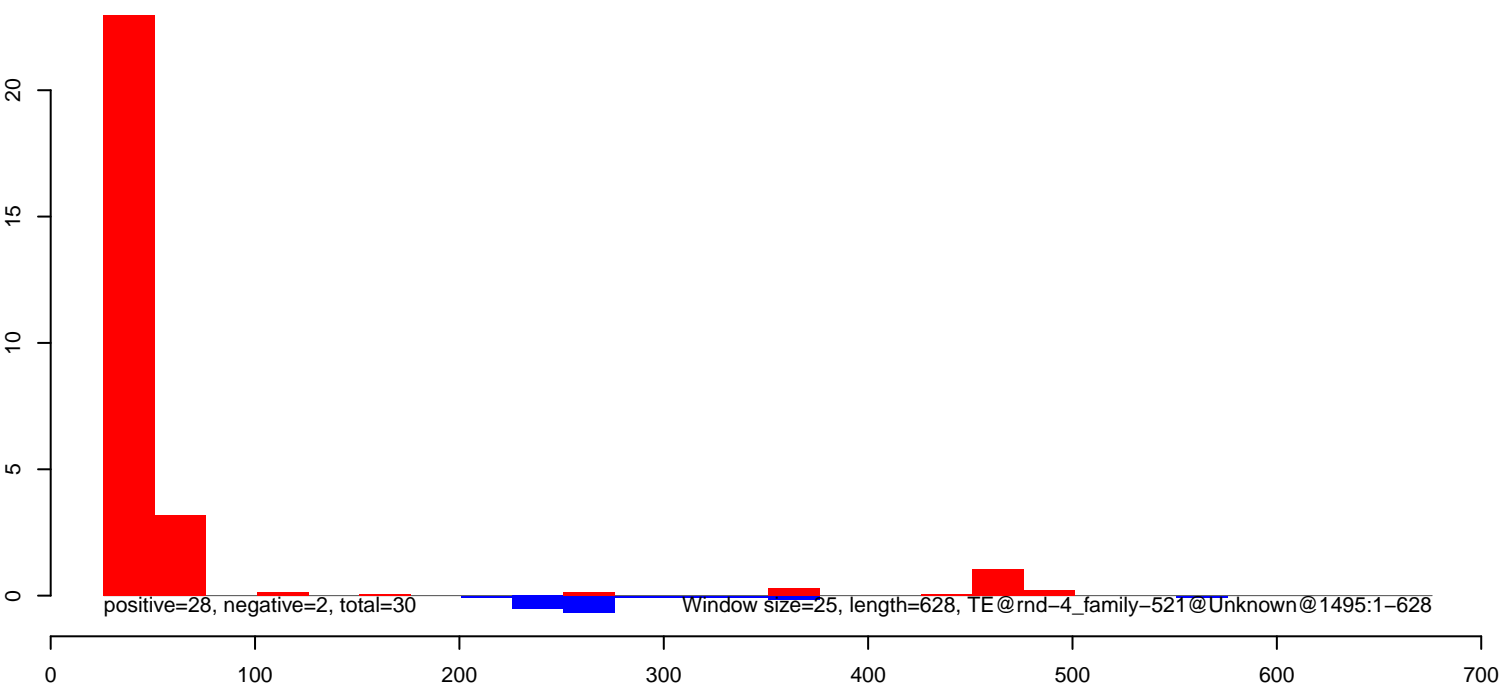
CuQuin_Testes_NL.18_23.rep



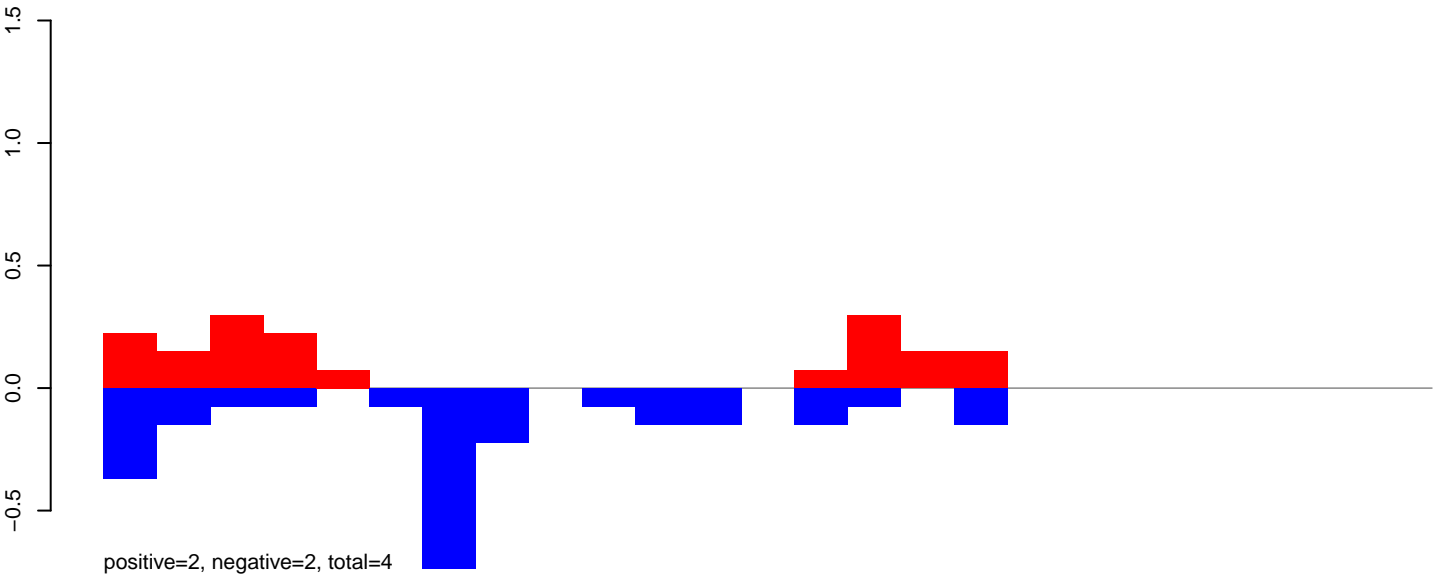
CuQuin_Testes_NL.24_35.rep



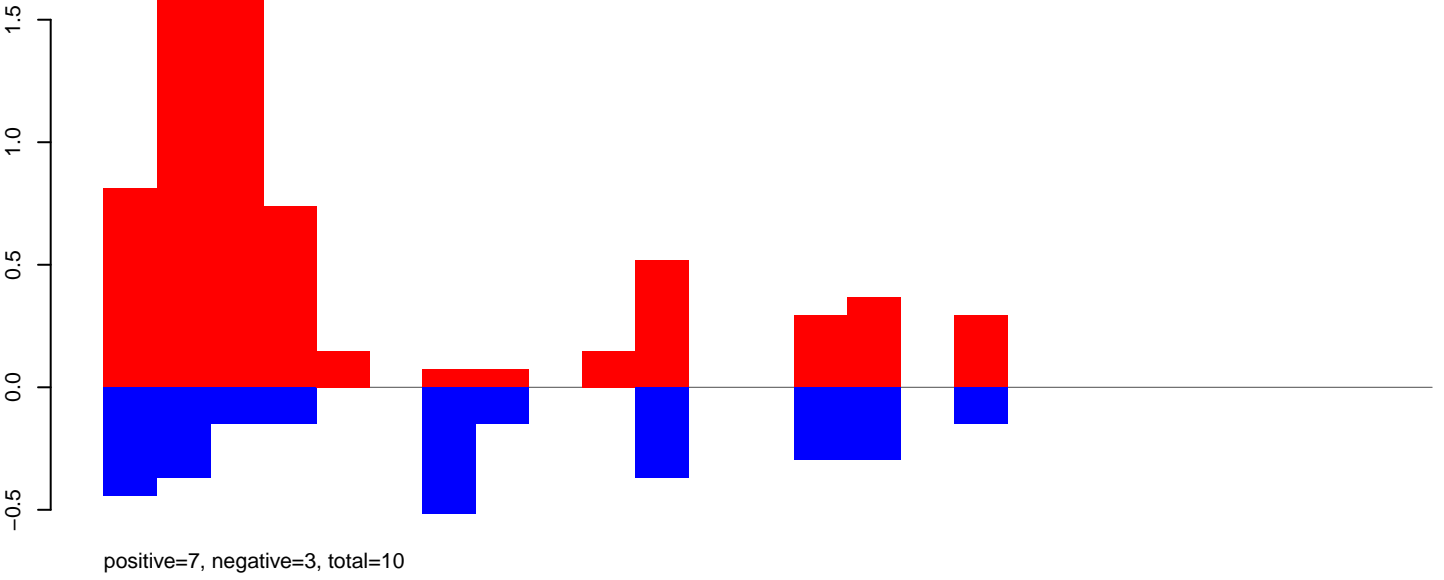
CuQuin_Testes_NL.rep



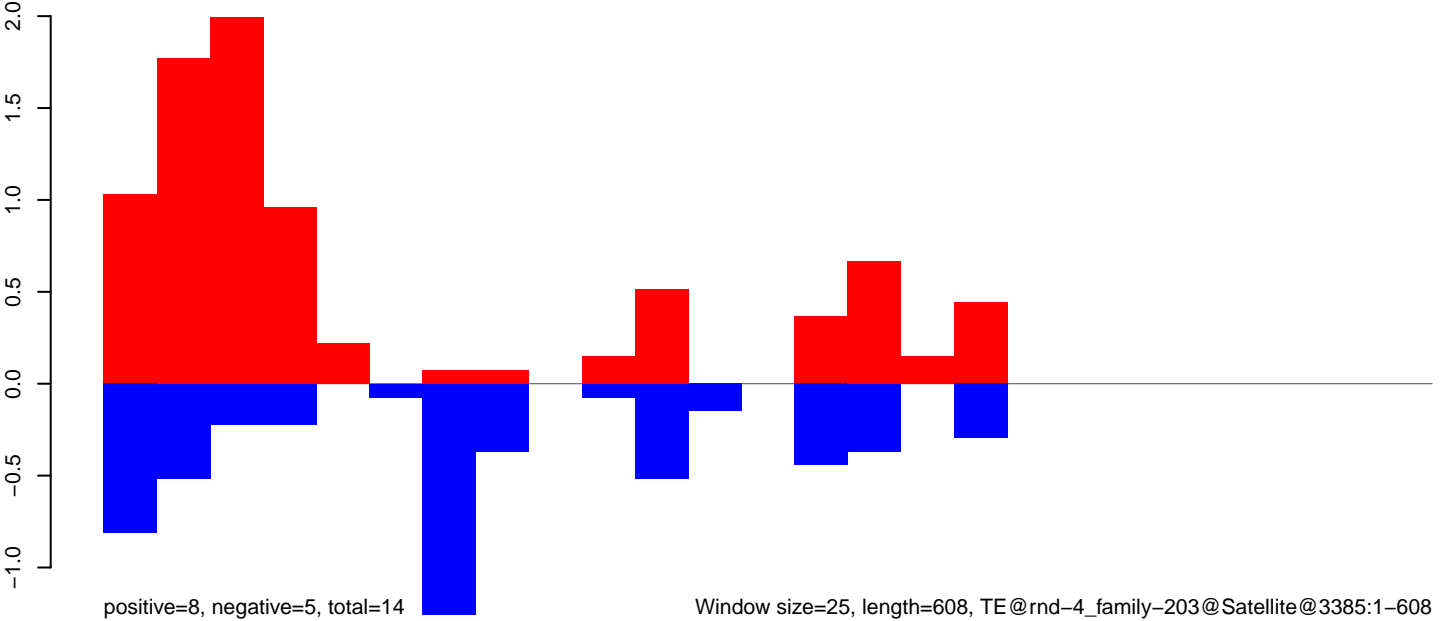
CuQuin_Testes_NL.18_23.rep



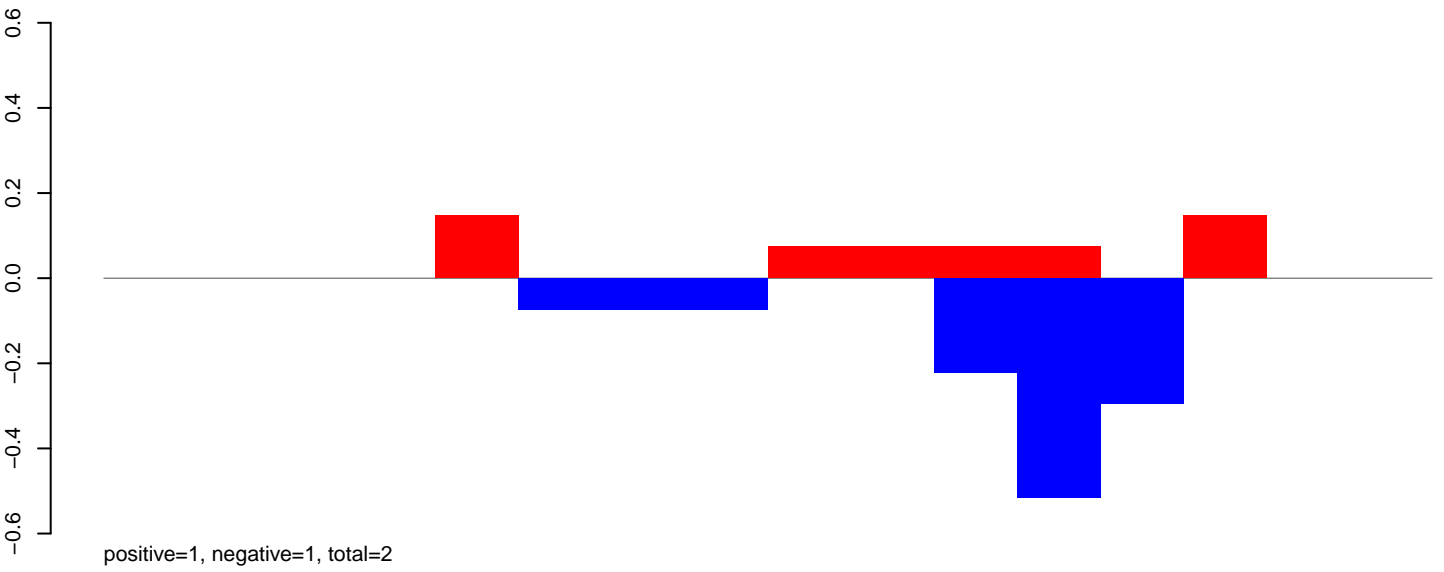
CuQuin_Testes_NL.24_35.rep



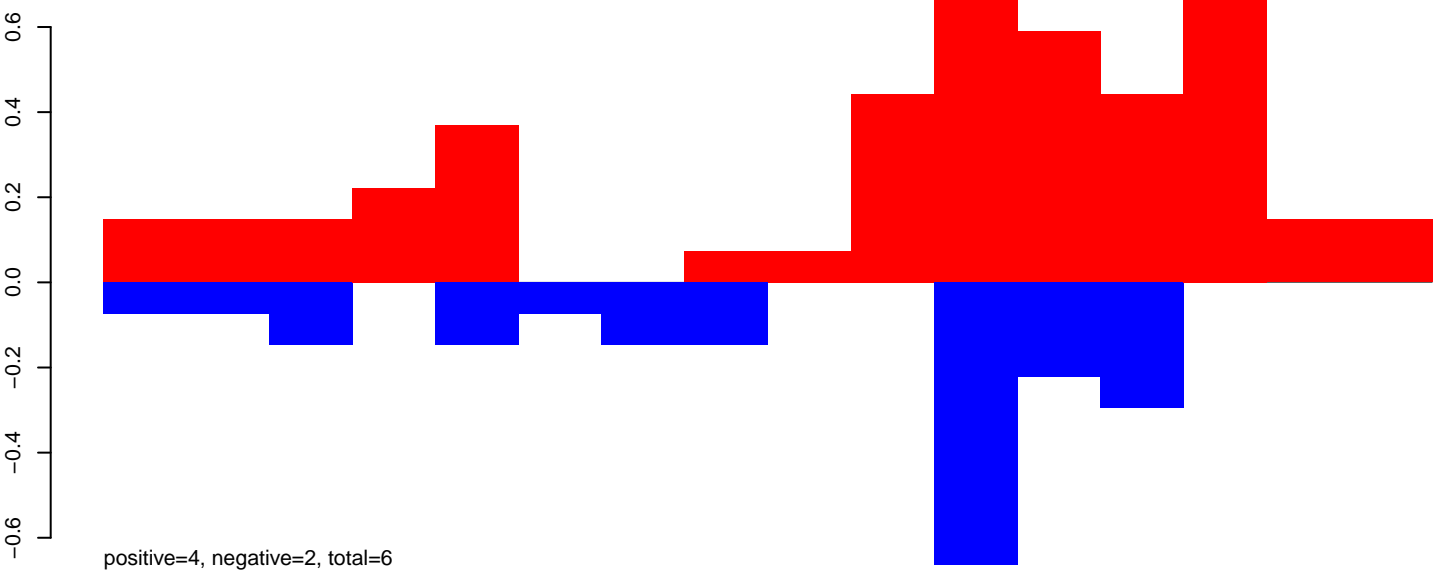
CuQuin_Testes_NL.rep



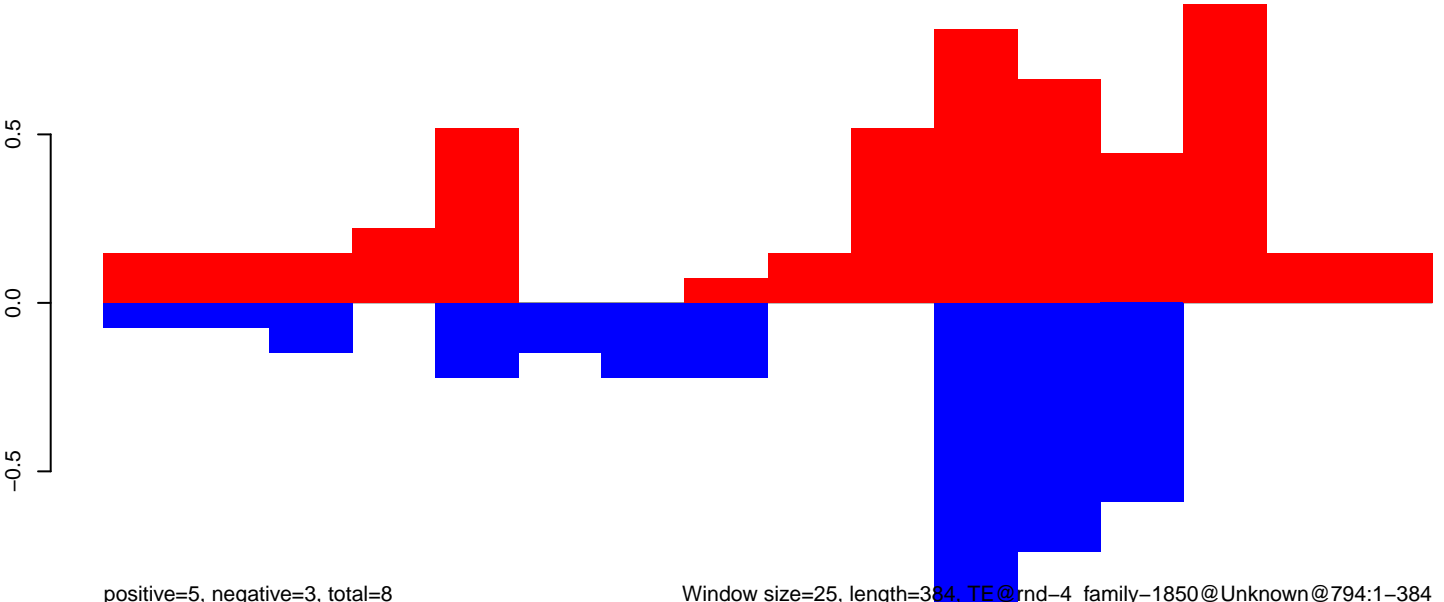
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



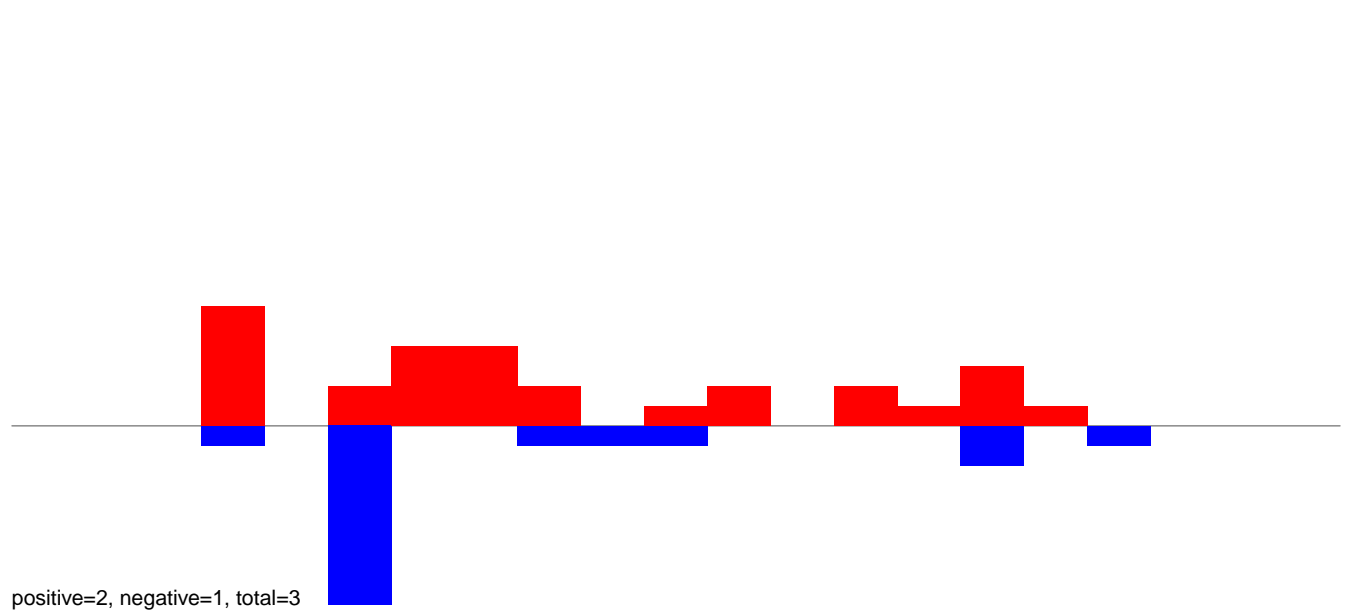
CuQuin_Testes_NL.rep



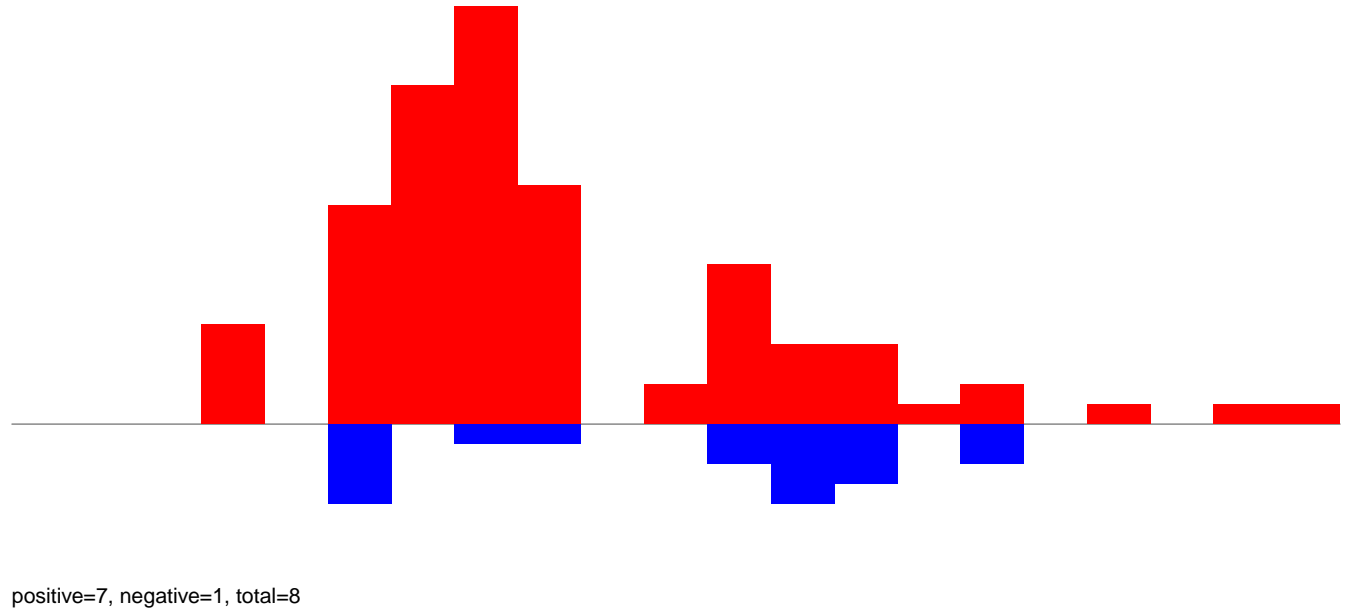
Window size=25, length=384, TE@rnd-4_family-1850@Unknown@794:1-384

100 200 300 400

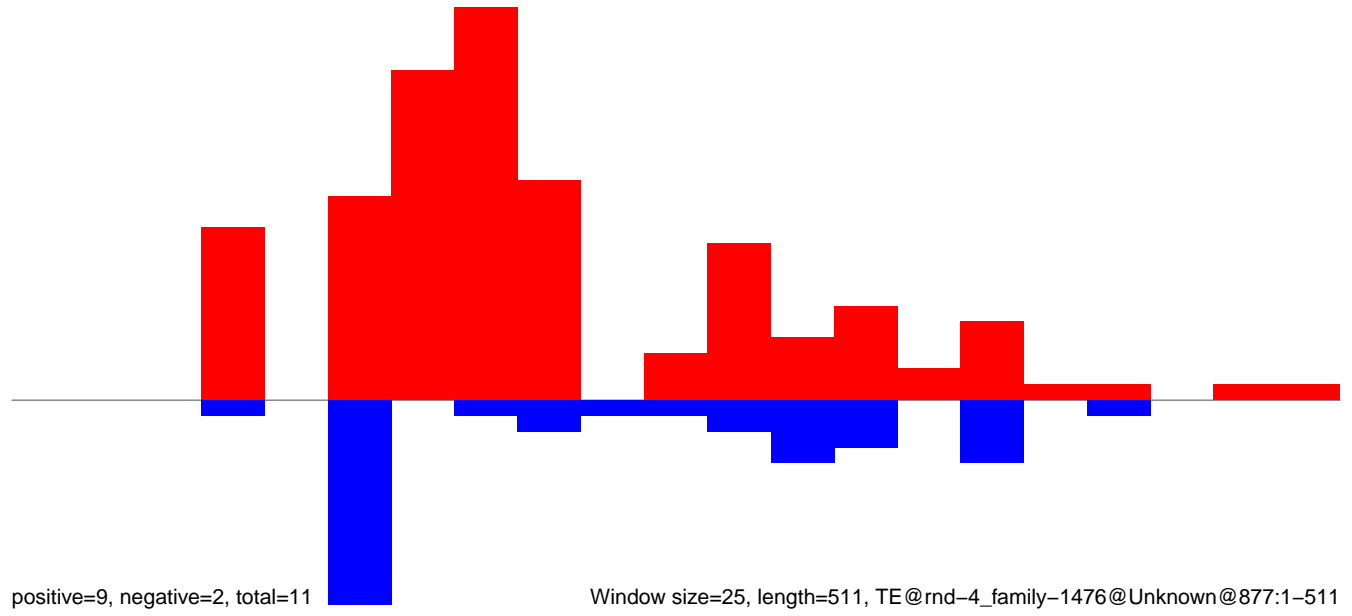
CuQuin_Testes_NL.18_23.rep



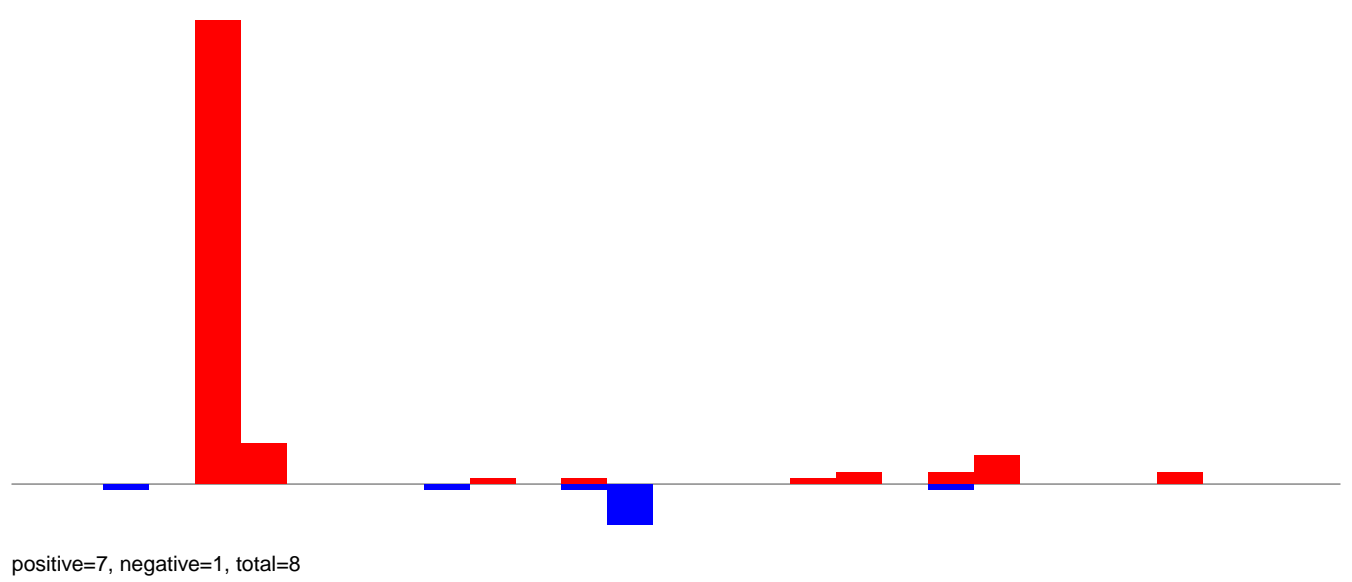
CuQuin_Testes_NL.24_35.rep



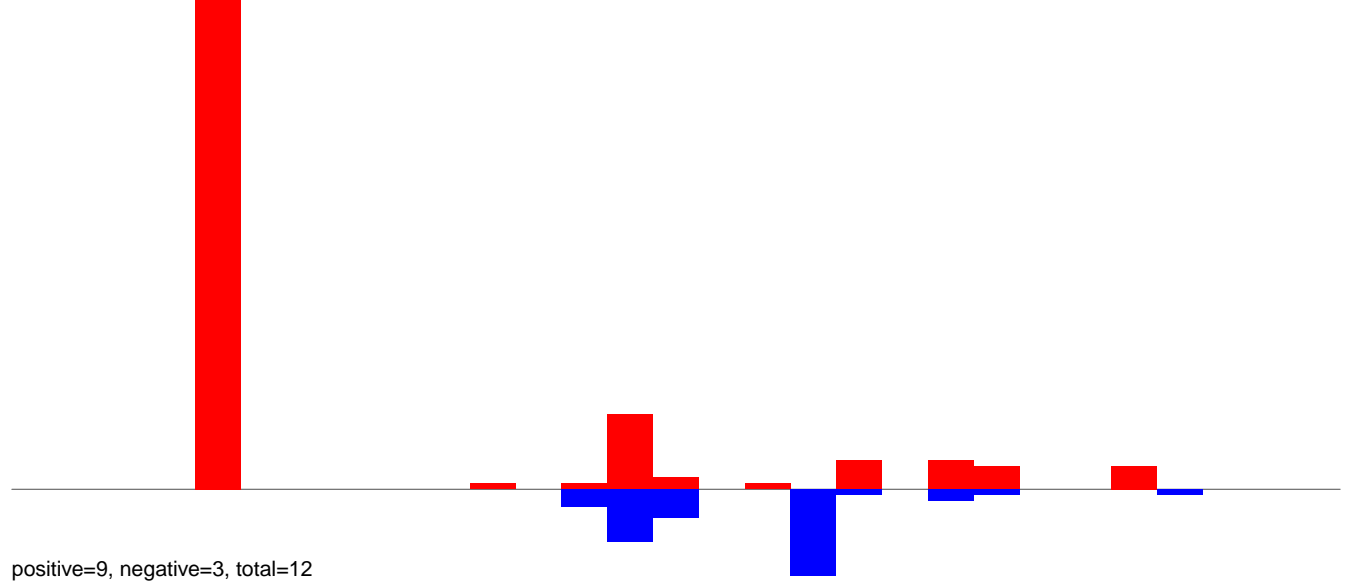
CuQuin_Testes_NL.rep



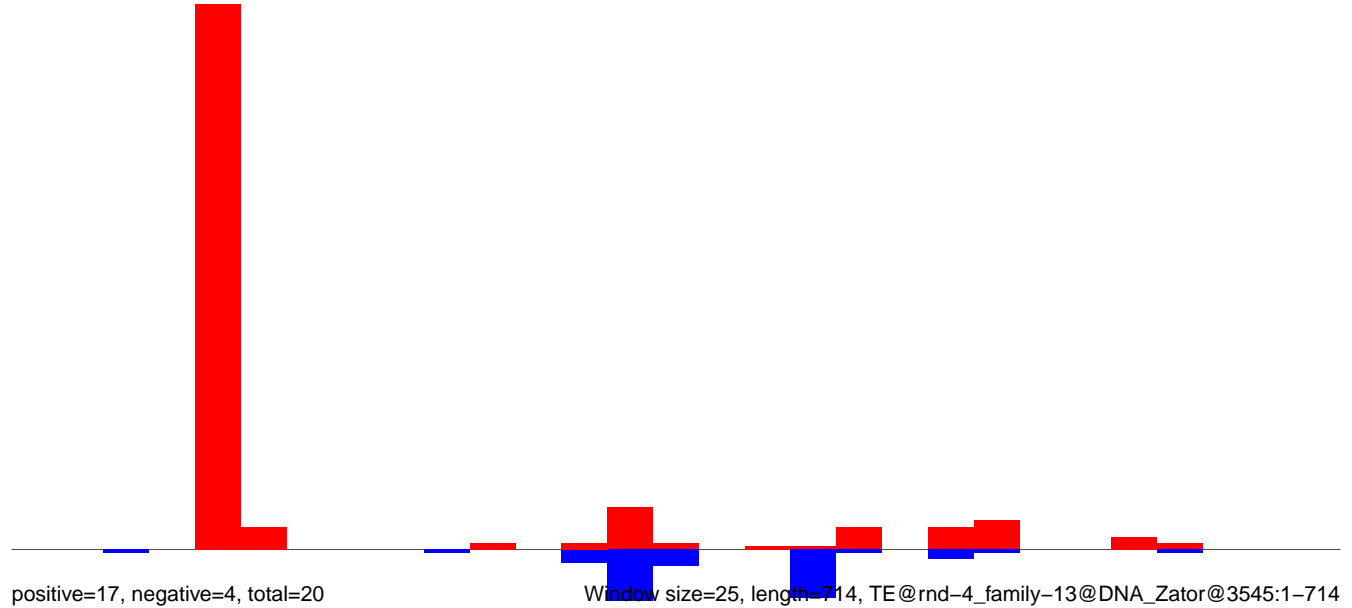
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

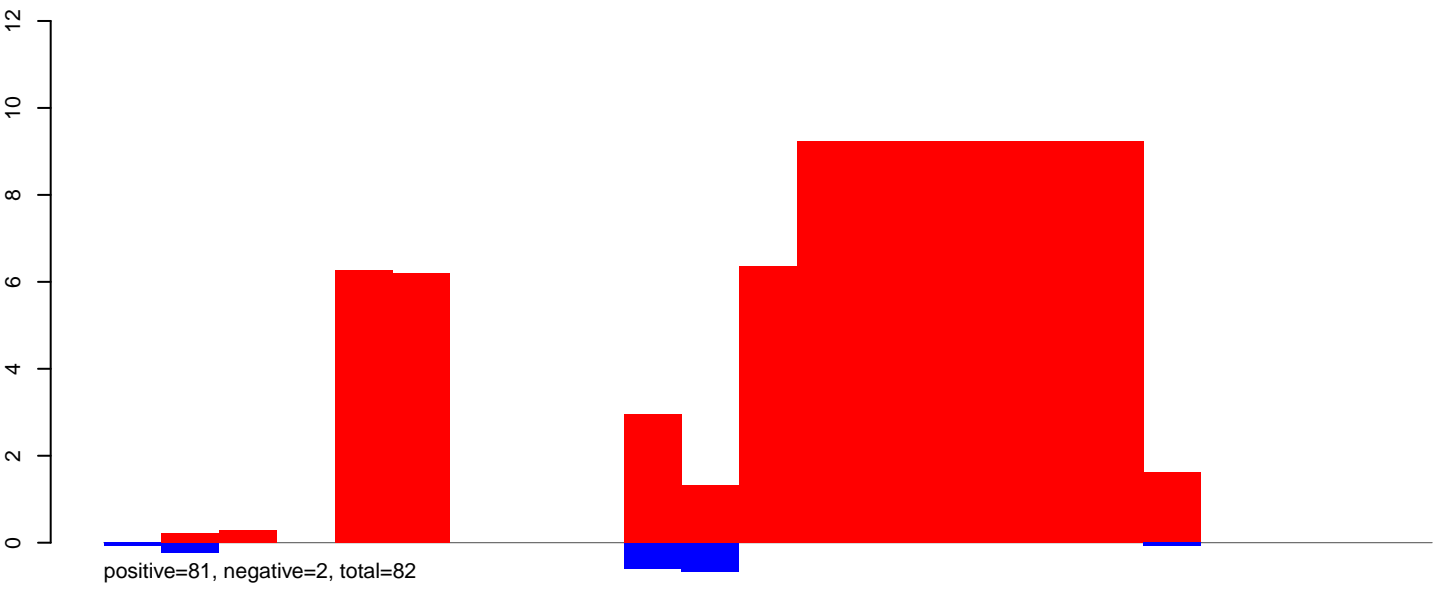


CuQuin_Testes_NL.rep

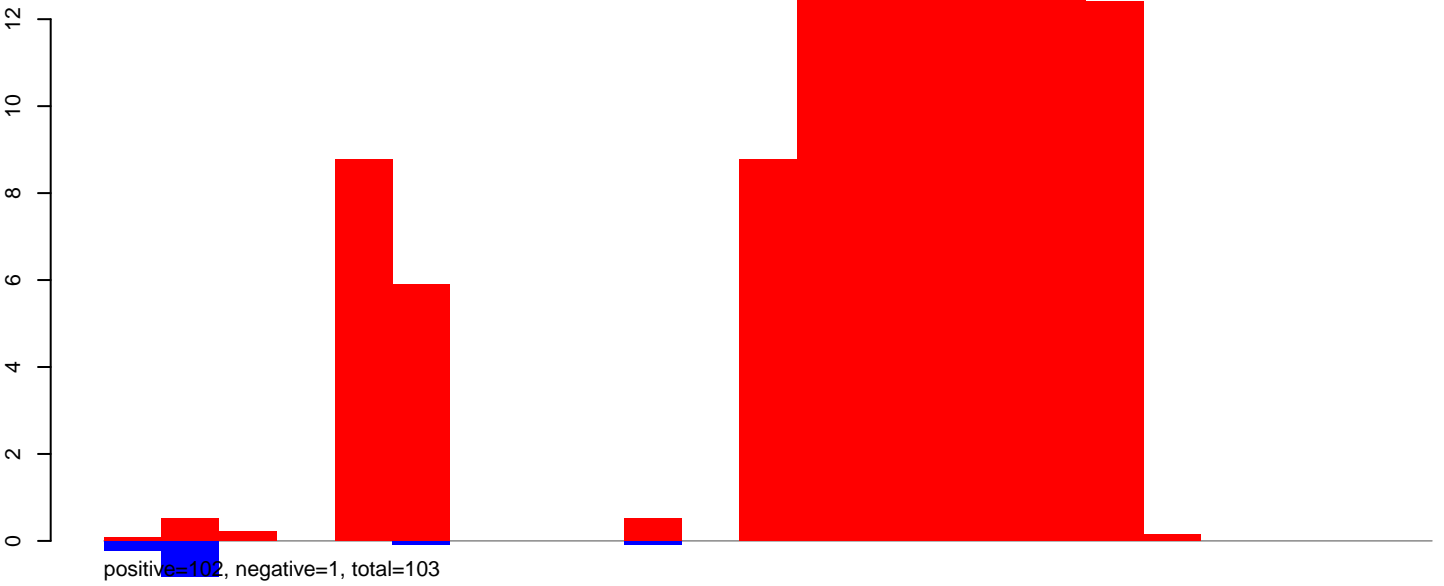


Window size=25, length=714, TE@rnd-4_family-13@DNA_Zator@3545:1-714

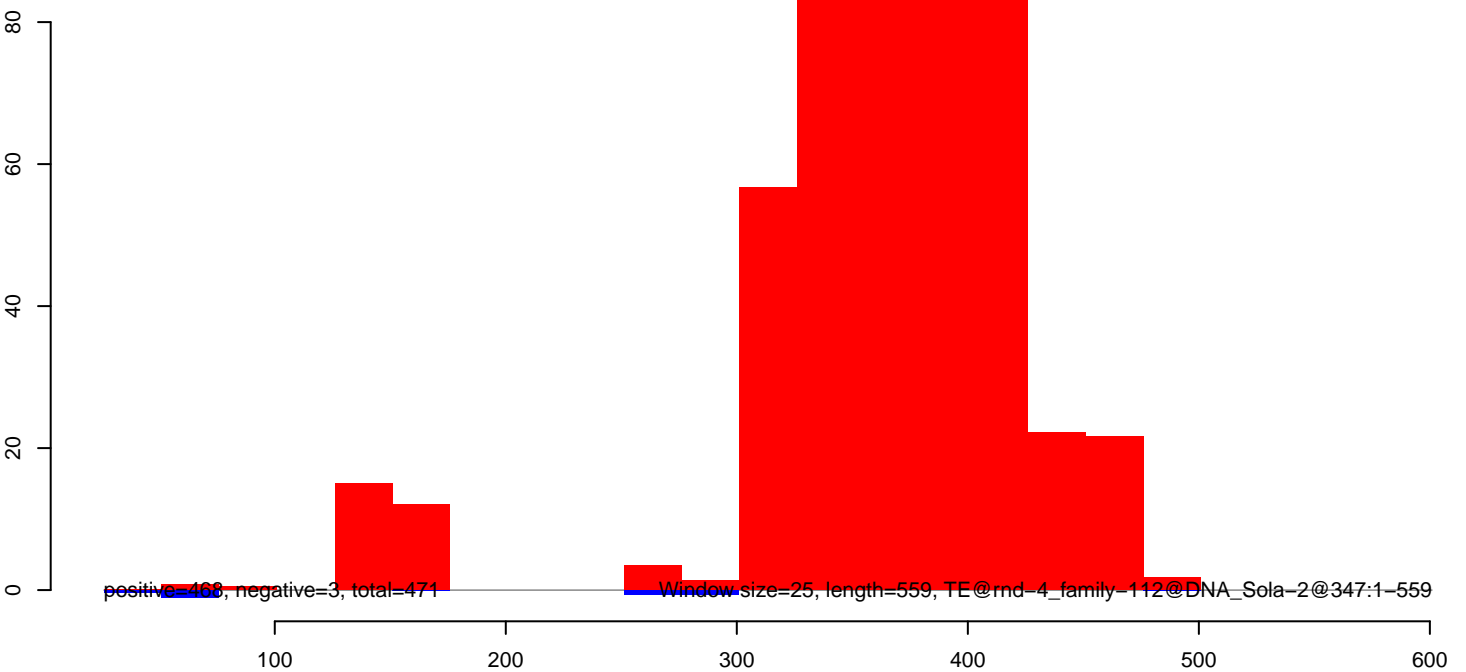
CuQuin_Testes_NL.18_23.rep



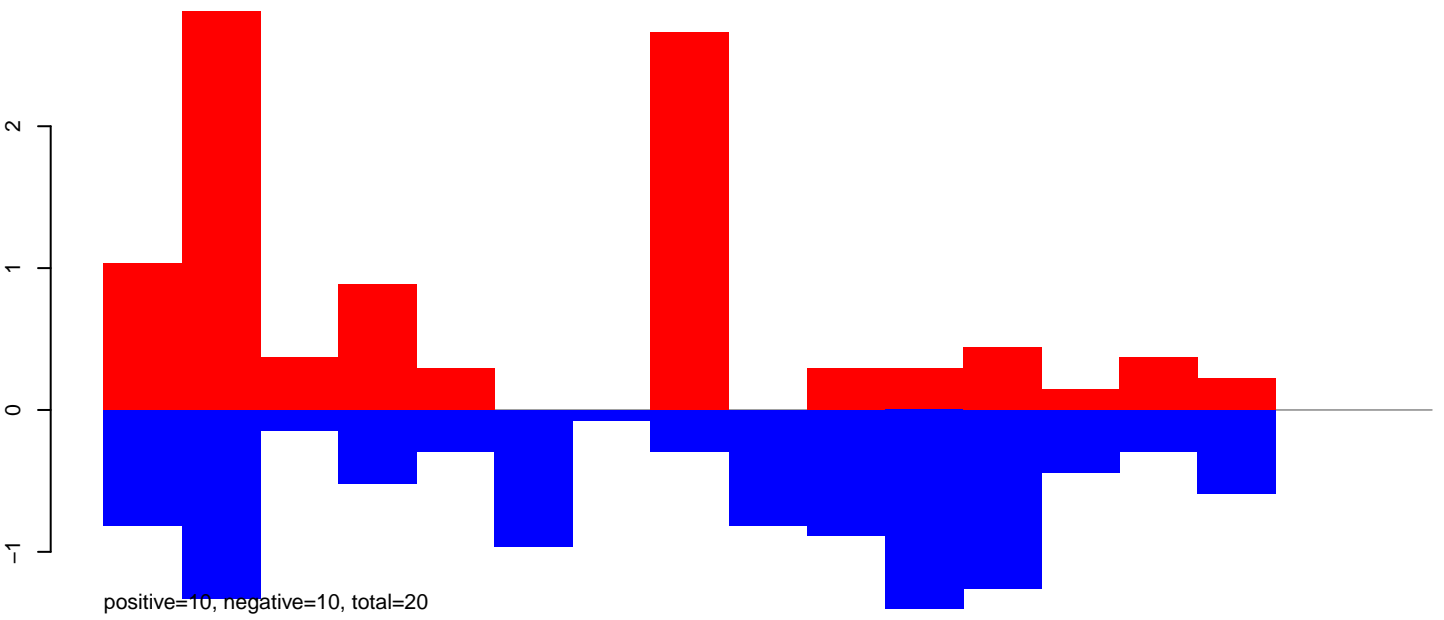
CuQuin_Testes_NL.24_35.rep



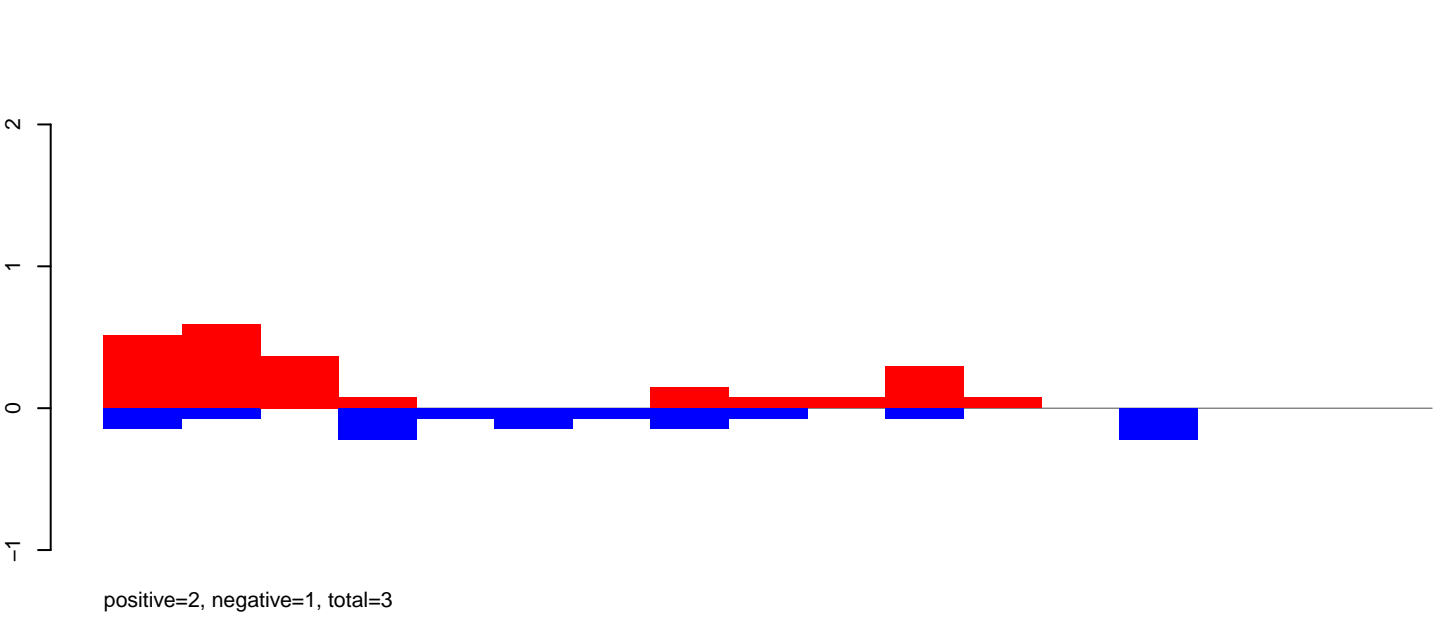
CuQuin_Testes_NL.rep



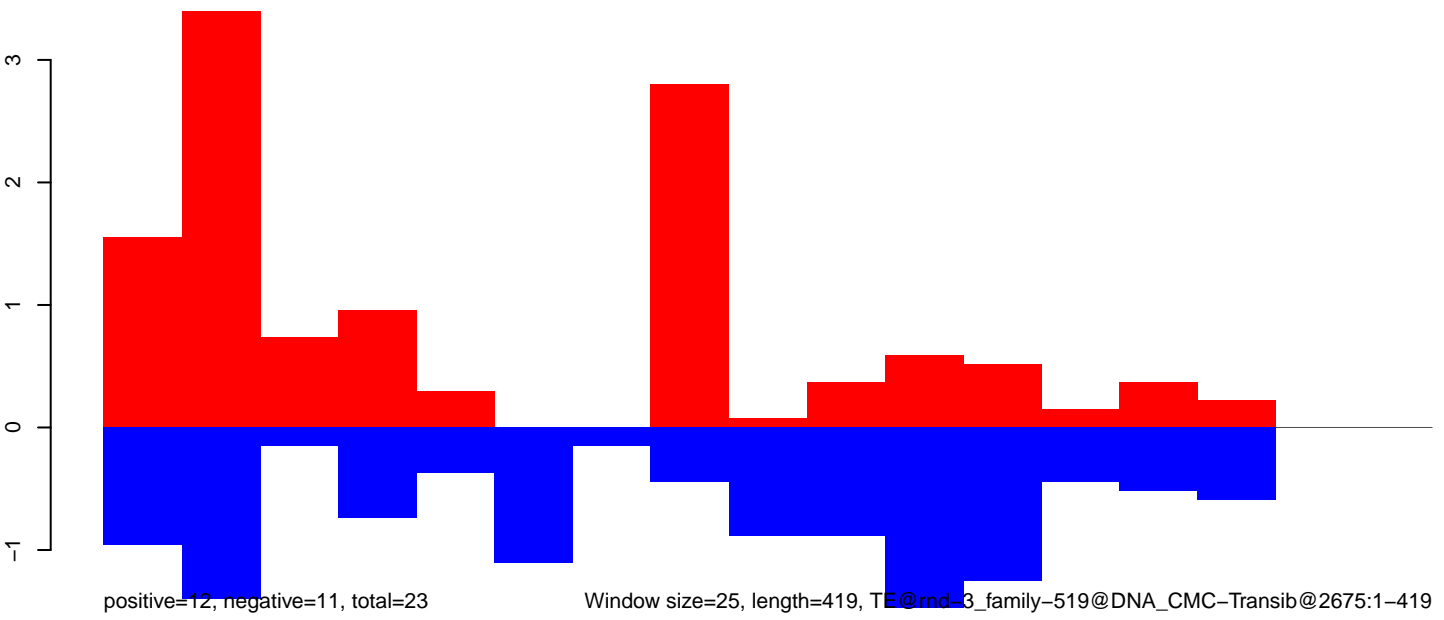
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



Window size=25, length=419, TE@rnd_3_family-519@DNA_CMC-Transib@2675:1-419

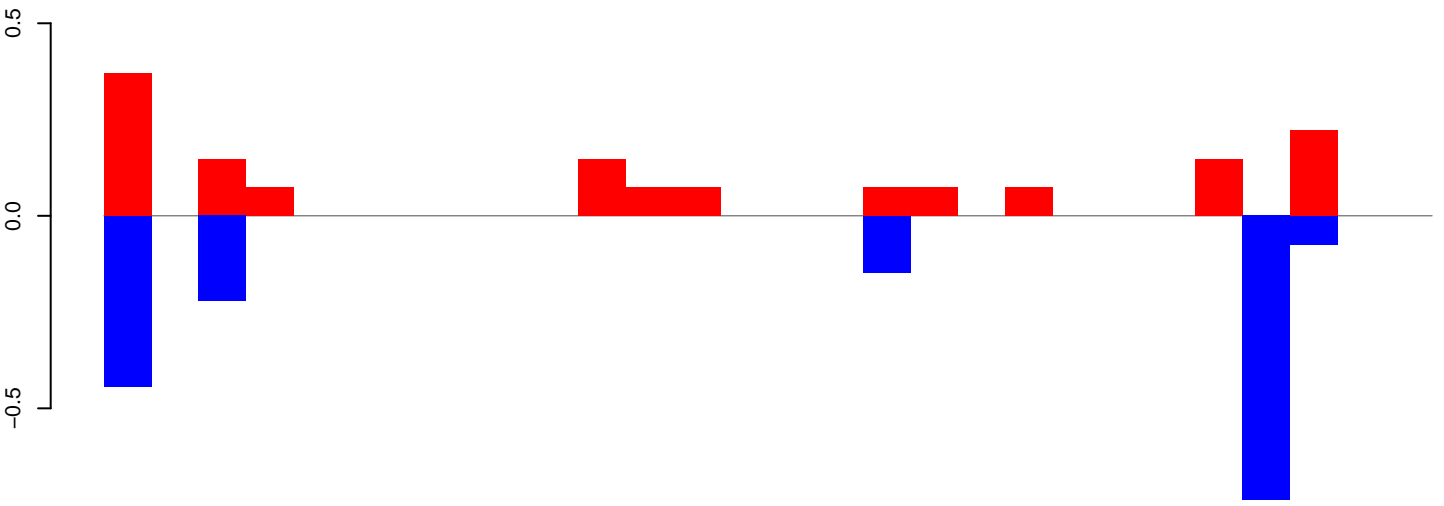
100

200

300

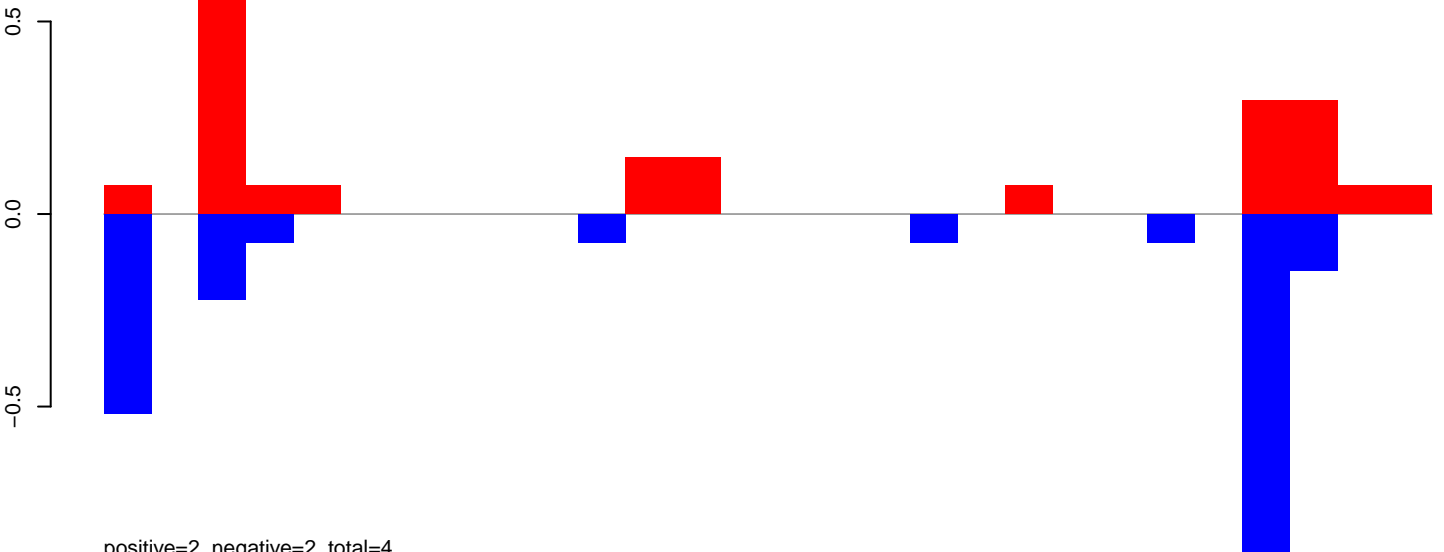
400

CuQuin_Testes_NL.18_23.rep



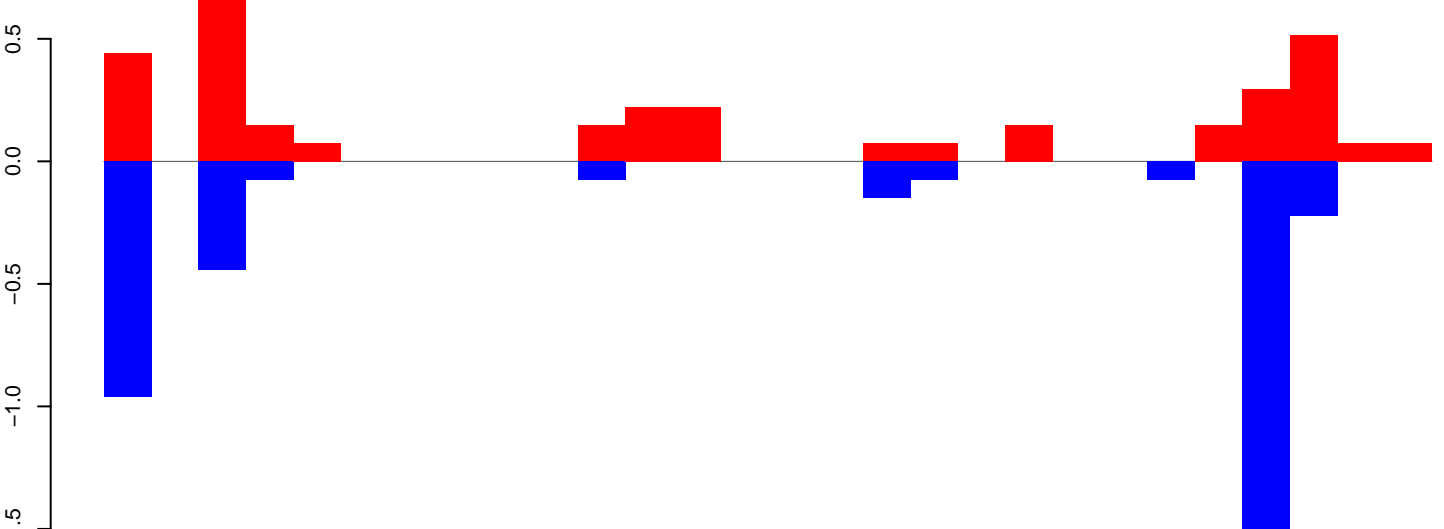
positive=1, negative=2, total=3

CuQuin_Testes_NL.24_35.rep



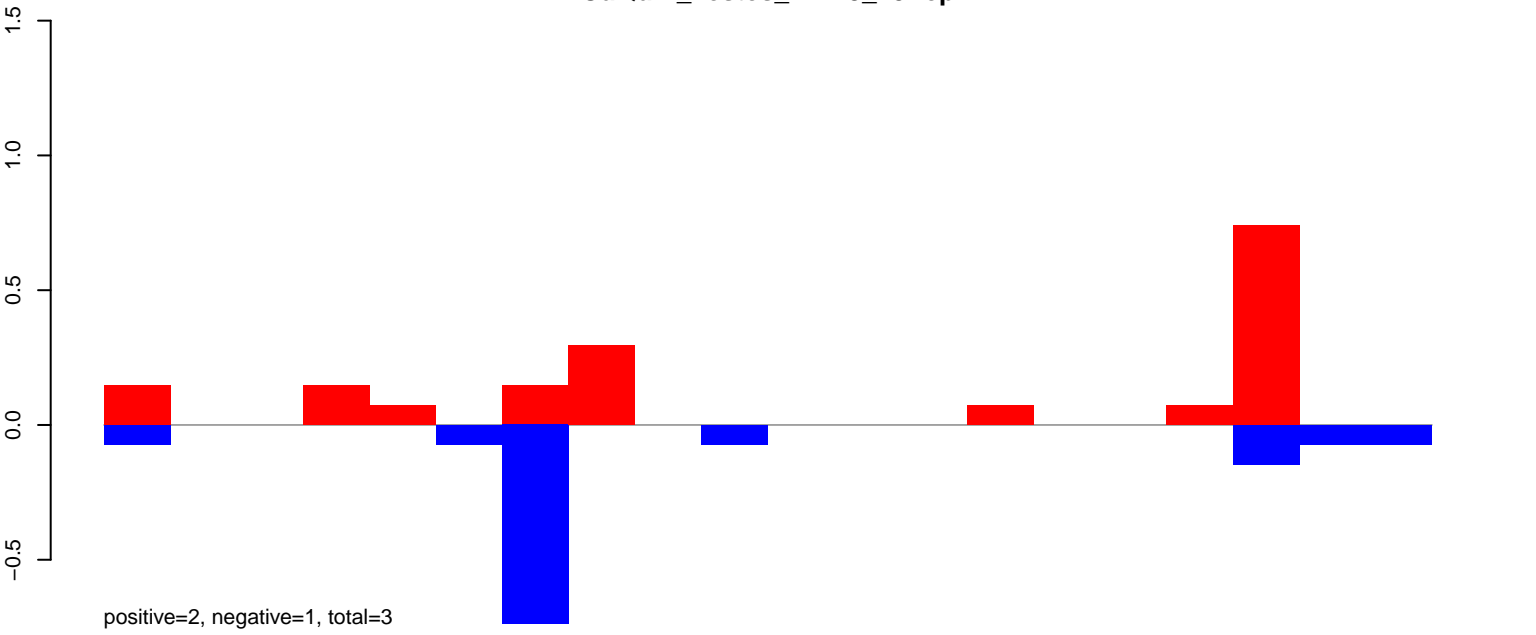
positive=2, negative=2, total=4

CuQuin_Testes_NL.rep

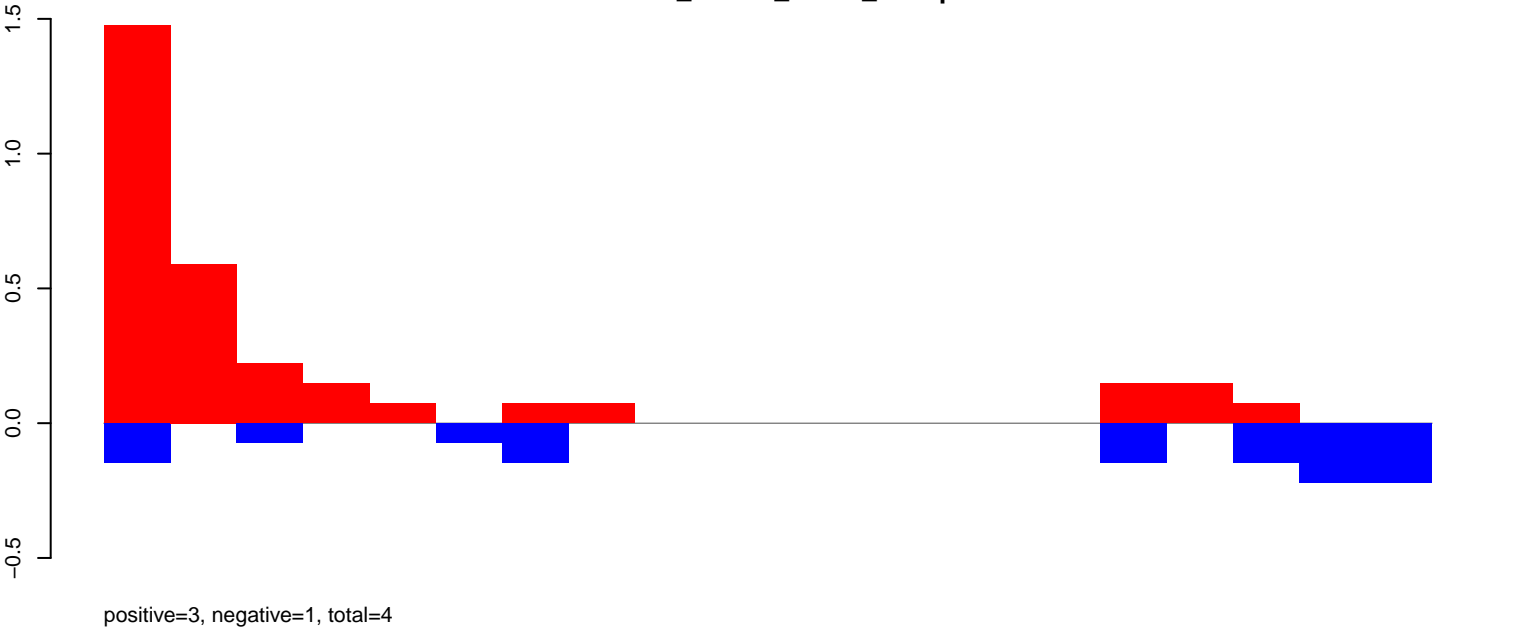


positive=3, negative=4, total=7

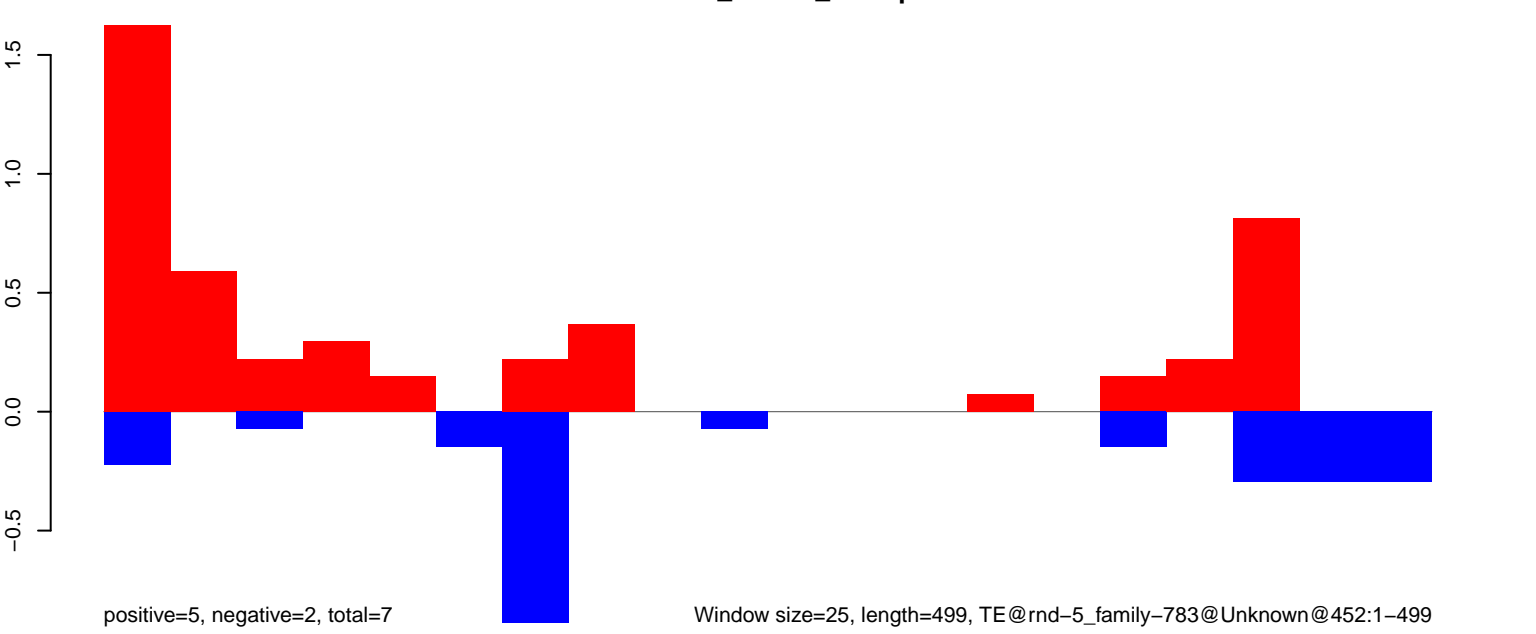
CuQuin_Testes_NL.18_23.rep



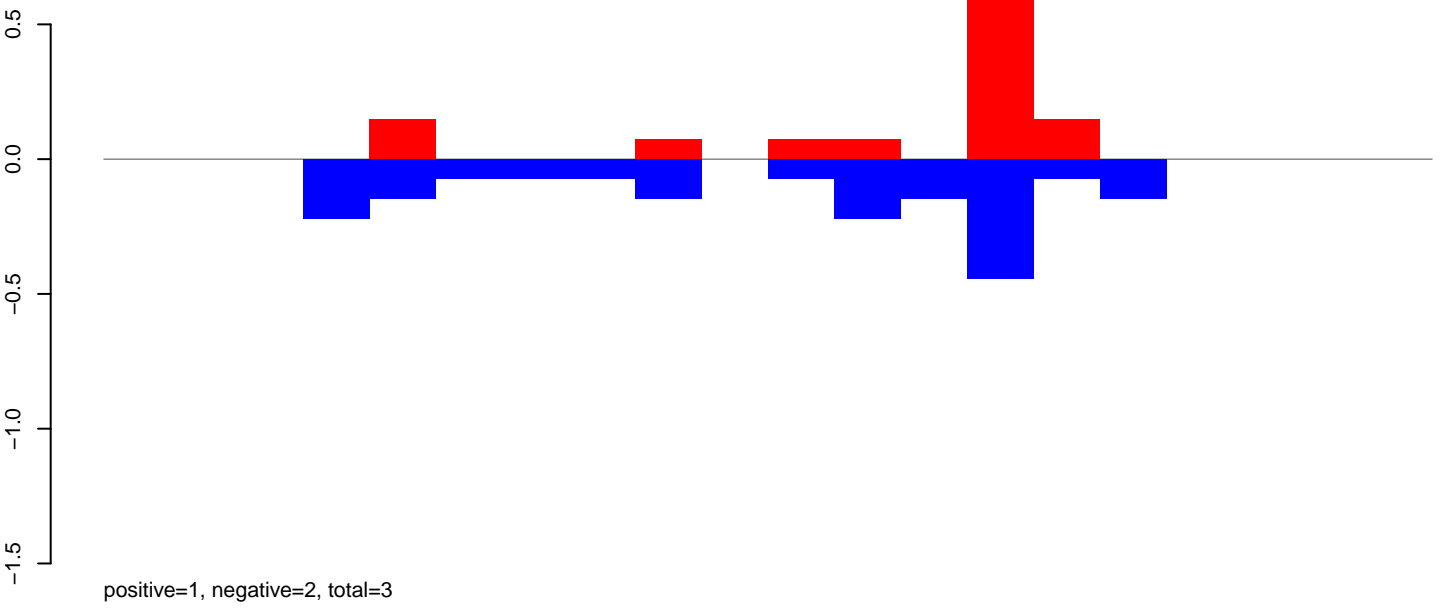
CuQuin_Testes_NL.24_35.rep



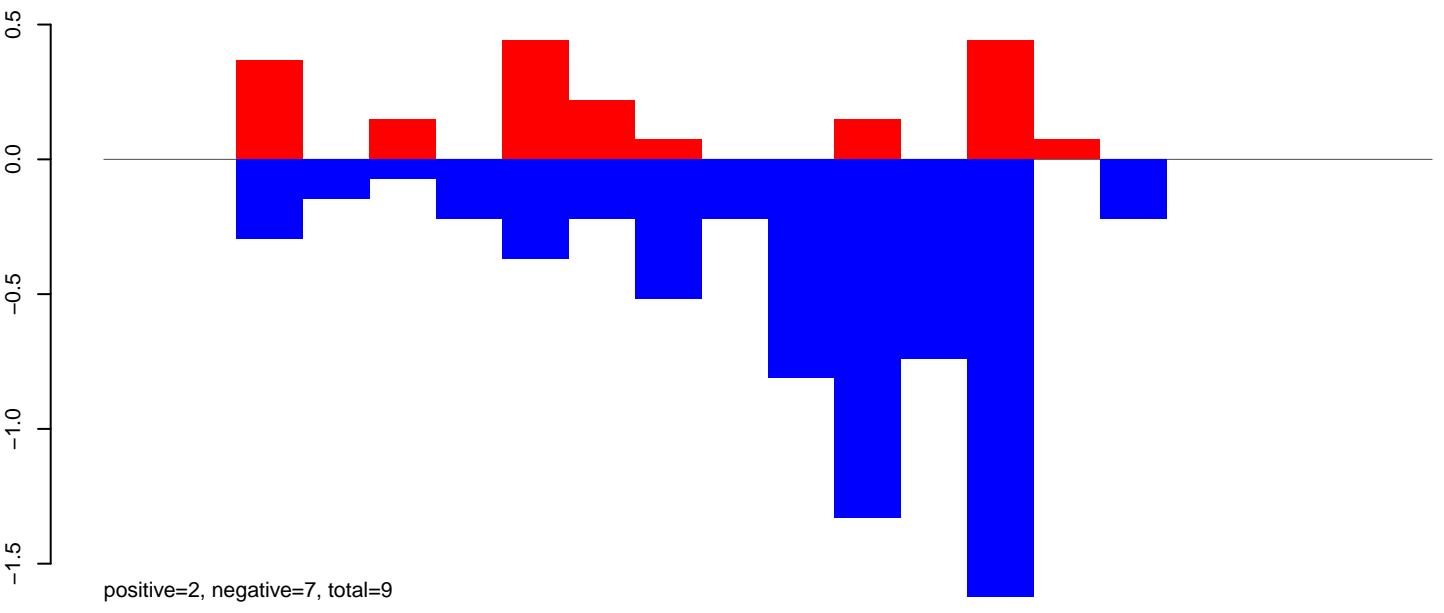
CuQuin_Testes_NL.rep



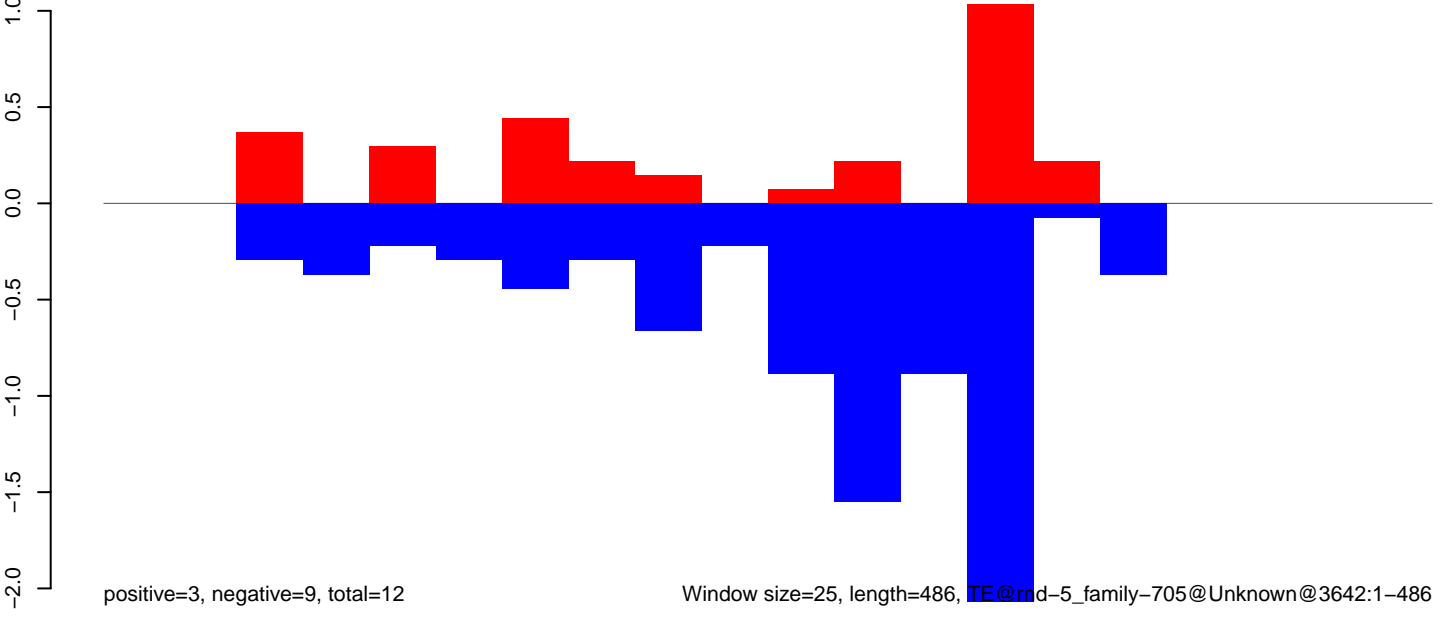
CuQuin_Testes_NL.18_23.rep



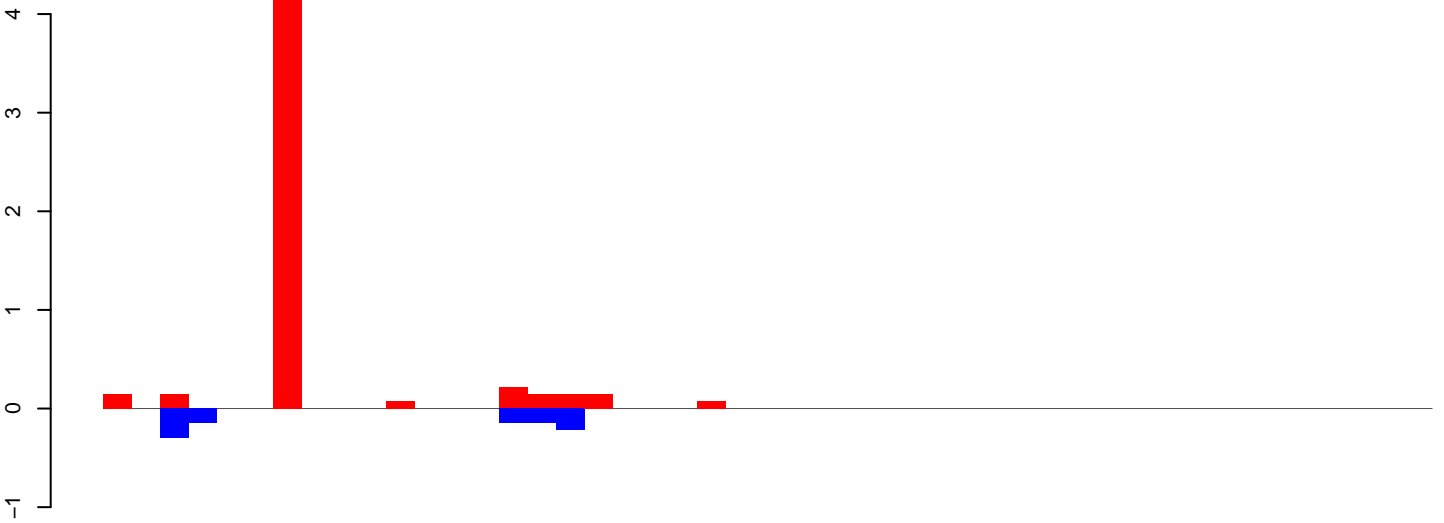
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

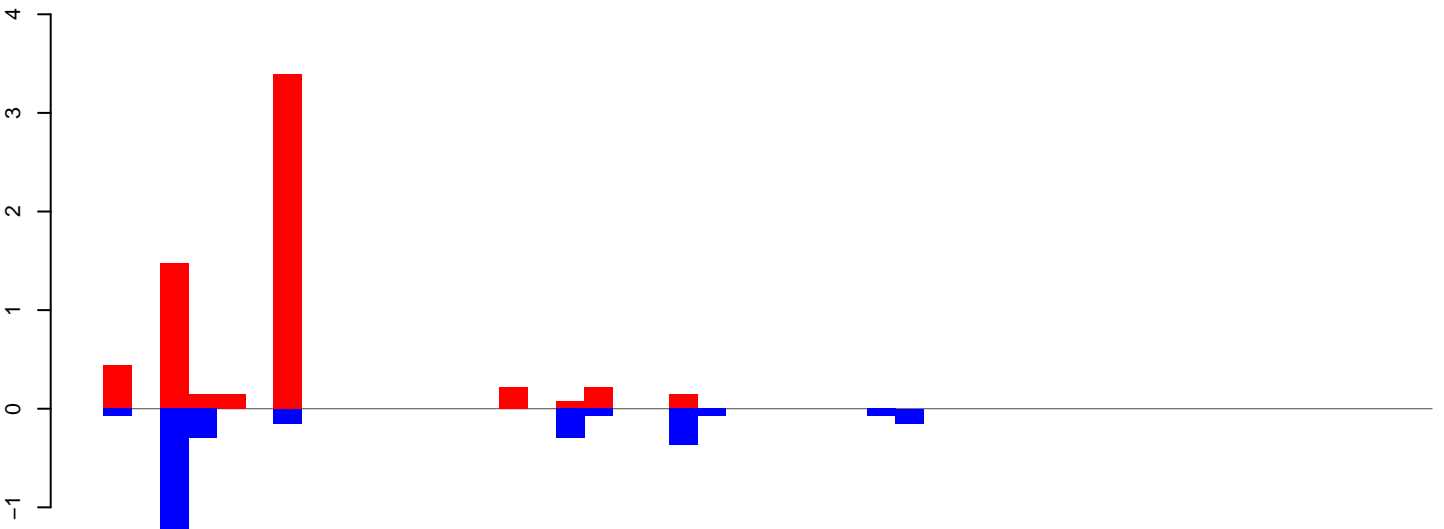


CuQuin_Testes_NL.18_23.rep



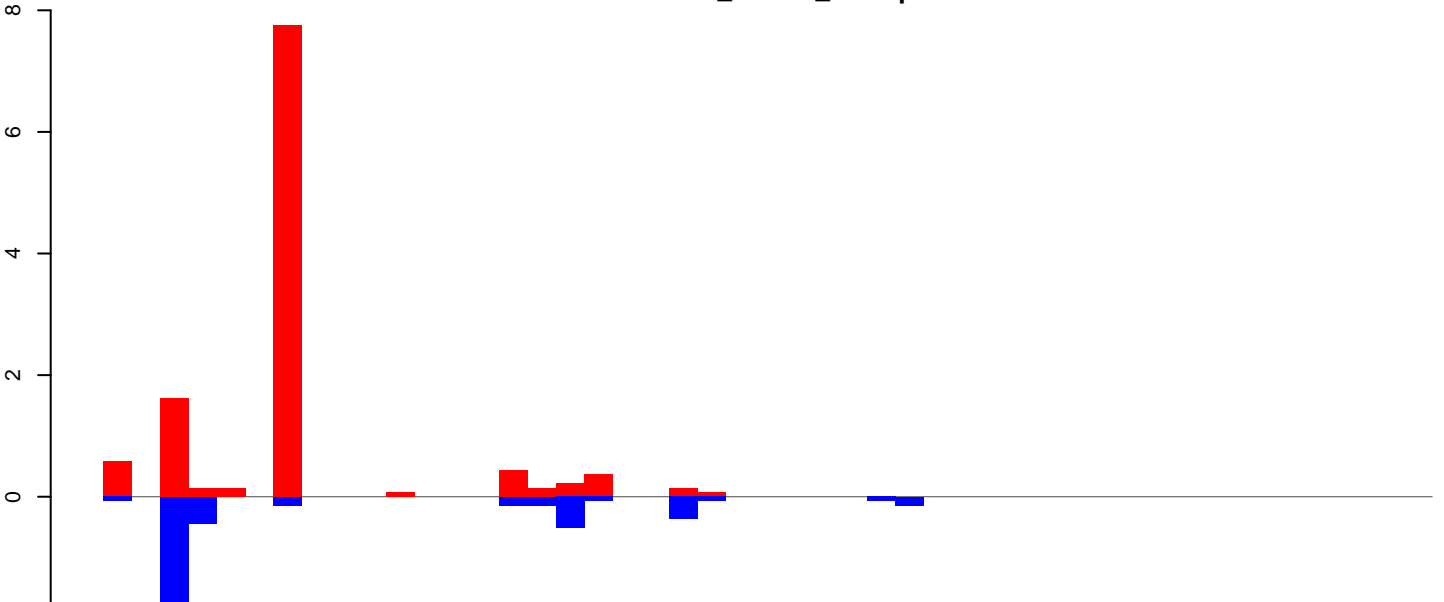
positive=5, negative=1, total=6

CuQuin_Testes_NL.24_35.rep



positive=6, negative=3, total=10

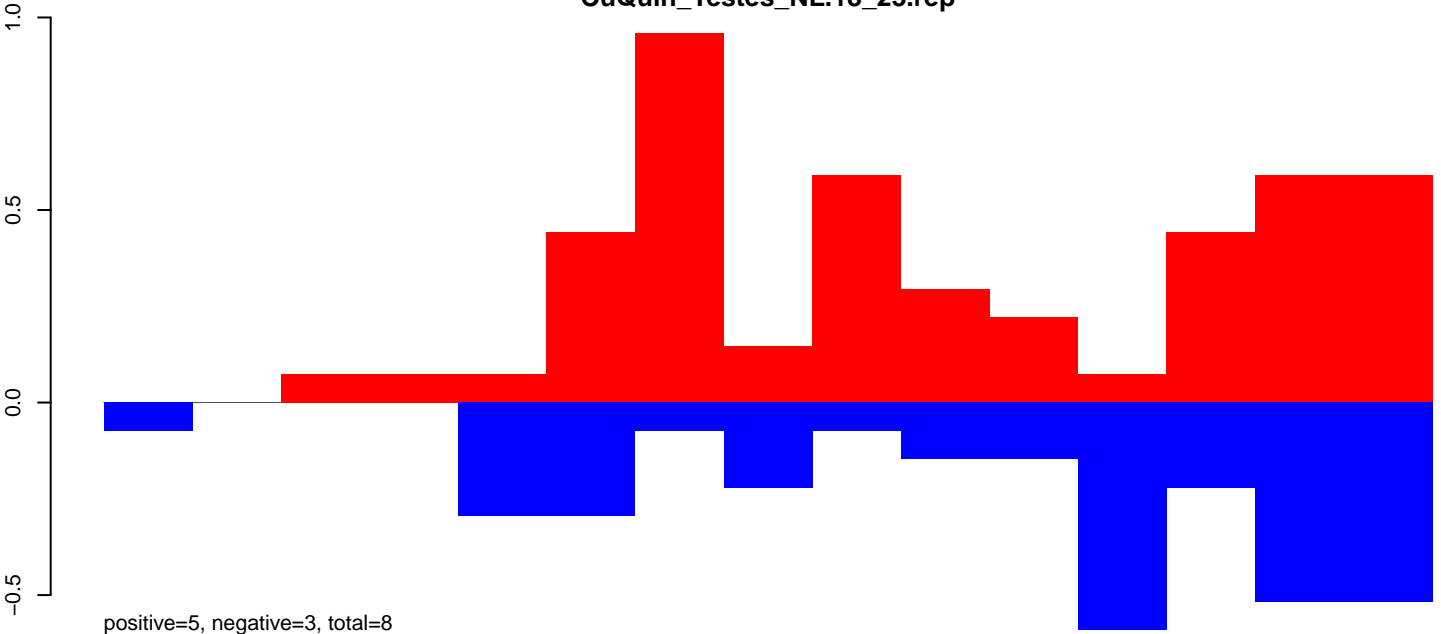
CuQuin_Testes_NL.rep



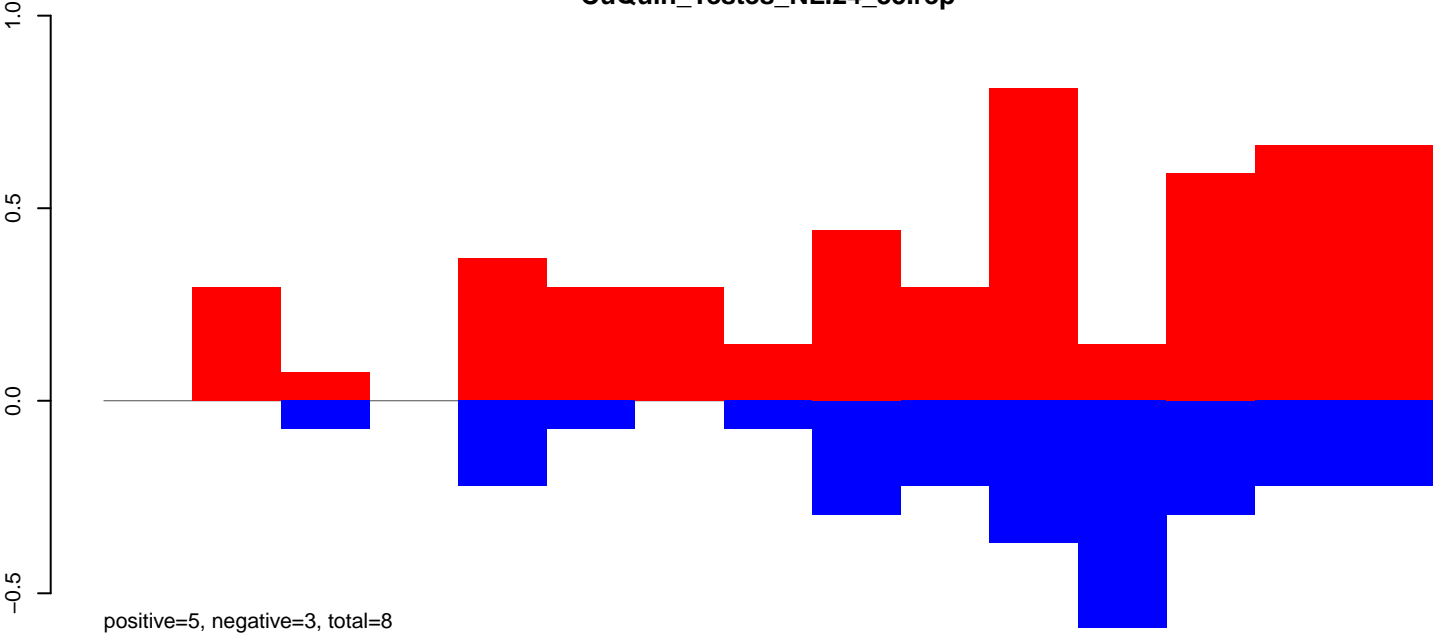
positive=12, negative=4, total=16

Window size=25, length=1151, TE@rnd-5_family-69@LINE_I-Jockey@510:1-1151

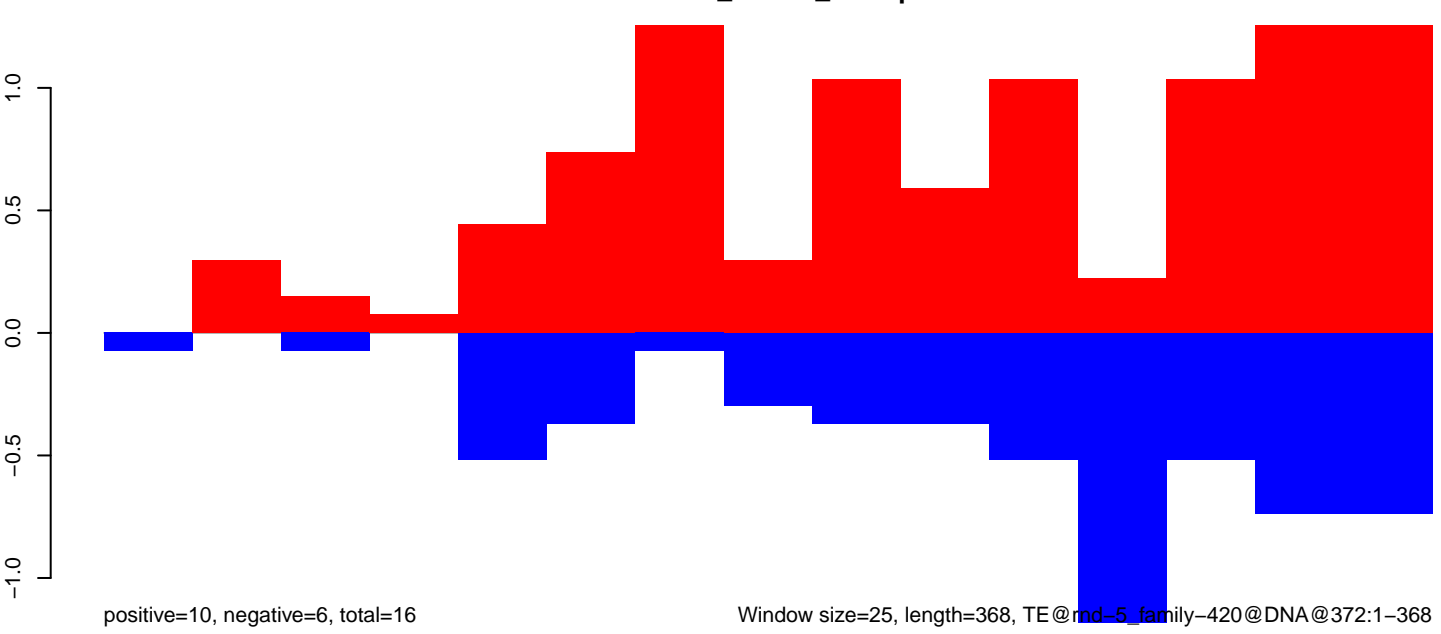
CuQuin_Testes_NL.18_23.rep



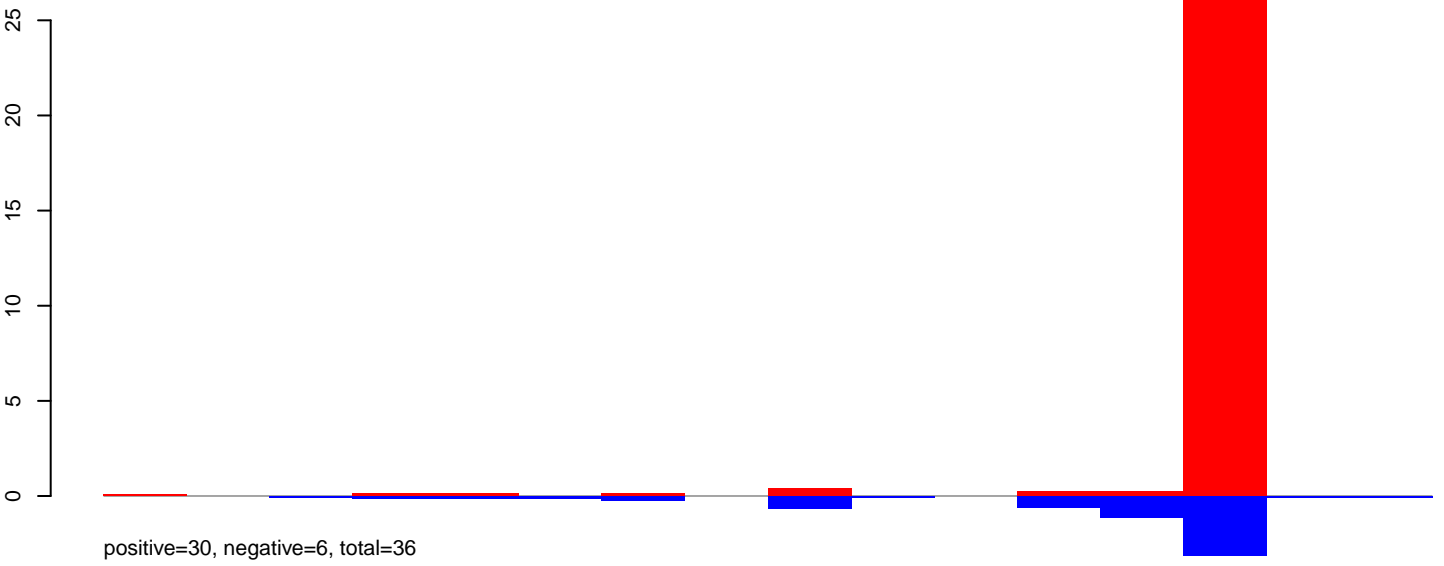
CuQuin_Testes_NL.24_35.rep



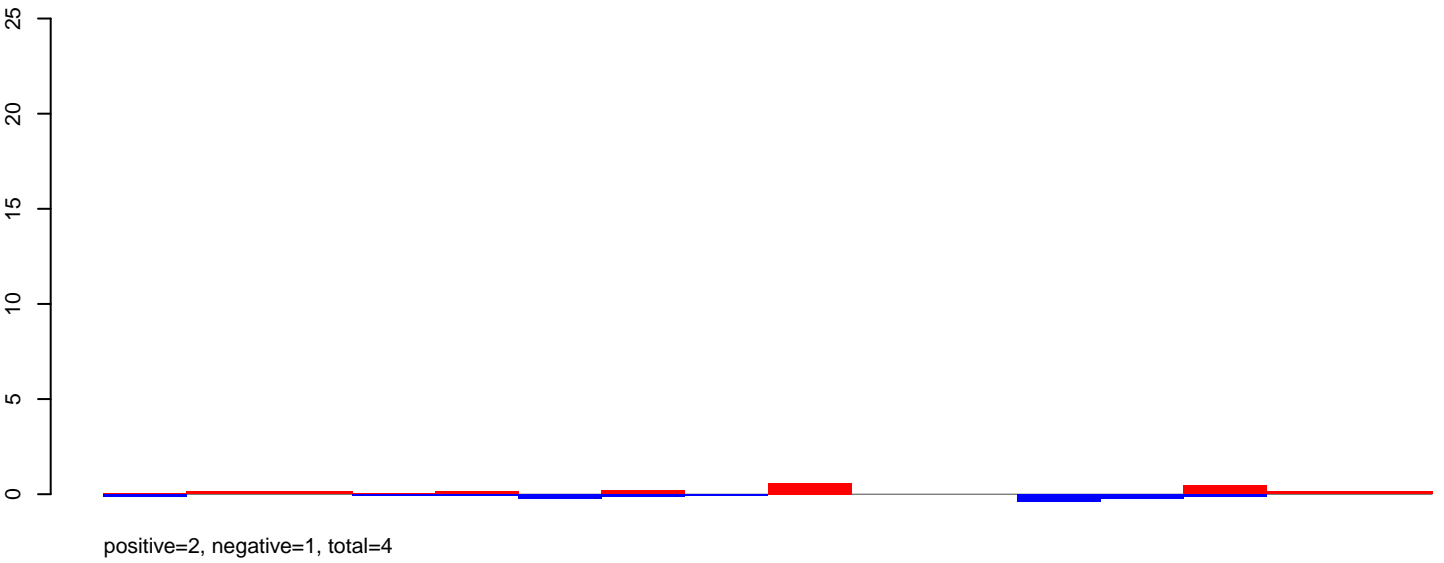
CuQuin_Testes_NL.rep



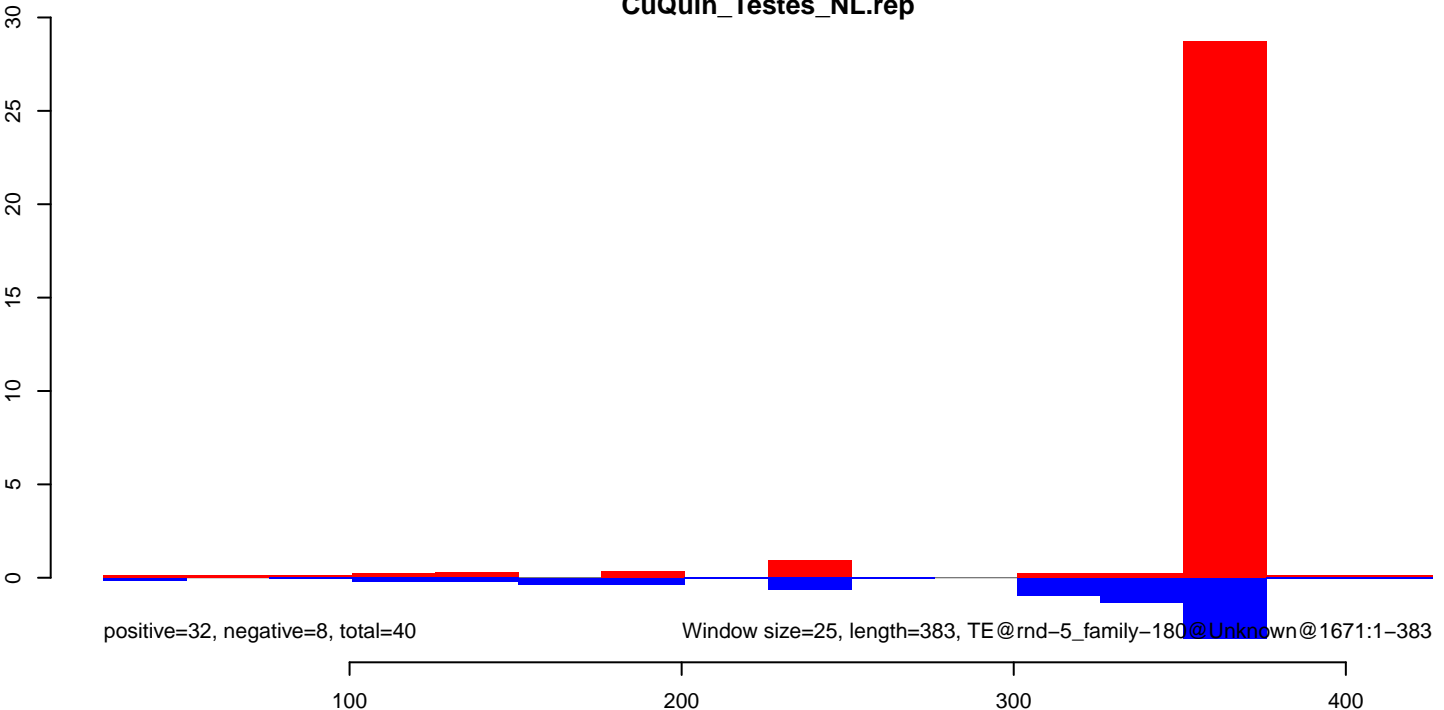
CuQuin_Testes_NL.18_23.rep



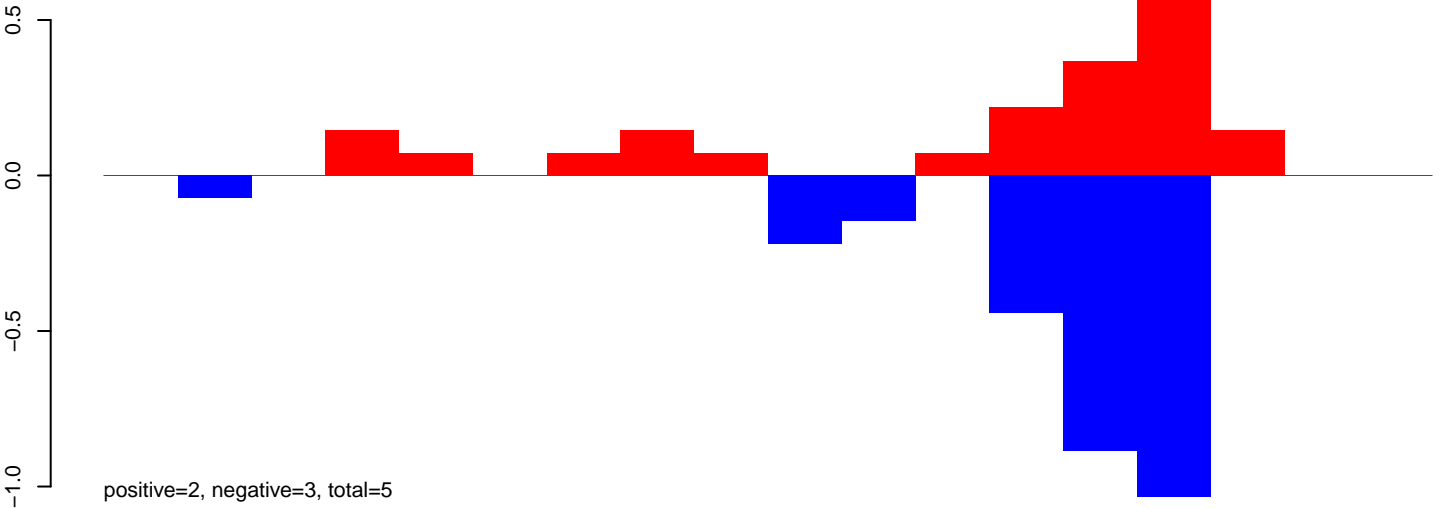
CuQuin_Testes_NL.24_35.rep



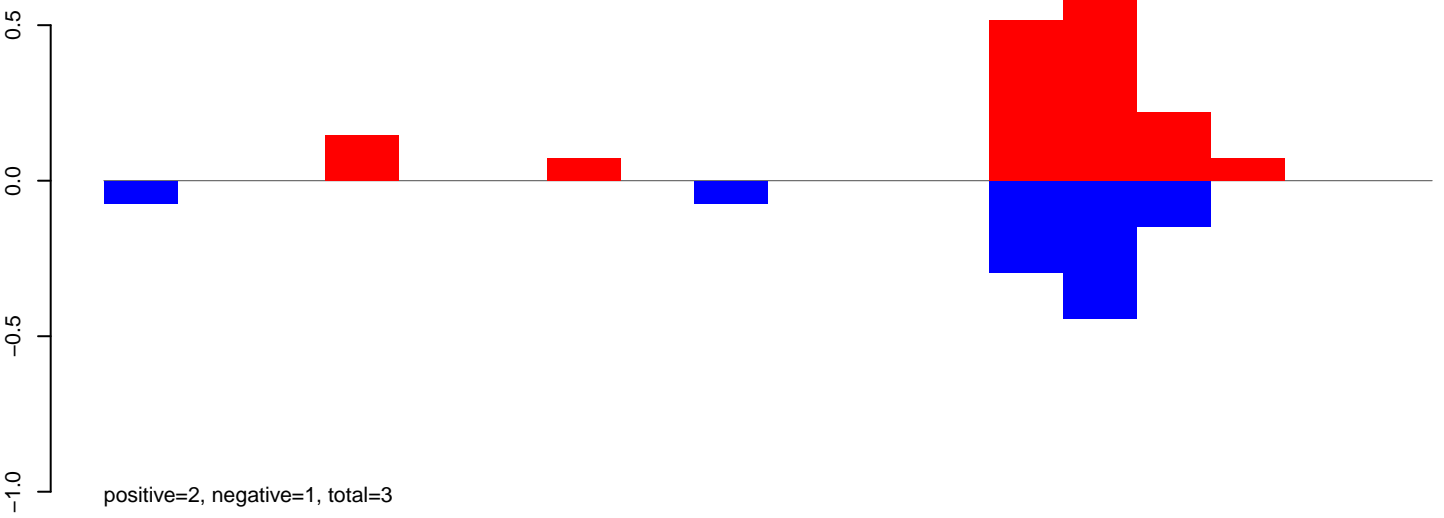
CuQuin_Testes_NL.rep



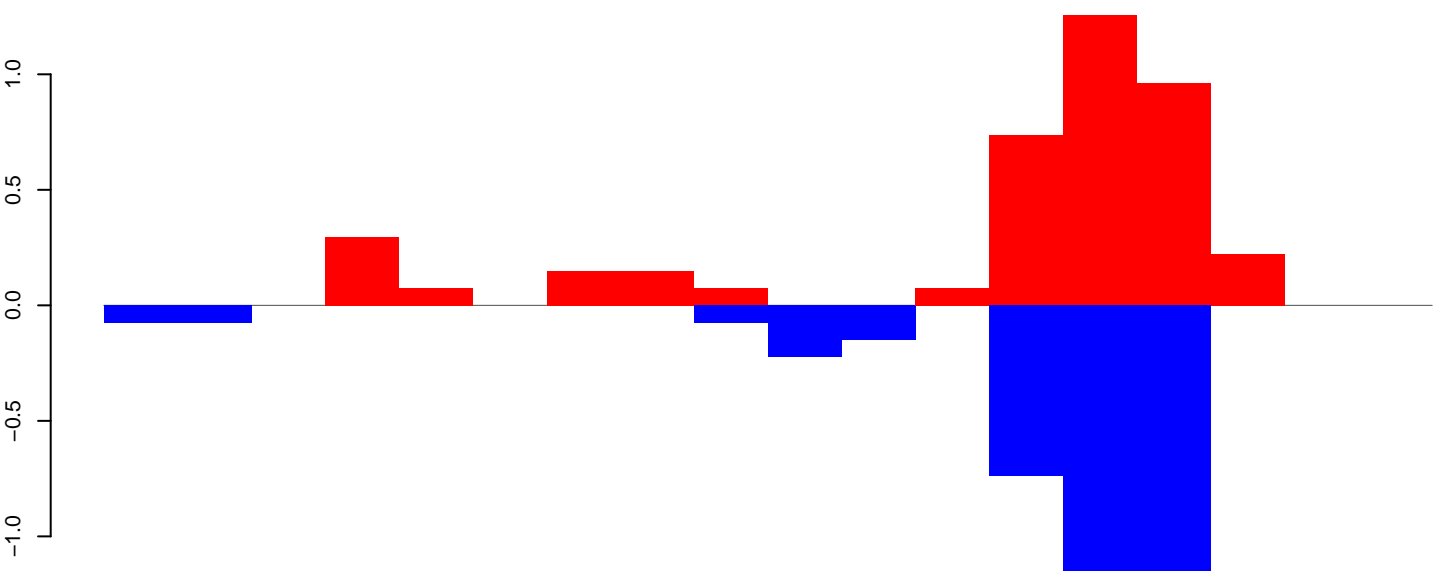
CuQuin_Testes_NL.18_23.rep



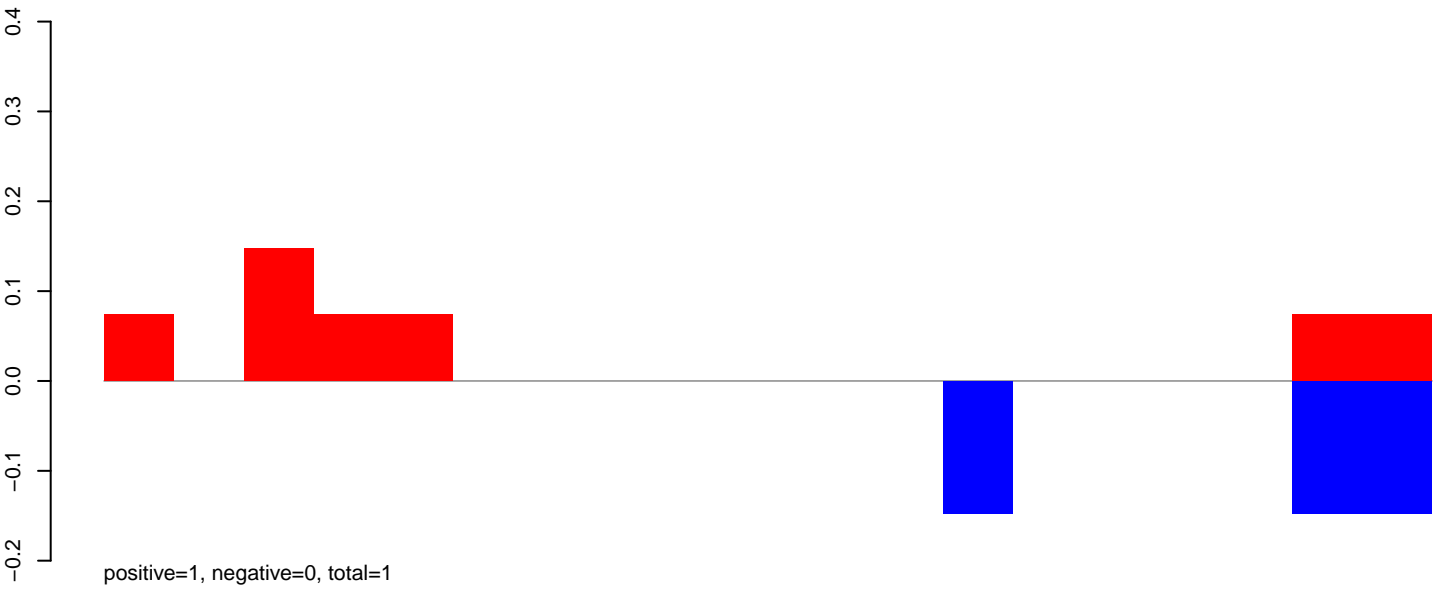
CuQuin_Testes_NL.24_35.rep



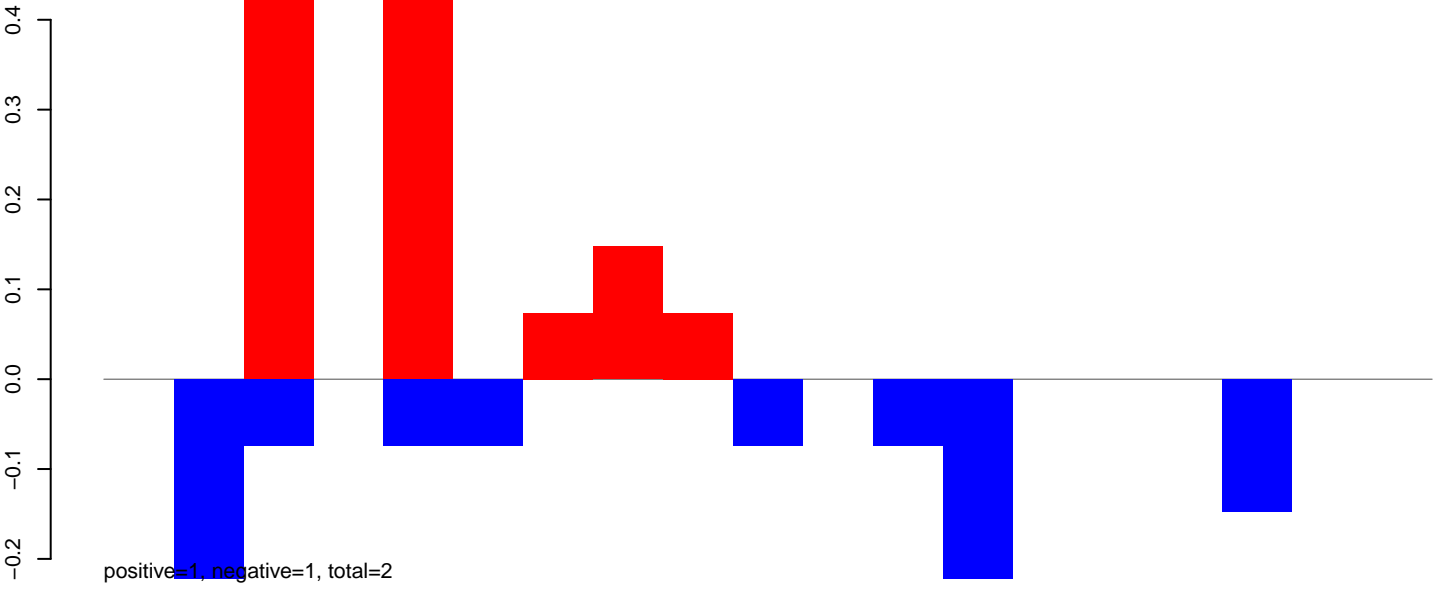
CuQuin_Testes_NL.rep



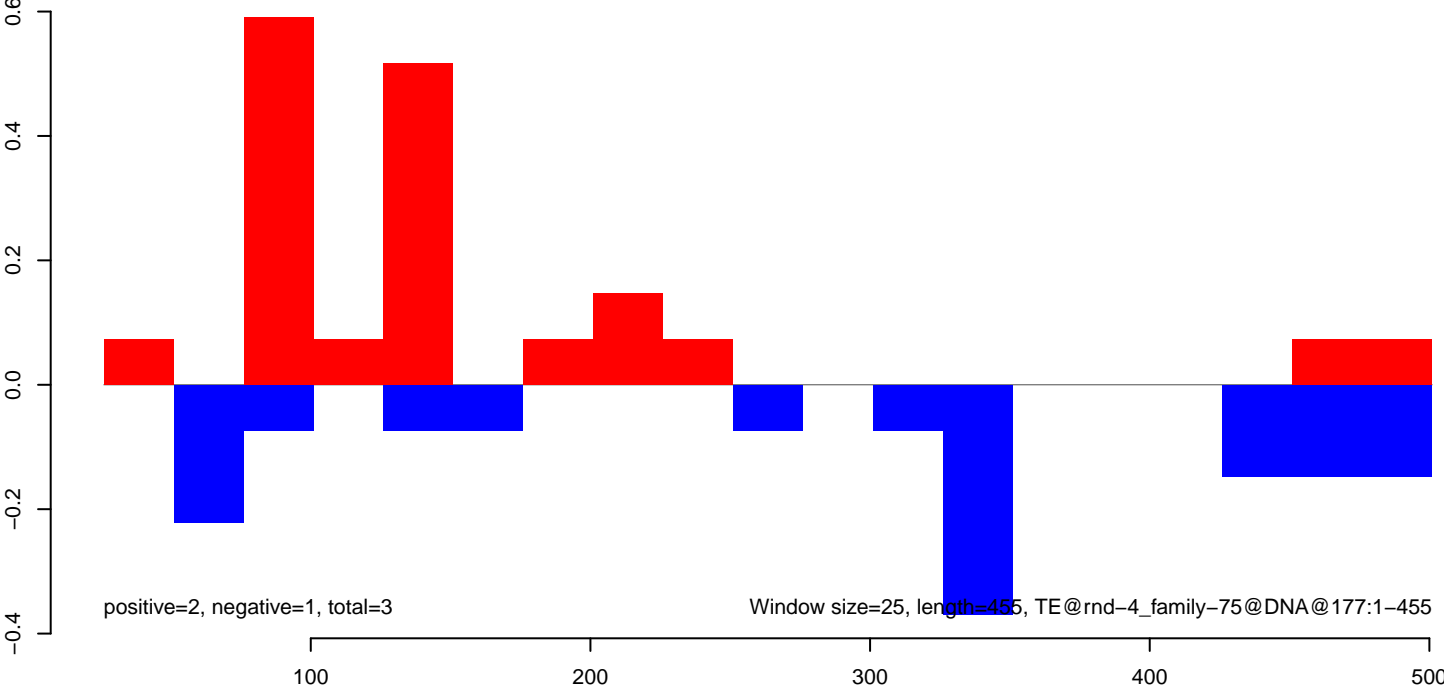
CuQuin_Testes_NL.18_23.rep



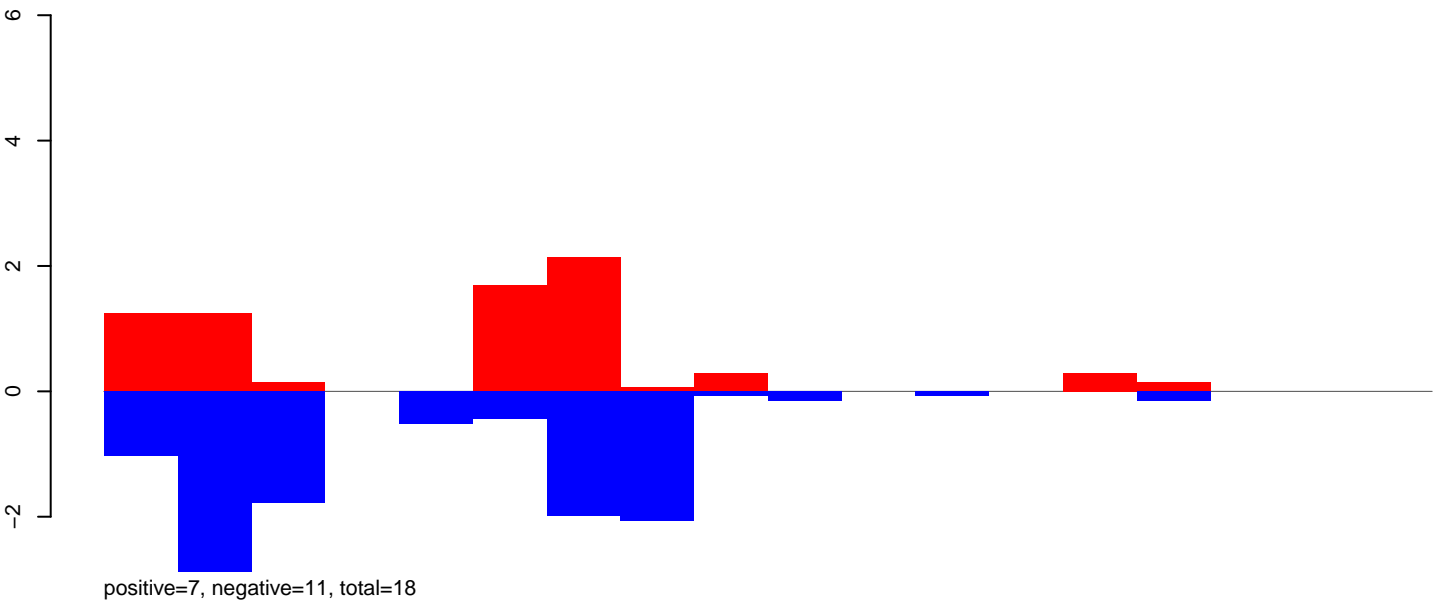
CuQuin_Testes_NL.24_35.rep



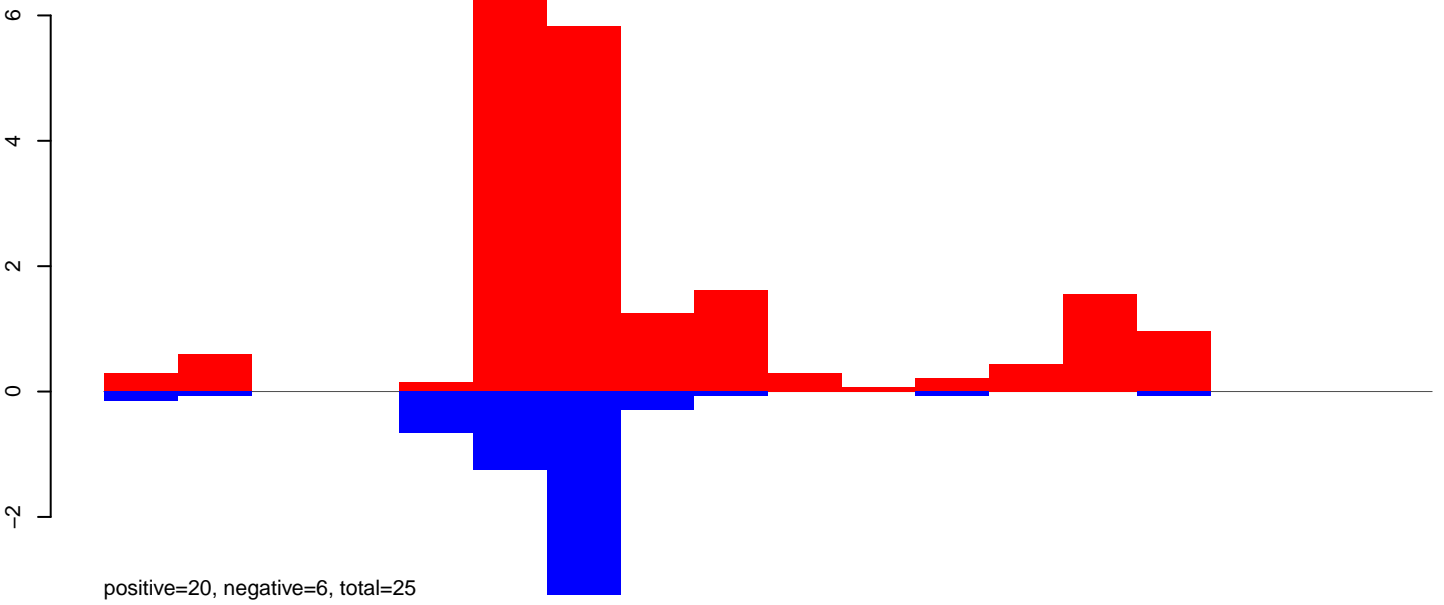
CuQuin_Testes_NL.rep



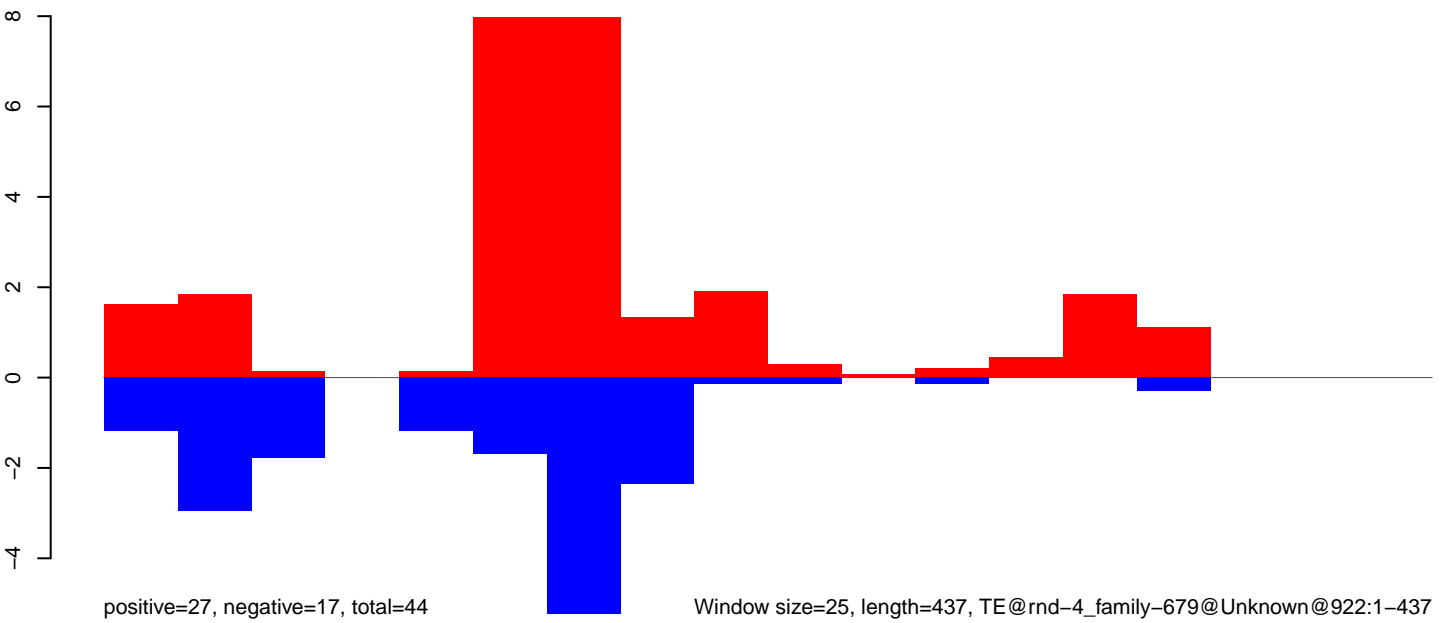
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

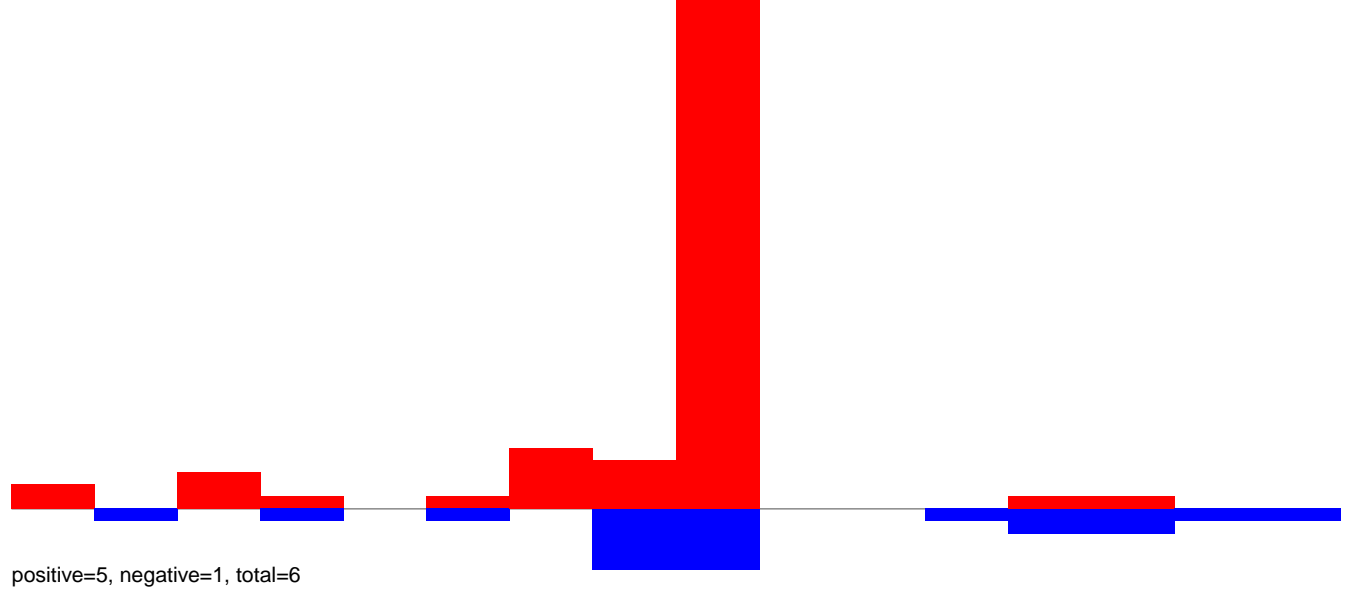


CuQuin_Testes_NL.rep

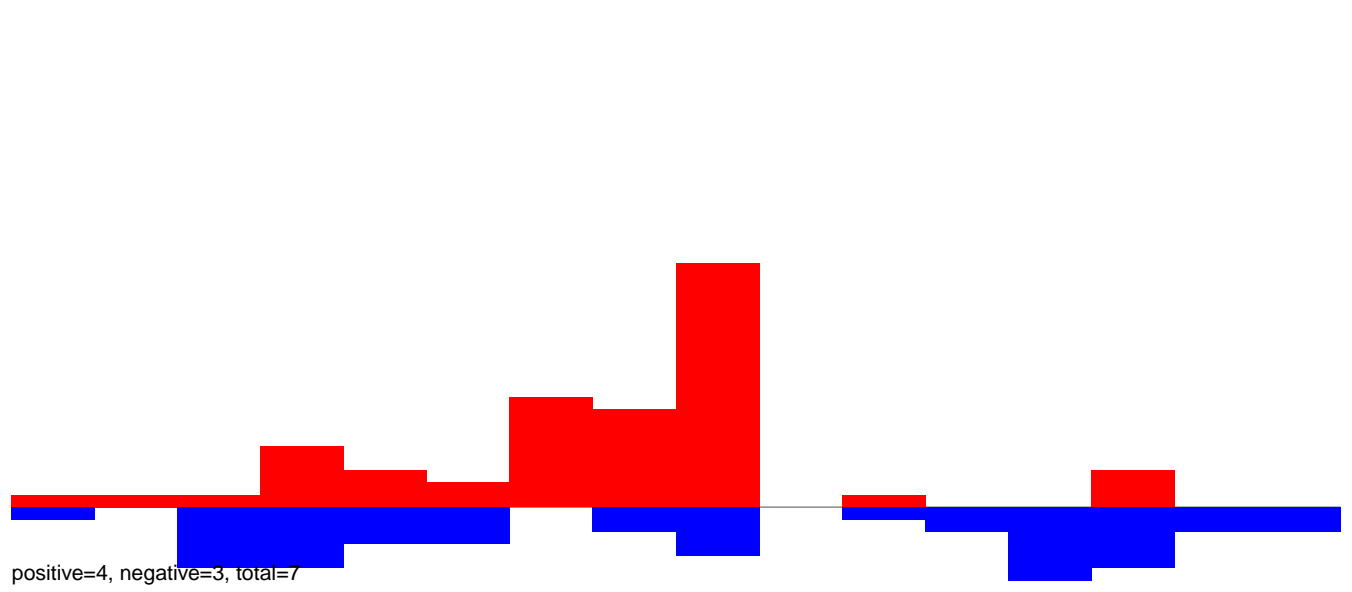


Window size=25, length=437, TE@rnd-4_family-679@Unknown@922:1-437

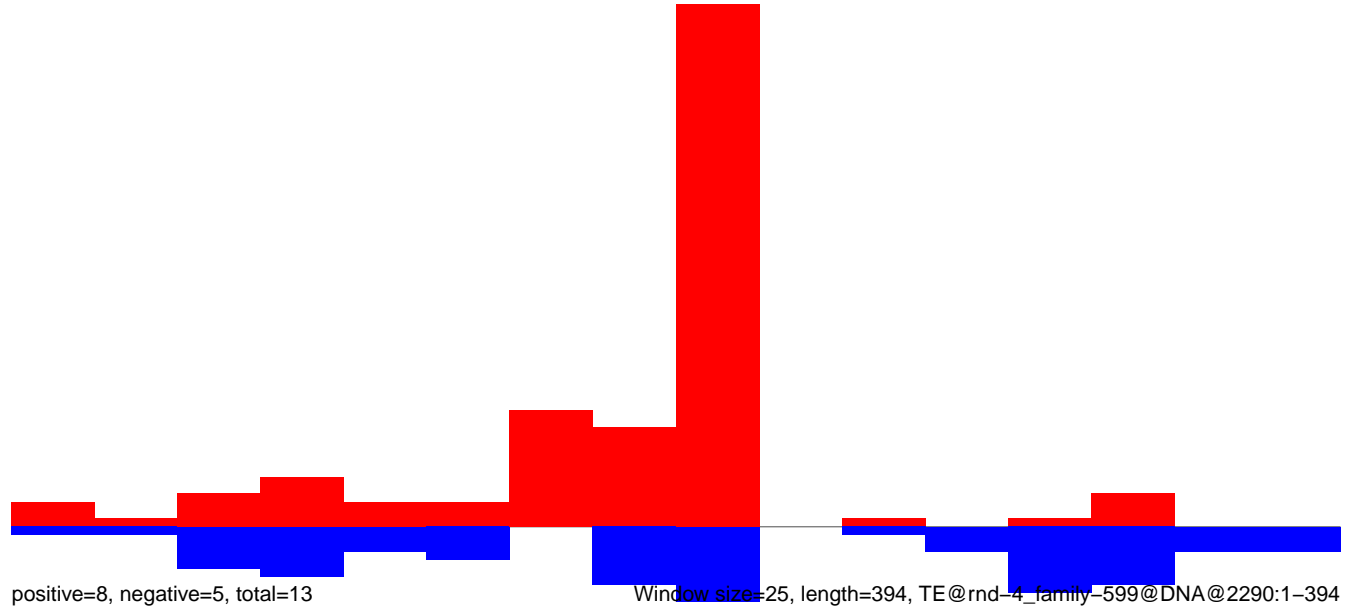
CuQuin_Testes_NL.18_23.rep



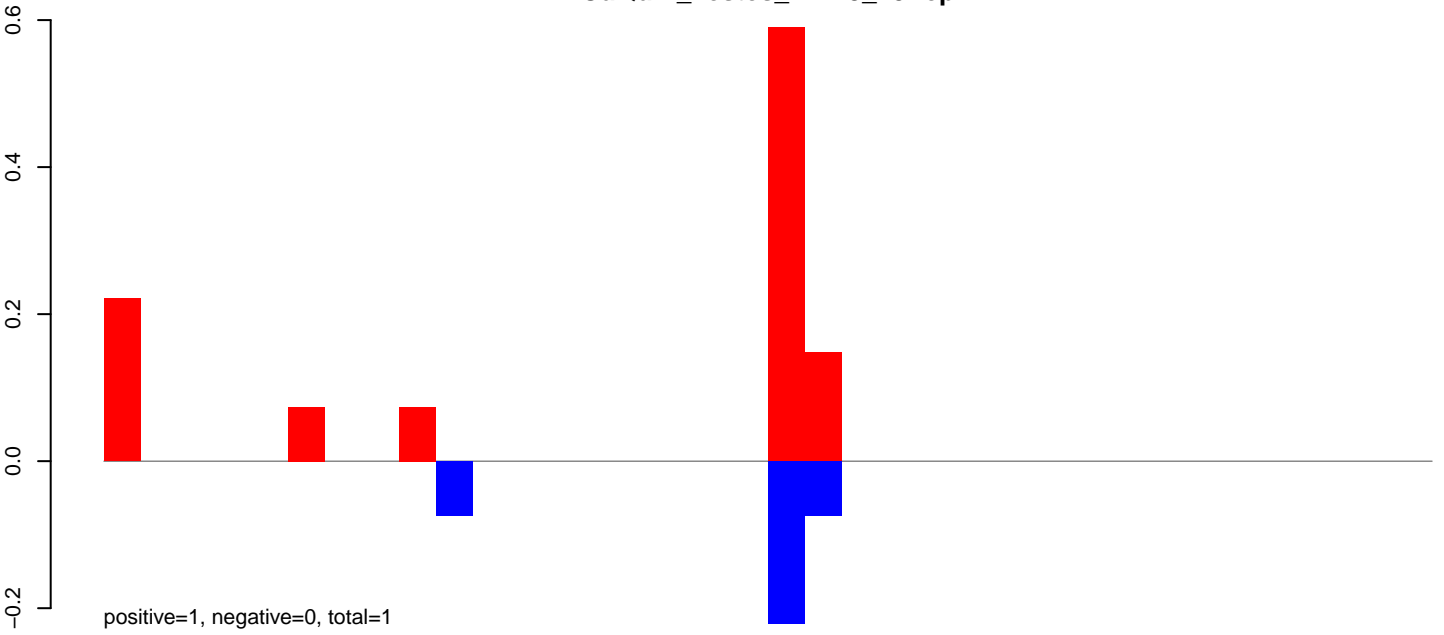
CuQuin_Testes_NL.24_35.rep



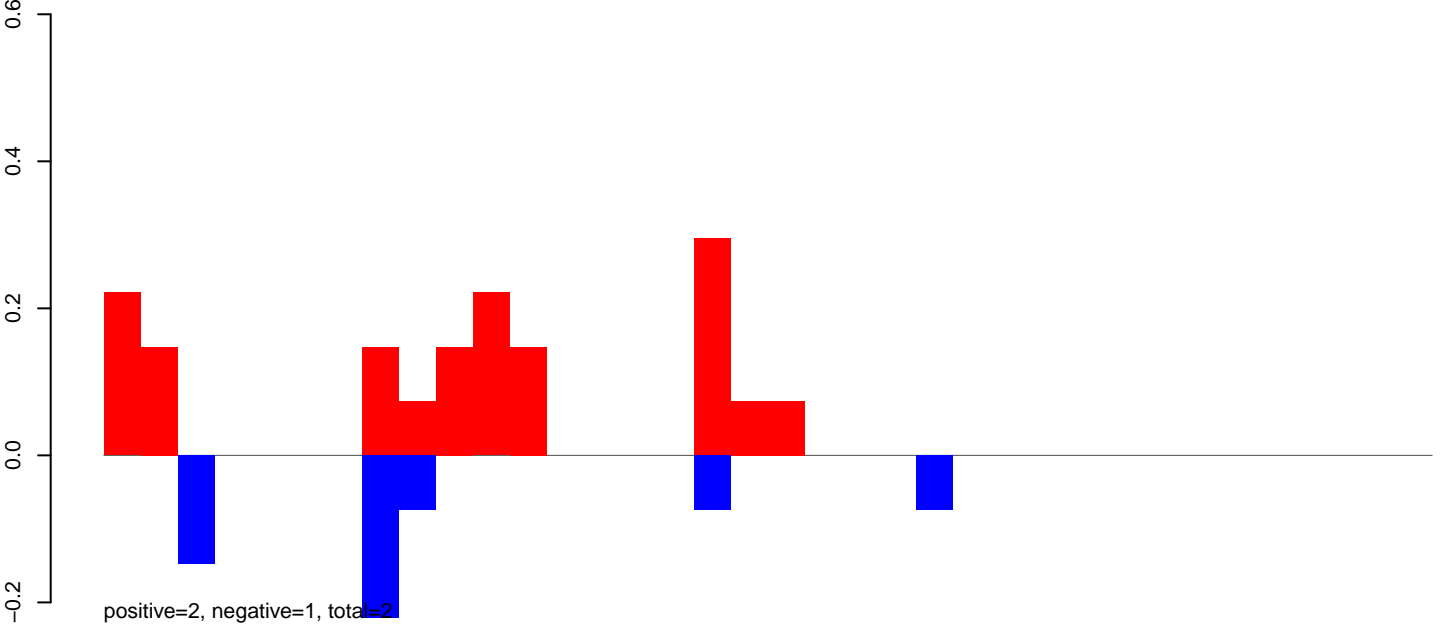
CuQuin_Testes_NL.rep



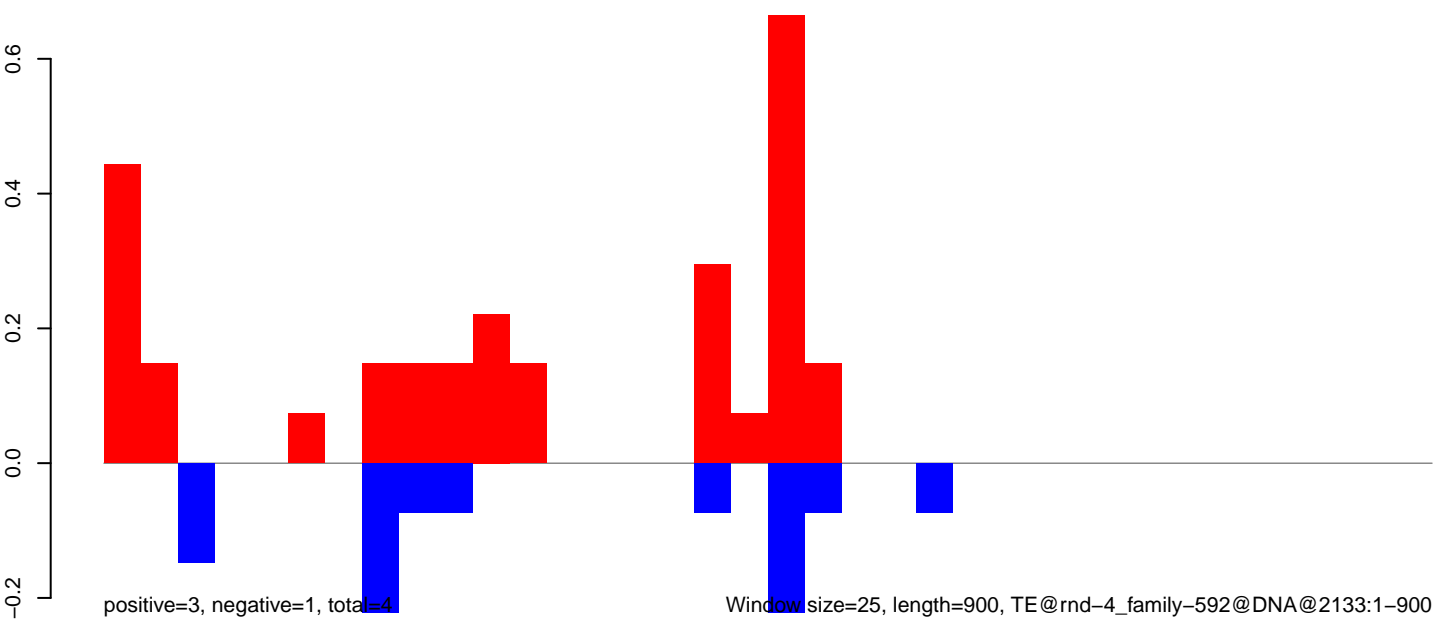
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



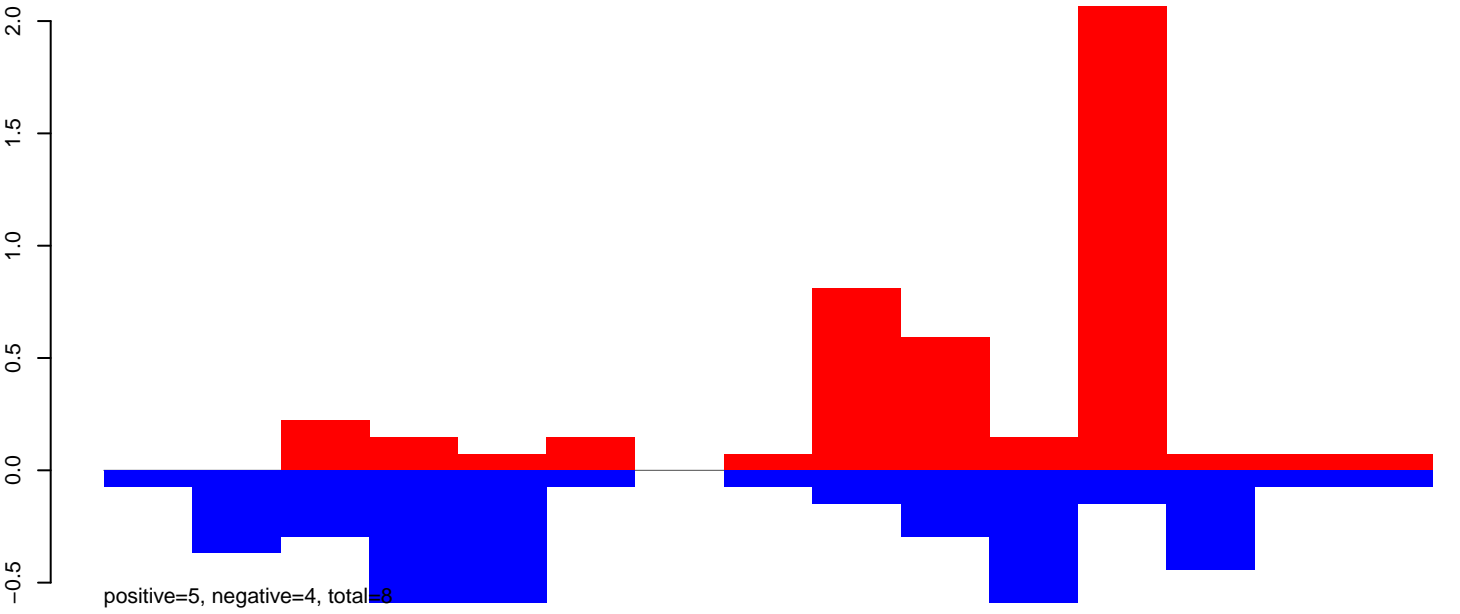
CuQuin_Testes_NL.rep



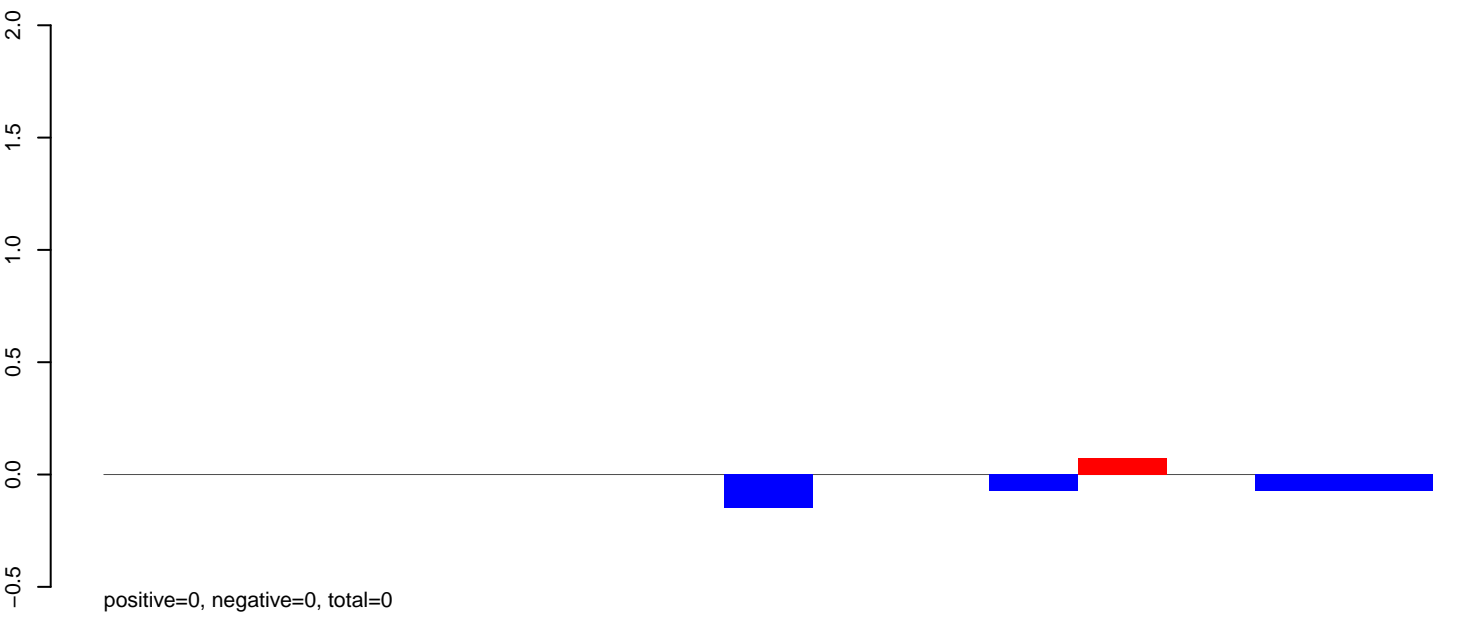
Window size=25, length=900, TE@rnd-4_family-592@DNA@2133:1-900

0 200 400 600 800

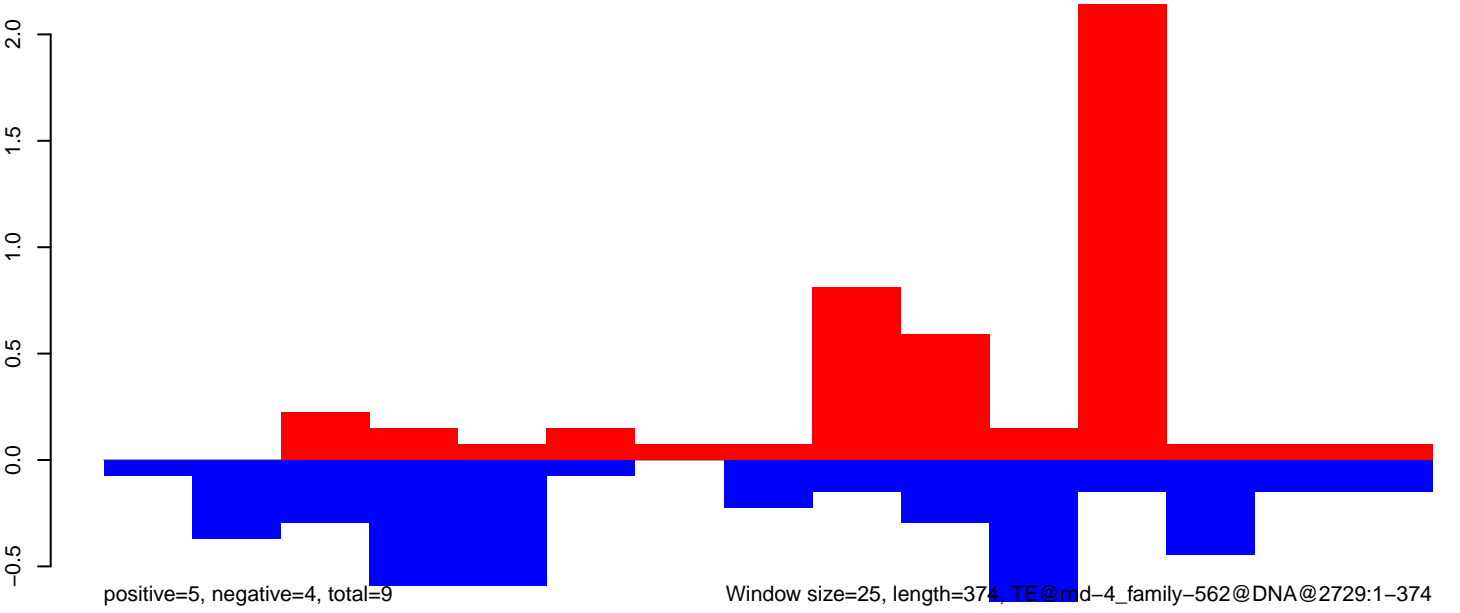
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

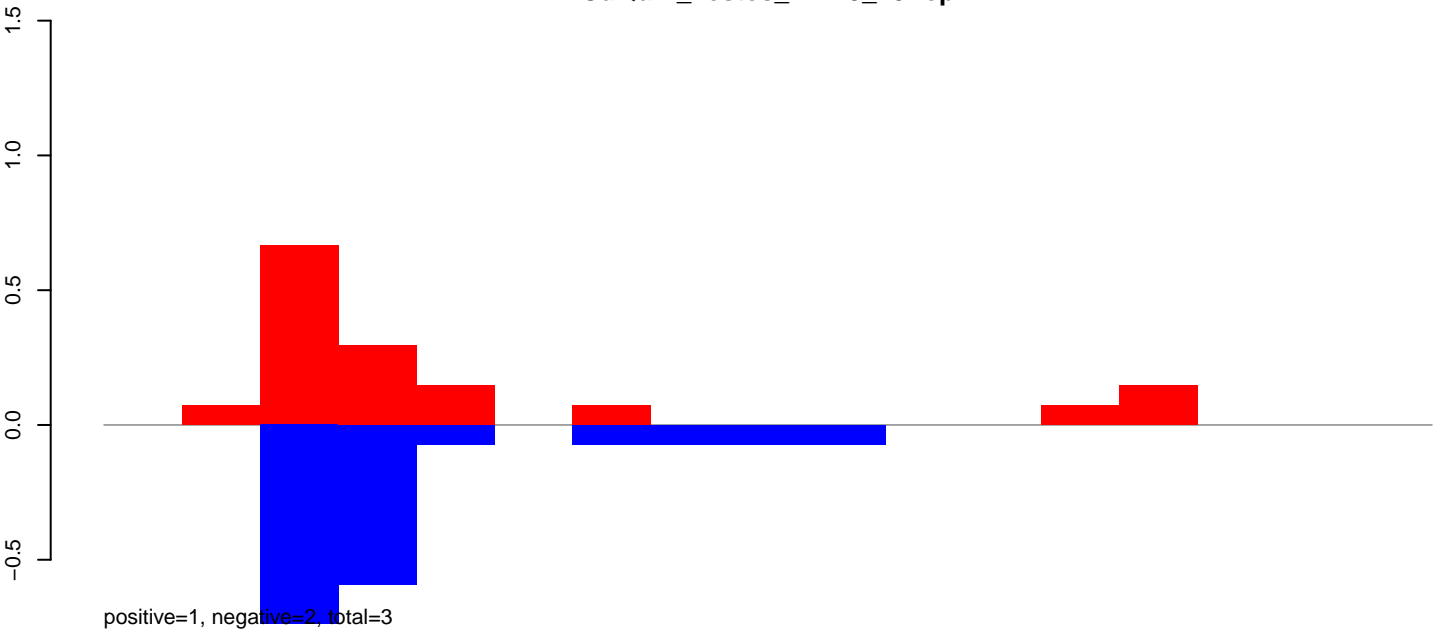


CuQuin_Testes_NL.rep

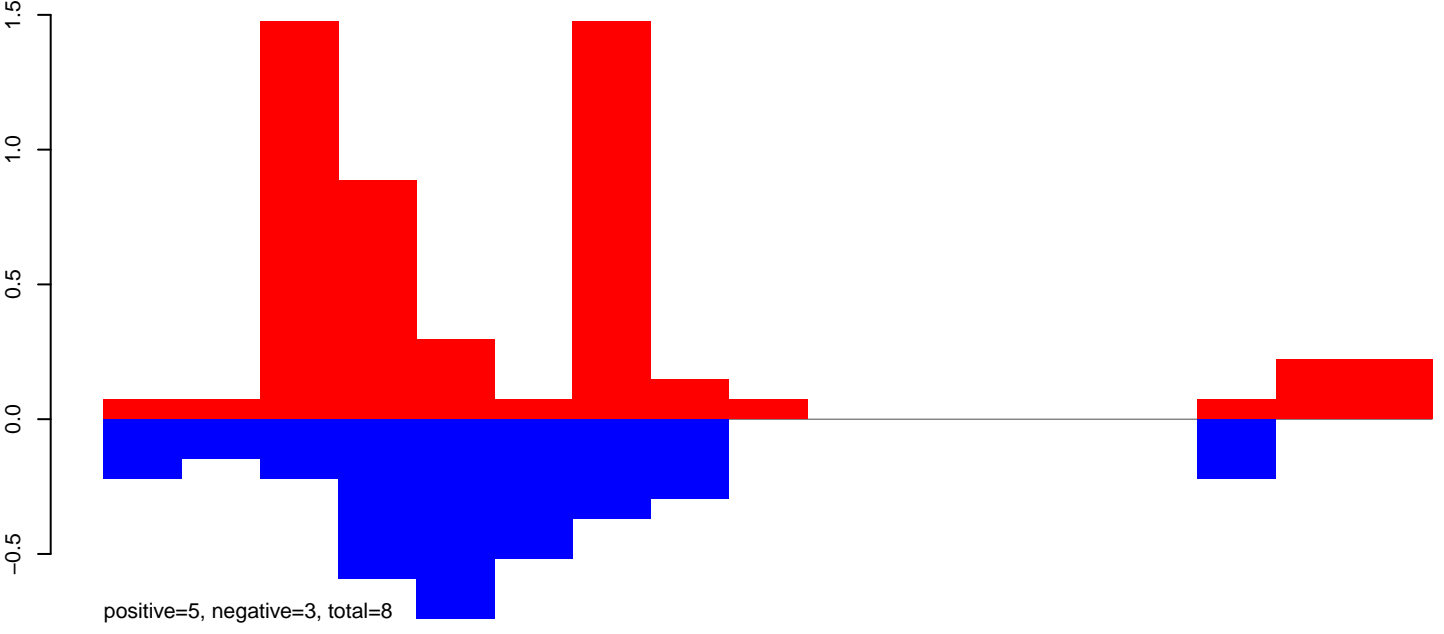


Window size=25, length=374, TE@rmd-4_family-562@DNA@2729:1-374

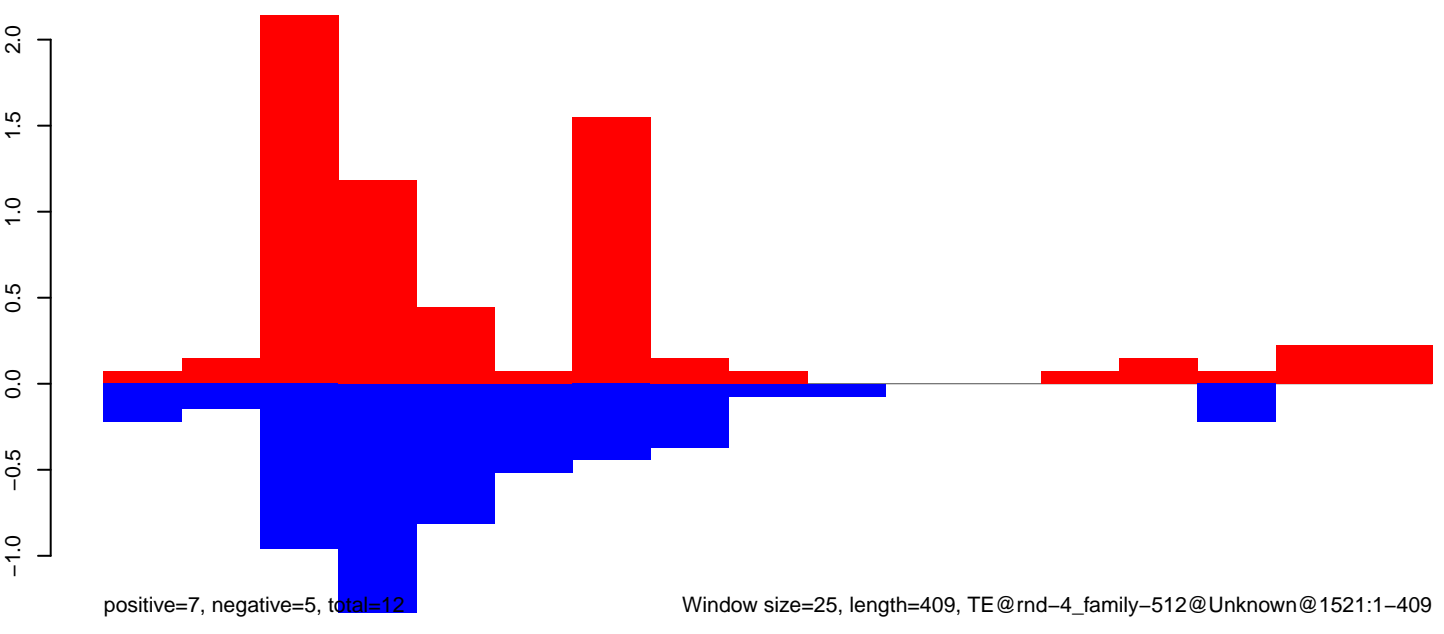
CuQuin_Testes_NL.18_23.rep



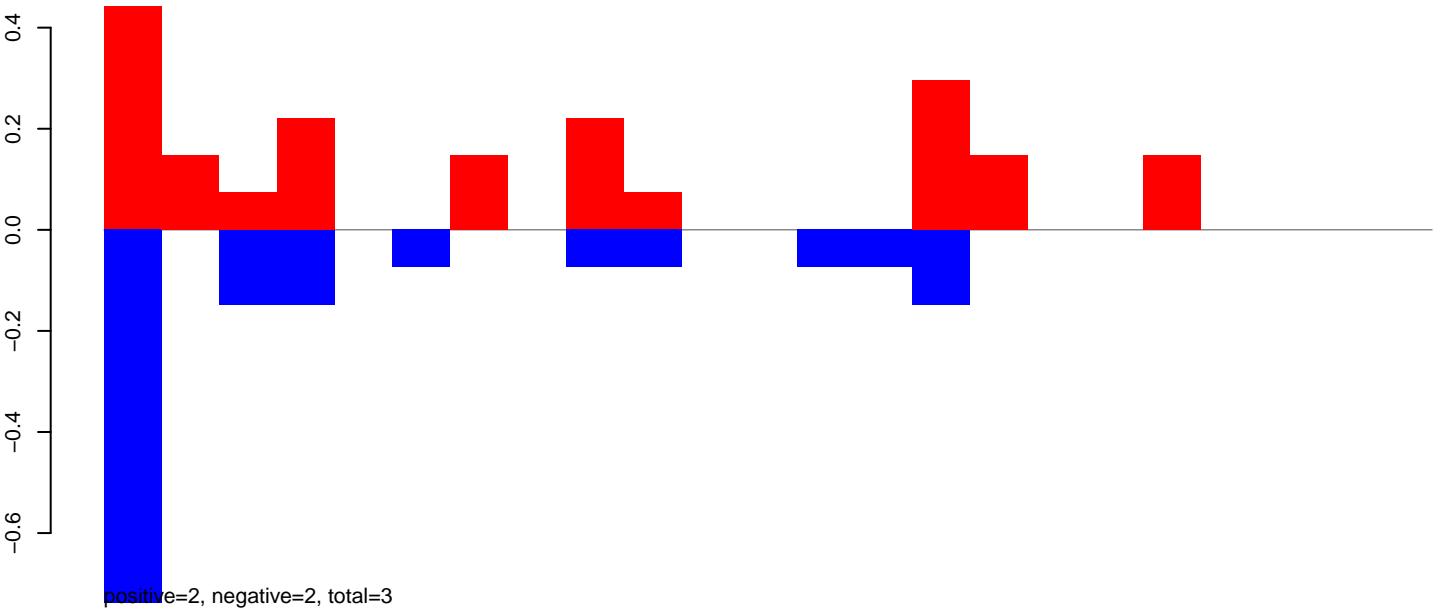
CuQuin_Testes_NL.24_35.rep



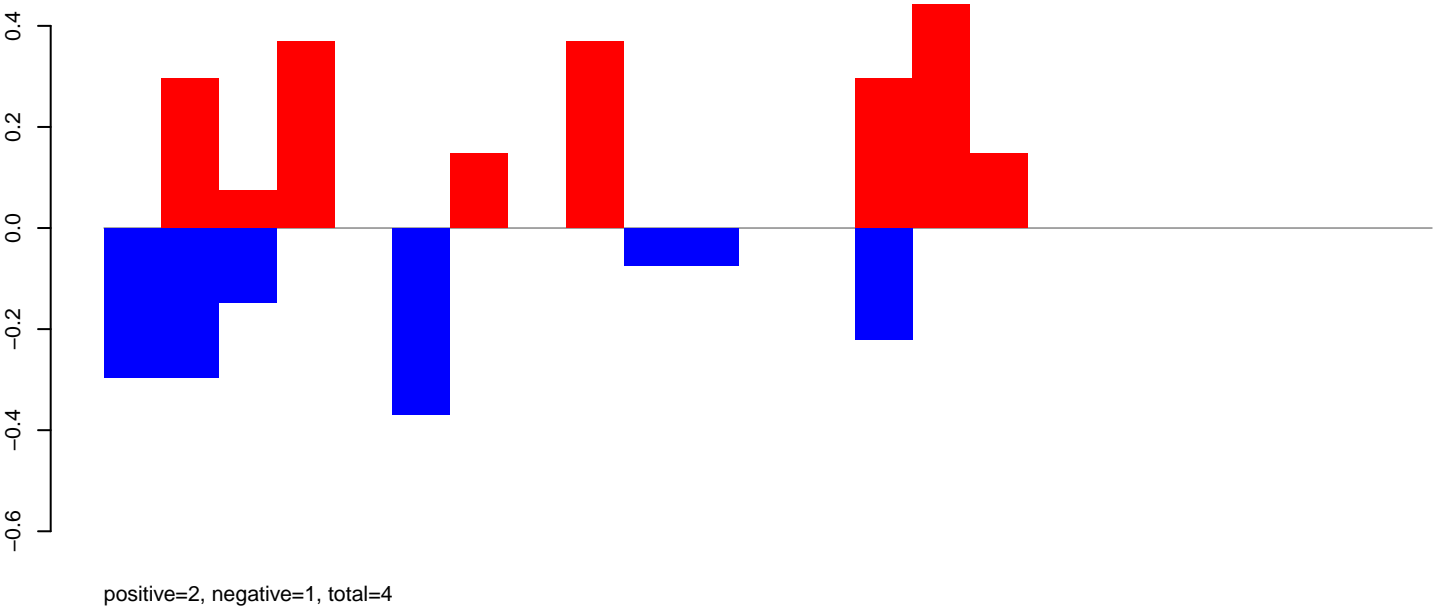
CuQuin_Testes_NL.rep



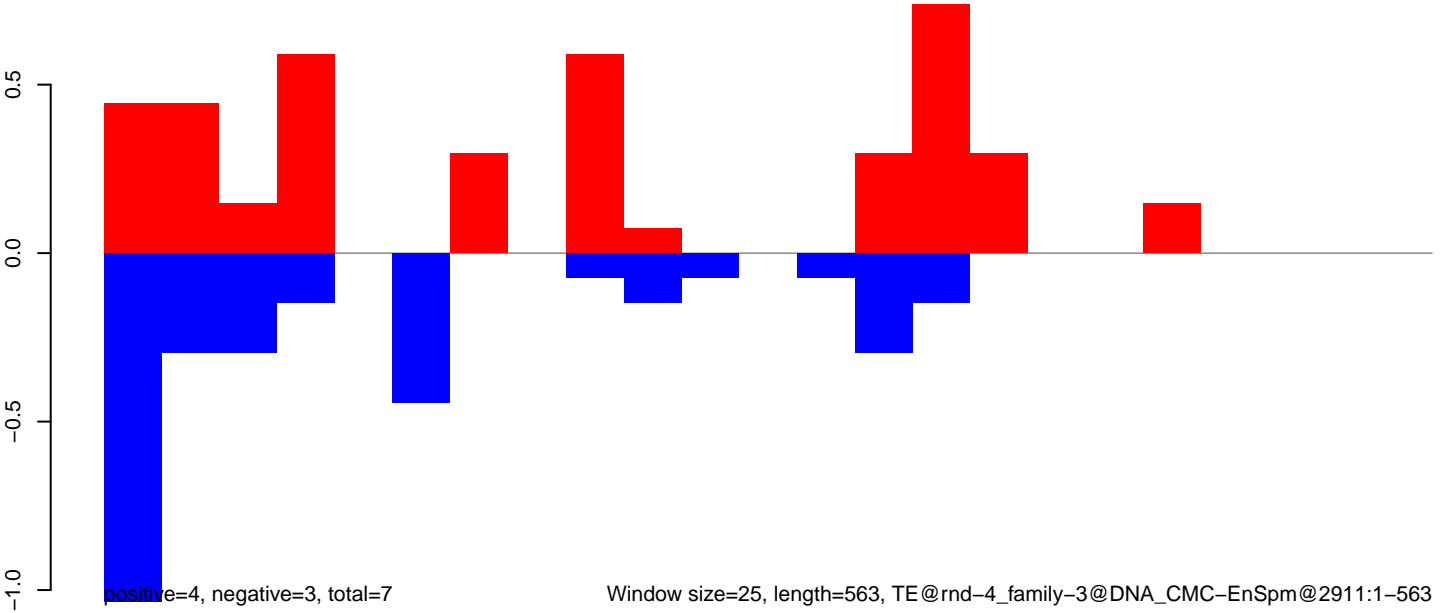
CuQuin_Testes_NL.18_23.rep



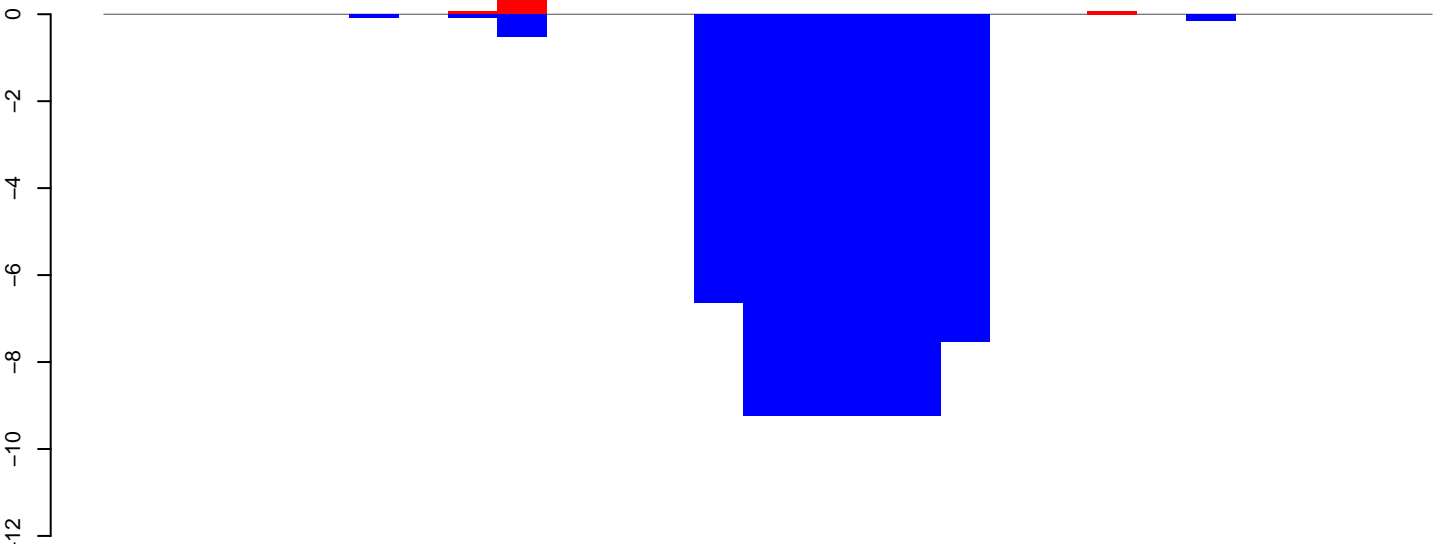
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

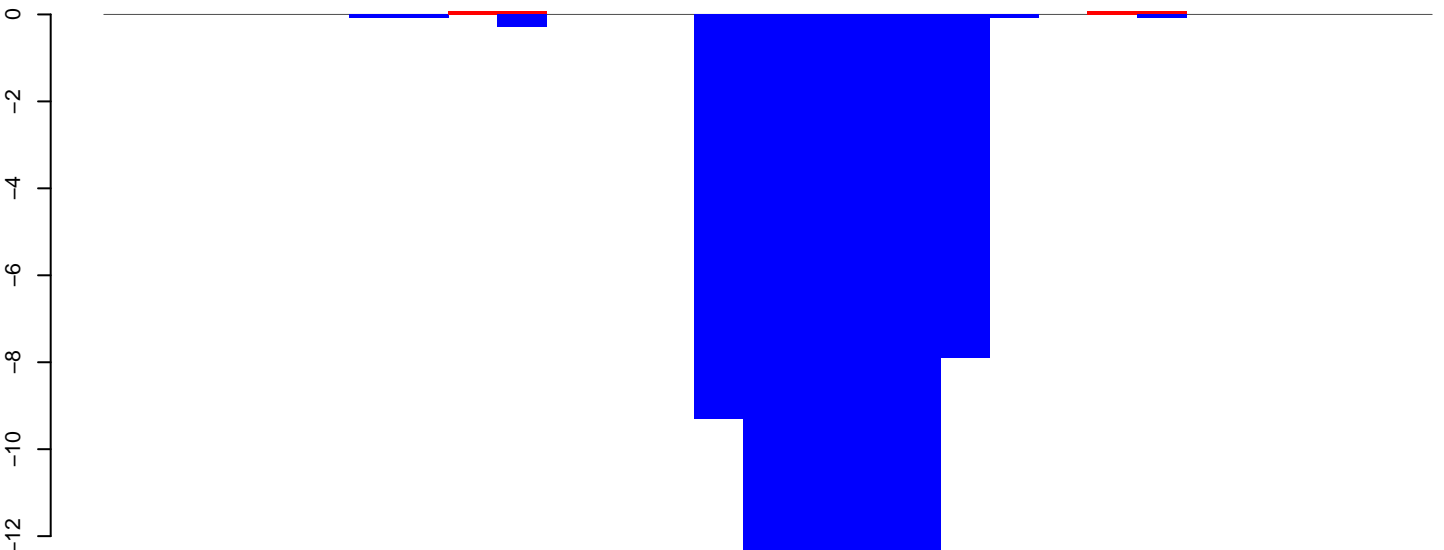


CuQuin_Testes_NL.18_23.rep



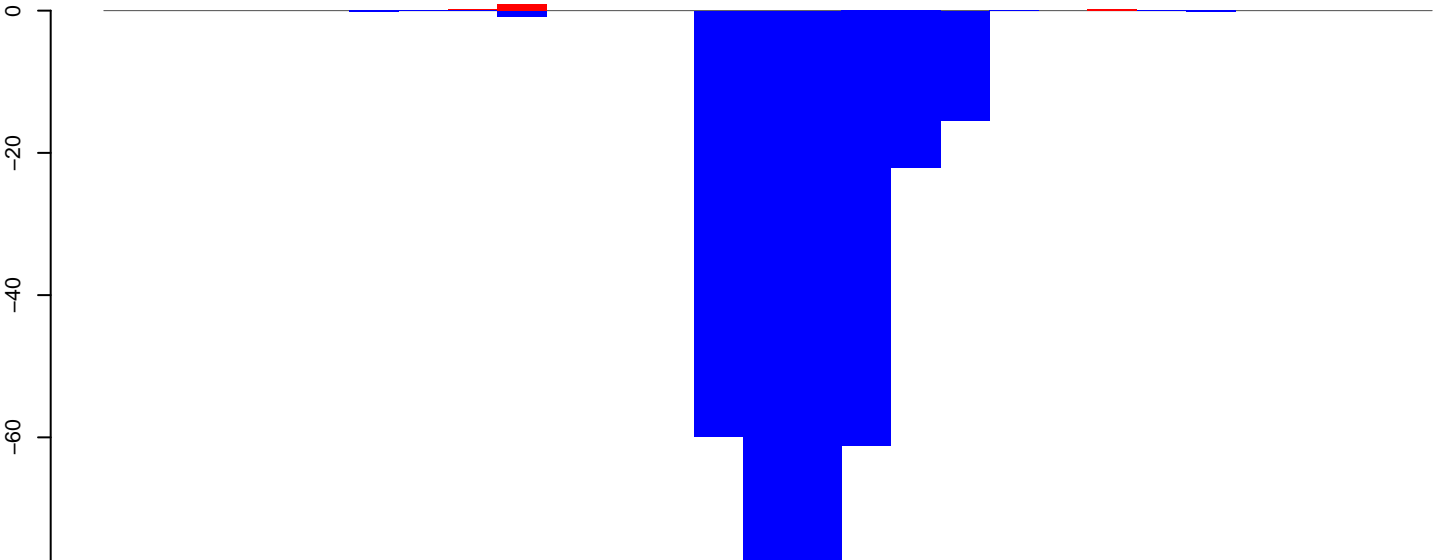
positive=1, negative=52, total=53

CuQuin_Testes_NL.24_35.rep



positive=0, negative=69, total=70

CuQuin_Testes_NL.rep

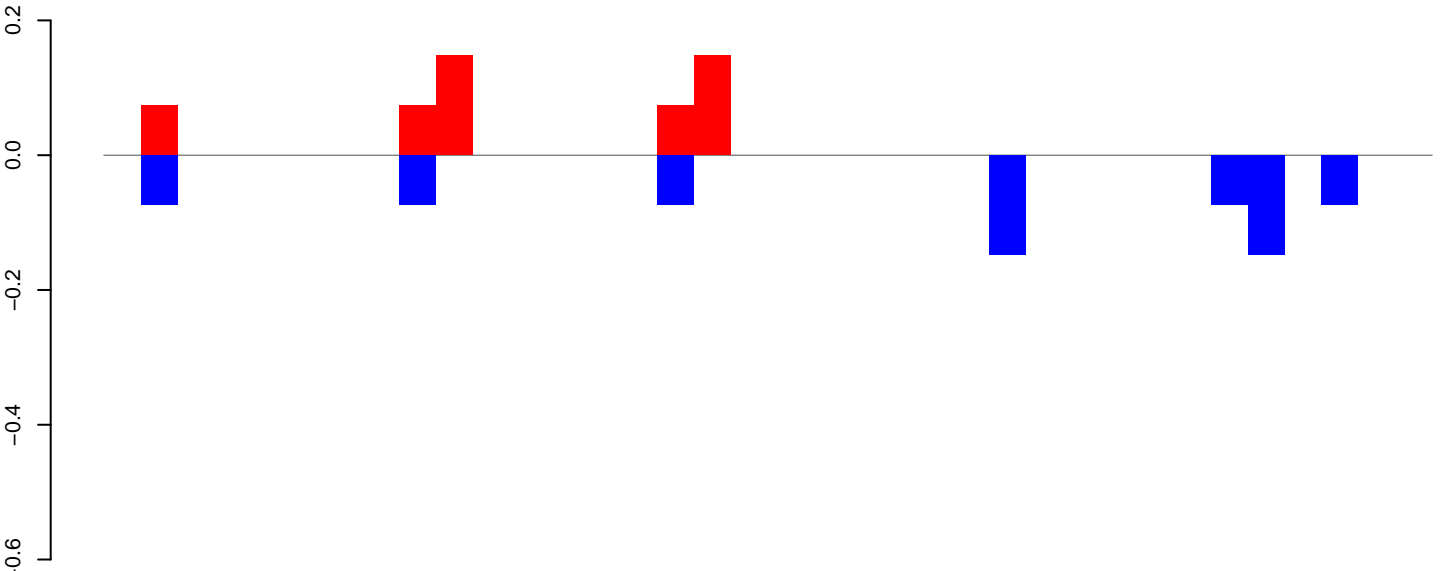


positive=1, negative=326, total=327

Window size=25, length=655, TE@rnd-4_family-149@DNA_MULE-MuDR@961:1-655

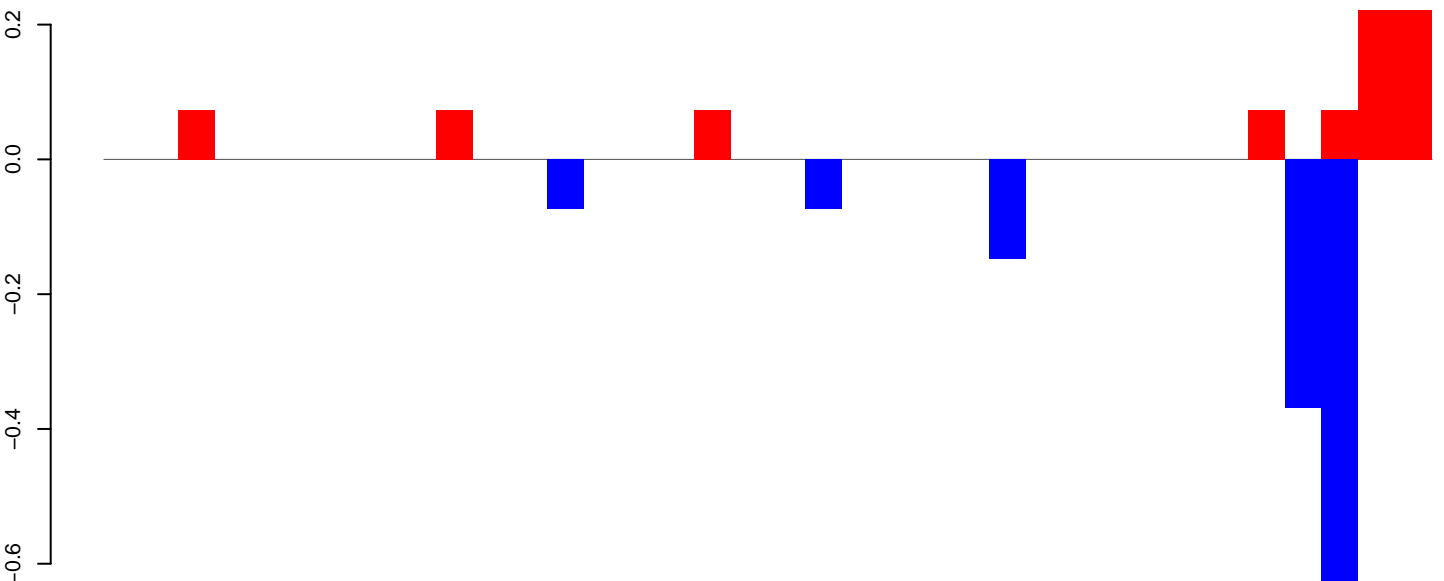
0 100 200 300 400 500 600 700

CuQuin_Testes_NL.18_23.rep



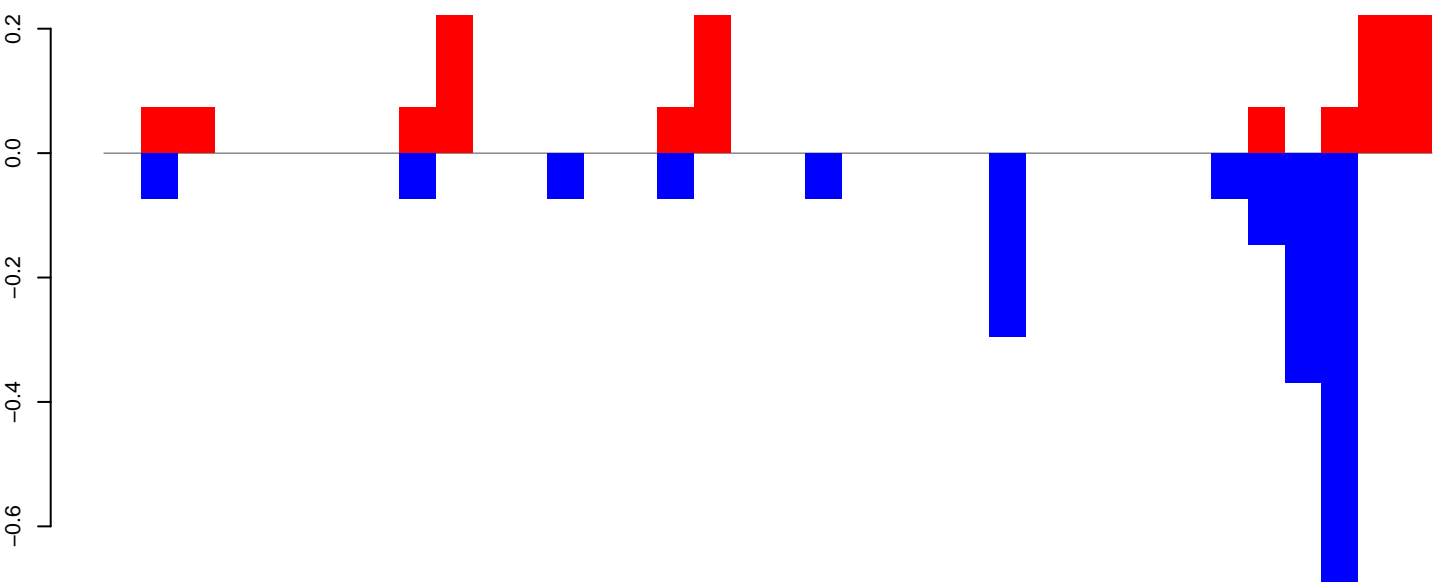
positive=1, negative=1, total=1

CuQuin_Testes_NL.24_35.rep



positive=1, negative=1, total=2

CuQuin_Testes_NL.rep

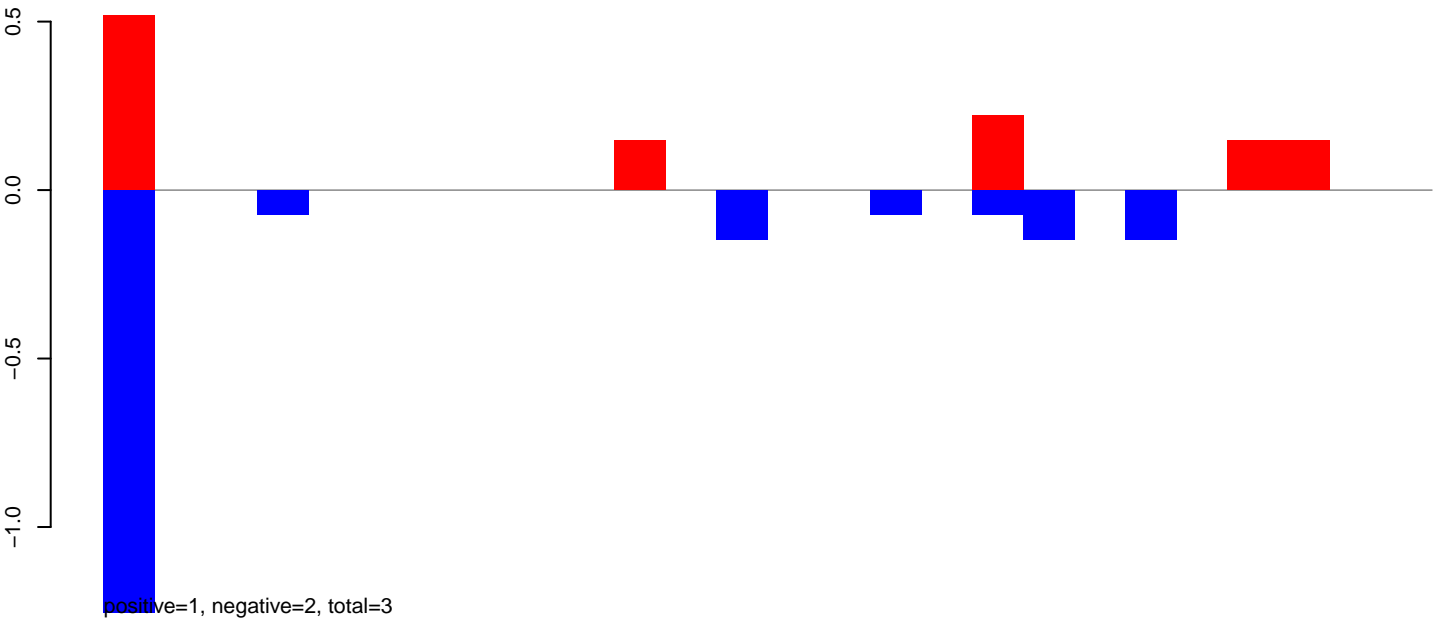


positive=1, negative=2, total=3

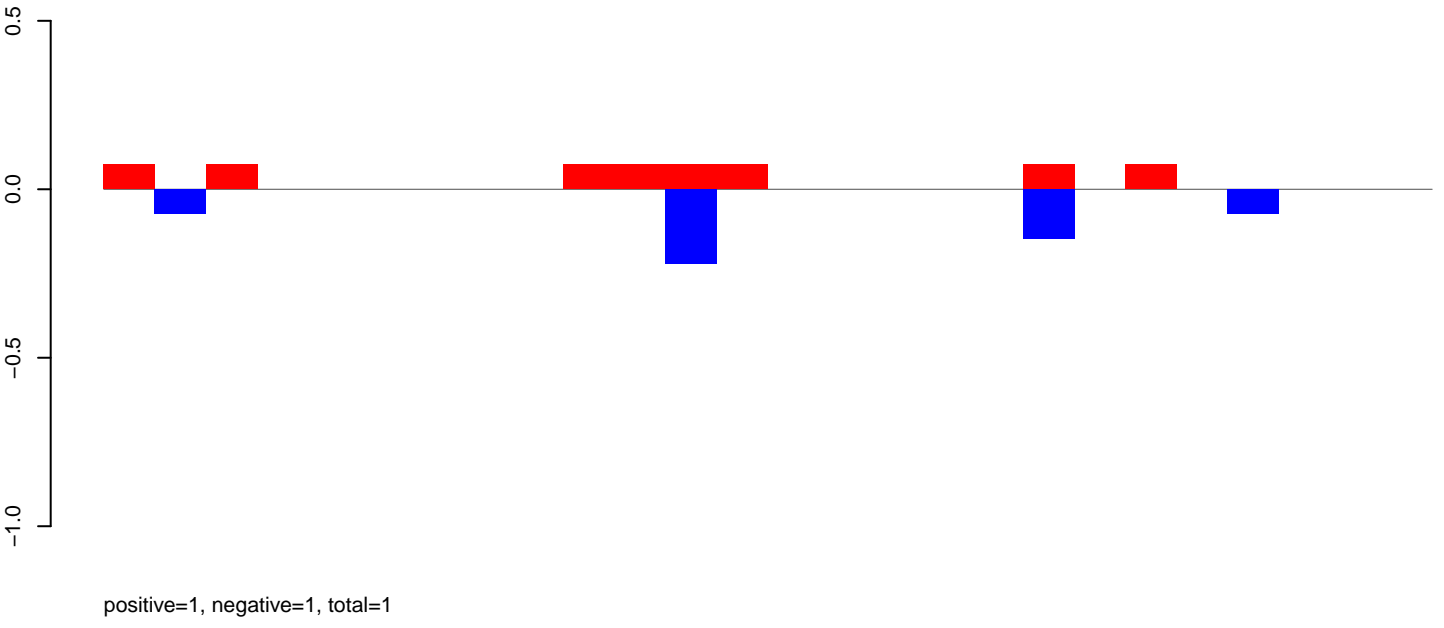
Window size=25, length=882, TE@rnd-4_family-1421@Unknown@328:1-882

0 200 400 600 800

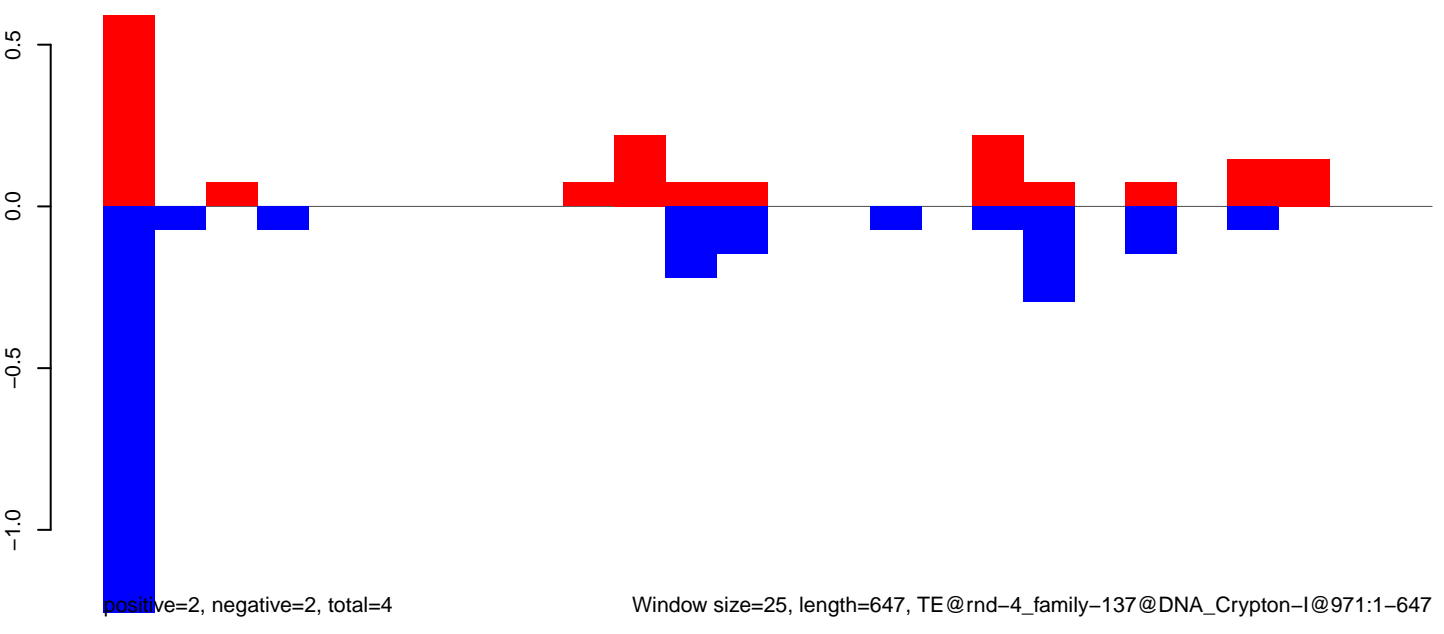
CuQuin_Testes_NL.18_23.rep



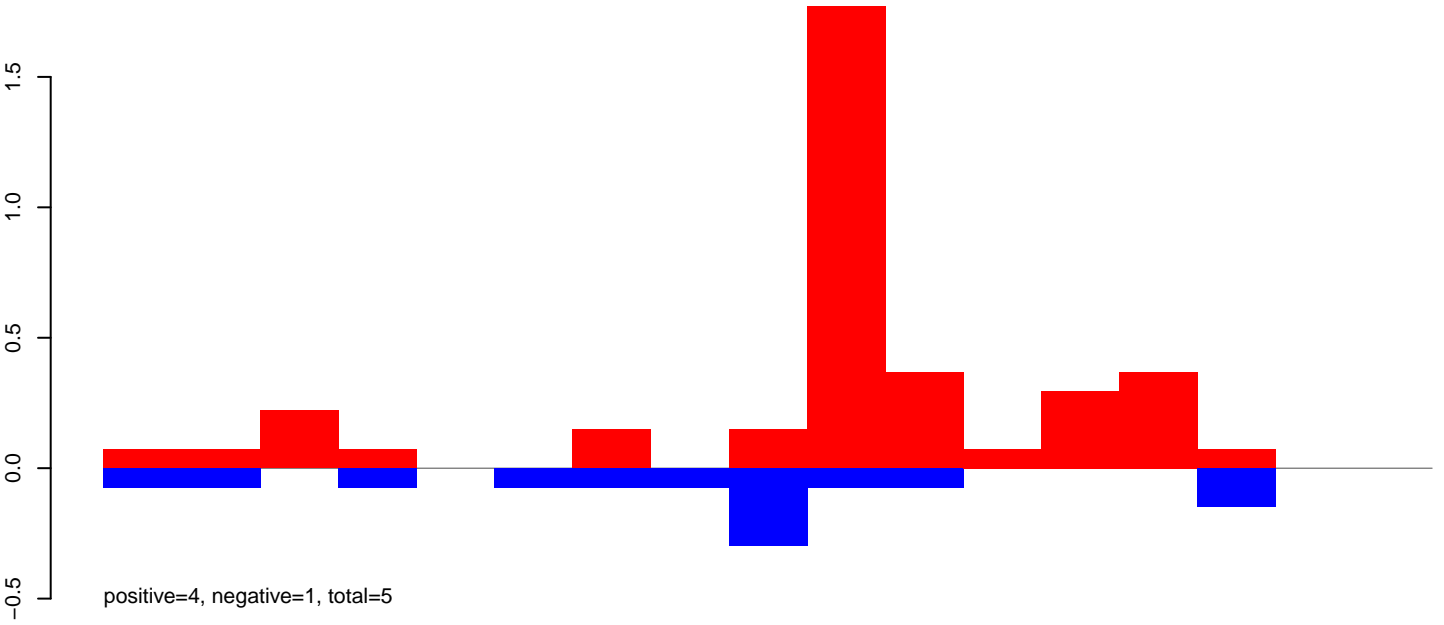
CuQuin_Testes_NL.24_35.rep



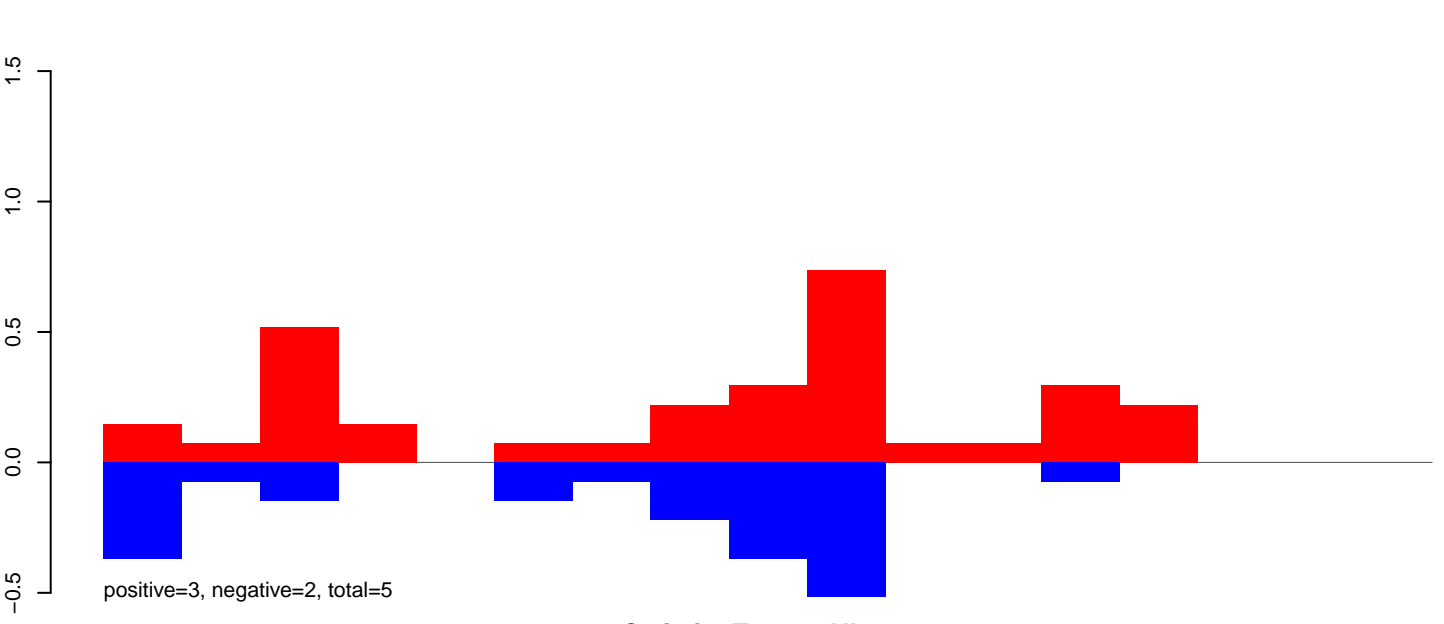
CuQuin_Testes_NL.rep



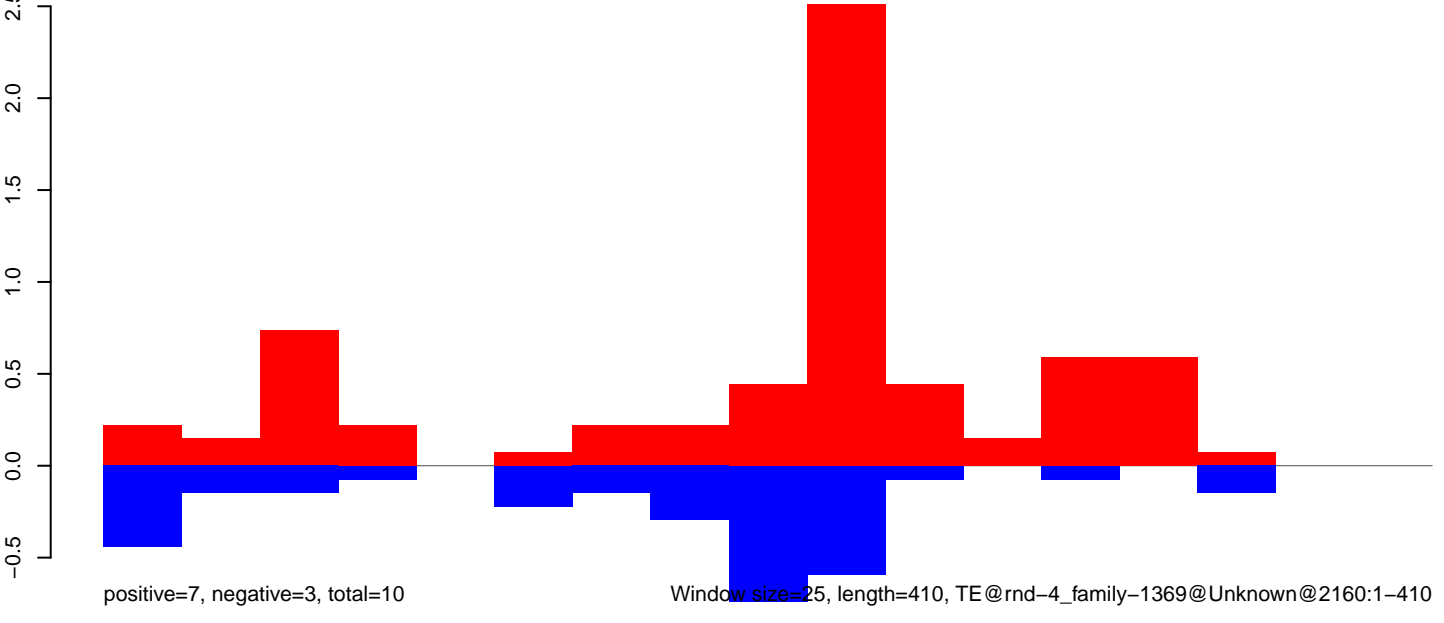
CuQuin_Testes_NL.18_23.rep



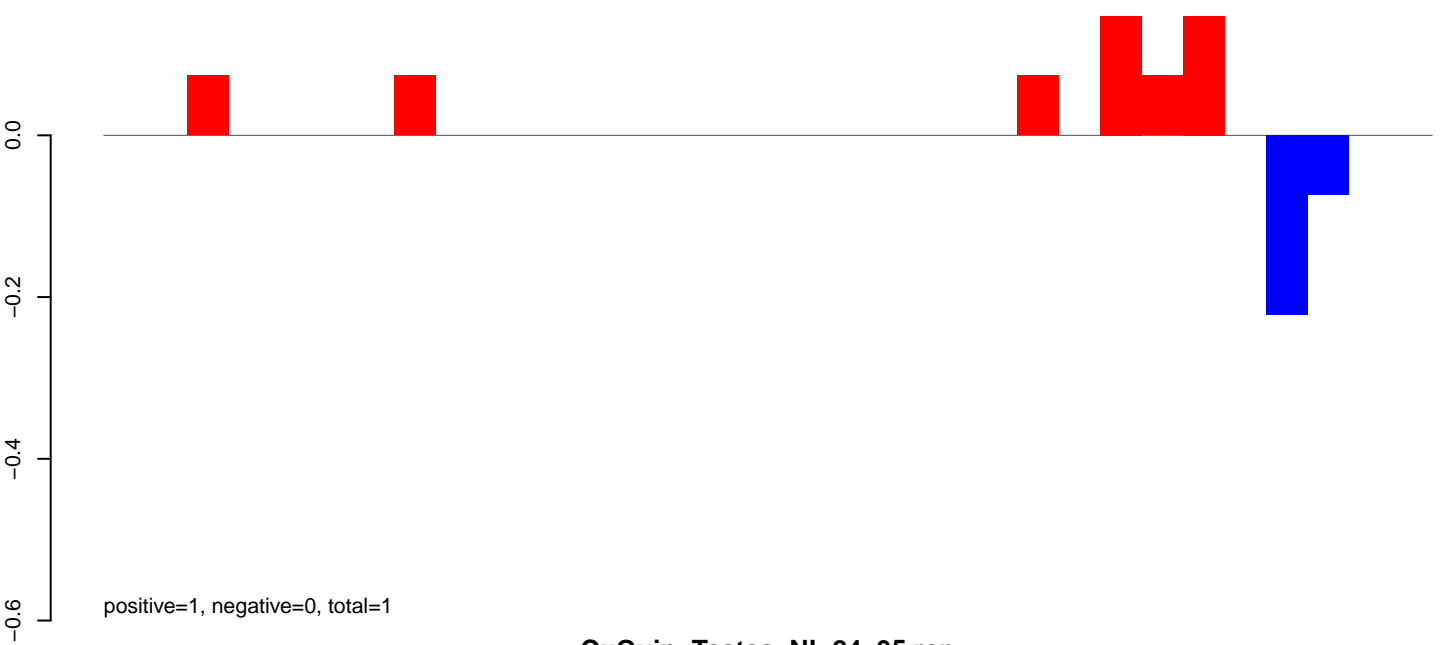
CuQuin_Testes_NL.24_35.rep



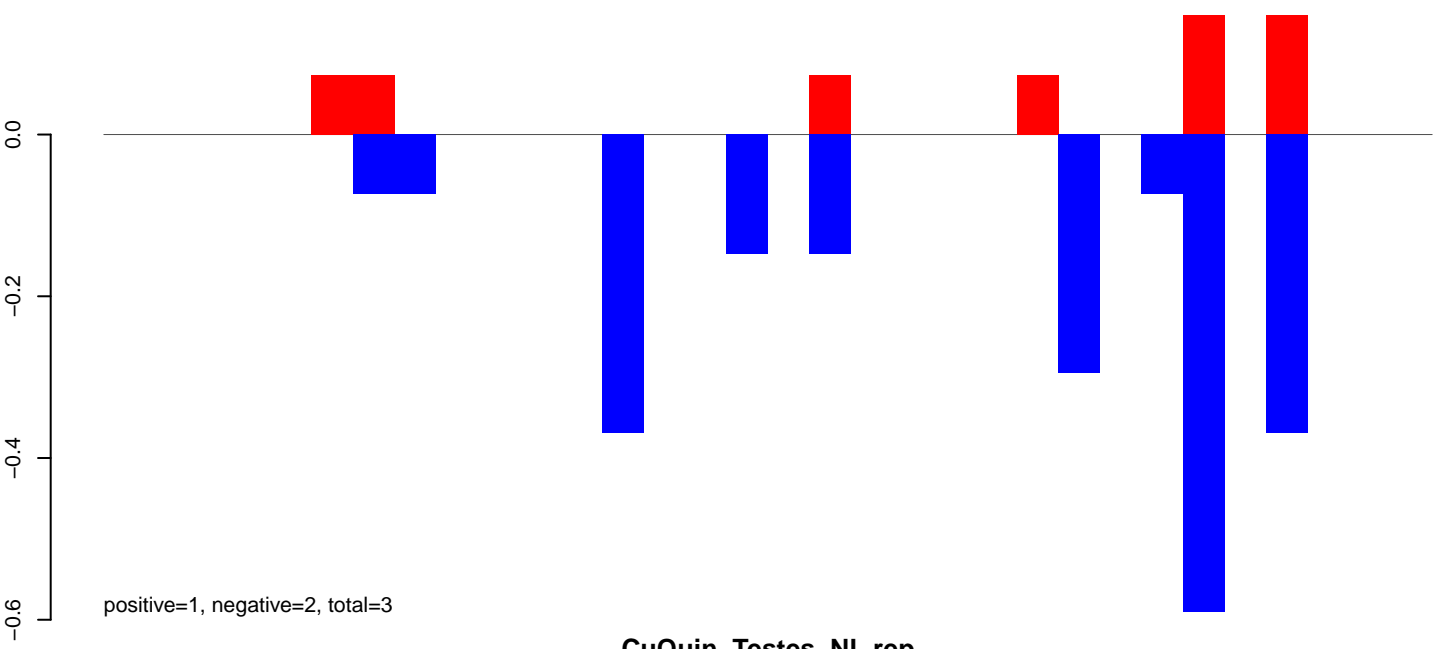
CuQuin_Testes_NL.rep



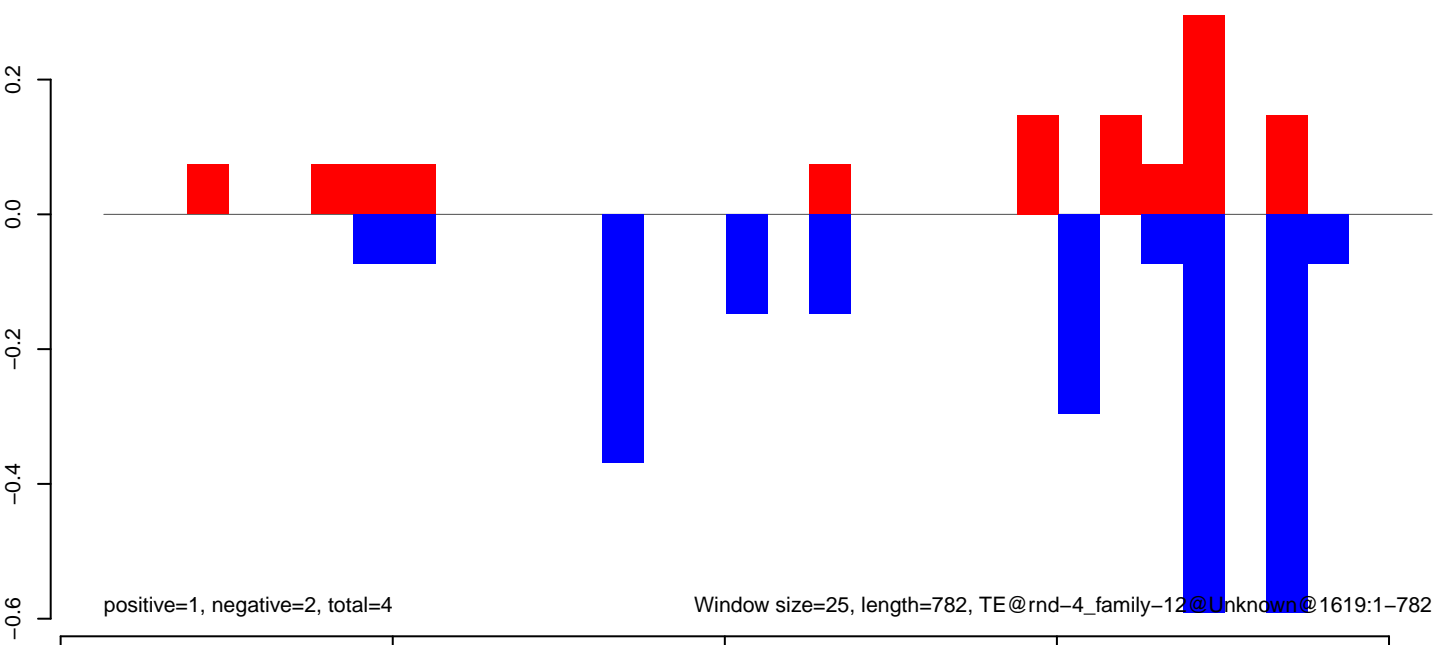
CuQuin_Testes_NL.18_23.rep



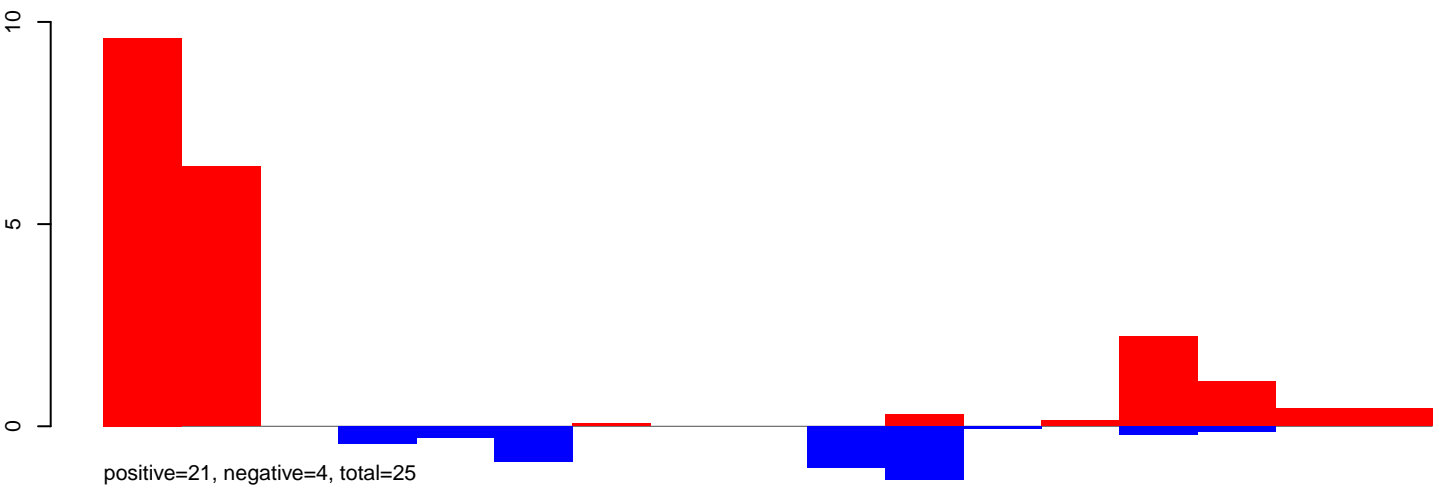
CuQuin_Testes_NL.24_35.rep



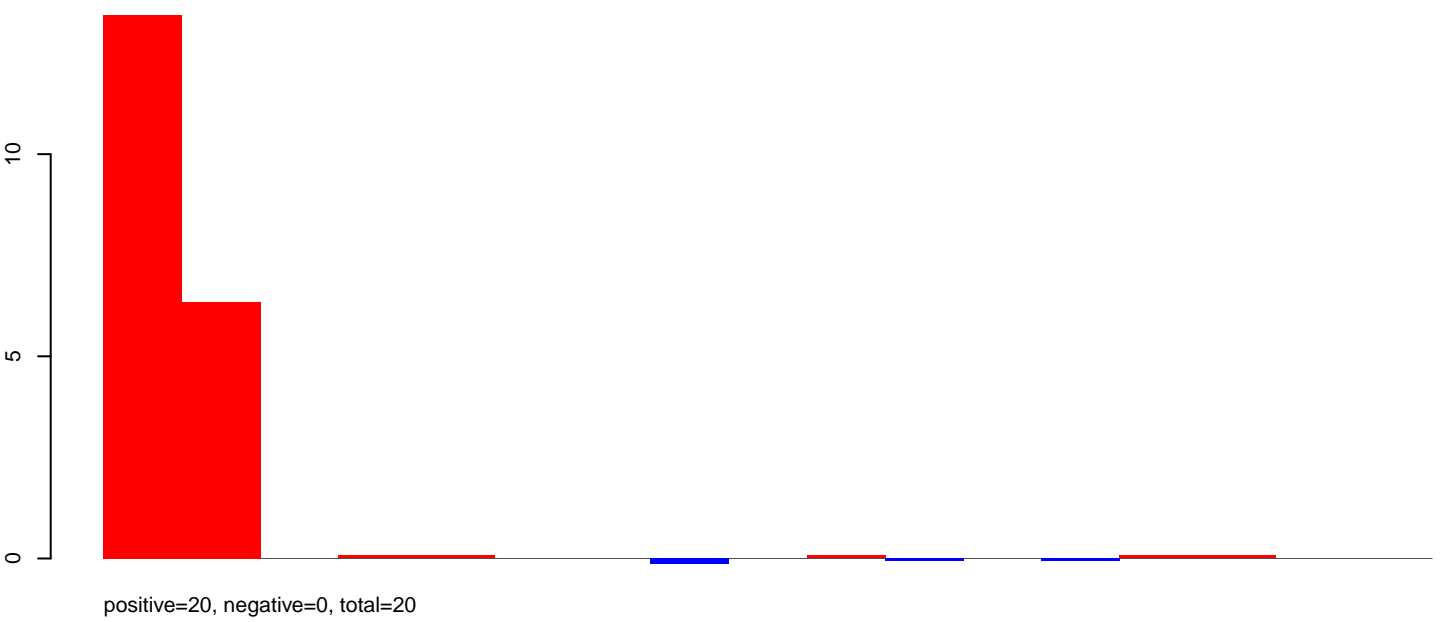
CuQuin_Testes_NL.rep



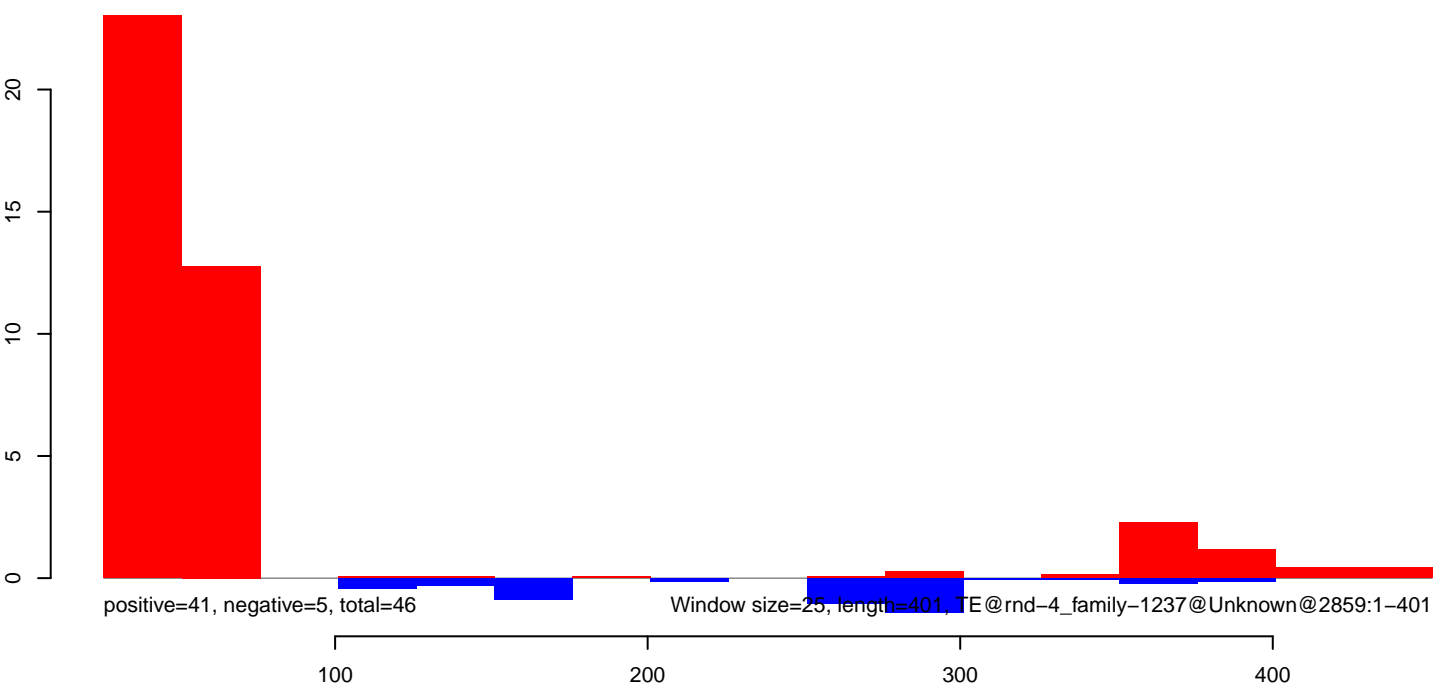
CuQuin_Testes_NL.18_23.rep



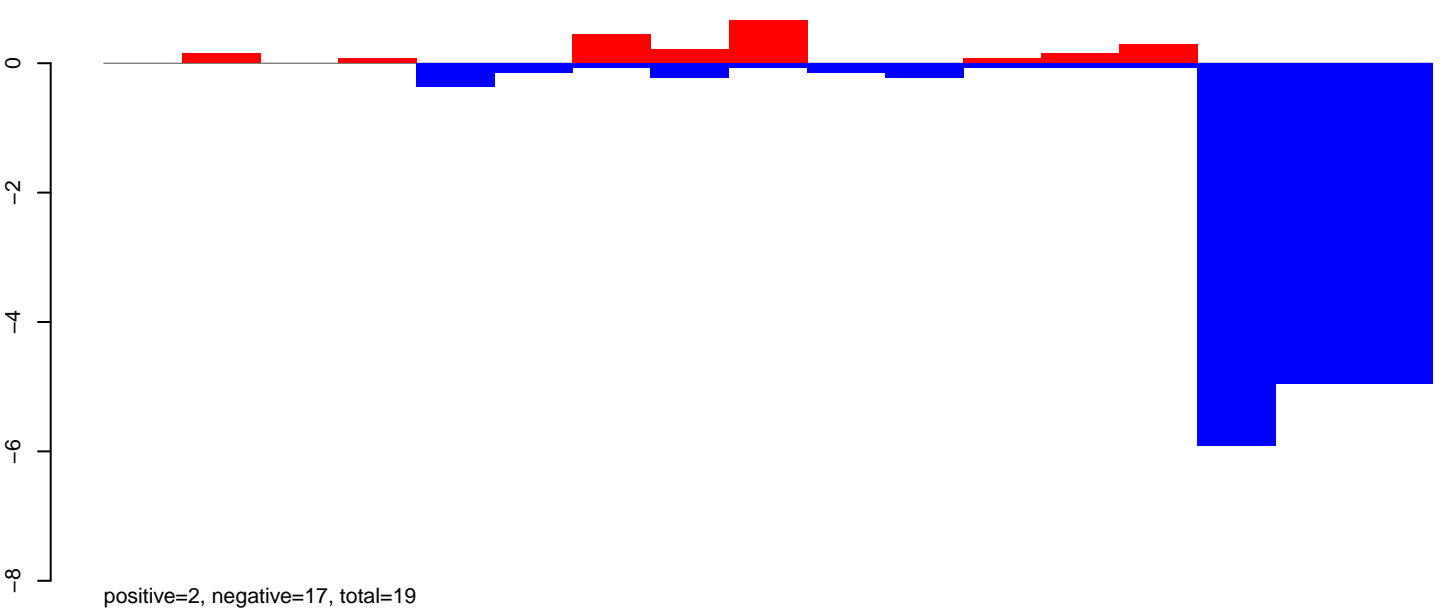
CuQuin_Testes_NL.24_35.rep



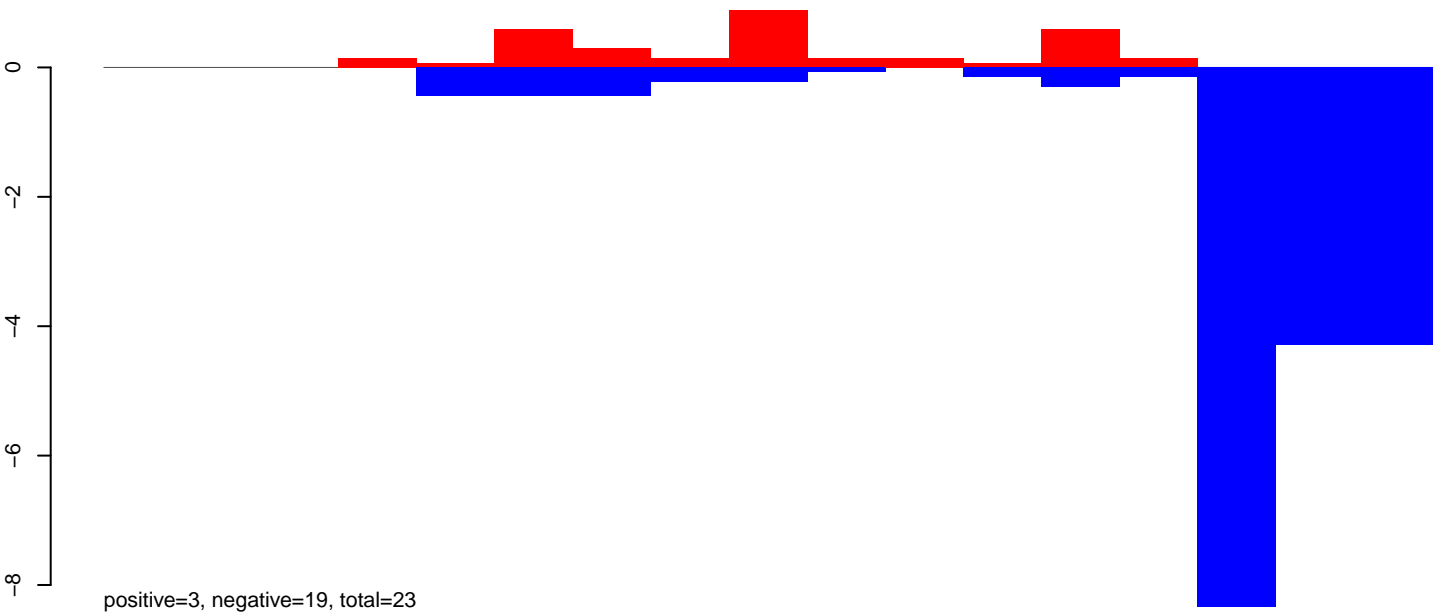
CuQuin_Testes_NL.rep



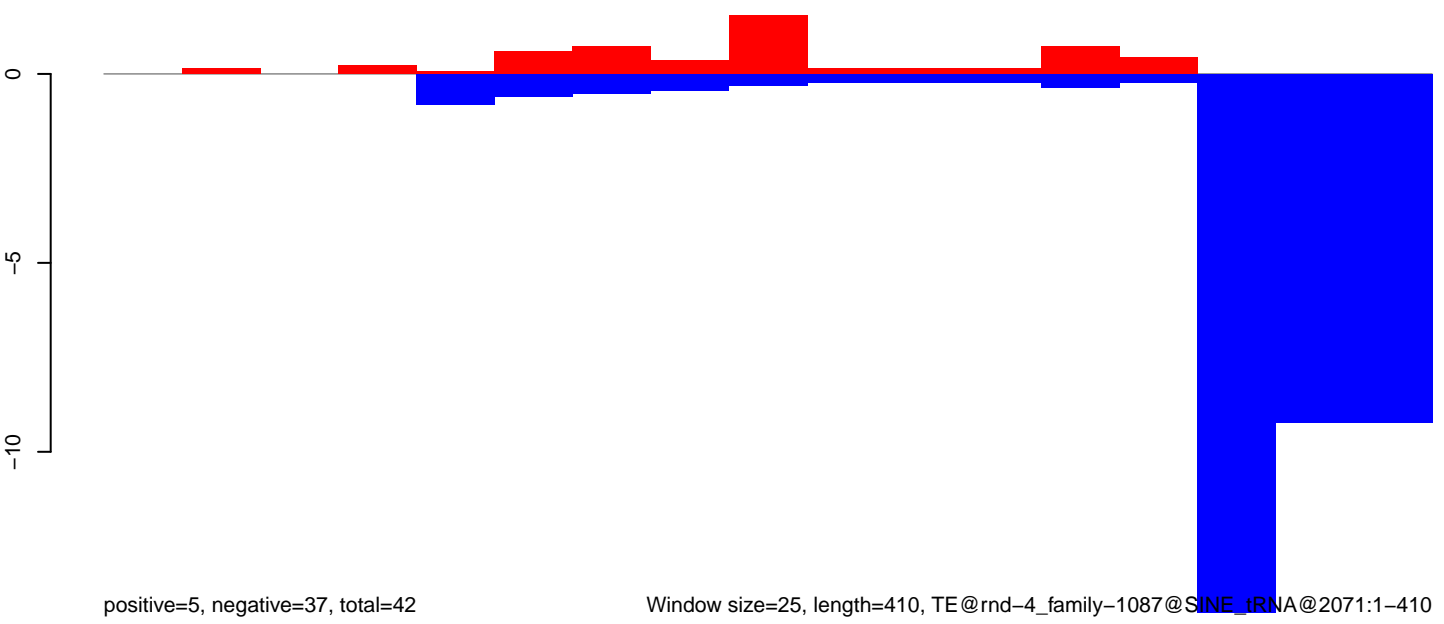
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



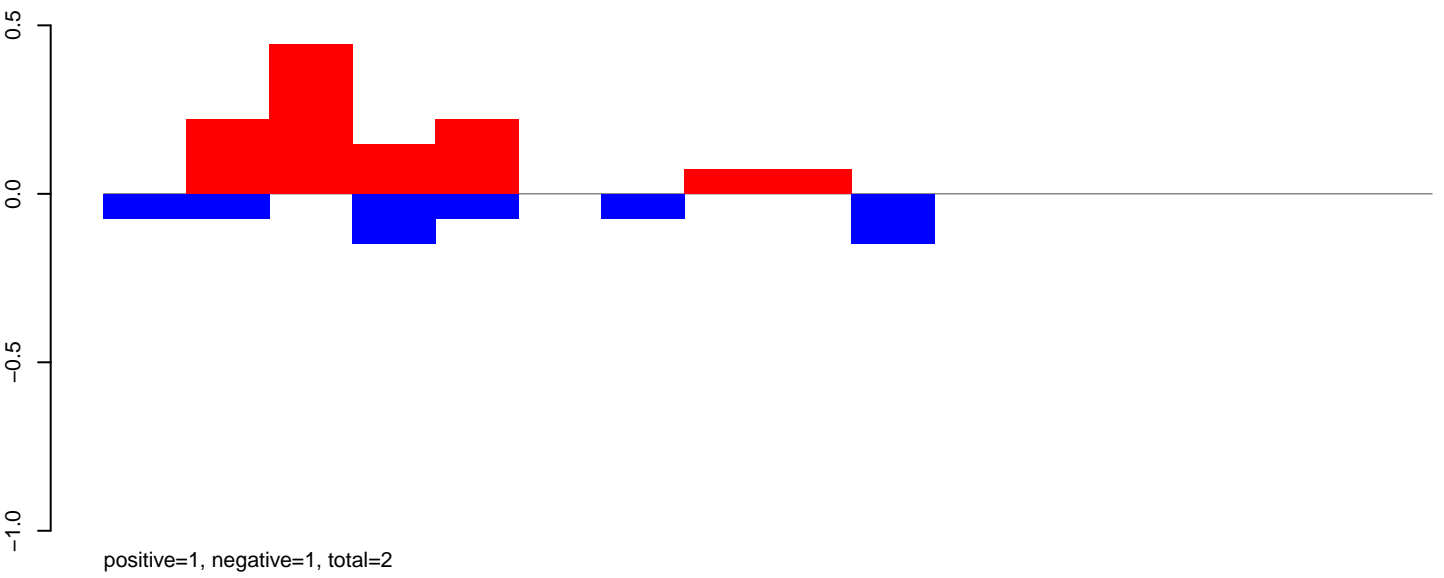
CuQuin_Testes_NL.rep



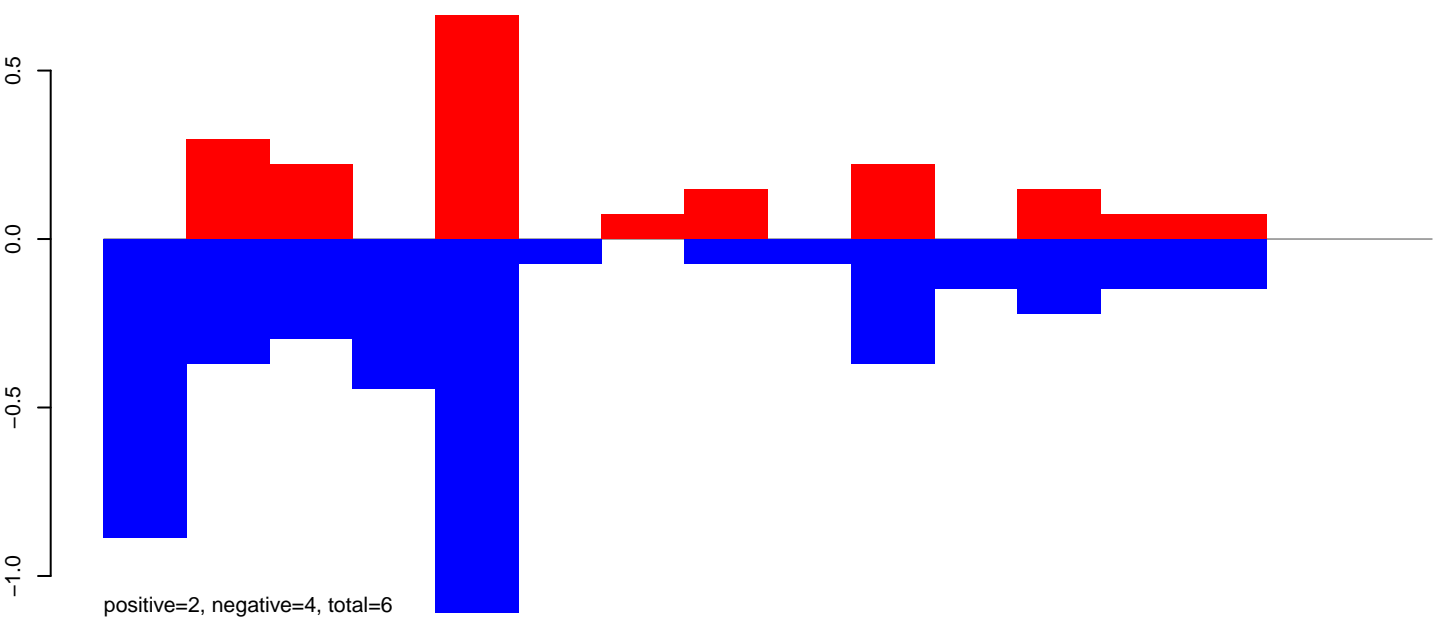
Window size=25, length=410, TE@rnd-4_family-1087@SINE_tRNA@2071:1-410

100 200 300 400

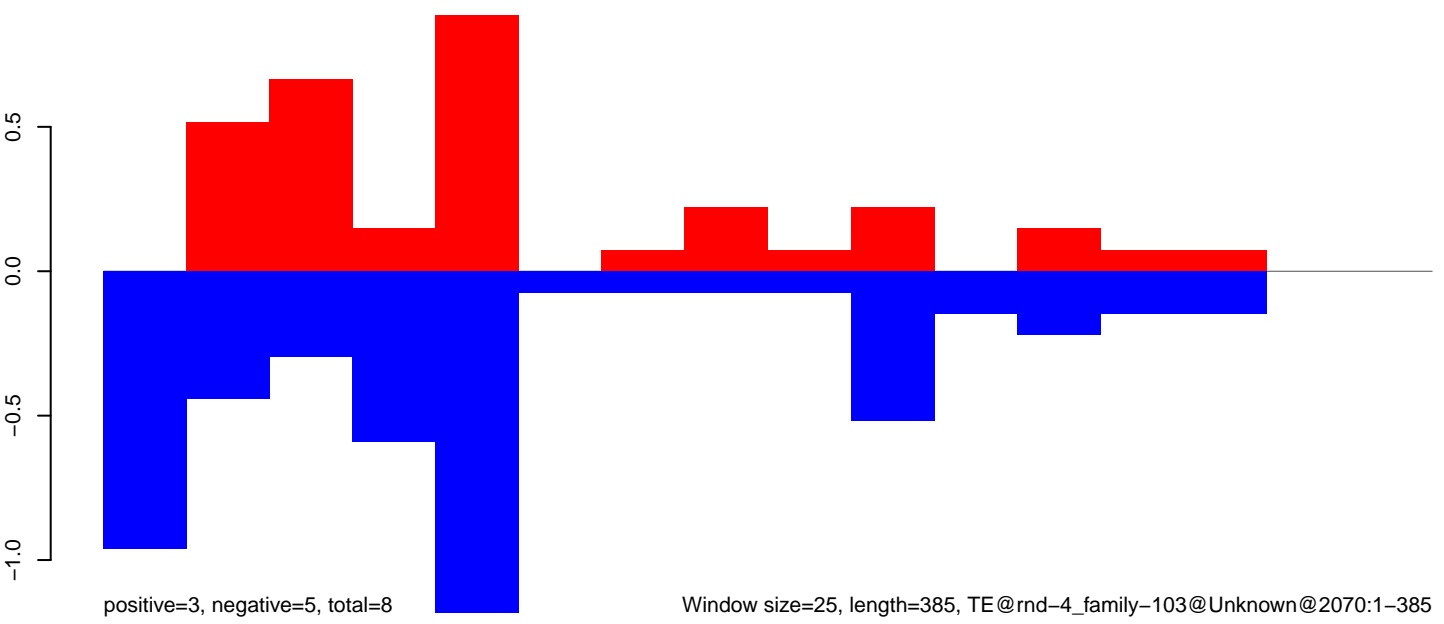
CuQuin_Testes_NL.18_23.rep



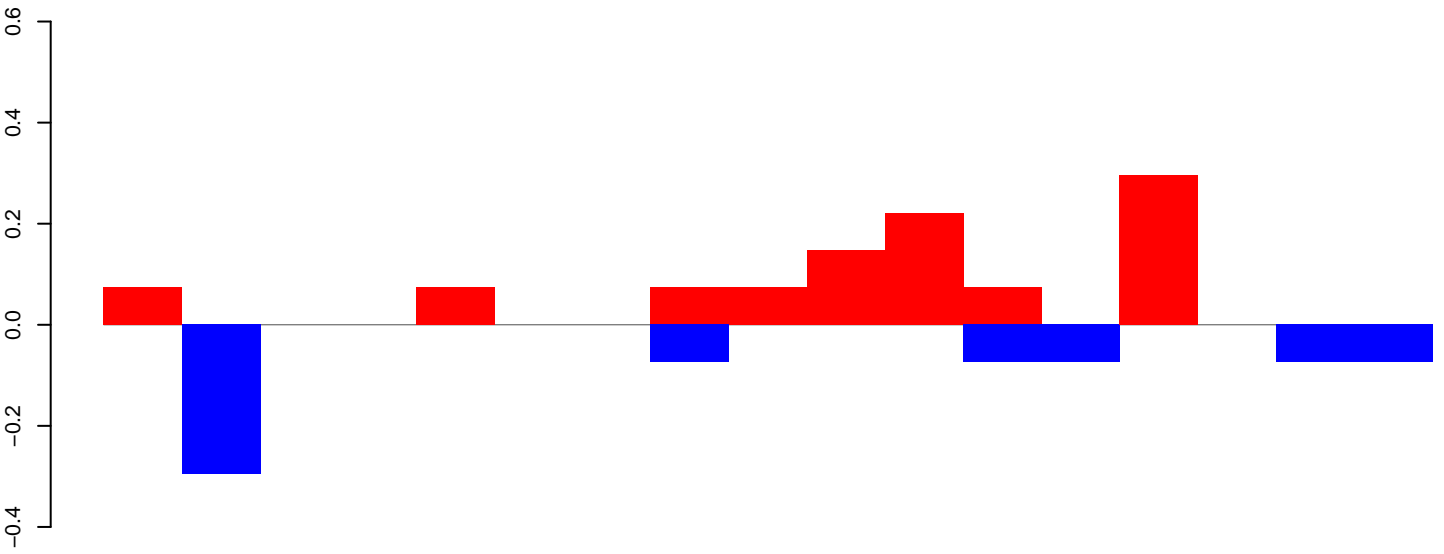
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

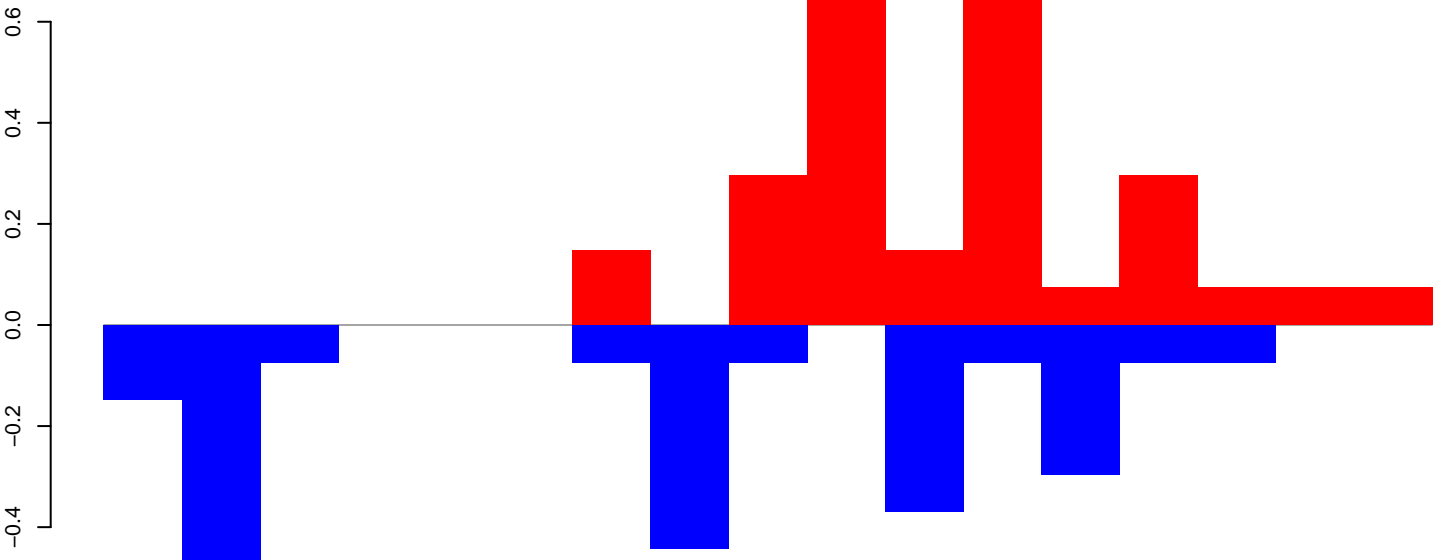


CuQuin_Testes_NL.18_23.rep



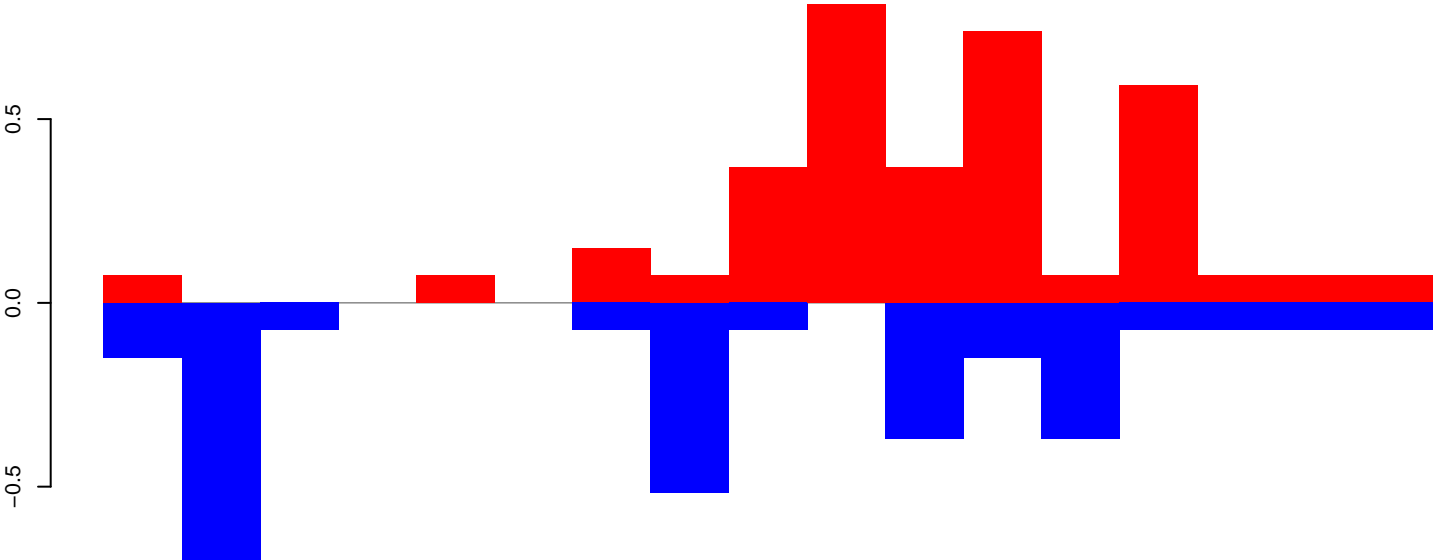
positive=1, negative=1, total=2

CuQuin_Testes_NL.24_35.rep



positive=3, negative=2, total=5

CuQuin_Testes_NL.rep

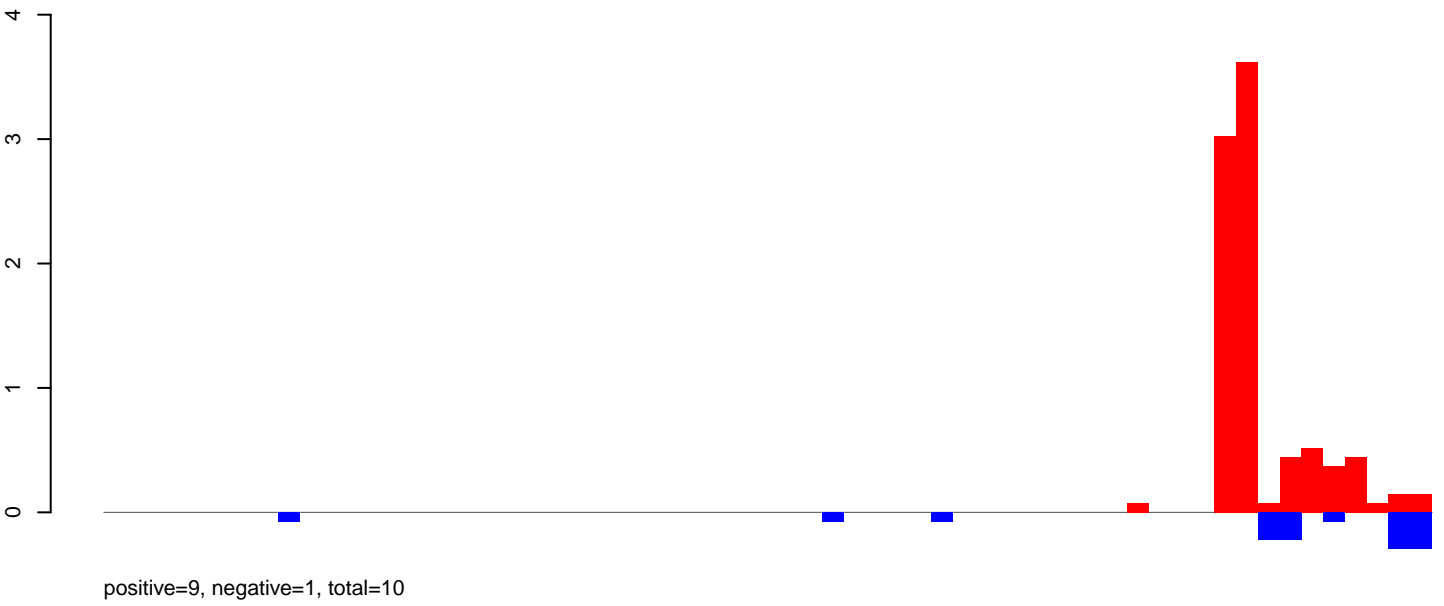


positive=4, negative=3, total=6

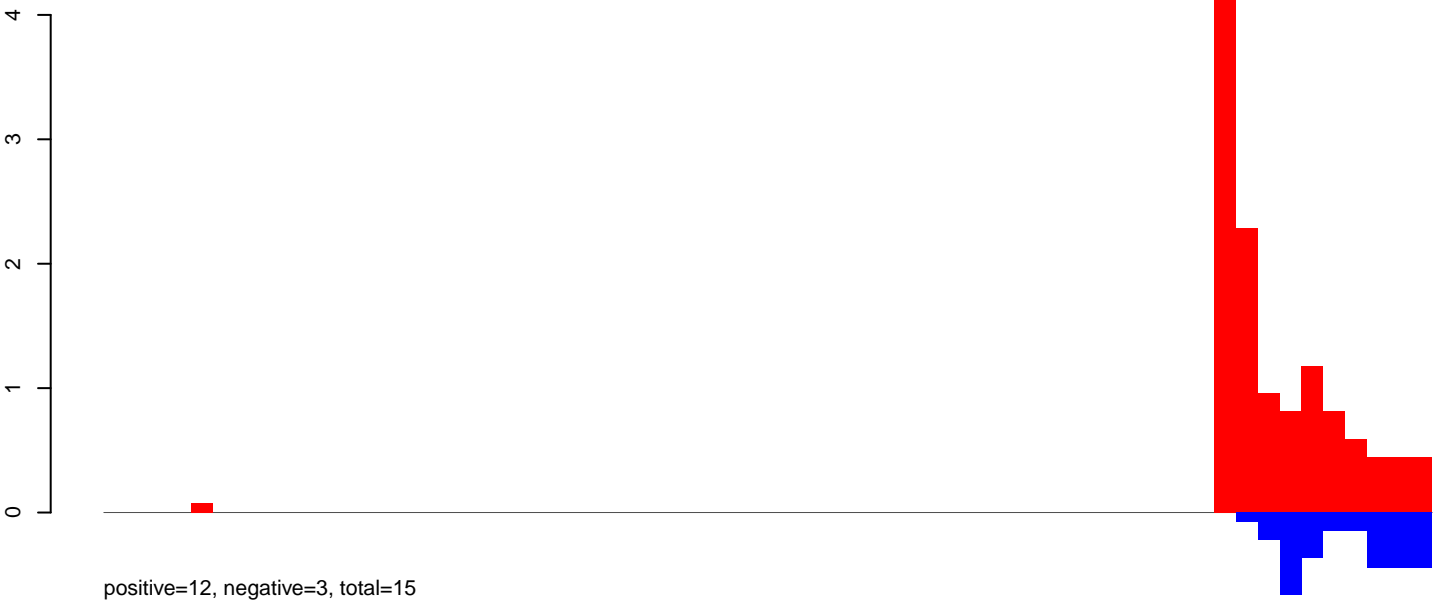
Window size=25, length=402, TE@rnd-3_family-708@LTR_Pao@139:1-402

100 200 300 400

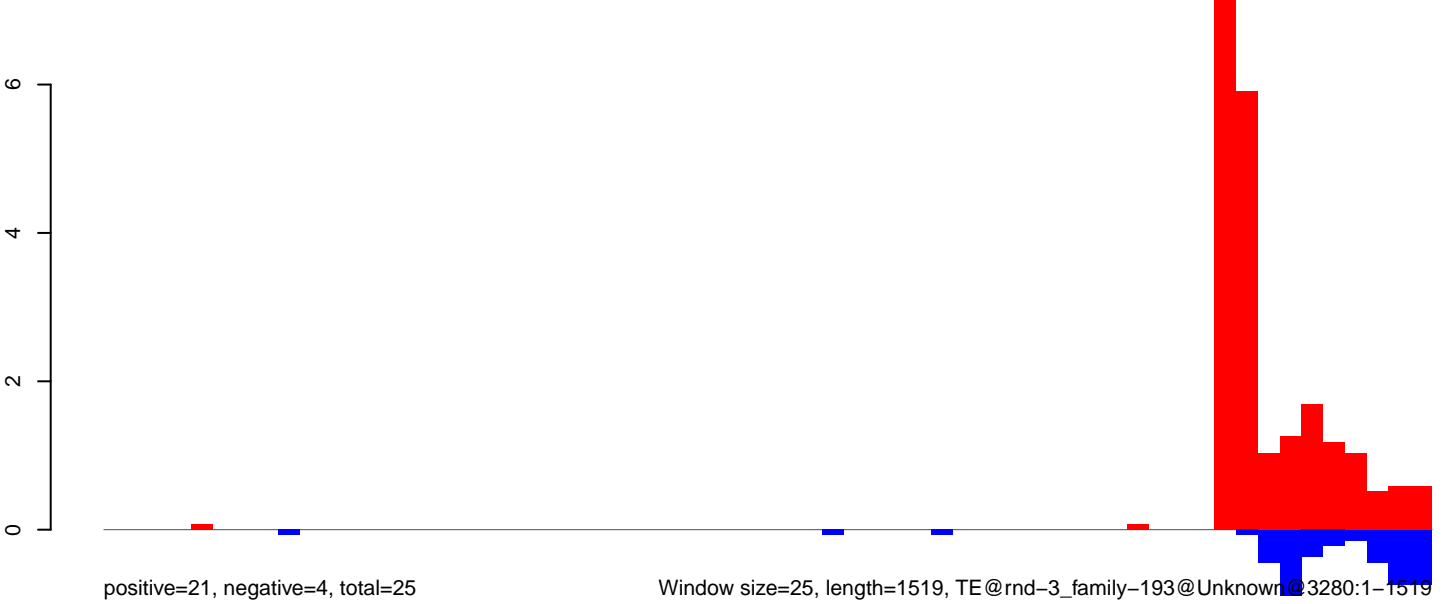
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

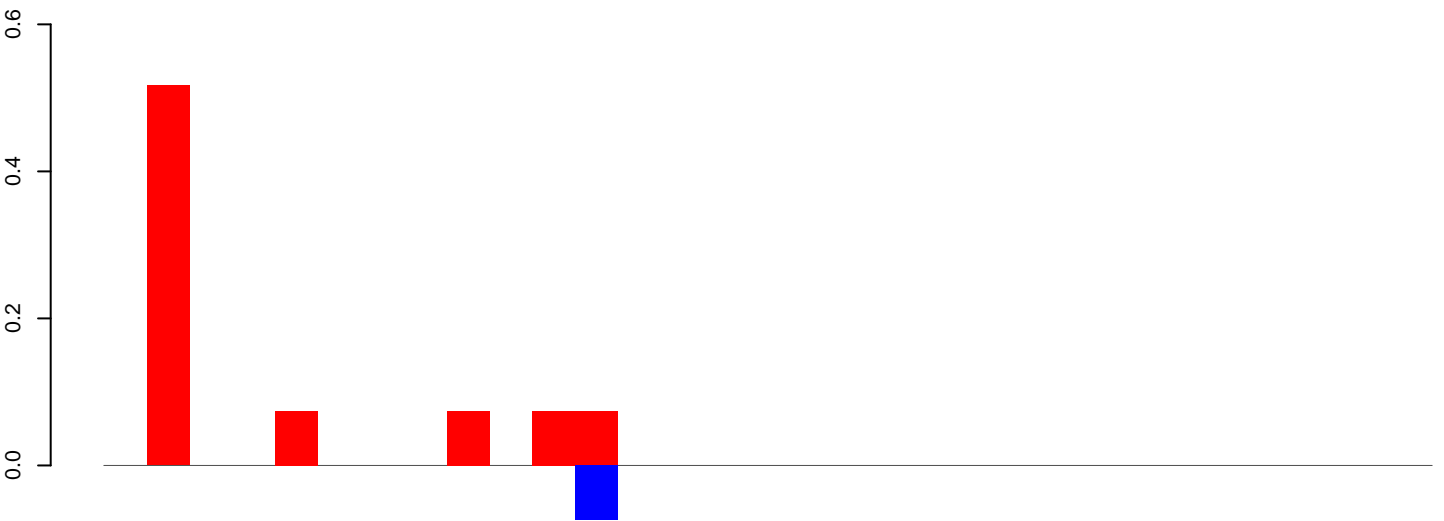


CuQuin_Testes_NL.rep



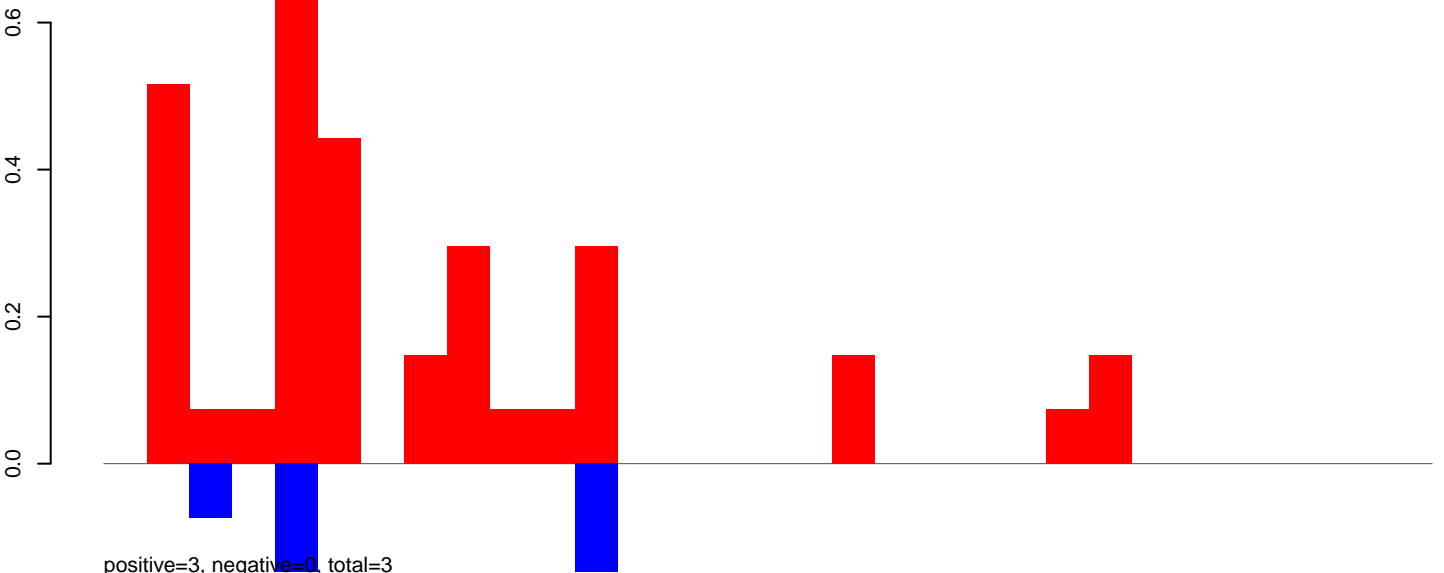
Window size=25, length=1519, TE@rnd-3_family-193@Unknown@3280:1-1519

CuQuin_Testes_NL.18_23.rep



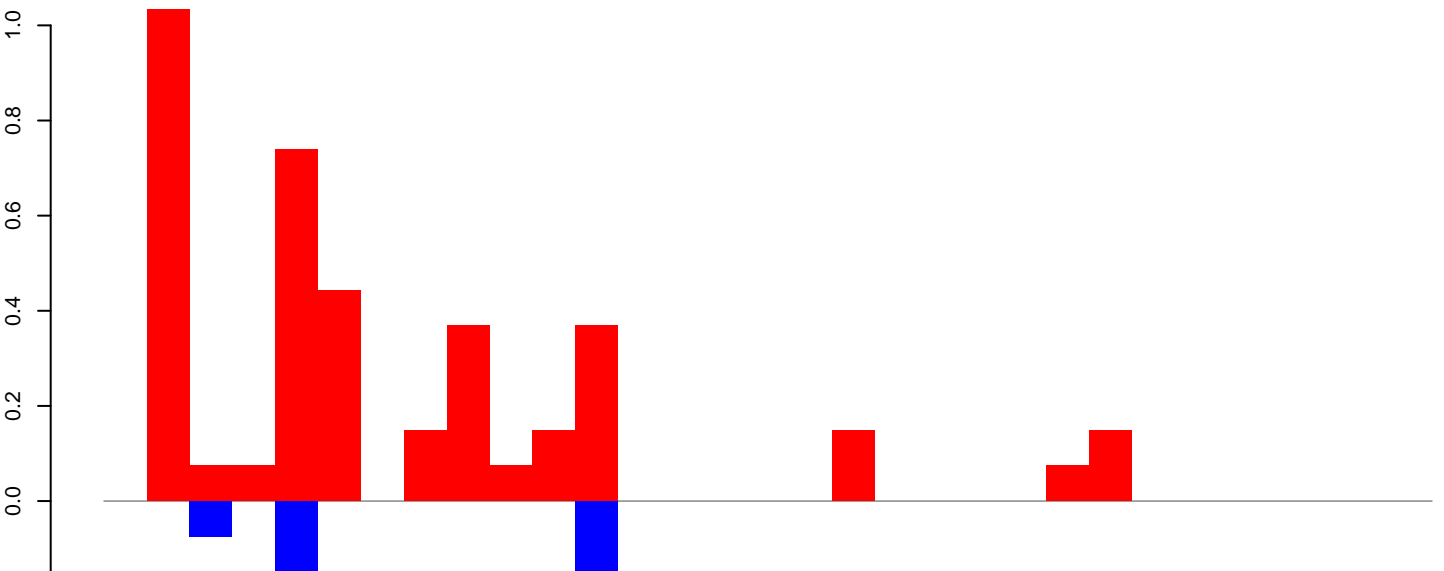
positive=1, negative=0, total=1

CuQuin_Testes_NL.24_35.rep



positive=3, negative=0, total=3

CuQuin_Testes_NL.rep

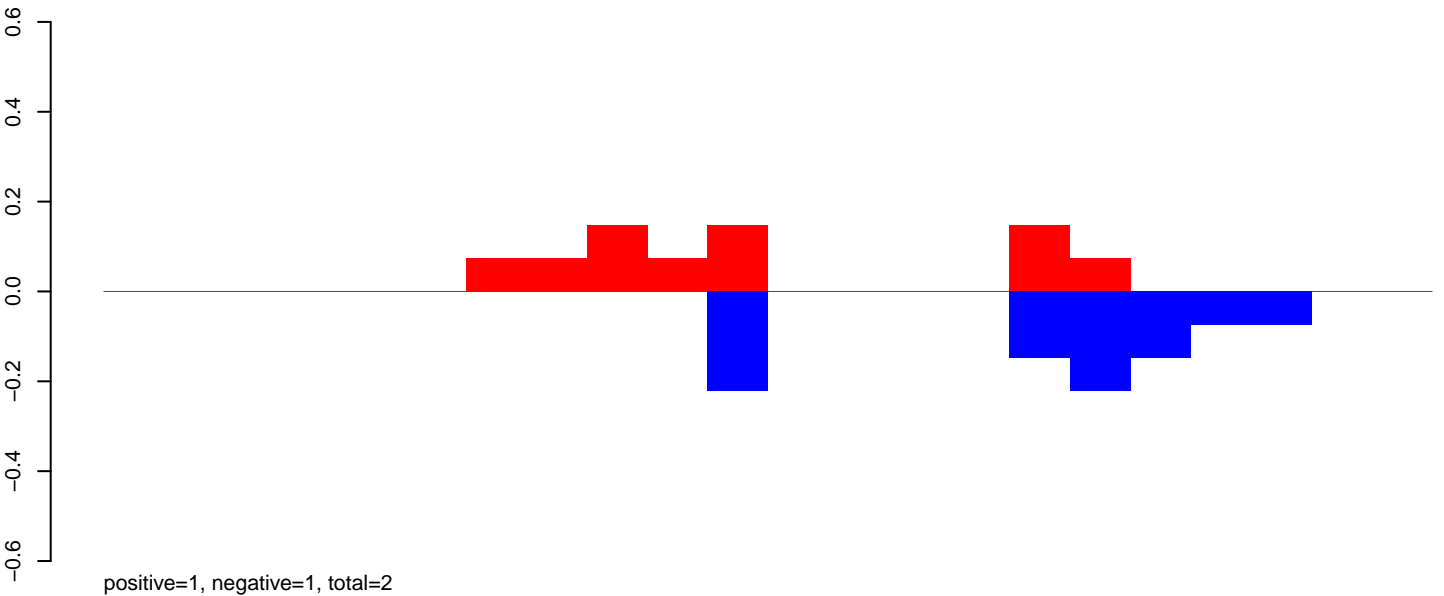


positive=4, negative=0, total=4

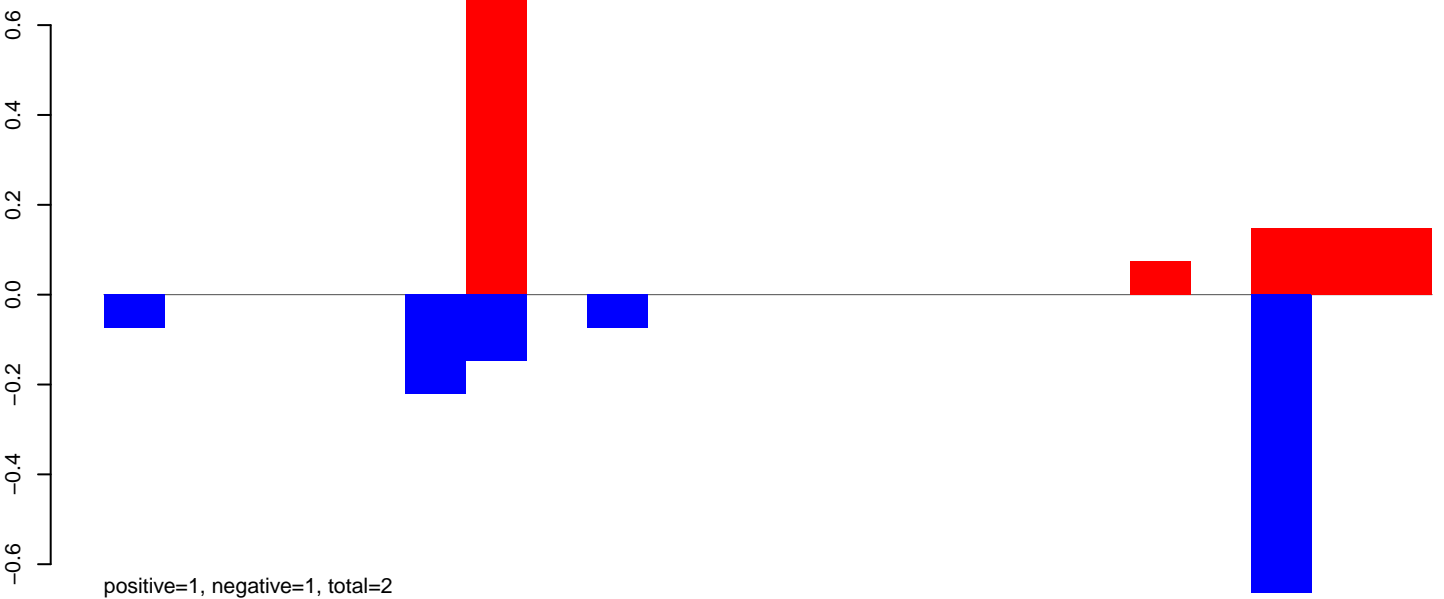
Window size=25, length=766, TE@rnd-5_family-976@Unknown@1849:1-766

0 200 400 600 800

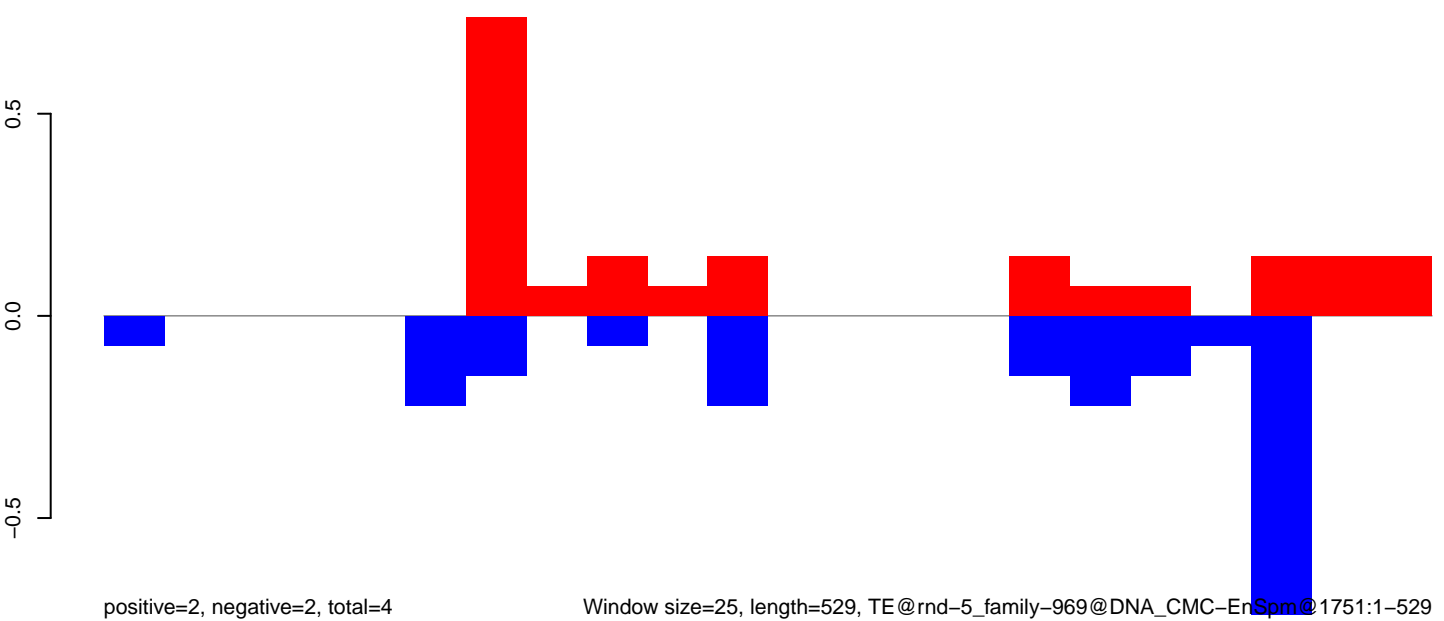
CuQuin_Testes_NL.18_23.rep



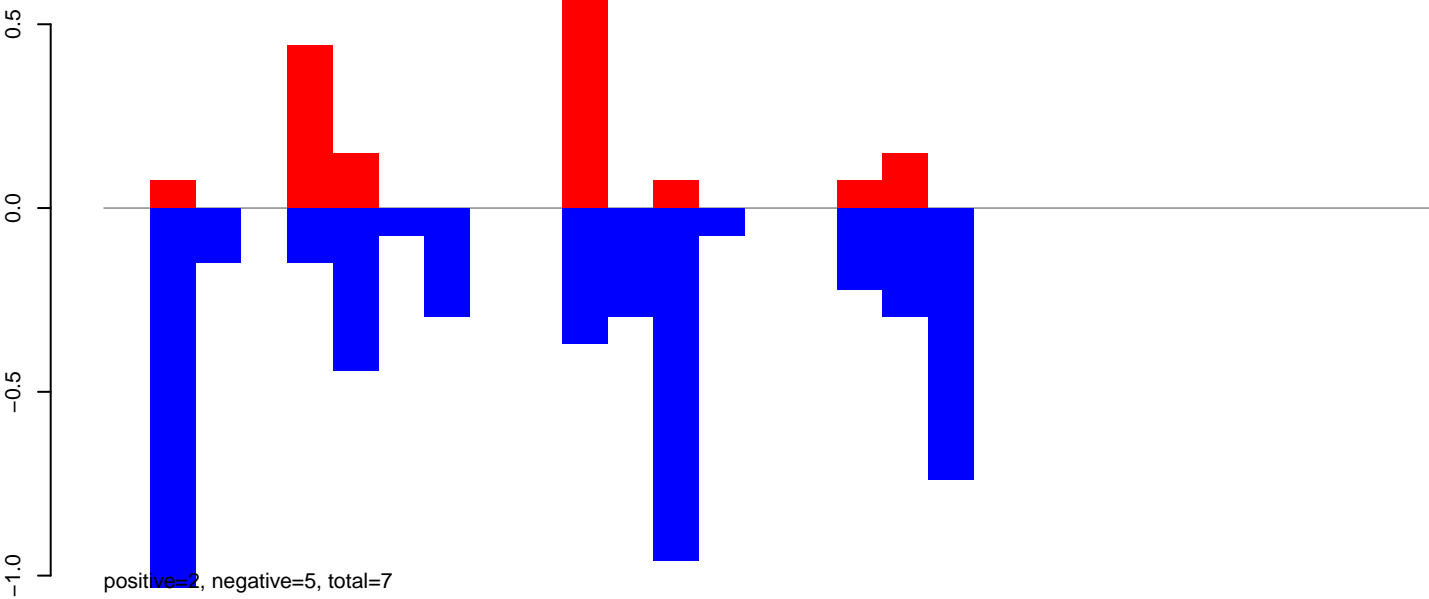
CuQuin_Testes_NL.24_35.rep



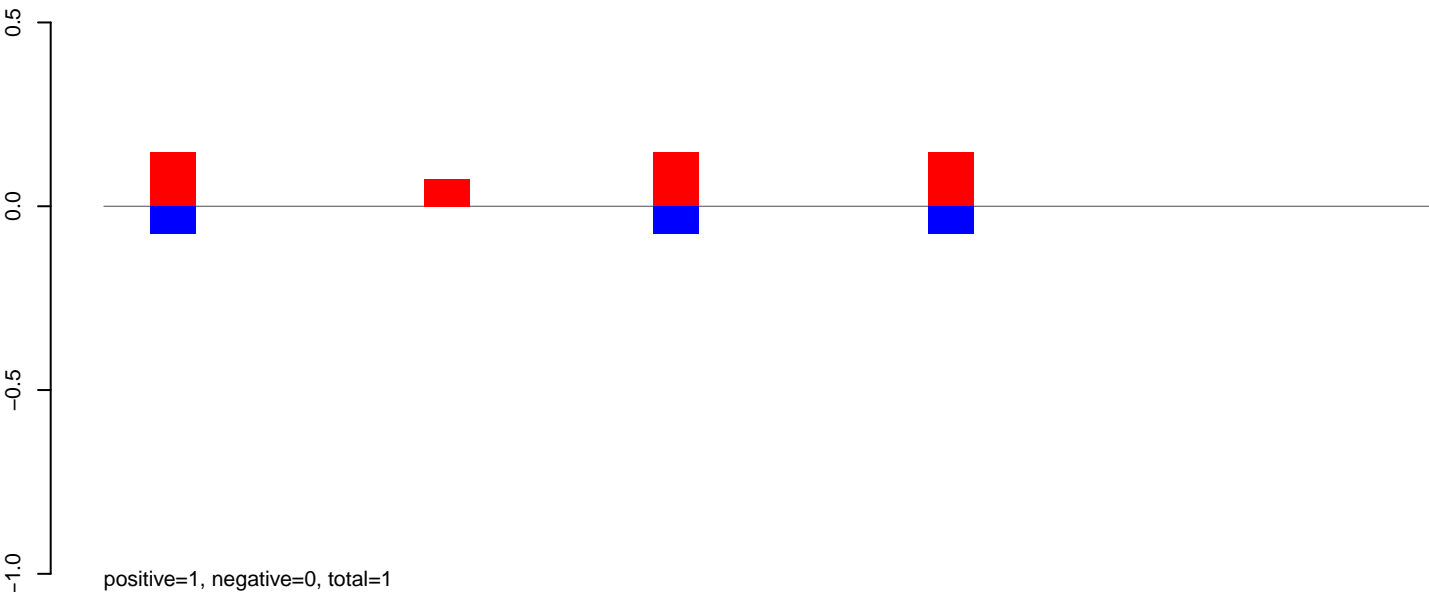
CuQuin_Testes_NL.rep



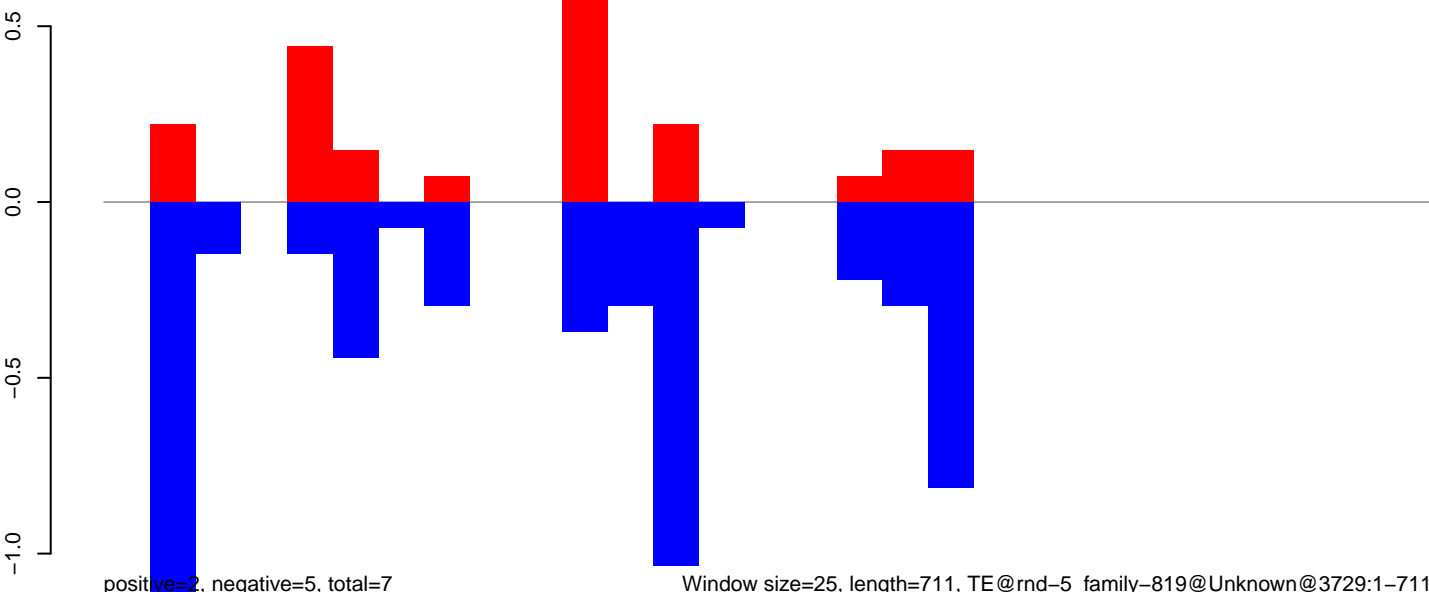
CuQuin_Testes_NL.18_23.rep



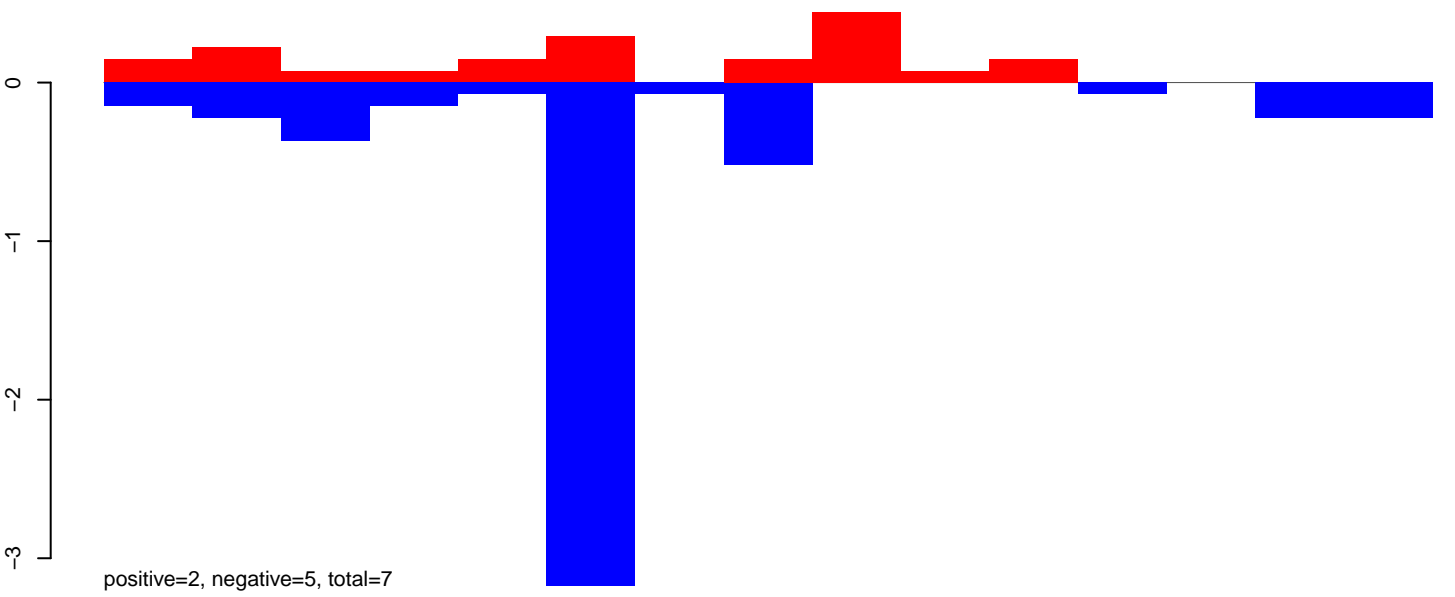
CuQuin_Testes_NL.24_35.rep



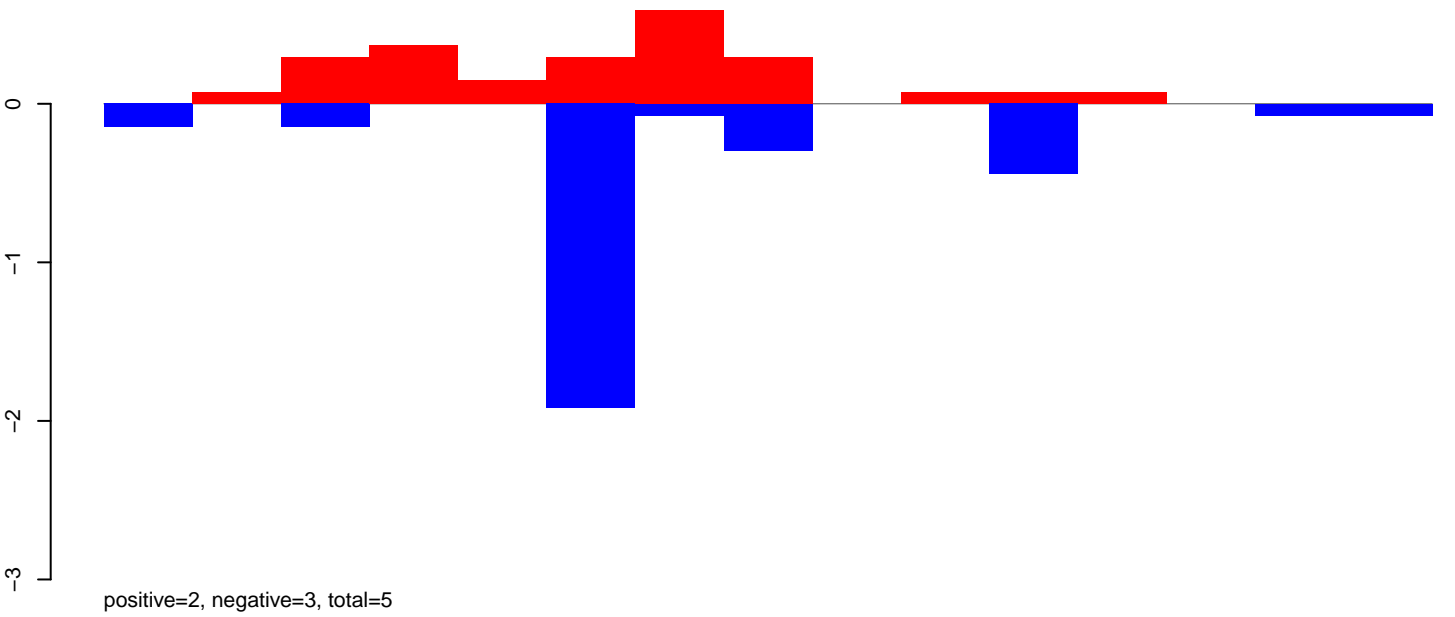
CuQuin_Testes_NL.rep



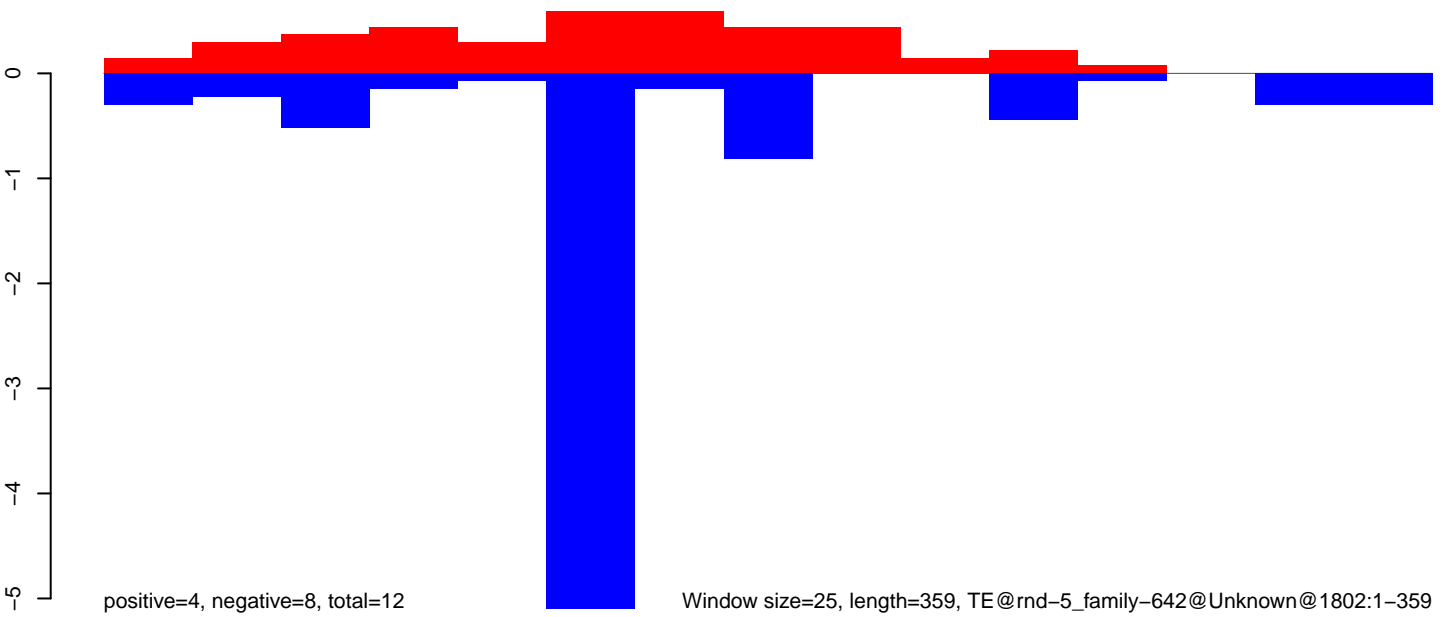
CuQuin_Testes_NL.18_23.rep



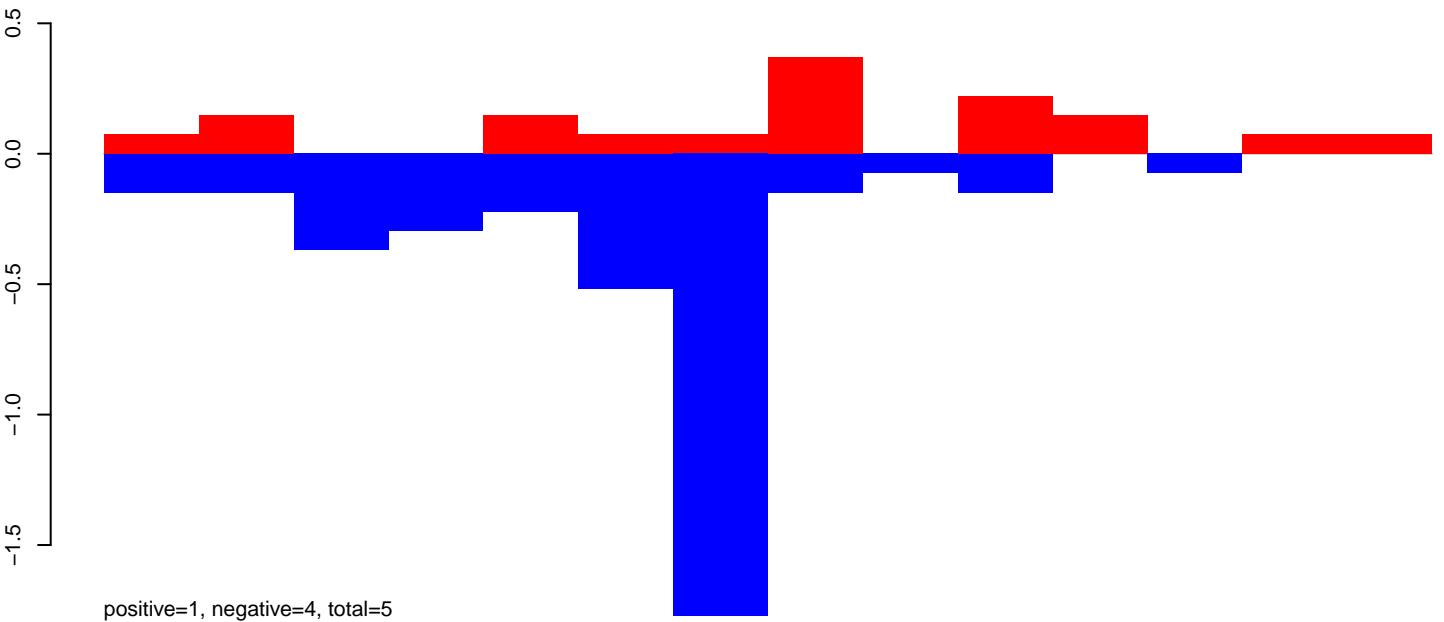
CuQuin_Testes_NL.24_35.rep



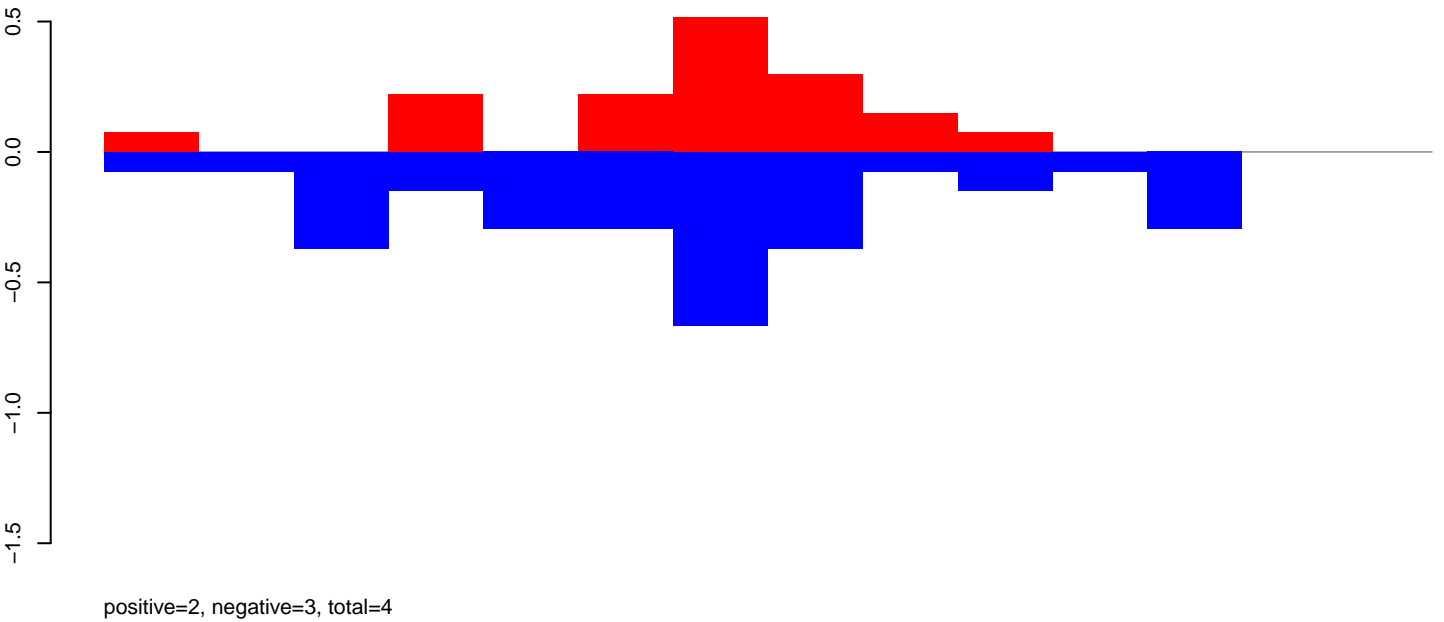
CuQuin_Testes_NL.rep



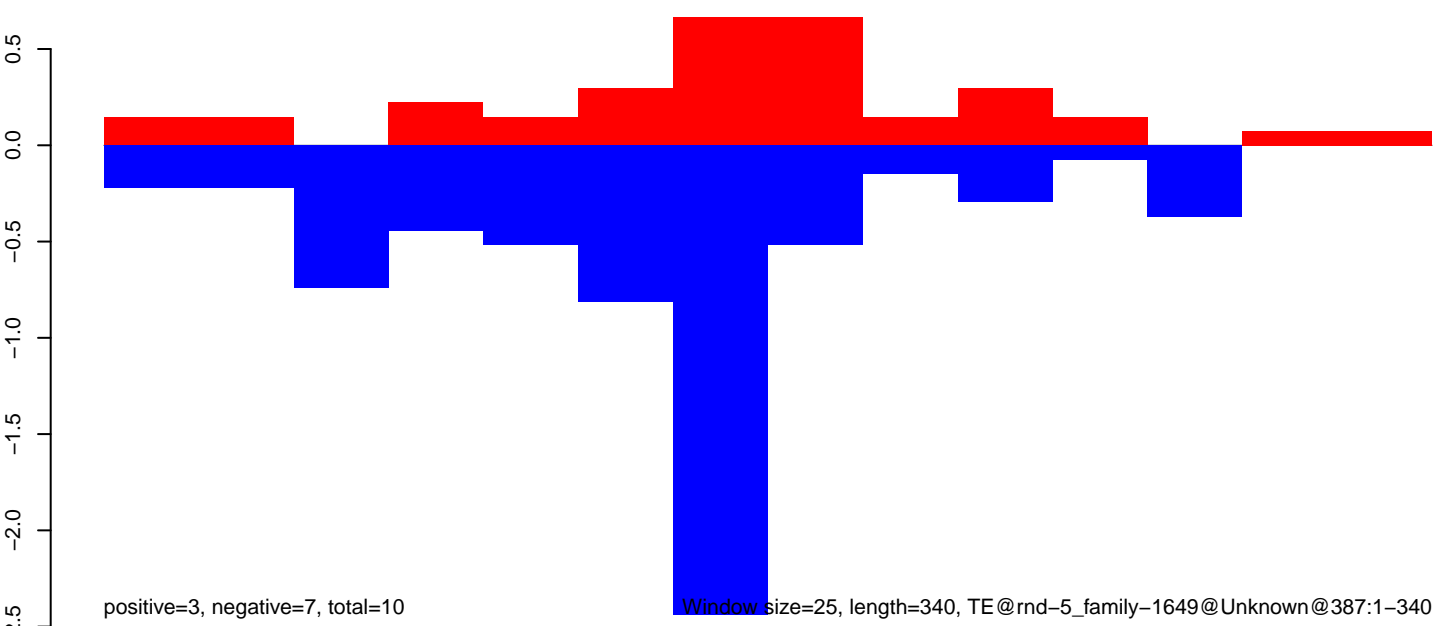
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

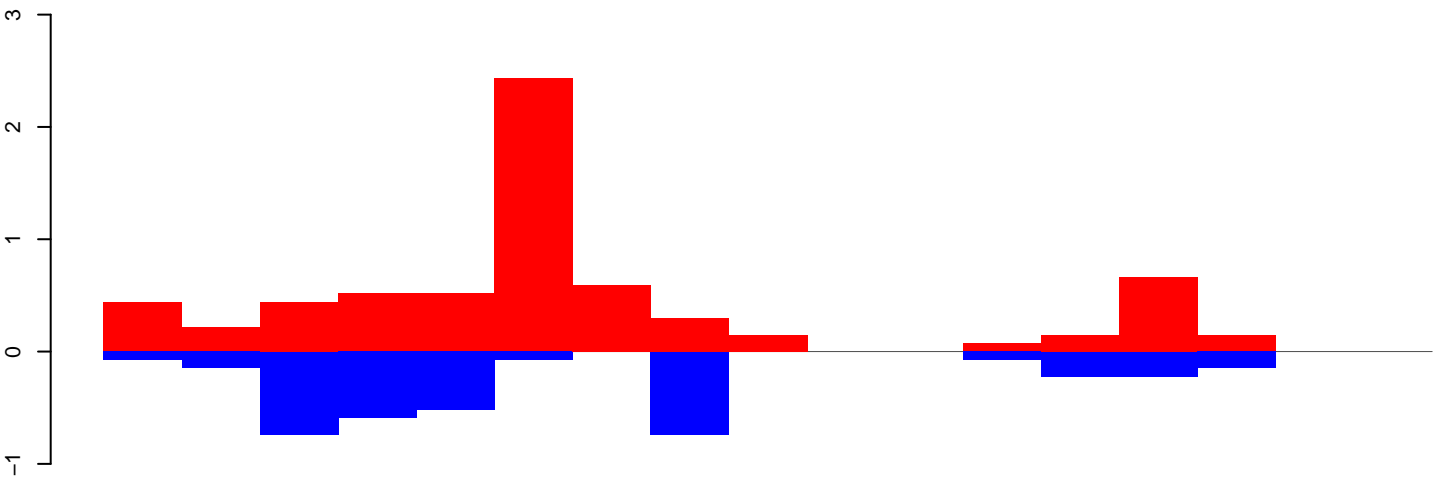


CuQuin_Testes_NL.rep



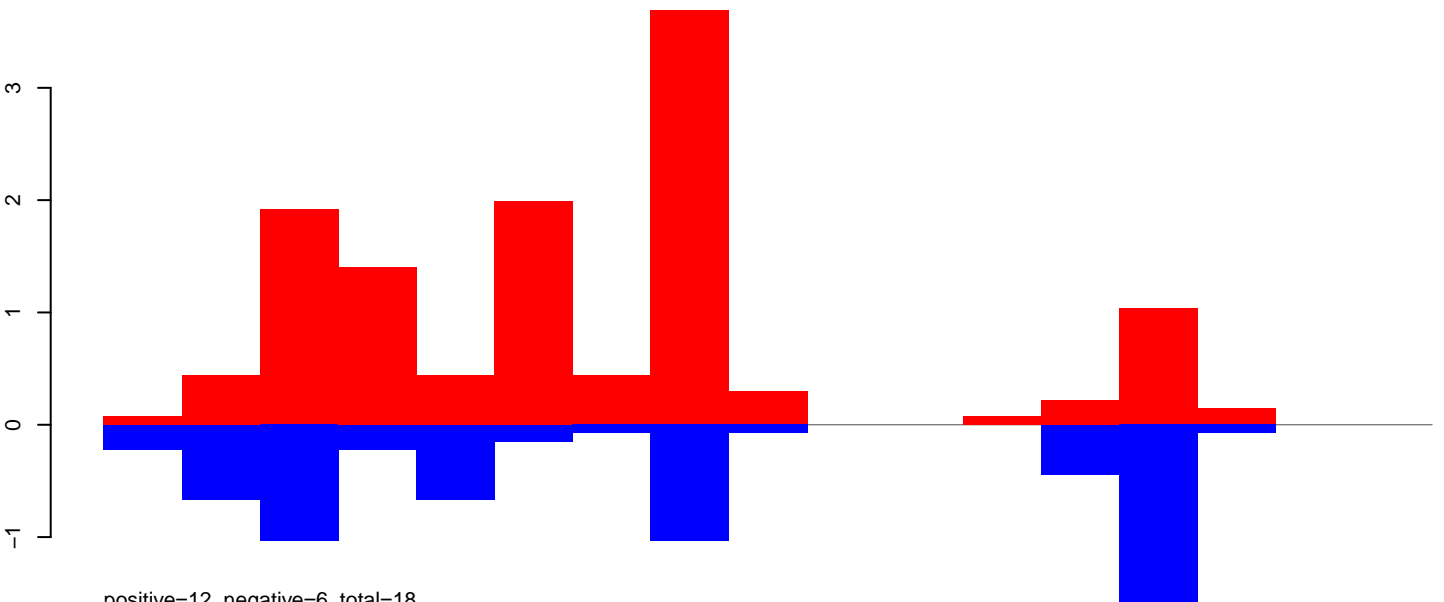
Window size=25, length=340, TE@rnd-5_family-1649@Unknown@387:1-340

CuQuin_Testes_NL.18_23.rep



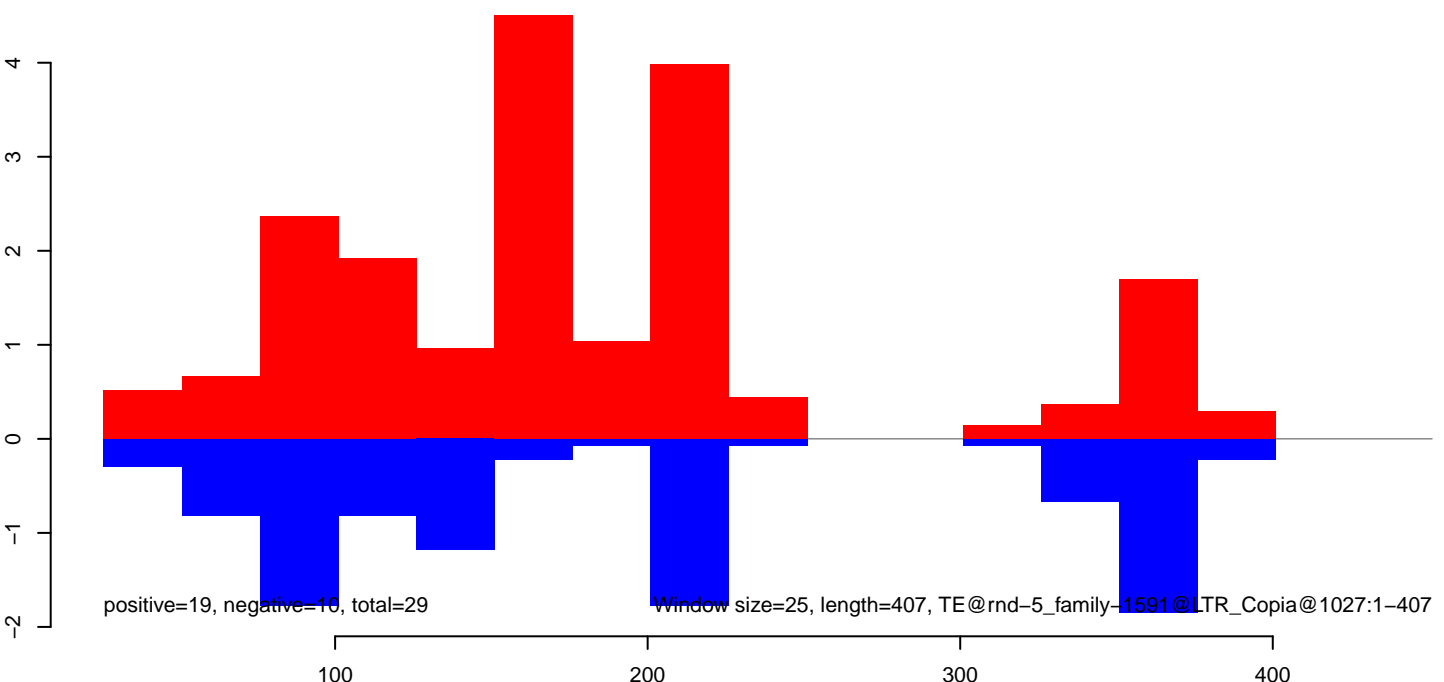
positive=7, negative=4, total=10

CuQuin_Testes_NL.24_35.rep



positive=12, negative=6, total=18

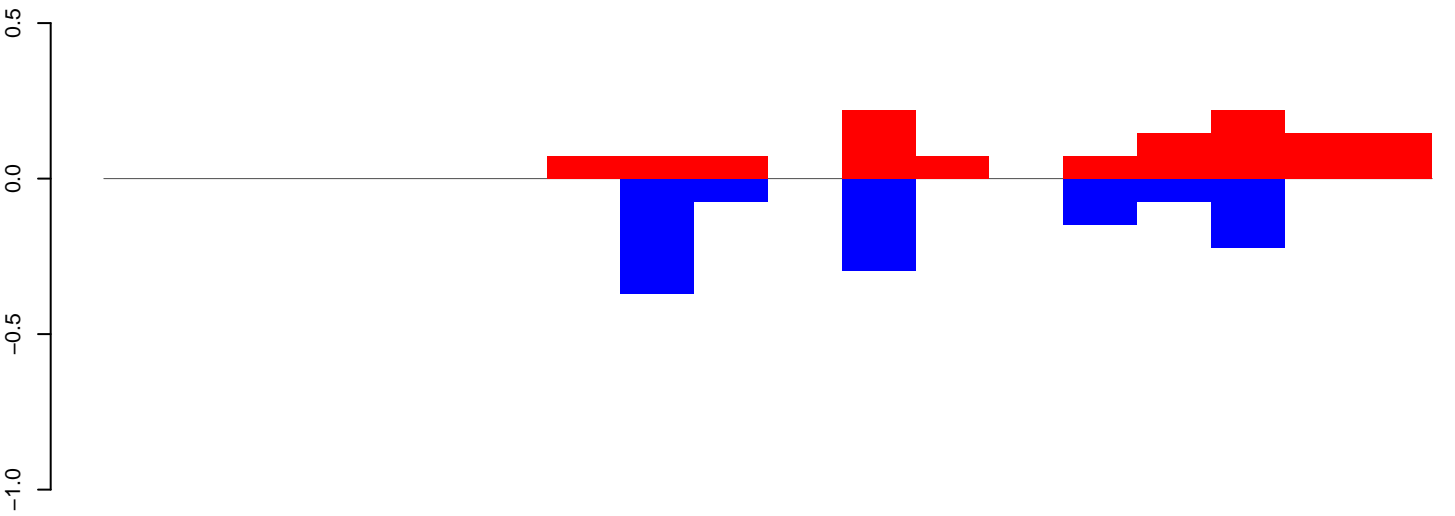
CuQuin_Testes_NL.rep



positive=19, negative=10, total=29

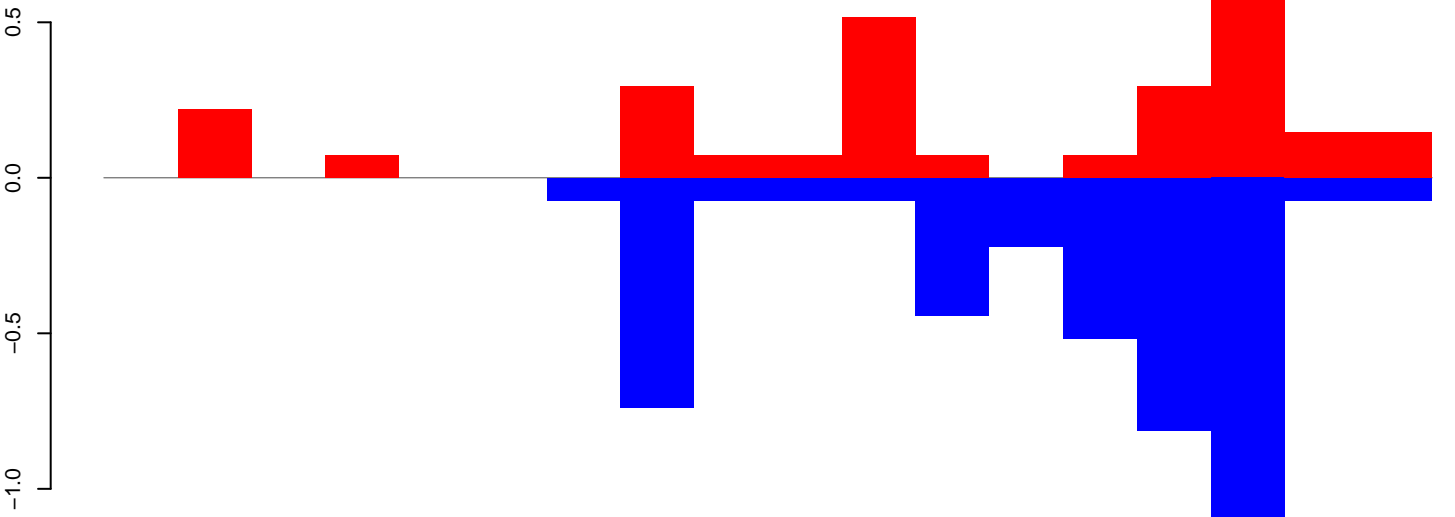
Window size=25, length=407, TE@rnd-5_family-1591@LTR_Copia@1027:1-407

CuQuin_Testes_NL.18_23.rep



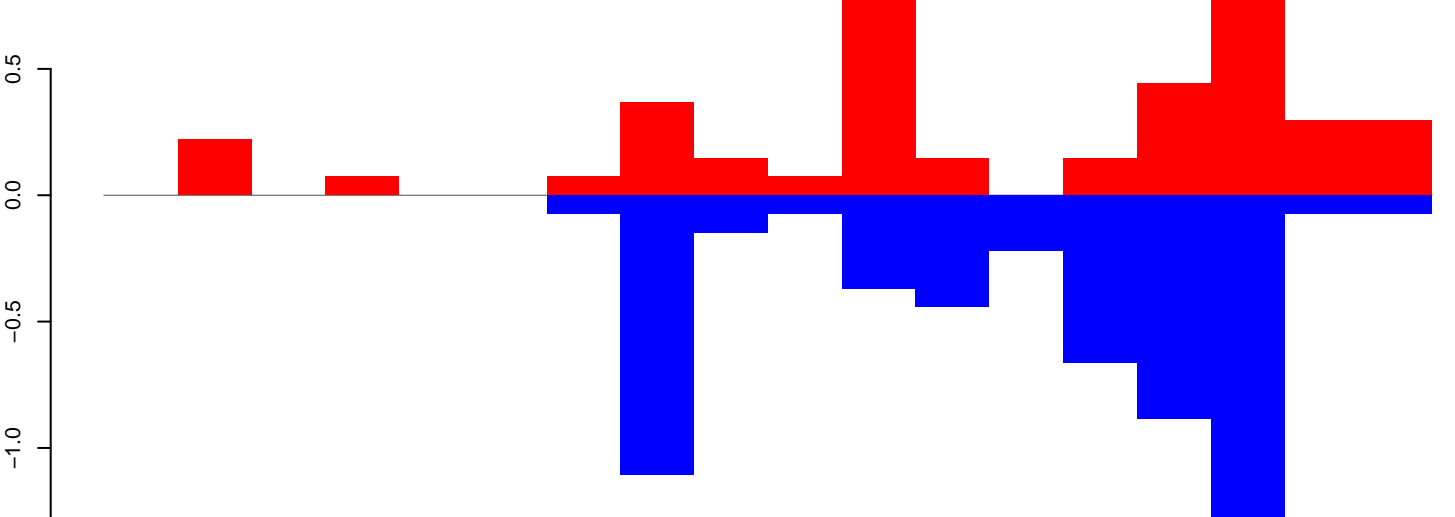
positive=1, negative=1, total=2

CuQuin_Testes_NL.24_35.rep



positive=3, negative=4, total=7

CuQuin_Testes_NL.rep



positive=4, negative=6, total=10

Window size=25, length=434, TE@rnd-5_family-1392@Unknown@1694:1-434

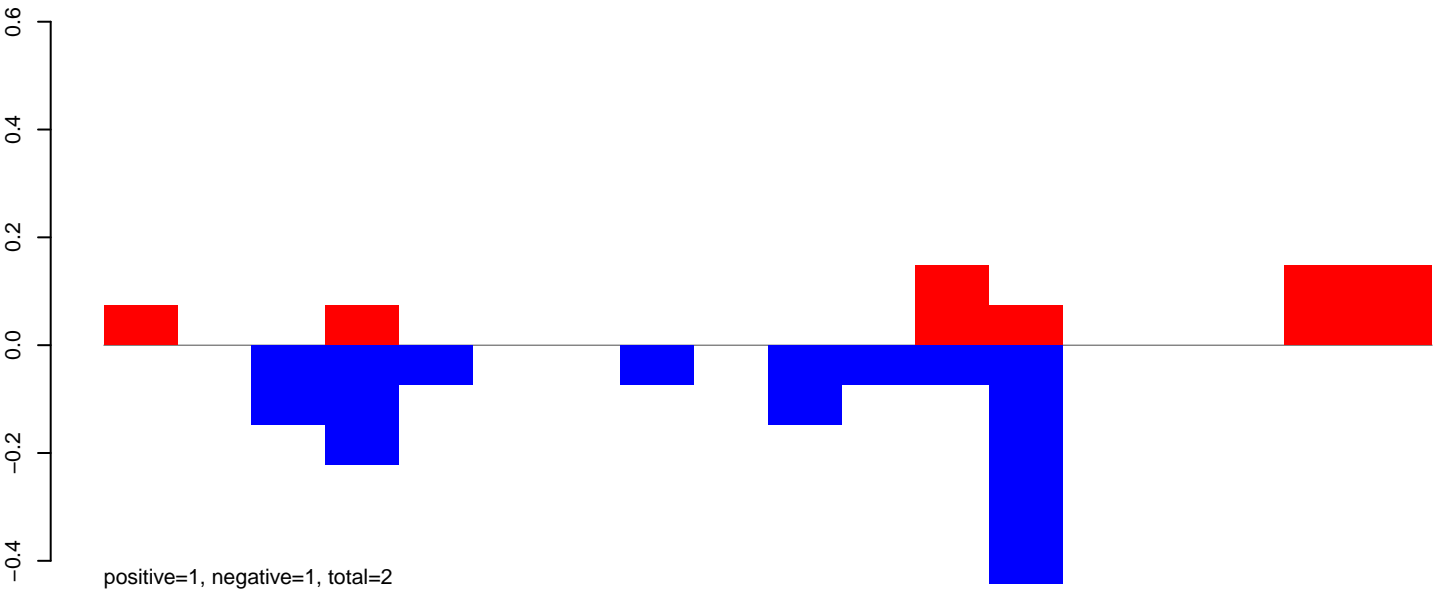
100

200

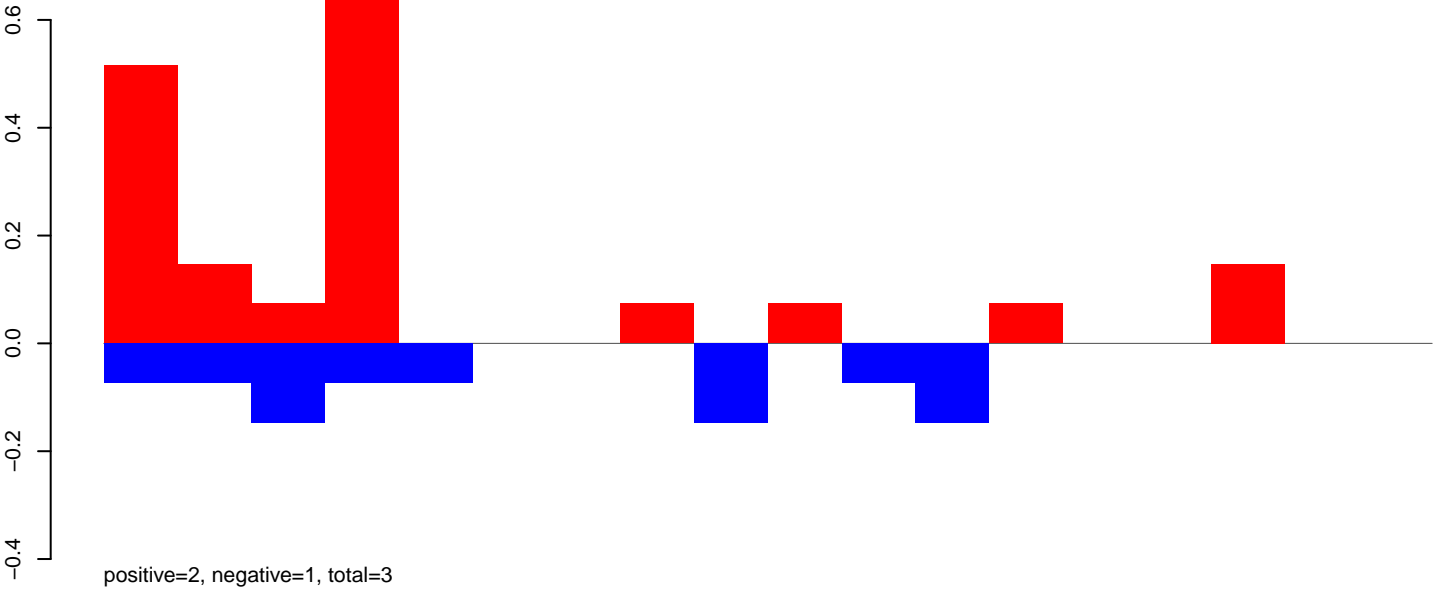
300

400

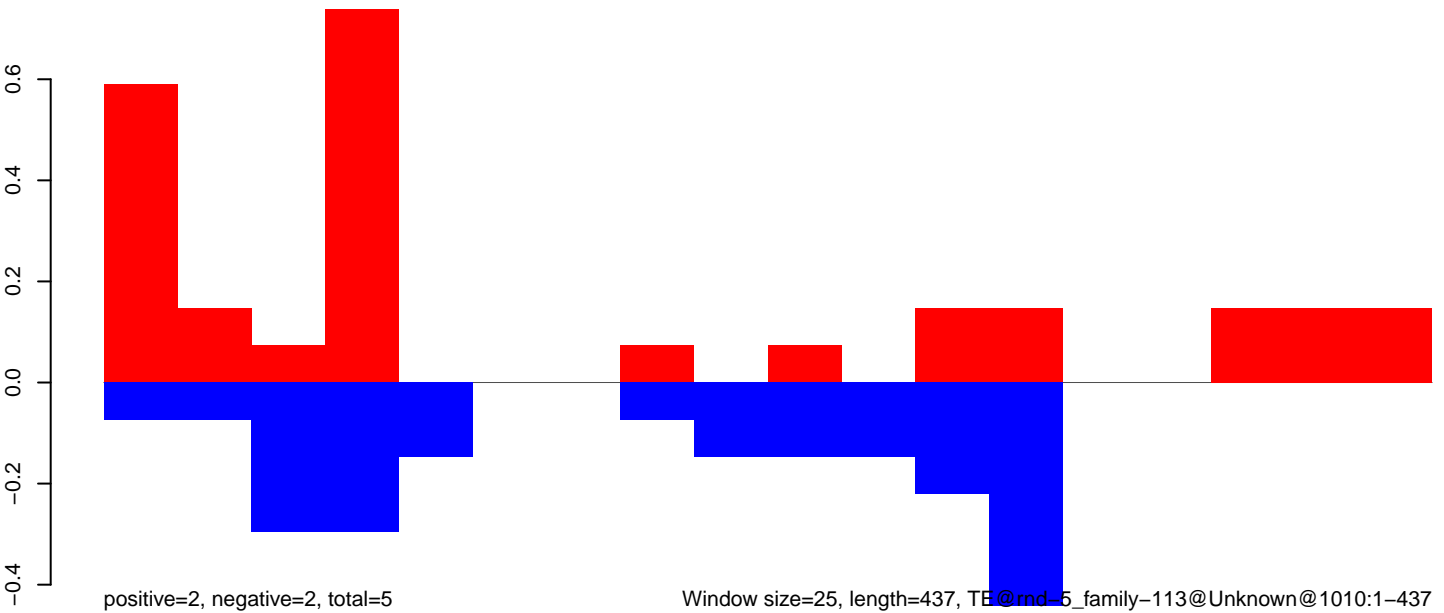
CuQuin_Testes_NL.18_23.rep



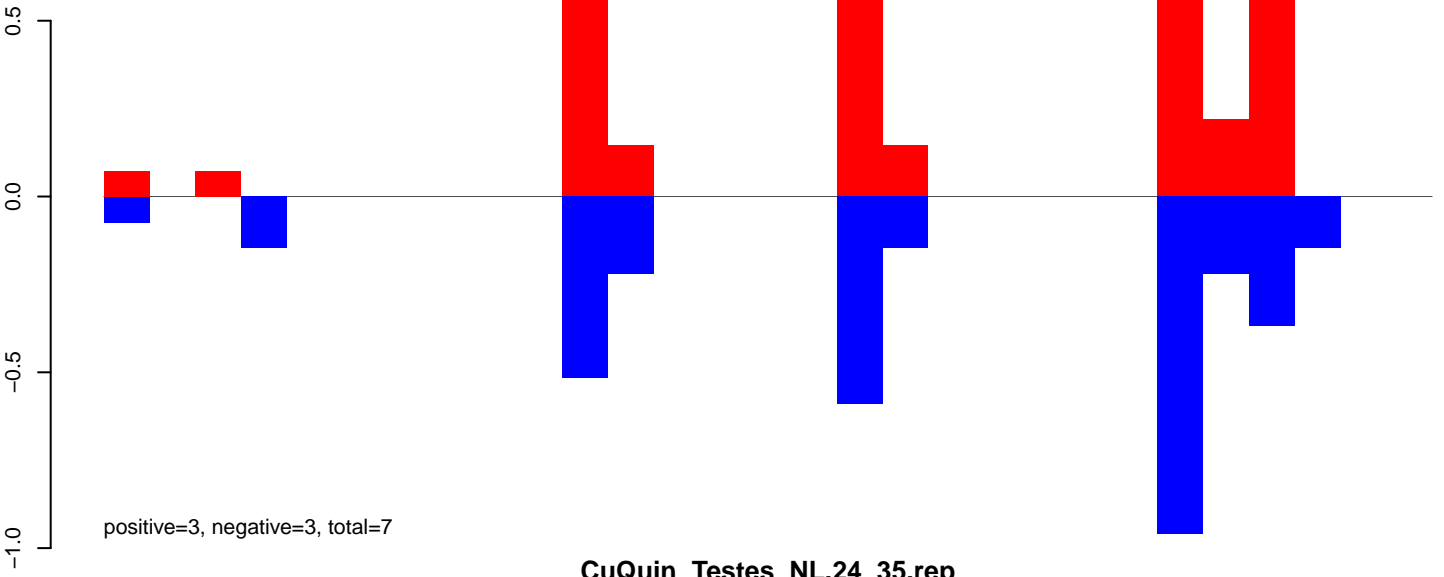
CuQuin_Testes_NL.24_35.rep



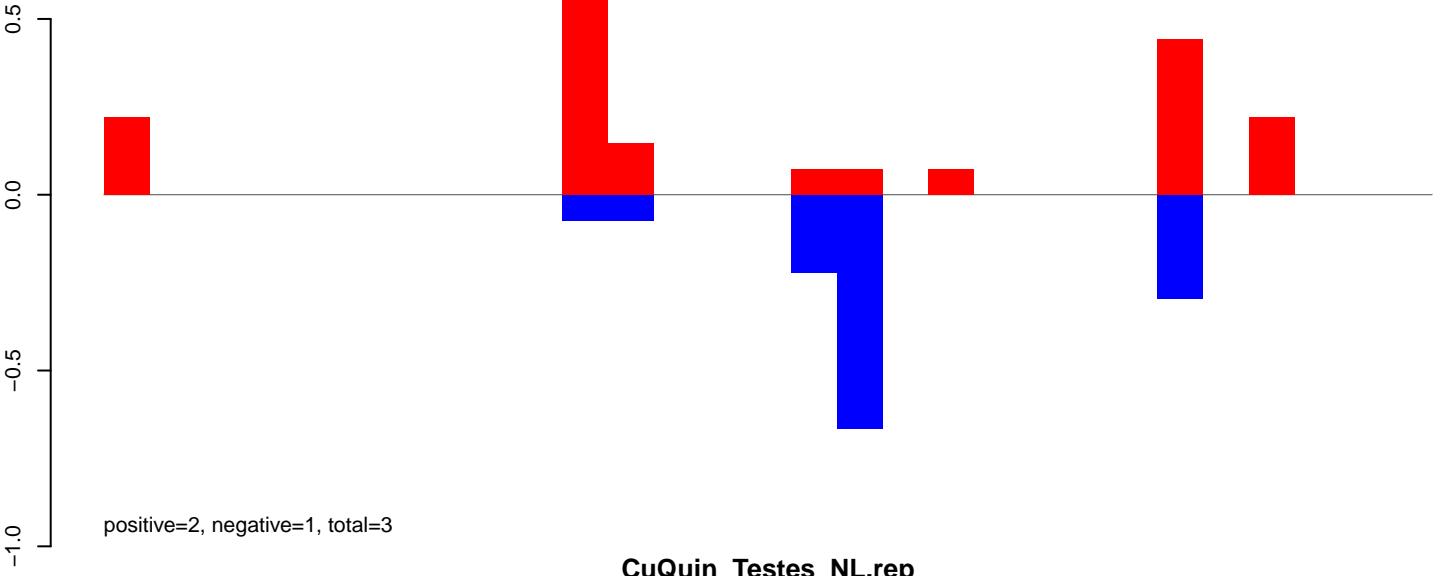
CuQuin_Testes_NL.rep



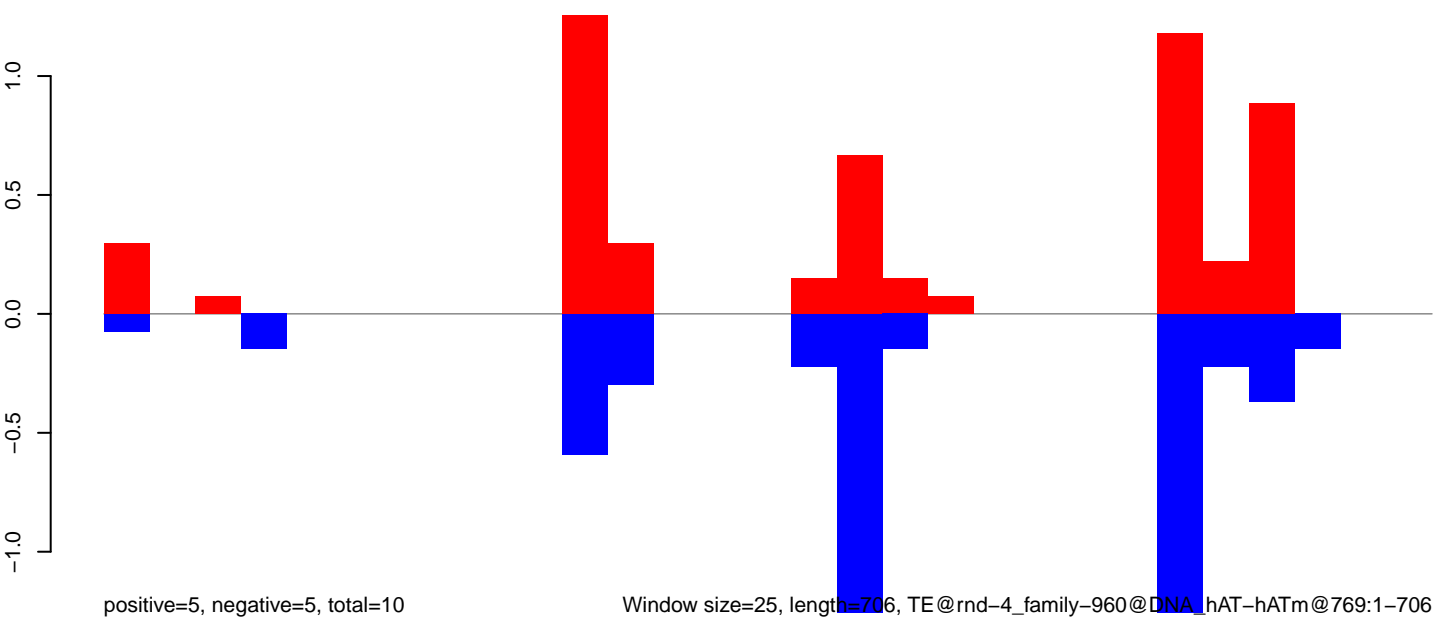
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

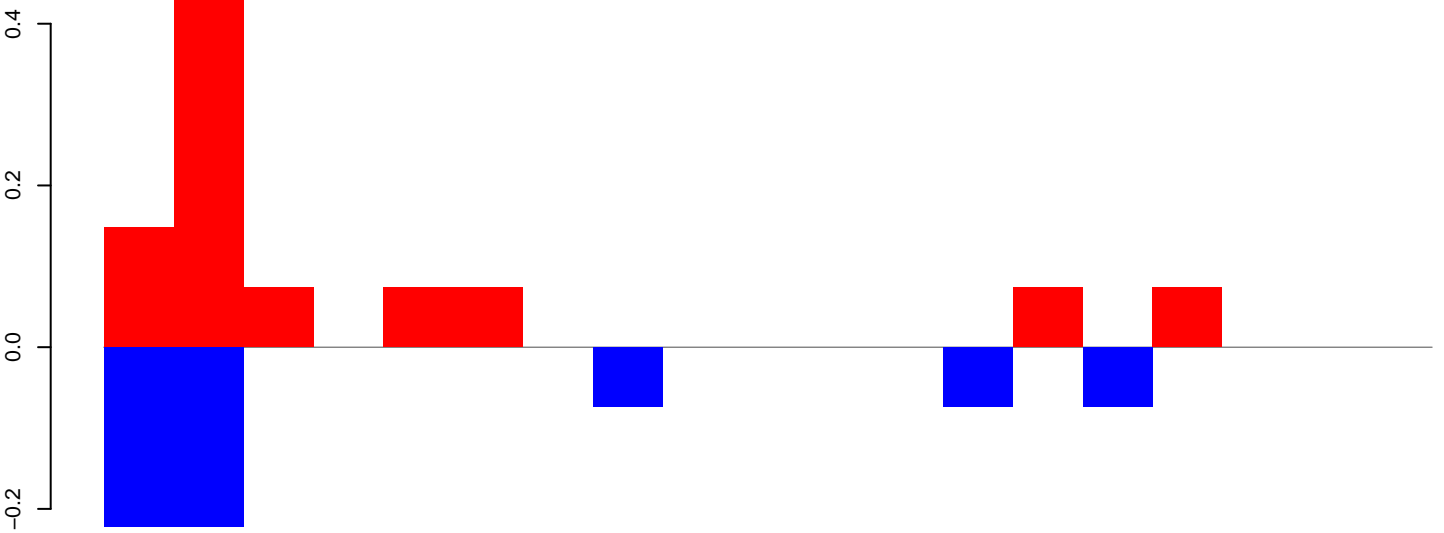


CuQuin_Testes_NL.rep



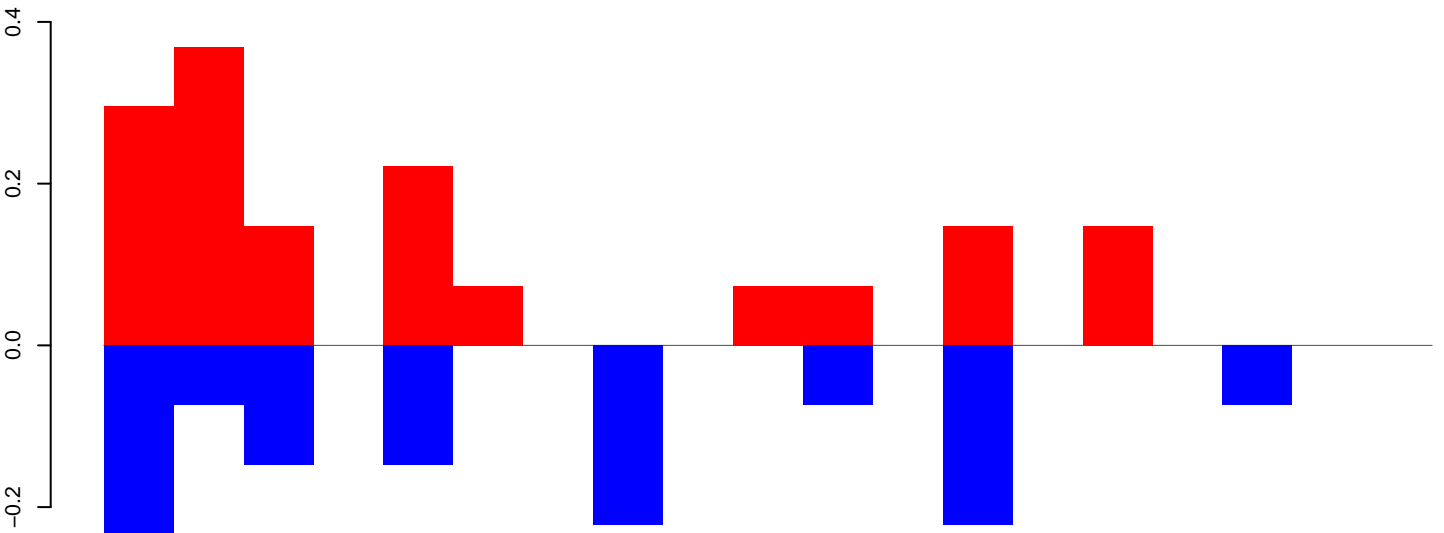
Window size=25, length=706, TE@rnd-4_family-960@DNA_hAT-hATm@769:1-706

CuQuin_Testes_NL.18_23.rep



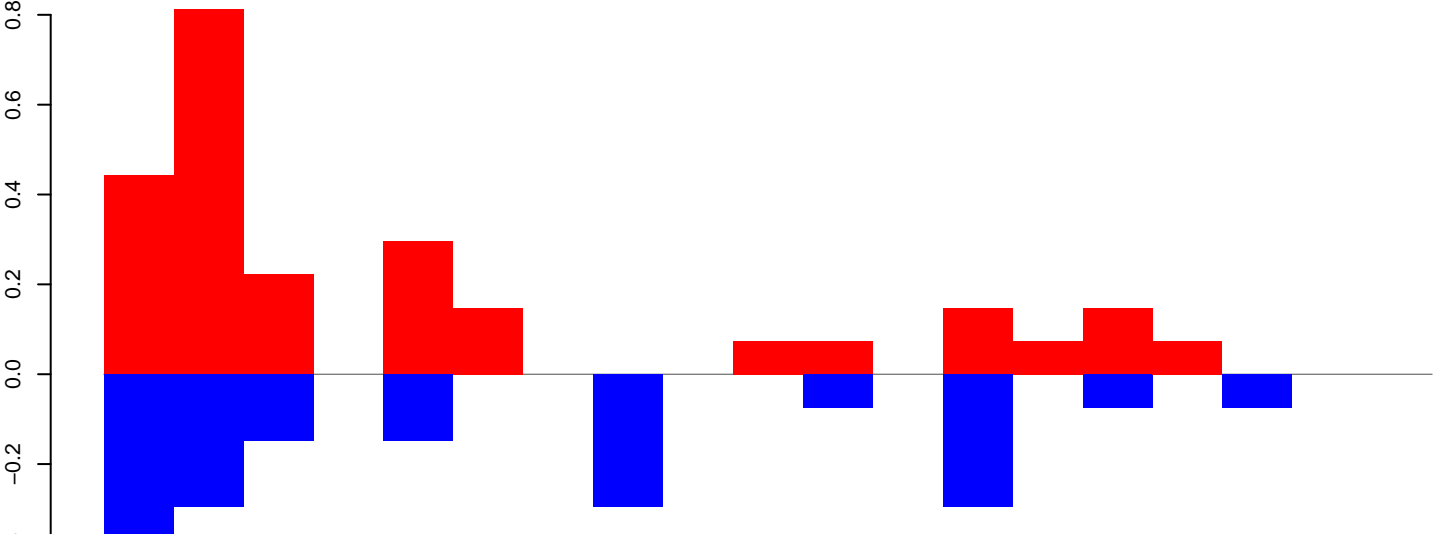
positive=1, negative=1, total=2

CuQuin_Testes_NL.24_35.rep



positive=2, negative=1, total=3

CuQuin_Testes_NL.rep

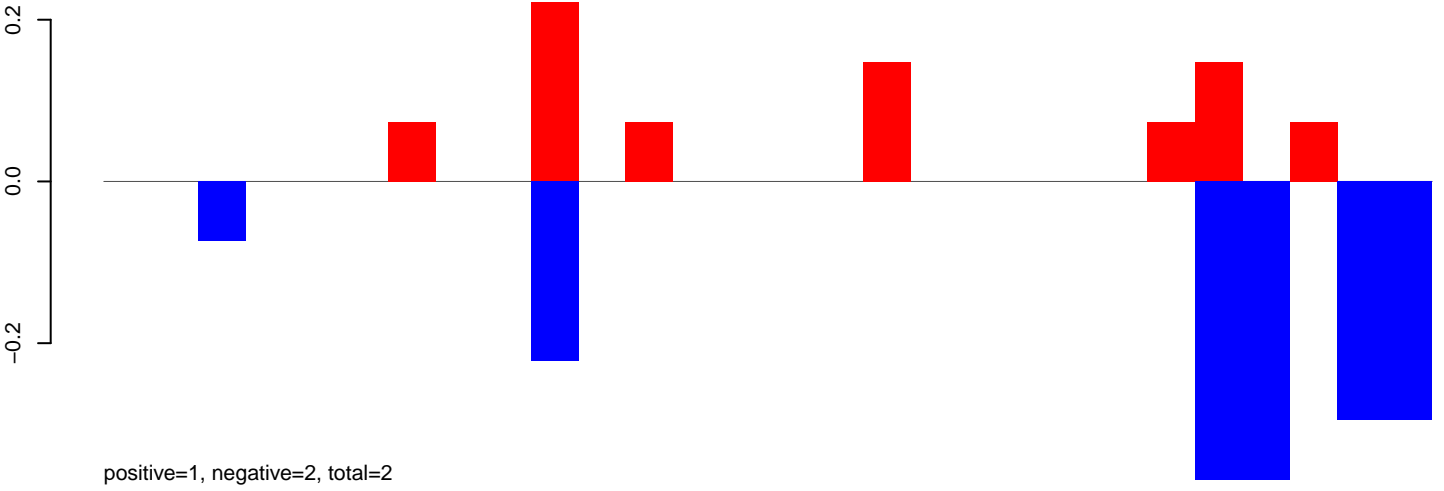


positive=3, negative=2, total=4

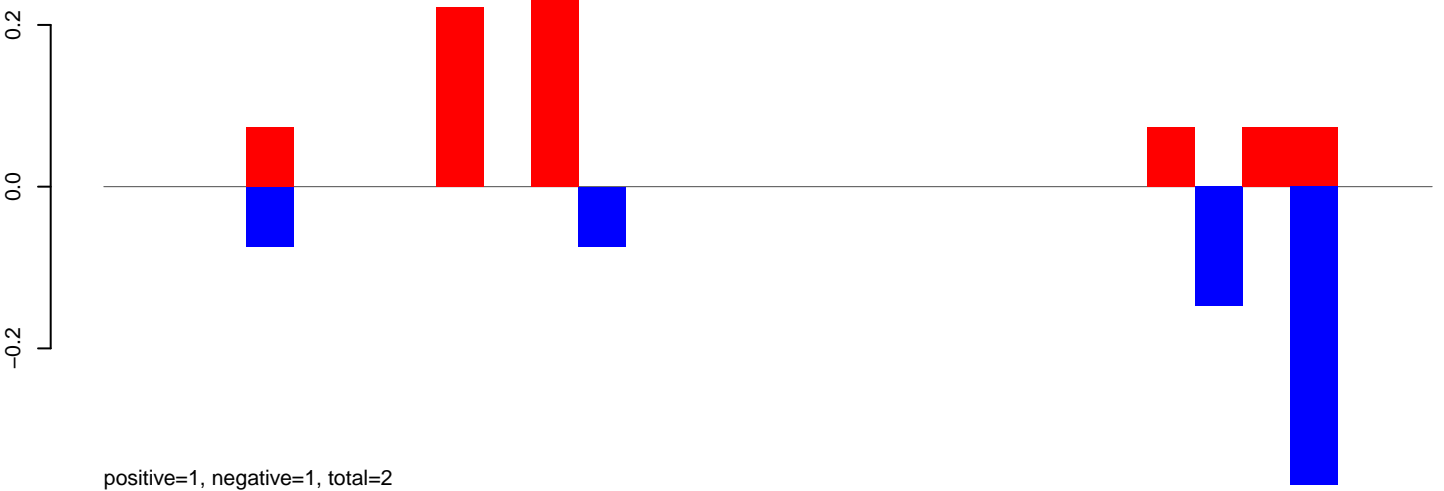
Window size=25, length=464, TE@rnd-4_family-527@Unknown@2827:1-464

100 200 300 400 500

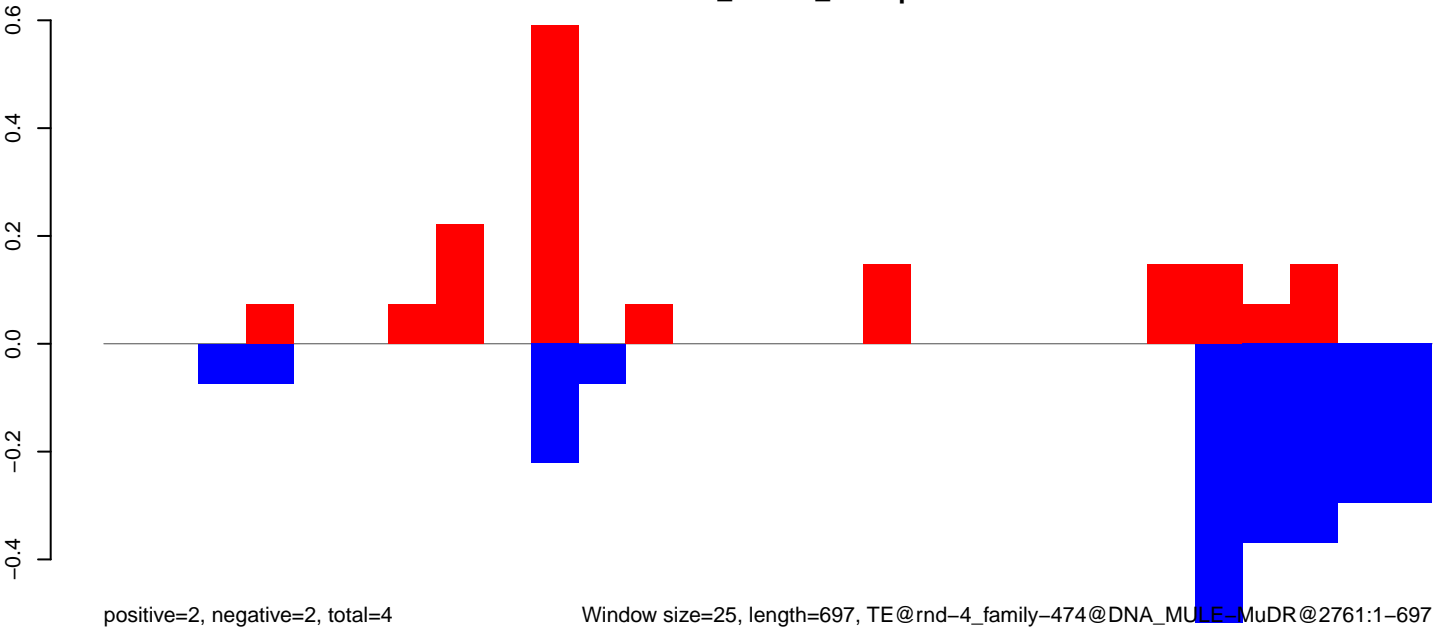
CuQuin_Testes_NL.18_23.rep



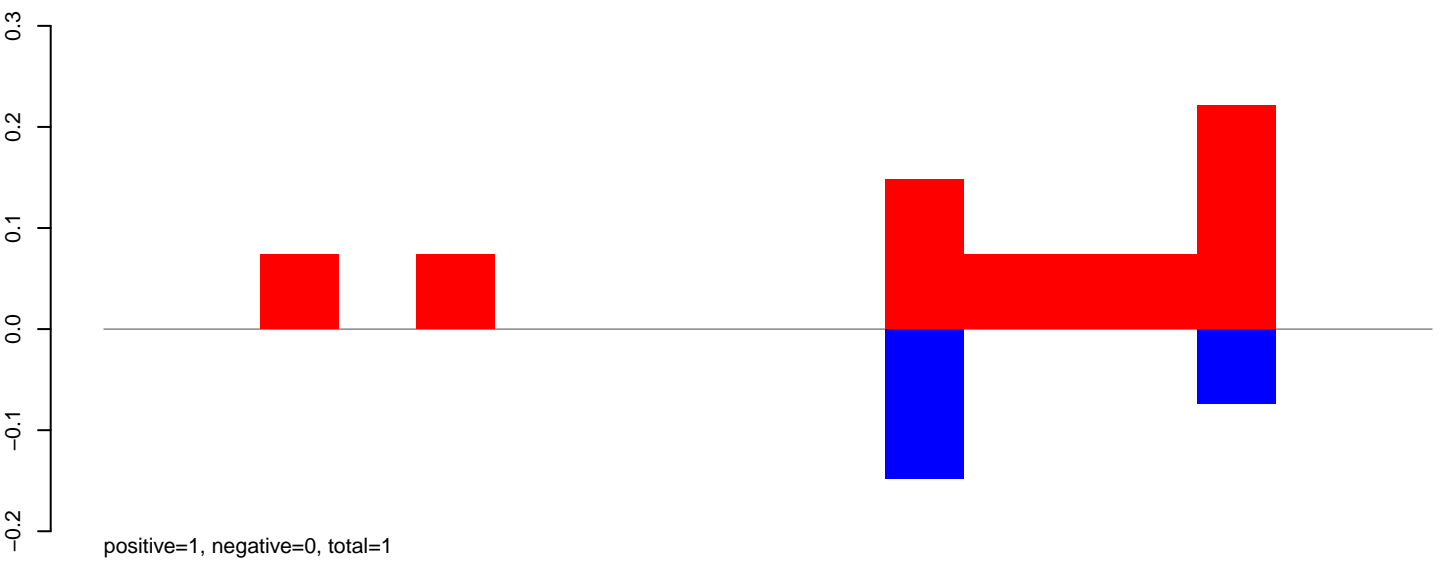
CuQuin_Testes_NL.24_35.rep



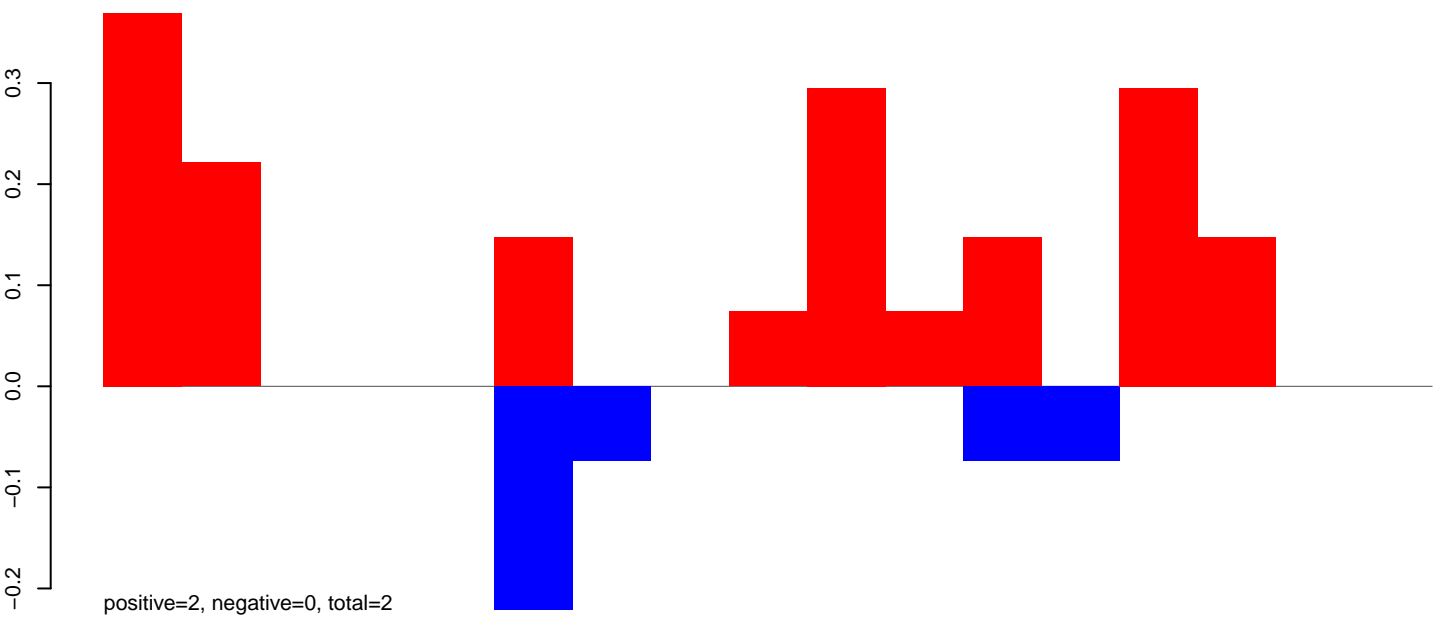
CuQuin_Testes_NL.rep



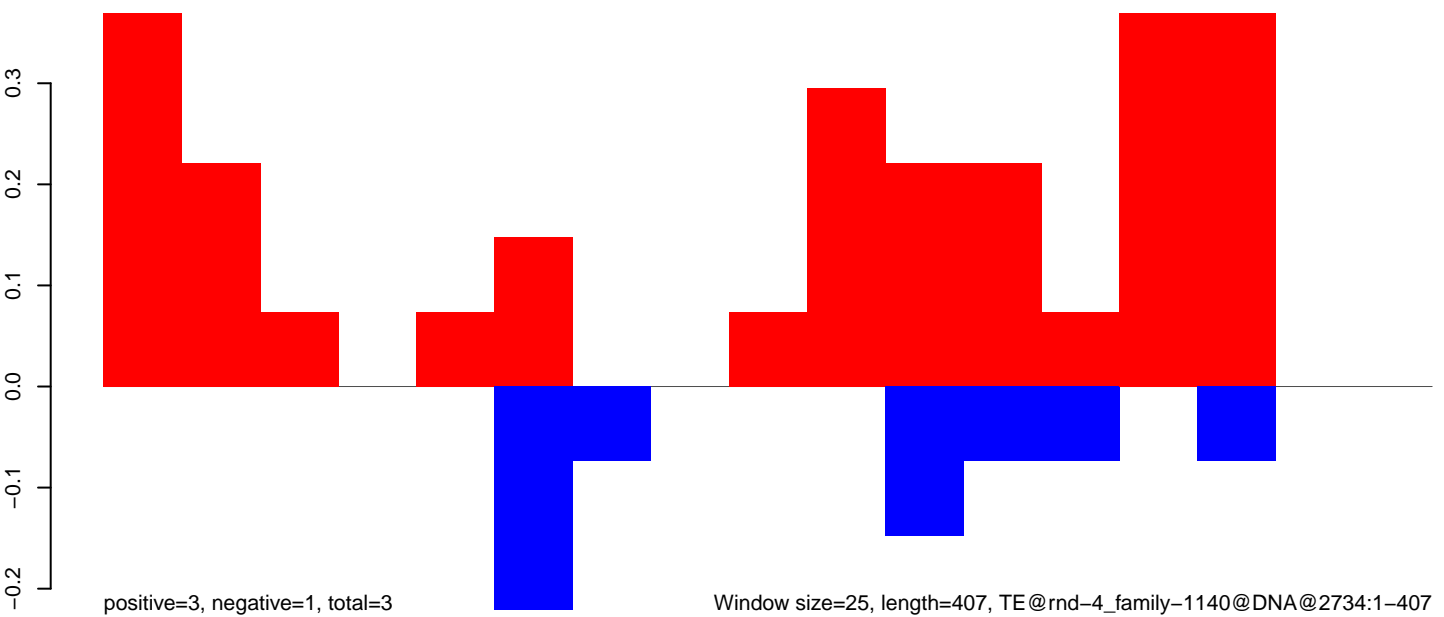
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



Window size=25, length=407, TE@rnd-4_family-1140@DNA@2734:1-407

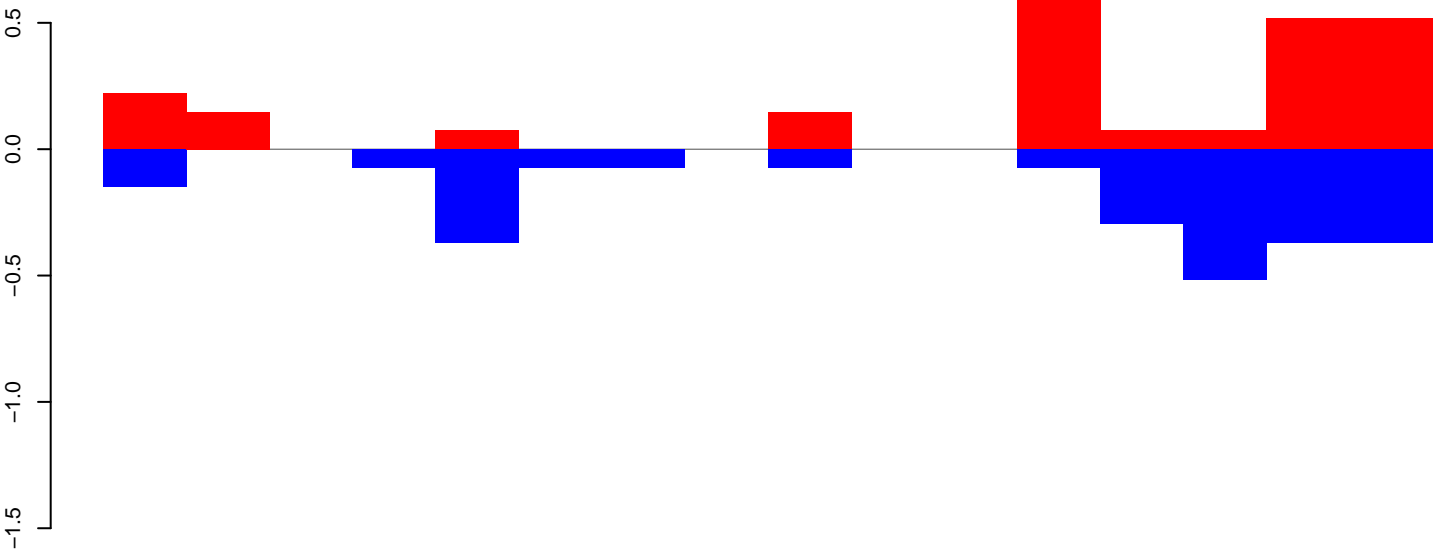
100

200

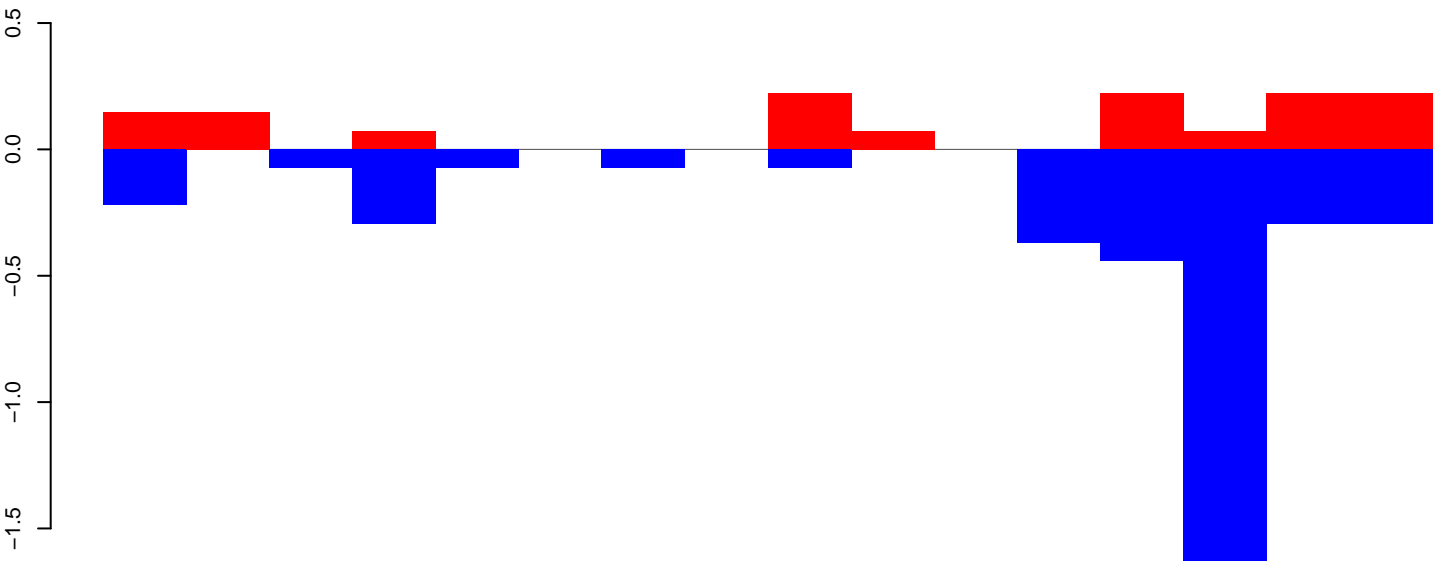
300

400

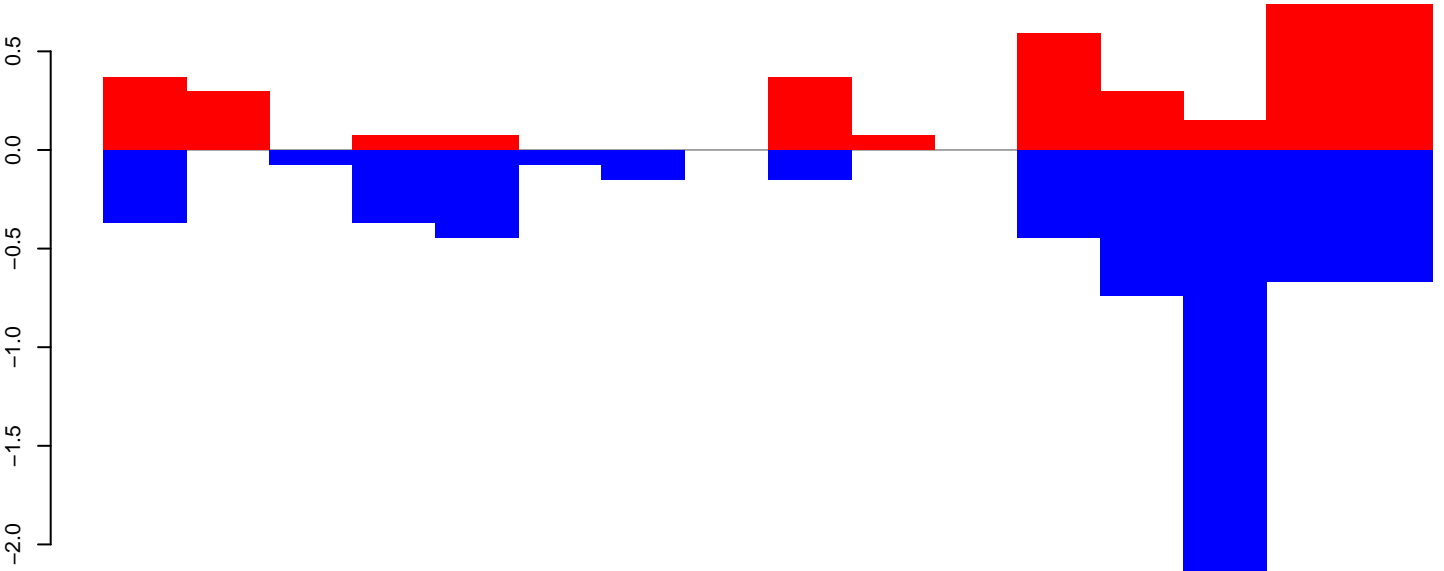
CuQuin_Testes_NL.18_23.rep



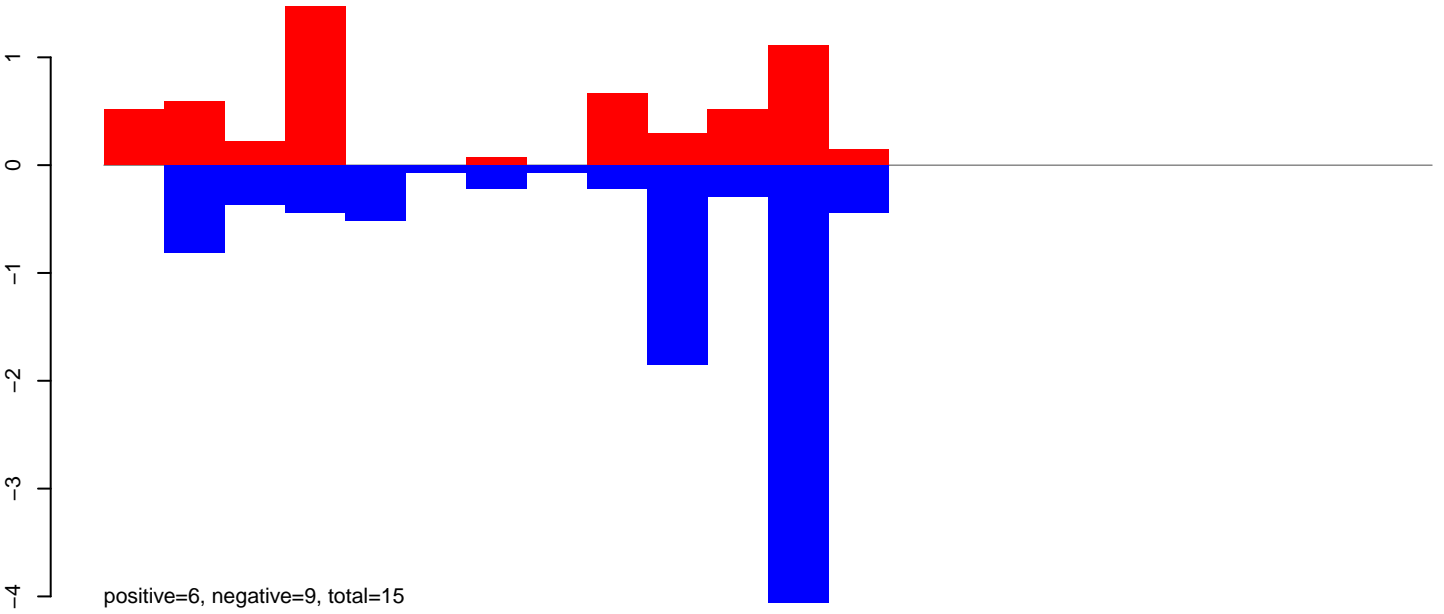
CuQuin_Testes_NL.24_35.rep



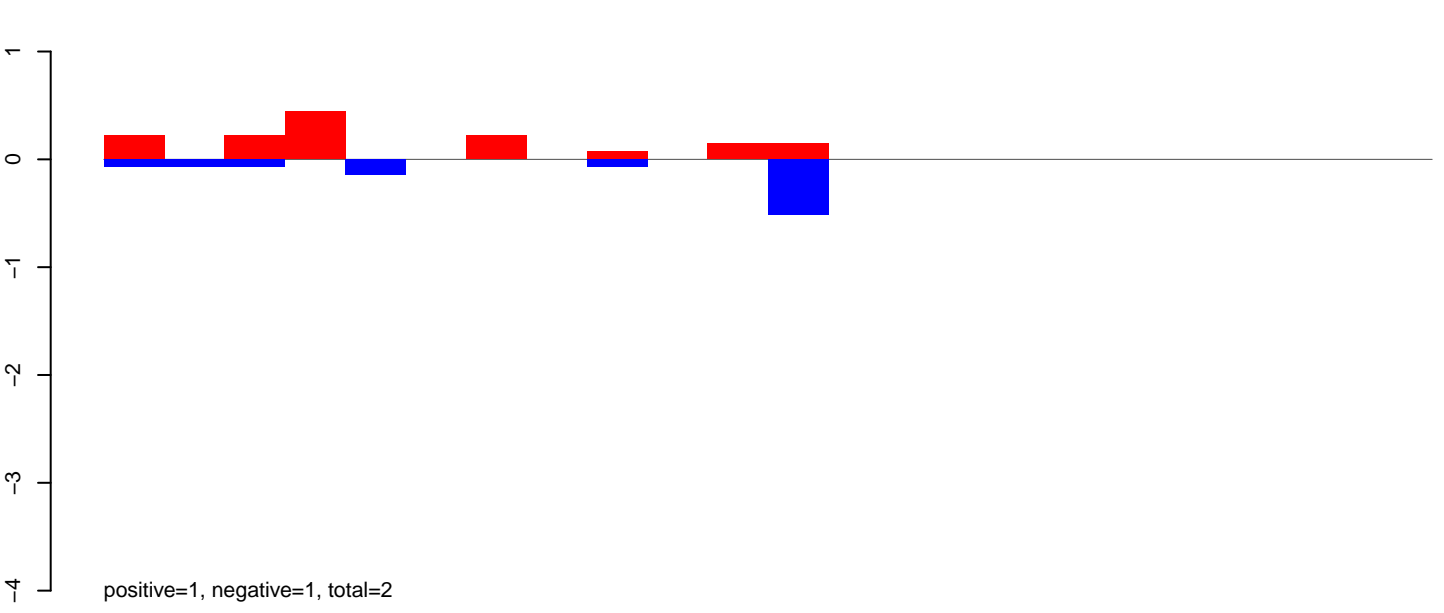
CuQuin_Testes_NL.rep



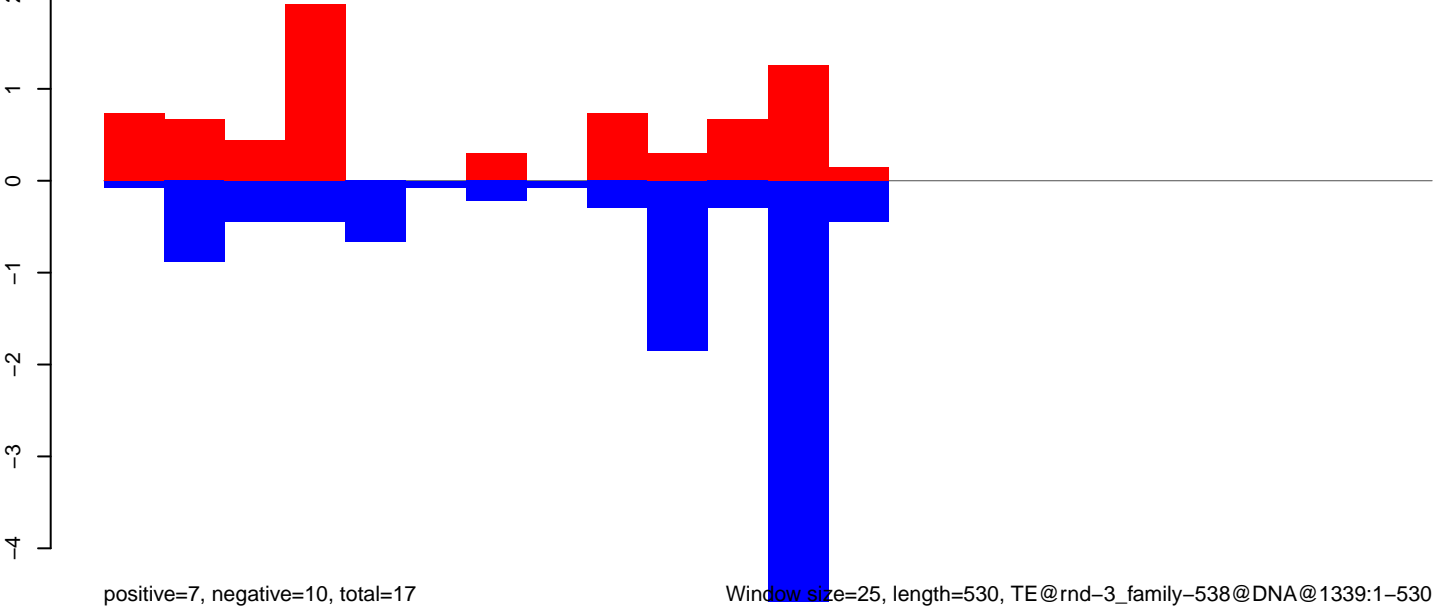
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

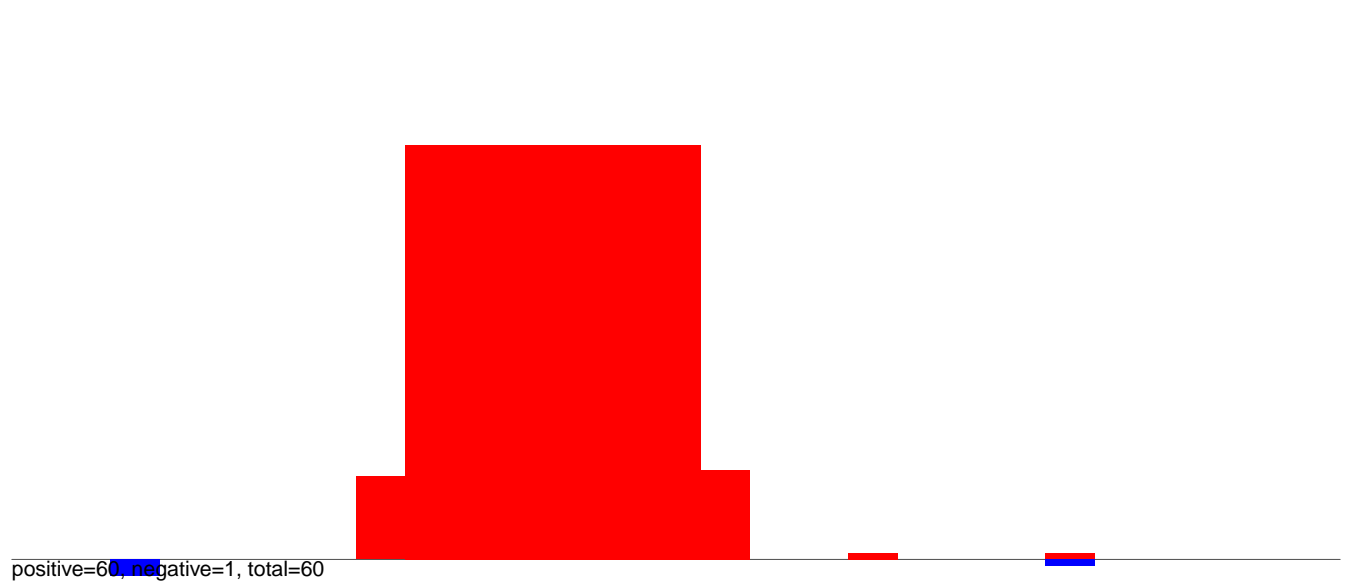


CuQuin_Testes_NL.rep

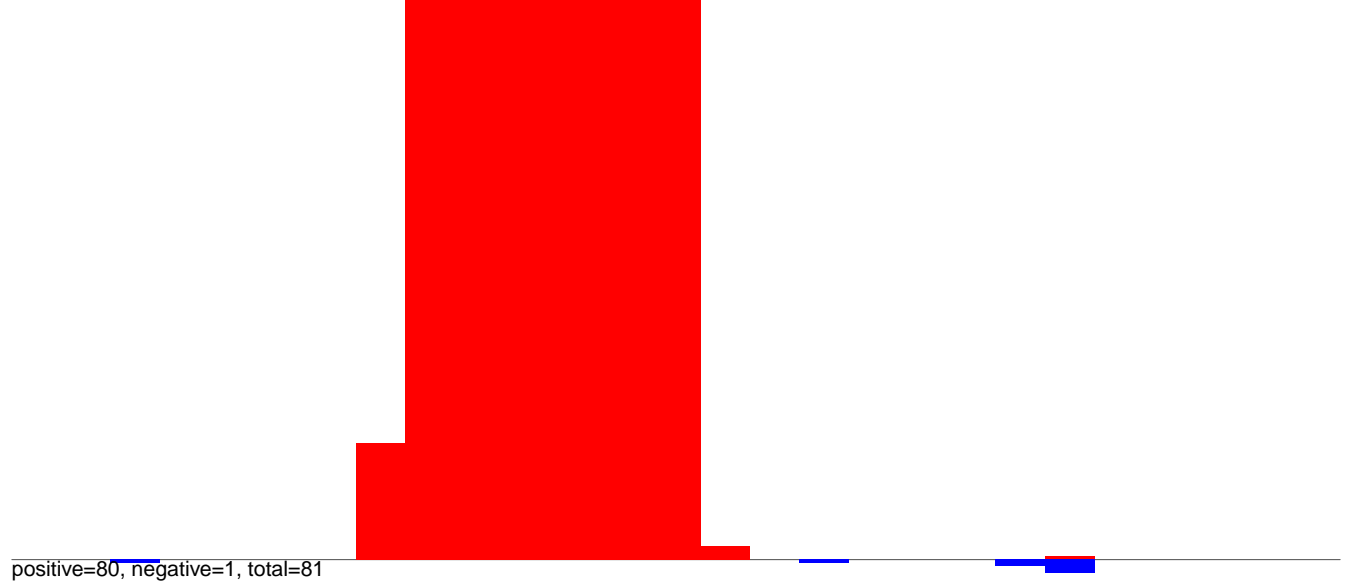


Window size=25, length=530, TE@rnd-3_family-538@DNA@1339:1-530

CuQuin_Testes_NL.18_23.rep



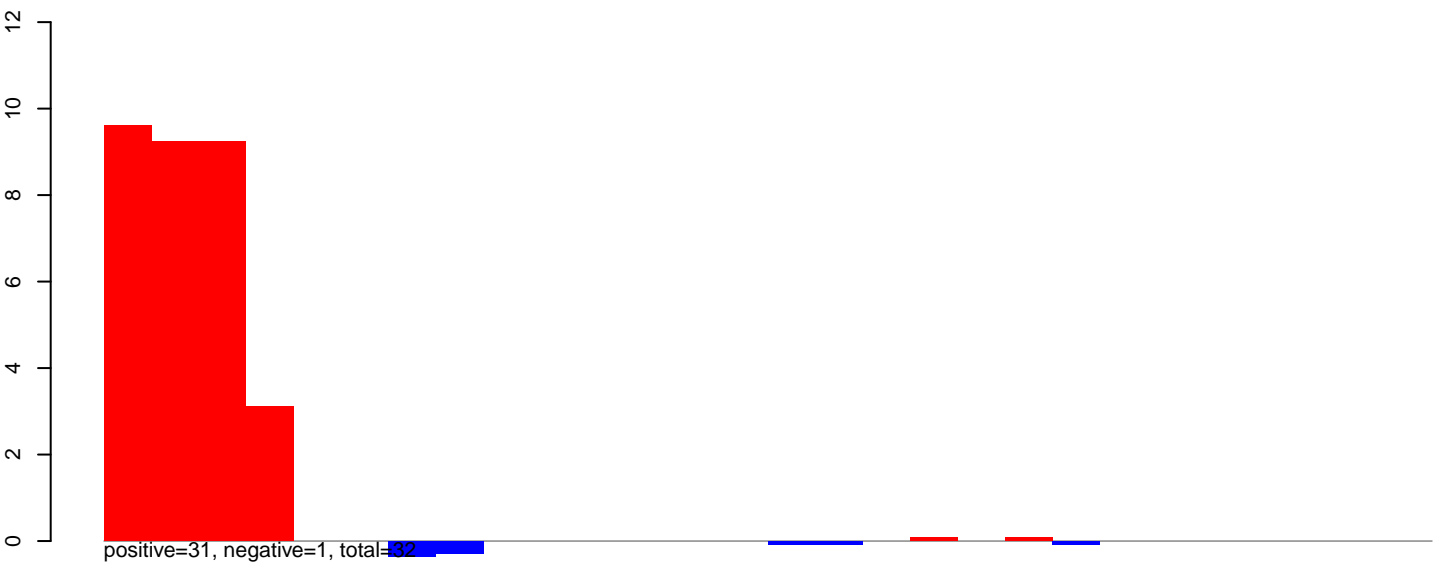
CuQuin_Testes_NL.24_35.rep



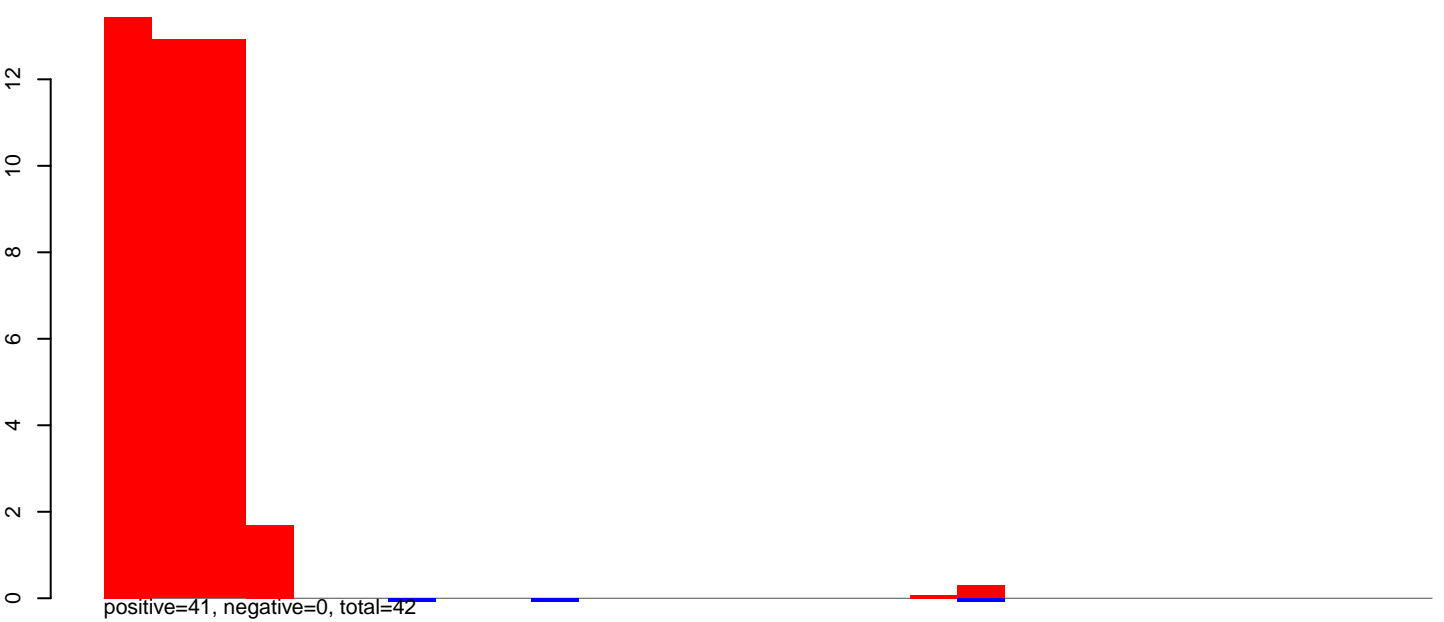
CuQuin_Testes_NL.rep



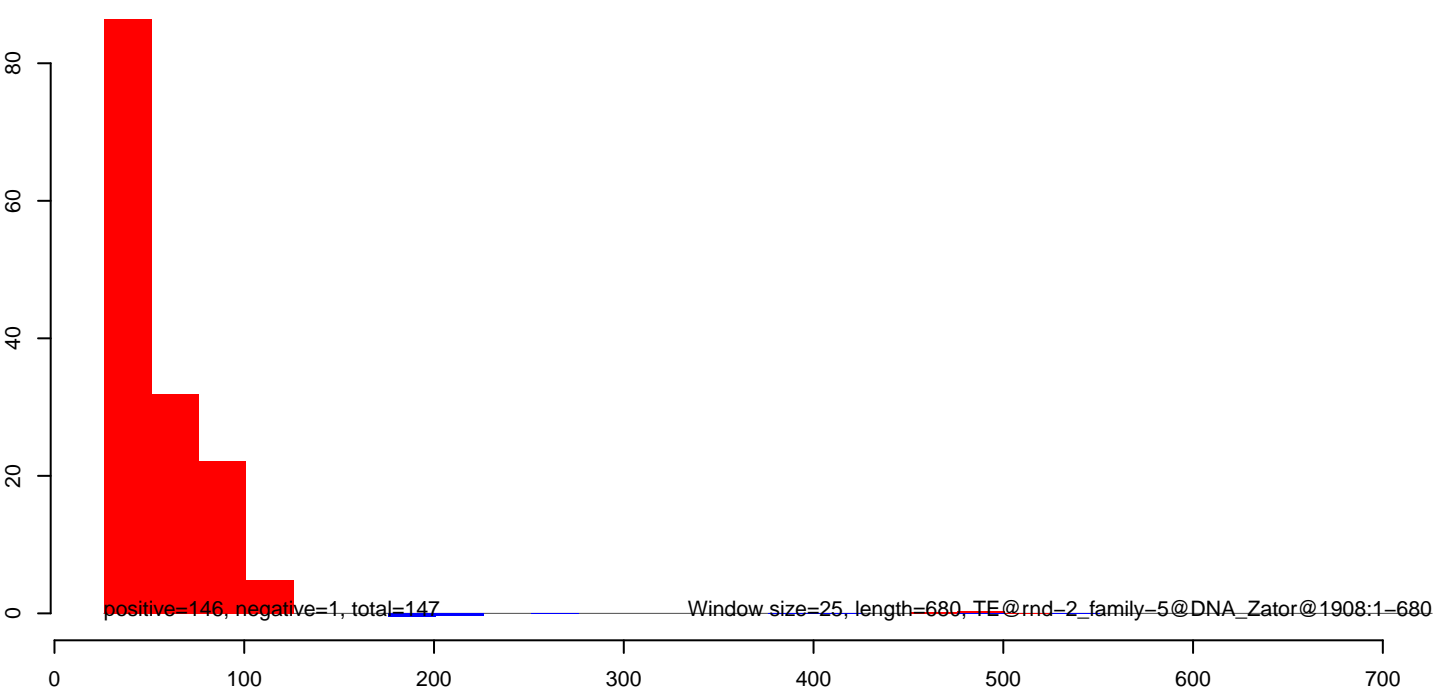
CuQuin_Testes_NL.18_23.rep



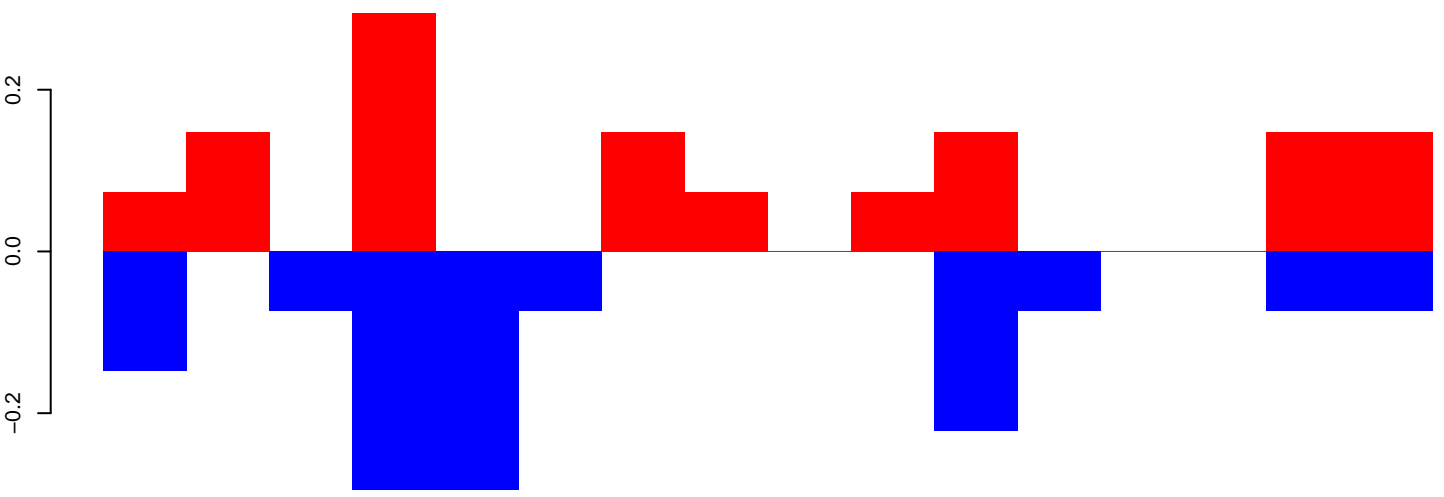
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

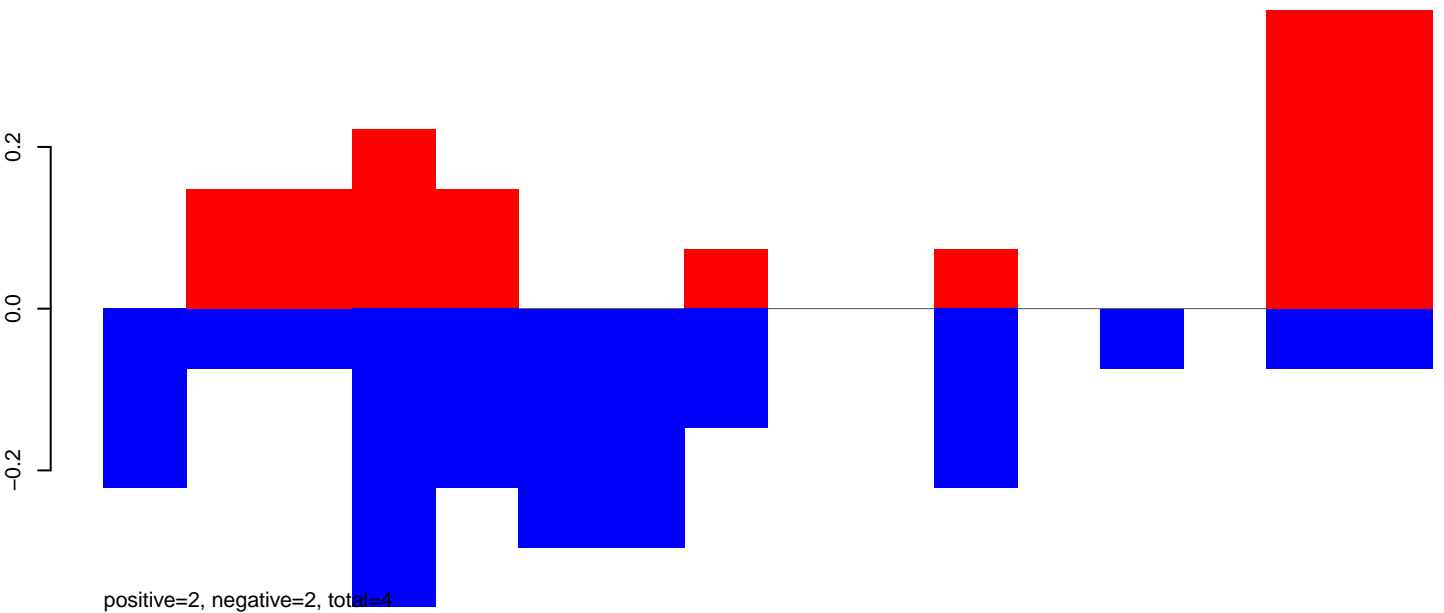


CuQuin_Testes_NL.18_23.rep



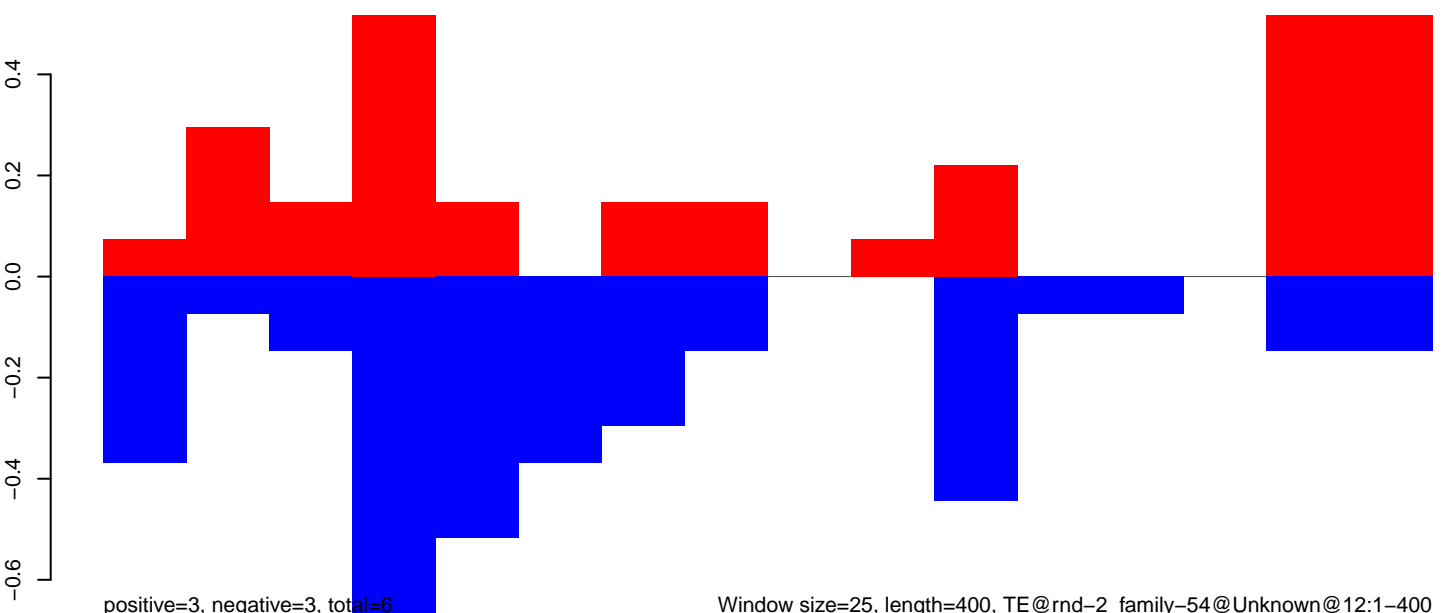
positive=1, negative=1, total=3

CuQuin_Testes_NL.24_35.rep



positive=2, negative=2, total=4

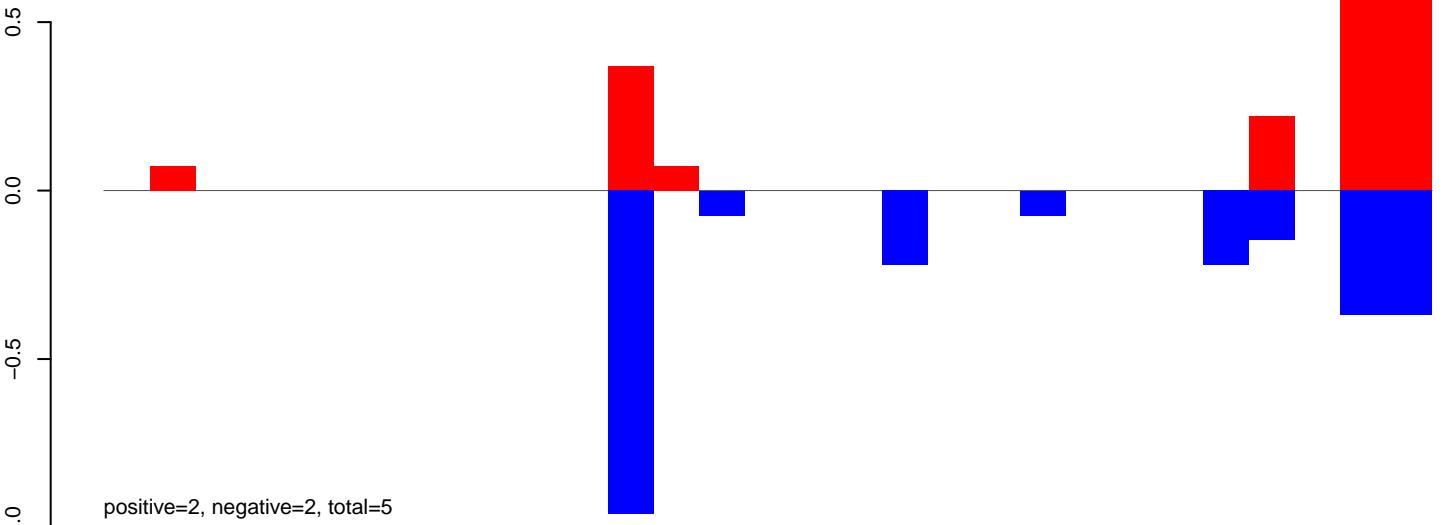
CuQuin_Testes_NL.rep



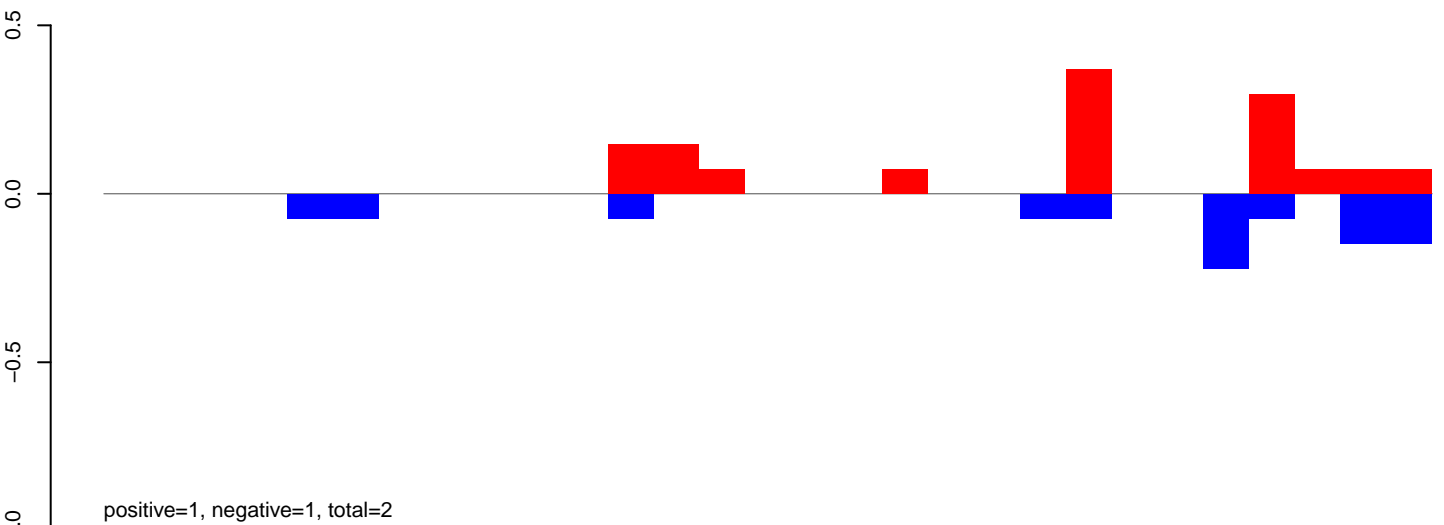
positive=3, negative=3, total=6

Window size=25, length=400, TE@rnd-2_family-54@Unknown@12:1-400

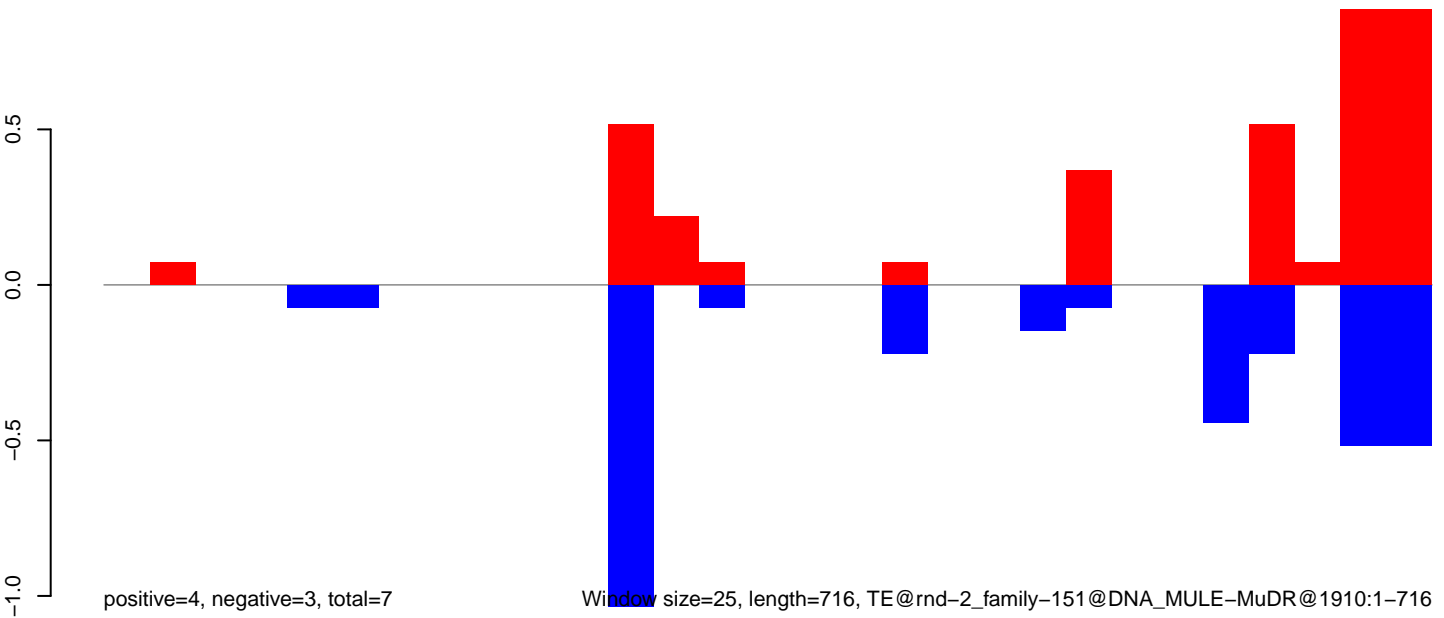
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

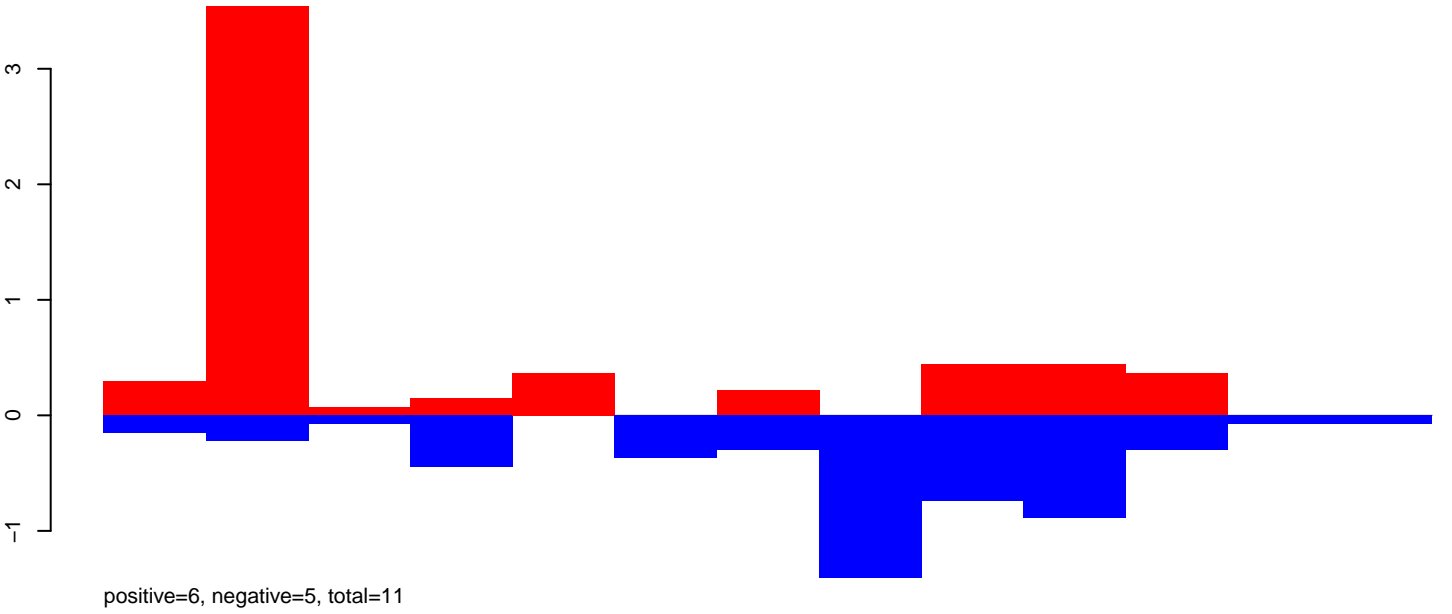


CuQuin_Testes_NL.rep

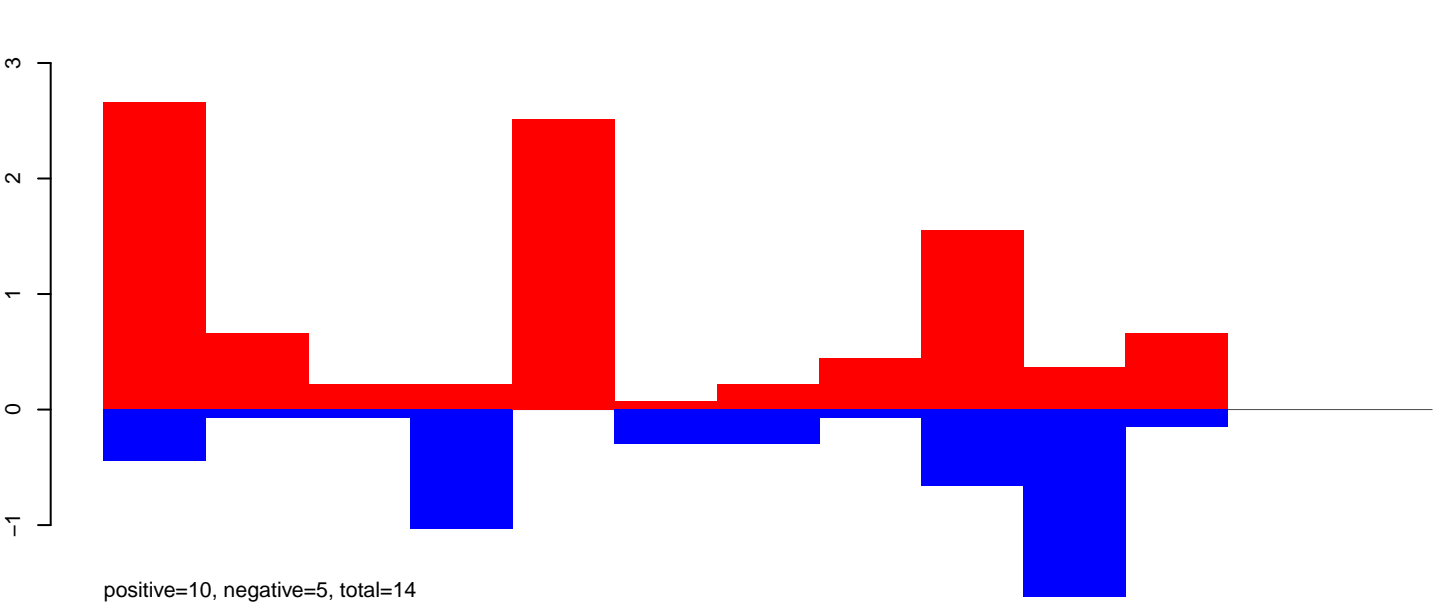


Window size=25, length=716, TE@rnd-2_family-151@DNA_MULE-MuDR@1910:1-716

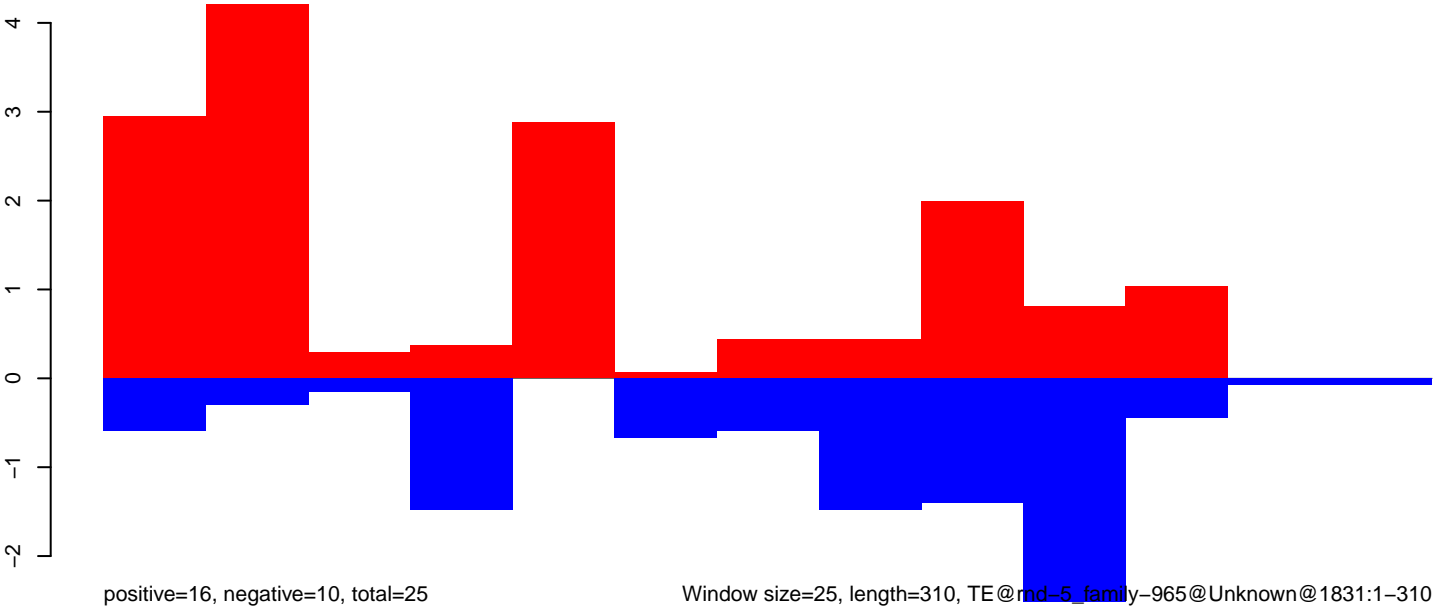
CuQuin_Testes_NL.18_23.rep



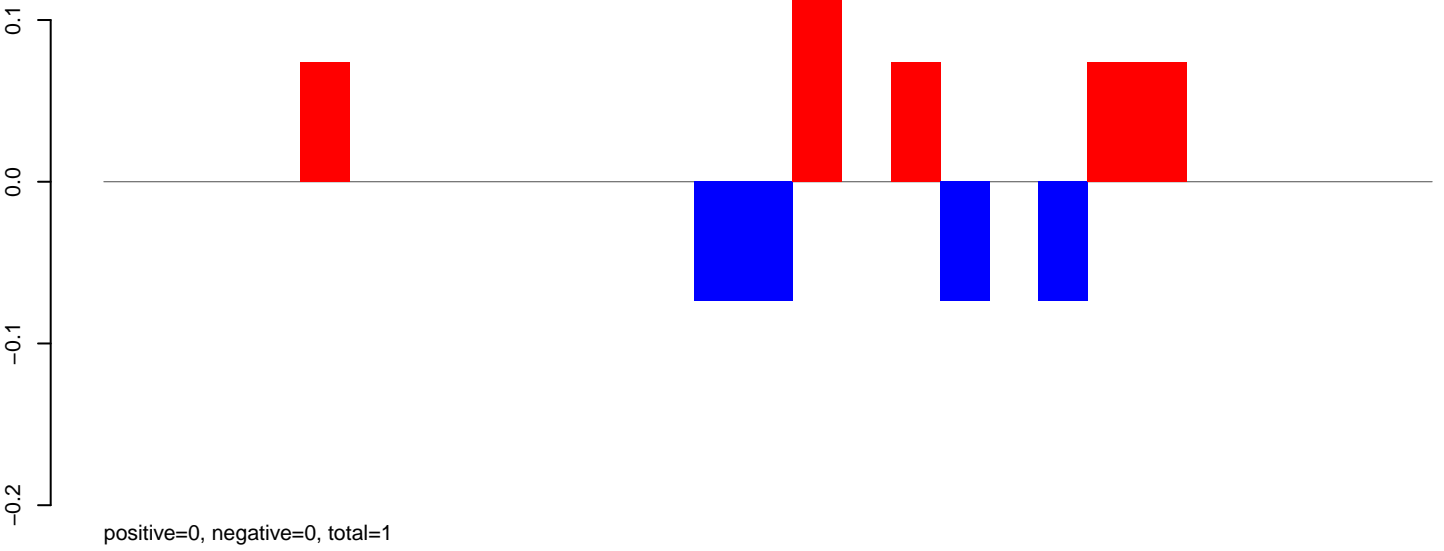
CuQuin_Testes_NL.24_35.rep



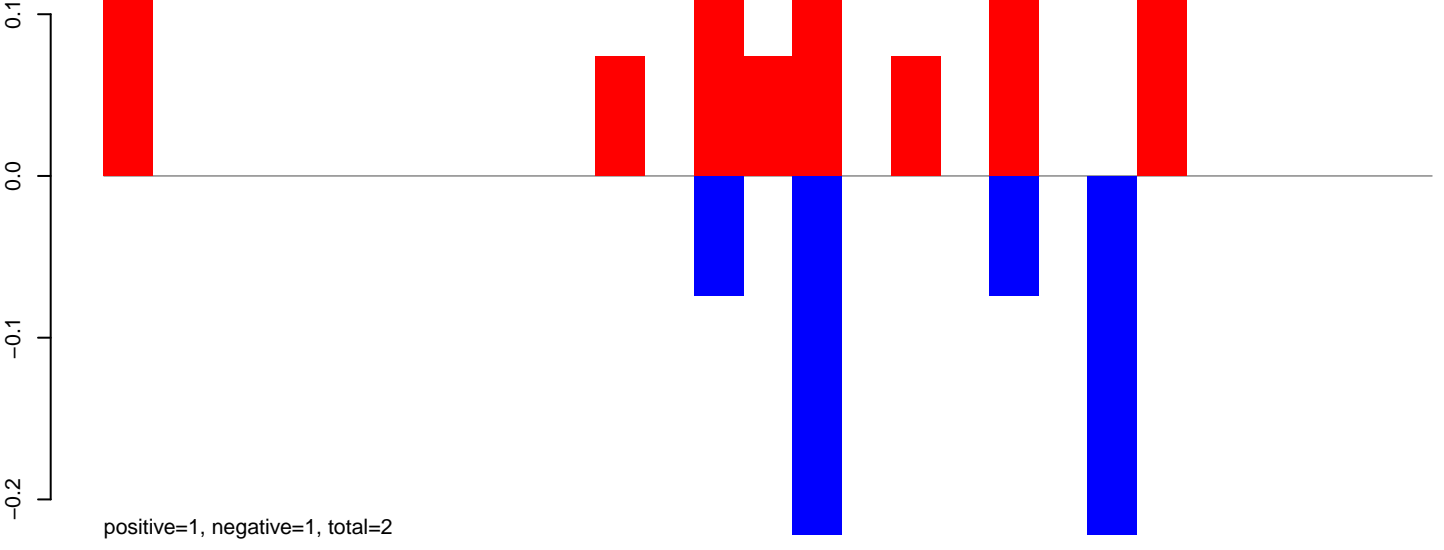
CuQuin_Testes_NL.rep



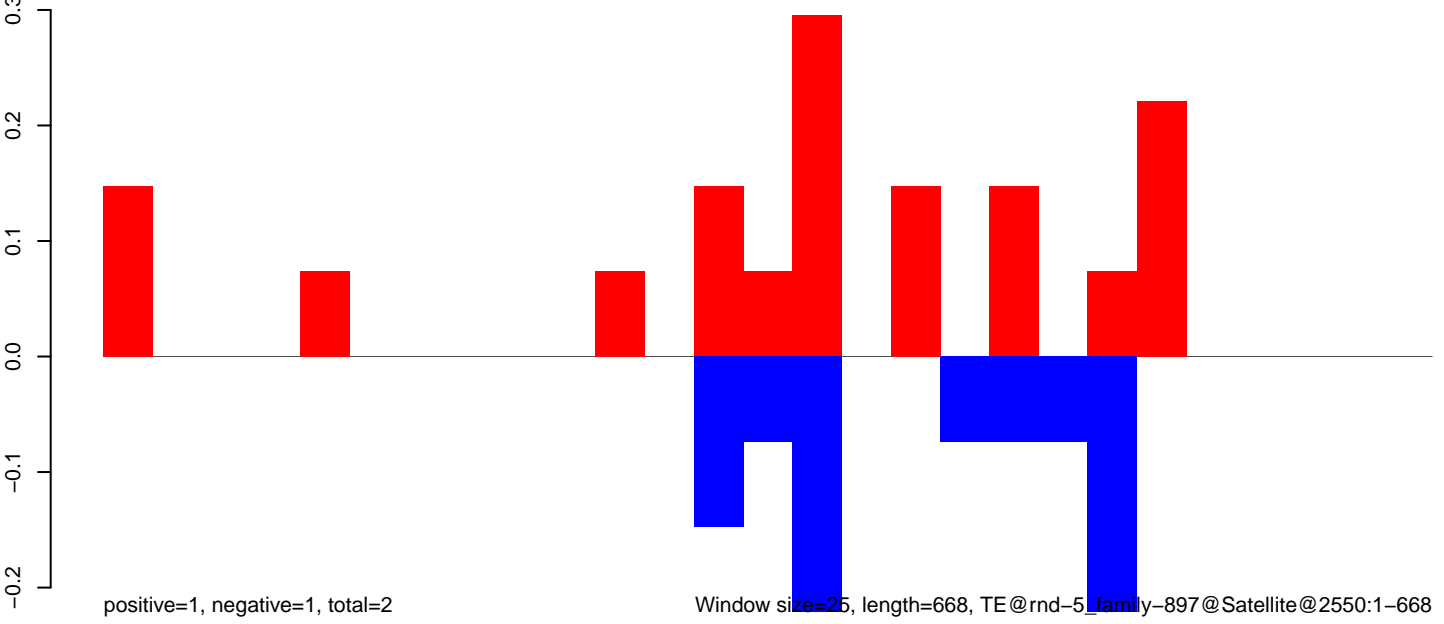
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

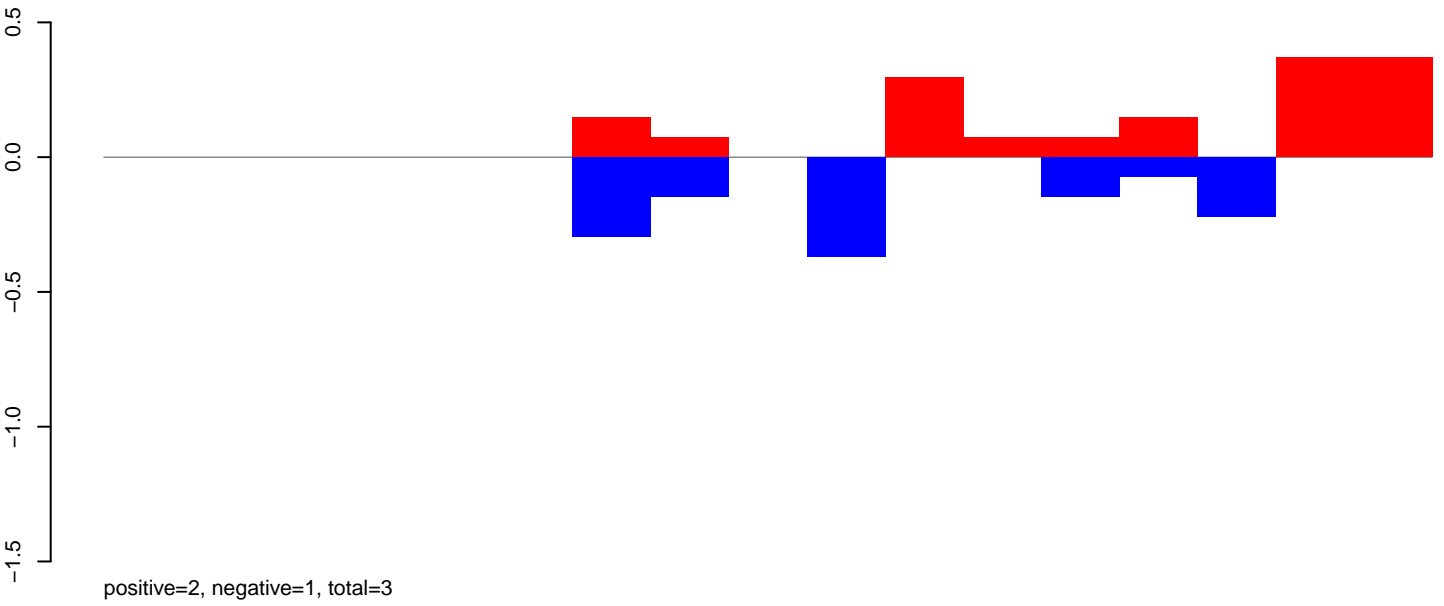


CuQuin_Testes_NL.rep

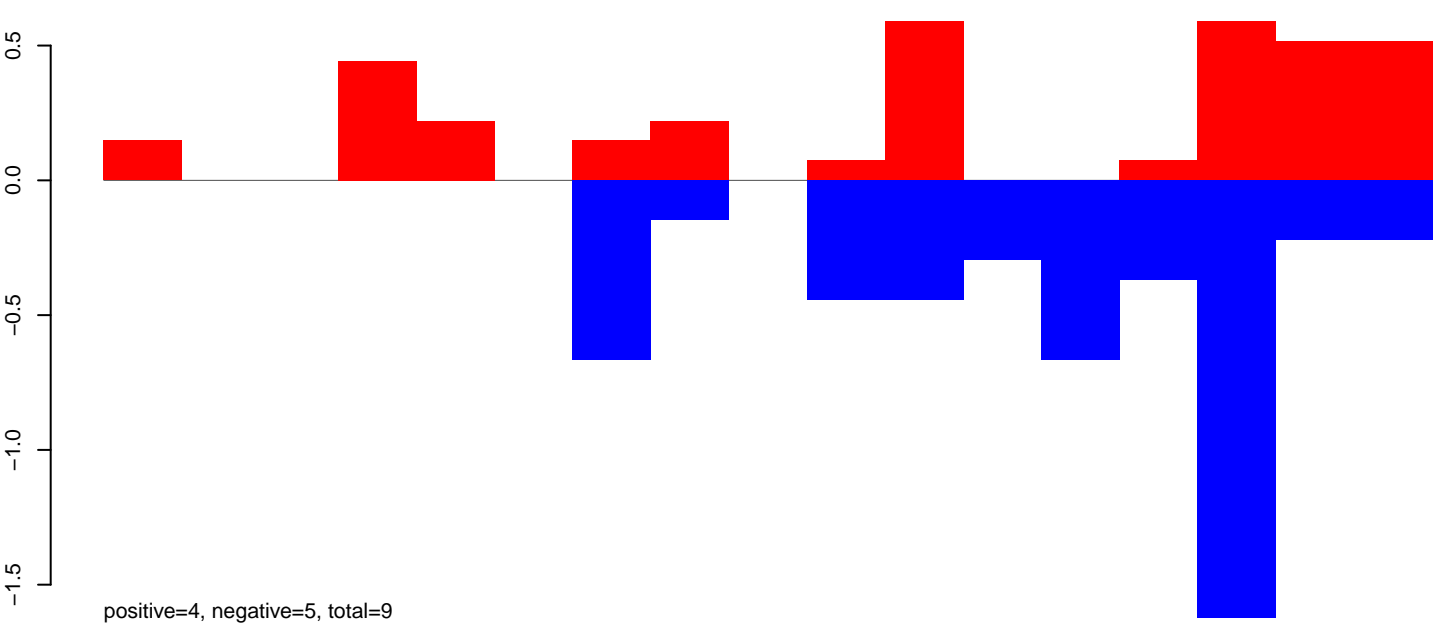


Window size=25, length=668, TE@rnd-5_family-897@Satellite@2550:1-668

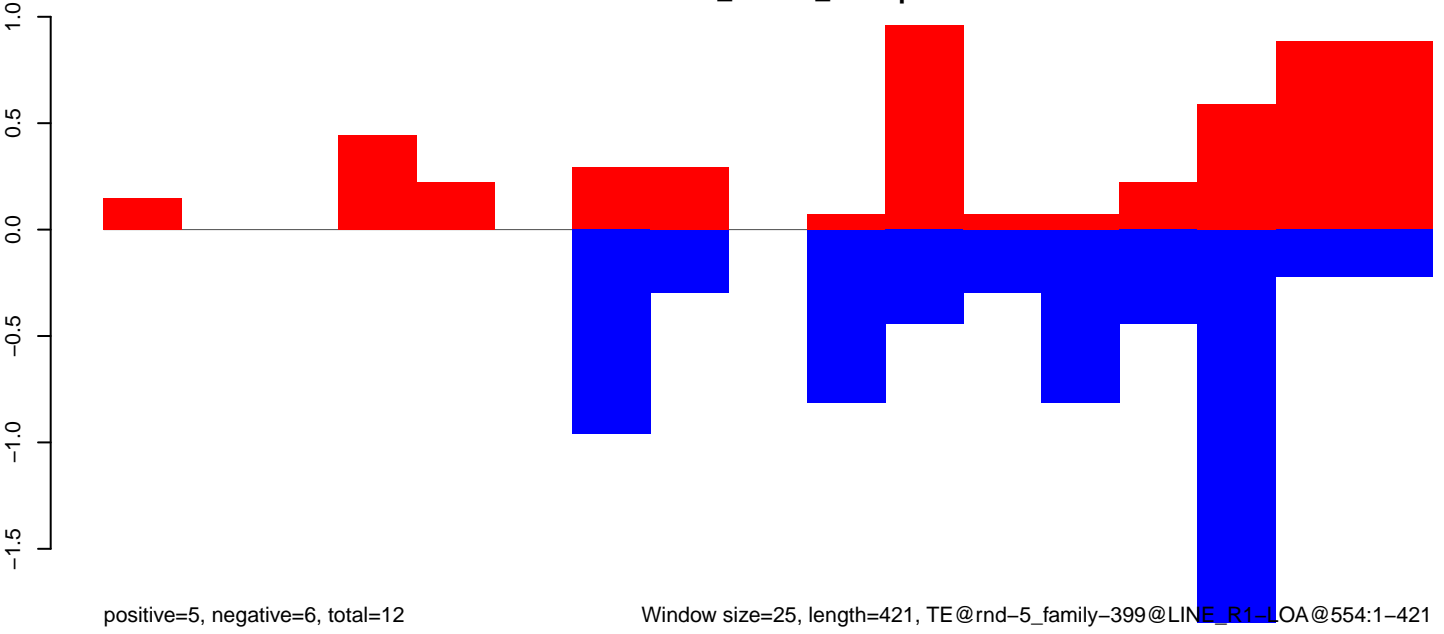
CuQuin_Testes_NL.18_23.rep



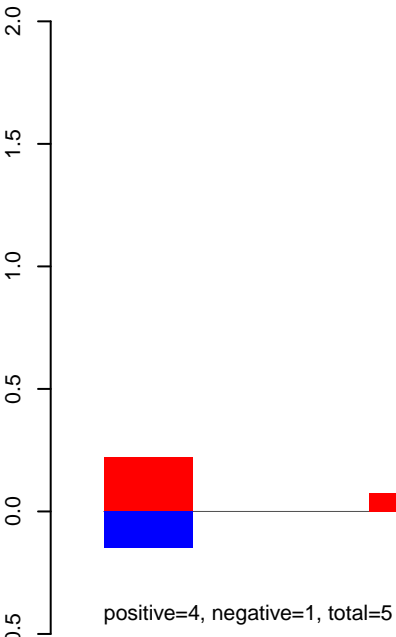
CuQuin_Testes_NL.24_35.rep



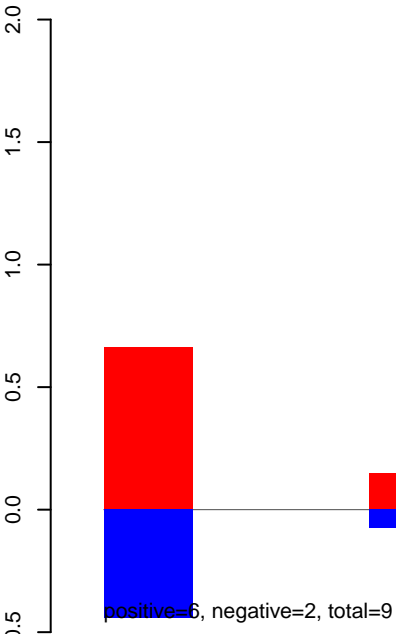
CuQuin_Testes_NL.rep



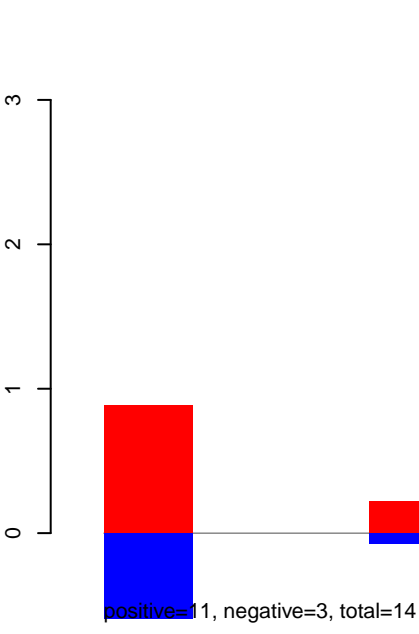
CuQuin_Testes_NL.18_23.rep



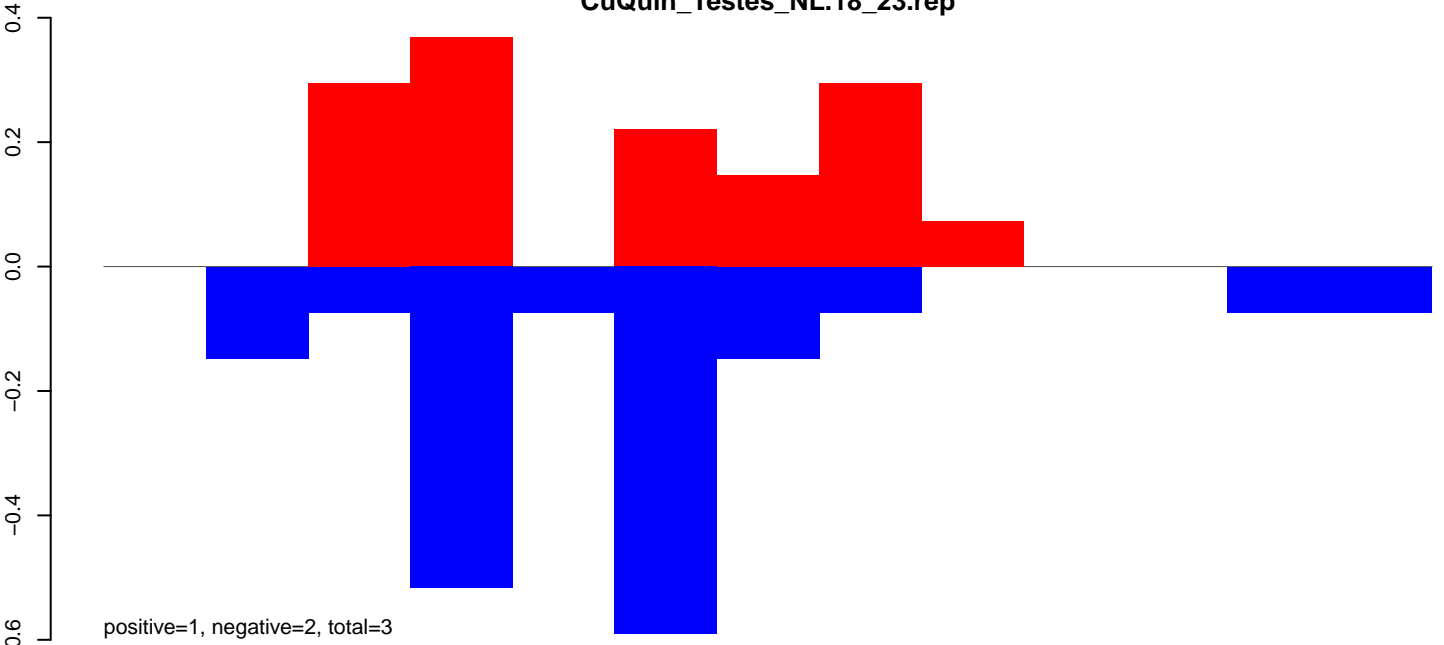
CuQuin_Testes_NL.24_35.rep



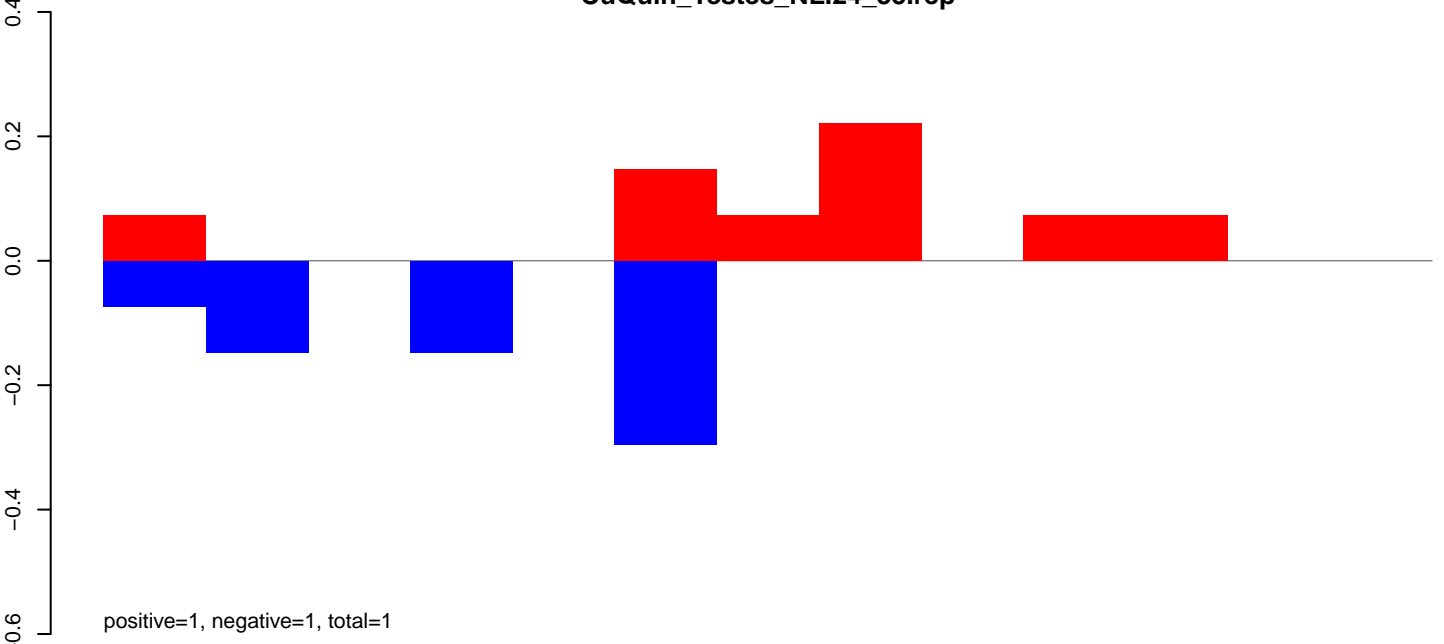
CuQuin_Testes_NL.rep



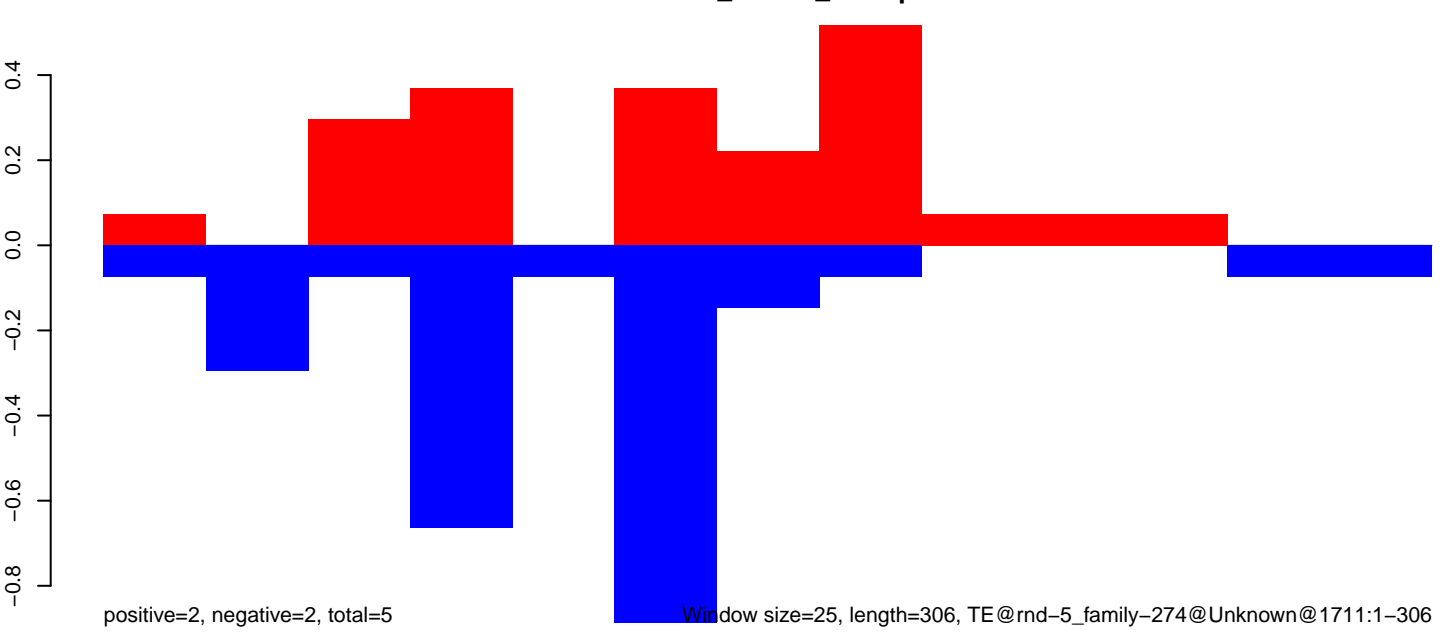
CuQuin_Testes_NL.18_23.rep



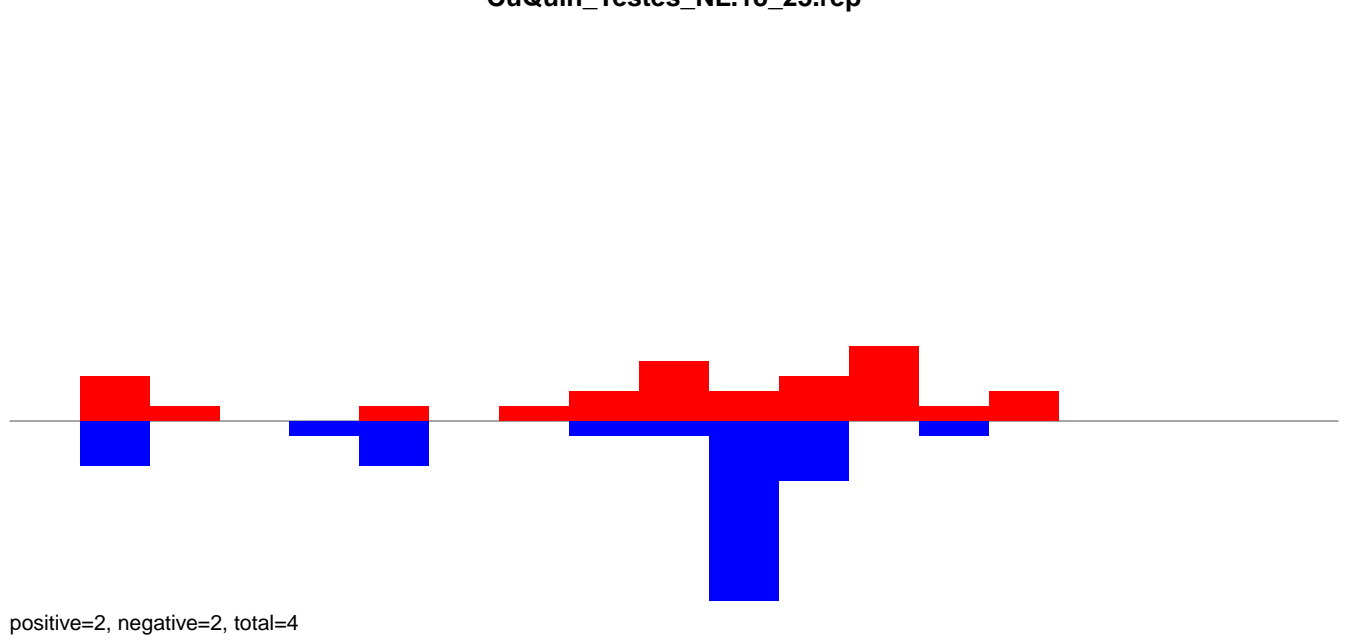
CuQuin_Testes_NL.24_35.rep



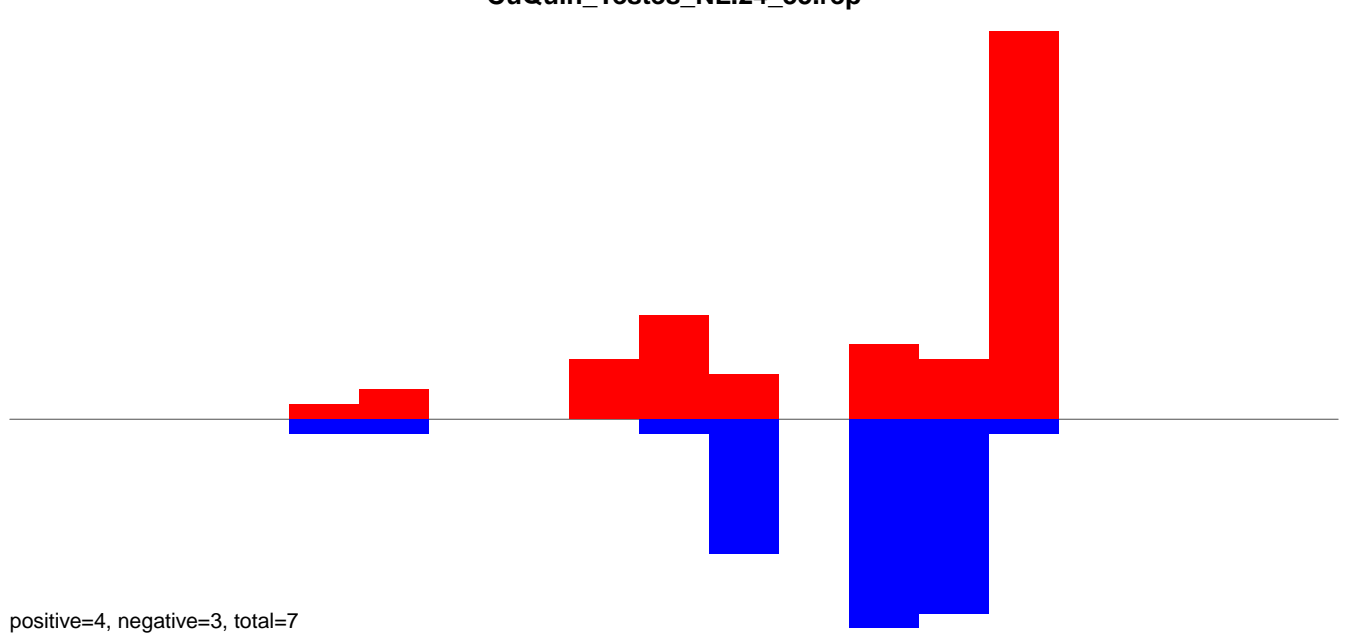
CuQuin_Testes_NL.rep



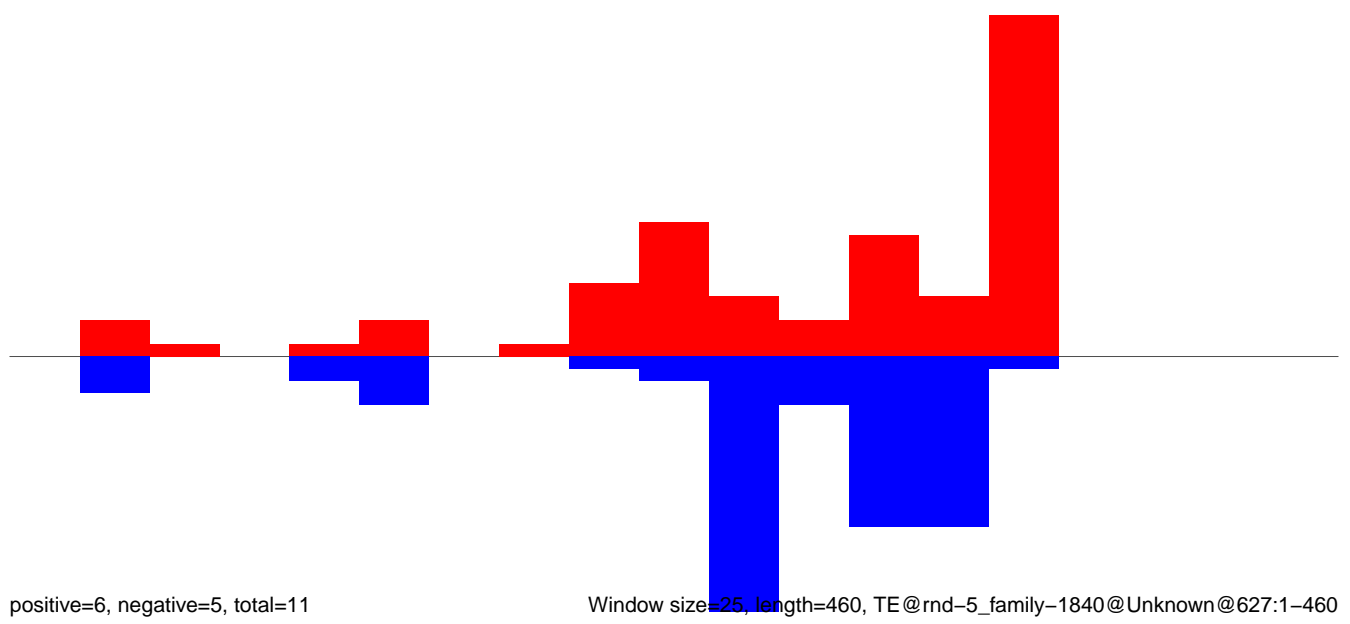
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

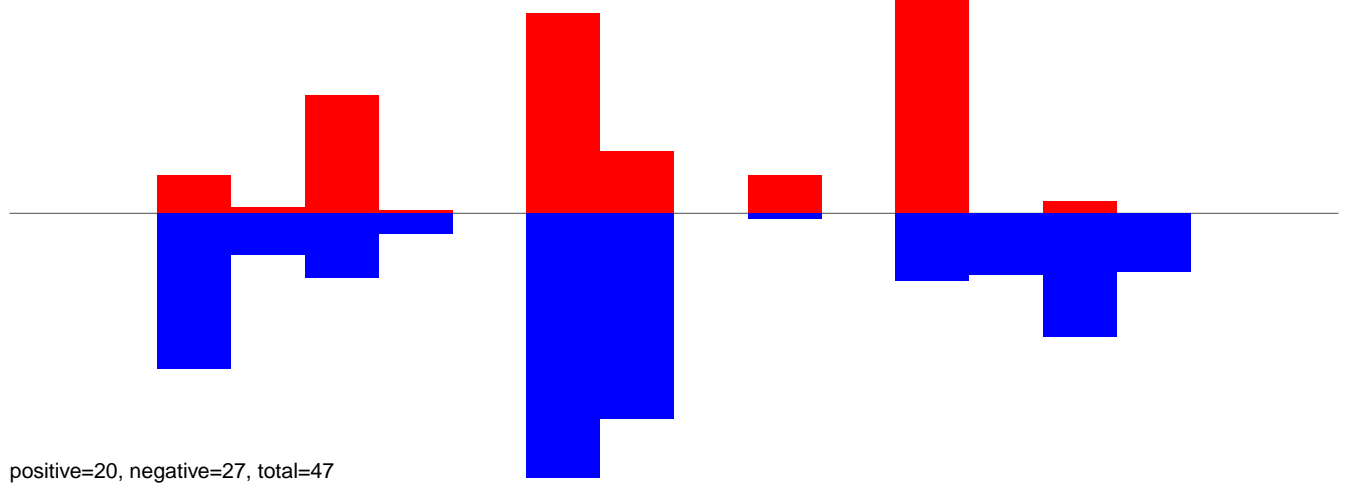


CuQuin_Testes_NL.rep

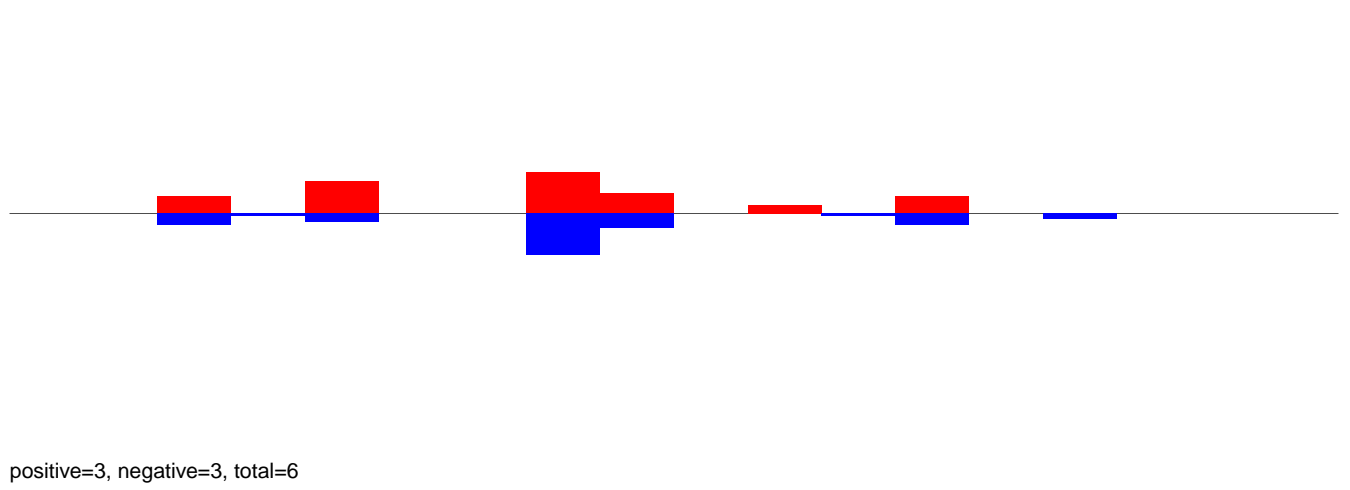


Window size=25, length=460, TE@rnd-5_family-1840@Unknown@627:1-460

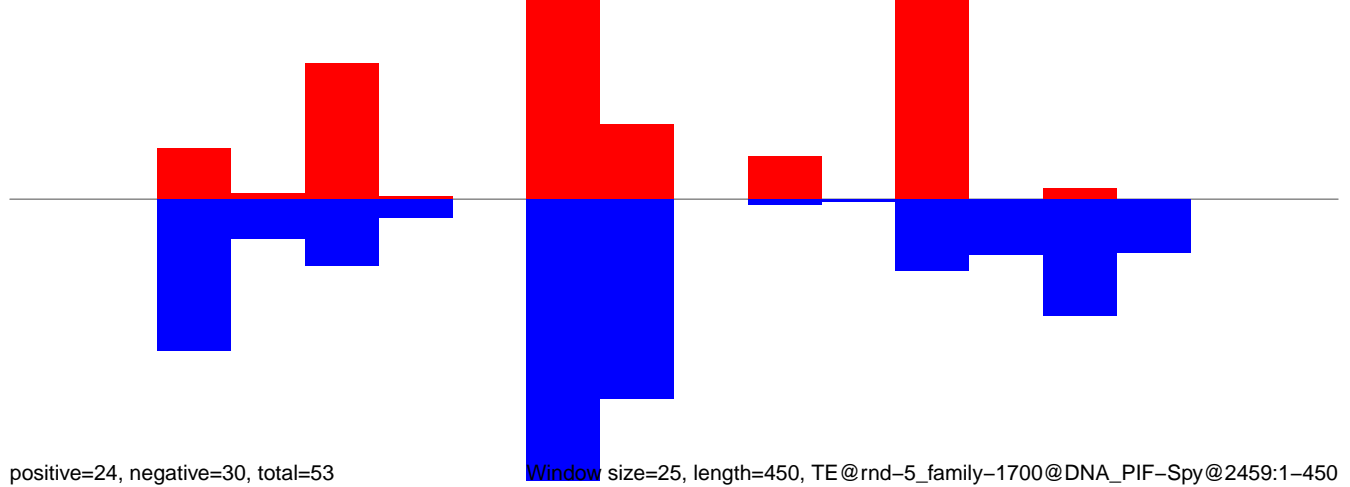
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

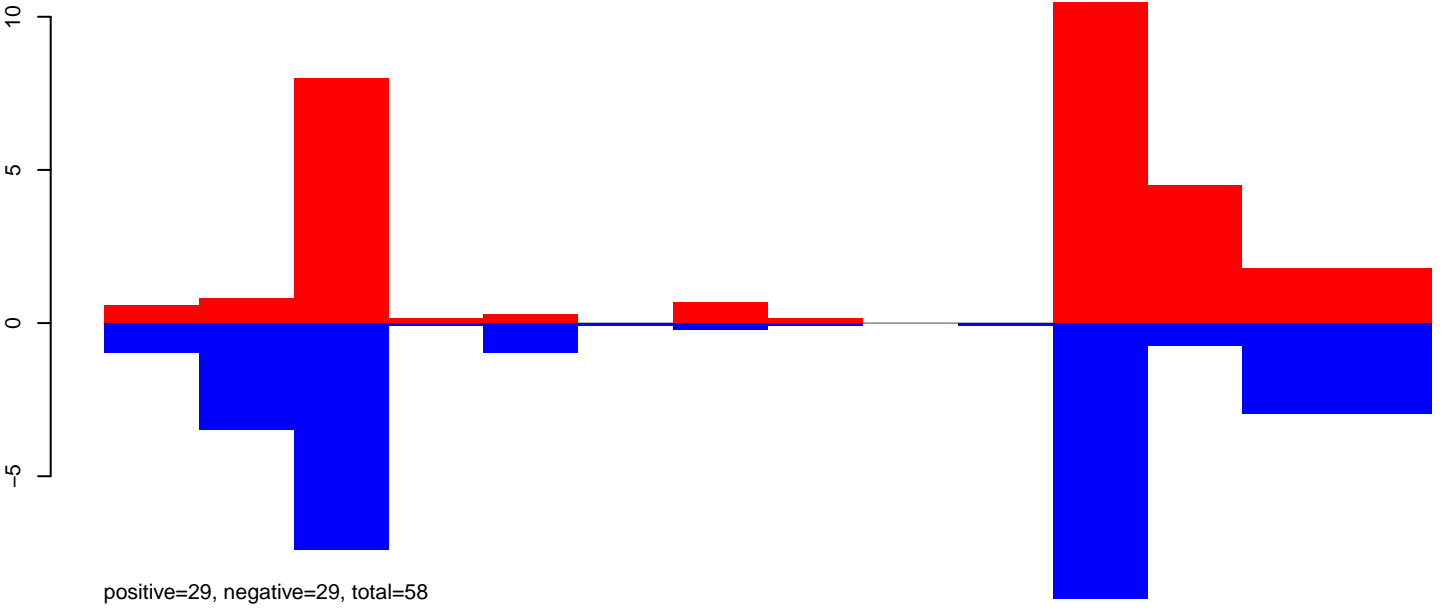


CuQuin_Testes_NL.rep

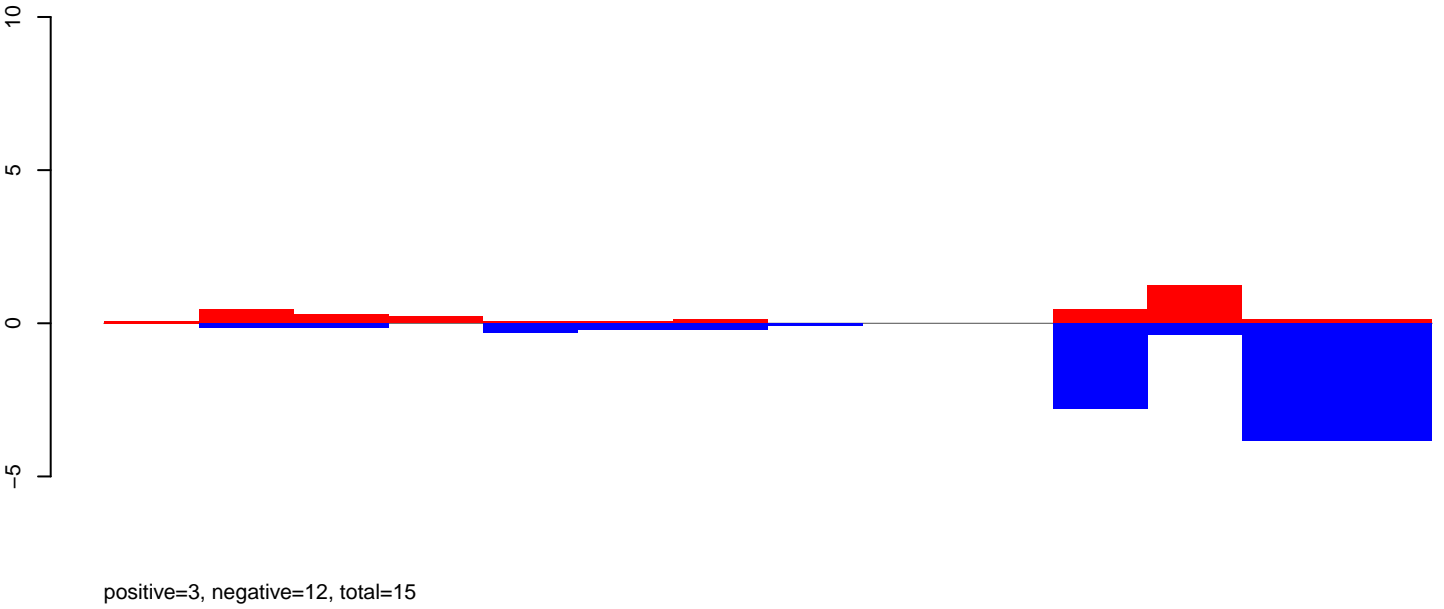


Window size=25, length=450, TE@rnd-5_family-1700@DNA_PIF-Spy@2459:1-450

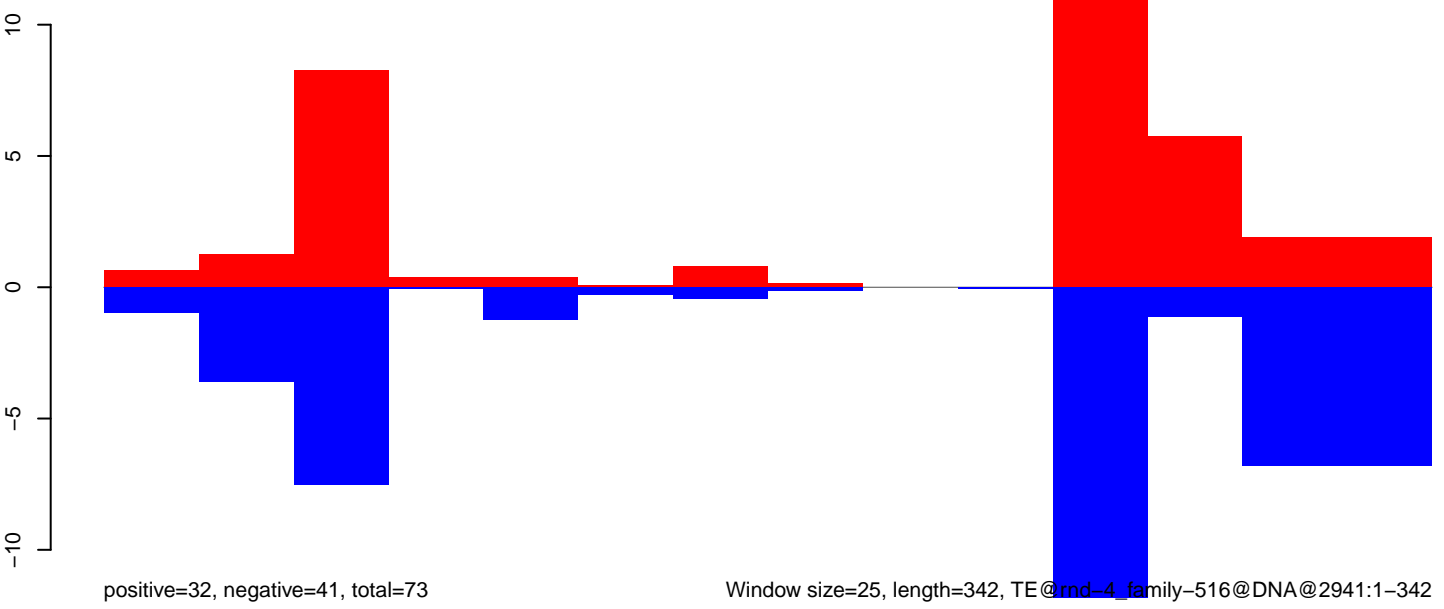
CuQuin_Testes_NL.18_23.rep



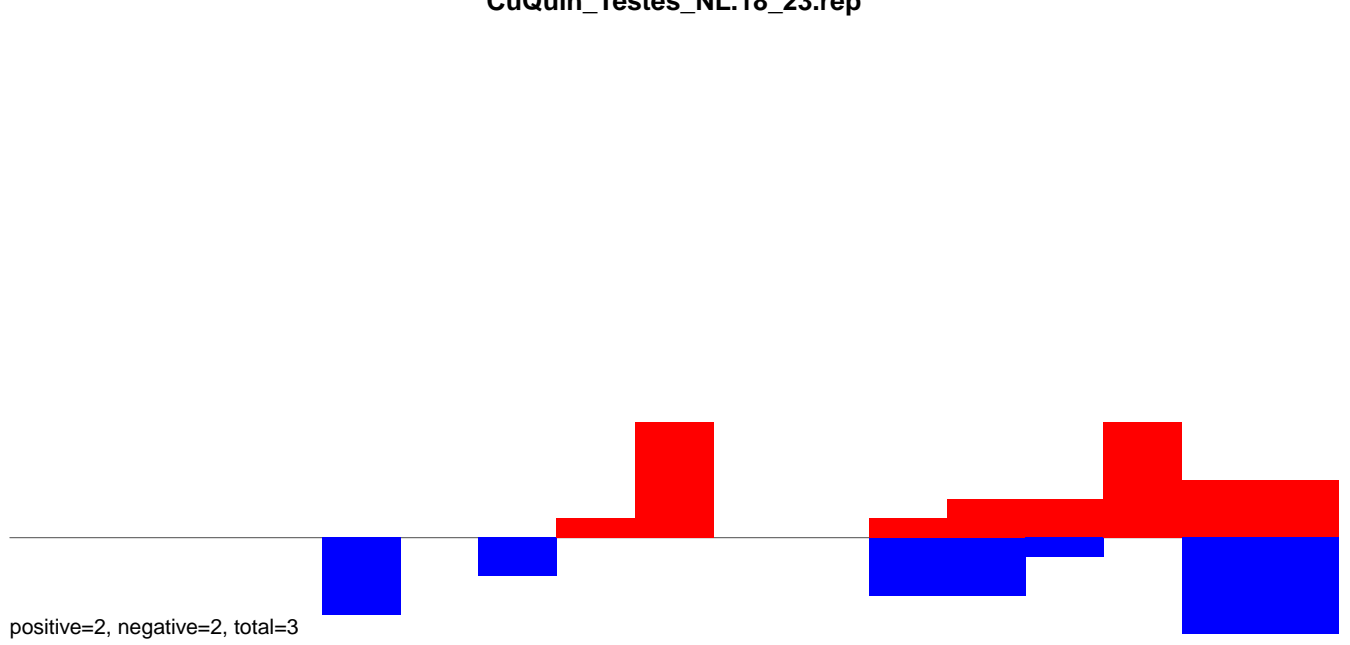
CuQuin_Testes_NL.24_35.rep



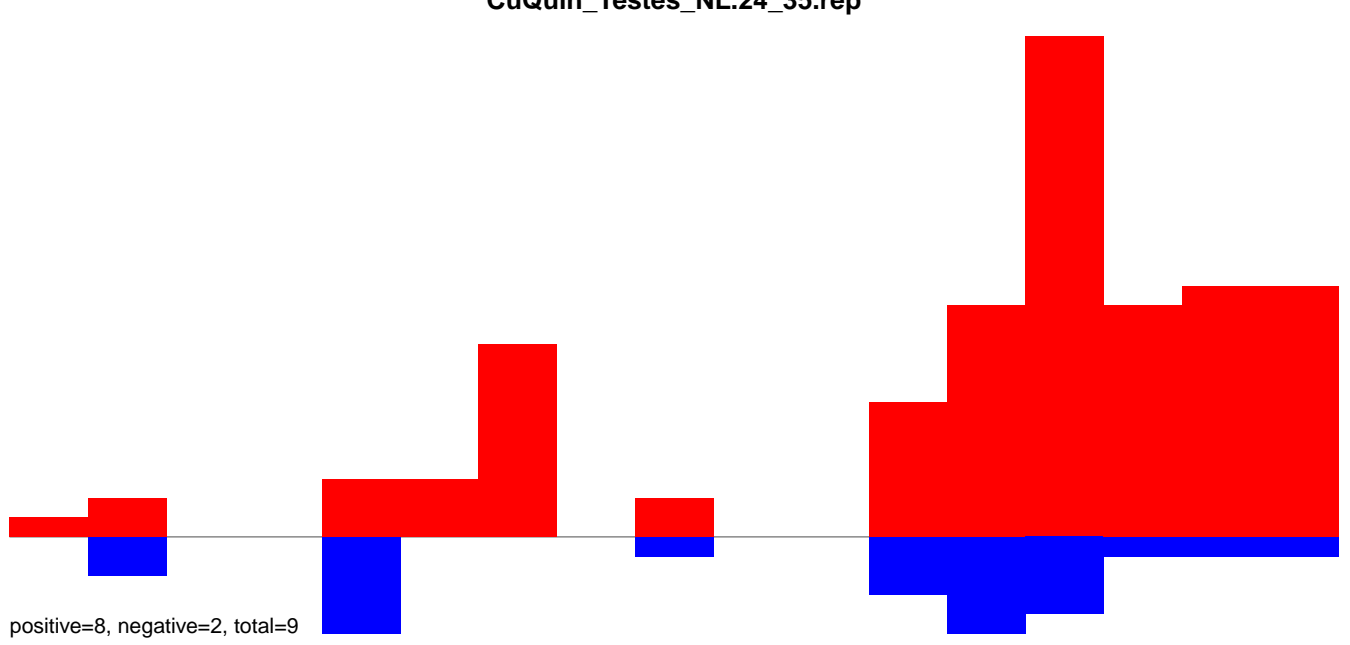
CuQuin_Testes_NL.rep



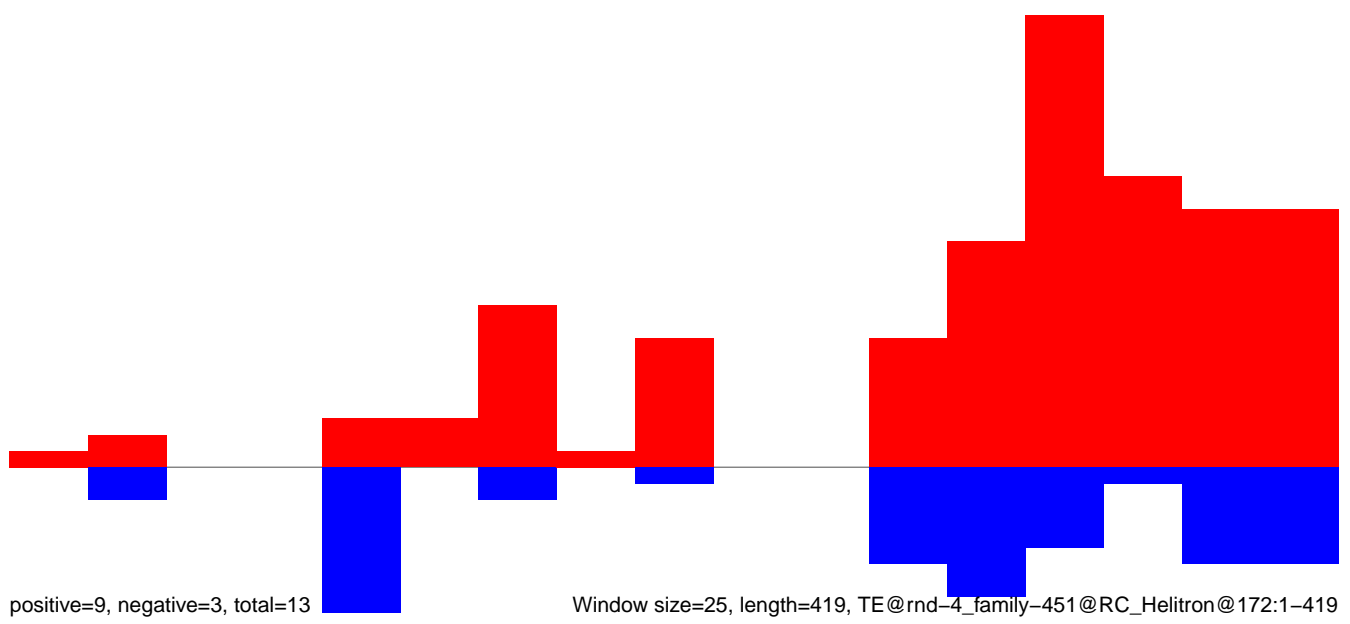
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

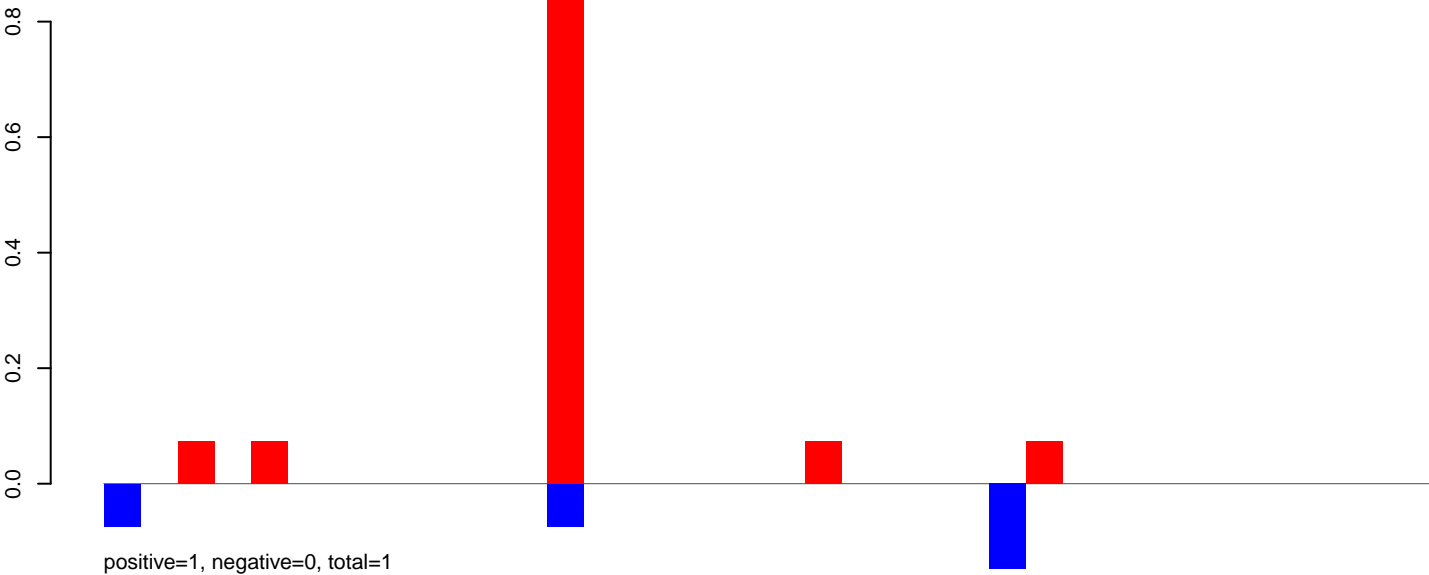


CuQuin_Testes_NL.rep

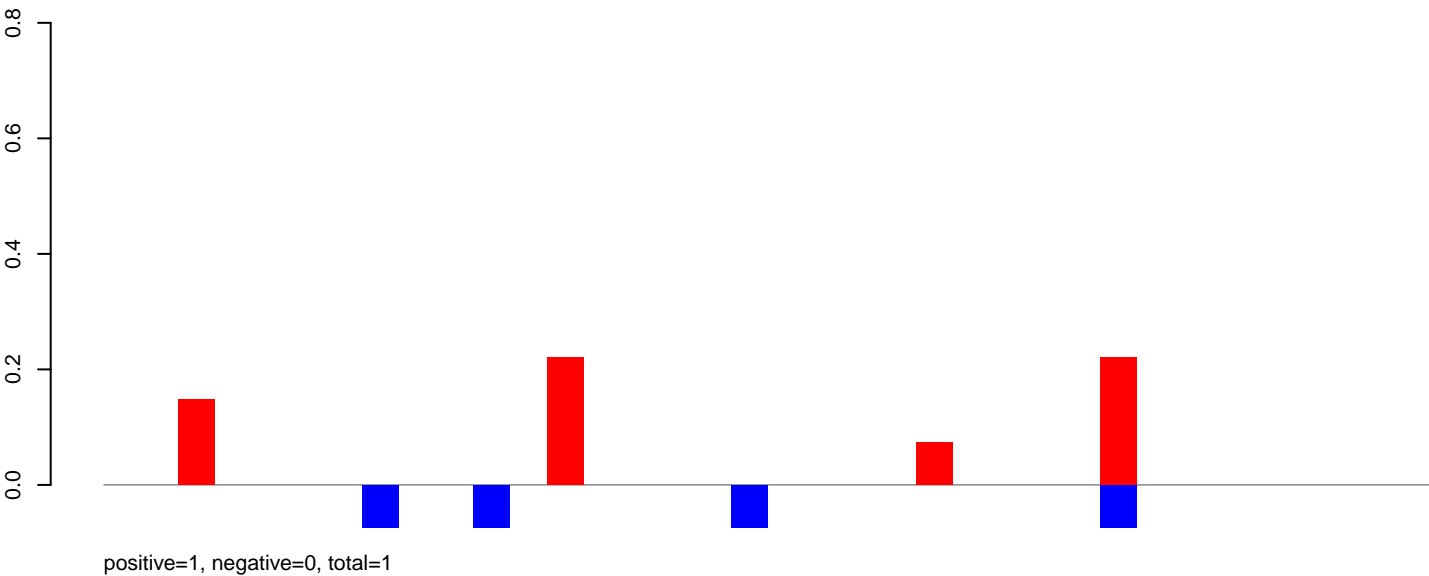


Window size=25, length=419, TE@rnd-4_family-451@RC_Helitron@172:1-419

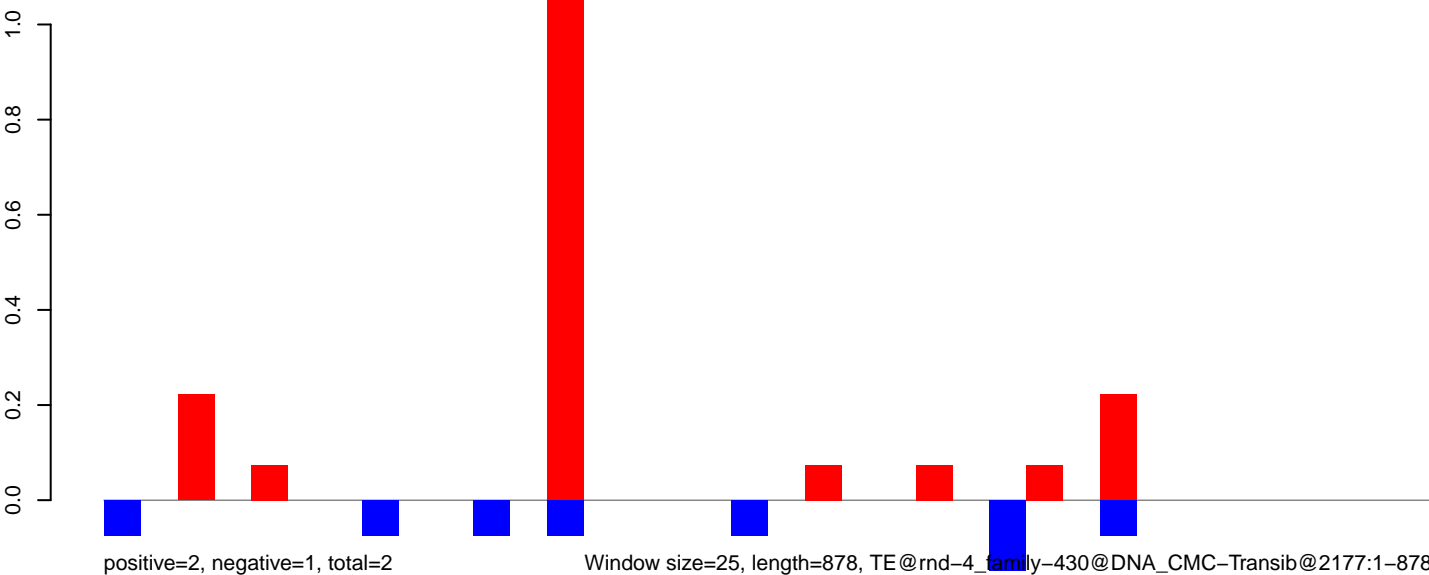
CuQuin_Testes_NL.18_23.rep



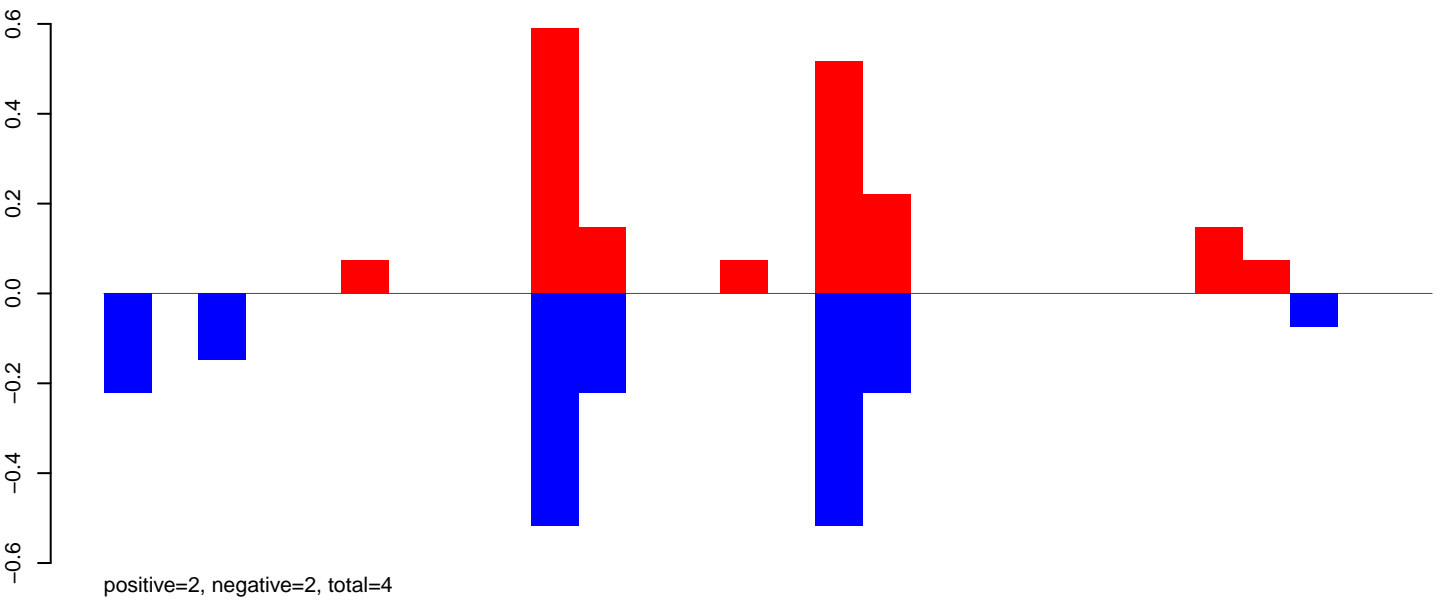
CuQuin_Testes_NL.24_35.rep



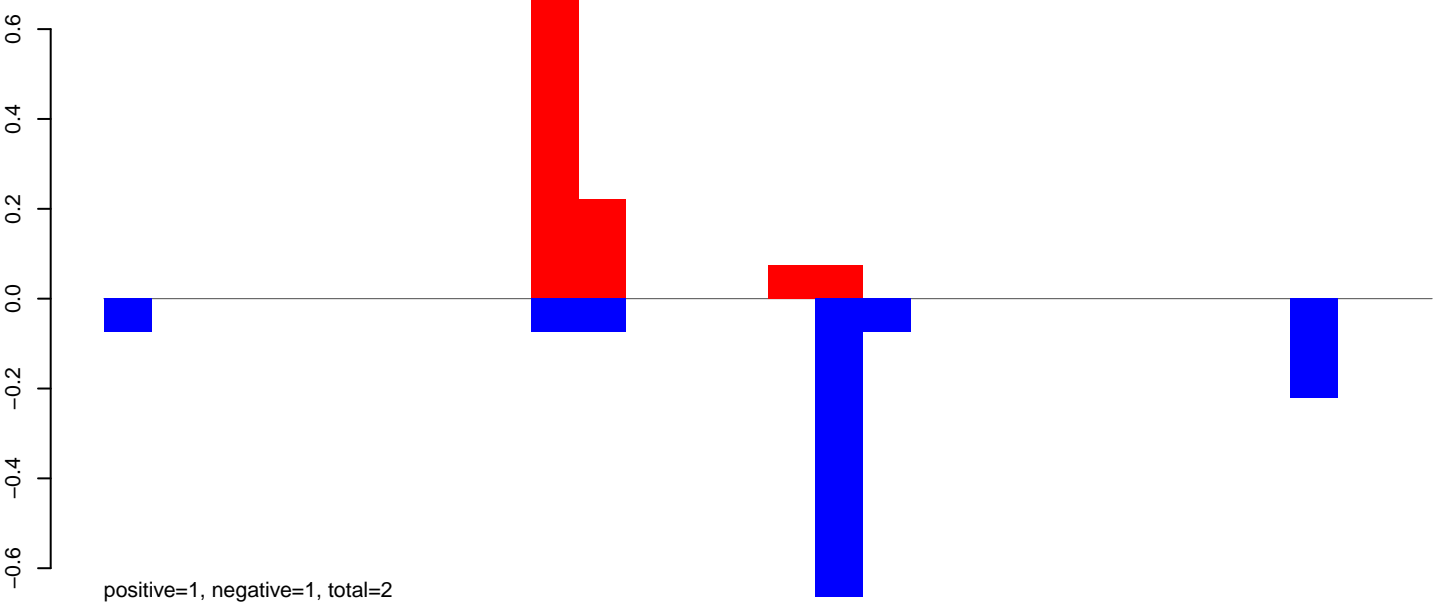
CuQuin_Testes_NL.rep



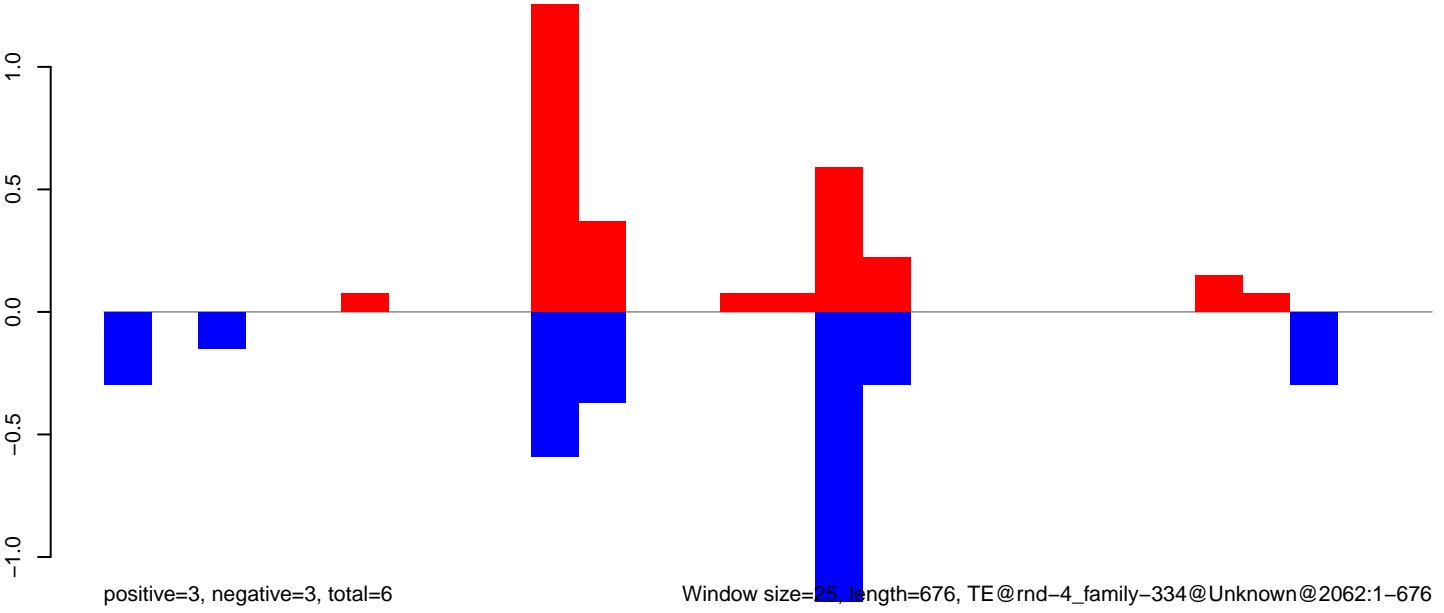
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

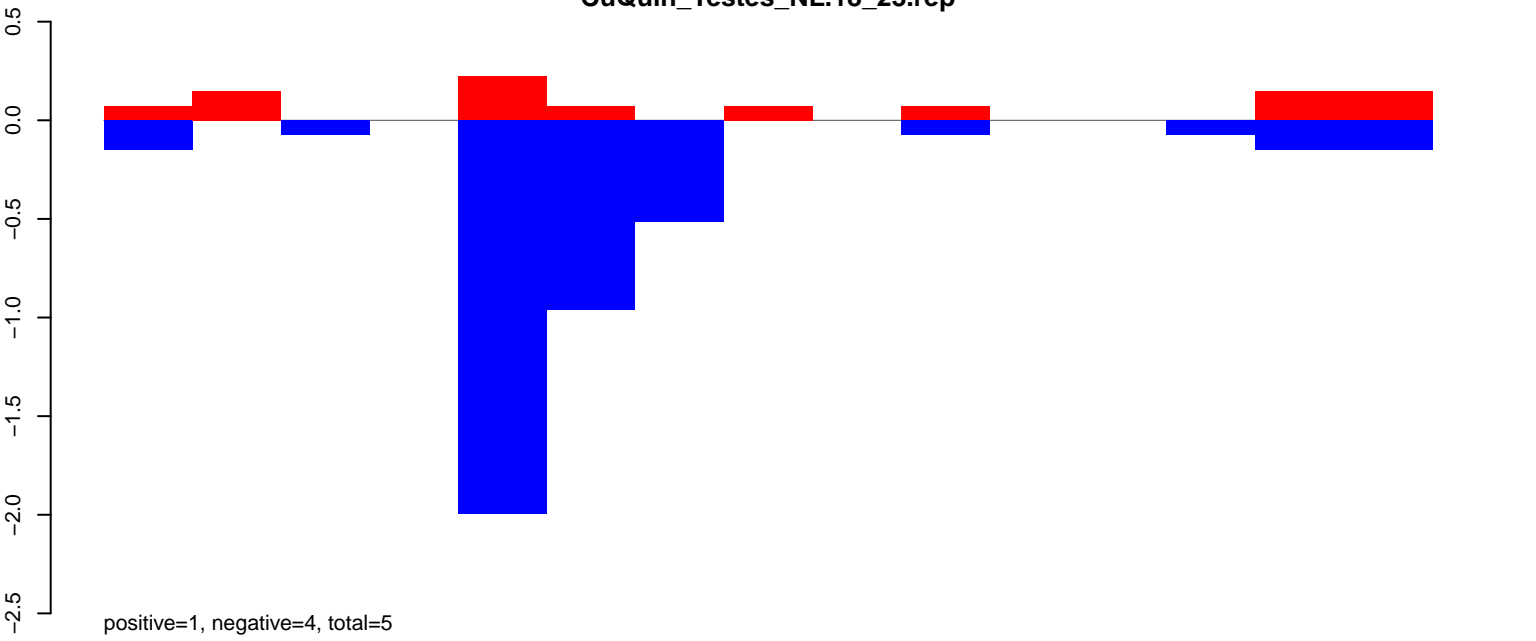


CuQuin_Testes_NL.rep

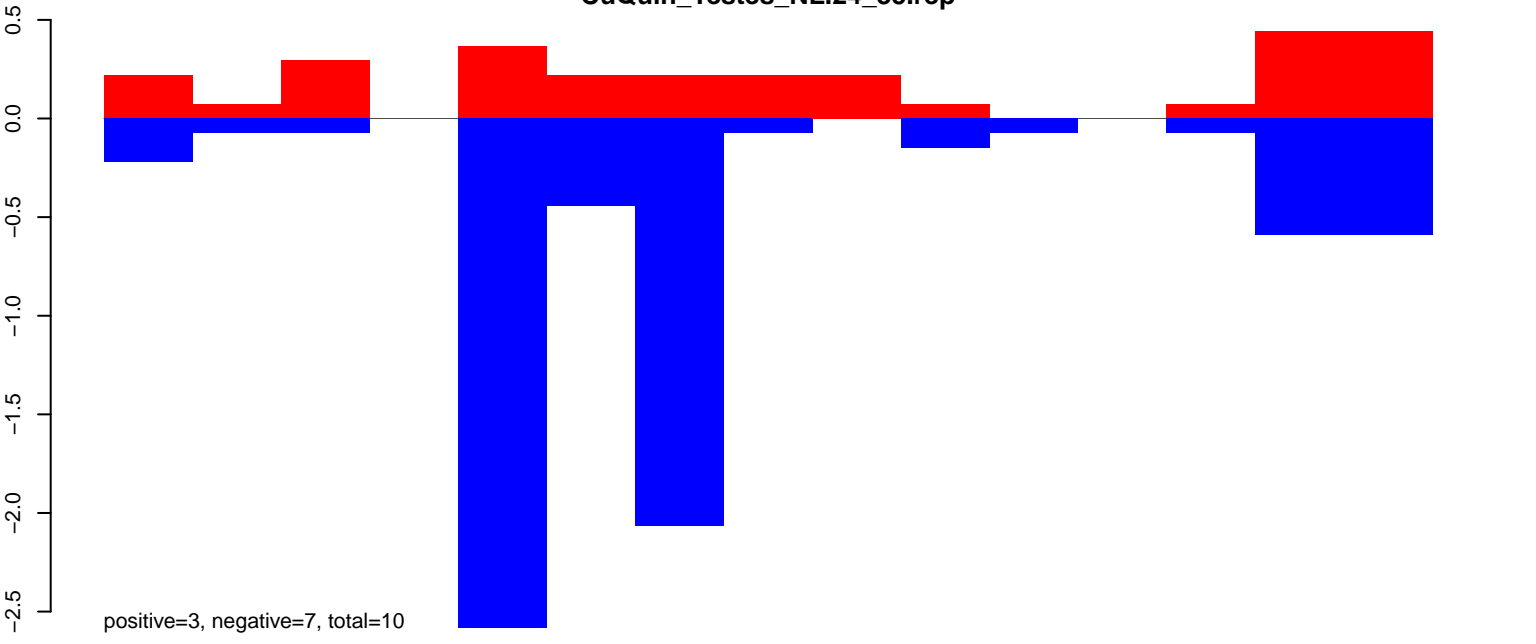


Window size=25, length=676, TE@rnd-4_family-334@Unknown@2062:1-676

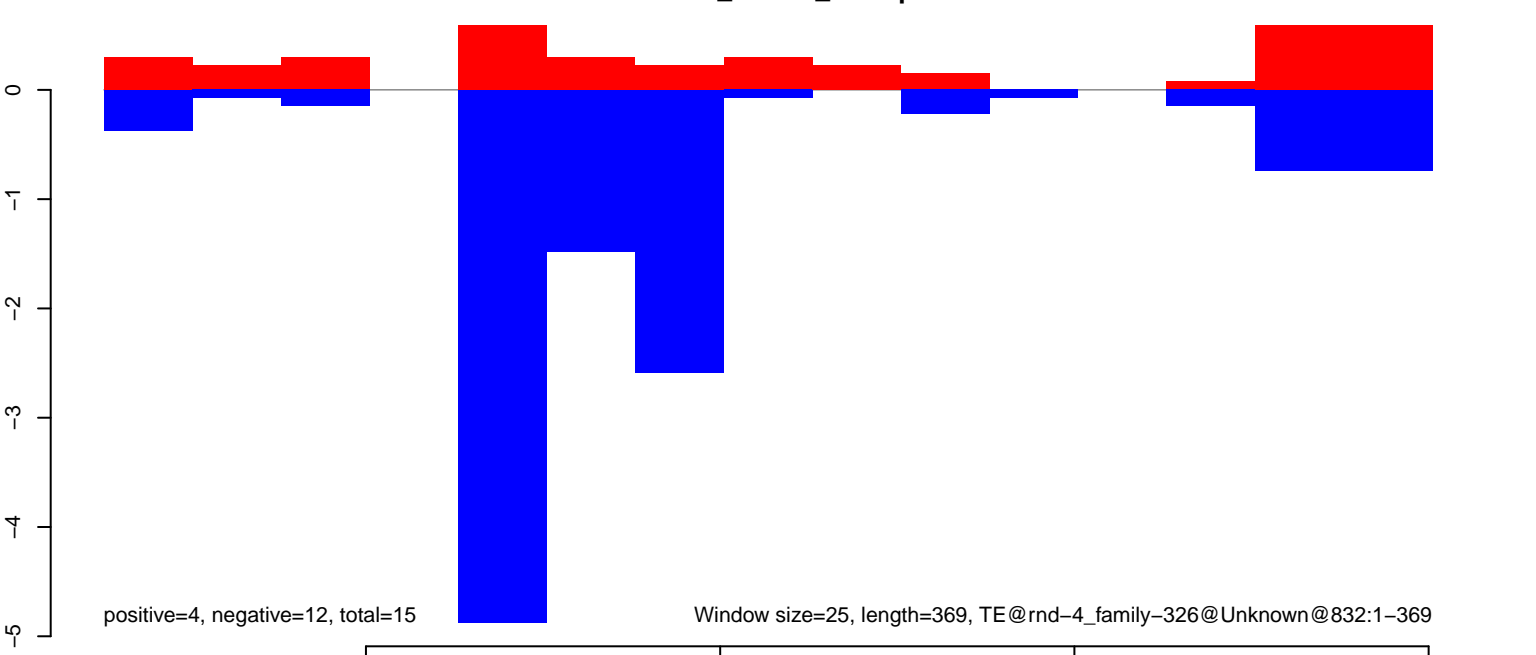
CuQuin_Testes_NL.18_23.rep



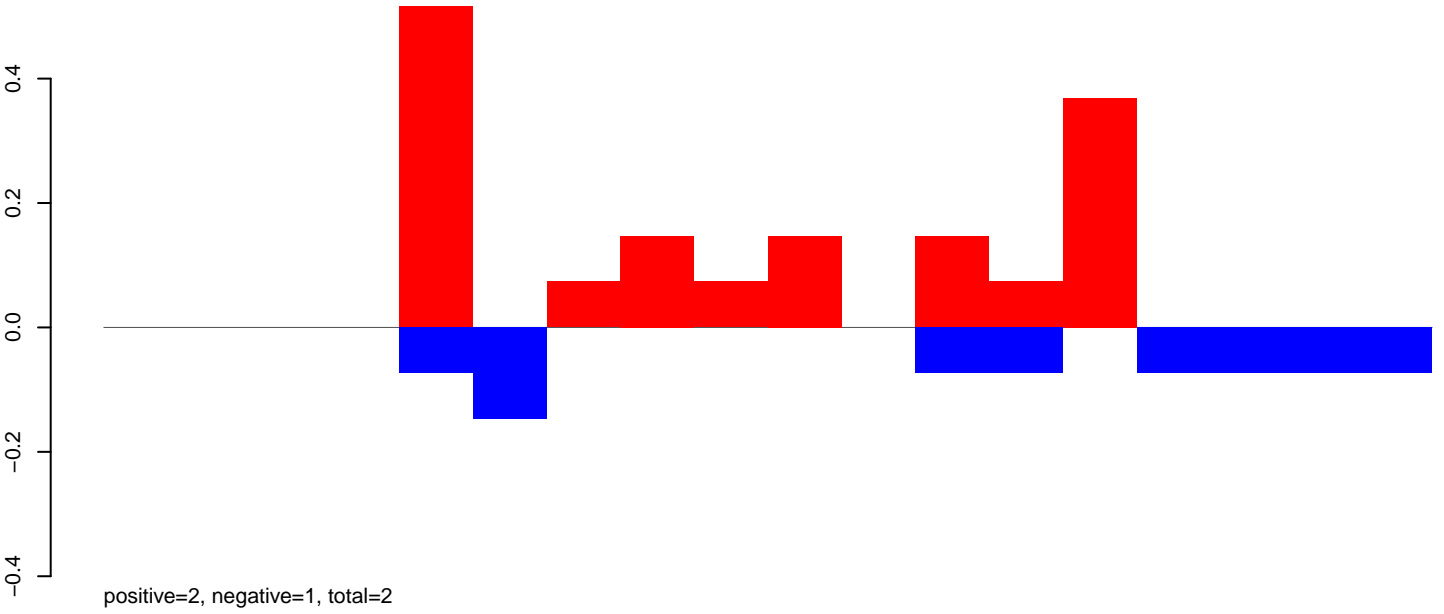
CuQuin_Testes_NL.24_35.rep



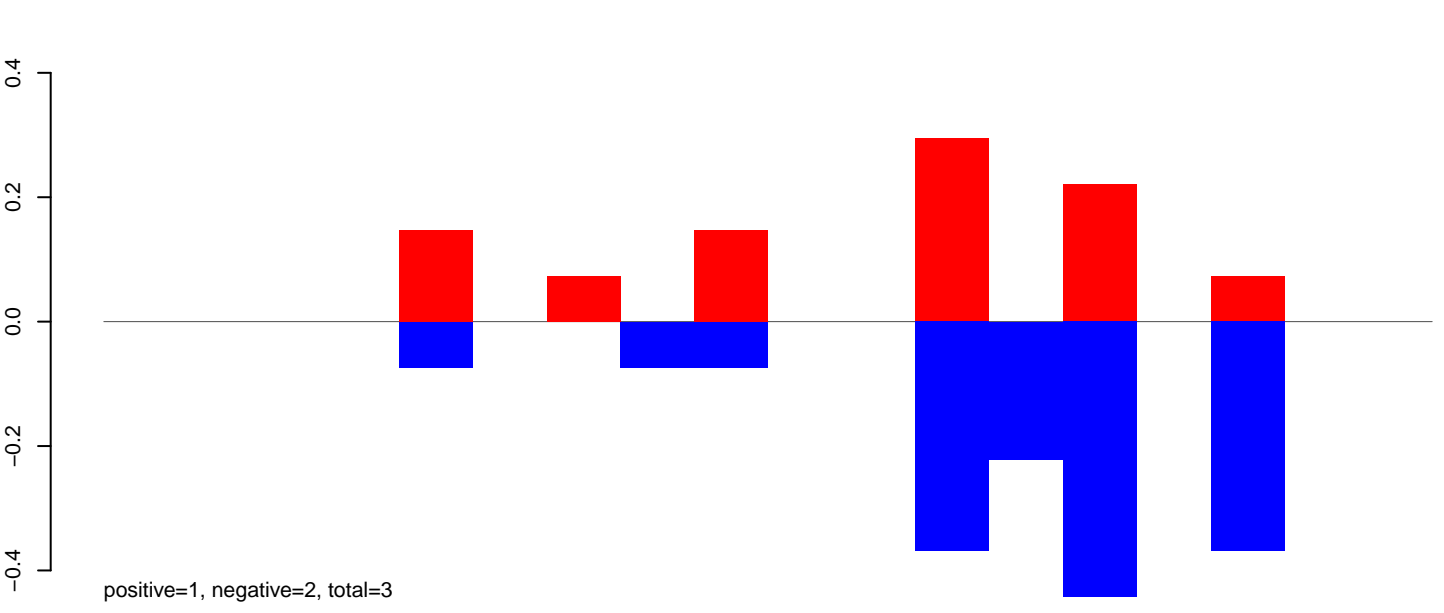
CuQuin_Testes_NL.rep



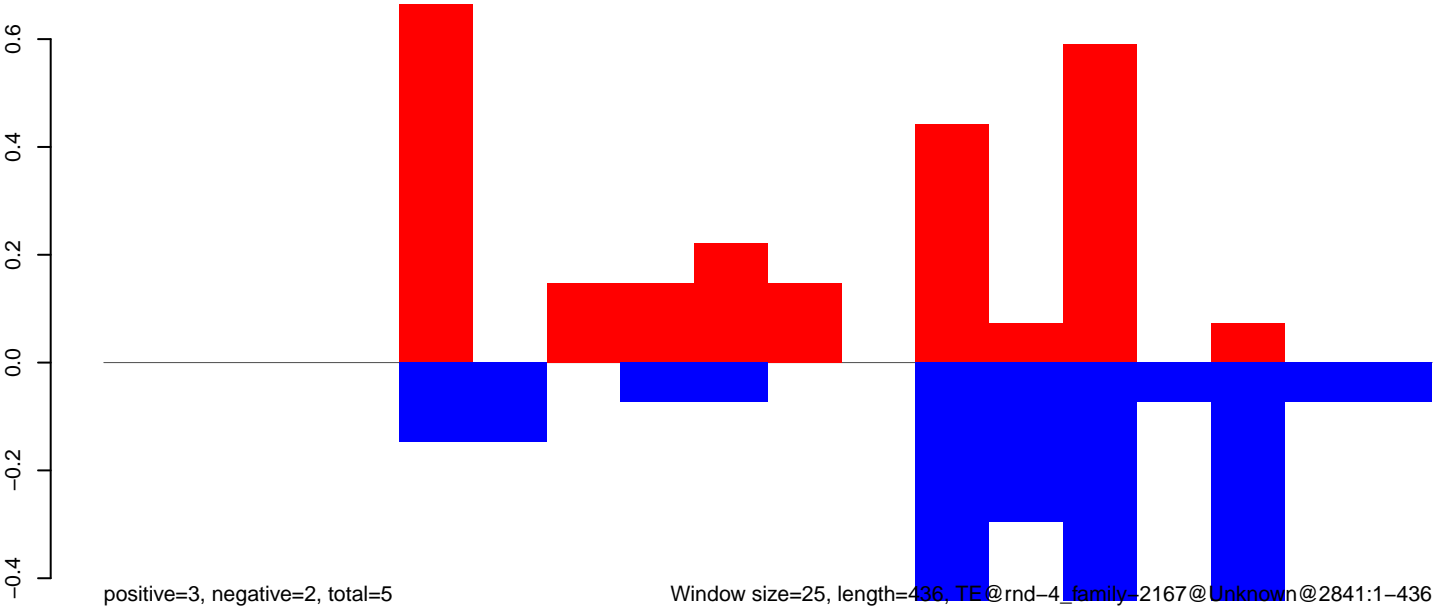
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

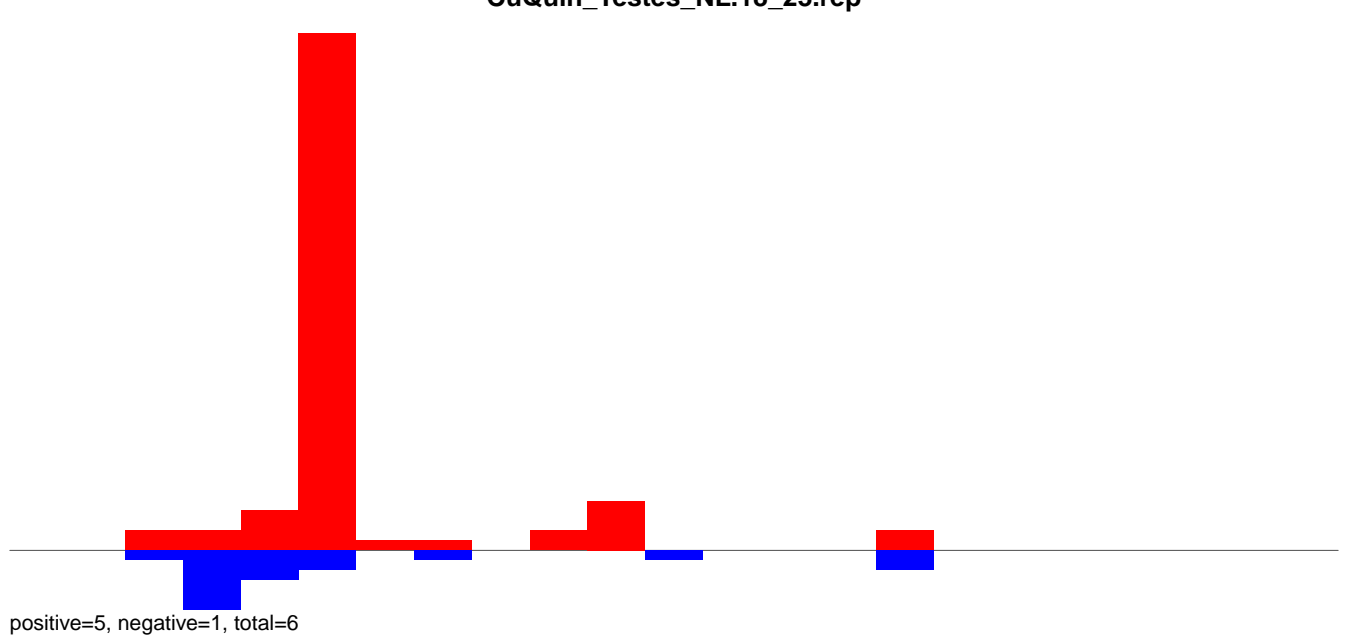


CuQuin_Testes_NL.rep

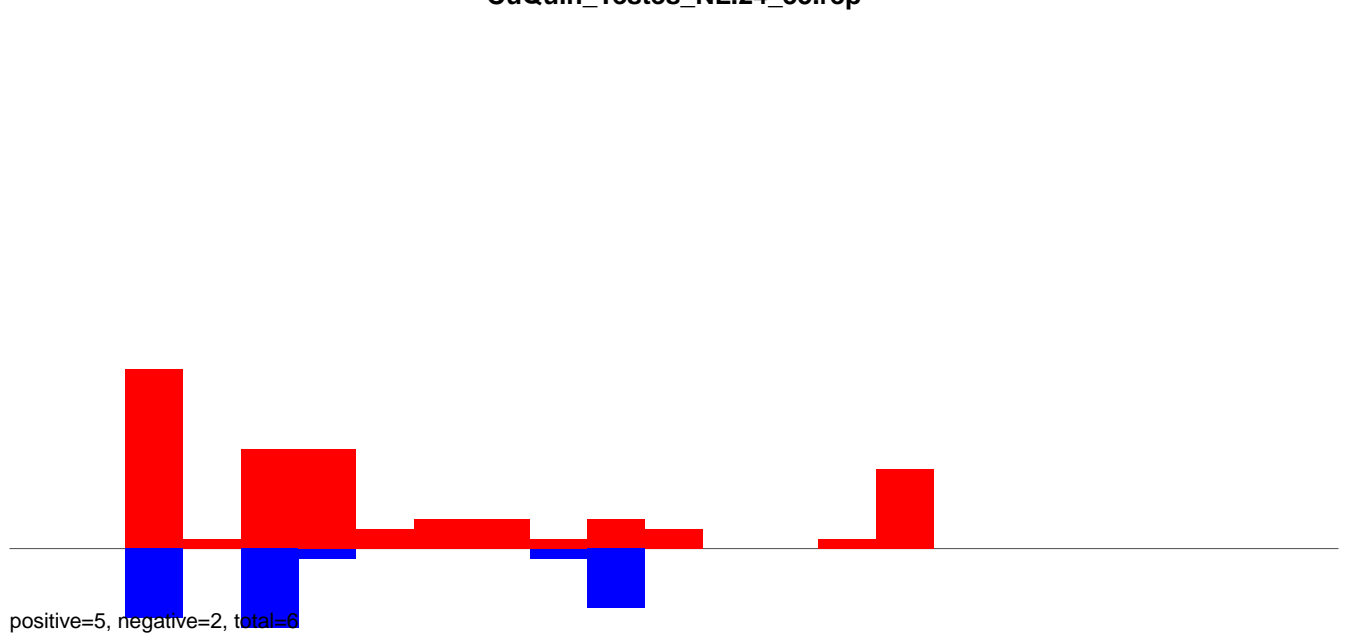


Window size=25, length=436, TE@rnd-4_family-2167@Unknown@2841:1-436

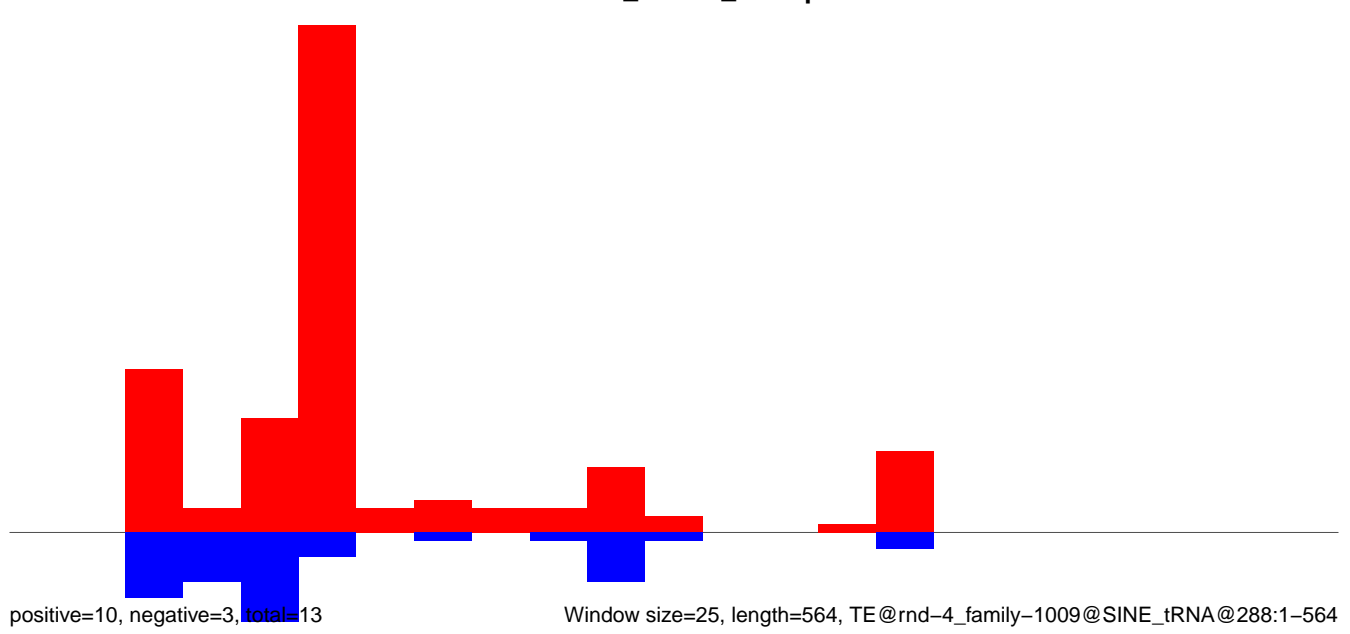
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



CuQuin_Testes_NL.18_23.rep



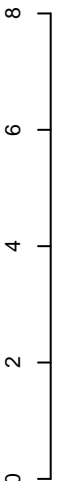
positive=3, negative=1, total=5

CuQuin_Testes_NL.24_35.rep



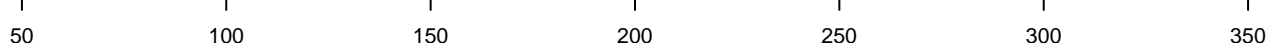
positive=11, negative=3, total=14

CuQuin_Testes_NL.rep

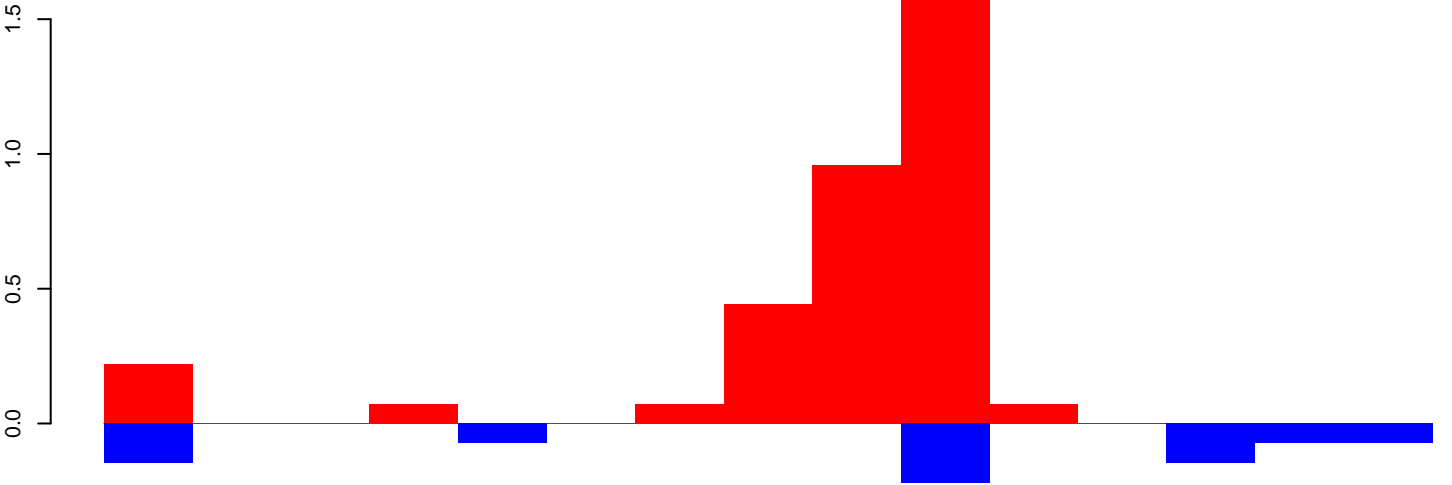


positive=19, negative=4, total=23

Window size=25, length=315, TE@rmd-3_family-74@LTR_Gypsy@697:1-315

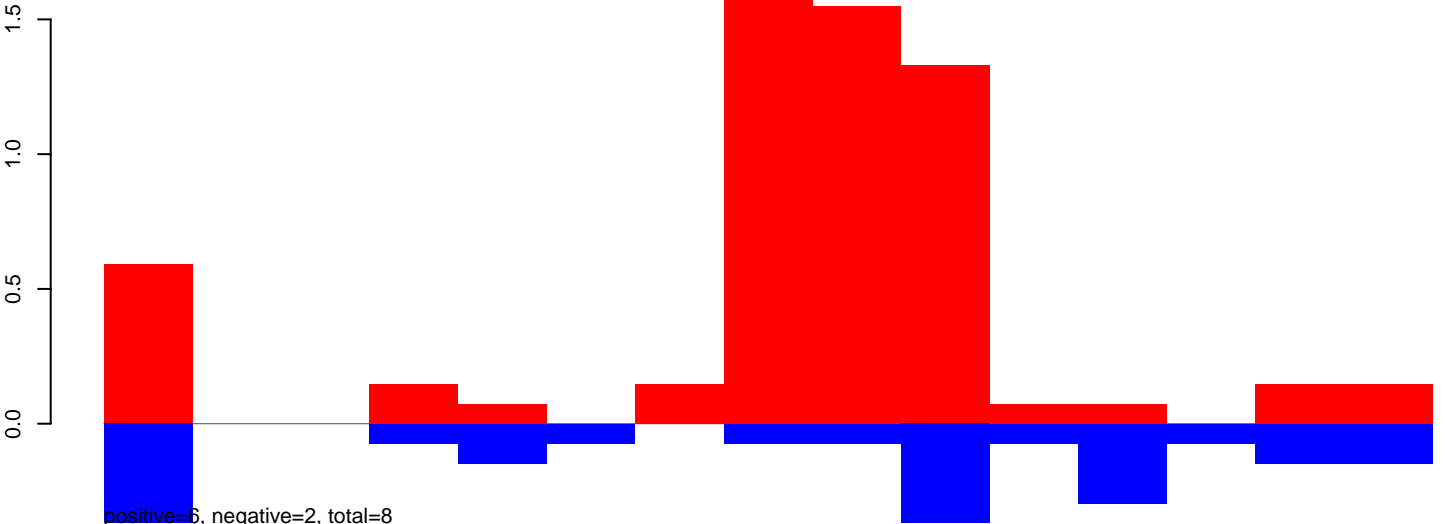


CuQuin_Testes_NL.18_23.rep



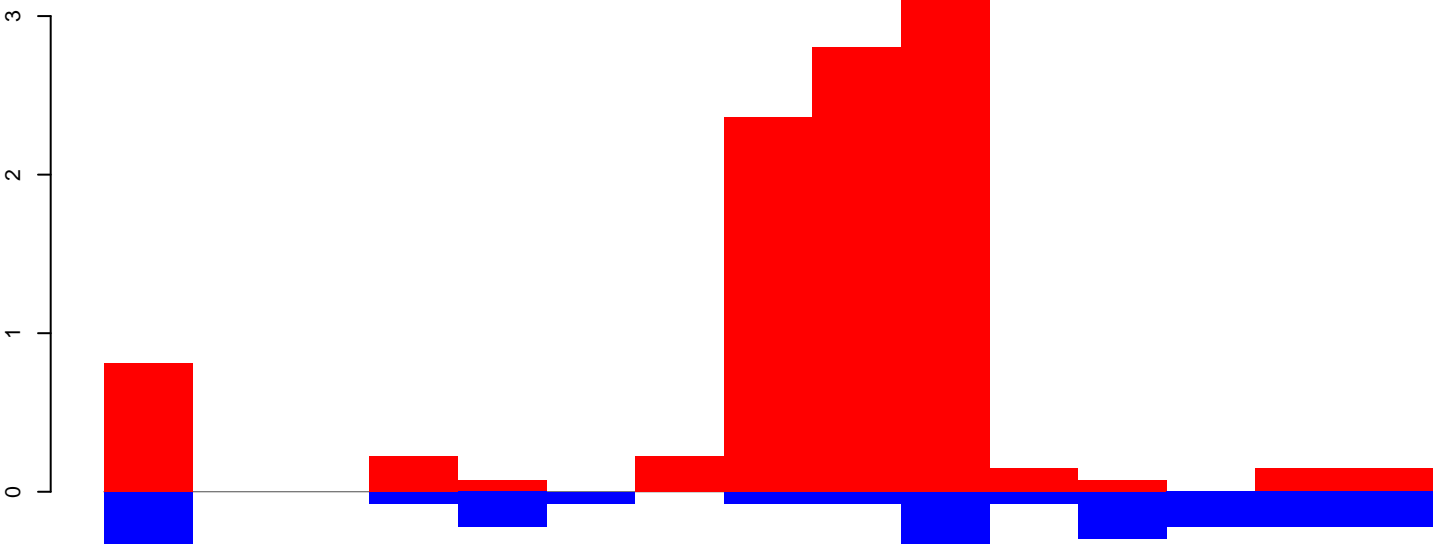
positive=4, negative=1, total=4

CuQuin_Testes_NL.24_35.rep



positive=6, negative=2, total=8

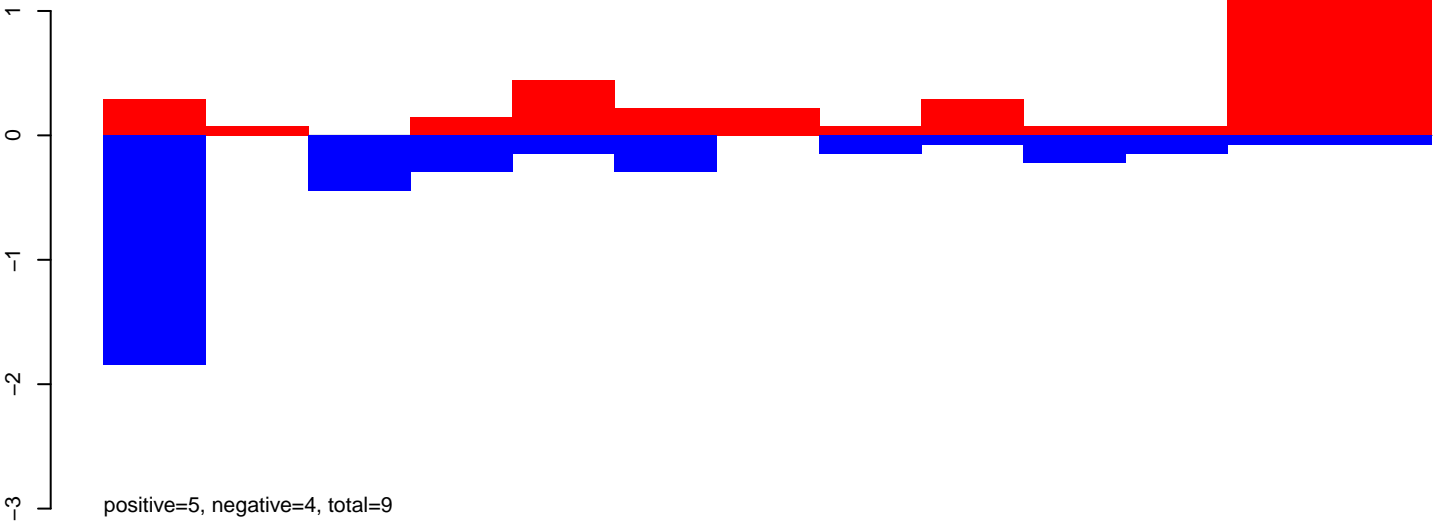
CuQuin_Testes_NL.rep



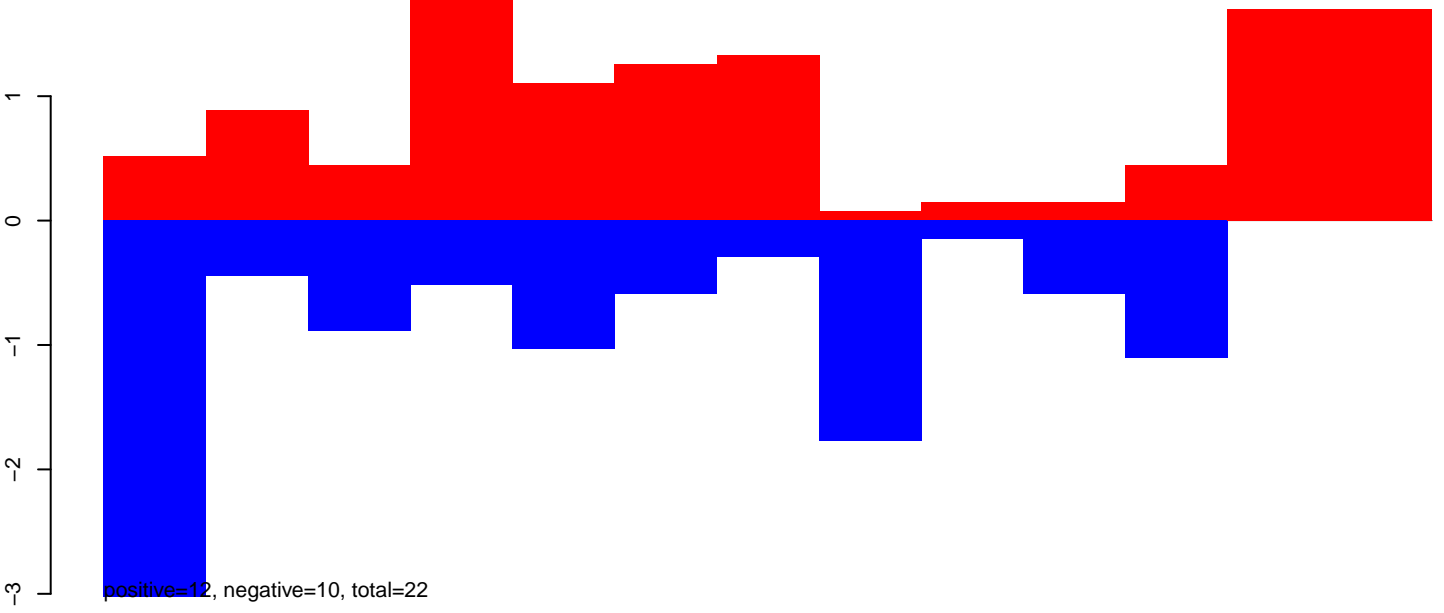
positive=10, negative=3, total=13

Window size=25, length=364, TE@rnd-3_family-749@RC_Helitron@653:1-364

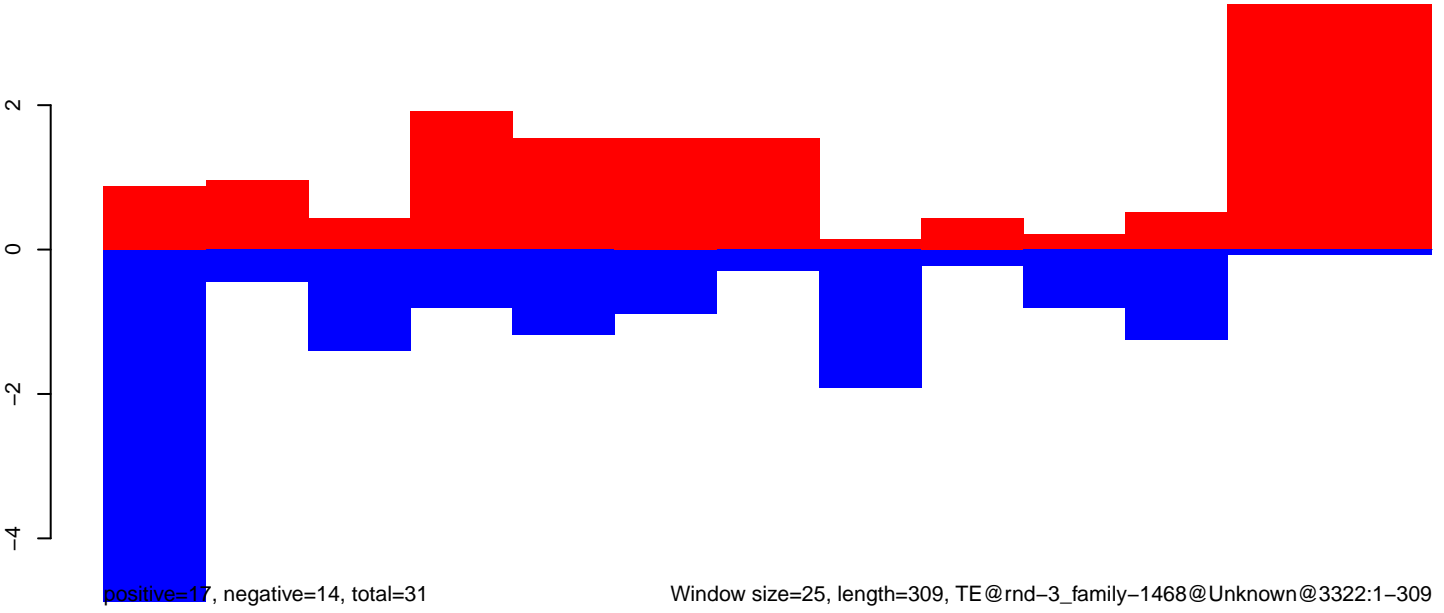
CuQuin_Testes_NL.18_23.rep



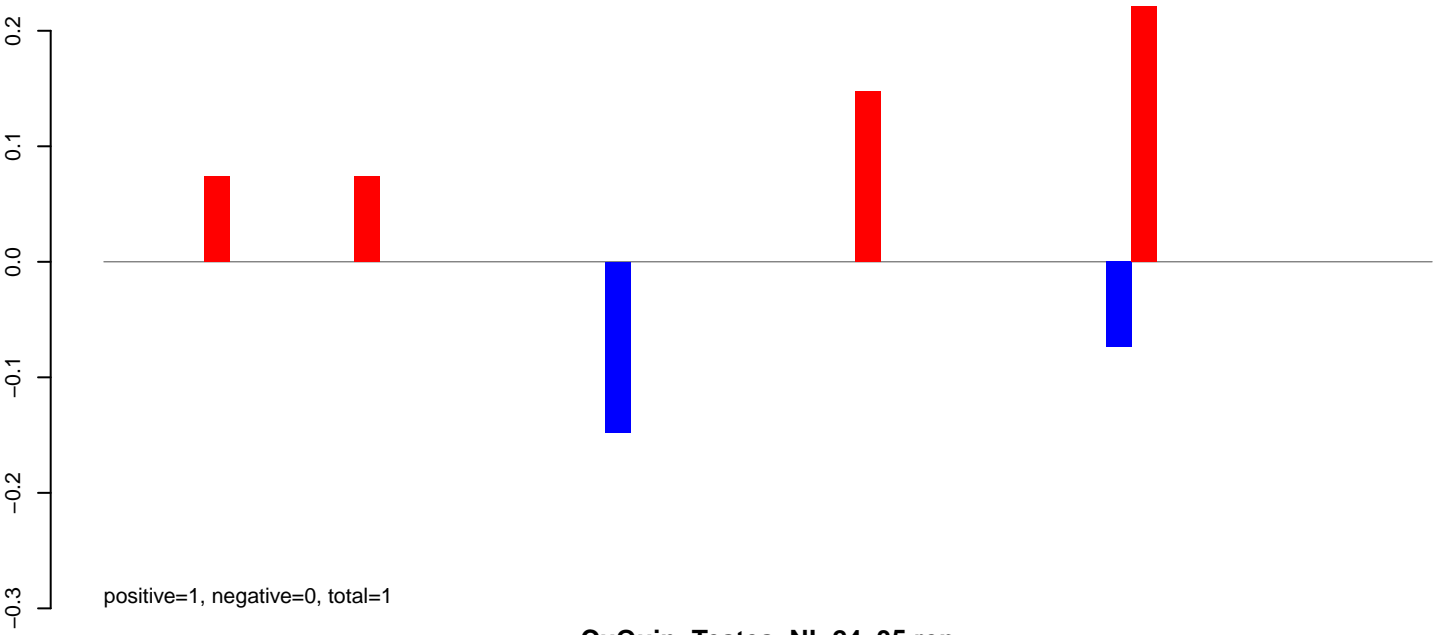
CuQuin_Testes_NL.24_35.rep



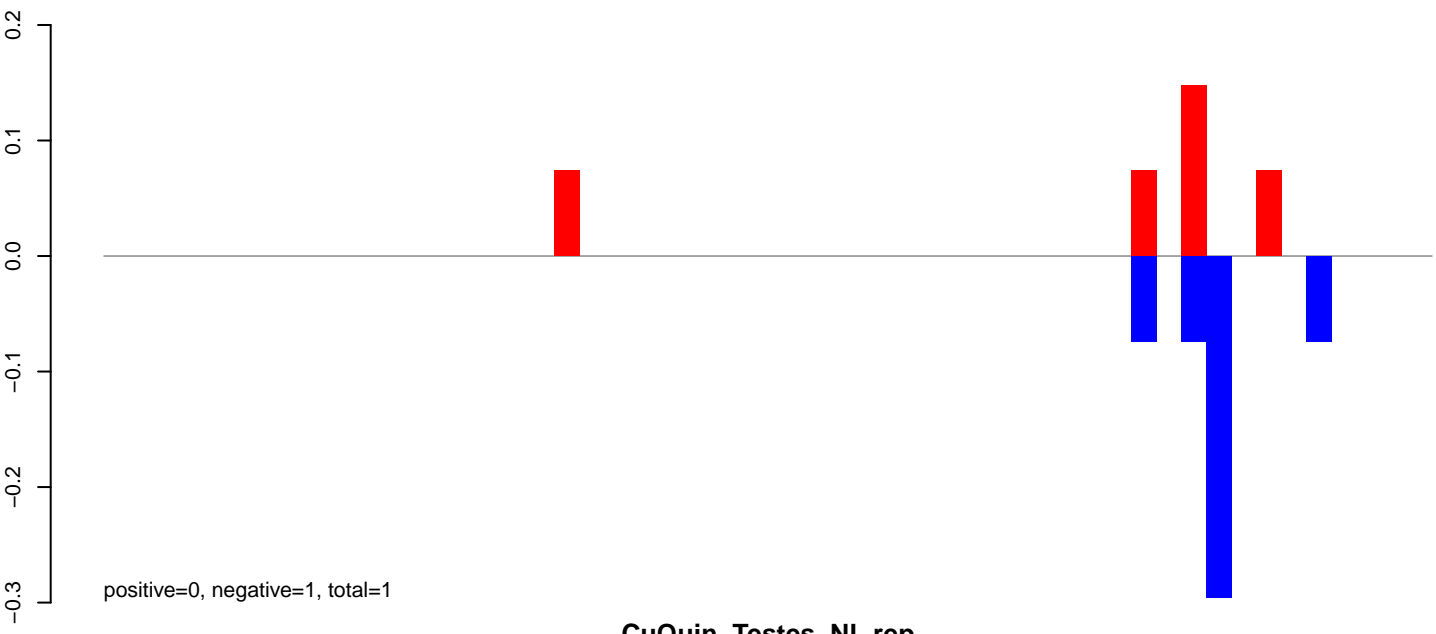
CuQuin_Testes_NL.rep



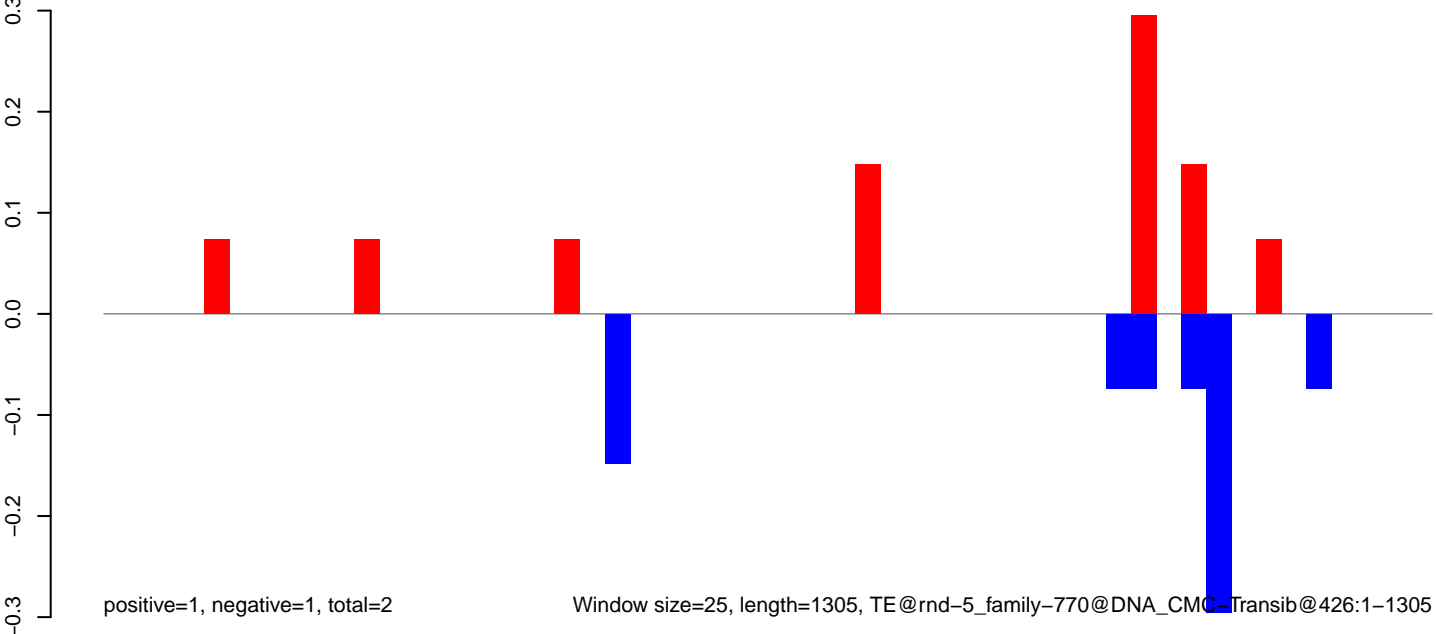
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



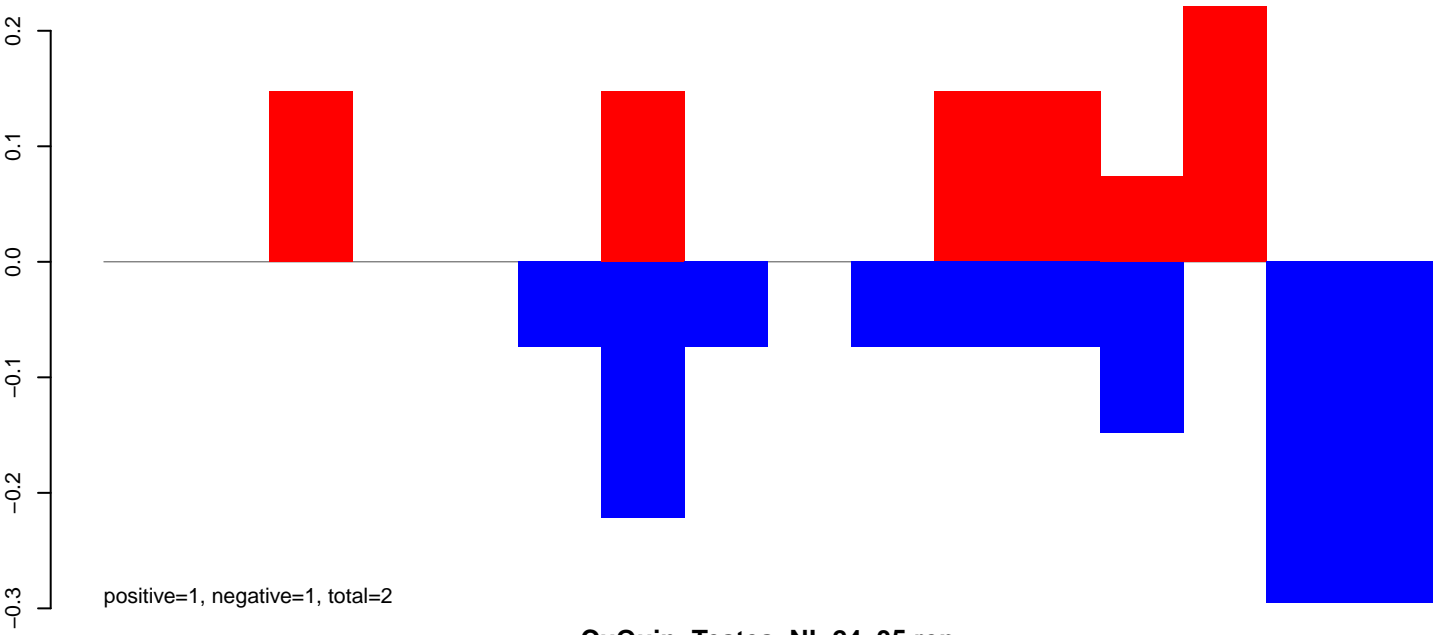
CuQuin_Testes_NL.rep



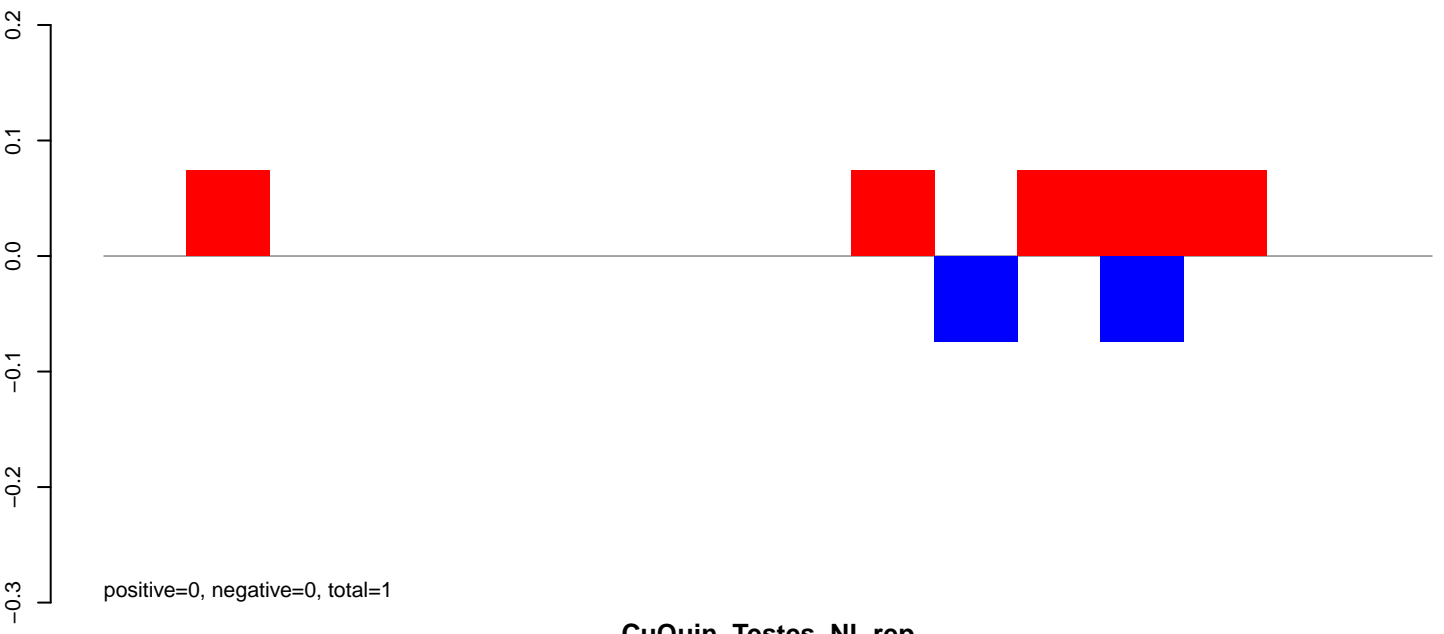
Window size=25, length=1305, TE@rnd-5_family-770@DNA_CMC_Transib@426:1-1305

0 200 400 600 800 1000 1200 1400

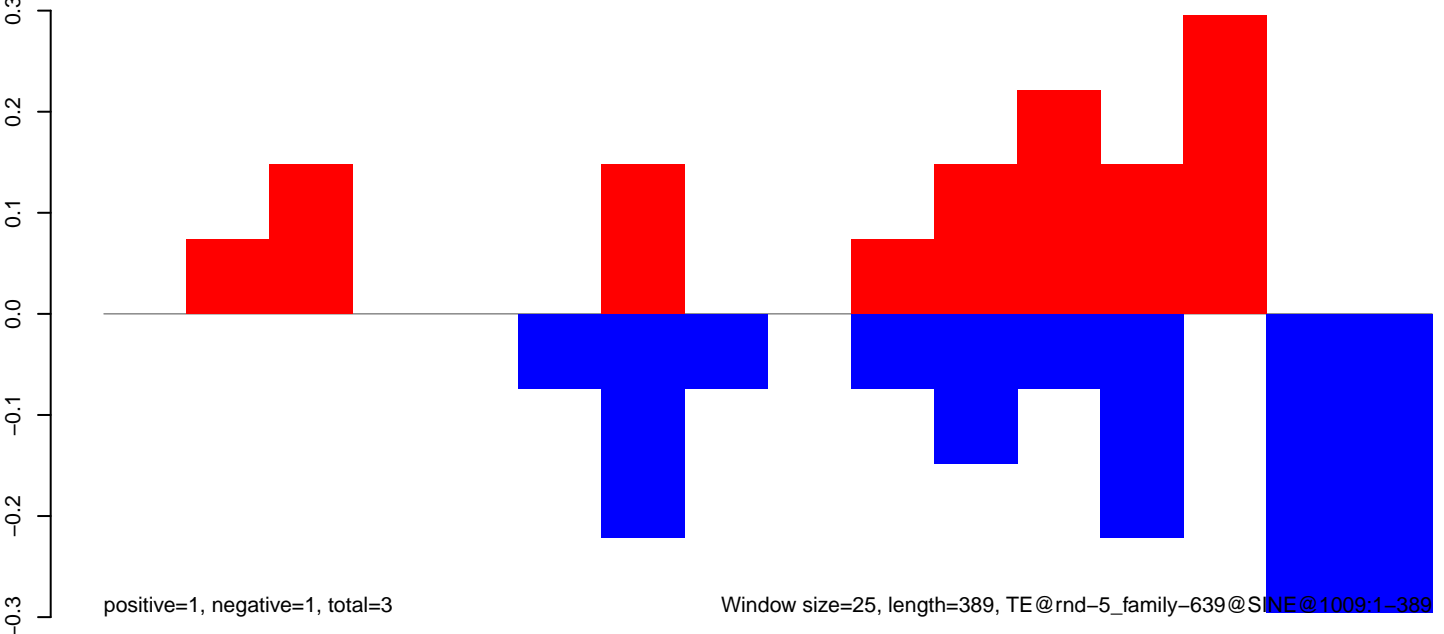
CuQuin_Testes_NL.18_23.rep



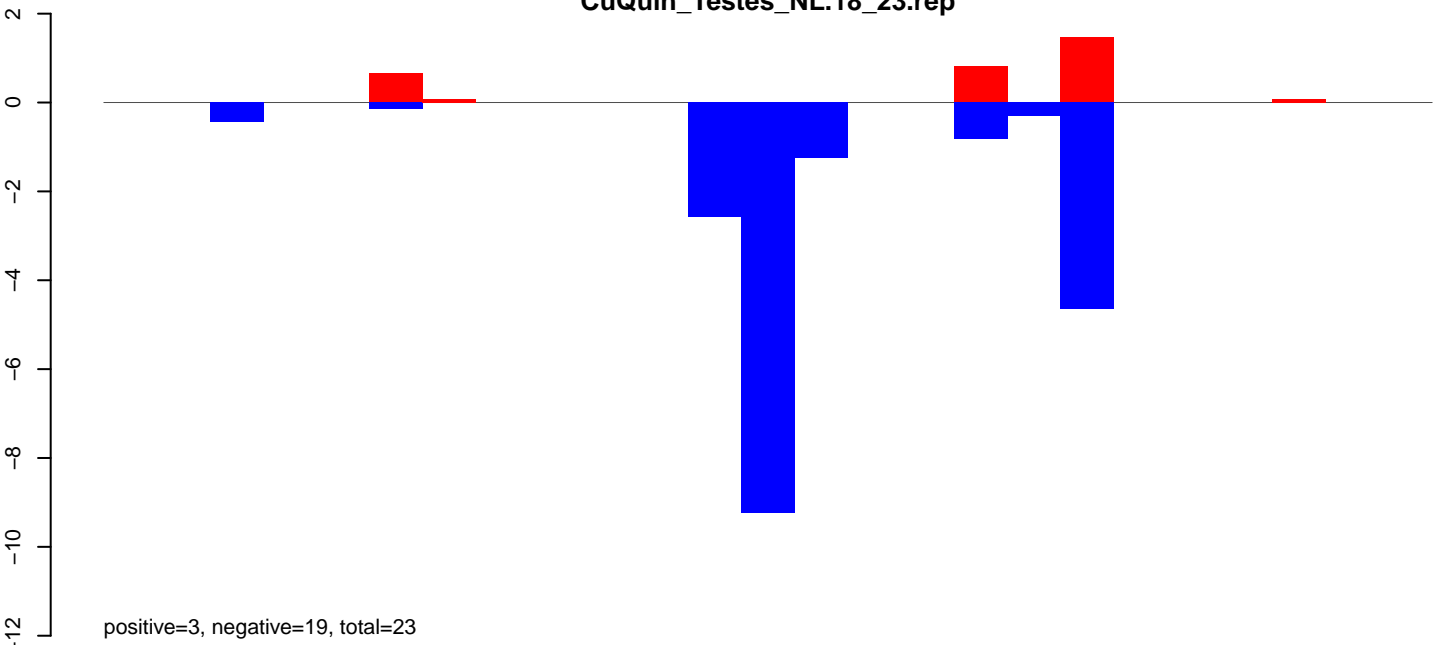
CuQuin_Testes_NL.24_35.rep



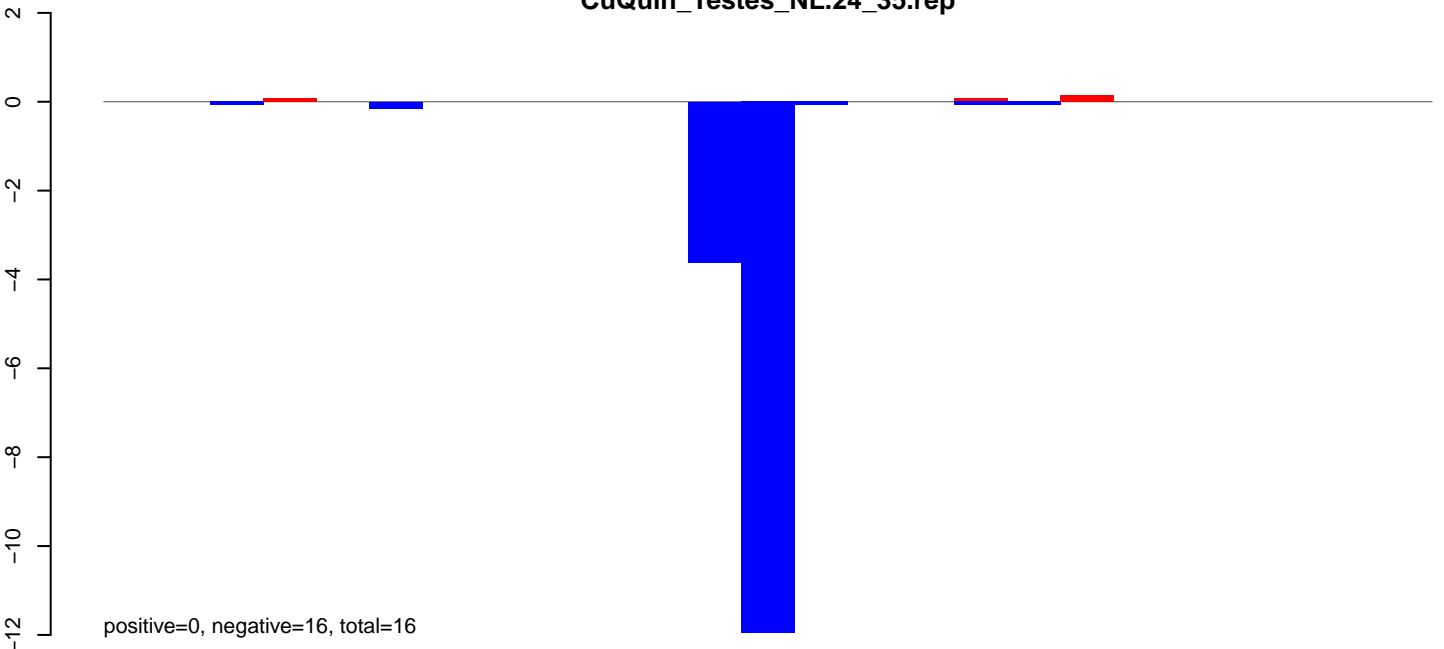
CuQuin_Testes_NL.rep



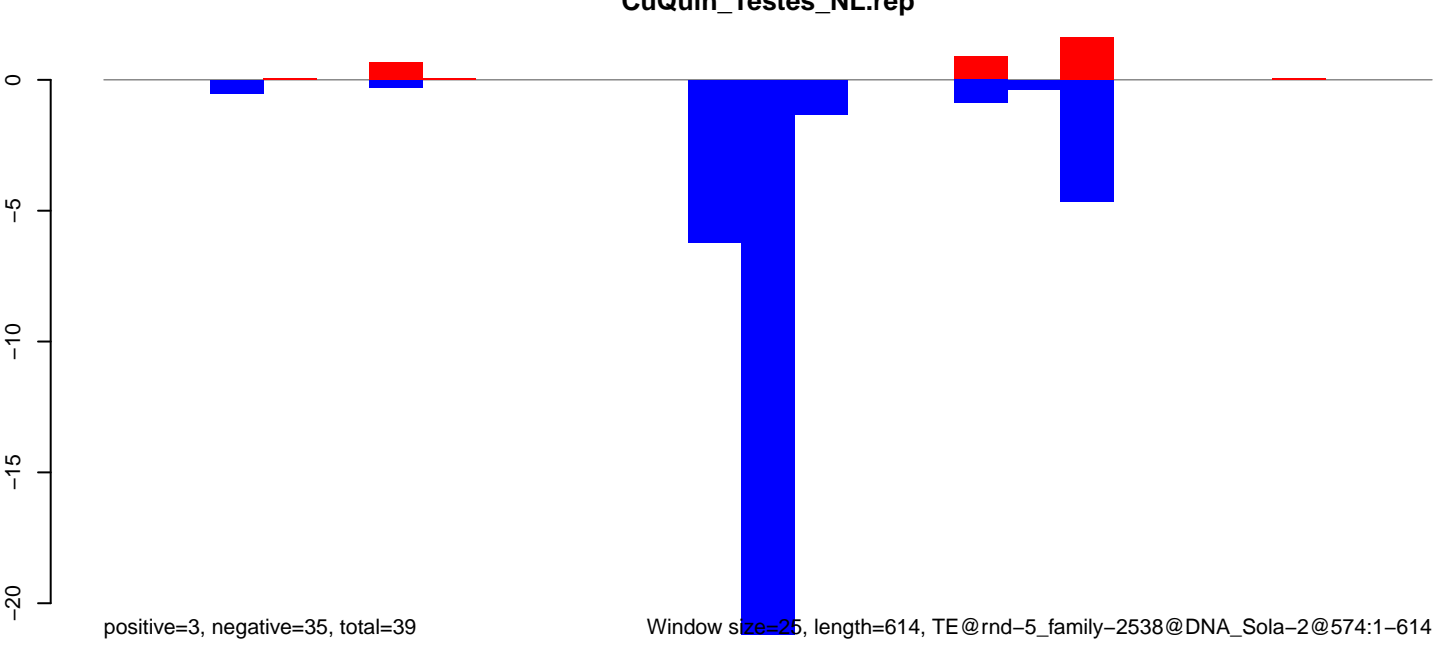
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

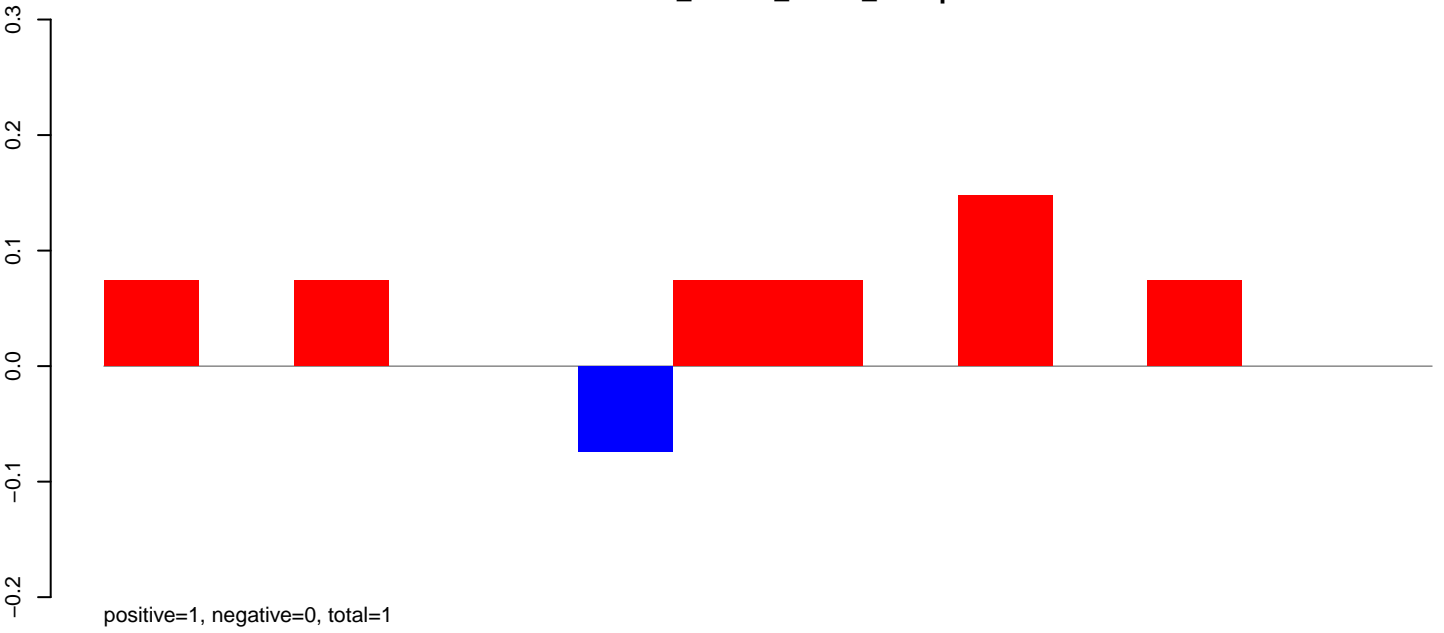


CuQuin_Testes_NL.rep

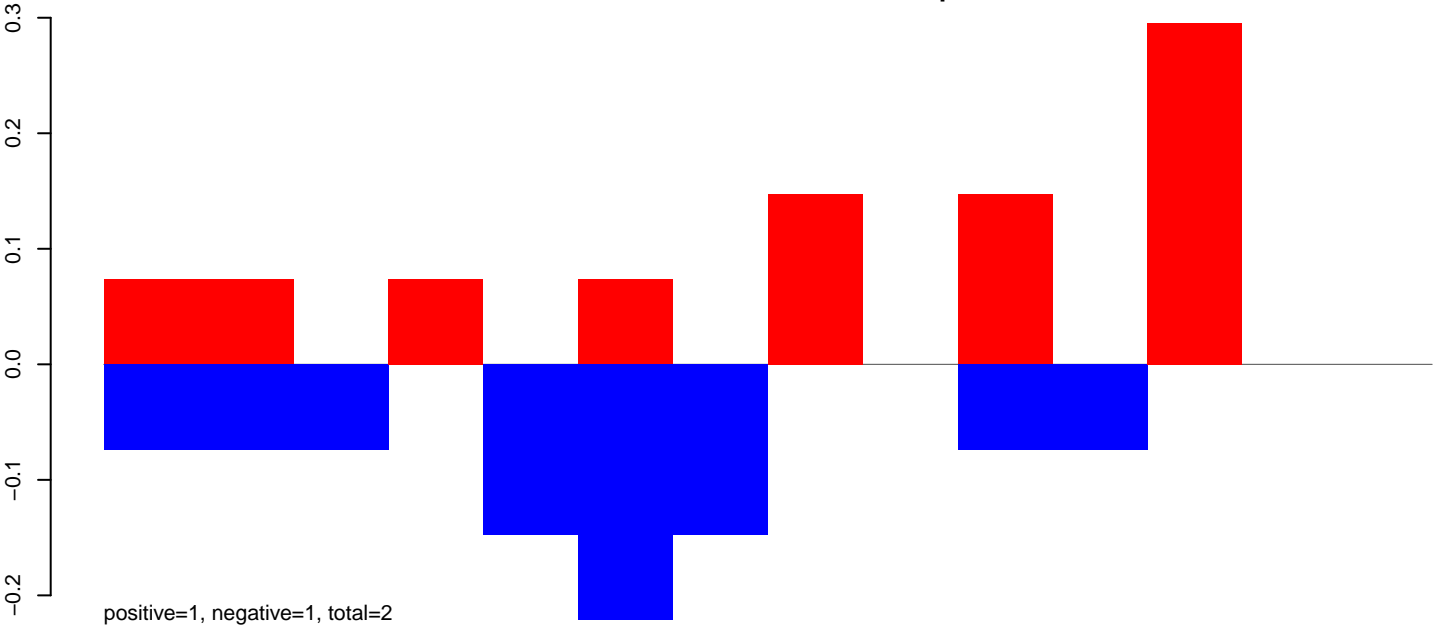


Window size=25, length=614, TE@rnd-5_family-2538@DNA_Sola-2@574:1-614

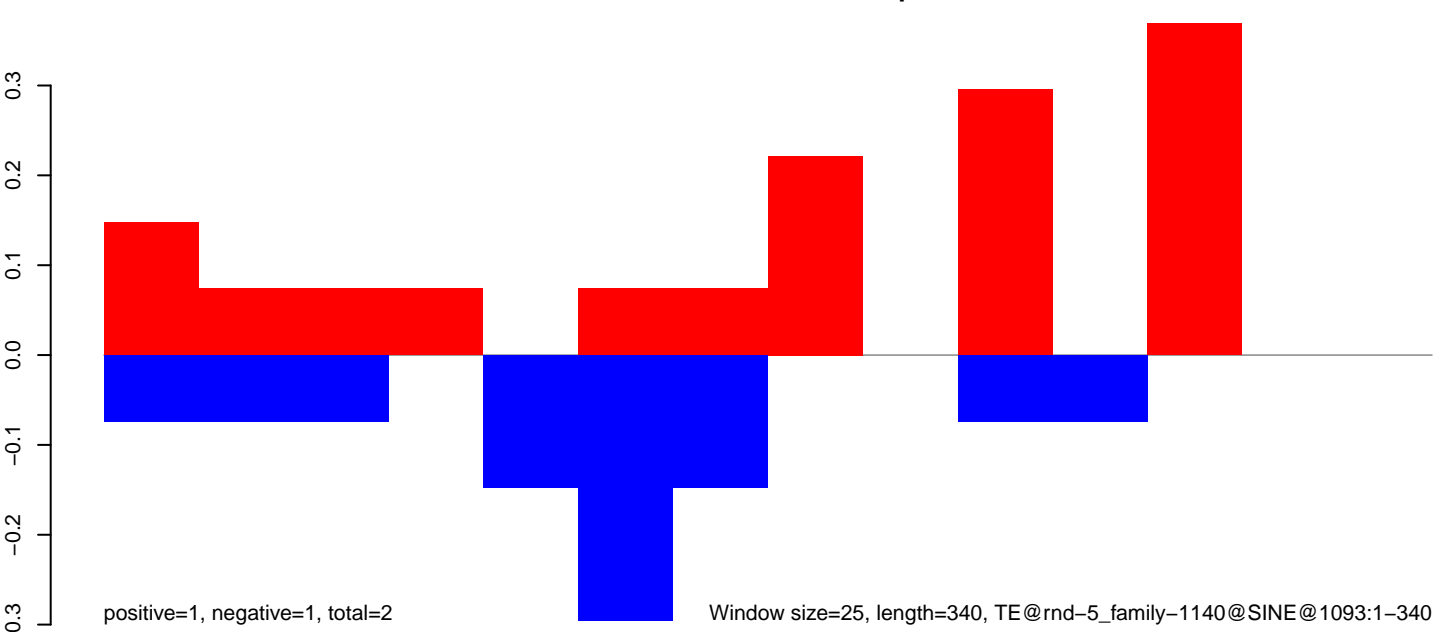
CuQuin_Testes_NL.18_23.rep



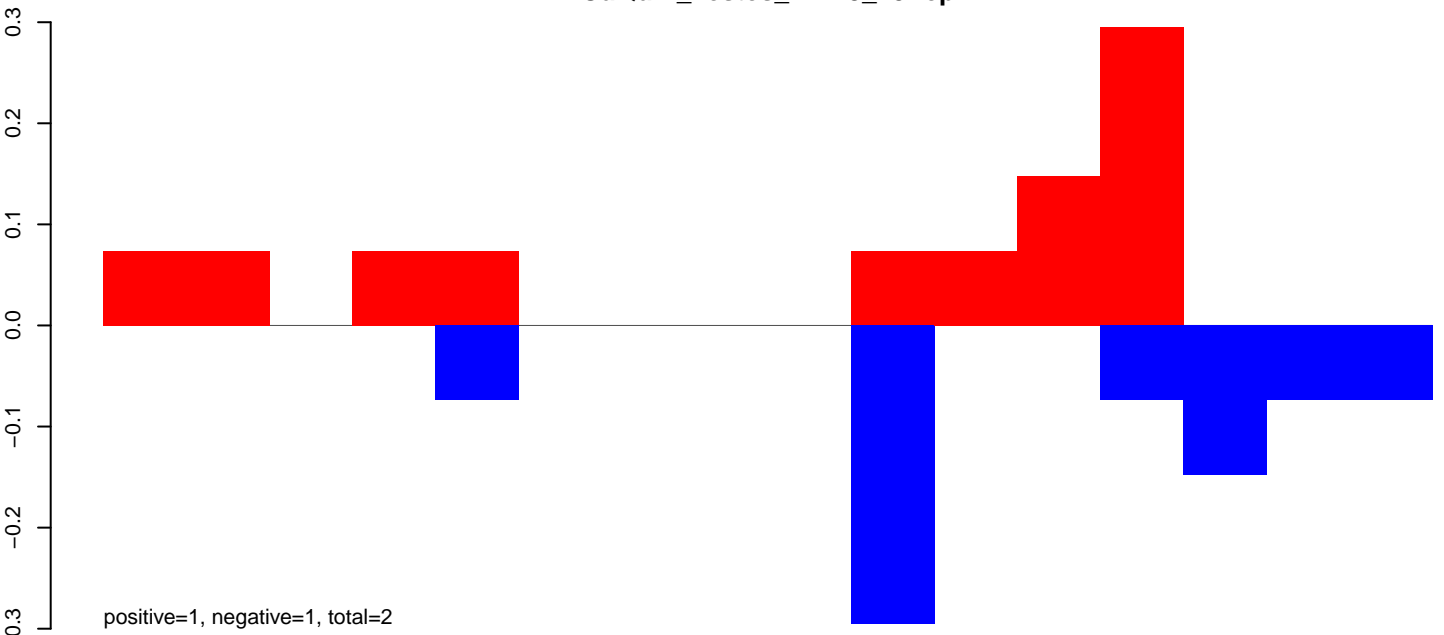
CuQuin_Testes_NL.24_35.rep



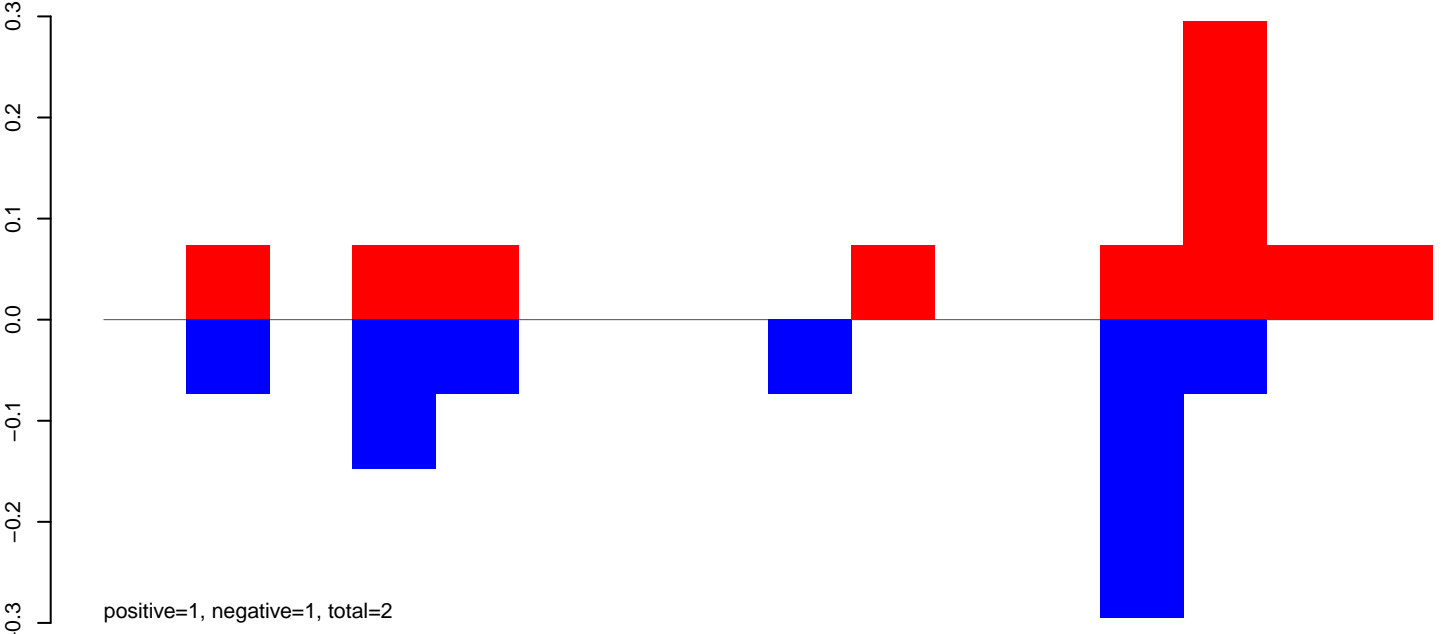
CuQuin_Testes_NL.rep



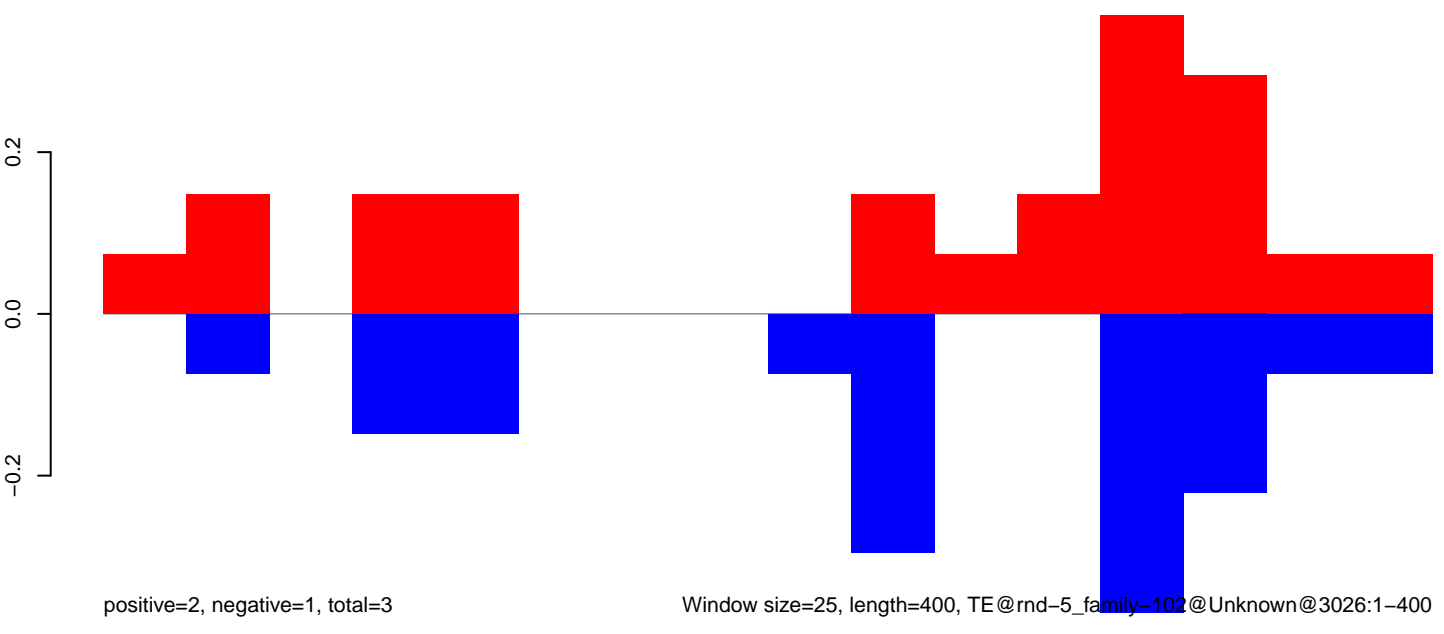
CuQuin_Testes_NL.18_23.rep



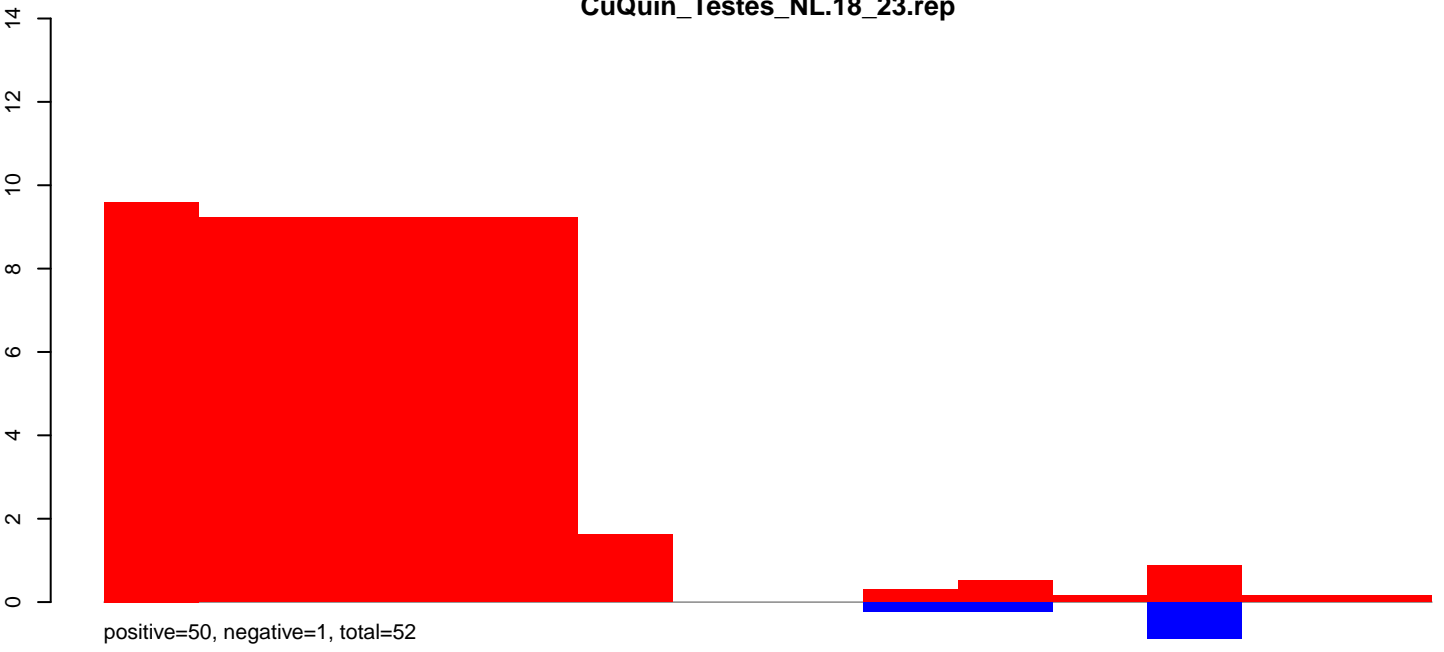
CuQuin_Testes_NL.24_35.rep



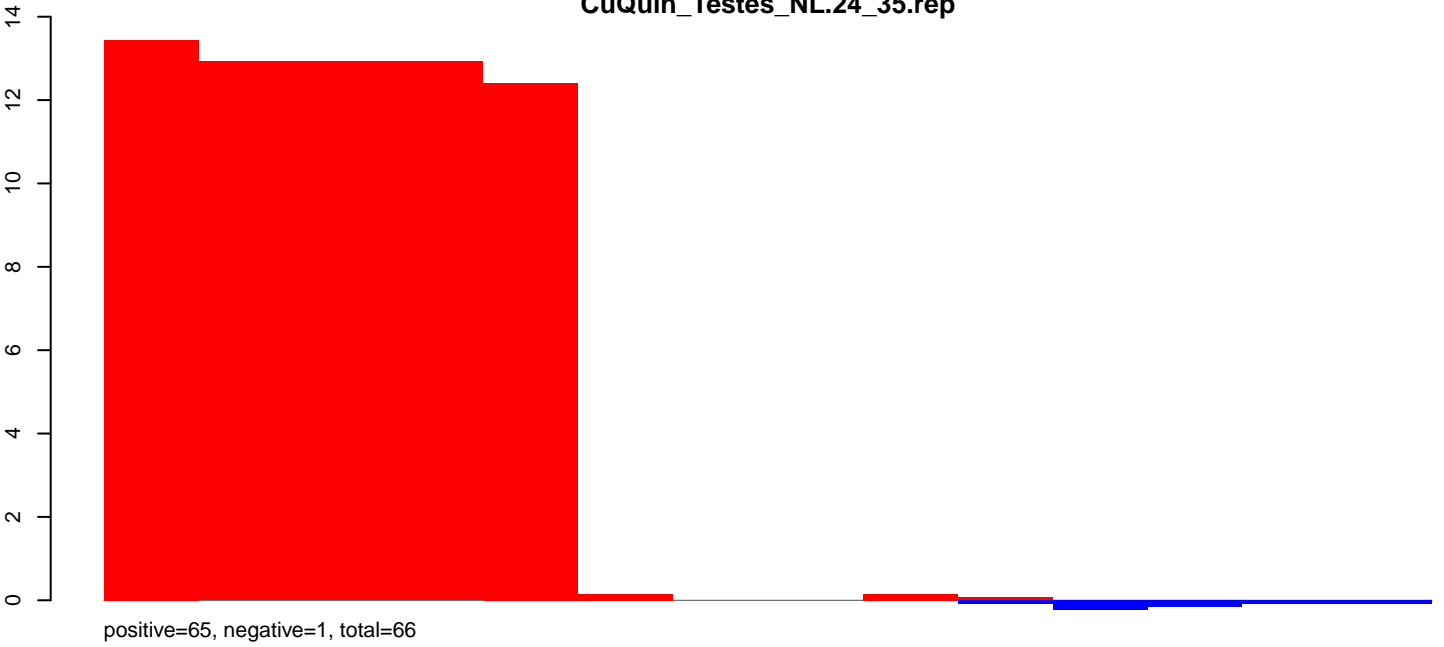
CuQuin_Testes_NL.rep



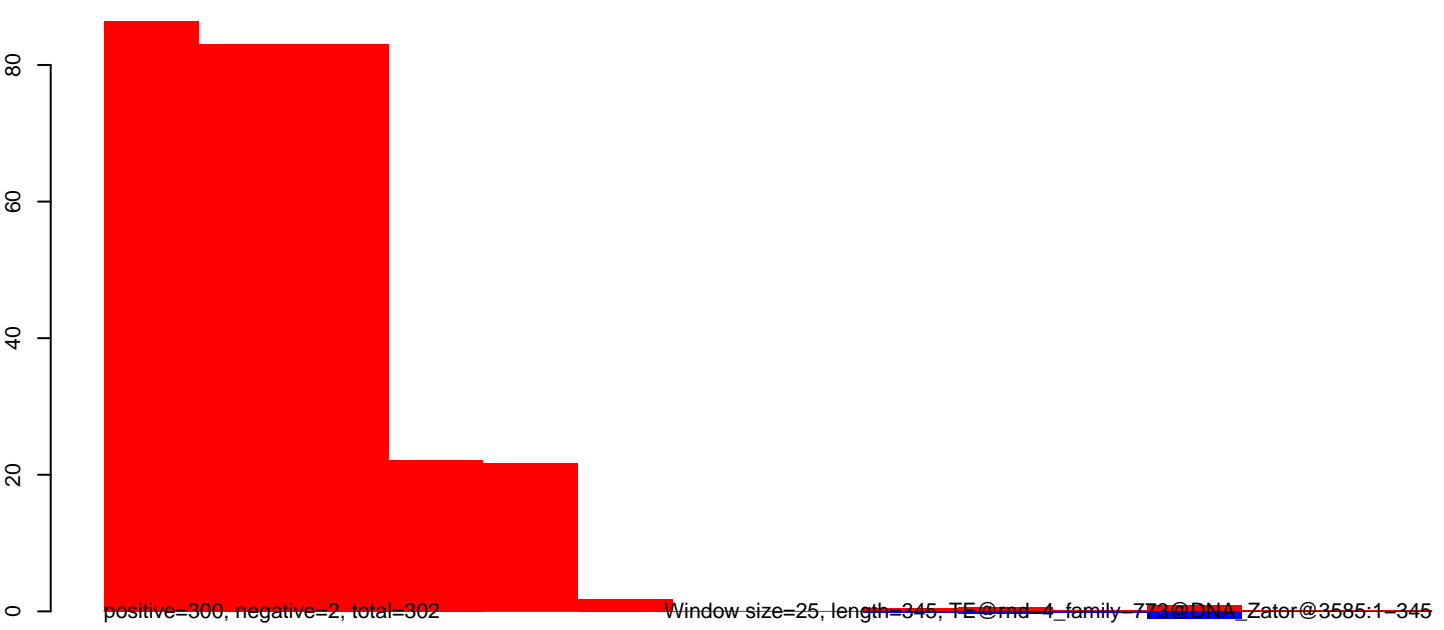
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

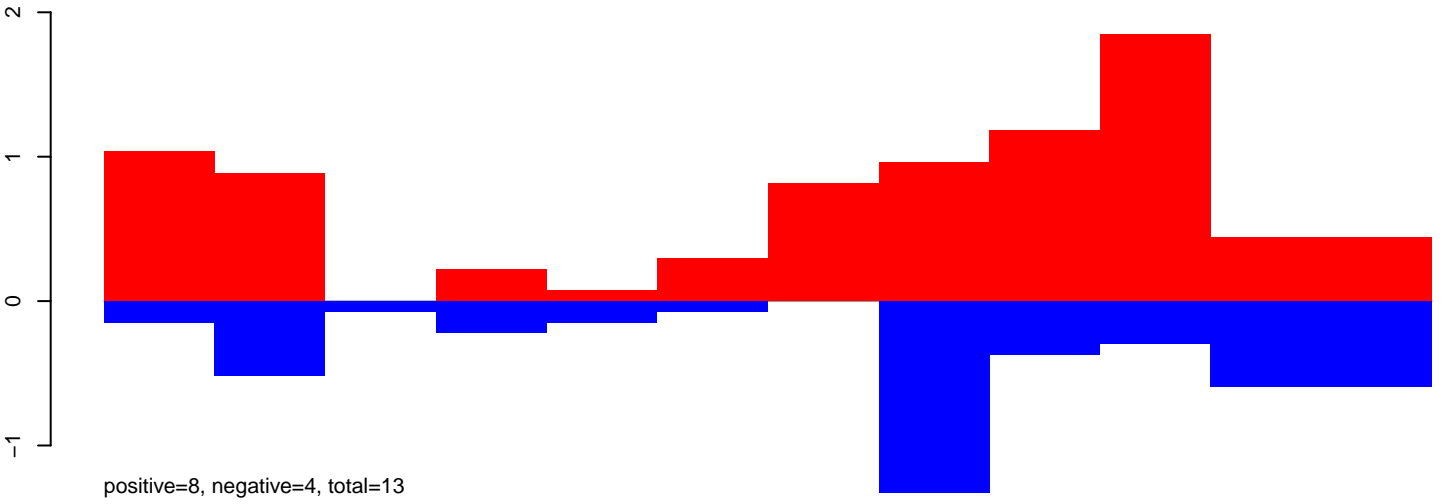


CuQuin_Testes_NL.rep

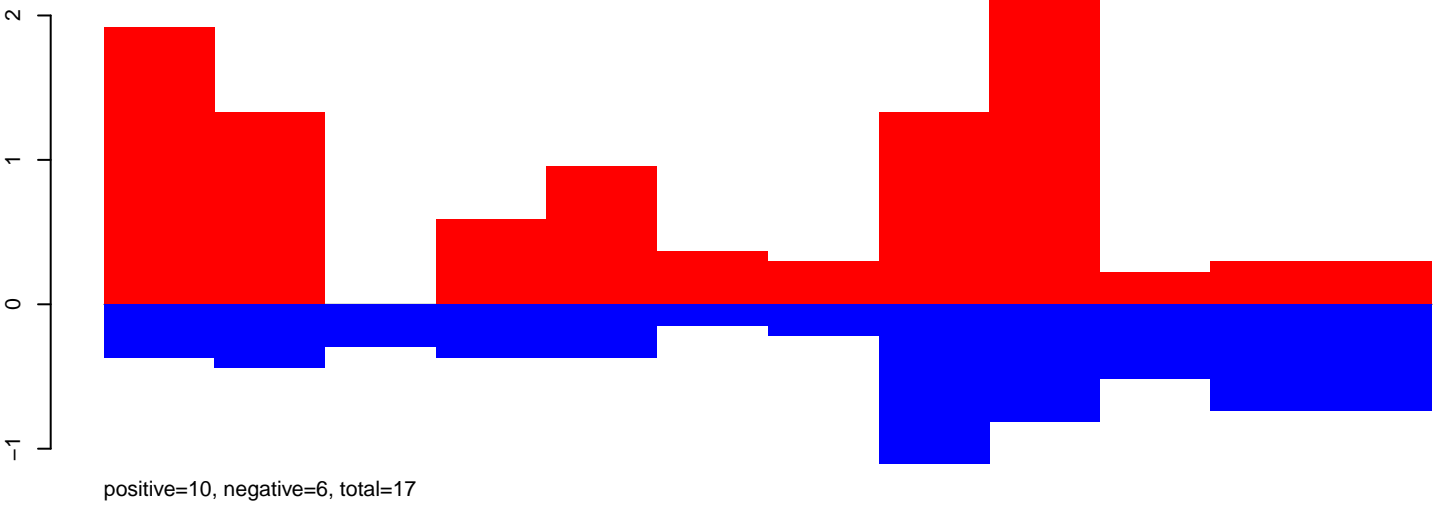


Window size=25, length=345, TE@md-4_family-773@DNA_Zator@3585:1-345

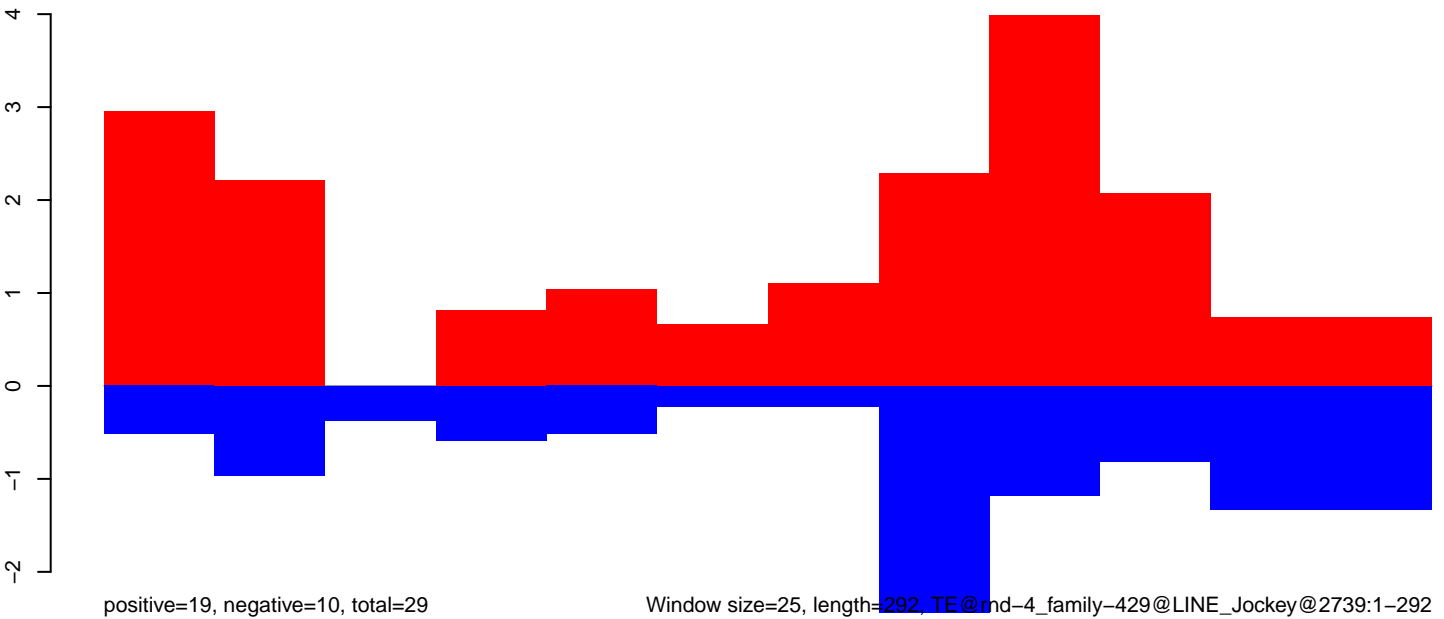
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



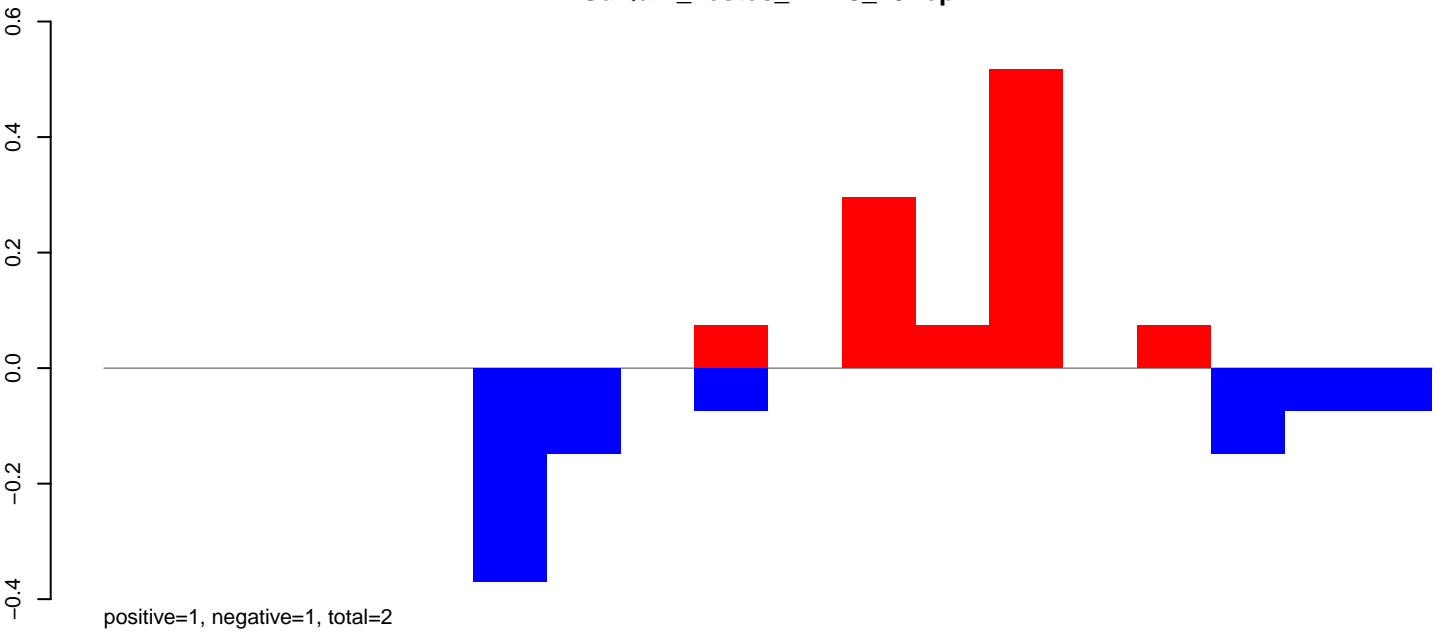
CuQuin_Testes_NL.rep



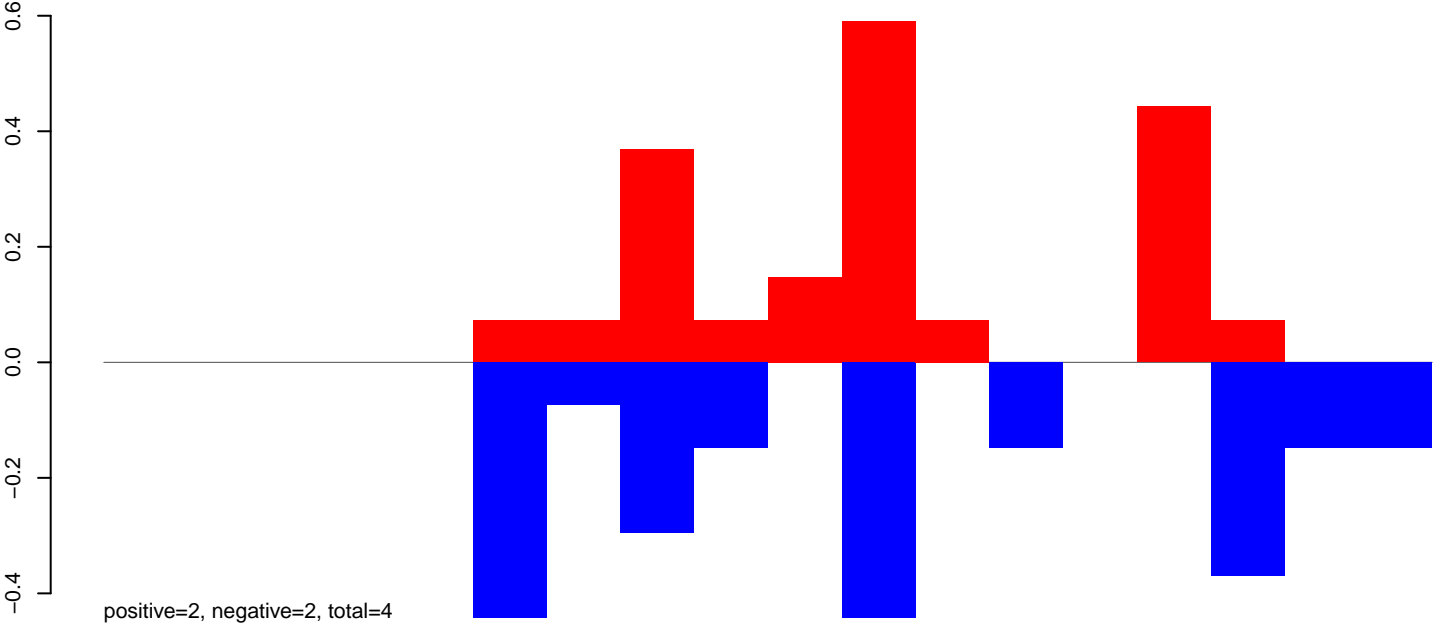
Window size=25, length=292, TE@rnd-4_family-429@LINE_Jockey@2739:1-292

50 100 150 200 250 300

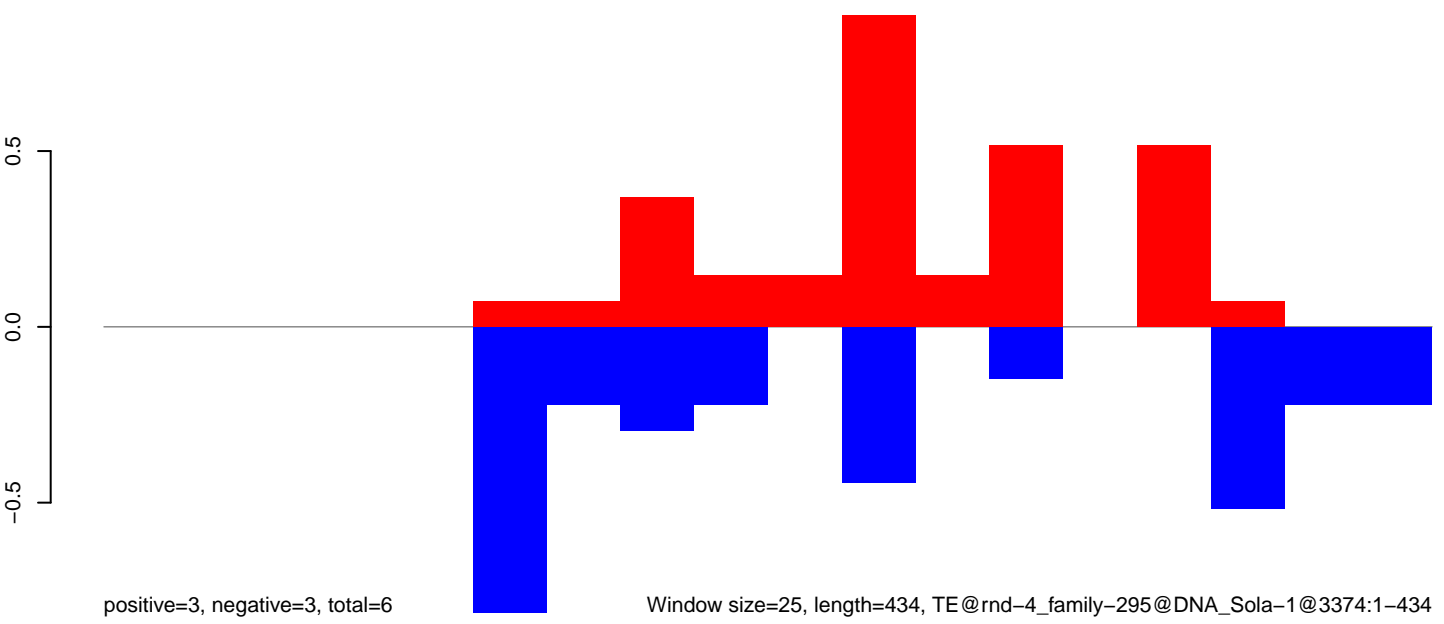
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

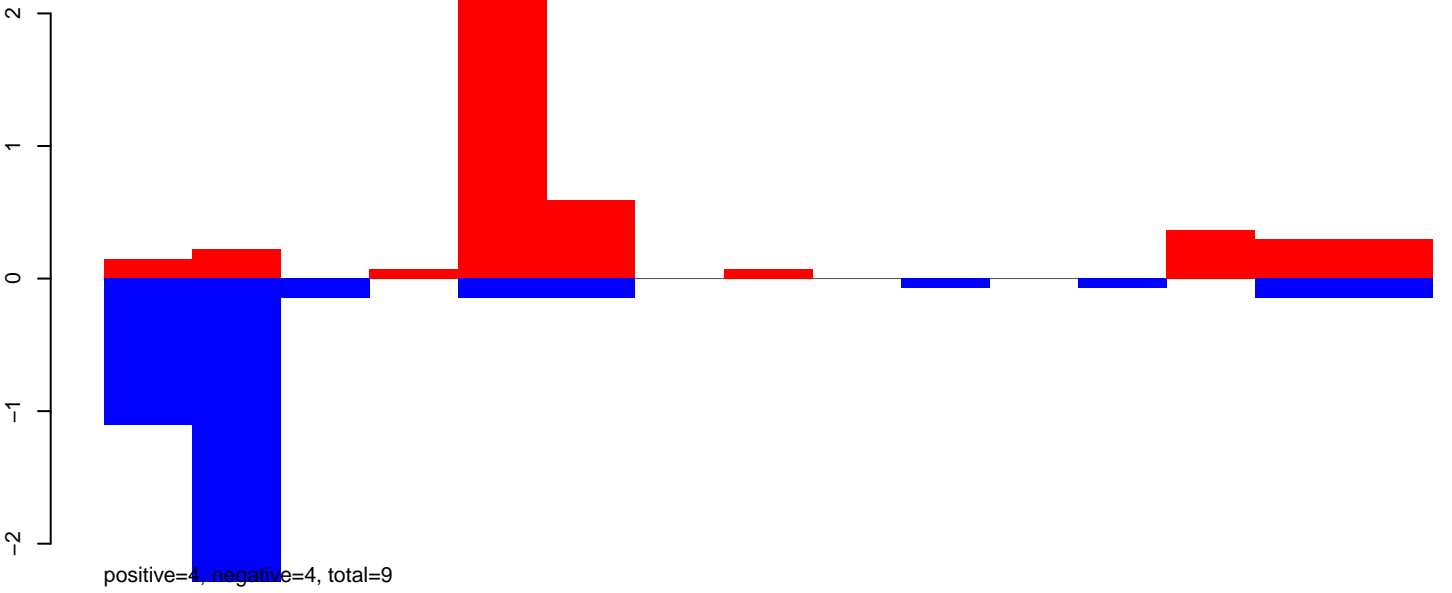


CuQuin_Testes_NL.rep

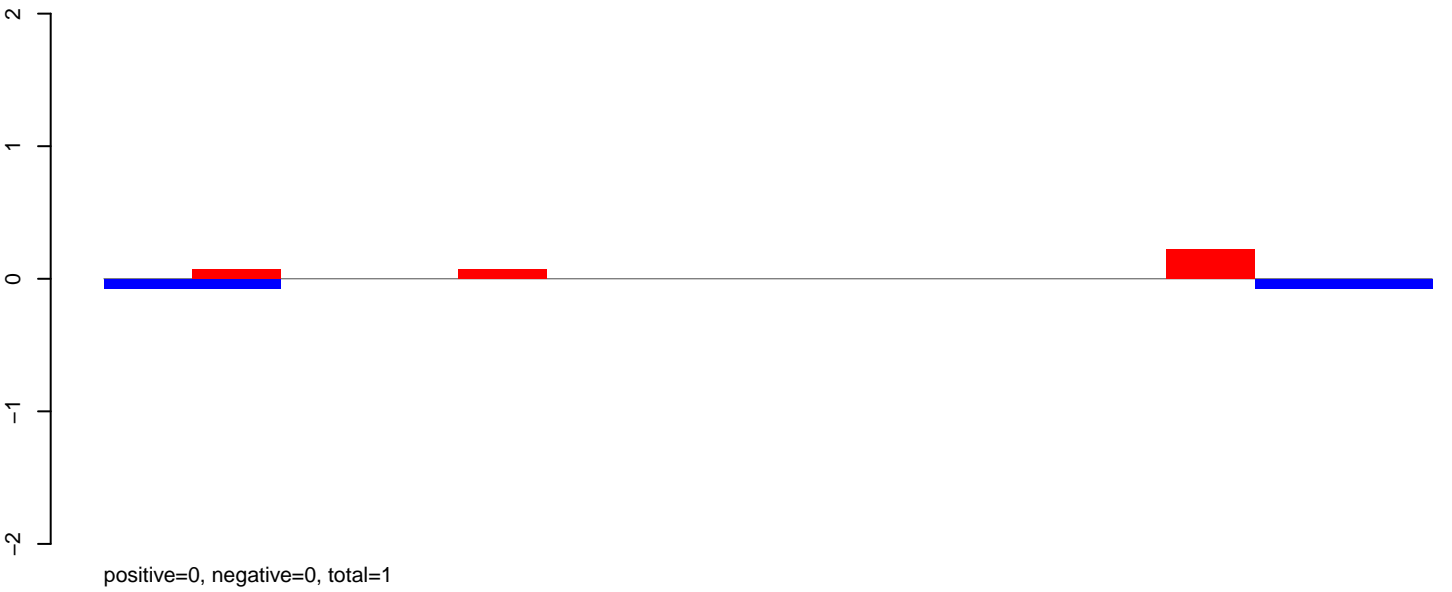


Window size=25, length=434, TE@rnd-4_family-295@DNA_Sola-1@3374:1-434

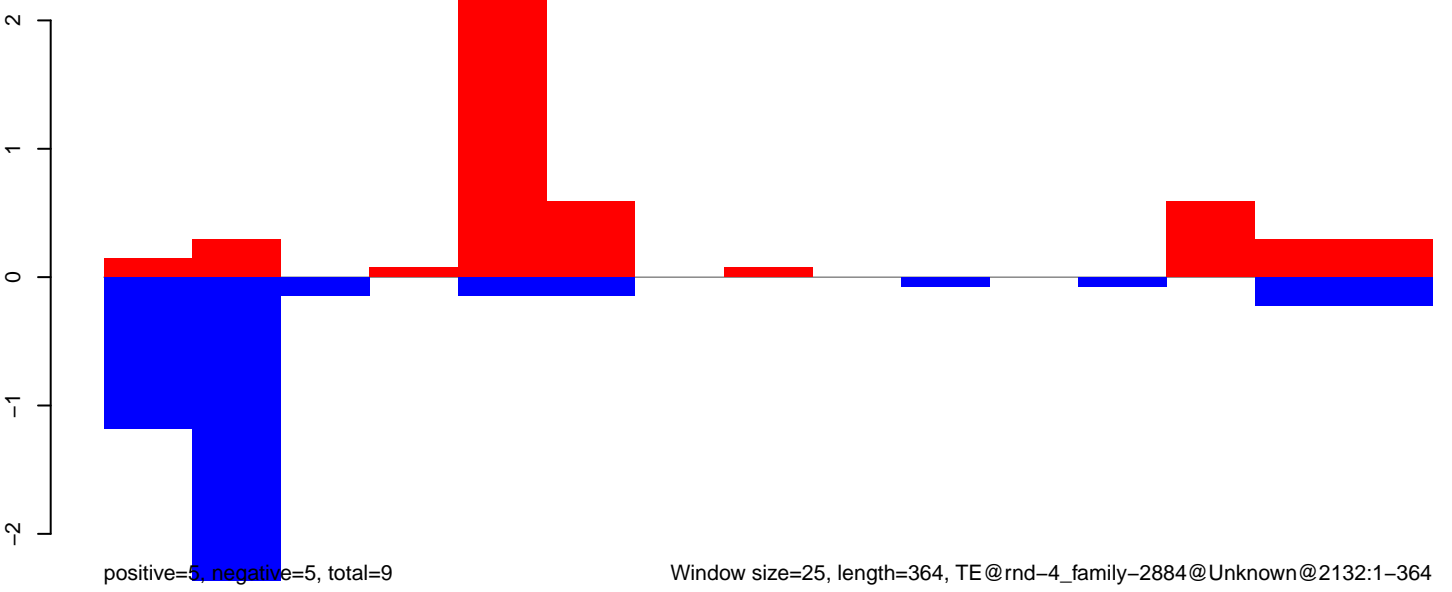
CuQuin_Testes_NL.18_23.rep



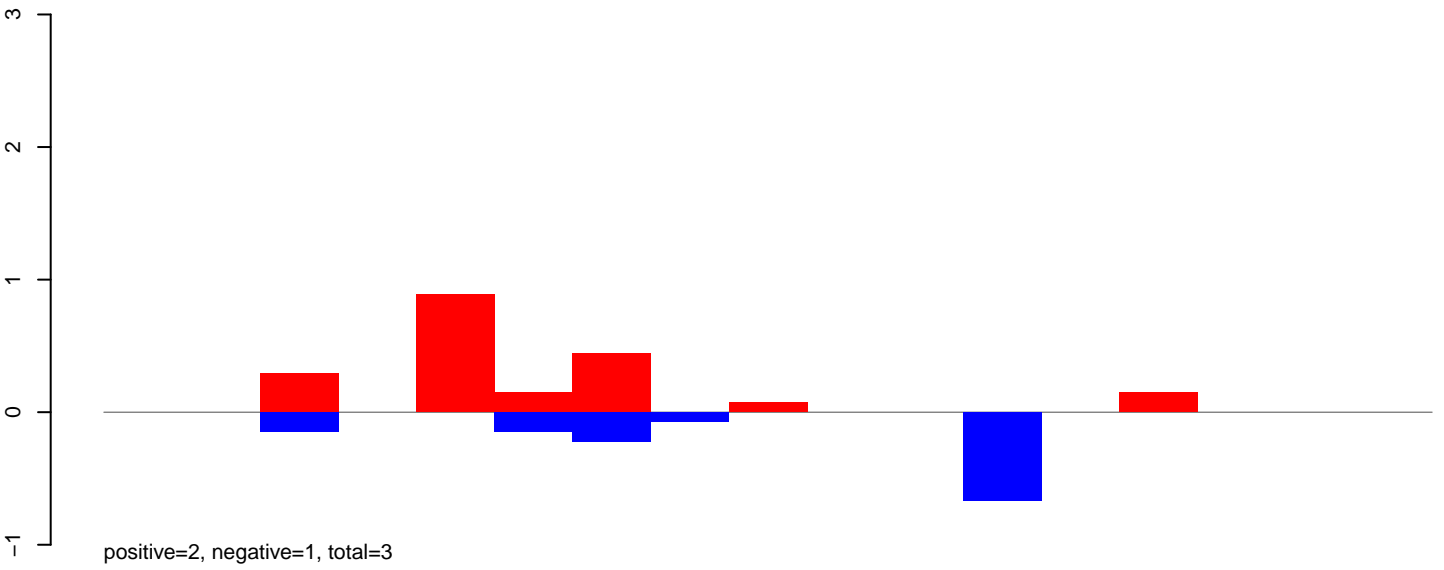
CuQuin_Testes_NL.24_35.rep



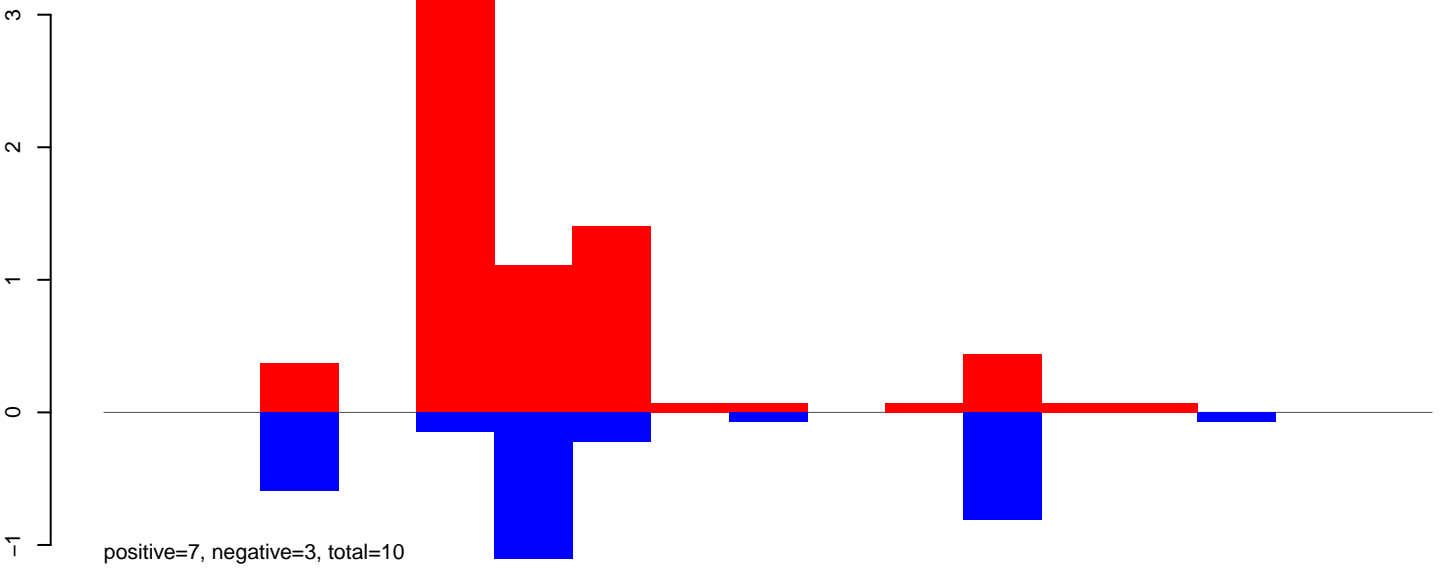
CuQuin_Testes_NL.rep



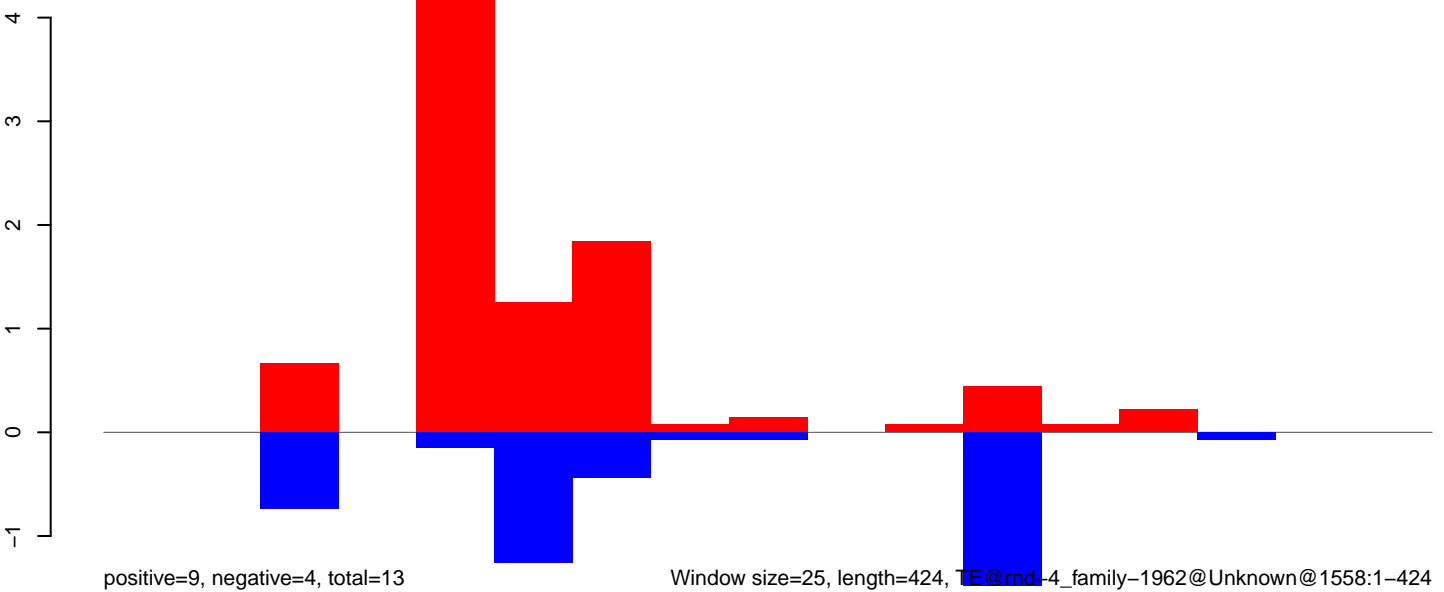
CuQuin_Testes_NL.18_23.rep



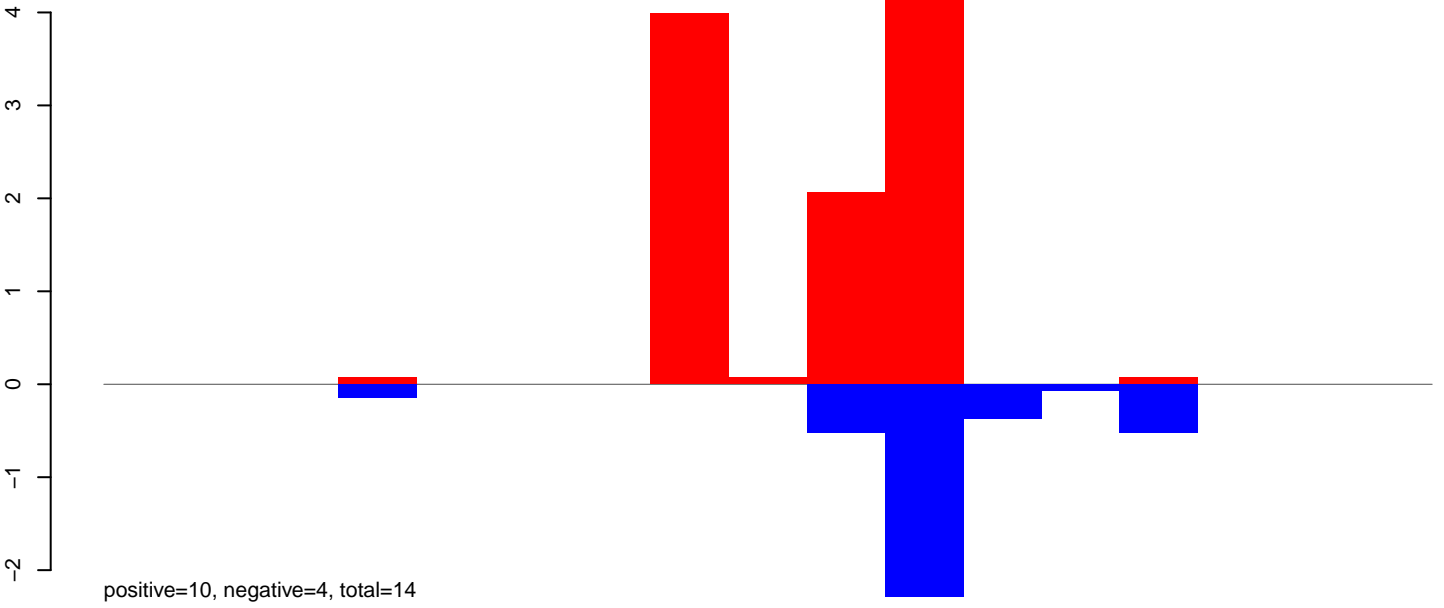
CuQuin_Testes_NL.24_35.rep



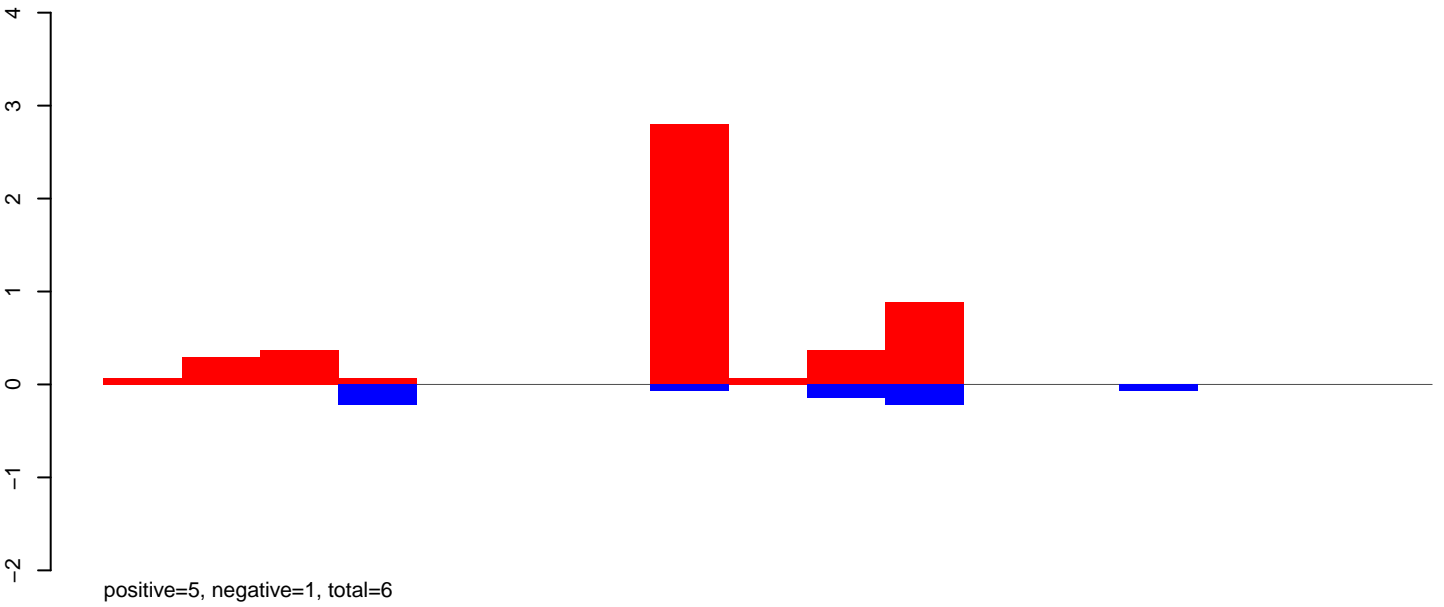
CuQuin_Testes_NL.rep



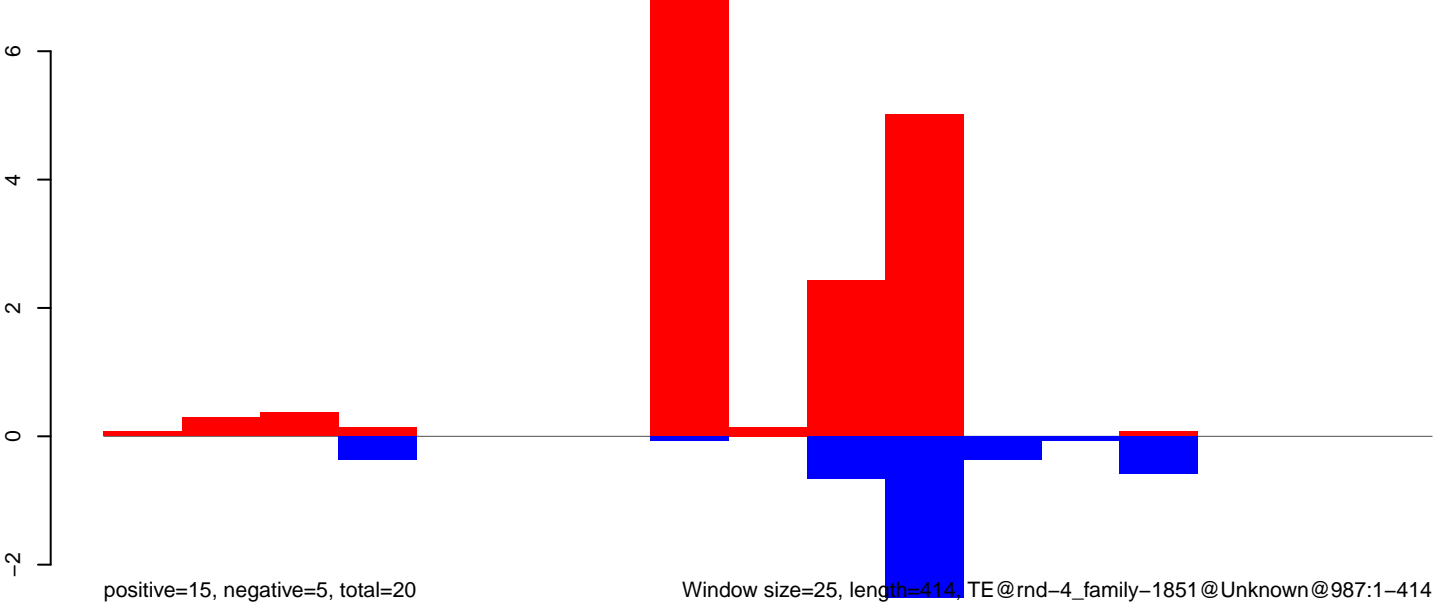
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

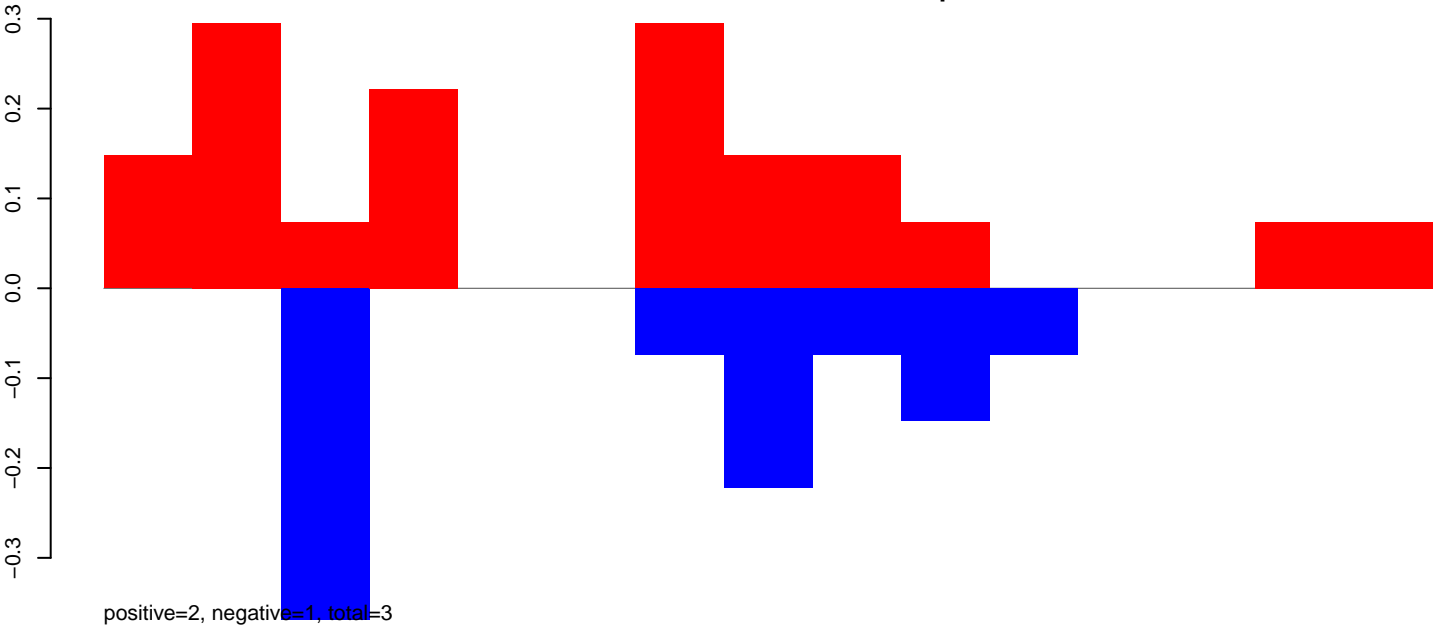


CuQuin_Testes_NL.rep

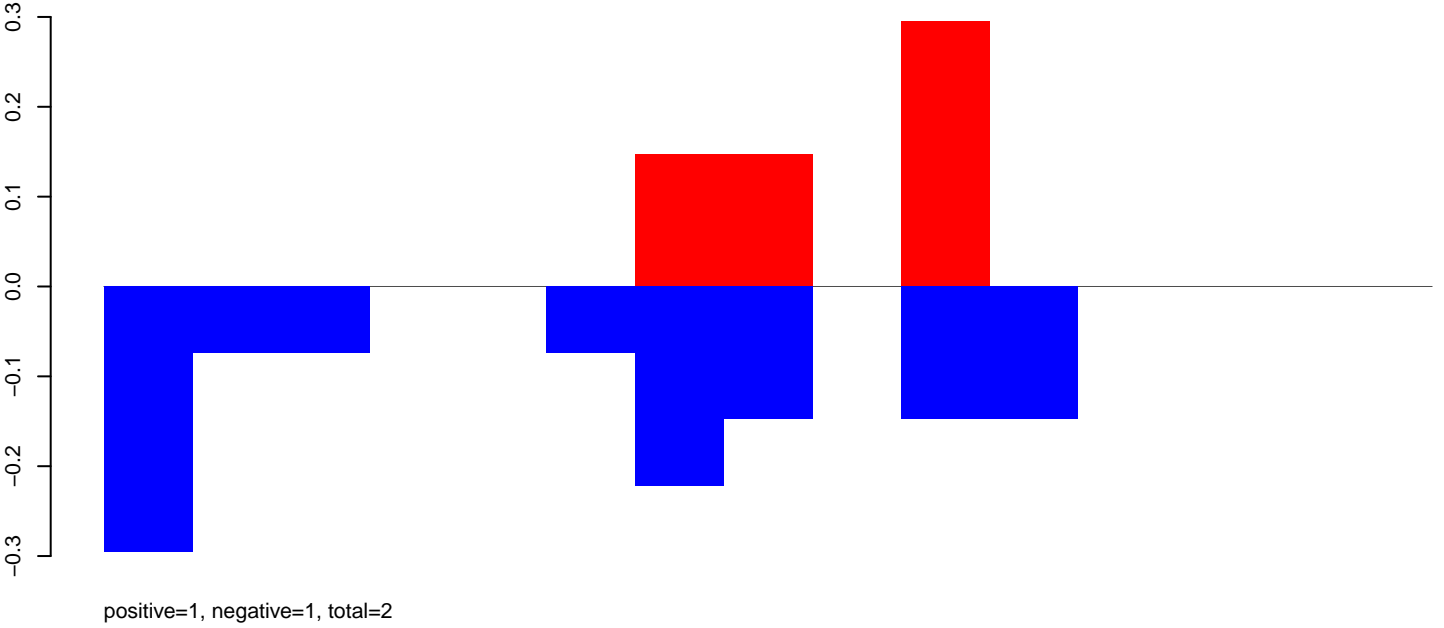


Window size=25, length=414, TE@rnd-4_family-1851@Unknown@987:1-414

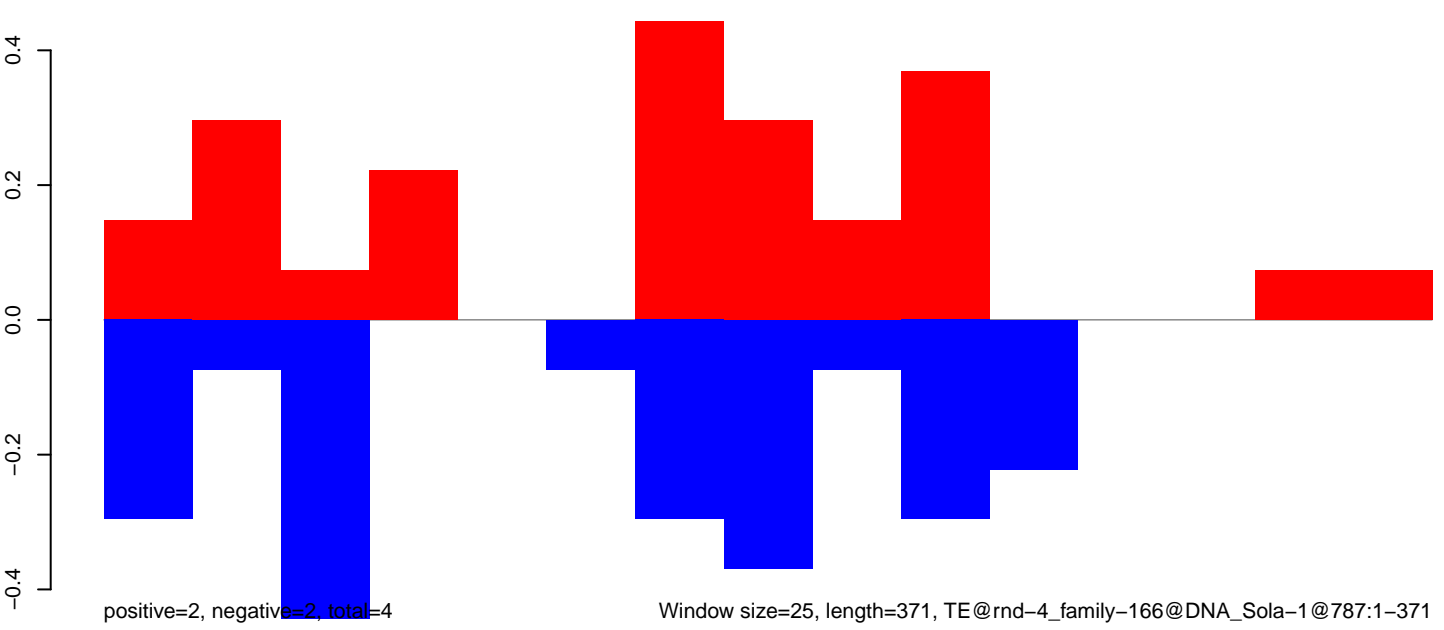
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



Window size=25, length=371, TE@rnd-4_family-166@DNA_Sola-1@787:1-371

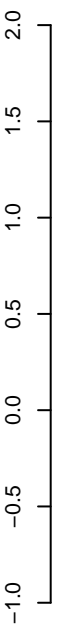
100

200

300

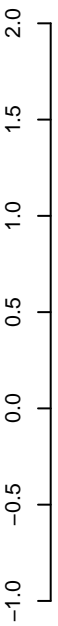
400

CuQuin_Testes_NL.18_23.rep



positive=3, negative=5, total=8

CuQuin_Testes_NL.24_35.rep



positive=10, negative=4, total=14

CuQuin_Testes_NL.rep

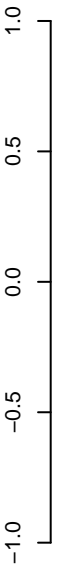


positive=13, negative=9, total=22

Window size=25, length=294, TE@rnd-3_family-50@Unknown@2606:1-294

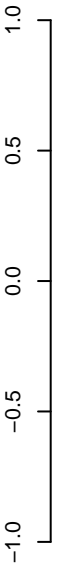


CuQuin_Testes_NL.18_23.rep



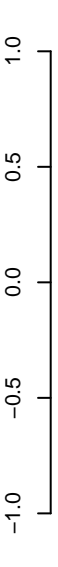
positive=1, negative=0, total=1

CuQuin_Testes_NL.24_35.rep



positive=4, negative=2, total=6

CuQuin_Testes_NL.rep

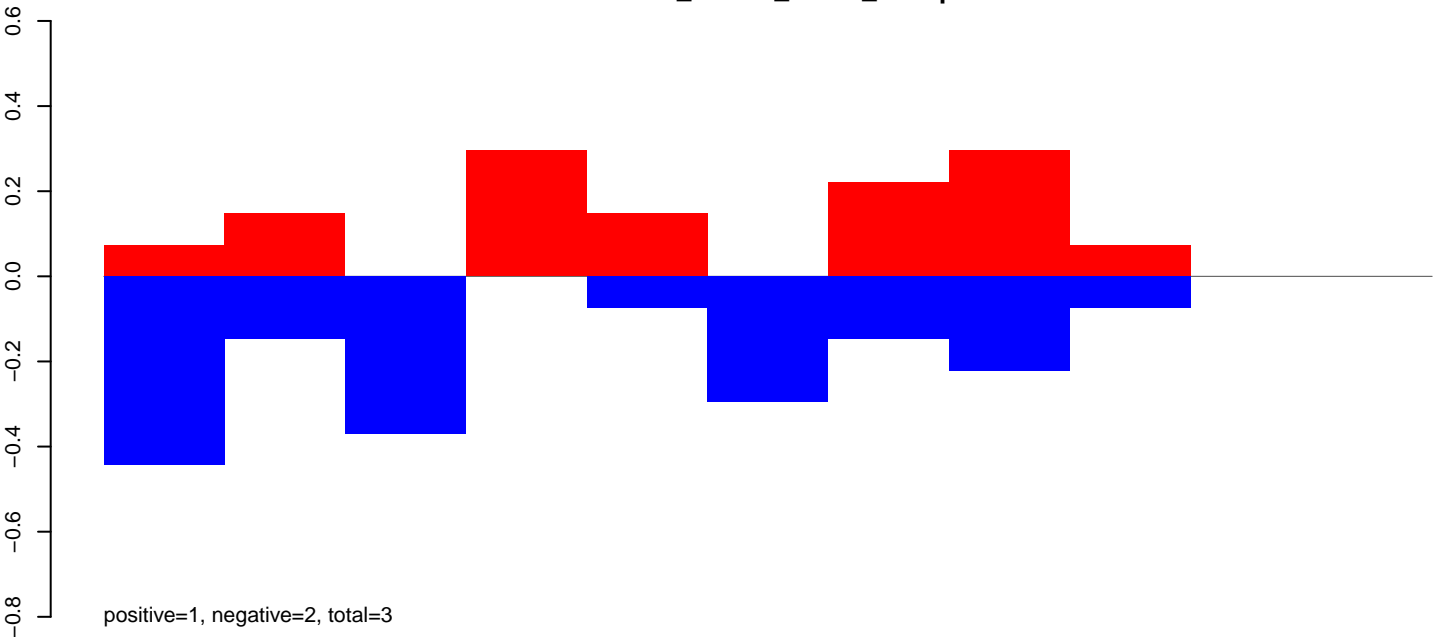


positive=5, negative=2, total=7

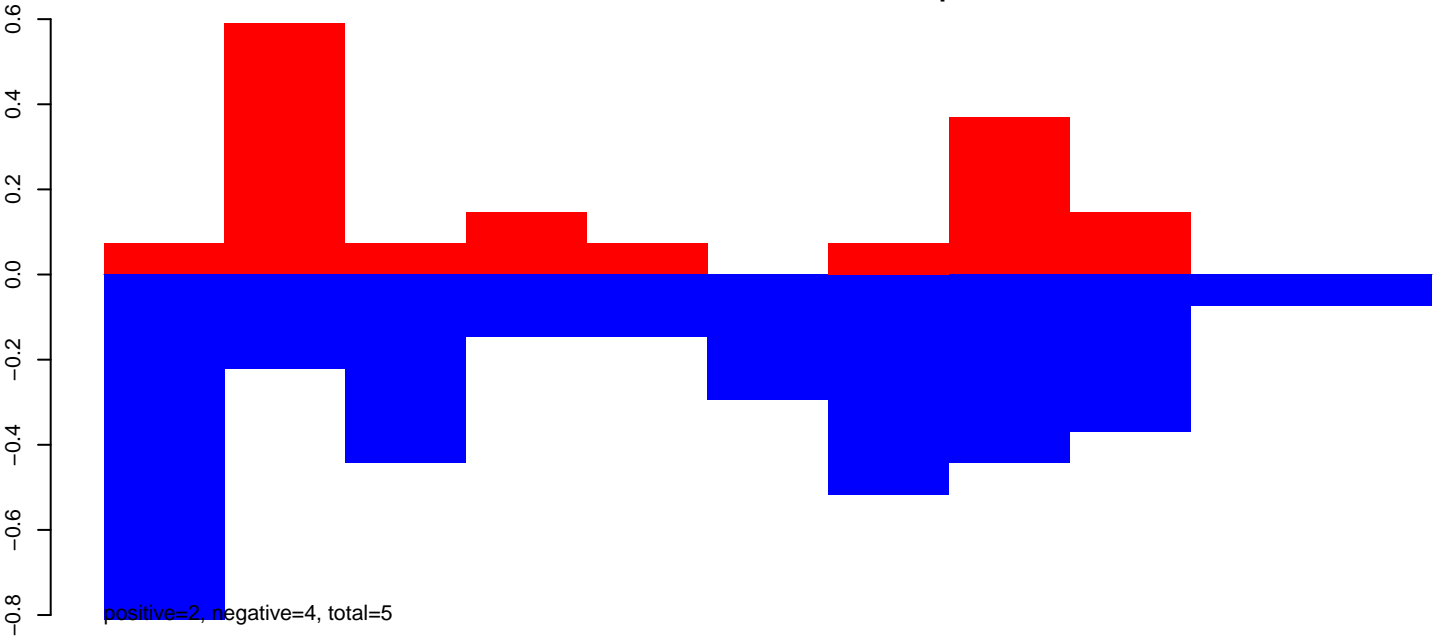
Window size=25, length=615, TE@rnd-3_family-114@DNA_Zator@2638:1-615

100 200 300 400 500 600

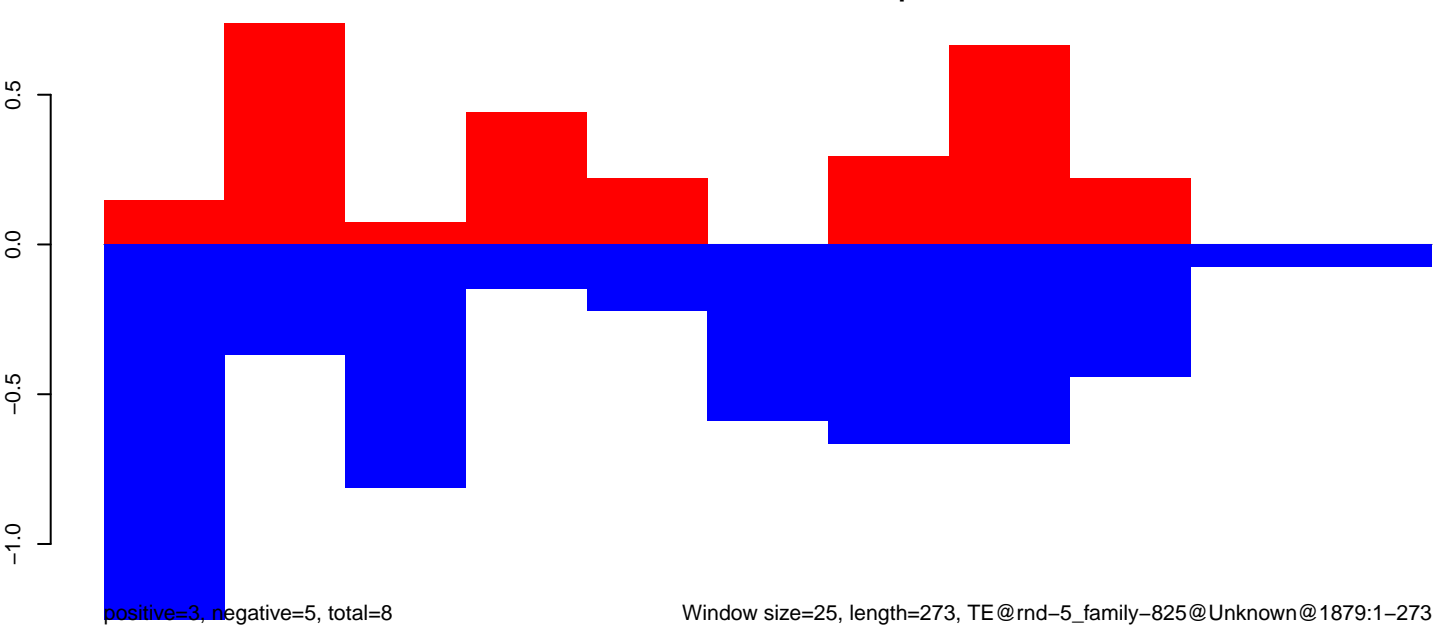
CuQuin_Testes_NL.18_23.rep



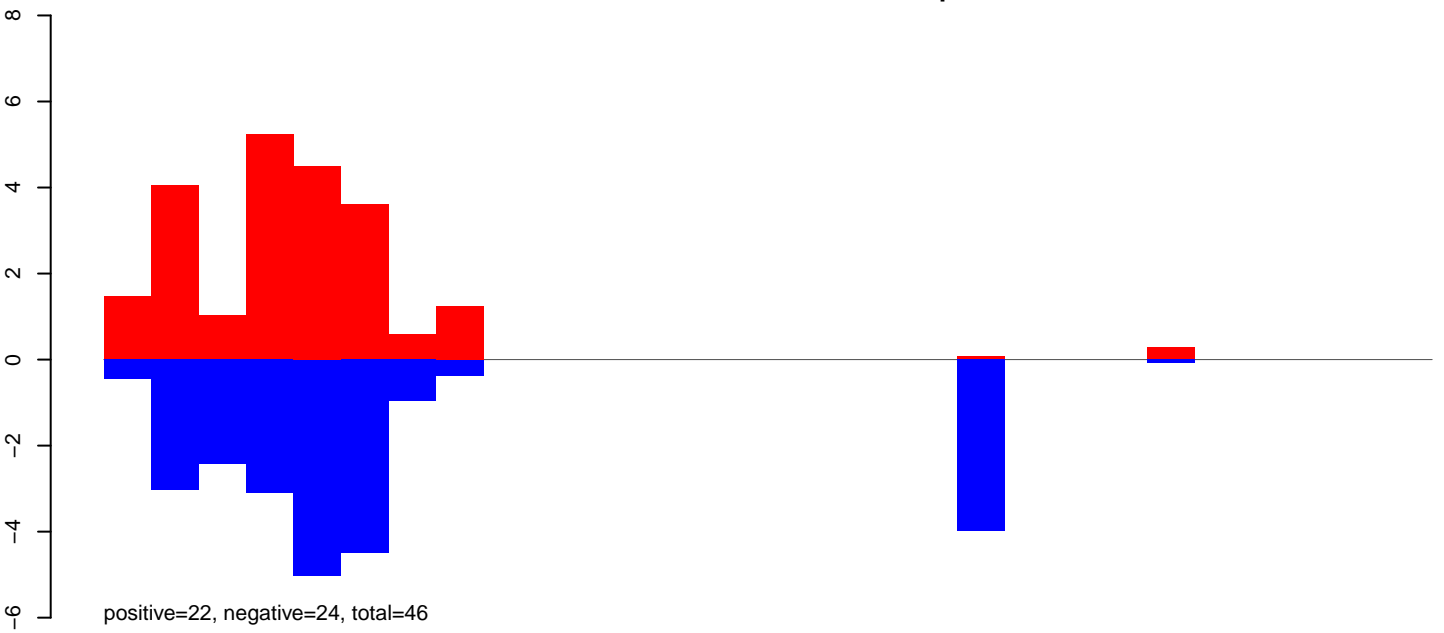
CuQuin_Testes_NL.24_35.rep



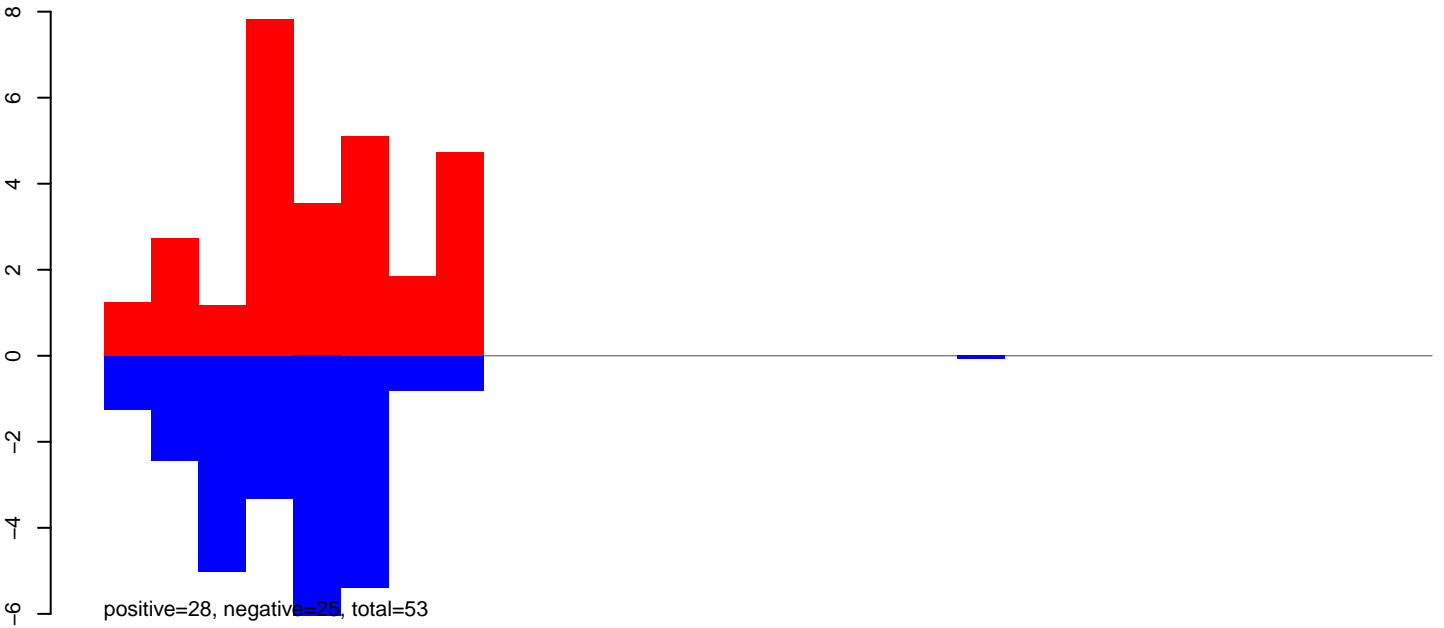
CuQuin_Testes_NL.rep



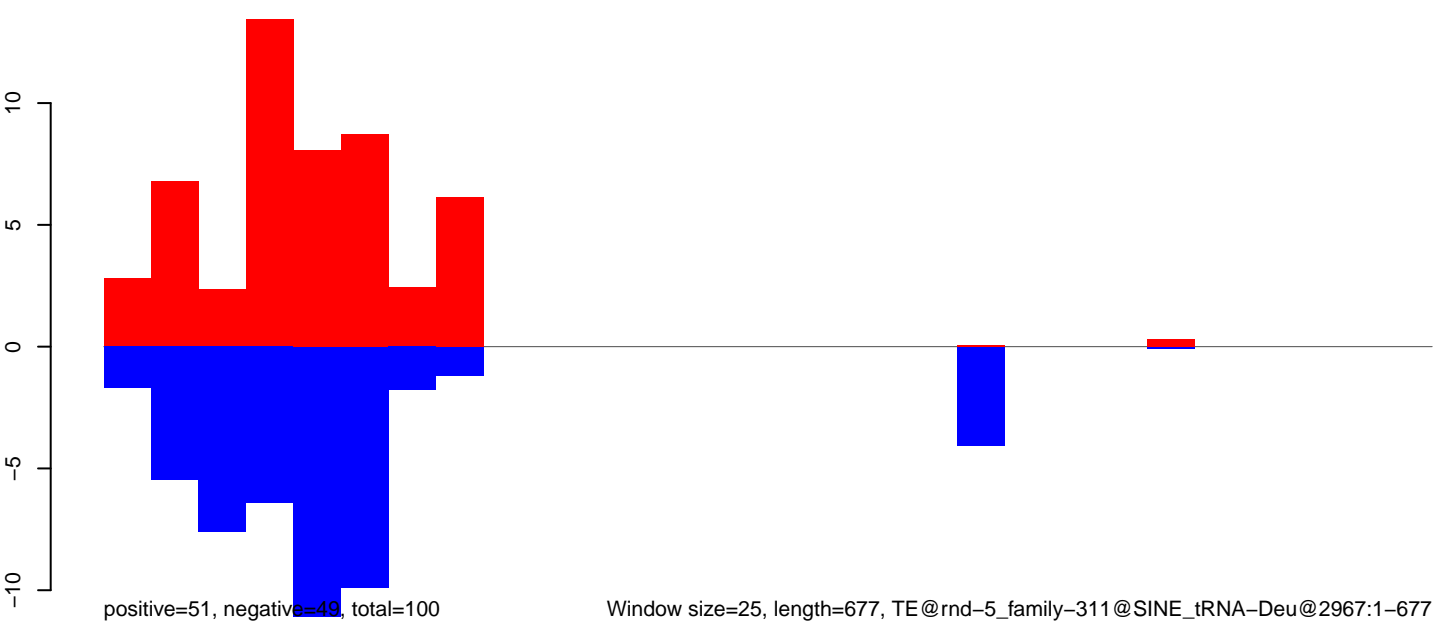
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



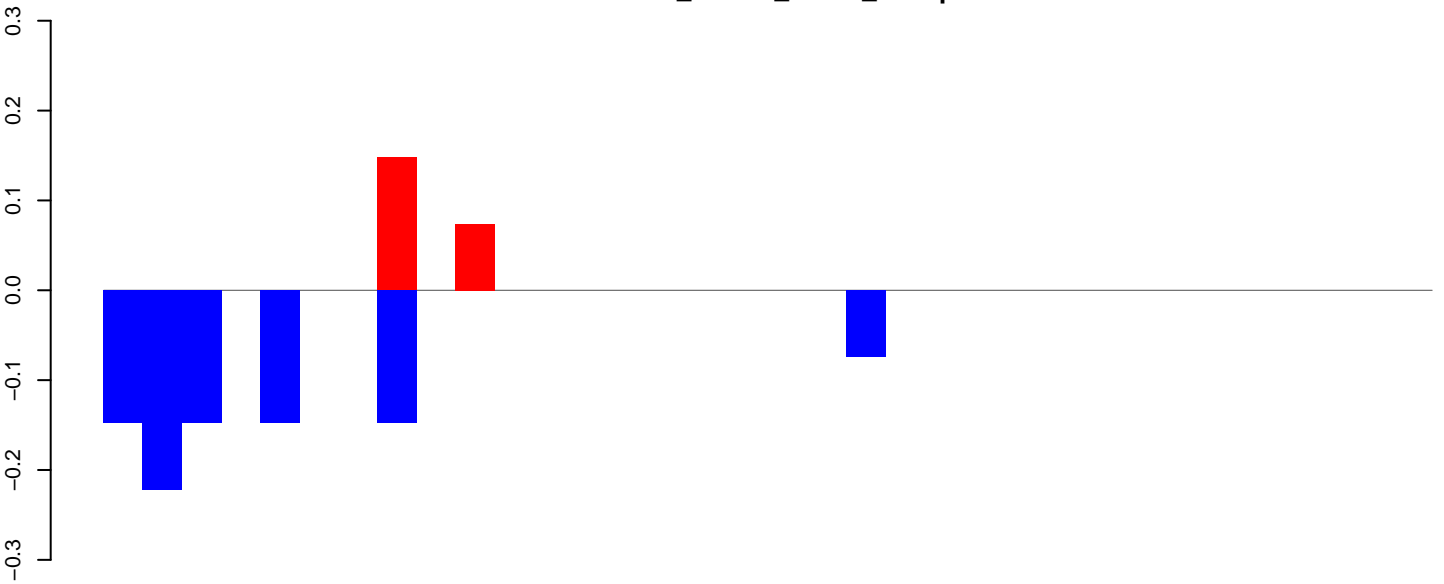
CuQuin_Testes_NL.rep



Window size=25, length=677, TE@rnd-5_family-311@SINE_tRNA-Deu@2967:1-677

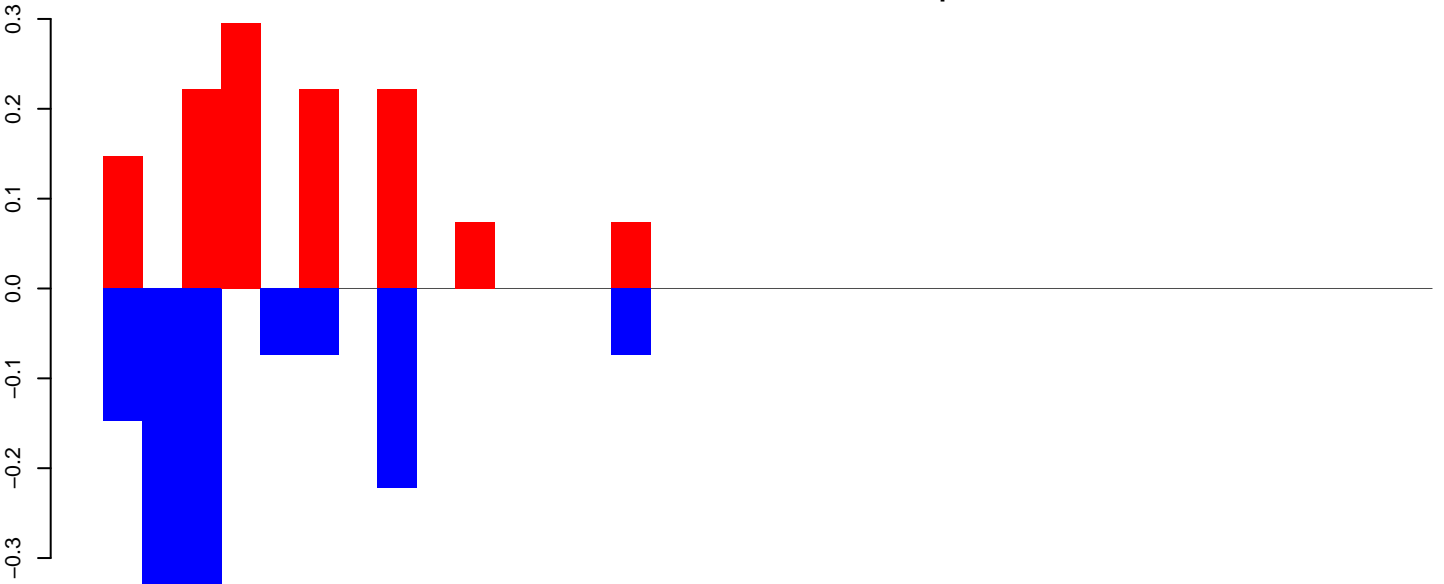
0 100 200 300 400 500 600 700

CuQuin_Testes_NL.18_23.rep



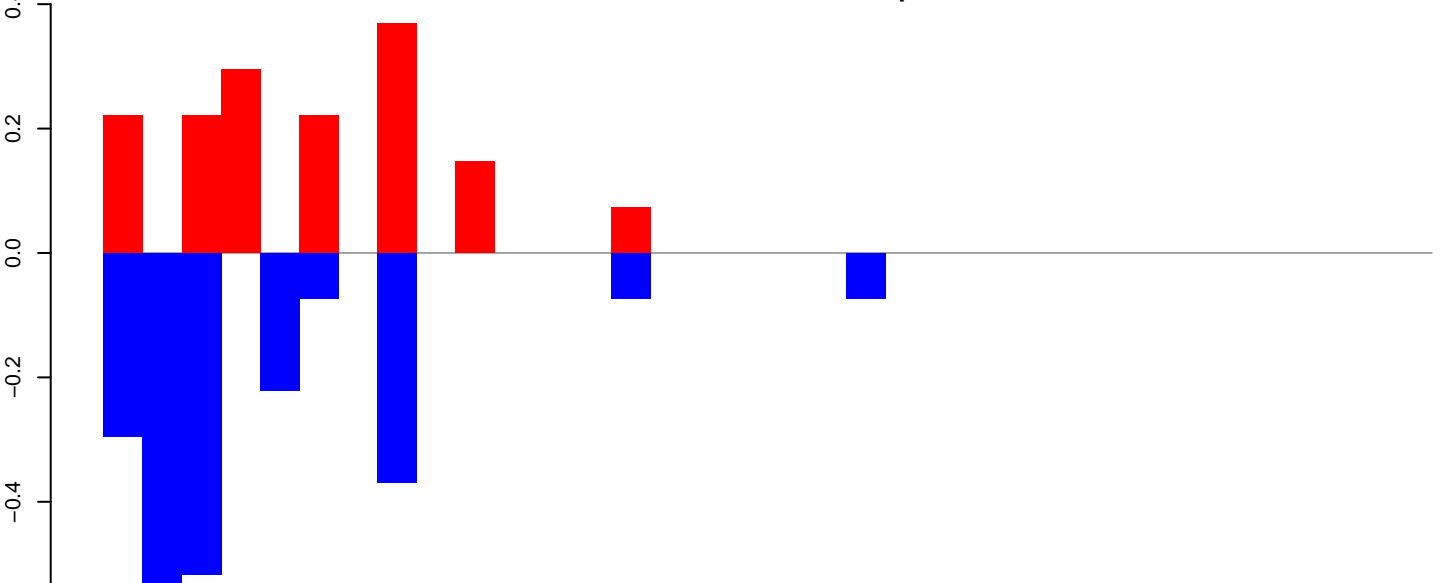
positive=0, negative=1, total=1

CuQuin_Testes_NL.24_35.rep



positive=1, negative=1, total=3

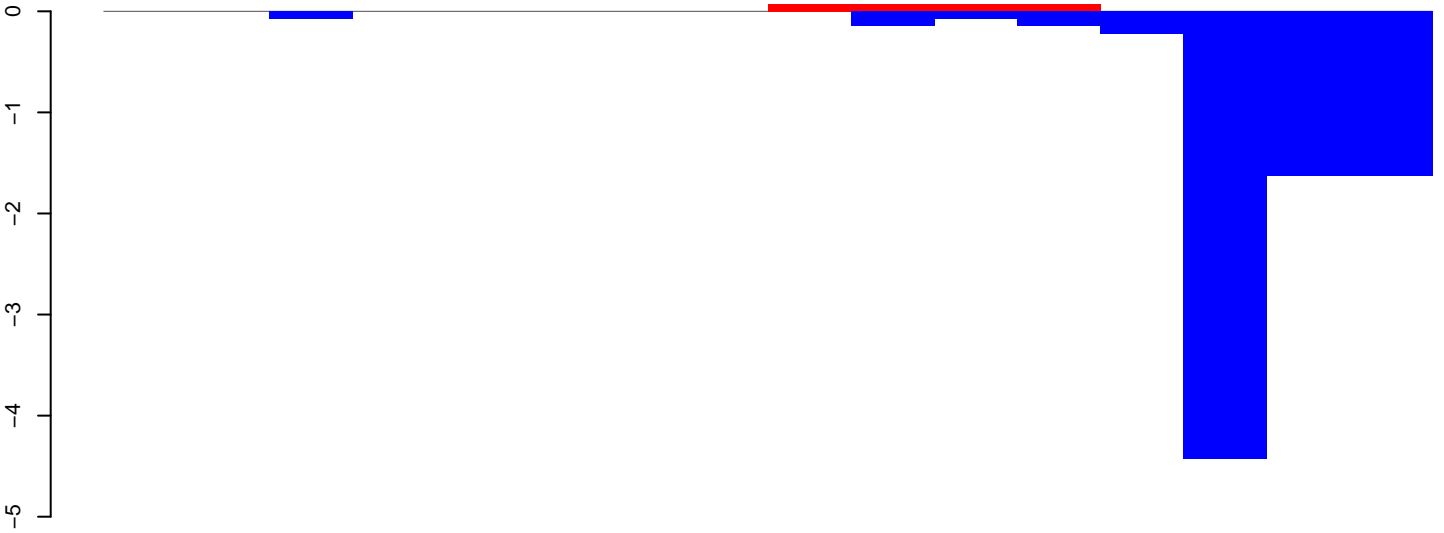
CuQuin_Testes_NL.rep



positive=2, negative=2, total=4

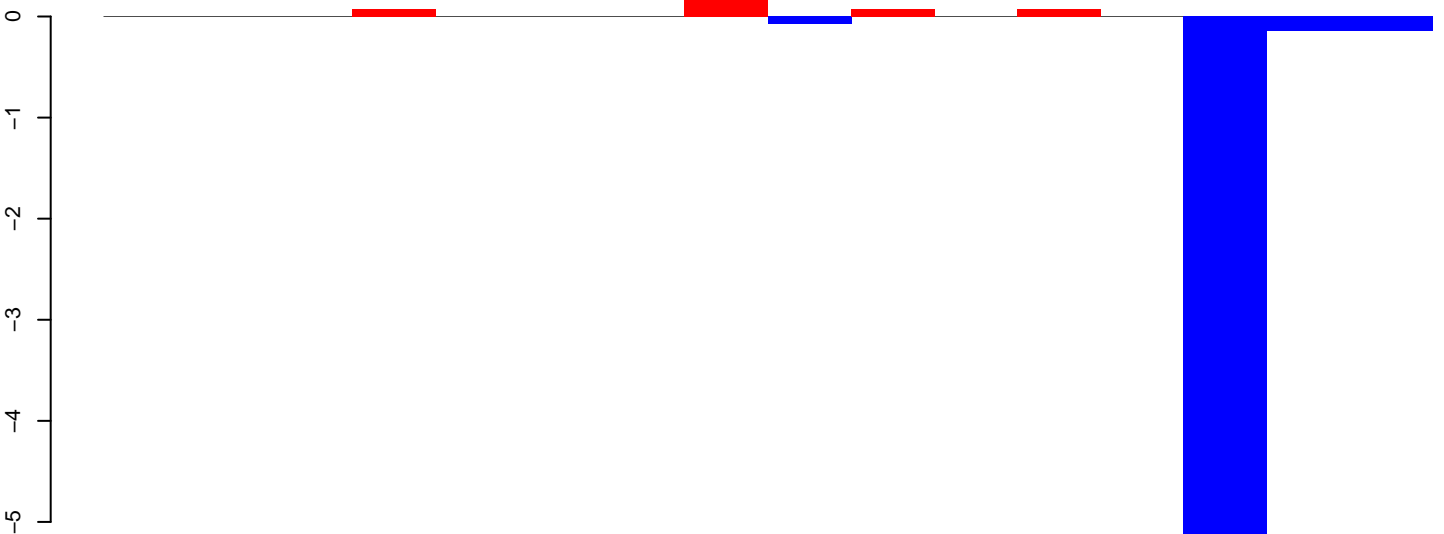
Window size=25, length=842, TE@rnd-4_family-873@DNA_PIF-Harbinger@3485:1-842

CuQuin_Testes_NL.18_23.rep



positive=0, negative=8, total=9

CuQuin_Testes_NL.24_35.rep



positive=0, negative=6, total=6

CuQuin_Testes_NL.rep

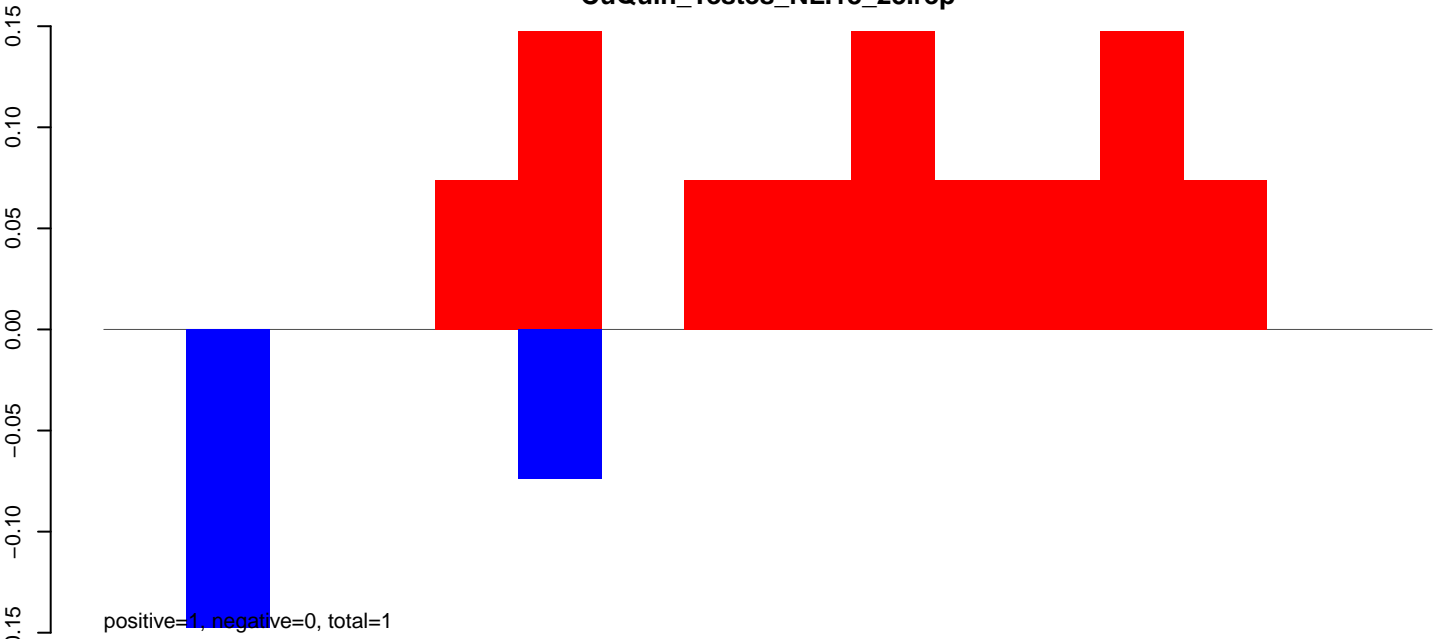


positive=1, negative=14, total=15

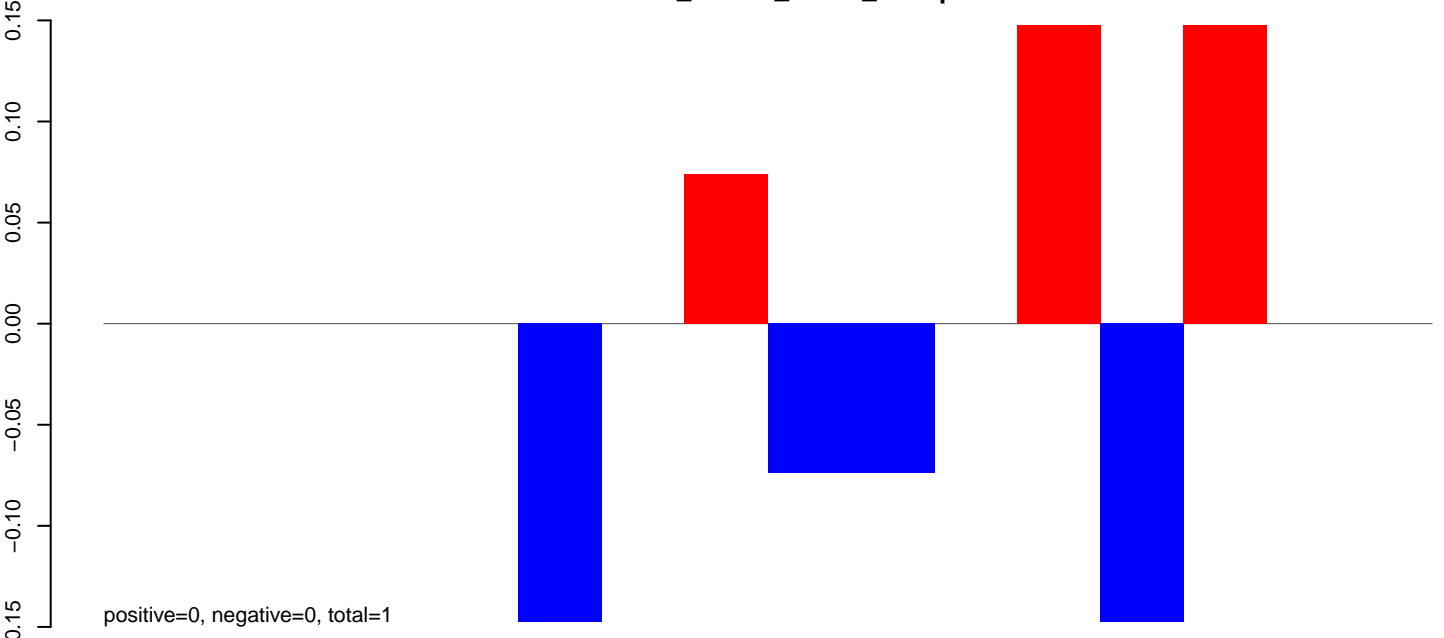
Window size=25, length=376, TE@rnd-4_family-830@DNA_TcMar_Tc4@3538:1-376

100 200 300 400

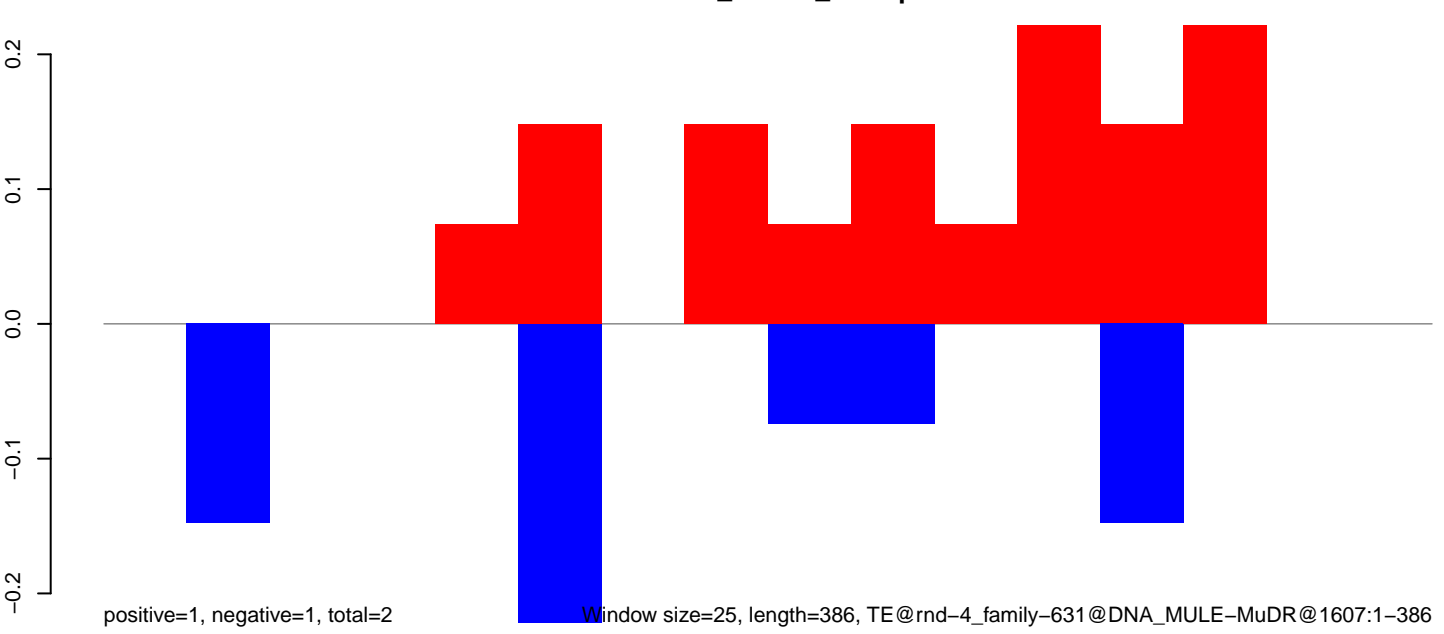
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



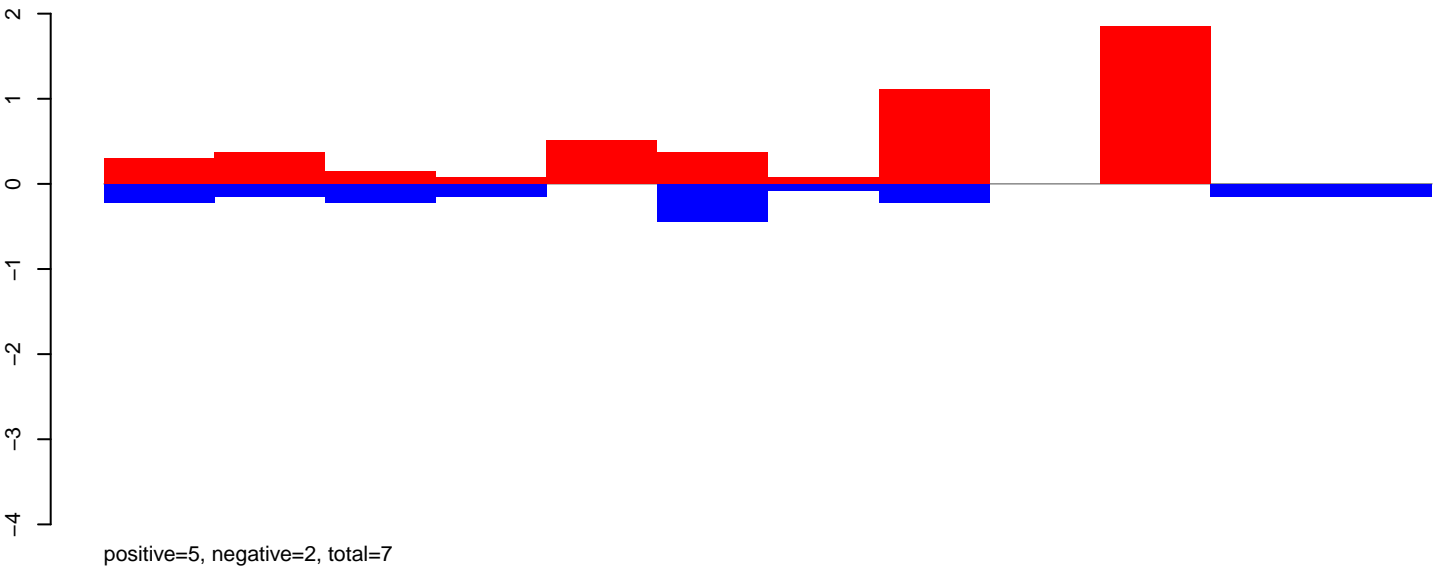
CuQuin_Testes_NL.rep



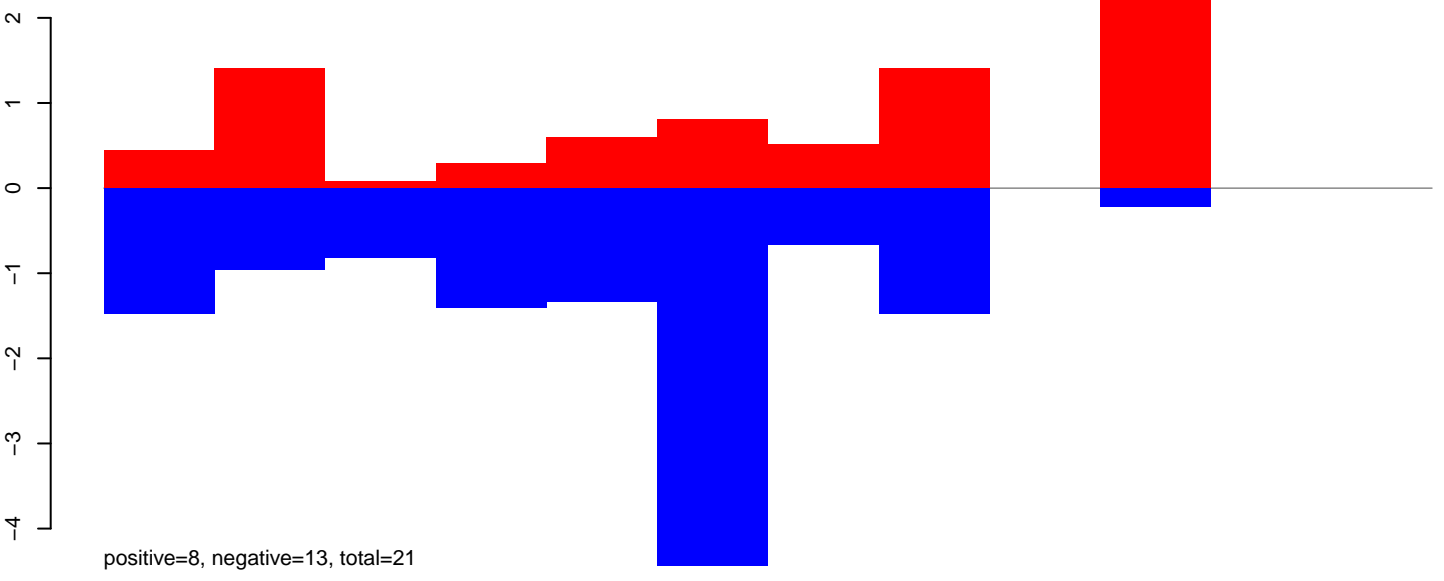
Window size=25, length=386, TE@rnd-4_family-631@DNA_MULE-MuDR@1607:1-386

100 200 300 400

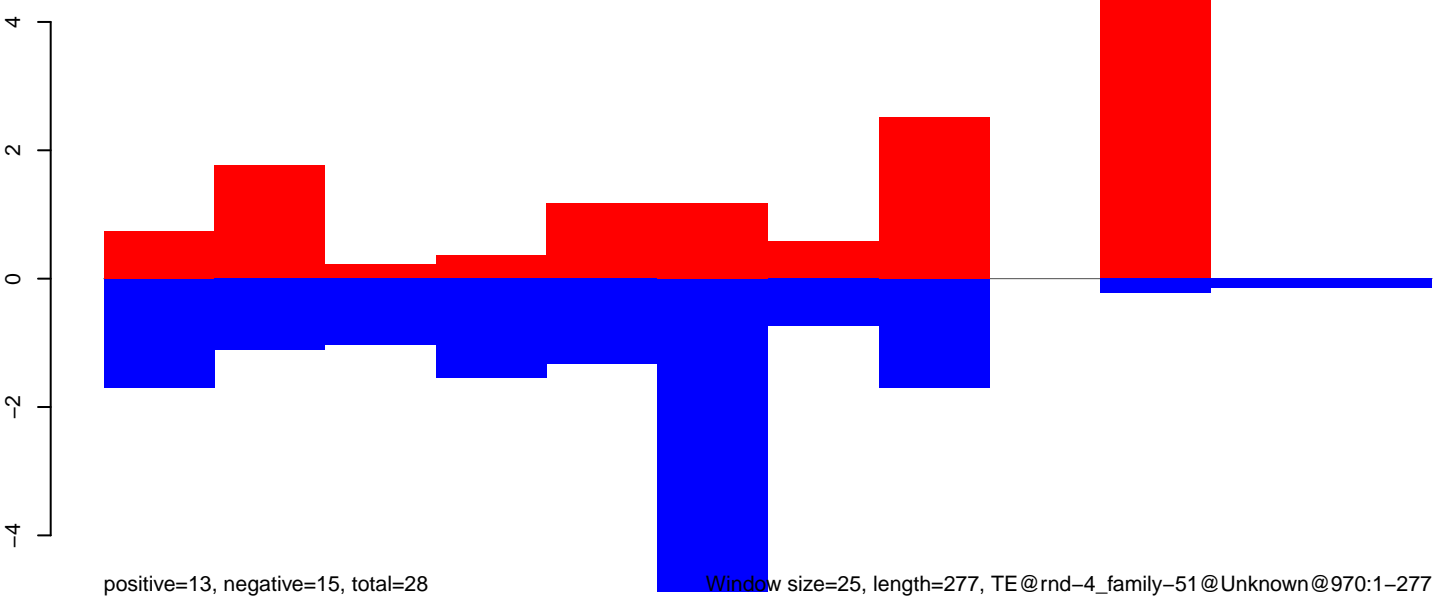
CuQuin_Testes_NL.18_23.rep



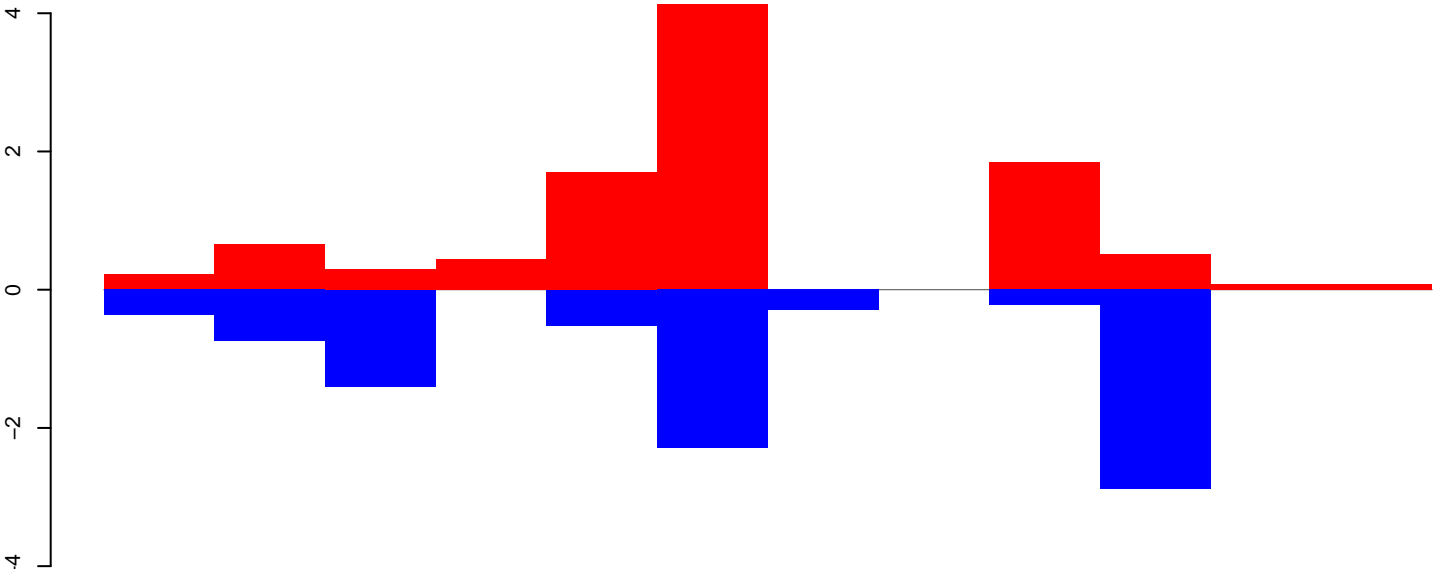
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

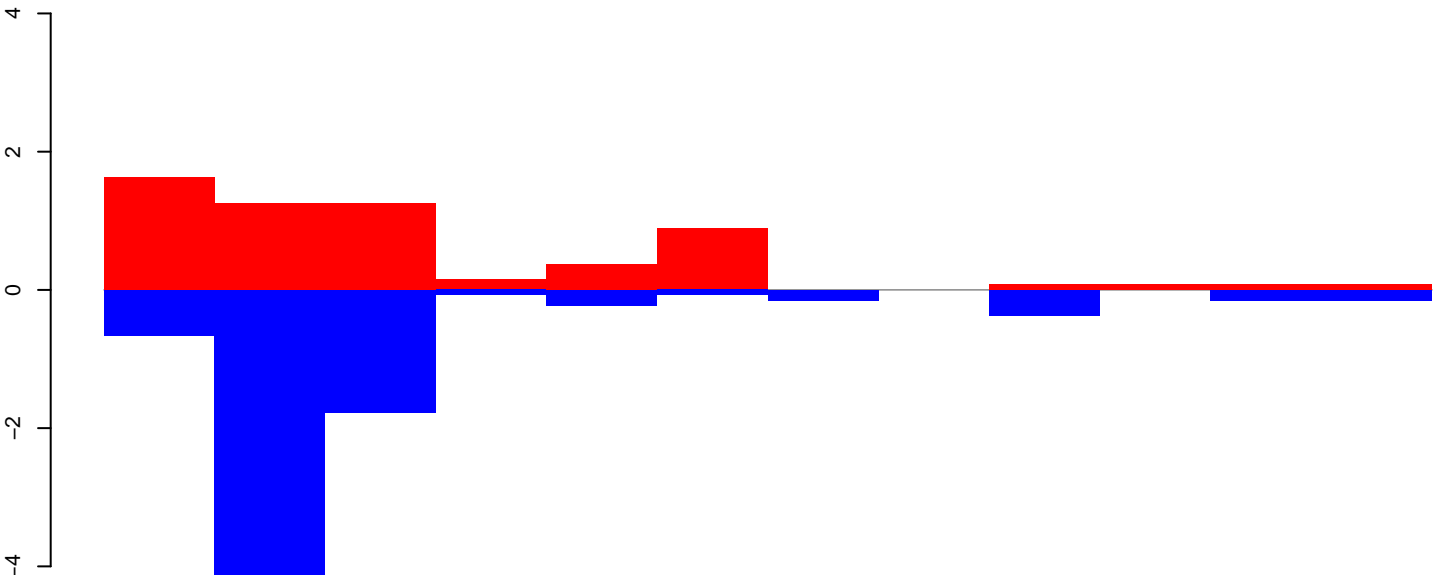


CuQuin_Testes_NL.18_23.rep



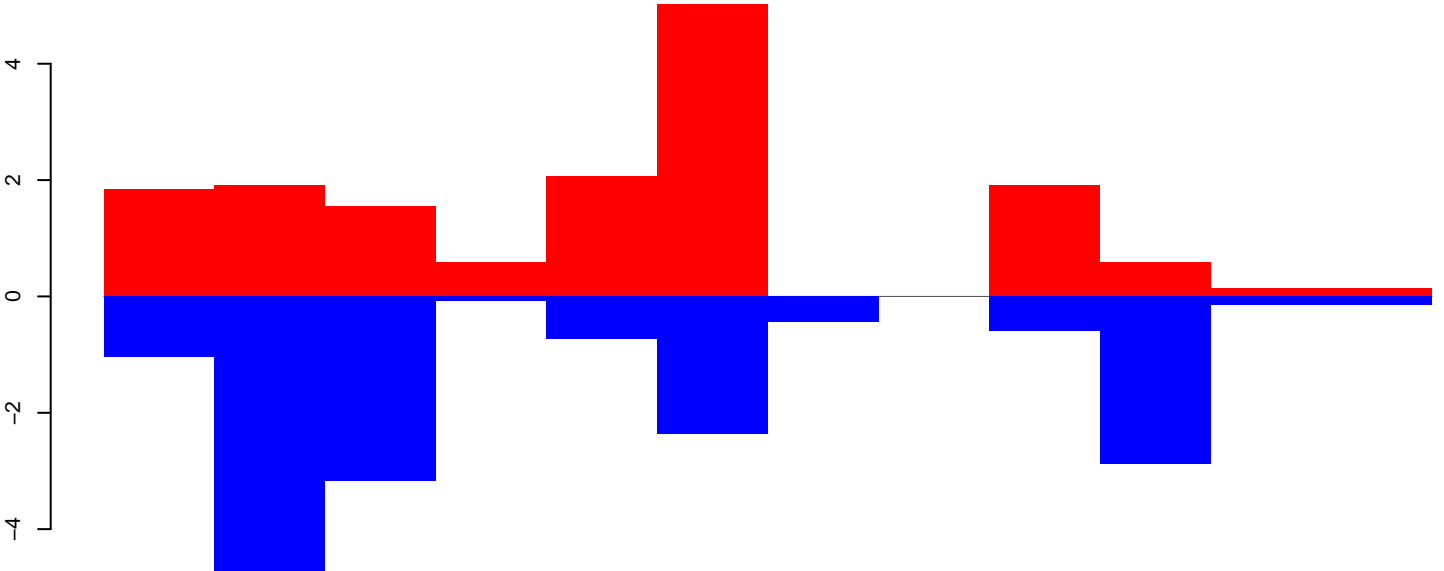
positive=10, negative=9, total=19

CuQuin_Testes_NL.24_35.rep



positive=6, negative=8, total=14

CuQuin_Testes_NL.rep

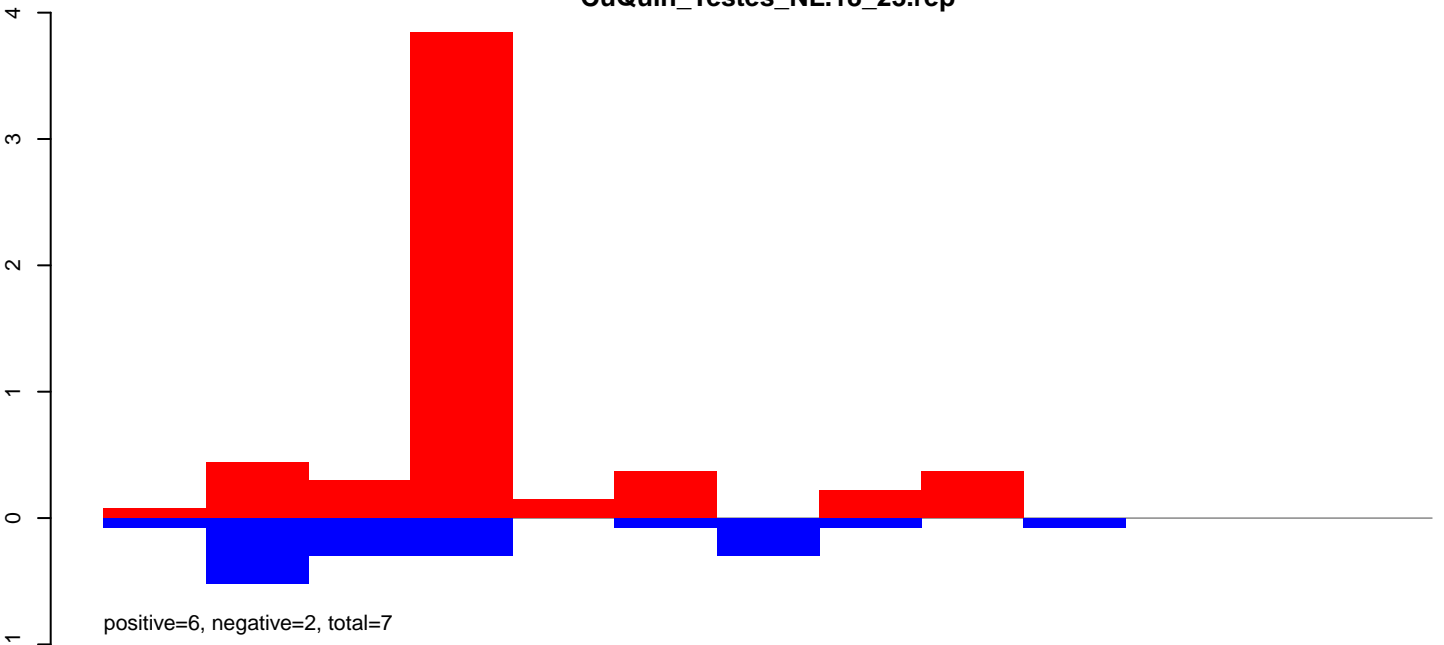


positive=16, negative=17, total=33

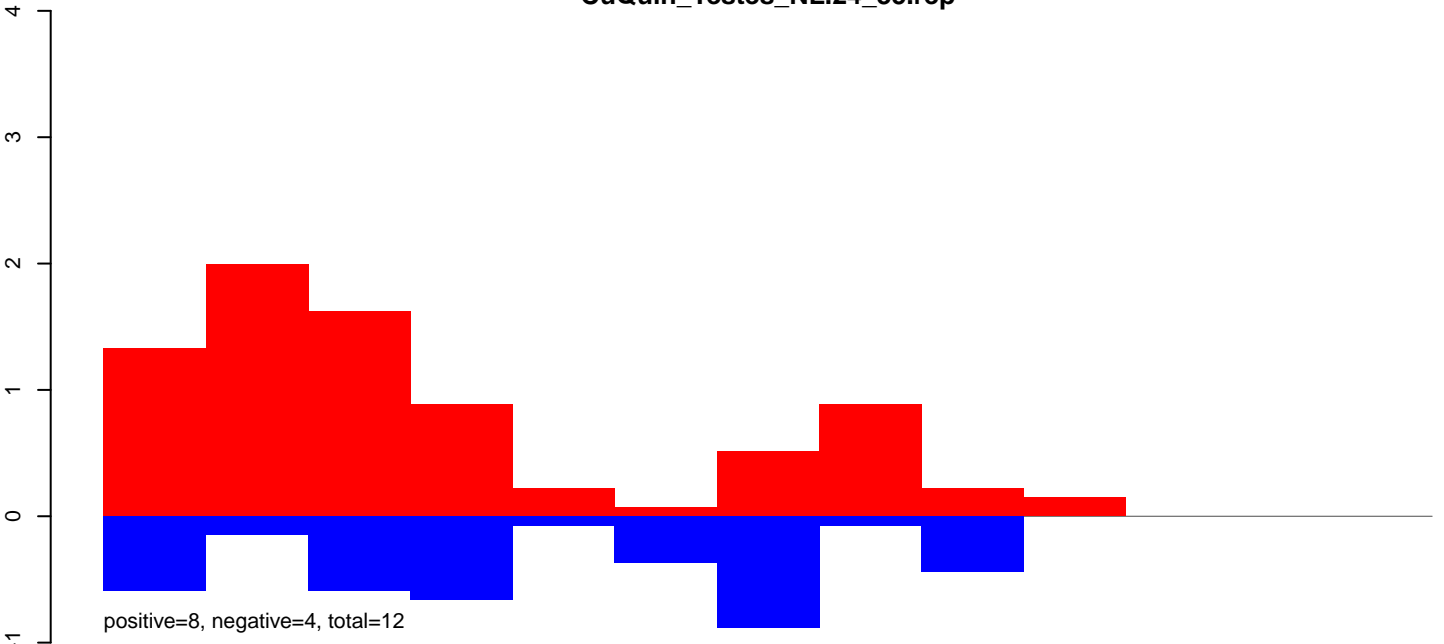
Window size=25, length=281, TE@rnd-4_family-437@SINE_tRNA-Deu@1443:1-281

50 100 150 200 250 300

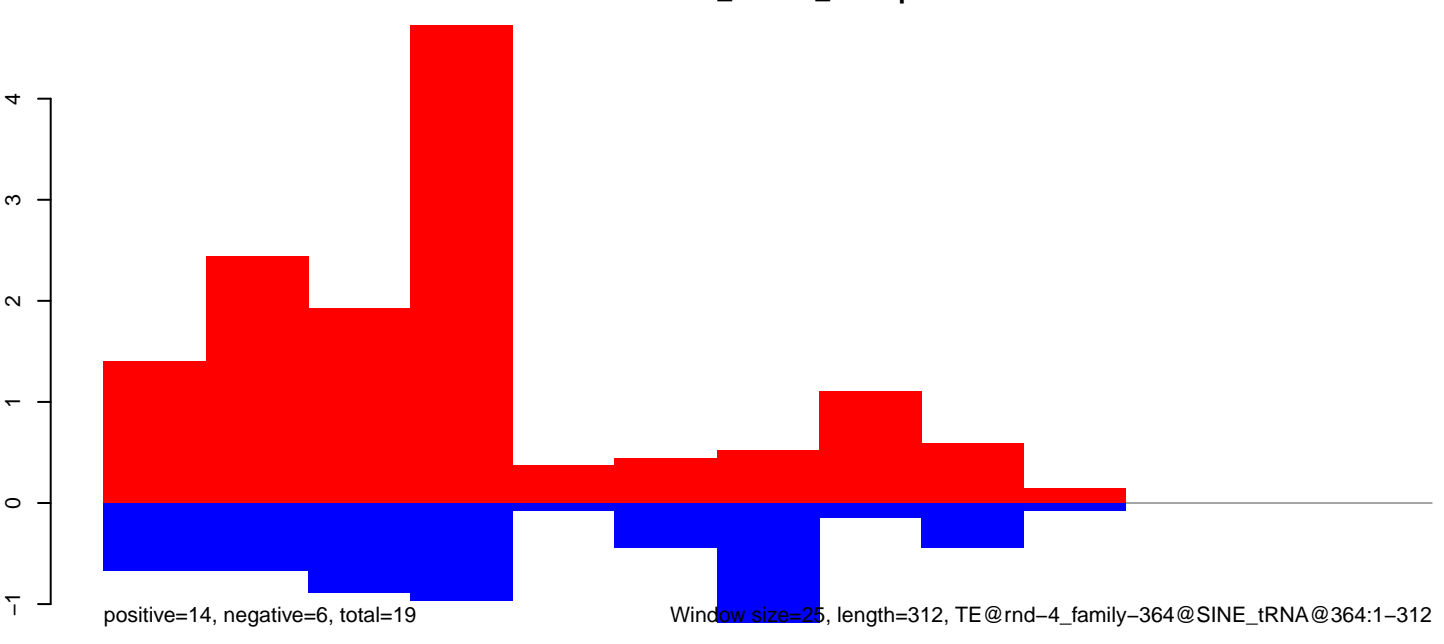
CuQuin_Testes_NL.18_23.rep



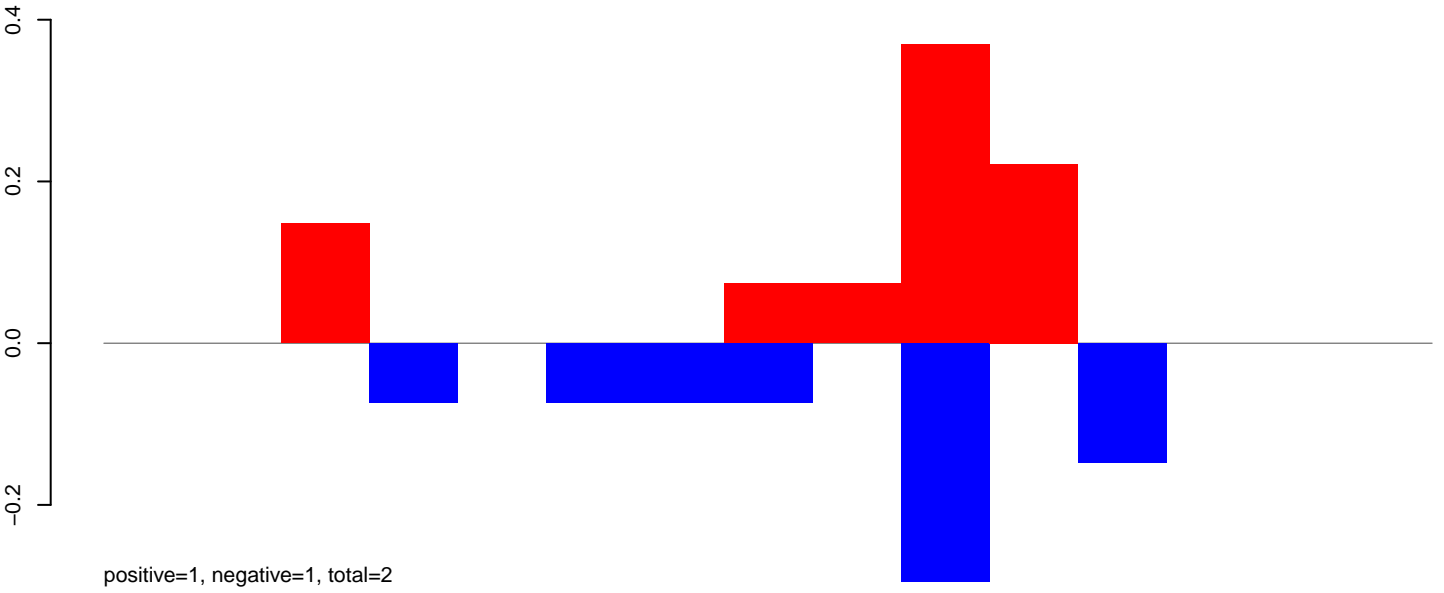
CuQuin_Testes_NL.24_35.rep



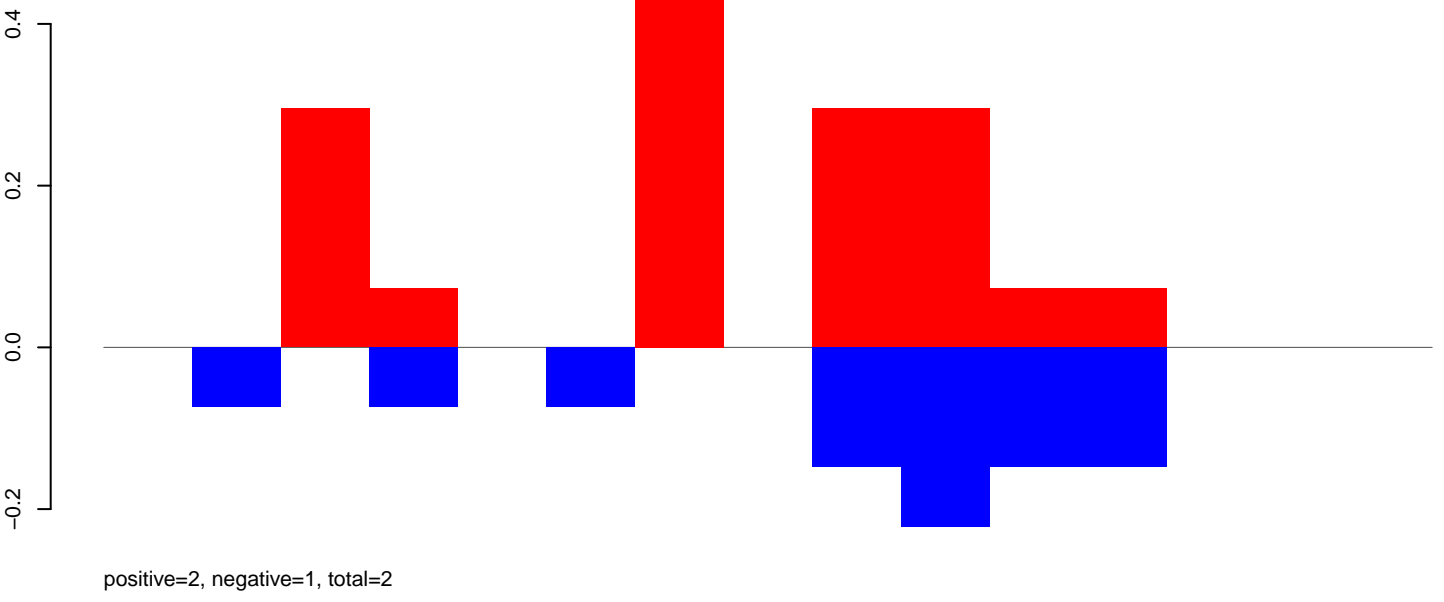
CuQuin_Testes_NL.rep



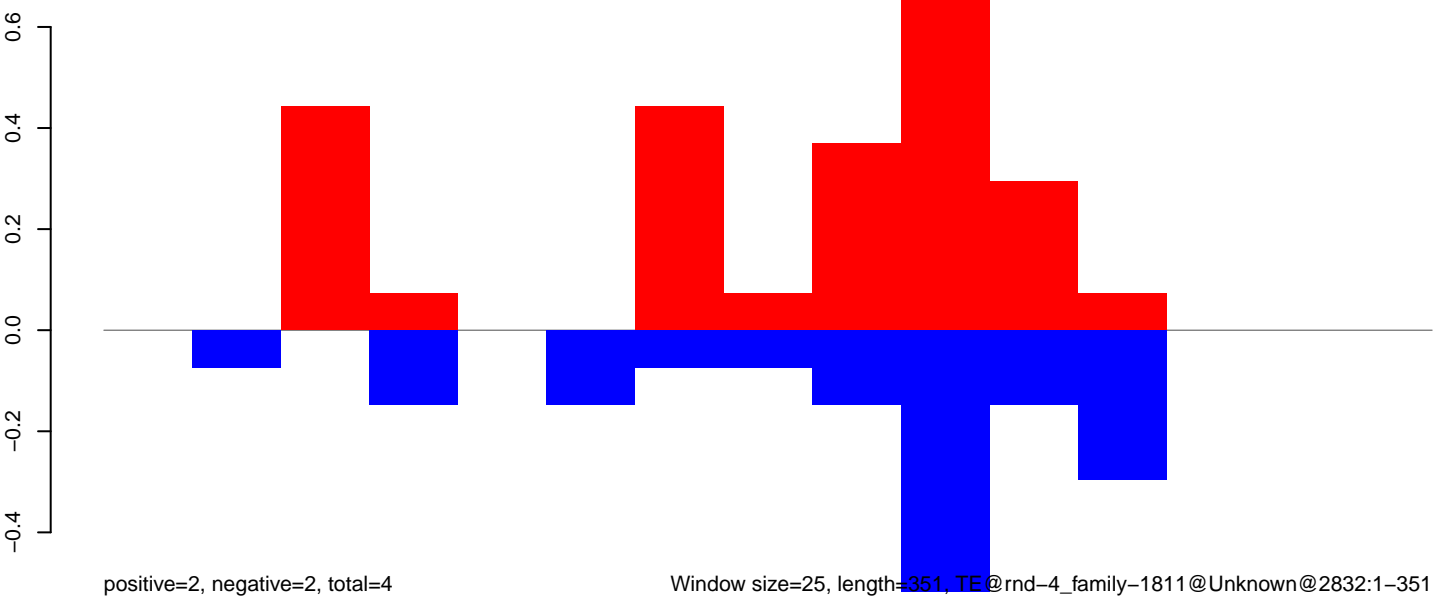
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

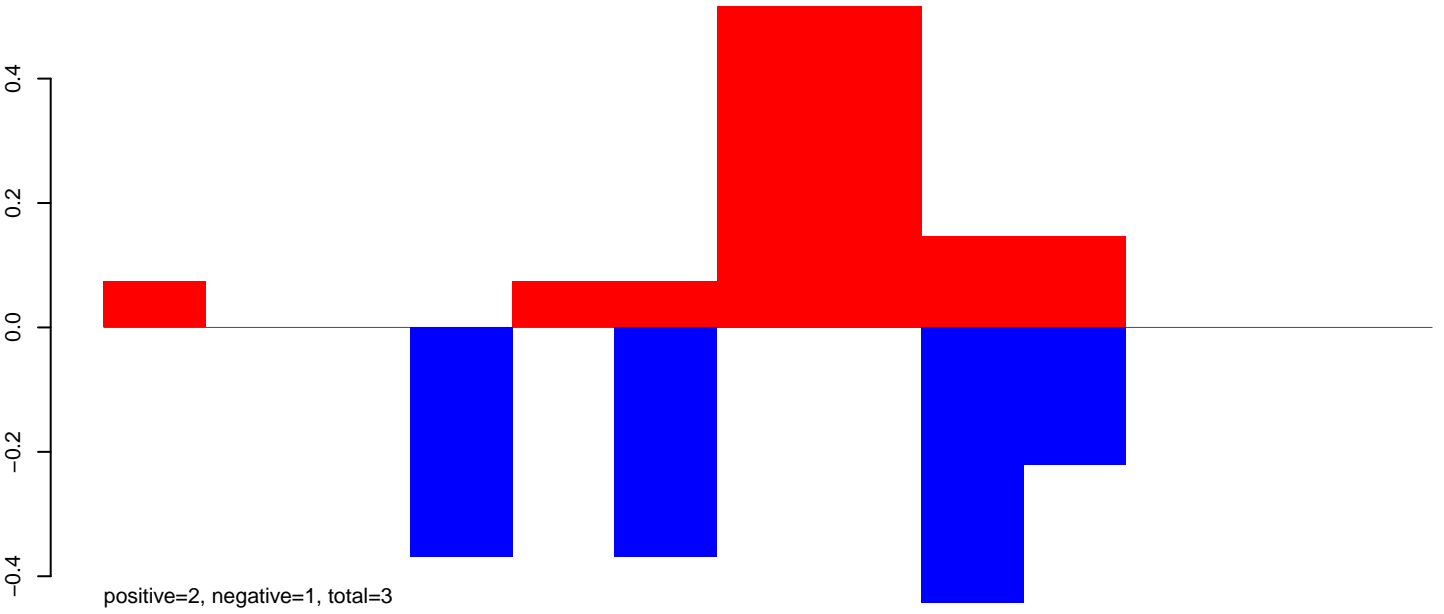


CuQuin_Testes_NL.rep

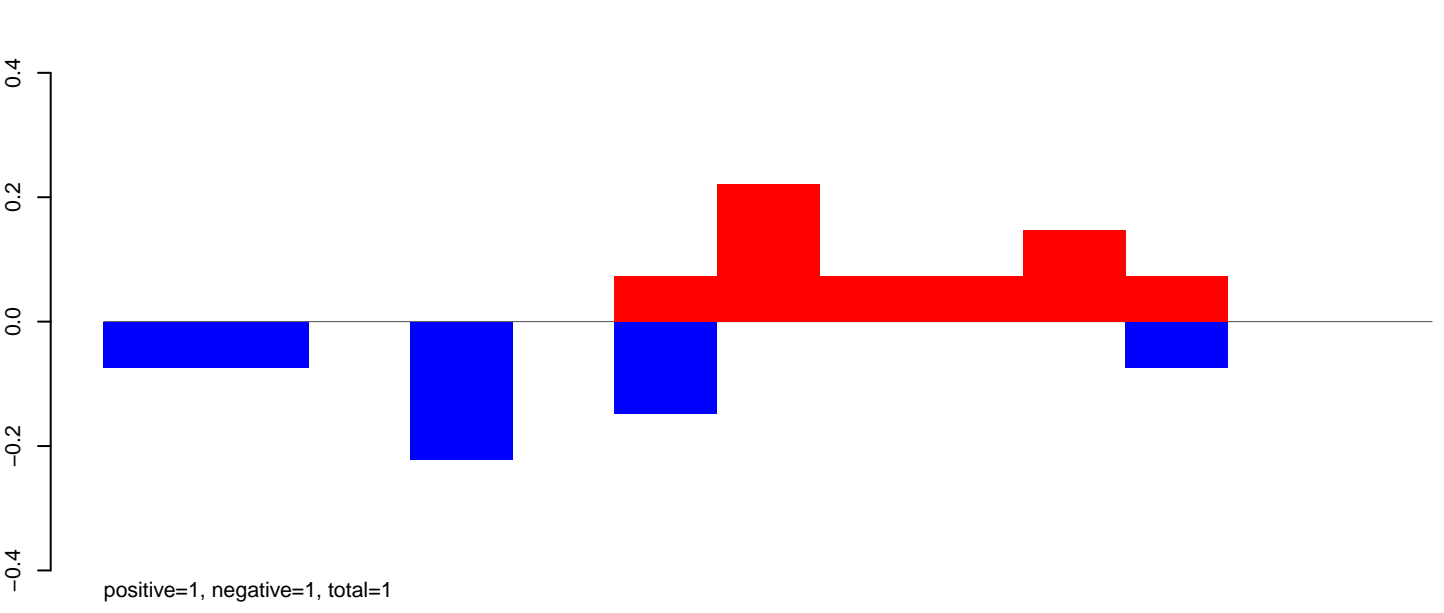


Window size=25, length=351, TE@rnd-4_family-1811@Unknown@2832:1-351

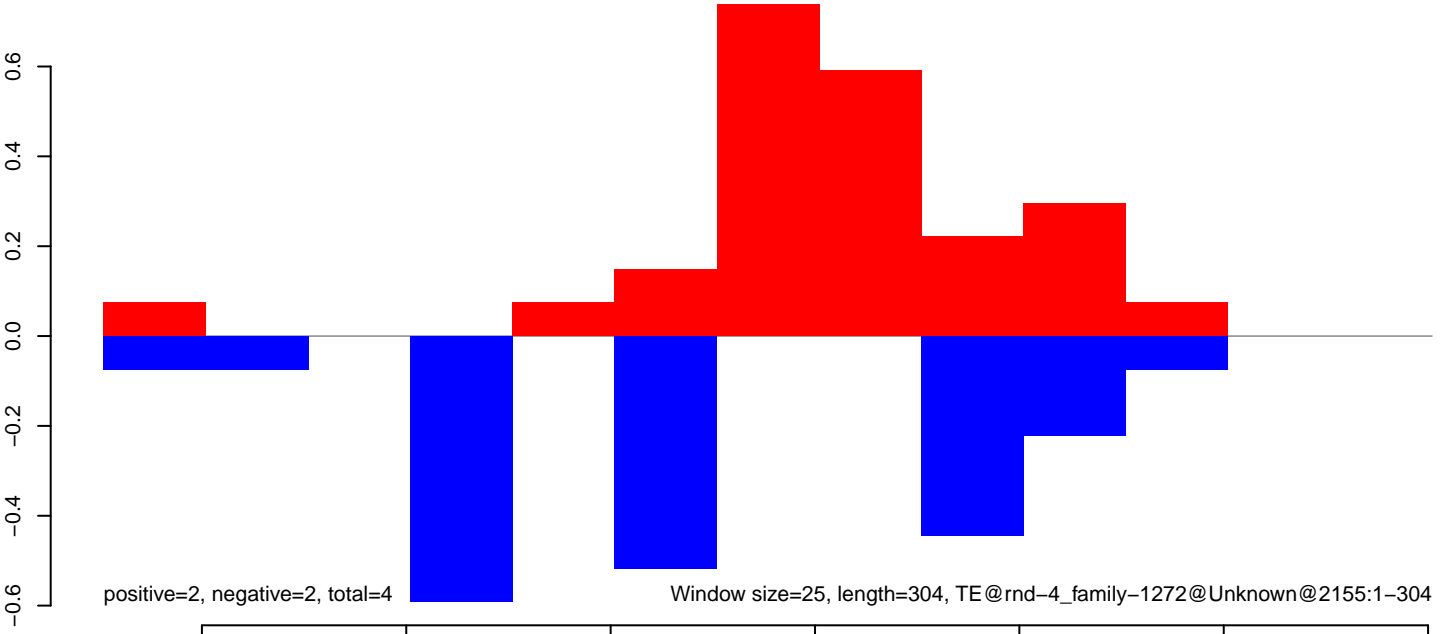
CuQuin_Testes_NL.18_23.rep



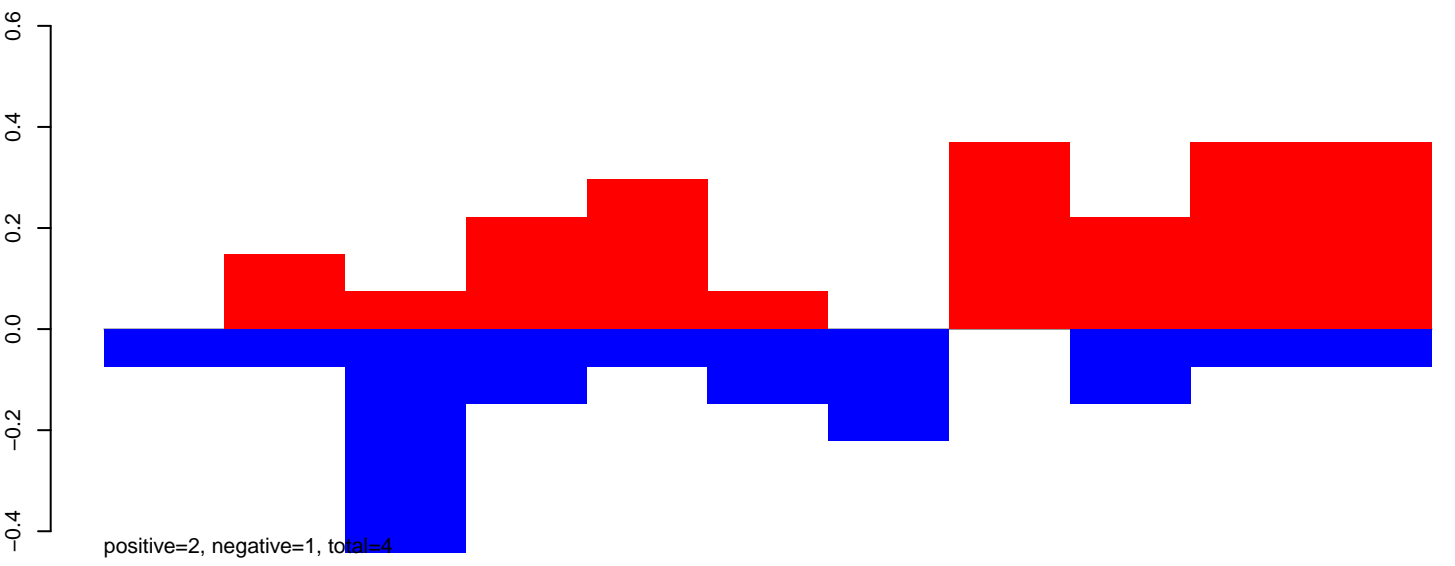
CuQuin_Testes_NL.24_35.rep



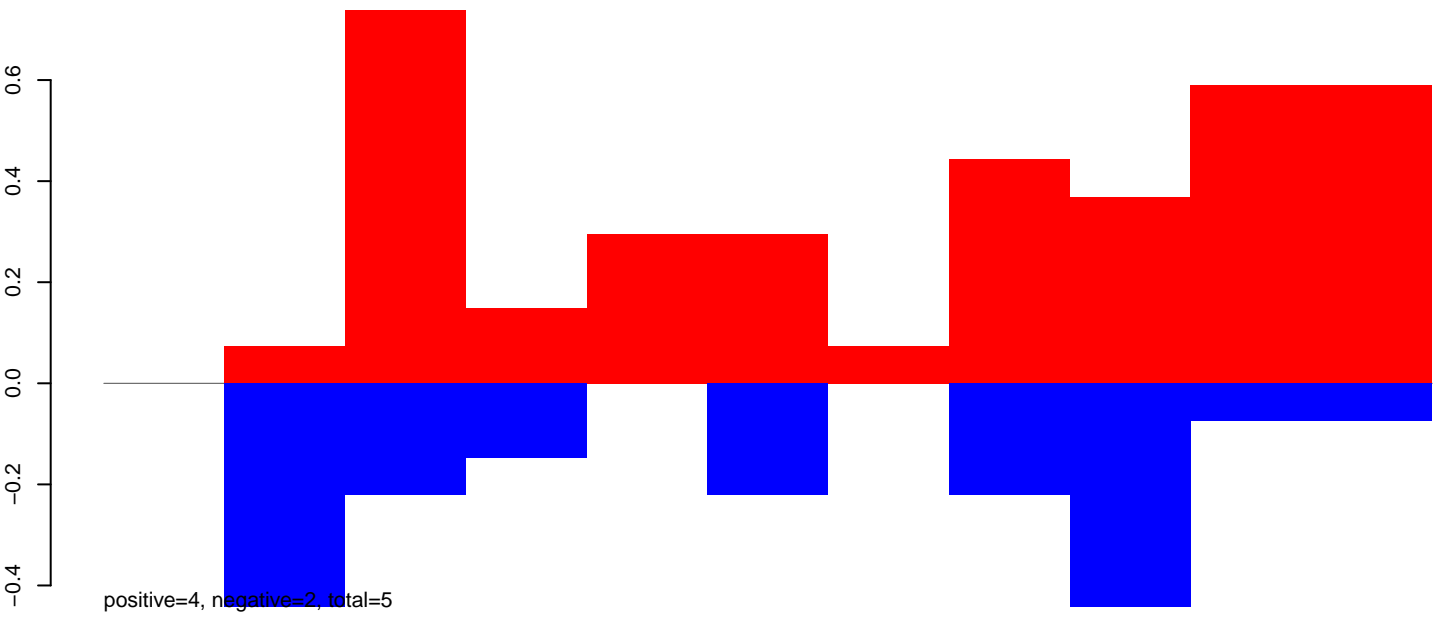
CuQuin_Testes_NL.rep



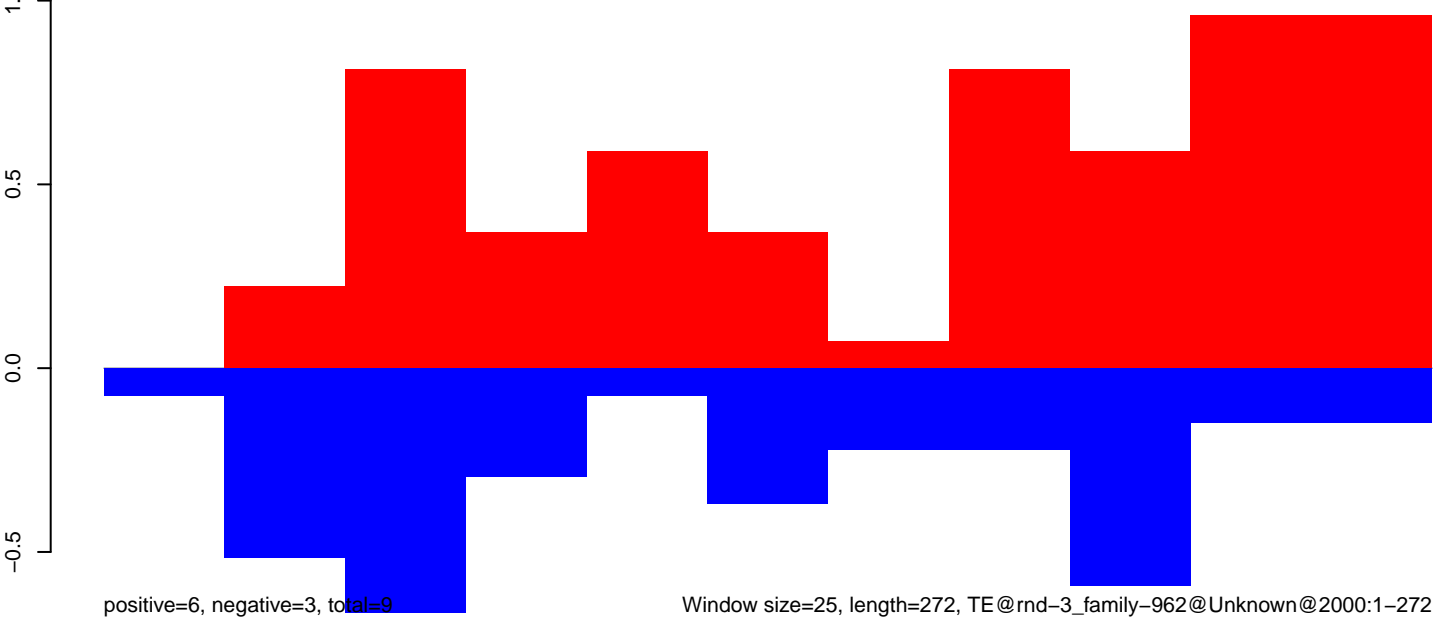
CuQuin_Testes_NL.18_23.rep



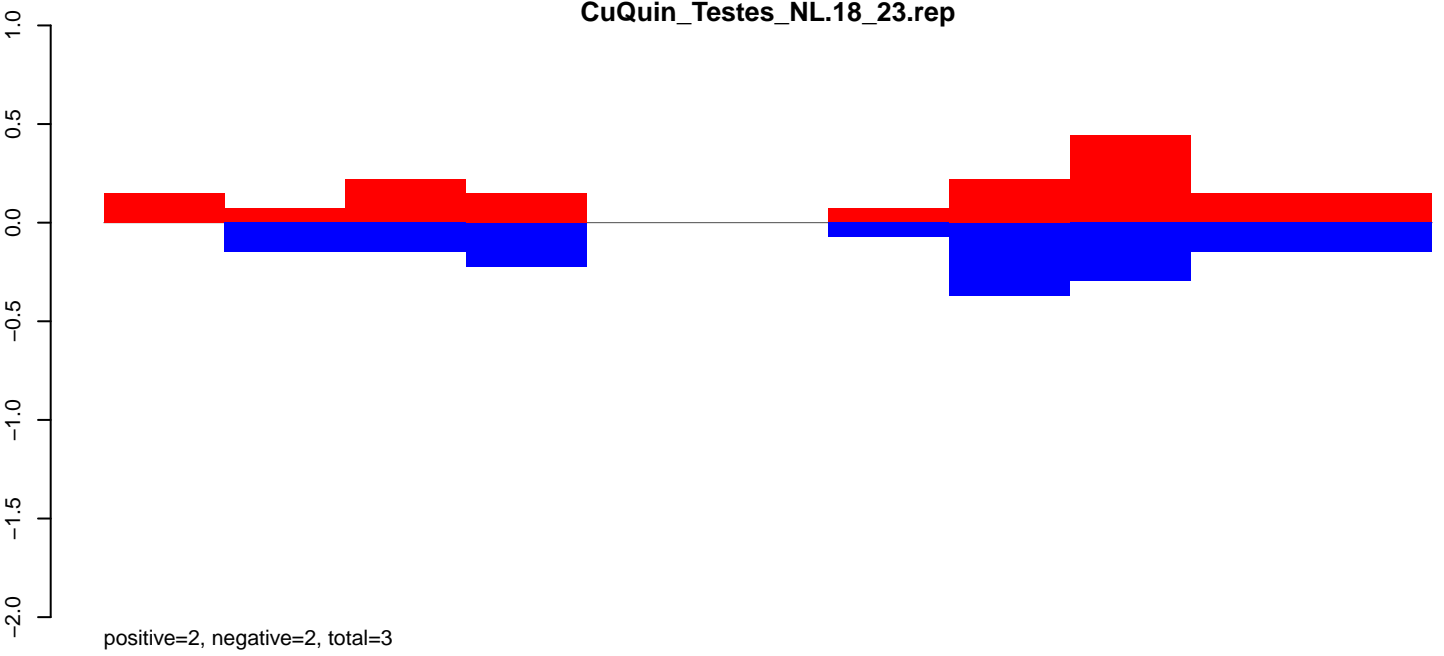
CuQuin_Testes_NL.24_35.rep



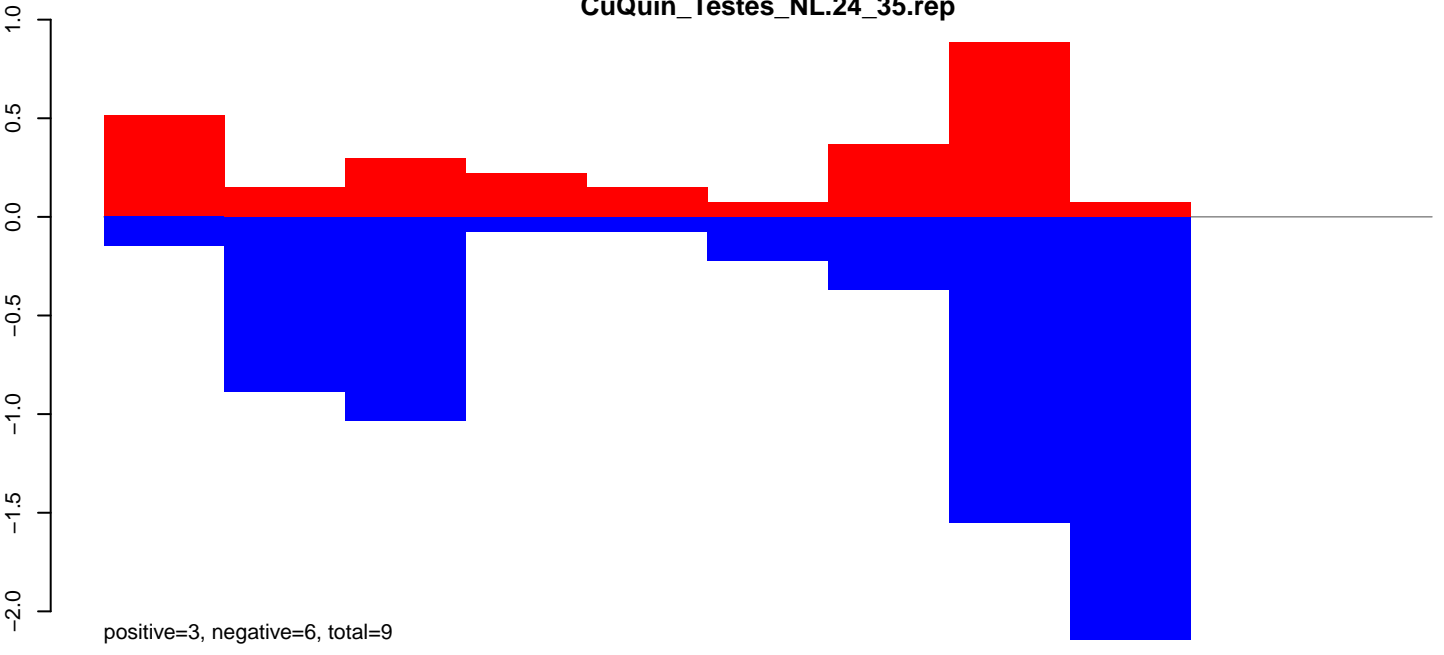
CuQuin_Testes_NL.rep



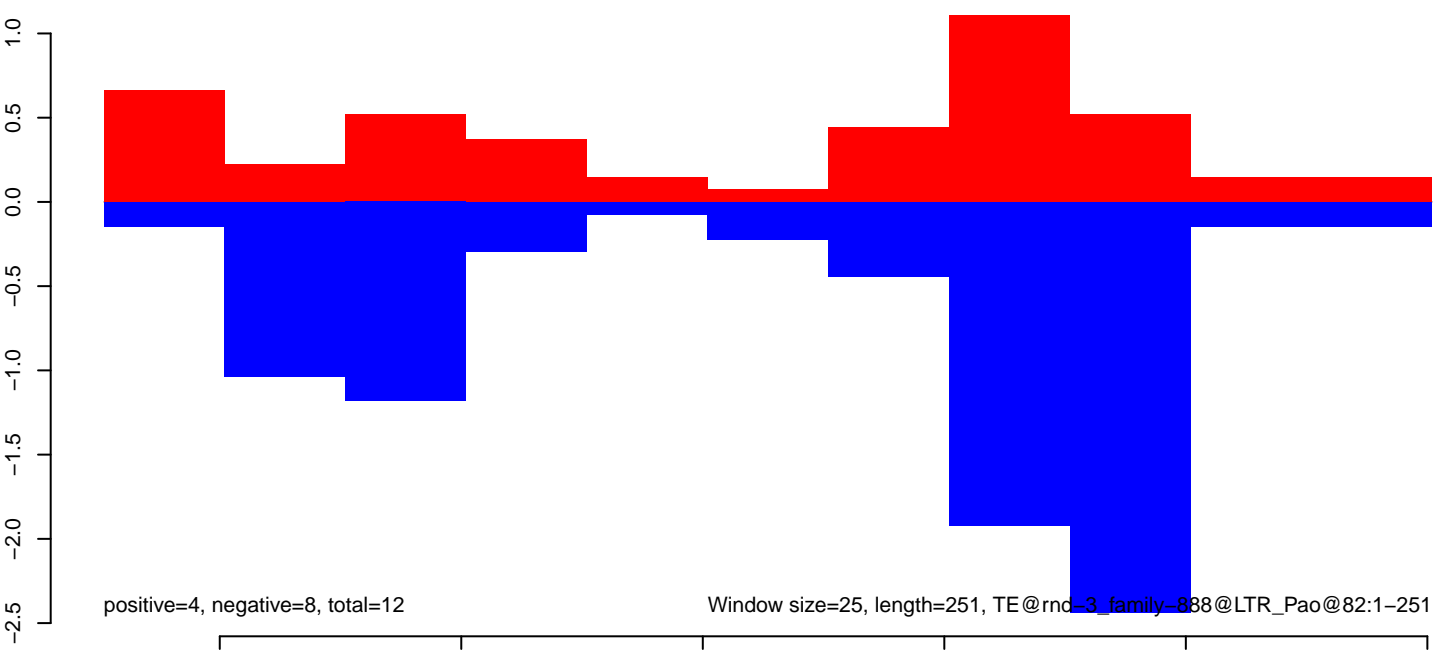
CuQuin_Testes_NL.18_23.rep



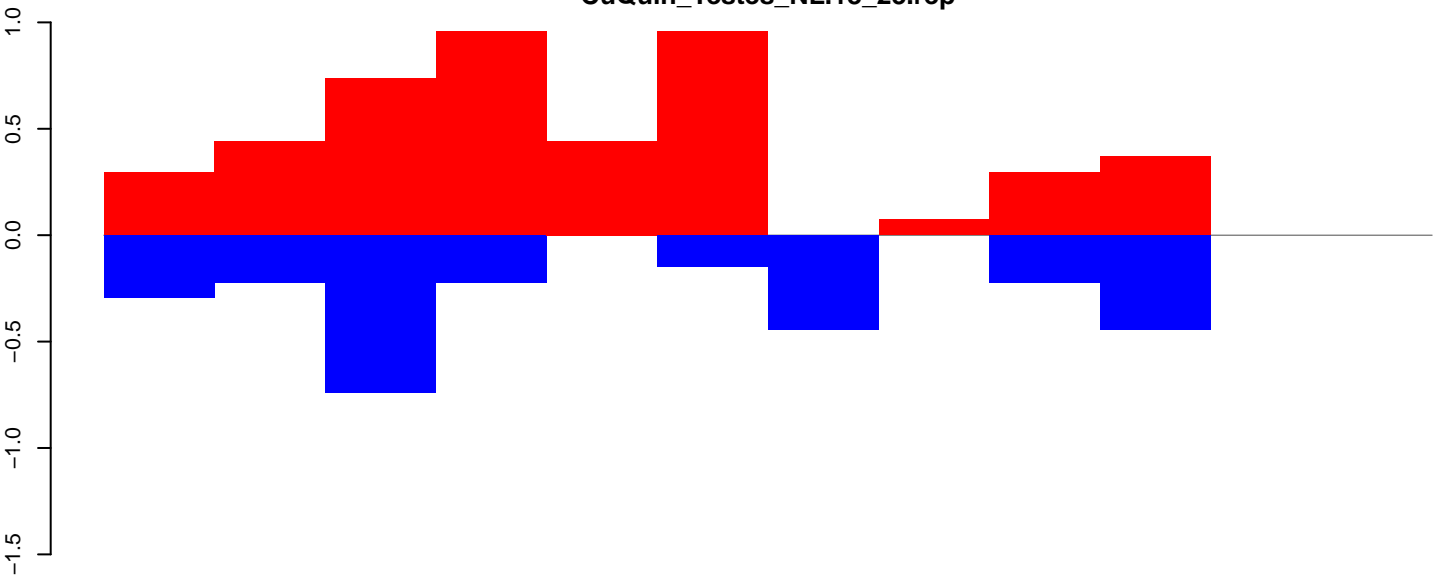
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

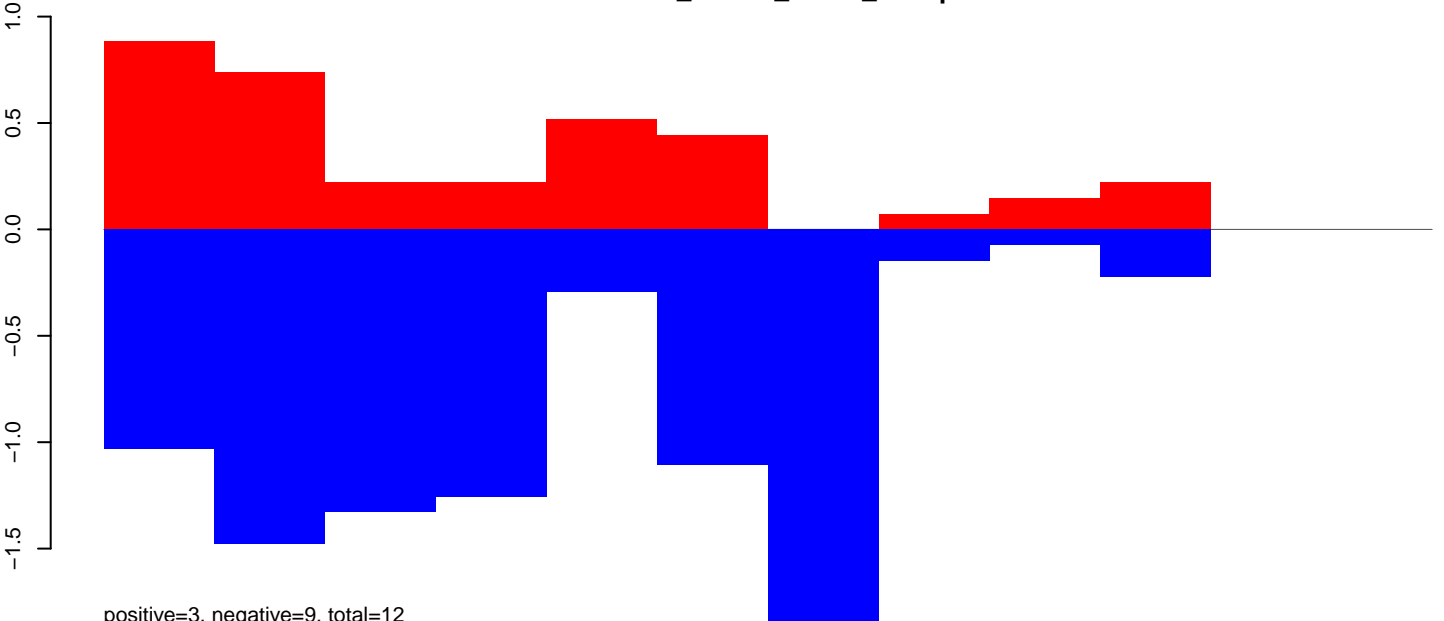


CuQuin_Testes_NL.18_23.rep



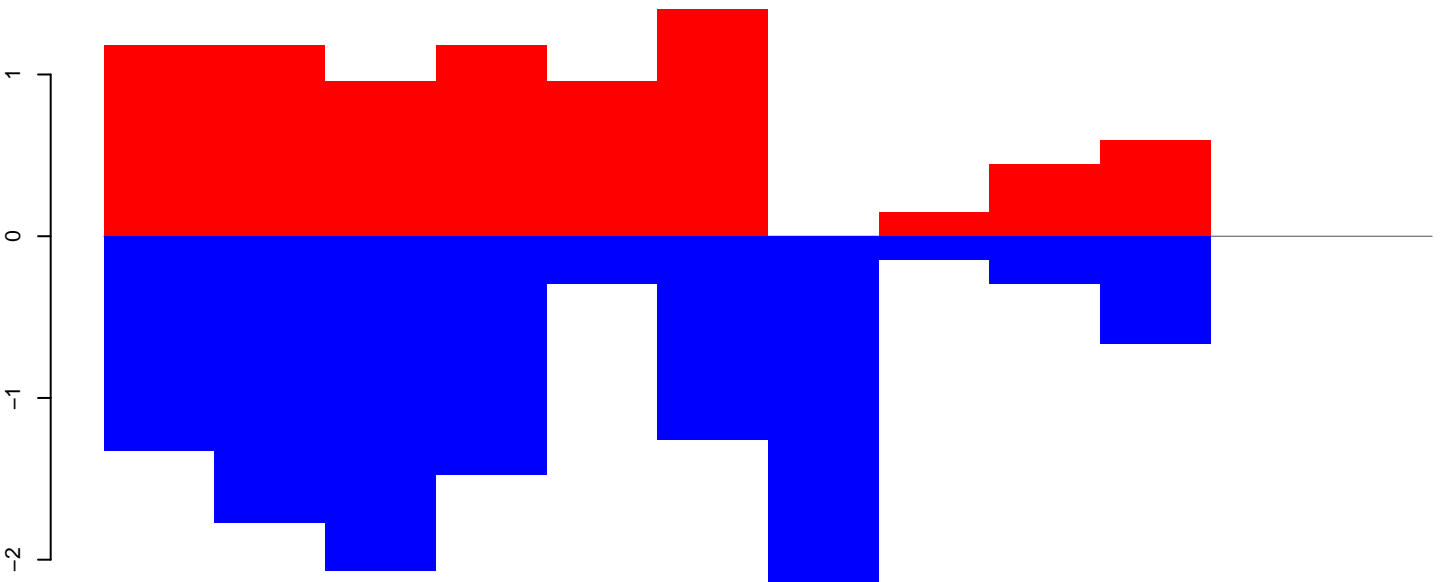
positive=5, negative=3, total=7

CuQuin_Testes_NL.24_35.rep



positive=3, negative=9, total=12

CuQuin_Testes_NL.rep

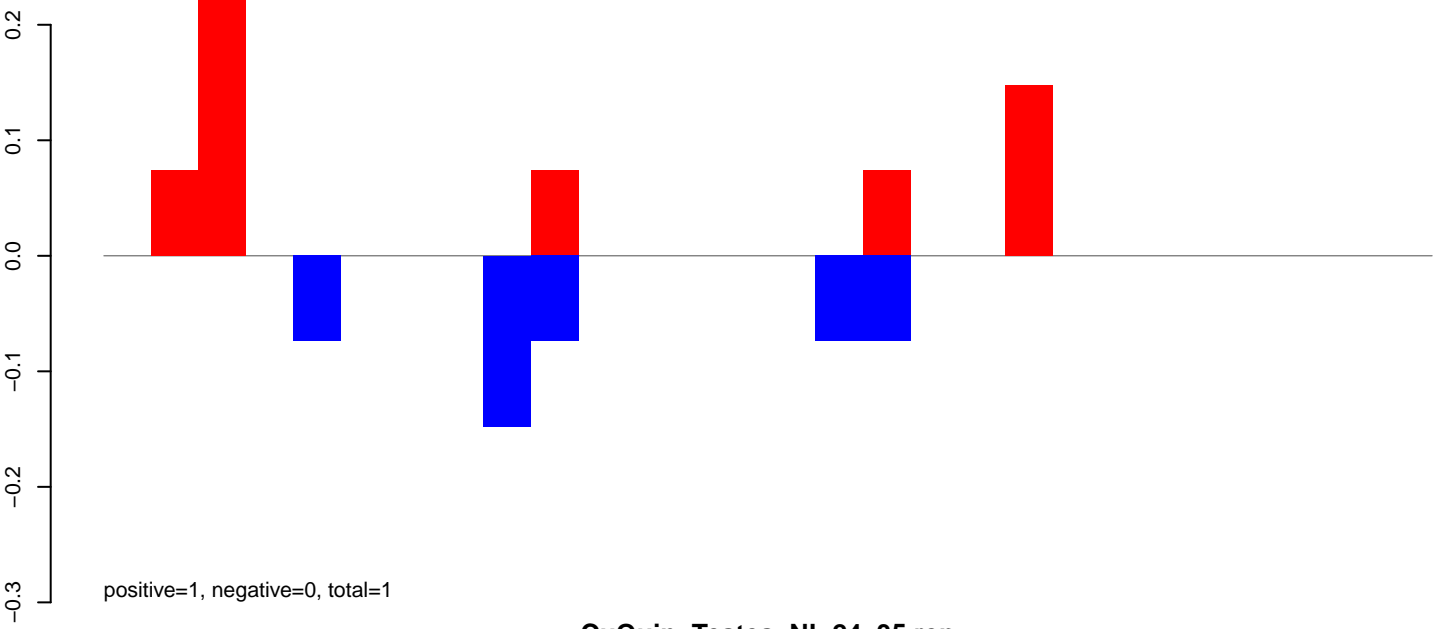


positive=8, negative=12, total=20

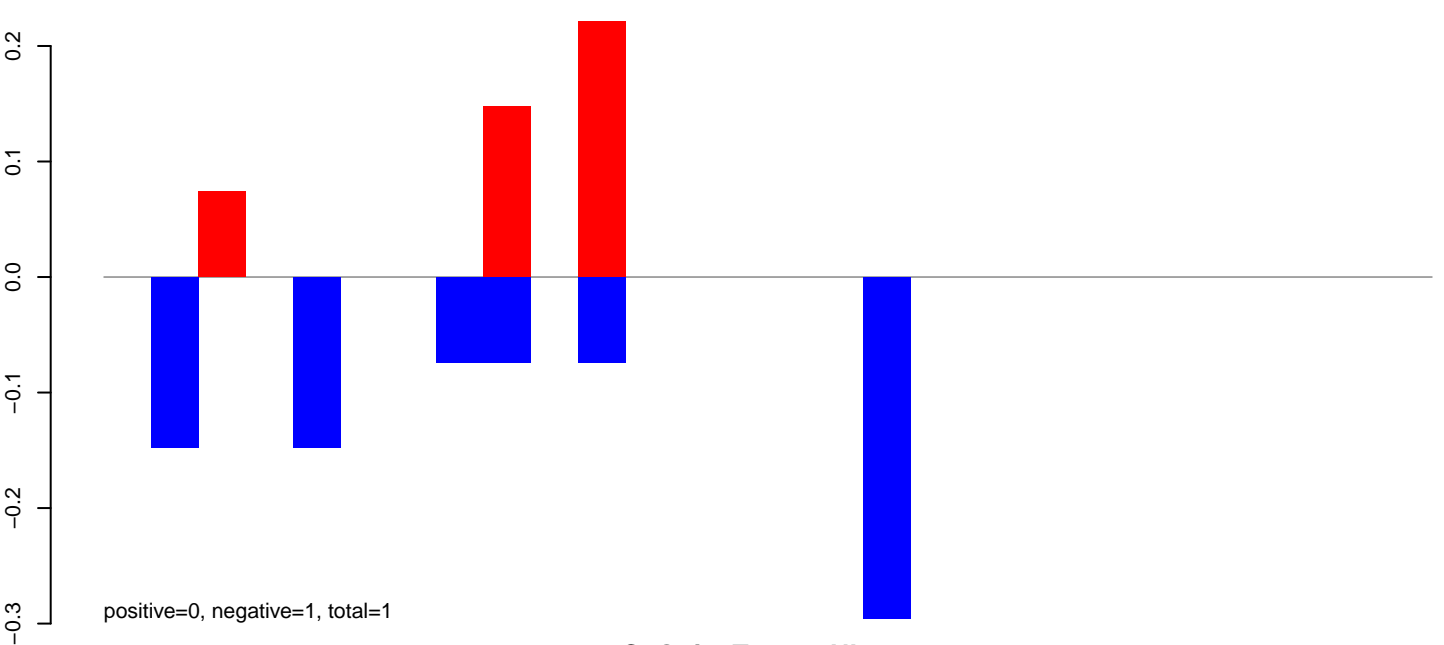
Window size=25, length=294, TE@rnd-3_family-131@Unknown@1295:1-294

50 100 150 200 250 300

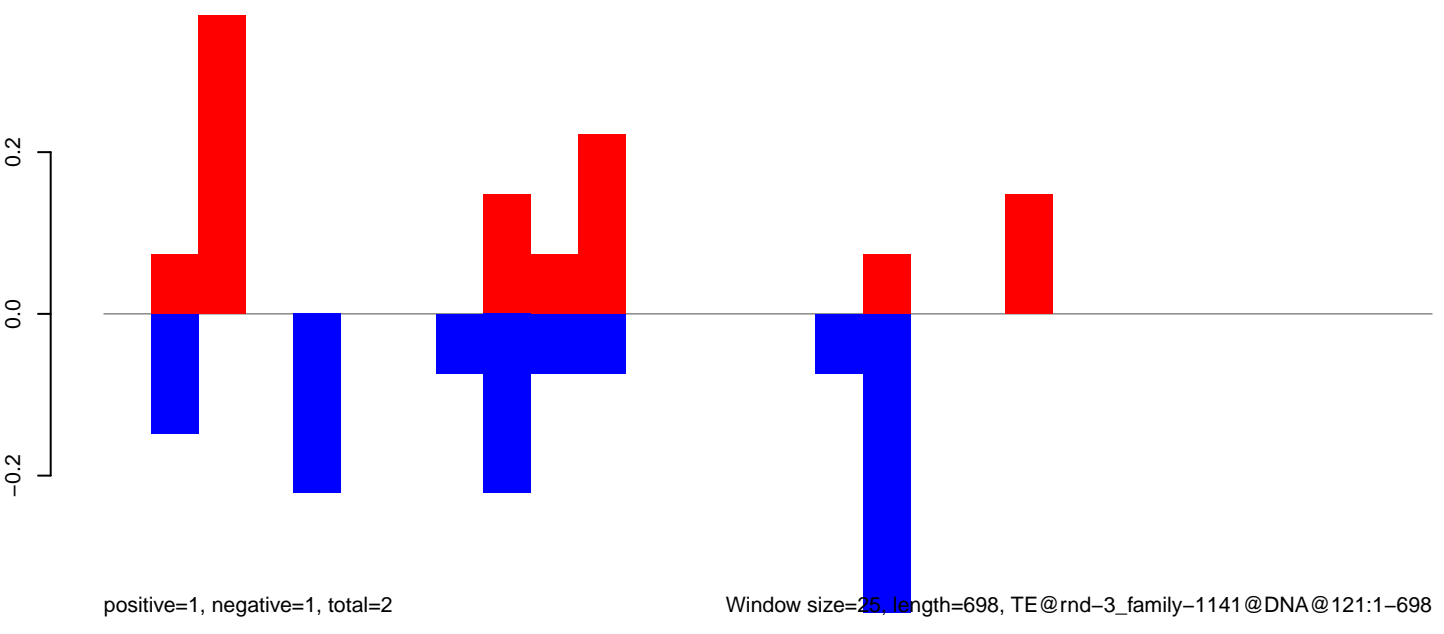
CuQuin_Testes_NL.18_23.rep



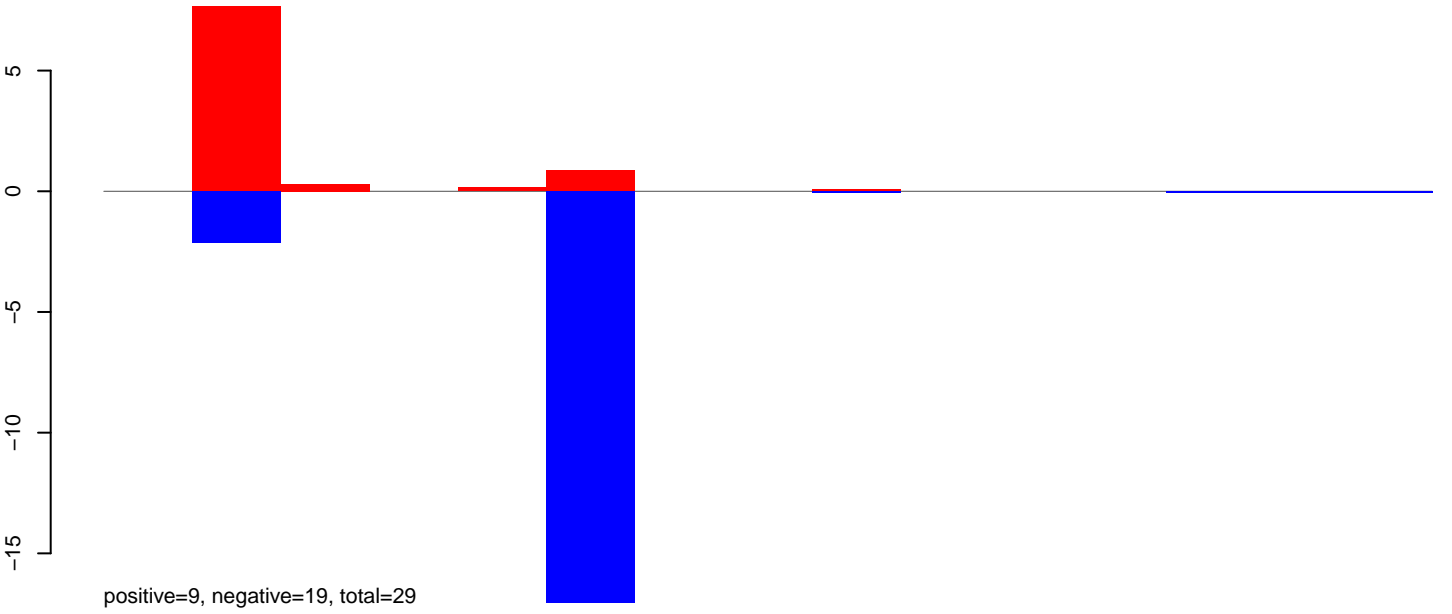
CuQuin_Testes_NL.24_35.rep



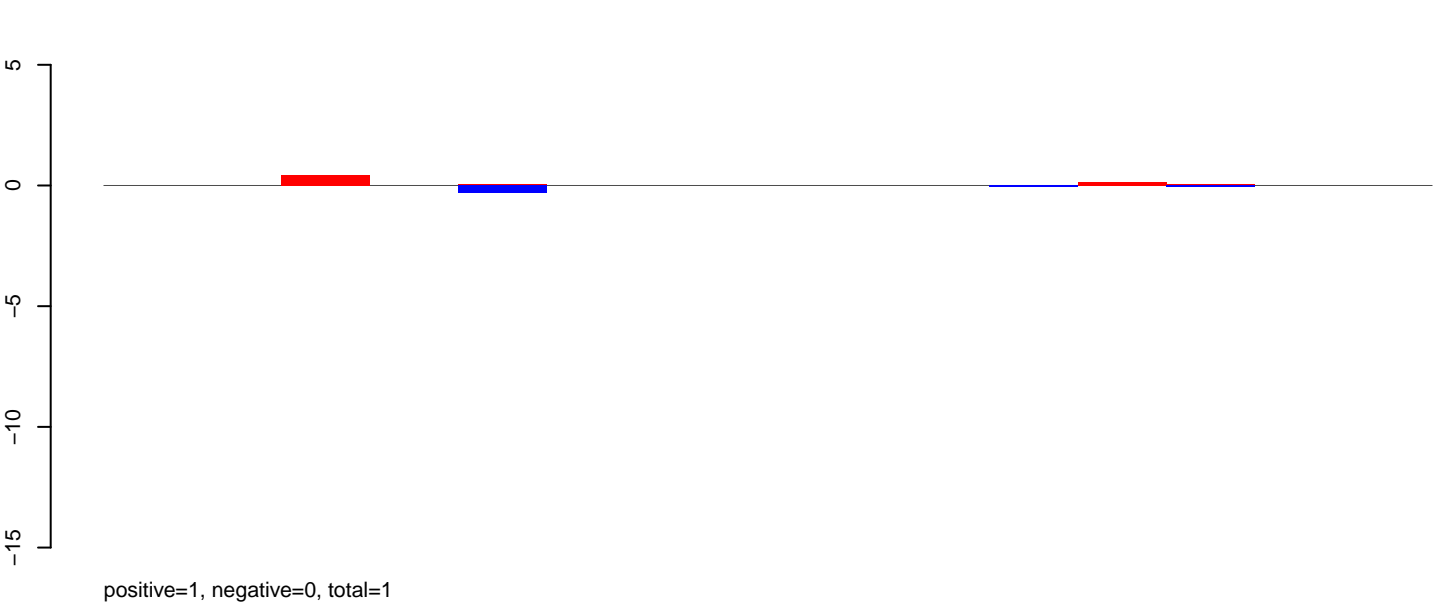
CuQuin_Testes_NL.rep



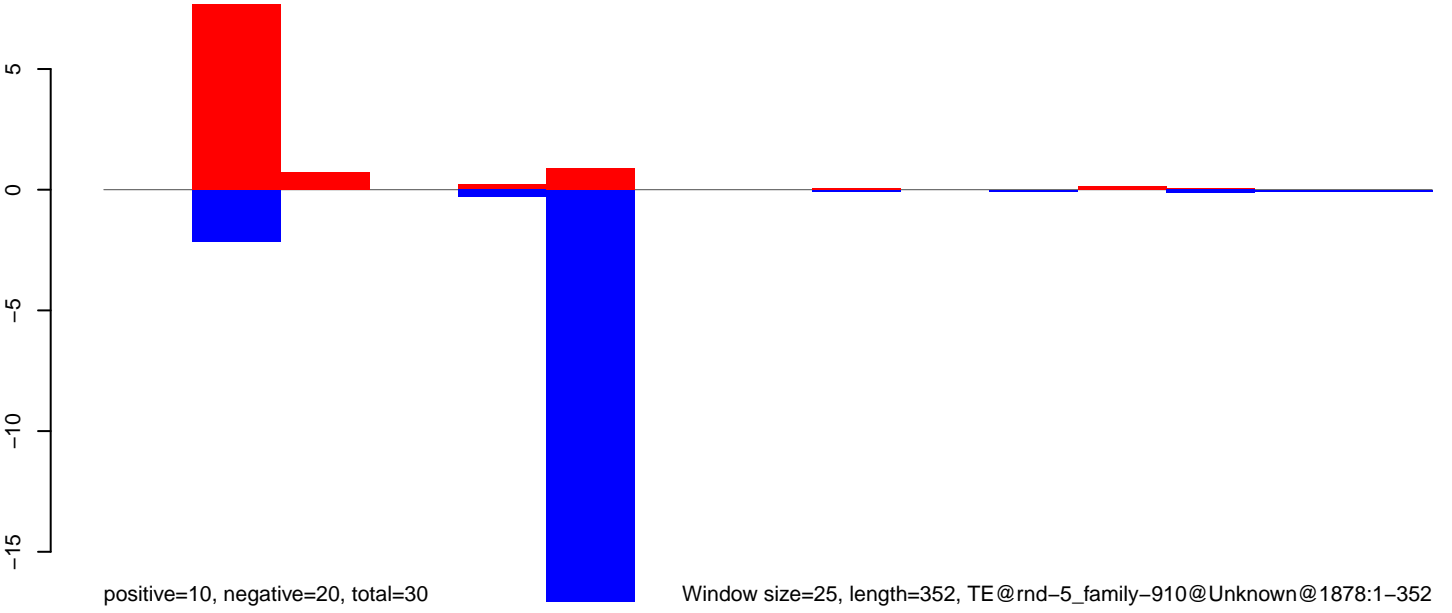
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



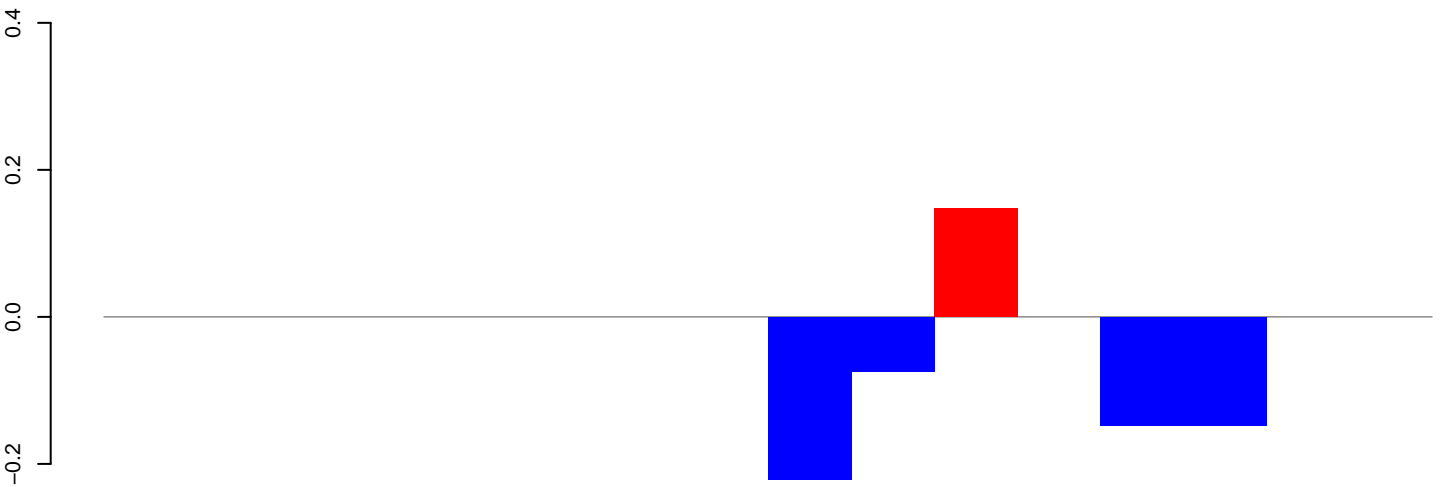
CuQuin_Testes_NL.rep



Window size=25, length=352, TE@rnd-5_family-910@Unknown@1878:1-352

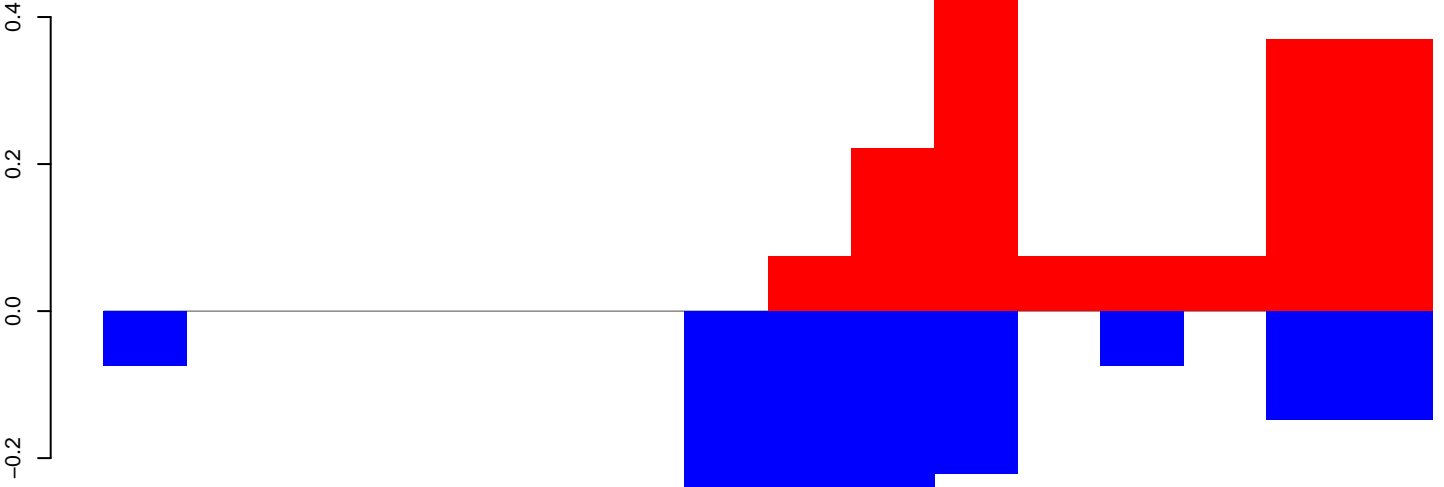
100 200 300 400

CuQuin_Testes_NL.18_23.rep



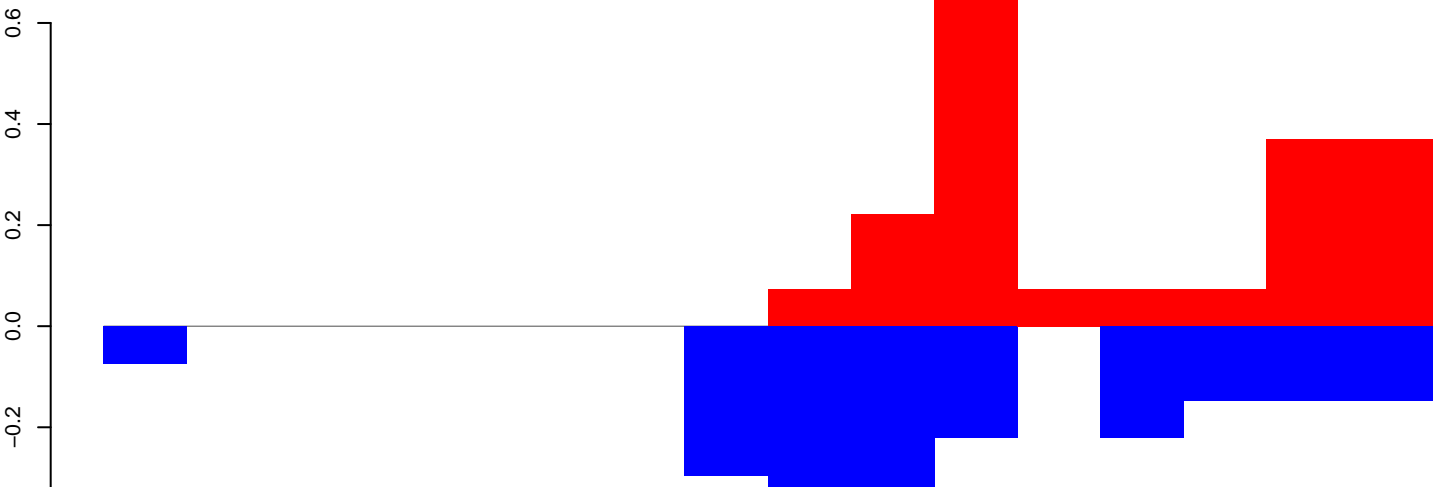
positive=0, negative=1, total=1

CuQuin_Testes_NL.24_35.rep



positive=2, negative=2, total=3

CuQuin_Testes_NL.rep

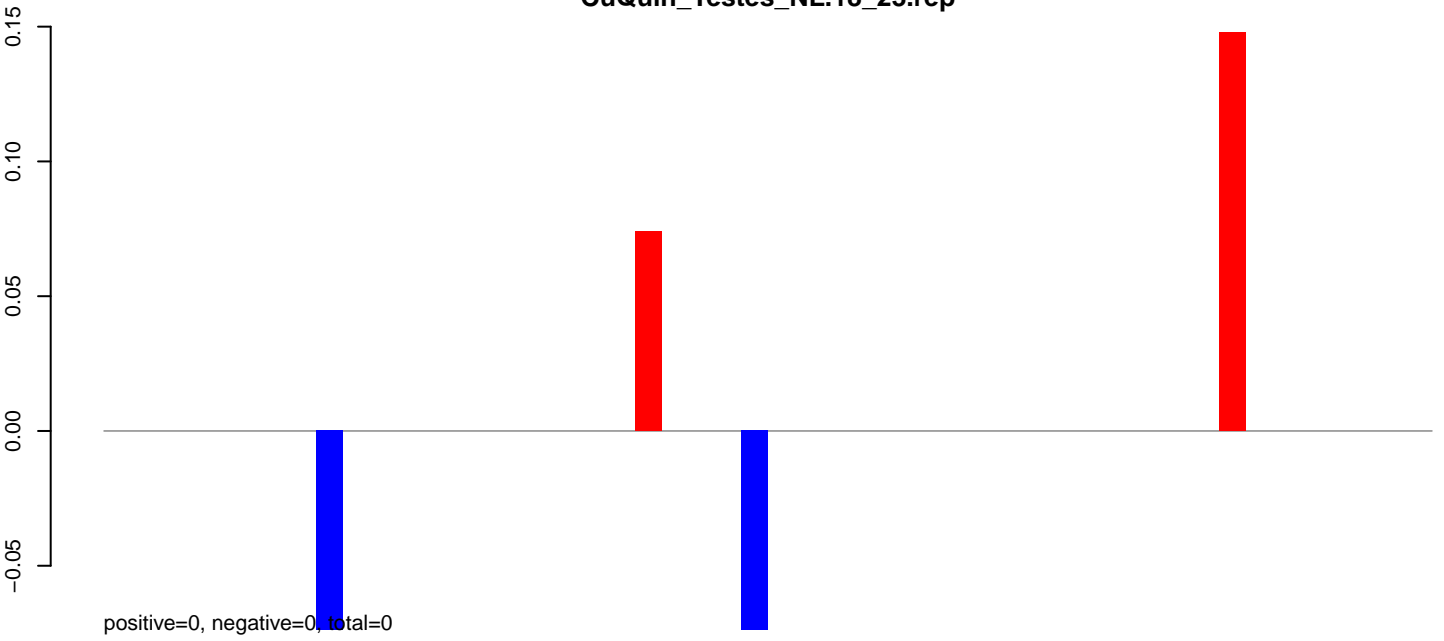


positive=2, negative=2, total=4

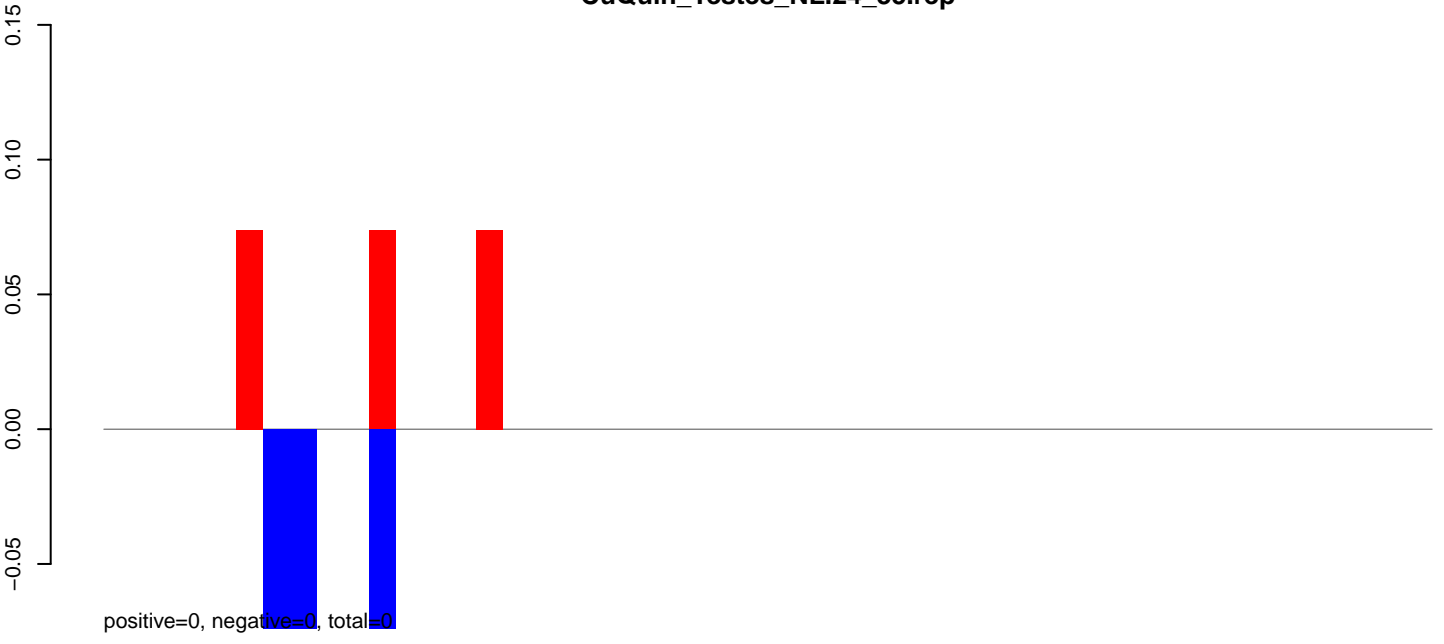
Window size=25, length=398, TE@rnd-5_family-860@DNA_hAT-Tip100@625:1-398

100 200 300 400

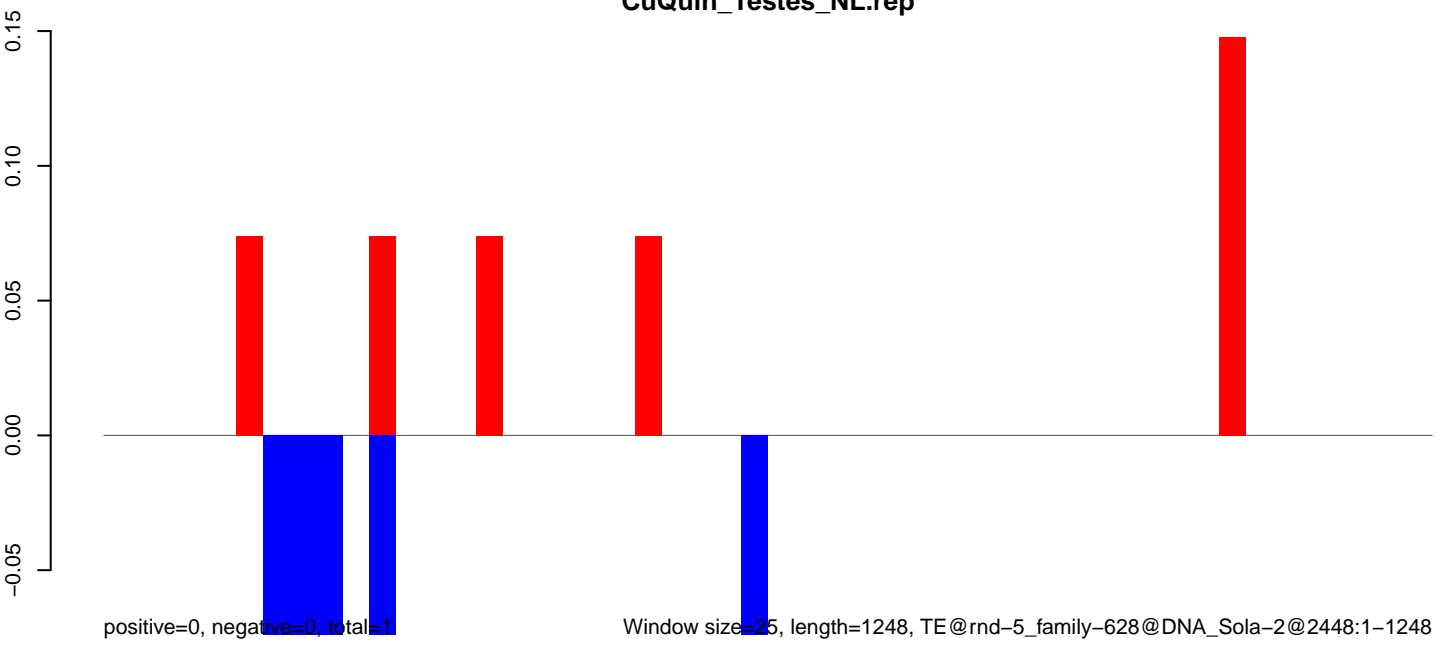
CuQuin_Testes_NL.18_23.rep



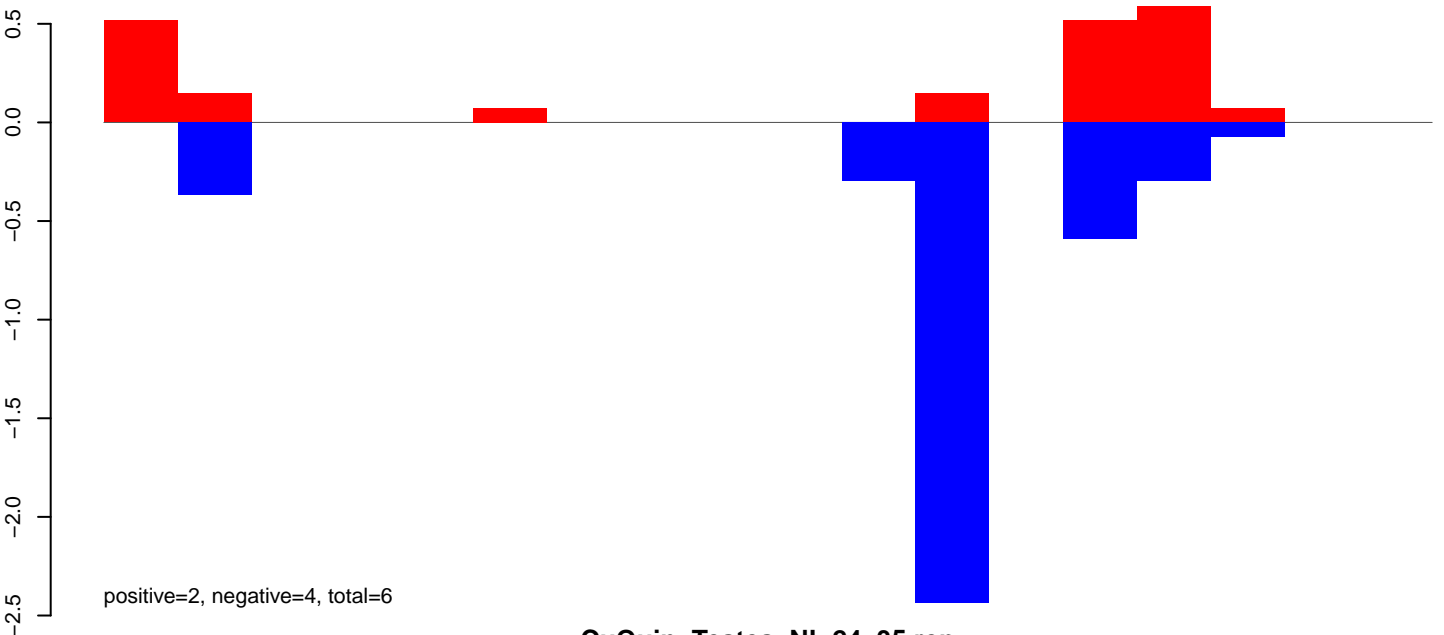
CuQuin_Testes_NL.24_35.rep



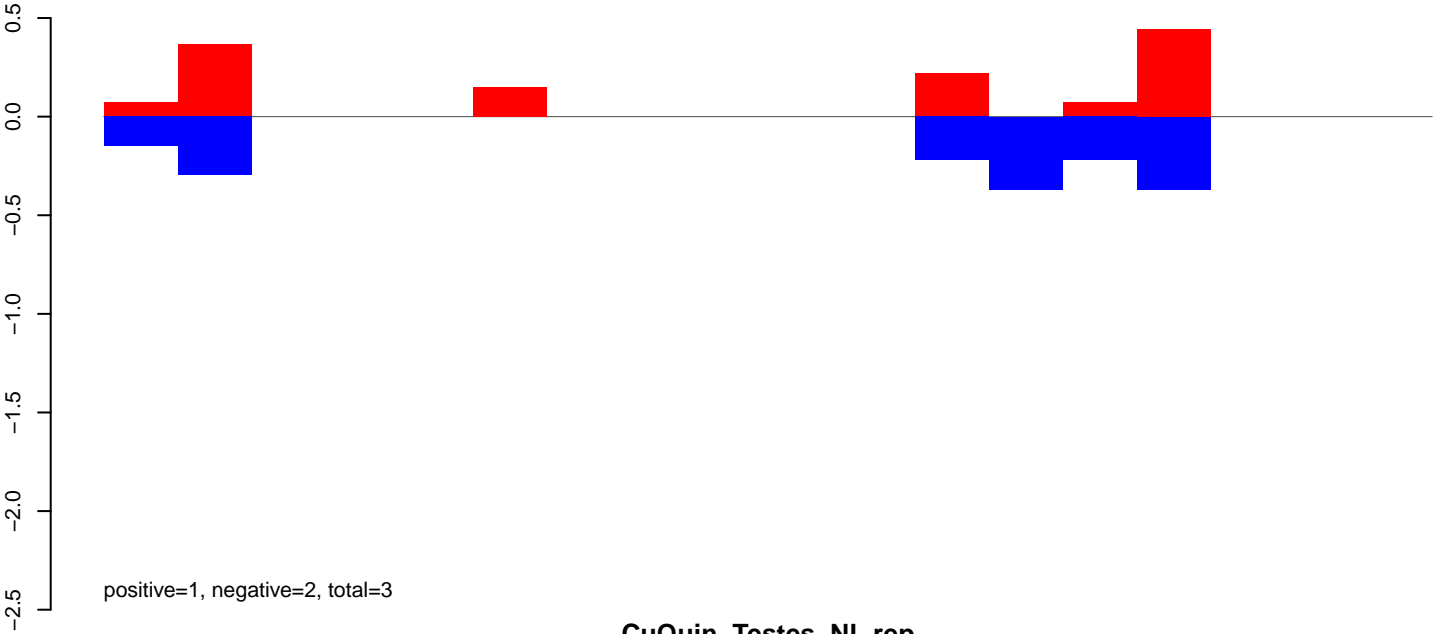
CuQuin_Testes_NL.rep



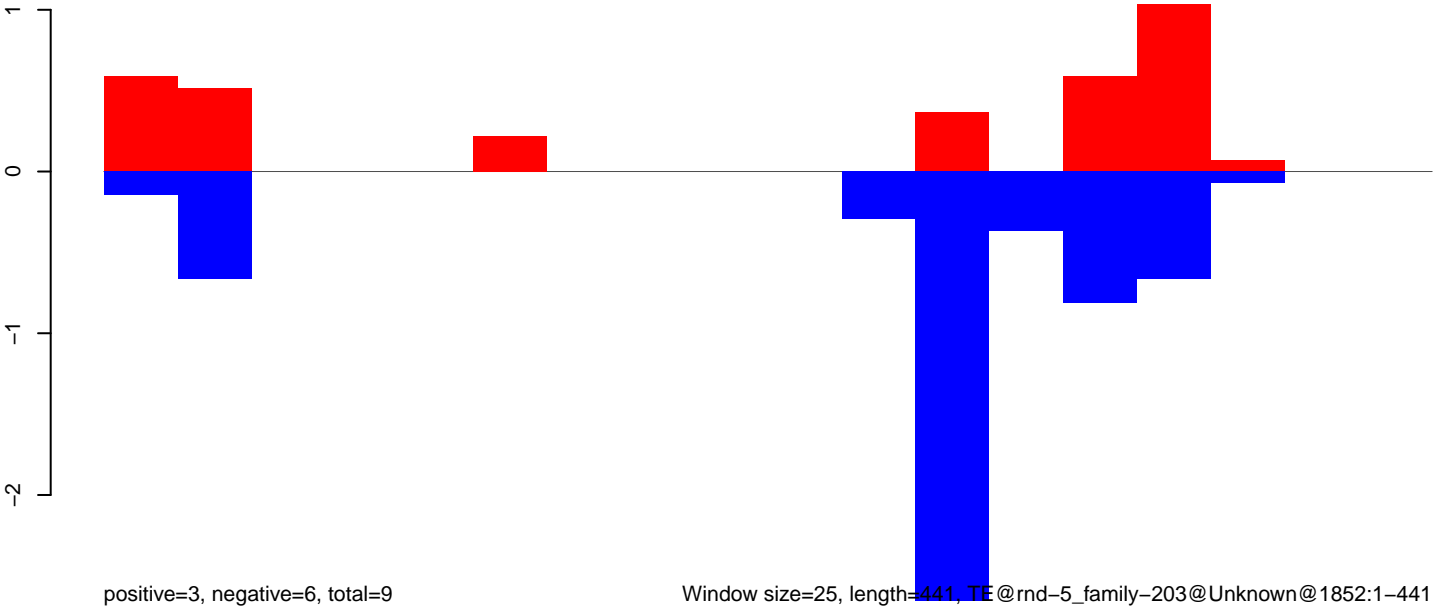
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



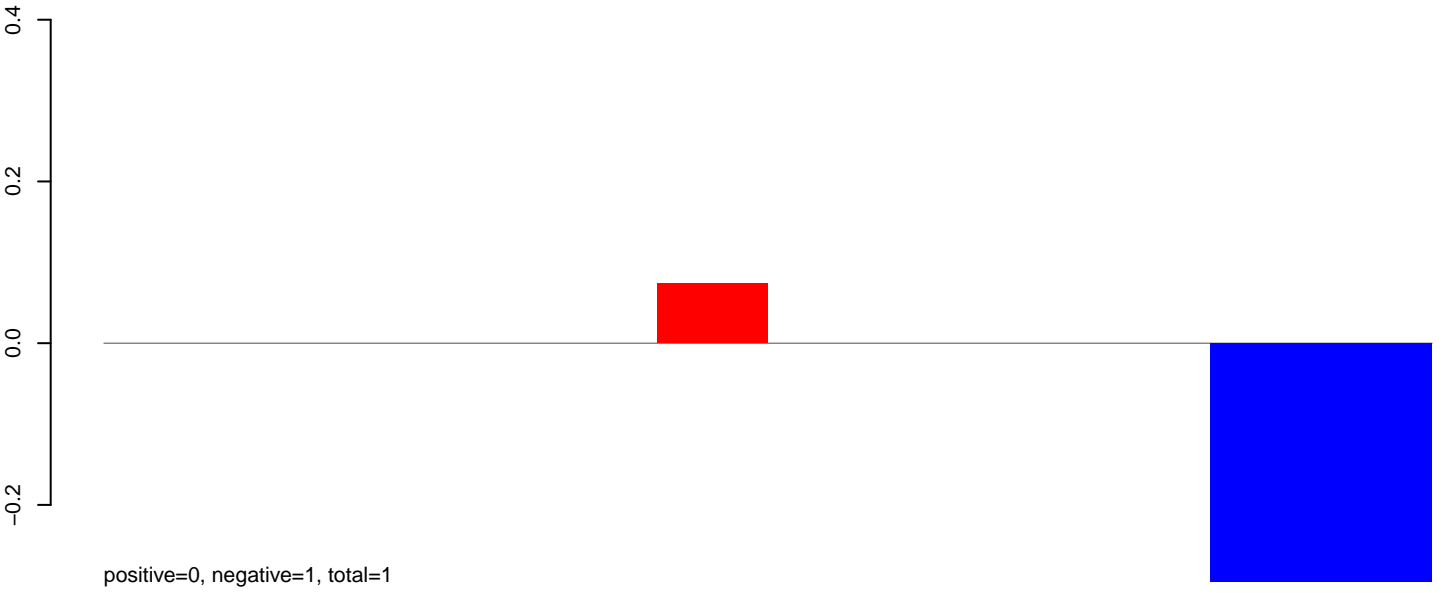
CuQuin_Testes_NL.rep



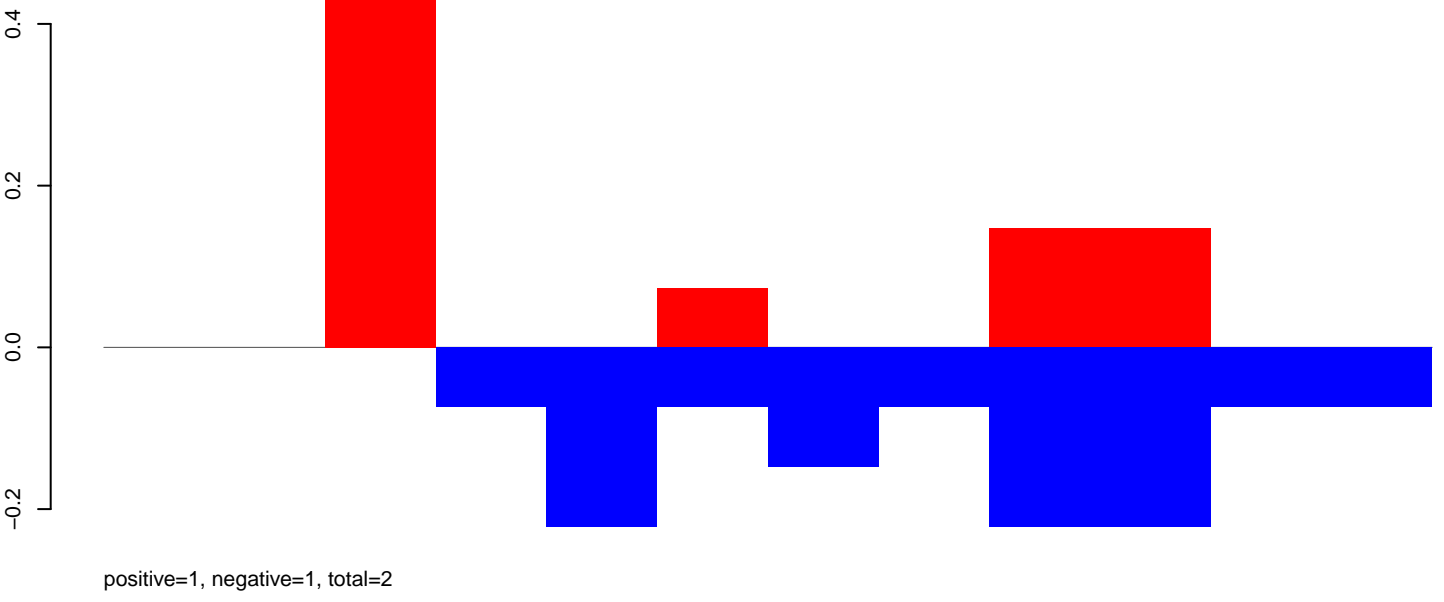
Window size=25, length=441, TE@rnd-5_family-203@Unknown@1852:1-441

100 200 300 400

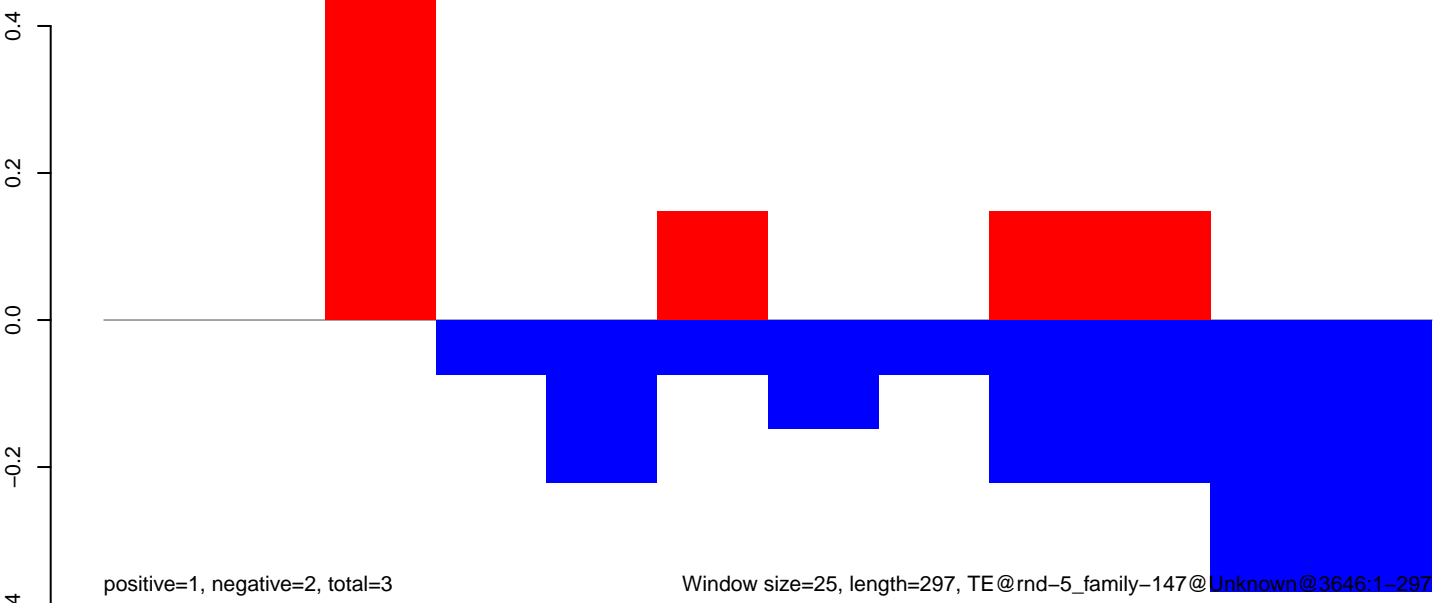
CuQuin_Testes_NL.18_23.rep



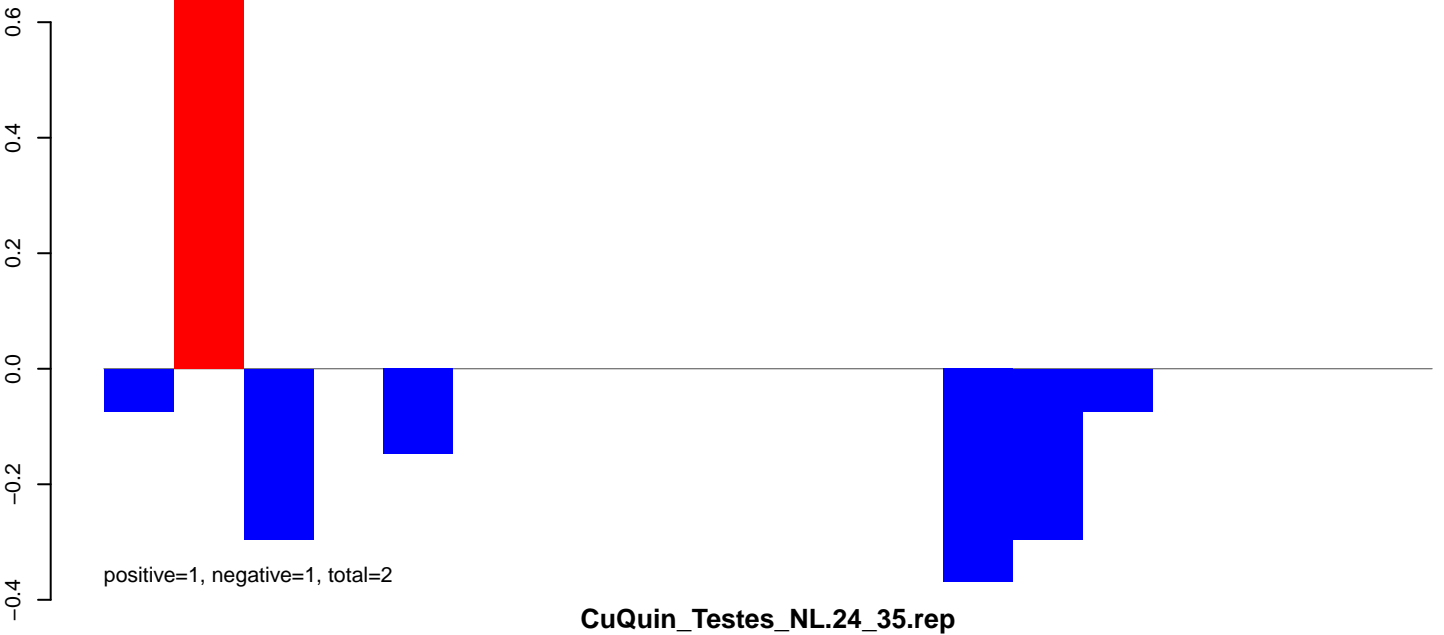
CuQuin_Testes_NL.24_35.rep



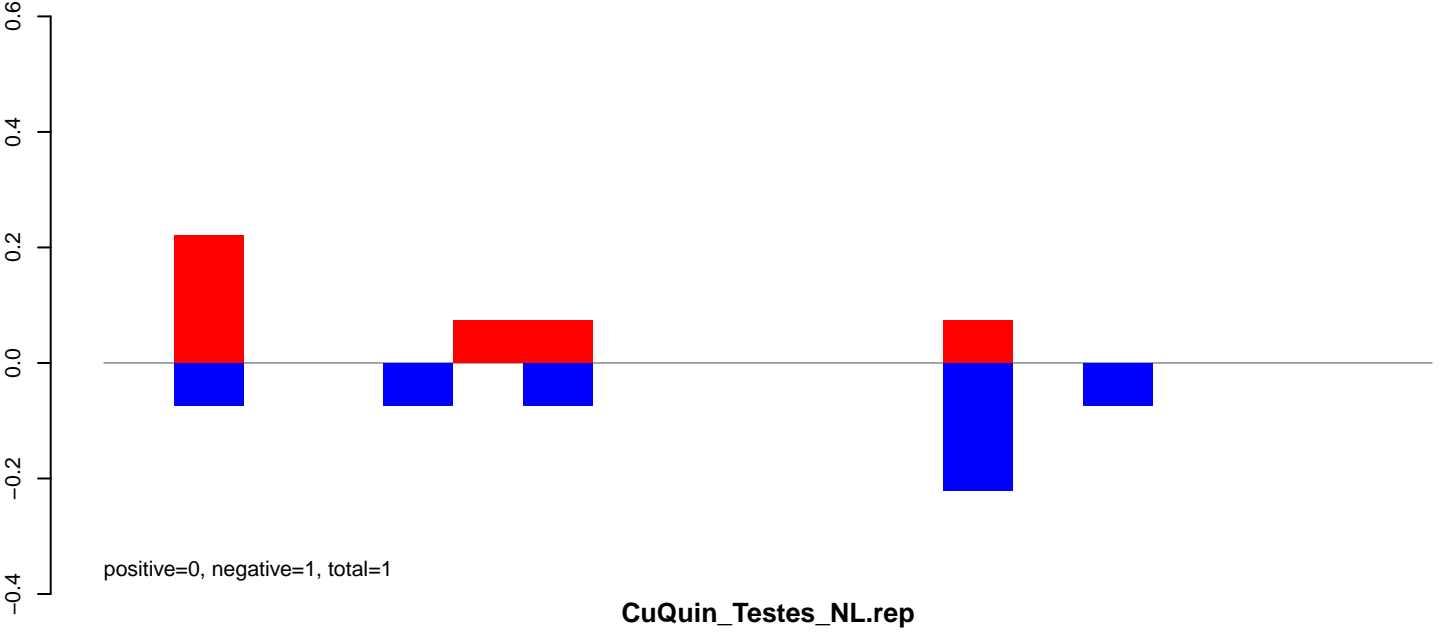
CuQuin_Testes_NL.rep



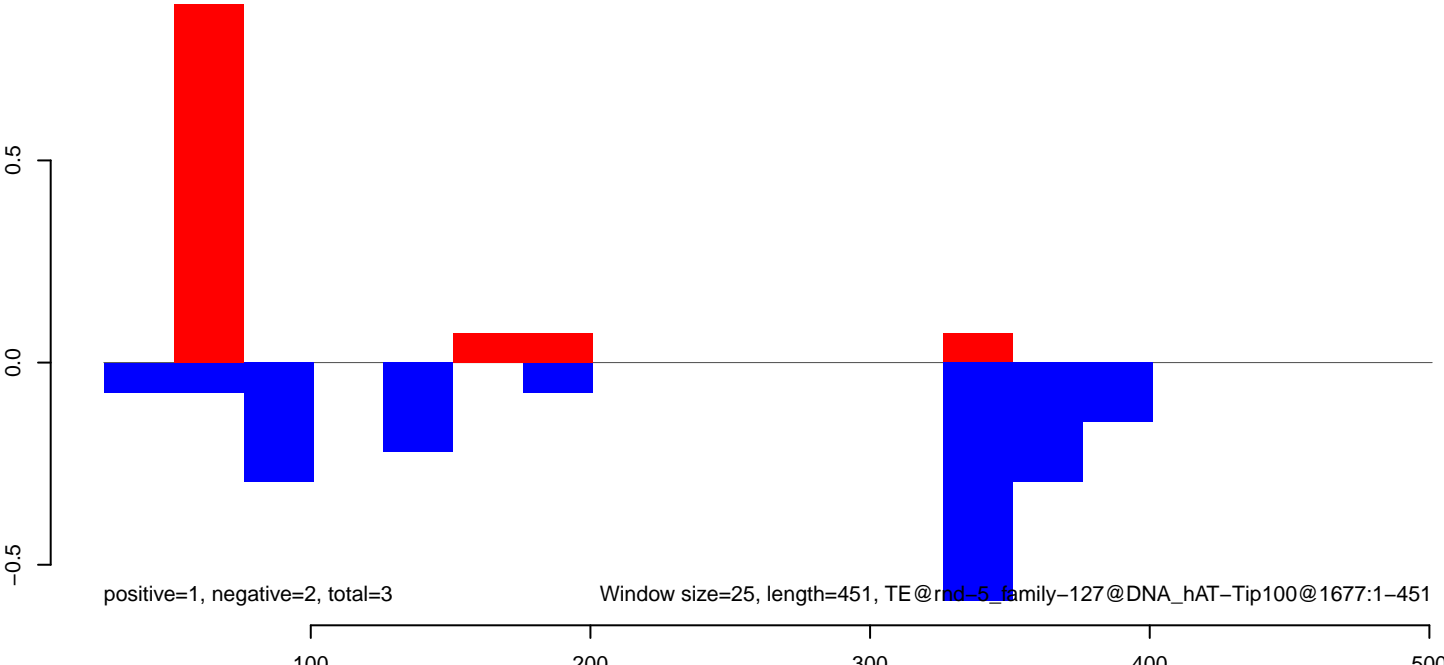
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

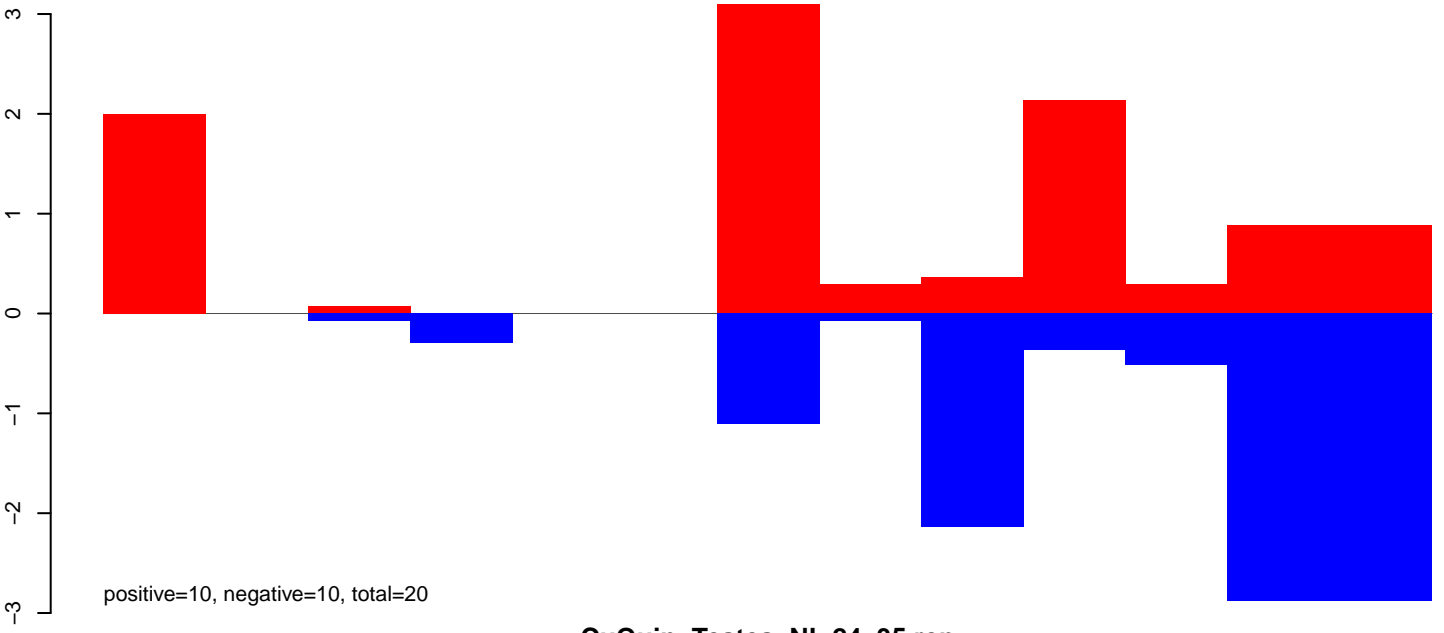


CuQuin_Testes_NL.rep

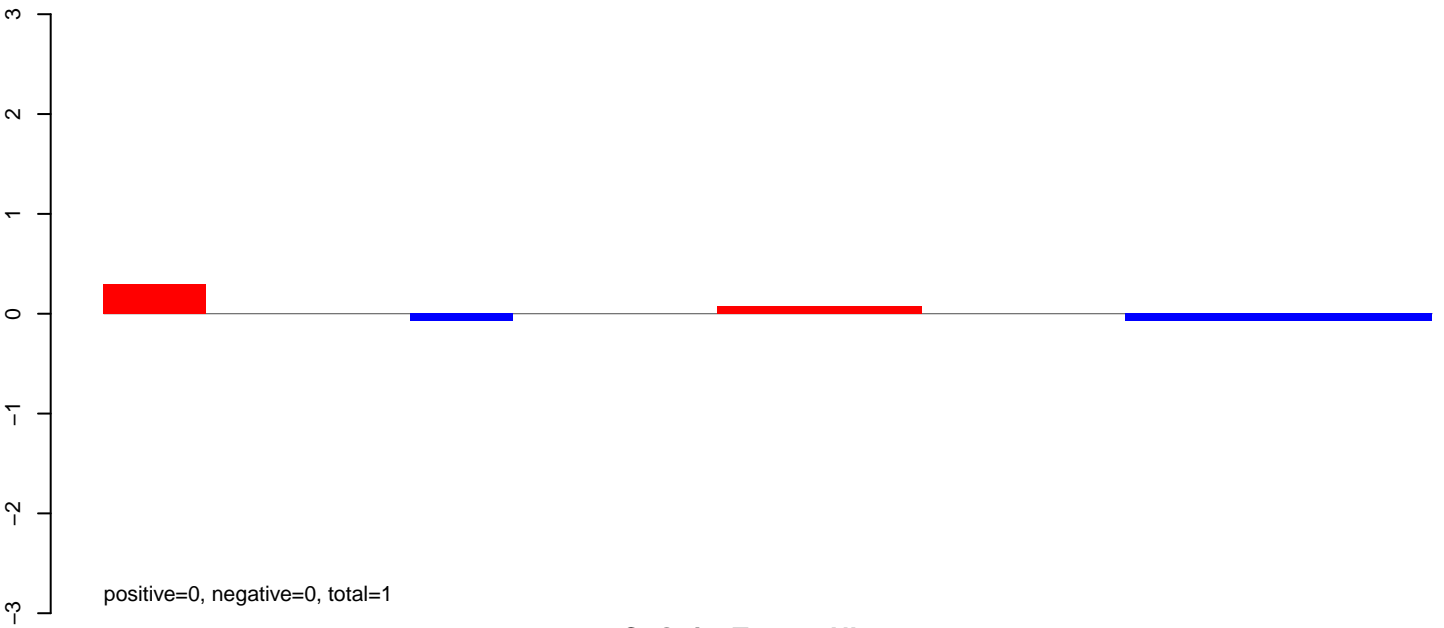


Window size=25, length=451, TE@rnd-5_family-127@DNA_hAT-Tip100@1677:1-451

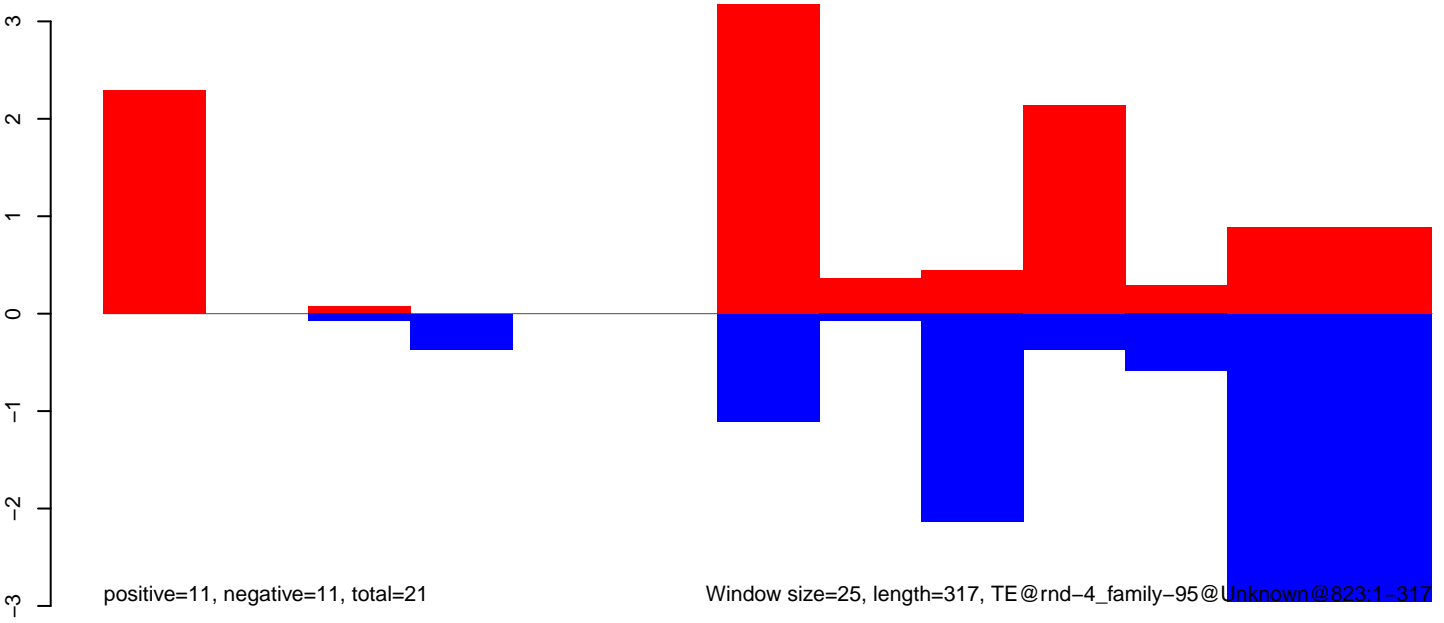
CuQuin_Testes_NL.18_23.rep



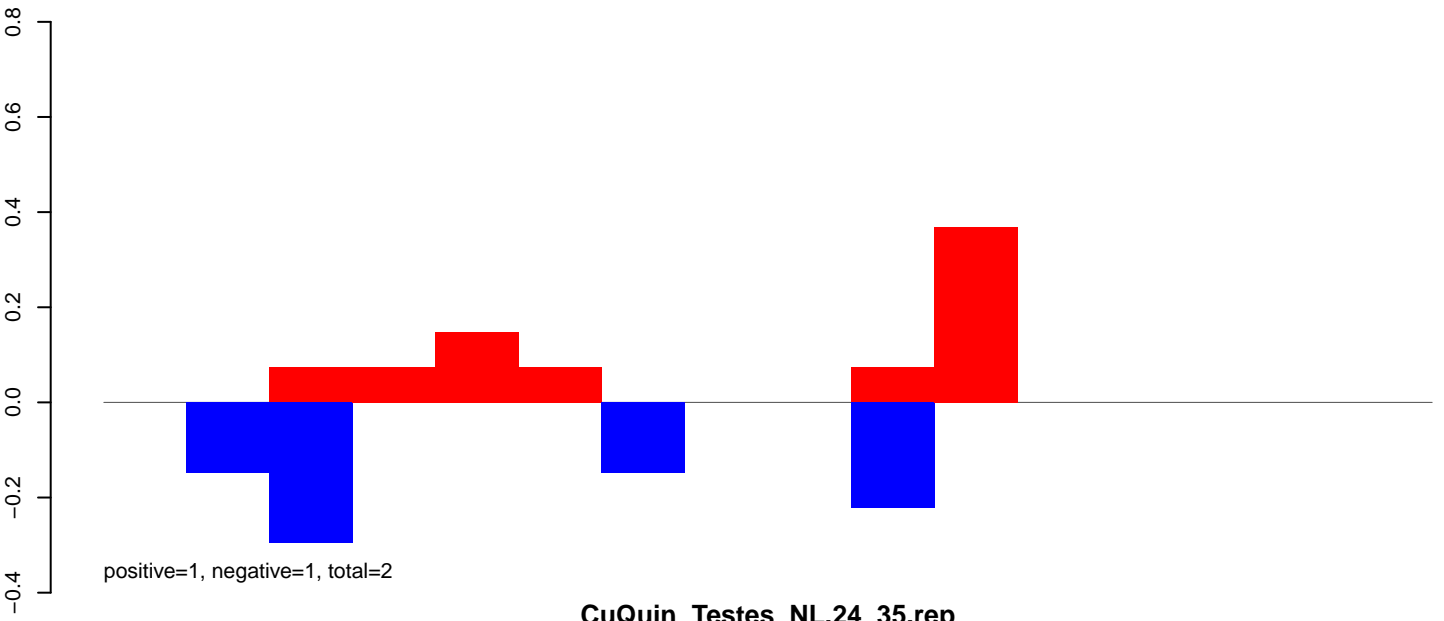
CuQuin_Testes_NL.24_35.rep



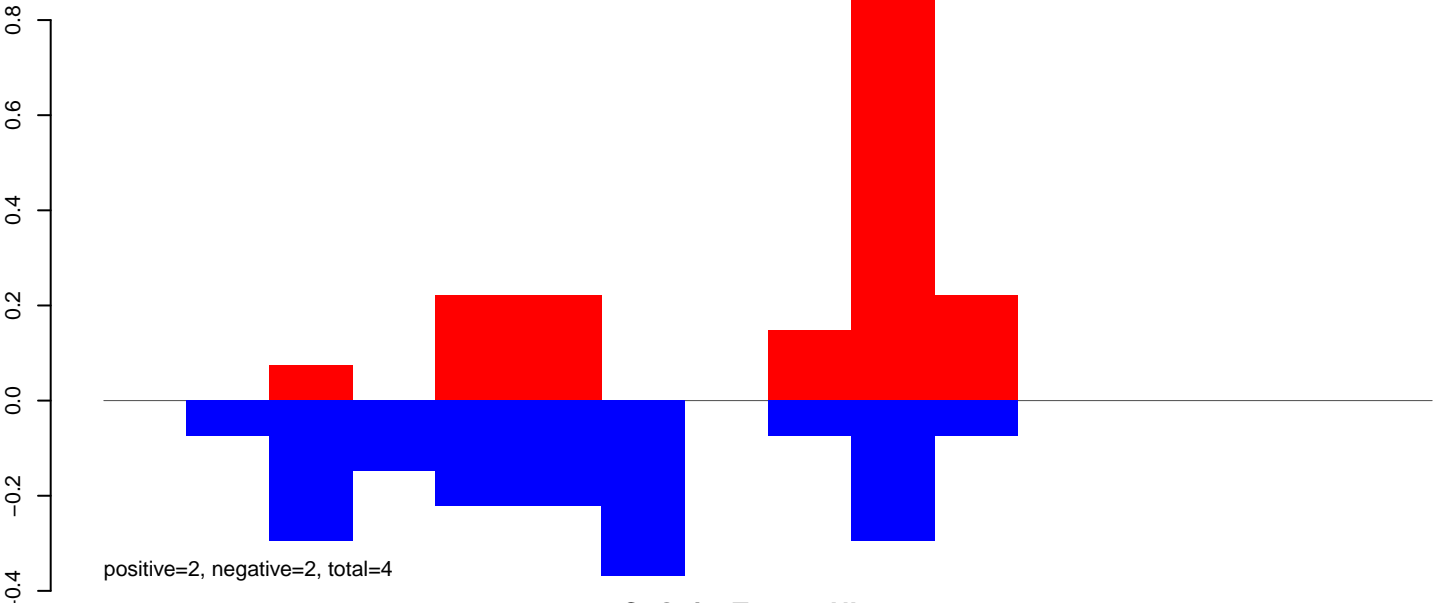
CuQuin_Testes_NL.rep



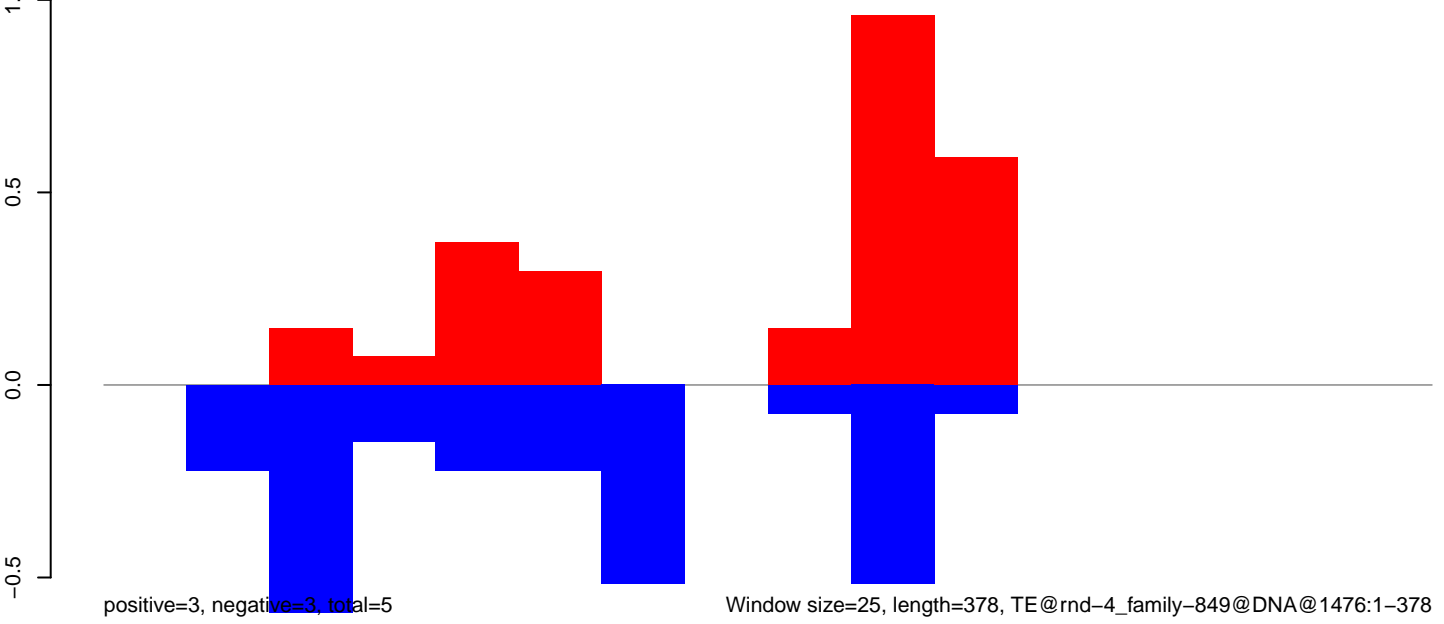
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

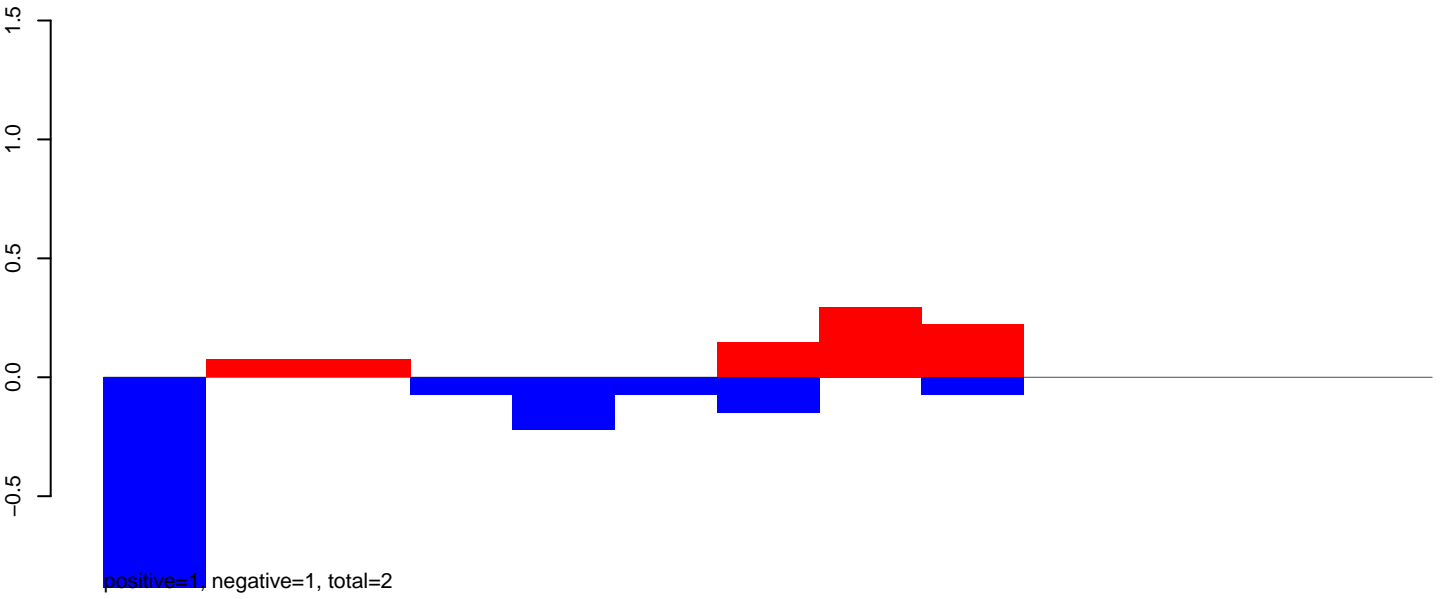


CuQuin_Testes_NL.rep

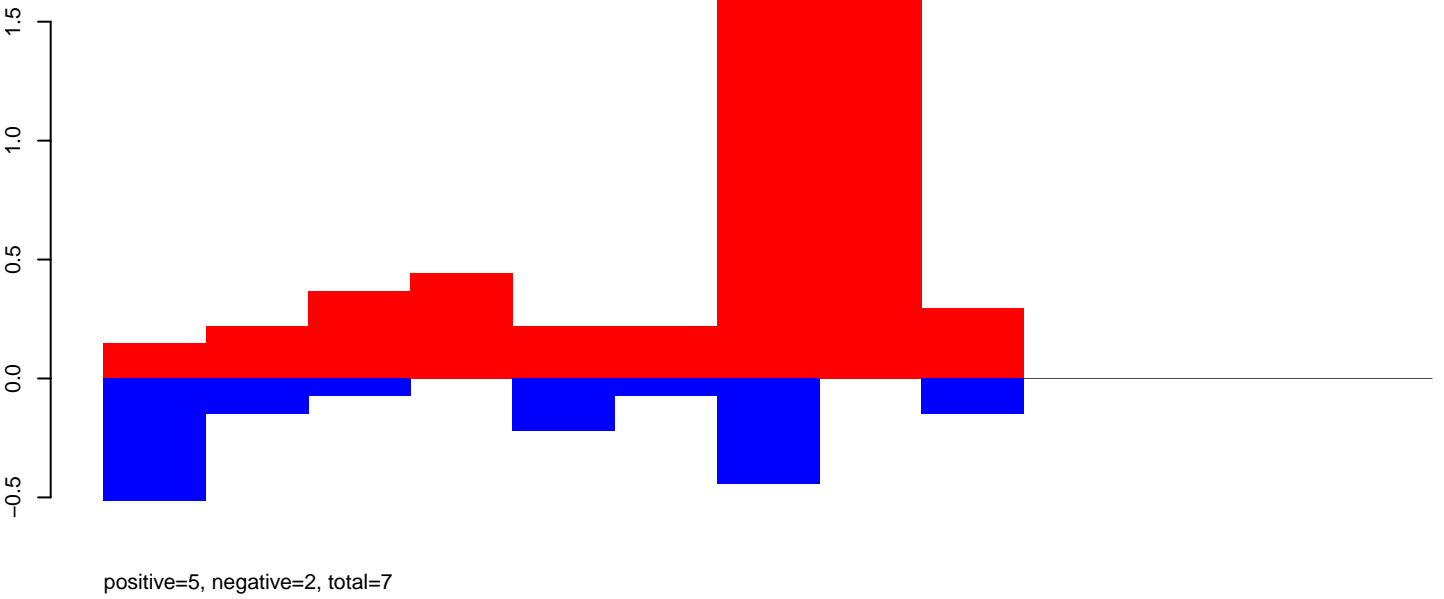


Window size=25, length=378, TE@rnd-4_family-849@DNA@1476:1-378

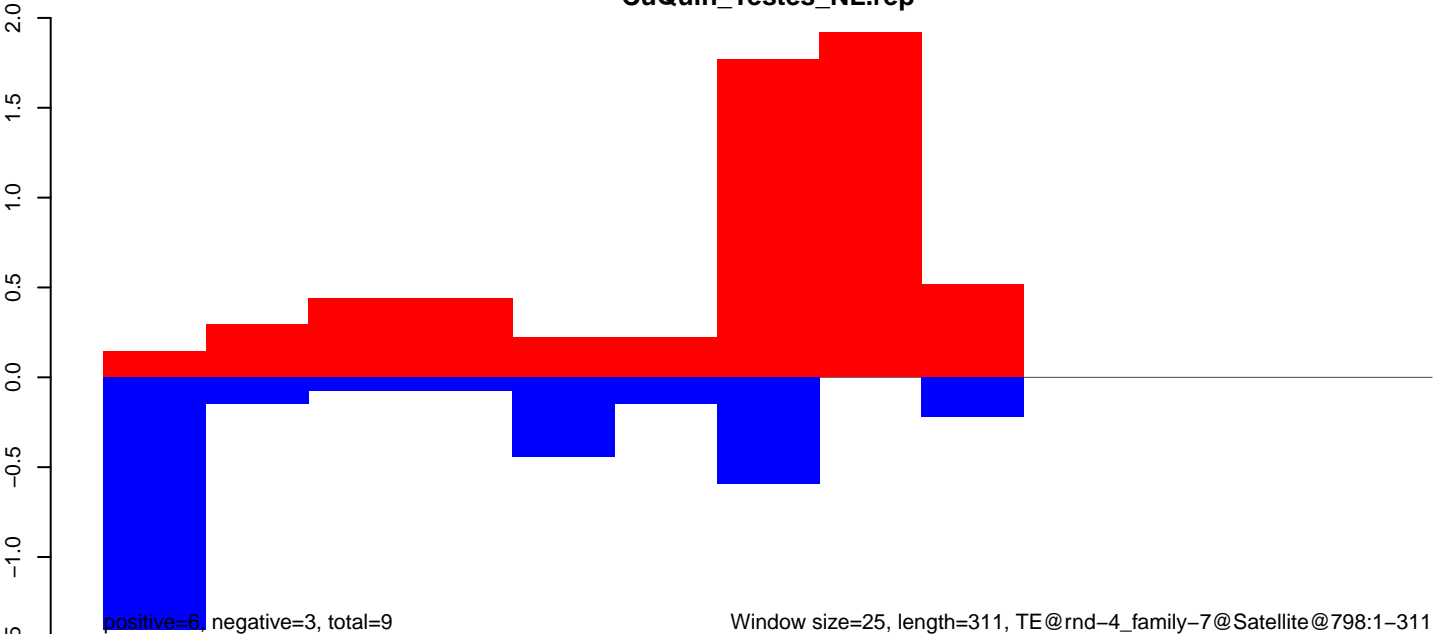
CuQuin_Testes_NL.18_23.rep



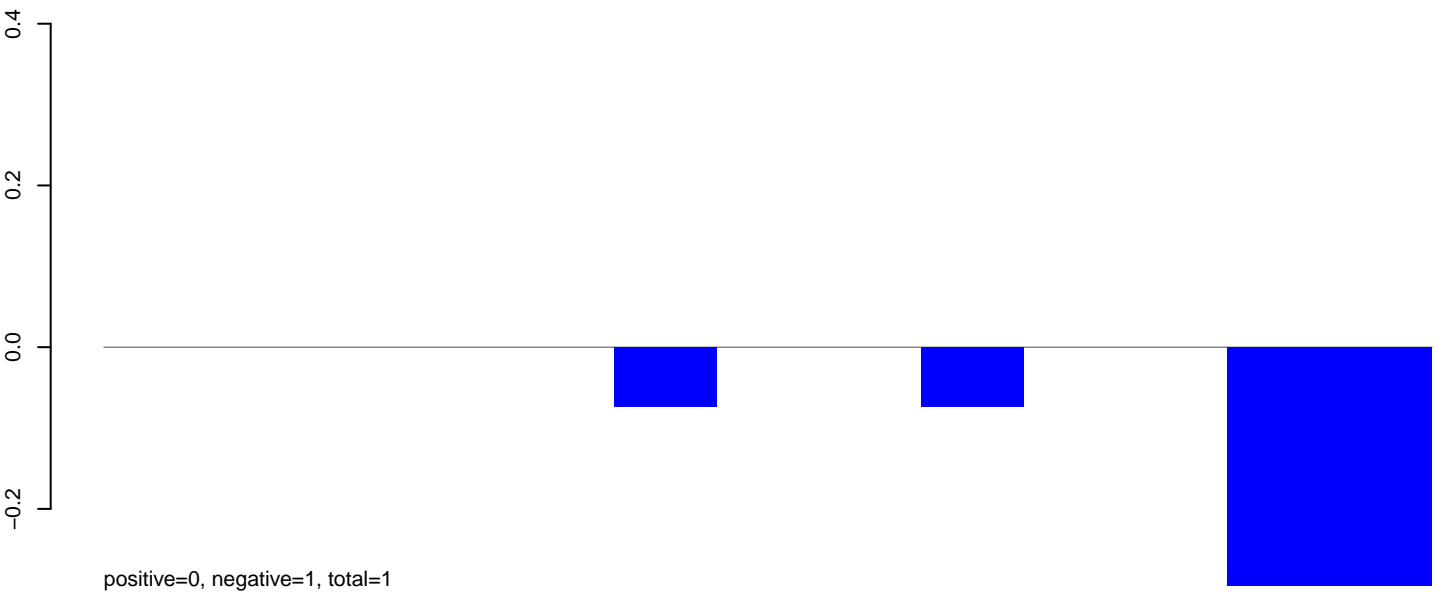
CuQuin_Testes_NL.24_35.rep



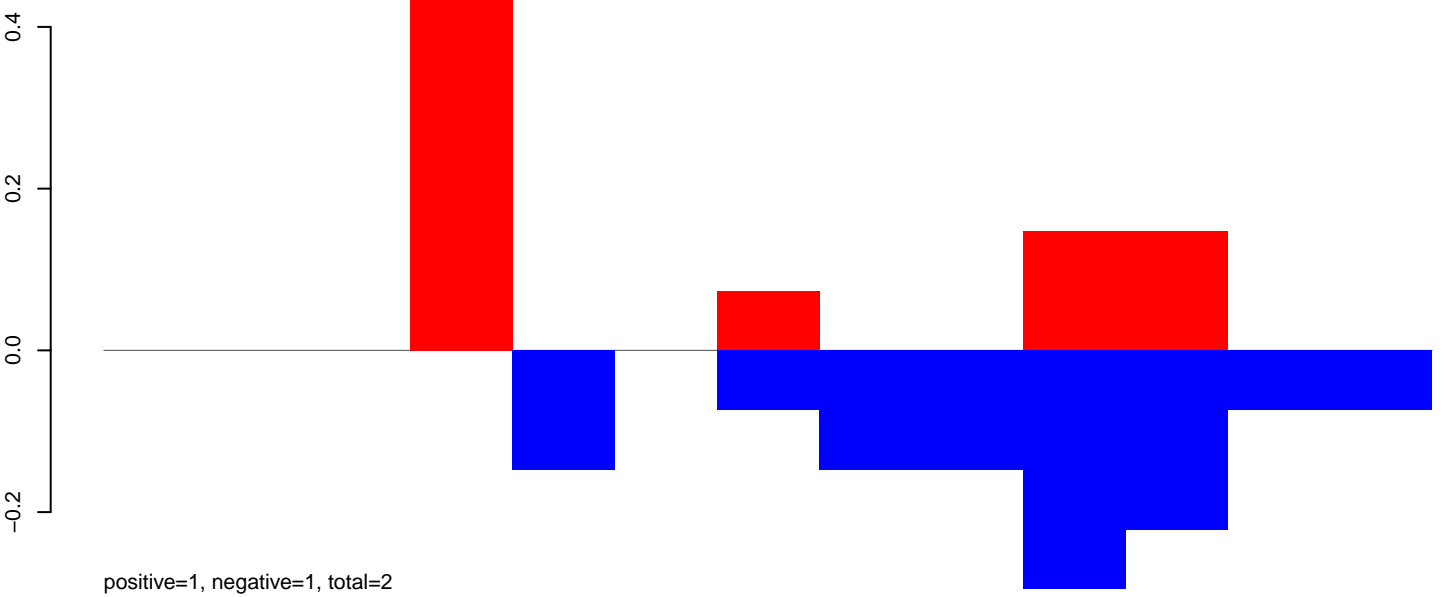
CuQuin_Testes_NL.rep



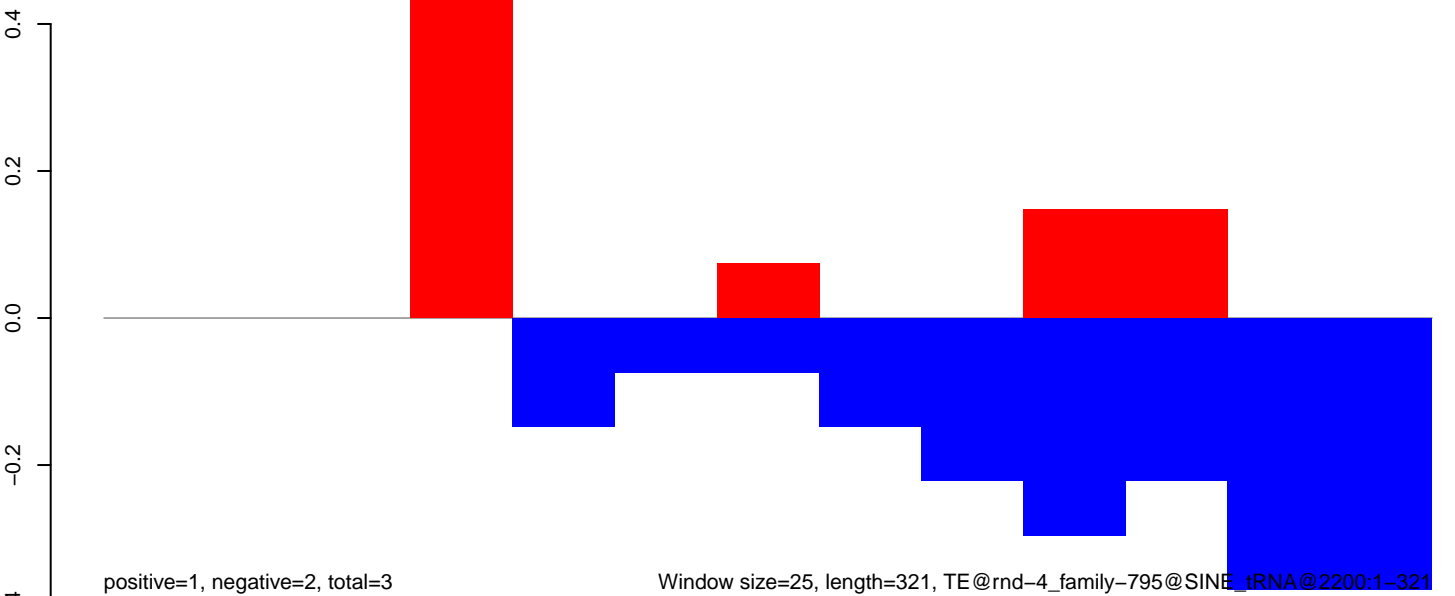
CuQuin_Testes_NL.18_23.rep



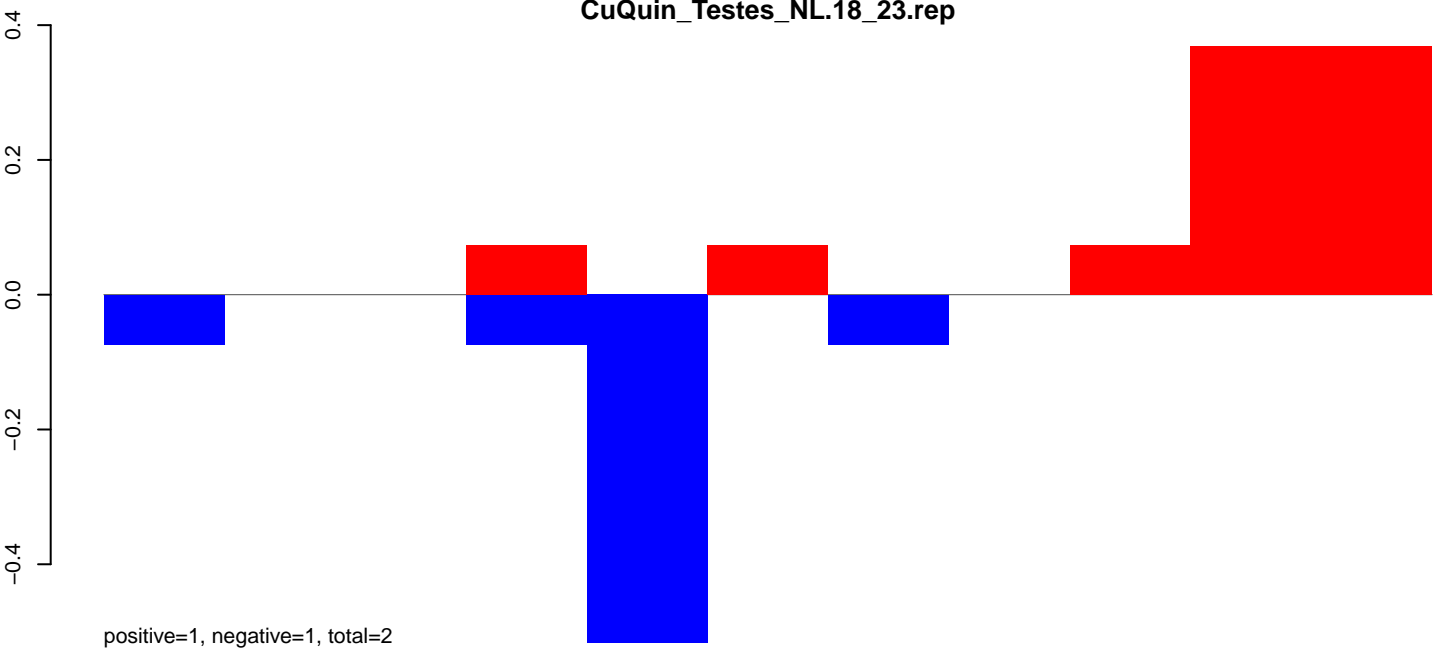
CuQuin_Testes_NL.24_35.rep



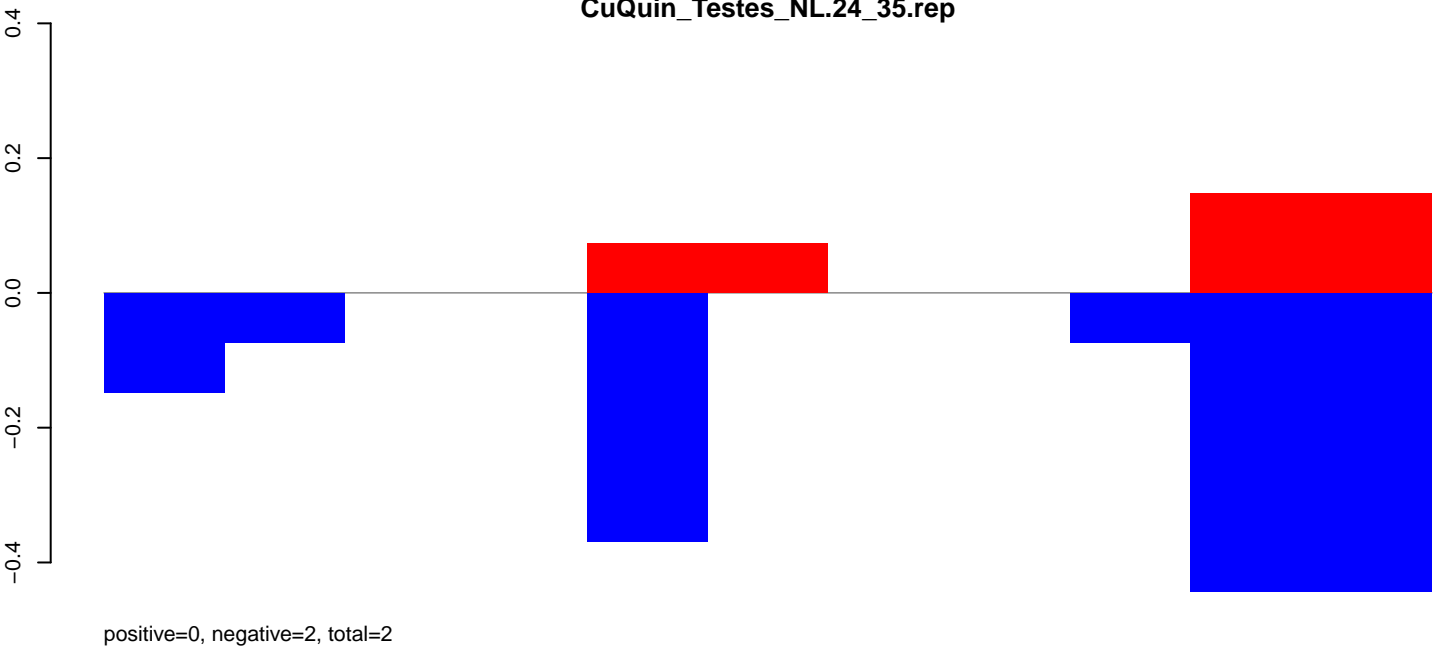
CuQuin_Testes_NL.rep



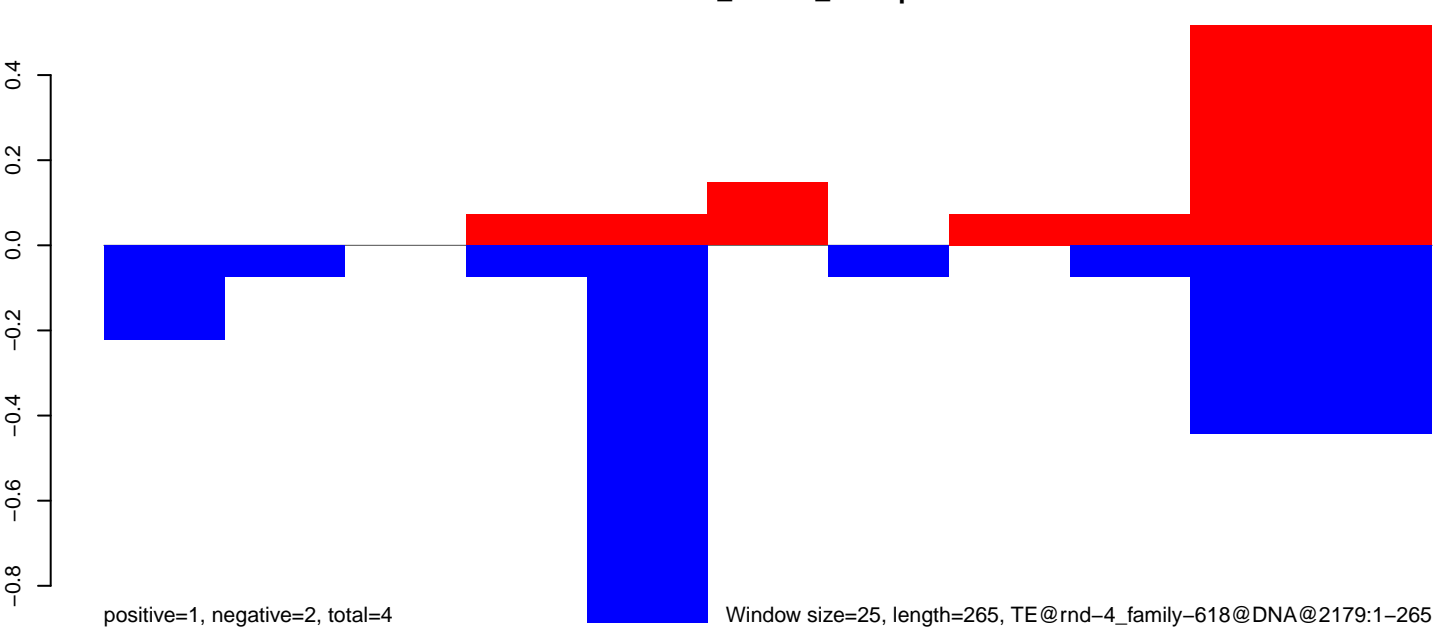
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

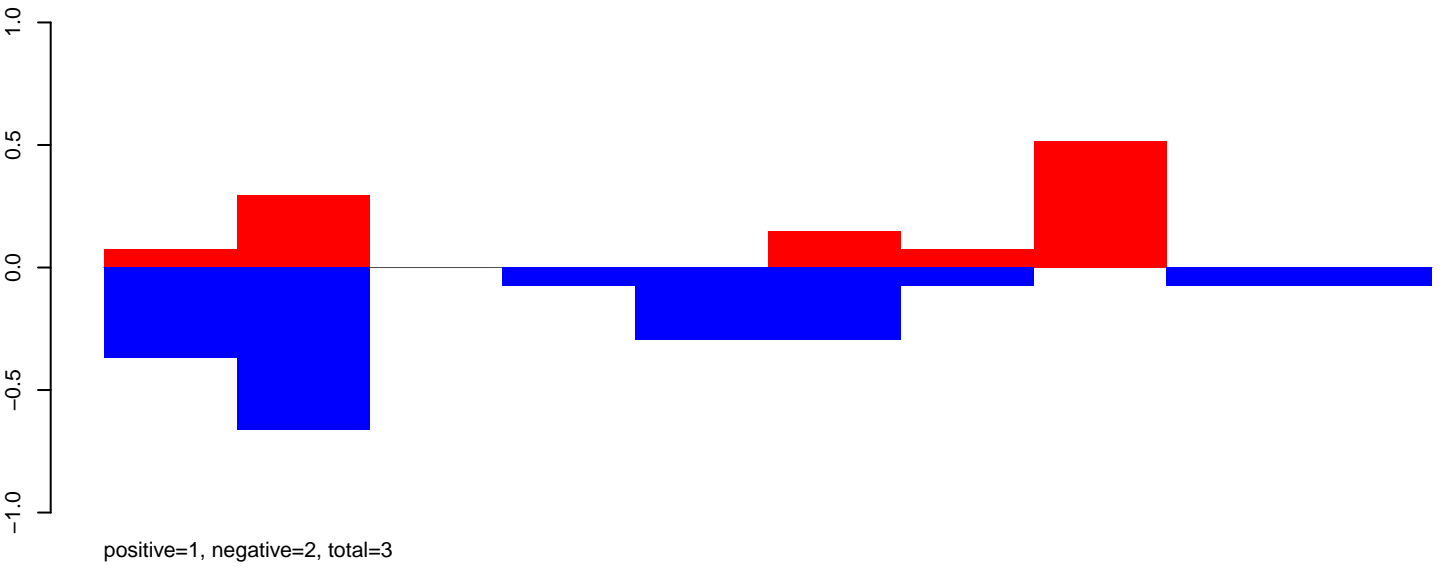


CuQuin_Testes_NL.rep

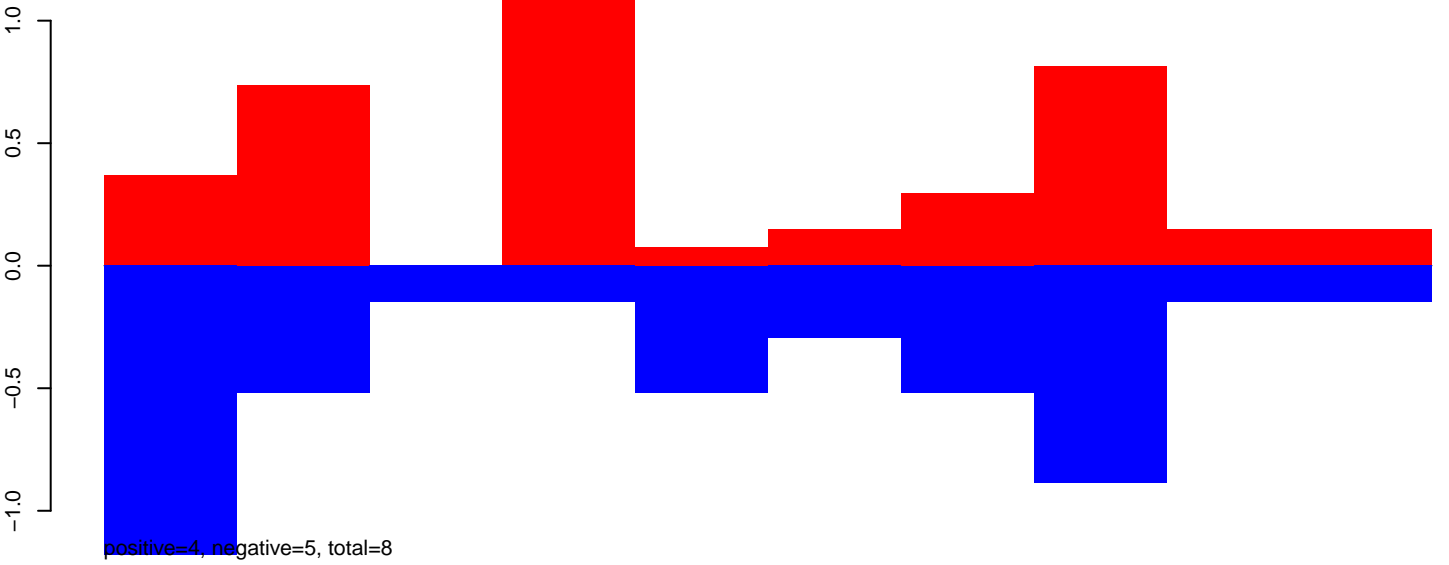


50 100 150 200 250 300

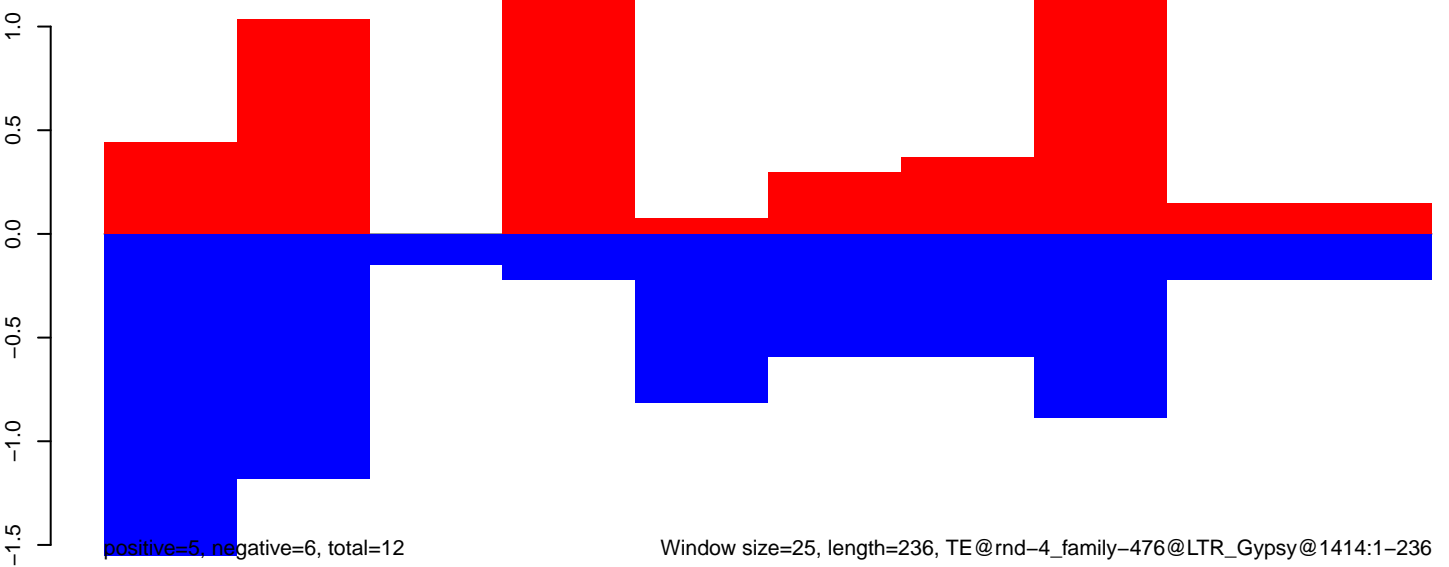
CuQuin_Testes_NL.18_23.rep



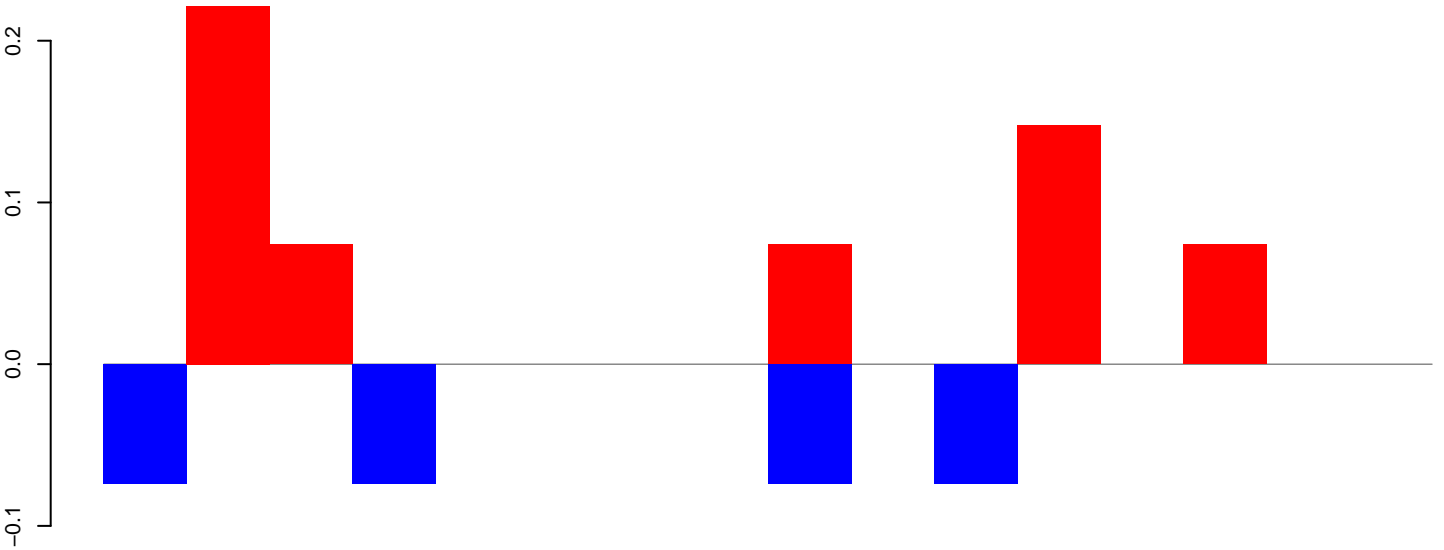
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

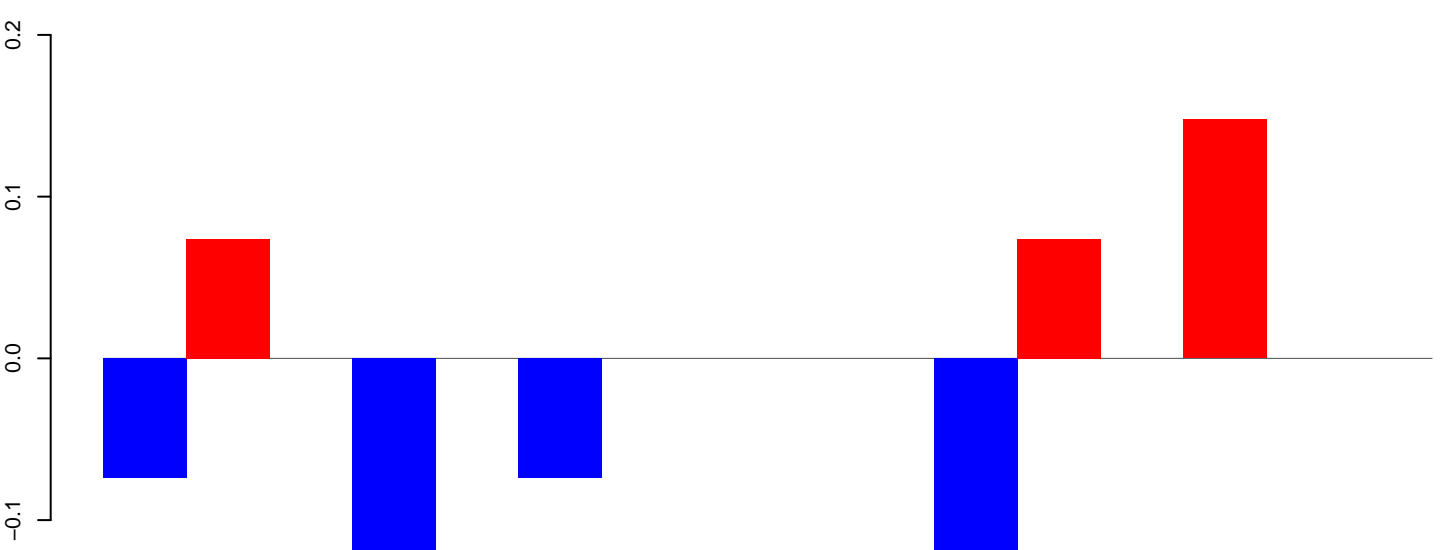


CuQuin_Testes_NL.18_23.rep



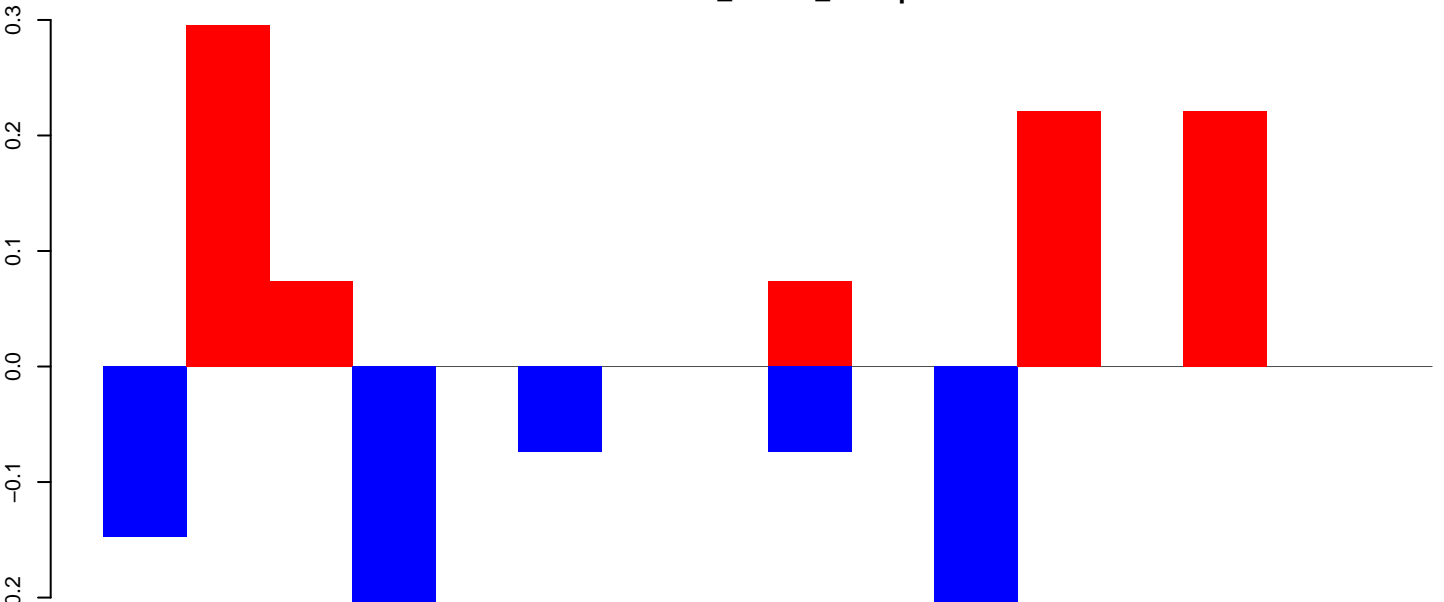
positive=1, negative=0, total=1

CuQuin_Testes_NL.24_35.rep



positive=0, negative=0, total=1

CuQuin_Testes_NL.rep



positive=1, negative=1, total=2

Window size=25, length=398, TE@ind-4_family-29@DNA_Zator@3525:1-398

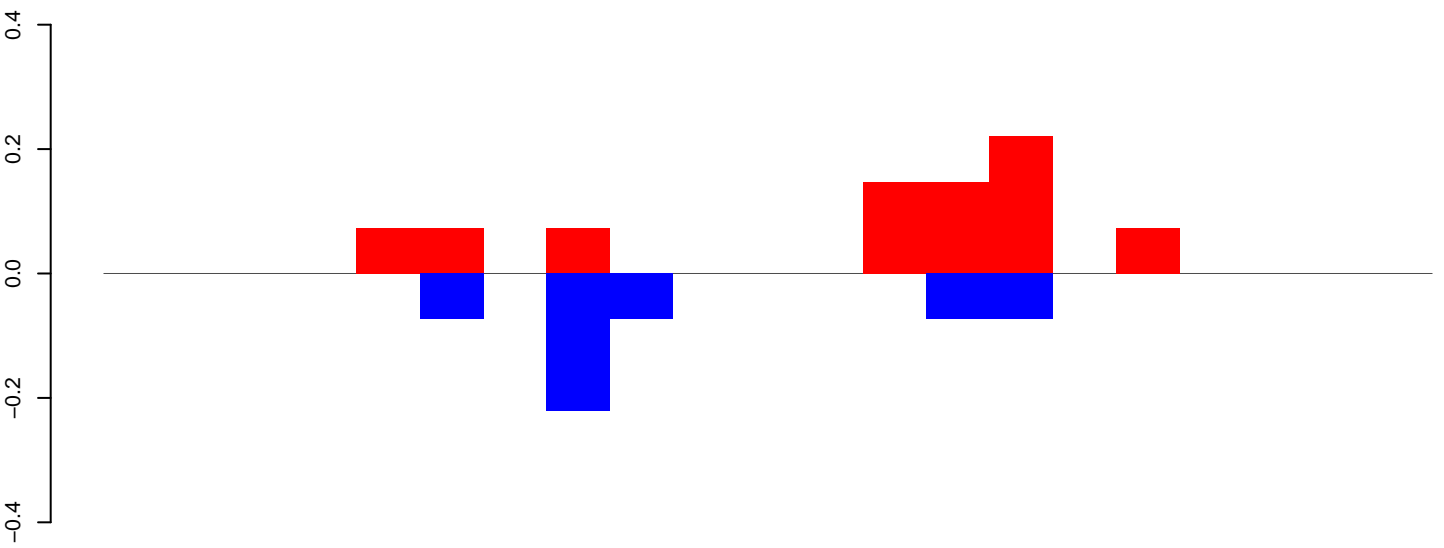
100

200

300

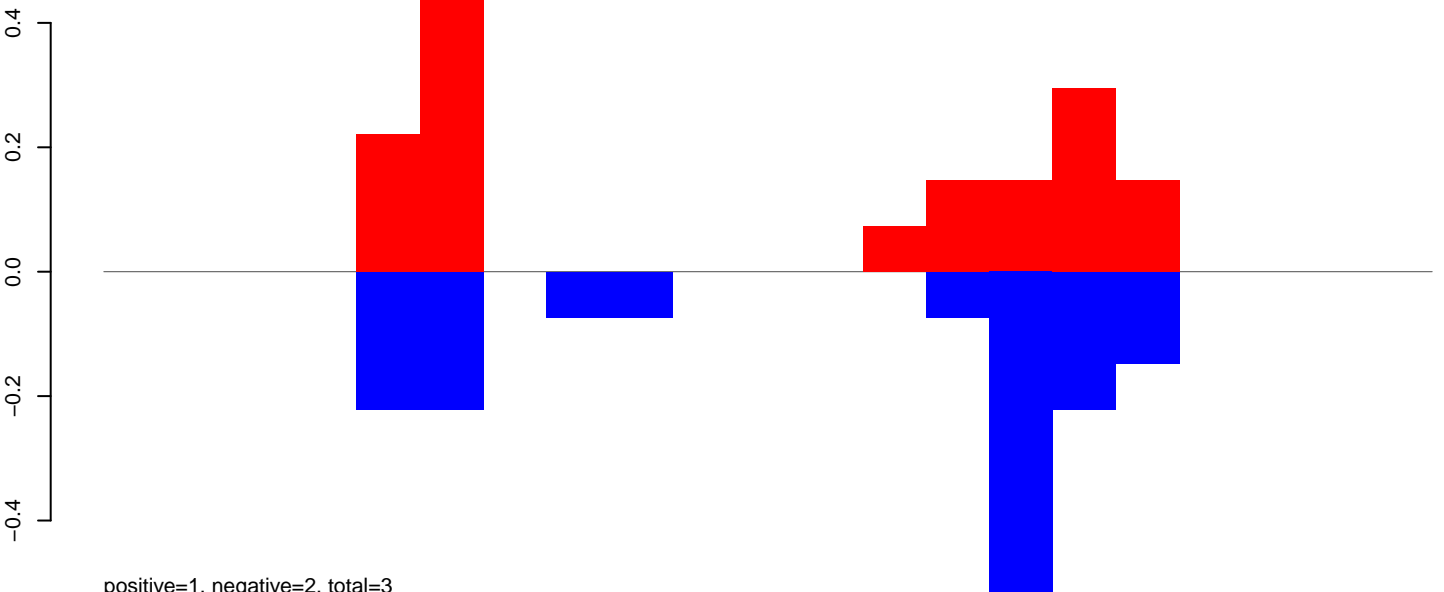
400

CuQuin_Testes_NL.18_23.rep



positive=1, negative=1, total=1

CuQuin_Testes_NL.24_35.rep



positive=1, negative=2, total=3

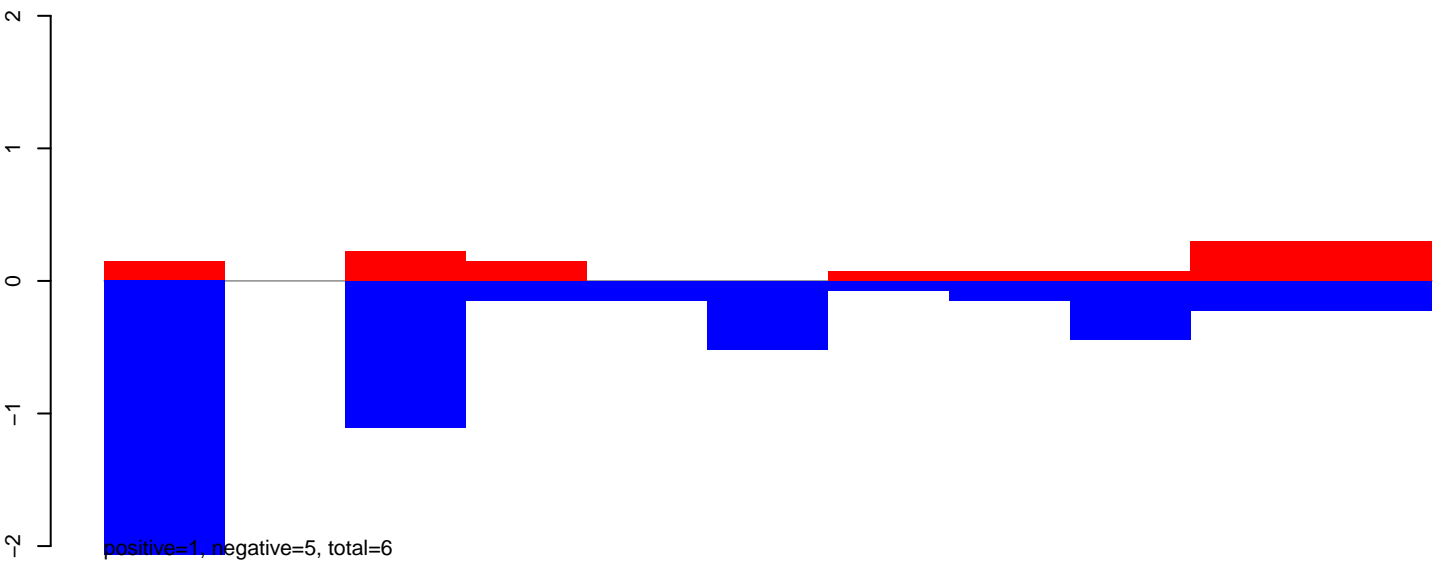
CuQuin_Testes_NL.rep



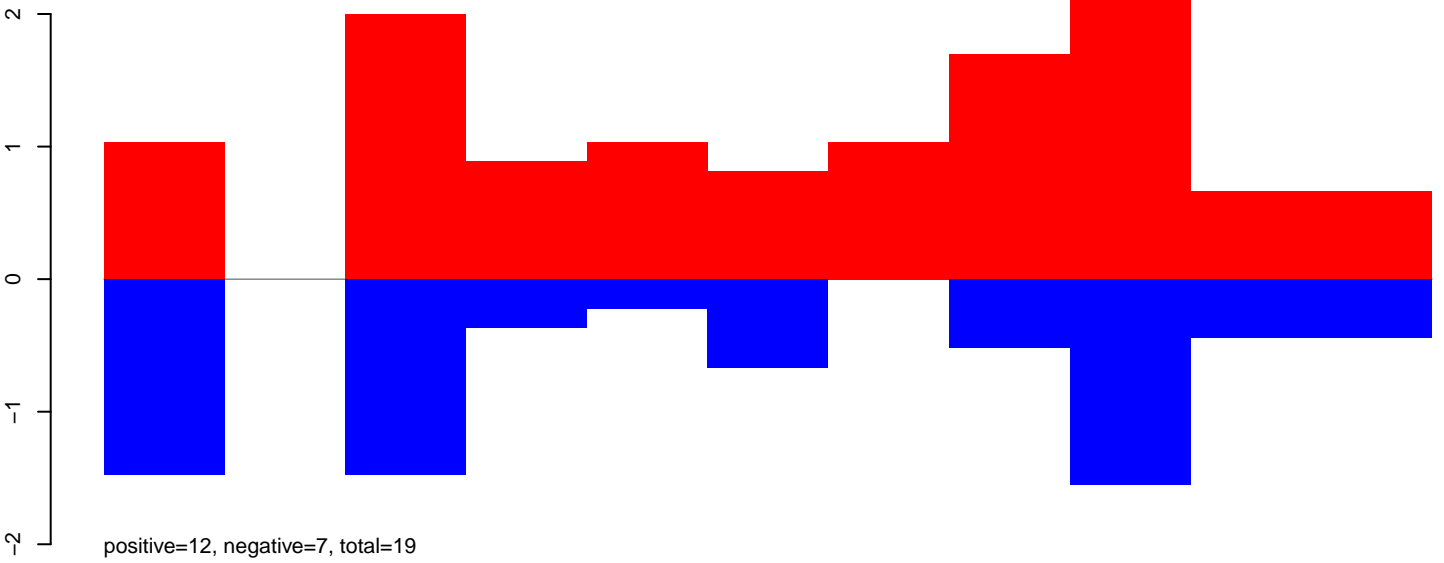
positive=2, negative=2, total=4

Window size=25, length=519, TE@rnd-4_family-298@Unknown@168:1-519

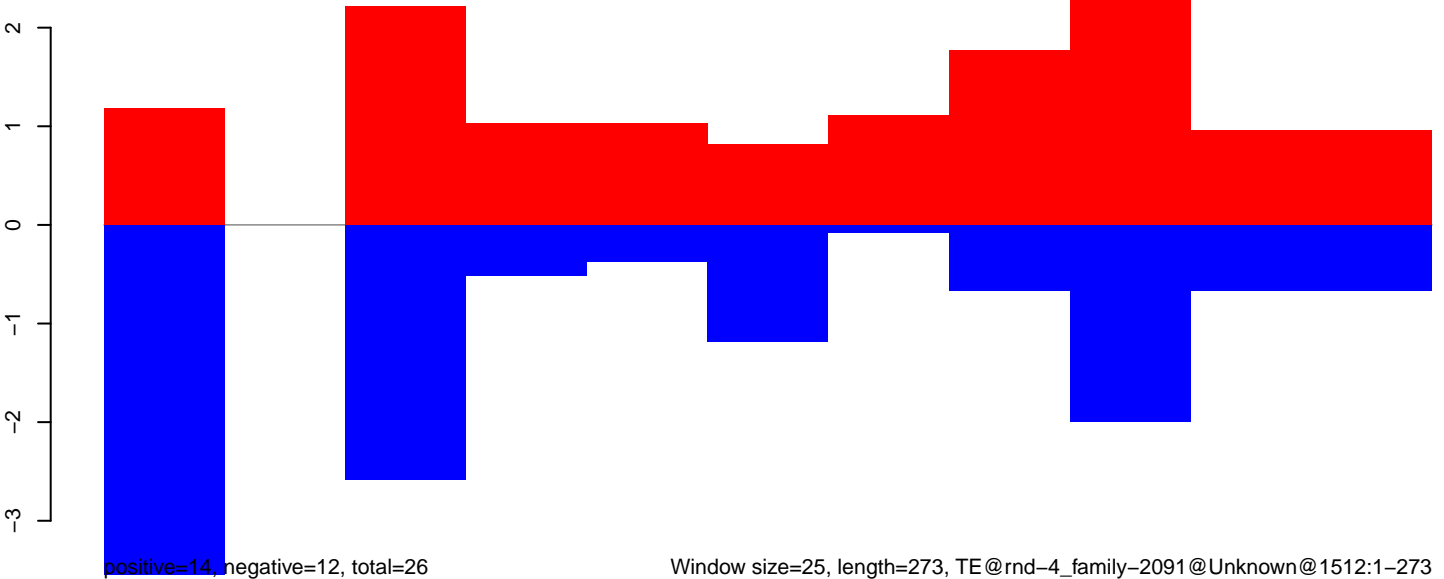
CuQuin_Testes_NL.18_23.rep



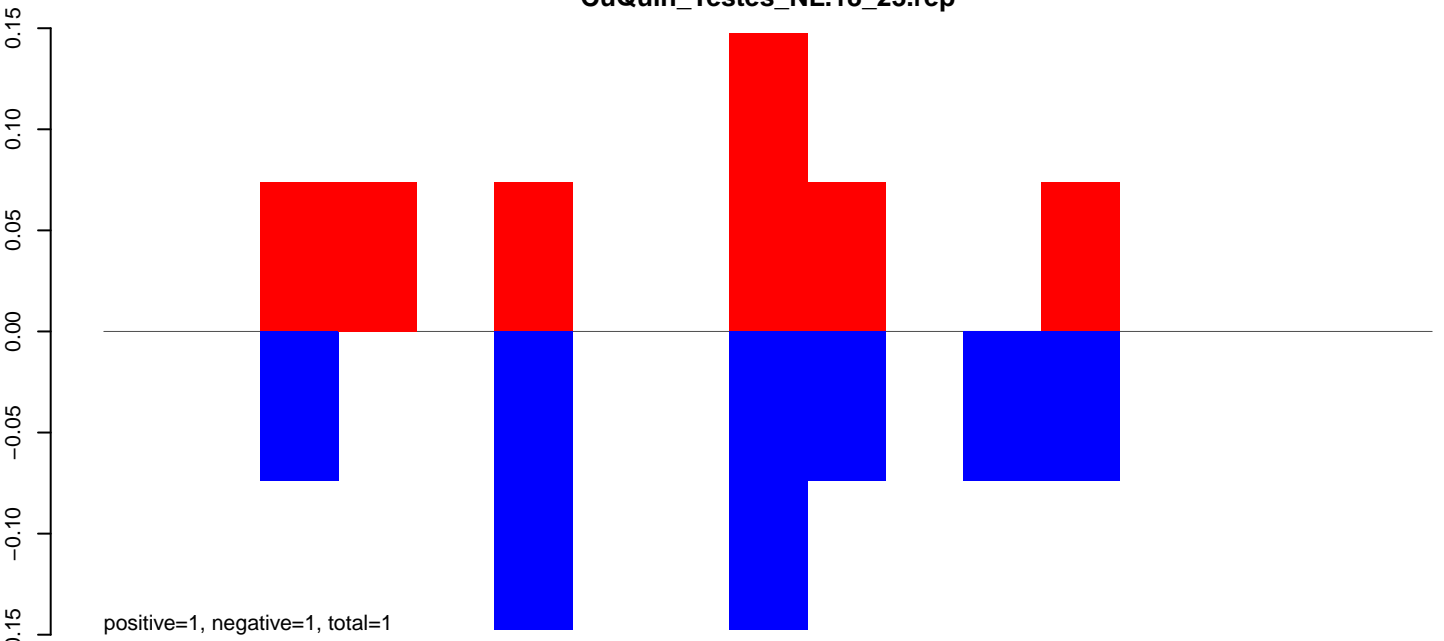
CuQuin_Testes_NL.24_35.rep



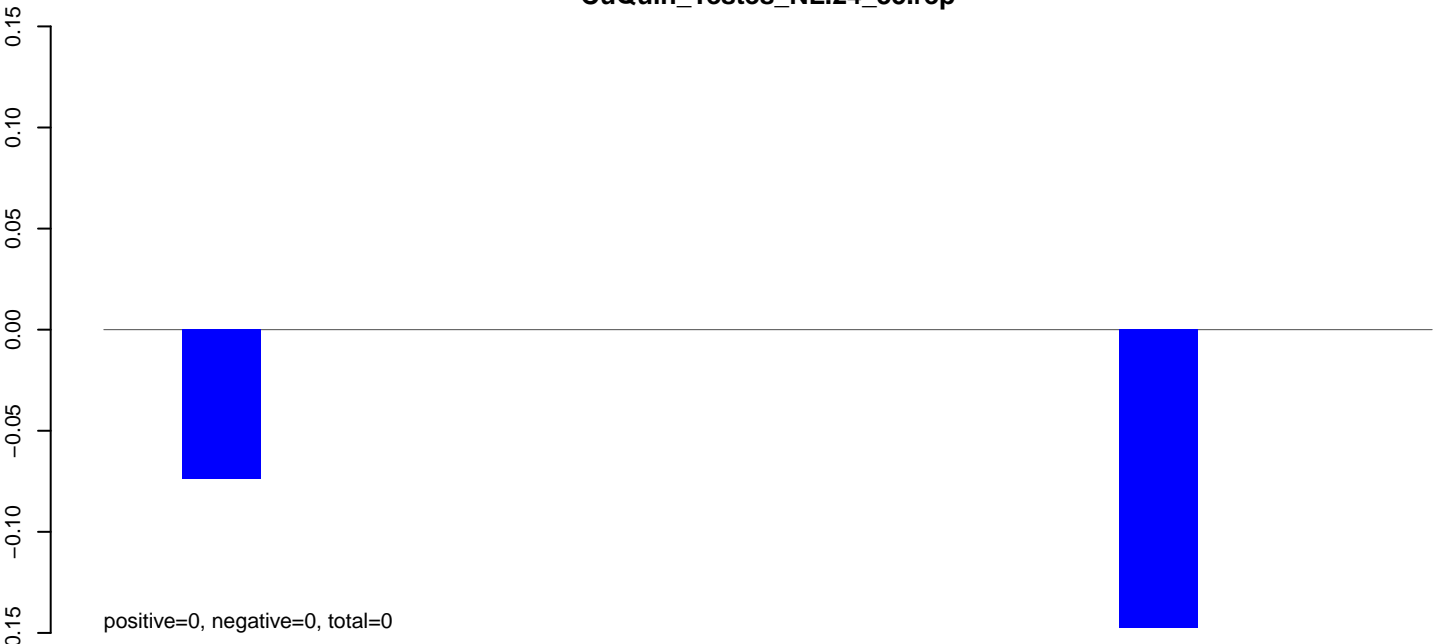
CuQuin_Testes_NL.rep



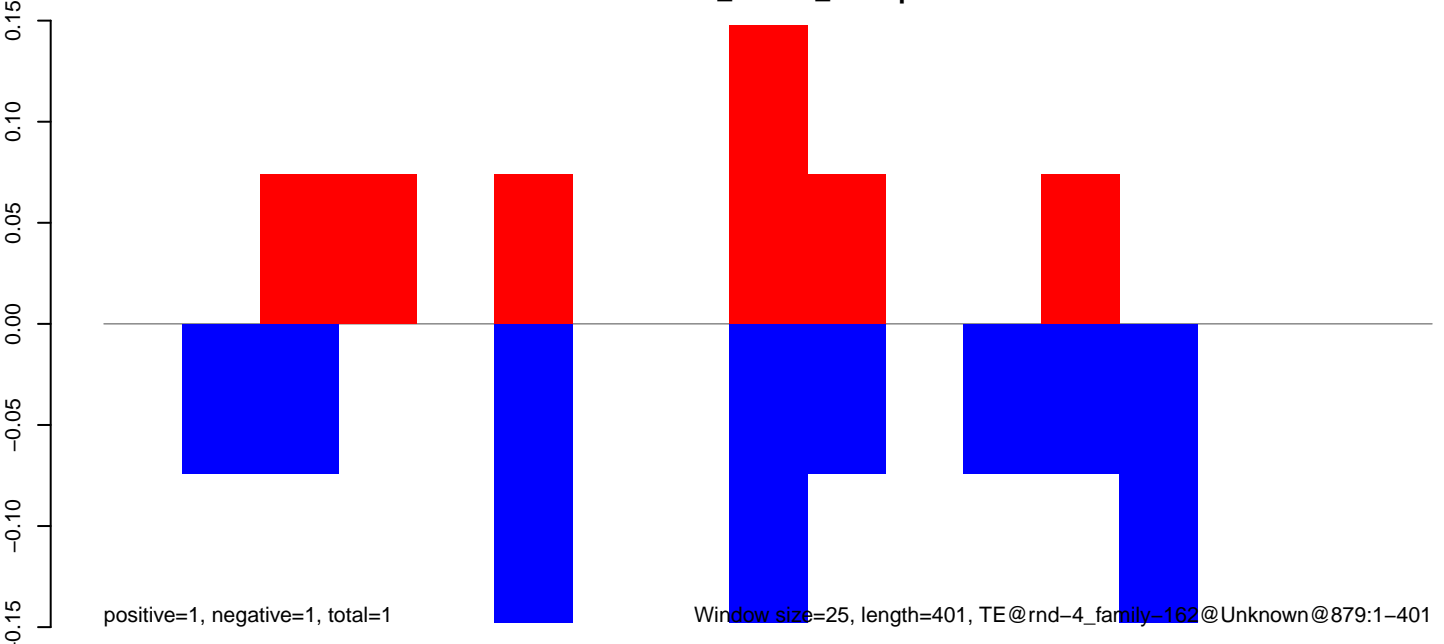
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



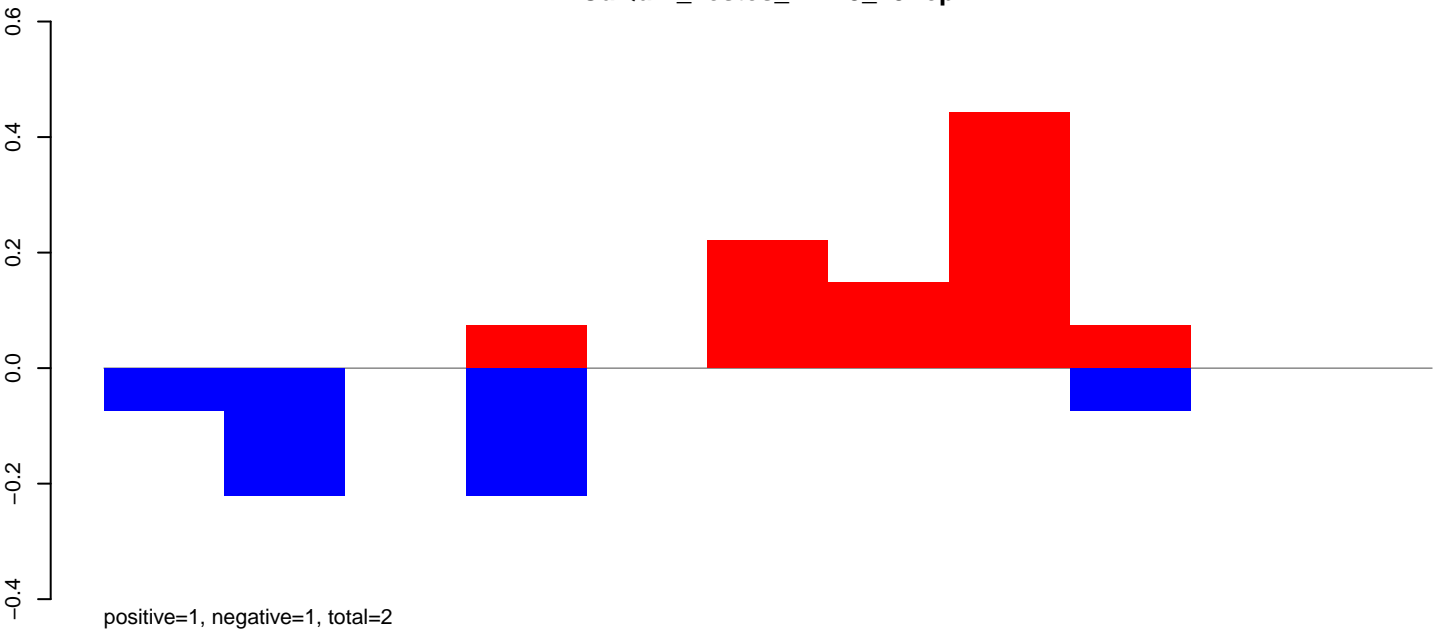
CuQuin_Testes_NL.rep



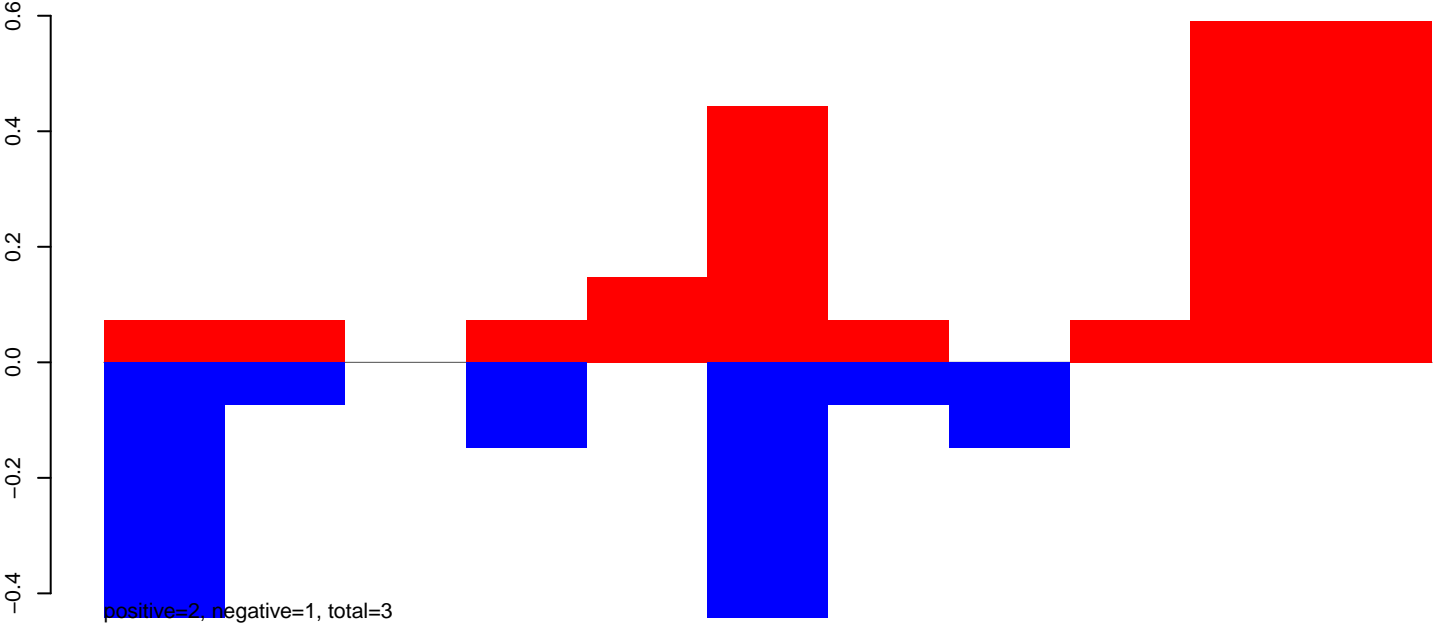
Window size=25, length=401, TE@rnd-4_family-162@Unknown@879:1-401

100 200 300 400

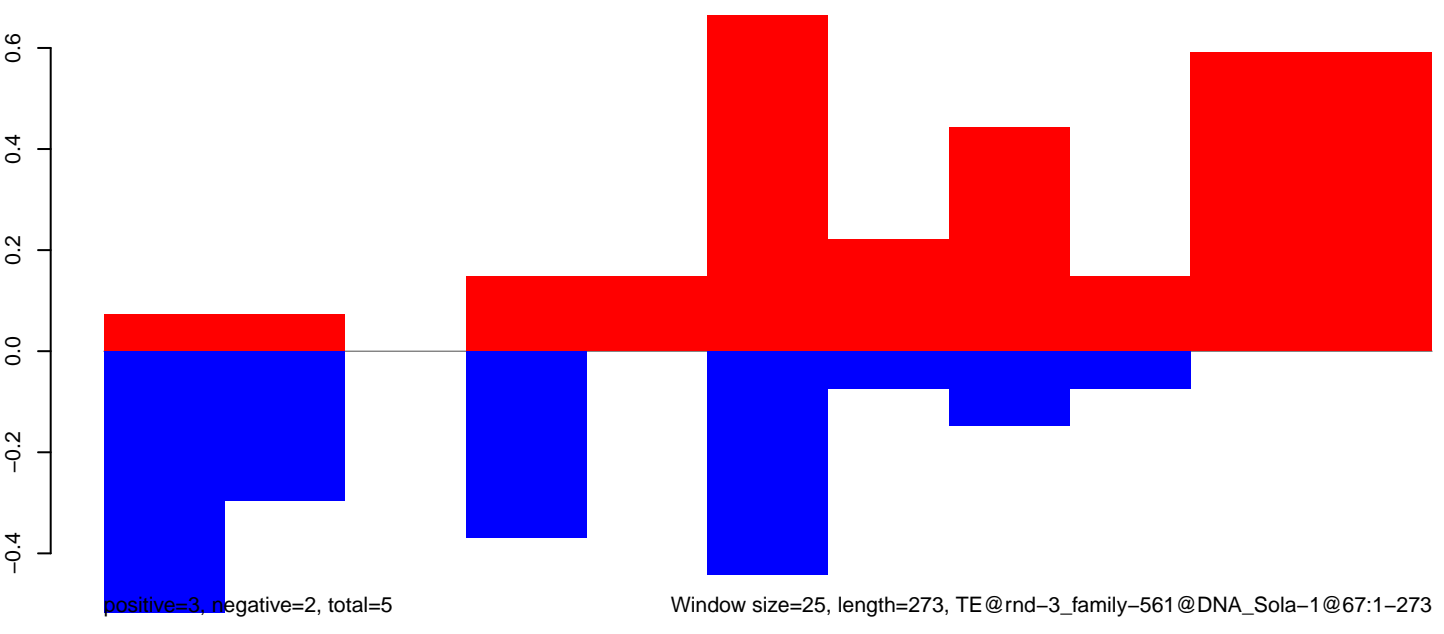
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



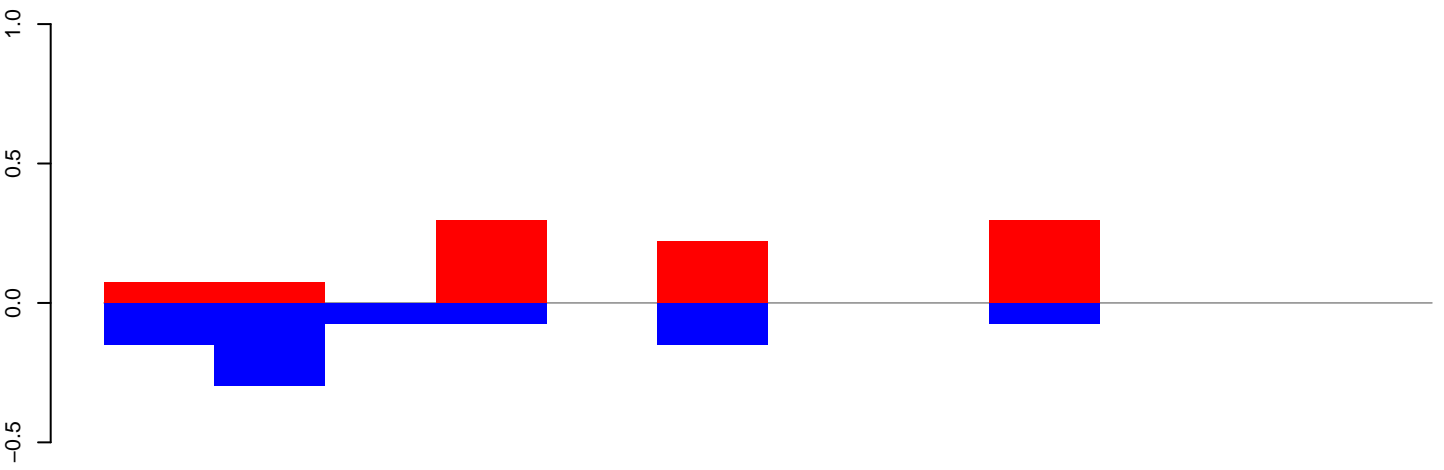
CuQuin_Testes_NL.rep



Window size=25, length=273, TE@rnd-3_family-561@DNA_Sola-1@67:1-273

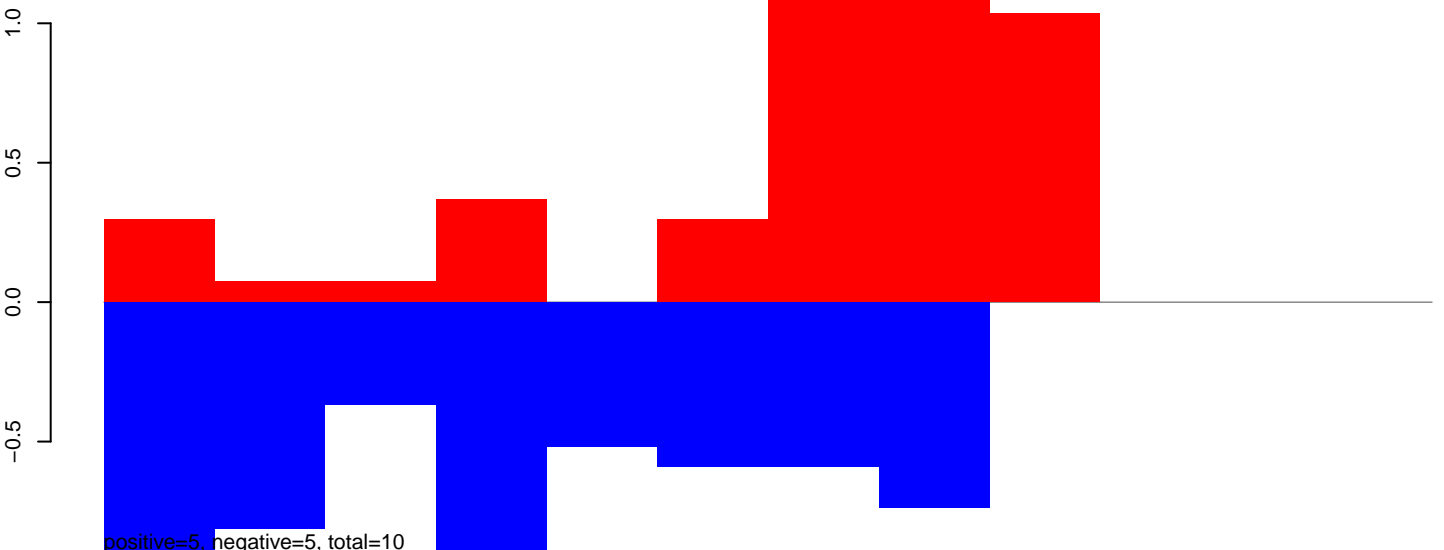
50 100 150 200 250 300

CuQuin_Testes_NL.18_23.rep



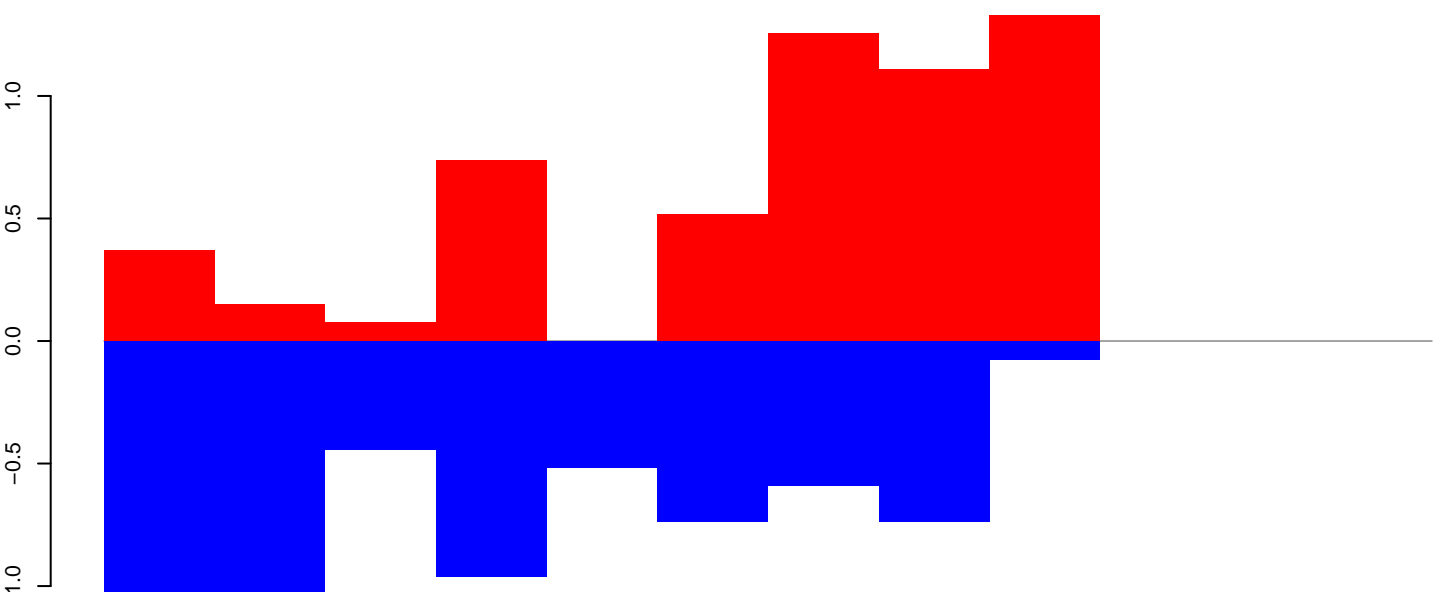
positive=1, negative=1, total=2

CuQuin_Testes_NL.24_35.rep



positive=5, negative=5, total=10

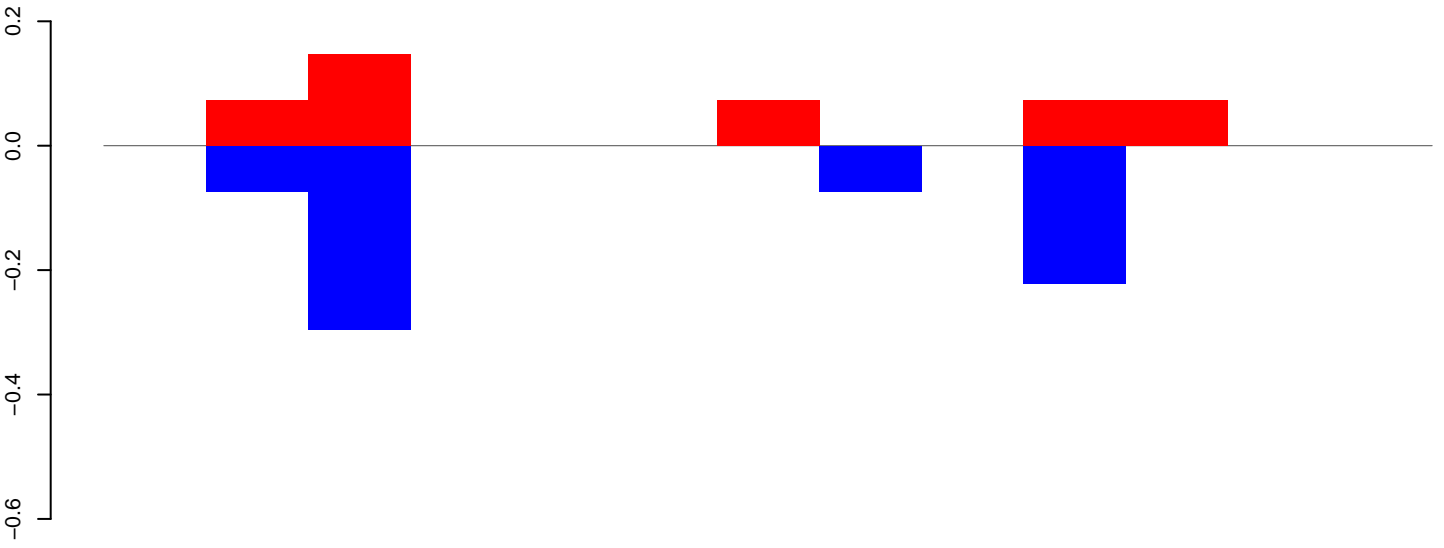
CuQuin_Testes_NL.rep



positive=6, negative=6, total=12

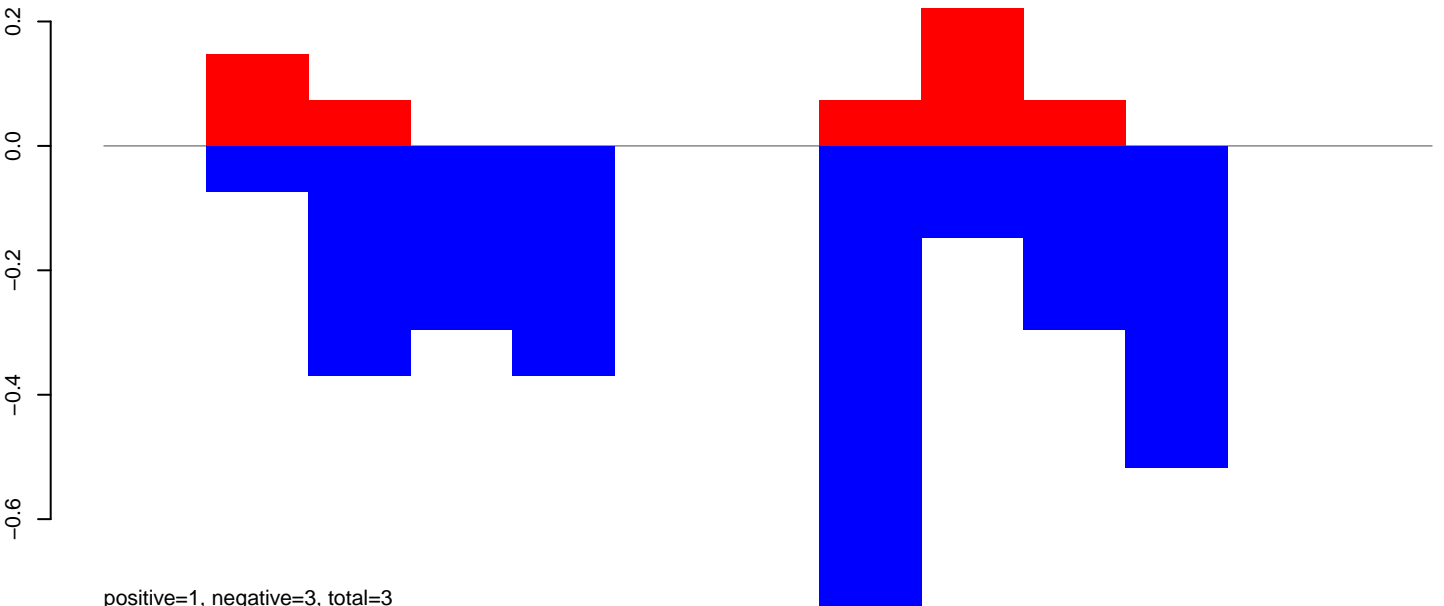
Window size=25, length=279, TE@rnd-3_family-36@Unknown@3340:1-279

CuQuin_Testes_NL.18_23.rep



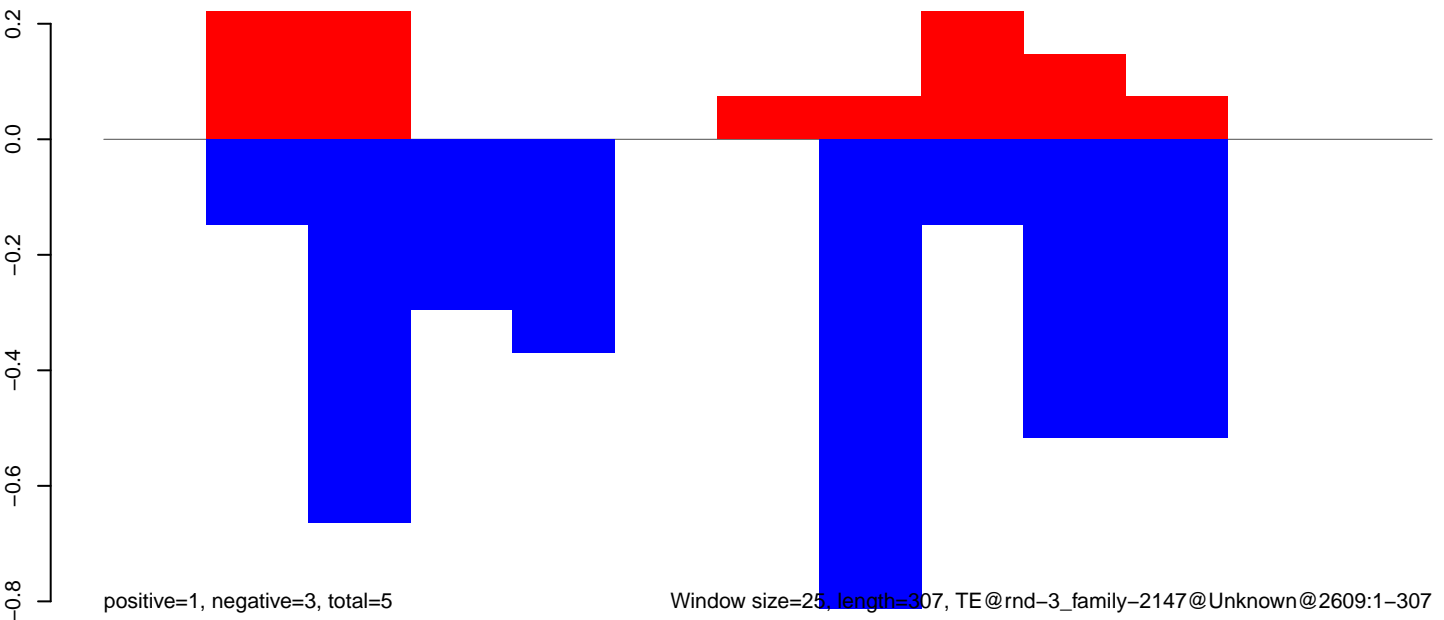
positive=0, negative=1, total=1

CuQuin_Testes_NL.24_35.rep



positive=1, negative=3, total=3

CuQuin_Testes_NL.rep

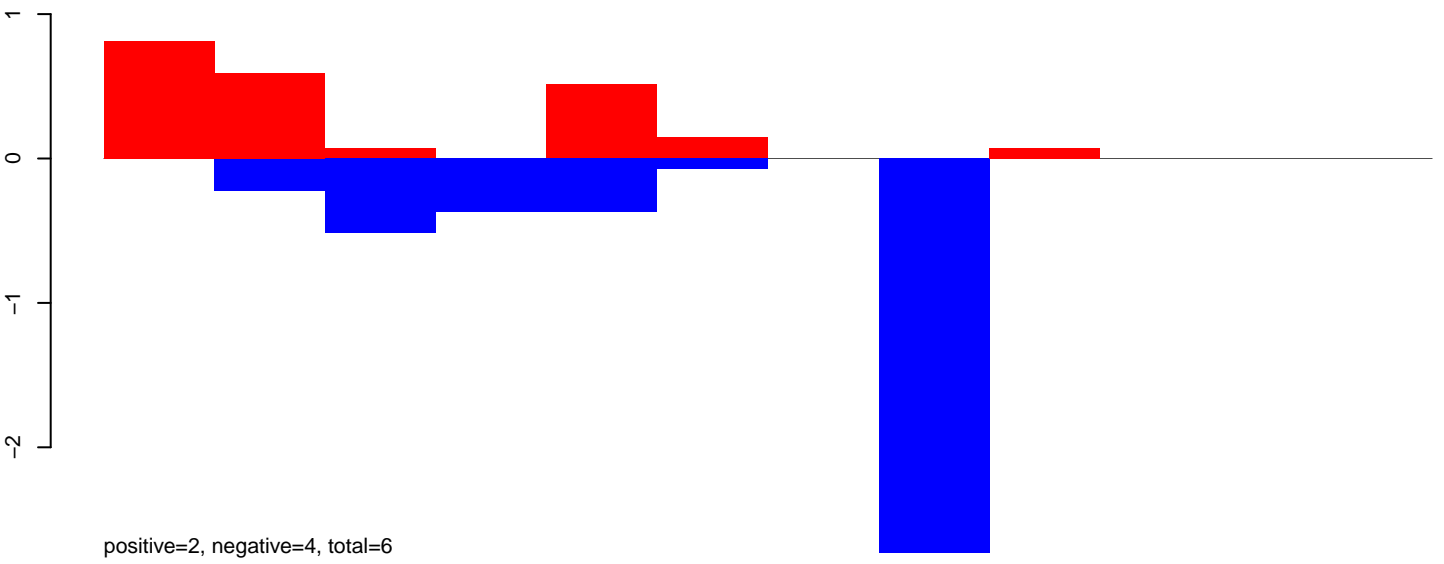


positive=1, negative=3, total=5

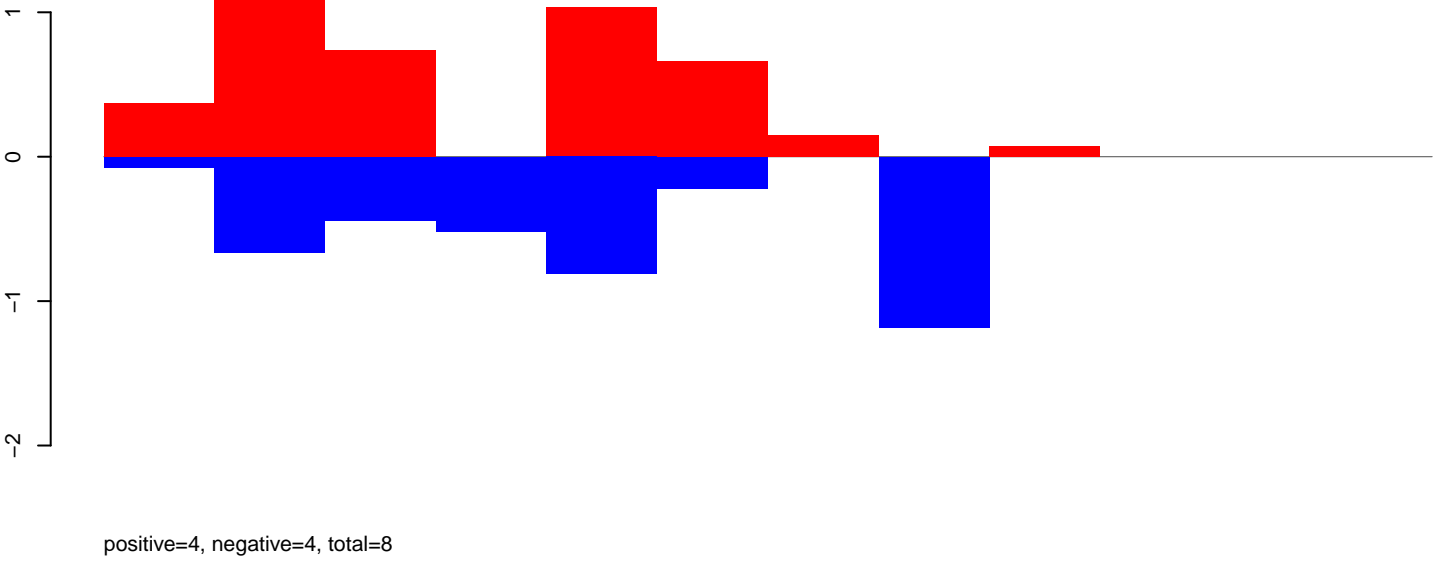
Window size=25, length=307, TE@rnd-3_family-2147@Unknown@2609:1-307

50 100 150 200 250 300 350

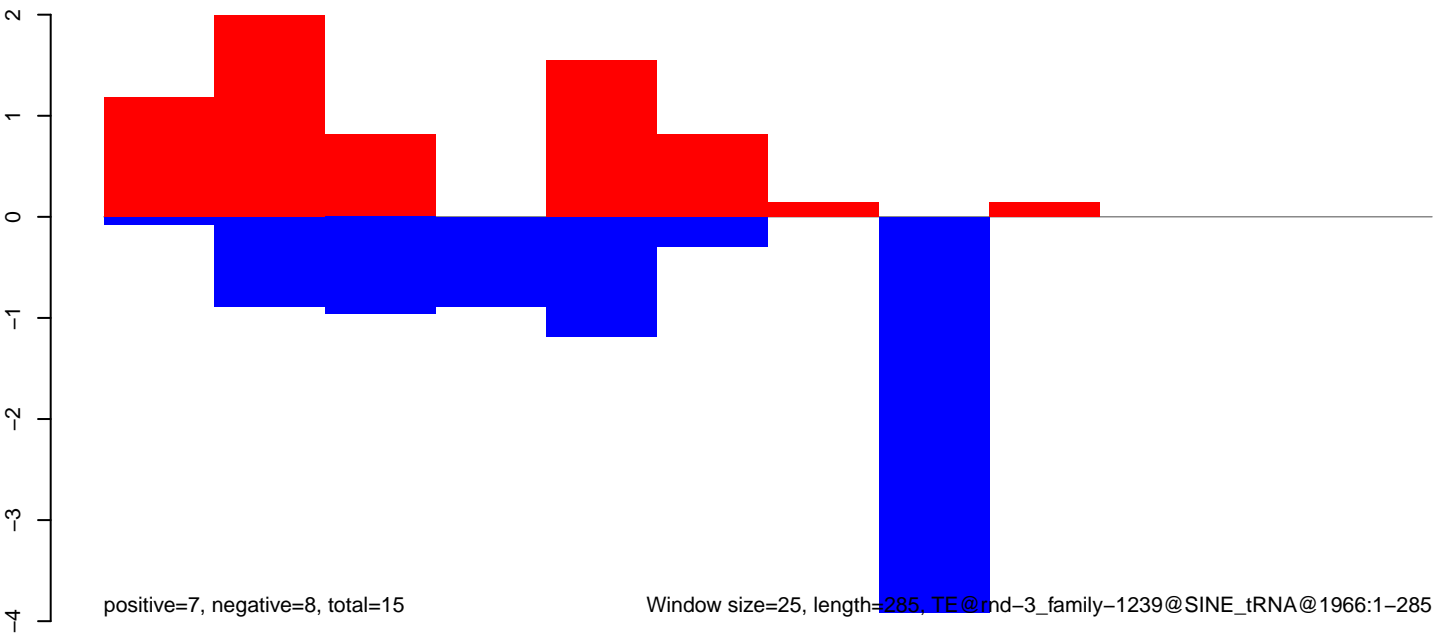
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

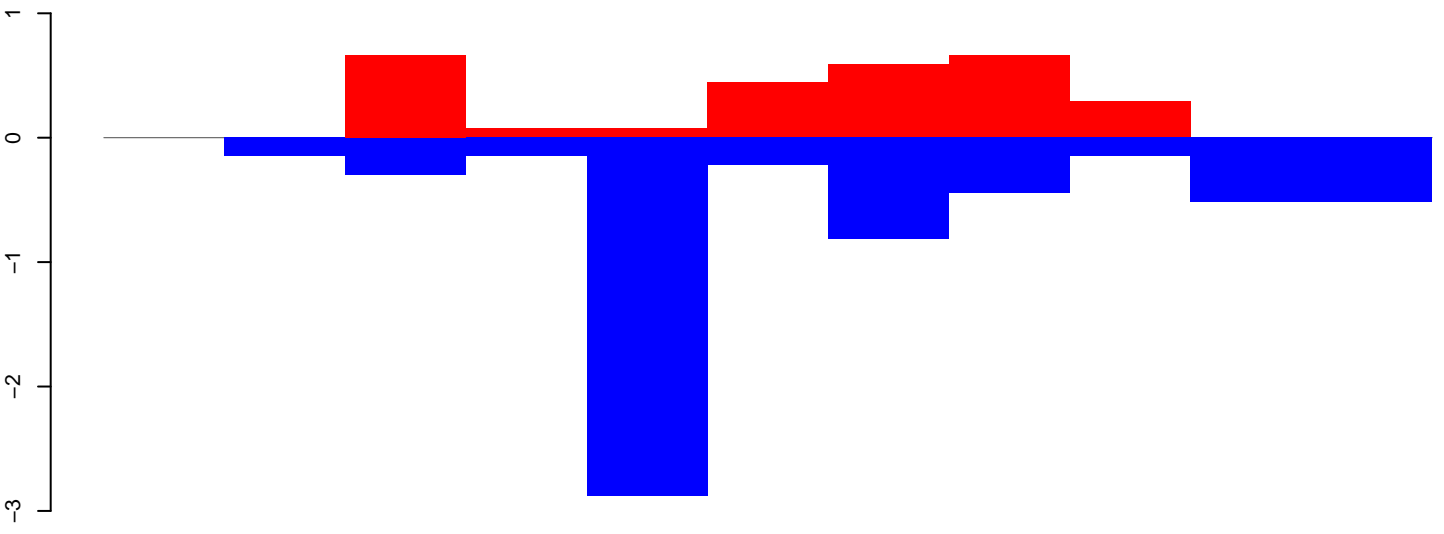


CuQuin_Testes_NL.rep



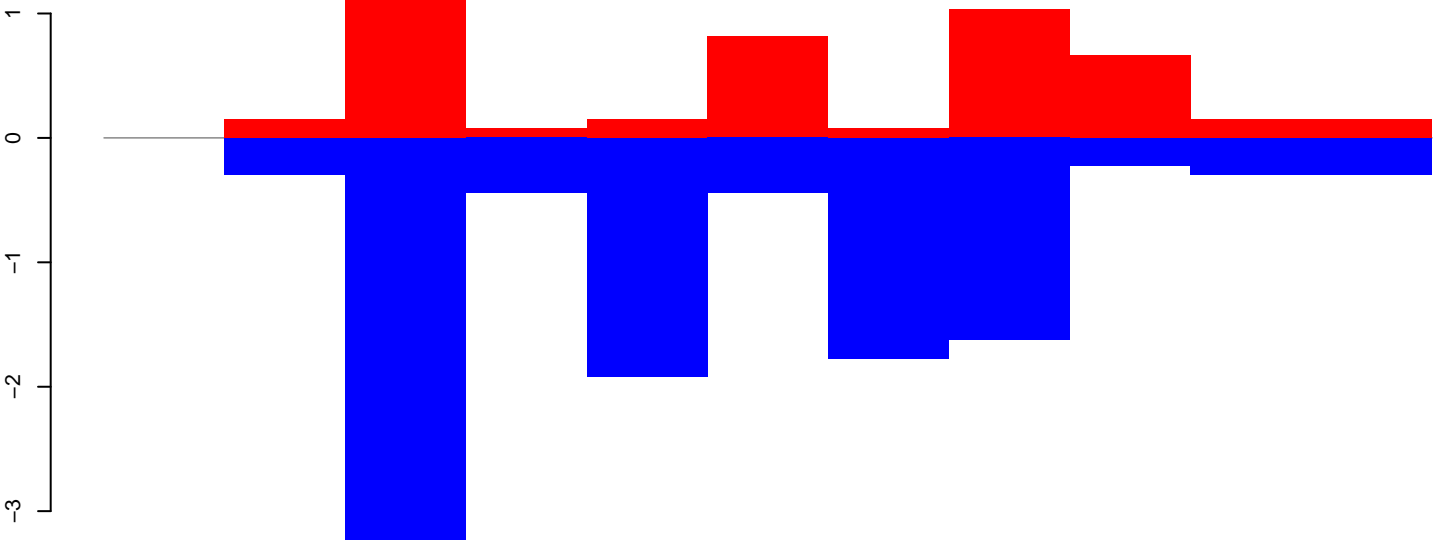
Window size=25, length=285, TE@rnd-3_family-1239@SINE_tRNA@1966:1-285

CuQuin_Testes_NL.18_23.rep



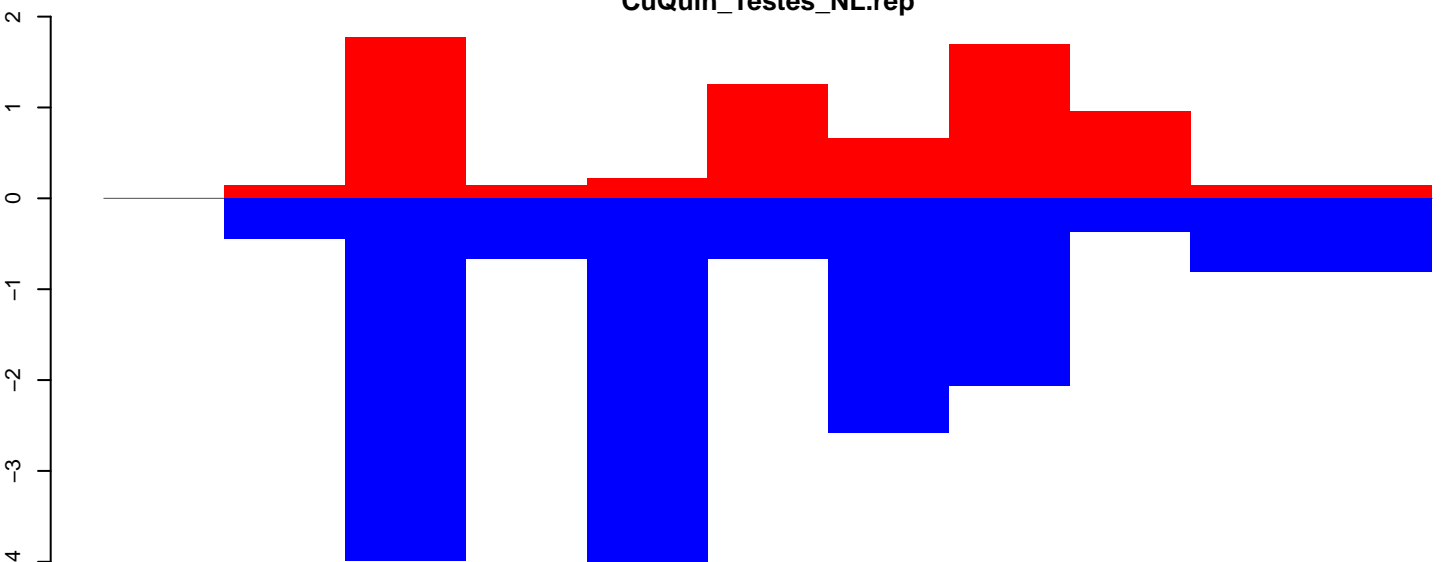
positive=3, negative=6, total=9

CuQuin_Testes_NL.24_35.rep



positive=4, negative=11, total=15

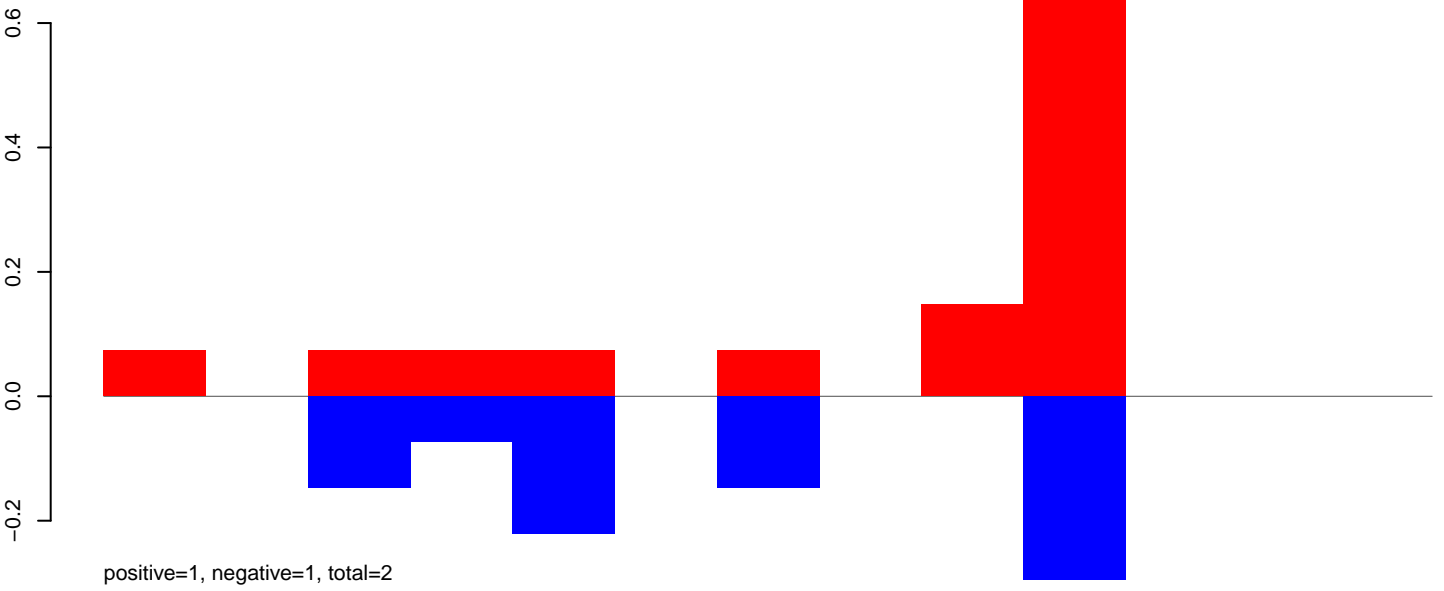
CuQuin_Testes_NL.rep



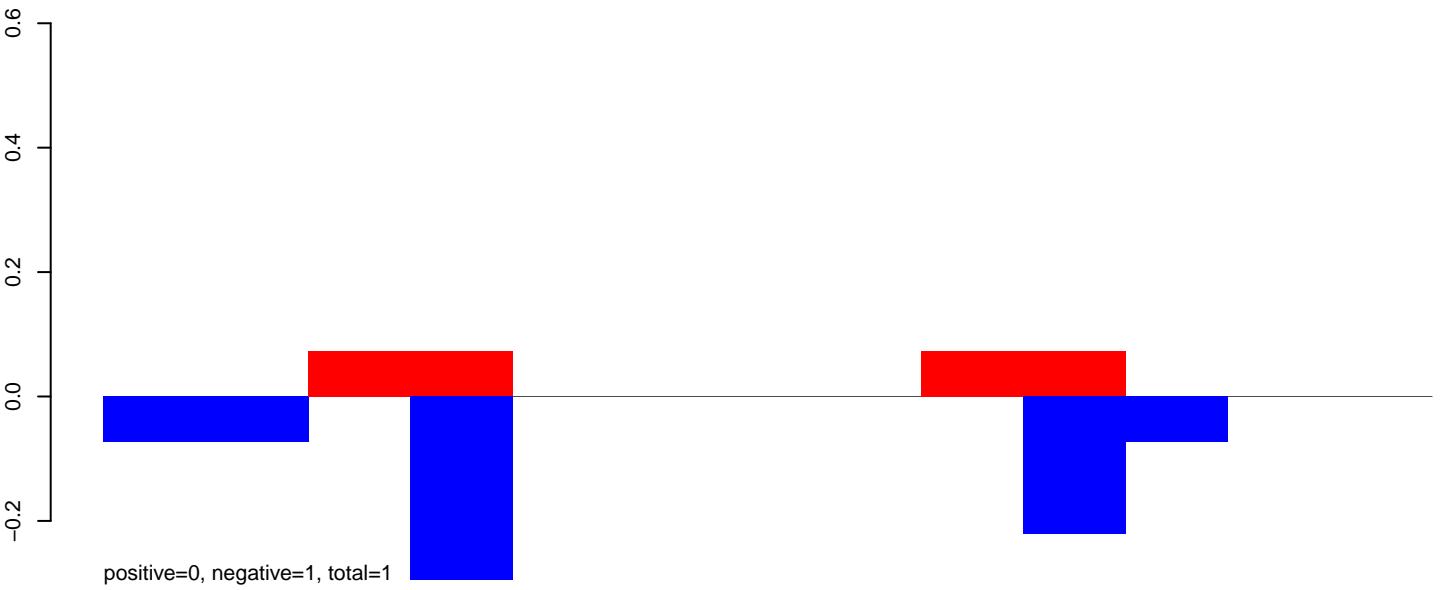
positive=7, negative=17, total=24

Window size=25, length=273, TE@rnd-3_family-1045@DNA@2025:1-273

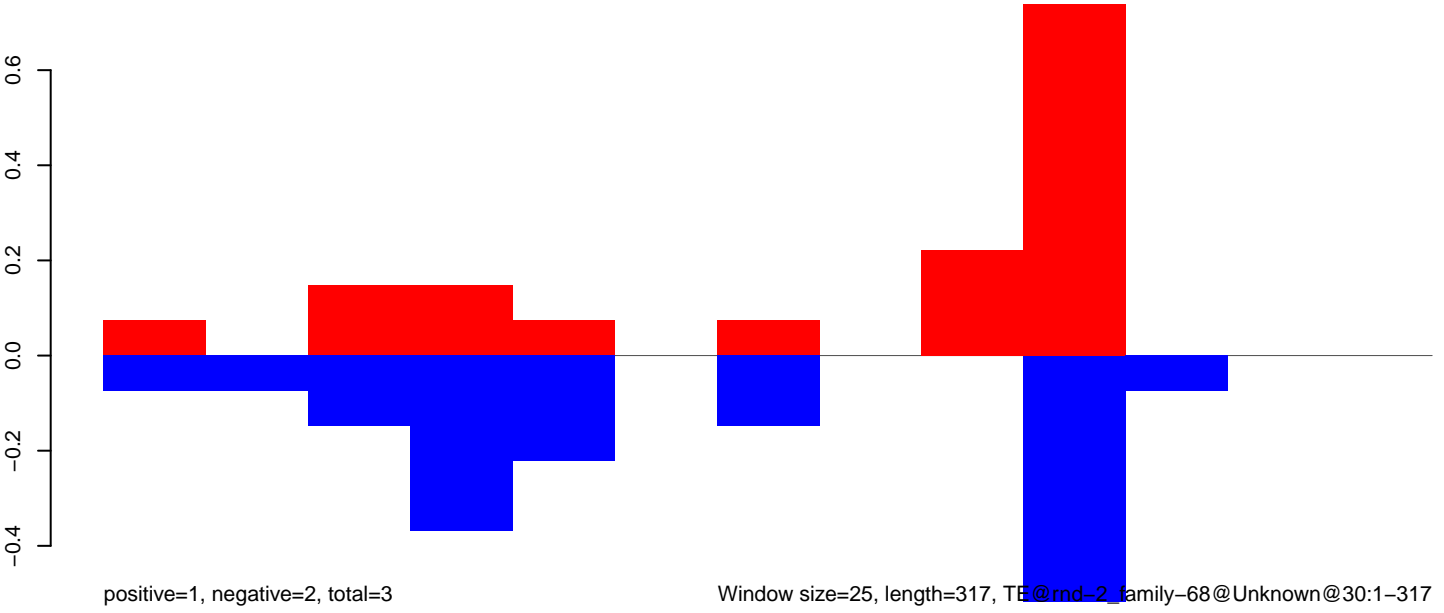
CuQuin_Testes_NL.18_23.rep



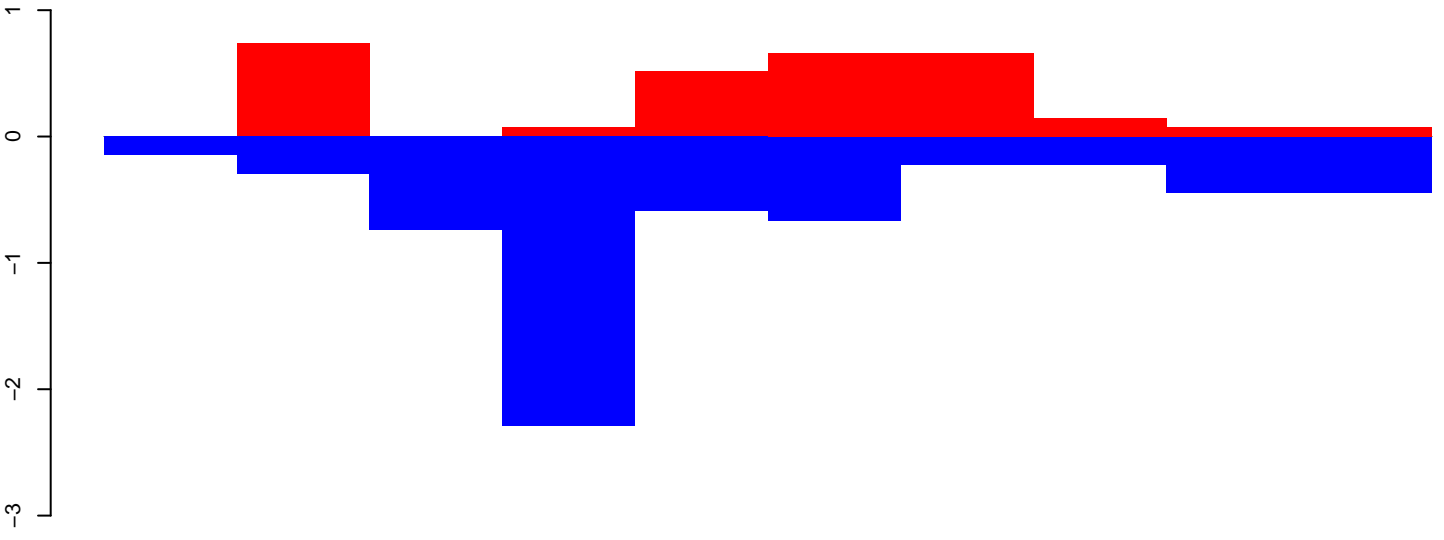
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

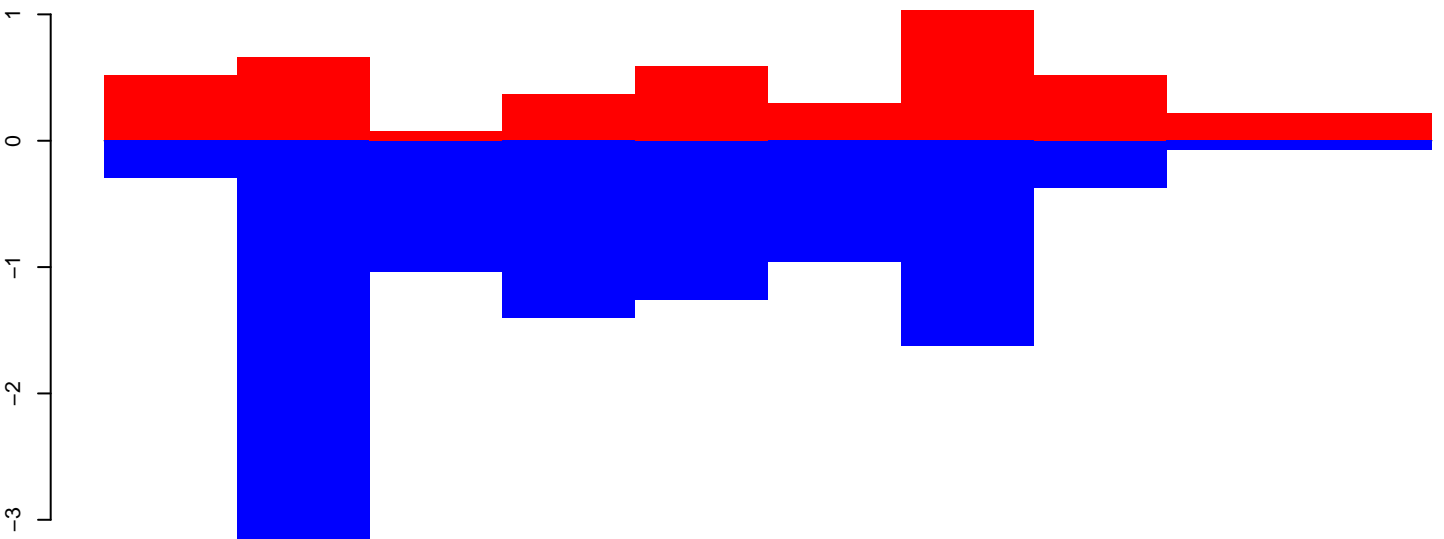


CuQuin_Testes_NL.18_23.rep



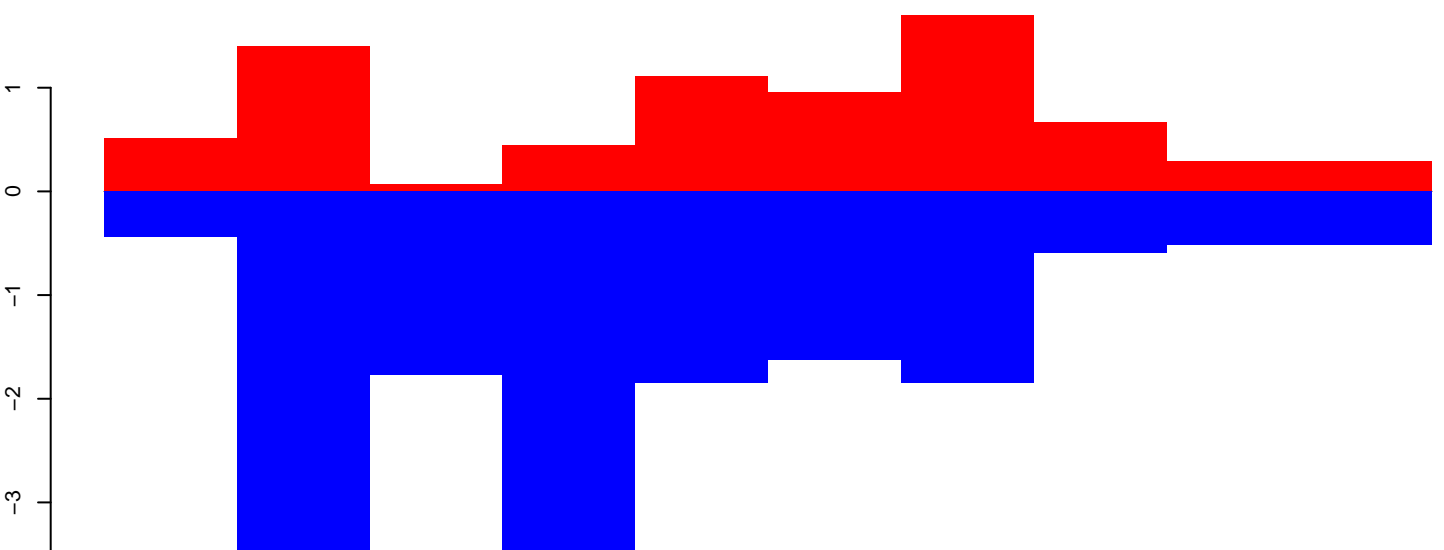
positive=3, negative=6, total=9

CuQuin_Testes_NL.24_35.rep



positive=5, negative=11, total=15

CuQuin_Testes_NL.rep

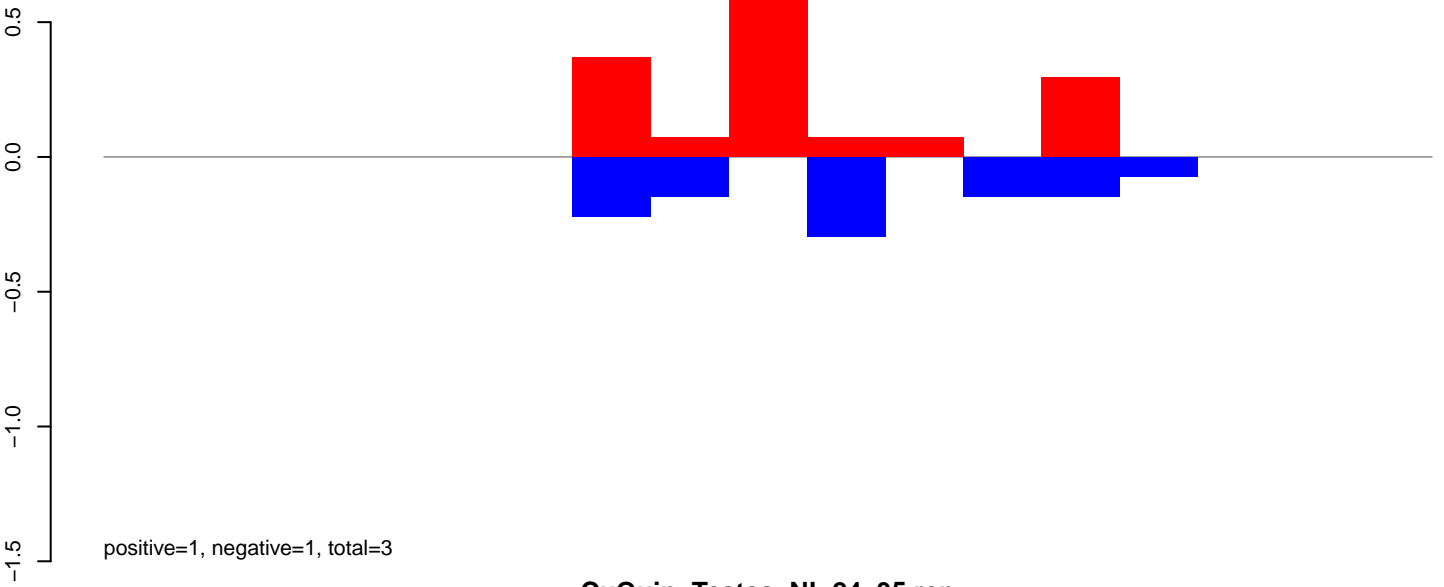


positive=7, negative=17, total=24

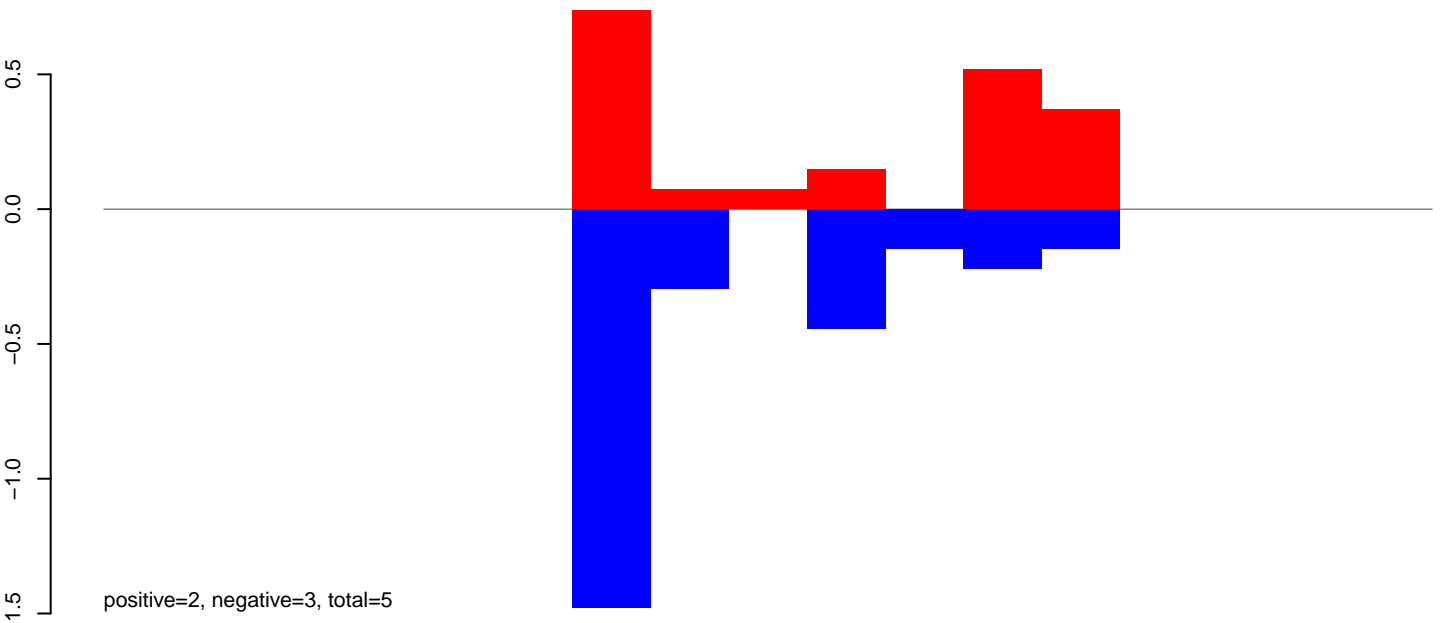
Window size=25, length=248, TE@rnd-2_family-271@LTR_Copia@3254:1-248

50 100 150 200 250

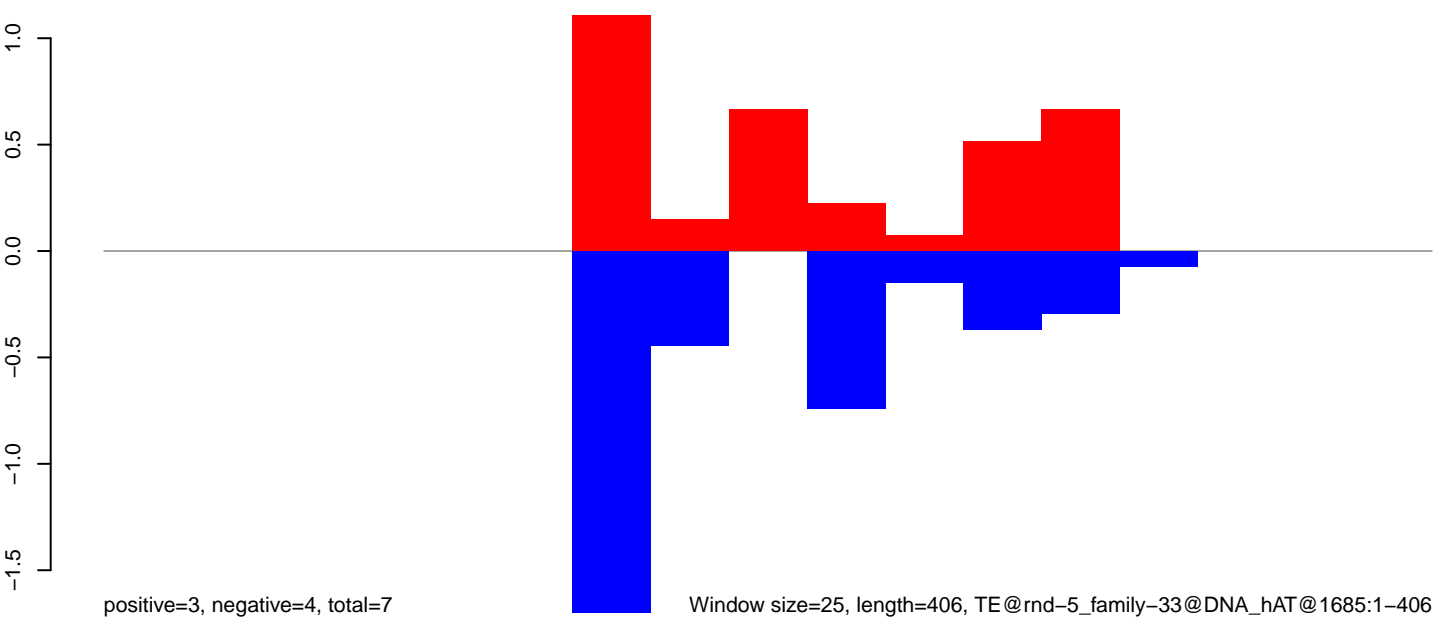
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



Window size=25, length=406, TE@rnd-5_family-33@DNA_hAT@1685:1-406

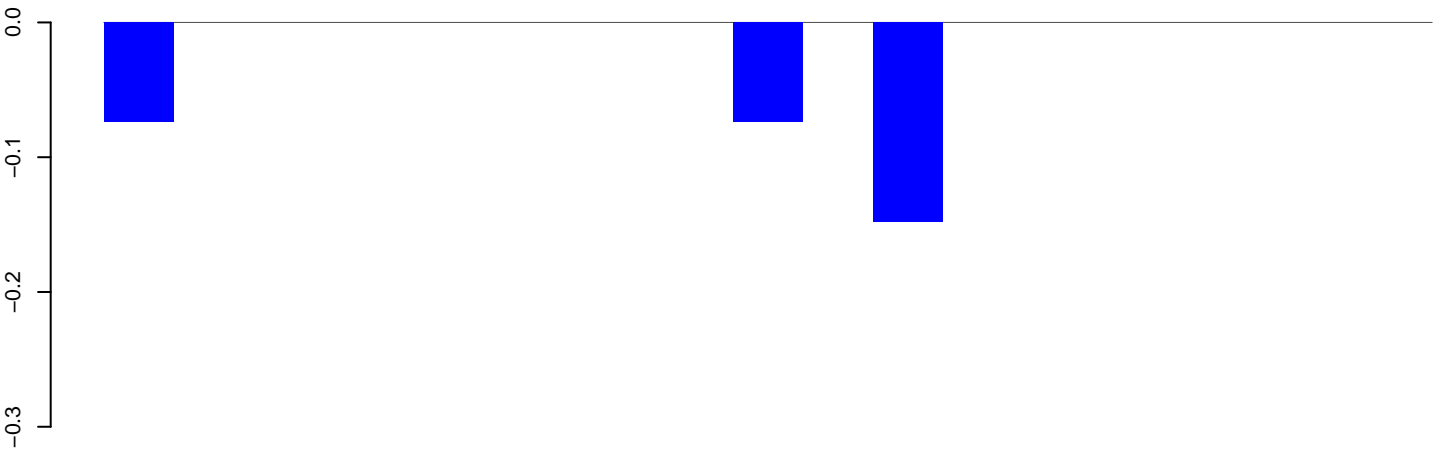
100

200

300

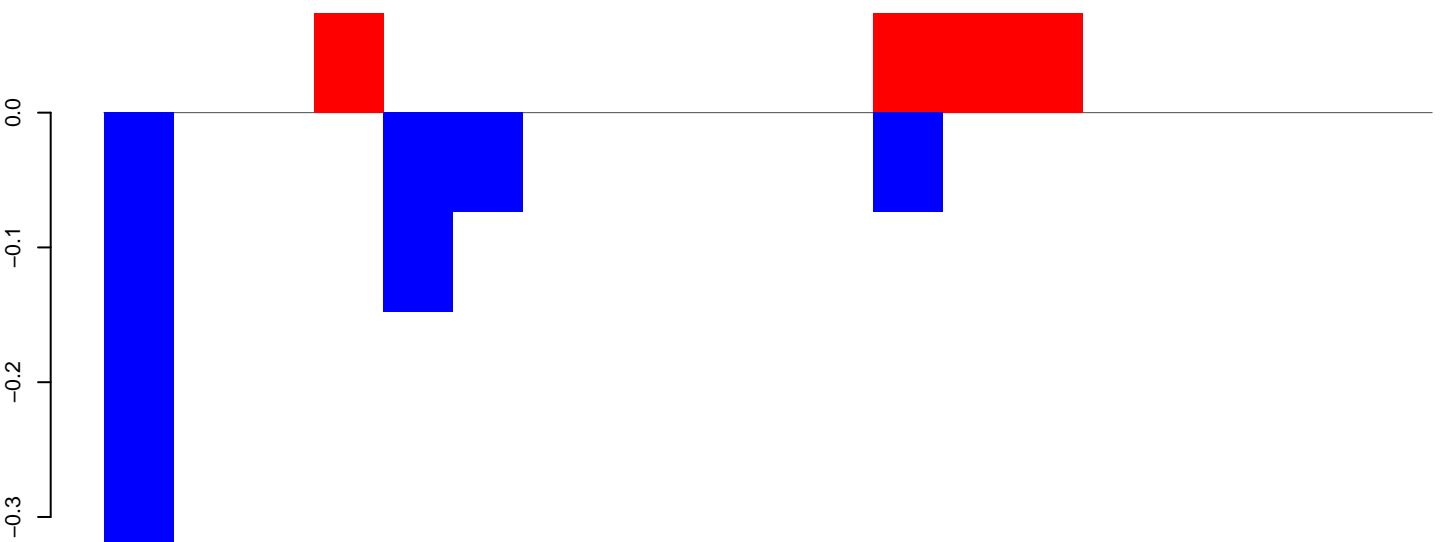
400

CuQuin_Testes_NL.18_23.rep



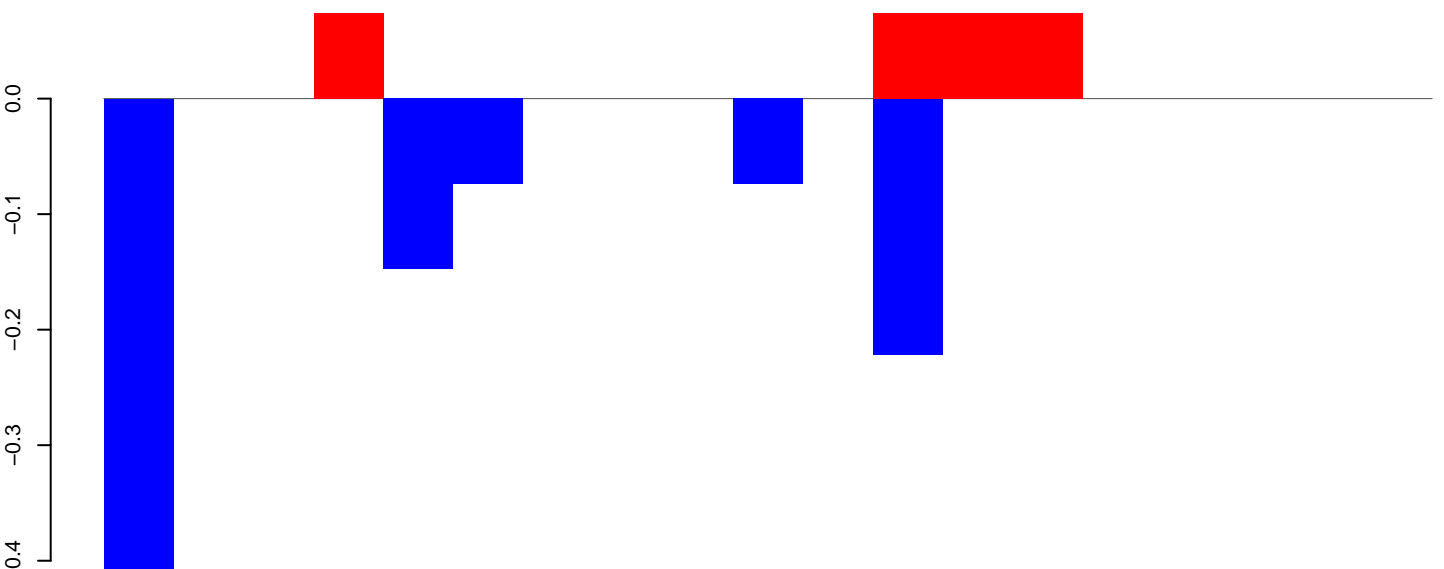
positive=0, negative=0, total=0

CuQuin_Testes_NL.24_35.rep



positive=0, negative=1, total=1

CuQuin_Testes_NL.rep

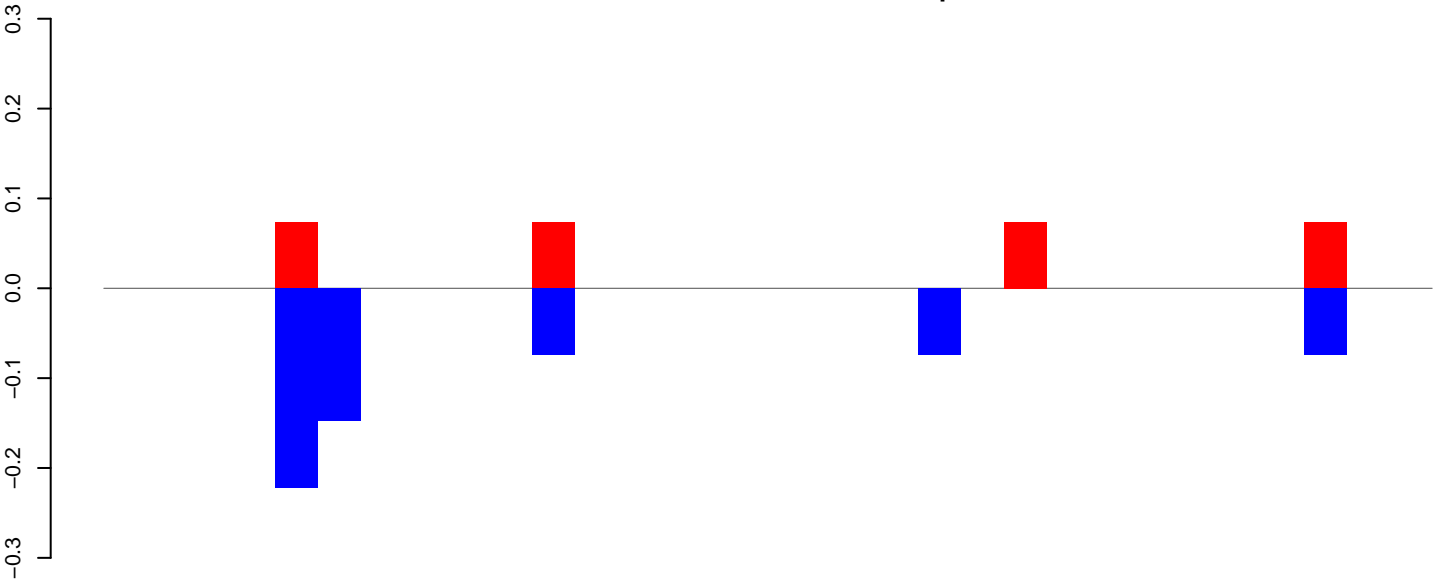


positive=0, negative=1, total=1

Window size=25, length=451, TE@rnd-5_family-21@DNA_MULE-MuDR@3707:1-451

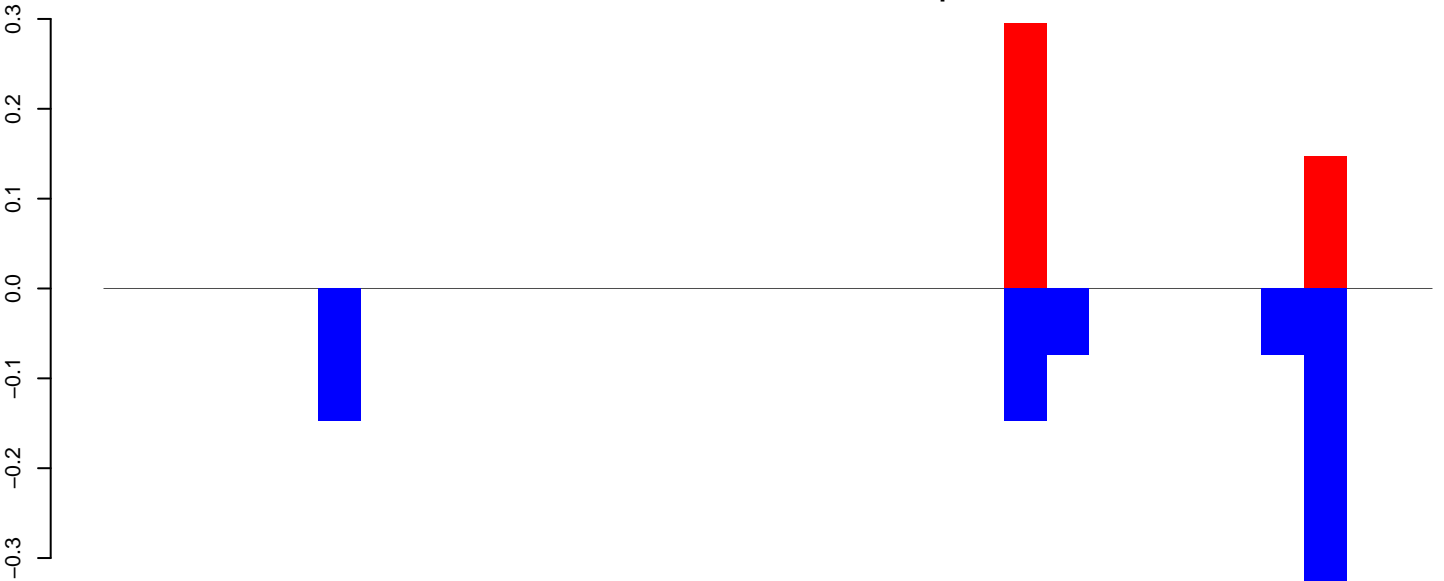
100 200 300 400 500

CuQuin_Testes_NL.18_23.rep



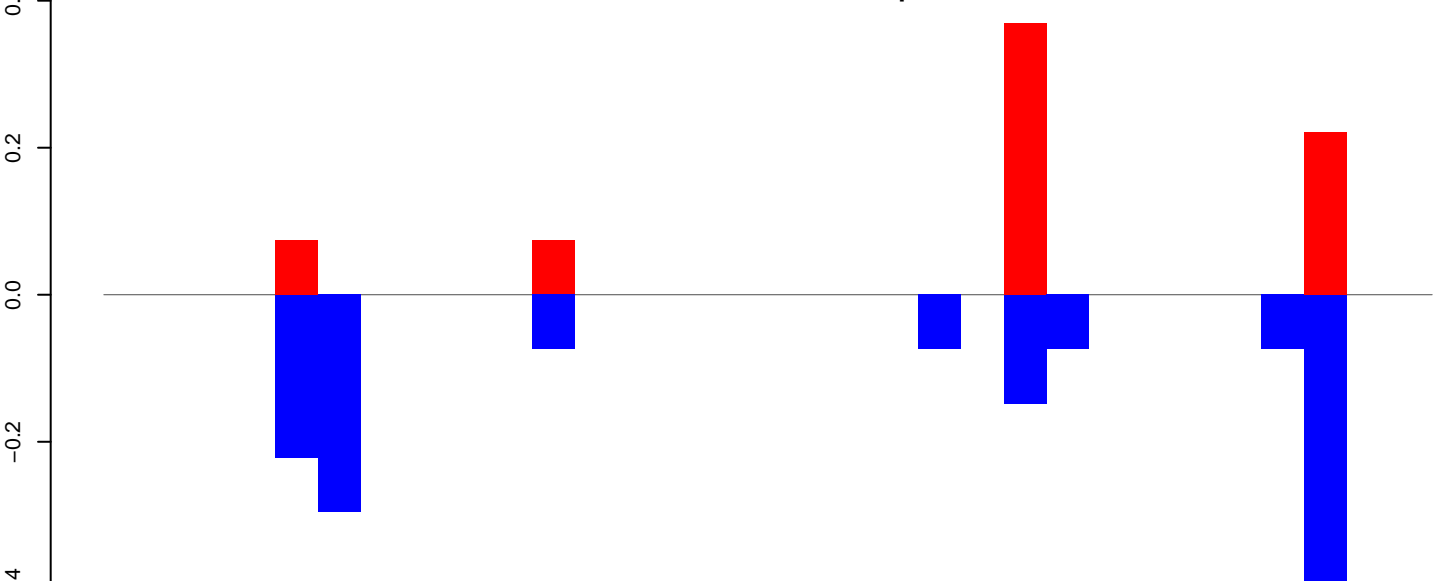
positive=0, negative=1, total=1

CuQuin_Testes_NL.24_35.rep



positive=0, negative=1, total=1

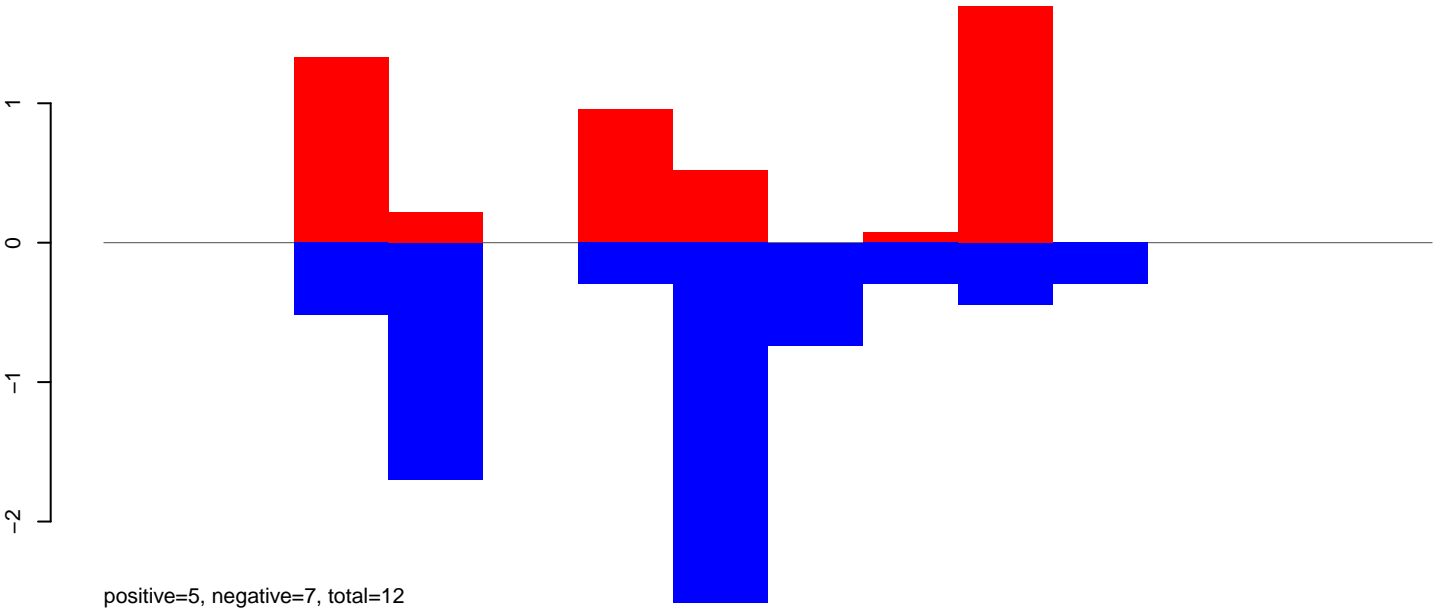
CuQuin_Testes_NL.rep



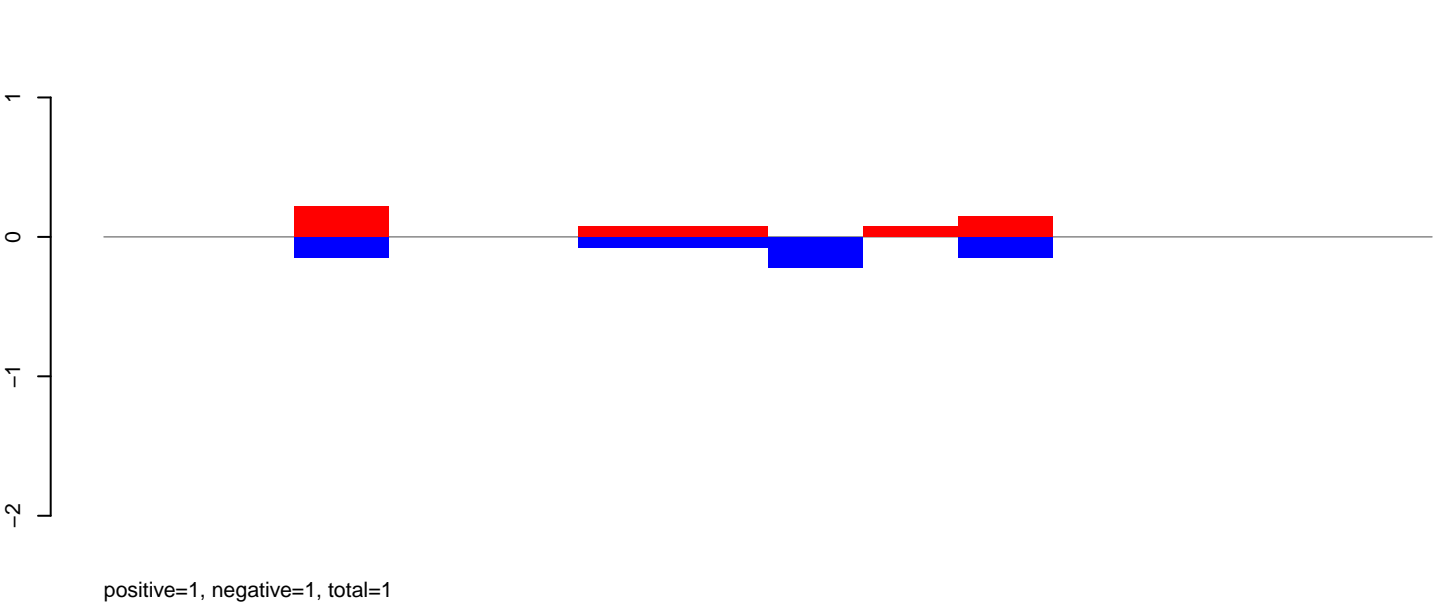
positive=1, negative=1, total=2

Window size=25, length=758, TE@rnd-5_family-1396@Satellite@1138:1-758

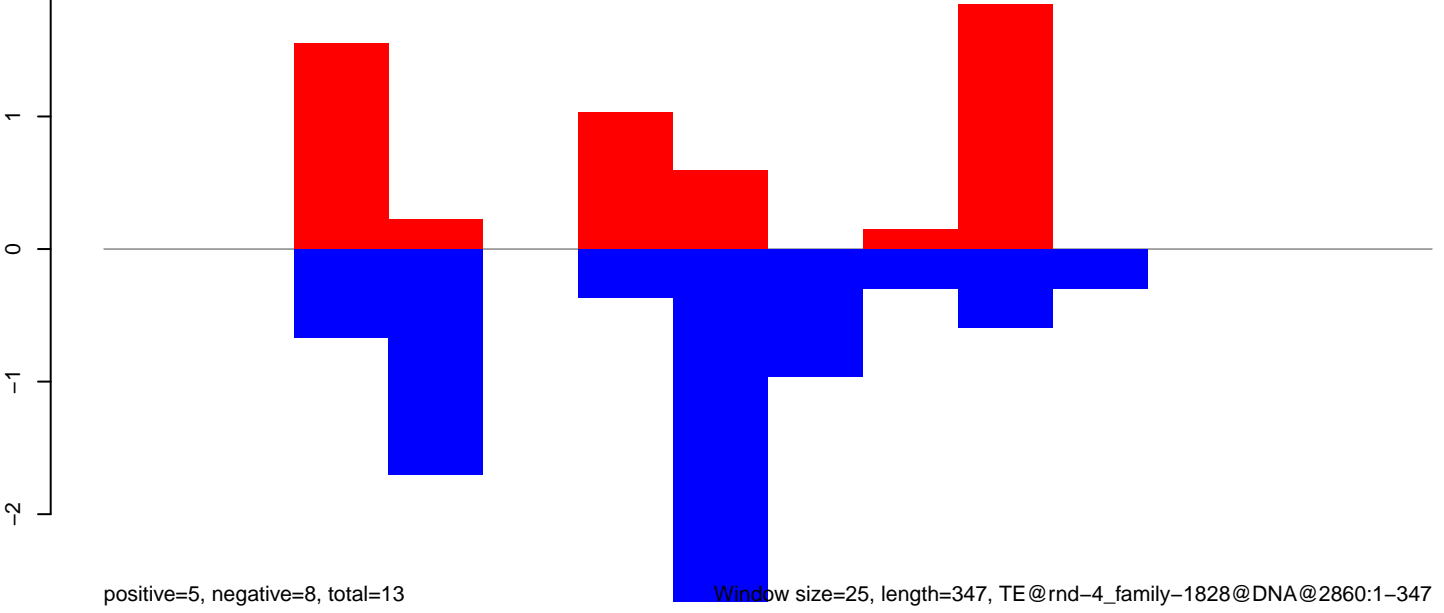
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



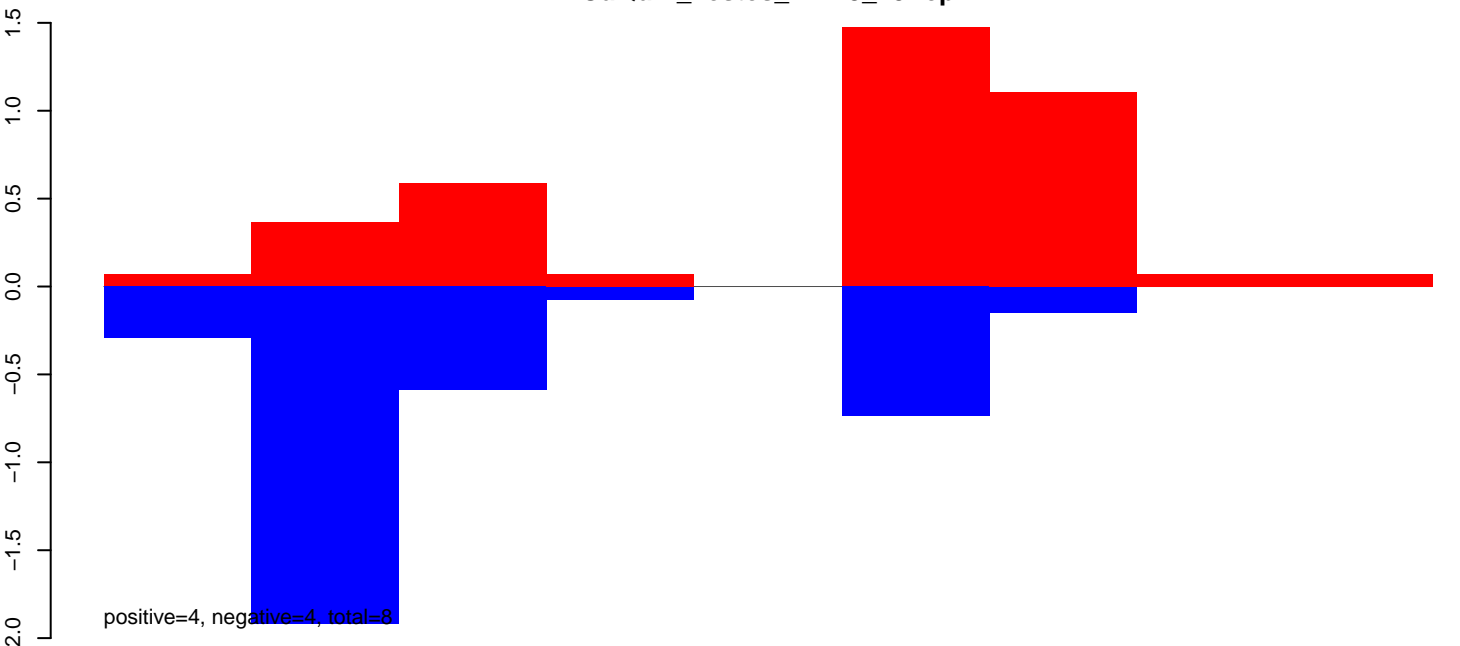
CuQuin_Testes_NL.rep



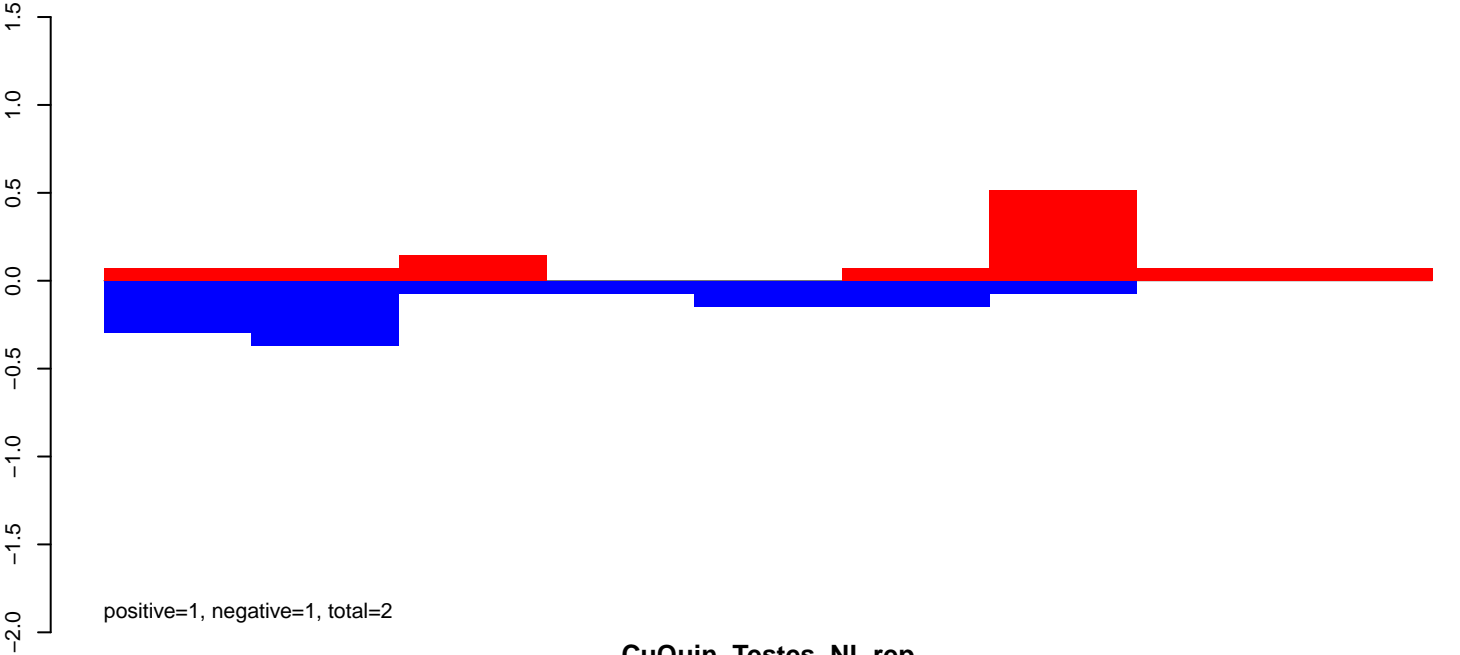
Window size=25, length=347, TE@rnd-4_family-1828@DNA@2860:1-347

50 100 150 200 250 300 350

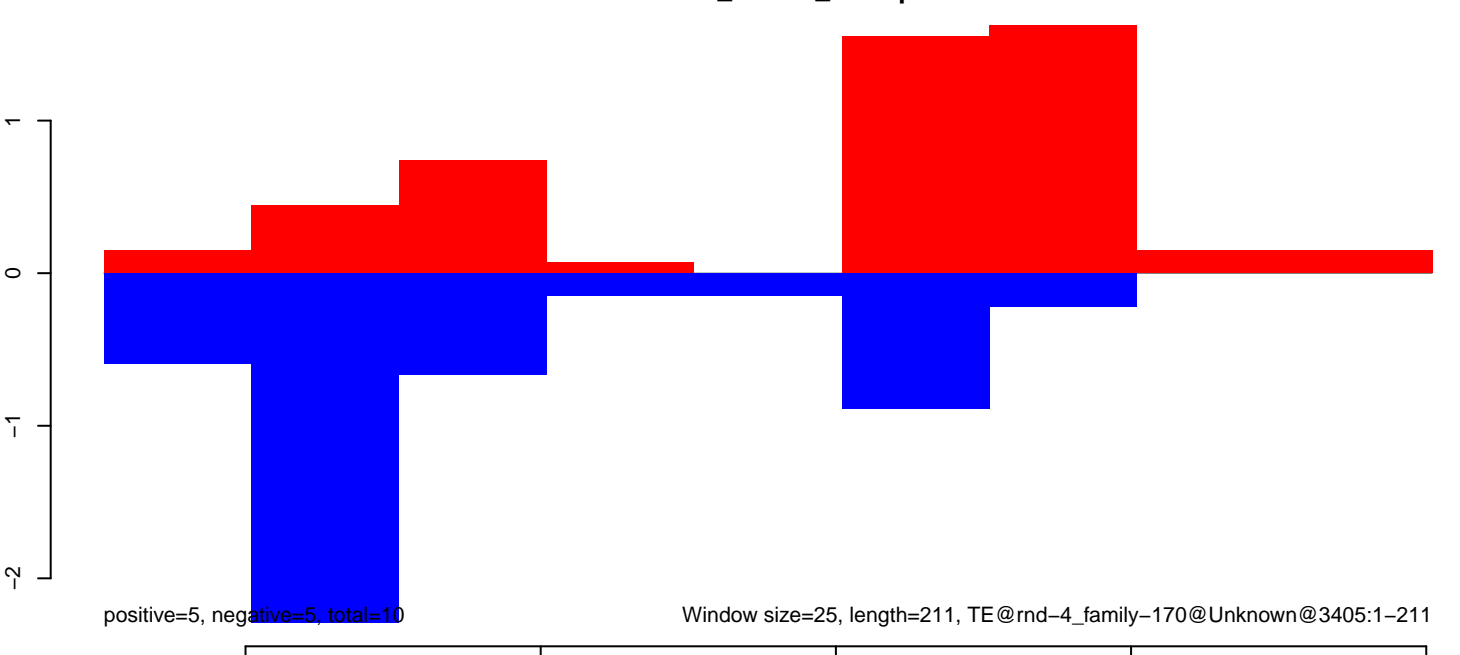
CuQuin_Testes_NL.18_23.rep



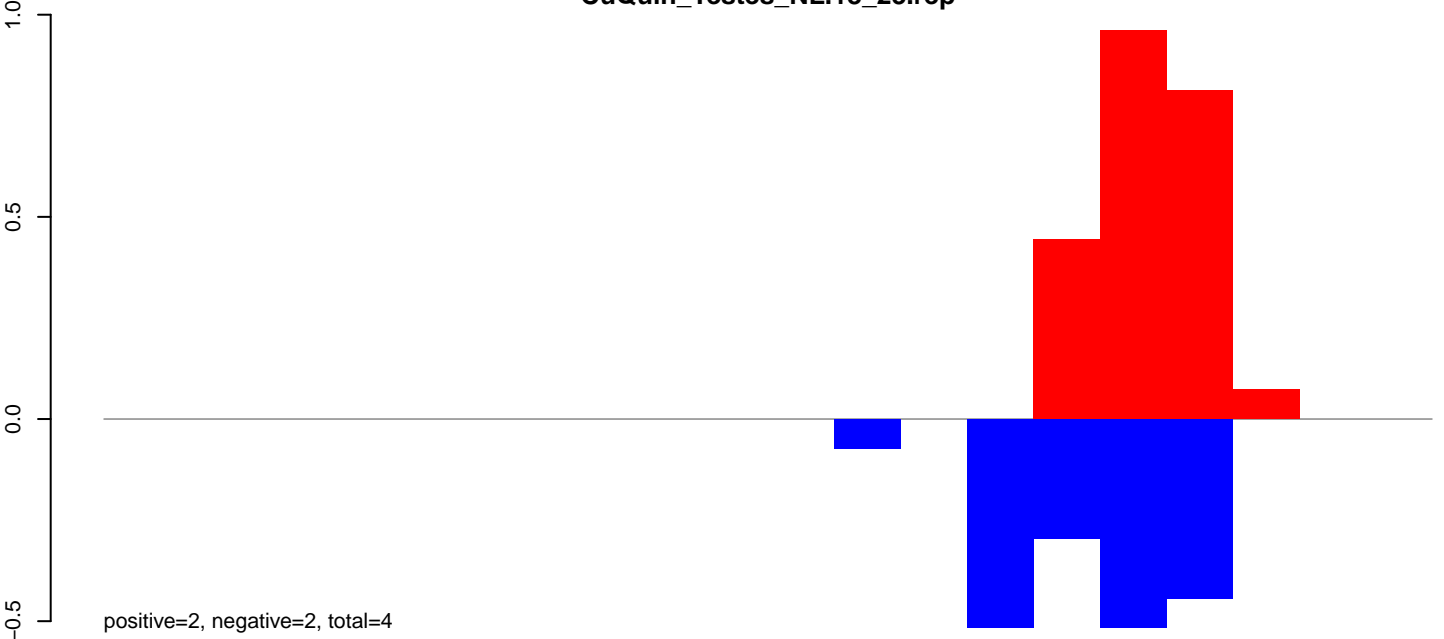
CuQuin_Testes_NL.24_35.rep



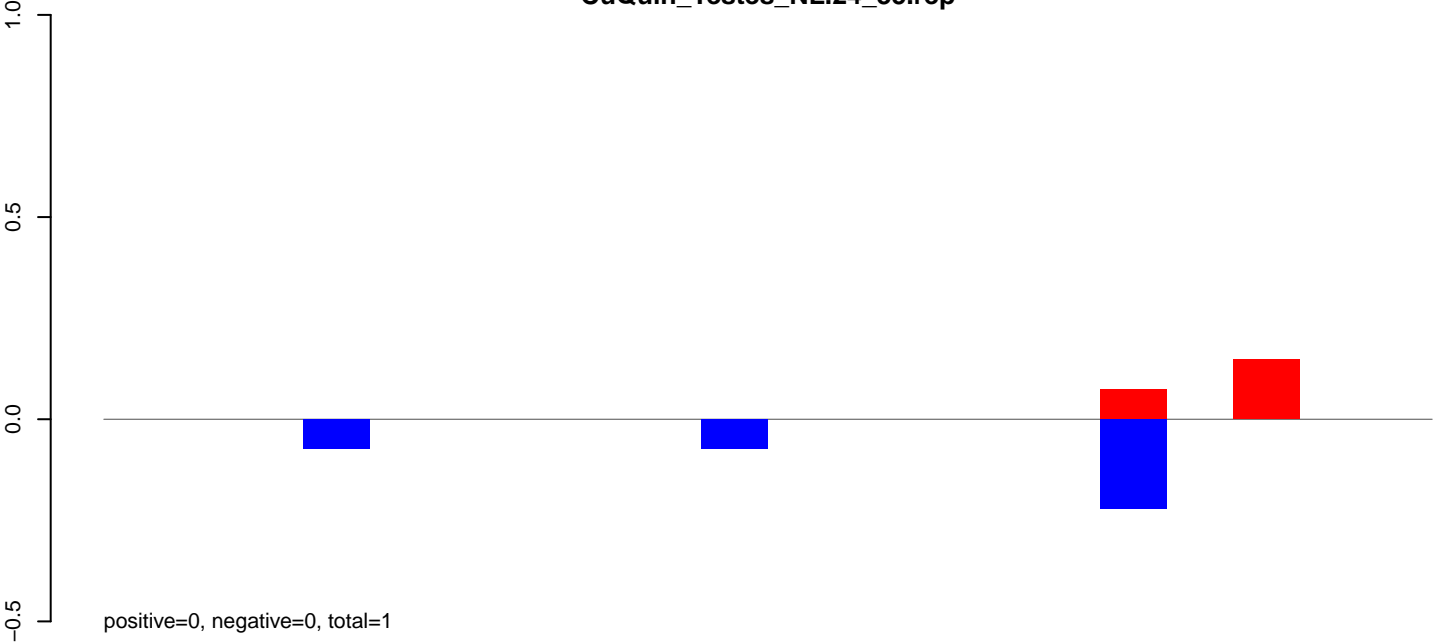
CuQuin_Testes_NL.rep



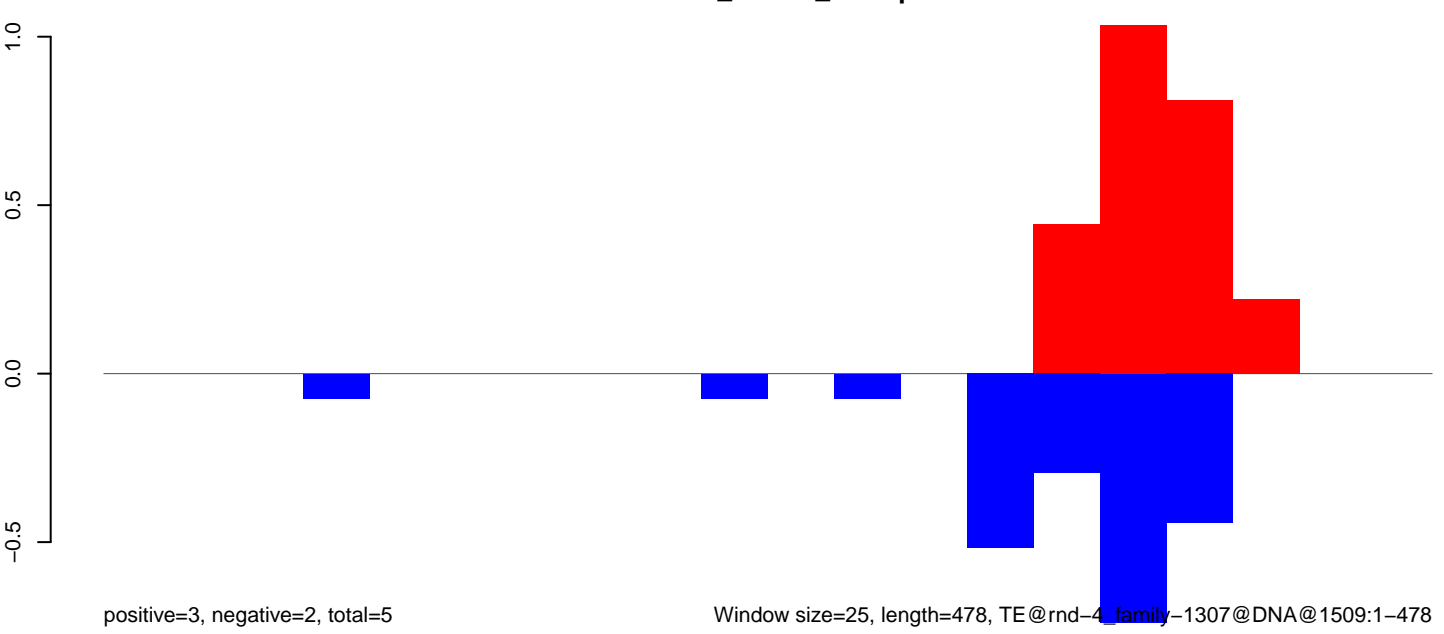
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

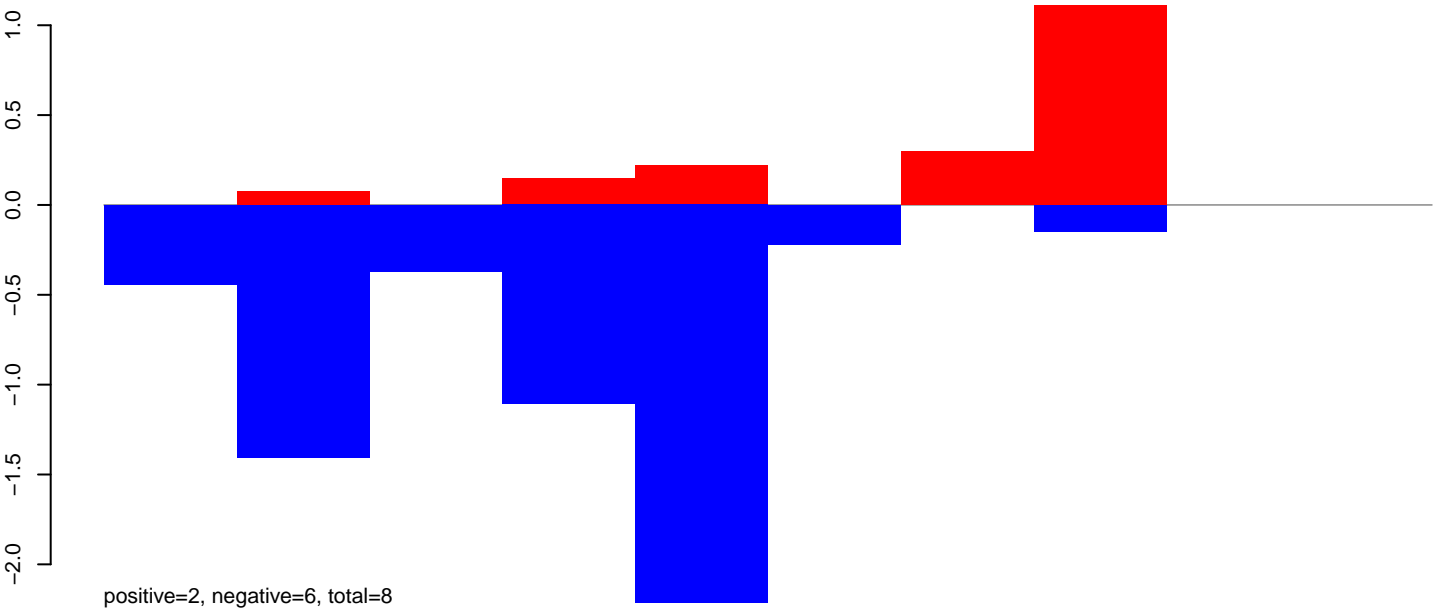


CuQuin_Testes_NL.rep

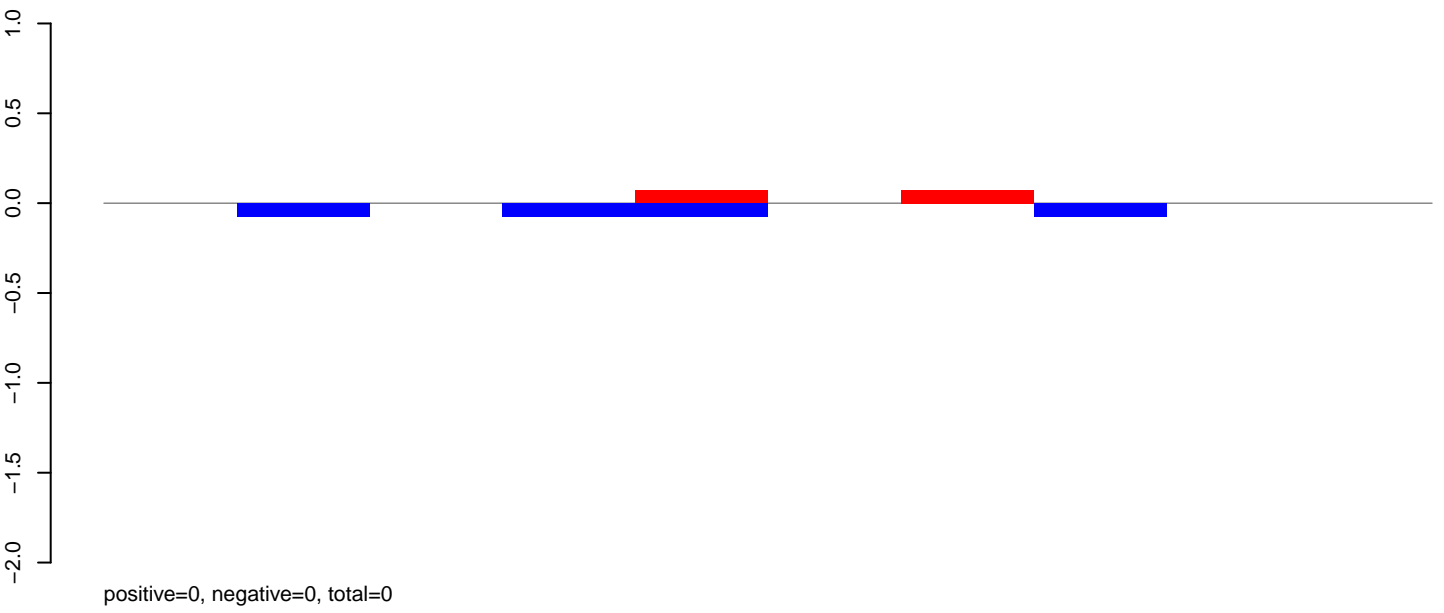


Window size=25, length=478, TE@rnd-4_family-1307@DNA@1509:1-478

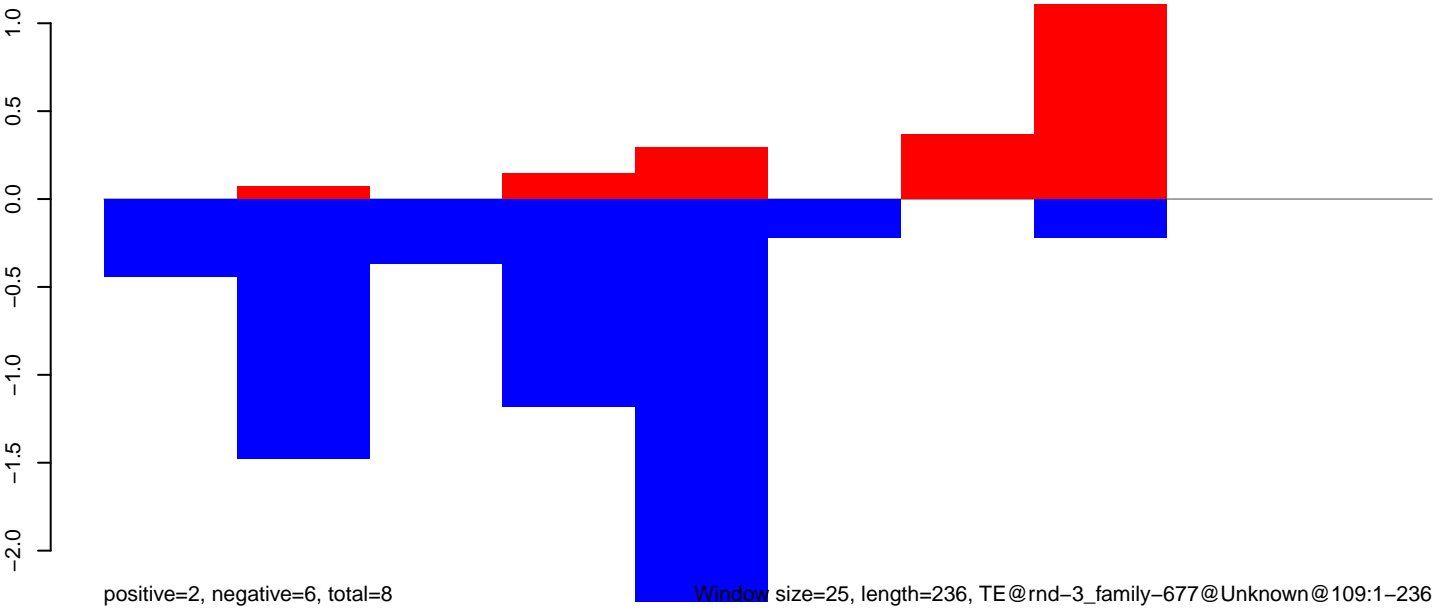
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



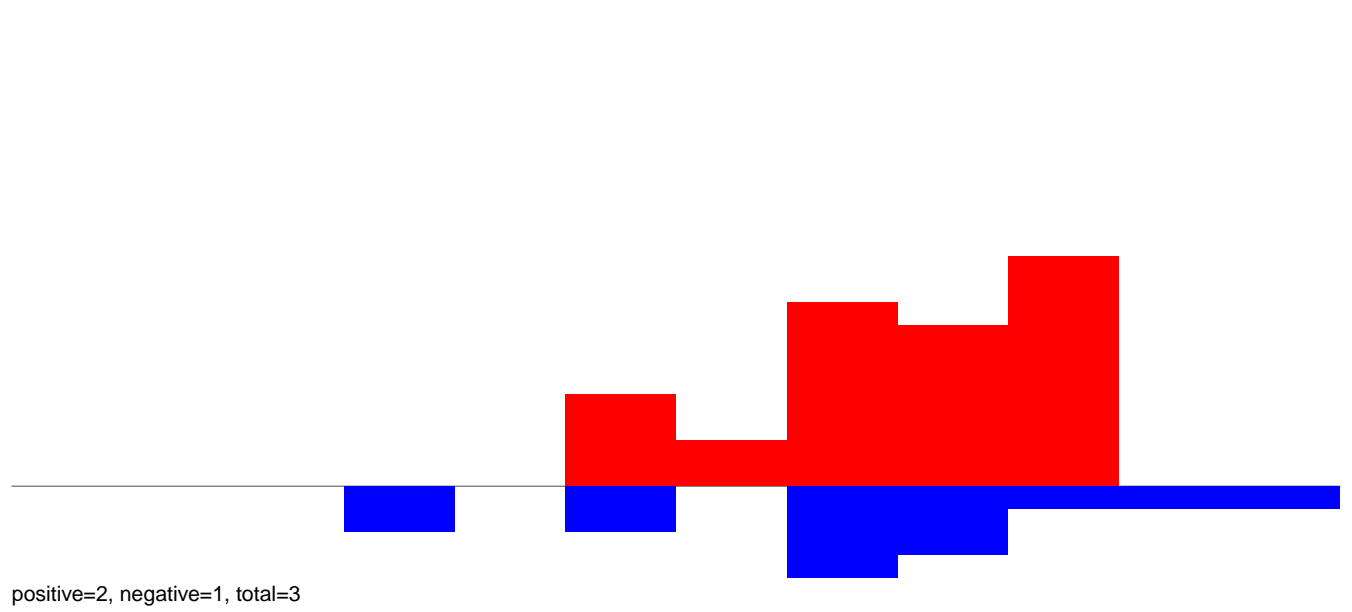
CuQuin_Testes_NL.rep



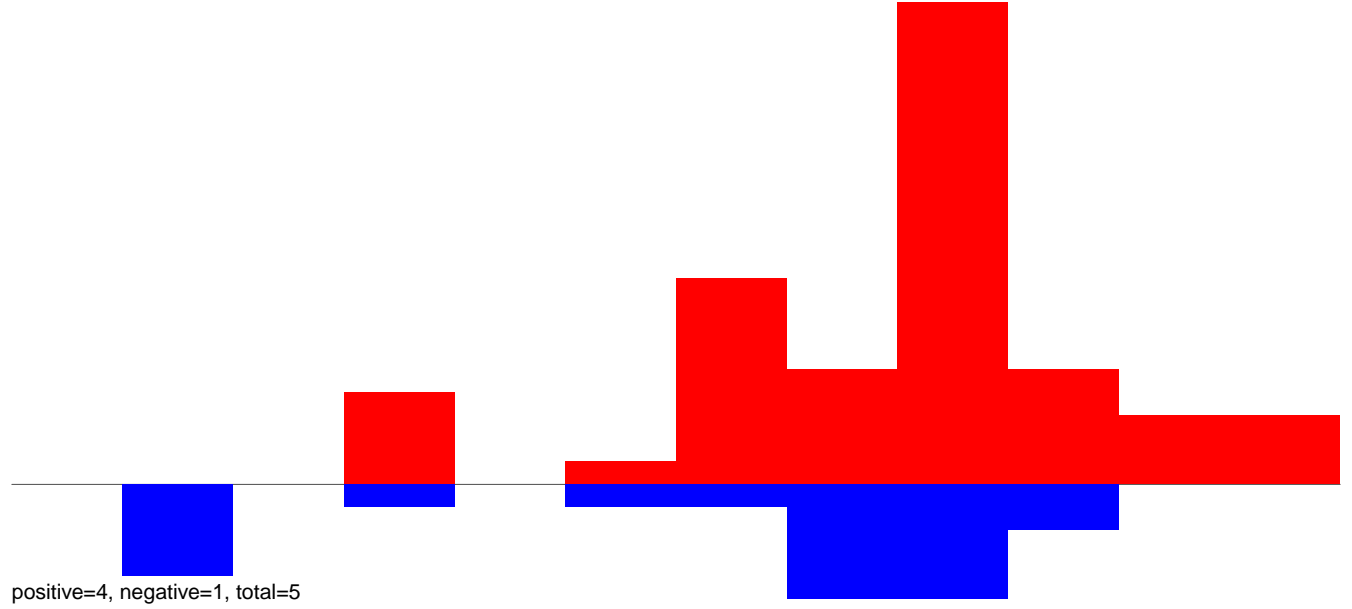
Window size=25, length=236, TE@rnd-3_family-677@Unknown@109:1-236

50 100 150 200 250

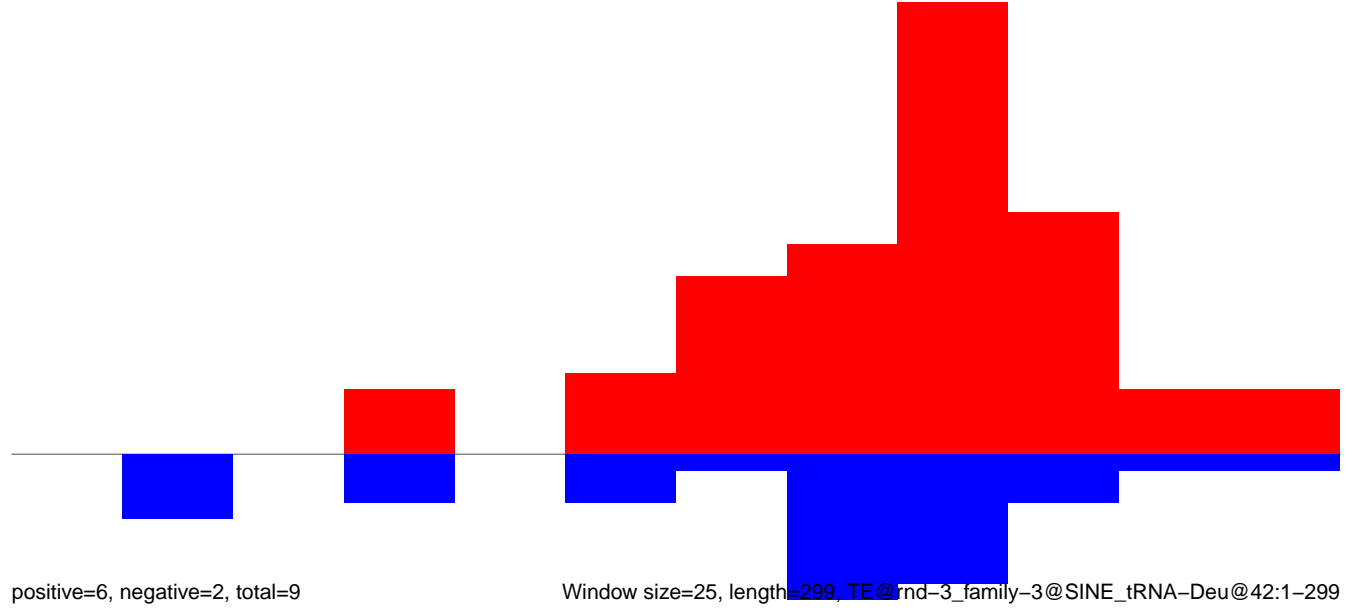
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

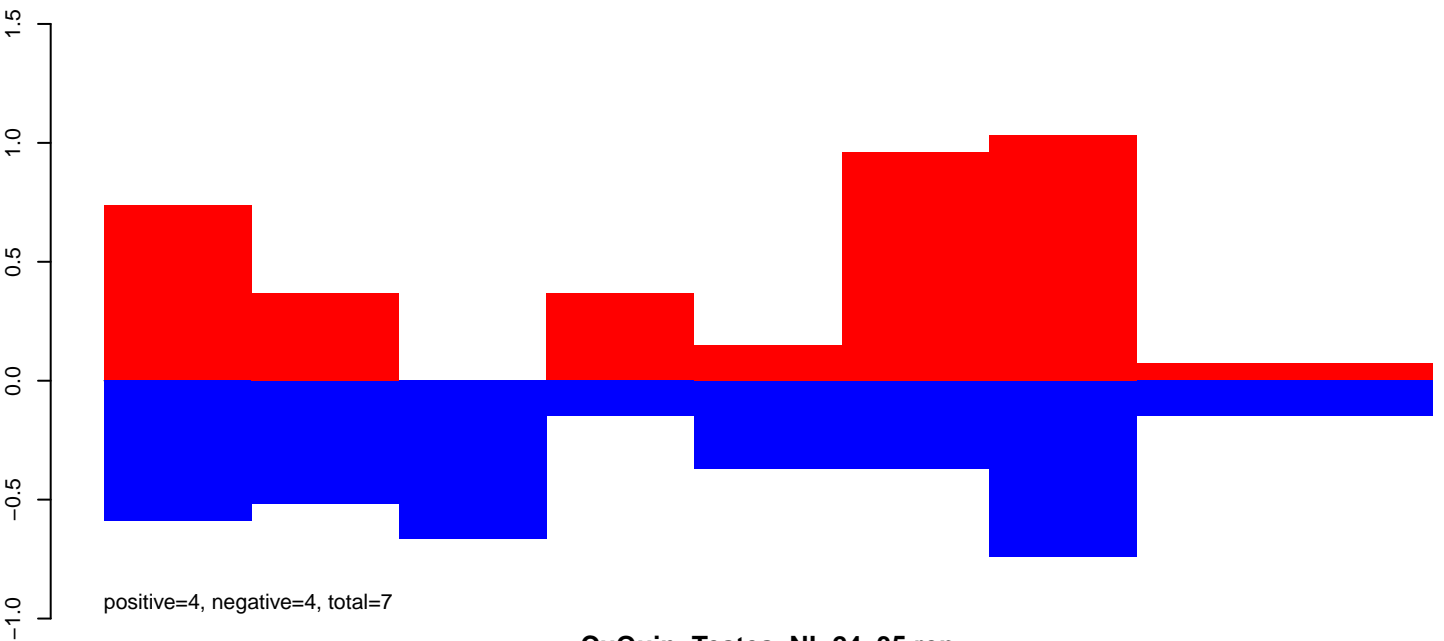


CuQuin_Testes_NL.rep

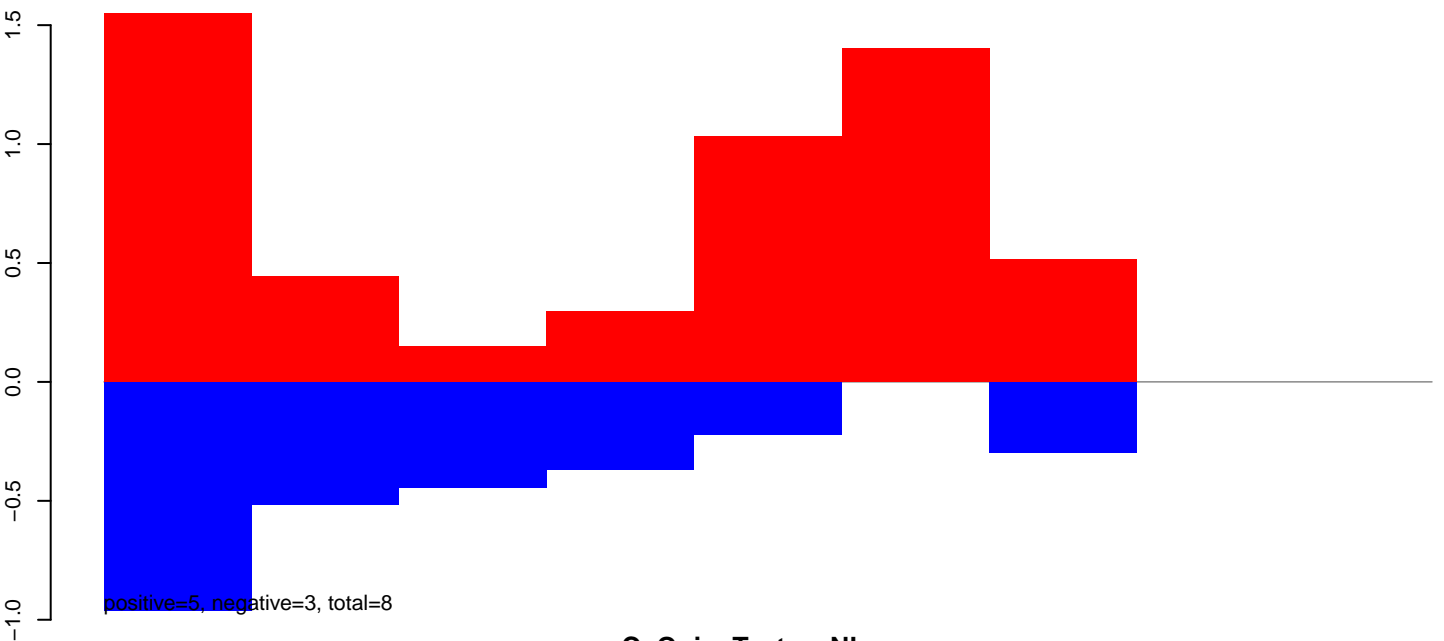


Window size=25, length=299, TE@rnd-3_family-3@SINE_tRNA-Deu@42:1-299

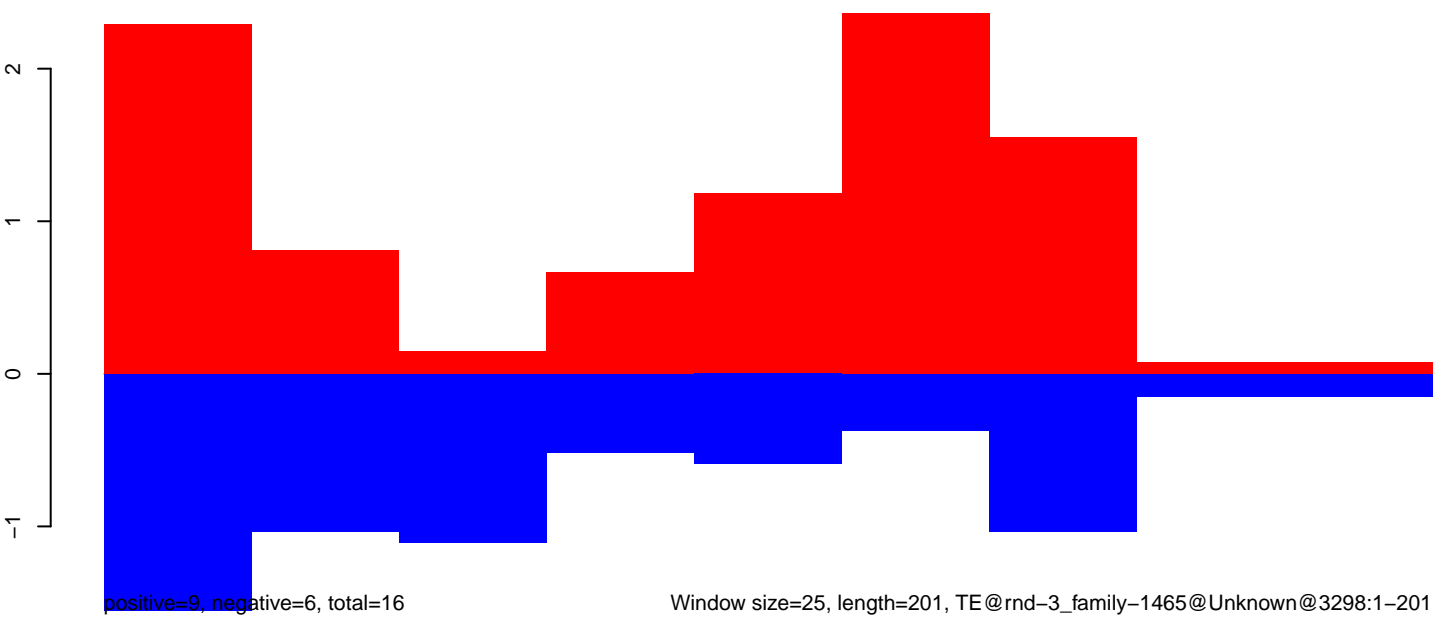
CuQuin_Testes_NL.18_23.rep



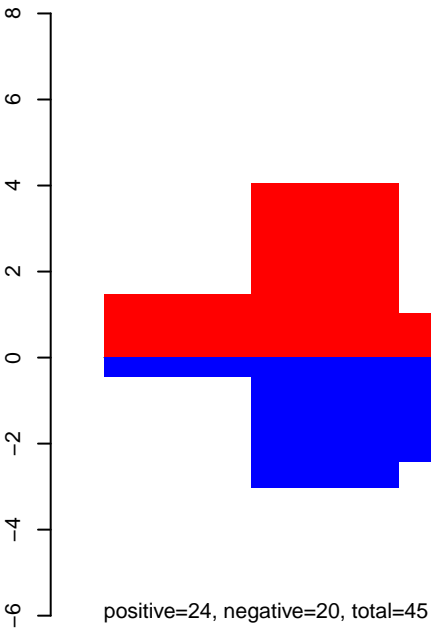
CuQuin_Testes_NL.24_35.rep



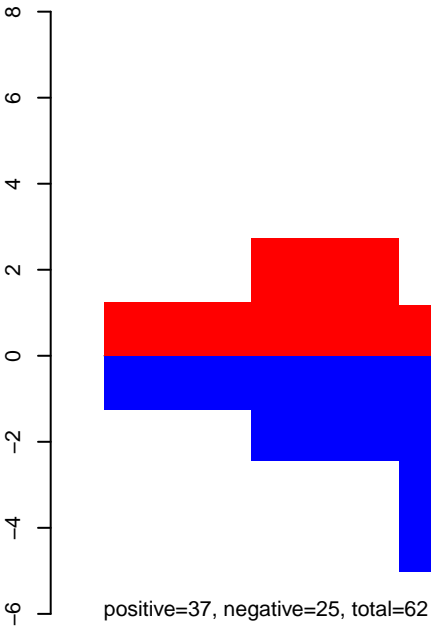
CuQuin_Testes_NL.rep



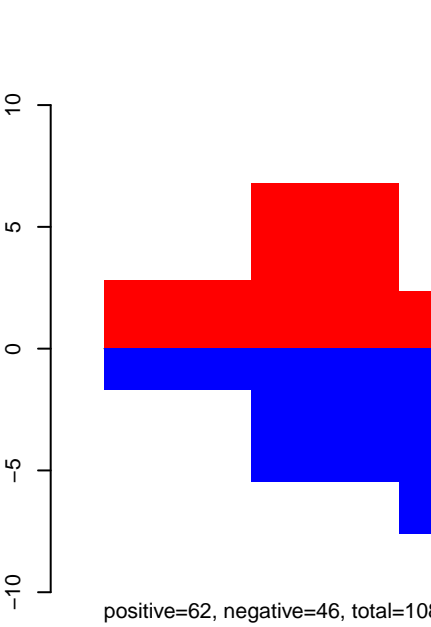
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



Window size=25, length=215, TE@rnd-2_family-554@SINE_tRNA-Deu@1915:1-215

50

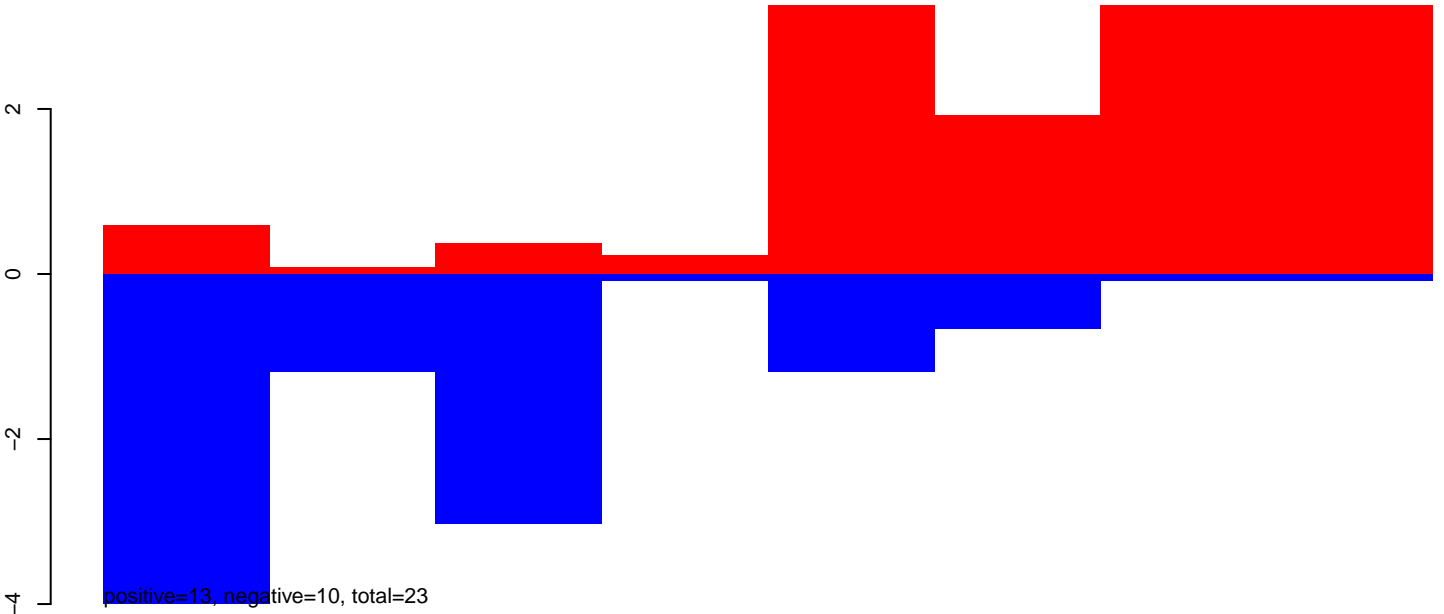
100

150

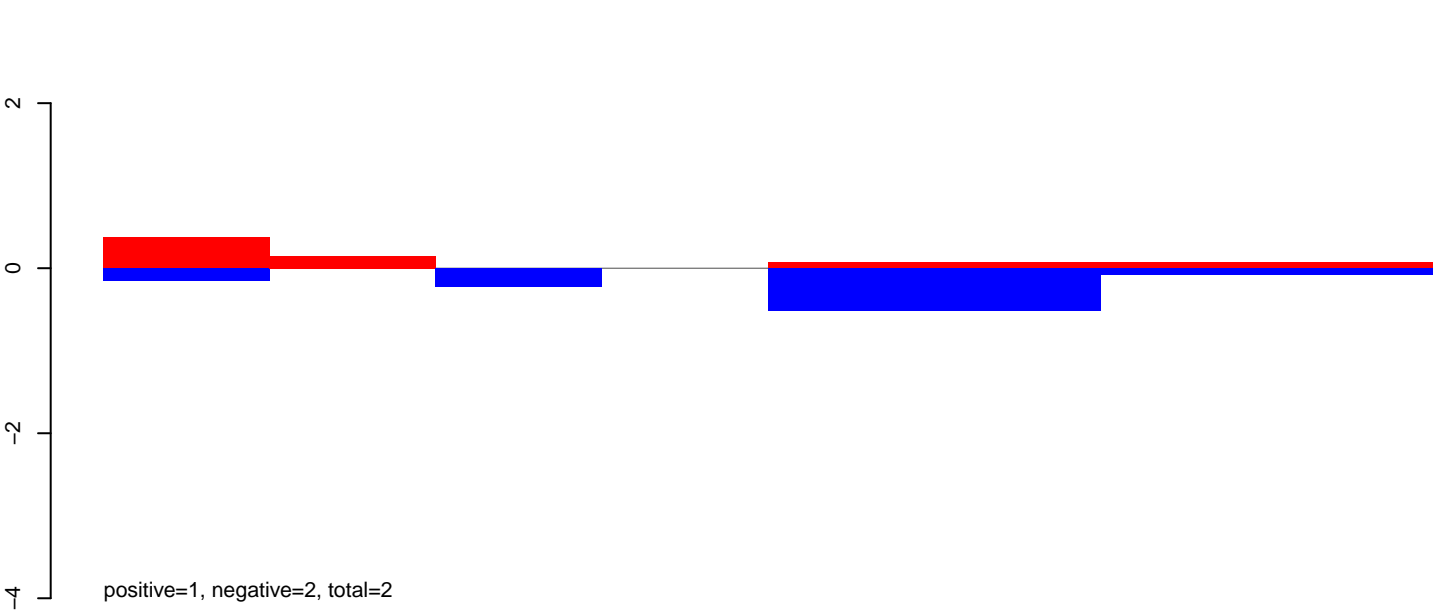
200

250

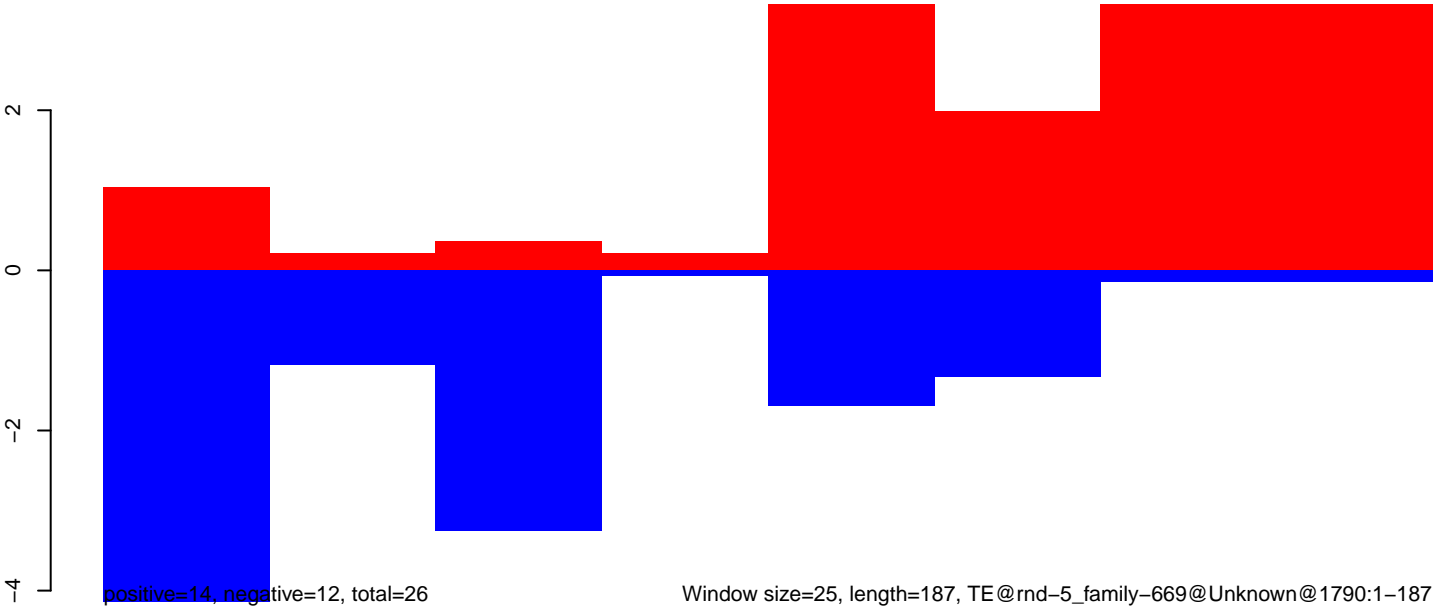
CuQuin_Testes_NL.18_23.rep



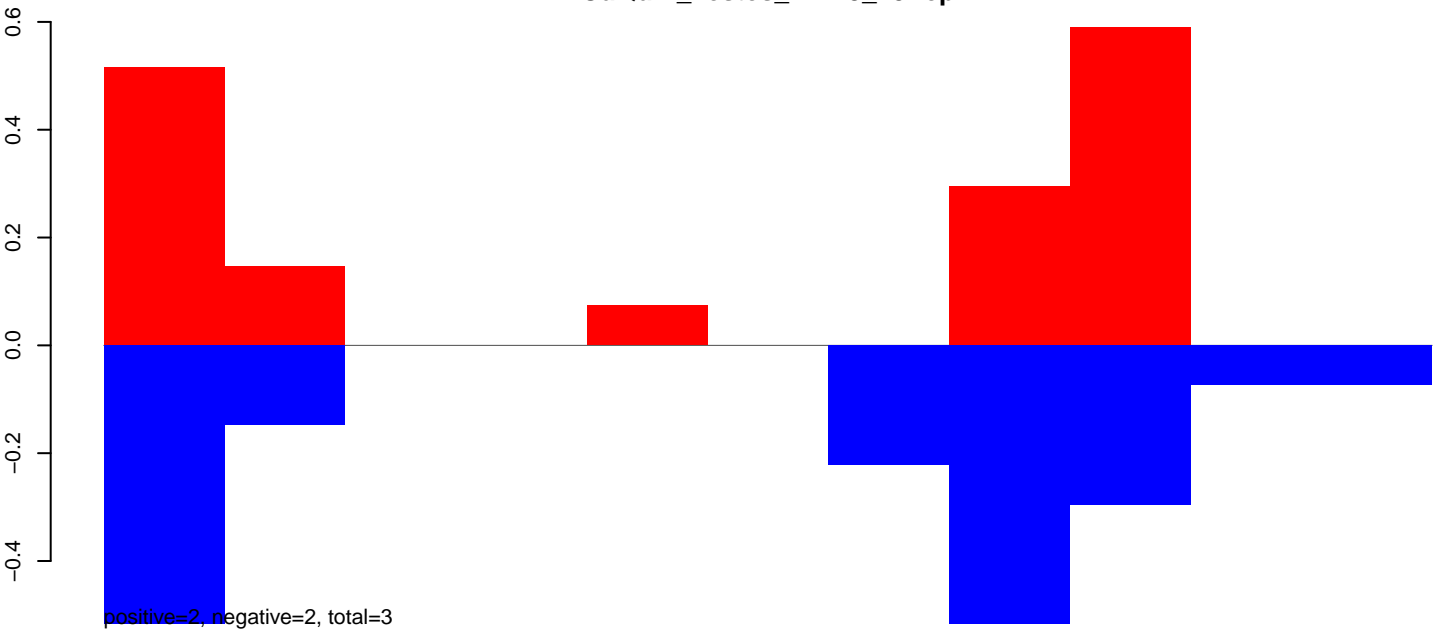
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



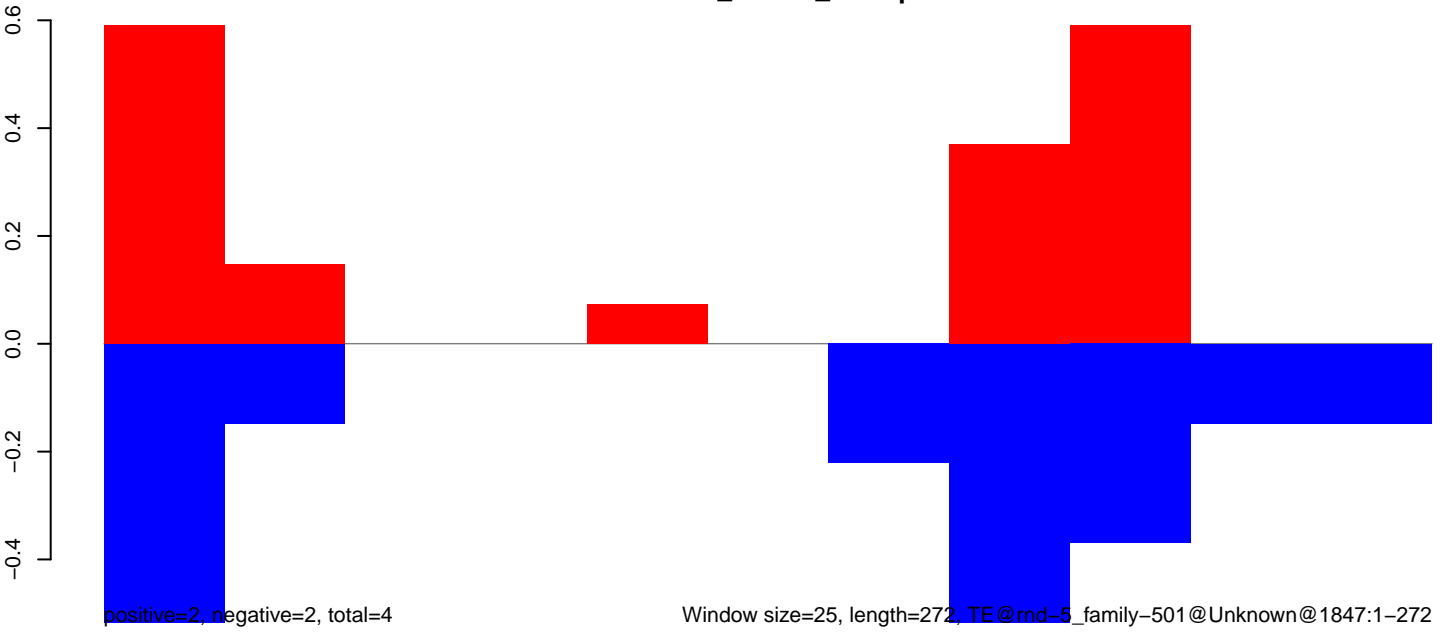
CuQuin_Testes_NL.18_23.rep



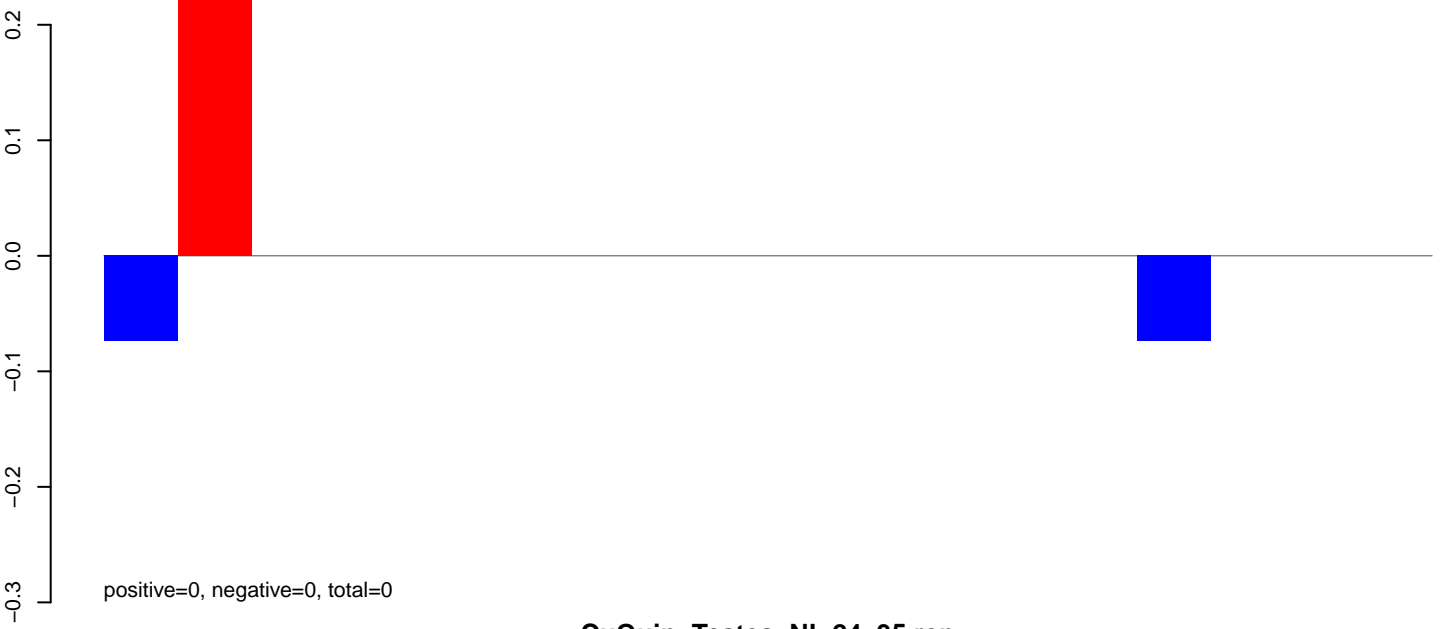
CuQuin_Testes_NL.24_35.rep



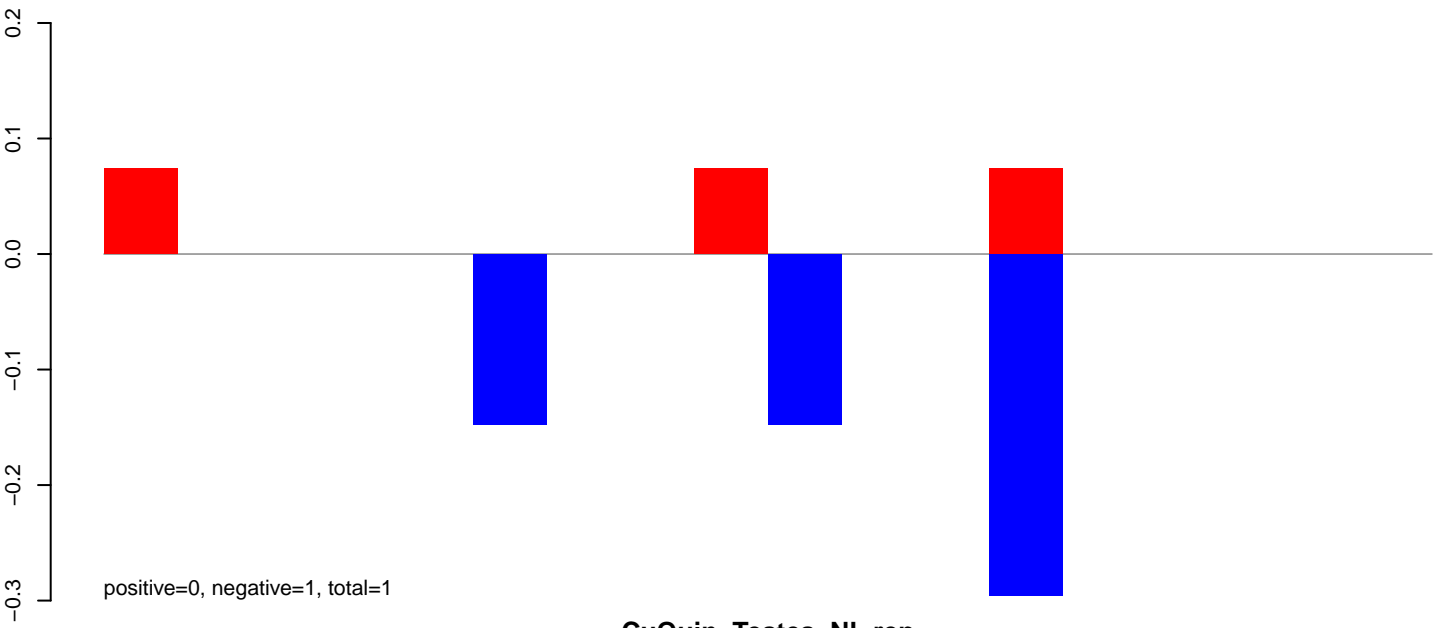
CuQuin_Testes_NL.rep



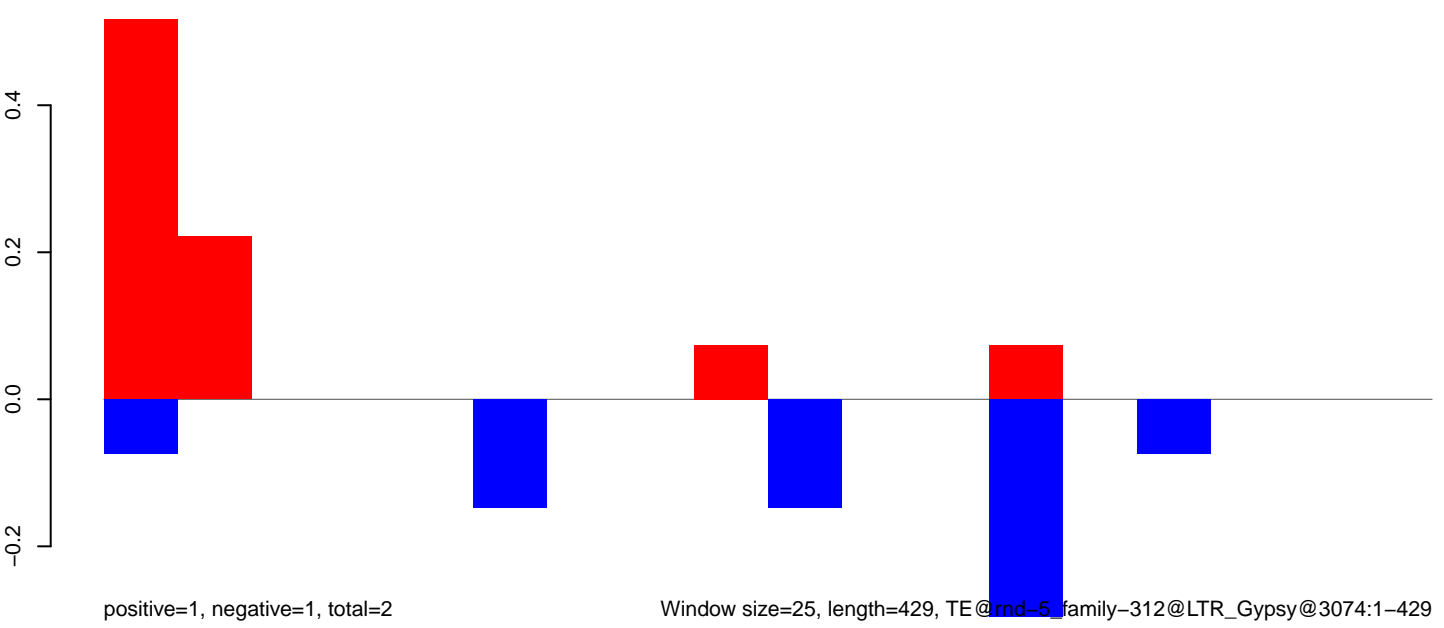
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

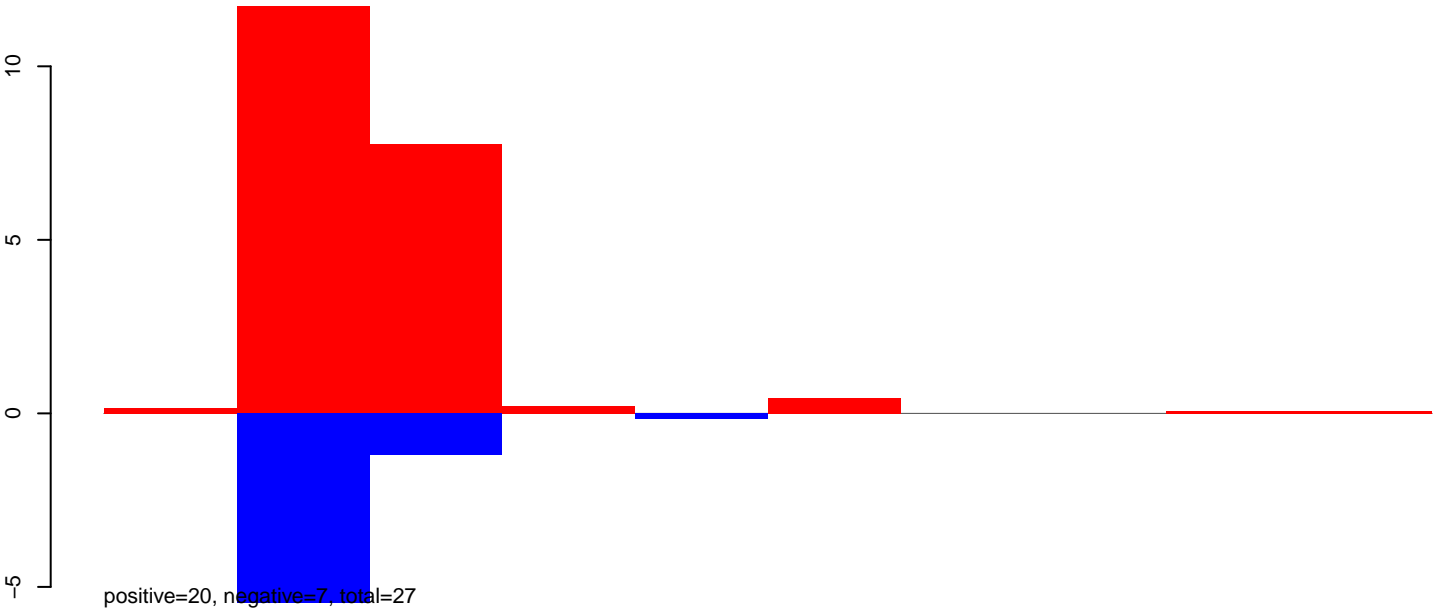


CuQuin_Testes_NL.rep

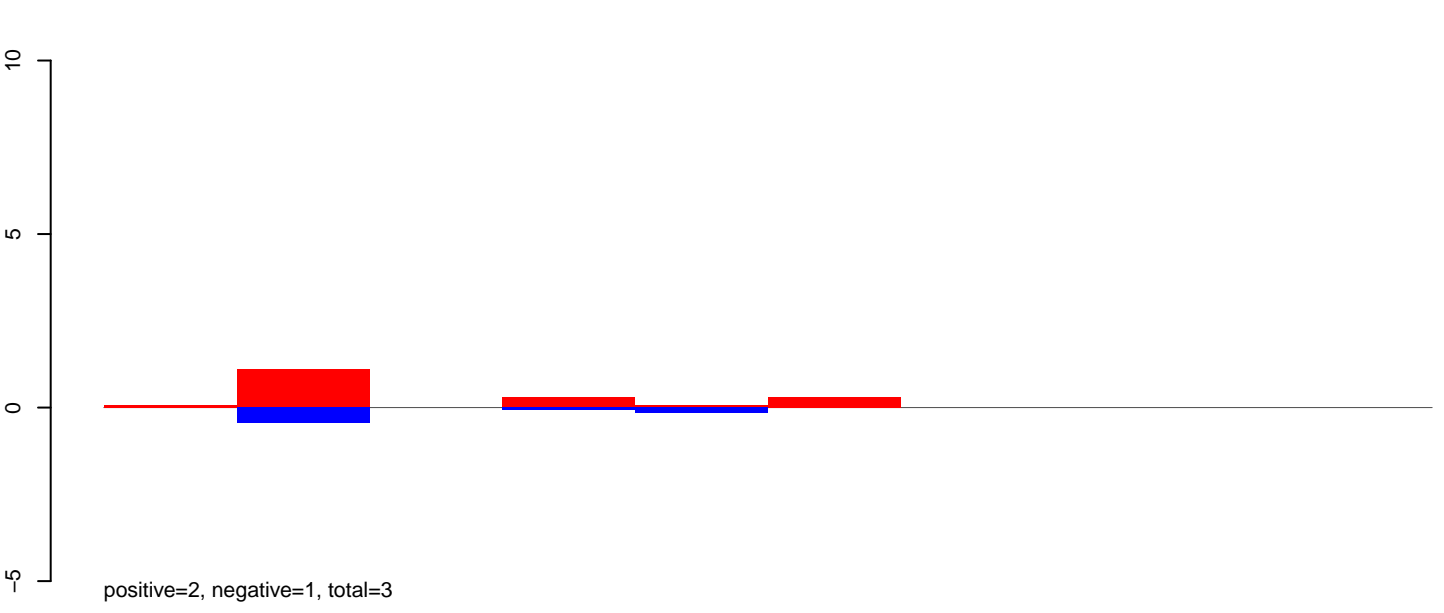


Window size=25, length=429, TE@rnd-5, family-312@LTR_Gypsy@3074:1-429

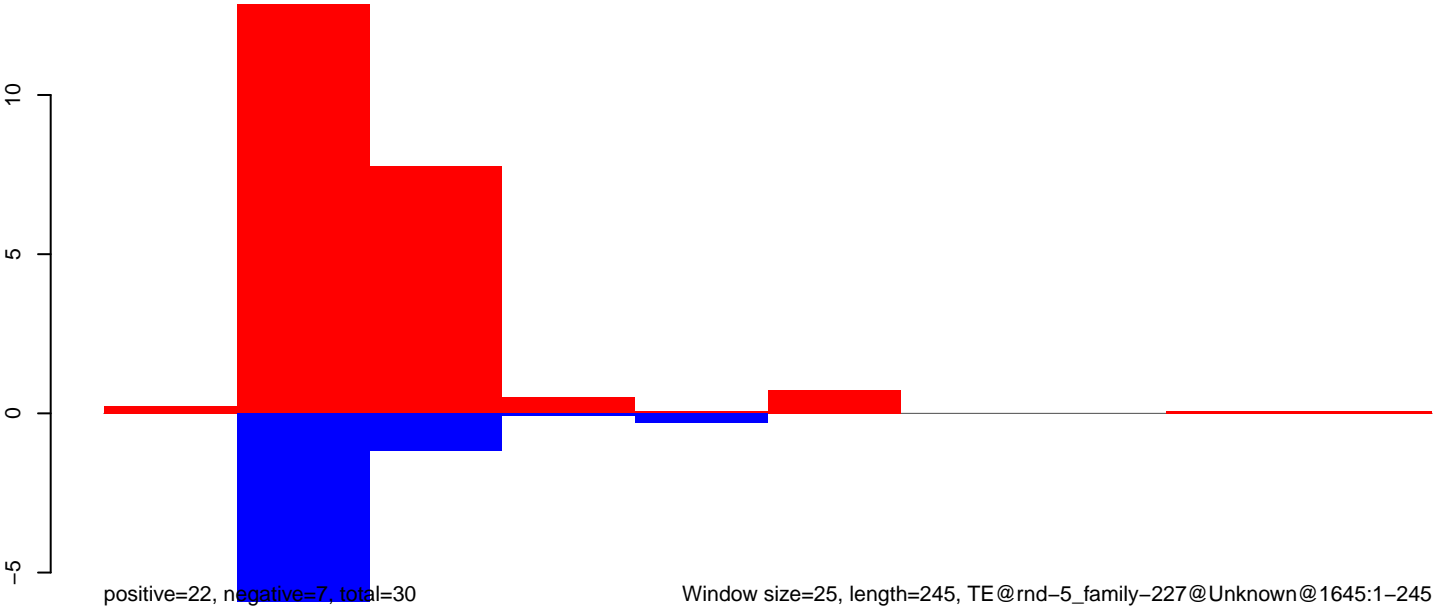
CuQuin_Testes_NL.18_23.rep



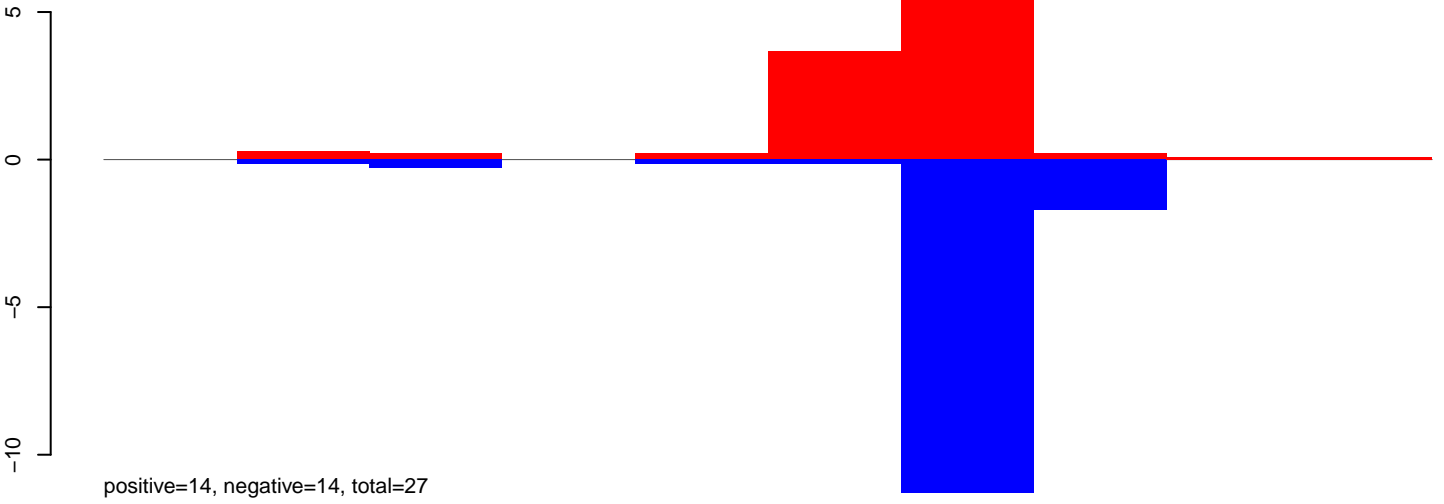
CuQuin_Testes_NL.24_35.rep



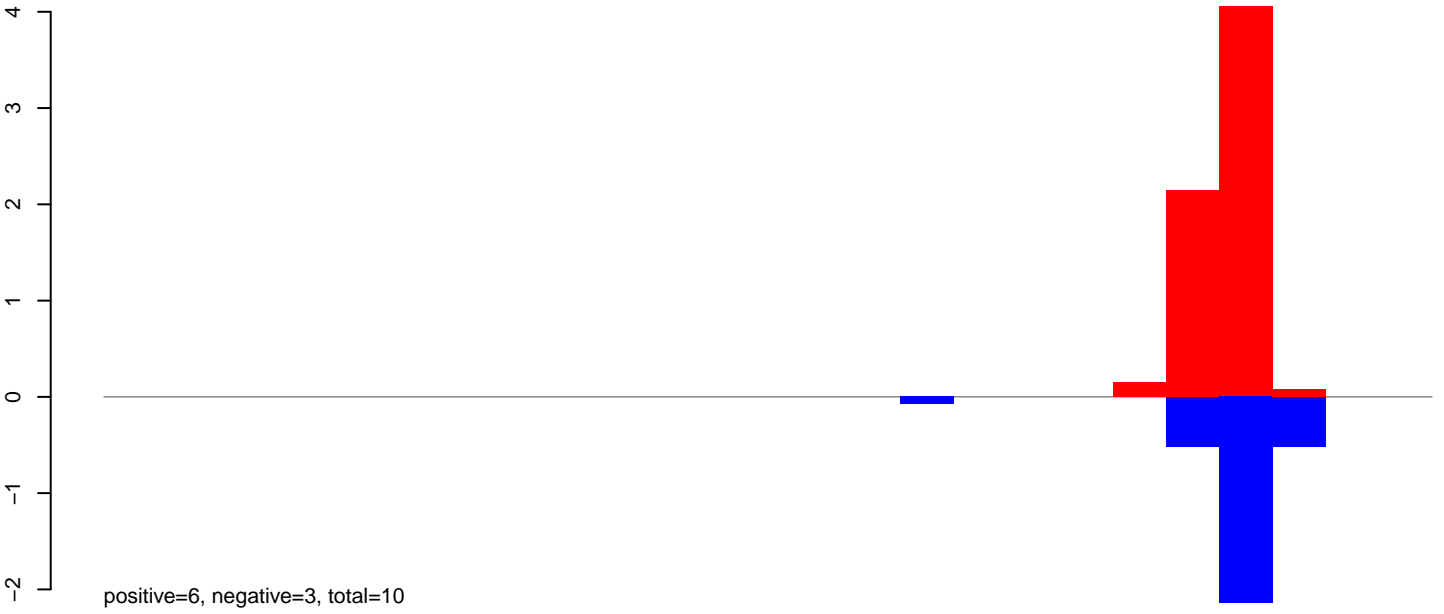
CuQuin_Testes_NL.rep



CuQuin_Testes_NL.18_23.rep



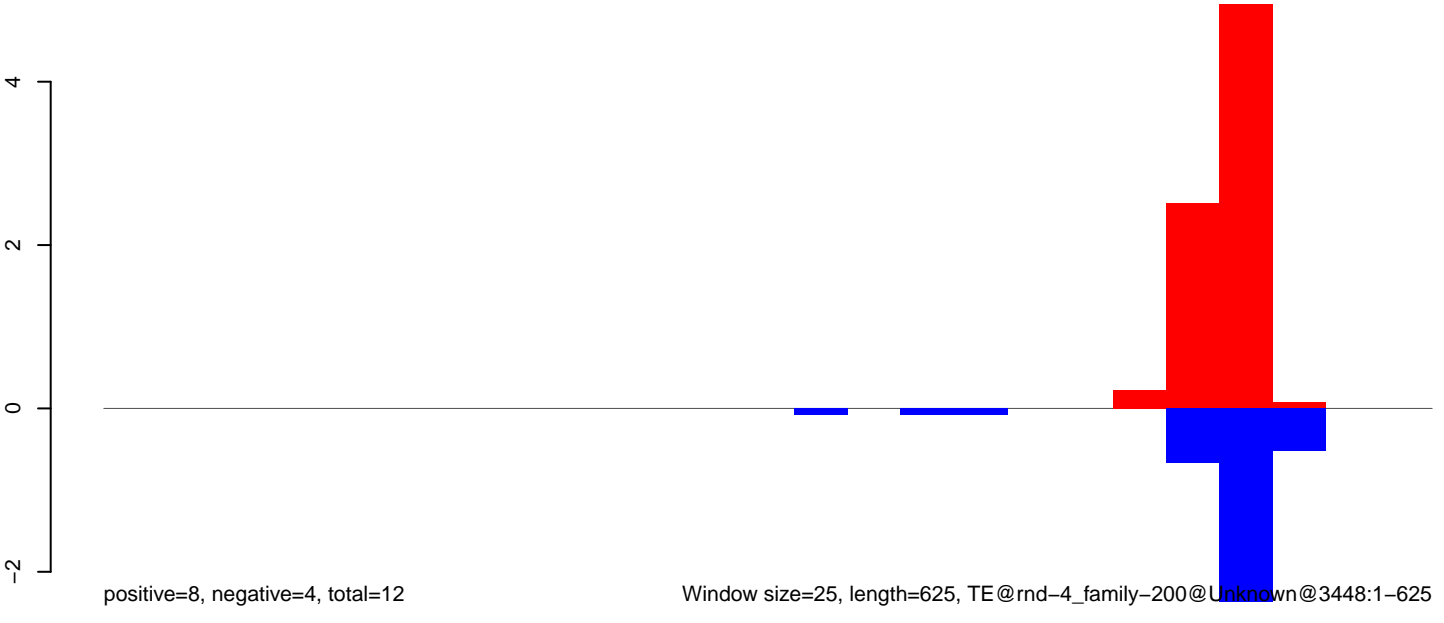
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



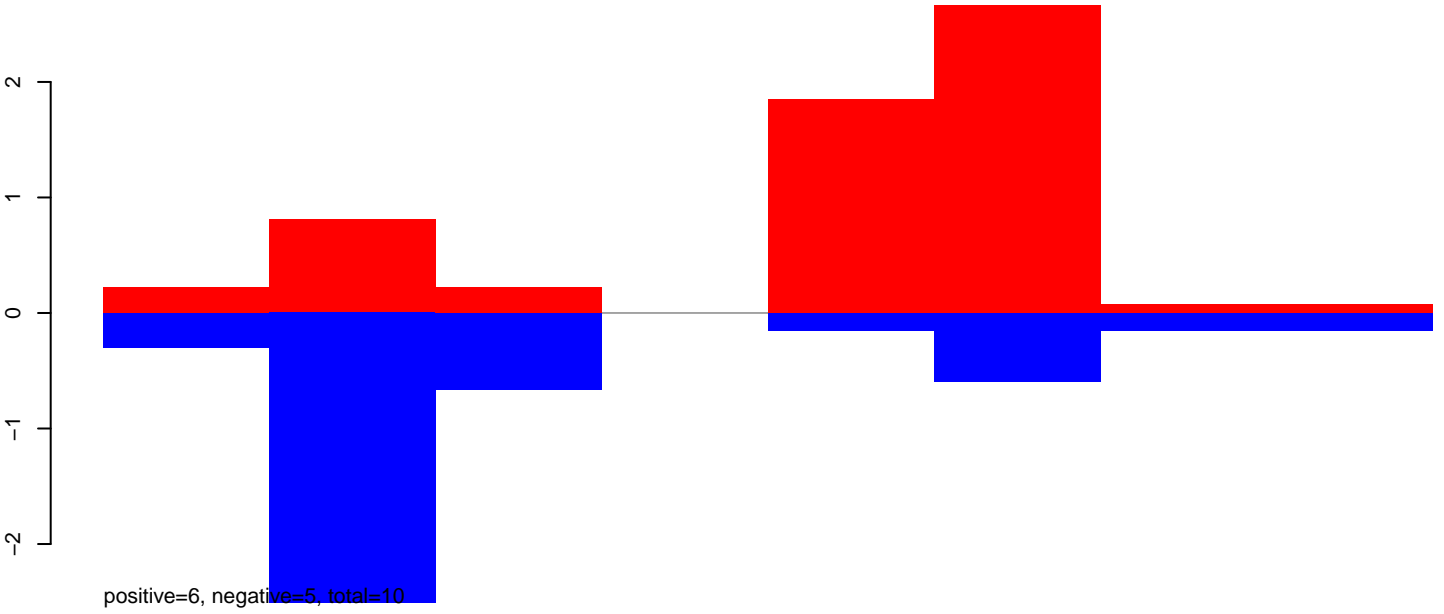
CuQuin_Testes_NL.rep



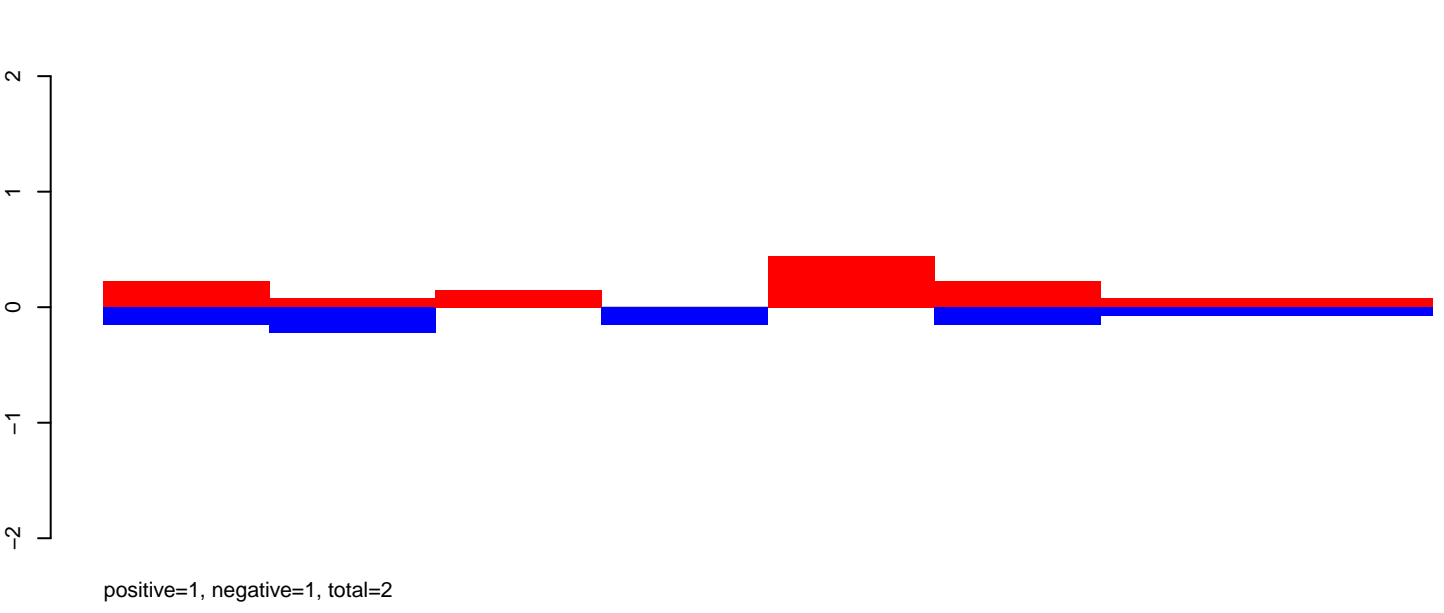
Window size=25, length=625, TE@rnd-4_family-200@Unknown@3448:1-625

100 200 300 400 500 600

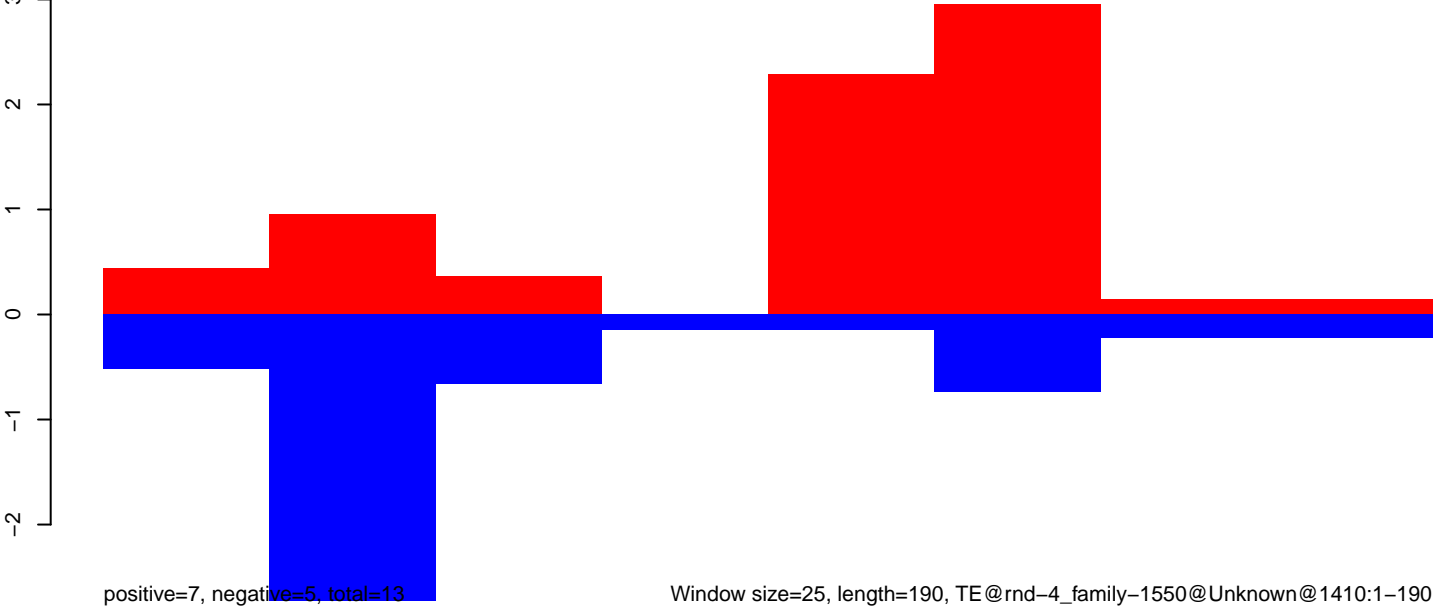
CuQuin_Testes_NL.18_23.rep



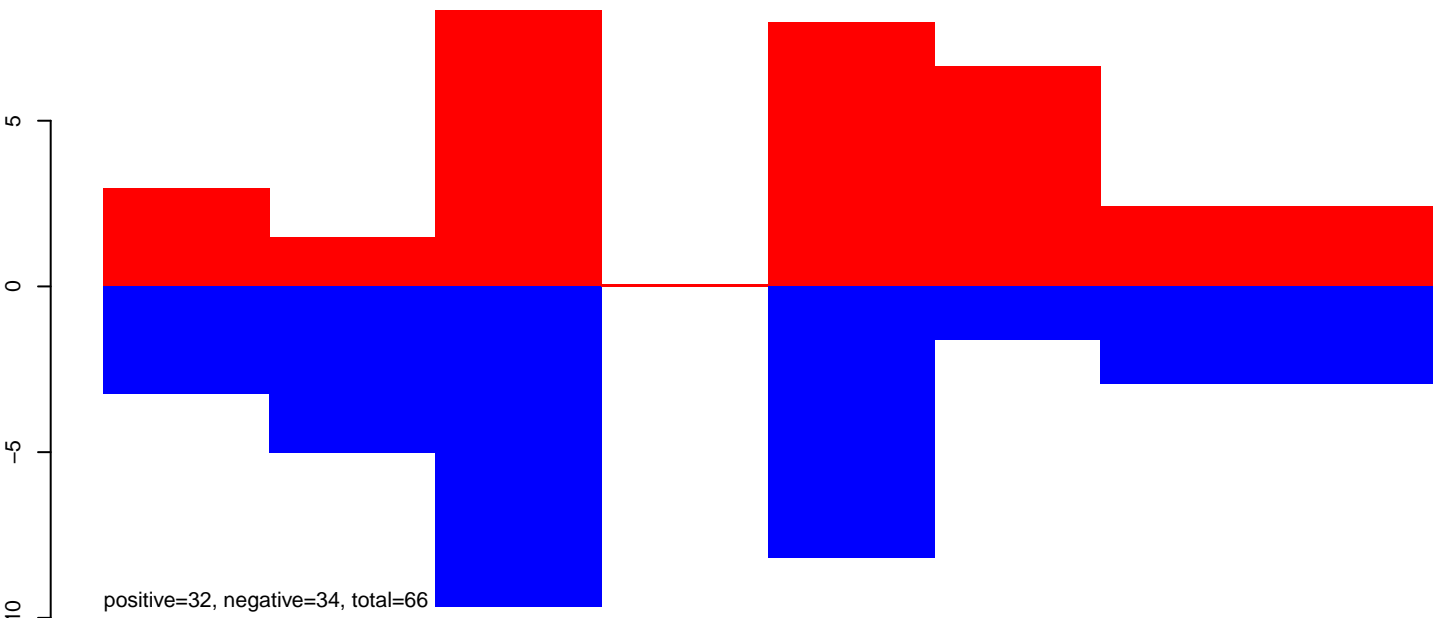
CuQuin_Testes_NL.24_35.rep



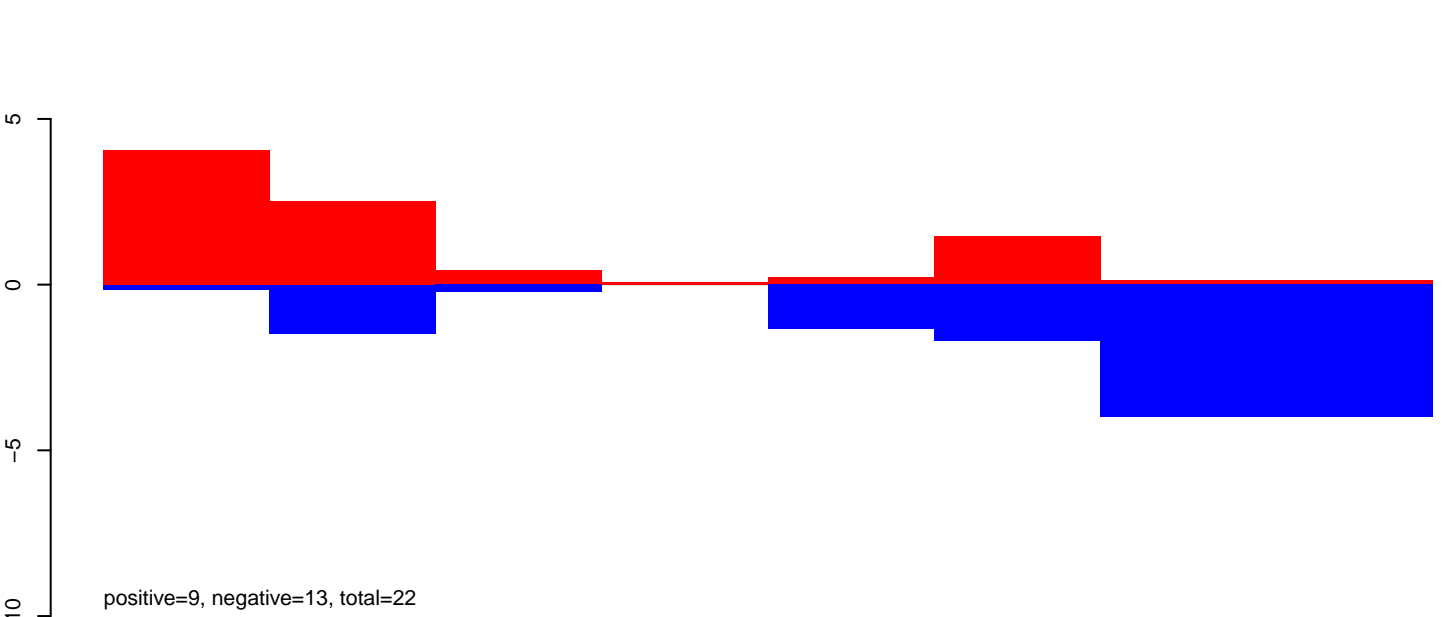
CuQuin_Testes_NL.rep



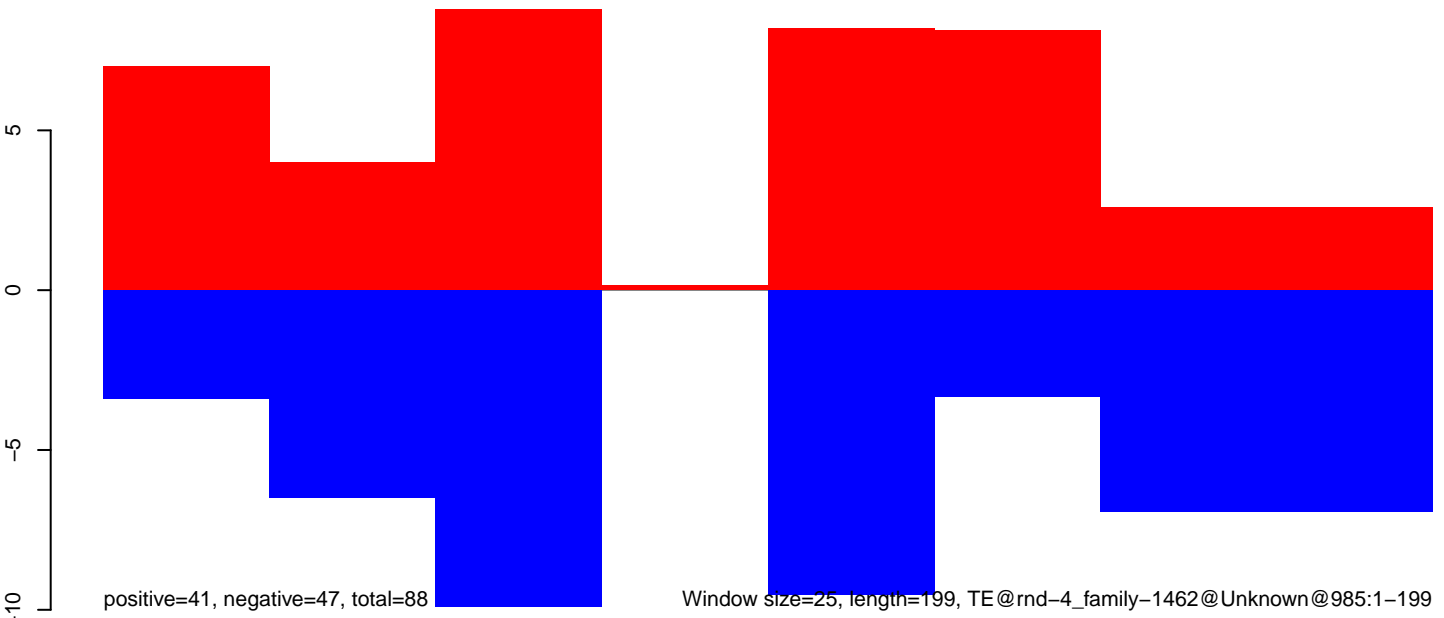
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

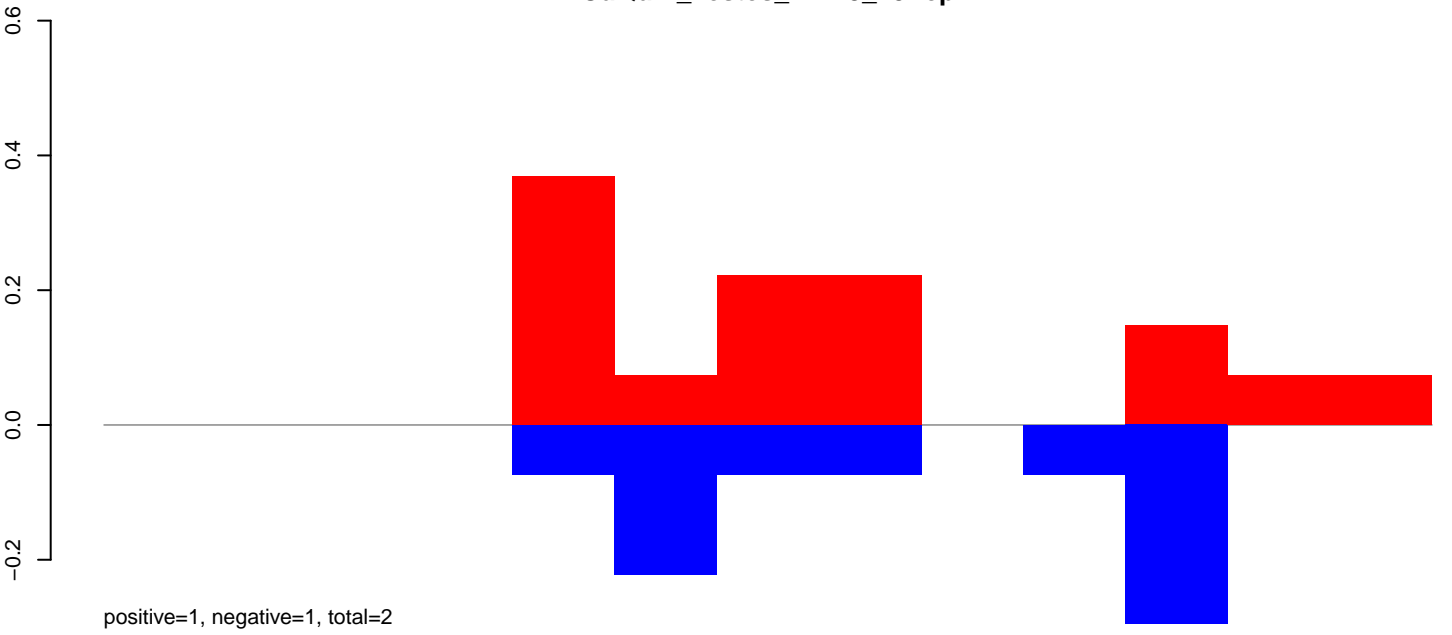


CuQuin_Testes_NL.rep

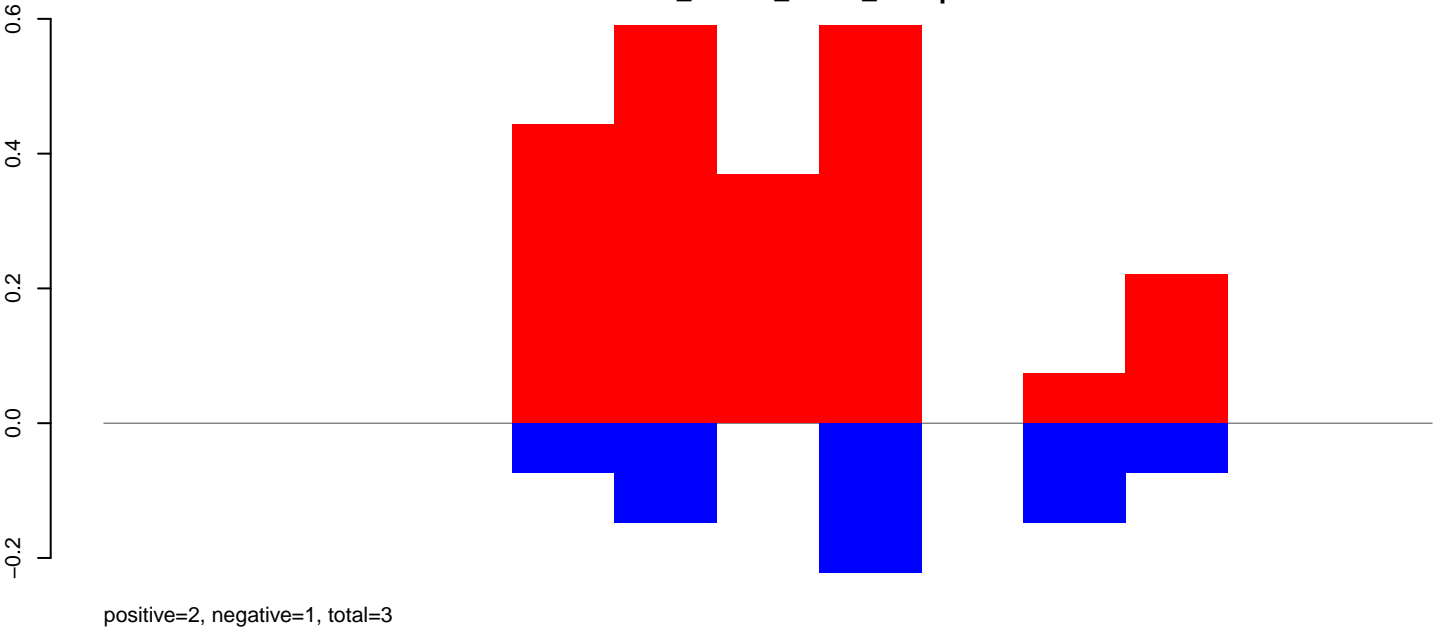


Window size=25, length=199, TE@rnd-4_family-1462@Unknown@985:1-199

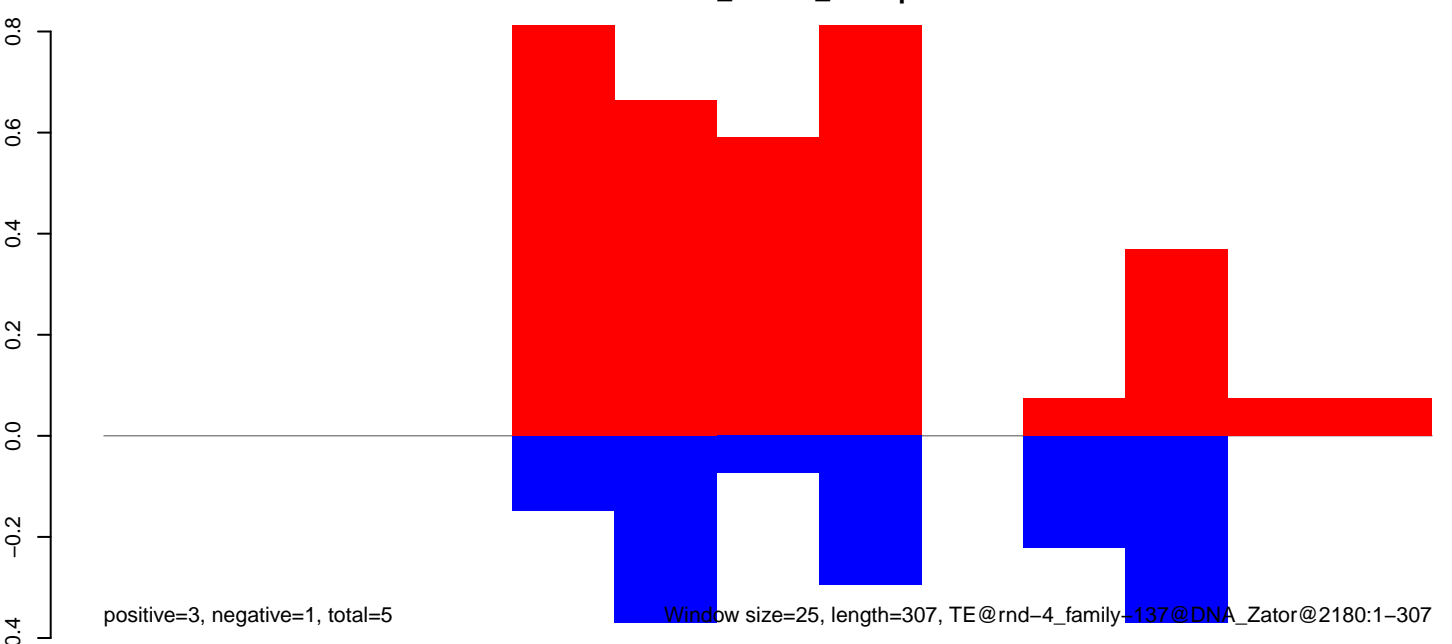
CuQuin_Testes_NL.18_23.rep



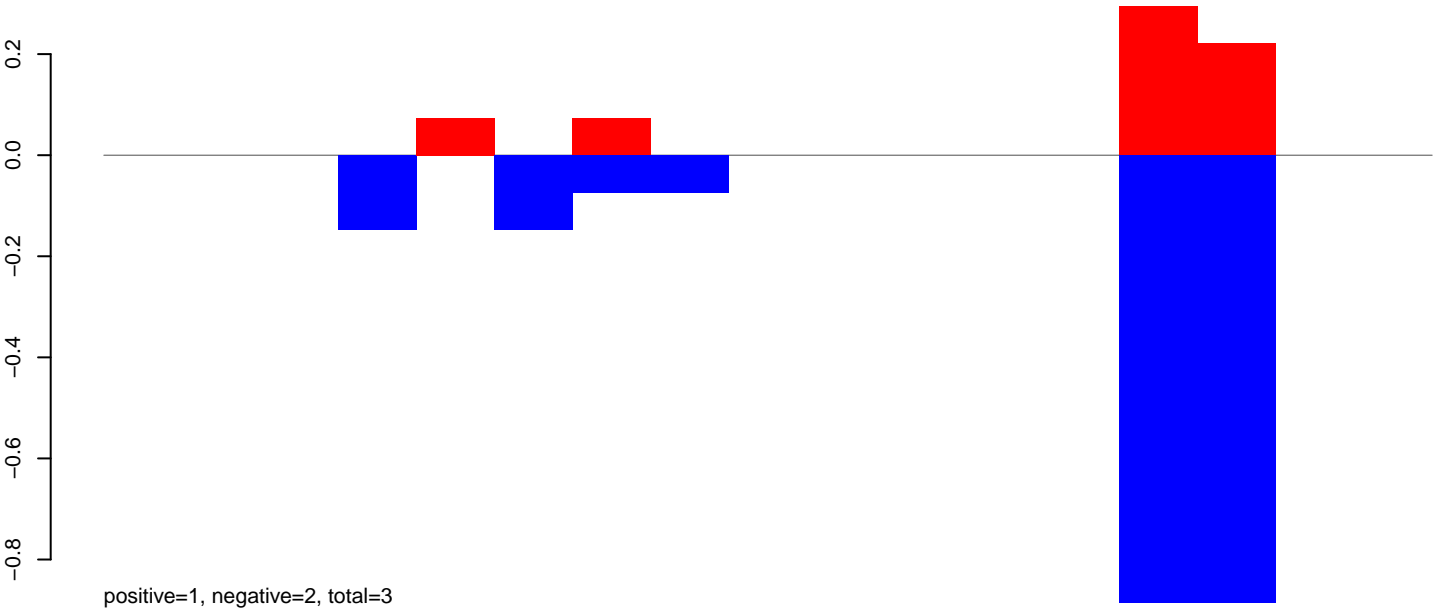
CuQuin_Testes_NL.24_35.rep



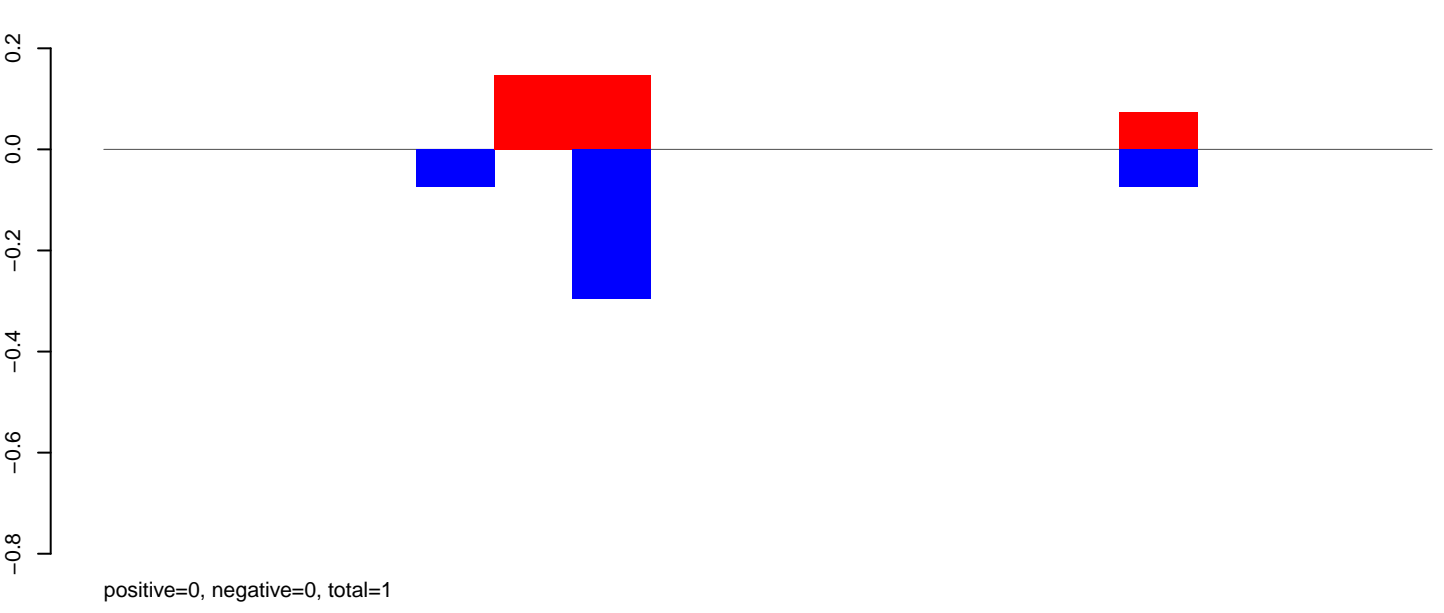
CuQuin_Testes_NL.rep



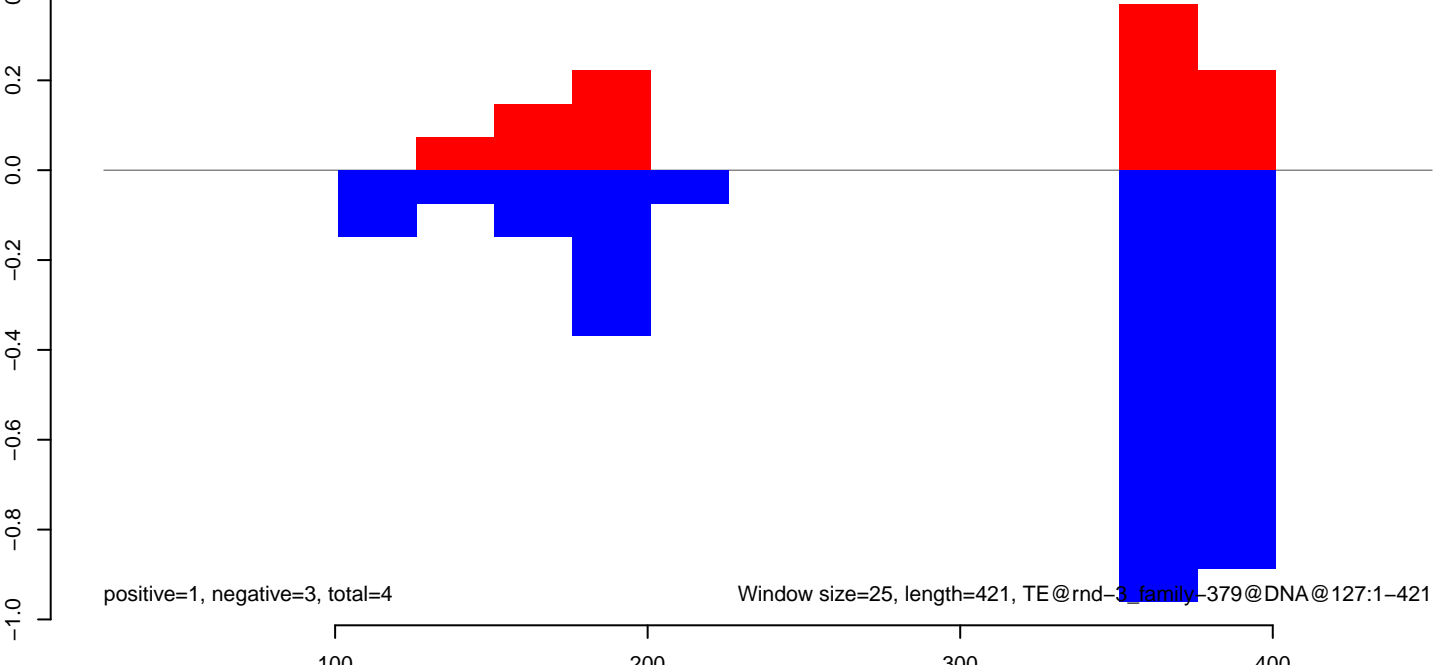
CuQuin_Testes_NL.18_23.rep



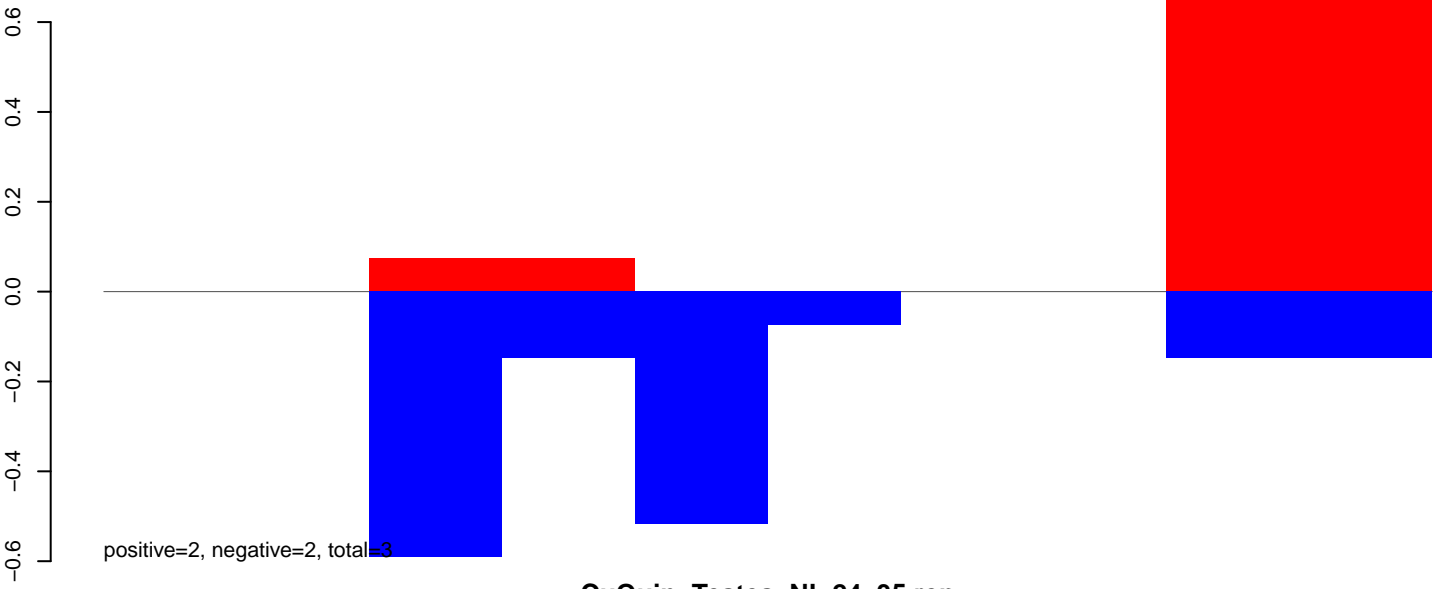
CuQuin_Testes_NL.24_35.rep



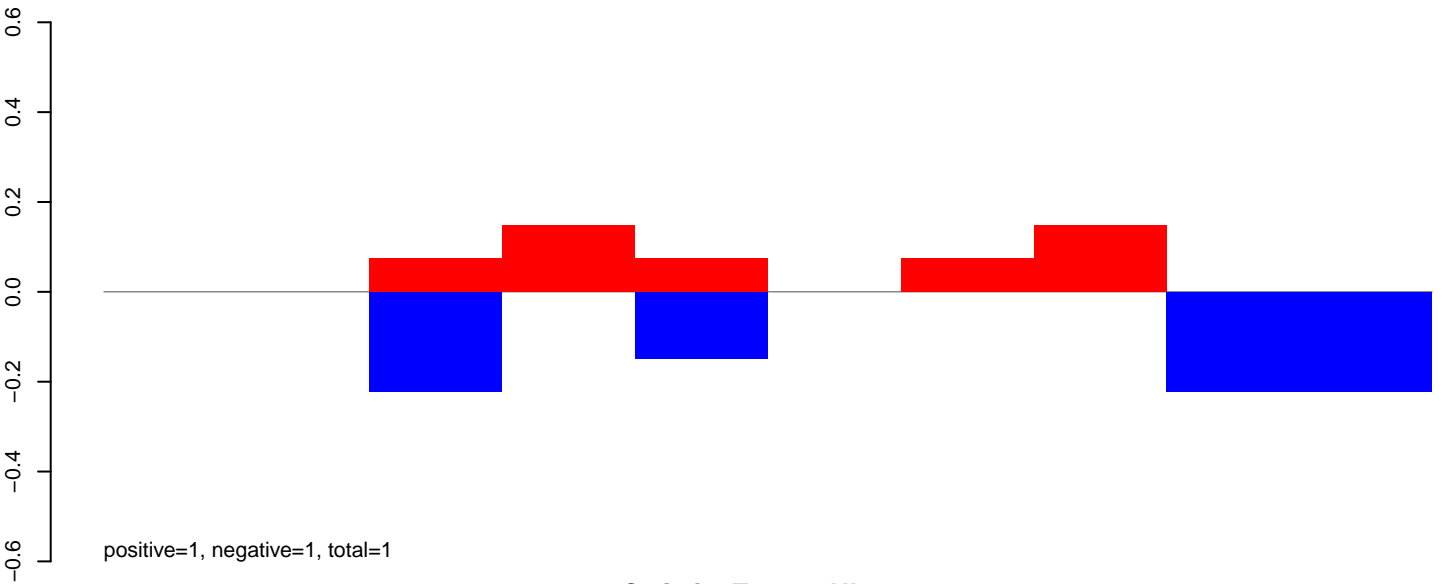
CuQuin_Testes_NL.rep



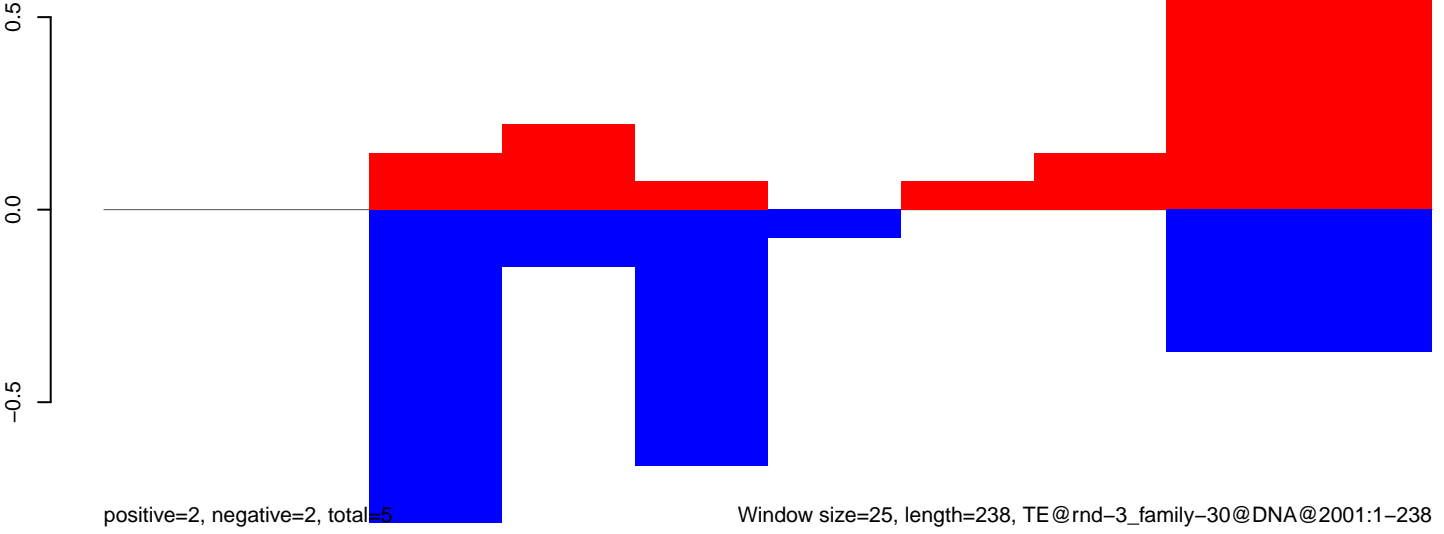
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



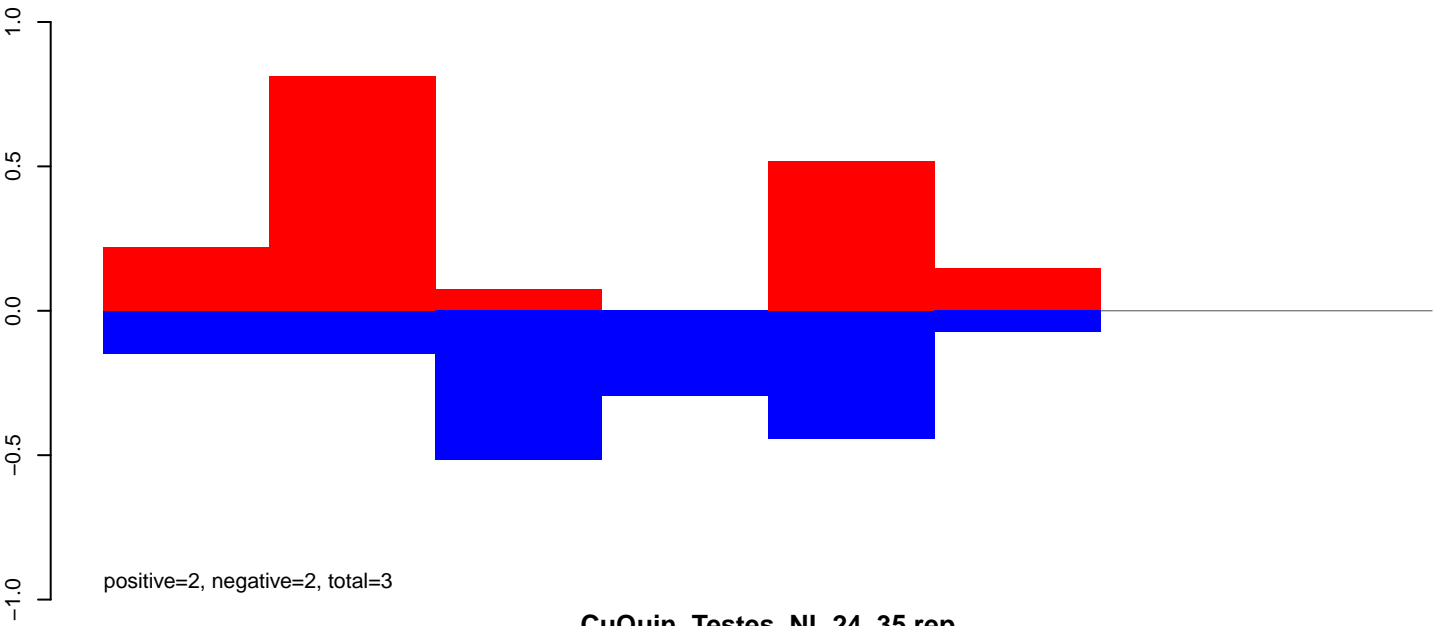
CuQuin_Testes_NL.rep



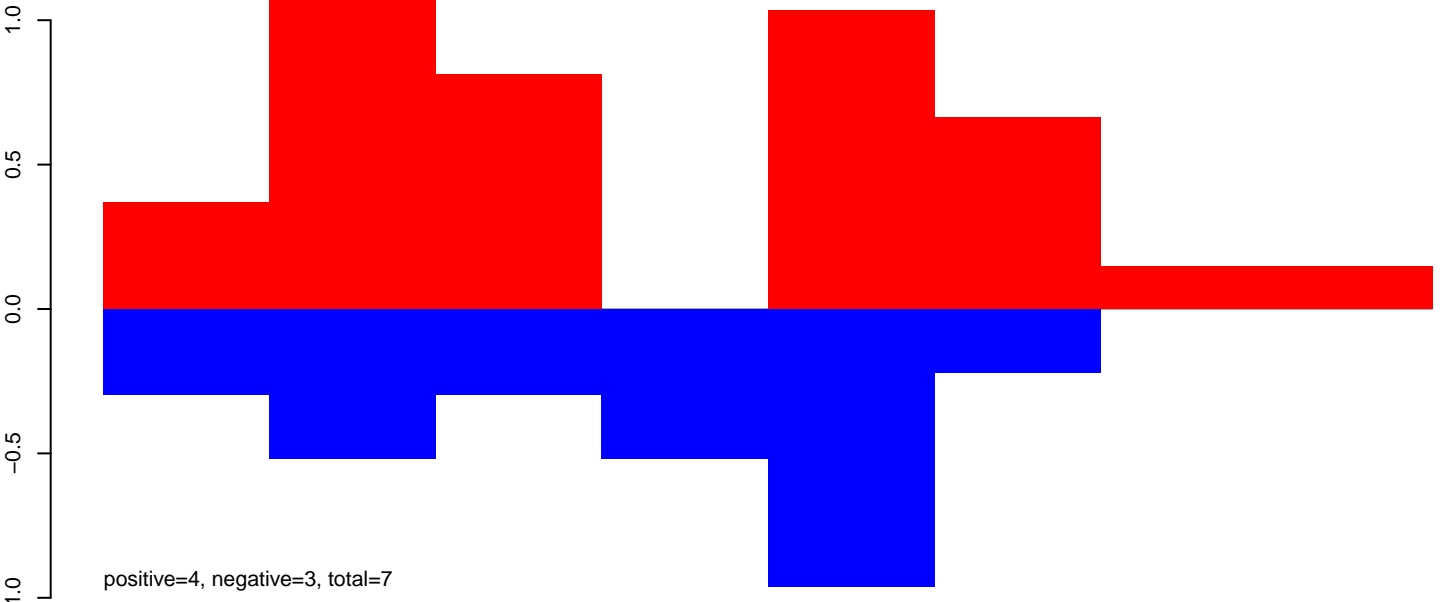
Window size=25, length=238, TE@rnd-3_family-30@DNA@2001:1-238

50 100 150 200 250

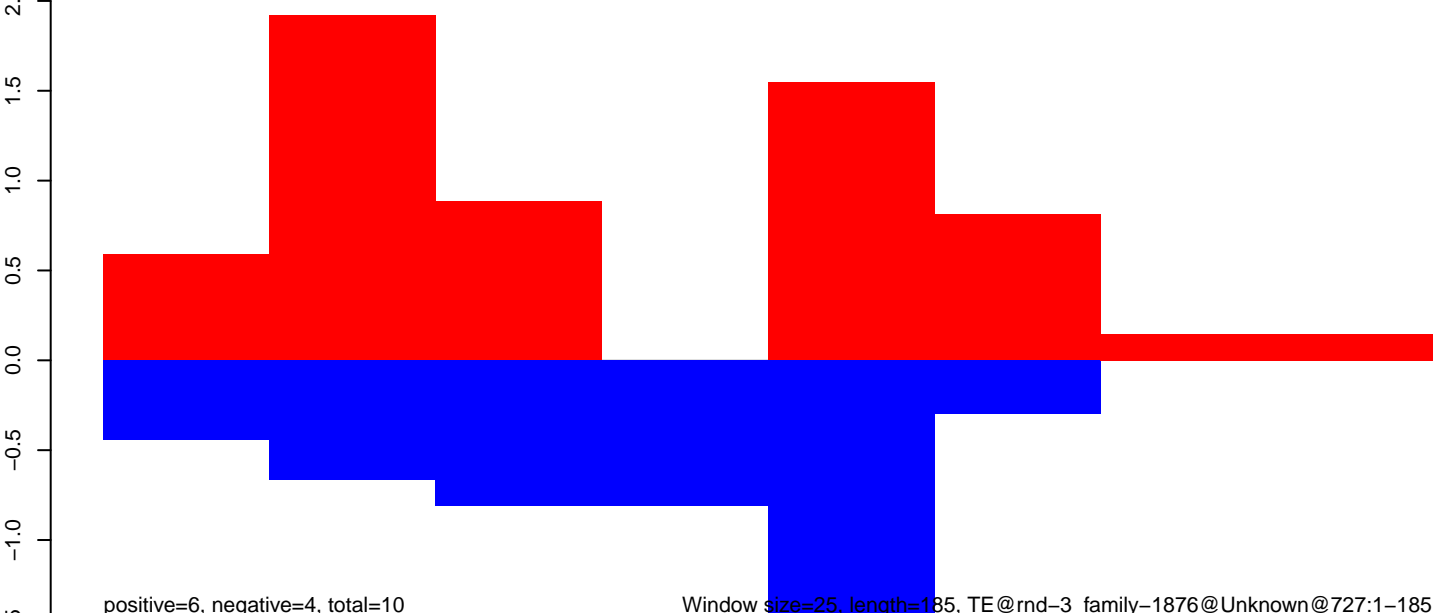
CuQuin_Testes_NL.18_23.rep



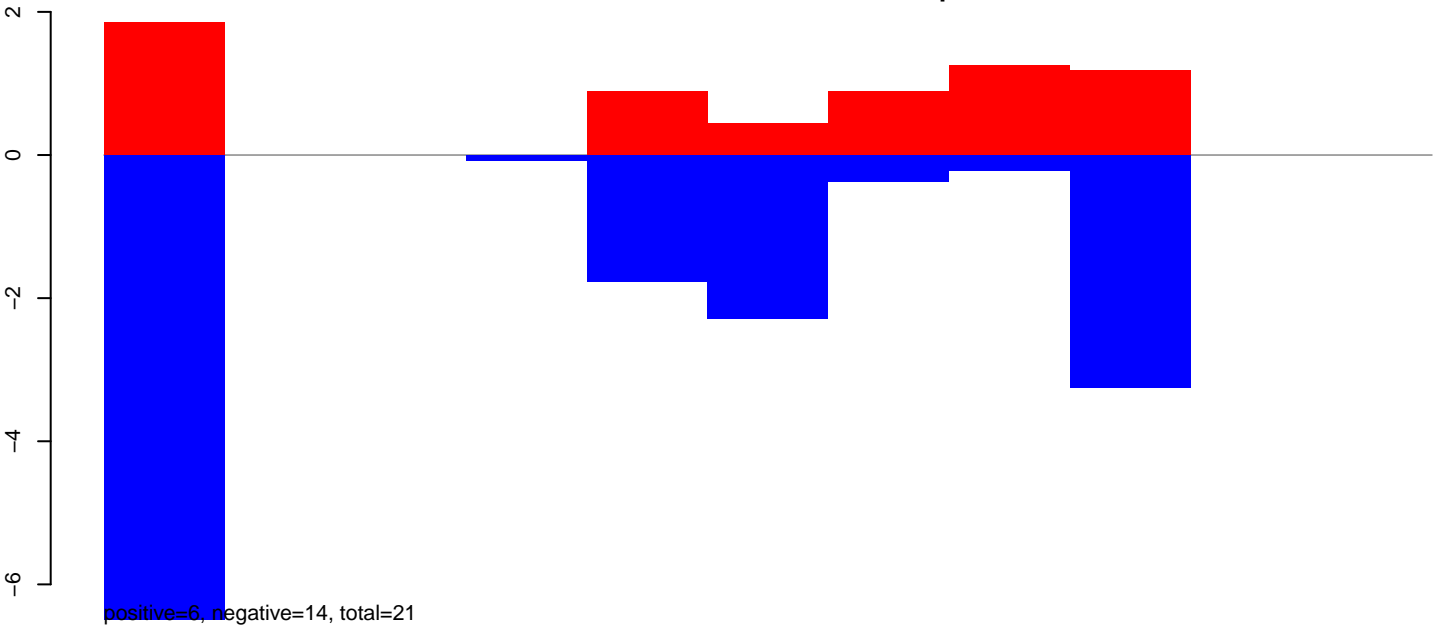
CuQuin_Testes_NL.24_35.rep



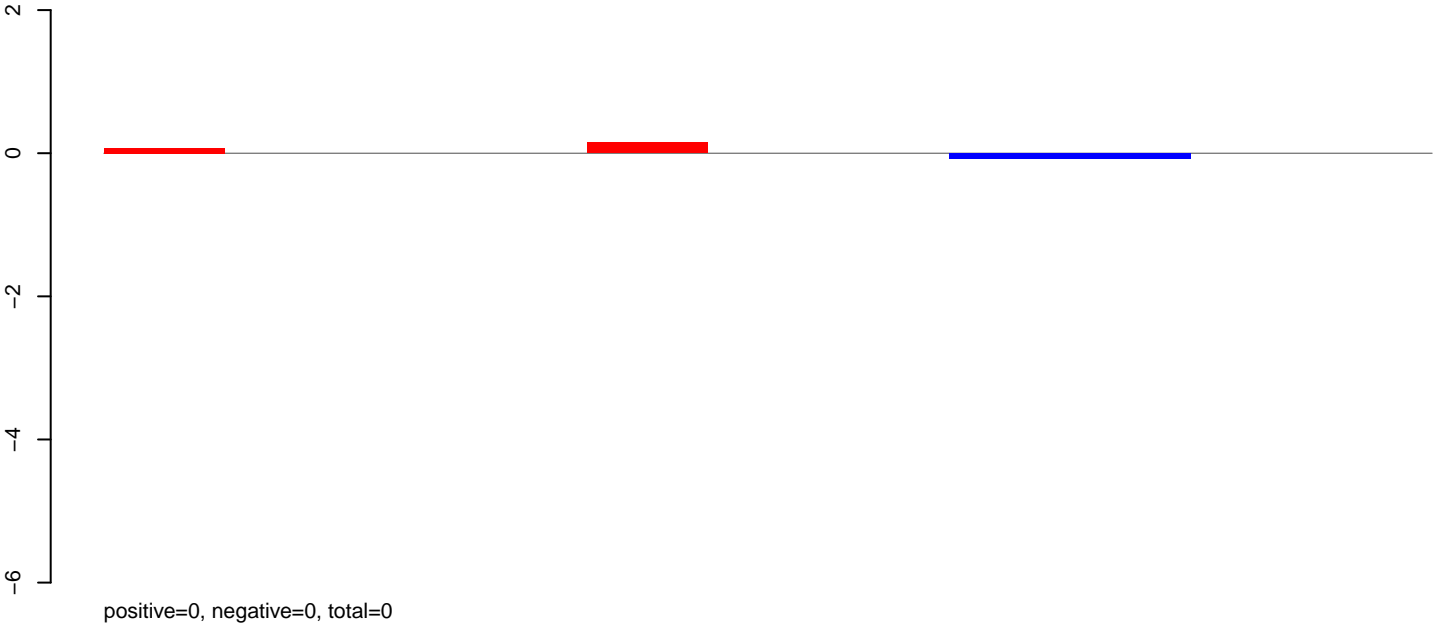
CuQuin_Testes_NL.rep



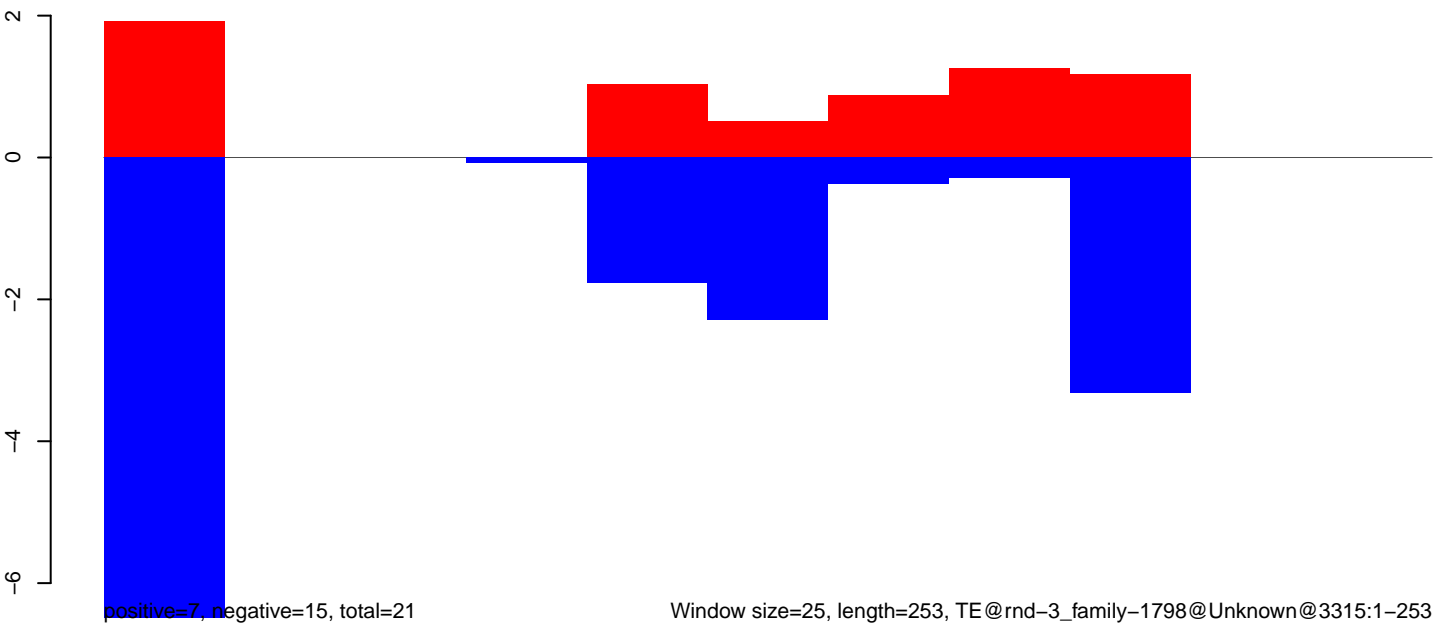
CuQuin_Testes_NL.18_23.rep



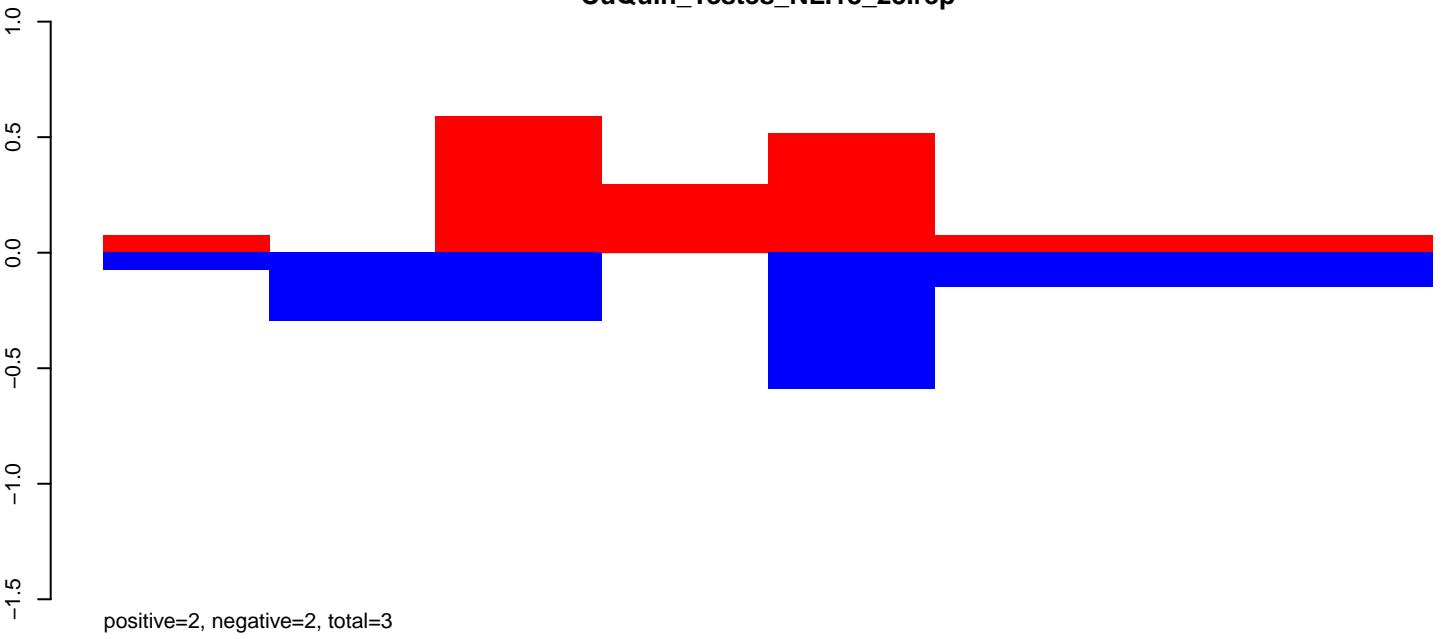
CuQuin_Testes_NL.24_35.rep



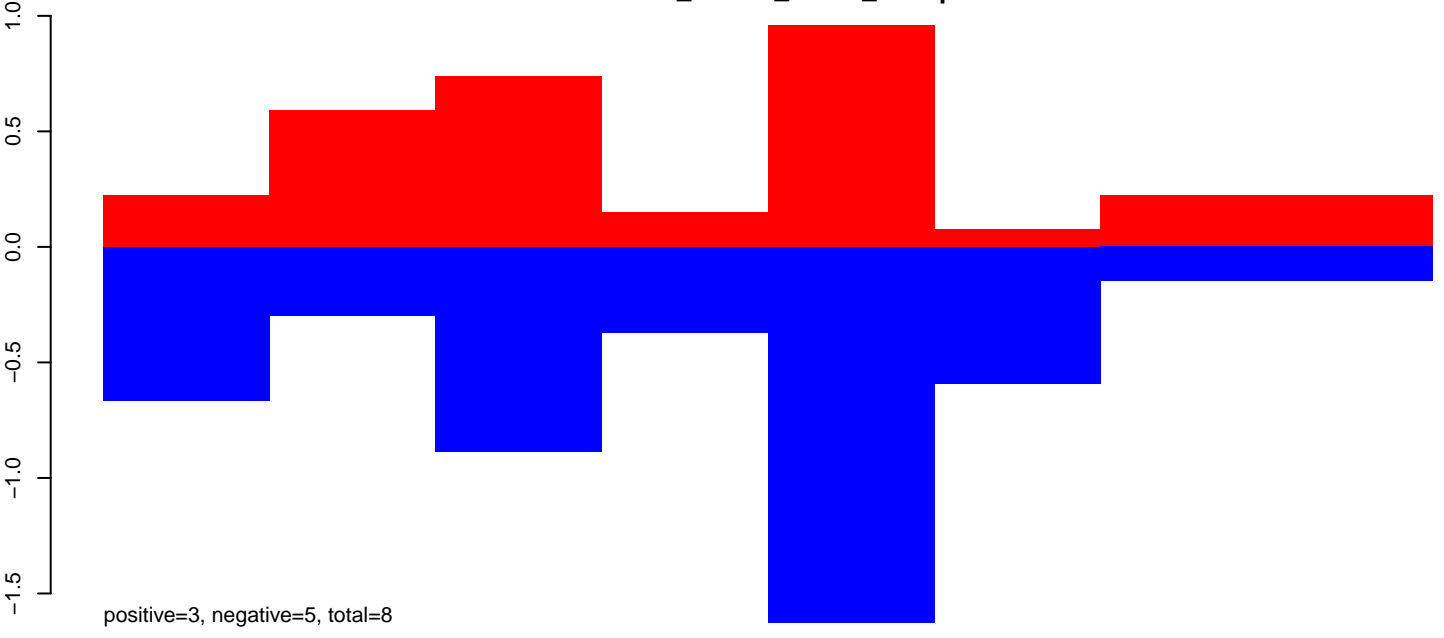
CuQuin_Testes_NL.rep



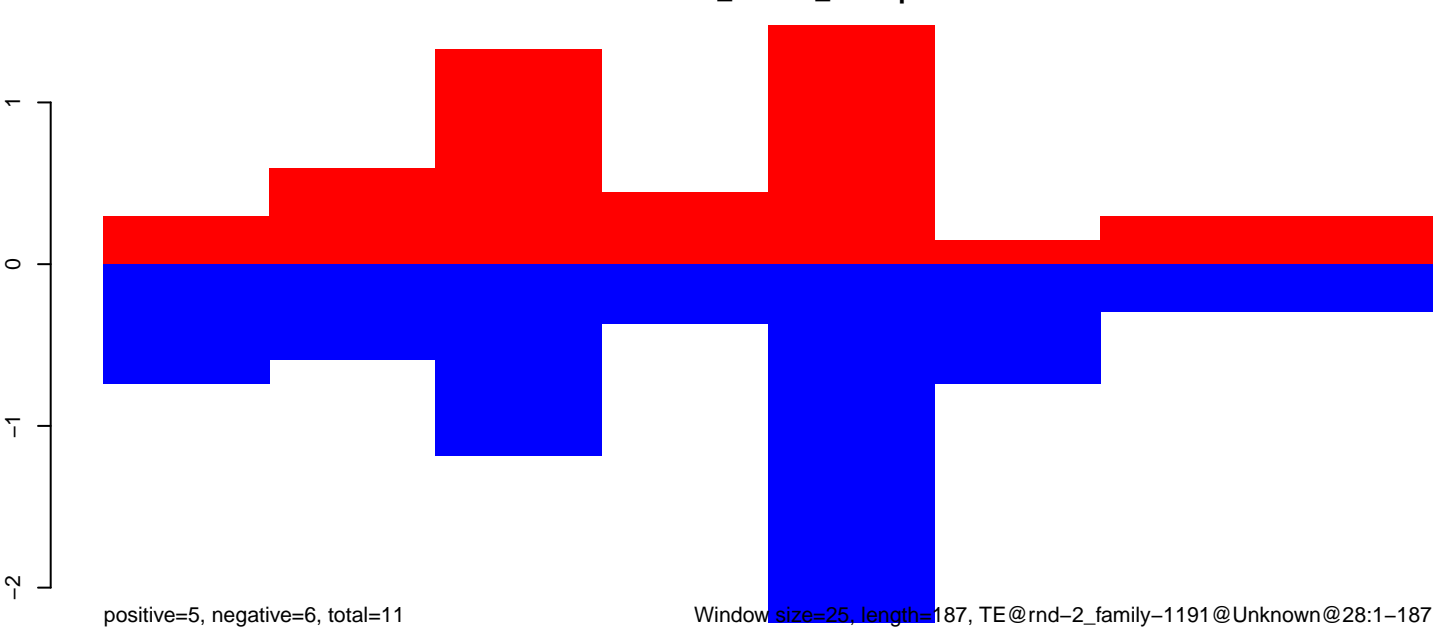
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



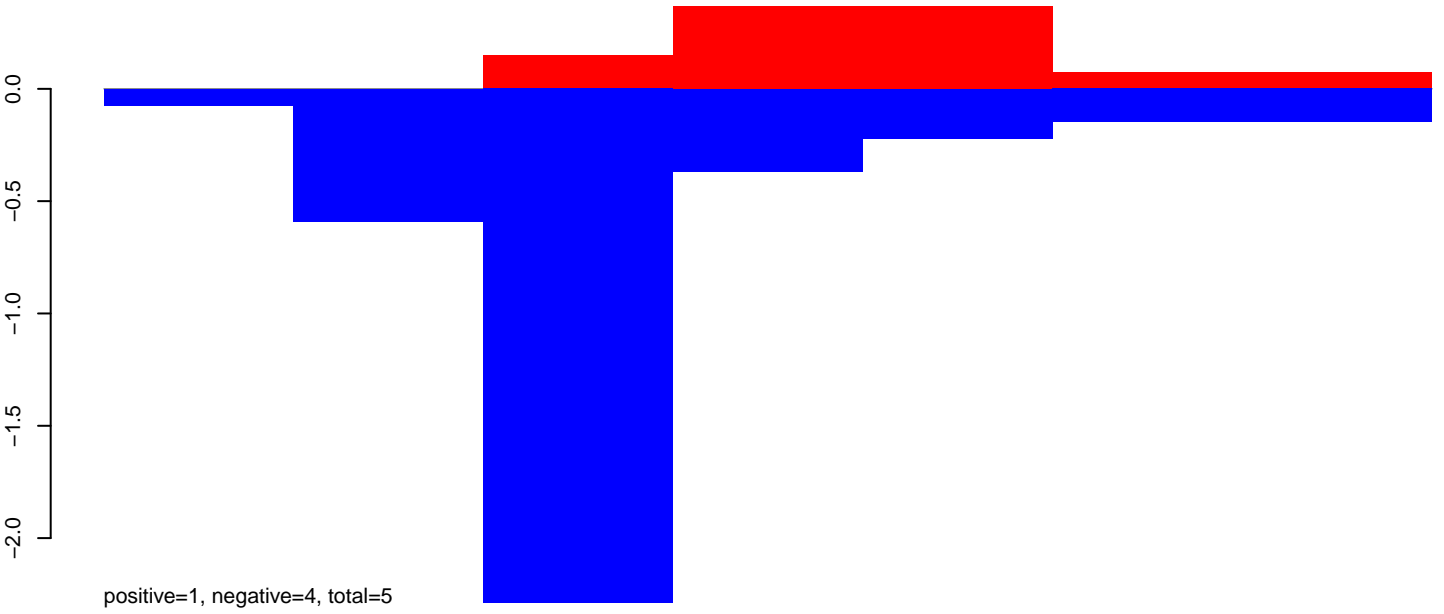
CuQuin_Testes_NL.rep



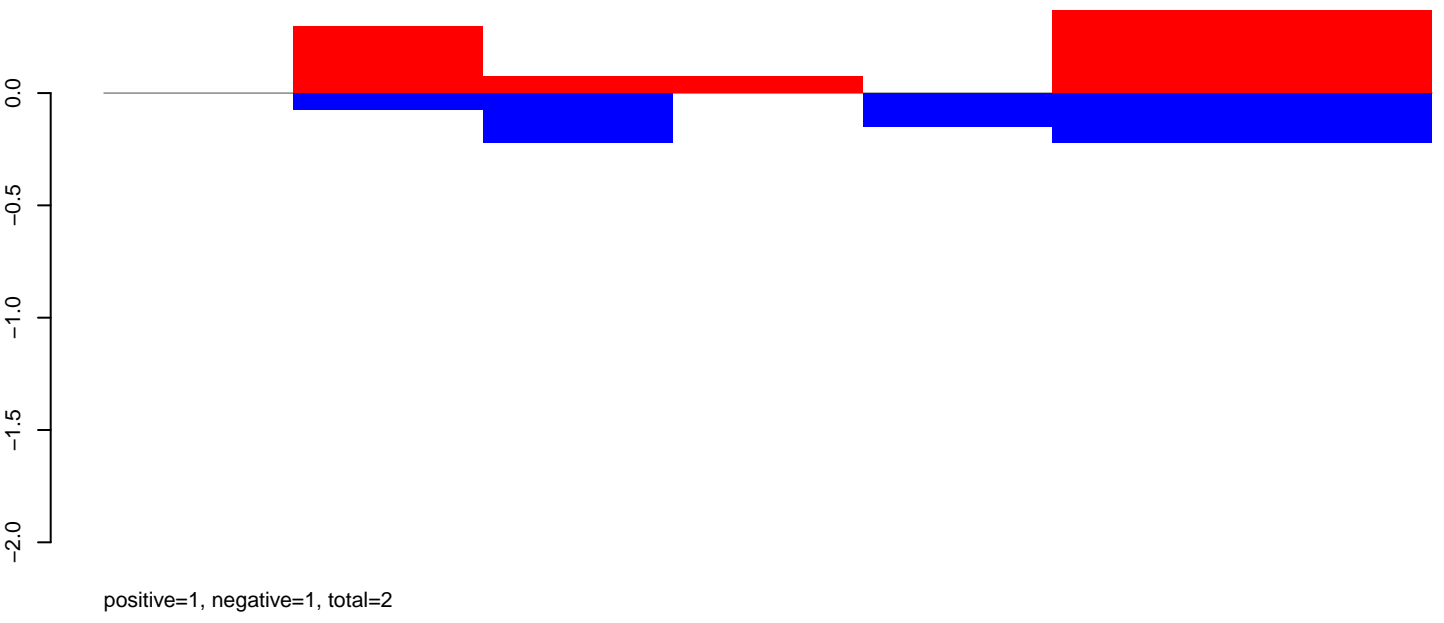
Window size=25, length=187, TE@rnd-2_family-1191@Unknown@28:1-187

50 100 150 200

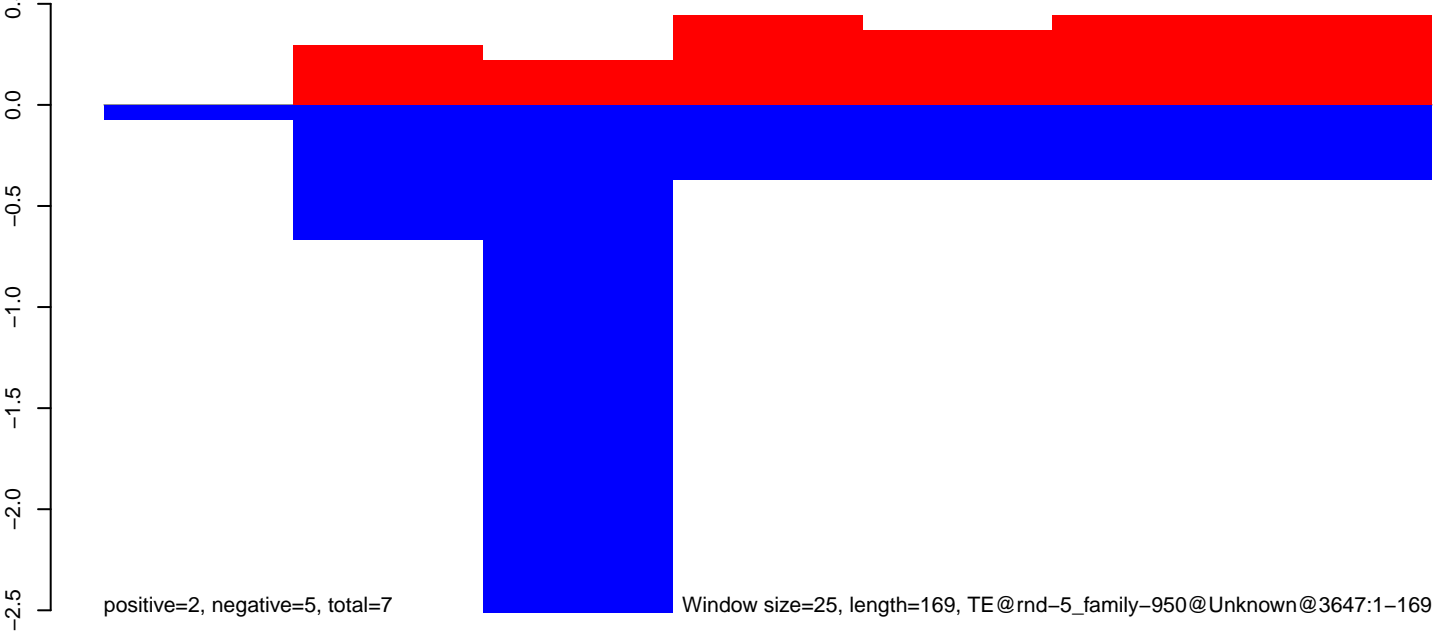
CuQuin_Testes_NL.18_23.rep



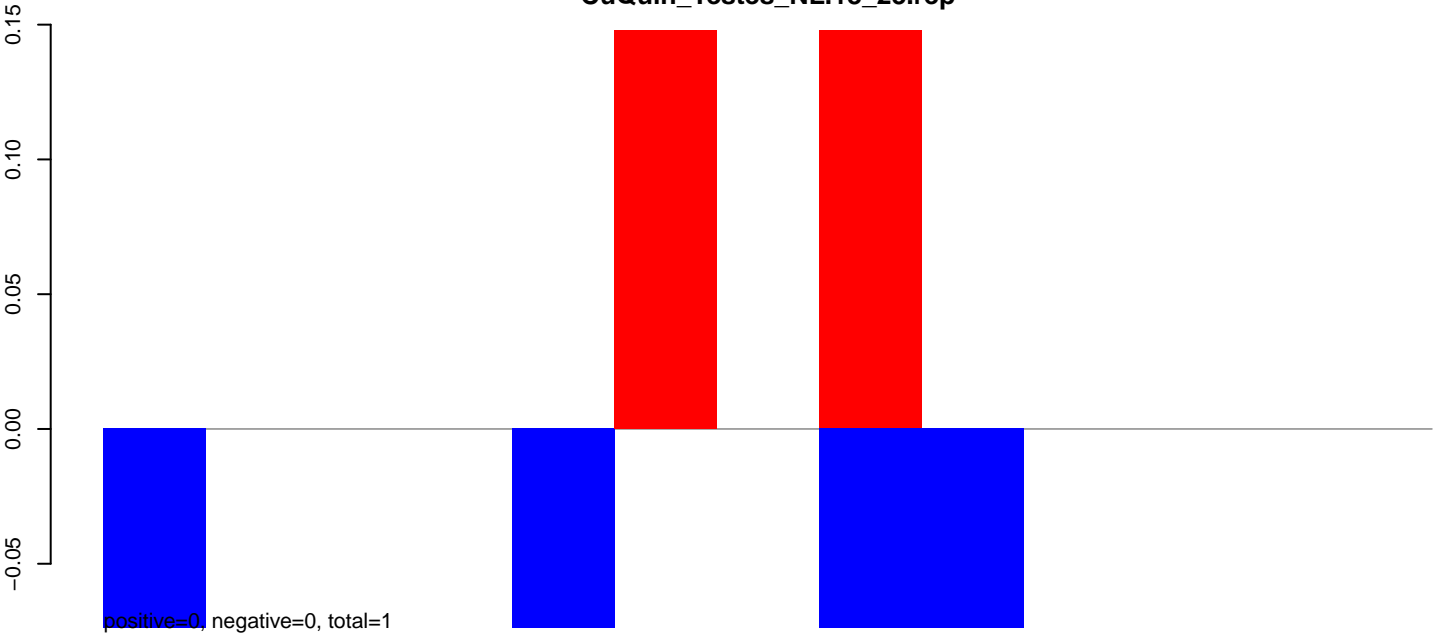
CuQuin_Testes_NL.24_35.rep



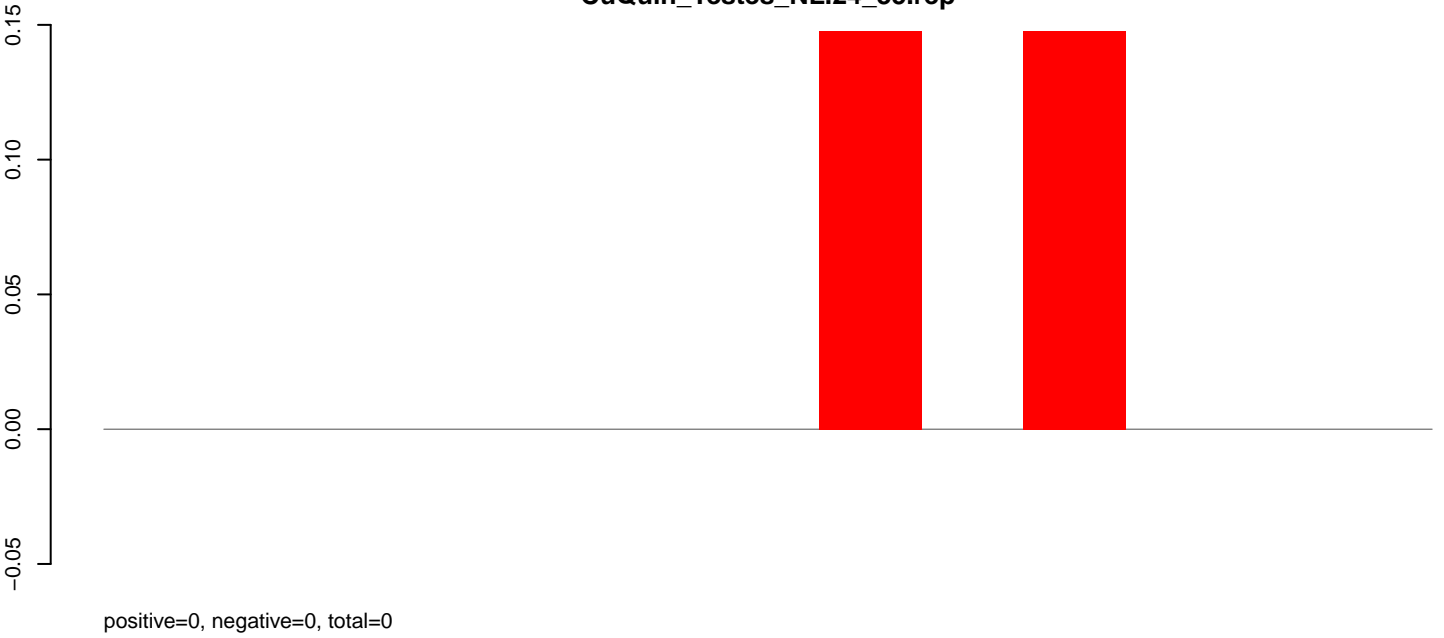
CuQuin_Testes_NL.rep



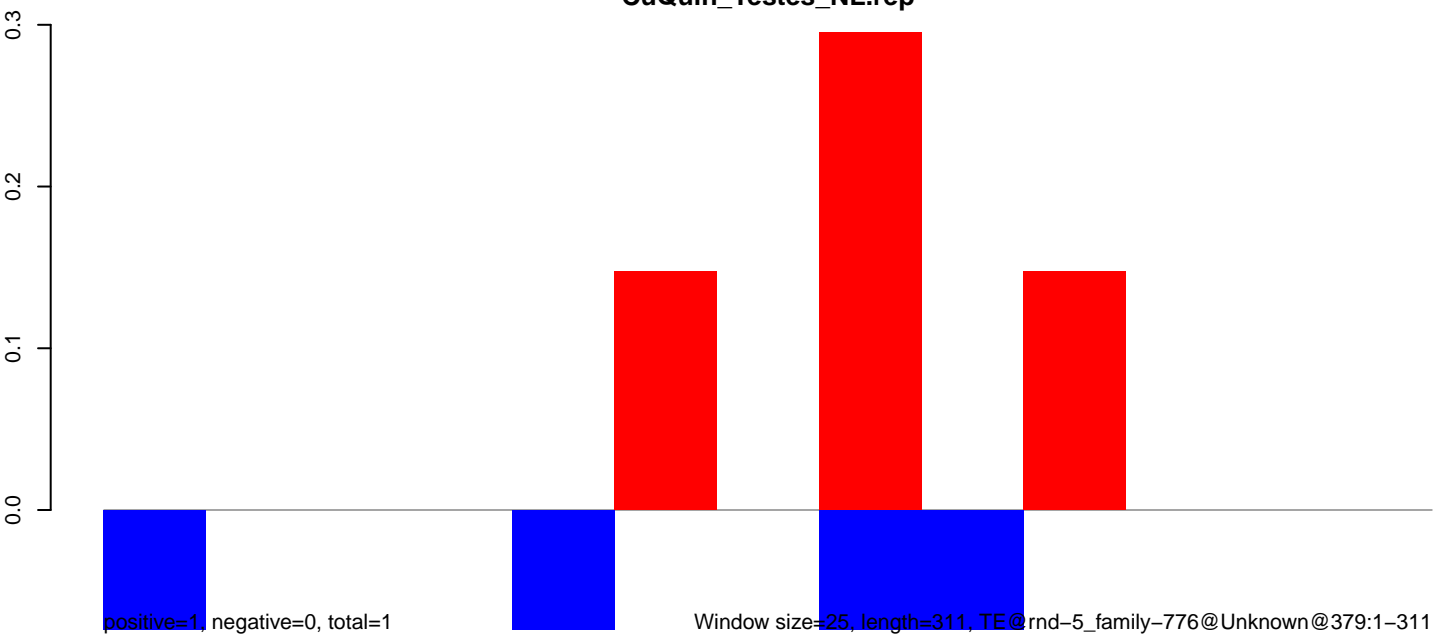
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



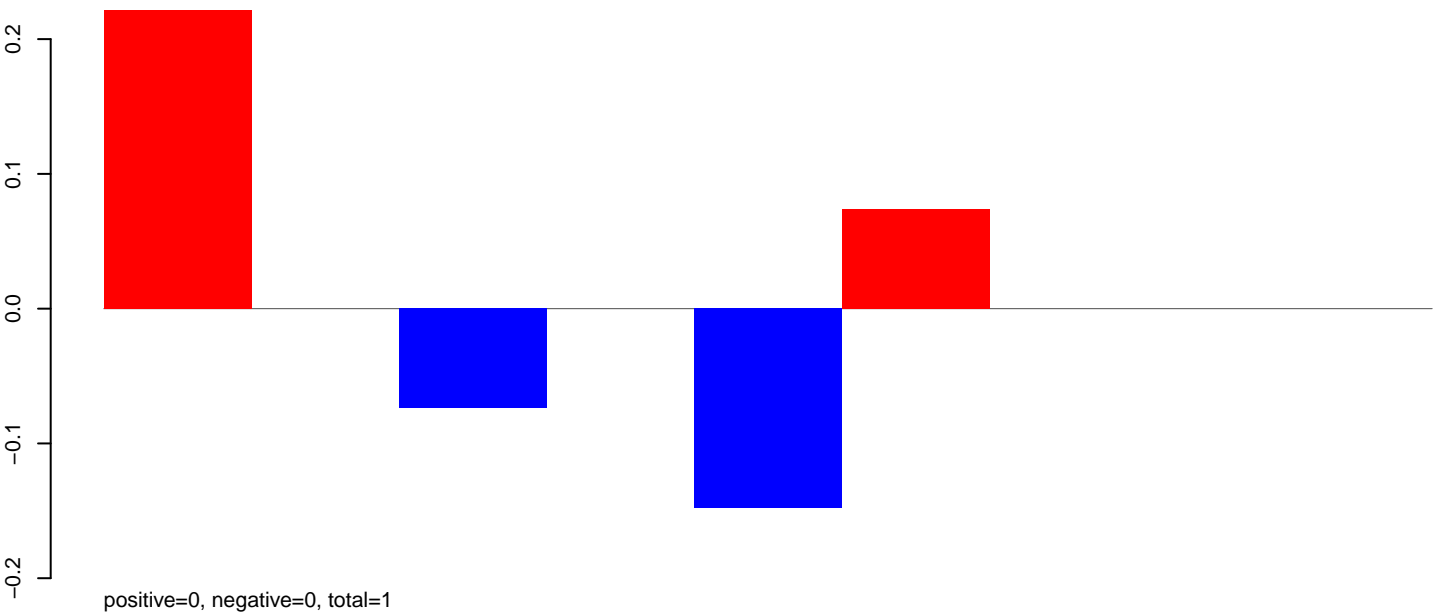
CuQuin_Testes_NL.rep



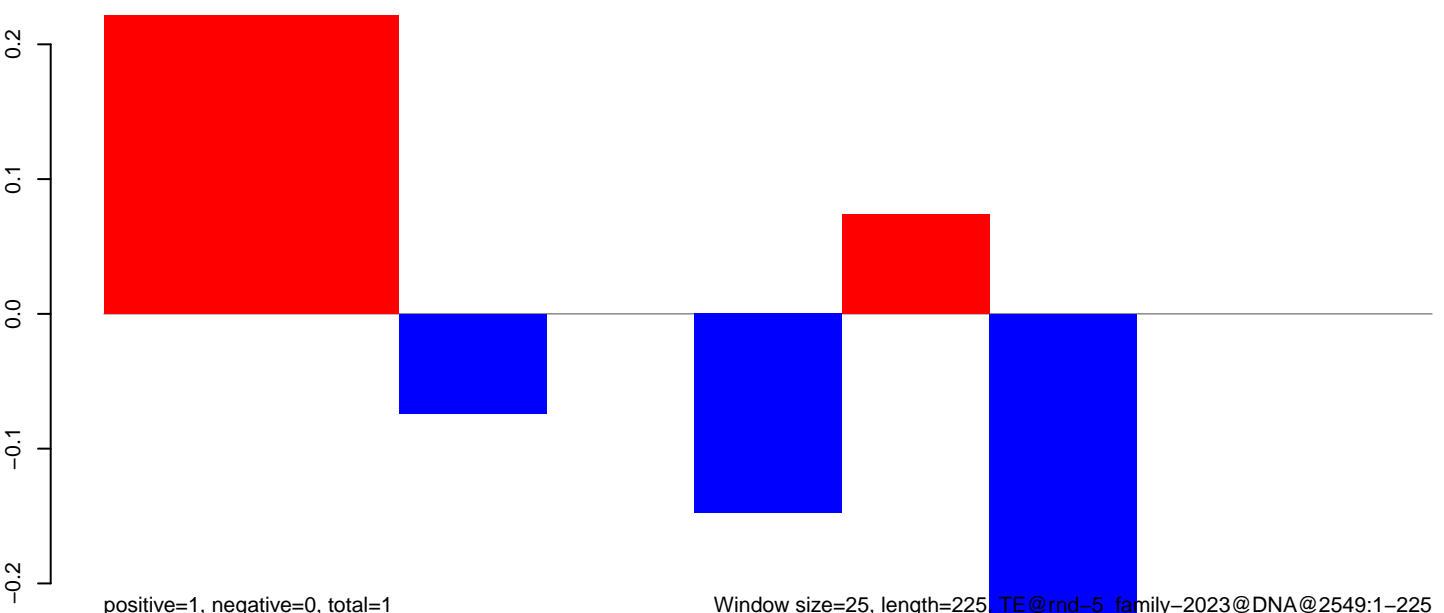
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



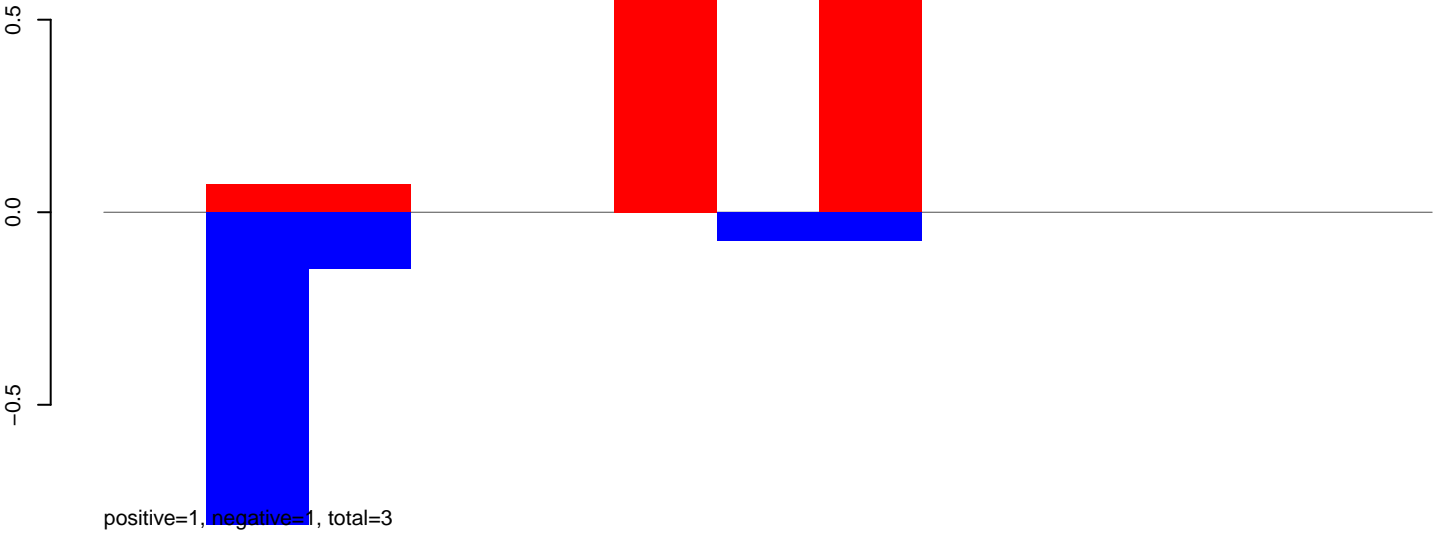
CuQuin_Testes_NL.rep



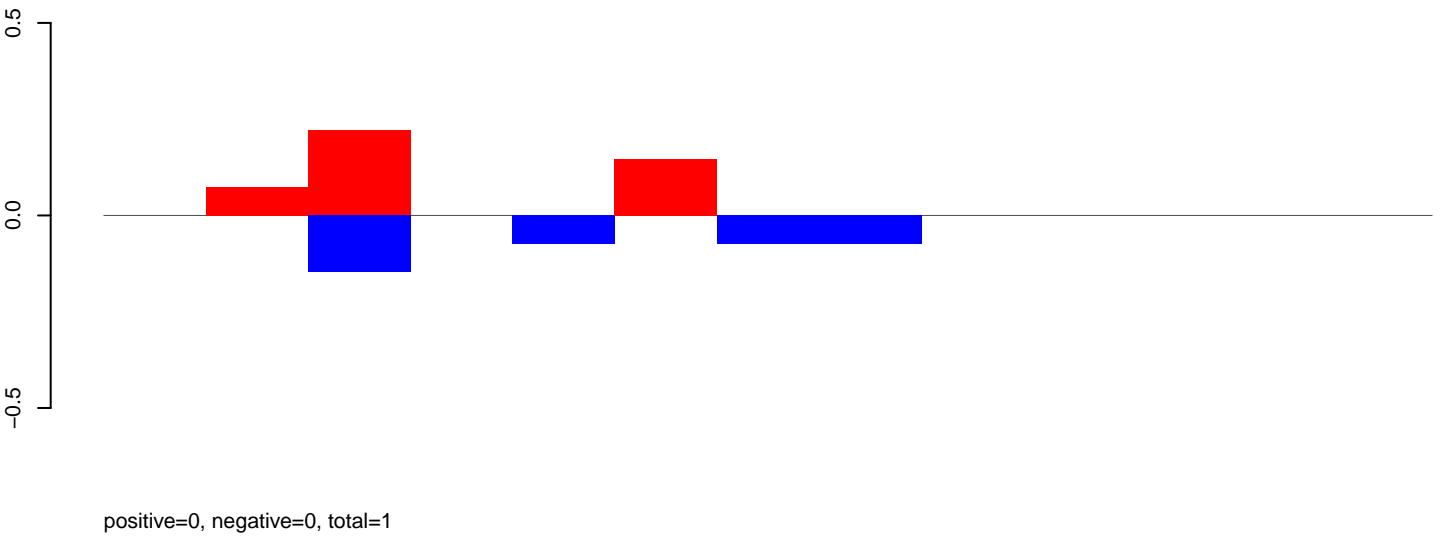
Window size=25, length=225, TE@rnd-5_family-2023@DNA@2549:1-225

50 100 150 200 250

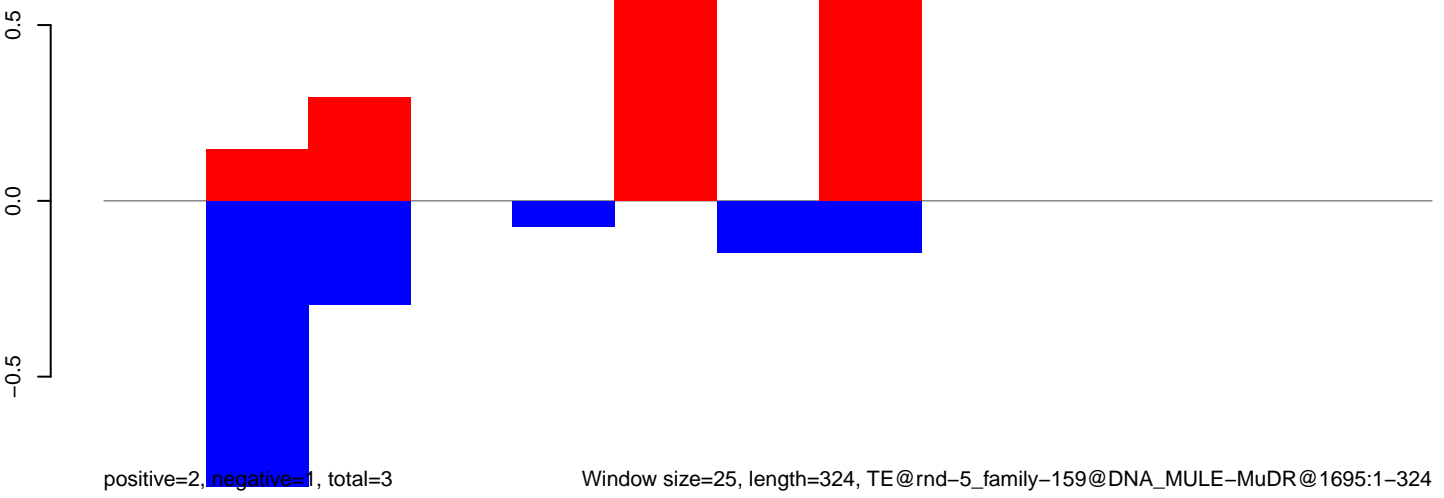
CuQuin_Testes_NL.18_23.rep



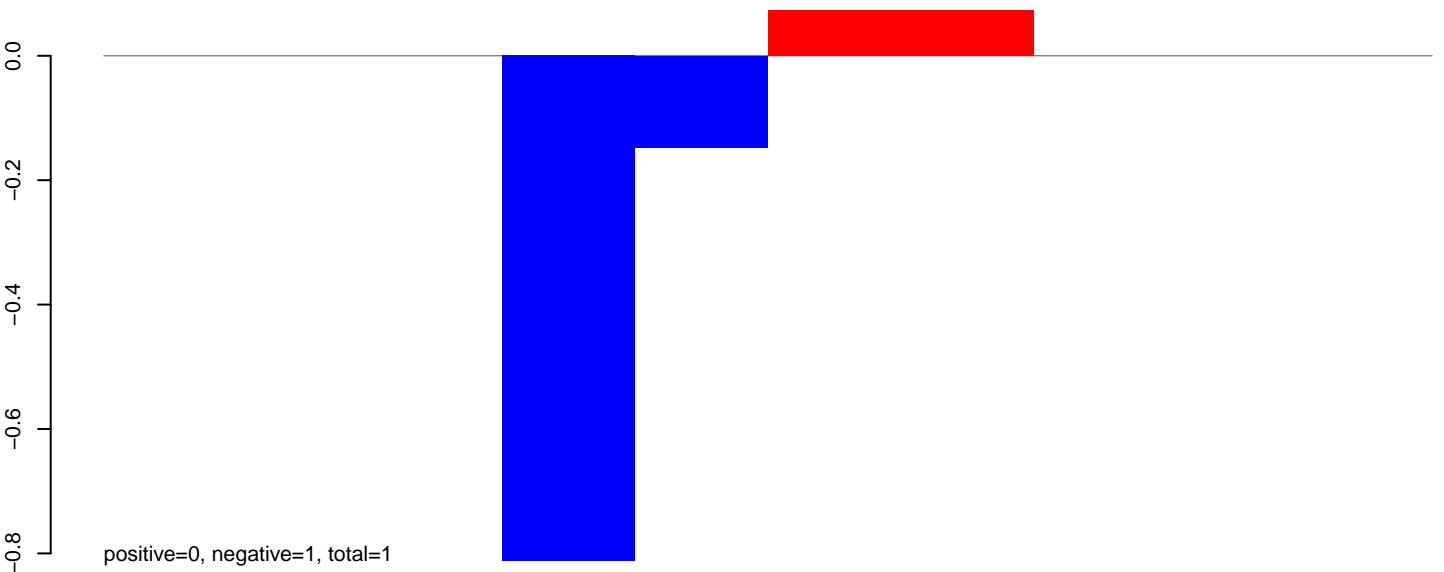
CuQuin_Testes_NL.24_35.rep



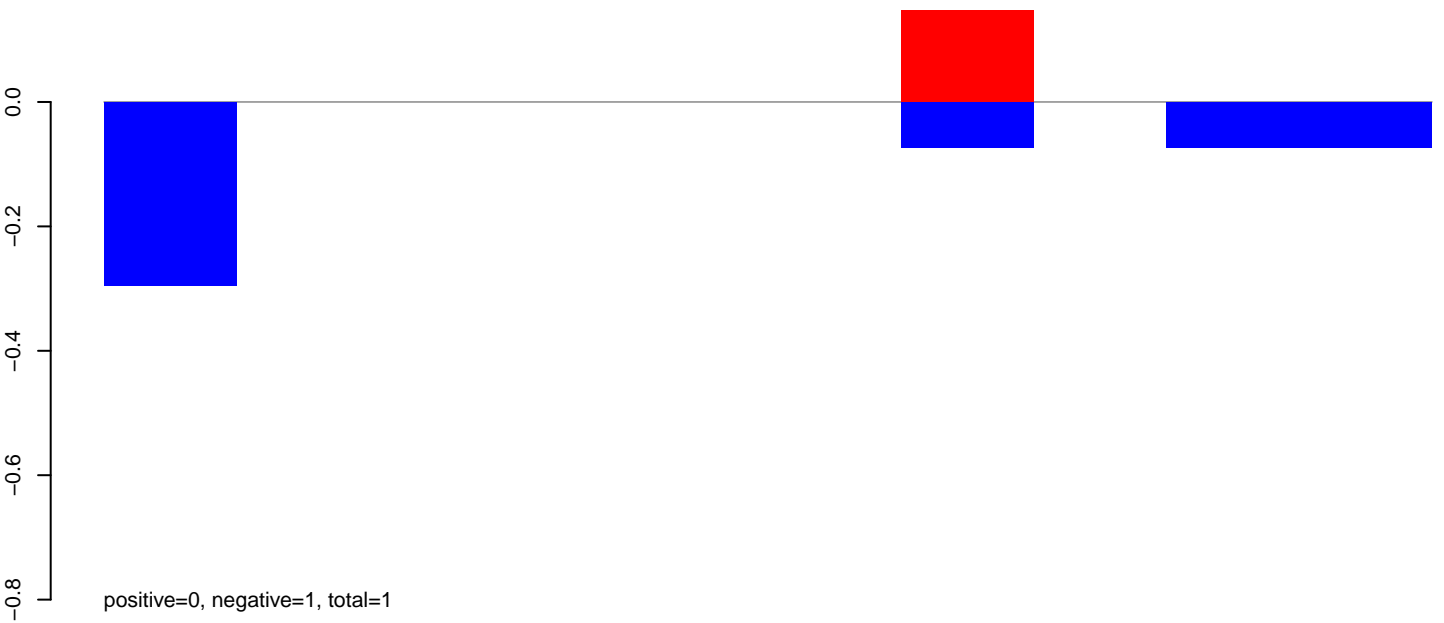
CuQuin_Testes_NL.rep



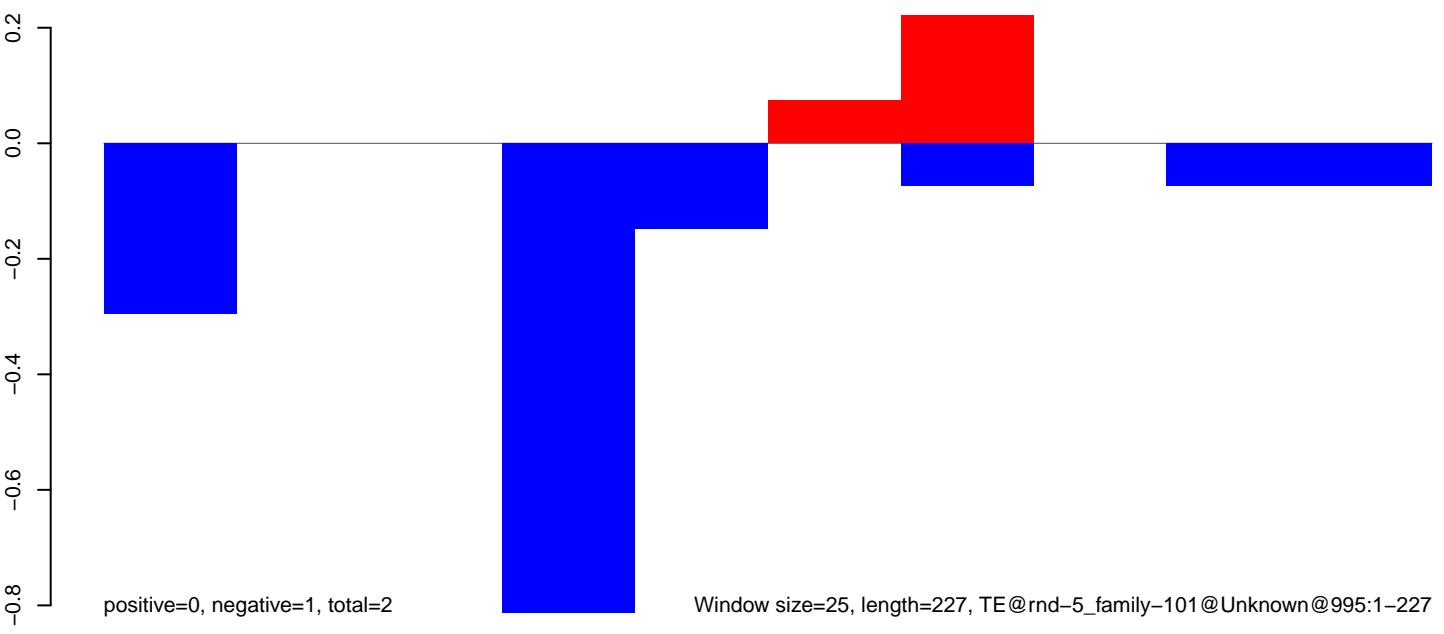
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

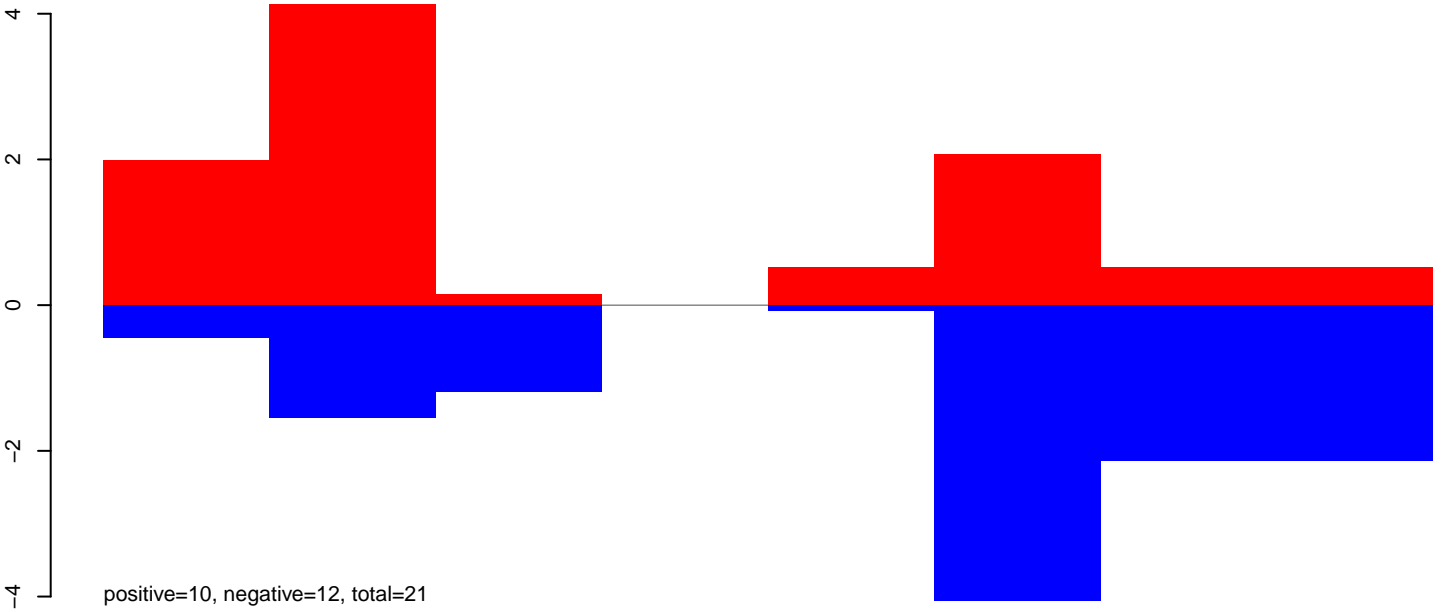


CuQuin_Testes_NL.rep

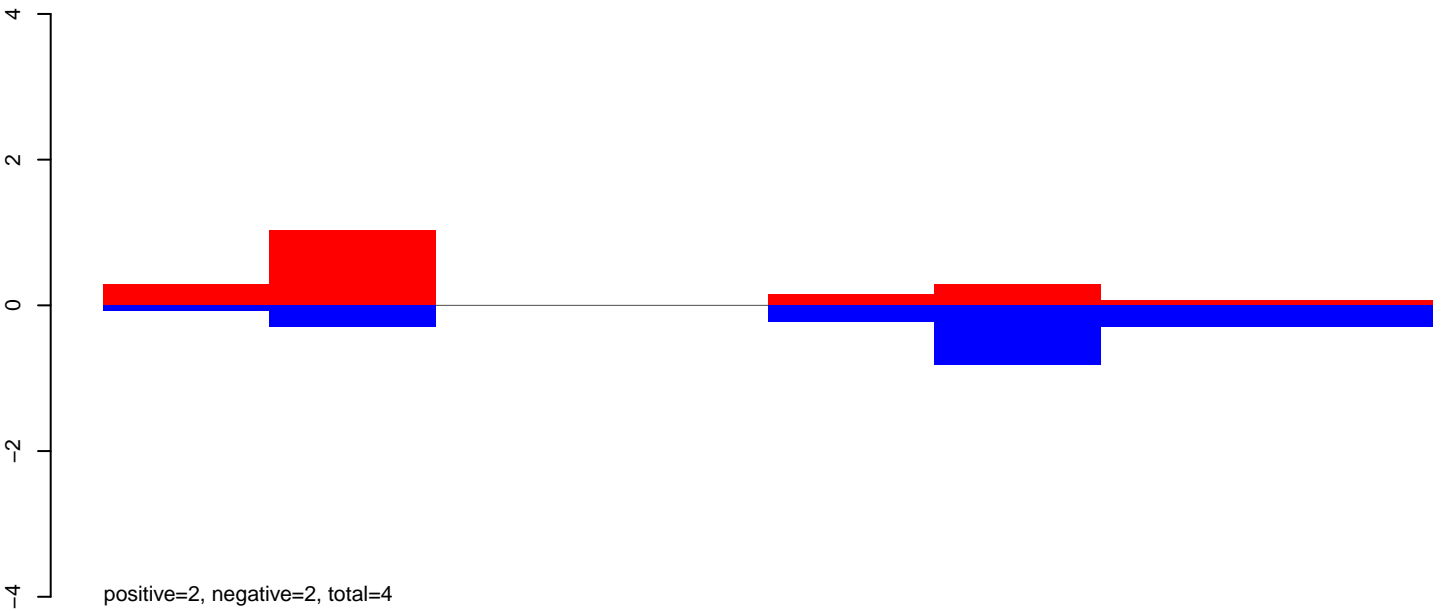


50 100 150 200 250

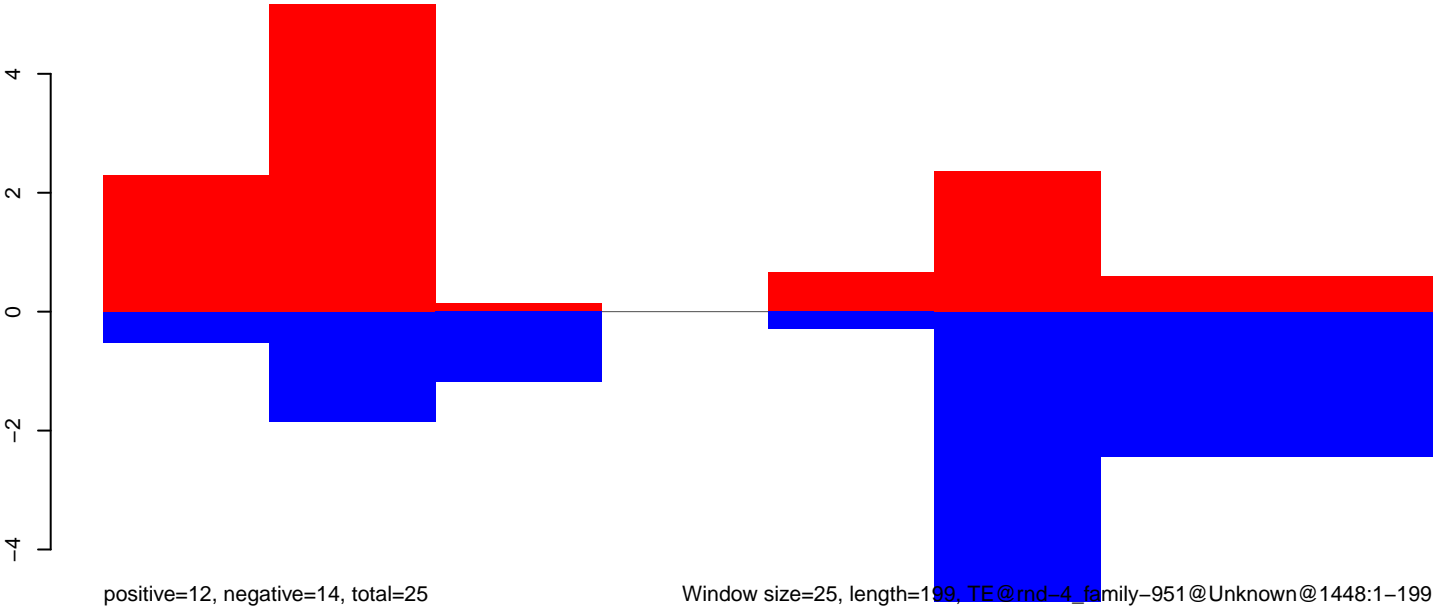
CuQuin_Testes_NL.18_23.rep



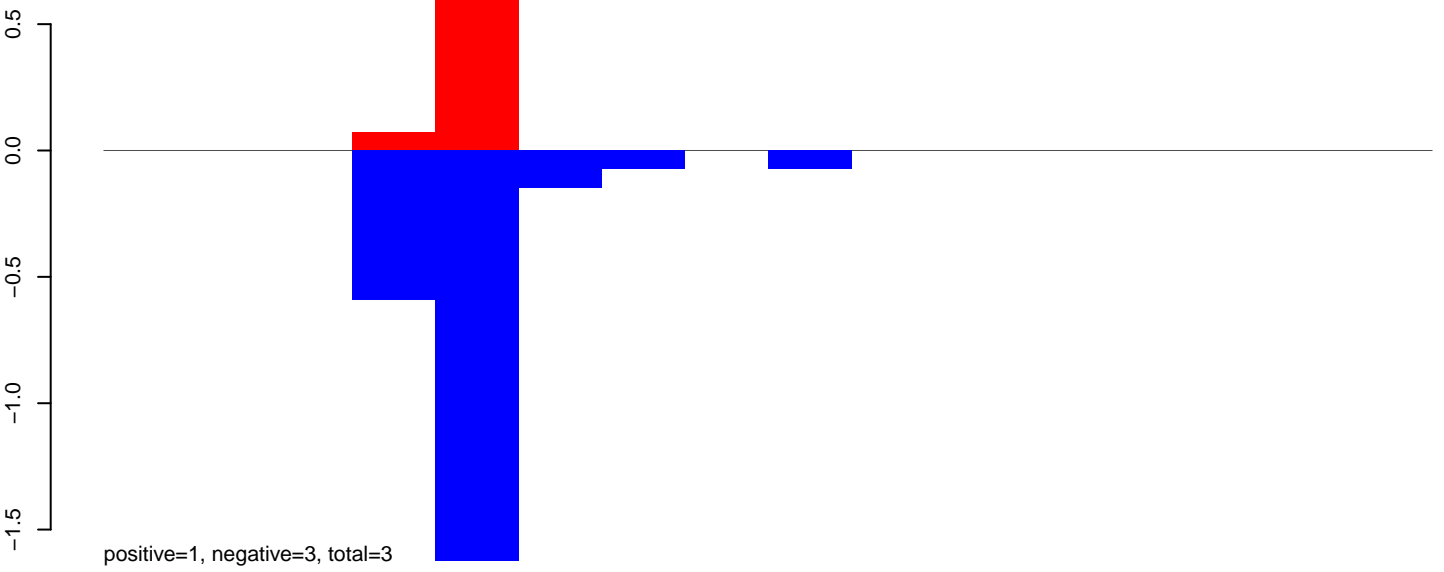
CuQuin_Testes_NL.24_35.rep



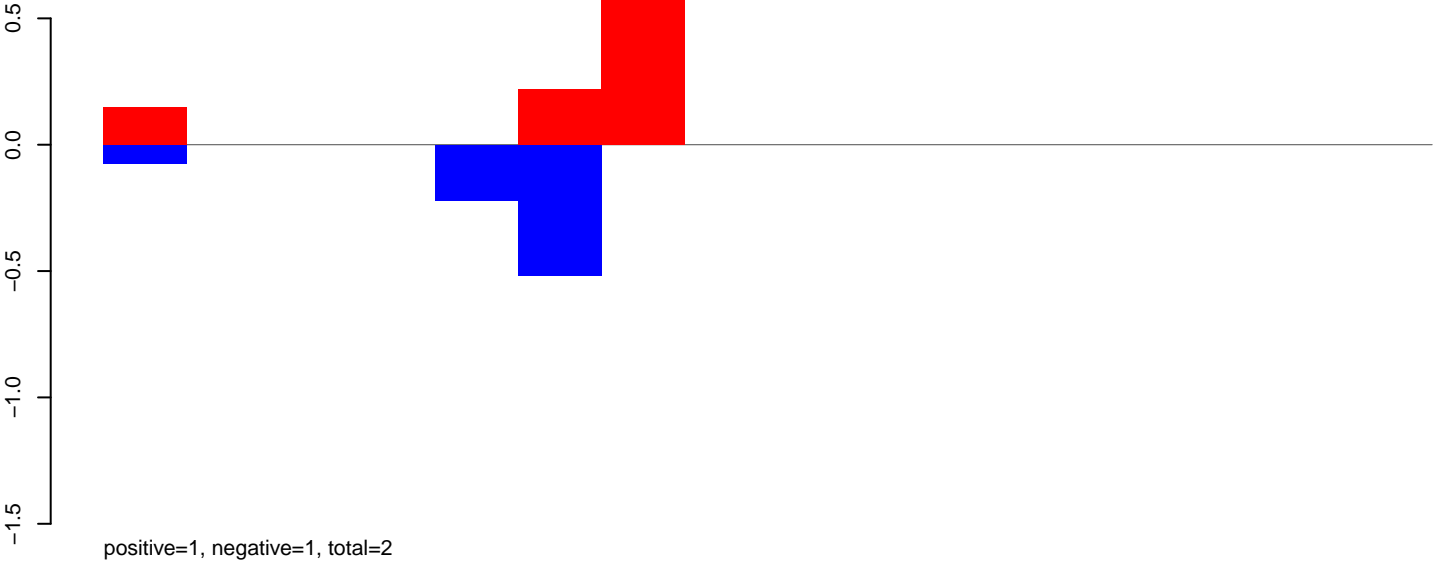
CuQuin_Testes_NL.rep



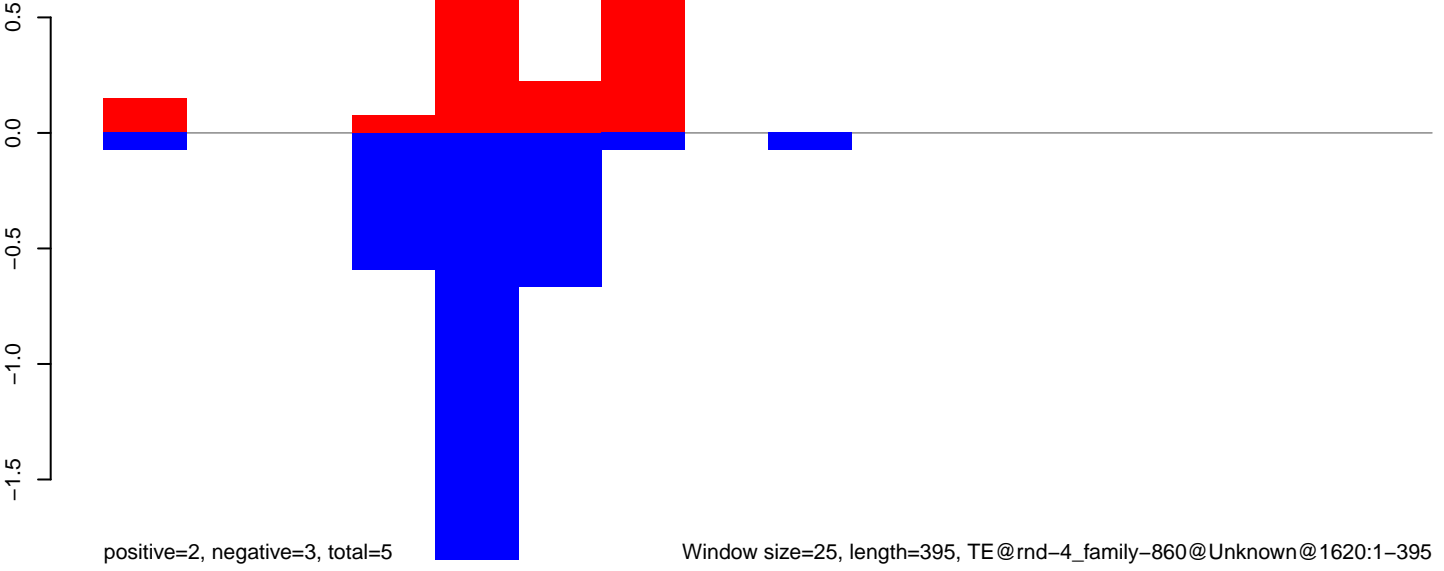
CuQuin_Testes_NL.18_23.rep



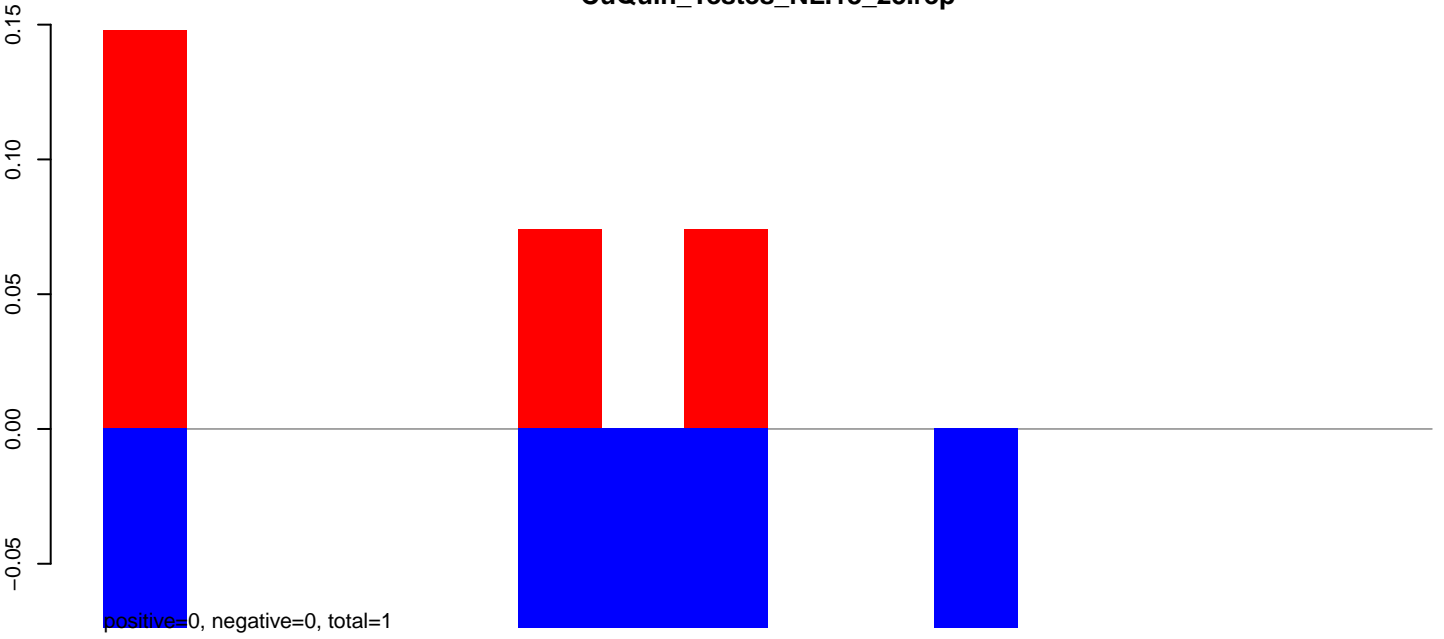
CuQuin_Testes_NL.24_35.rep



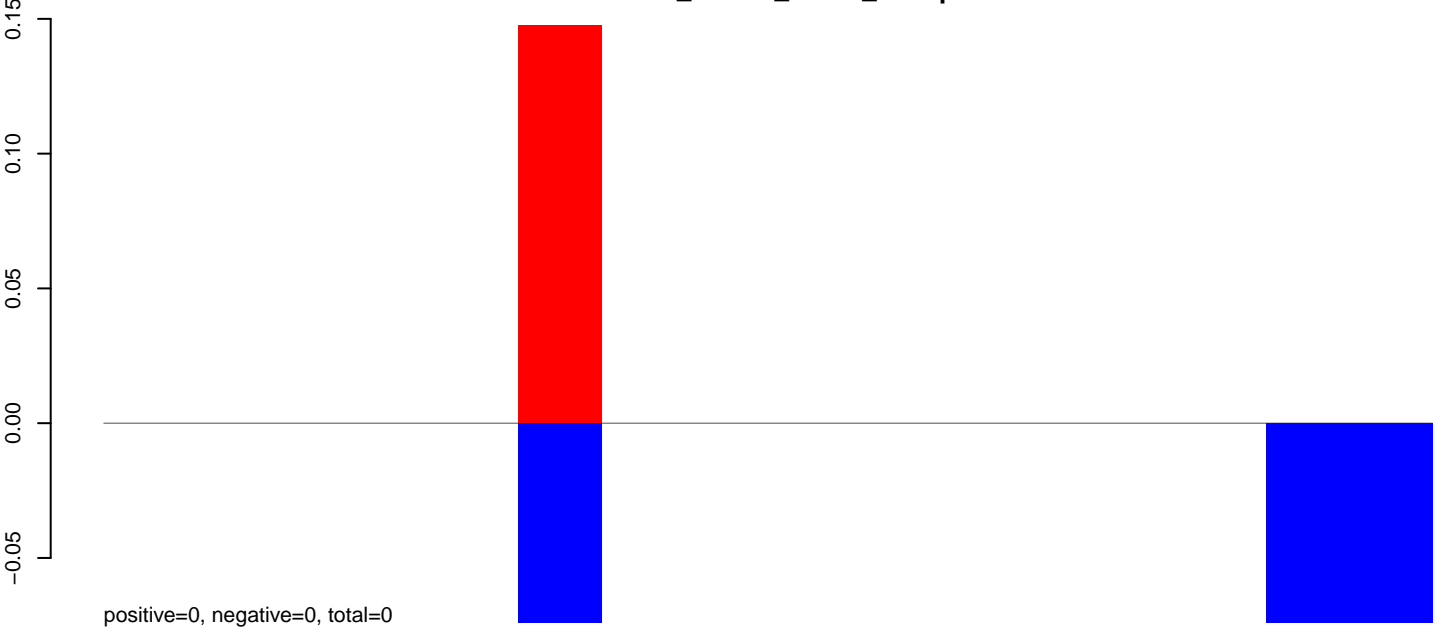
CuQuin_Testes_NL.rep



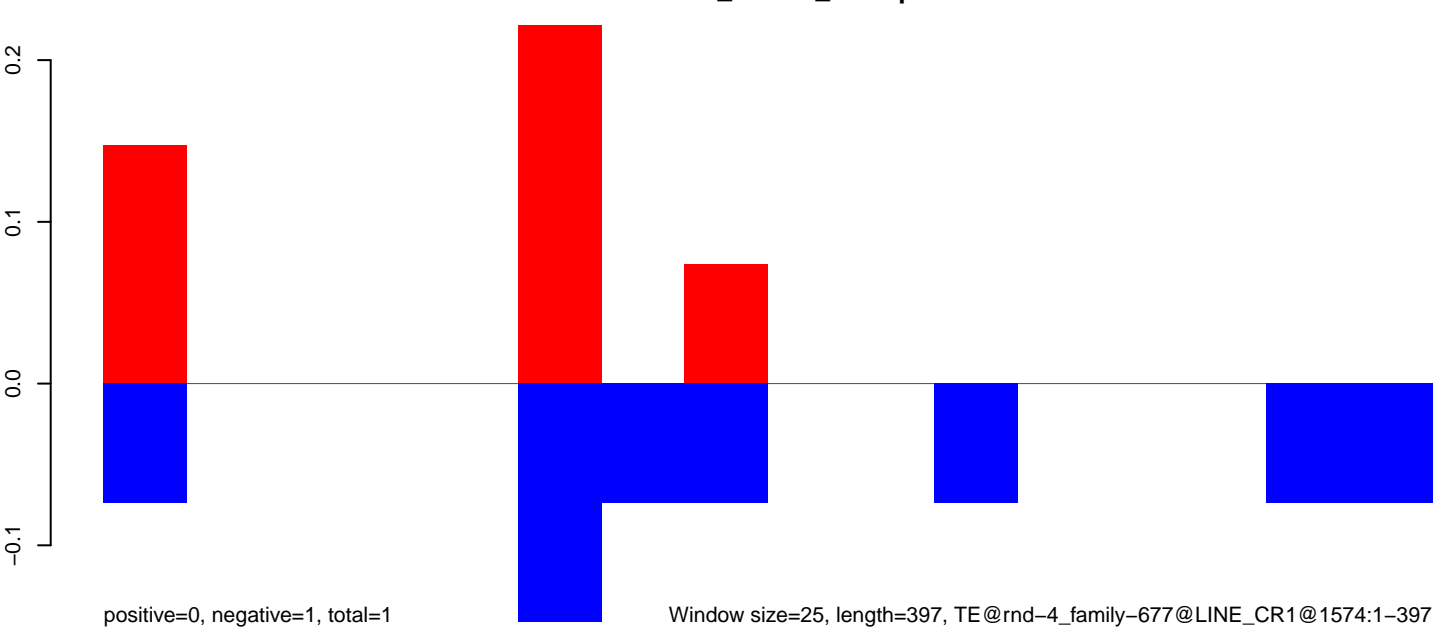
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



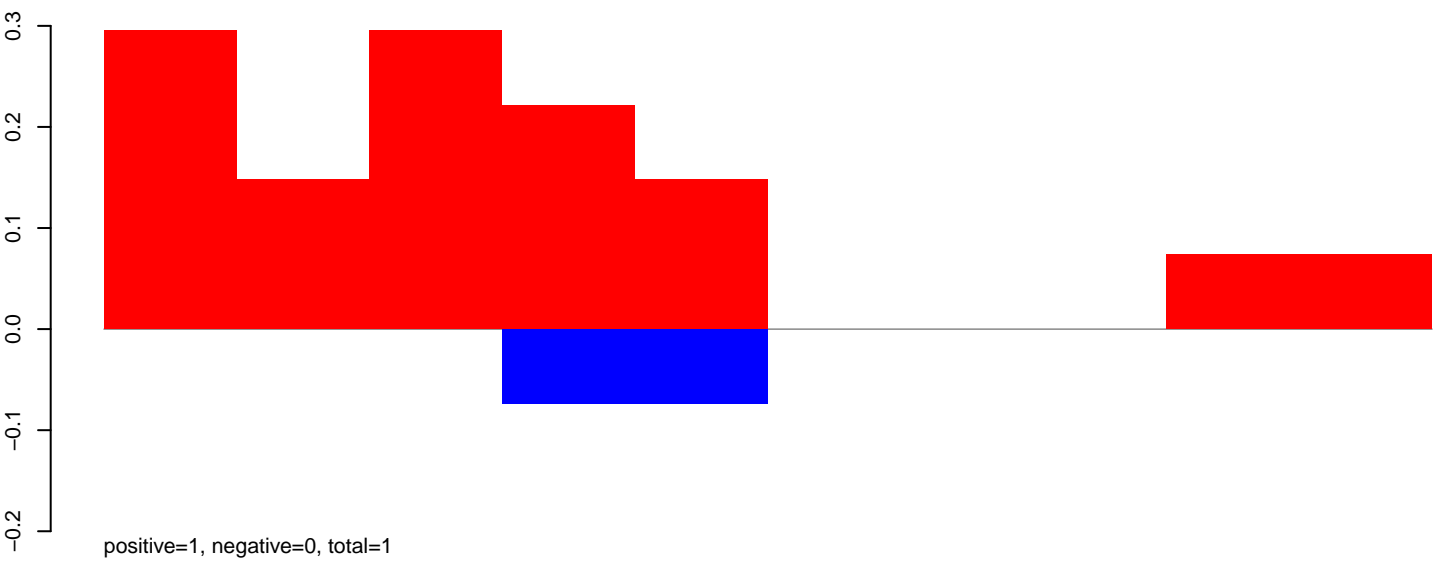
CuQuin_Testes_NL.rep



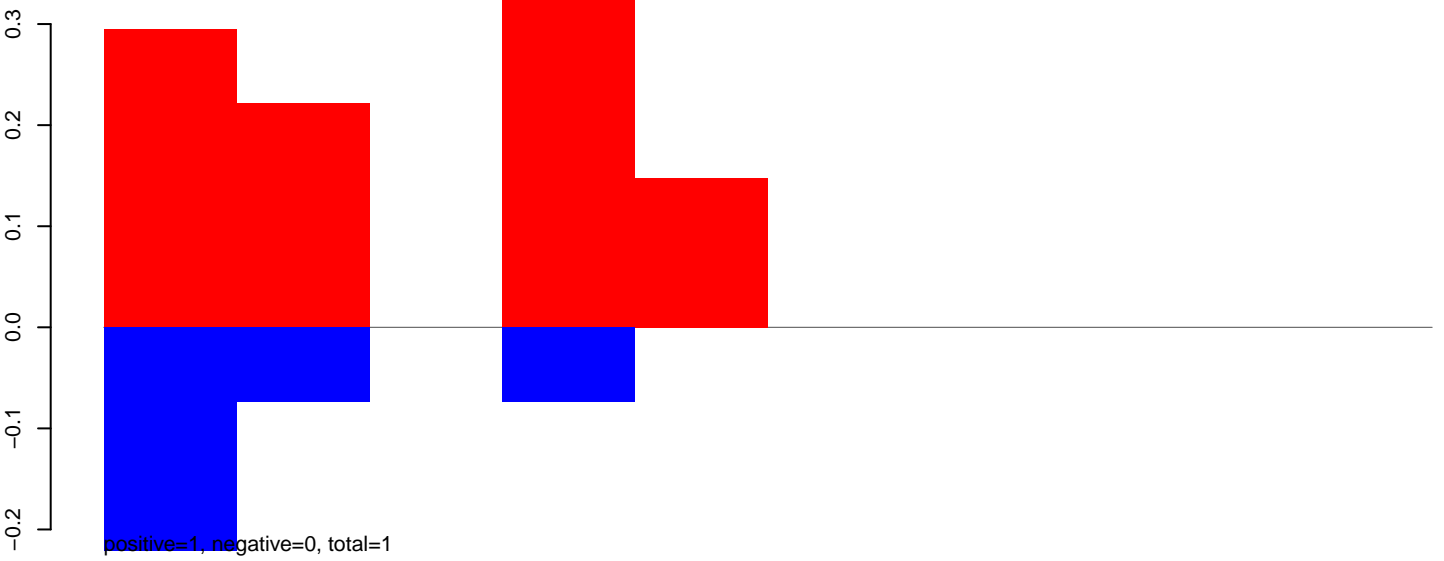
Window size=25, length=397, TE@rnd-4_family-677@LINE_CR1@1574:1-397

100 200 300 400

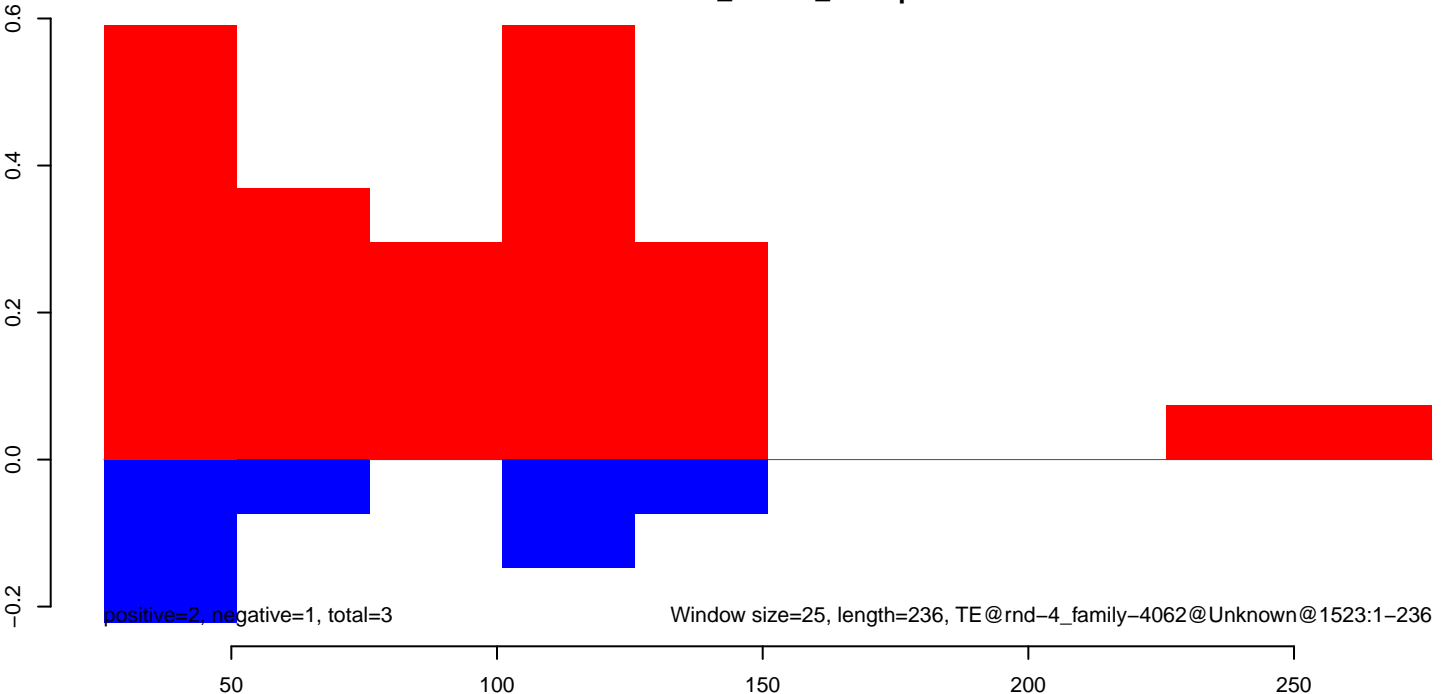
CuQuin_Testes_NL.18_23.rep



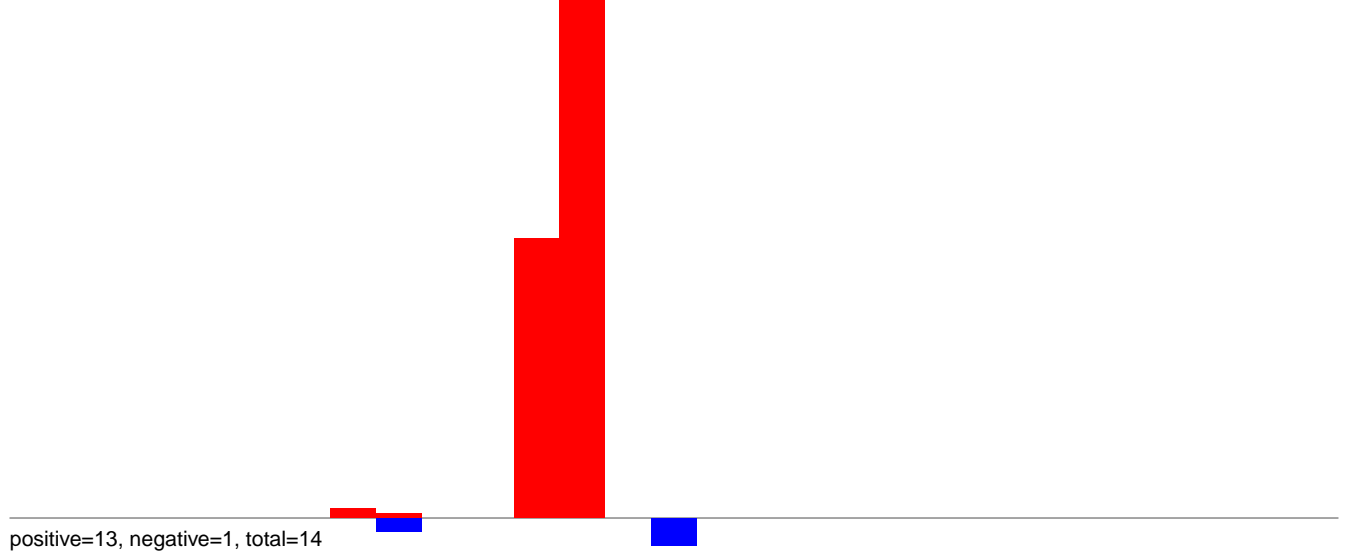
CuQuin_Testes_NL.24_35.rep



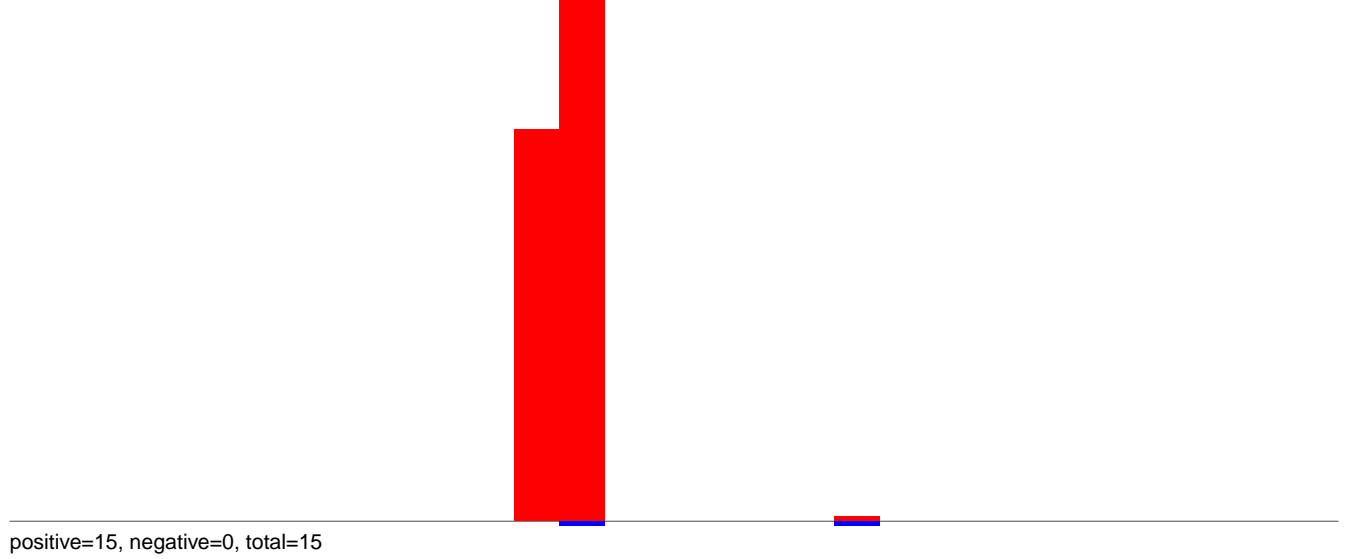
CuQuin_Testes_NL.rep



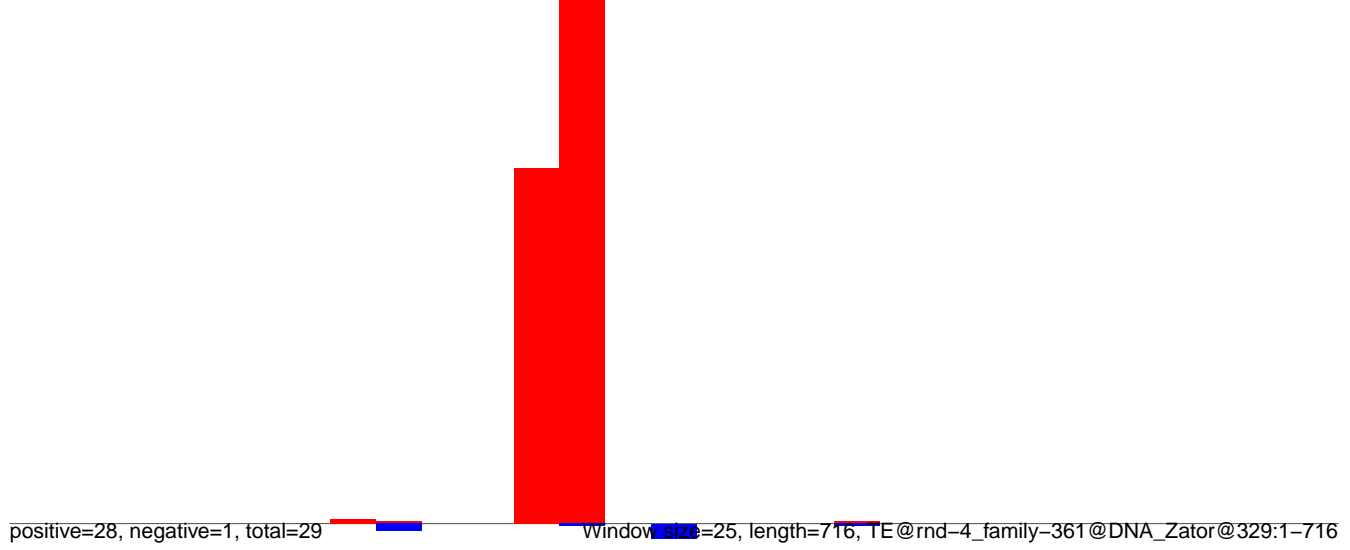
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

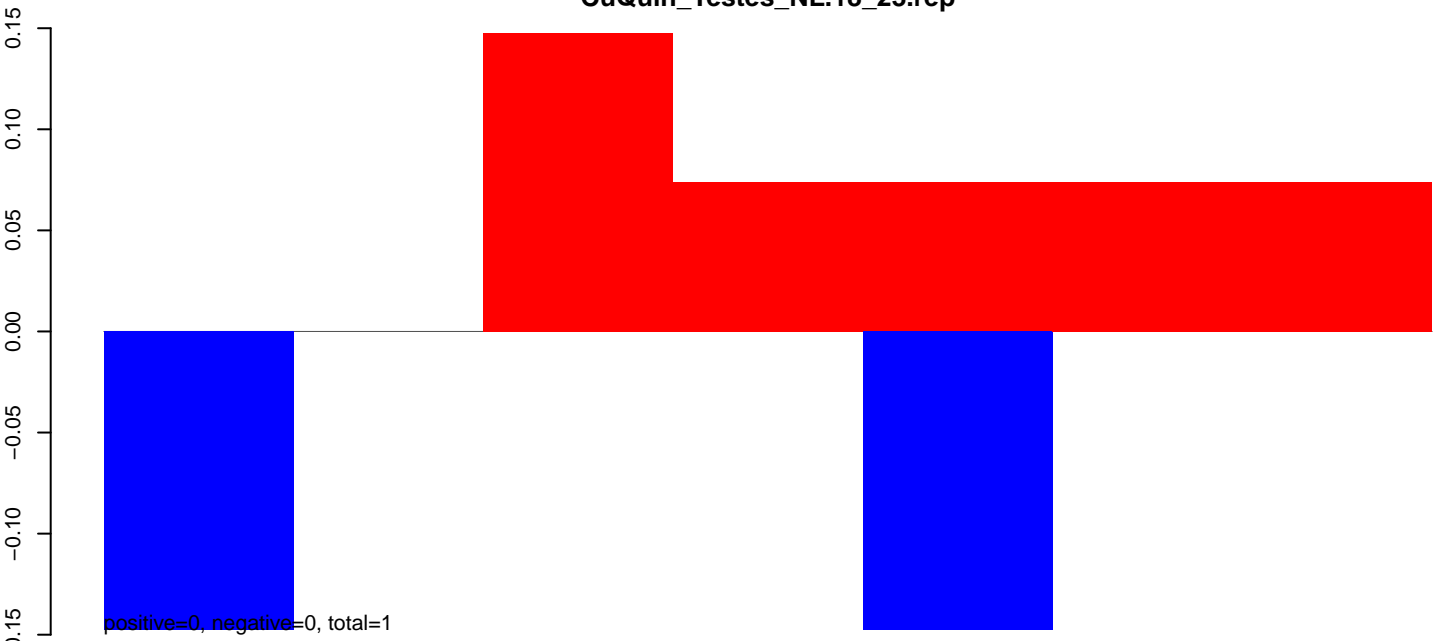


CuQuin_Testes_NL.rep

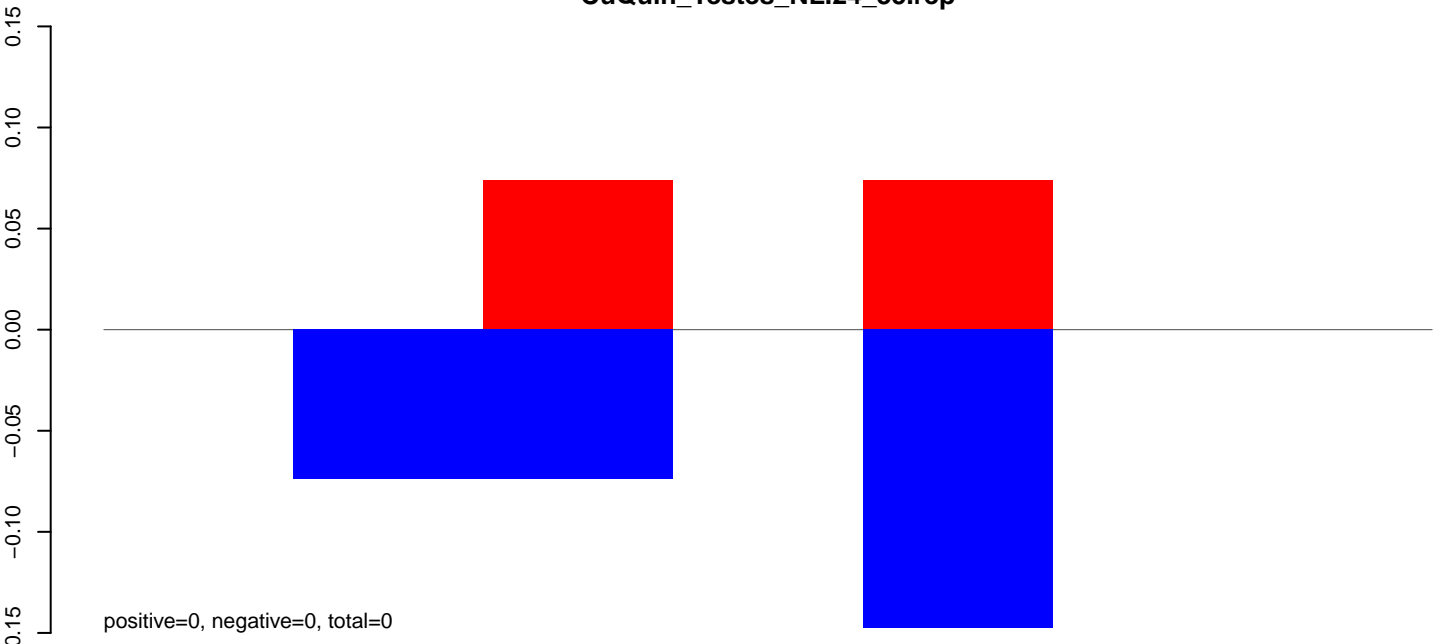


Window size=25, length=716, TE@rnd-4_family-361@DNA_Zator@329:1-716

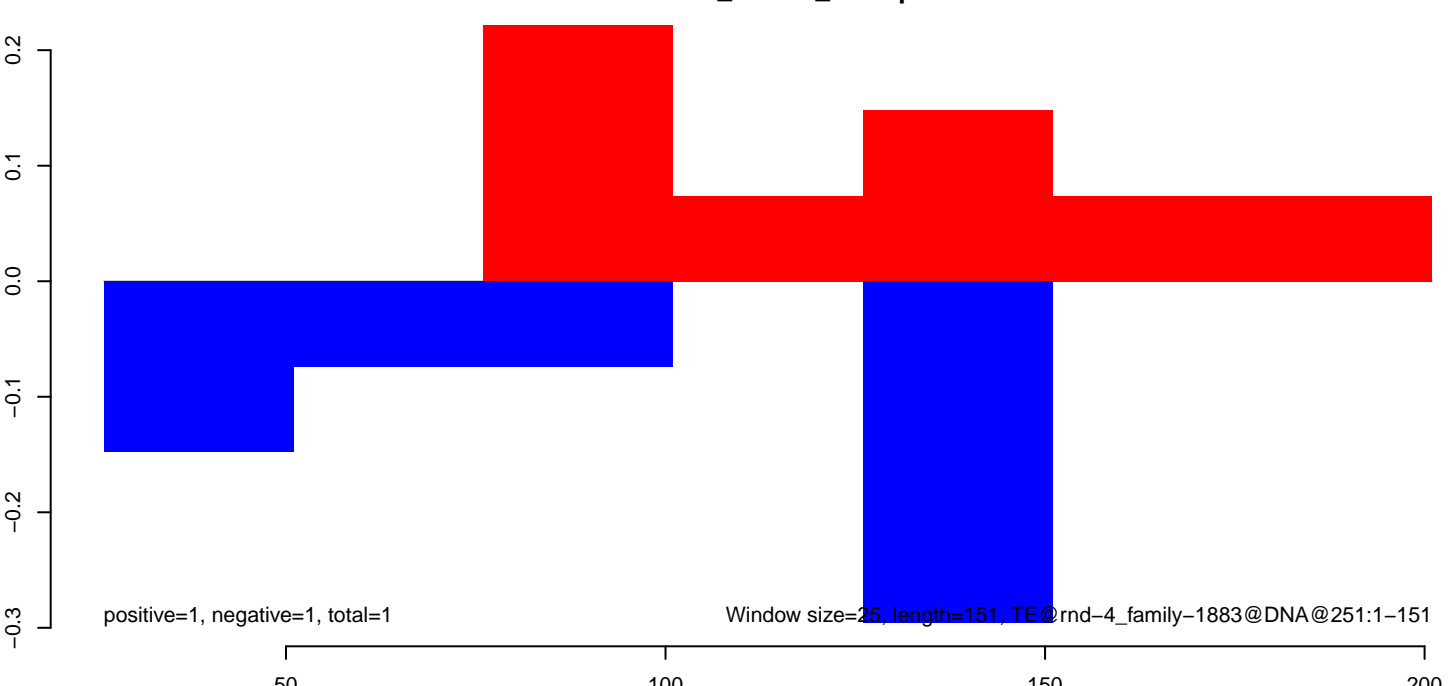
CuQuin_Testes_NL.18_23.rep



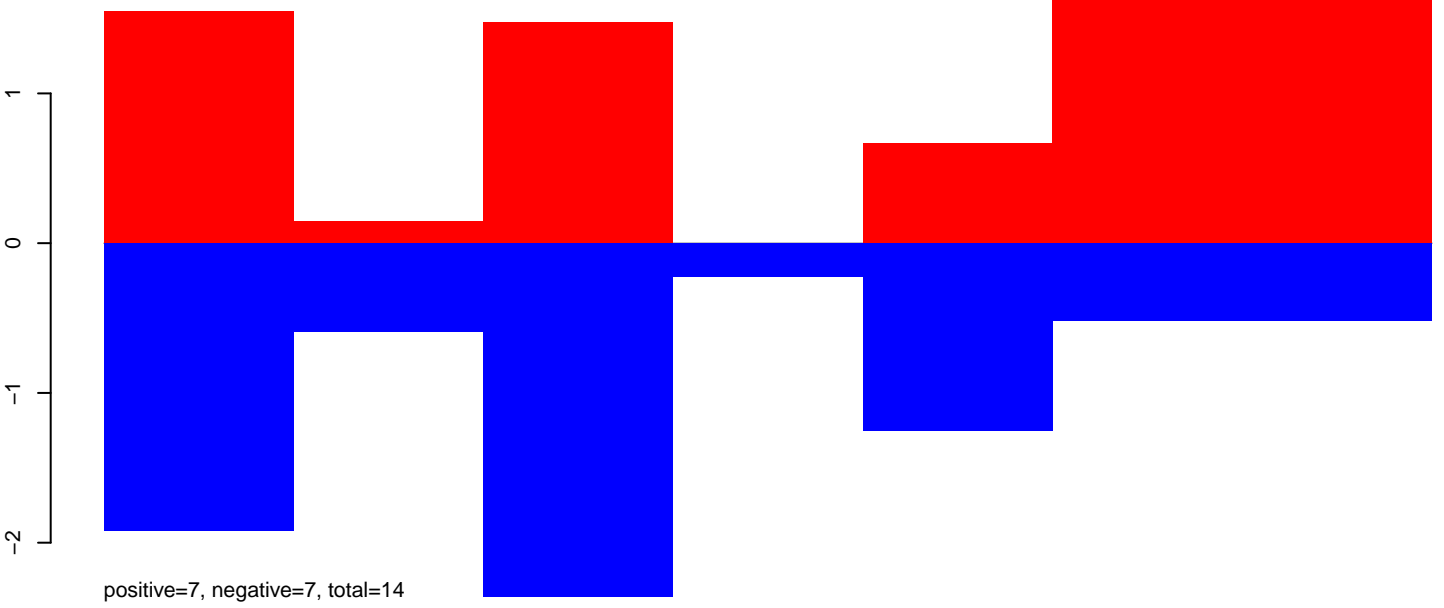
CuQuin_Testes_NL.24_35.rep



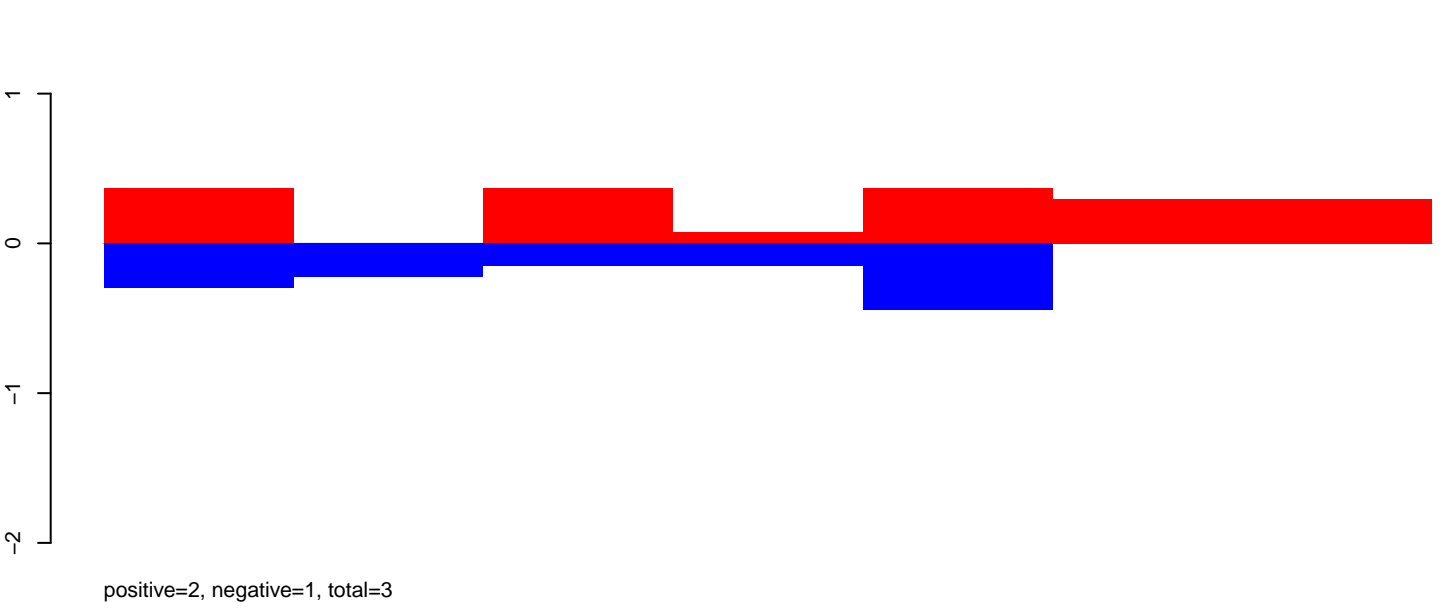
CuQuin_Testes_NL.rep



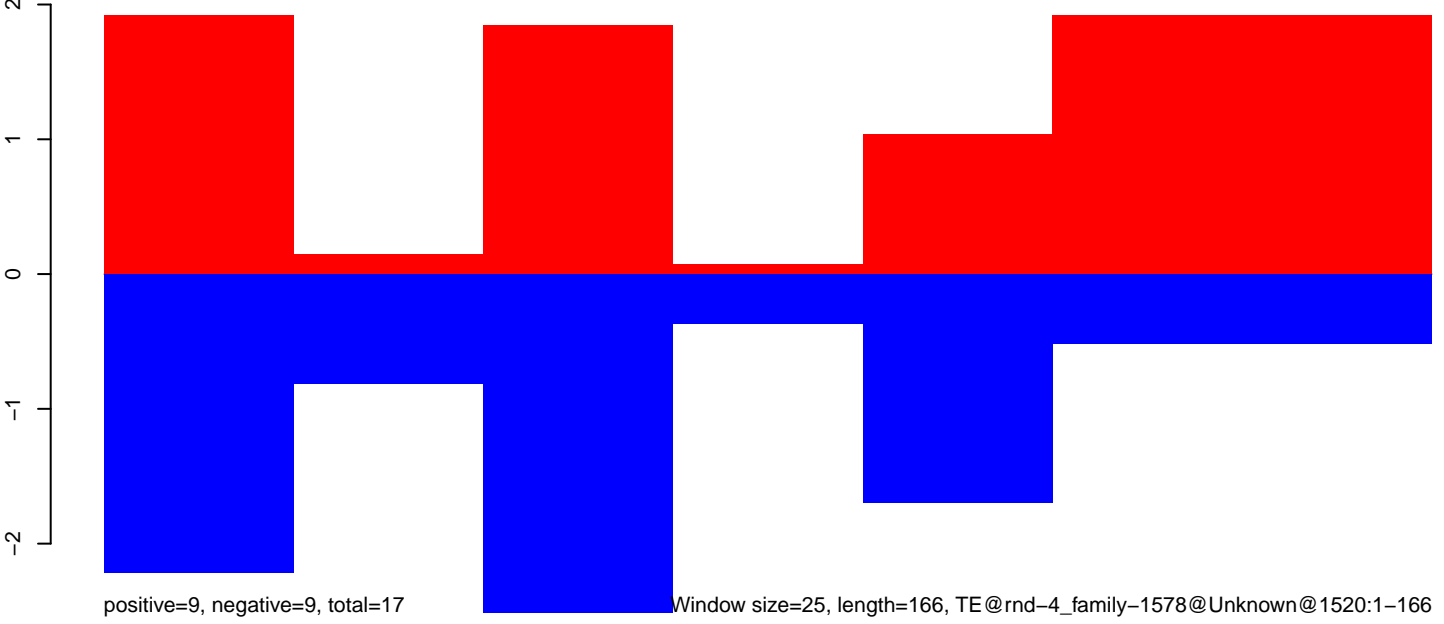
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep

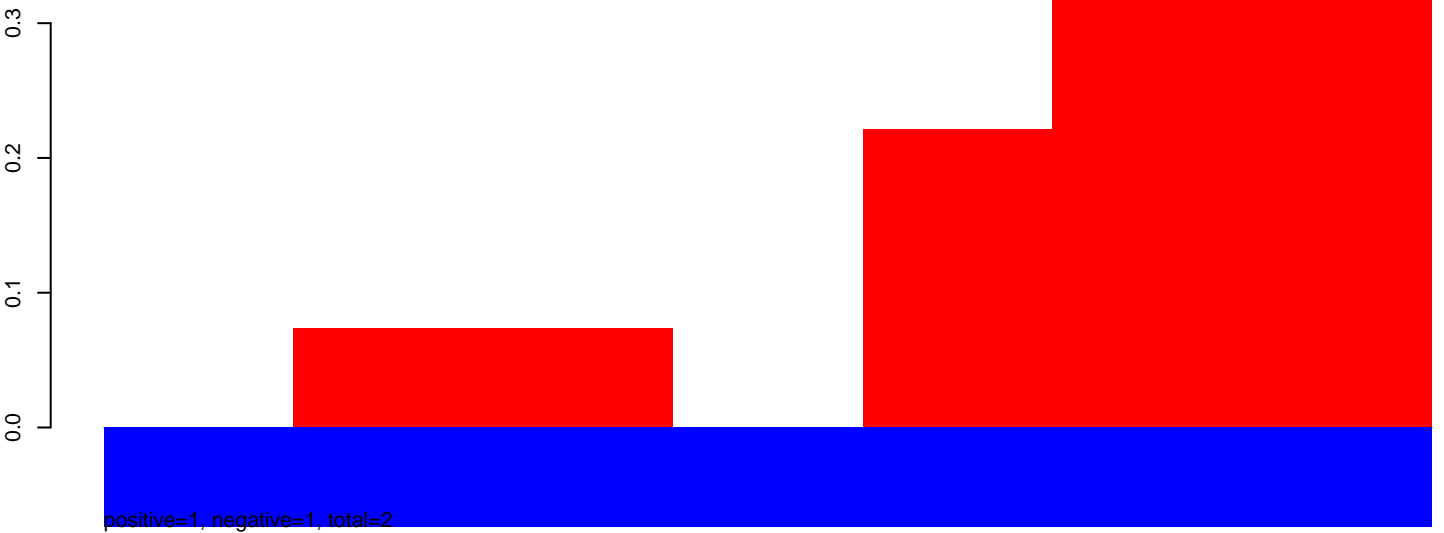


CuQuin_Testes_NL.rep

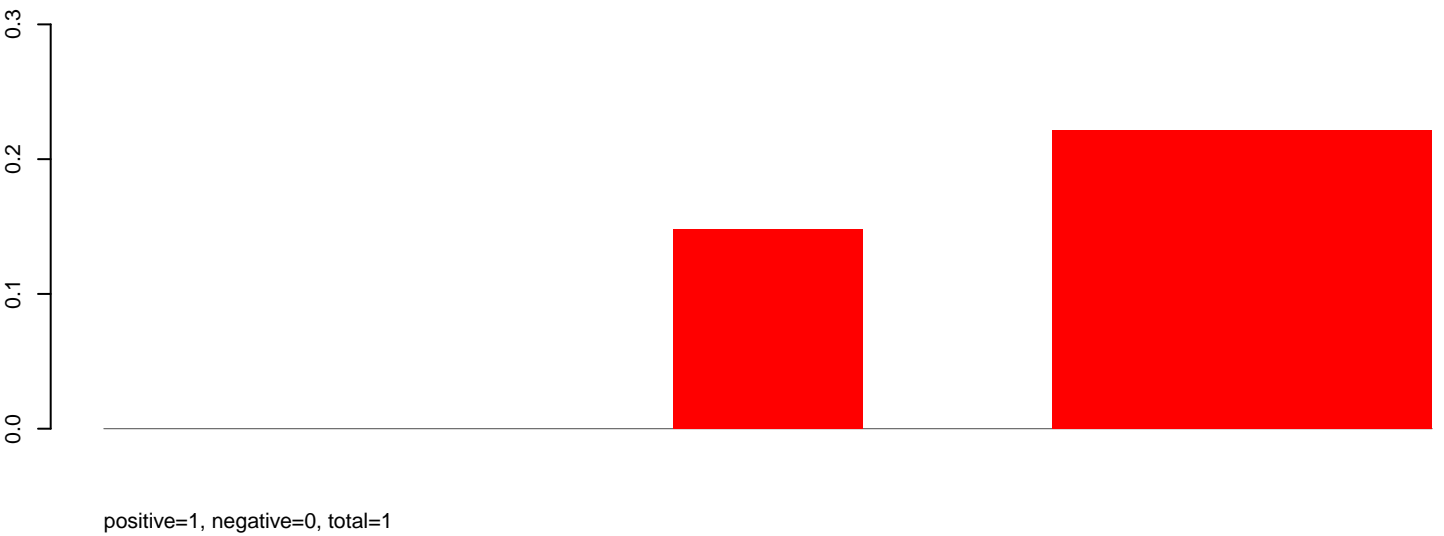


Window size=25, length=166, TE@rnd-4_family-1578@Unknown@1520:1-166

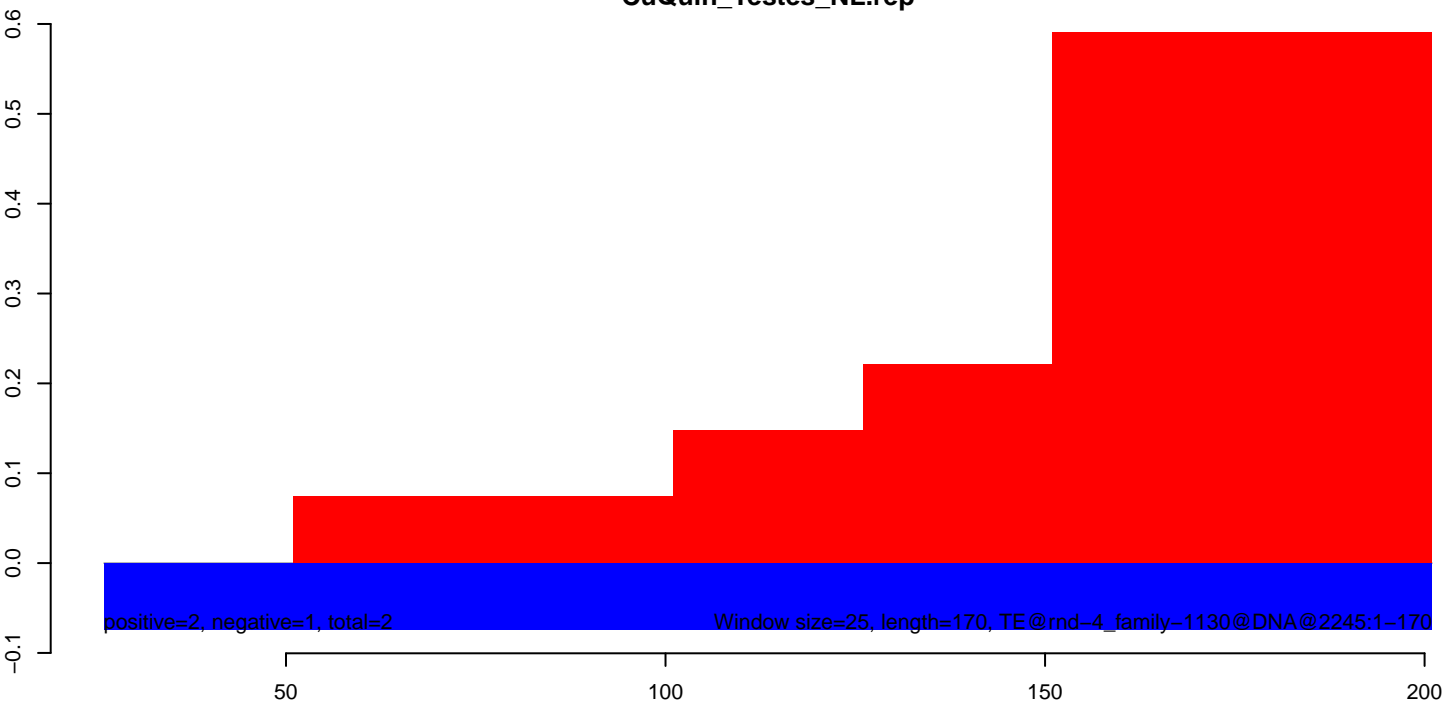
CuQuin_Testes_NL.18_23.rep



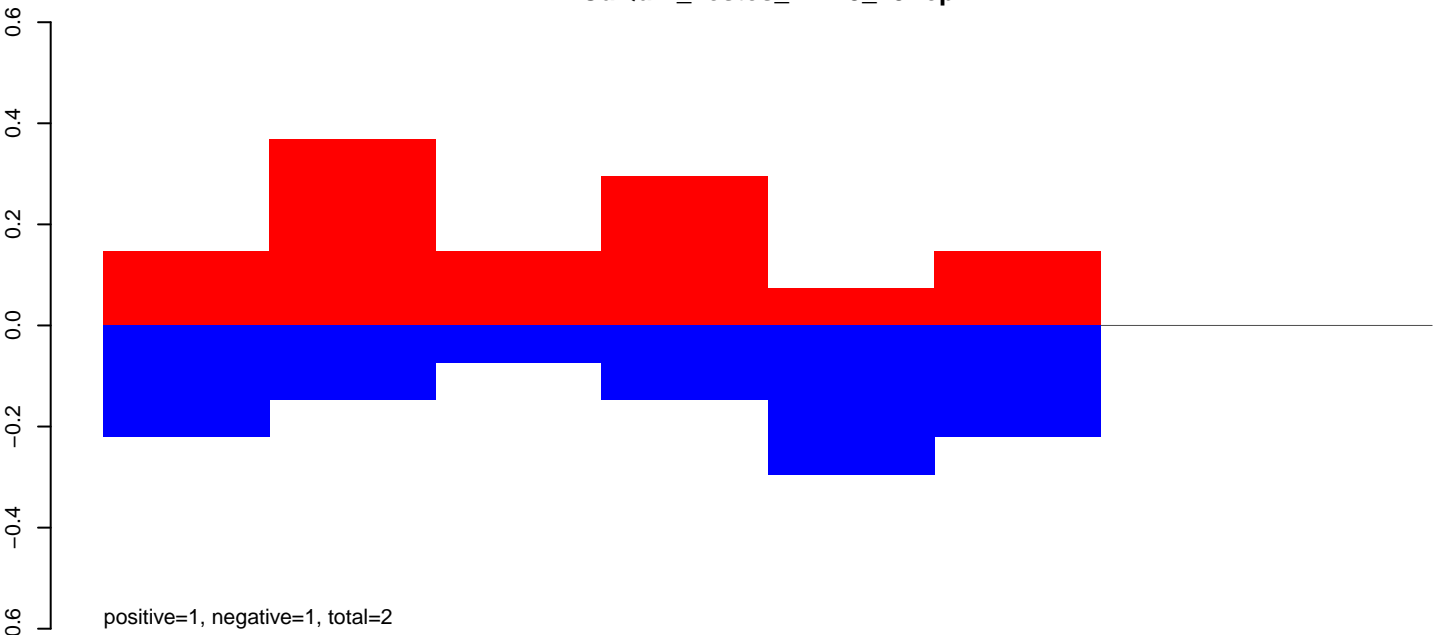
CuQuin_Testes_NL.24_35.rep



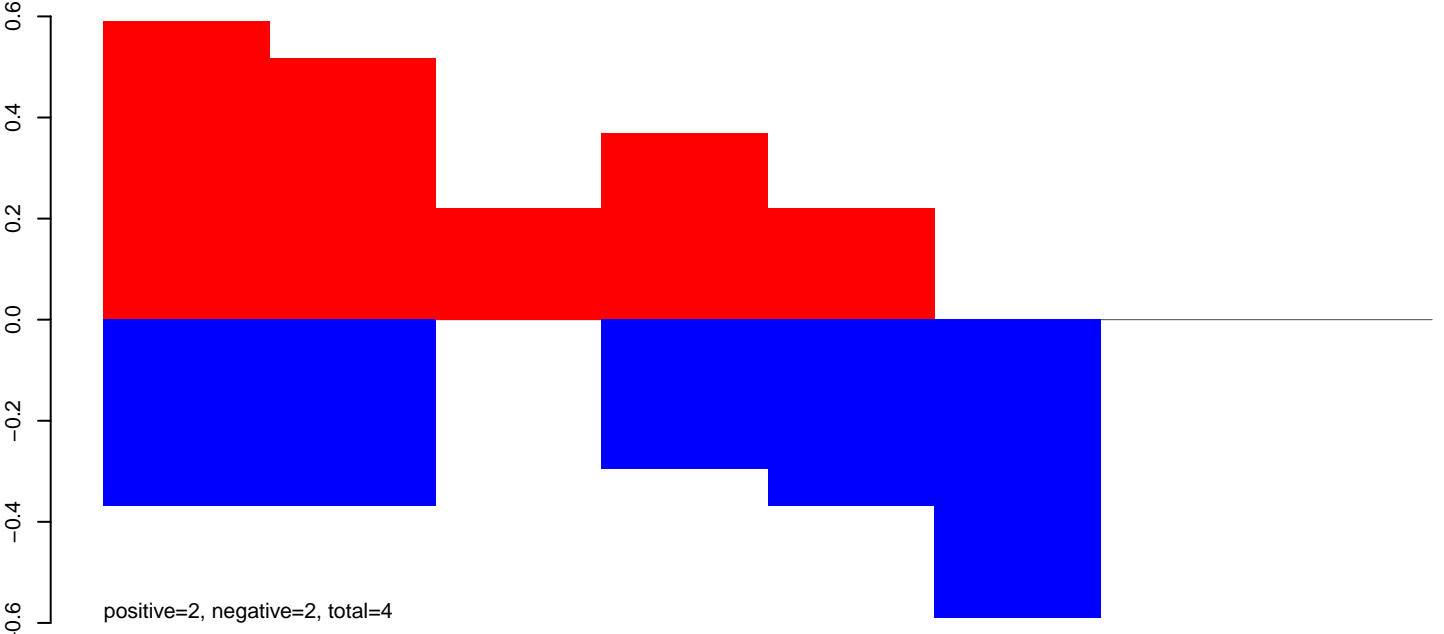
CuQuin_Testes_NL.rep



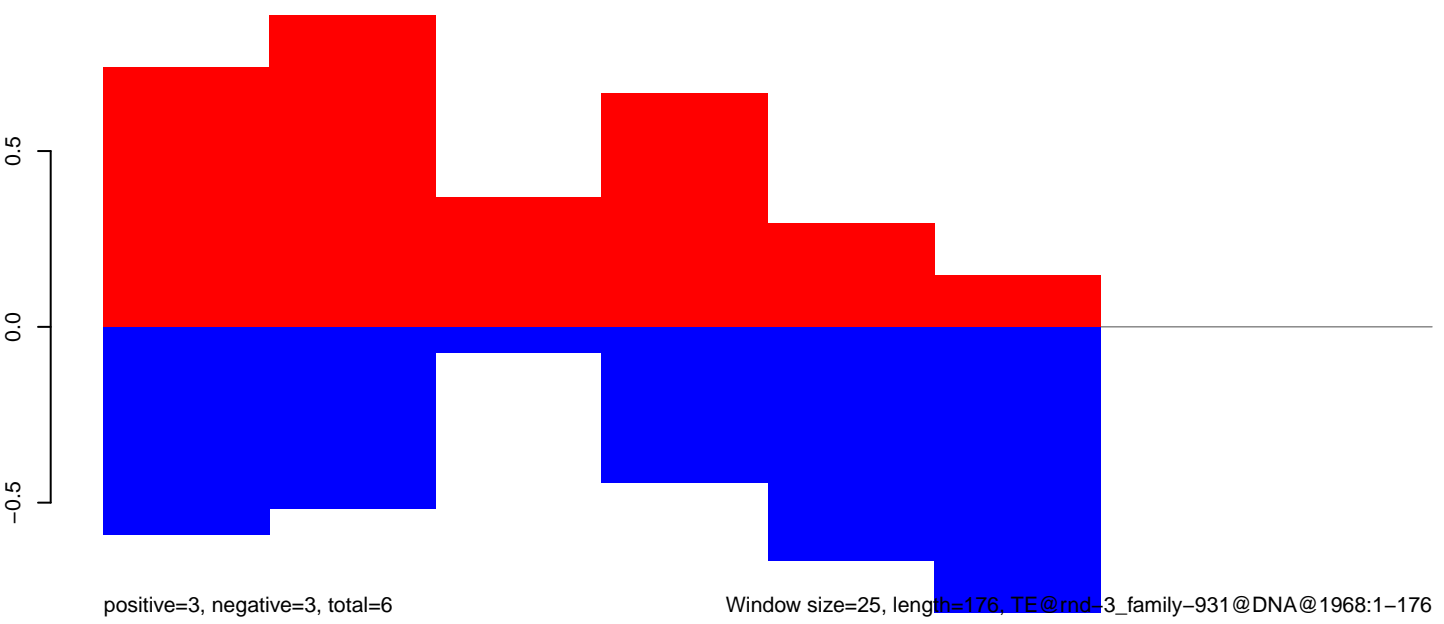
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



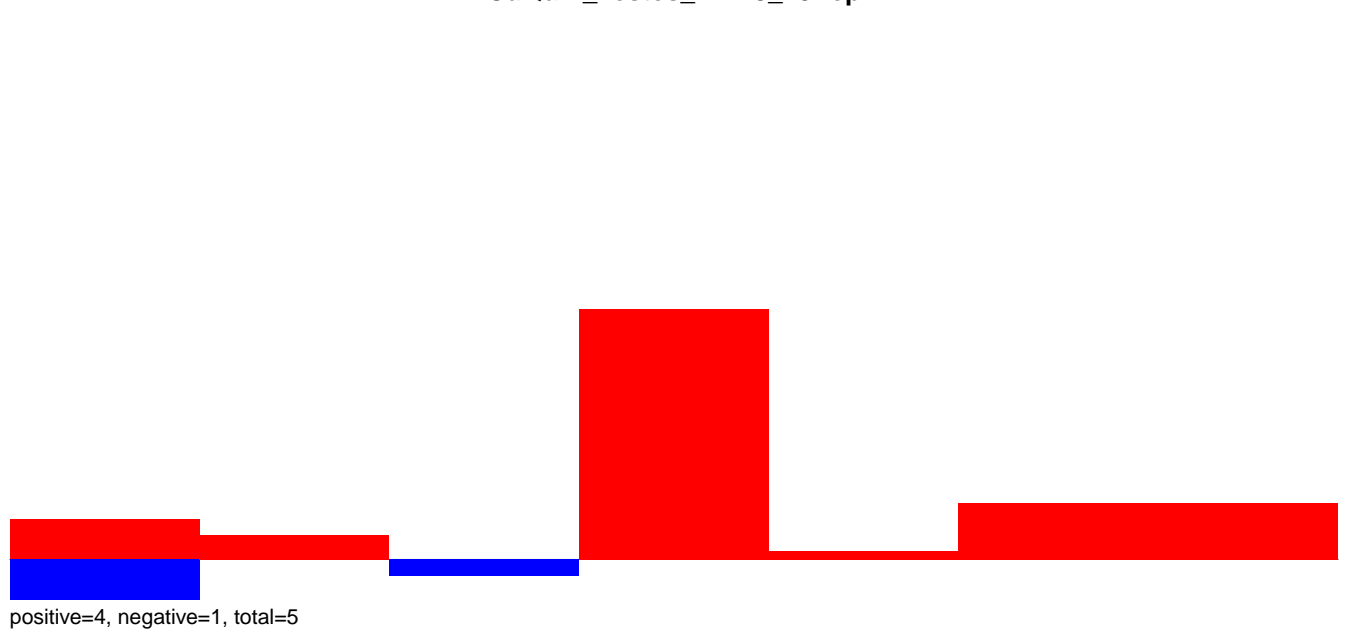
CuQuin_Testes_NL.rep



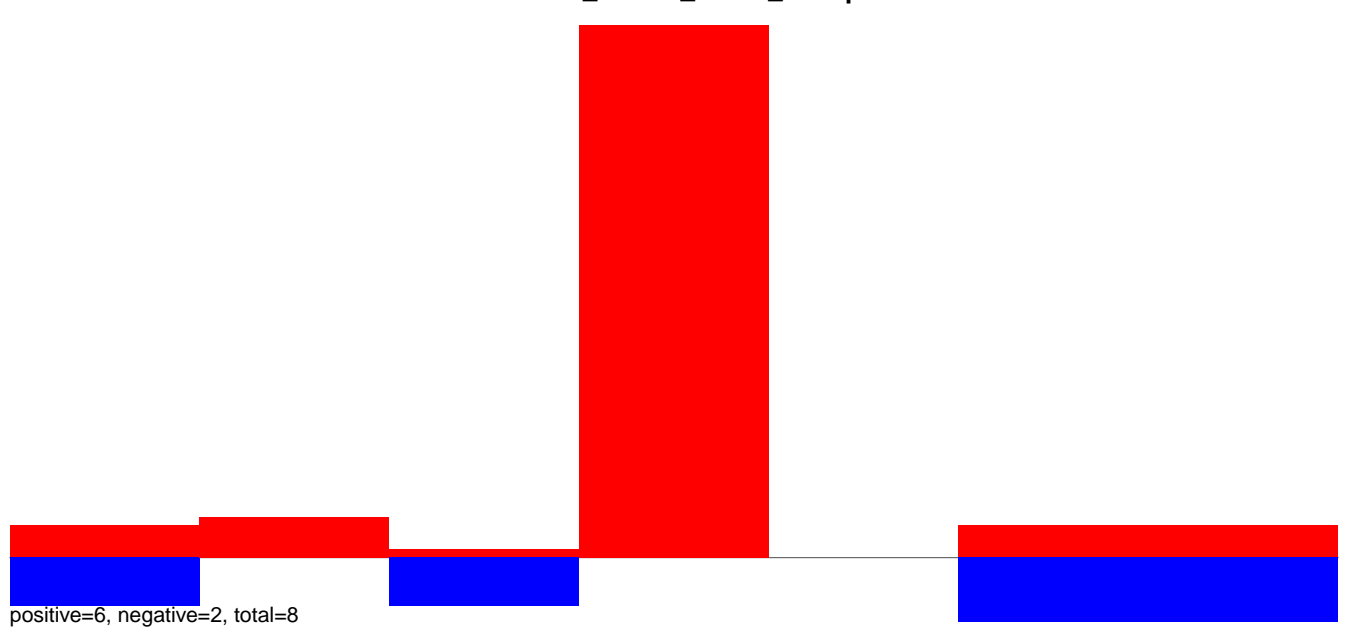
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50 100 150 200

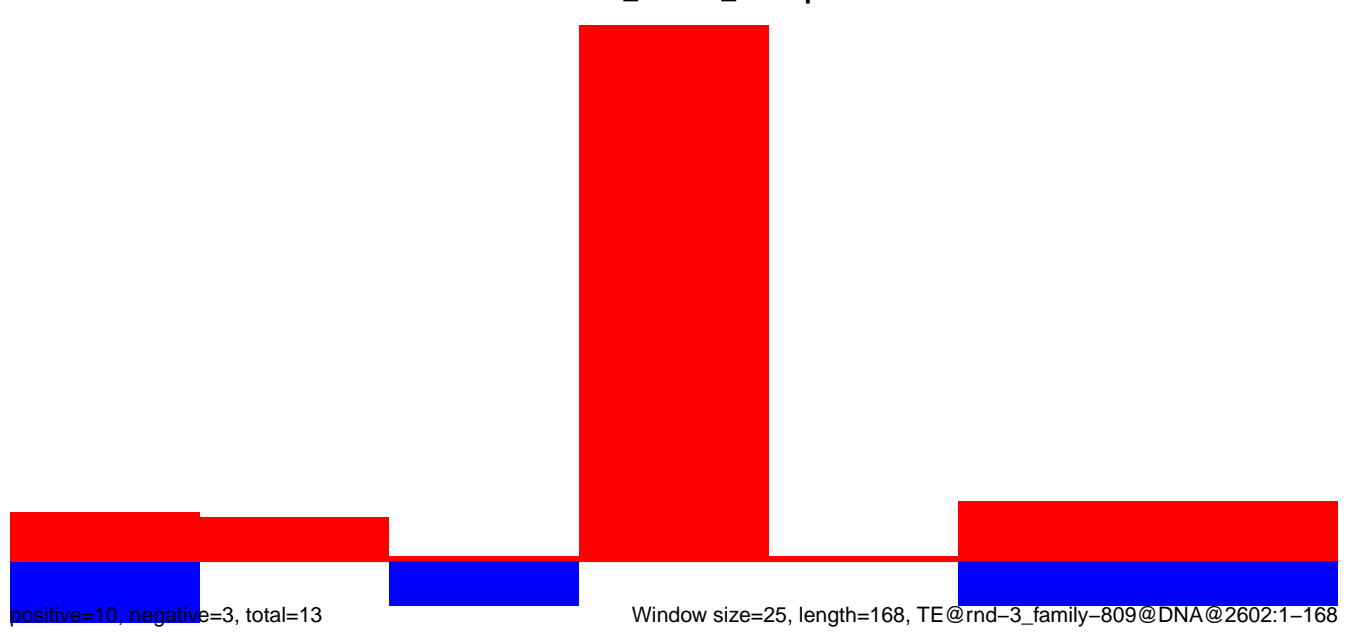
CuQuin_Testes_NL.18_23.rep



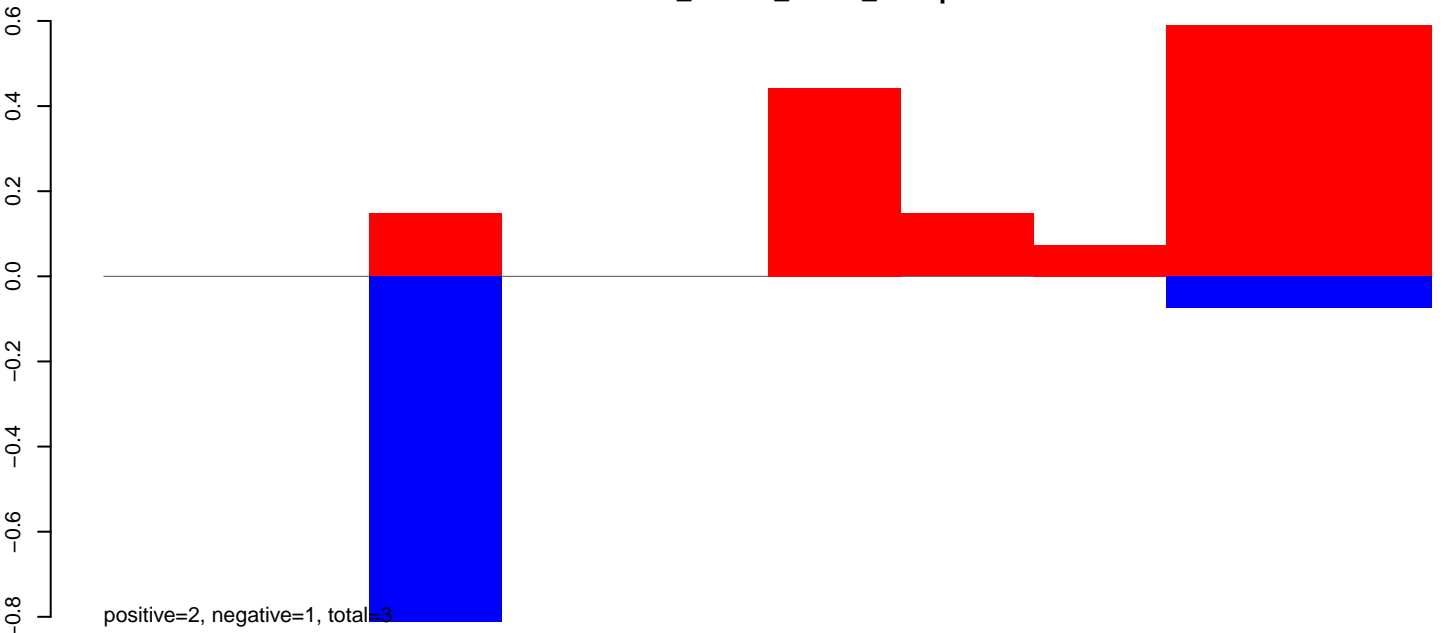
CuQuin_Testes_NL.24_35.rep



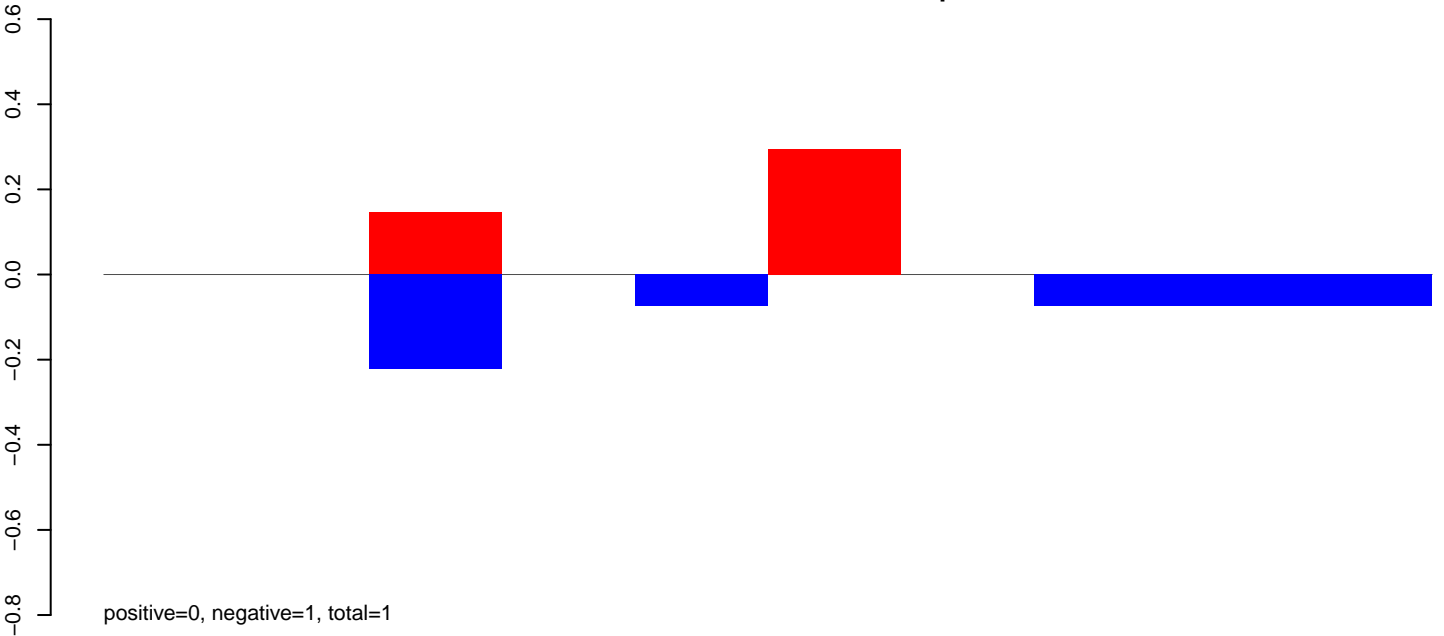
CuQuin_Testes_NL.rep



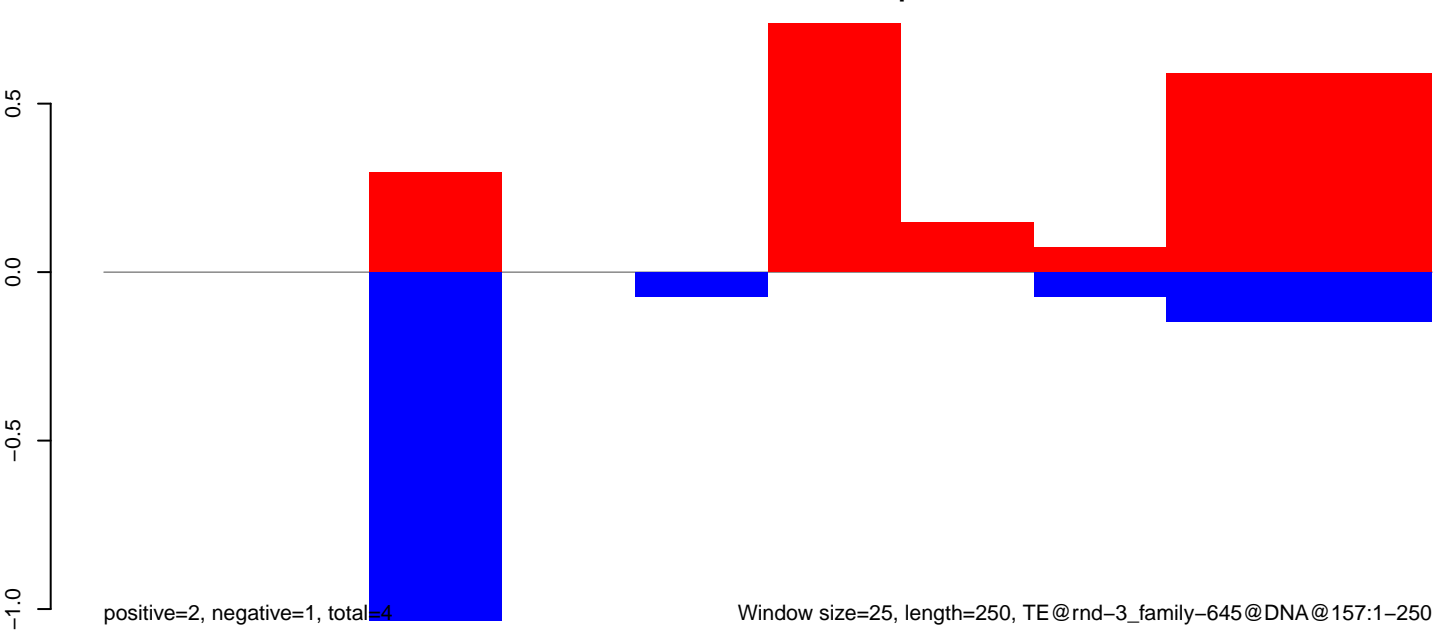
CuQuin_Testes_NL.18_23.rep



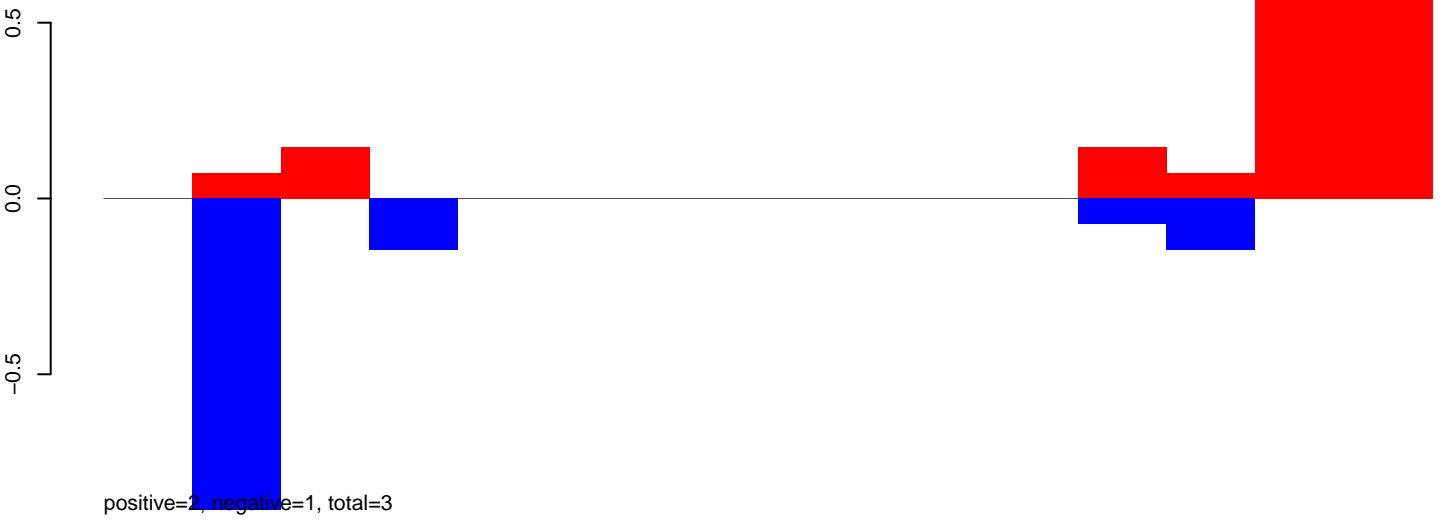
CuQuin_Testes_NL.24_35.rep



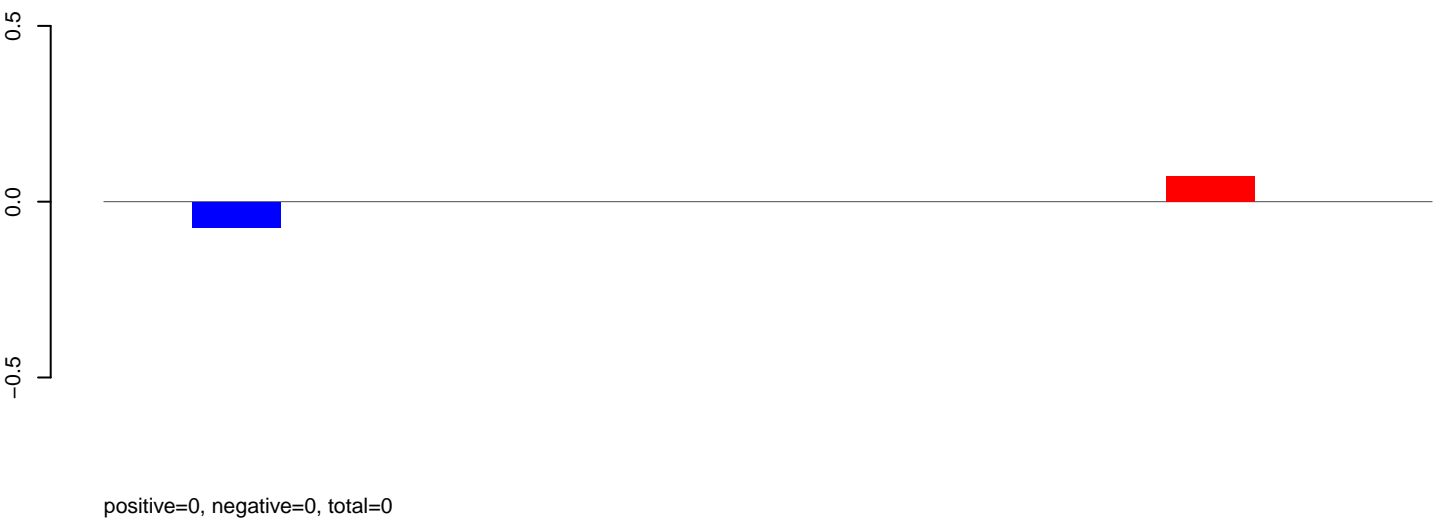
CuQuin_Testes_NL.rep



CuQuin_Testes_NL.18_23.rep



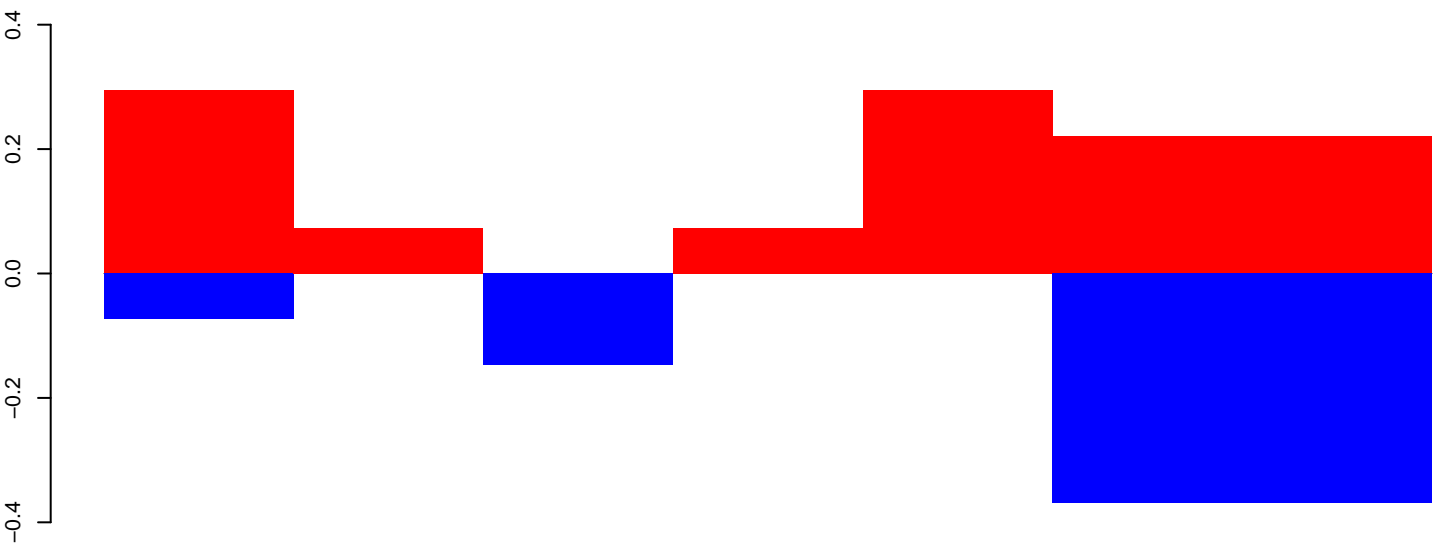
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

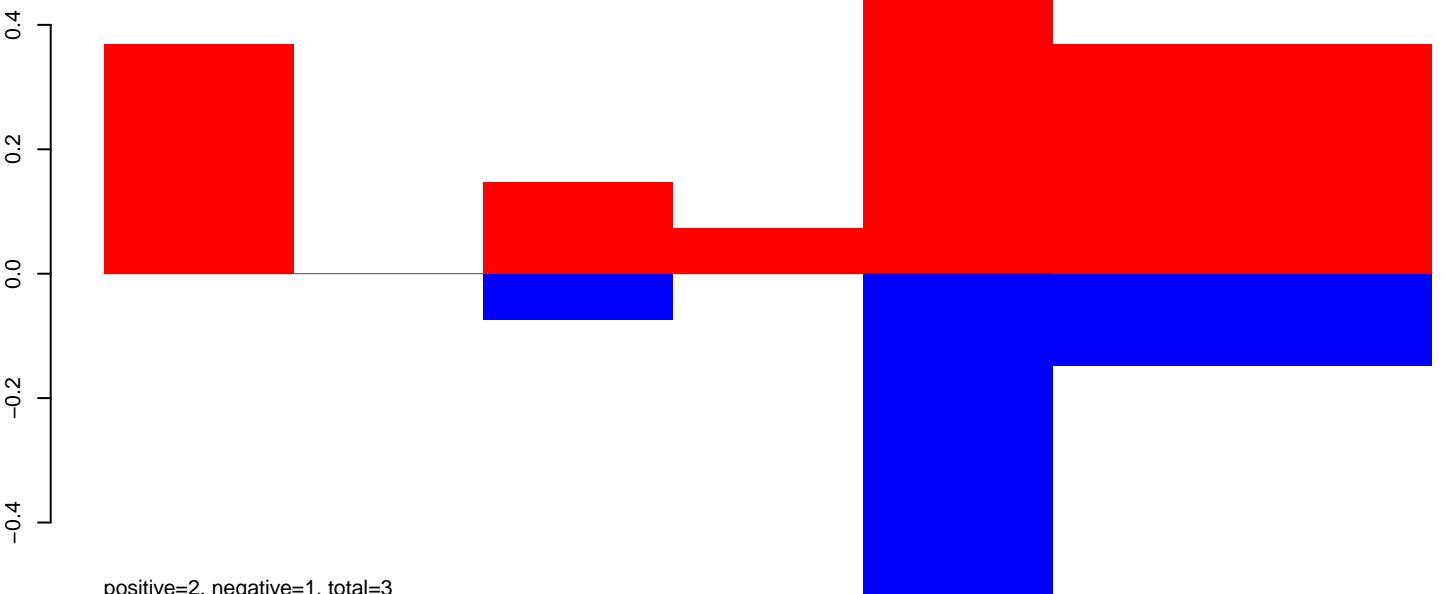


CuQuin_Testes_NL.18_23.rep



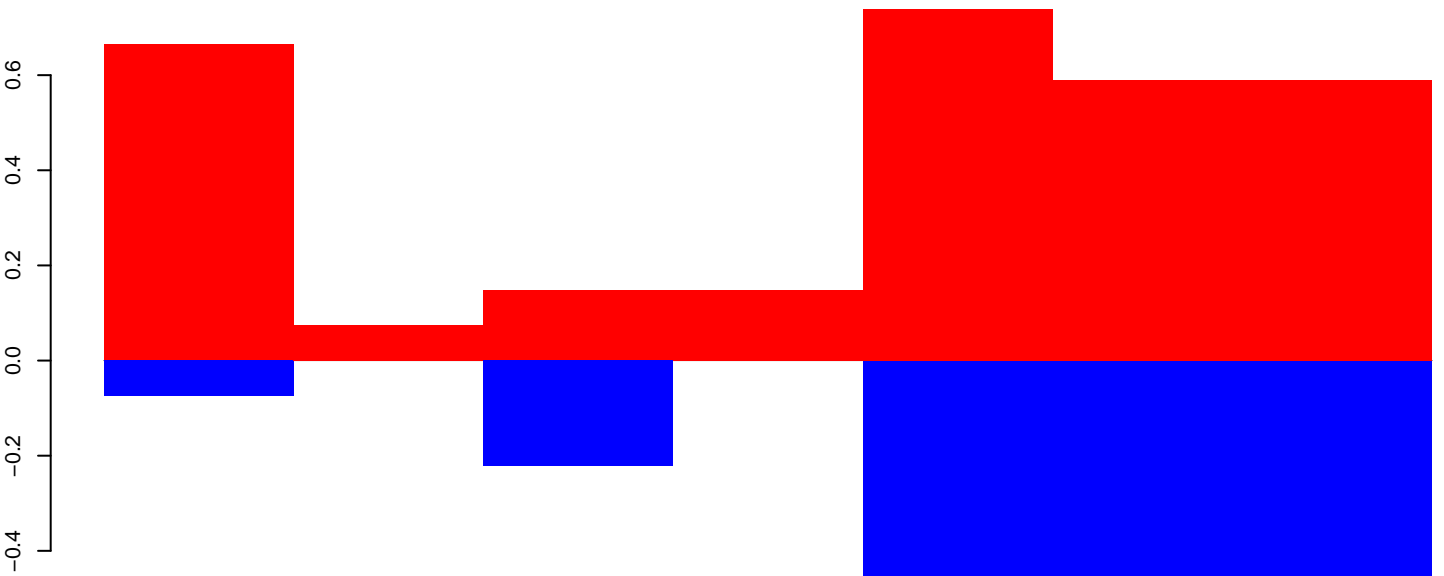
positive=1, negative=1, total=2

CuQuin_Testes_NL.24_35.rep



positive=2, negative=1, total=3

CuQuin_Testes_NL.rep



positive=3, negative=2, total=5

Window size=25, length=162, TE@rnd-3_family-1019@DNA_TcMar@112:1-162

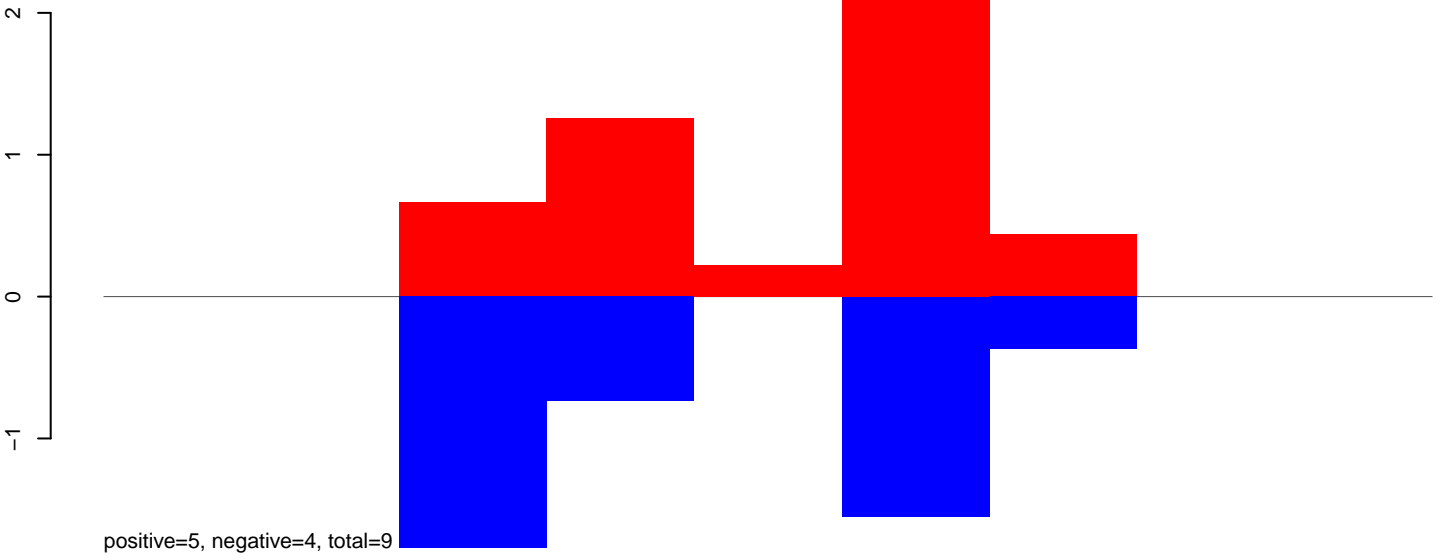
50

100

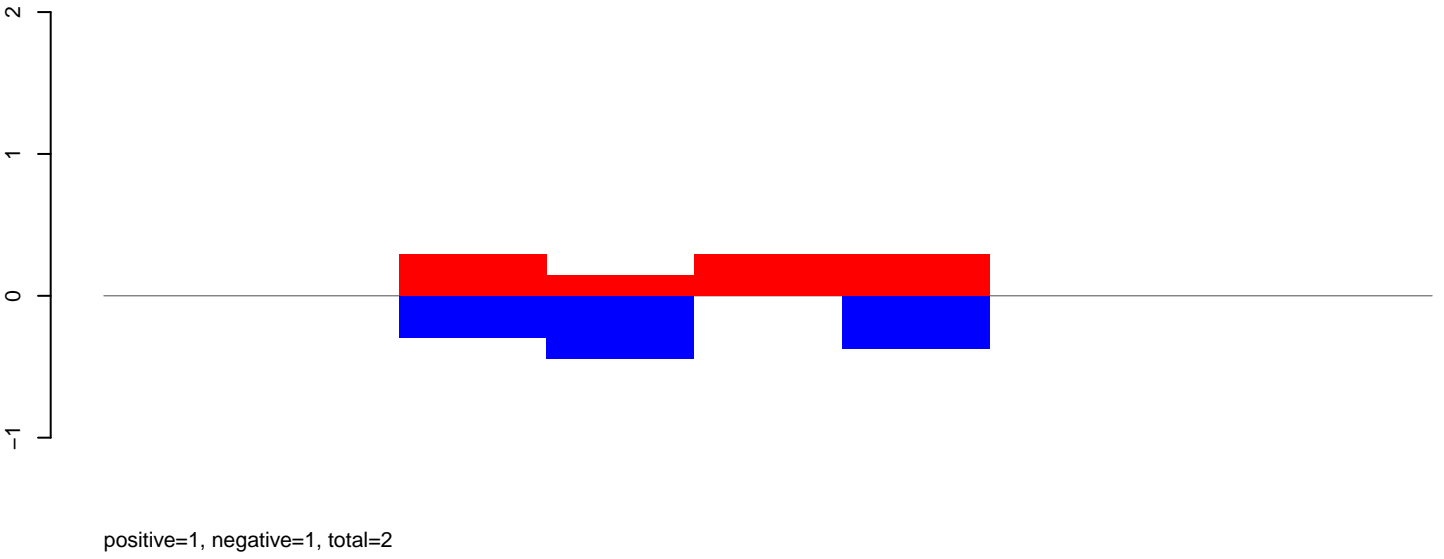
150

200

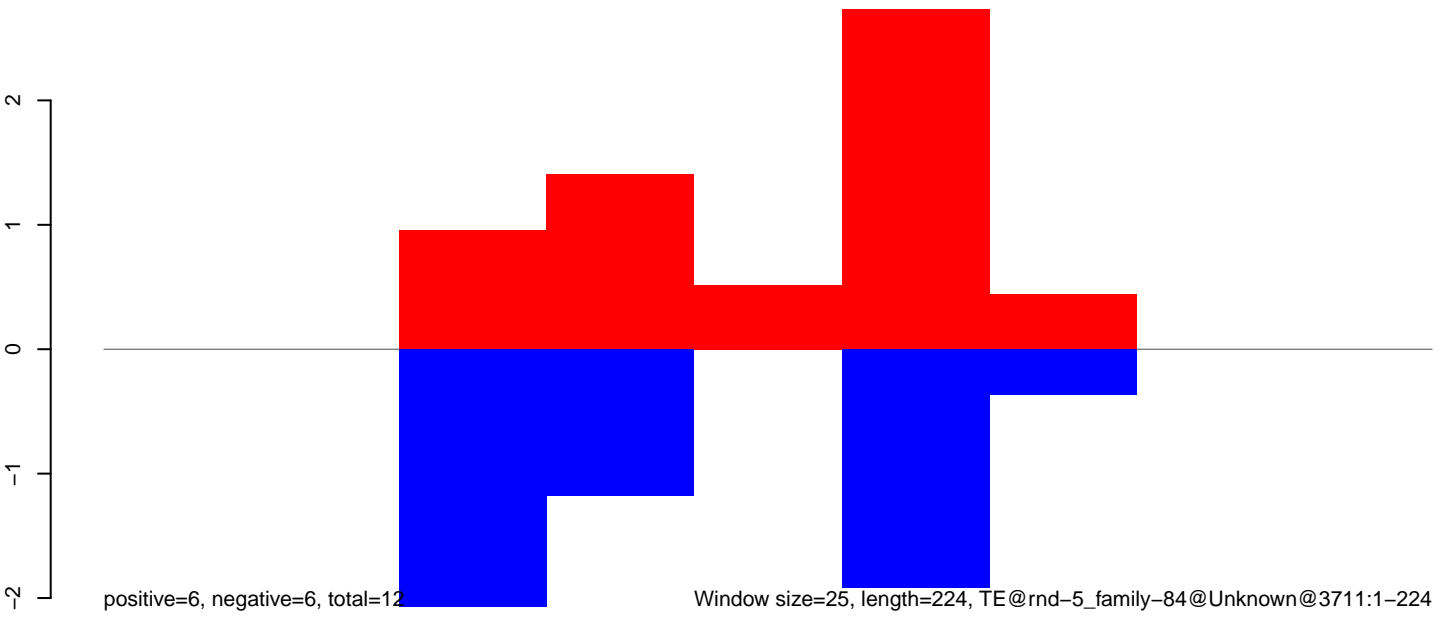
CuQuin_Testes_NL.18_23.rep



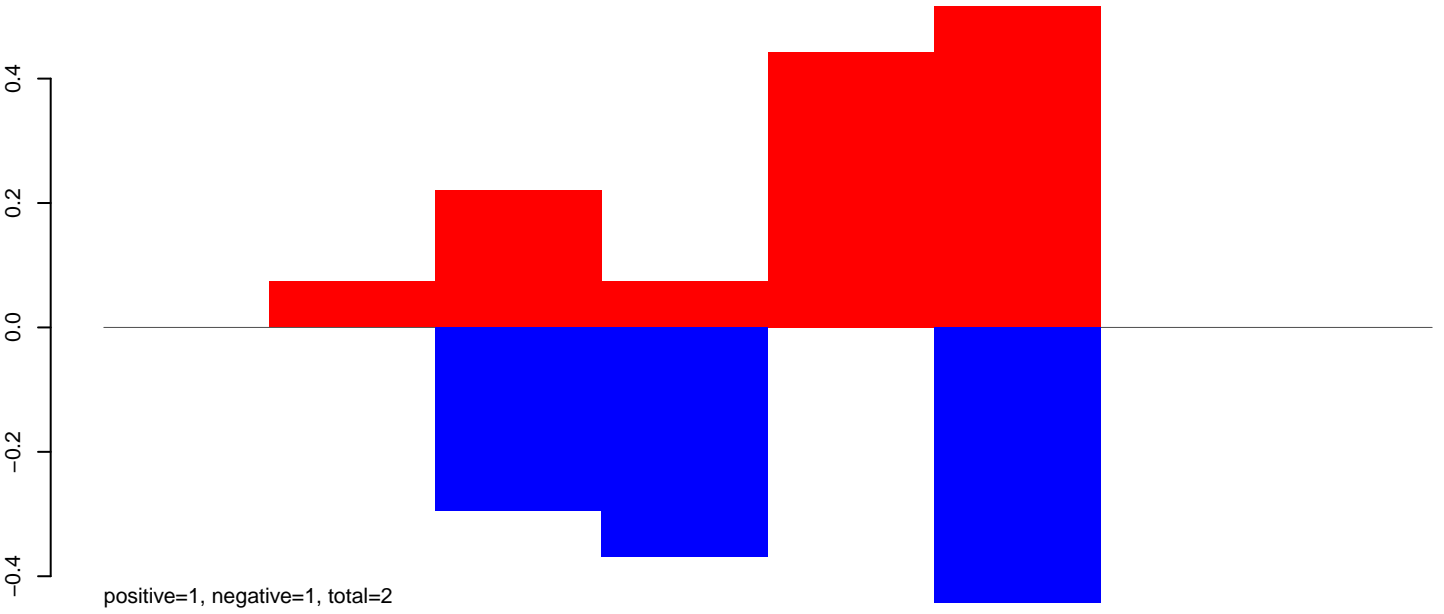
CuQuin_Testes_NL.24_35.rep



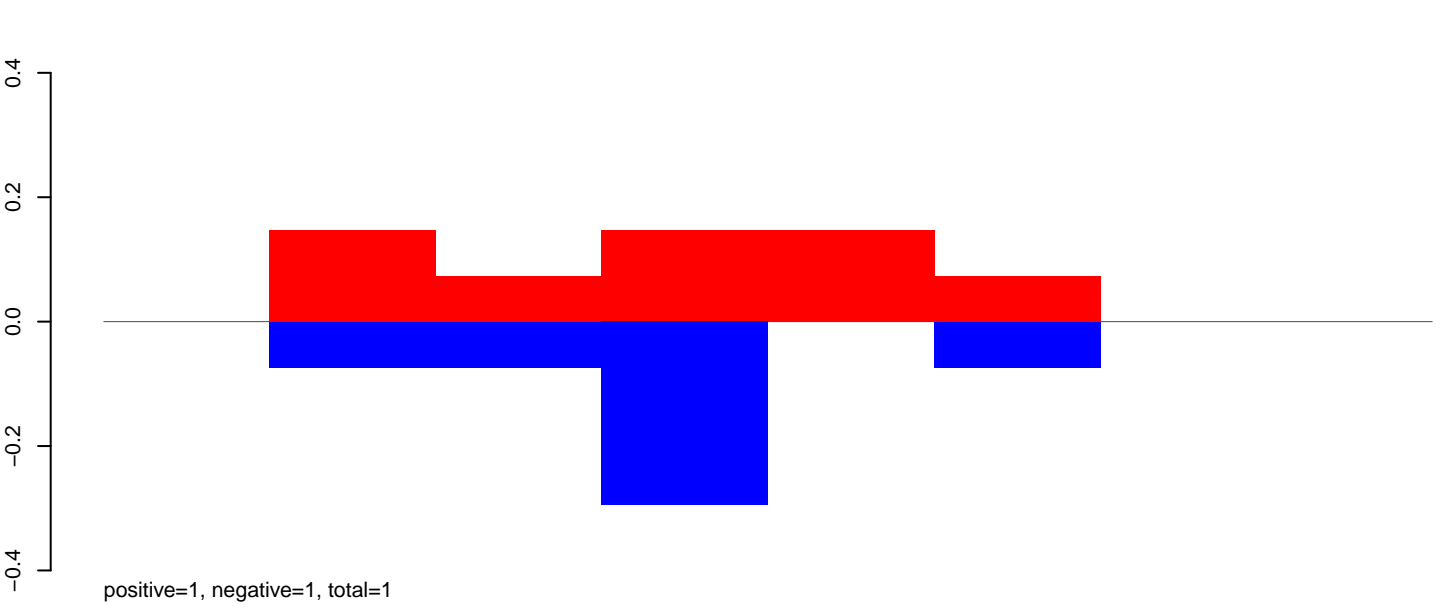
CuQuin_Testes_NL.rep



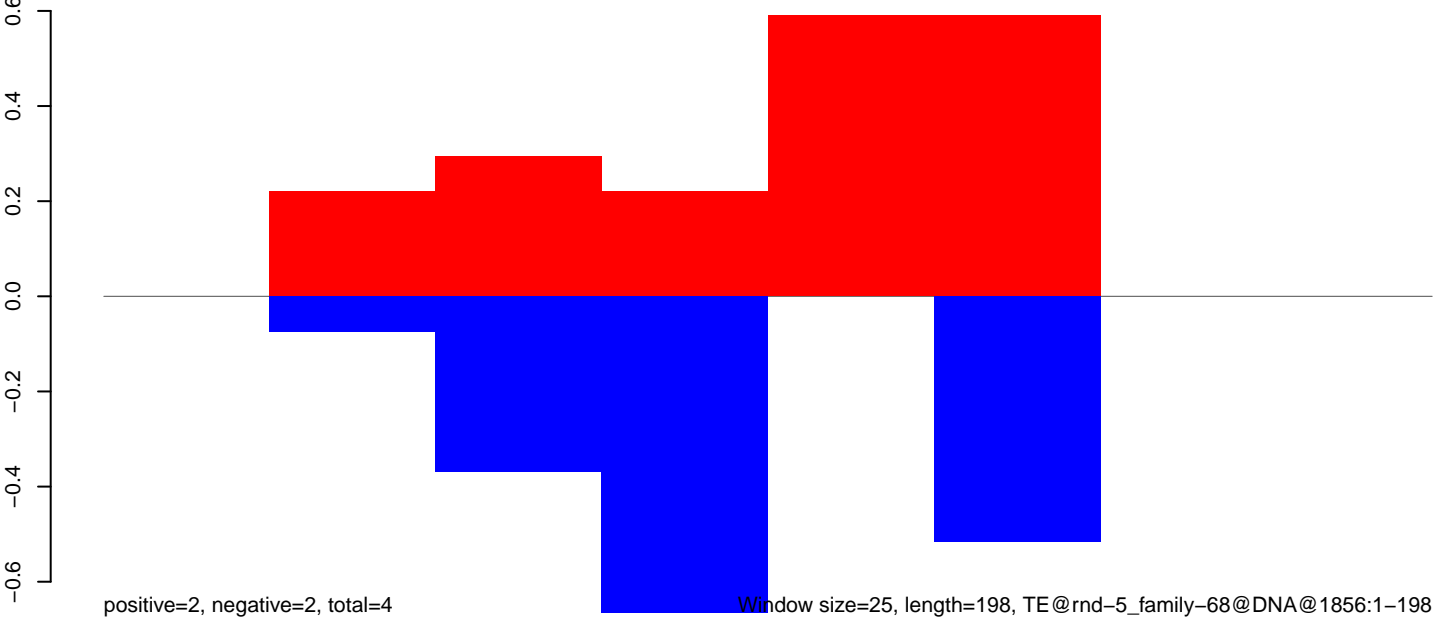
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



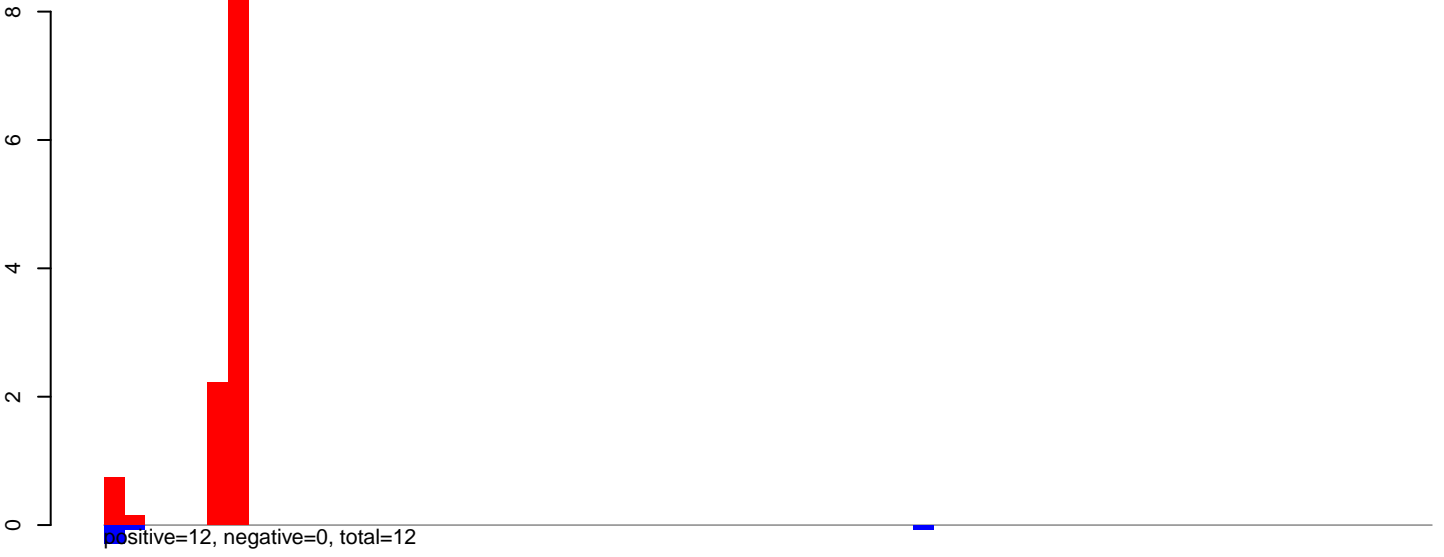
CuQuin_Testes_NL.rep



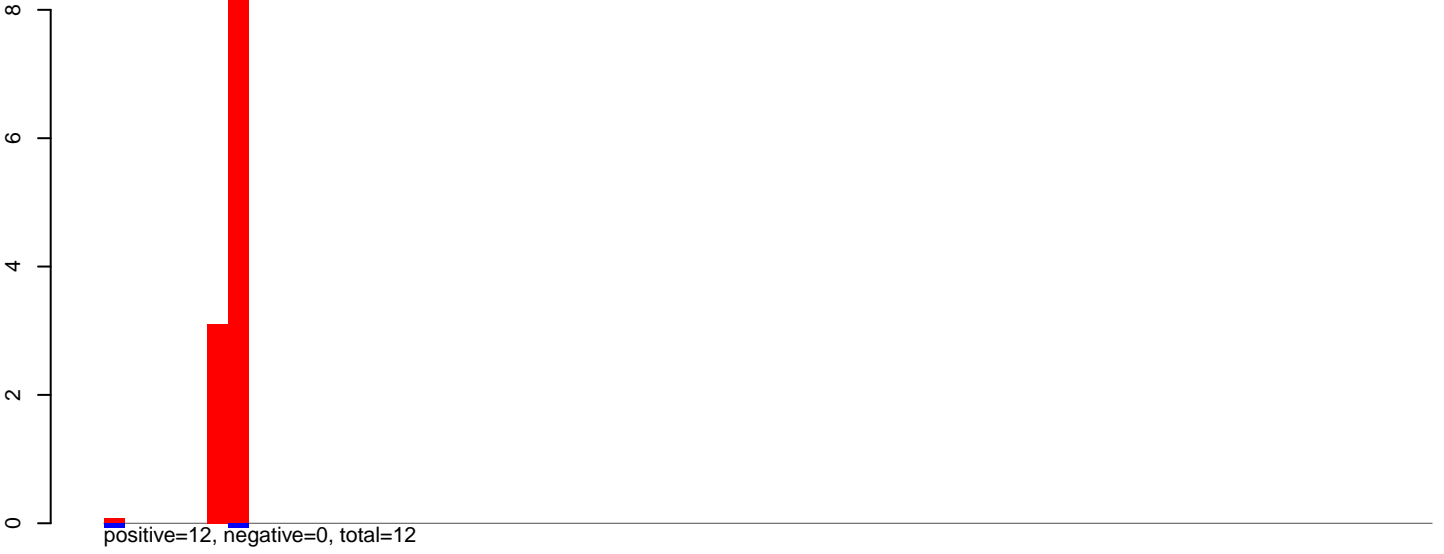
Window size=25, length=198, TE@rnd-5_family-68@DNA@1856:1-198

50 100 150 200

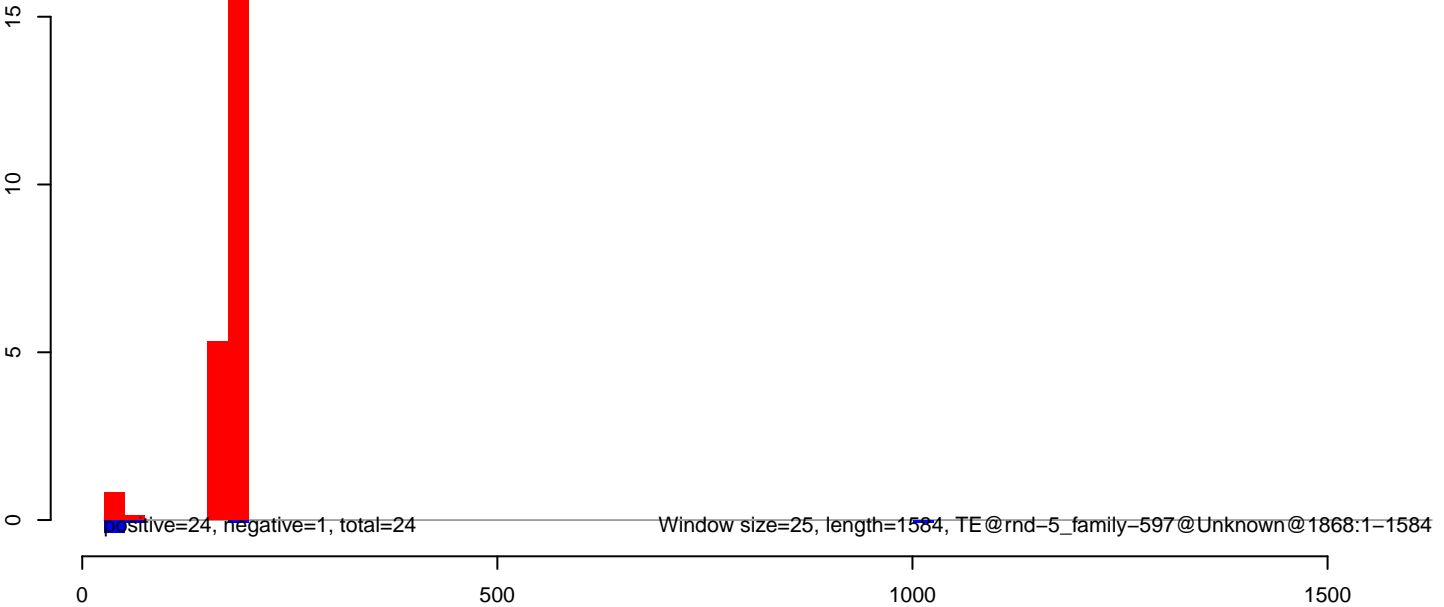
CuQuin_Testes_NL.18_23.rep



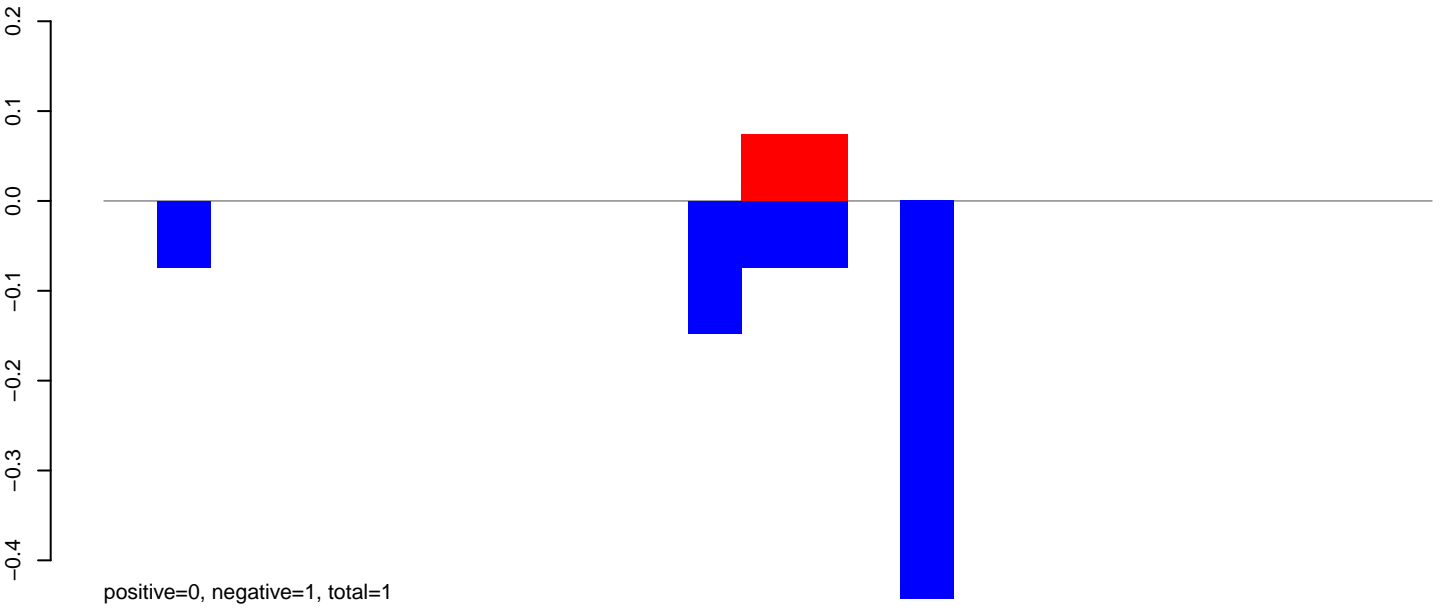
CuQuin_Testes_NL.24_35.rep



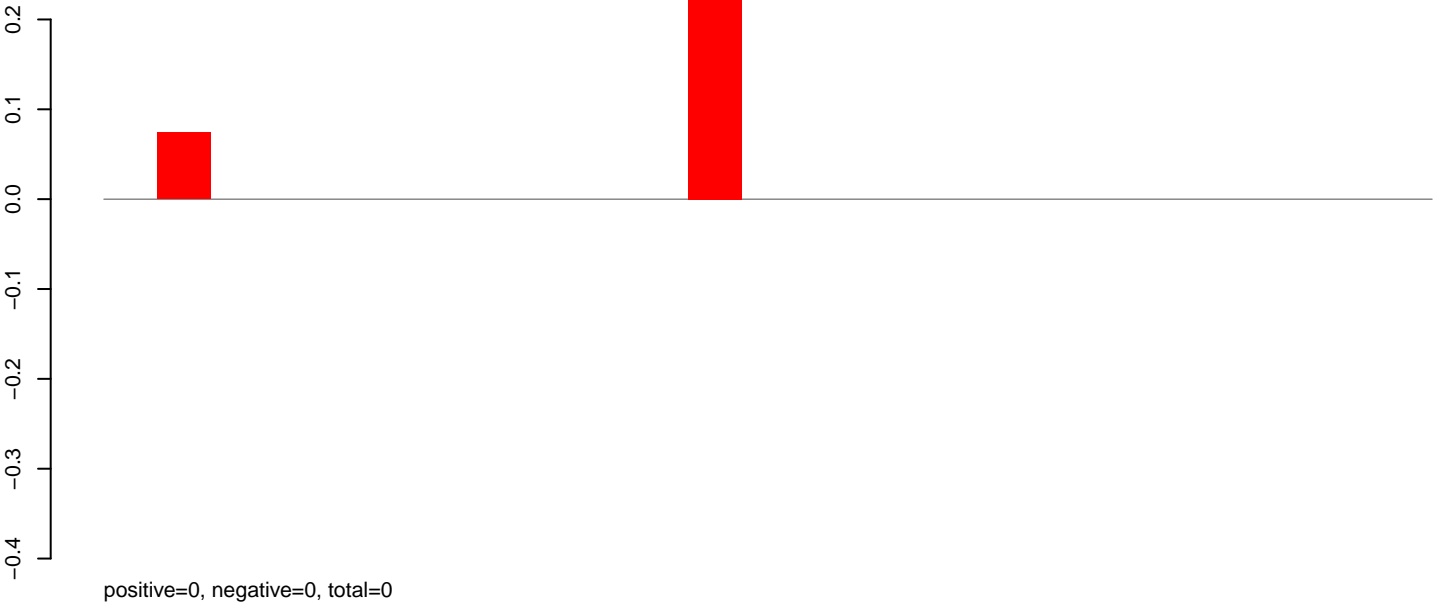
CuQuin_Testes_NL.rep



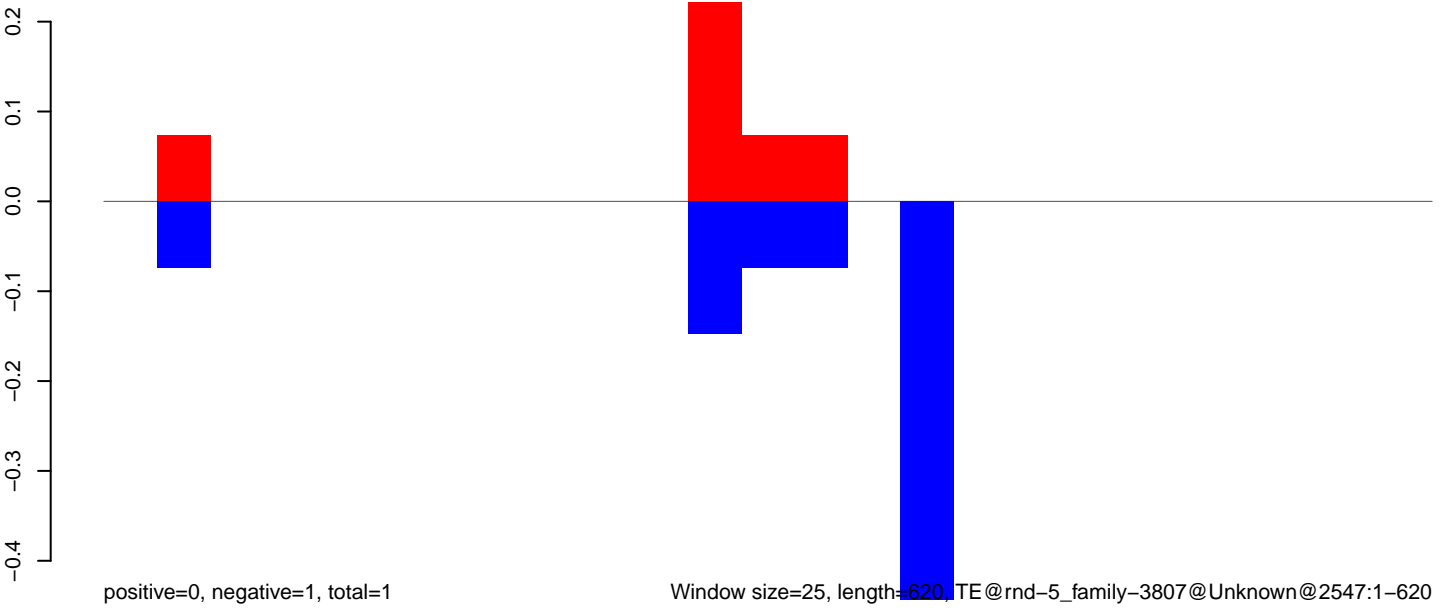
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



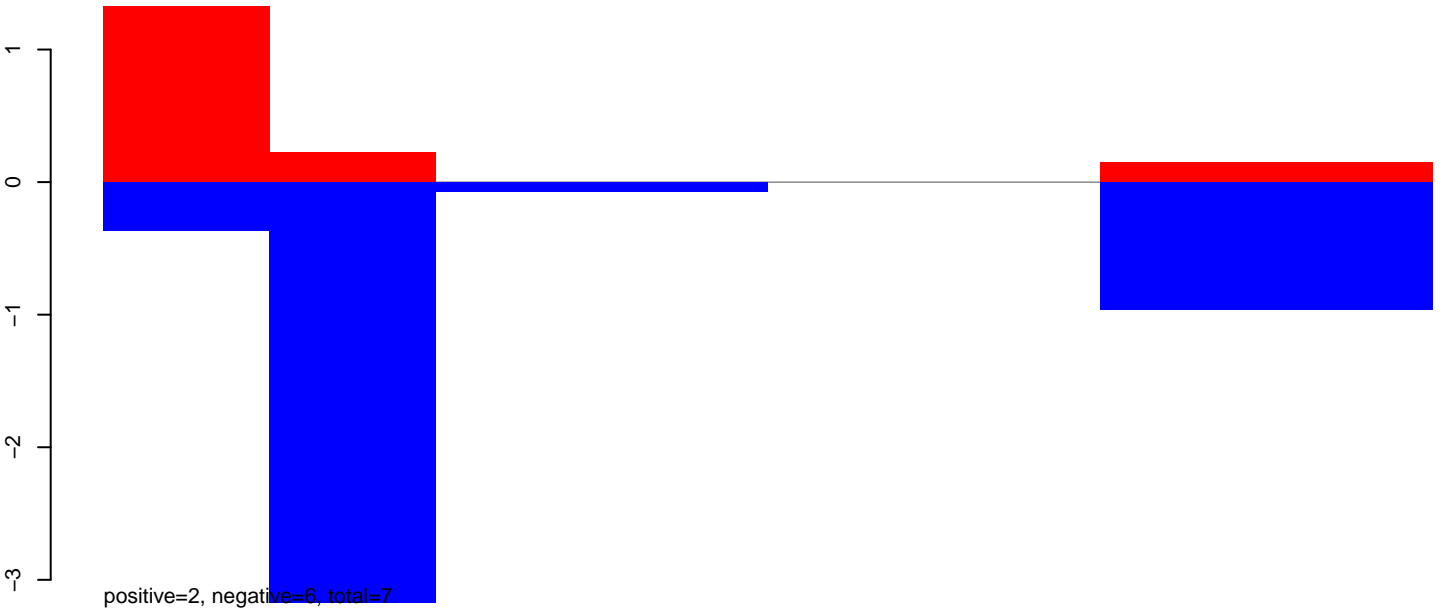
CuQuin_Testes_NL.rep



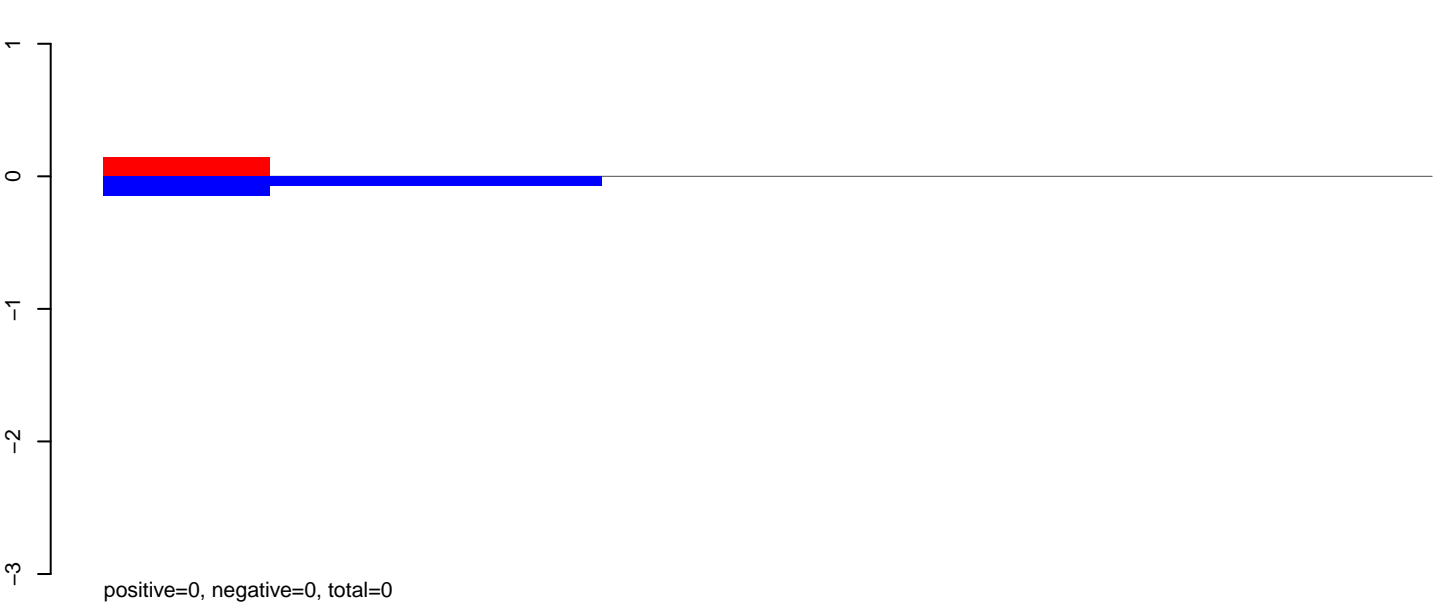
Window size=25, length=620, TE@rnd-5_family-3807@Unknown@2547:1-620

100 200 300 400 500 600

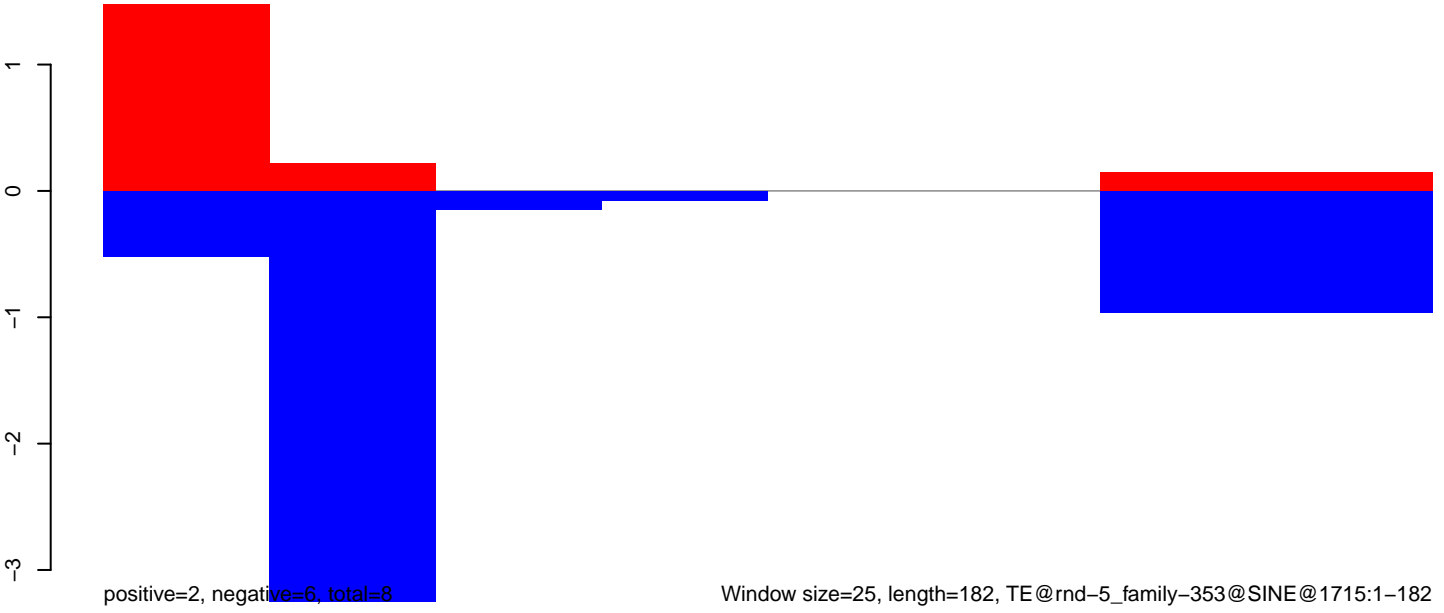
CuQuin_Testes_NL.18_23.rep



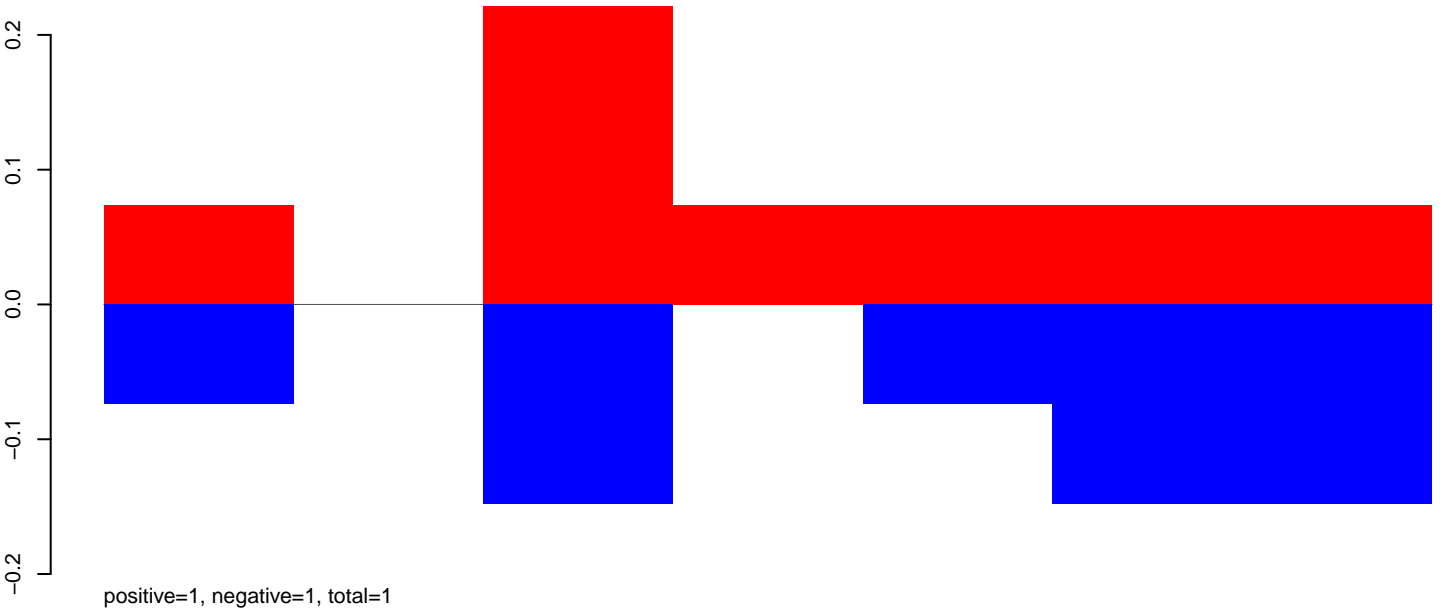
CuQuin_Testes_NL.24_35.rep



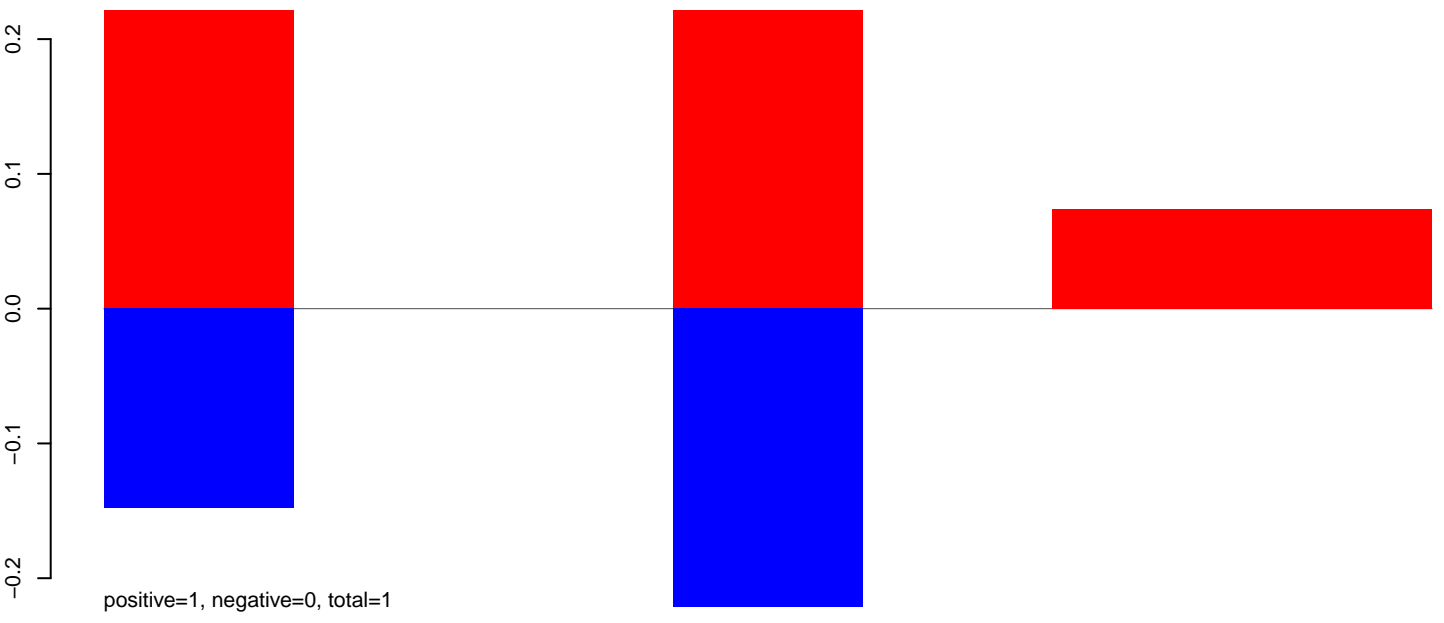
CuQuin_Testes_NL.rep



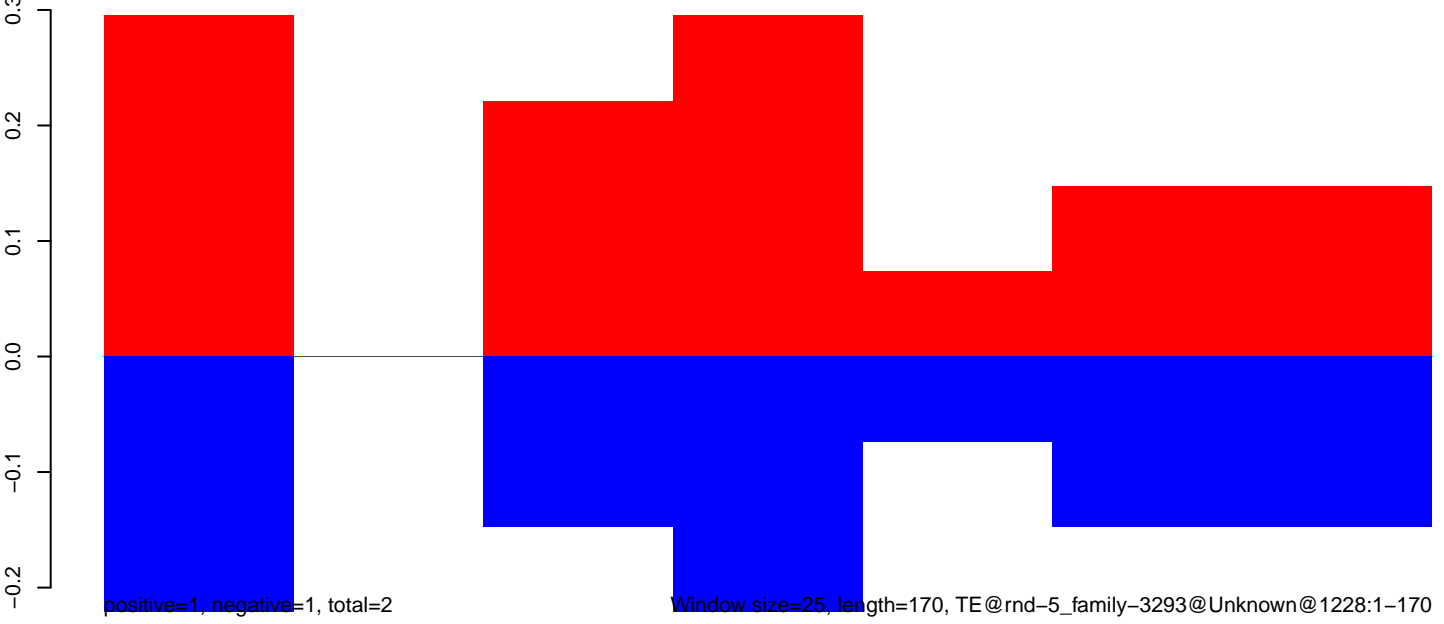
CuQuin_Testes_NL.18_23.rep



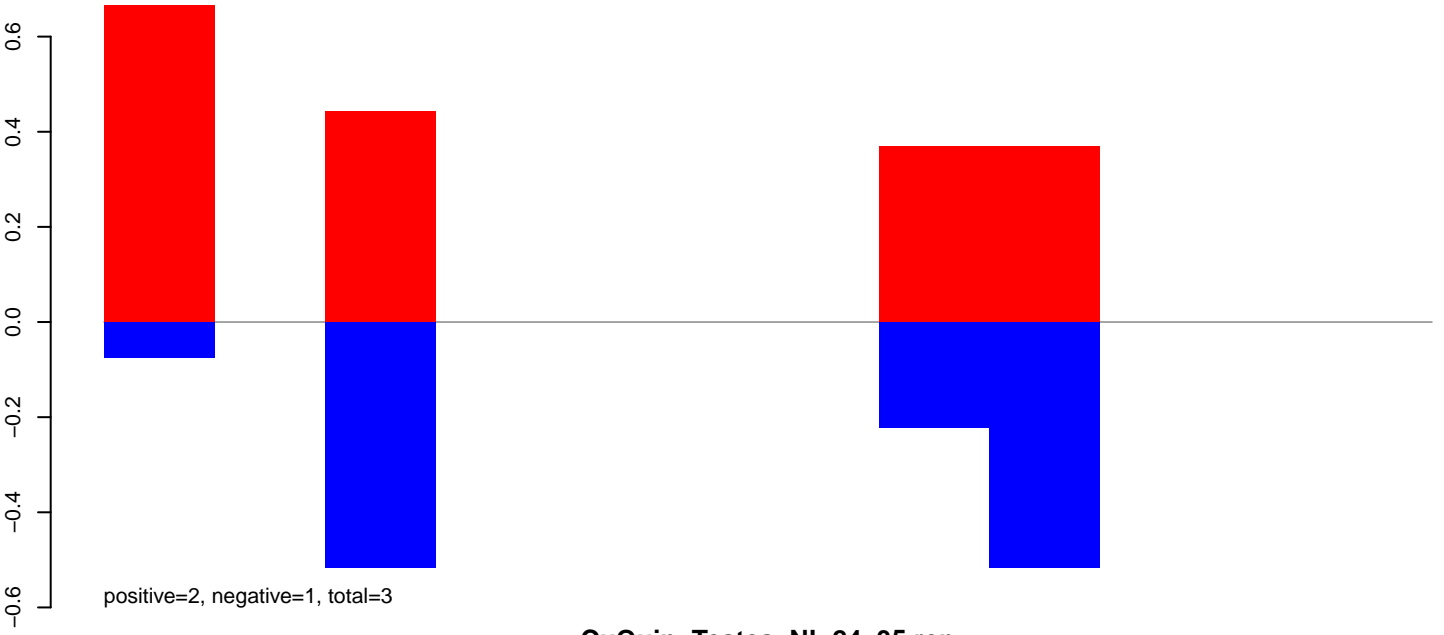
CuQuin_Testes_NL.24_35.rep



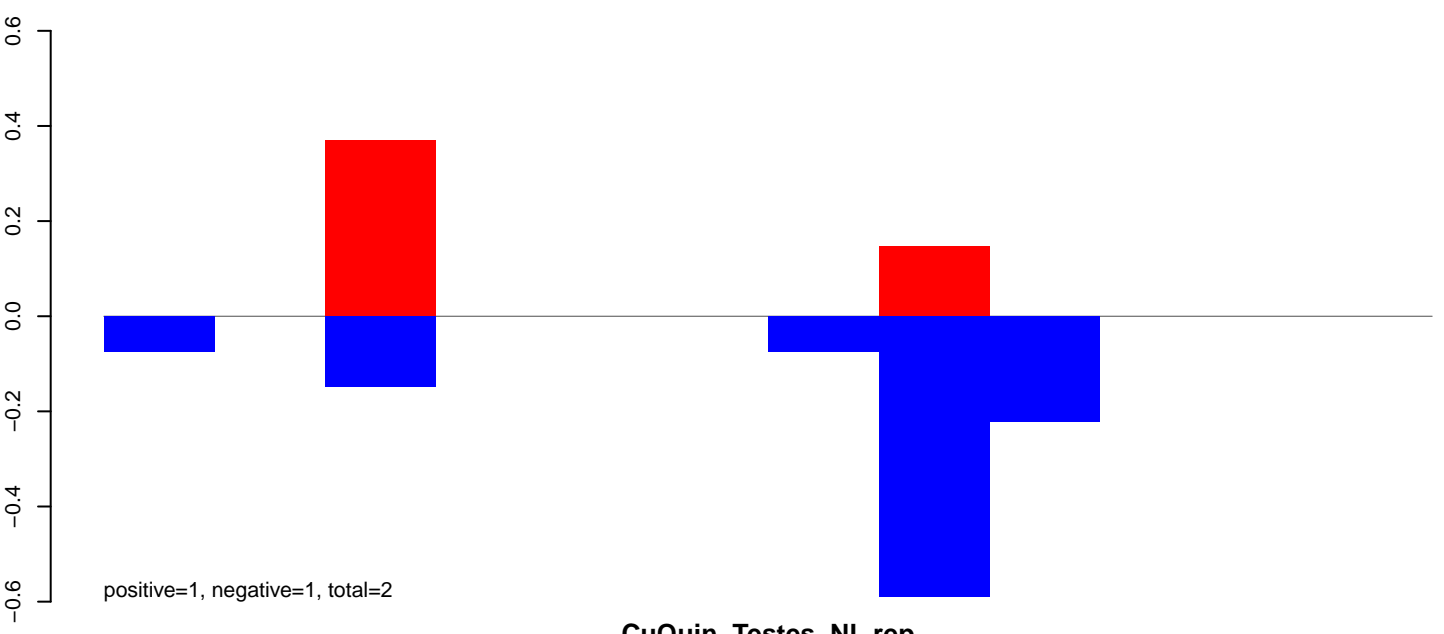
CuQuin_Testes_NL.rep



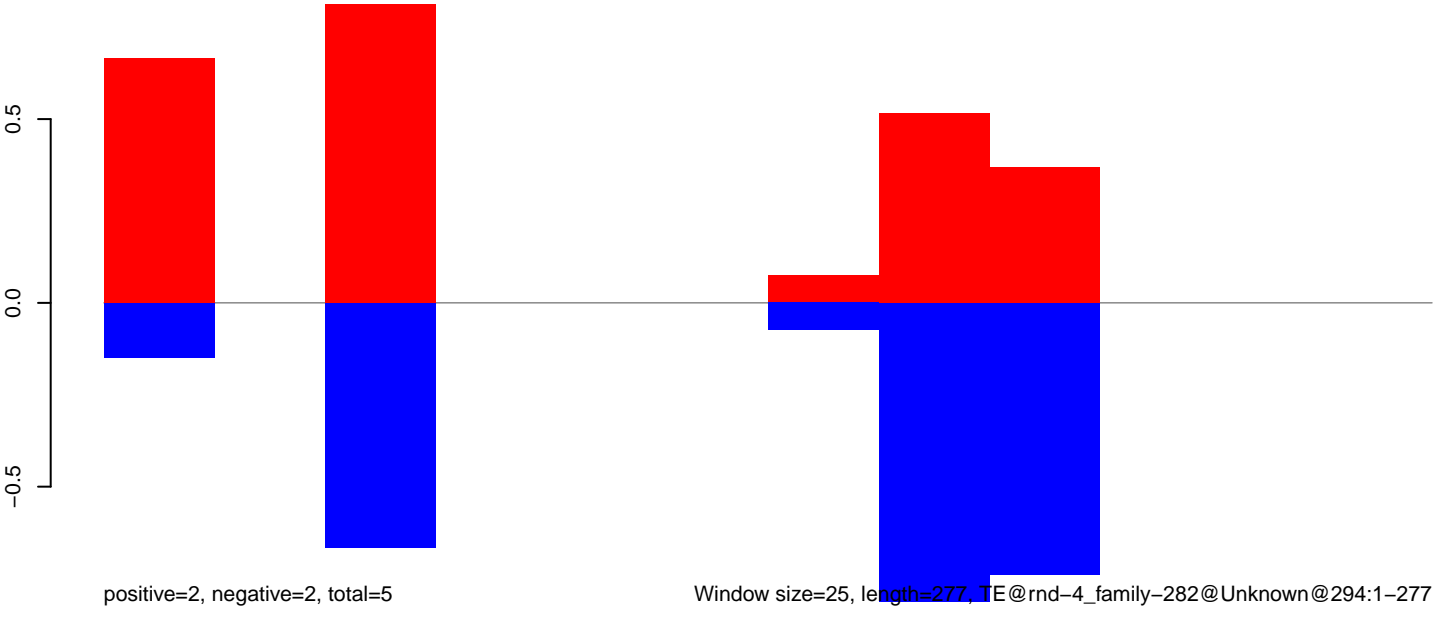
CuQuin_Testes_NL.18_23.rep



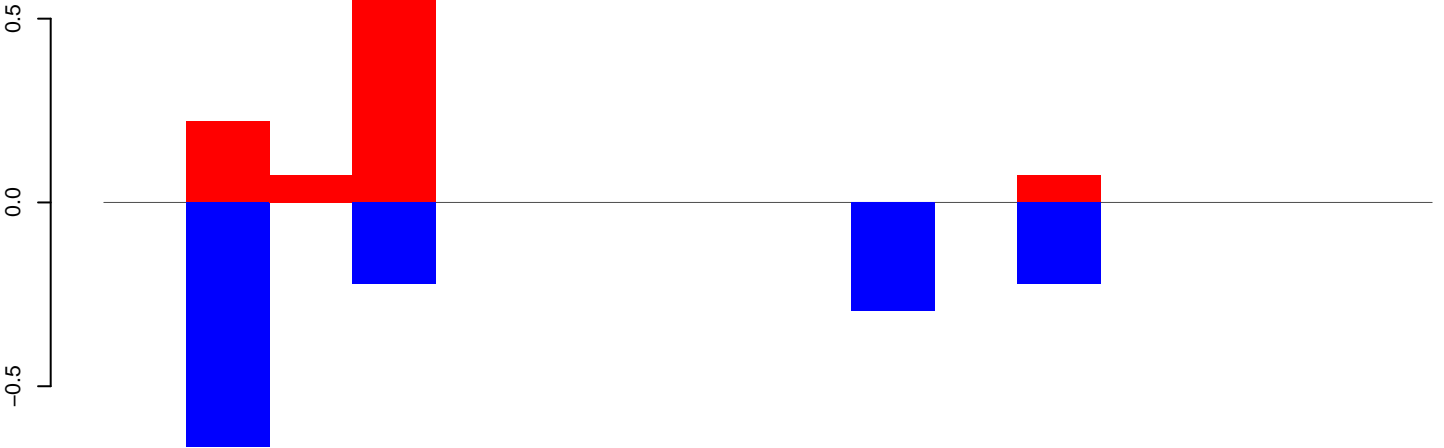
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

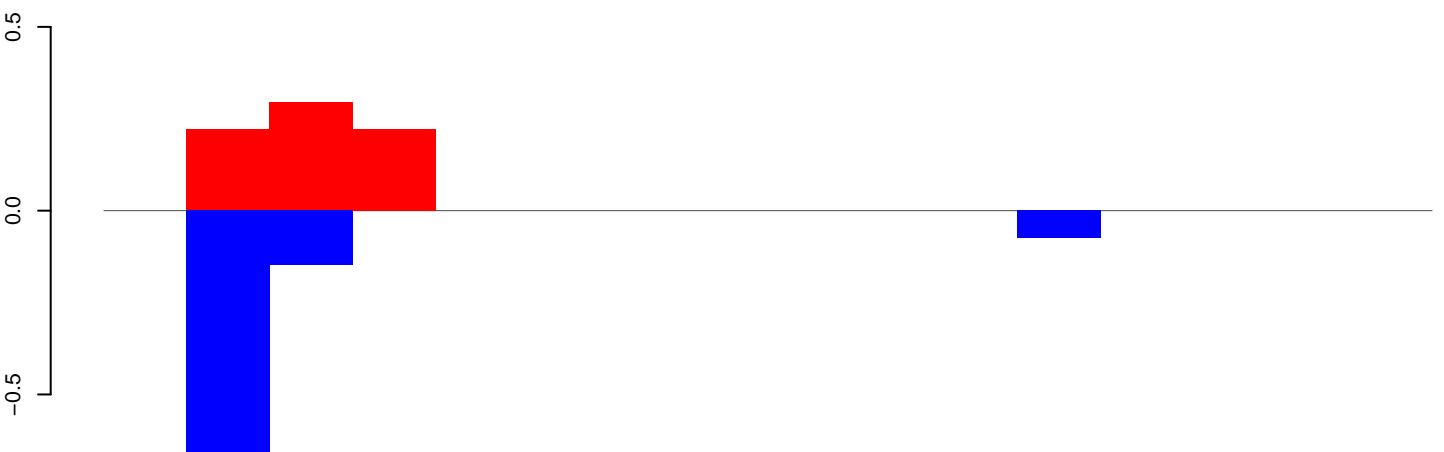


CuQuin_Testes_NL.18_23.rep



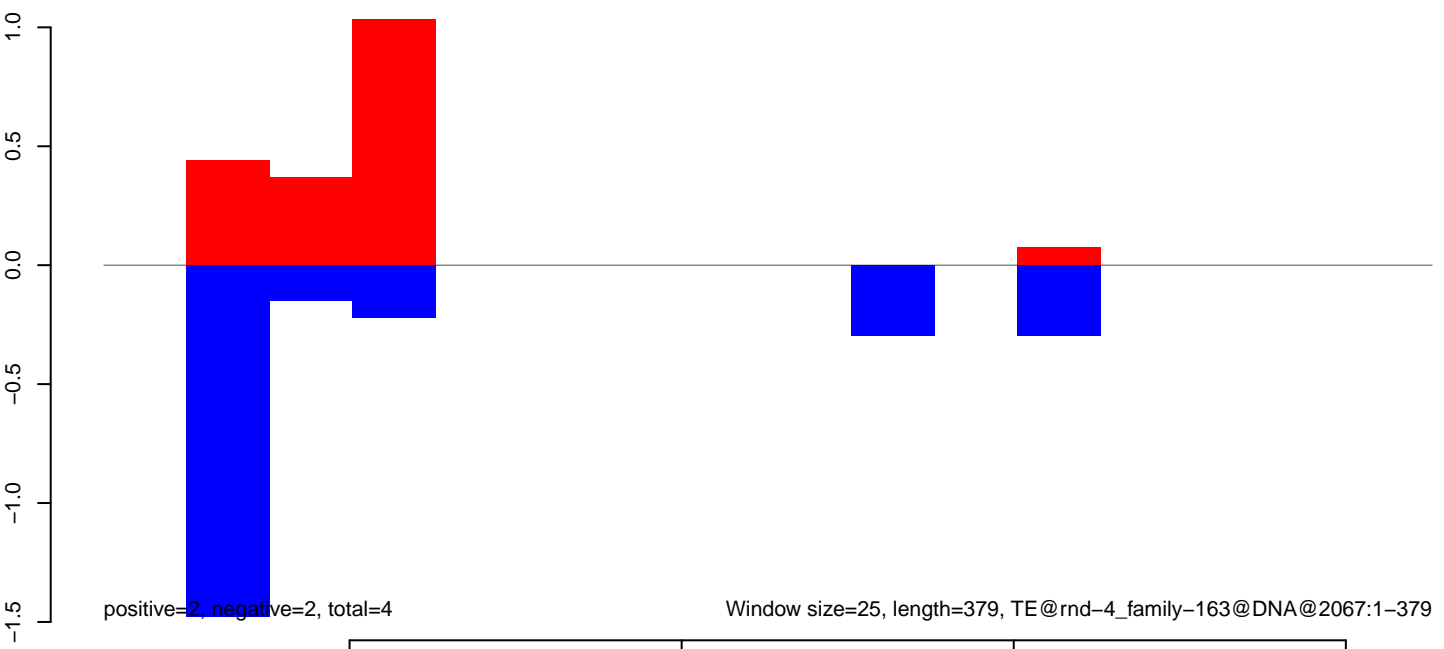
positive=1, negative=1, total=3

CuQuin_Testes_NL.24_35.rep



positive=1, negative=1, total=2

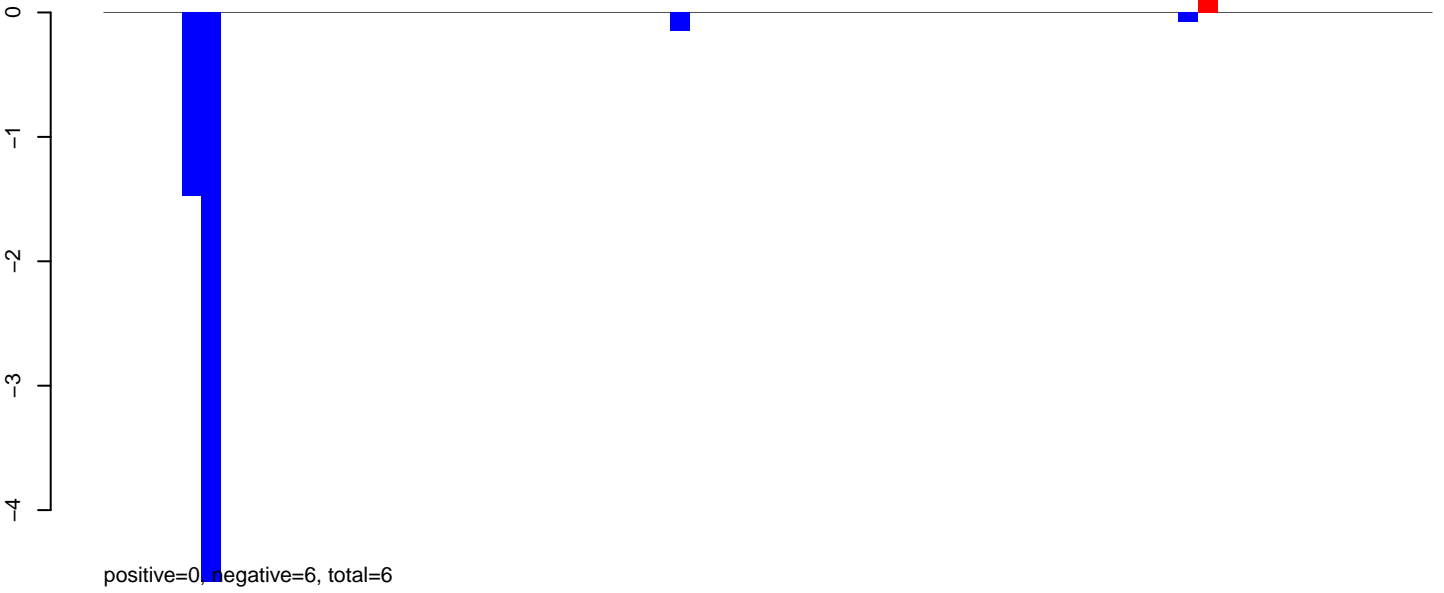
CuQuin_Testes_NL.rep



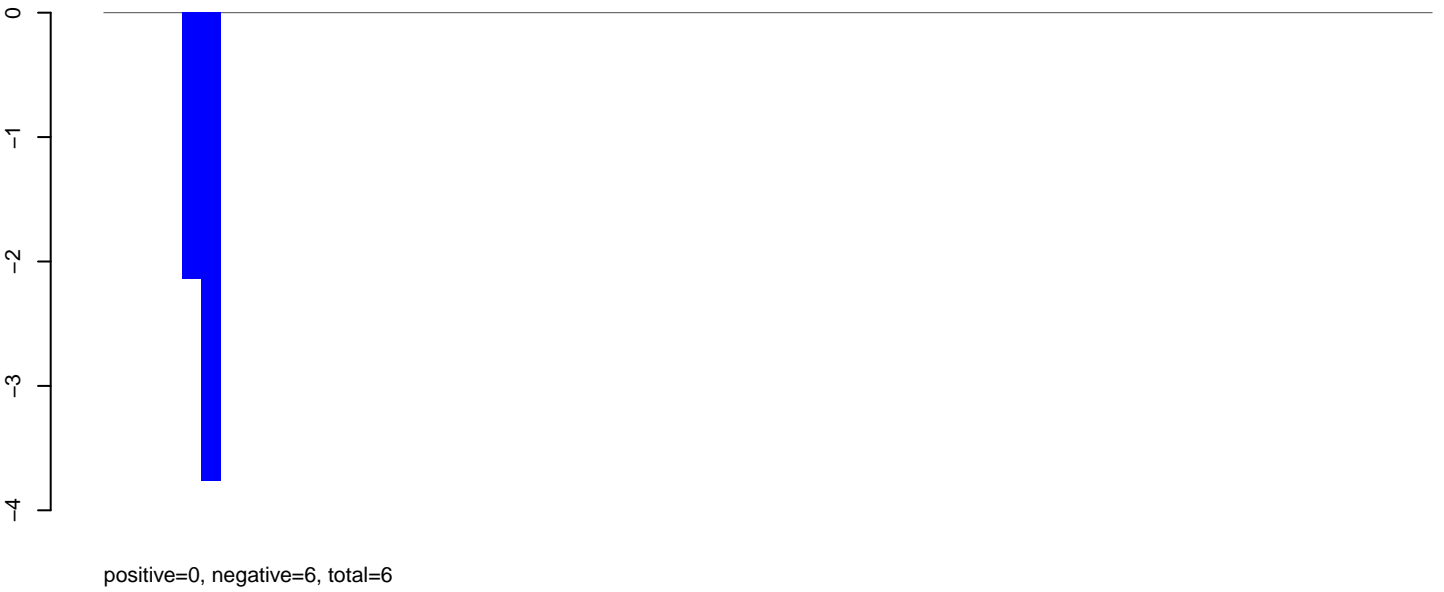
positive=2, negative=2, total=4

Window size=25, length=379, TE@rnd-4_family-163@DNA@2067:1-379

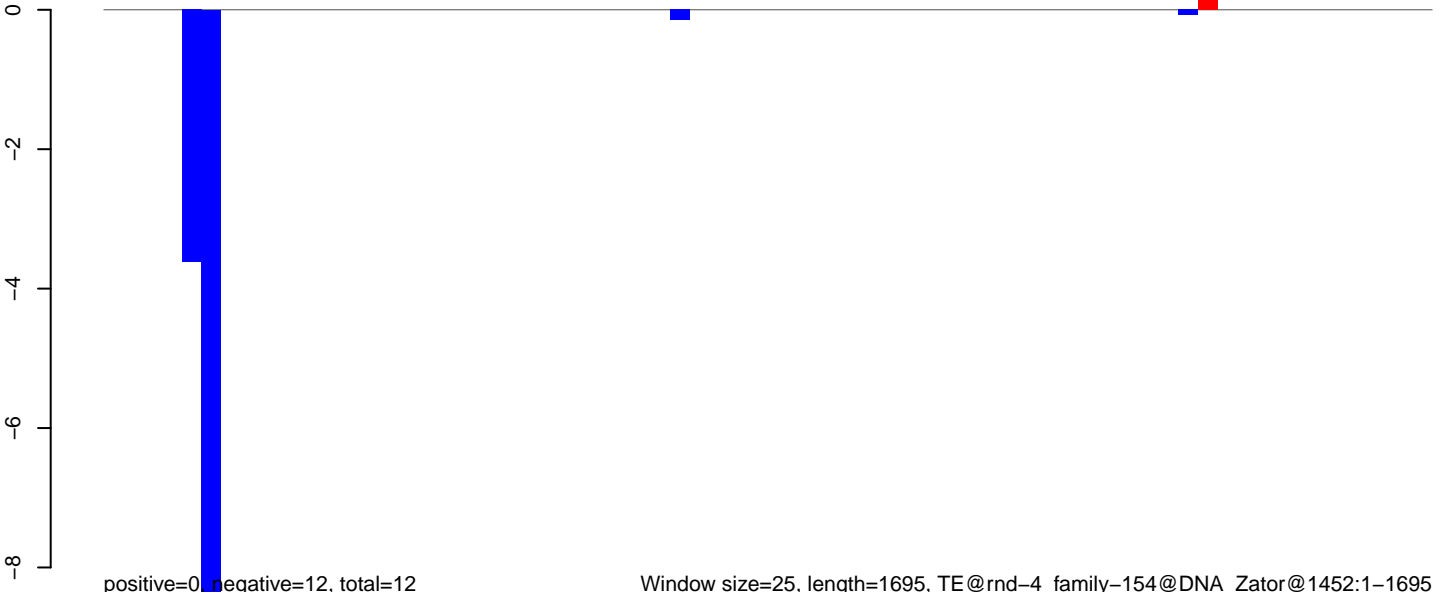
CuQuin_Testes_NL.18_23.rep



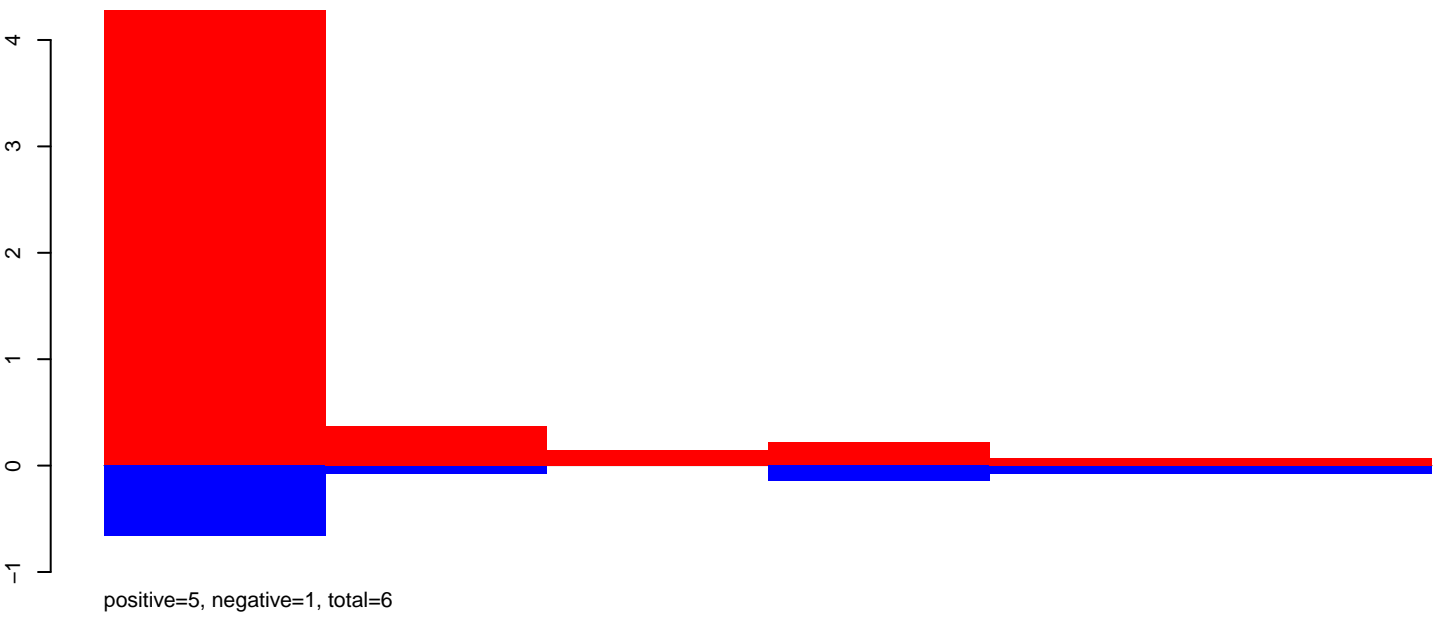
CuQuin_Testes_NL.24_35.rep



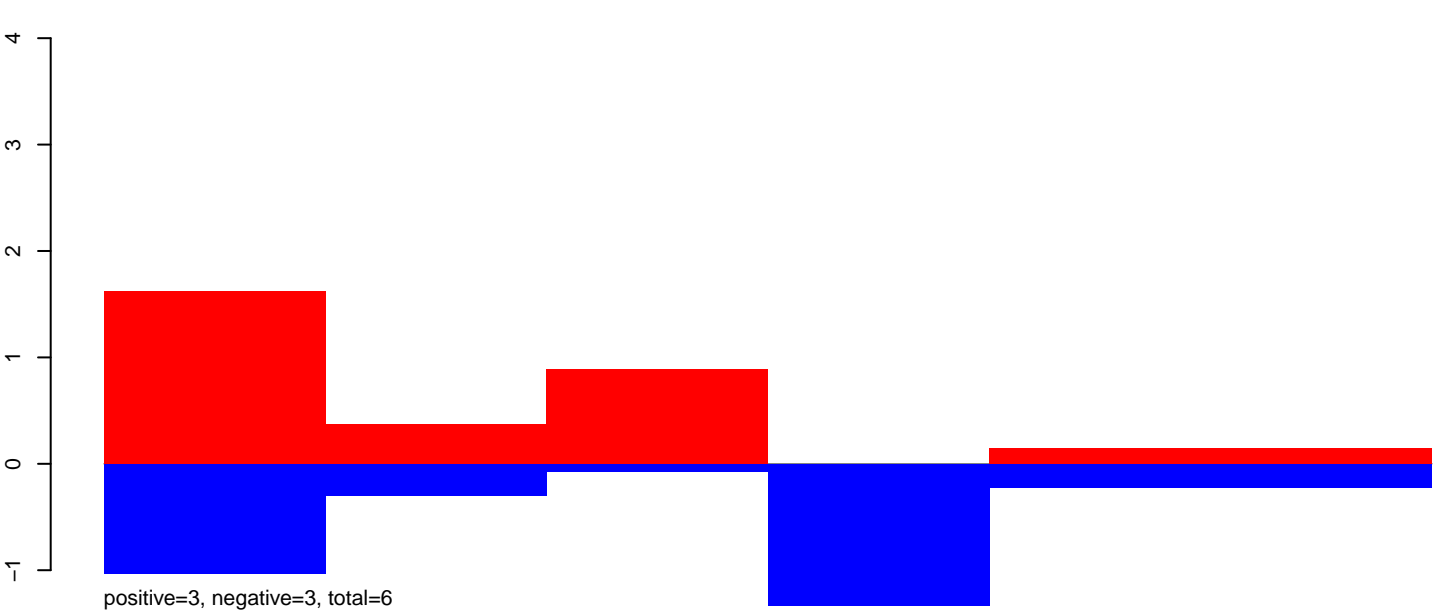
CuQuin_Testes_NL.rep



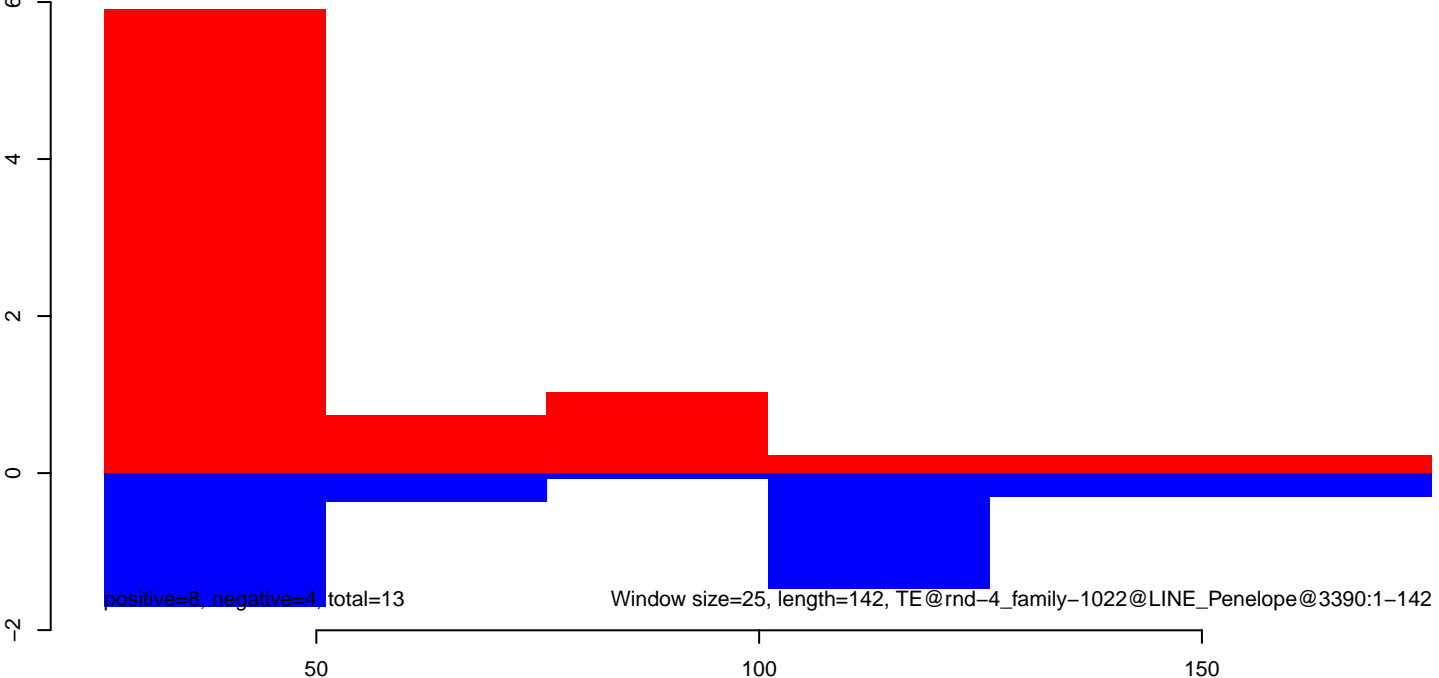
CuQuin_Testes_NL.18_23.rep



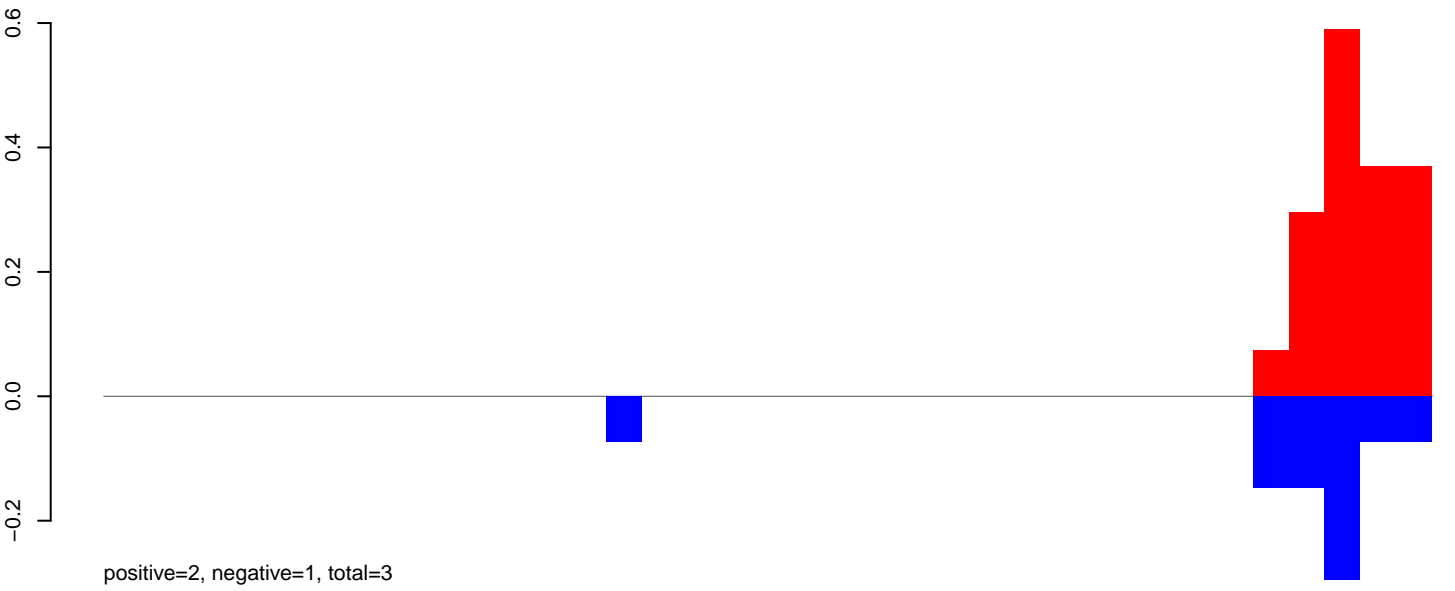
CuQuin_Testes_NL.24_35.rep



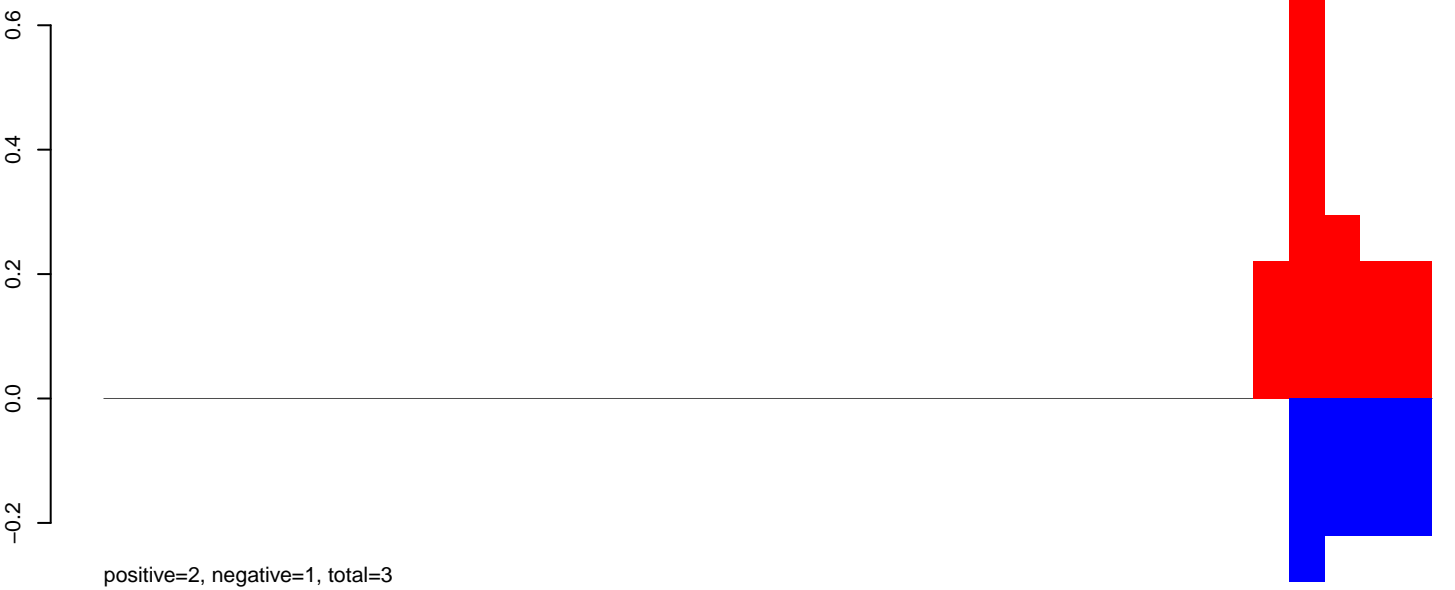
CuQuin_Testes_NL.rep



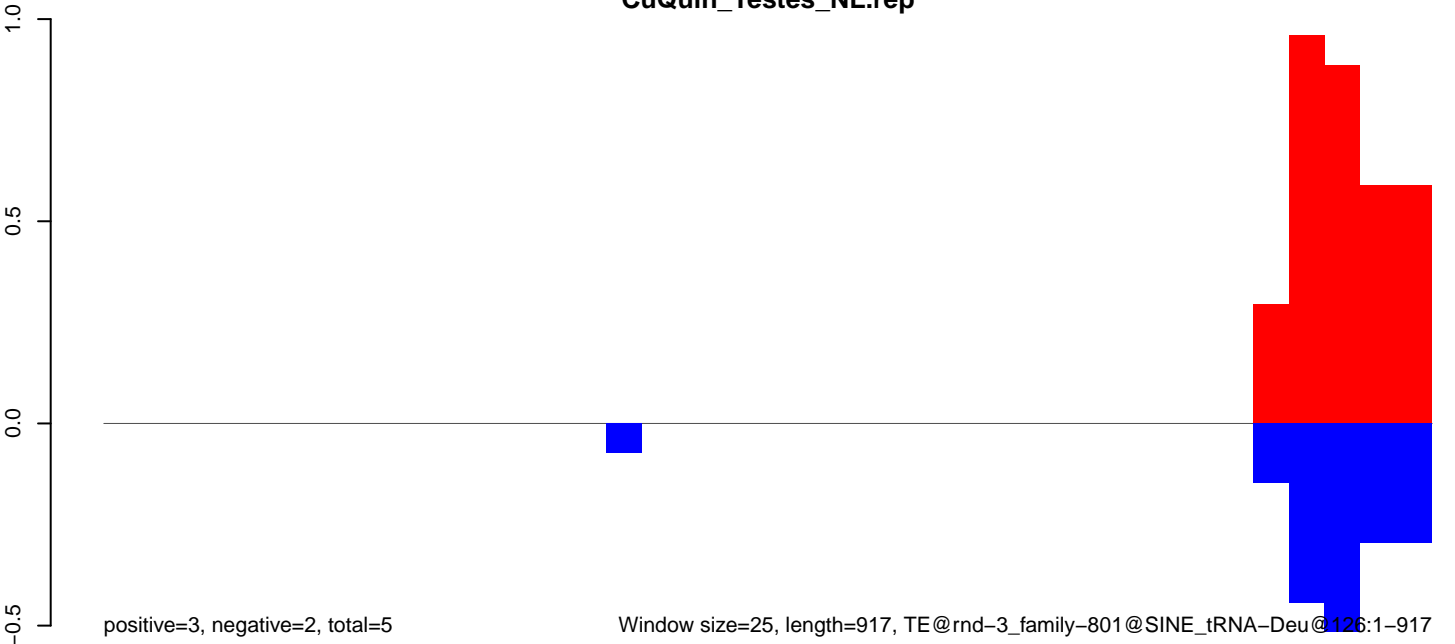
CuQuin_Testes_NL.18_23.rep



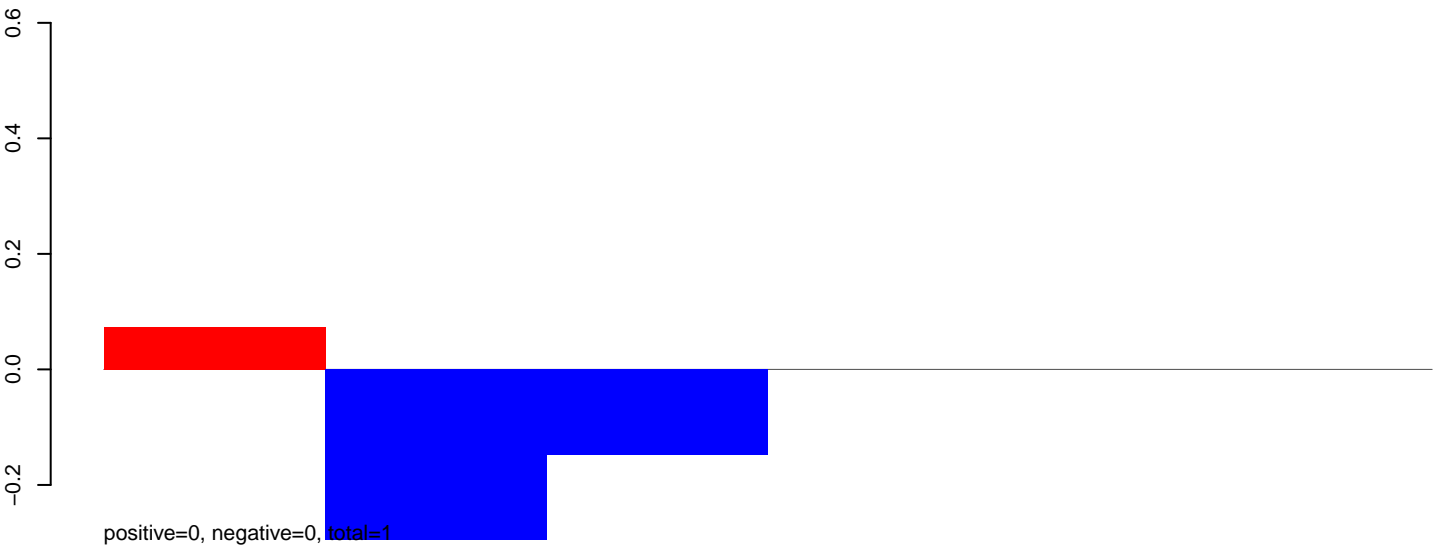
CuQuin_Testes_NL.24_35.rep



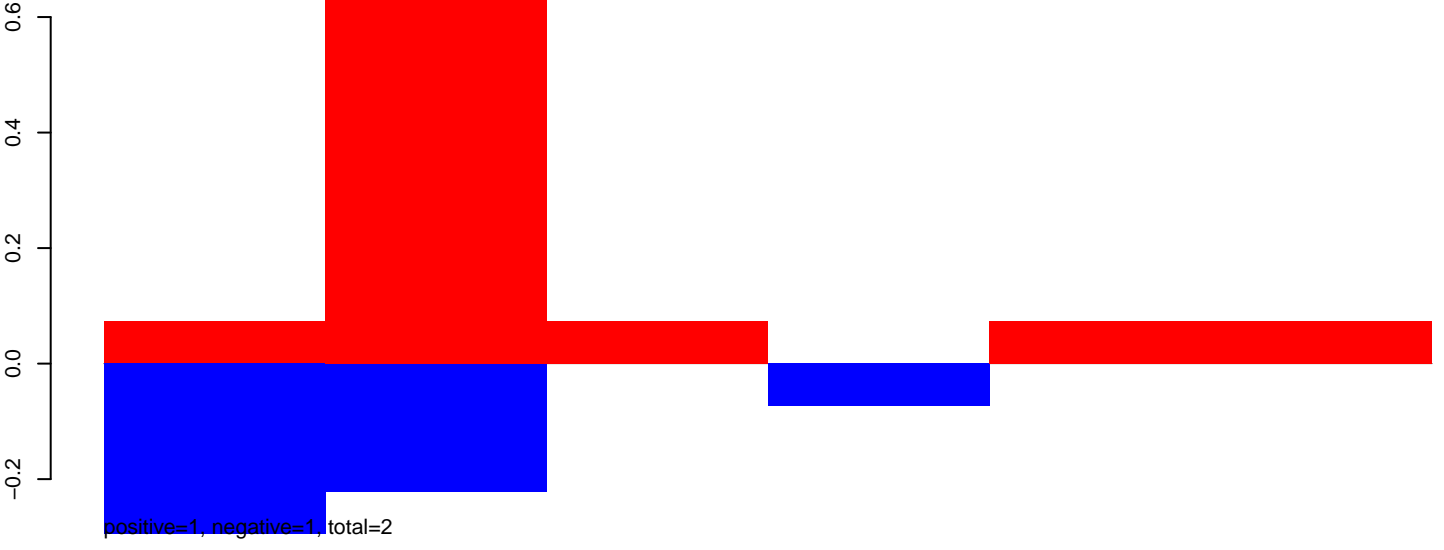
CuQuin_Testes_NL.rep



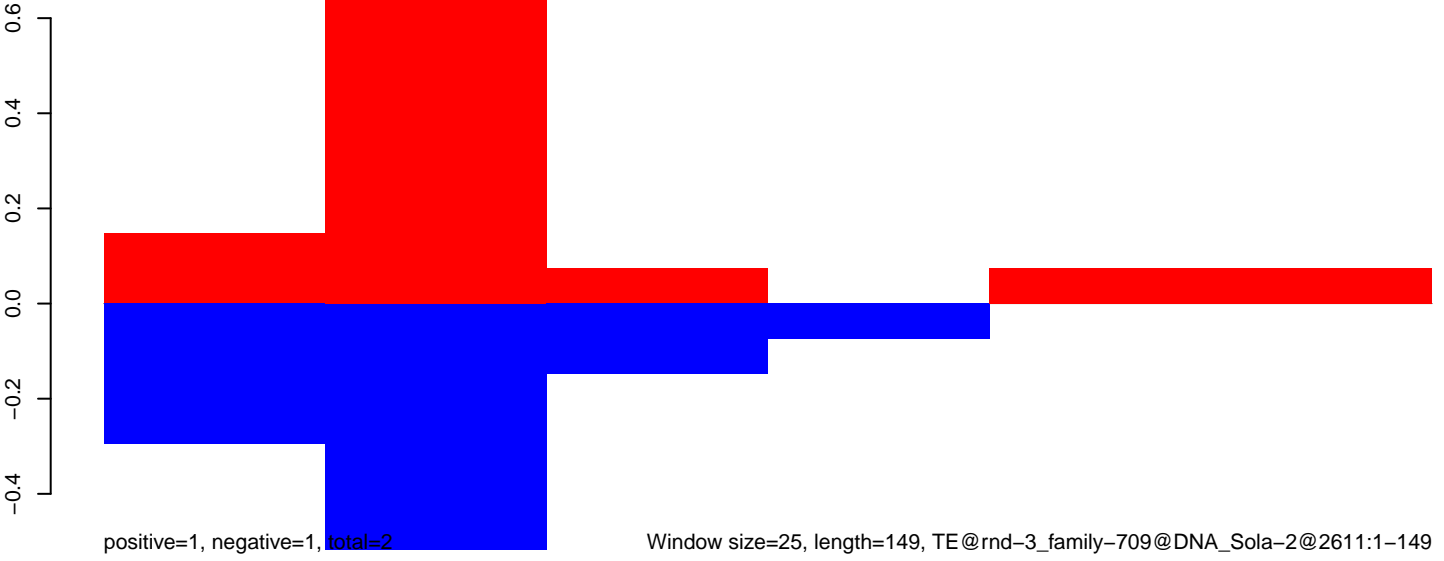
CuQuin_Testes_NL.18_23.rep



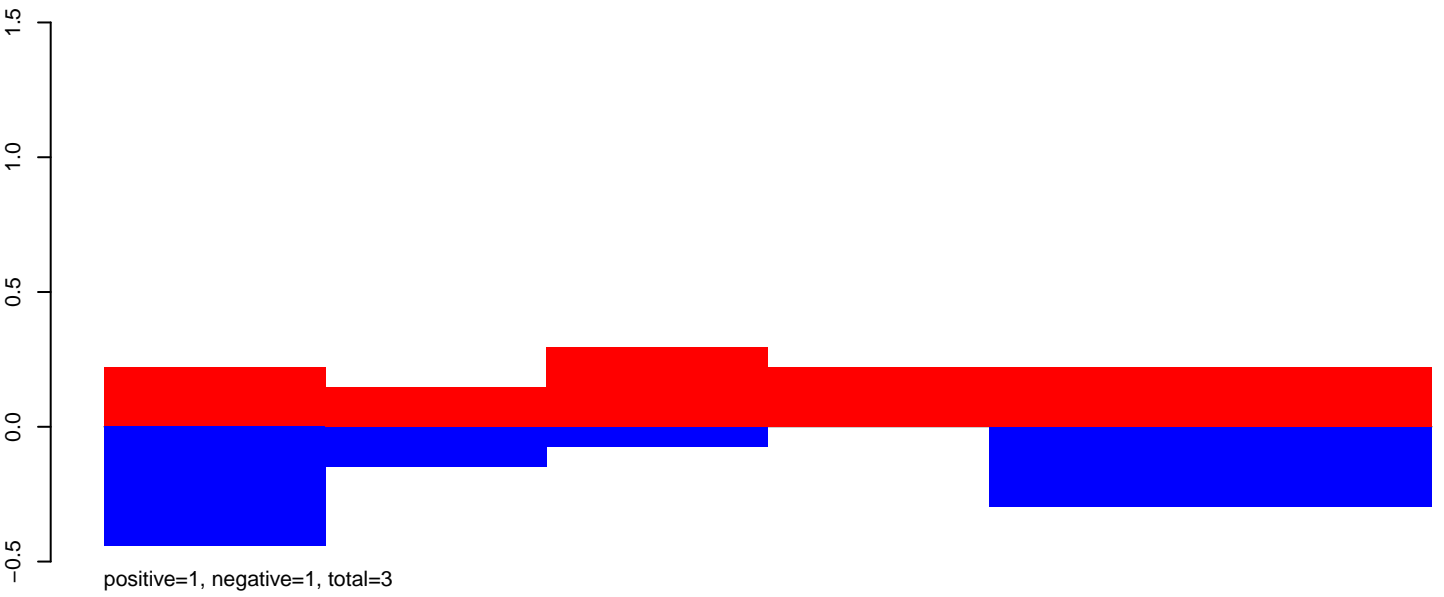
CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



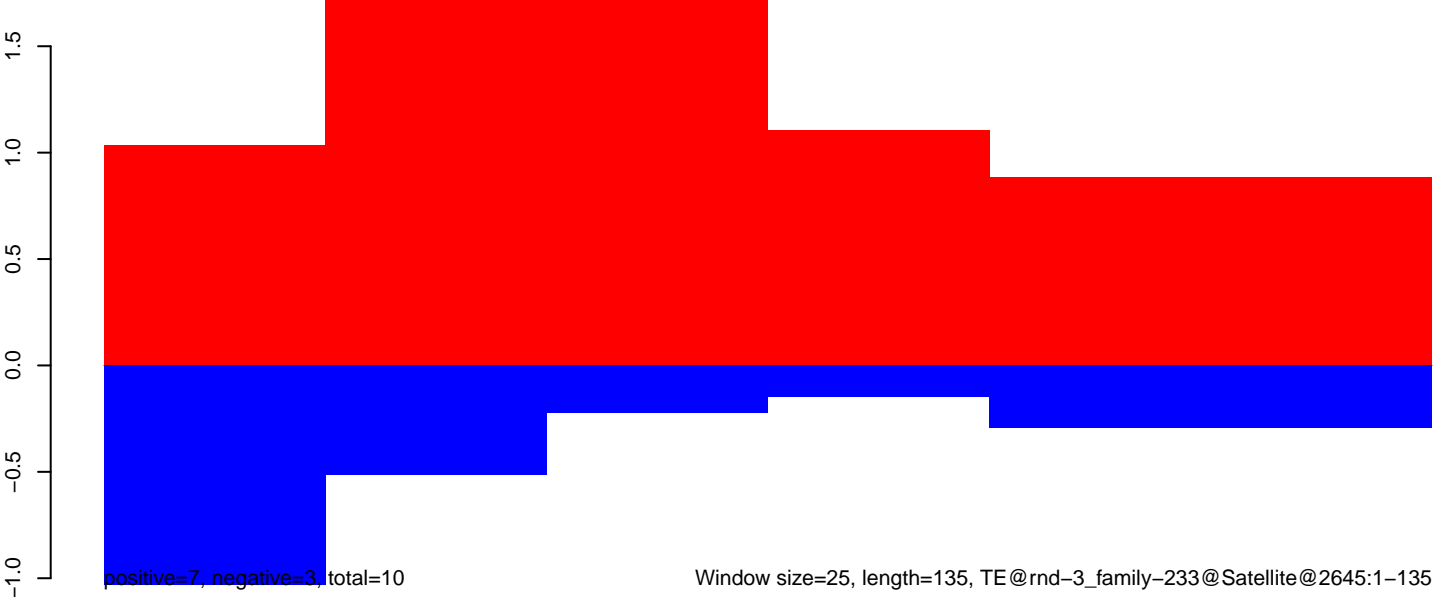
CuQuin_Testes_NL.18_23.rep



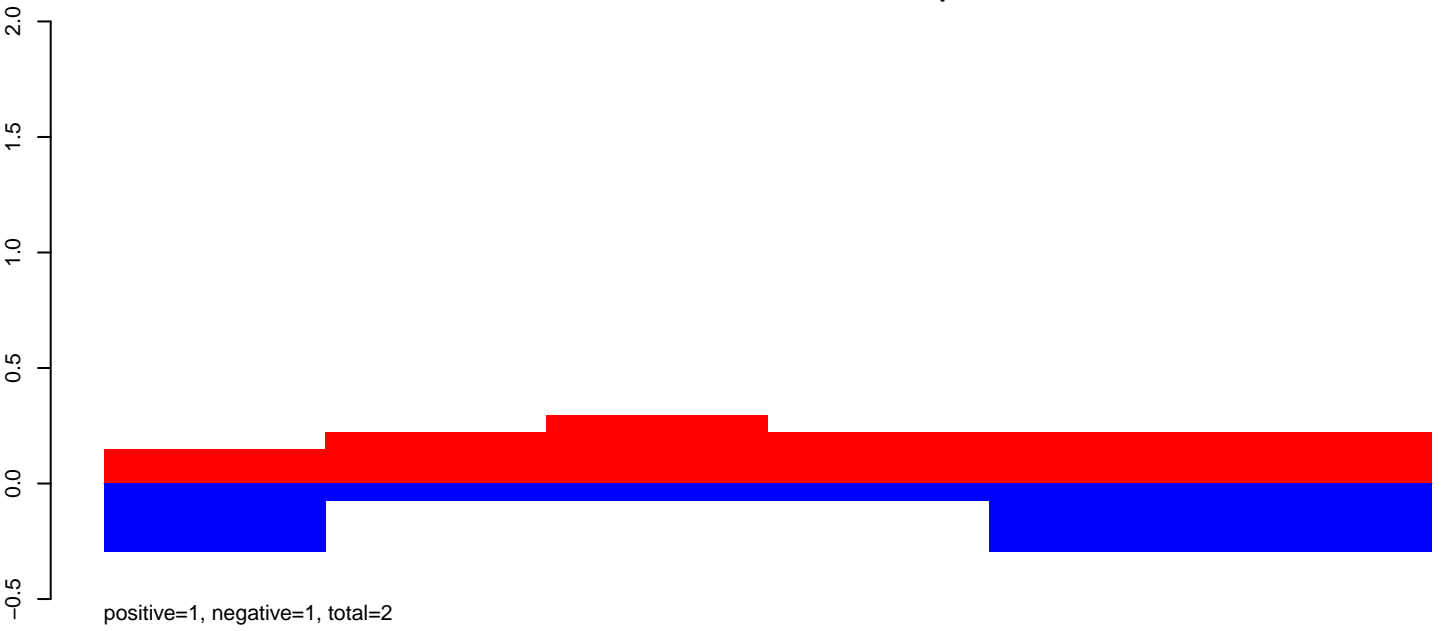
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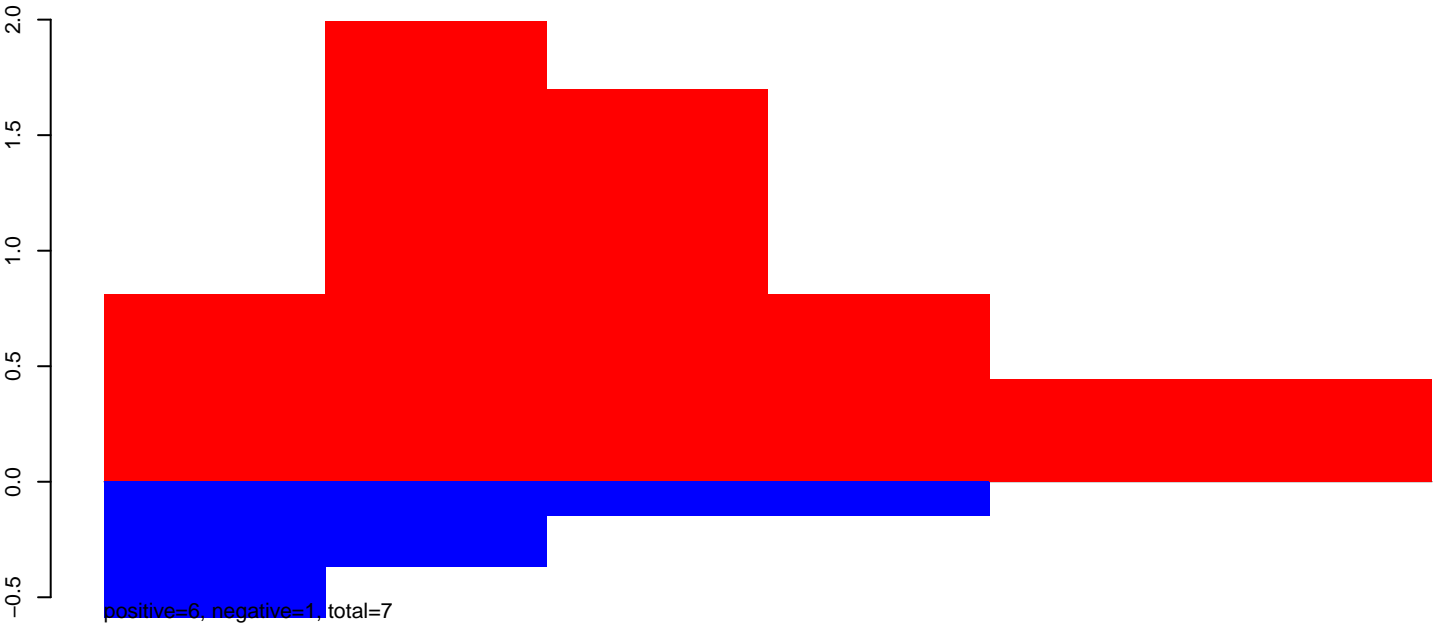
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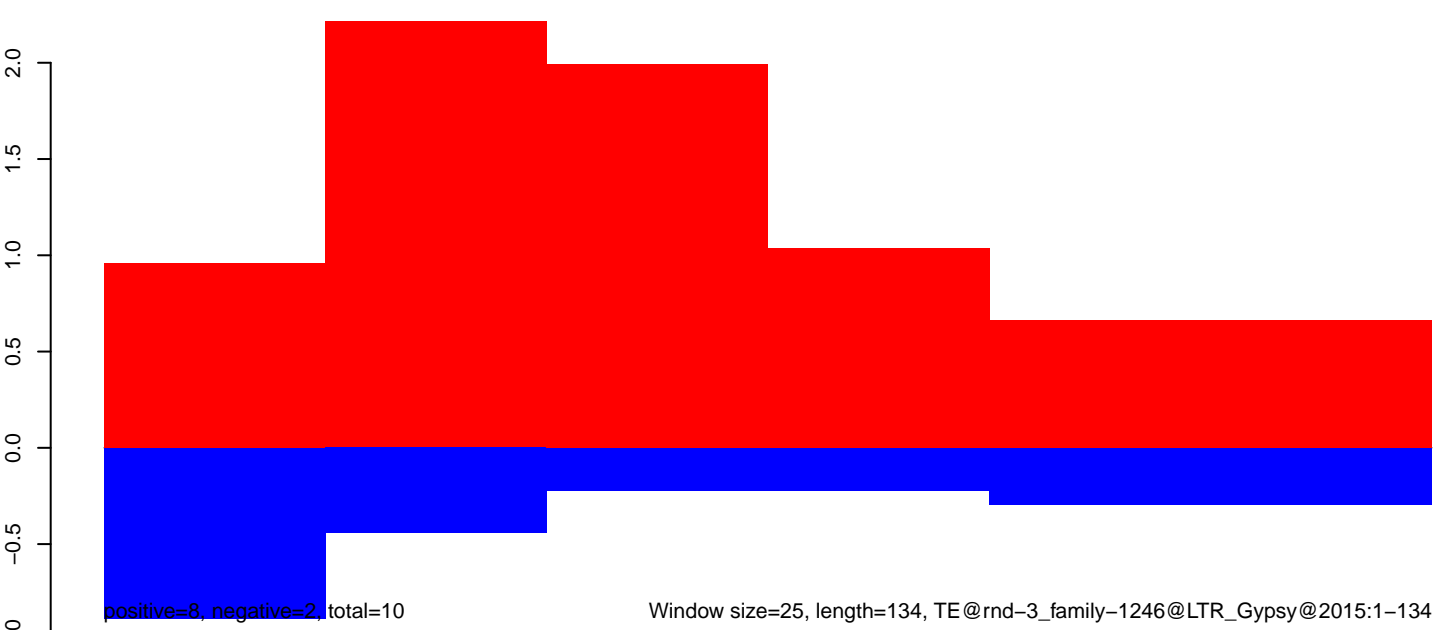
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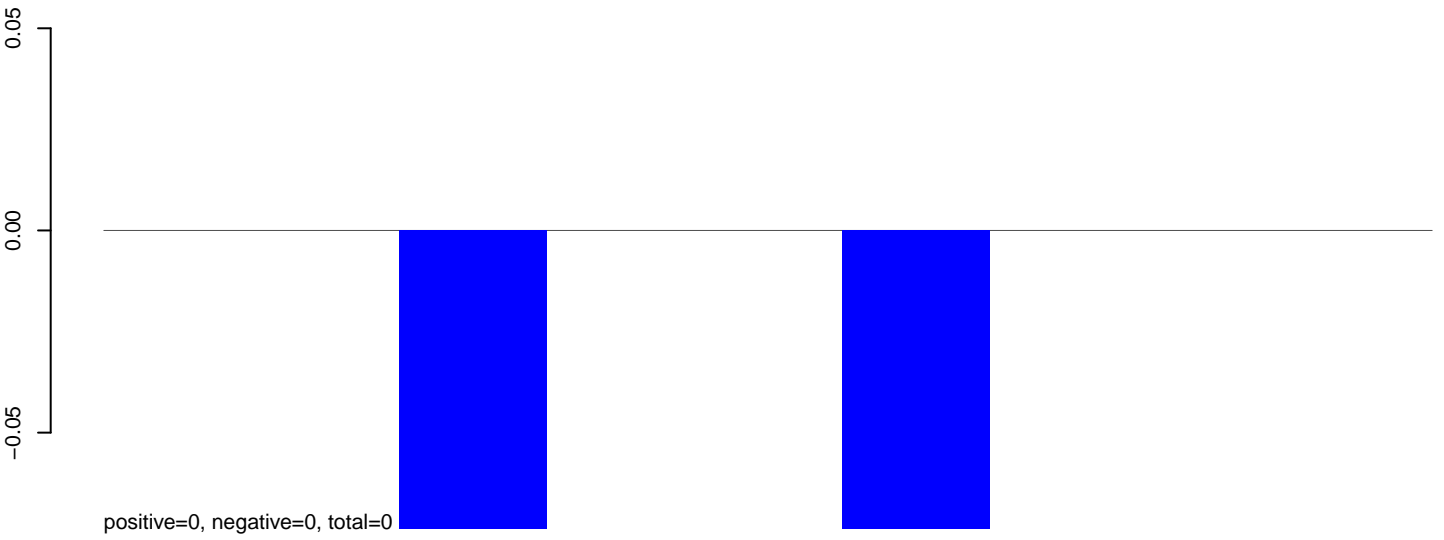
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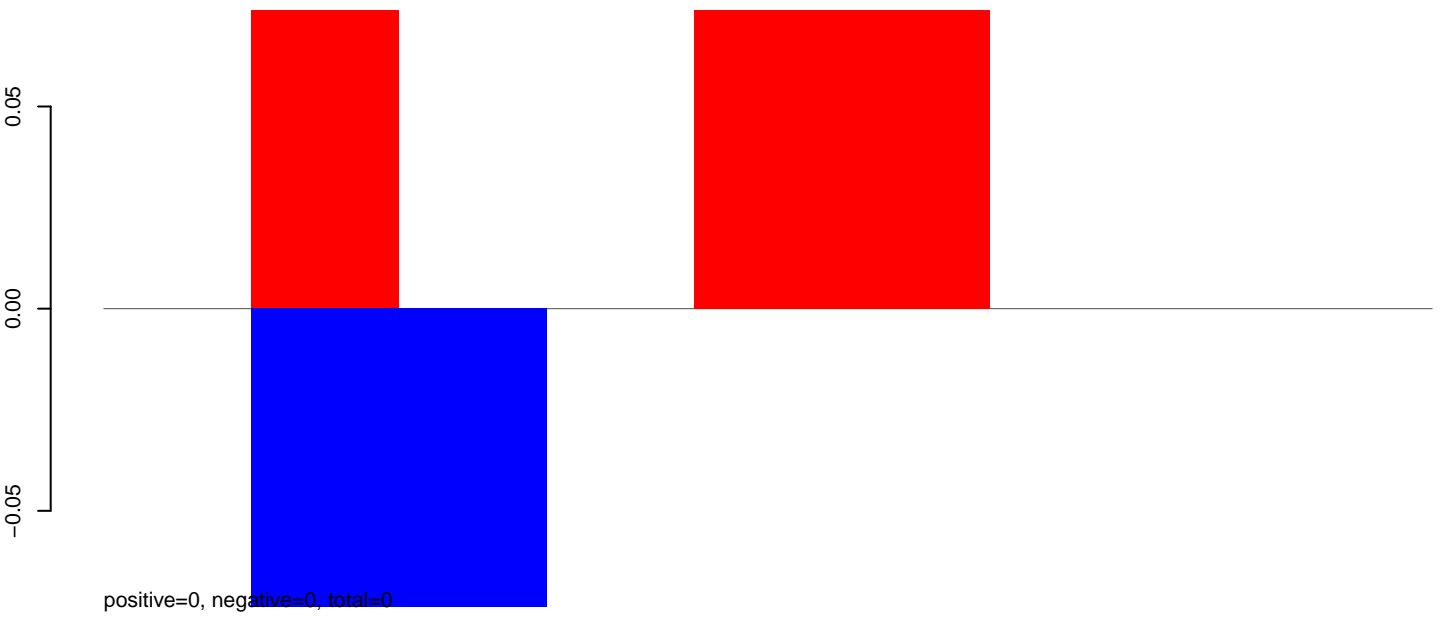
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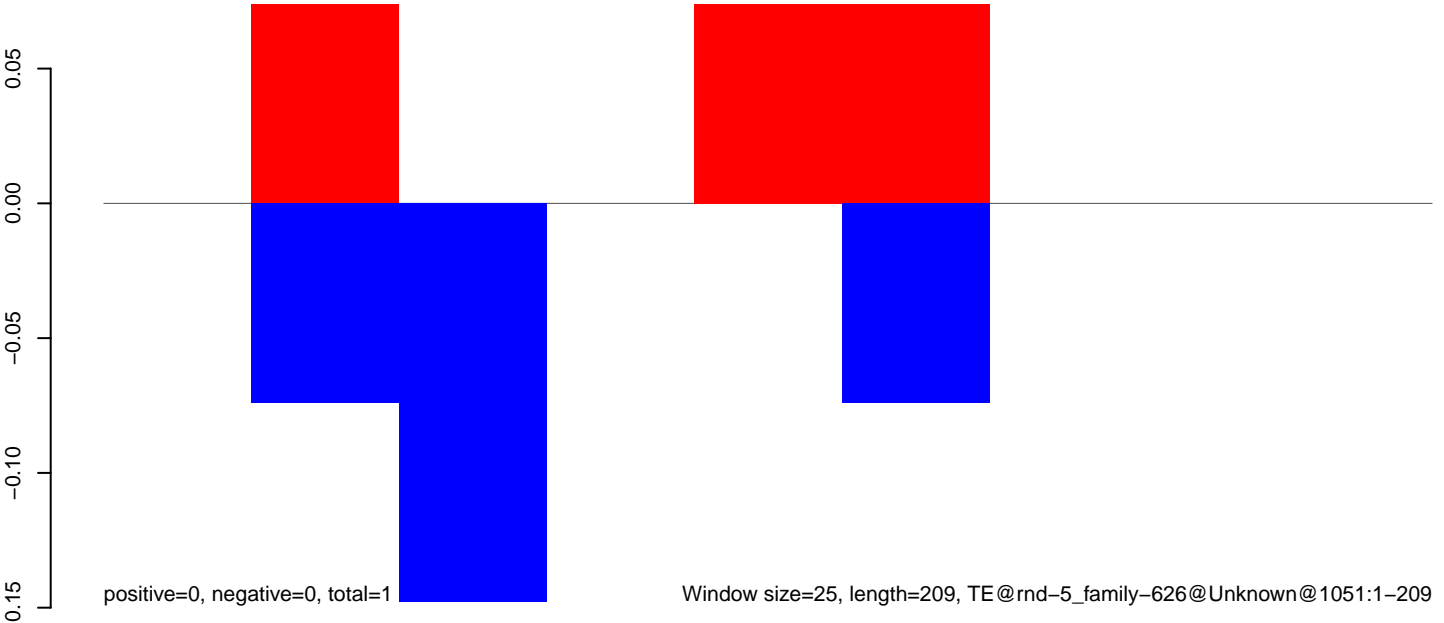
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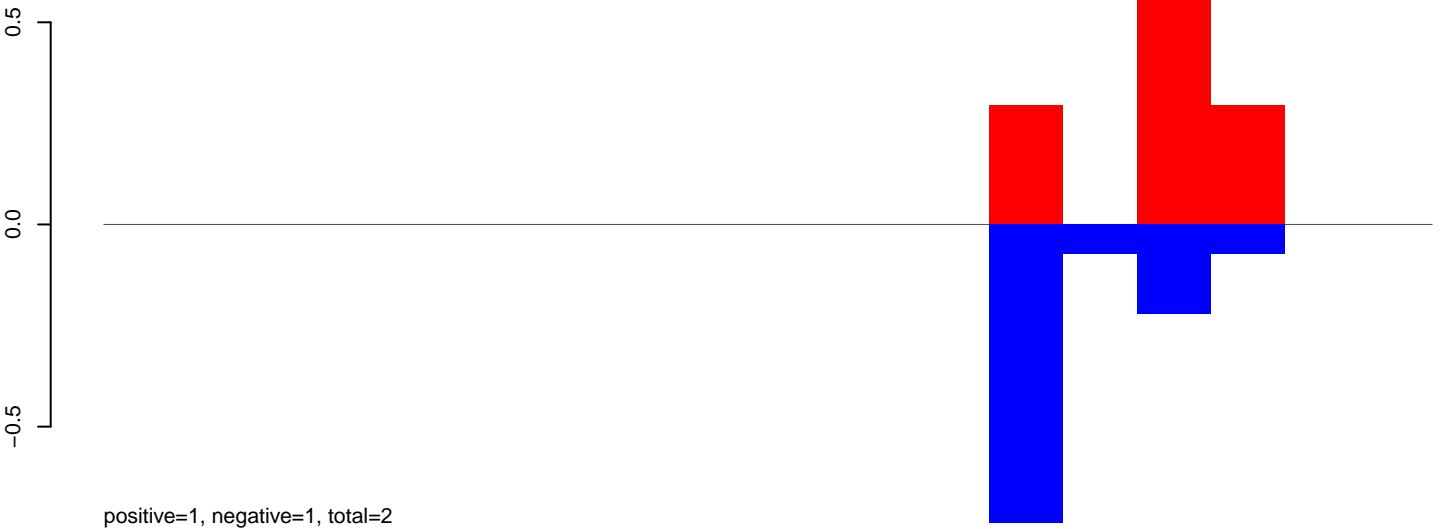
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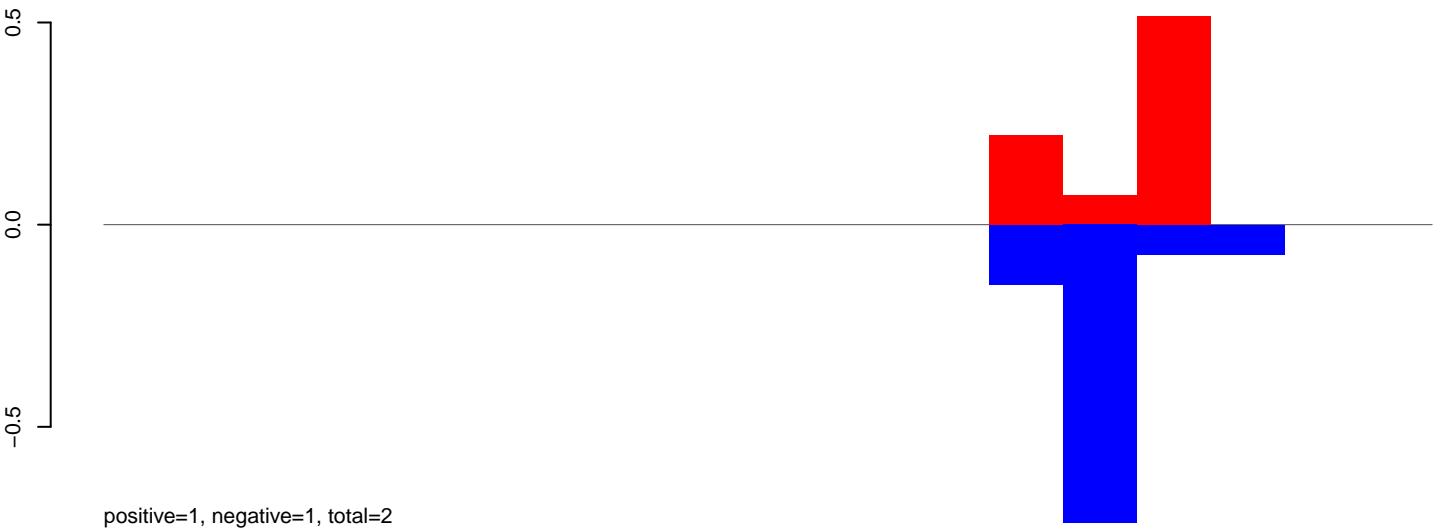
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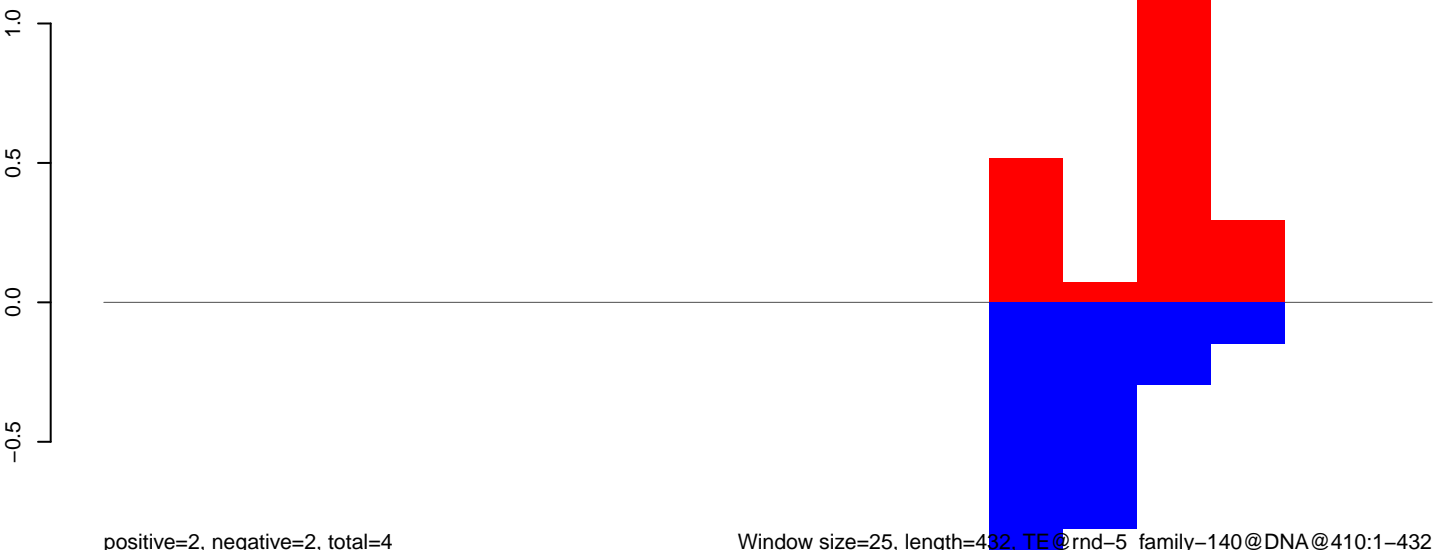
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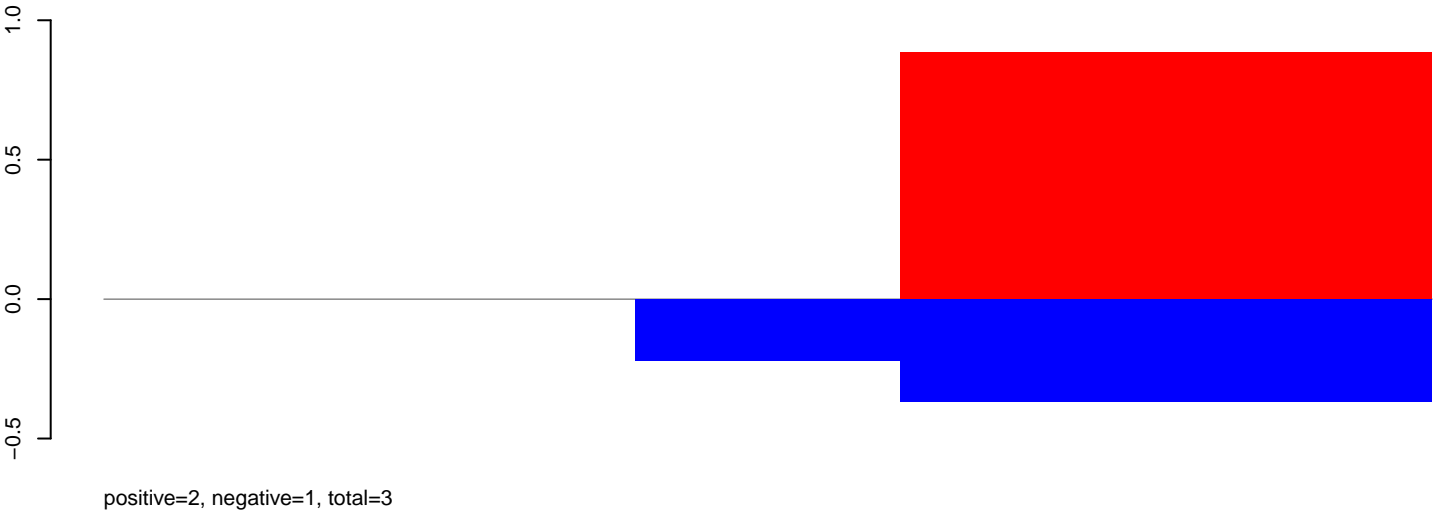
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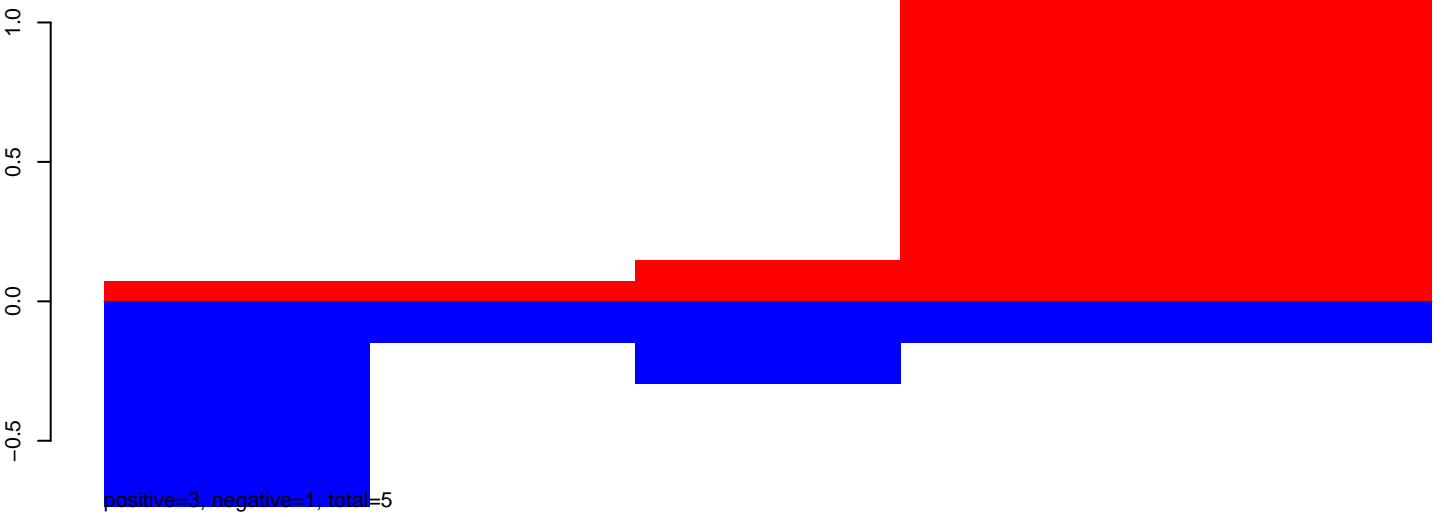
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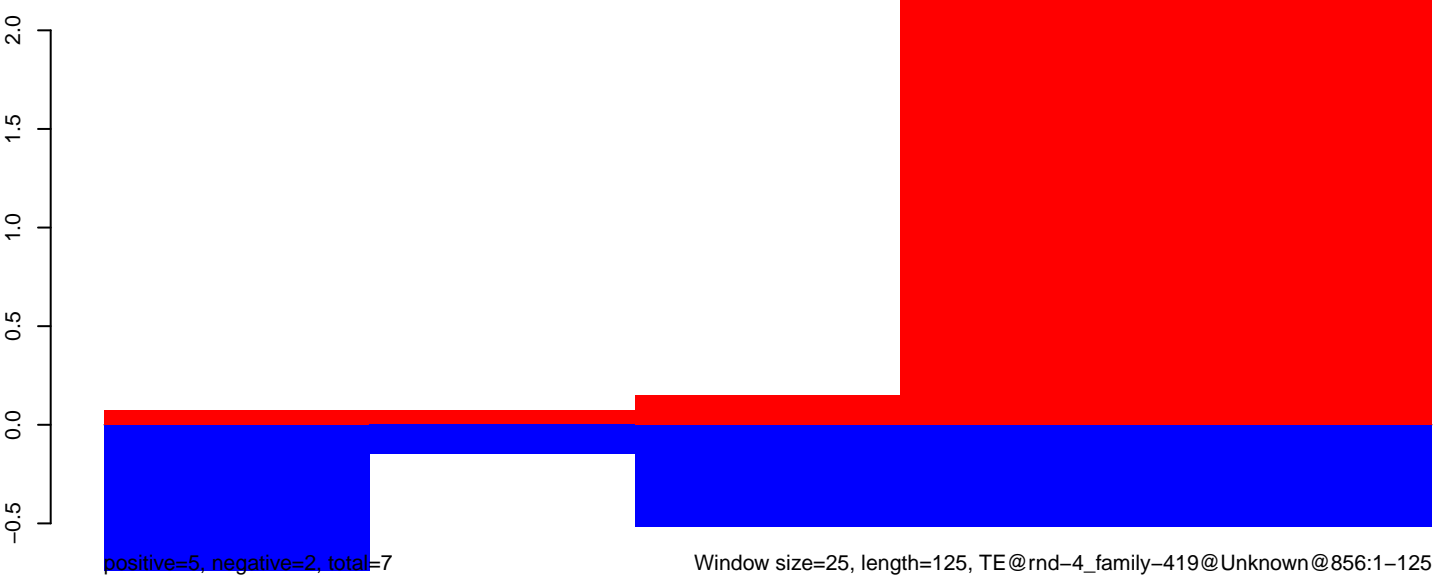
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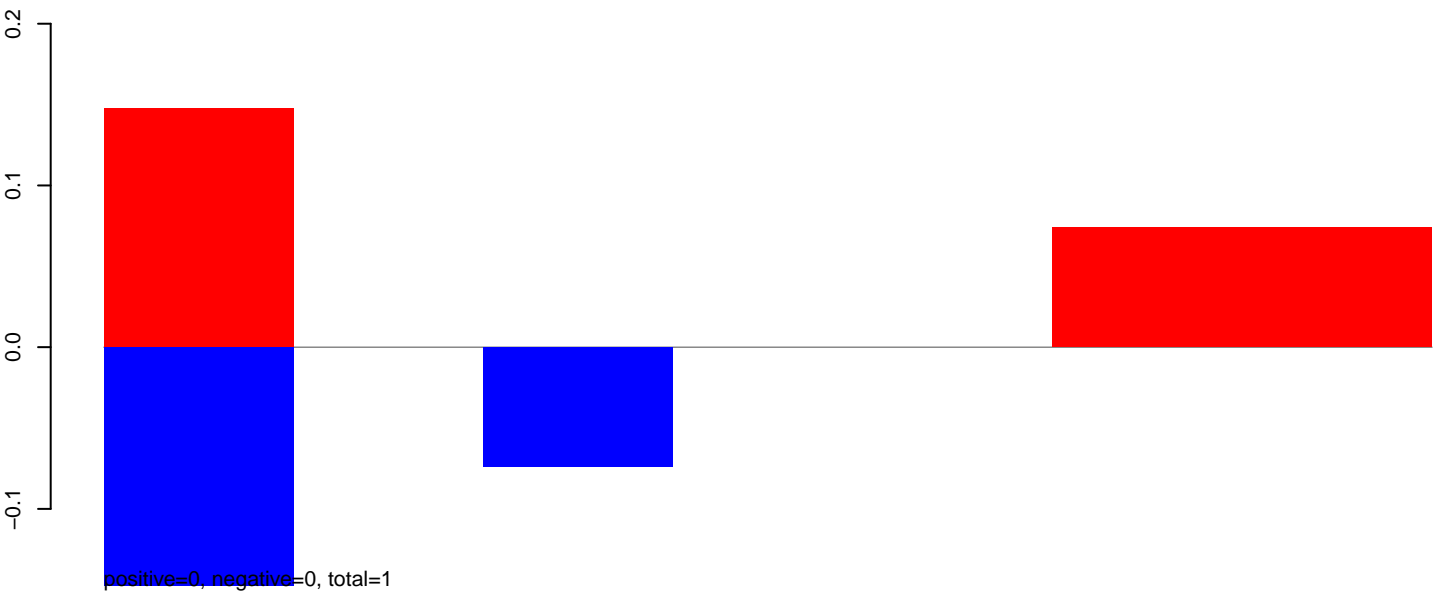
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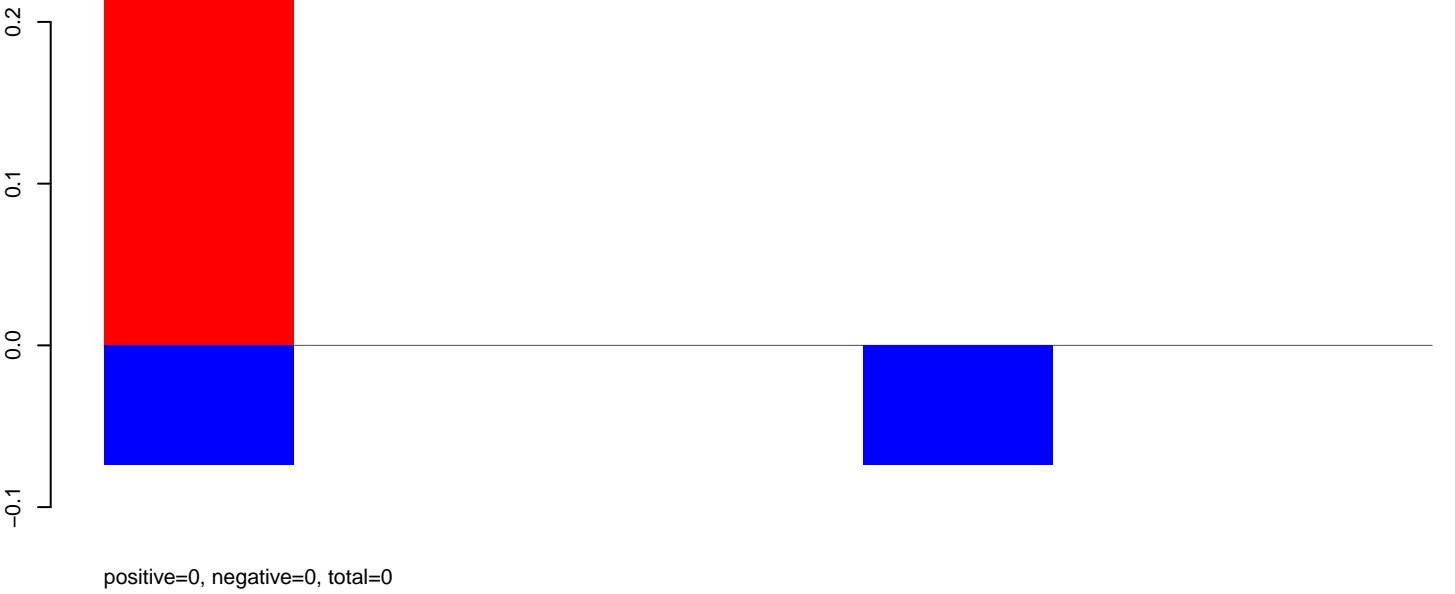
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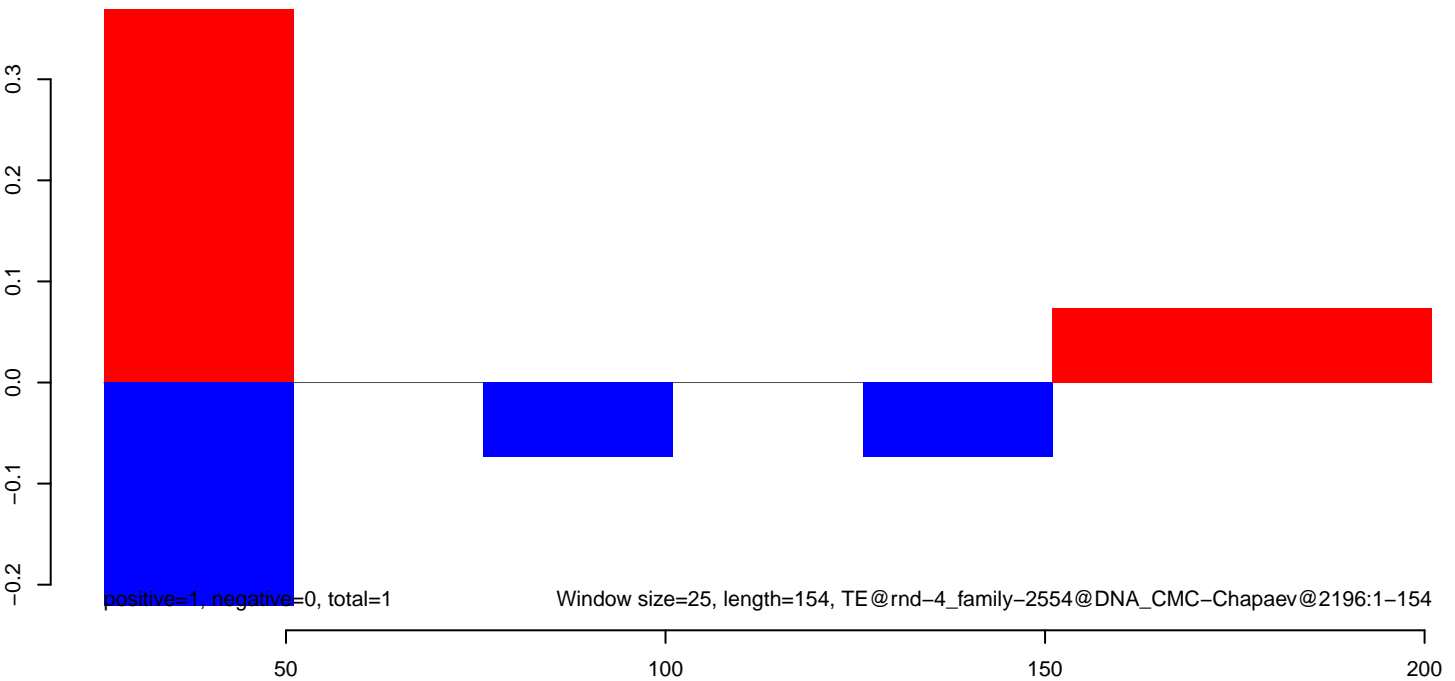
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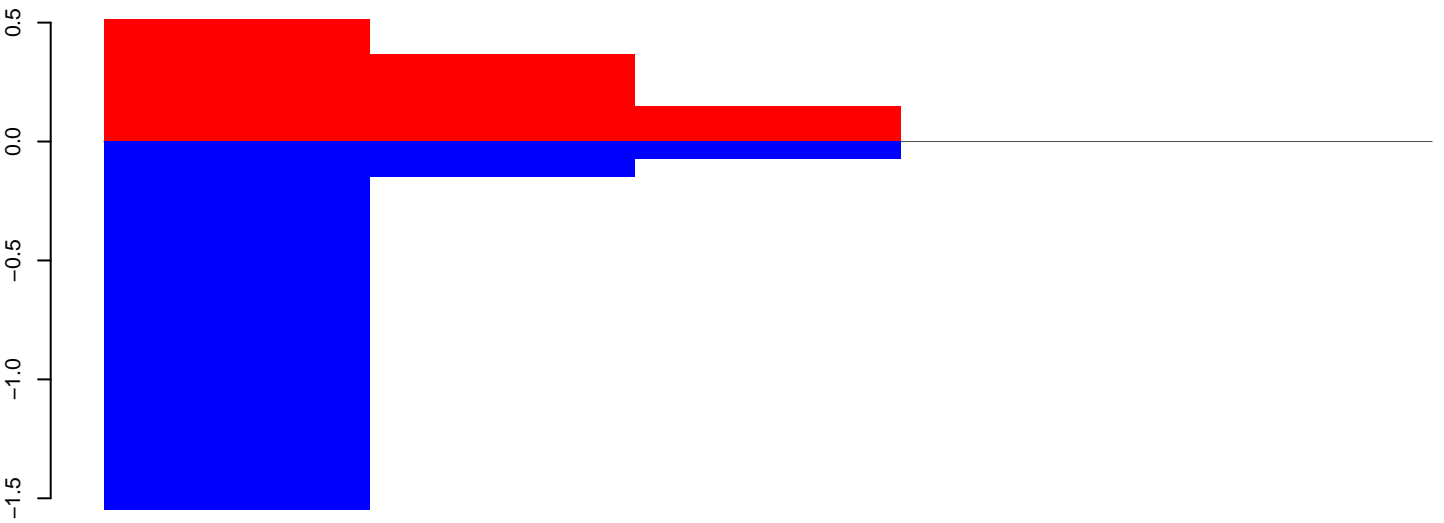
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CuQuin_Testes_NL.rep

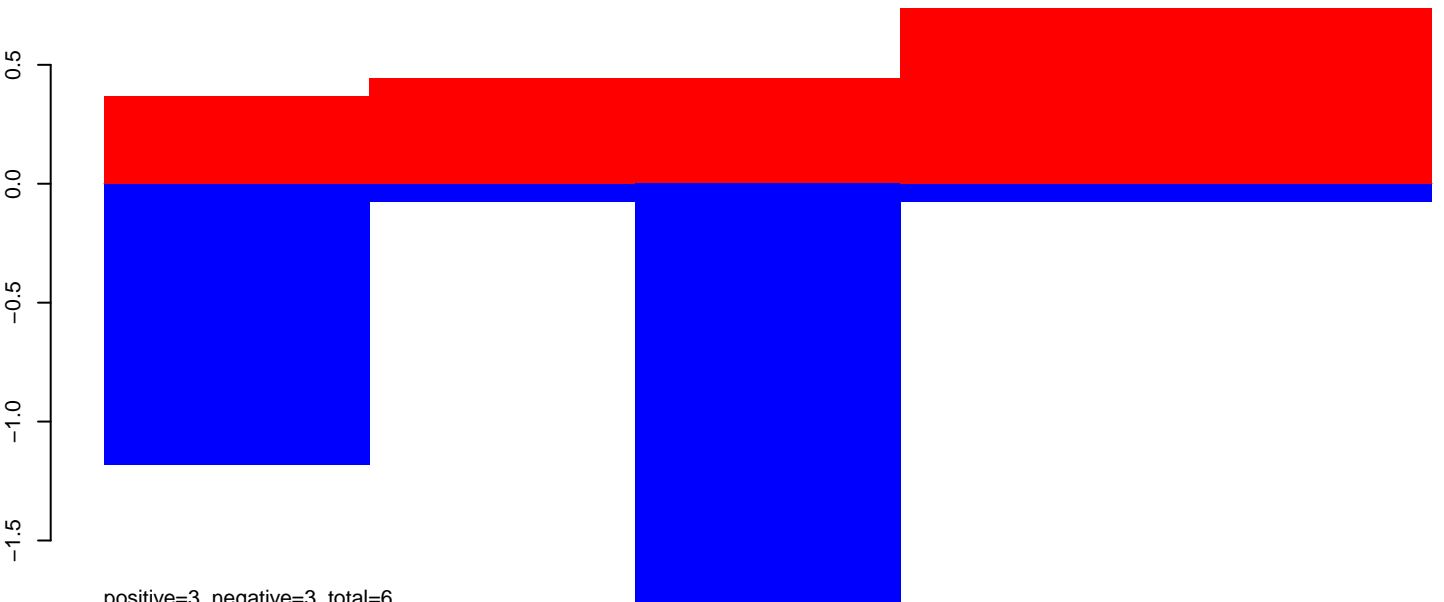


CuQuin_Testes_NL.18_23.rep



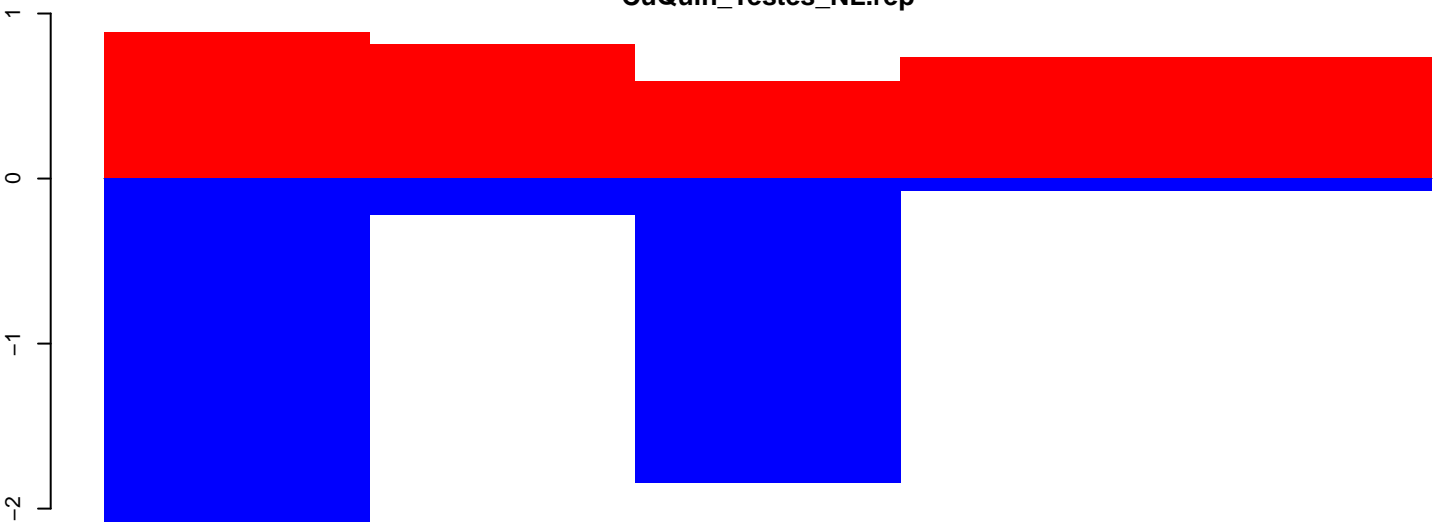
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CuQuin_Testes_NL.24_35.rep



positive=3, negative=3, total=6

CuQuin_Testes_NL.rep

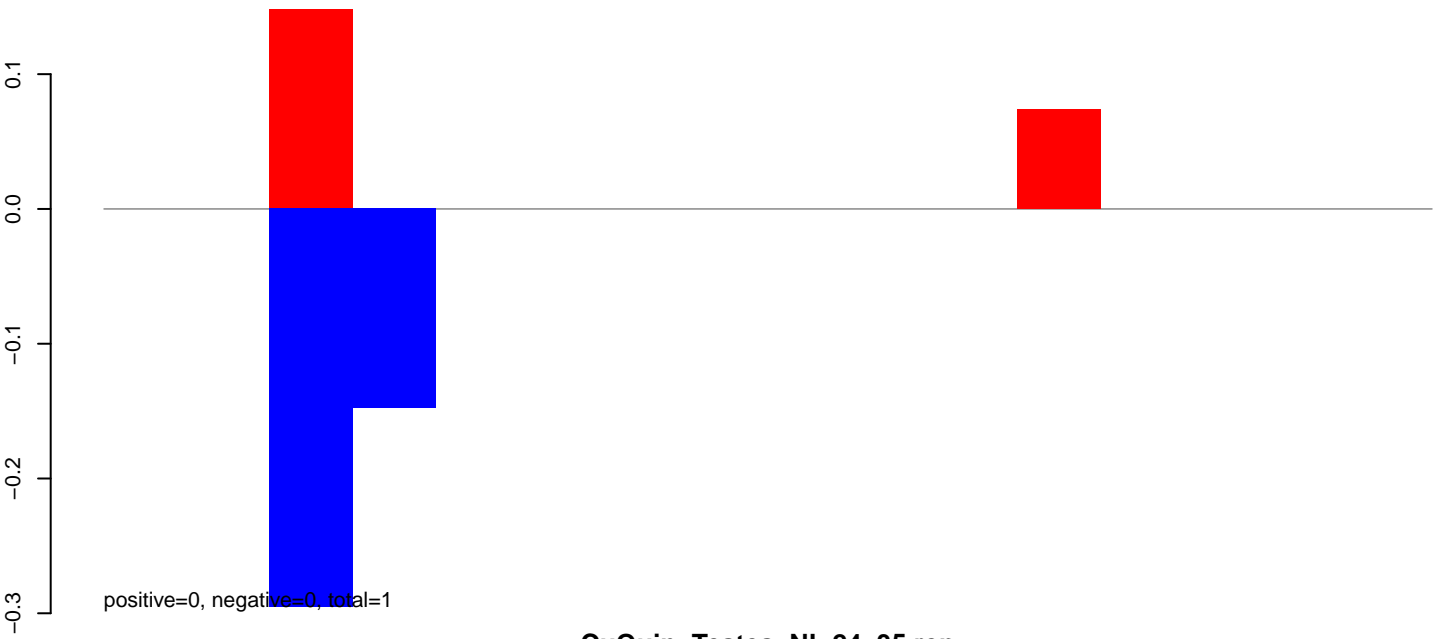


positive=4, negative=5, total=9

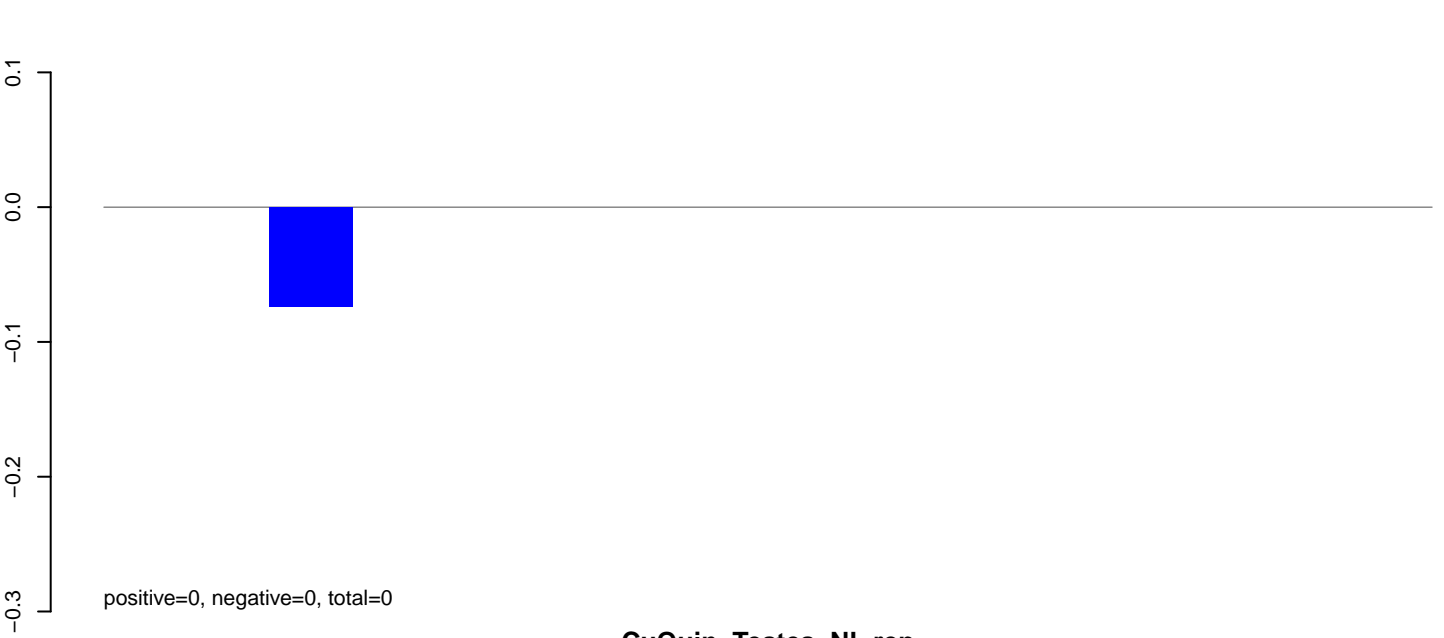
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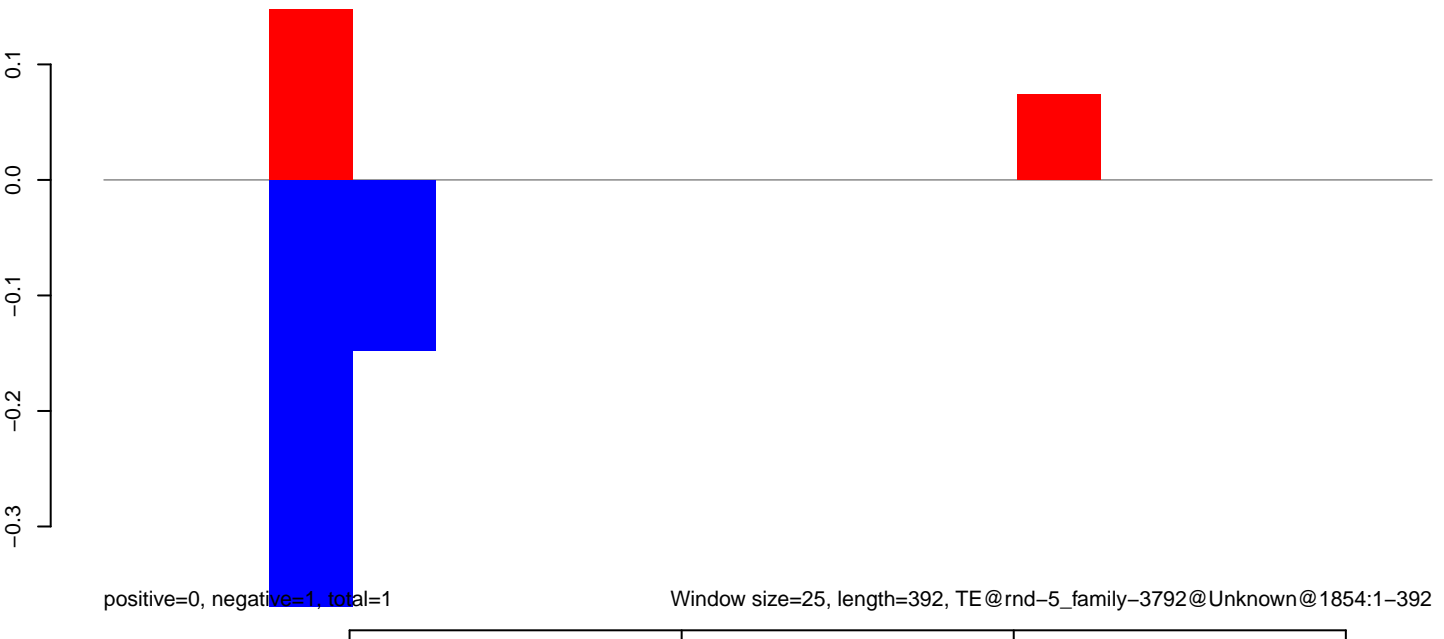
CuQuin_Testes_NL.18_23.rep



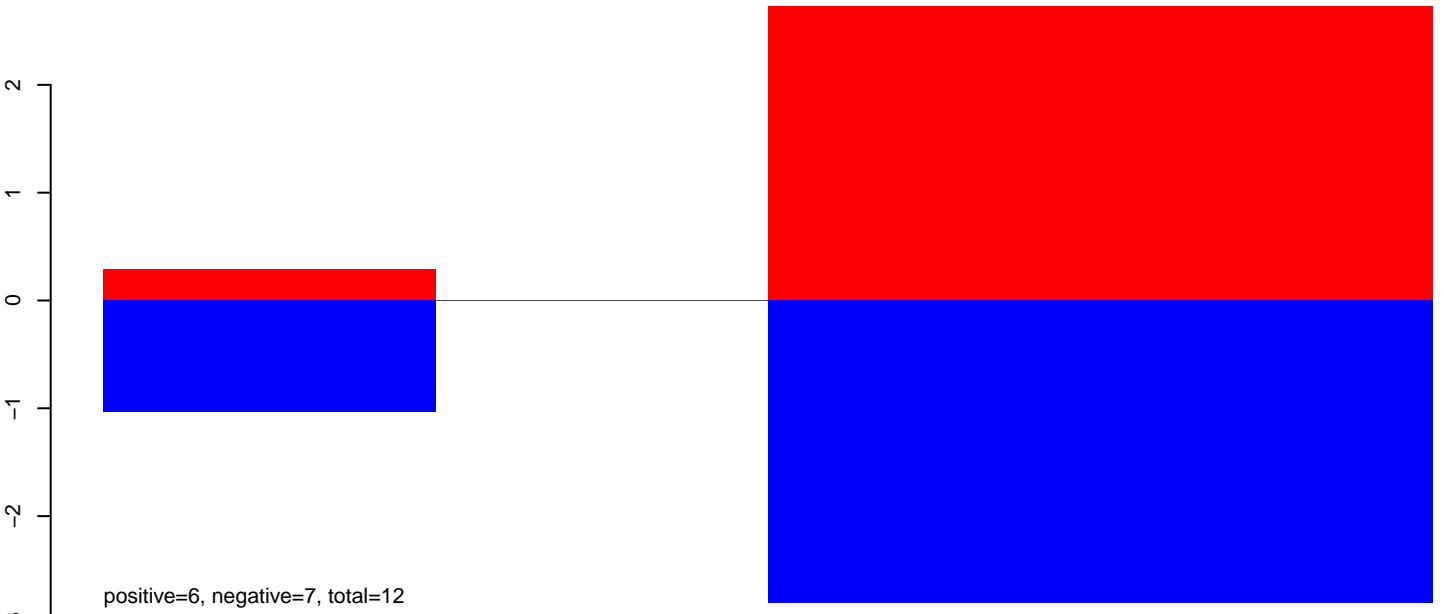
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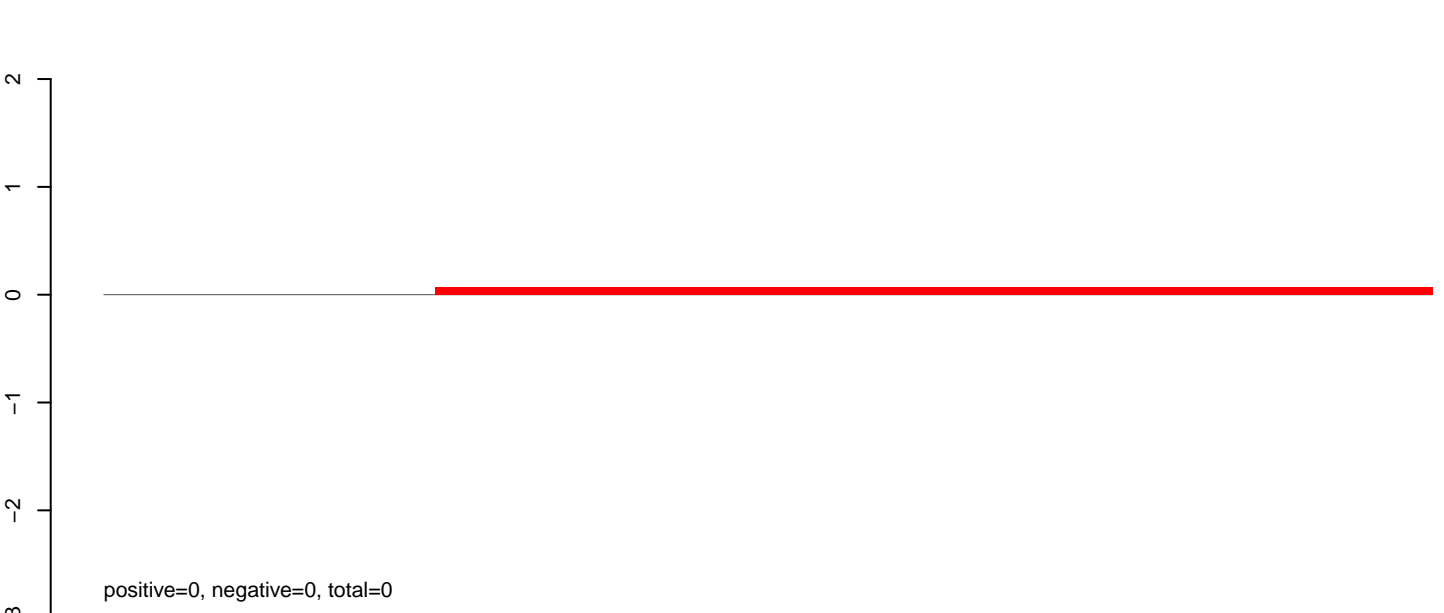
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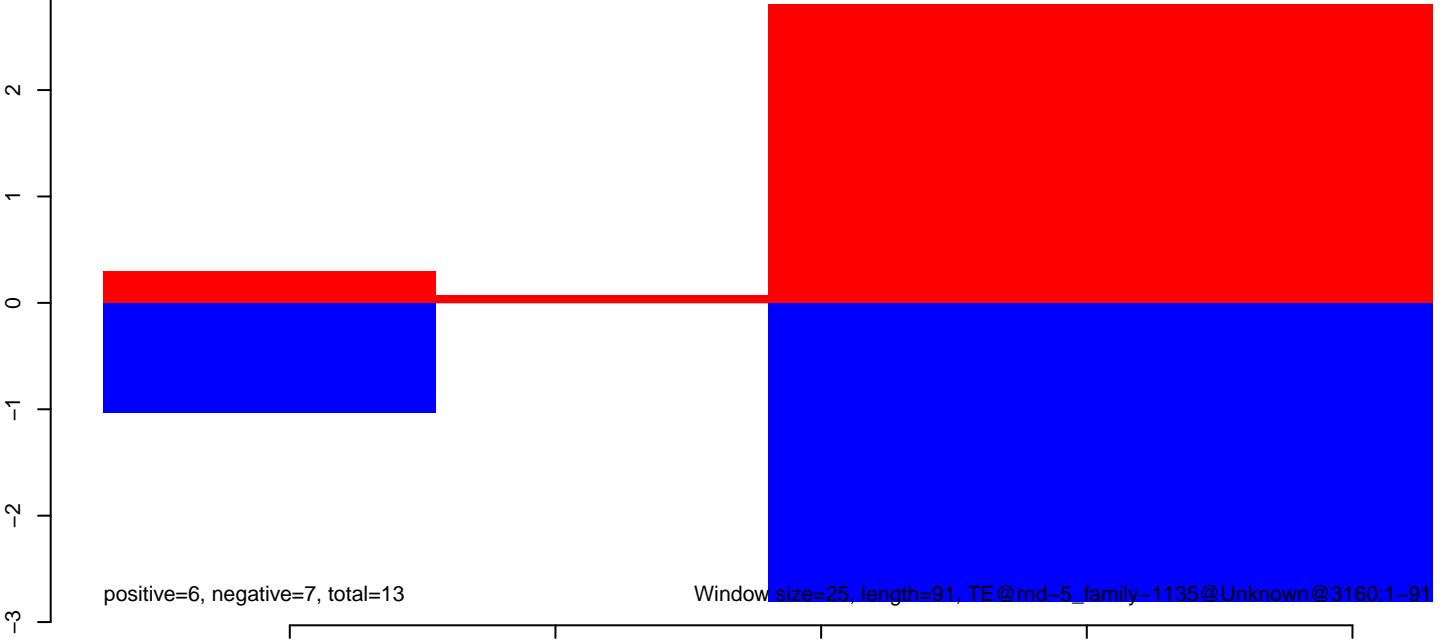
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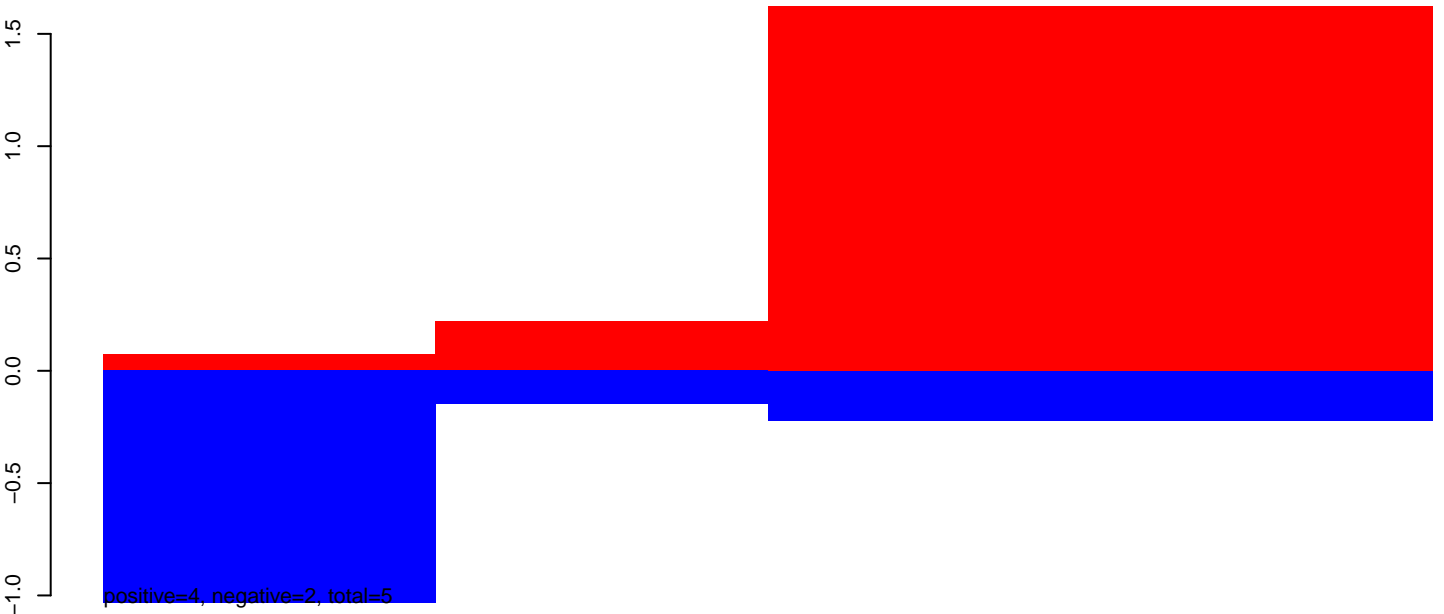
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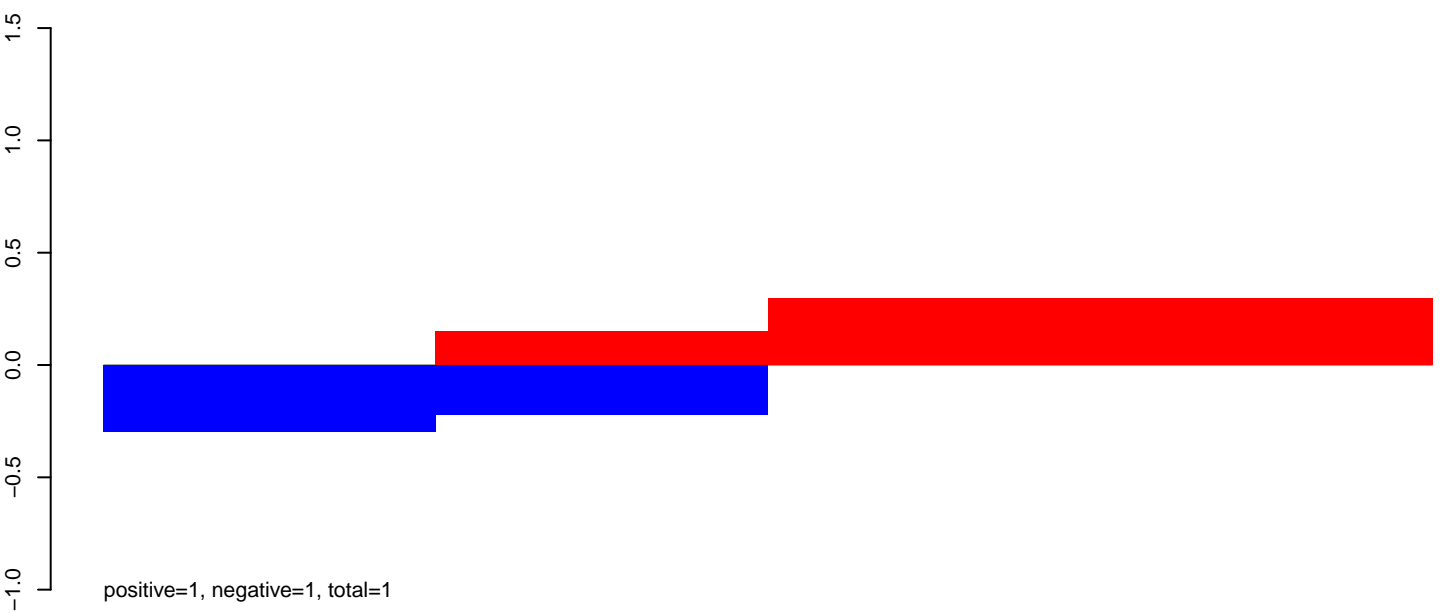
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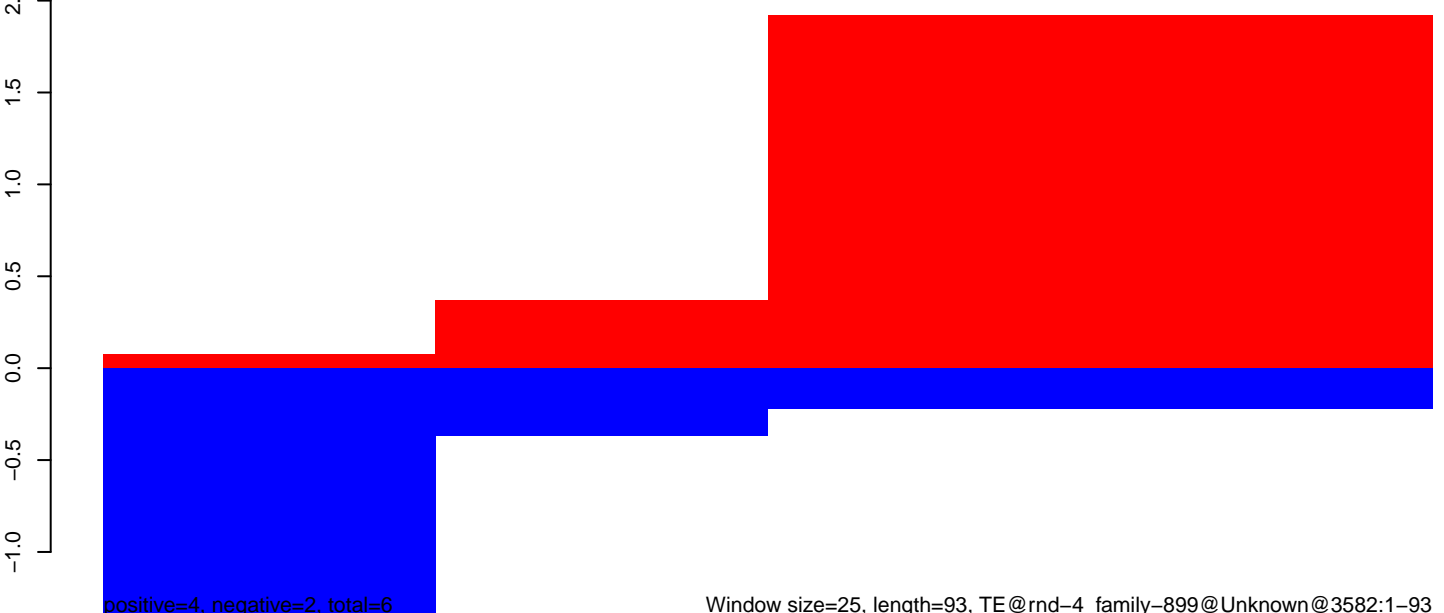
CuQuin_Testes_NL.18_23.rep



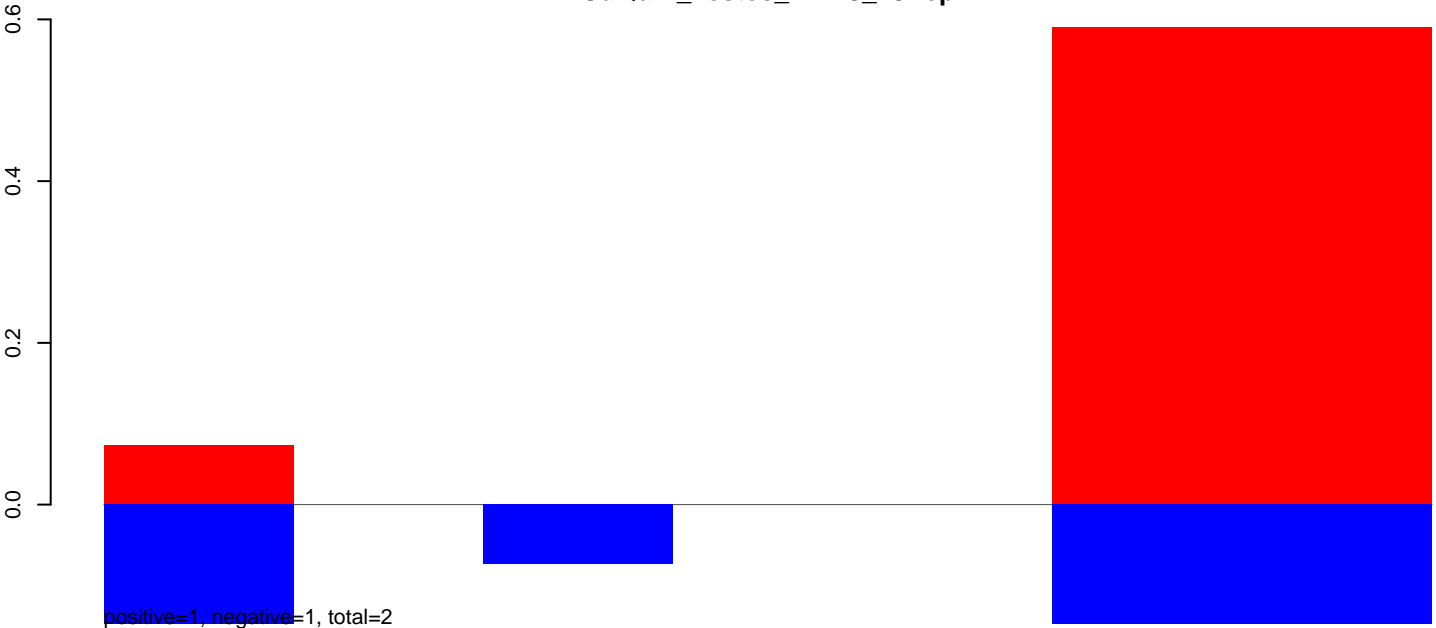
CuQuin_Testes_NL.24_35.rep



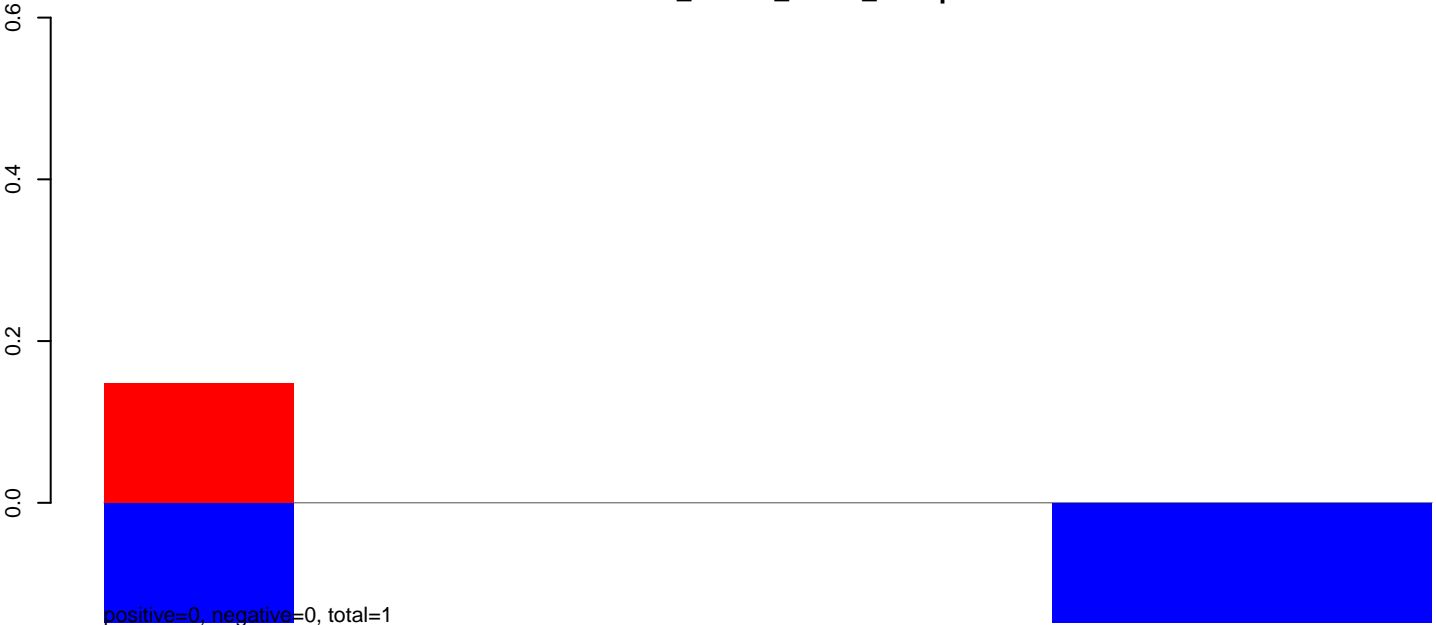
CuQuin_Testes_NL.rep



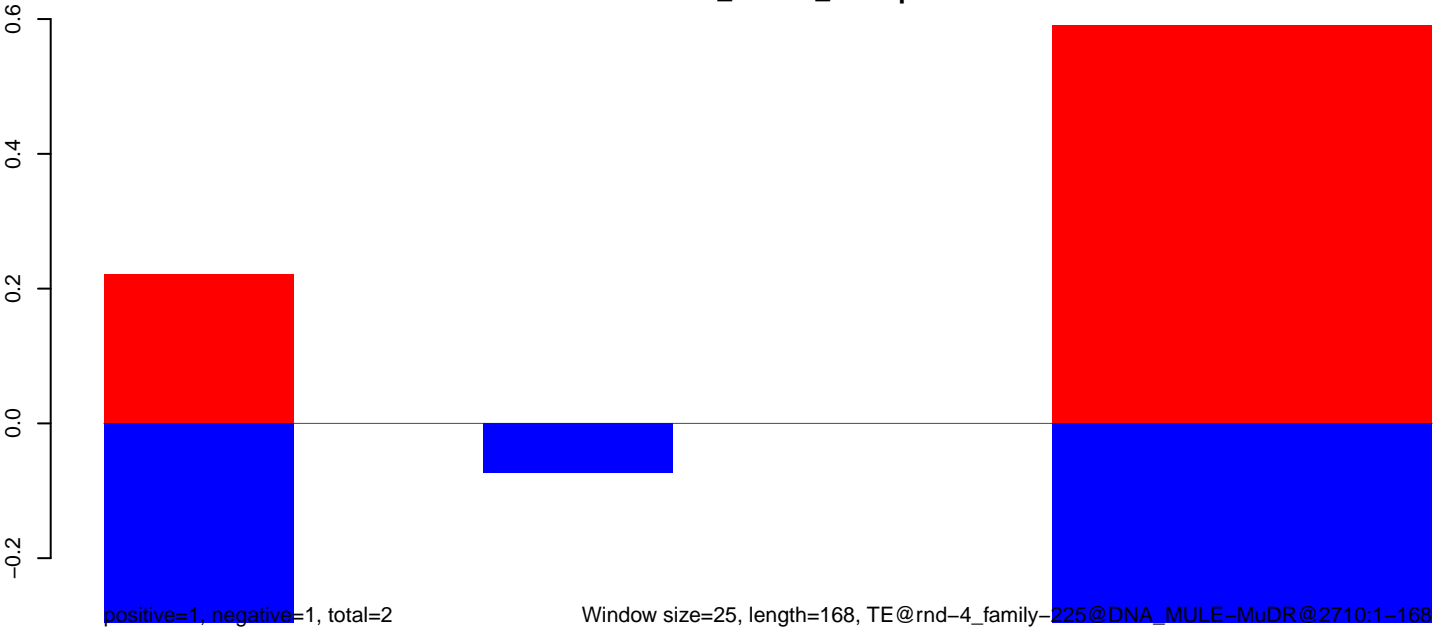
CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep



positive=1, negative=1, total=2

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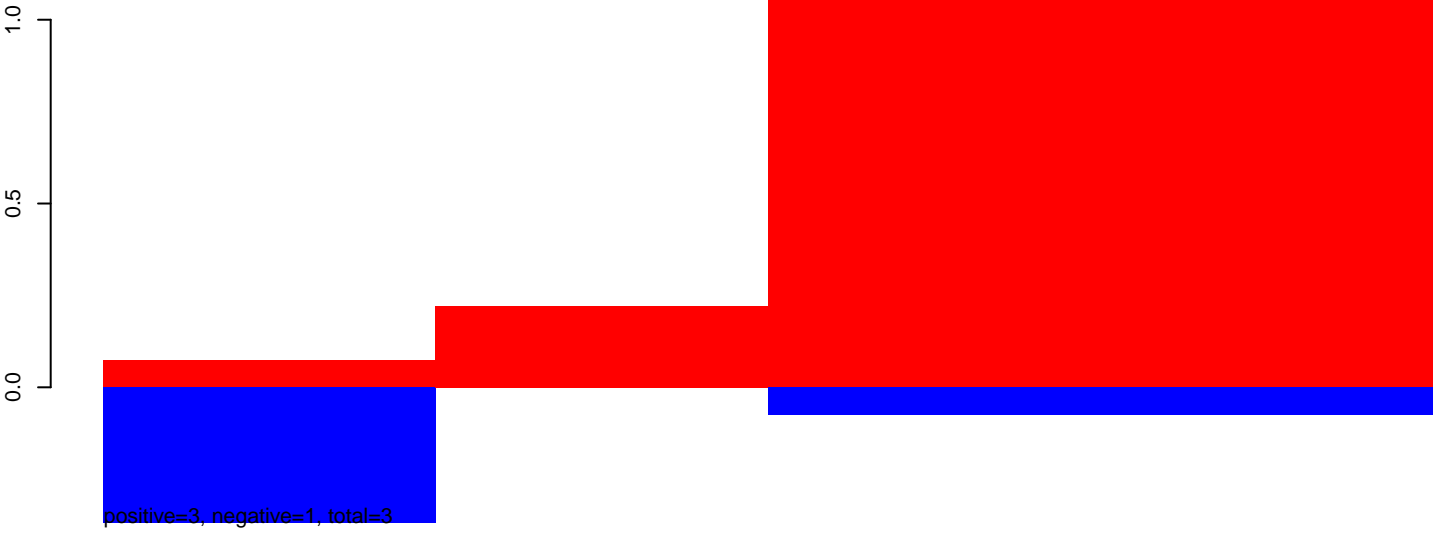
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100

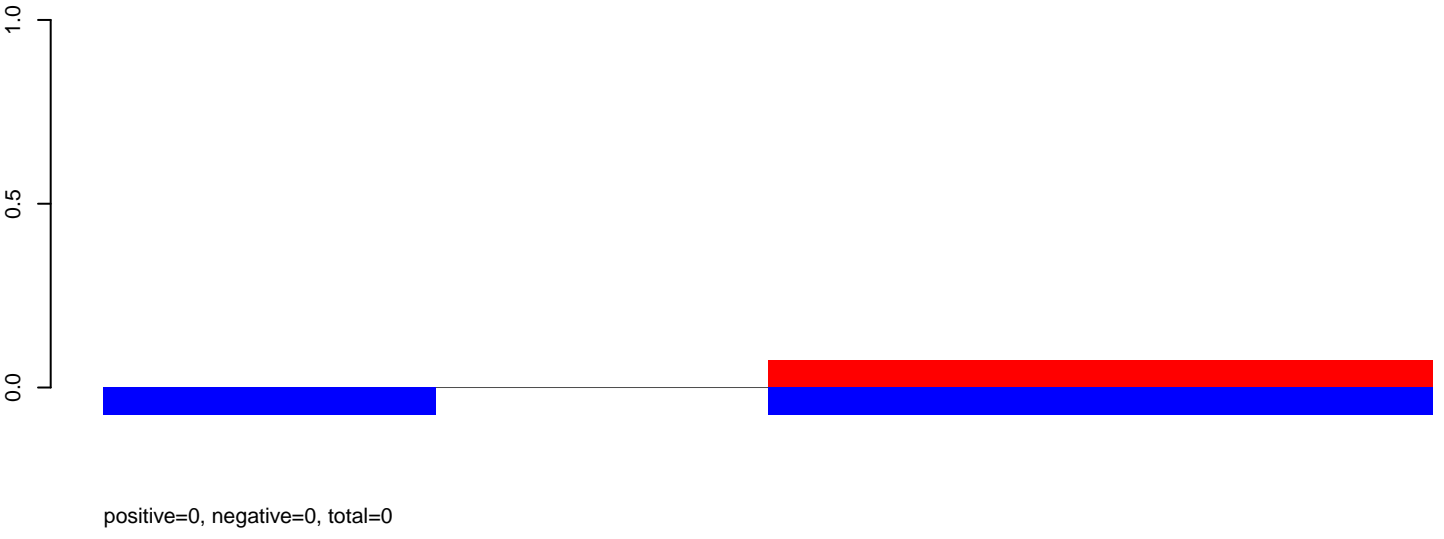
150

200

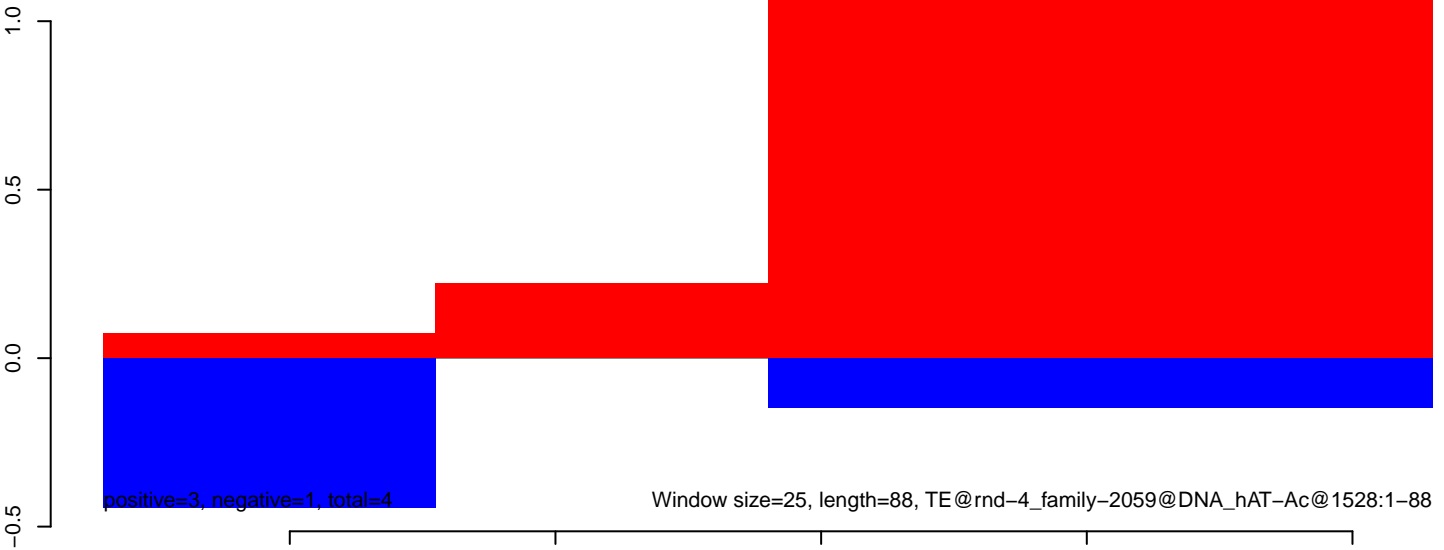
CuQuin_Testes_NL.18_23.rep



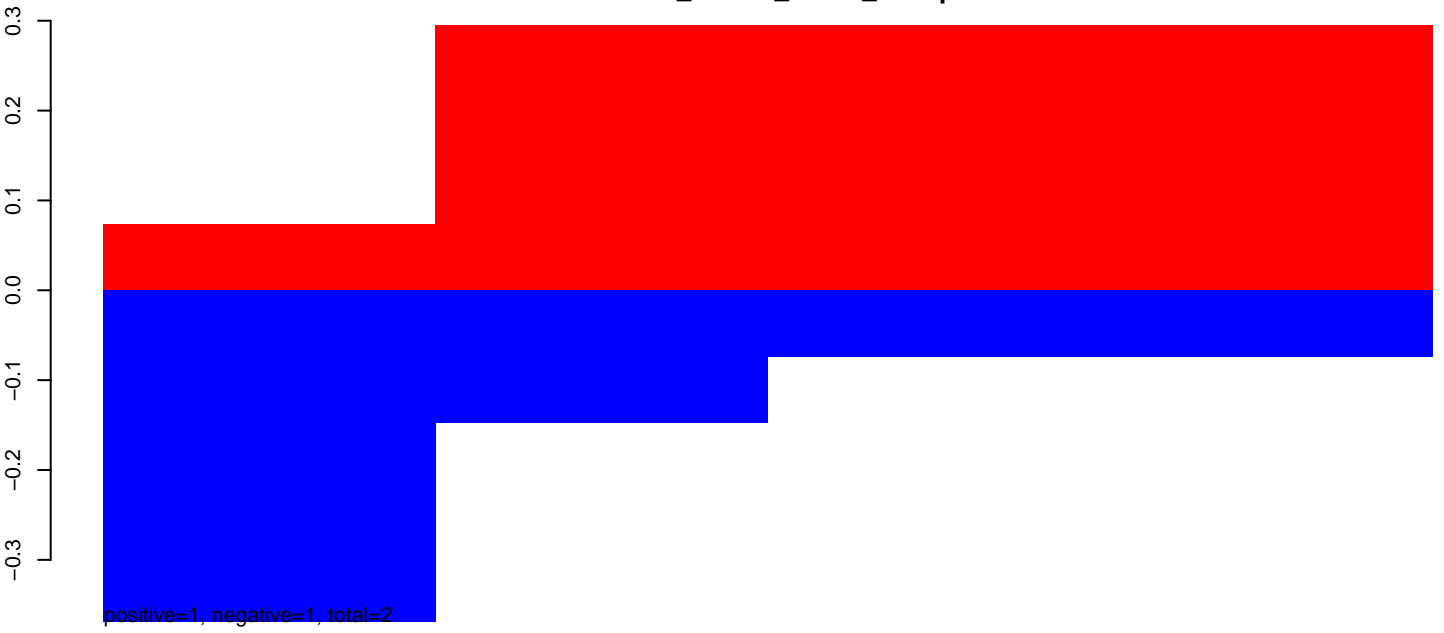
CuQuin_Testes_NL.24_35.rep



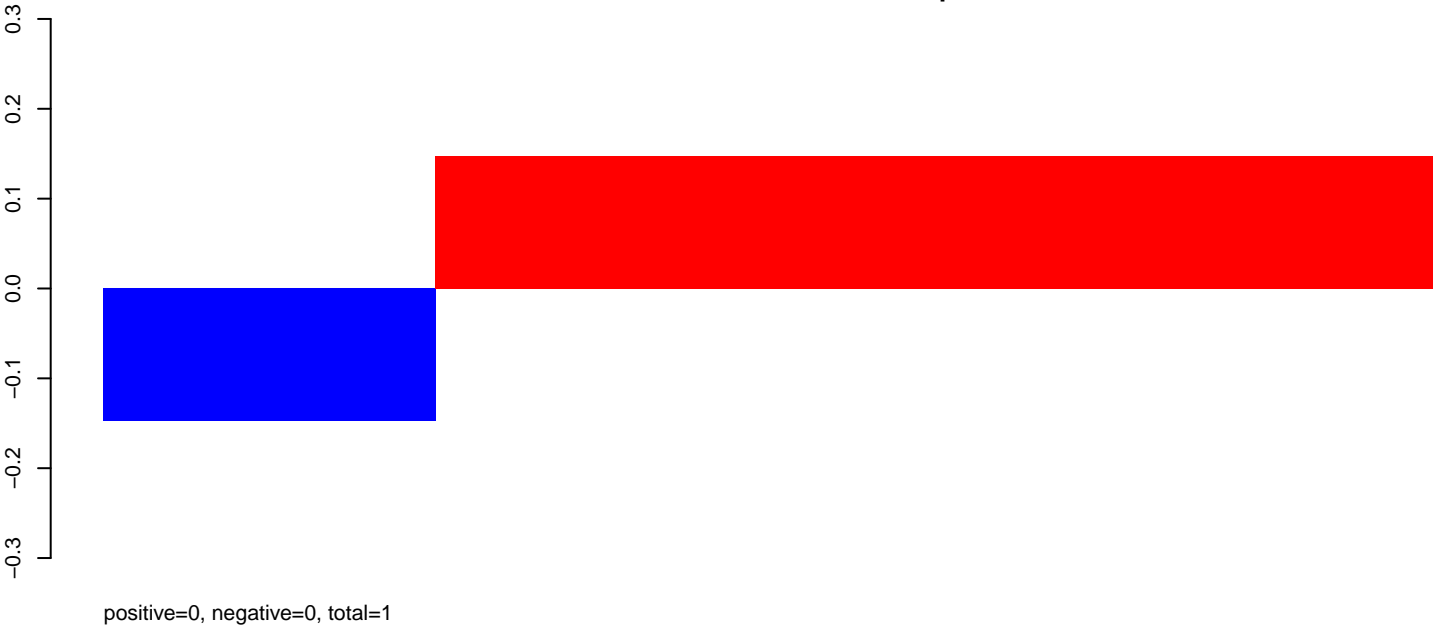
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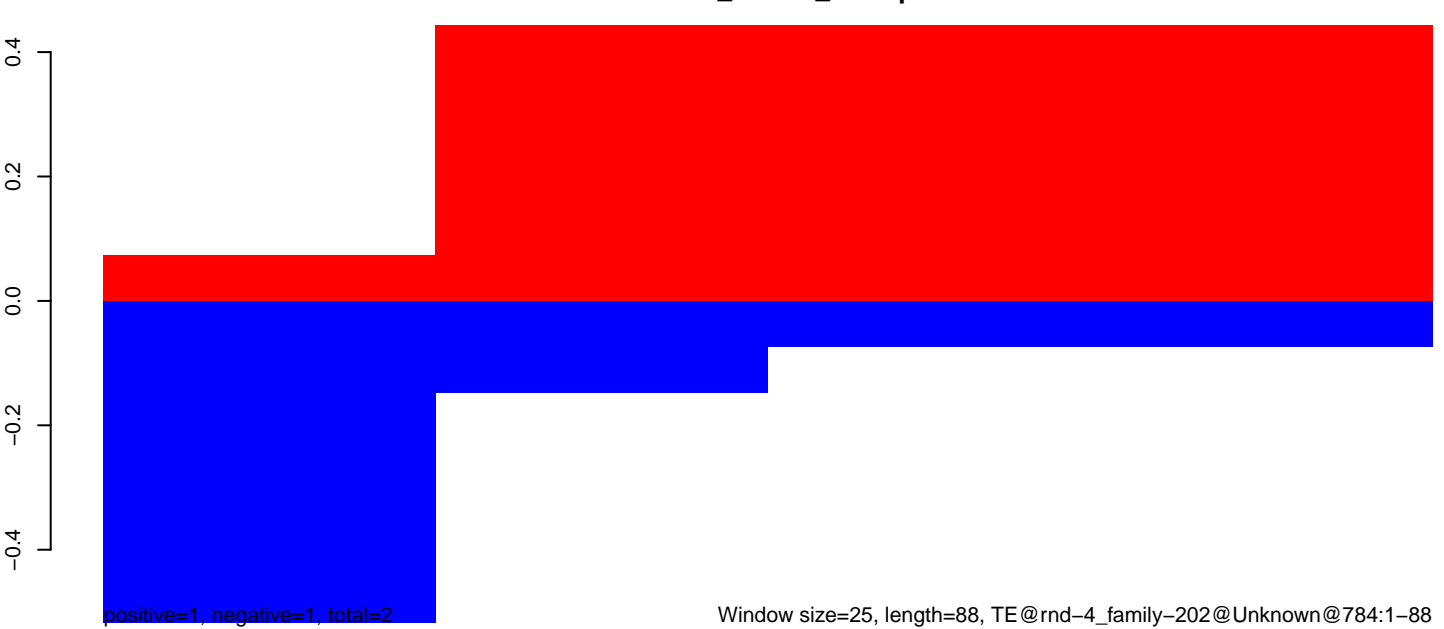
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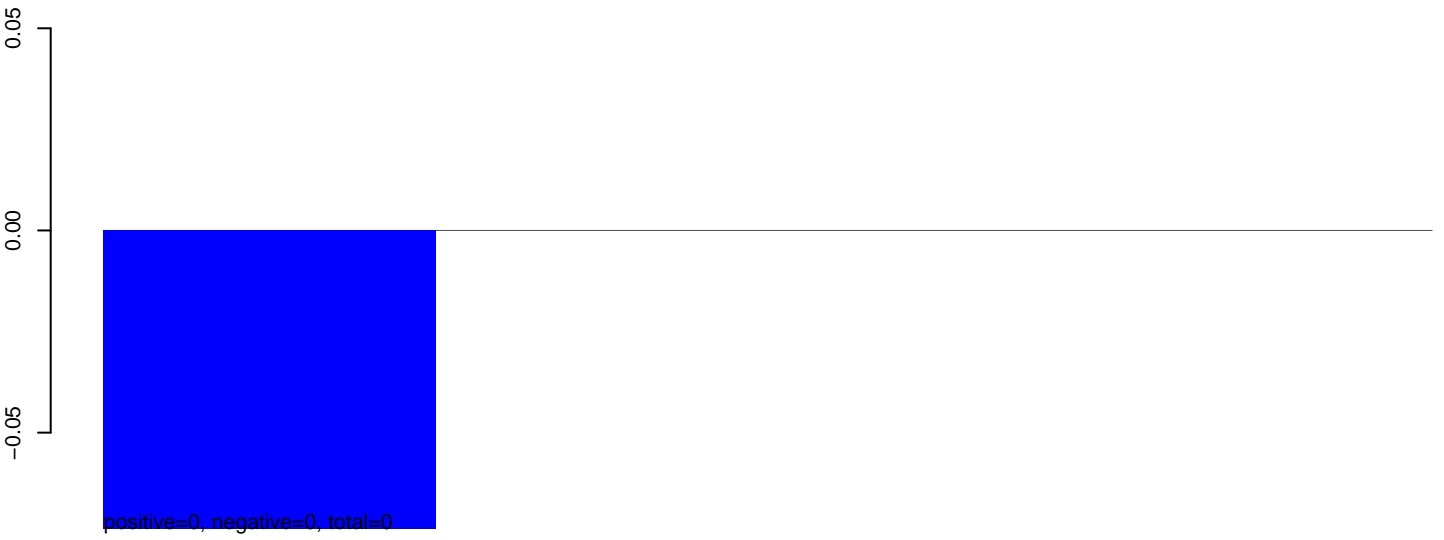
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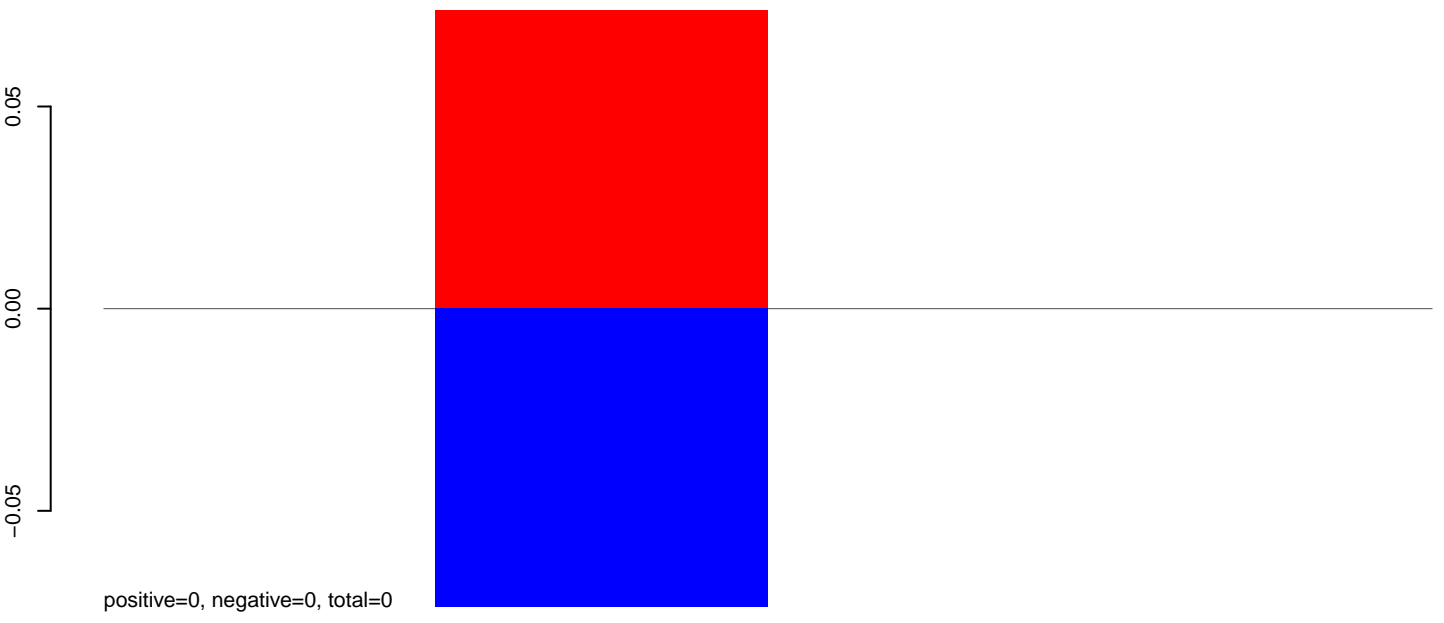
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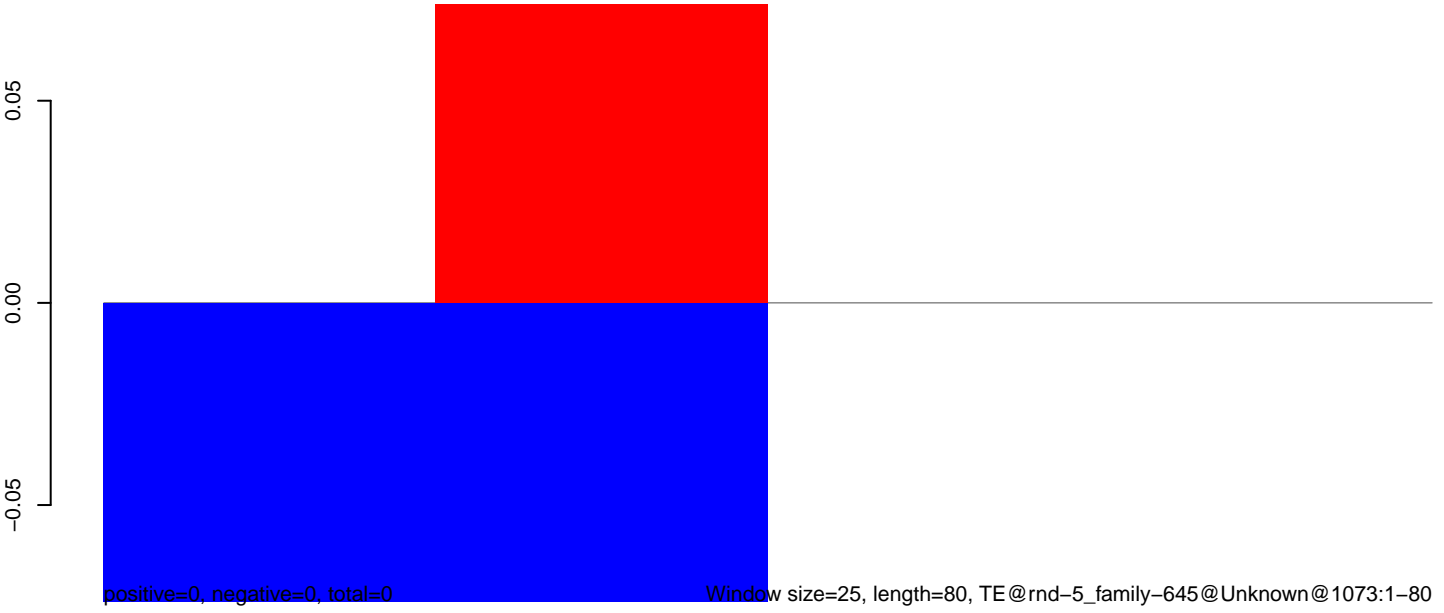
CuQuin_Testes_NL.18_23.rep



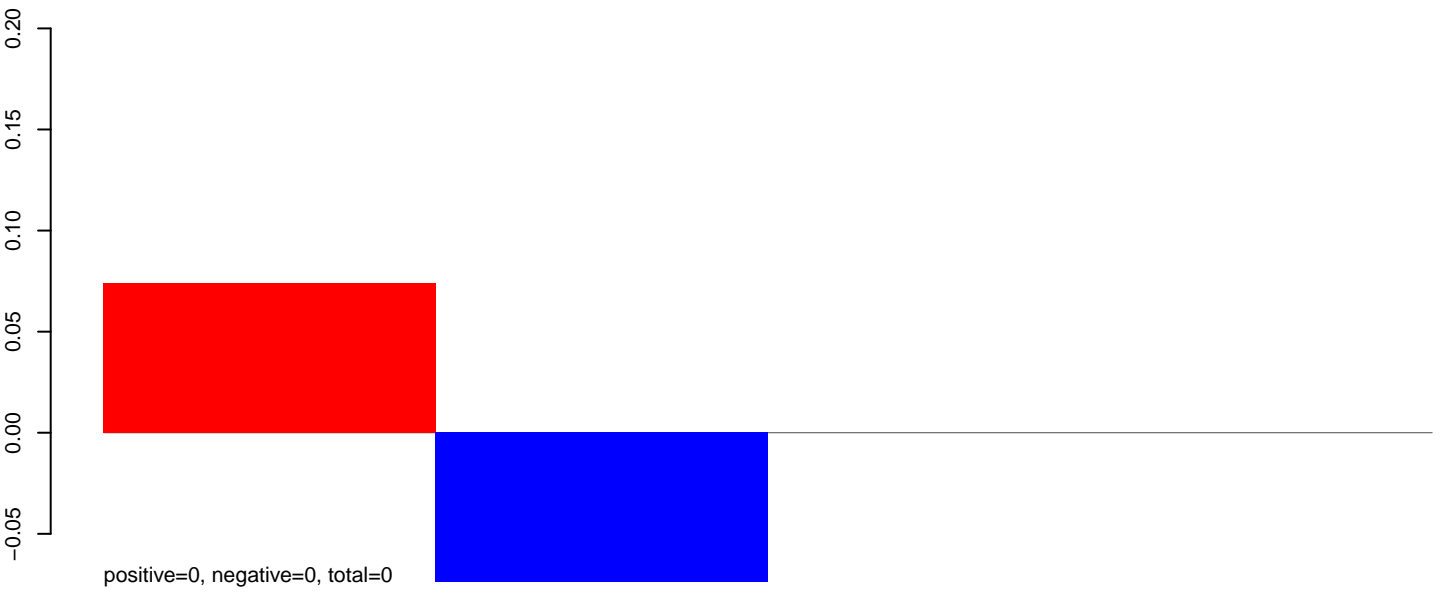
CuQuin_Testes_NL.24_35.rep



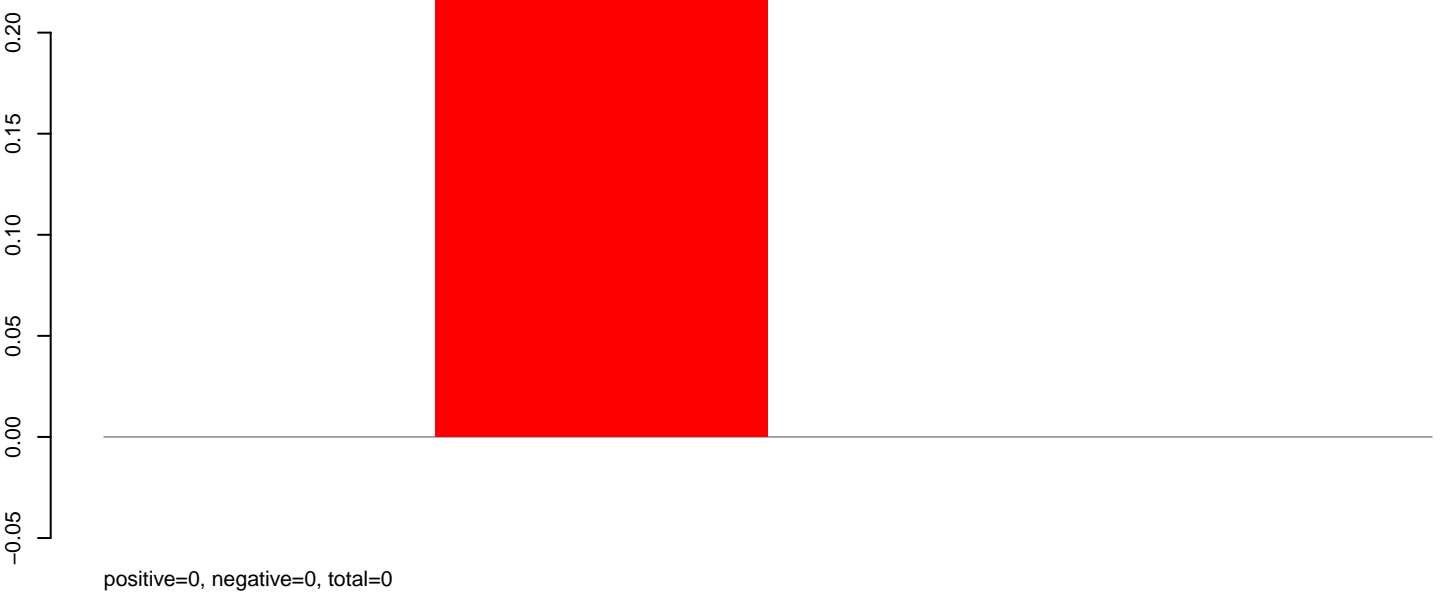
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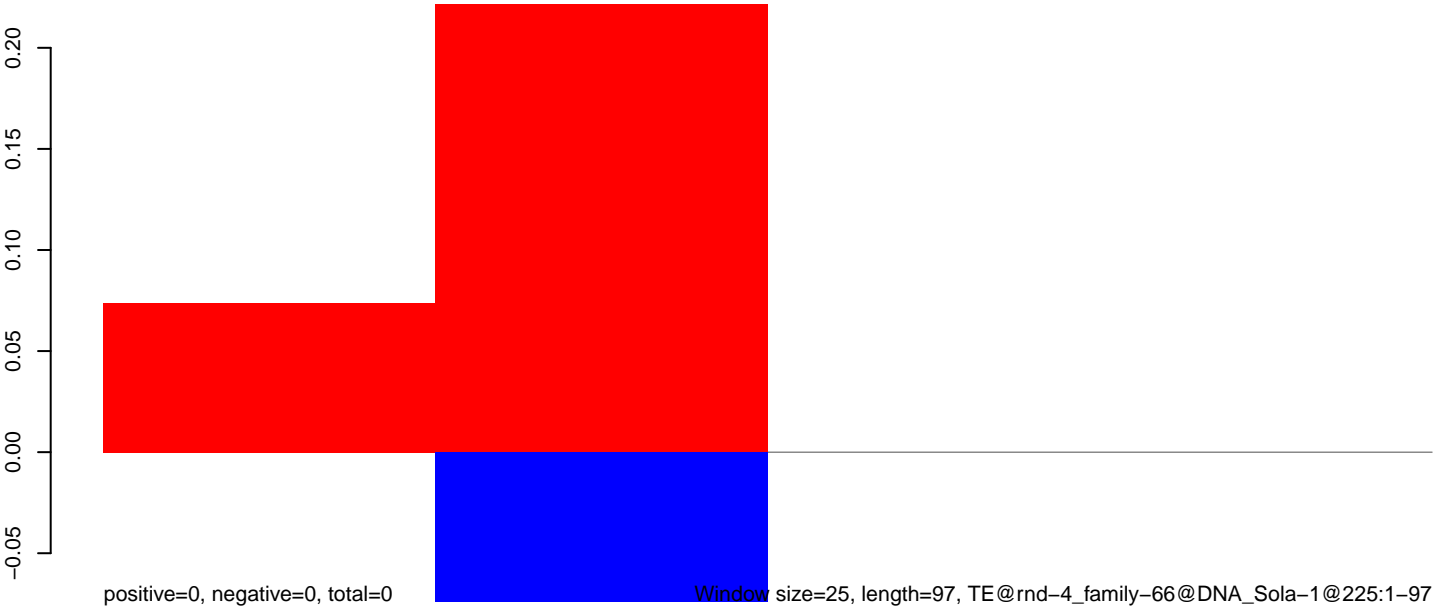
CuQuin_Testes_NL.18_23.rep



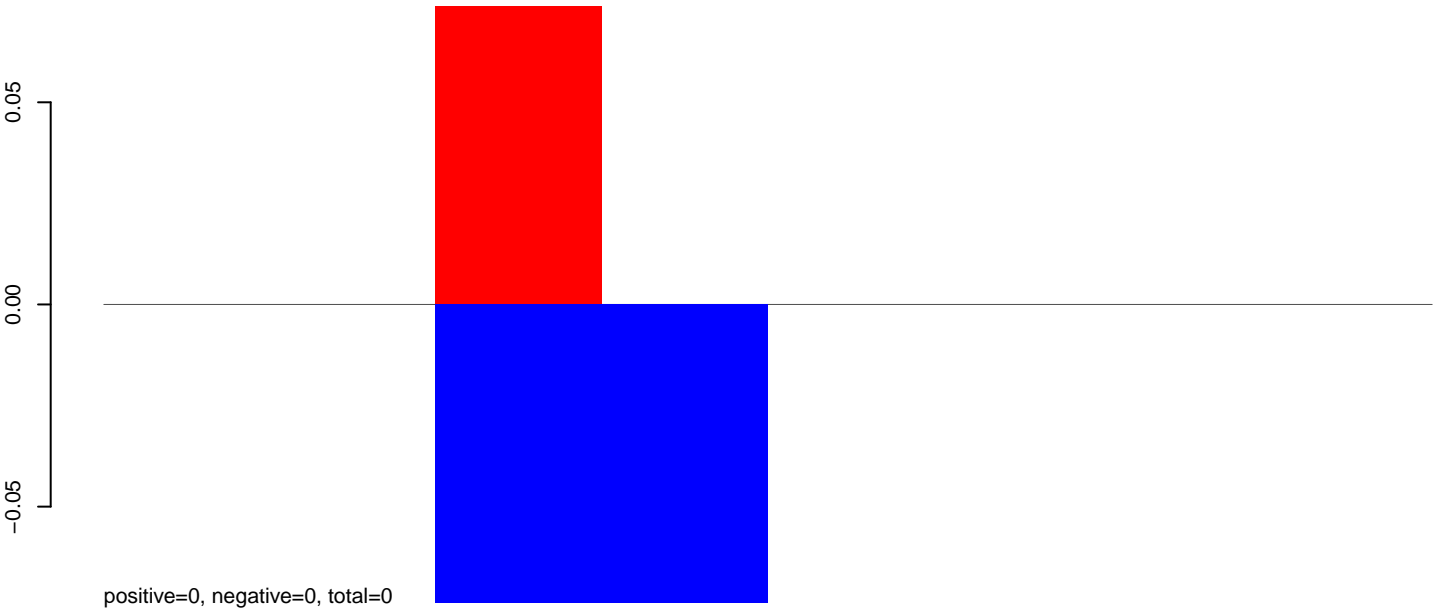
CuQuin_Testes_NL.24_35.rep



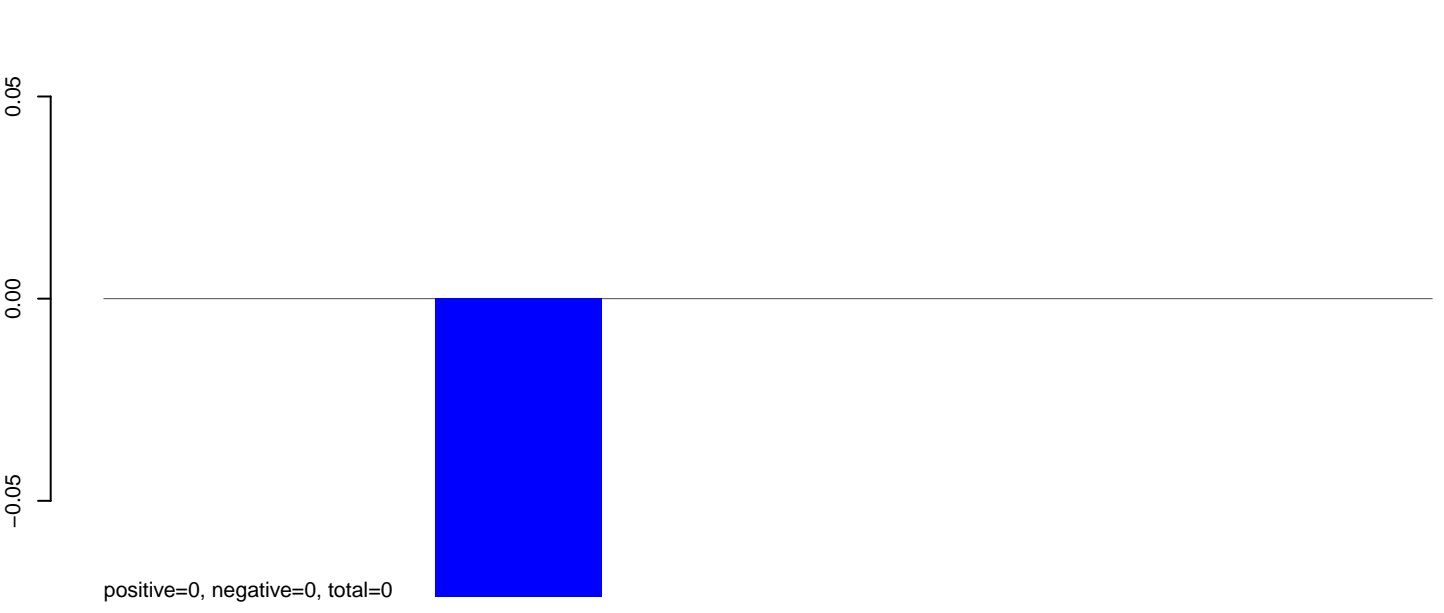
CuQuin_Testes_NL.rep



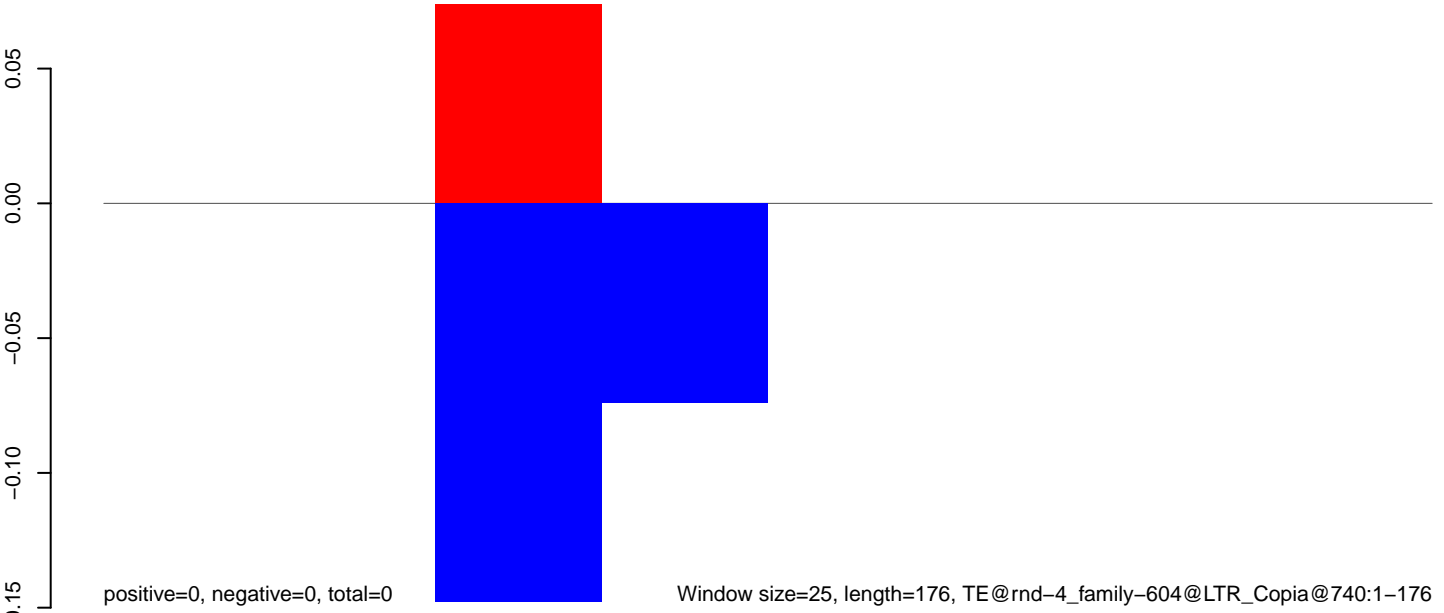
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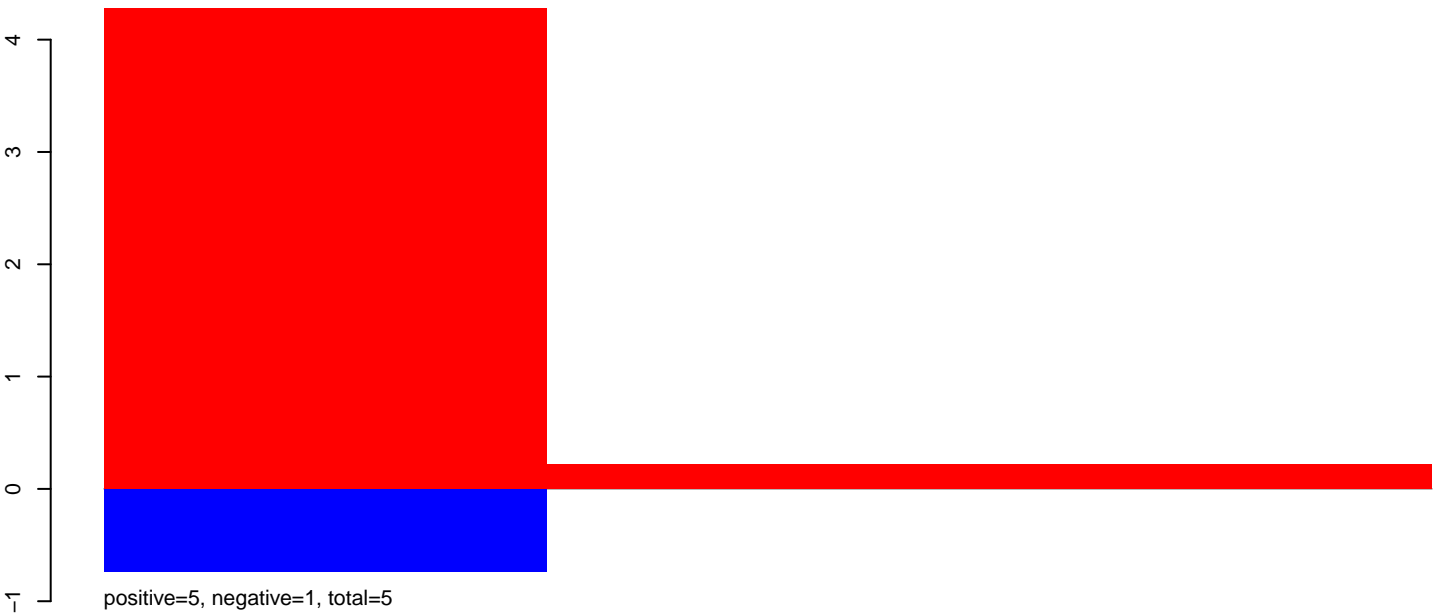
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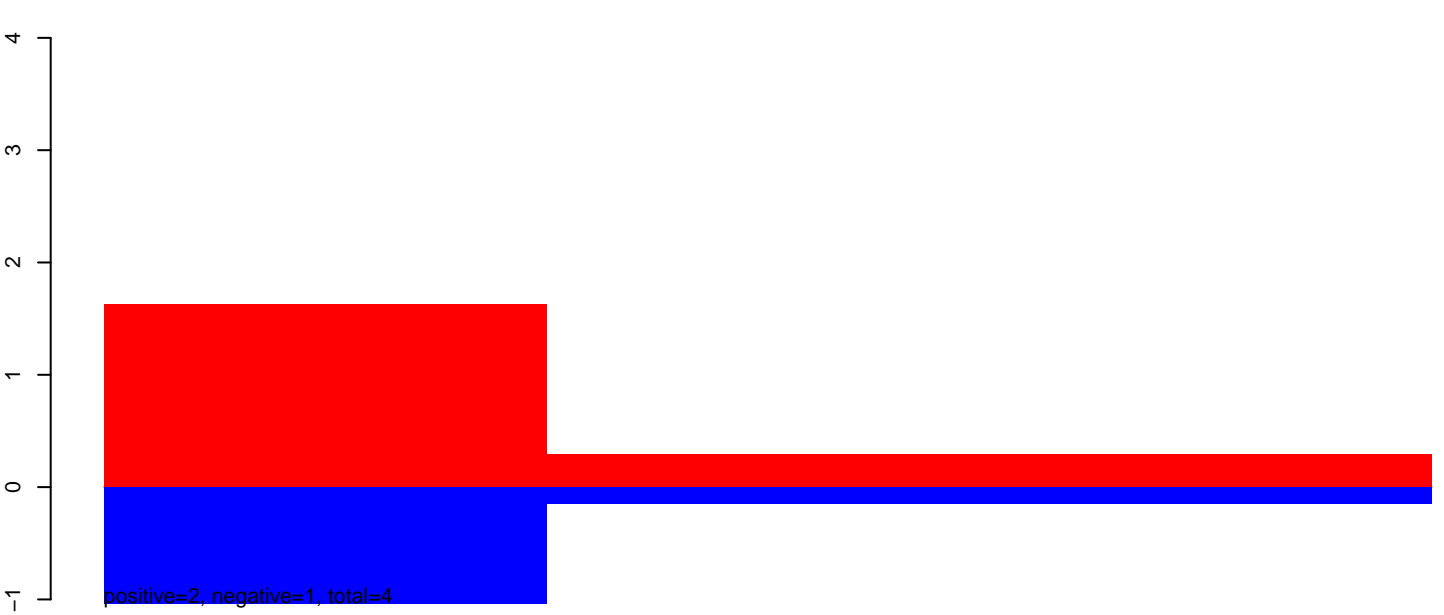
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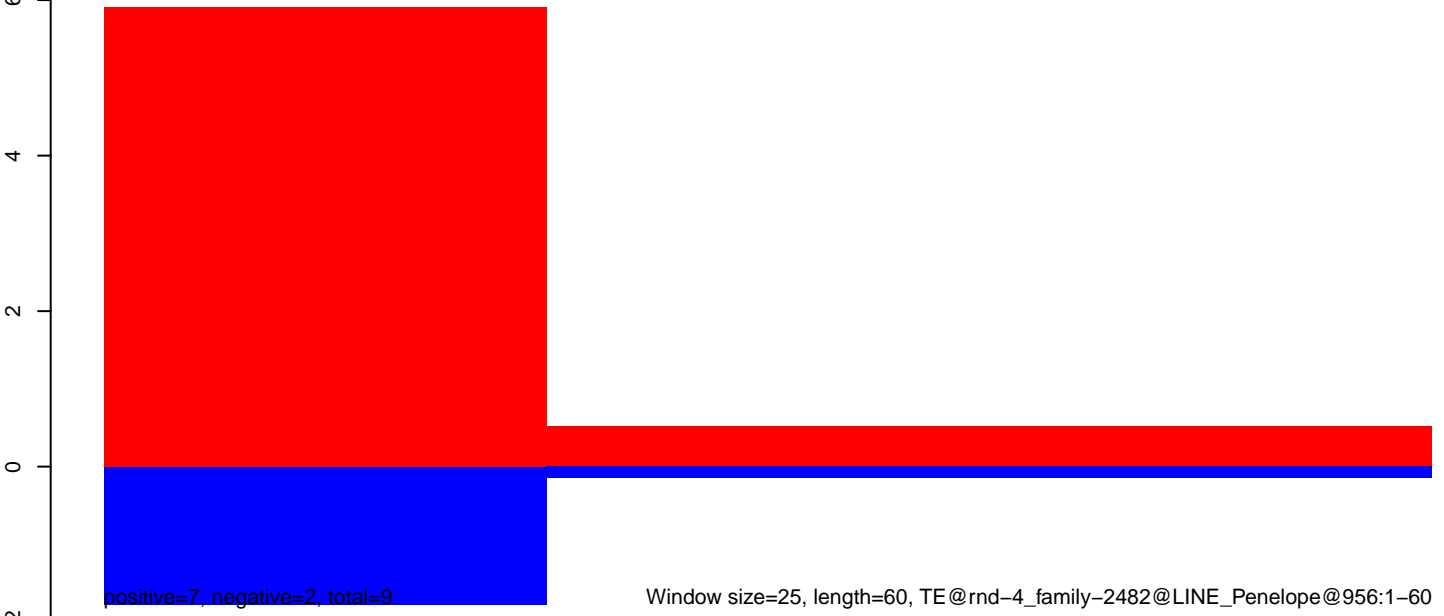
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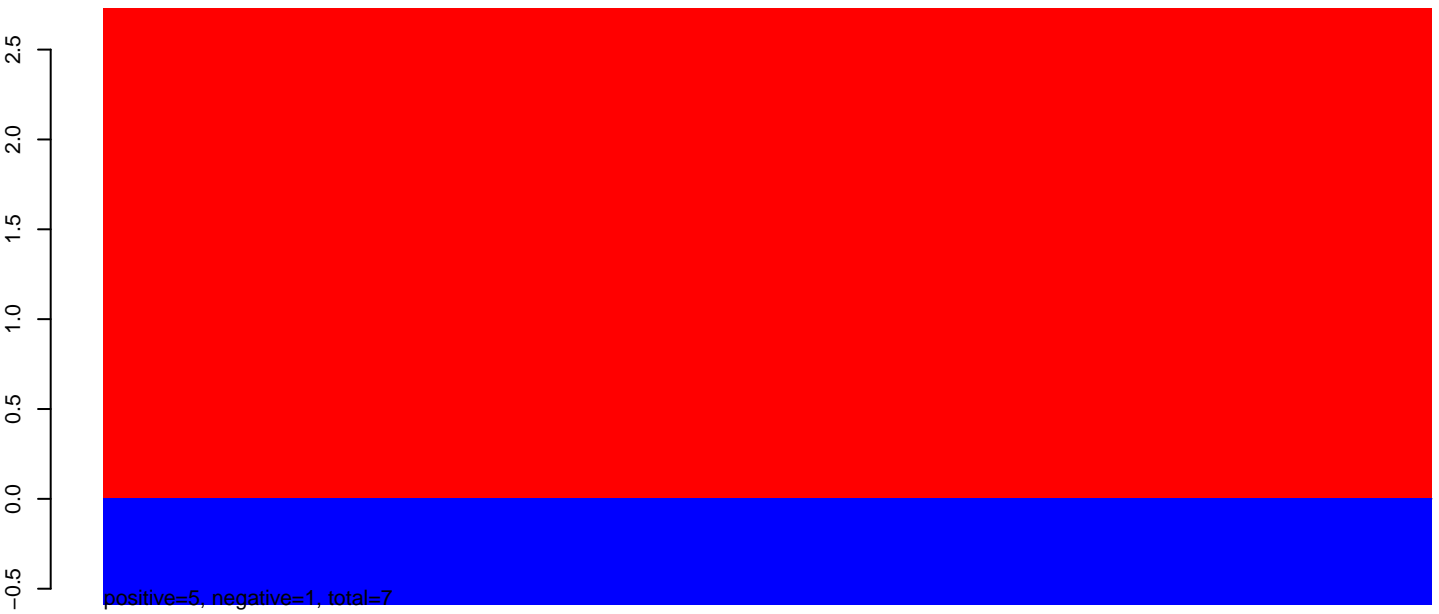
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CuQuin_Testes_NL.rep



CuQuin_Testes_NL.18_23.rep



CuQuin_Testes_NL.24_35.rep



CuQuin_Testes_NL.rep

