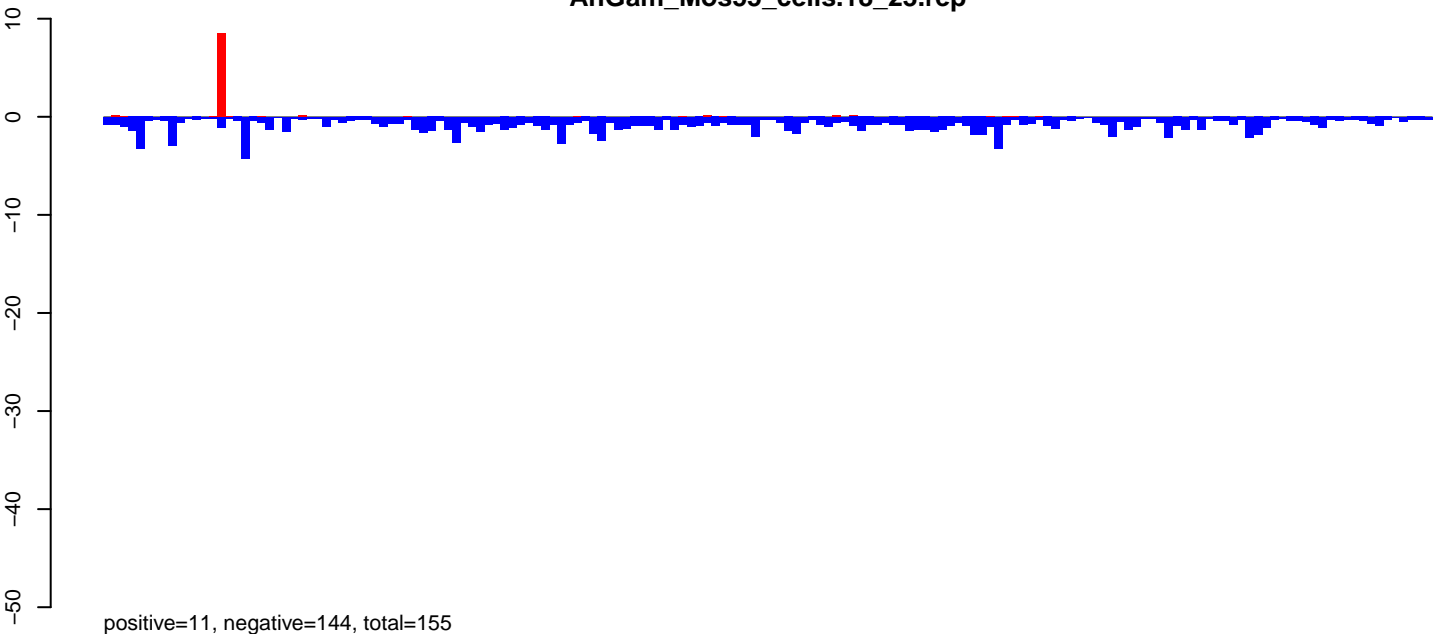
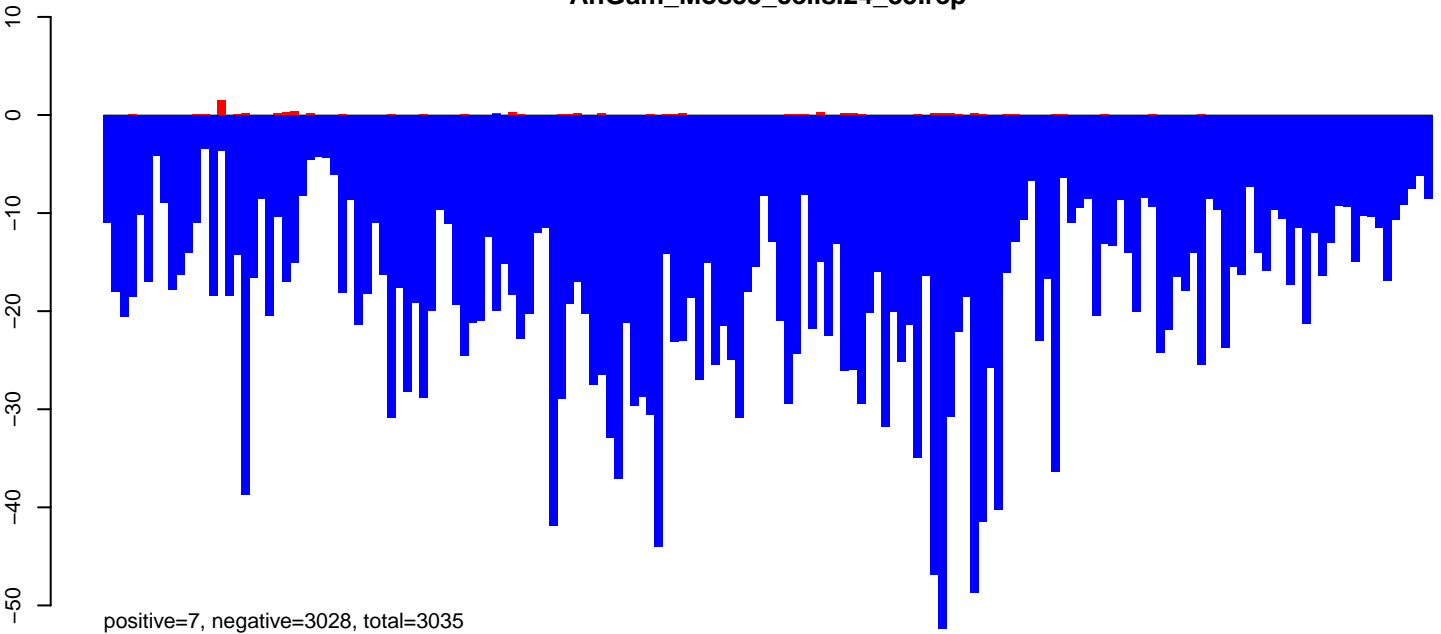


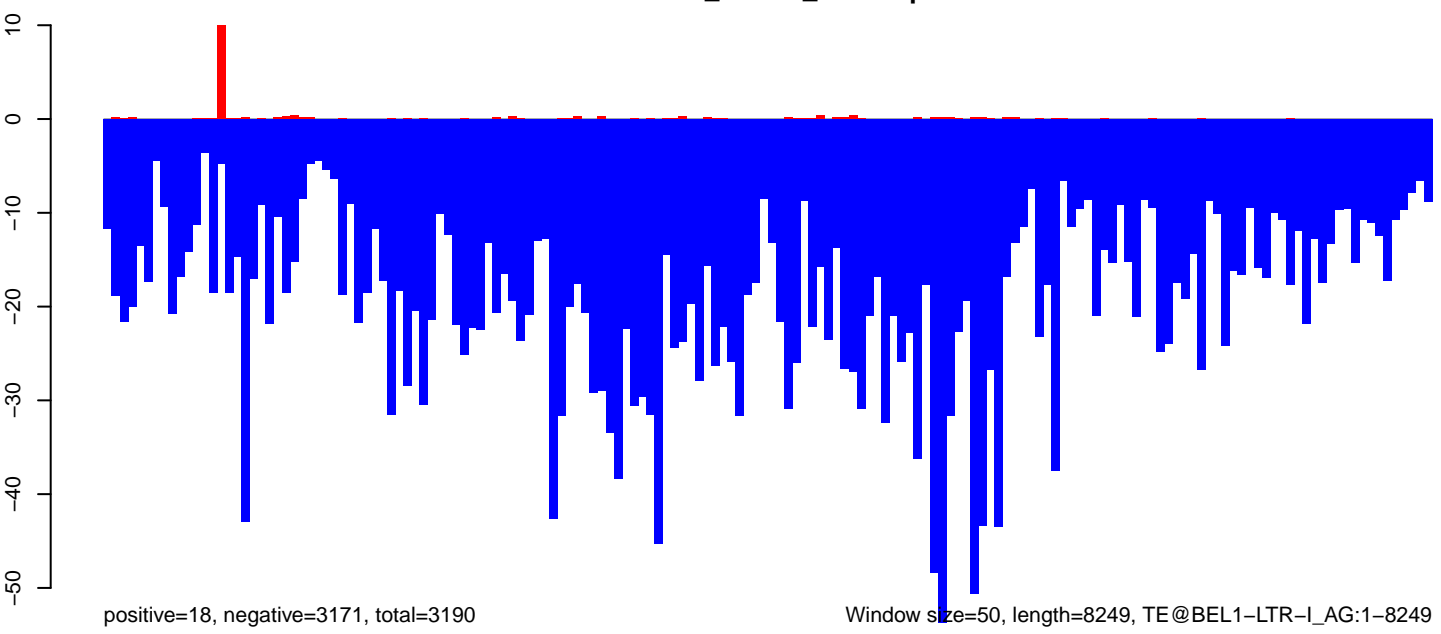
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



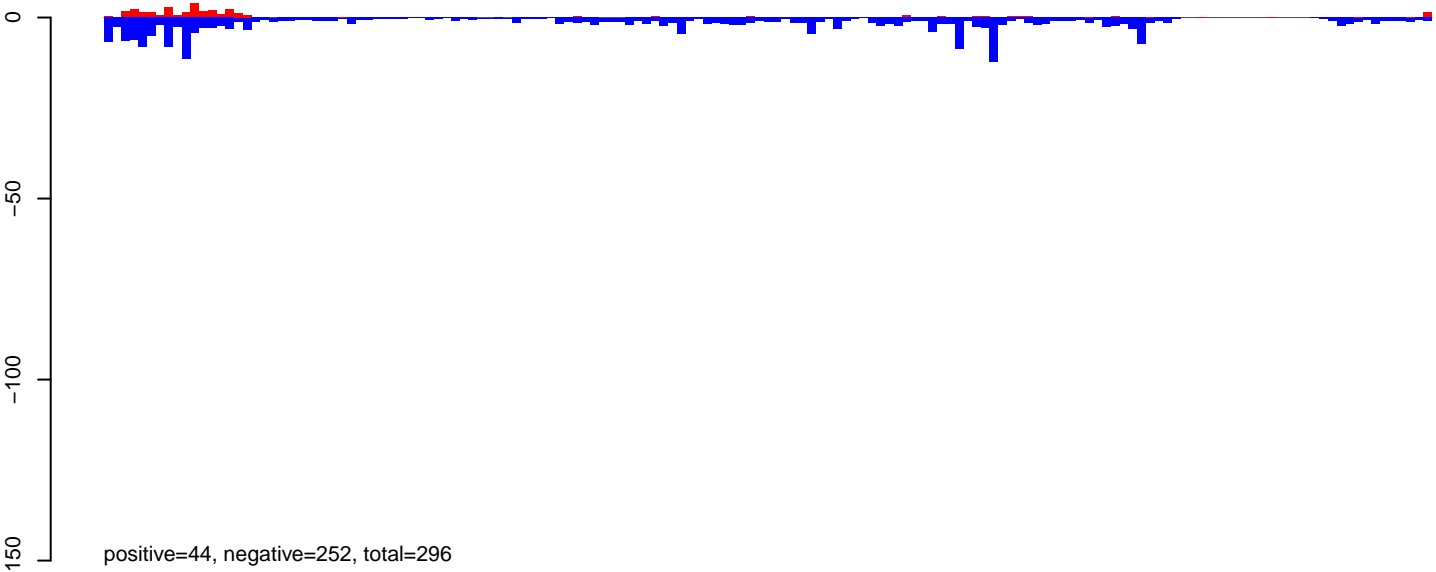
AnGam_Mos55_cells.rep



Window size=50, length=8249, TE@BEL1-LTR-I_AG:1-8249

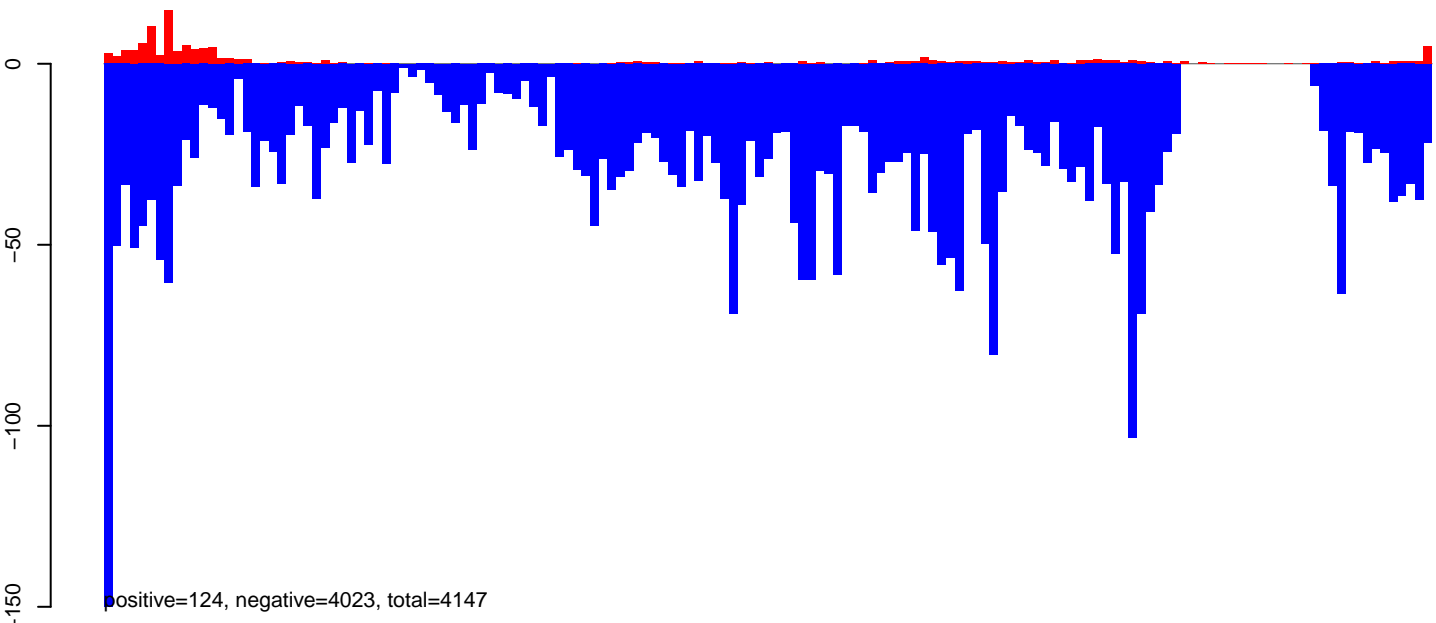
0 2000 4000 6000 8000

AnGam_Mos55_cells.18_23.rep



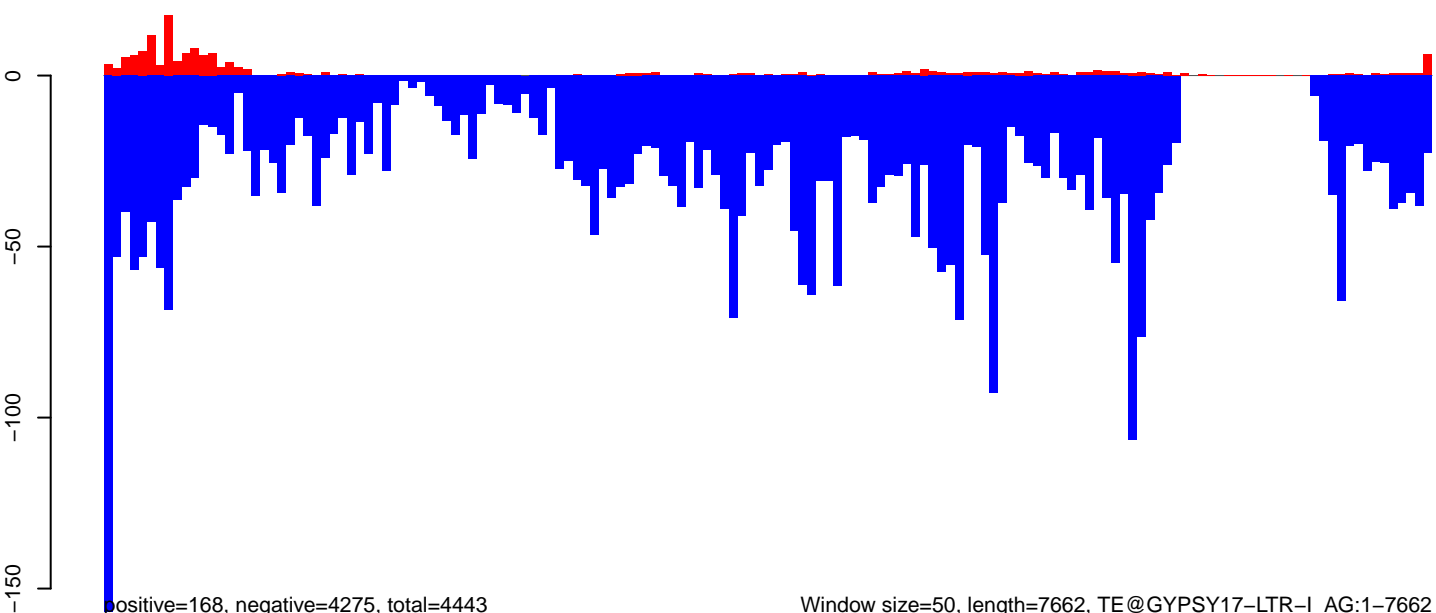
positive=44, negative=252, total=296

AnGam_Mos55_cells.24_35.rep



positive=124, negative=4023, total=4147

AnGam_Mos55_cells.rep

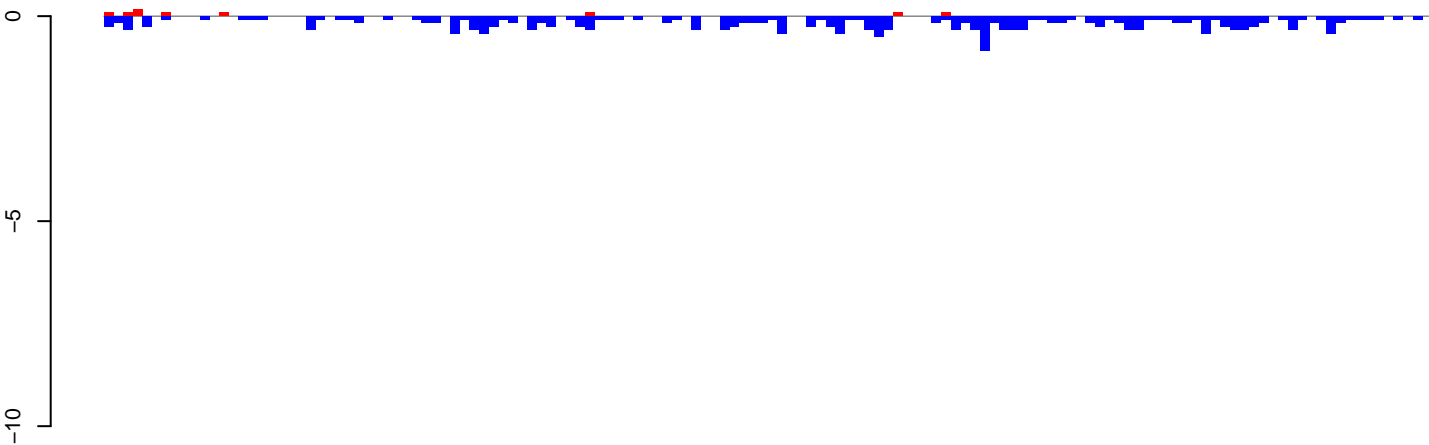


positive=168, negative=4275, total=4443

Window size=50, length=7662, TE@GYPSY17-LTR-I_AG:1-7662

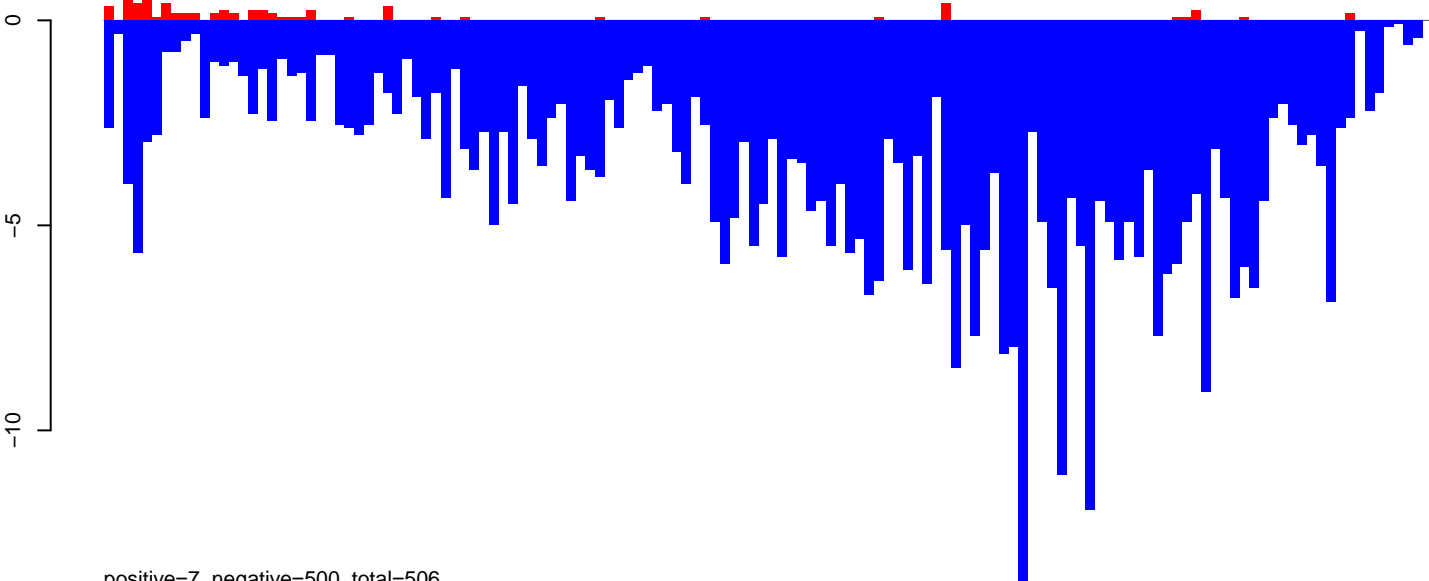
0 2000 4000 6000 8000

AnGam_Mos55_cells.18_23.rep



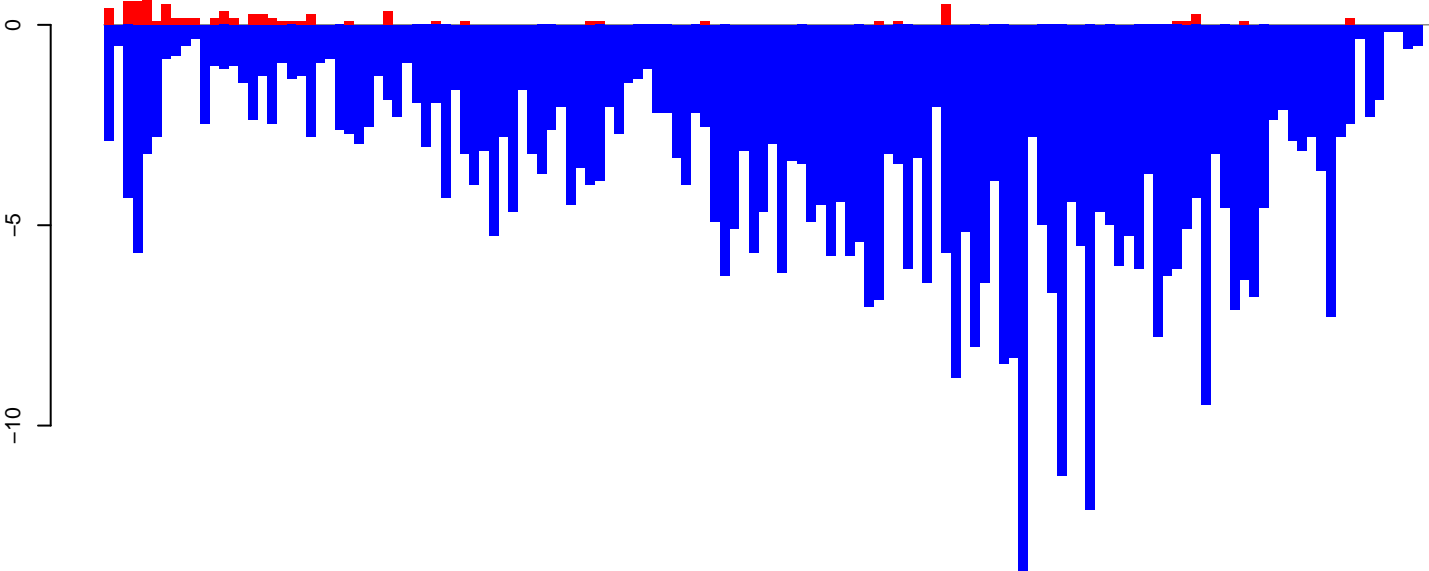
positive=1, negative=20, total=21

AnGam_Mos55_cells.24_35.rep



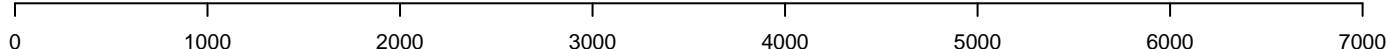
positive=7, negative=500, total=506

AnGam_Mos55_cells.rep

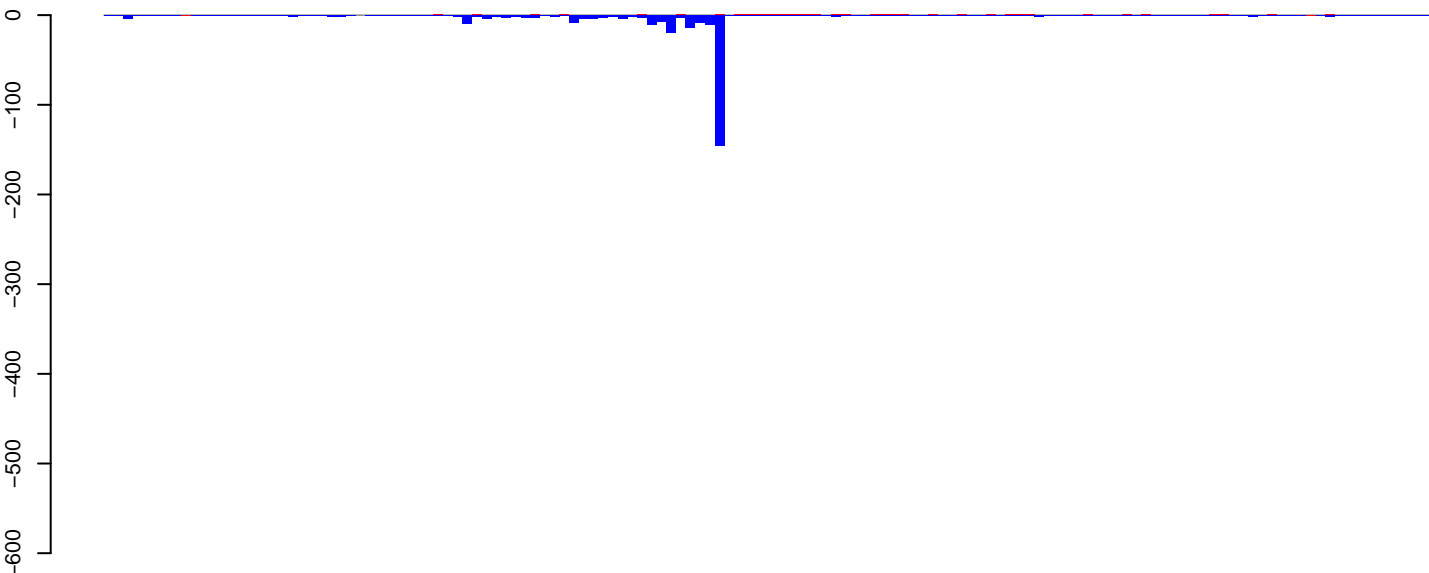


positive=8, negative=520, total=527

Window size=50, length=6912, TE@GYPSY41-LTR-L_AG:1-6912

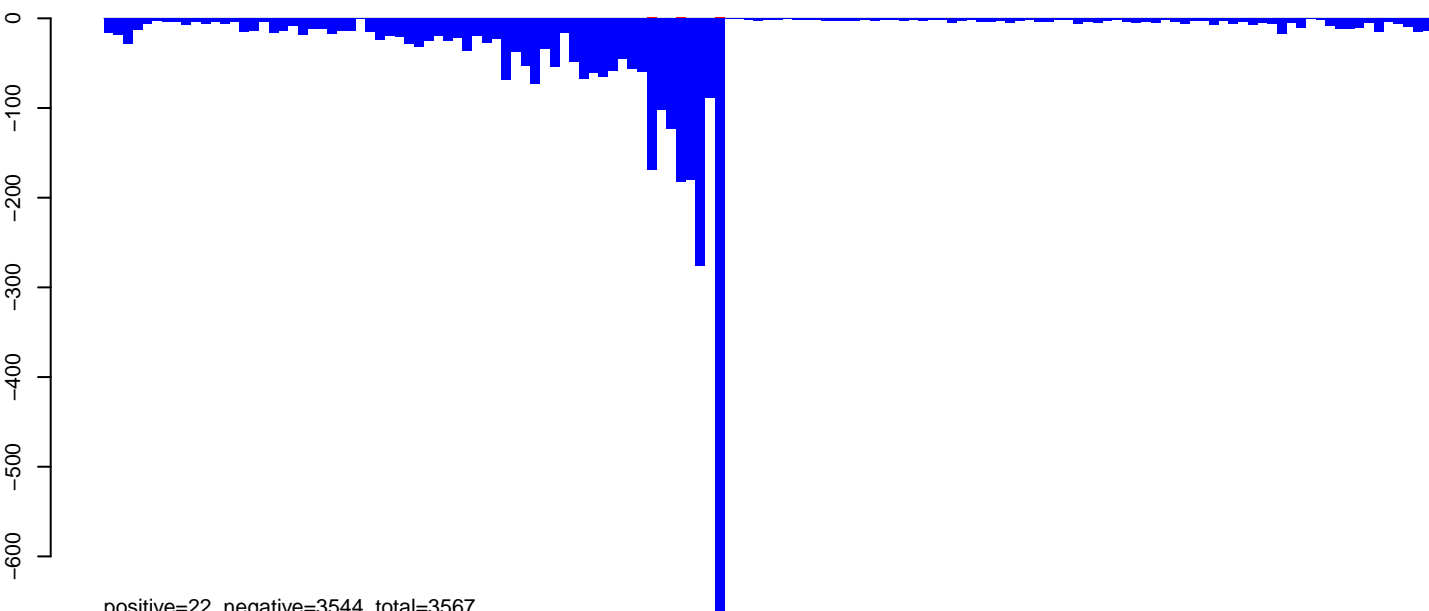


AnGam_Mos55_cells.18_23.rep



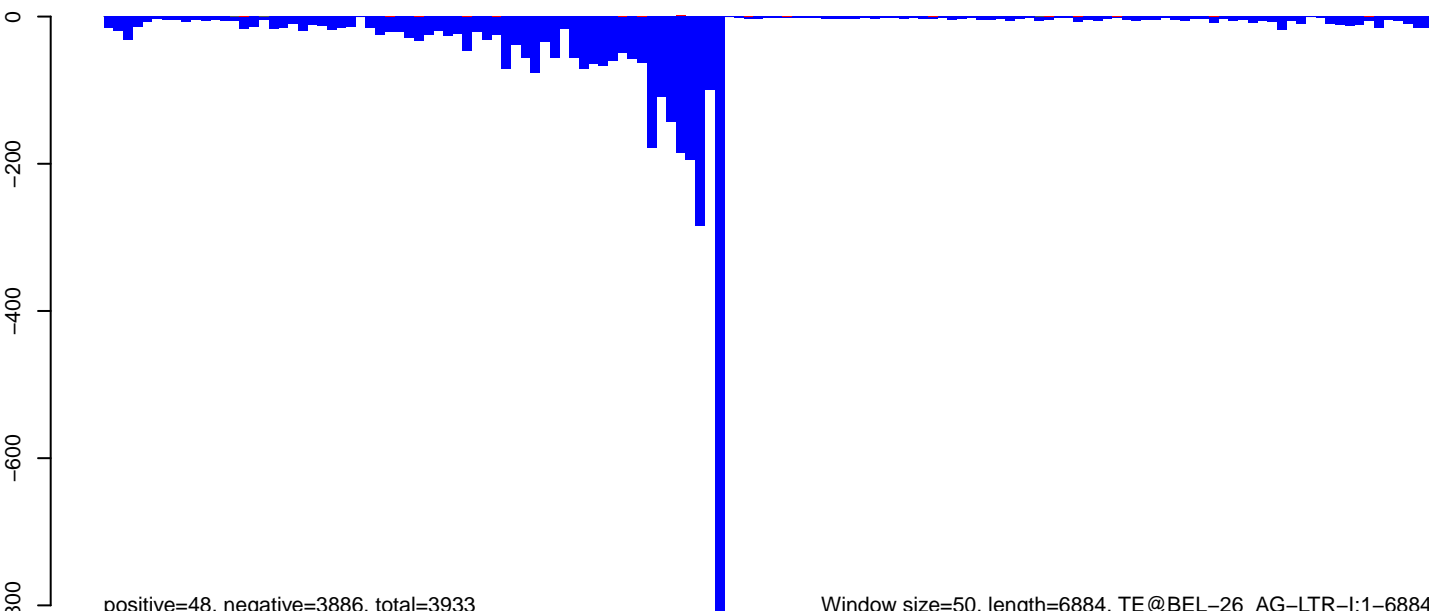
positive=26, negative=341, total=367

AnGam_Mos55_cells.24_35.rep



positive=22, negative=3544, total=3567

AnGam_Mos55_cells.rep



positive=48, negative=3886, total=3933

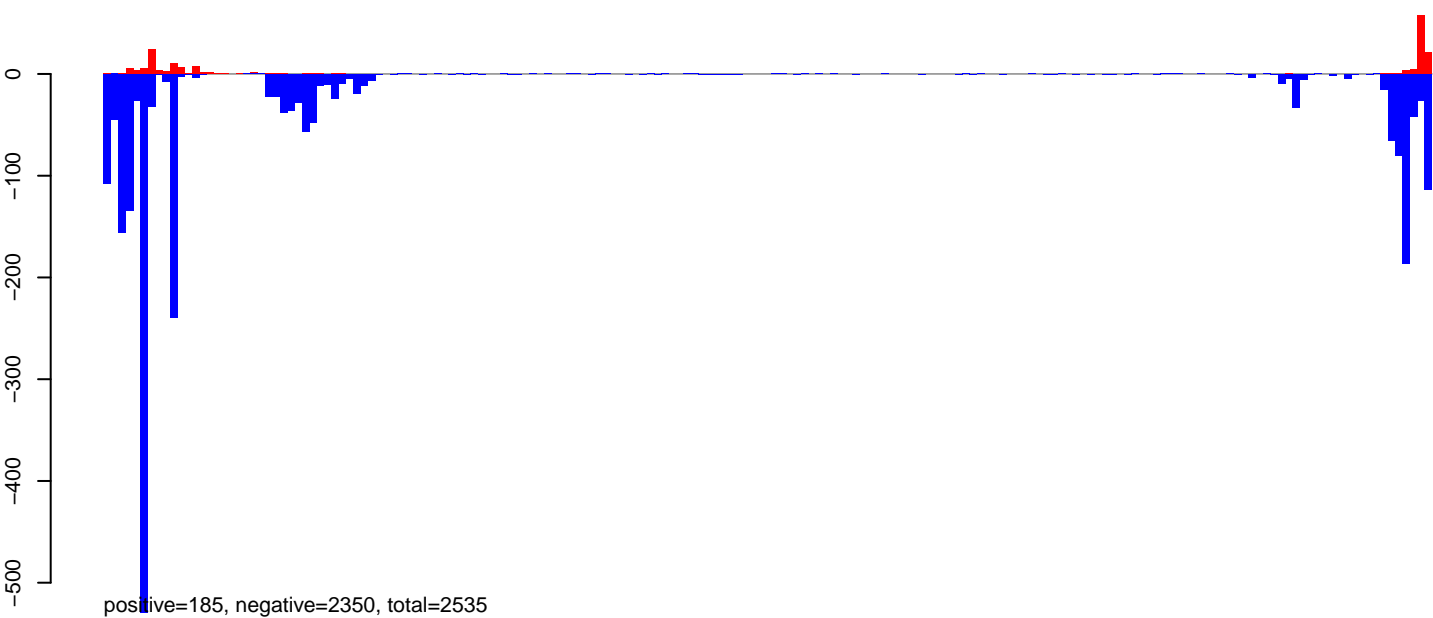
Window size=50, length=6884, TE@BEL-26_AG-LTR-I:1-6884

0 1000 2000 3000 4000 5000 6000 7000

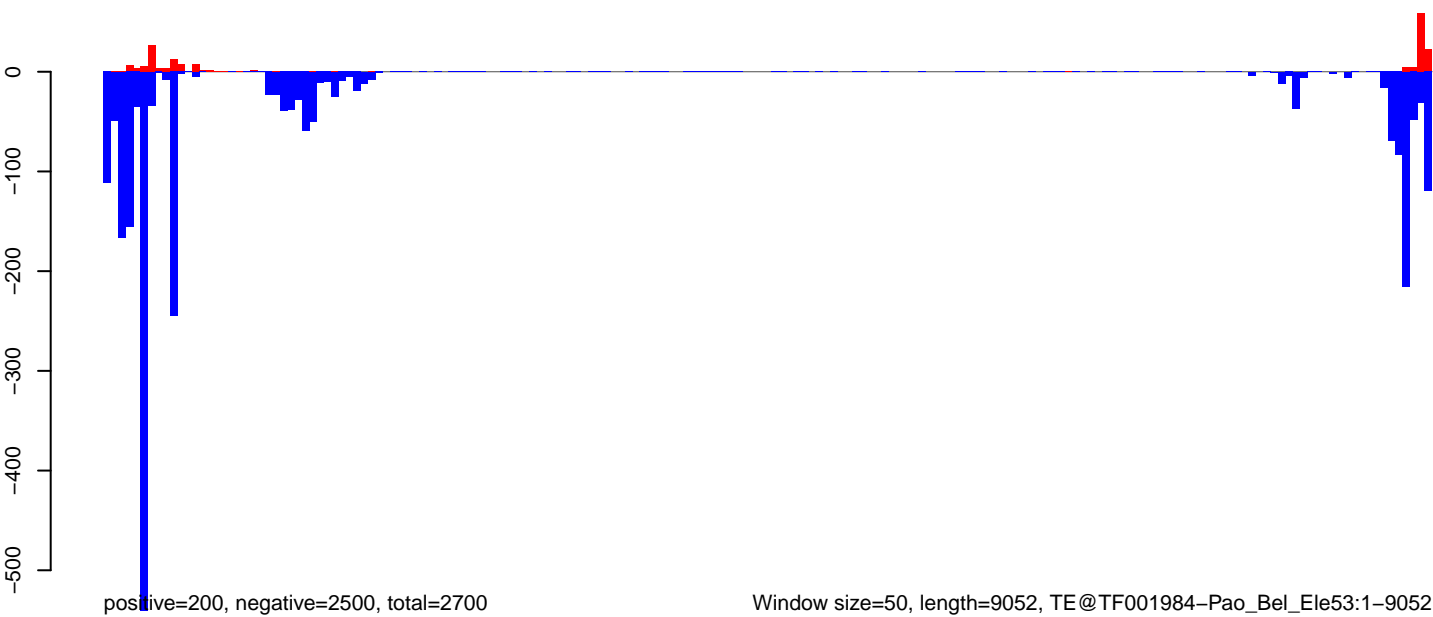
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



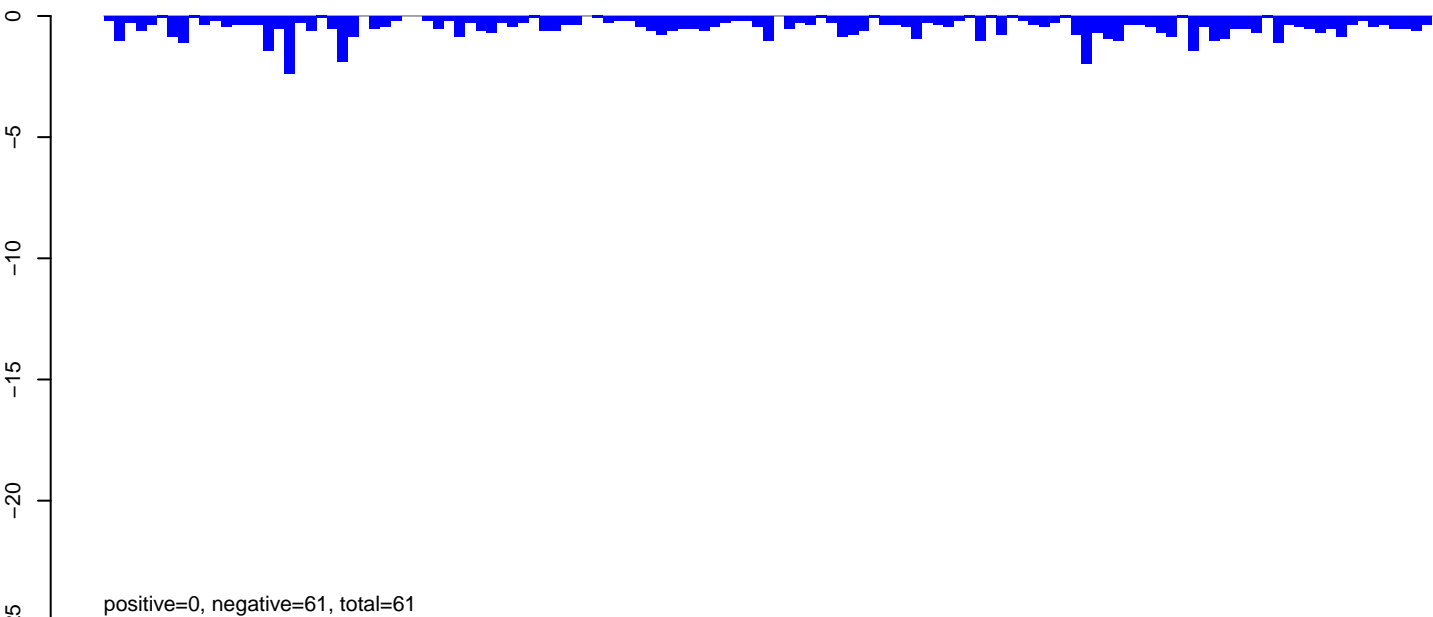
AnGam_Mos55_cells.rep



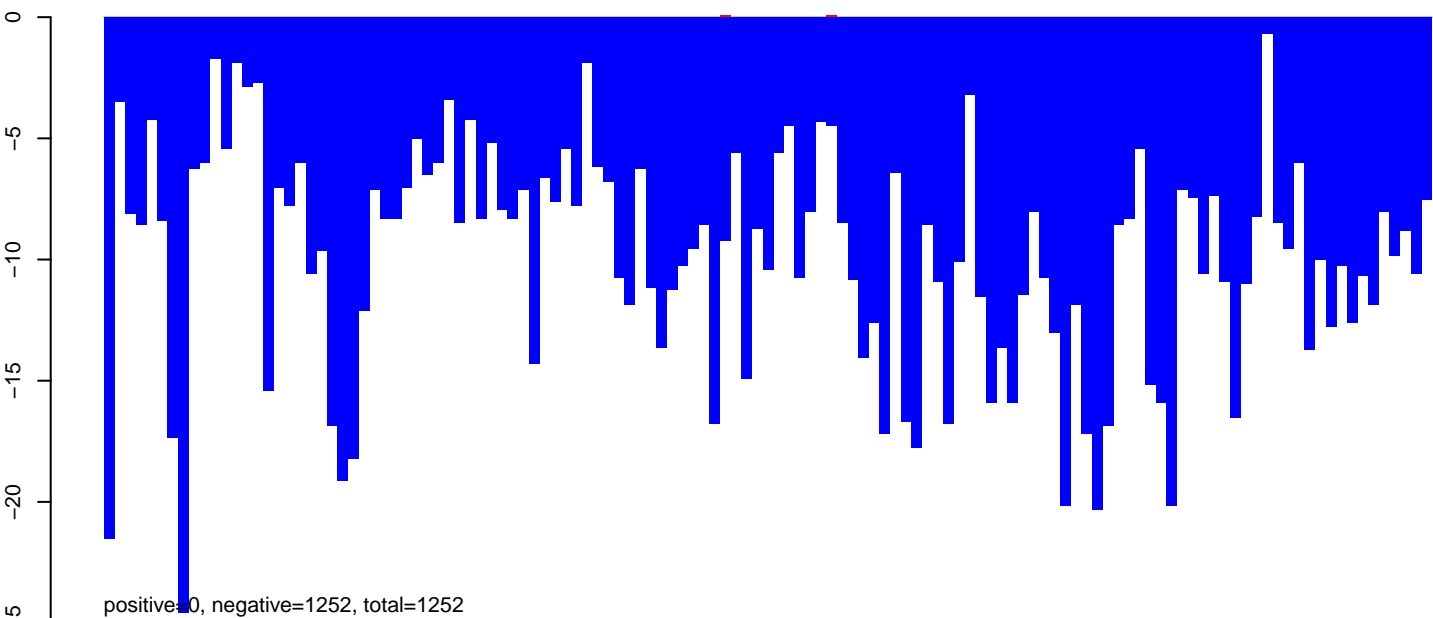
Window size=50, length=9052, TE@TF001984-Pao_Bel_Ele53:1-9052

0 2000 4000 6000 8000

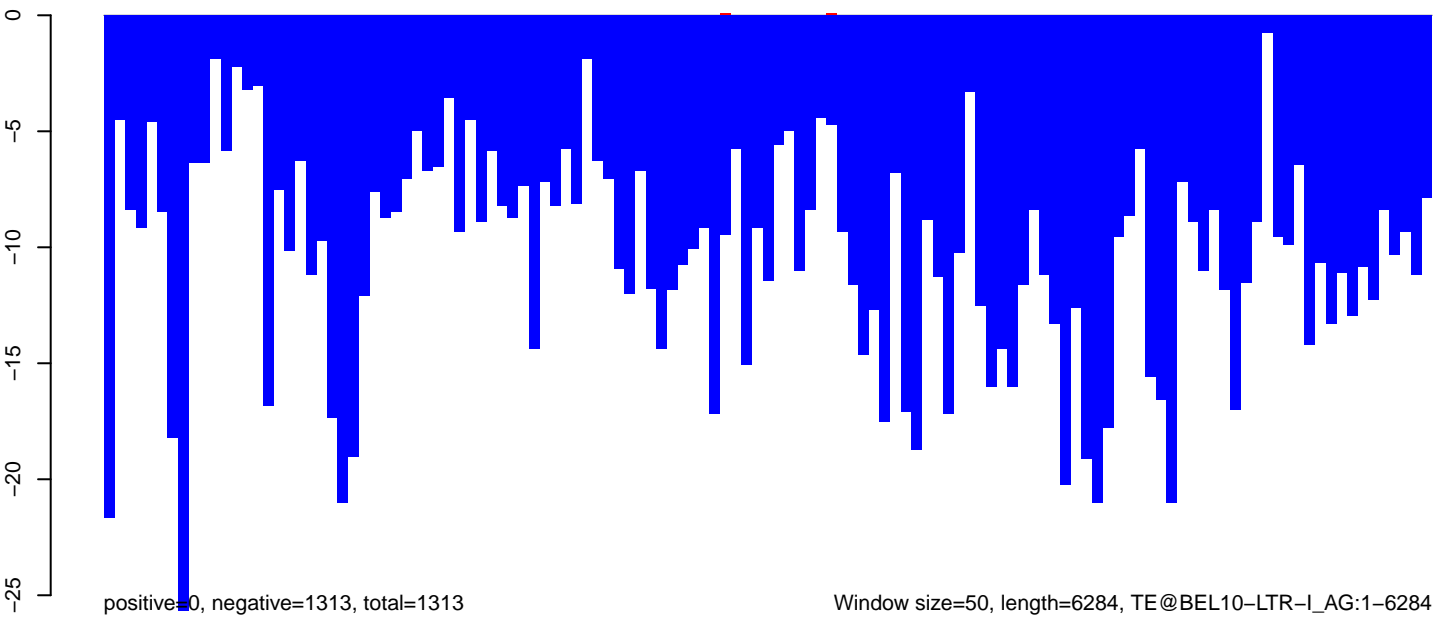
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

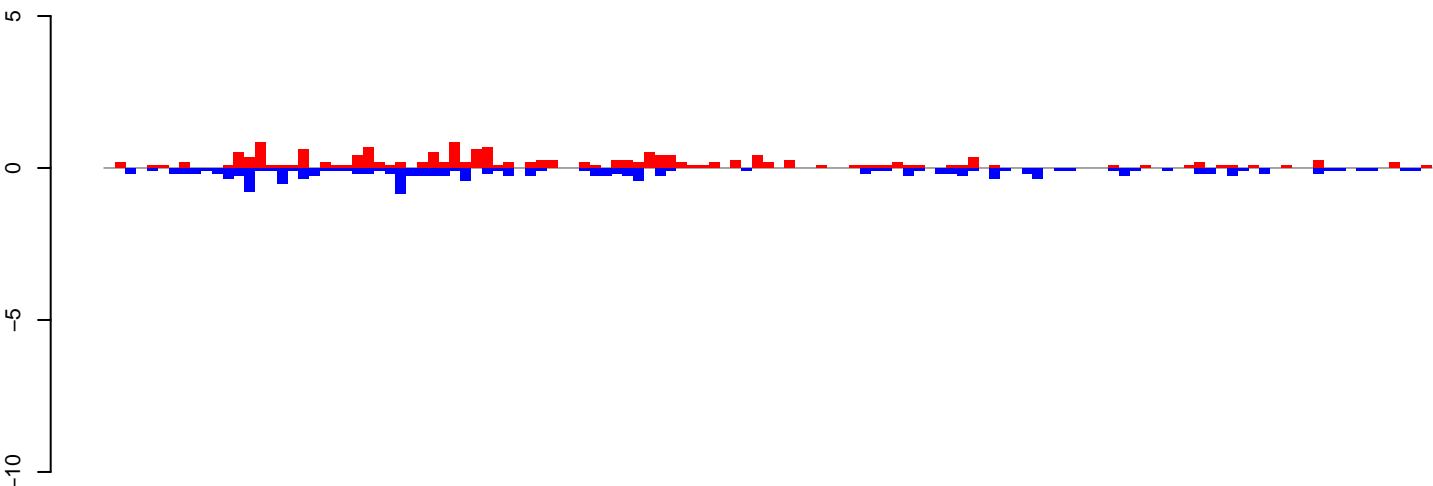


AnGam_Mos55_cells.rep



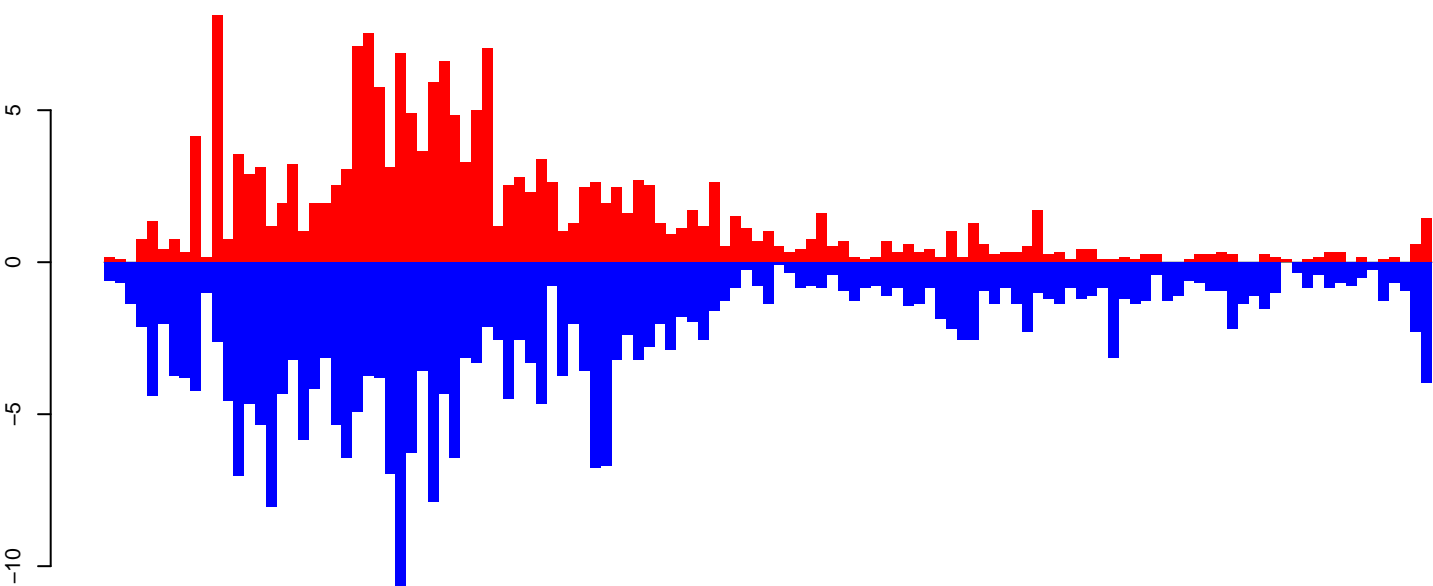
Window size=50, length=6284, TE@BEL10-LTR-I_AG:1-6284

AnGam_Mos55_cells.18_23.rep



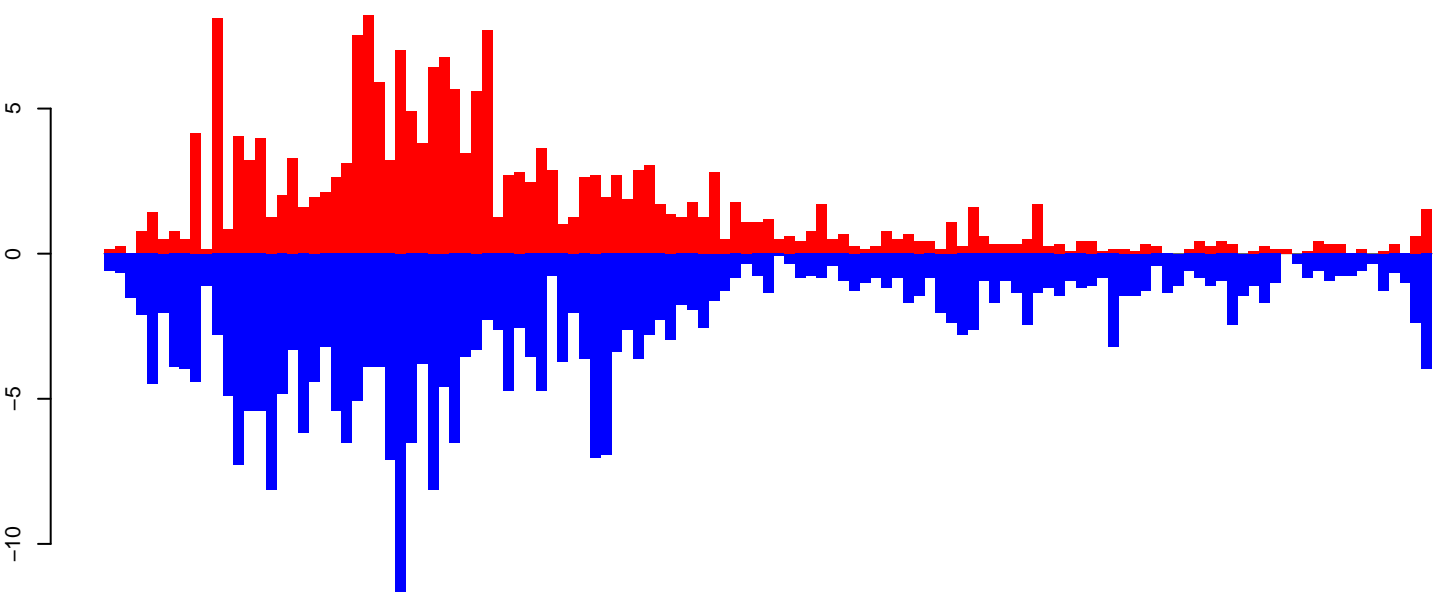
positive=16, negative=14, total=30

AnGam_Mos55_cells.24_35.rep



positive=185, negative=300, total=486

AnGam_Mos55_cells.rep

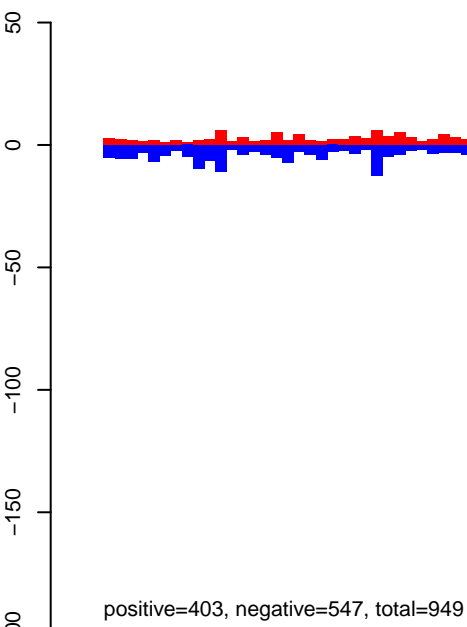


positive=201, negative=315, total=516

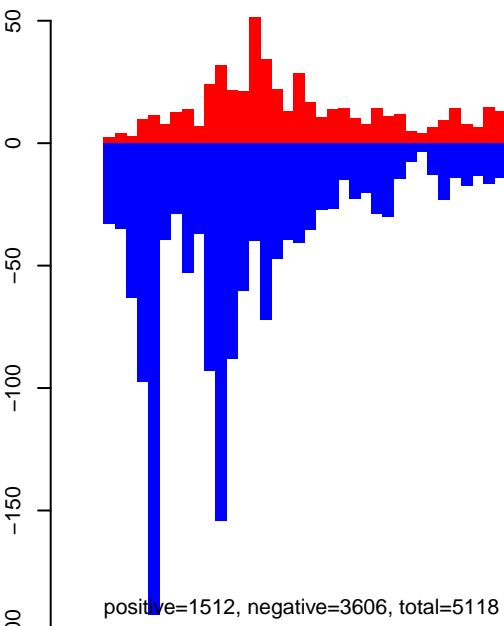
Window size=50, length=6196, TE@TF001952-Pao_Bel_Ele21:1-6196

0 1000 2000 3000 4000 5000 6000

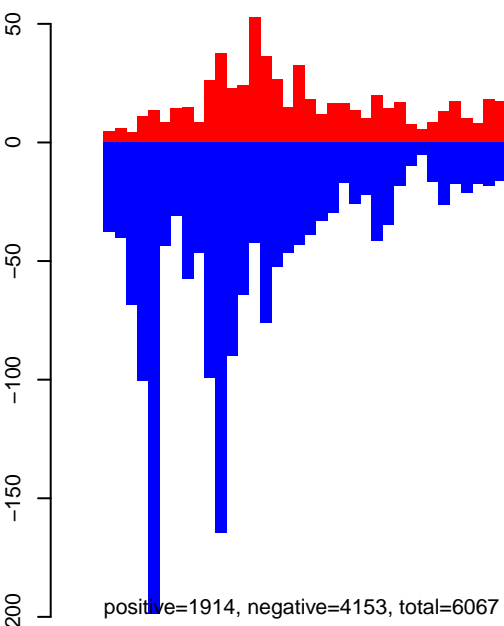
AnGam_Mos55_cells.18_23.rep



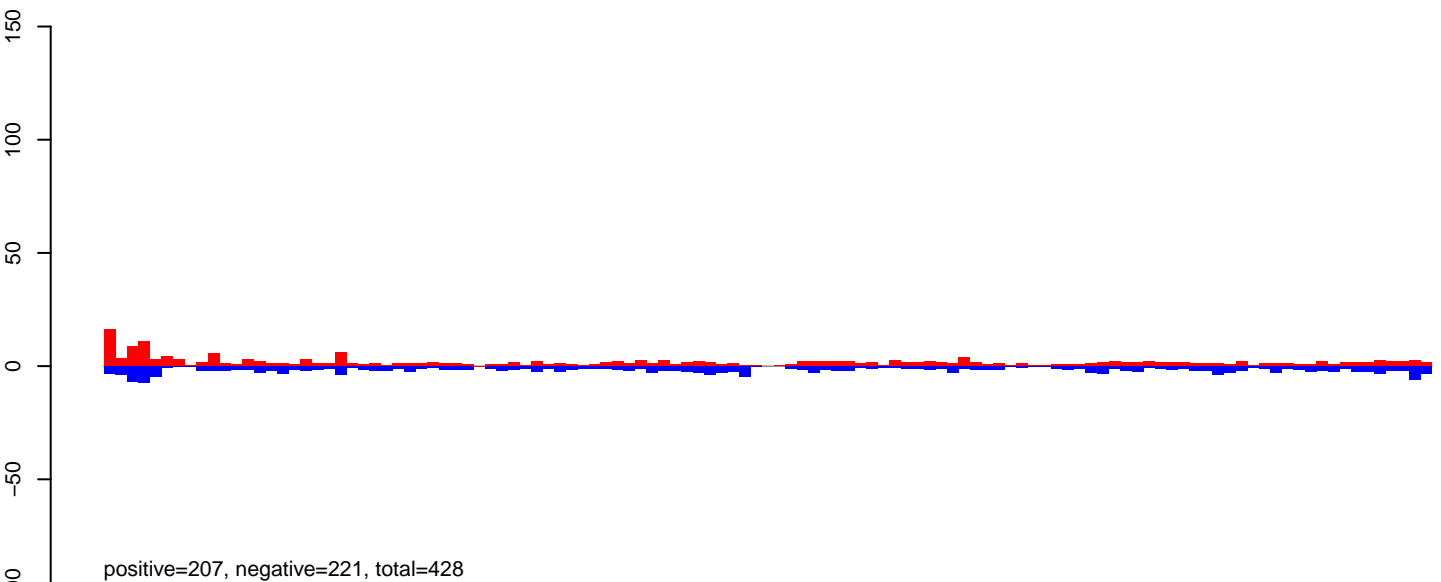
AnGam_Mos55_cells.24_35.rep



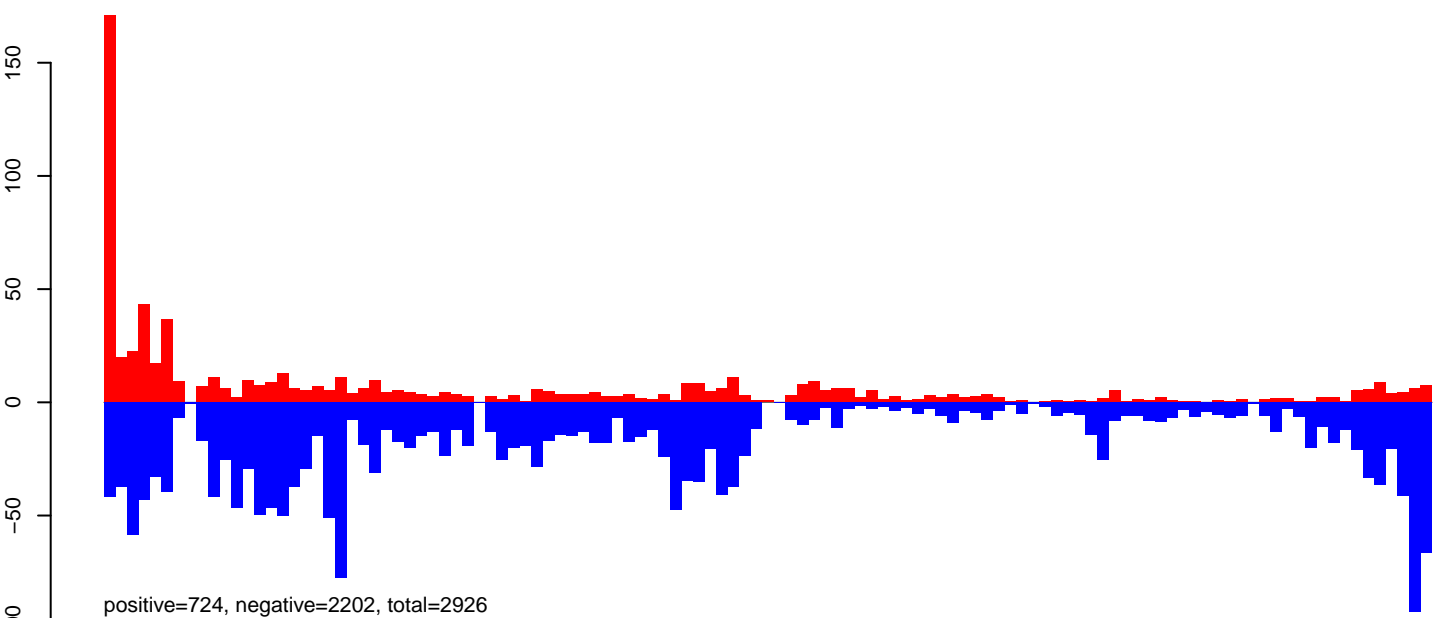
AnGam_Mos55_cells.rep



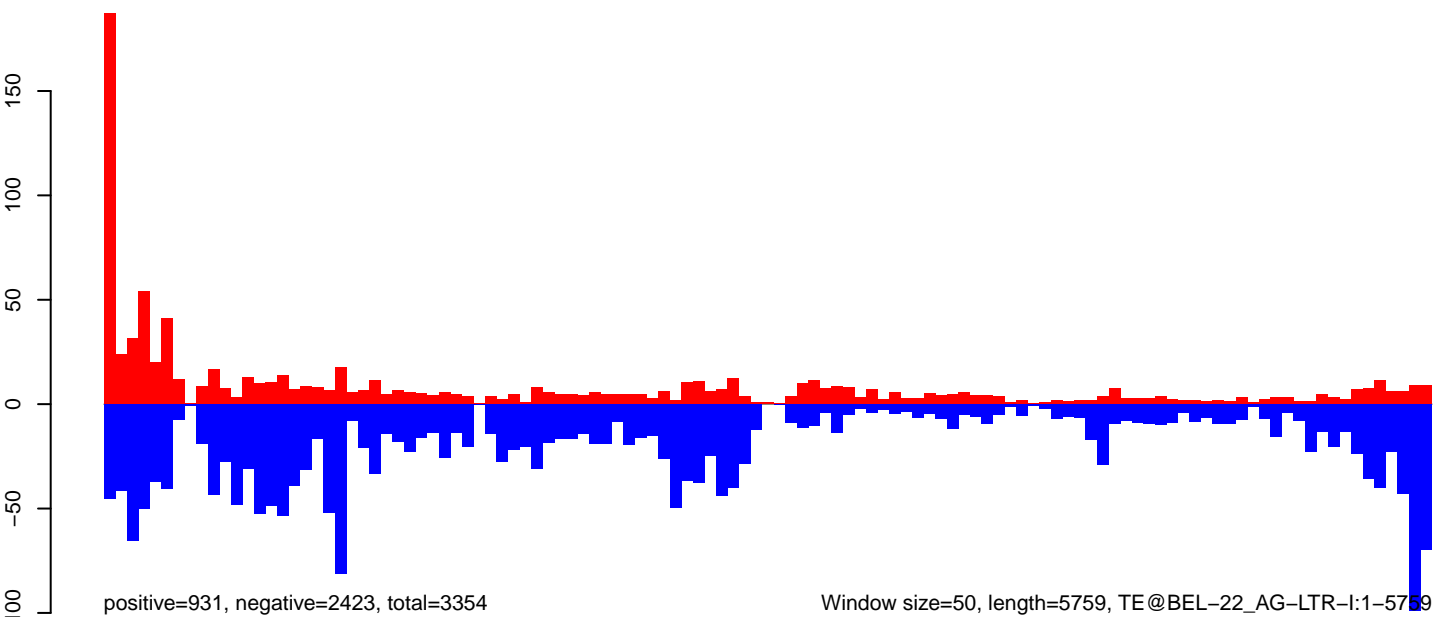
AnGam_Mos55_cells.18_23.rep



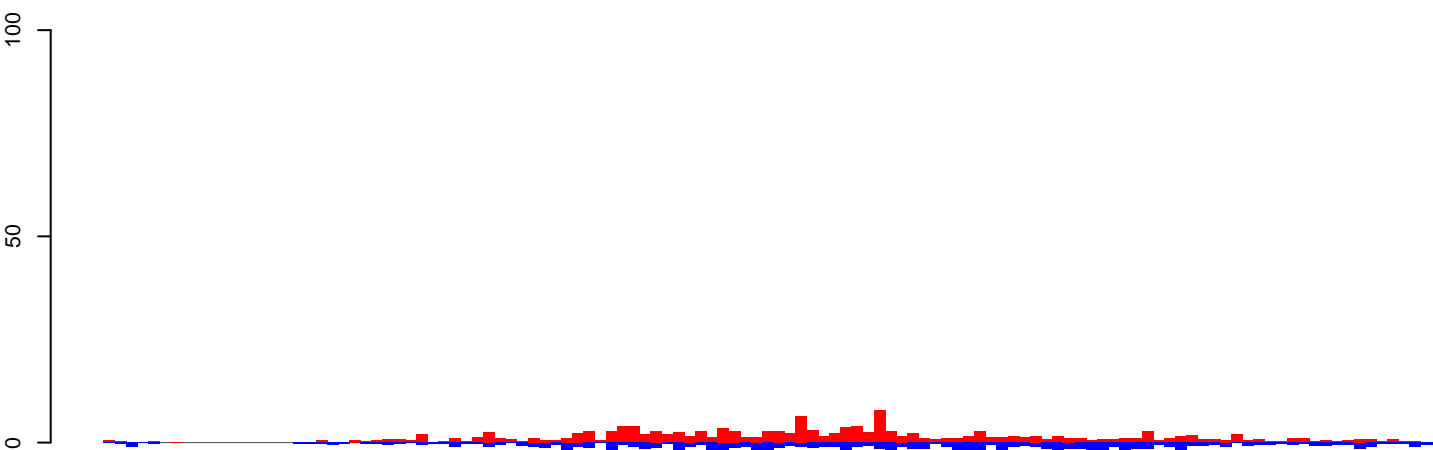
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

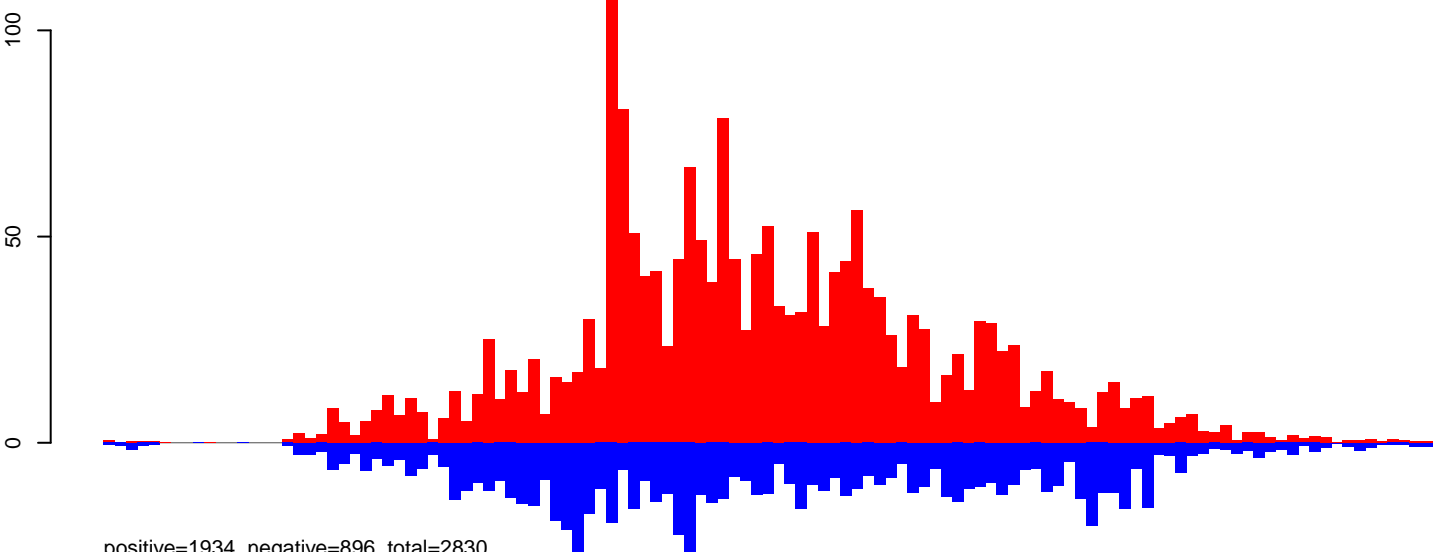


AnGam_Mos55_cells.18_23.rep



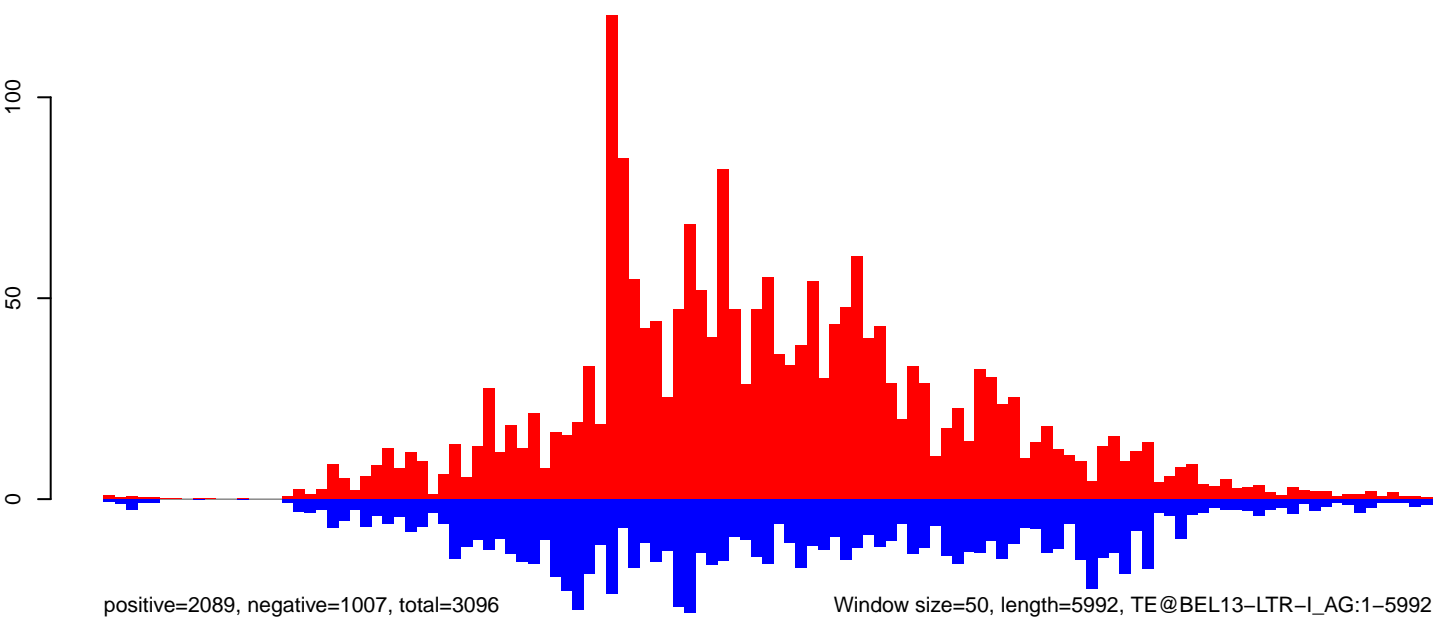
positive=155, negative=111, total=266

AnGam_Mos55_cells.24_35.rep



positive=1934, negative=896, total=2830

AnGam_Mos55_cells.rep

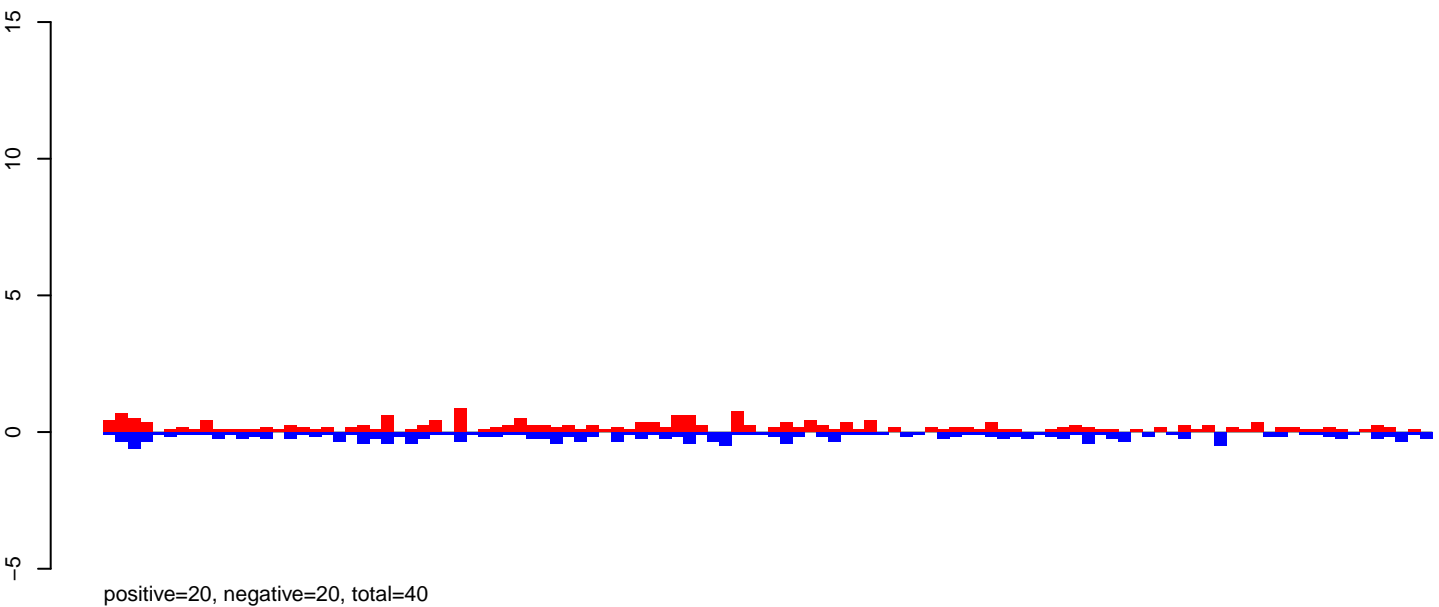


positive=2089, negative=1007, total=3096

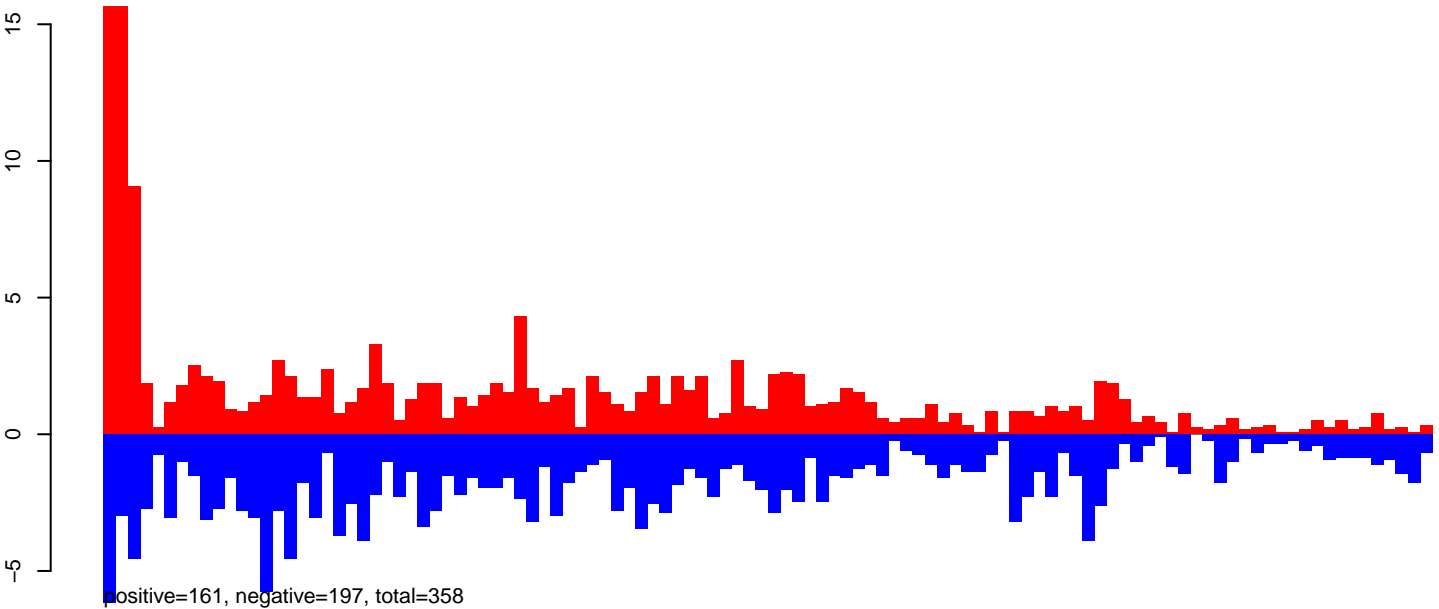
Window size=50, length=5992, TE@BEL13-LTR-I_AG:1-5992

0 1000 2000 3000 4000 5000 6000

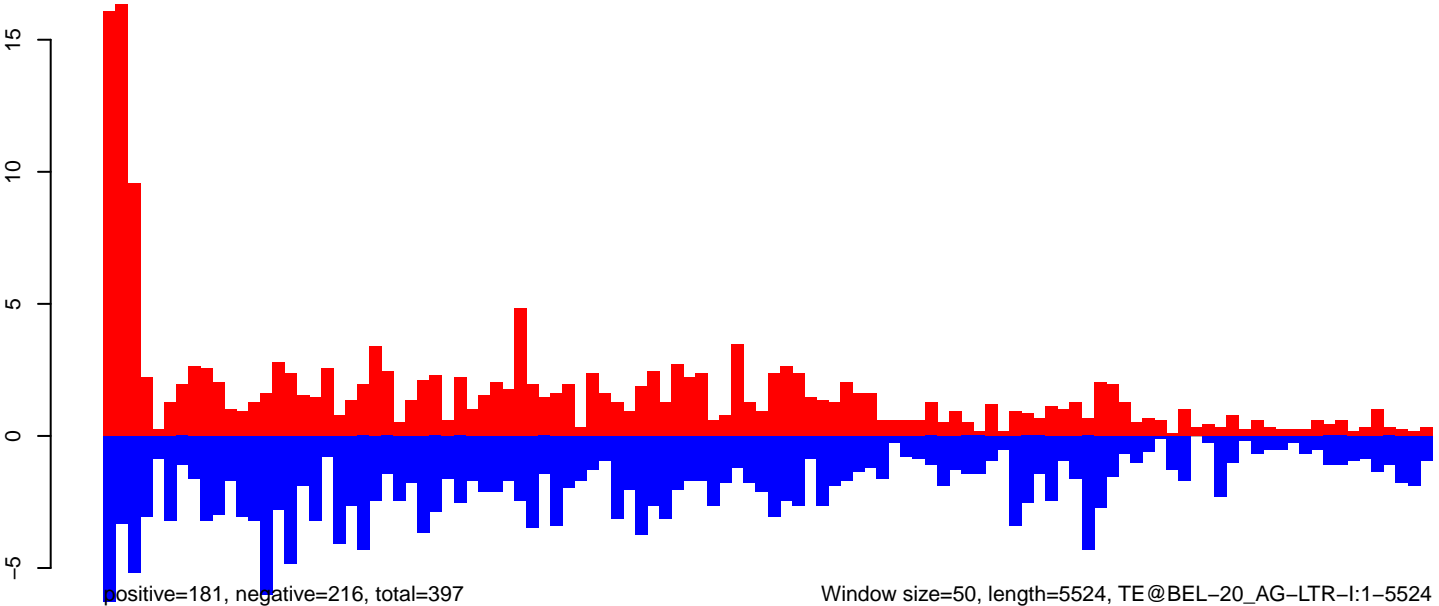
AnGam_Mos55_cells.18_23.rep



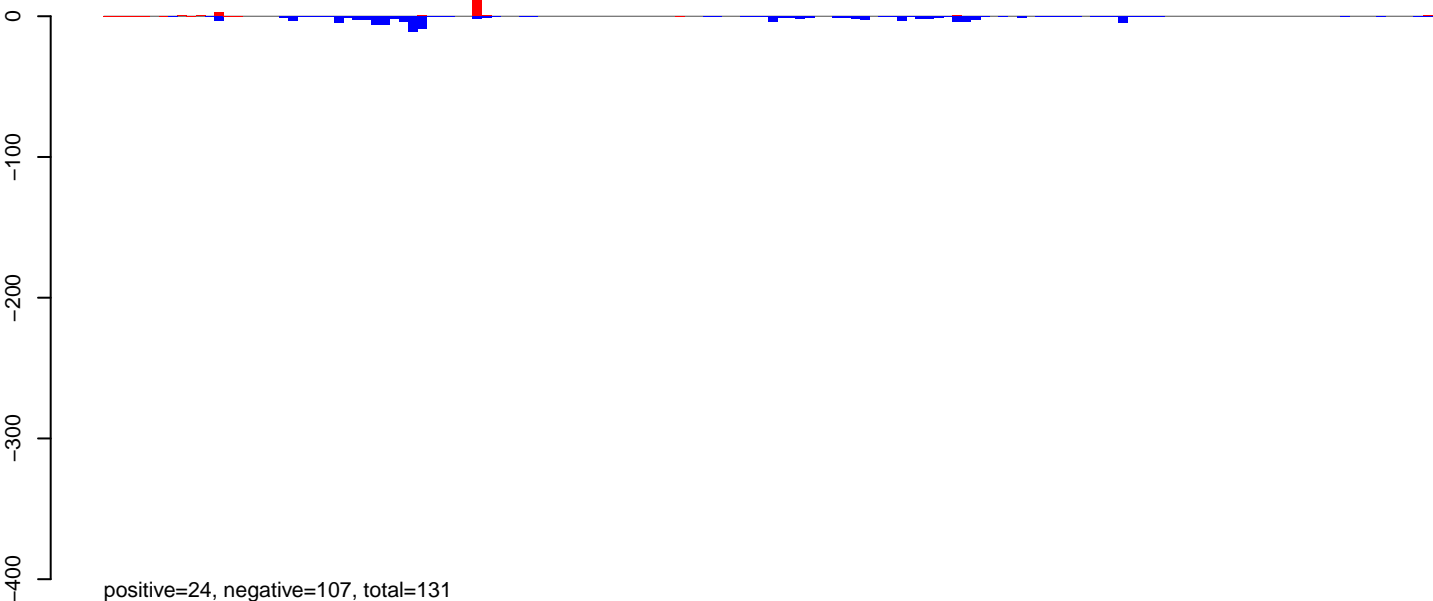
AnGam_Mos55_cells.24_35.rep



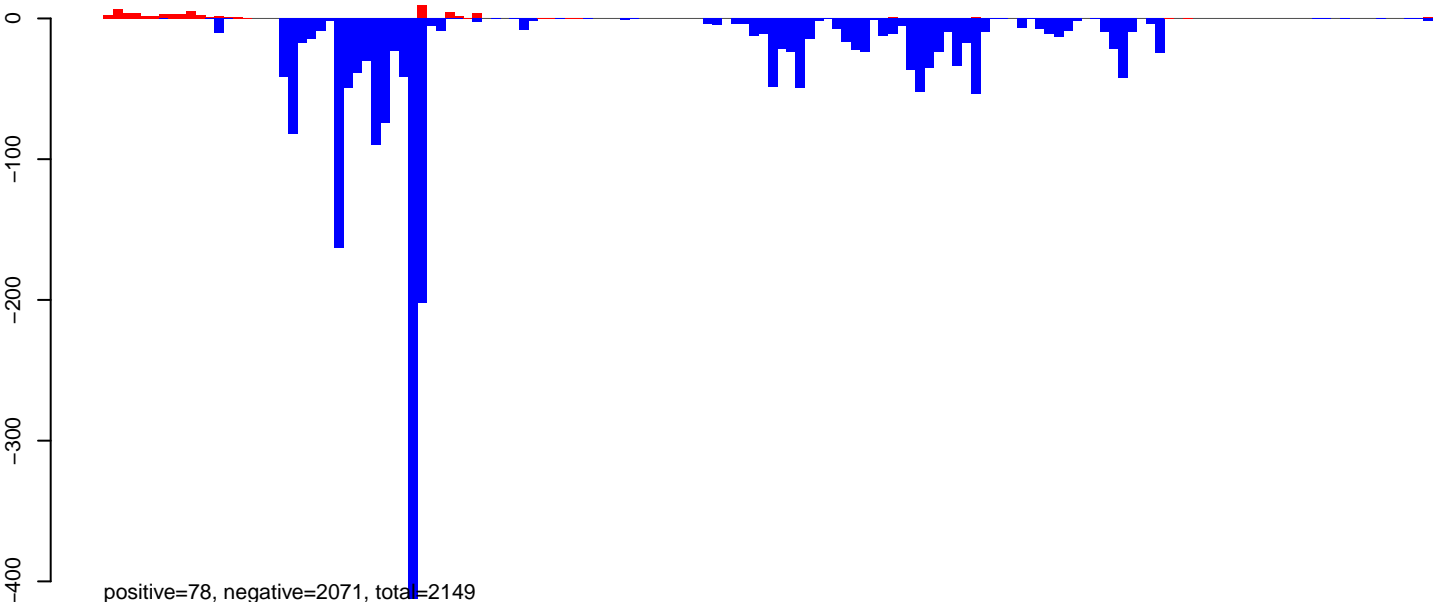
AnGam_Mos55_cells.rep



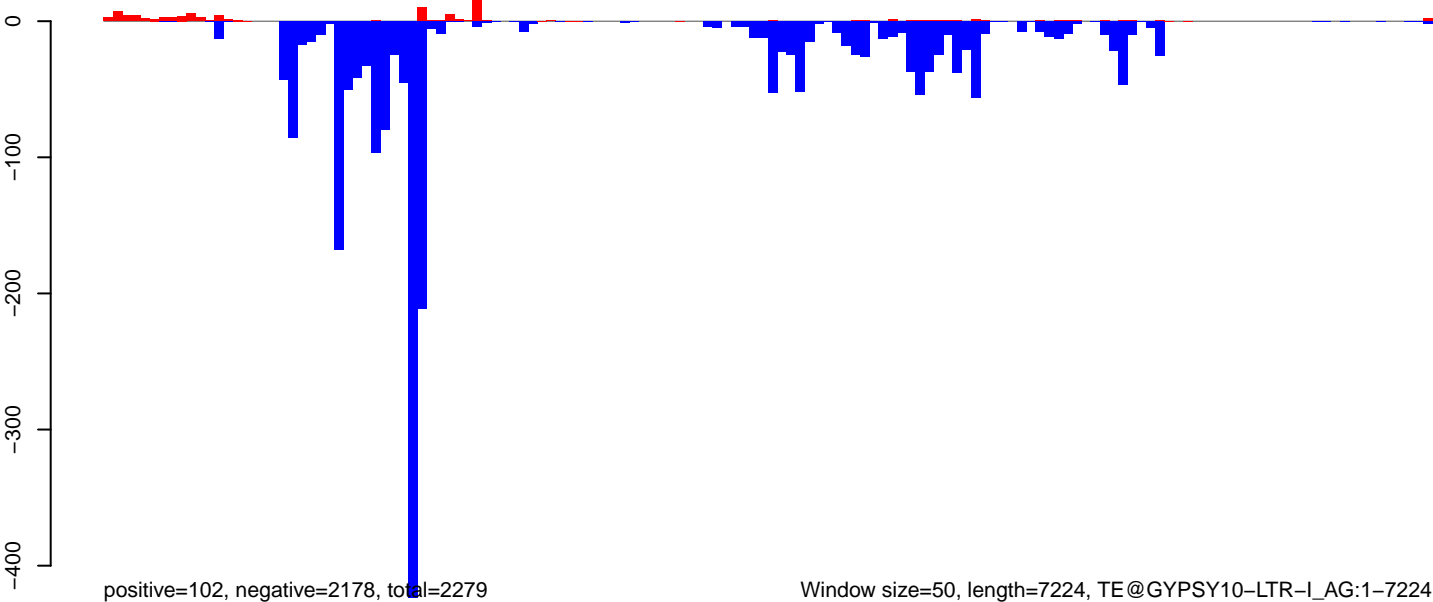
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



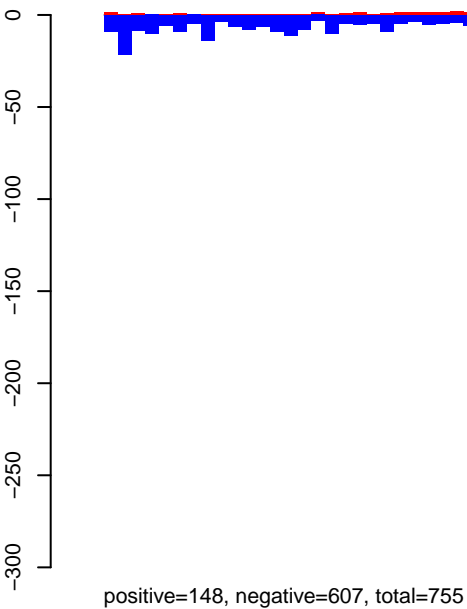
AnGam_Mos55_cells.rep



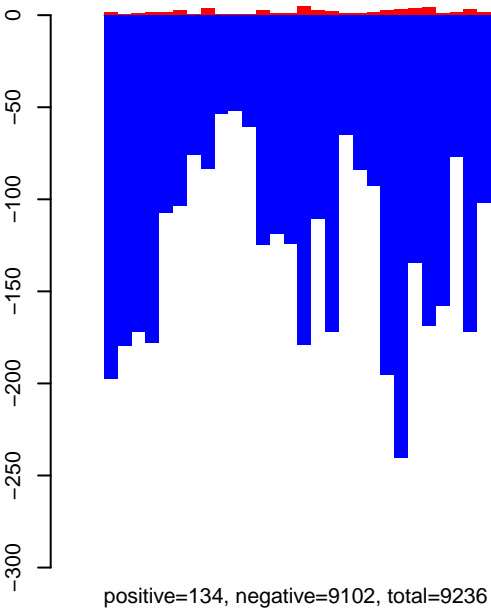
Window size=50, length=7224, TE@GYPSY10-LTR-I_AG:1-7224

0 1000 2000 3000 4000 5000 6000 7000

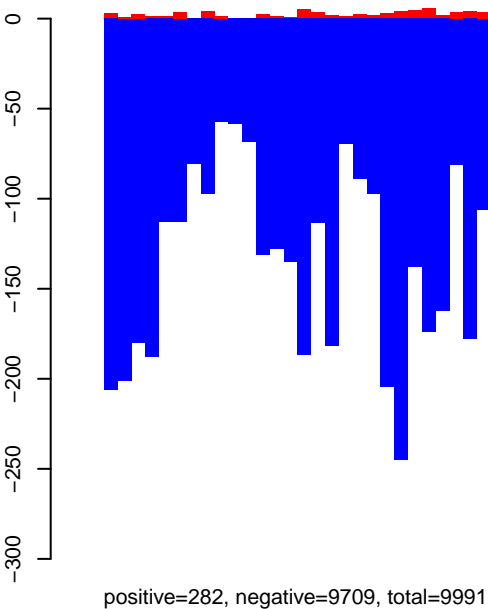
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



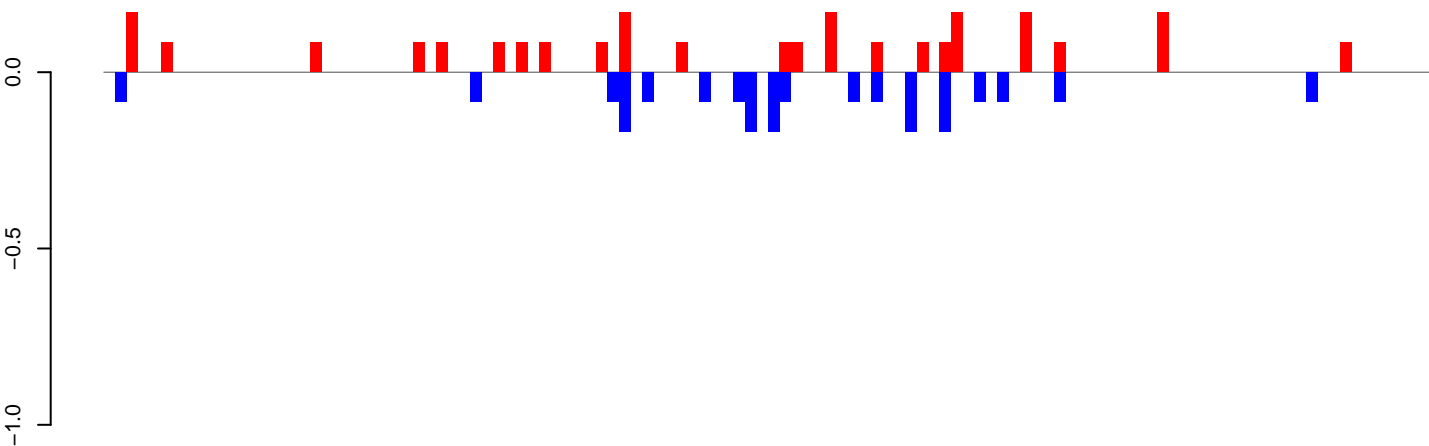
AnGam_Mos55_cells.rep



Window size=50, length=4848, TE@GYPSY60-LTR-I_AG:1-4848

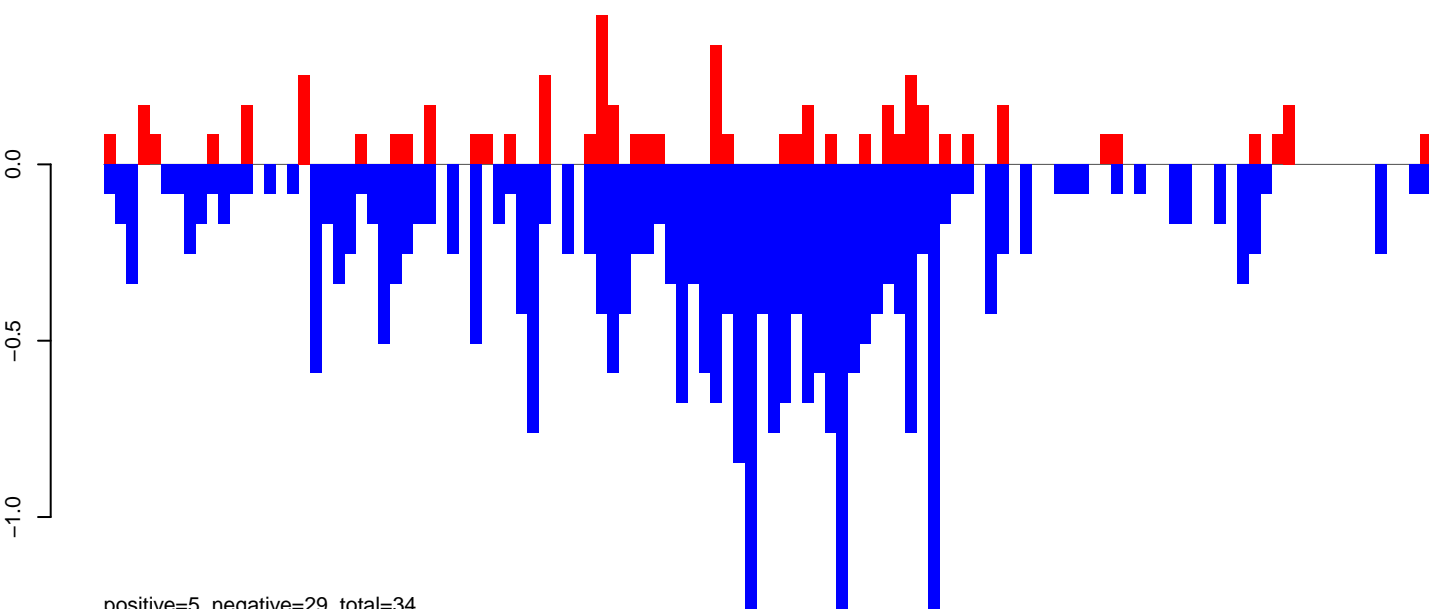
0 1000 2000 3000 4000 5000

AnGam_Mos55_cells.18_23.rep



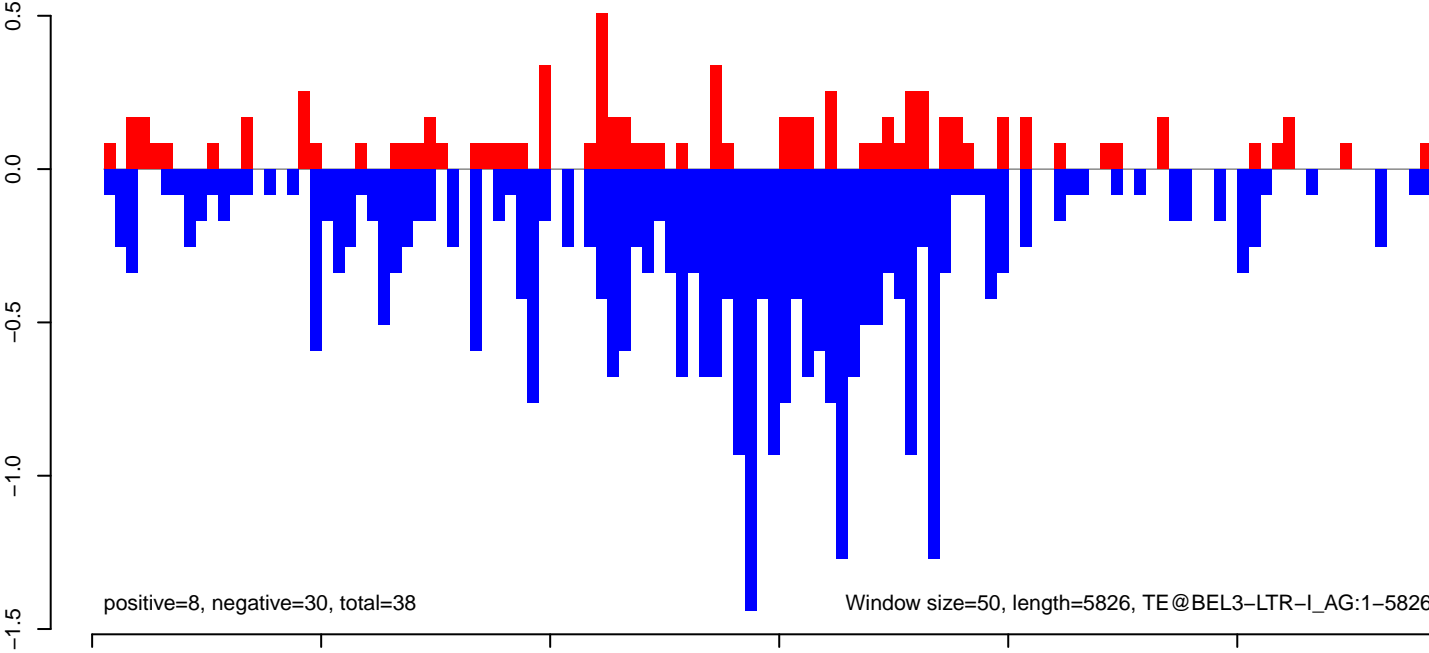
positive=2, negative=2, total=4

AnGam_Mos55_cells.24_35.rep



positive=5, negative=29, total=34

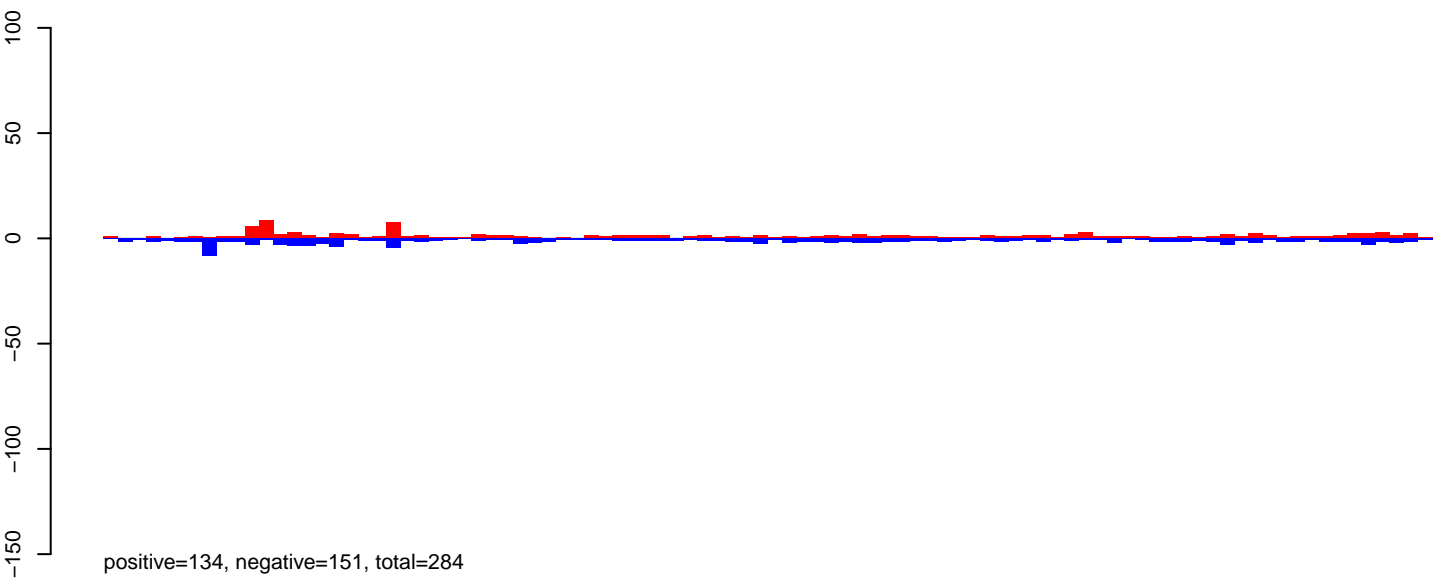
AnGam_Mos55_cells.rep



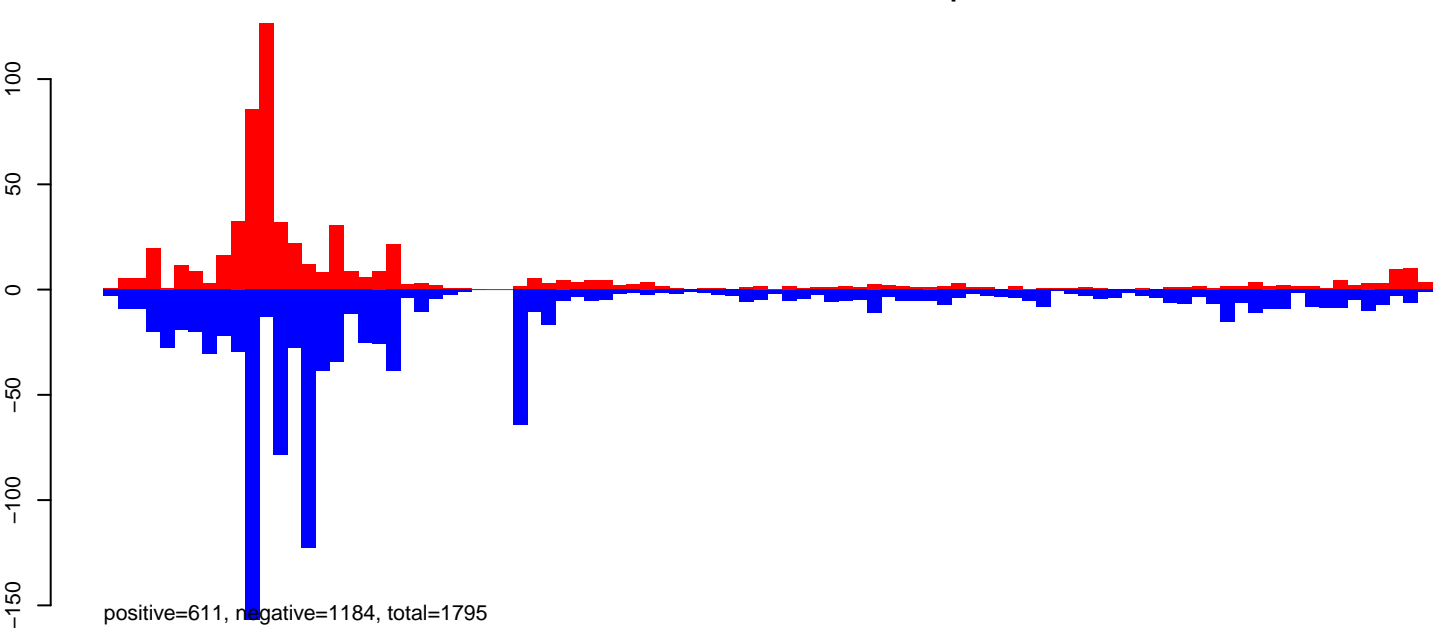
positive=8, negative=30, total=38

Window size=50, length=5826, TE@BEL3-LTR-I_AG:1-5826

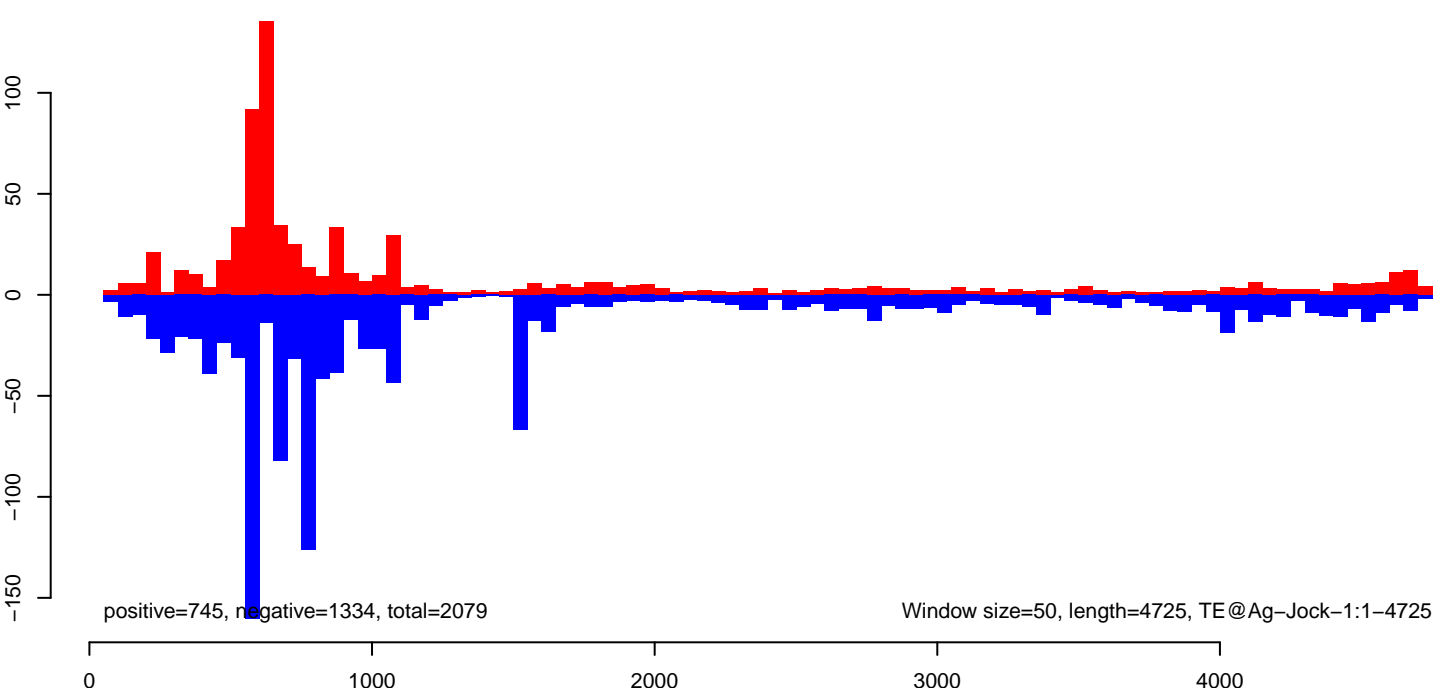
AnGam_Mos55_cells.18_23.rep



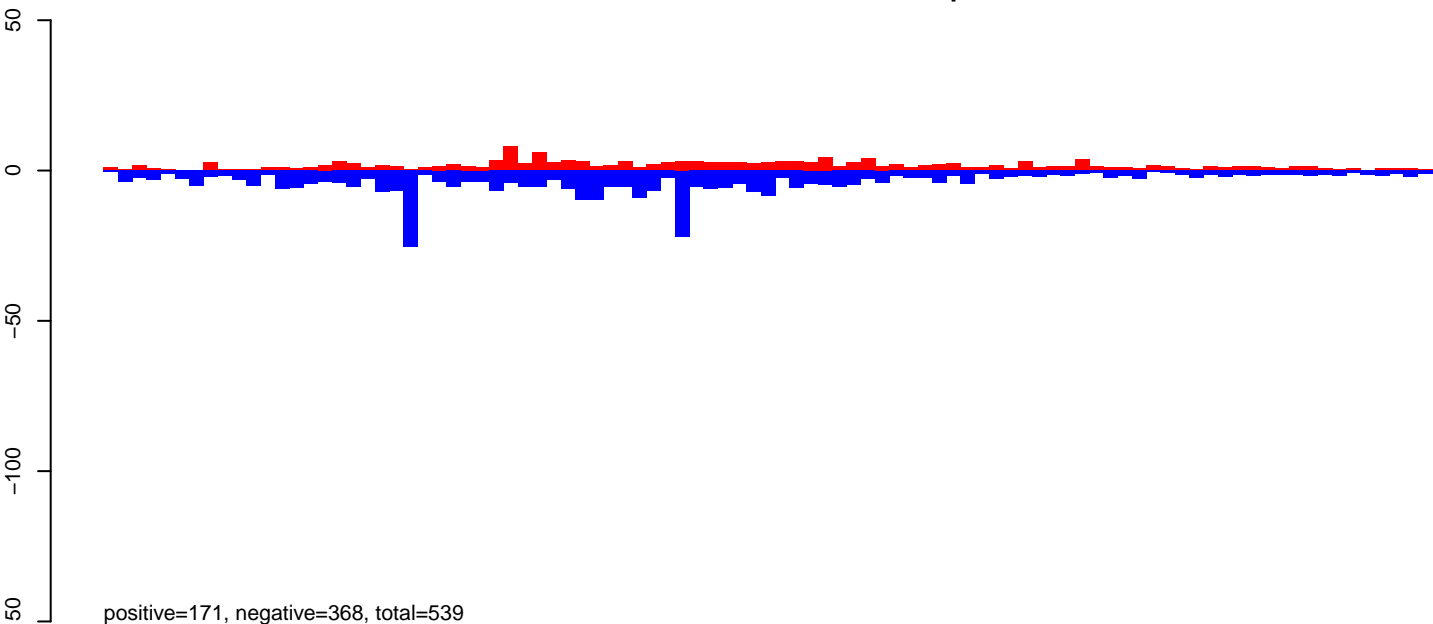
AnGam_Mos55_cells.24_35.rep



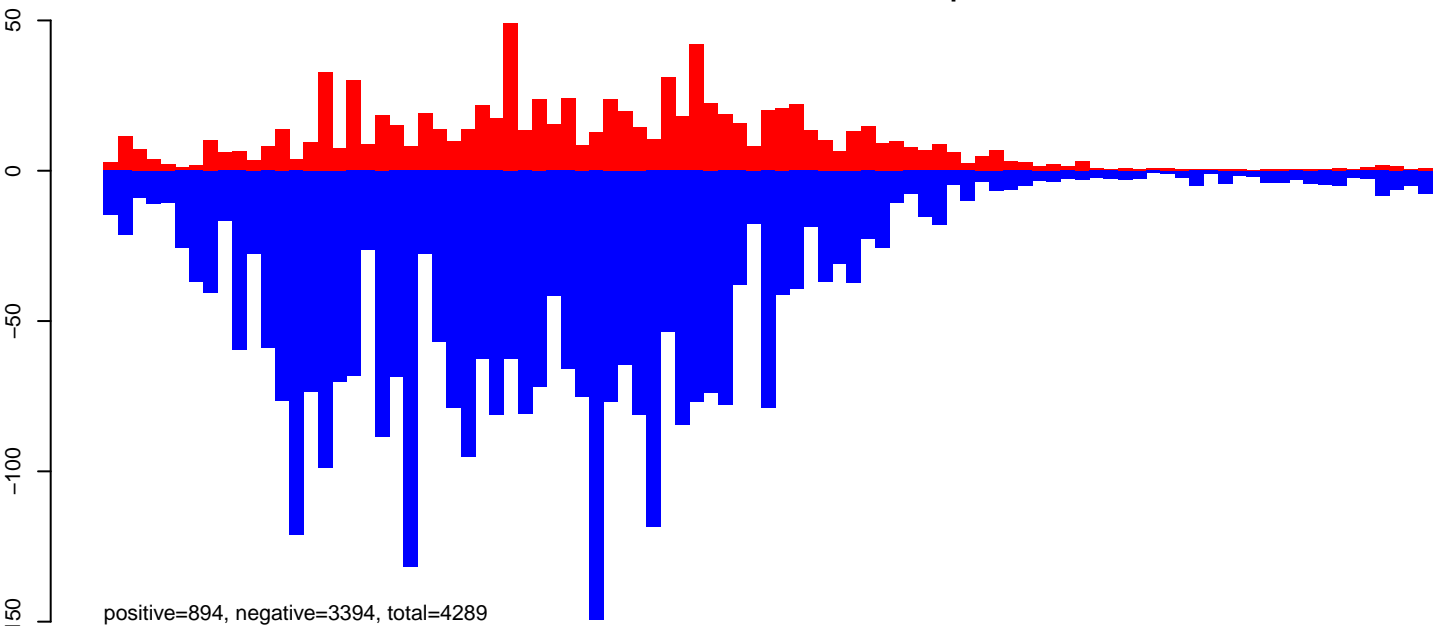
AnGam_Mos55_cells.rep



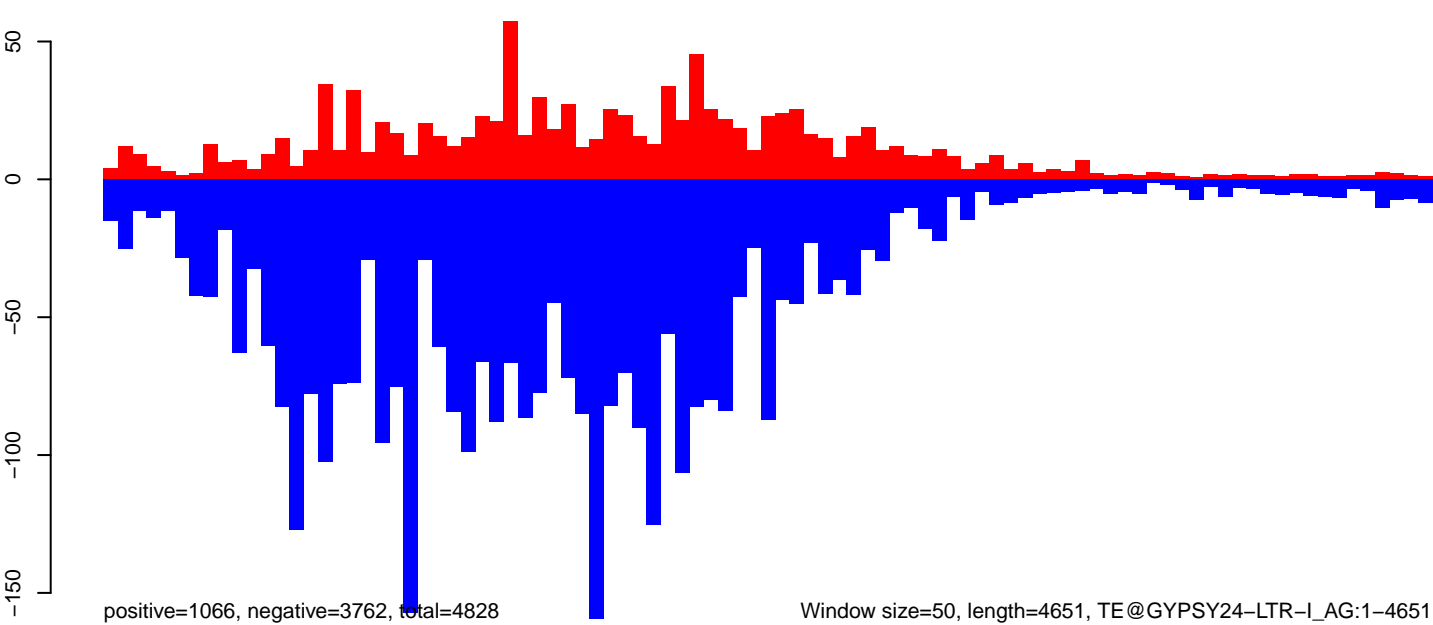
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



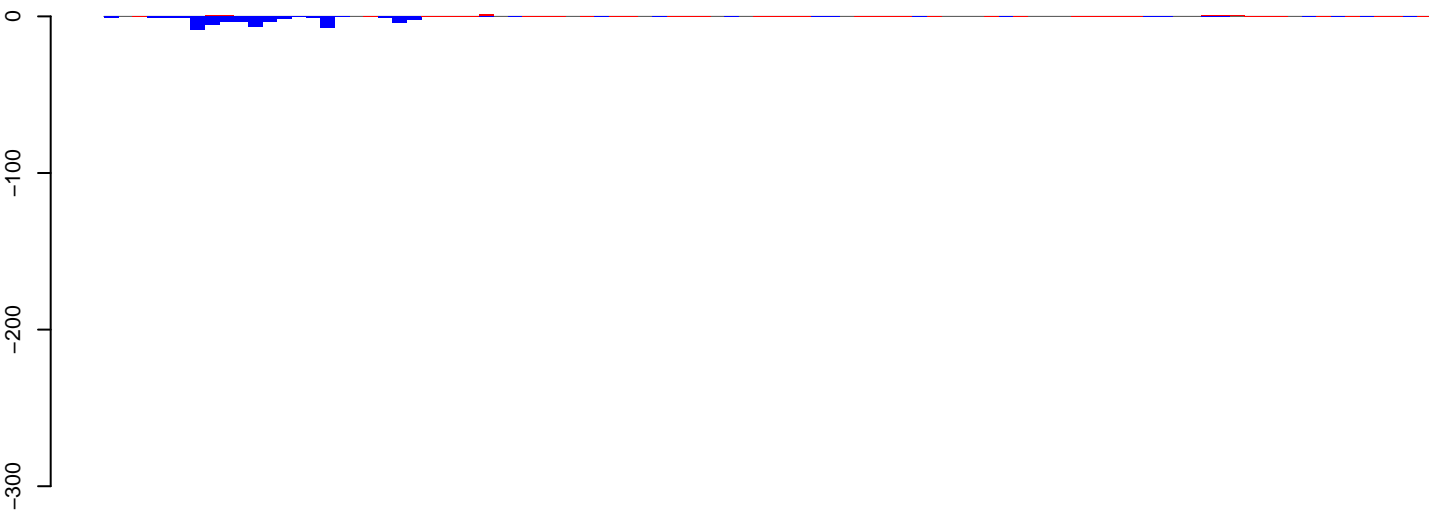
AnGam_Mos55_cells.rep



Window size=50, length=4651, TE@GYPSY24-LTR-I_AG:1-4651

0 1000 2000 3000 4000

AnGam_Mos55_cells.18_23.rep



positive=14, negative=54, total=68

AnGam_Mos55_cells.24_35.rep



positive=236, negative=779, total=1015

AnGam_Mos55_cells.rep

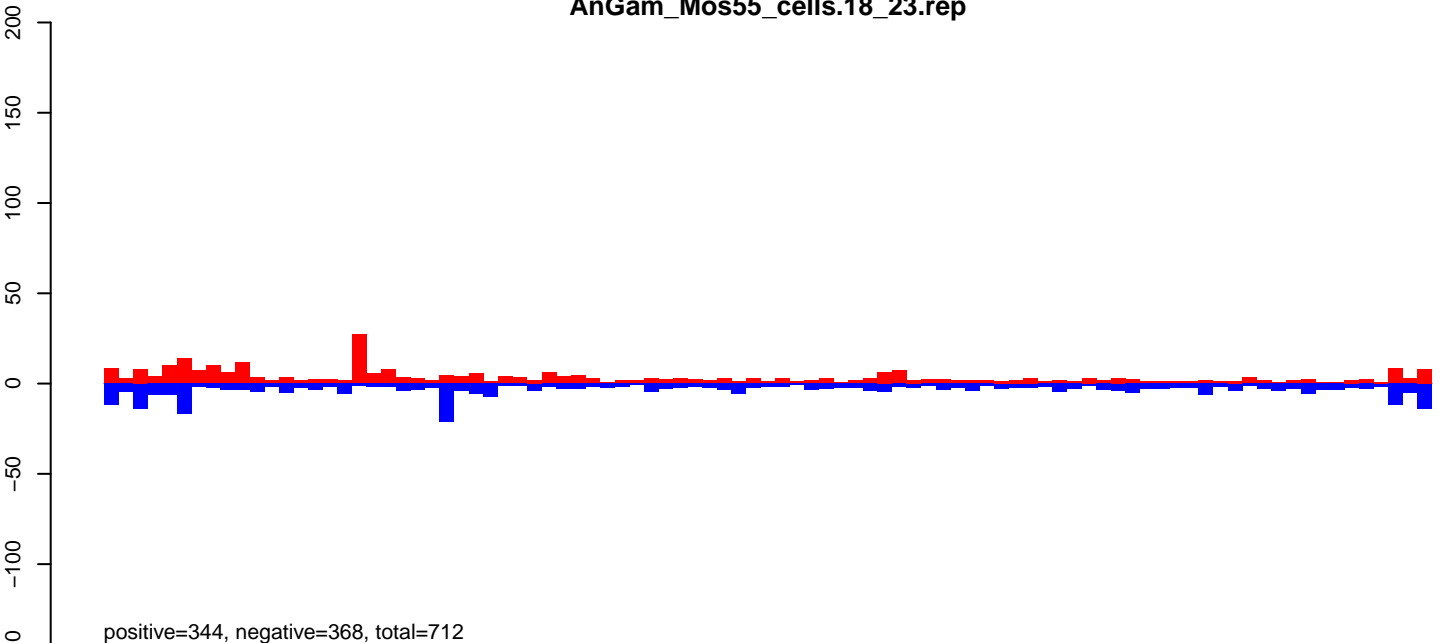


positive=250, negative=833, total=1083

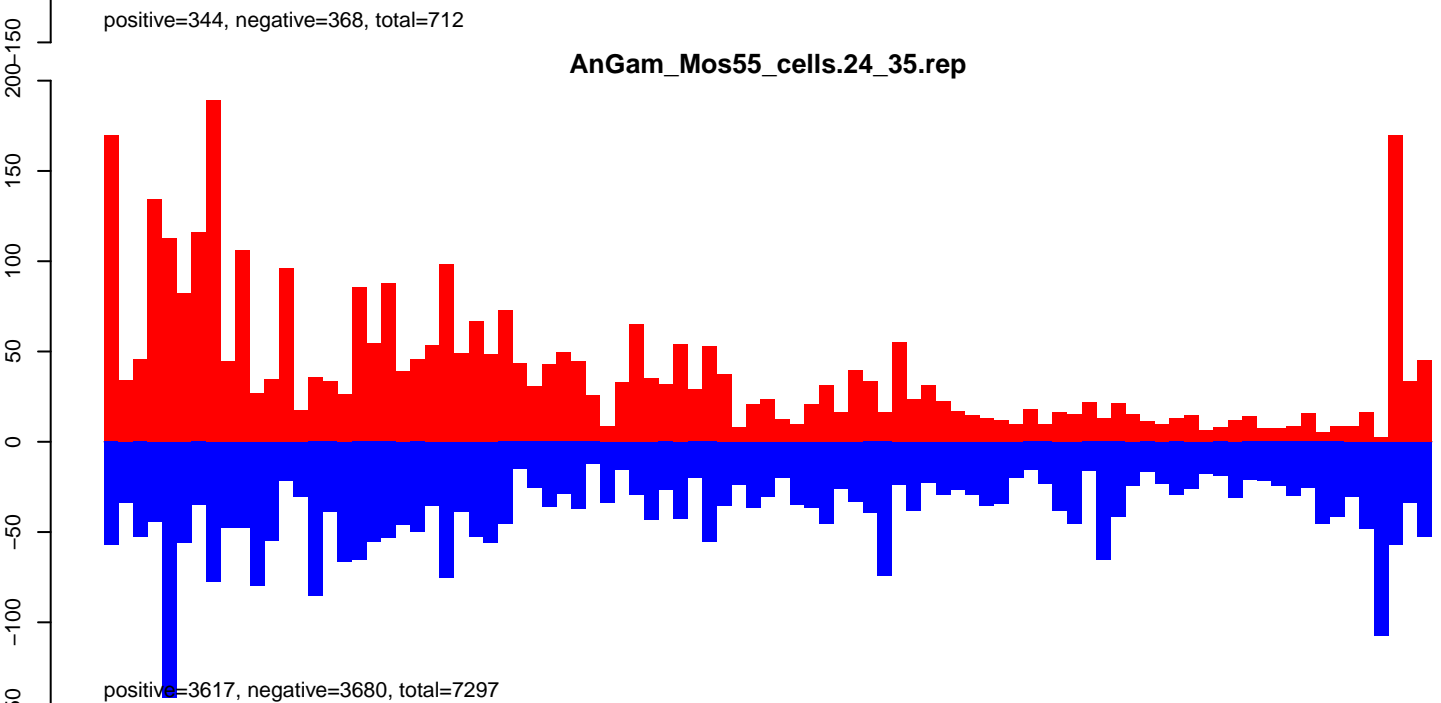
Window size=50, length=4634, TE@T1:1-4634

0 1000 2000 3000 4000

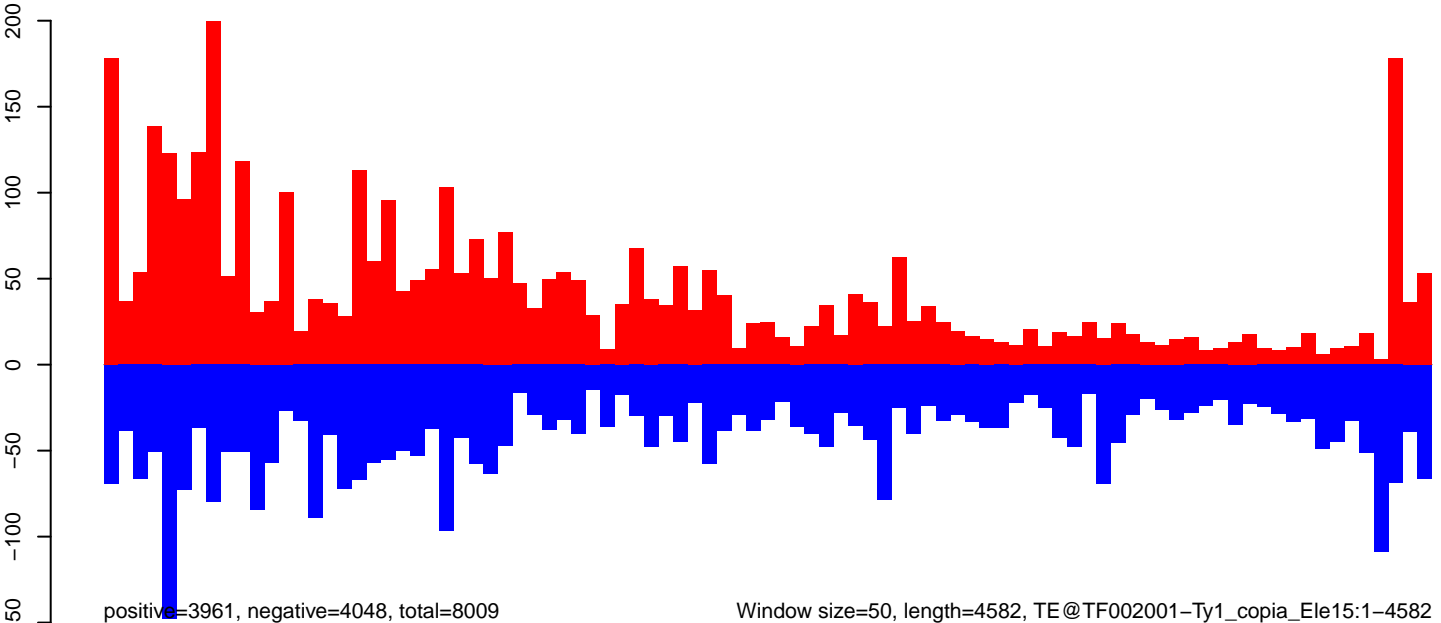
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



Window size=50, length=4582, TE@TF002001-Ty1_copia_Ele15:1-4582

AnGam_Mos55_cells.18_23.rep



positive=1, negative=18, total=19

AnGam_Mos55_cells.24_35.rep



positive=1, negative=424, total=425

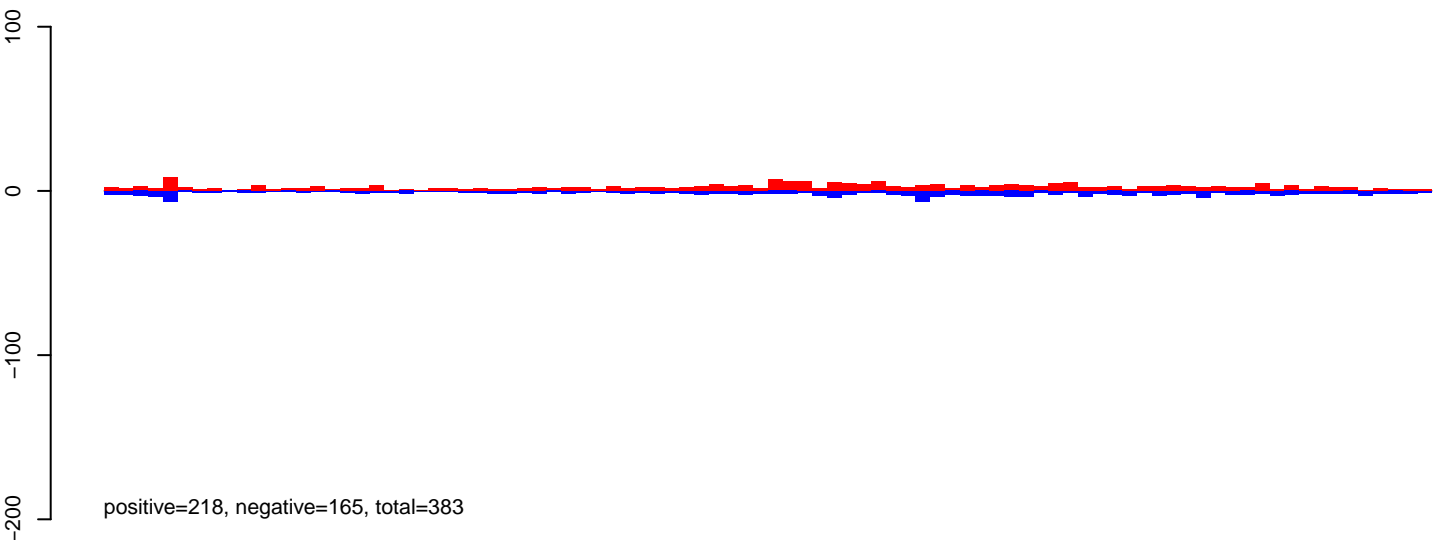
AnGam_Mos55_cells.rep



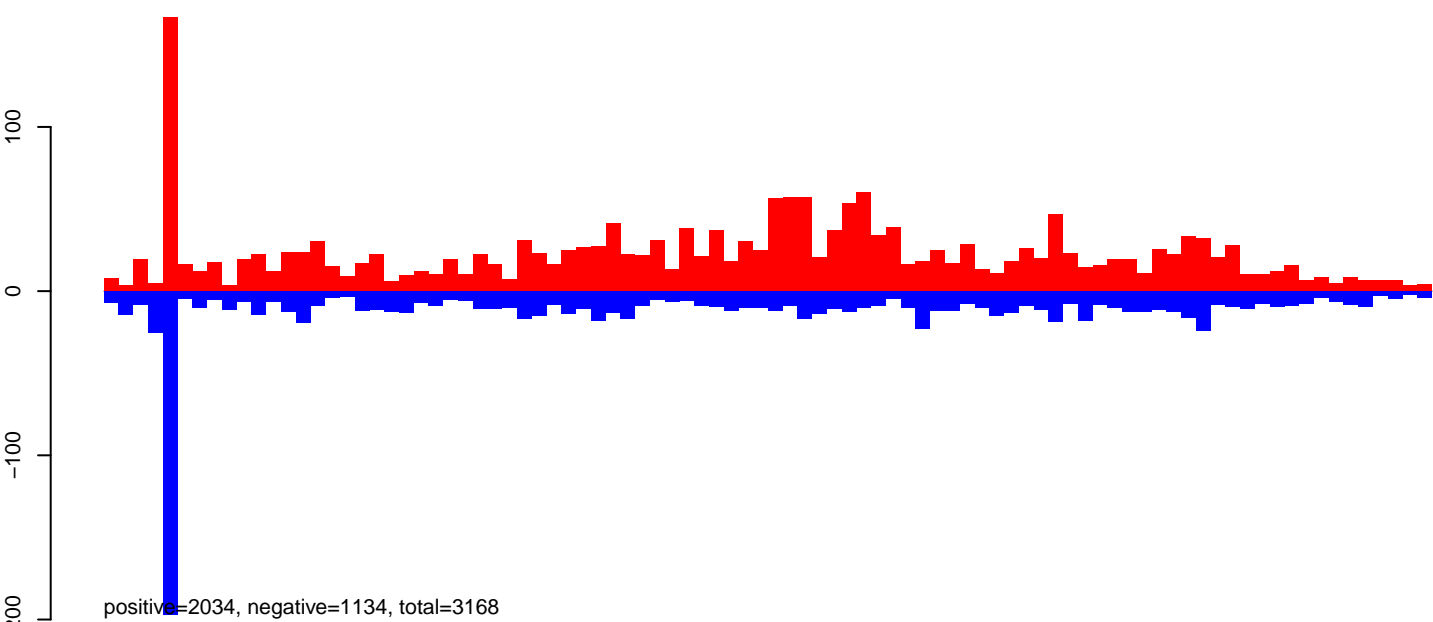
positive=2, negative=442, total=443

Window size=50, length=6690, TE@GYPSY39-LTR-I_AG:1-6690

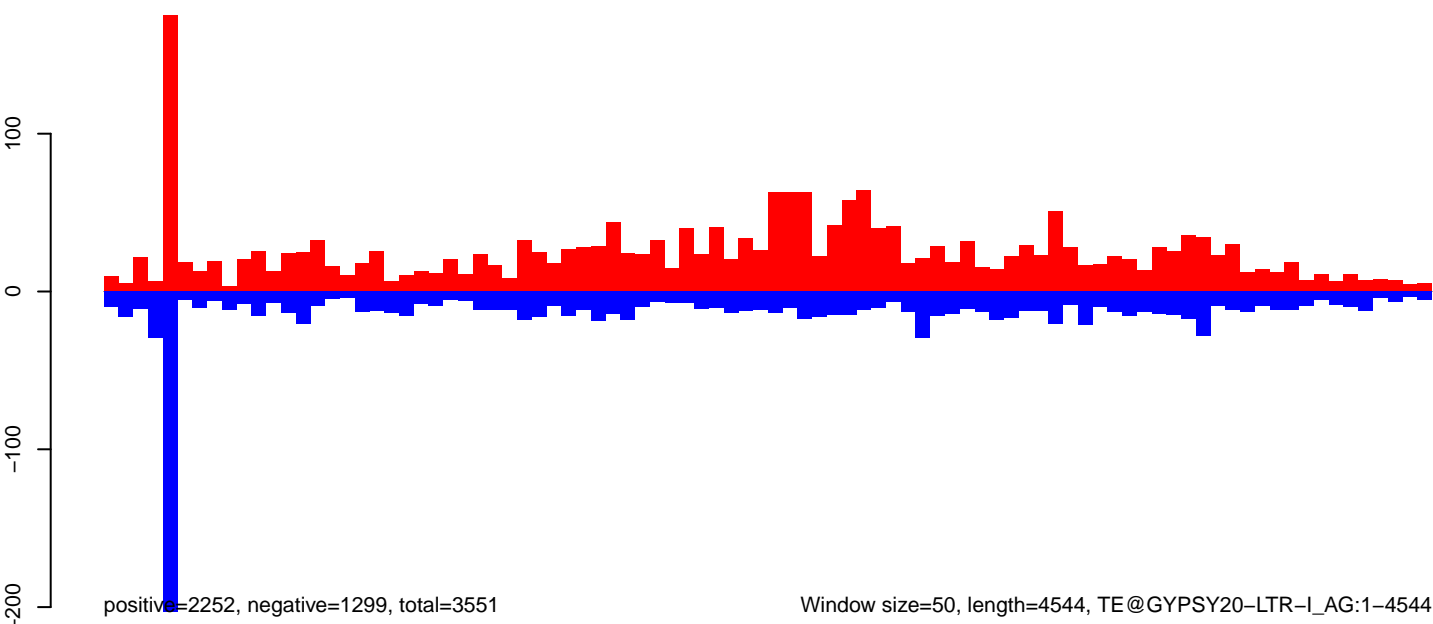
AnGam_Mos55_cells.18_23.rep



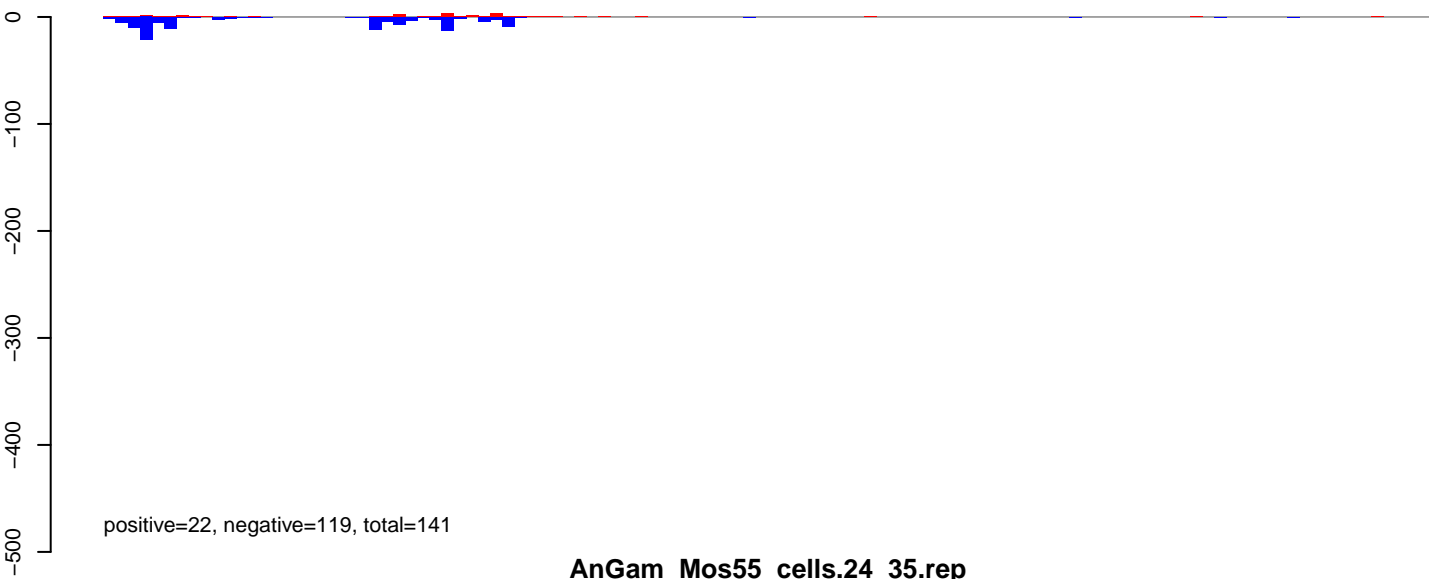
AnGam_Mos55_cells.24_35.rep



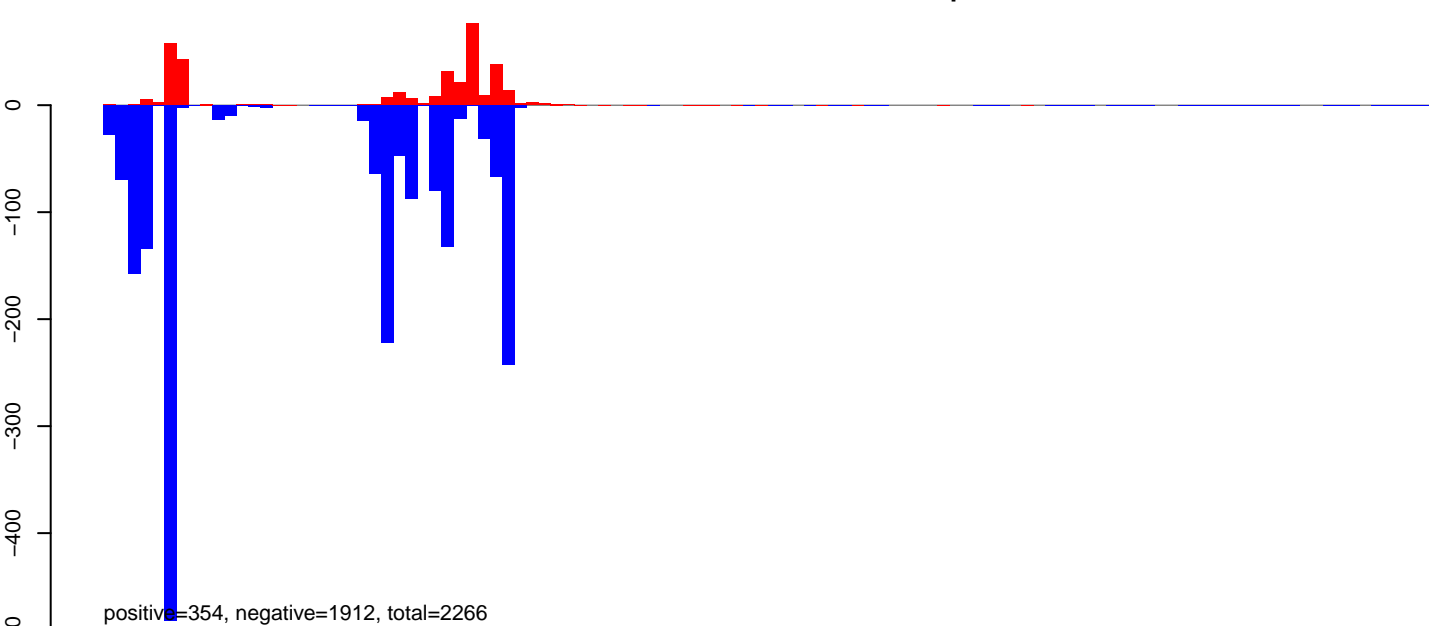
AnGam_Mos55_cells.rep



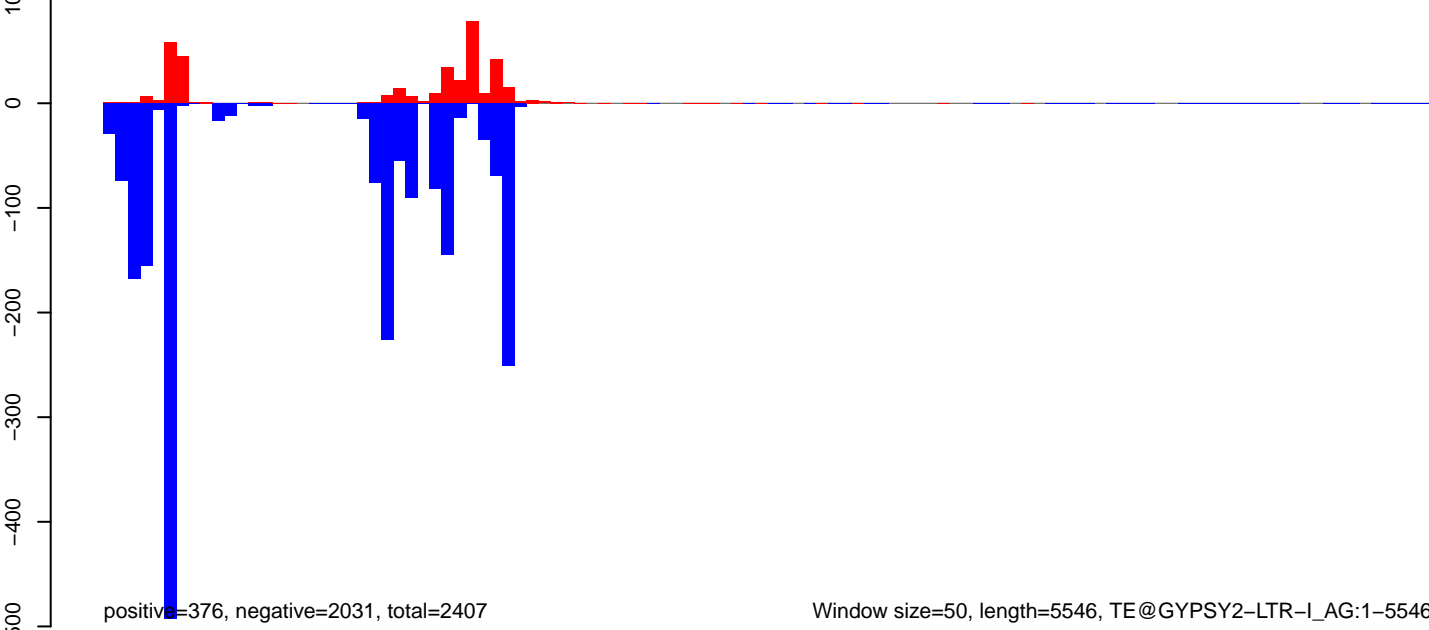
AnGam_Mos55_cells.18_23.rep



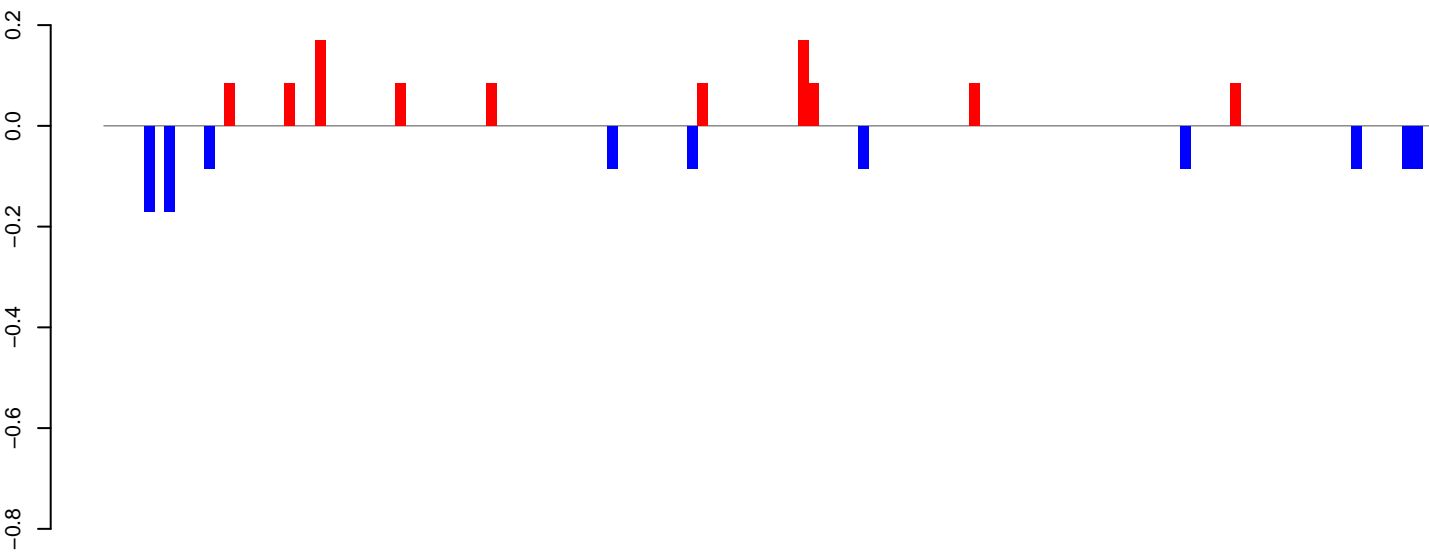
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

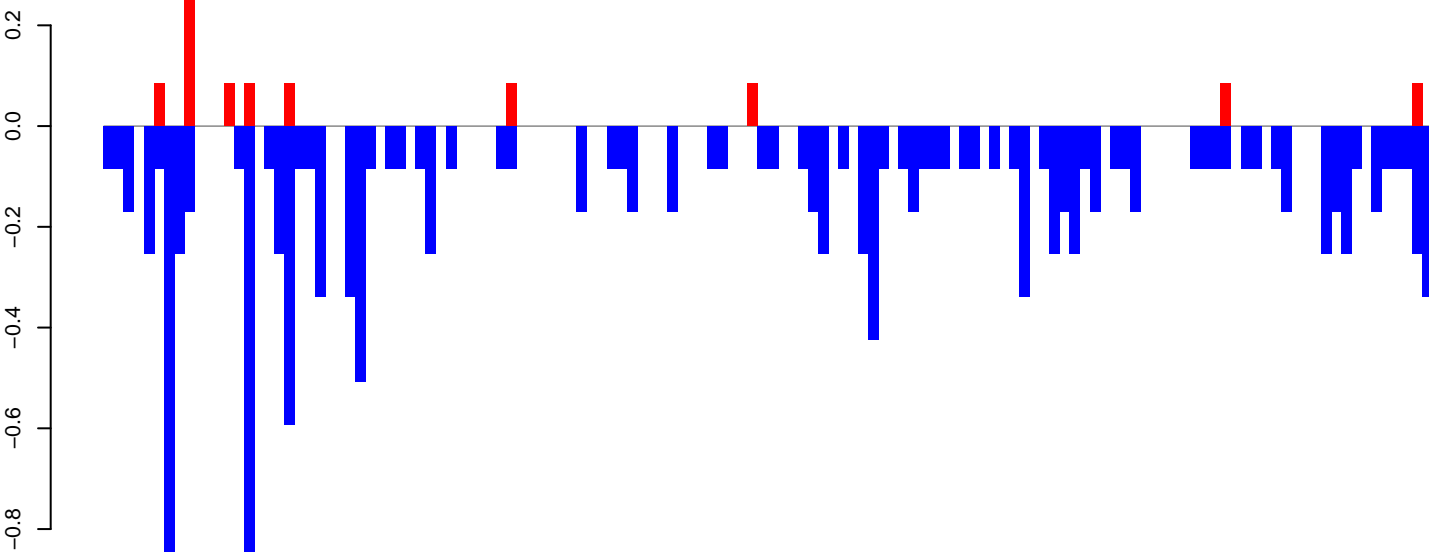


AnGam_Mos55_cells.18_23.rep



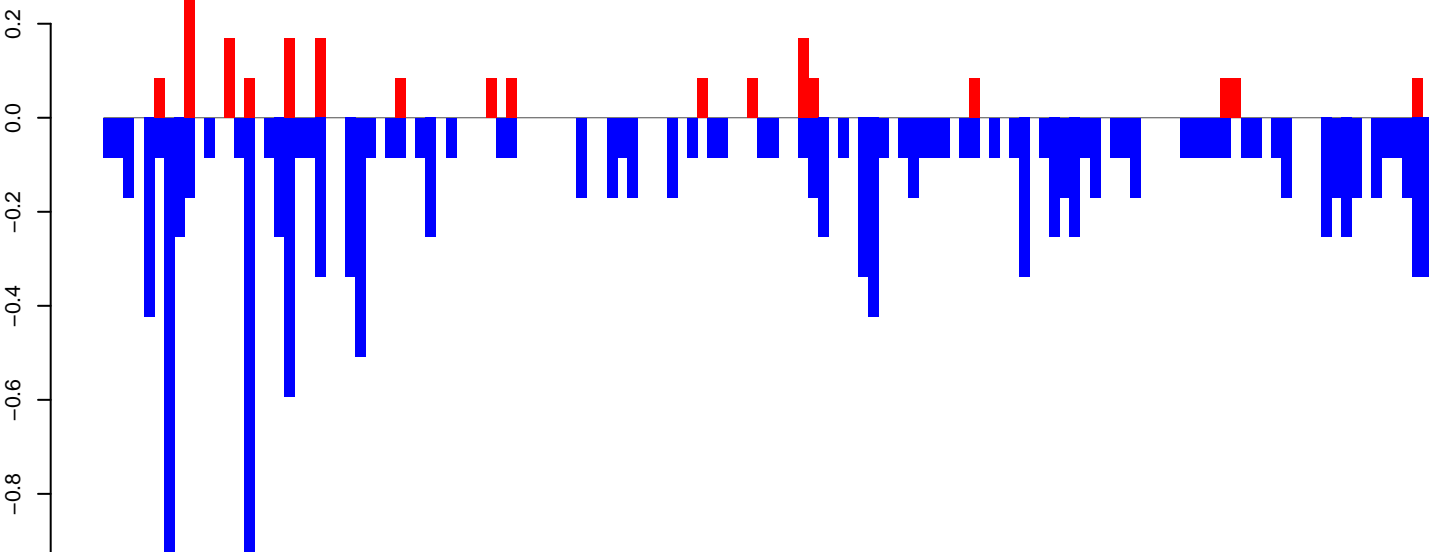
positive=1, negative=1, total=2

AnGam_Mos55_cells.24_35.rep



positive=1, negative=14, total=15

AnGam_Mos55_cells.rep

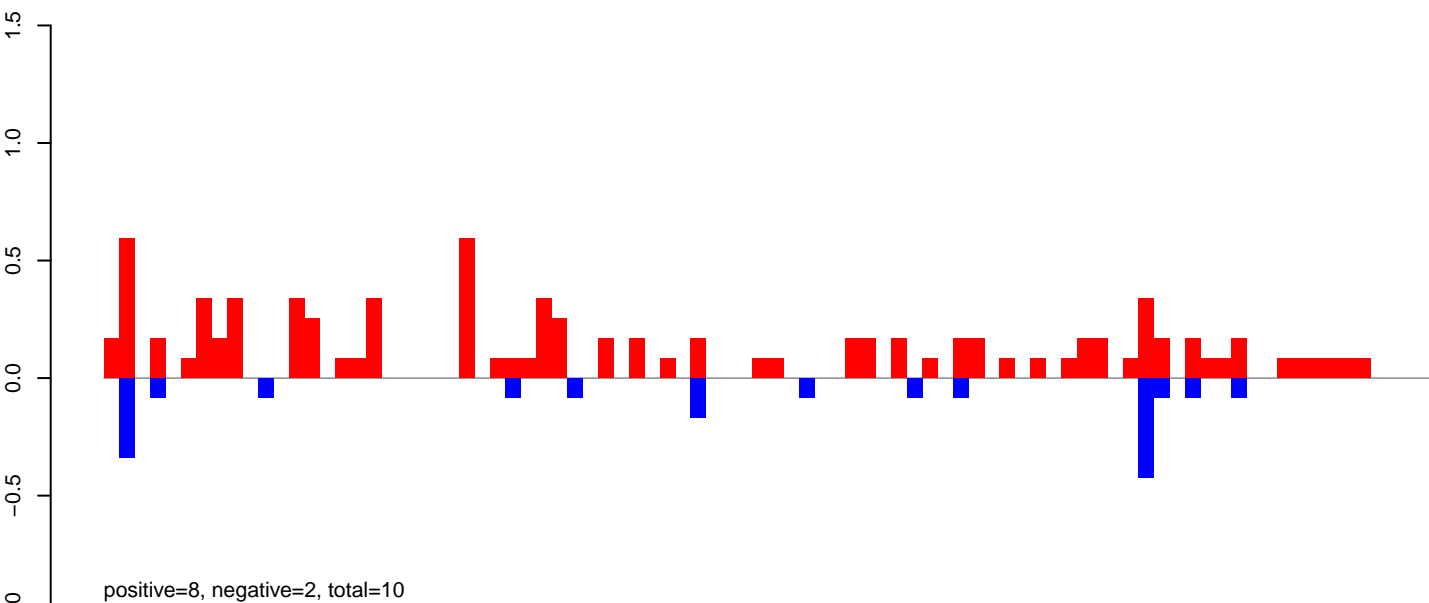


positive=2, negative=15, total=17

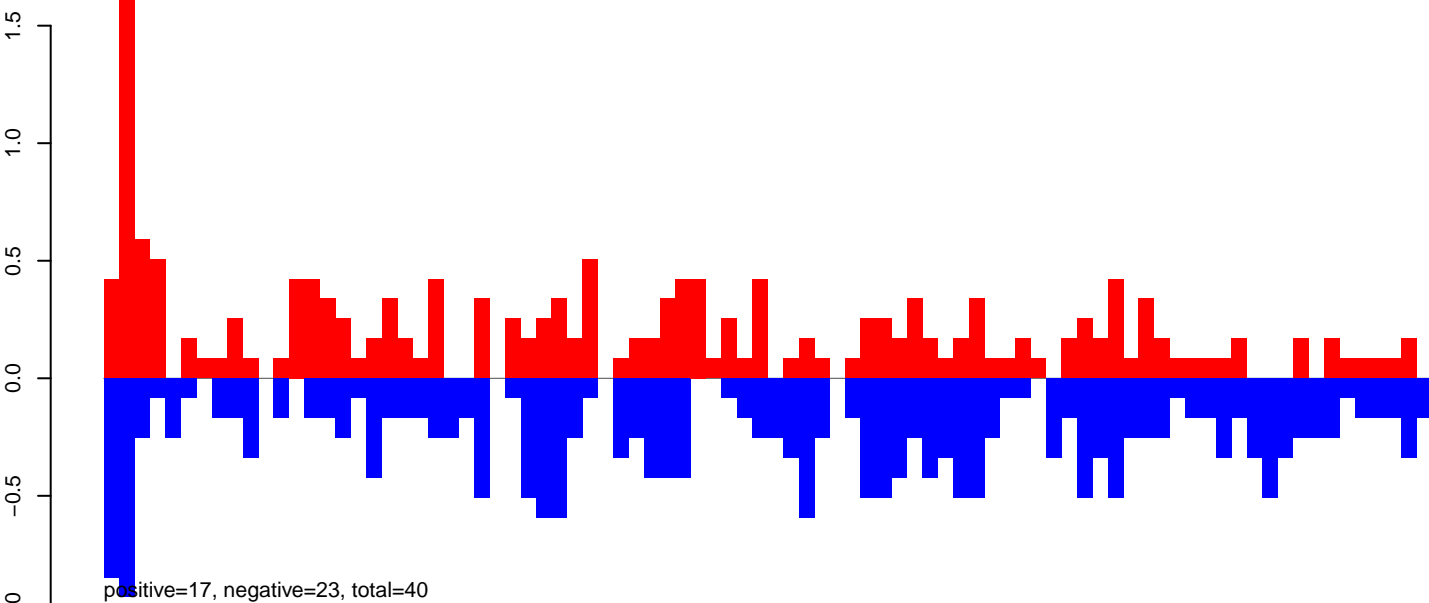
Window size=50, length=6638, TE@TF001937-Pao_Bel_Ele6:1-6638

0 1000 2000 3000 4000 5000 6000

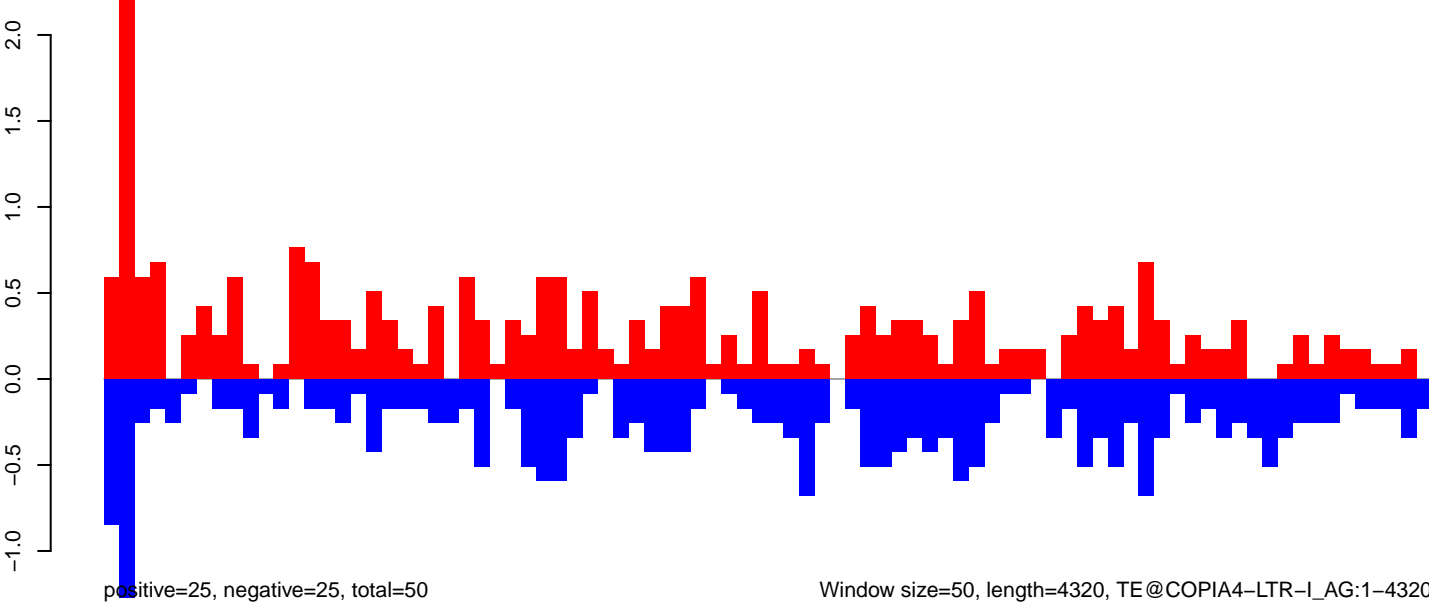
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



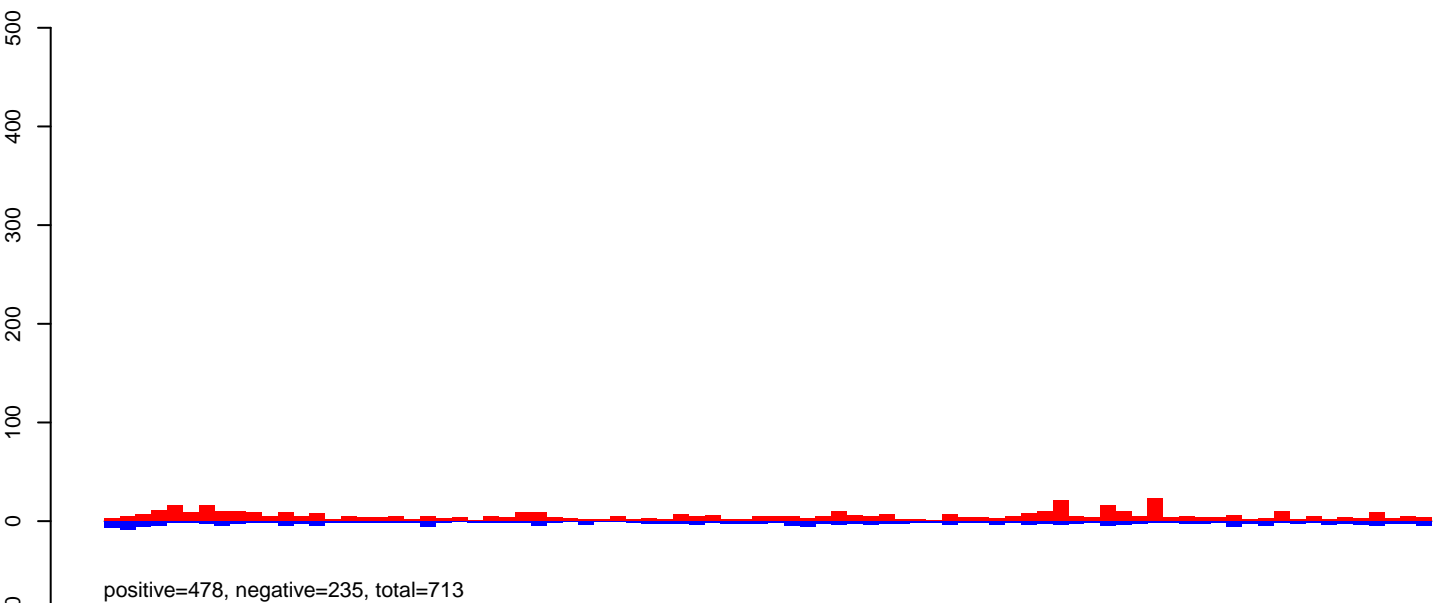
AnGam_Mos55_cells.rep



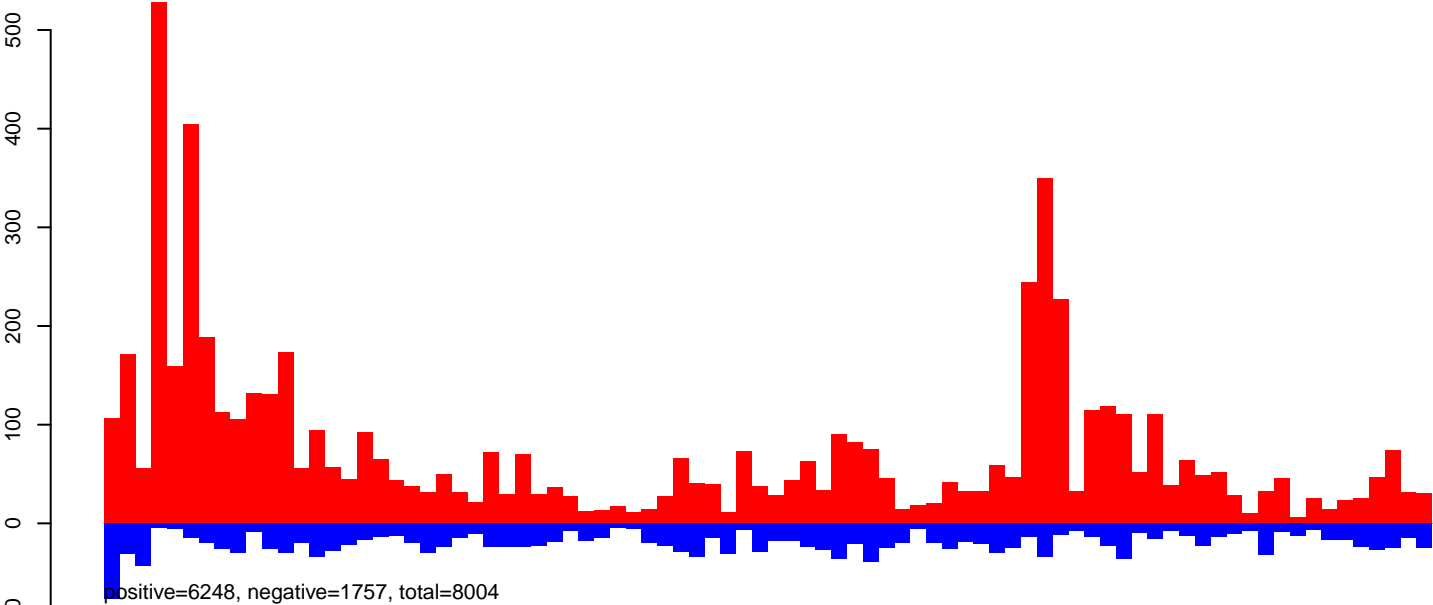
Window size=50, length=4320, TE@COPIA4-LTR-I_AG:1-4320

0 1000 2000 3000 4000

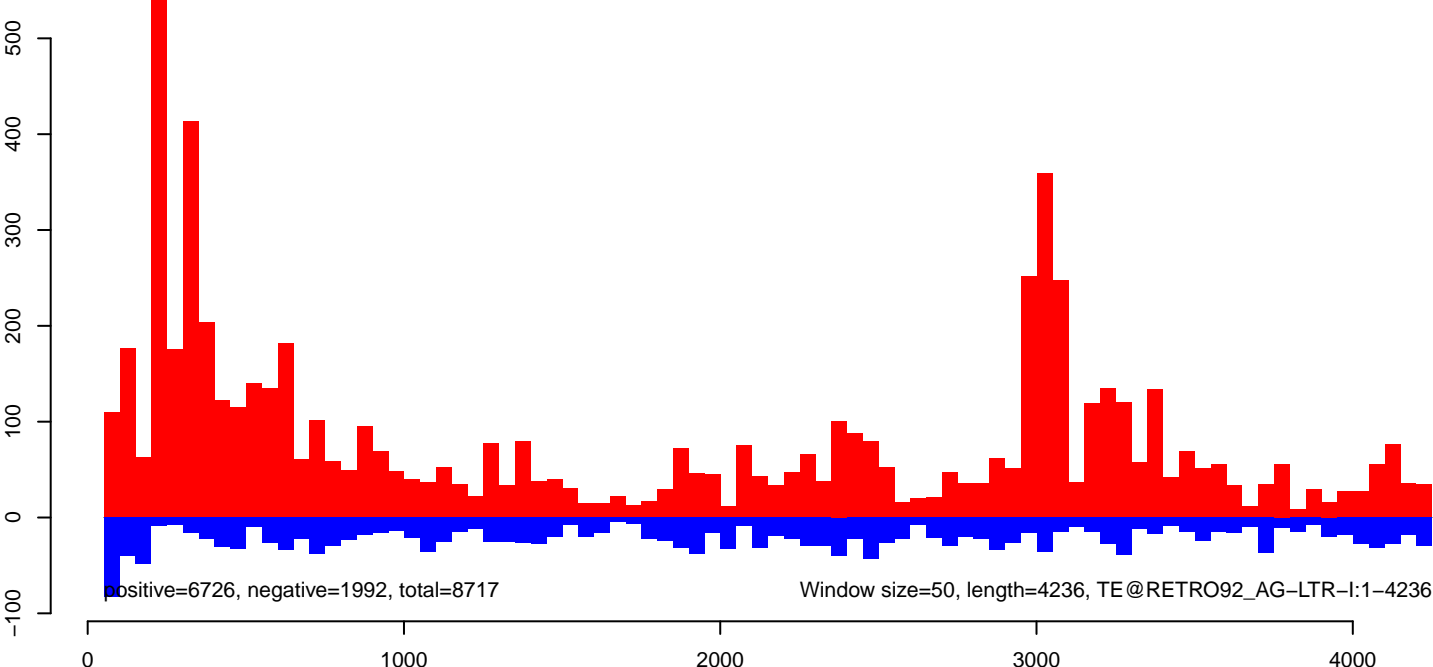
AnGam_Mos55_cells.18_23.rep



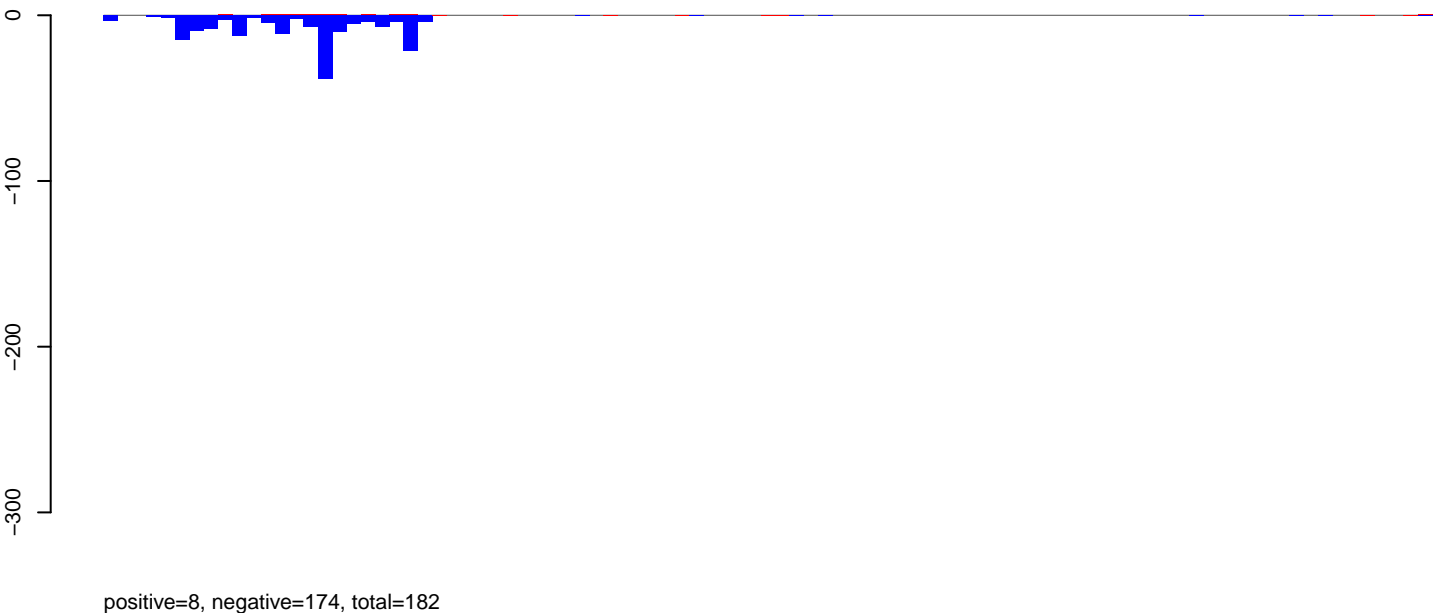
AnGam_Mos55_cells.24_35.rep



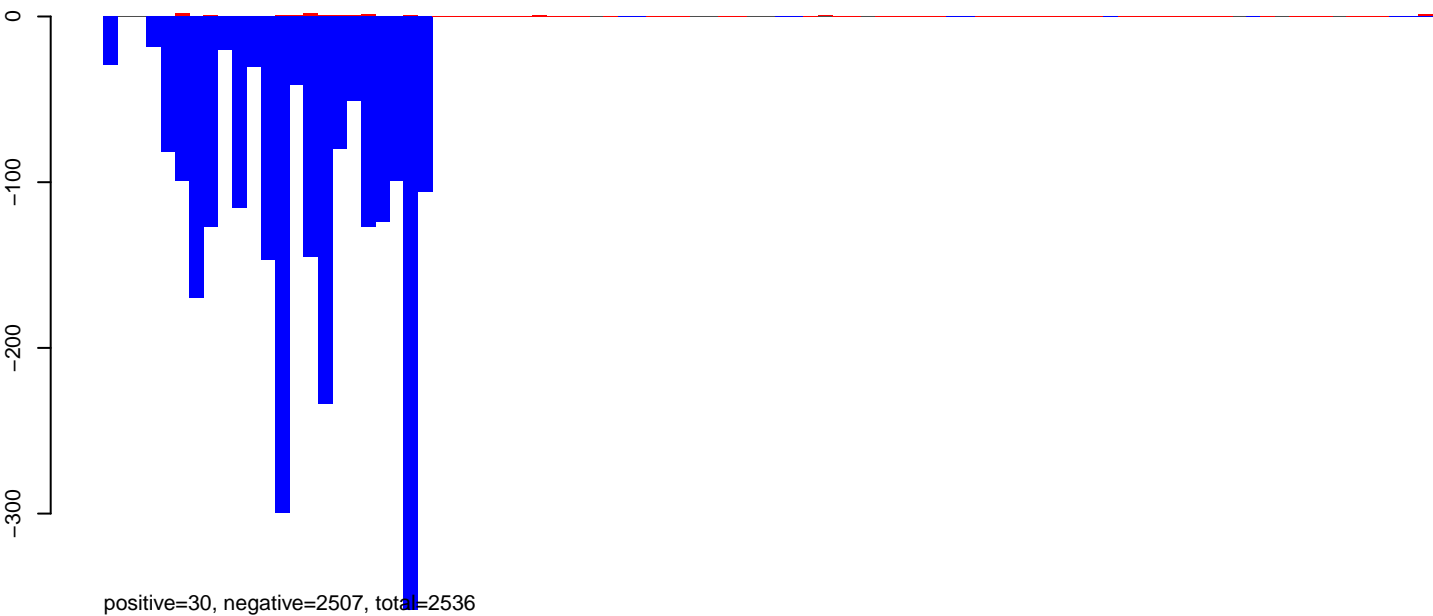
AnGam_Mos55_cells.rep



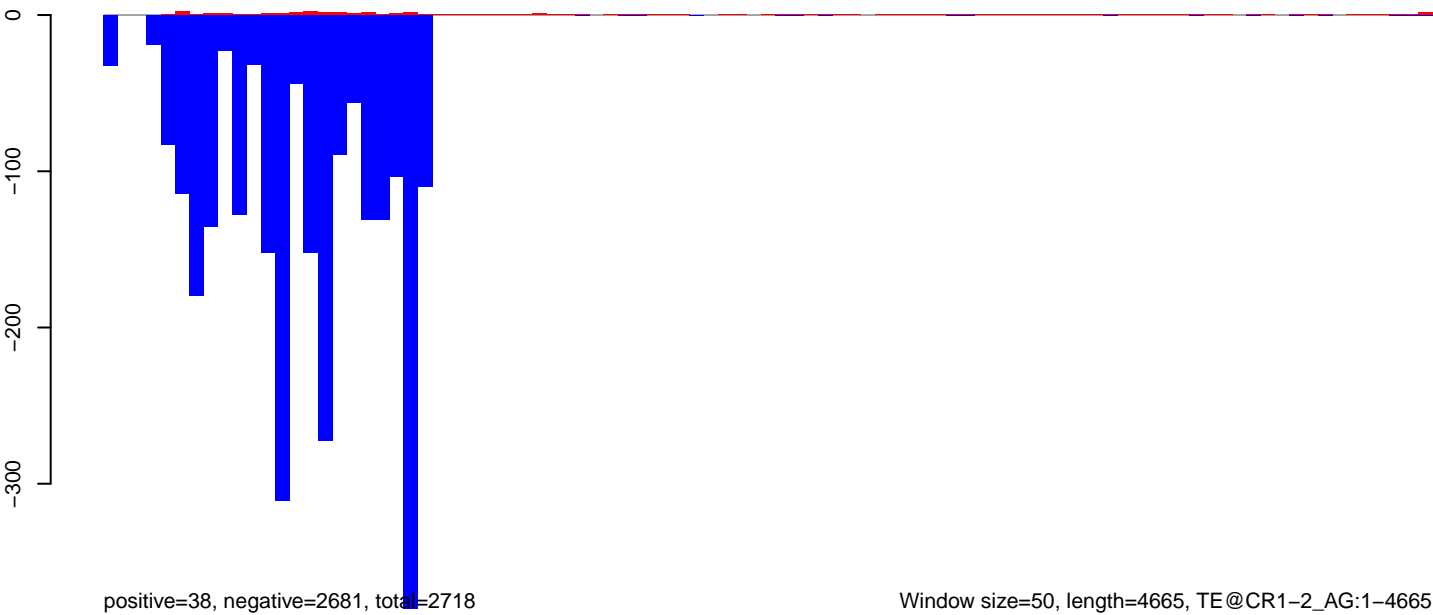
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

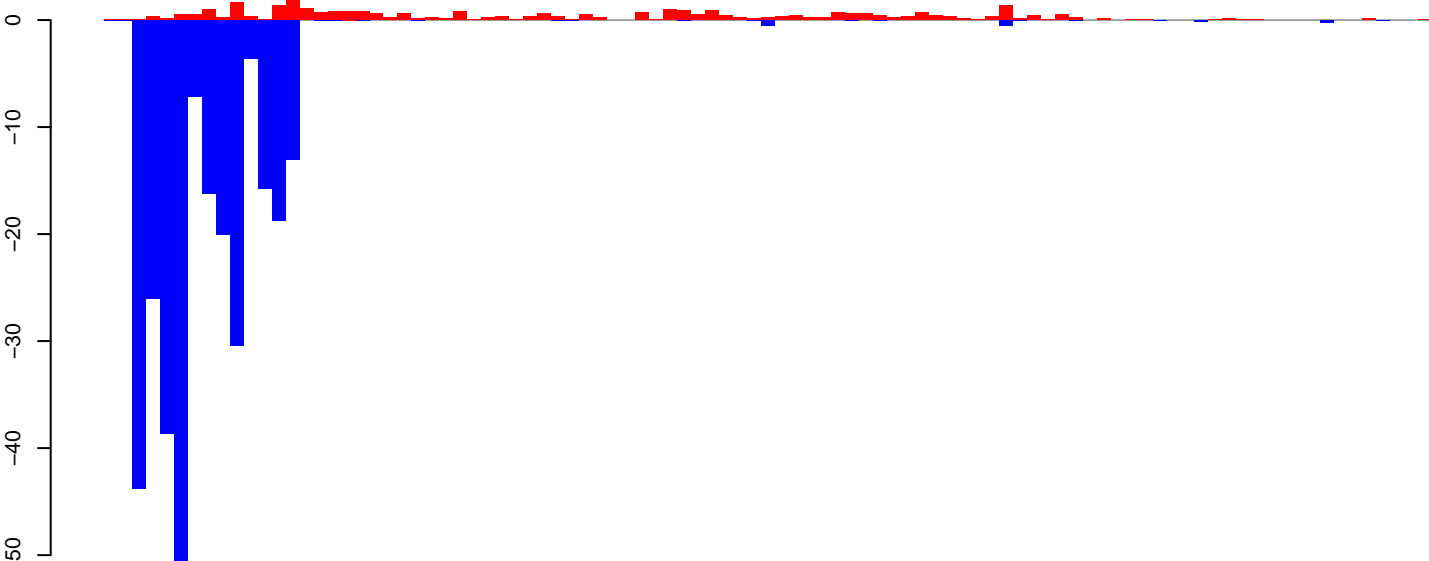


AnGam_Mos55_cells.18_23.rep



positive=3, negative=19, total=22

AnGam_Mos55_cells.24_35.rep



positive=34, negative=290, total=324

AnGam_Mos55_cells.rep

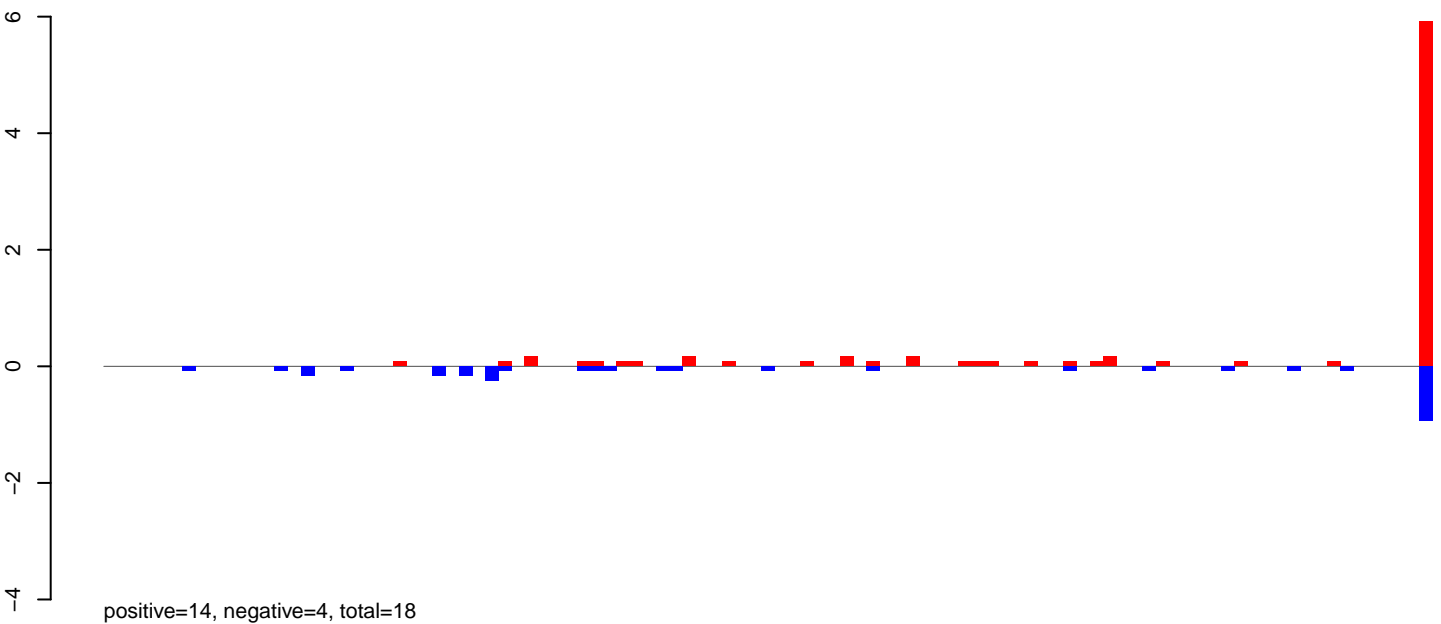


positive=37, negative=309, total=346

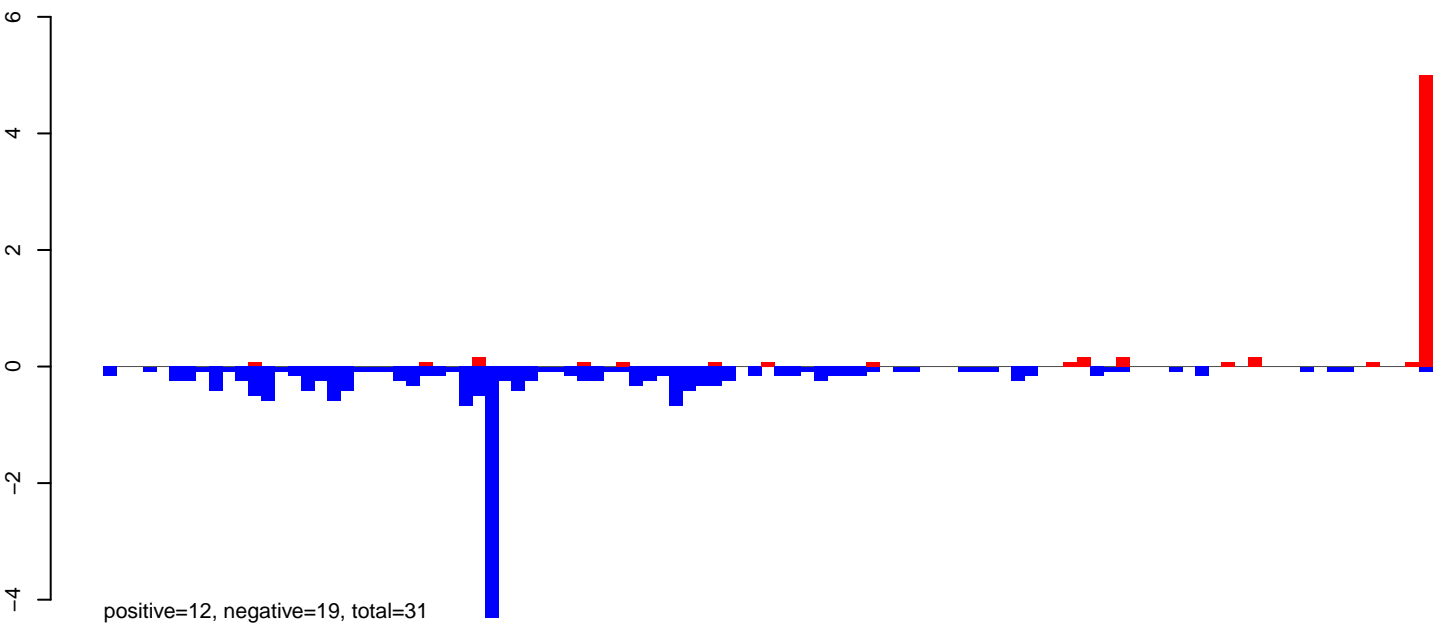
Window size=50, length=4796, TE@GYPSY32-LTR-I_AG:1-4796

0 1000 2000 3000 4000

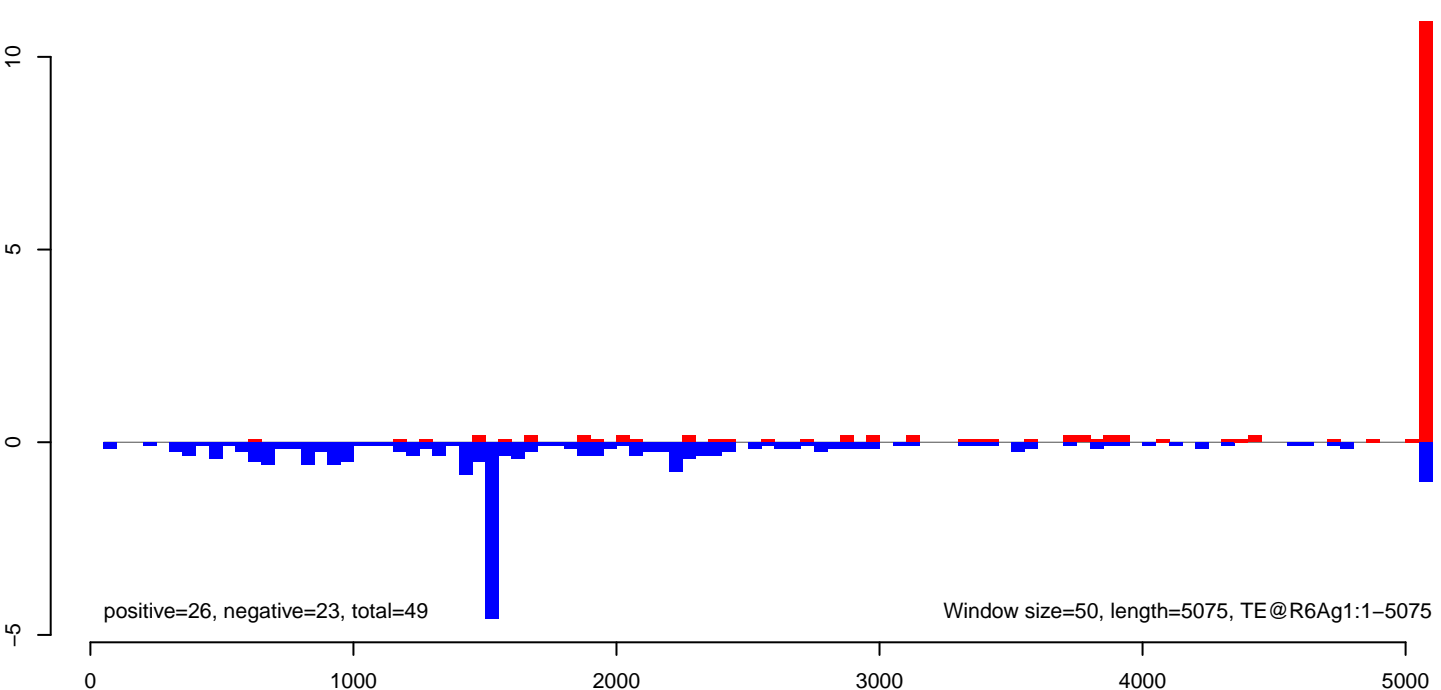
AnGam_Mos55_cells.18_23.rep



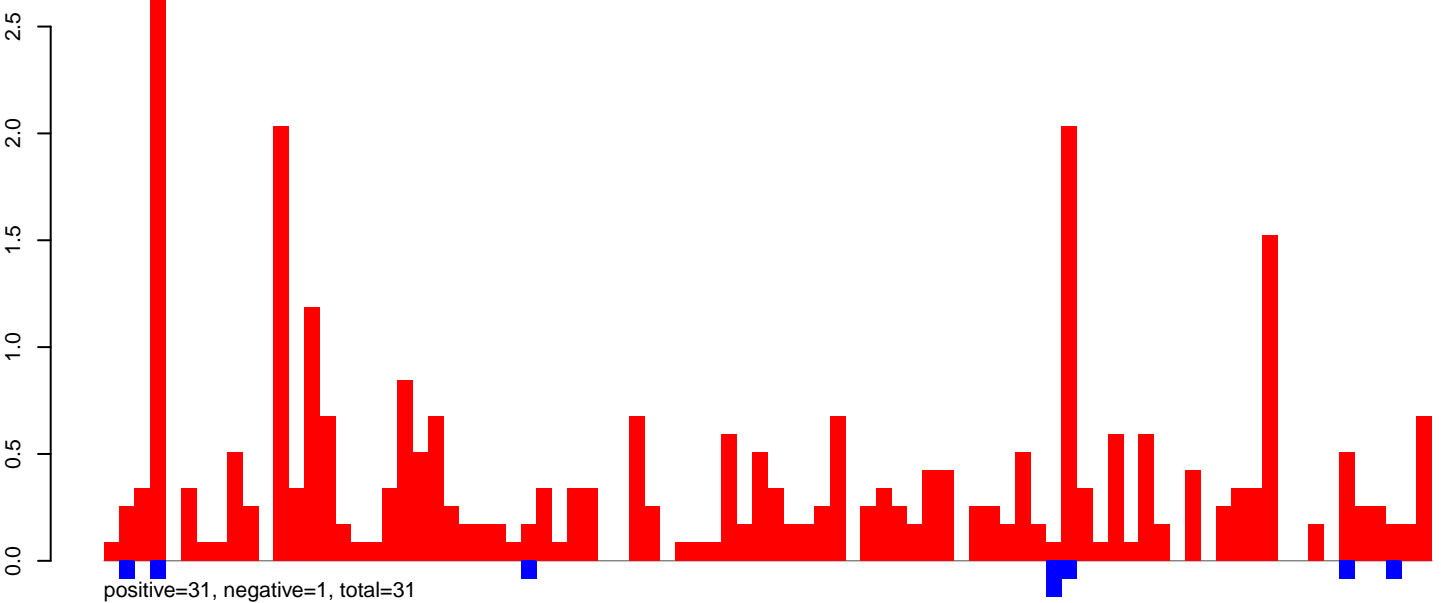
AnGam_Mos55_cells.24_35.rep



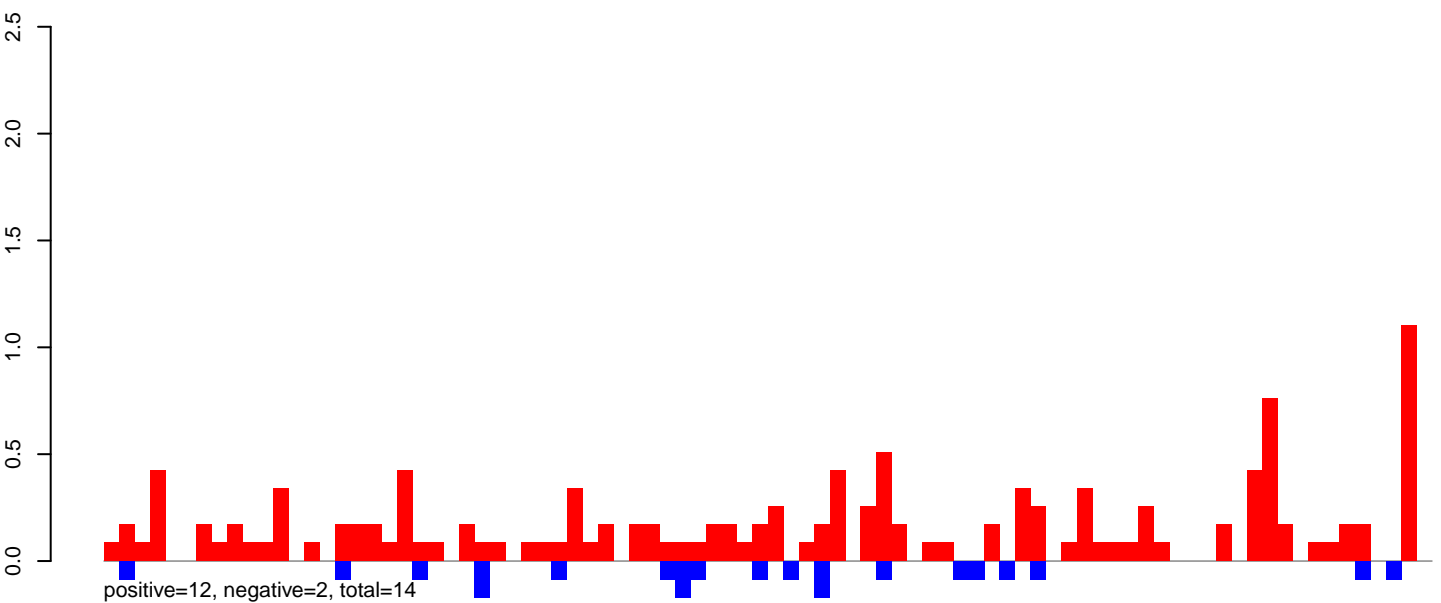
AnGam_Mos55_cells.rep



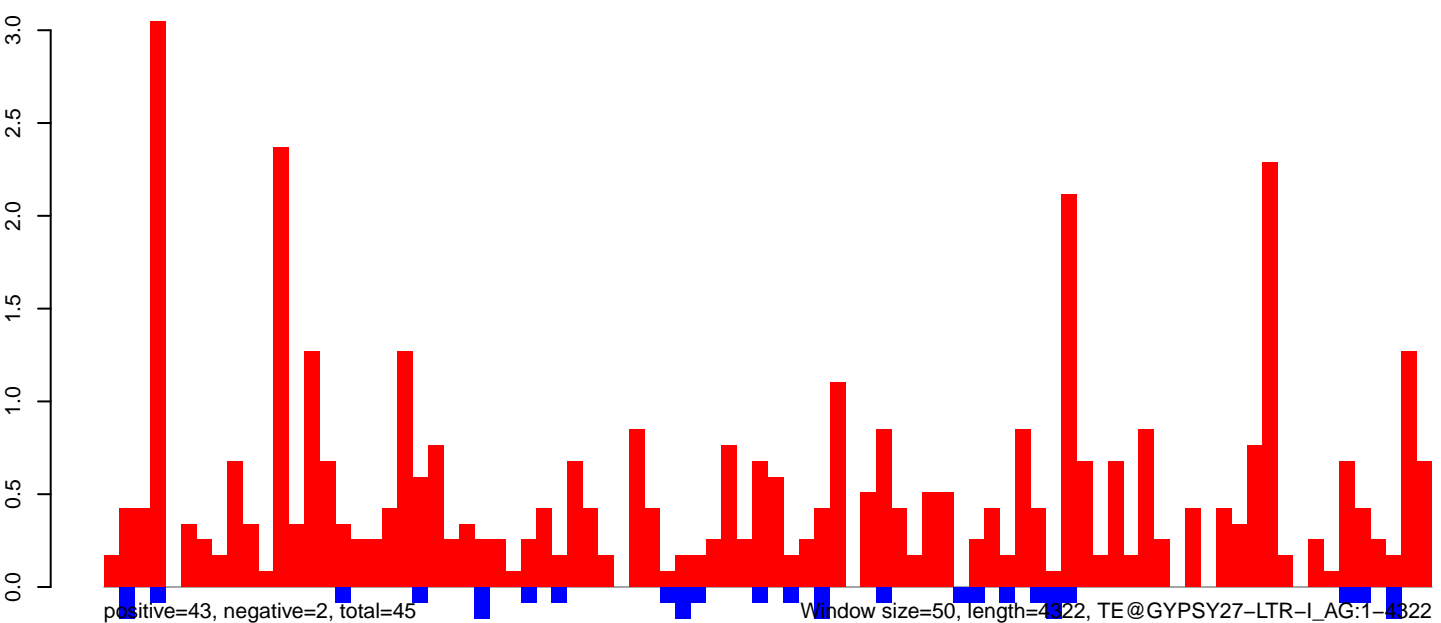
AnGam_Mos55_cells.18_23.rep



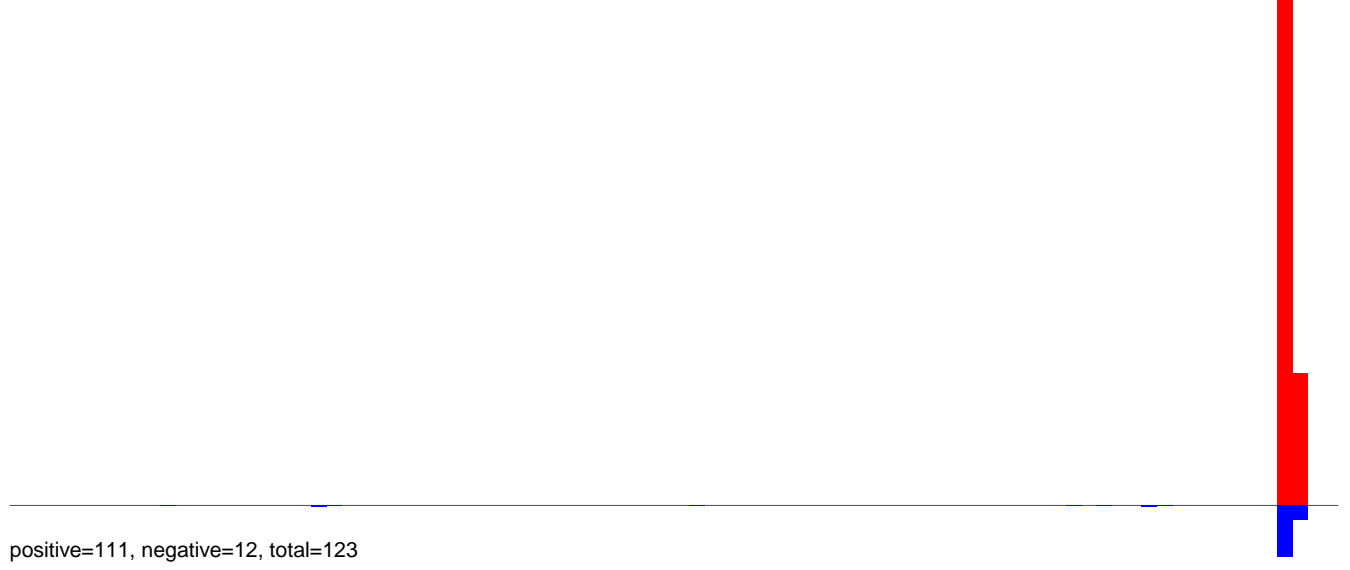
AnGam_Mos55_cells.24_35.rep



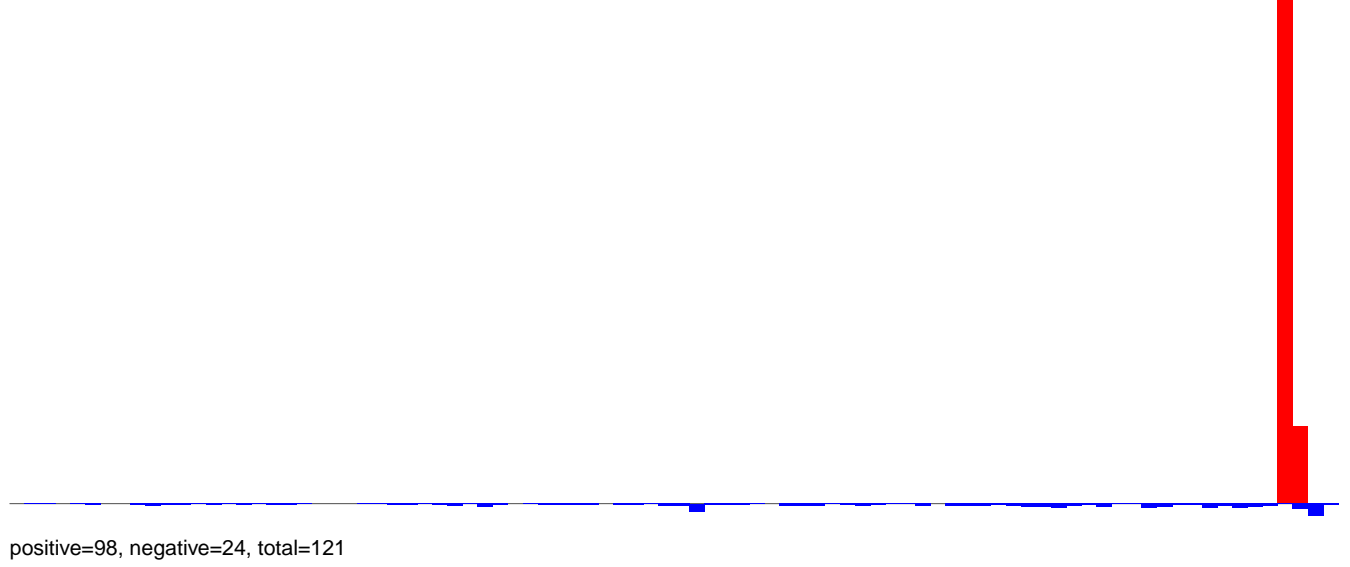
AnGam_Mos55_cells.rep



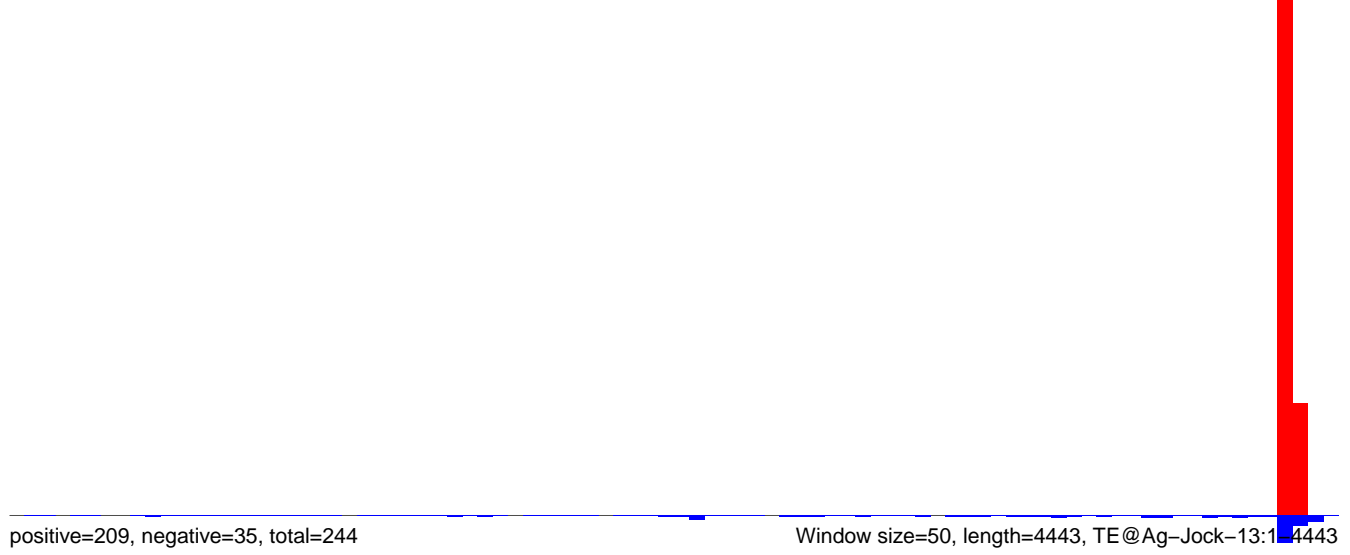
AnGam_Mos55_cells.18_23.rep



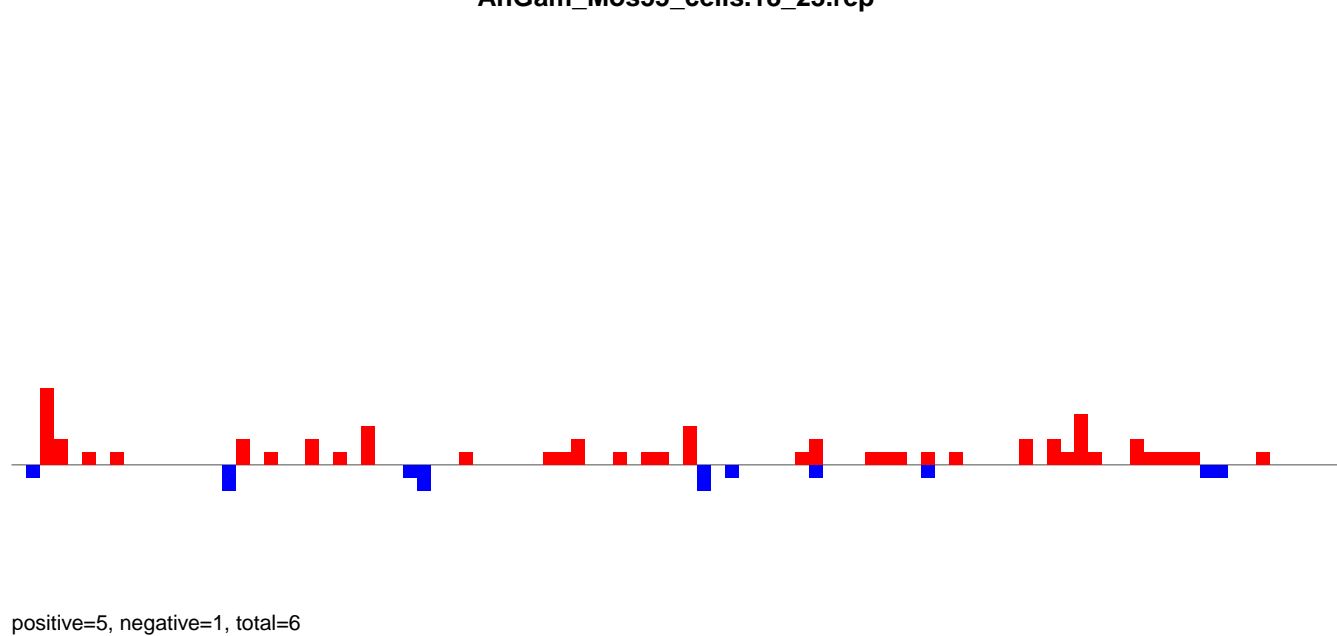
AnGam_Mos55_cells.24_35.rep



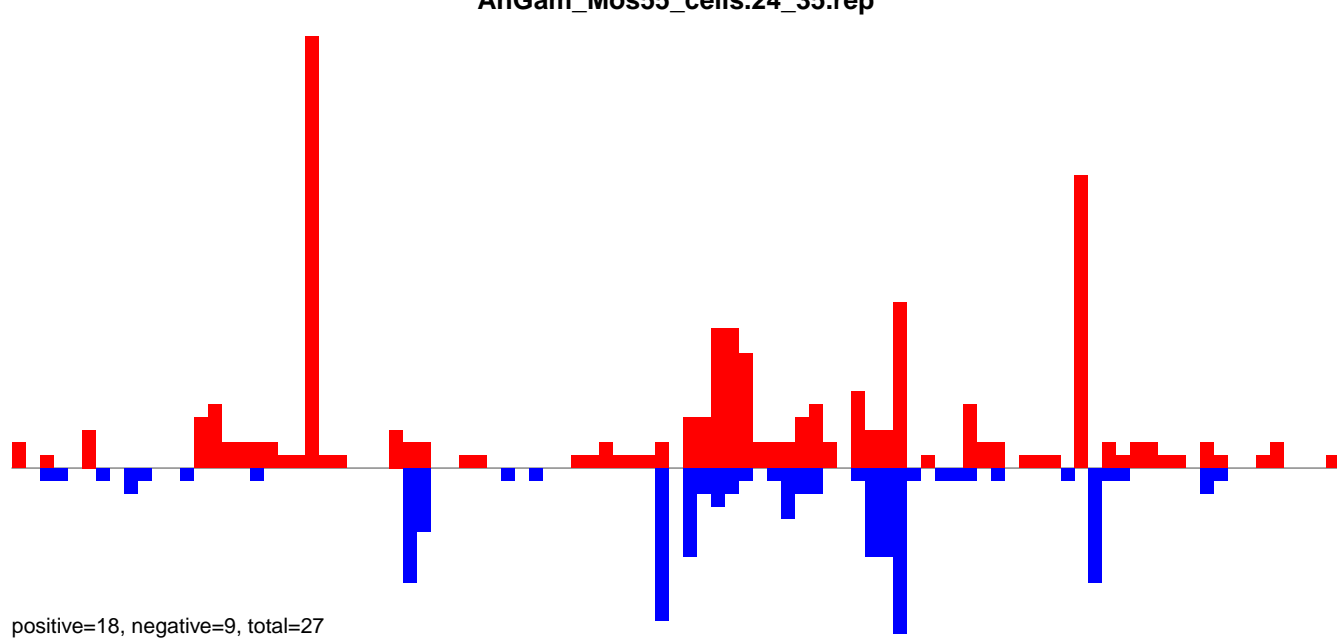
AnGam_Mos55_cells.rep



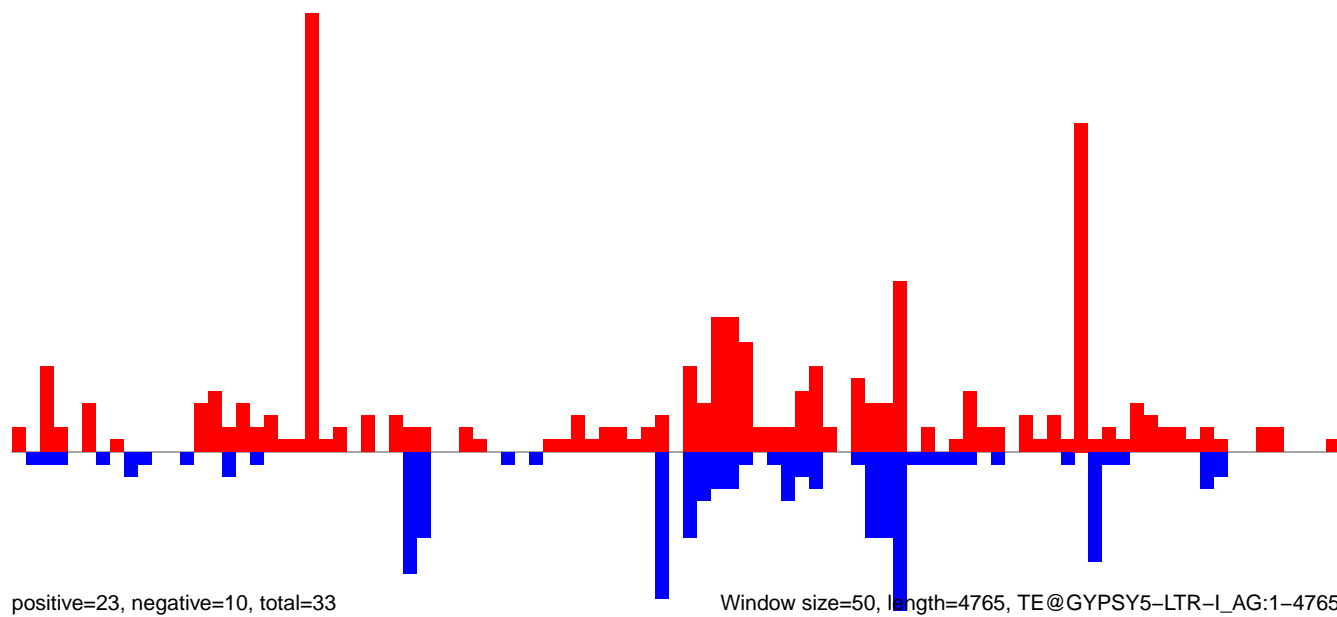
AnGam_Mos55_cells.18_23.rep



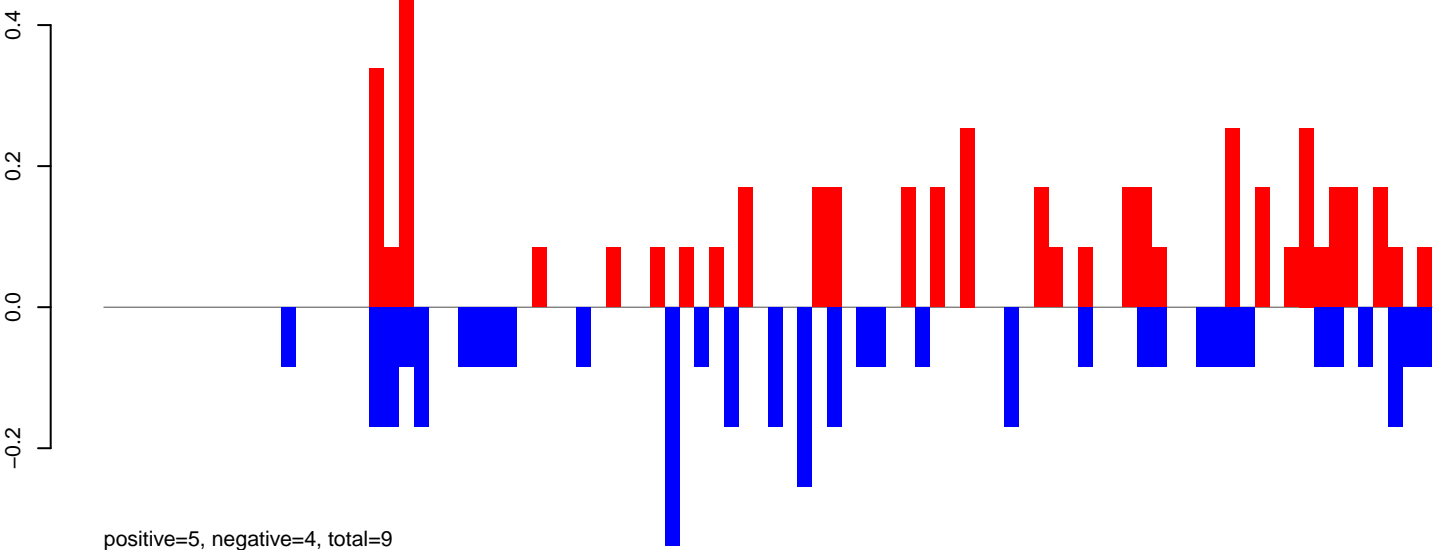
AnGam_Mos55_cells.24_35.rep



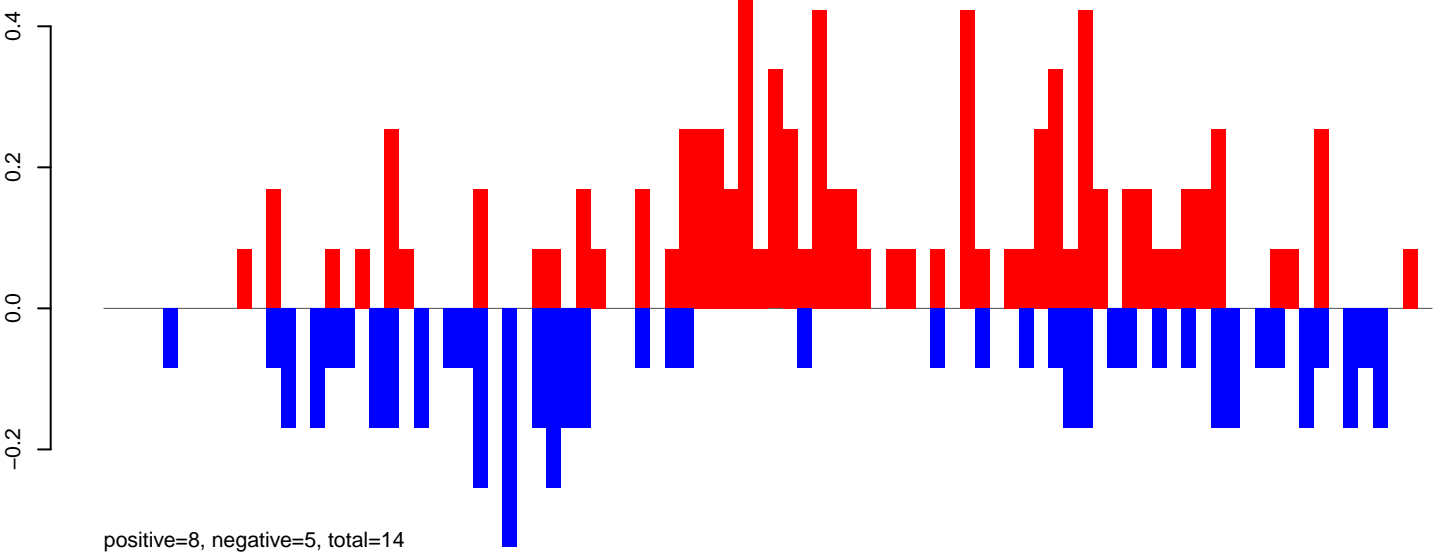
AnGam_Mos55_cells.rep



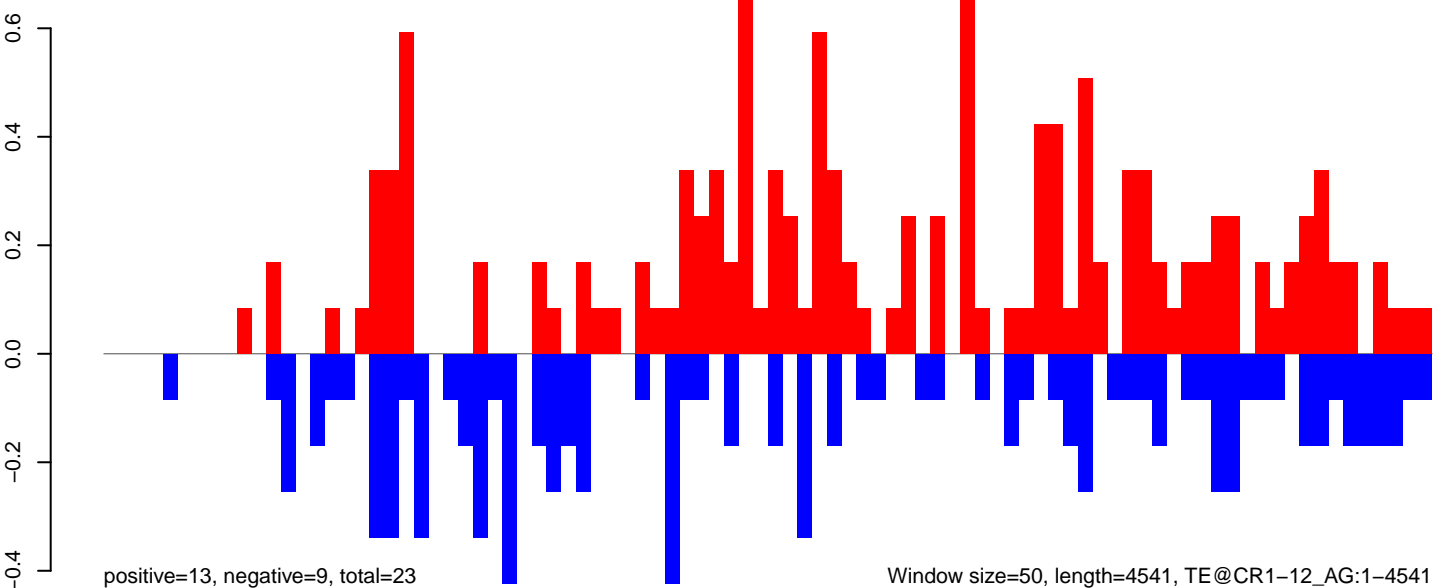
AnGam_Mos55_cells.18_23.rep



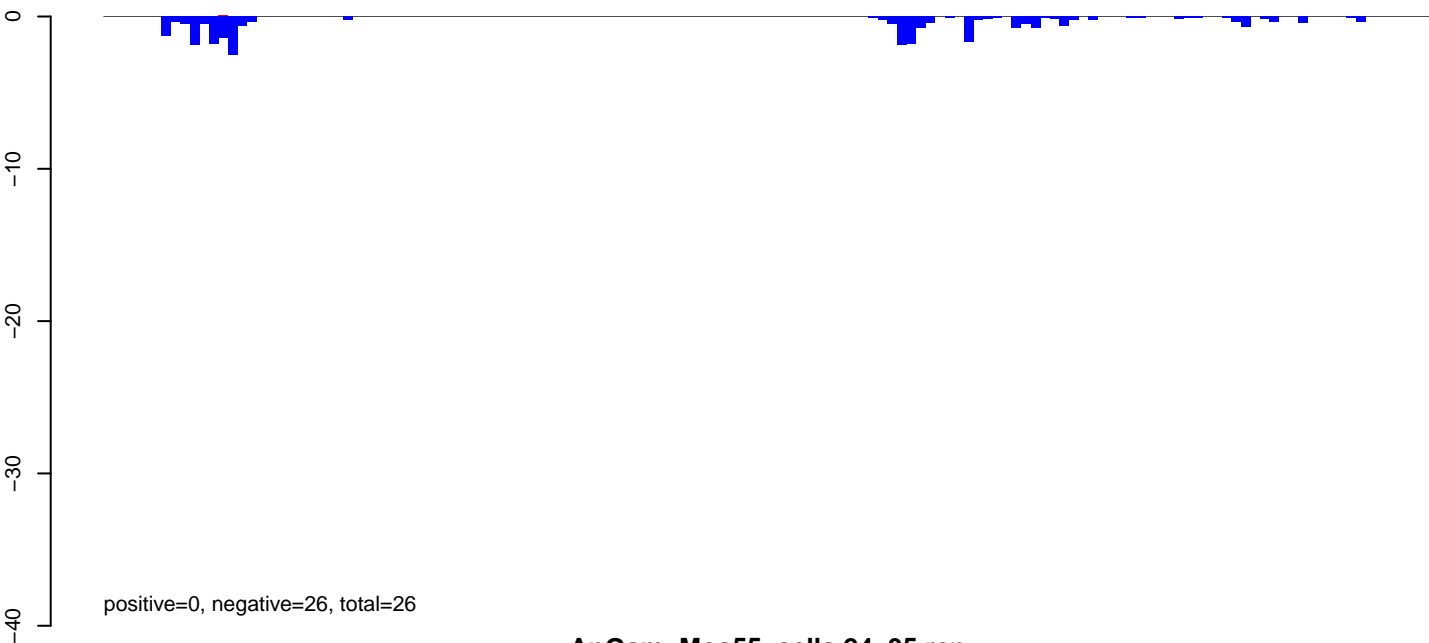
AnGam_Mos55_cells.24_35.rep



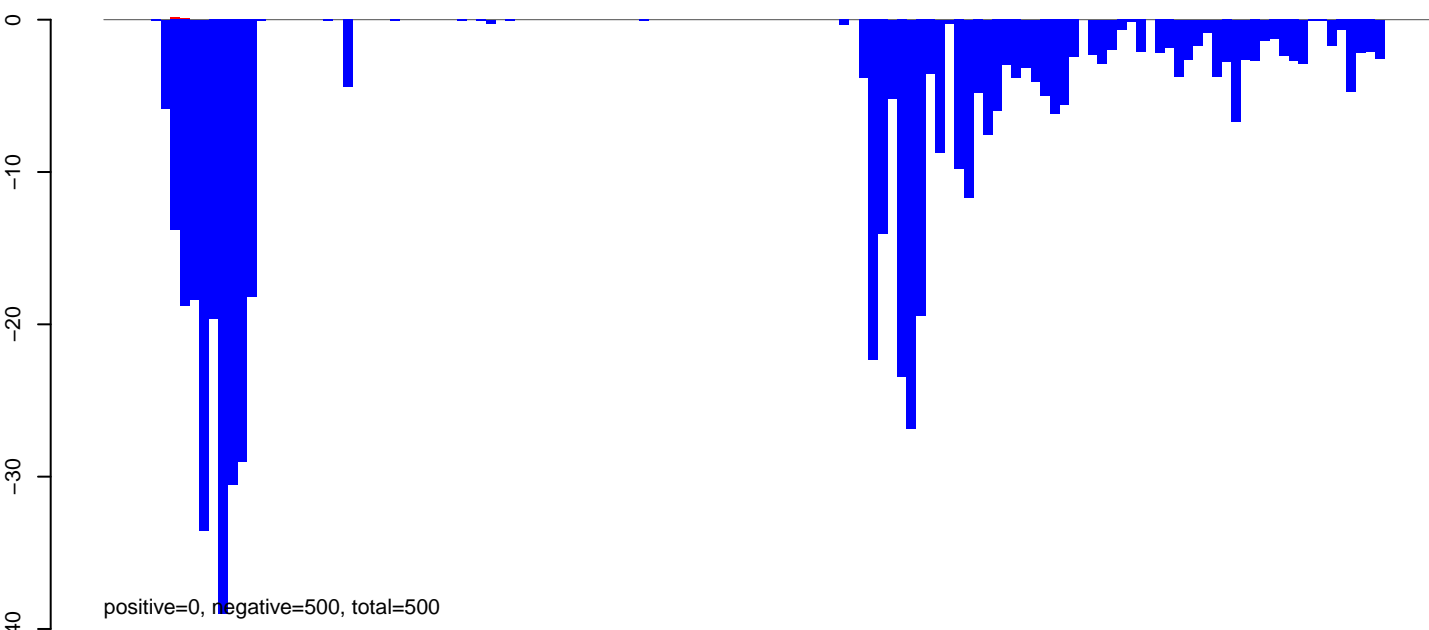
AnGam_Mos55_cells.rep



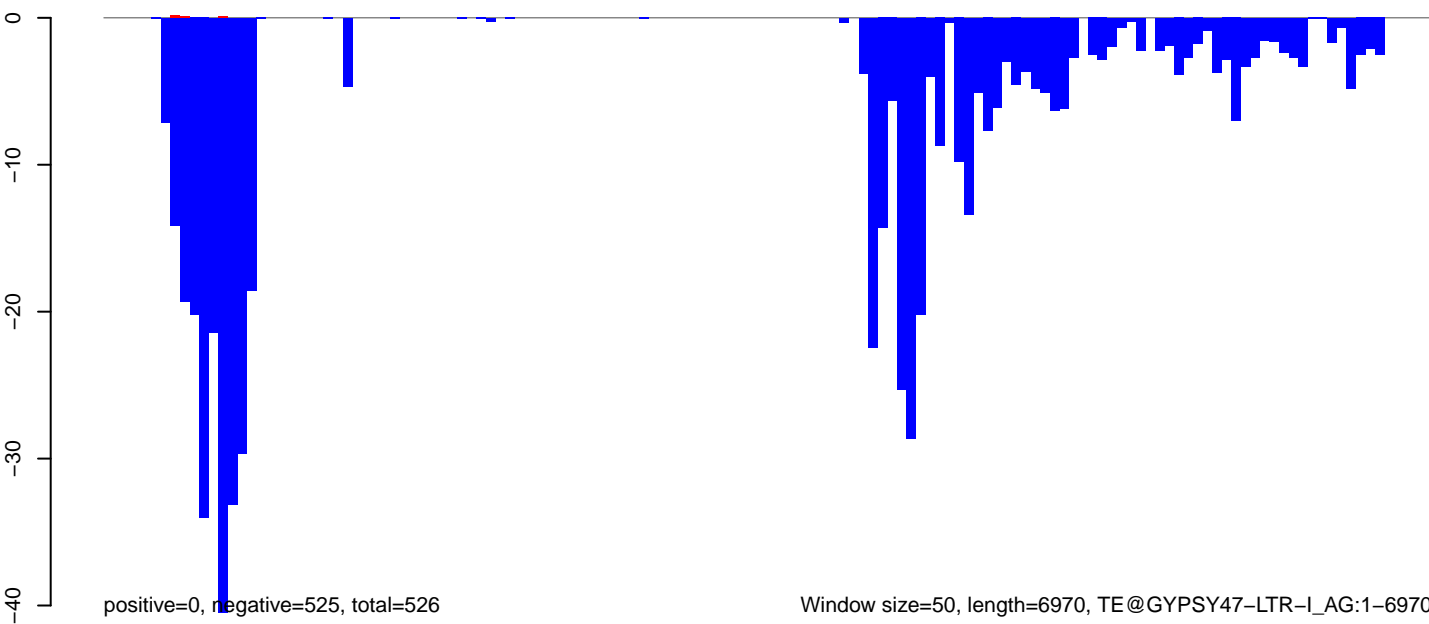
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



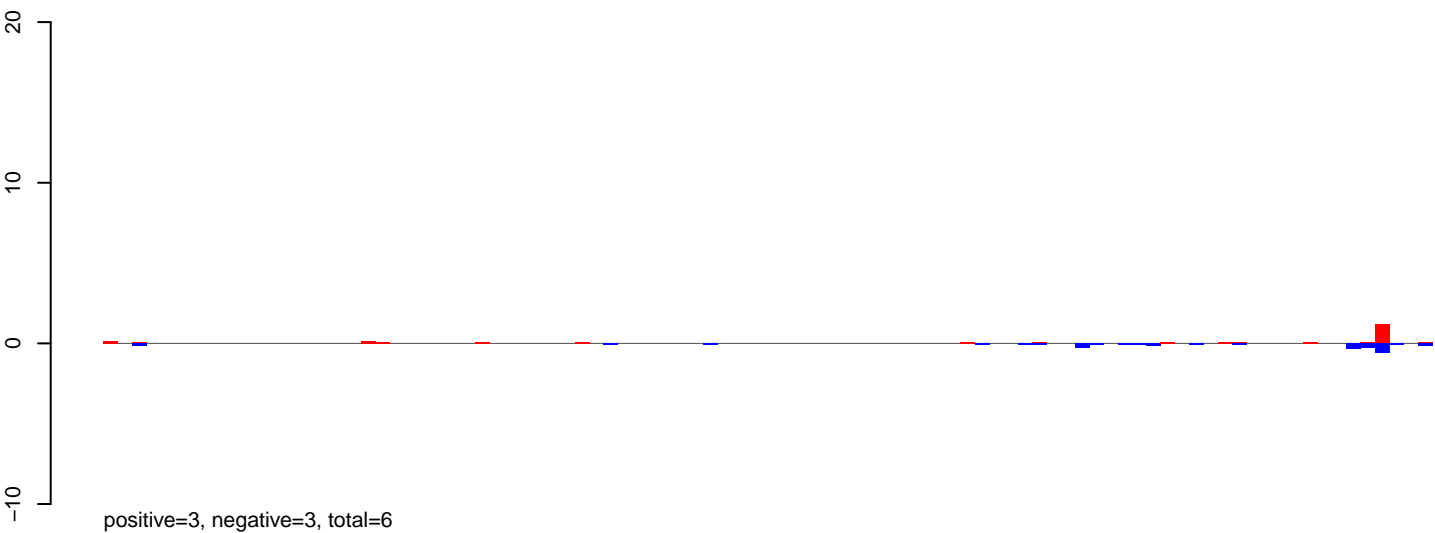
AnGam_Mos55_cells.rep



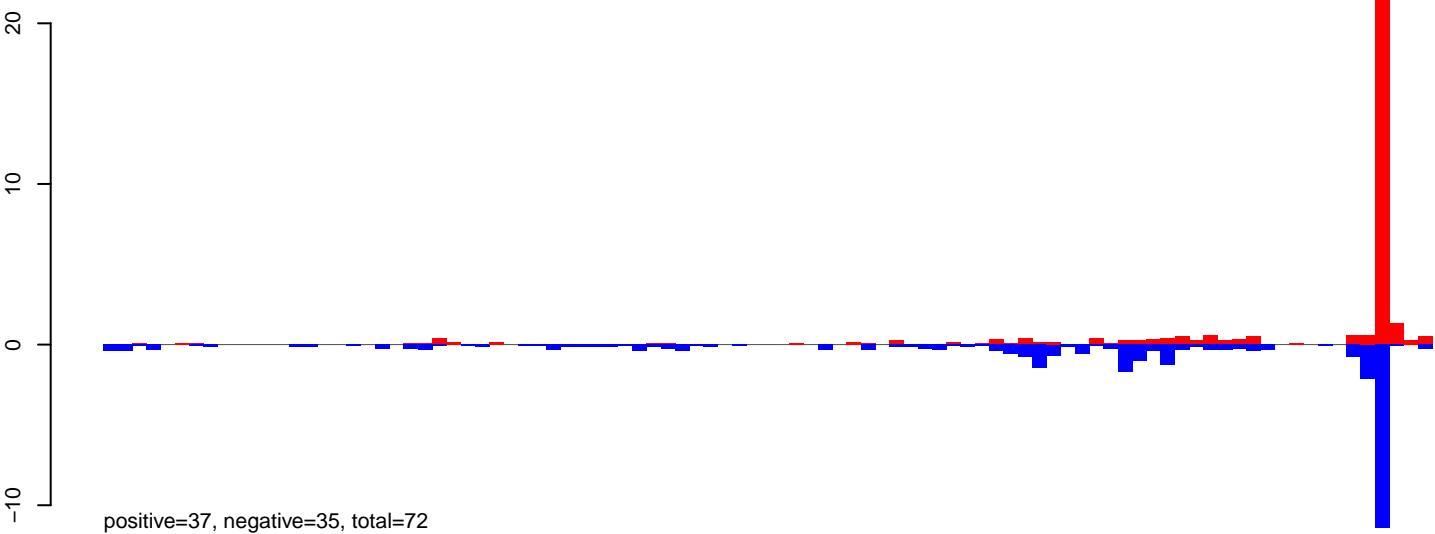
Window size=50, length=6970, TE@GYPSY47-LTR-I_AG:1-6970

0 1000 2000 3000 4000 5000 6000 7000

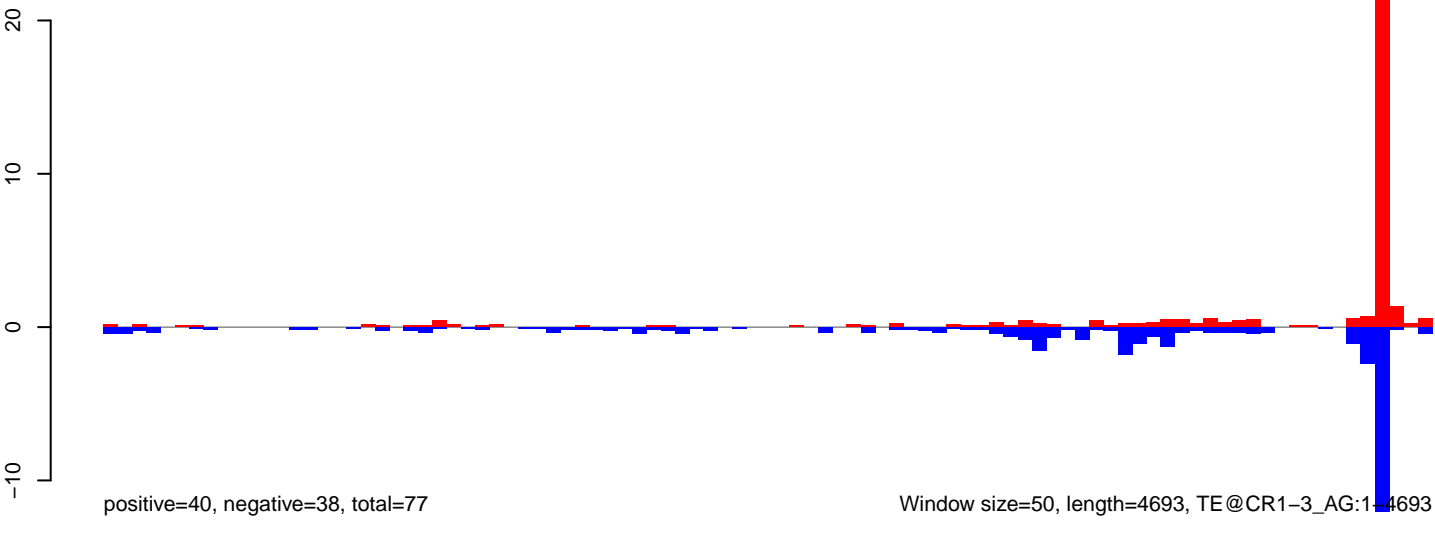
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep

1.5
1.0
0.5
0.0
-0.5

positive=1, negative=2, total=3

AnGam_Mos55_cells.24_35.rep

1.5
1.0
0.5
0.0
-0.5

positive=3, negative=13, total=15

AnGam_Mos55_cells.rep

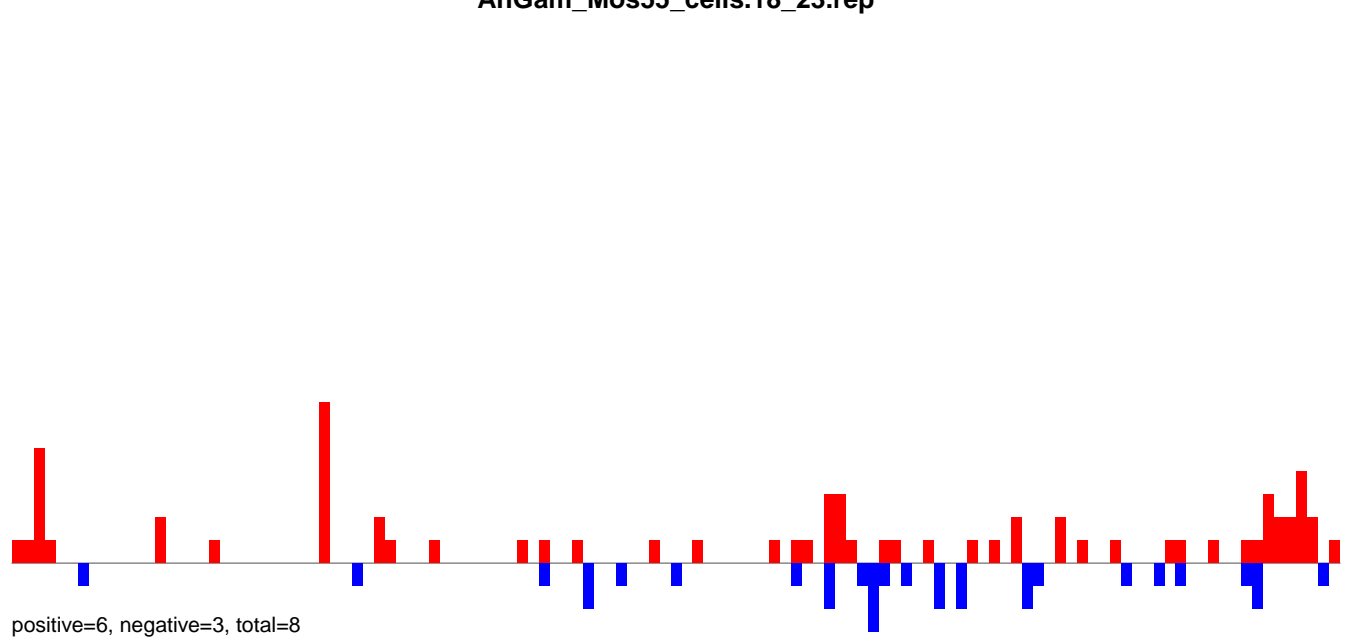
1.5
1.0
0.5
0.0
-0.5

positive=4, negative=15, total=18

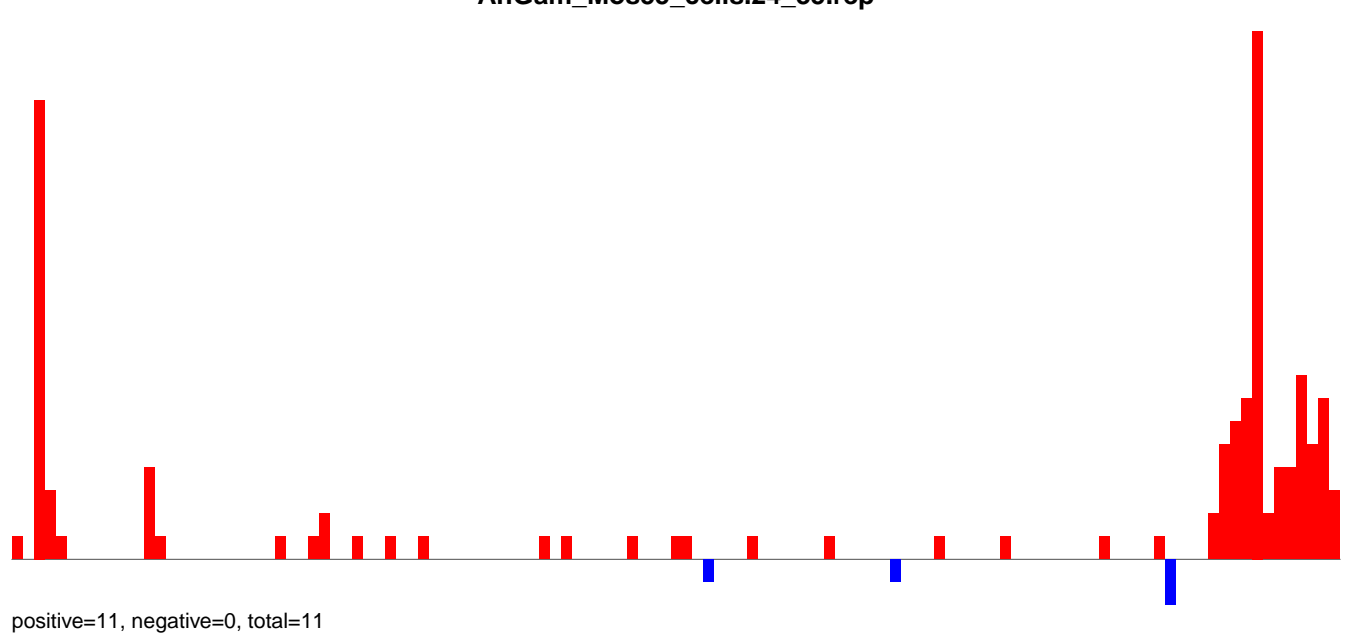
Window size=50, length=6766, TE@TF001975-Pao_Bel_Ele44:1-6766

0 1000 2000 3000 4000 5000 6000 7000

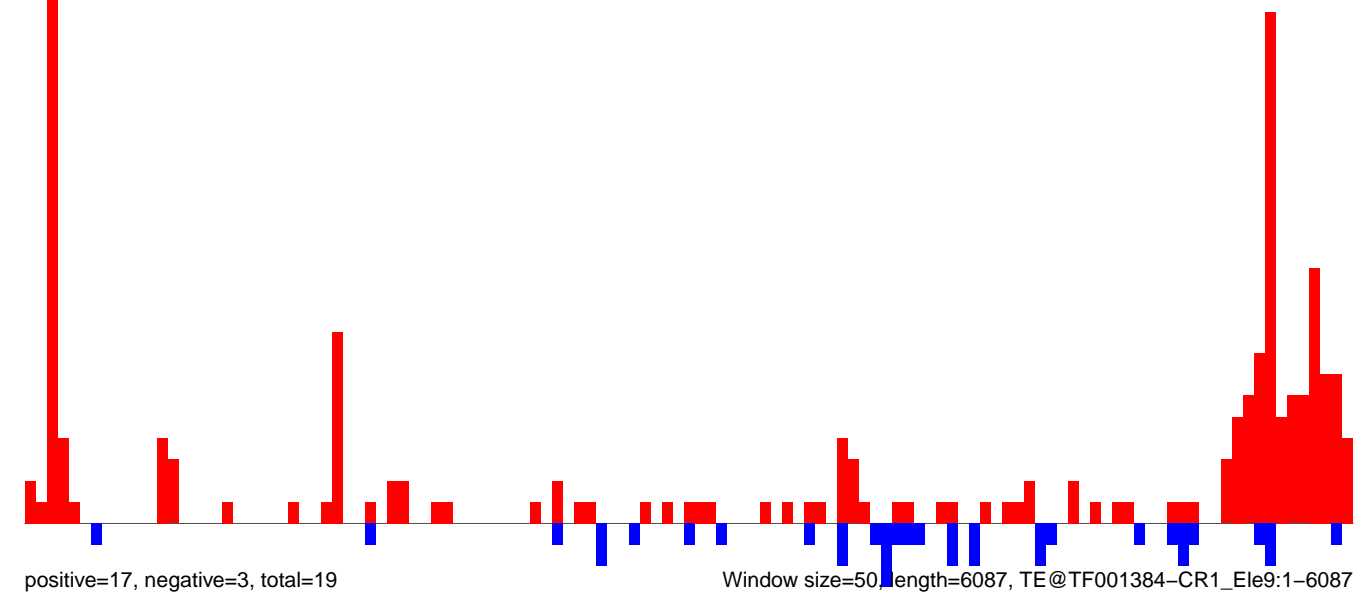
AnGam_Mos55_cells.18_23.rep



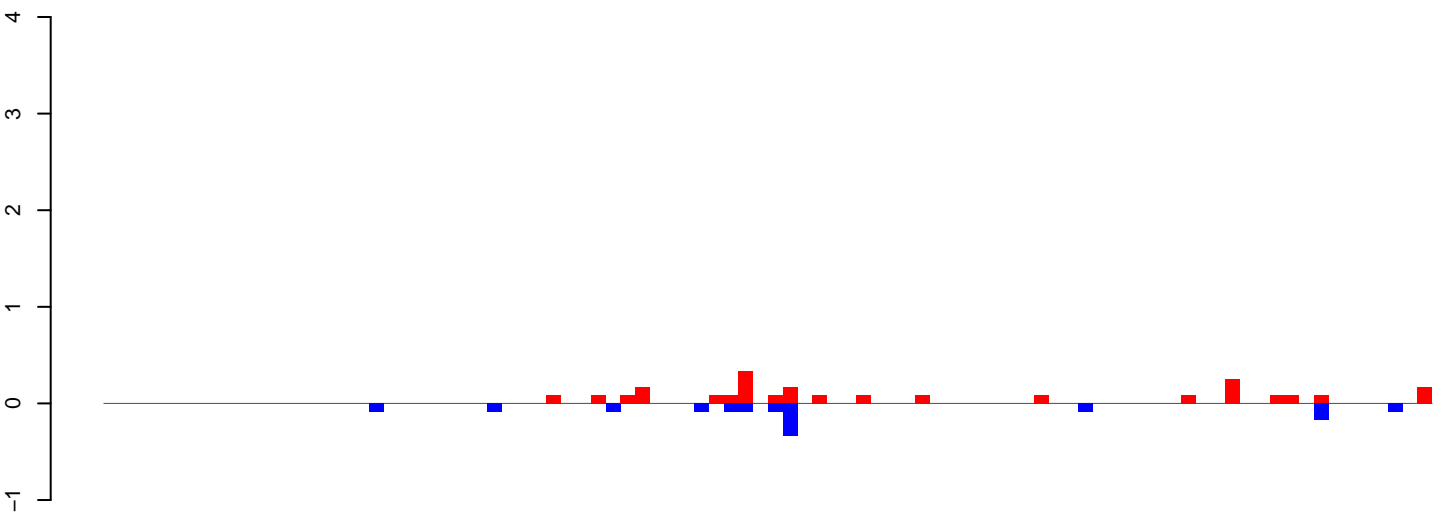
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

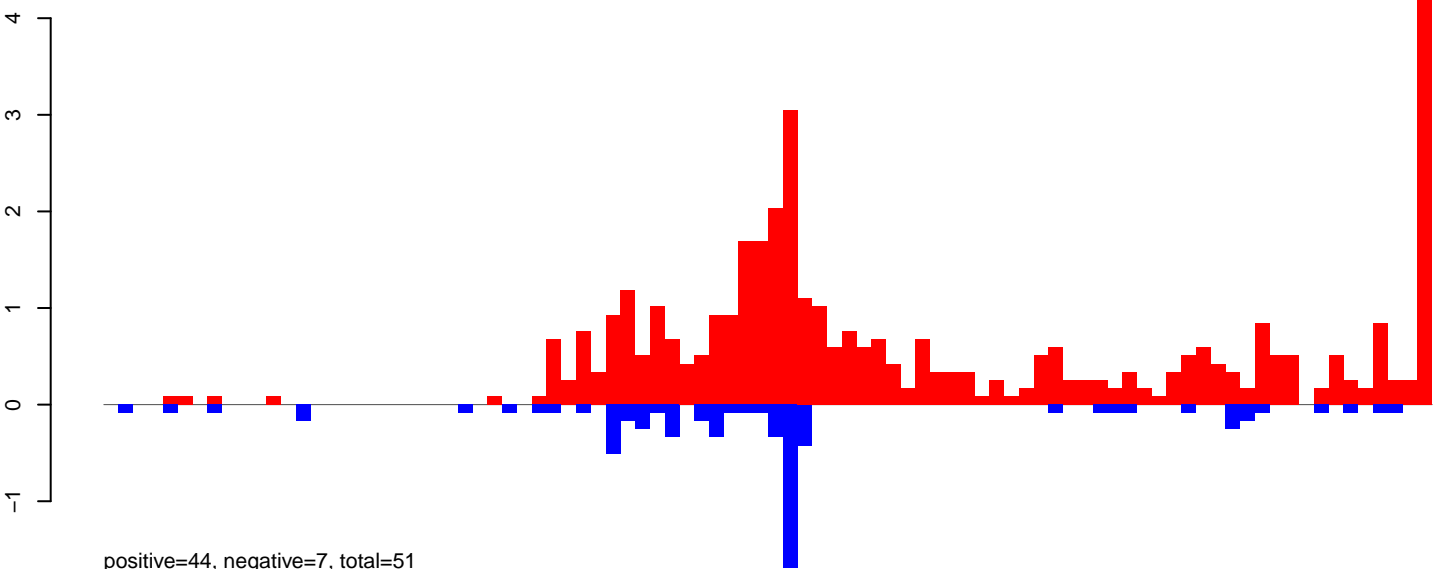


AnGam_Mos55_cells.18_23.rep



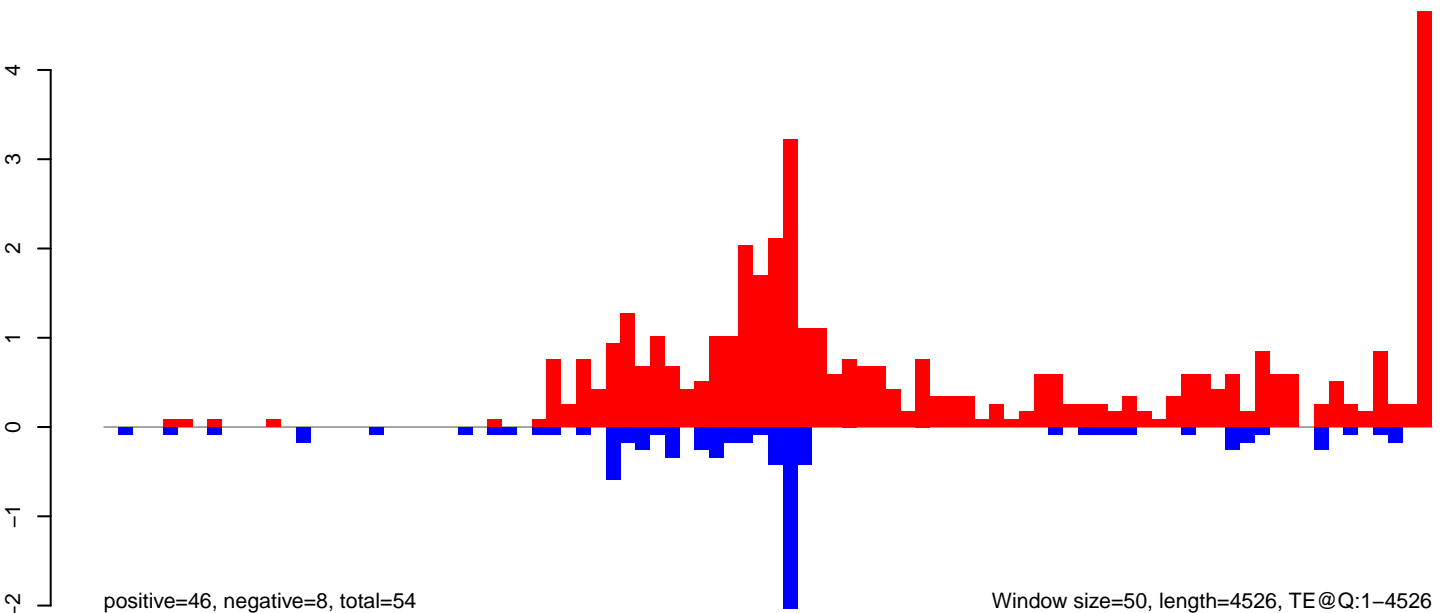
positive=2, negative=1, total=4

AnGam_Mos55_cells.24_35.rep



positive=44, negative=7, total=51

AnGam_Mos55_cells.rep

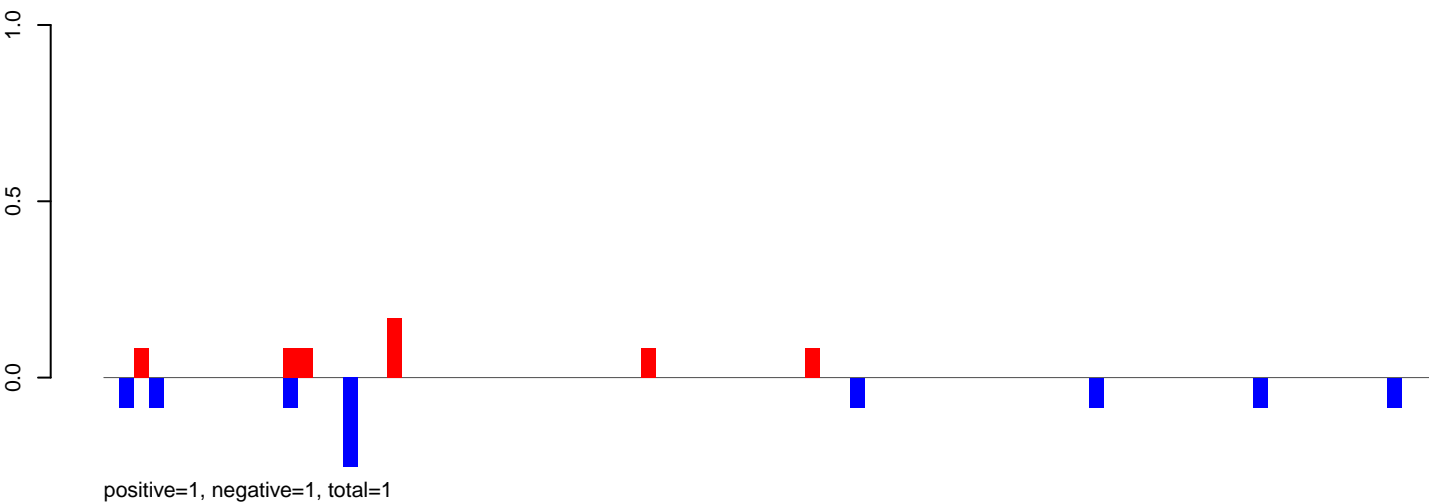


positive=46, negative=8, total=54

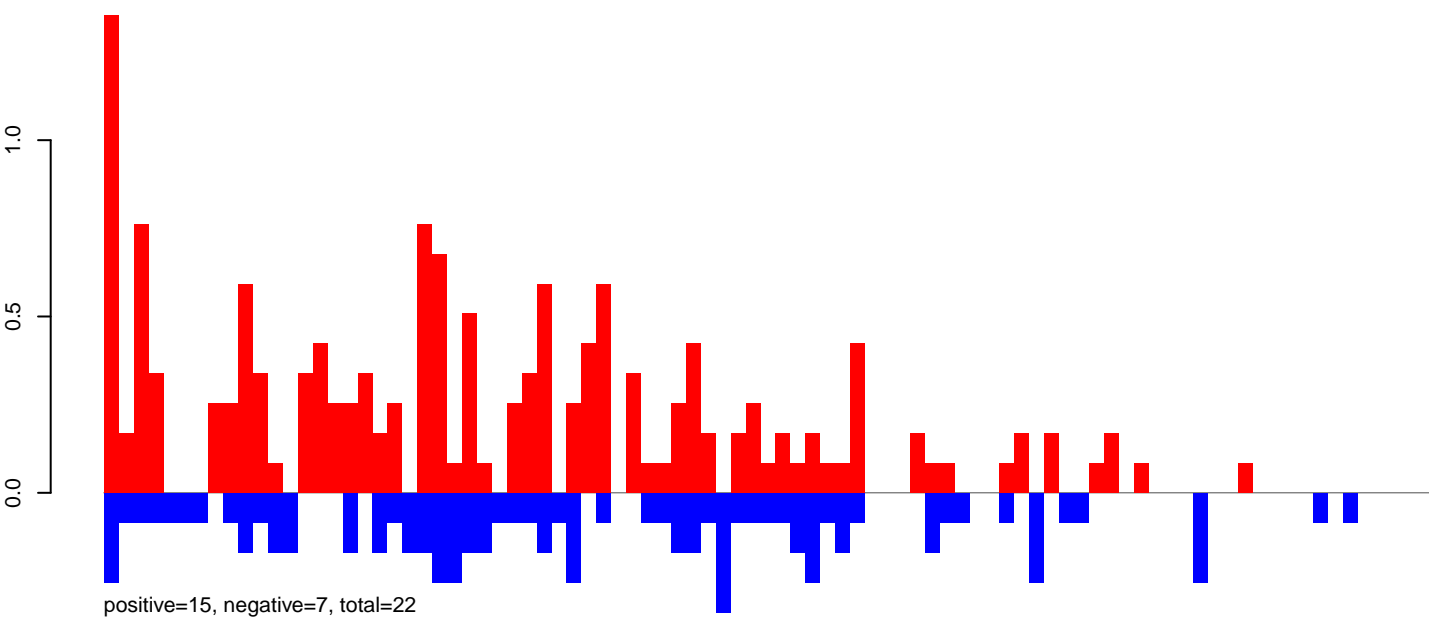
Window size=50, length=4526, TE@Q:1-4526

0 1000 2000 3000 4000

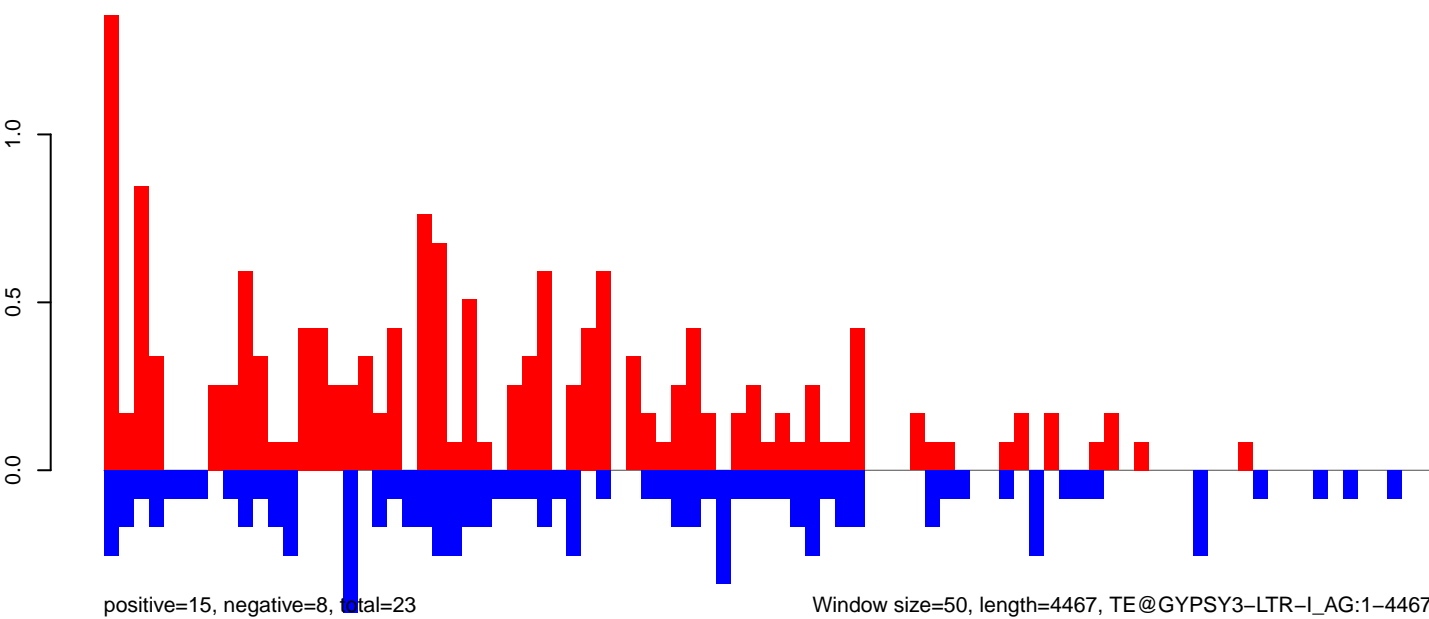
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

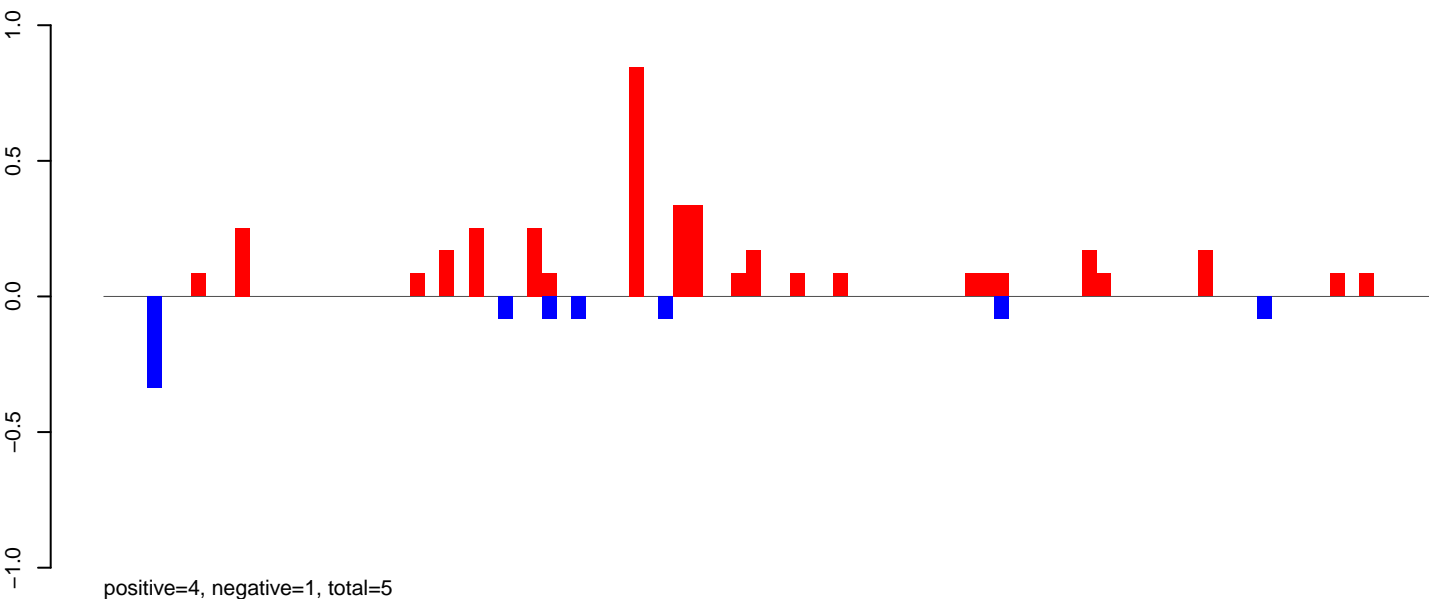


AnGam_Mos55_cells.rep

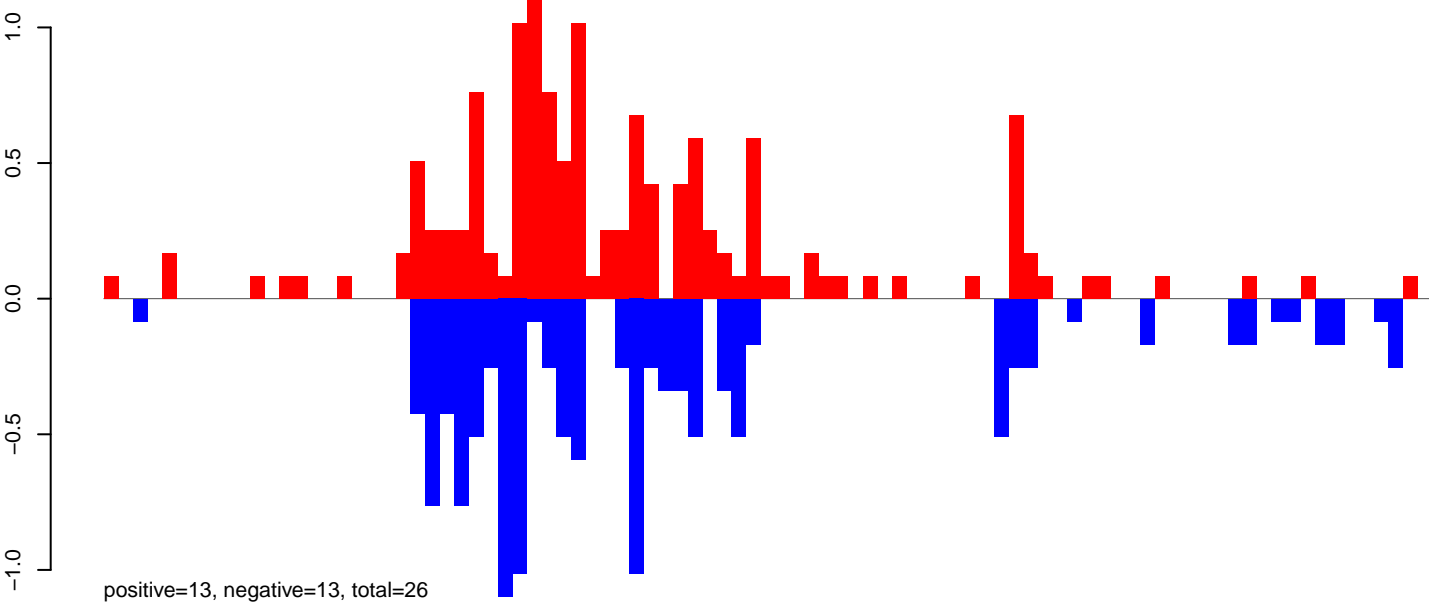


Window size=50, length=4467, TE@GYPSY3-LTR-L_AG:1-4467

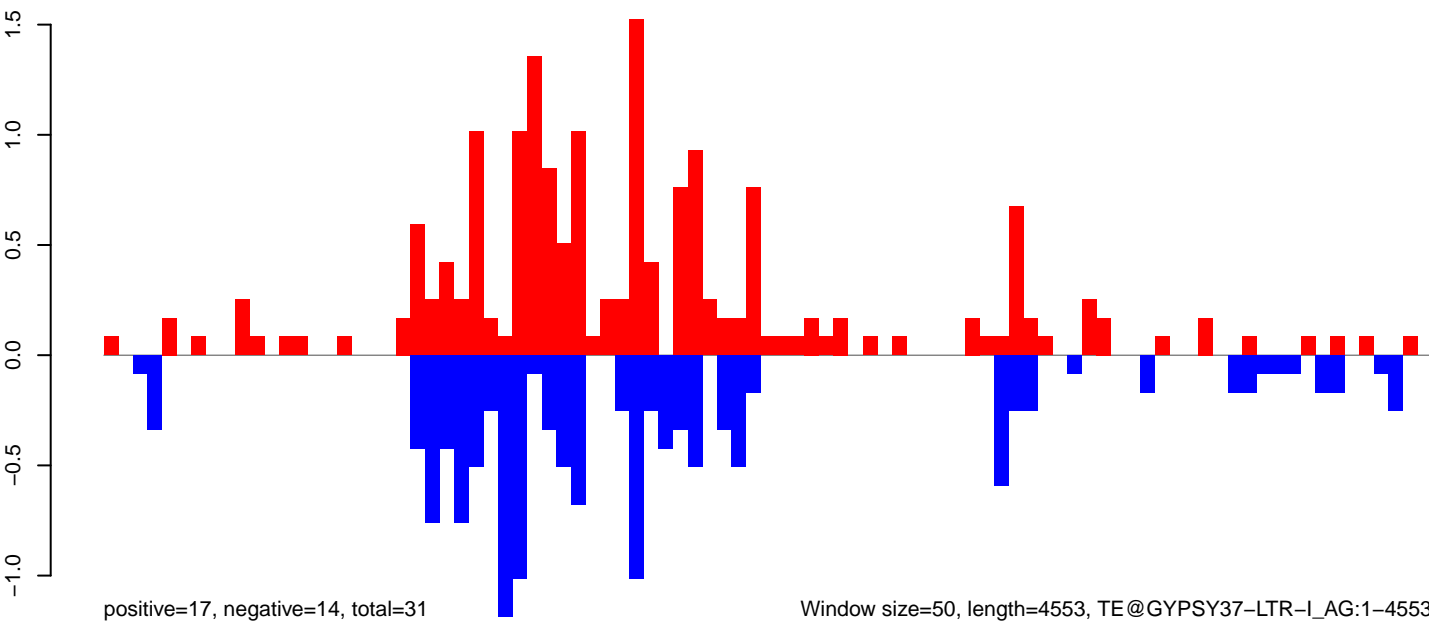
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



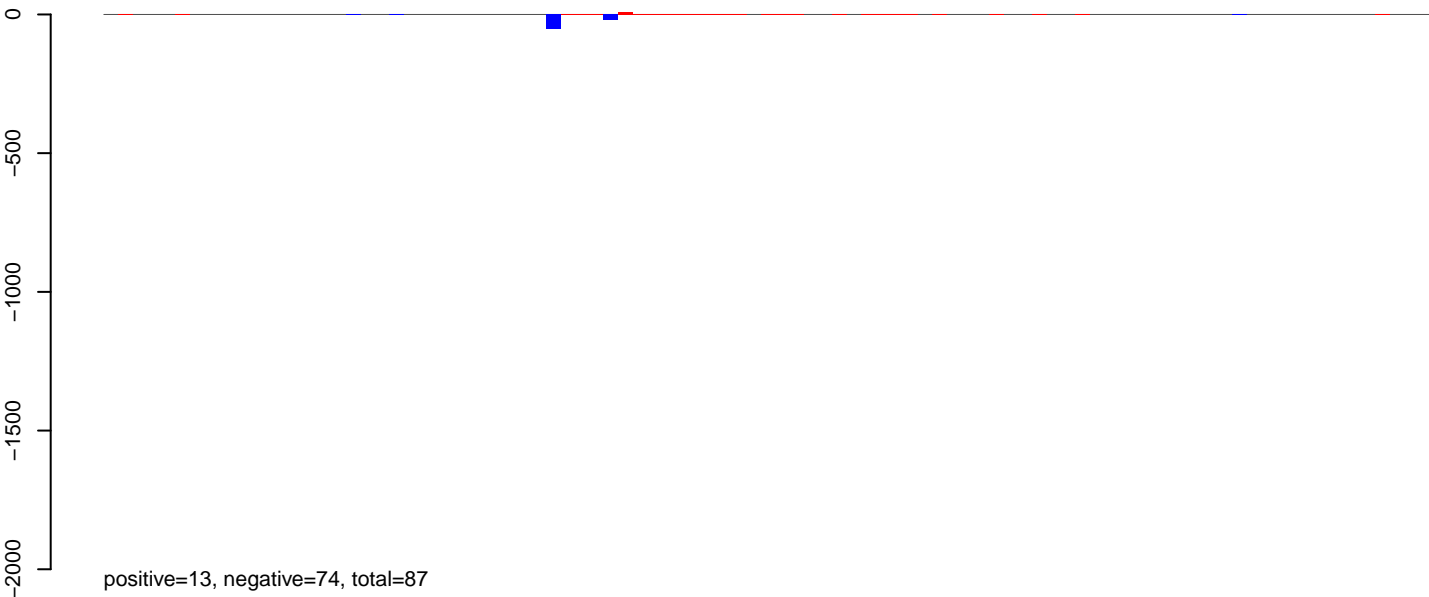
AnGam_Mos55_cells.rep



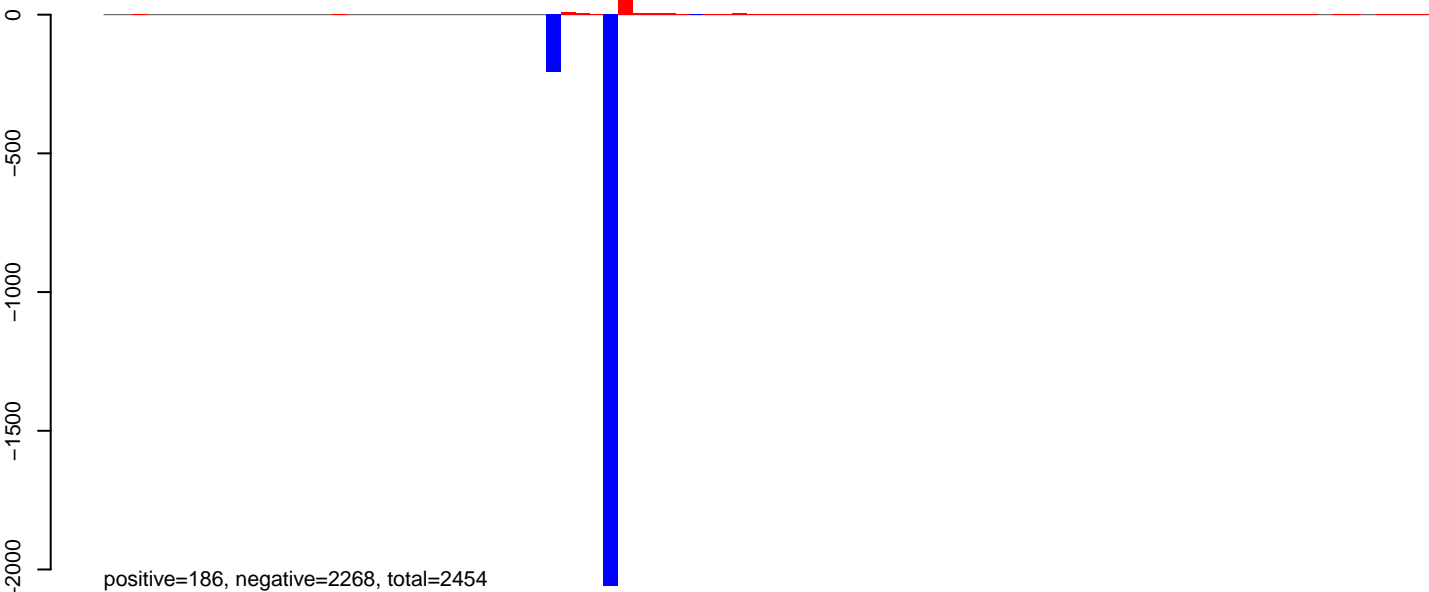
Window size=50, length=4553, TE@GYPSY37-LTR-I_AG:1-4553

0 1000 2000 3000 4000

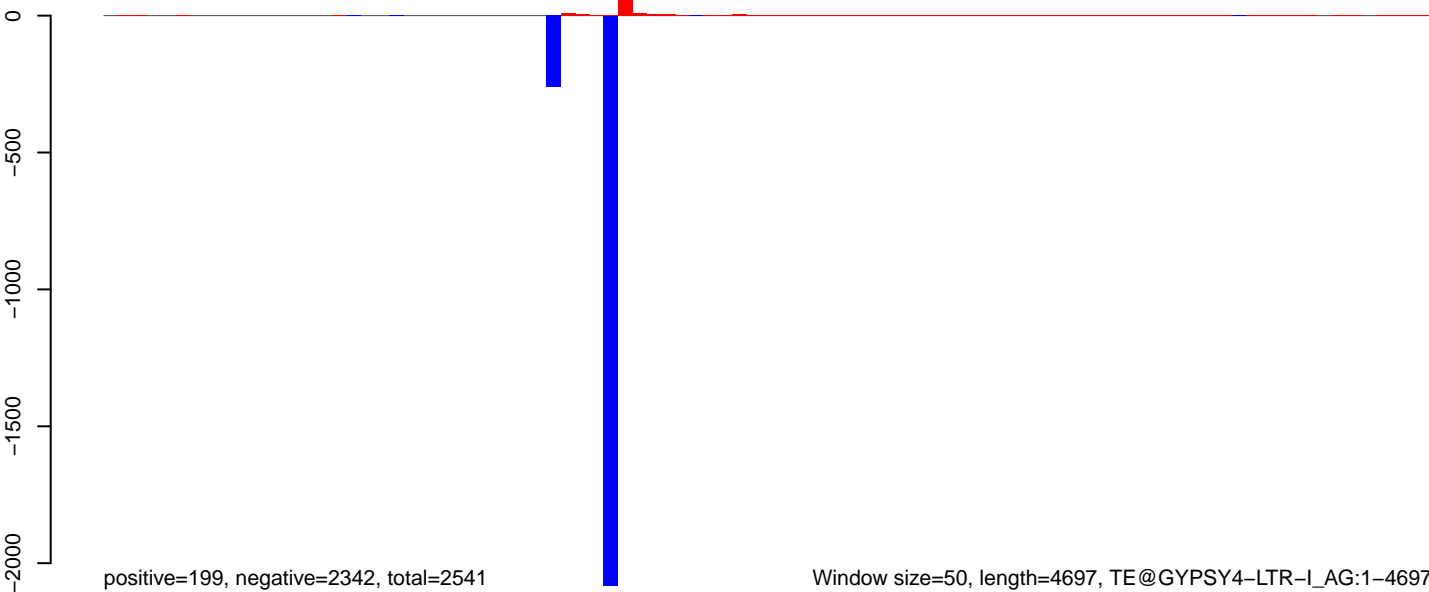
AnGam_Mos55_cells.18_23.rep



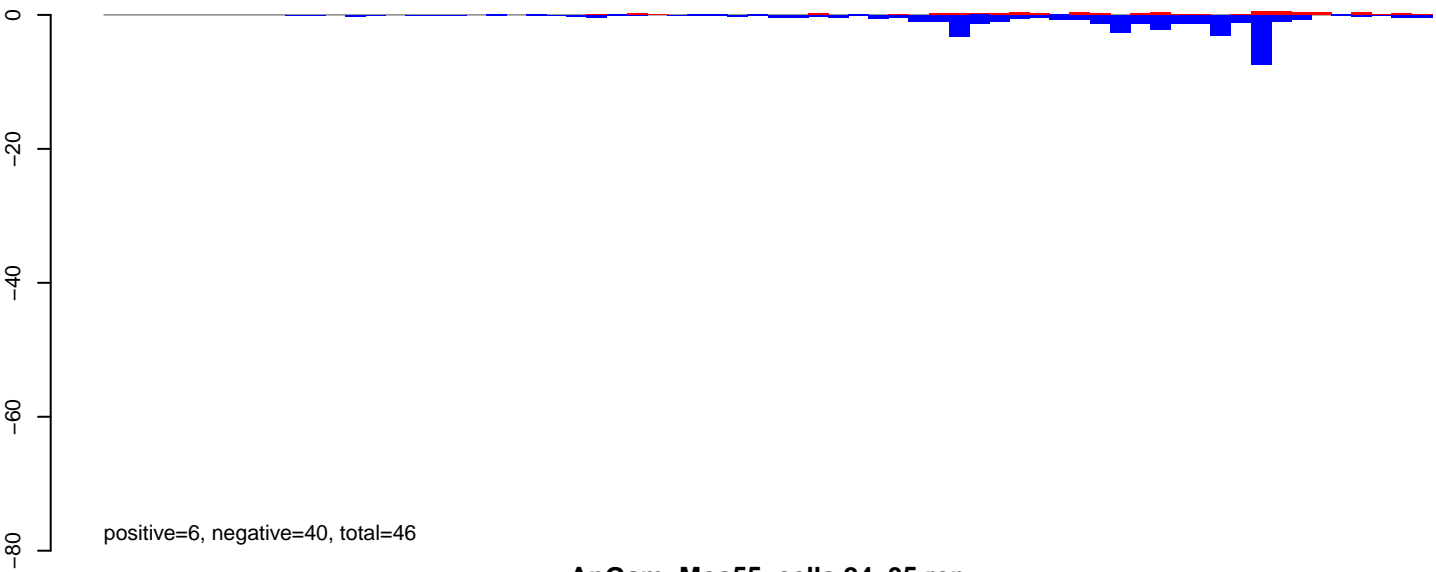
AnGam_Mos55_cells.24_35.rep



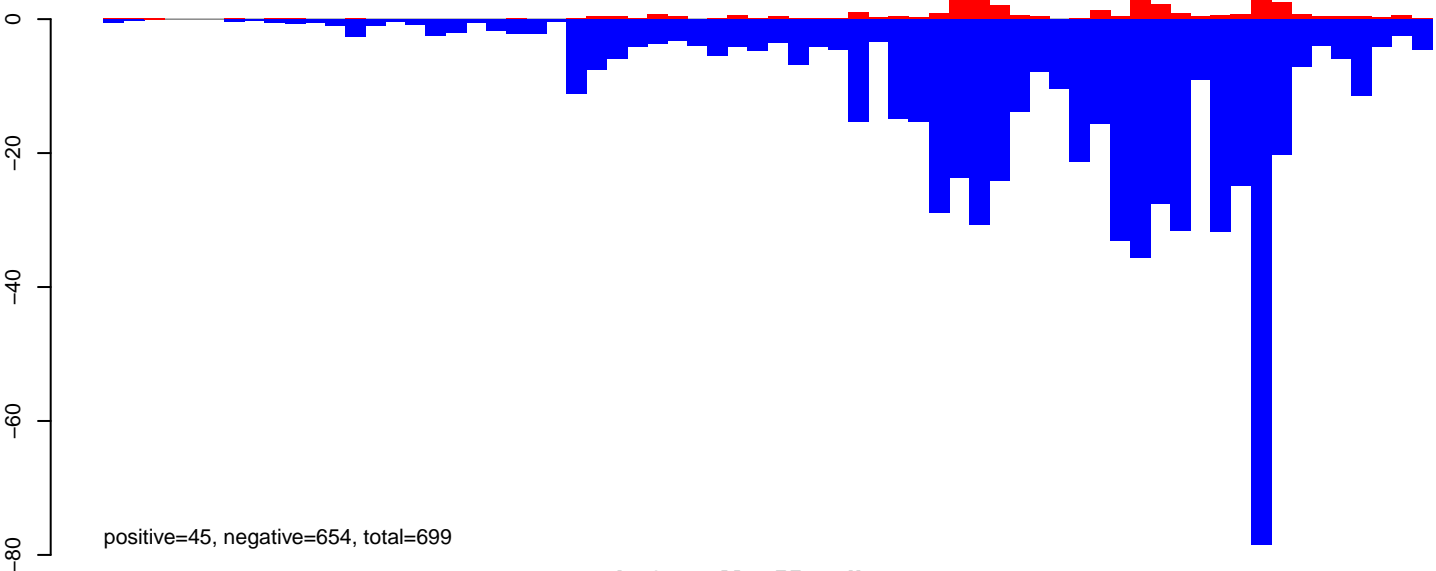
AnGam_Mos55_cells.rep



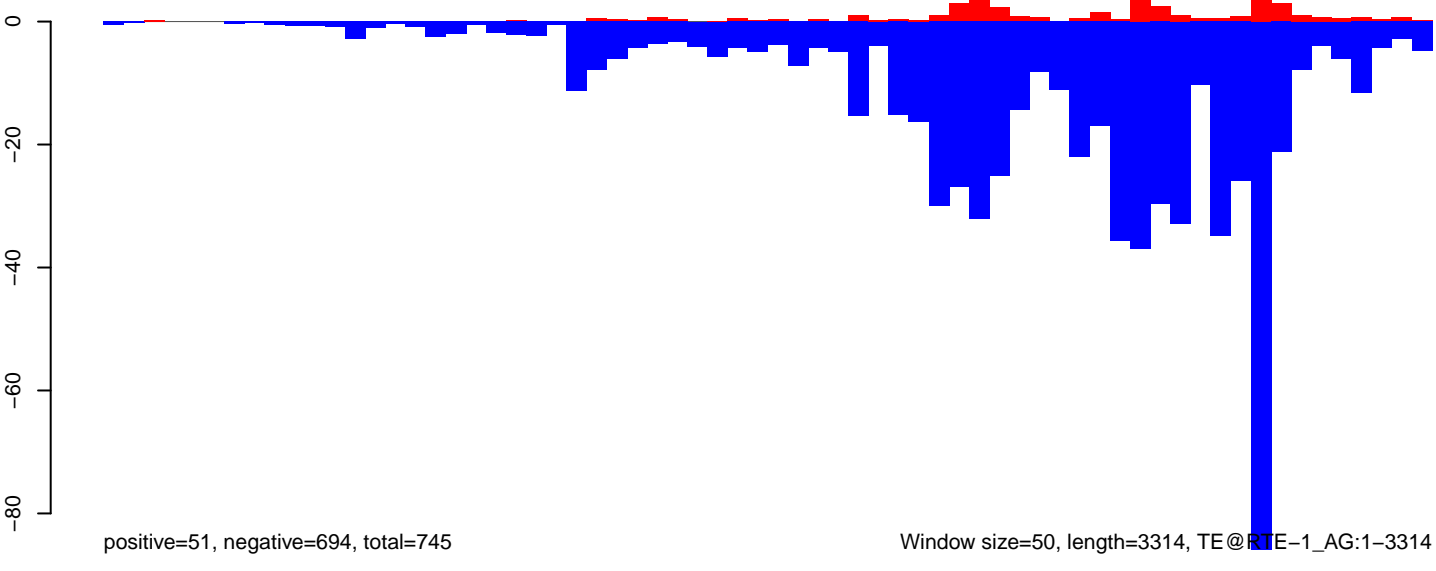
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



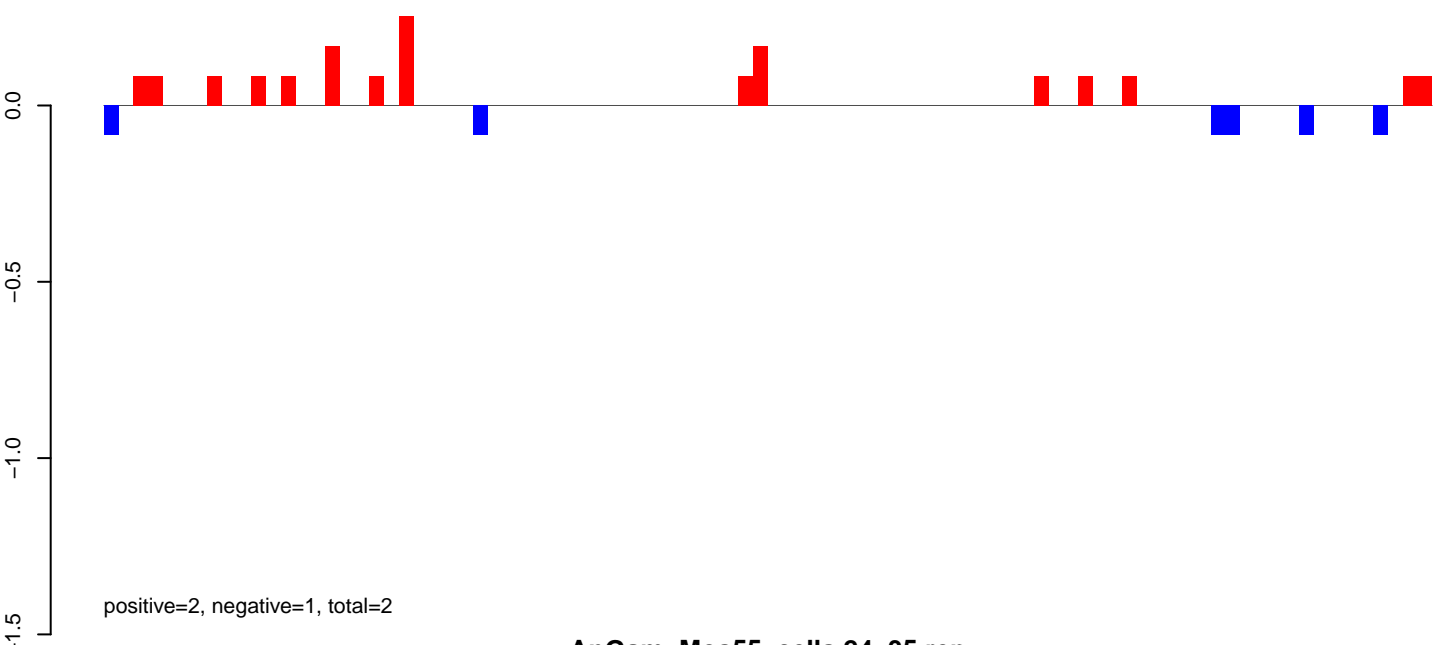
AnGam_Mos55_cells.rep



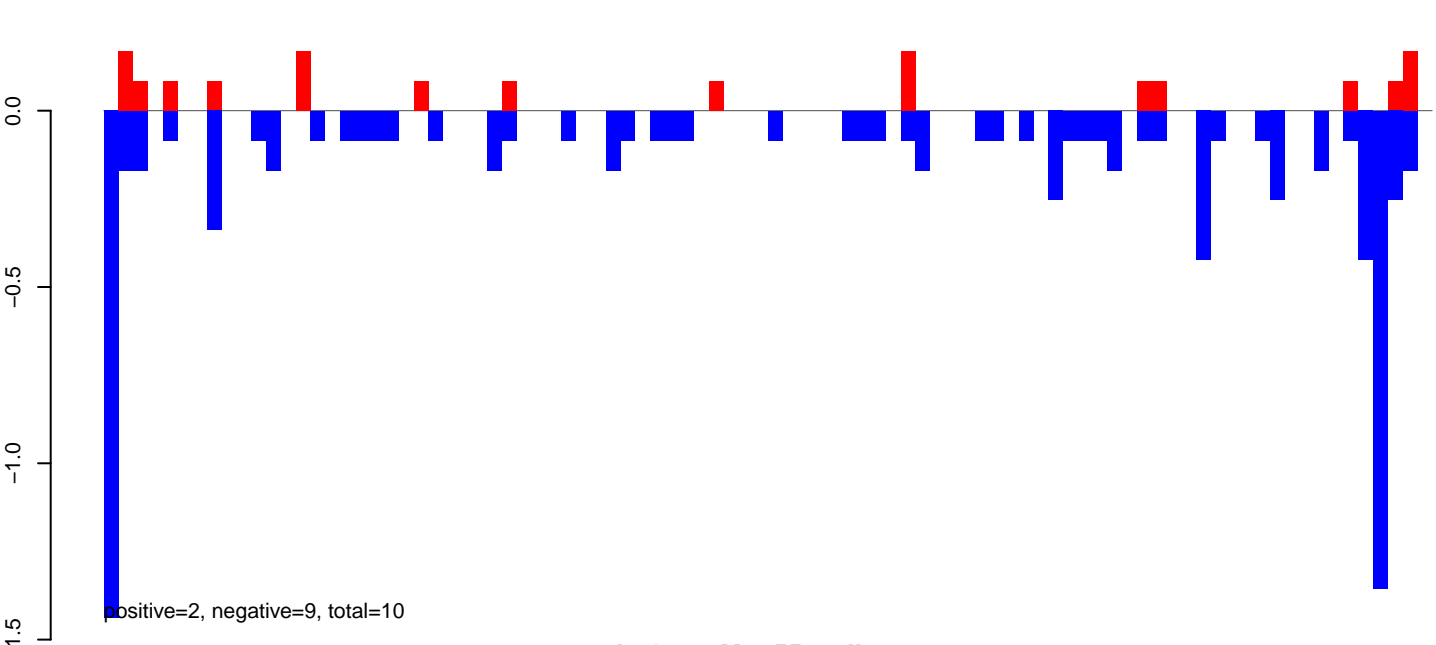
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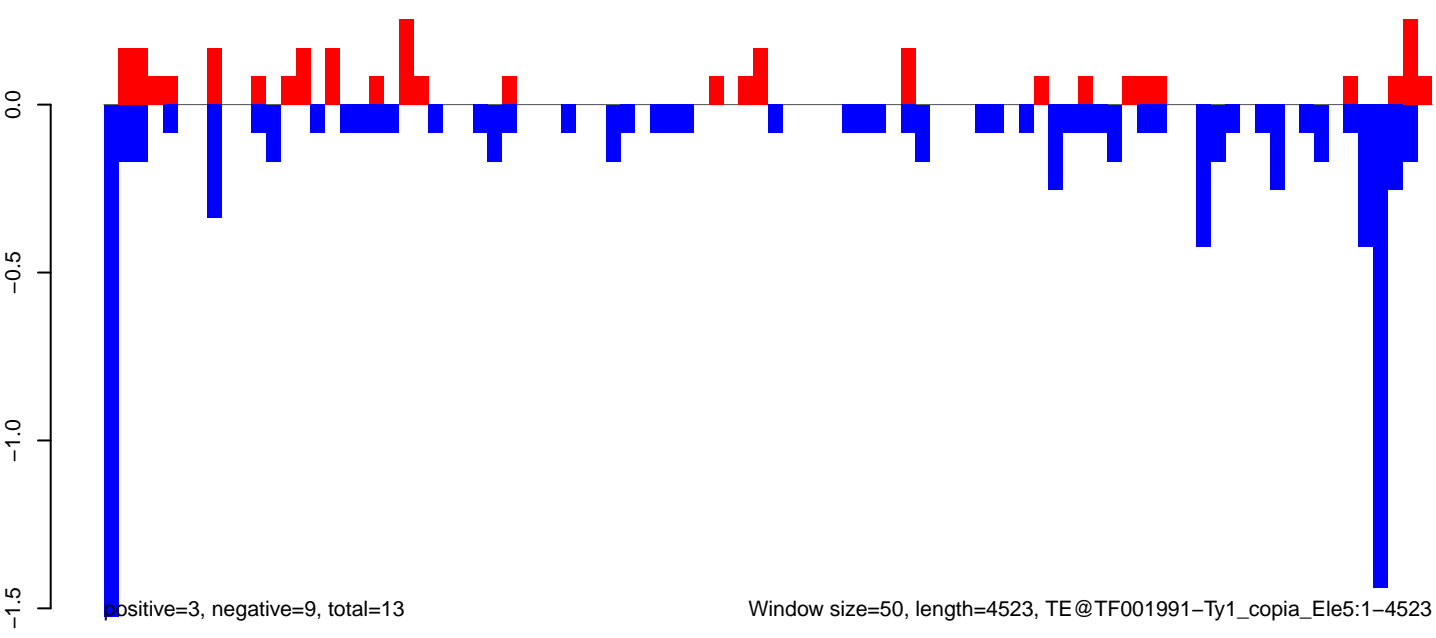
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

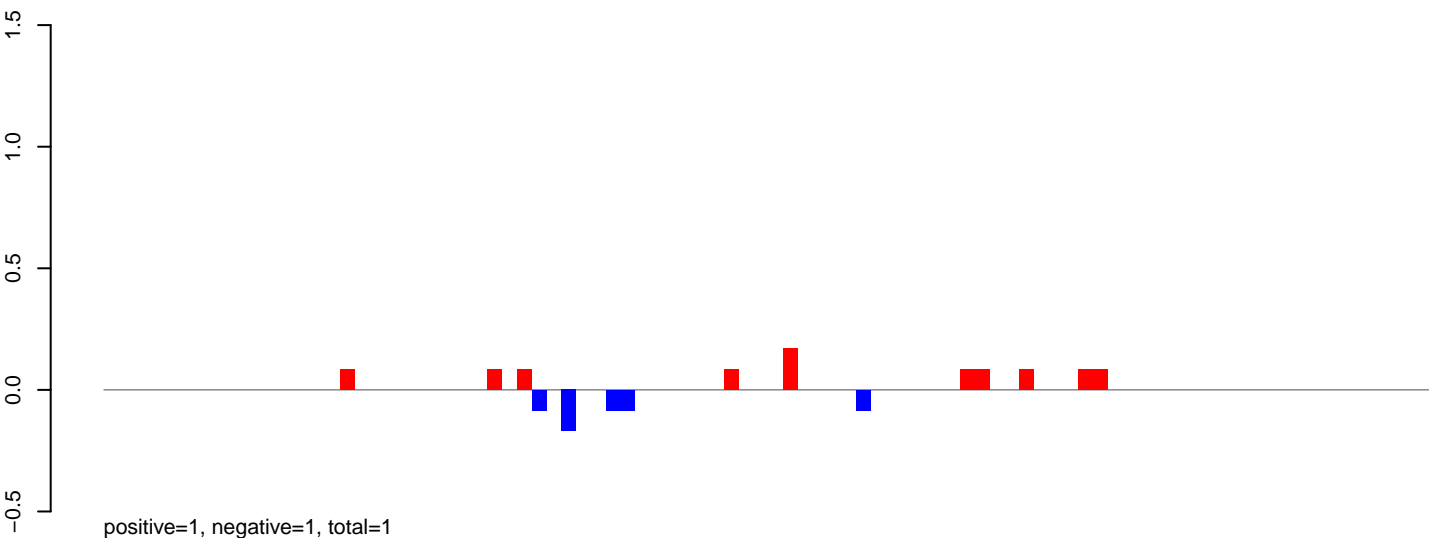


AnGam_Mos55_cells.rep

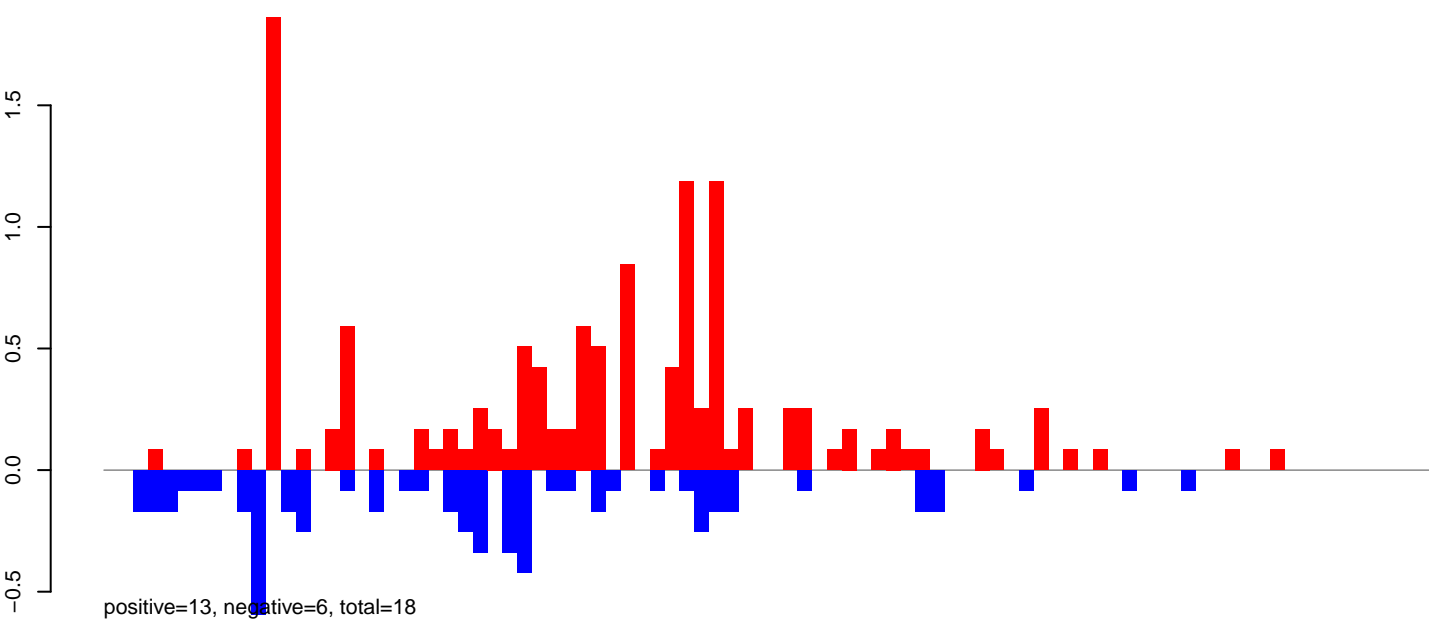


Window size=50, length=4523, TE@TF001991-Ty1_copia_Ele5:1-4523

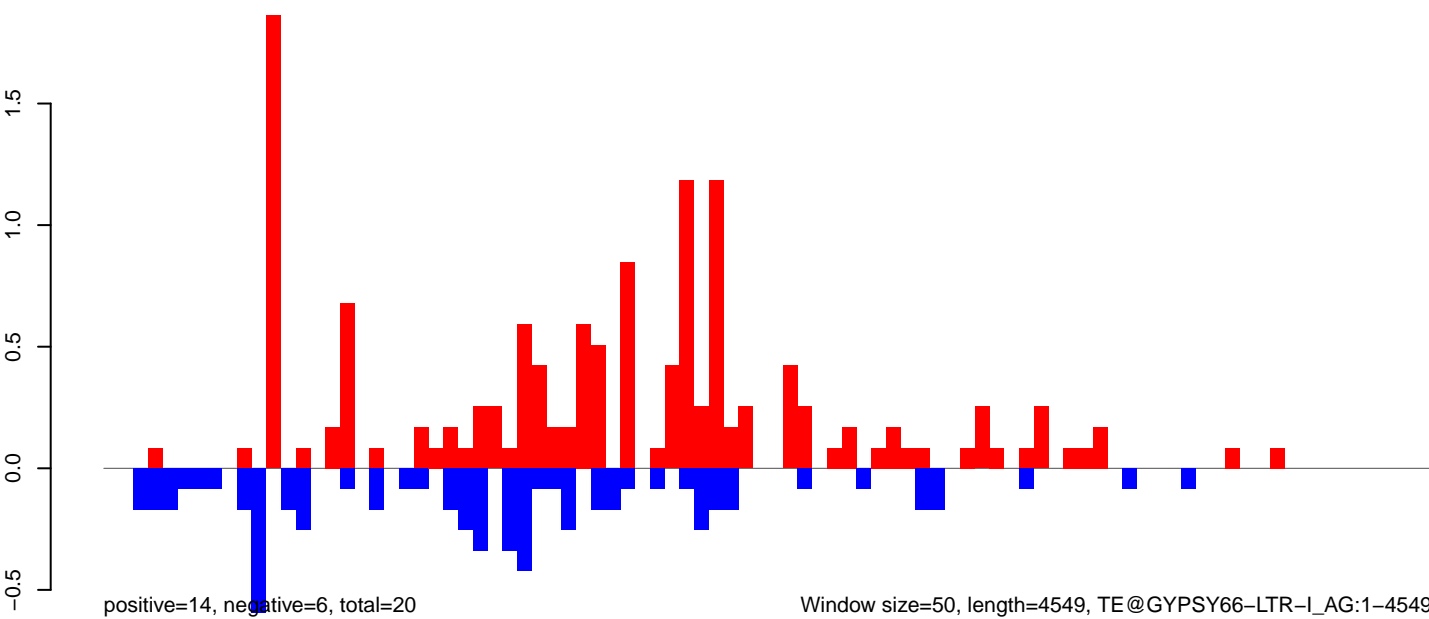
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



Window size=50, length=4549, TE@GYPSY66-LTR-I_AG:1-4549

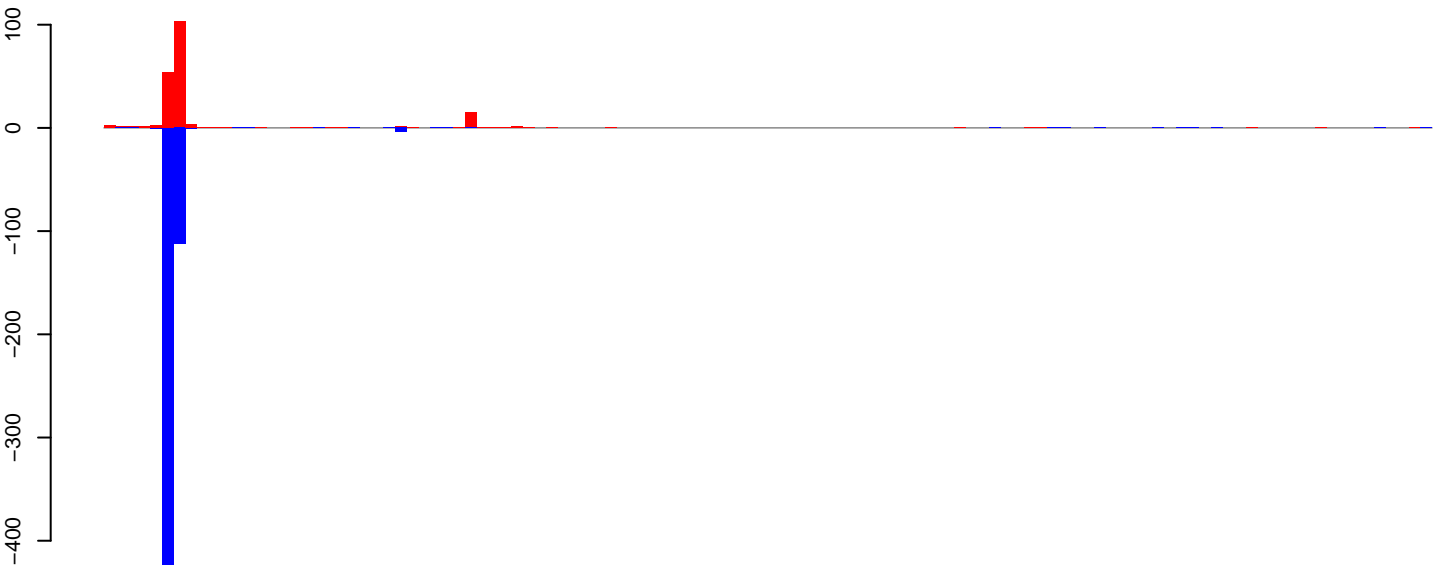
0 1000 2000 3000 4000

AnGam_Mos55_cells.18_23.rep



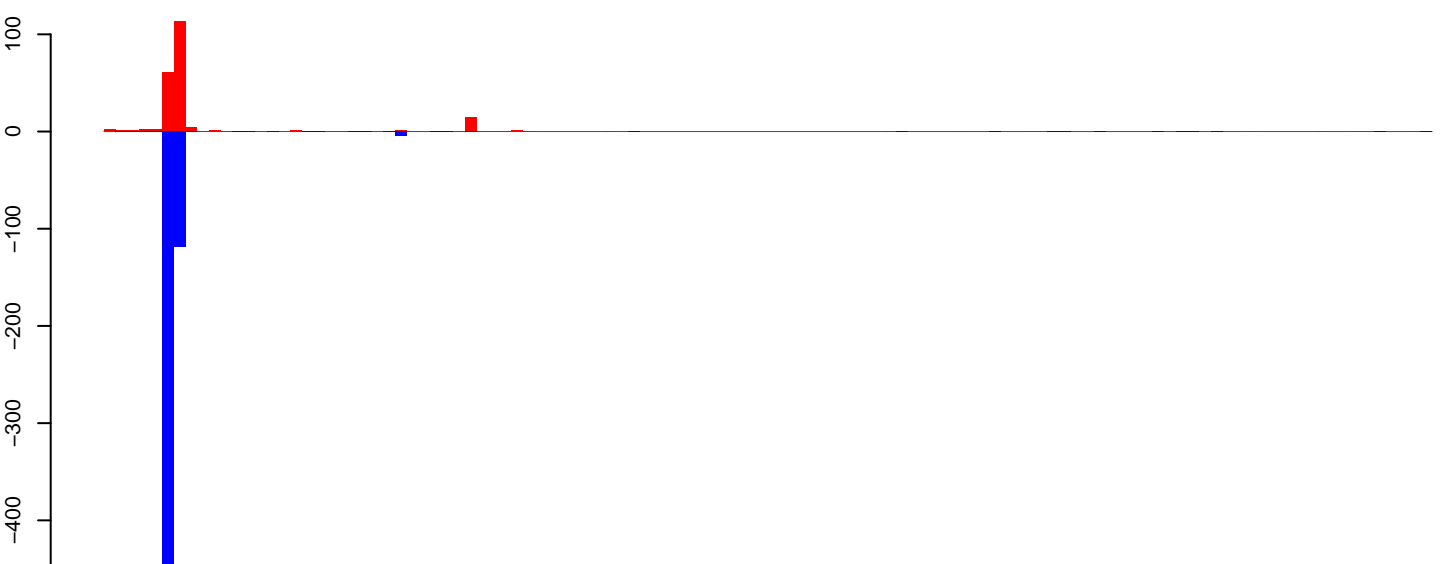
positive=25, negative=35, total=60

AnGam_Mos55_cells.24_35.rep



positive=193, negative=595, total=787

AnGam_Mos55_cells.rep

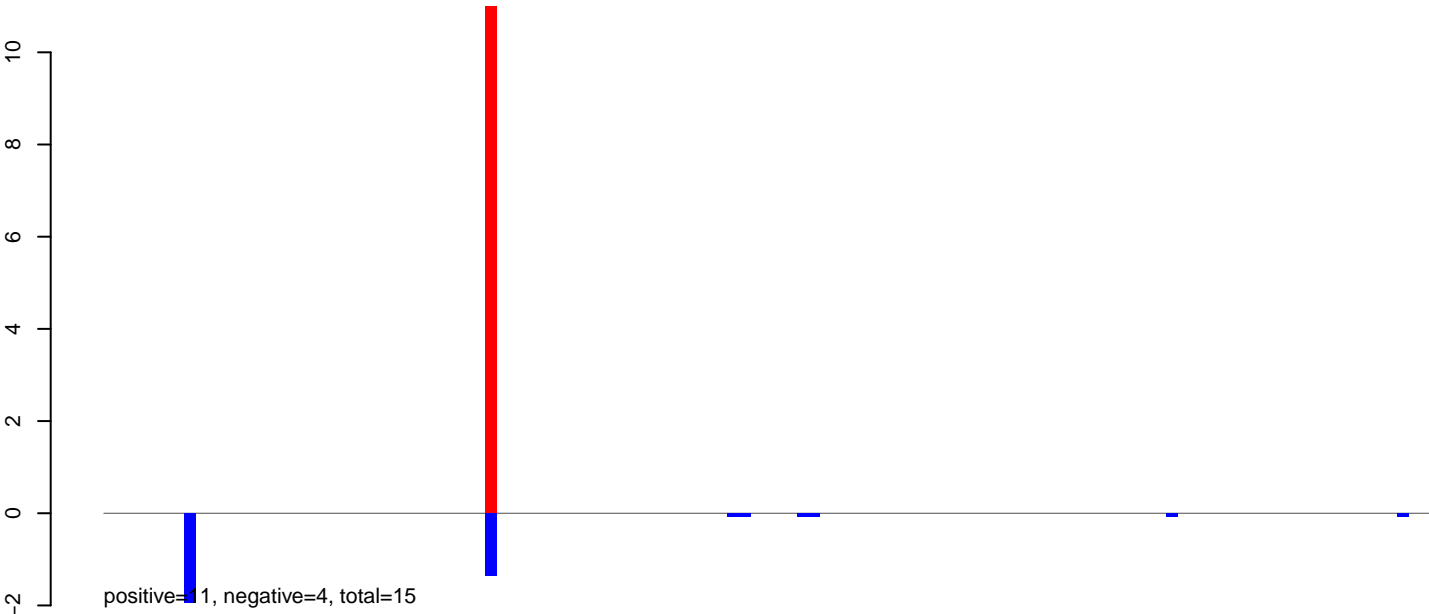


positive=217, negative=629, total=847

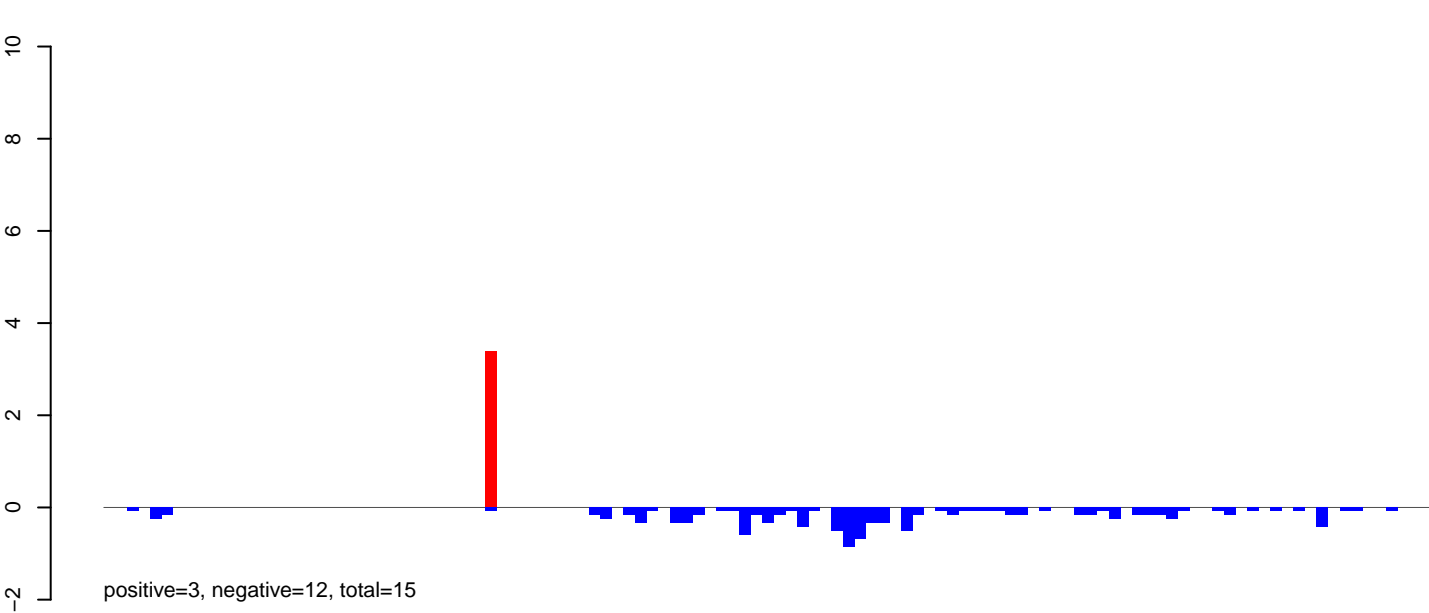
Window size=50, length=5734, TE@GYPSY52-LTR-I_AG:1-5734

0 1000 2000 3000 4000 5000

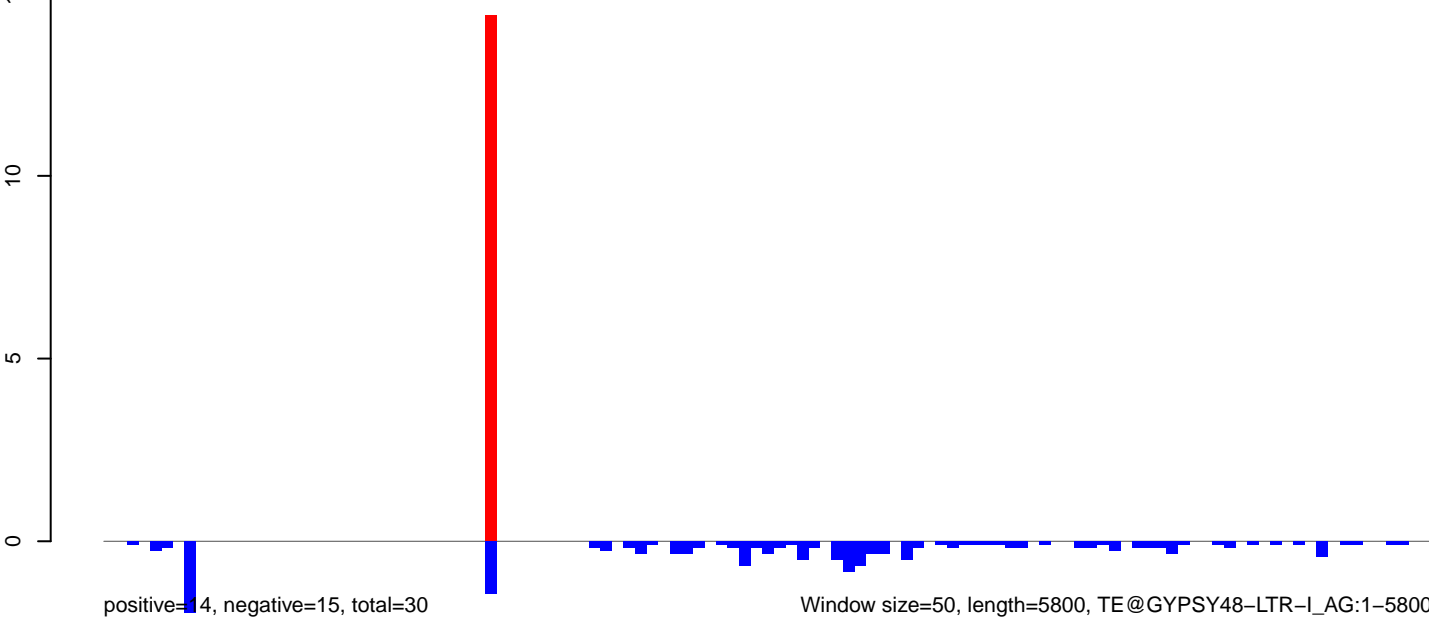
AnGam_Mos55_cells.18_23.rep



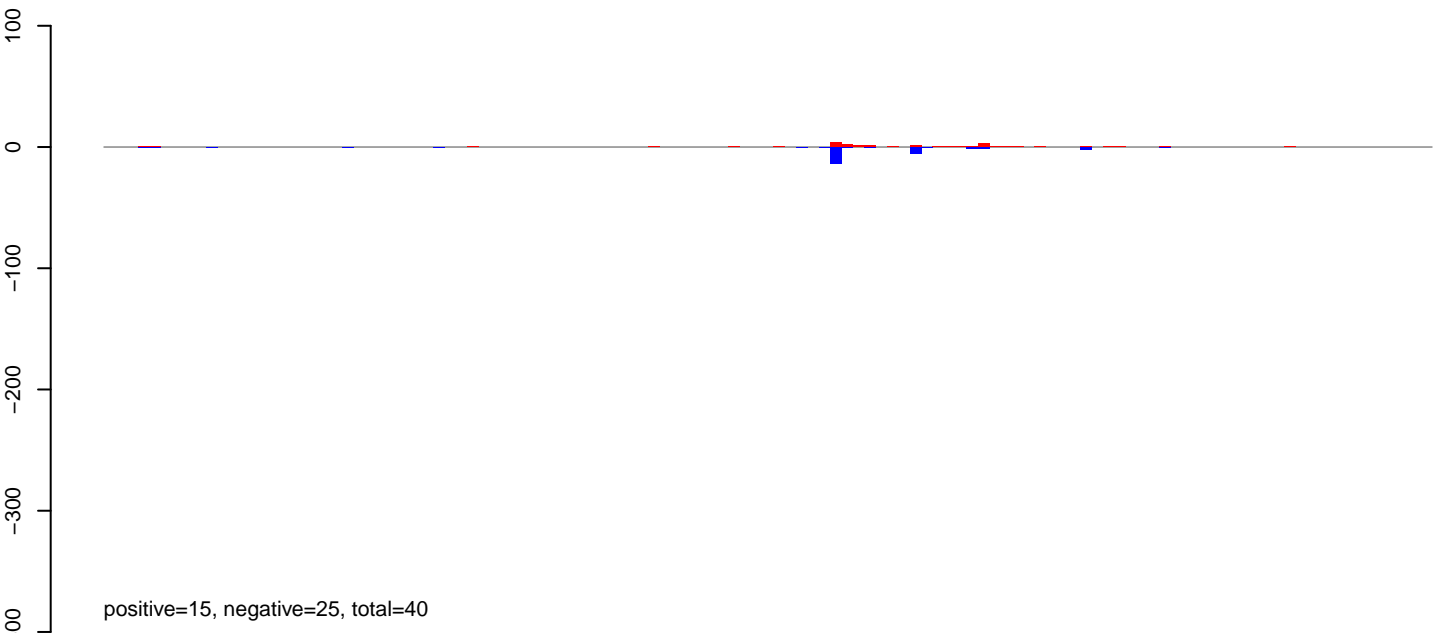
AnGam_Mos55_cells.24_35.rep



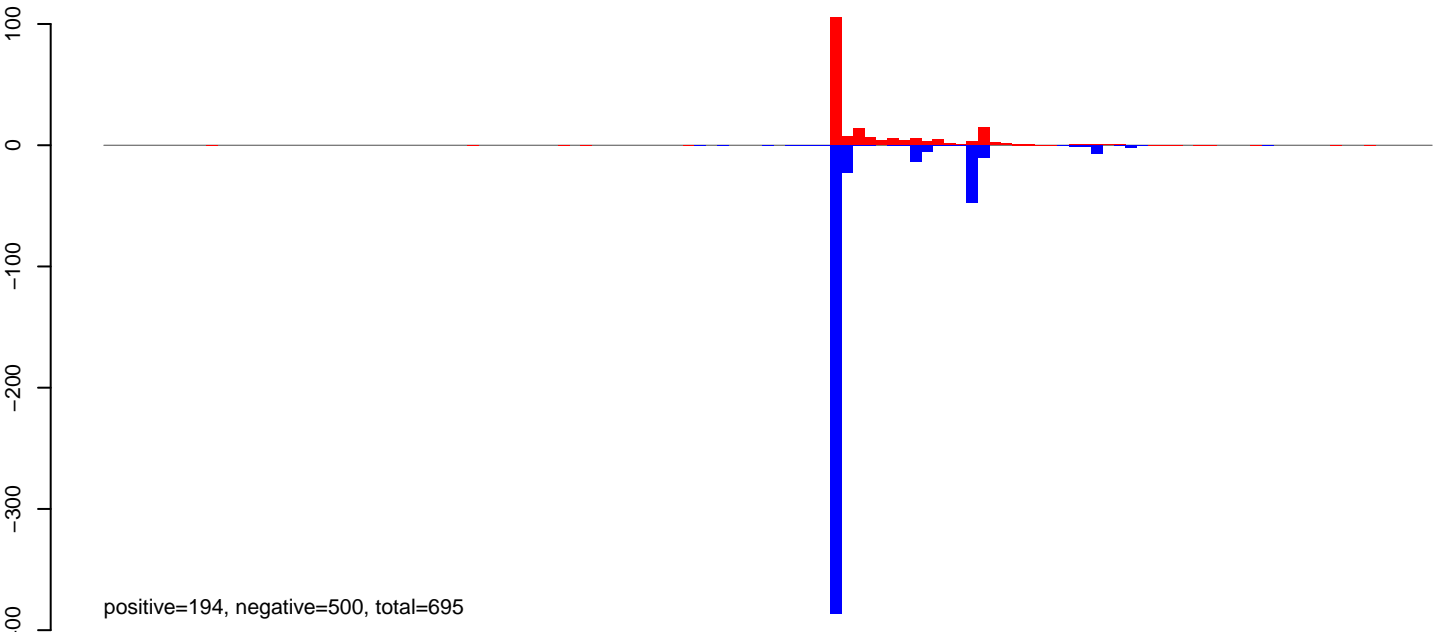
AnGam_Mos55_cells.rep



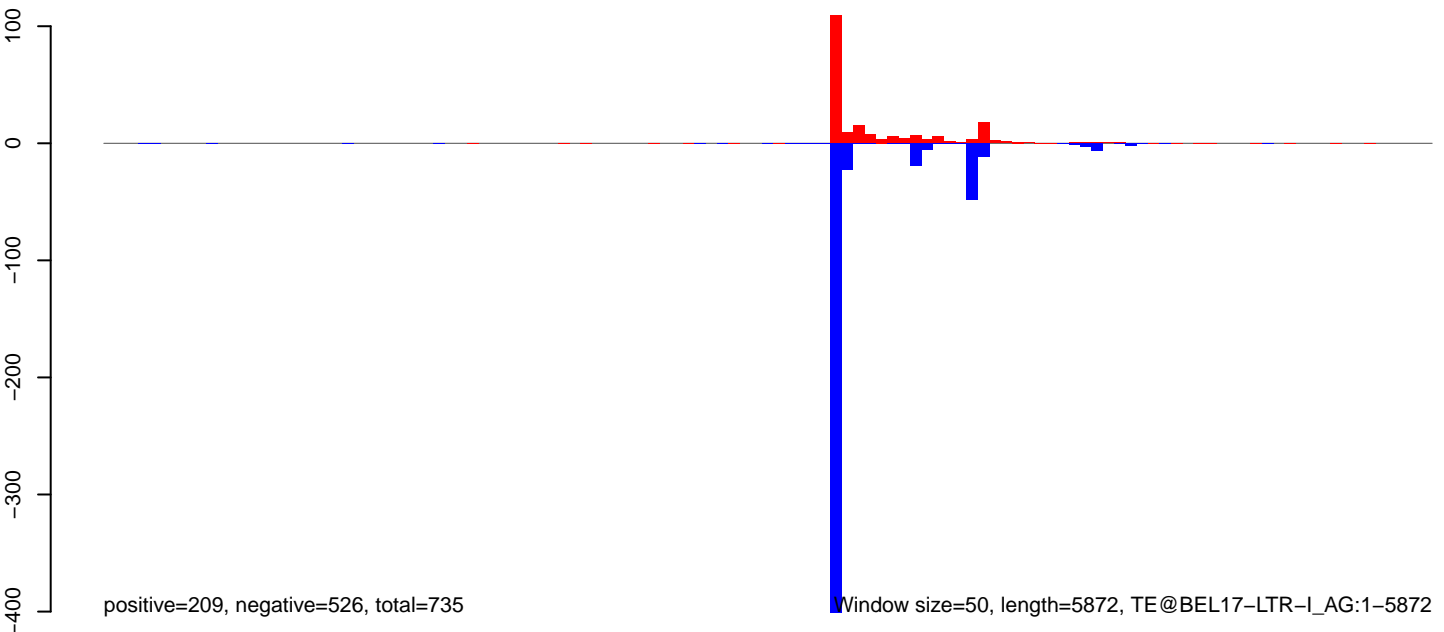
AnGam_Mos55_cells.18_23.rep



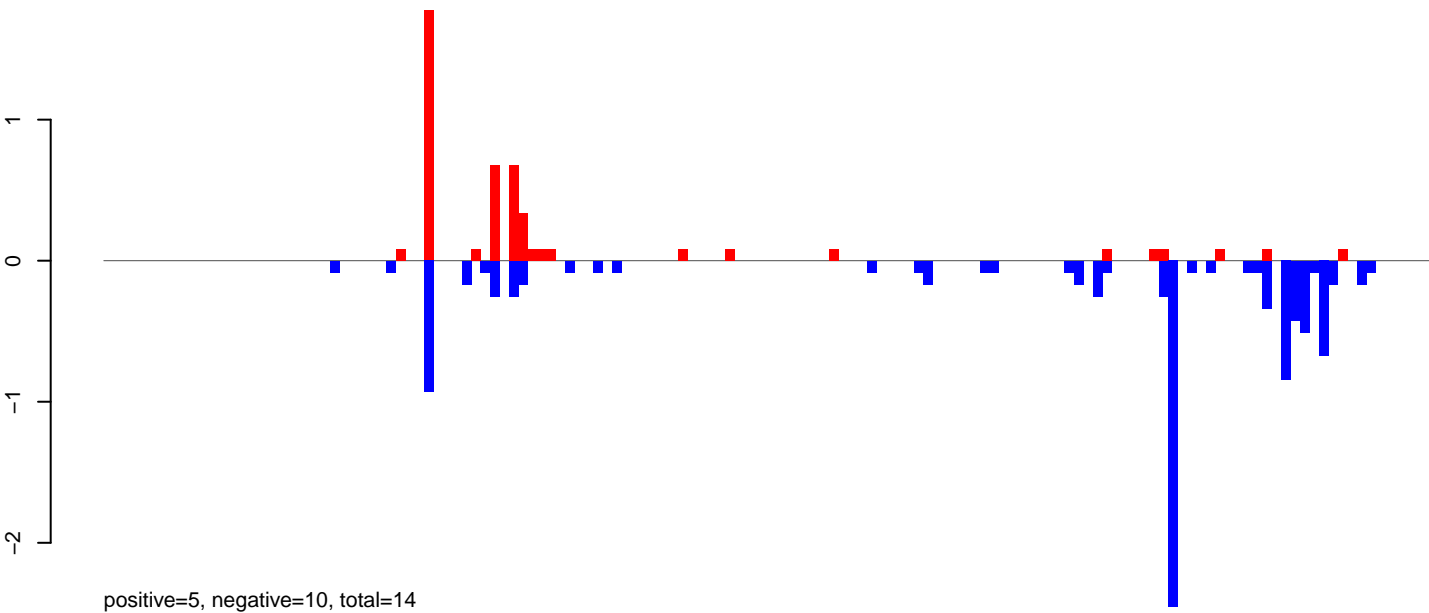
AnGam_Mos55_cells.24_35.rep



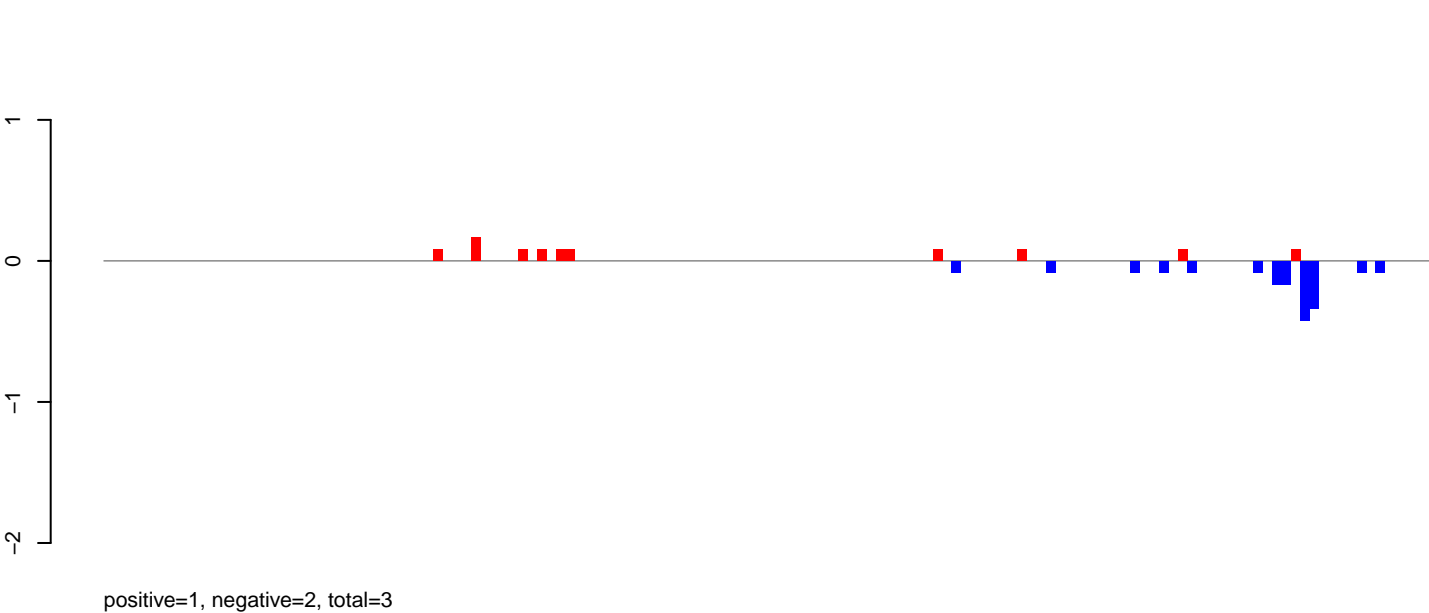
AnGam_Mos55_cells.rep



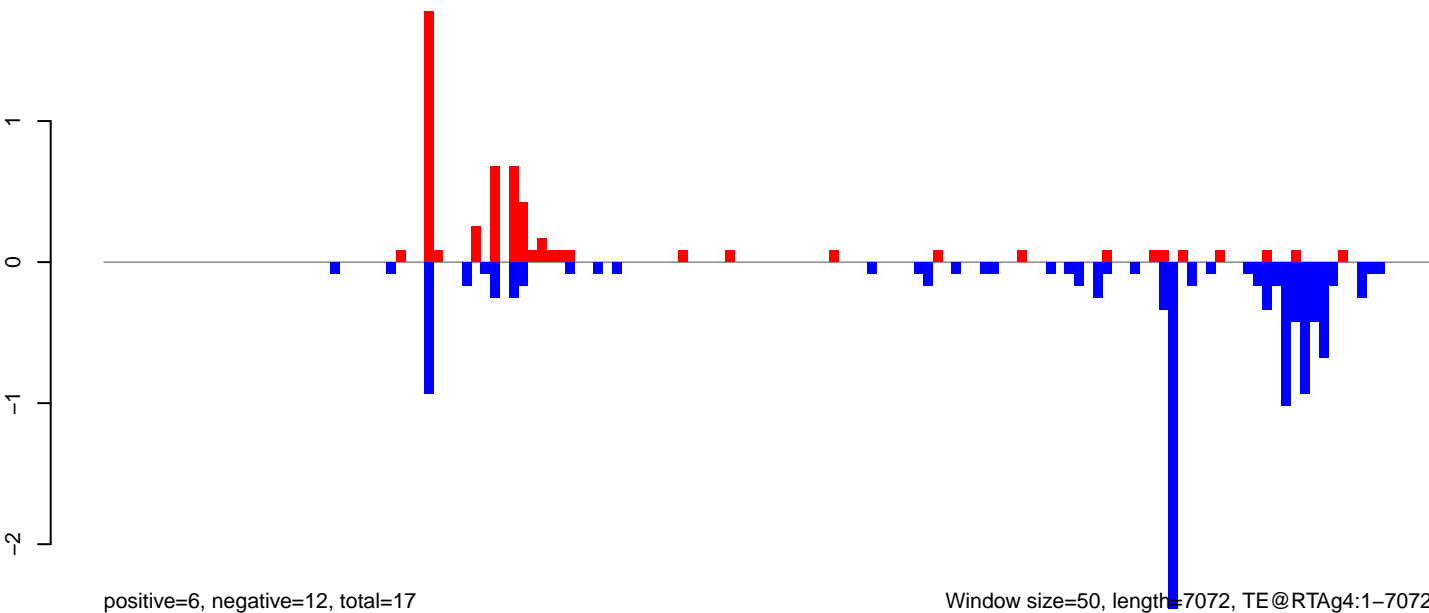
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



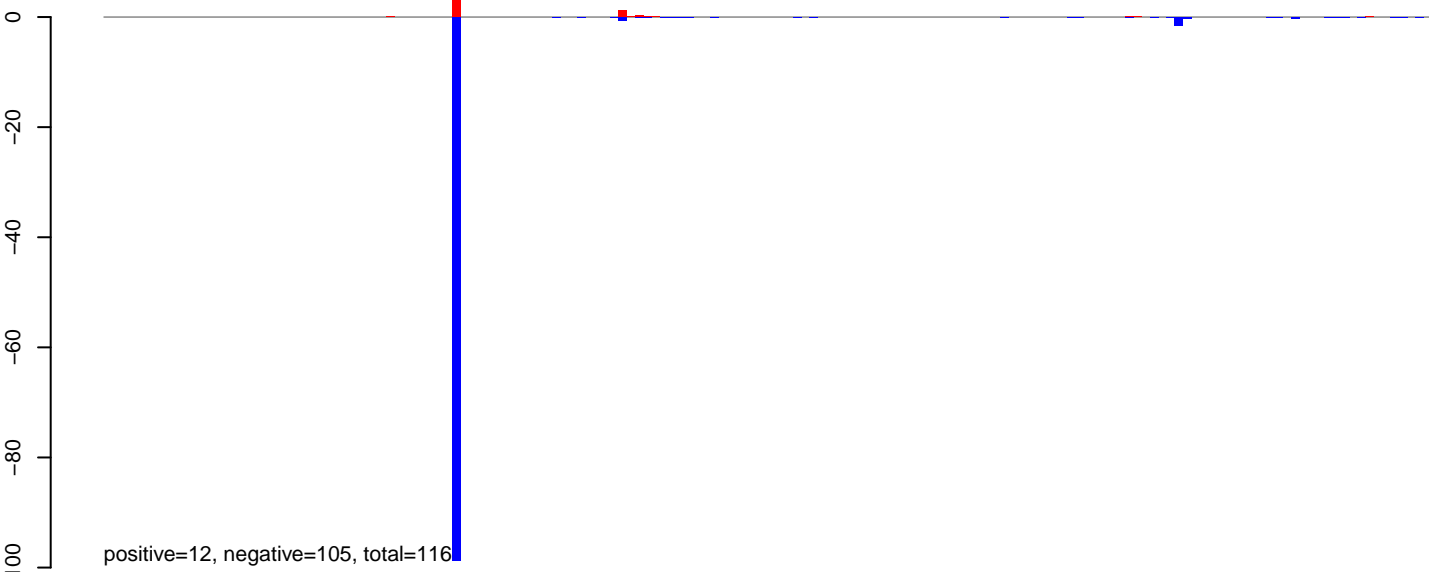
AnGam_Mos55_cells.rep



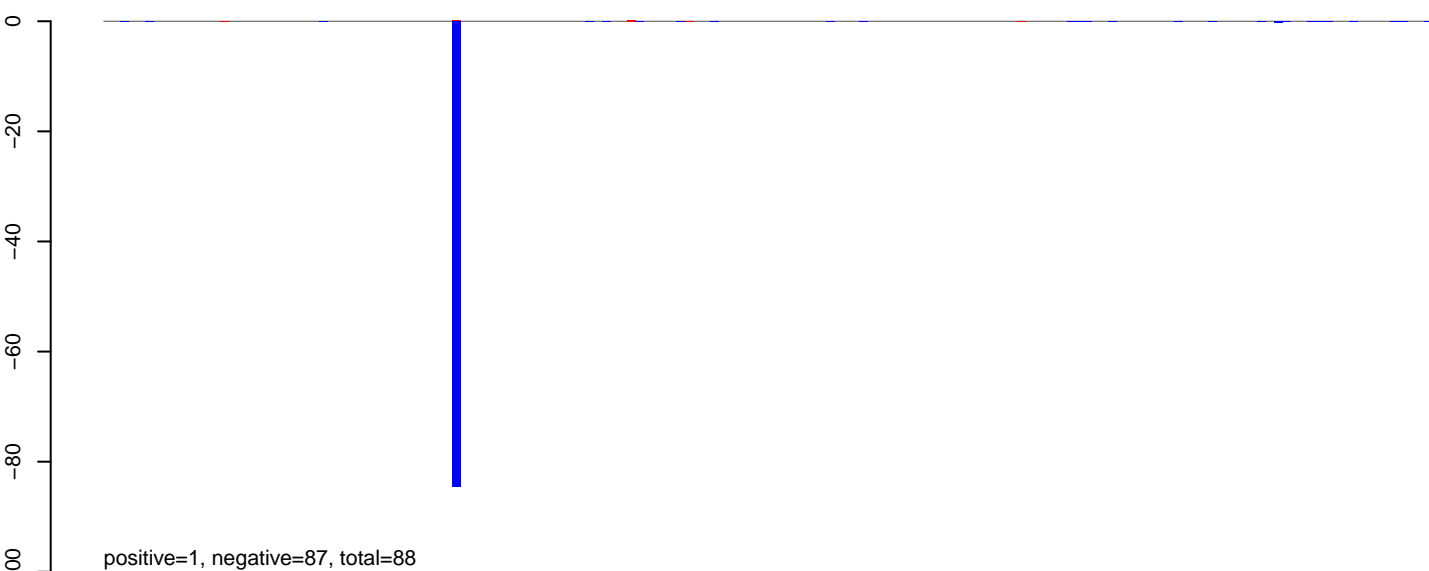
Window size=50, length=7072, TE@RTAg4:1-7072

0 1000 2000 3000 4000 5000 6000 7000

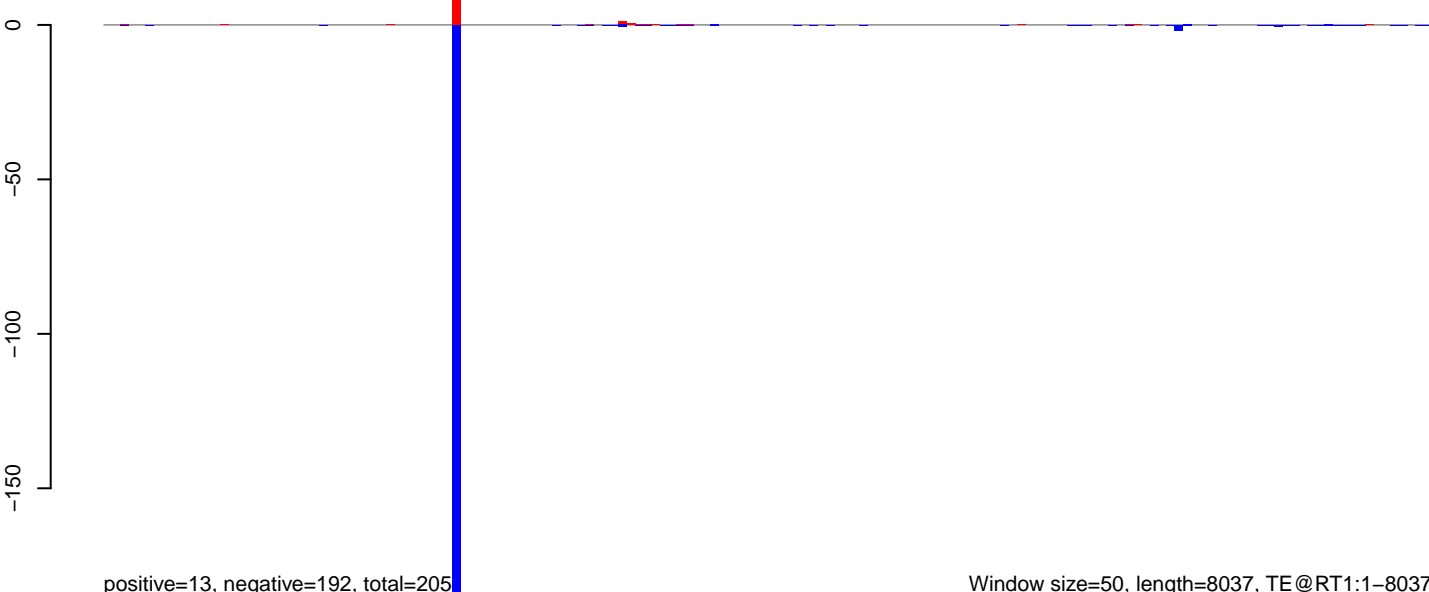
AnGam_Mos55_cells.18_23.rep



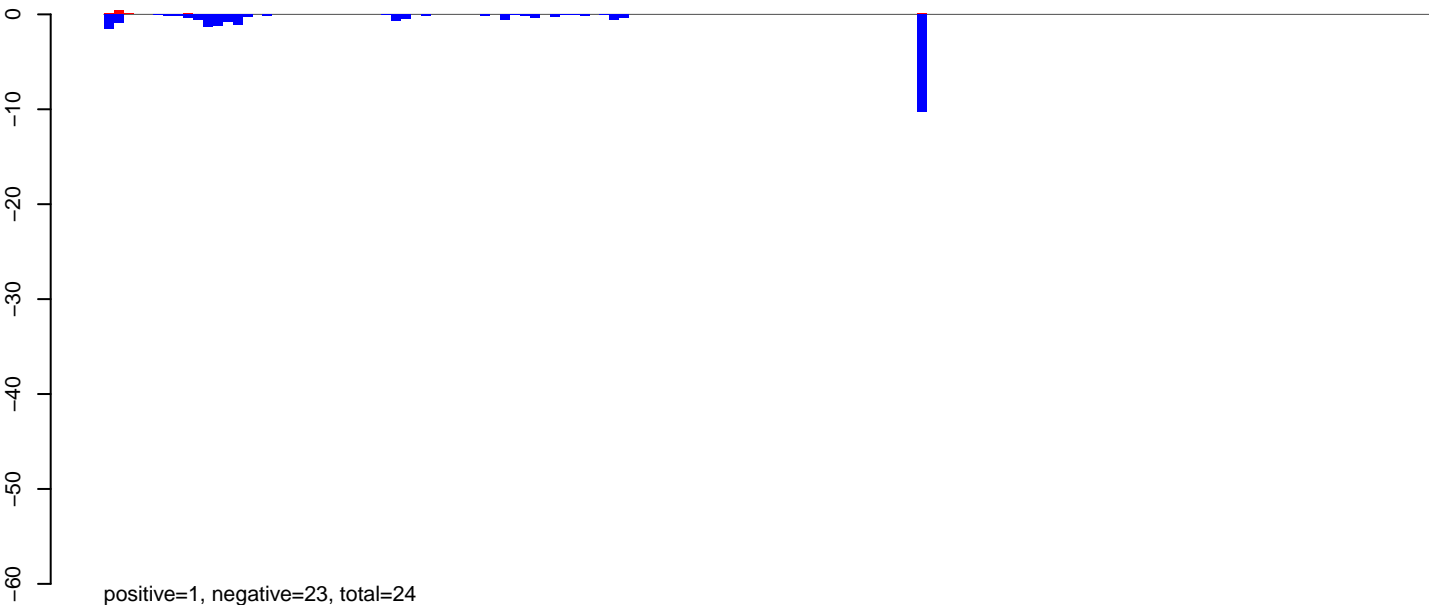
AnGam_Mos55_cells.24_35.rep



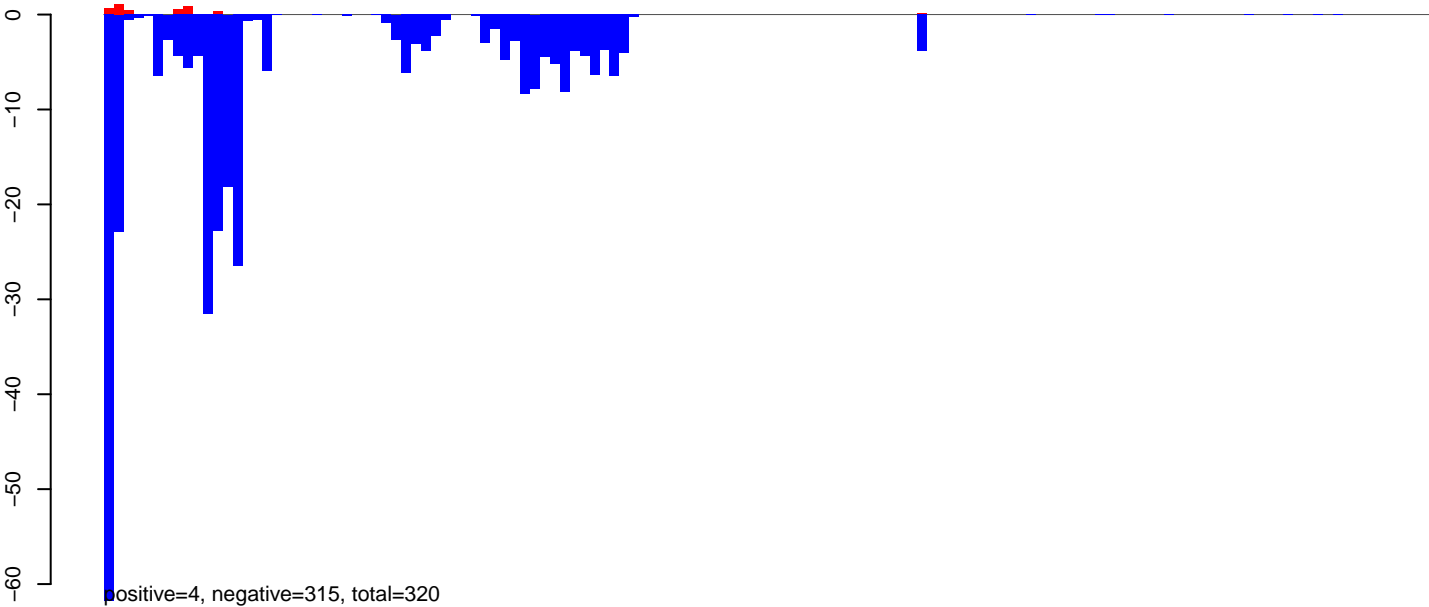
AnGam_Mos55_cells.rep



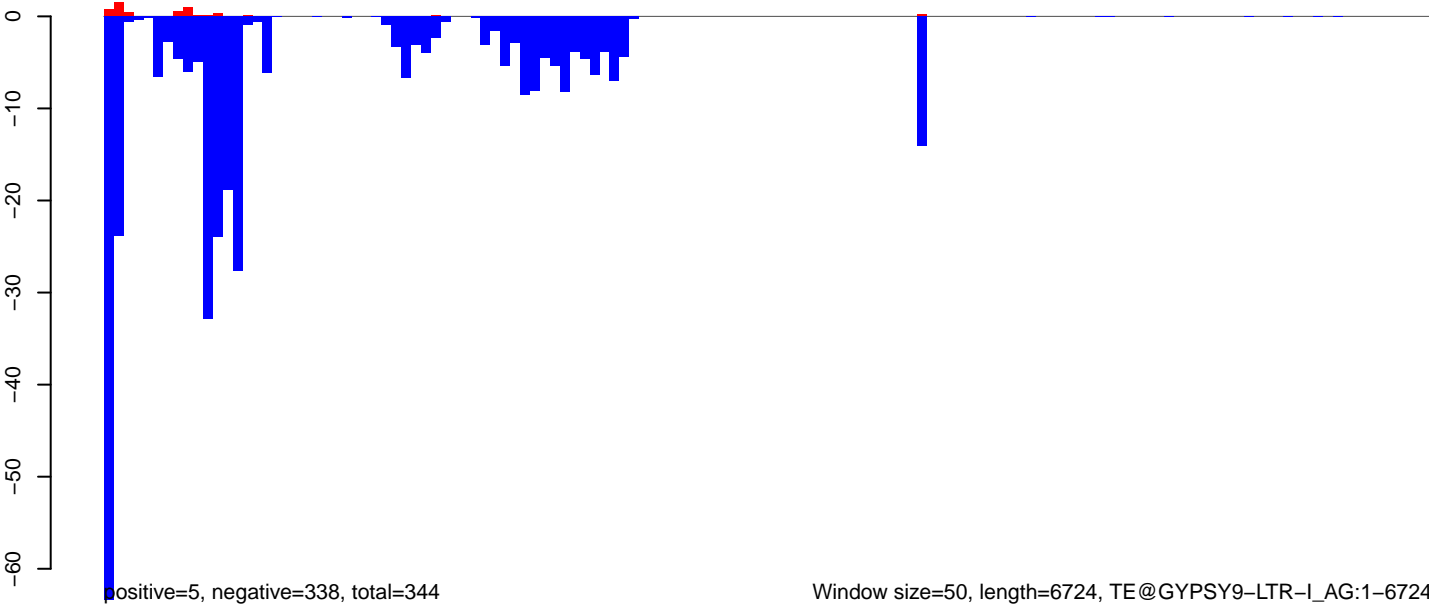
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



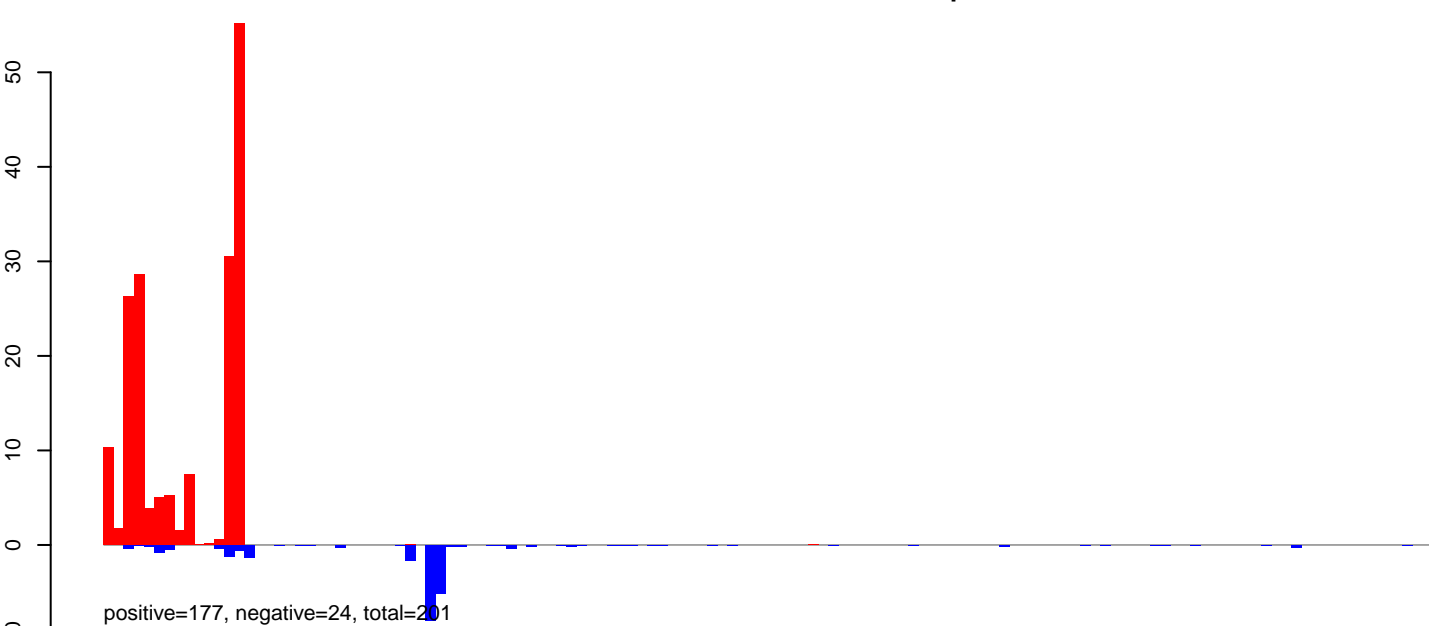
Window size=50, length=6724, TE@GYPSY9-LTR-I_AG:1-6724

0 1000 2000 3000 4000 5000 6000 7000

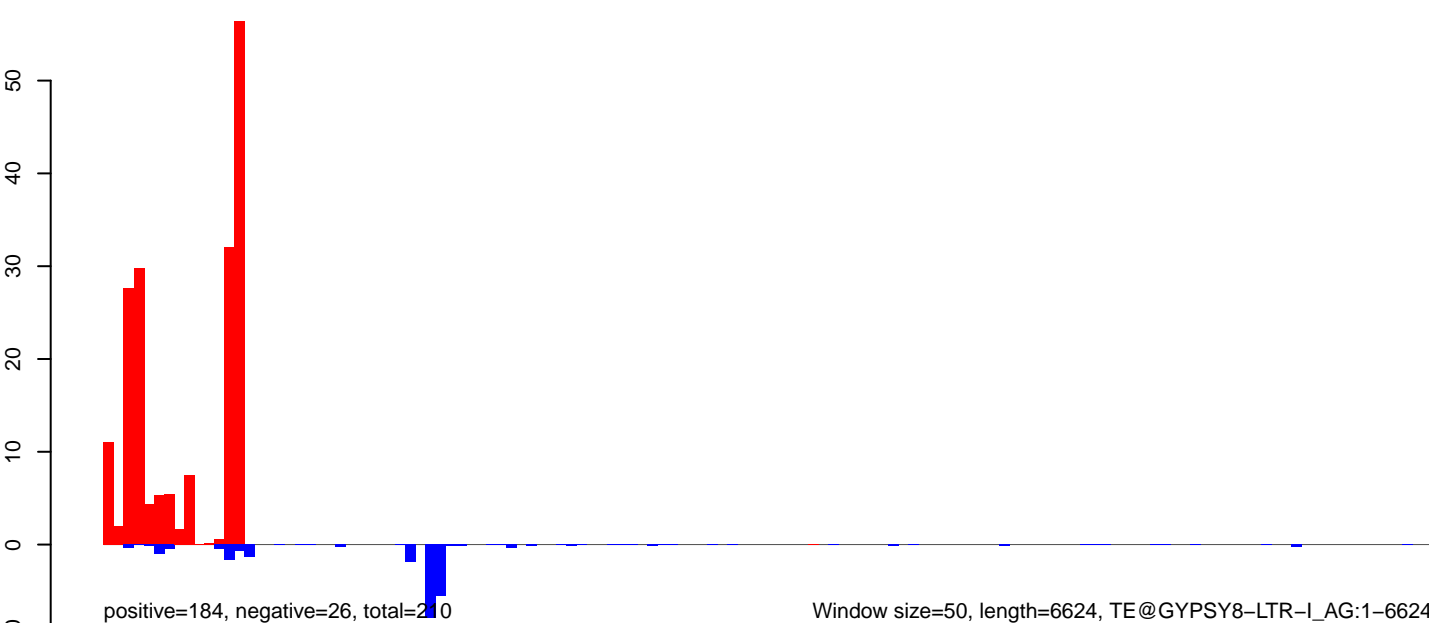
AnGam_Mos55_cells.18_23.rep



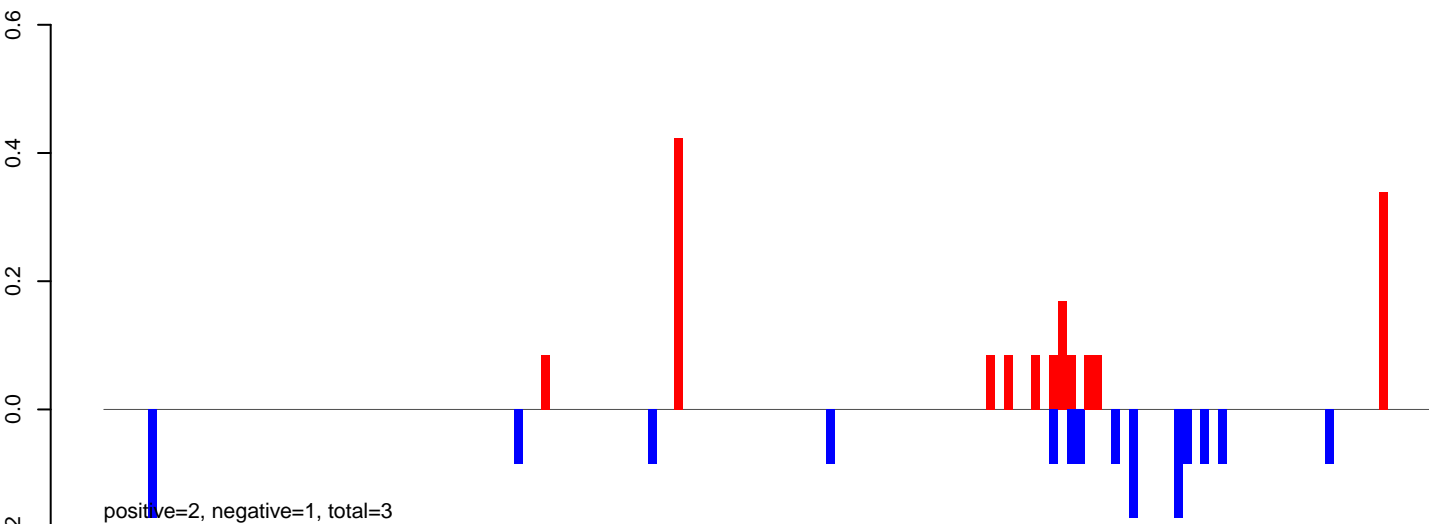
AnGam_Mos55_cells.24_35.rep



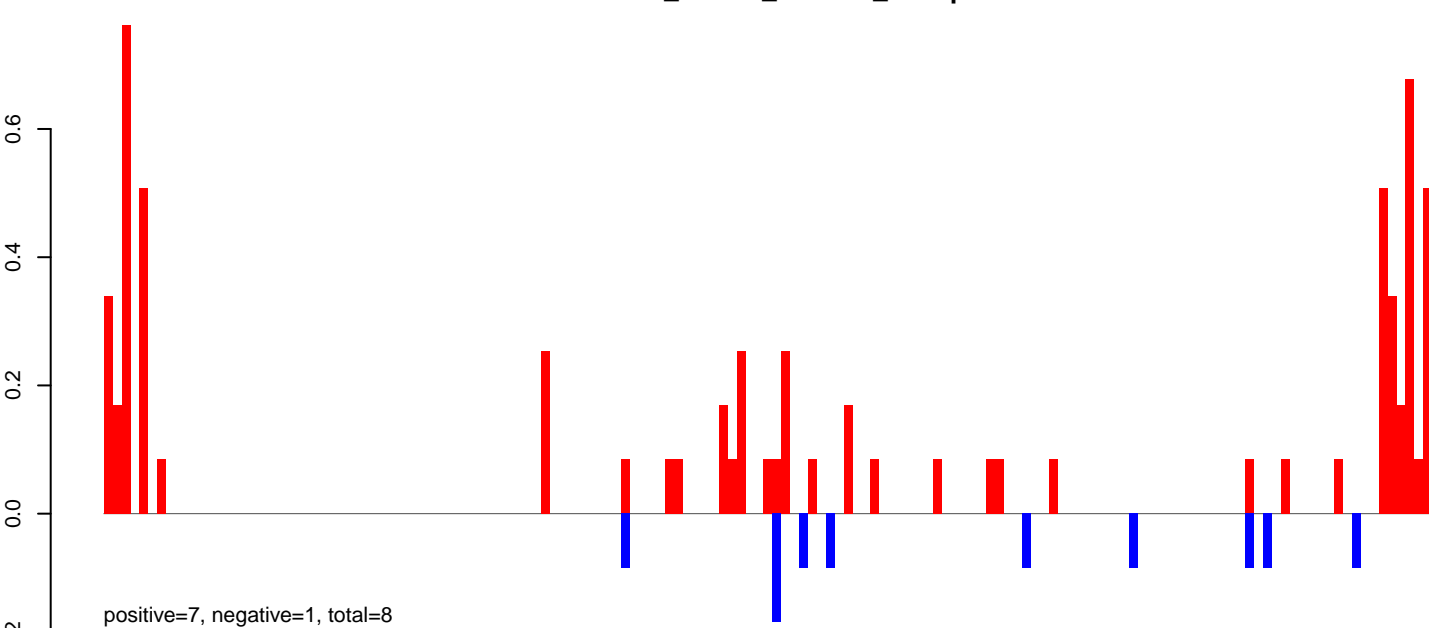
AnGam_Mos55_cells.rep



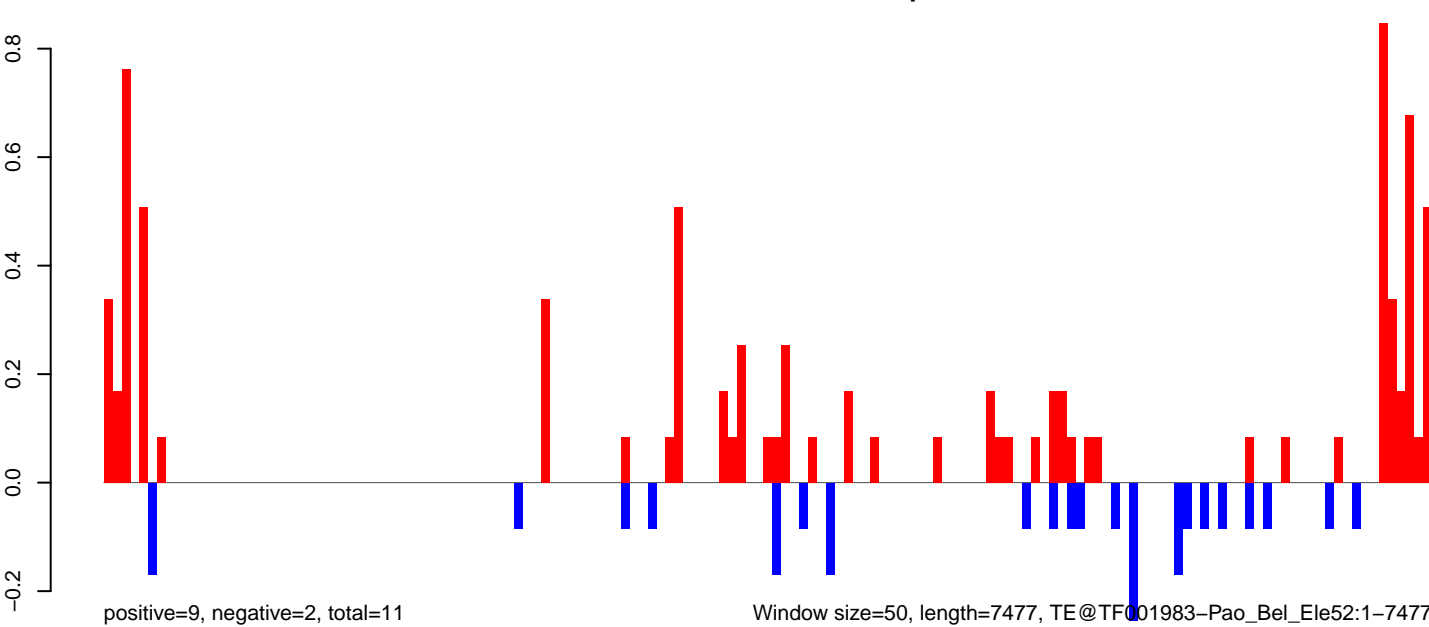
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

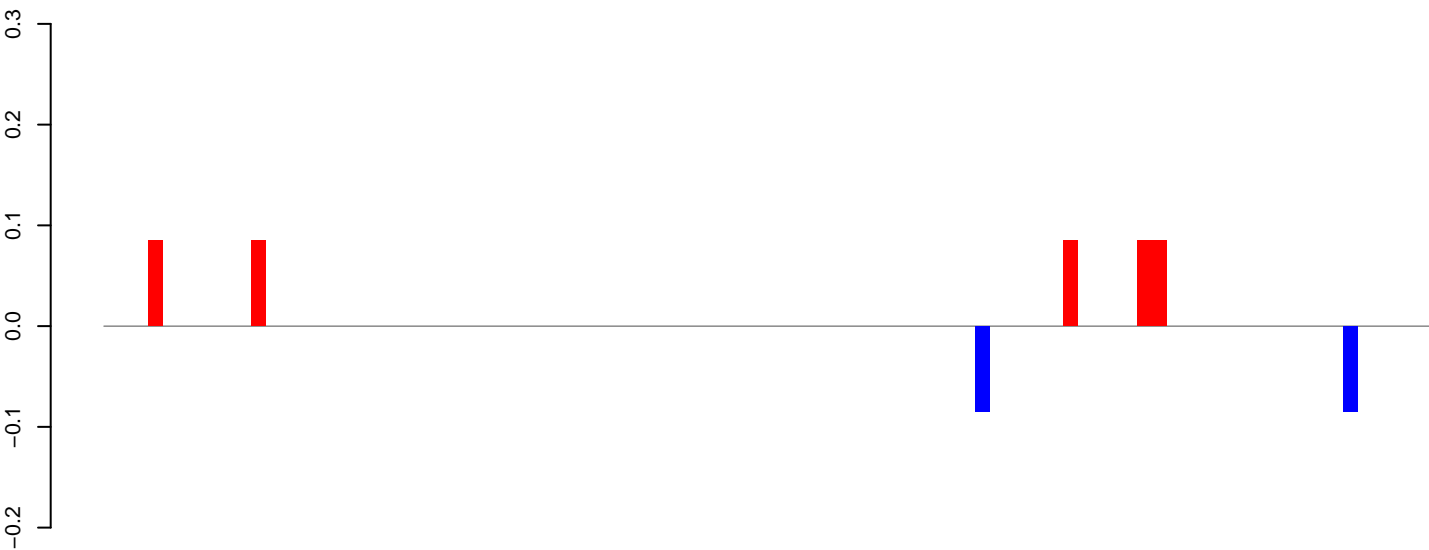


AnGam_Mos55_cells.rep



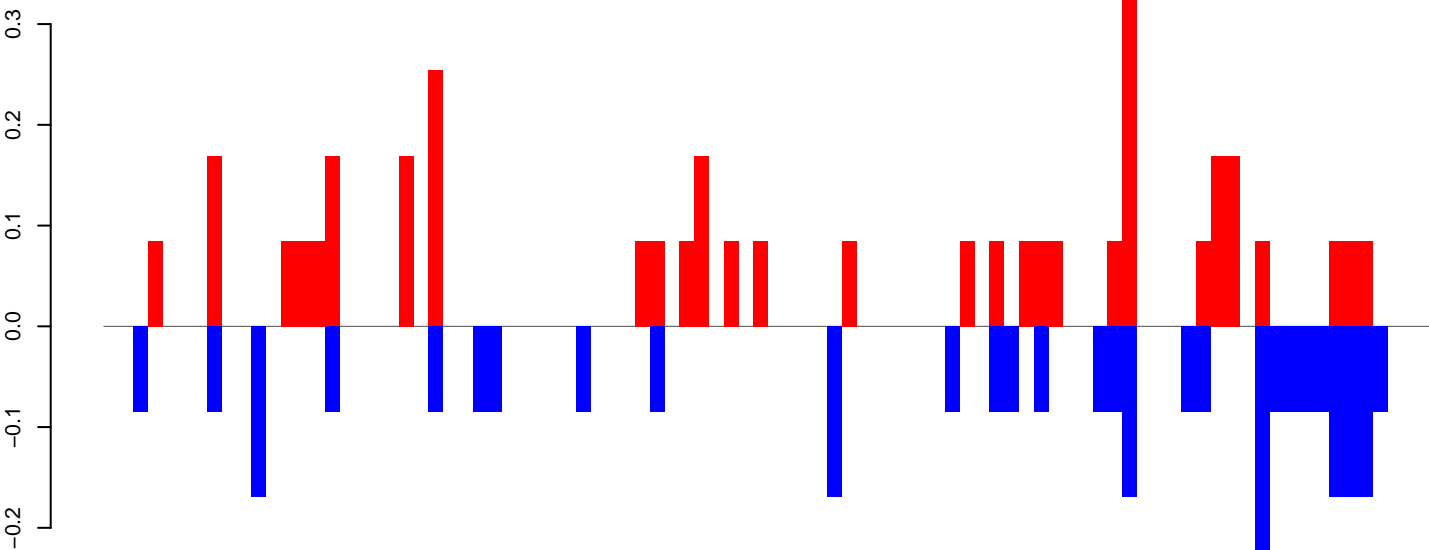
0 2000 4000 6000

AnGam_Mos55_cells.18_23.rep



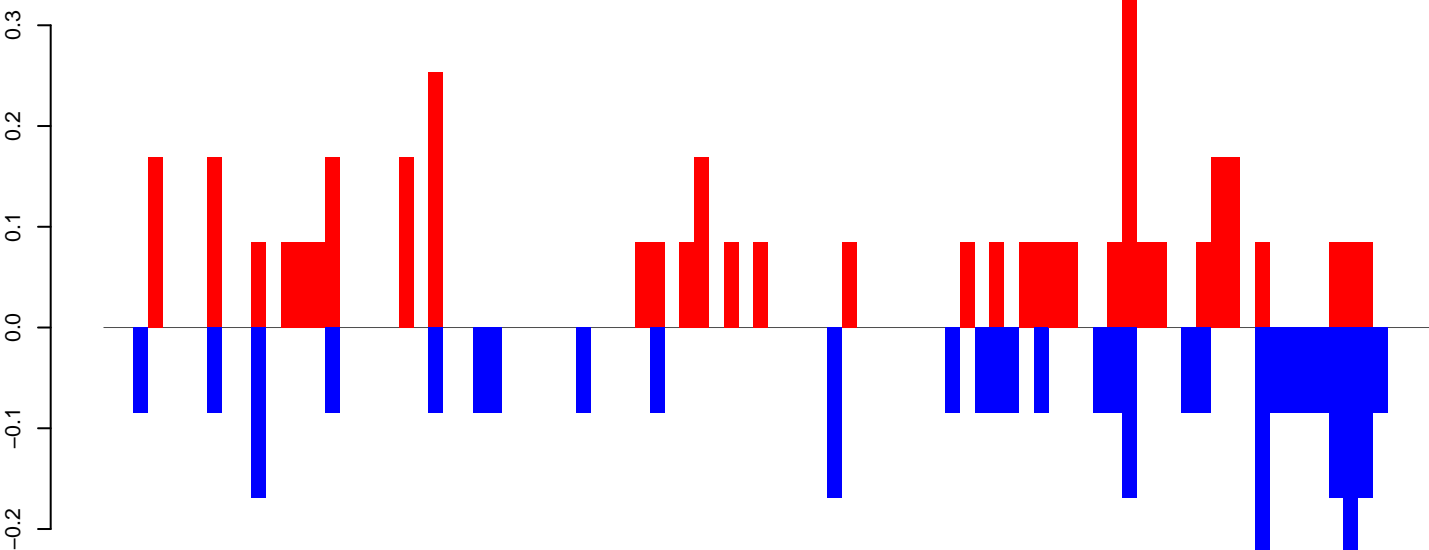
positive=0, negative=0, total=1

AnGam_Mos55_cells.24_35.rep



positive=3, negative=3, total=6

AnGam_Mos55_cells.rep

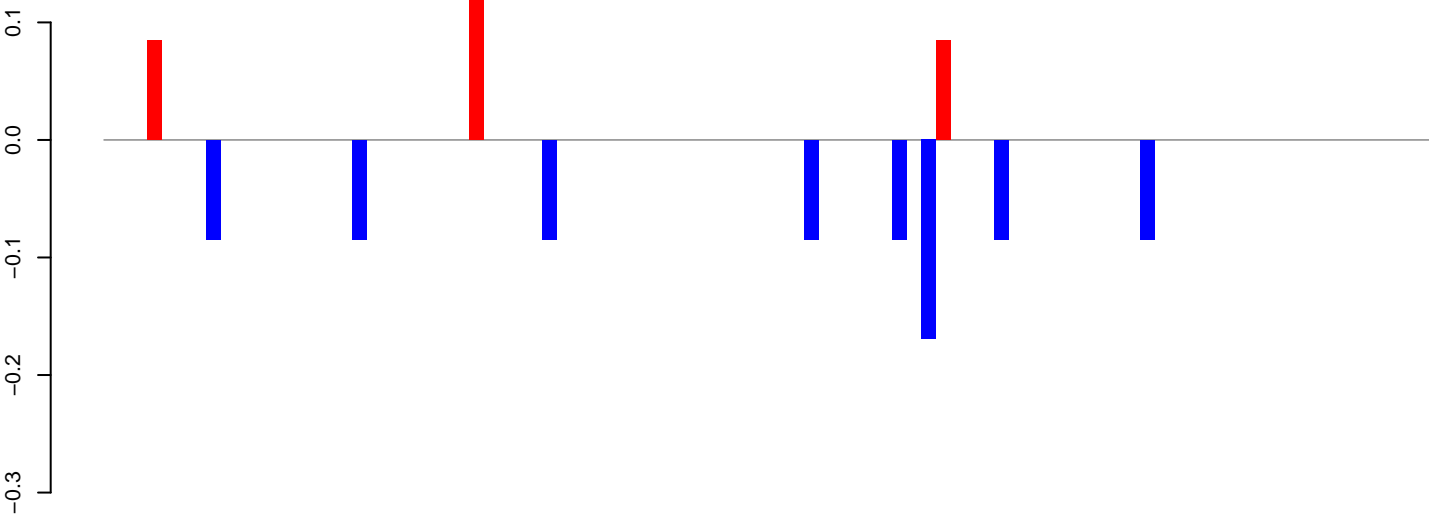


positive=4, negative=3, total=7

Window size=50, length=4546, TE@GYPSY64-LTR-L_AG:1-4546

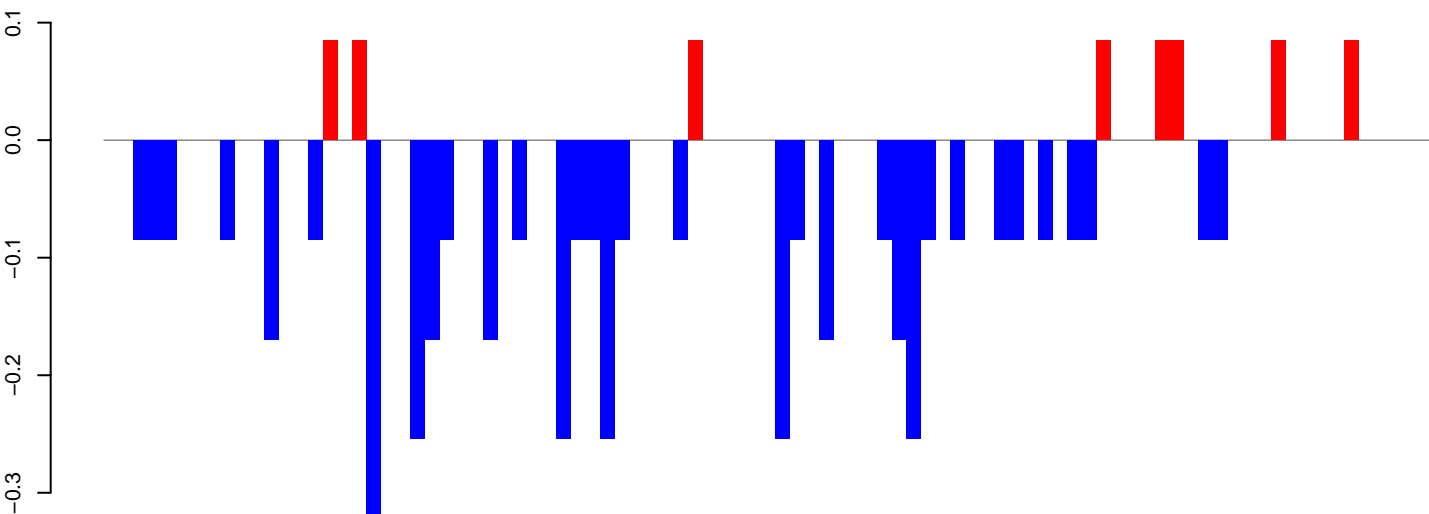
0 1000 2000 3000 4000

AnGam_Mos55_cells.18_23.rep



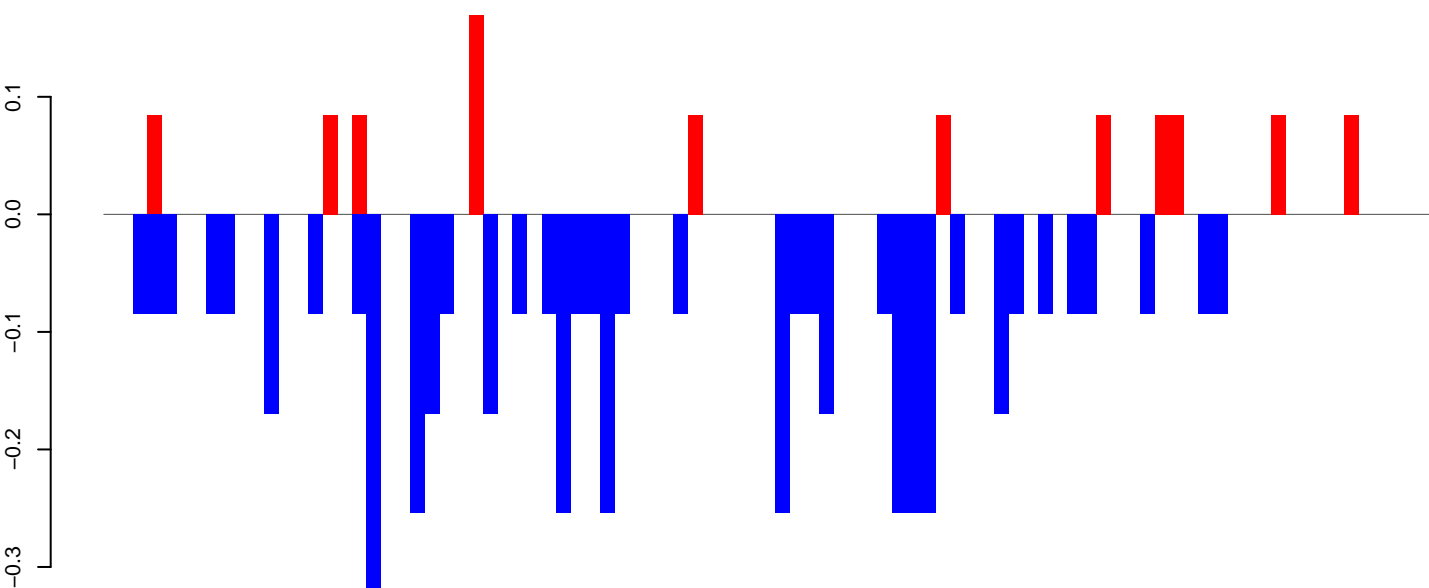
positive=0, negative=1, total=1

AnGam_Mos55_cells.24_35.rep



positive=1, negative=4, total=5

AnGam_Mos55_cells.rep



positive=1, negative=5, total=6

Window size=50, length=4556, TE@GYPSY23-LTR-I_AG:1-4556

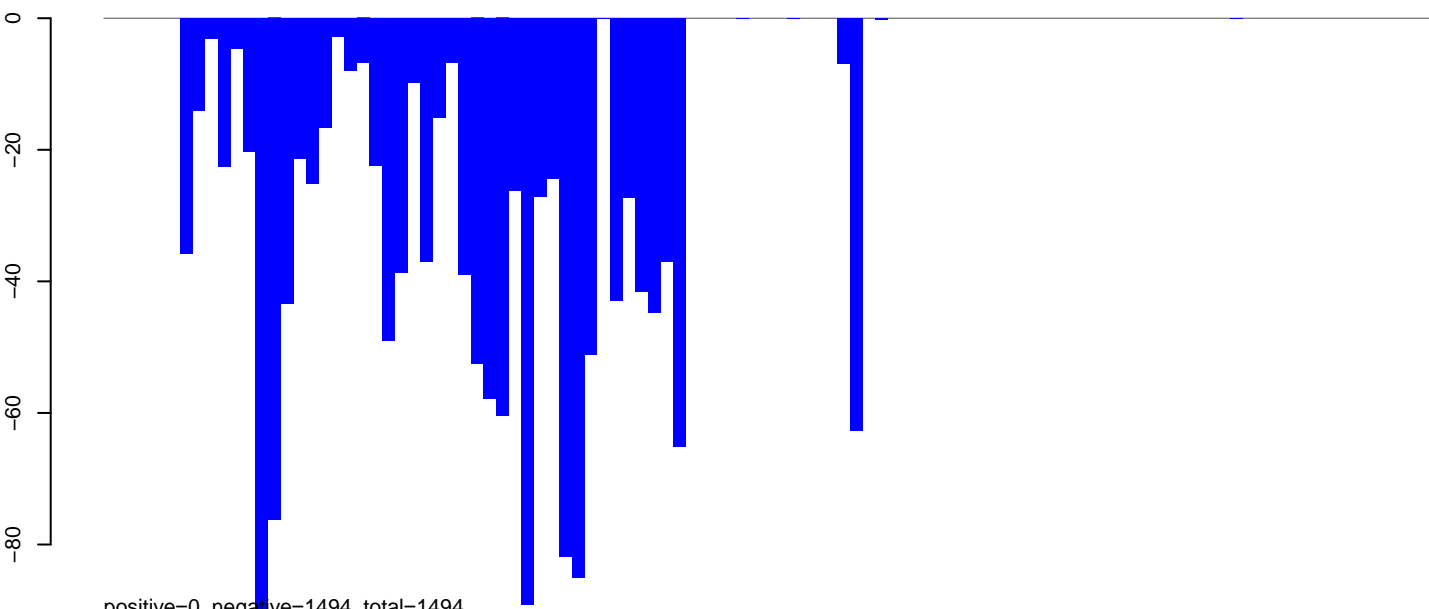
0 1000 2000 3000 4000

AnGam_Mos55_cells.18_23.rep



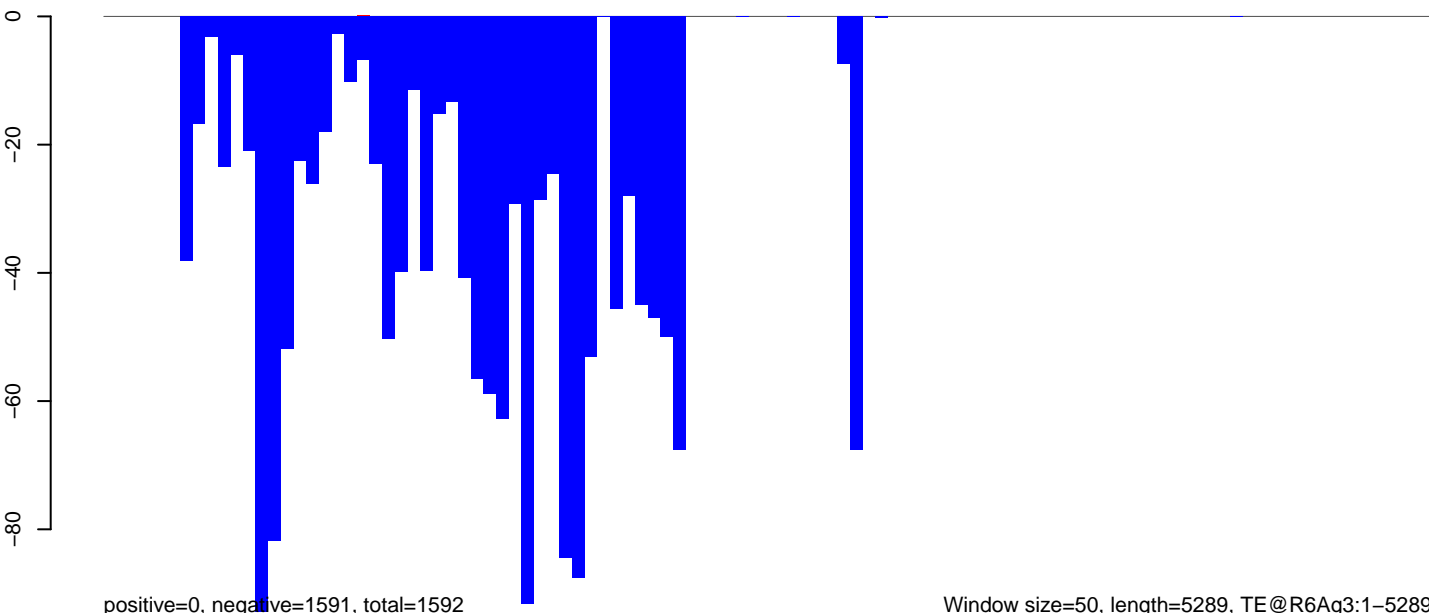
positive=0, negative=98, total=98

AnGam_Mos55_cells.24_35.rep



positive=0, negative=1494, total=1494

AnGam_Mos55_cells.rep

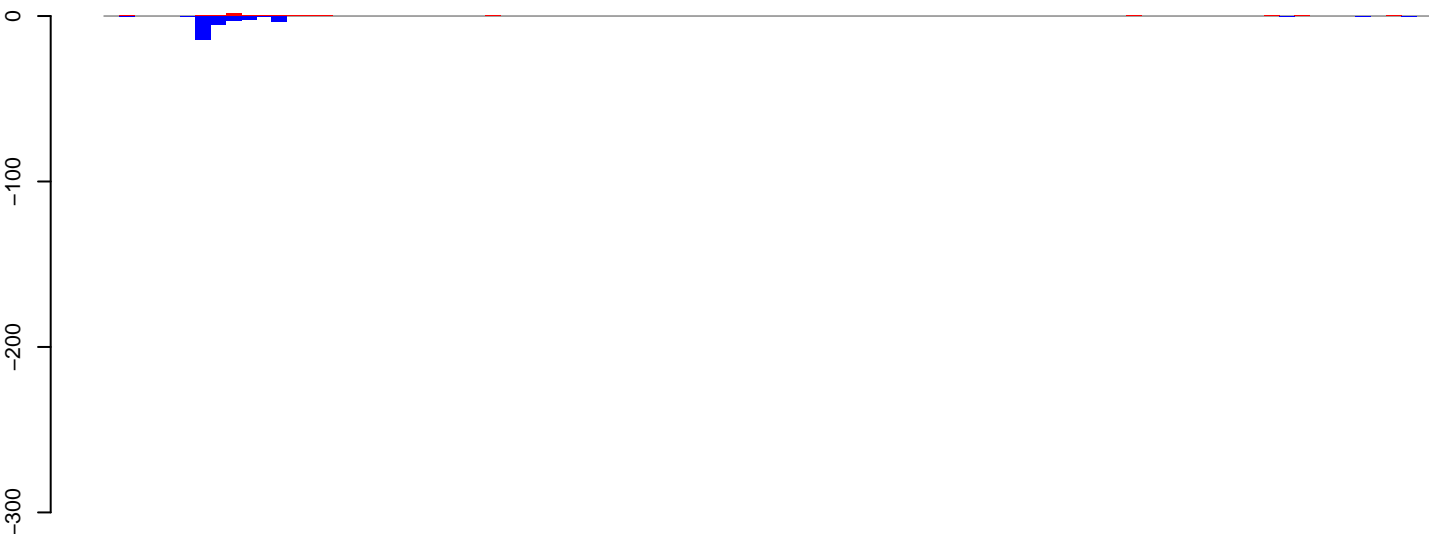


positive=0, negative=1591, total=1592

Window size=50, length=5289, TE@R6Ag3:1-5289

0 1000 2000 3000 4000 5000

AnGam_Mos55_cells.18_23.rep



positive=4, negative=29, total=32

AnGam_Mos55_cells.24_35.rep



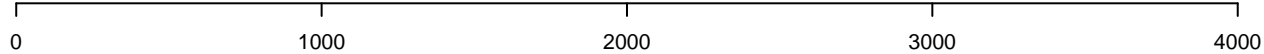
positive=34, negative=1063, total=1098

AnGam_Mos55_cells.rep

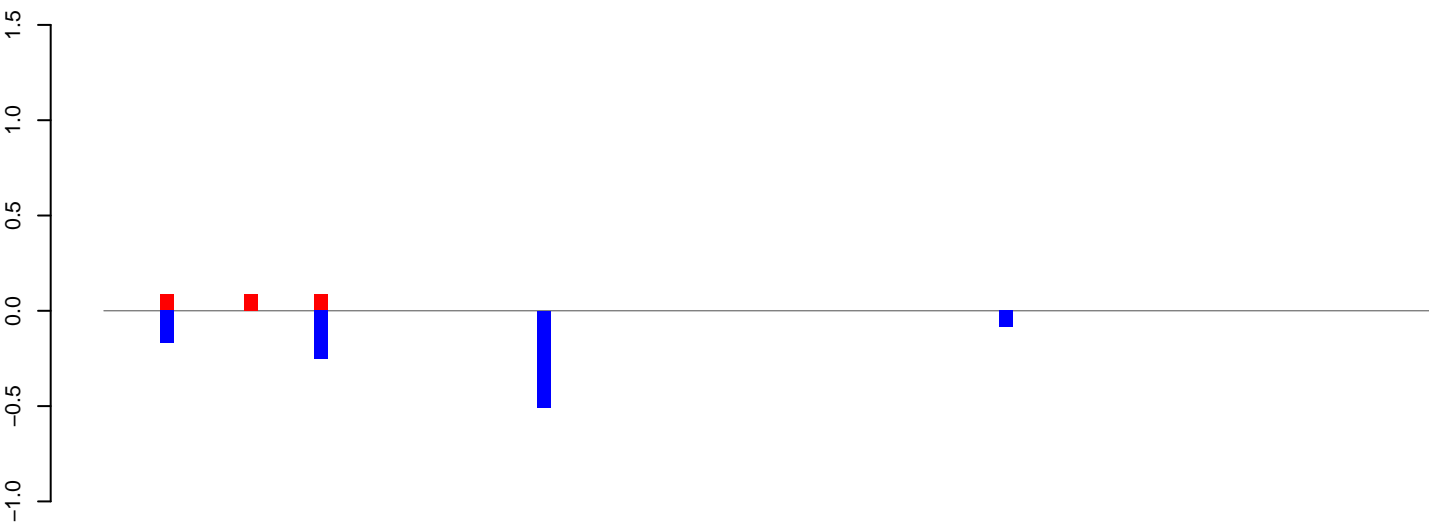


positive=38, negative=1092, total=1130

Window size=50, length=4352, TE@TF001996-Ty1_copia_Ele10:1-4352

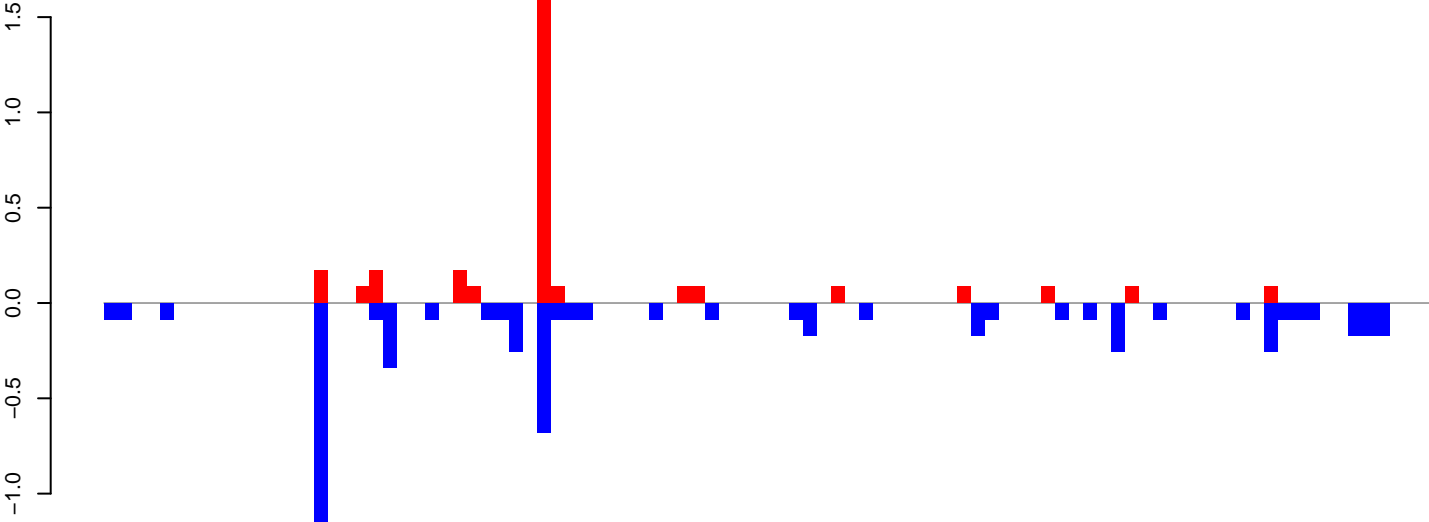


AnGam_Mos55_cells.18_23.rep



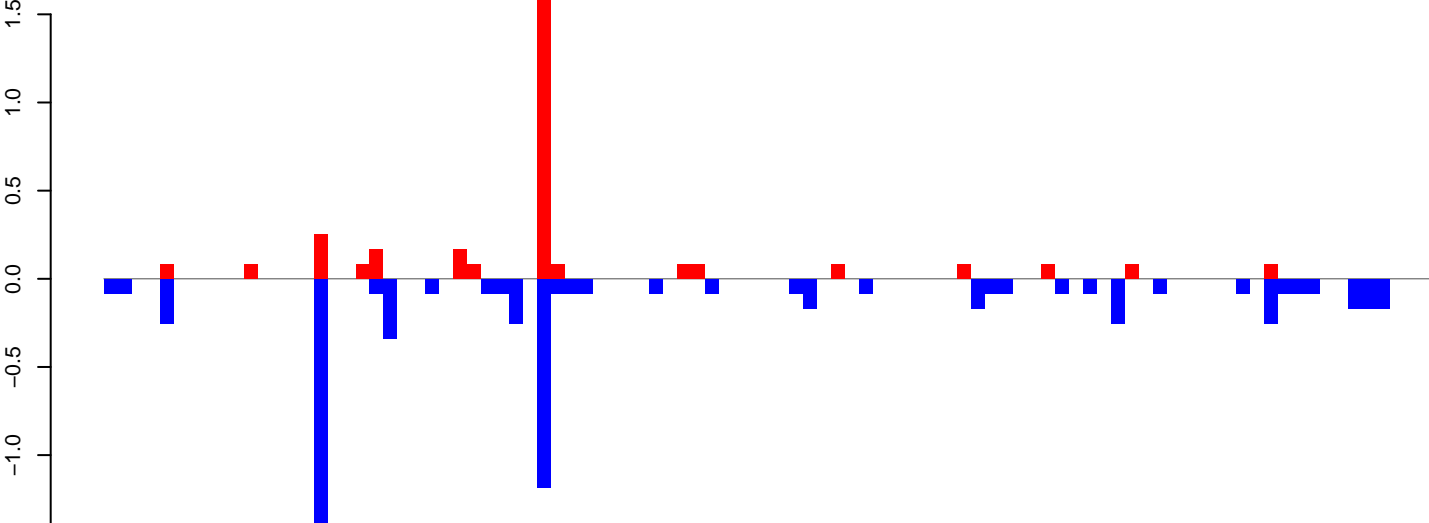
positive=0, negative=1, total=1

AnGam_Mos55_cells.24_35.rep



positive=3, negative=6, total=9

AnGam_Mos55_cells.rep

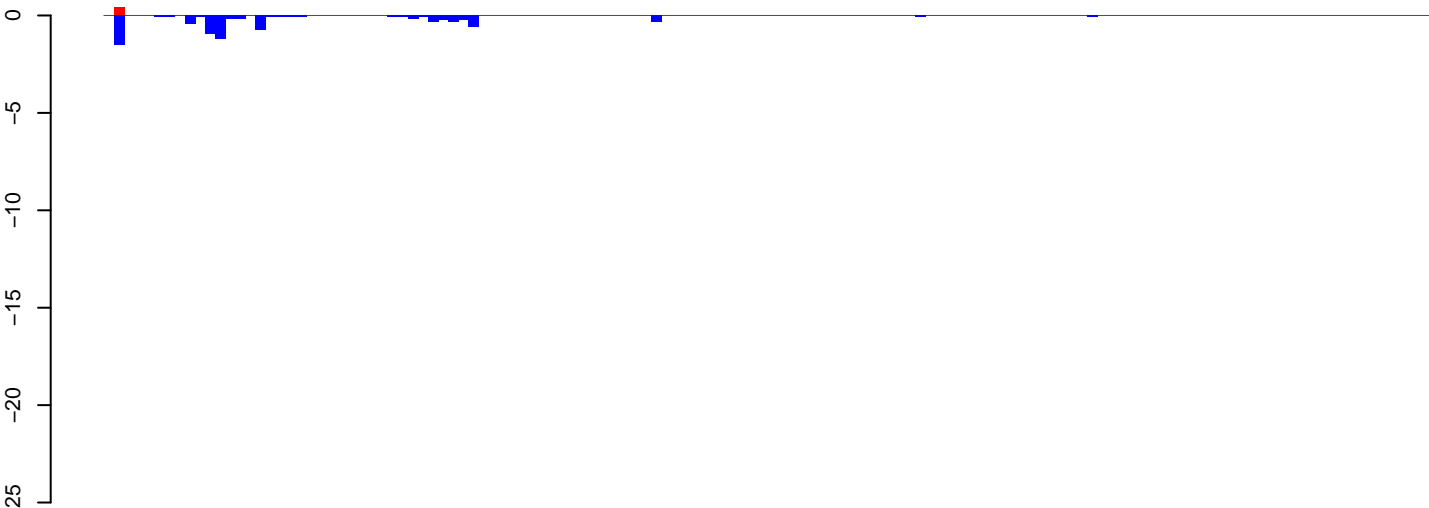


positive=3, negative=7, total=10

Window size=50, length=4799, TE@GYPSY33-LTR-L_AG:1-4799

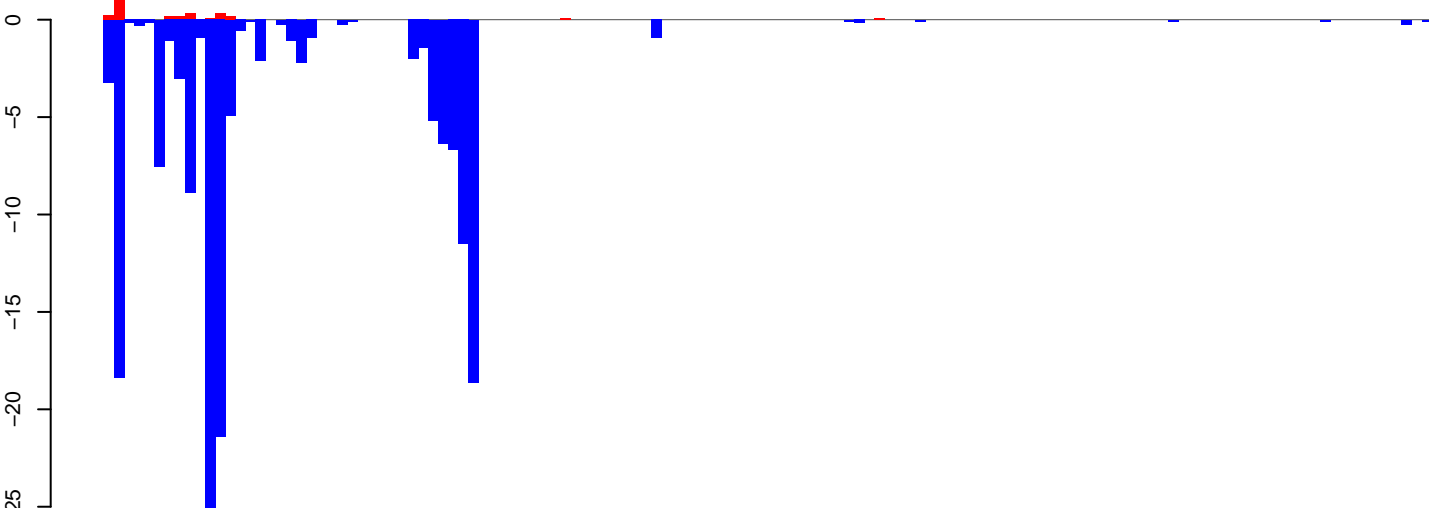
0 1000 2000 3000 4000

AnGam_Mos55_cells.18_23.rep



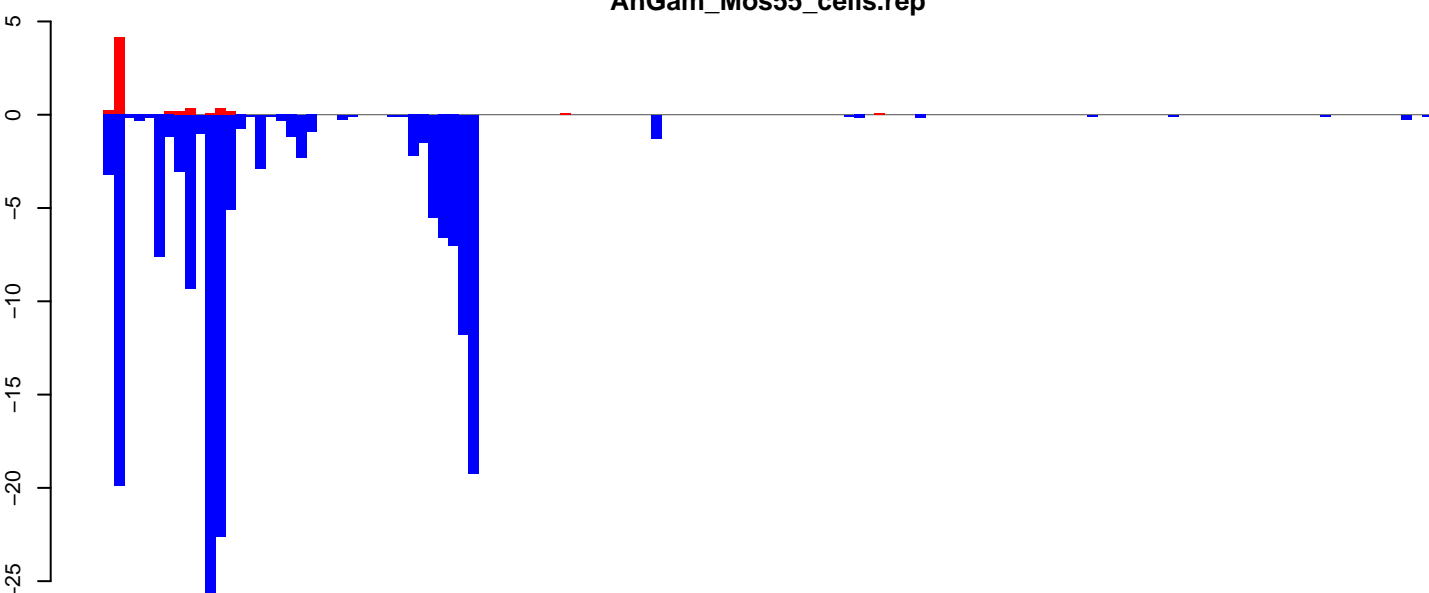
positive=0, negative=8, total=9

AnGam_Mos55_cells.24_35.rep



positive=5, negative=158, total=164

AnGam_Mos55_cells.rep

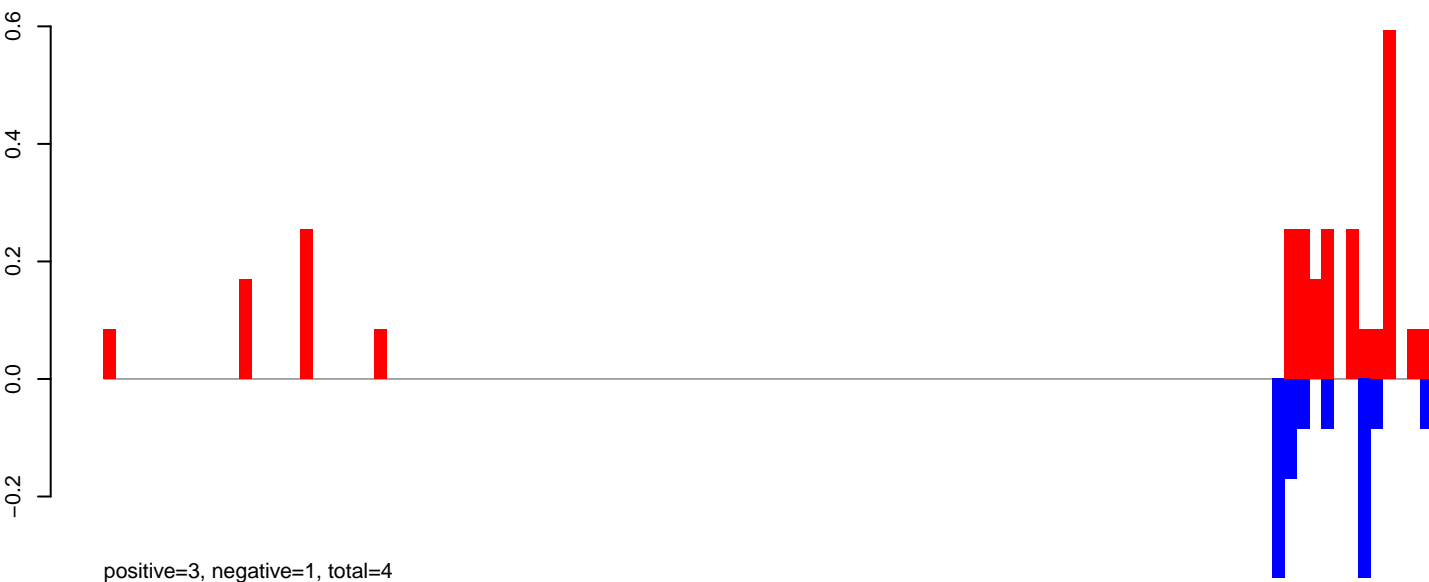


positive=6, negative=167, total=173

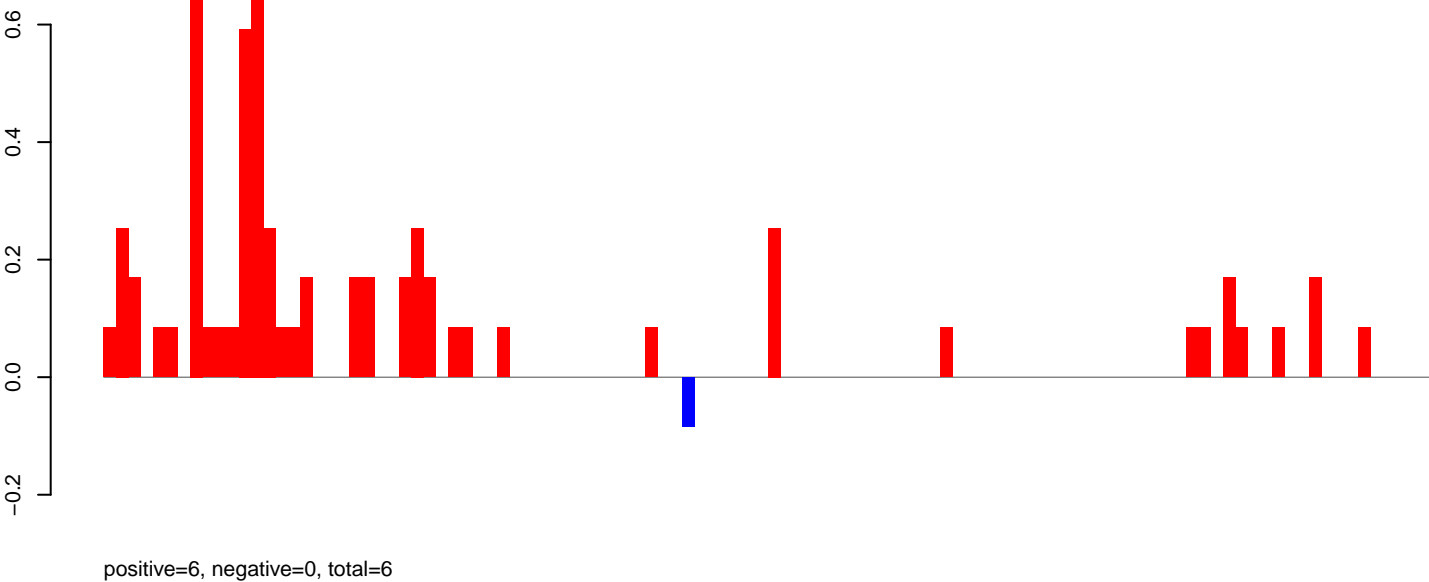
Window size=50, length=6600, TE@GYPSY15-LTR-I_AG:1-6600

0 1000 2000 3000 4000 5000 6000

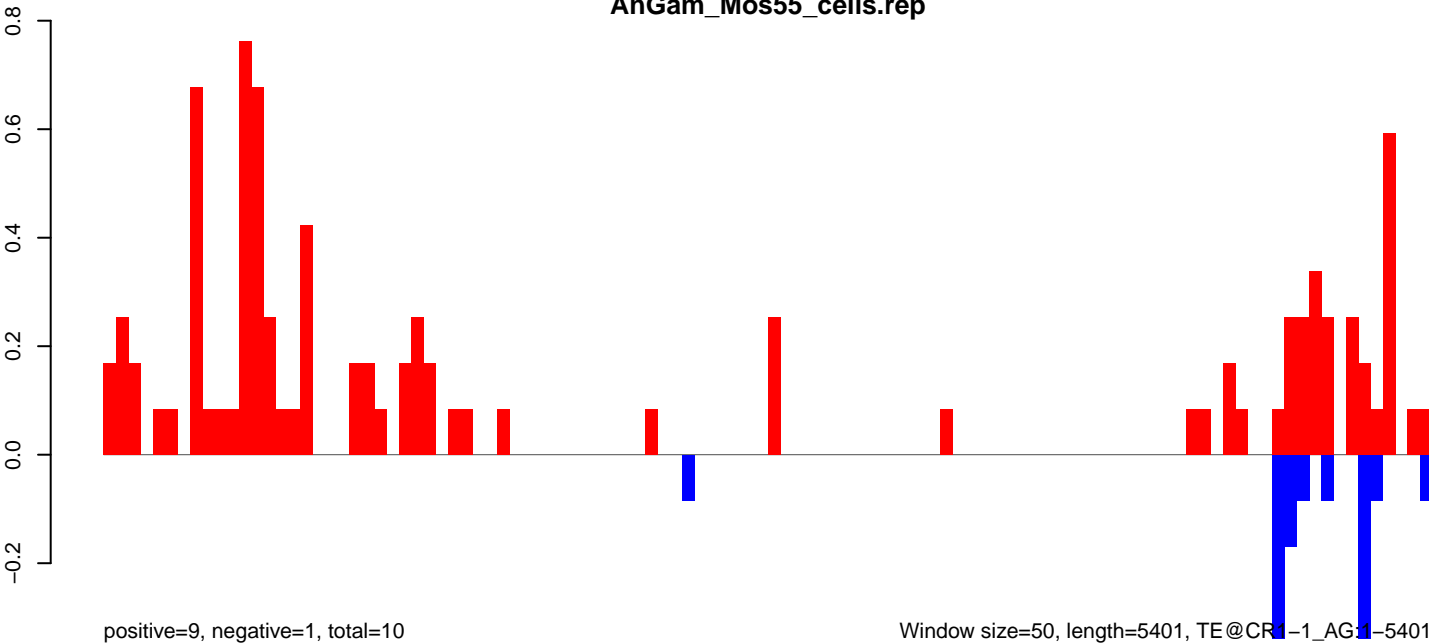
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

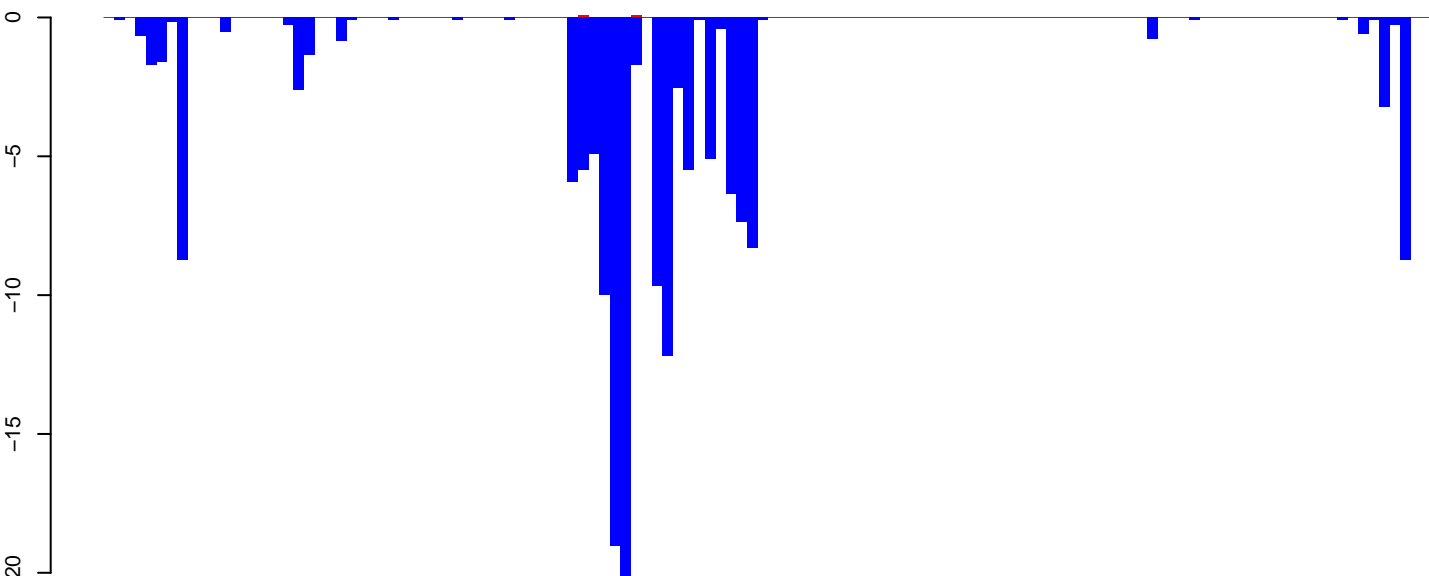


AnGam_Mos55_cells.18_23.rep



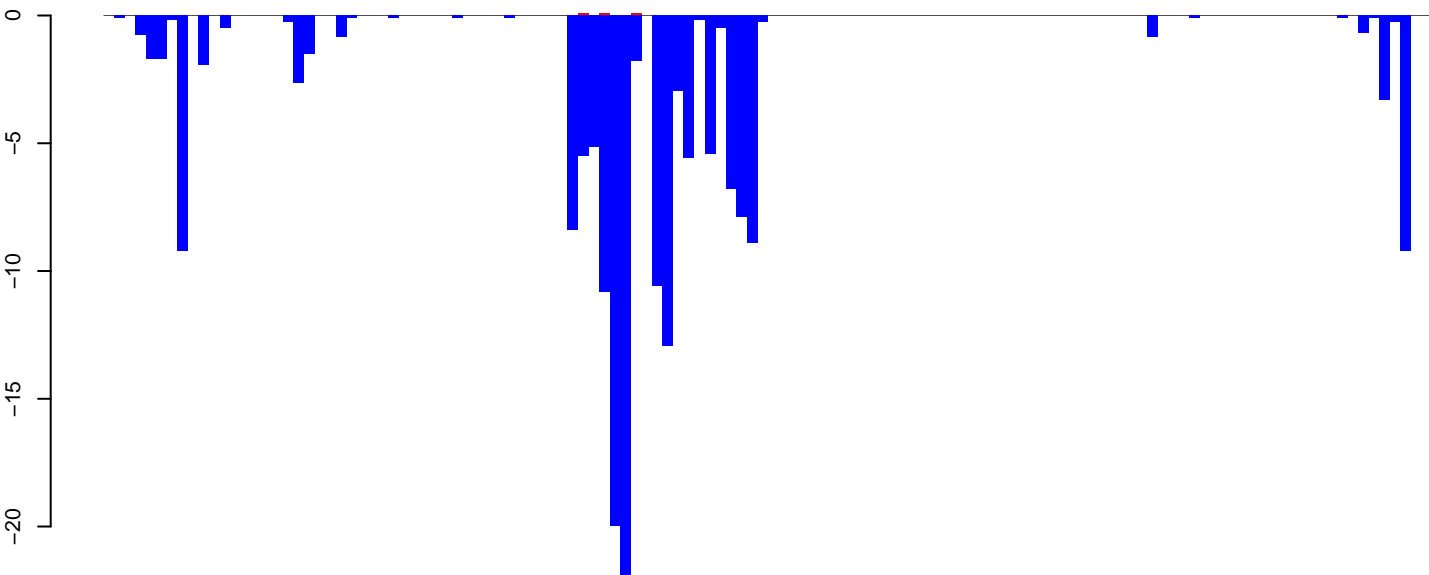
positive=0, negative=14, total=14

AnGam_Mos55_cells.24_35.rep



positive=0, negative=159, total=159

AnGam_Mos55_cells.rep

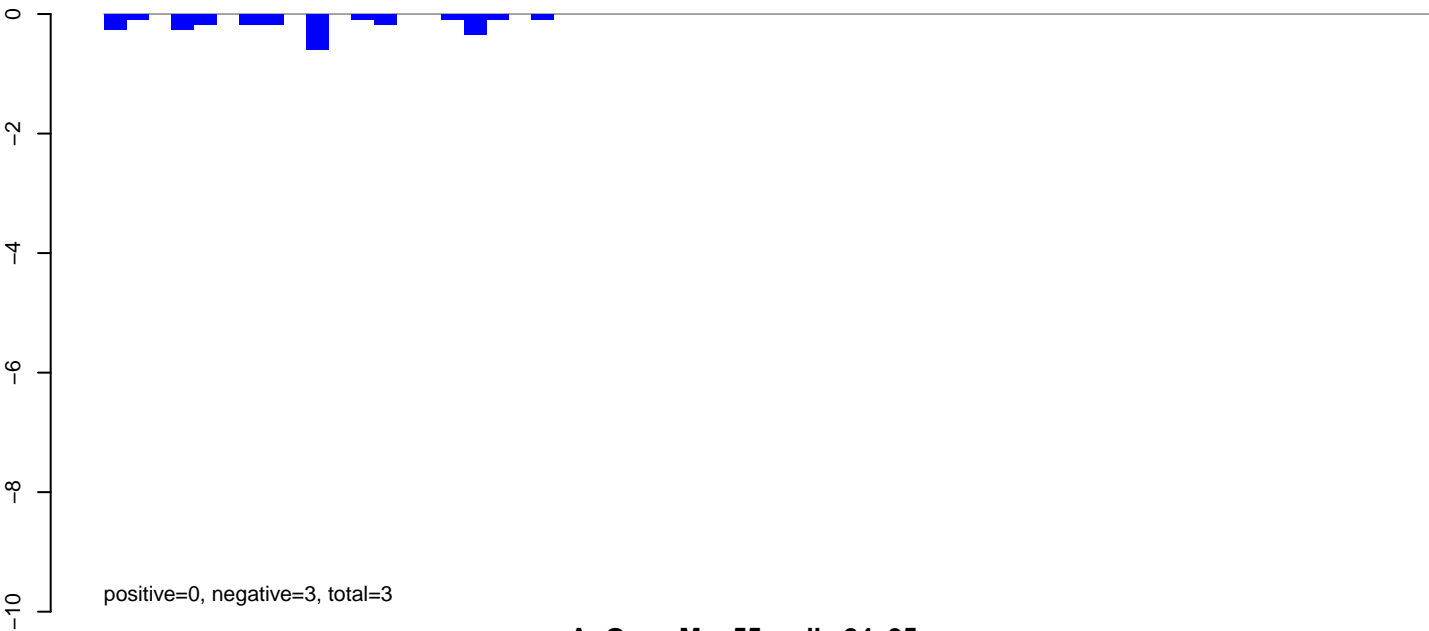


positive=0, negative=173, total=173

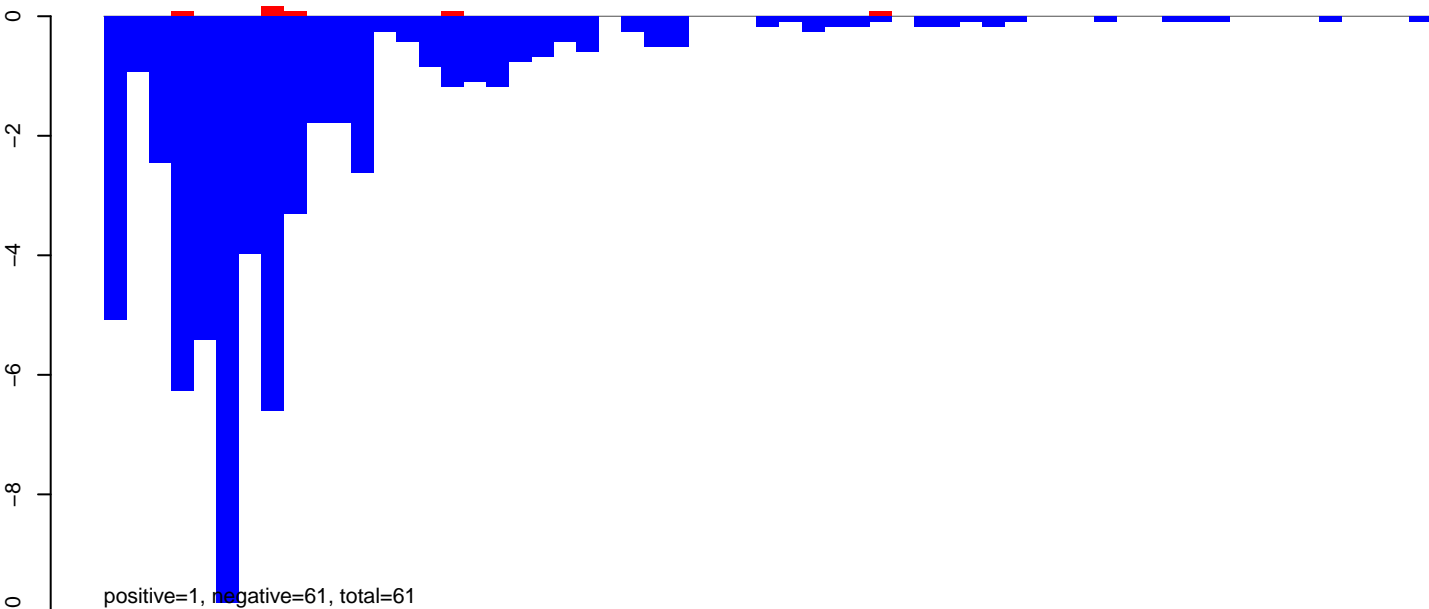
Window size=50, length=6301, TE@TF001949-Pao_Bel_Ele18:1-6301

0 1000 2000 3000 4000 5000 6000

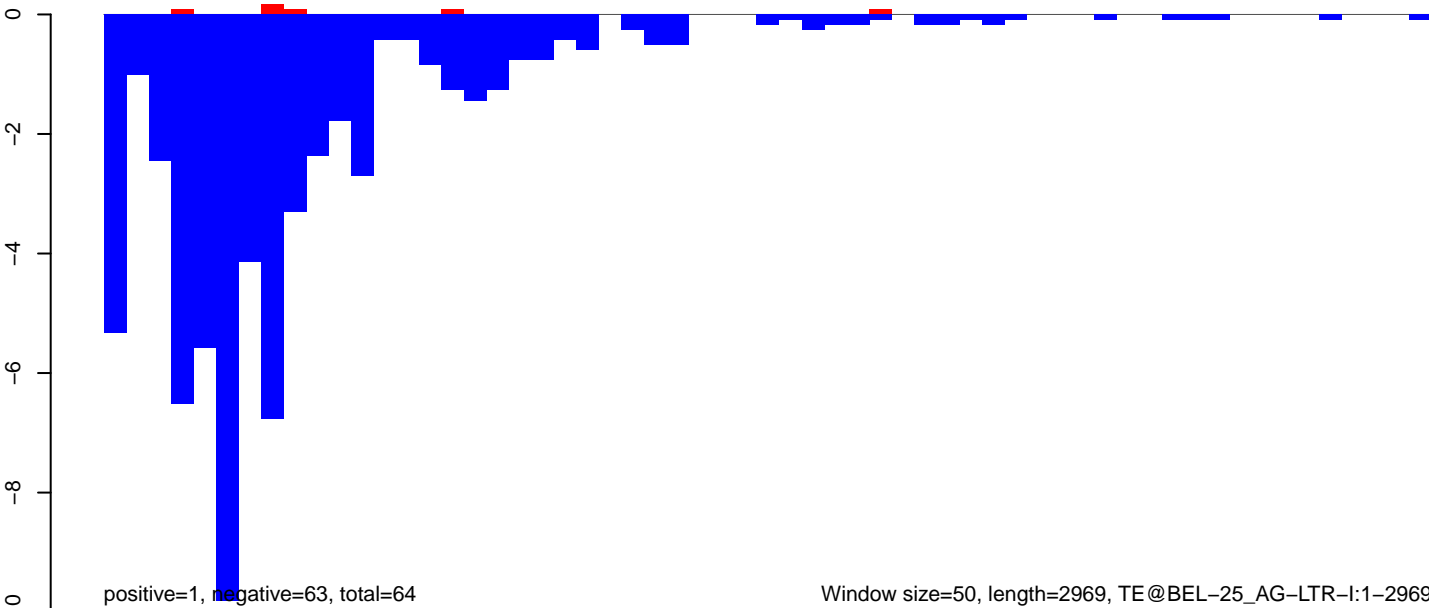
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

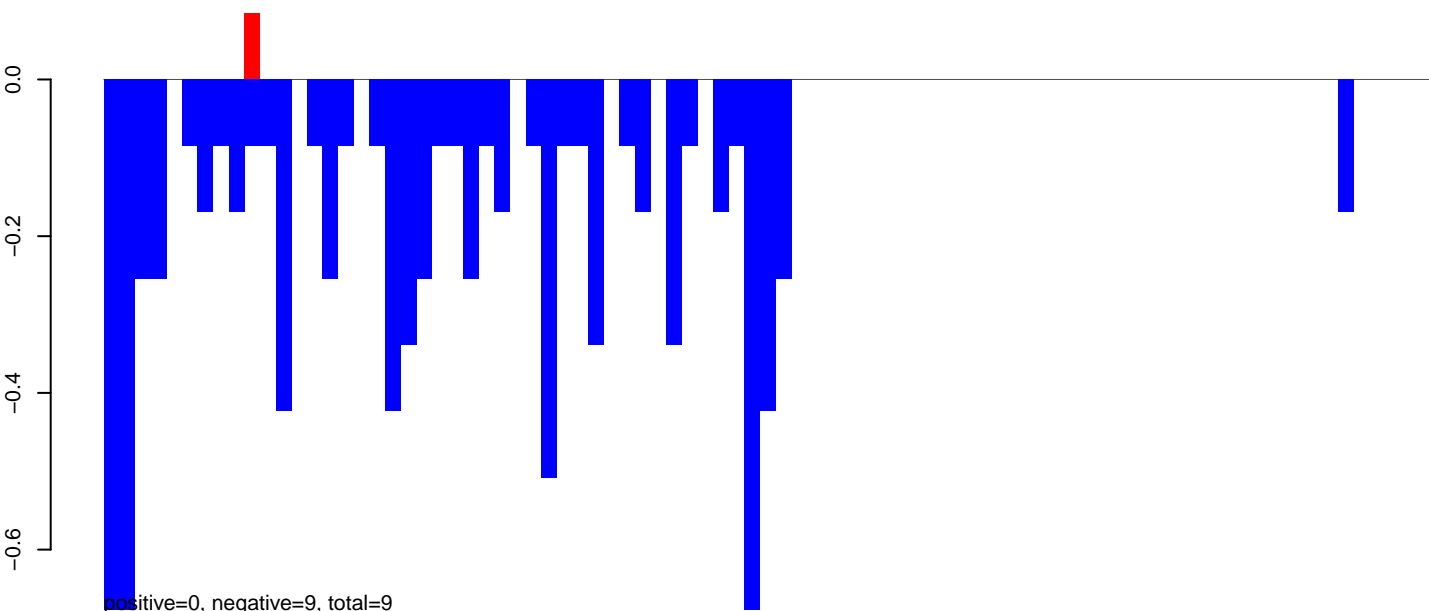


AnGam_Mos55_cells.18_23.rep



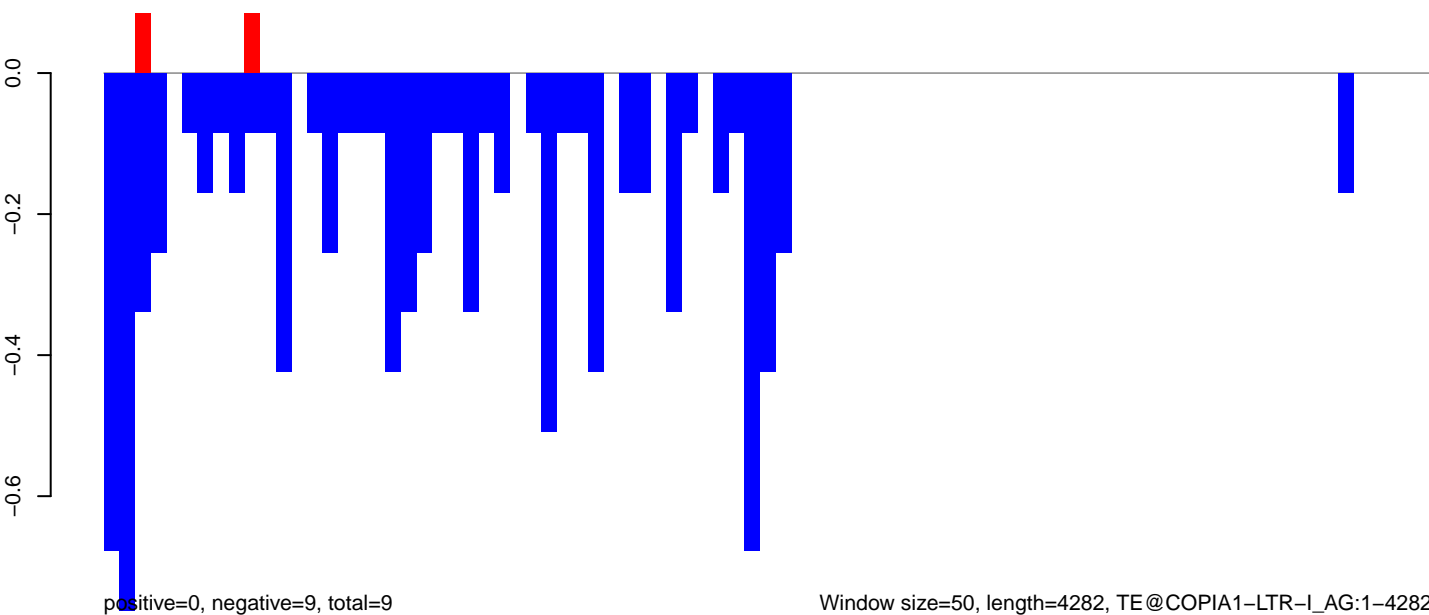
positive=0, negative=1, total=1

AnGam_Mos55_cells.24_35.rep



positive=0, negative=9, total=9

AnGam_Mos55_cells.rep

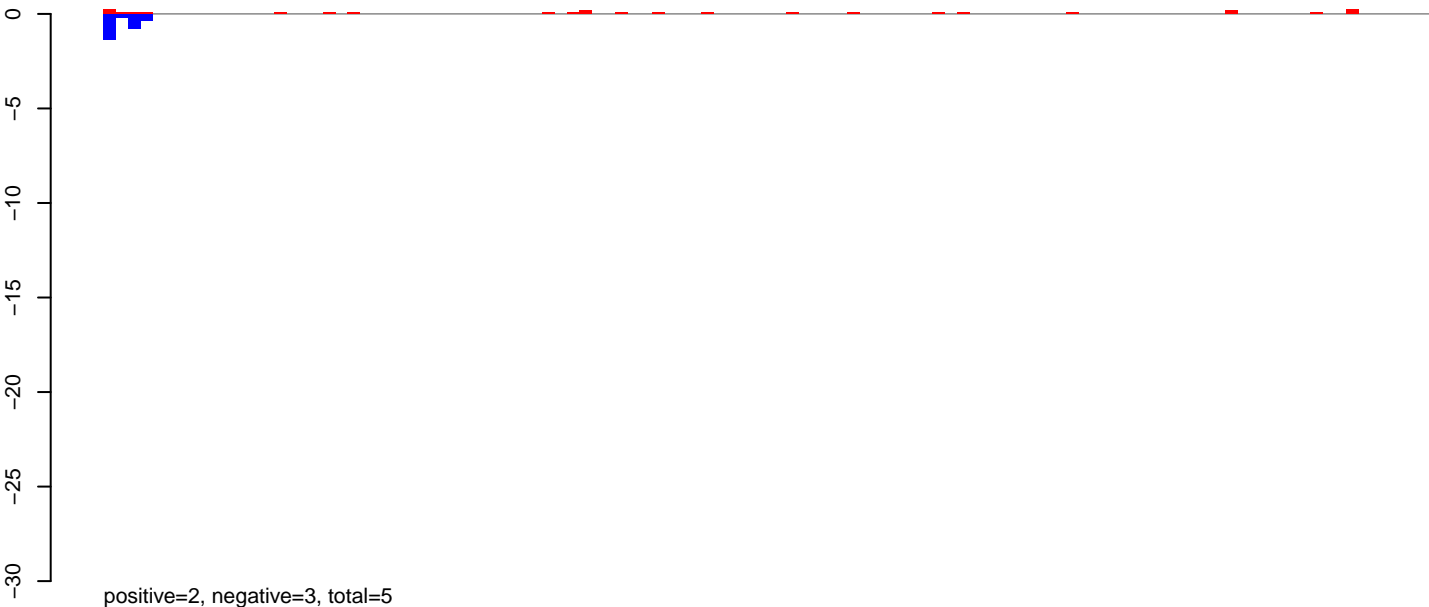


positive=0, negative=9, total=9

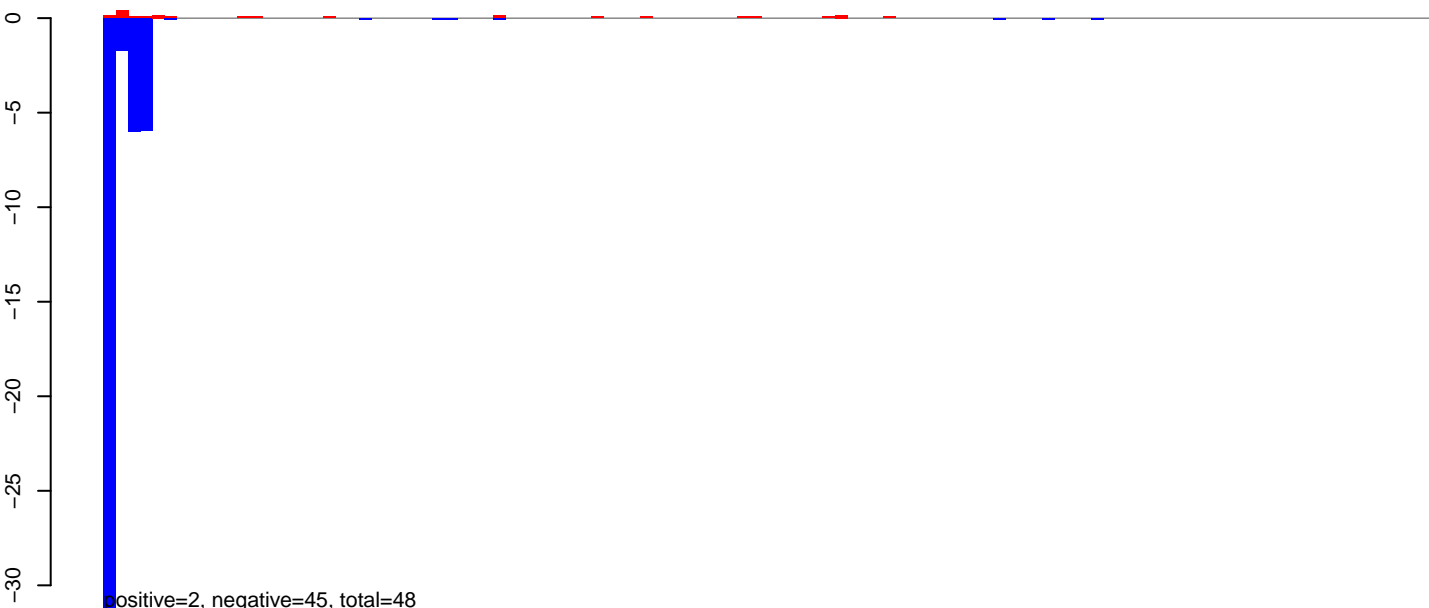
Window size=50, length=4282, TE@COPIA1-LTR-I_AG:1-4282

0 1000 2000 3000 4000

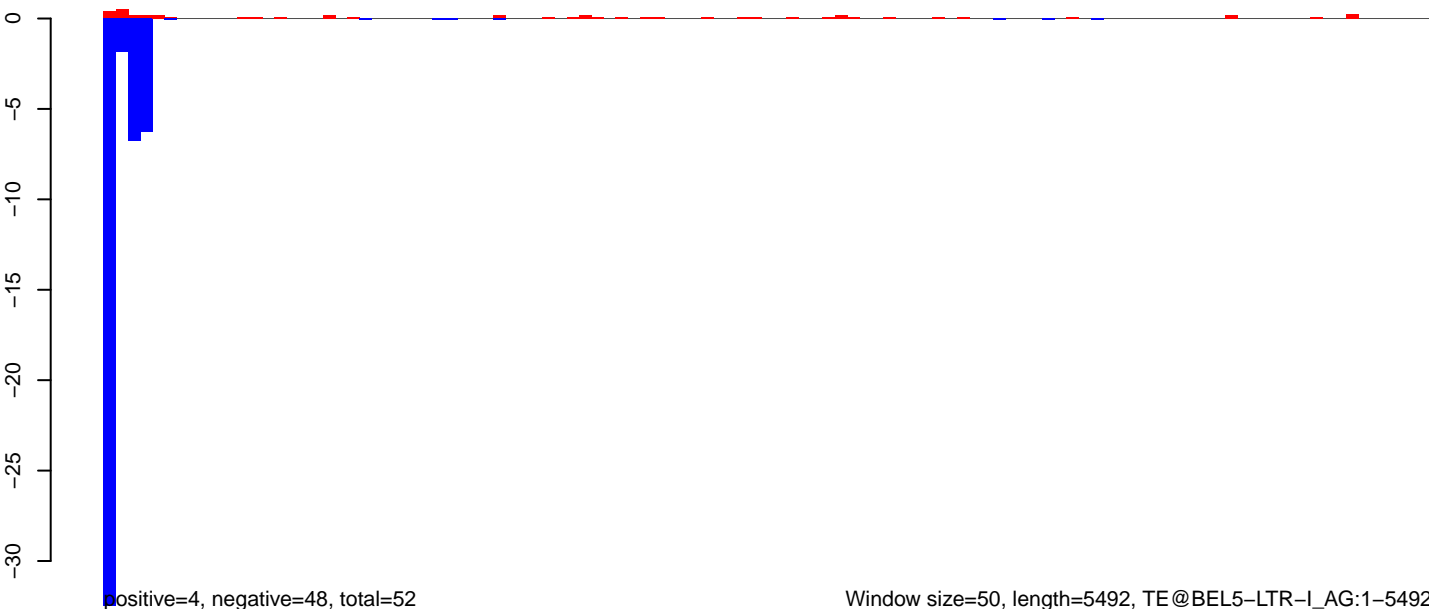
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



Window size=50, length=5492, TE@BEL5-LTR-I_AG:1-5492

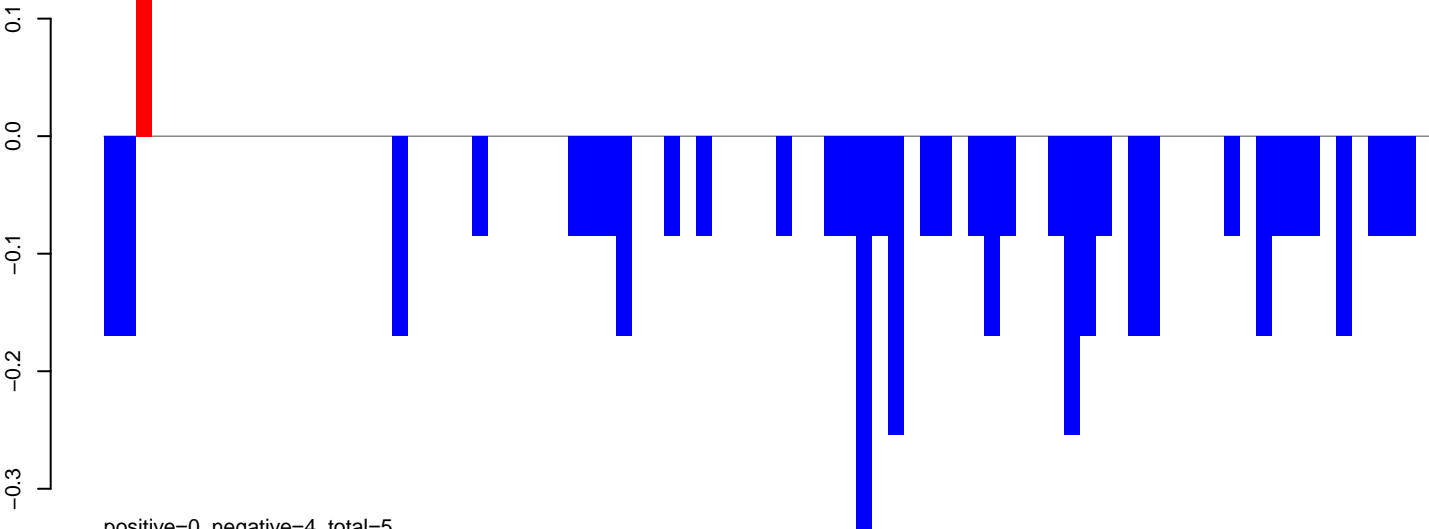
0 1000 2000 3000 4000 5000

AnGam_Mos55_cells.18_23.rep



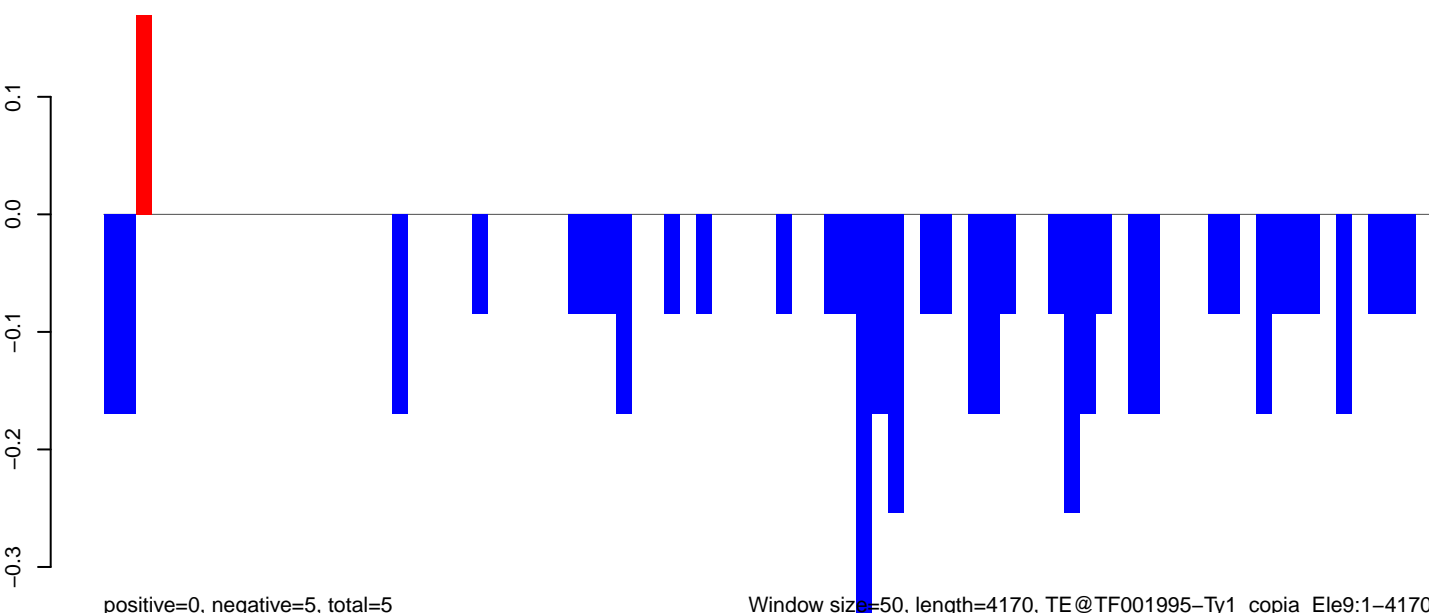
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=4, total=5

AnGam_Mos55_cells.rep

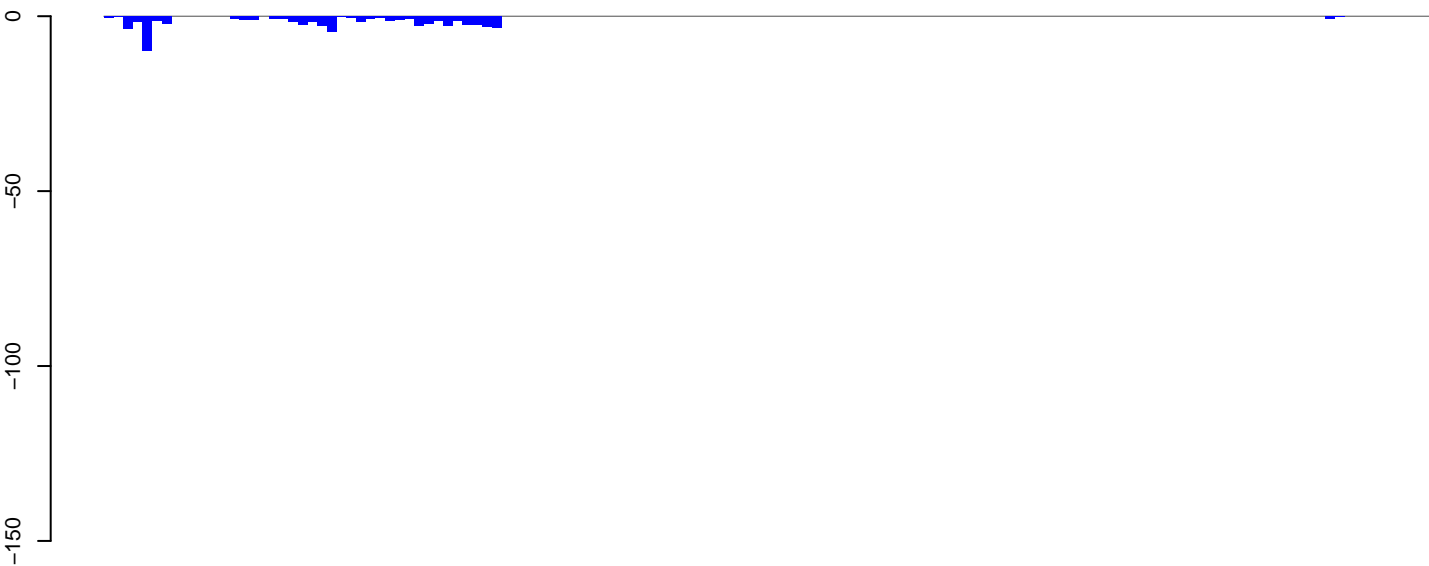


positive=0, negative=5, total=5

Window size=50, length=4170, TE@TF001995-Ty1_copia_Ele9:1-4170

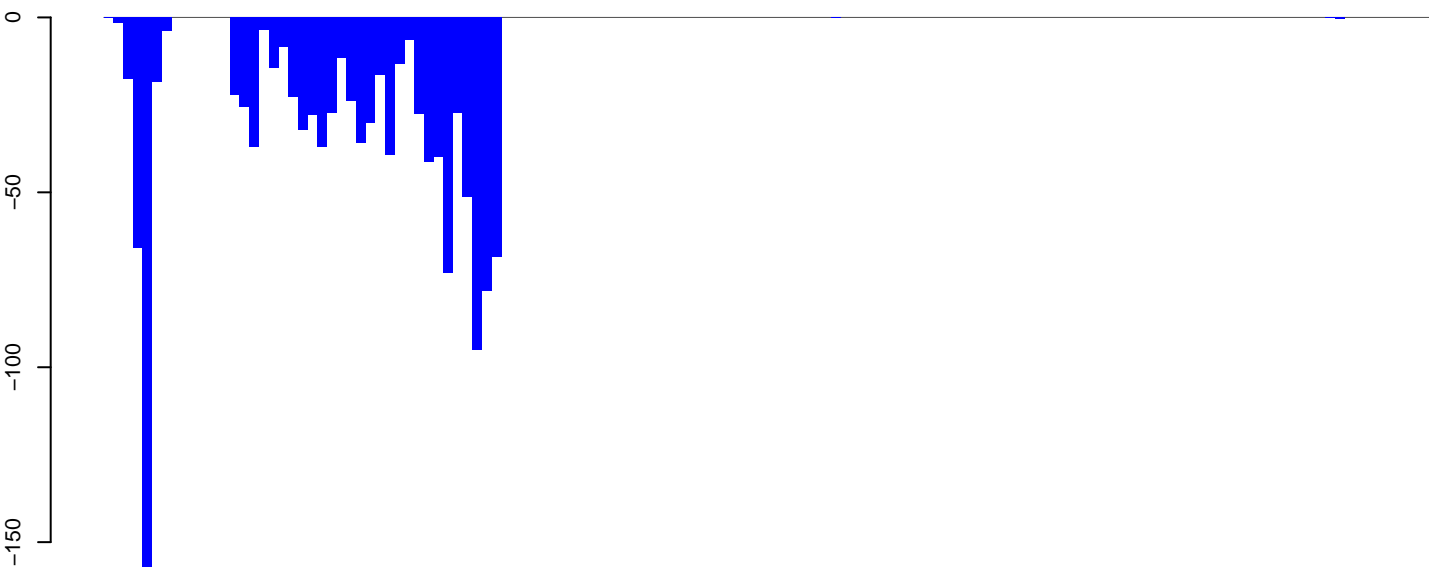
0 1000 2000 3000 4000

AnGam_Mos55_cells.18_23.rep



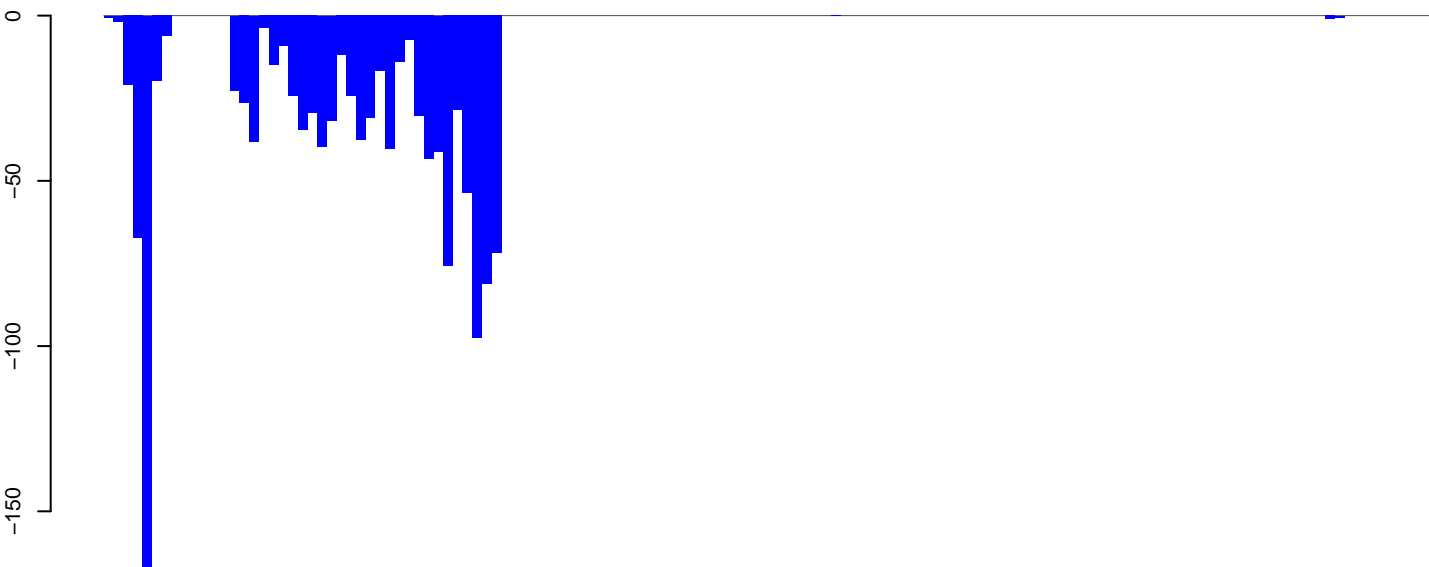
positive=0, negative=64, total=64

AnGam_Mos55_cells.24_35.rep



positive=0, negative=1217, total=1217

AnGam_Mos55_cells.rep

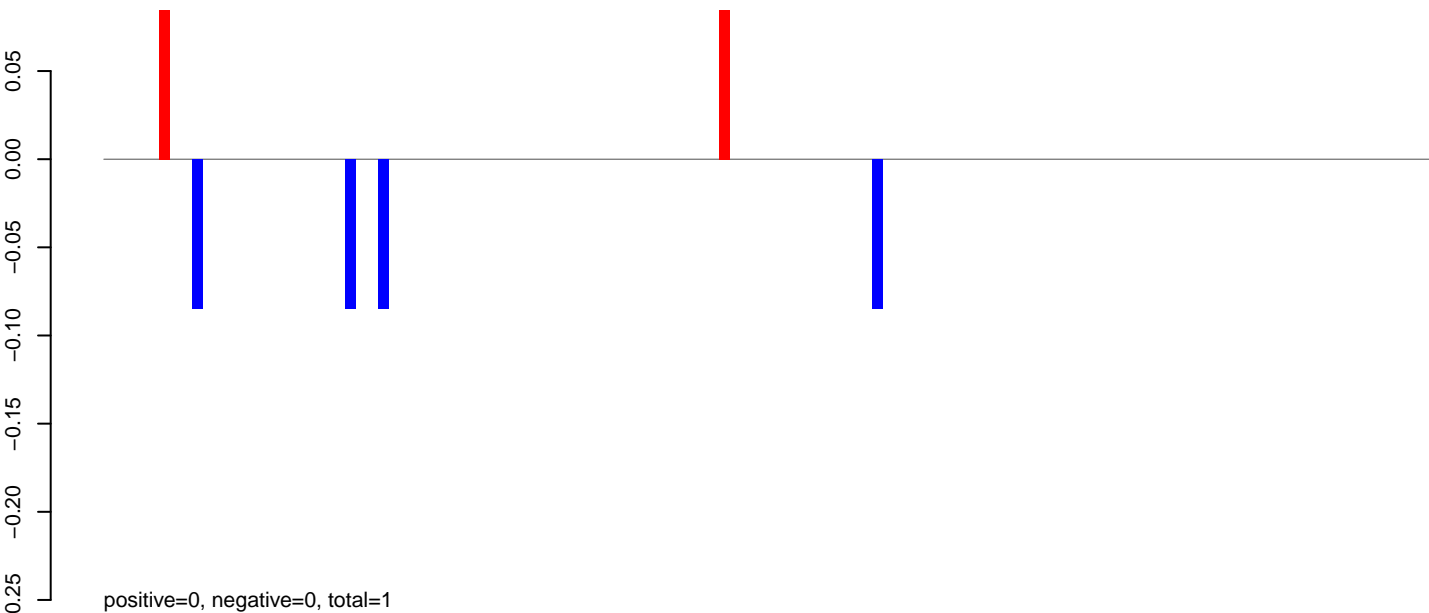


positive=0, negative=1281, total=1281

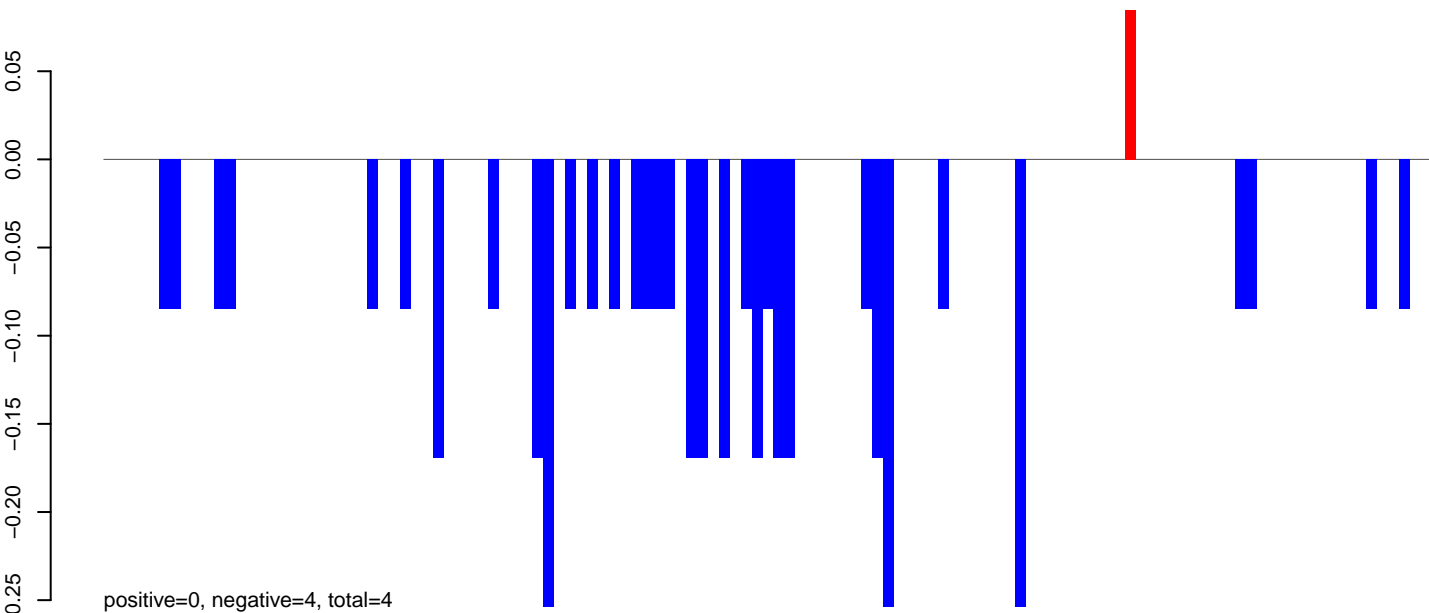
Window size=50, length=6861, TE@GYPSY7-LTR-I_AG:1-6861

0 1000 2000 3000 4000 5000 6000 7000

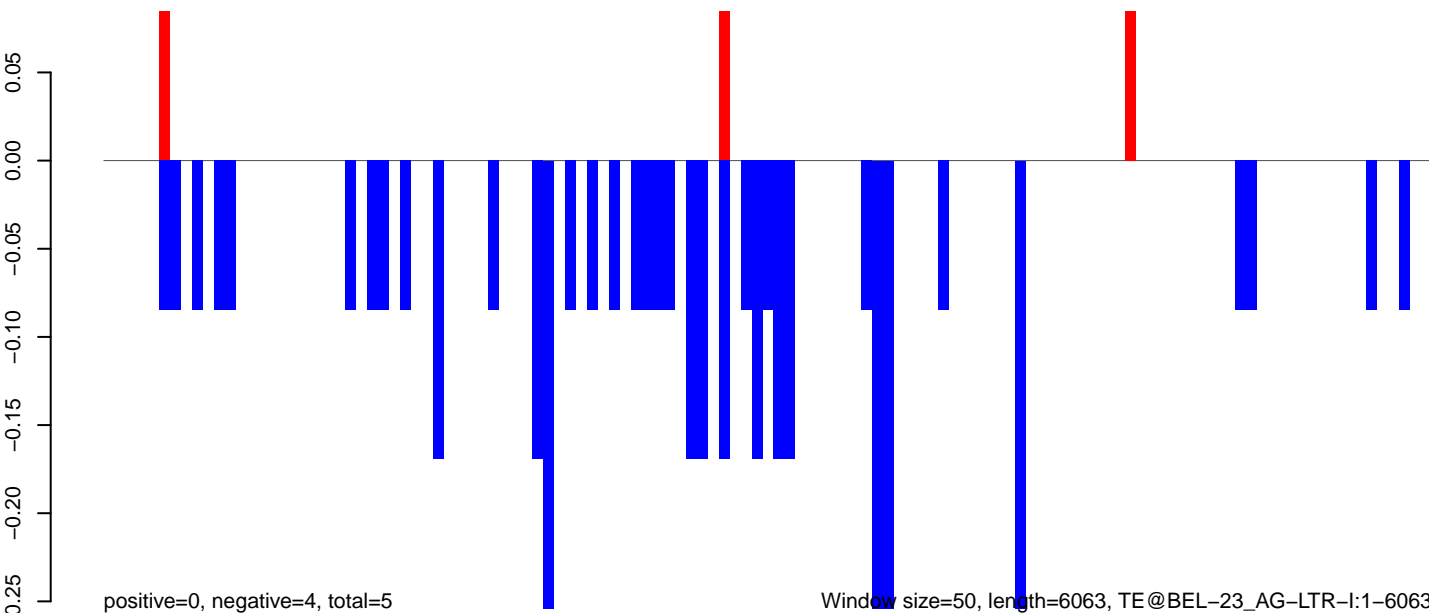
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



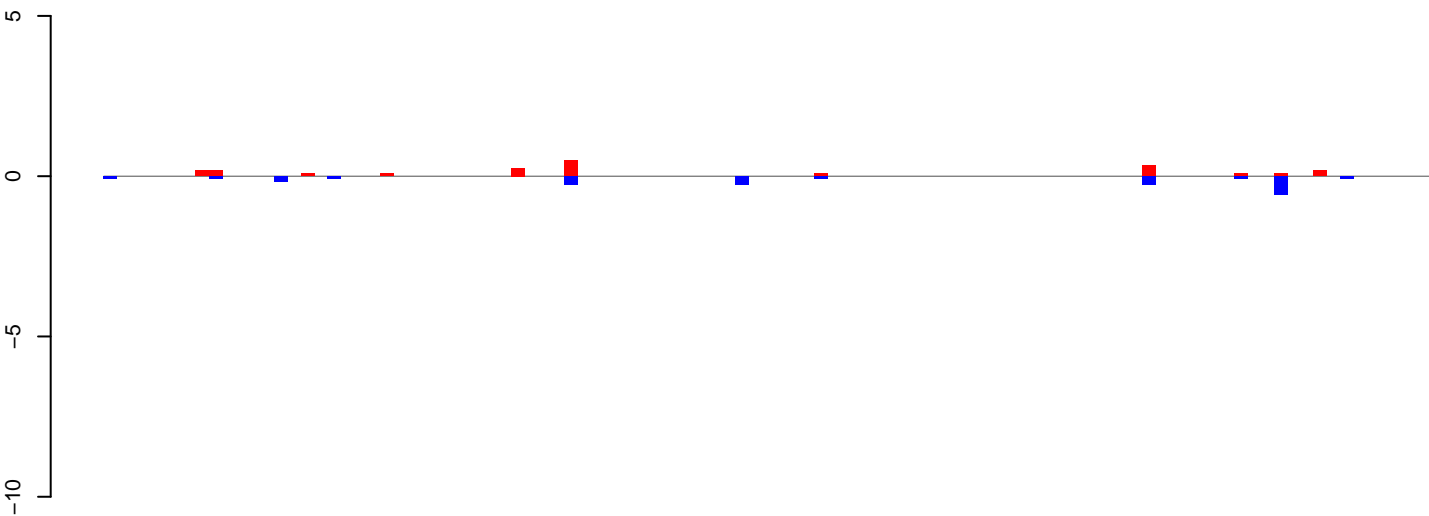
AnGam_Mos55_cells.rep



Window size=50, length=6063, TE@BEL-23_AG-LTR-I:1-6063

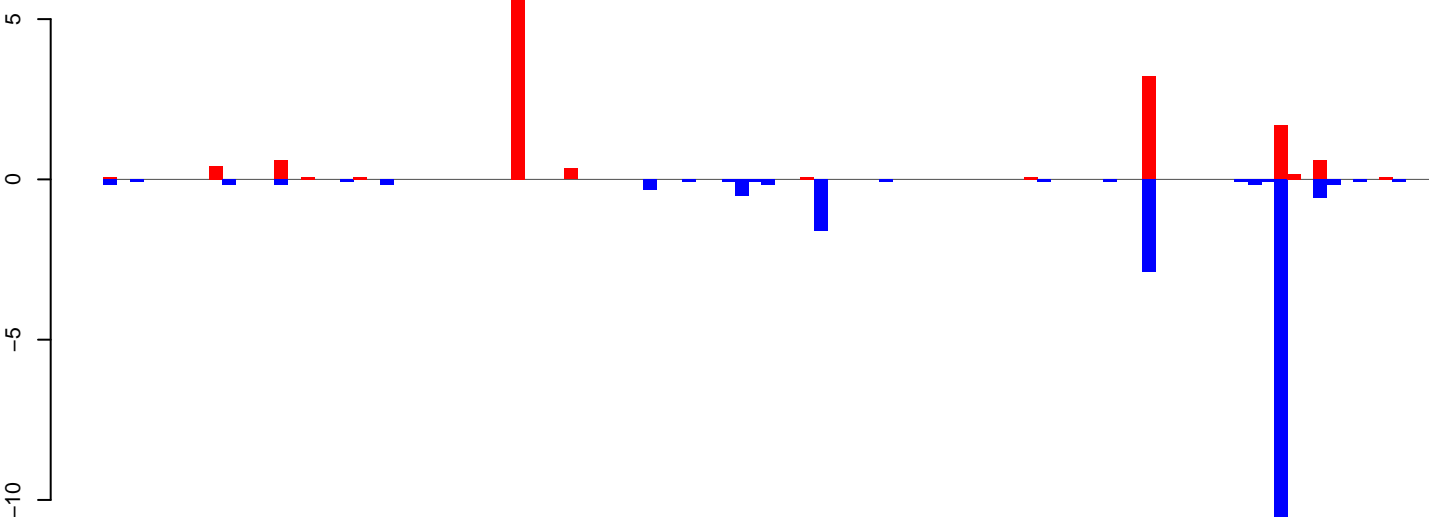
0 1000 2000 3000 4000 5000 6000

AnGam_Mos55_cells.18_23.rep



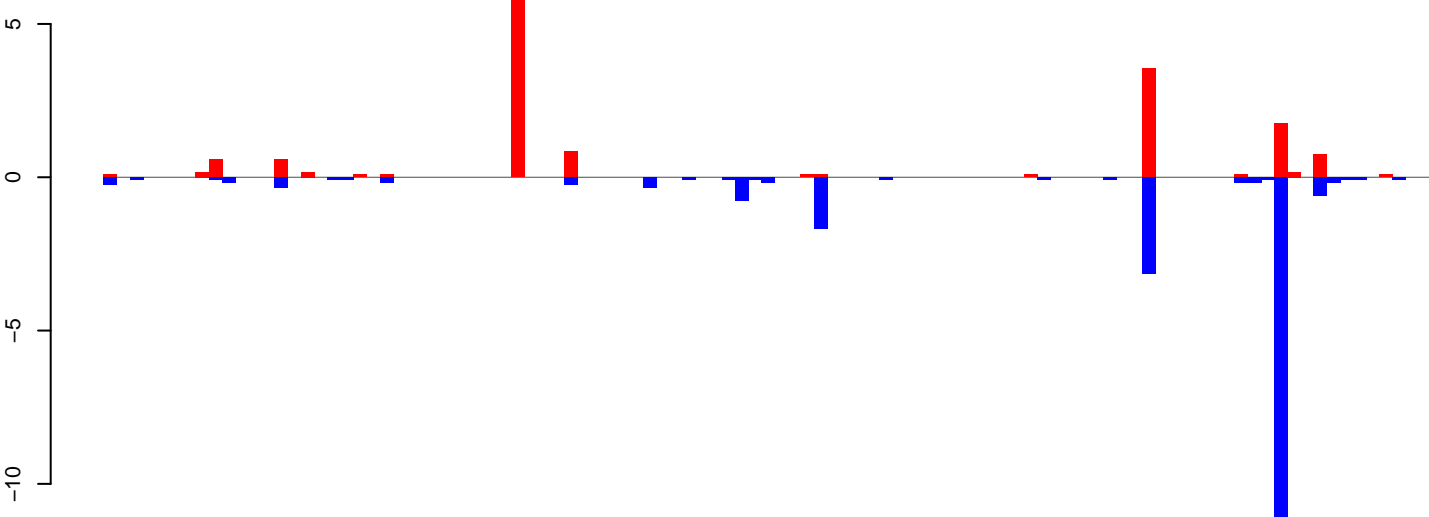
positive=2, negative=2, total=4

AnGam_Mos55_cells.24_35.rep



positive=13, negative=21, total=34

AnGam_Mos55_cells.rep

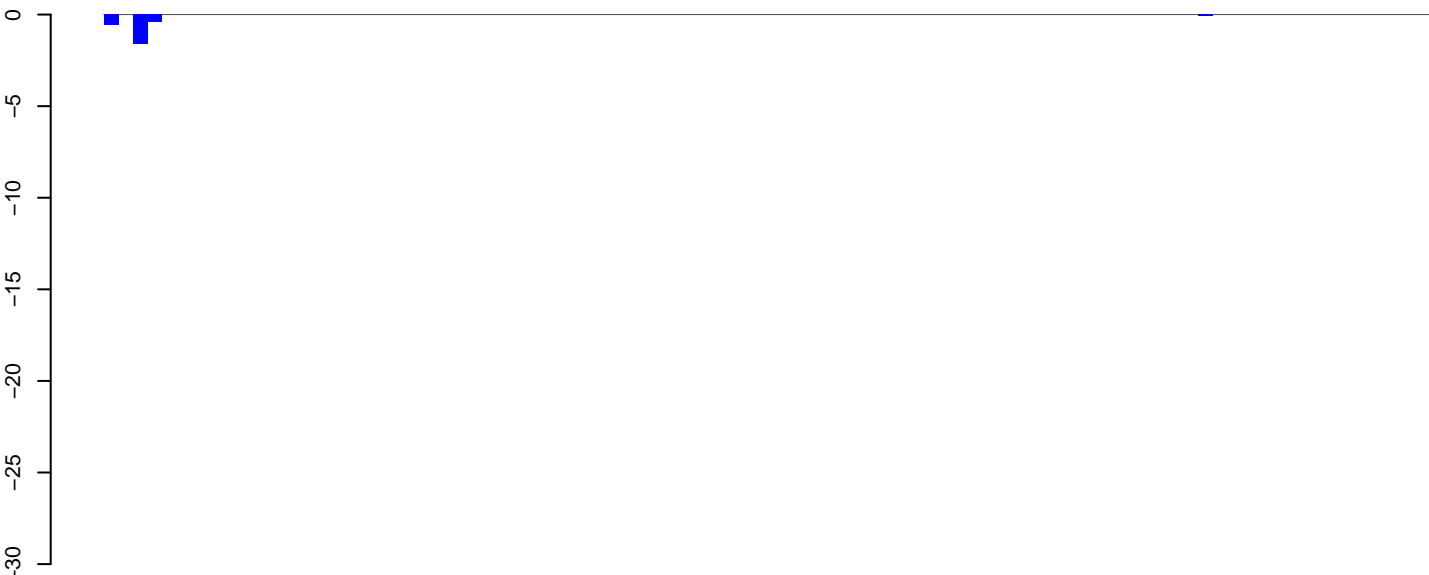


positive=15, negative=23, total=38

Window size=50, length=5083, TE@GYPSY69-LTR-L_AG:1-5083

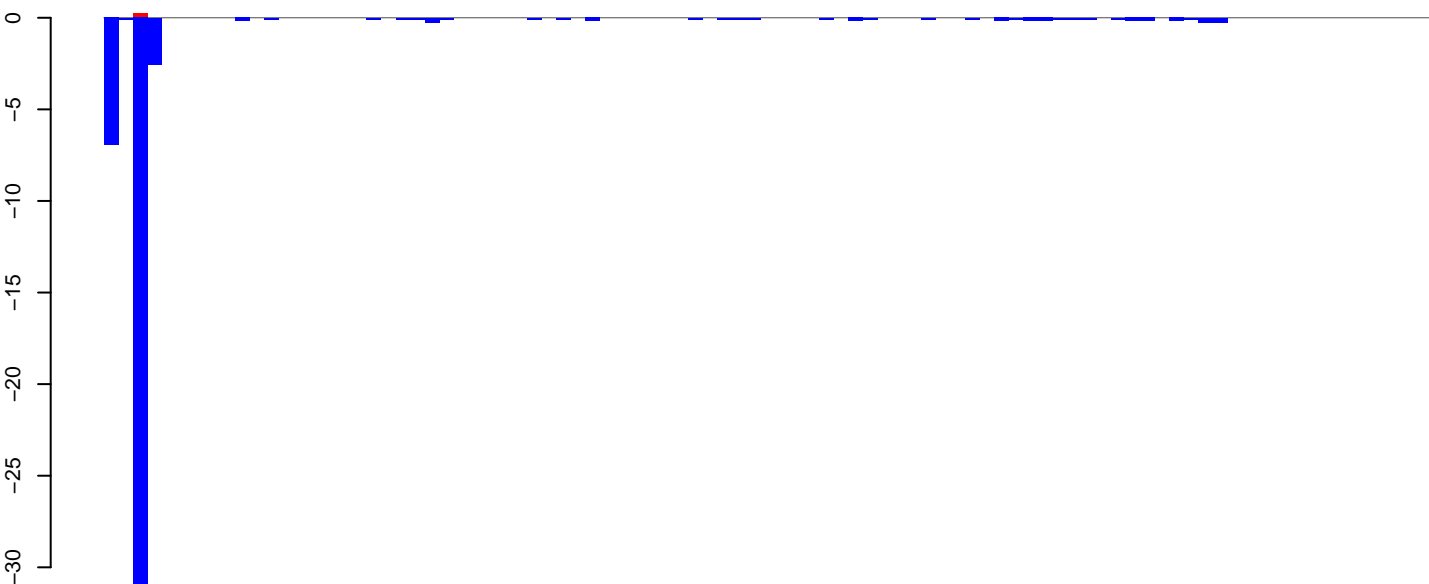
0 1000 2000 3000 4000 5000

AnGam_Mos55_cells.18_23.rep



positive=0, negative=3, total=3

AnGam_Mos55_cells.24_35.rep



positive=0, negative=46, total=46

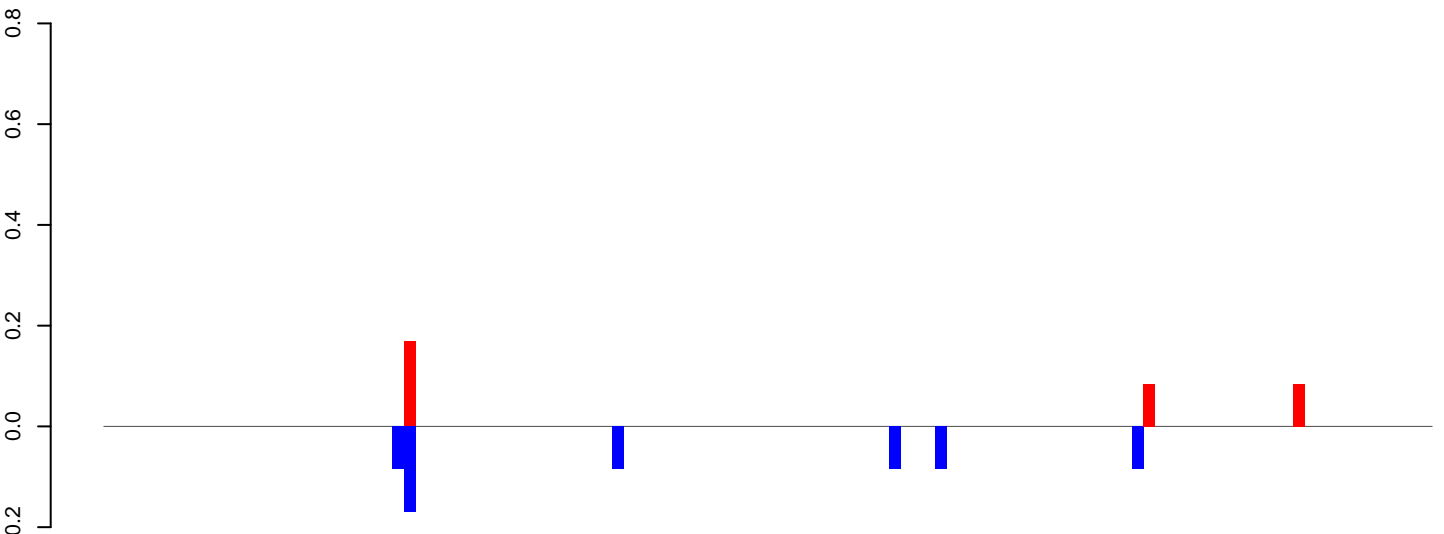
AnGam_Mos55_cells.rep



positive=0, negative=49, total=49

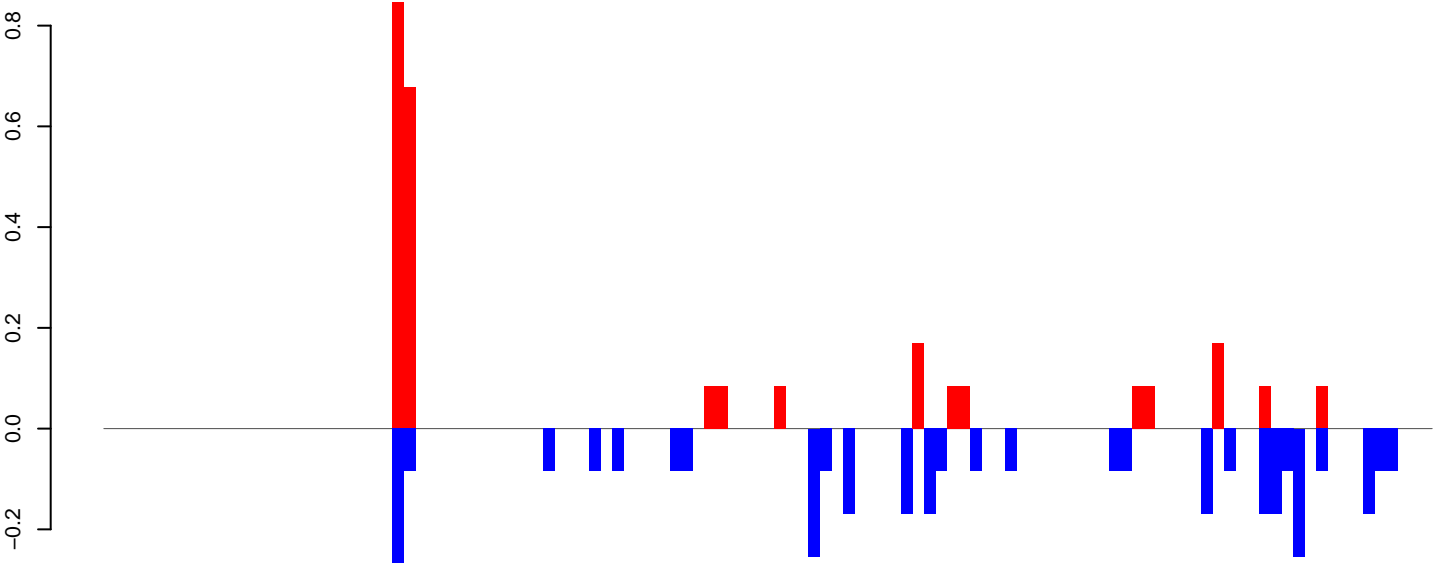
Window size=50, length=4560, TE@GYPSY62-LTR-I_AG:1-4560

AnGam_Mos55_cells.18_23.rep



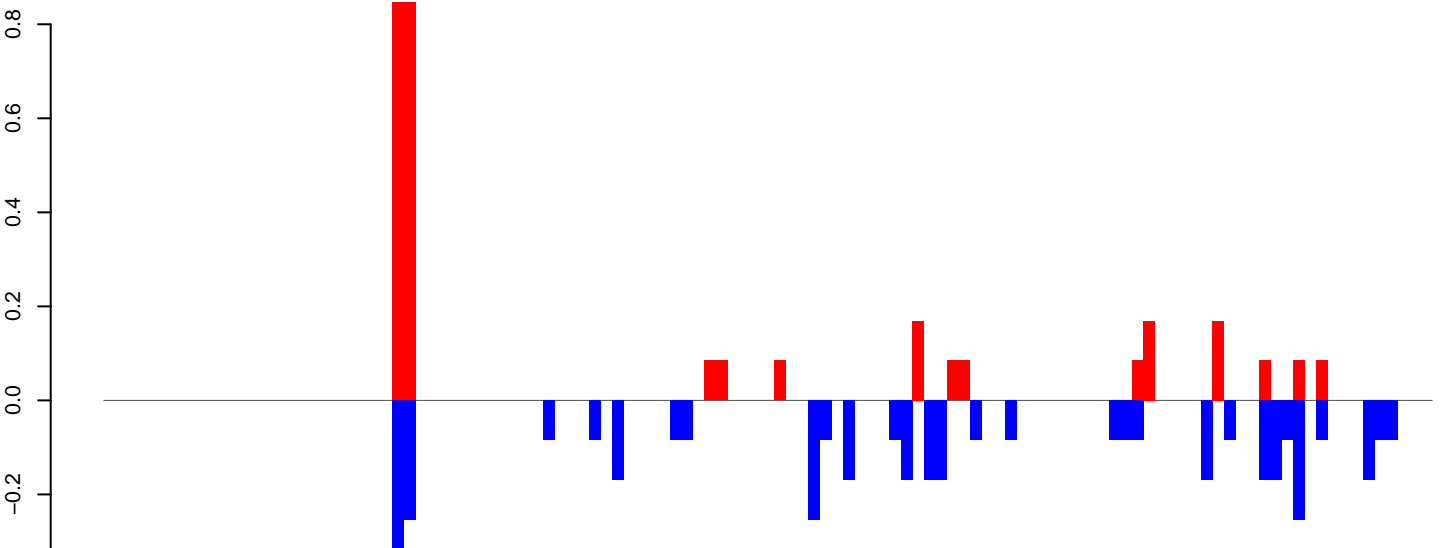
positive=0, negative=1, total=1

AnGam_Mos55_cells.24_35.rep



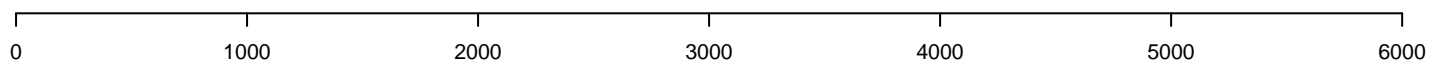
positive=3, negative=3, total=6

AnGam_Mos55_cells.rep

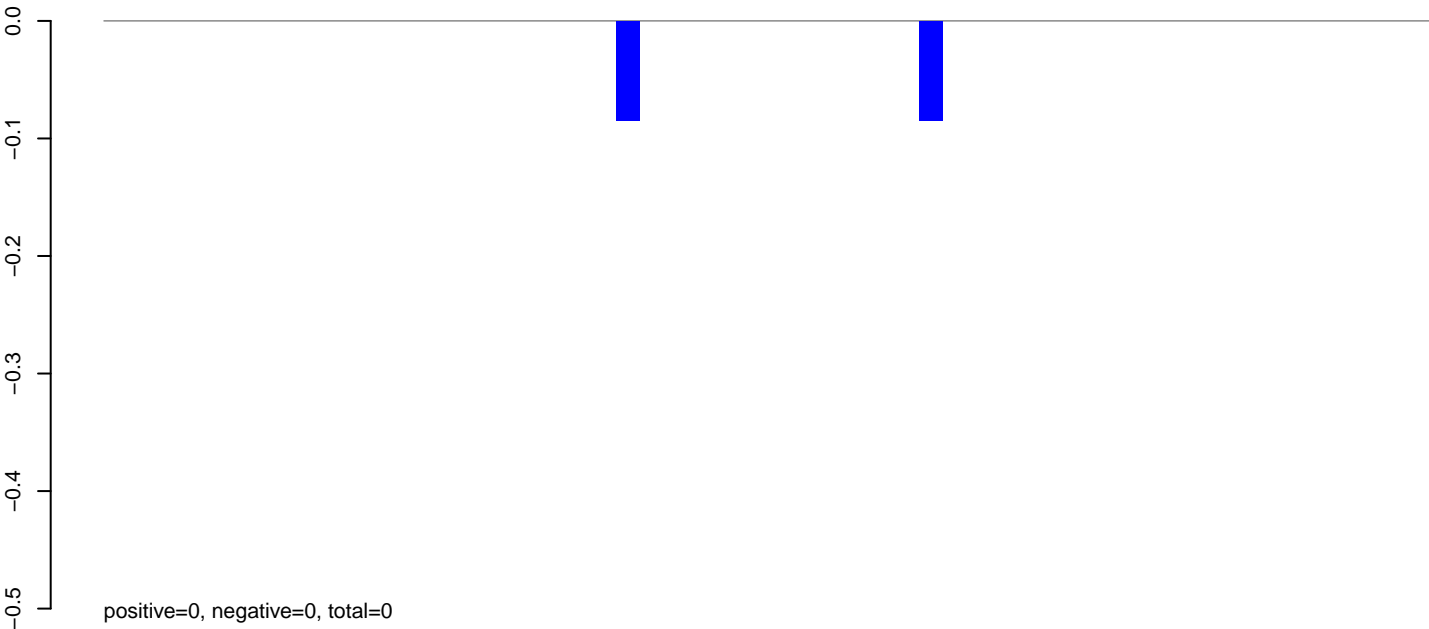


positive=3, negative=4, total=7

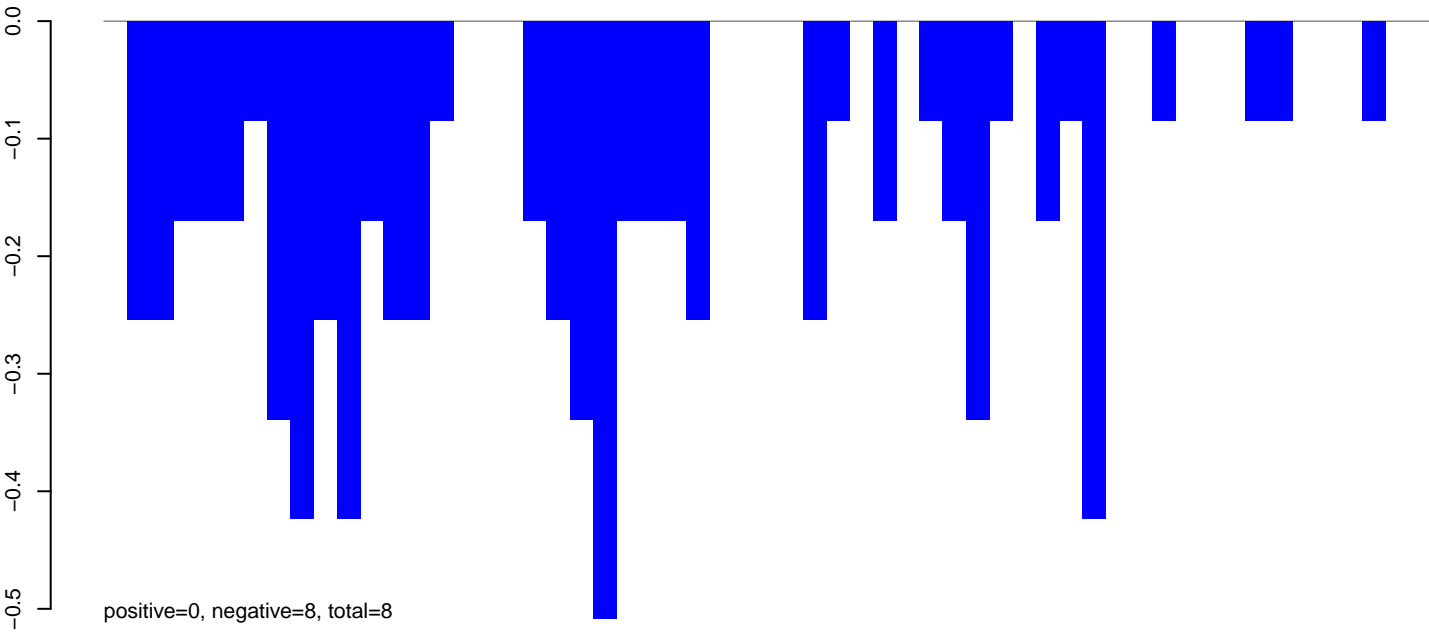
Window size=50, length=5773, TE@GYPSY49-LTR-L_AG:1-5773



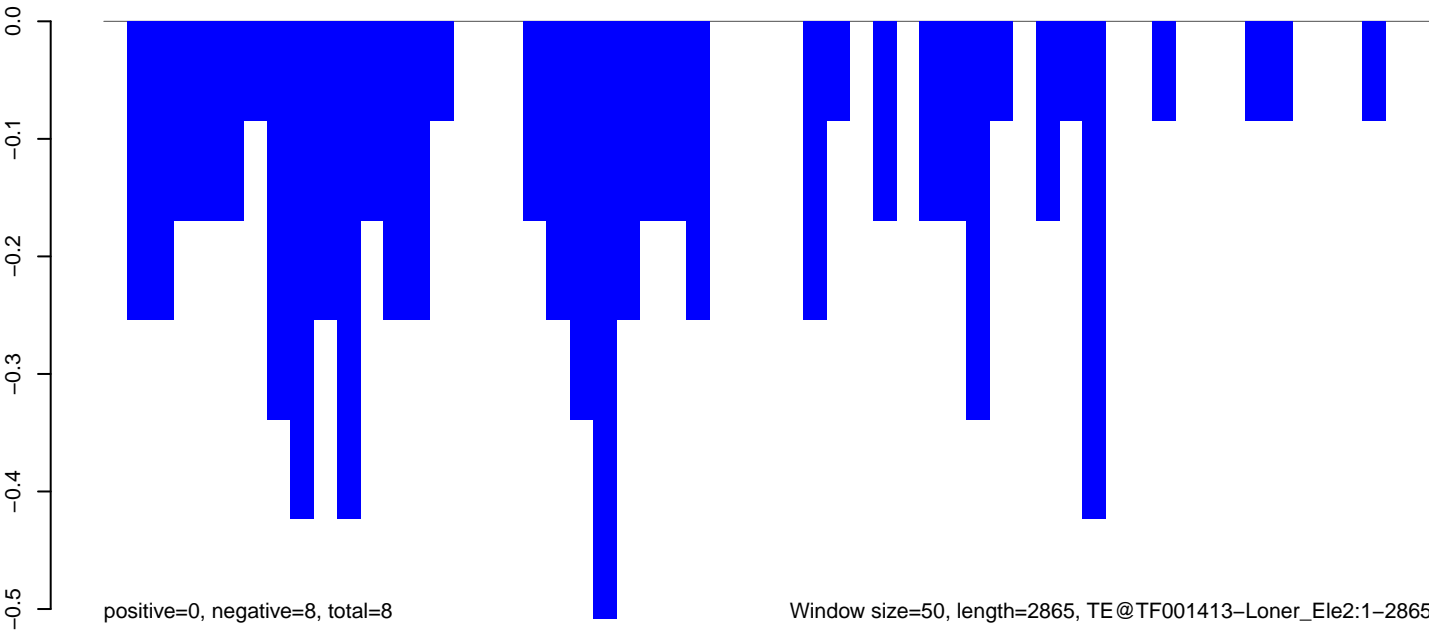
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

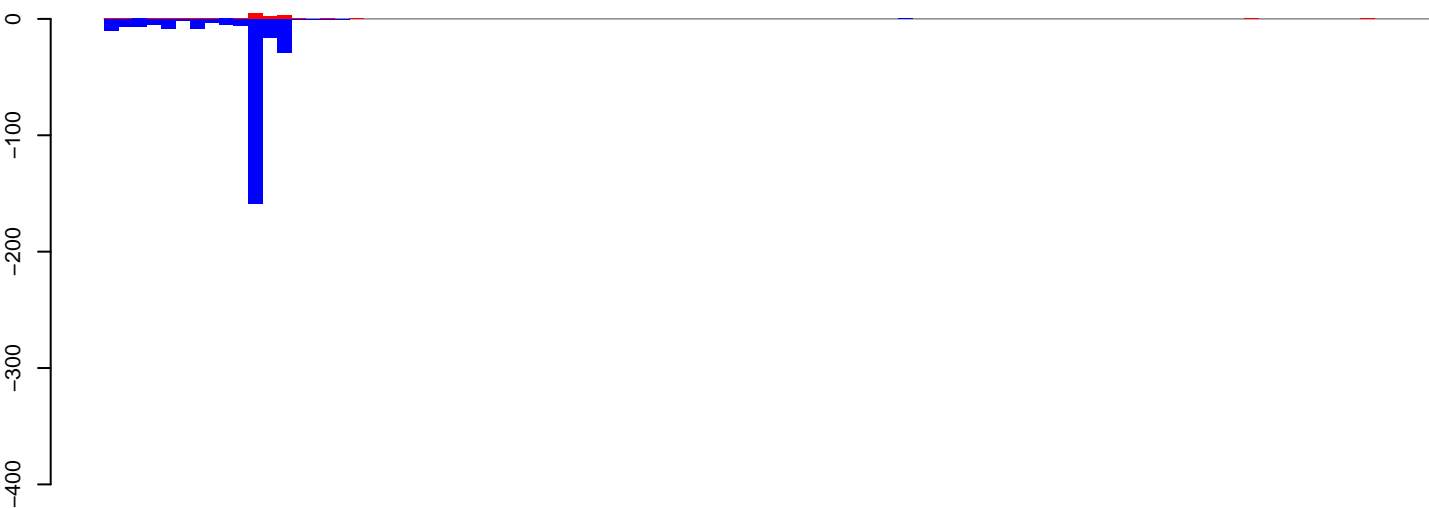


AnGam_Mos55_cells.rep



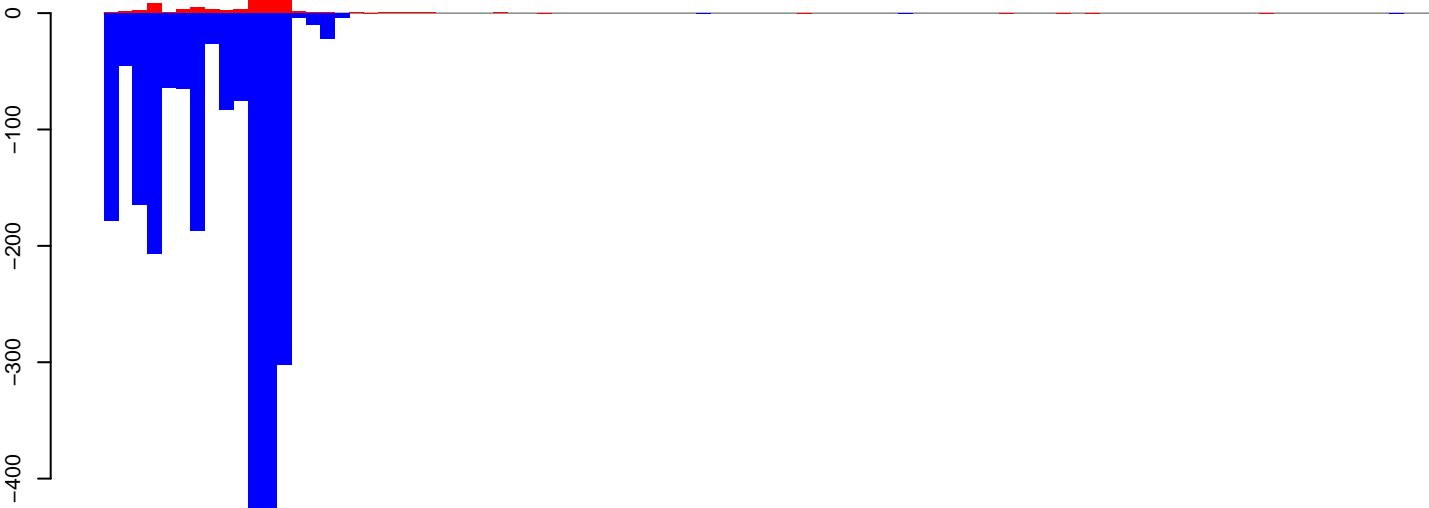
Window size=50, length=2865, TE@TF001413-Loner_Ele2:1-2865

AnGam_Mos55_cells.18_23.rep



positive=14, negative=267, total=281

AnGam_Mos55_cells.24_35.rep



positive=118, negative=2357, total=2475

AnGam_Mos55_cells.rep

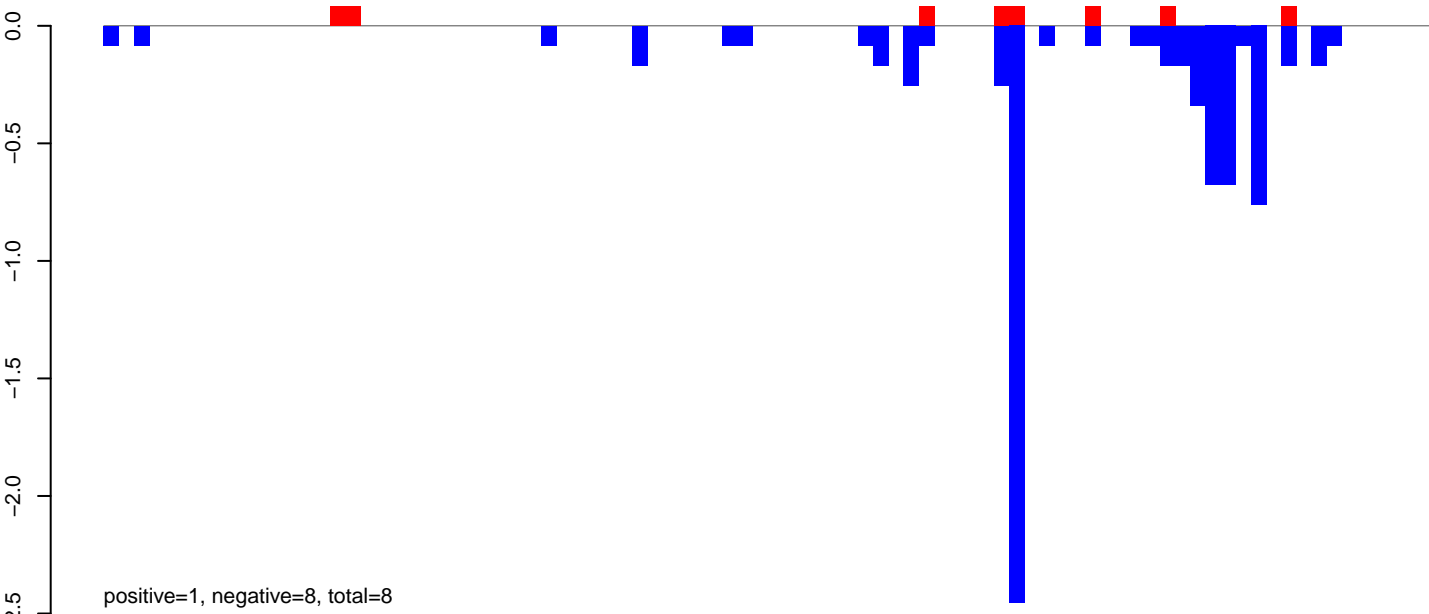


positive=132, negative=2624, total=2756

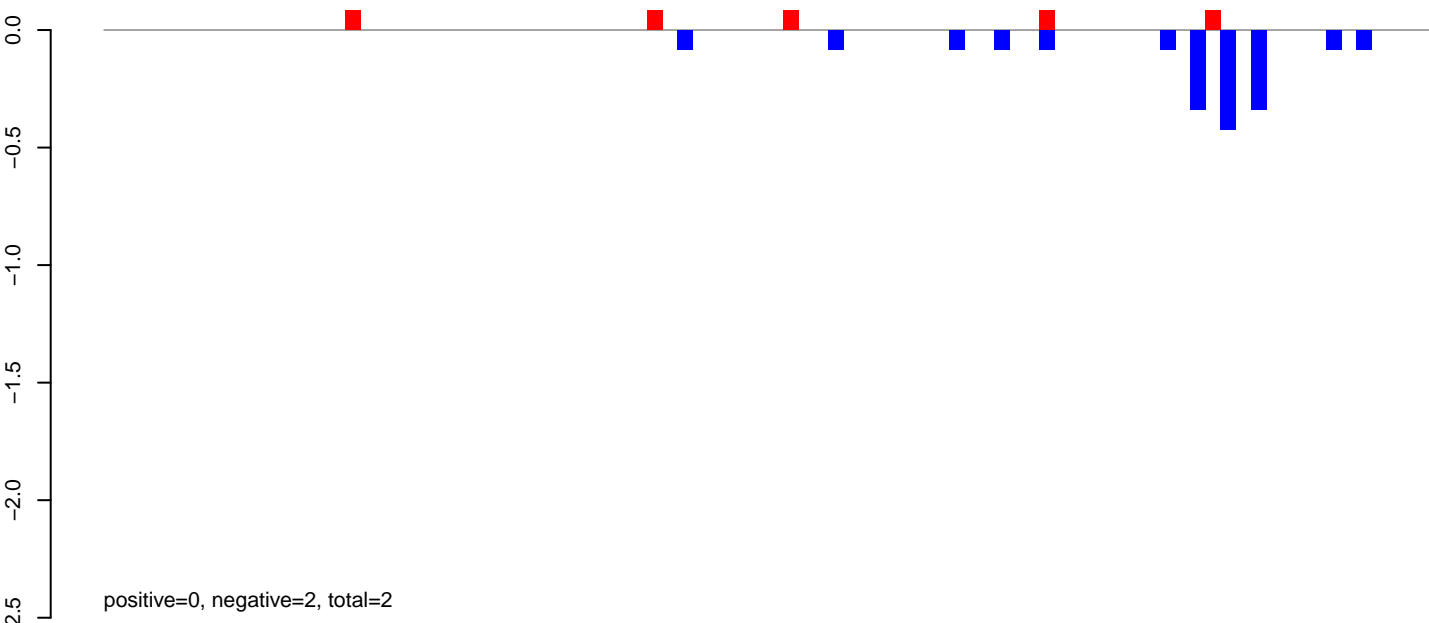
Window size=50, length=4641, TE@GYPSY35-LTR-I_AG:1-4641

0 1000 2000 3000 4000

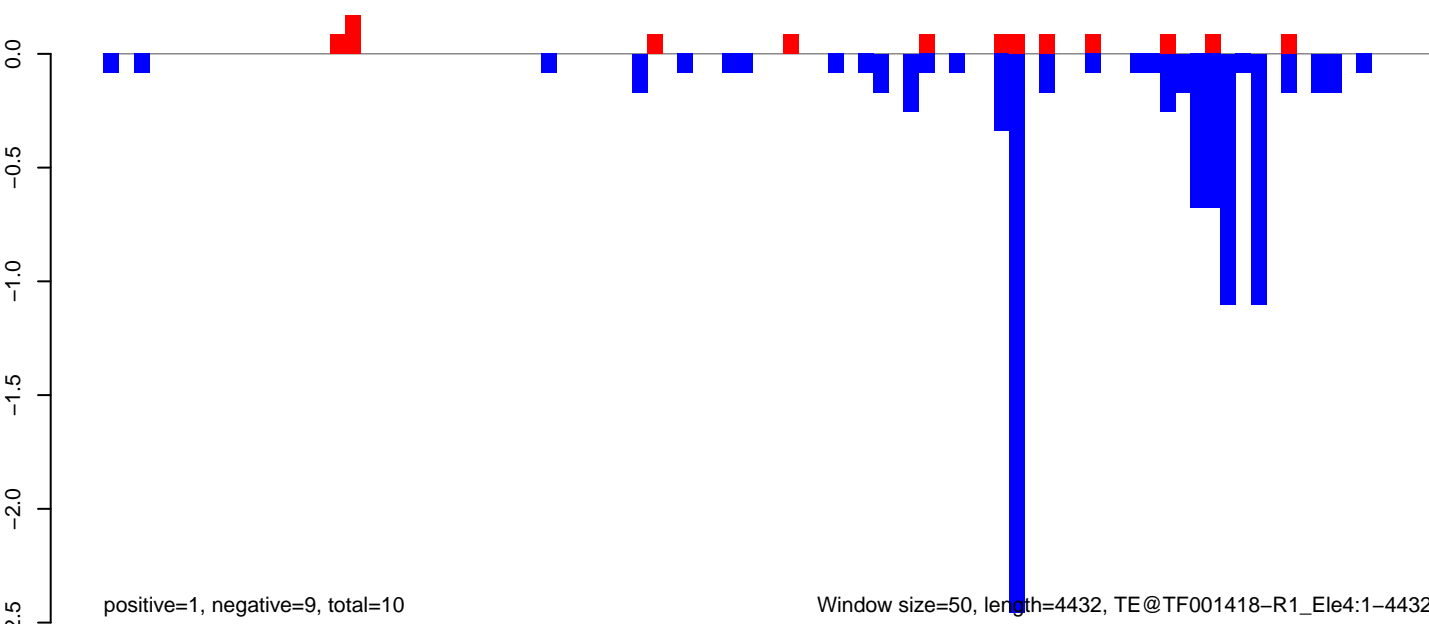
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

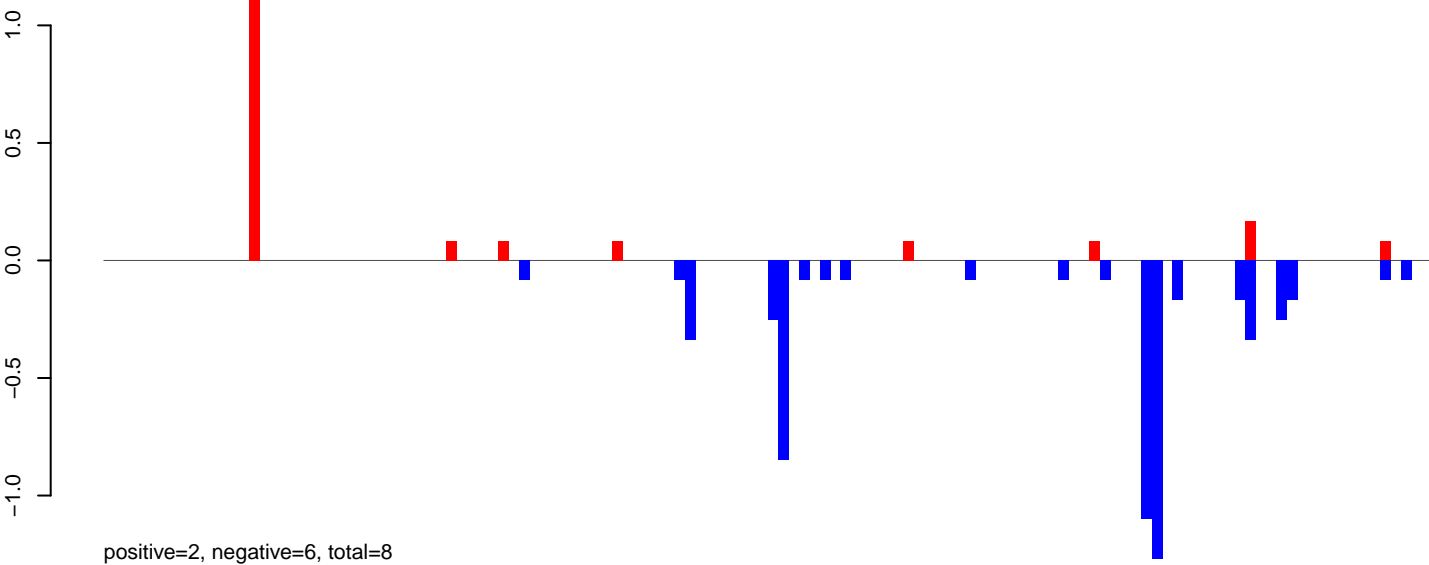


AnGam_Mos55_cells.rep

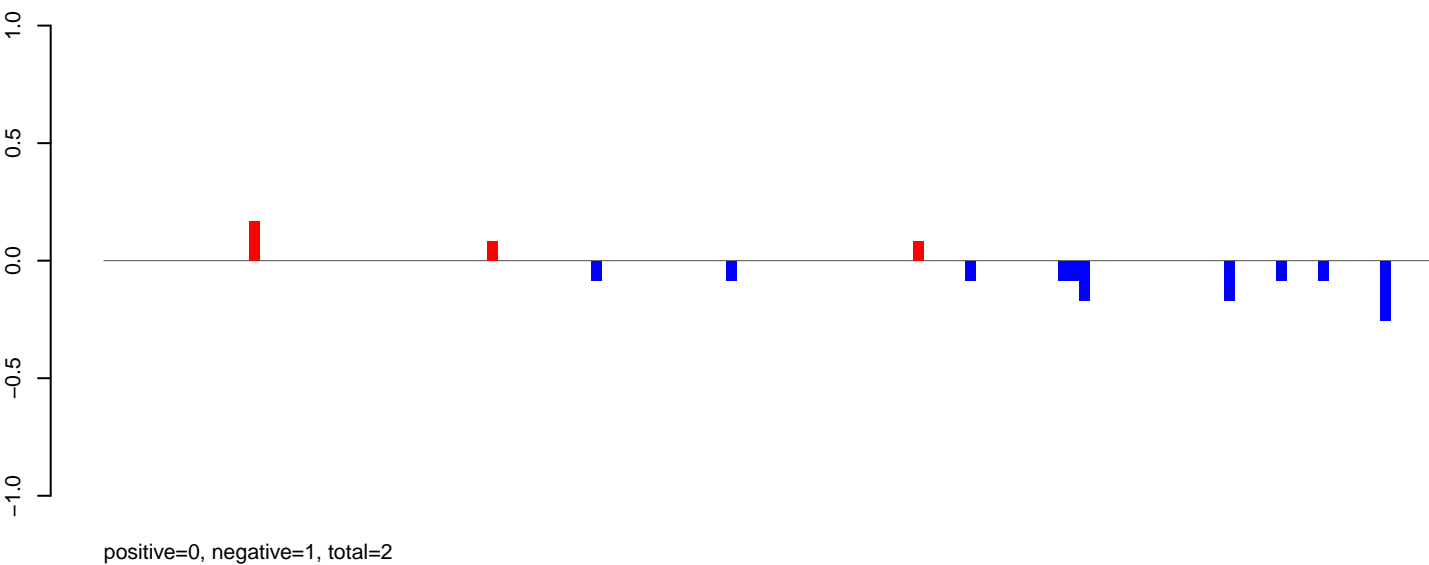


Window size=50, length=4432, TE@TF001418-R1_Ele4:1-4432

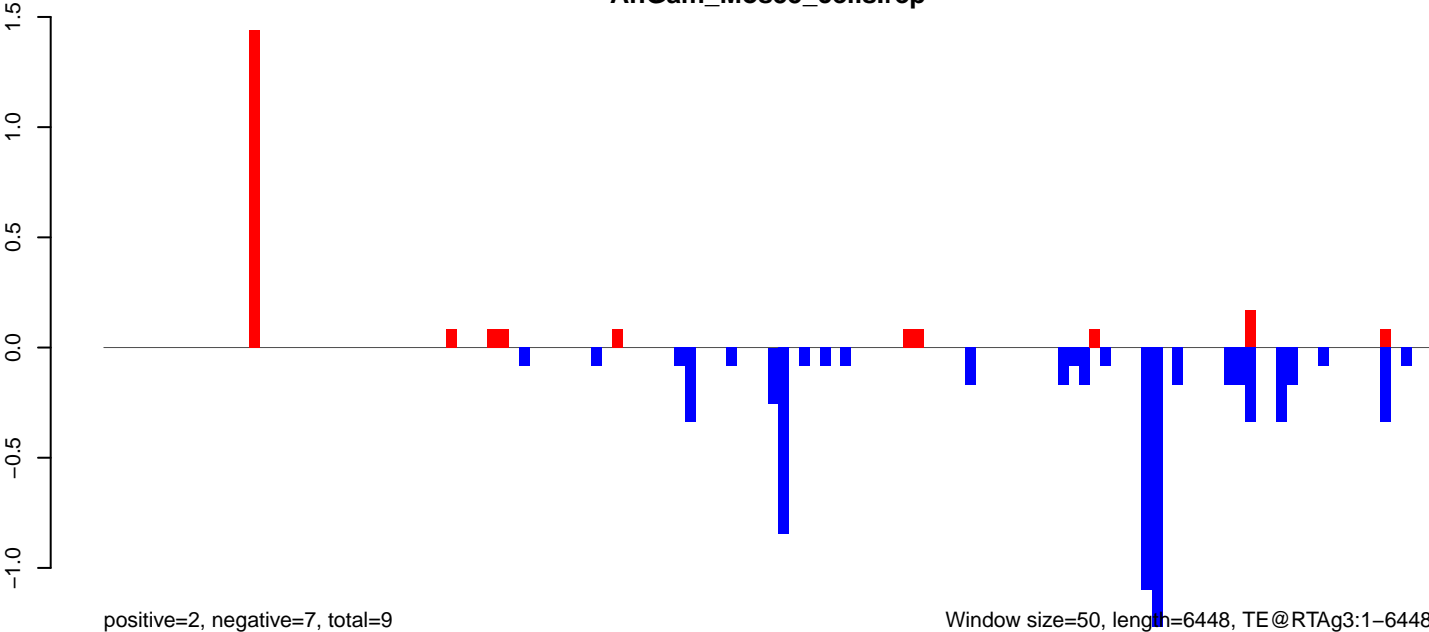
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

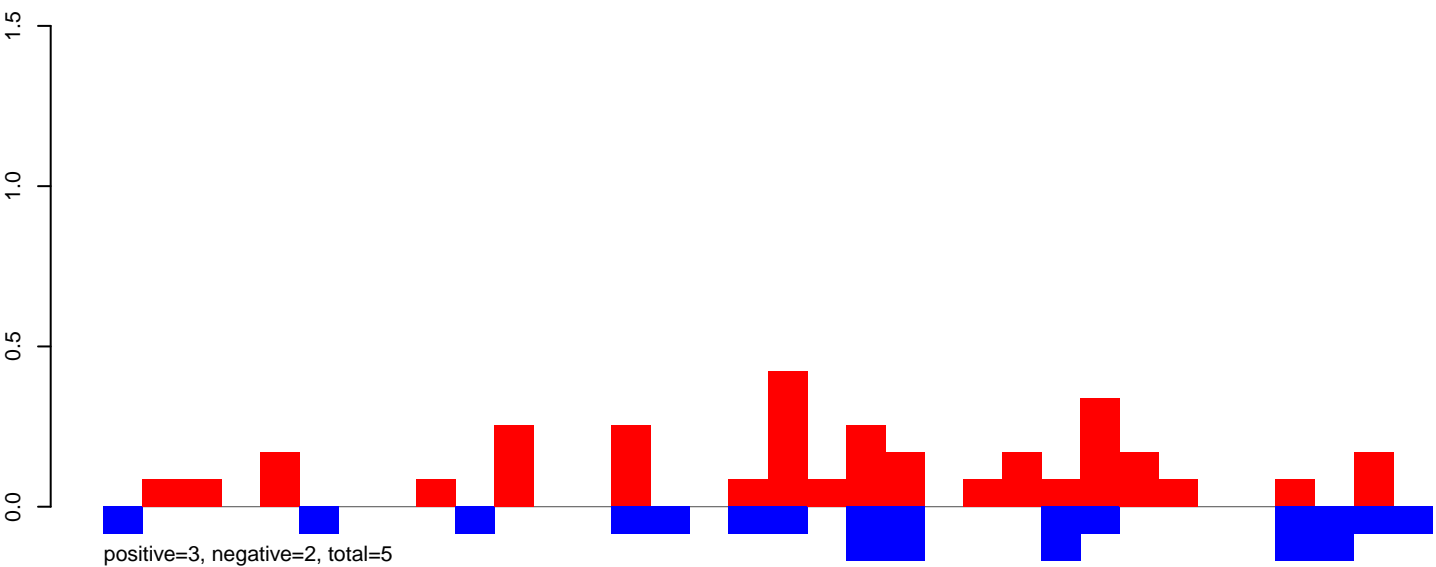


AnGam_Mos55_cells.rep

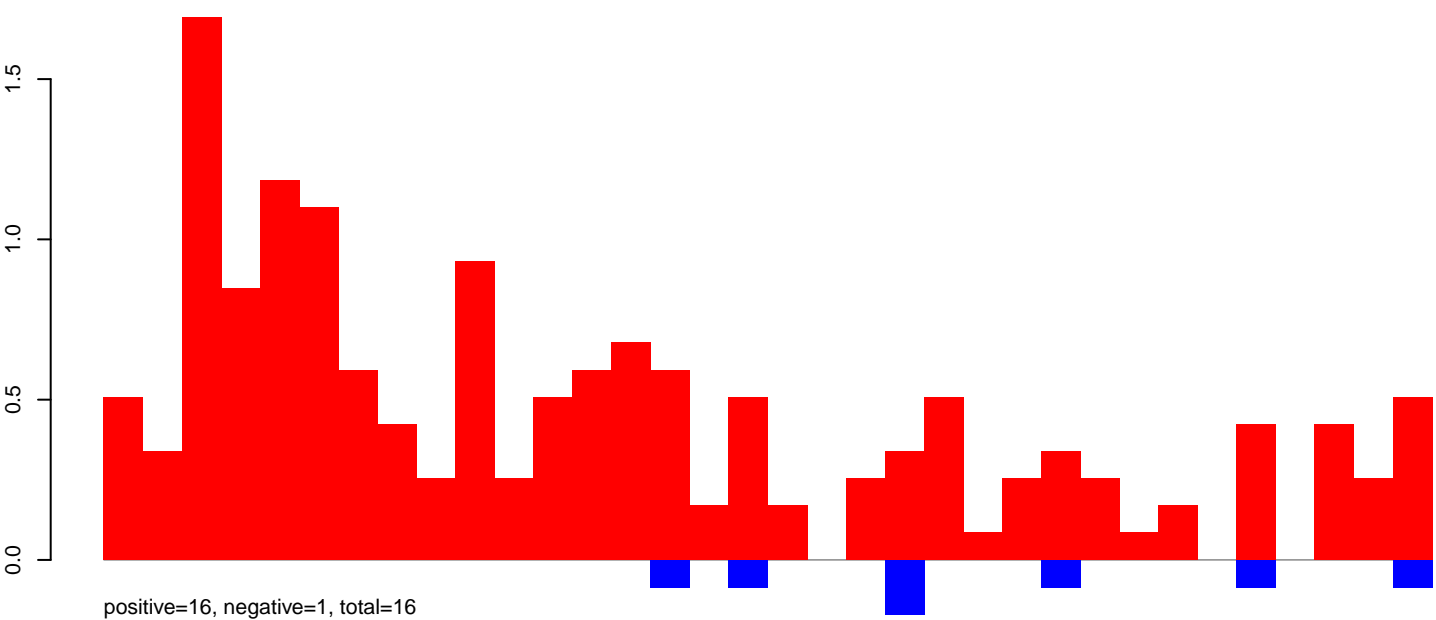


Window size=50, length=6448, TE@RTAg3:1-6448

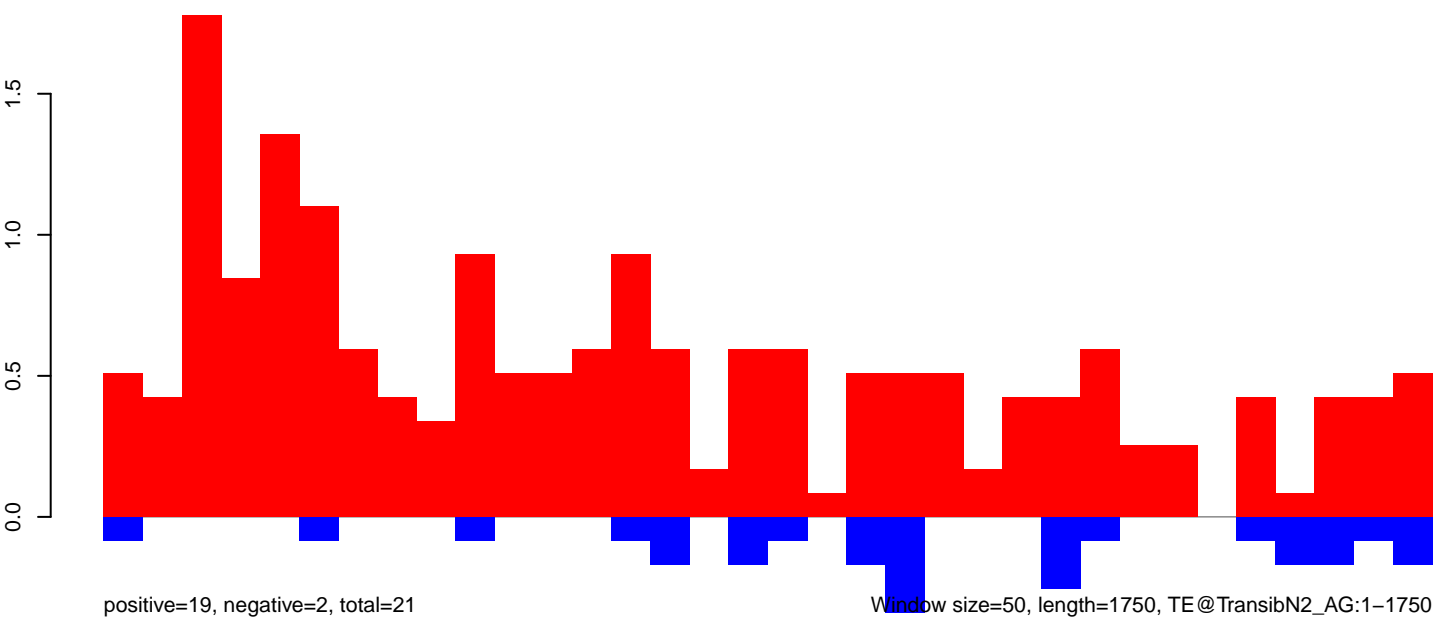
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



0

500

1000

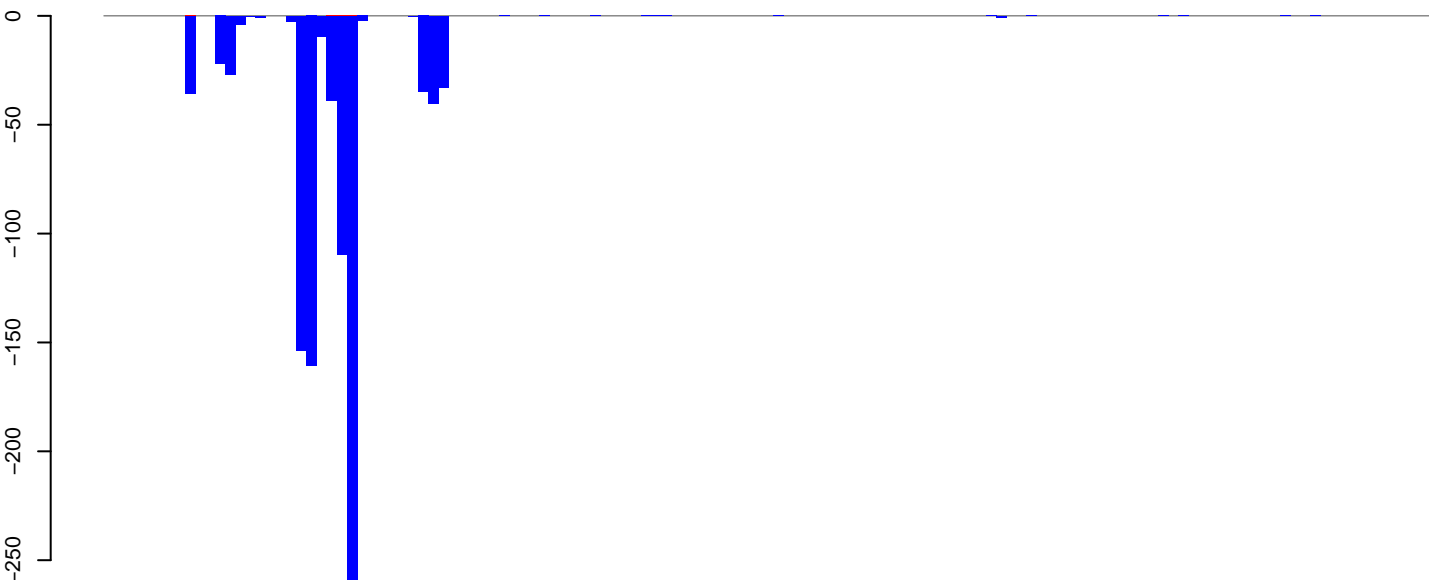
1500

AnGam_Mos55_cells.18_23.rep



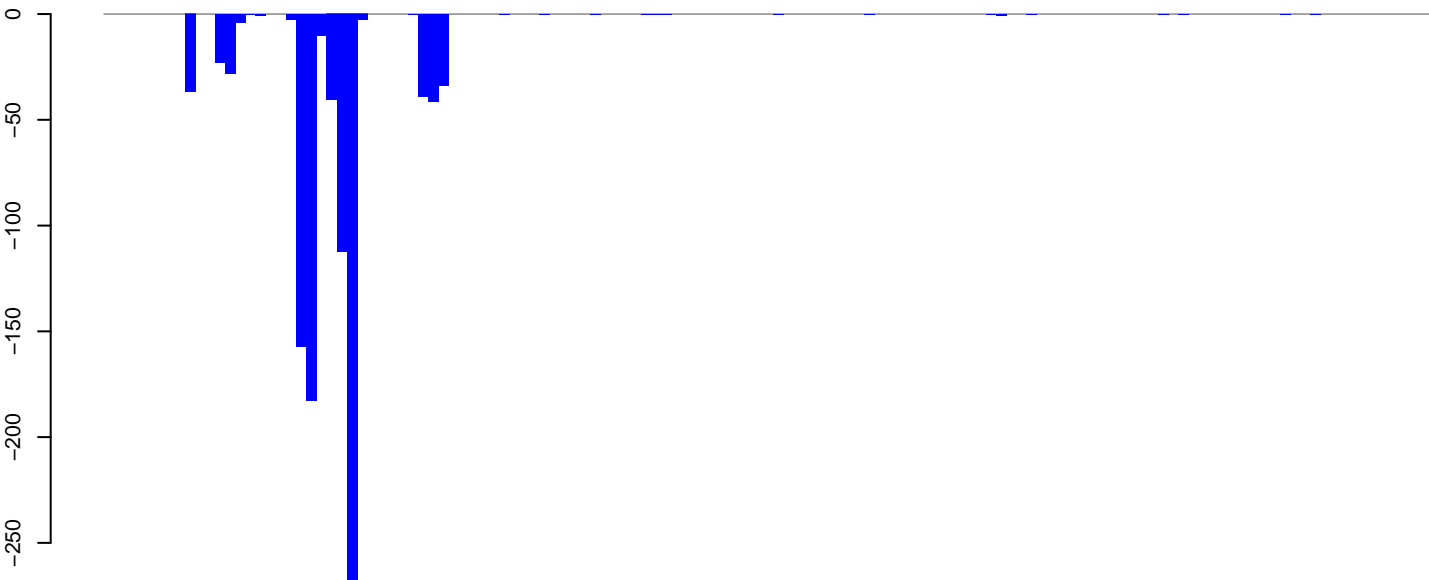
positive=0, negative=48, total=48

AnGam_Mos55_cells.24_35.rep



positive=1, negative=952, total=952

AnGam_Mos55_cells.rep

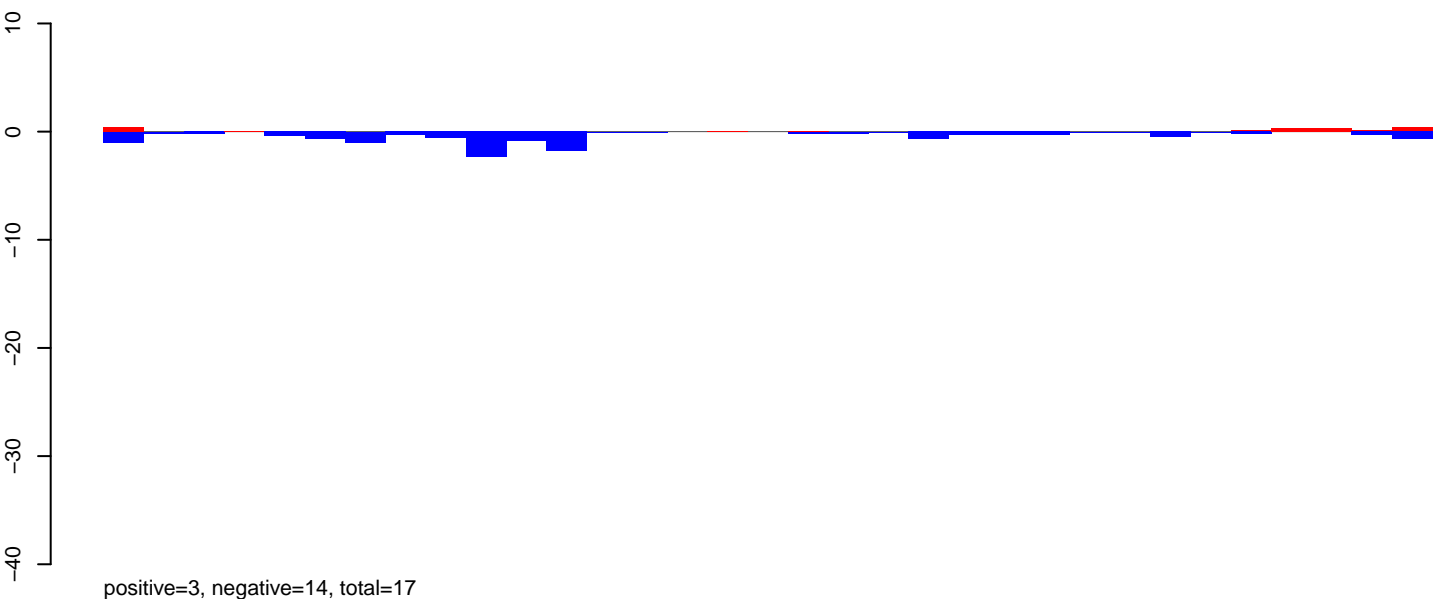


positive=1, negative=999, total=1000

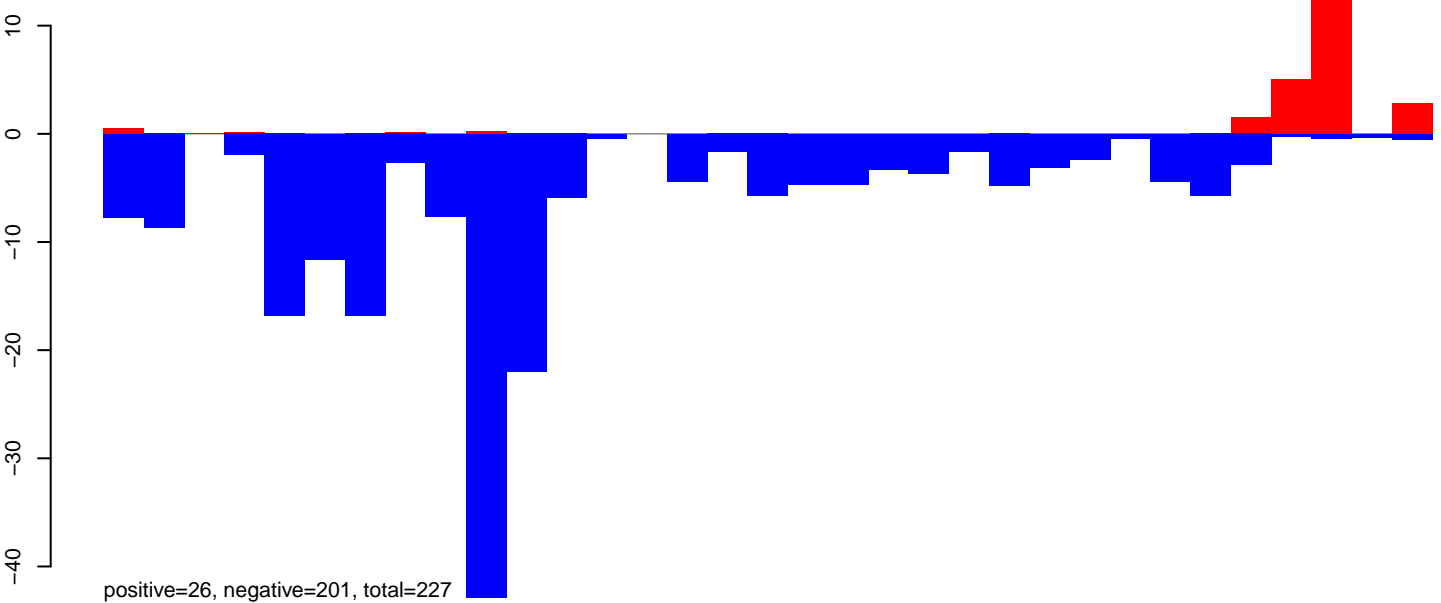
Window size=50, length=6565, TE@GYPSY12-LTR-L_AG:1-6565

0 1000 2000 3000 4000 5000 6000

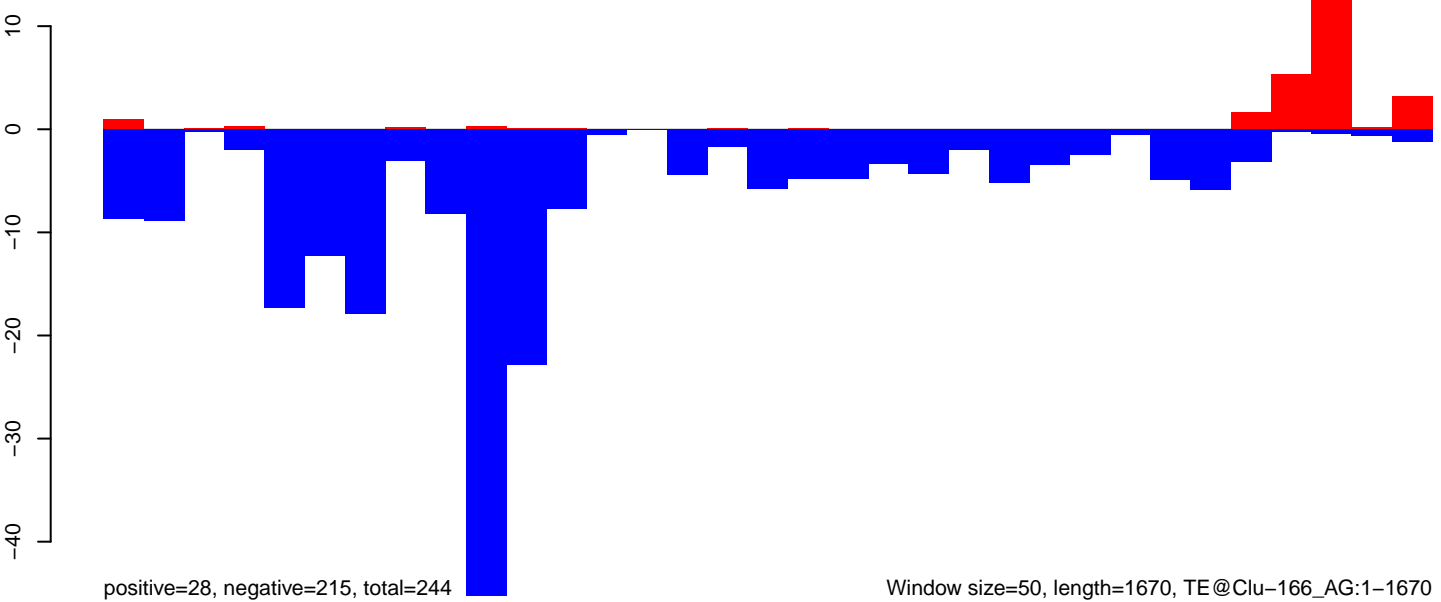
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



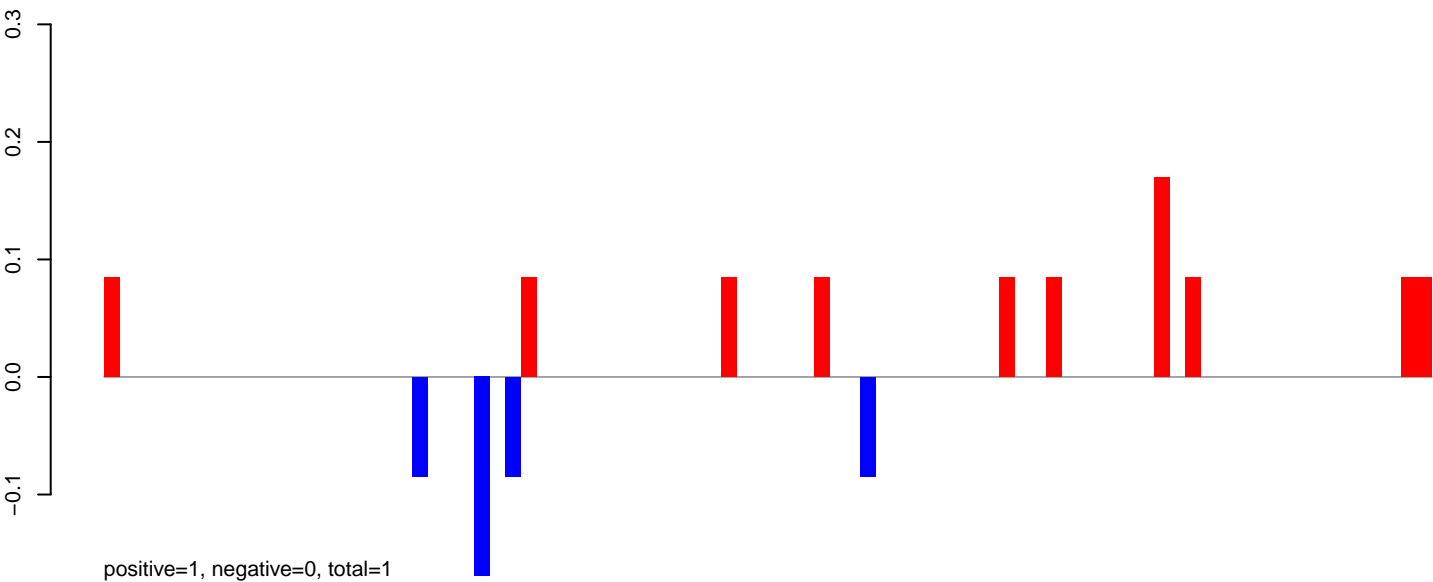
AnGam_Mos55_cells.rep



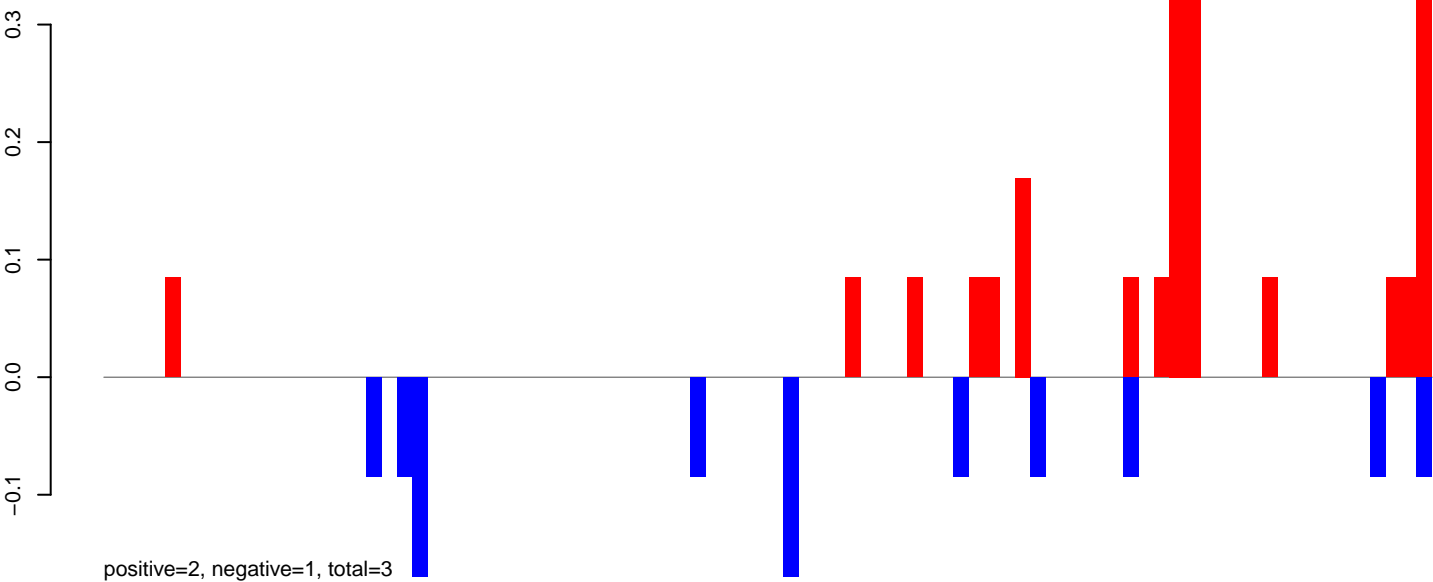
Window size=50, length=1670, TE@Clu-166_AG:1-1670

0 500 1000 1500

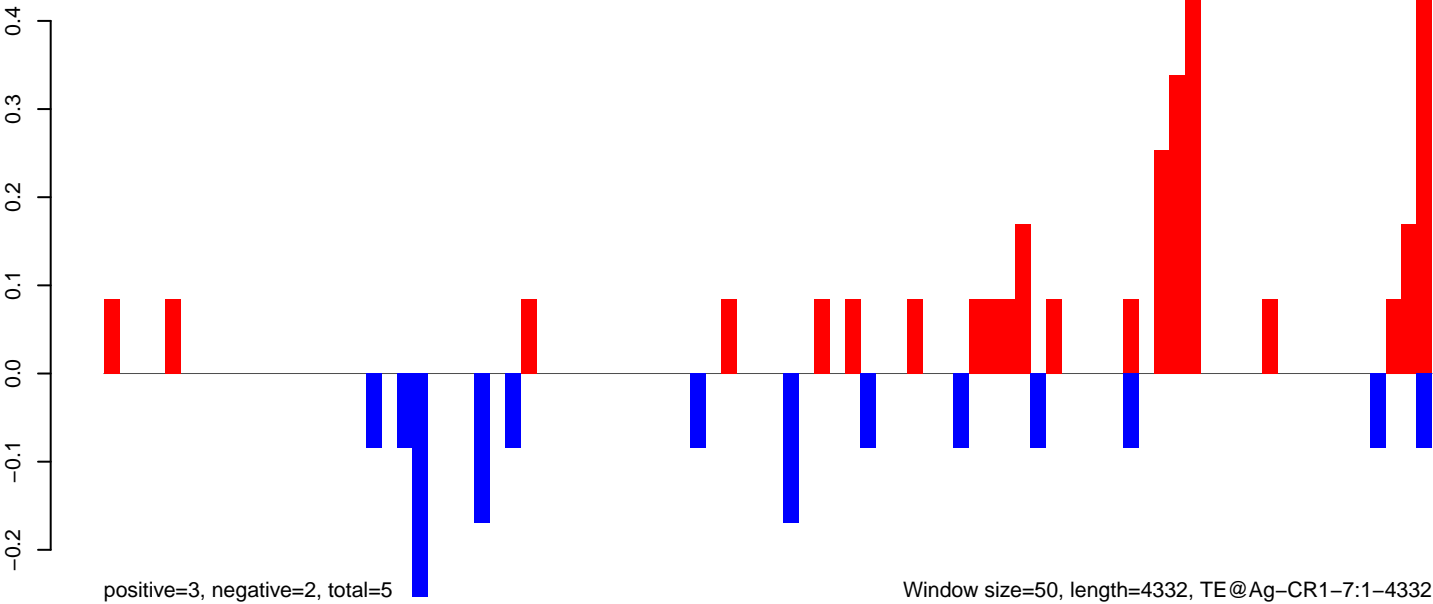
AnGam_Mos55_cells.18_23.rep



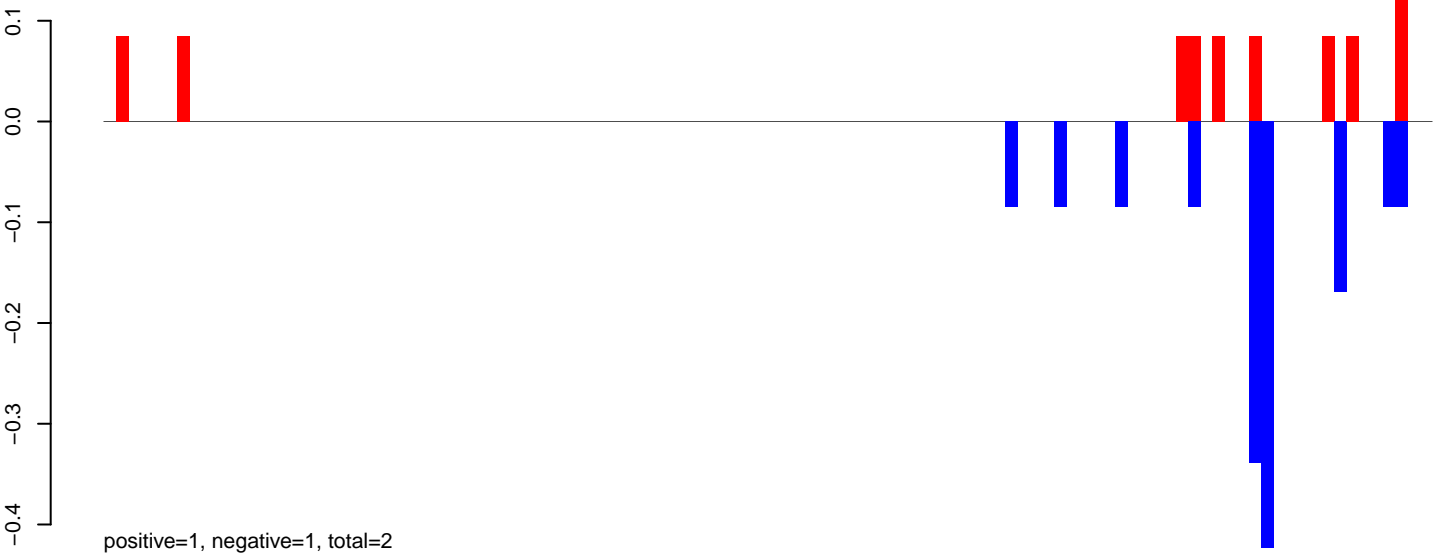
AnGam_Mos55_cells.24_35.rep



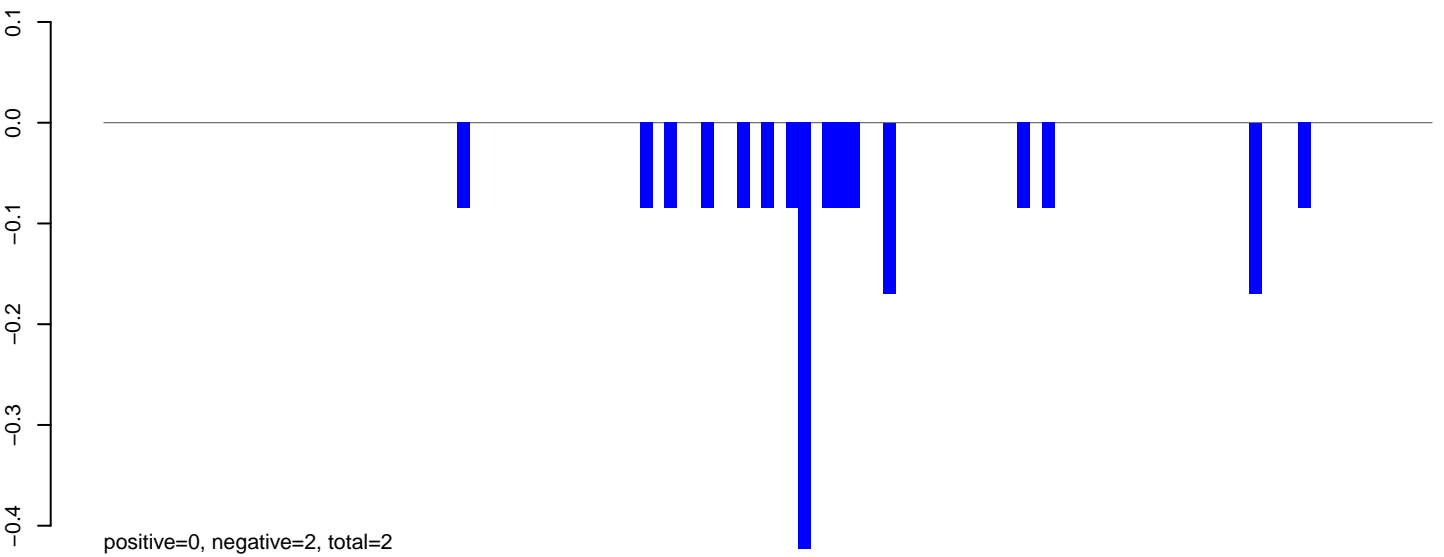
AnGam_Mos55_cells.rep



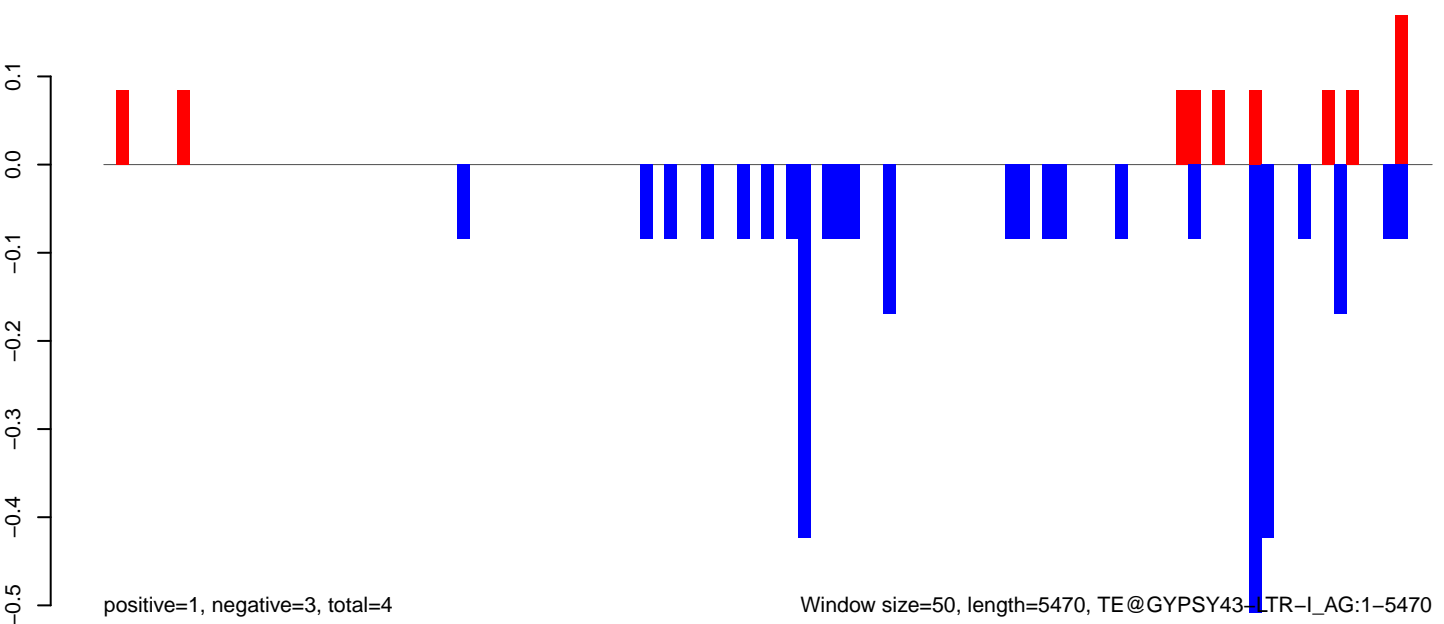
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

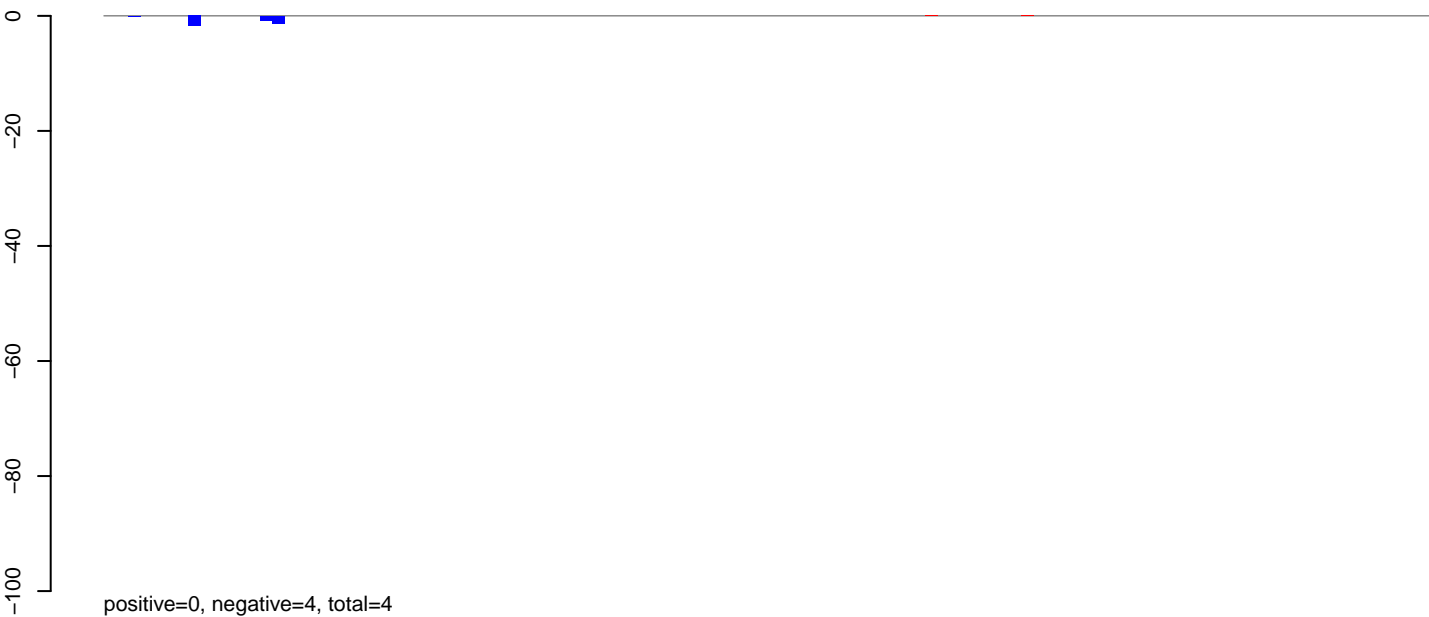


AnGam_Mos55_cells.rep

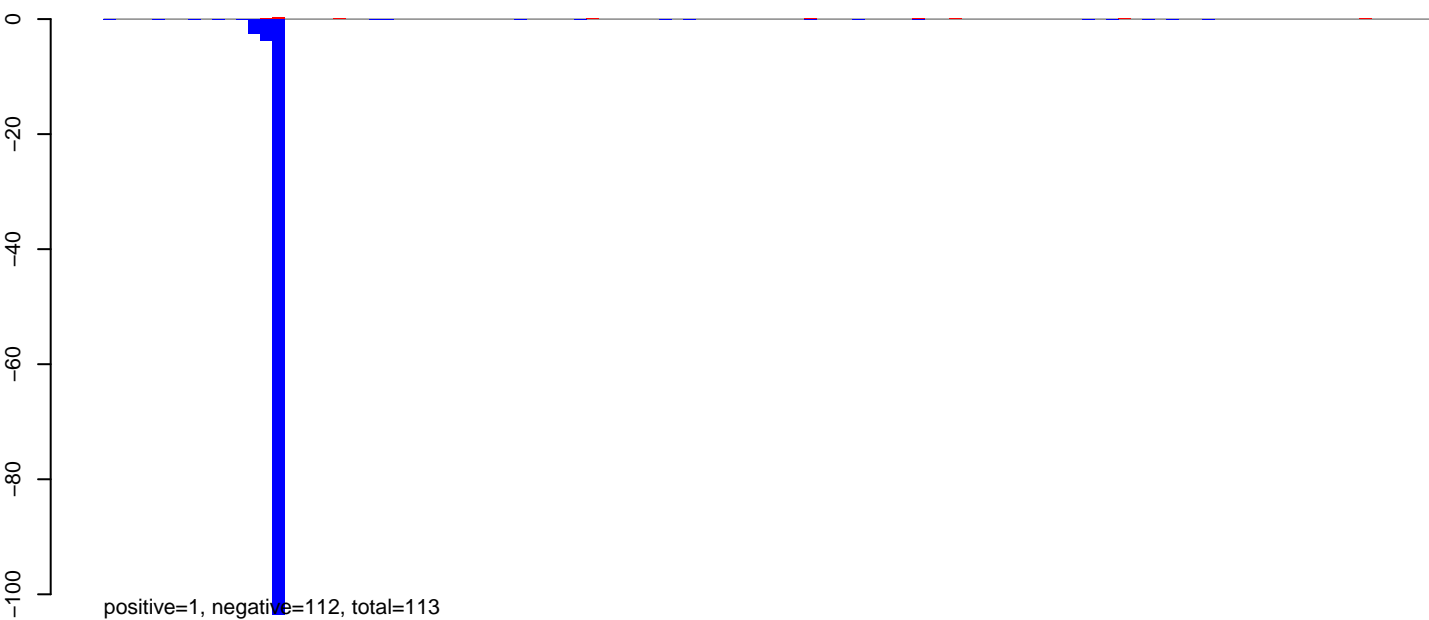


Window size=50, length=5470, TE@GYPSY43-LTR-I_AG:1-5470

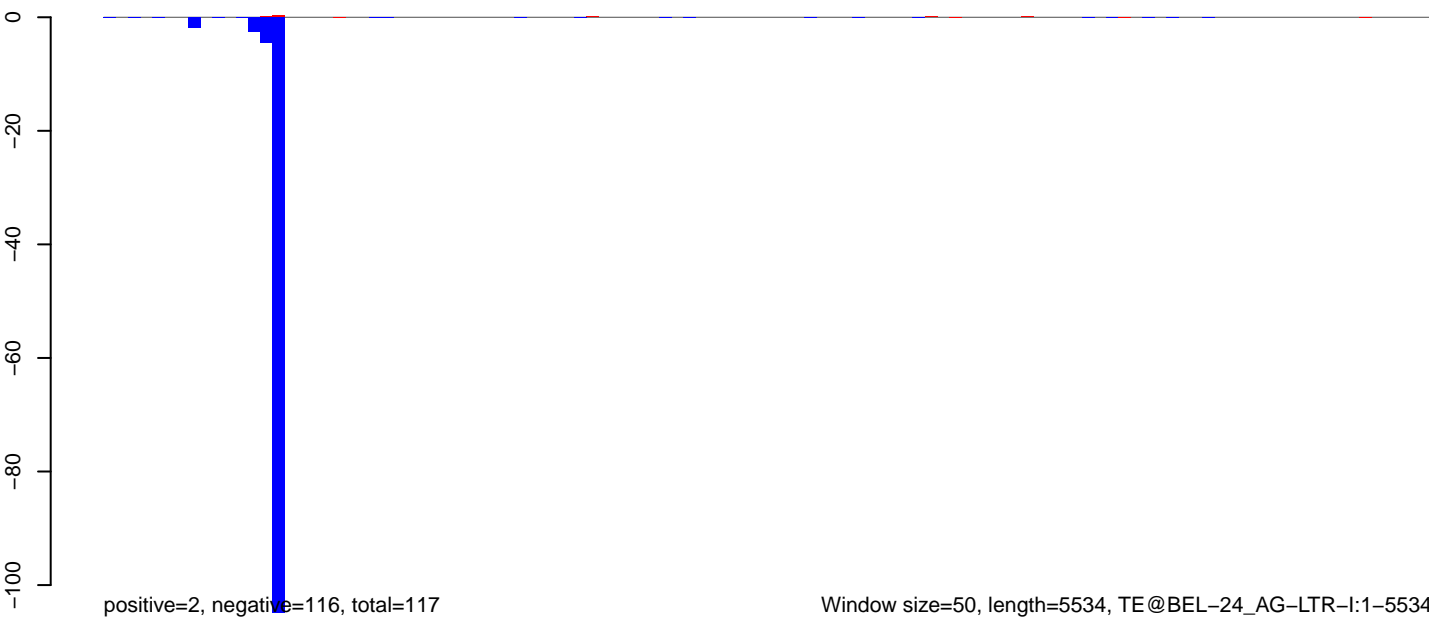
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

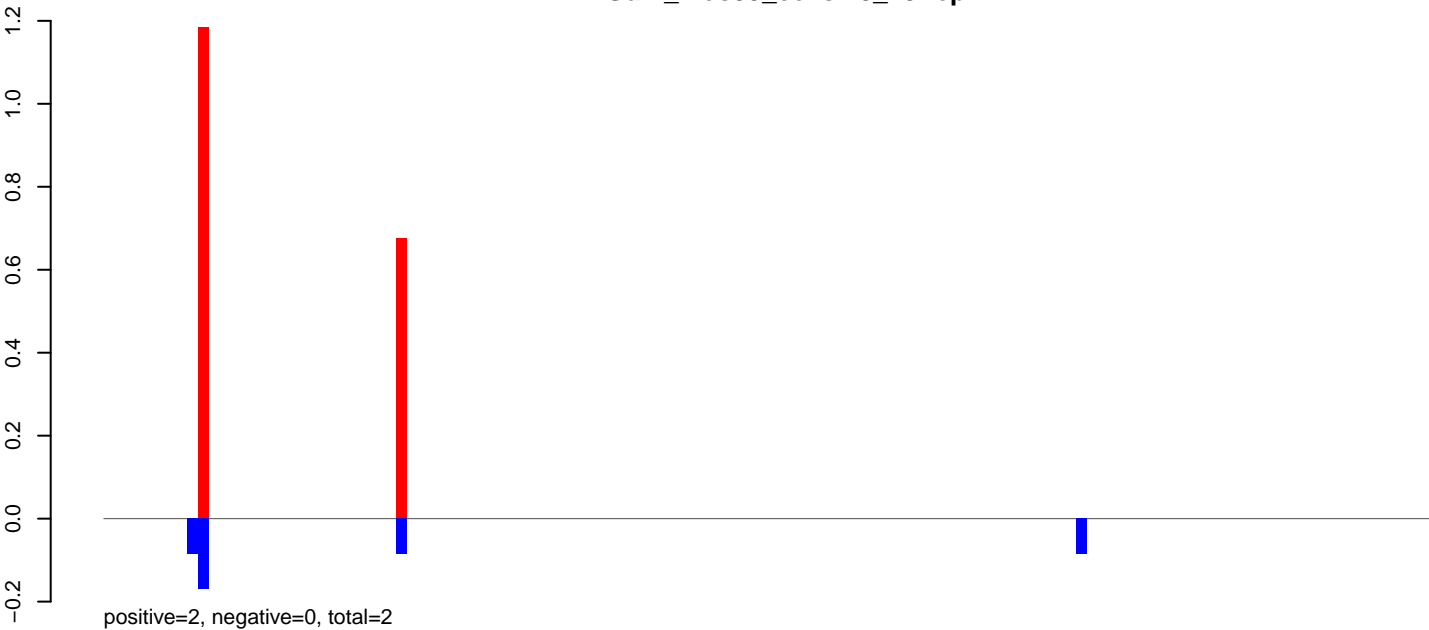


AnGam_Mos55_cells.rep

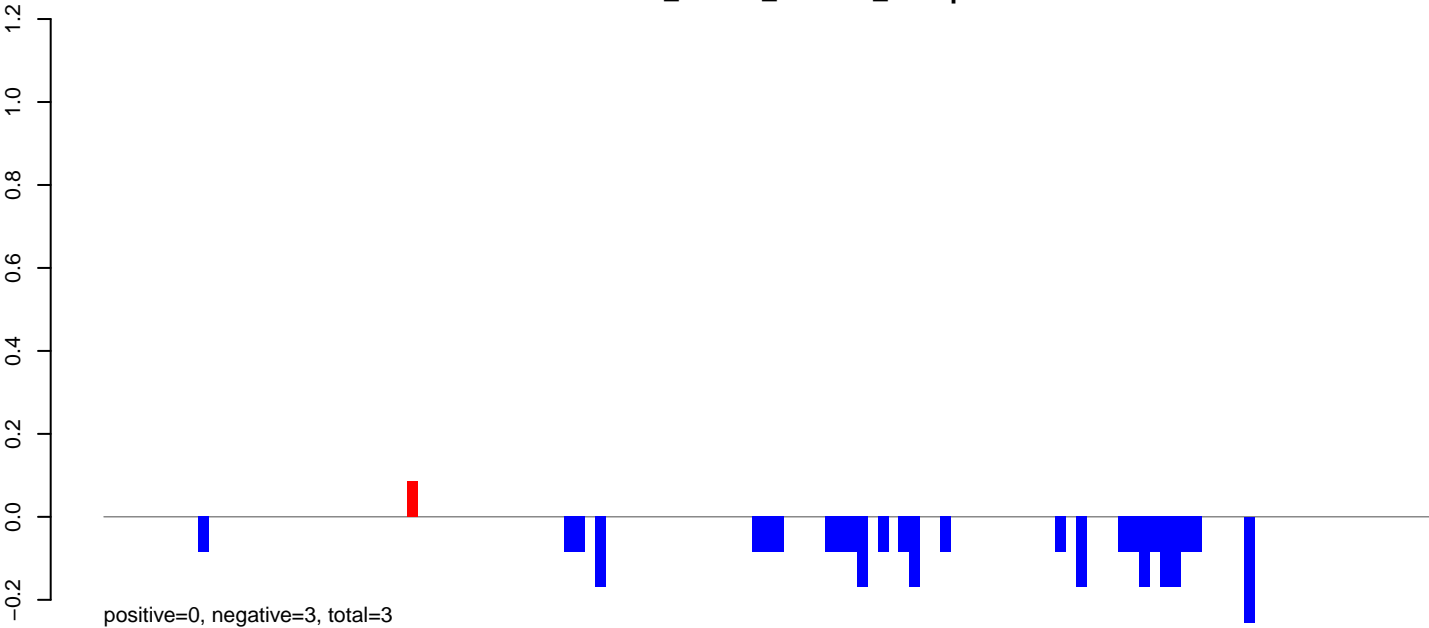


Window size=50, length=5534, TE@BEL-24_AG-LTR-I:1-5534

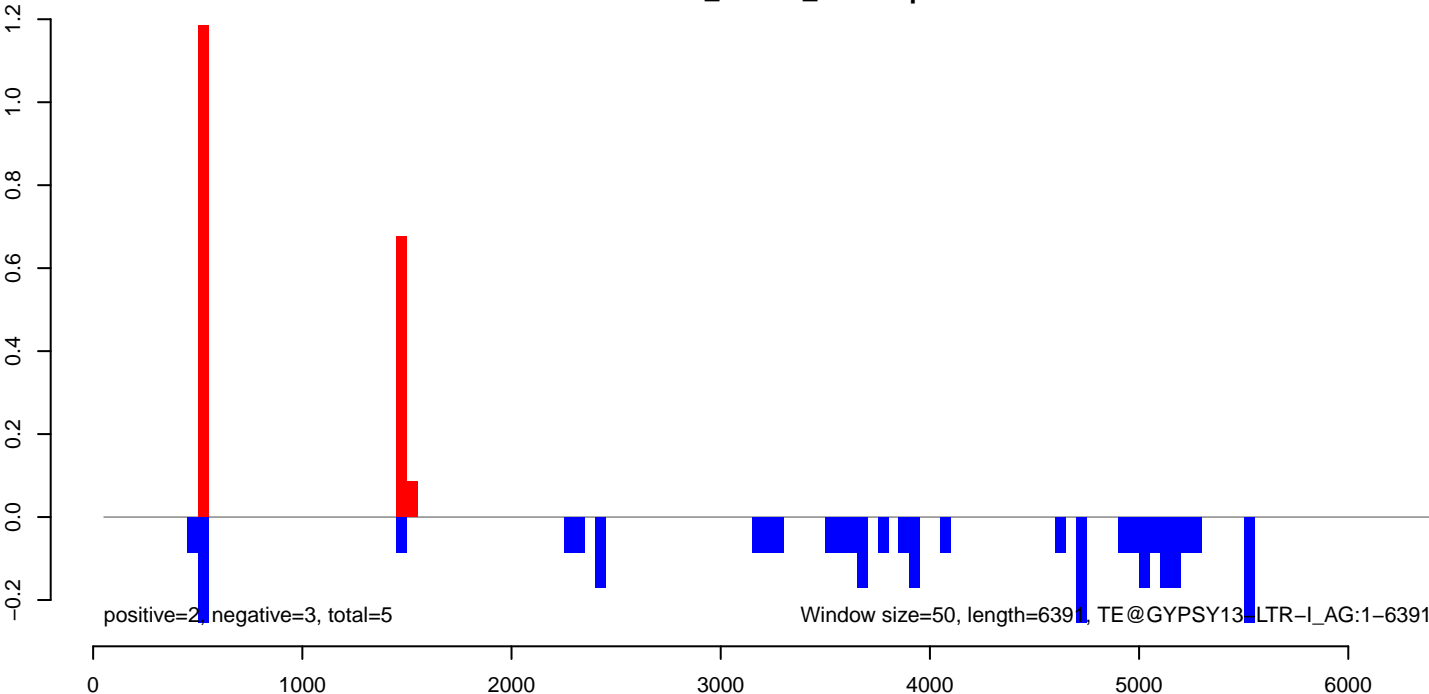
AnGam_Mos55_cells.18_23.rep



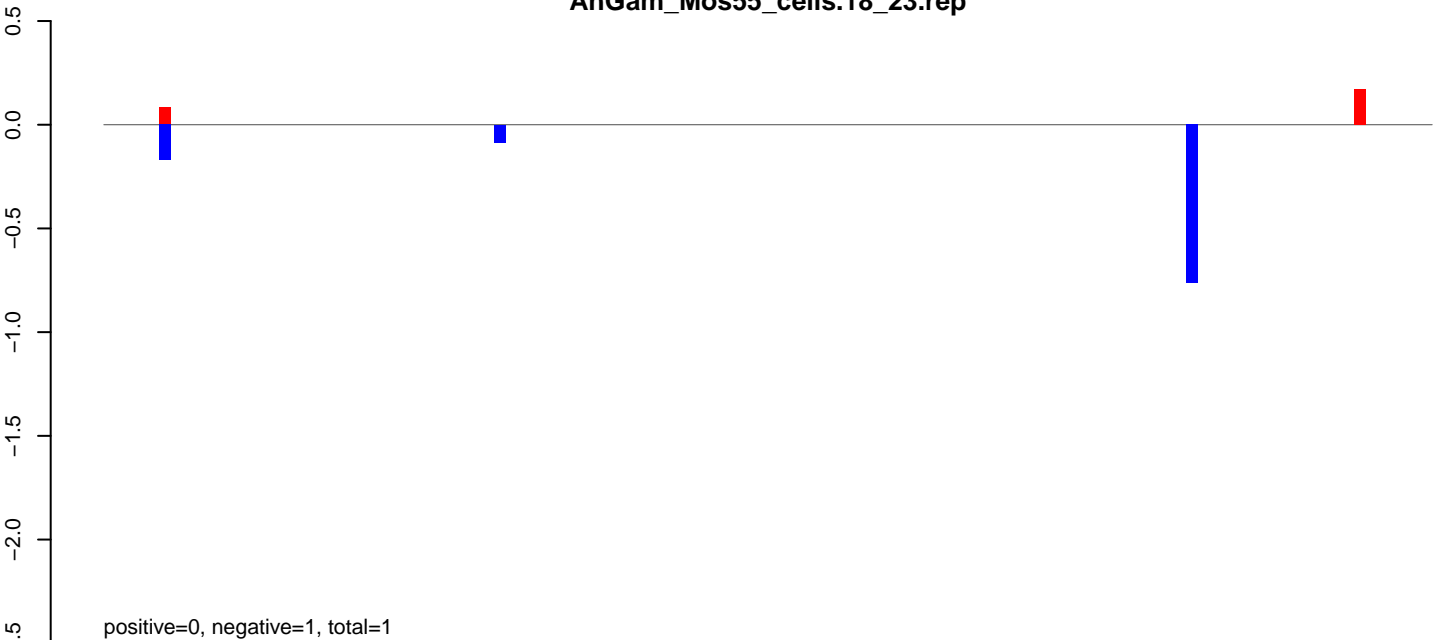
AnGam_Mos55_cells.24_35.rep



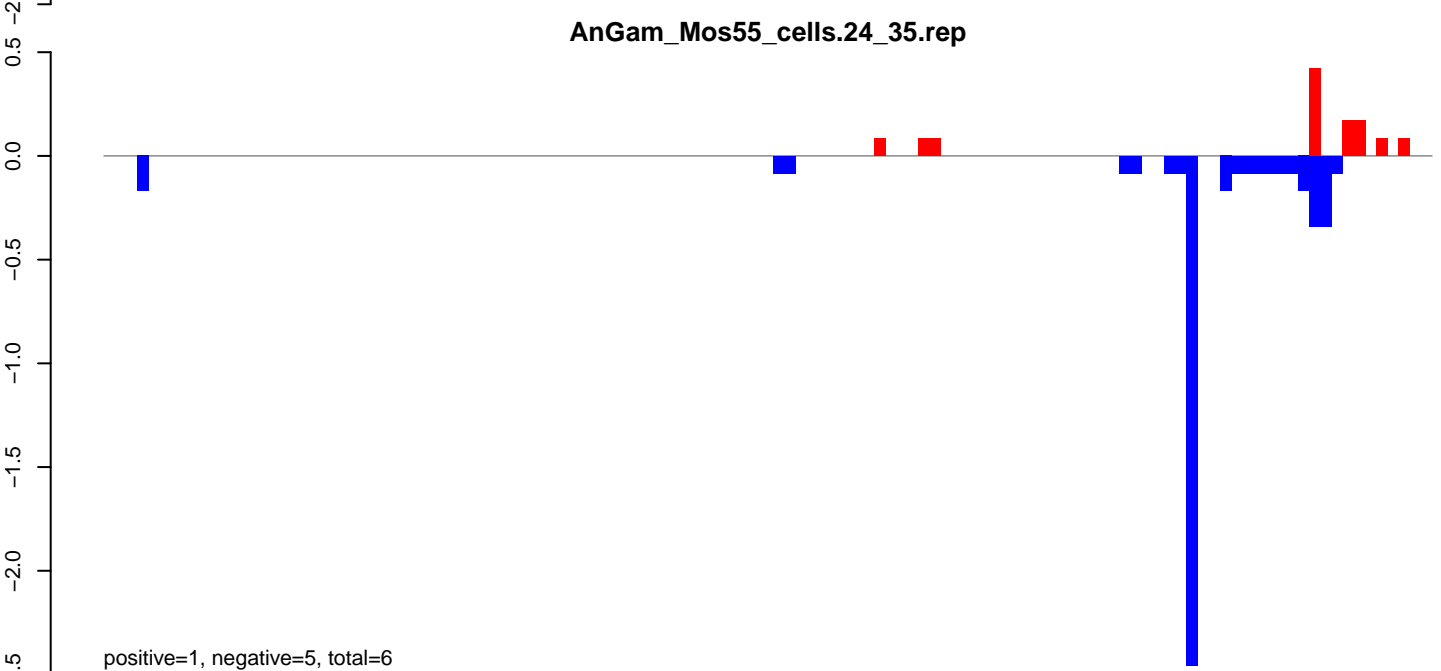
AnGam_Mos55_cells.rep



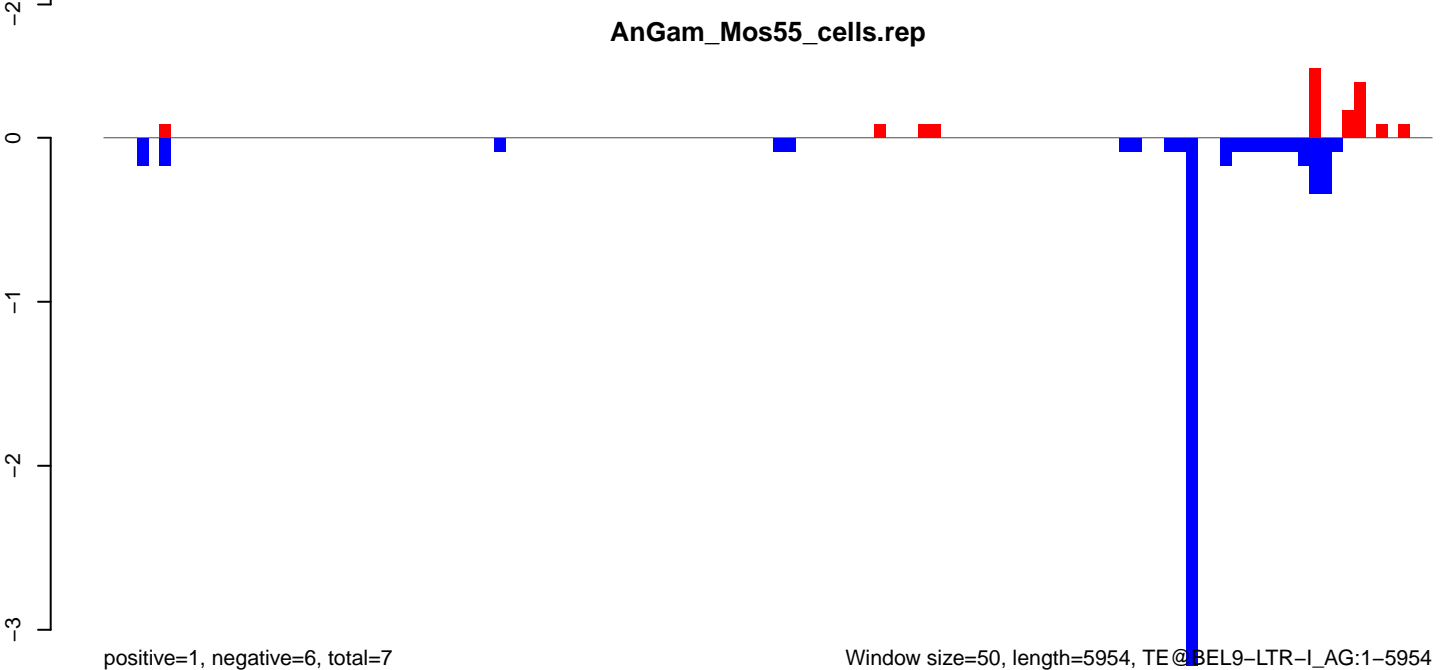
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

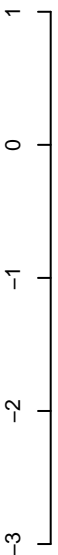


AnGam_Mos55_cells.rep



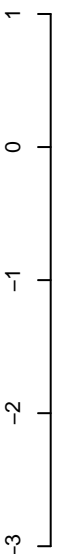
Window size=50, length=5954, TE@BEL9-LTR-I_AG:1-5954

AnGam_Mos55_cells.18_23.rep



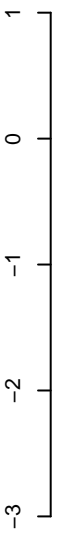
positive=2, negative=2, total=5

AnGam_Mos55_cells.24_35.rep



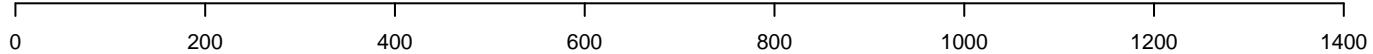
positive=3, negative=8, total=11

AnGam_Mos55_cells.rep

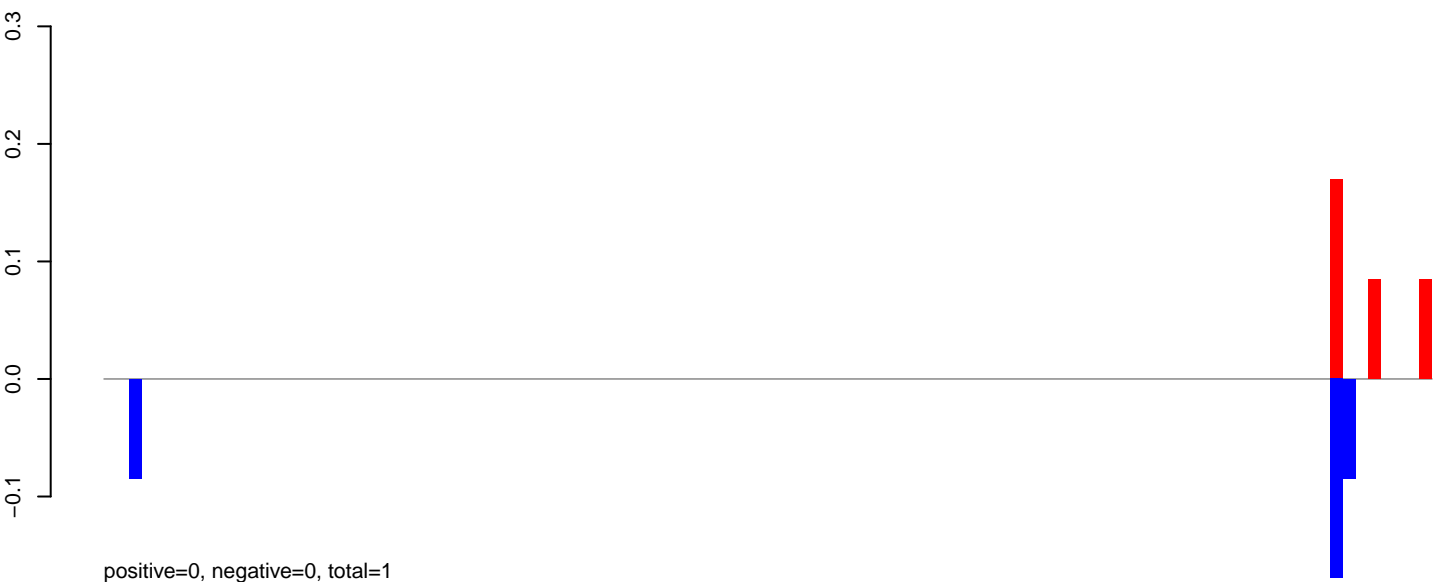


positive=5, negative=10, total=15

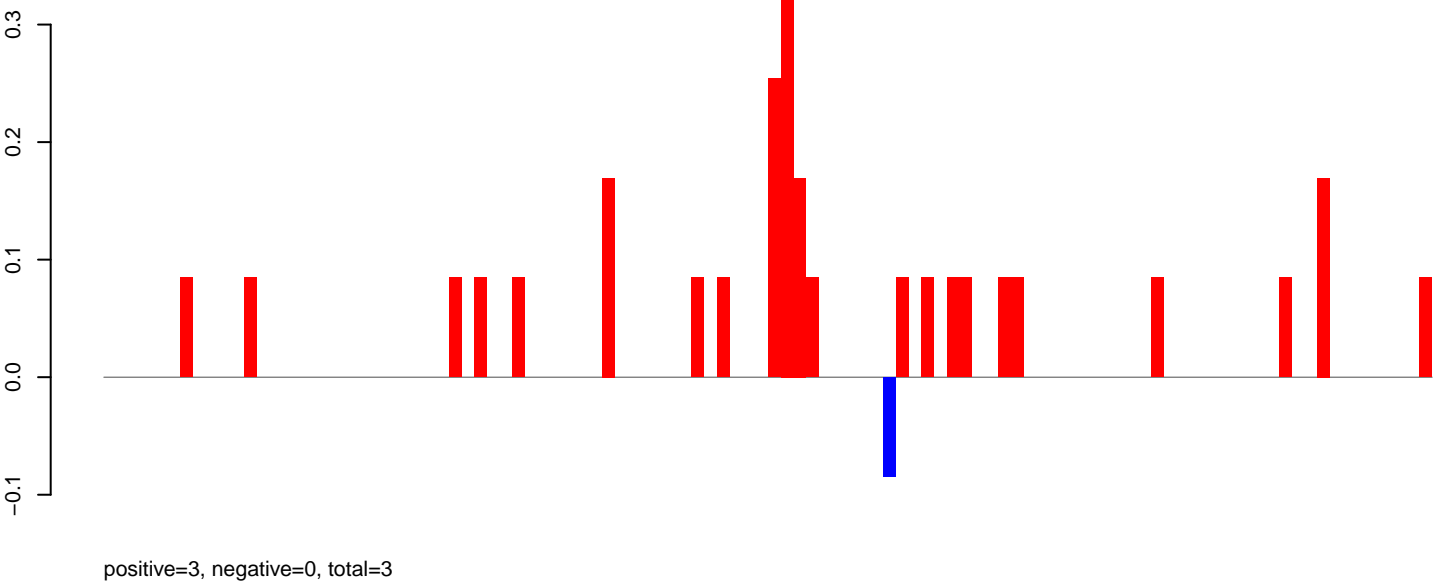
Window size=50, length=1410, TE@Mariner-N22_AG:1-1410



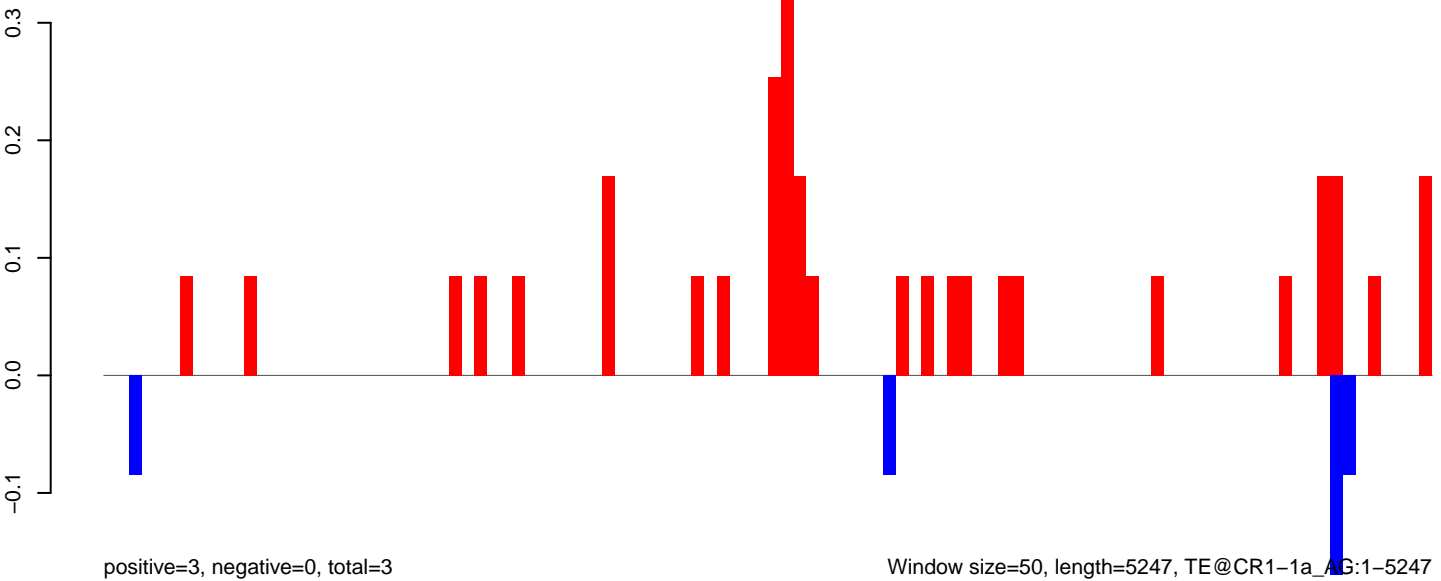
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



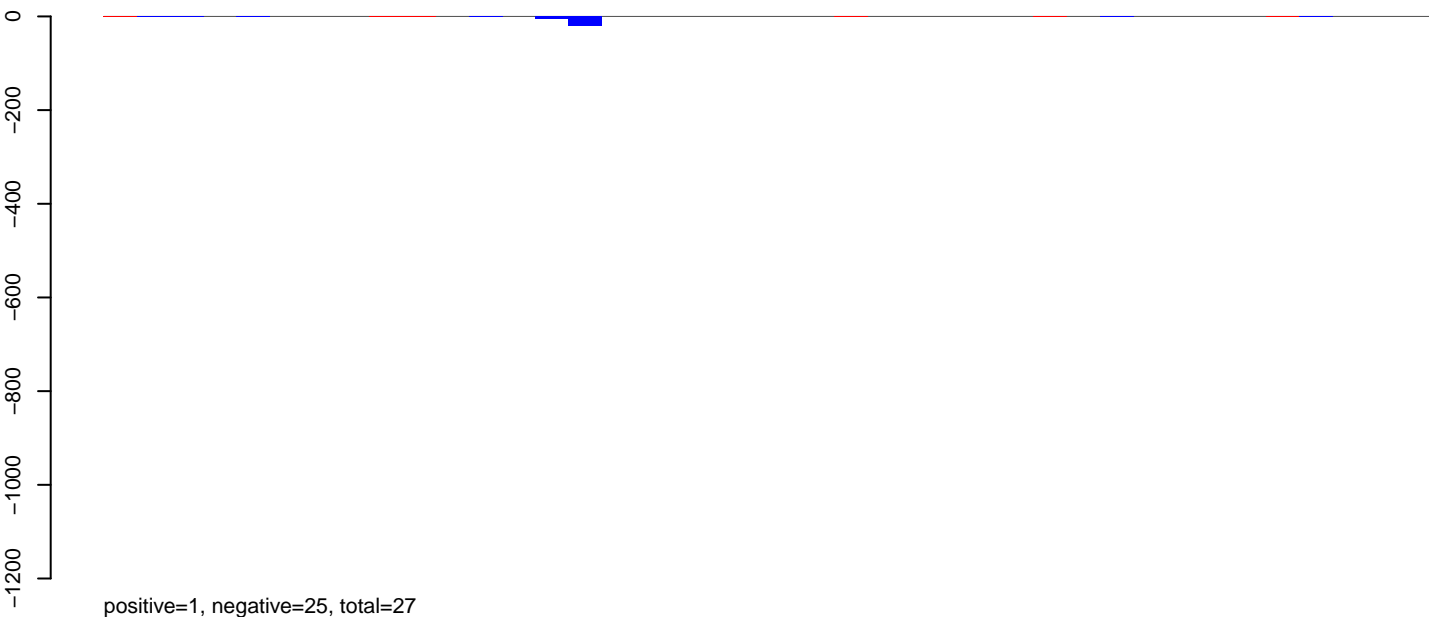
AnGam_Mos55_cells.rep



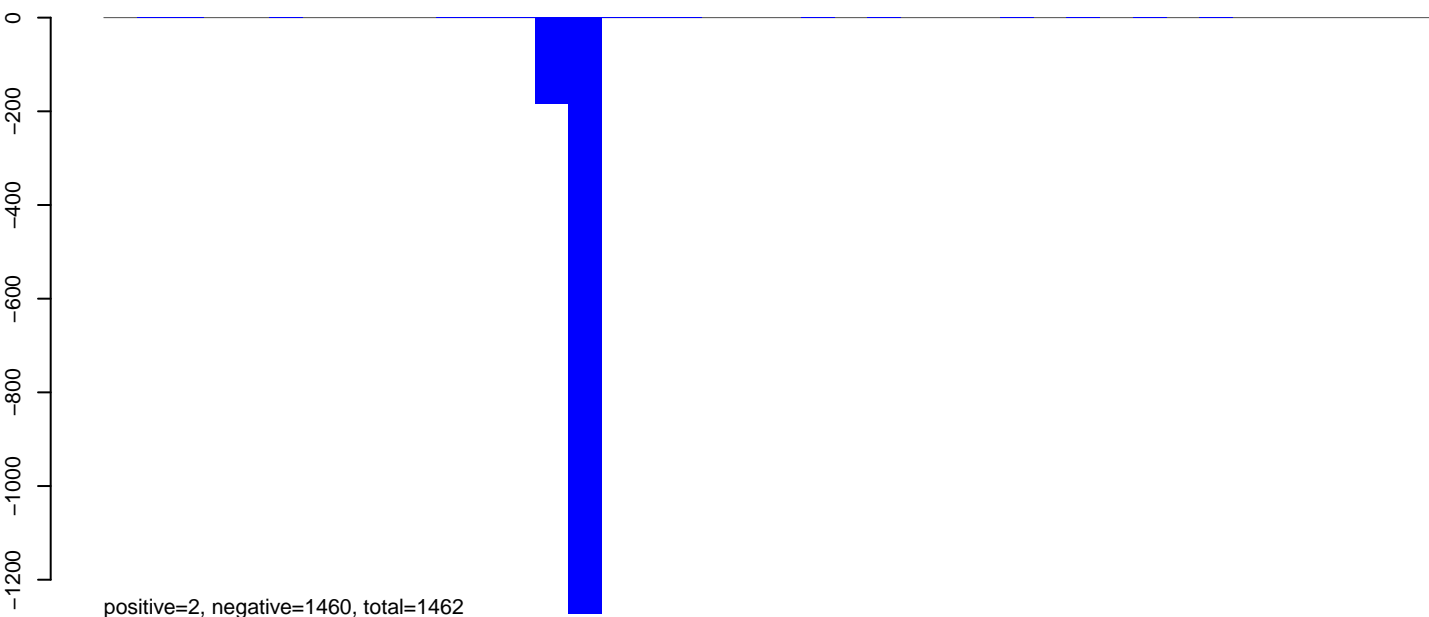
Window size=50, length=5247, TE@CR1-1a_AG:1-5247

0 1000 2000 3000 4000 5000

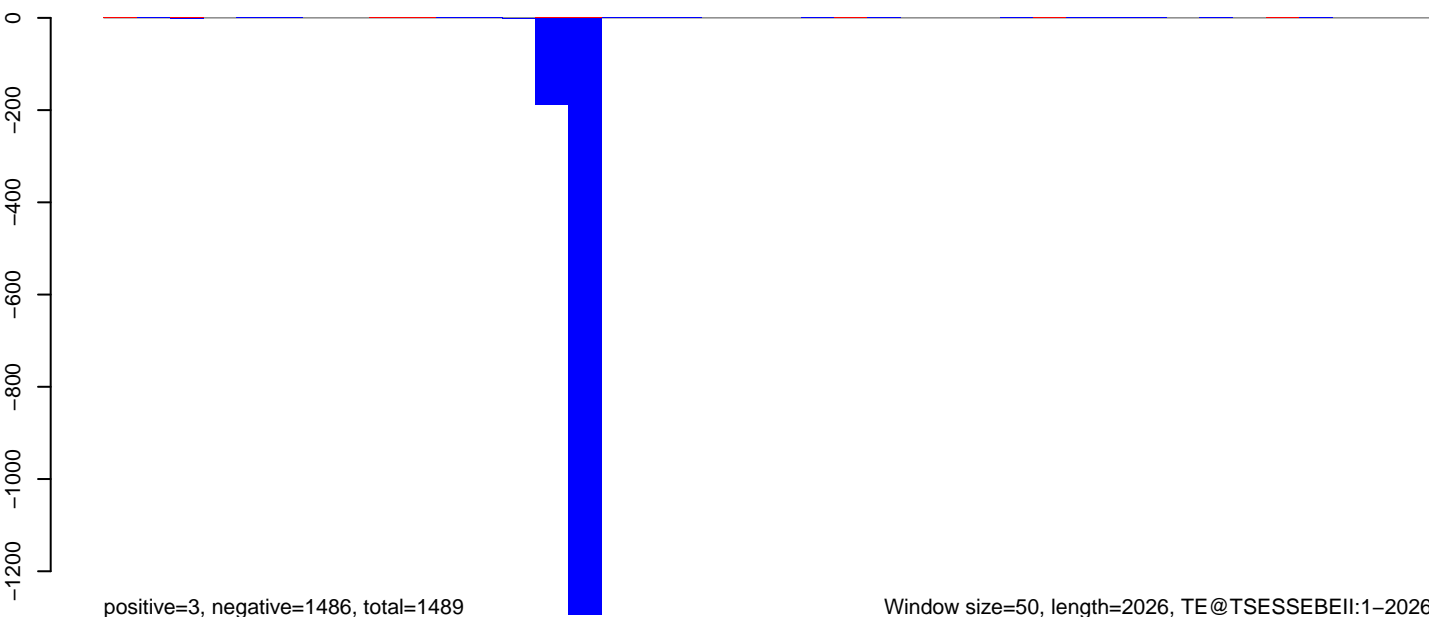
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



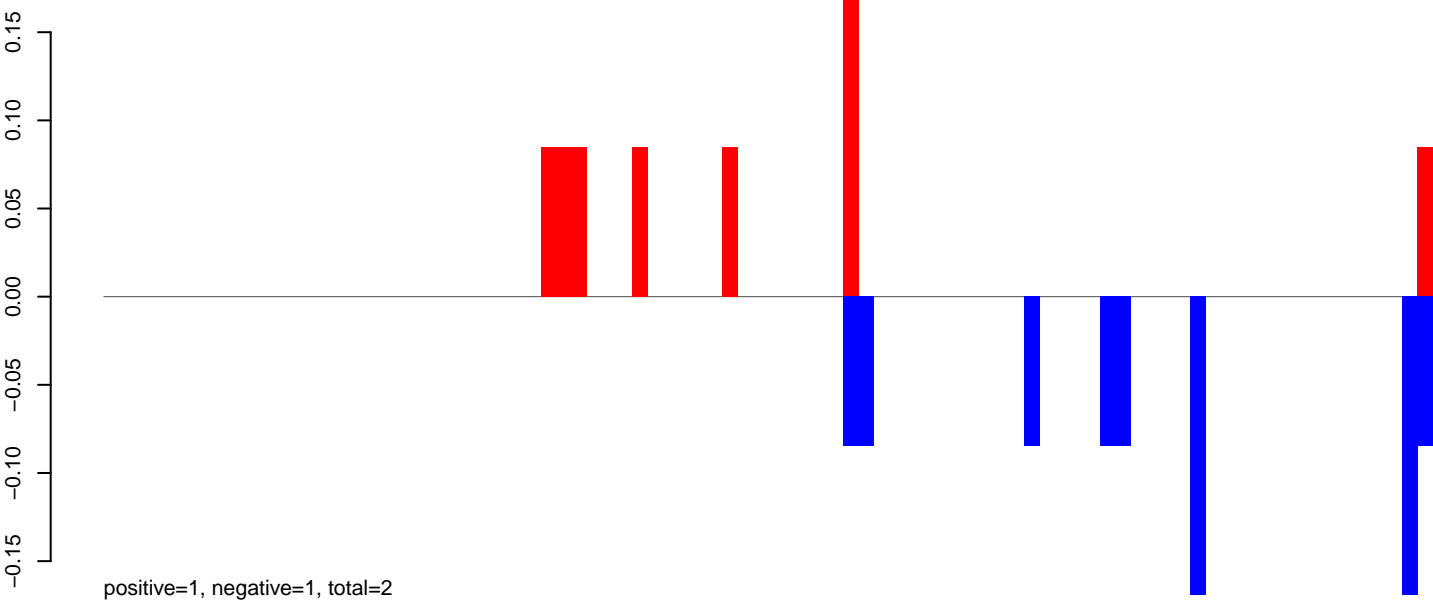
Window size=50, length=2026, TE@TSESSEBEII:1-2026

0 500 1000 1500 2000

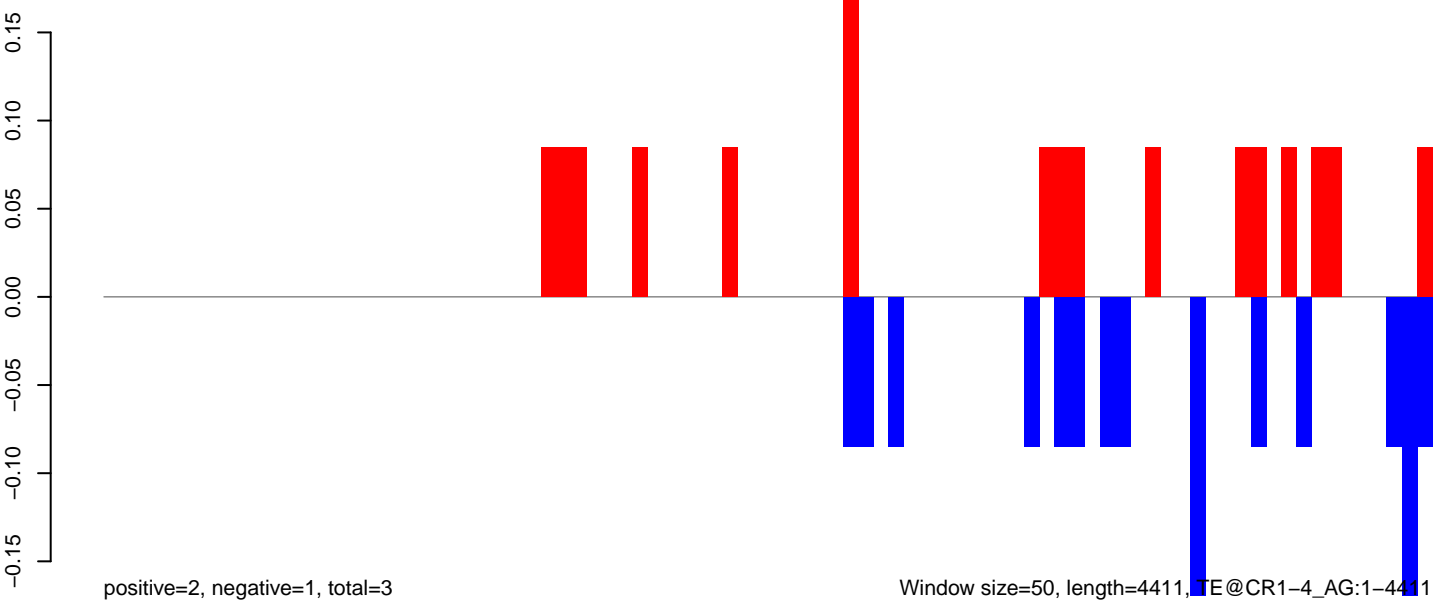
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

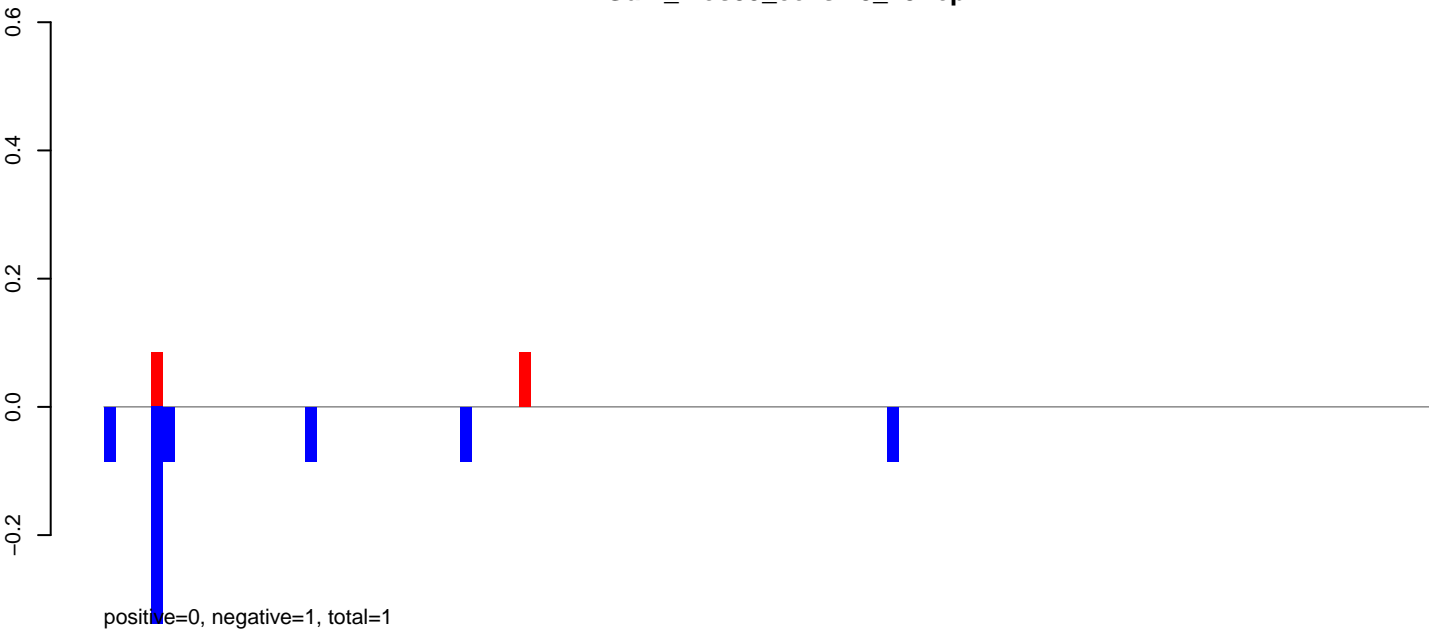


AnGam_Mos55_cells.rep

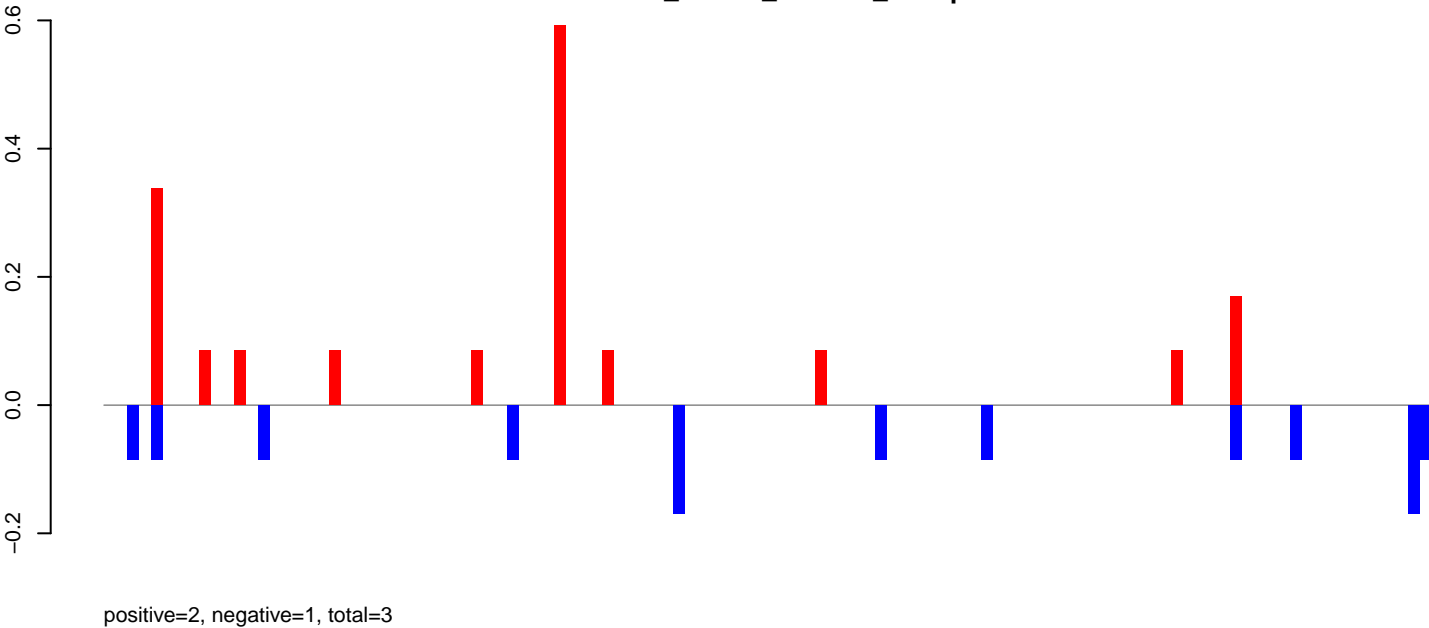


Window size=50, length=4411, TE@CR1-4_AG:1-4411

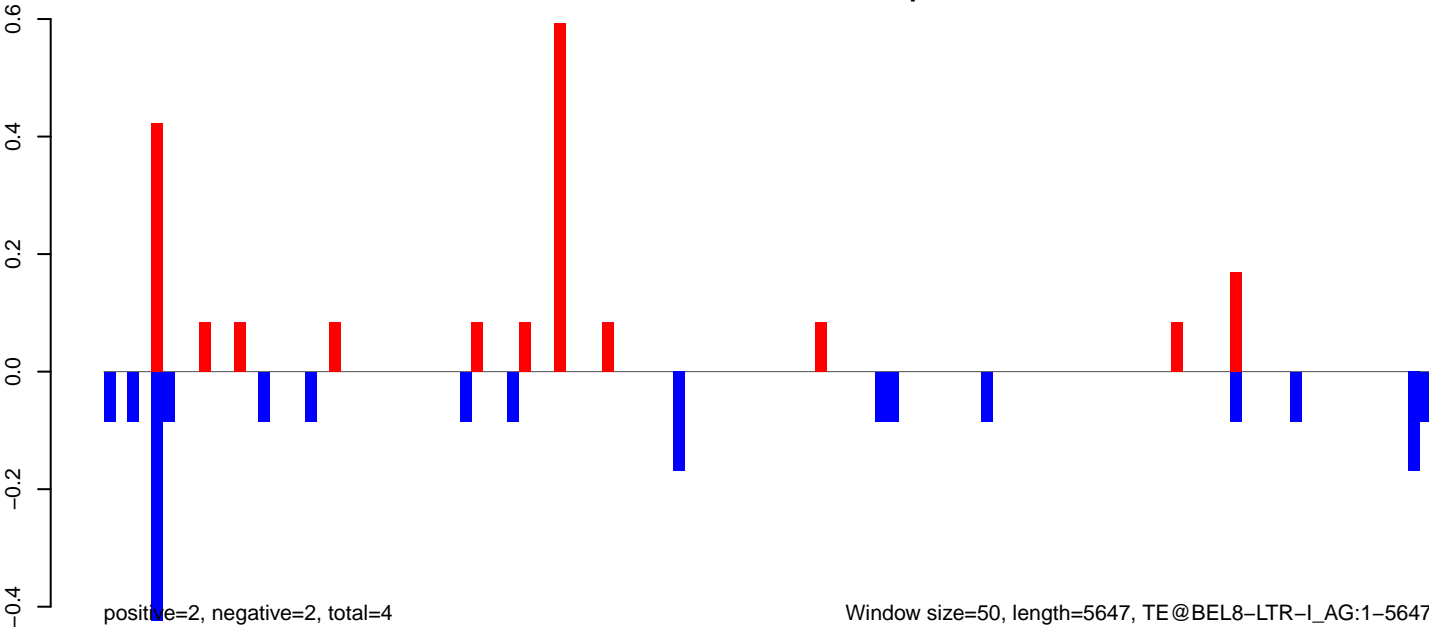
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

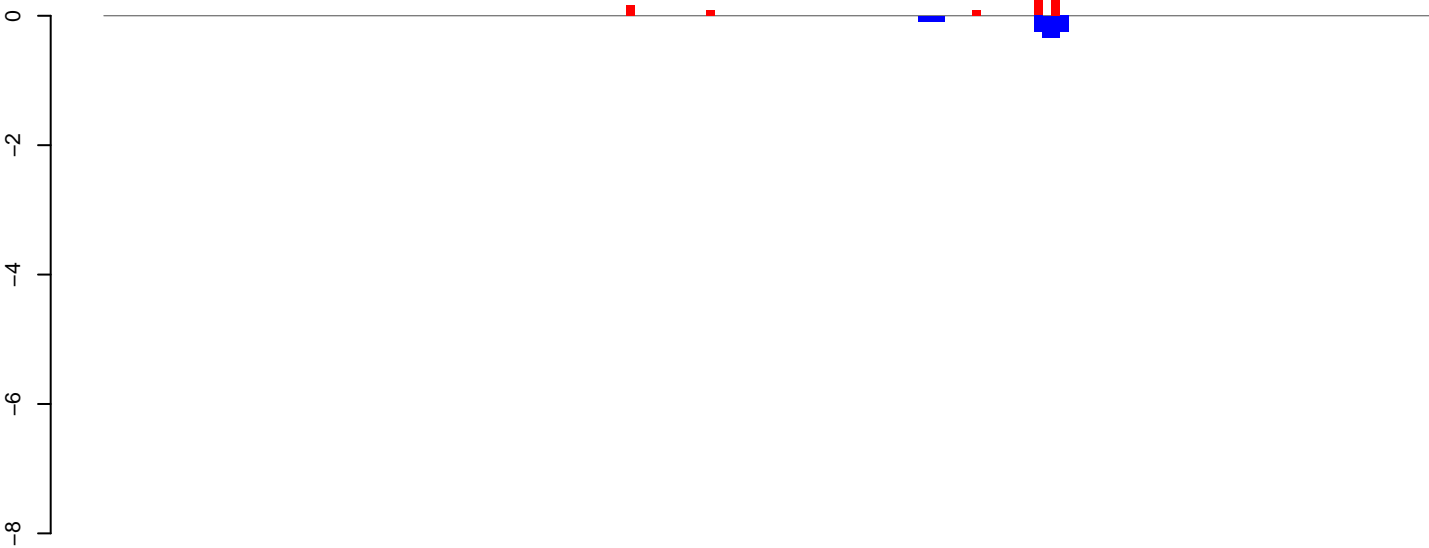


AnGam_Mos55_cells.rep



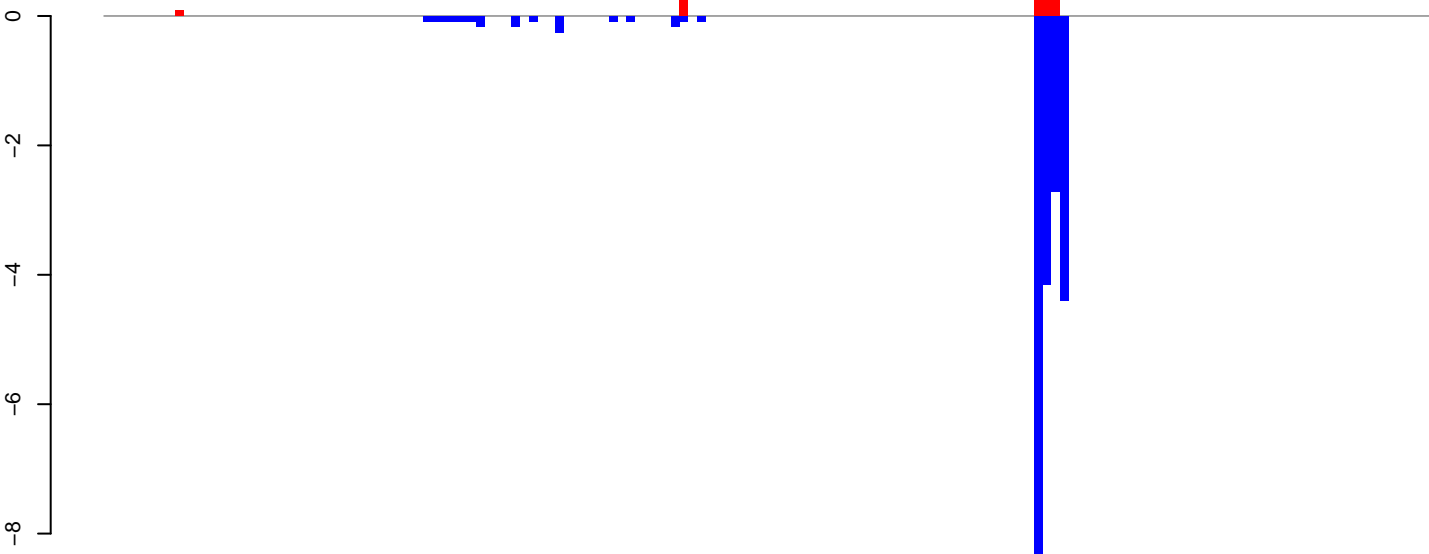
Window size=50, length=5647, TE@BEL8-LTR-I_AG:1-5647

AnGam_Mos55_cells.18_23.rep



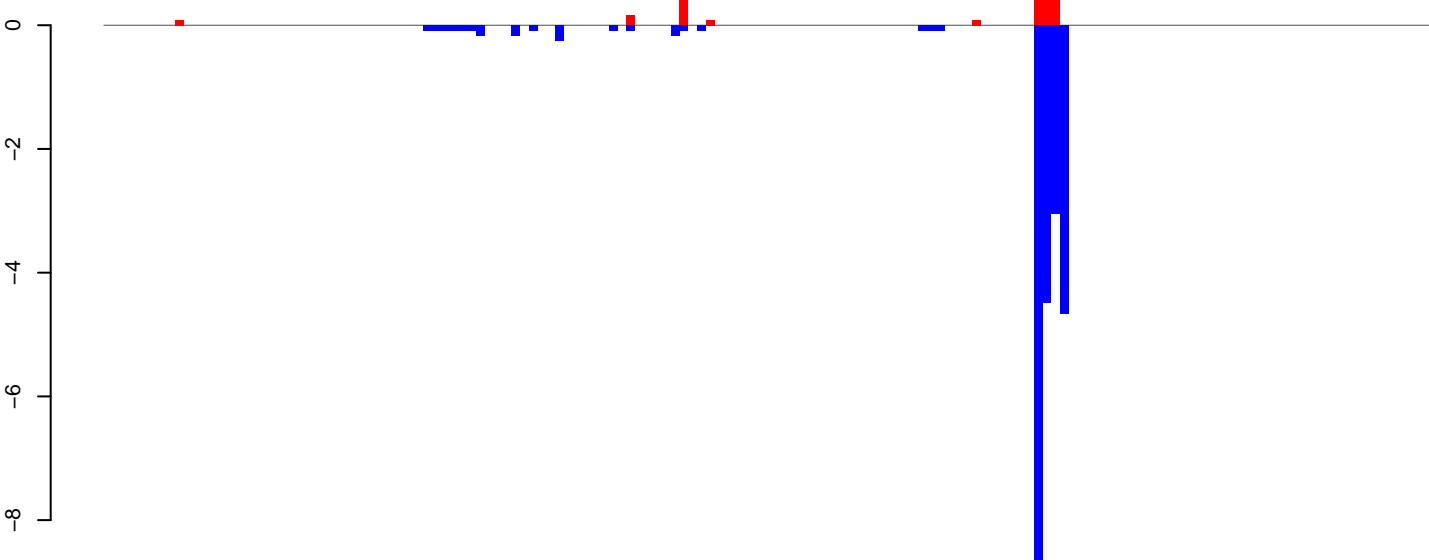
positive=1, negative=1, total=2

AnGam_Mos55_cells.24_35.rep



positive=2, negative=22, total=24

AnGam_Mos55_cells.rep

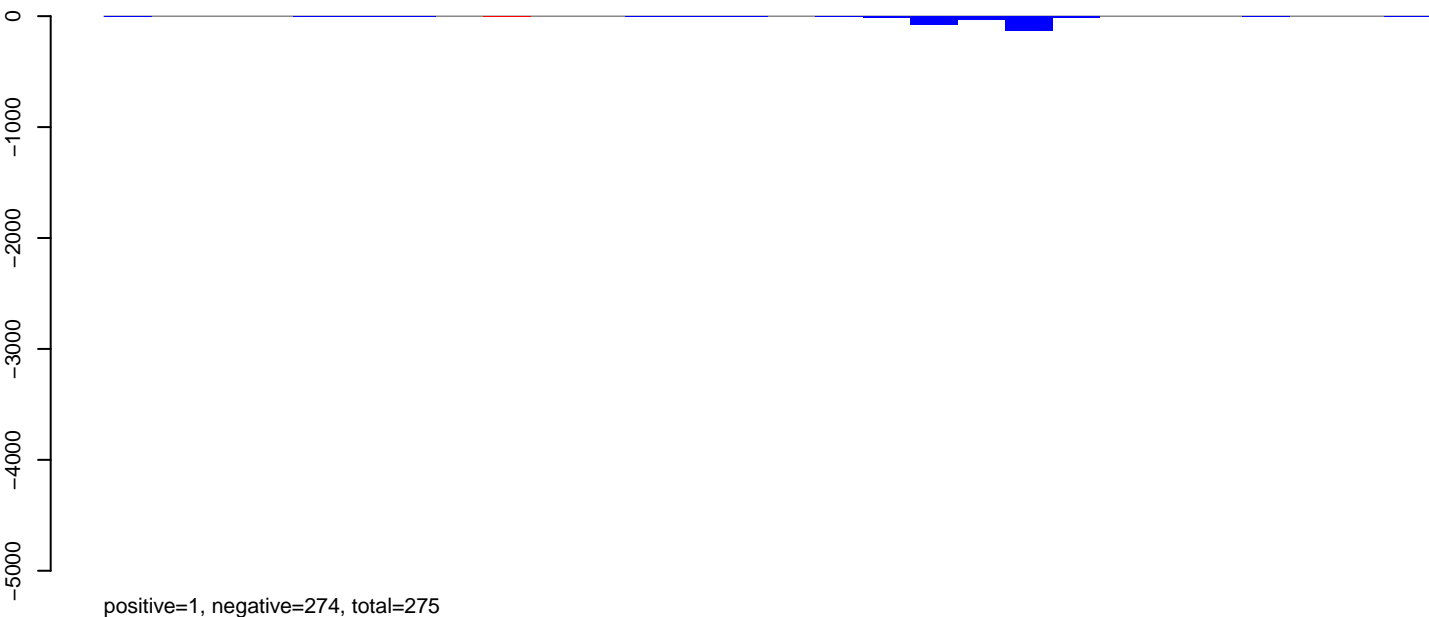


positive=3, negative=23, total=26

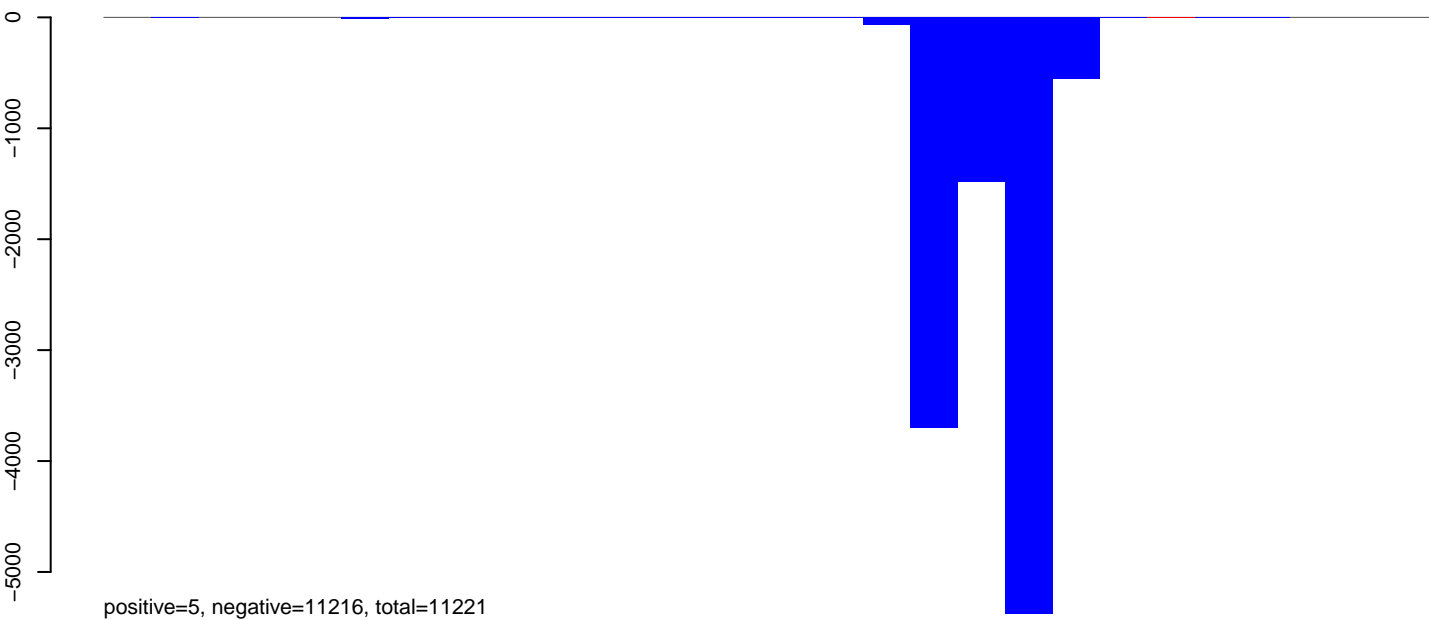
Window size=50, length=7508, TE@AgaP8:1-7508

0 2000 4000 6000

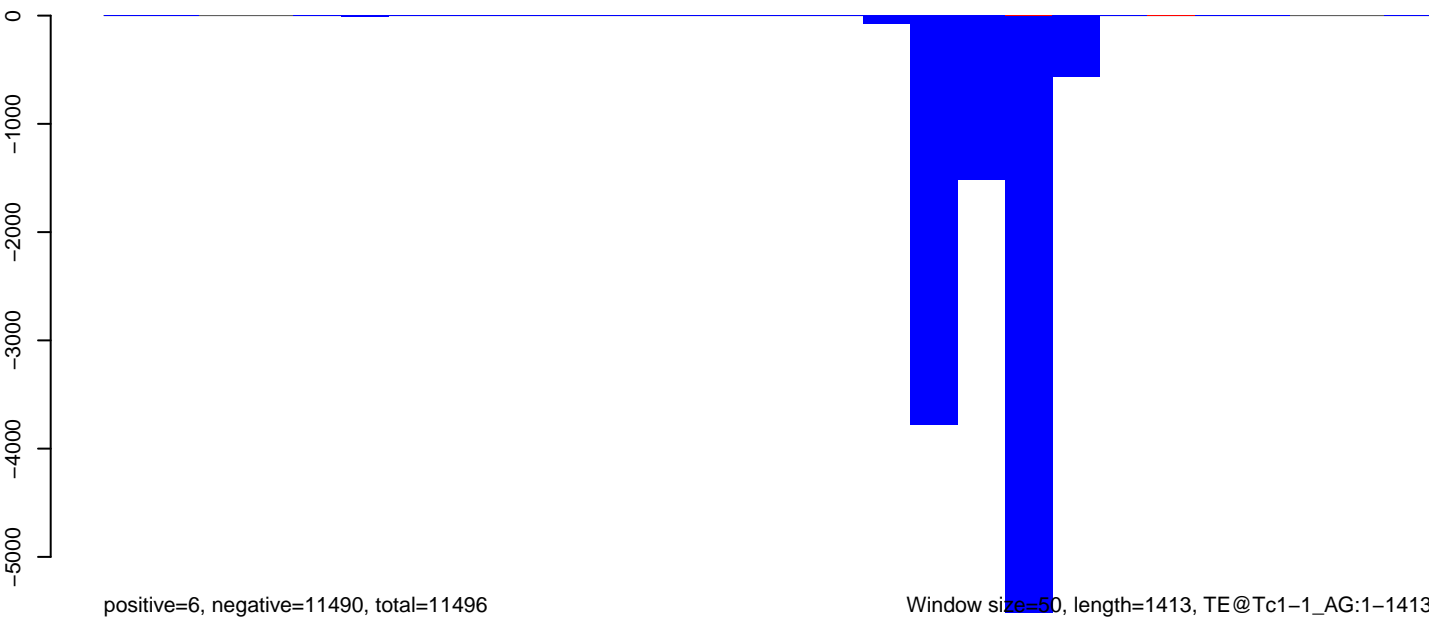
AnGam_Mos55_cells.18_23.rep



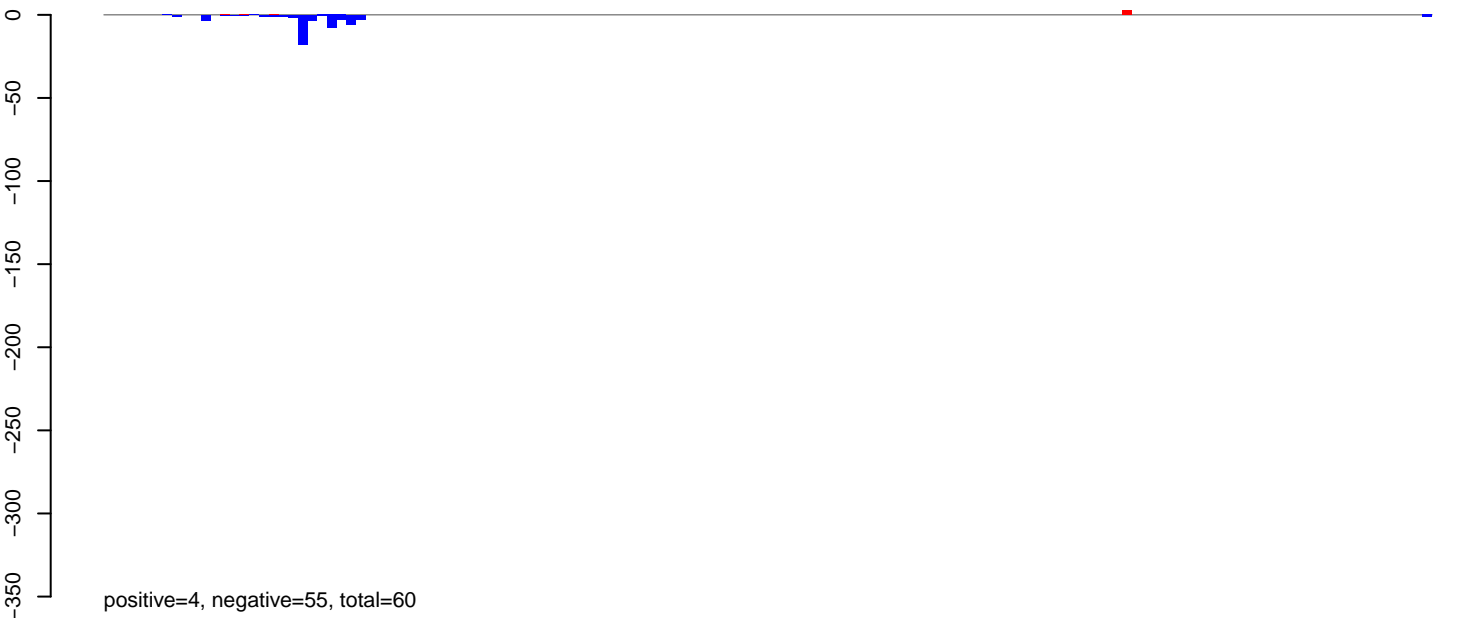
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



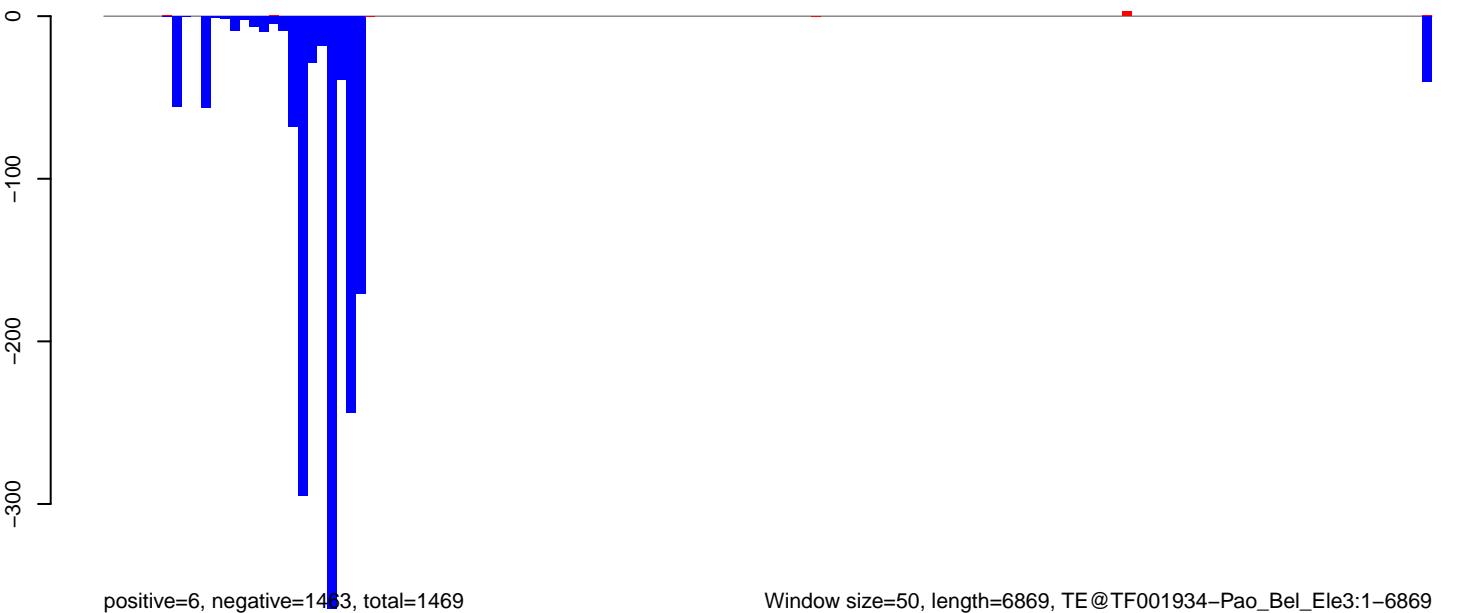
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



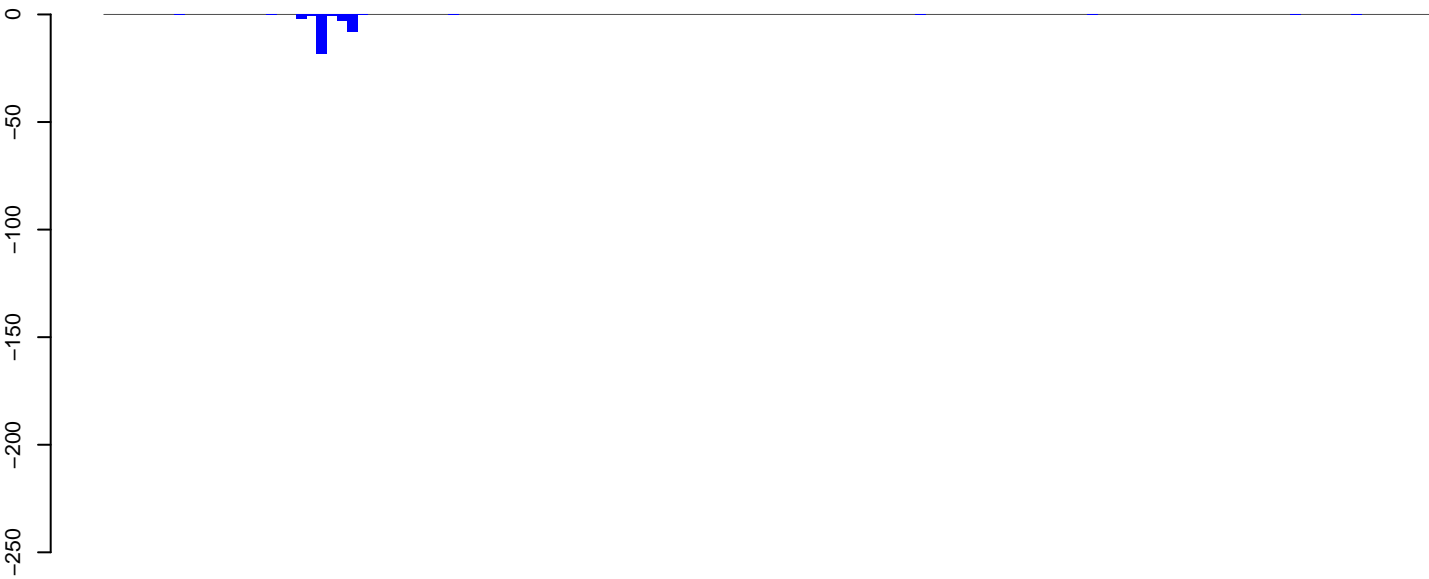
AnGam_Mos55_cells.rep



Window size=50, length=6869, TE@TF001934-Pao_Bel_Ele3:1-6869

0 1000 2000 3000 4000 5000 6000 7000

AnGam_Mos55_cells.18_23.rep



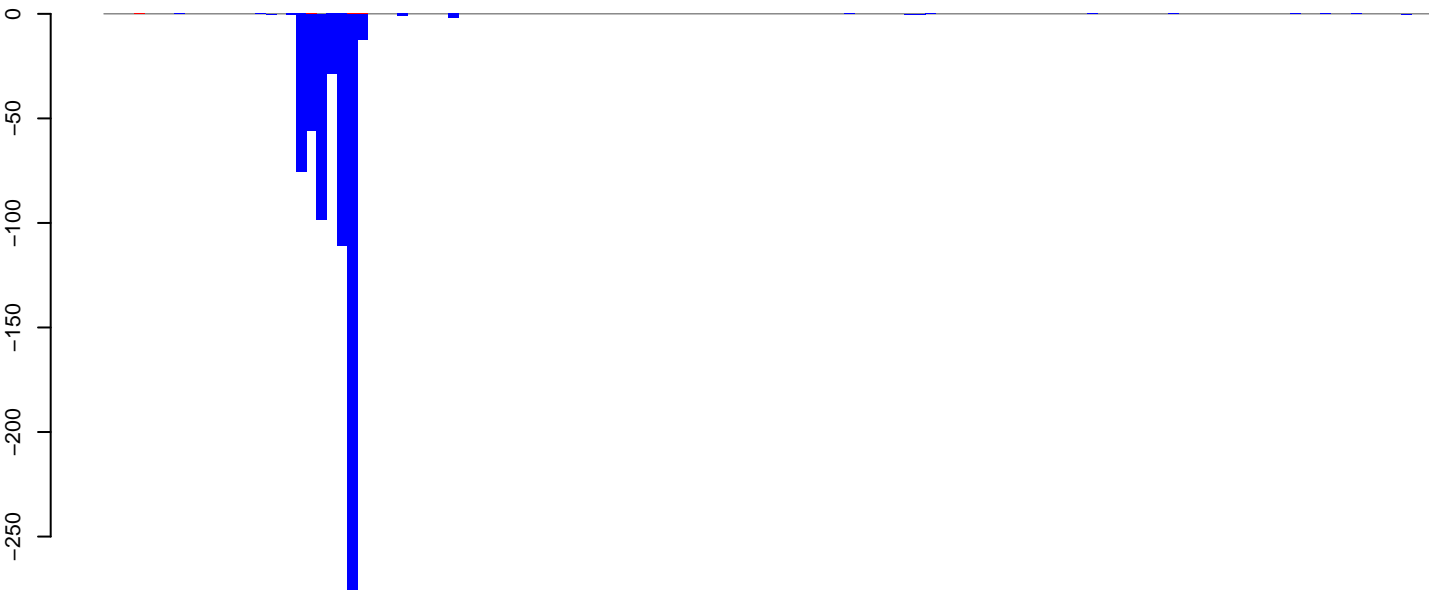
positive=0, negative=34, total=34

AnGam_Mos55_cells.24_35.rep



positive=1, negative=638, total=639

AnGam_Mos55_cells.rep

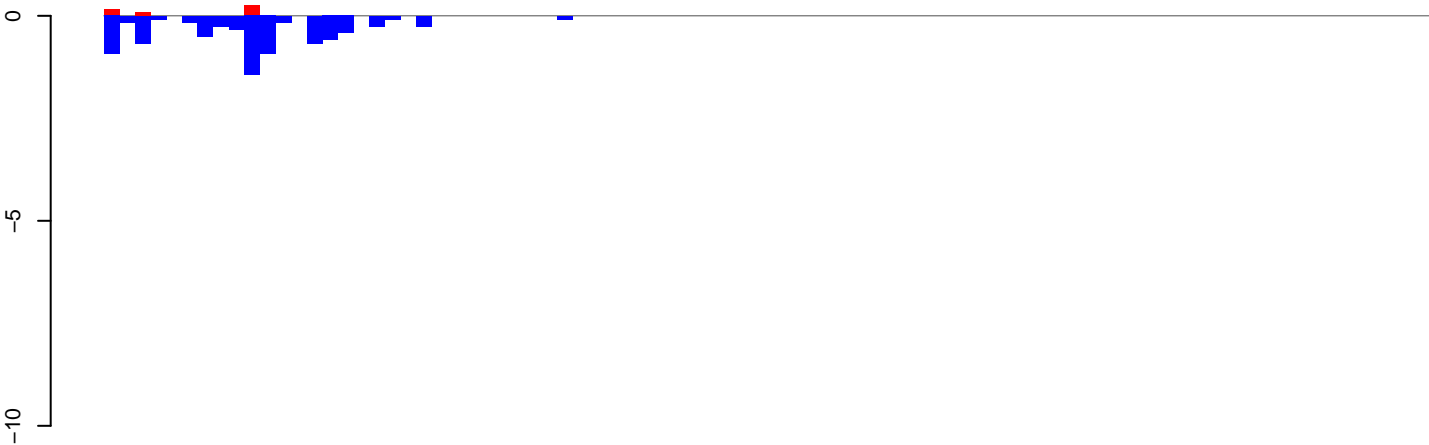


positive=1, negative=672, total=673

Window size=50, length=6592, TE@GYPSY11-LTR-L_AG:1-6592

0 1000 2000 3000 4000 5000 6000

AnGam_Mos55_cells.18_23.rep



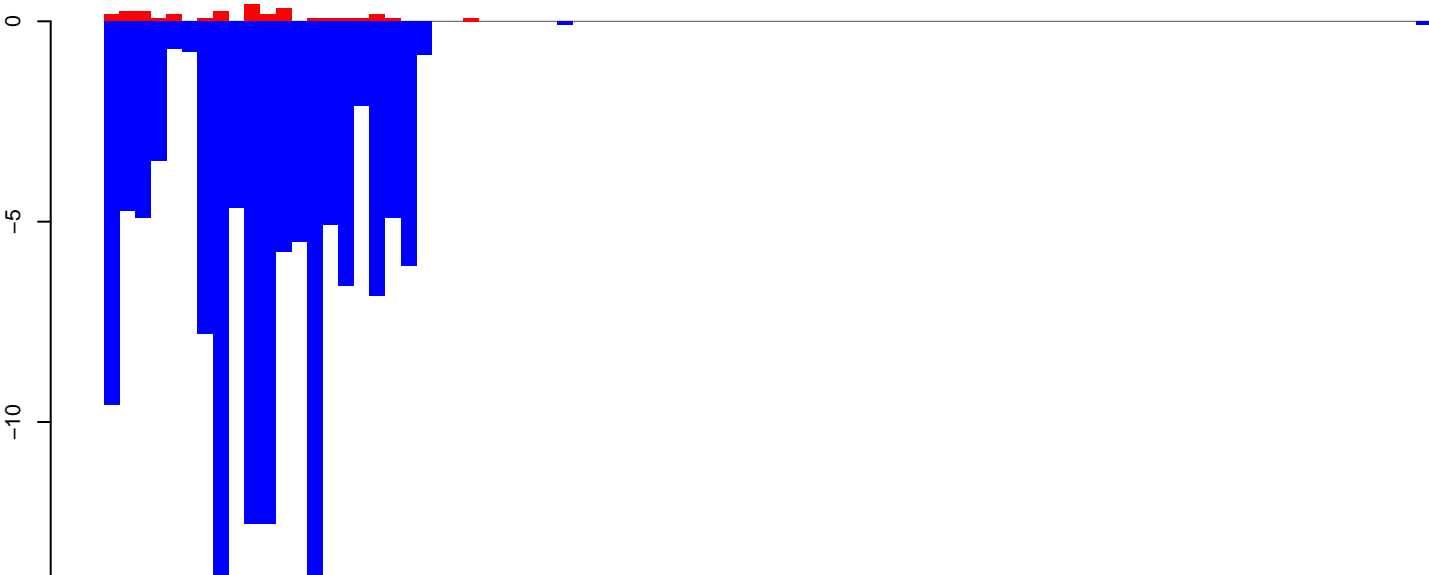
positive=1, negative=8, total=9

AnGam_Mos55_cells.24_35.rep



positive=2, negative=127, total=129

AnGam_Mos55_cells.rep

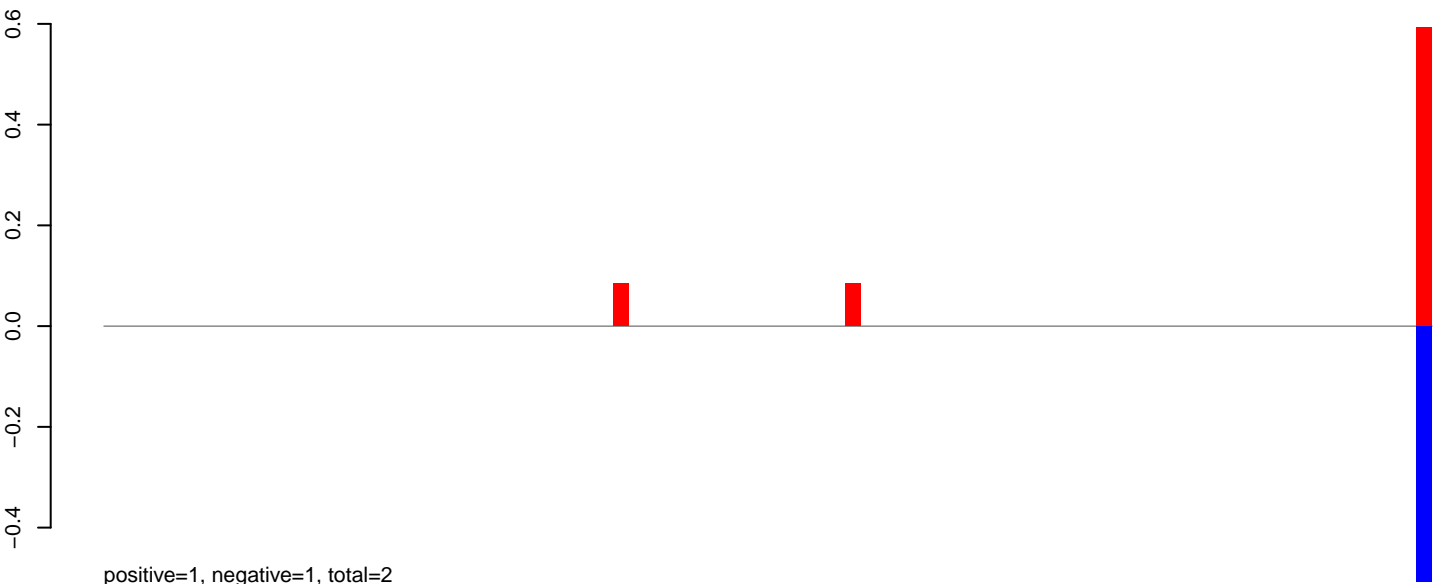


positive=3, negative=135, total=138

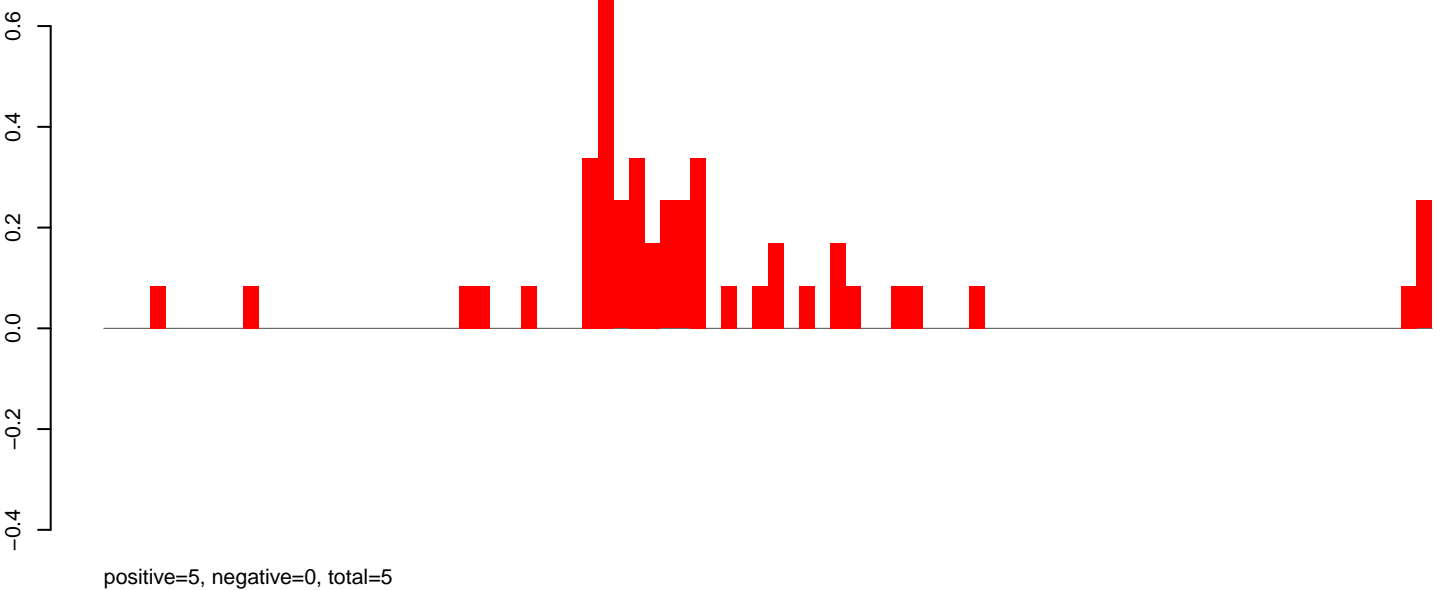
Window size=50, length=4269, TE@Ag-Jock-12:1-4269

0 1000 2000 3000 4000

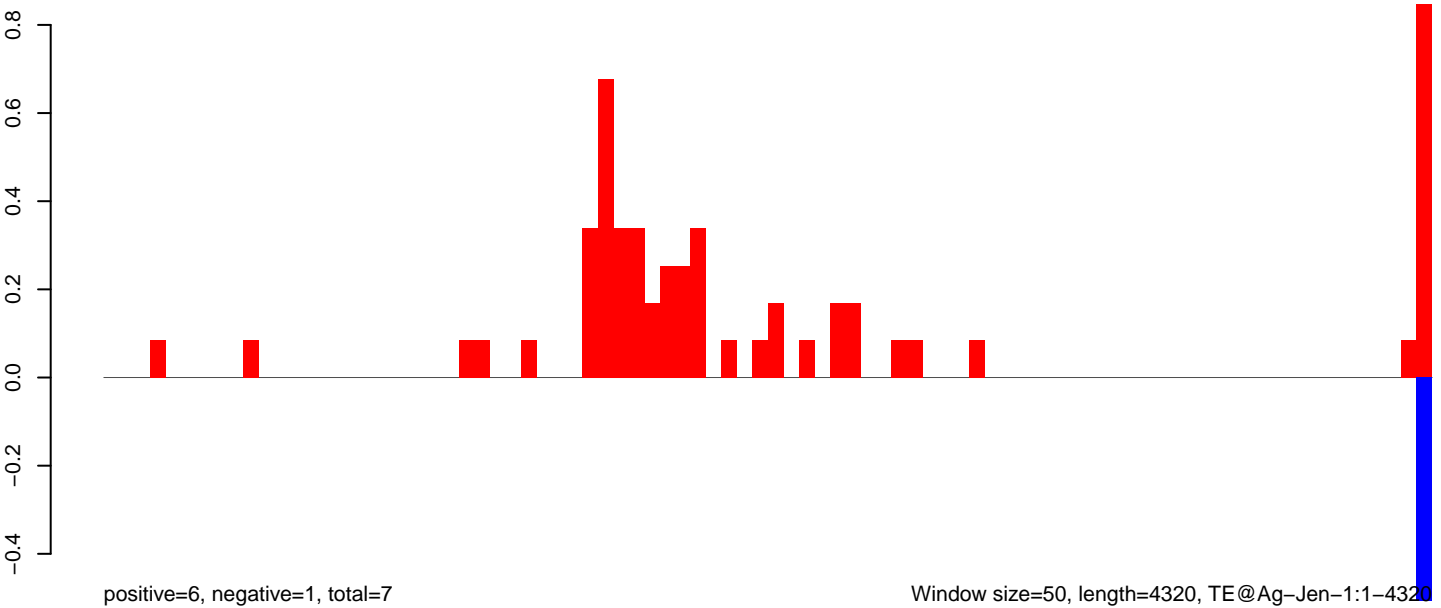
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



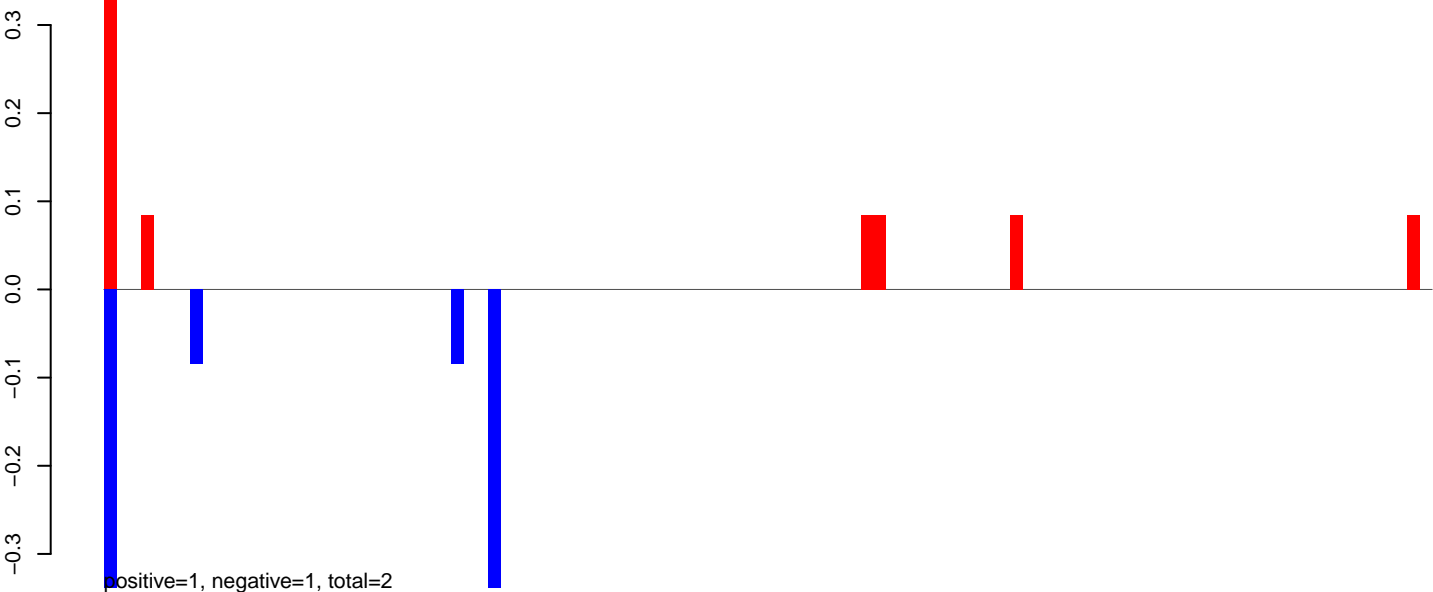
AnGam_Mos55_cells.rep



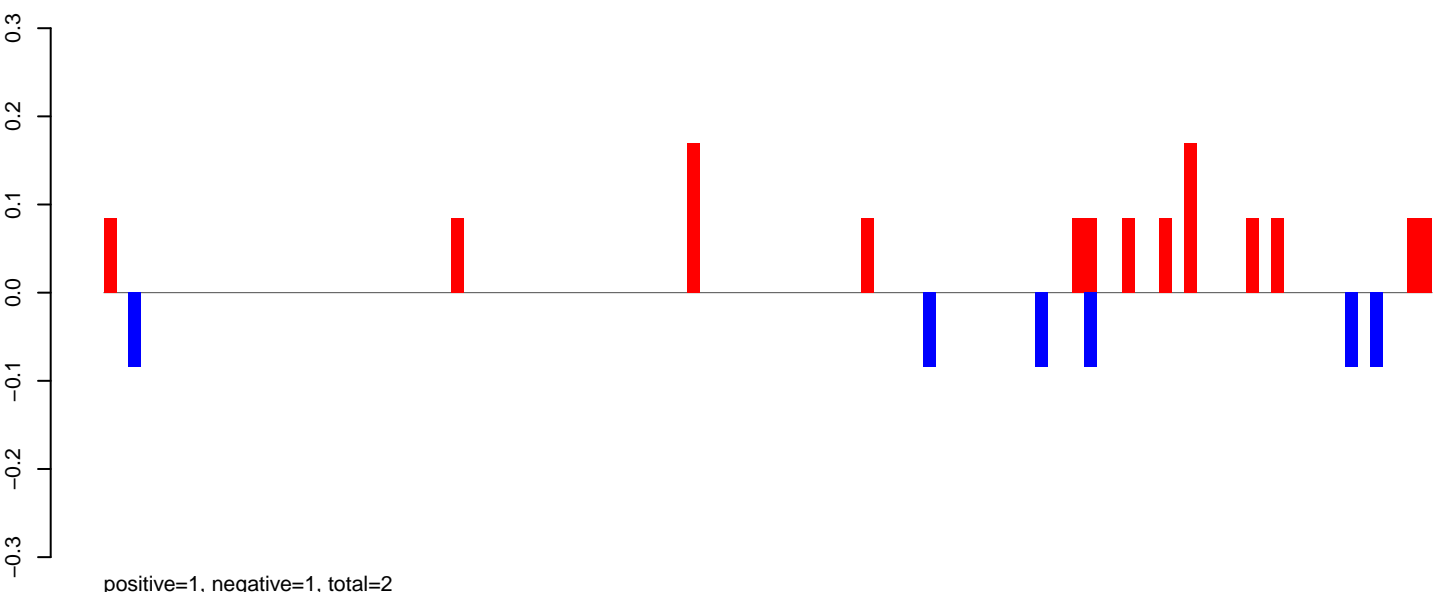
Window size=50, length=4320, TE@Ag-Jen-1:1-4320

0 1000 2000 3000 4000

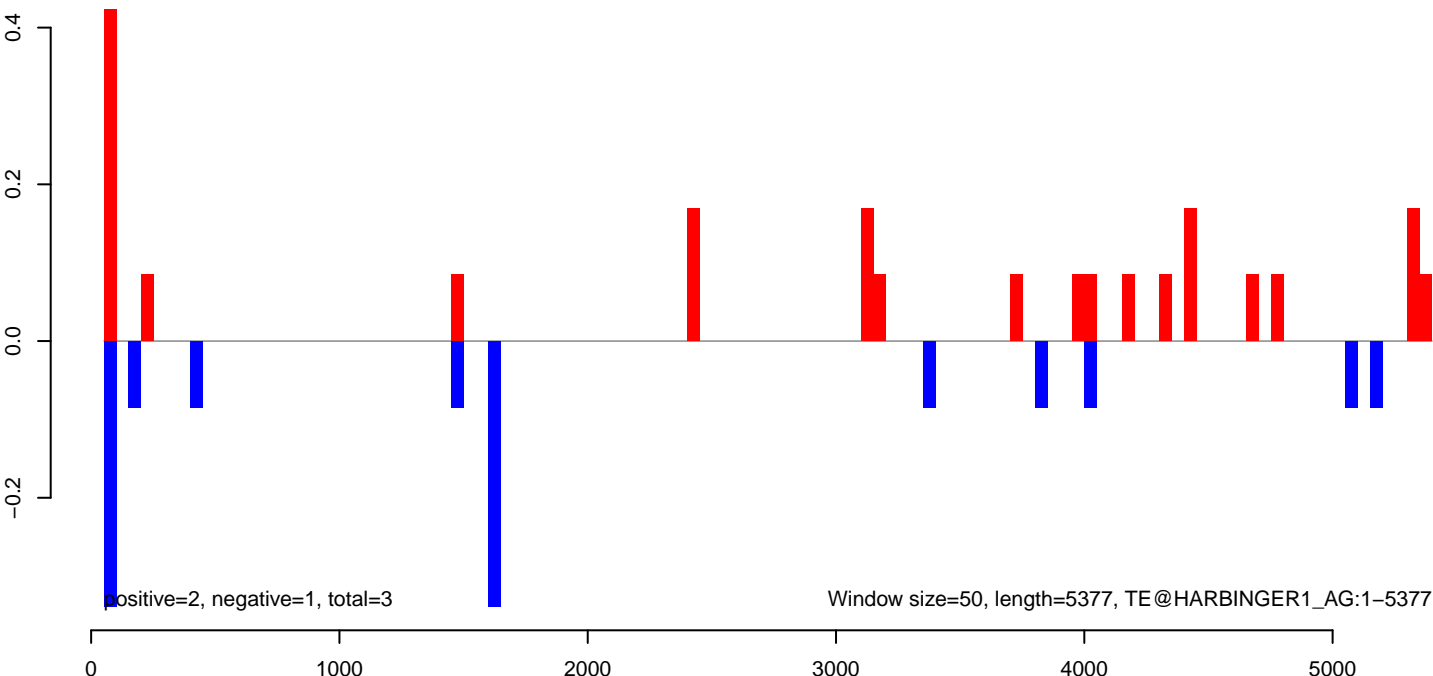
AnGam_Mos55_cells.18_23.rep



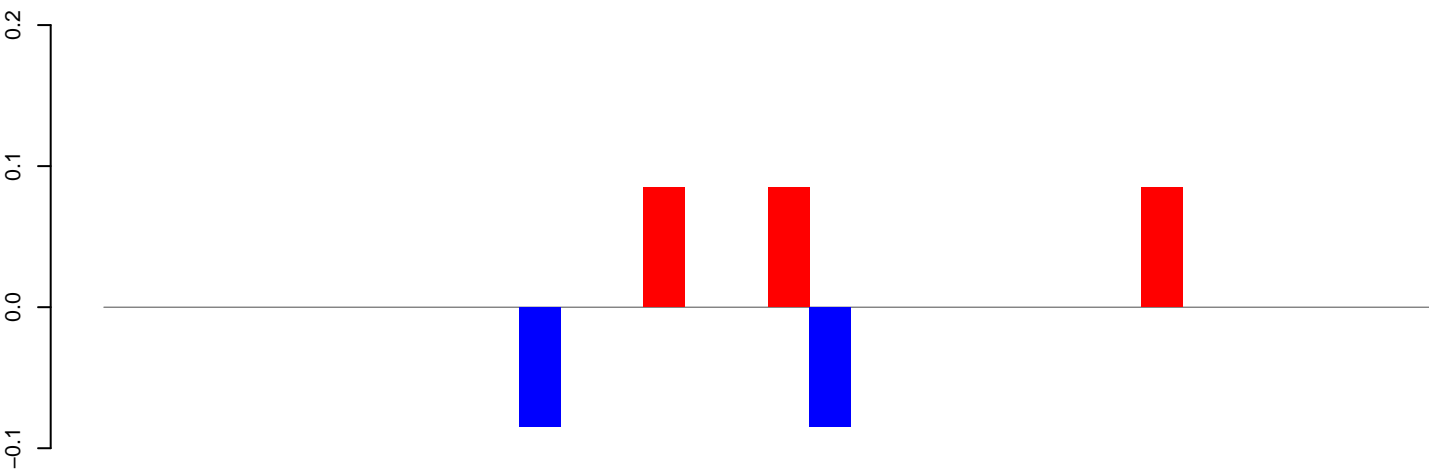
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

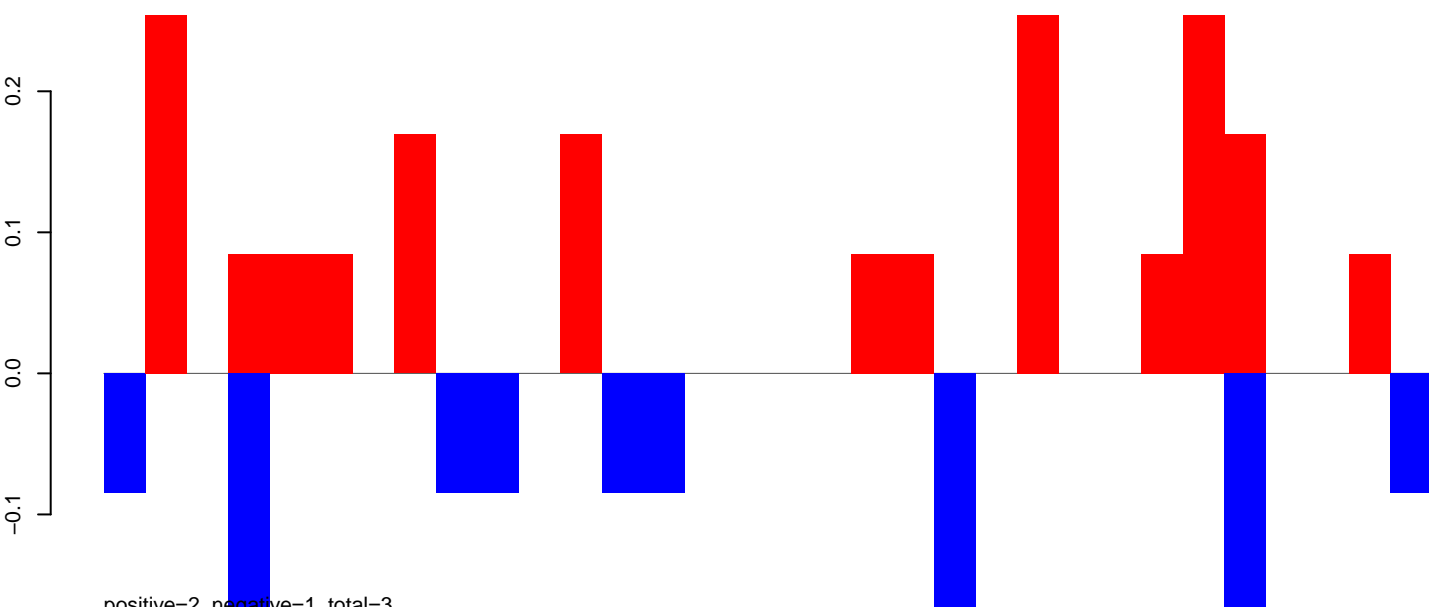


AnGam_Mos55_cells.18_23.rep



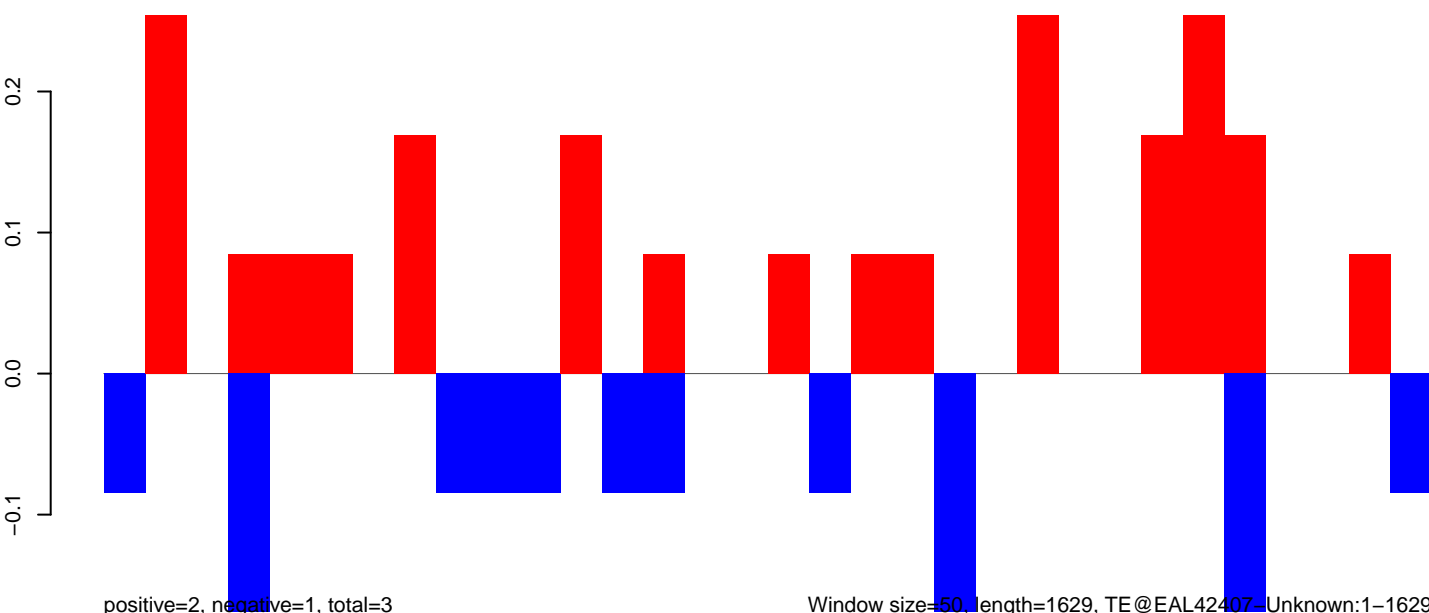
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=2, negative=1, total=3

AnGam_Mos55_cells.rep

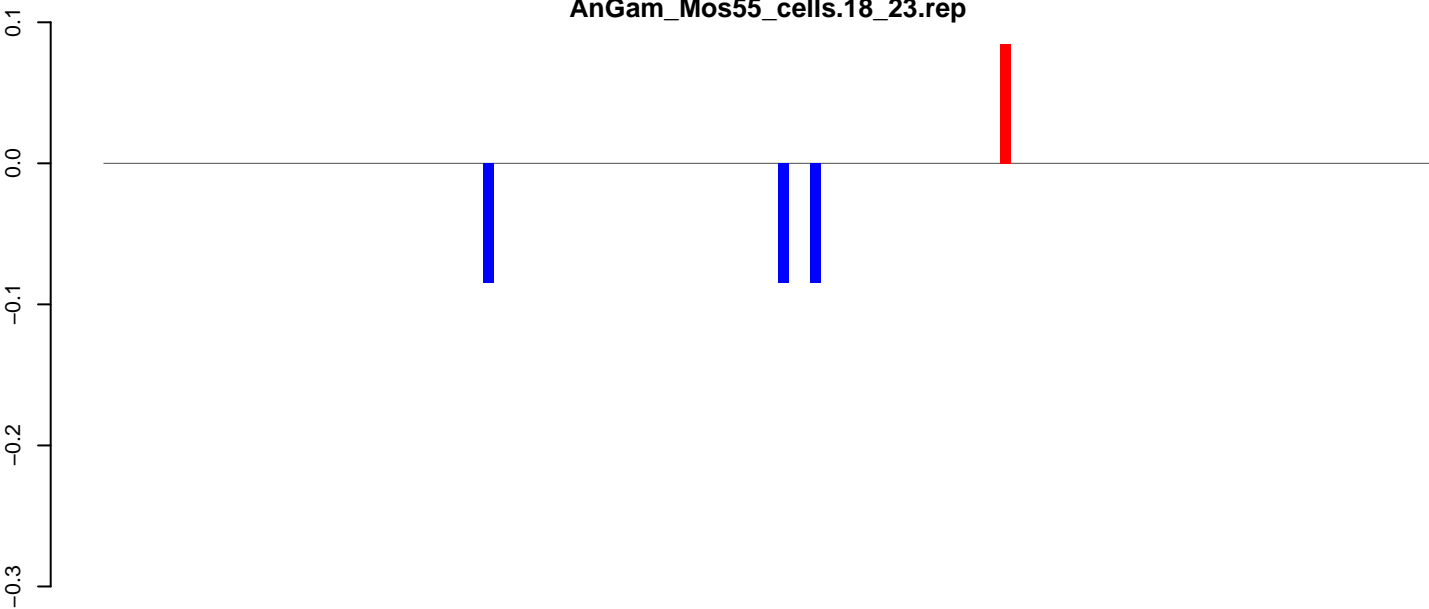


positive=2, negative=1, total=3

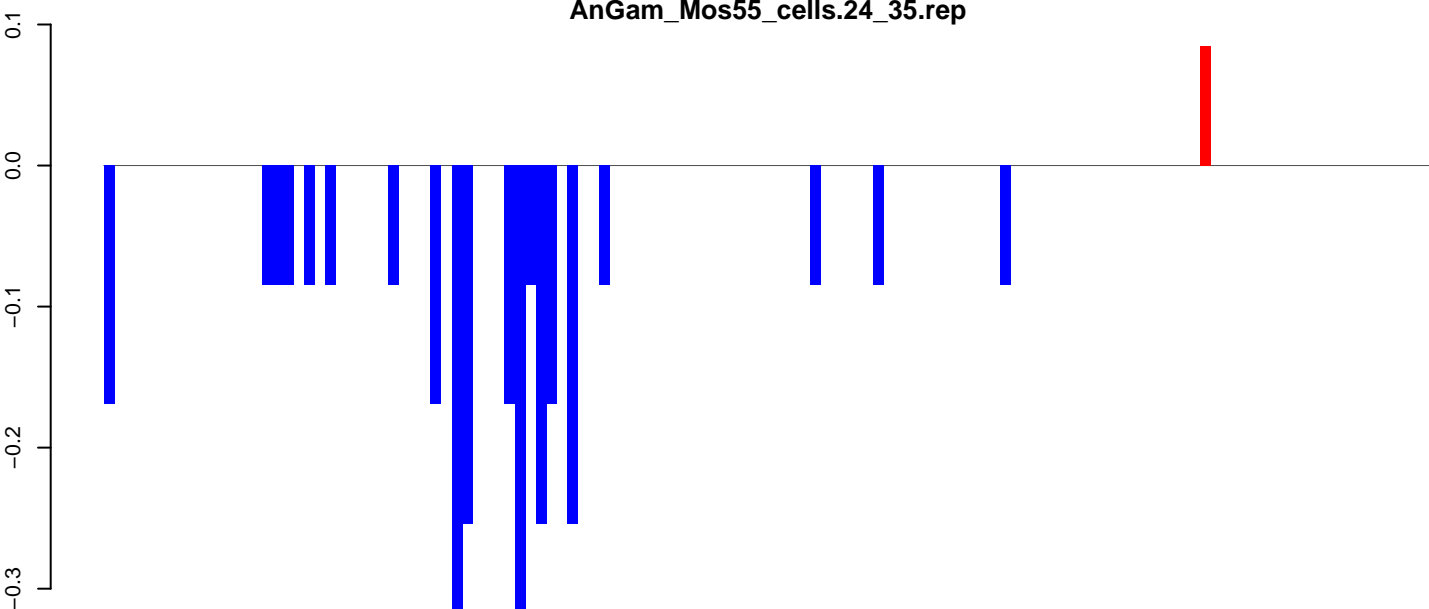
Window size=50, length=1629, TE@EAL42407-Unknown:1-1629

0 500 1000 1500

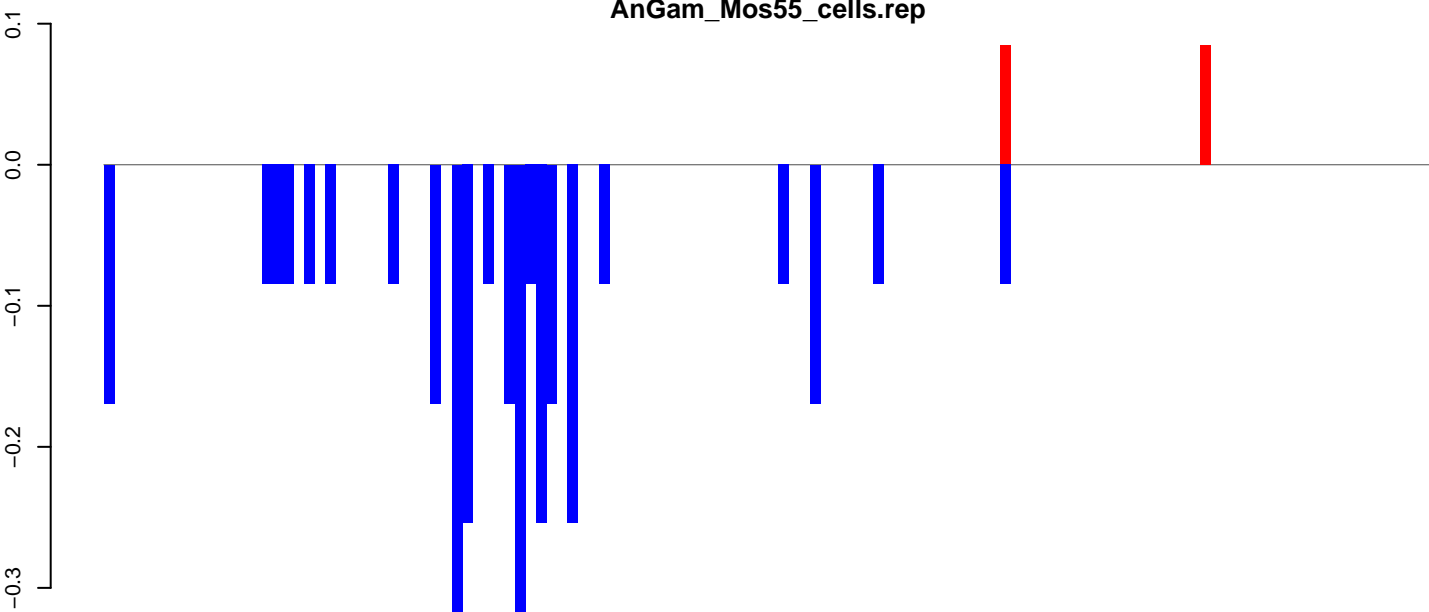
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



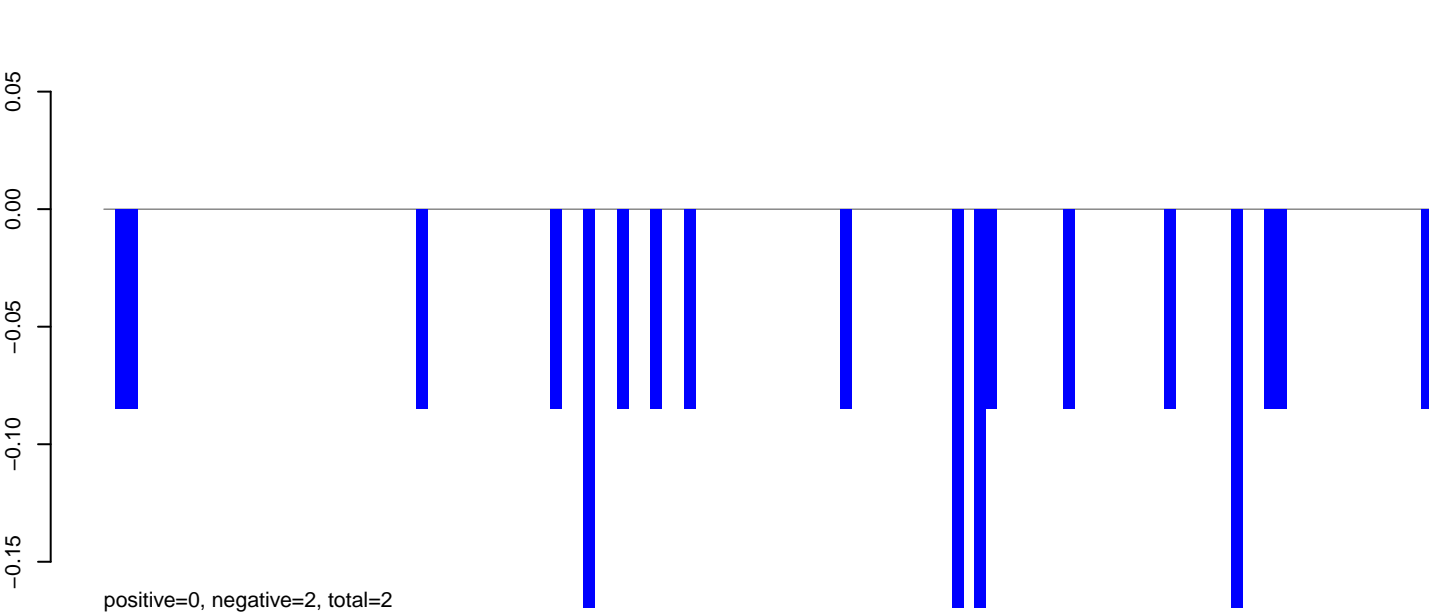
Window size=50, length=6343, TE@Ag-Loner-1:1-6343

0 1000 2000 3000 4000 5000 6000

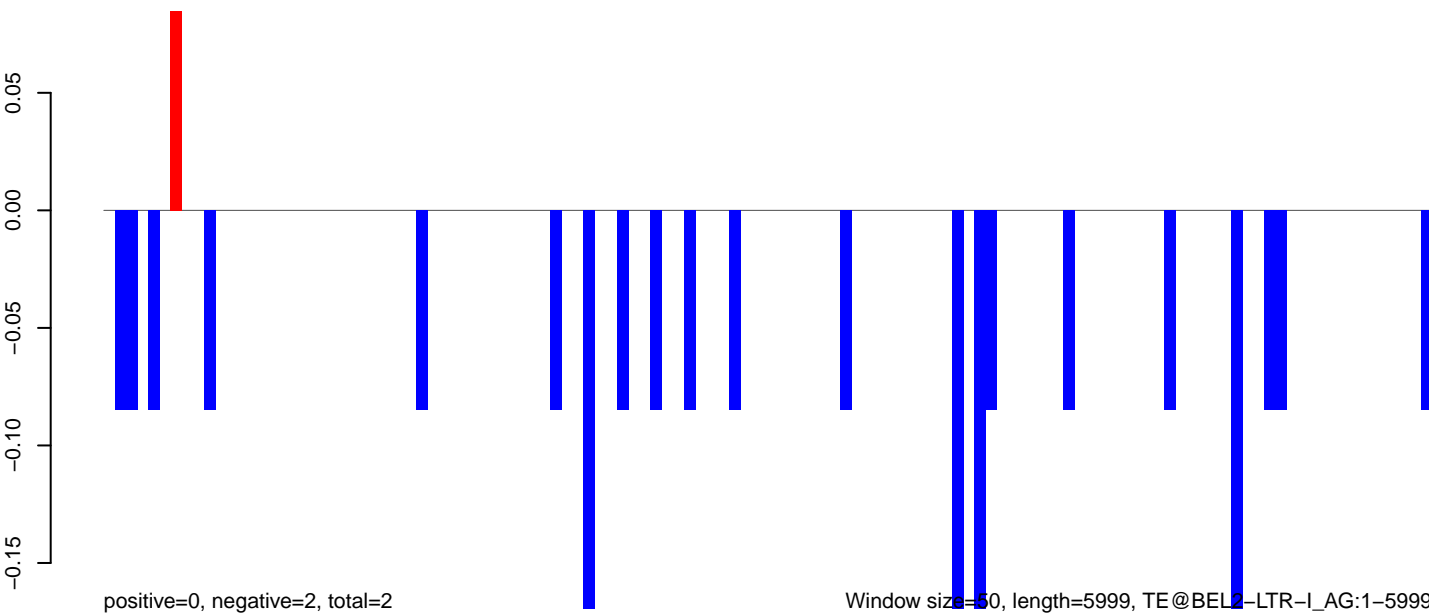
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



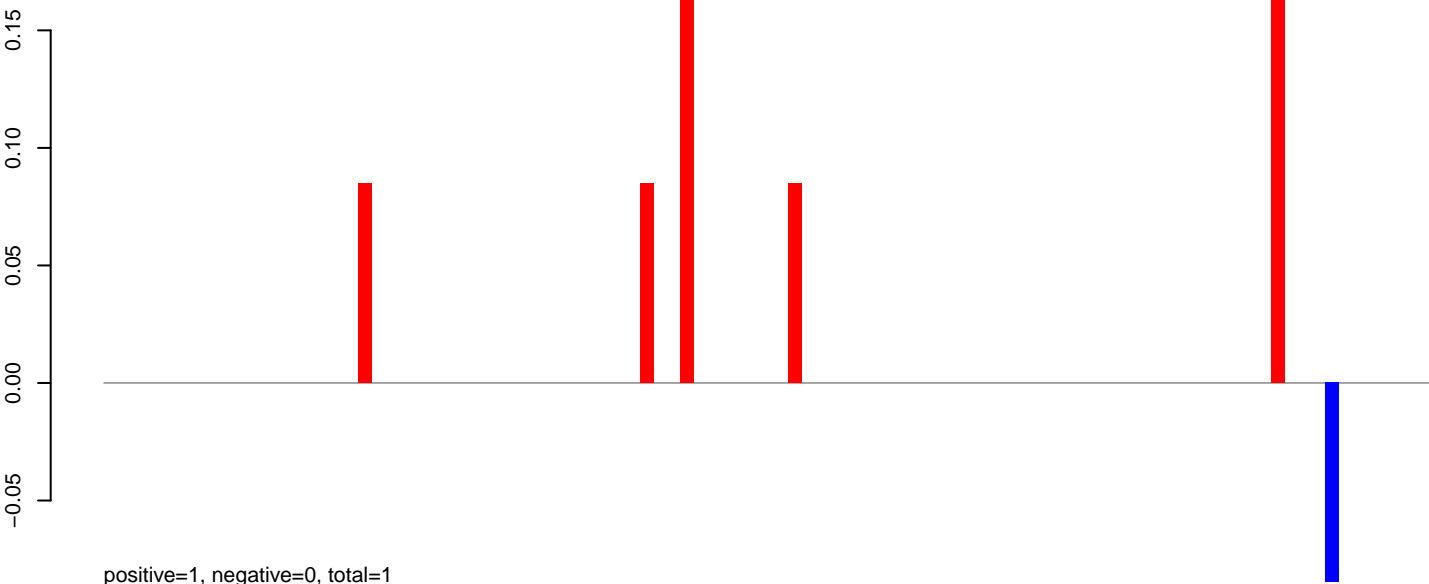
AnGam_Mos55_cells.rep



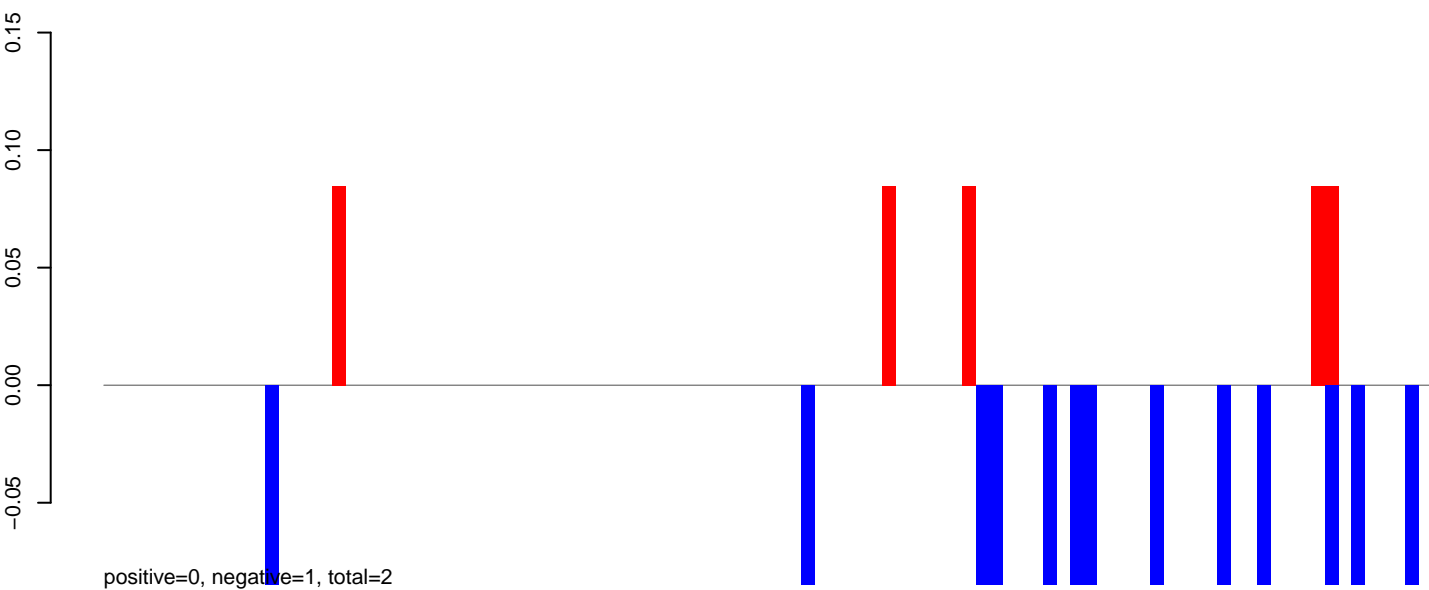
Window size=50, length=5999, TE@BEL2-LTR-I_AG:1-5999

0 1000 2000 3000 4000 5000 6000

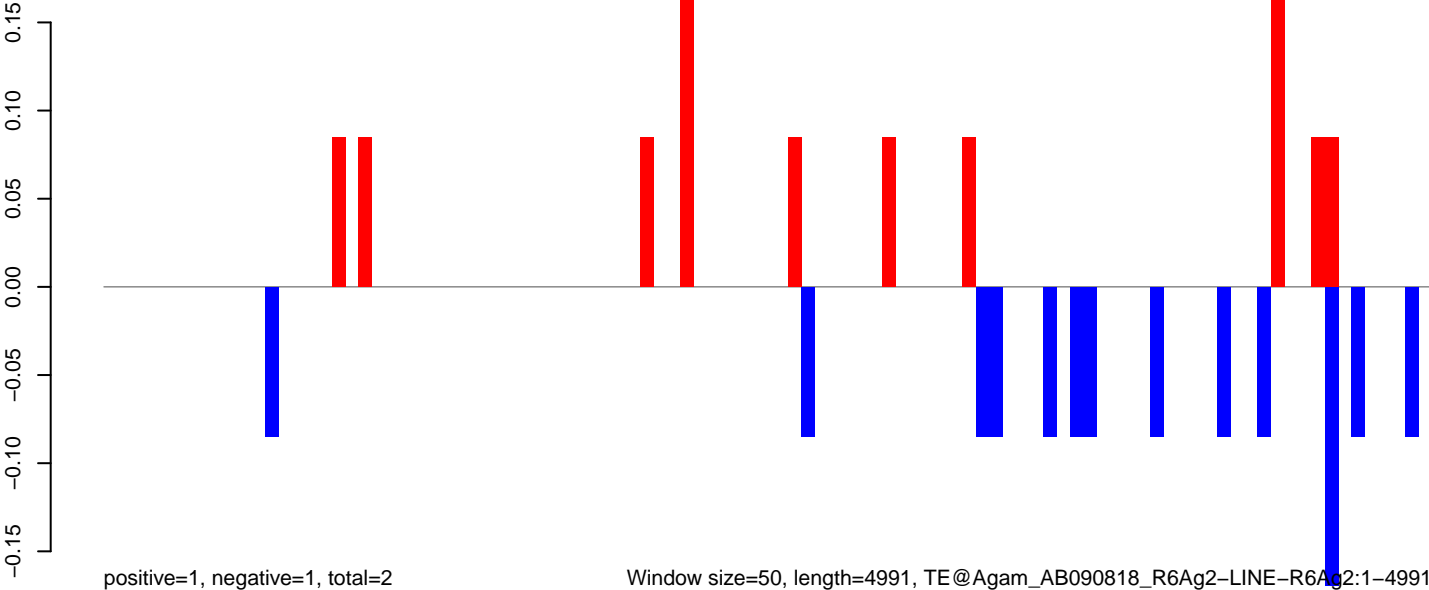
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

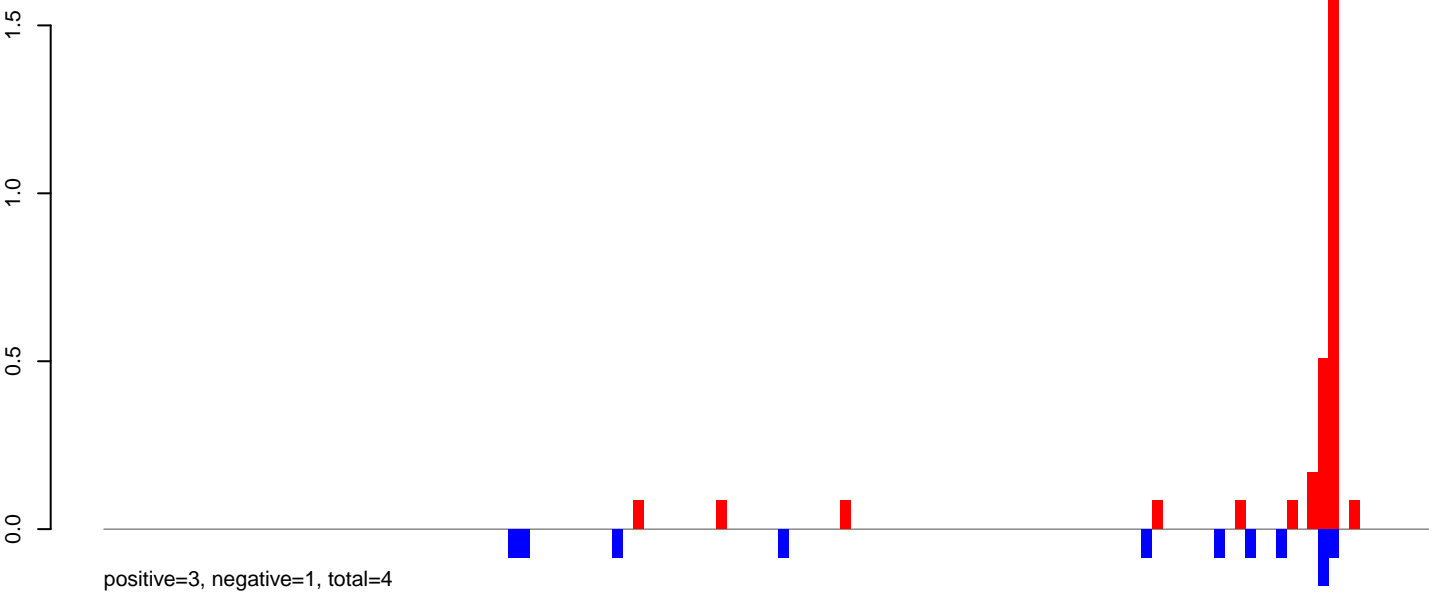


AnGam_Mos55_cells.rep

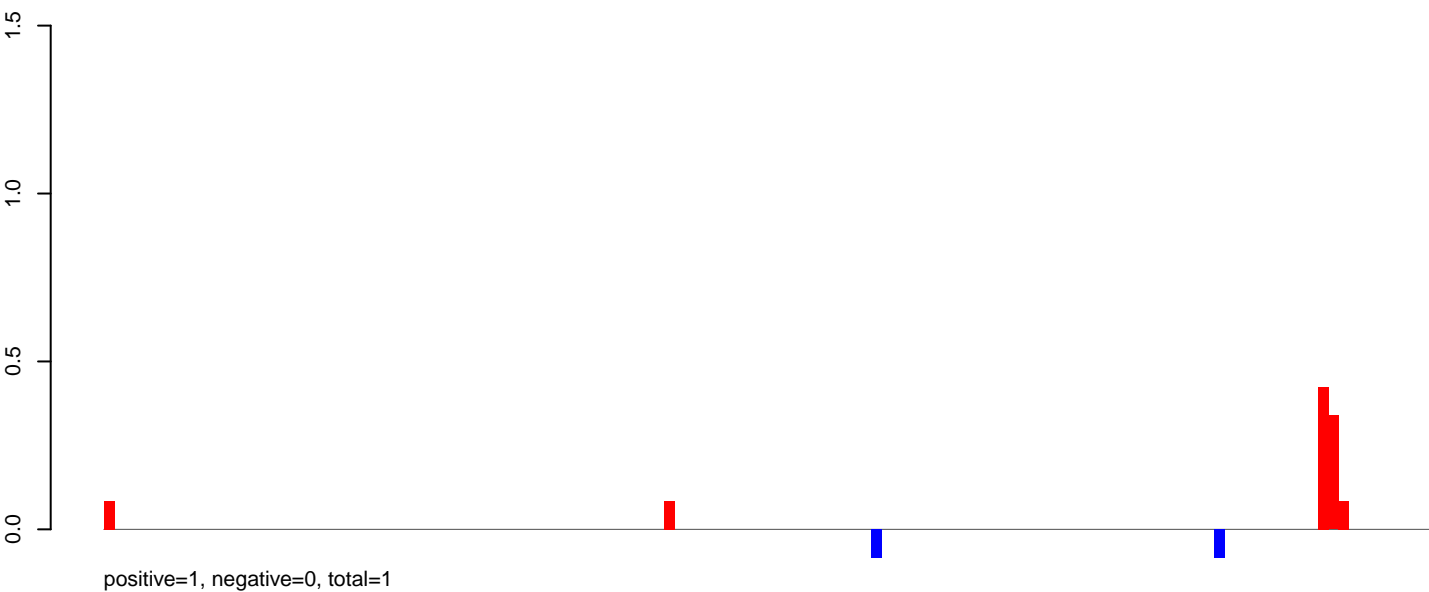


Window size=50, length=4991, TE@Agam_AB090818_R6Ag2-LINE-R6Ag2:1-4991

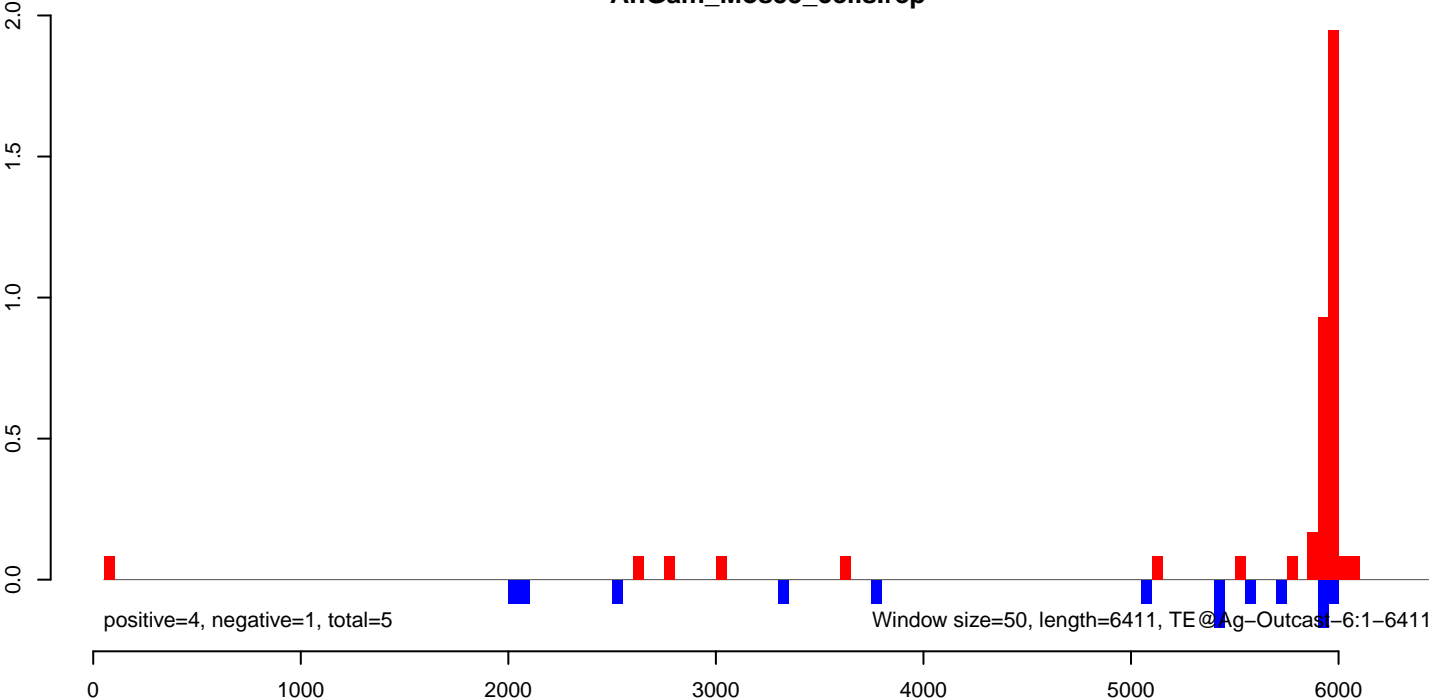
AnGam_Mos55_cells.18_23.rep



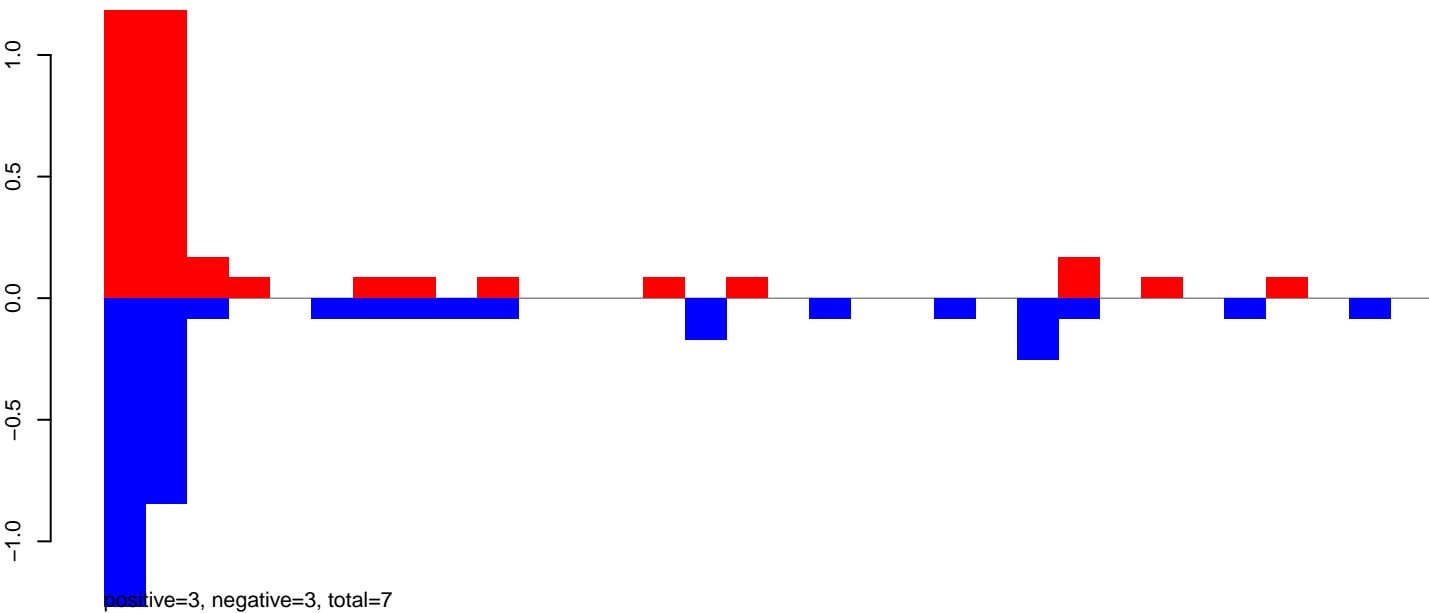
AnGam_Mos55_cells.24_35.rep



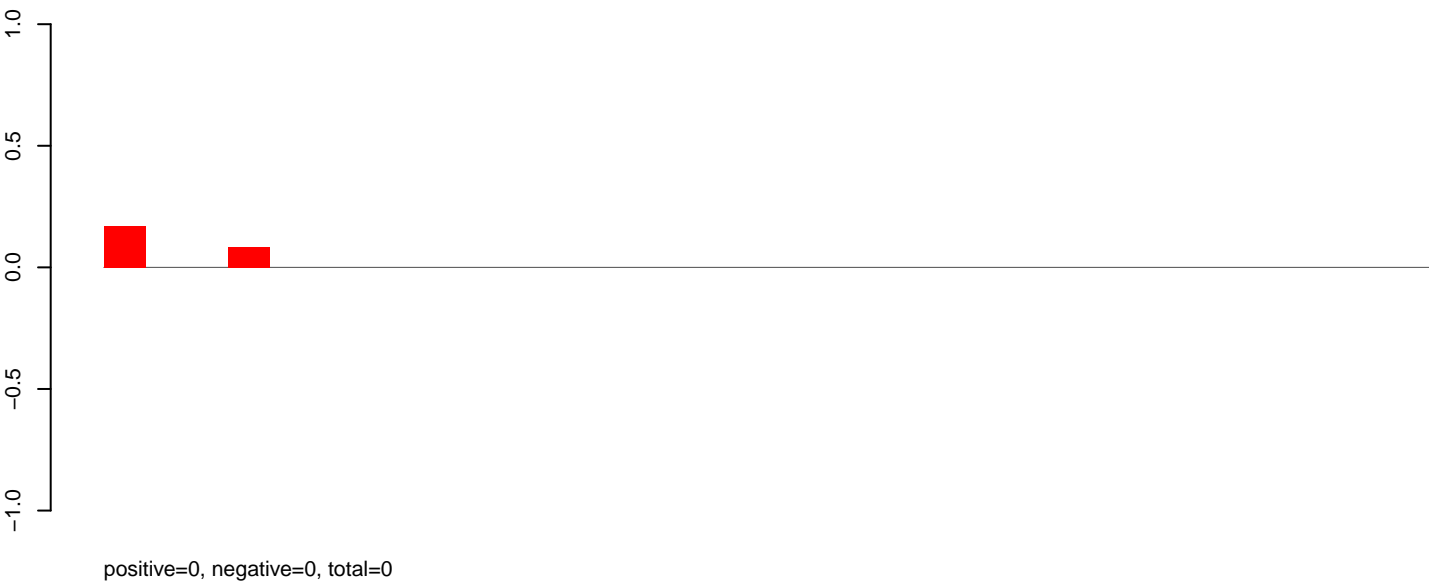
AnGam_Mos55_cells.rep



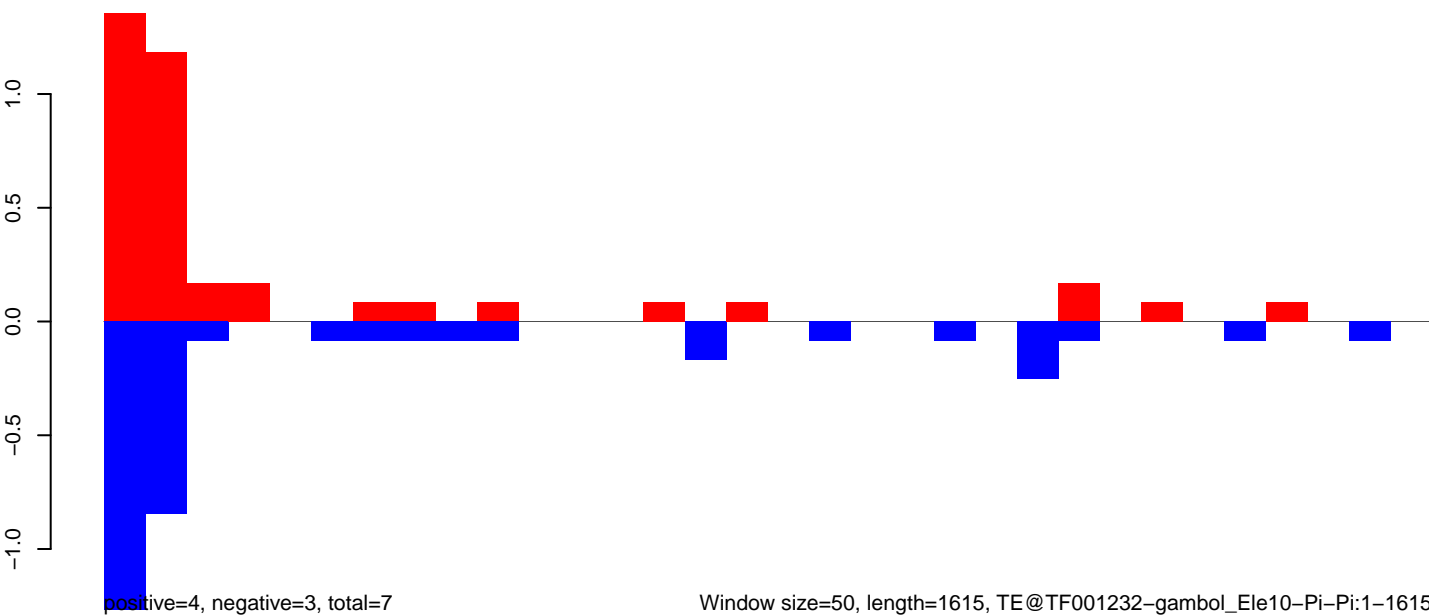
AnGam_Mos55_cells.18_23.rep



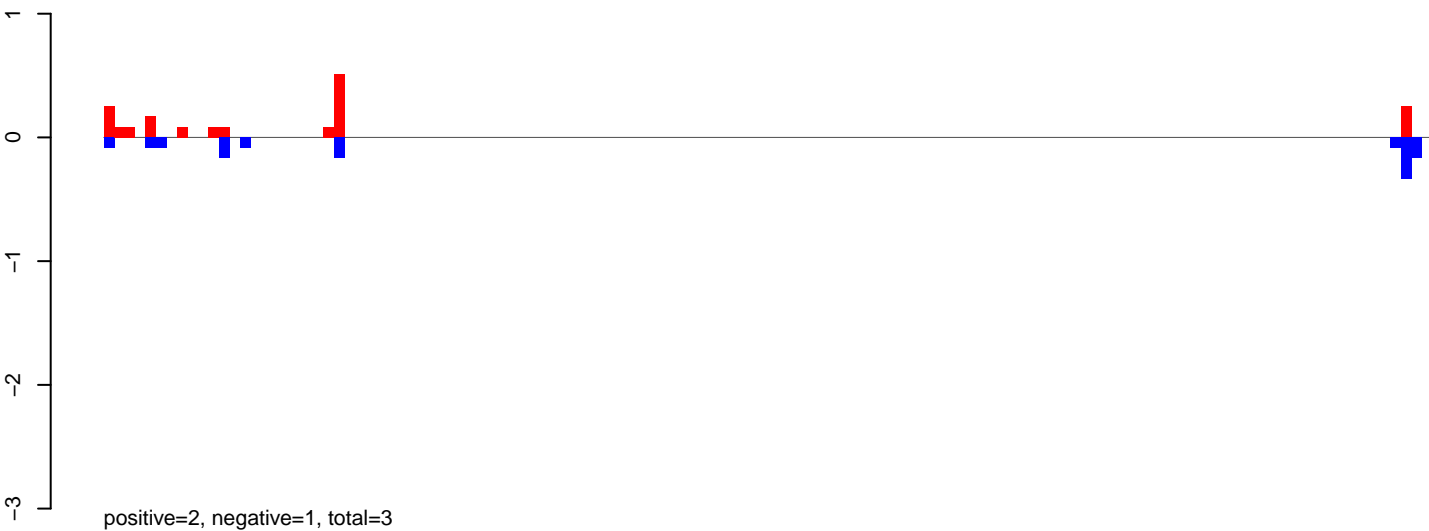
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



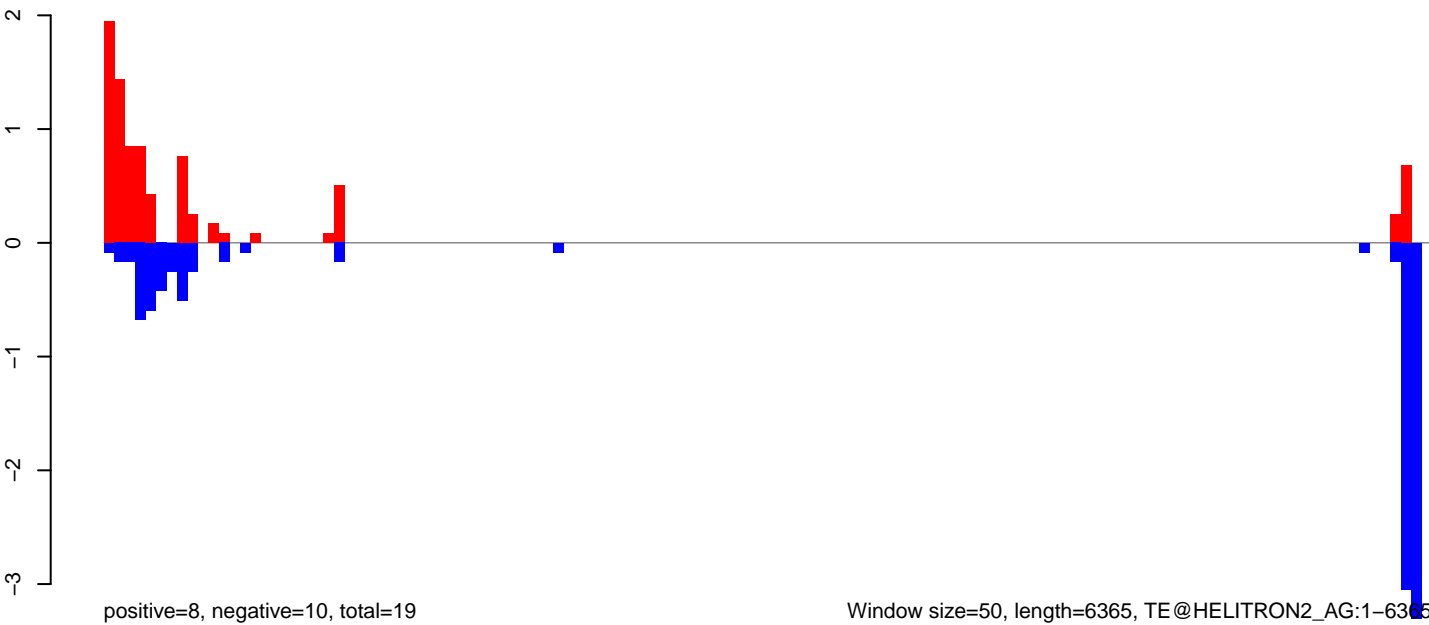
AnGam_Mos55_cells.18_23.rep



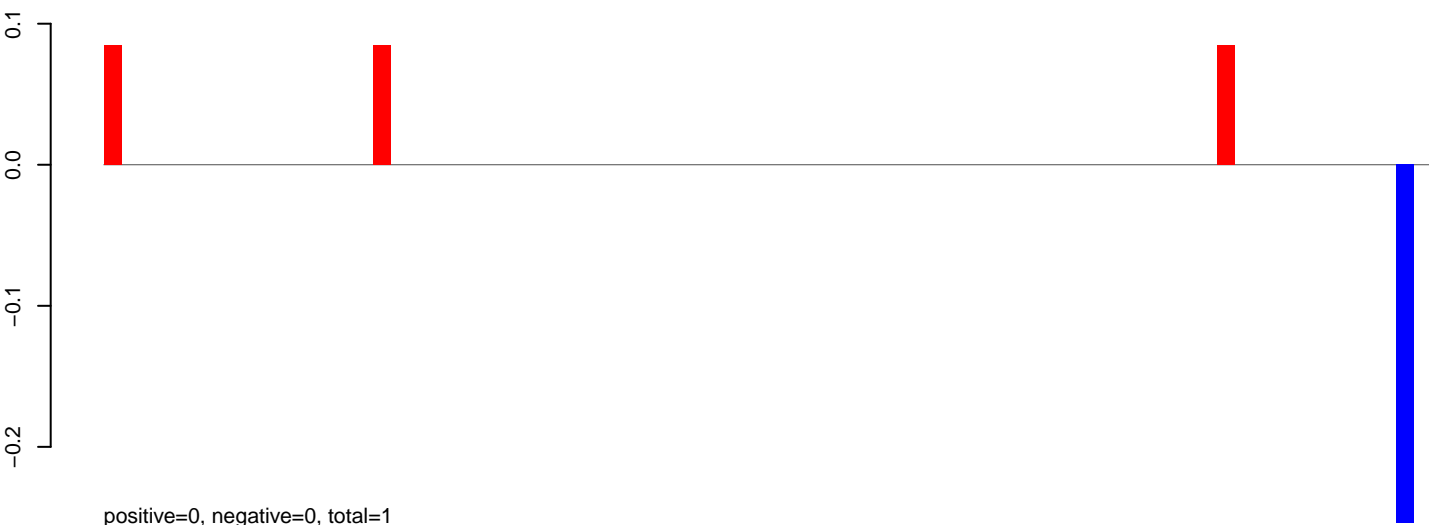
AnGam_Mos55_cells.24_35.rep



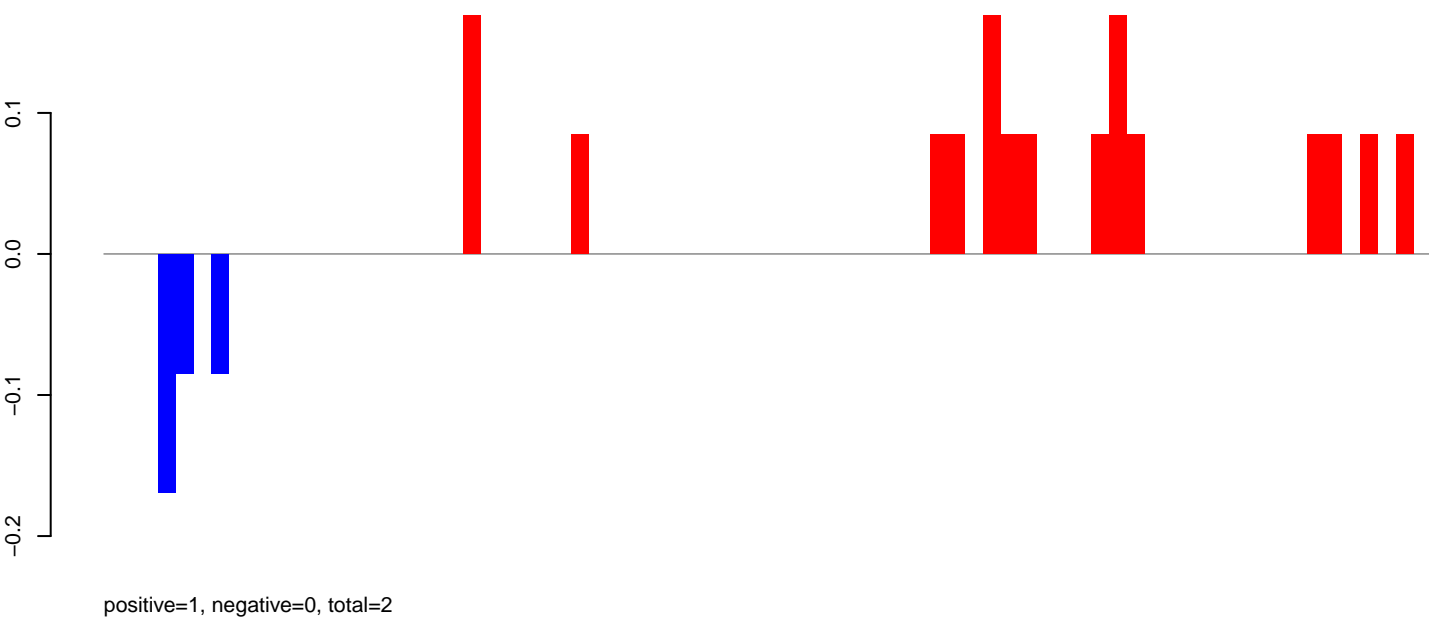
AnGam_Mos55_cells.rep



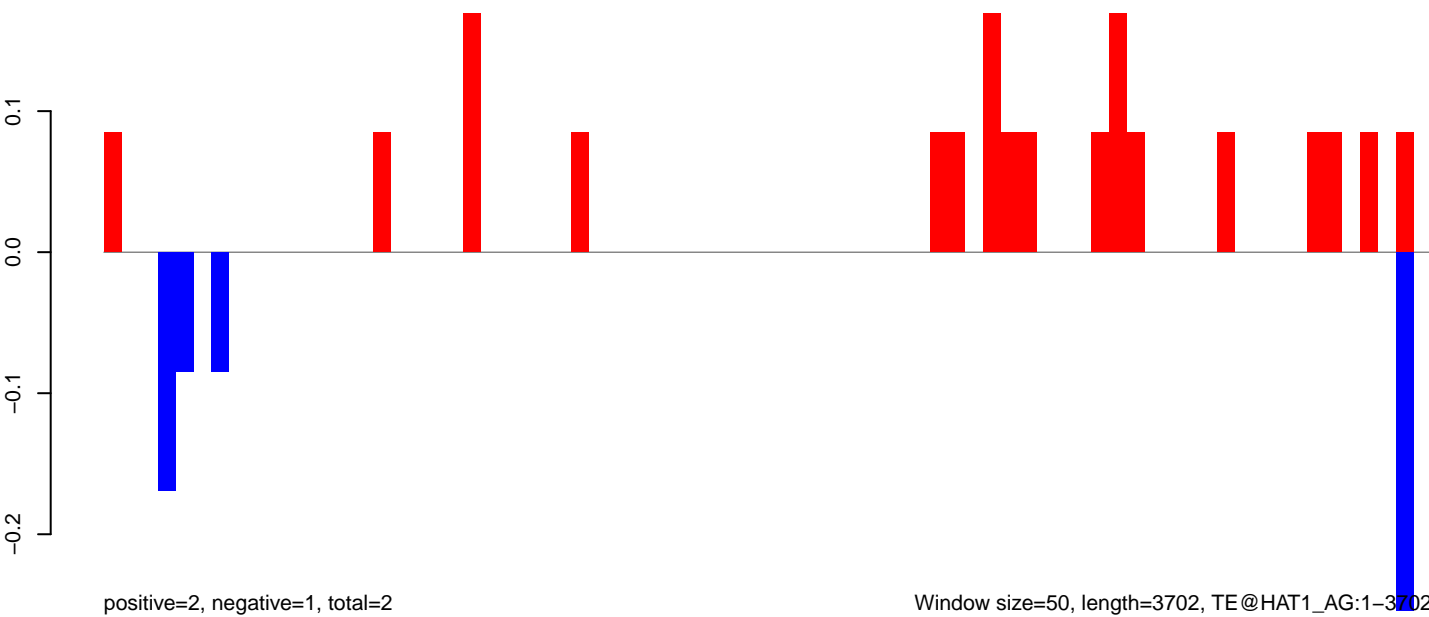
AnGam_Mos55_cells.18_23.rep



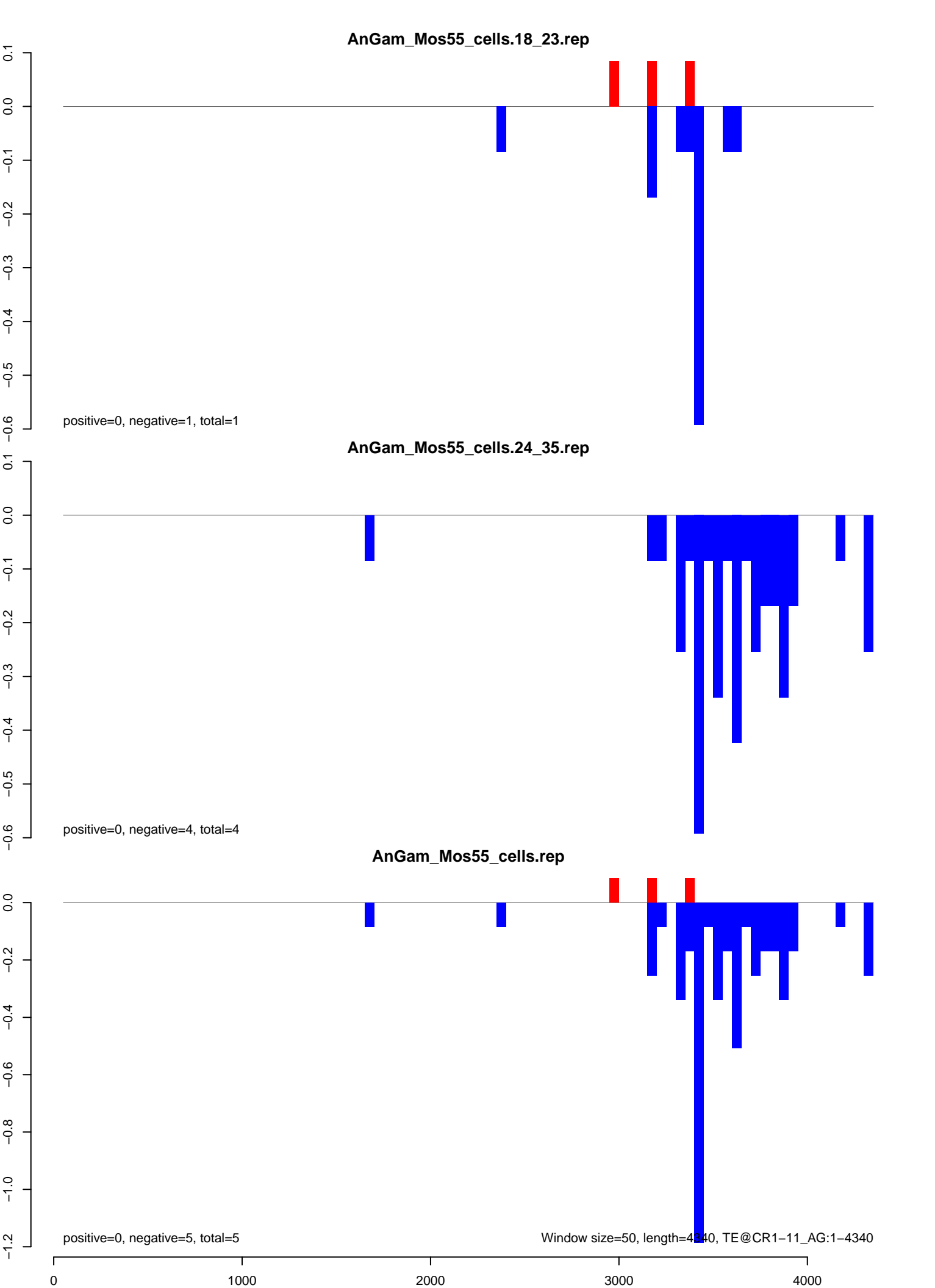
AnGam_Mos55_cells.24_35.rep



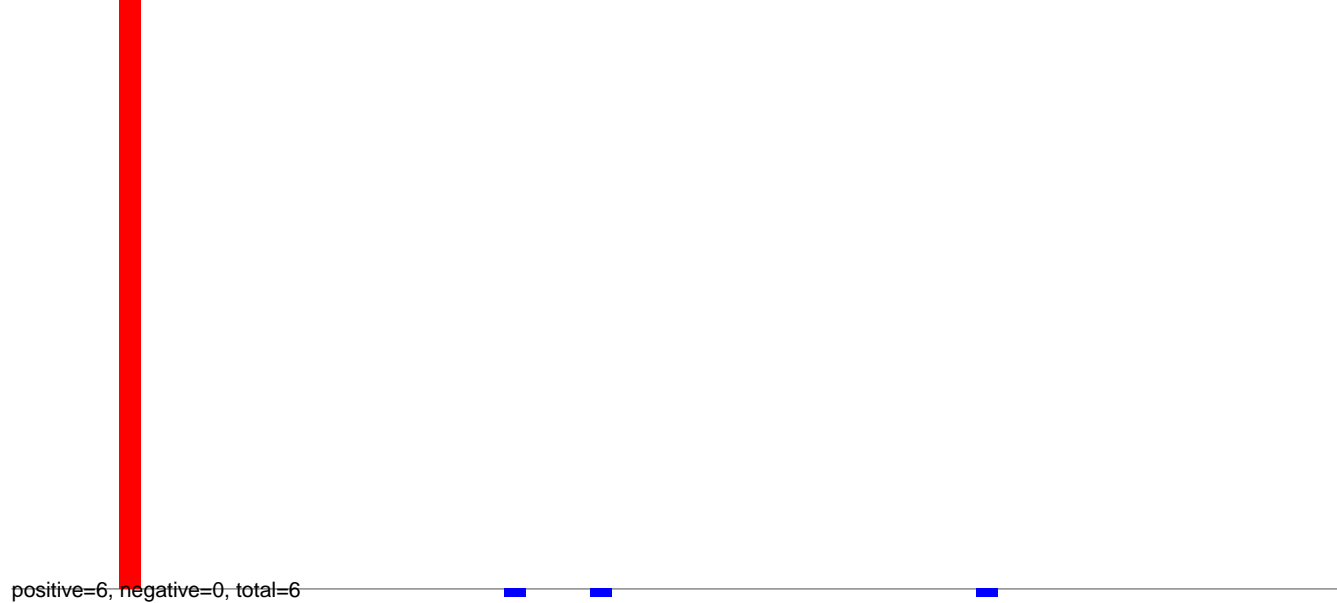
AnGam_Mos55_cells.rep



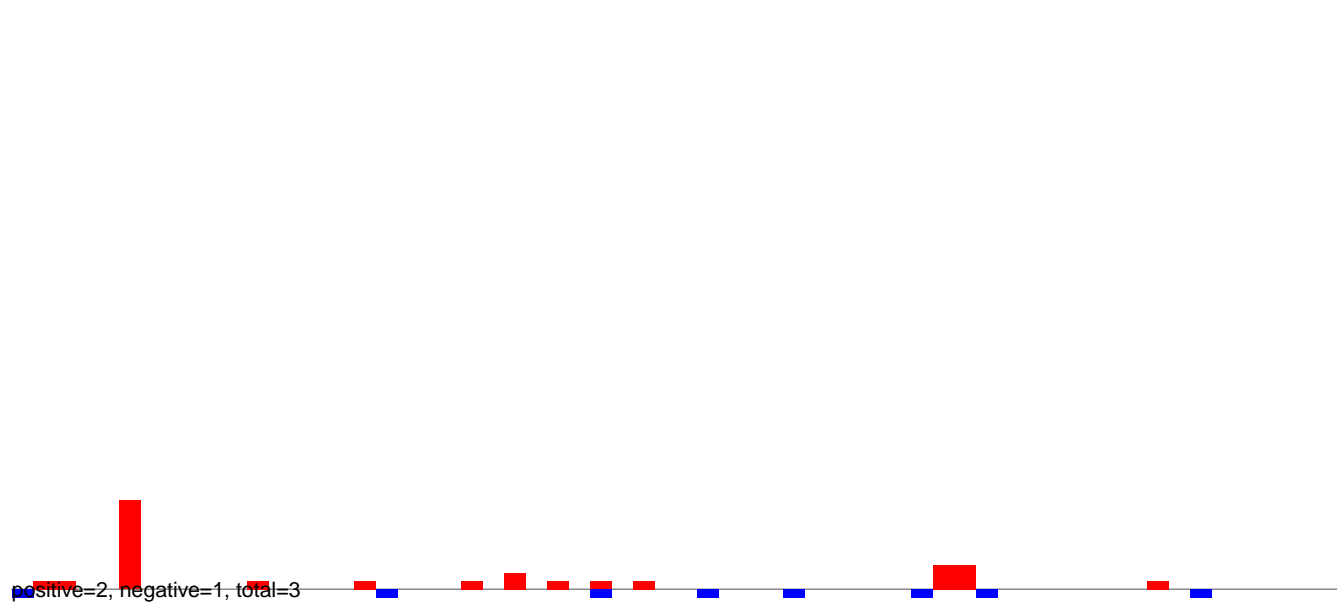
Window size=50, length=3702, TE@HAT1_AG:1-3702



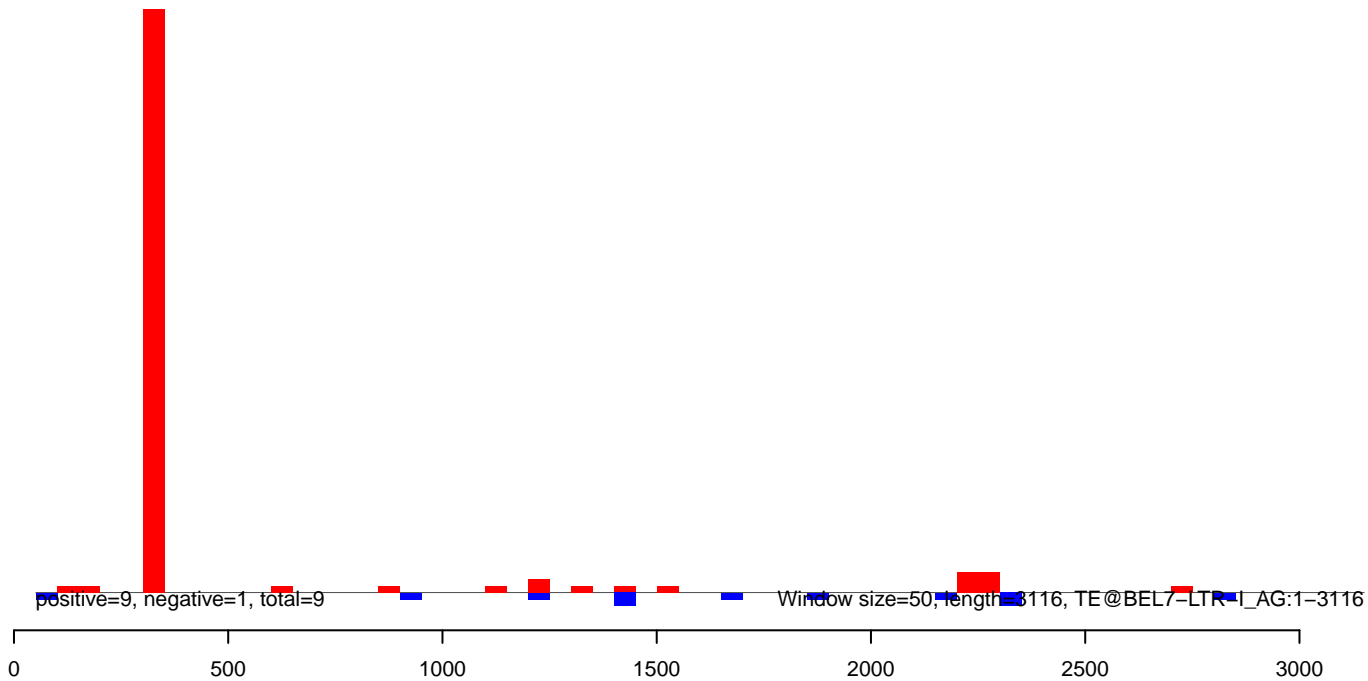
AnGam_Mos55_cells.18_23.rep



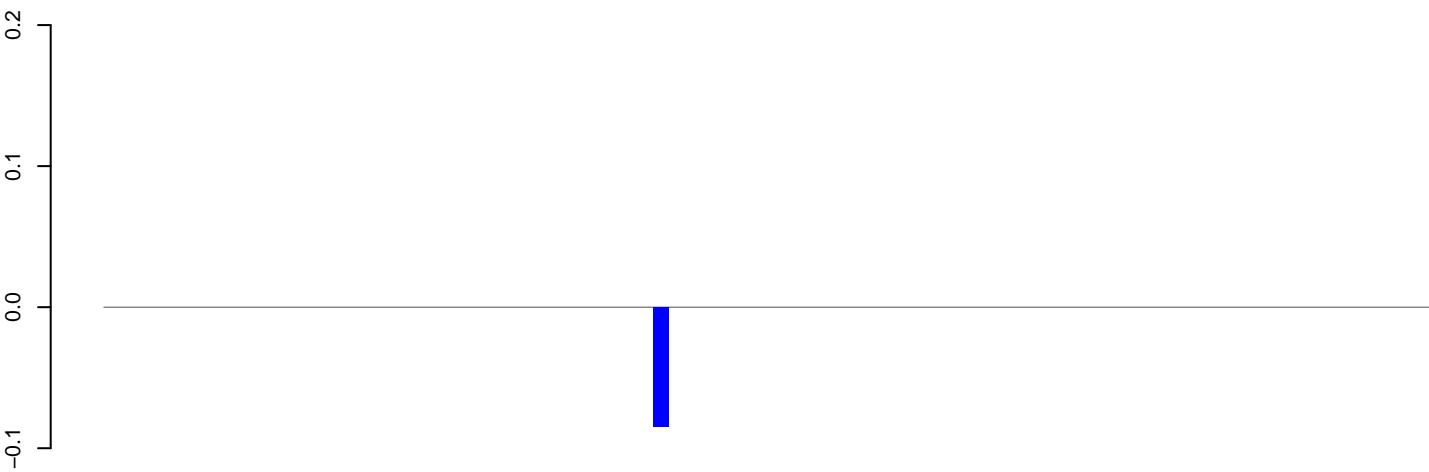
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

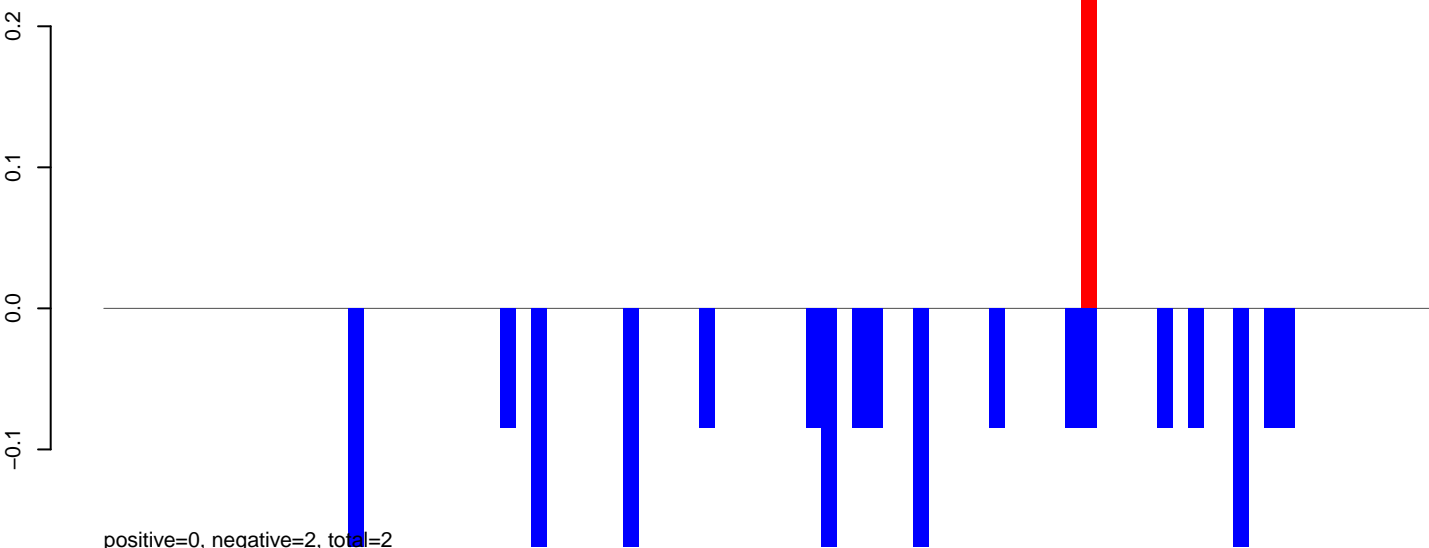


AnGam_Mos55_cells.18_23.rep



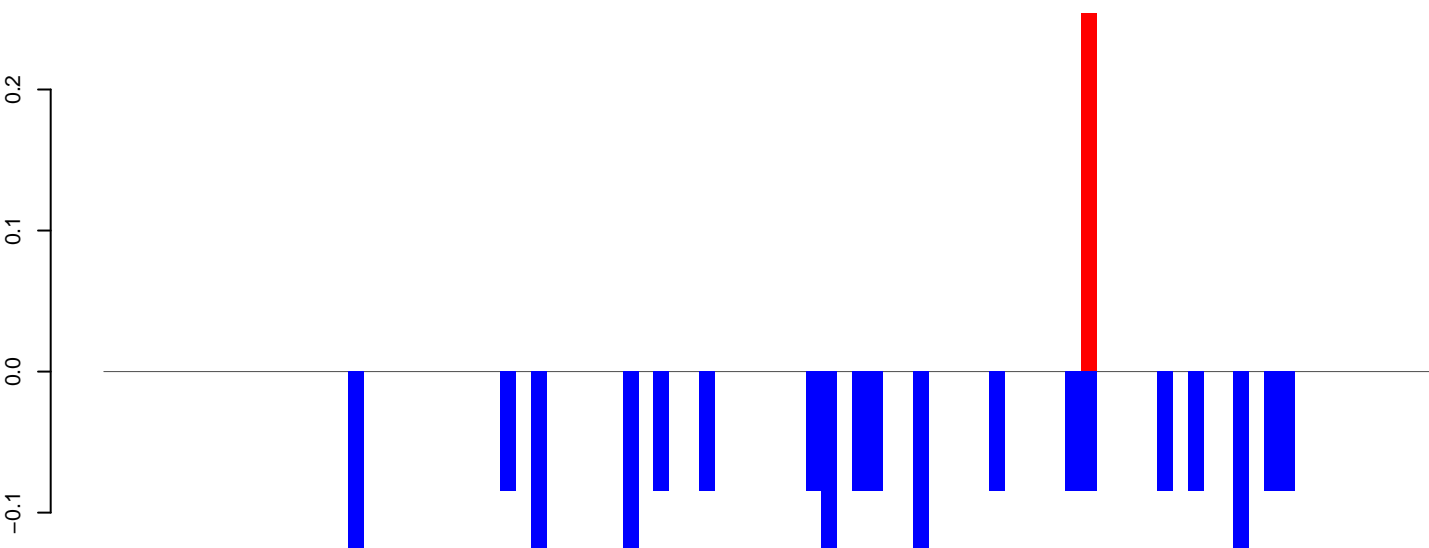
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=2, total=2

AnGam_Mos55_cells.rep

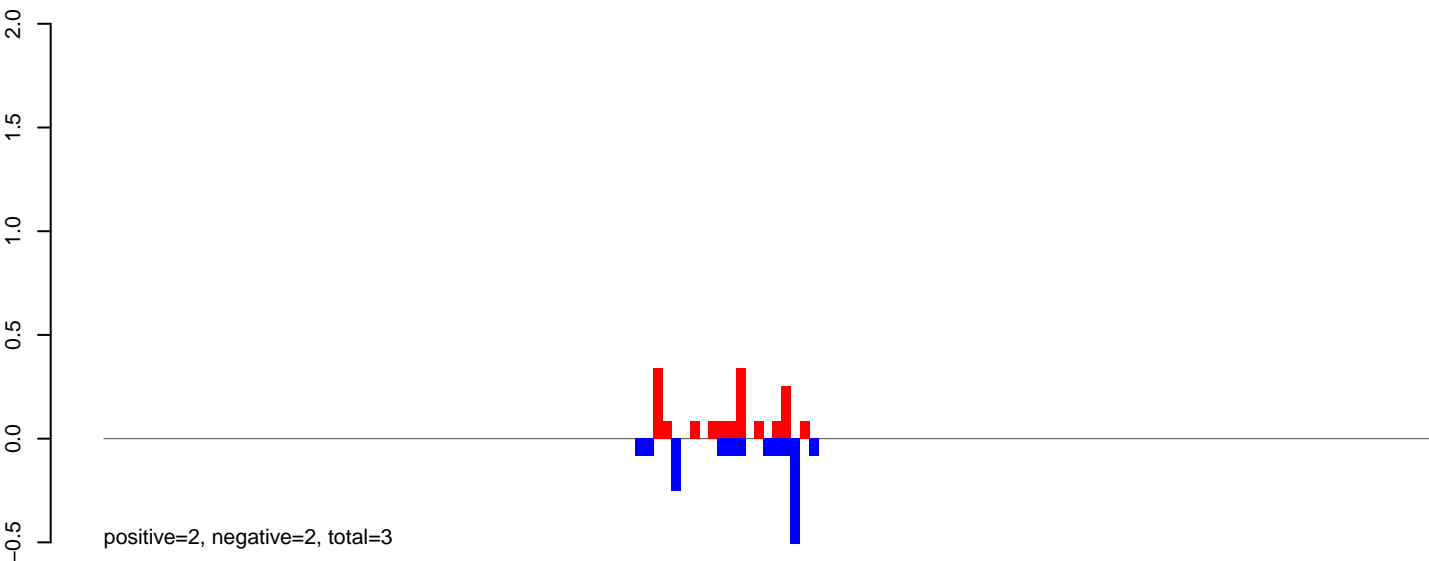


positive=0, negative=2, total=2

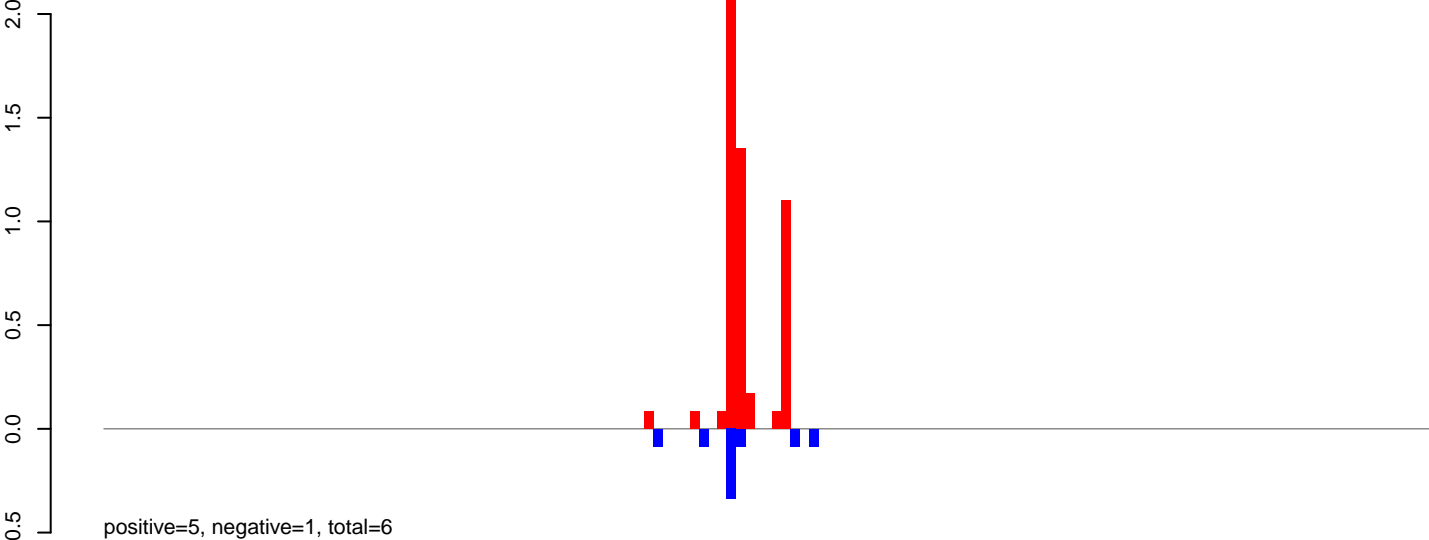
Window size=50, length=4383, TE@TF002002-Ty1_copia_Ele16:1-4383

0 1000 2000 3000 4000

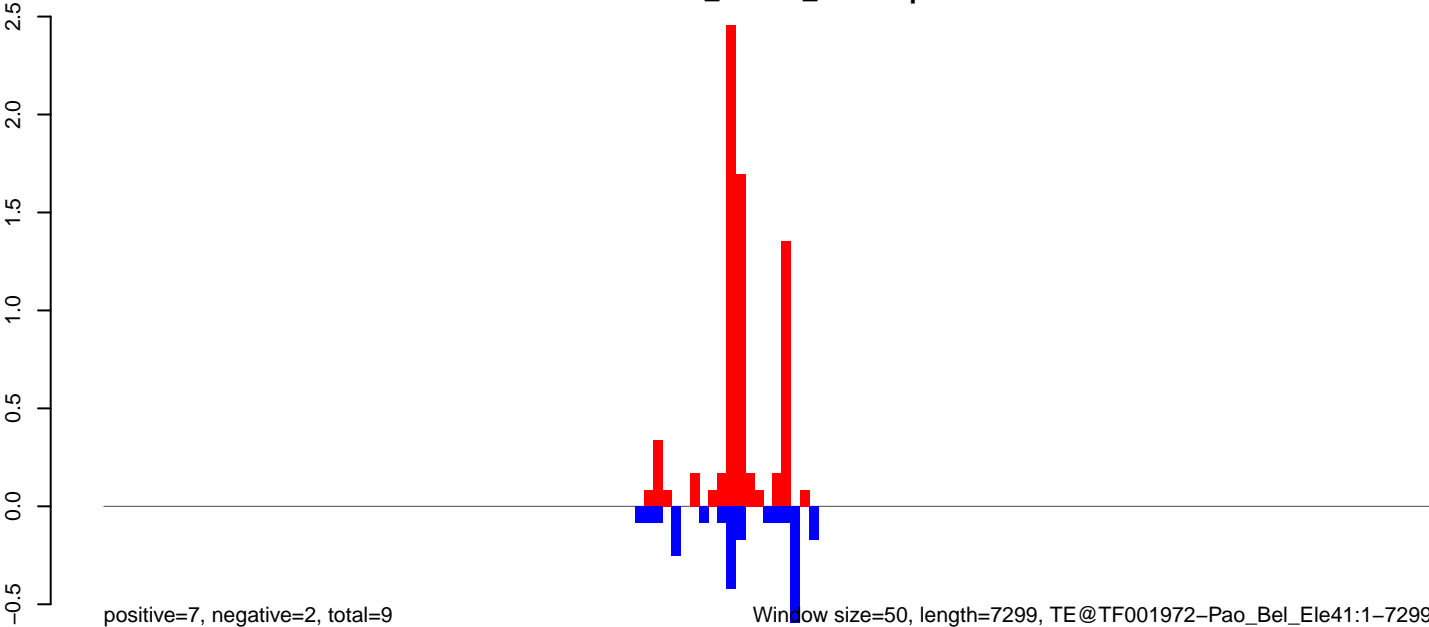
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



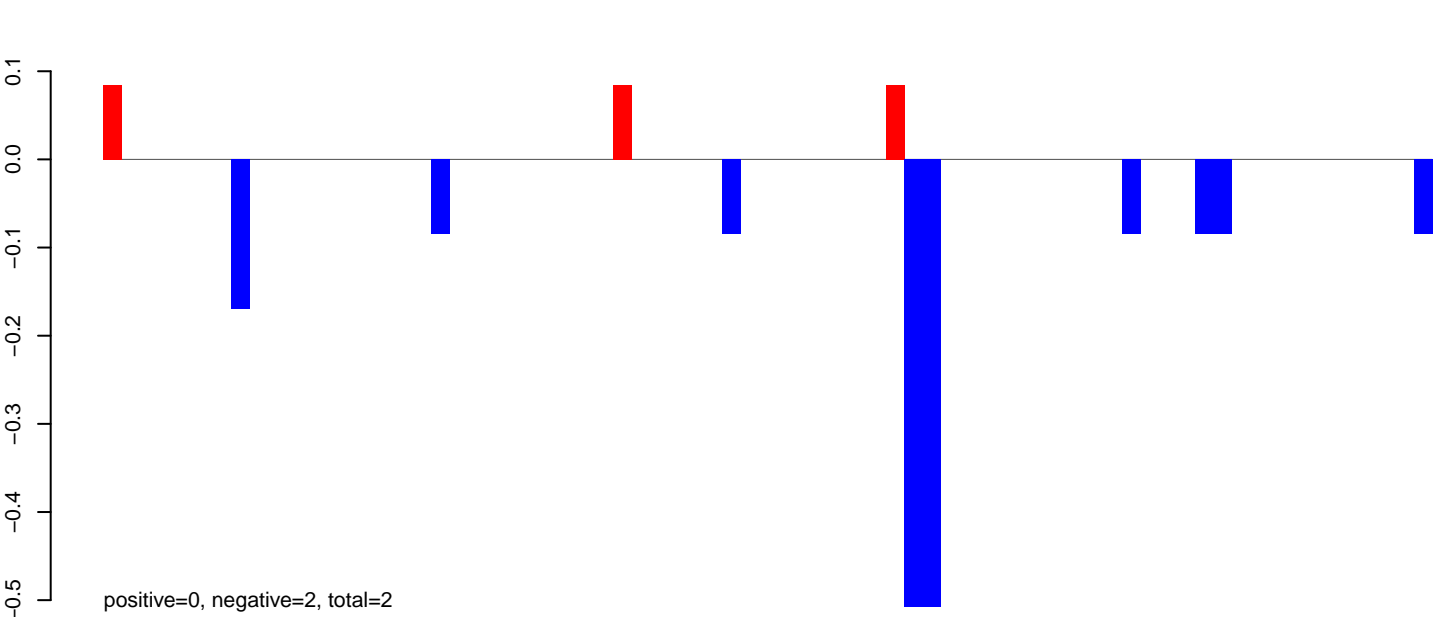
AnGam_Mos55_cells.rep



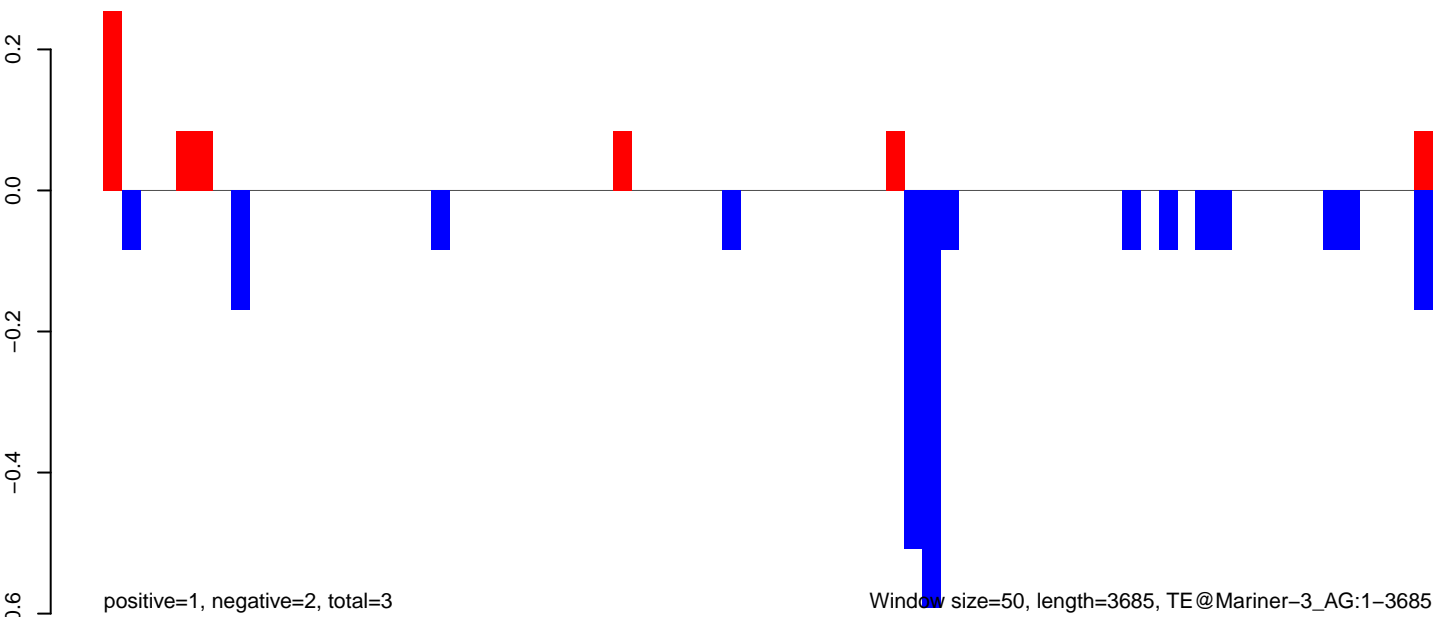
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

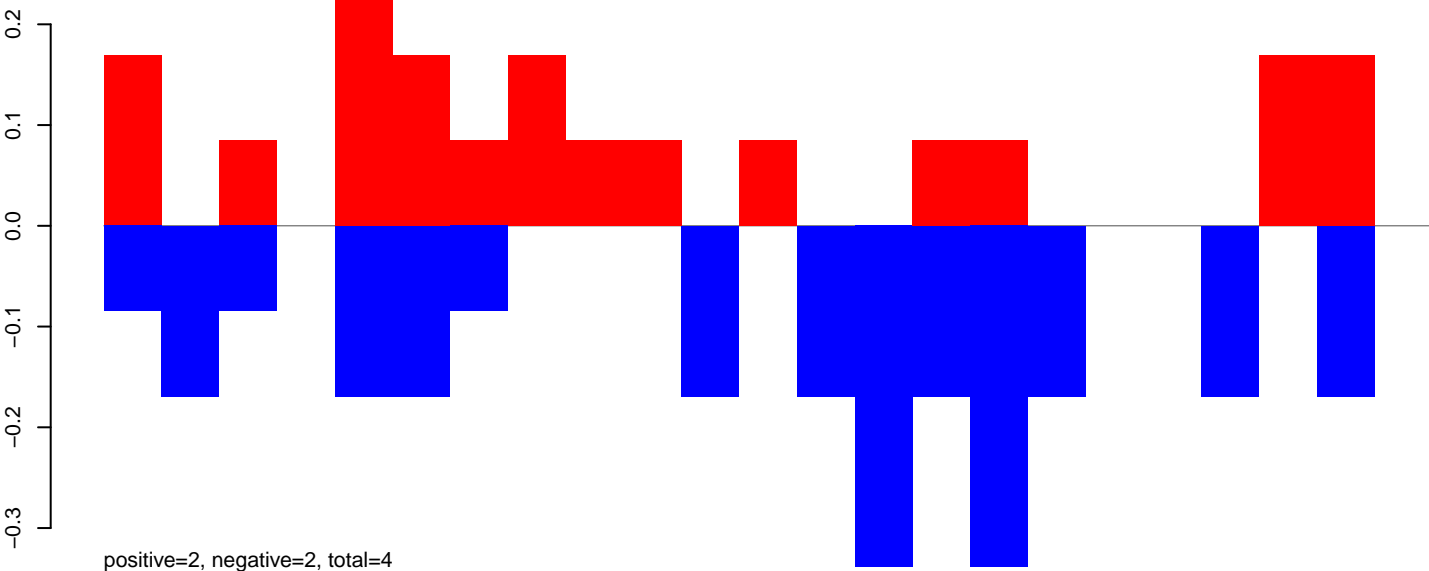


AnGam_Mos55_cells.rep



Window size=50, length=3685, TE@Mariner-3_AG:1-3685

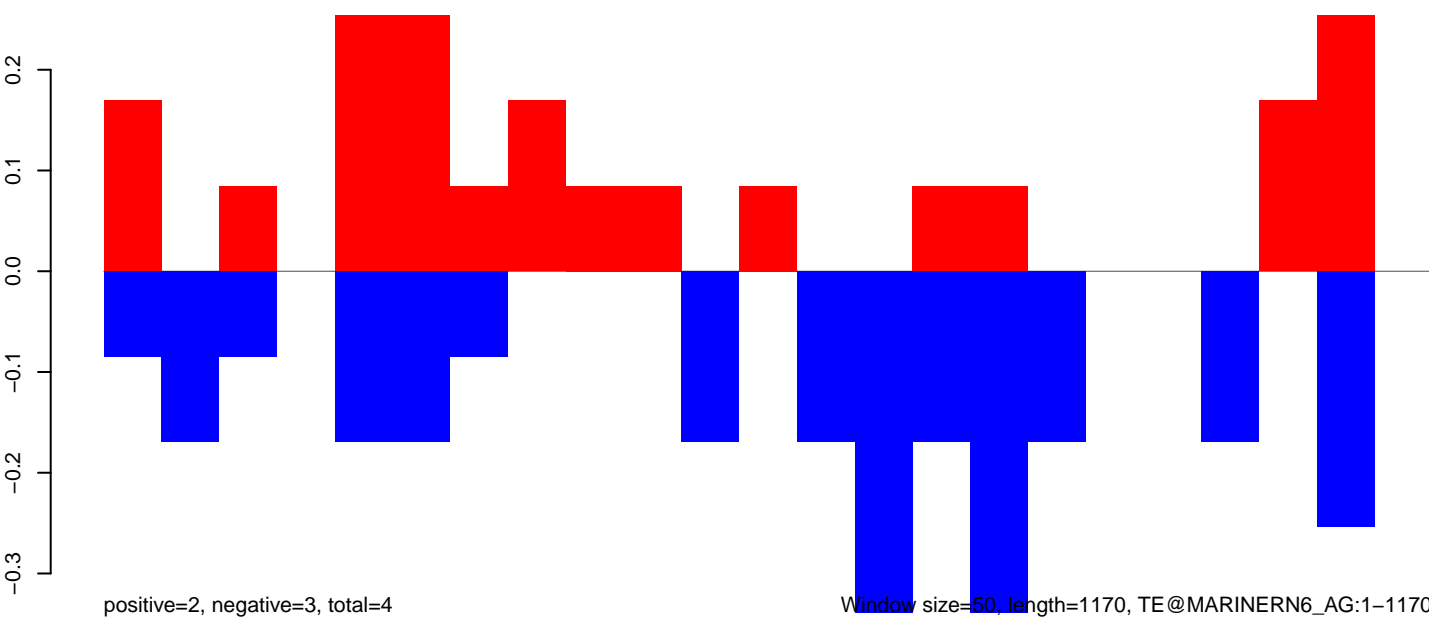
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

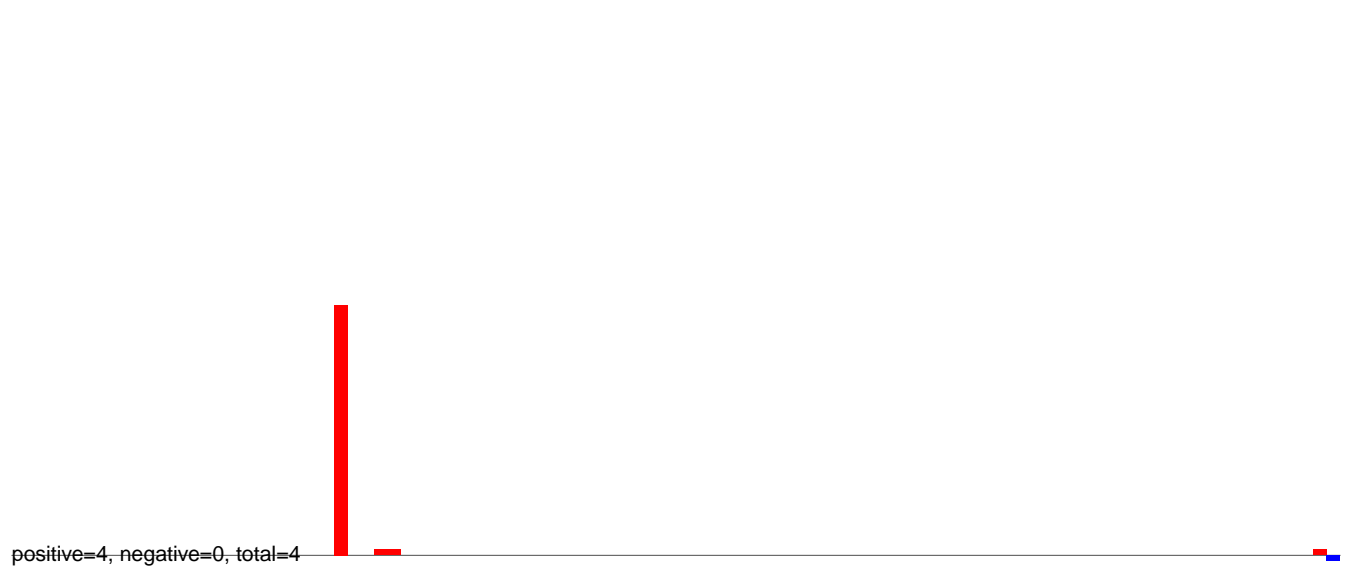


AnGam_Mos55_cells.rep

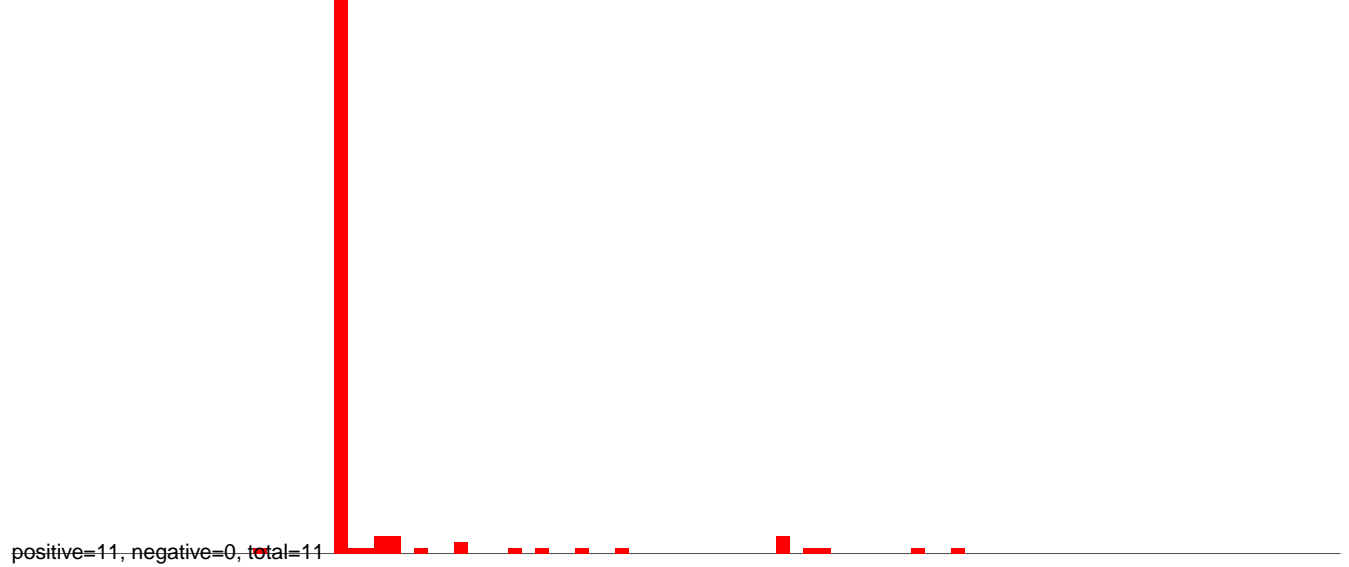


Window size=50, length=1170, TE@MARINERN6_AG:1-1170

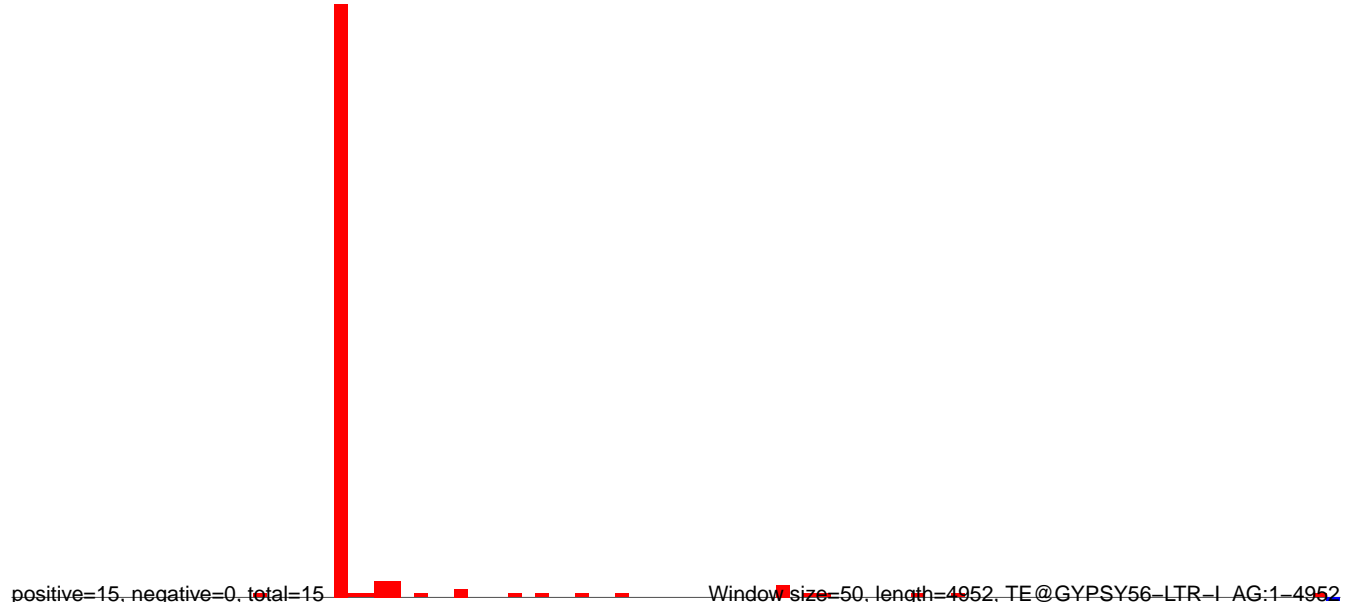
AnGam_Mos55_cells.18_23.rep



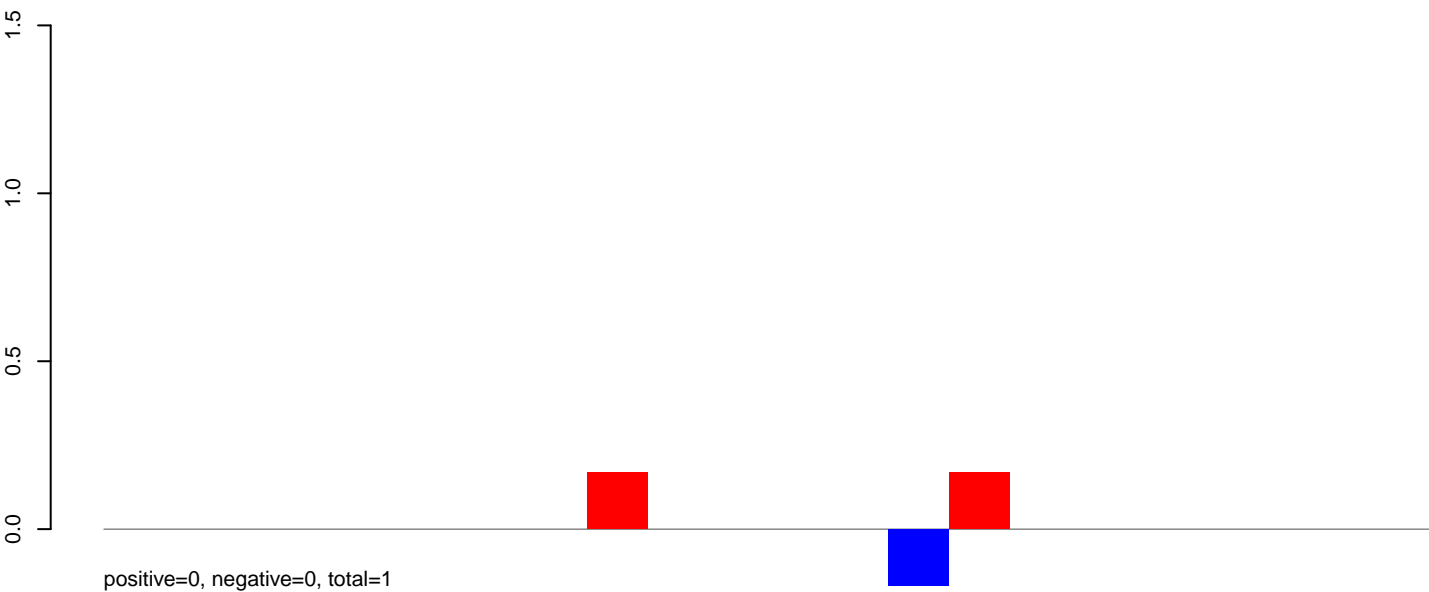
AnGam_Mos55_cells.24_35.rep



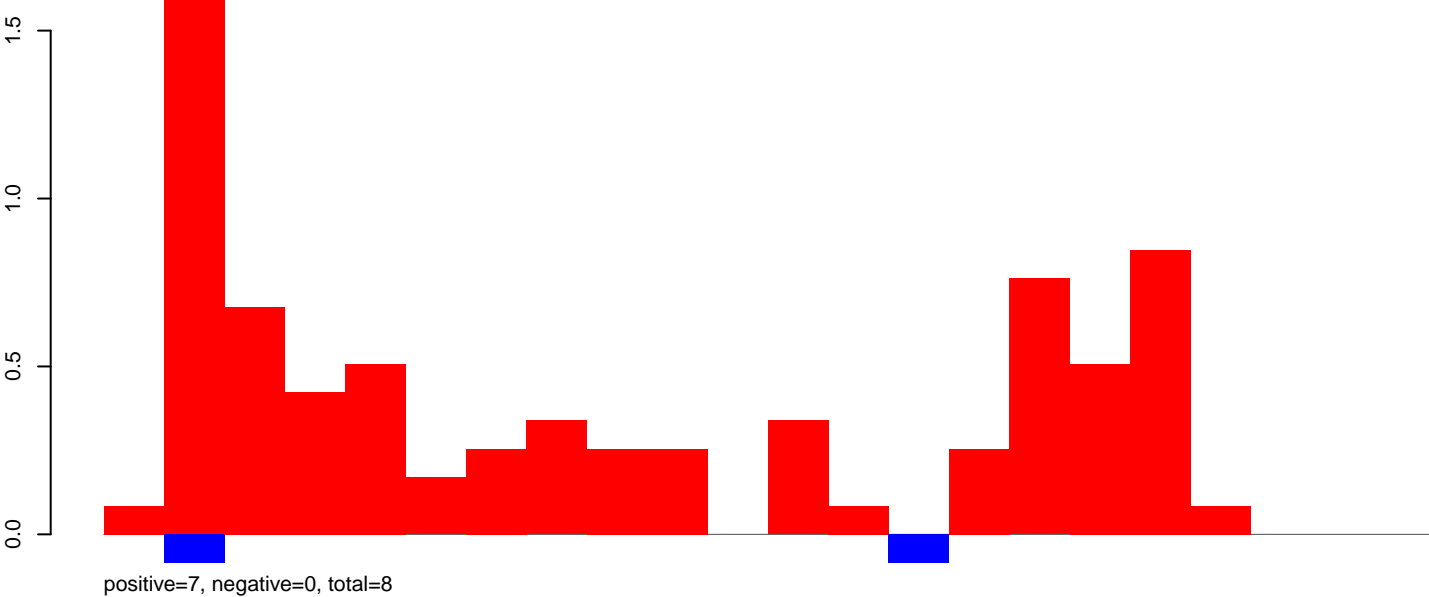
AnGam_Mos55_cells.rep



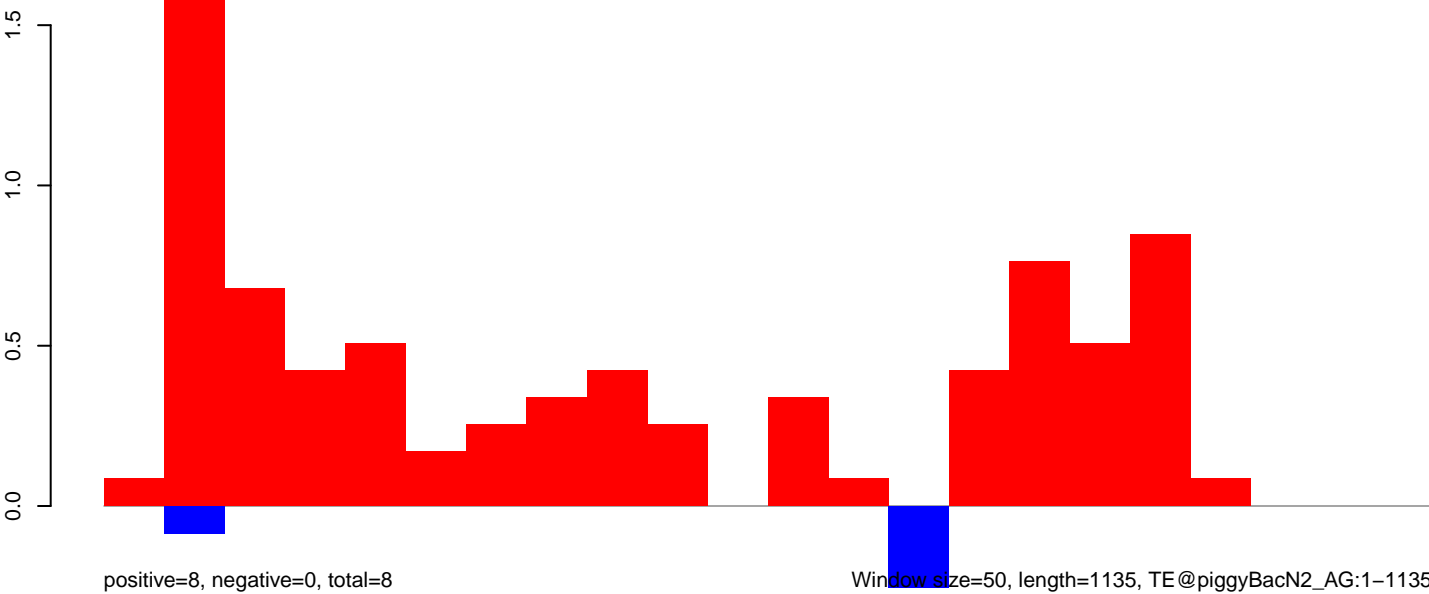
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

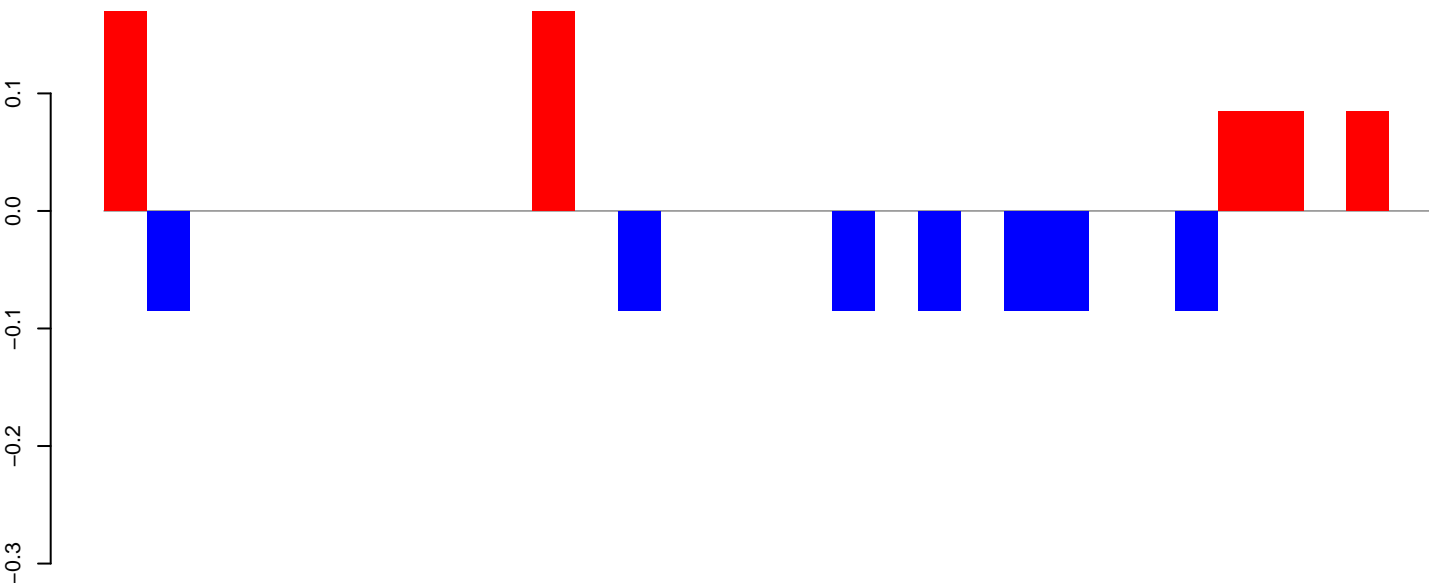


AnGam_Mos55_cells.rep



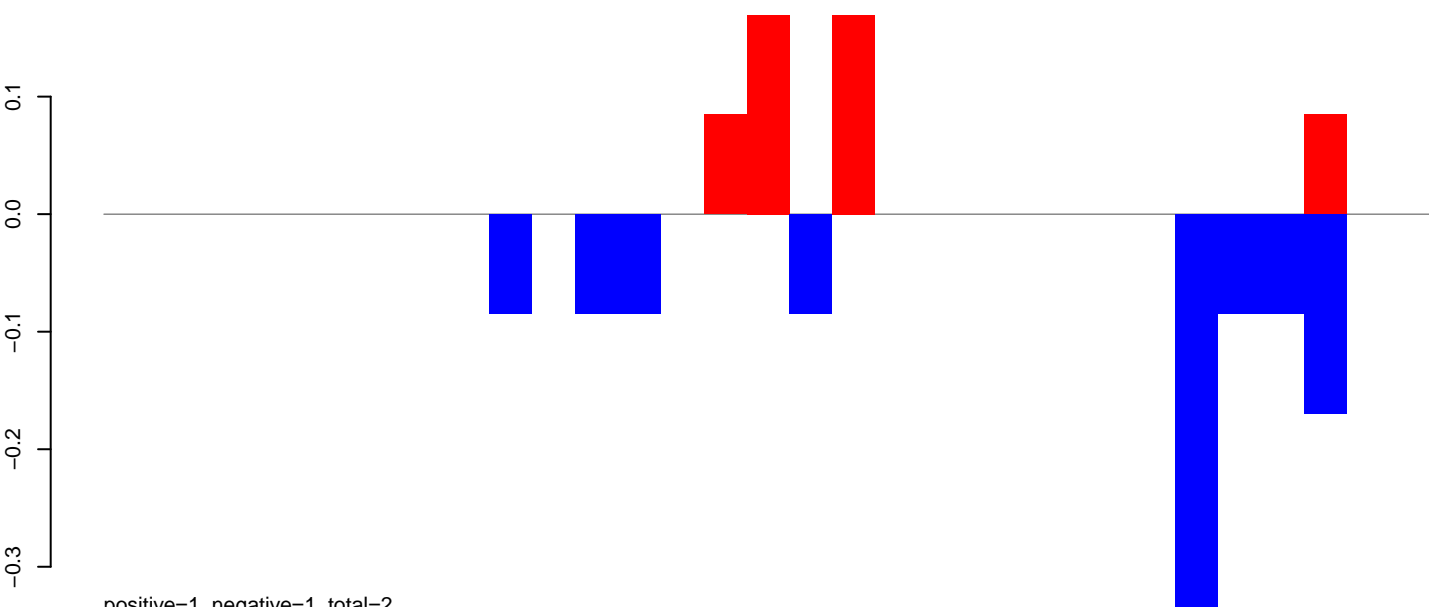
Window size=50, length=1135, TE@piggyBacN2_AG:1-1135

AnGam_Mos55_cells.18_23.rep



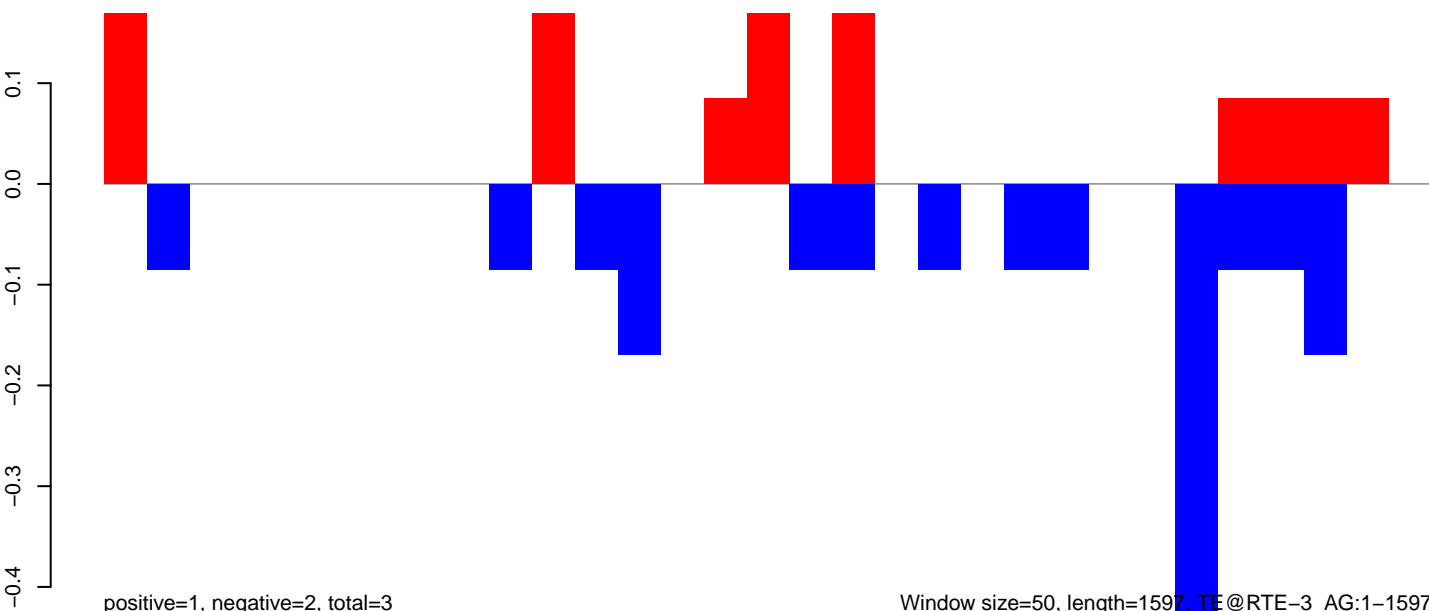
positive=1, negative=1, total=1

AnGam_Mos55_cells.24_35.rep



positive=1, negative=1, total=2

AnGam_Mos55_cells.rep

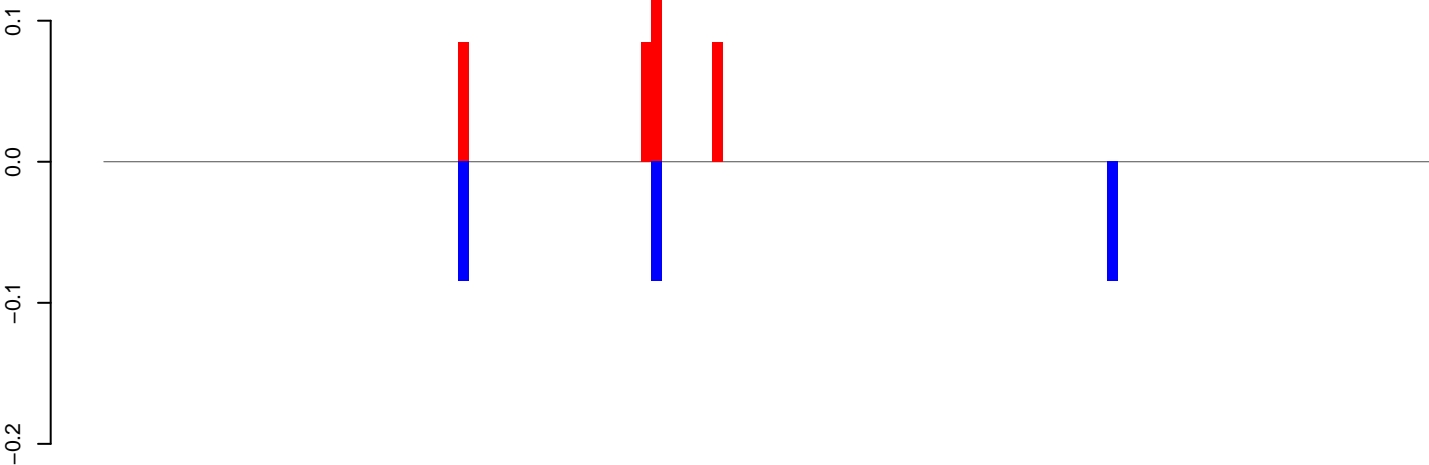


positive=1, negative=2, total=3

Window size=50, length=1597, TE@RTE-3_AG:1-1597

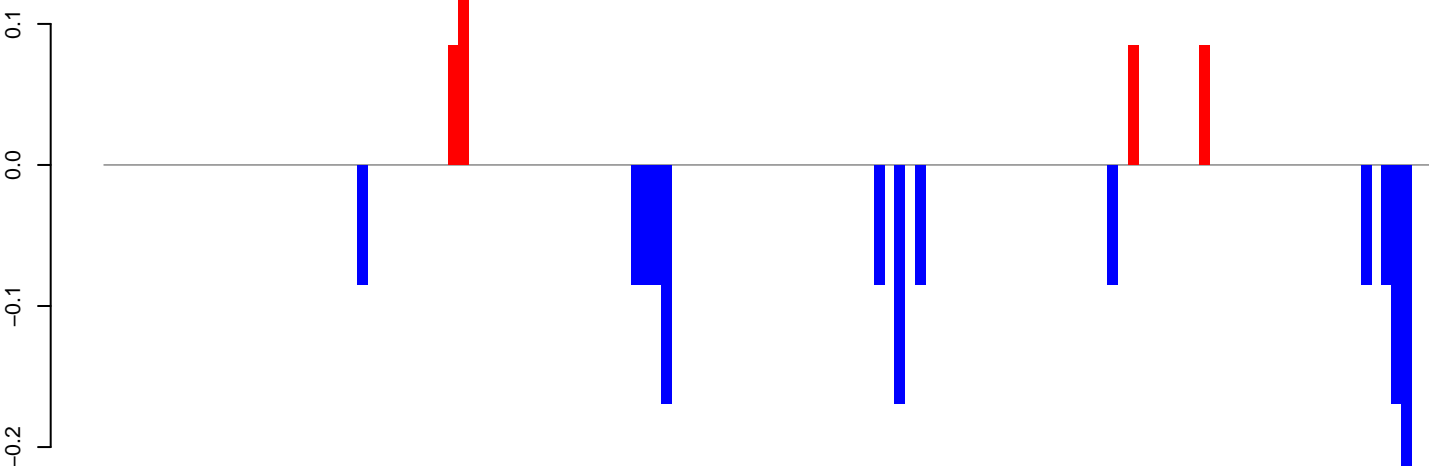
0 500 1000 1500

AnGam_Mos55_cells.18_23.rep



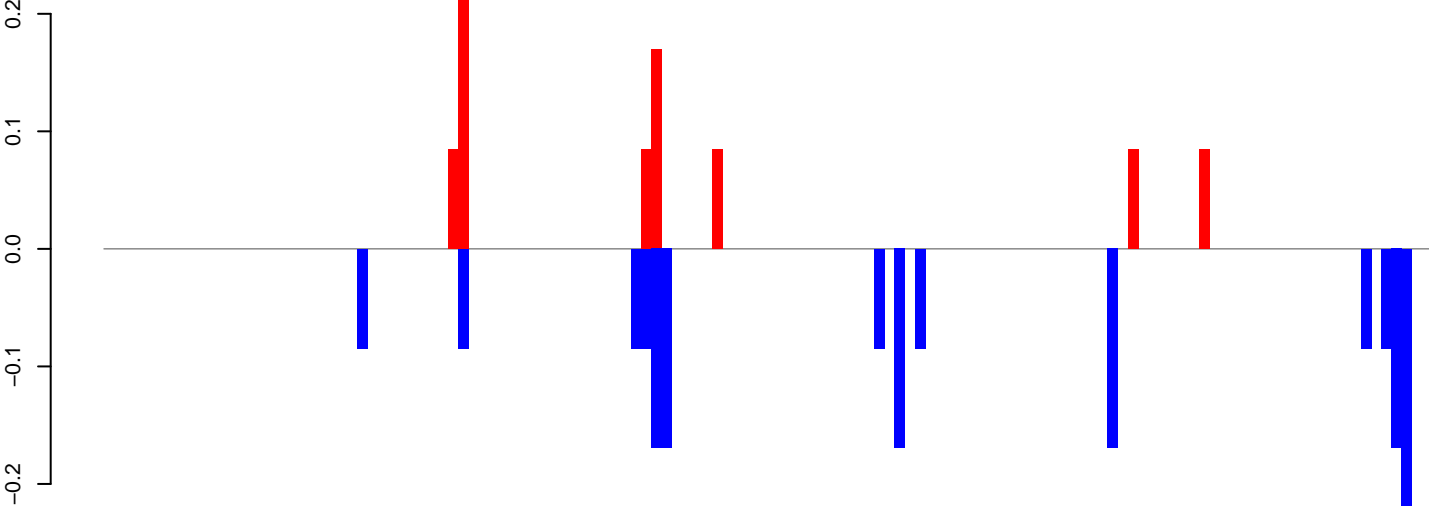
positive=0, negative=0, total=1

AnGam_Mos55_cells.24_35.rep



positive=0, negative=2, total=2

AnGam_Mos55_cells.rep

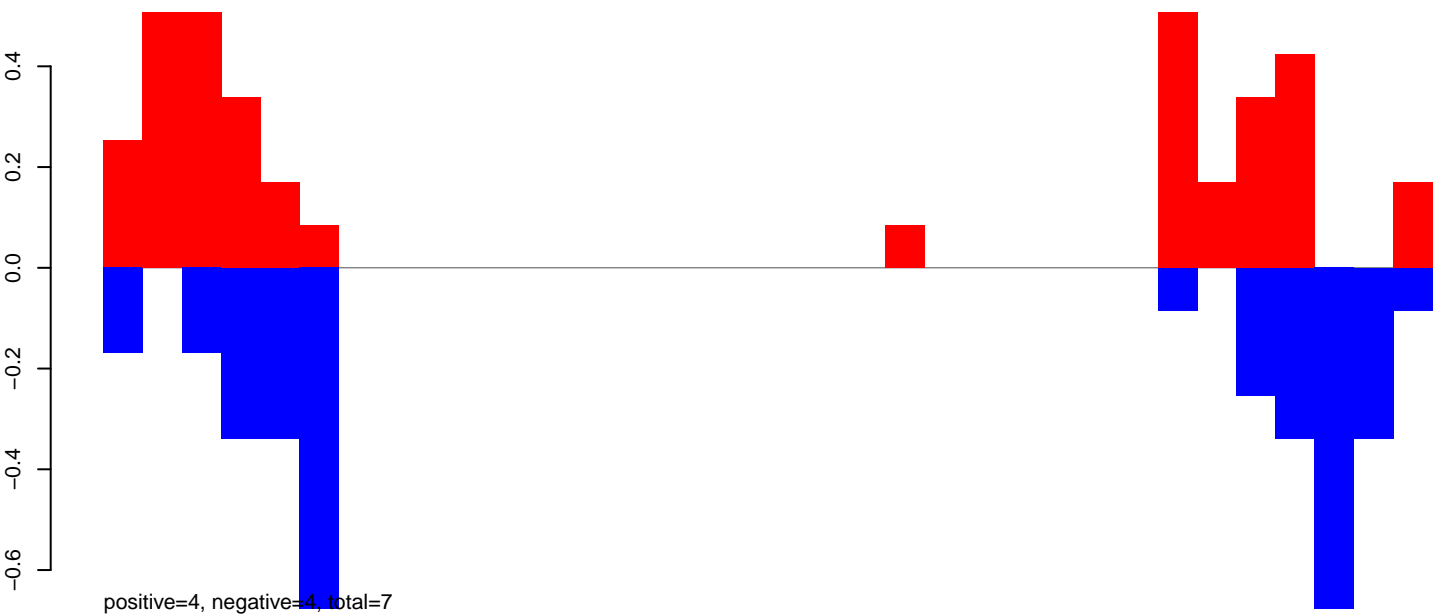


positive=1, negative=2, total=3

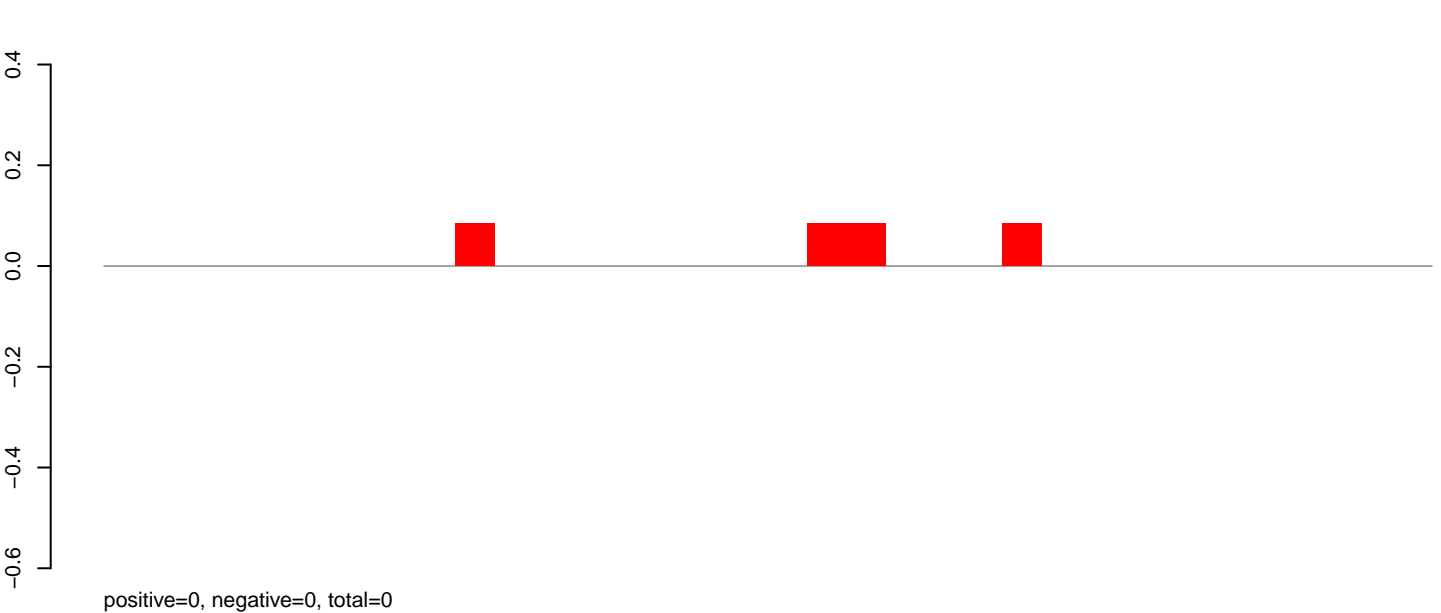
Window size=50, length=6590, TE@R7Ag2:1-6590

0 1000 2000 3000 4000 5000 6000

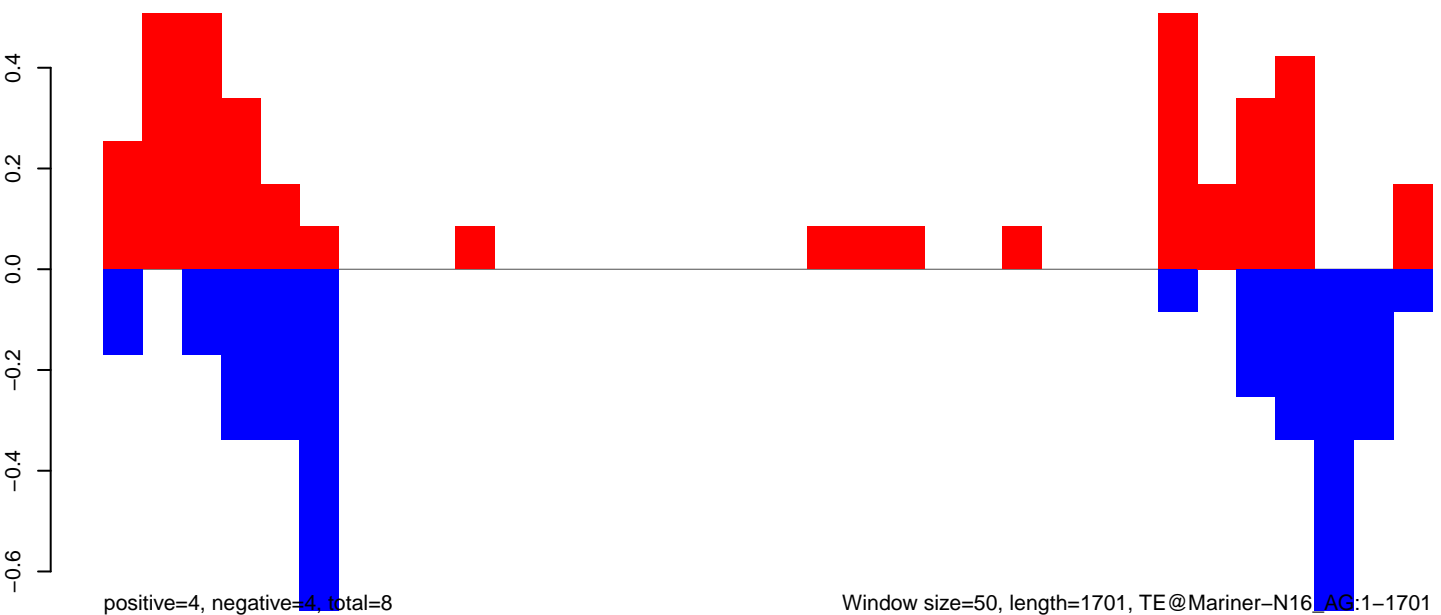
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



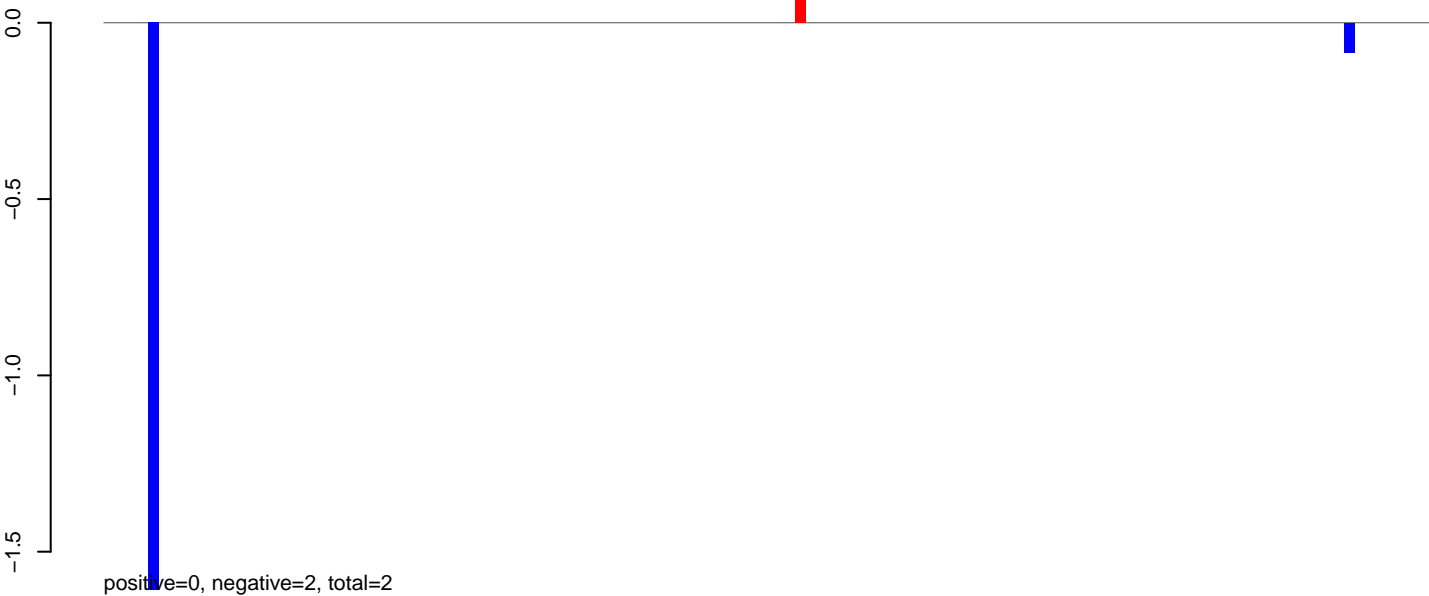
AnGam_Mos55_cells.rep



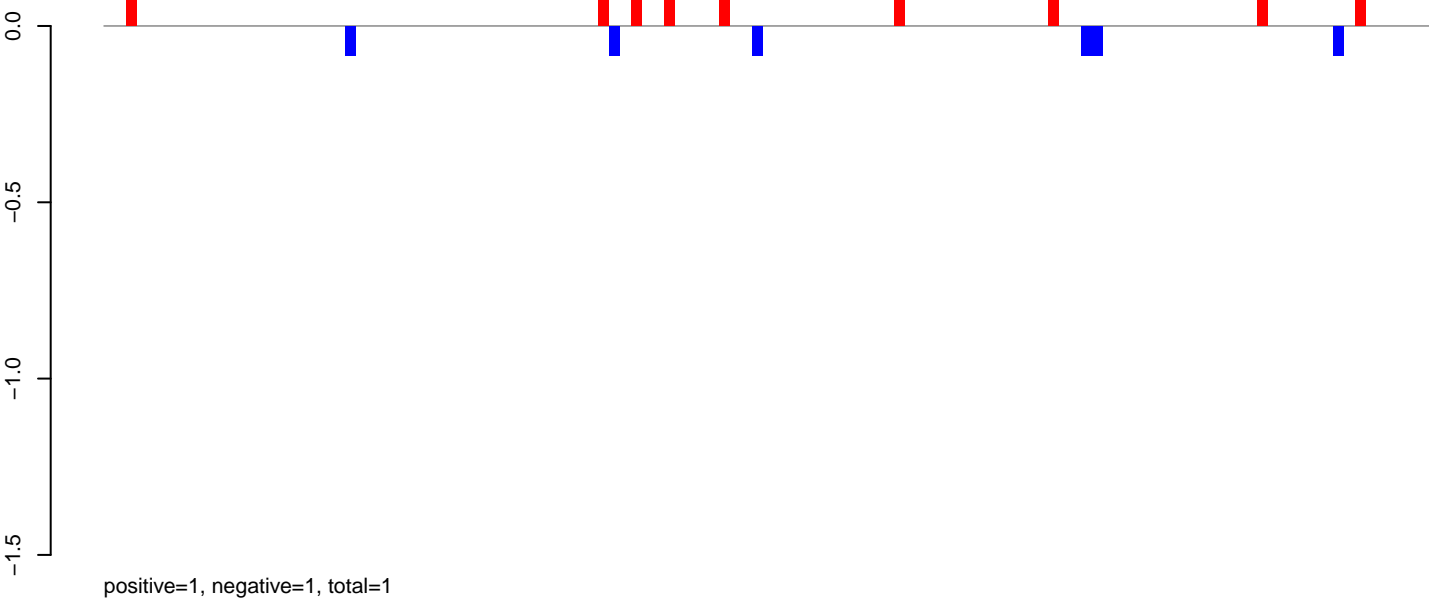
Window size=50, length=1701, TE@Mariner-N16, AG:1-1701

0 500 1000 1500

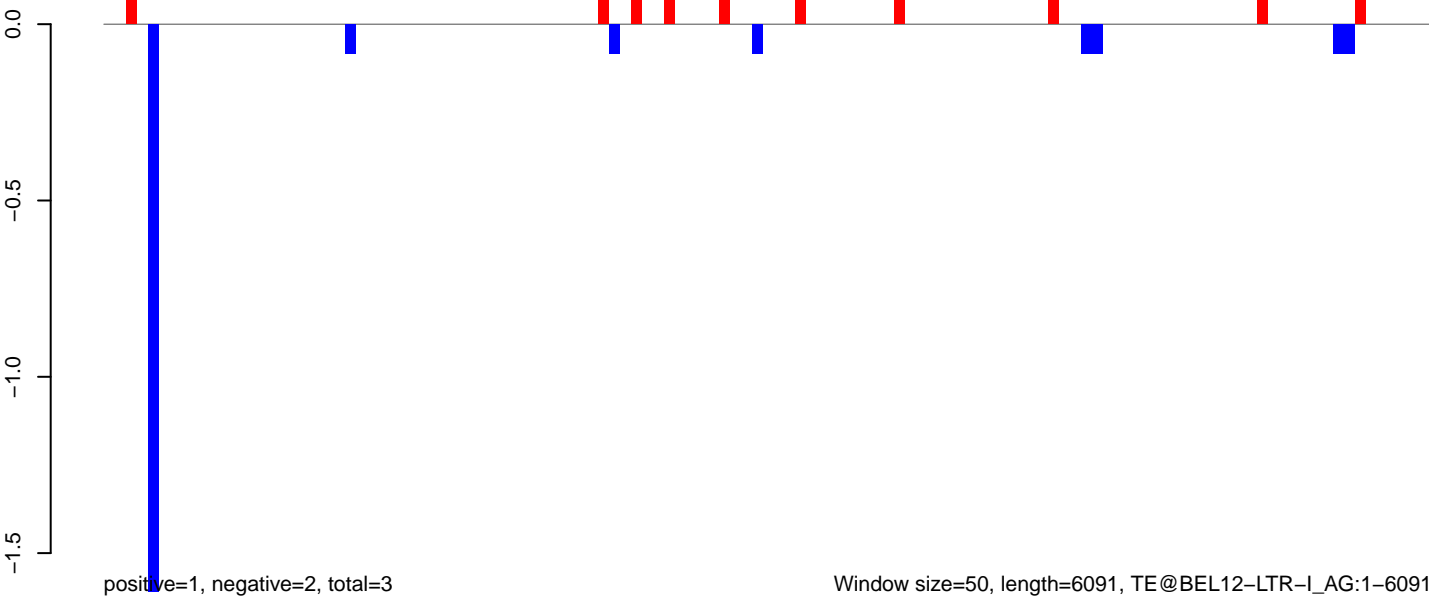
AnGam_Mos55_cells.18_23.rep



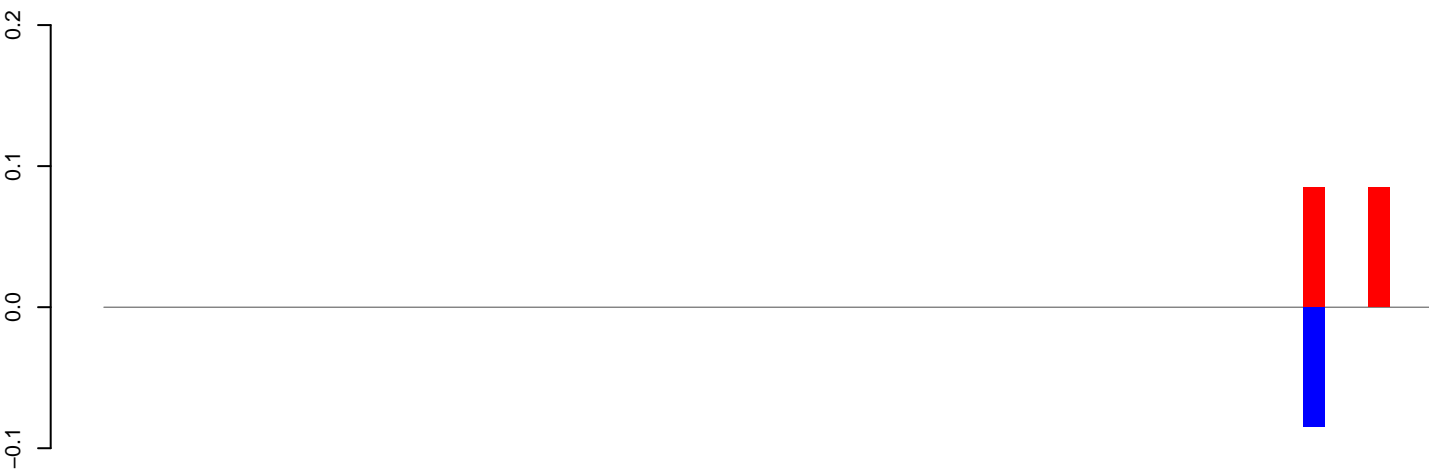
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

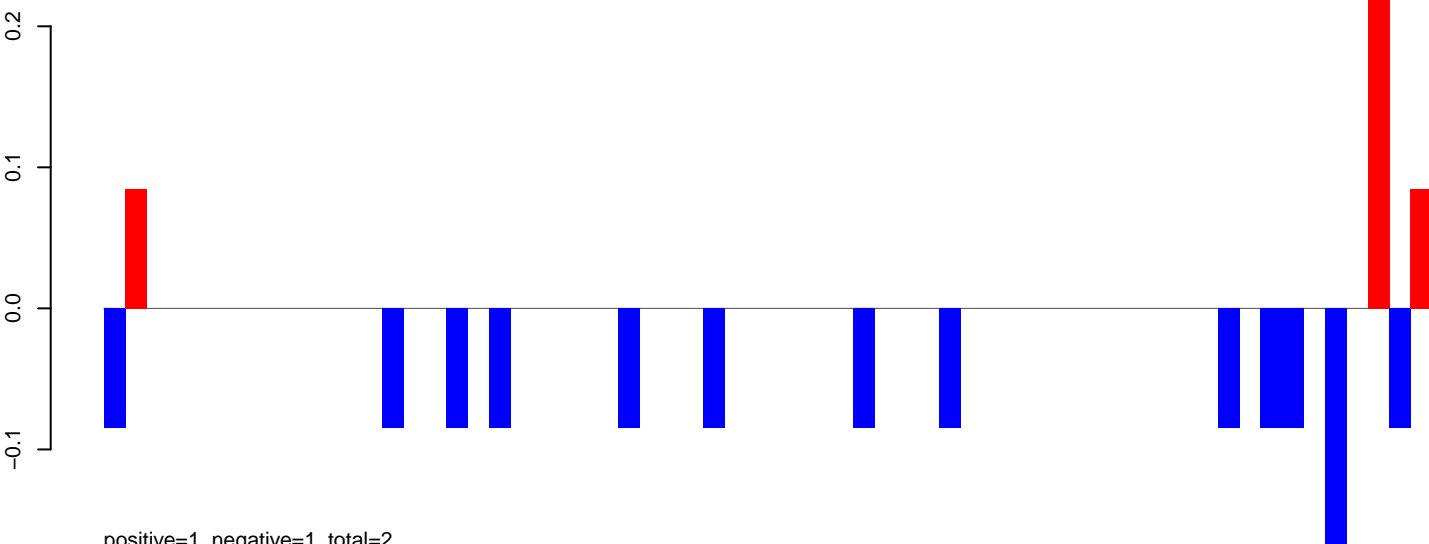


AnGam_Mos55_cells.18_23.rep



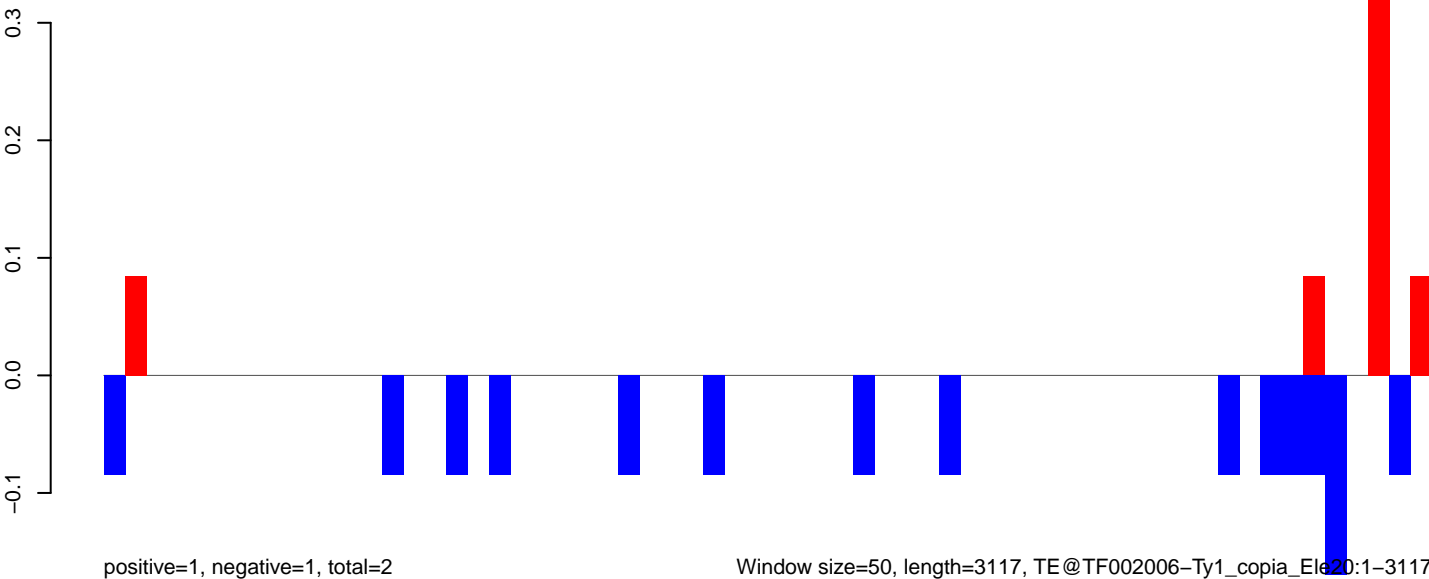
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=1, negative=1, total=2

AnGam_Mos55_cells.rep

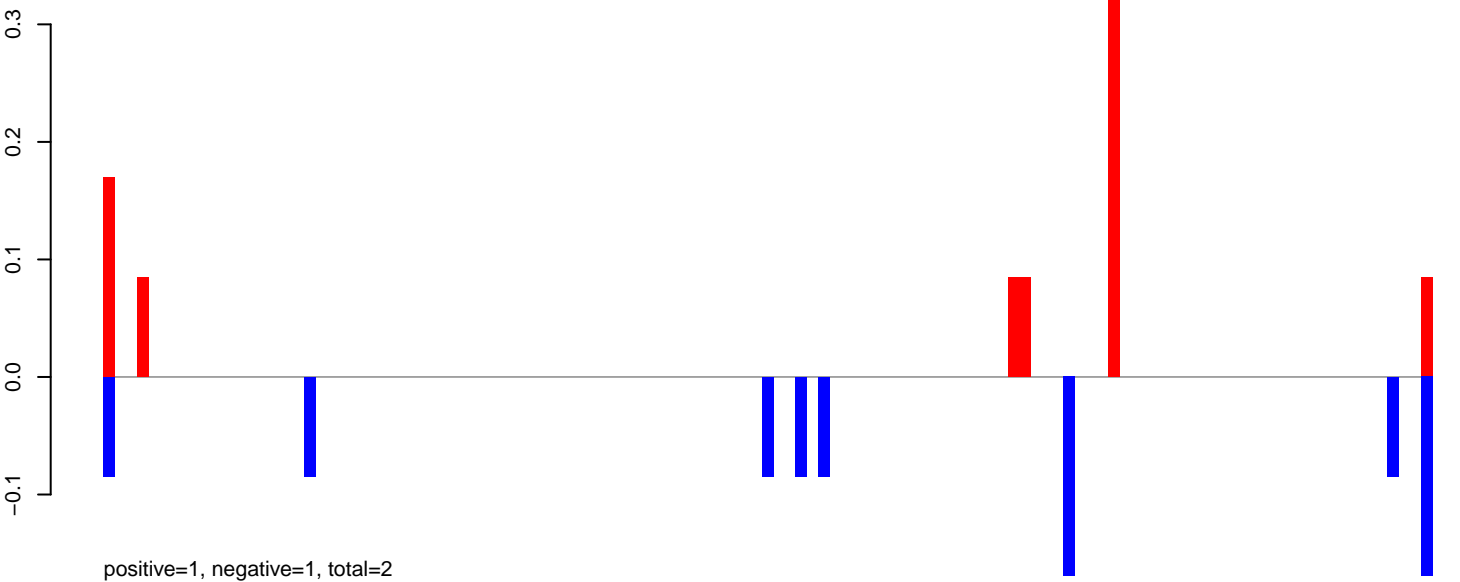


positive=1, negative=1, total=2

Window size=50, length=3117, TE@TF002006-Ty1_copia_Ele20:1-3117

0 500 1000 1500 2000 2500 3000

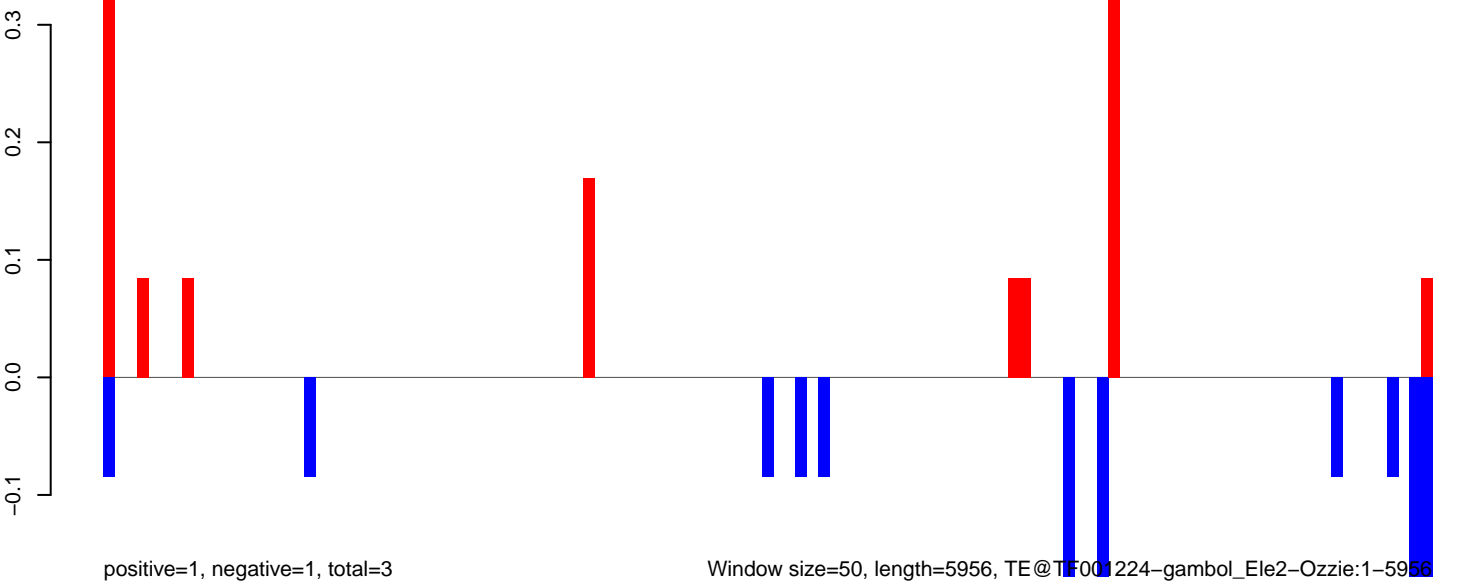
AnGam_Mos55_cells.18_23.rep



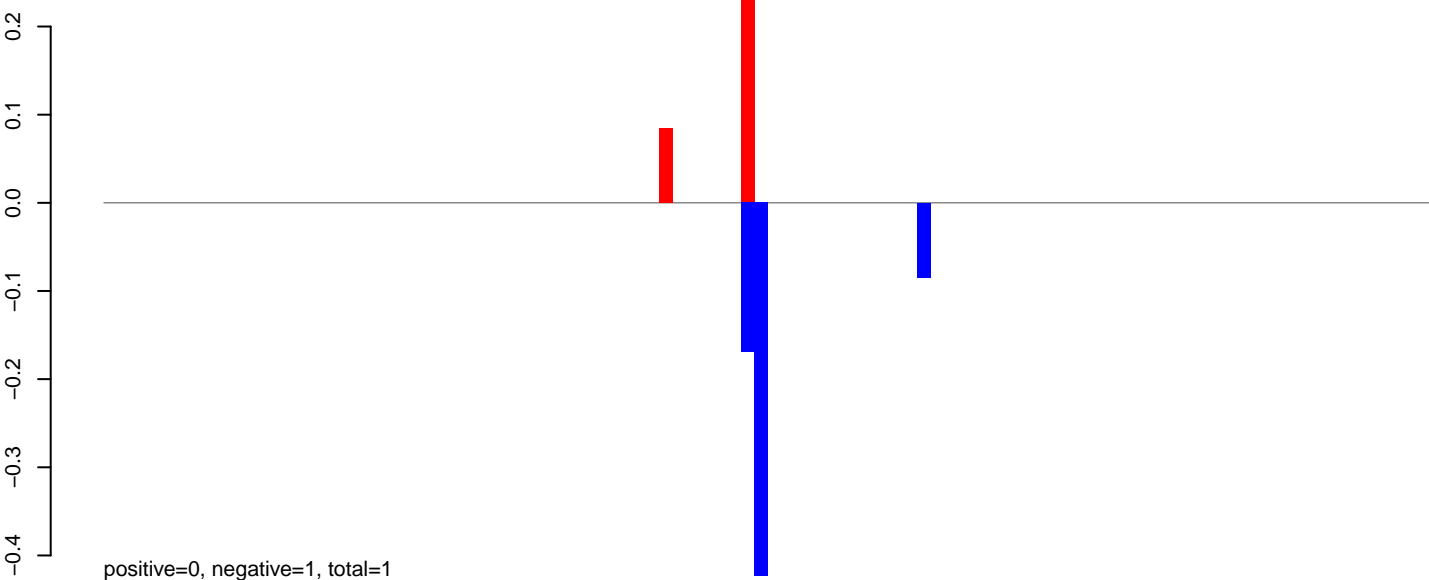
AnGam_Mos55_cells.24_35.rep



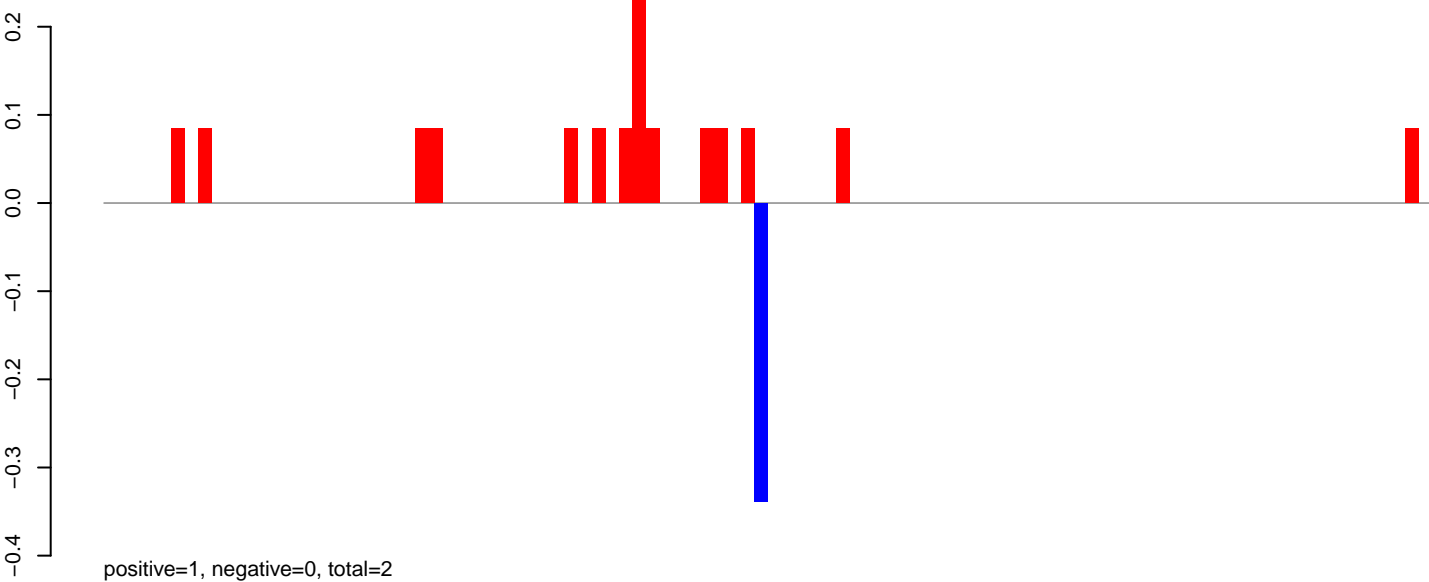
AnGam_Mos55_cells.rep



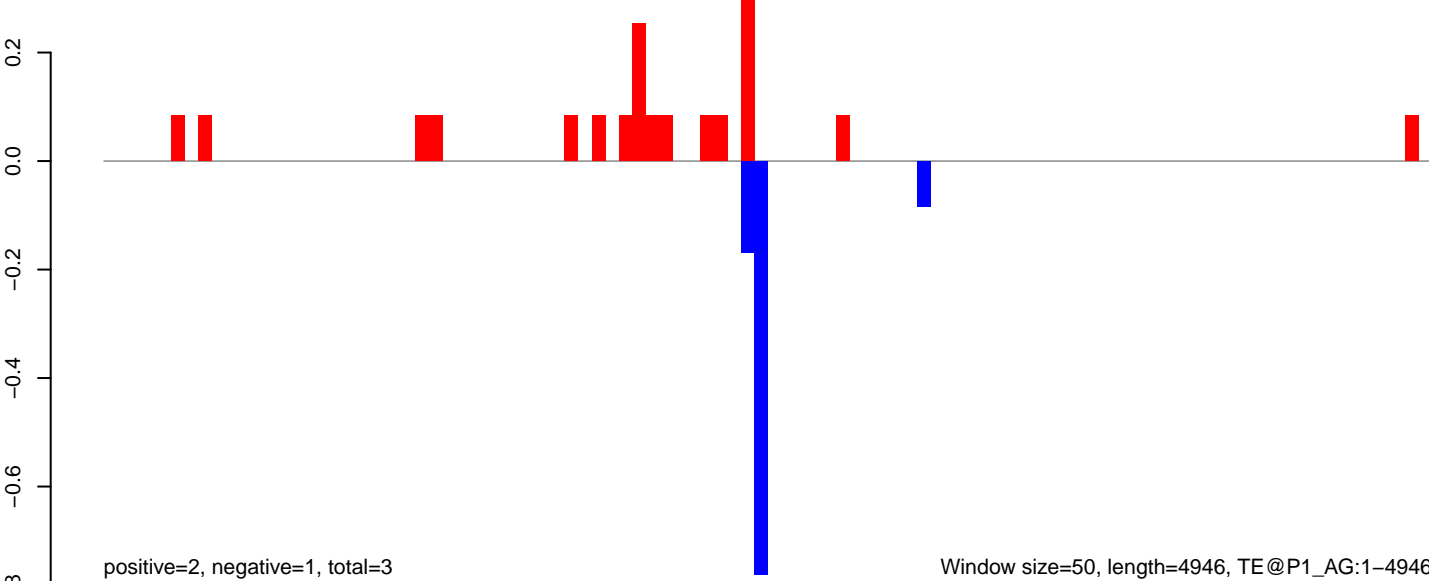
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

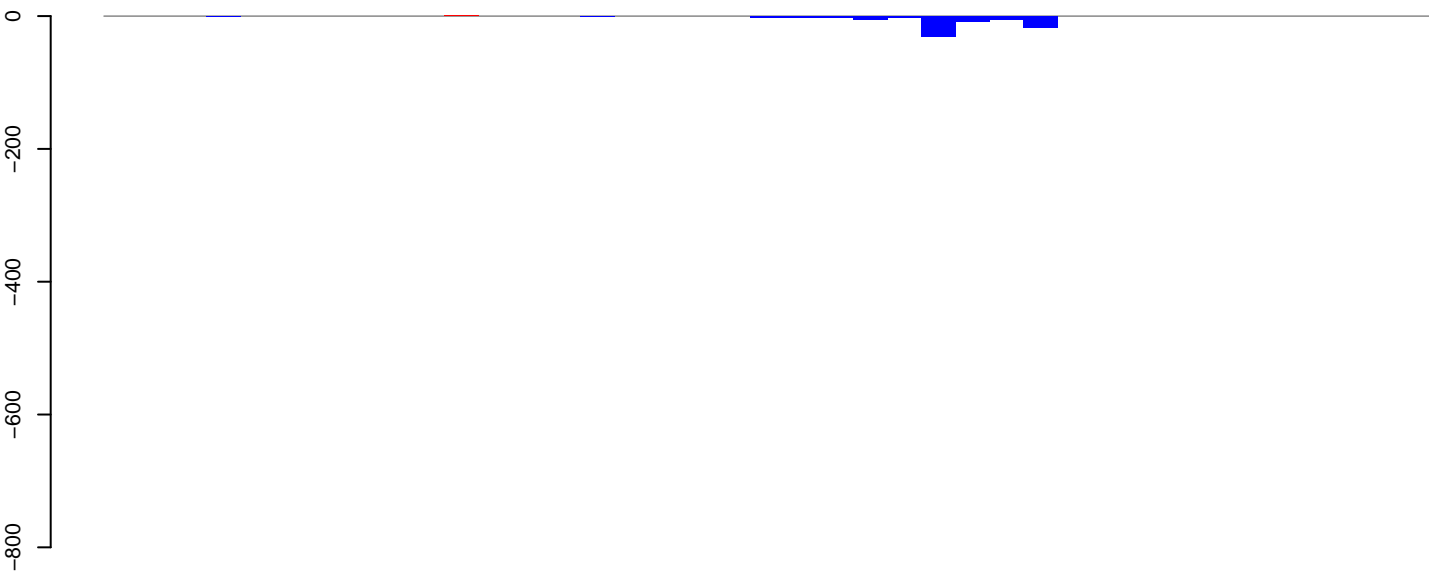


AnGam_Mos55_cells.rep



Window size=50, length=4946, TE@P1_AG:1-4946

AnGam_Mos55_cells.18_23.rep



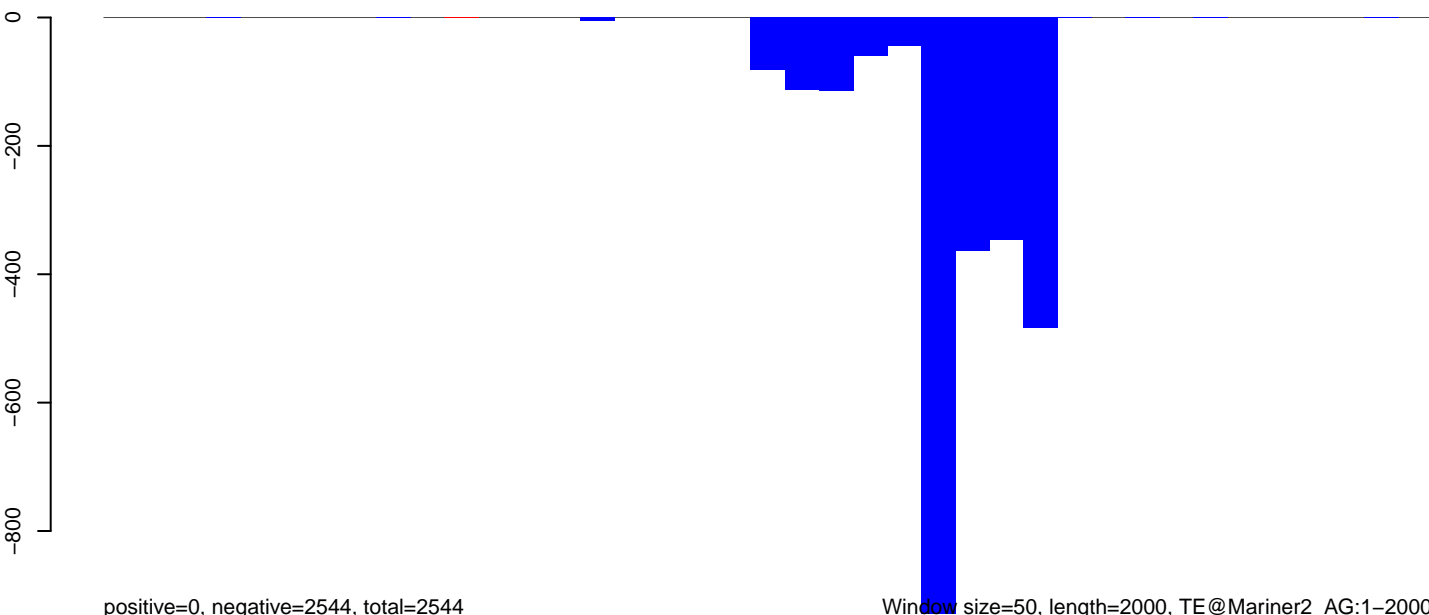
positive=0, negative=77, total=77

AnGam_Mos55_cells.24_35.rep



positive=0, negative=2467, total=2467

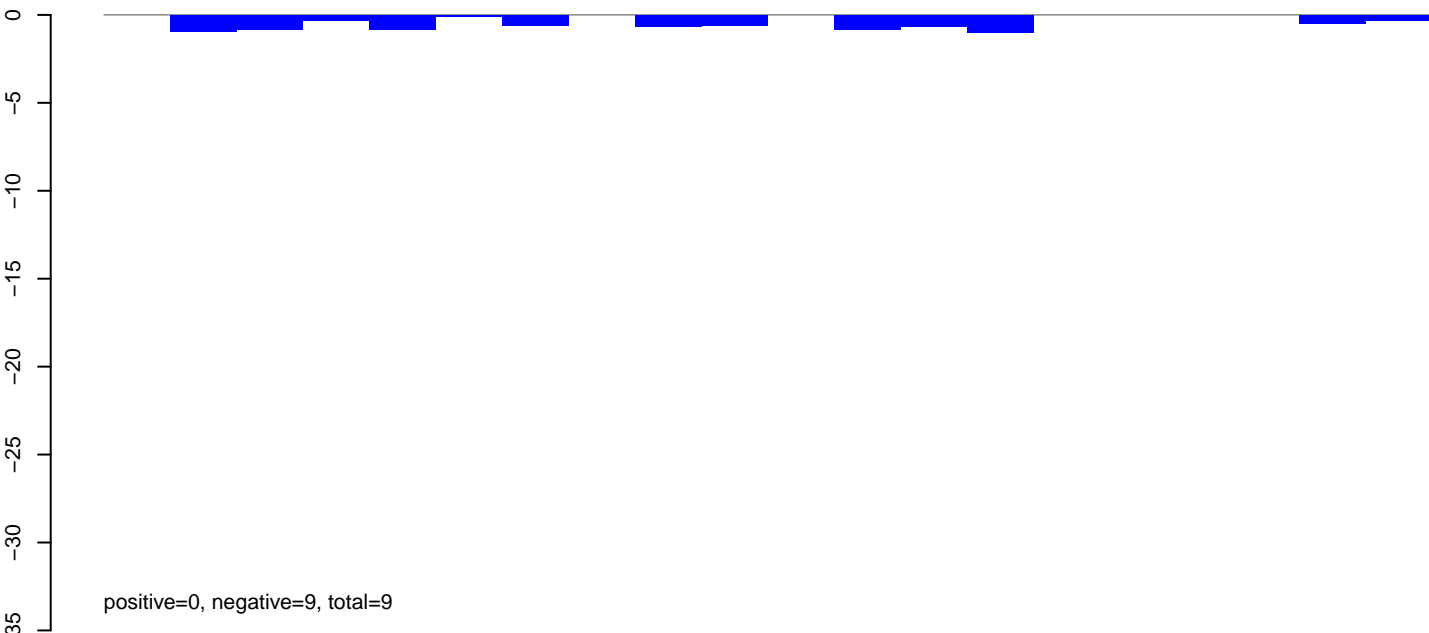
AnGam_Mos55_cells.rep



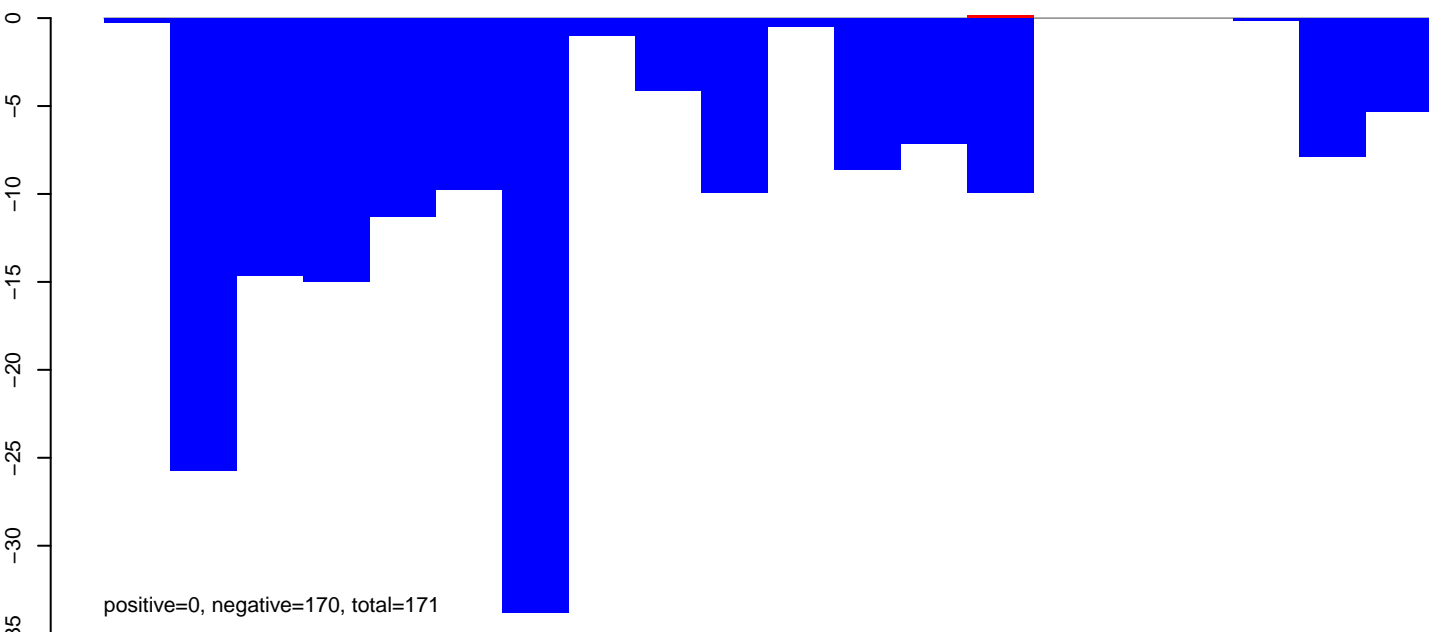
positive=0, negative=2544, total=2544

Window size=50, length=2000, TE@Mariner2_AG:1-2000

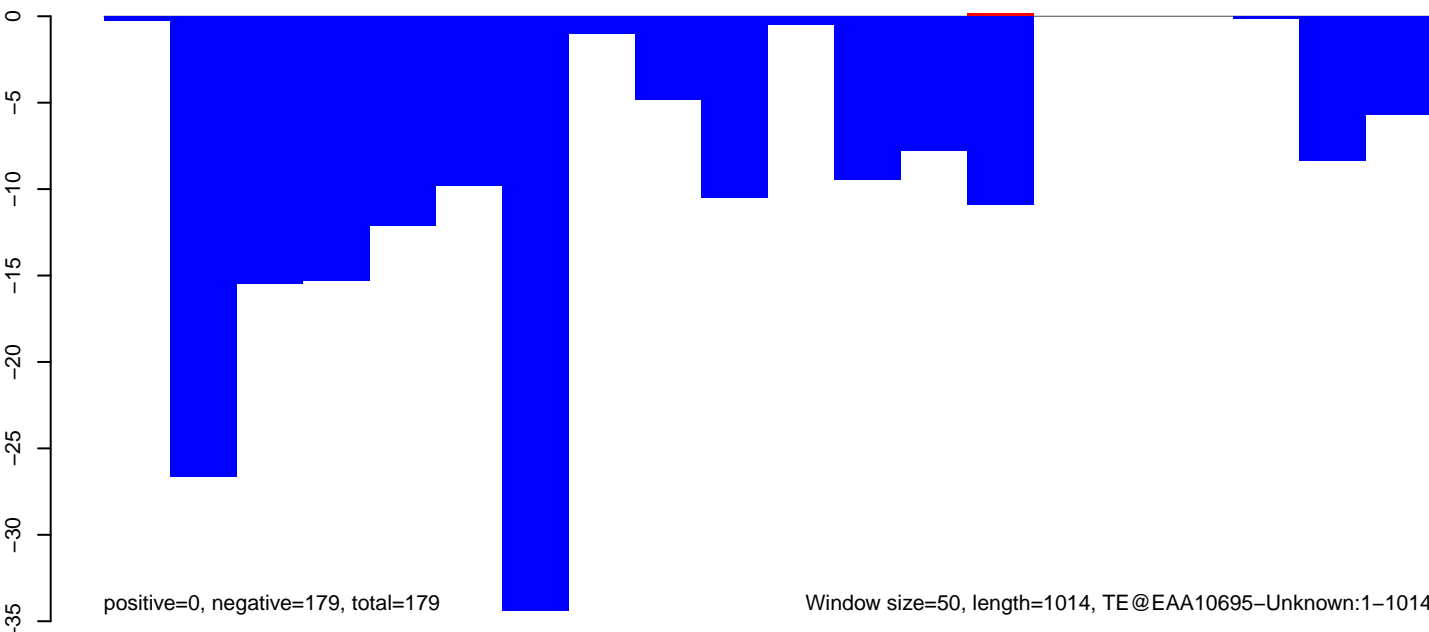
AnGam_Mos55_cells.18_23.rep



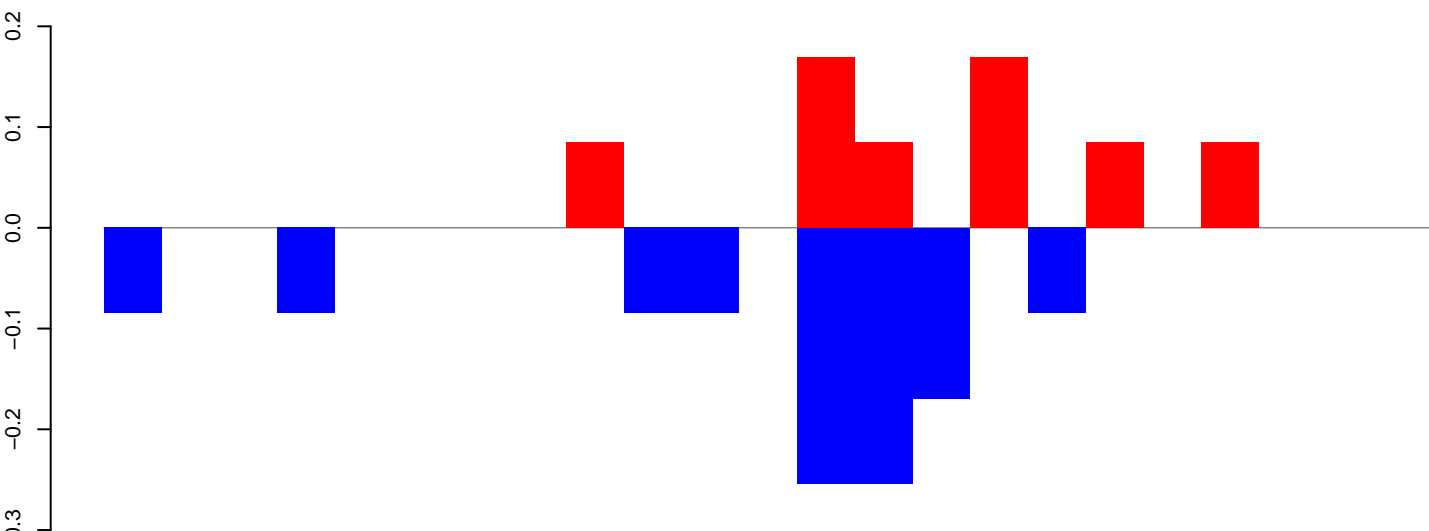
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

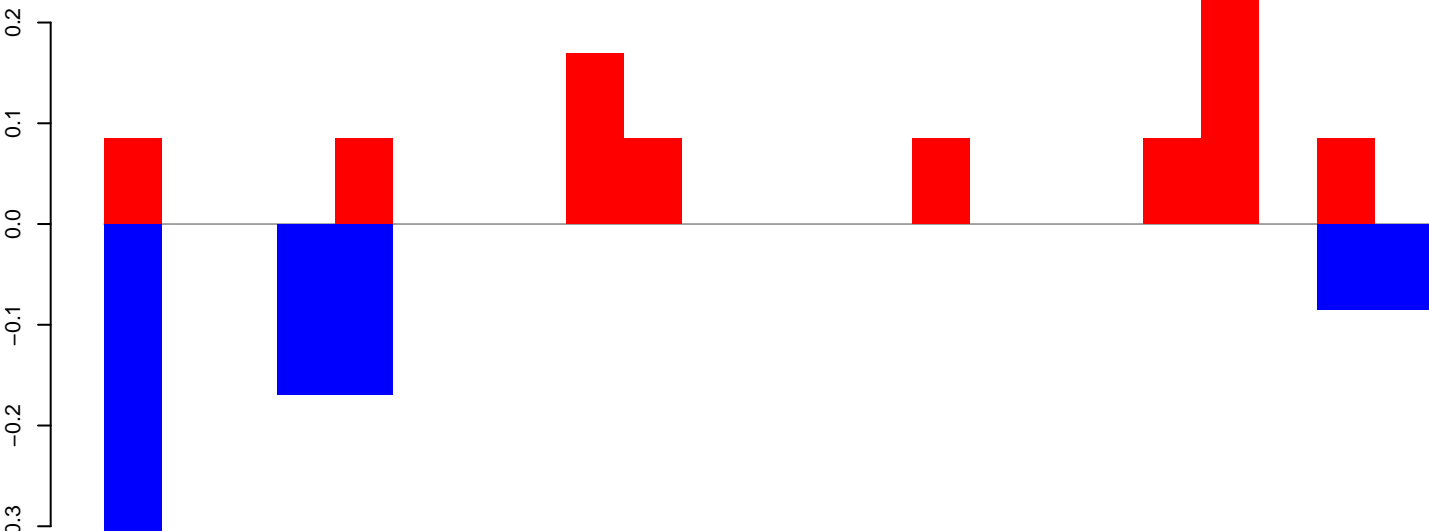


AnGam_Mos55_cells.18_23.rep



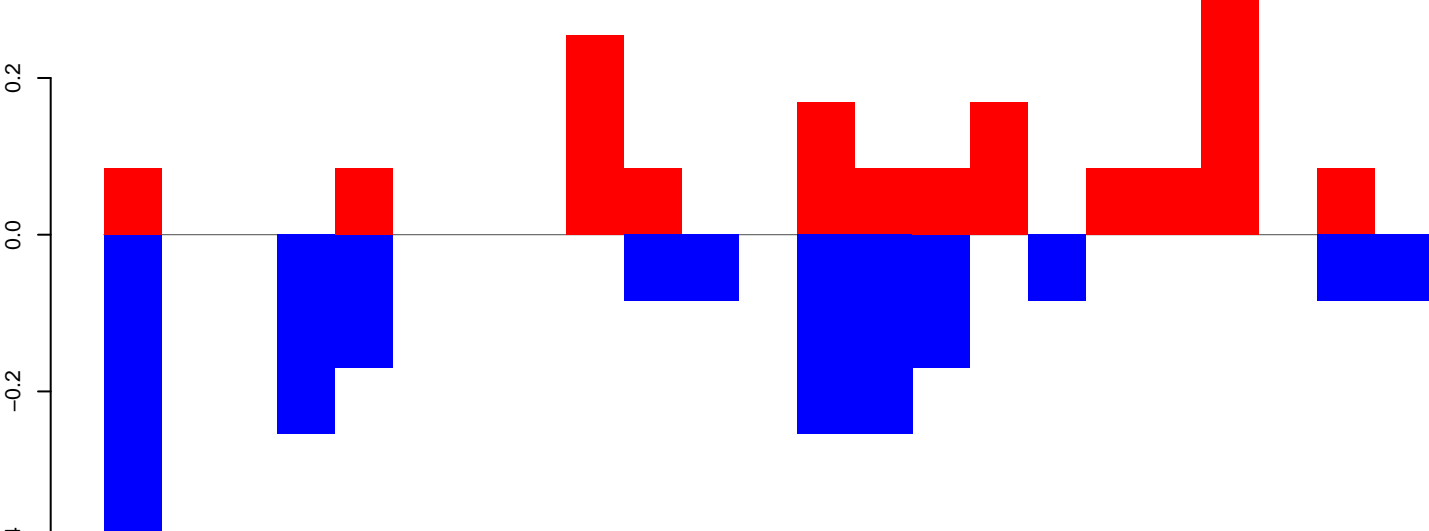
positive=1, negative=1, total=2

AnGam_Mos55_cells.24_35.rep



positive=1, negative=1, total=2

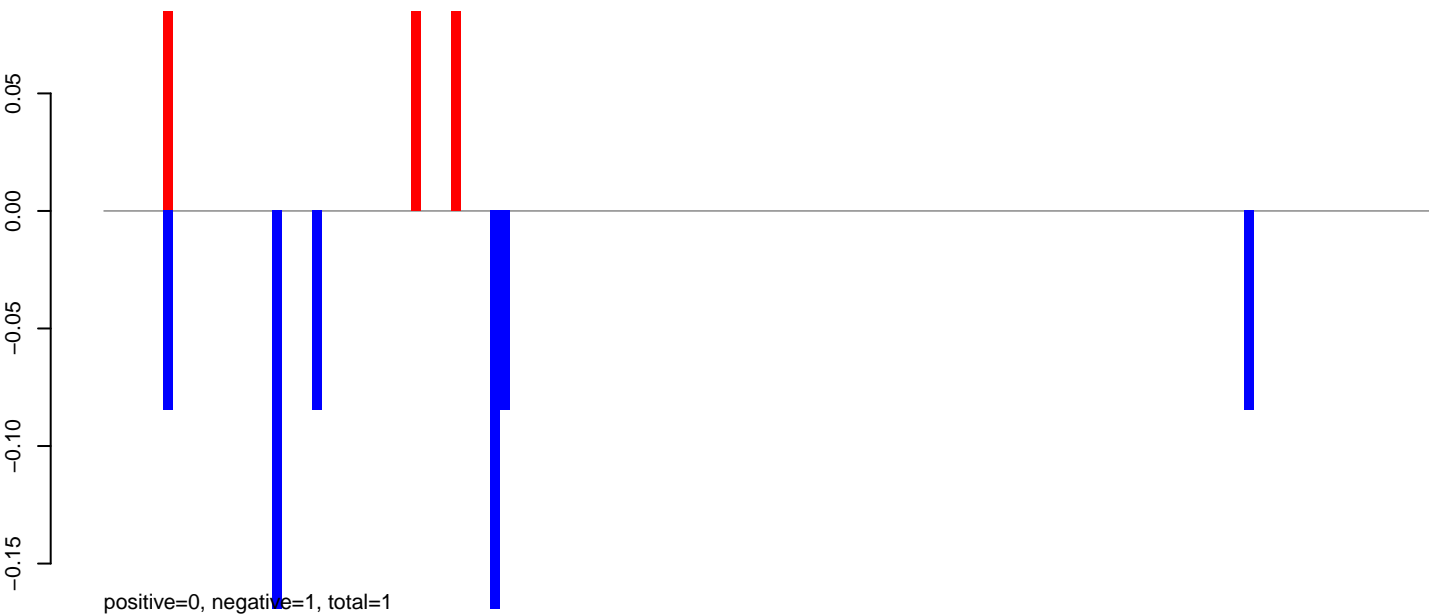
AnGam_Mos55_cells.rep



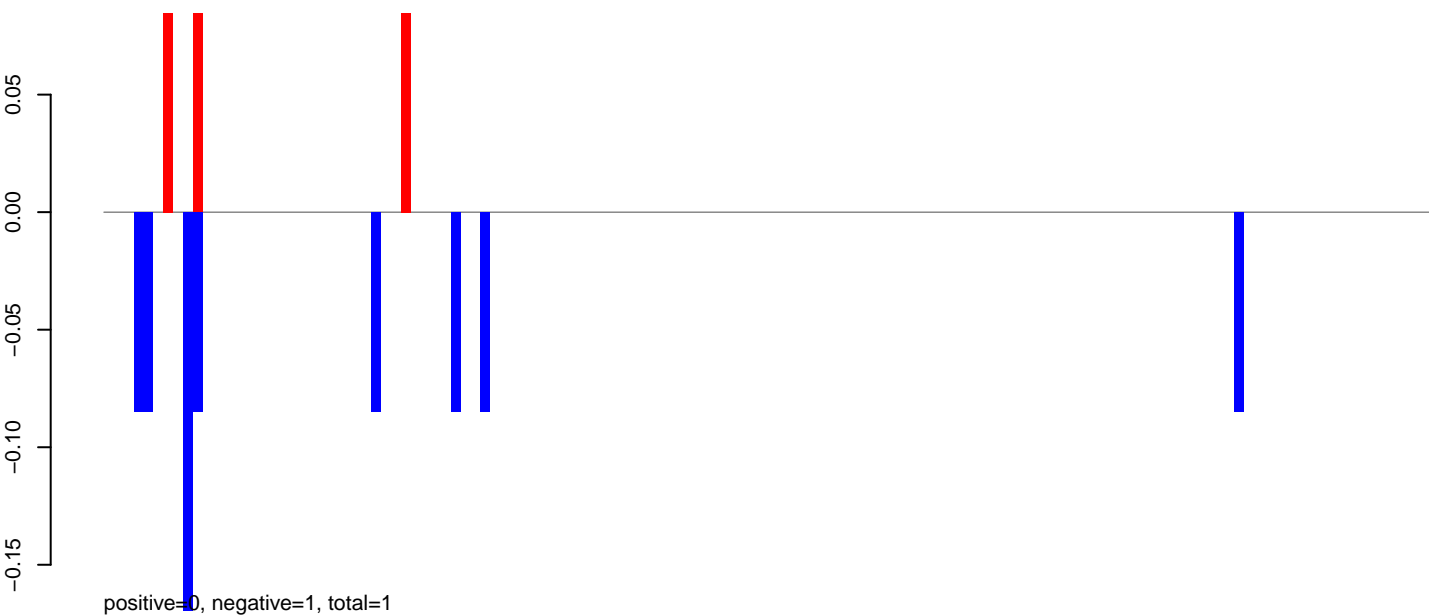
positive=2, negative=2, total=4

Window size=50, length=1158, TE@TF003667-m8bp_Ele6:1-1158

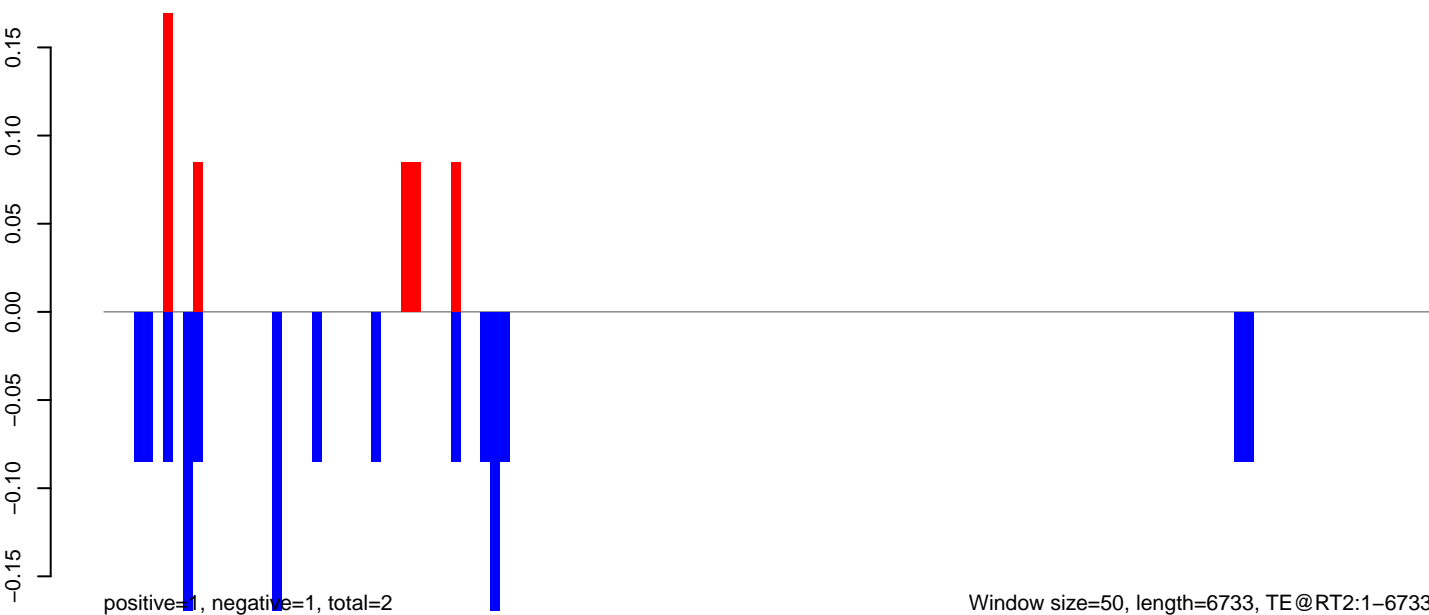
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

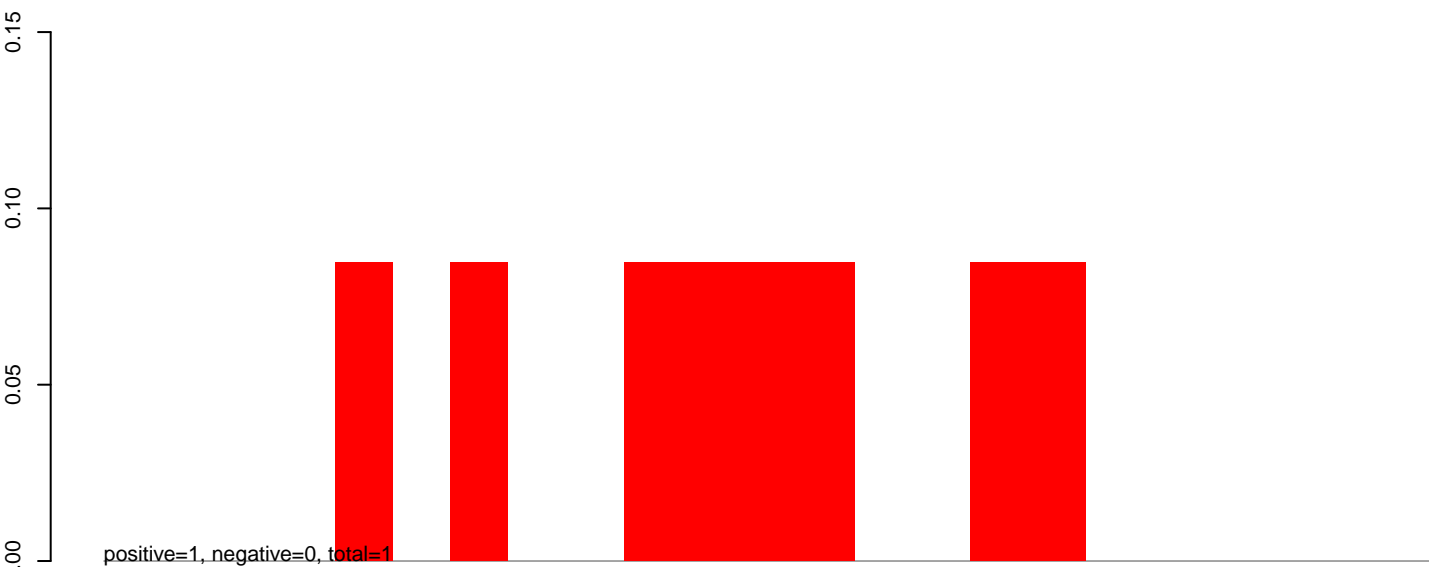


AnGam_Mos55_cells.rep

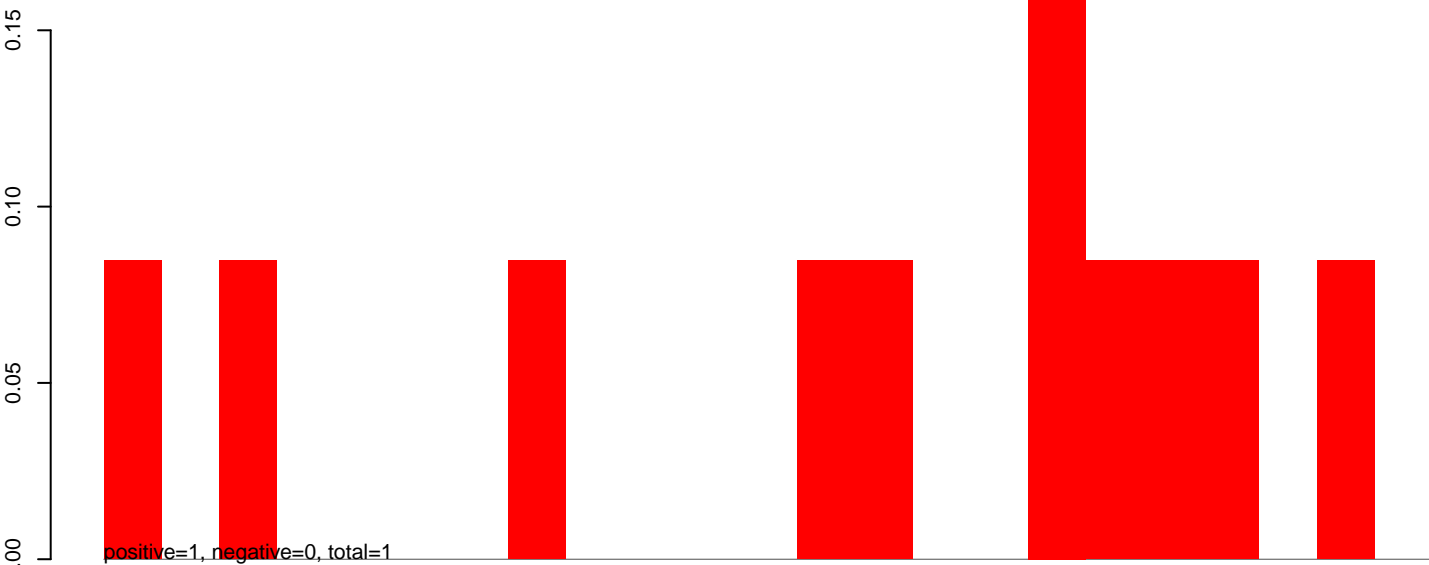


Window size=50, length=6733, TE@RT2:1-6733

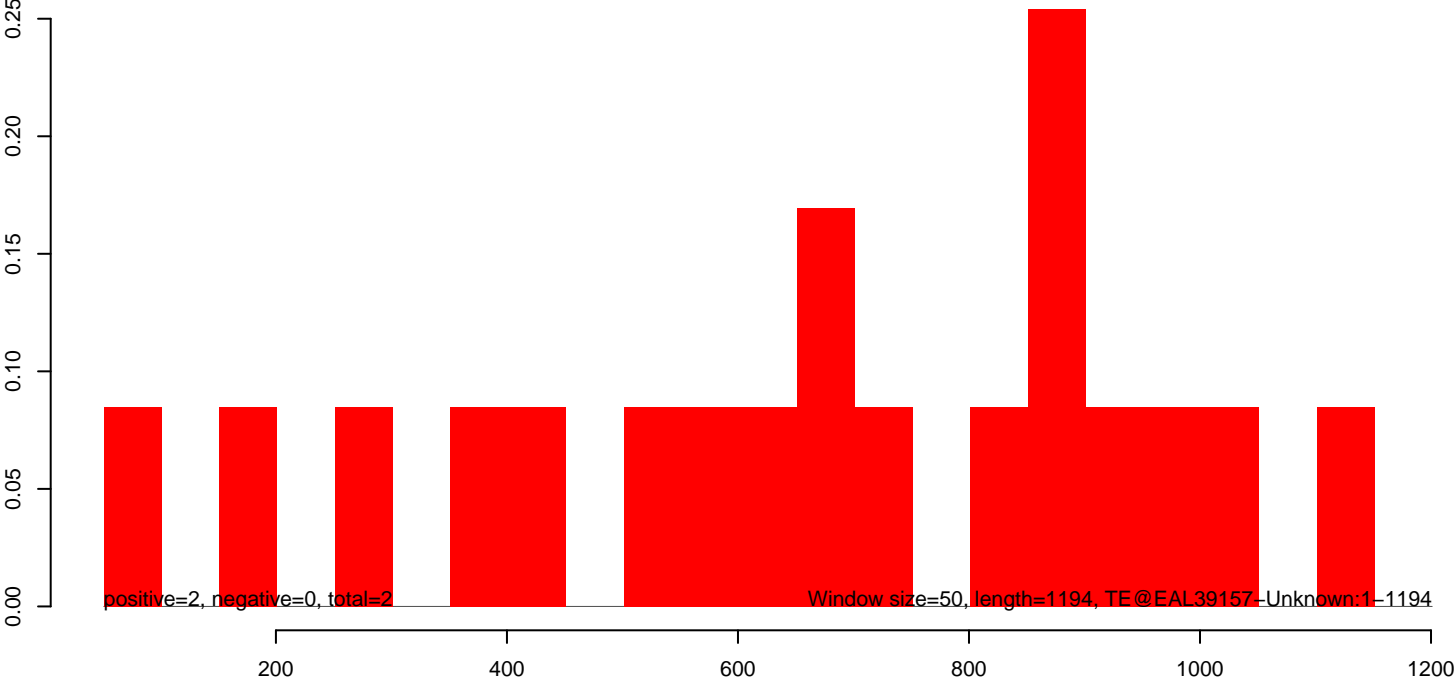
AnGam_Mos55_cells.18_23.rep



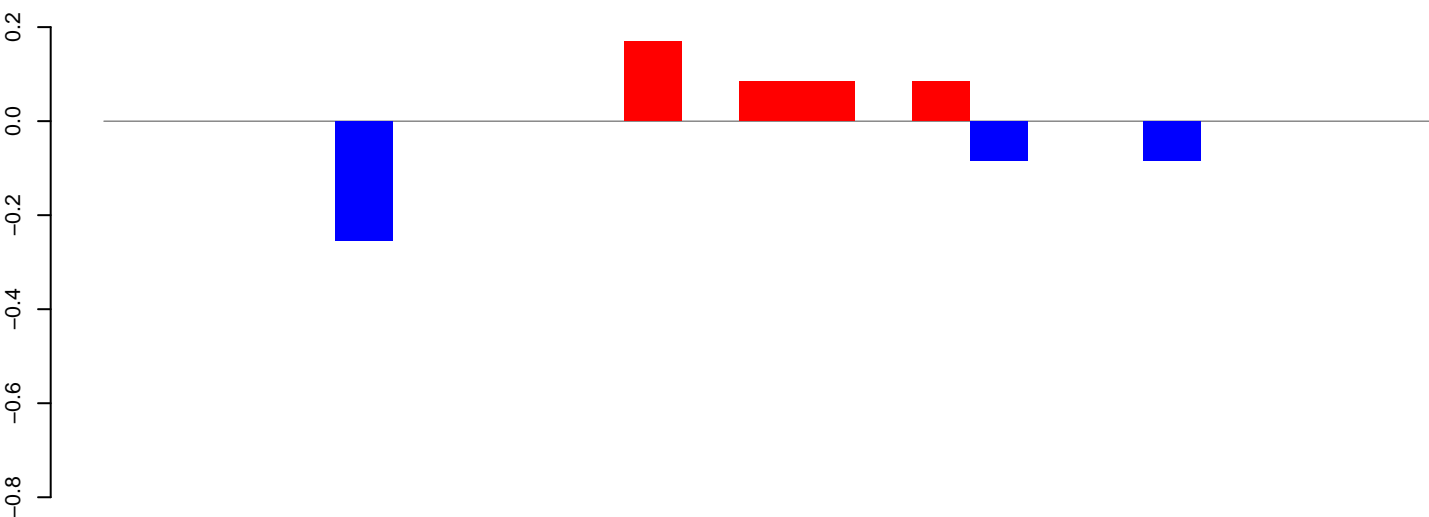
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

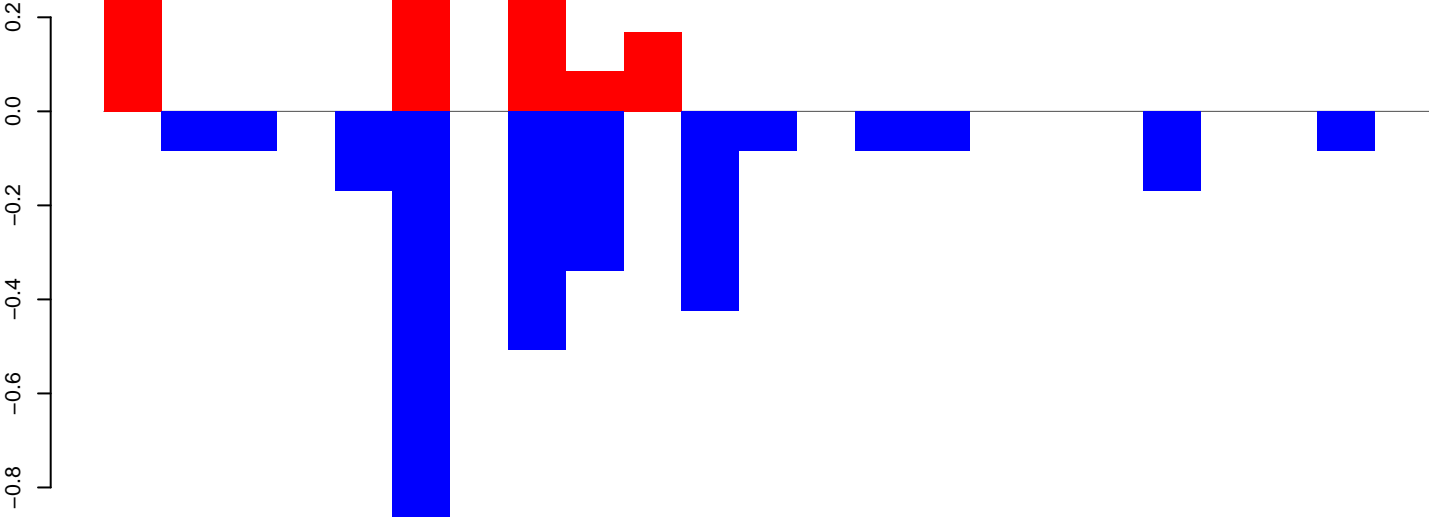


AnGam_Mos55_cells.18_23.rep



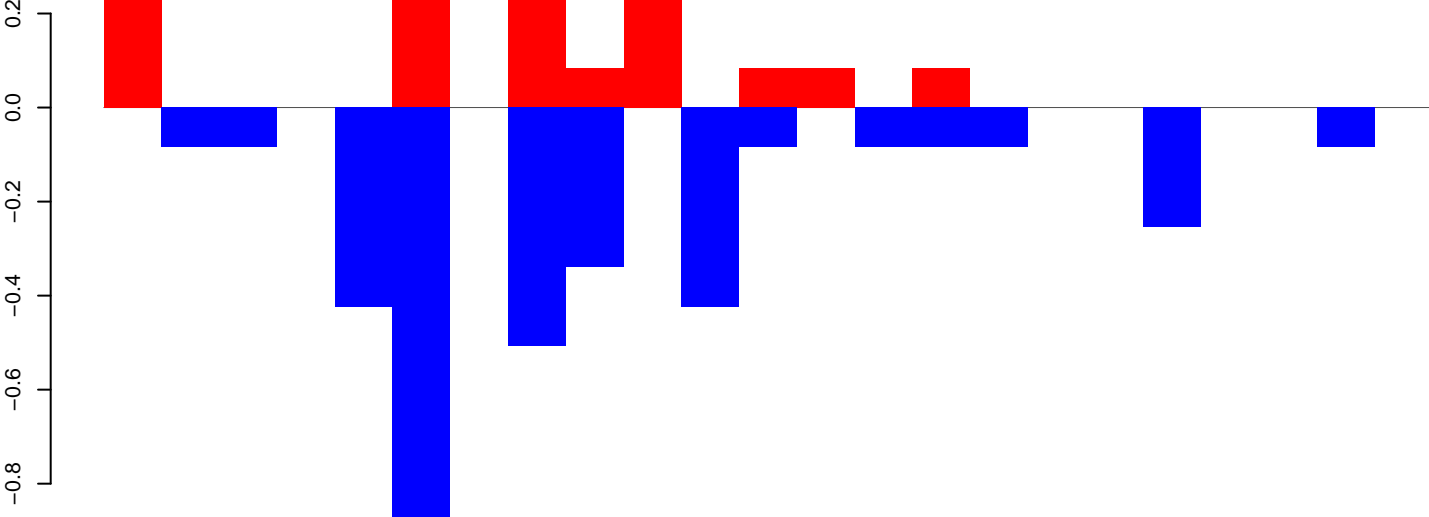
positive=0, negative=0, total=1

AnGam_Mos55_cells.24_35.rep



positive=1, negative=3, total=4

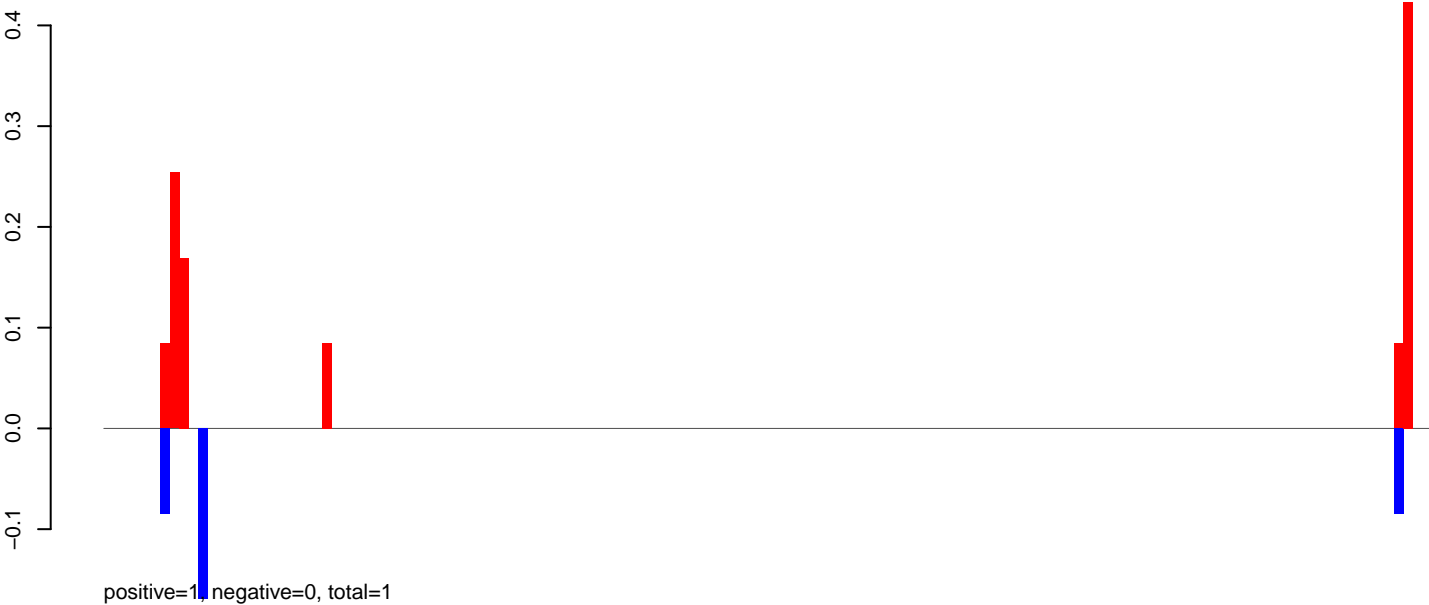
AnGam_Mos55_cells.rep



positive=2, negative=3, total=5

Window size=50, length=1195, TE@Clu-47B_AG:1-1195

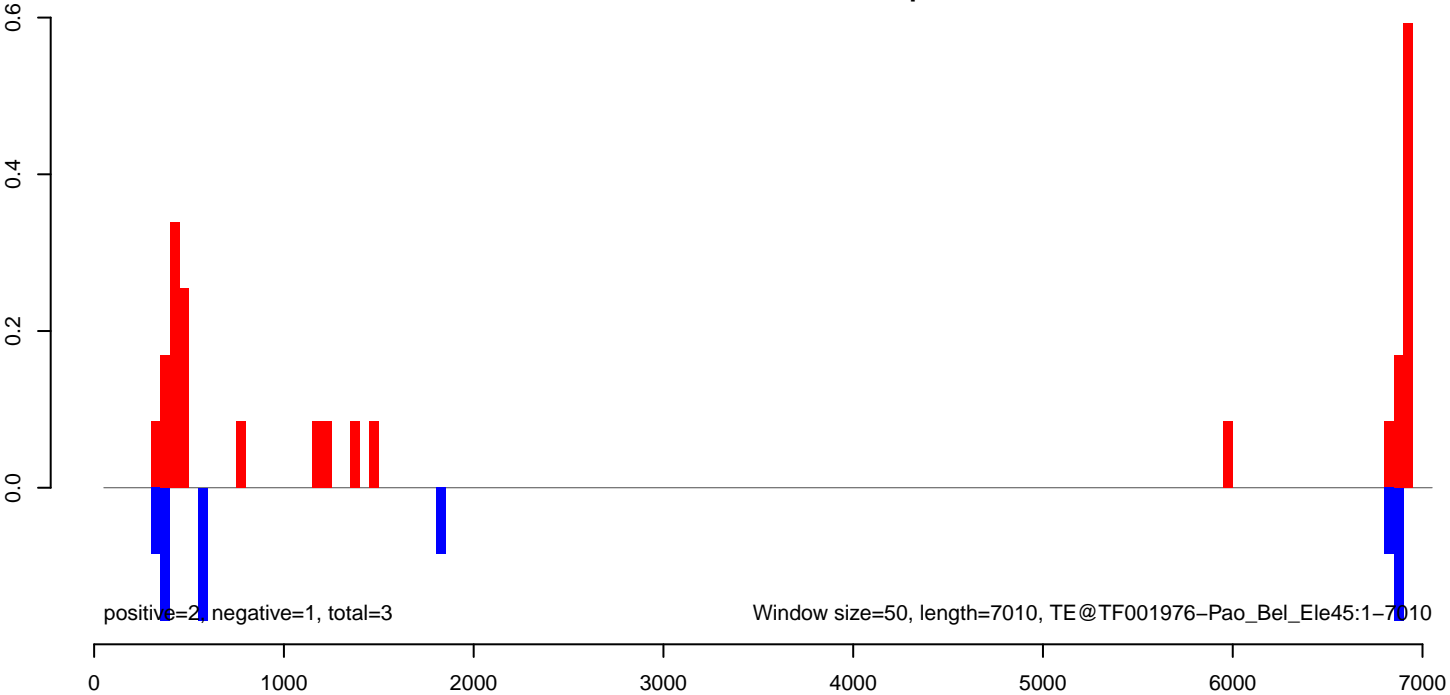
AnGam_Mos55_cells.18_23.rep



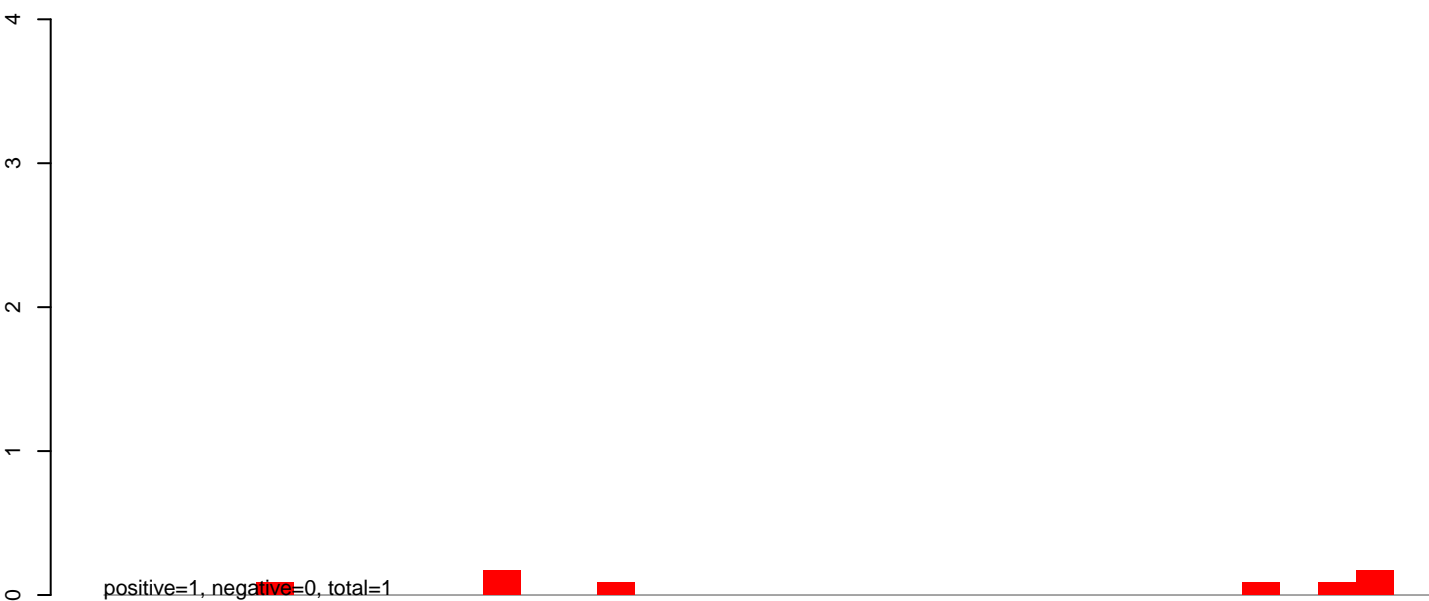
AnGam_Mos55_cells.24_35.rep



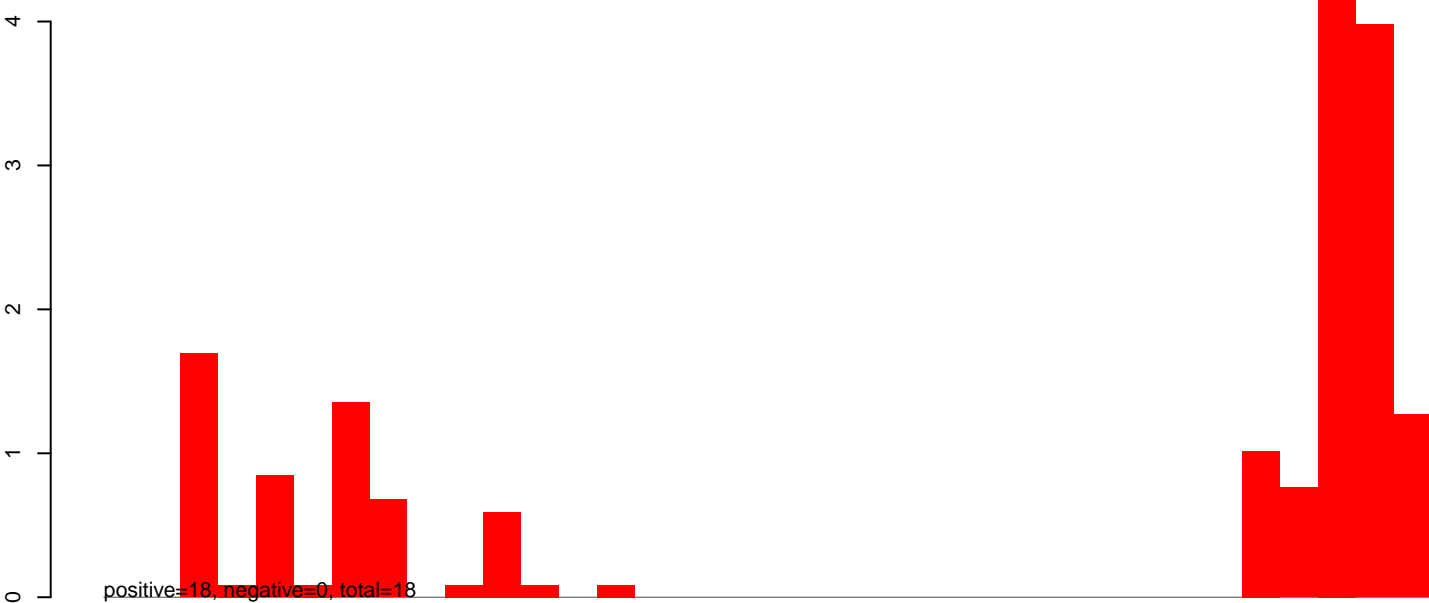
AnGam_Mos55_cells.rep



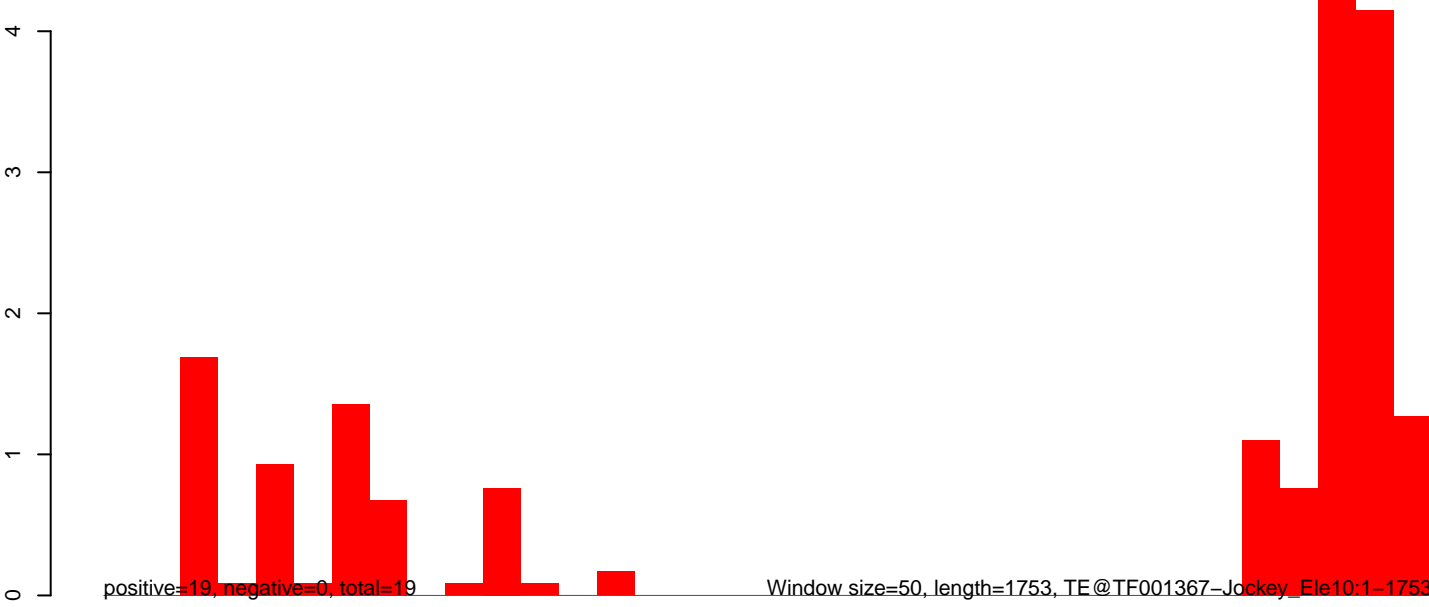
AnGam_Mos55_cells.18_23.rep



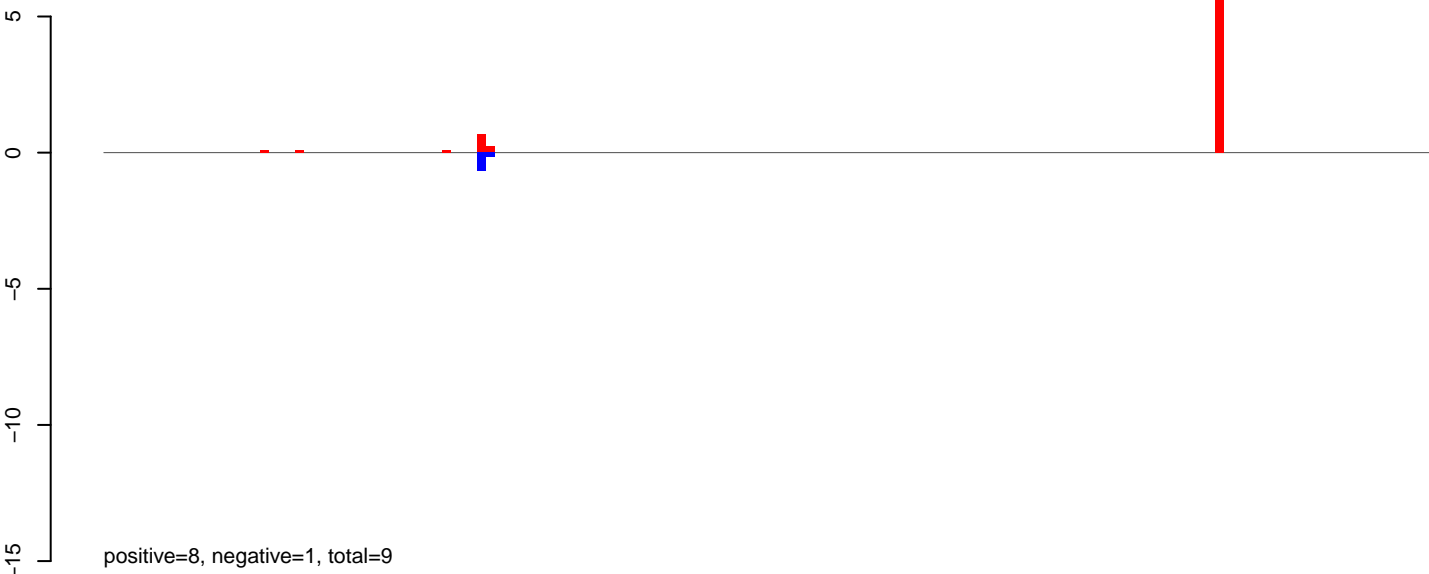
AnGam_Mos55_cells.24_35.rep



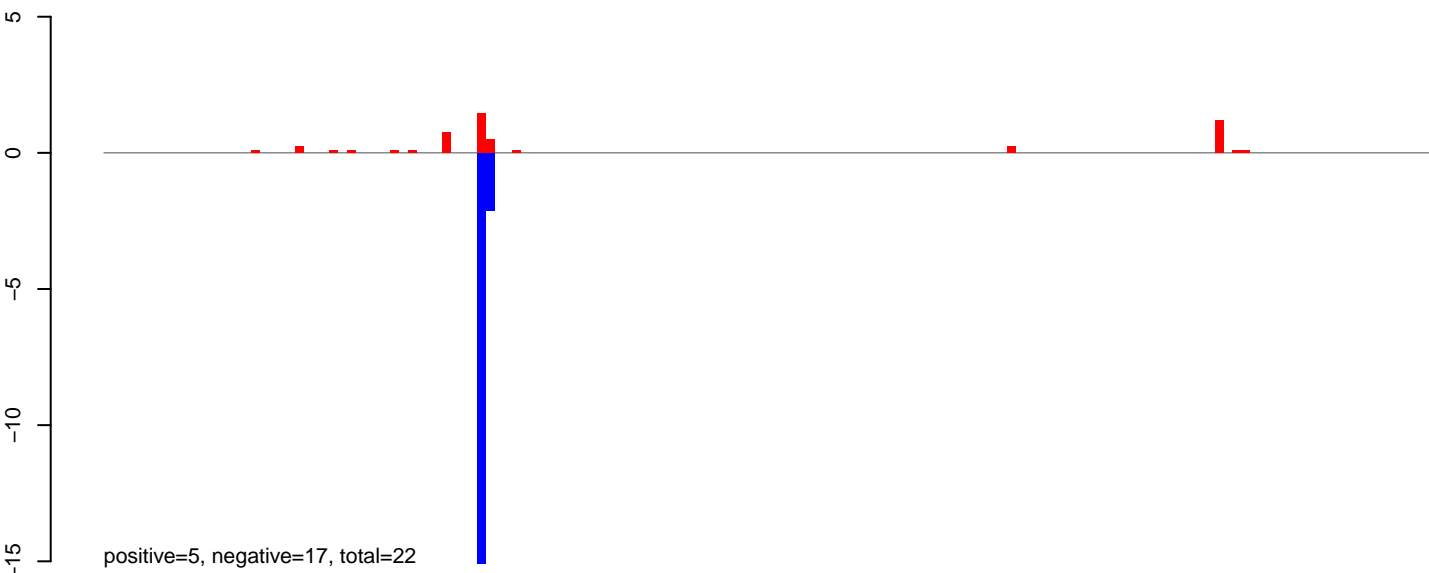
AnGam_Mos55_cells.rep



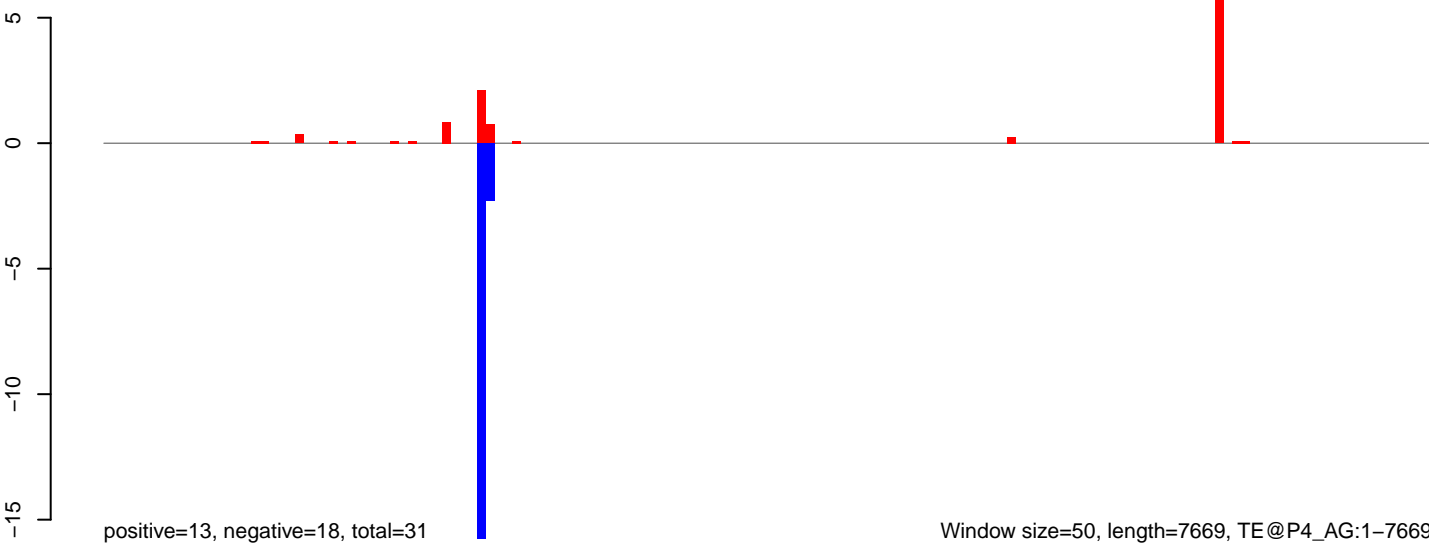
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



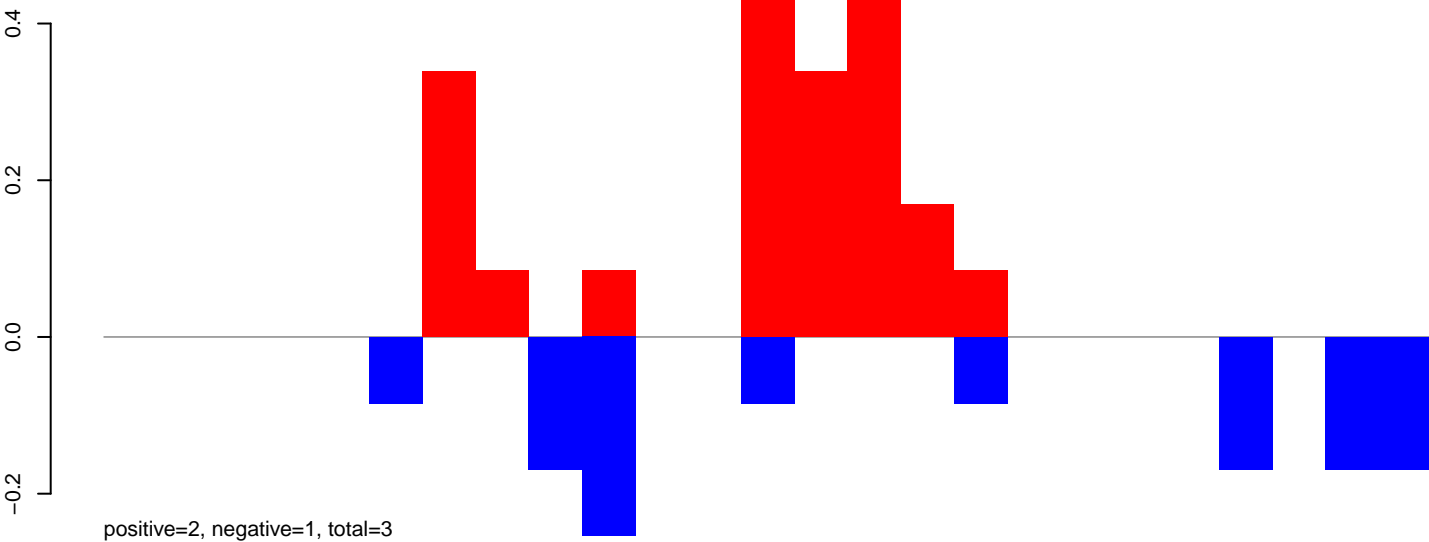
AnGam_Mos55_cells.rep



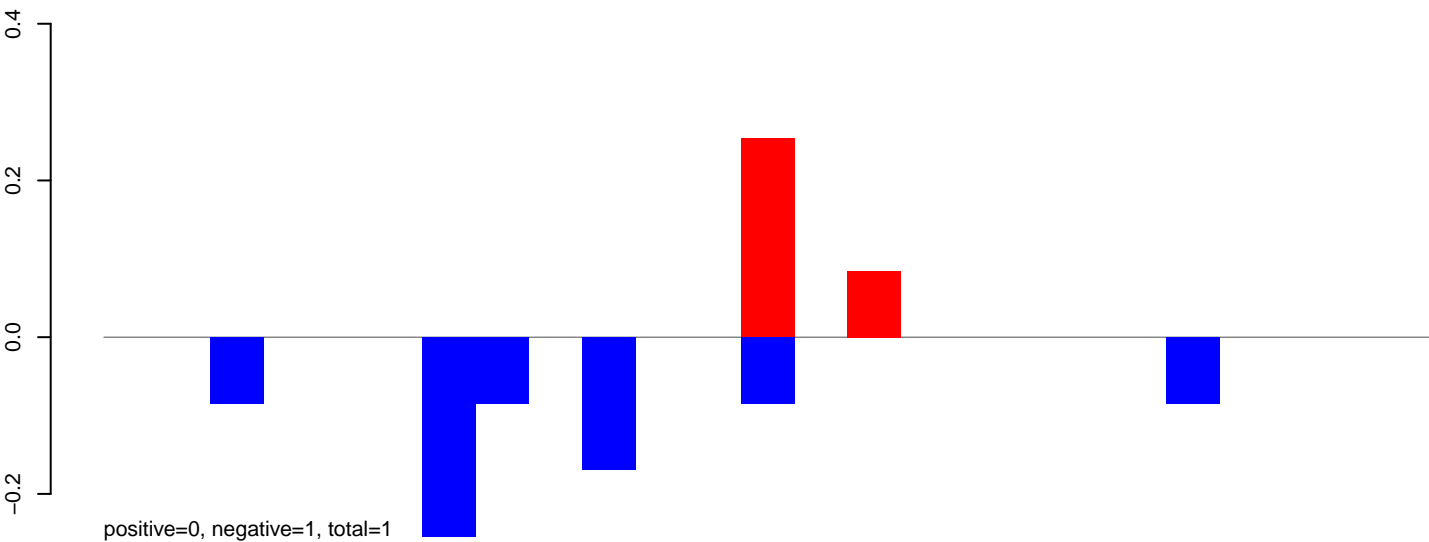
Window size=50, length=7669, TE@P4_AG:1-7669

0 2000 4000 6000 8000

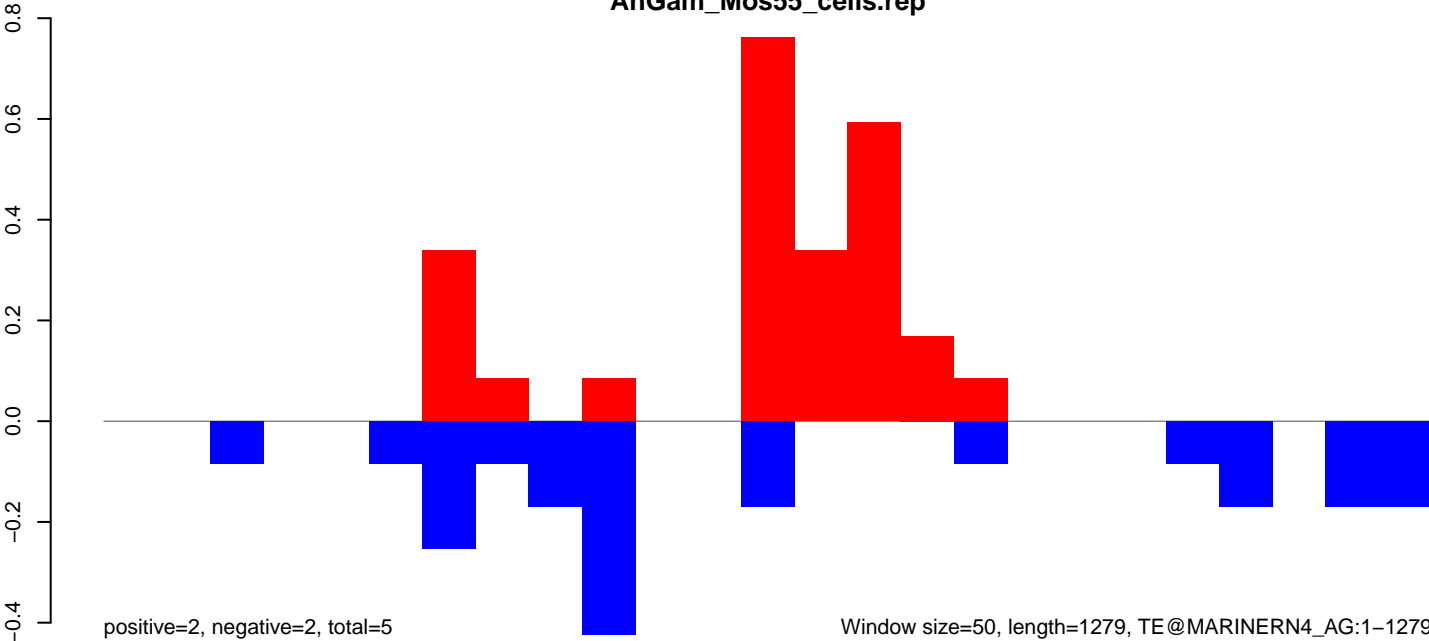
AnGam_Mos55_cells.18_23.rep



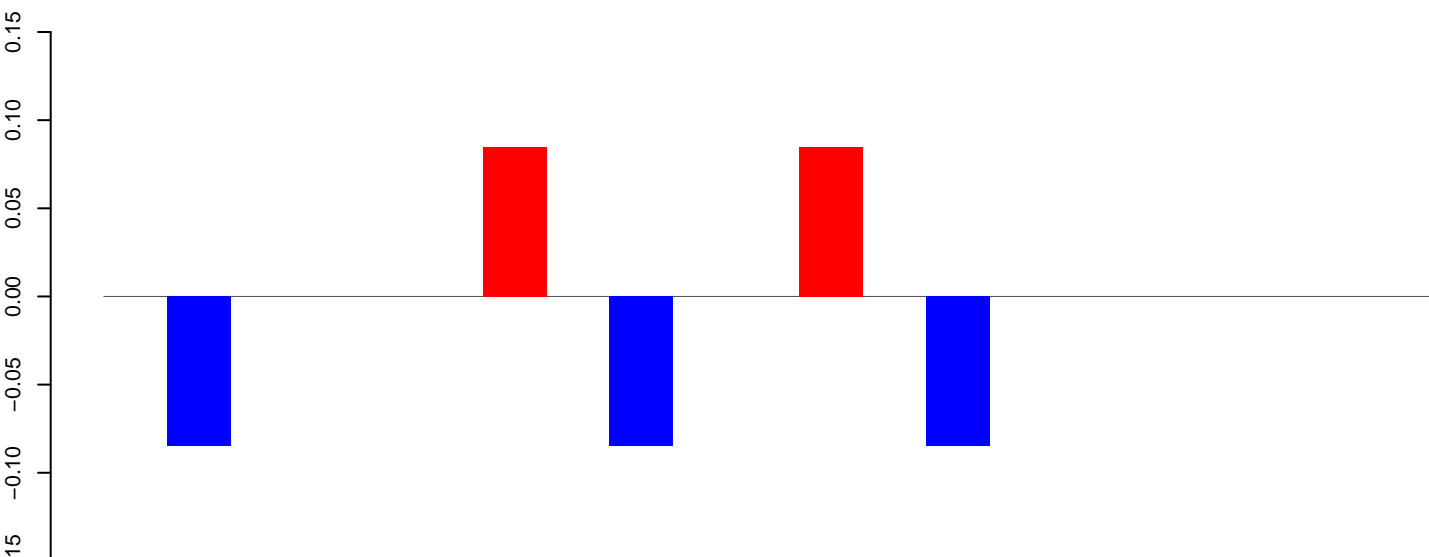
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

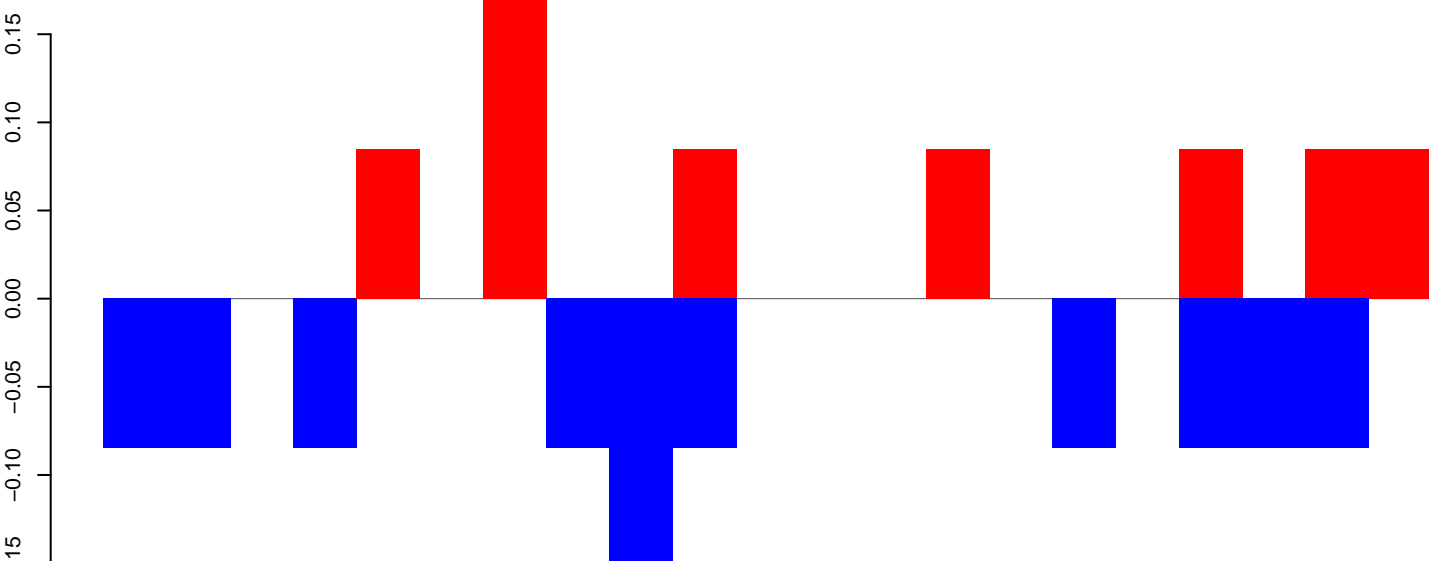


AnGam_Mos55_cells.18_23.rep



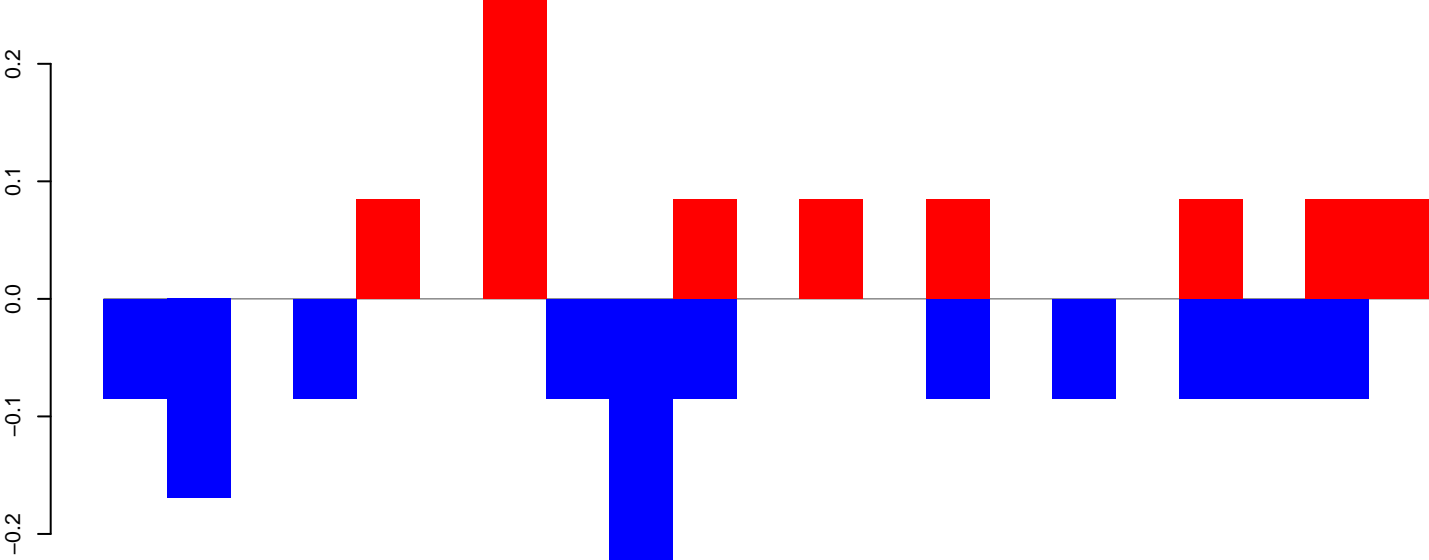
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=1, negative=1, total=2

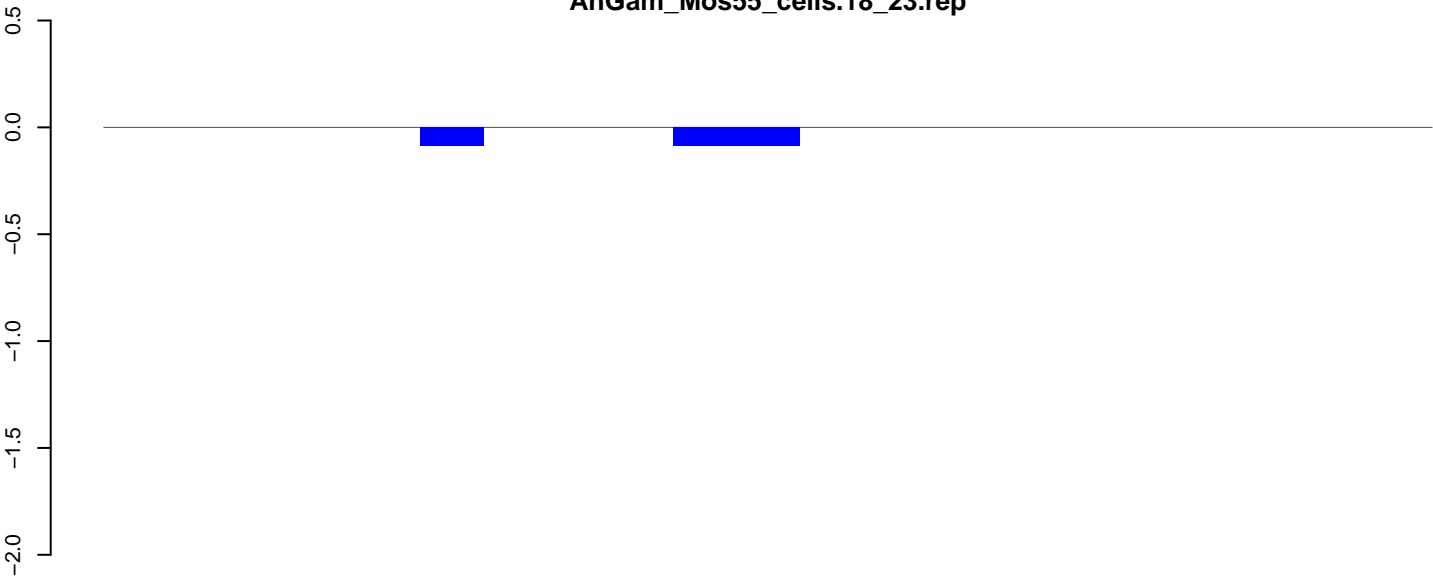
AnGam_Mos55_cells.rep



positive=1, negative=1, total=2

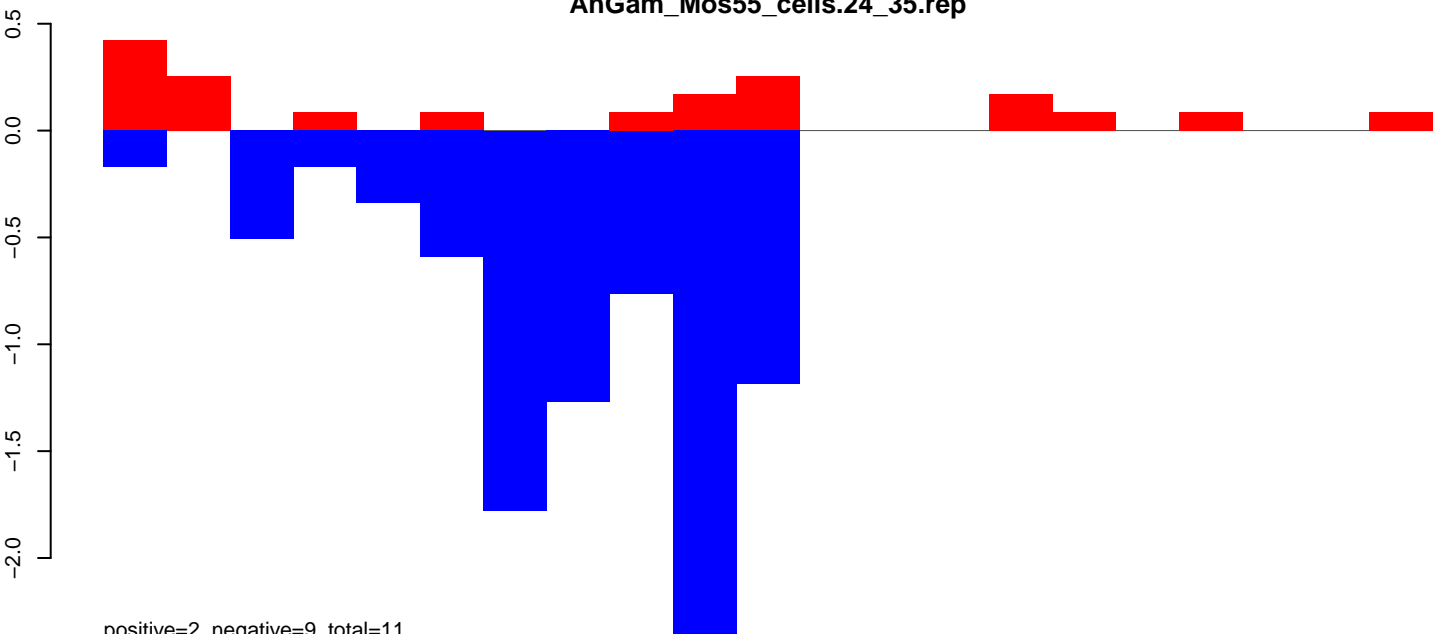
Window size=50, length=1056, TE@MARINERN1_AG:1-1056

AnGam_Mos55_cells.18_23.rep



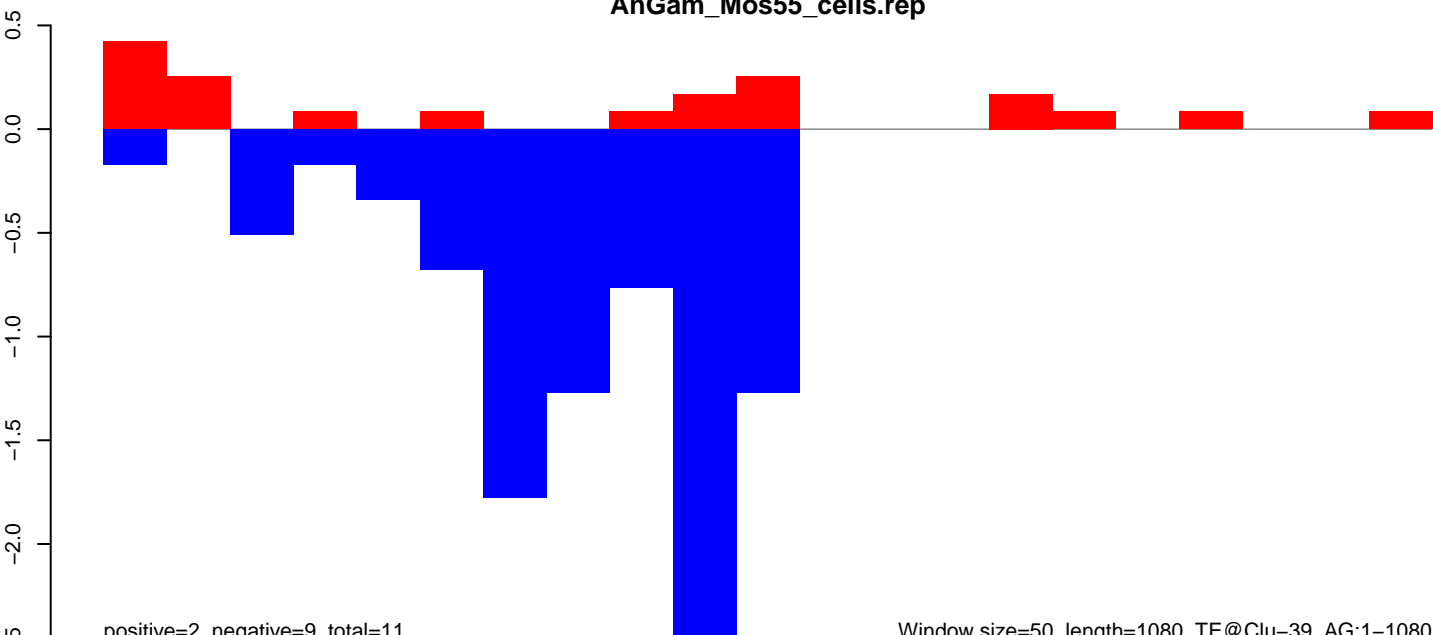
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=2, negative=9, total=11

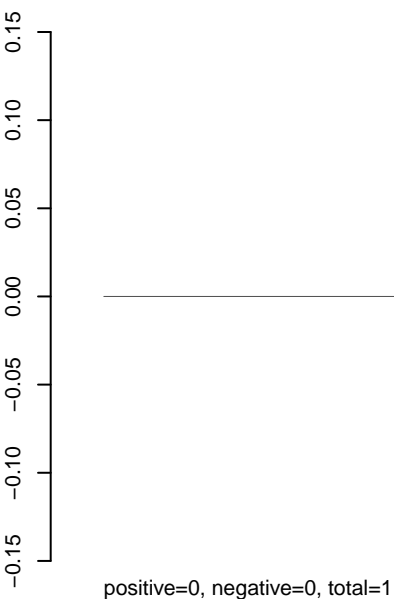
AnGam_Mos55_cells.rep



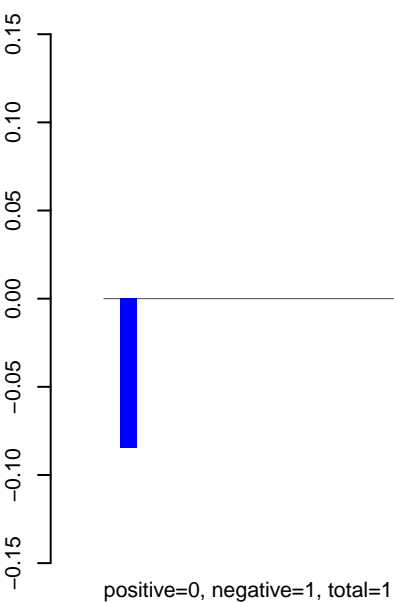
positive=2, negative=9, total=11

Window size=50, length=1080, TE@Clu-39_AG:1-1080

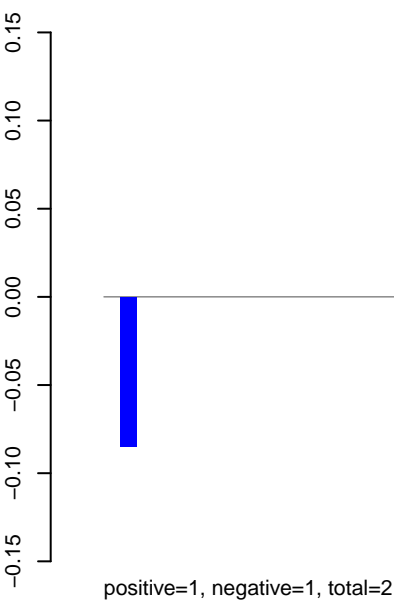
AnGam_Mos55_cells.18_23.rep



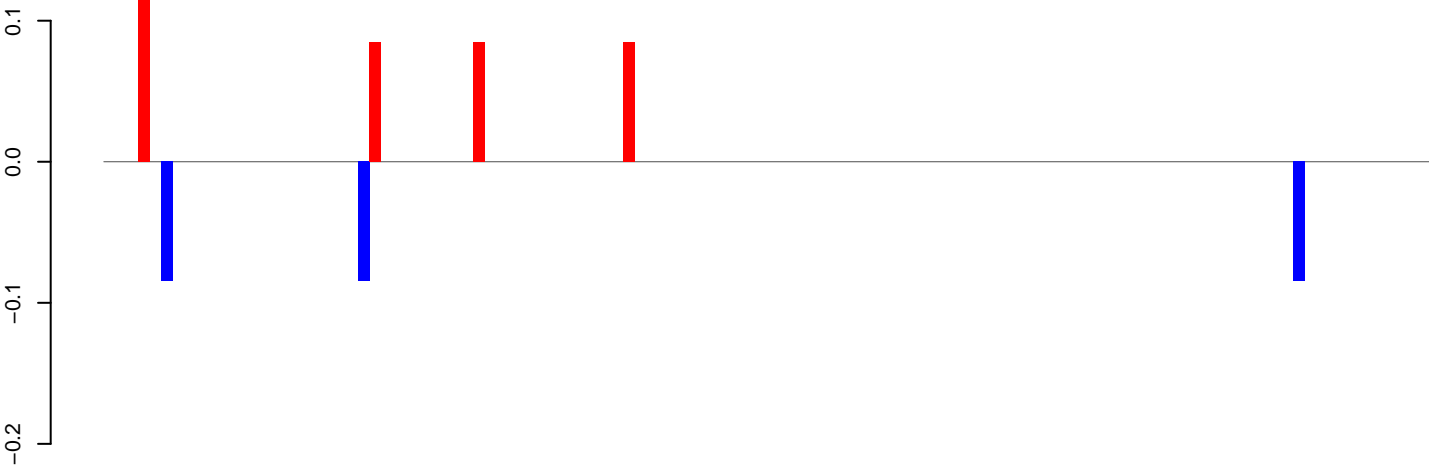
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



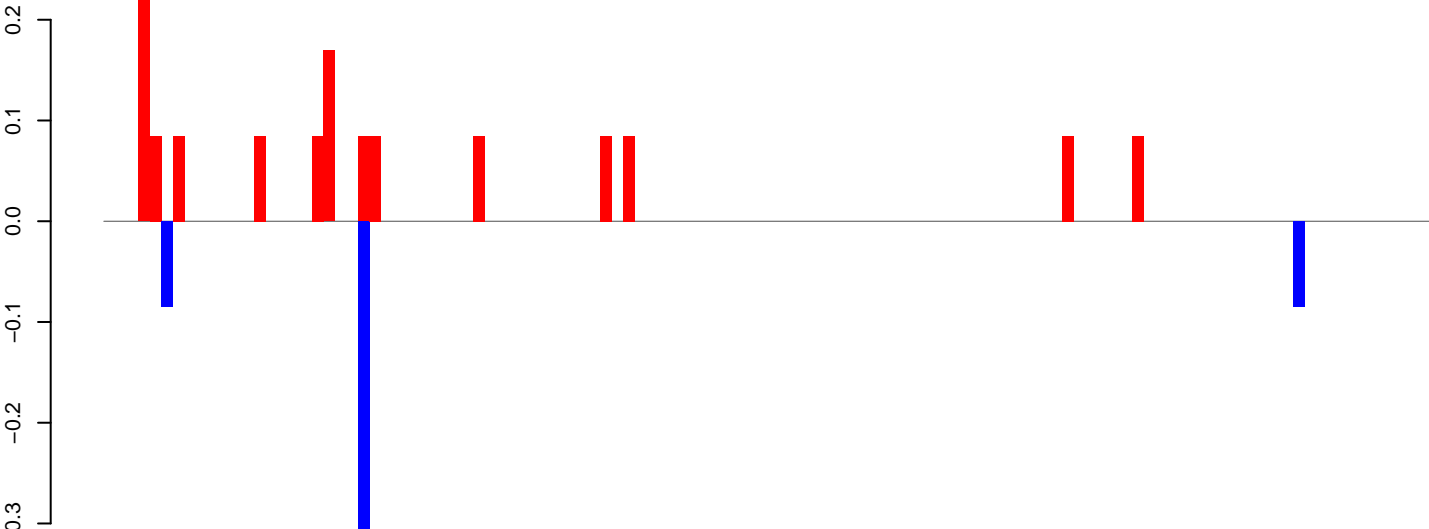
positive=0, negative=0, total=1

AnGam_Mos55_cells.24_35.rep



positive=1, negative=0, total=1

AnGam_Mos55_cells.rep

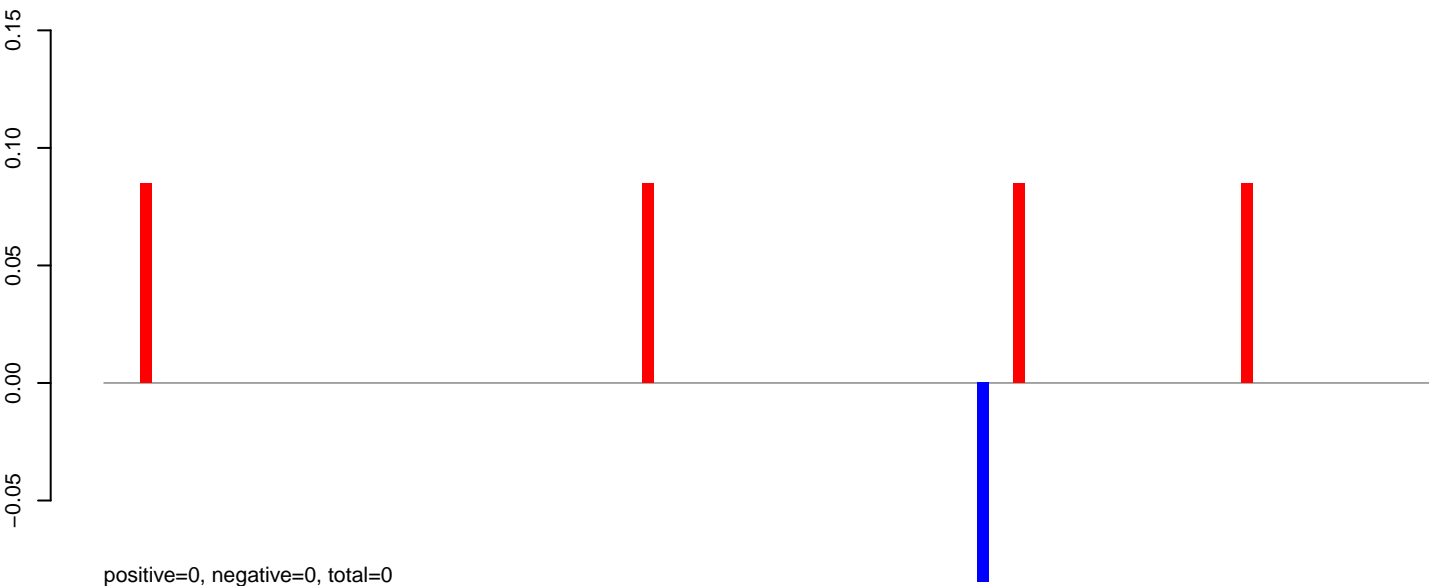


positive=1, negative=1, total=2

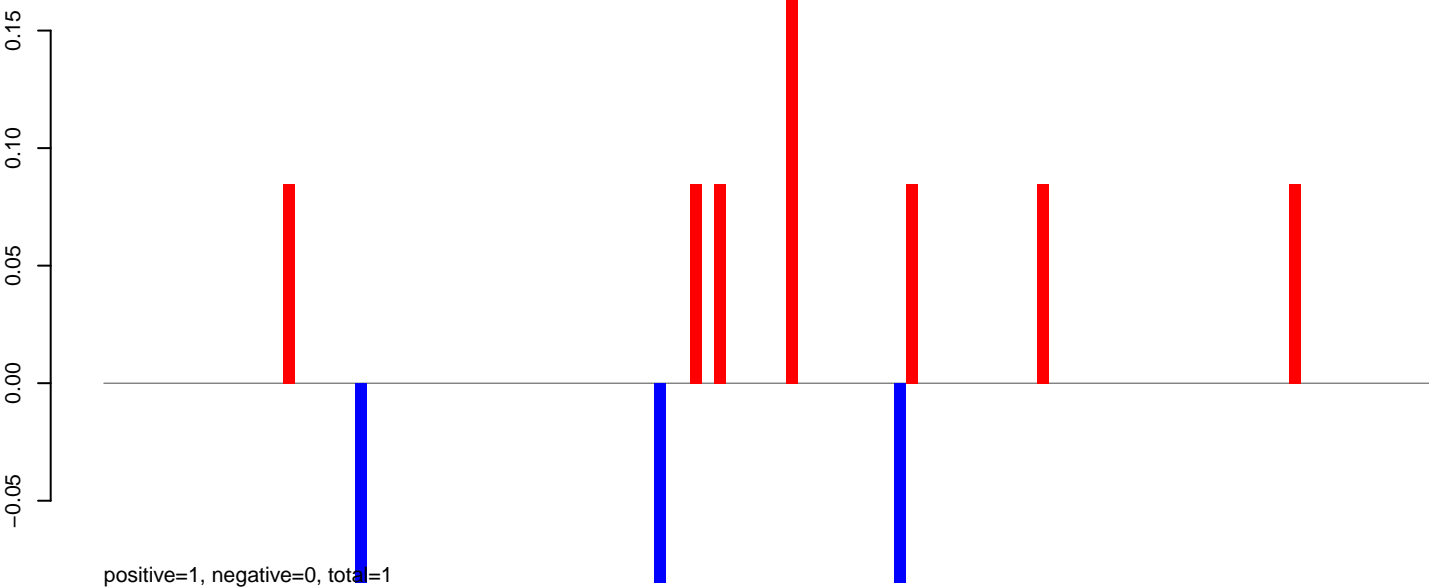
Window size=50, length=5797, TE@BEL16-LTR-I_AG:1-5797

0 1000 2000 3000 4000 5000 6000

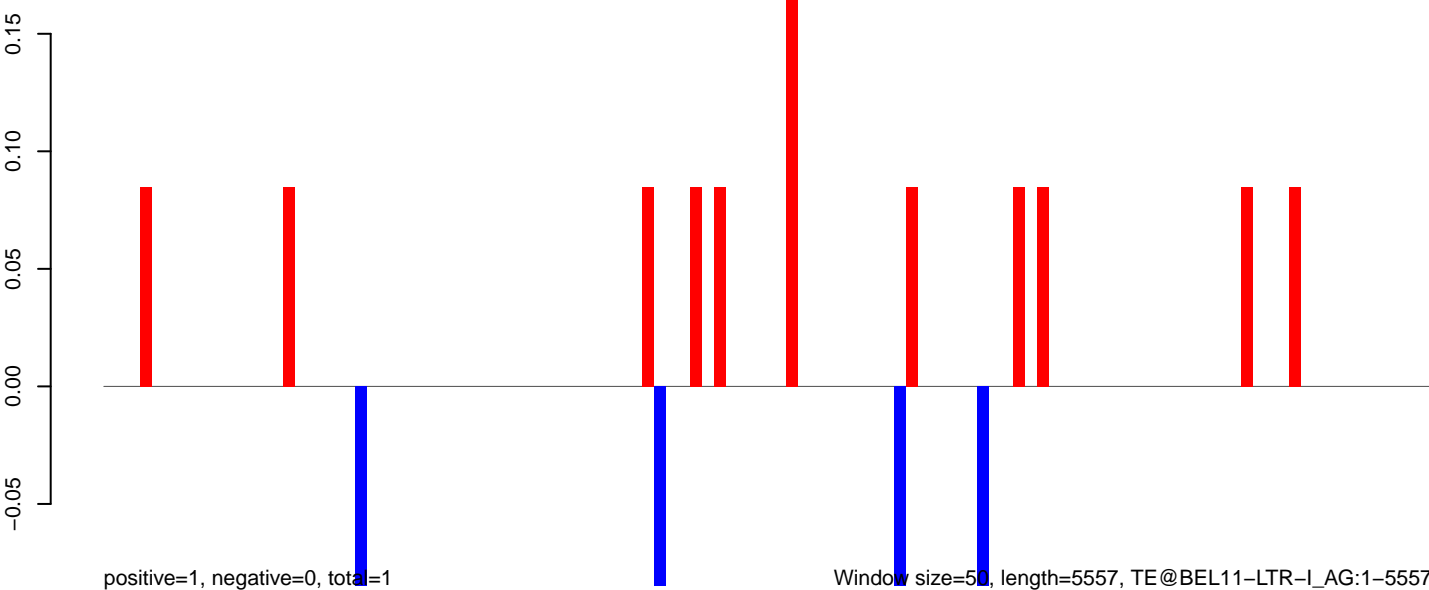
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

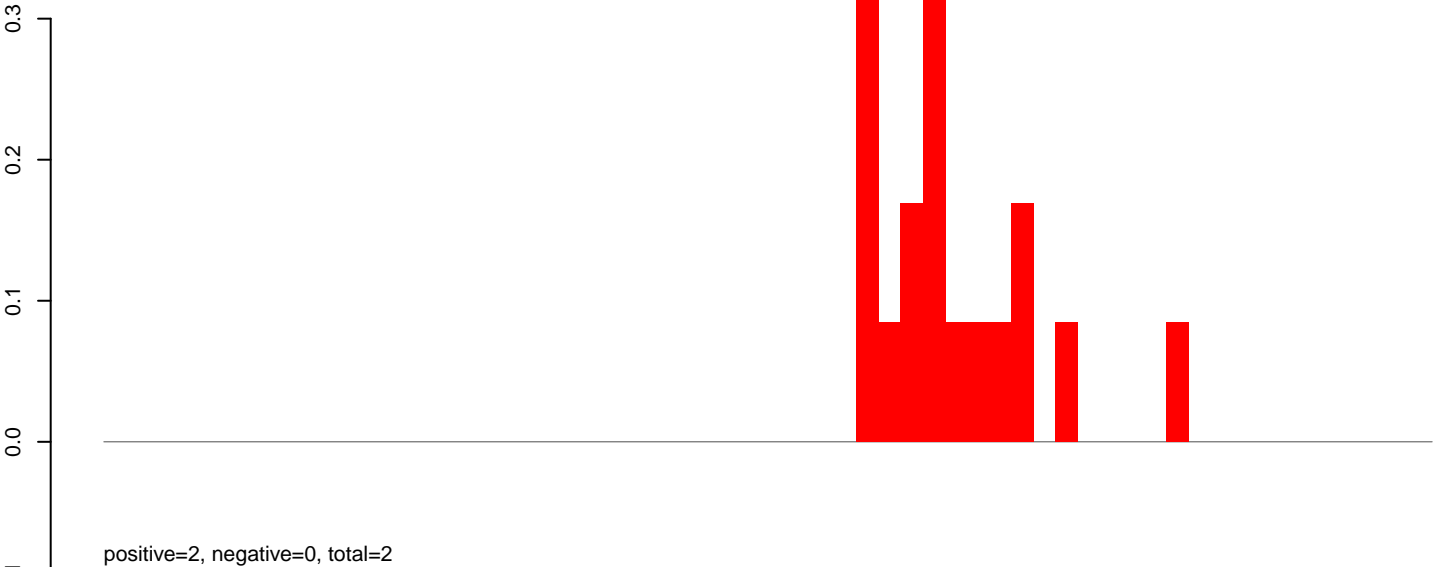


0 1000 2000 3000 4000 5000

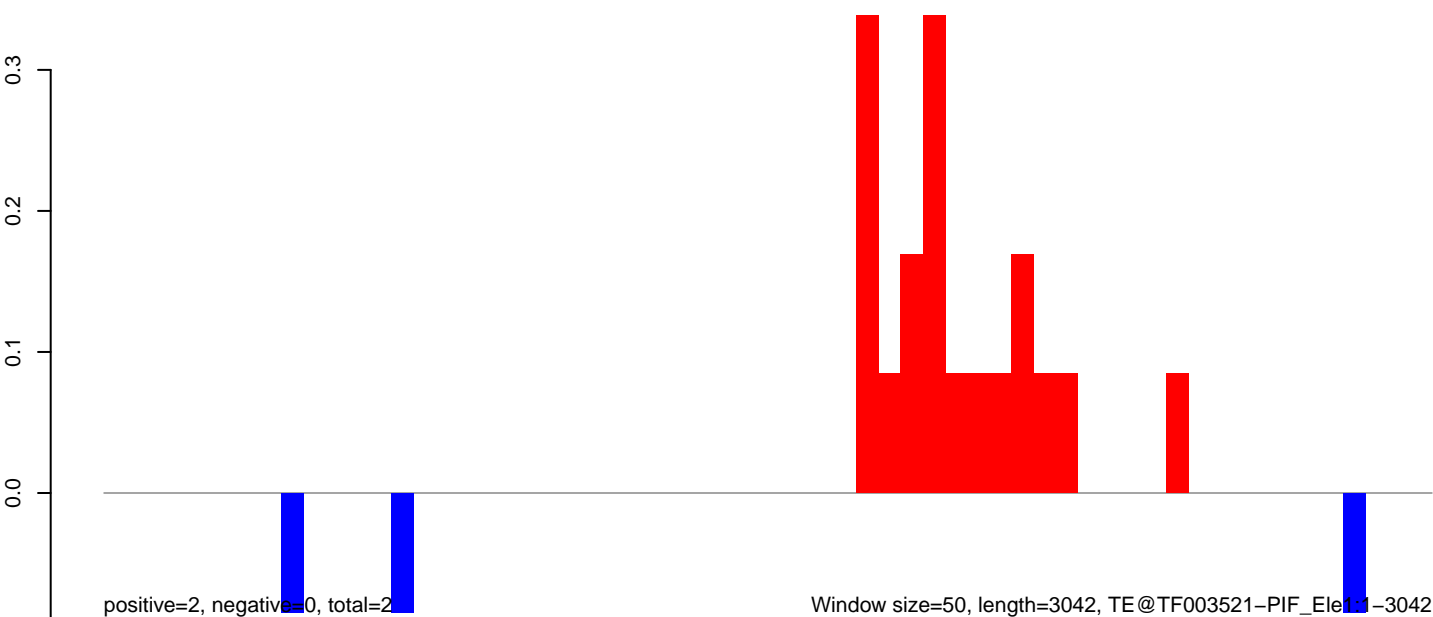
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

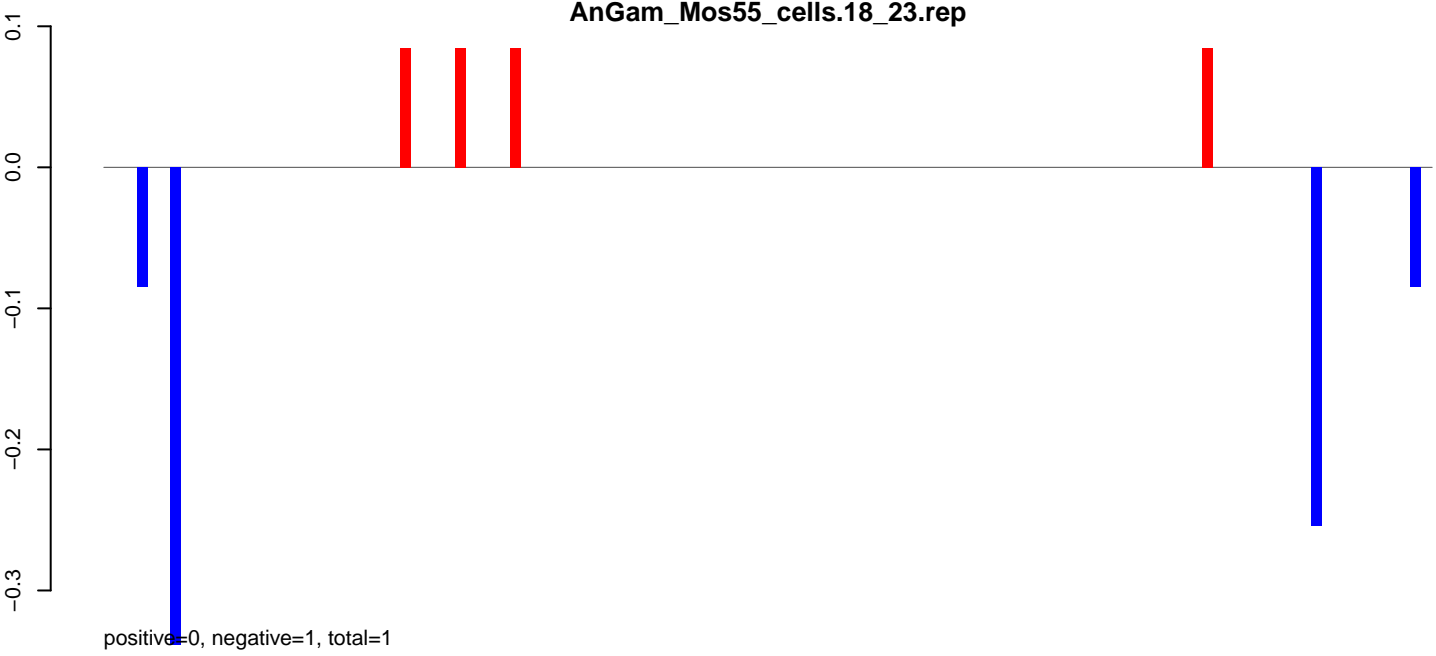


AnGam_Mos55_cells.rep

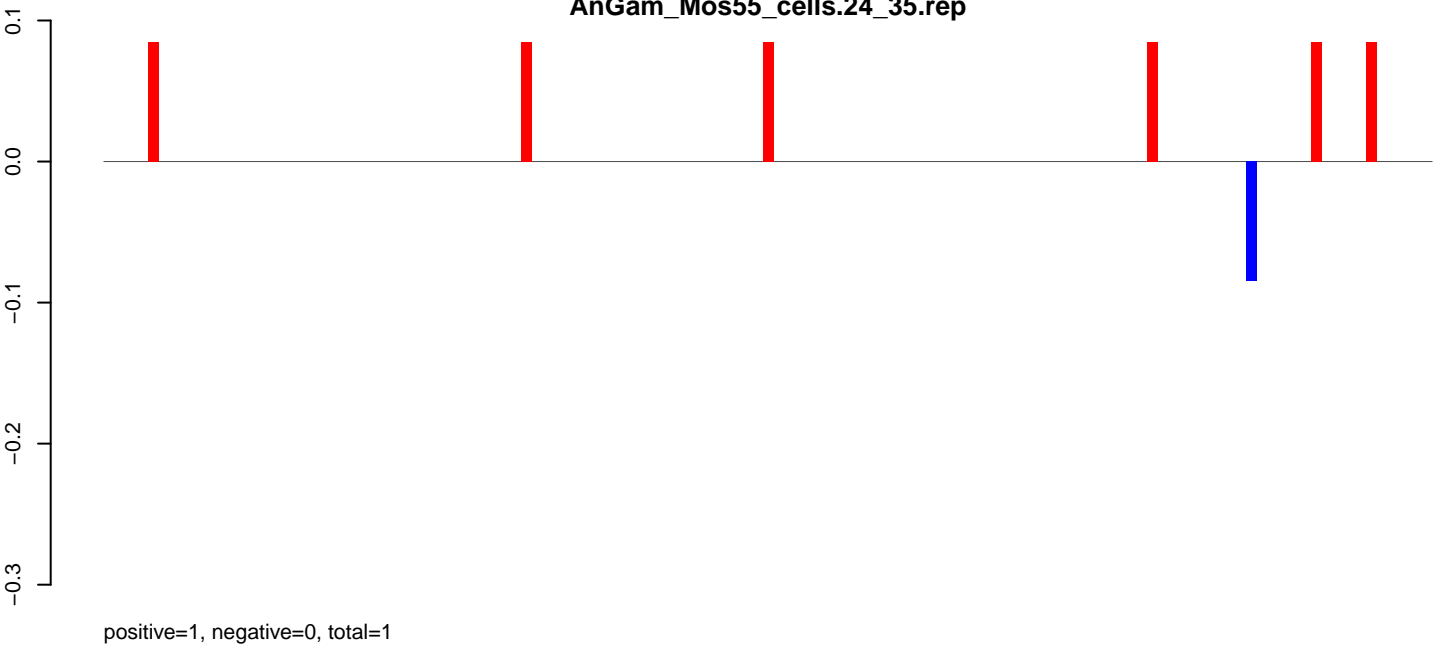


Window size=50, length=3042, TE@TF003521-PIF_Element-3042

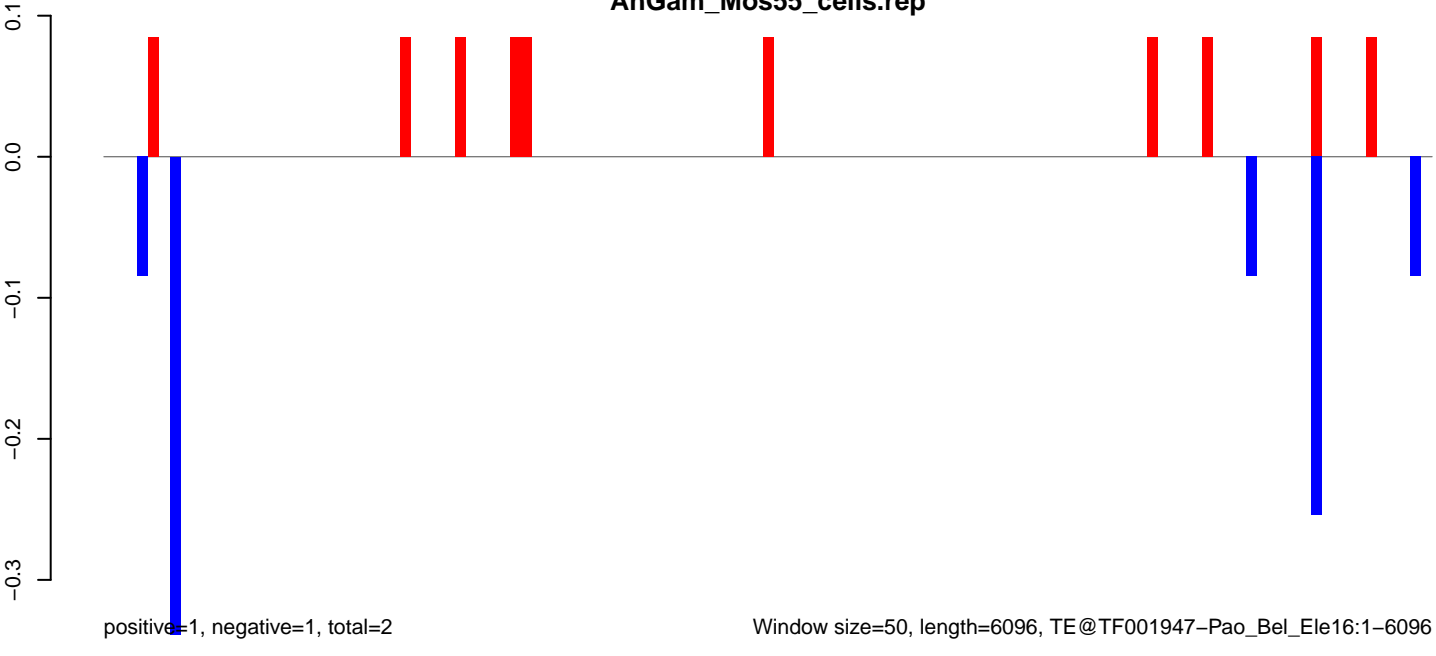
AnGam_Mos55_cells.18_23.rep



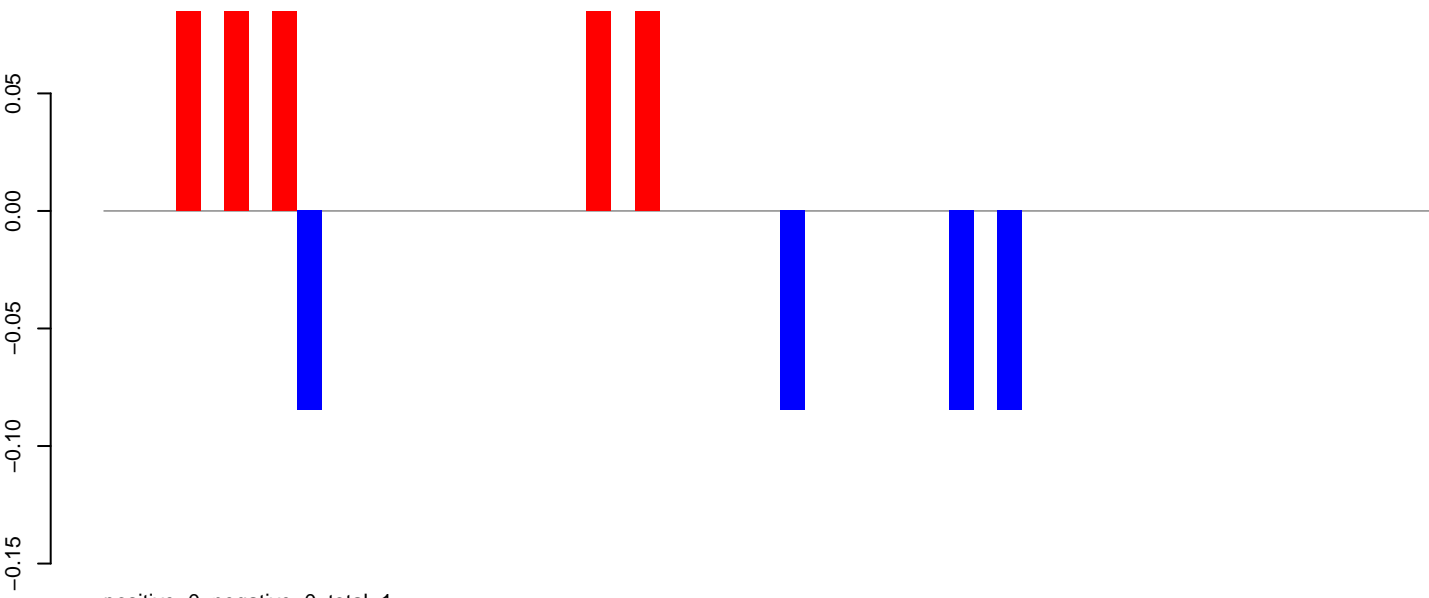
AnGam_Mos55_cells.24_35.rep



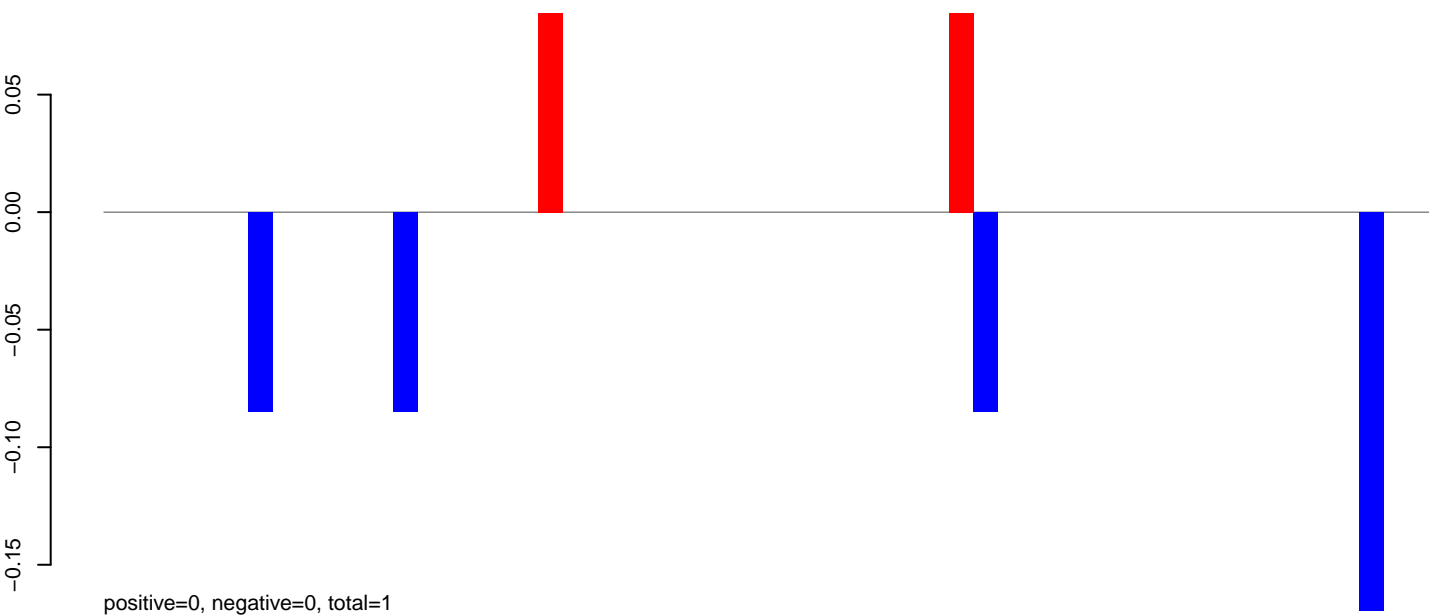
AnGam_Mos55_cells.rep



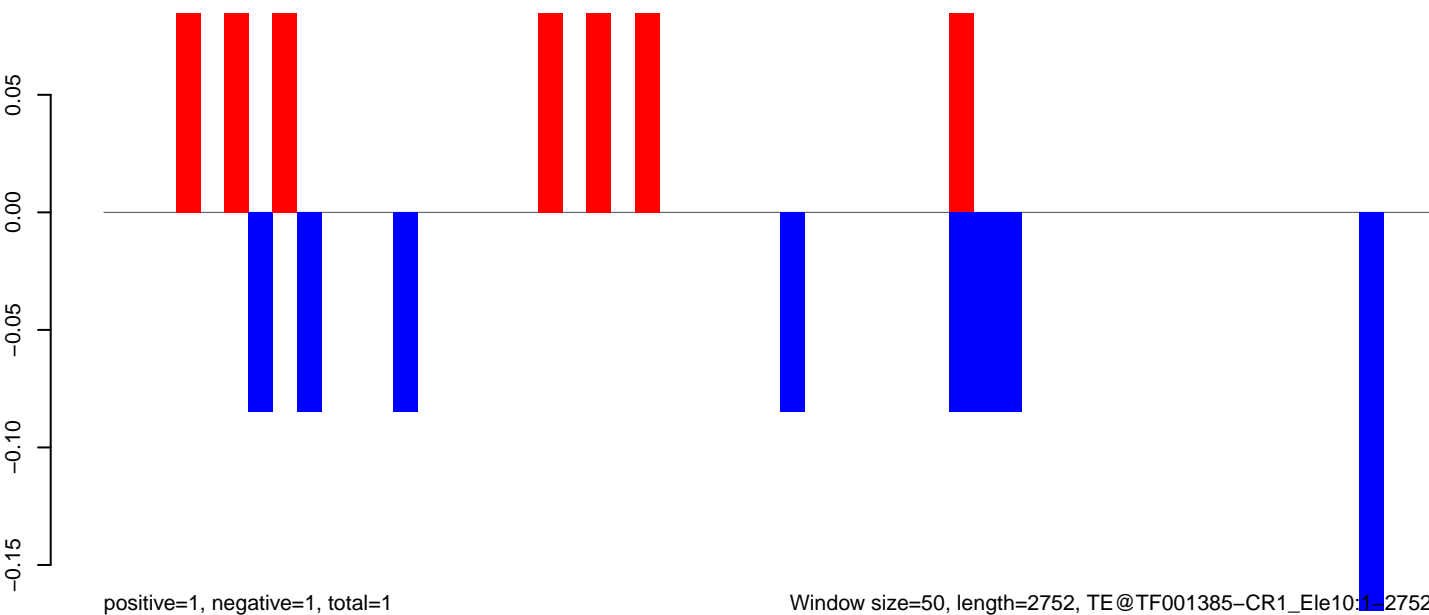
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



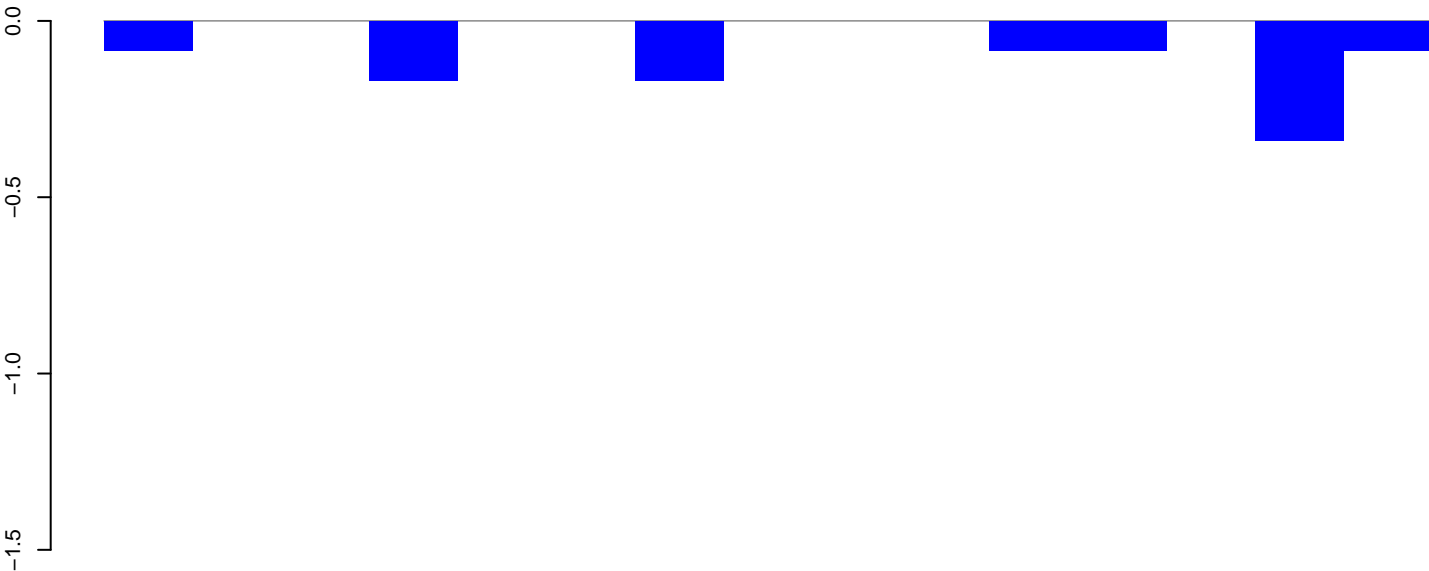
AnGam_Mos55_cells.rep



Window size=50, length=2752, TE@TF001385-CR1_Ele10:1-2752

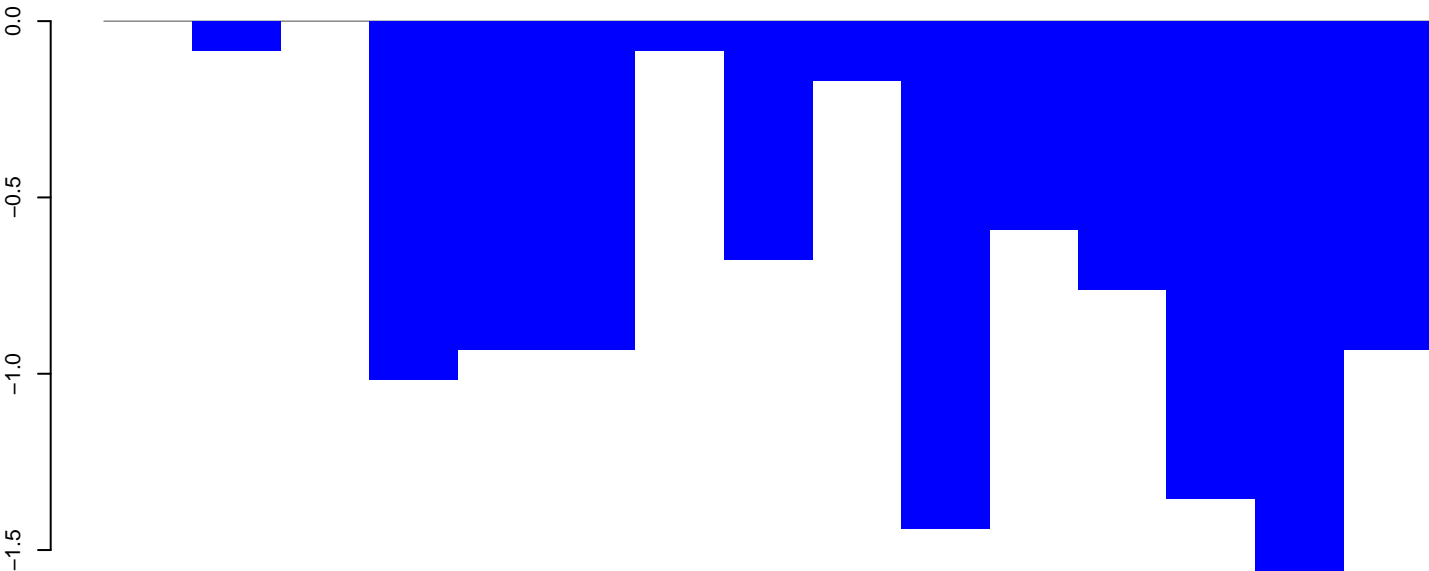
0 500 1000 1500 2000 2500

AnGam_Mos55_cells.18_23.rep



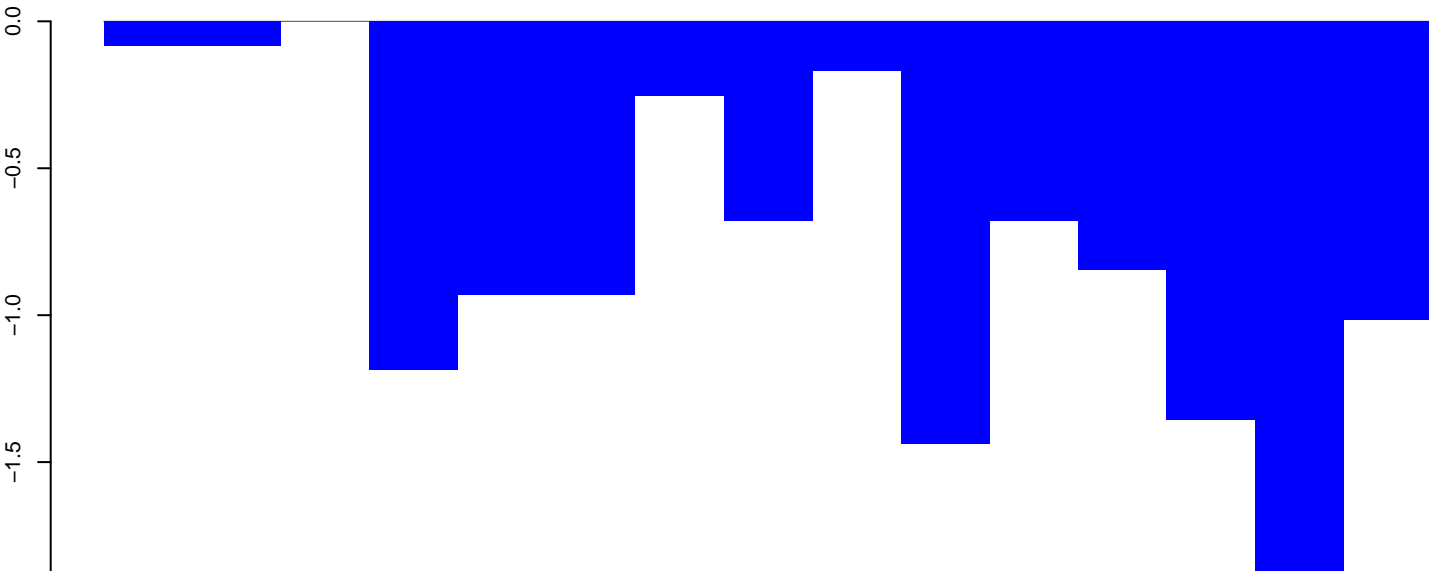
positive=0, negative=1, total=1

AnGam_Mos55_cells.24_35.rep



positive=0, negative=12, total=12

AnGam_Mos55_cells.rep

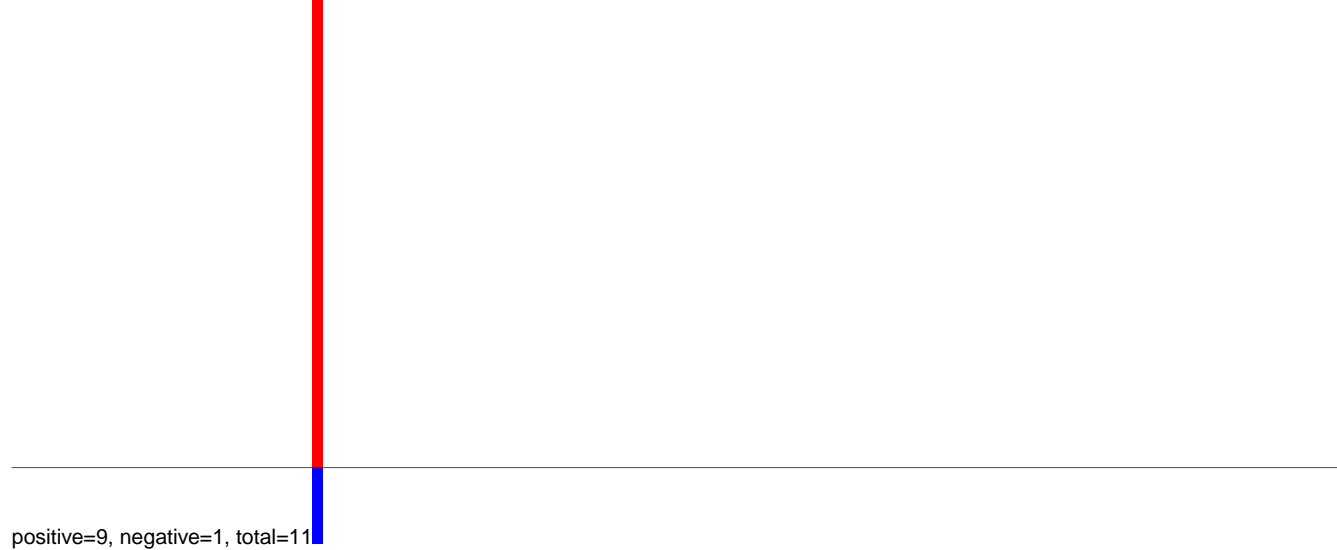


positive=0, negative=13, total=13

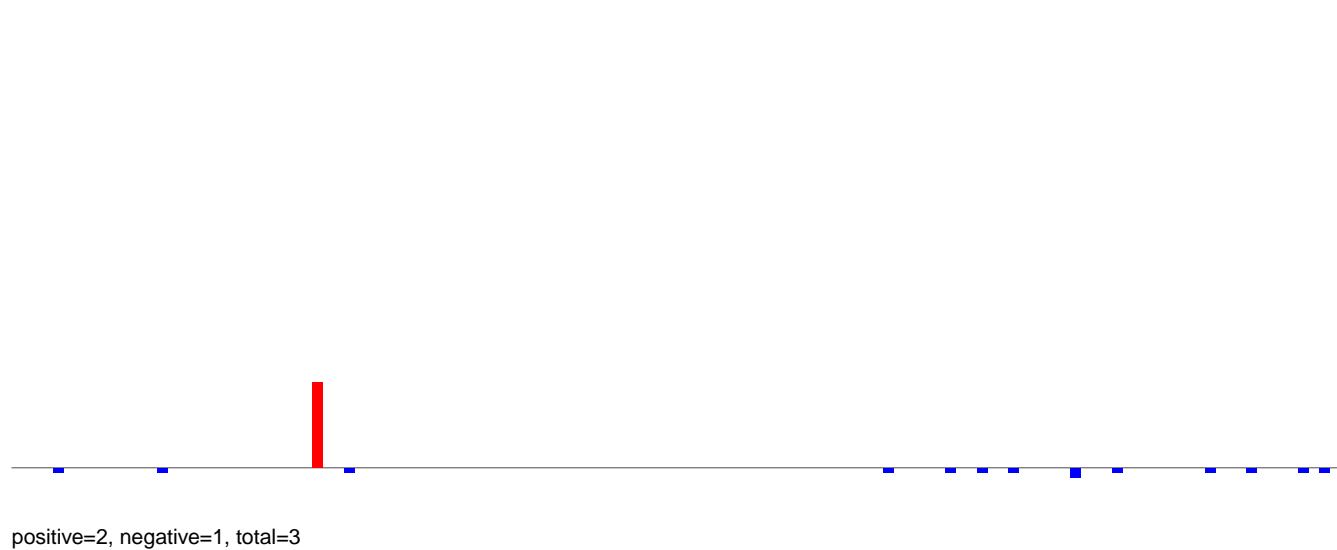
Window size=50, length=762, TE@HATN4, AG:1-762

200 400 600 800

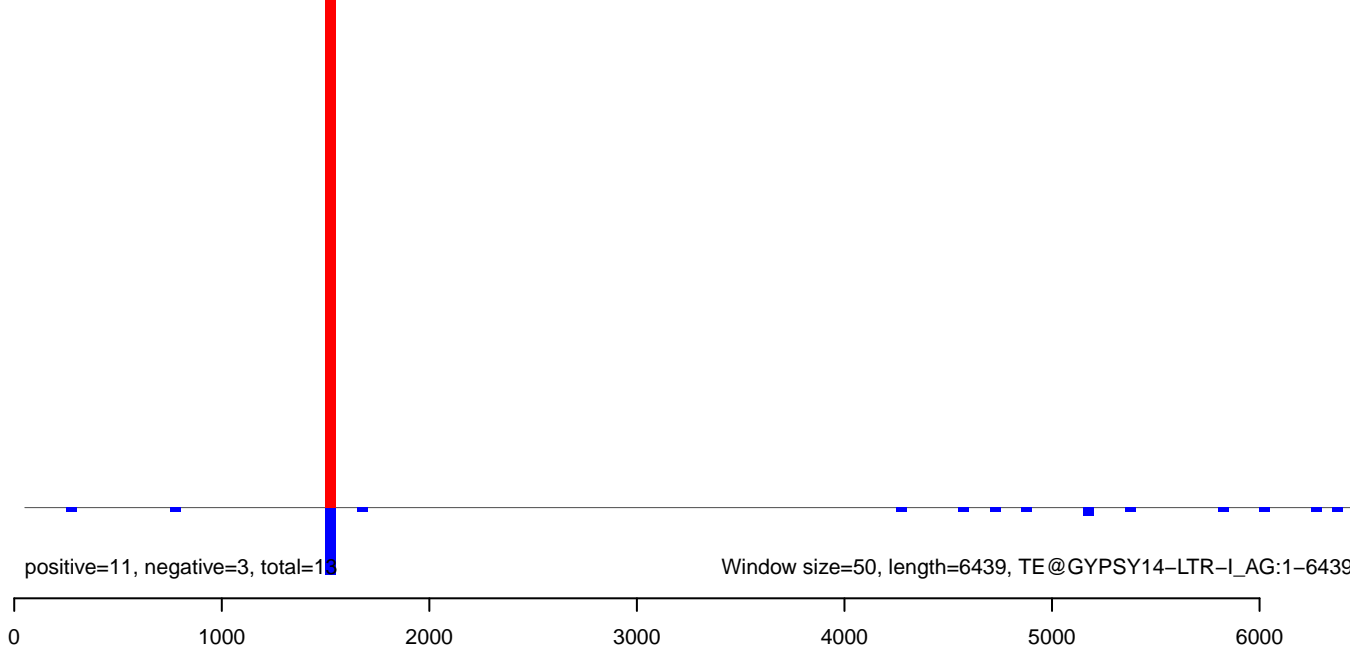
AnGam_Mos55_cells.18_23.rep



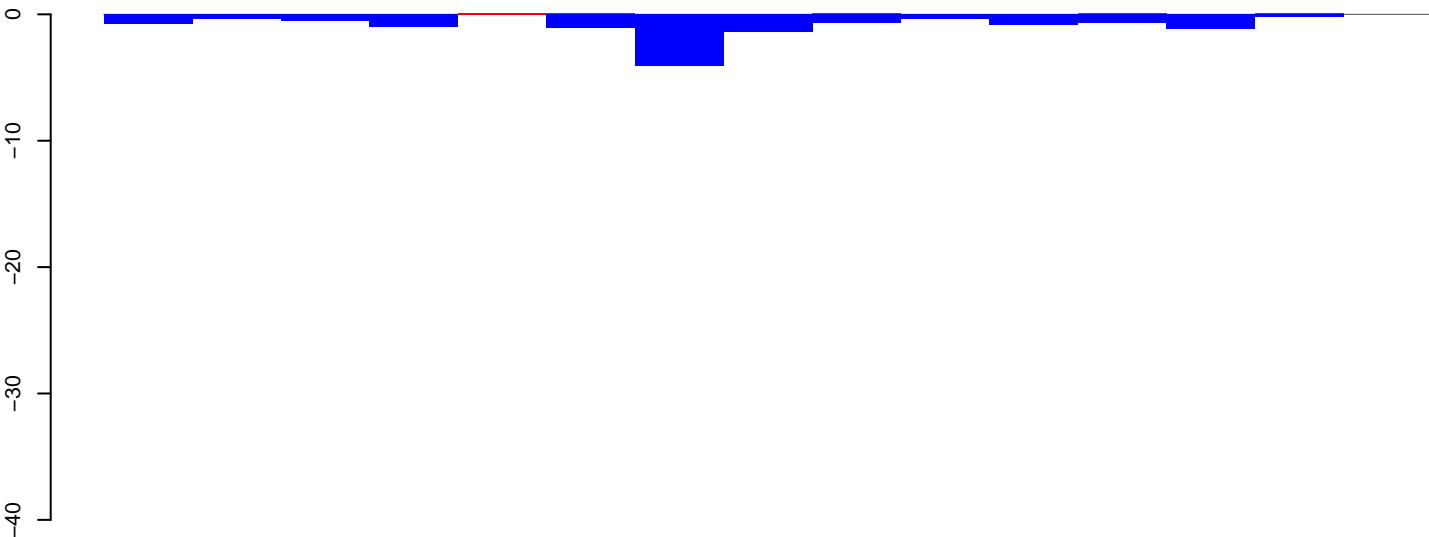
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

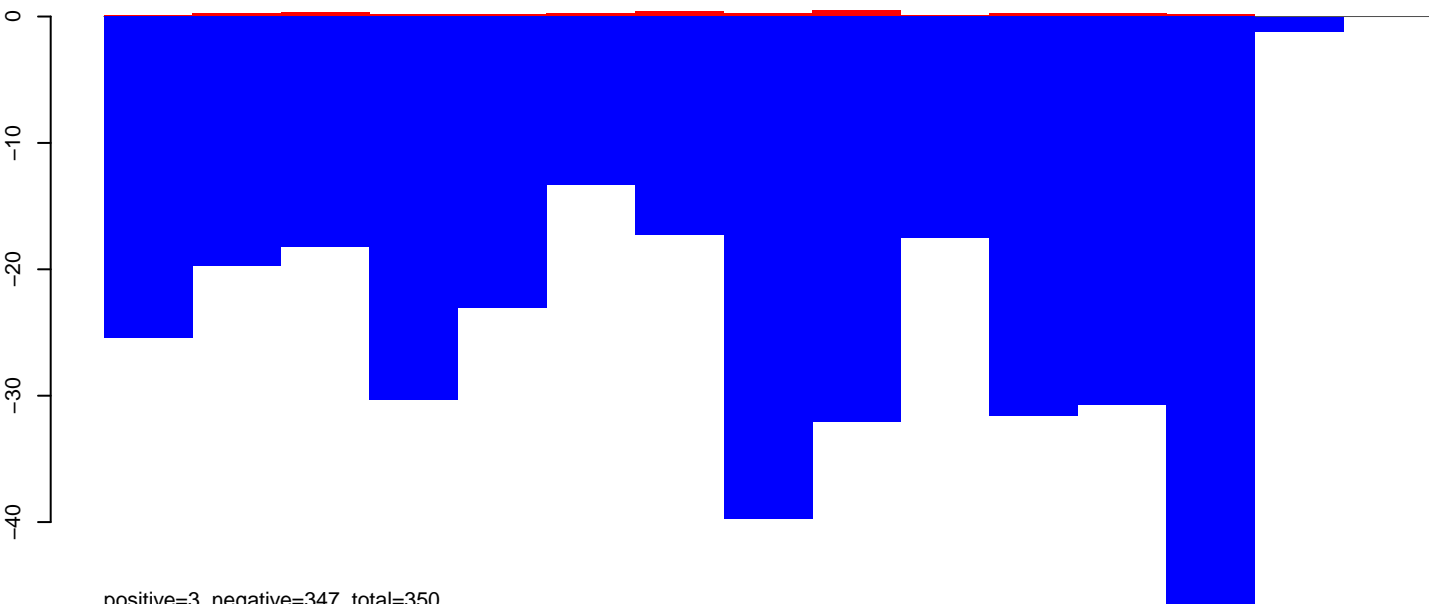


AnGam_Mos55_cells.18_23.rep



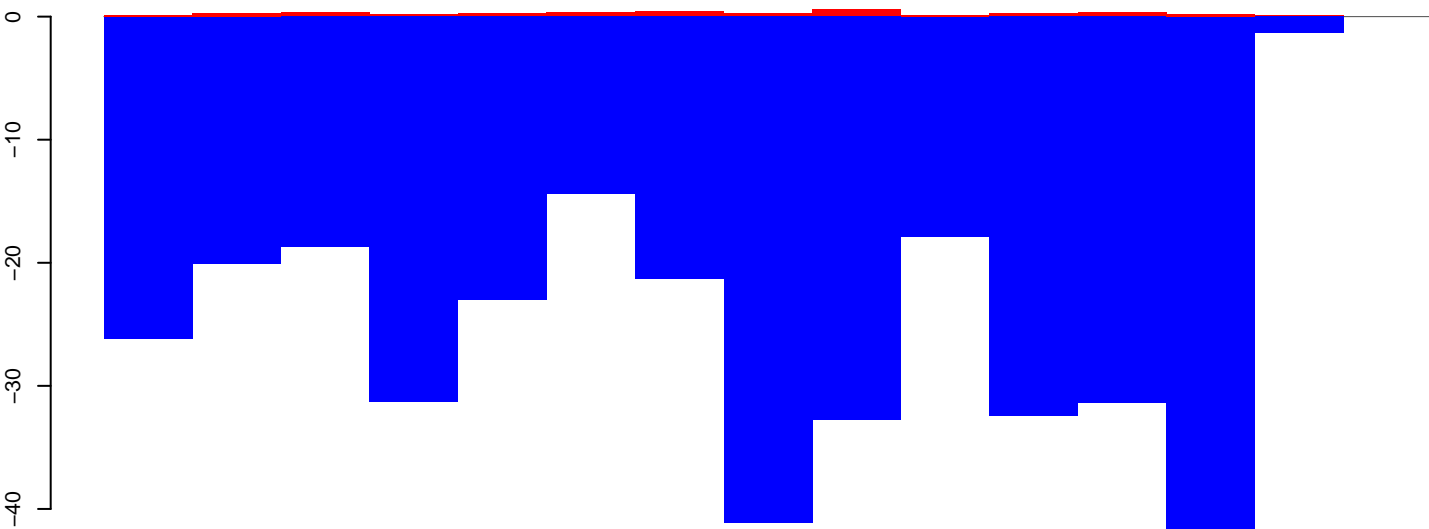
positive=0, negative=13, total=13

AnGam_Mos55_cells.24_35.rep



positive=3, negative=347, total=350

AnGam_Mos55_cells.rep

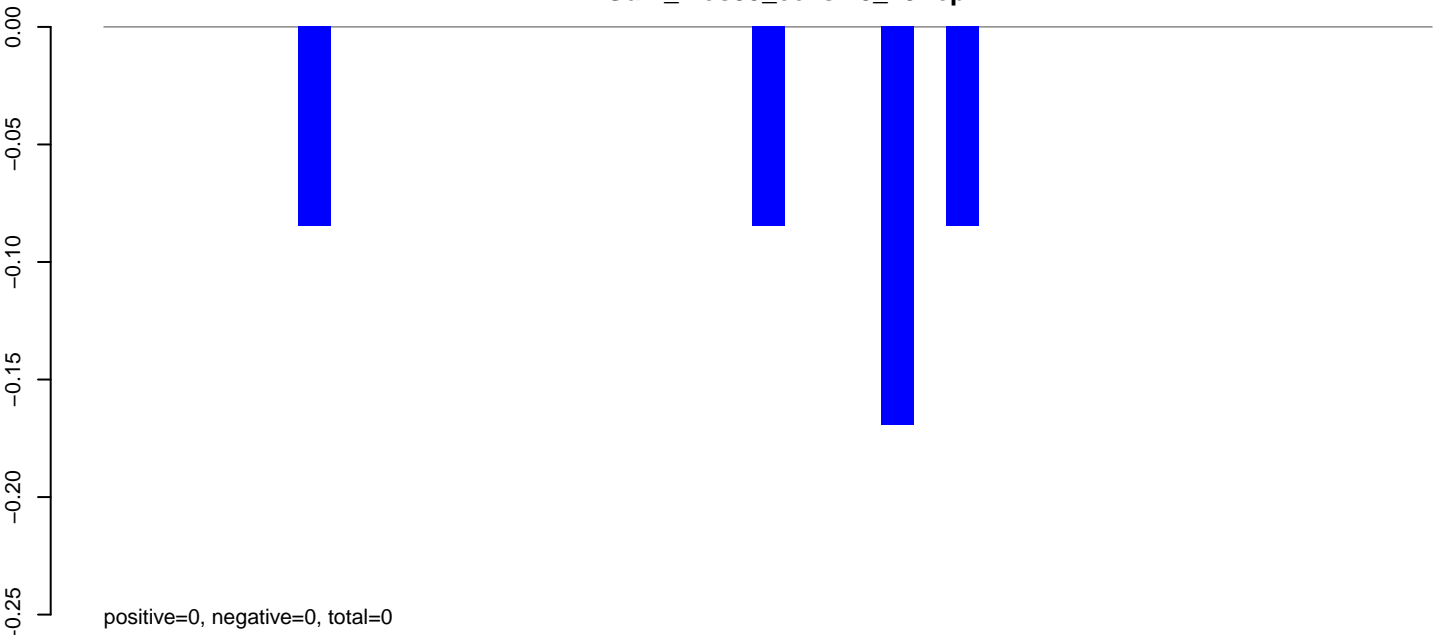


positive=4, negative=360, total=364

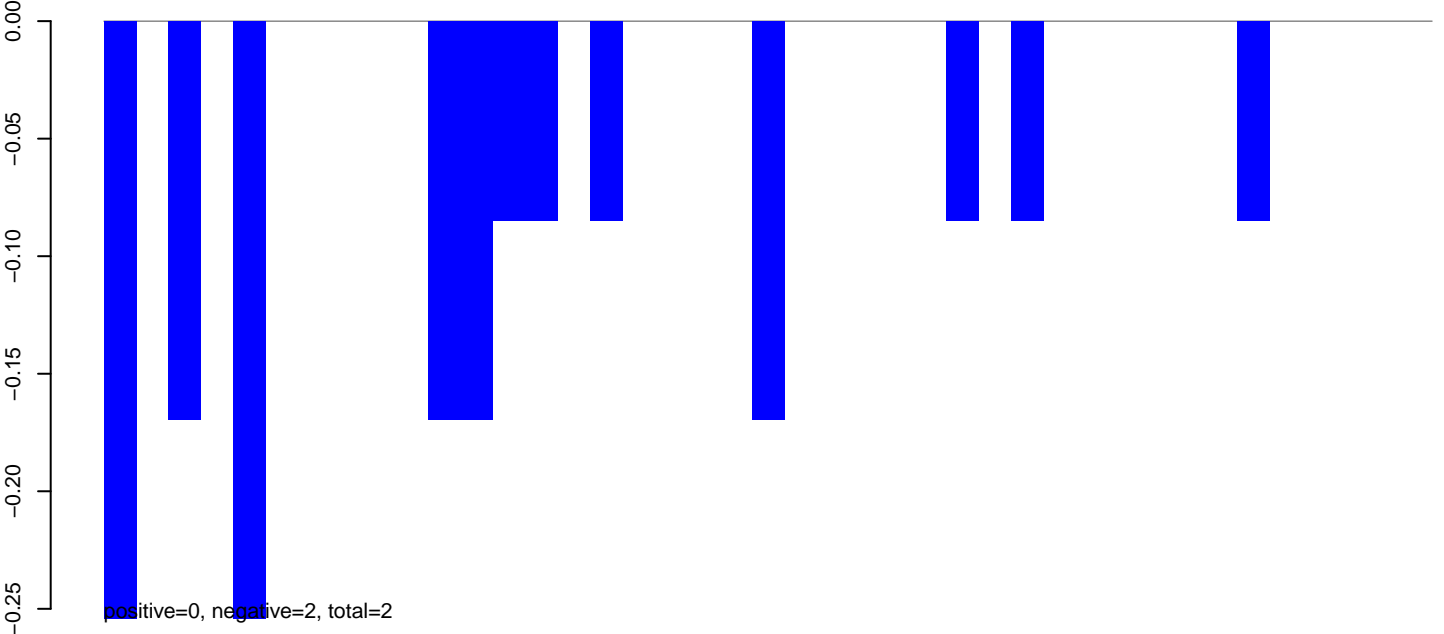
Window size=50, length=756, TE@EAA44976-Unknown:1-756

200 400 600 800

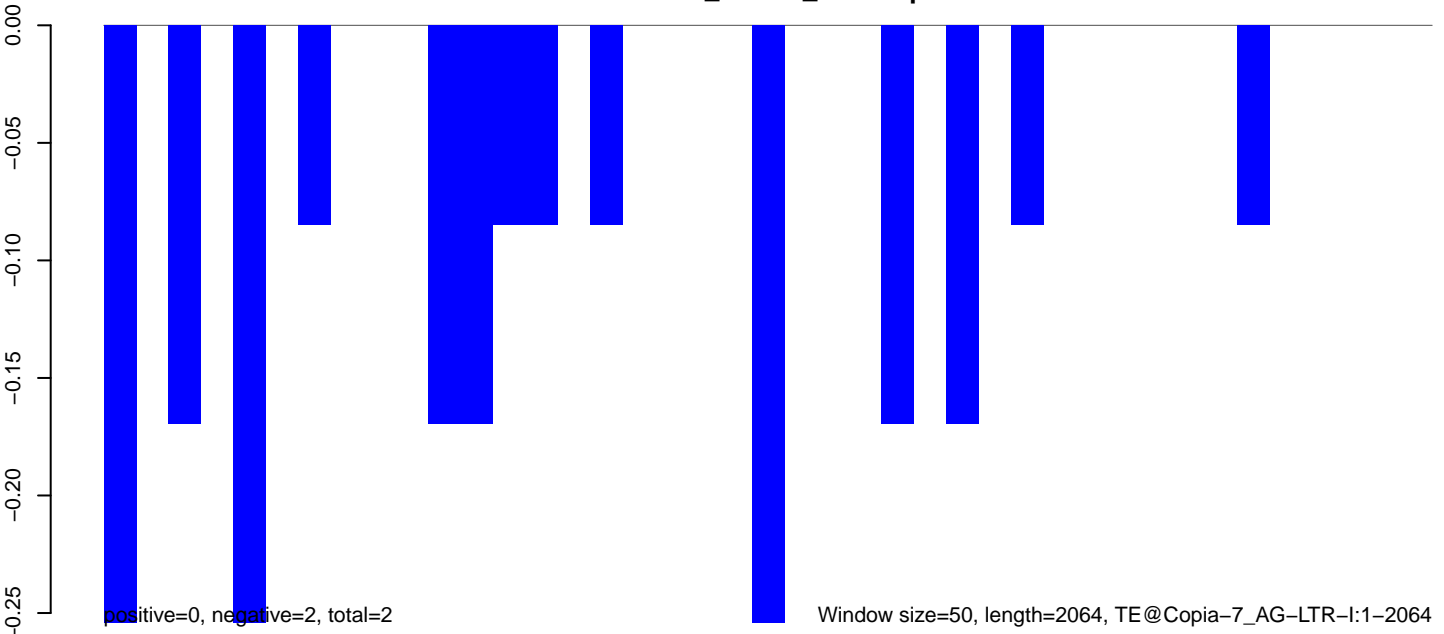
AnGam_Mos55_cells.18_23.rep



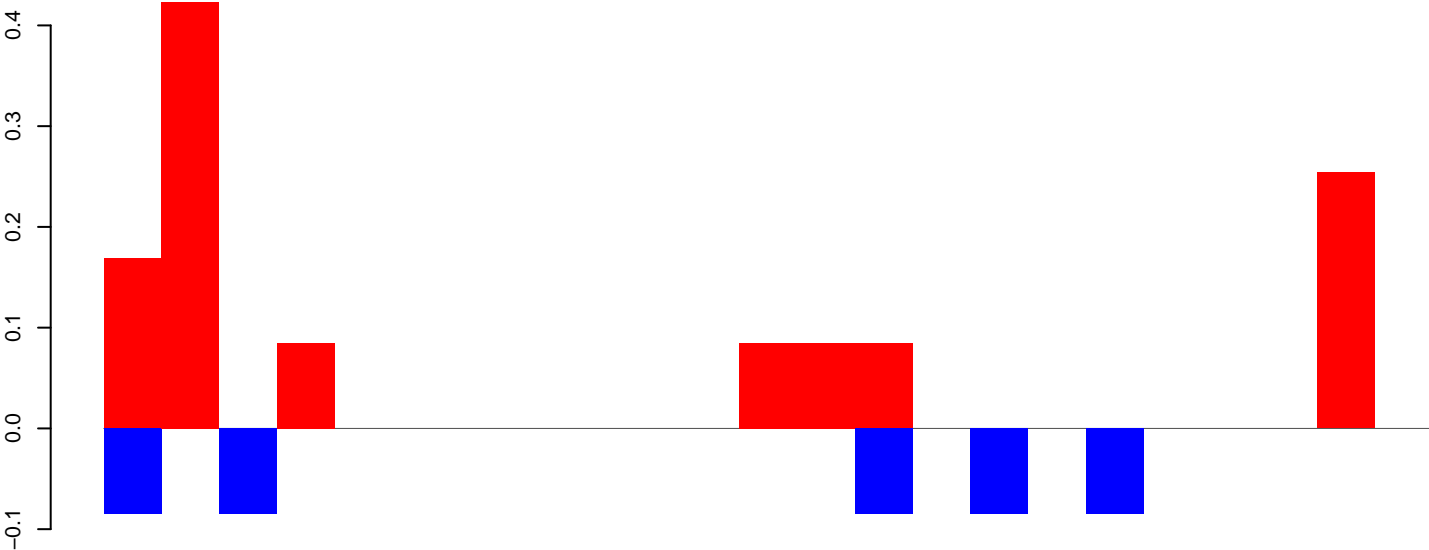
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

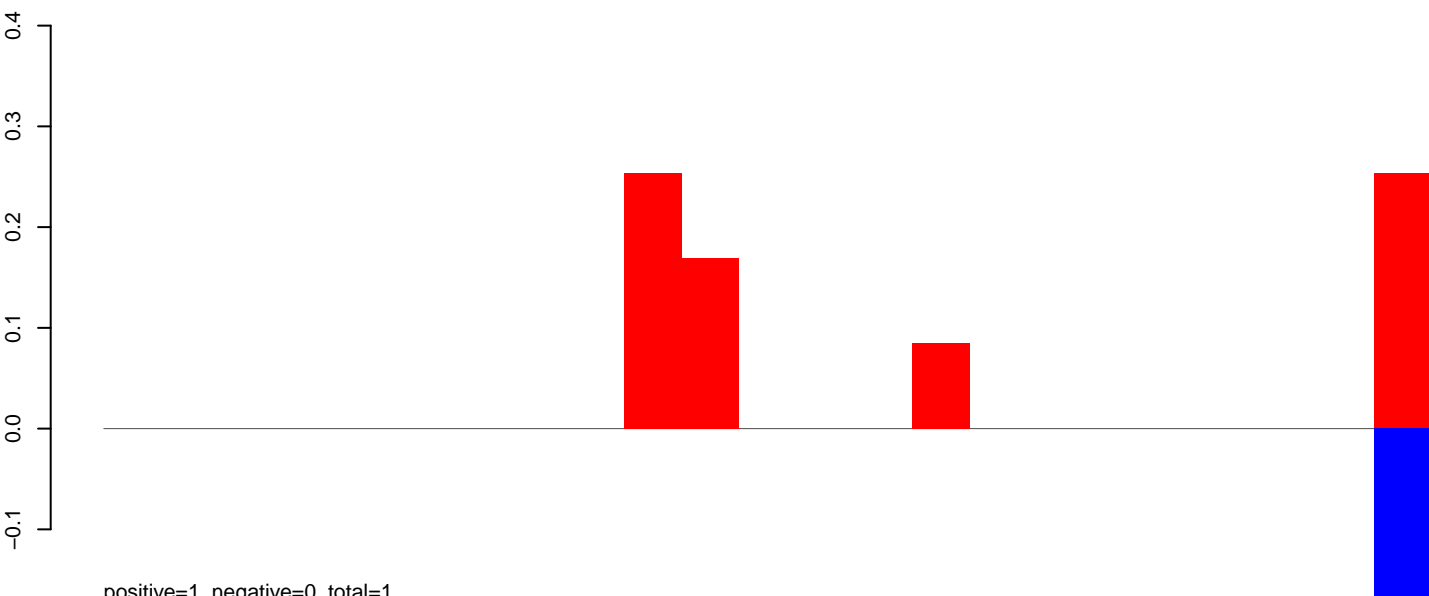


AnGam_Mos55_cells.18_23.rep



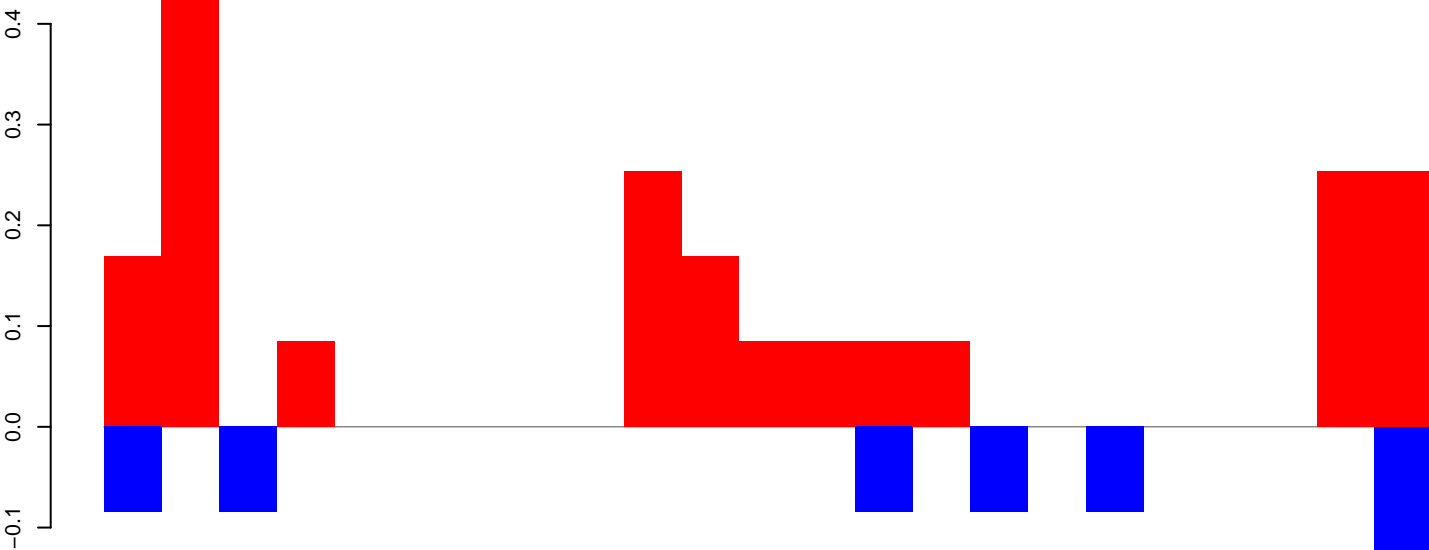
positive=1, negative=0, total=2

AnGam_Mos55_cells.24_35.rep



positive=1, negative=0, total=1

AnGam_Mos55_cells.rep



positive=2, negative=1, total=3

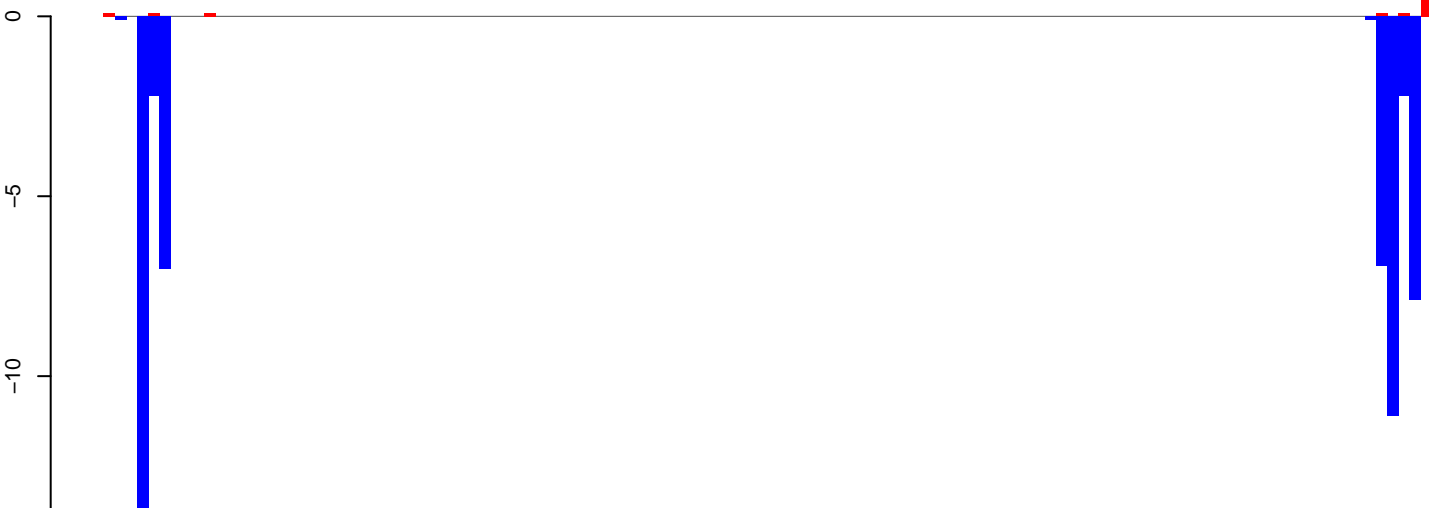
Window size=50, length=1175, TE@Clu-137_AG:1-1175

AnGam_Mos55_cells.18_23.rep



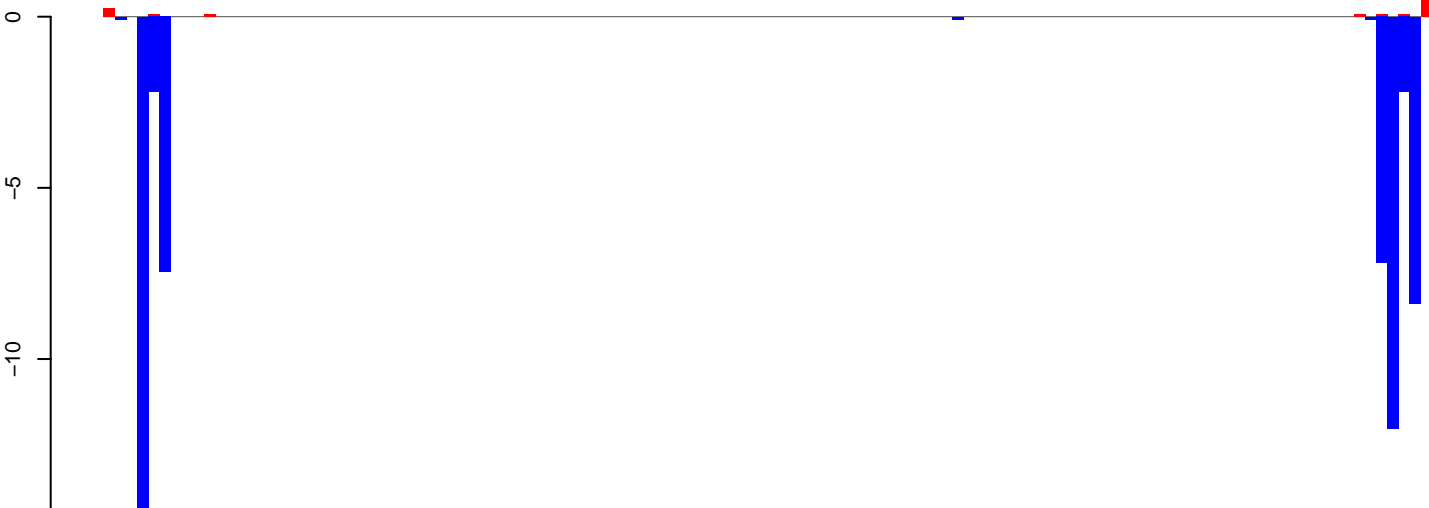
positive=0, negative=3, total=3

AnGam_Mos55_cells.24_35.rep



positive=5, negative=52, total=57

AnGam_Mos55_cells.rep

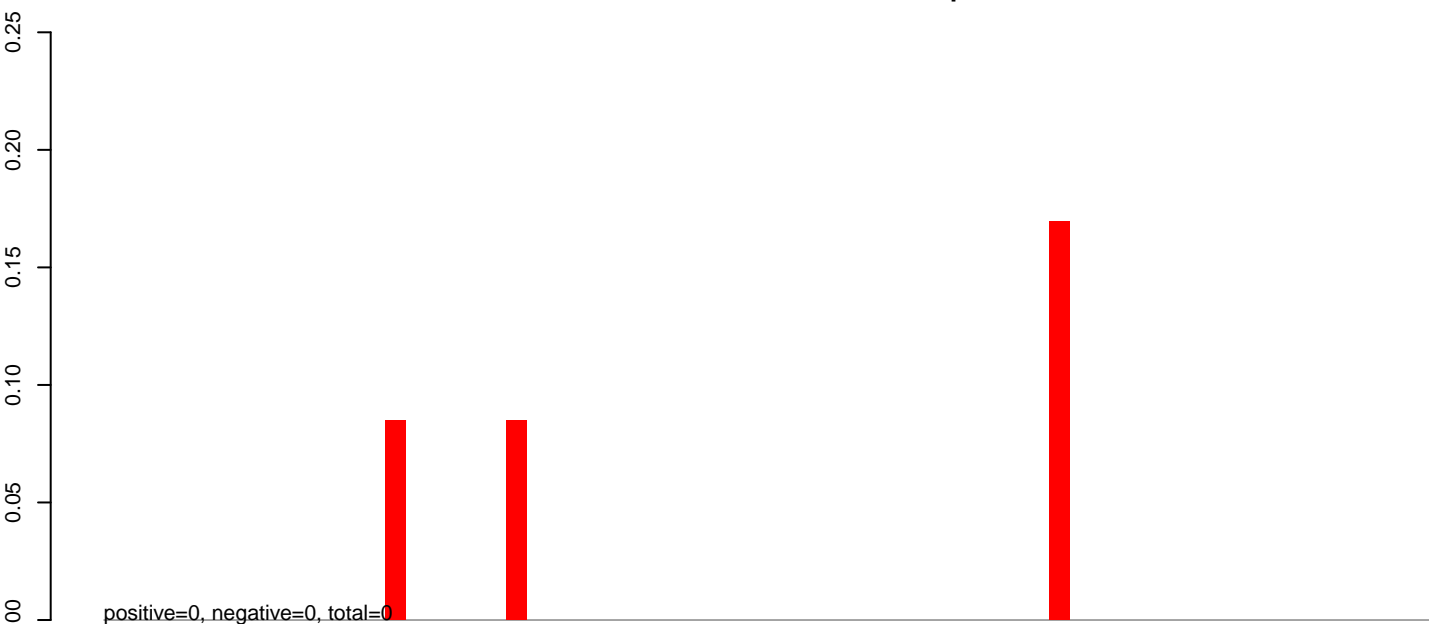


positive=5, negative=55, total=60

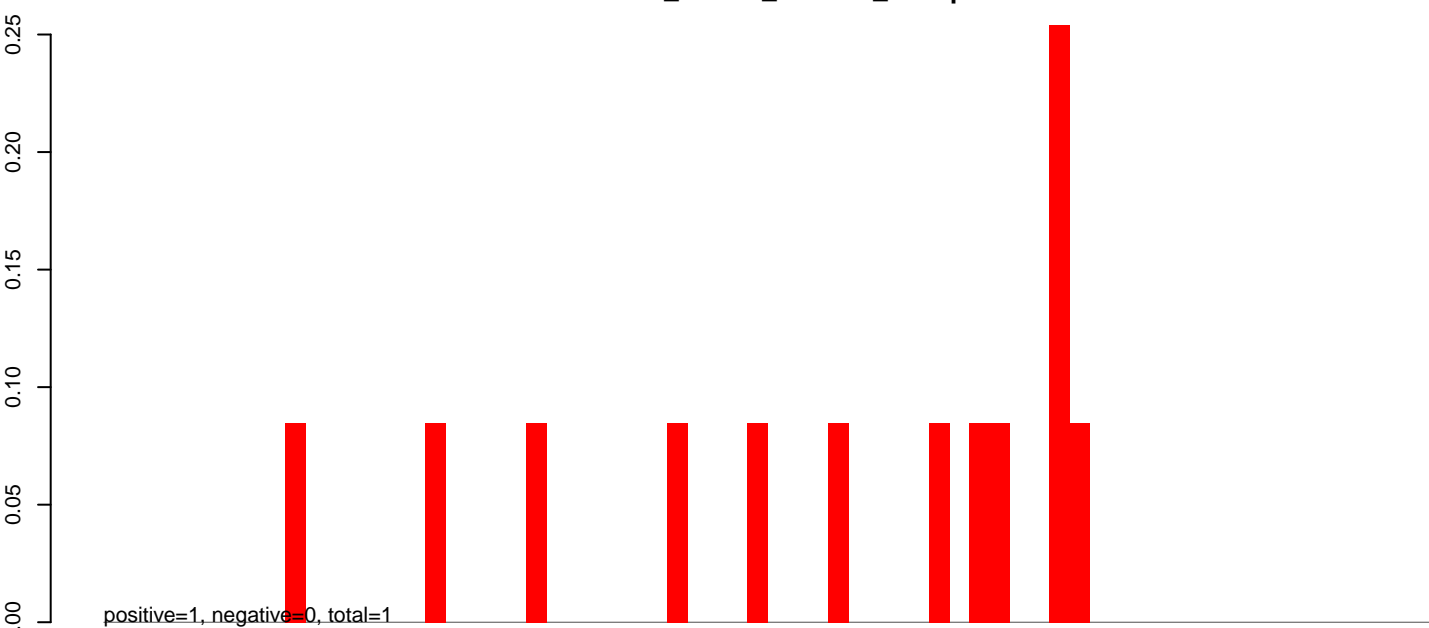
Window size=50, length=5983, TE@AGM1:1-5983

0 1000 2000 3000 4000 5000 6000

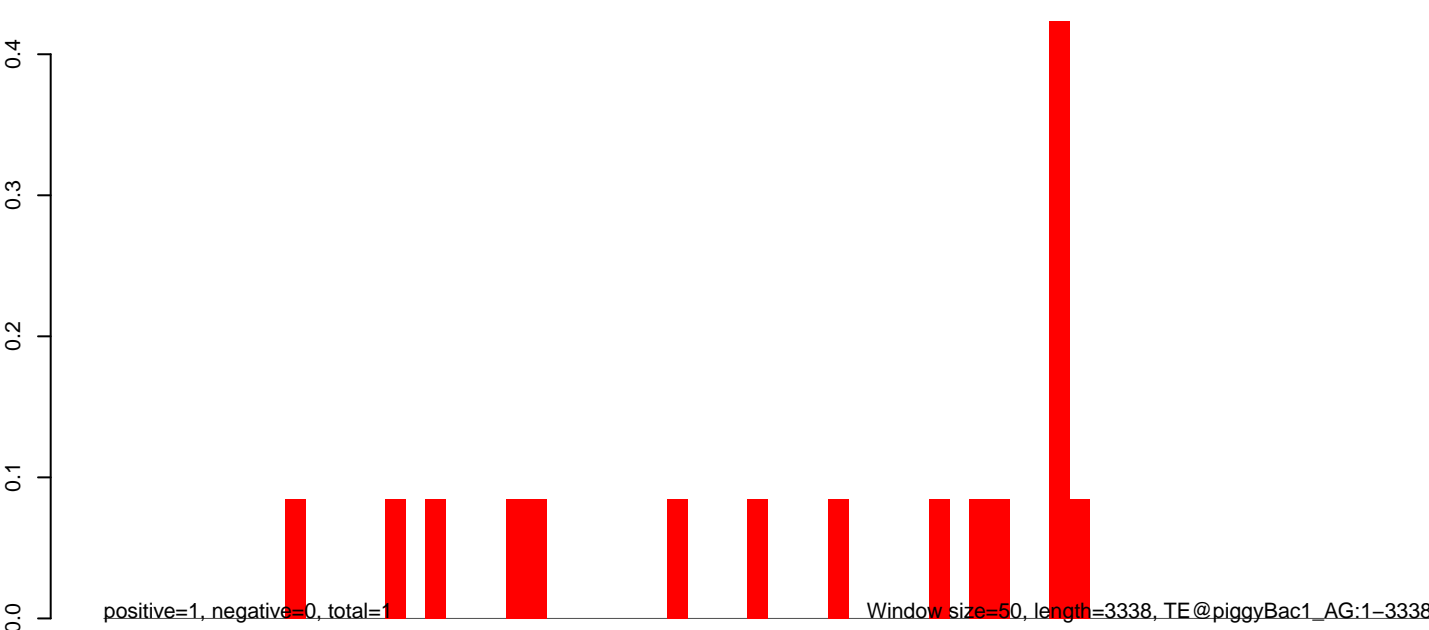
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

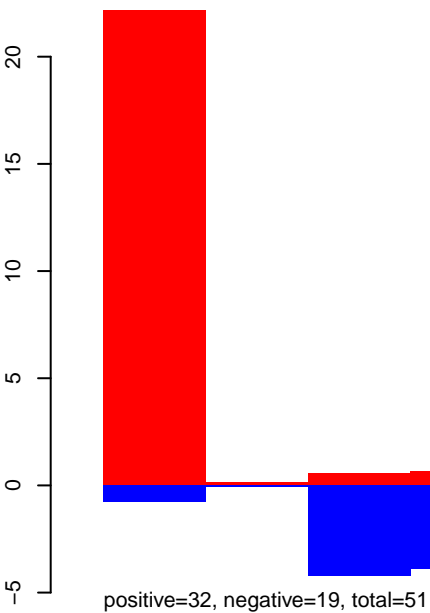


AnGam_Mos55_cells.rep

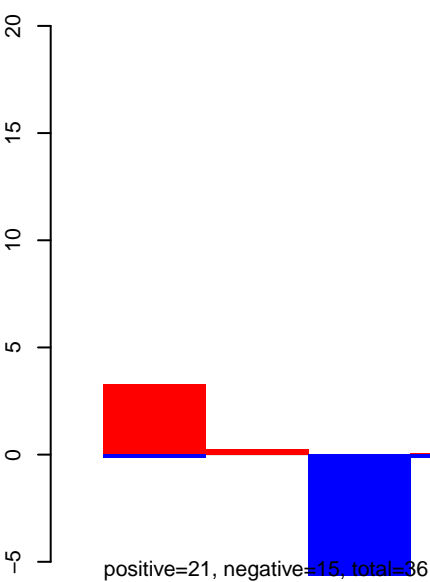


0 500 1000 1500 2000 2500 3000

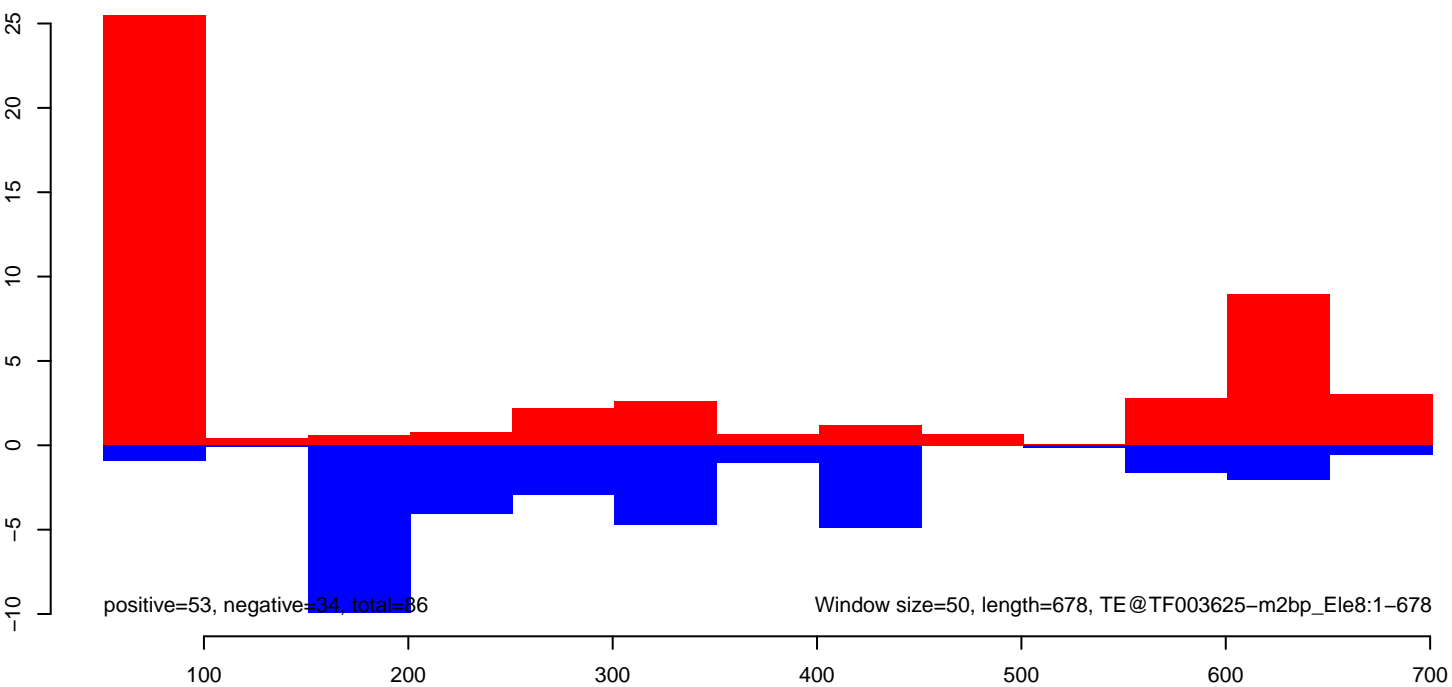
AnGam_Mos55_cells.18_23.rep



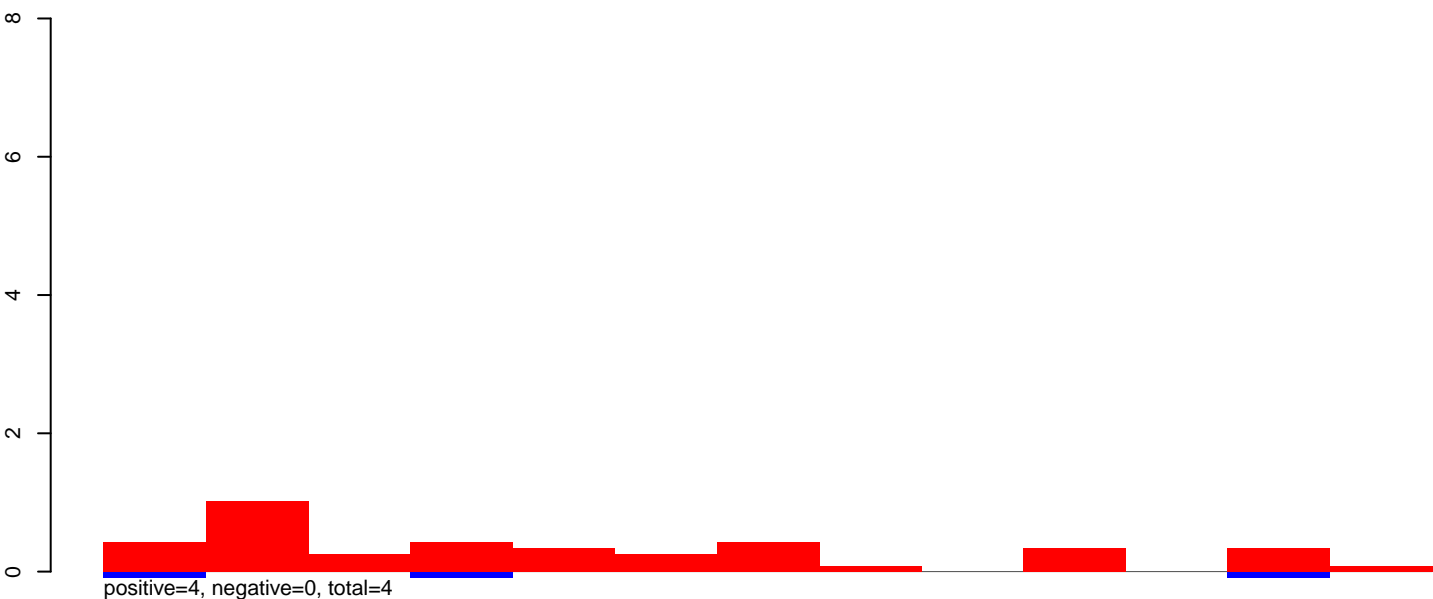
AnGam_Mos55_cells.24_35.rep



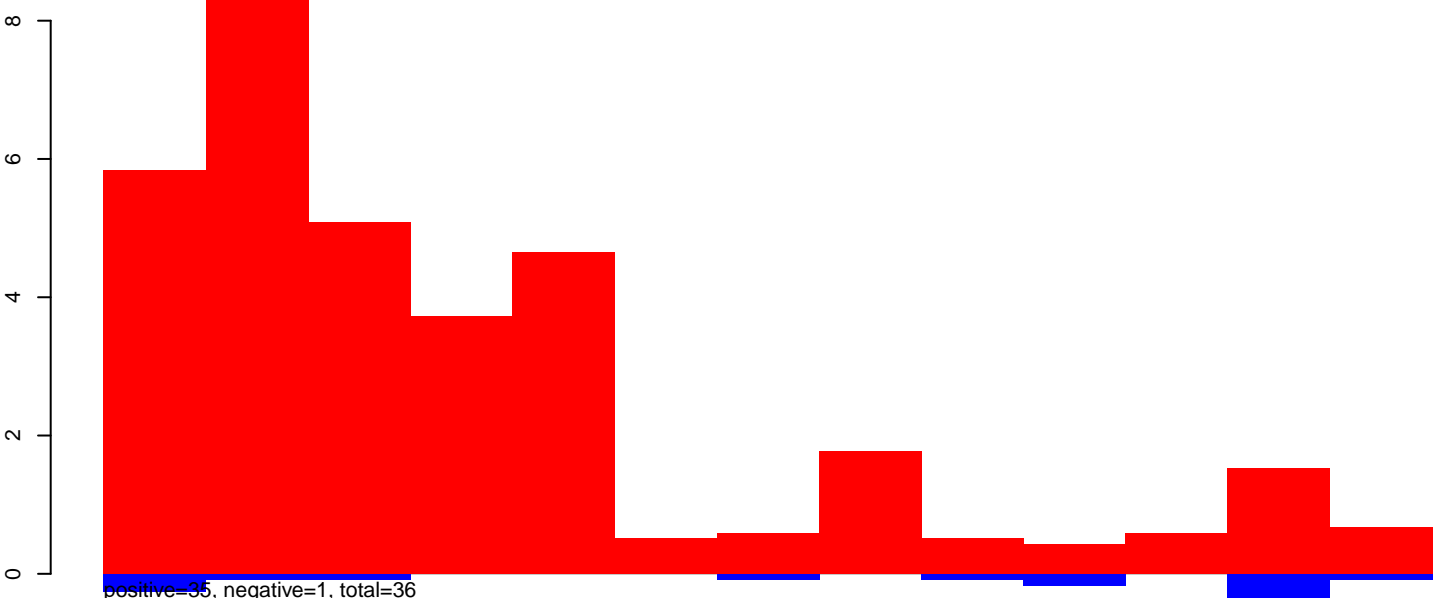
AnGam_Mos55_cells.rep



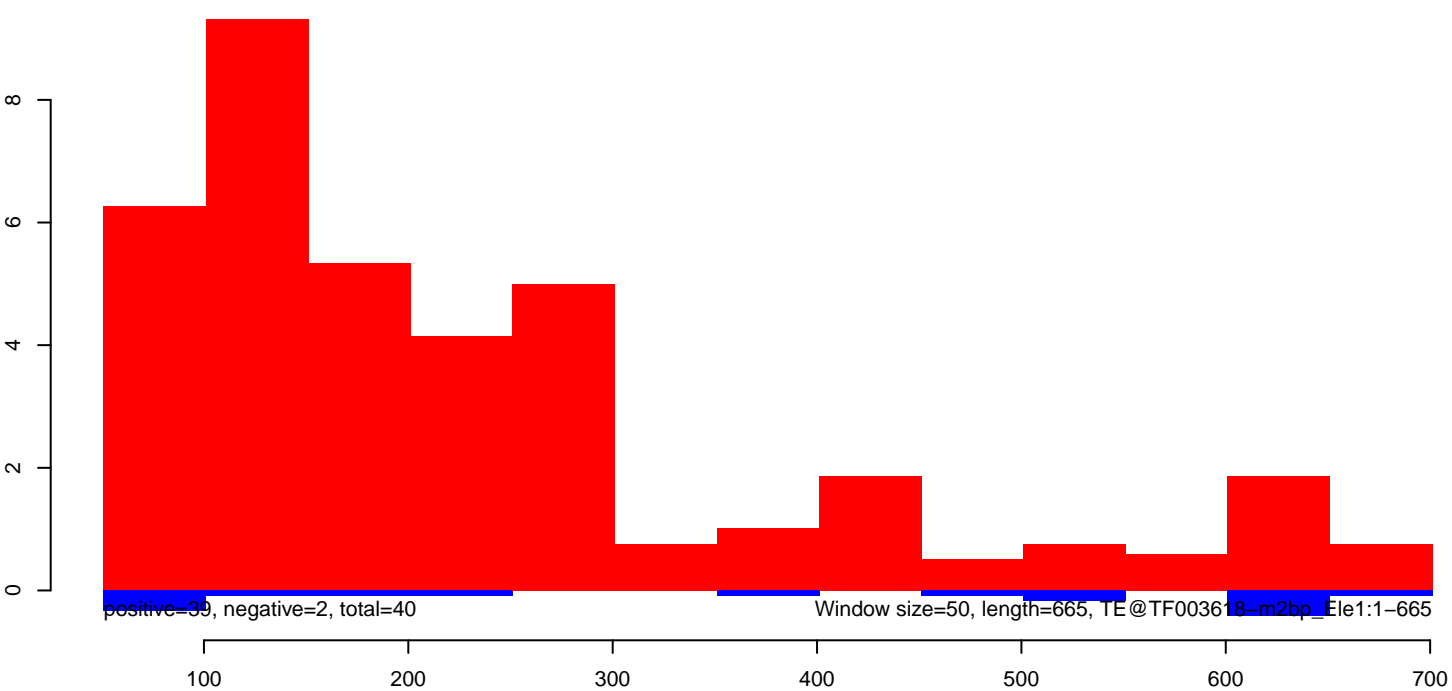
AnGam_Mos55_cells.18_23.rep



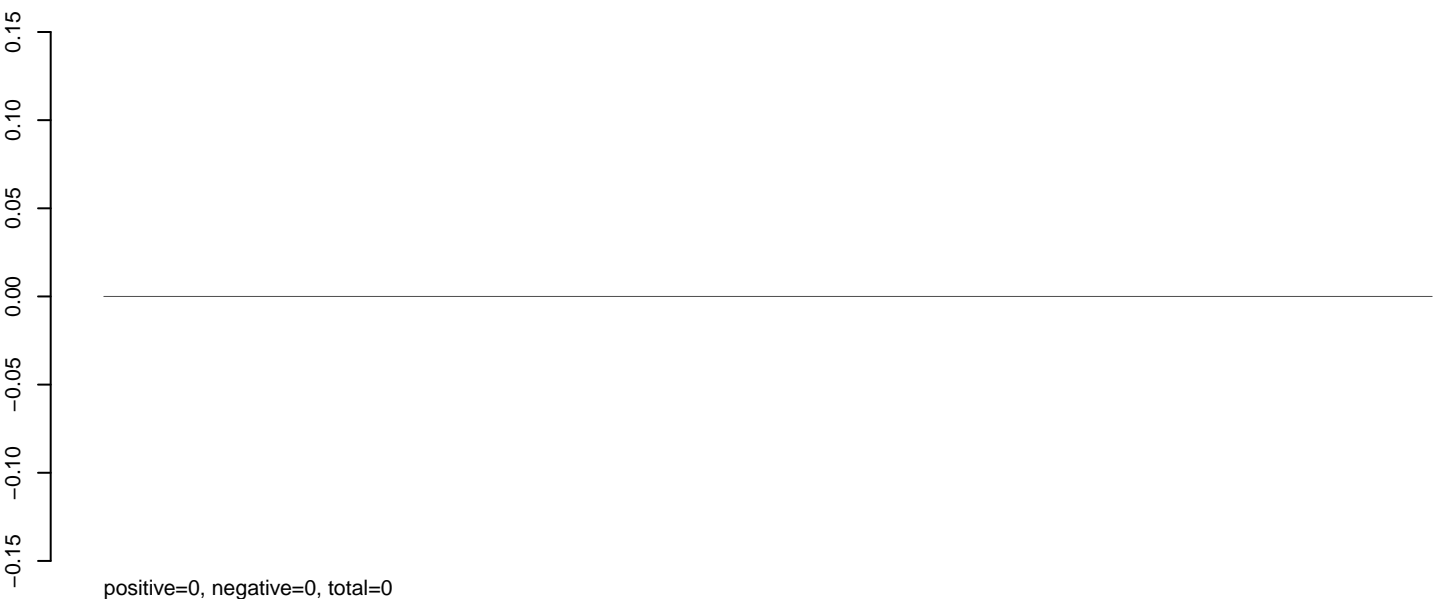
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



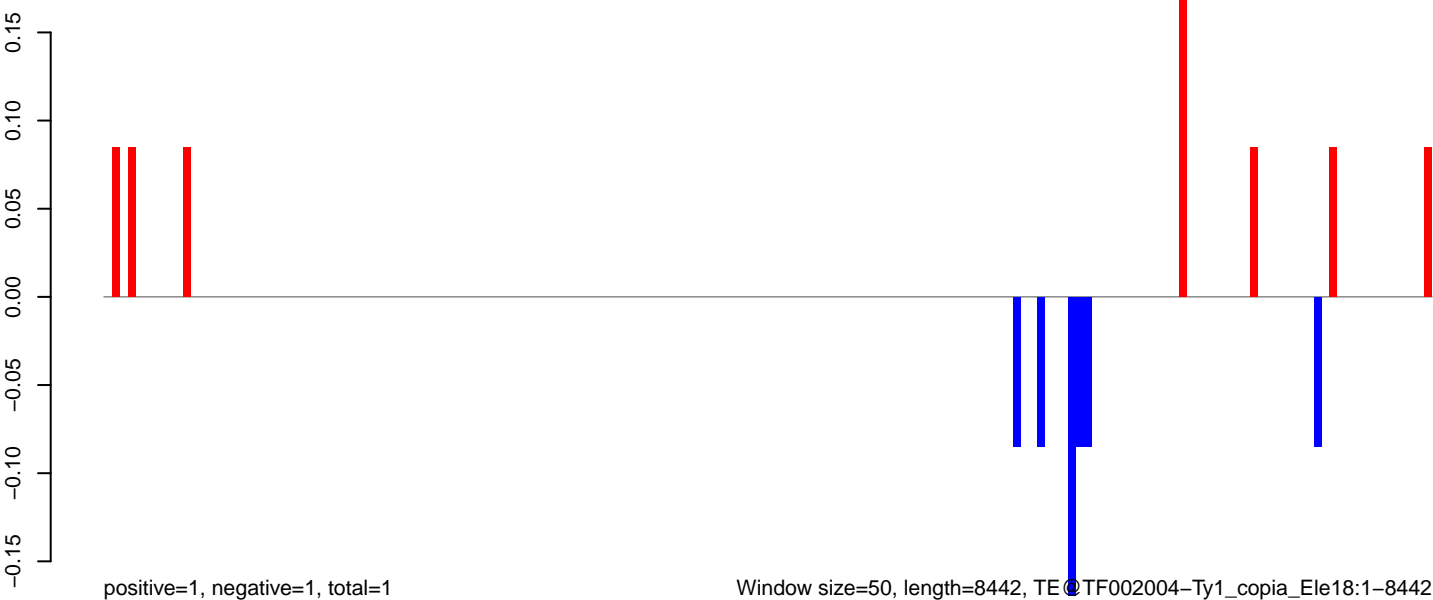
AnGam_Mos55_cells.18_23.rep



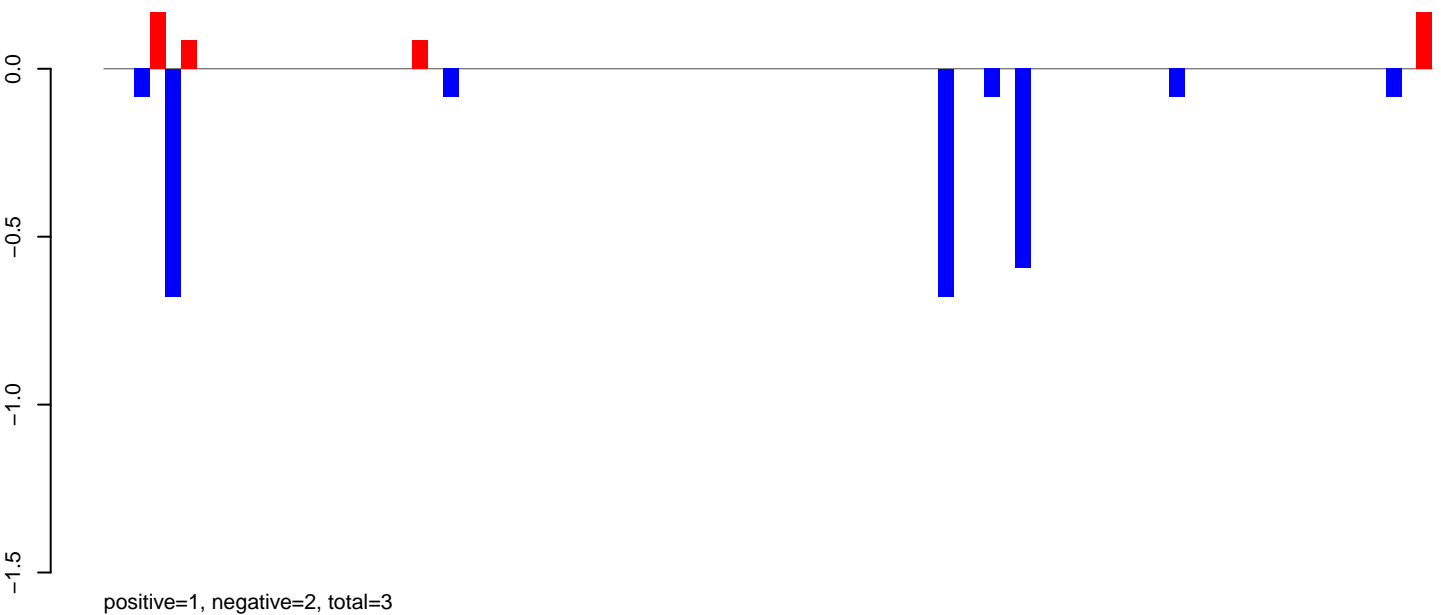
AnGam_Mos55_cells.24_35.rep



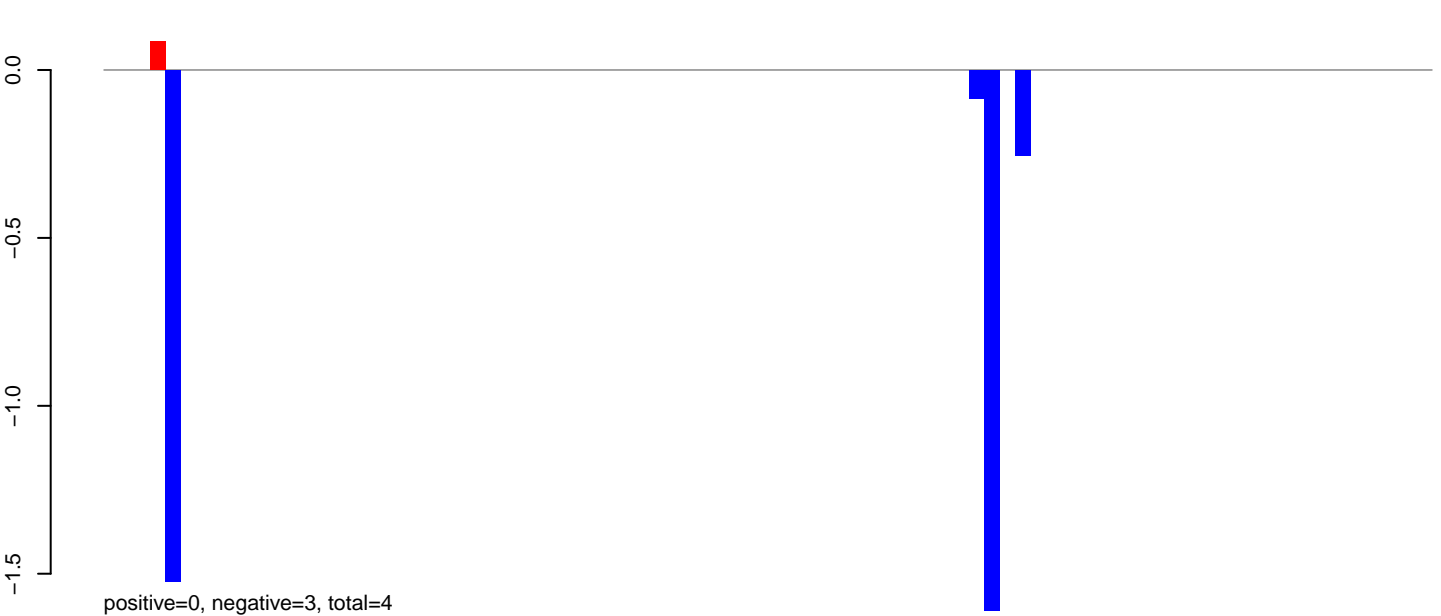
AnGam_Mos55_cells.rep



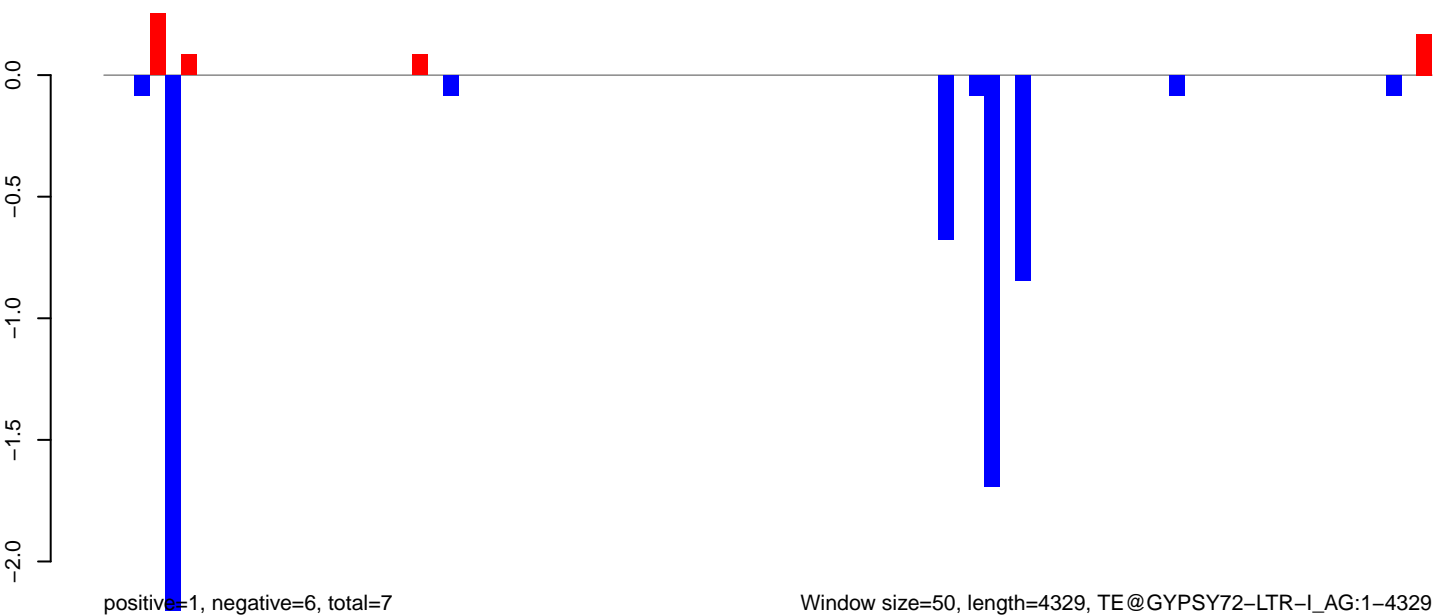
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

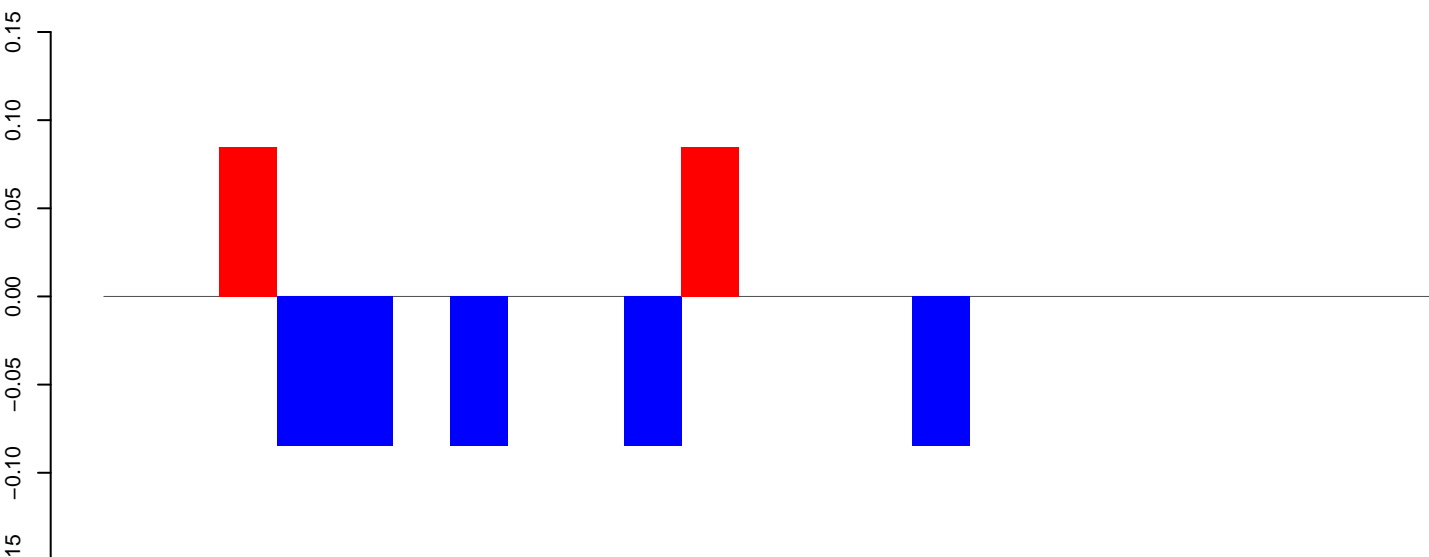


AnGam_Mos55_cells.rep



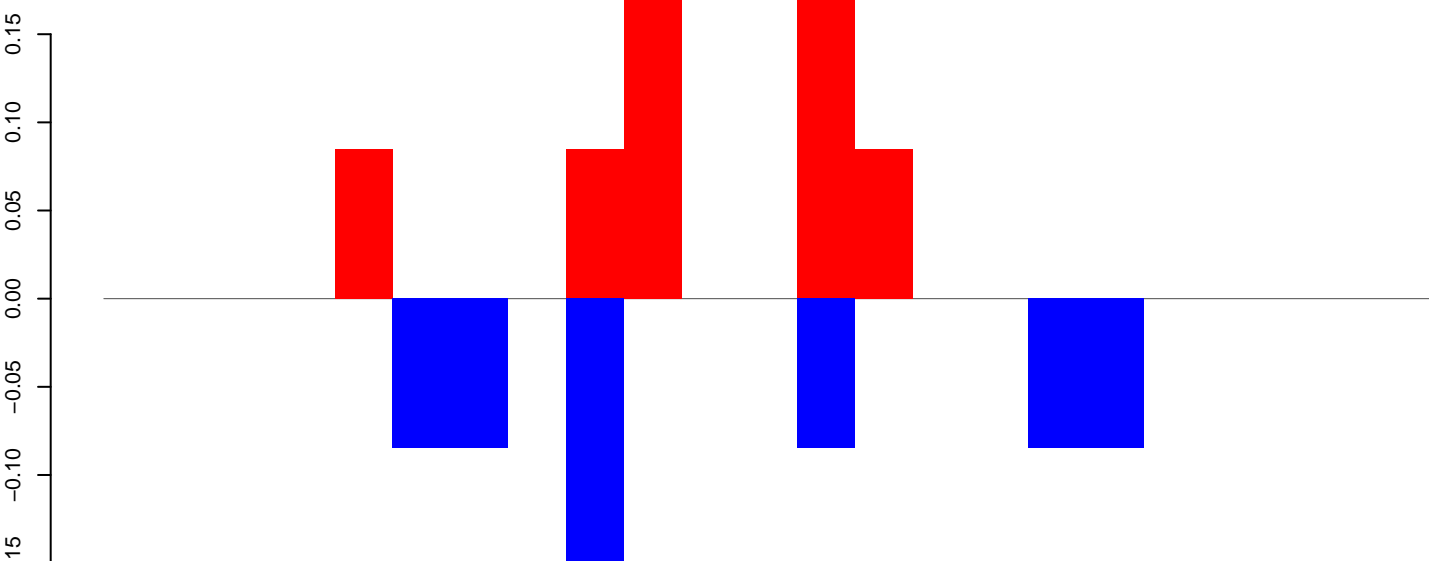
Window size=50, length=4329, TE@GYPSY72-LTR-L_AG:1-4329

AnGam_Mos55_cells.18_23.rep



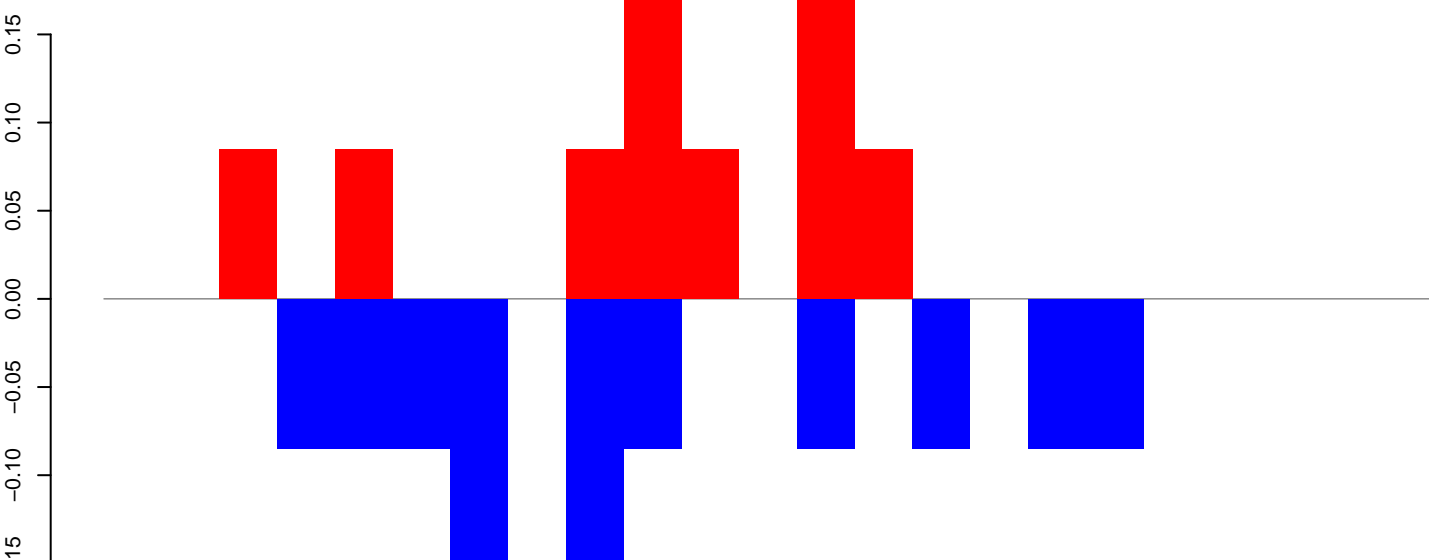
positive=0, negative=0, total=1

AnGam_Mos55_cells.24_35.rep



positive=1, negative=1, total=1

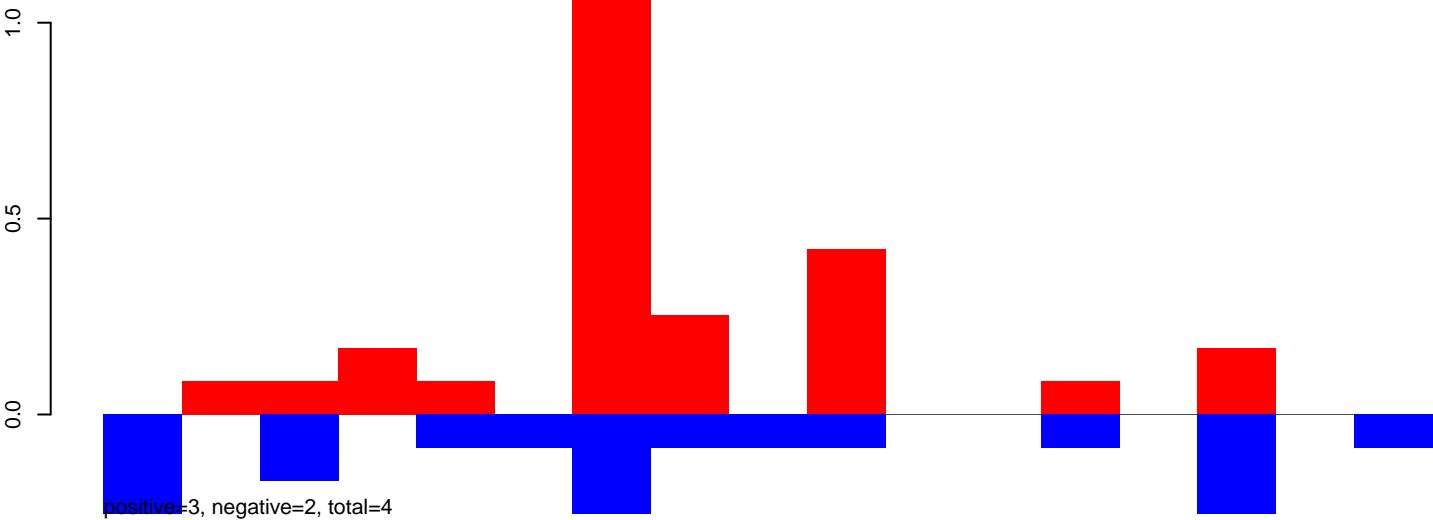
AnGam_Mos55_cells.rep



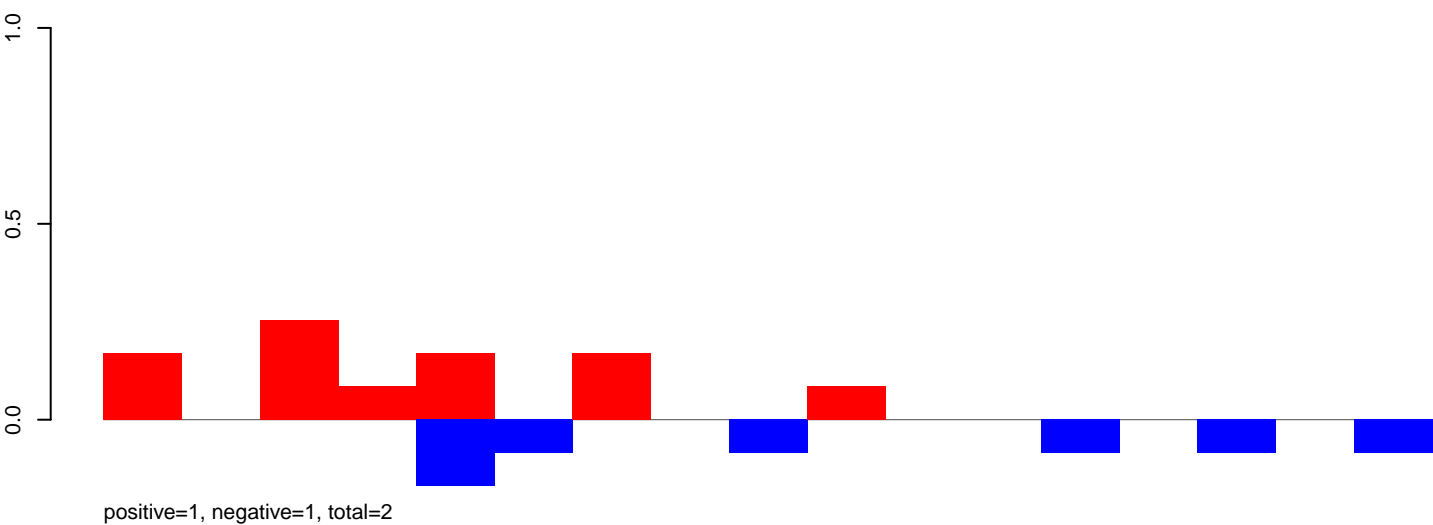
positive=1, negative=1, total=2

Window size=50, length=1172, TE@EnSpm-N1_AG:1-1172

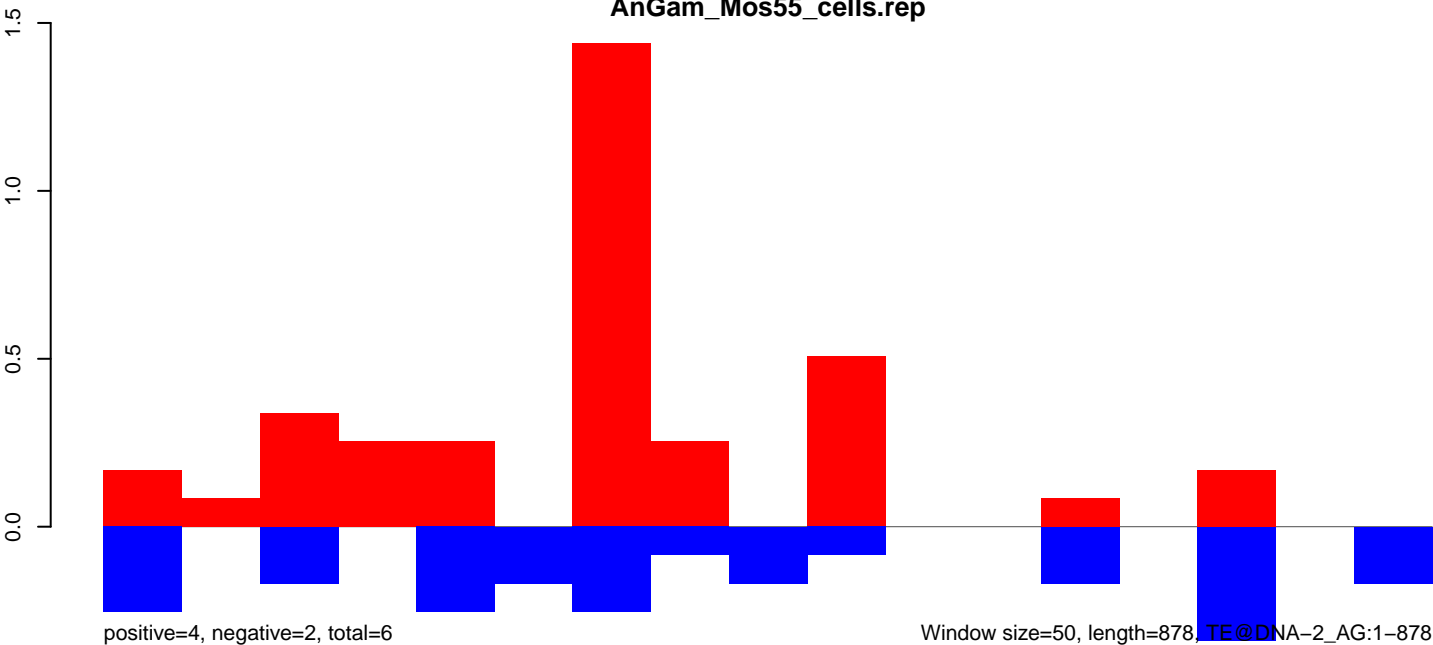
AnGam_Mos55_cells.18_23.rep



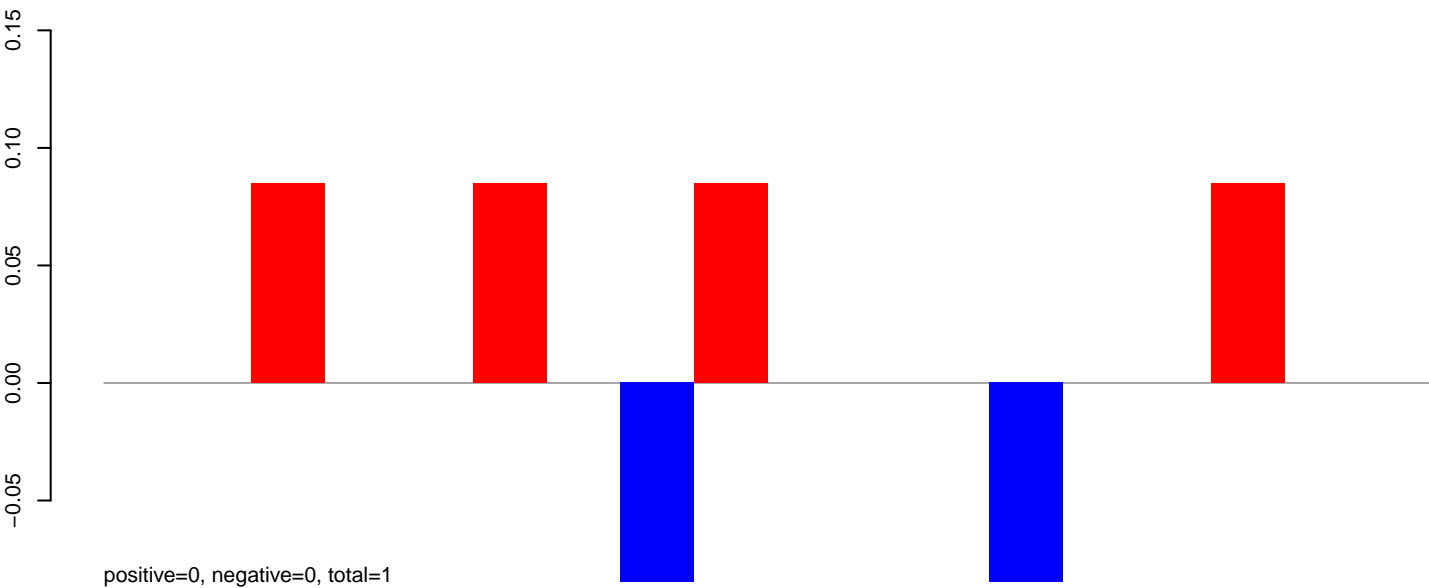
AnGam_Mos55_cells.24_35.rep



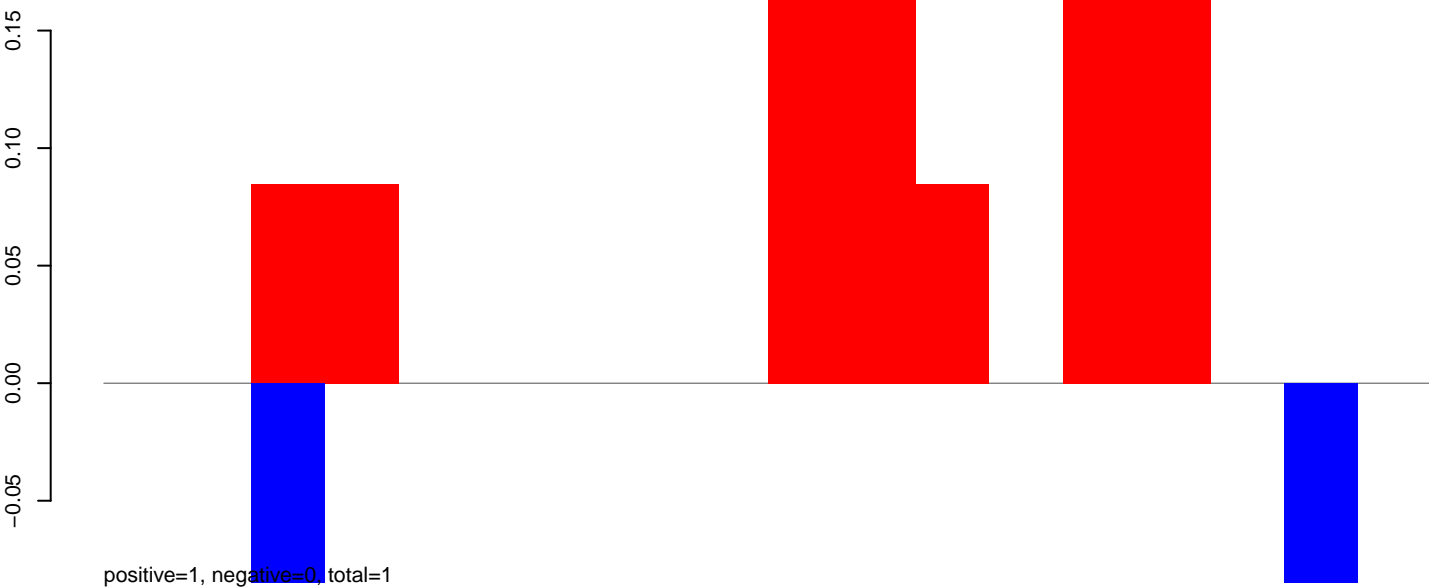
AnGam_Mos55_cells.rep



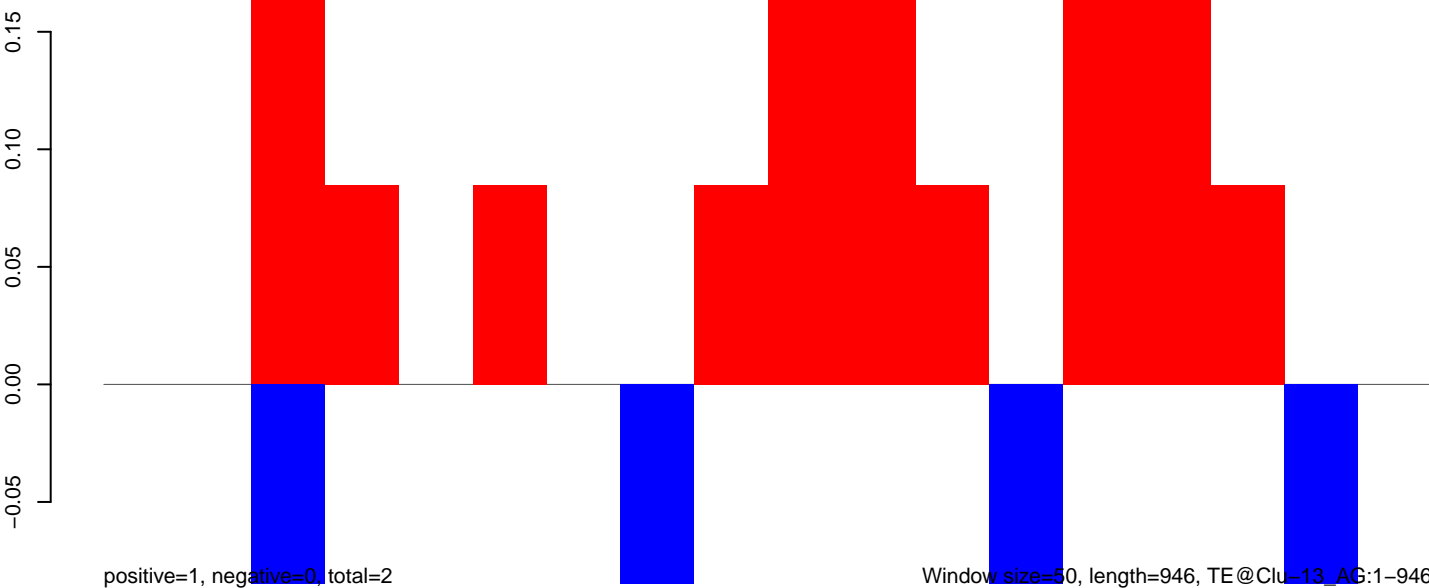
AnGam_Mos55_cells.18_23.rep



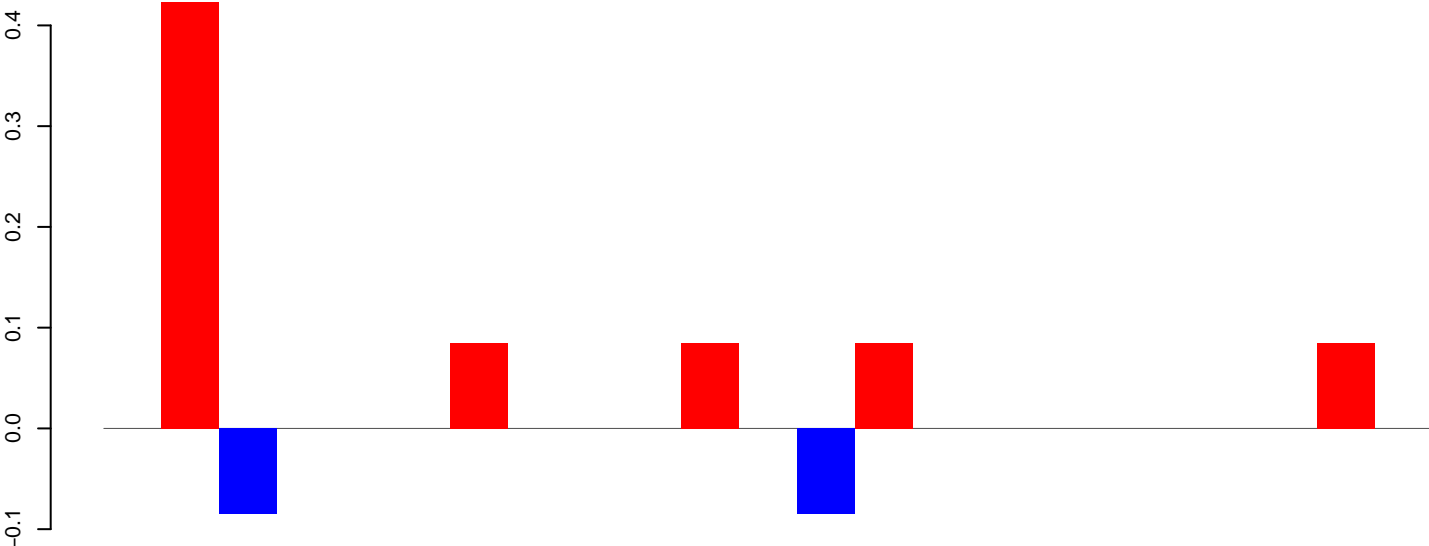
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

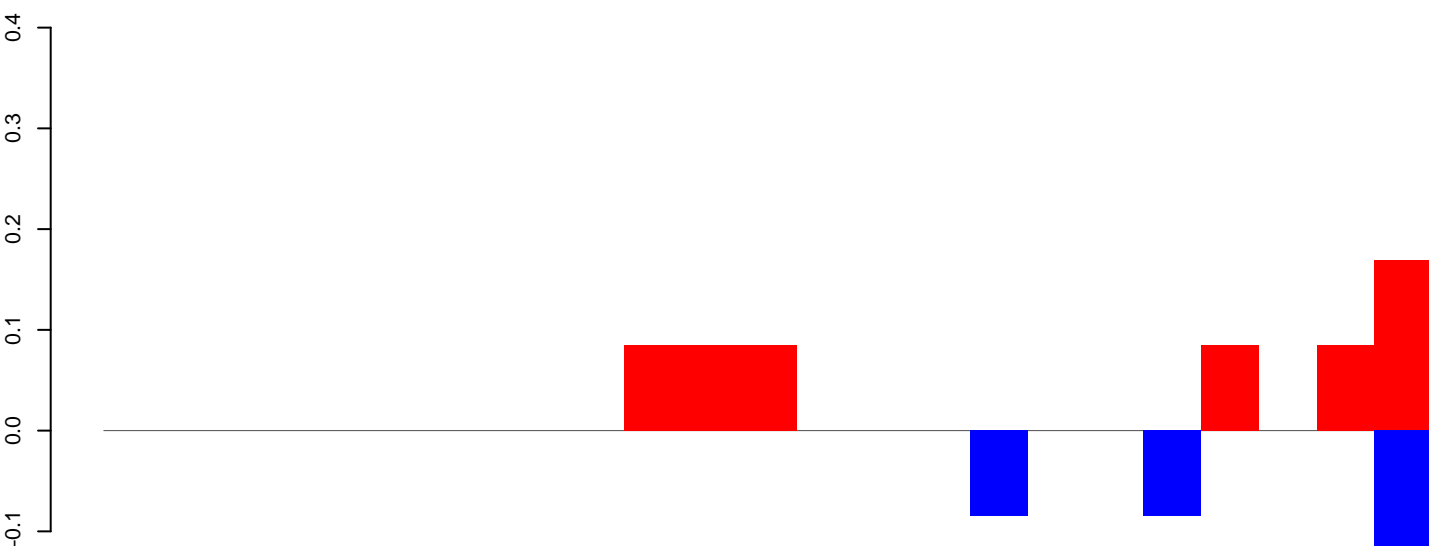


AnGam_Mos55_cells.18_23.rep



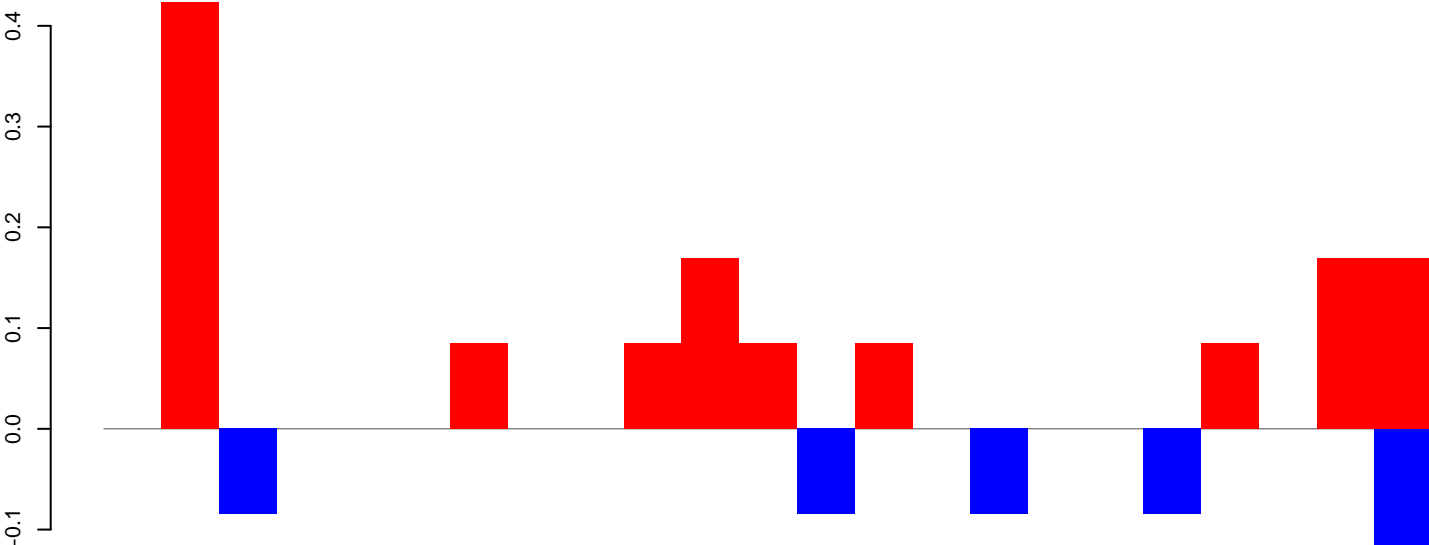
positive=1, negative=0, total=1

AnGam_Mos55_cells.24_35.rep



positive=1, negative=1, total=1

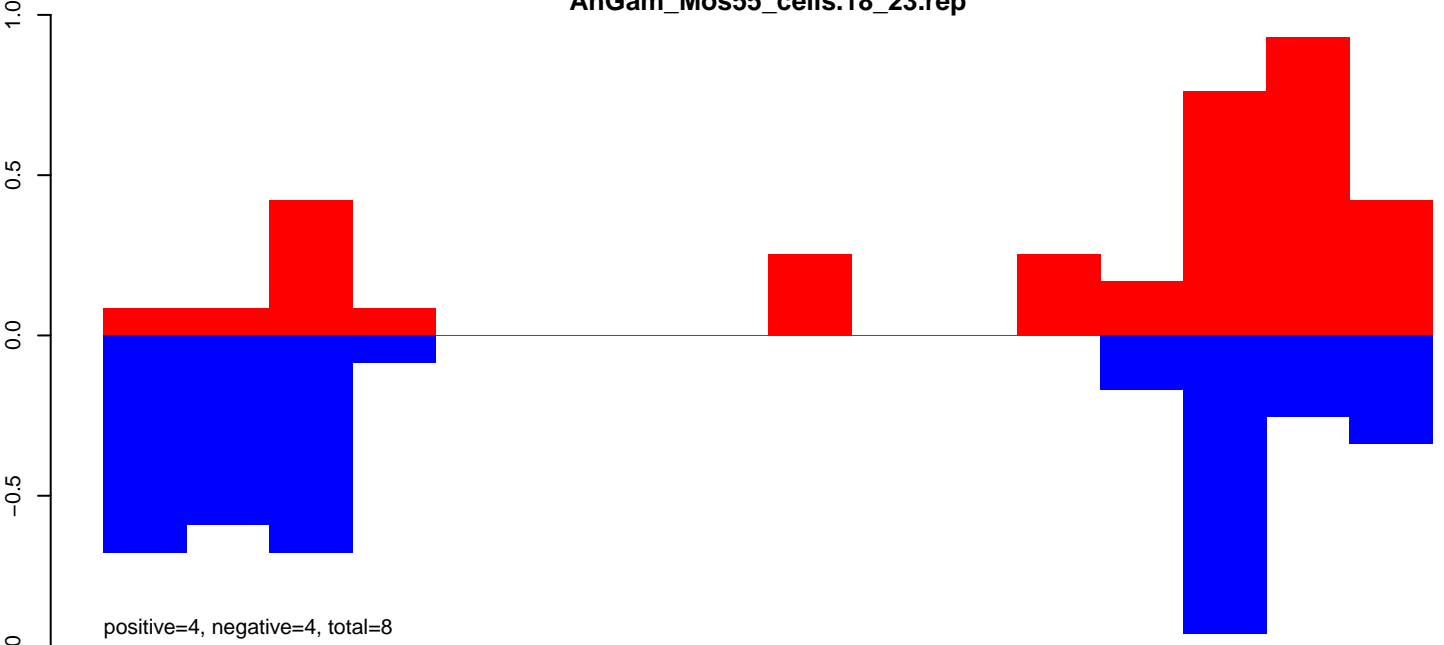
AnGam_Mos55_cells.rep



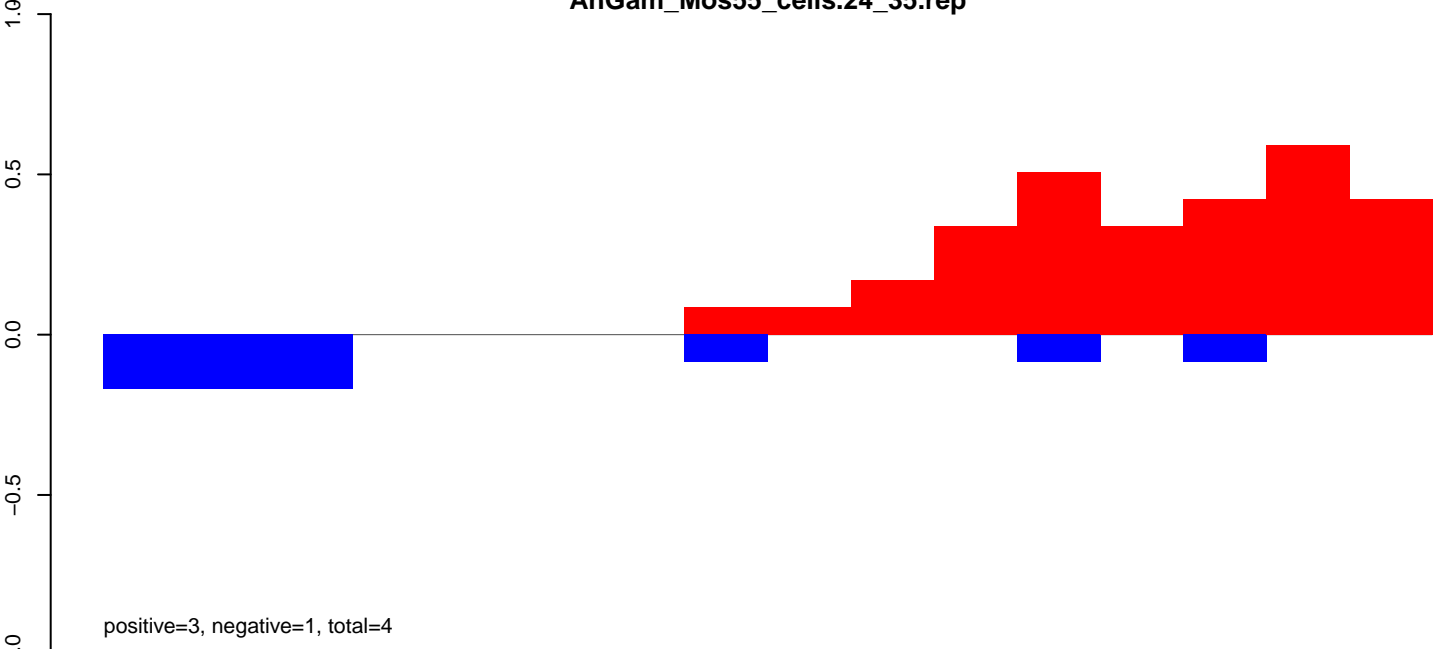
positive=2, negative=1, total=2

Window size=50, length=1168, TE@Clu-137B_AG:1-1168

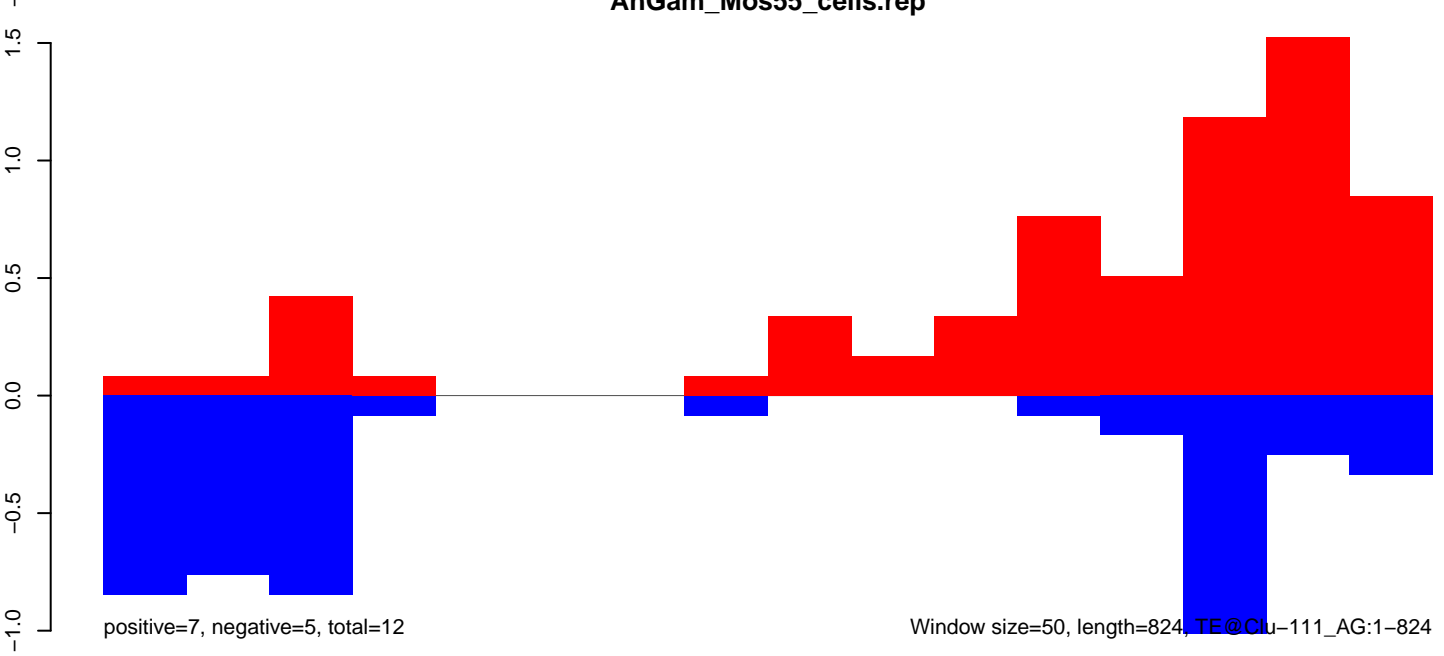
AnGam_Mos55_cells.18_23.rep



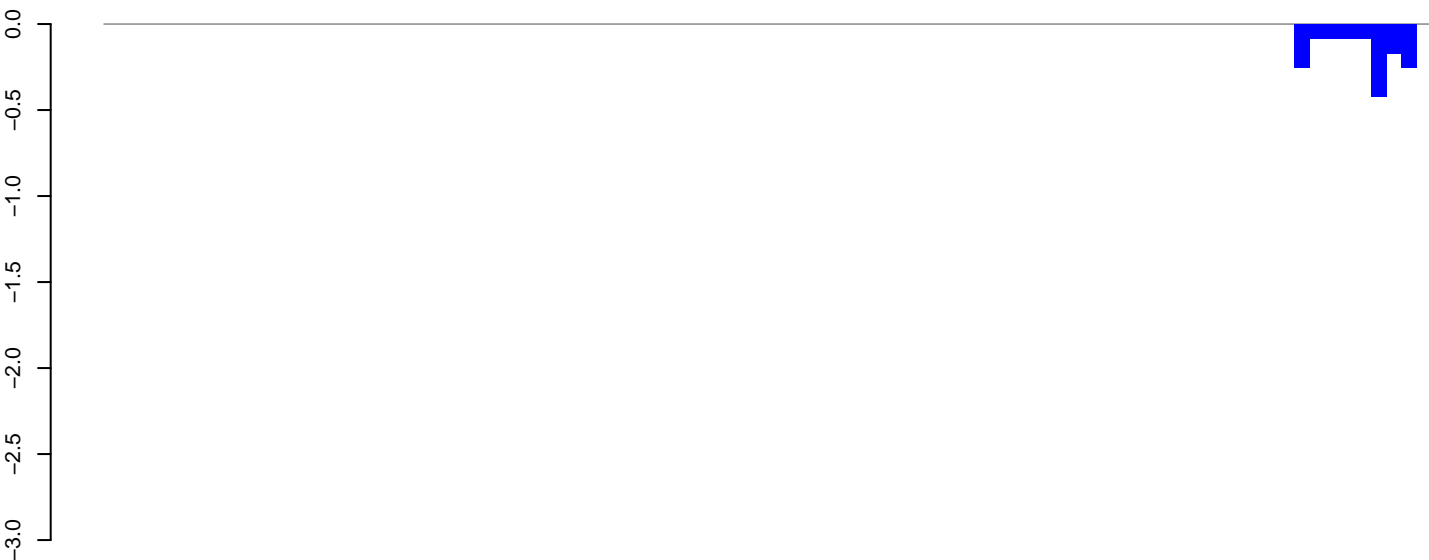
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

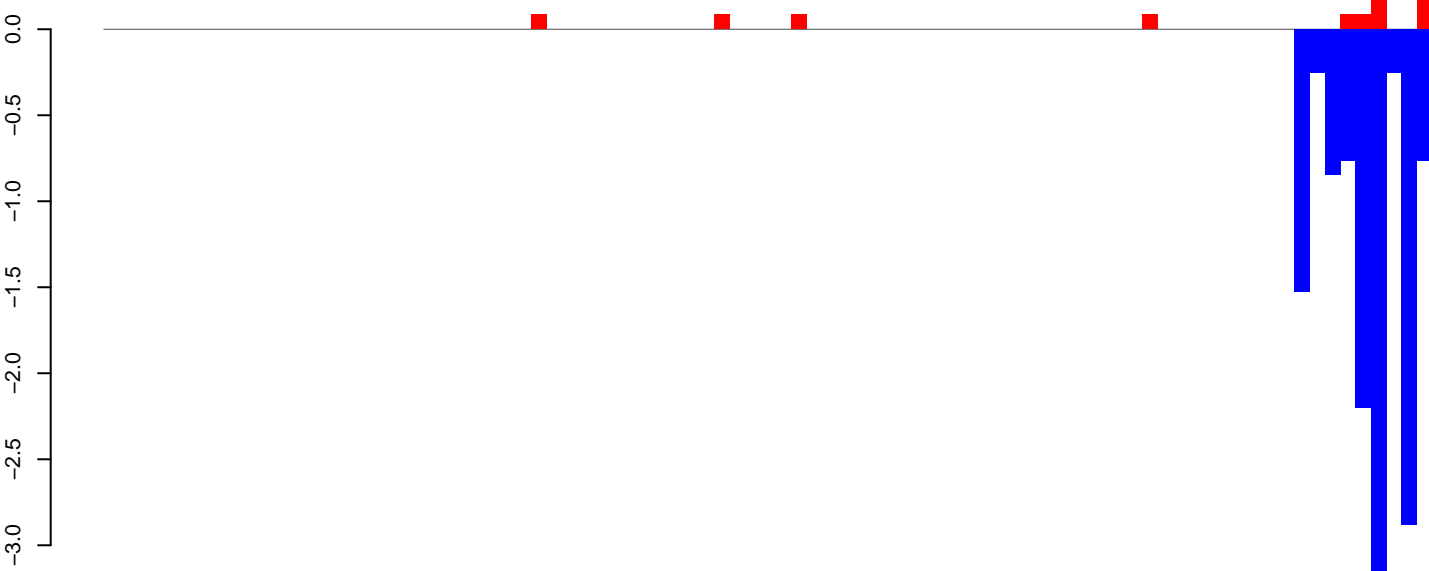


AnGam_Mos55_cells.18_23.rep



positive=0, negative=1, total=1

AnGam_Mos55_cells.24_35.rep



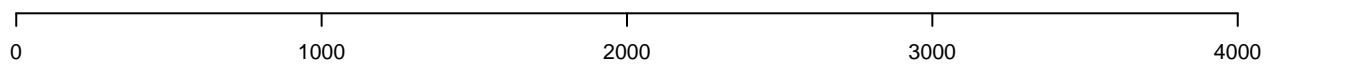
positive=1, negative=14, total=15

AnGam_Mos55_cells.rep

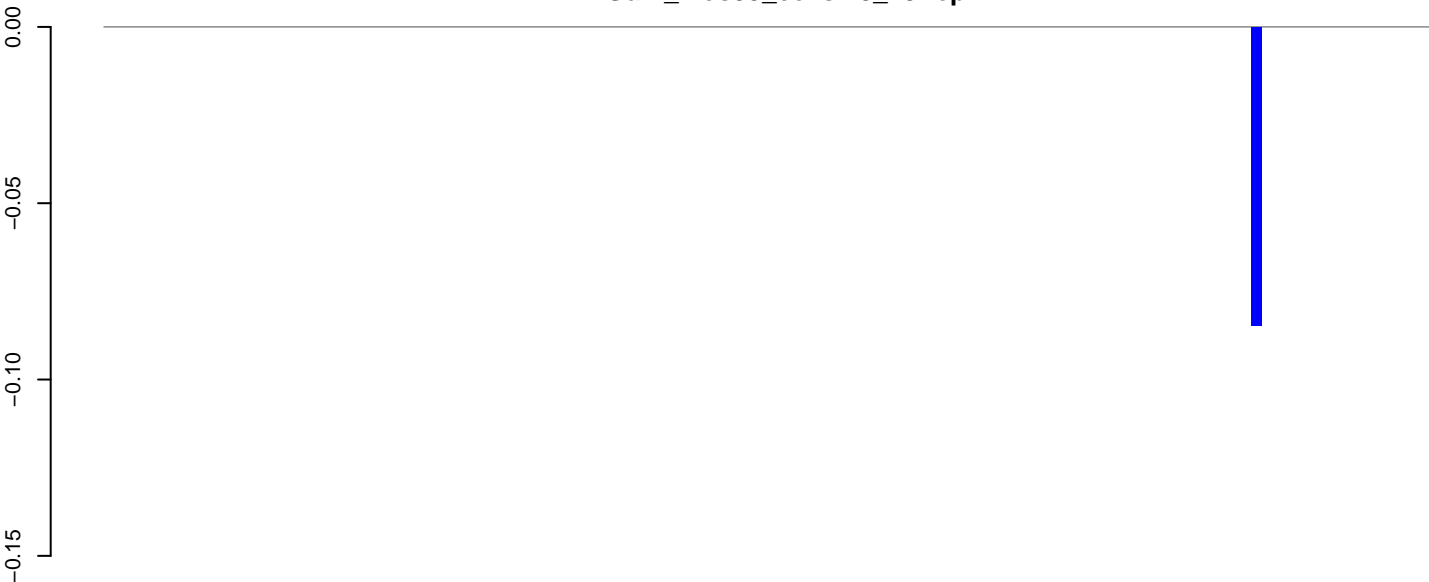


positive=1, negative=15, total=16

Window size=50, length=4397, TE@CR1-9_AG:1-4397

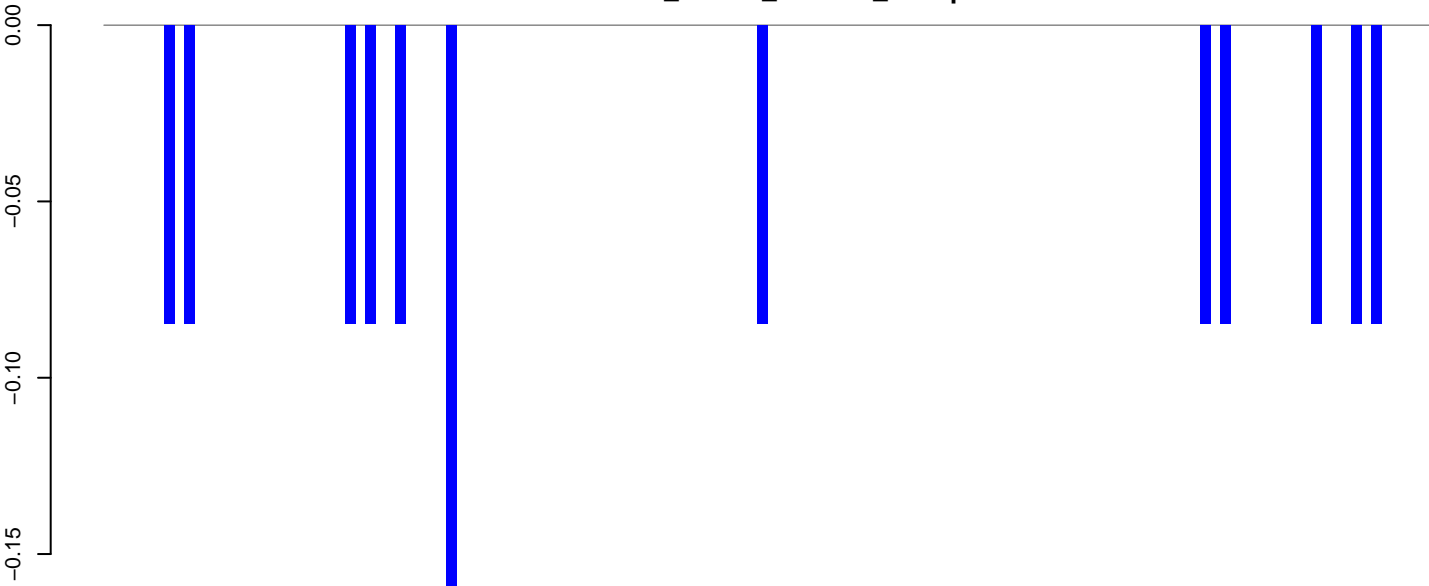


AnGam_Mos55_cells.18_23.rep



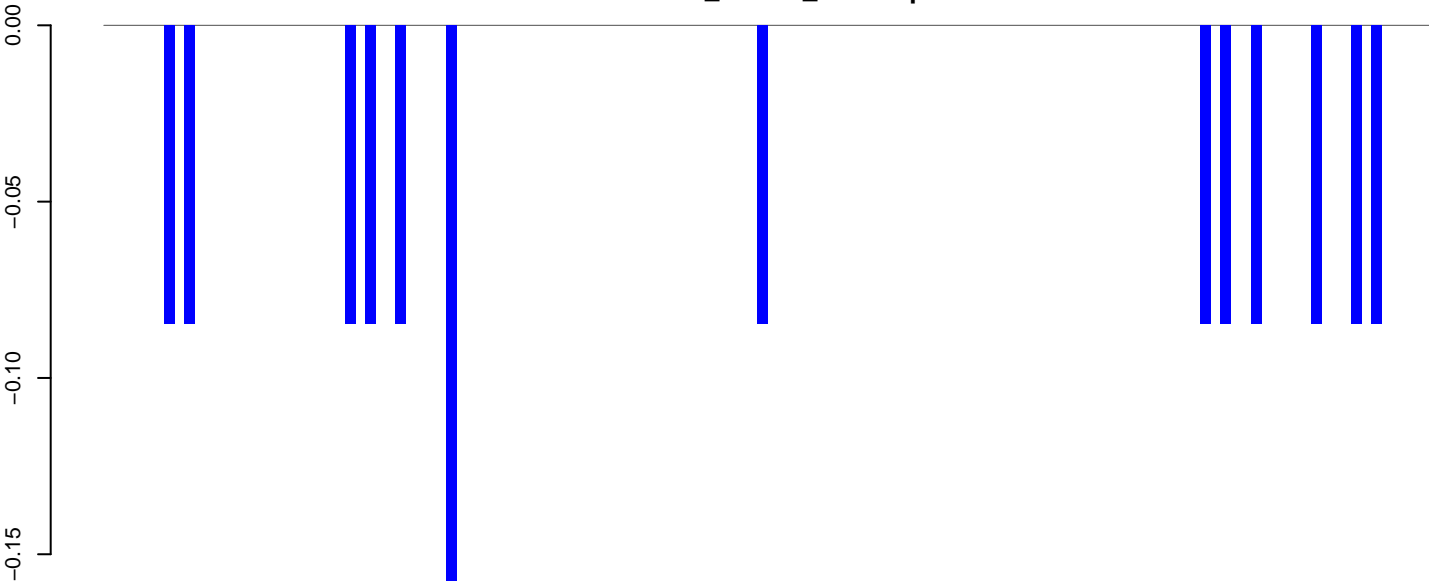
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



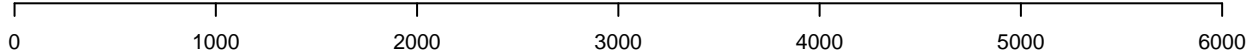
positive=0, negative=1, total=1

AnGam_Mos55_cells.rep

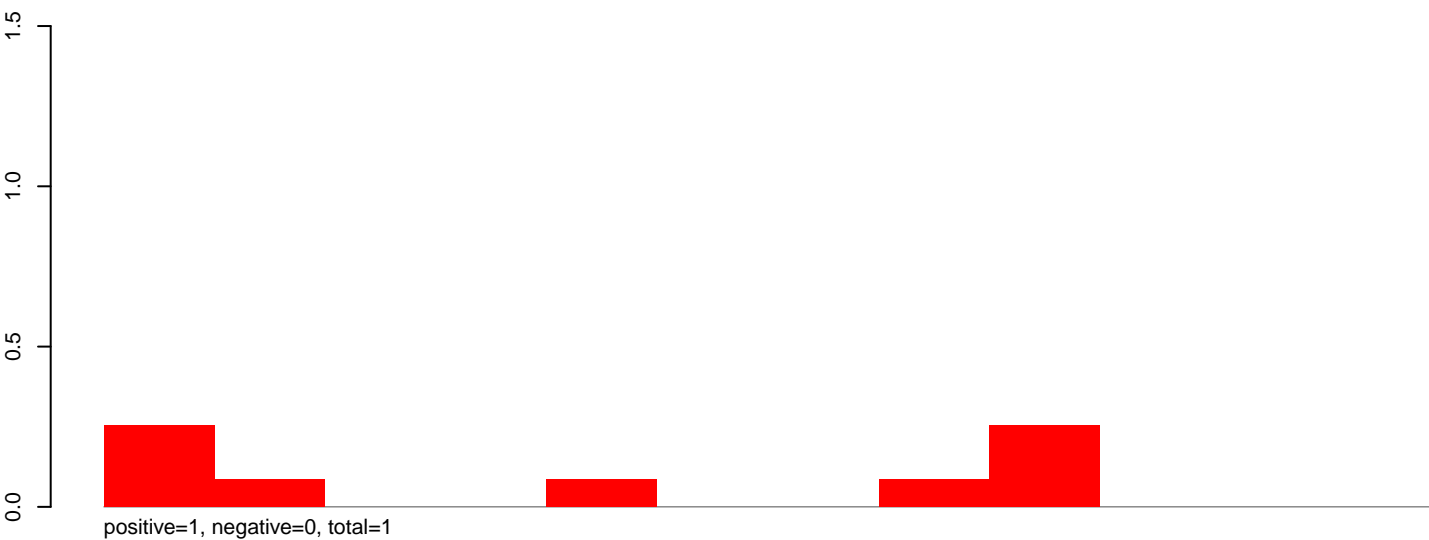


positive=0, negative=1, total=1

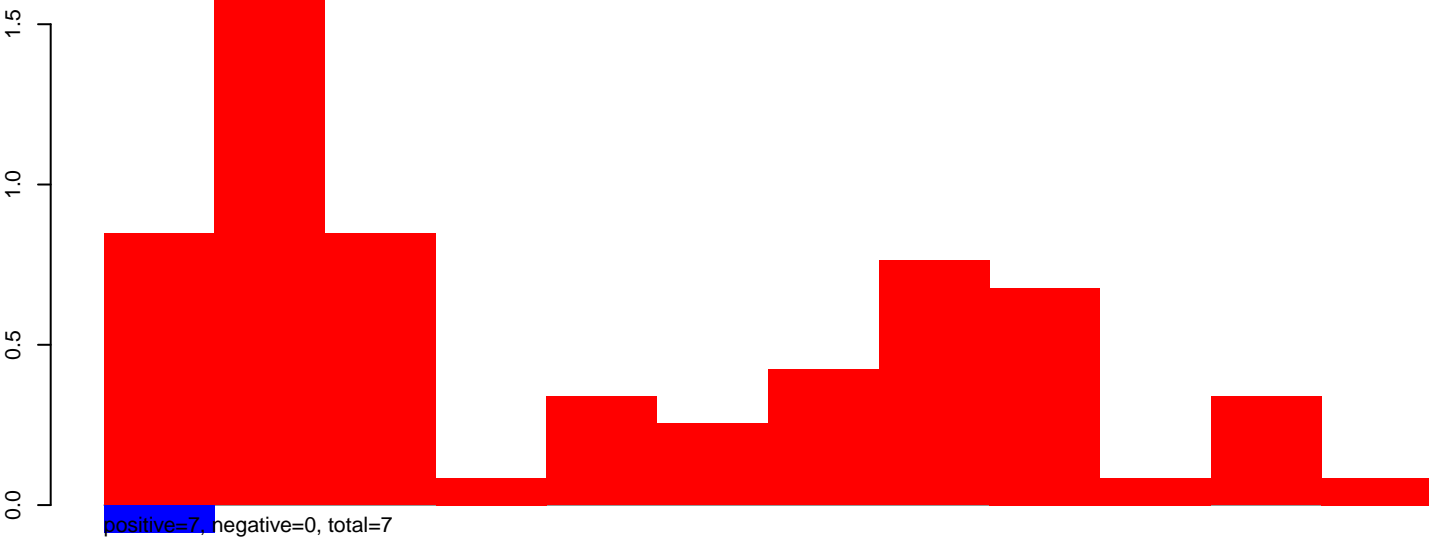
Window size=50, length=6608, TE@BEL18-LTR-I_AG:1-6608



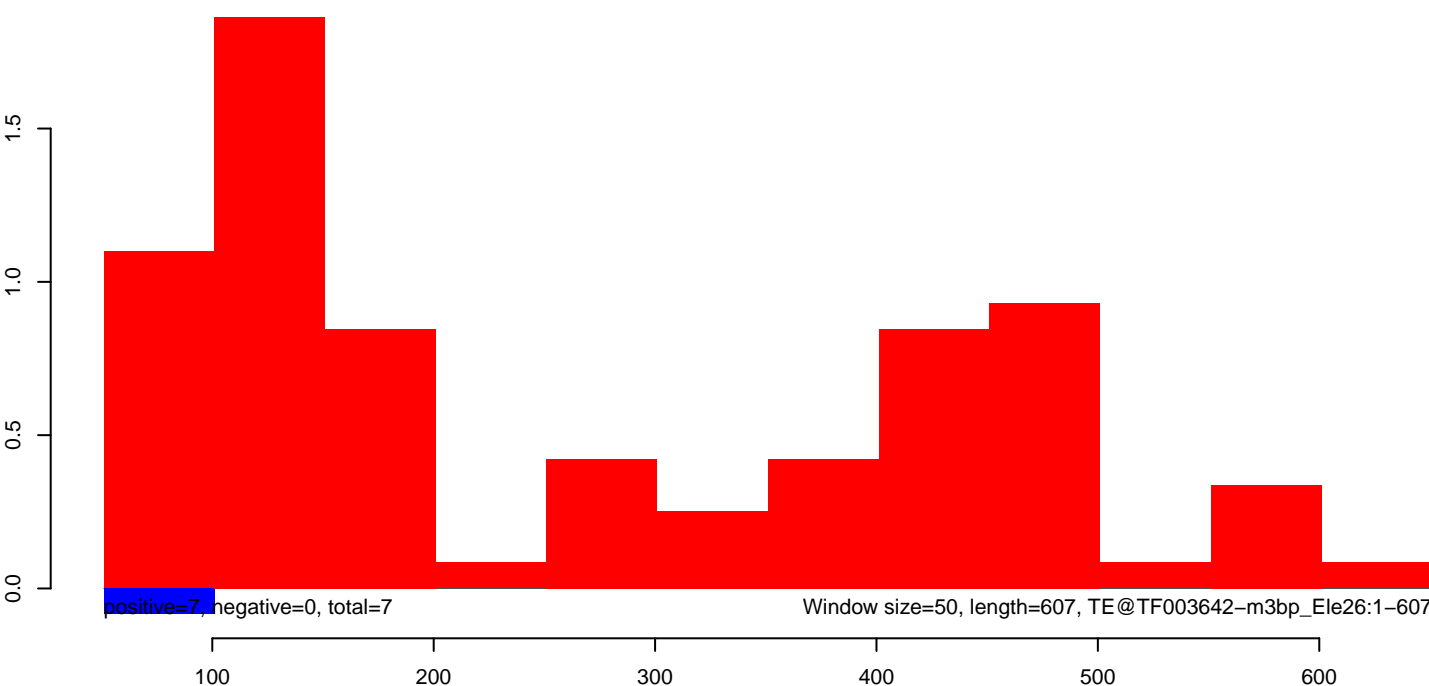
AnGam_Mos55_cells.18_23.rep



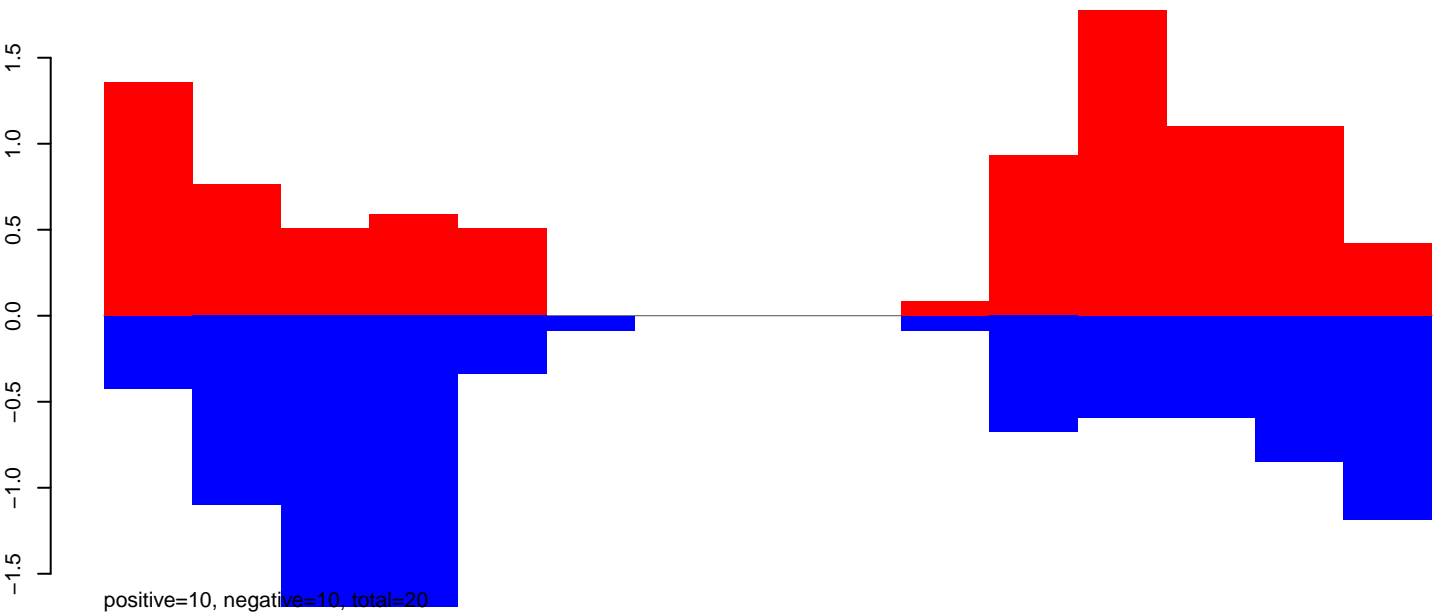
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



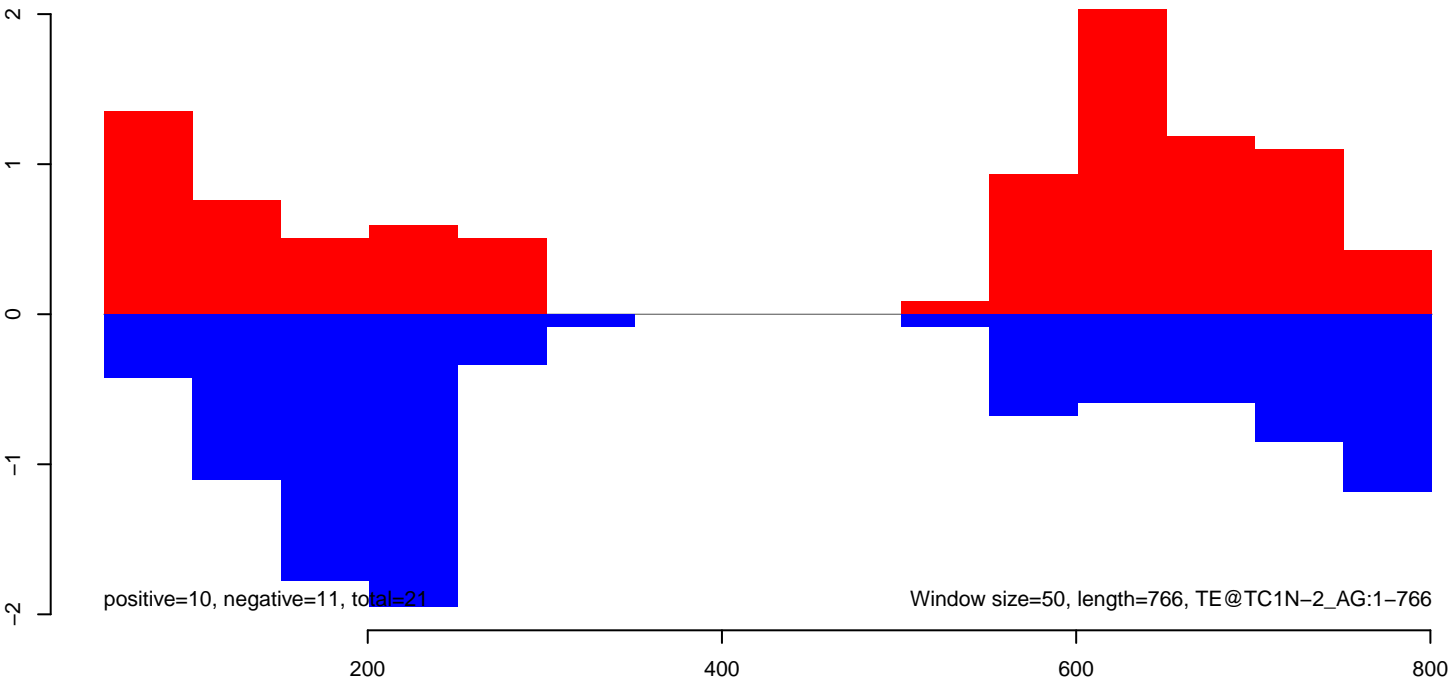
AnGam_Mos55_cells.18_23.rep



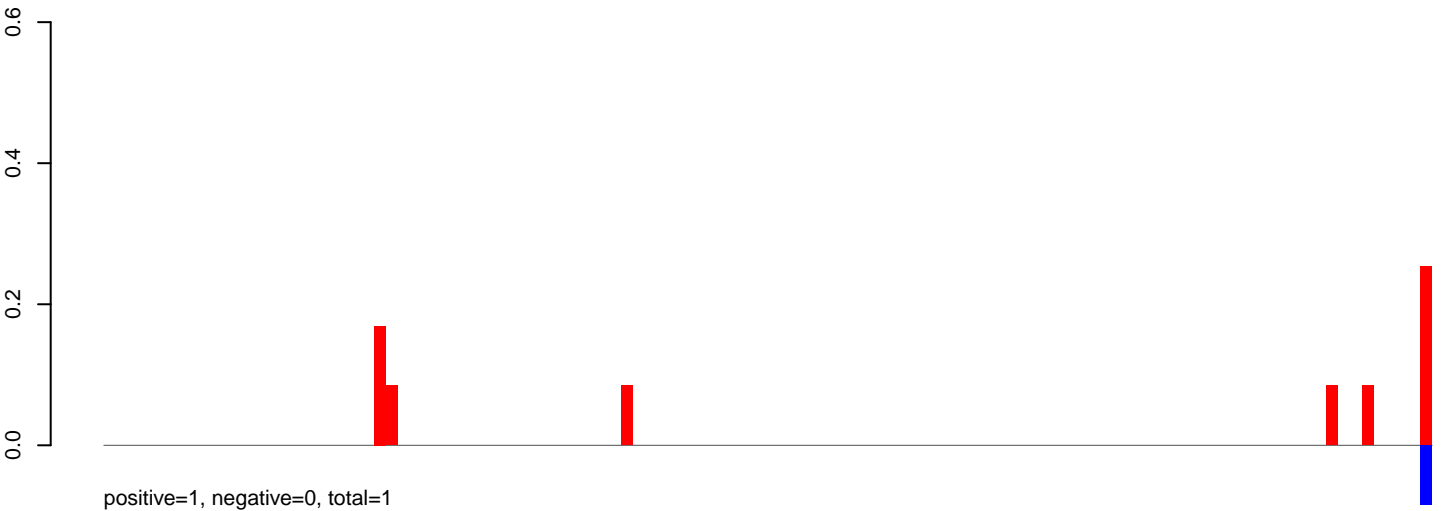
AnGam_Mos55_cells.24_35.rep



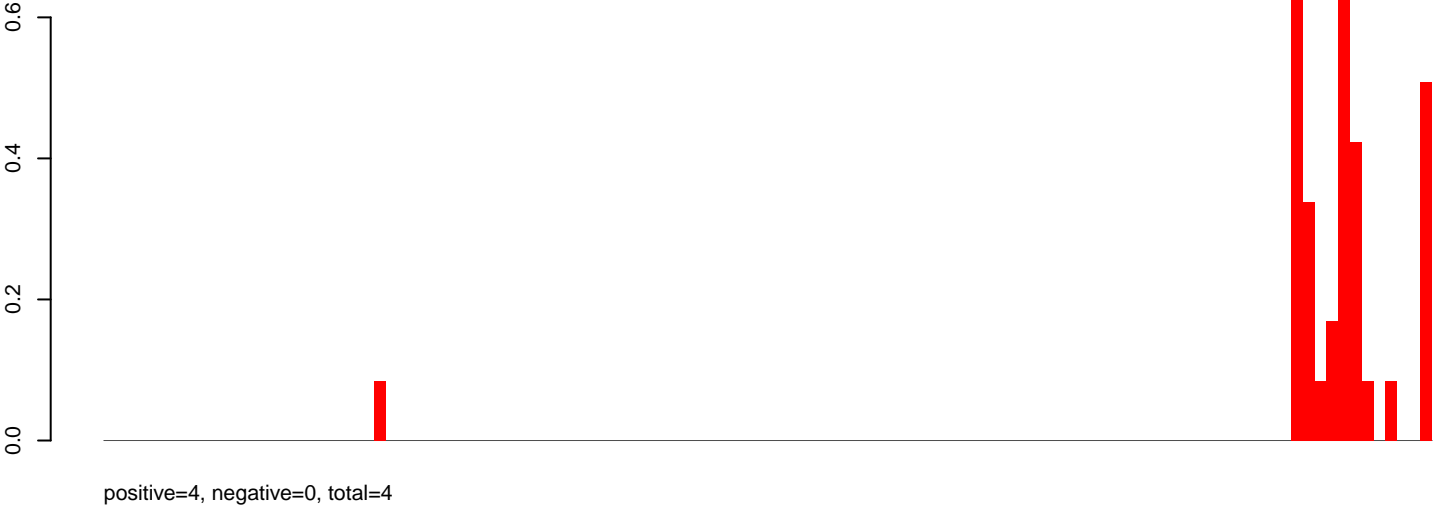
AnGam_Mos55_cells.rep



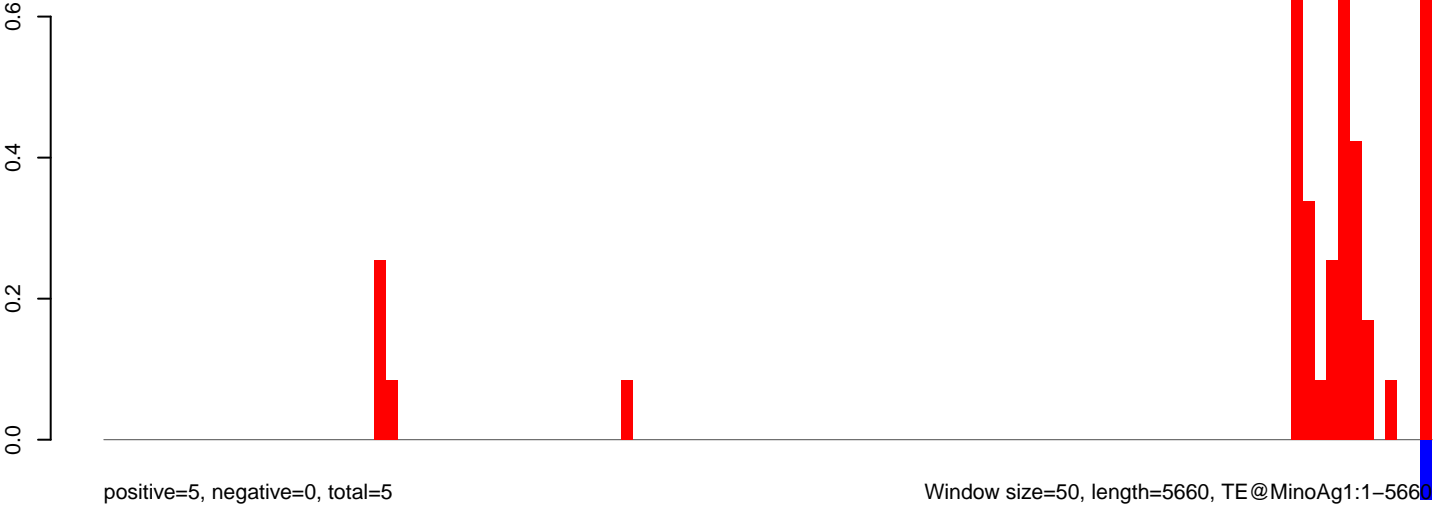
AnGam_Mos55_cells.18_23.rep



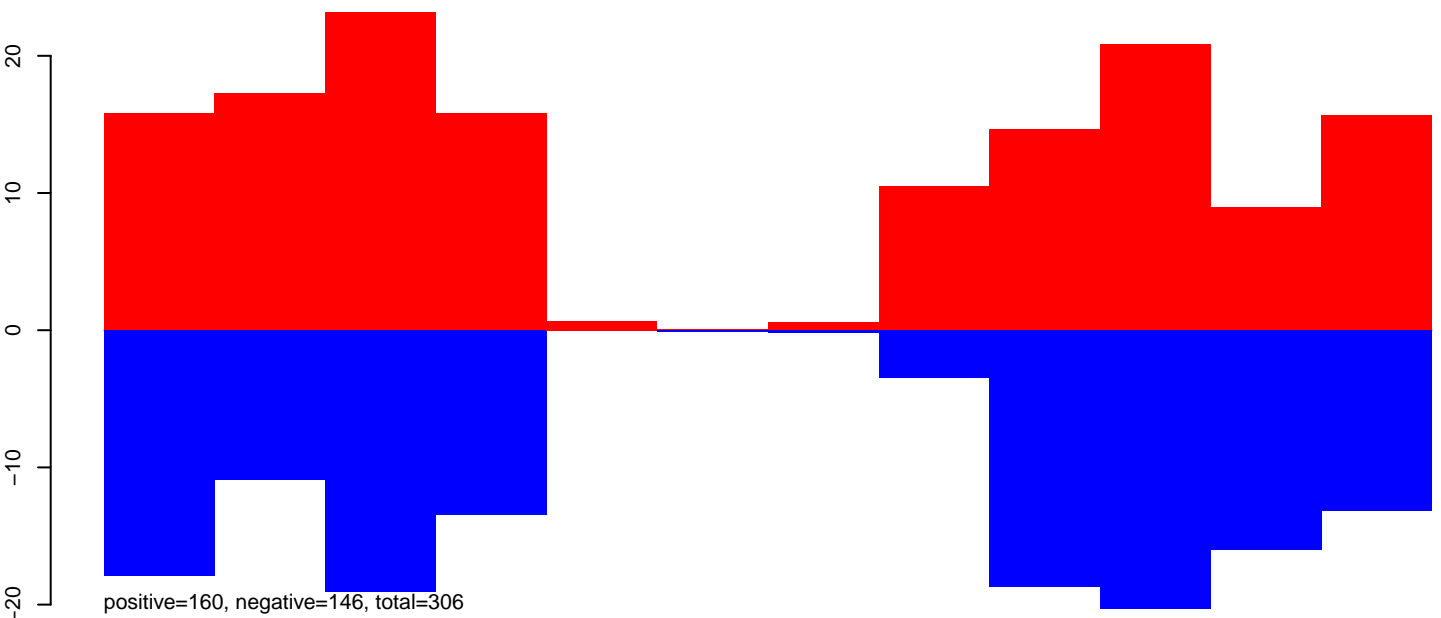
AnGam_Mos55_cells.24_35.rep



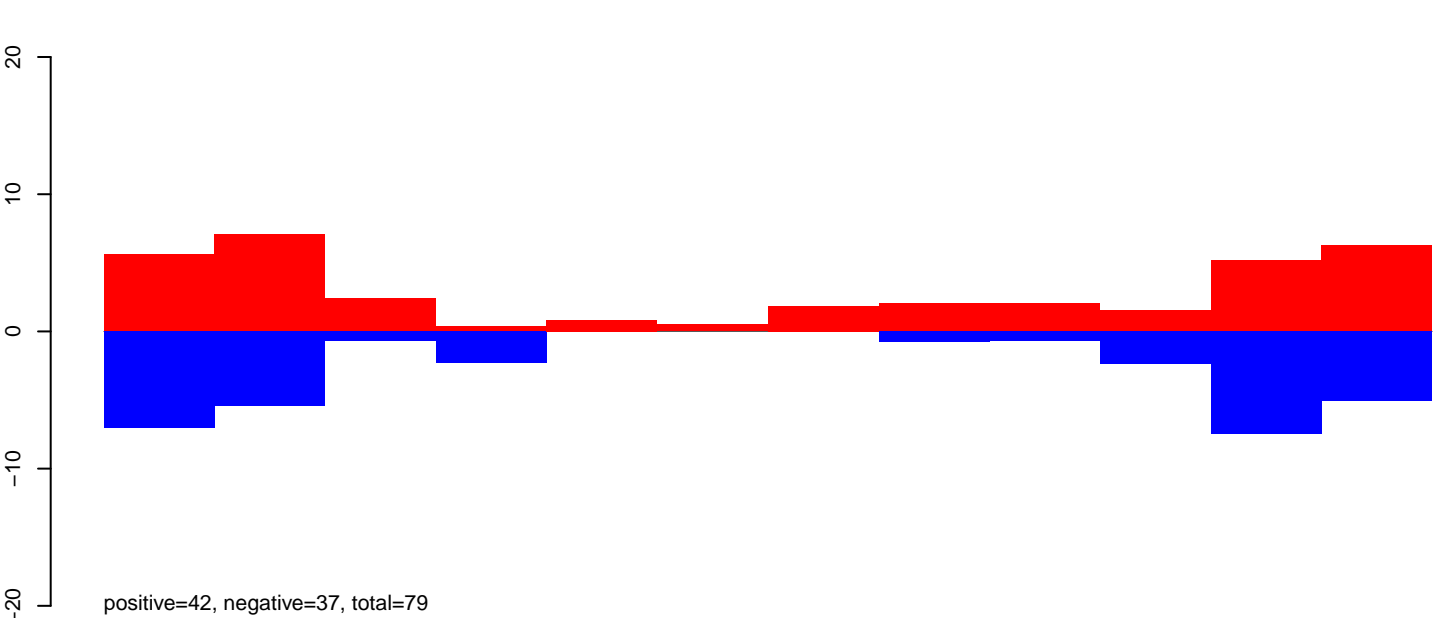
AnGam_Mos55_cells.rep



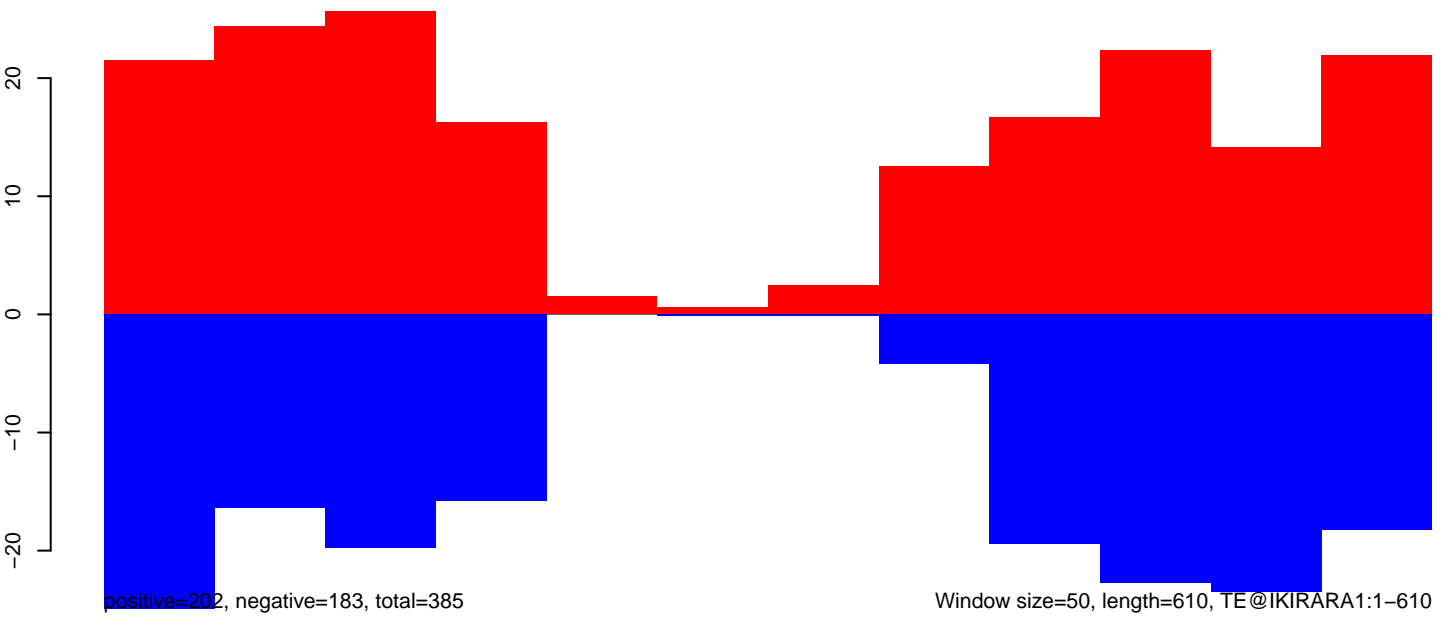
AnGam_Mos55_cells.18_23.rep



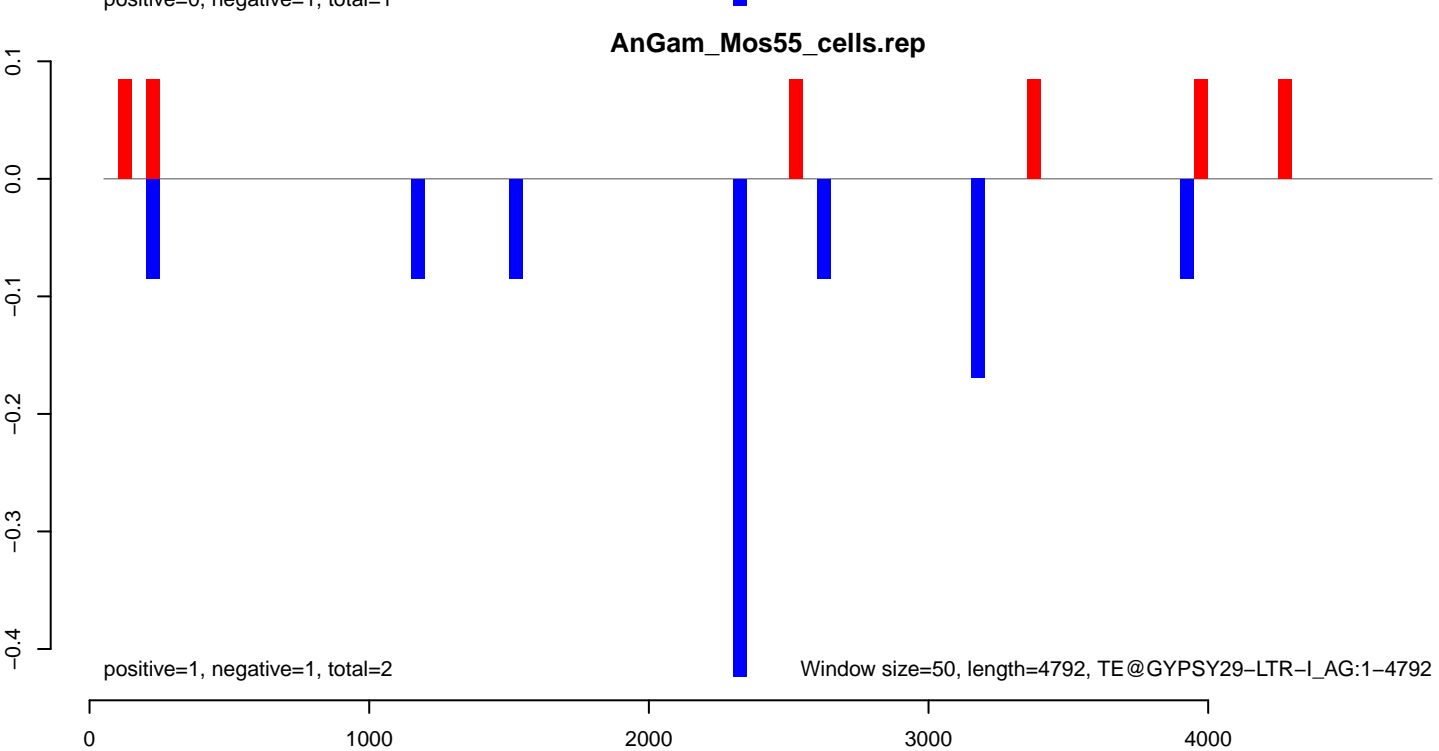
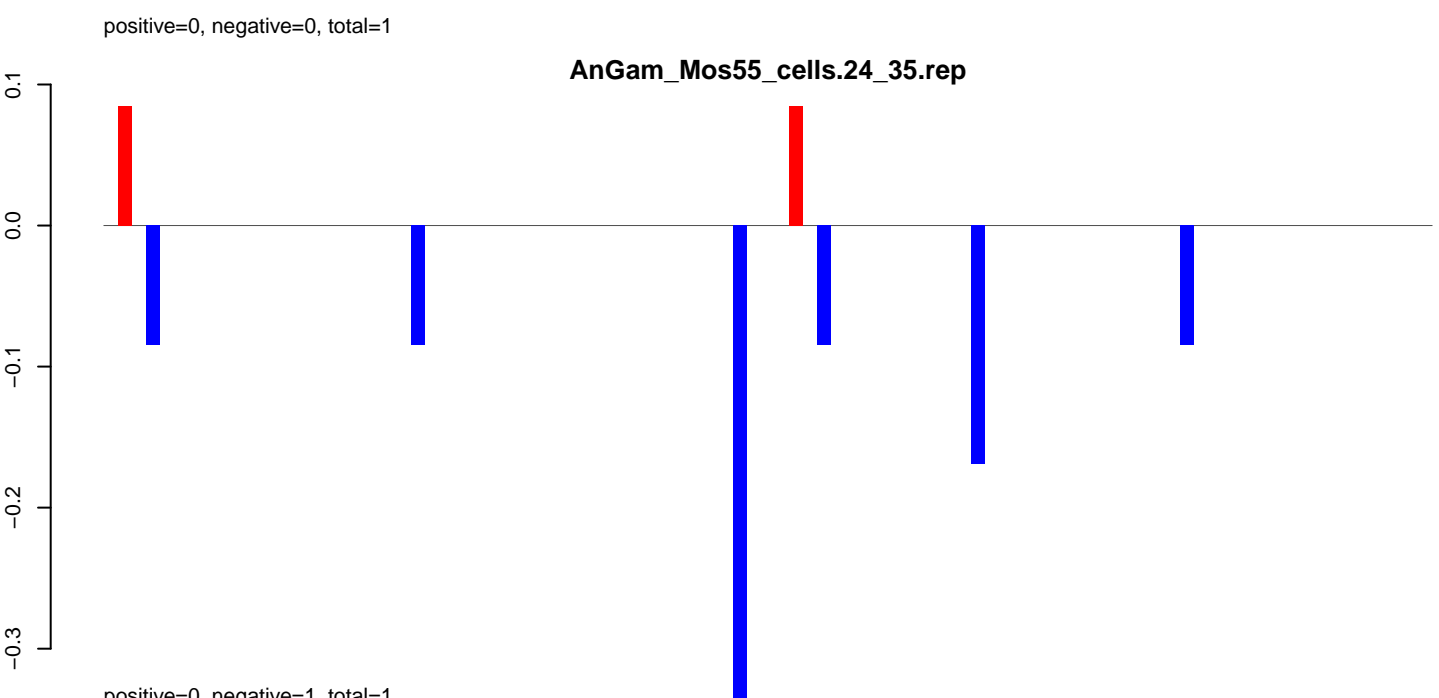
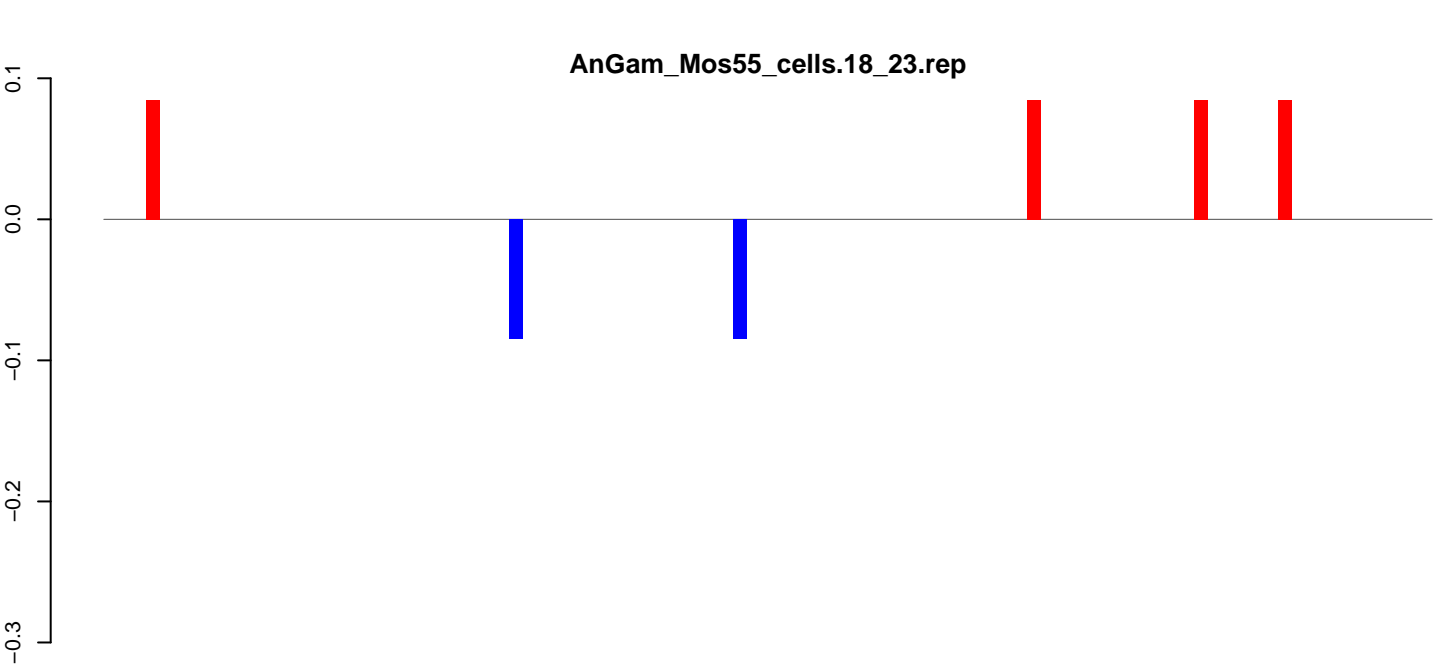
AnGam_Mos55_cells.24_35.rep



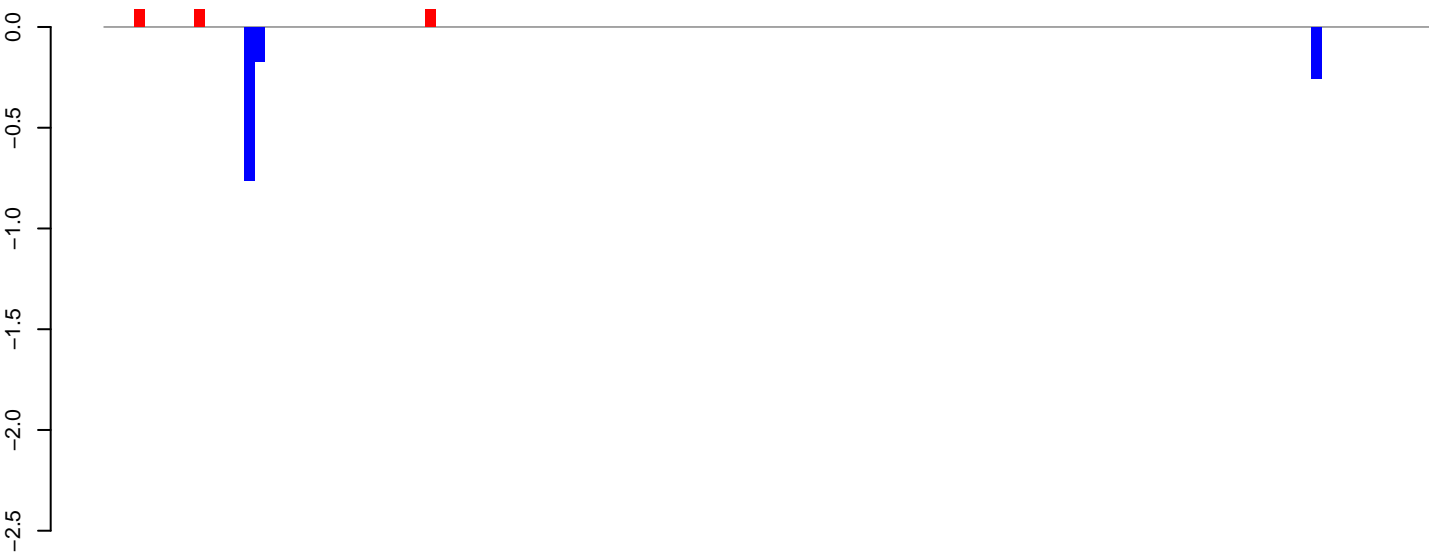
AnGam_Mos55_cells.rep



Window size=50, length=610, TE@IKIRARA1:1-610



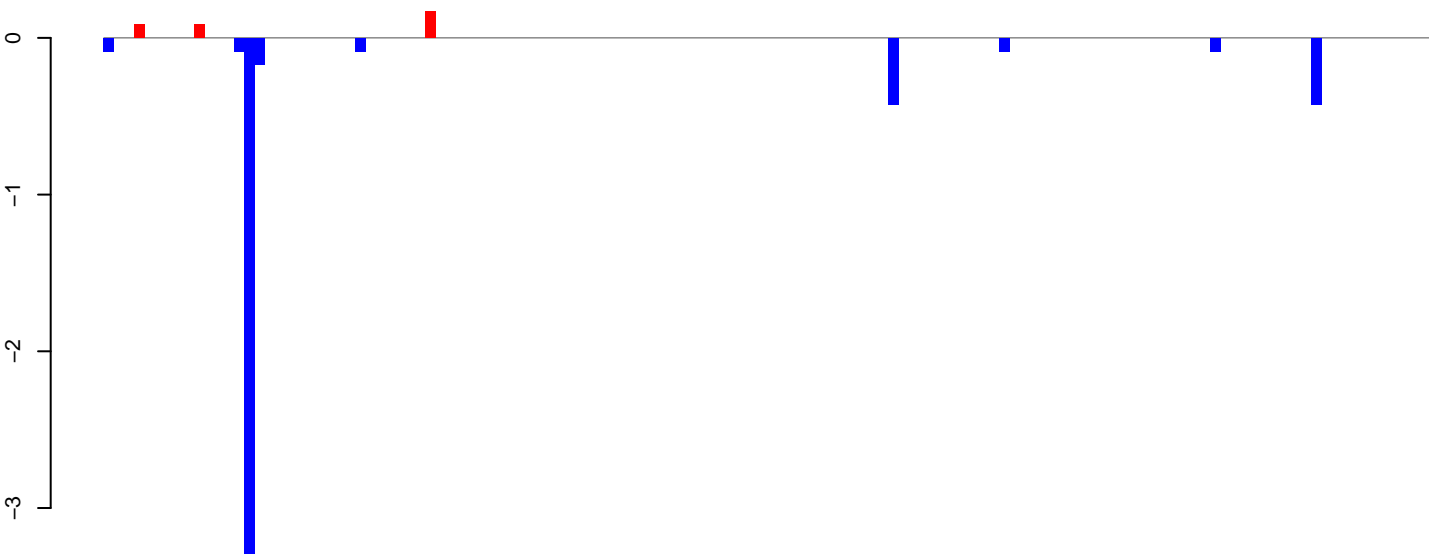
AnGam_Mos55_cells.18_23.rep



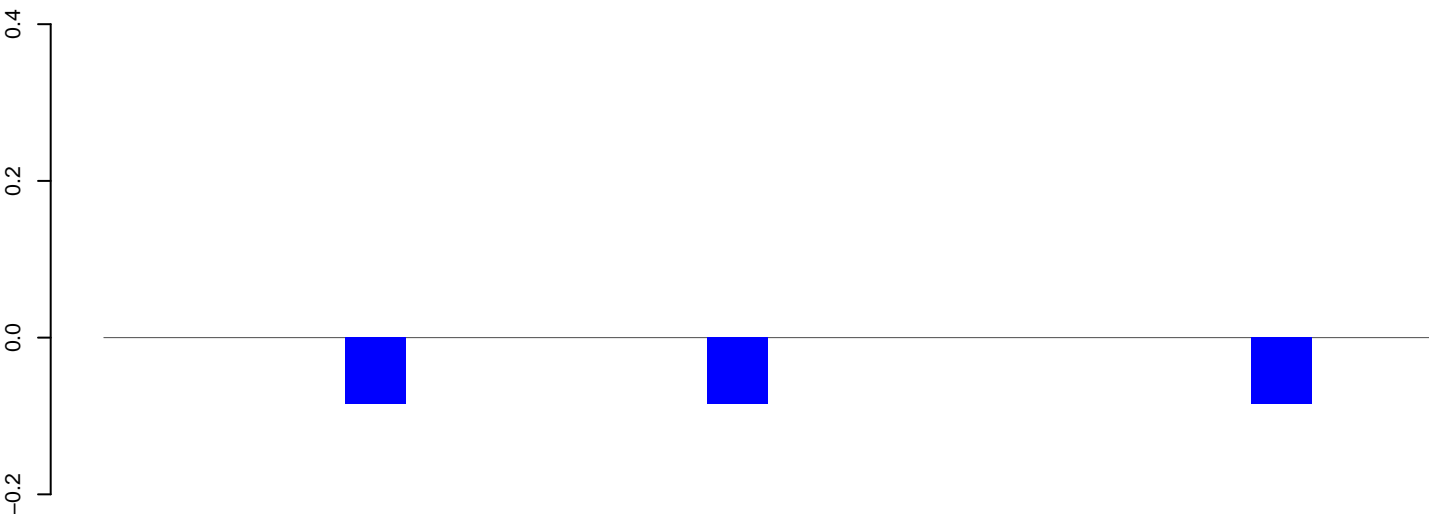
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

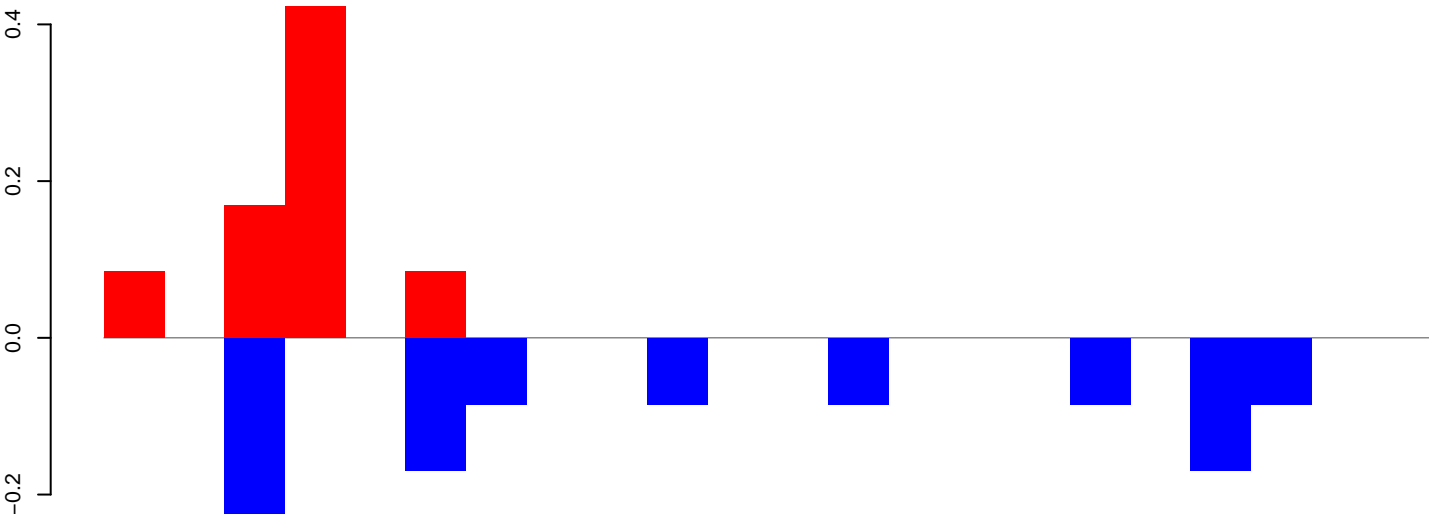


AnGam_Mos55_cells.18_23.rep



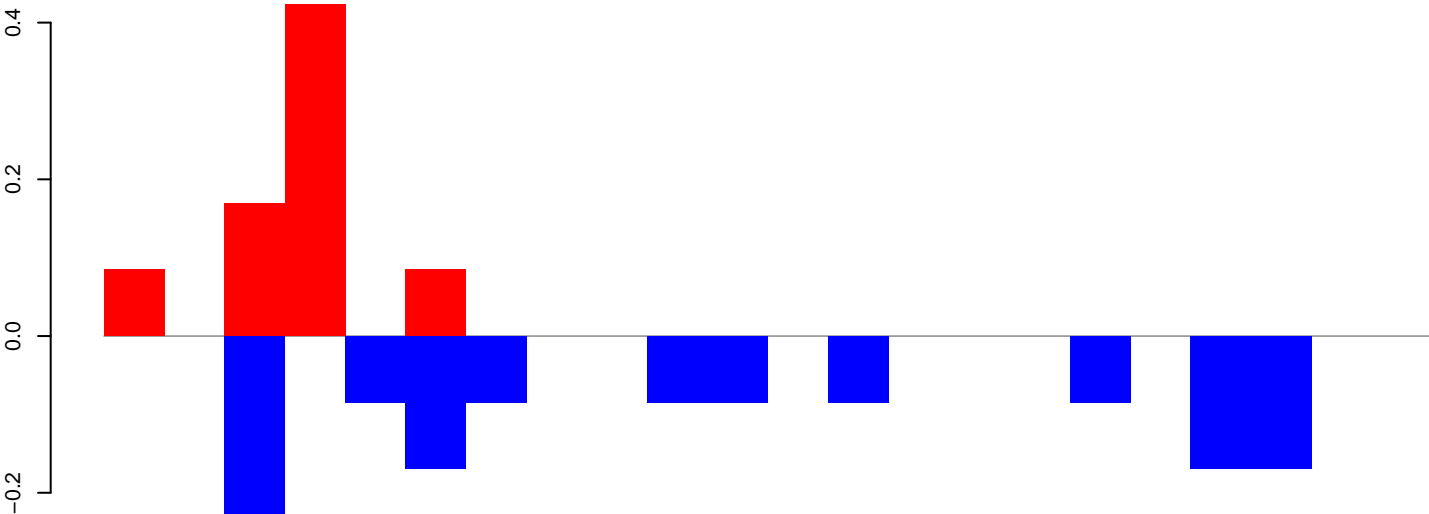
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=1, negative=1, total=2

AnGam_Mos55_cells.rep



positive=1, negative=1, total=2

Window size=50, length=1110, TE@Clu-39B_AG:1-1110

AnGam_Mos55_cells.18_23.rep

5
0
-5
-10

positive=17, negative=28, total=44

AnGam_Mos55_cells.24_35.rep

5
0
-5
-10

positive=3, negative=5, total=8

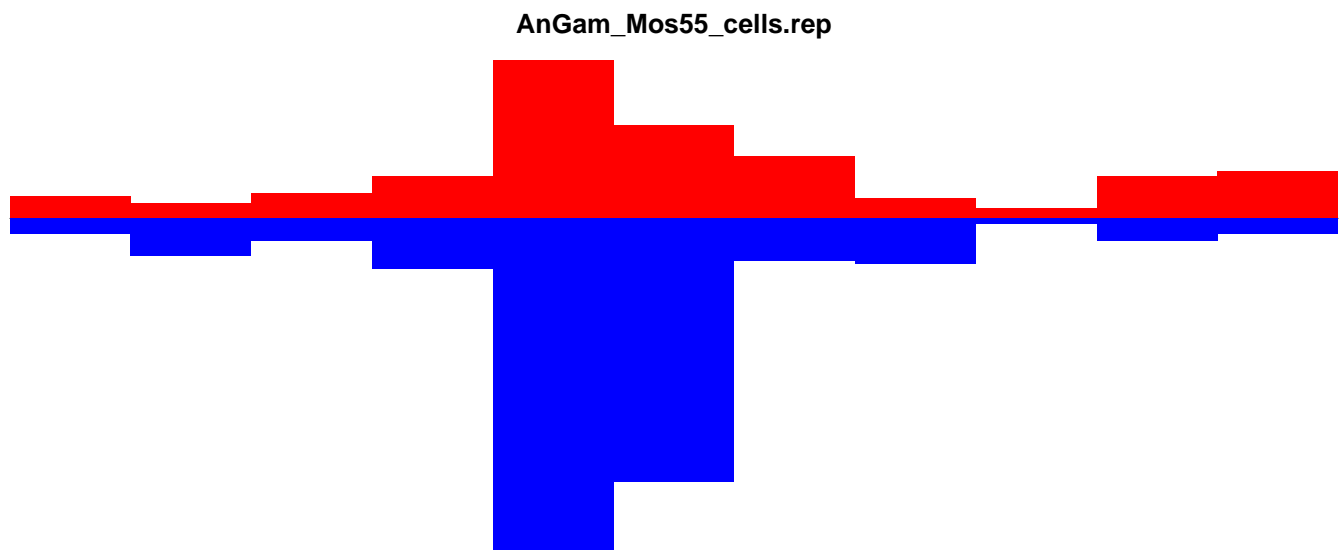
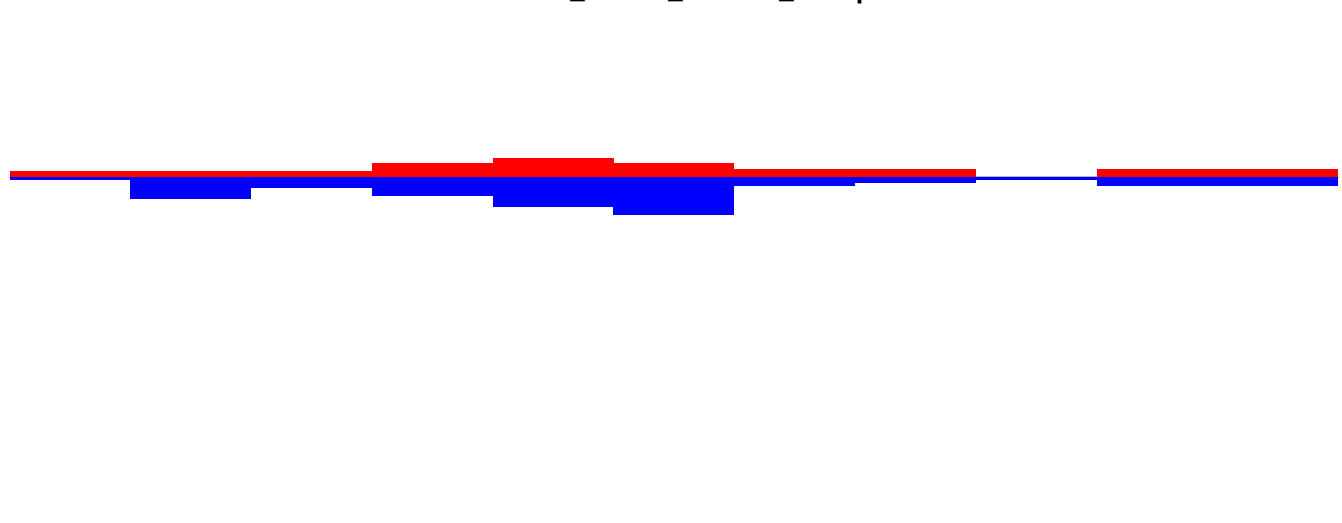
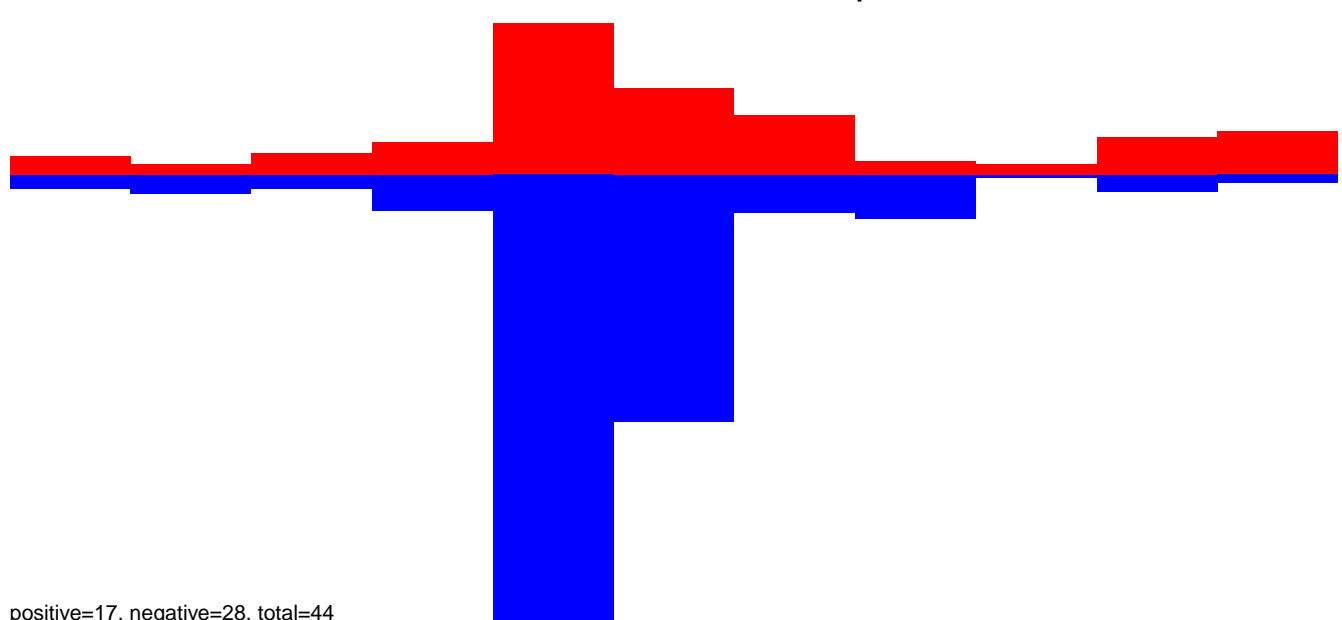
AnGam_Mos55_cells.rep

5
0
-5
-10
-15

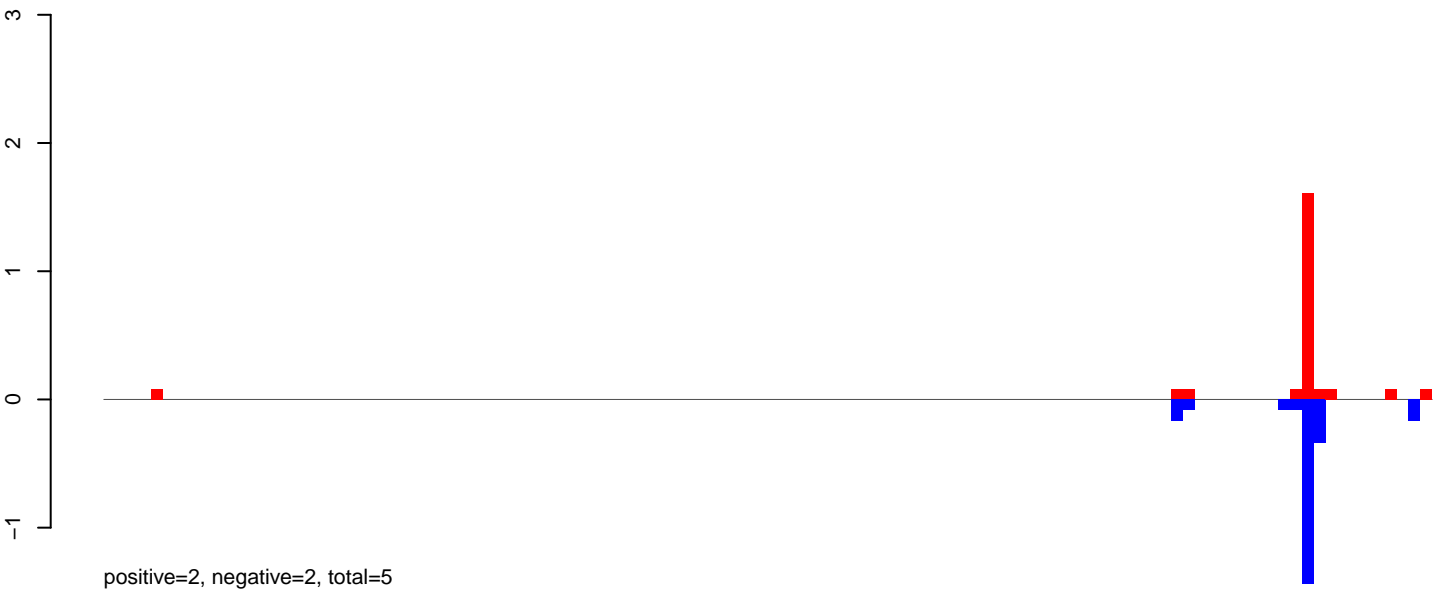
positive=20, negative=33, total=53

Window size=50, length=588, TE@TF003619-m2bp_Ele10:1-588

100 200 300 400 500 600



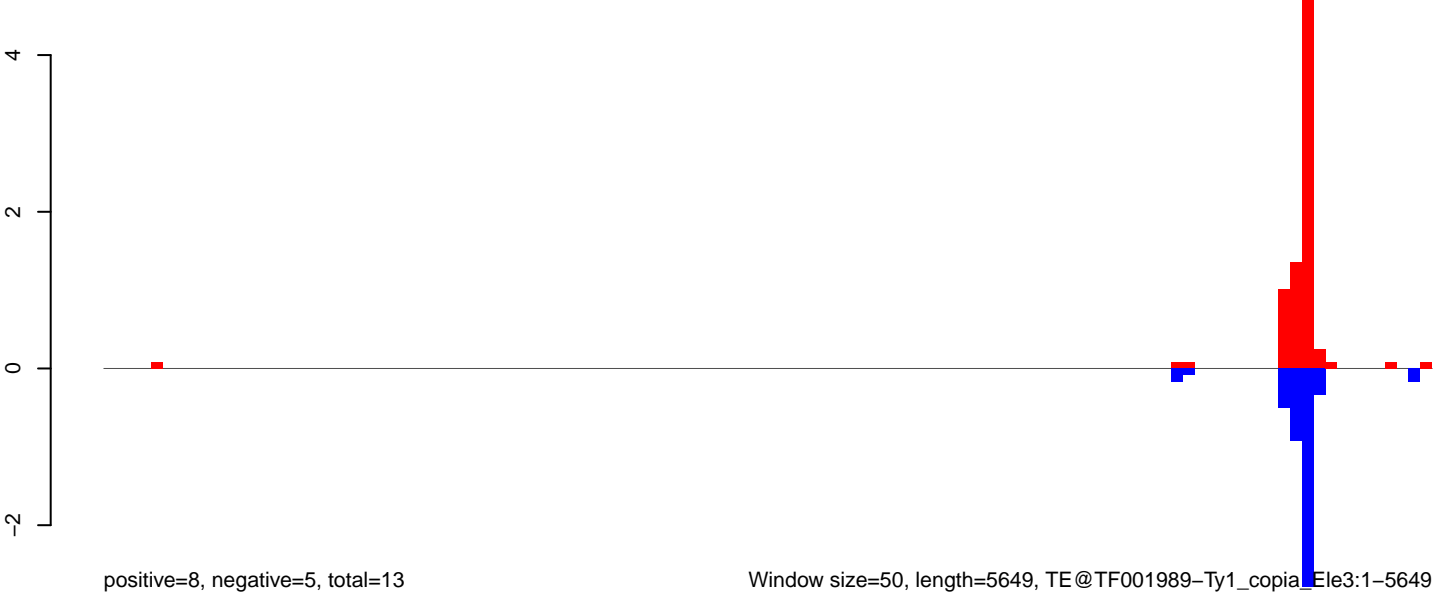
AnGam_Mos55_cells.18_23.rep



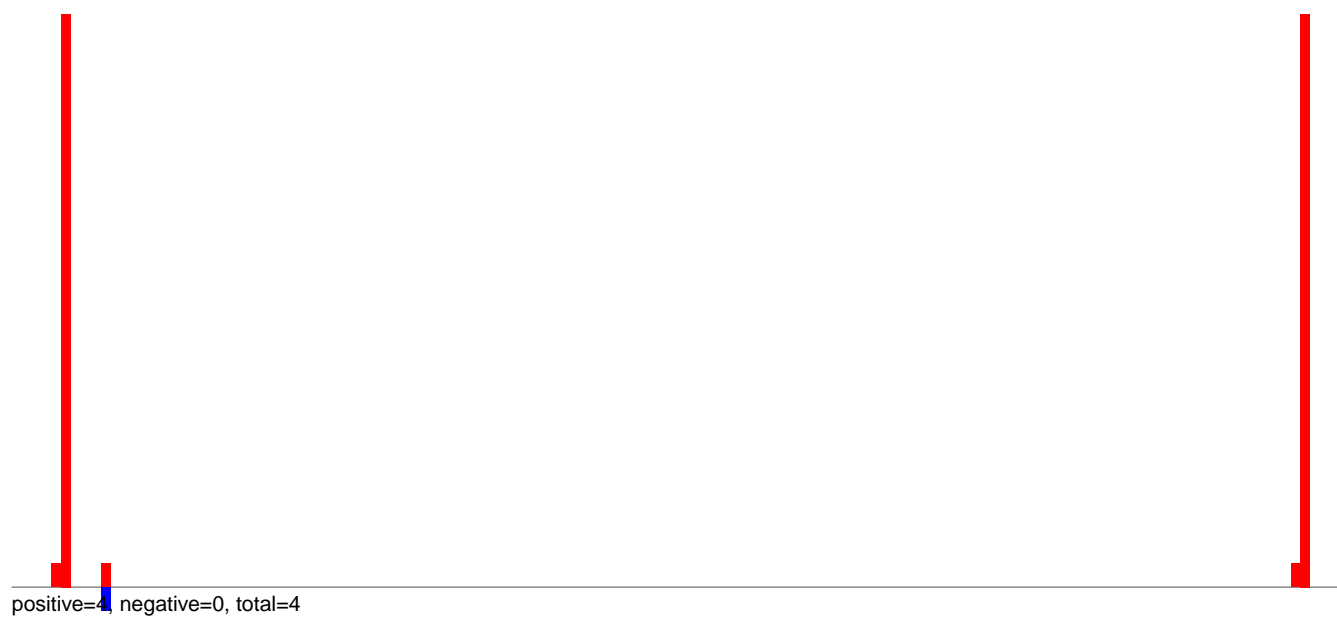
AnGam_Mos55_cells.24_35.rep



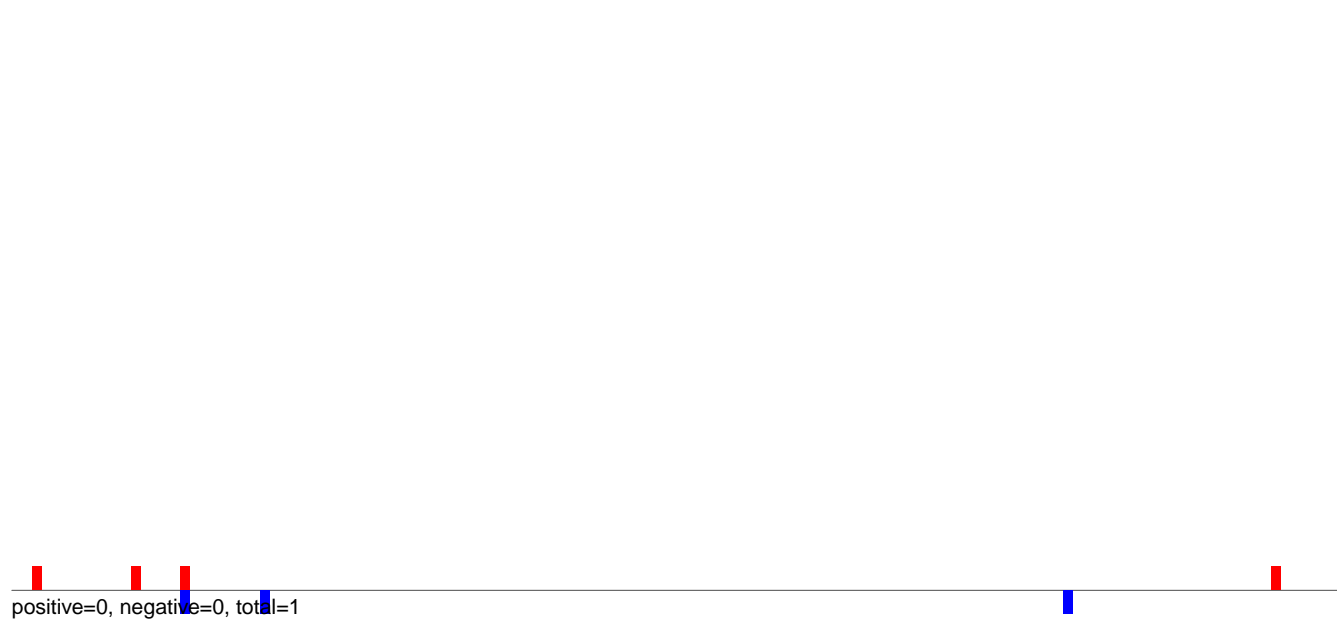
AnGam_Mos55_cells.rep



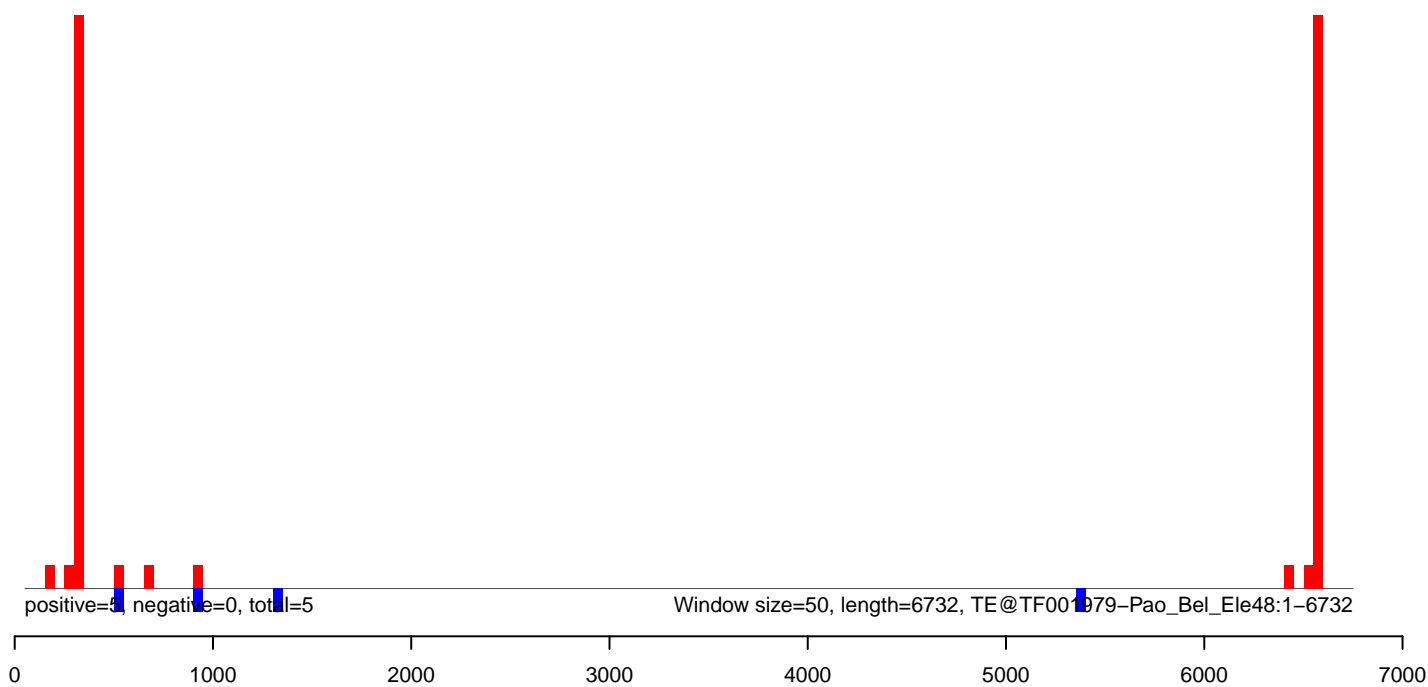
AnGam_Mos55_cells.18_23.rep



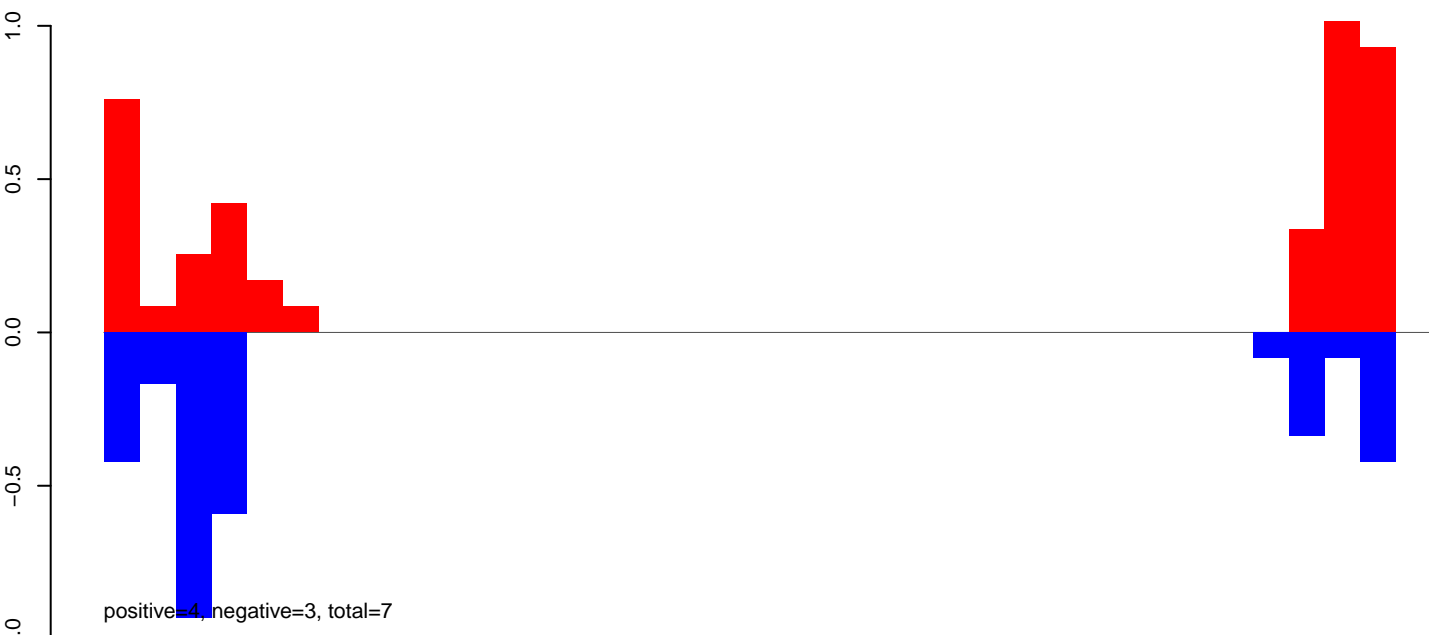
AnGam_Mos55_cells.24_35.rep



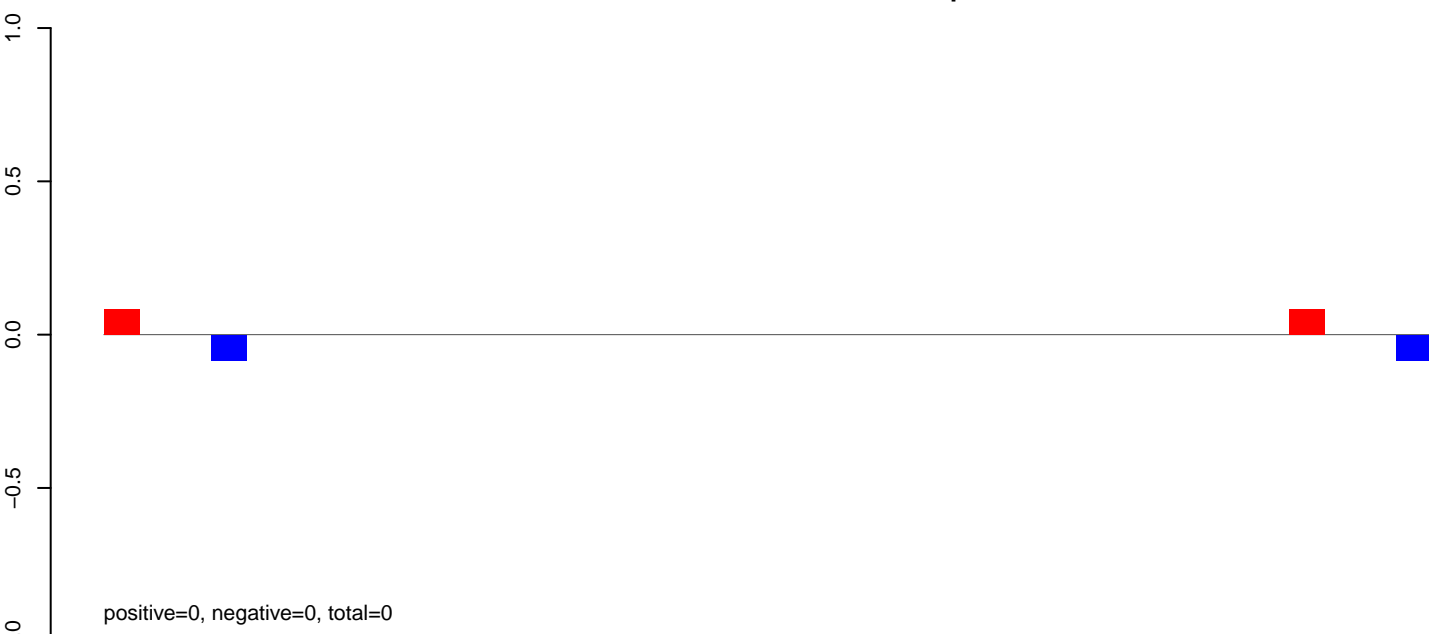
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



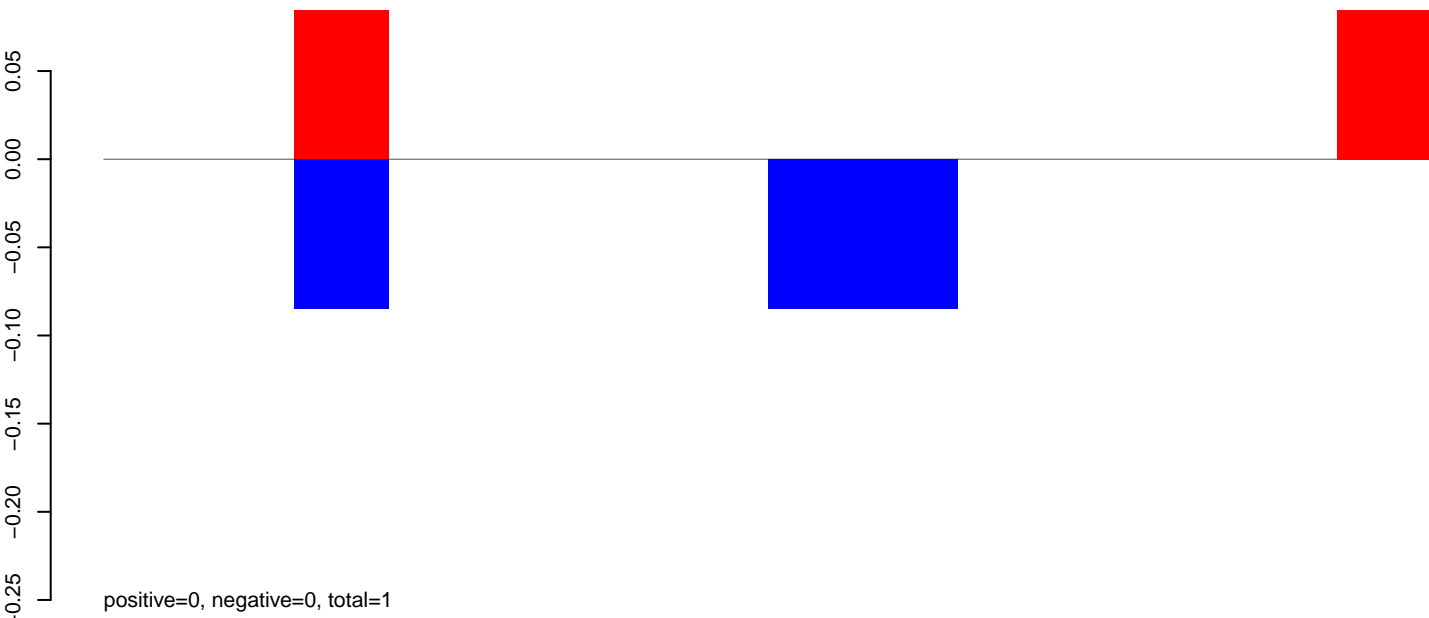
AnGam_Mos55_cells.24_35.rep



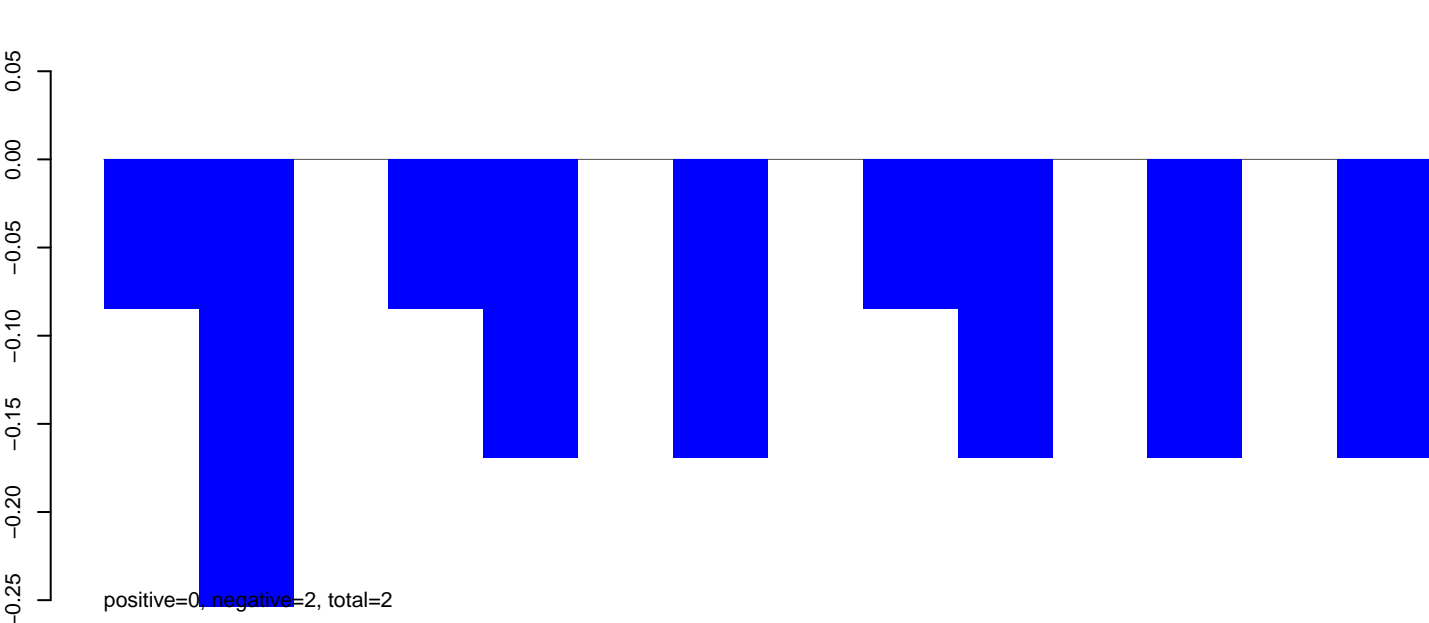
AnGam_Mos55_cells.rep



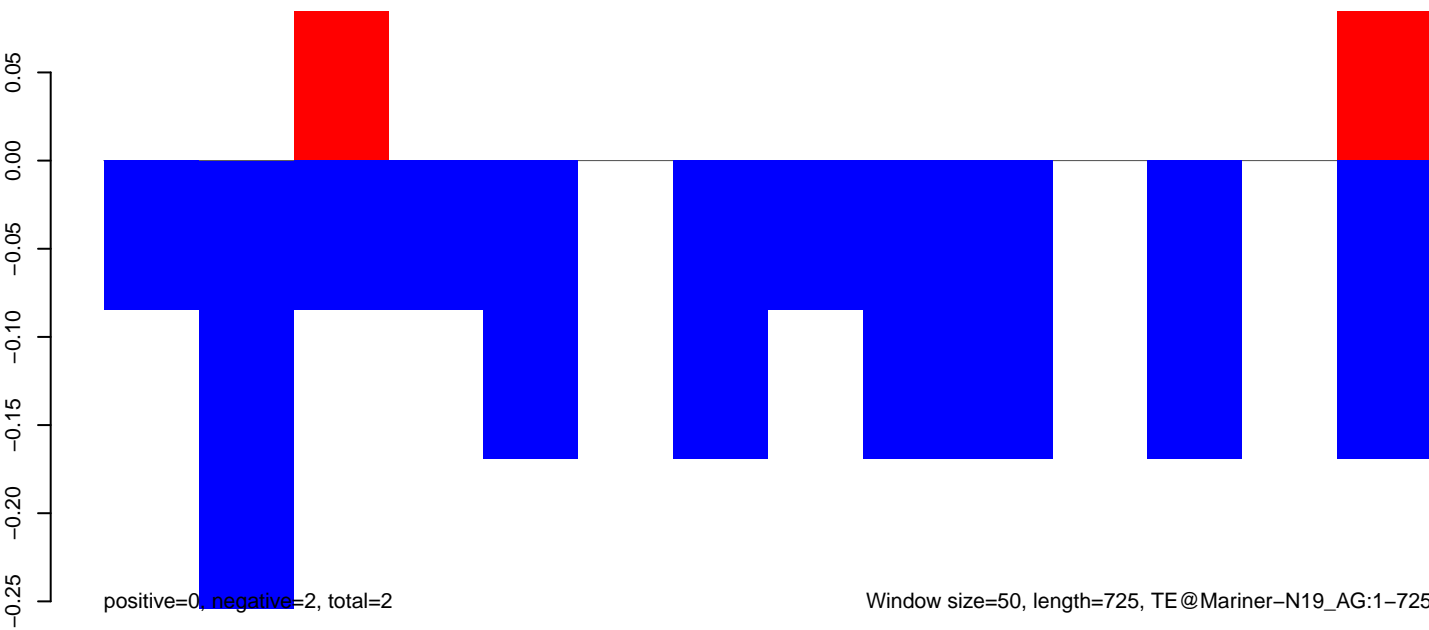
AnGam_Mos55_cells.18_23.rep



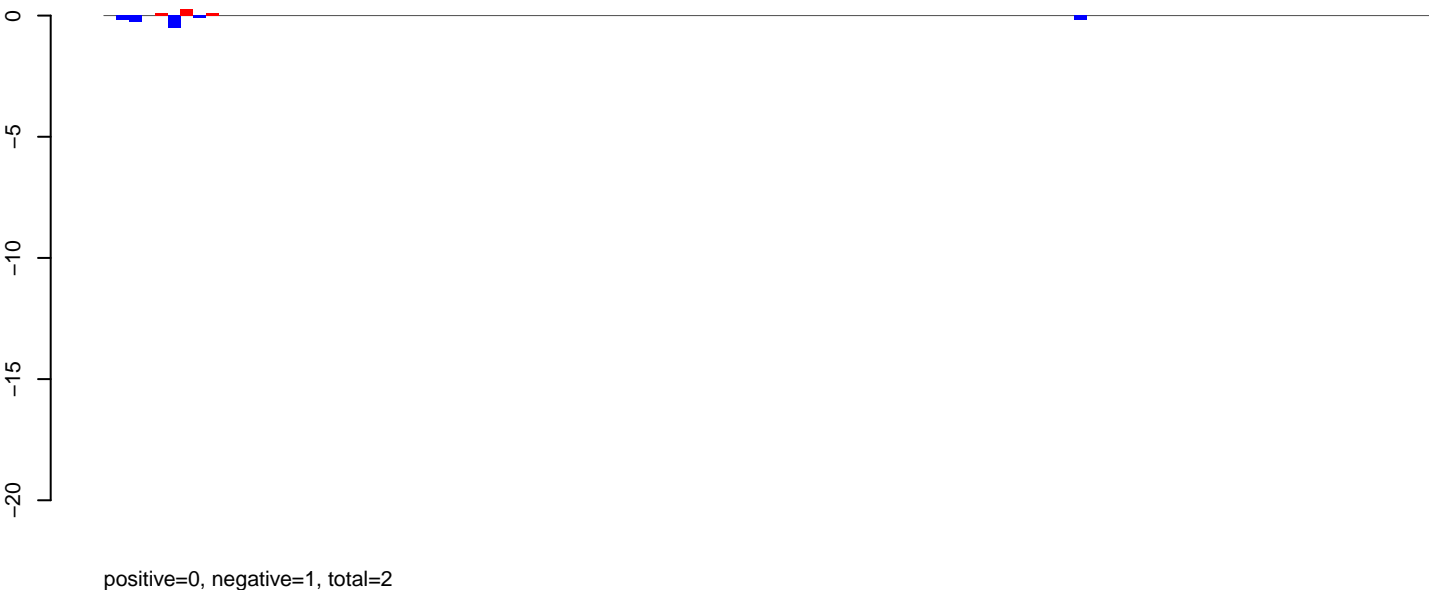
AnGam_Mos55_cells.24_35.rep



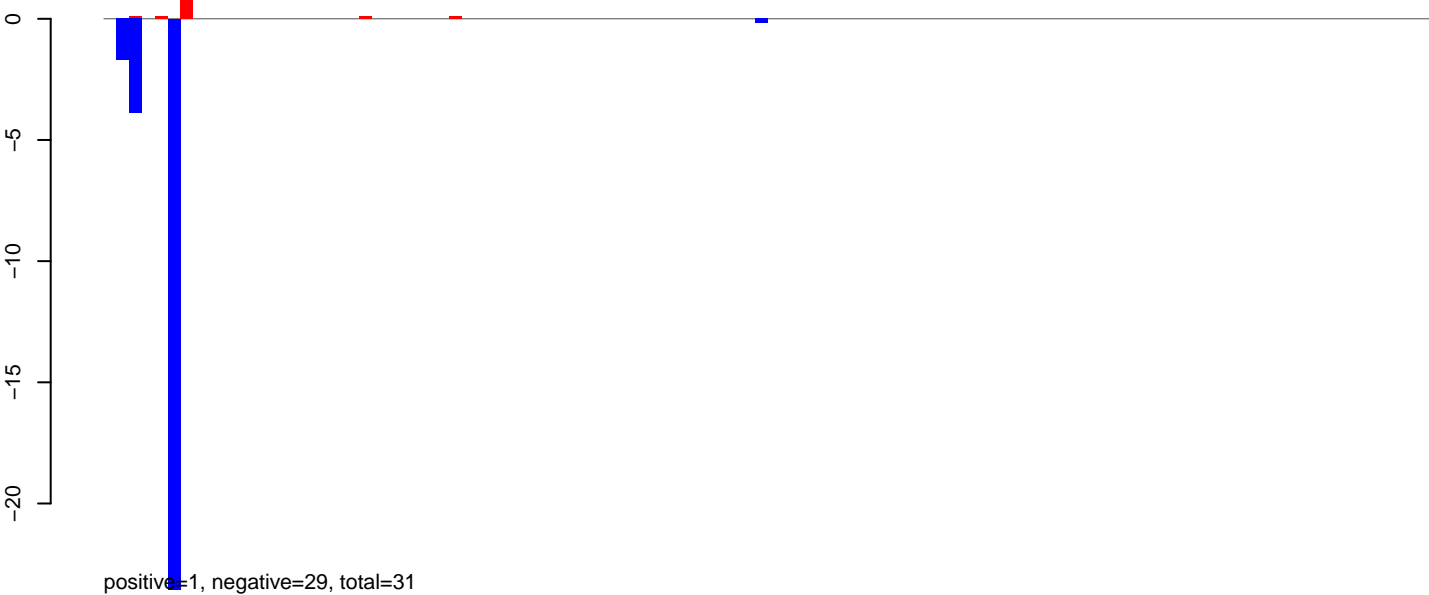
AnGam_Mos55_cells.rep



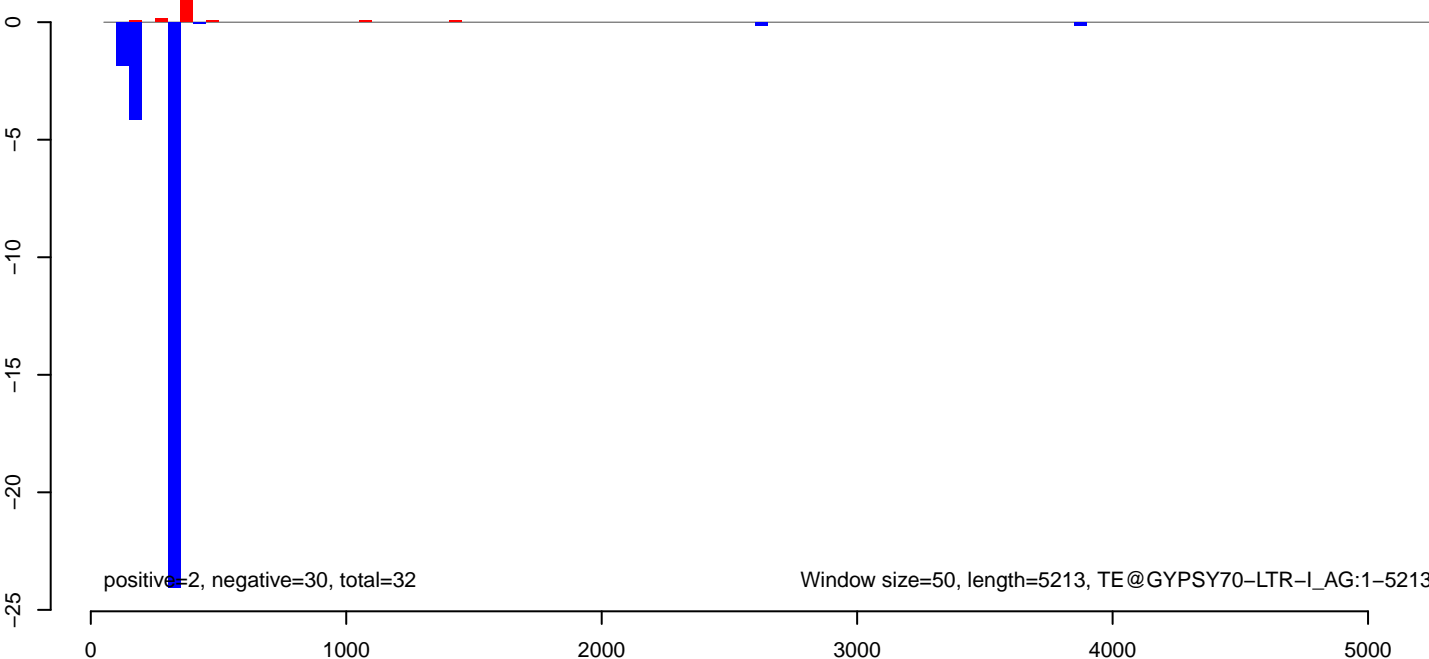
AnGam_Mos55_cells.18_23.rep



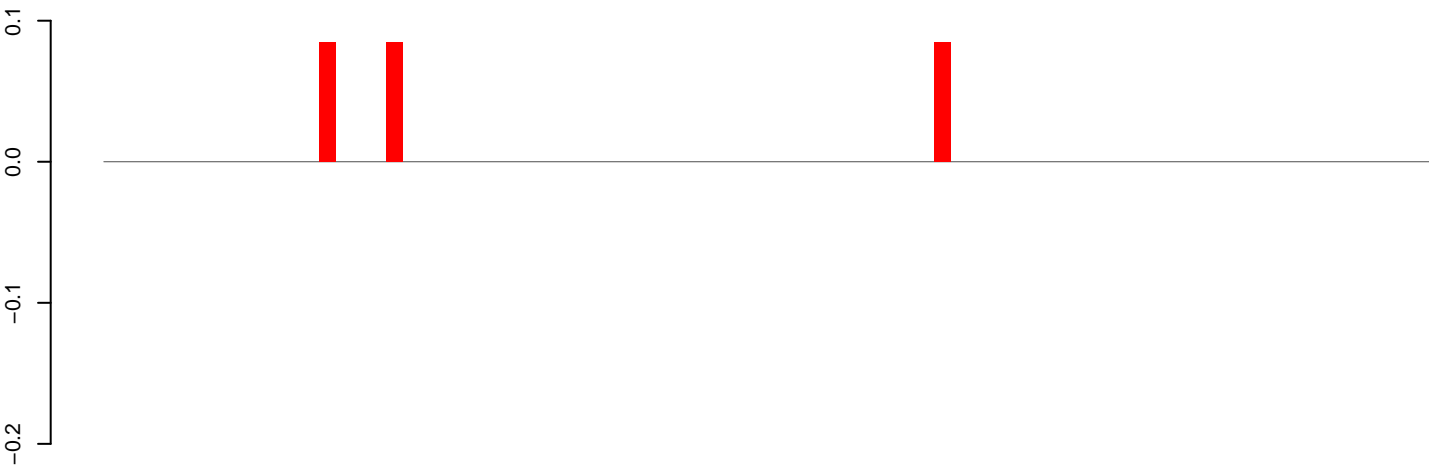
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

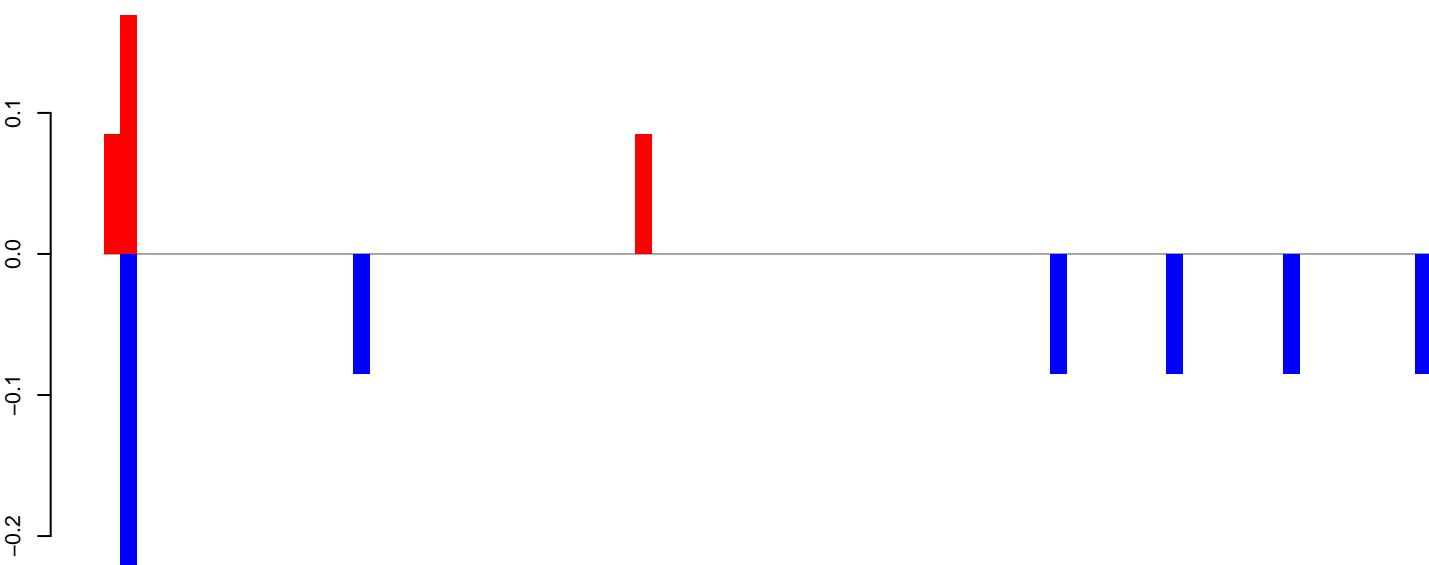


AnGam_Mos55_cells.18_23.rep



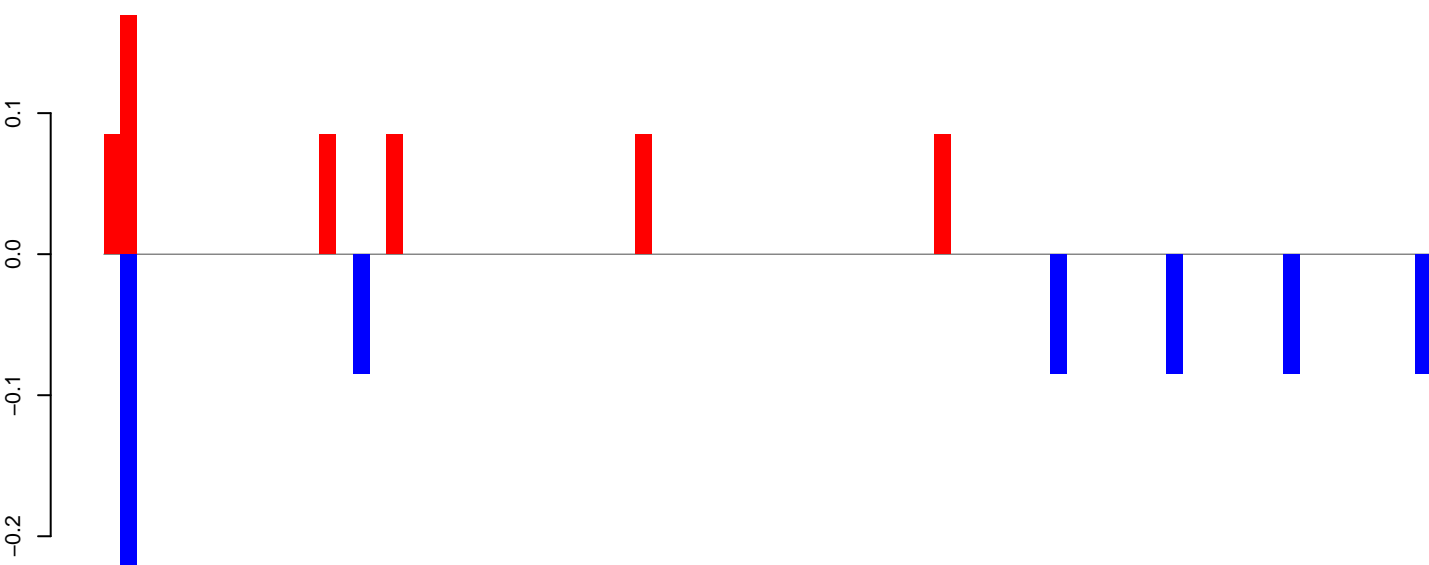
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=1, total=1

AnGam_Mos55_cells.rep



positive=1, negative=1, total=1

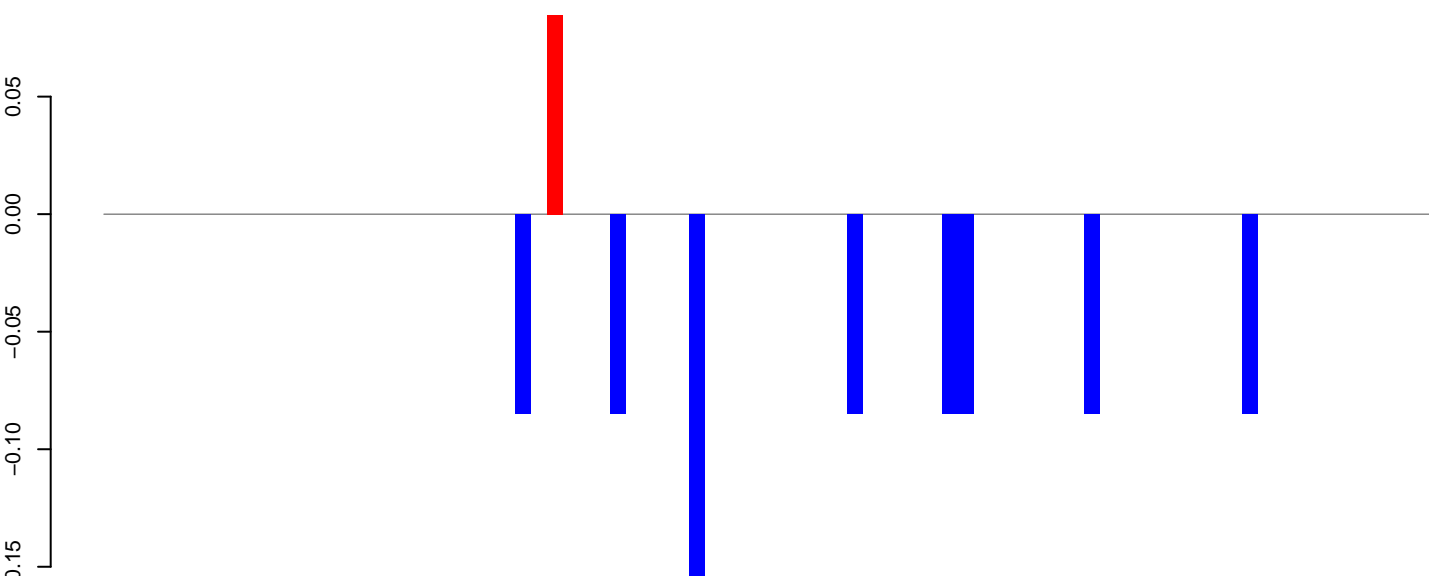
Window size=50, length=4031, TE@GYPSY1-LTR-L_AG:1-4031

AnGam_Mos55_cells.18_23.rep



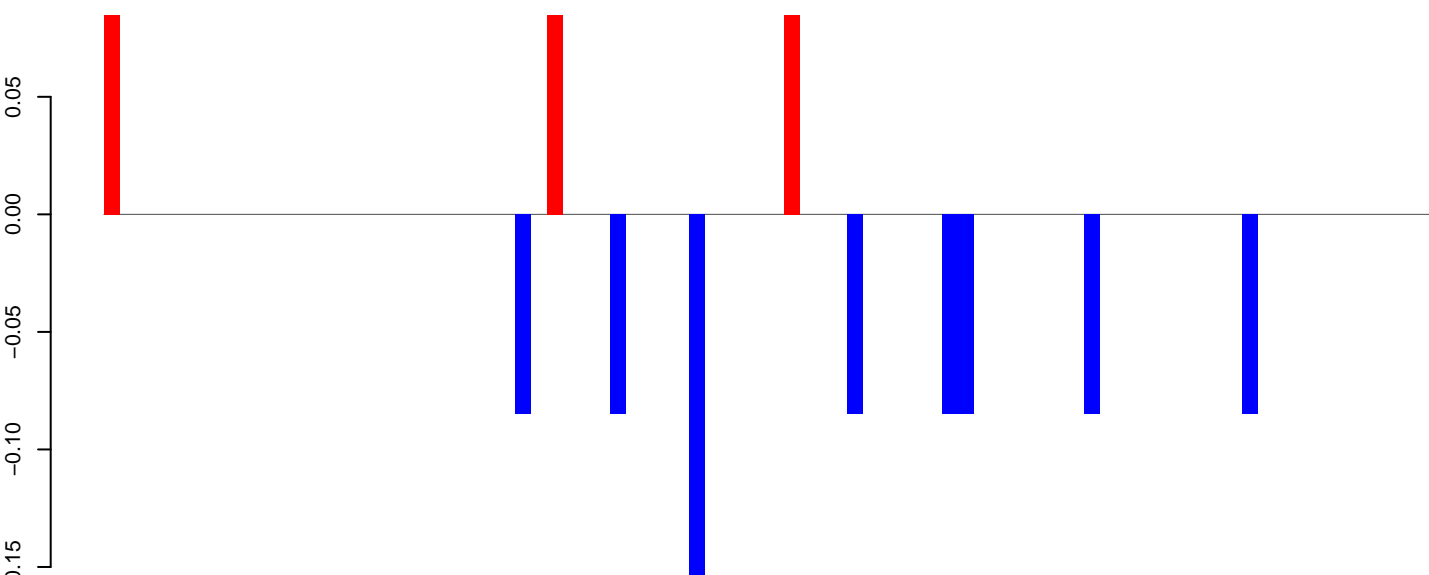
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=1, total=1

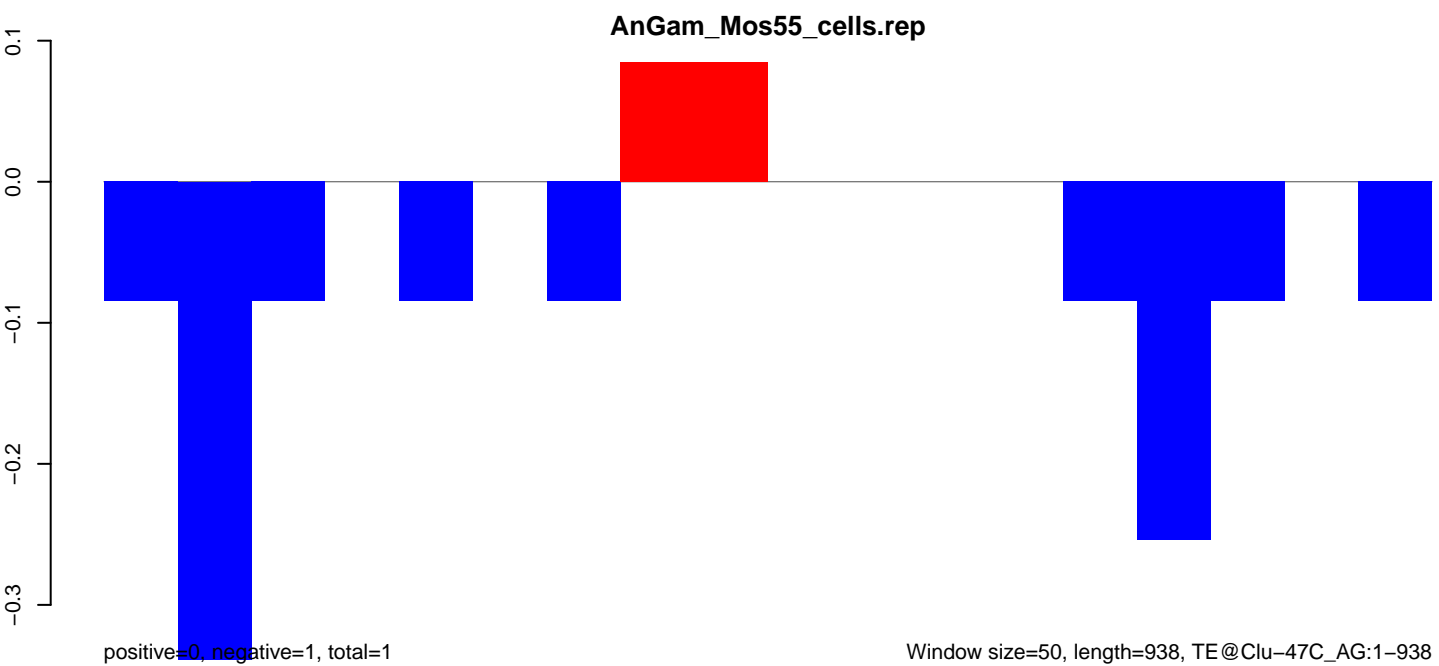
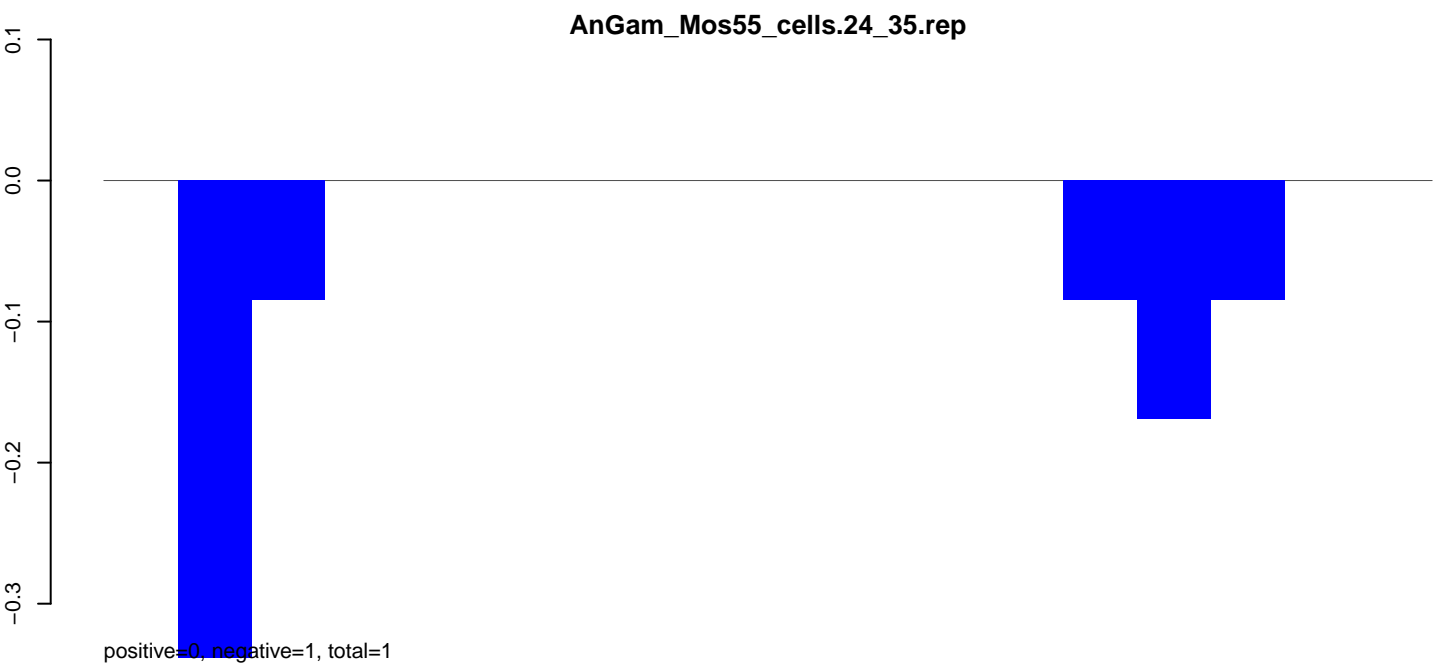
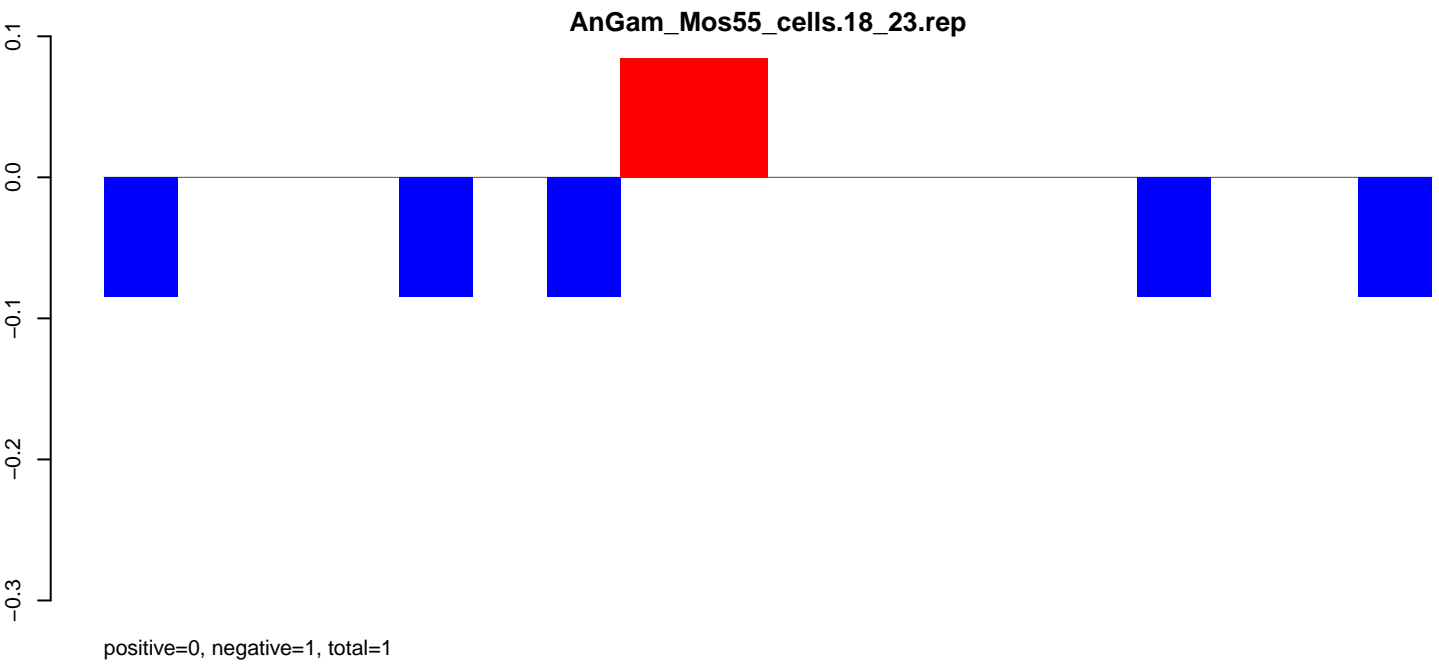
AnGam_Mos55_cells.rep



positive=0, negative=1, total=1

Window size=50, length=4247, TE@Copia-6_AG-LTR-I:1-4247

0 1000 2000 3000 4000

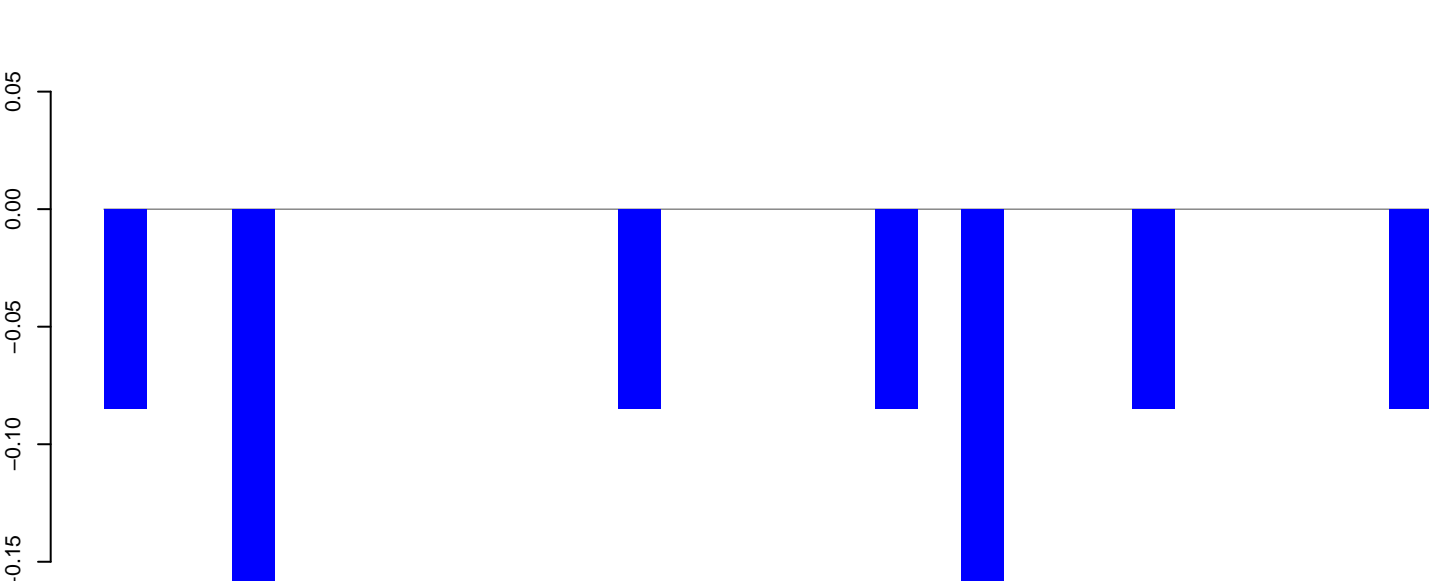


AnGam_Mos55_cells.18_23.rep



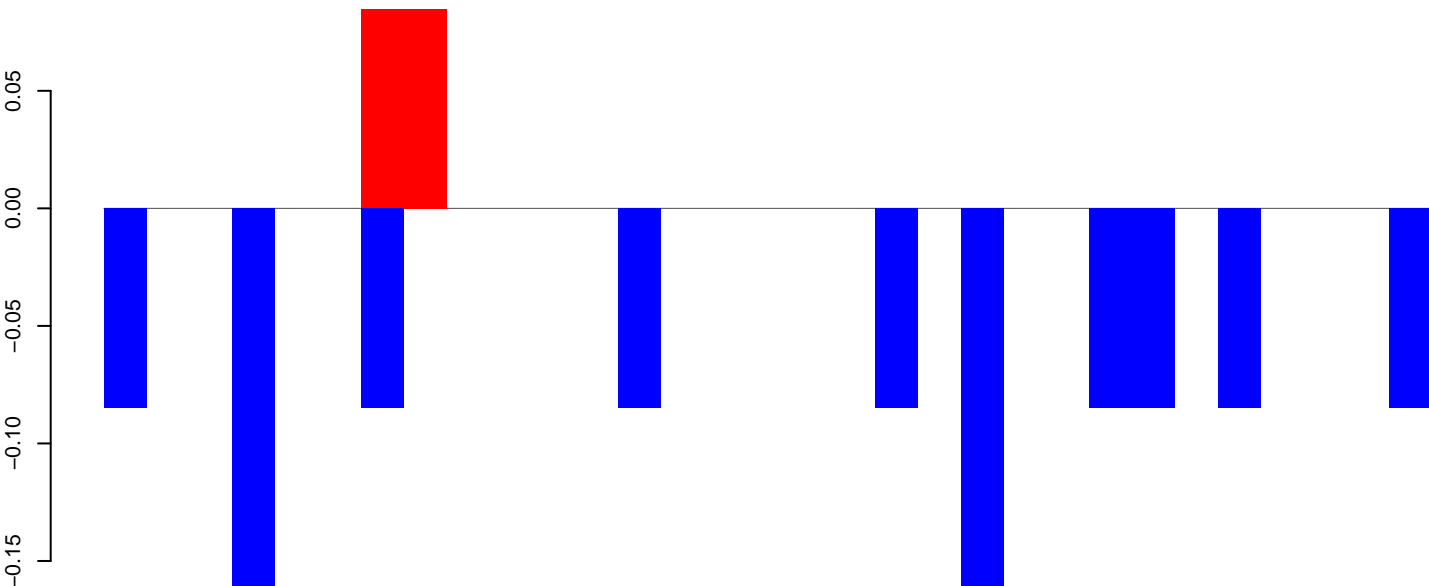
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=1, total=1

AnGam_Mos55_cells.rep



positive=0, negative=1, total=1

Window size=50, length=1558, TE@Clu-240_AG:1-1558

0 500 1000 1500

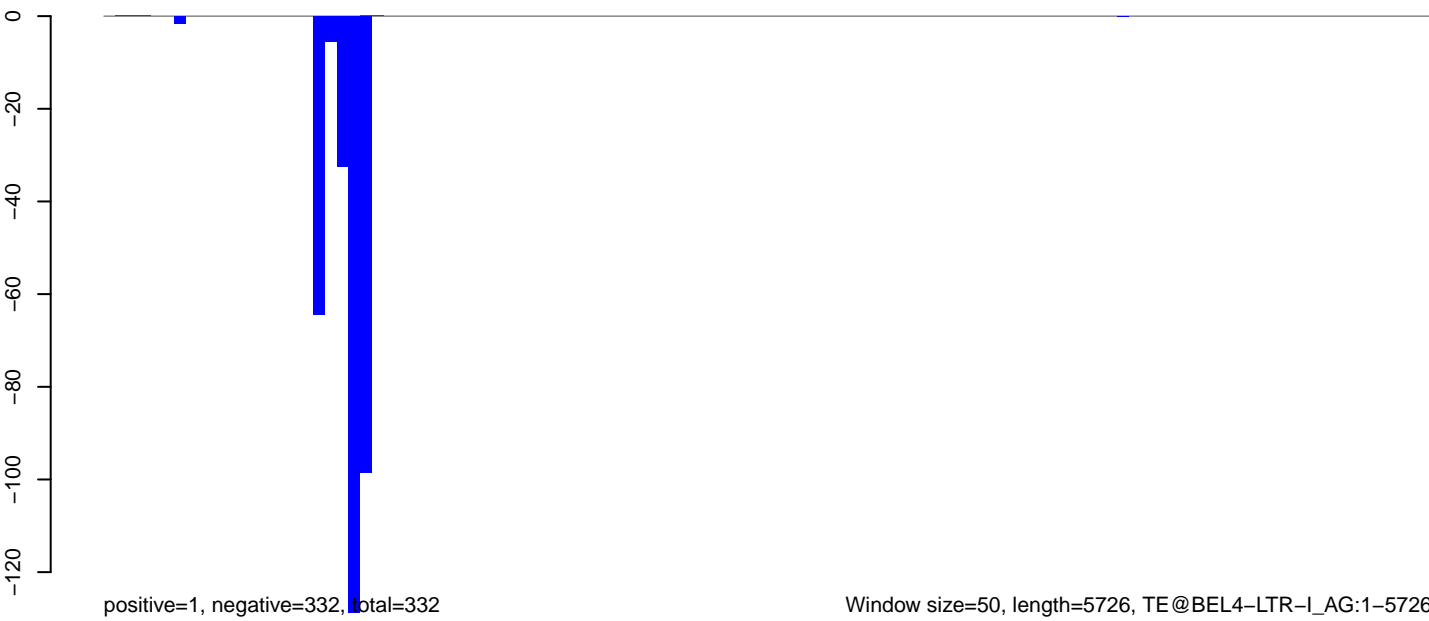
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



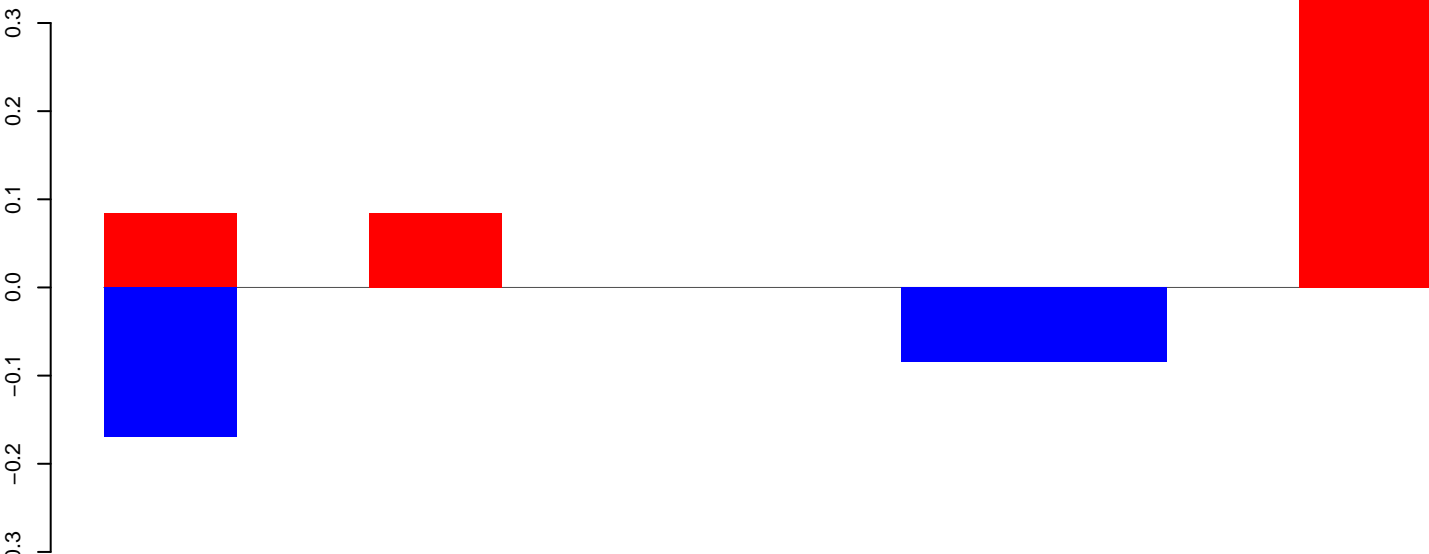
AnGam_Mos55_cells.rep



Window size=50, length=5726, TE@BEL4-LTR-I_AG:1-5726

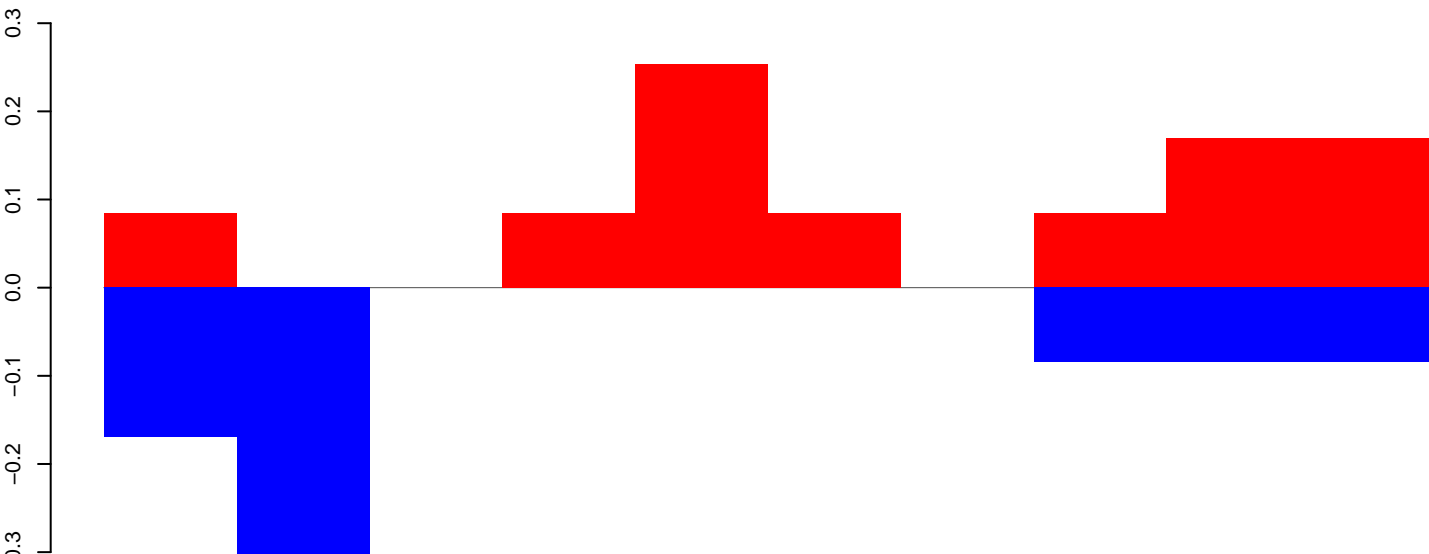
0 1000 2000 3000 4000 5000

AnGam_Mos55_cells.18_23.rep



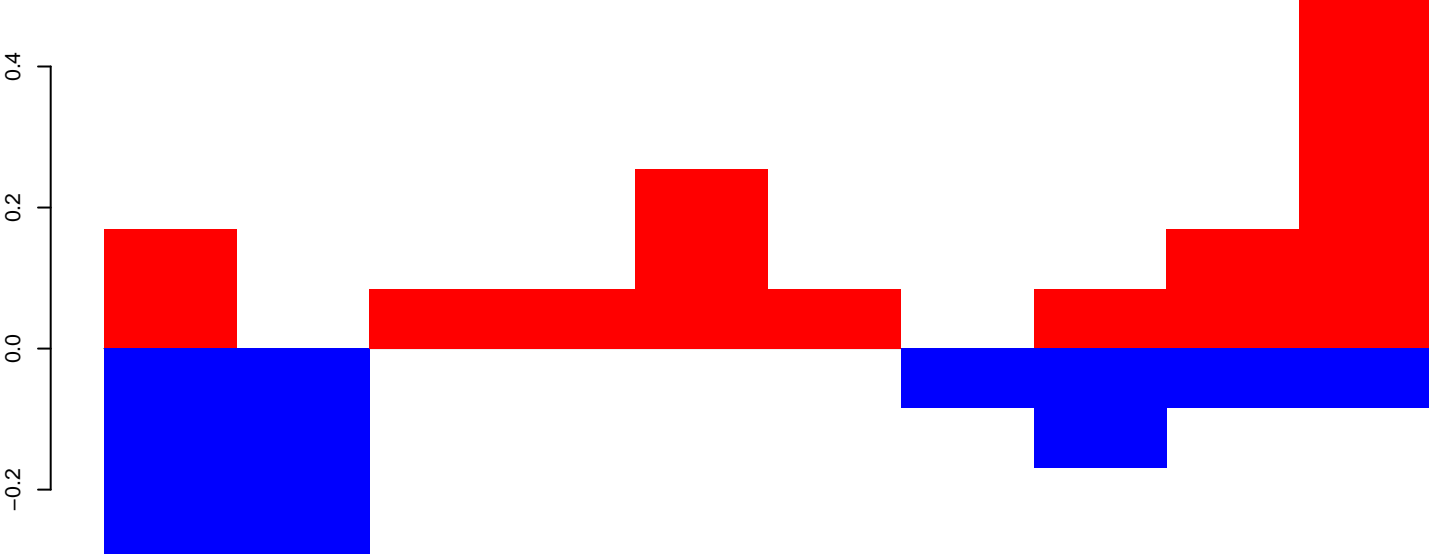
positive=1, negative=0, total=1

AnGam_Mos55_cells.24_35.rep



positive=1, negative=1, total=2

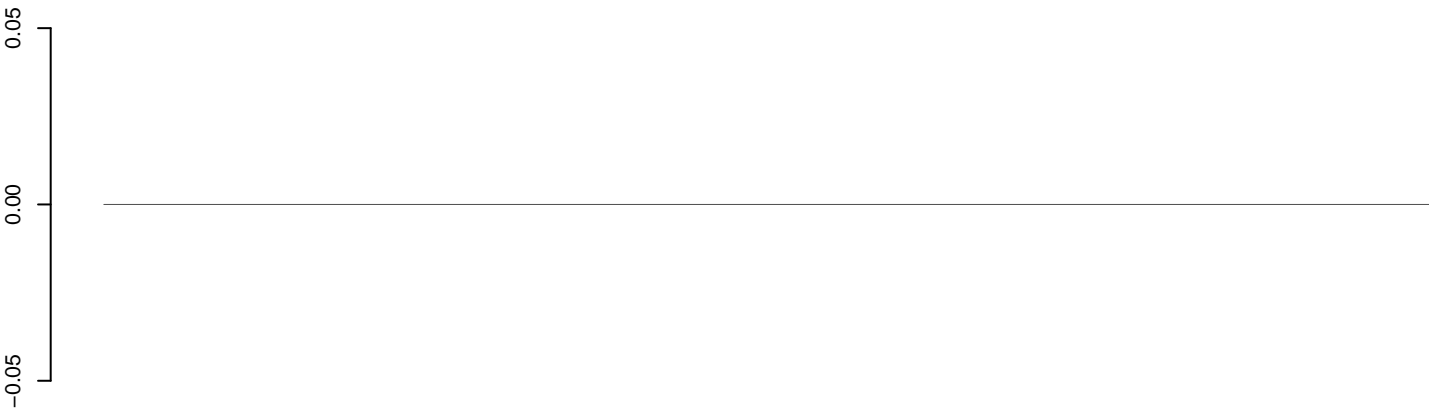
AnGam_Mos55_cells.rep



positive=2, negative=1, total=3

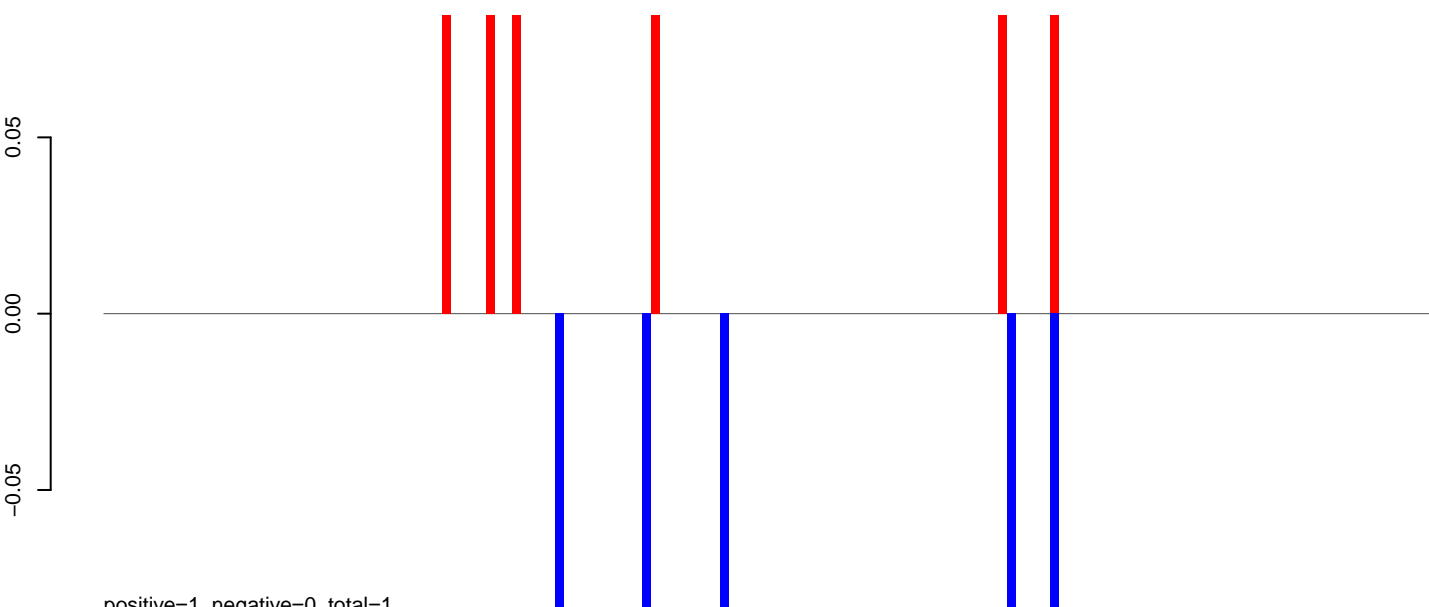
Window size=50, length=528, TE@TF003653-m4bp_Ele1:1-528

AnGam_Mos55_cells.18_23.rep



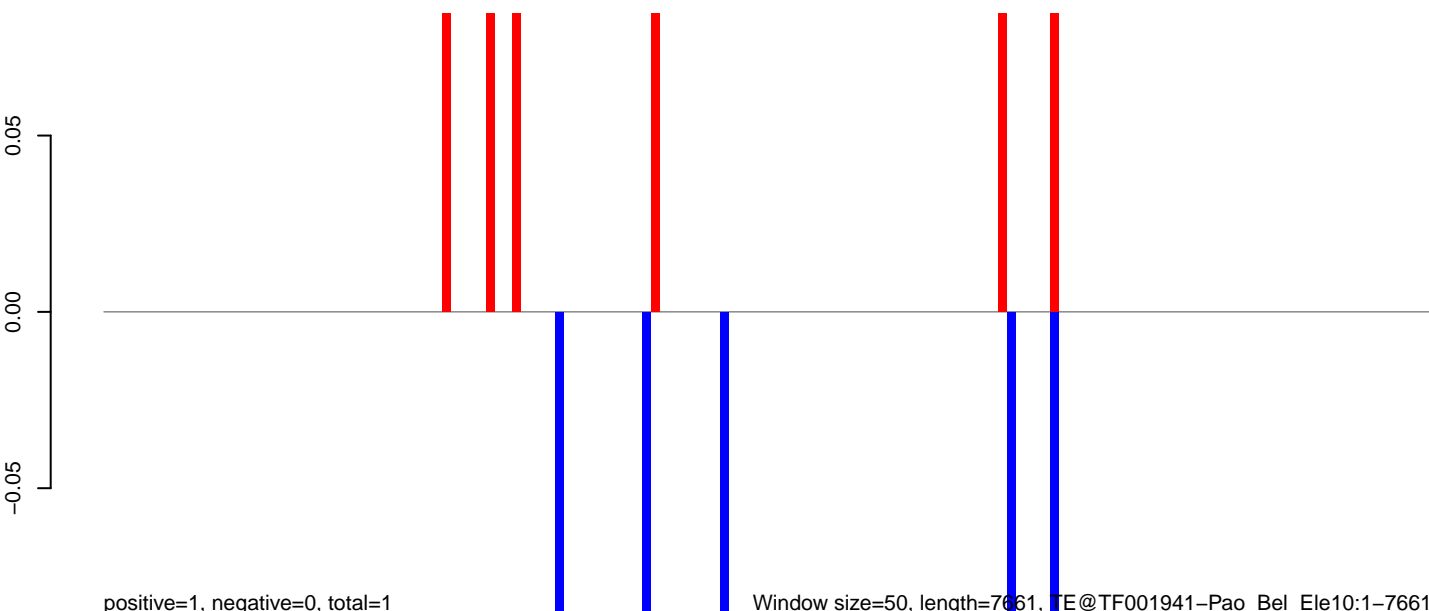
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=1, negative=0, total=1

AnGam_Mos55_cells.rep

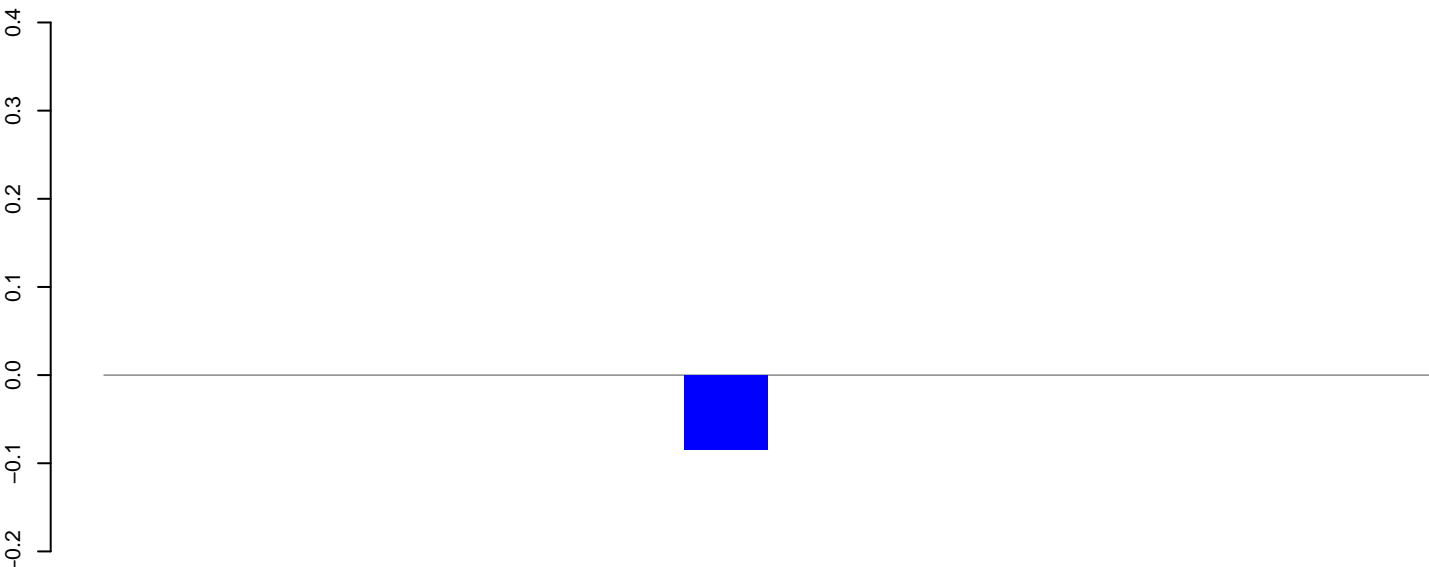


positive=1, negative=0, total=1

Window size=50, length=7661, TE@TF001941-Pao_Bel_Ele10:1-7661

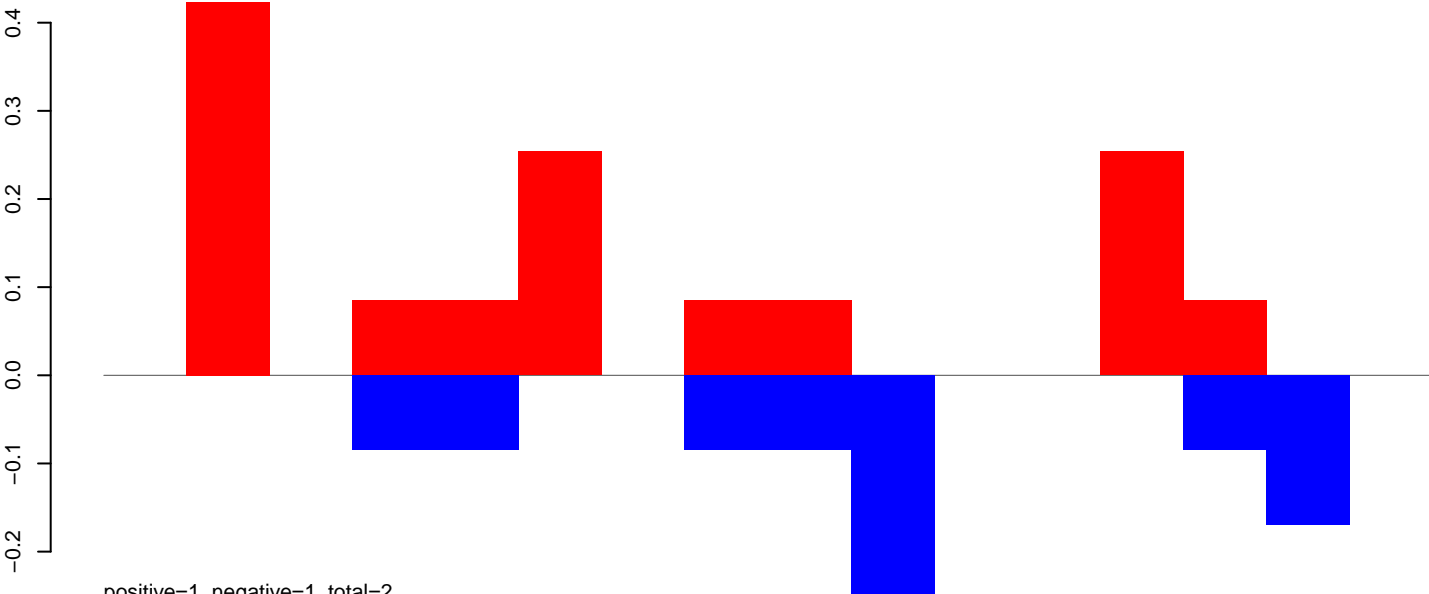
0 2000 4000 6000 8000

AnGam_Mos55_cells.18_23.rep



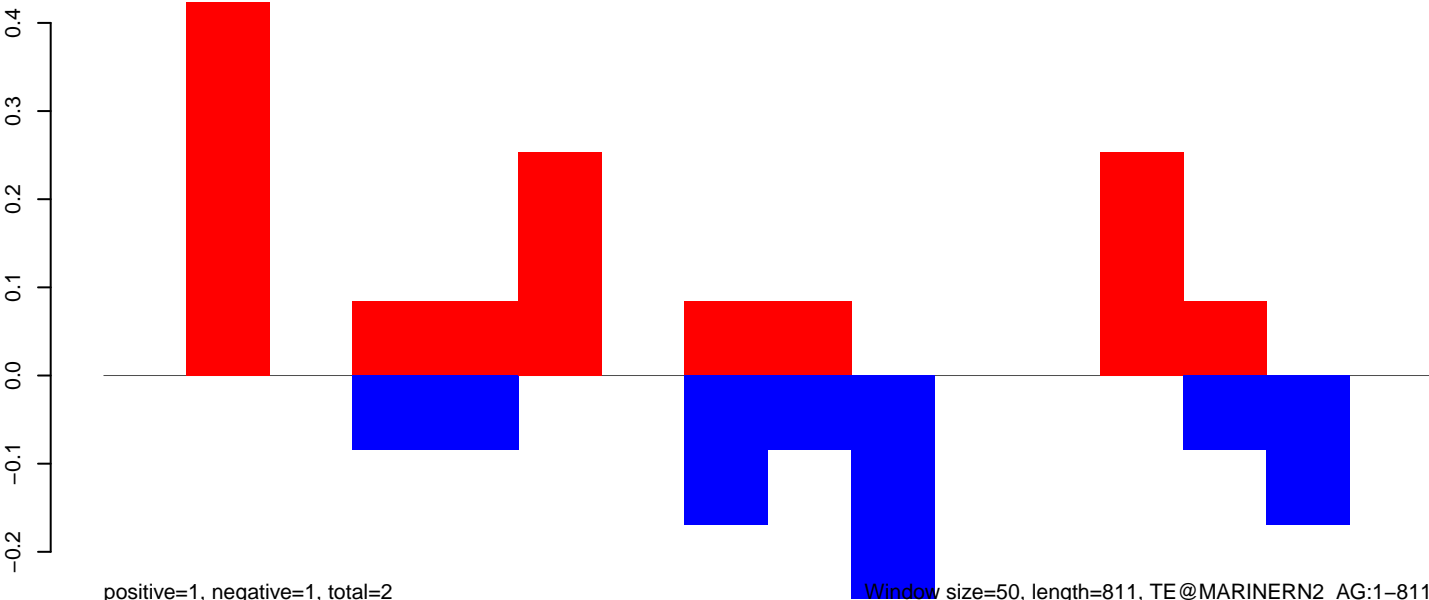
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=1, negative=1, total=2

AnGam_Mos55_cells.rep



positive=1, negative=1, total=2

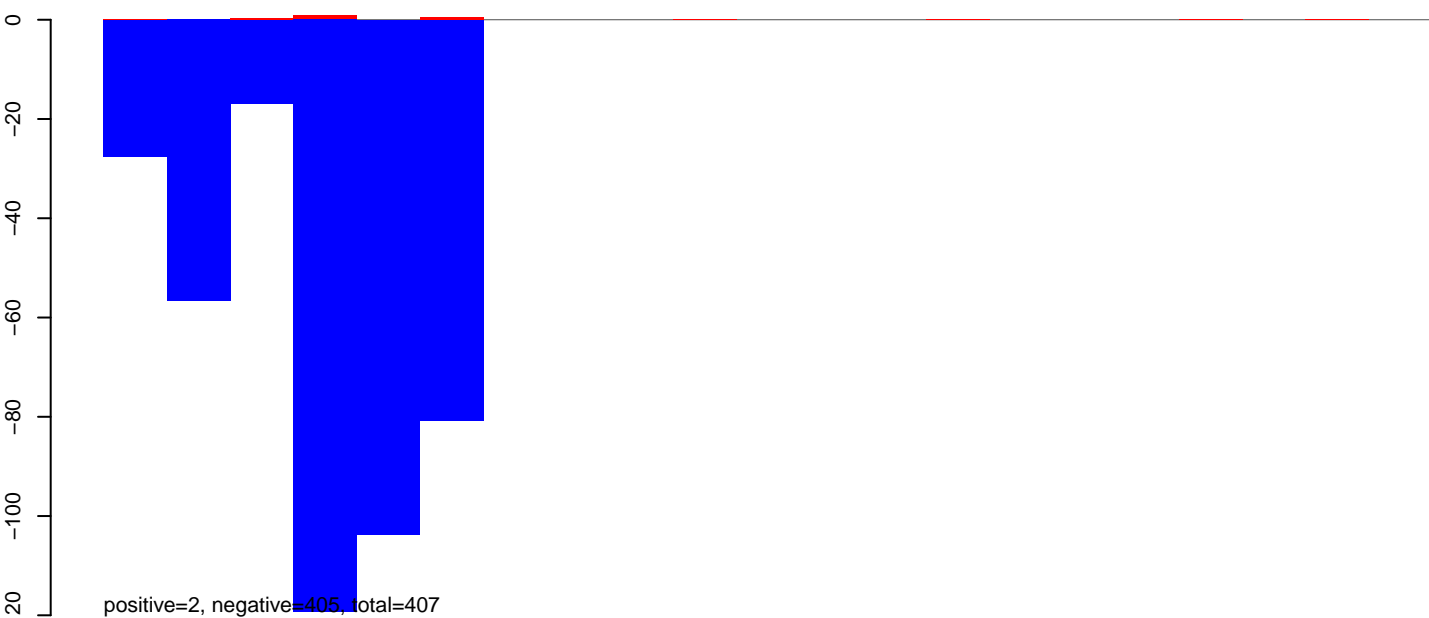
Window size=50, length=811, TE@MARINERN2_AG:1-811

200 400 600 800

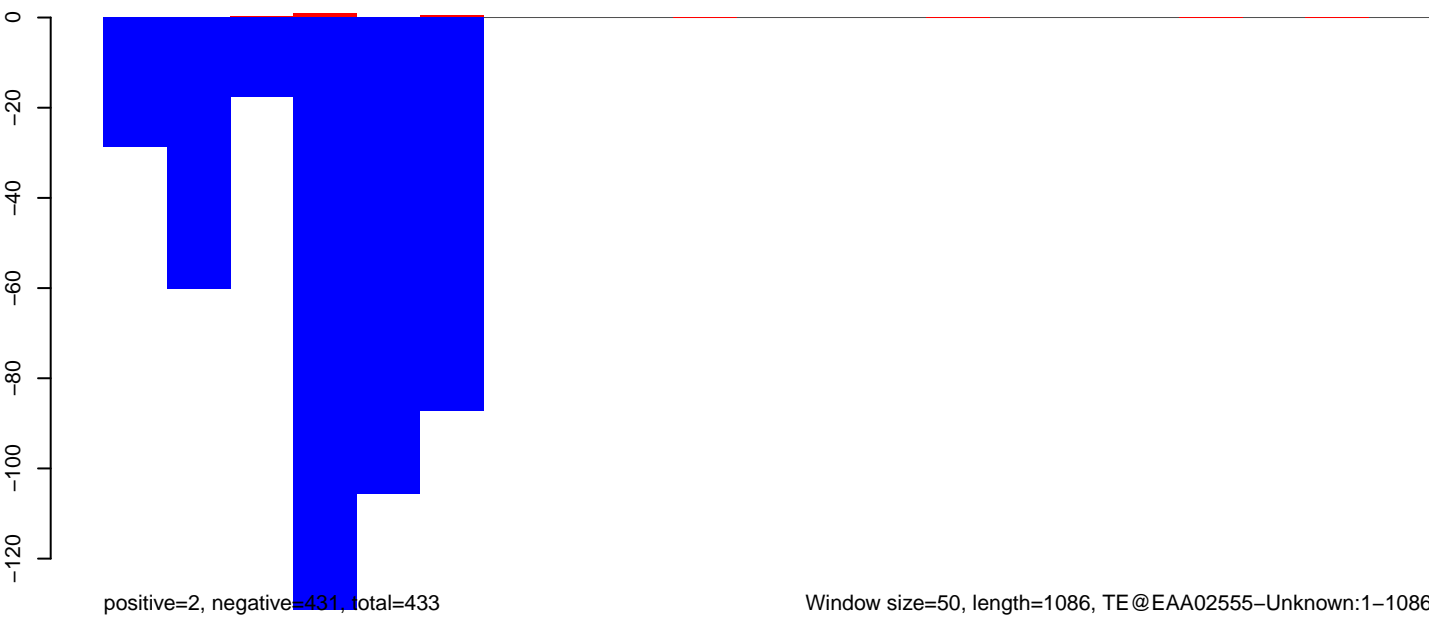
AnGam_Mos55_cells.18_23.rep



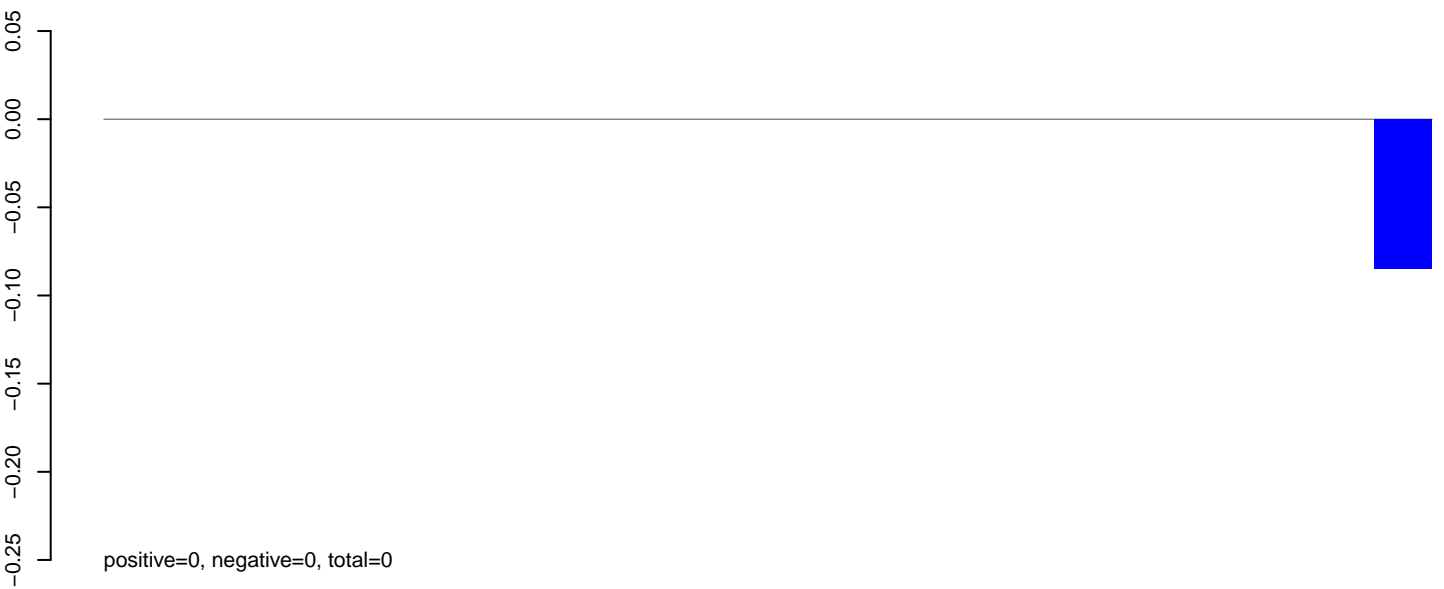
AnGam_Mos55_cells.24_35.rep



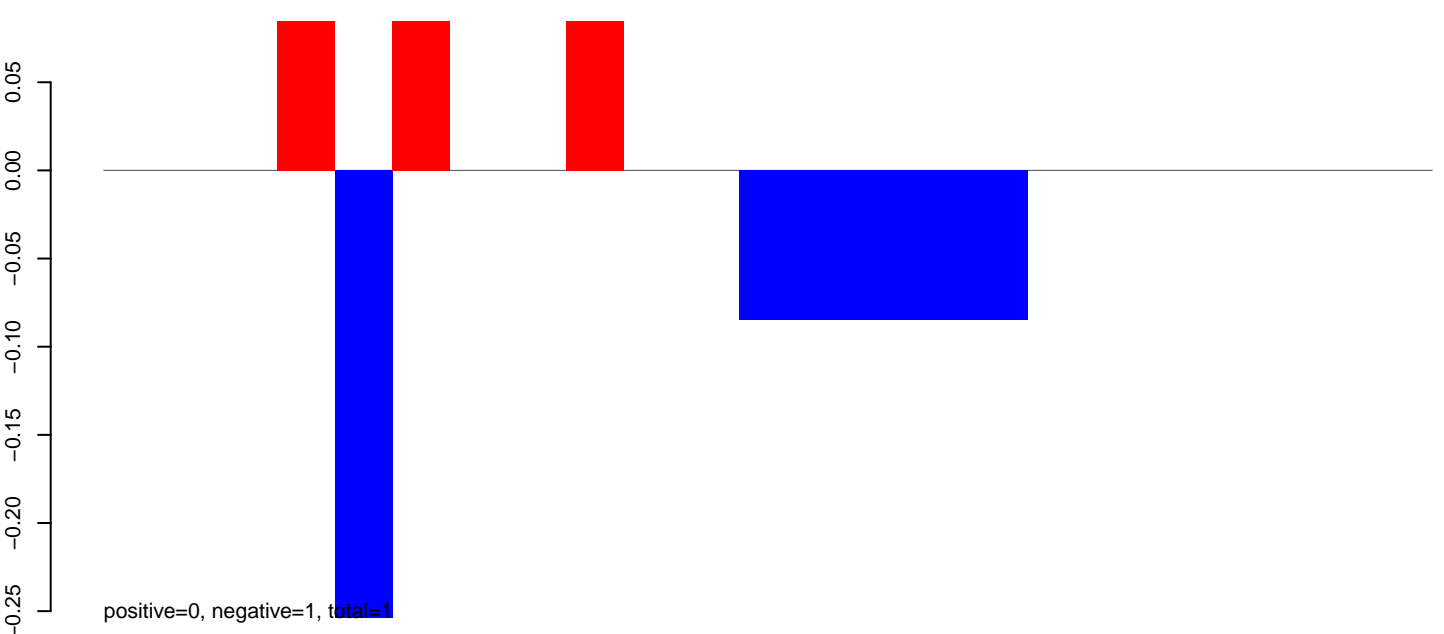
AnGam_Mos55_cells.rep



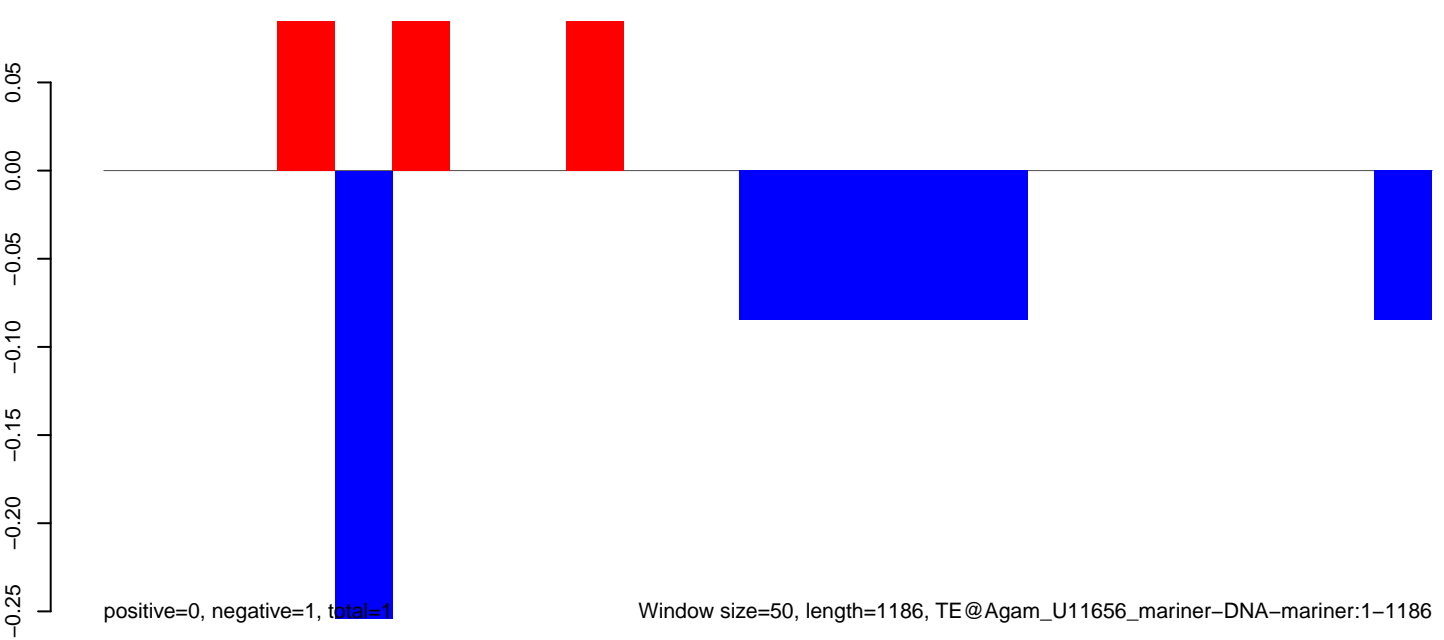
AnGam_Mos55_cells.18_23.rep



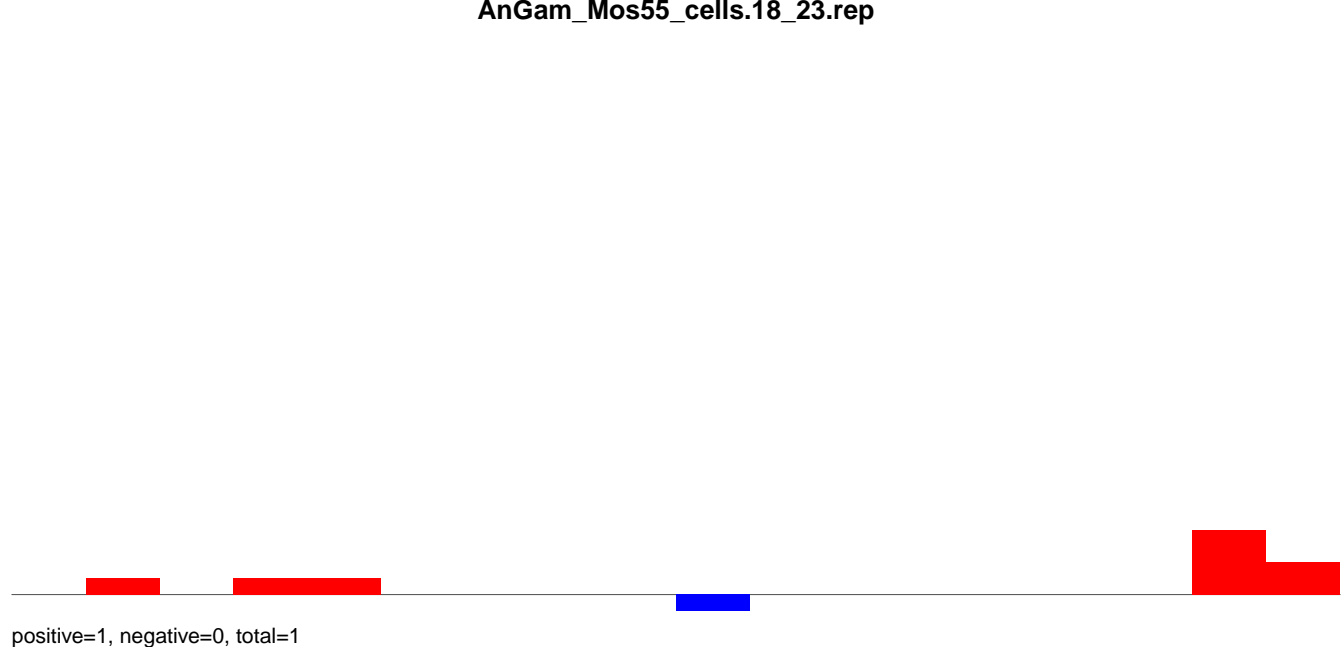
AnGam_Mos55_cells.24_35.rep



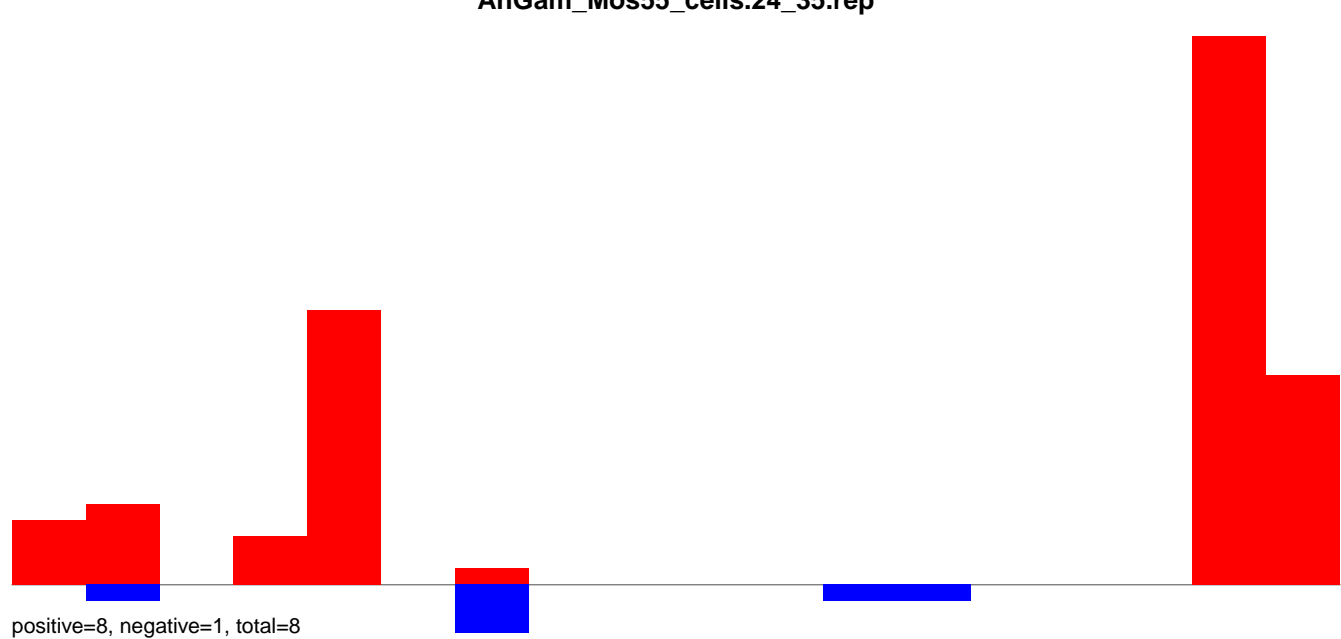
AnGam_Mos55_cells.rep



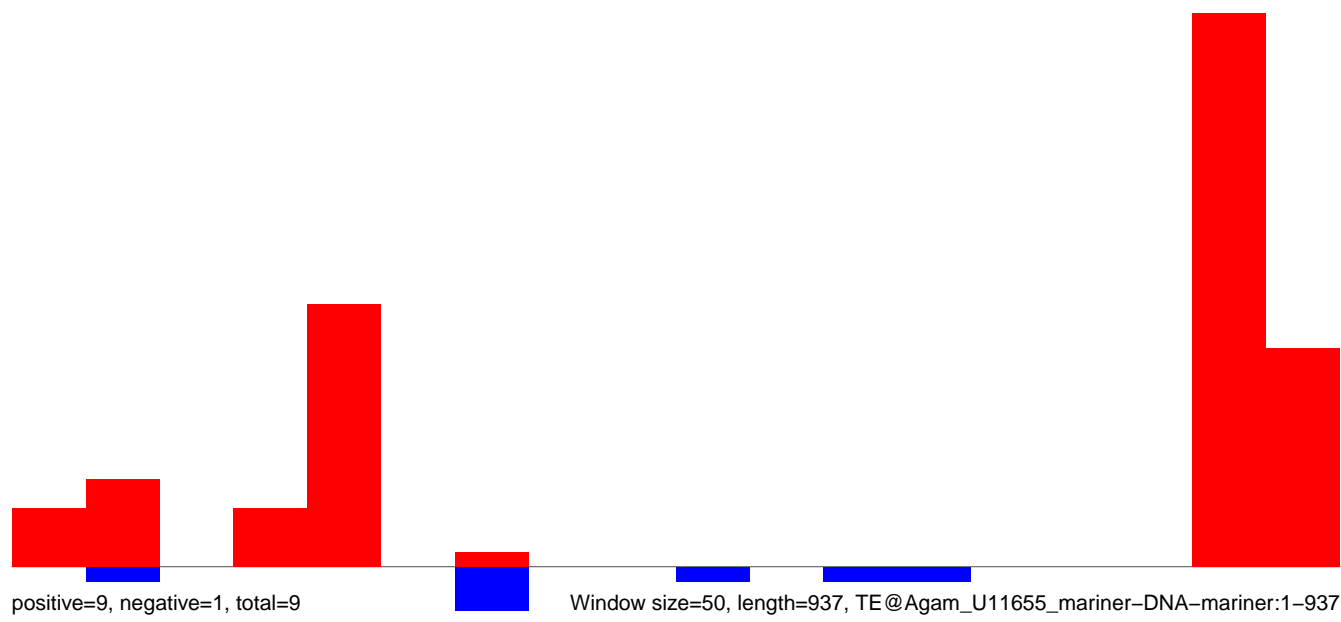
AnGam_Mos55_cells.18_23.rep



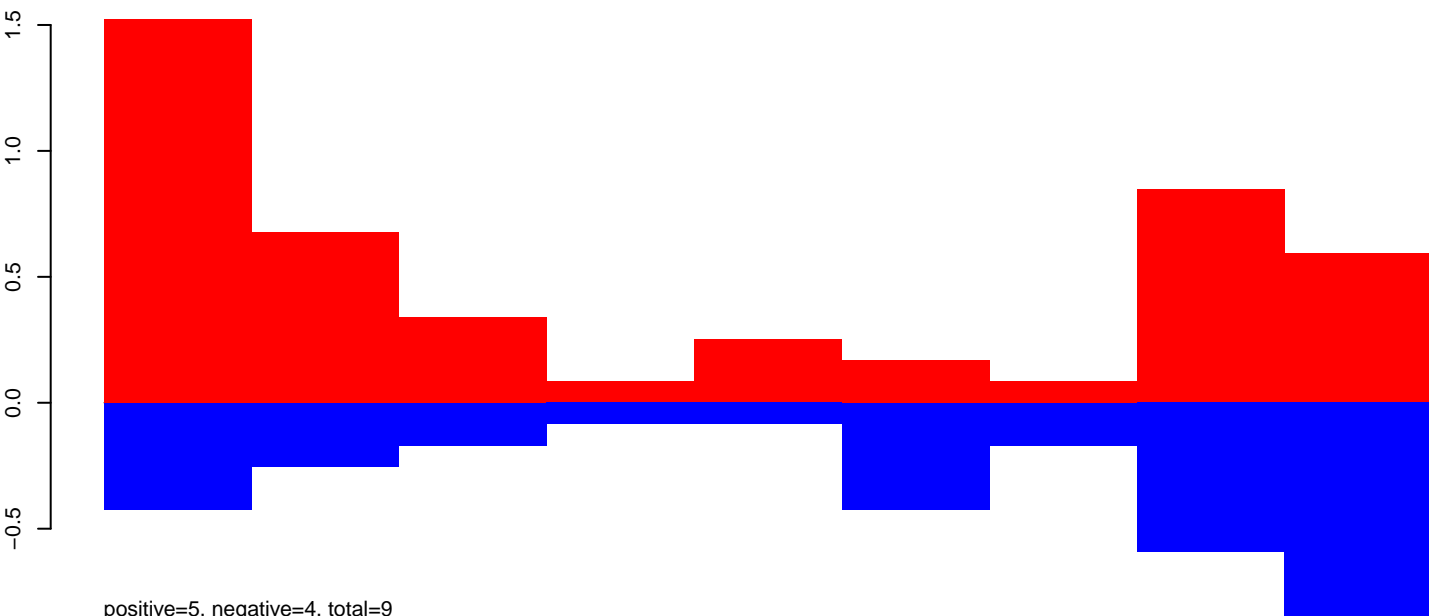
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



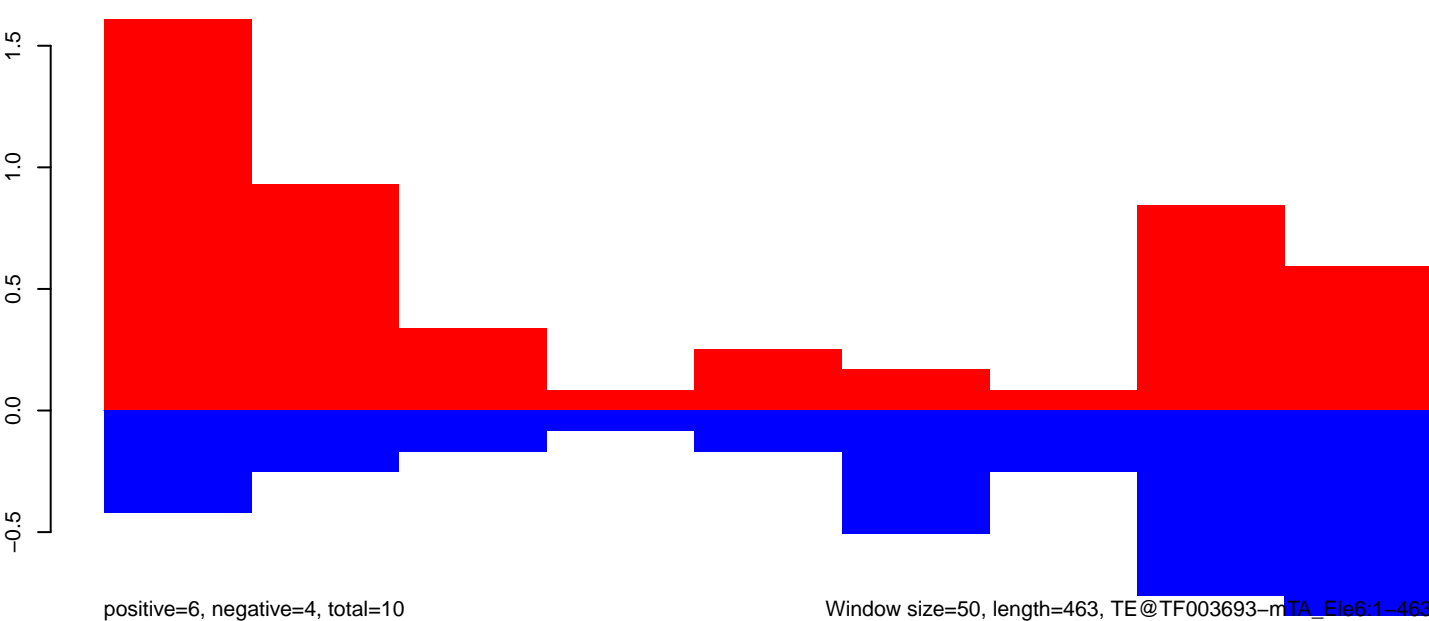
AnGam_Mos55_cells.18_23.rep



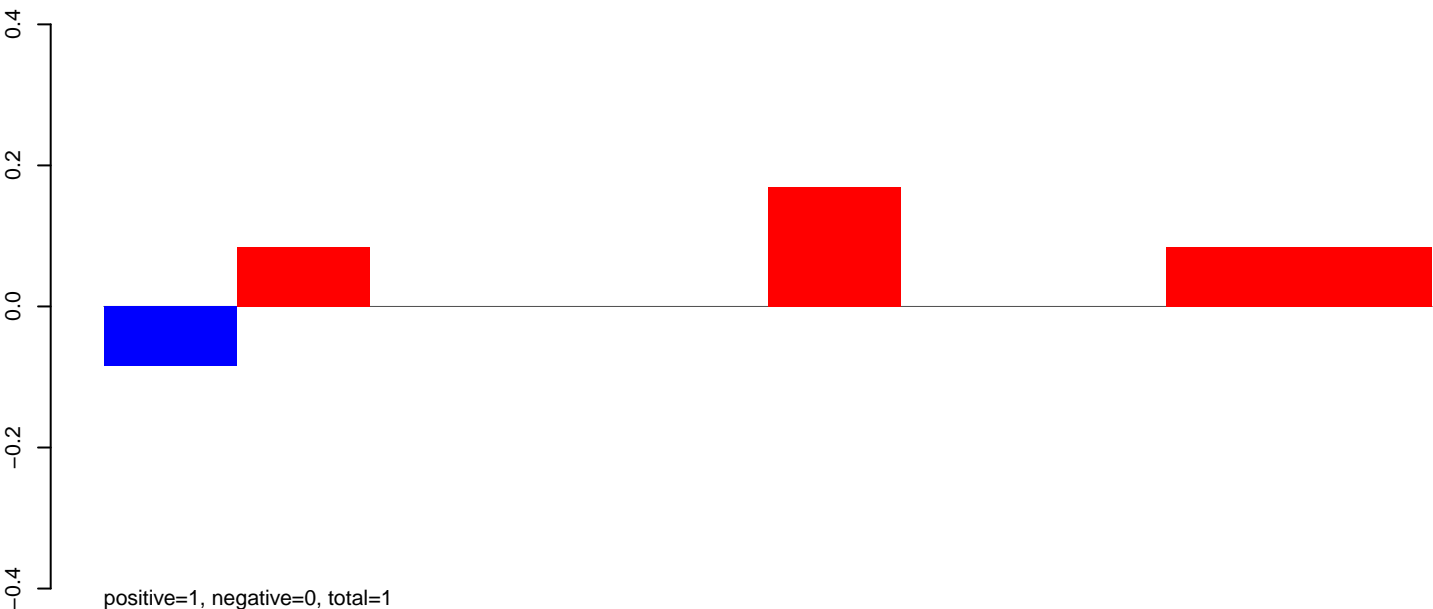
AnGam_Mos55_cells.24_35.rep



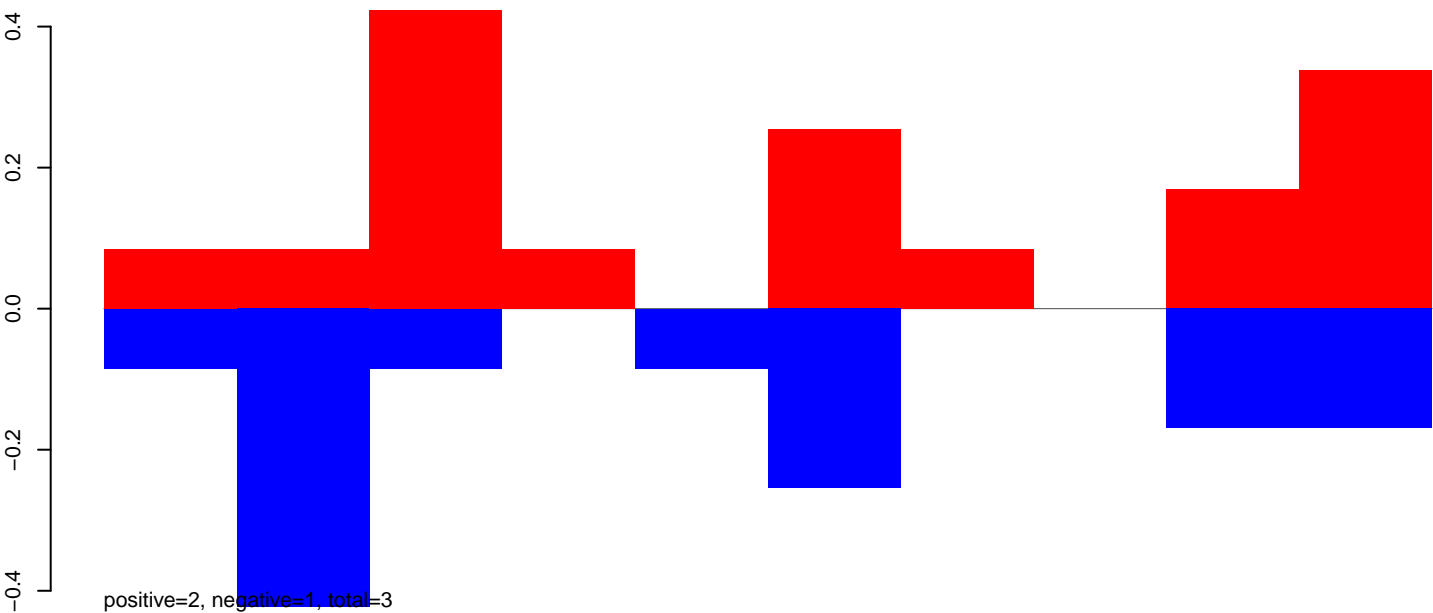
AnGam_Mos55_cells.rep



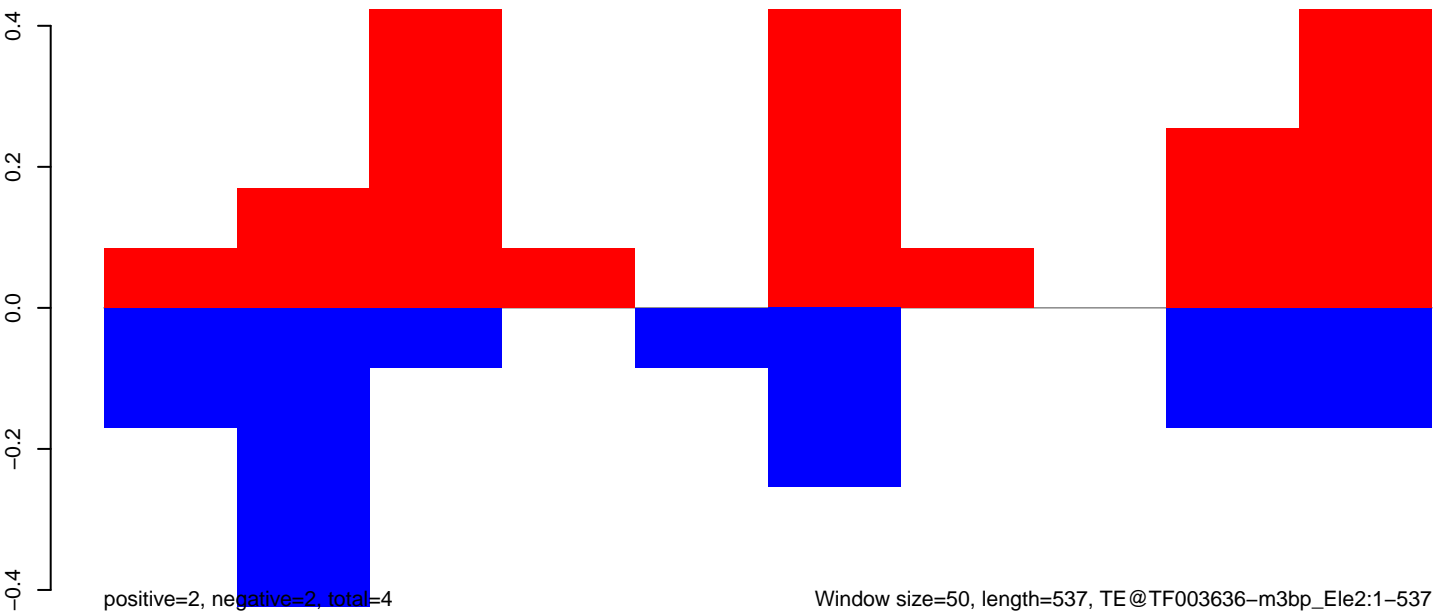
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



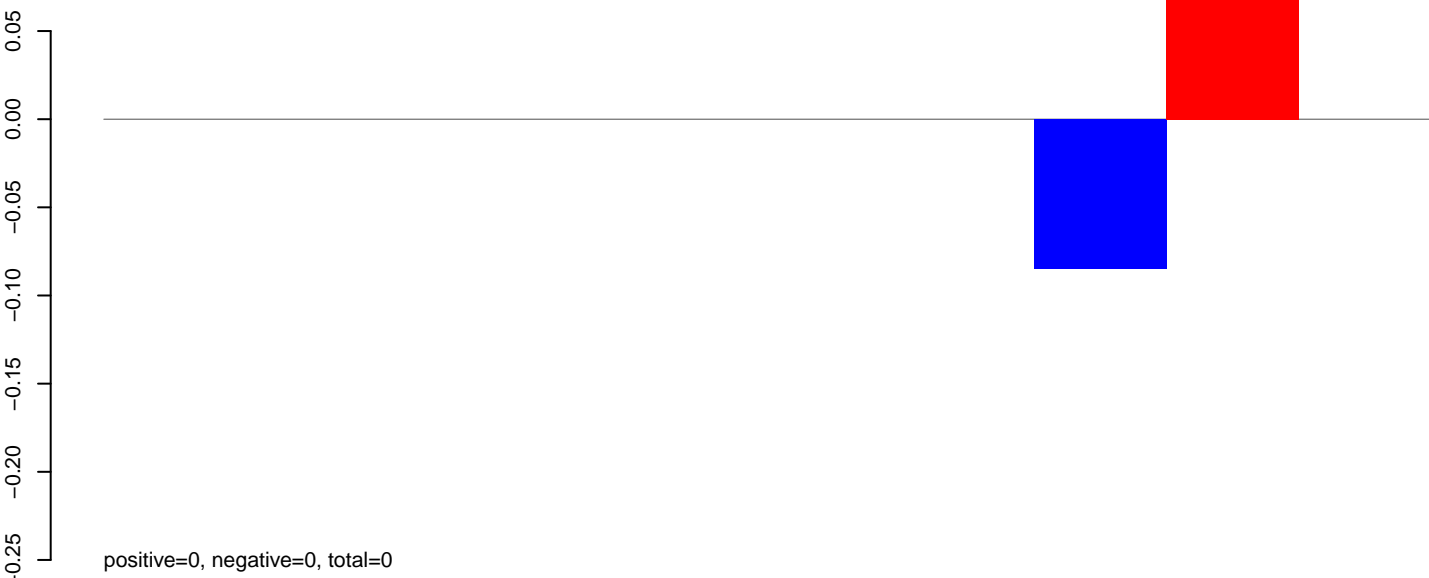
AnGam_Mos55_cells.rep



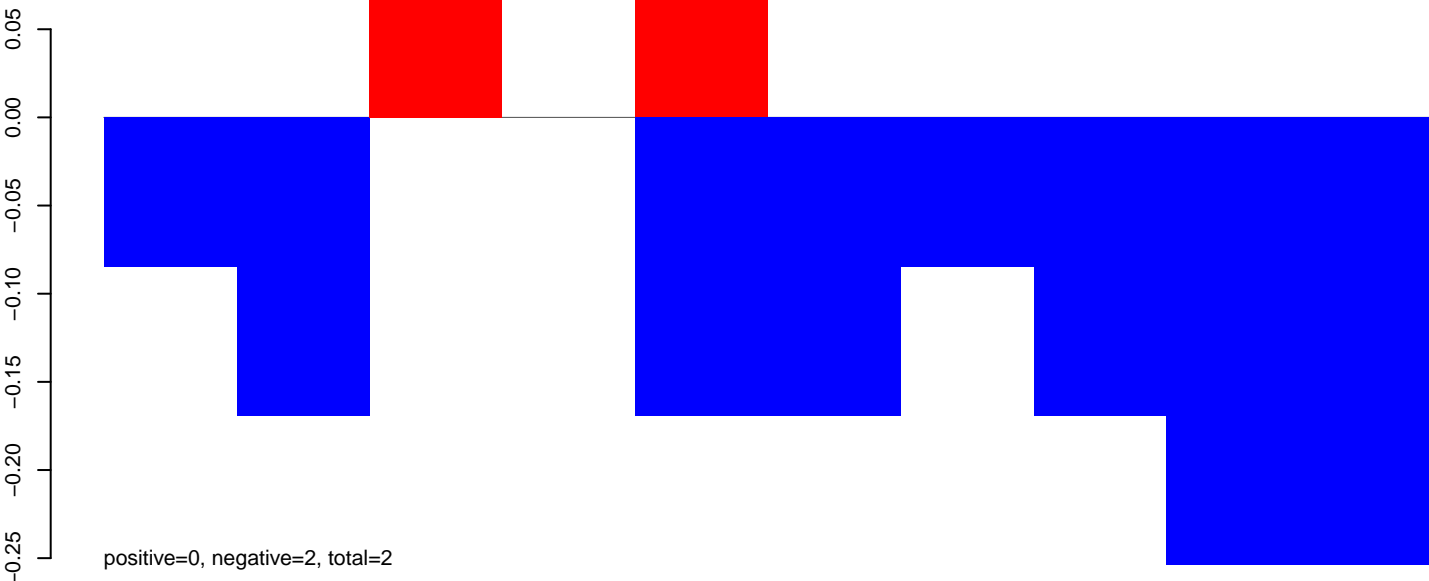
Window size=50, length=537, TE@TF003636-m3bp_Ele2:1-537

100 200 300 400 500

AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



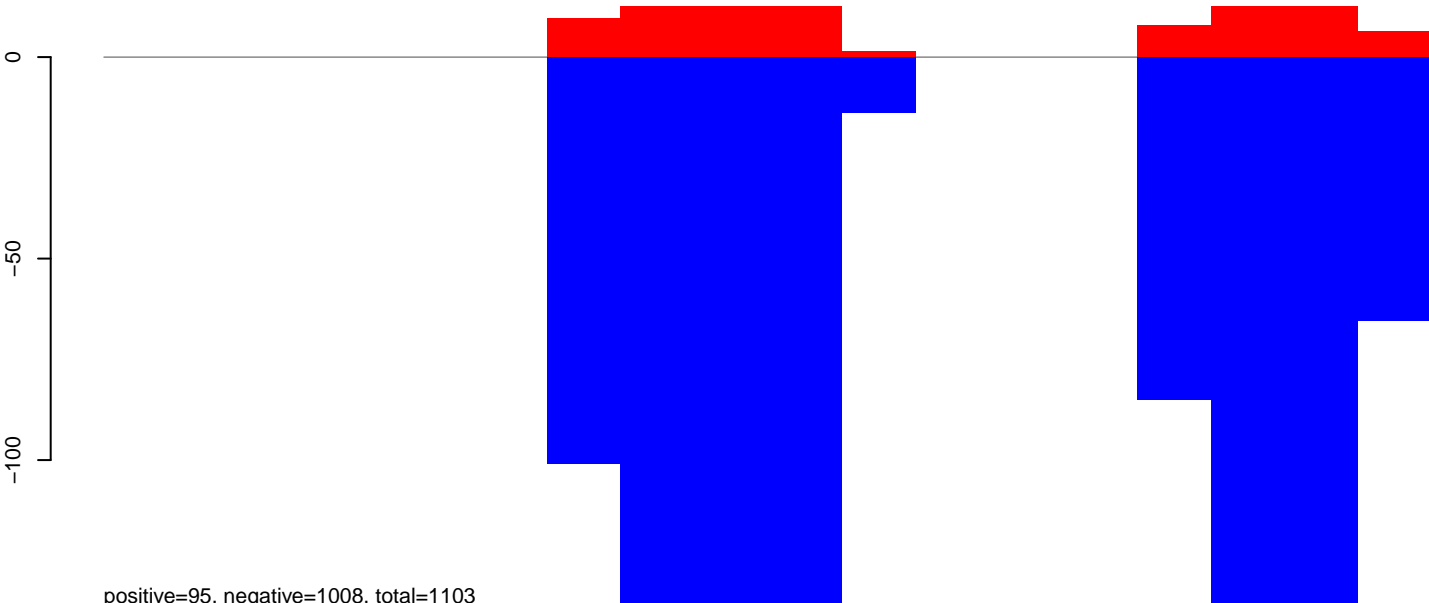
AnGam_Mos55_cells.rep



Window size=50, length=546, TE@TF003620-m2bp_Ele11:1-546

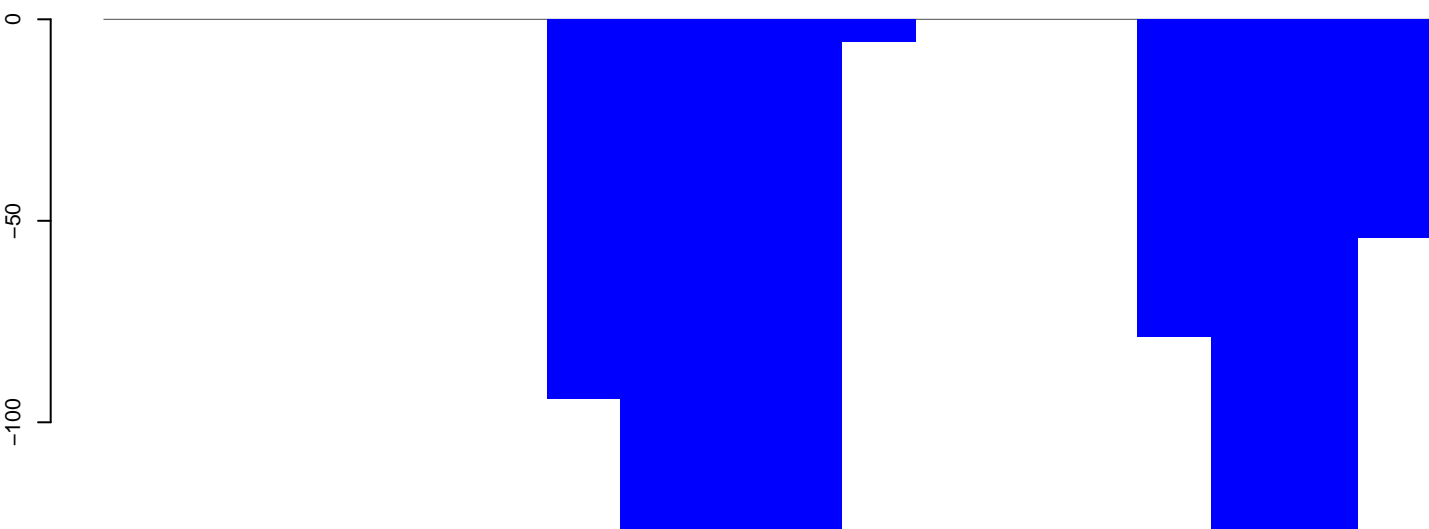
100 200 300 400 500

AnGam_Mos55_cells.18_23.rep



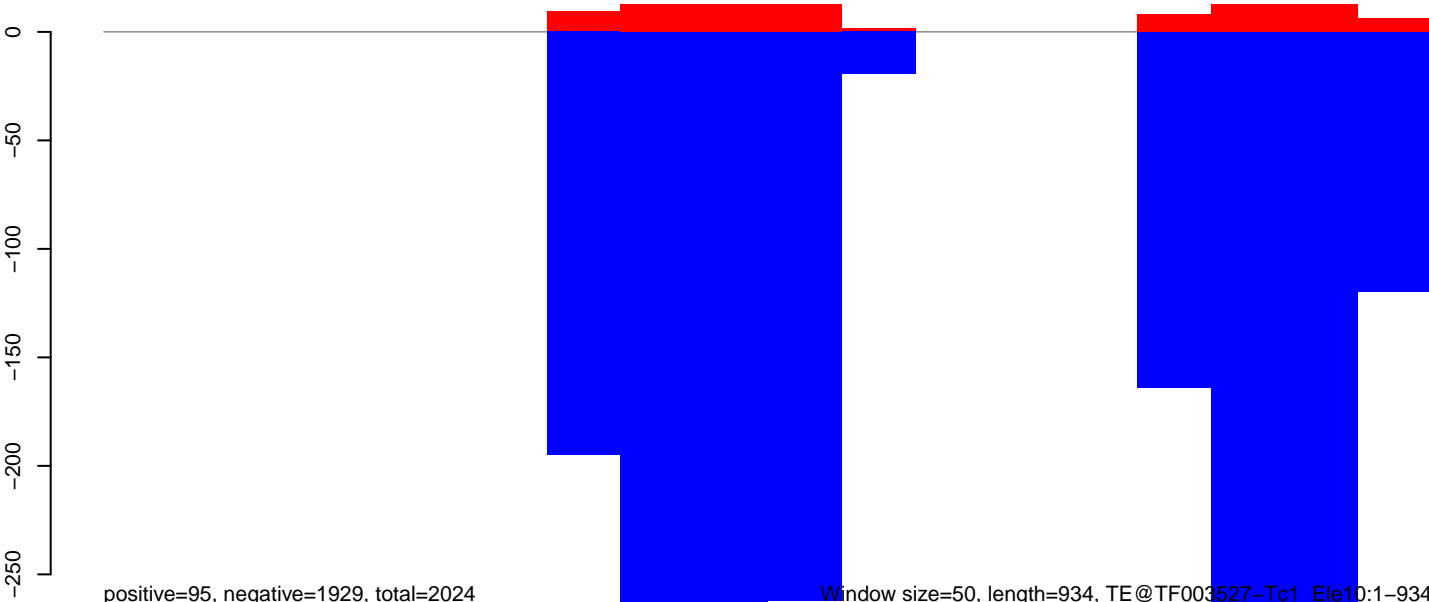
positive=95, negative=1008, total=1103

AnGam_Mos55_cells.24_35.rep



positive=0, negative=921, total=921

AnGam_Mos55_cells.rep



positive=95, negative=1929, total=2024

Window size=50, length=934, TE@TF003527-Tc1_Ele10:1-934

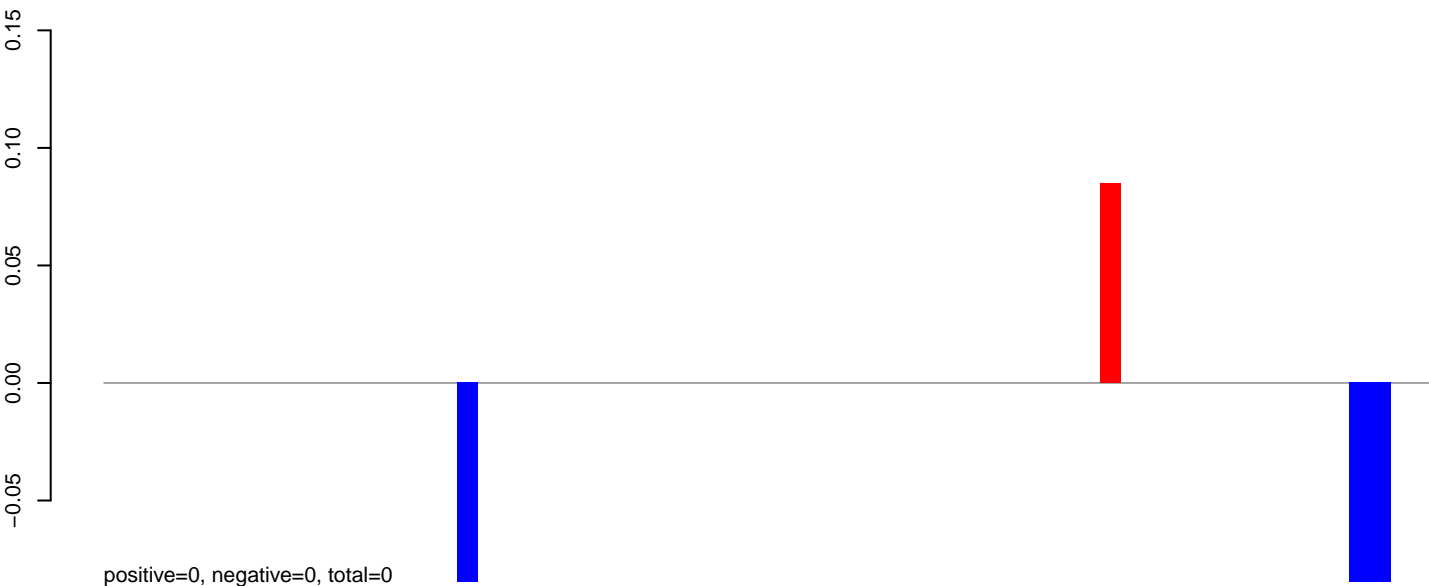
200

400

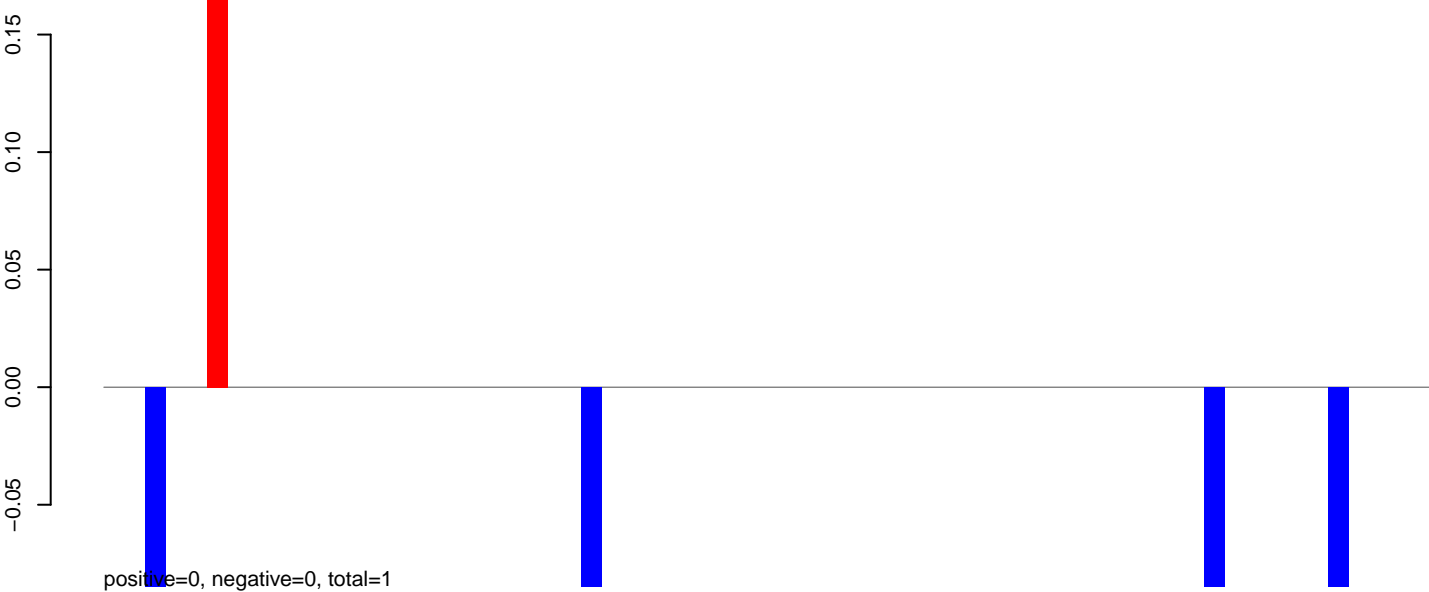
600

800

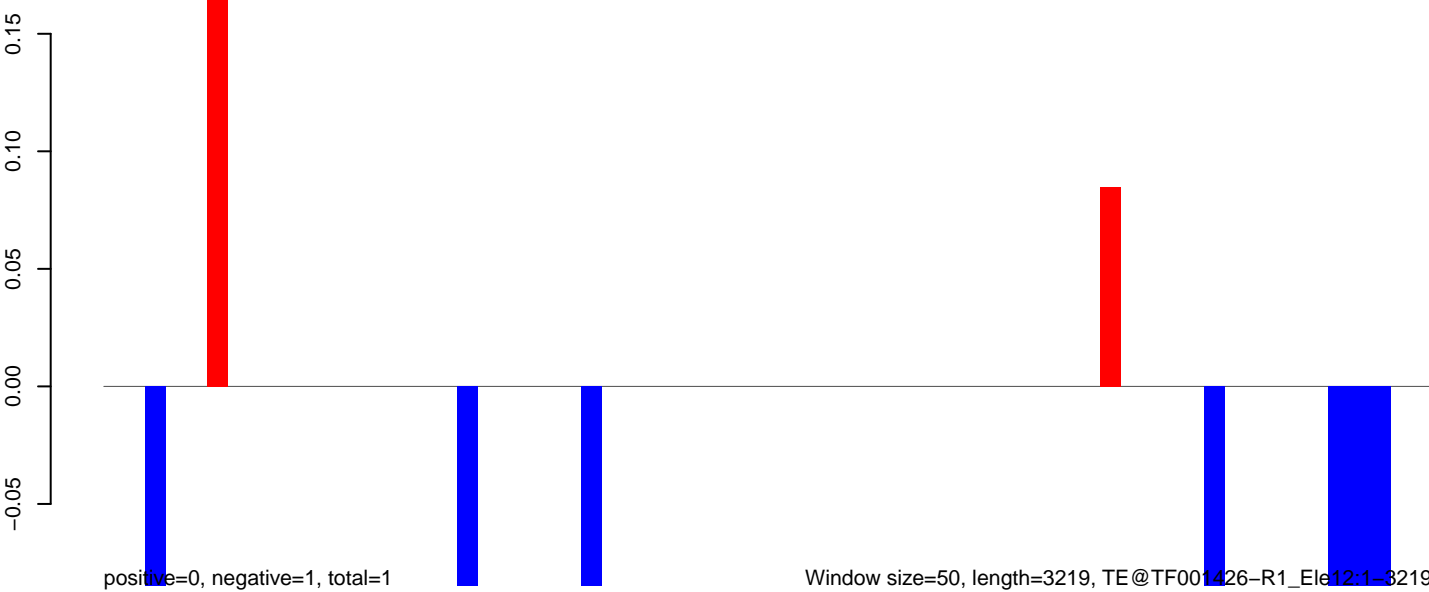
AnGam_Mos55_cells.18_23.rep



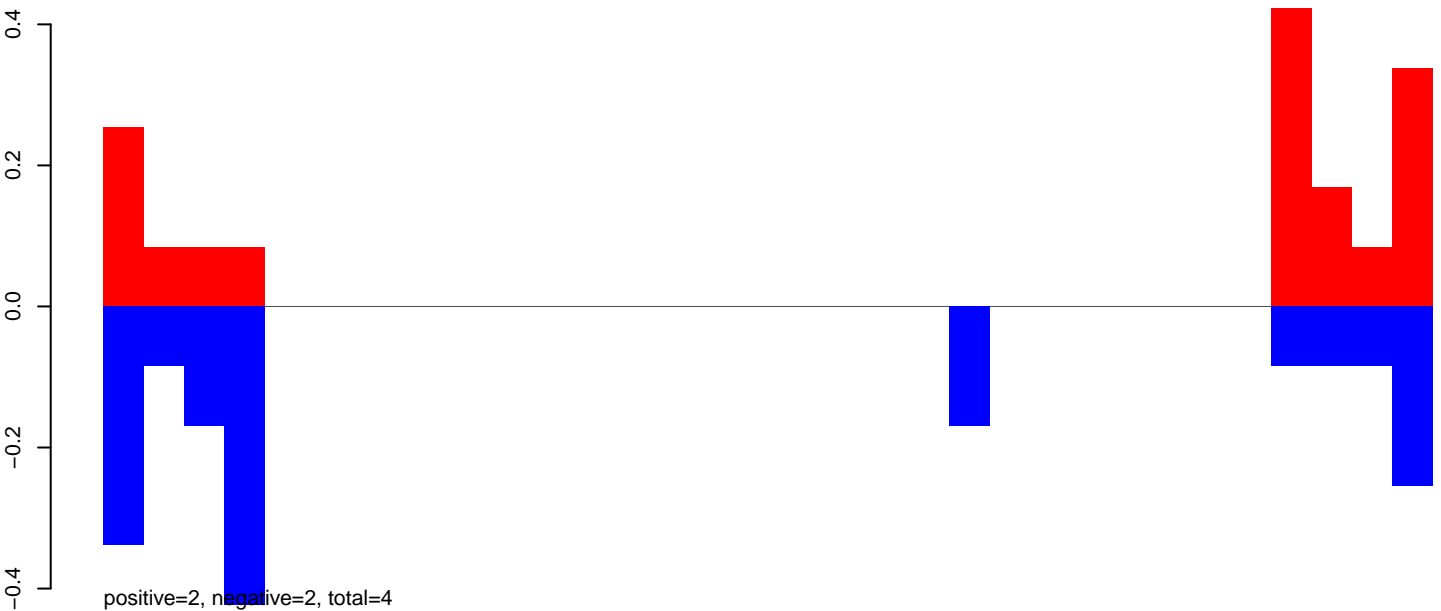
AnGam_Mos55_cells.24_35.rep



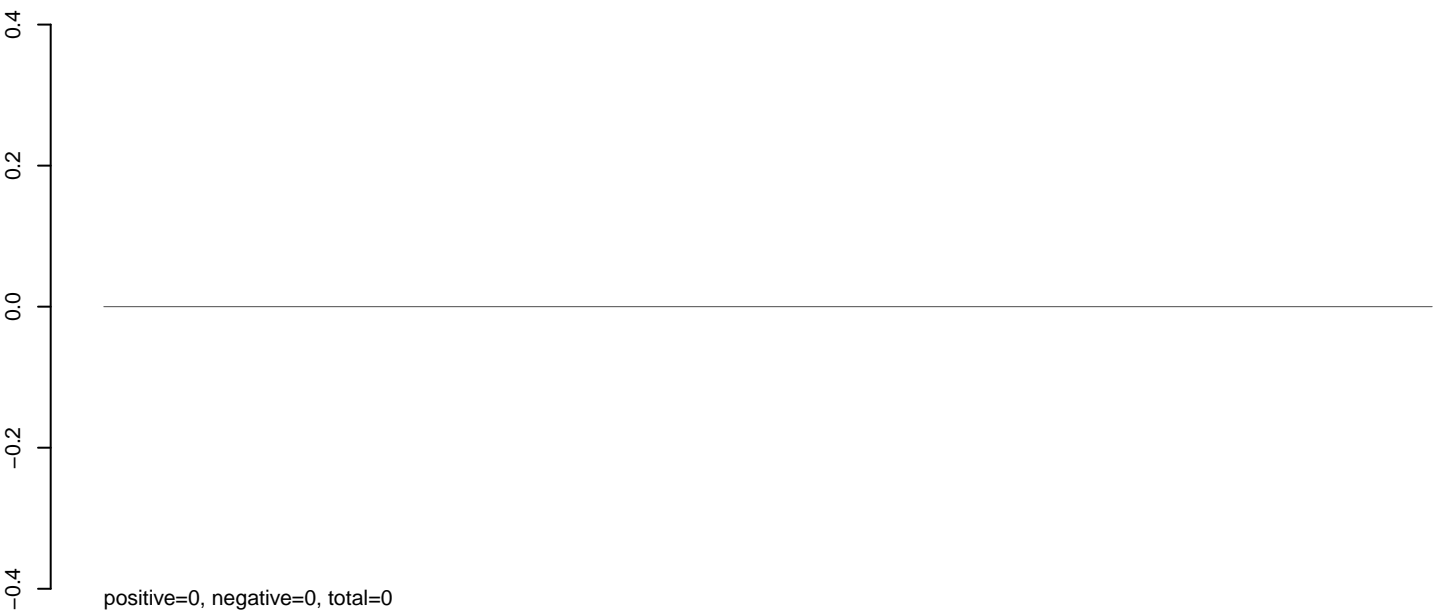
AnGam_Mos55_cells.rep



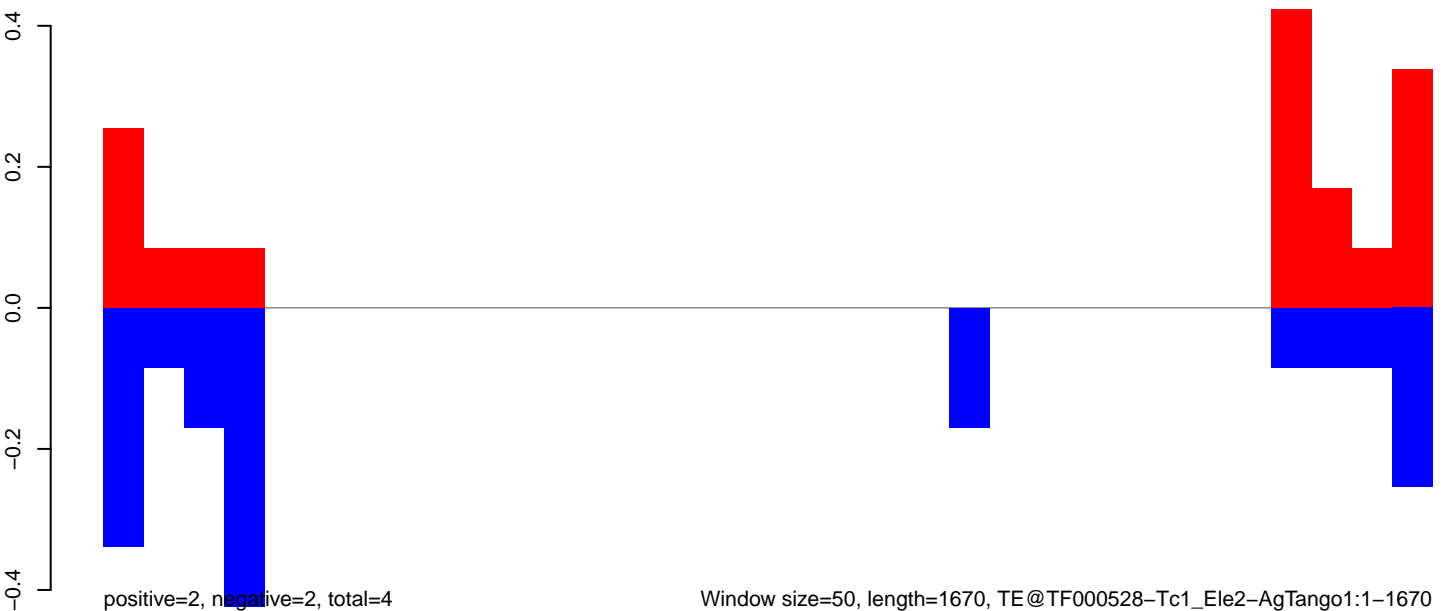
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



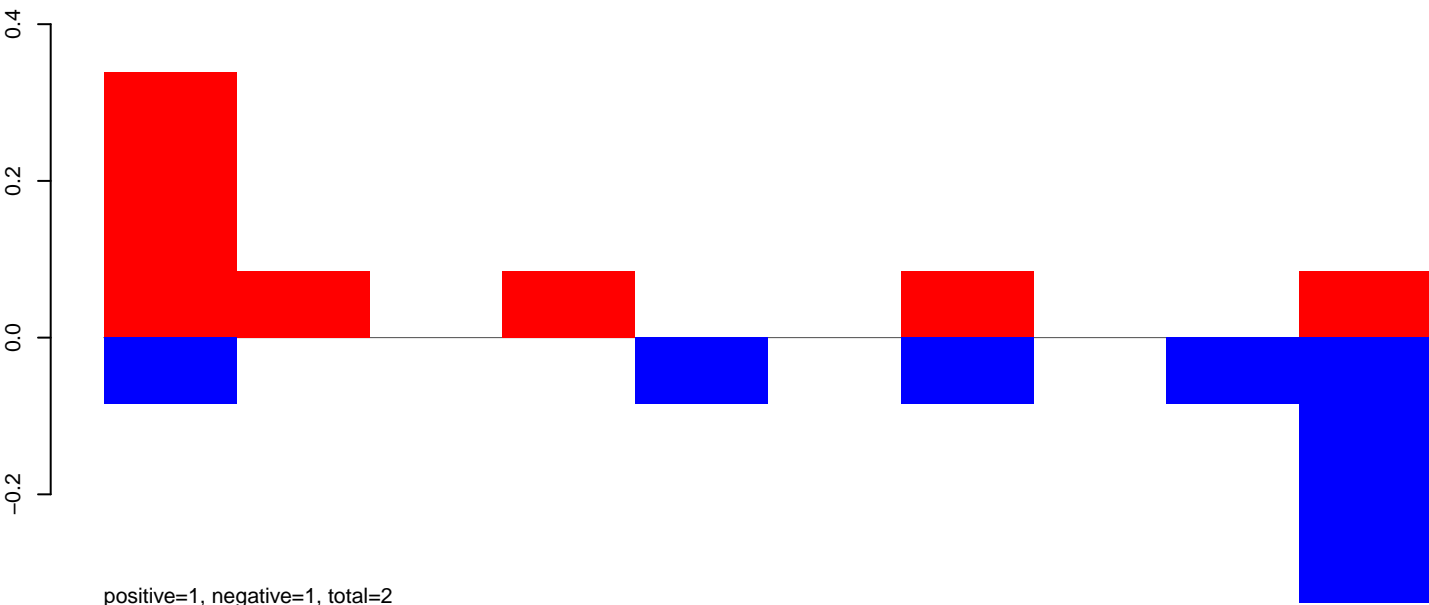
AnGam_Mos55_cells.rep



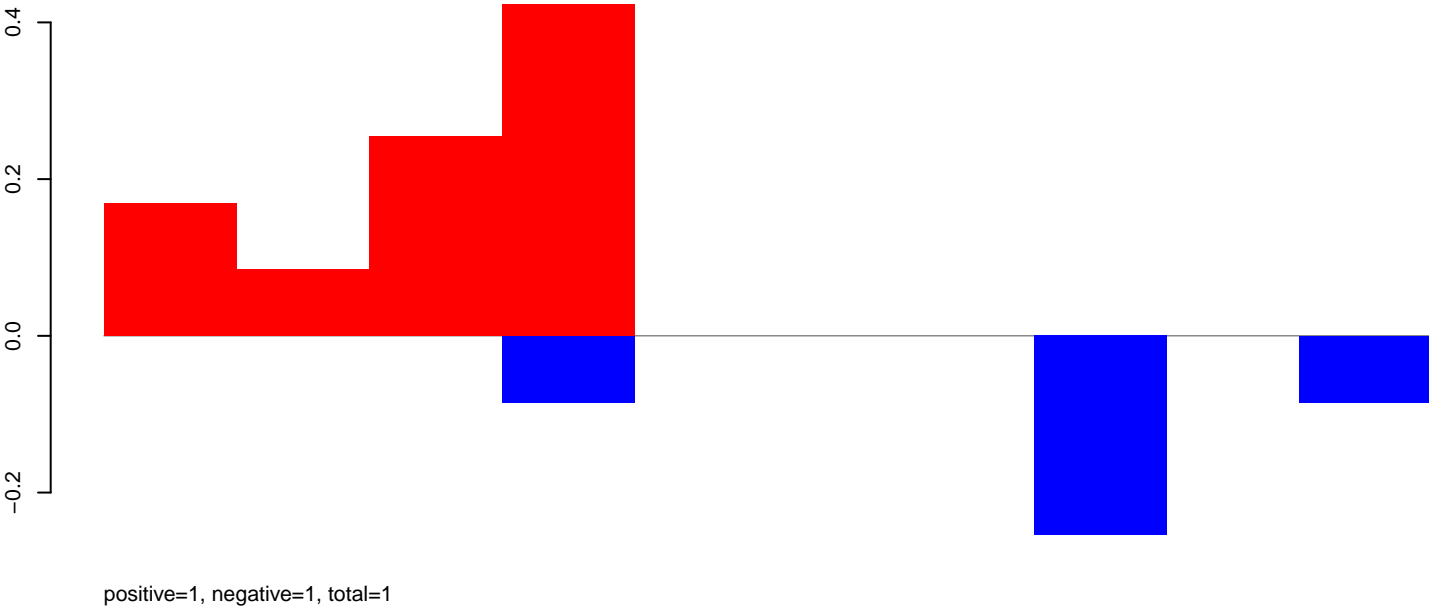
Window size=50, length=1670, TE@TF000528-Tc1_Ele2-AgTango1:1-1670

0 500 1000 1500

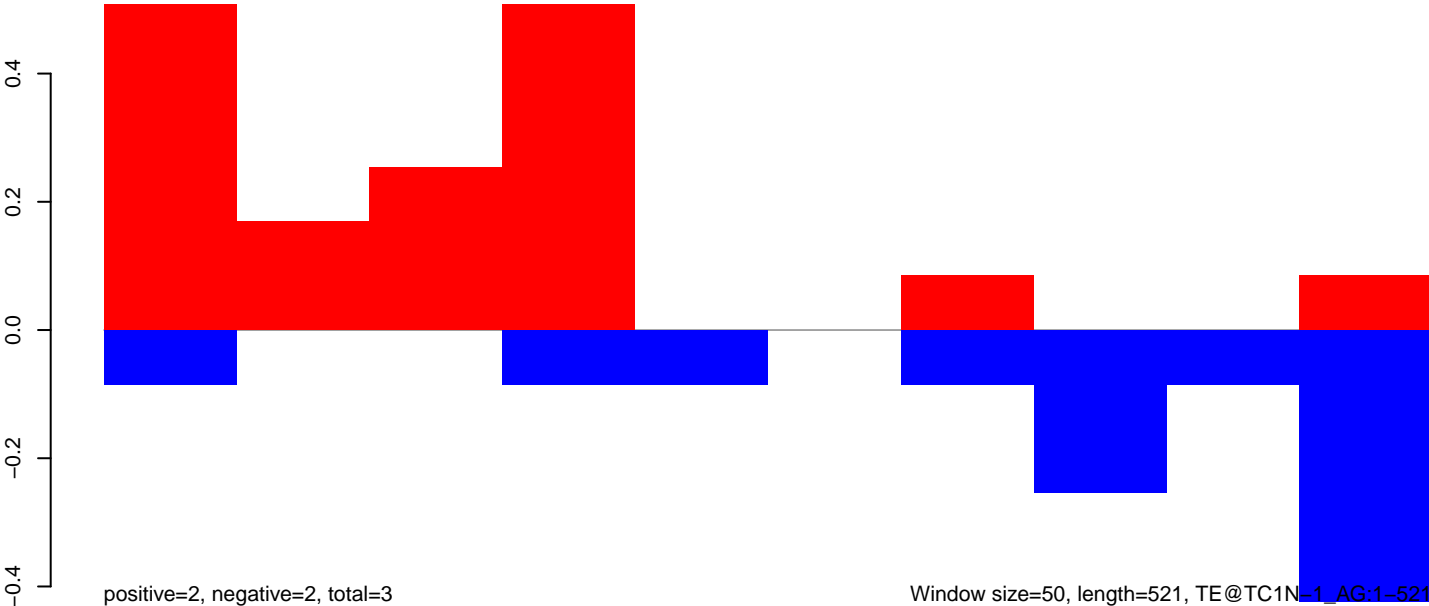
AnGam_Mos55_cells.18_23.rep



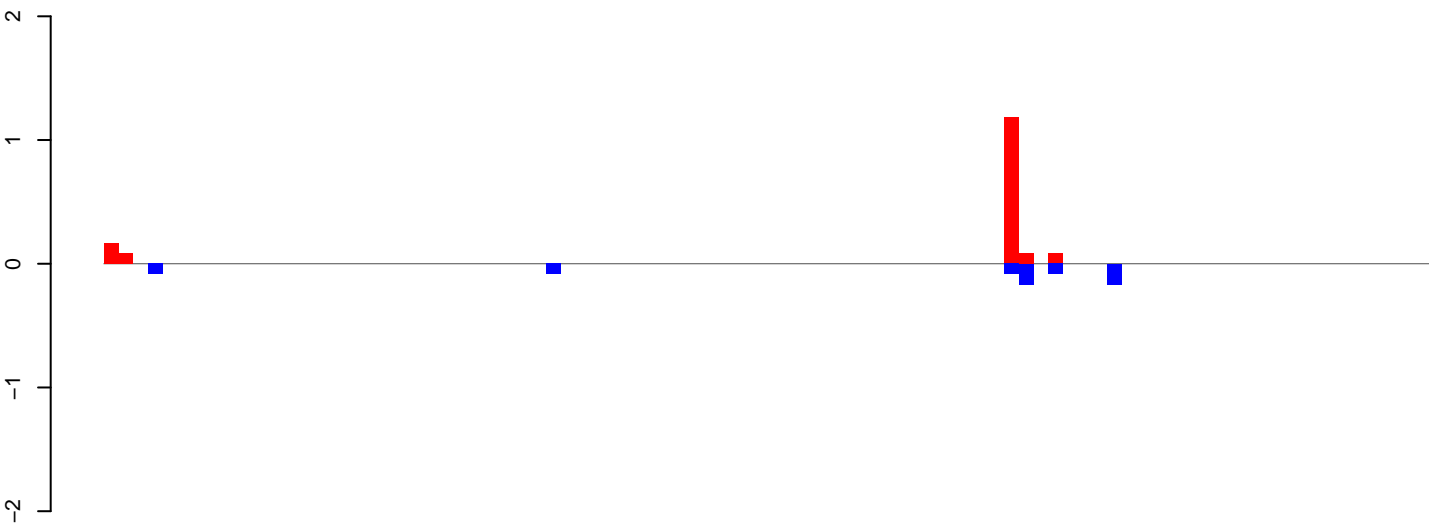
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

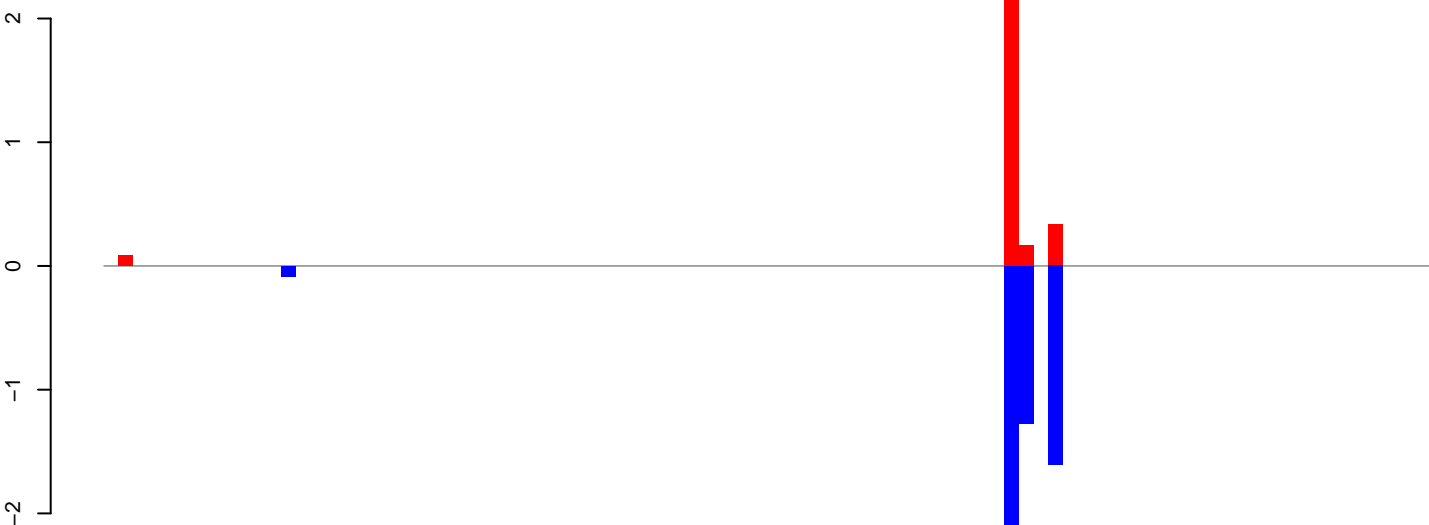


AnGam_Mos55_cells.18_23.rep



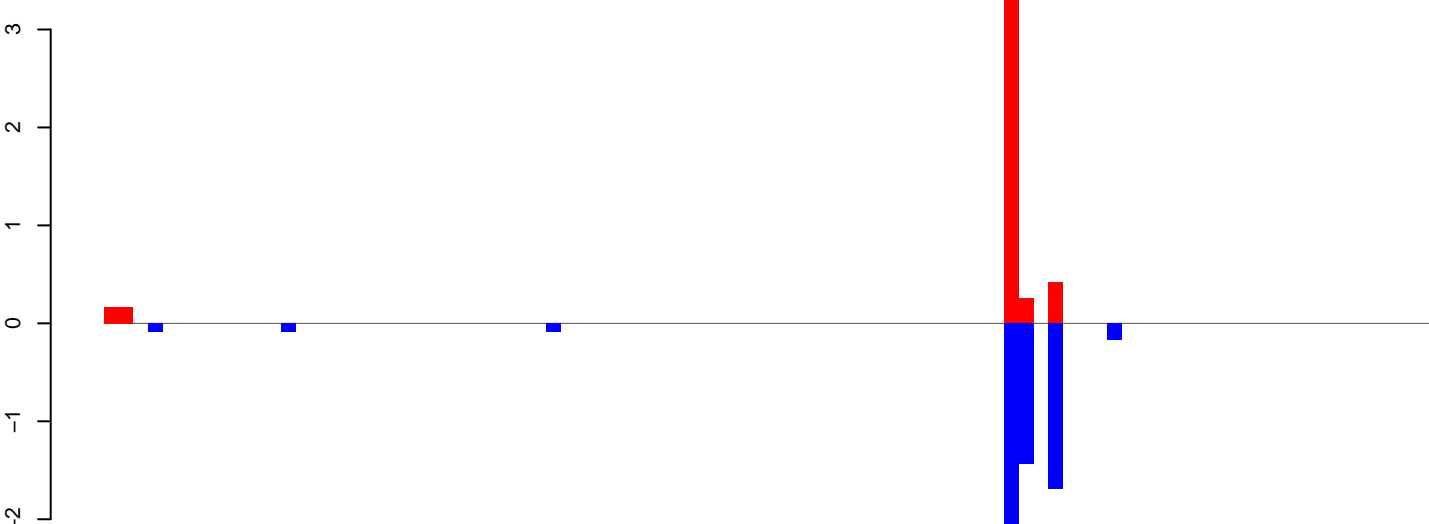
positive=2, negative=1, total=2

AnGam_Mos55_cells.24_35.rep



positive=3, negative=6, total=8

AnGam_Mos55_cells.rep

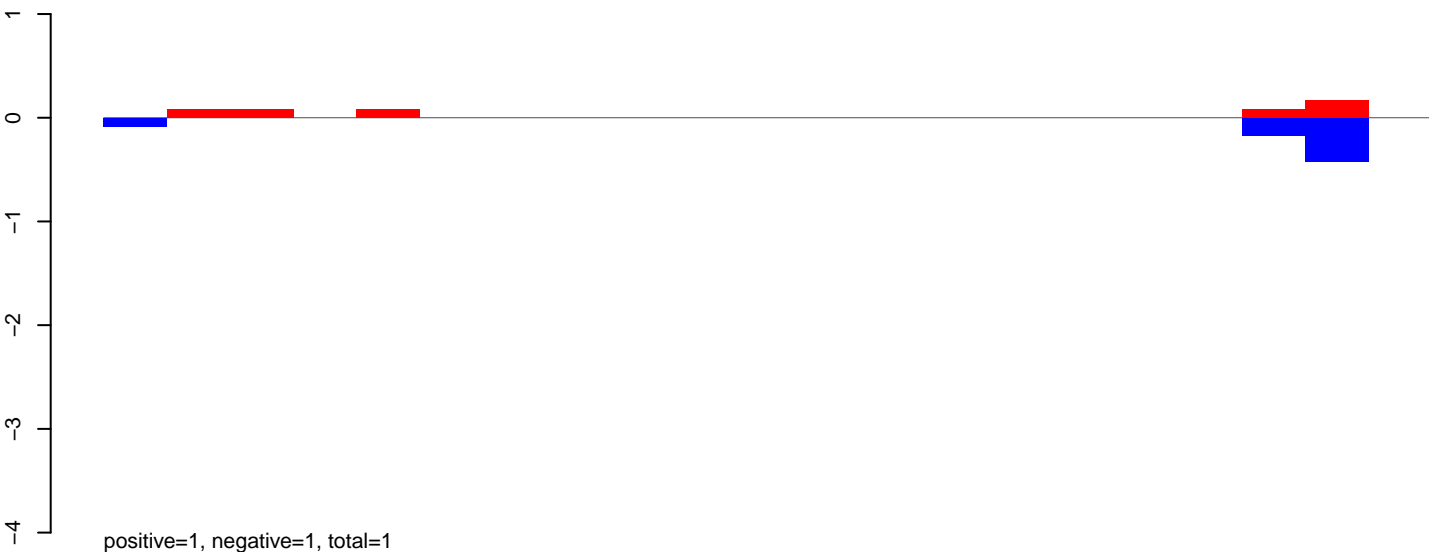


positive=4, negative=6, total=11

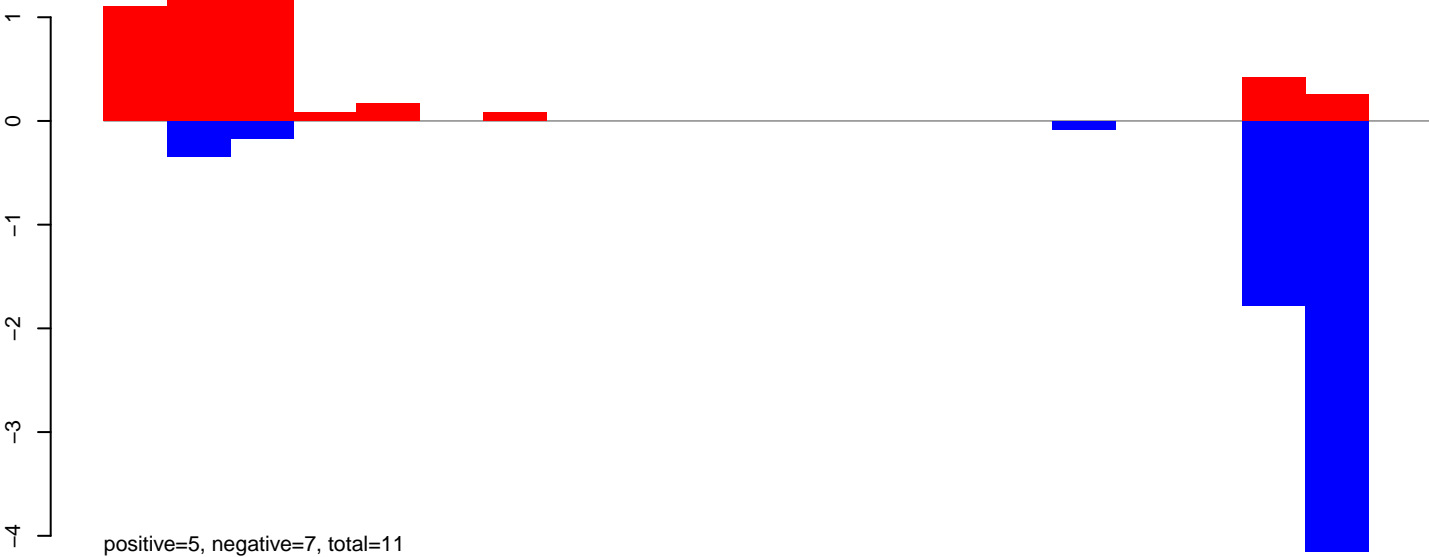
Window size=50, length=4519, TE@Harbinger-2_AG:1-4519

0 1000 2000 3000 4000

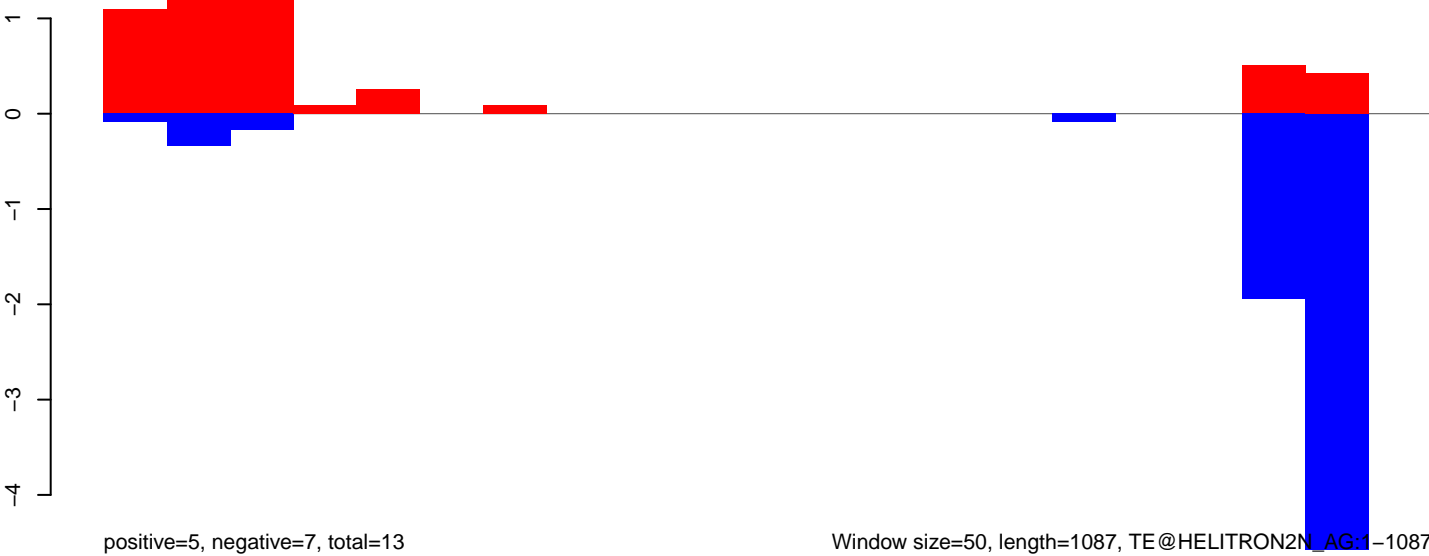
AnGam_Mos55_cells.18_23.rep



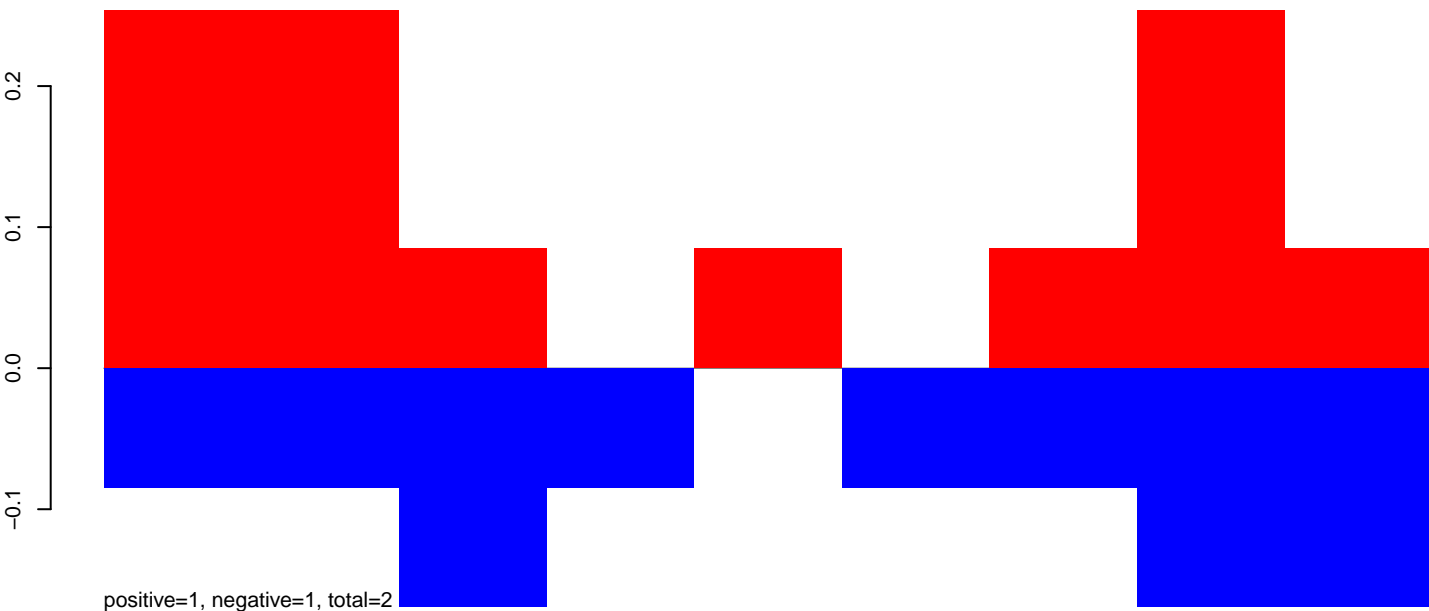
AnGam_Mos55_cells.24_35.rep



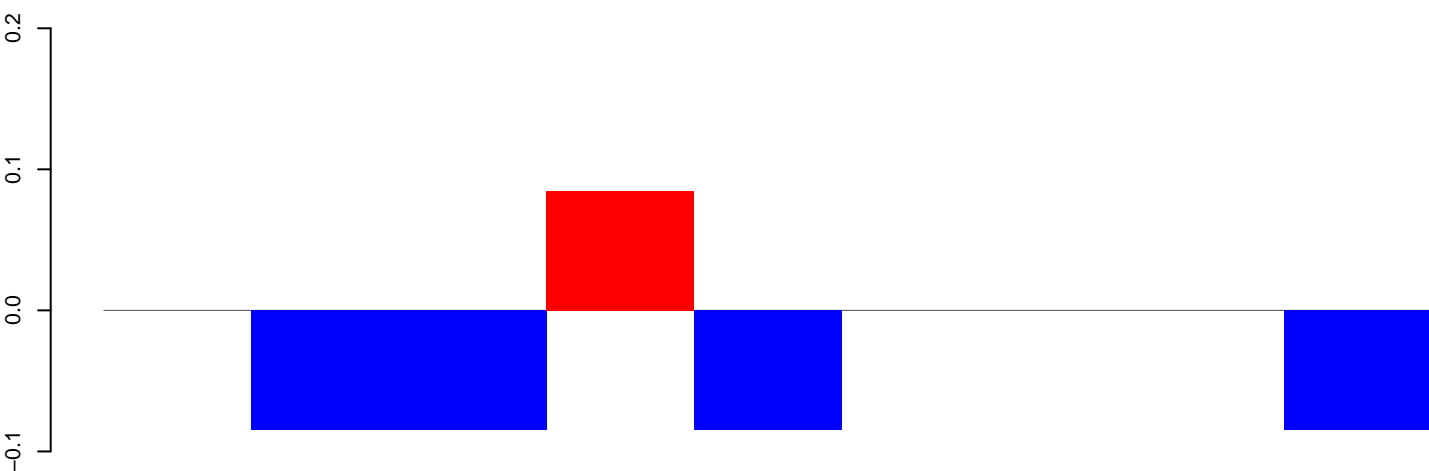
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep

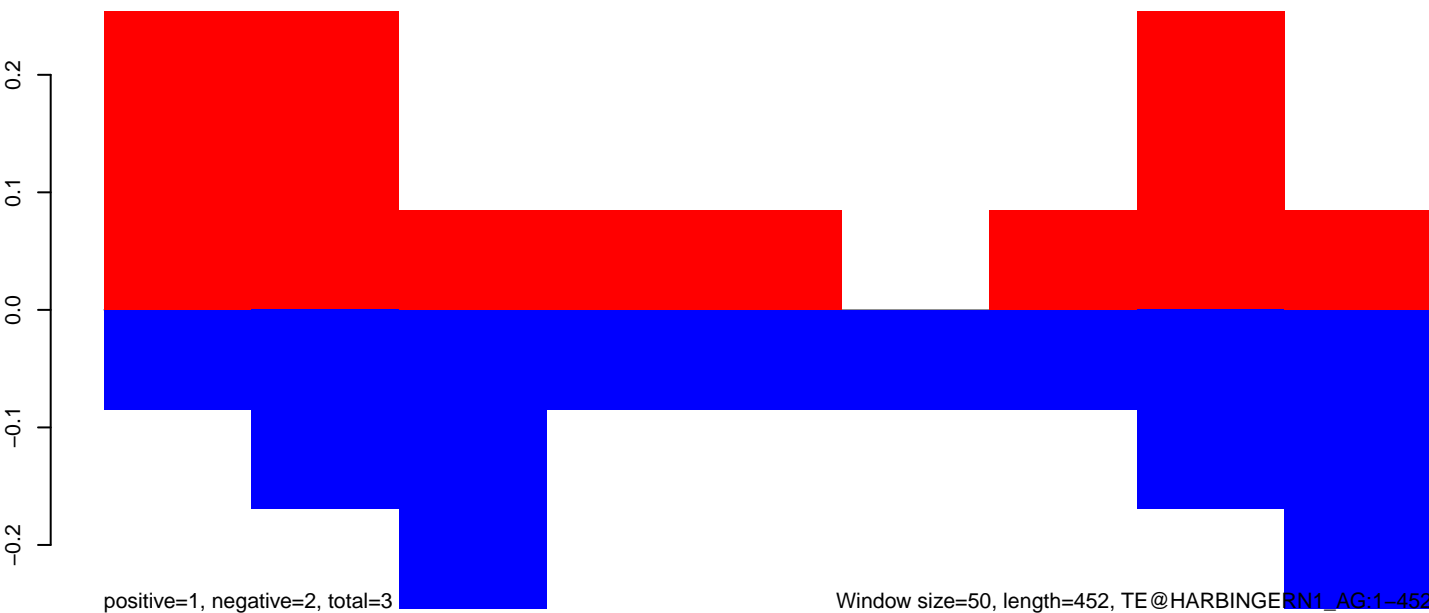


AnGam_Mos55_cells.24_35.rep



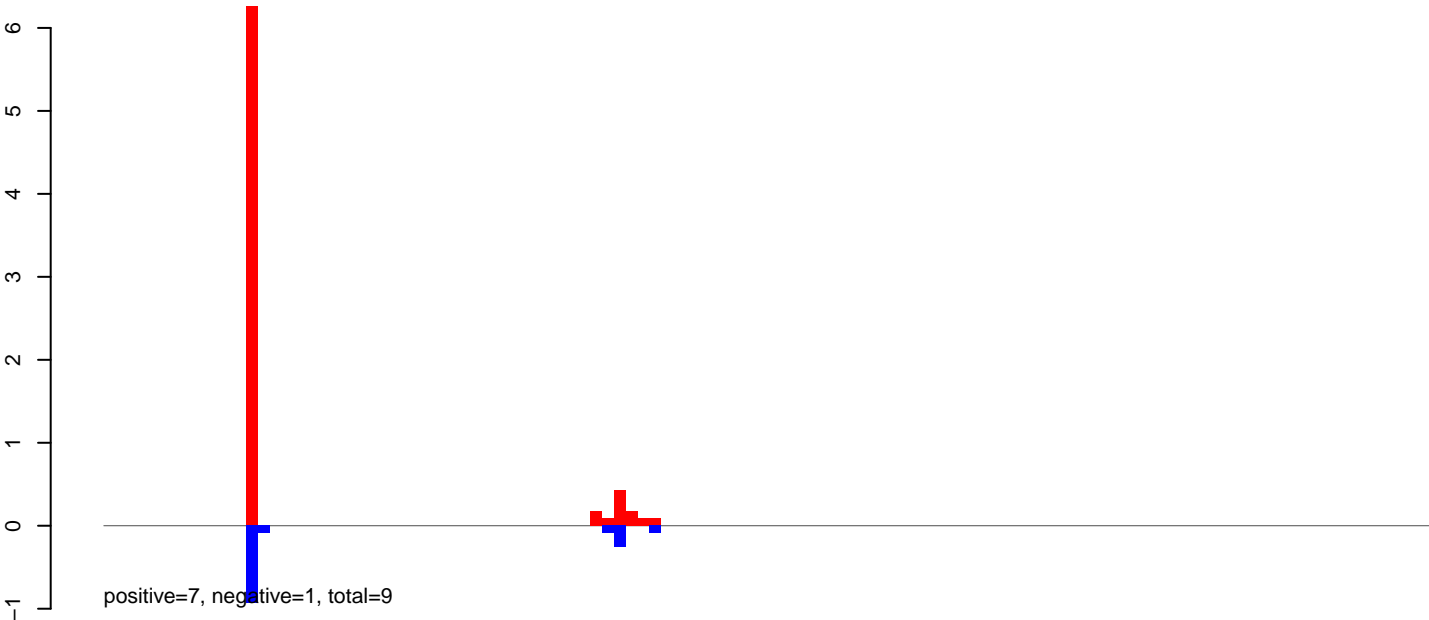
positive=0, negative=0, total=1

AnGam_Mos55_cells.rep

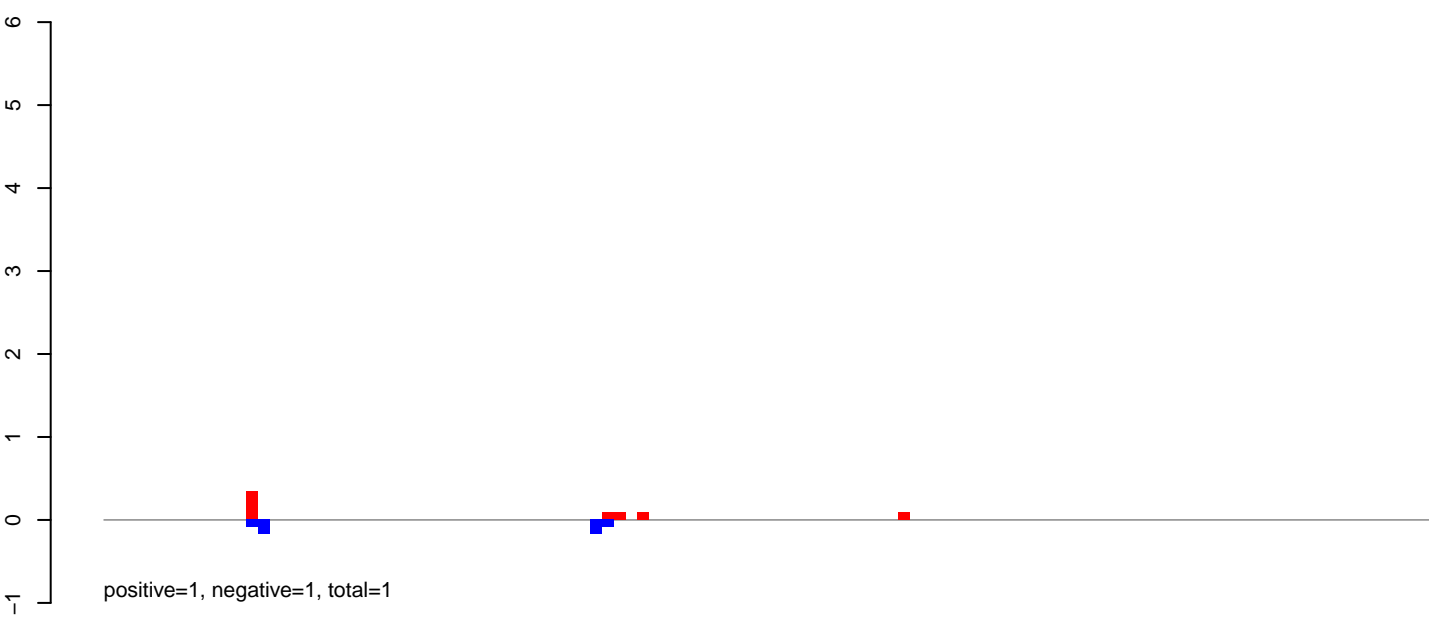


100 200 300 400 500

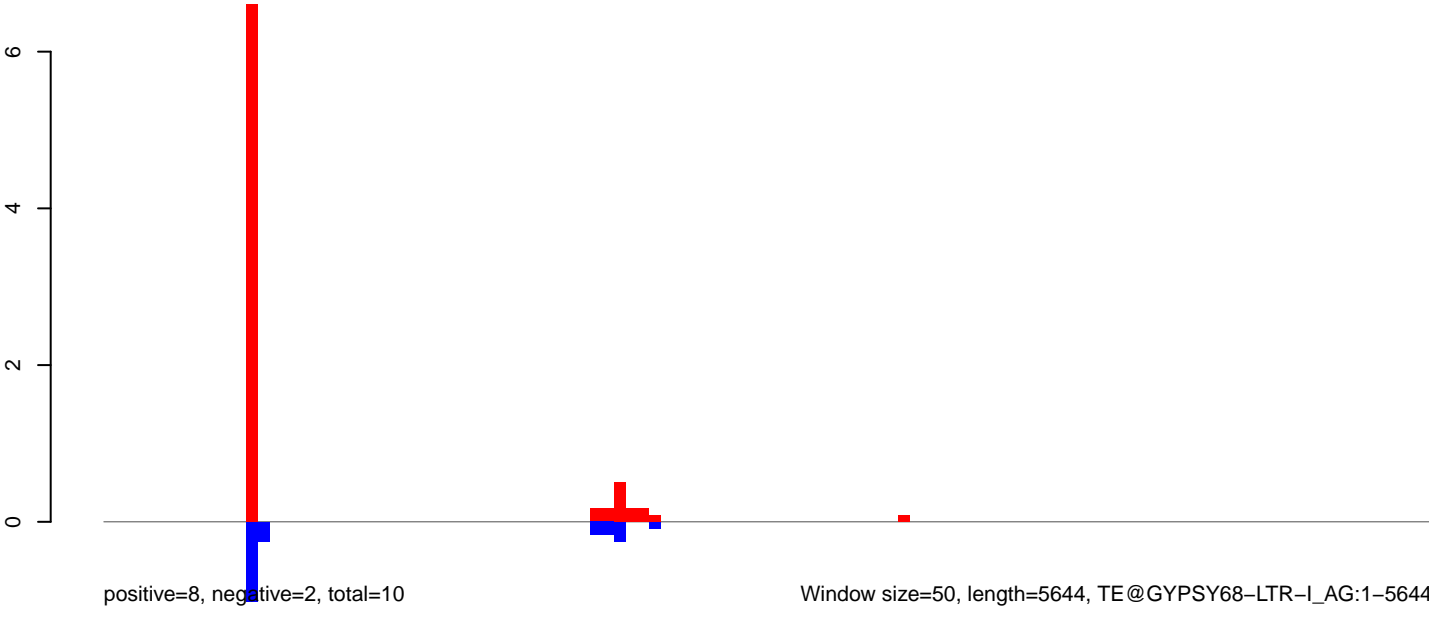
AnGam_Mos55_cells.18_23.rep



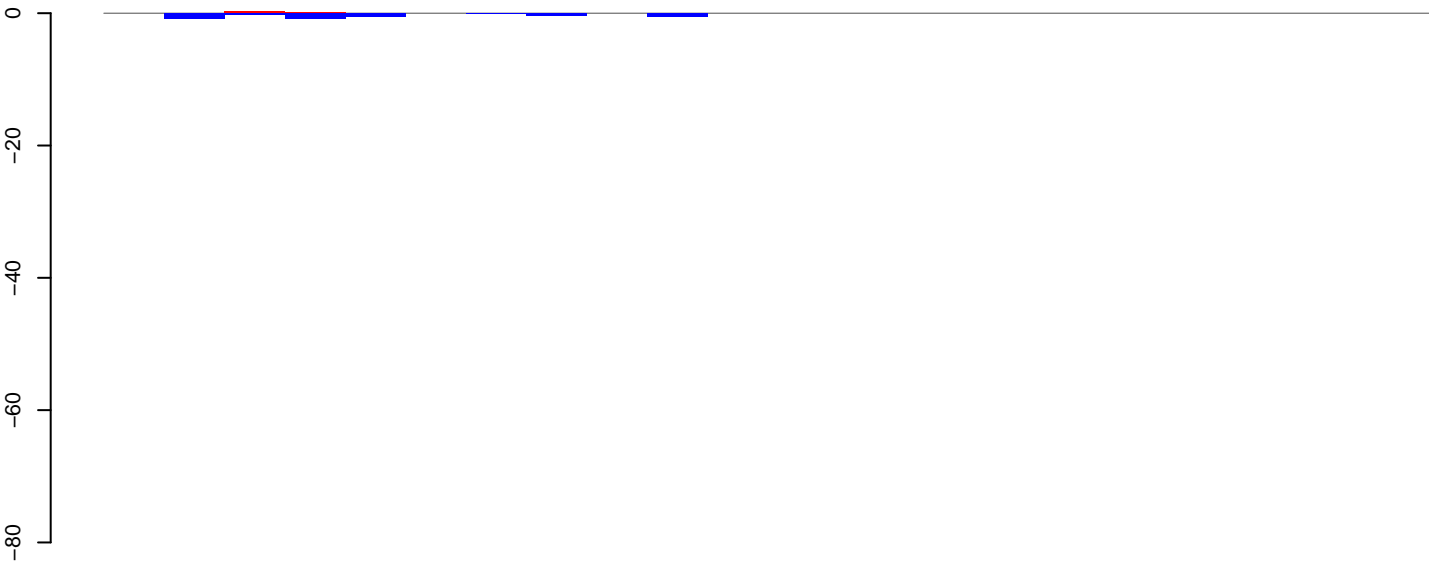
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

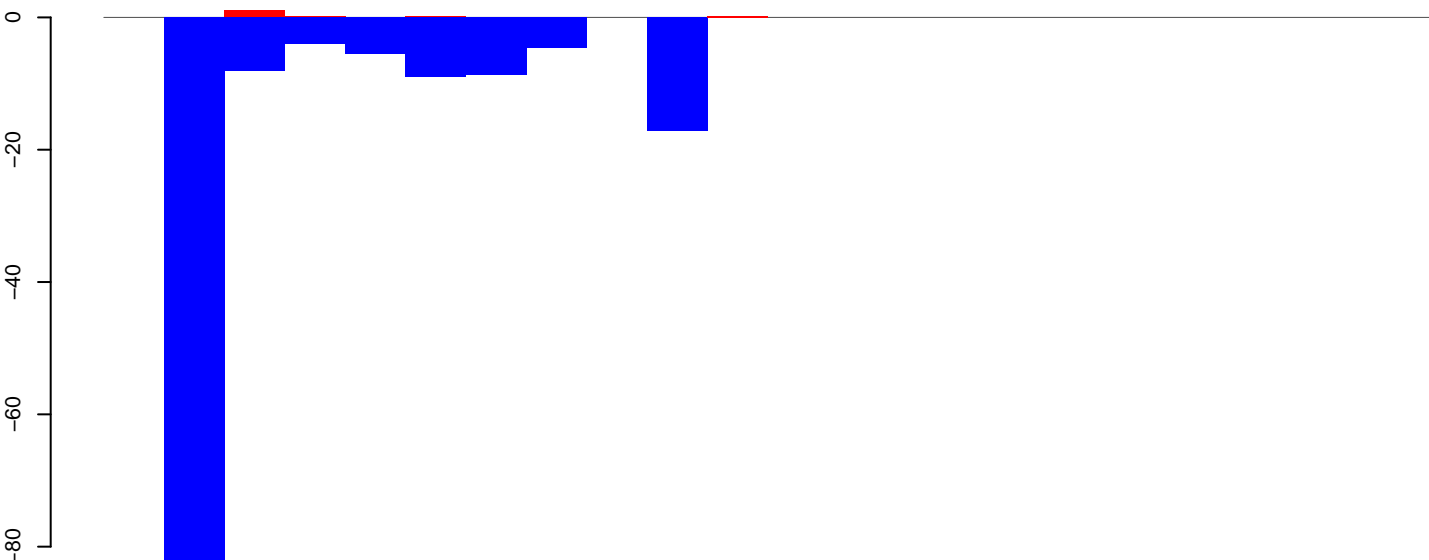


AnGam_Mos55_cells.18_23.rep



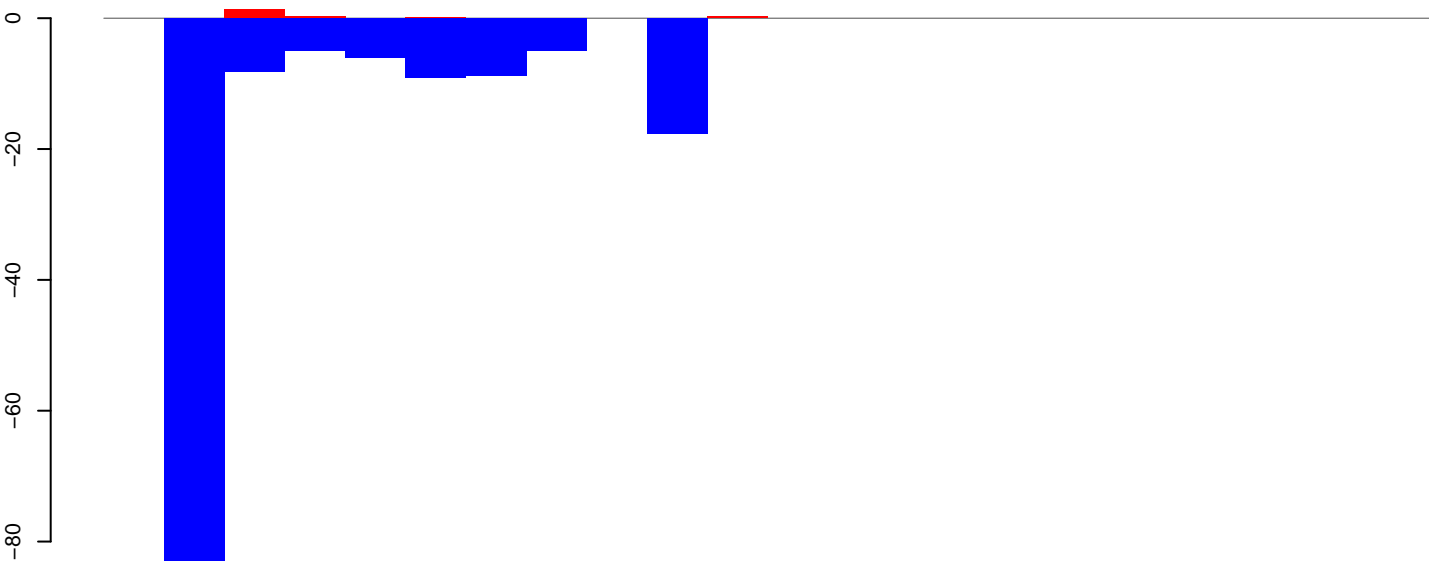
positive=0, negative=3, total=4

AnGam_Mos55_cells.24_35.rep



positive=2, negative=146, total=148

AnGam_Mos55_cells.rep

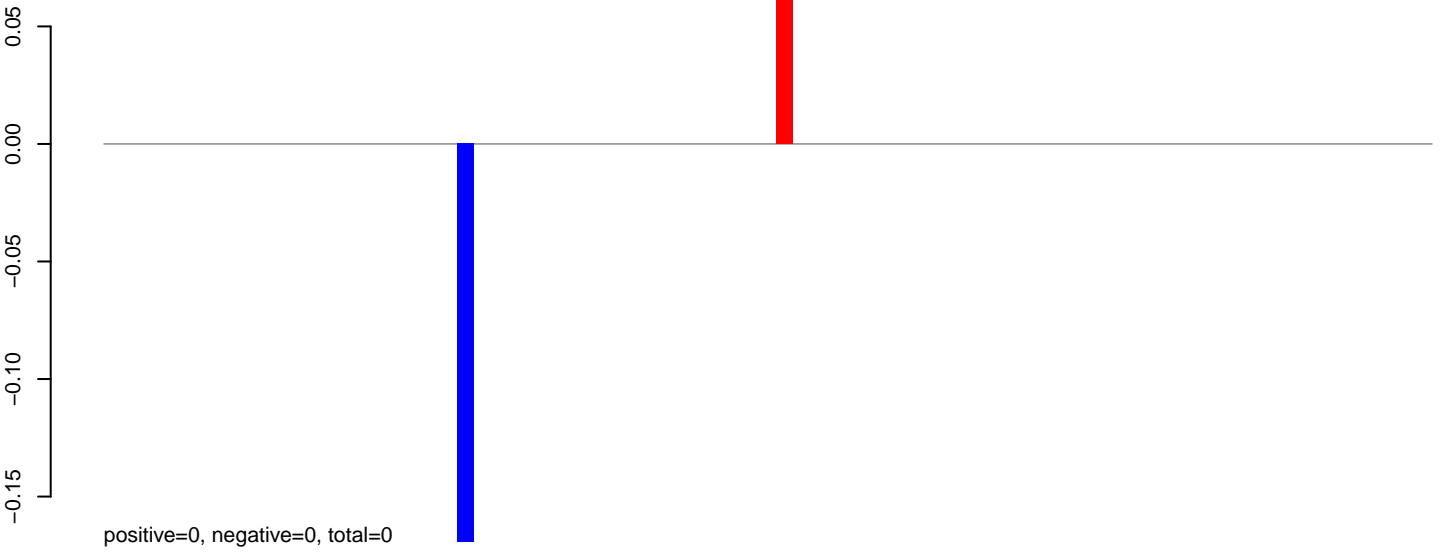


positive=2, negative=150, total=152

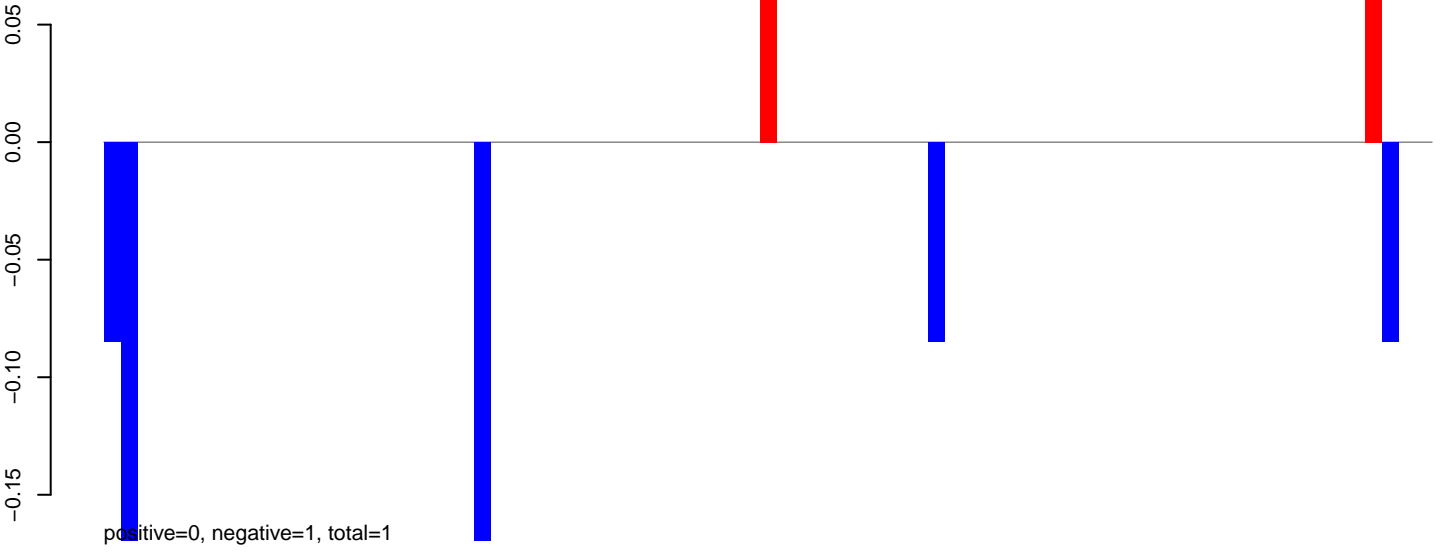
Window size=50, length=1143, TE@EAL40144-Unknown:1-1143

200 400 600 800 1000

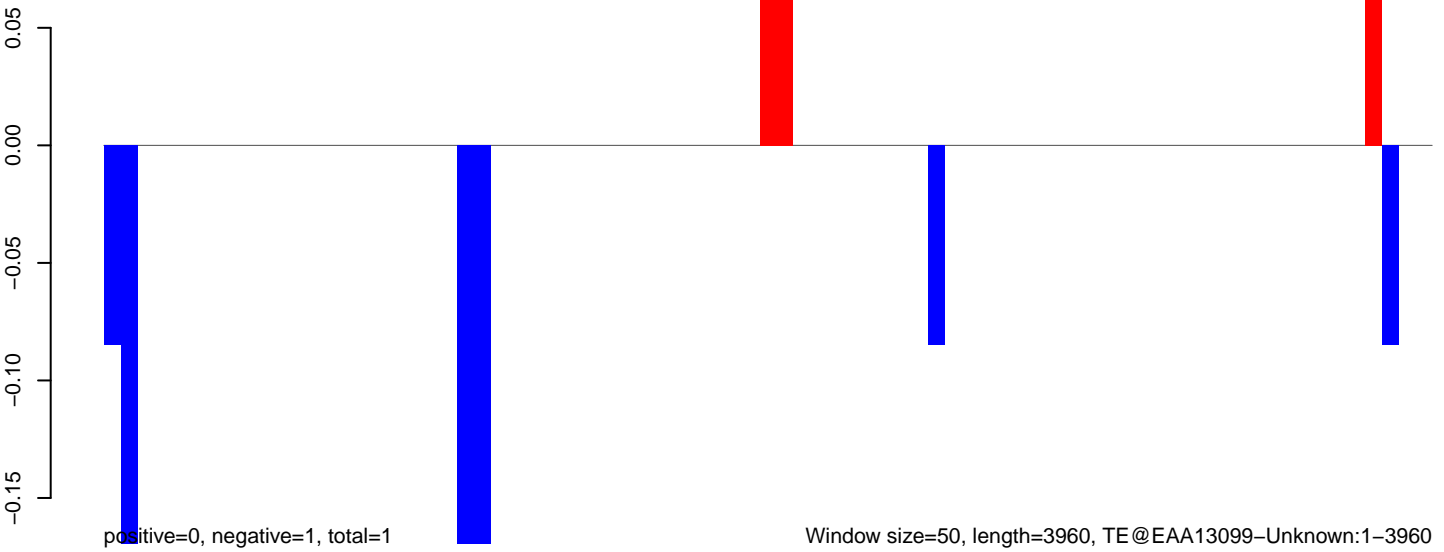
AnGam_Mos55_cells.18_23.rep



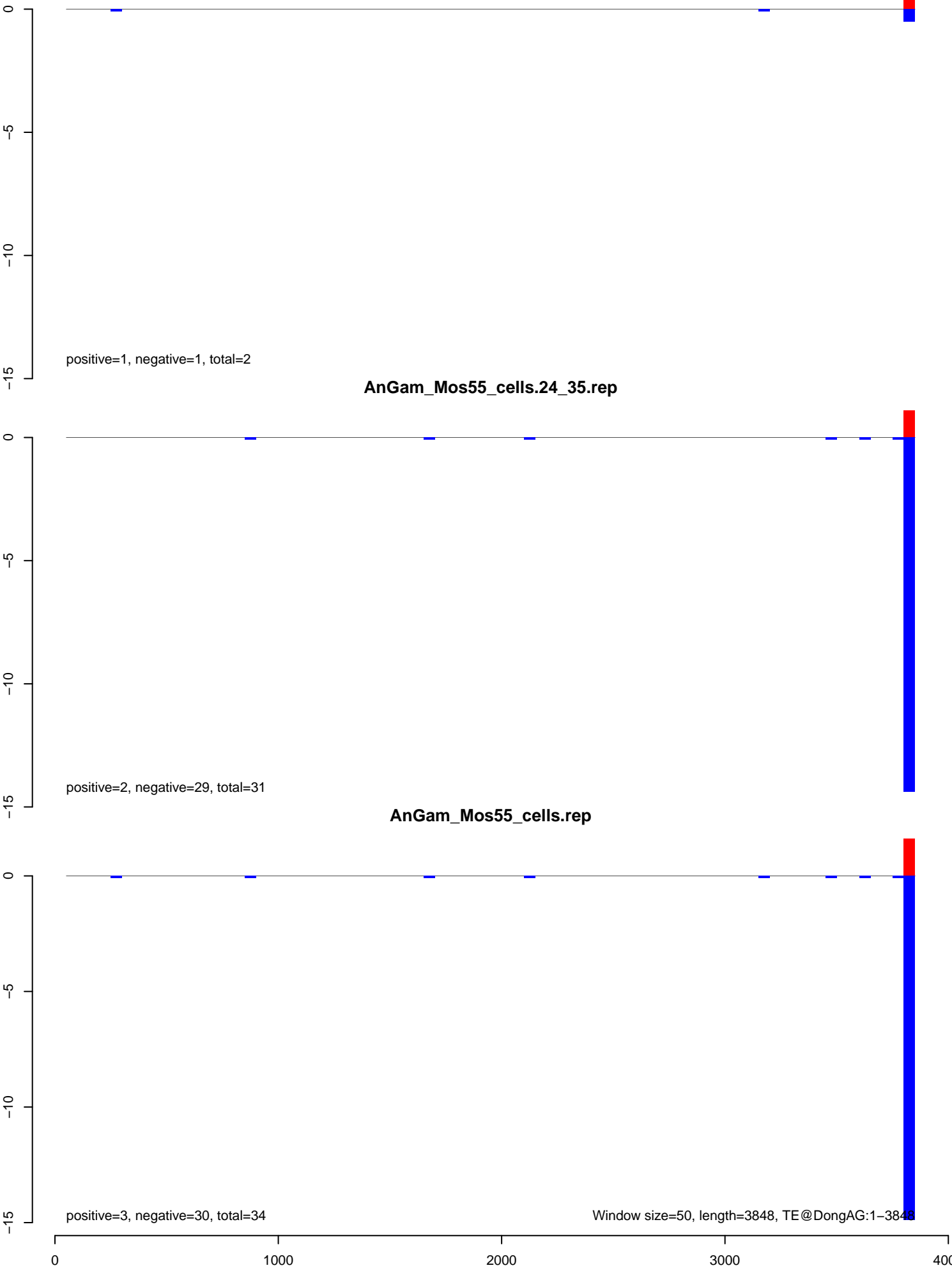
AnGam_Mos55_cells.24_35.rep



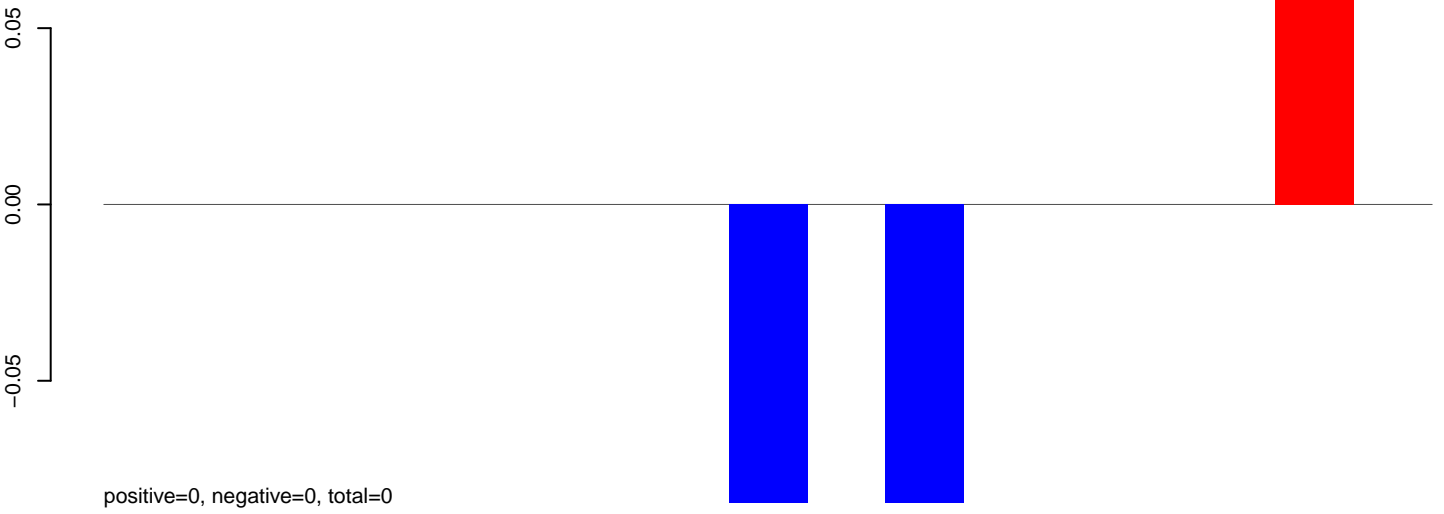
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep

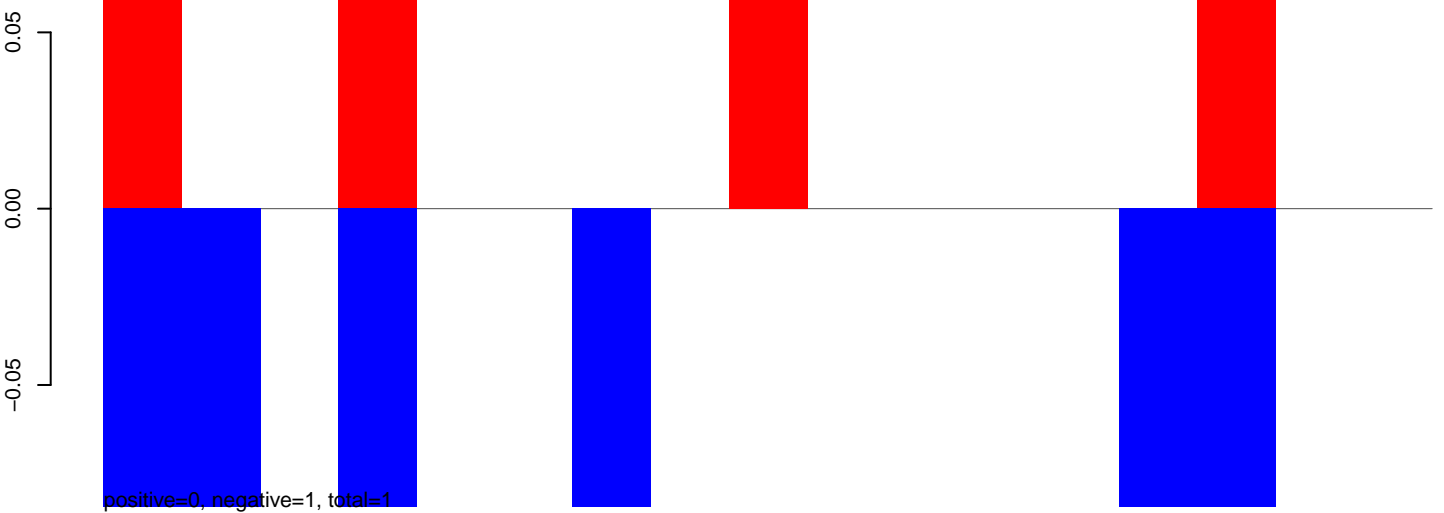


AnGam_Mos55_cells.18_23.rep



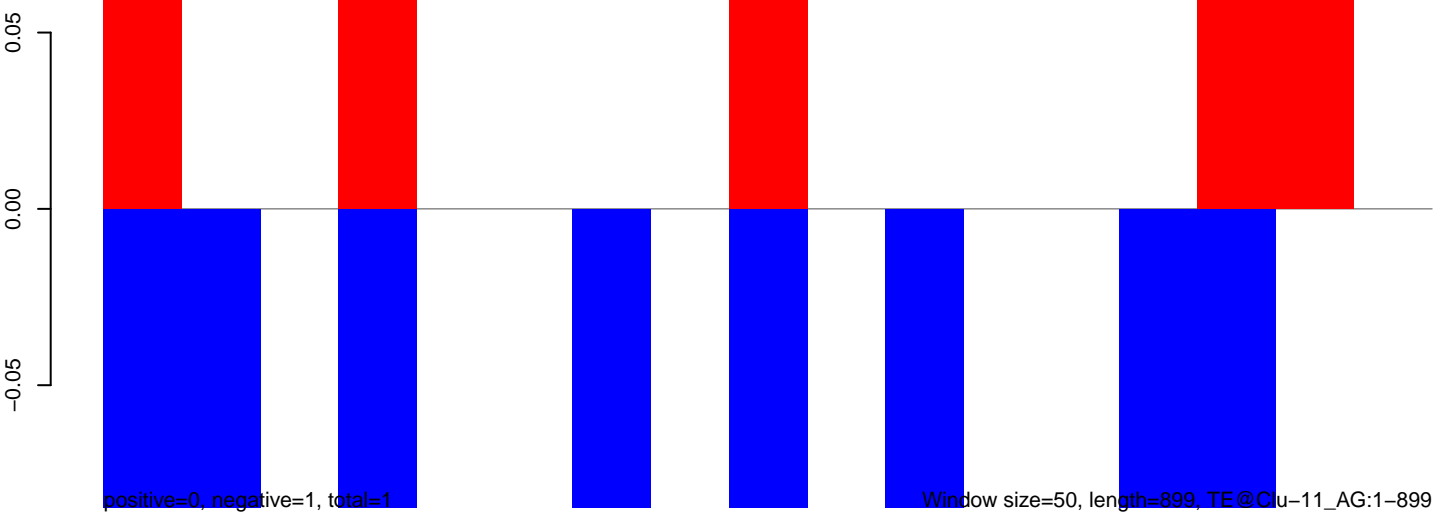
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=1, total=1

AnGam_Mos55_cells.rep

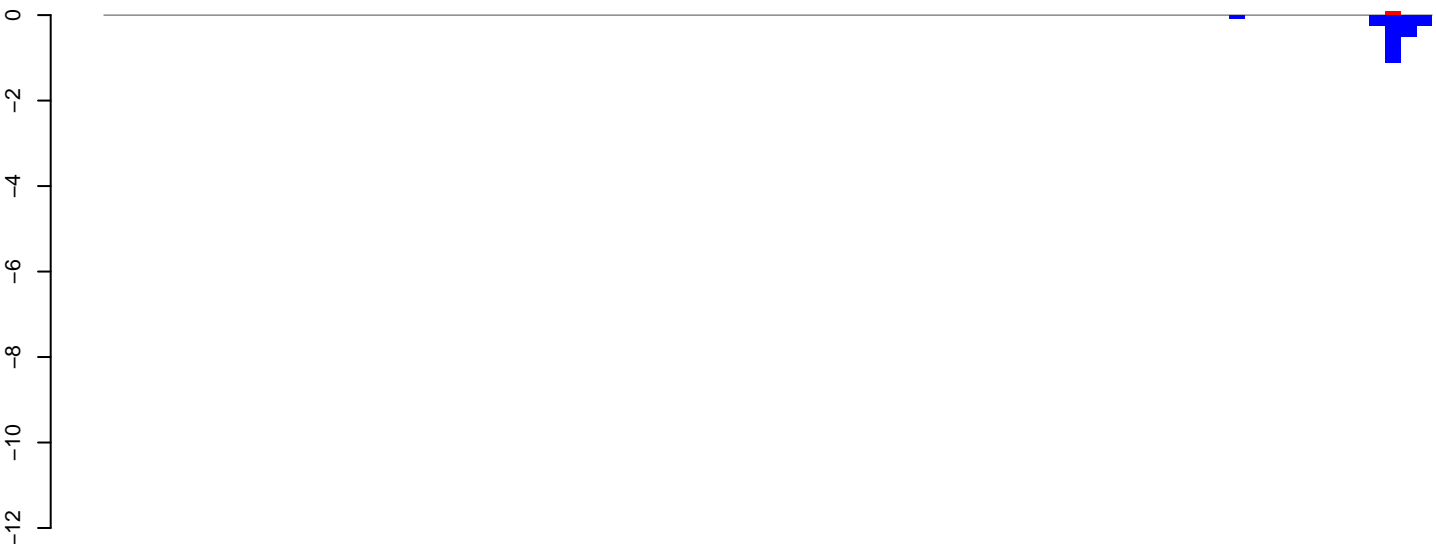


positive=0, negative=1, total=1

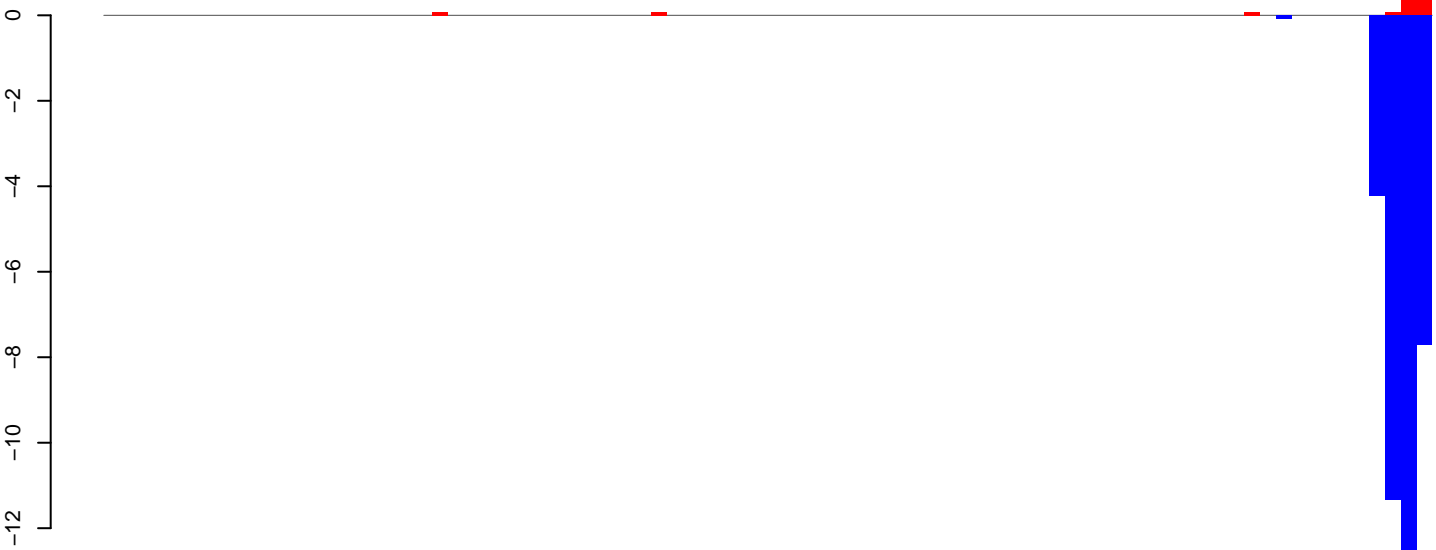
Window size=50, length=899, TE@Clu-11_AG:1-899

200 400 600 800

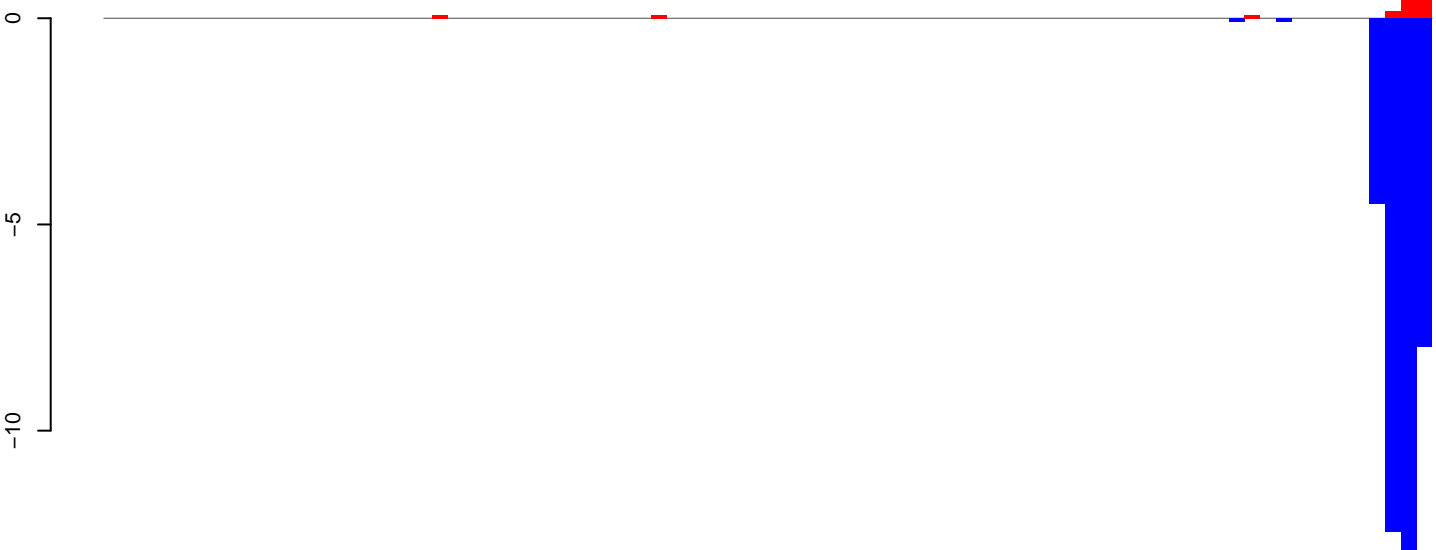
AnGam_Mos55_cells.18_23.rep



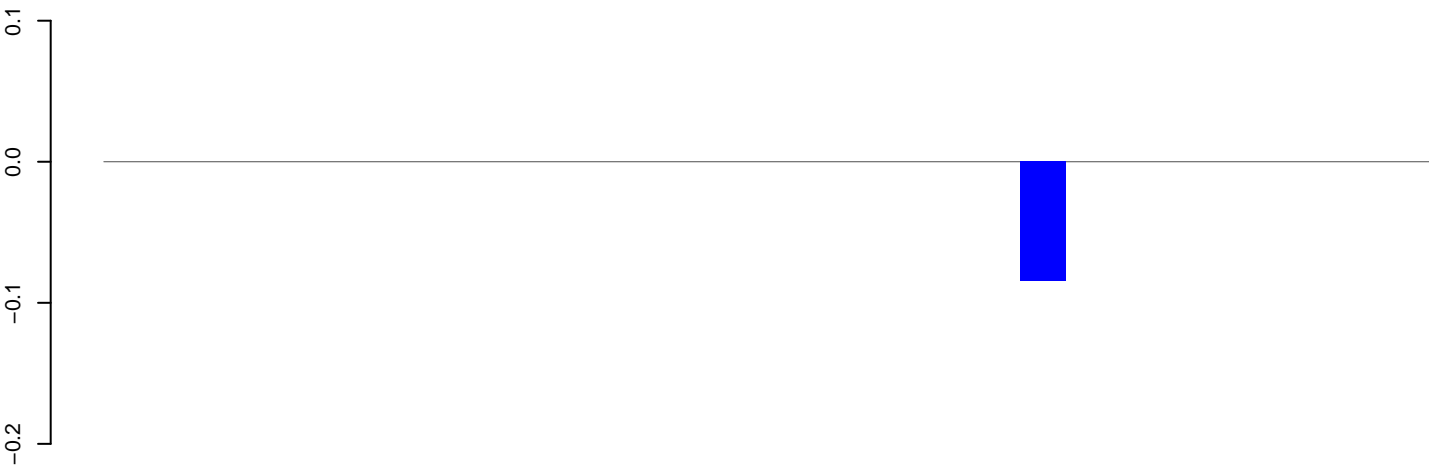
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

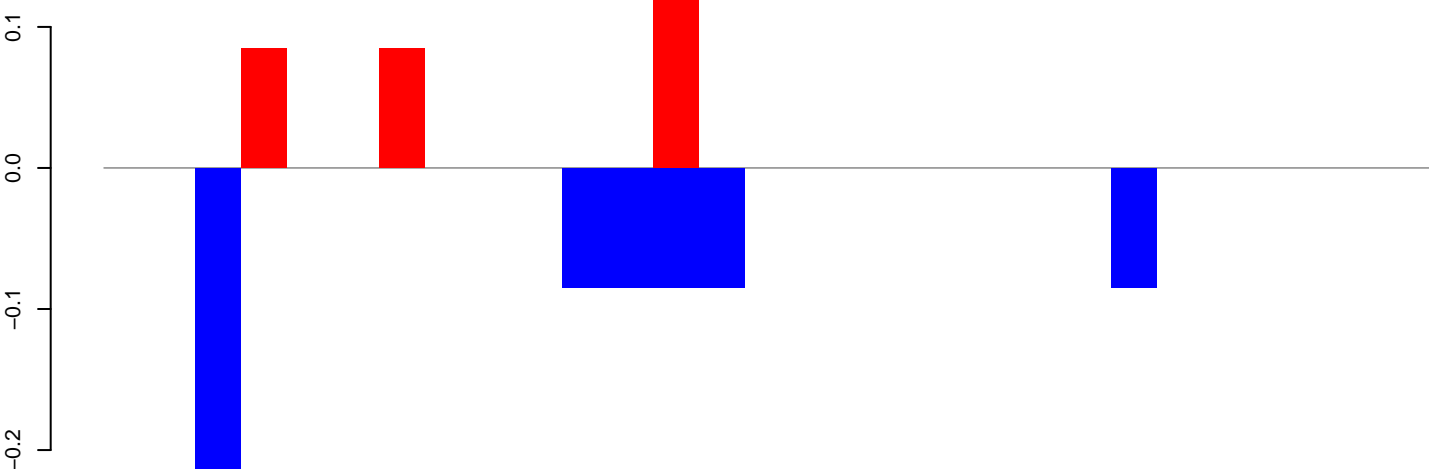


AnGam_Mos55_cells.18_23.rep



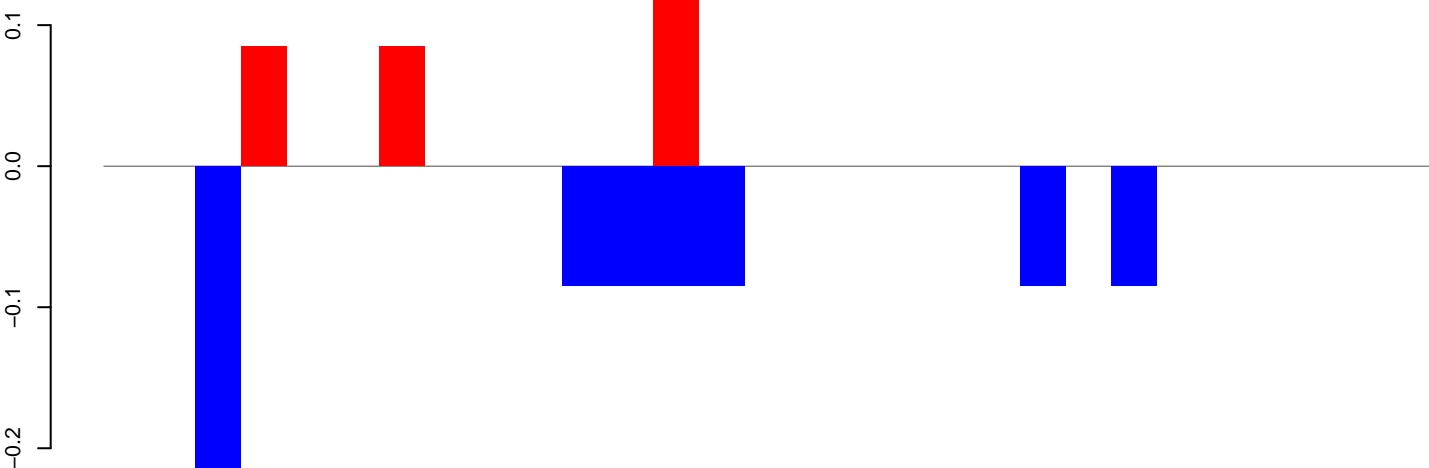
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=1, total=1

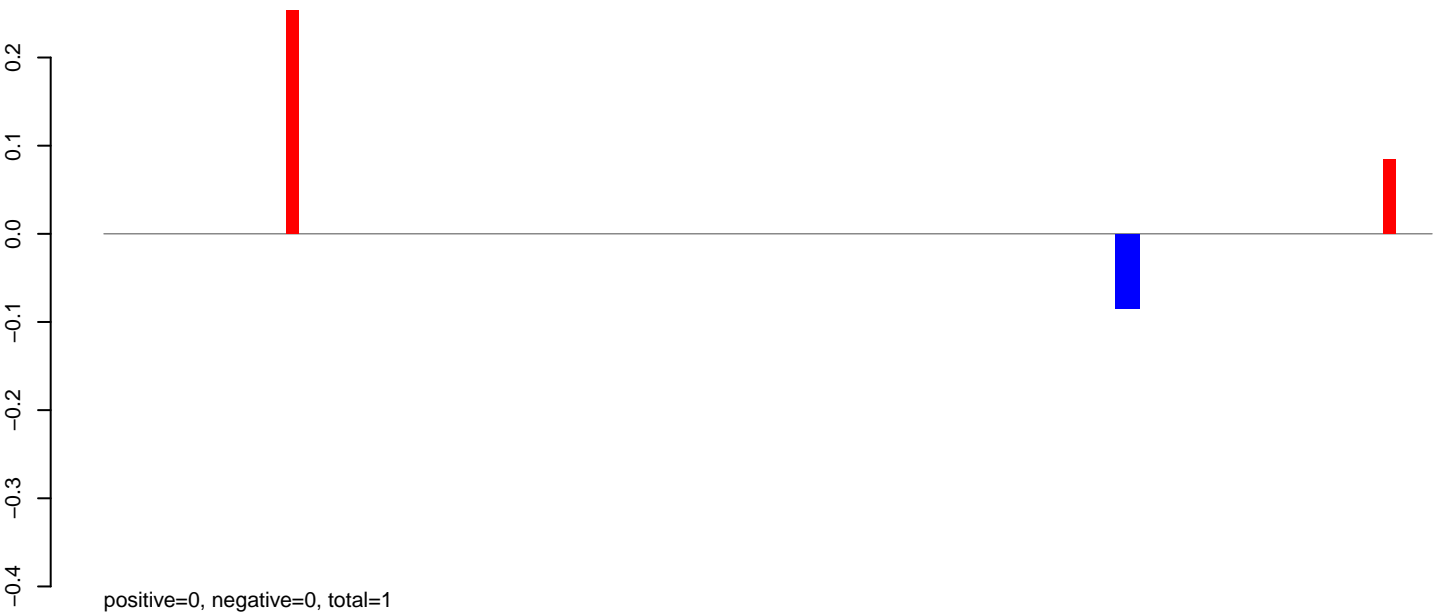
AnGam_Mos55_cells.rep



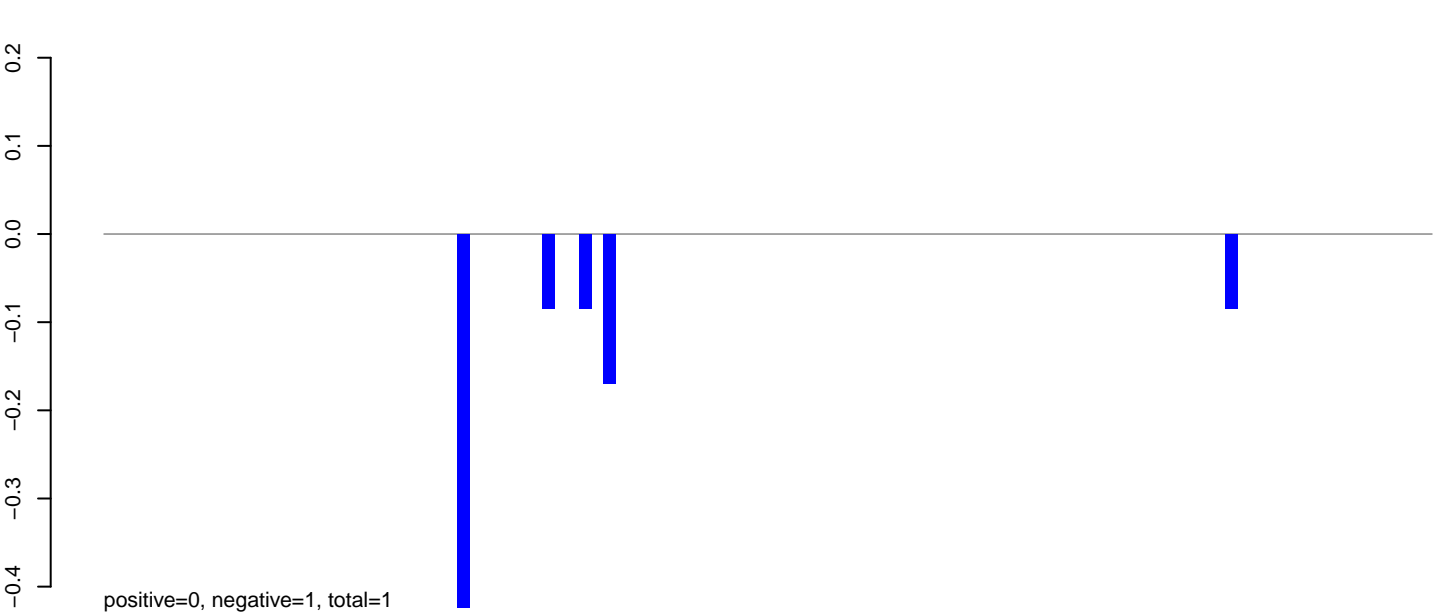
positive=0, negative=1, total=1

Window size=50, length=1492, TE@Agam_U11659_mariner-DNA-mariner:1-1492

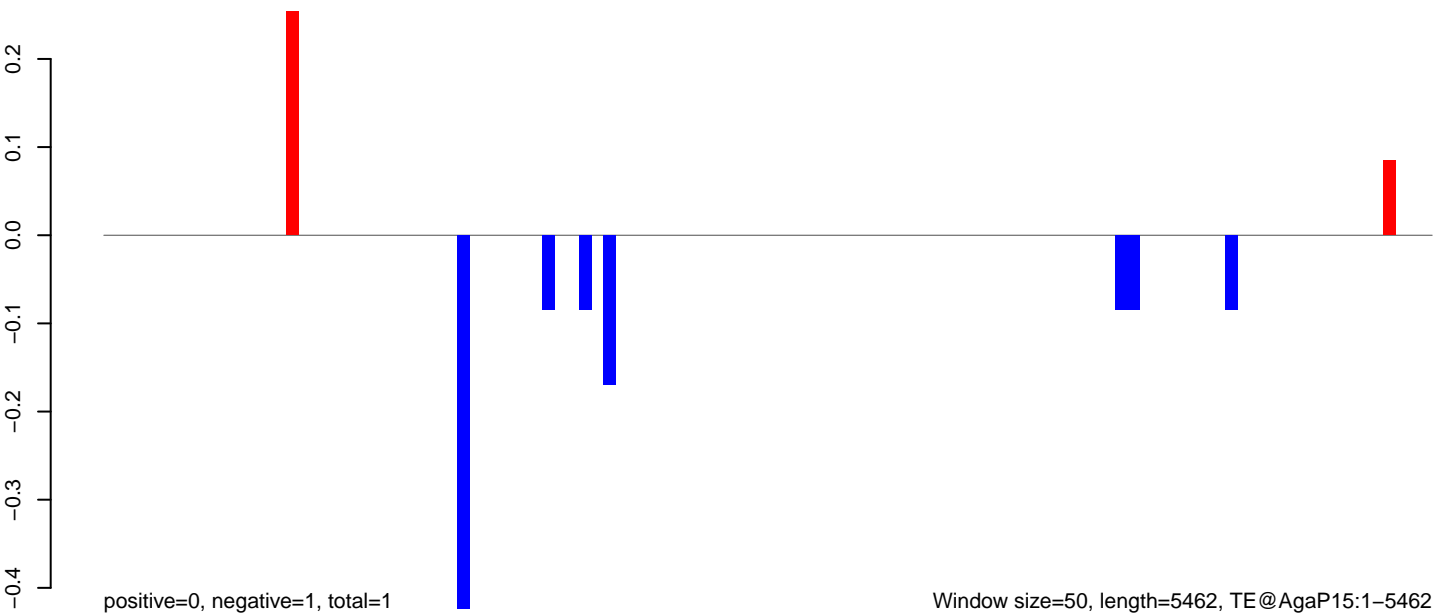
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



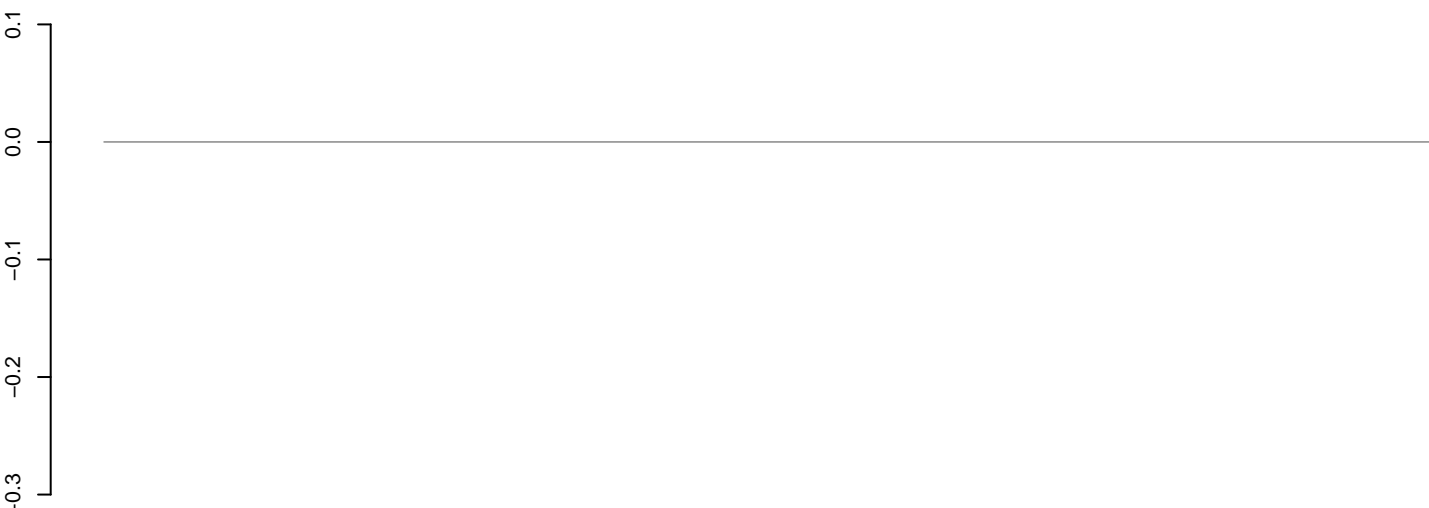
AnGam_Mos55_cells.rep



Window size=50, length=5462, TE@AgaP15:1-5462

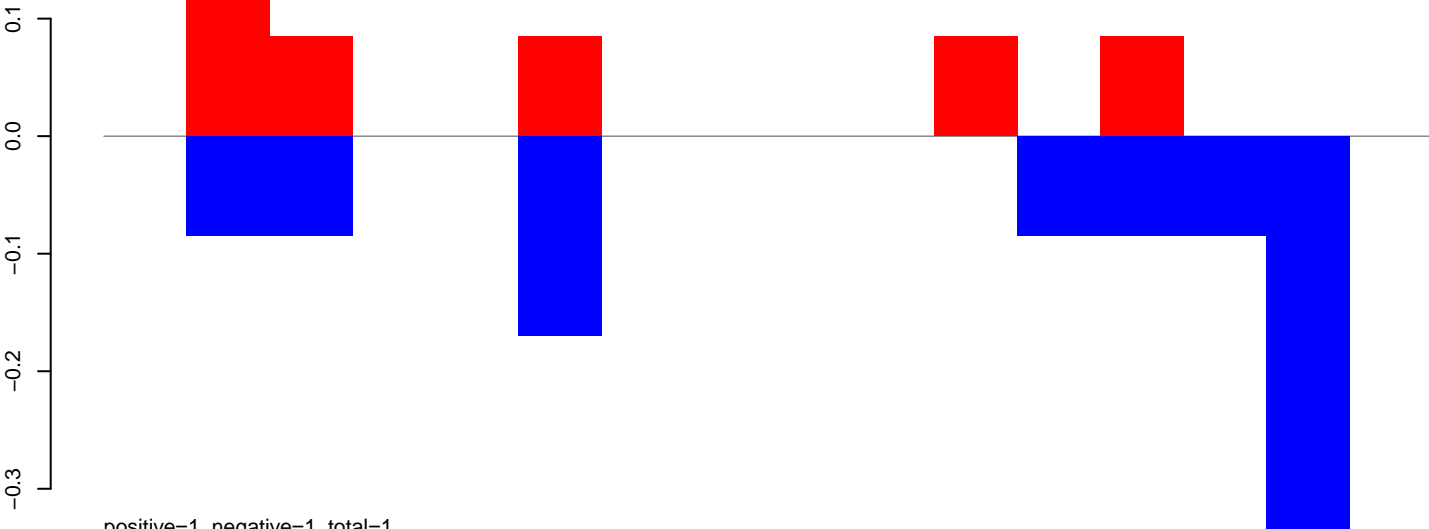
0 1000 2000 3000 4000 5000

AnGam_Mos55_cells.18_23.rep



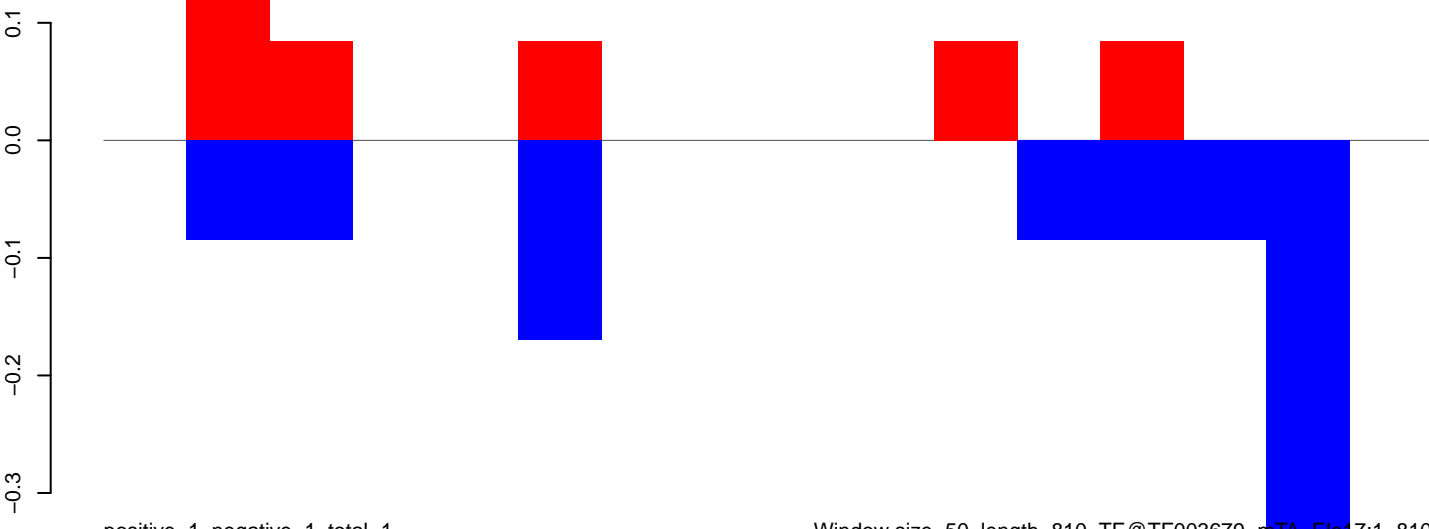
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=1, negative=1, total=1

AnGam_Mos55_cells.rep



positive=1, negative=1, total=1

Window size=50, length=810, TE@TF003679-nTA_Ele17:1-810

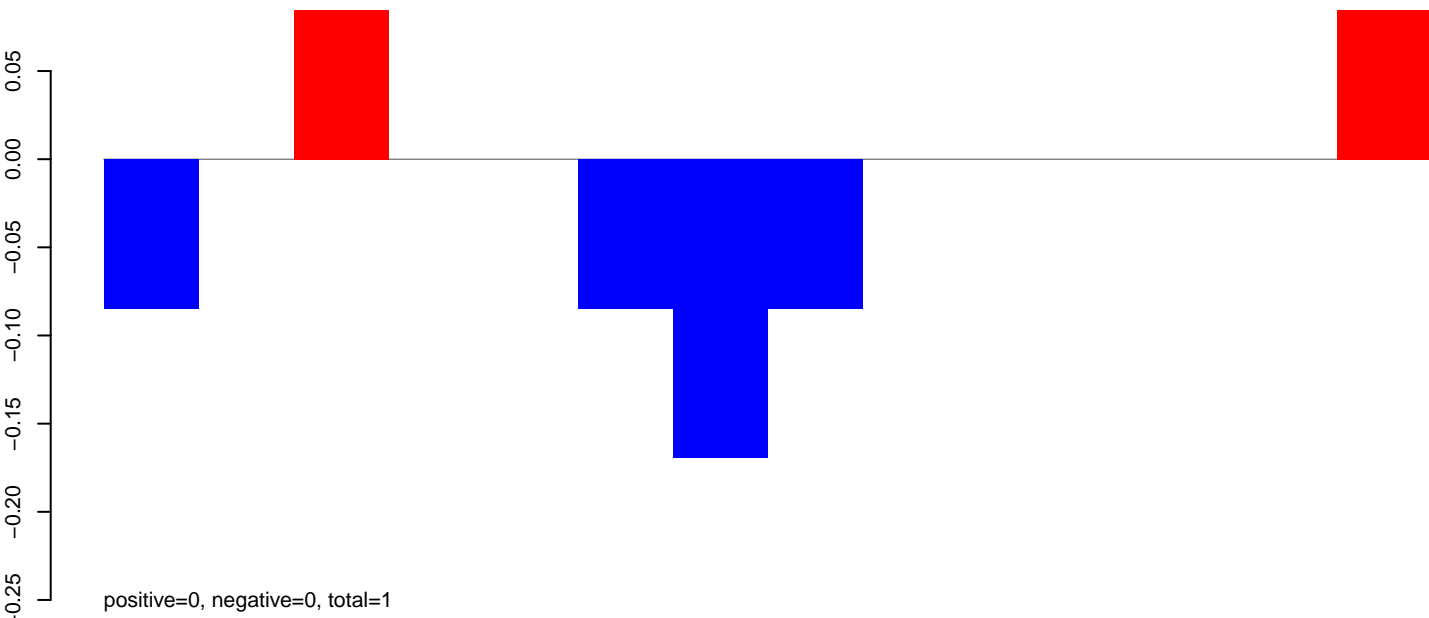
200

400

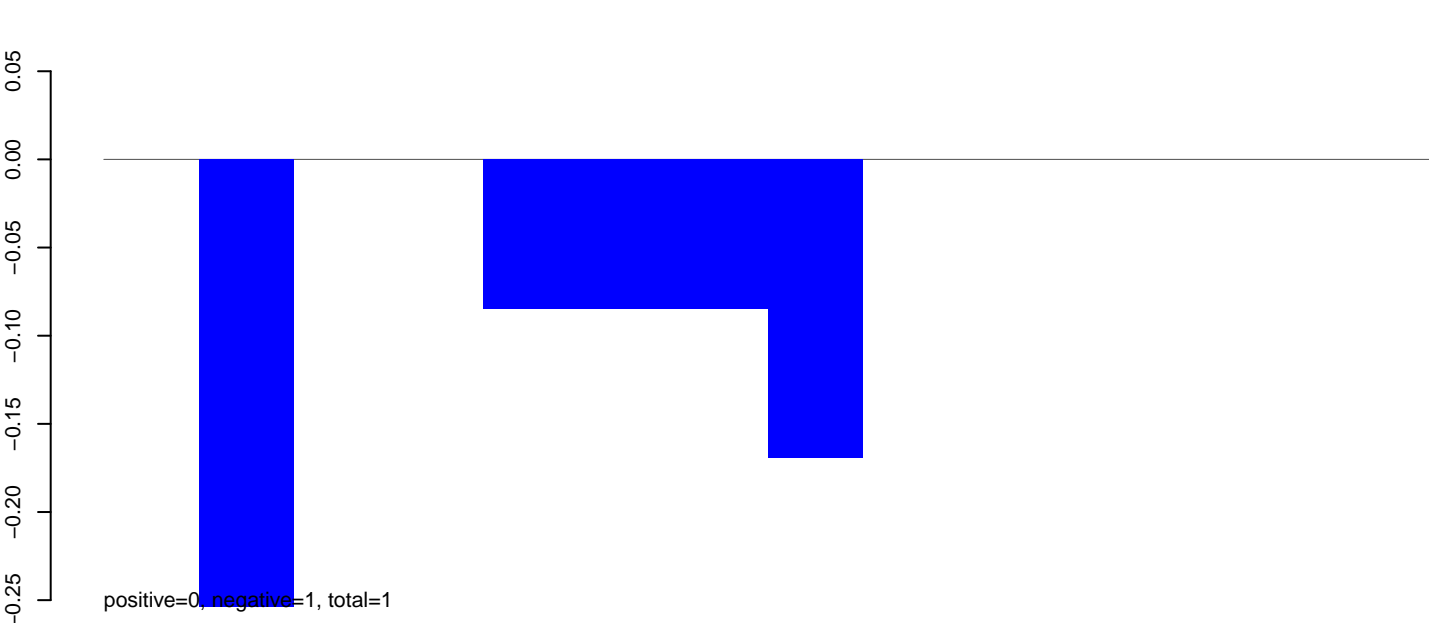
600

800

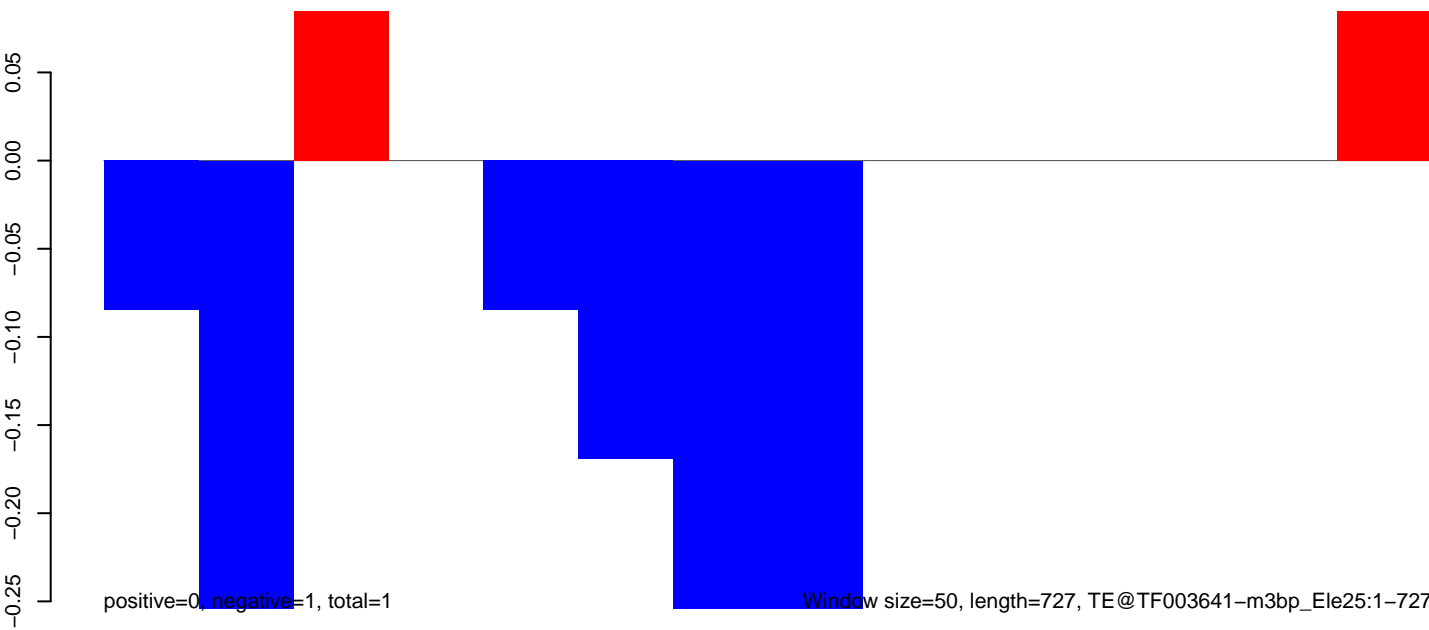
AnGam_Mos55_cells.18_23.rep



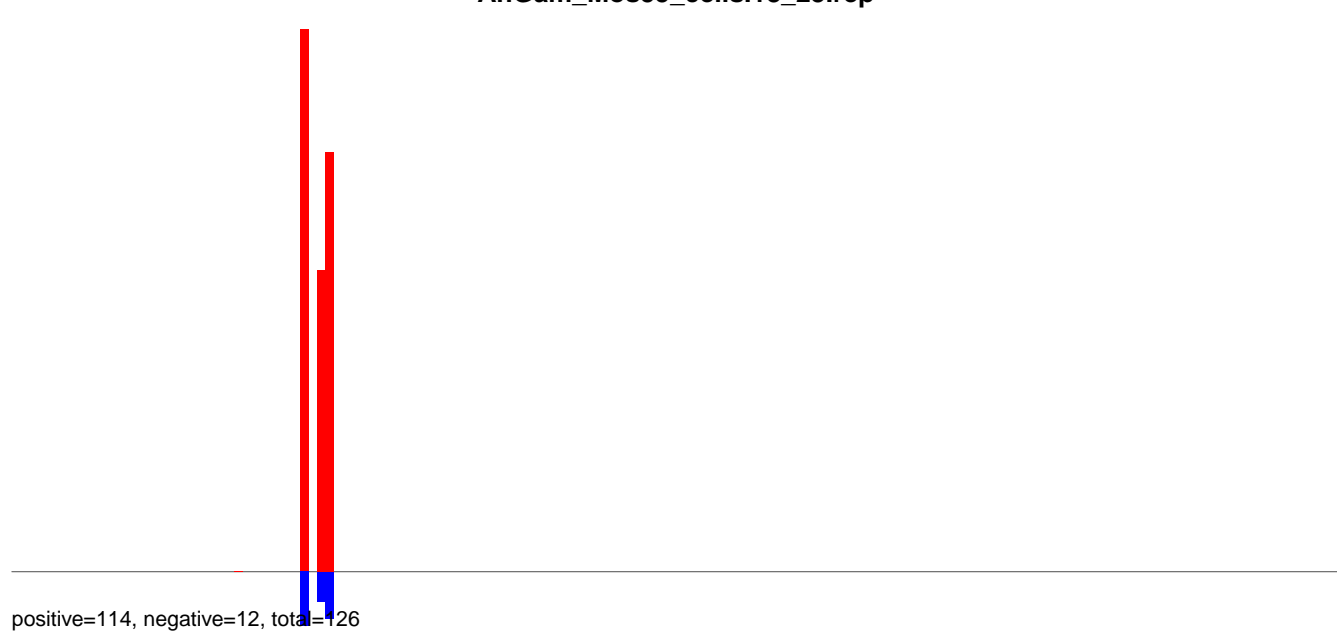
AnGam_Mos55_cells.24_35.rep



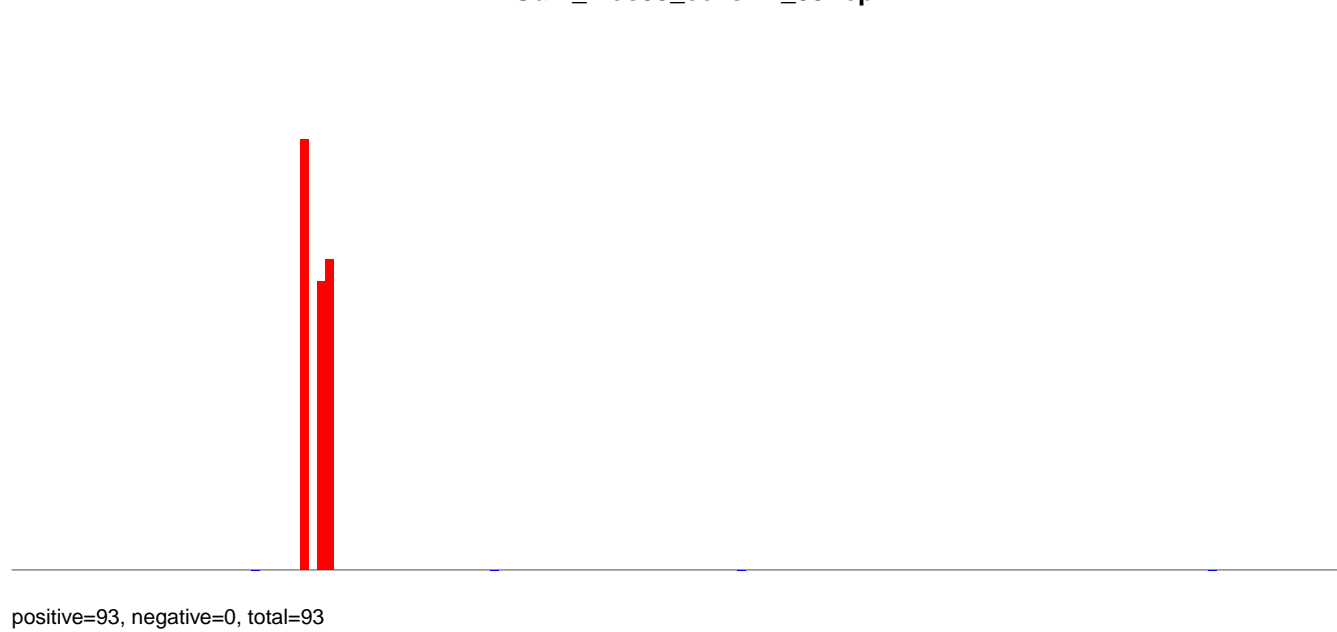
AnGam_Mos55_cells.rep



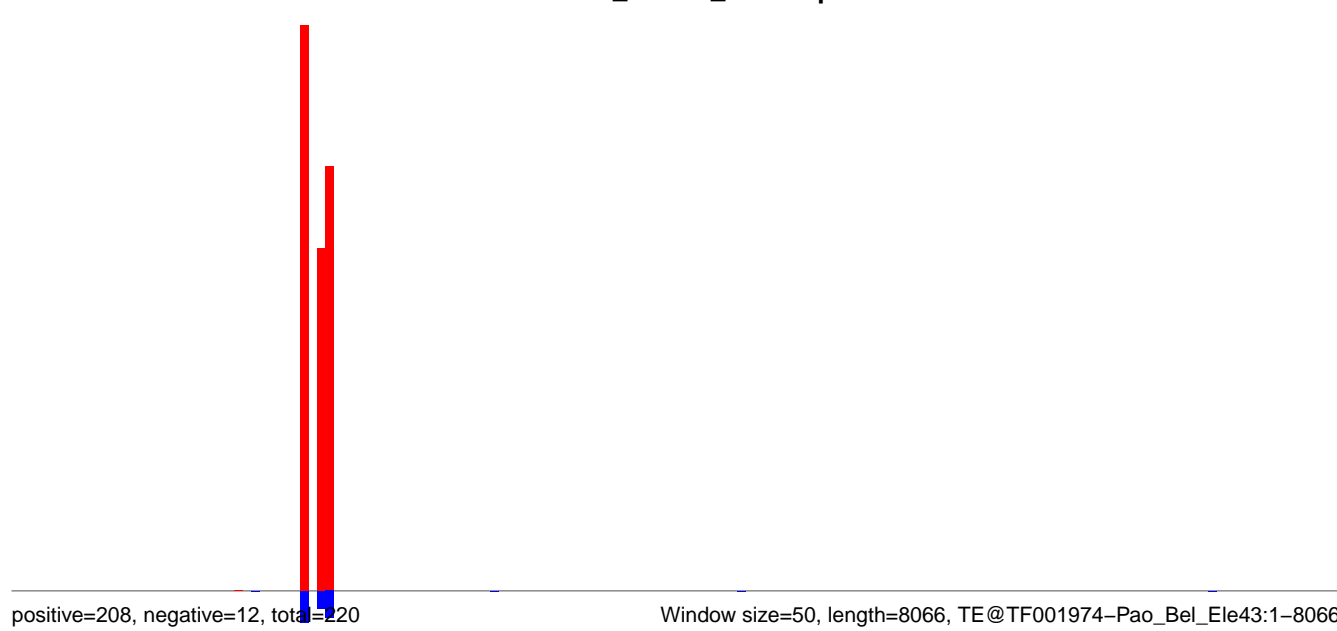
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



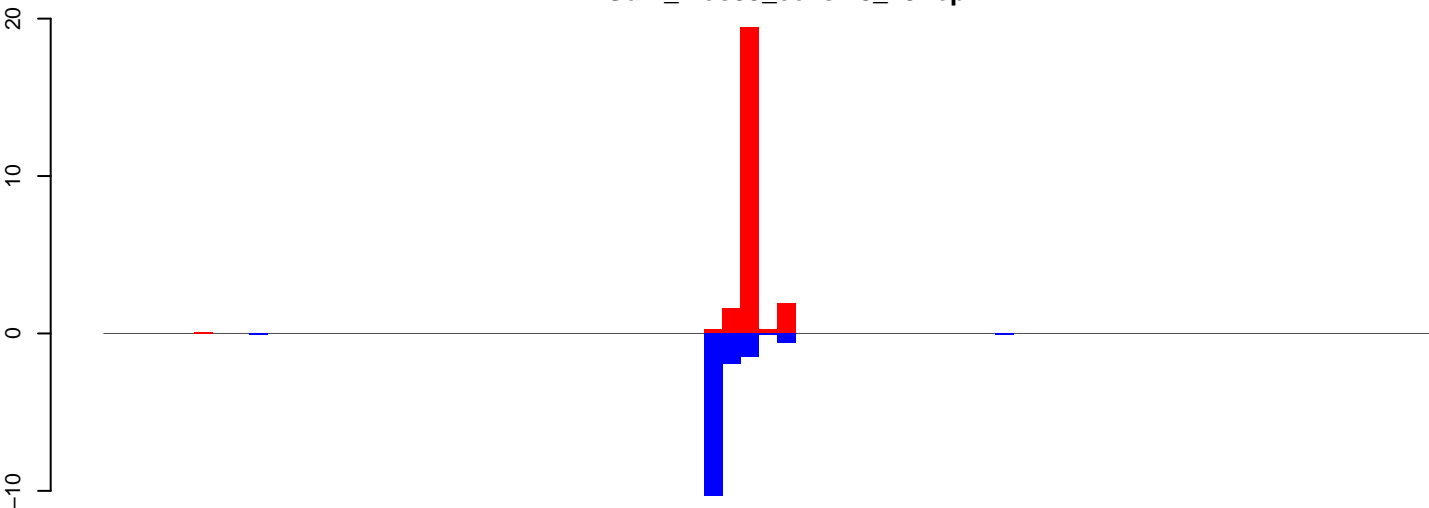
AnGam_Mos55_cells.rep



Window size=50, length=8066, TE@TF001974-Pao_Bel_Ele43:1-8066

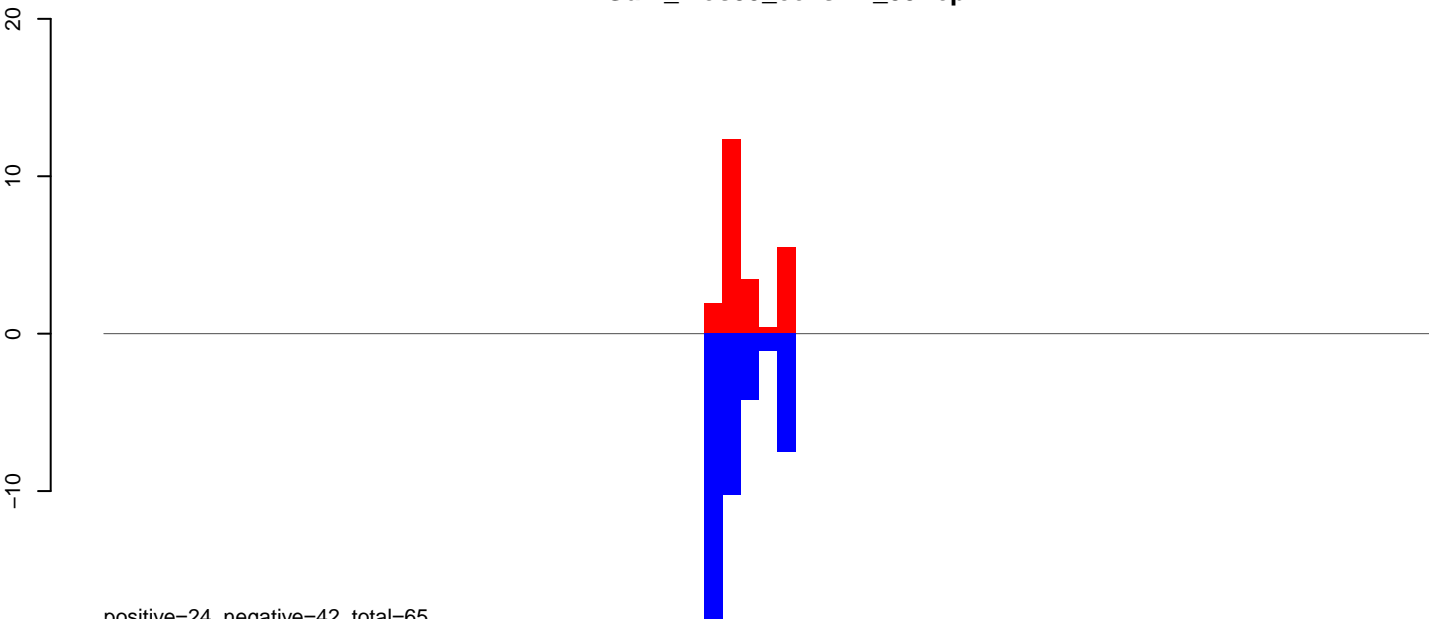
0 2000 4000 6000 8000

AnGam_Mos55_cells.18_23.rep



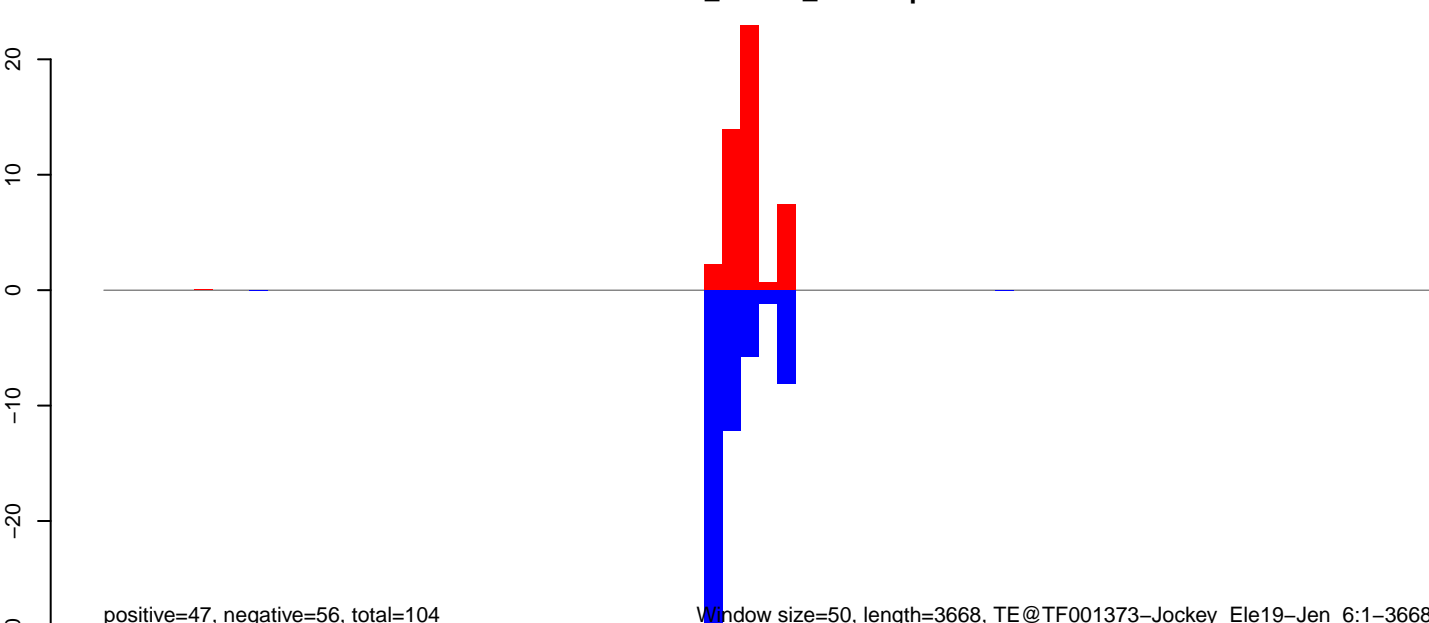
positive=24, negative=15, total=38

AnGam_Mos55_cells.24_35.rep



positive=24, negative=42, total=65

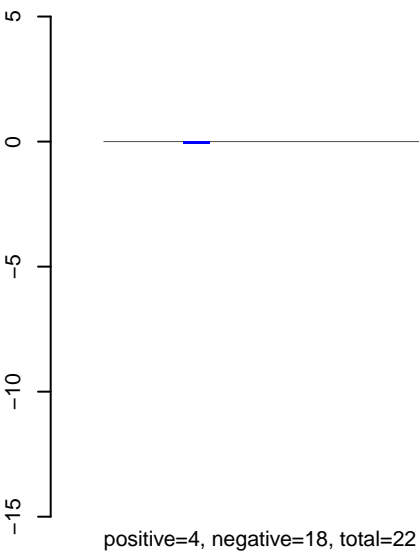
AnGam_Mos55_cells.rep



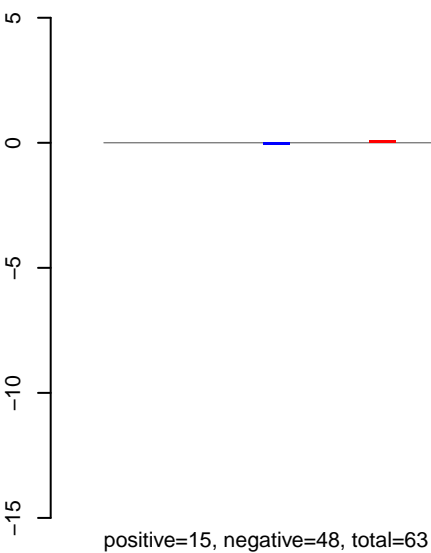
positive=47, negative=56, total=104

Window size=50, length=3668, TE@TF001373-Jockey_Ele19-Jen_6:1-3668

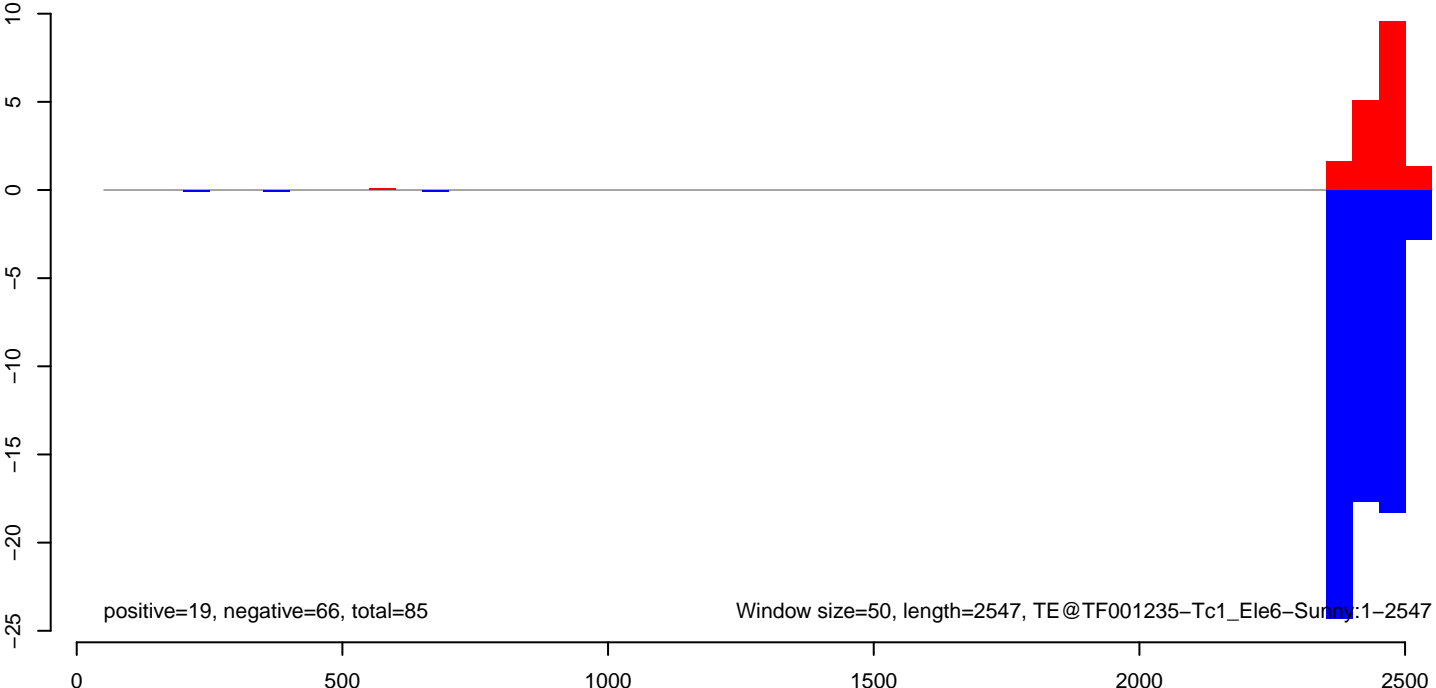
AnGam_Mos55_cells.18_23.rep



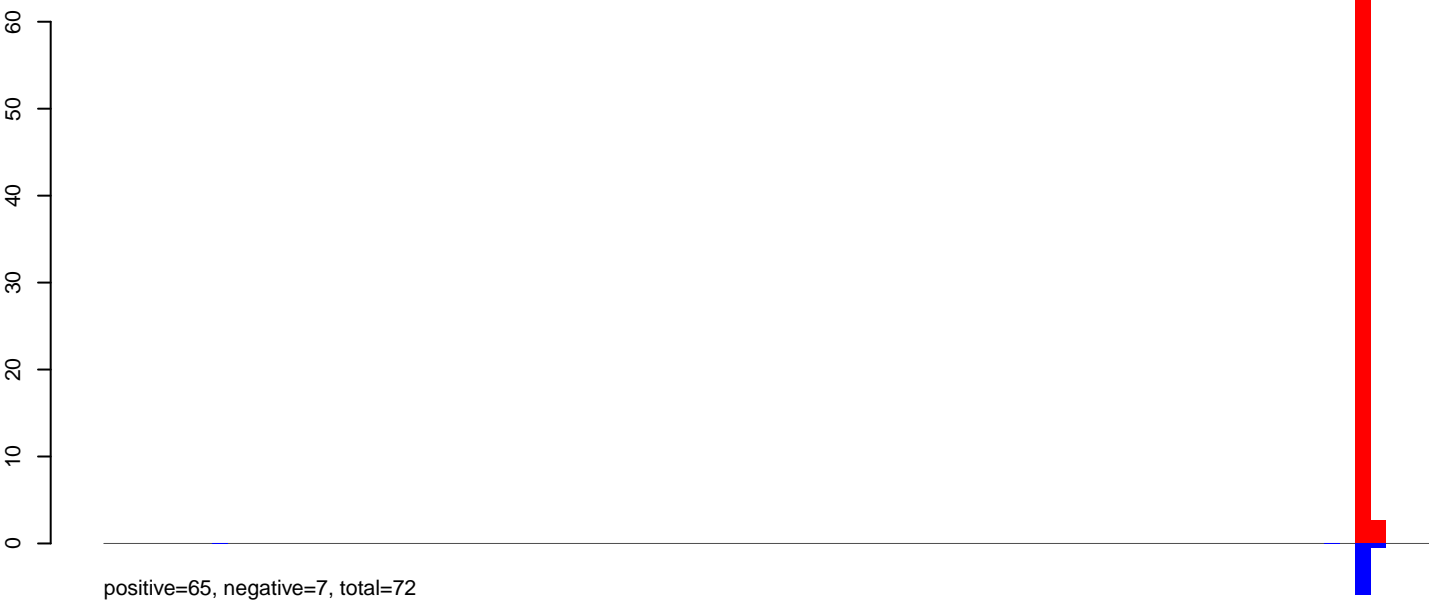
AnGam_Mos55_cells.24_35.rep



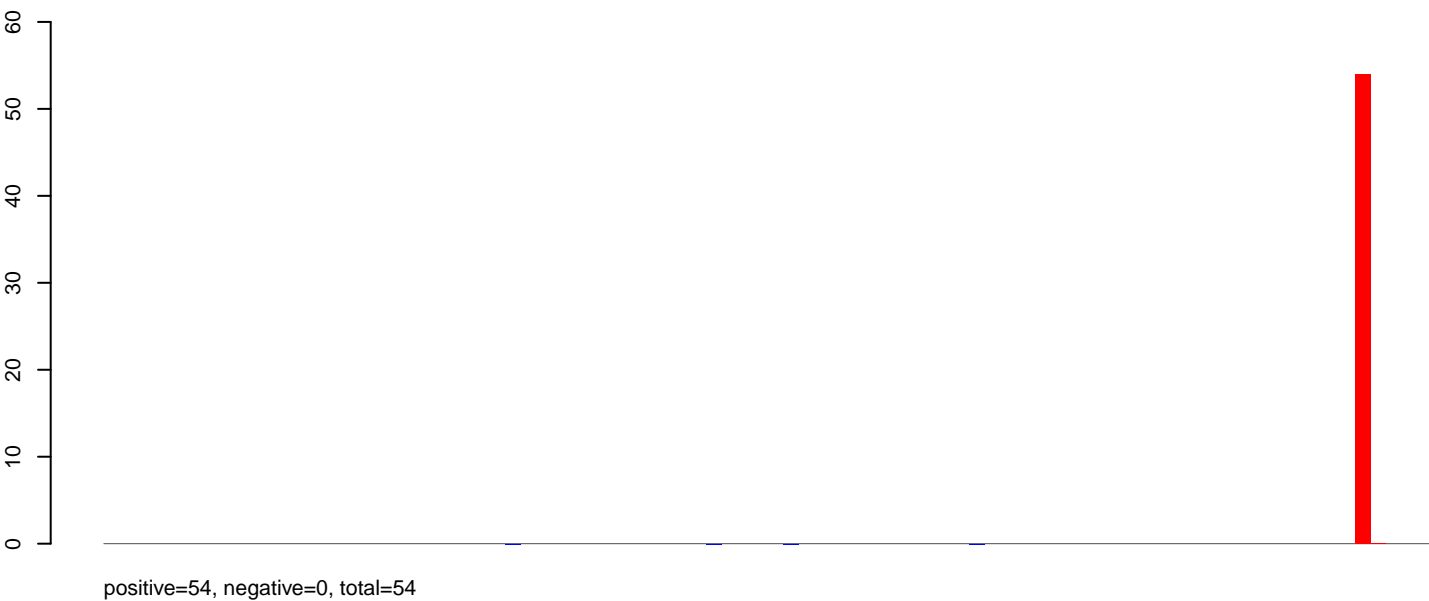
AnGam_Mos55_cells.rep



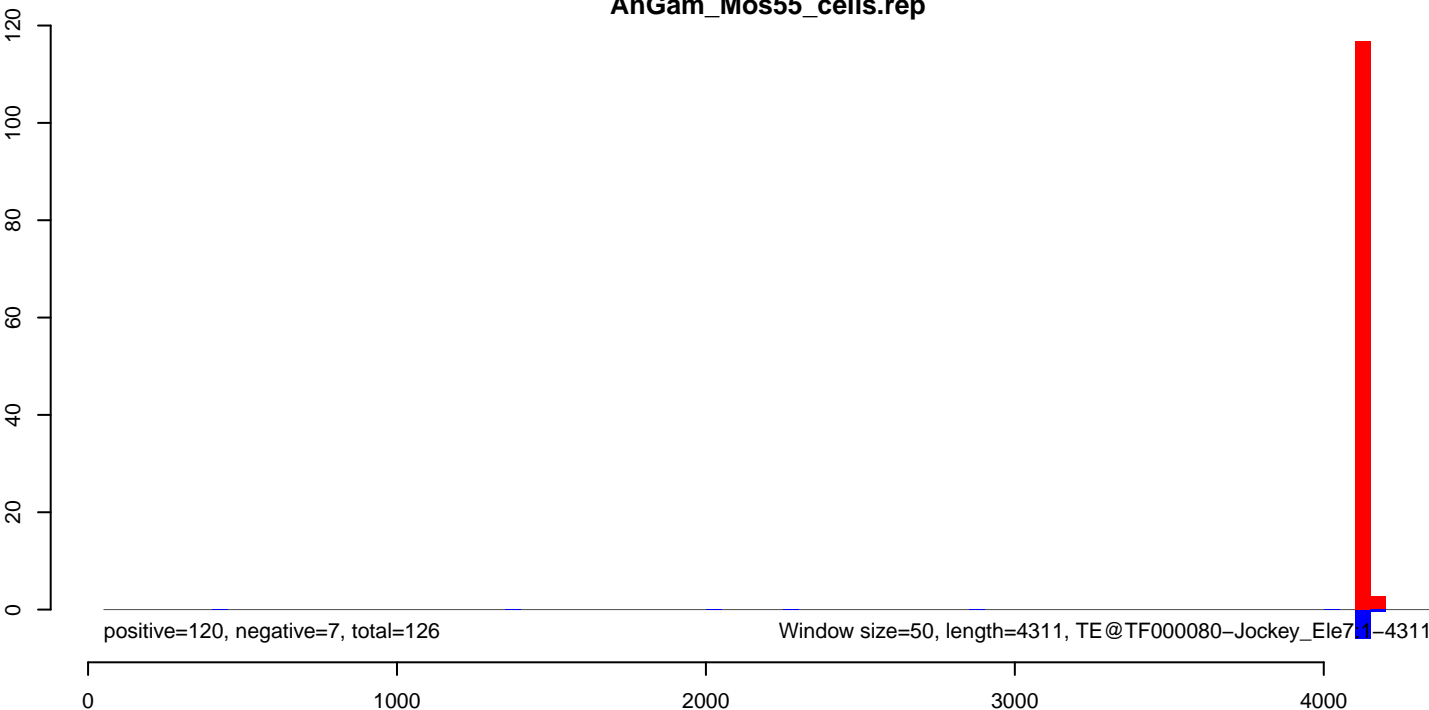
AnGam_Mos55_cells.18_23.rep



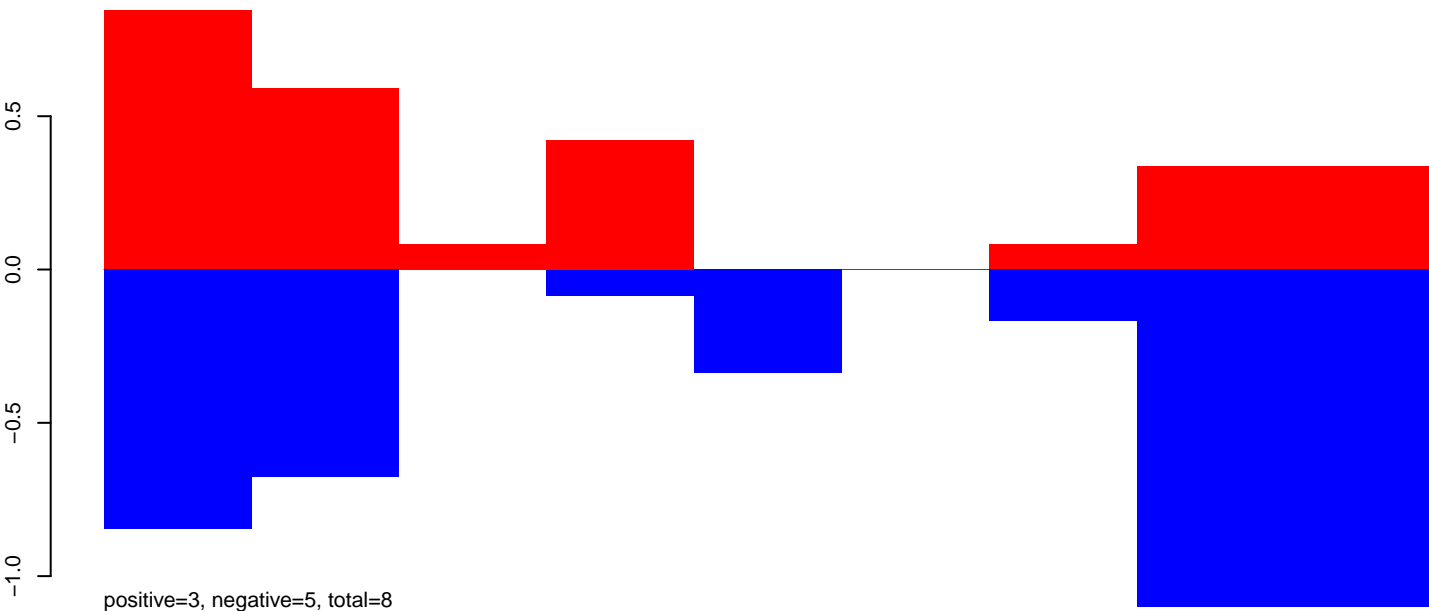
AnGam_Mos55_cells.24_35.rep



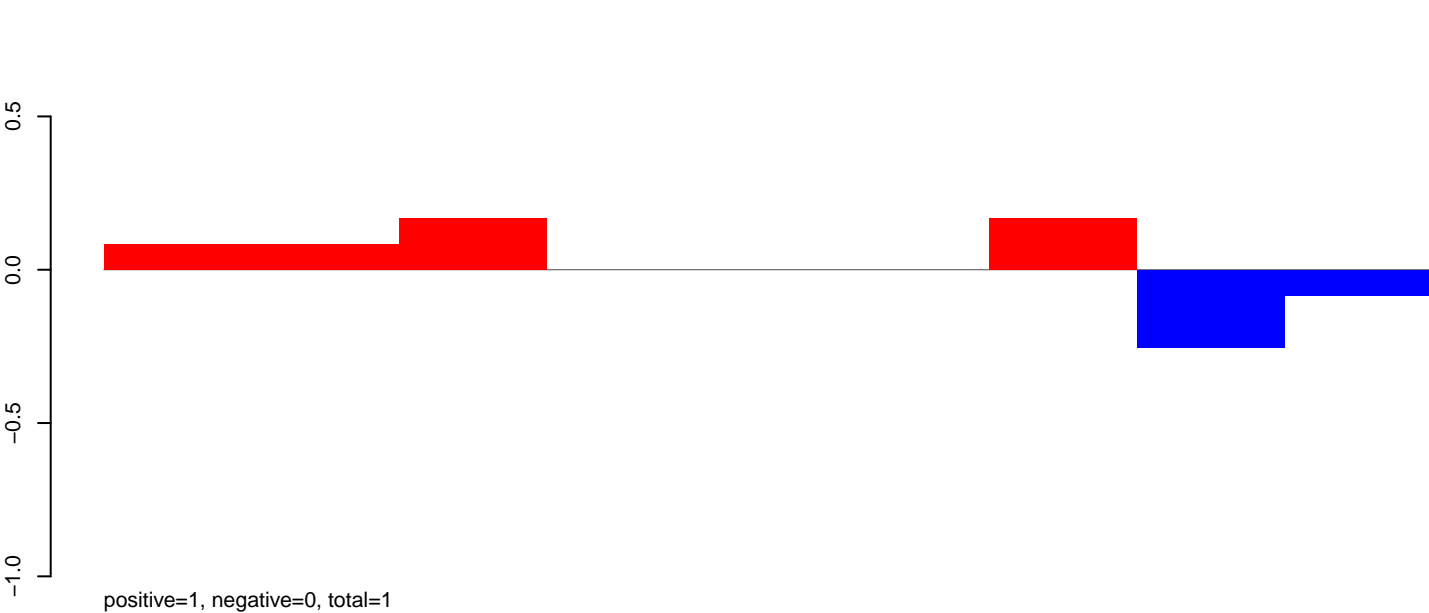
AnGam_Mos55_cells.rep



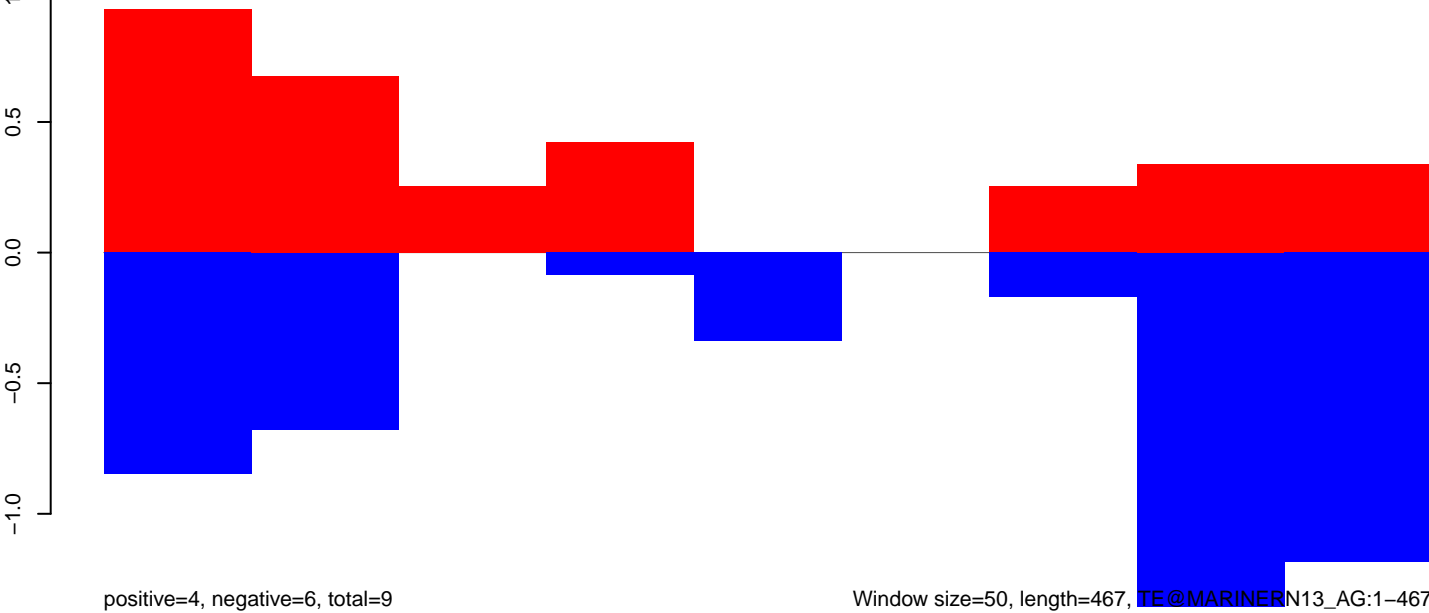
AnGam_Mos55_cells.18_23.rep



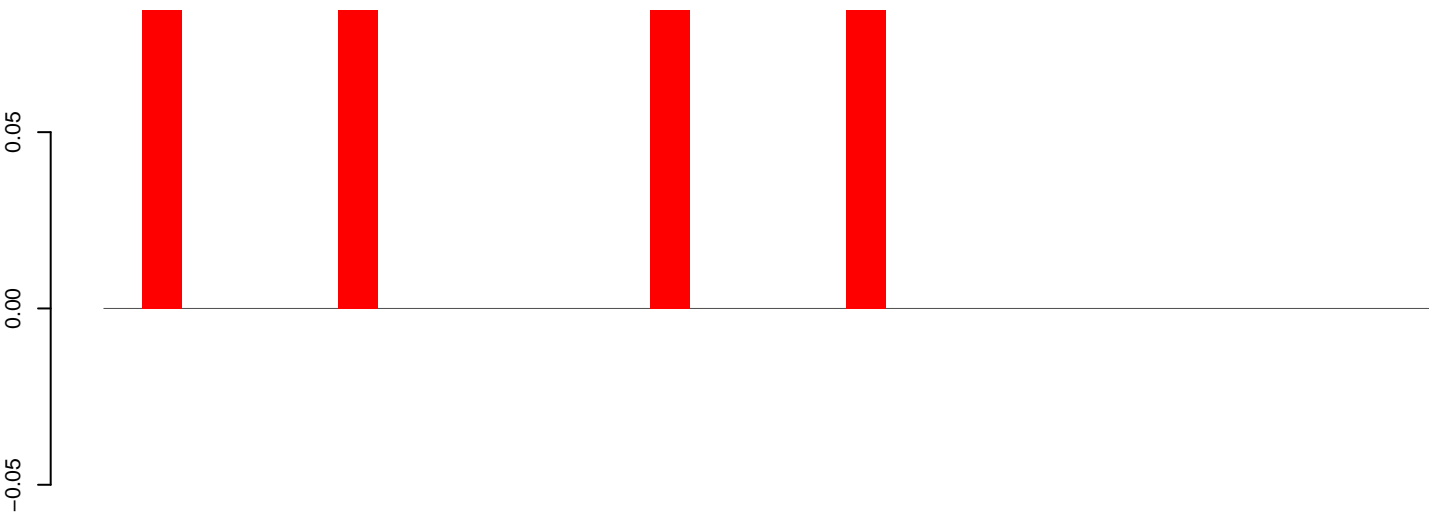
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

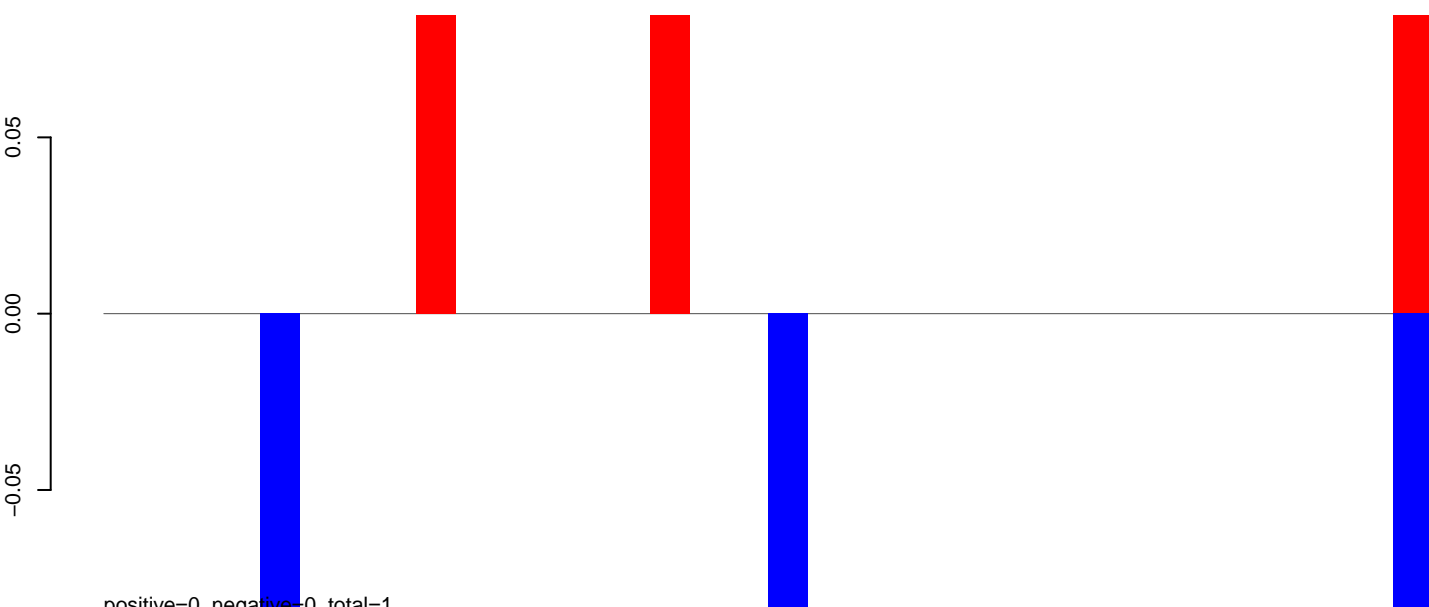


AnGam_Mos55_cells.18_23.rep



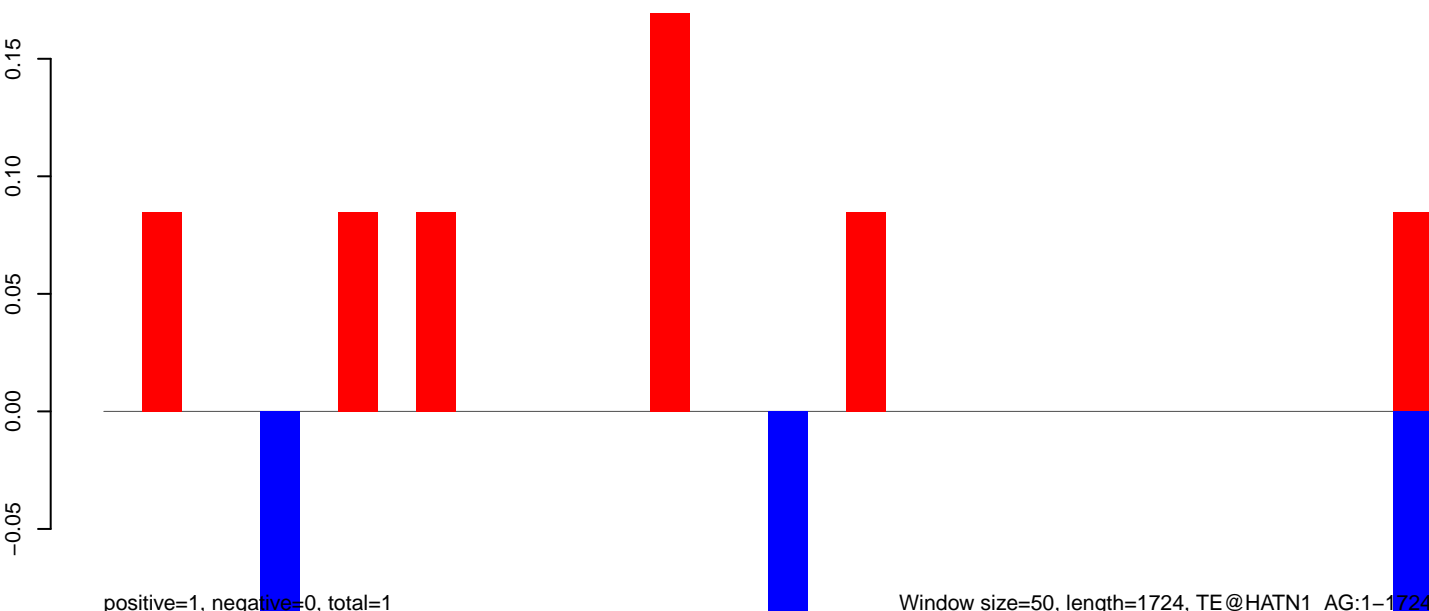
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



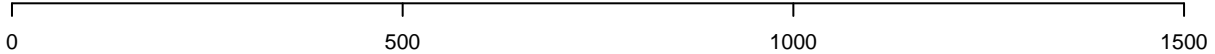
positive=0, negative=0, total=1

AnGam_Mos55_cells.rep

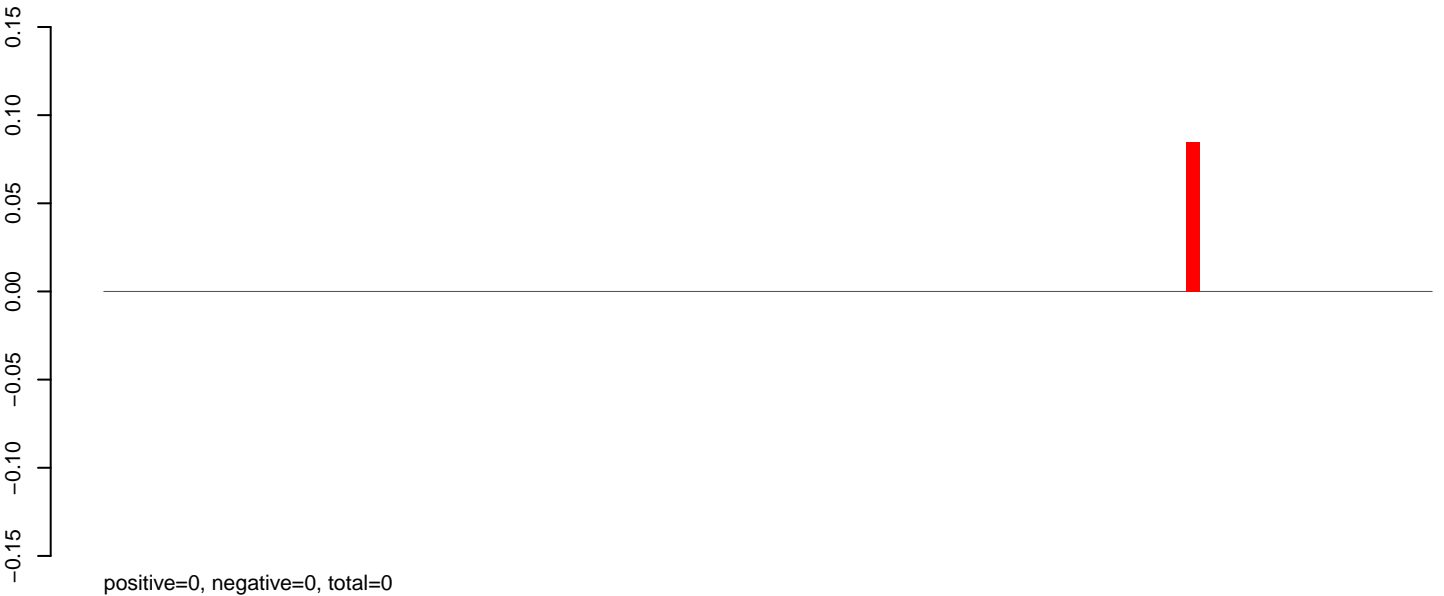


positive=1, negative=0, total=1

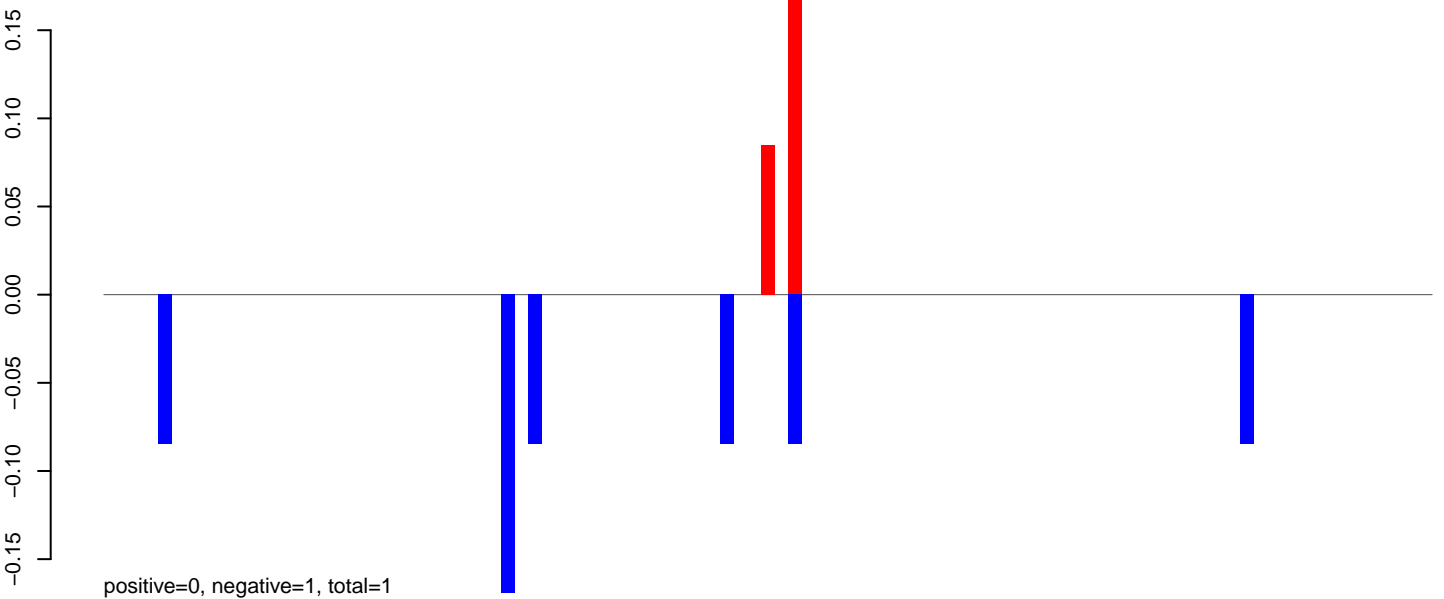
Window size=50, length=1724, TE@HATN1_AG:1-1724



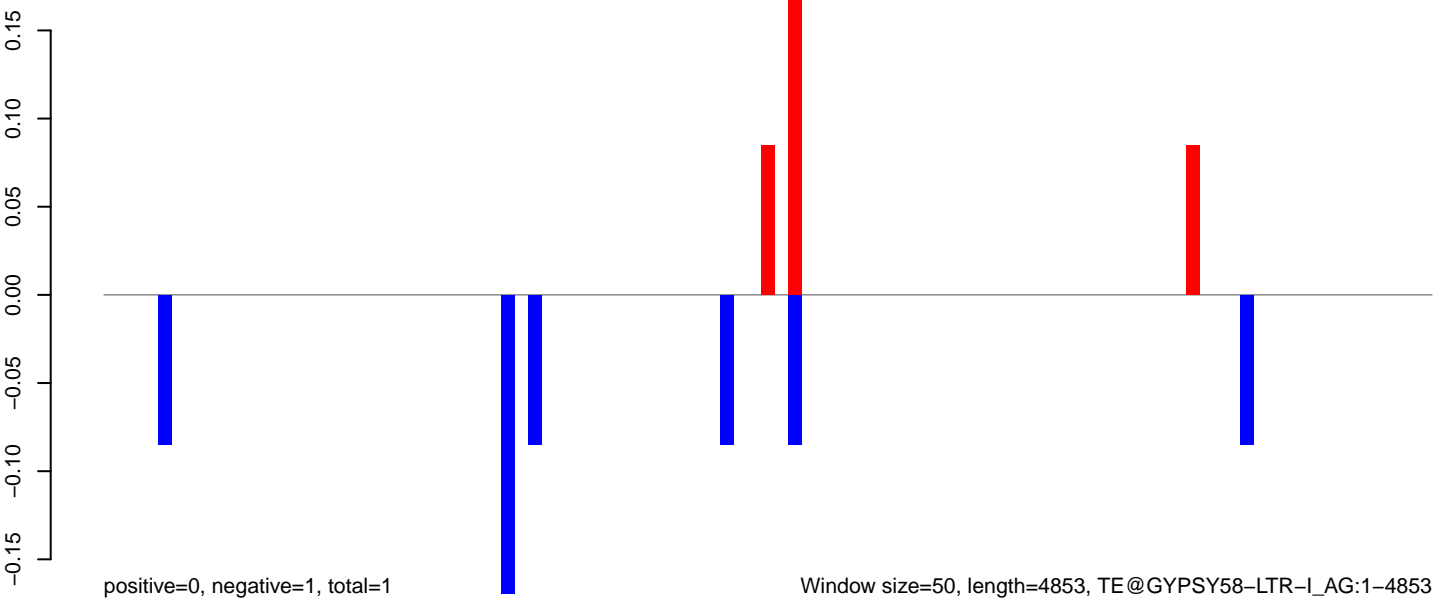
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



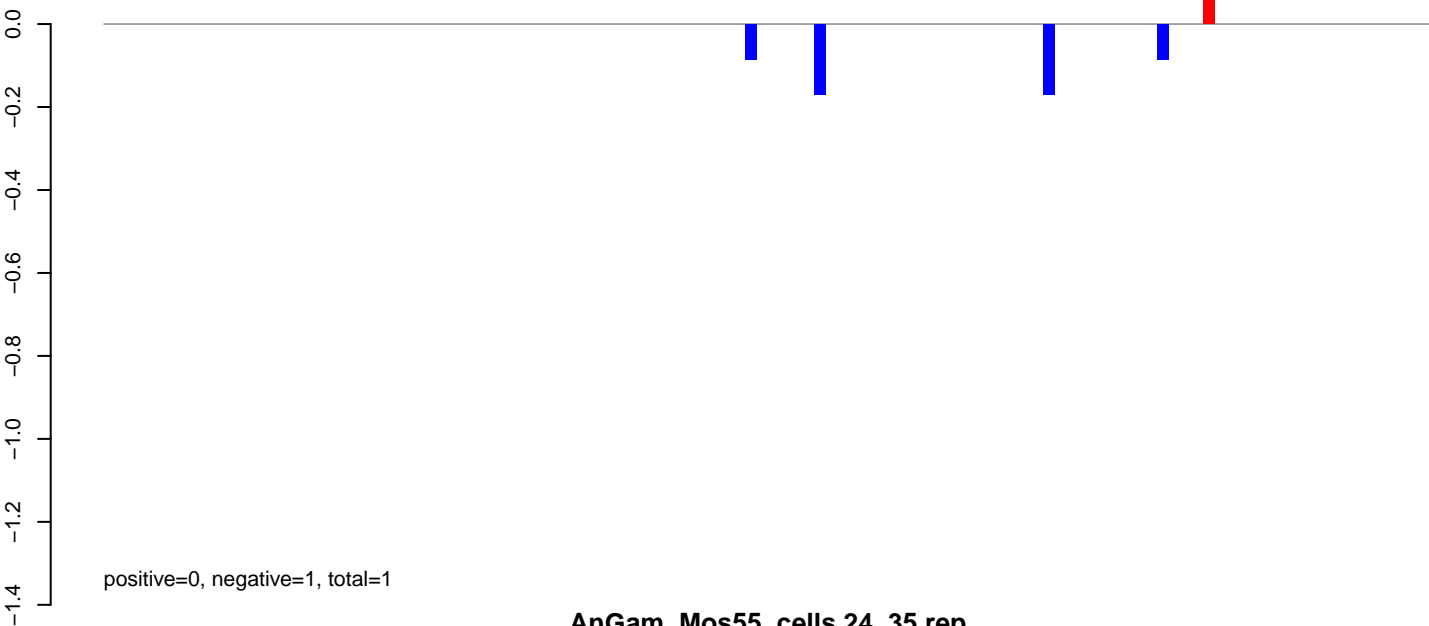
AnGam_Mos55_cells.rep



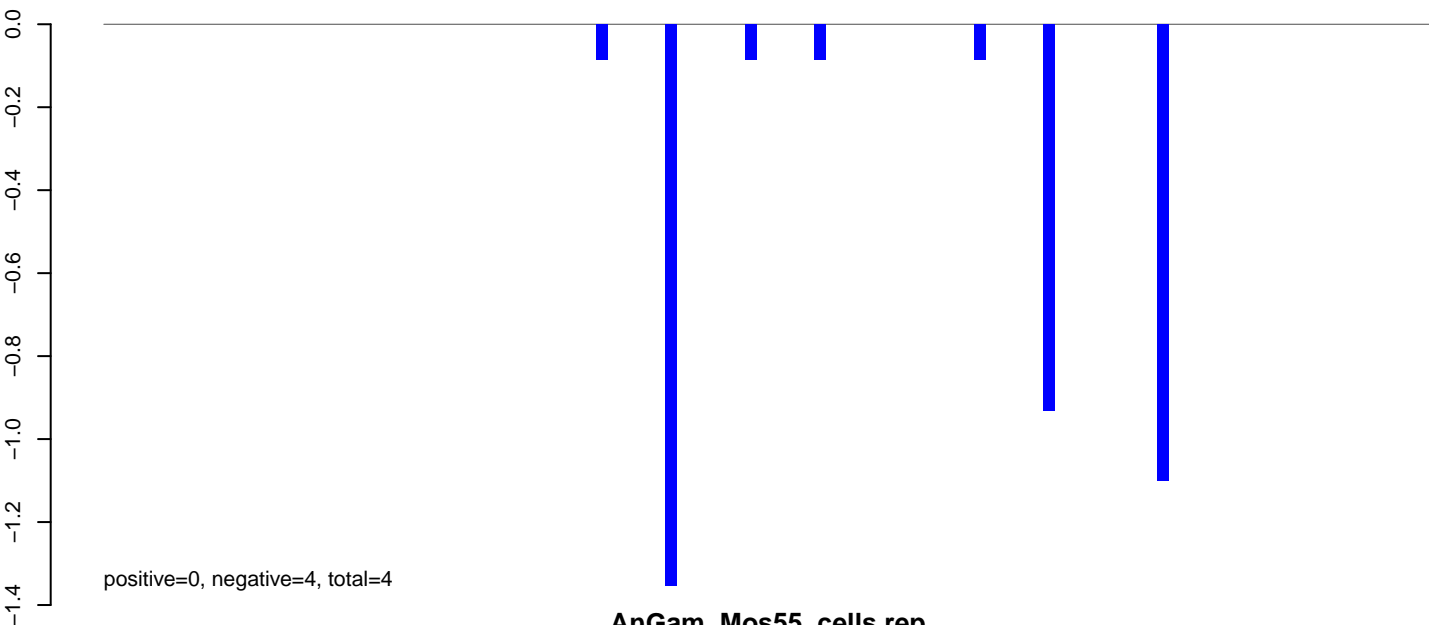
Window size=50, length=4853, TE@GYPSY58-LTR-L_AG:1-4853

0 1000 2000 3000 4000 5000

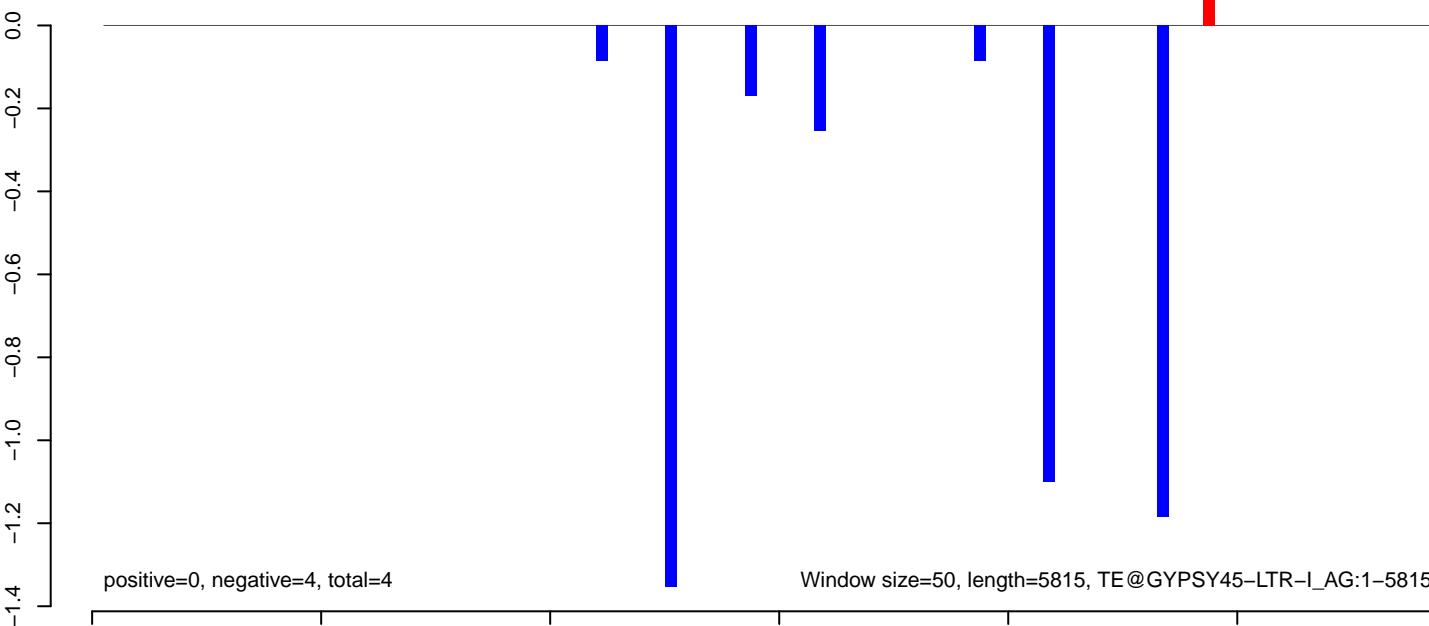
AnGam_Mos55_cells.18_23.rep



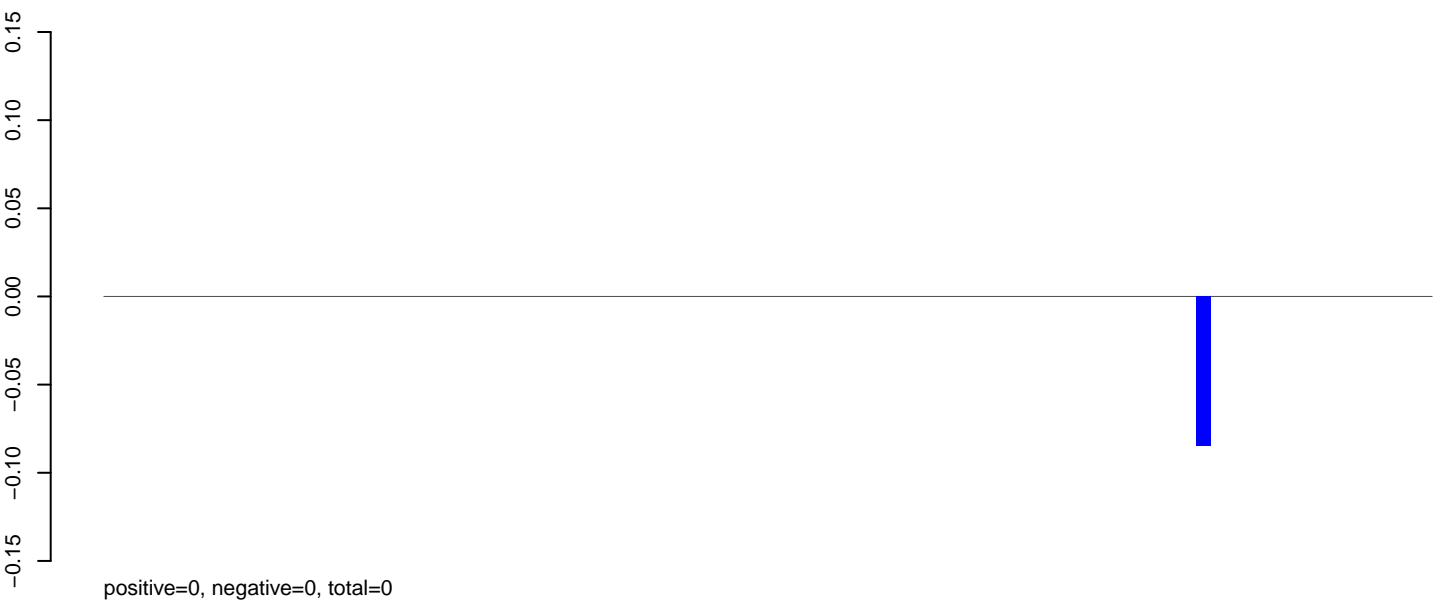
AnGam_Mos55_cells.24_35.rep



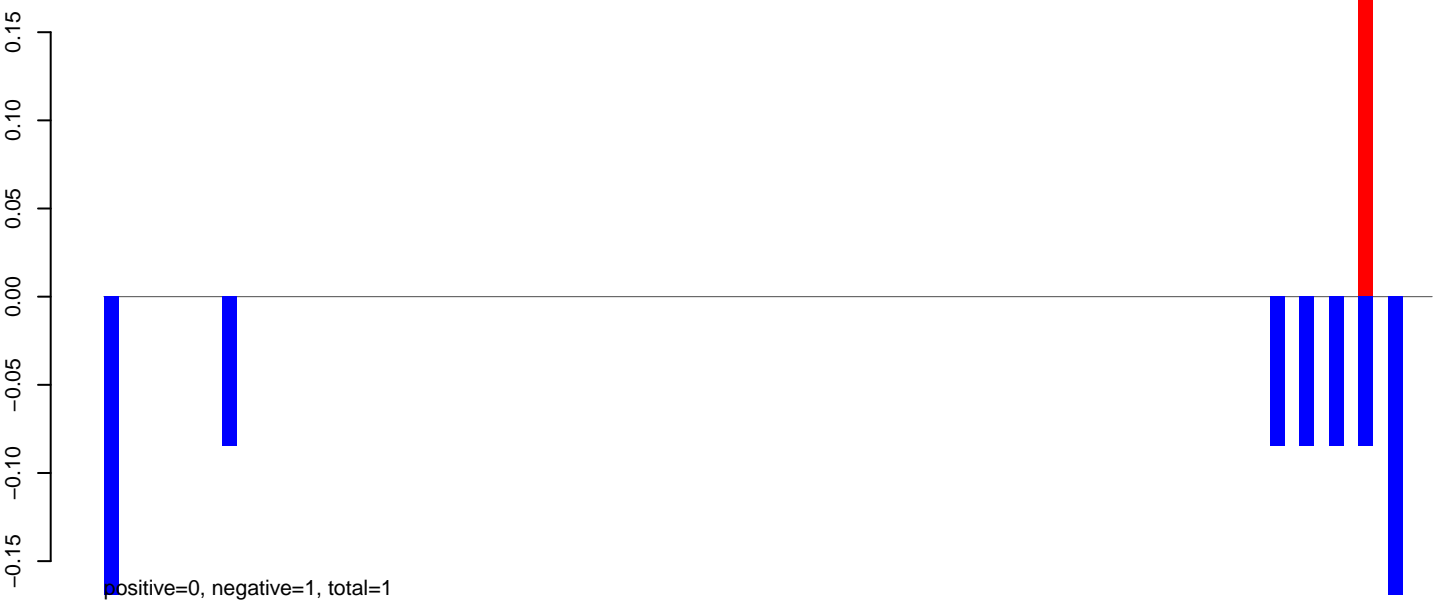
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



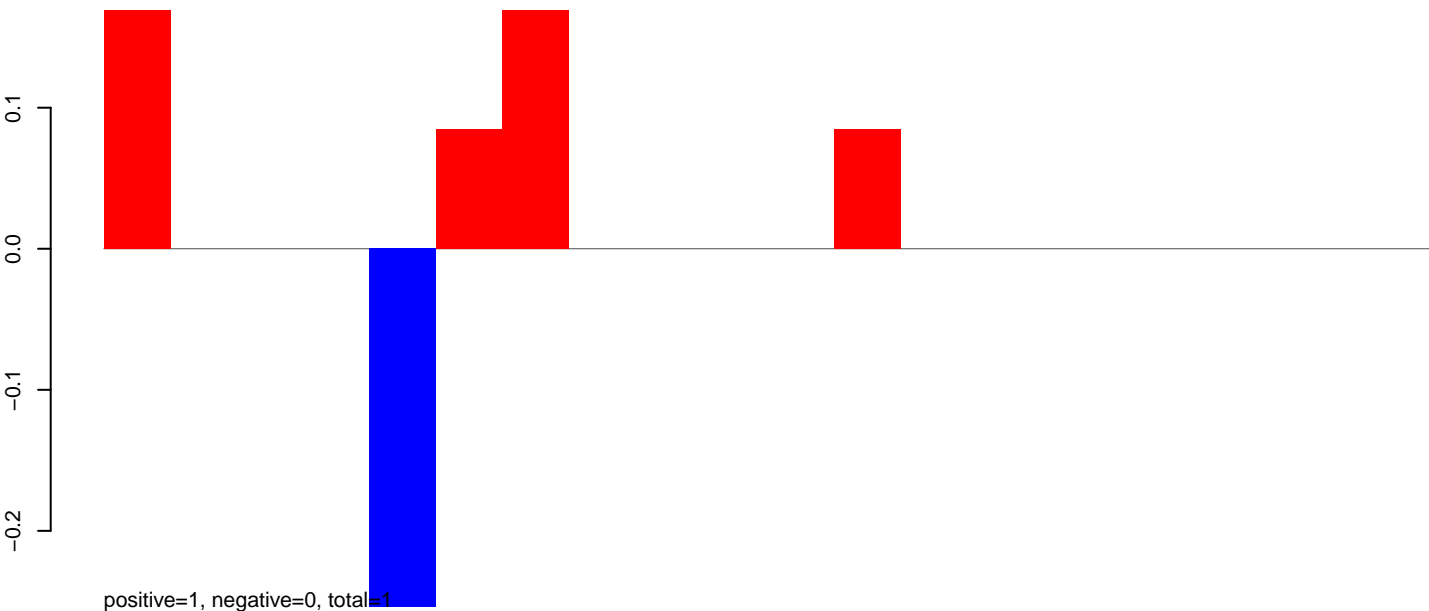
AnGam_Mos55_cells.24_35.rep



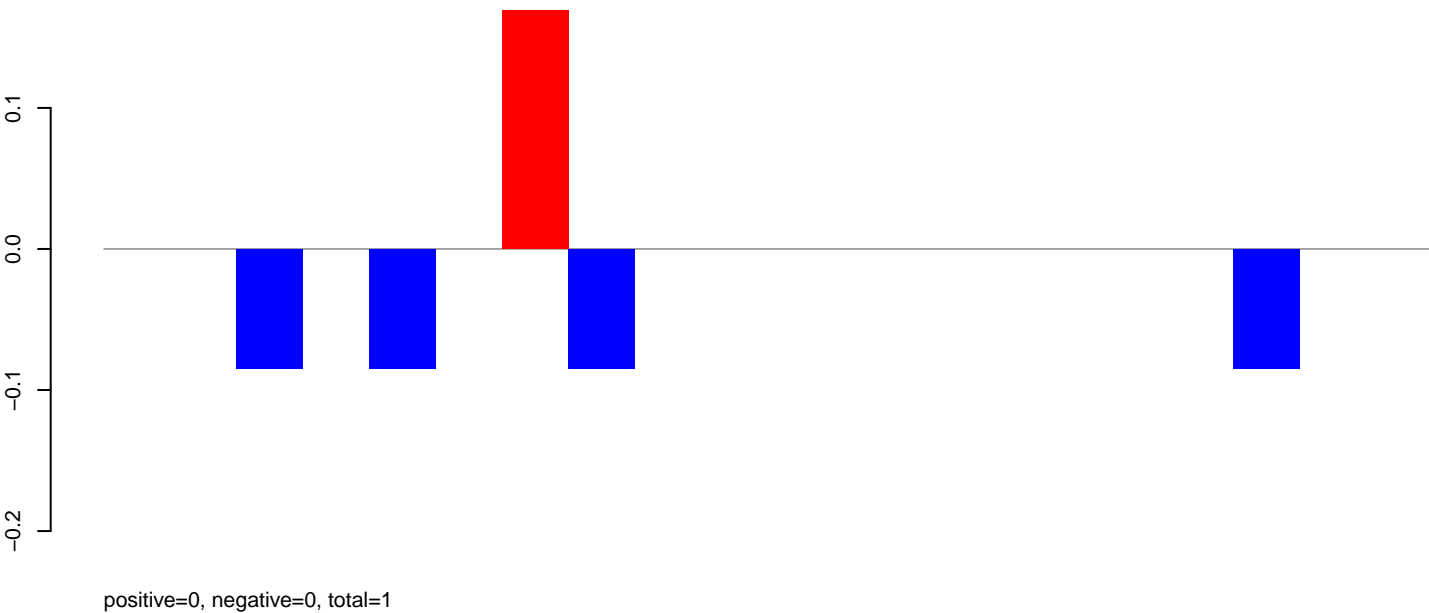
AnGam_Mos55_cells.rep



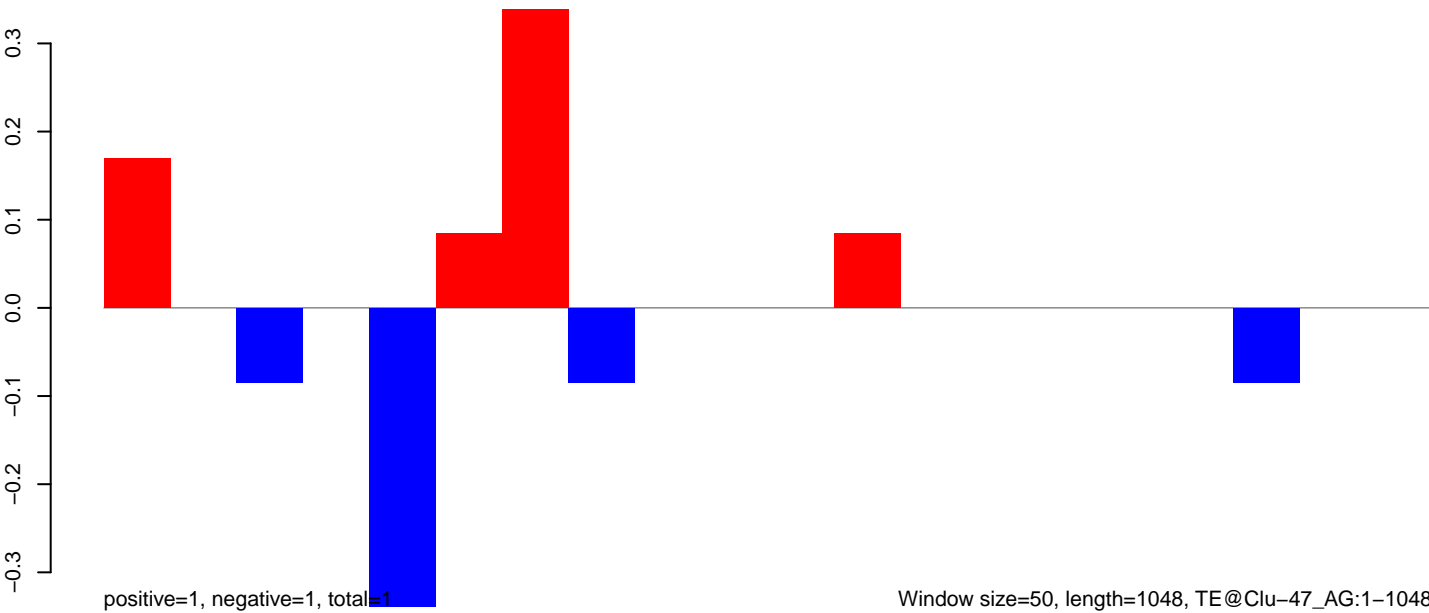
AnGam_Mos55_cells.18_23.rep



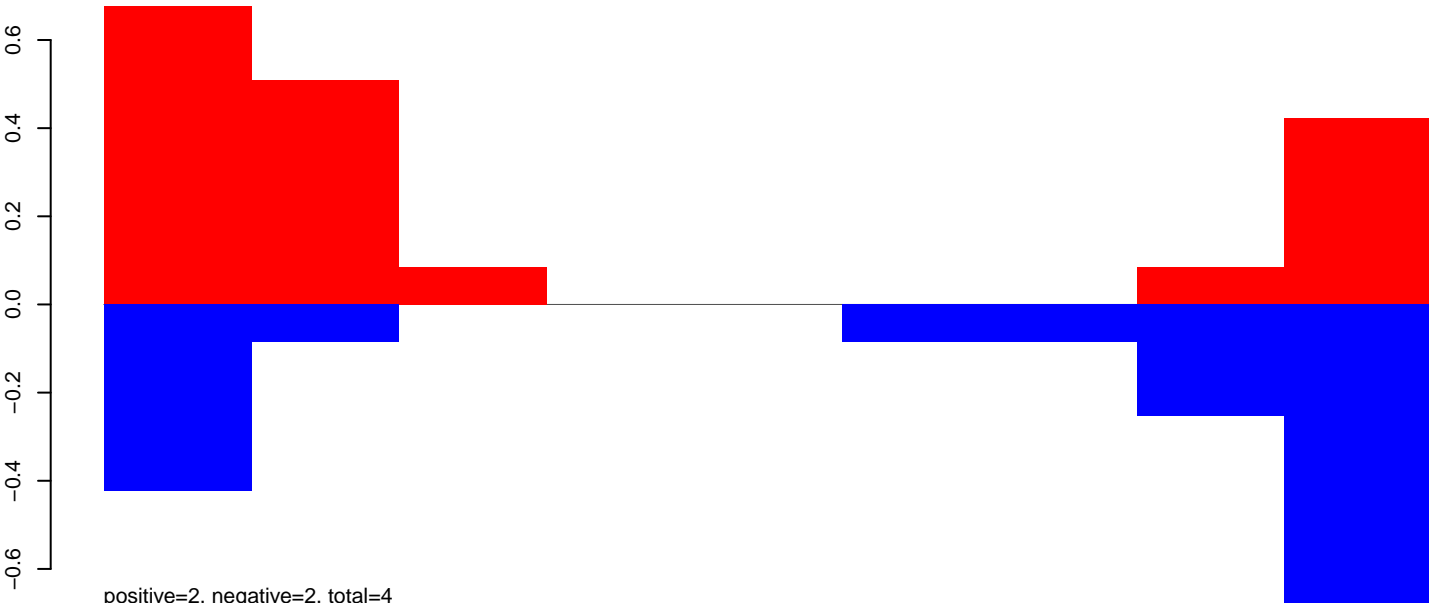
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



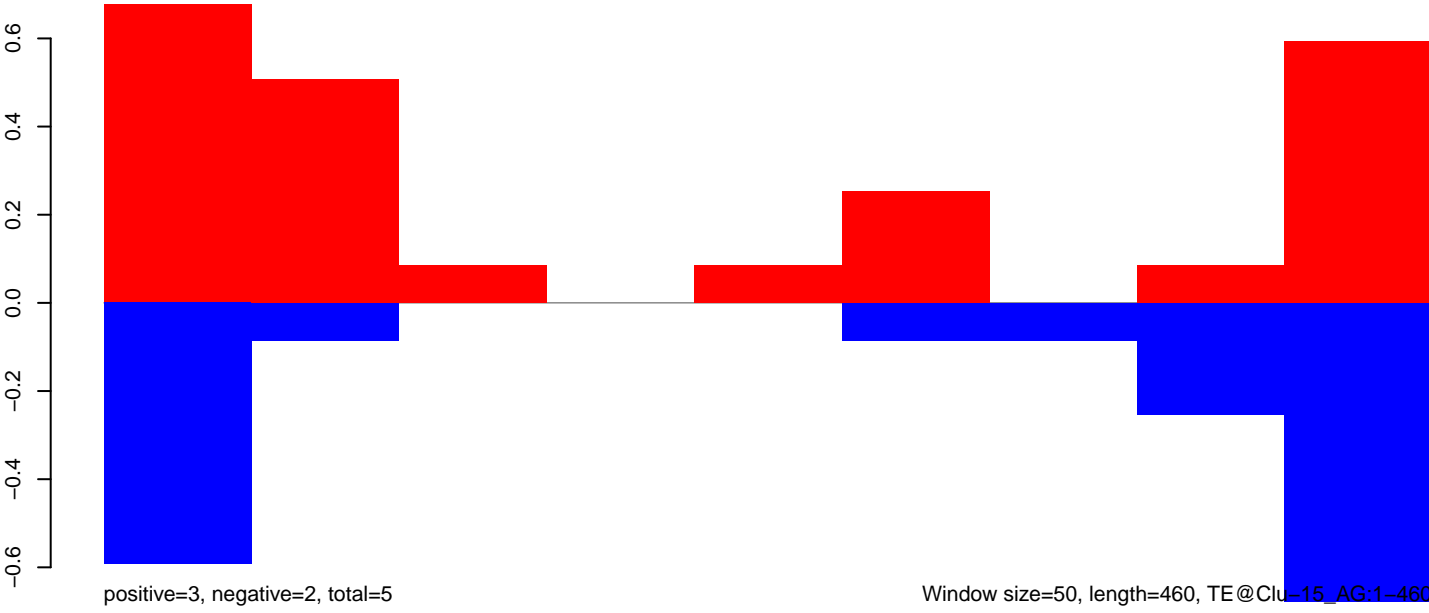
AnGam_Mos55_cells.18_23.rep



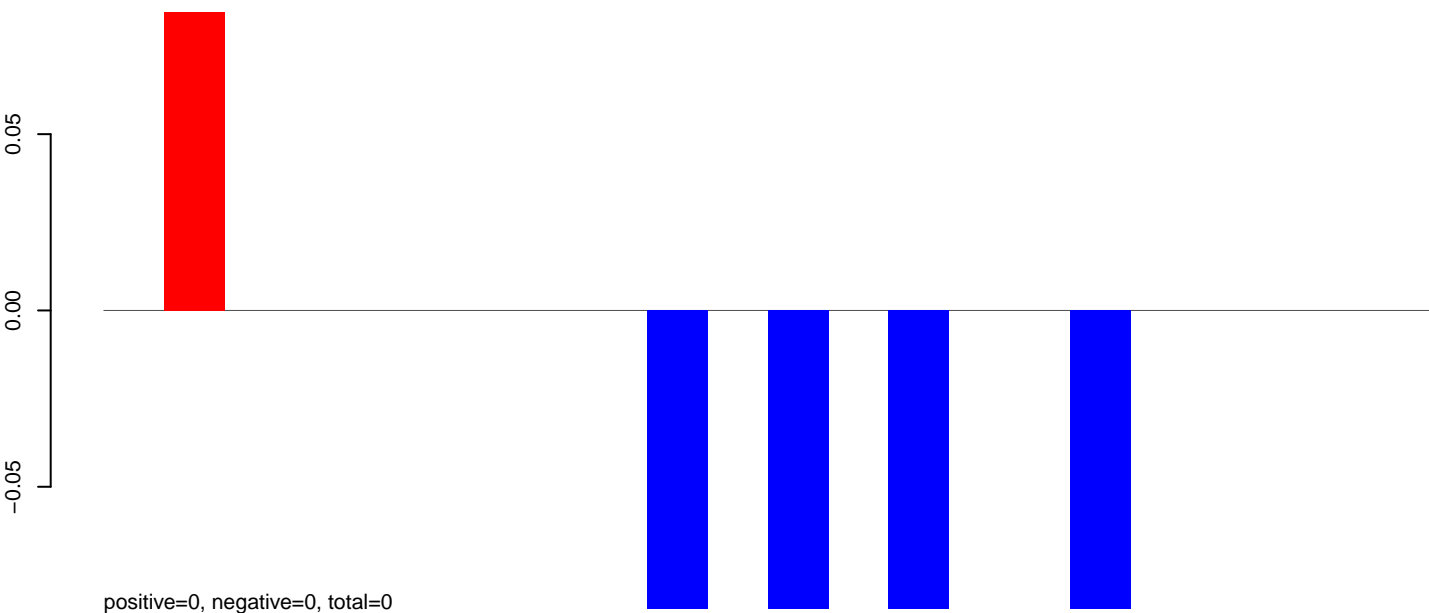
AnGam_Mos55_cells.24_35.rep



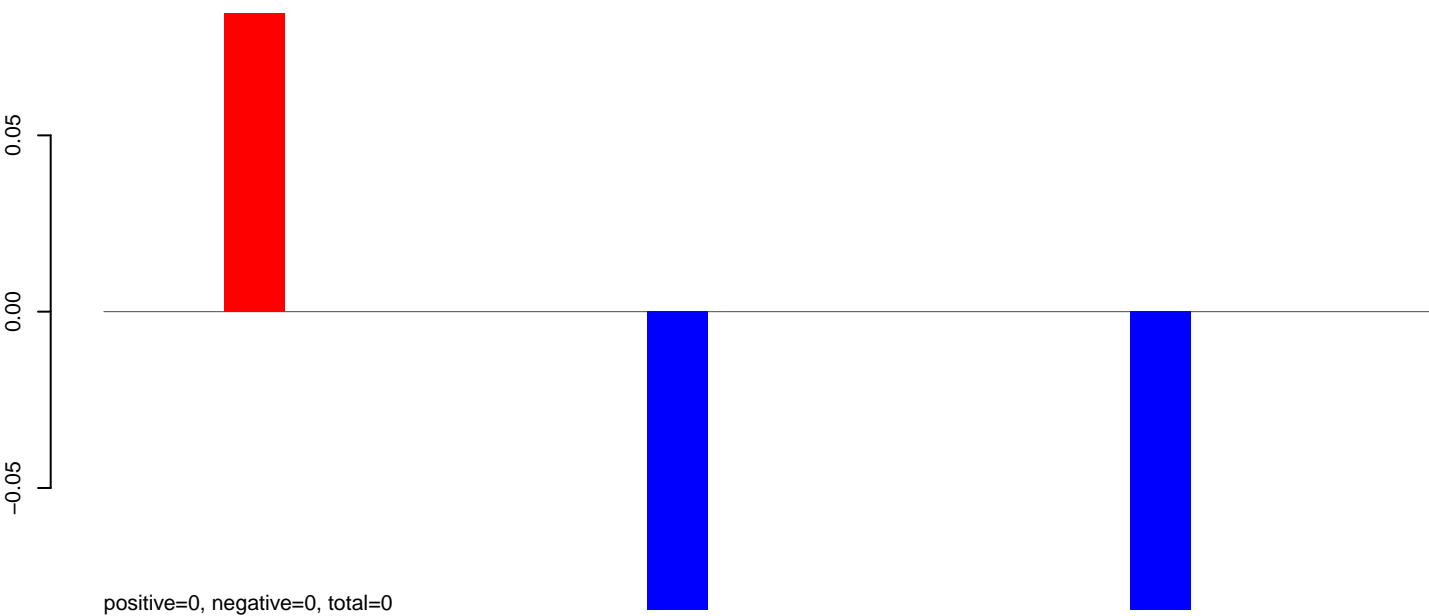
AnGam_Mos55_cells.rep



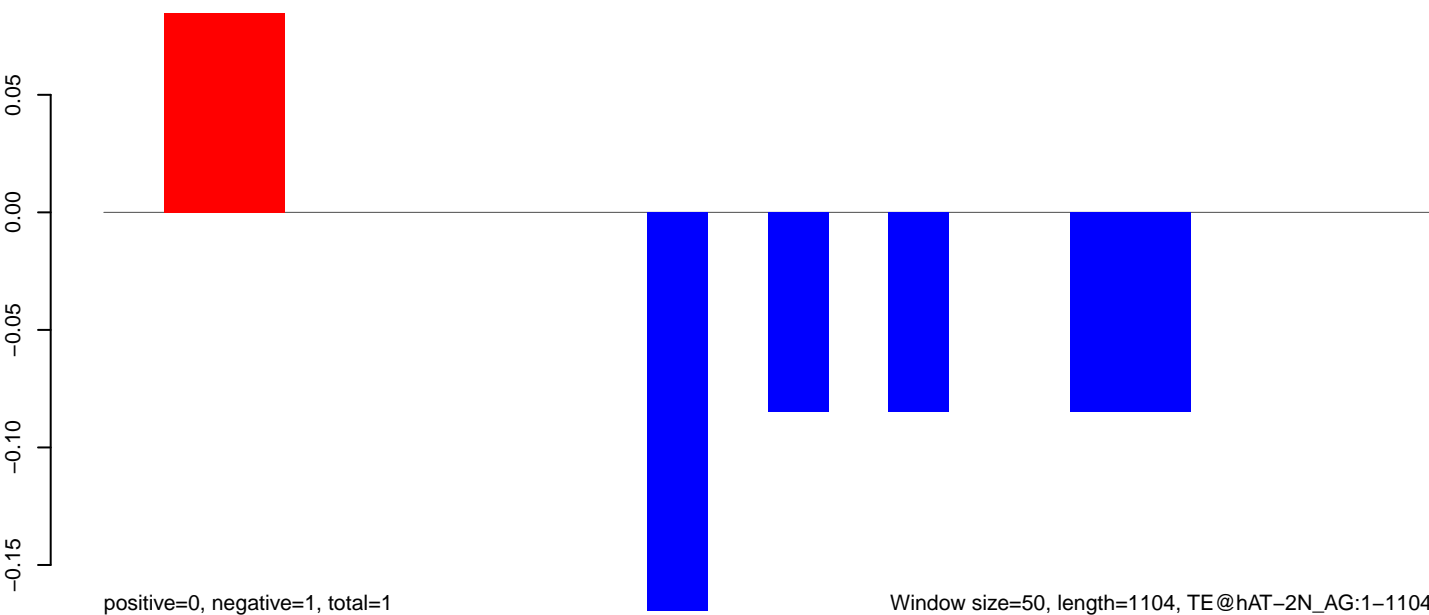
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



Window size=50, length=1104, TE@hAT-2N_AG:1-1104

AnGam_Mos55_cells.18_23.rep

0.2
0.1
0.0
-0.1
-0.2
-0.3

positive=0, negative=0, total=1

AnGam_Mos55_cells.24_35.rep

0.2
0.1
0.0
-0.1
-0.2
-0.3

positive=1, negative=1, total=2

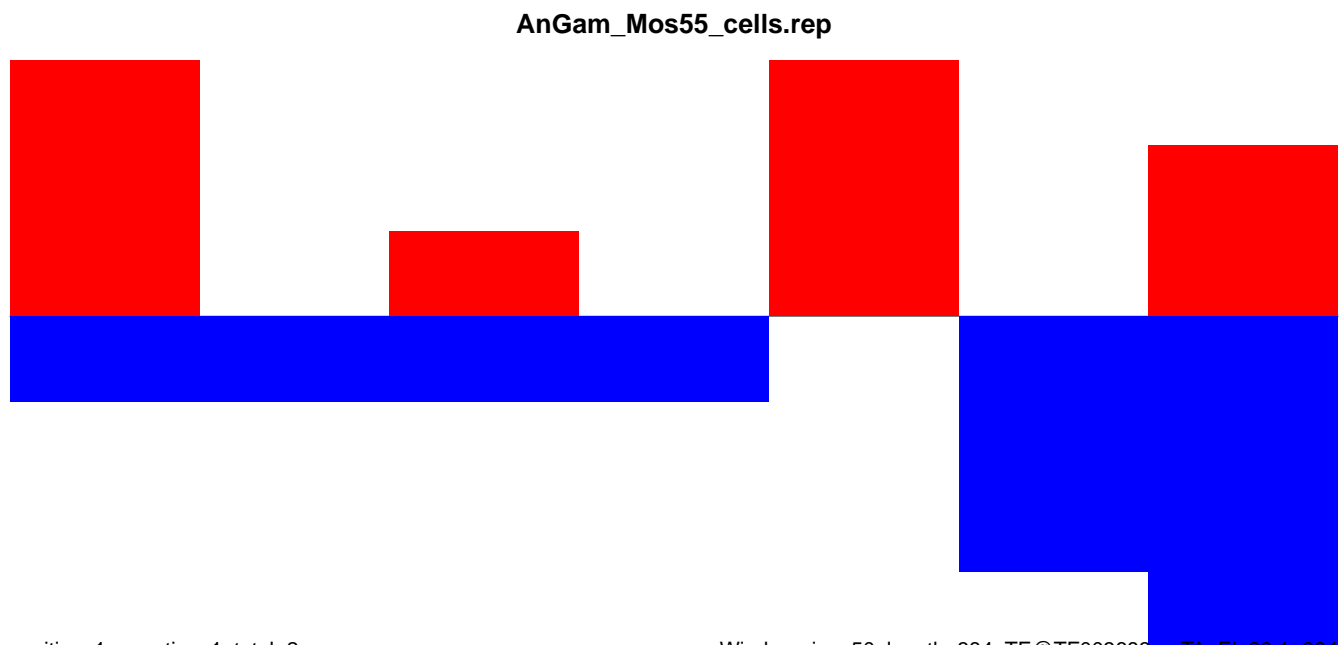
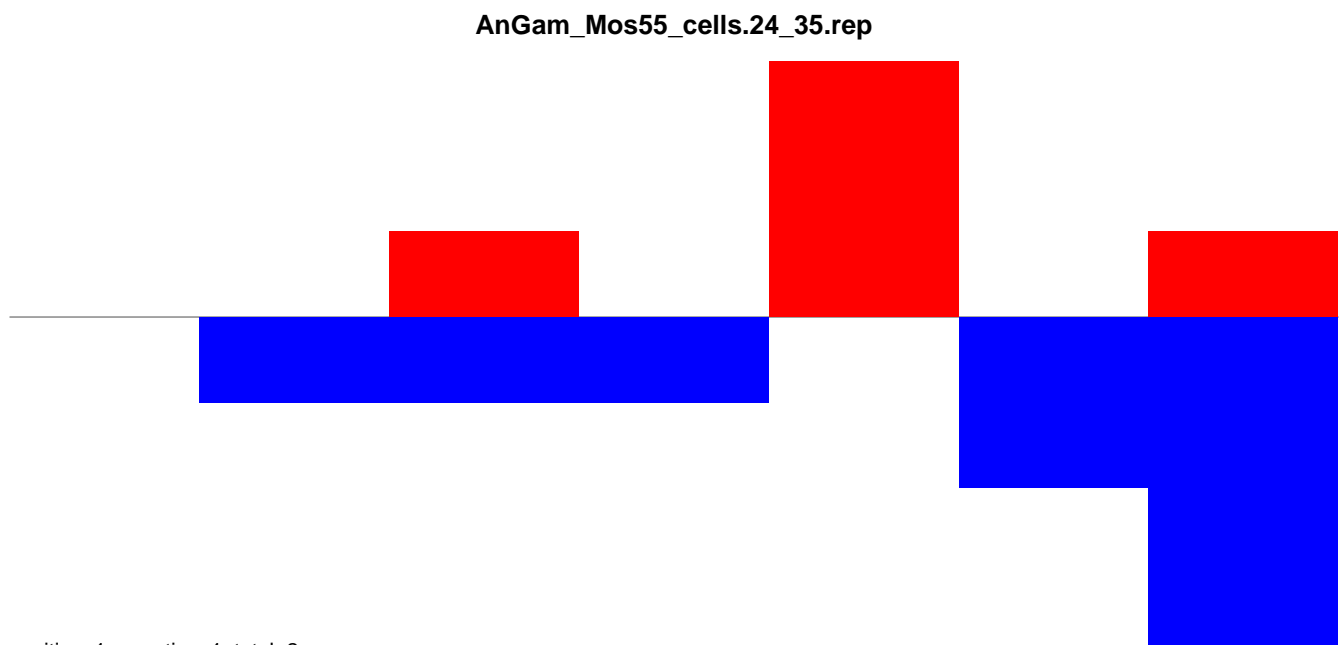
AnGam_Mos55_cells.rep

0.2
0.1
0.0
-0.1
-0.2
-0.3

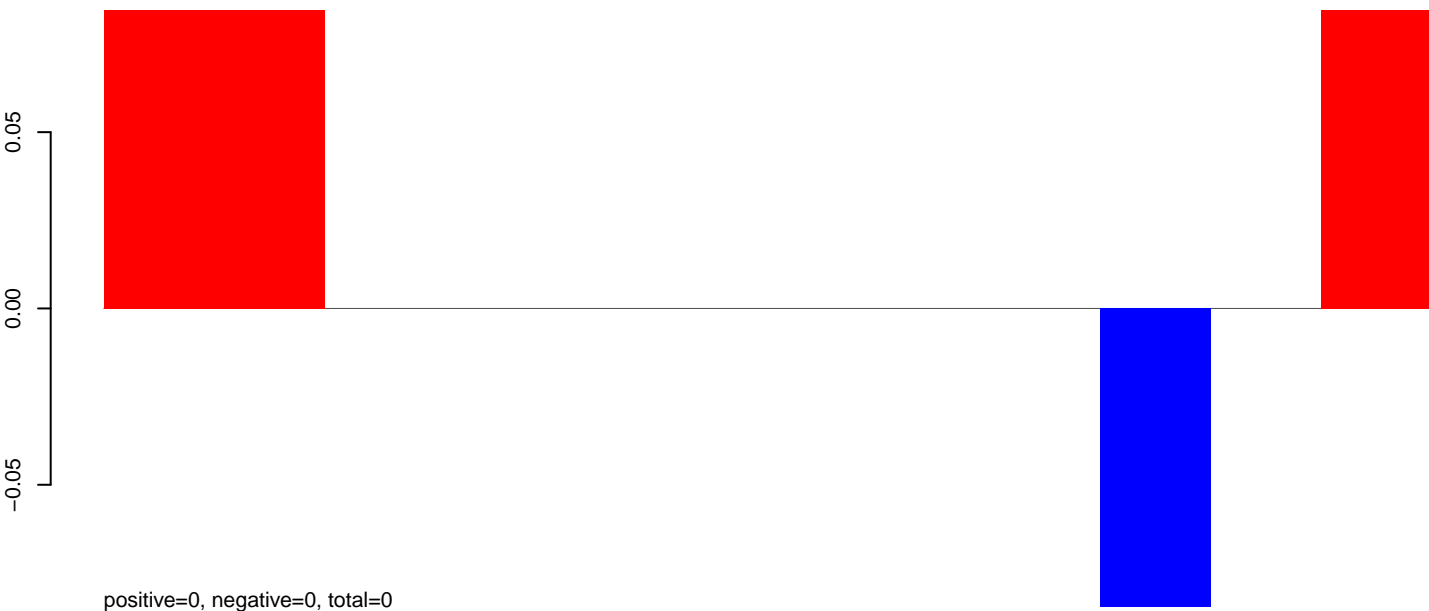
positive=1, negative=1, total=2

Window size=50, length=384, TE@TF003683-mTA_Ele20:1-384

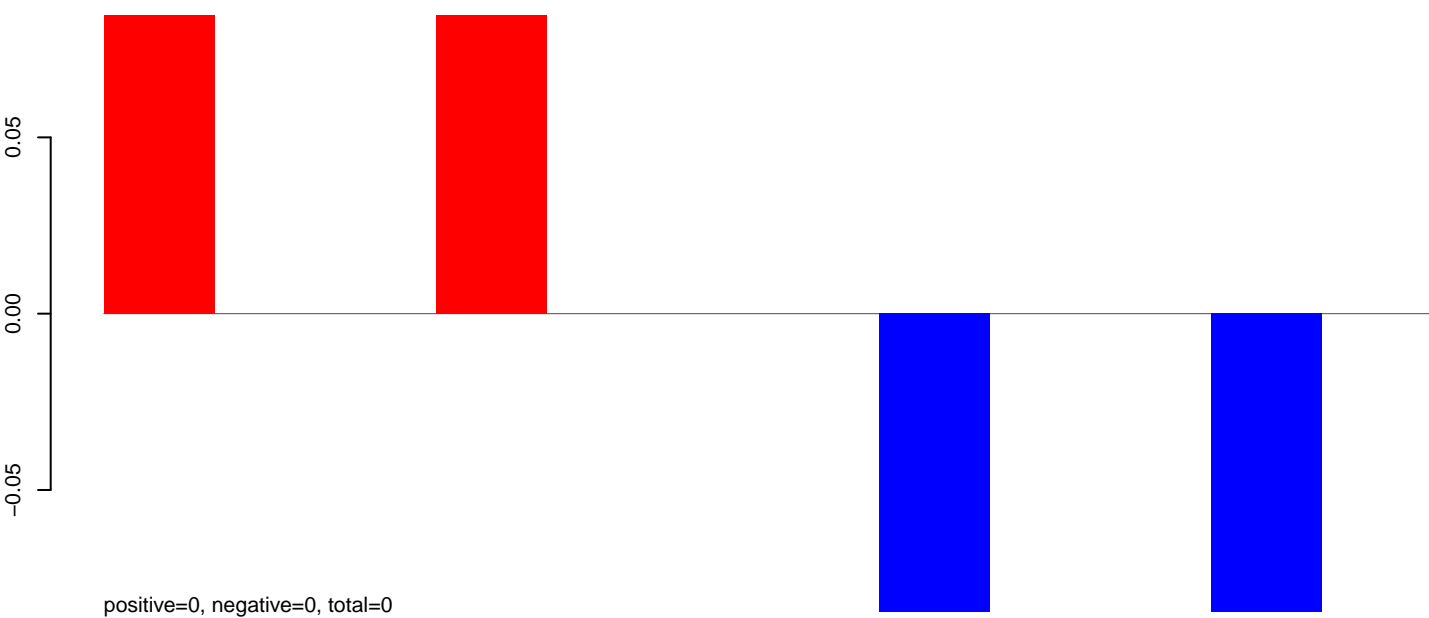
50 100 150 200 250 300 350 400



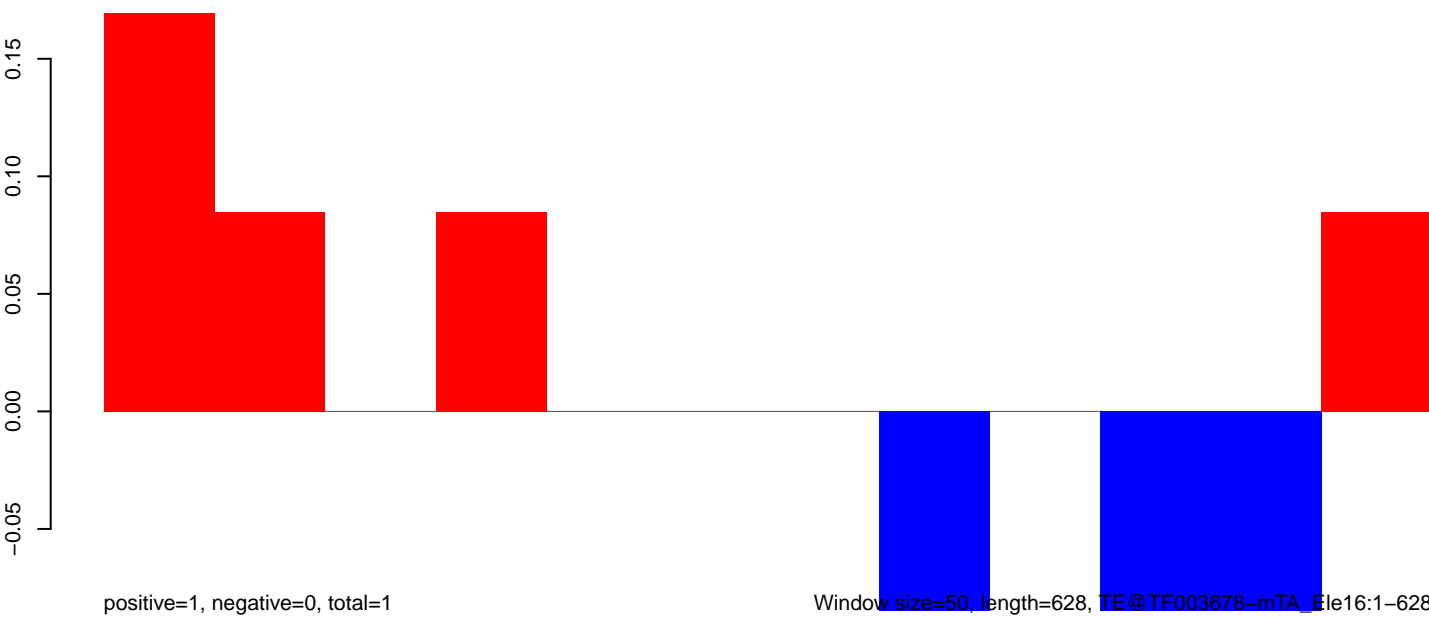
AnGam_Mos55_cells.18_23.rep



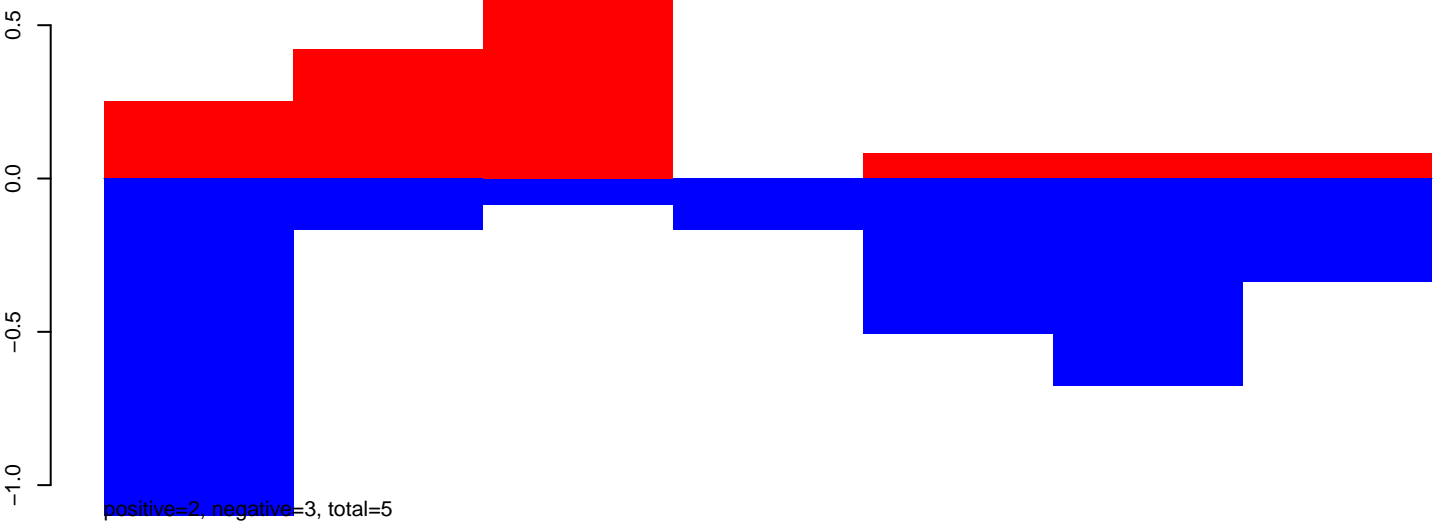
AnGam_Mos55_cells.24_35.rep



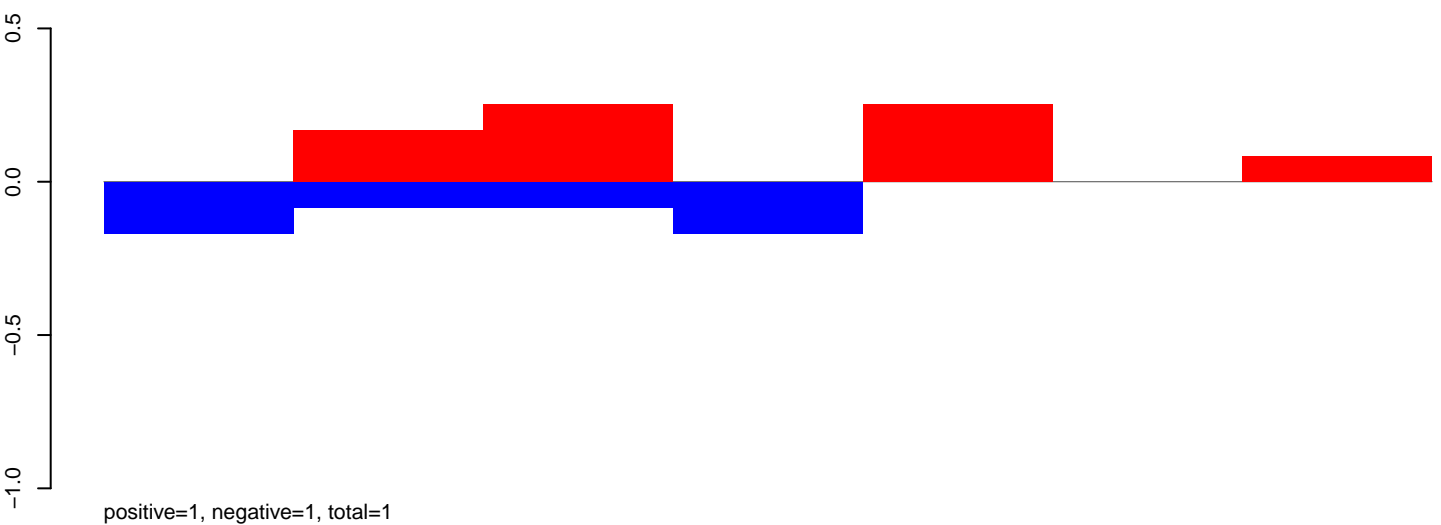
AnGam_Mos55_cells.rep



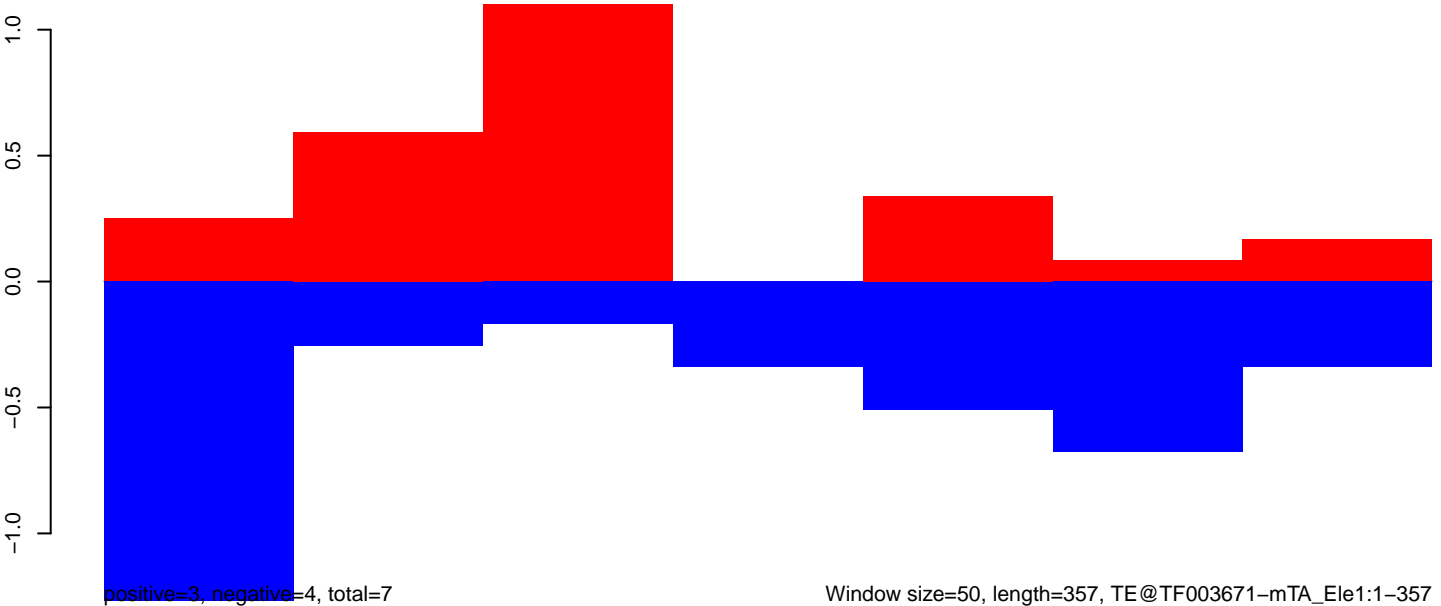
AnGam_Mos55_cells.18_23.rep



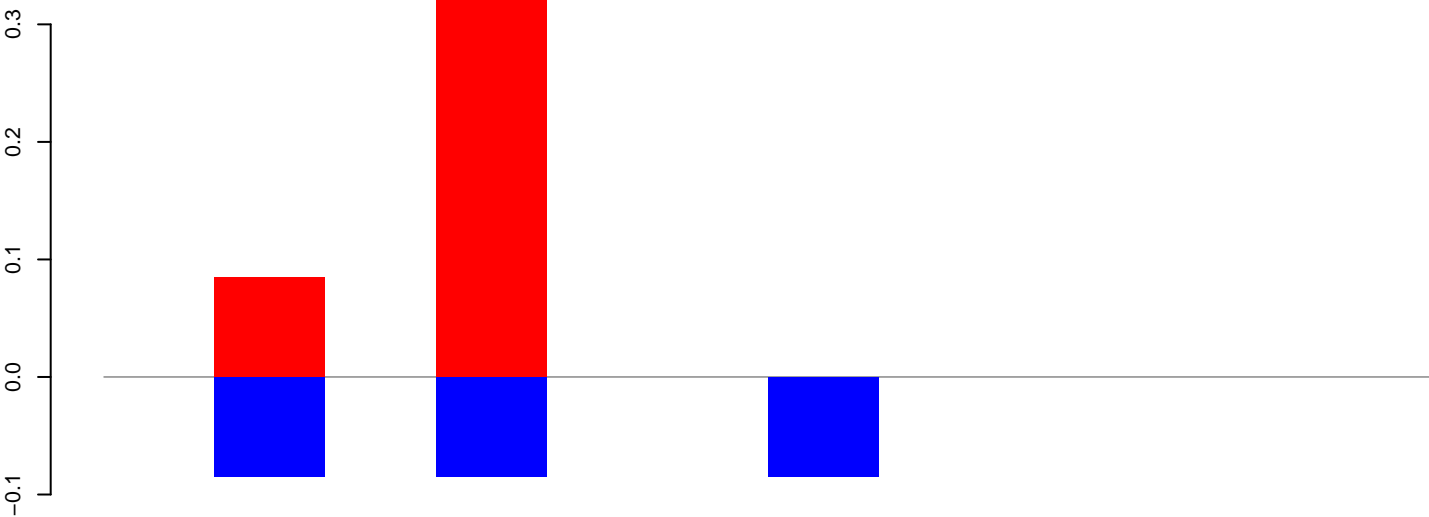
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

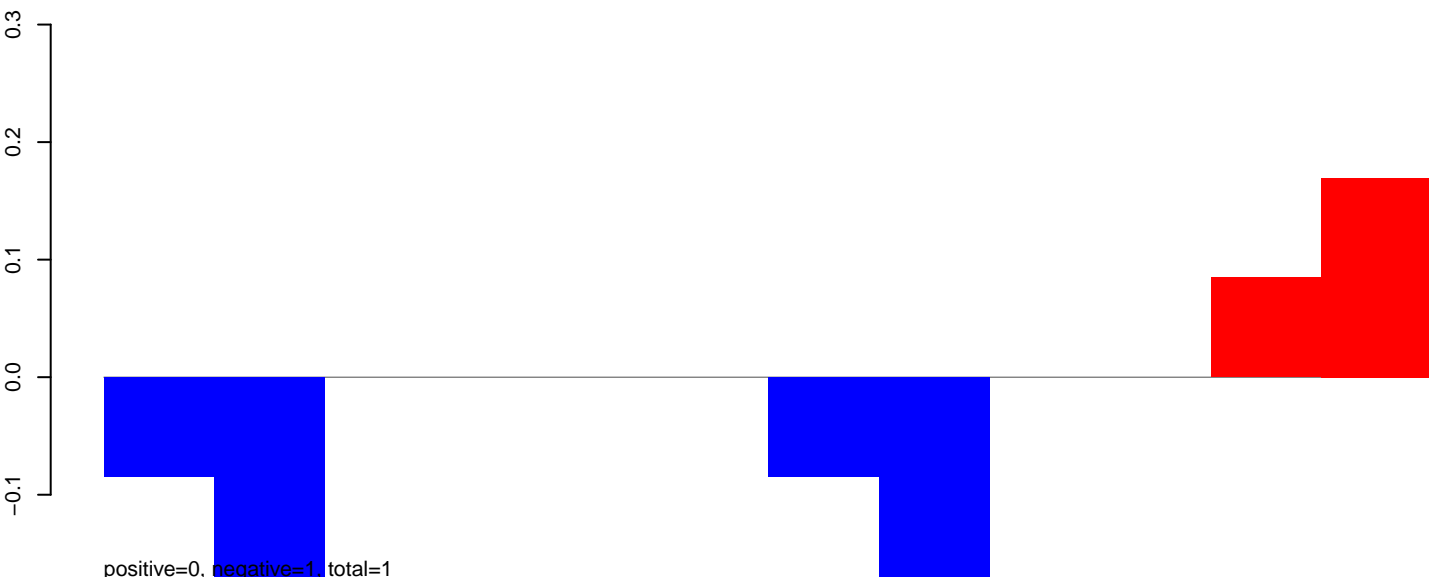


AnGam_Mos55_cells.18_23.rep



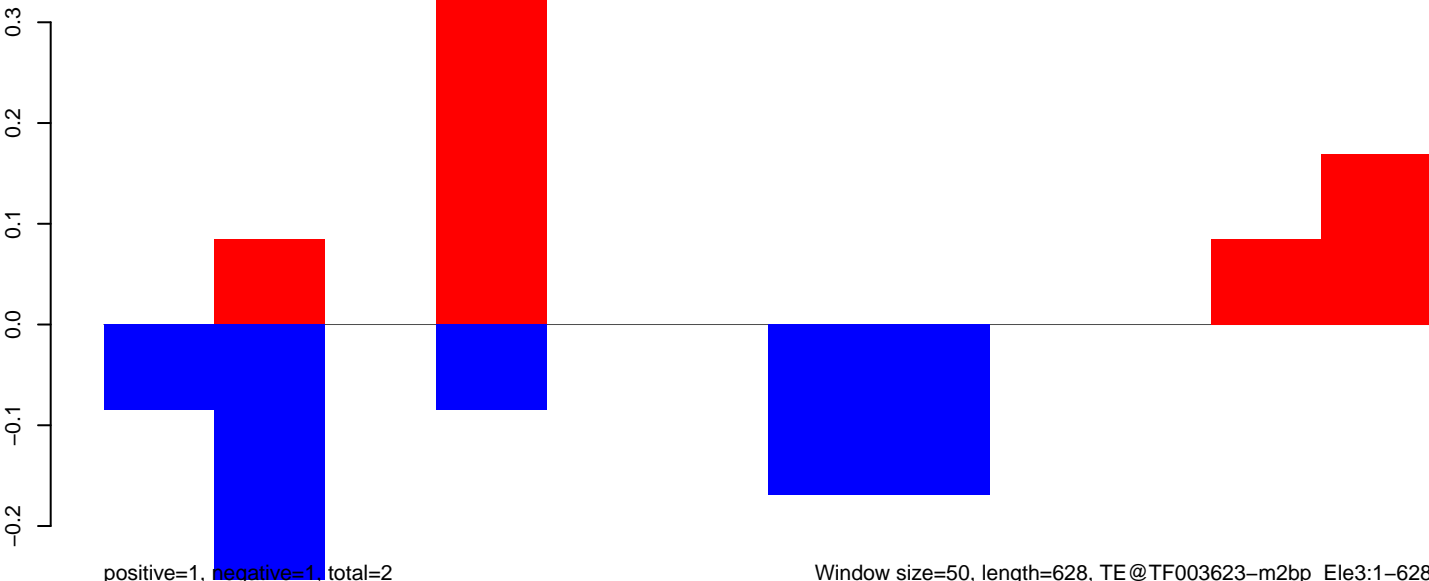
positive=0, negative=0, total=1

AnGam_Mos55_cells.24_35.rep



positive=0, negative=1, total=1

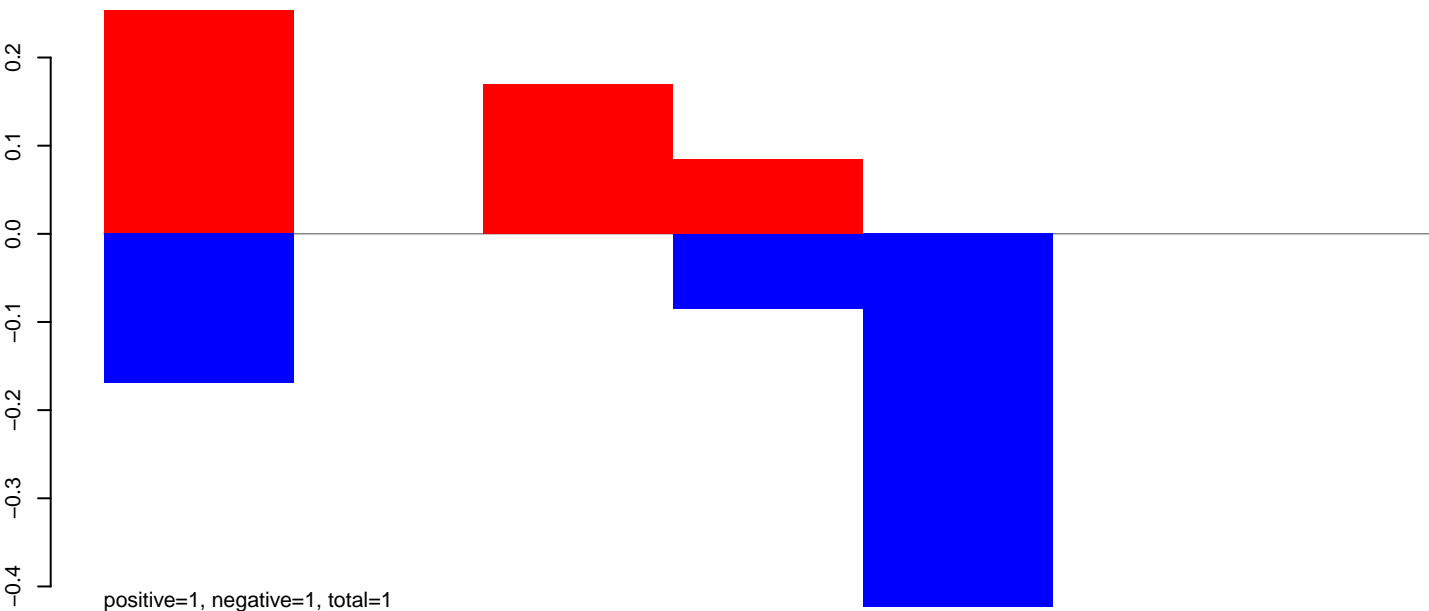
AnGam_Mos55_cells.rep



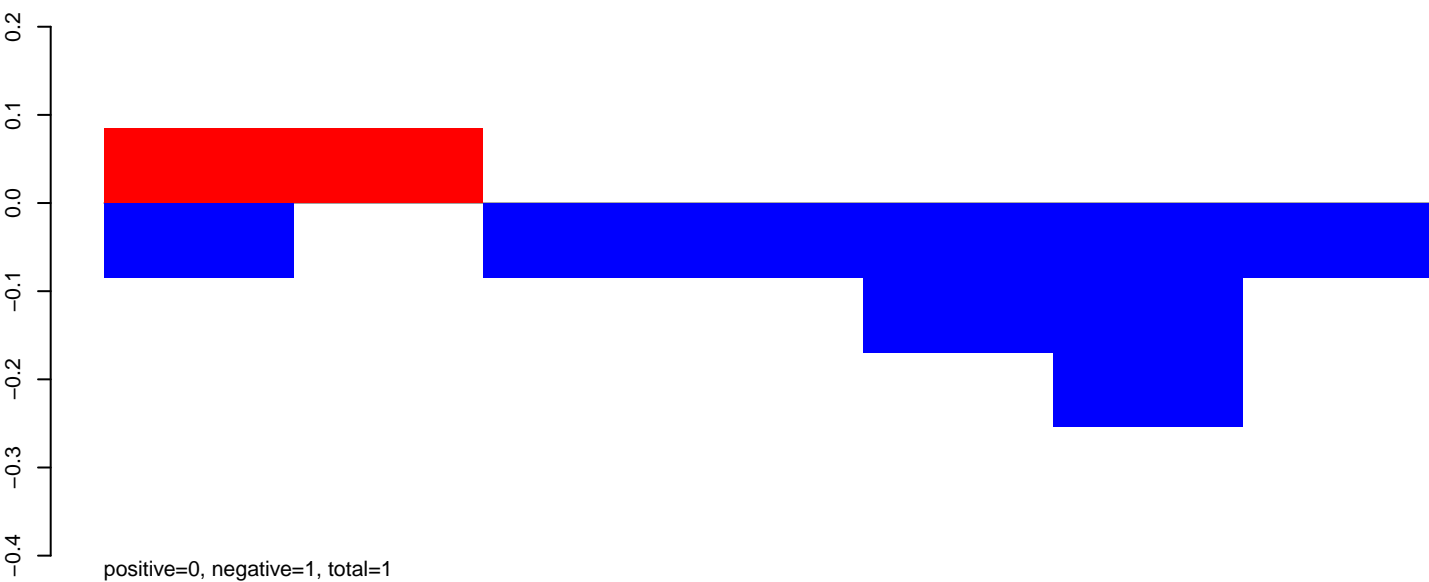
positive=1, negative=1, total=2

Window size=50, length=628, TE@TF003623-m2bp_Ele3:1-628

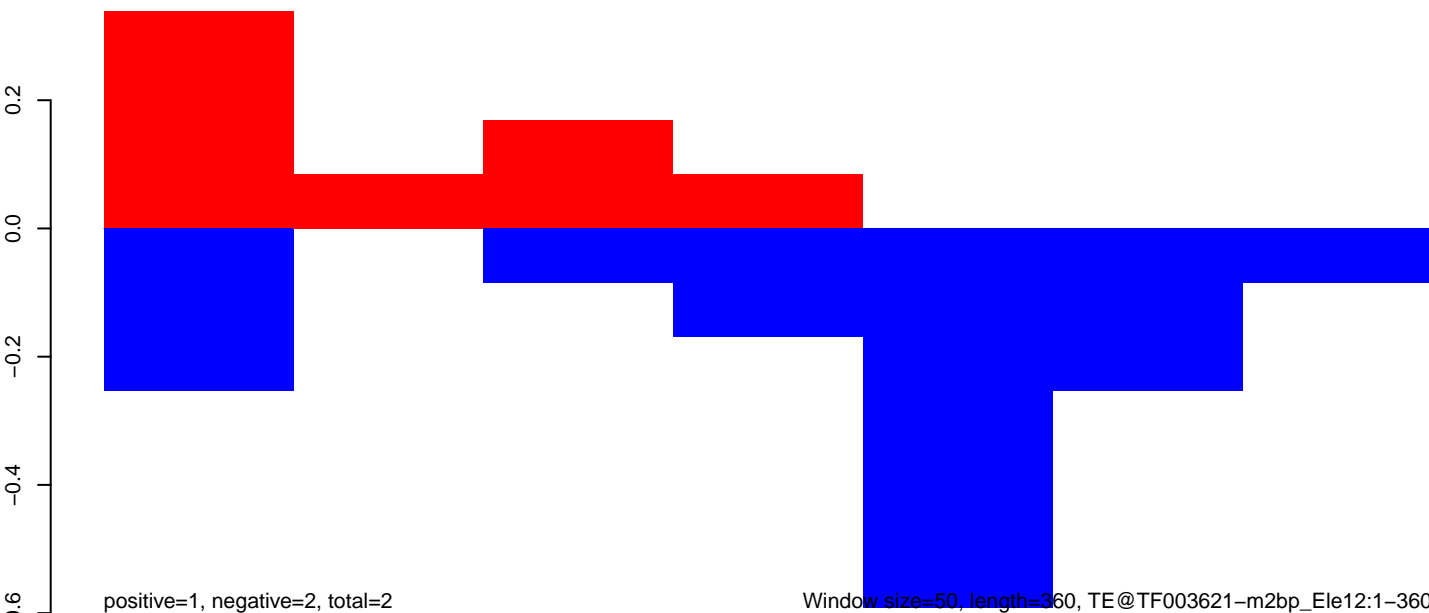
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

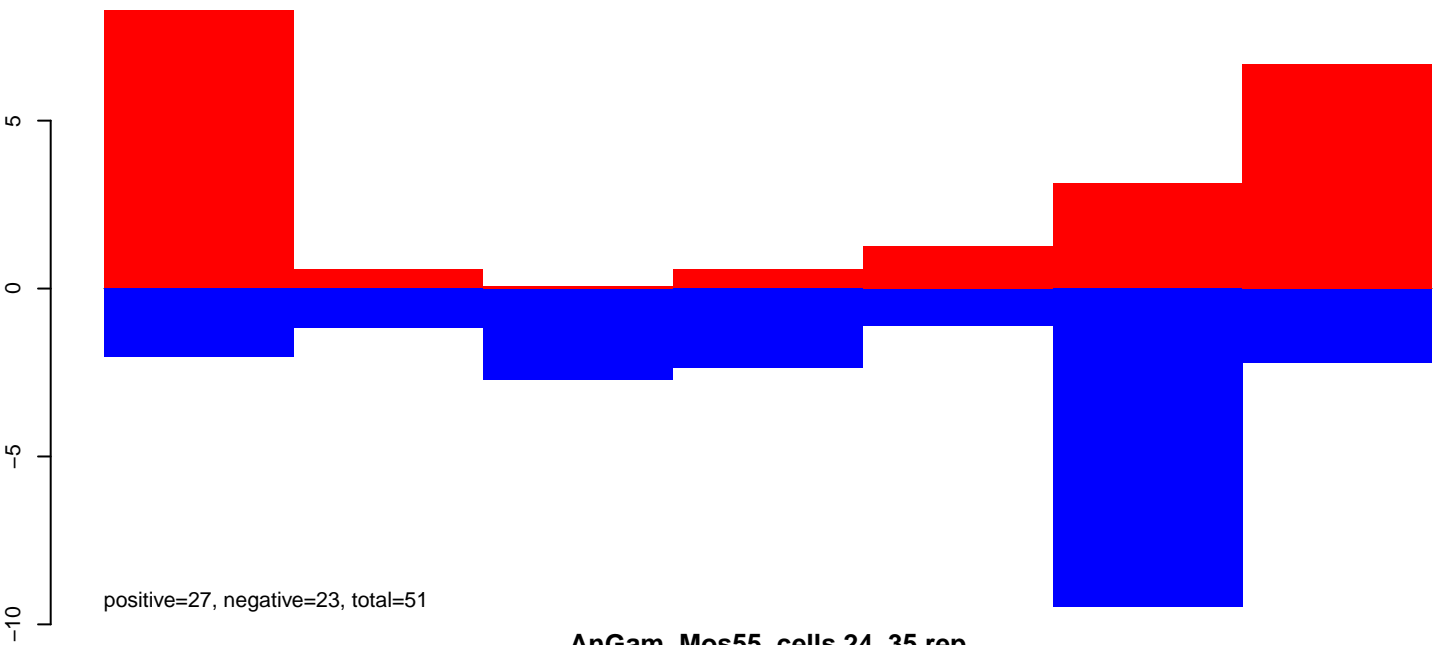


AnGam_Mos55_cells.rep

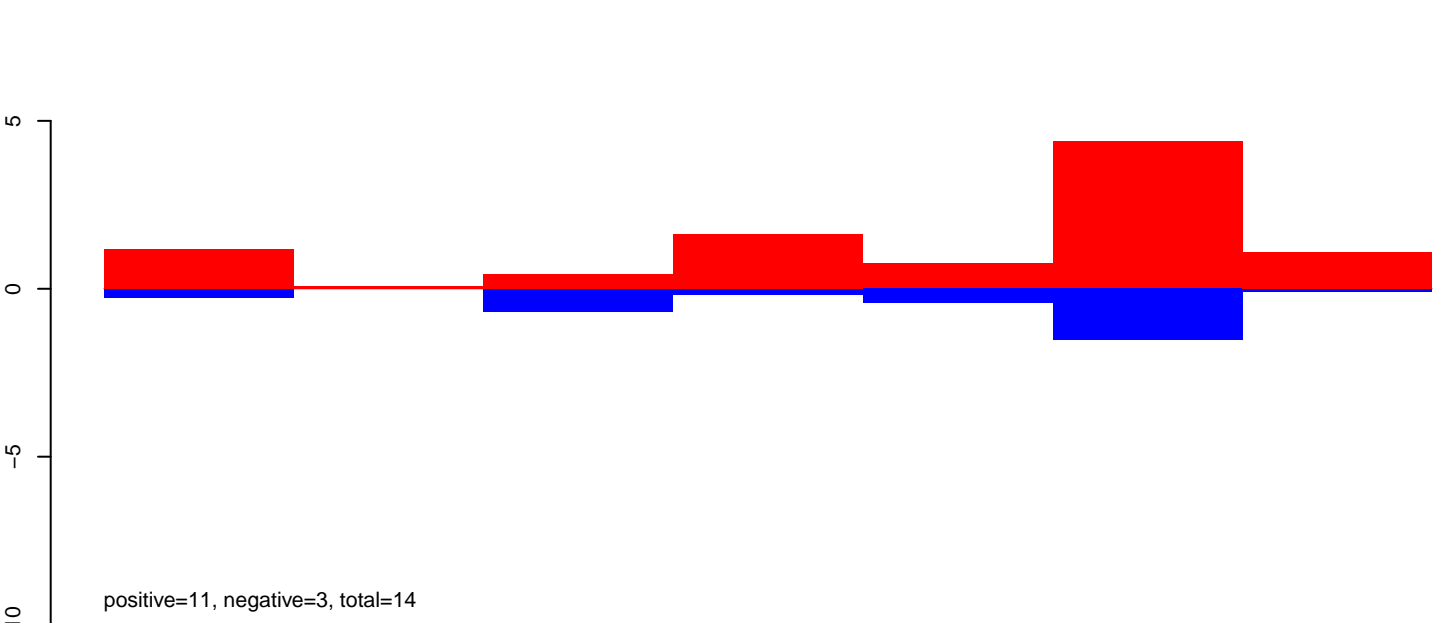


Window size=50, length=360, TE@TF003621-m2bp_Ele12:1-360

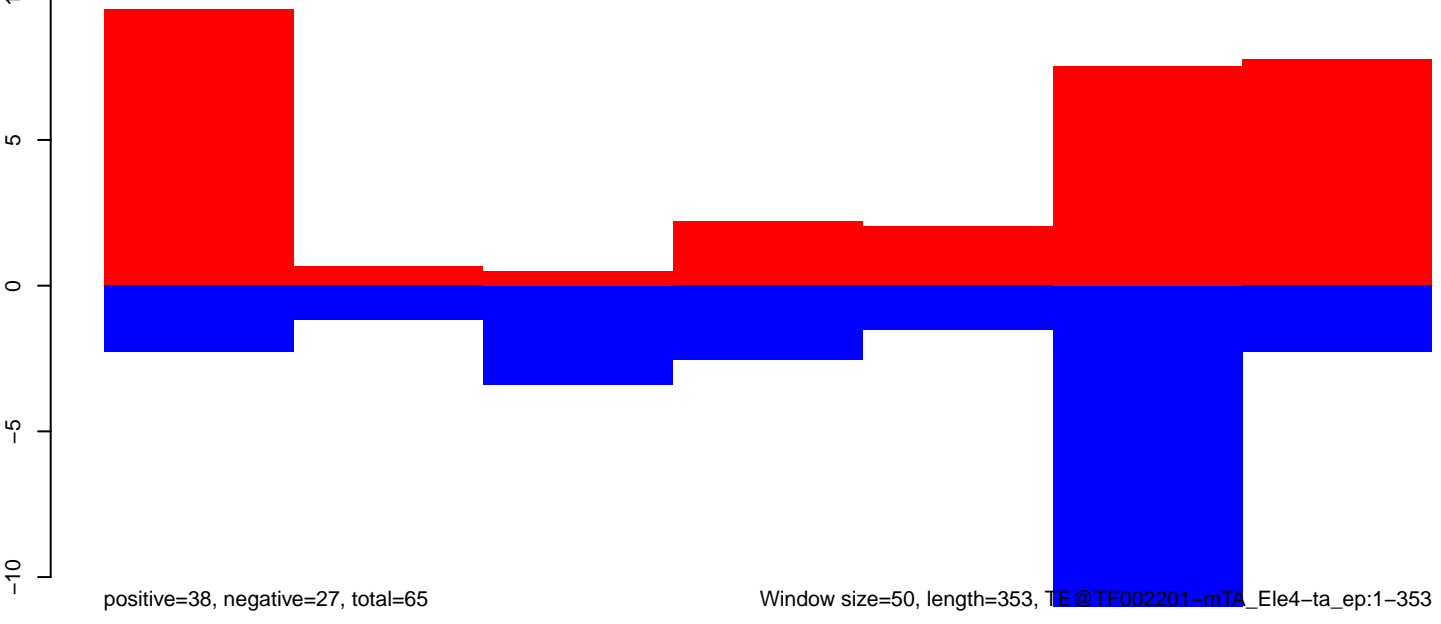
AnGam_Mos55_cells.18_23.rep



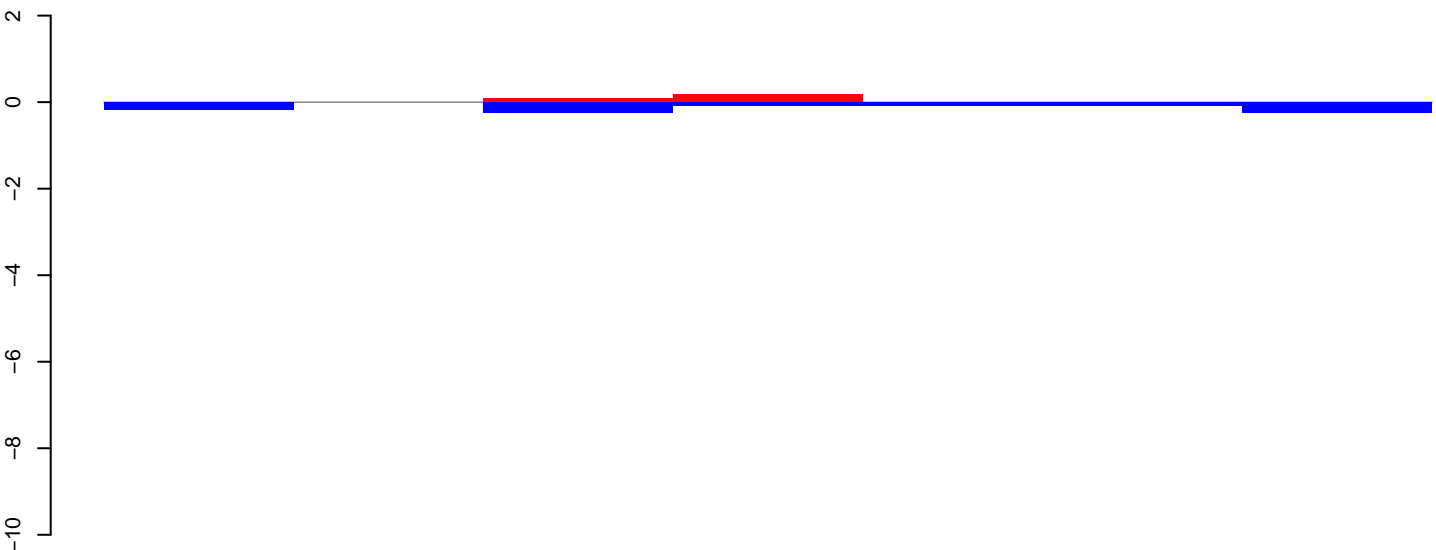
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

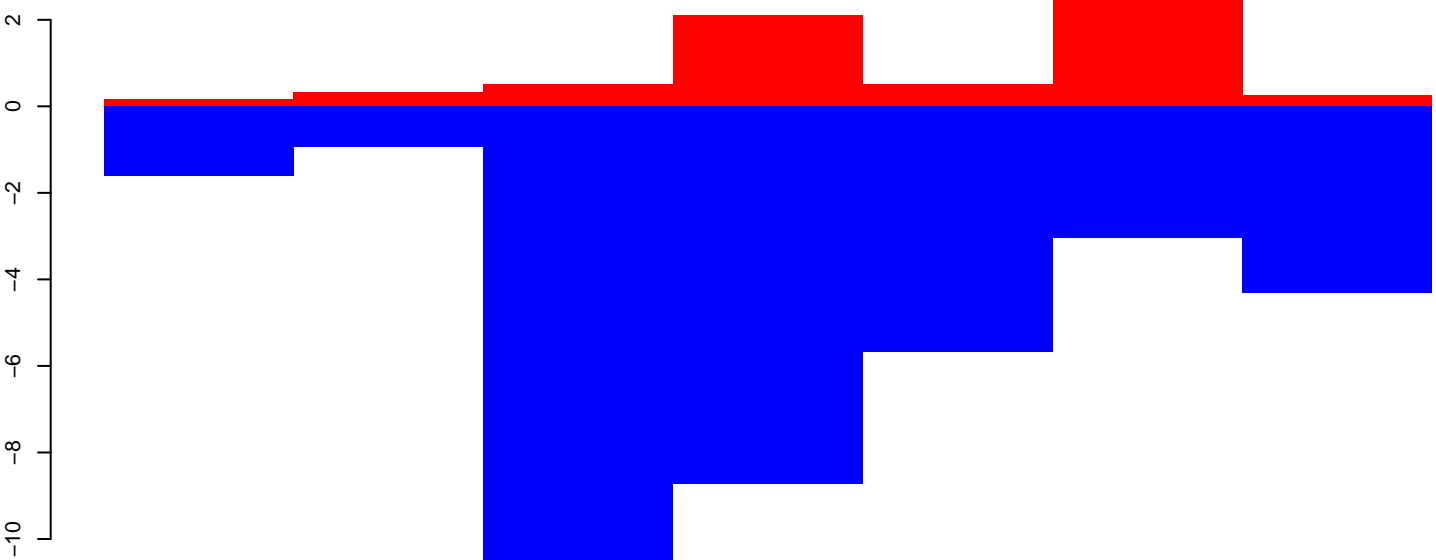


AnGam_Mos55_cells.18_23.rep



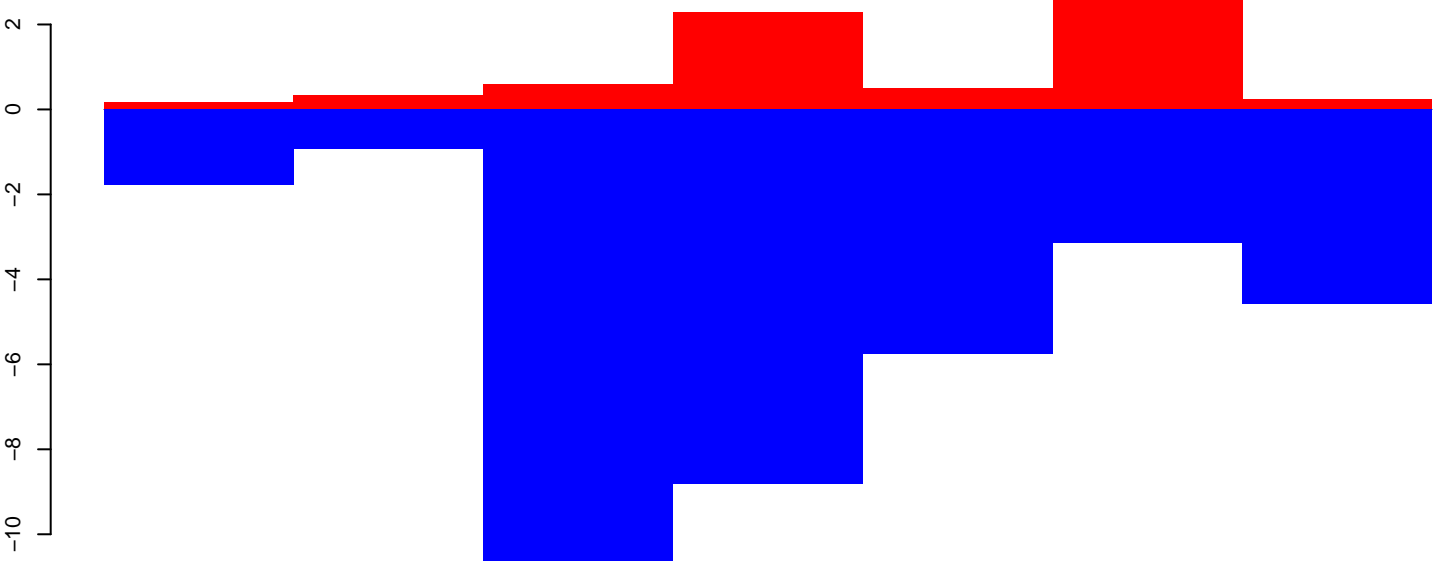
positive=0, negative=1, total=1

AnGam_Mos55_cells.24_35.rep



positive=7, negative=40, total=47

AnGam_Mos55_cells.rep

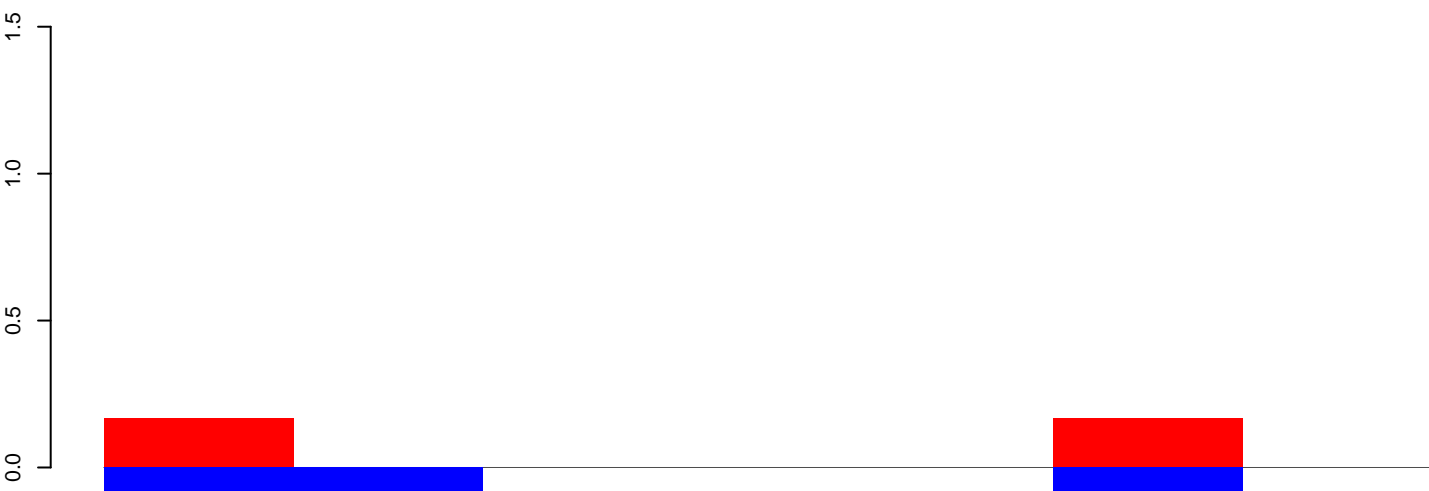


positive=7, negative=41, total=48

Window size=50, length=376, TE@TF002200-mTA_Ele3-ta_delta:1-376

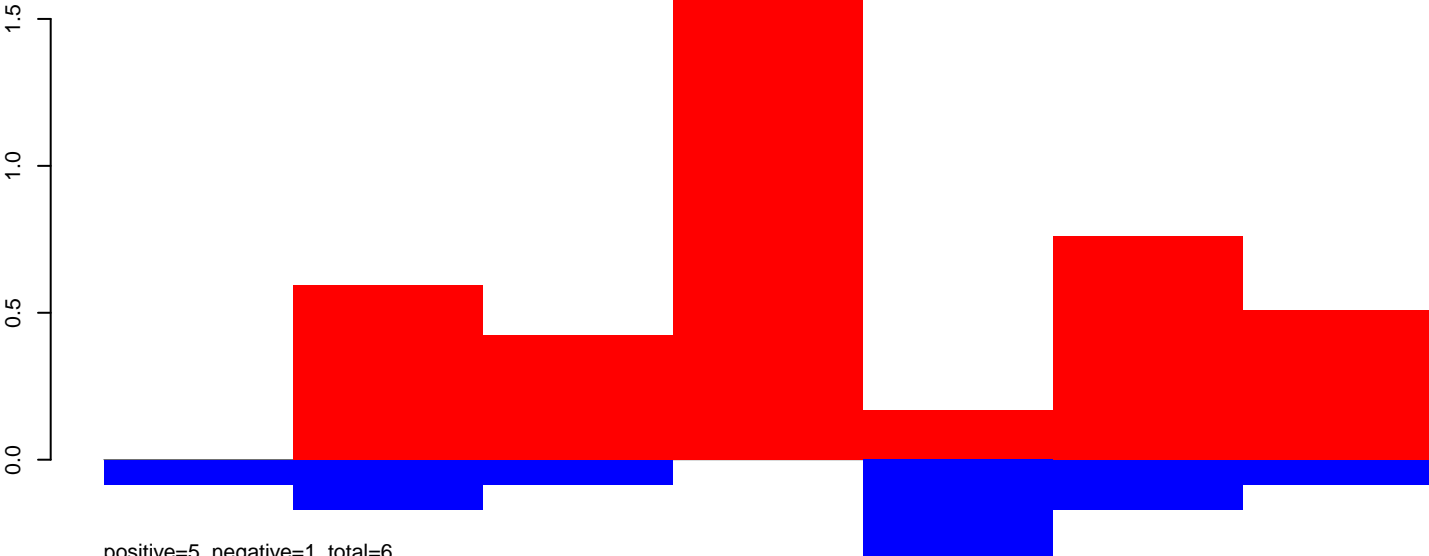
50 100 150 200 250 300 350 400

AnGam_Mos55_cells.18_23.rep



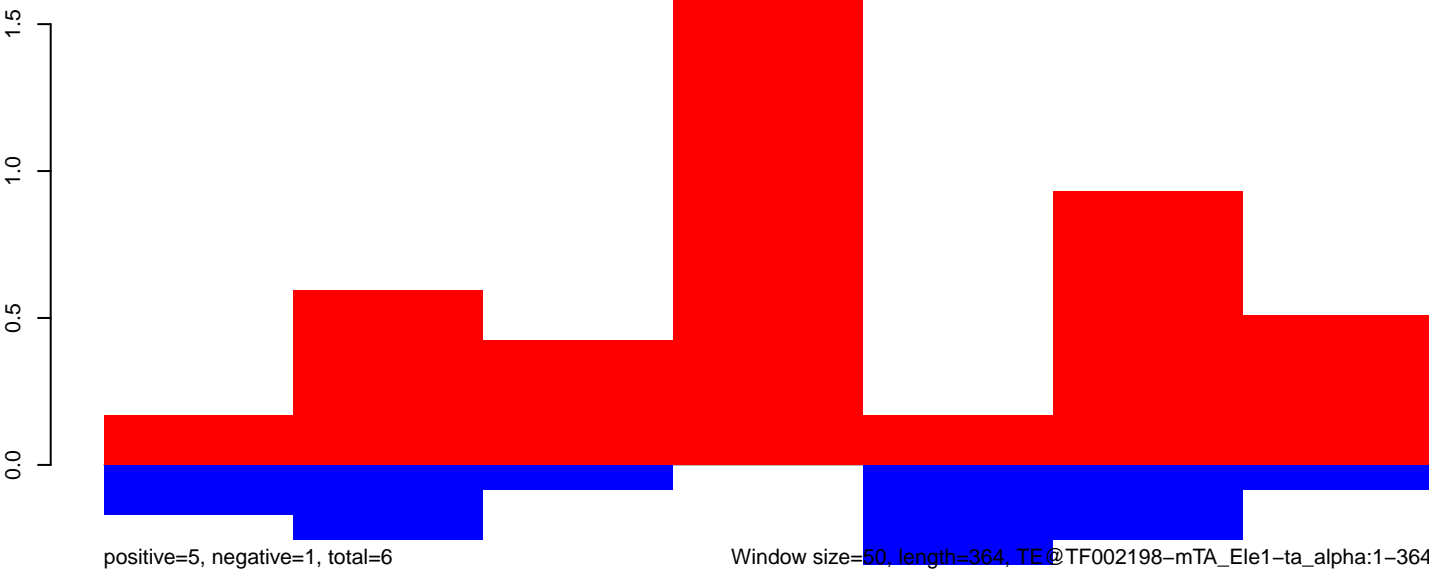
positive=0, negative=0, total=1

AnGam_Mos55_cells.24_35.rep



positive=5, negative=1, total=6

AnGam_Mos55_cells.rep

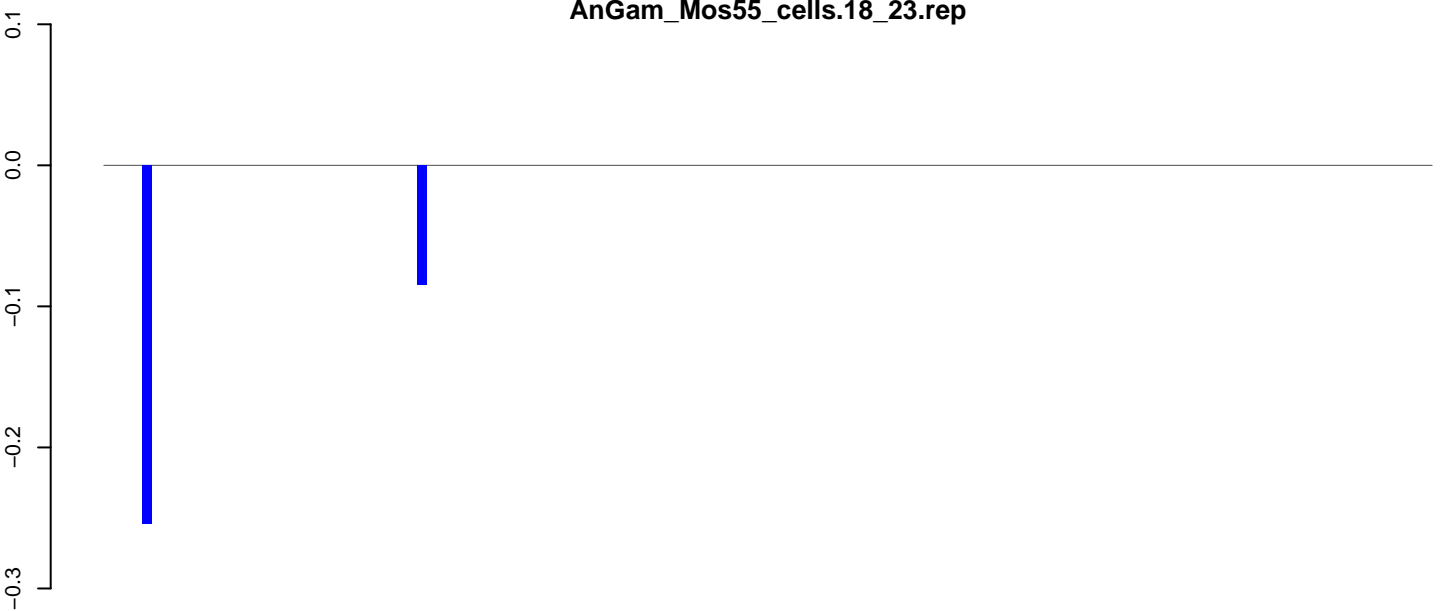


positive=5, negative=1, total=6

Window size=50, length=364, TE@TF002198-mTA_Ele1-ta_alpha:1-364

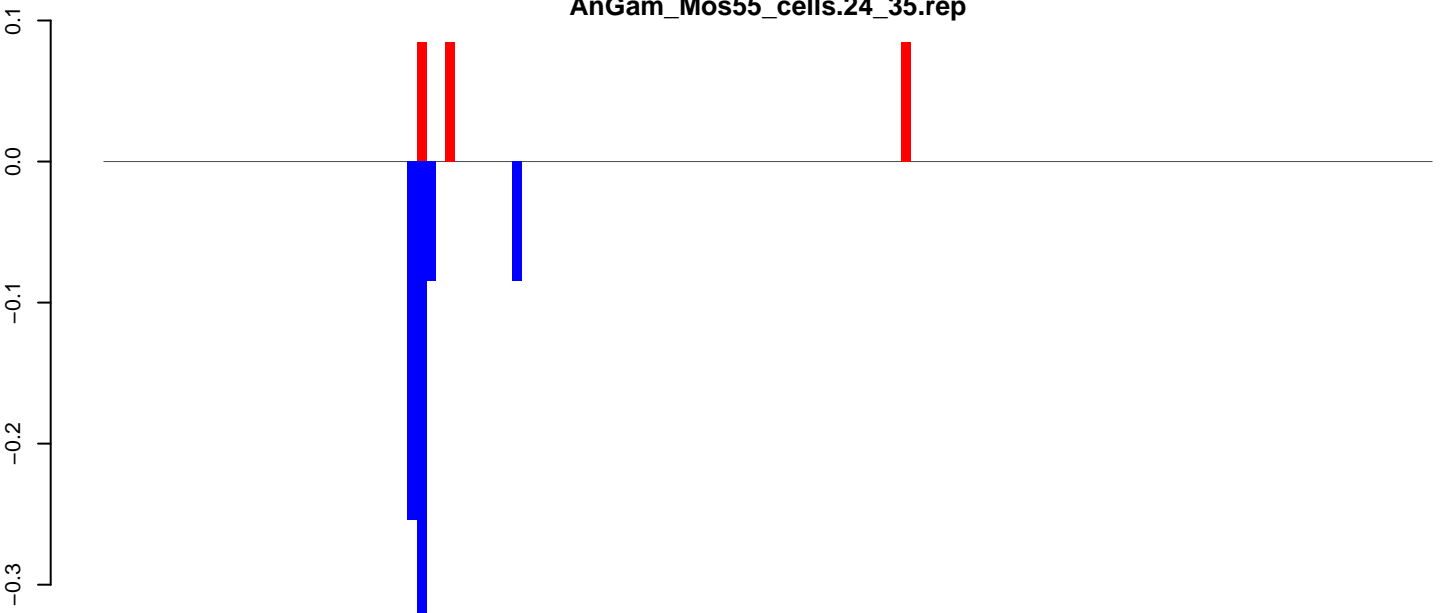
50 100 150 200 250 300 350 400

AnGam_Mos55_cells.18_23.rep



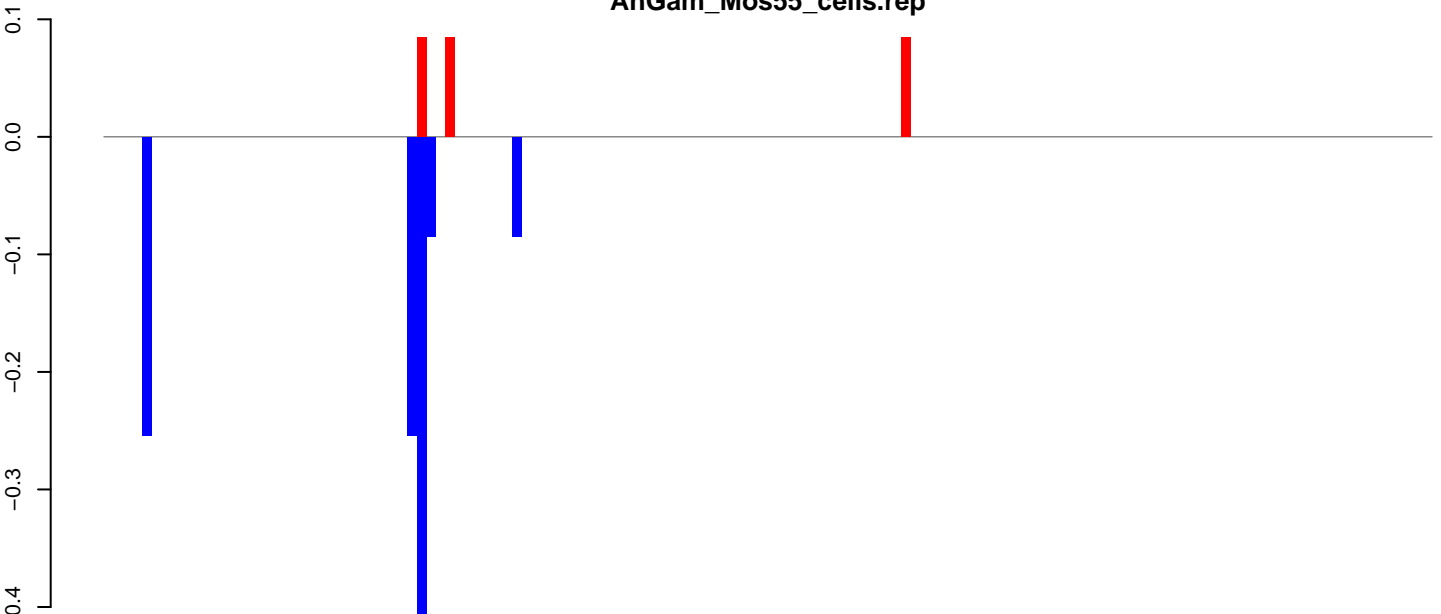
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=1, total=1

AnGam_Mos55_cells.rep

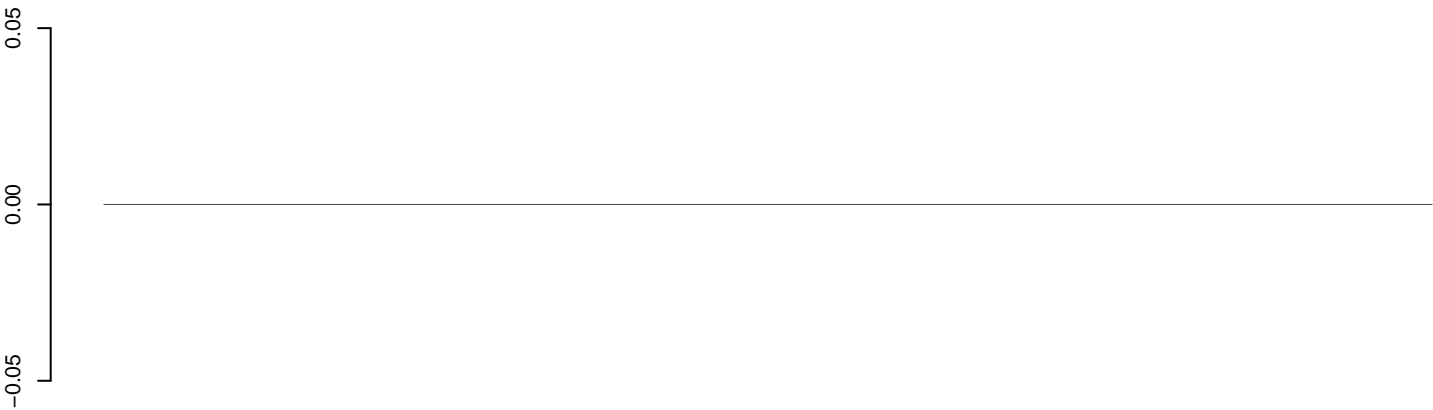


positive=0, negative=1, total=1

Window size=50, length=7048, TE@TF001977-Pao_Bel_Ele46:1-7048

0 1000 2000 3000 4000 5000 6000 7000

AnGam_Mos55_cells.18_23.rep



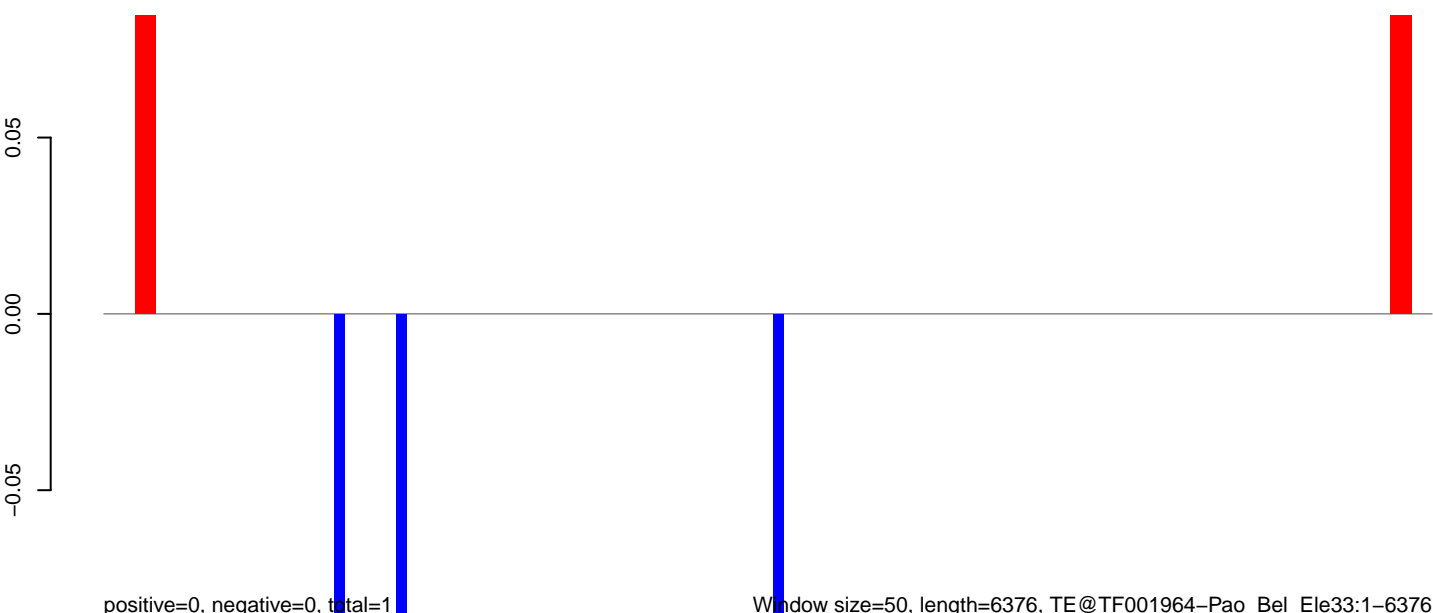
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=1

AnGam_Mos55_cells.rep

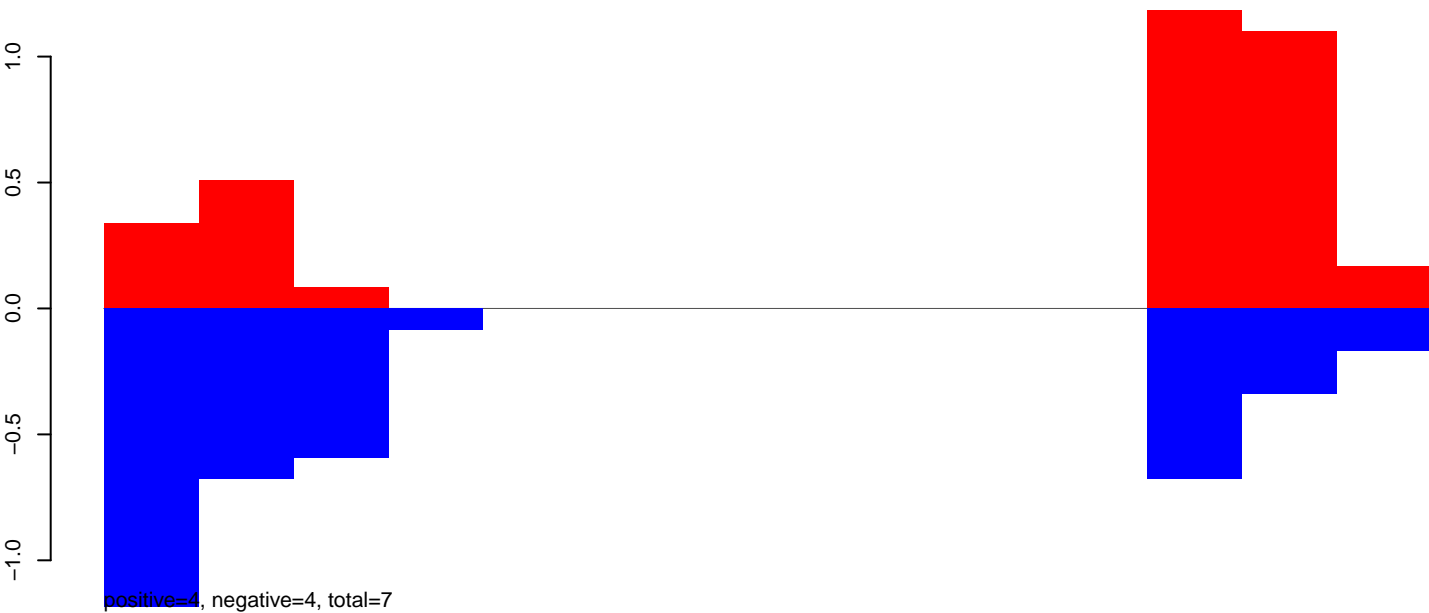


positive=0, negative=0, total=1

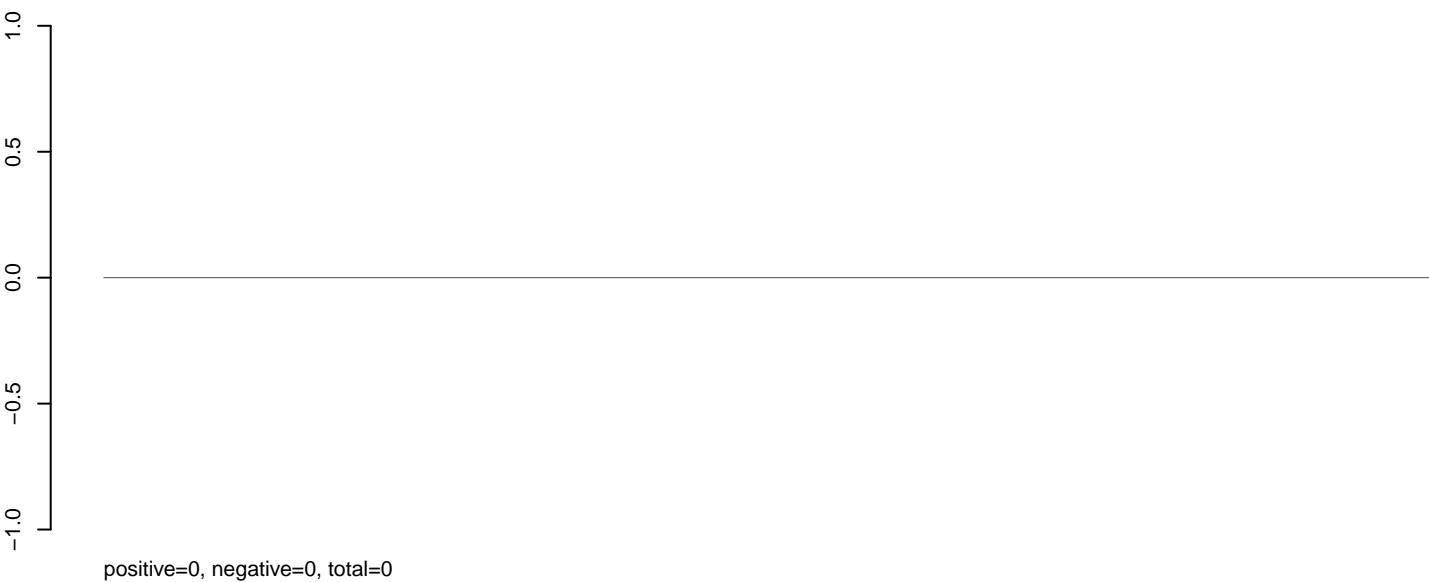
Window size=50, length=6376, TE@TF001964-Pao_Bel_Ele33:1-6376

0 1000 2000 3000 4000 5000 6000

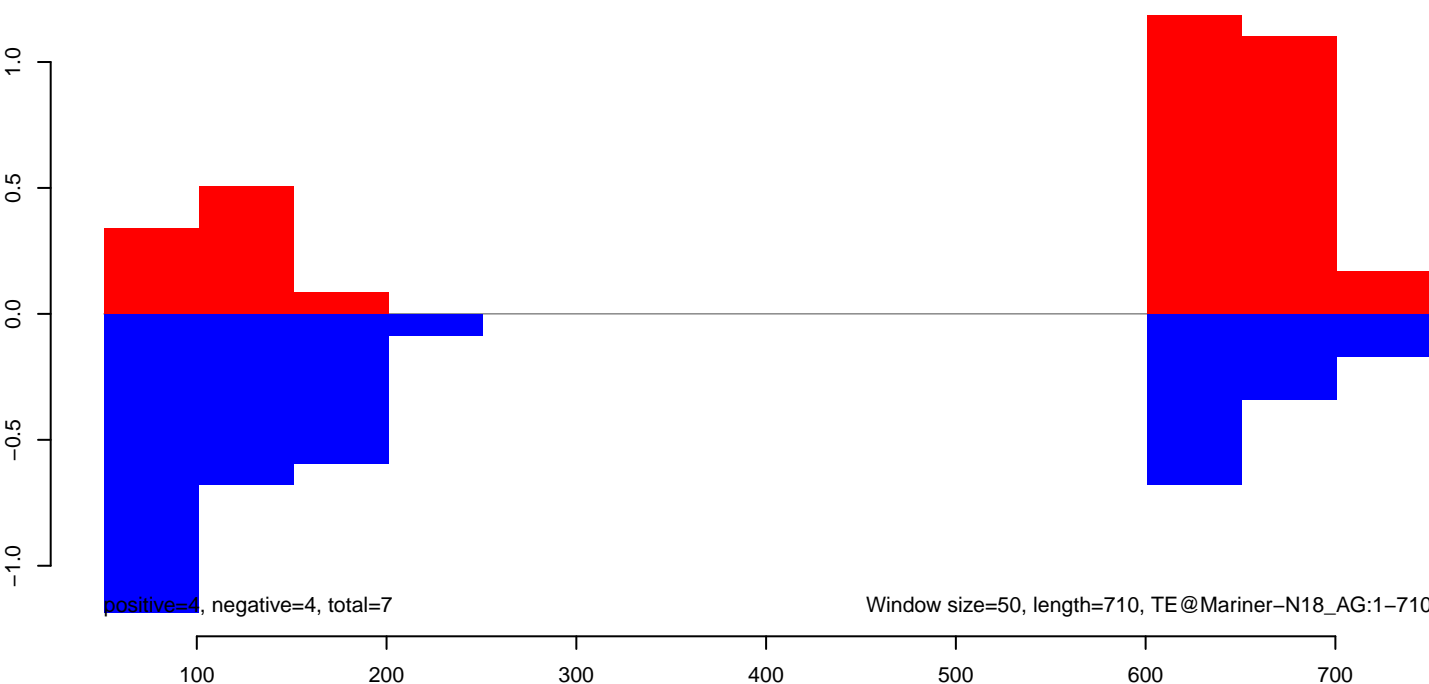
AnGam_Mos55_cells.18_23.rep



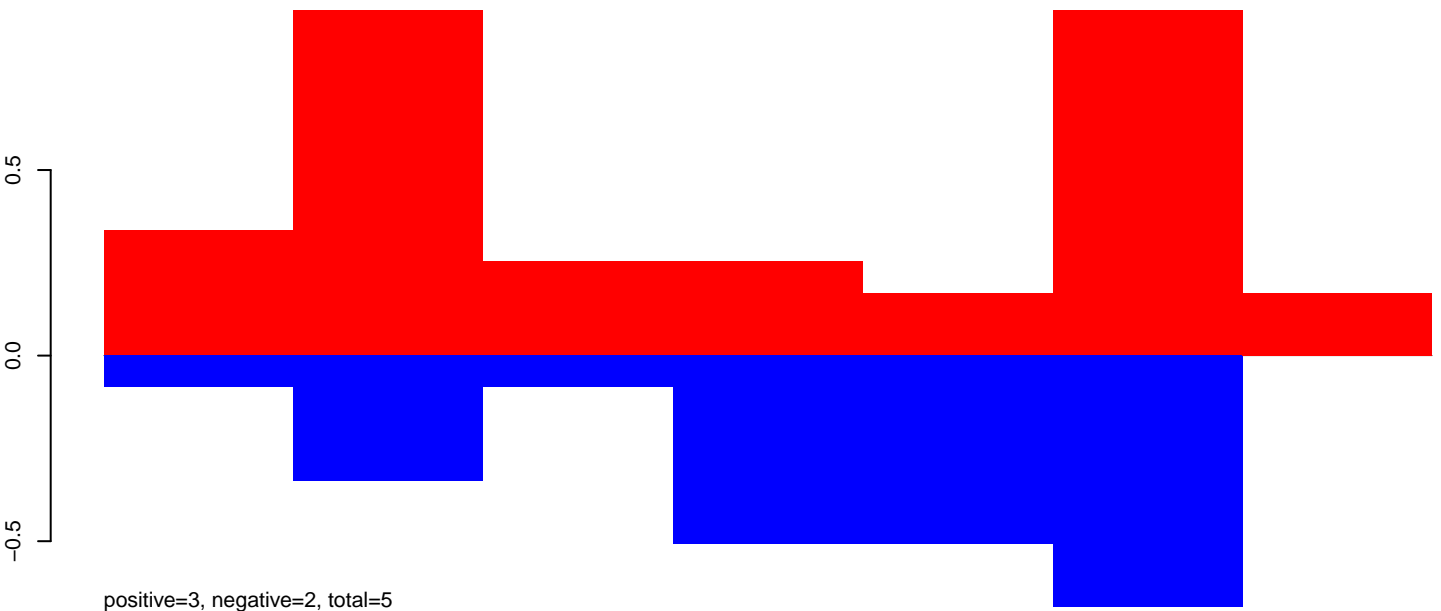
AnGam_Mos55_cells.24_35.rep



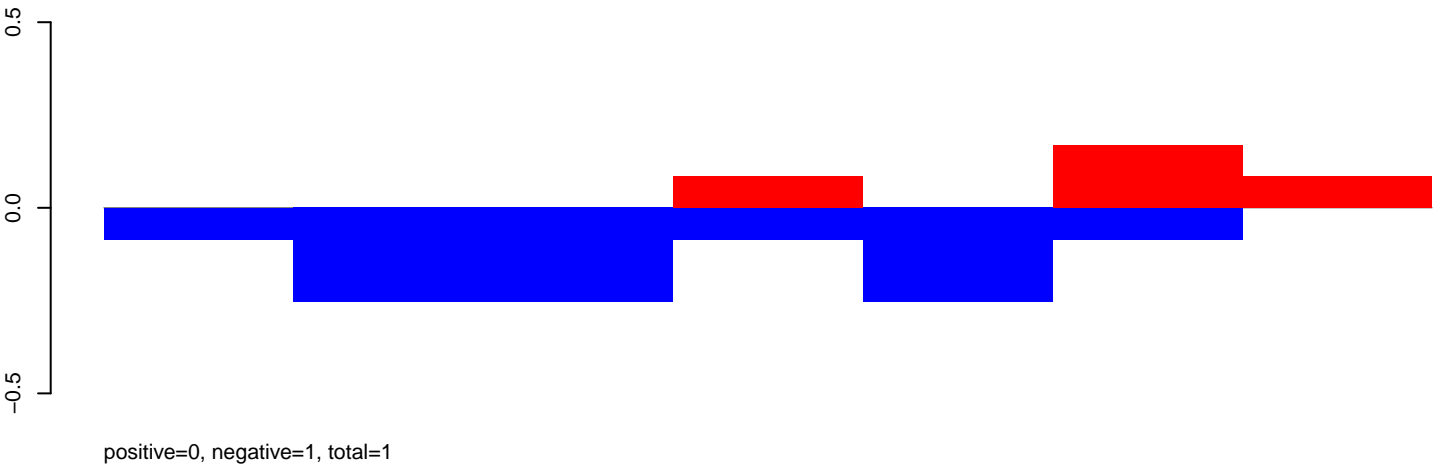
AnGam_Mos55_cells.rep



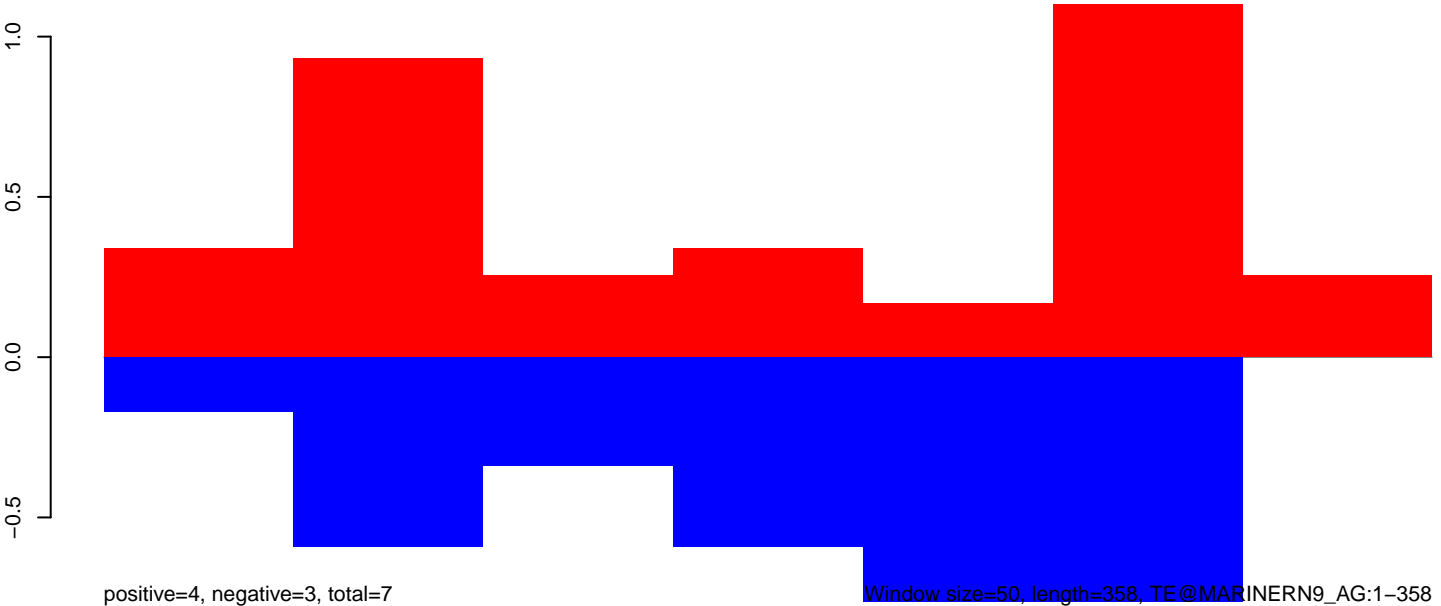
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

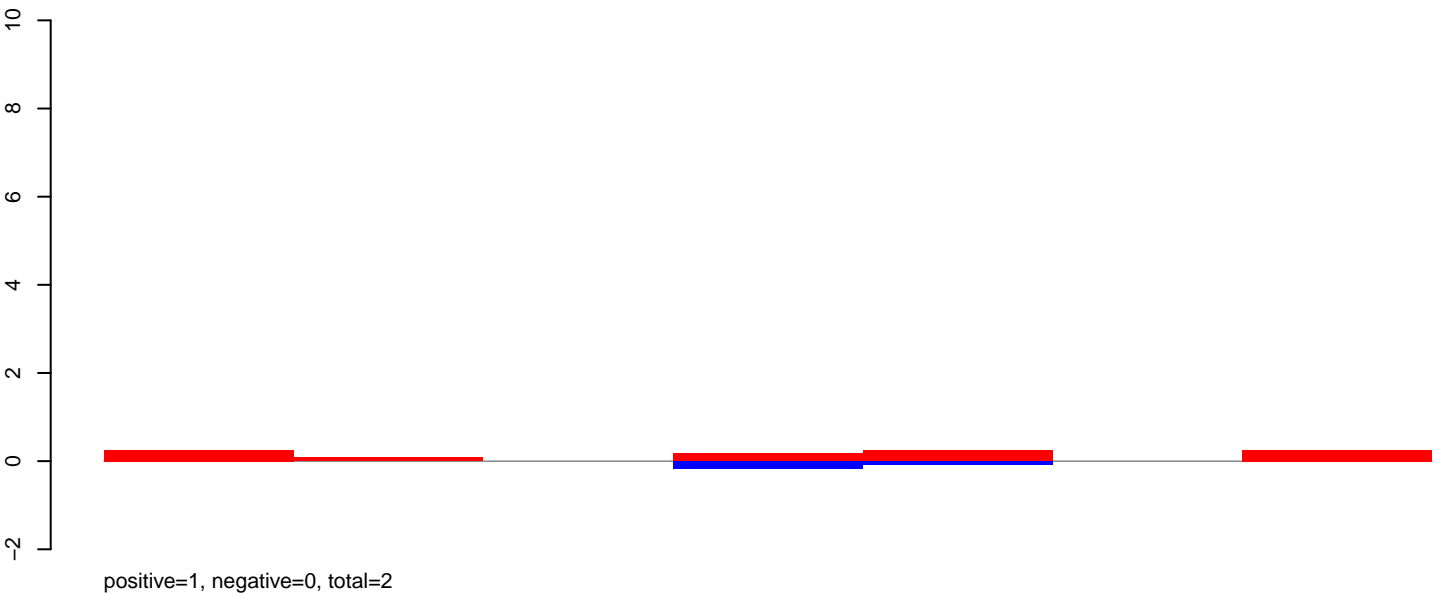


AnGam_Mos55_cells.rep



Window size=50, length=358, TE@MARINERN9_AG:1-358

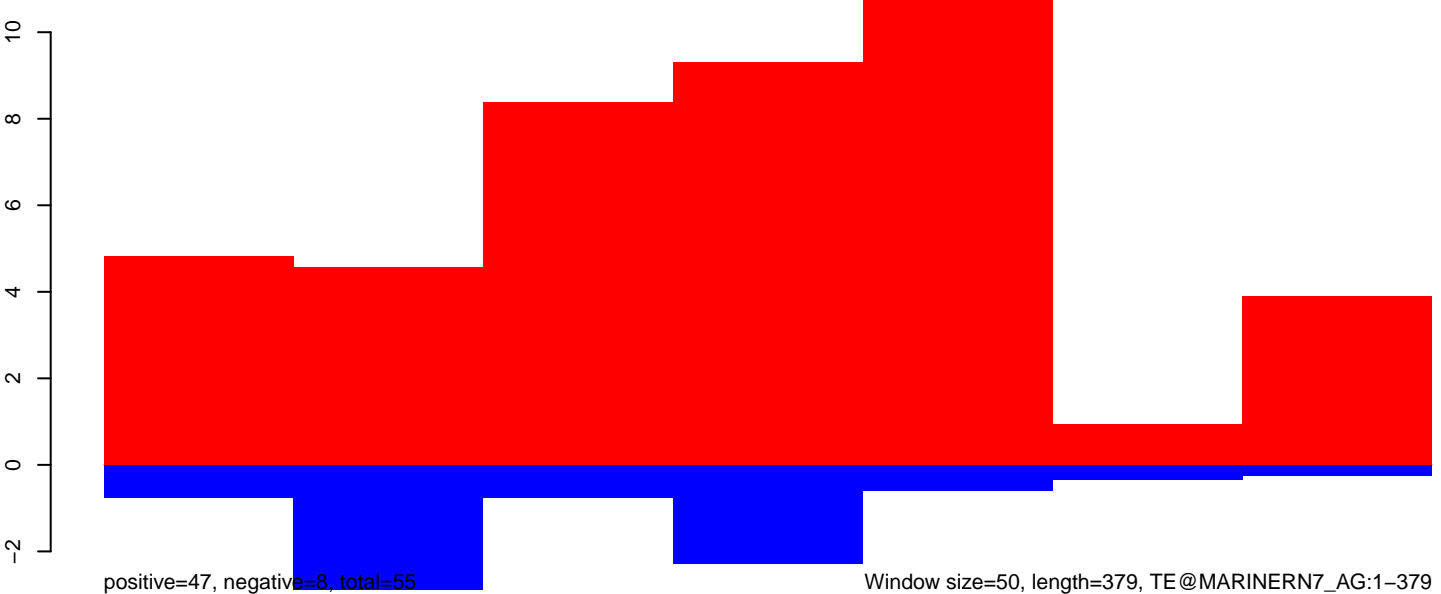
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



Window size=50, length=379, TE@MARINERN7_AG:1-379

AnGam_Mos55_cells.18_23.rep

0.05
0.00
-0.05
-0.10
-0.15
-0.20
-0.25

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.05
0.00
-0.05
-0.10
-0.15
-0.20
-0.25

positive=0, negative=1, total=1

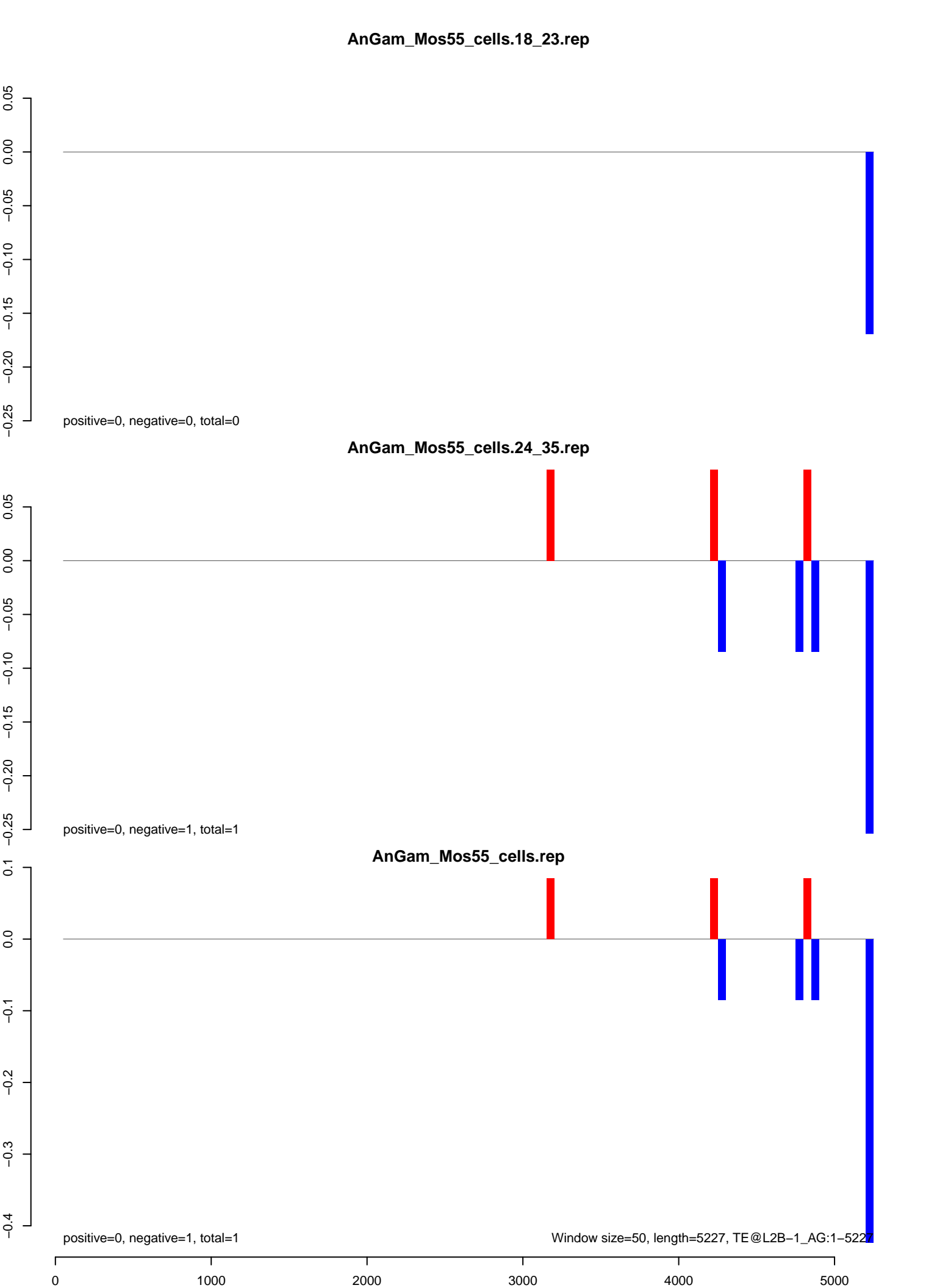
AnGam_Mos55_cells.rep

0.1
0.0
-0.1
-0.2
-0.3
-0.4

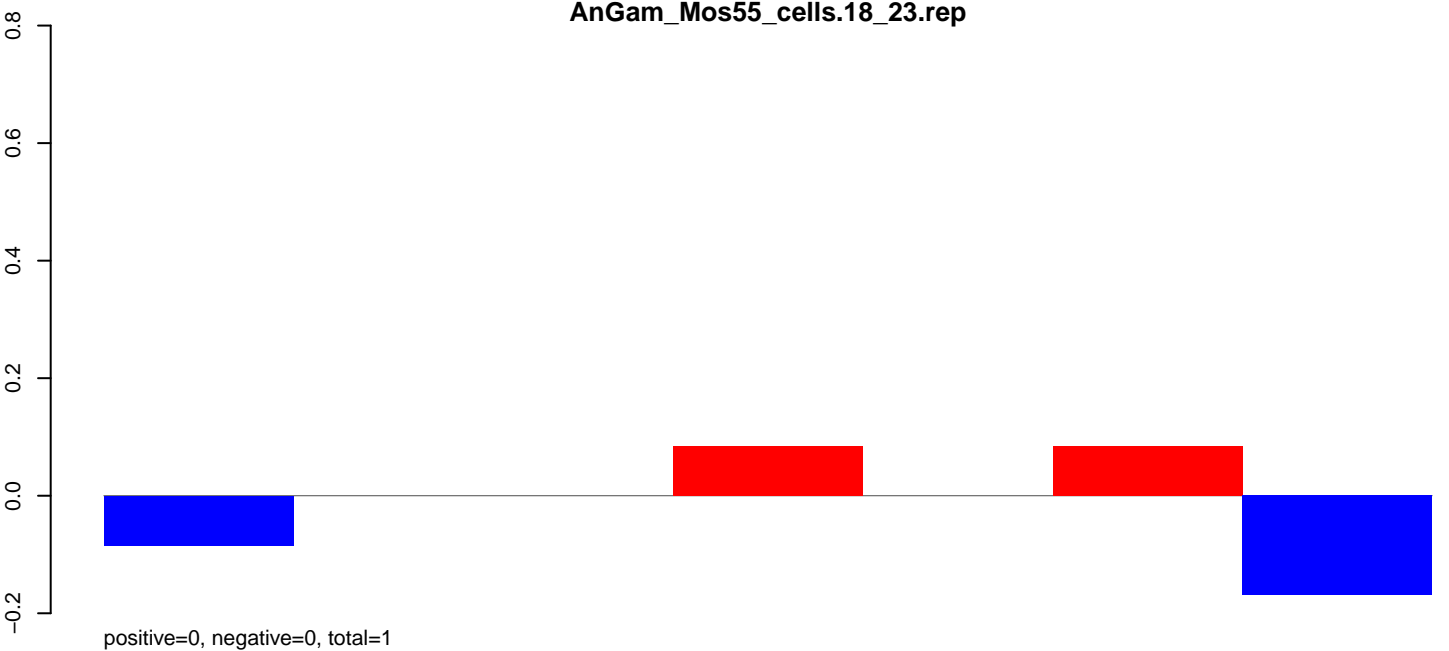
positive=0, negative=1, total=1

Window size=50, length=5227, TE@L2B-1_AG:1-5227

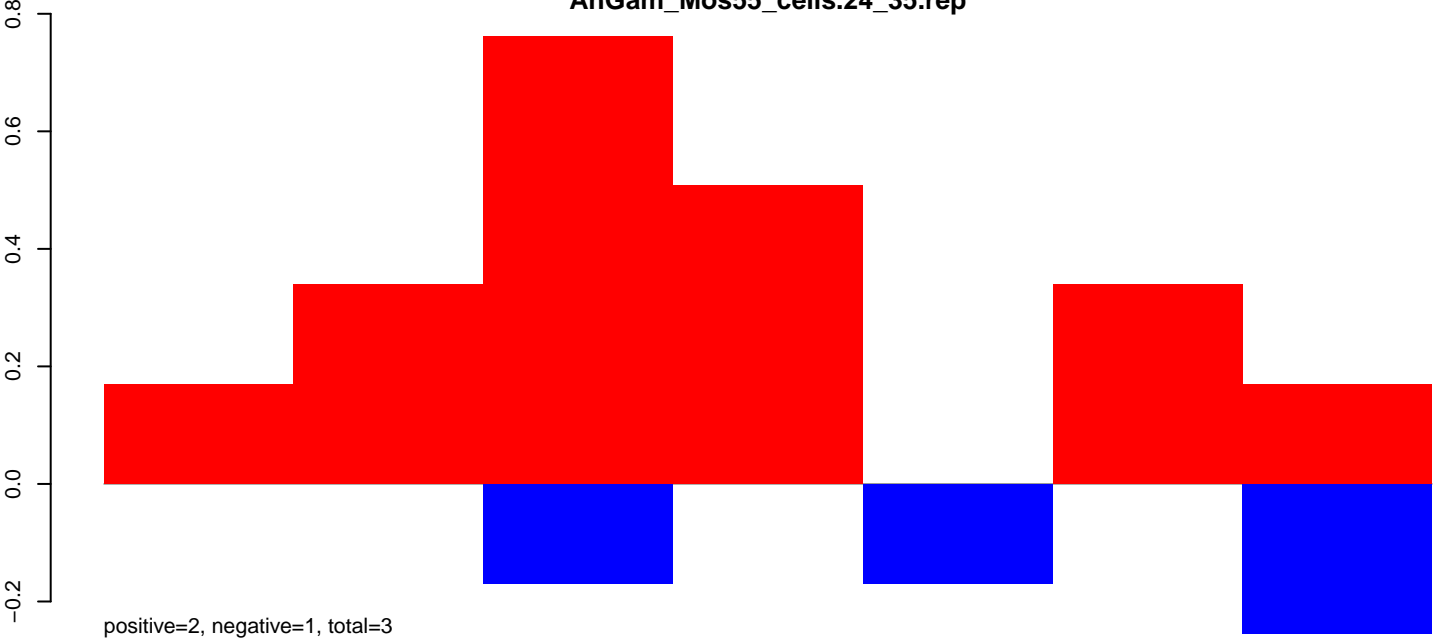
0 1000 2000 3000 4000 5000



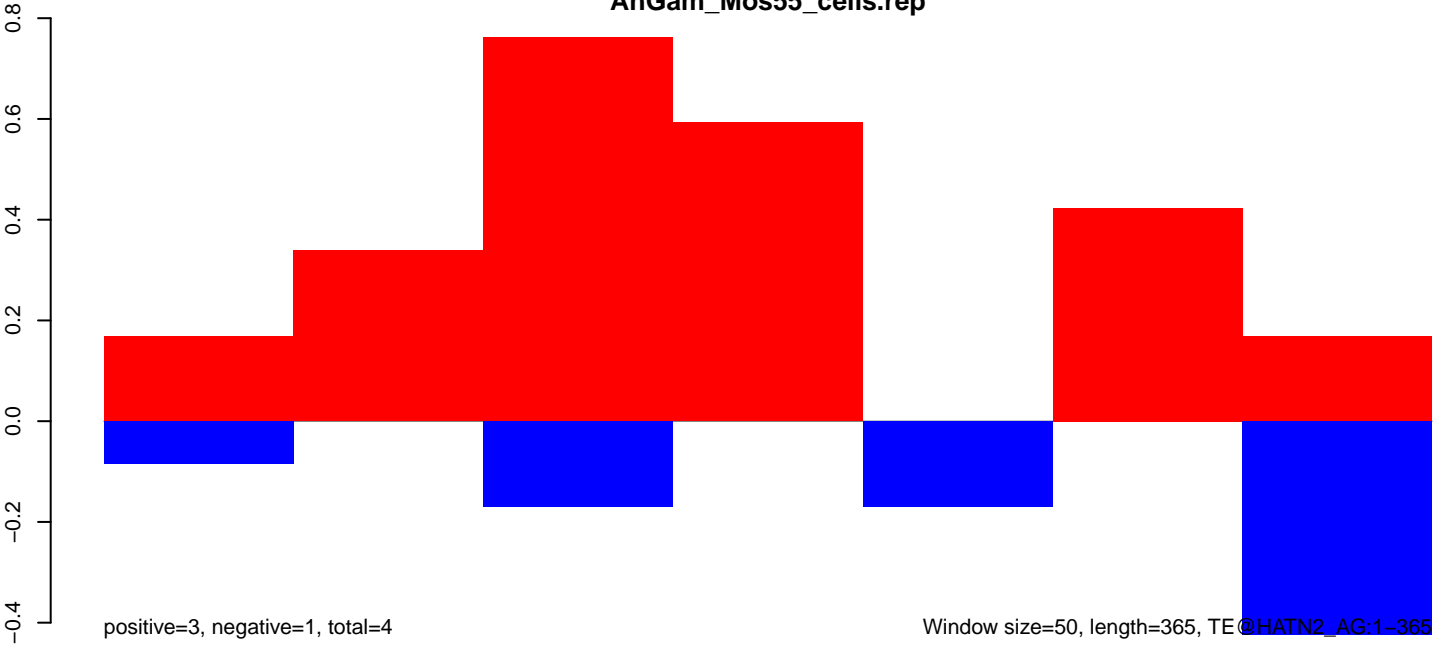
AnGam_Mos55_cells.18_23.rep



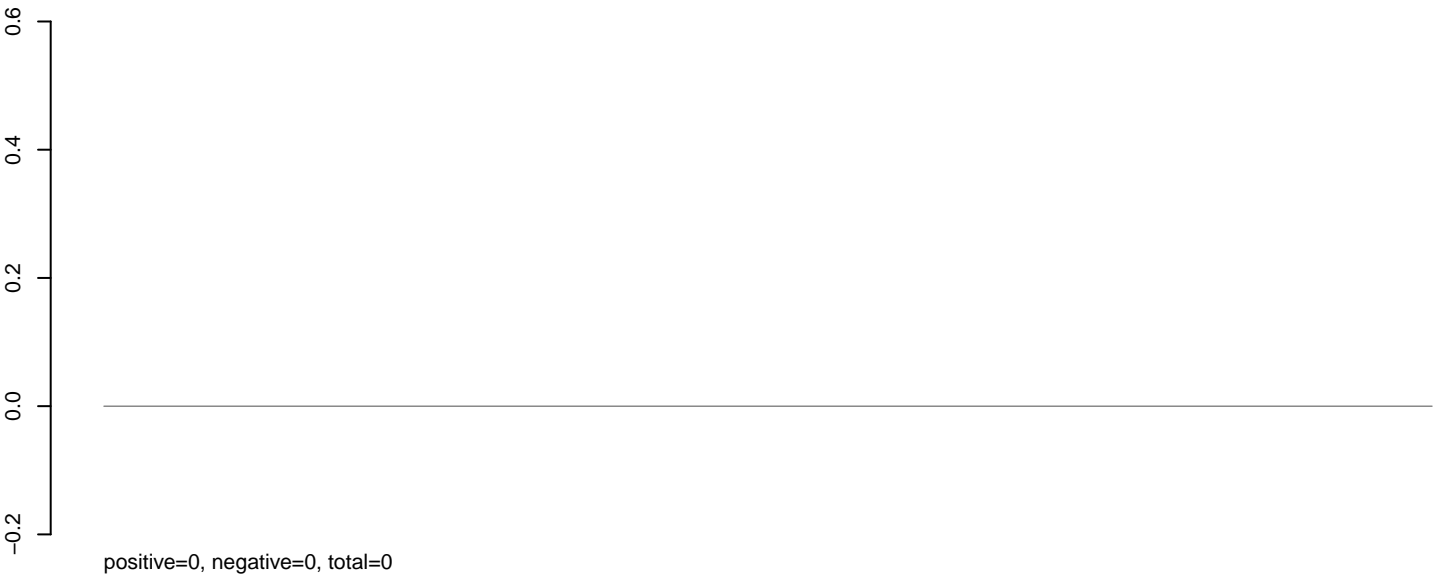
AnGam_Mos55_cells.24_35.rep



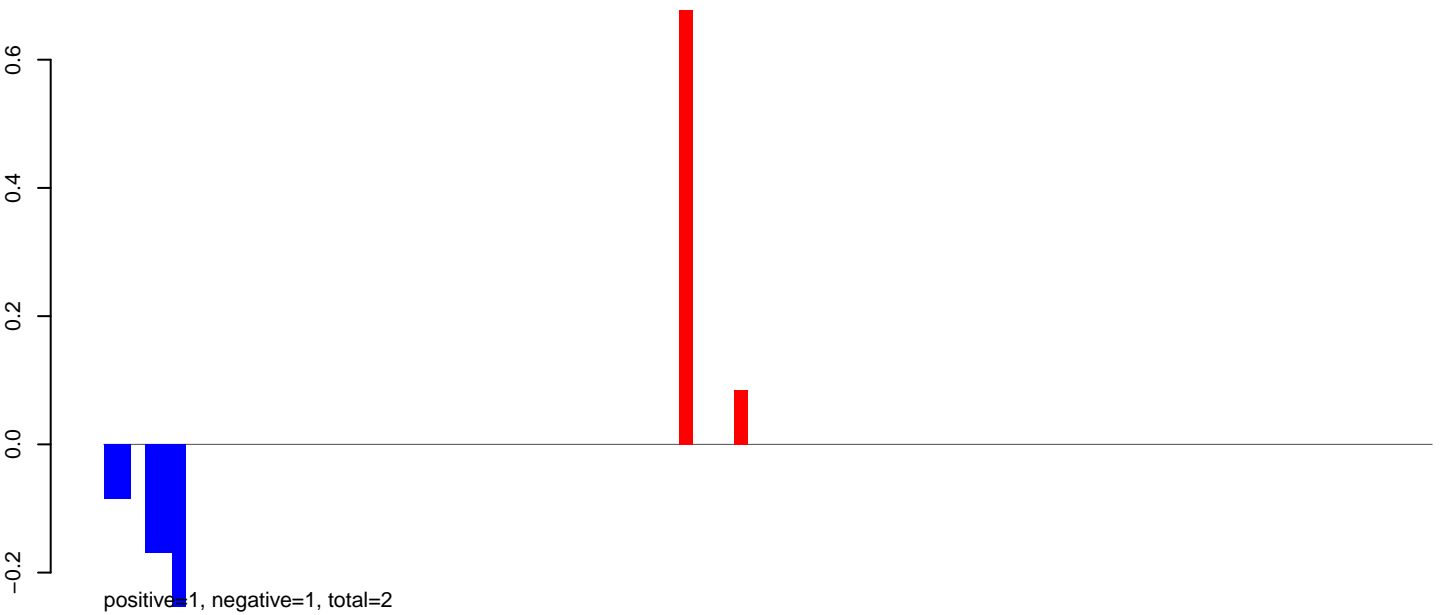
AnGam_Mos55_cells.rep



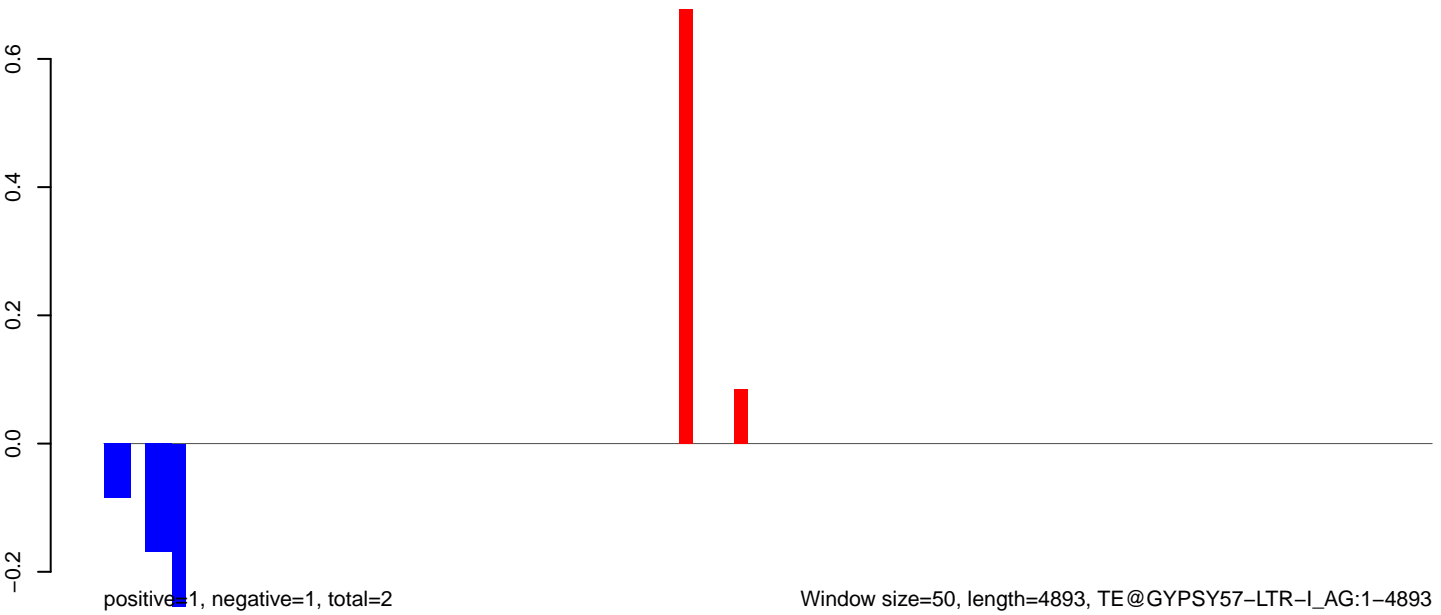
AnGam_Mos55_cells.18_23.rep



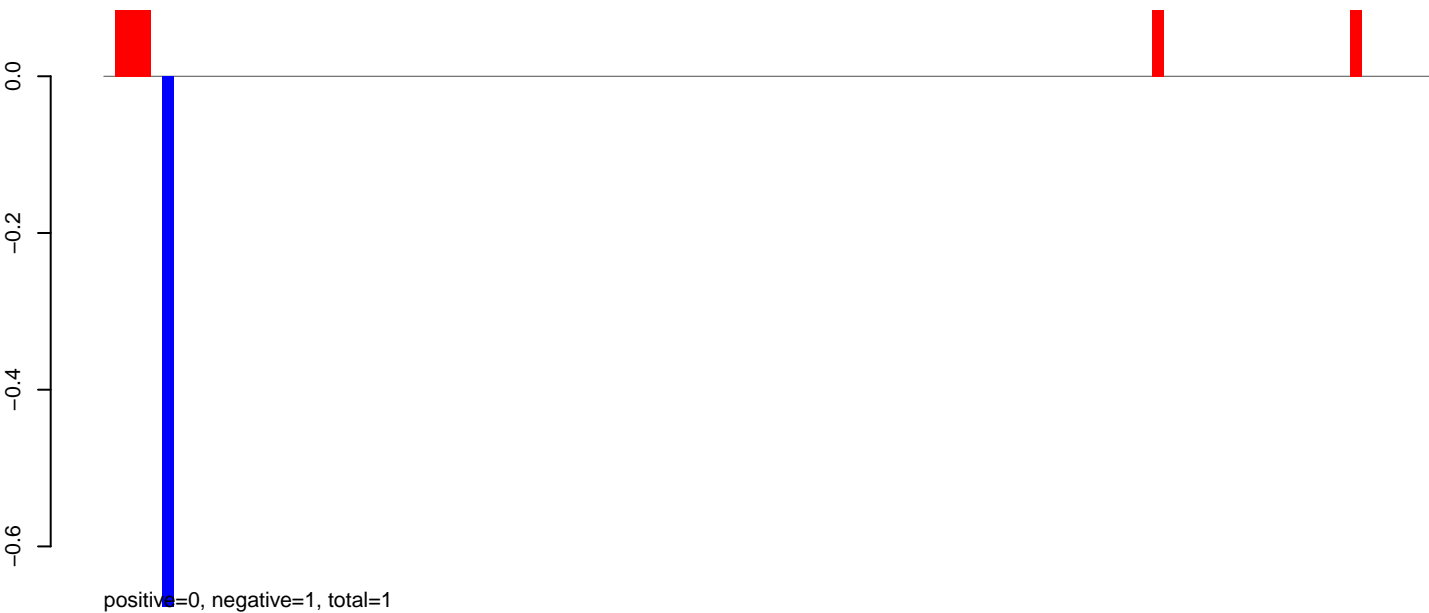
AnGam_Mos55_cells.24_35.rep



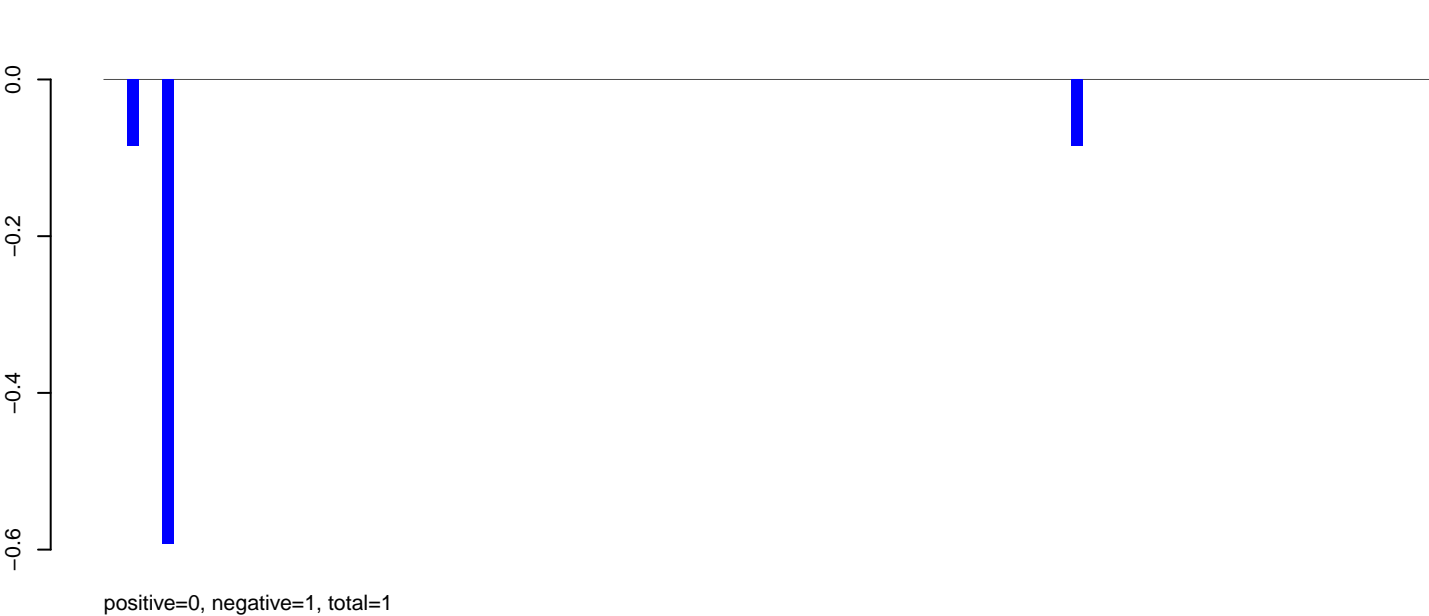
AnGam_Mos55_cells.rep



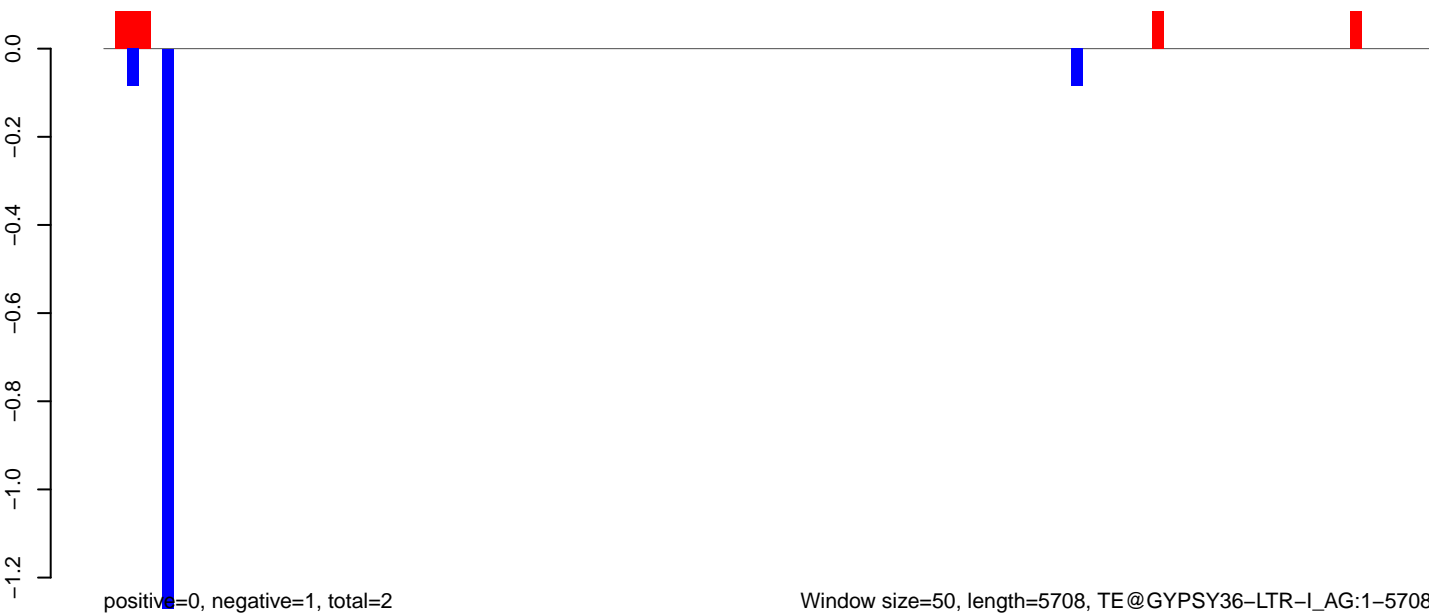
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

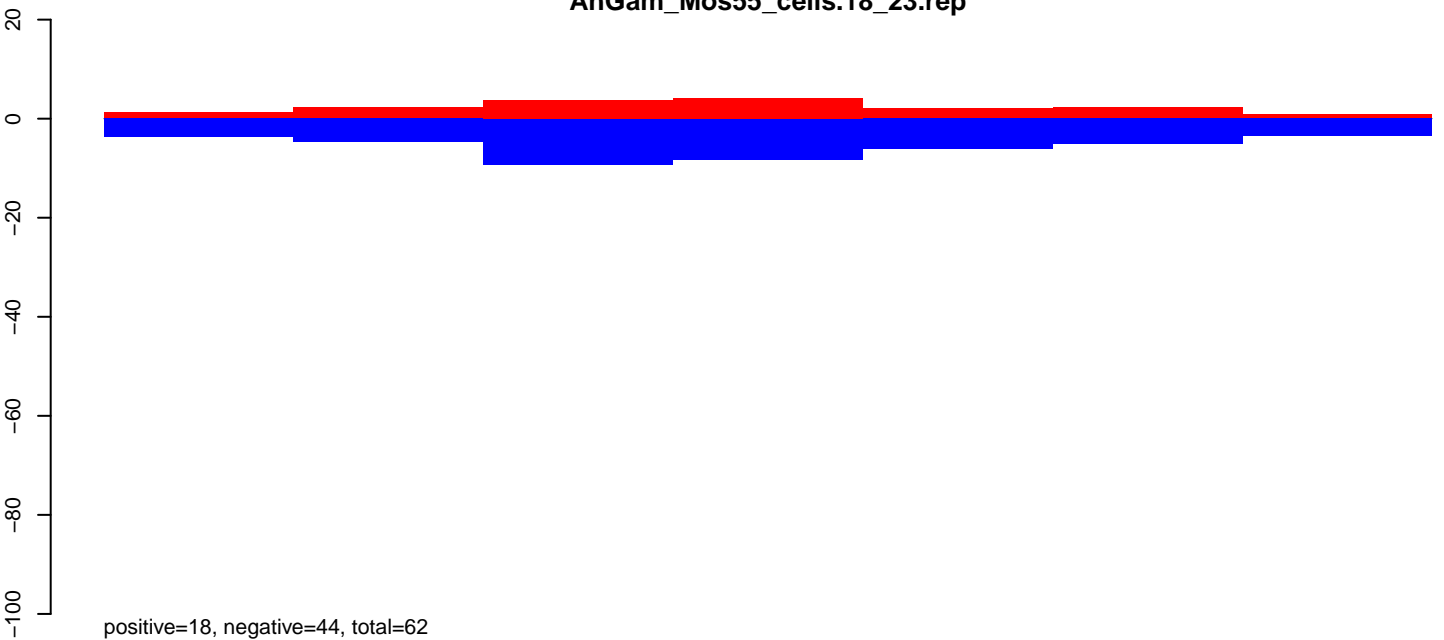


AnGam_Mos55_cells.rep

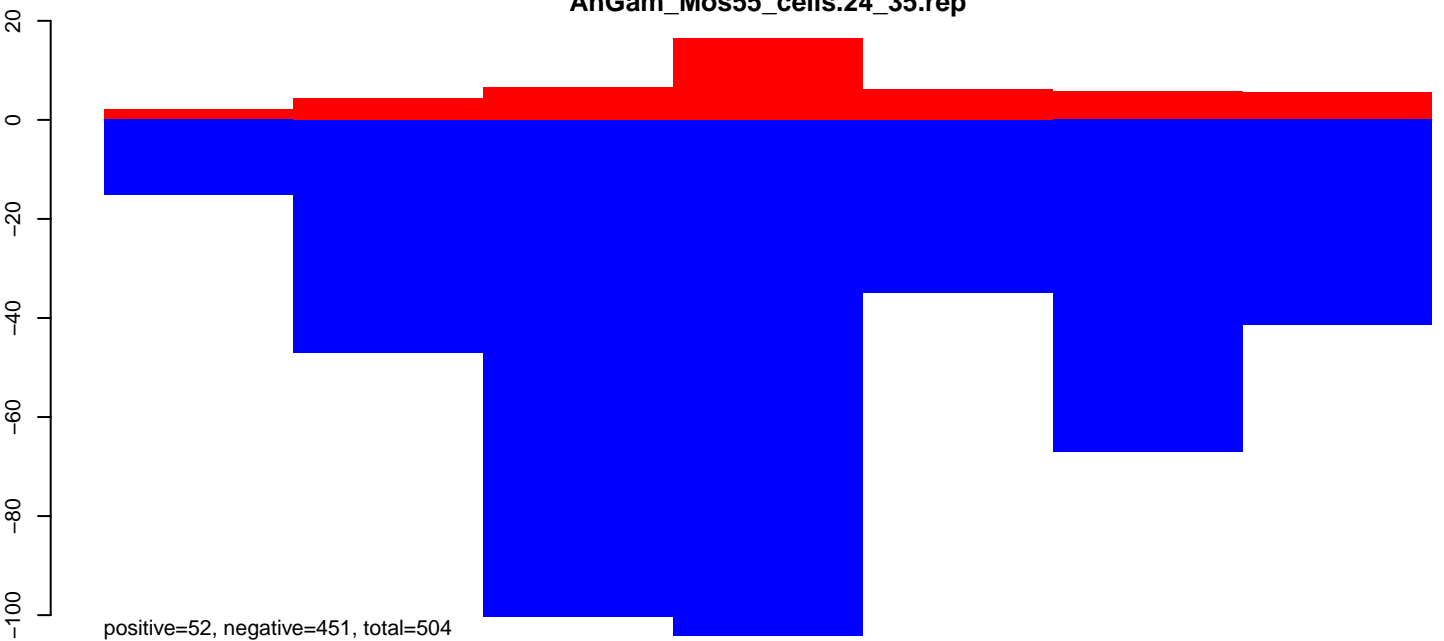


Window size=50, length=5708, TE@GYPSY36-LTR-L_AG:1-5708

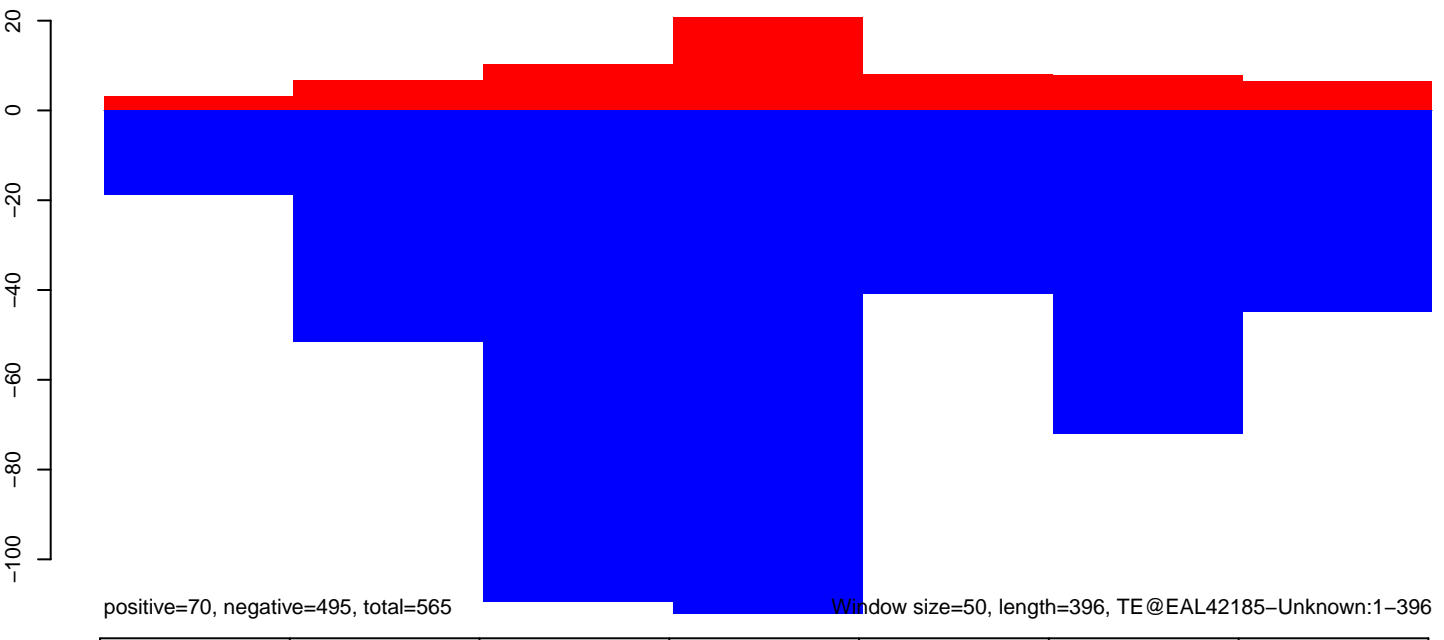
AnGam_Mos55_cells.18_23.rep



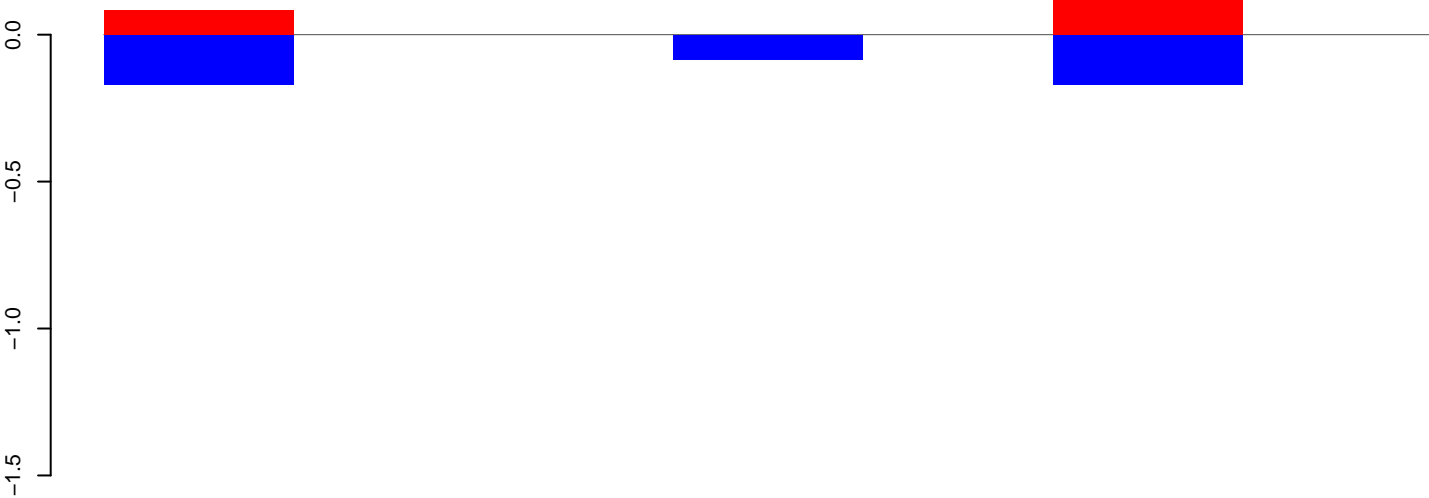
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

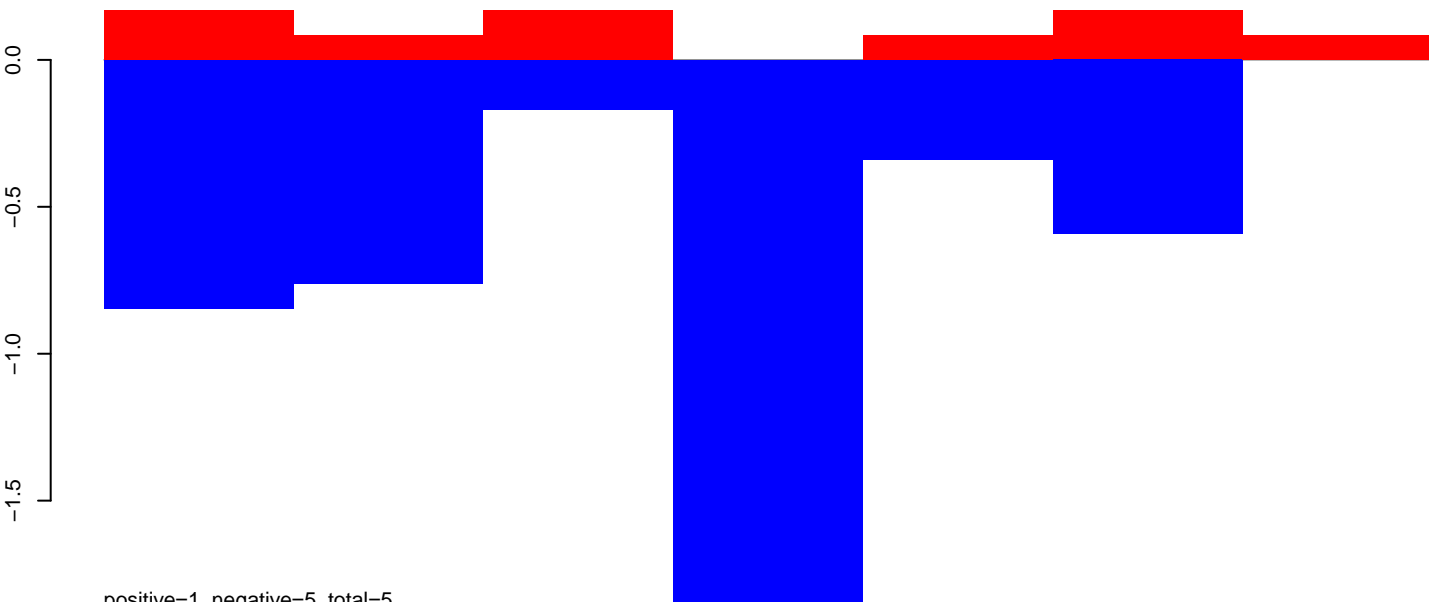


AnGam_Mos55_cells.18_23.rep



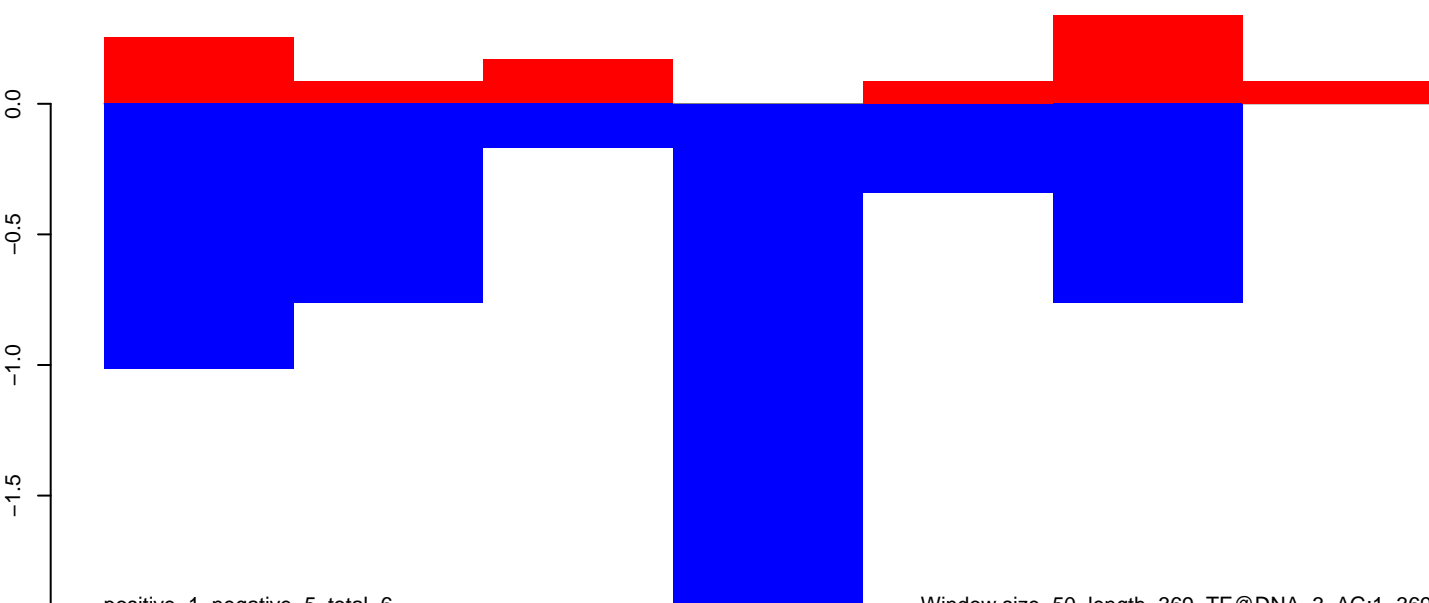
positive=0, negative=0, total=1

AnGam_Mos55_cells.24_35.rep



positive=1, negative=5, total=5

AnGam_Mos55_cells.rep

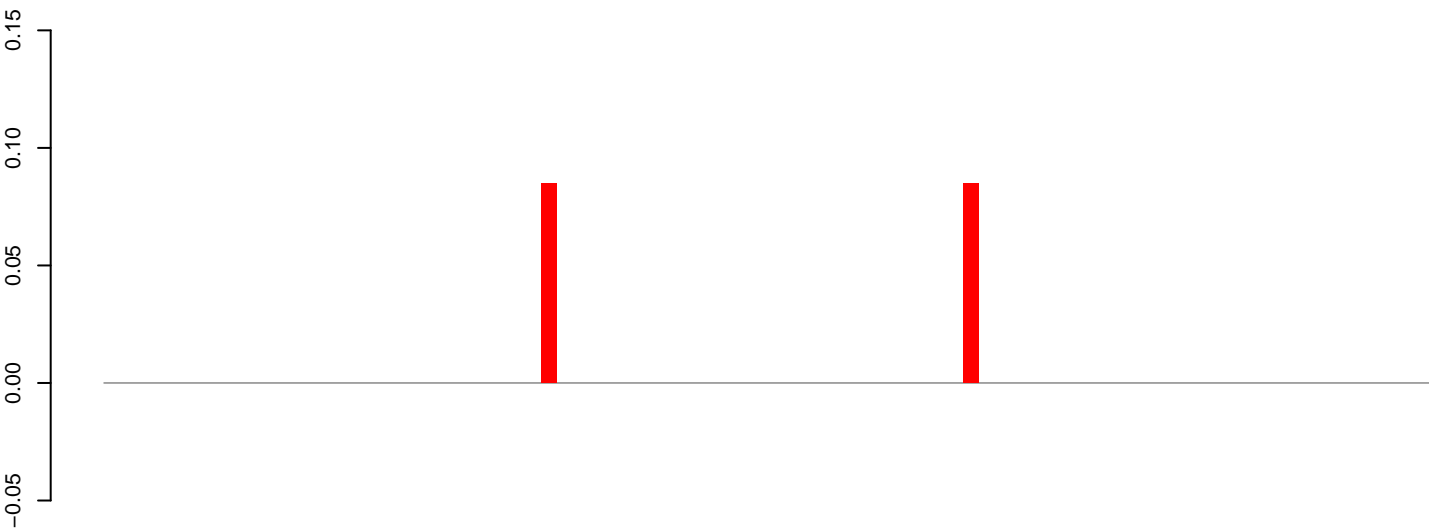


positive=1, negative=5, total=6

Window size=50, length=369, TE@DNA-3_AG:1-369

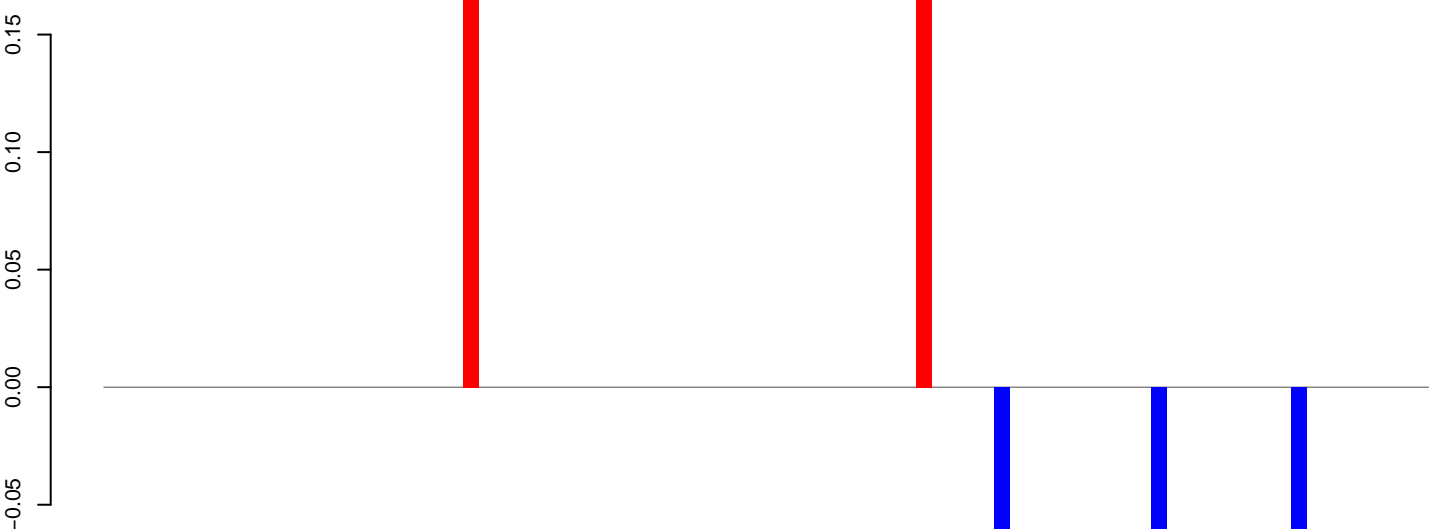
50 100 150 200 250 300 350 400

AnGam_Mos55_cells.18_23.rep



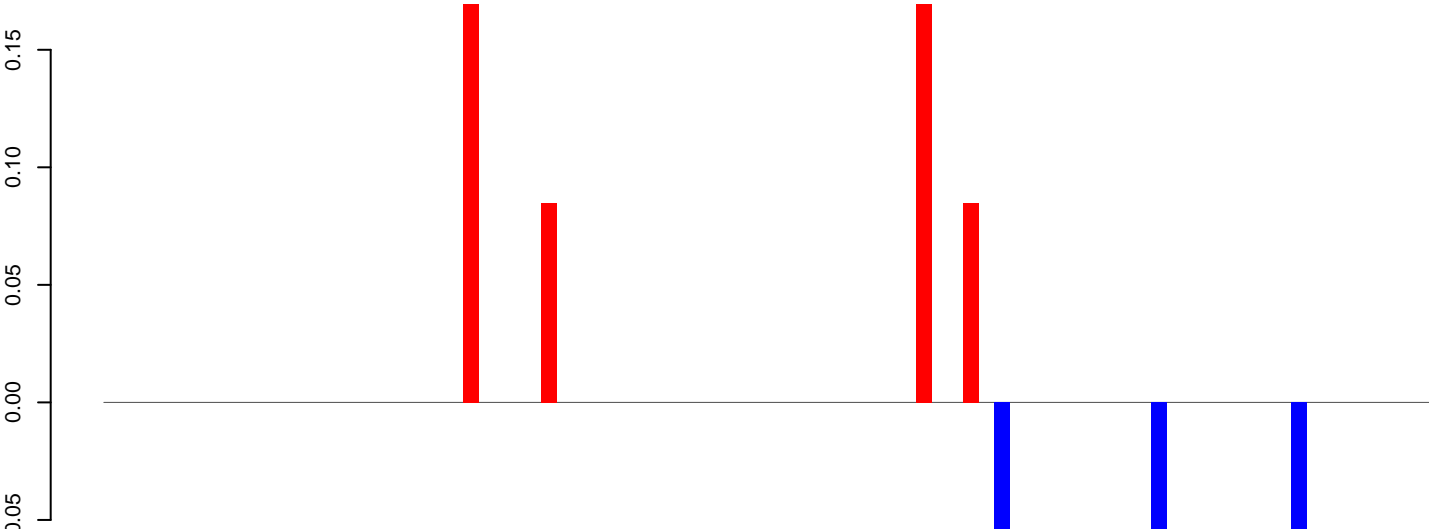
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=1

AnGam_Mos55_cells.rep

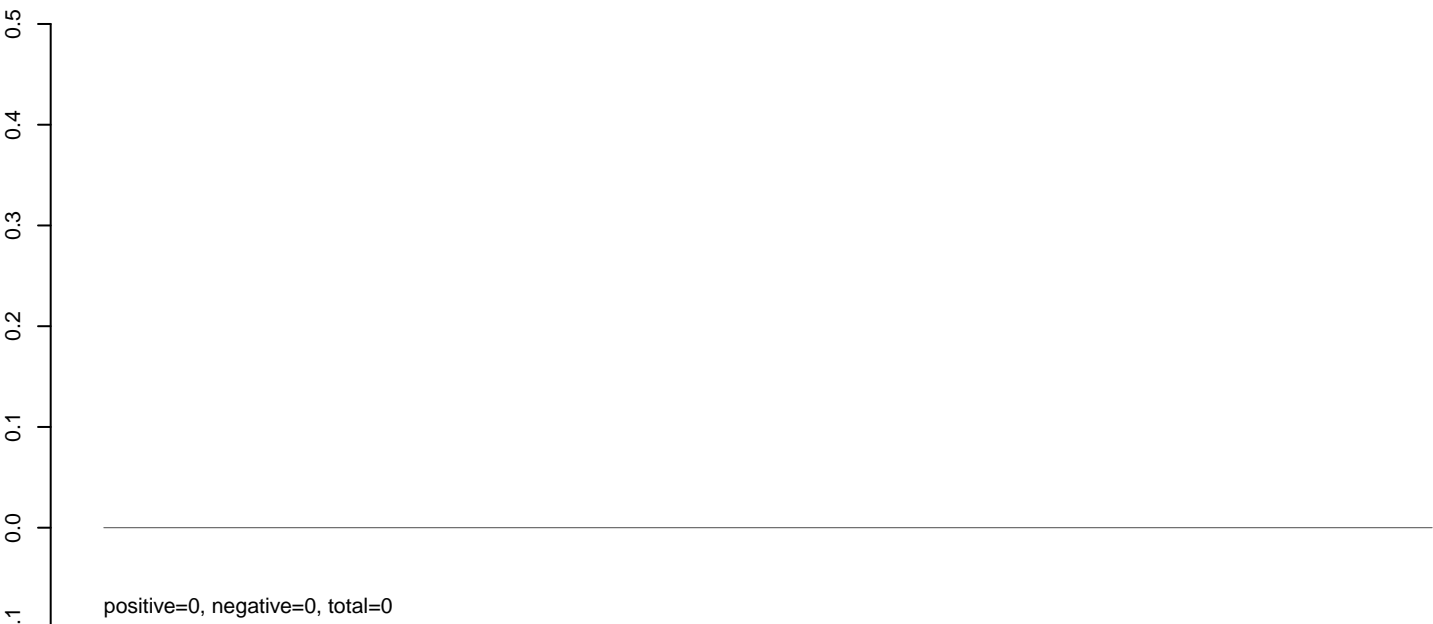


positive=1, negative=0, total=1

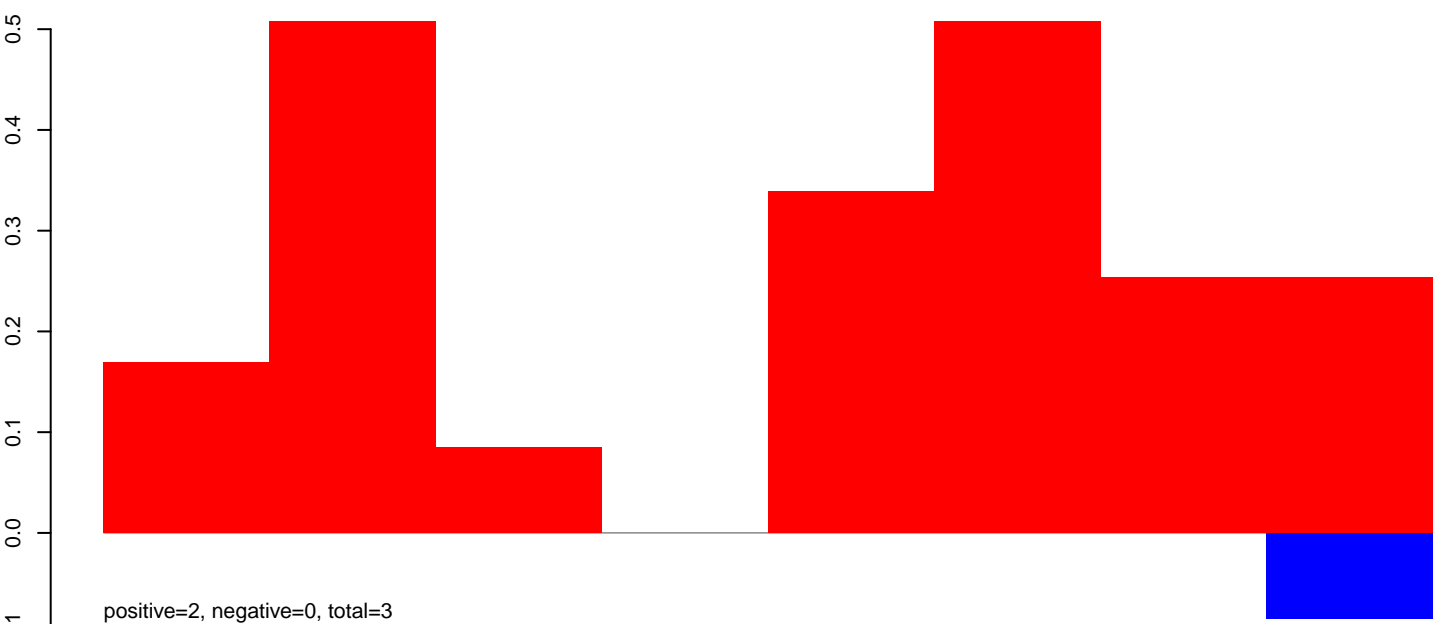
Window size=50, length=4295, TE@Copia-8_AG-LTR-I:1-4295

0 1000 2000 3000 4000

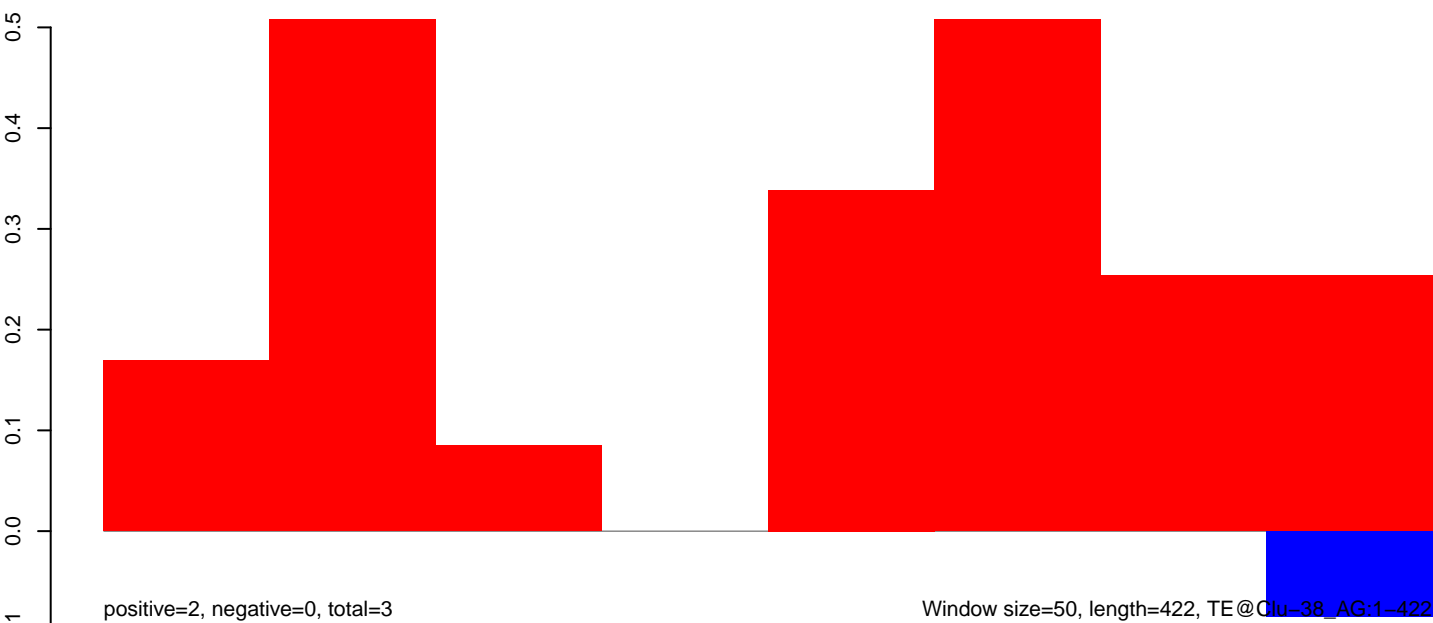
AnGam_Mos55_cells.18_23.rep



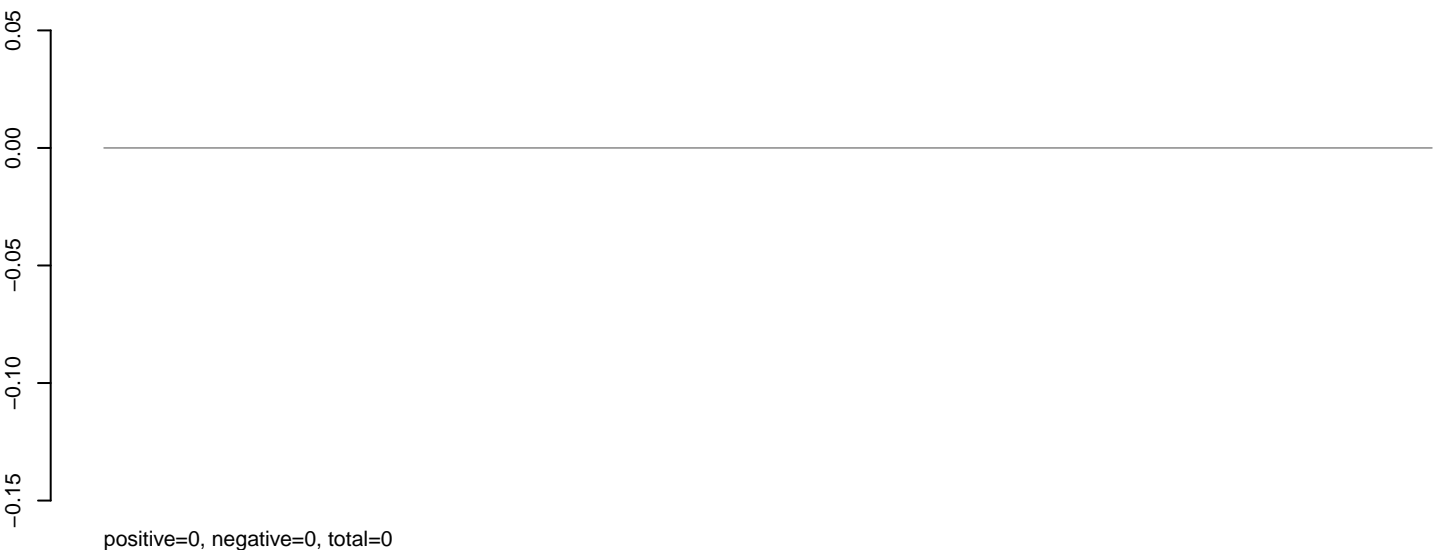
AnGam_Mos55_cells.24_35.rep



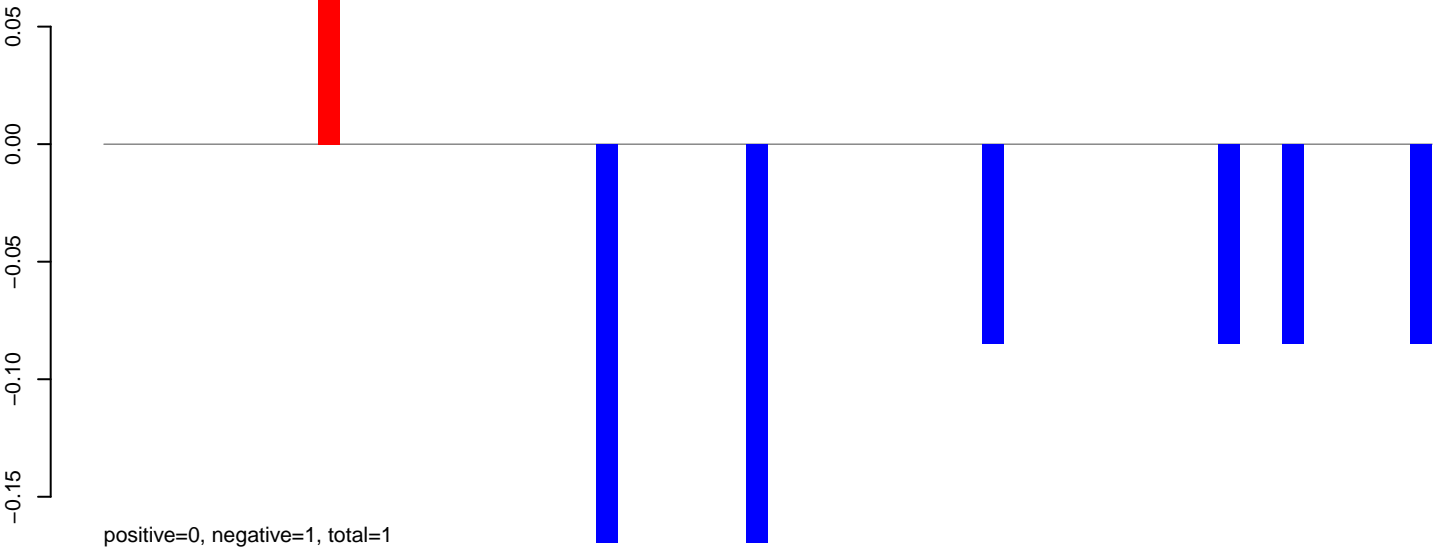
AnGam_Mos55_cells.rep



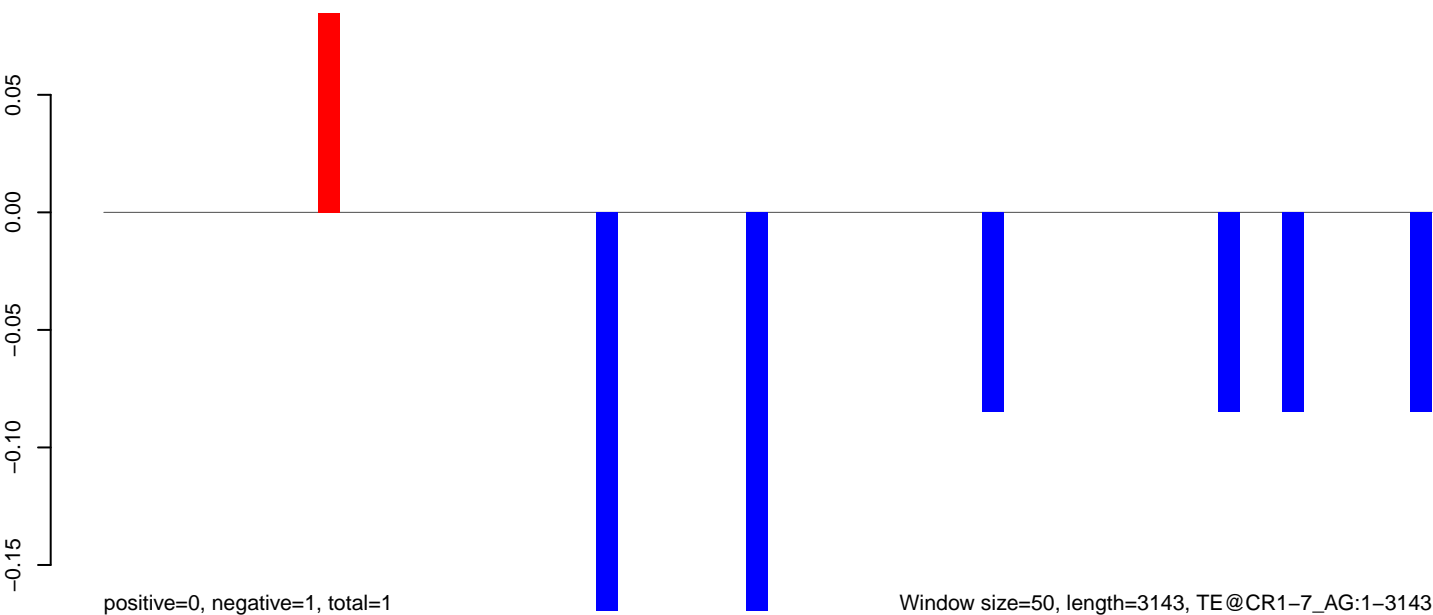
AnGam_Mos55_cells.18_23.rep



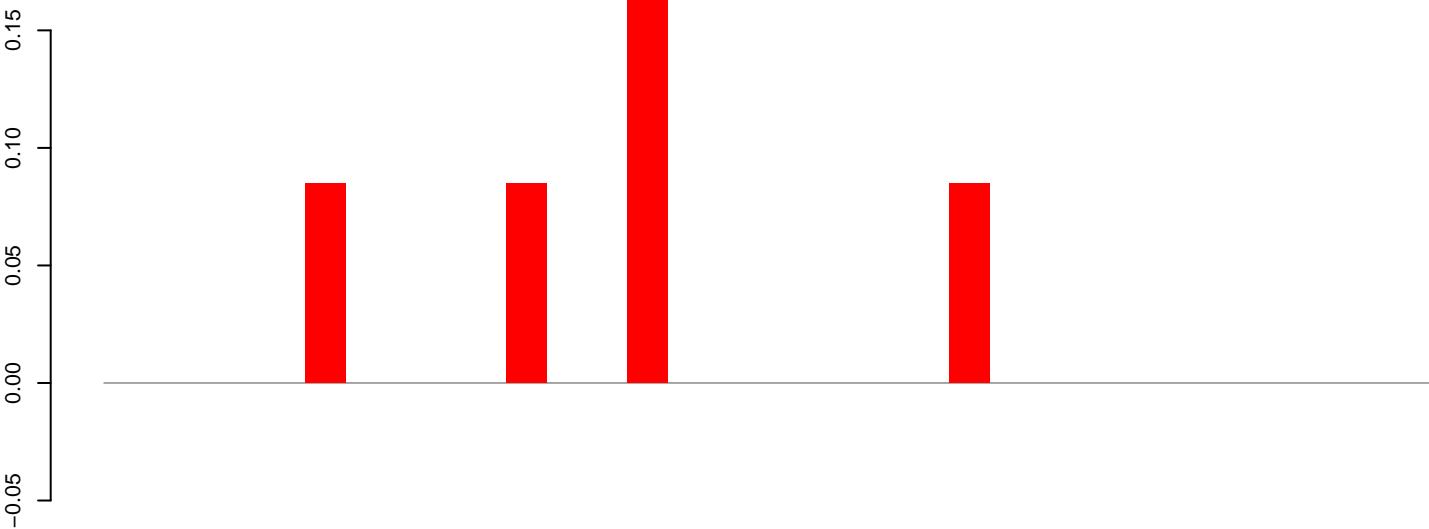
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

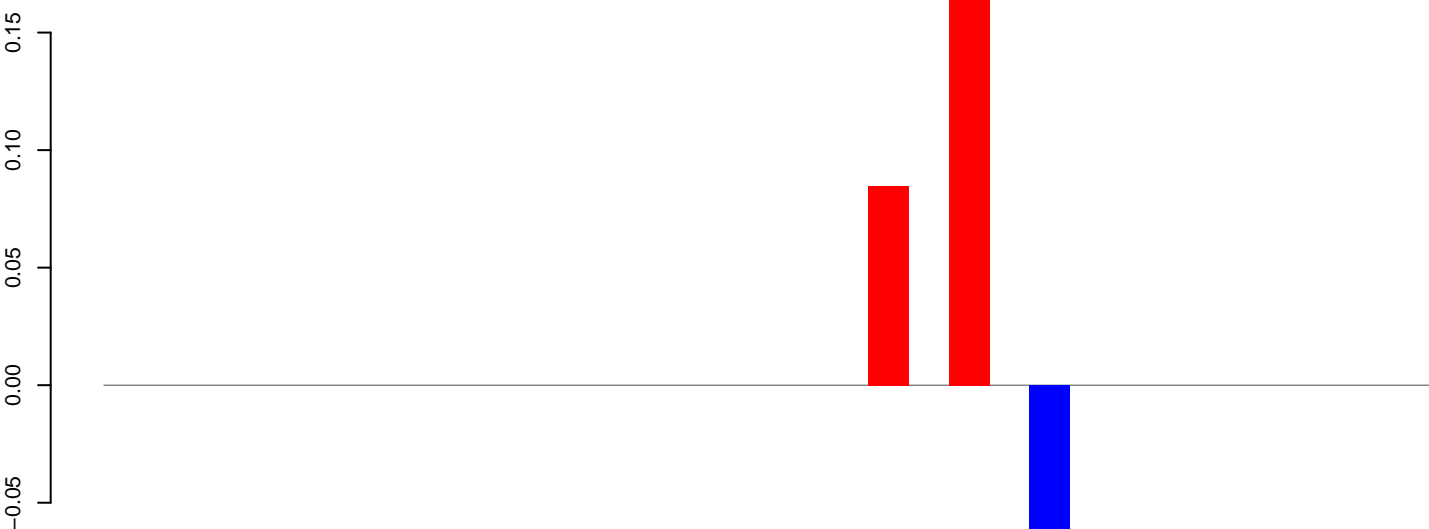


AnGam_Mos55_cells.18_23.rep



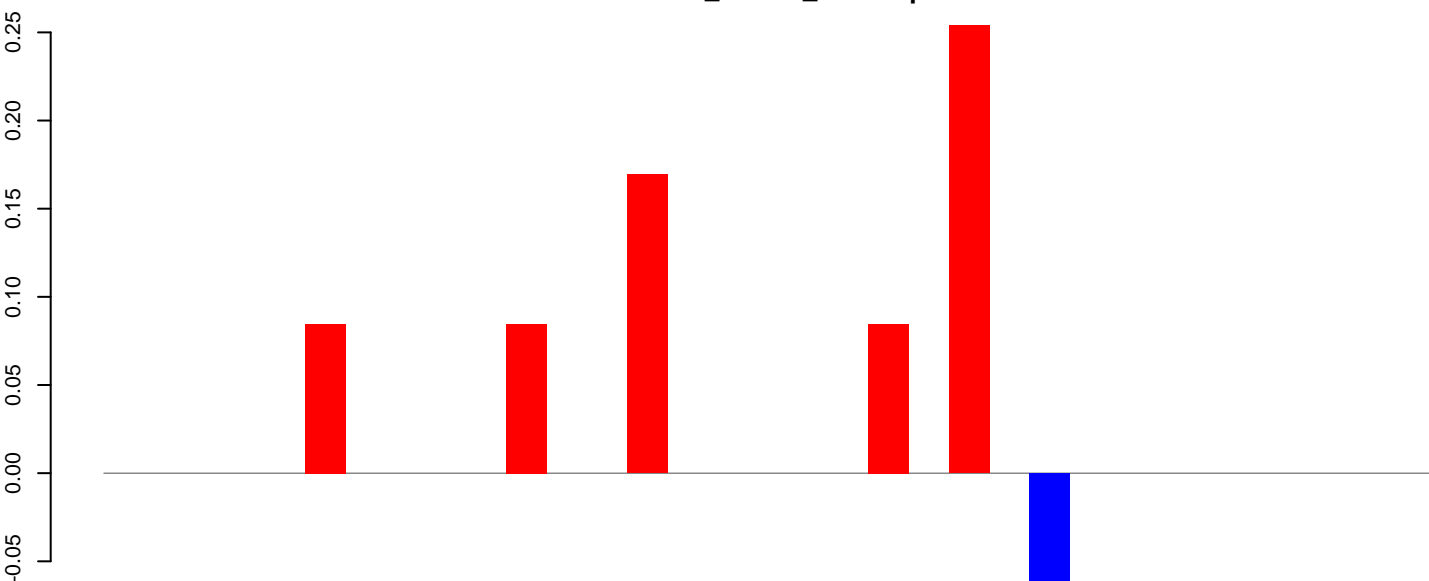
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

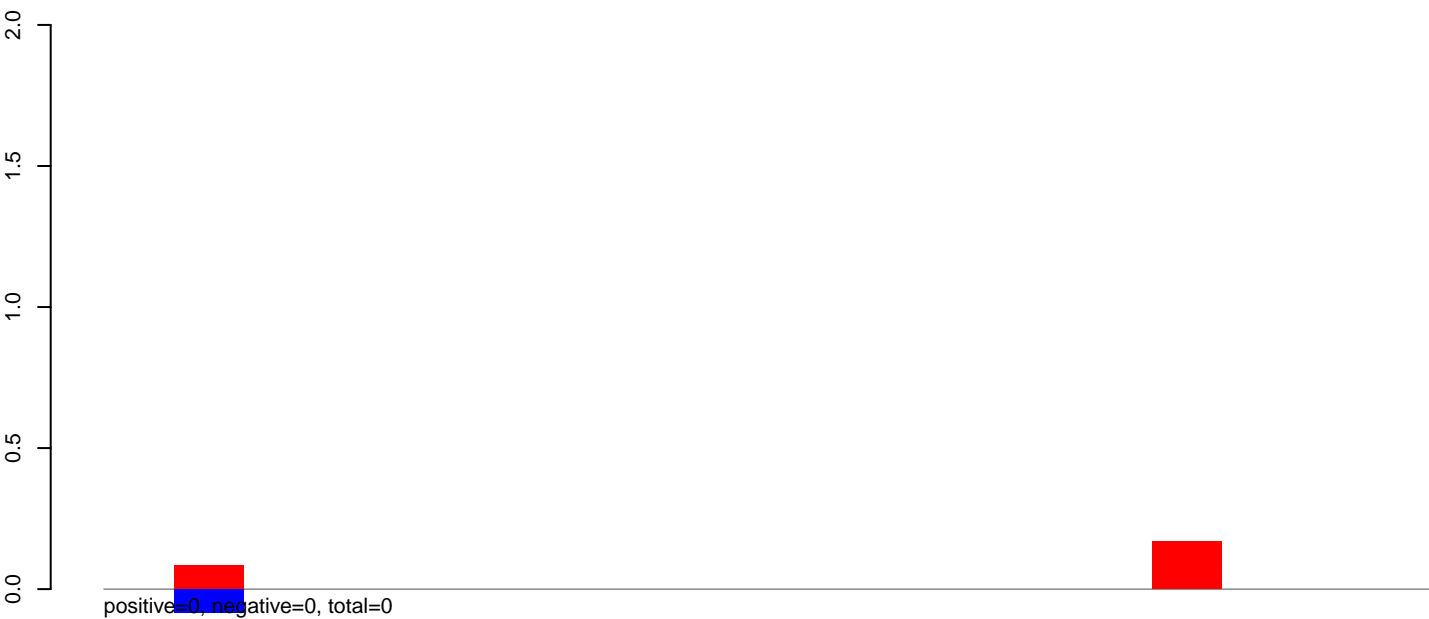
AnGam_Mos55_cells.rep



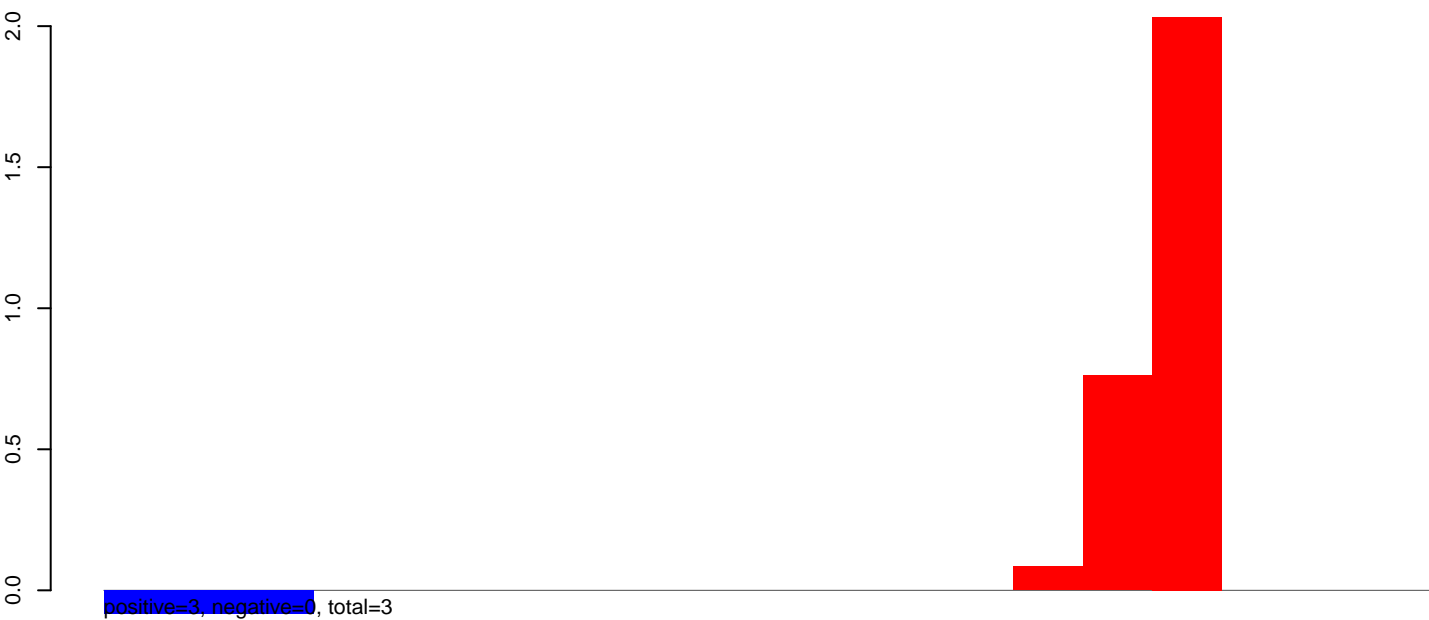
positive=1, negative=0, total=1

Window size=50, length=1661, TE@piggyBacN1_AG:1-1661

AnGam_Mos55_cells.18_23.rep



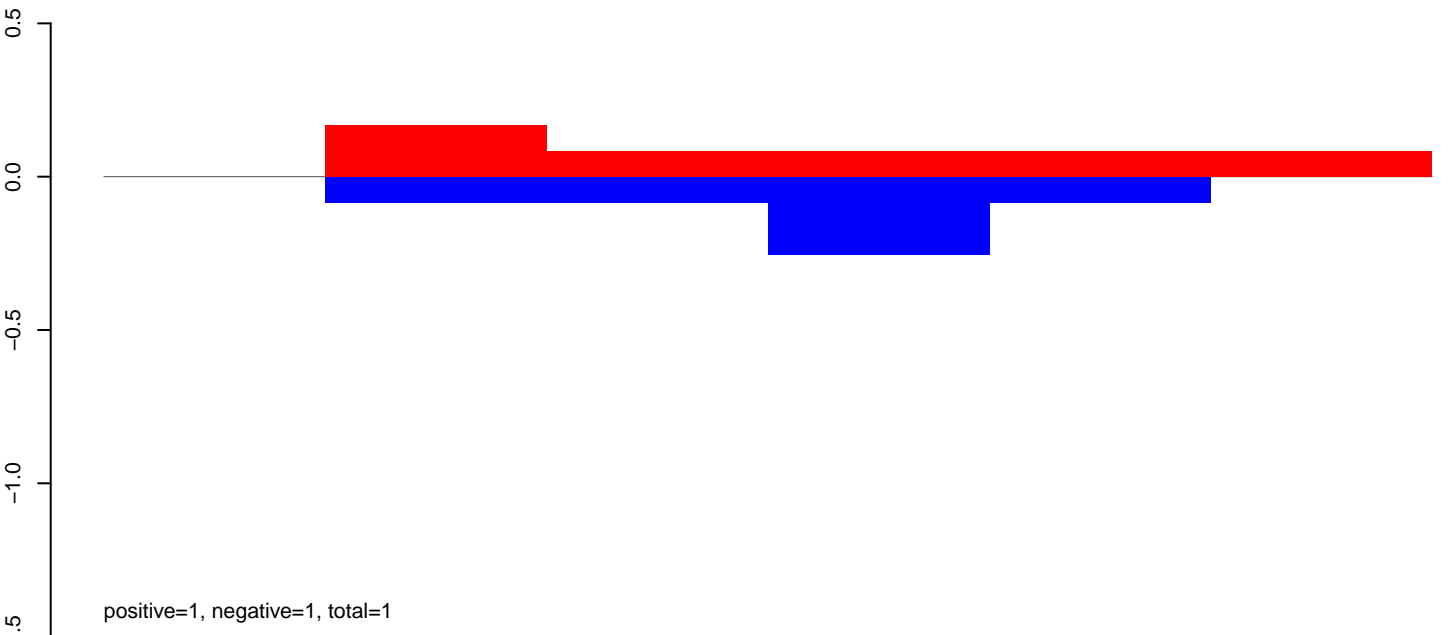
AnGam_Mos55_cells.24_35.rep



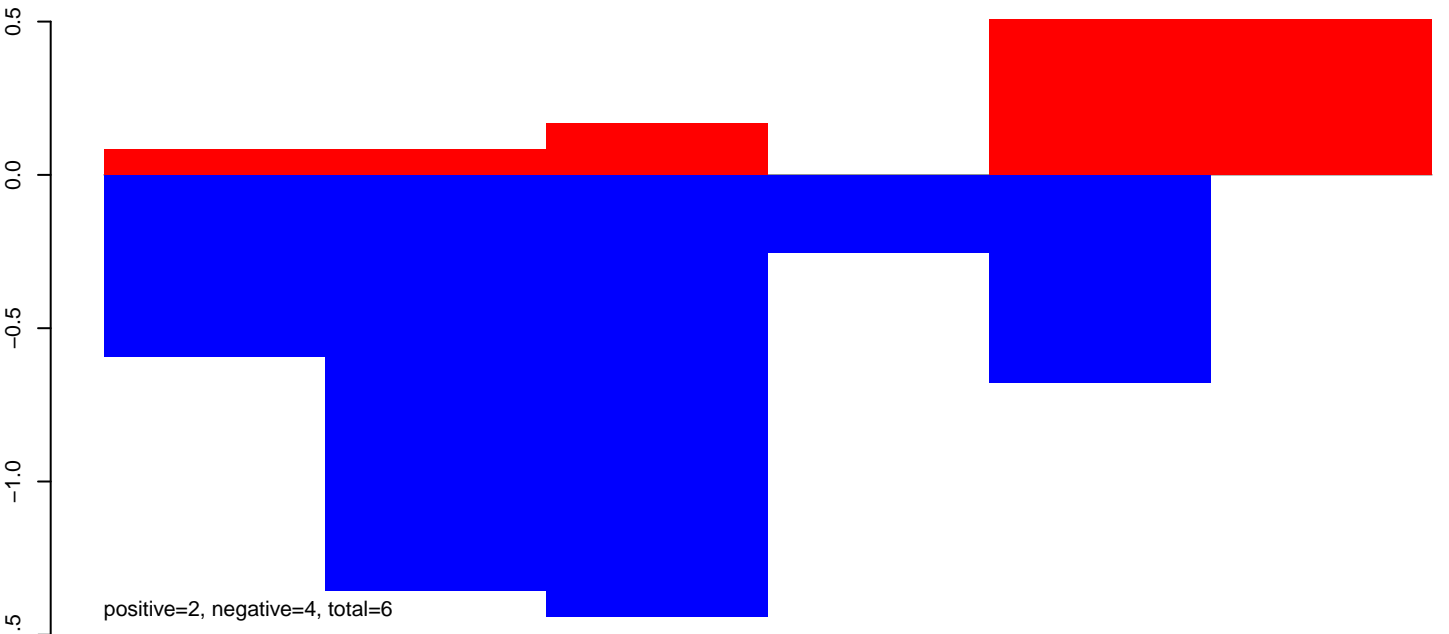
AnGam_Mos55_cells.rep



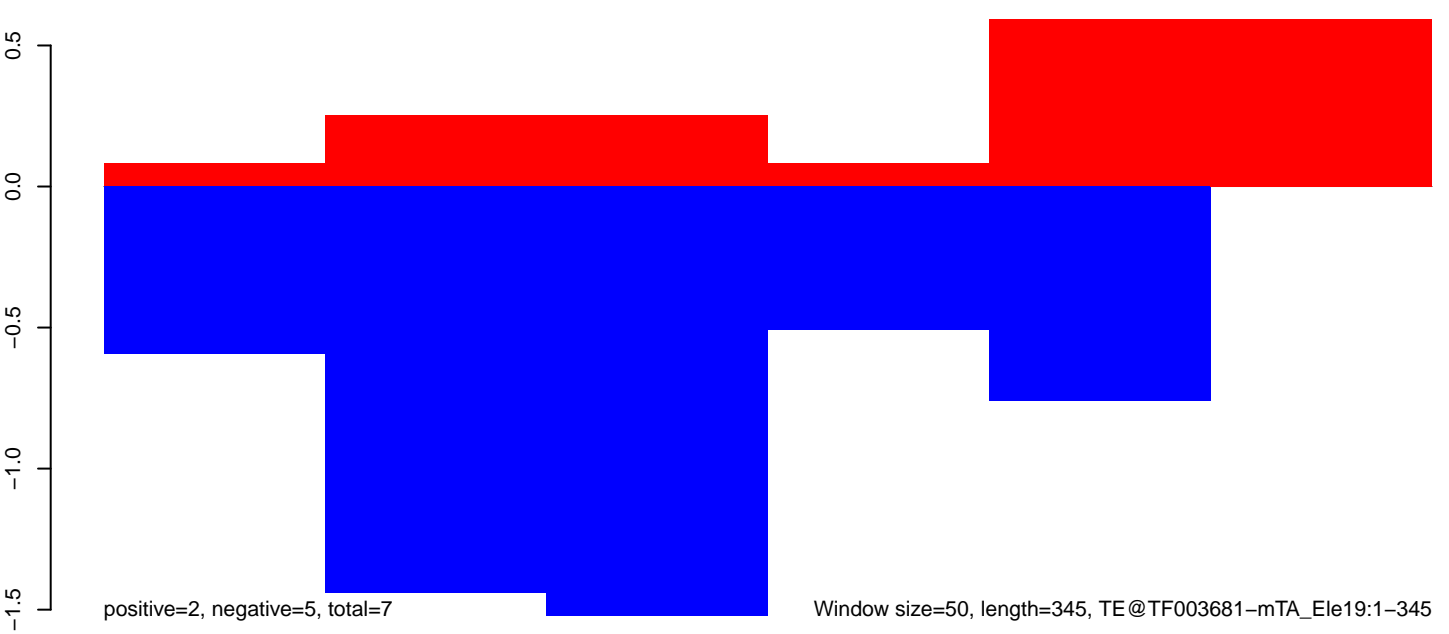
AnGam_Mos55_cells.18_23.rep



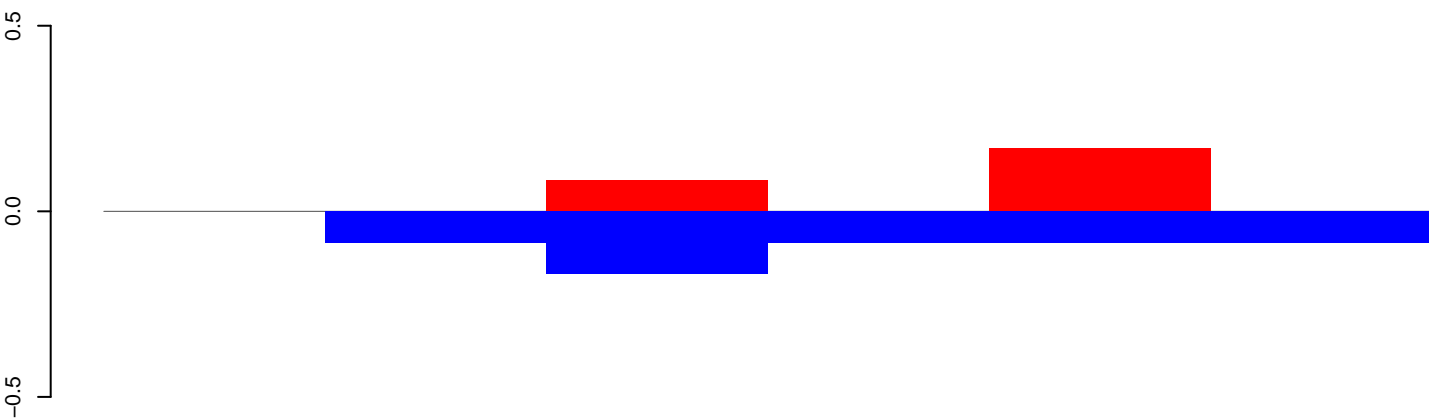
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

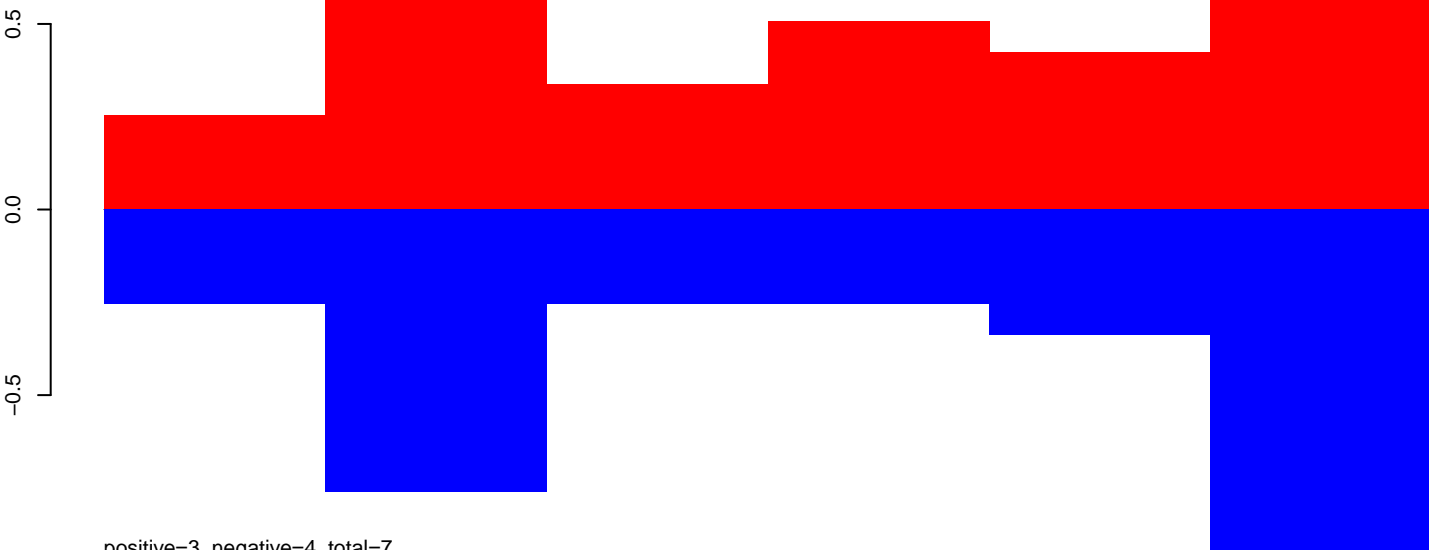


AnGam_Mos55_cells.18_23.rep



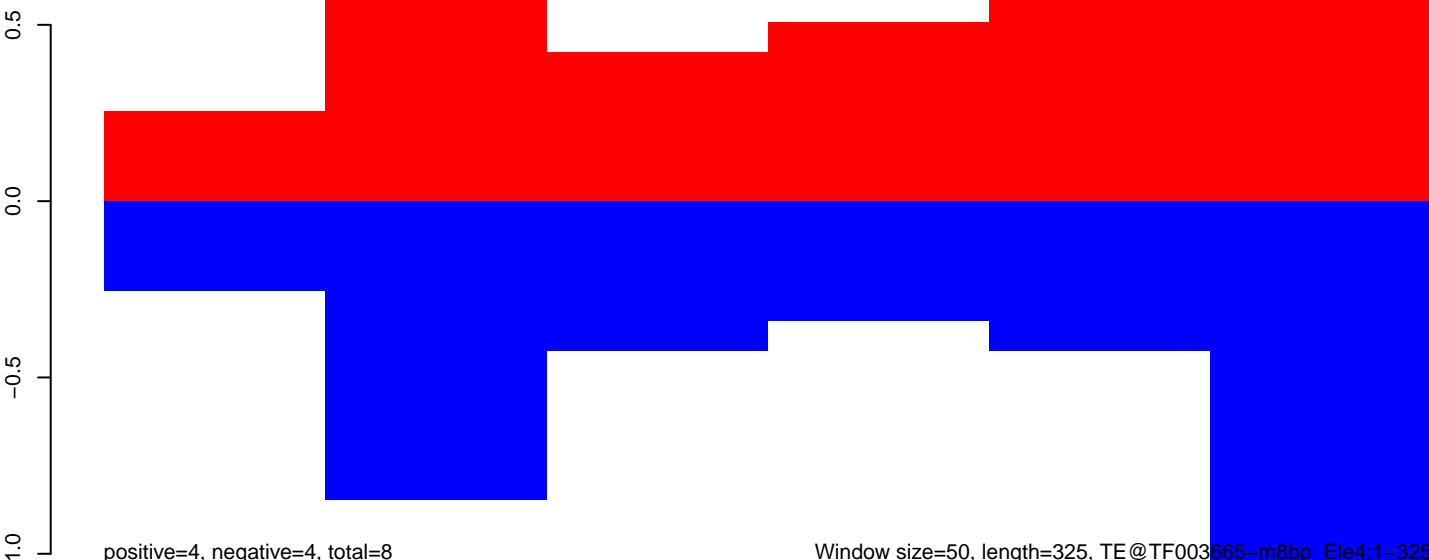
positive=0, negative=1, total=1

AnGam_Mos55_cells.24_35.rep



positive=3, negative=4, total=7

AnGam_Mos55_cells.rep



positive=4, negative=4, total=8

Window size=50, length=325, TE@TF003665-m8bp_Ele4:1-325

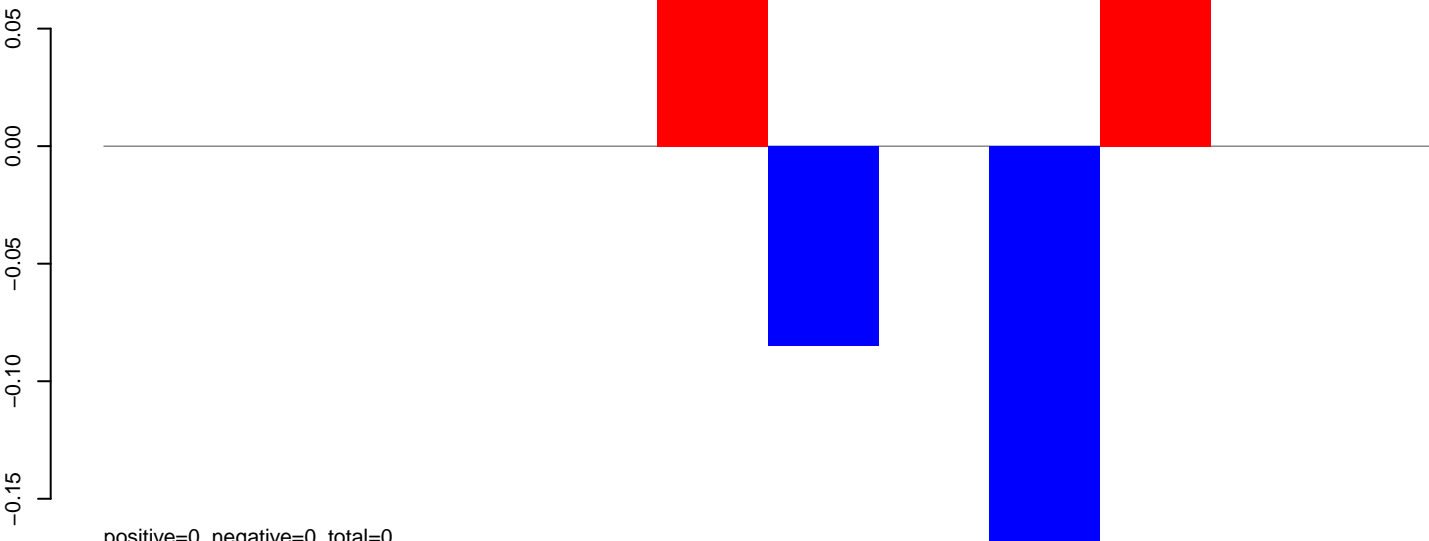
50 100 150 200 250 300 350

AnGam_Mos55_cells.18_23.rep



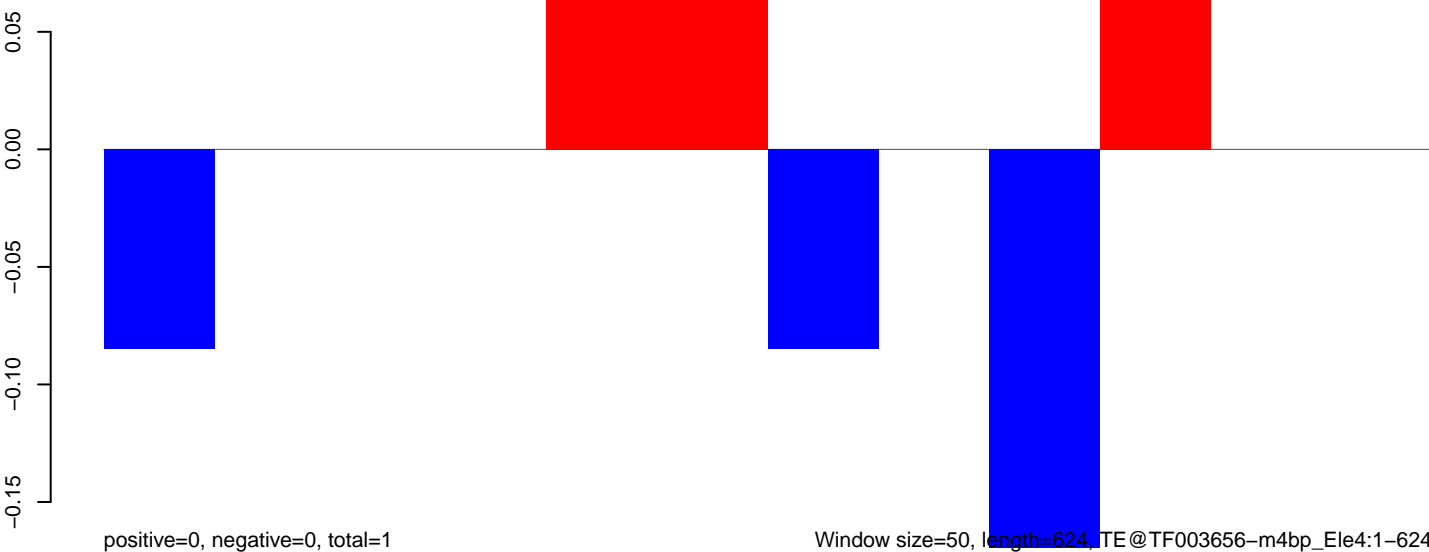
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.rep



positive=0, negative=0, total=1

Window size=50, length=624, TE@TF003656-m4bp_Ele4:1-624

100 200 300 400 500 600

AnGam_Mos55_cells.18_23.rep

0.0
-0.5
-1.0
-1.5

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.0
-0.5
-1.0
-1.5

positive=0, negative=2, total=3

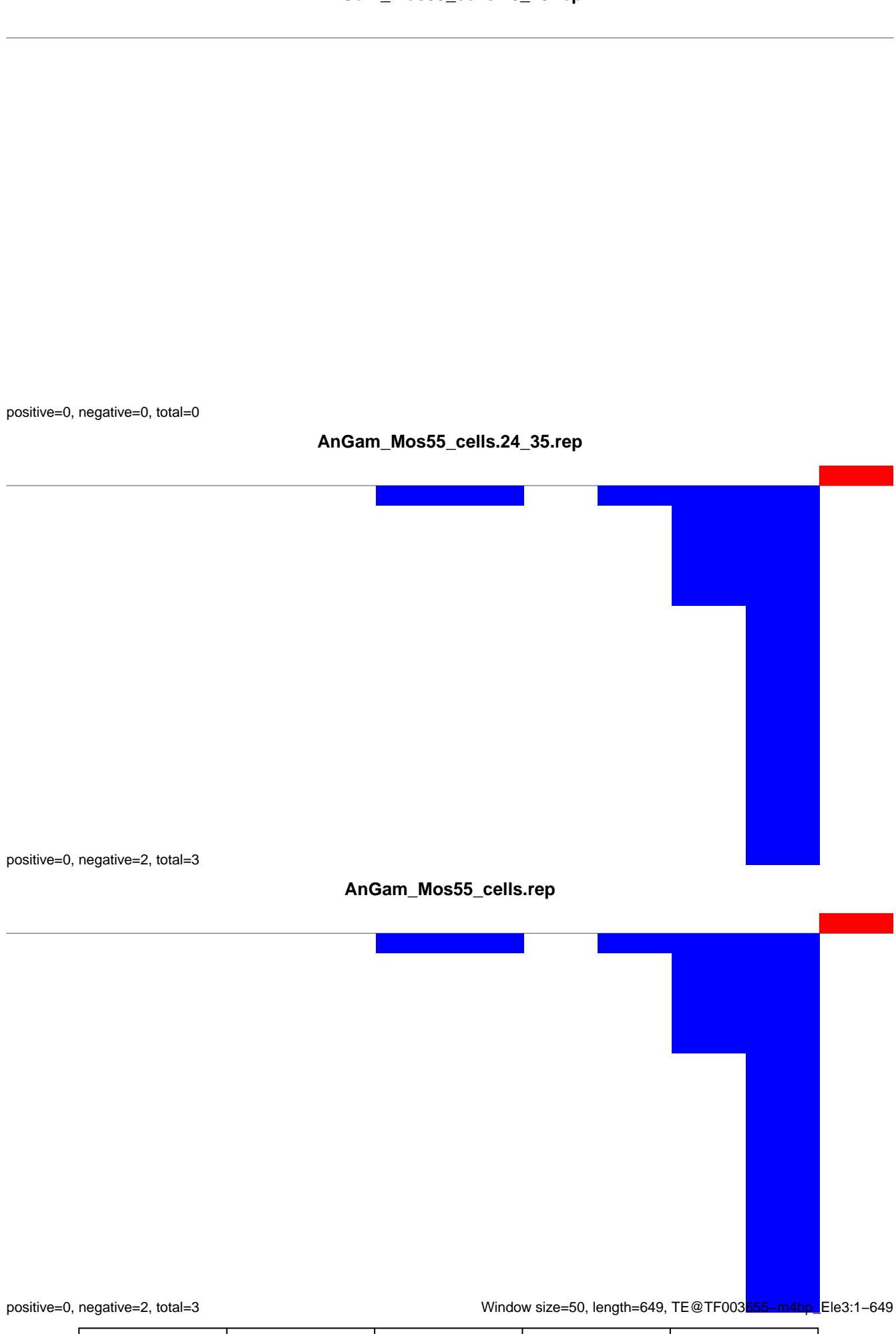
AnGam_Mos55_cells.rep

0.0
-0.5
-1.0
-1.5

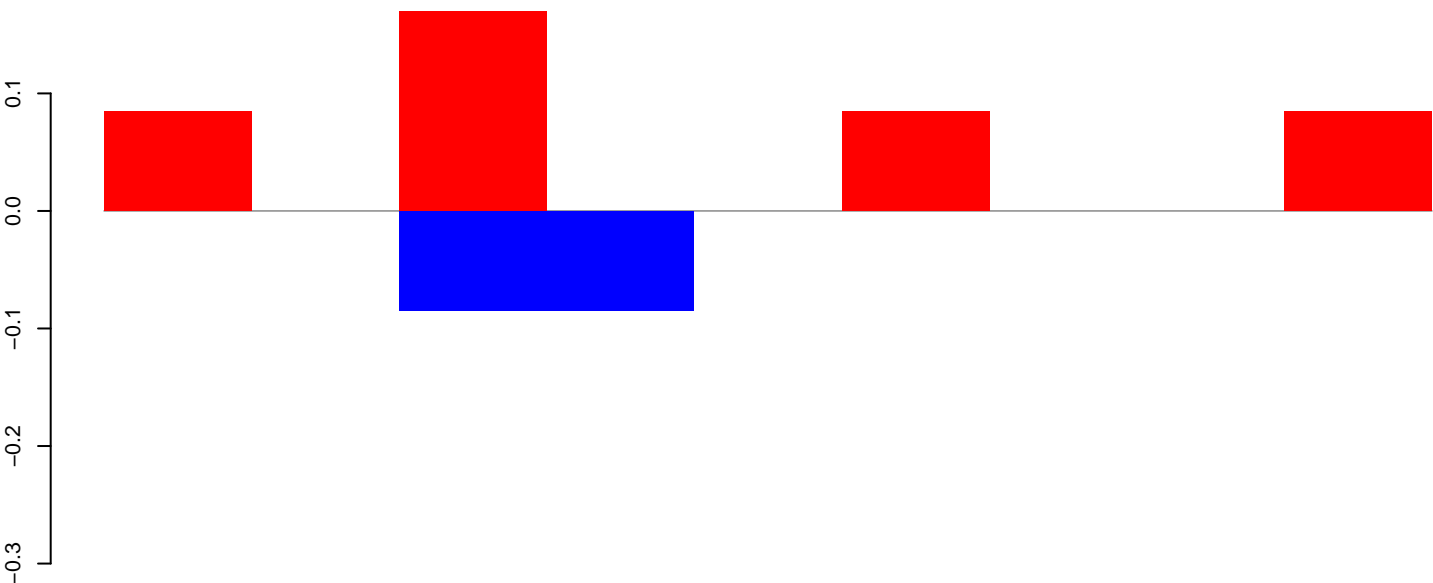
positive=0, negative=2, total=3

Window size=50, length=649, TE@TF003655-m4bp, Ele3:1-649

100 200 300 400 500 600

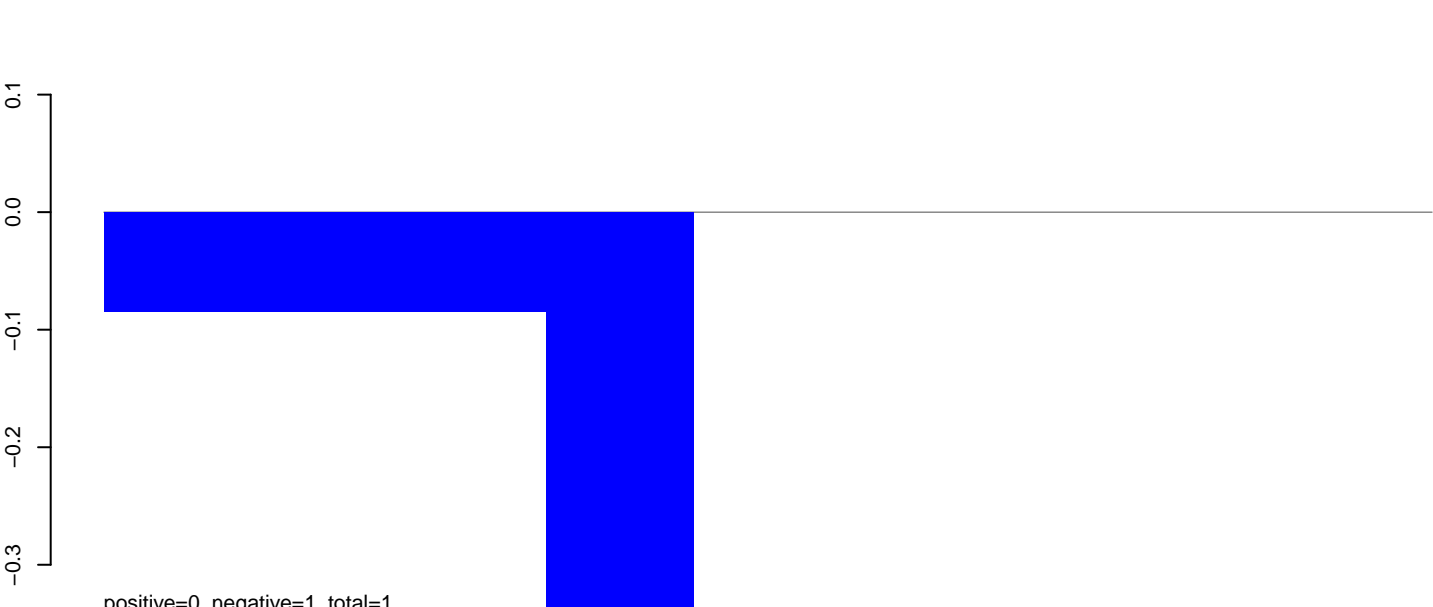


AnGam_Mos55_cells.18_23.rep



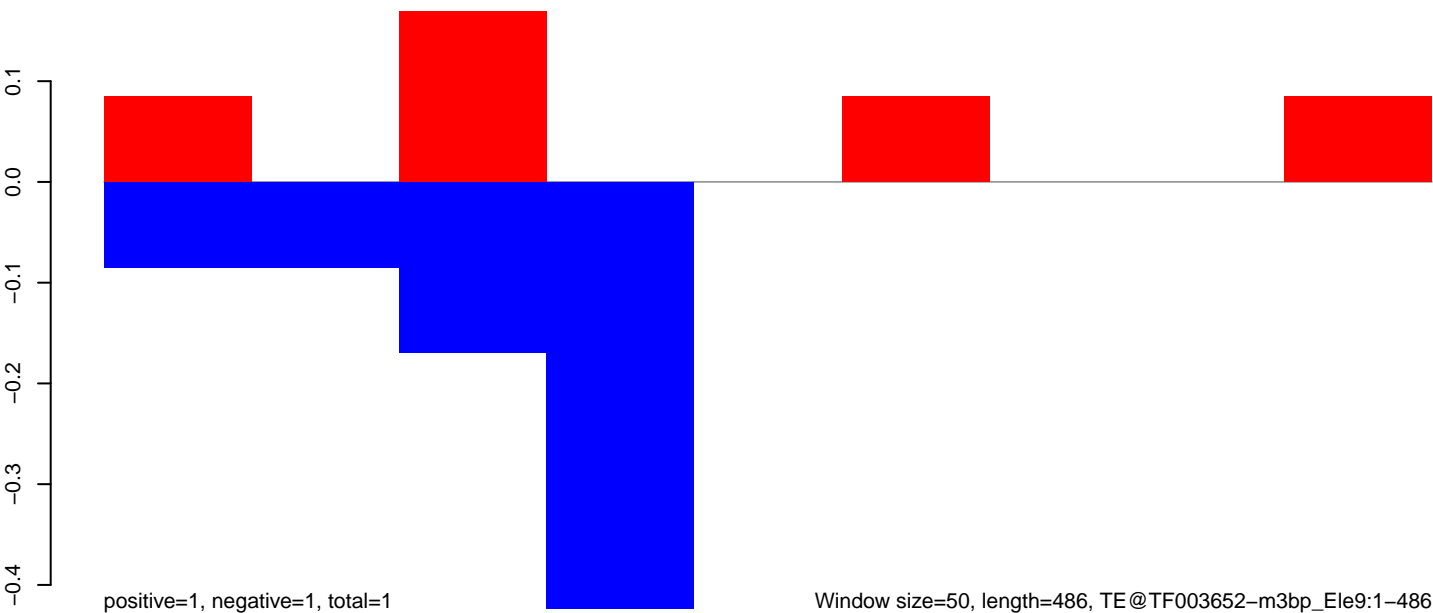
positive=1, negative=0, total=1

AnGam_Mos55_cells.24_35.rep



positive=0, negative=1, total=1

AnGam_Mos55_cells.rep



positive=1, negative=1, total=1

Window size=50, length=486, TE@TF003652-m3bp_Ele9:1-486

100

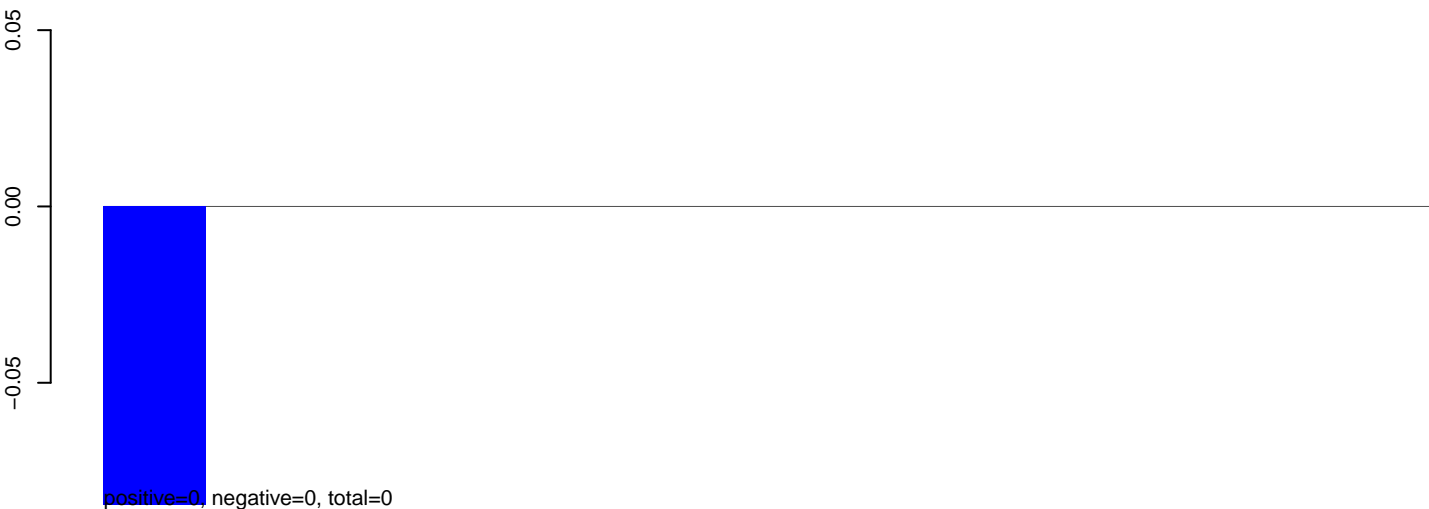
200

300

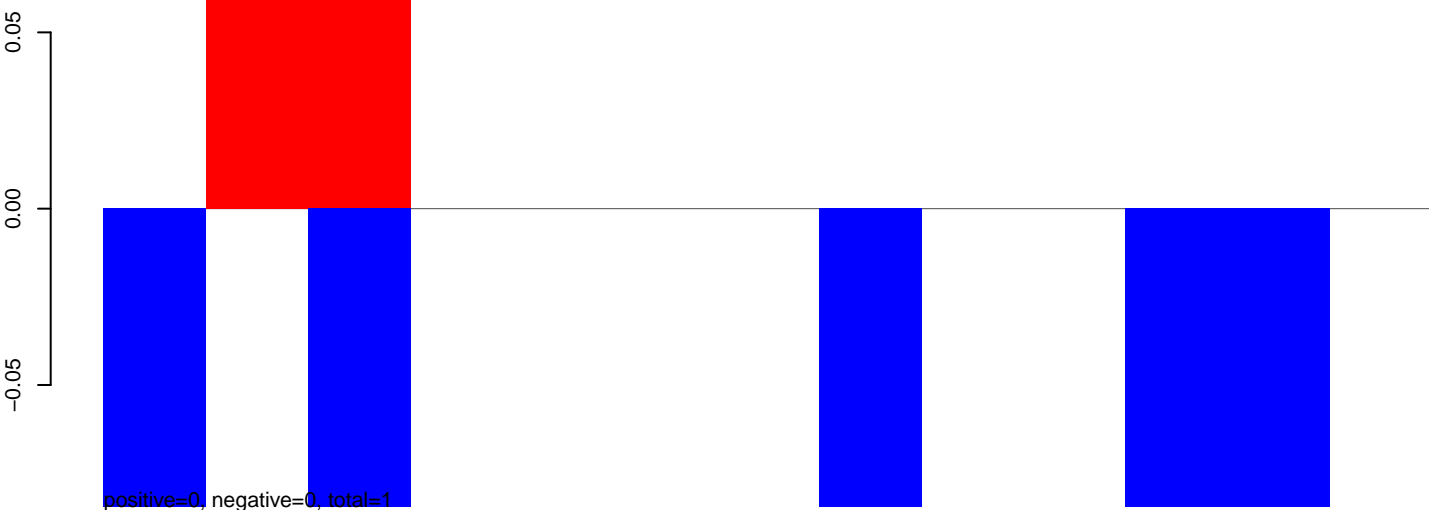
400

500

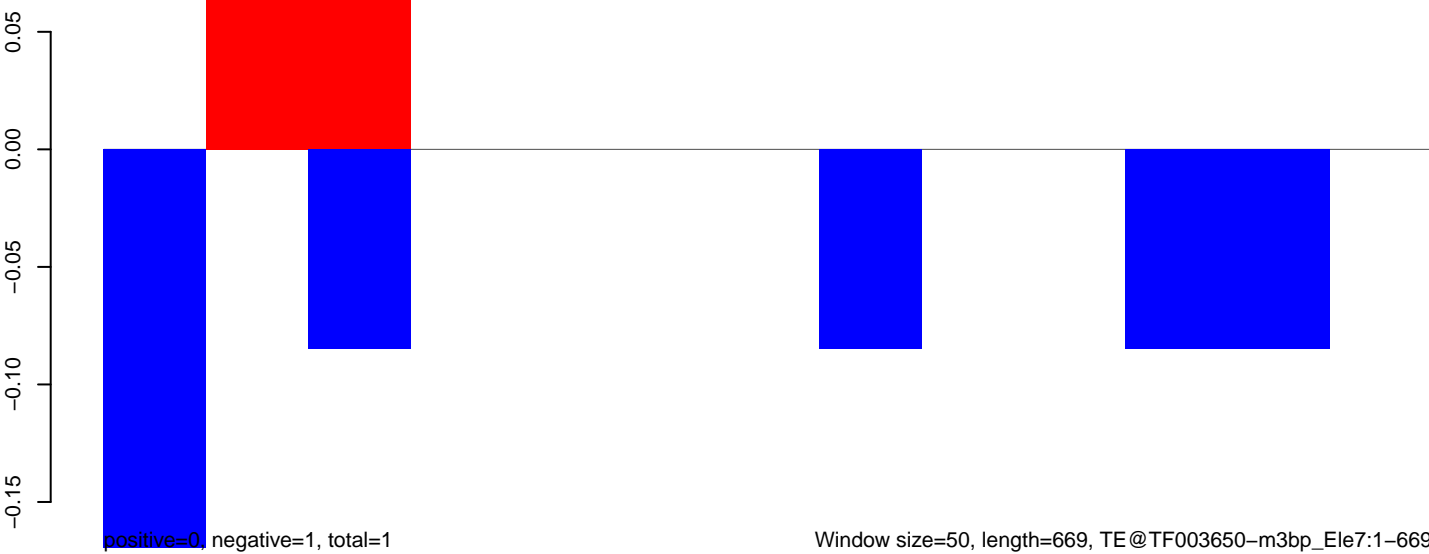
AnGam_Mos55_cells.18_23.rep



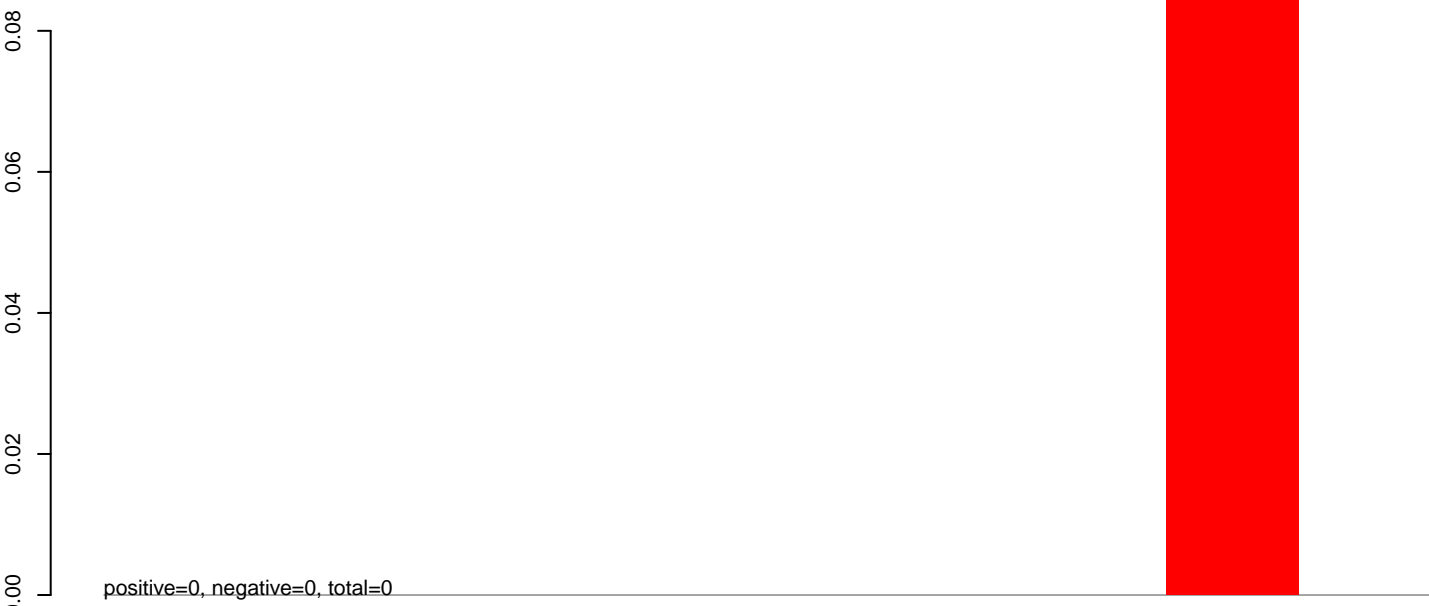
AnGam_Mos55_cells.24_35.rep



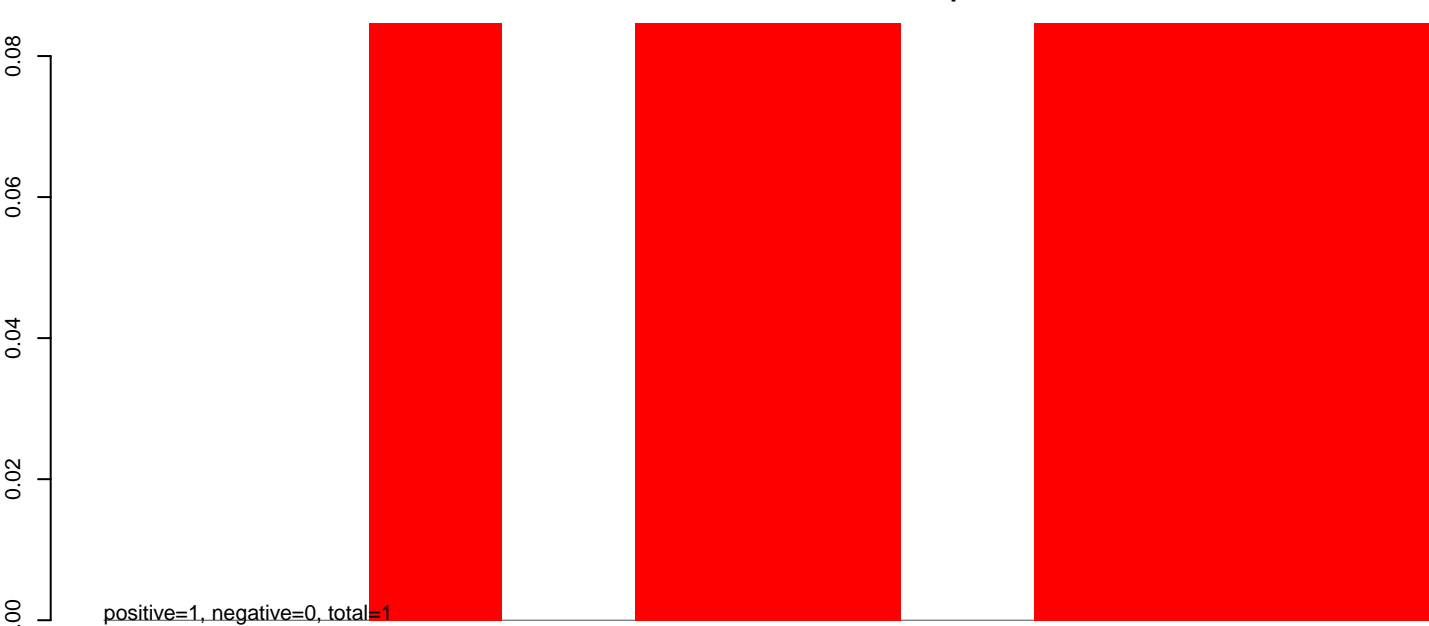
AnGam_Mos55_cells.rep



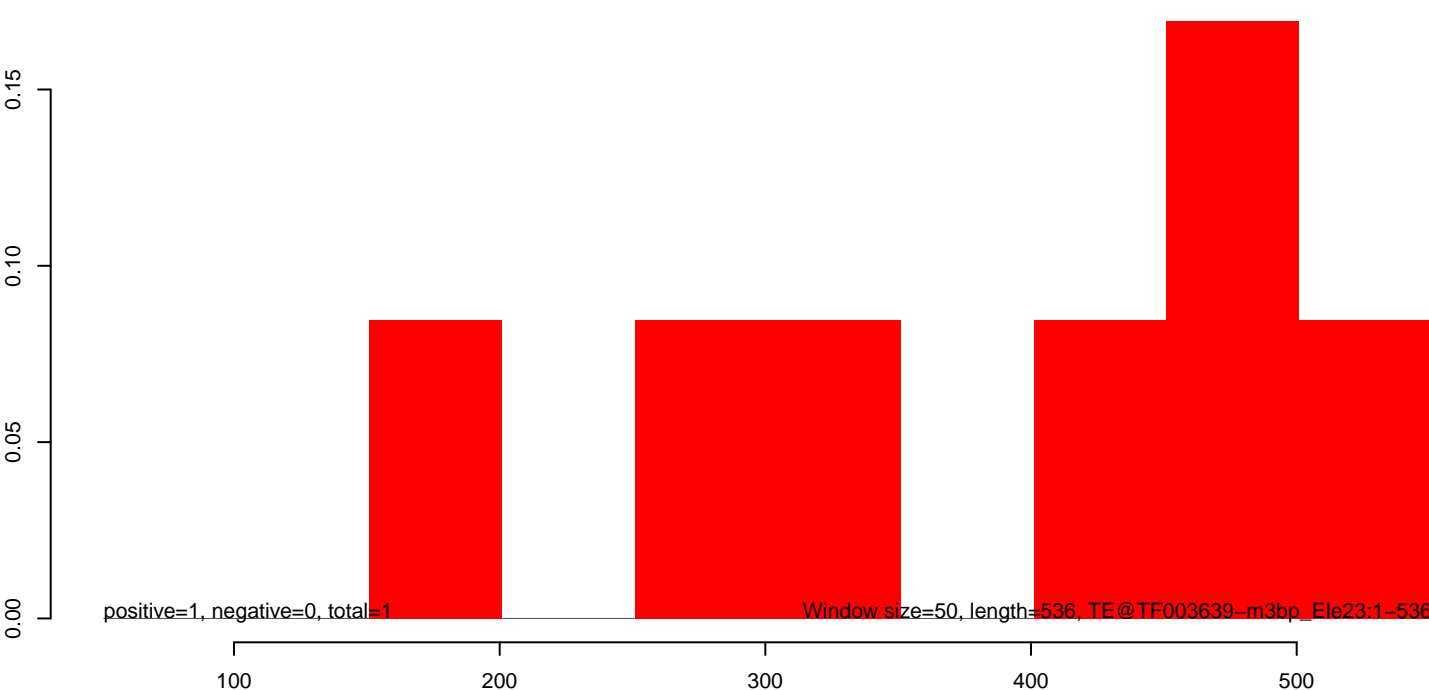
AnGam_Mos55_cells.18_23.rep



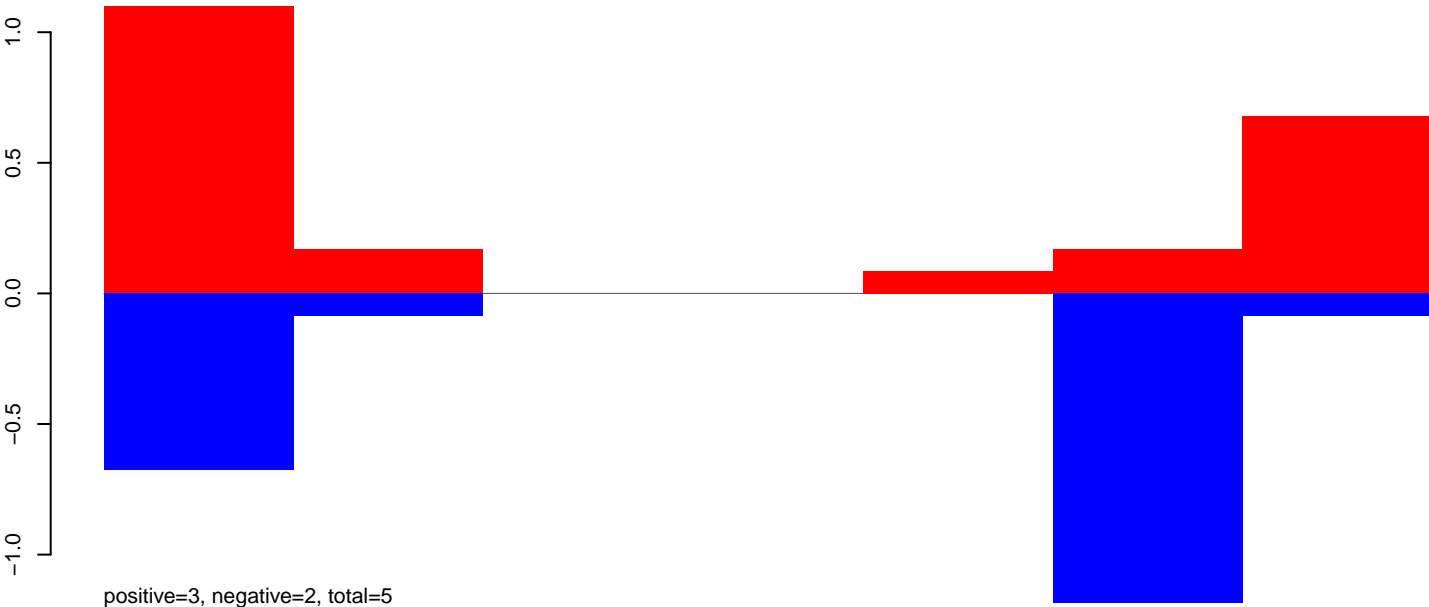
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



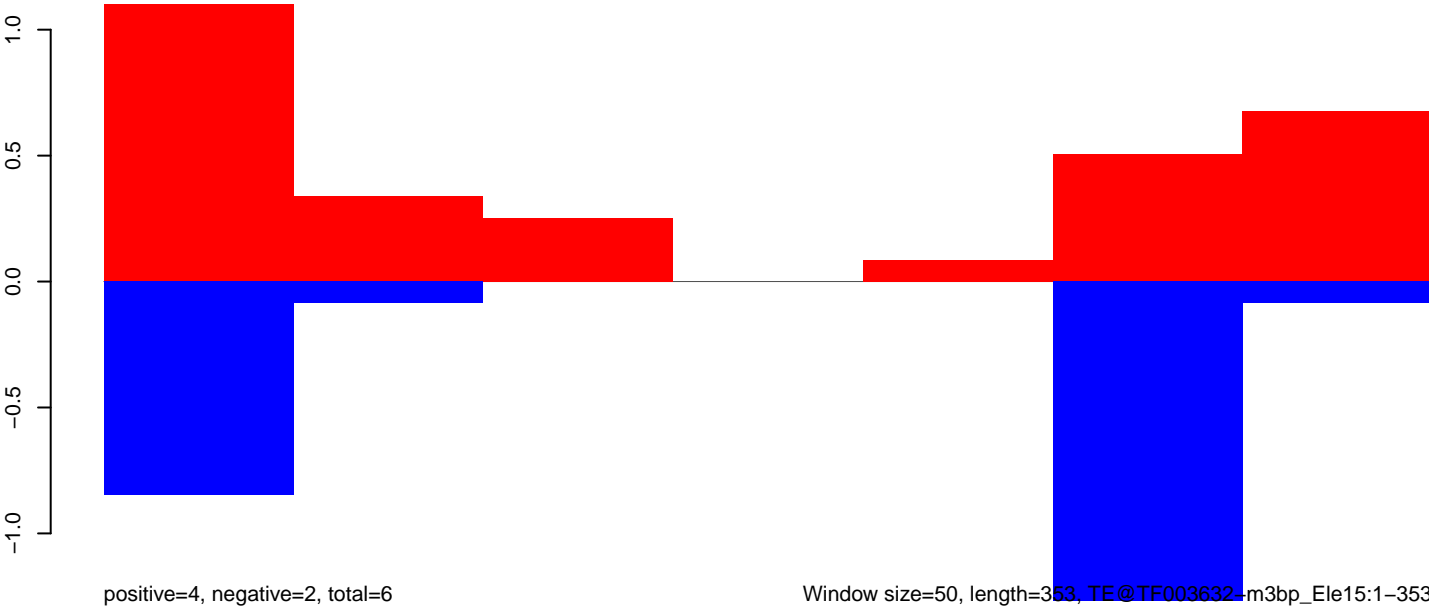
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



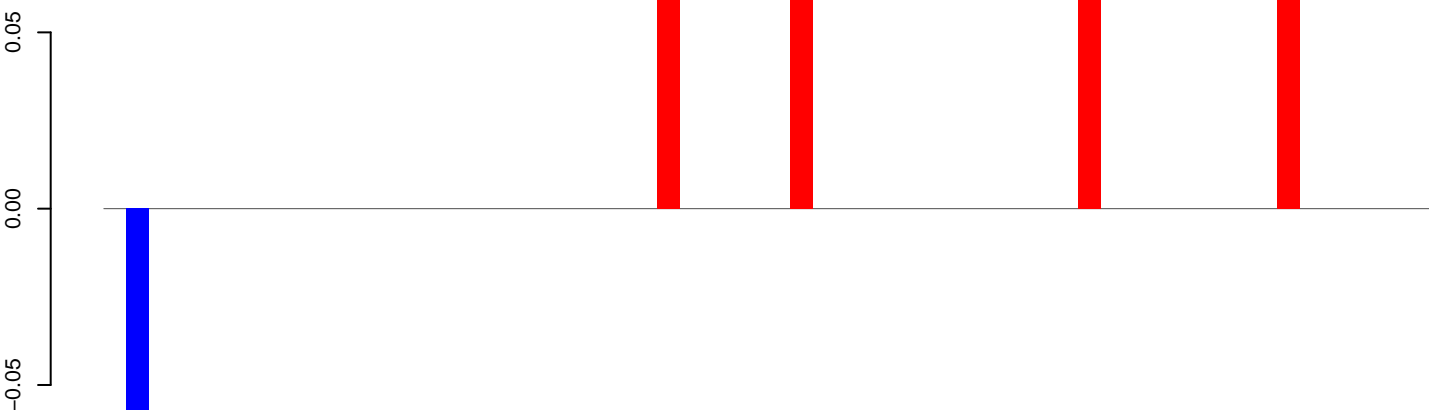
Window size=50, length=353, TE@TF003632-m3bp_Ele15:1-353

AnGam_Mos55_cells.18_23.rep



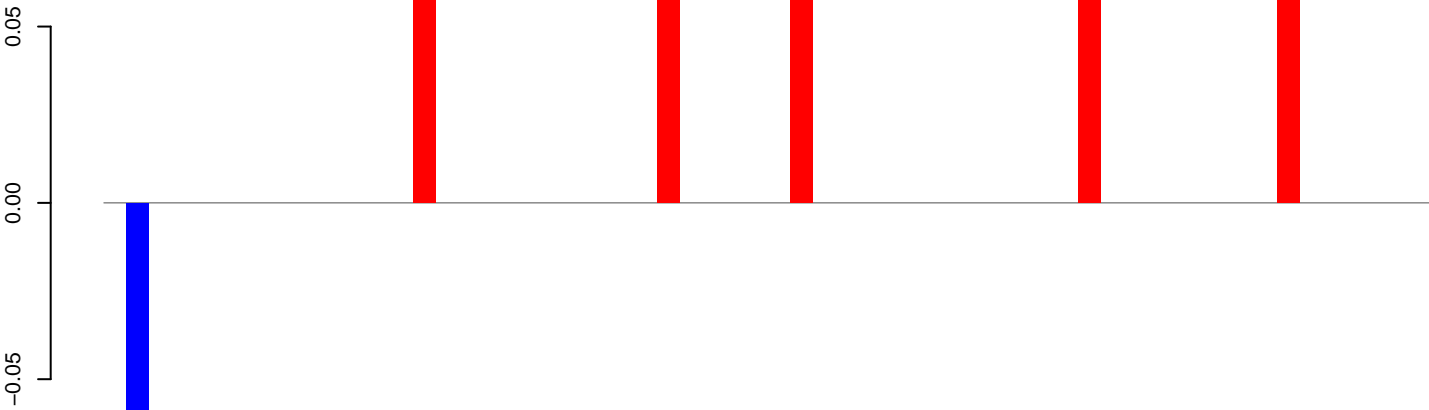
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

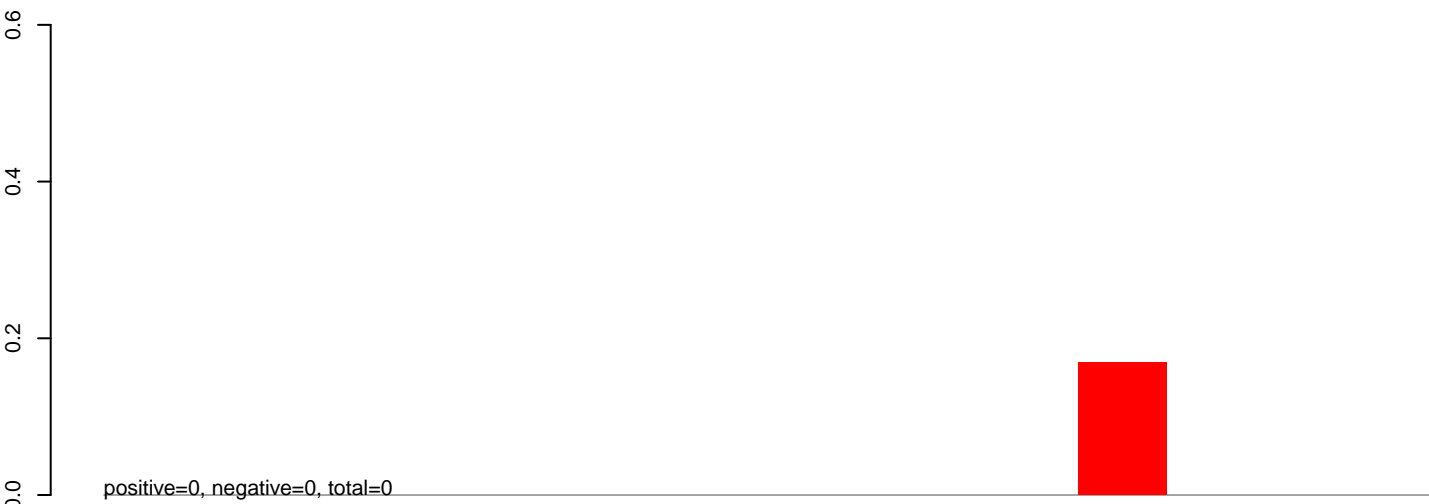


positive=0, negative=0, total=1

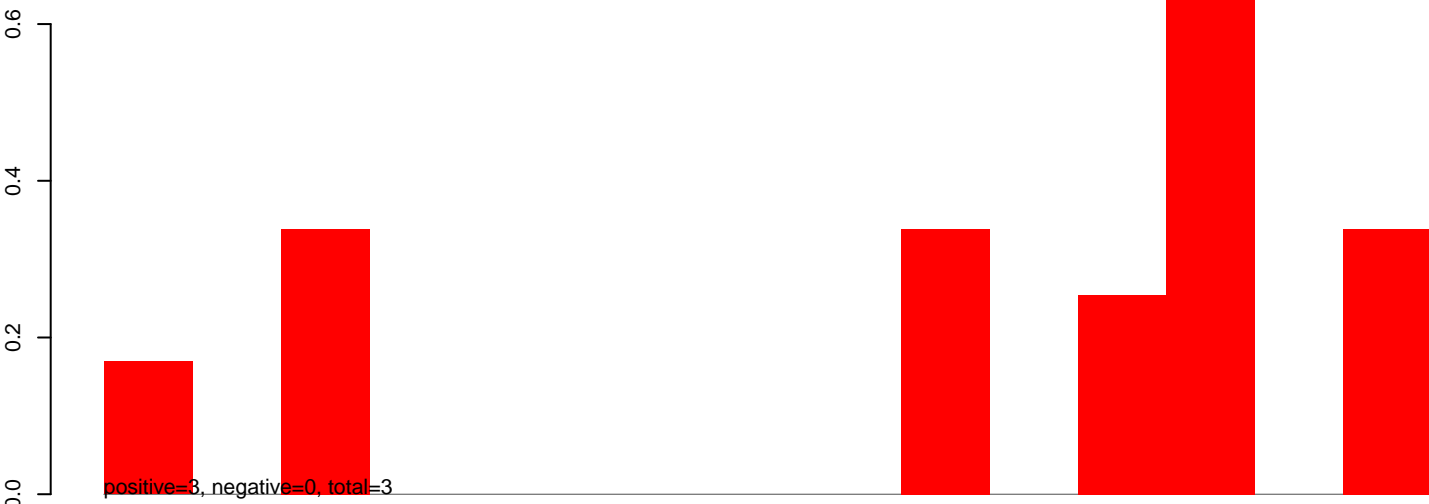
Window size=50, length=3010, TE@TF003522-PIF_Ele2:1-3010

0 500 1000 1500 2000 2500 3000

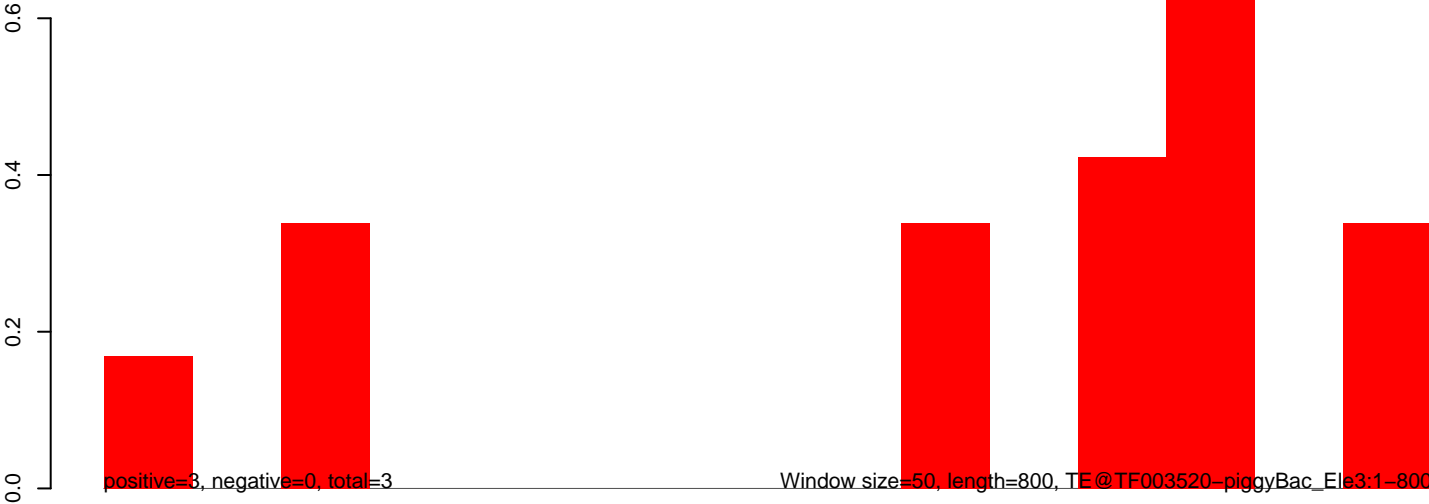
AnGam_Mos55_cells.18_23.rep



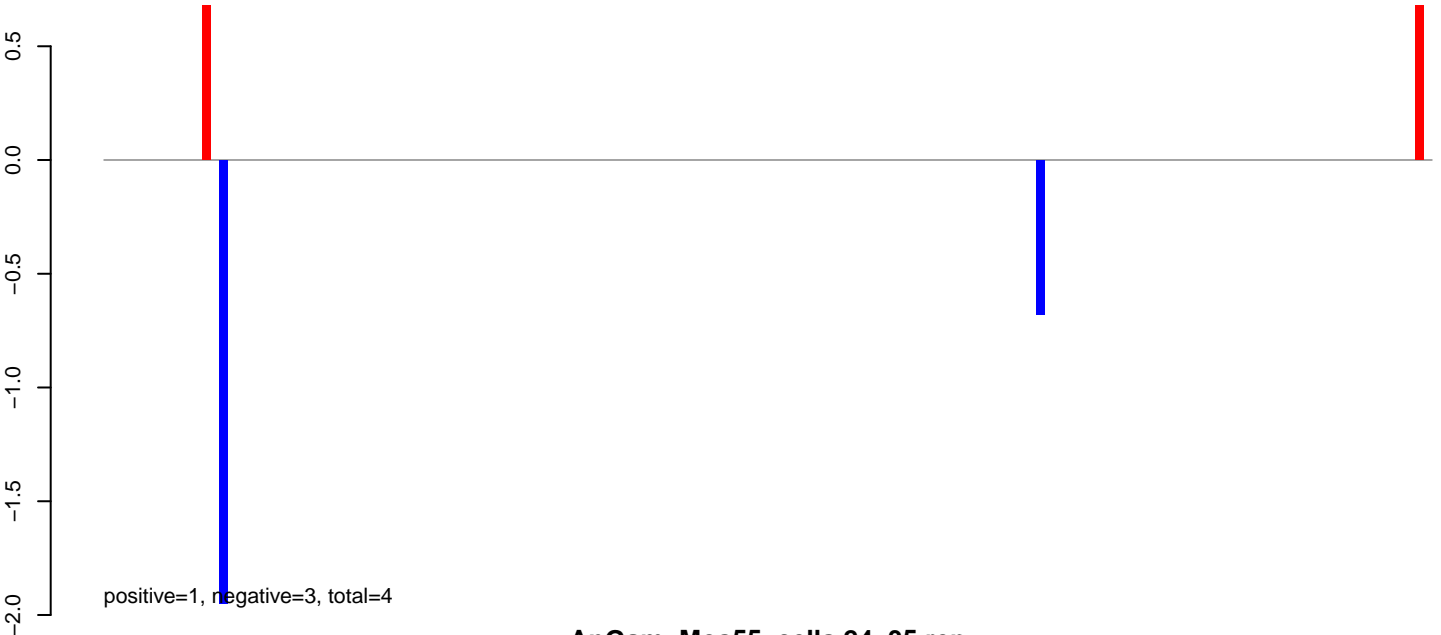
AnGam_Mos55_cells.24_35.rep



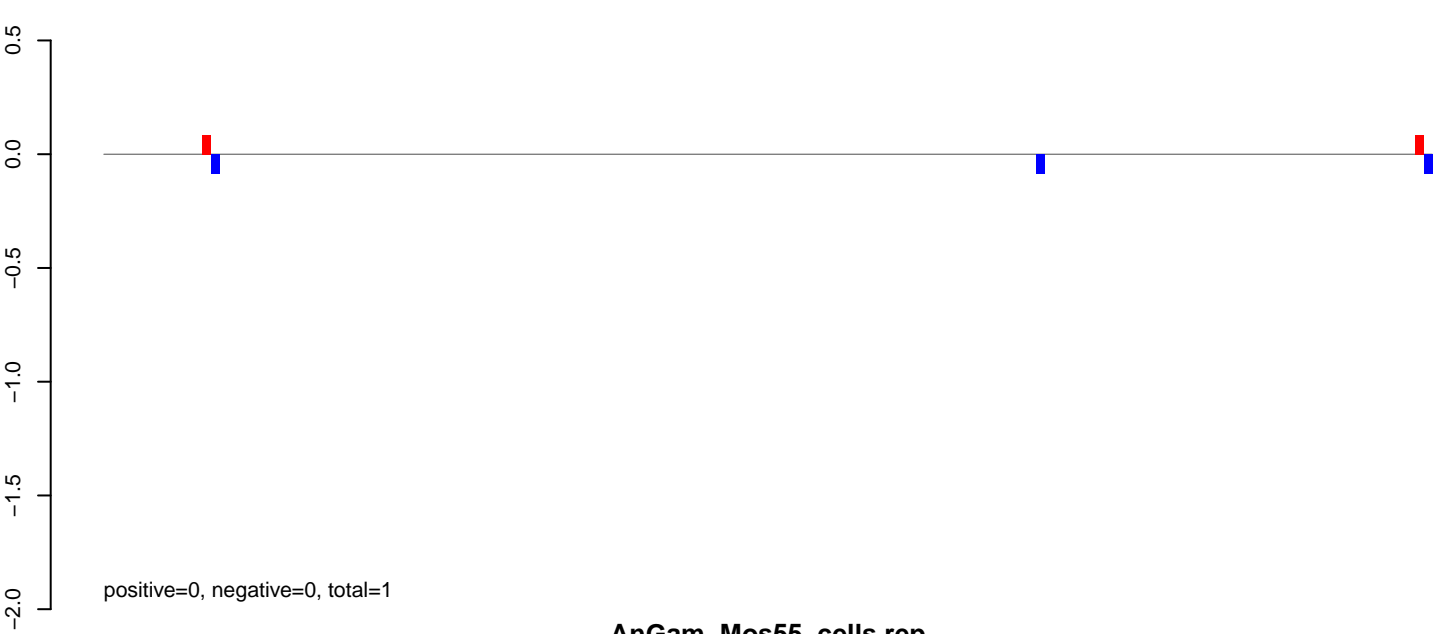
AnGam_Mos55_cells.rep



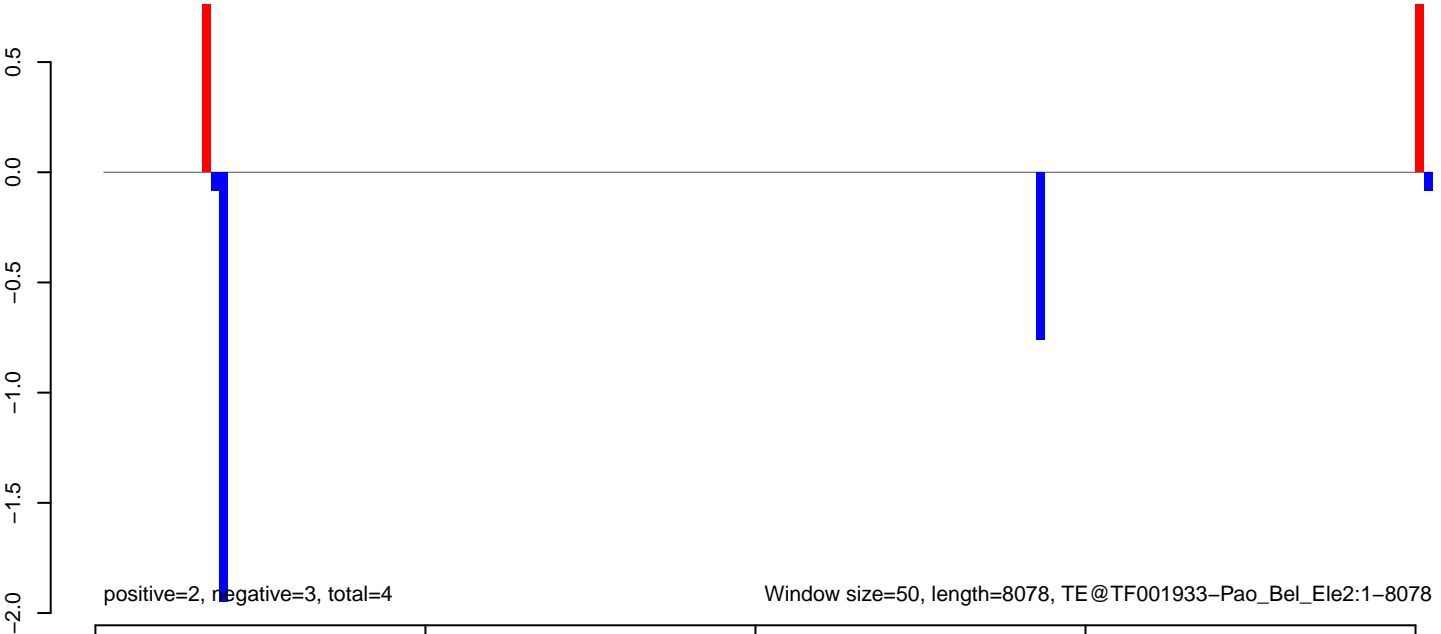
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

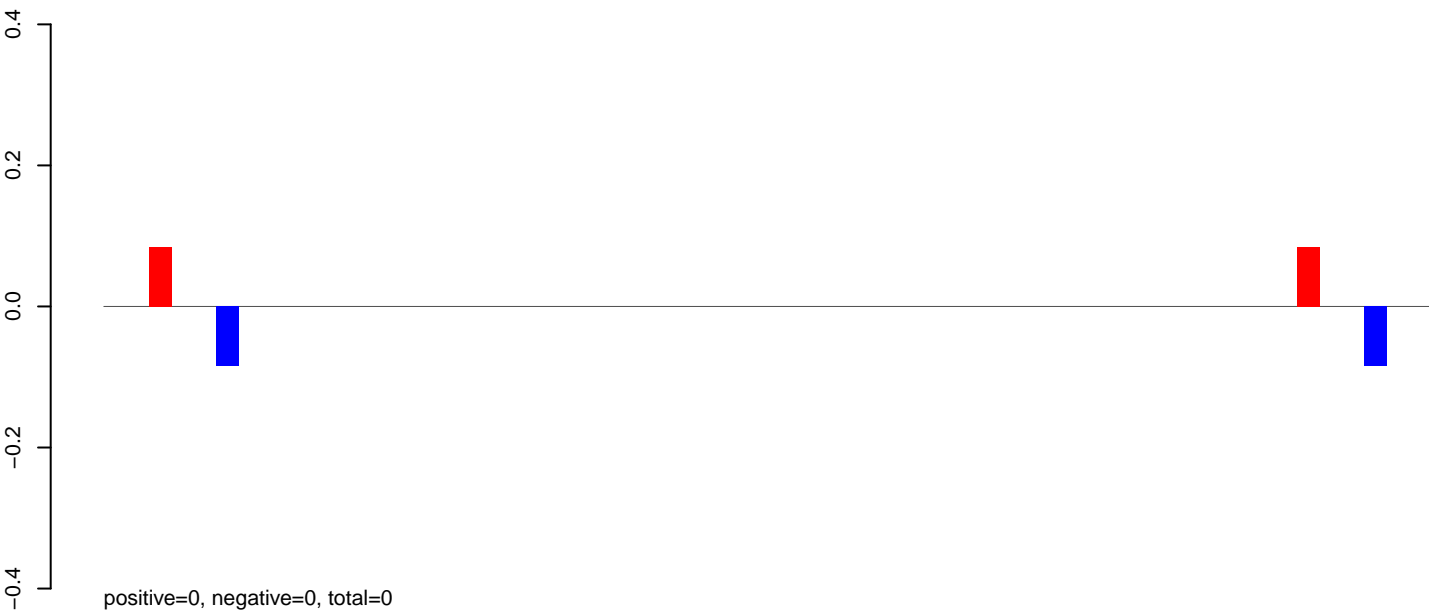


AnGam_Mos55_cells.rep

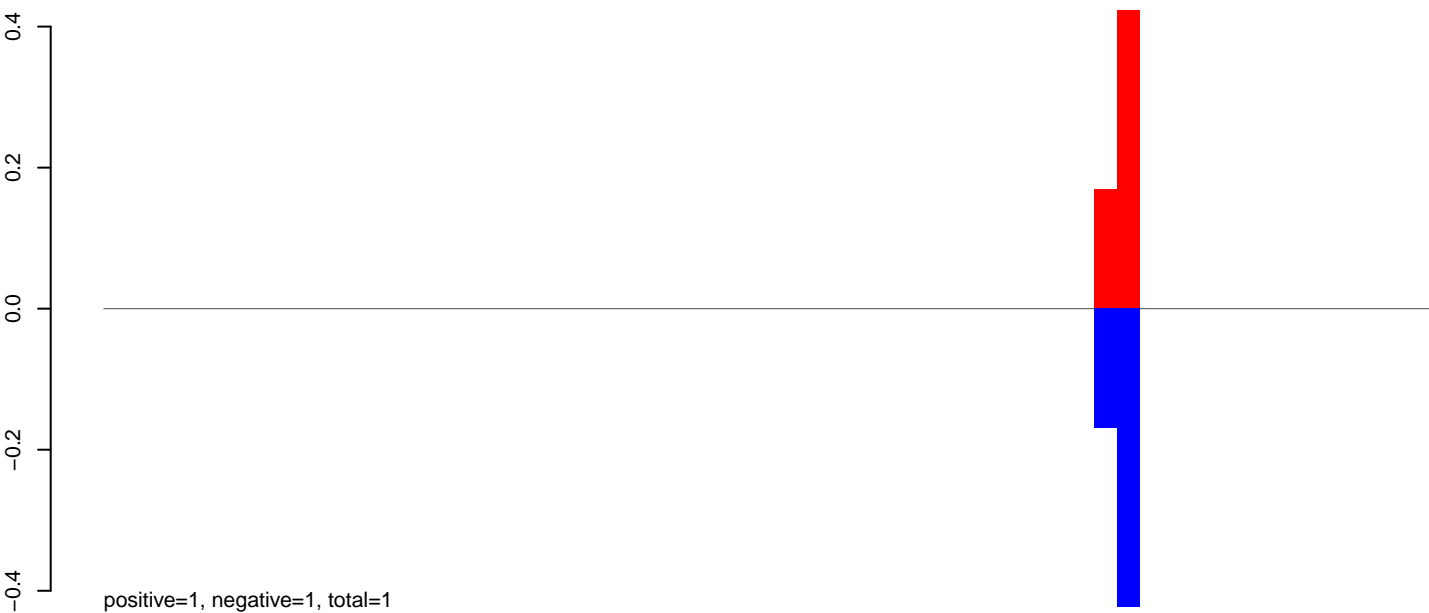


Window size=50, length=8078, TE@TF001933-Pao_Bel_Ele2:1-8078

AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



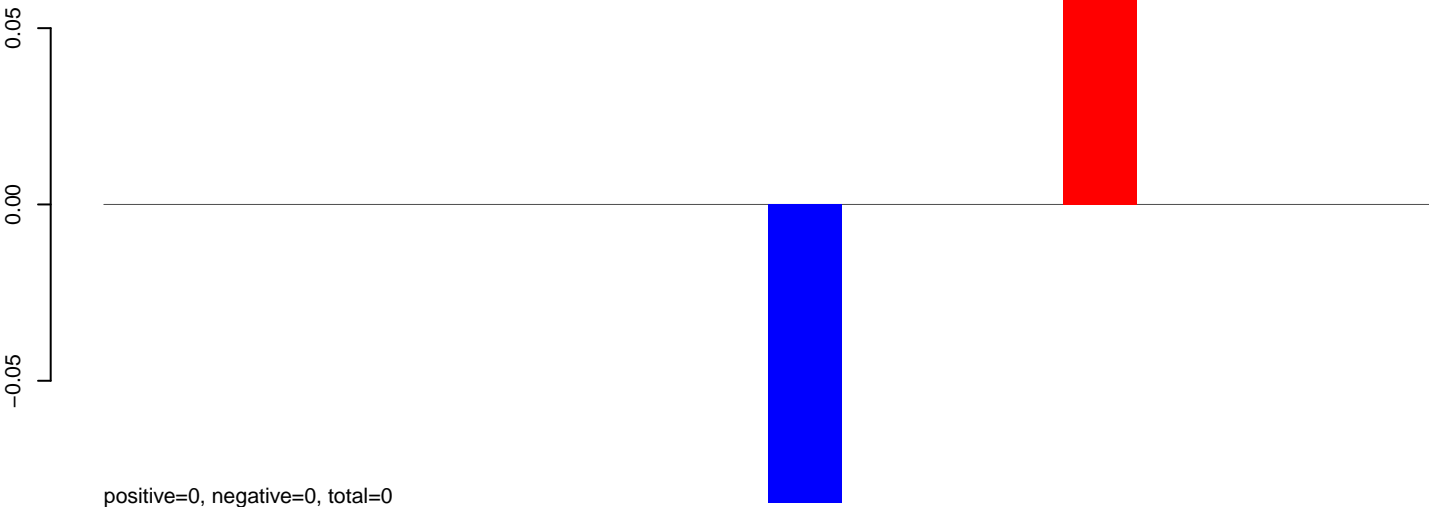
AnGam_Mos55_cells.rep



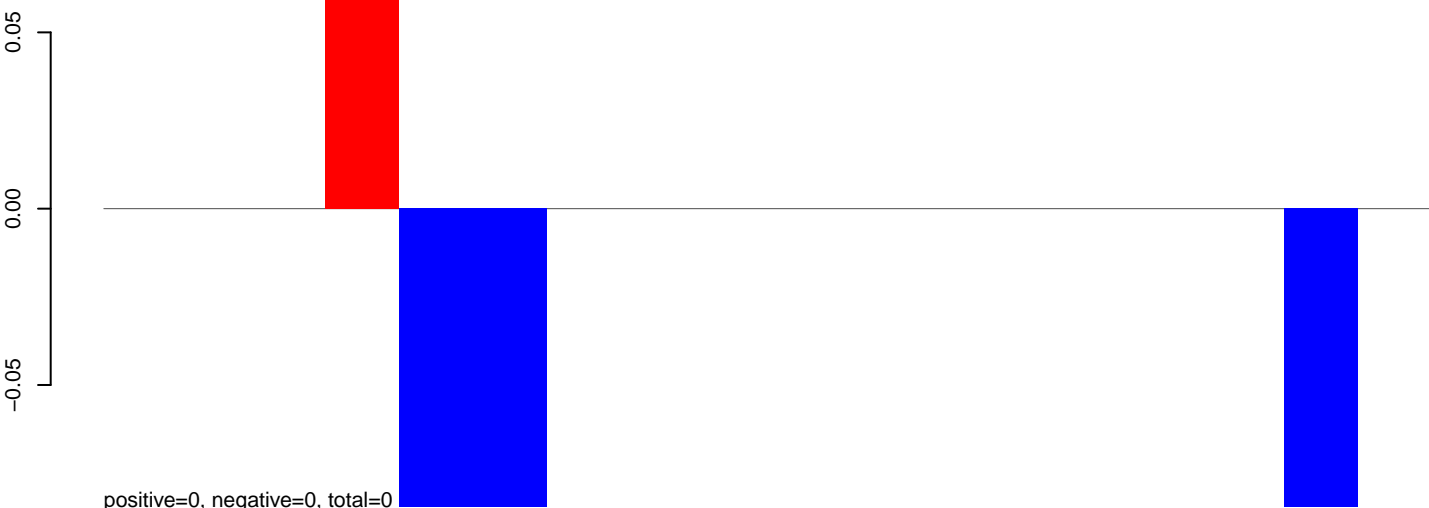
Window size=50, length=2989, TE@TF001233-gambol_Ele11-Piper:1-2989

0 500 1000 1500 2000 2500 3000

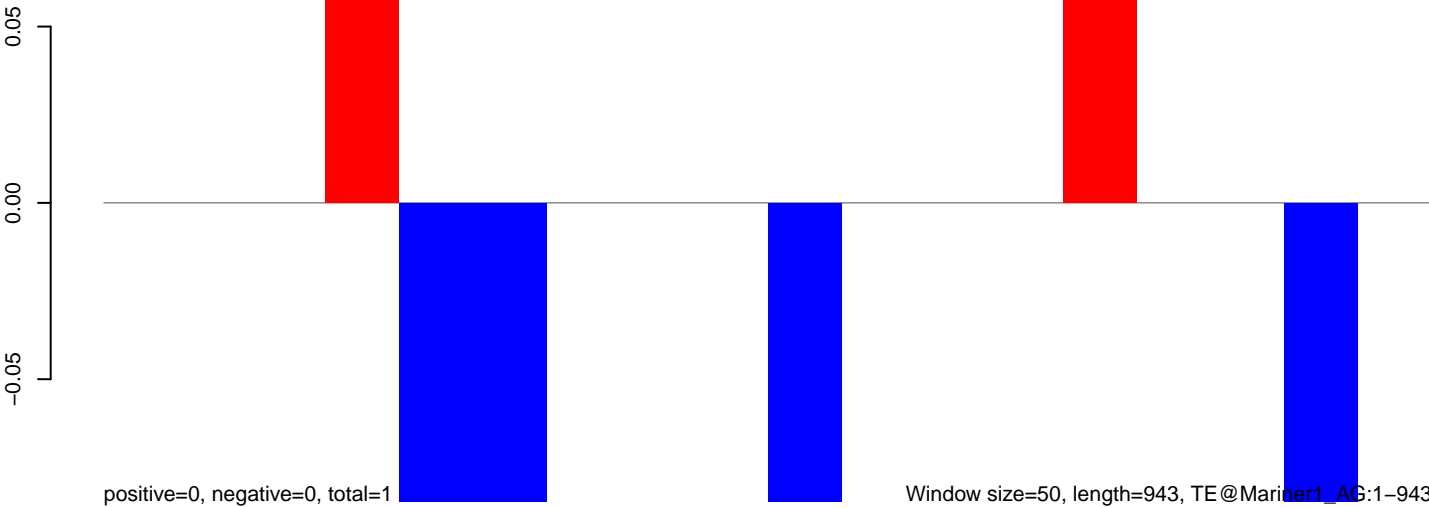
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



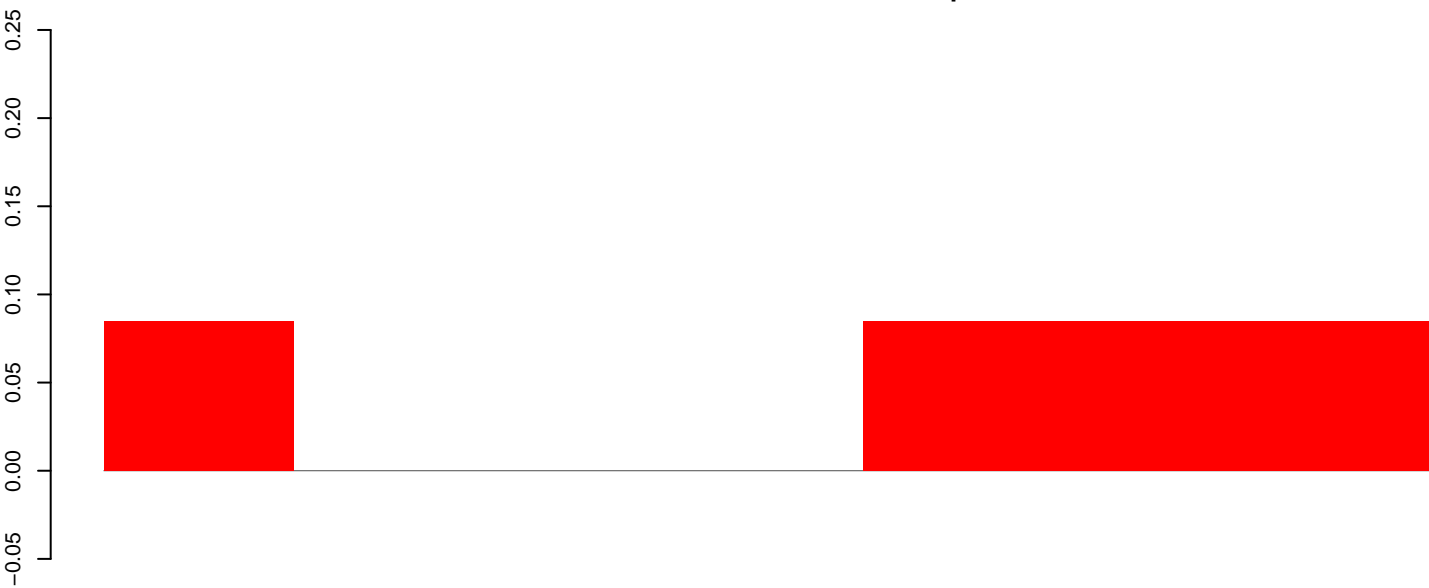
200

400

600

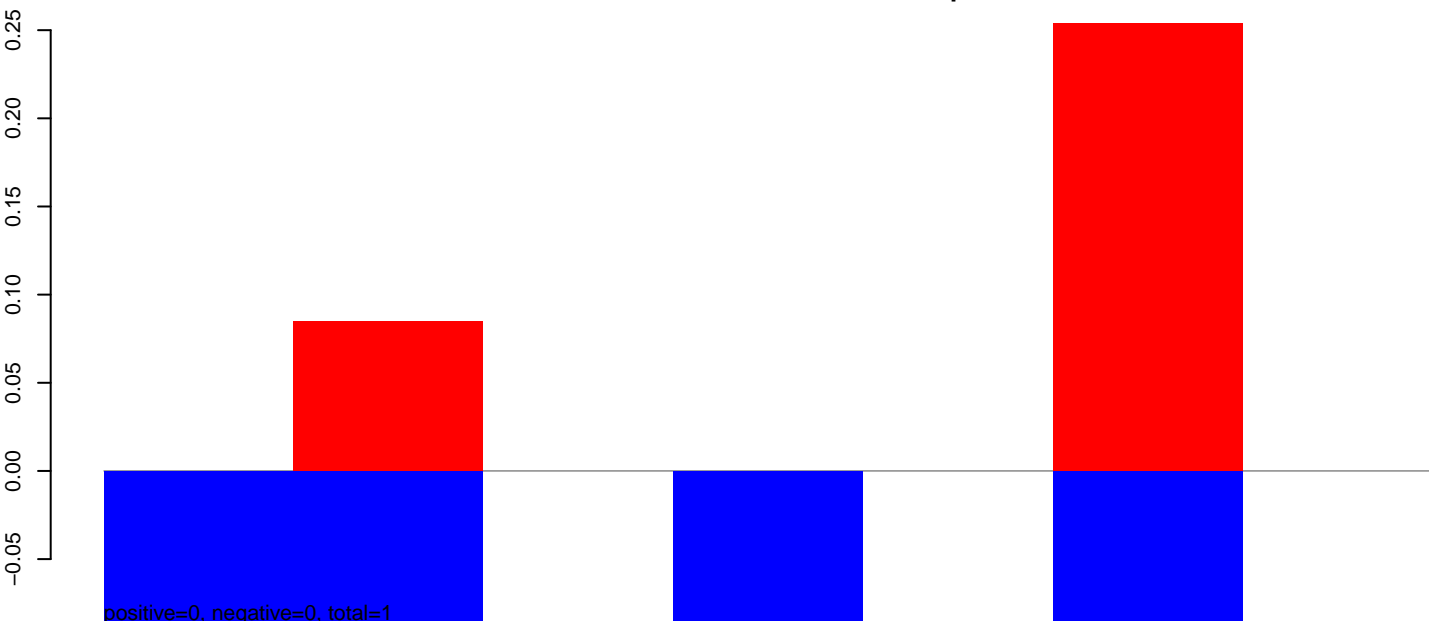
800

AnGam_Mos55_cells.18_23.rep



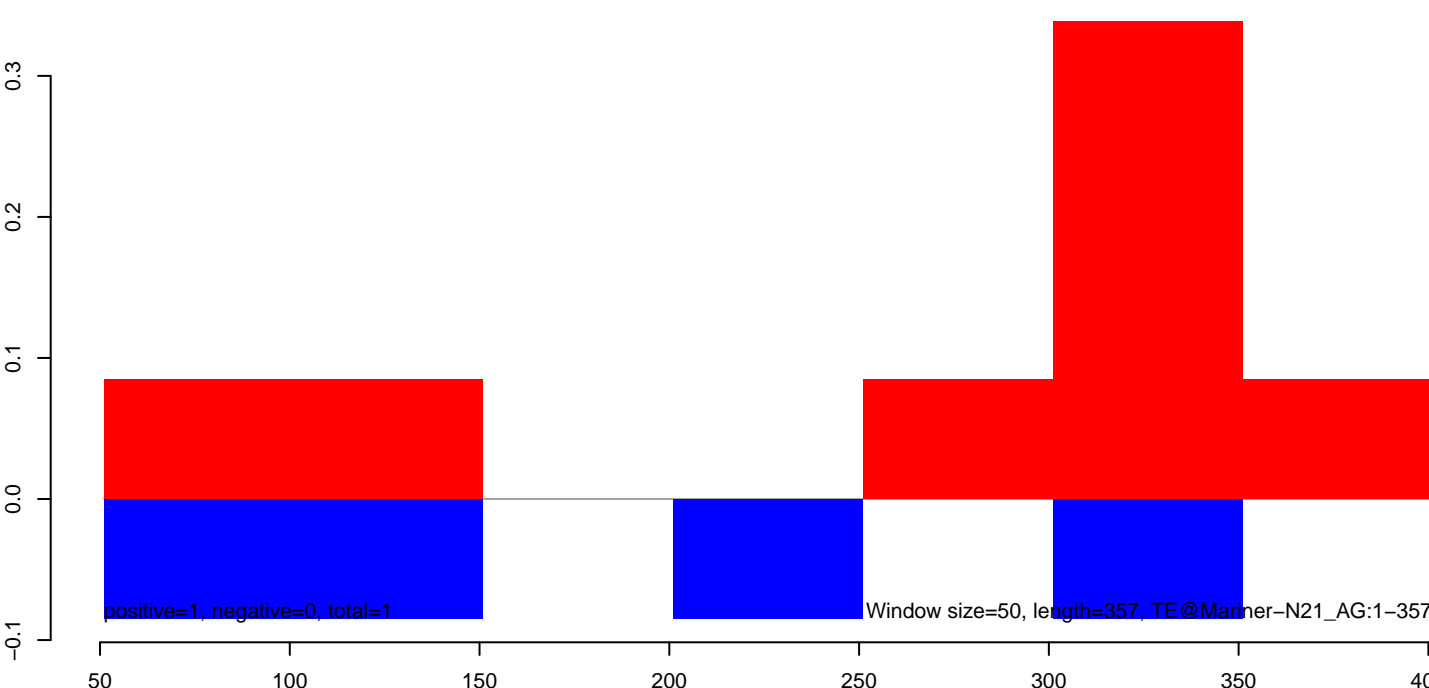
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=1

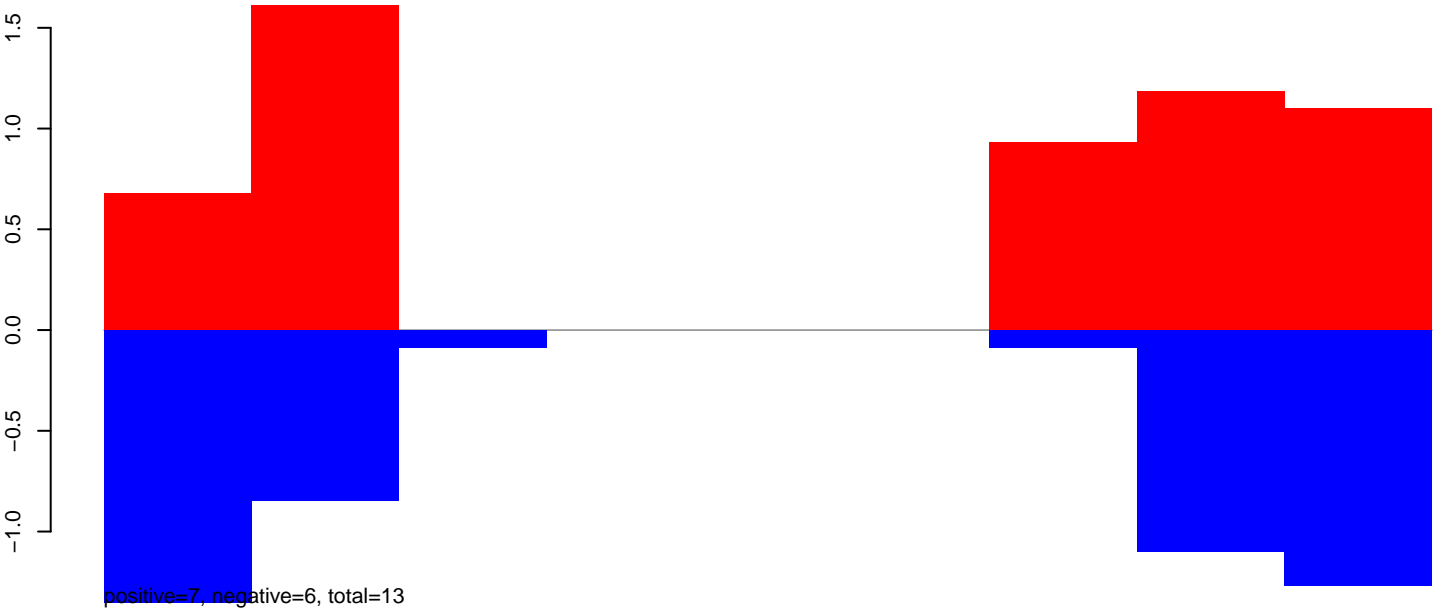
AnGam_Mos55_cells.rep



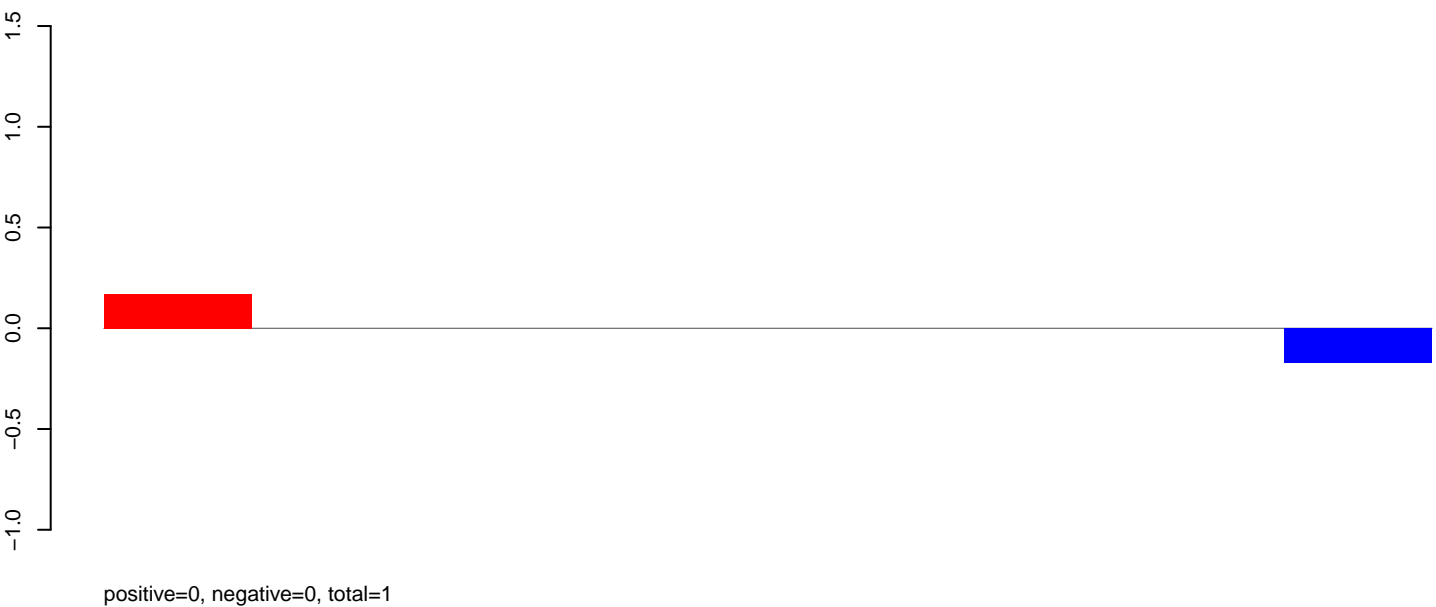
positive=1, negative=0, total=1

Window size=50, length=357, TE@Manner-N21_AG:1-357

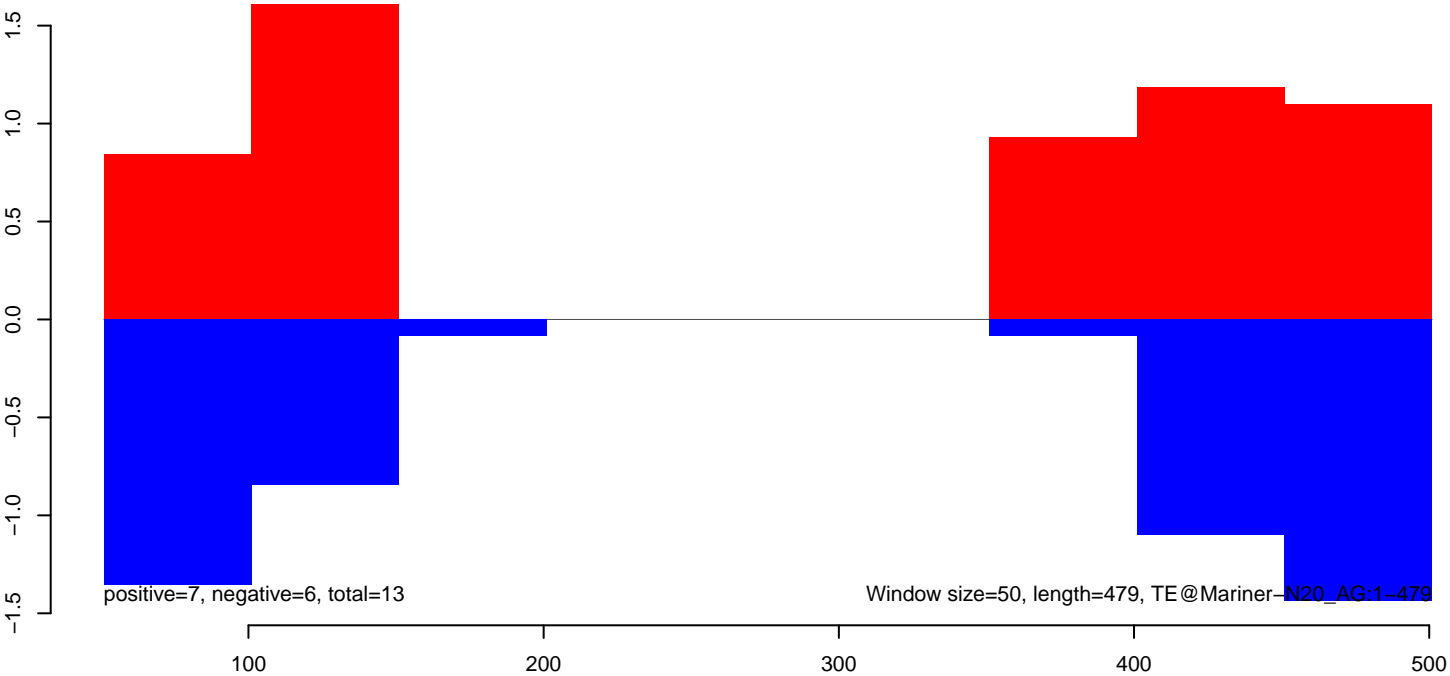
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



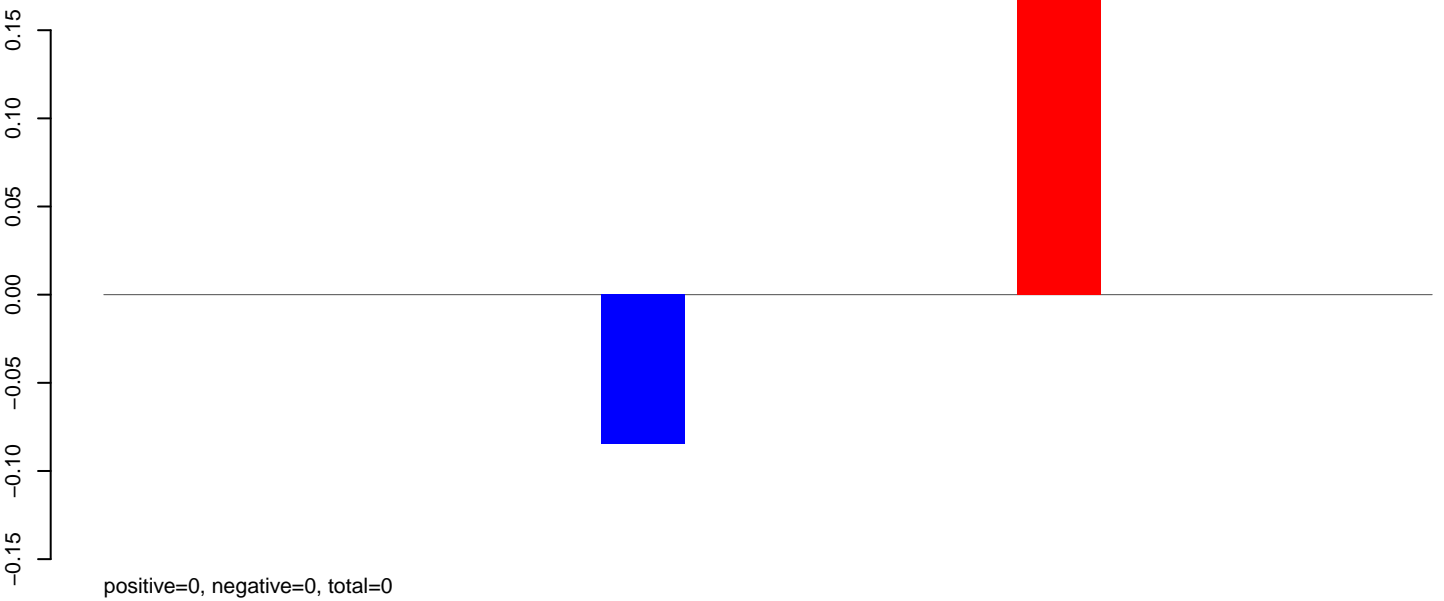
AnGam_Mos55_cells.rep



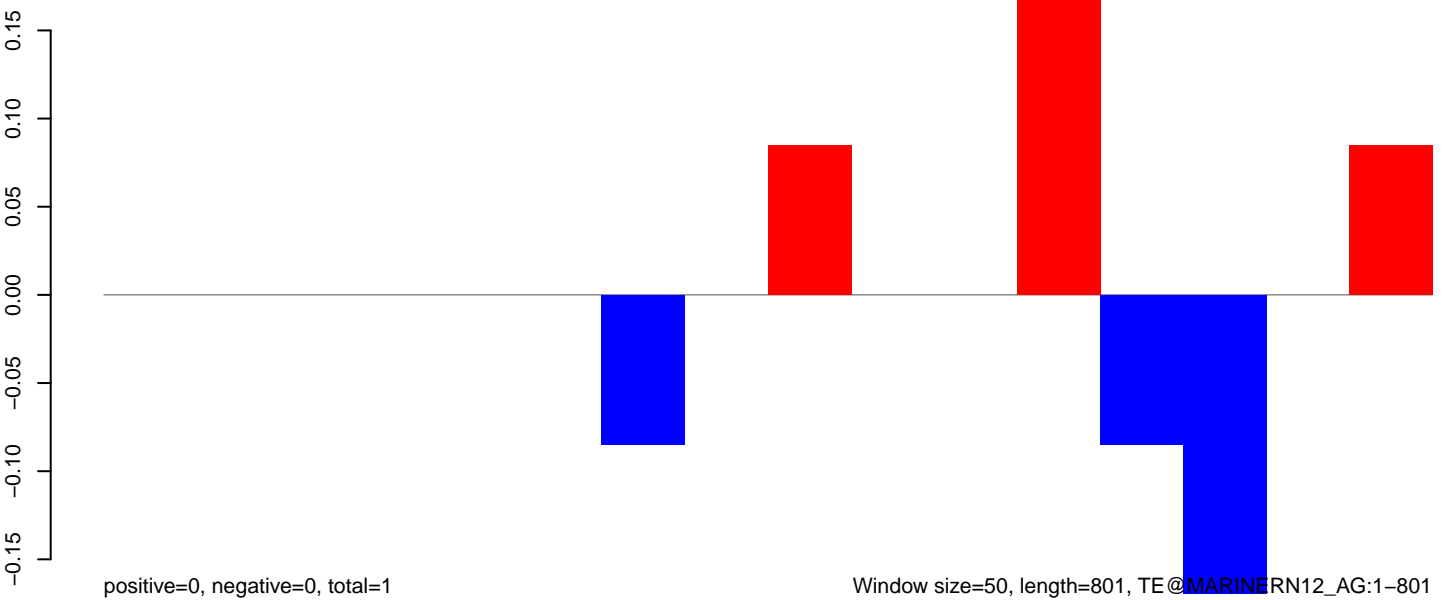
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



Window size=50, length=801, TE@MARINERN12_AG:1-801

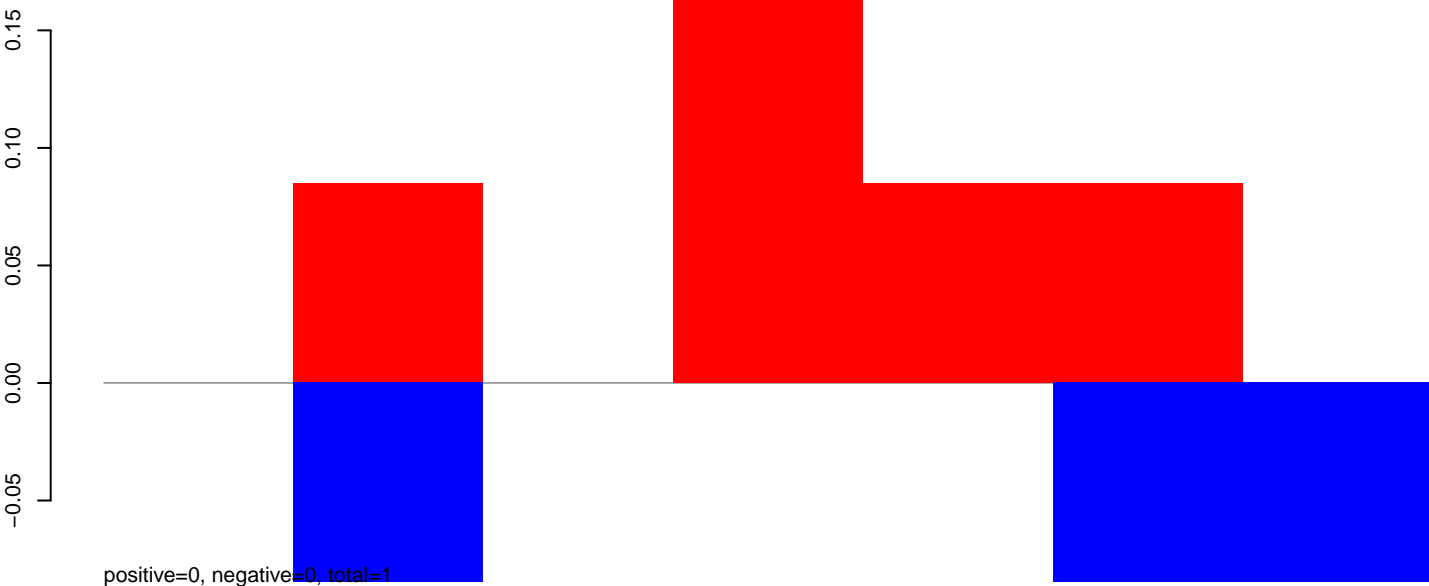
200

400

600

800

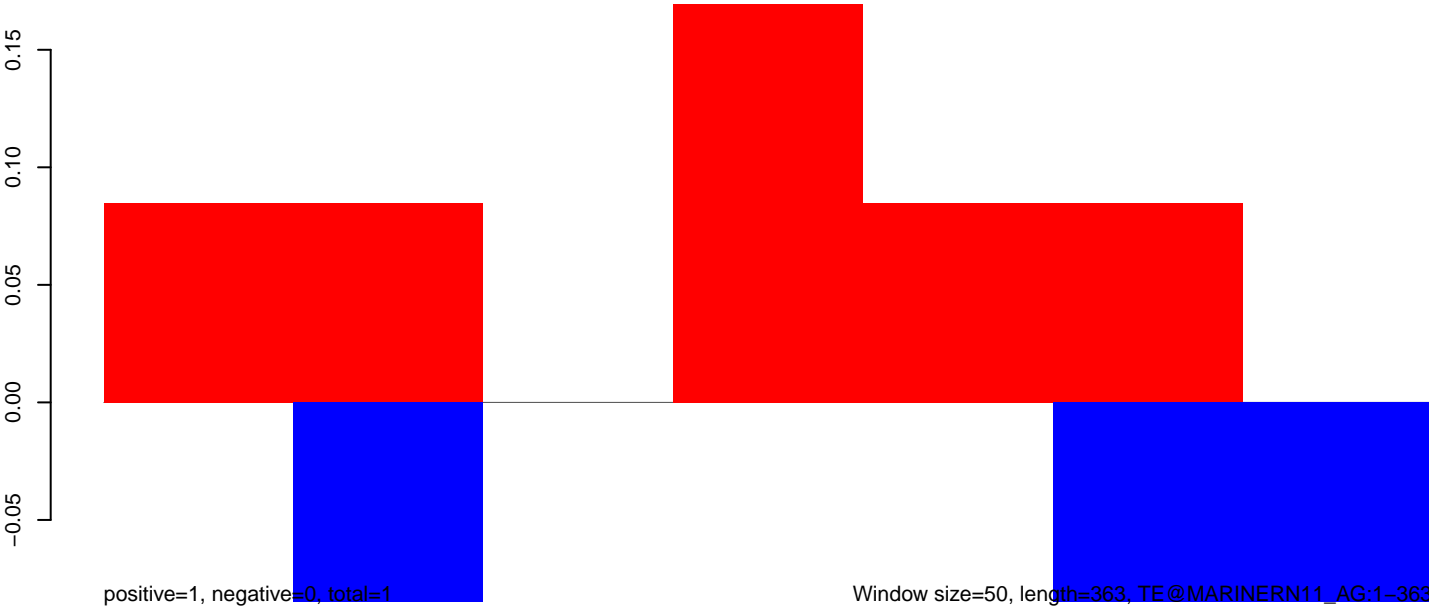
AnGam_Mos55_cells.18_23.rep



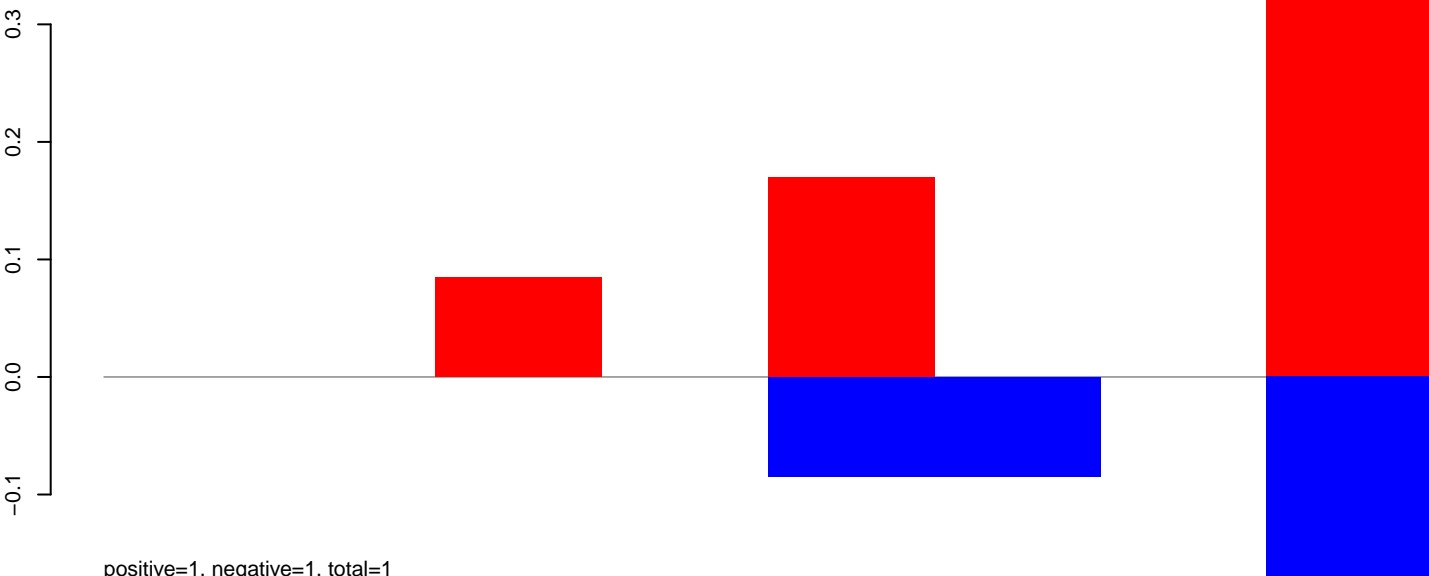
AnGam_Mos55_cells.24_35.rep



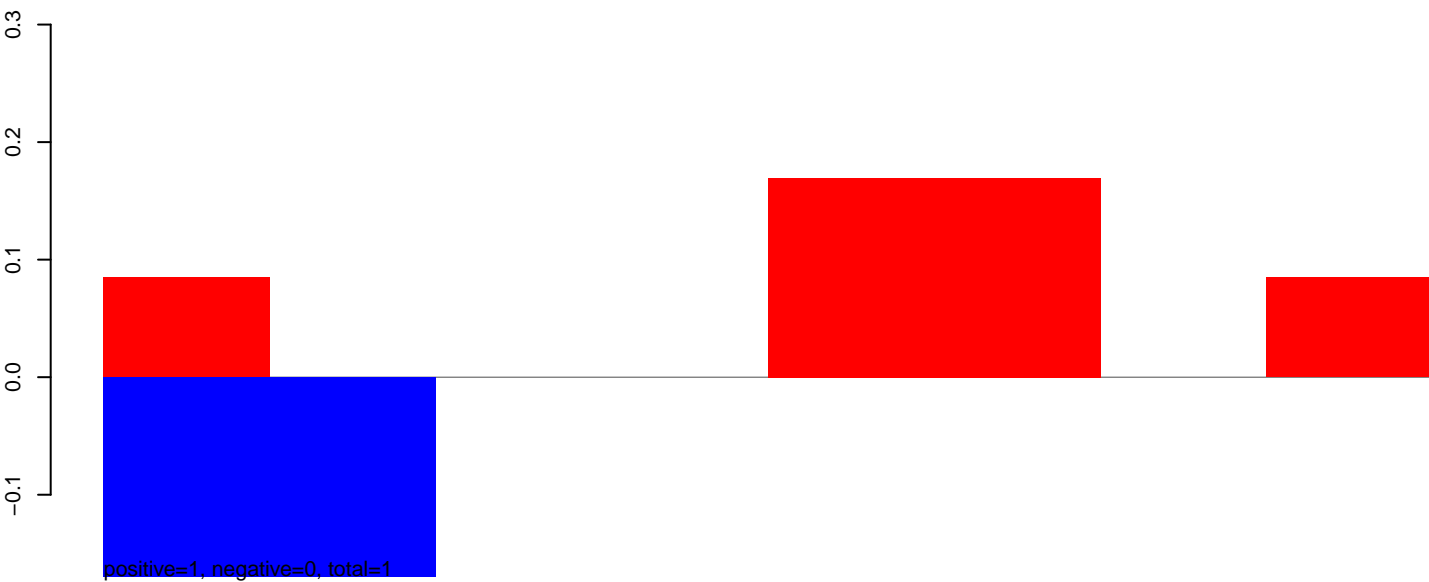
AnGam_Mos55_cells.rep



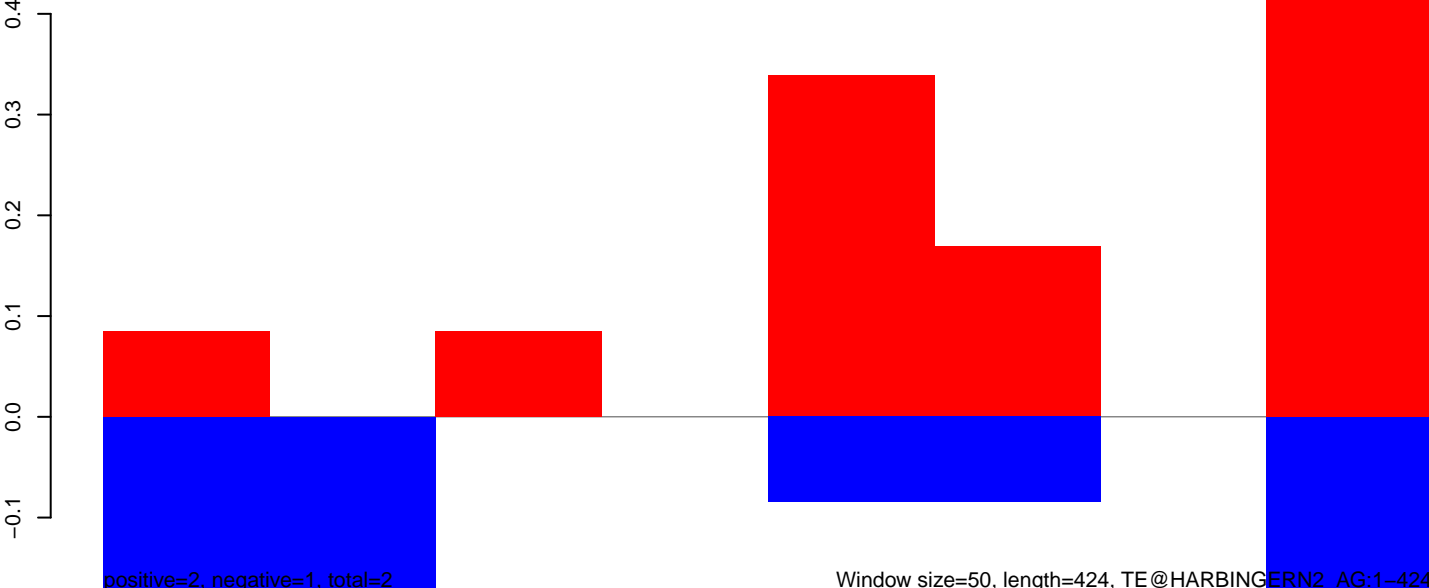
AnGam_Mos55_cells.18_23.rep



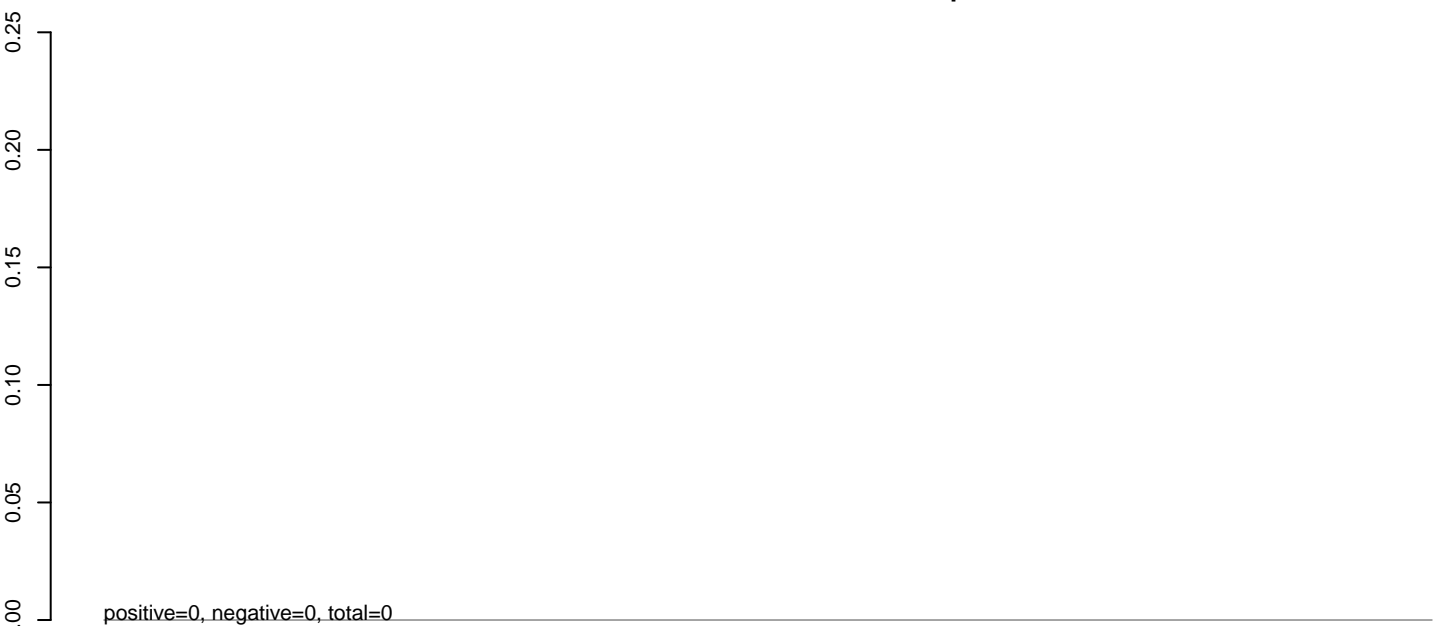
AnGam_Mos55_cells.24_35.rep



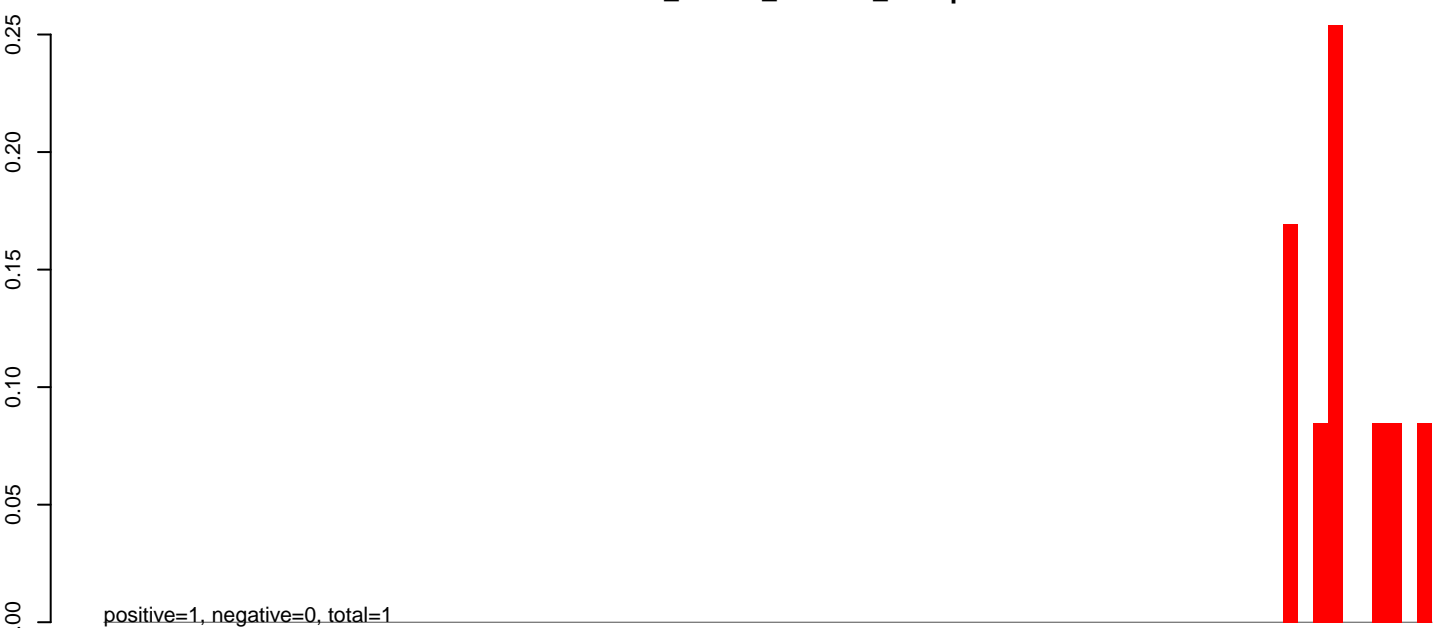
AnGam_Mos55_cells.rep



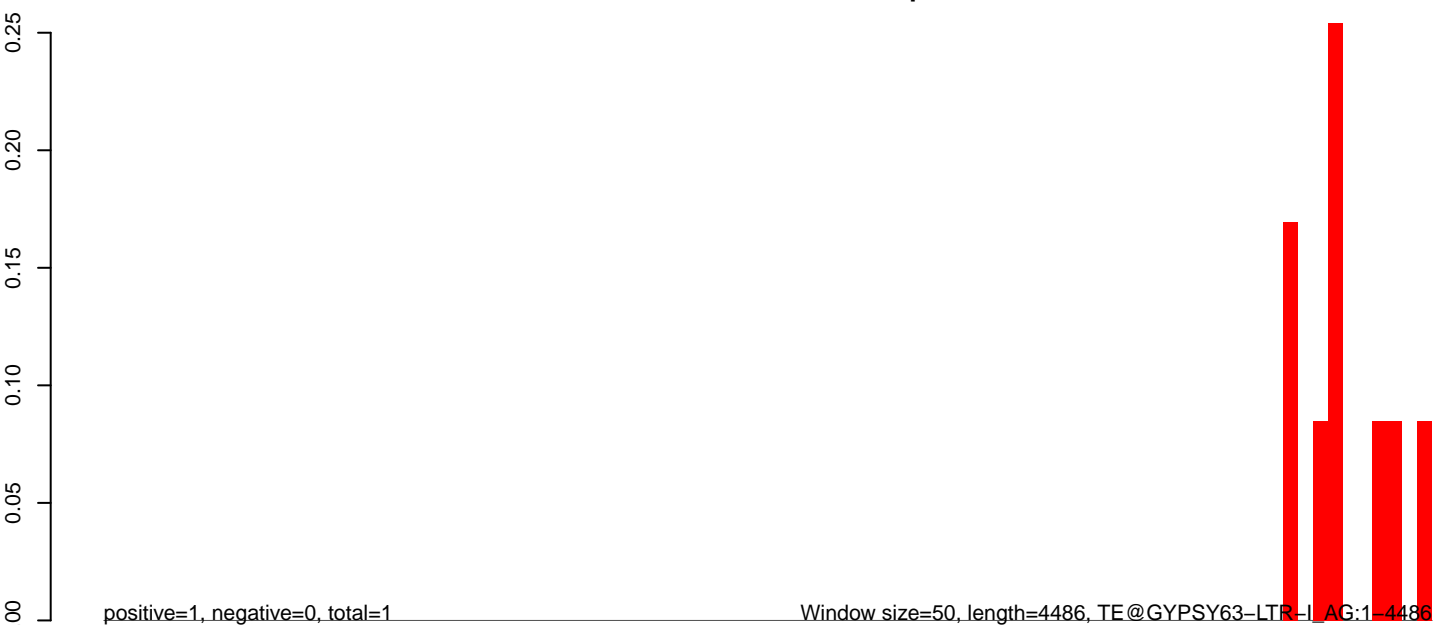
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

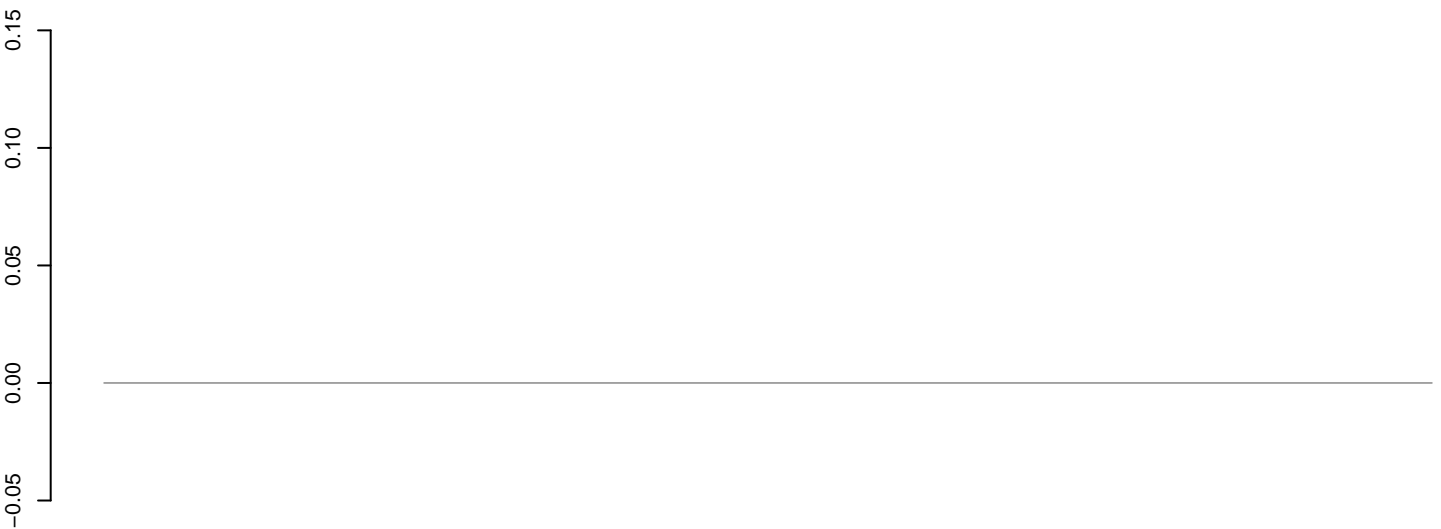


AnGam_Mos55_cells.rep



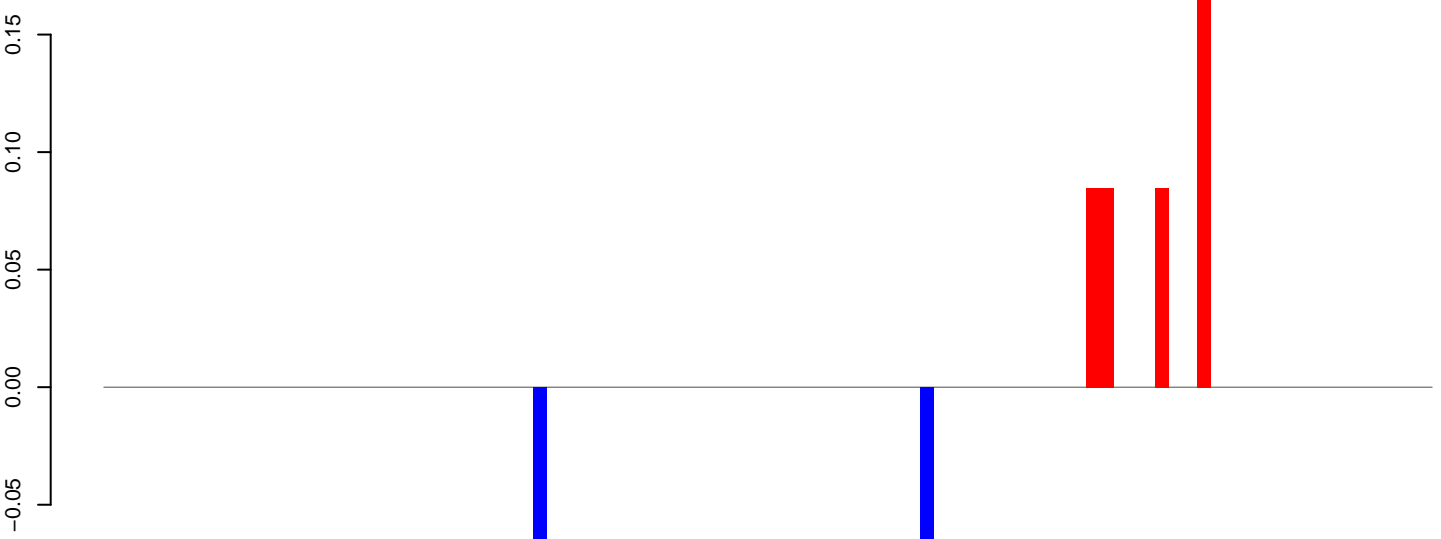
0 1000 2000 3000 4000

AnGam_Mos55_cells.18_23.rep



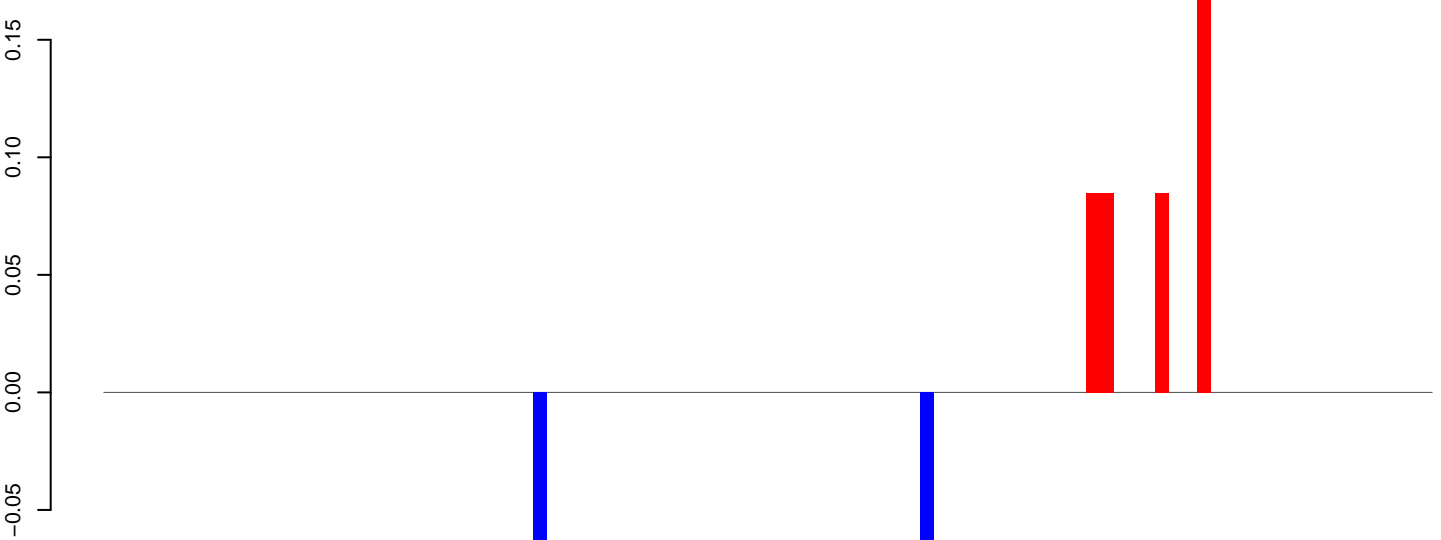
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



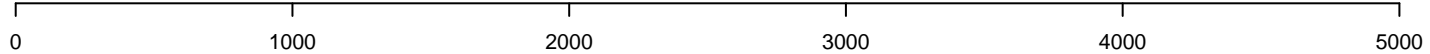
positive=0, negative=0, total=1

AnGam_Mos55_cells.rep

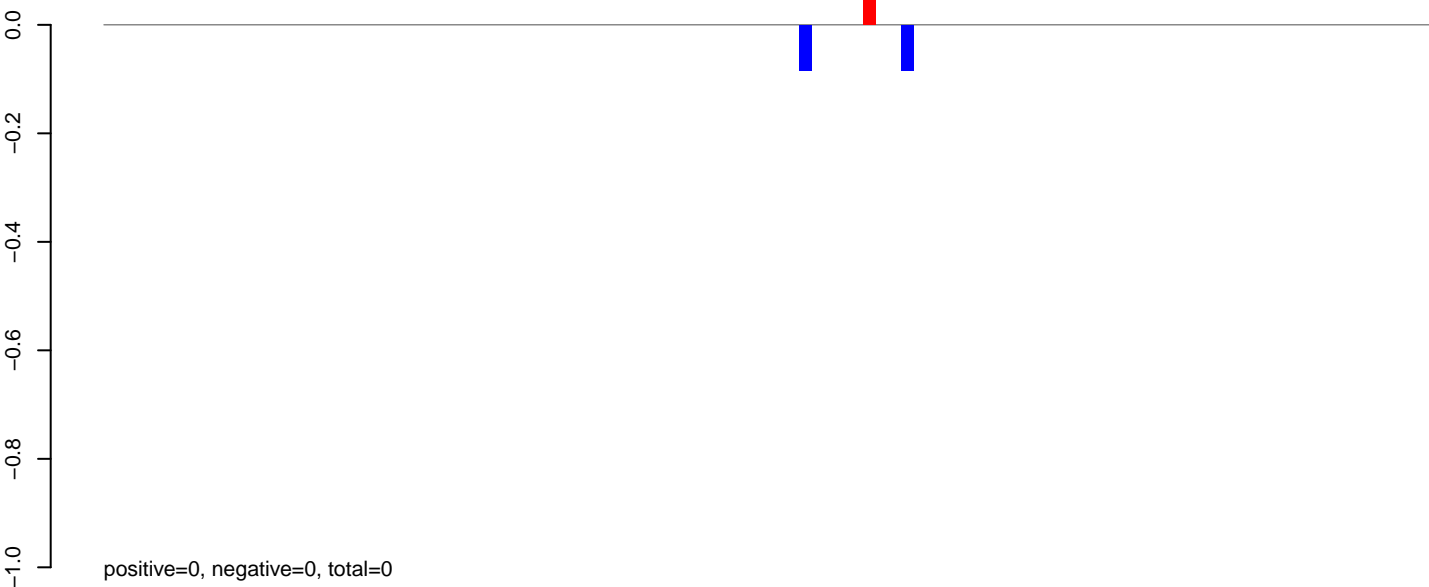


positive=0, negative=0, total=1

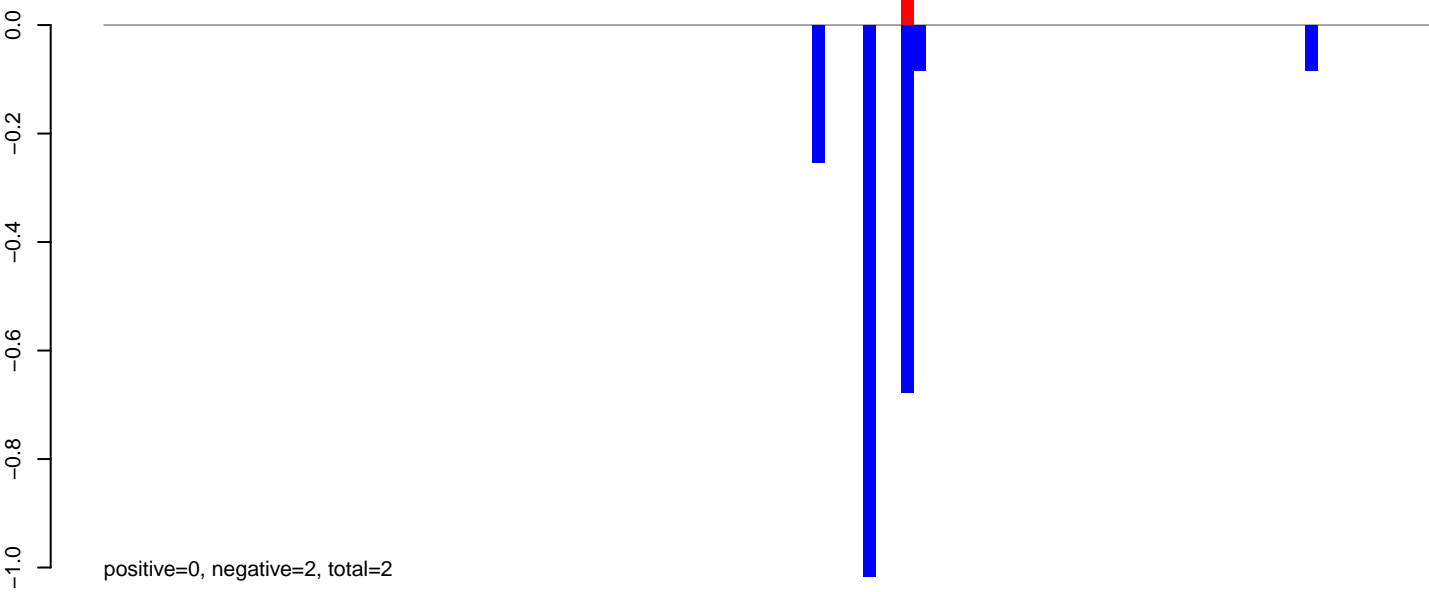
Window size=50, length=4840, TE@GYPSY59-LTR-L_AG:1-4840



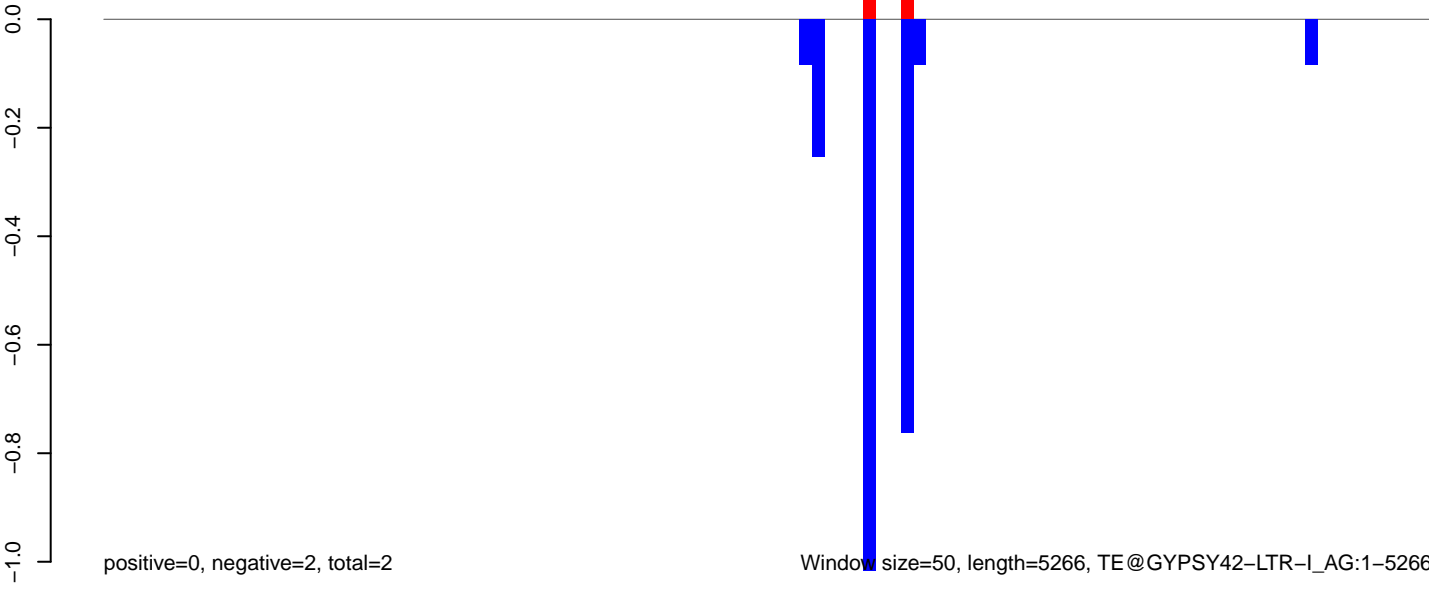
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

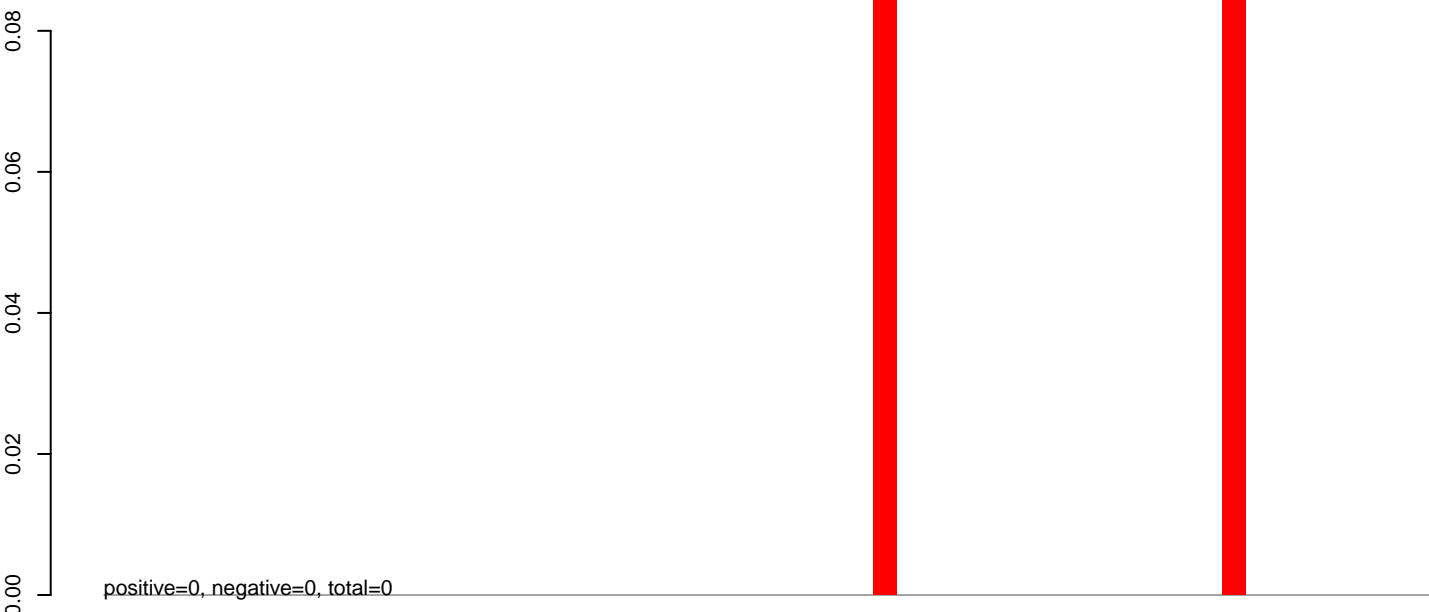


AnGam_Mos55_cells.rep

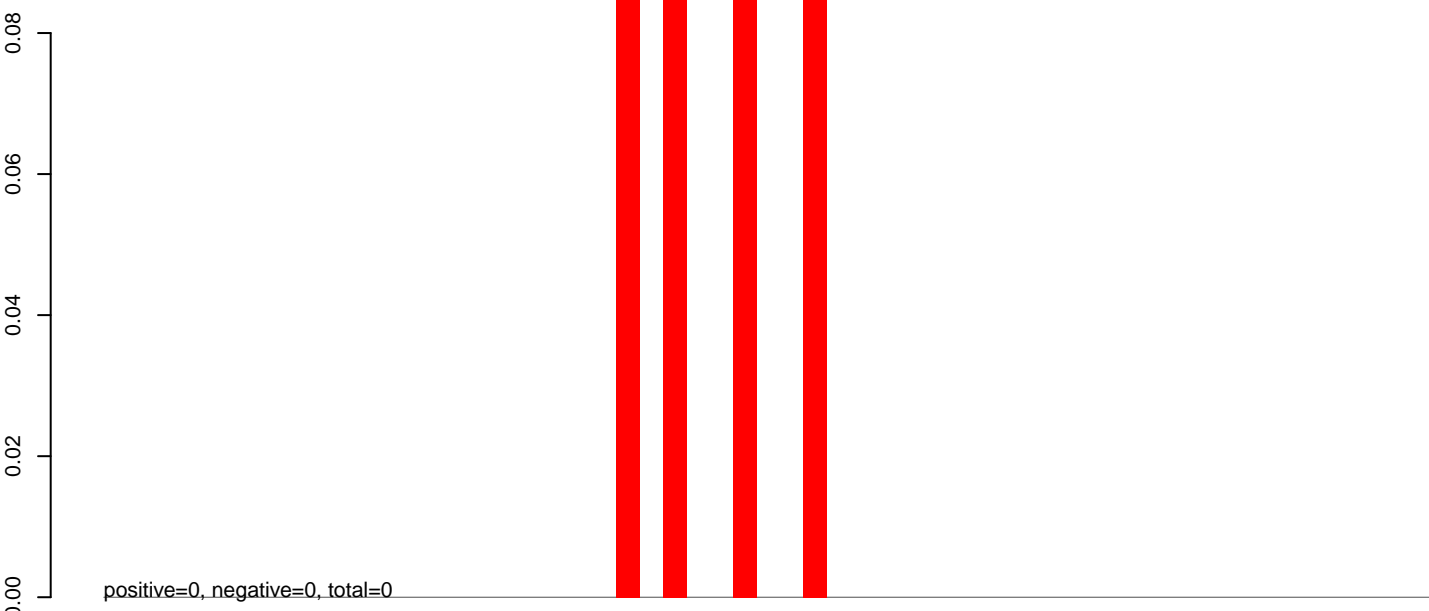


0 1000 2000 3000 4000 5000

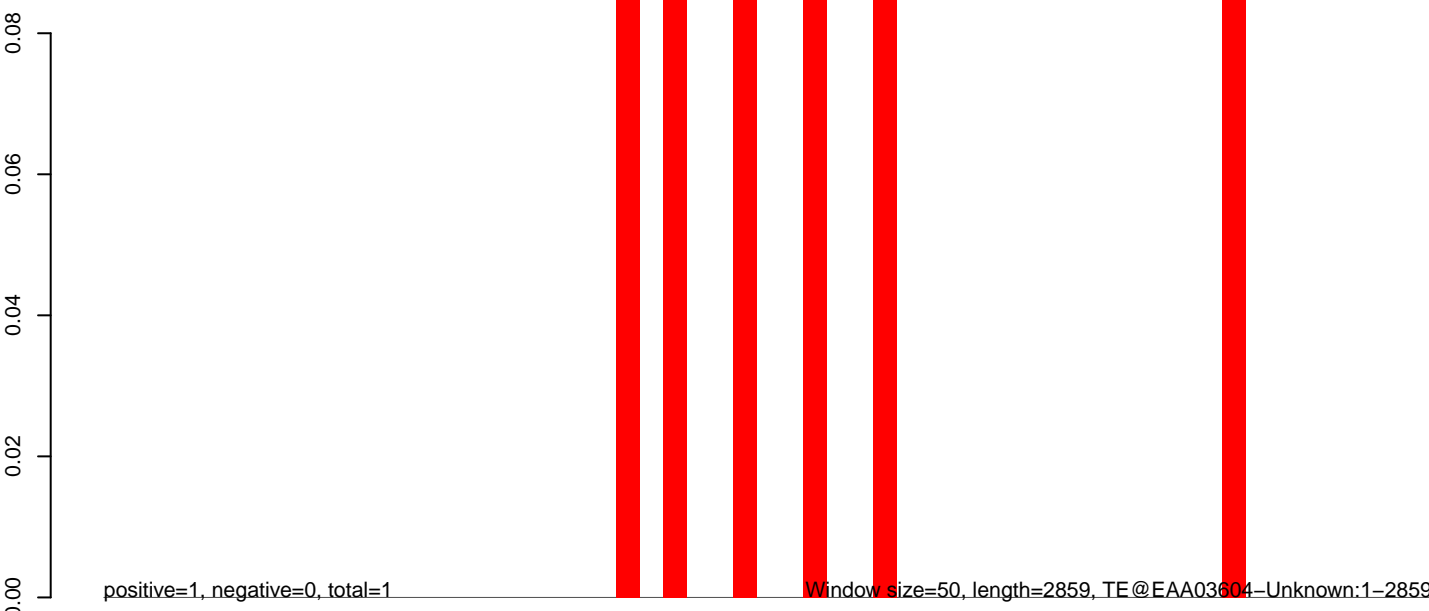
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



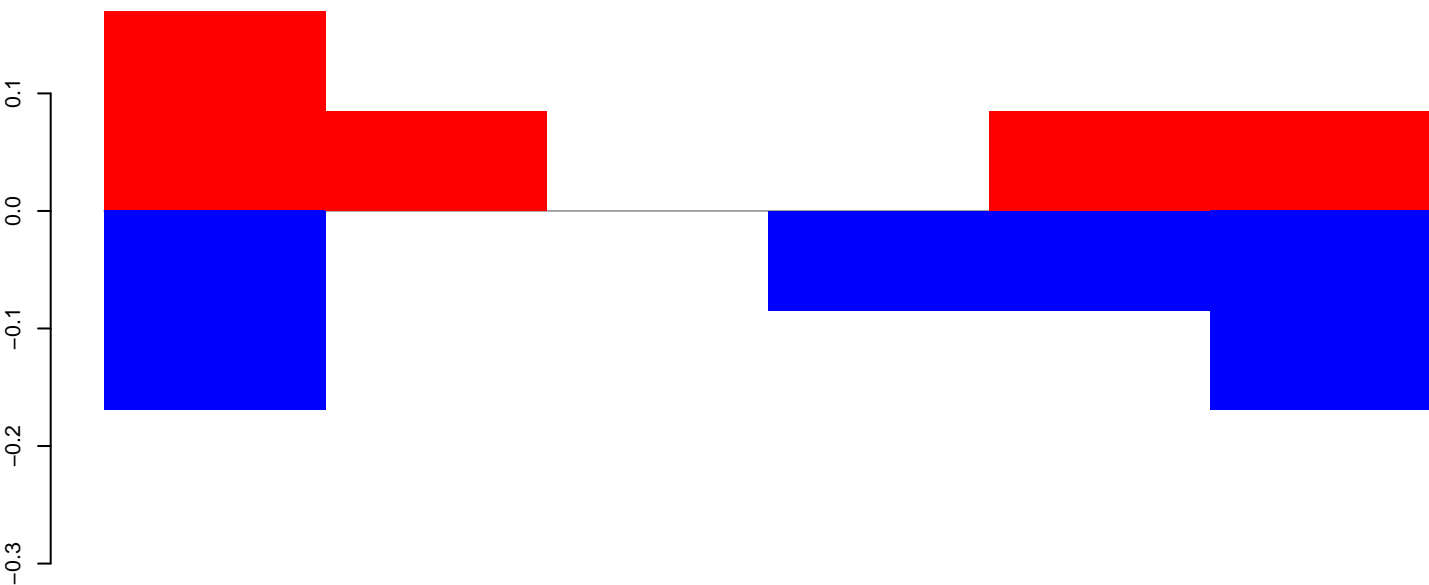
AnGam_Mos55_cells.rep



Window size=50, length=2859, TE@EAA03604-Unknown:1-2859

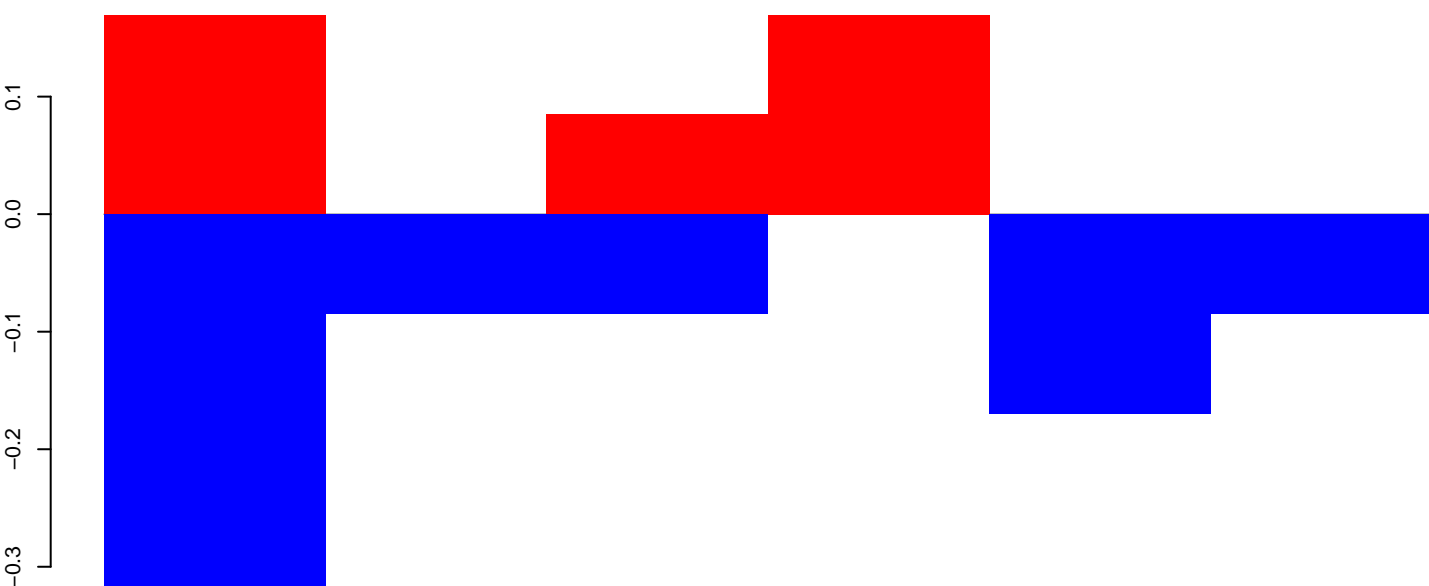
0 500 1000 1500 2000 2500 3000

AnGam_Mos55_cells.18_23.rep



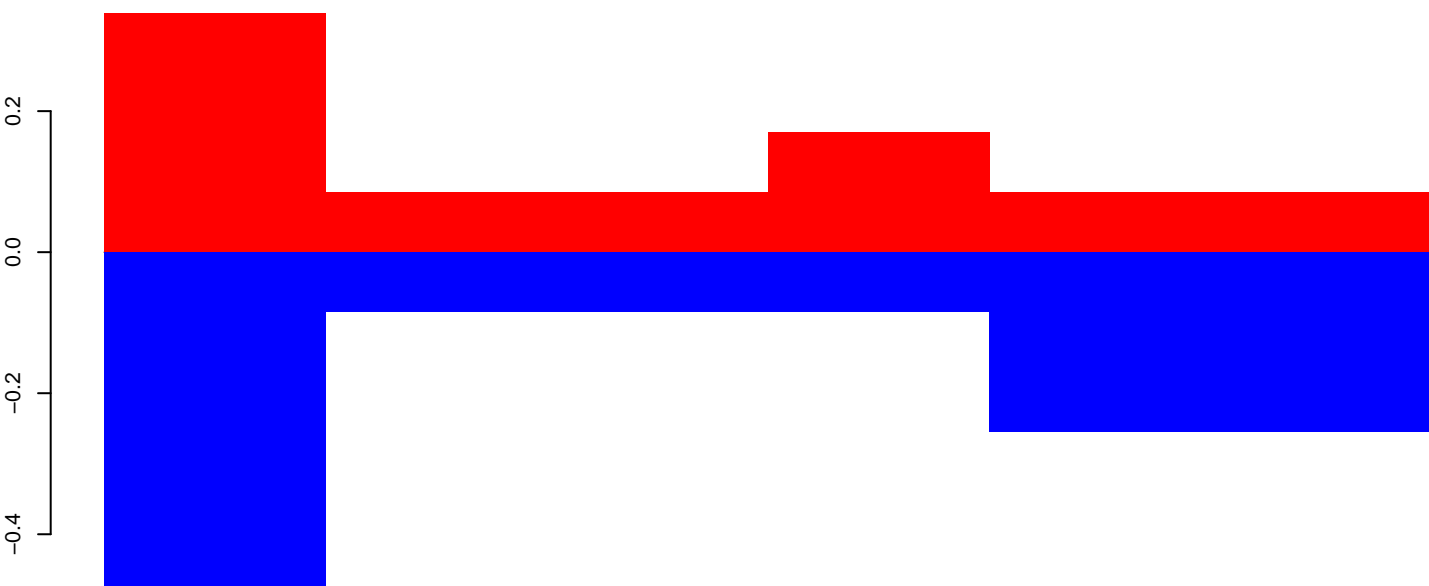
positive=1, negative=1, total=1

AnGam_Mos55_cells.24_35.rep



positive=0, negative=1, total=1

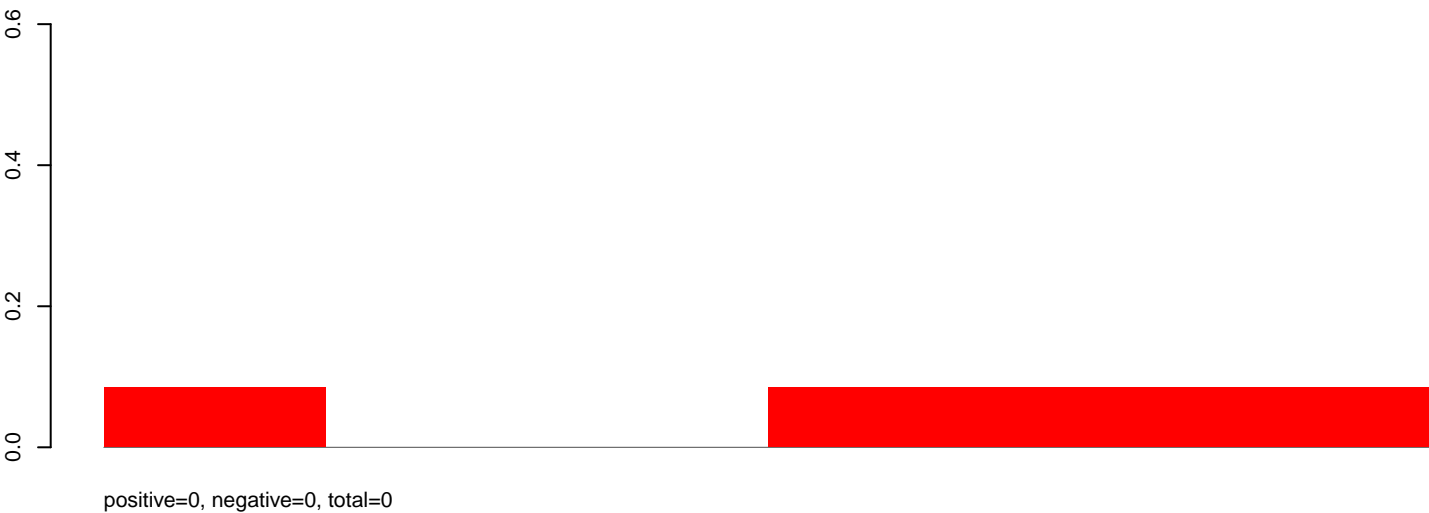
AnGam_Mos55_cells.rep



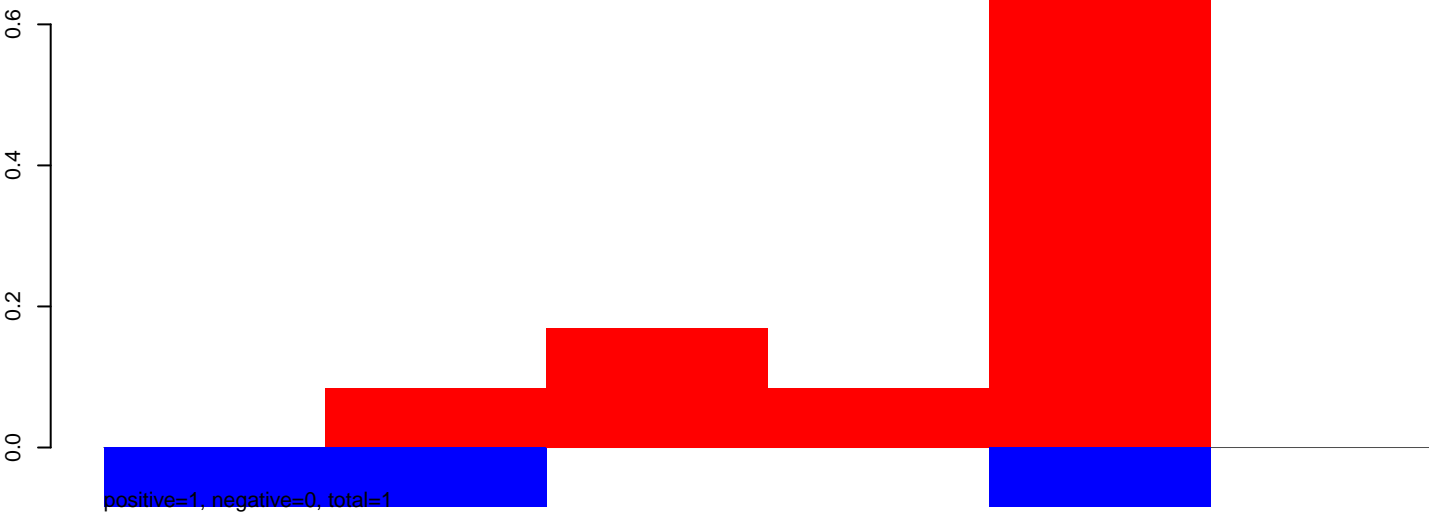
positive=1, negative=2, total=2

Window size=50, length=349, TE@DNA-5_AG:1-349

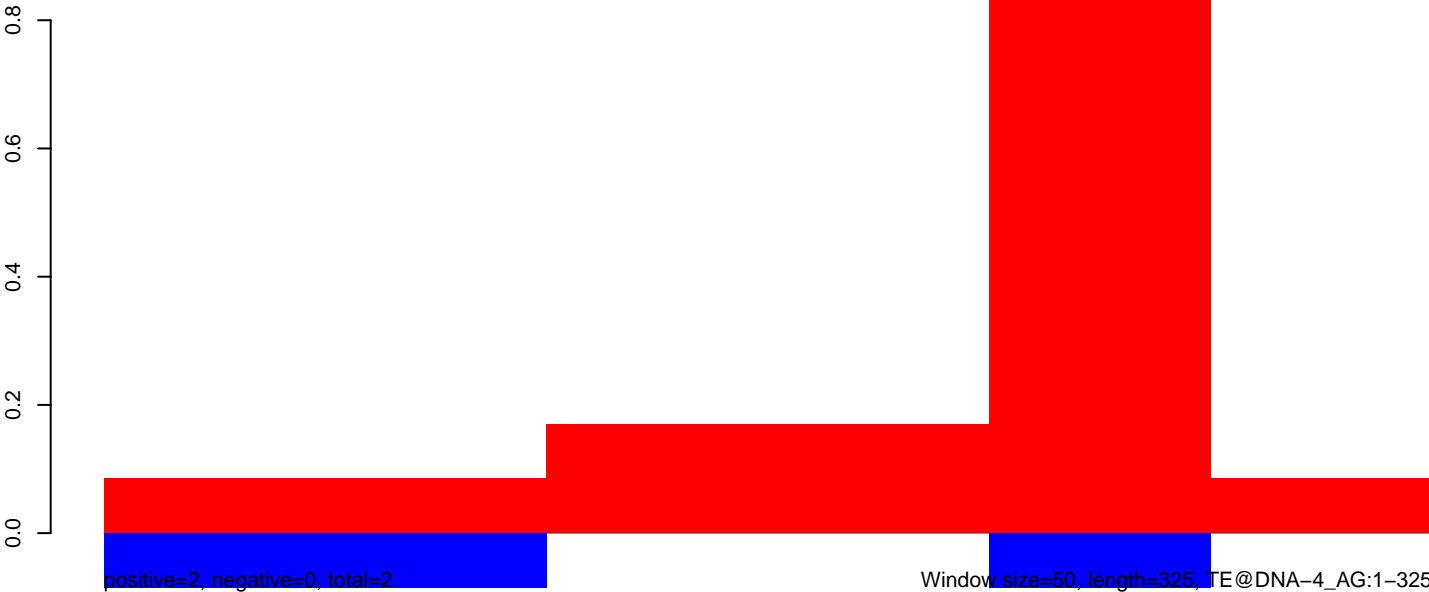
AnGam_Mos55_cells.18_23.rep



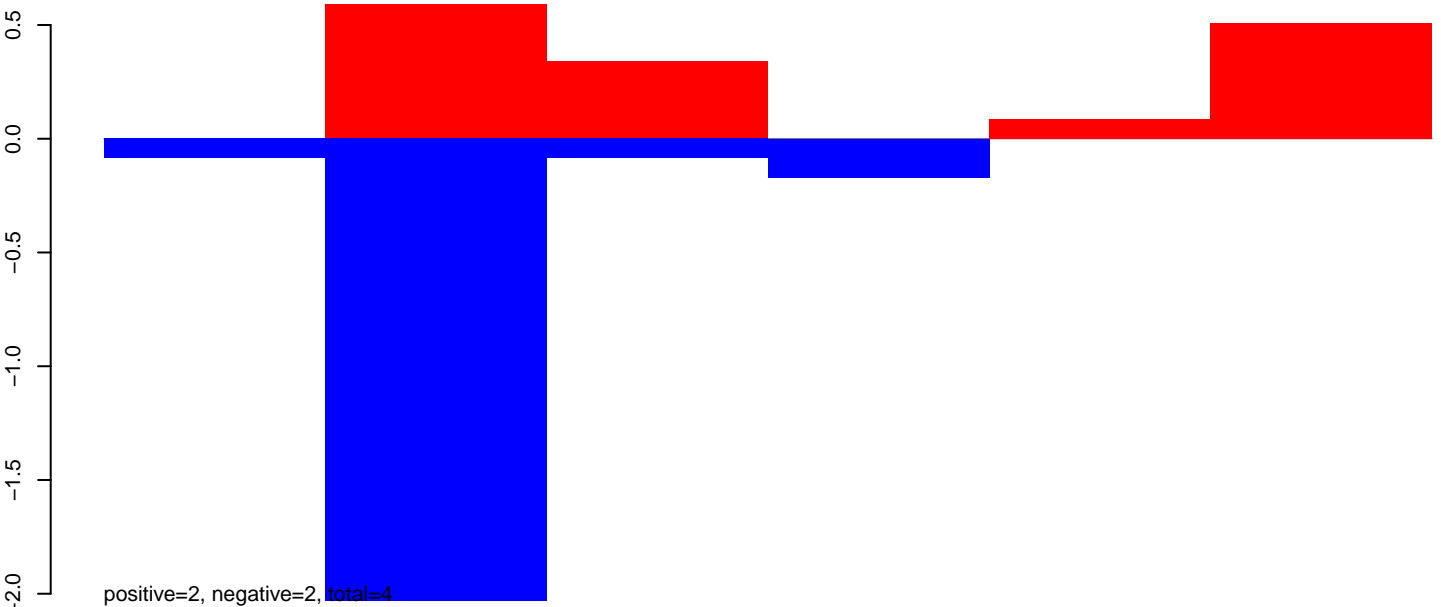
AnGam_Mos55_cells.24_35.rep



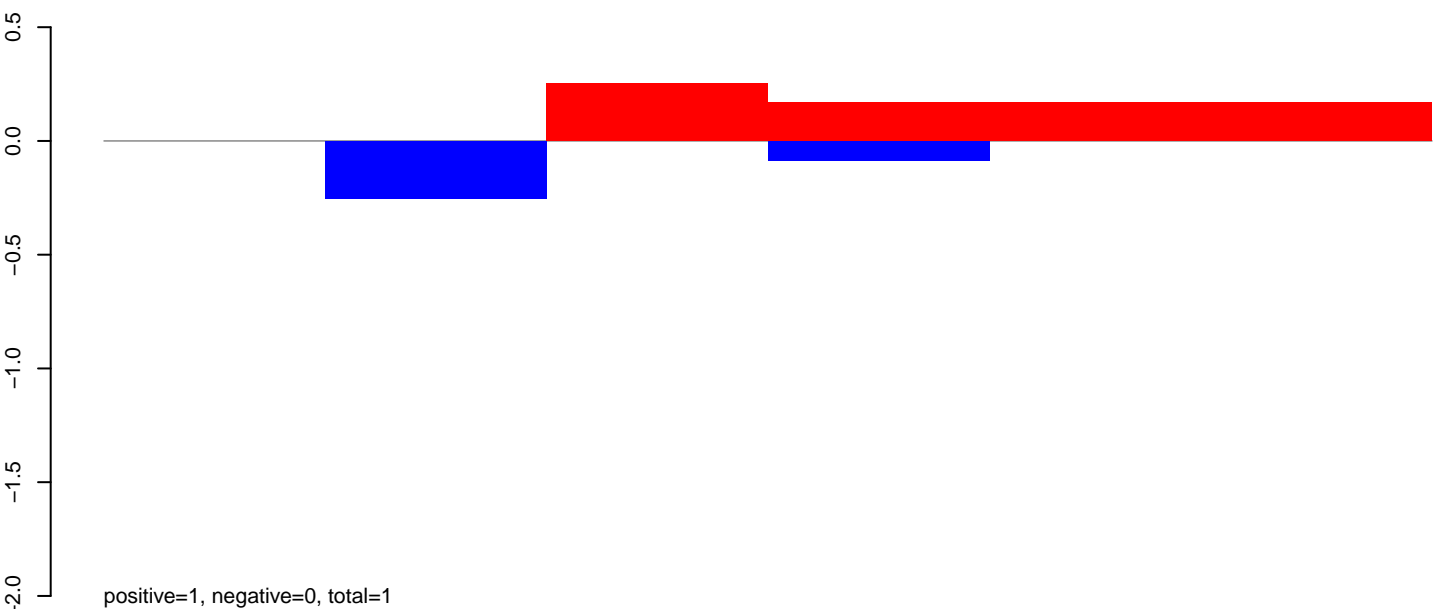
AnGam_Mos55_cells.rep



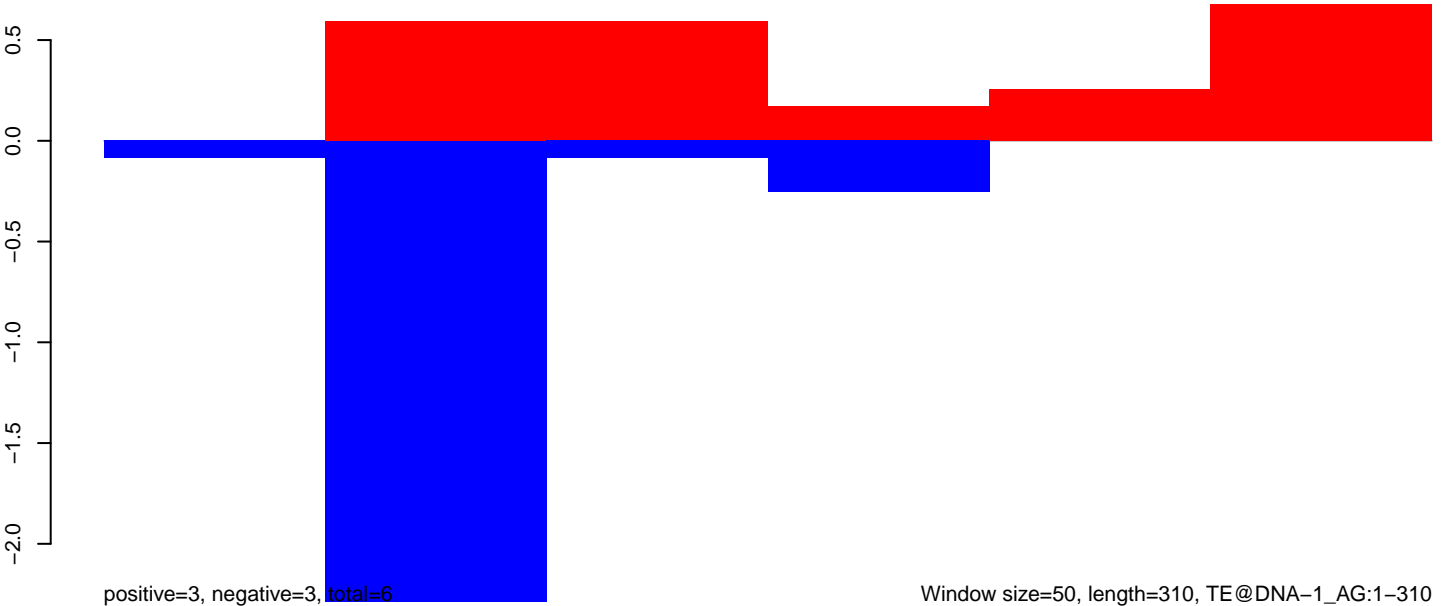
AnGam_Mos55_cells.18_23.rep



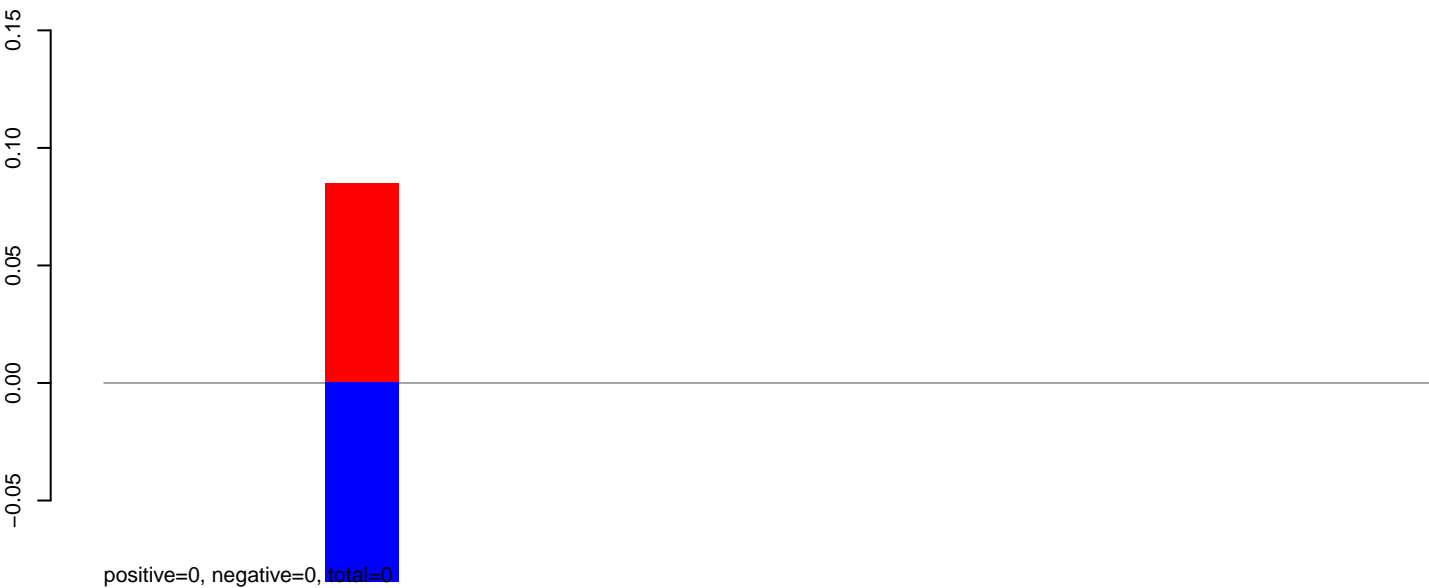
AnGam_Mos55_cells.24_35.rep



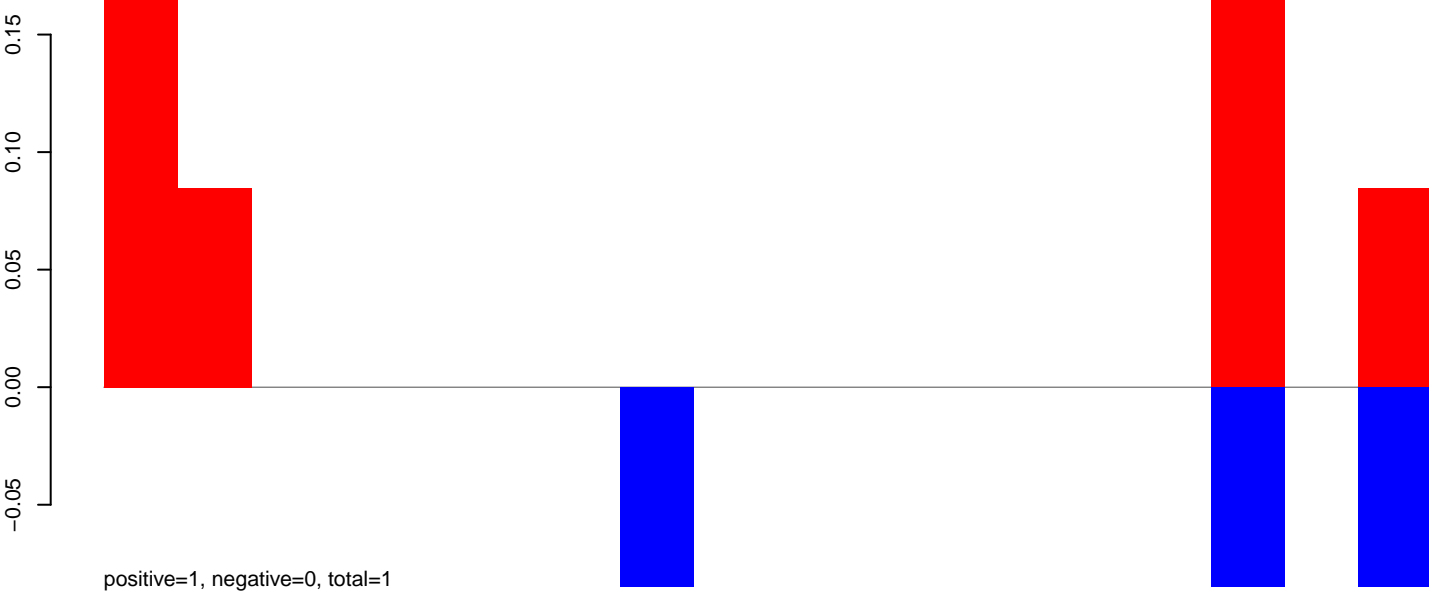
AnGam_Mos55_cells.rep



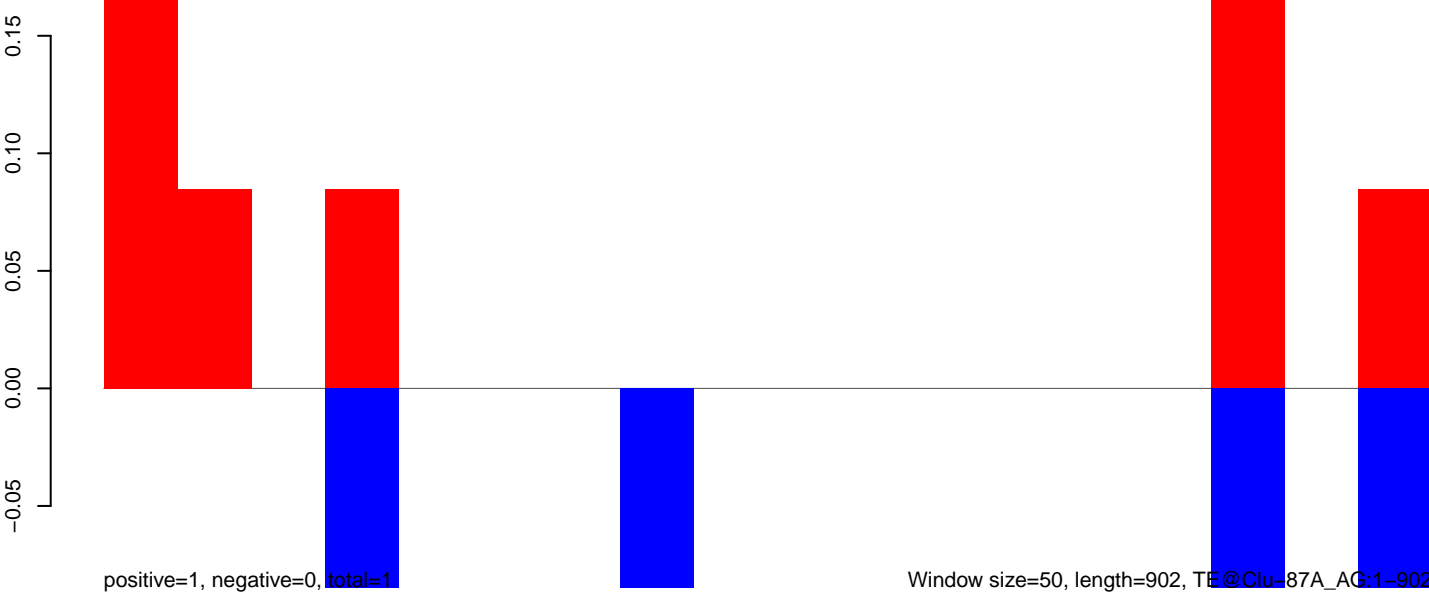
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



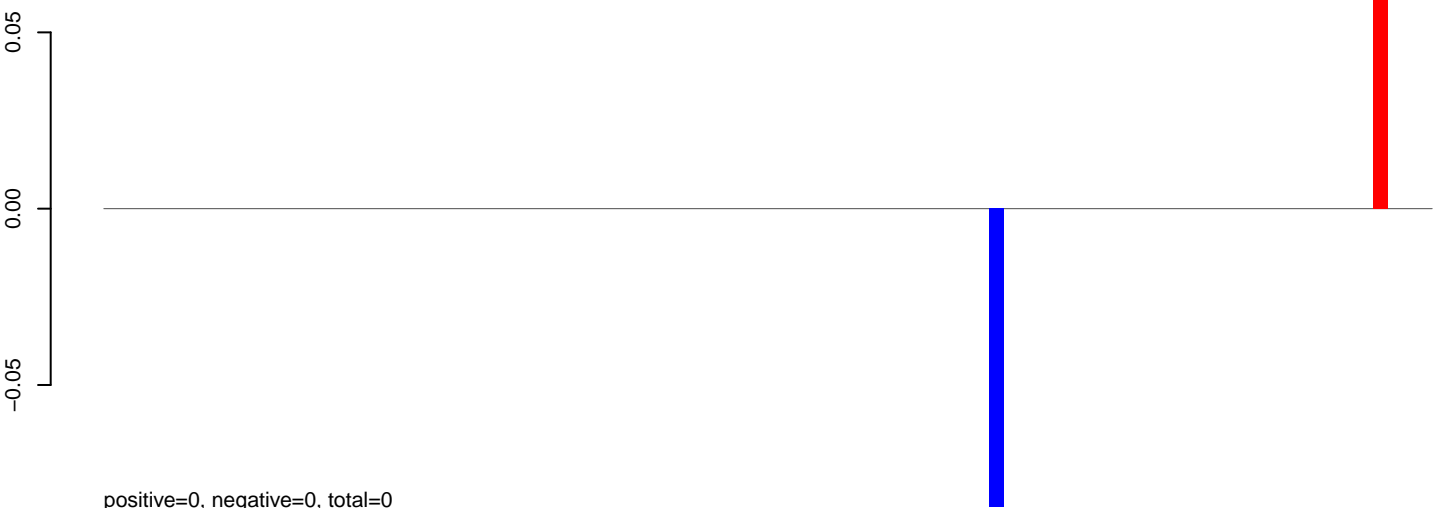
AnGam_Mos55_cells.rep



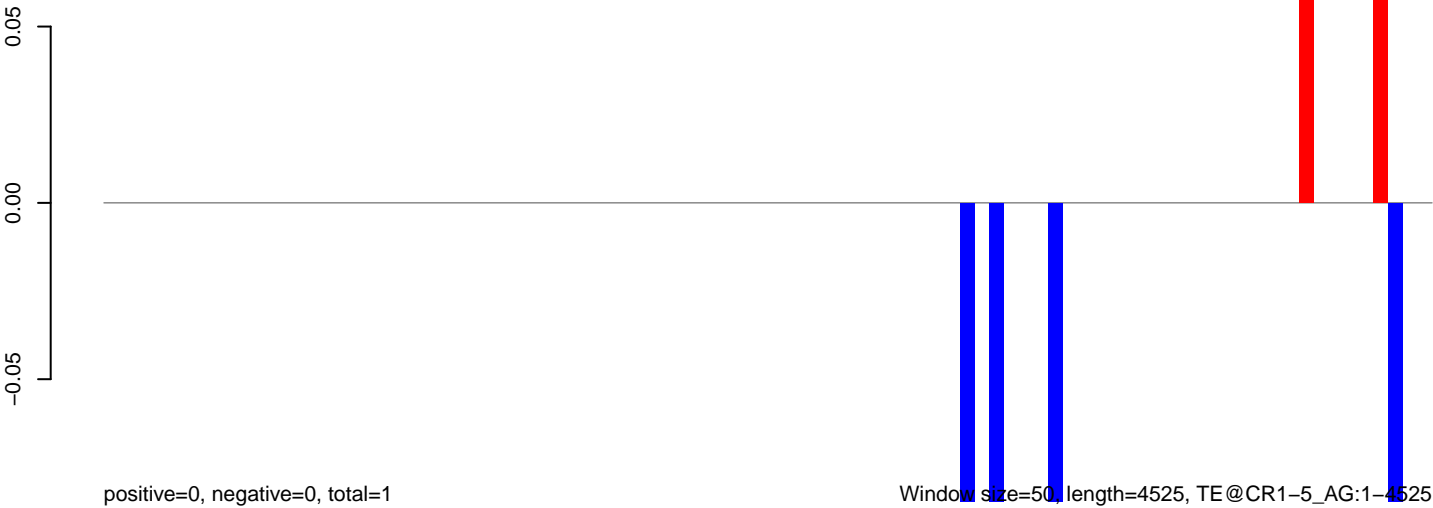
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



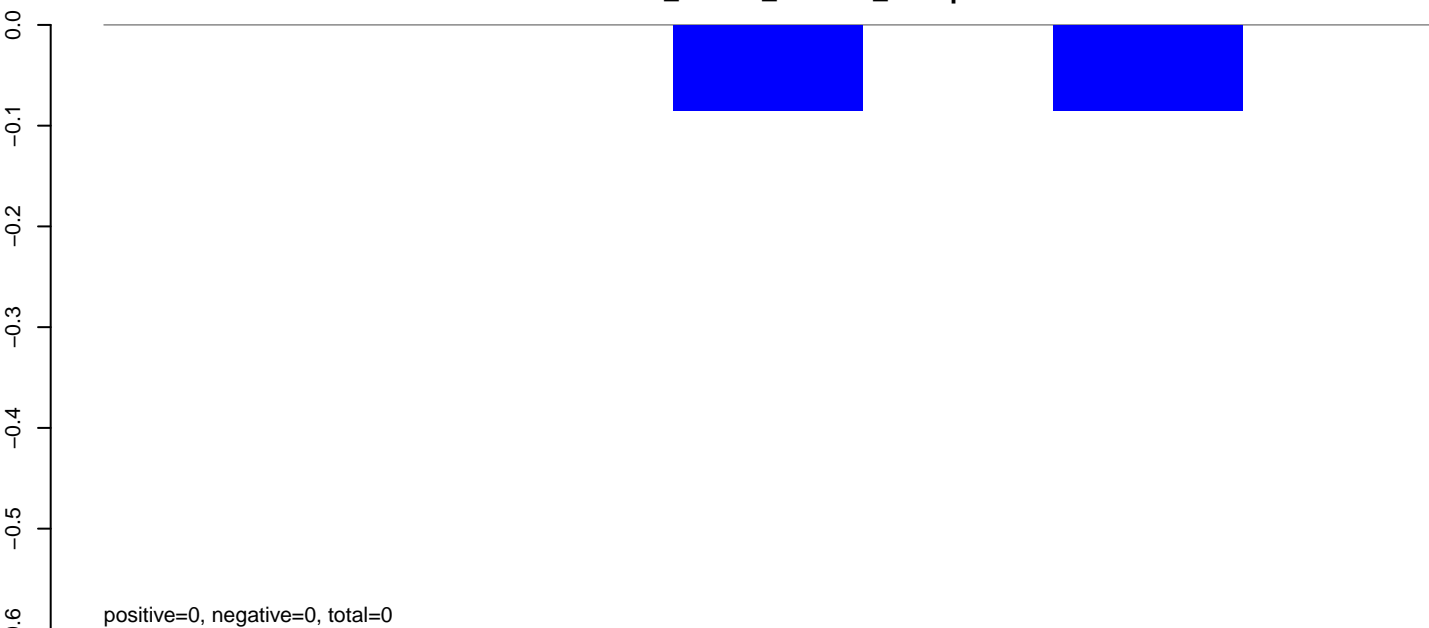
AnGam_Mos55_cells.rep



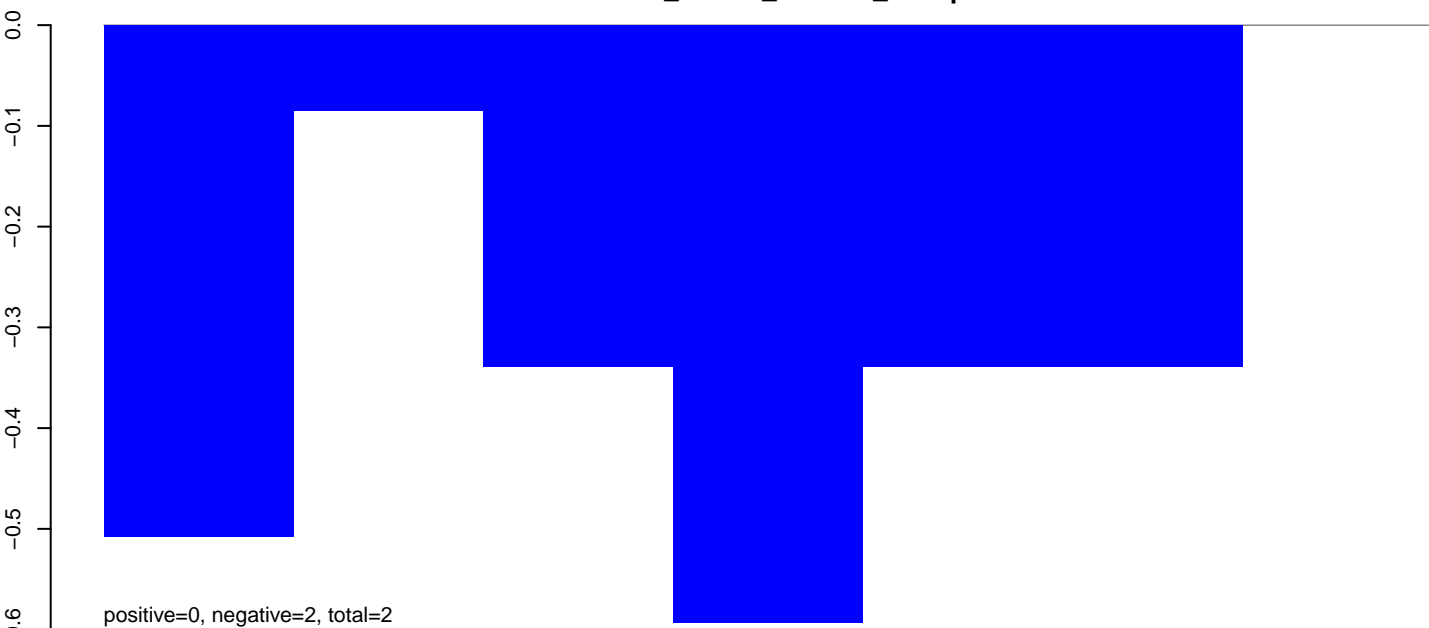
Window size=50, length=4525, TE@CR1-5_AG:1-4525

0 1000 2000 3000 4000

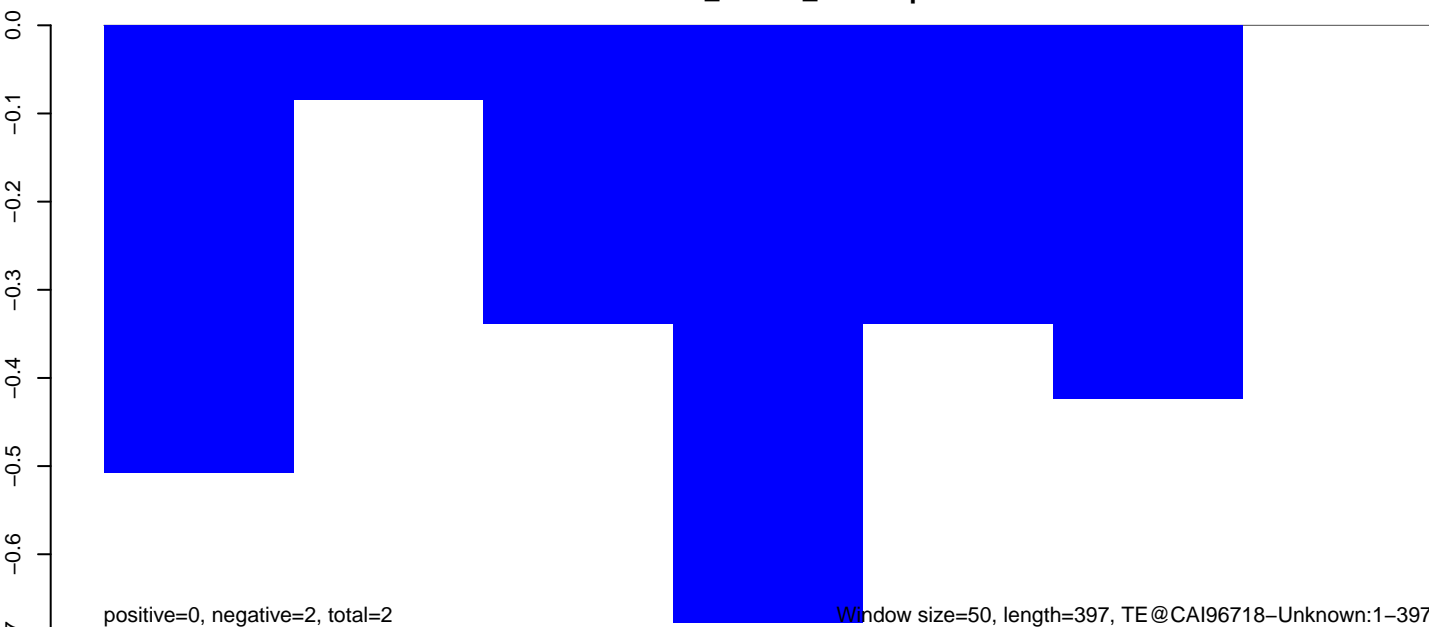
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



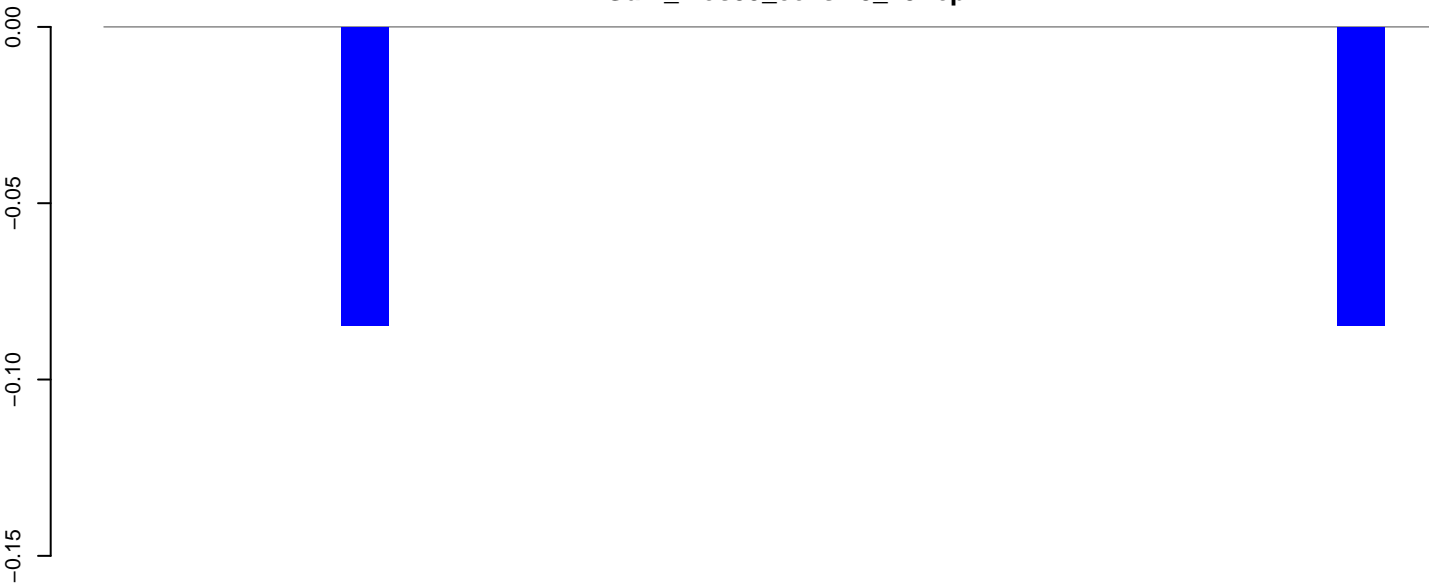
AnGam_Mos55_cells.rep



Window size=50, length=397, TE@CAI96718-Unknown:1-397

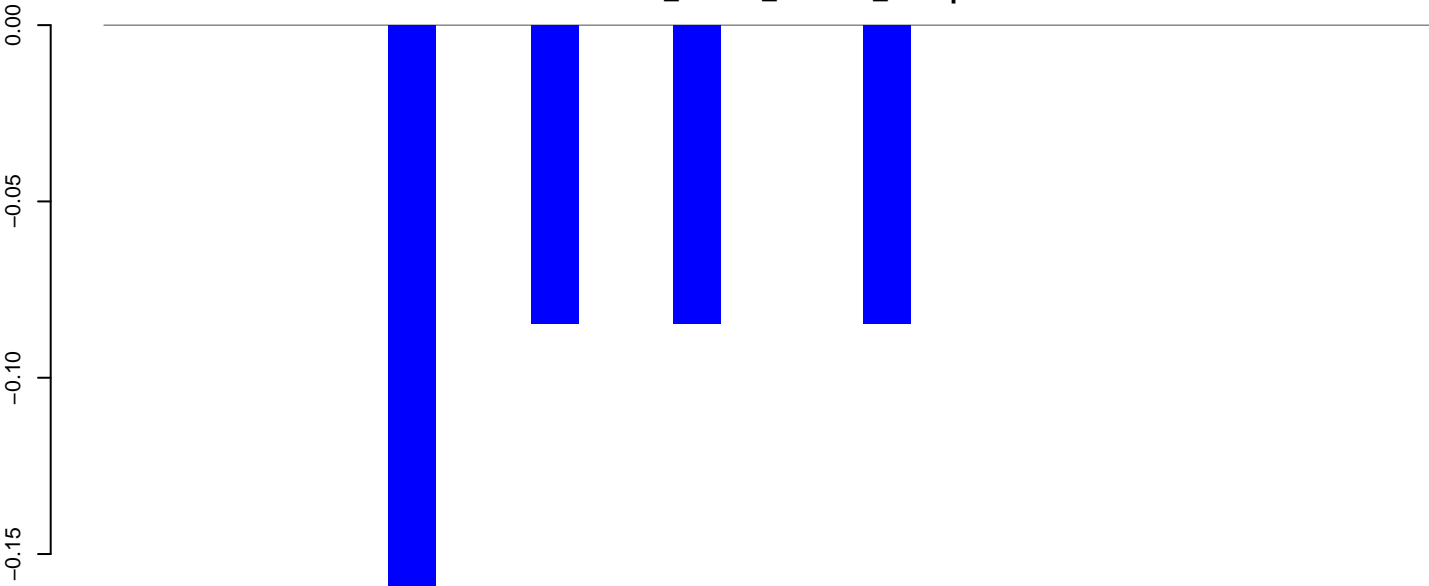
50 100 150 200 250 300 350 400

AnGam_Mos55_cells.18_23.rep



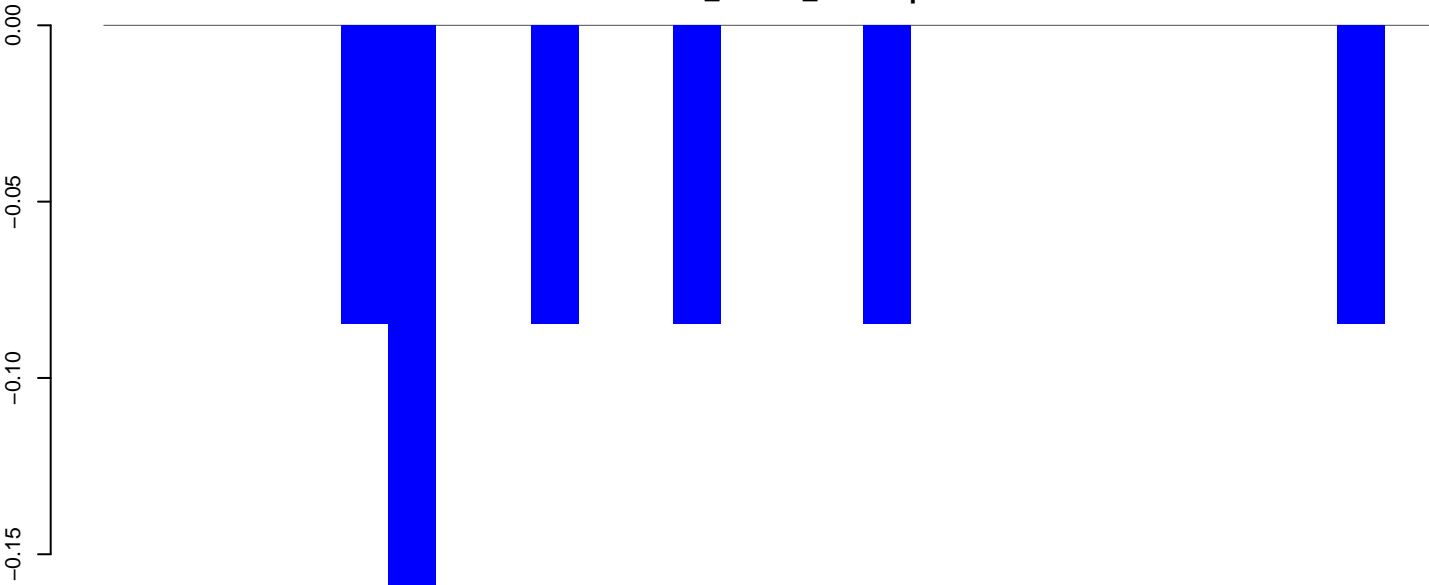
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



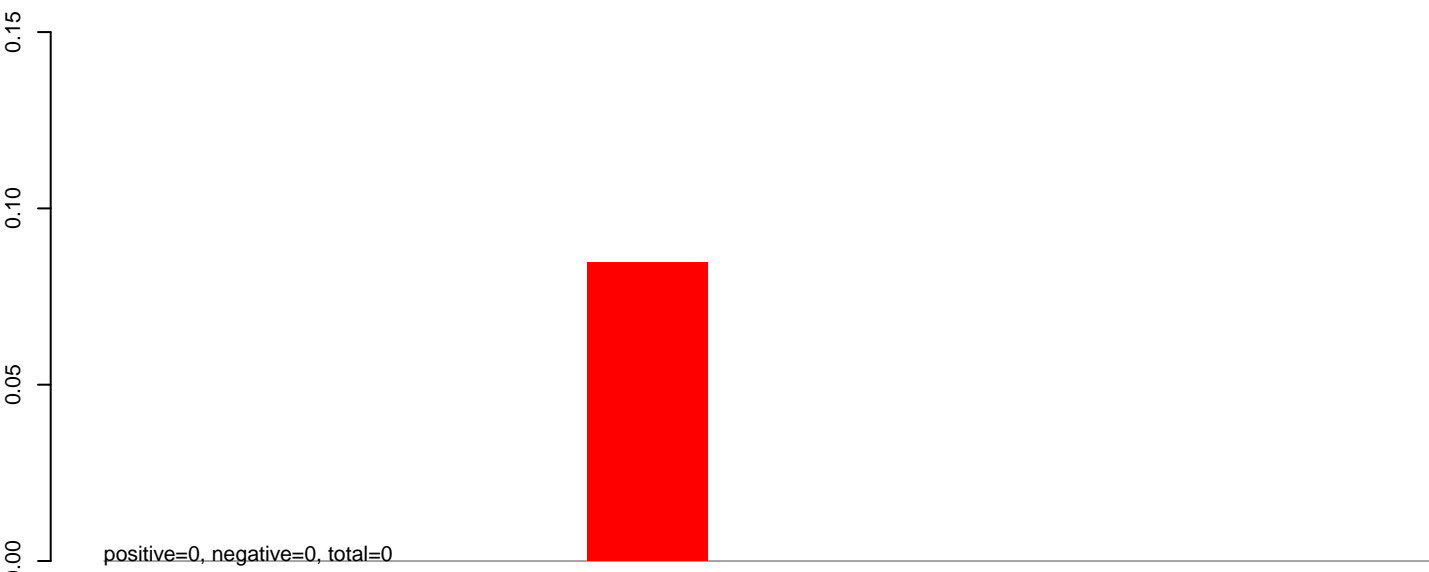
positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

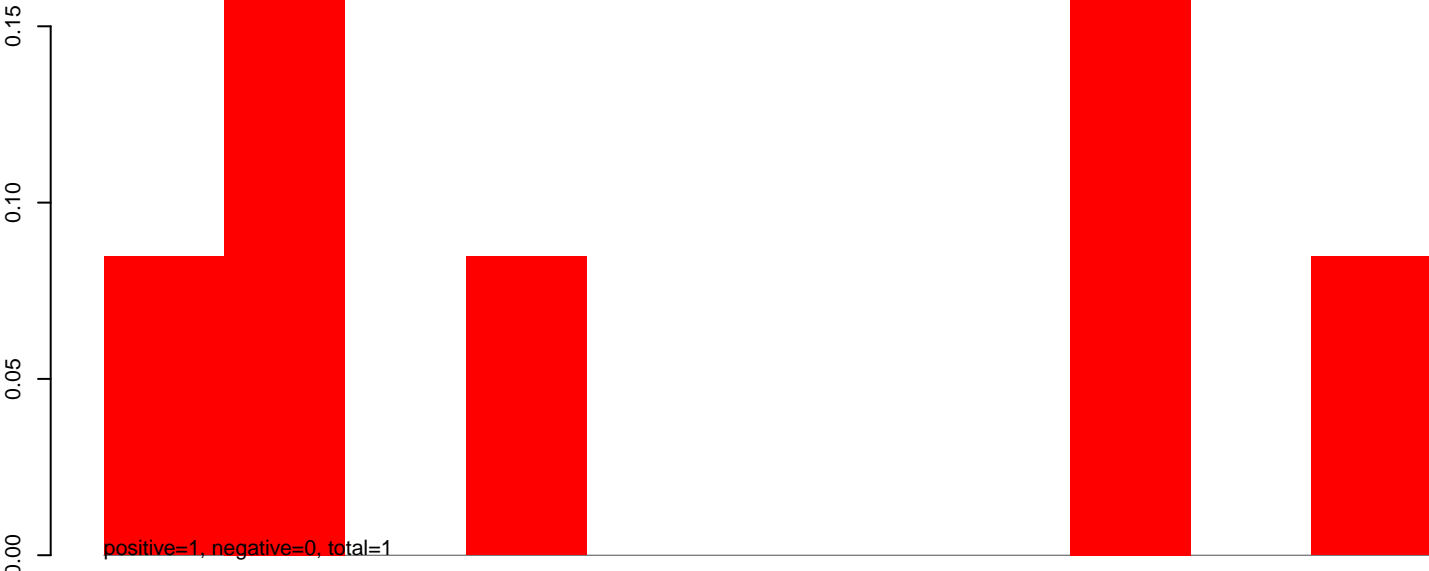


positive=0, negative=1, total=1

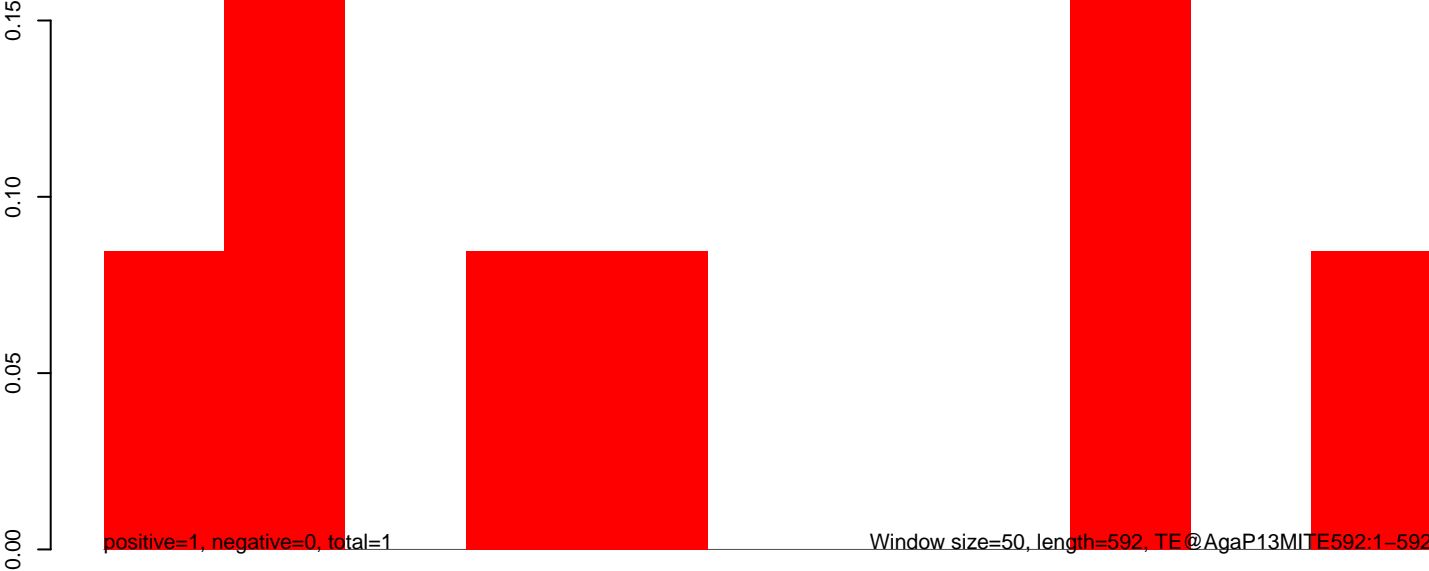
AnGam_Mos55_cells.18_23.rep



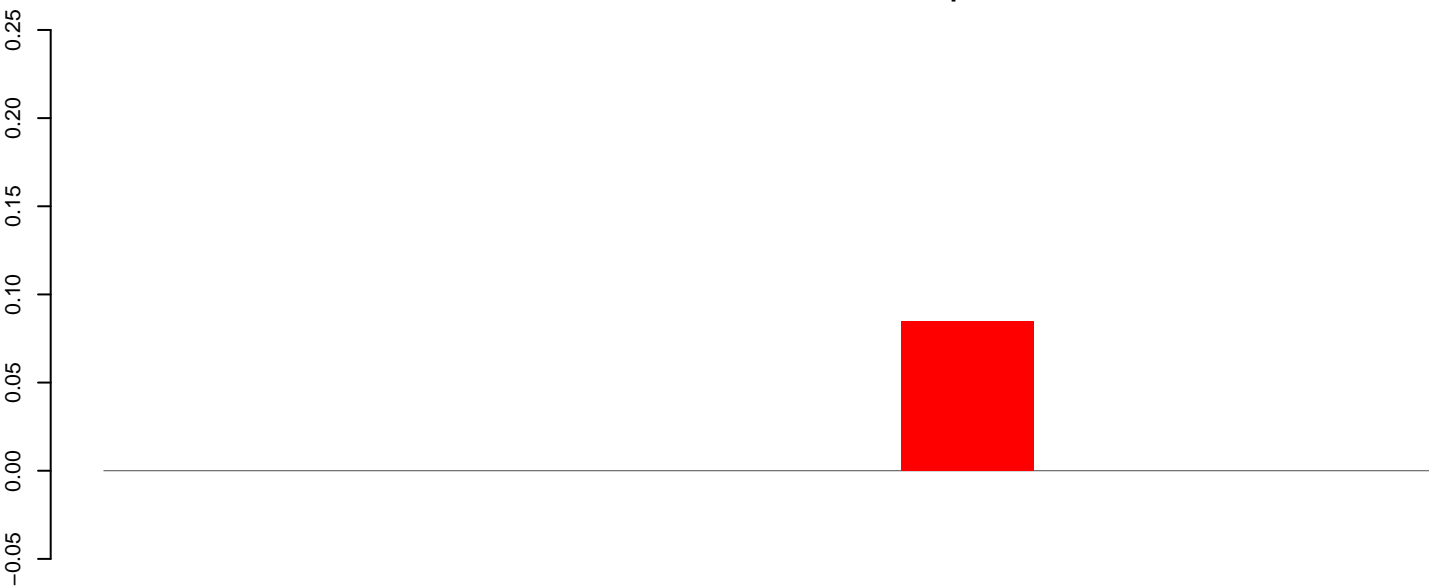
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

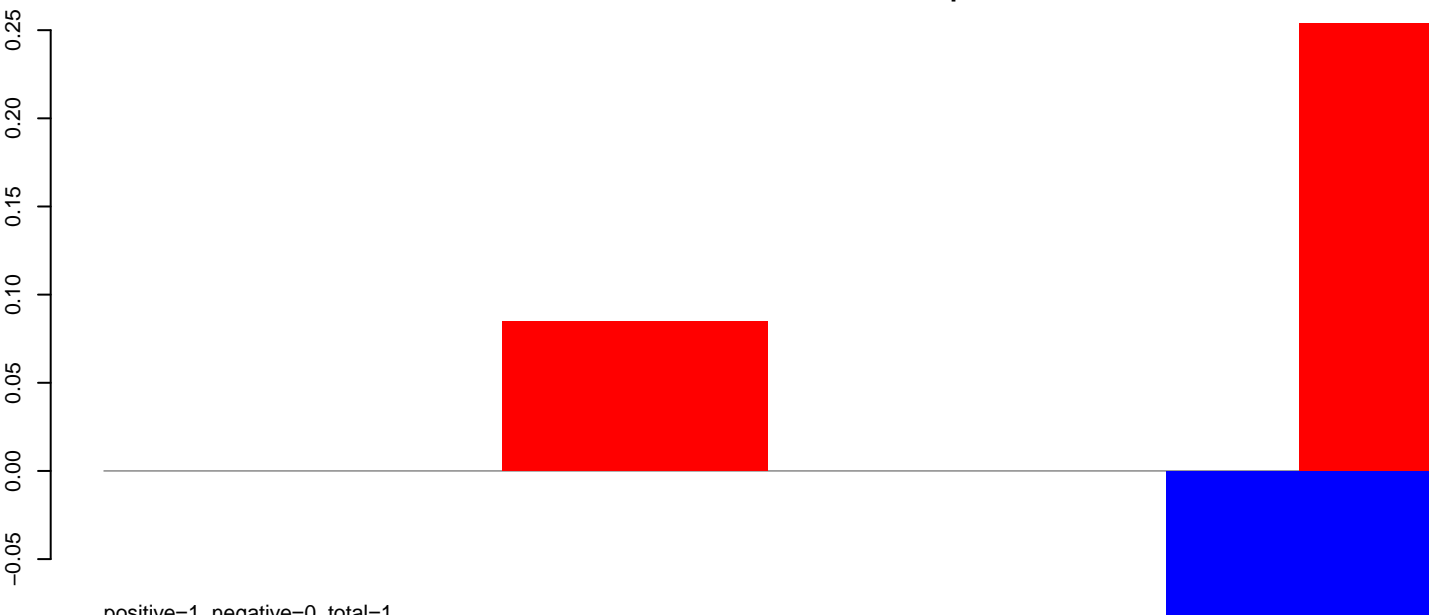


AnGam_Mos55_cells.18_23.rep



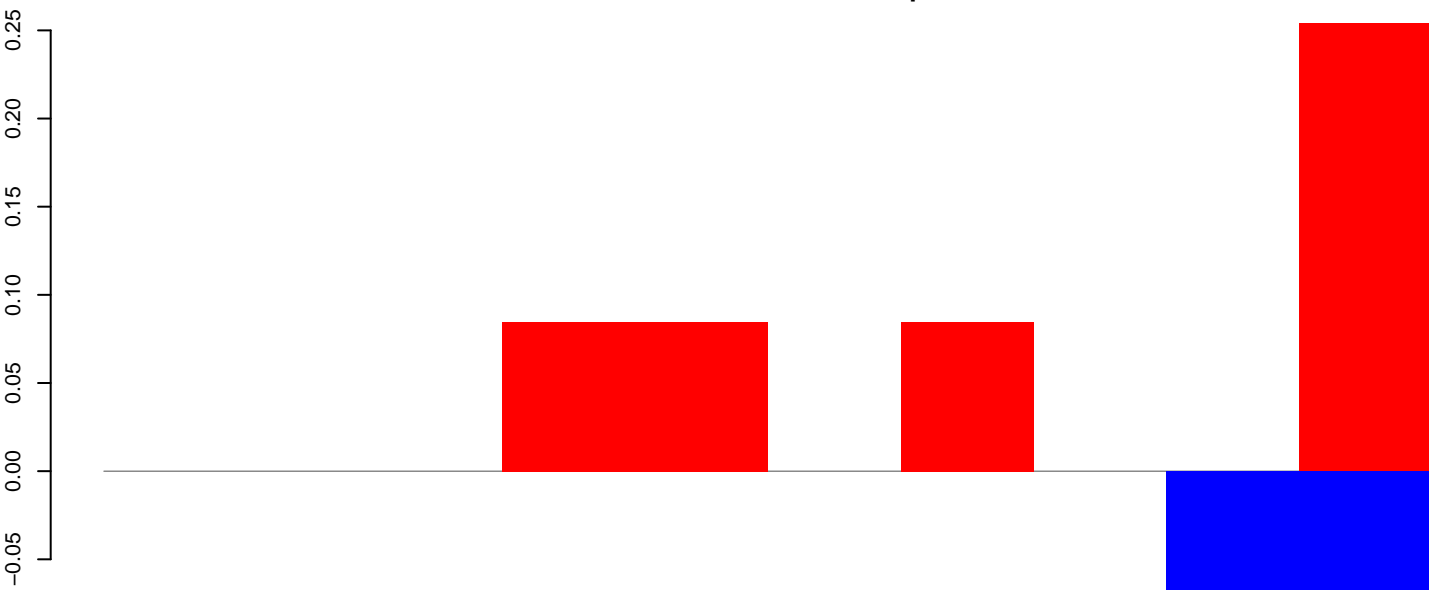
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=1, negative=0, total=1

AnGam_Mos55_cells.rep

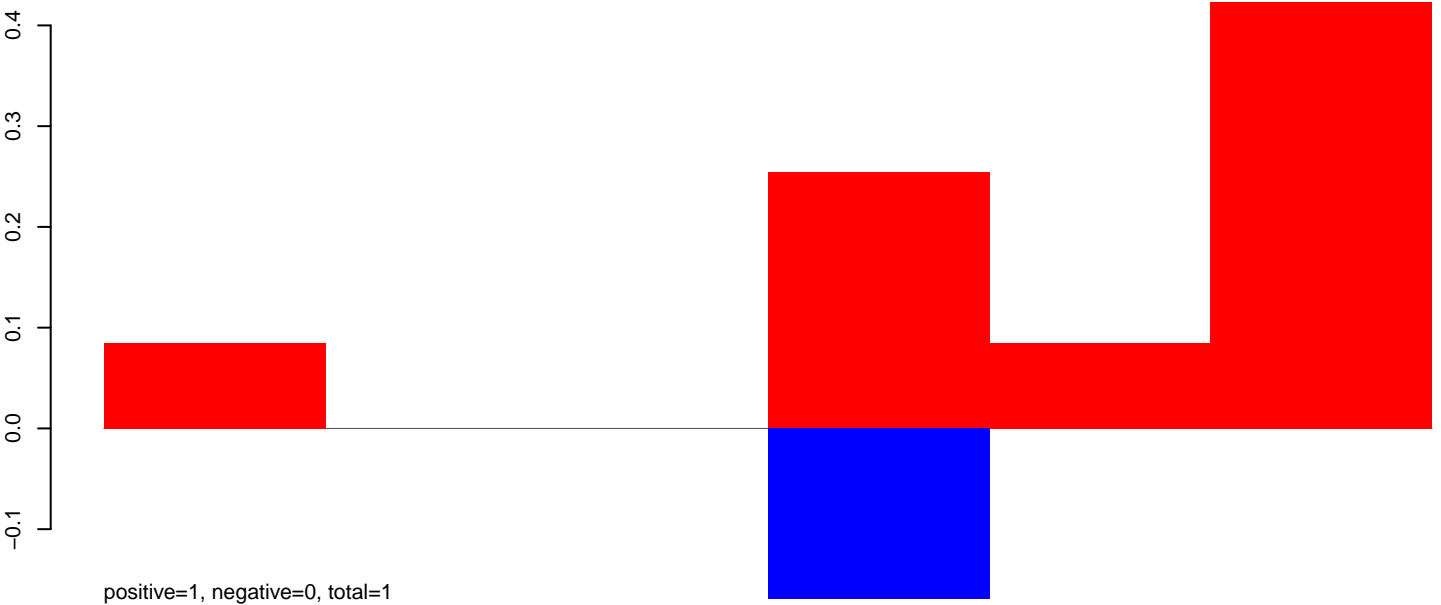


positive=1, negative=0, total=1

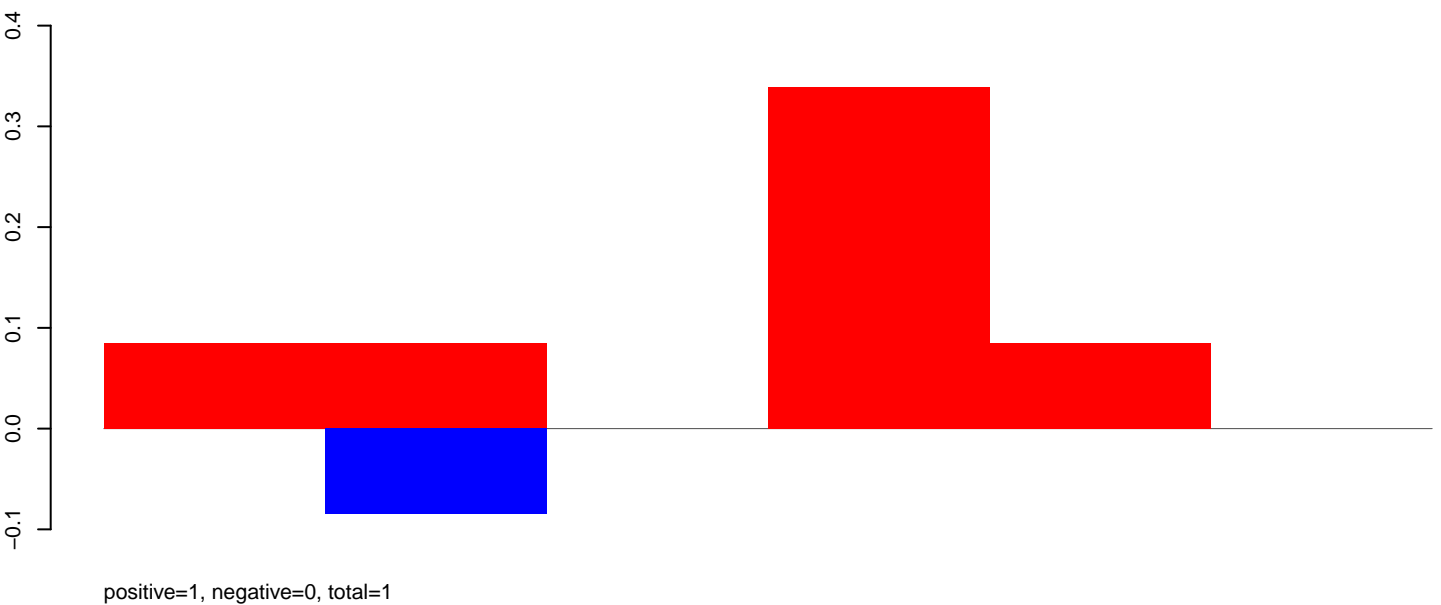
Window size=50, length=546, TE@TF003691-mTA_Ele4:1-546

100 200 300 400 500

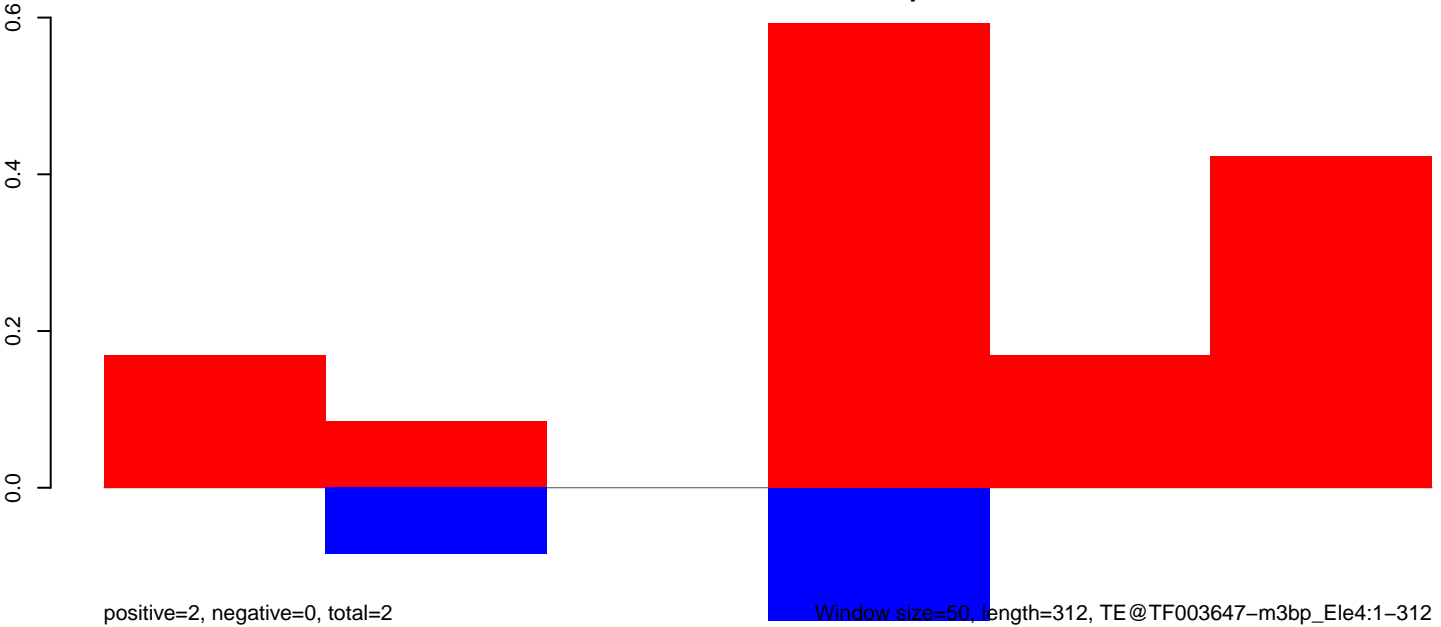
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



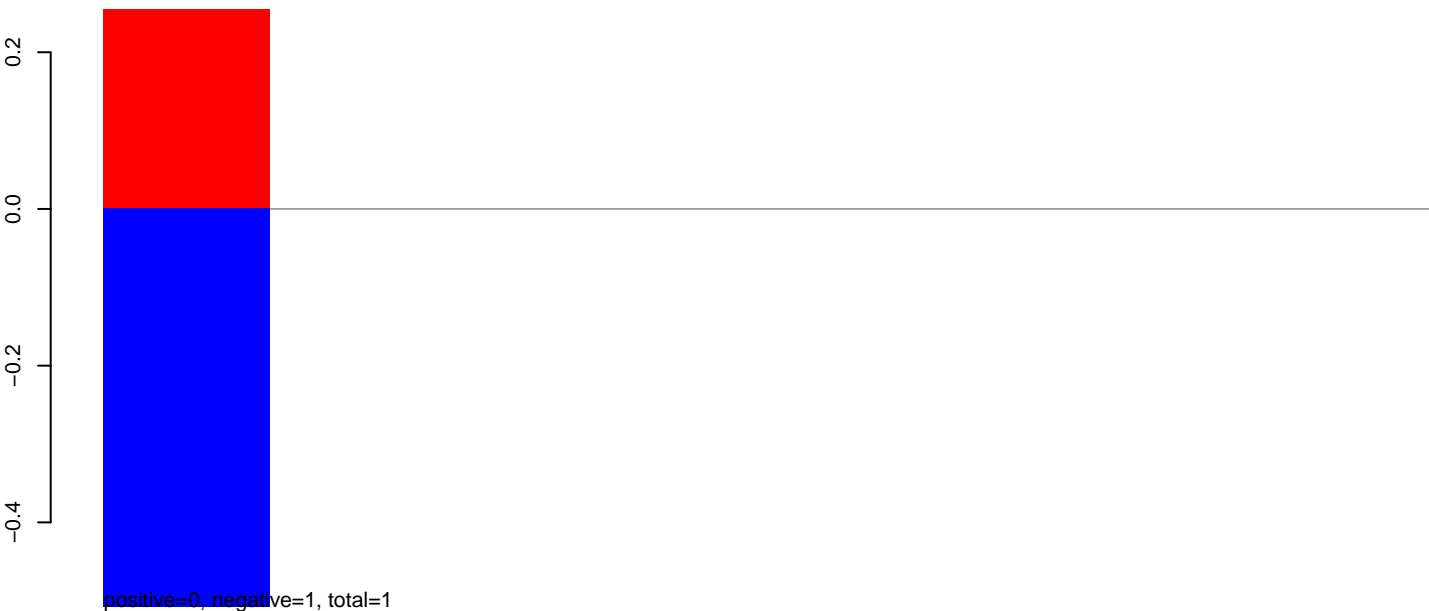
AnGam_Mos55_cells.rep



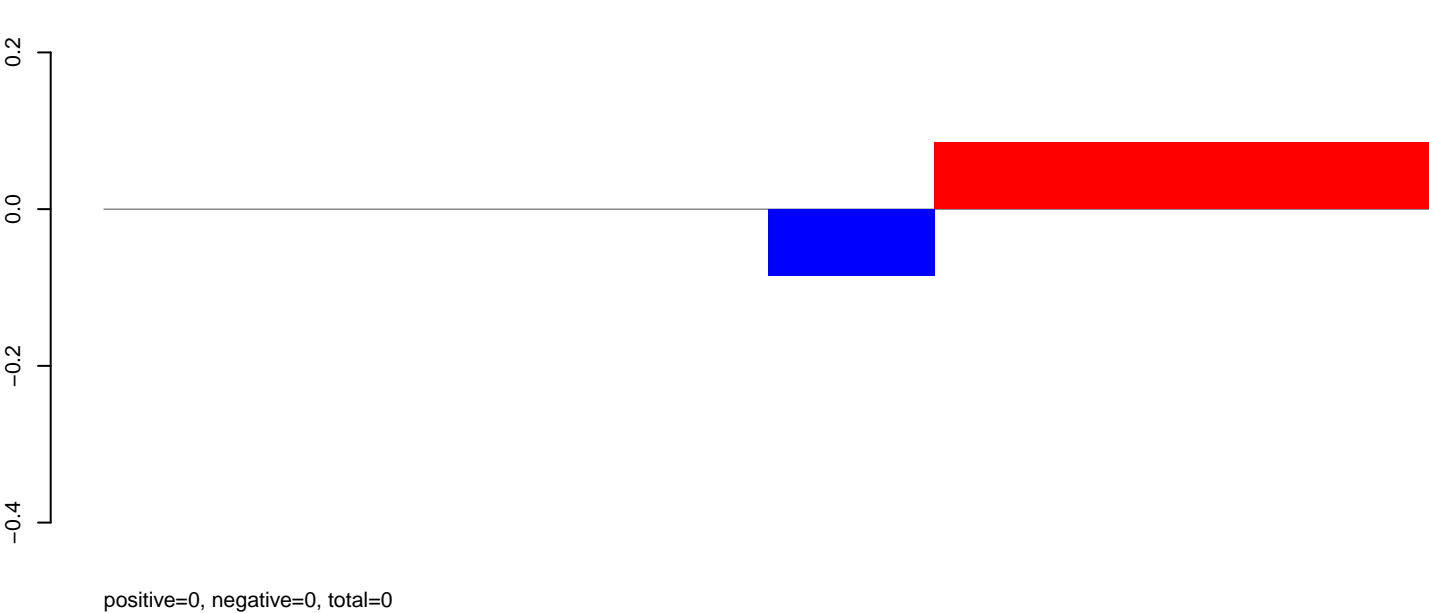
Window size=50, length=312, TE@TF003647-m3bp_Ele4:1-312

50 100 150 200 250 300 350

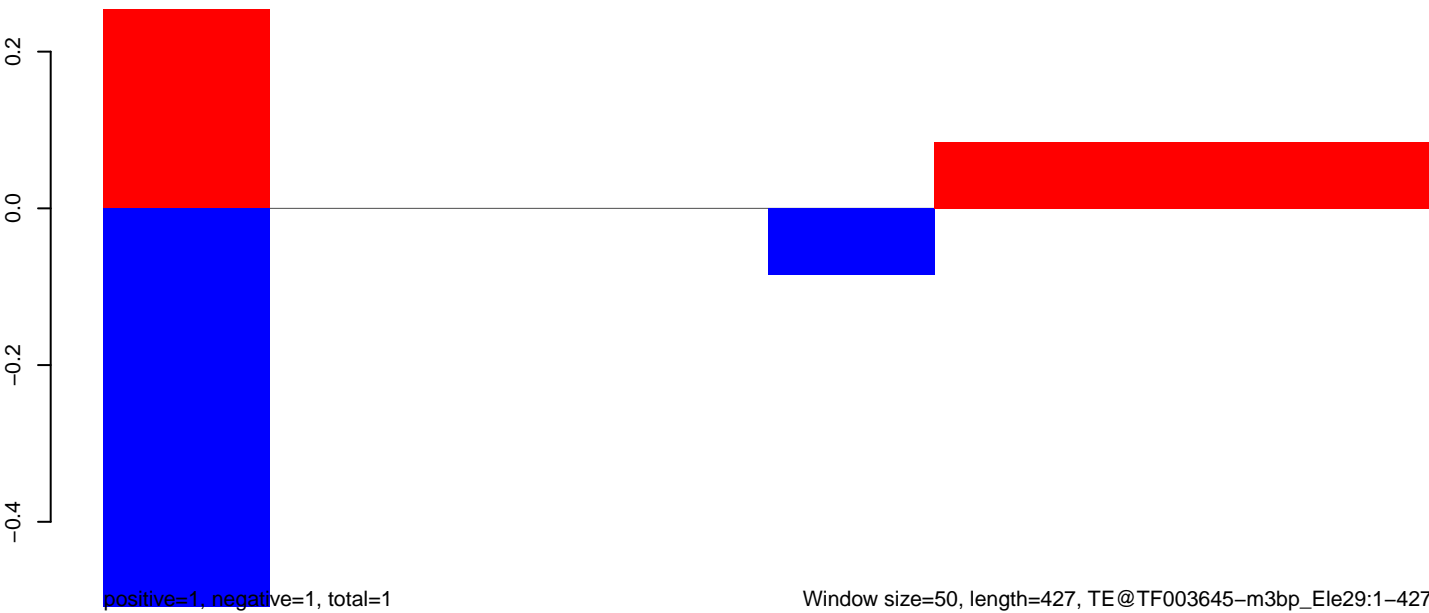
AnGam_Mos55_cells.18_23.rep



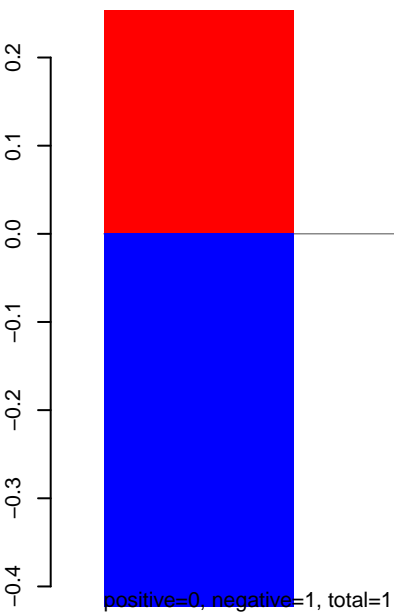
AnGam_Mos55_cells.24_35.rep



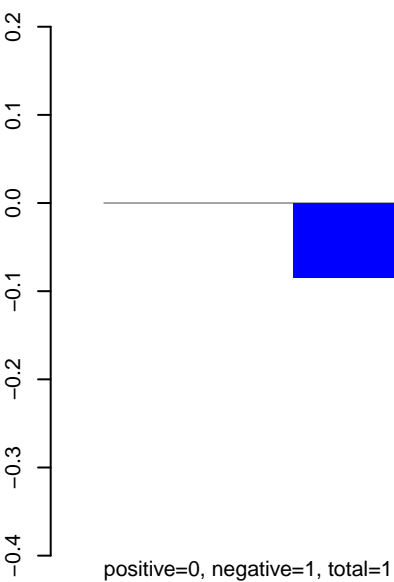
AnGam_Mos55_cells.rep



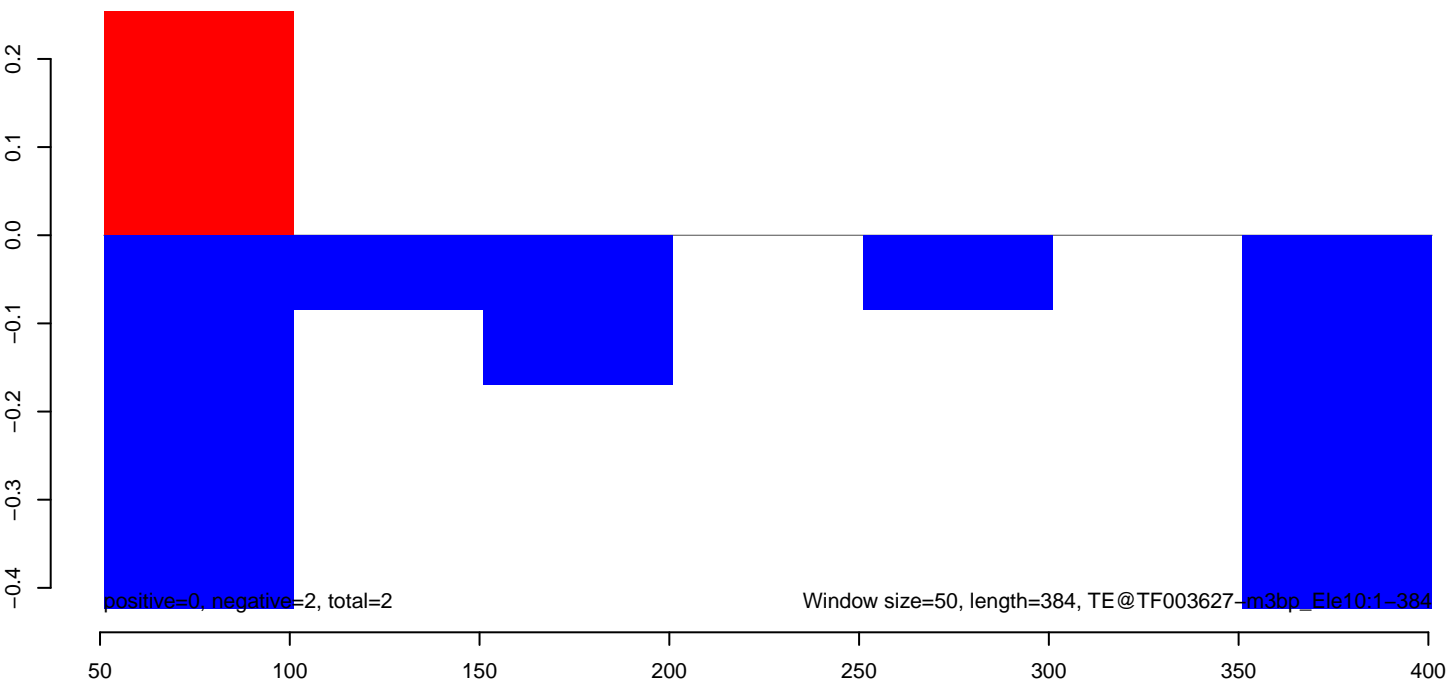
AnGam_Mos55_cells.18_23.rep



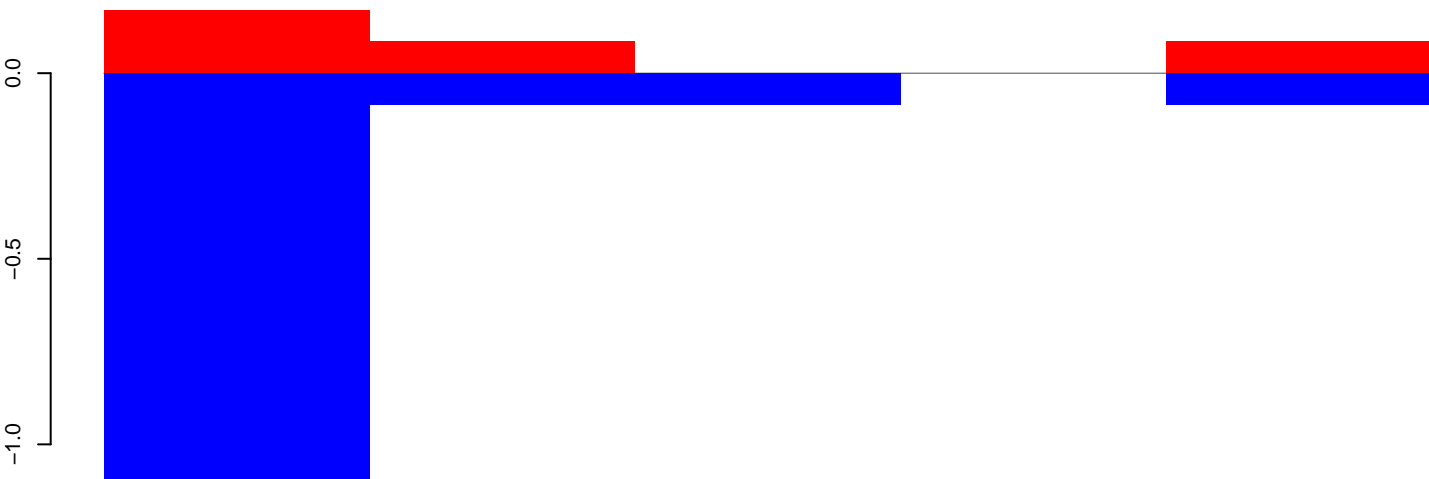
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

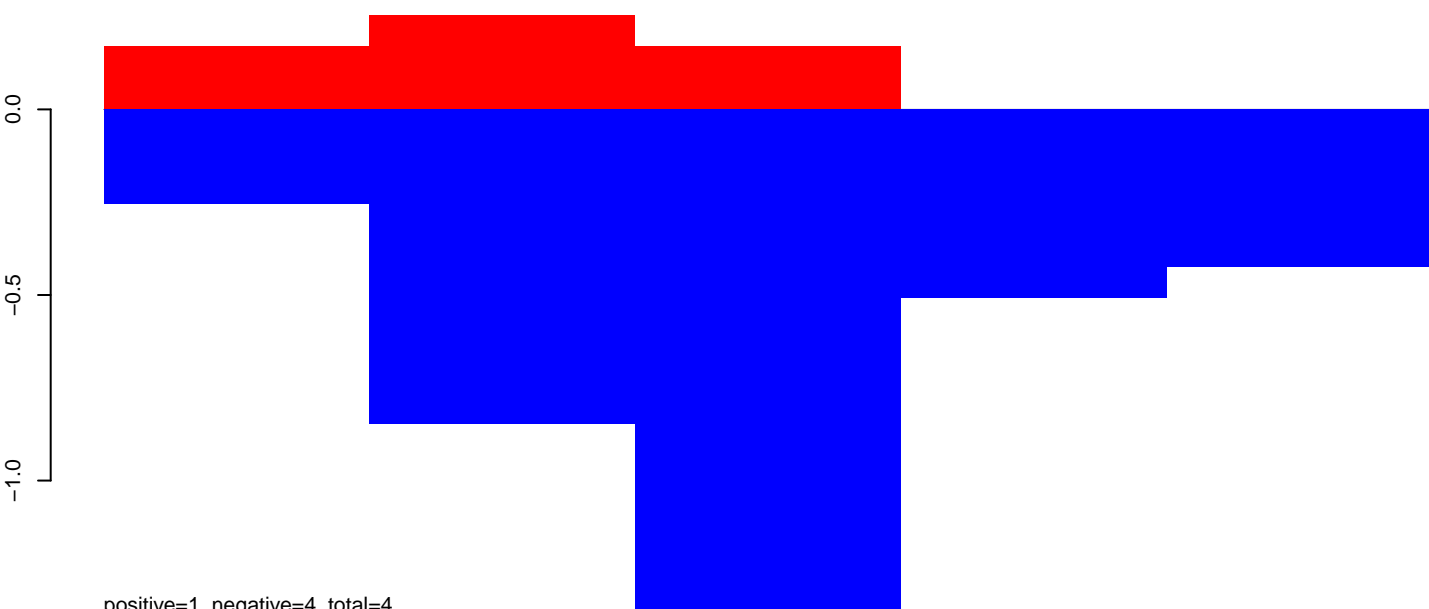


AnGam_Mos55_cells.18_23.rep



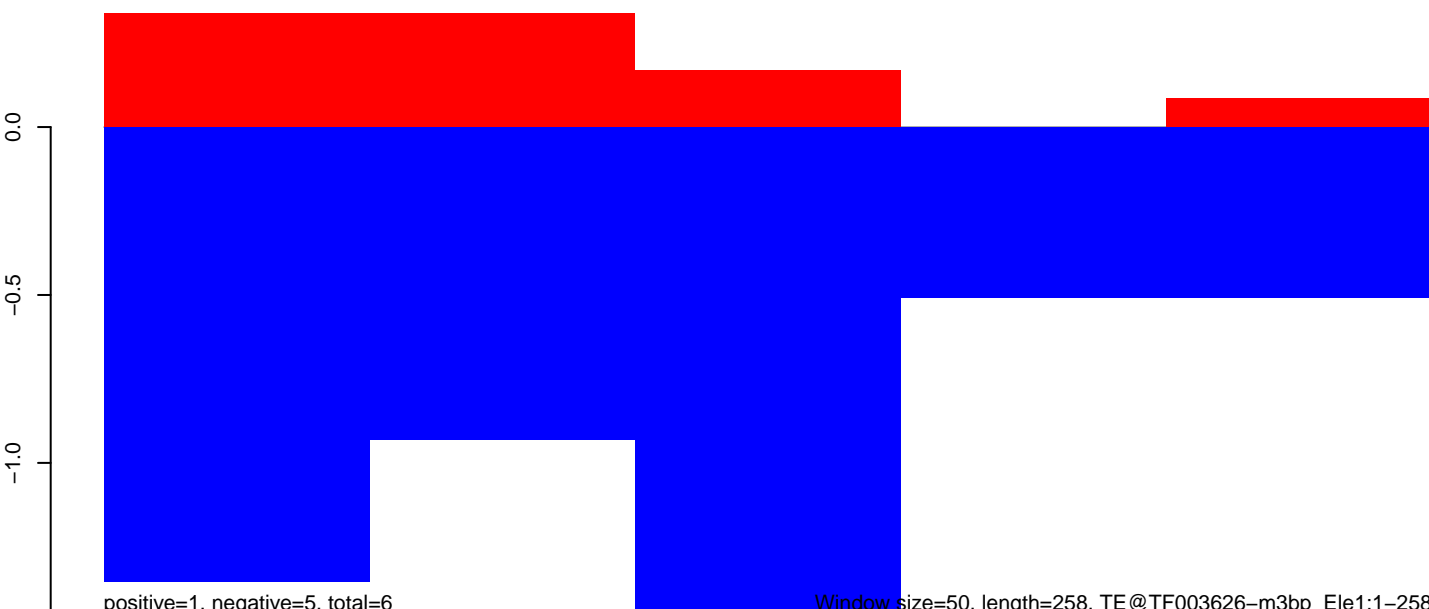
positive=0, negative=1, total=2

AnGam_Mos55_cells.24_35.rep



positive=1, negative=4, total=4

AnGam_Mos55_cells.rep



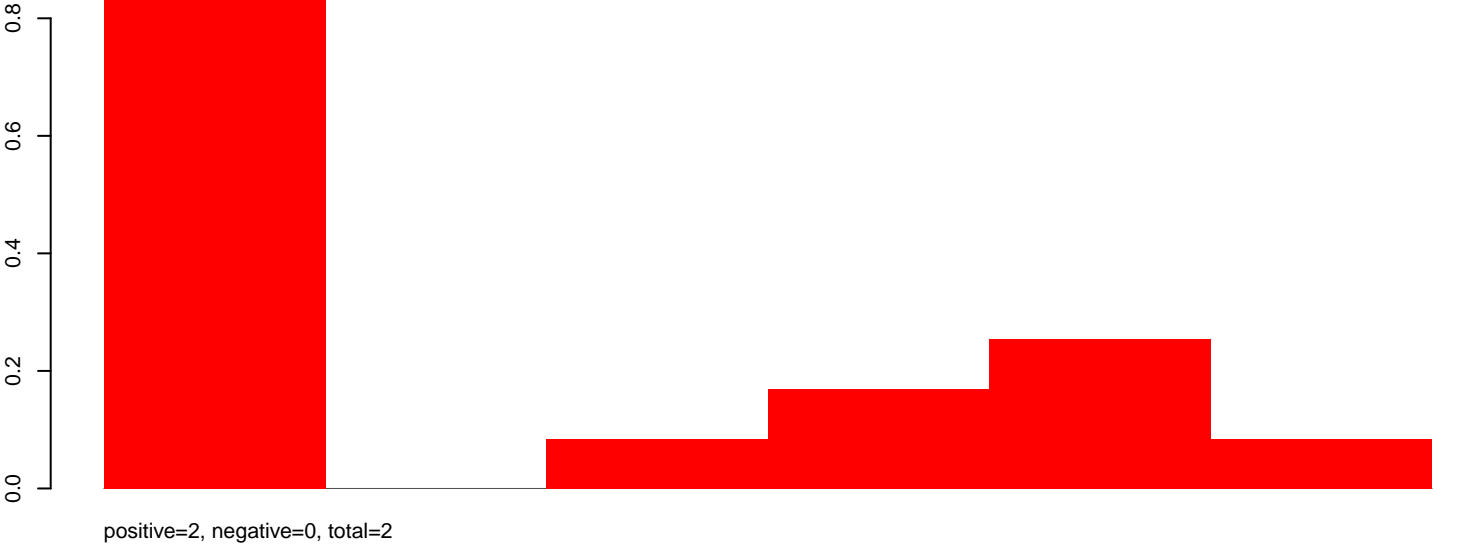
positive=1, negative=5, total=6

Window size=50, length=258, TE@TF003626-m3bp_Ele1:1-258

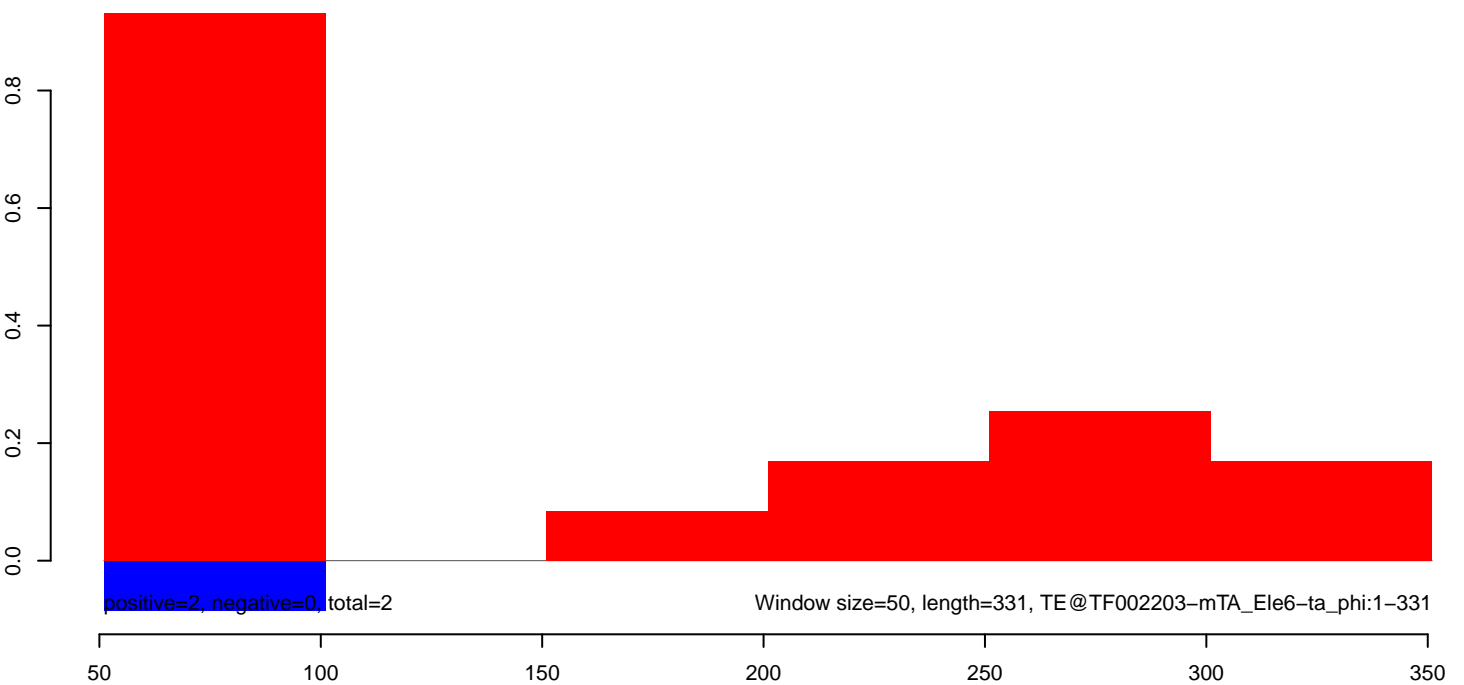
AnGam_Mos55_cells.18_23.rep



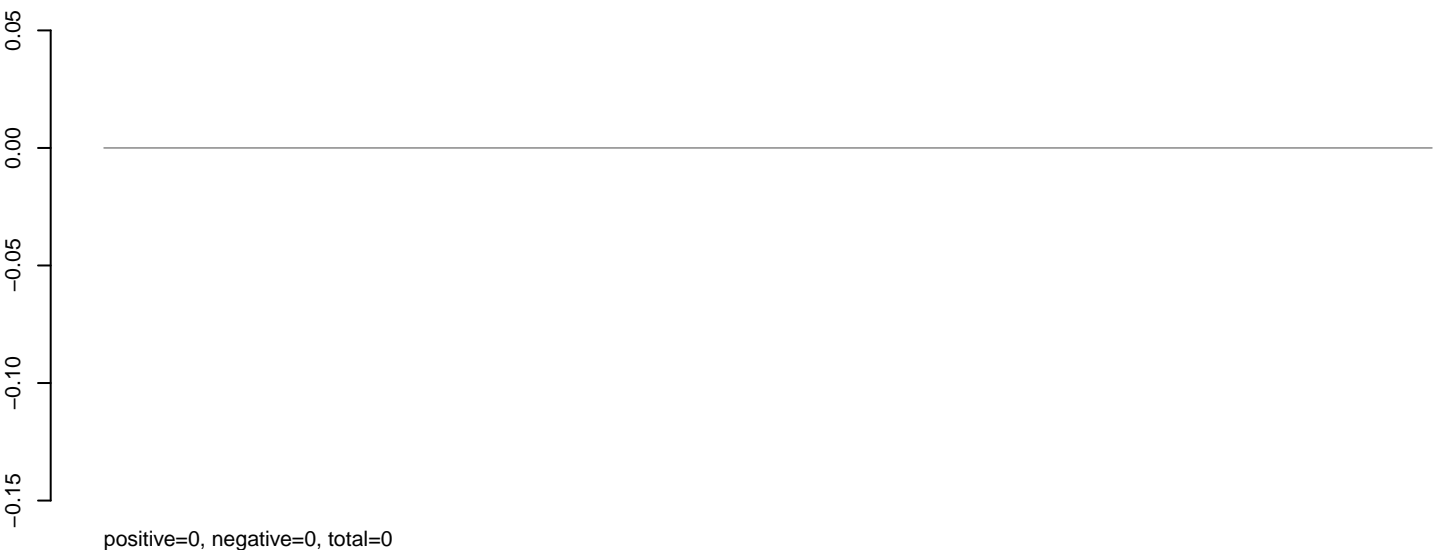
AnGam_Mos55_cells.24_35.rep



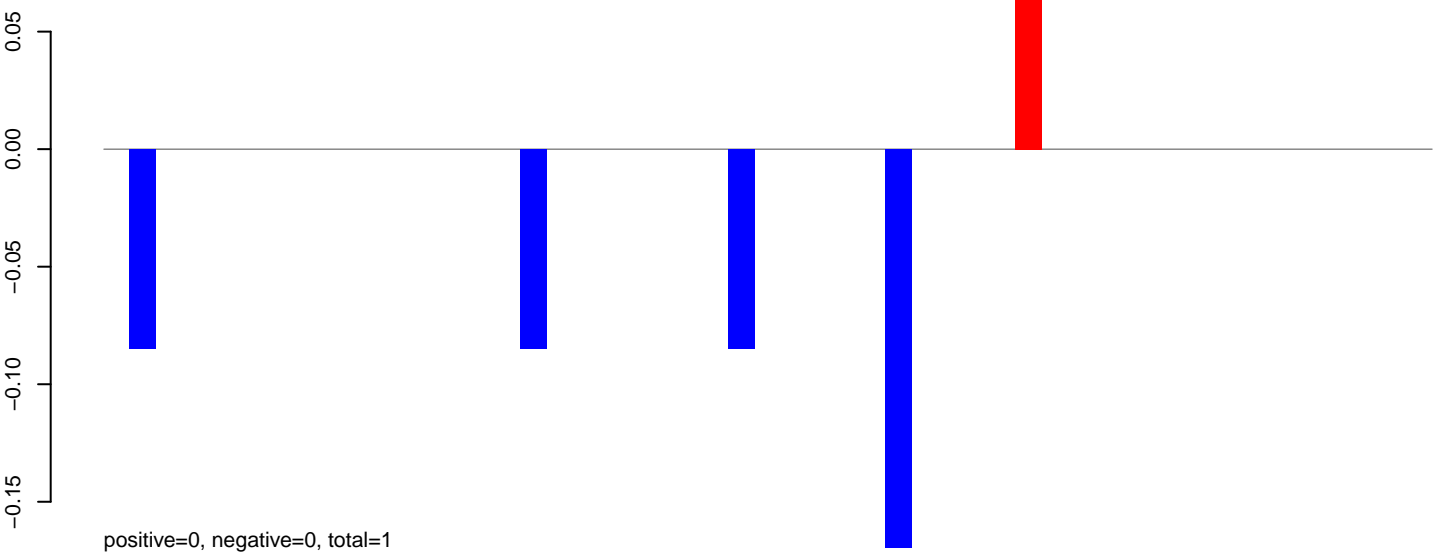
AnGam_Mos55_cells.rep



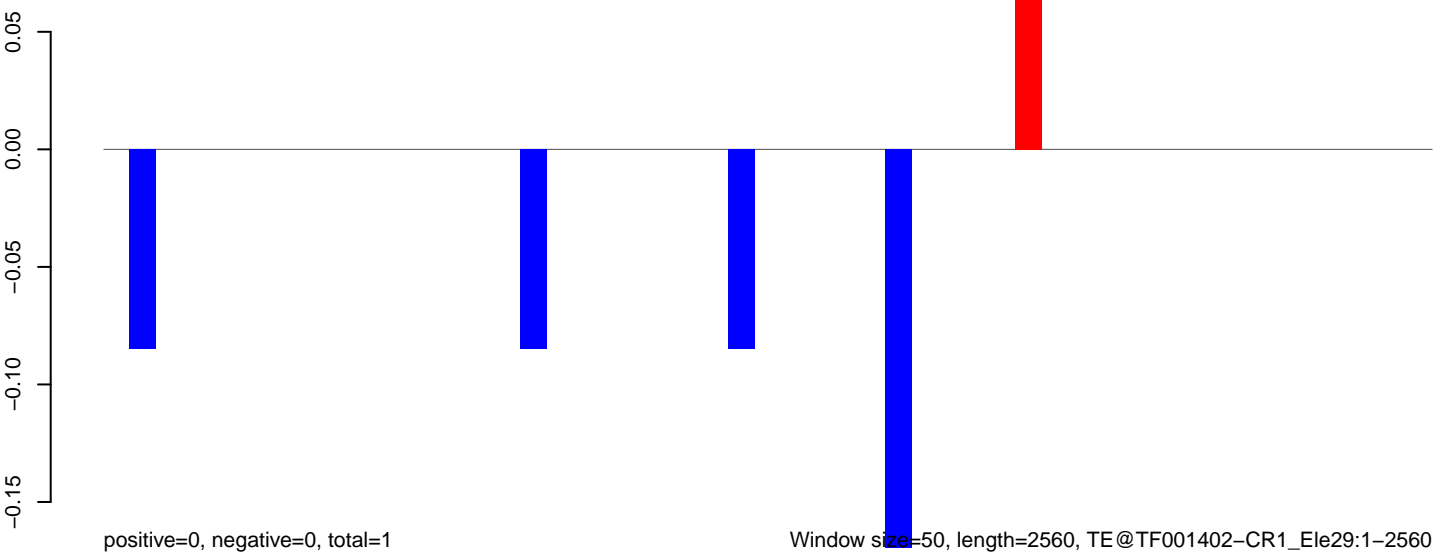
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

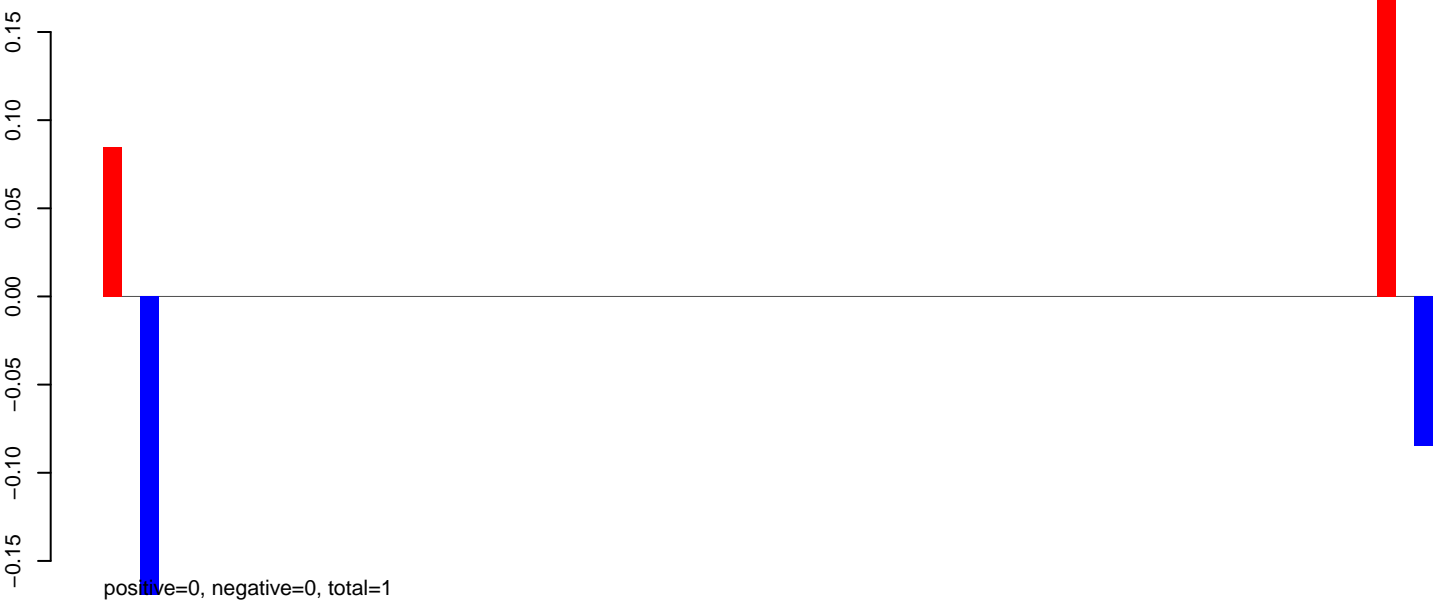


AnGam_Mos55_cells.rep

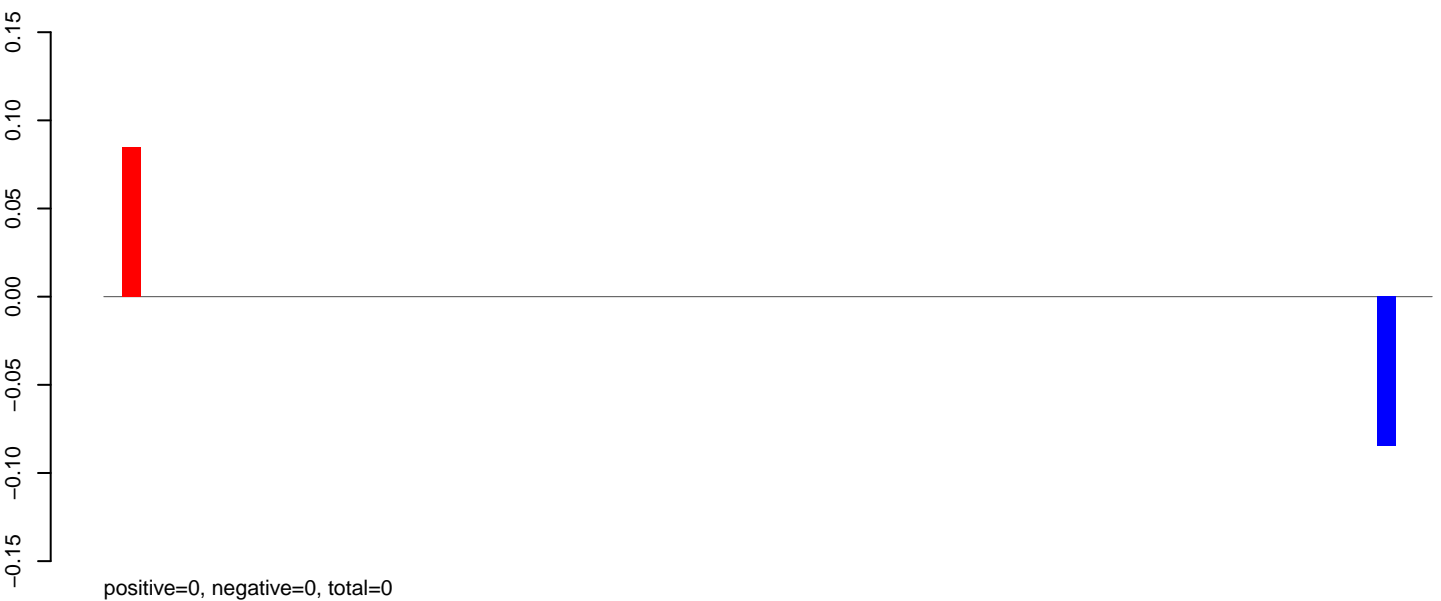


0 500 1000 1500 2000 2500

AnGam_Mos55_cells.18_23.rep



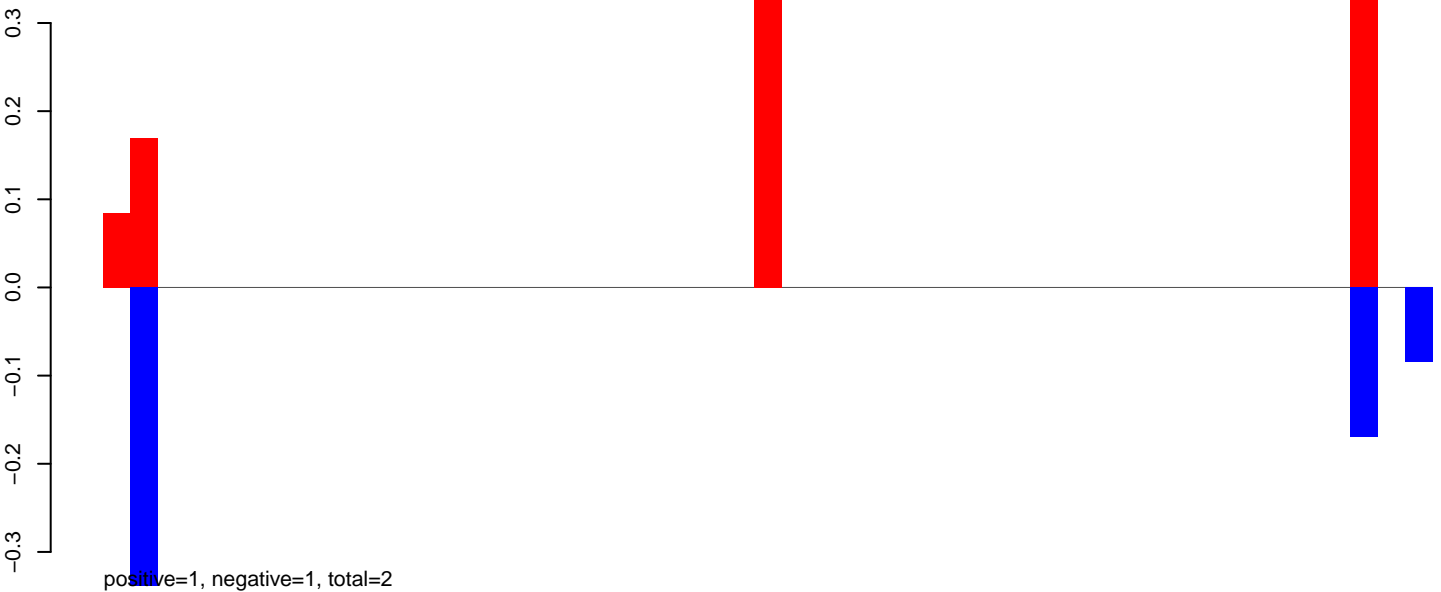
AnGam_Mos55_cells.24_35.rep



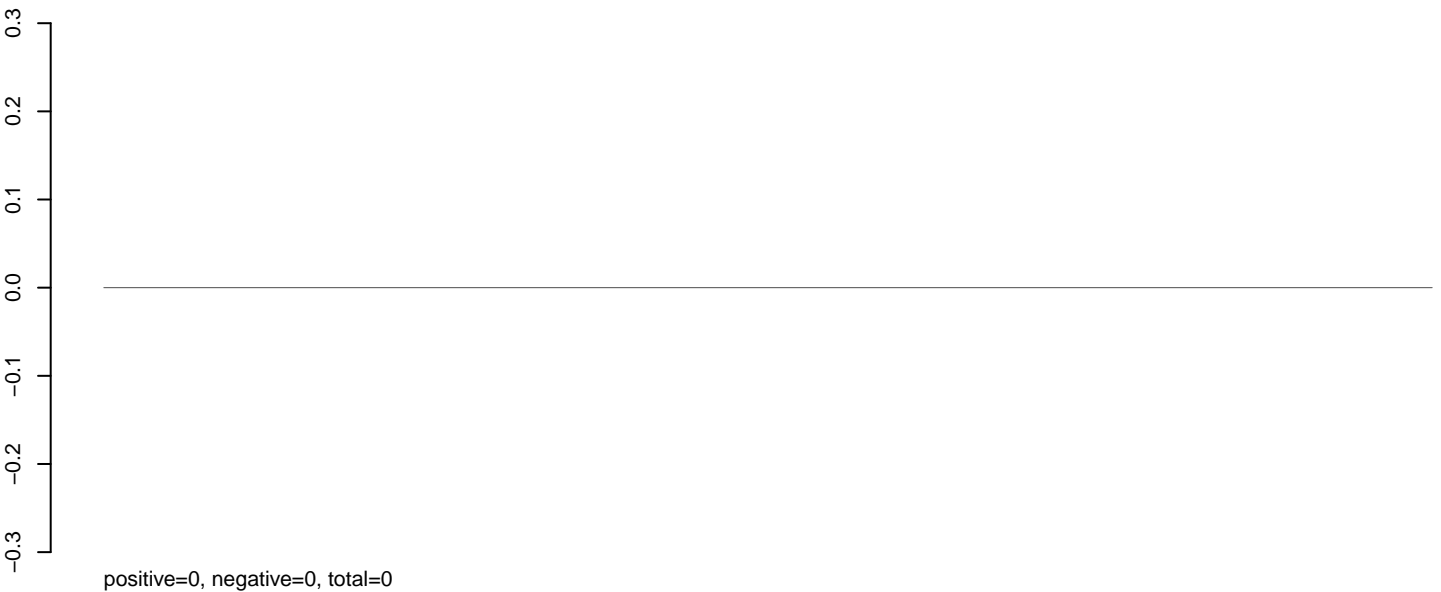
AnGam_Mos55_cells.rep



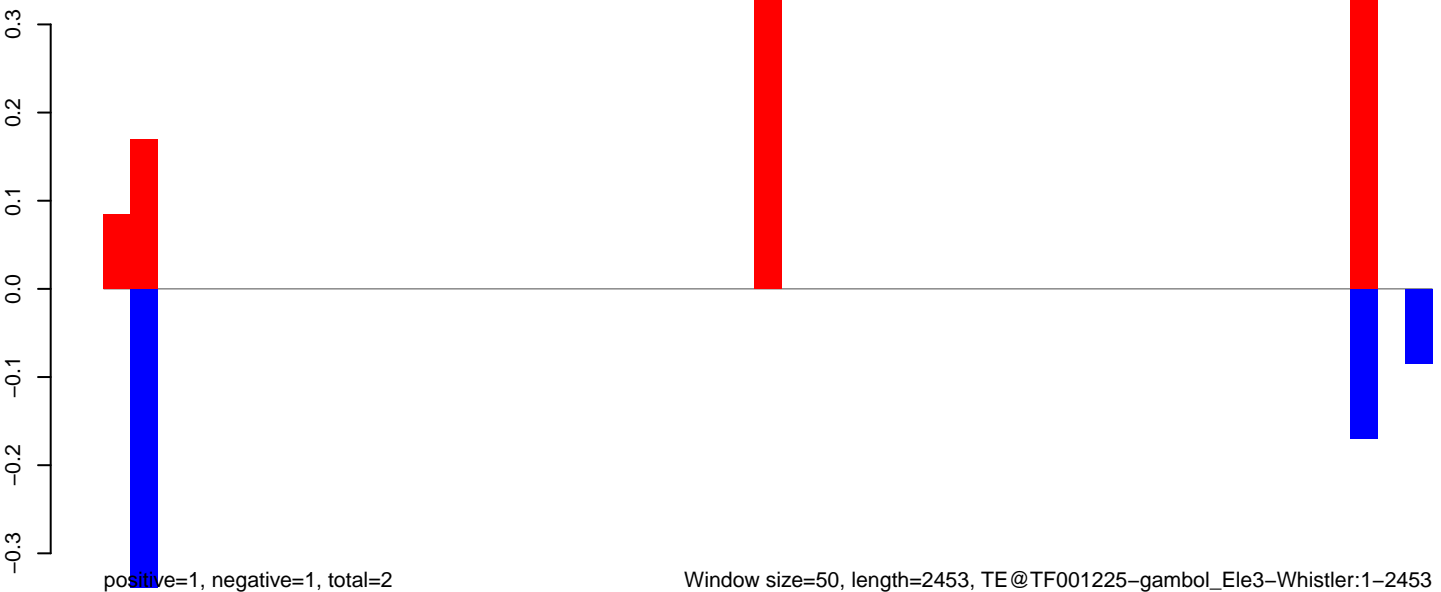
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

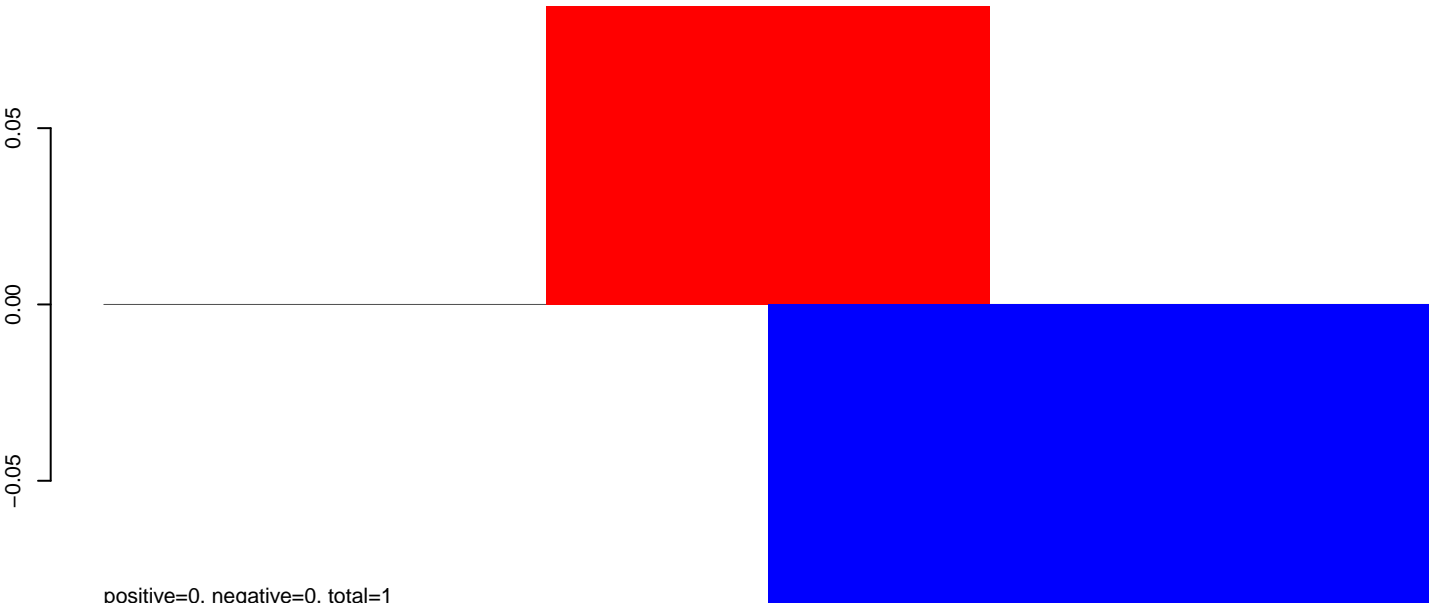


AnGam_Mos55_cells.rep



Window size=50, length=2453, TE@TF001225-gambol_Ele3-Whistler:1-2453

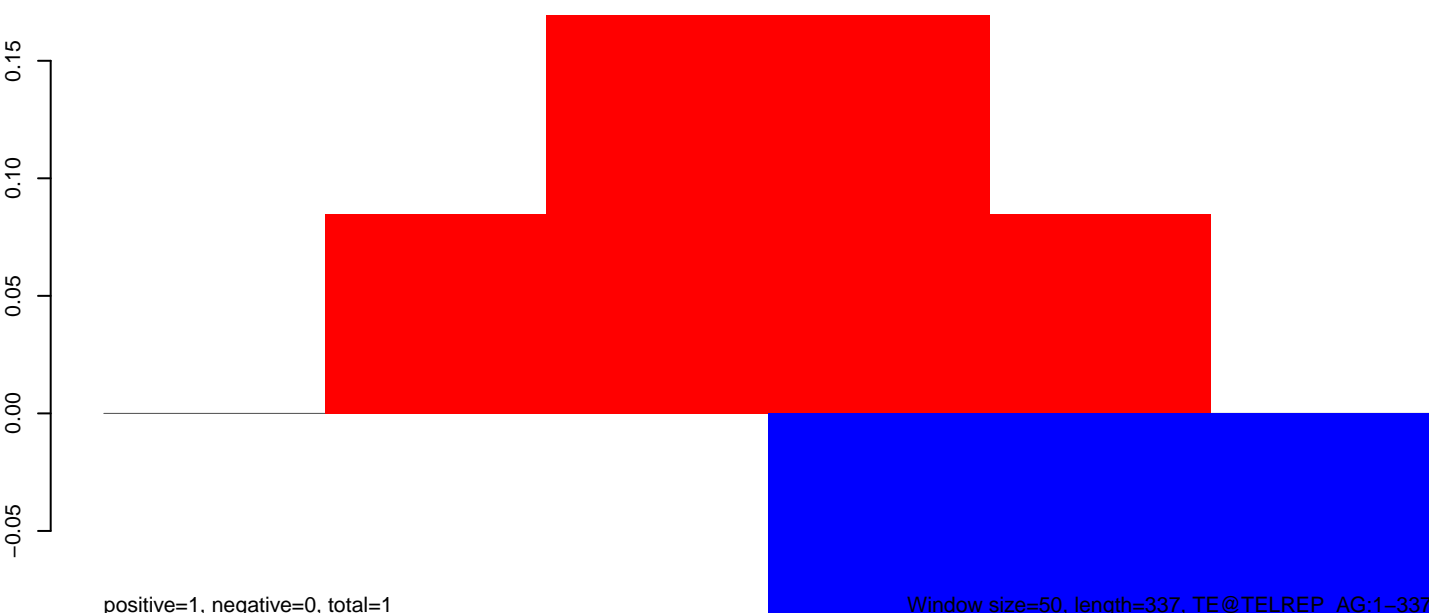
AnGam_Mos55_cells.18_23.rep



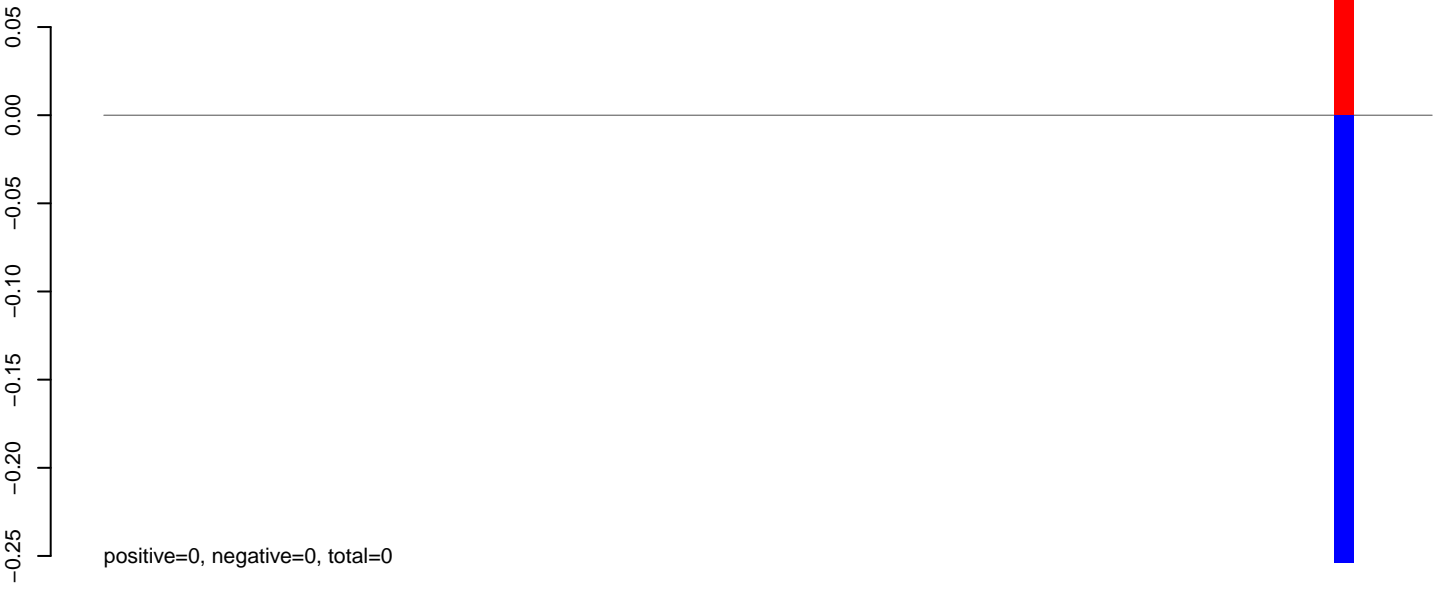
AnGam_Mos55_cells.24_35.rep



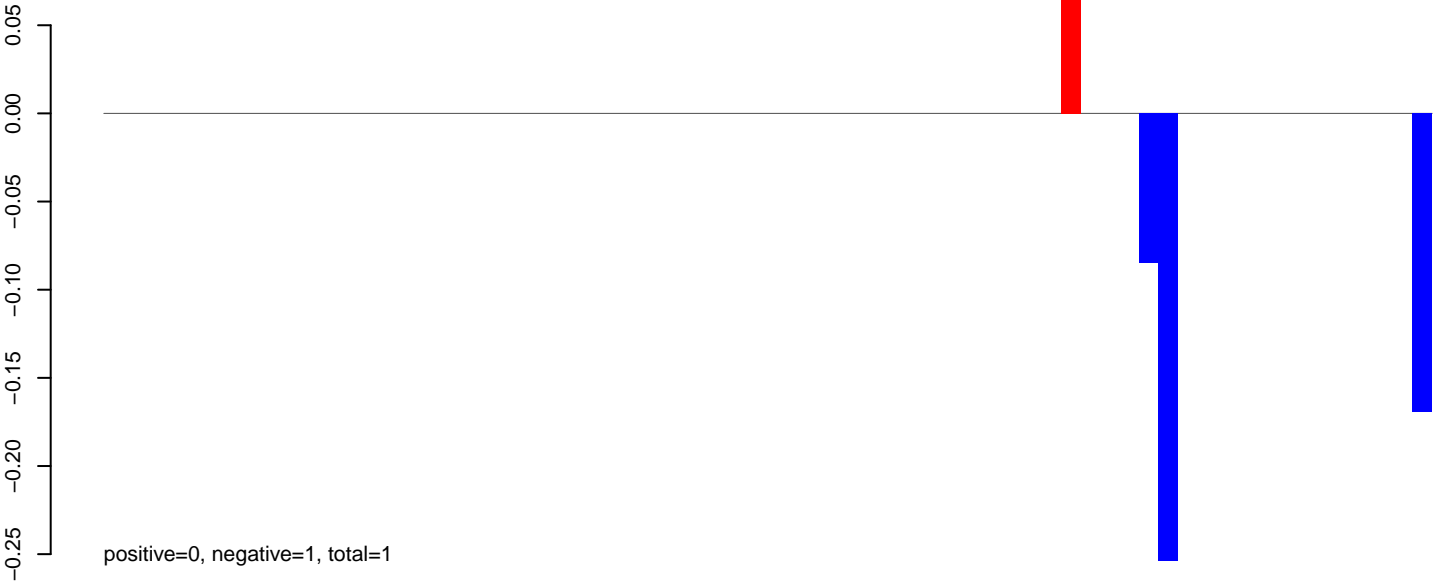
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



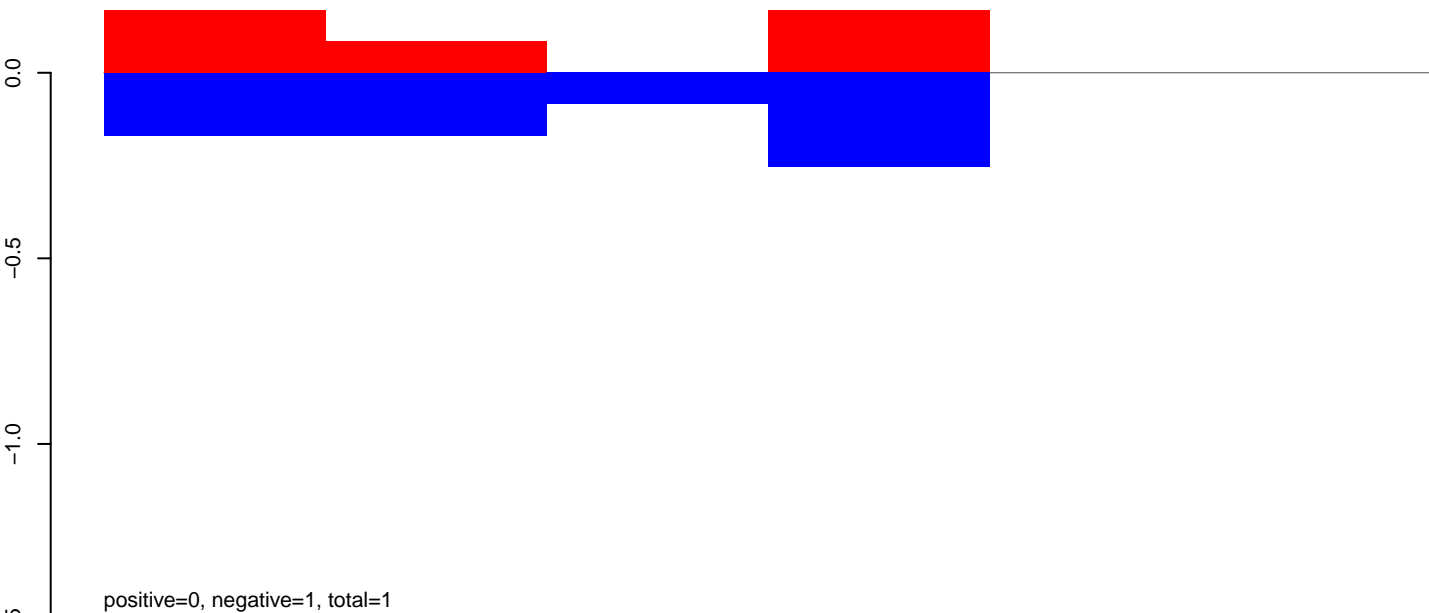
AnGam_Mos55_cells.24_35.rep



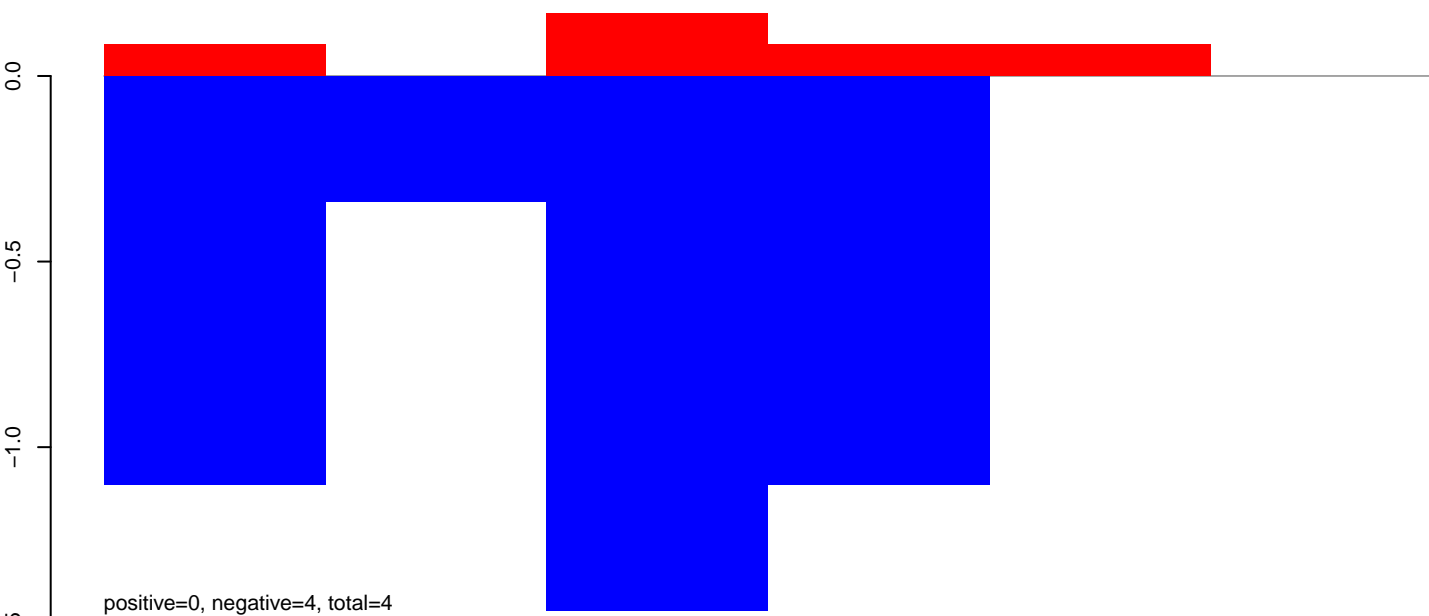
AnGam_Mos55_cells.rep



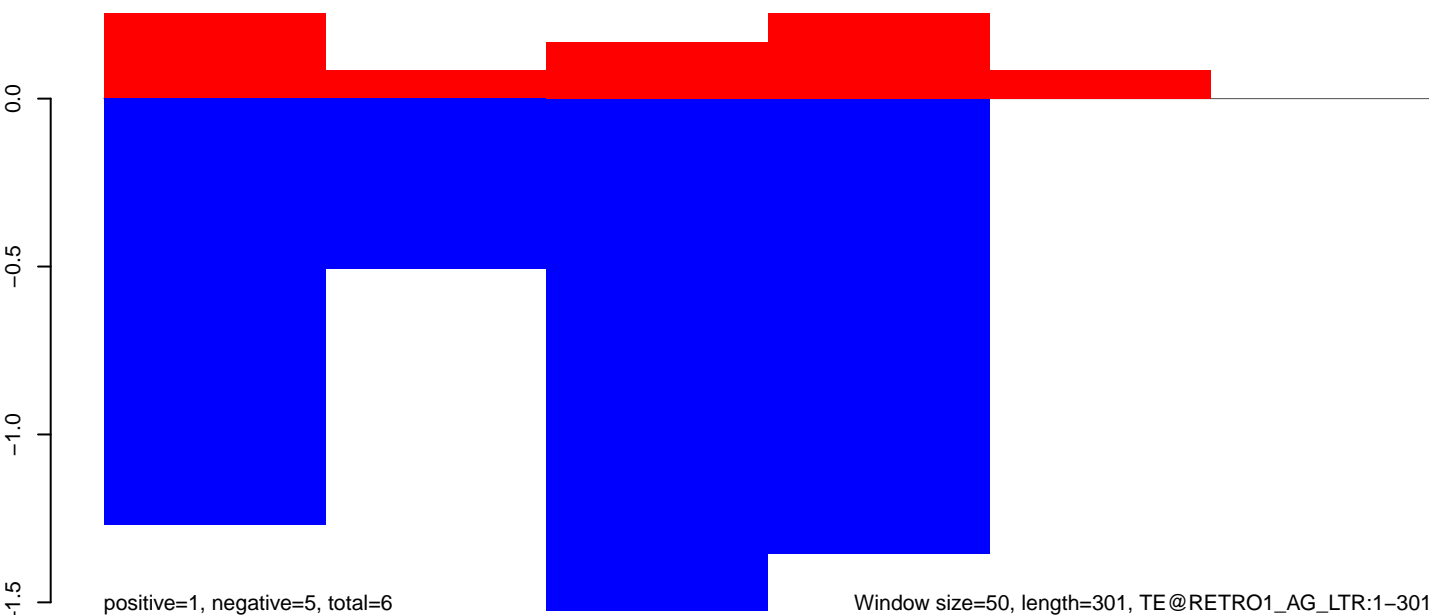
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



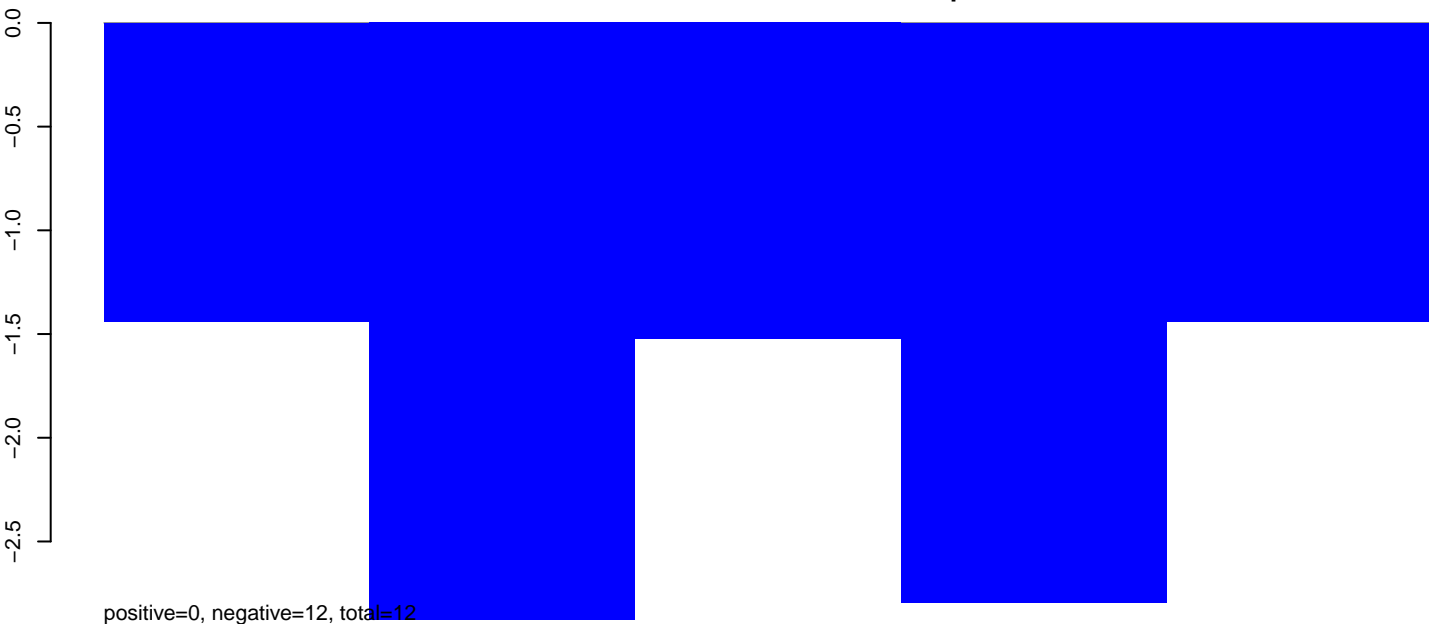
AnGam_Mos55_cells.rep



Window size=50, length=301, TE@RETRO1_AG_LTR:1-301

50 100 150 200 250 300 350

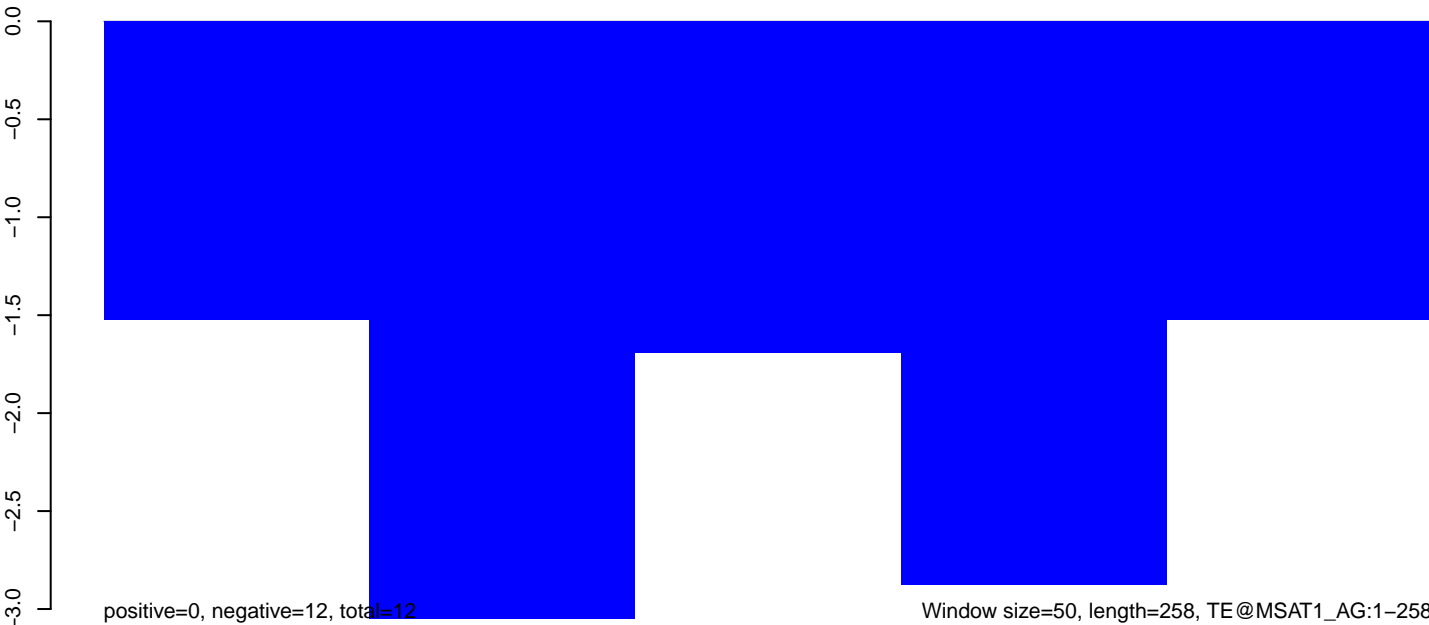
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

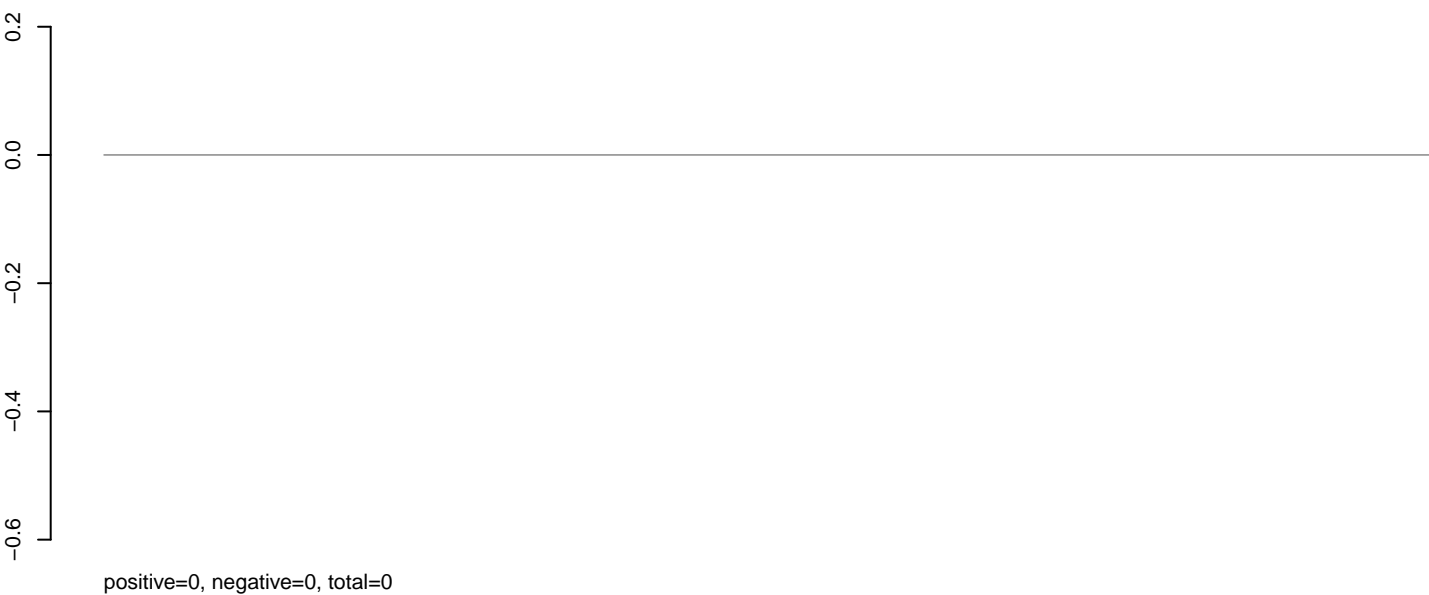


Window size=50, length=258, TE@MSAT1_AG:1-258

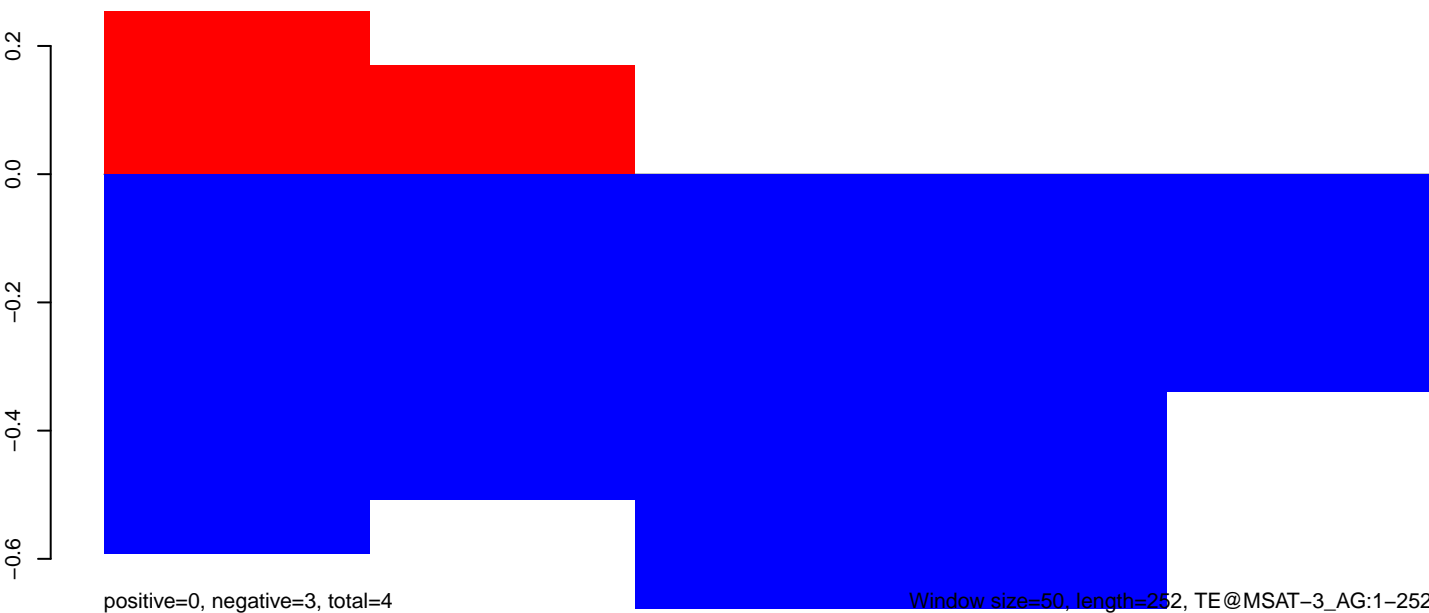
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



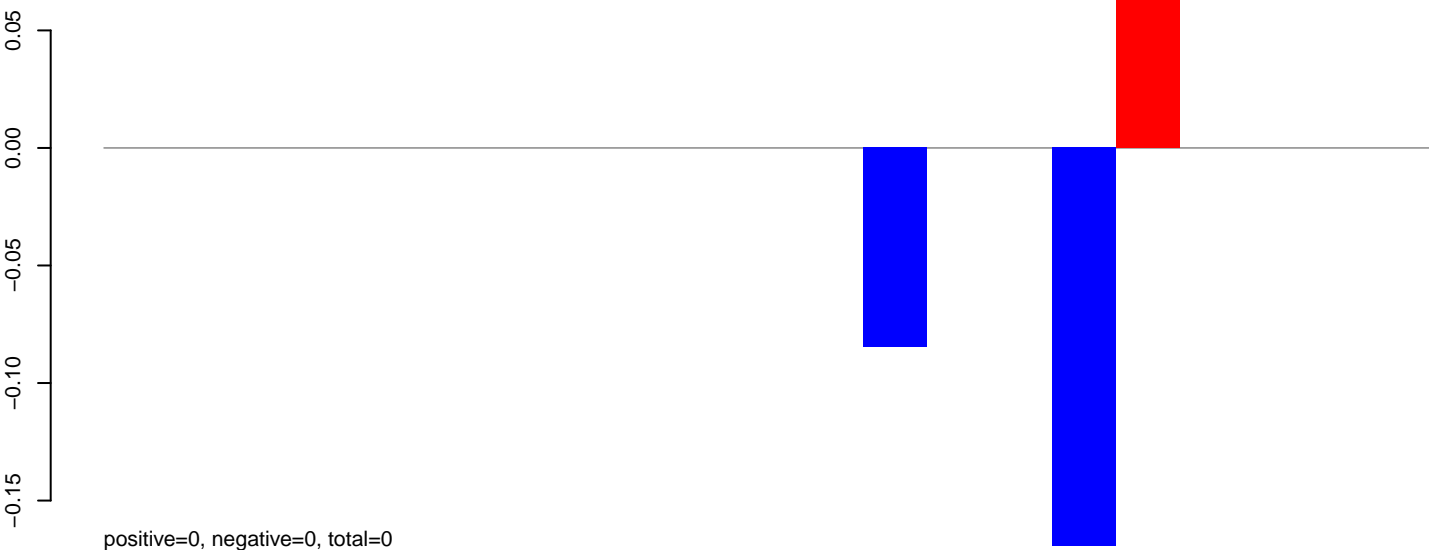
AnGam_Mos55_cells.rep



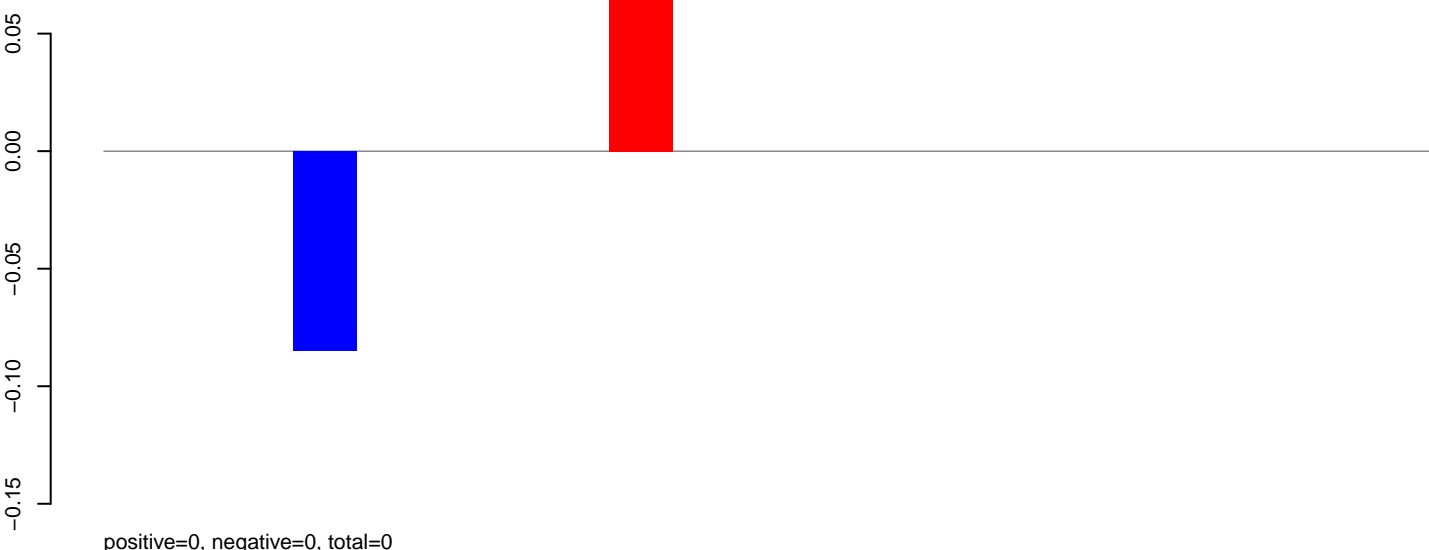
Window size=50, length=252, TE@MSAT-3_AG:1-252

50 100 150 200 250 300

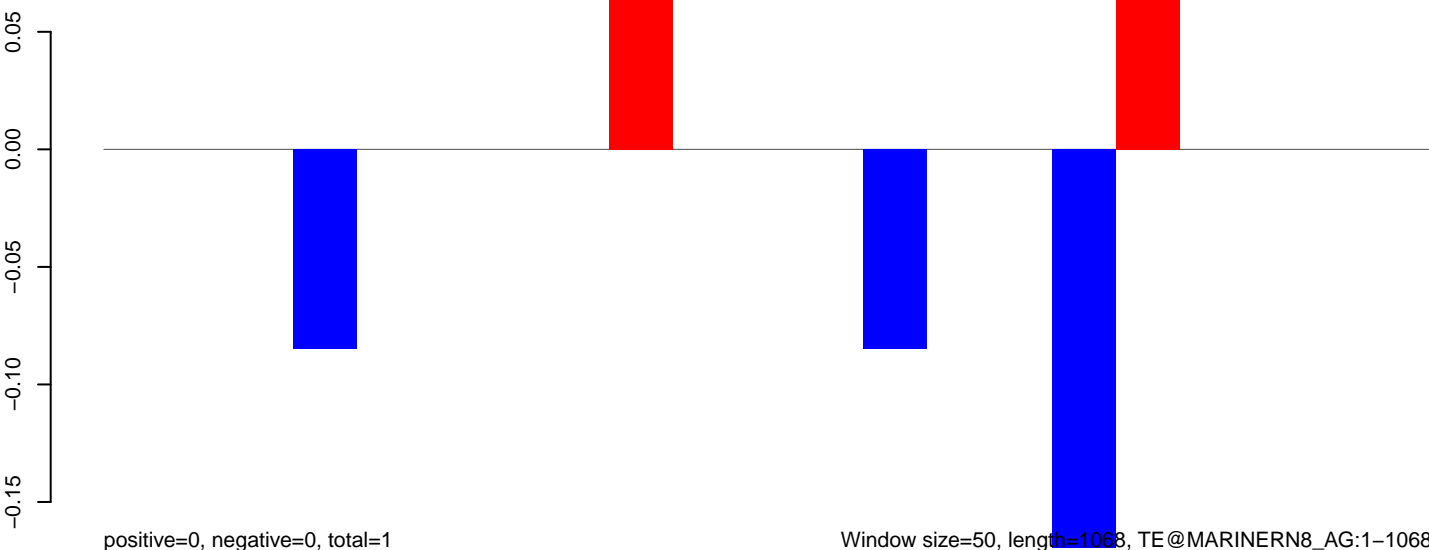
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



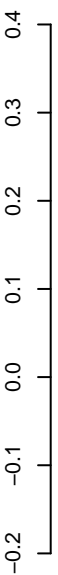
AnGam_Mos55_cells.rep



Window size=50, length=1068, TE@MARINERN8_AG:1-1068

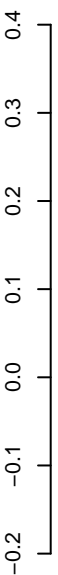
200 400 600 800 1000

AnGam_Mos55_cells.18_23.rep



positive=1, negative=1, total=2

AnGam_Mos55_cells.24_35.rep



positive=1, negative=0, total=1

AnGam_Mos55_cells.rep

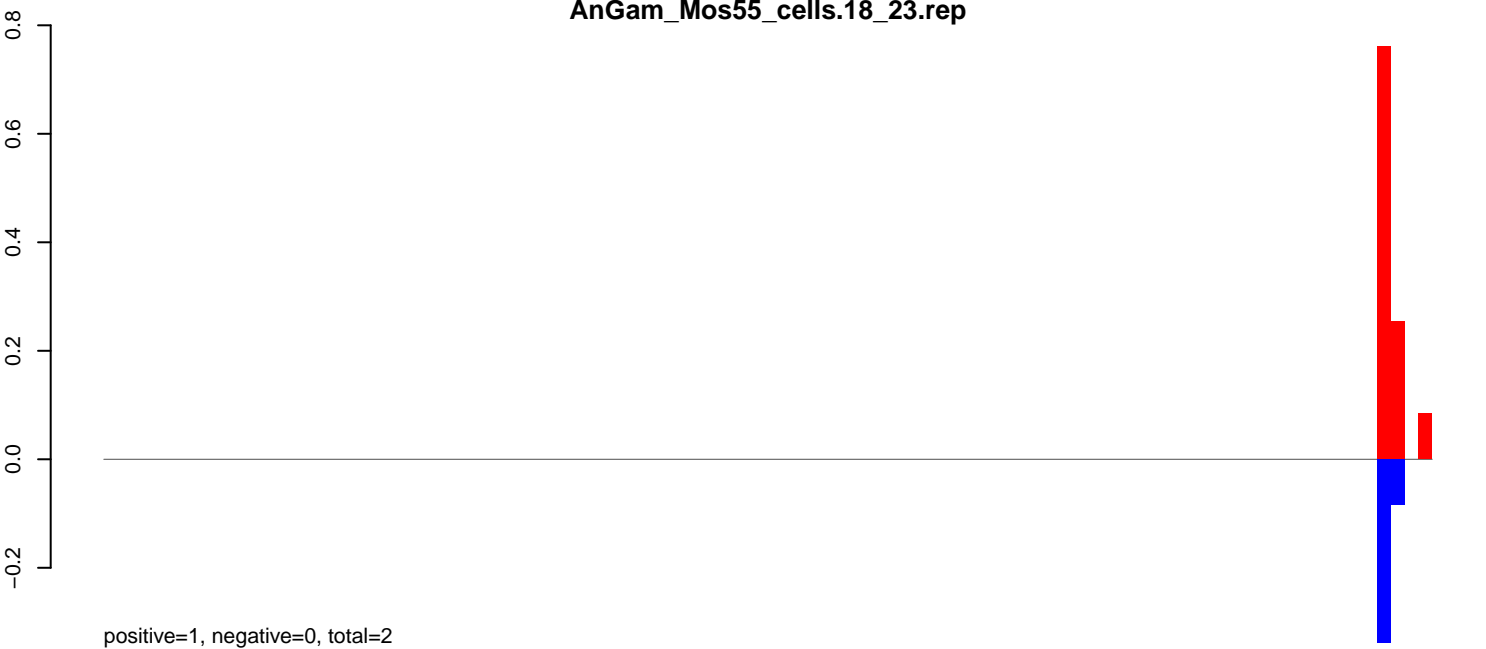


positive=2, negative=1, total=2

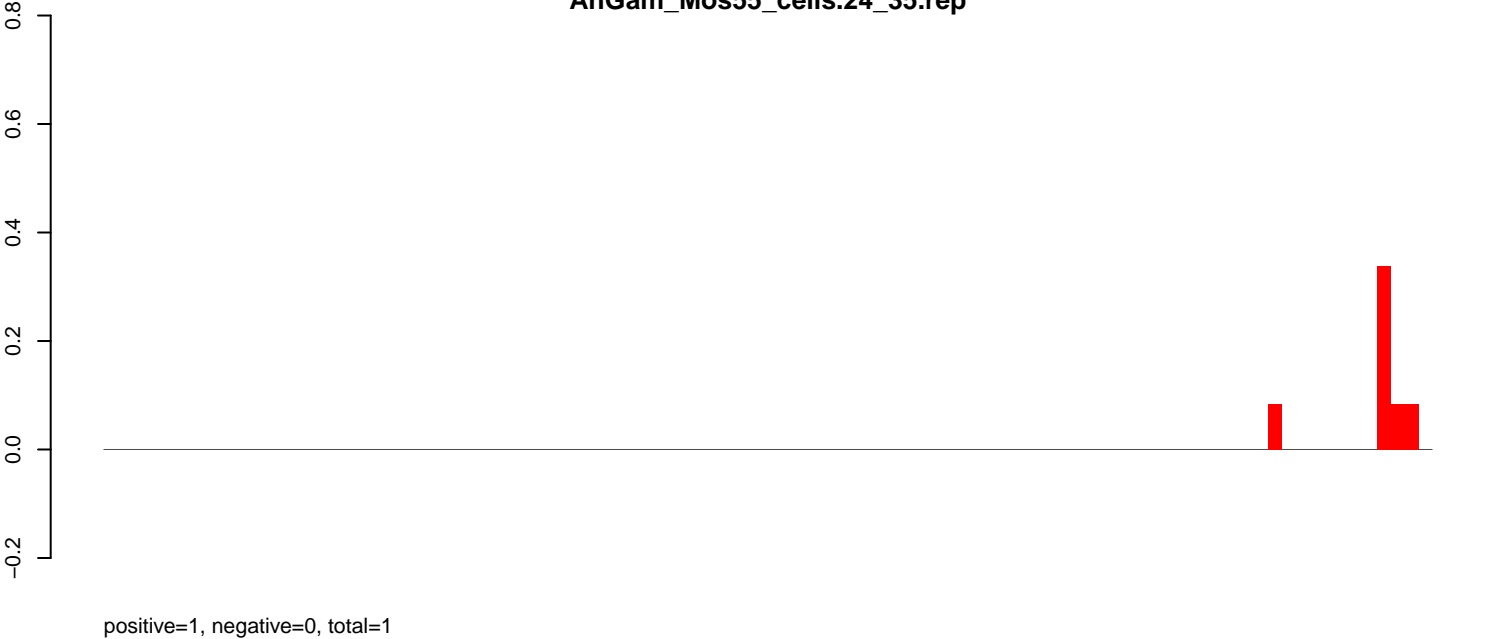
Window size=50, length=270, TE@MARINERN10_AG:1-270



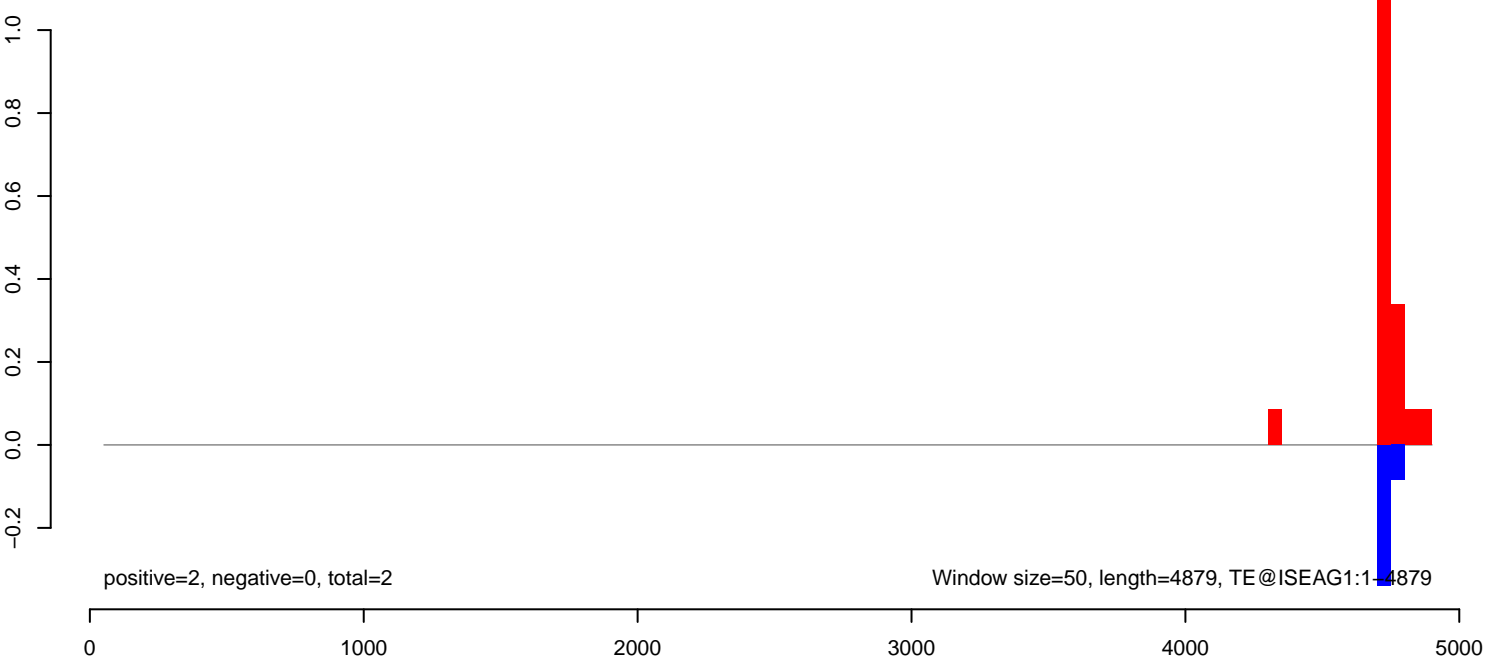
AnGam_Mos55_cells.18_23.rep



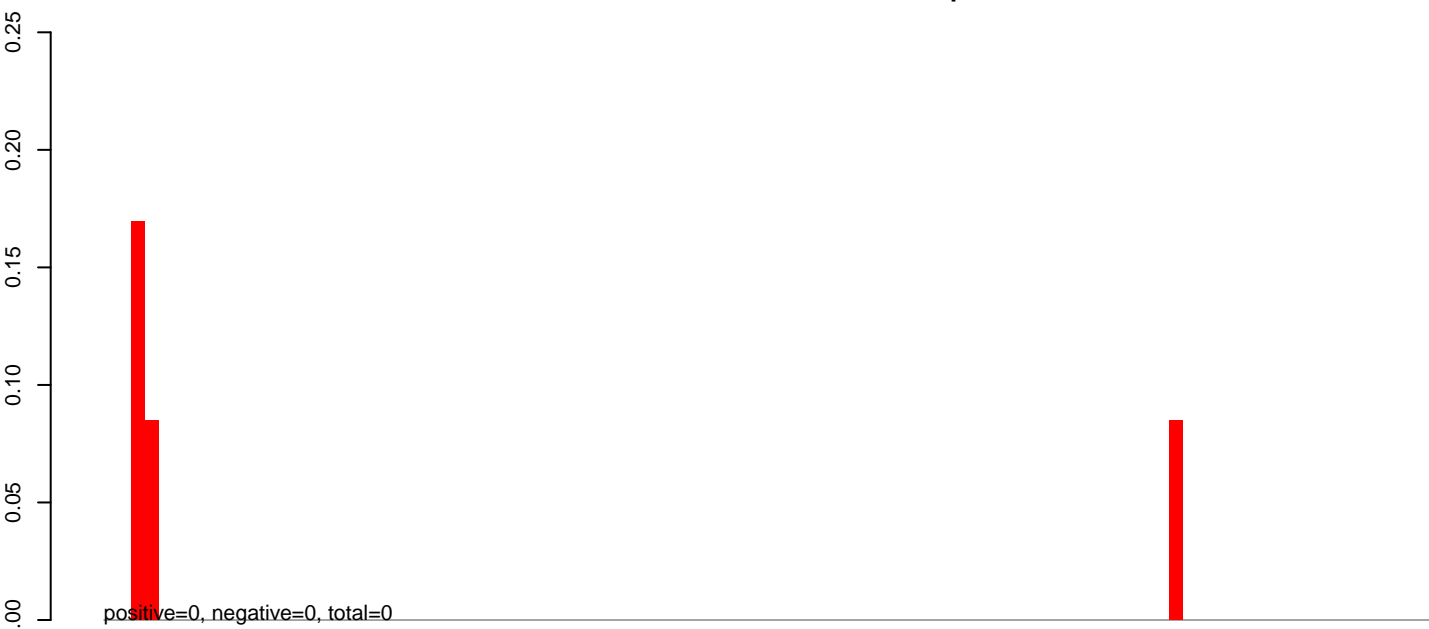
AnGam_Mos55_cells.24_35.rep



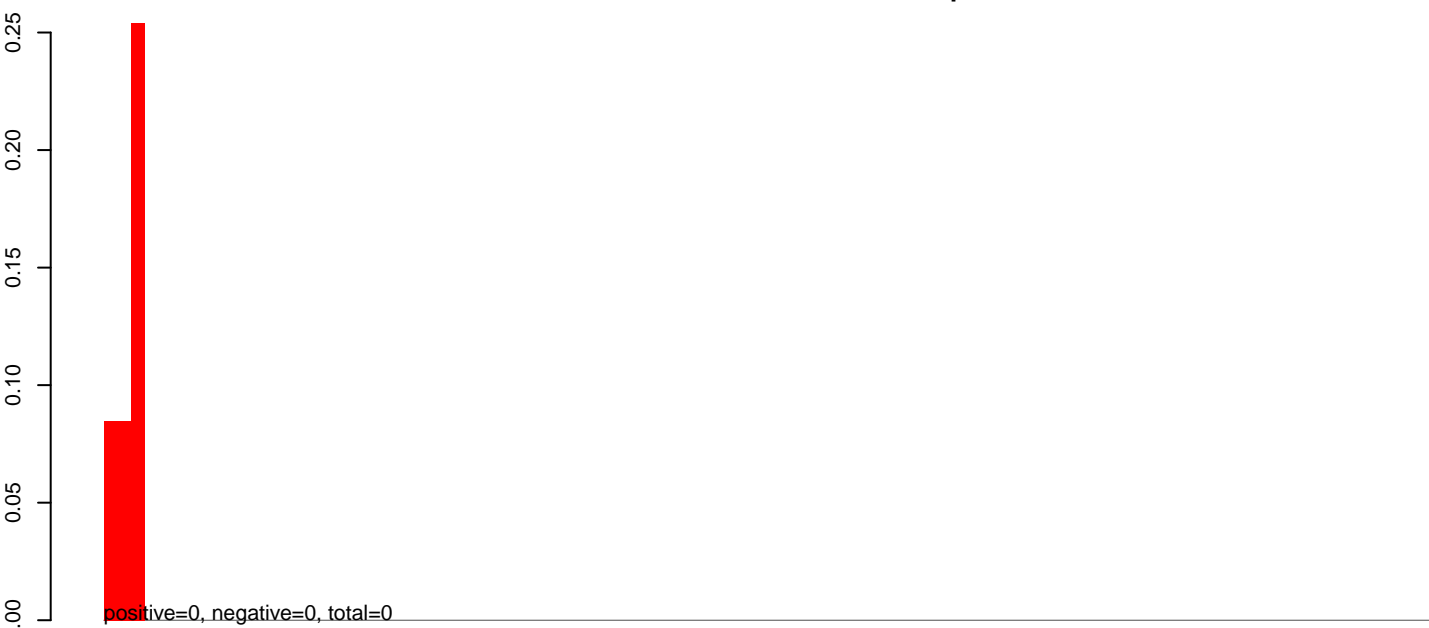
AnGam_Mos55_cells.rep



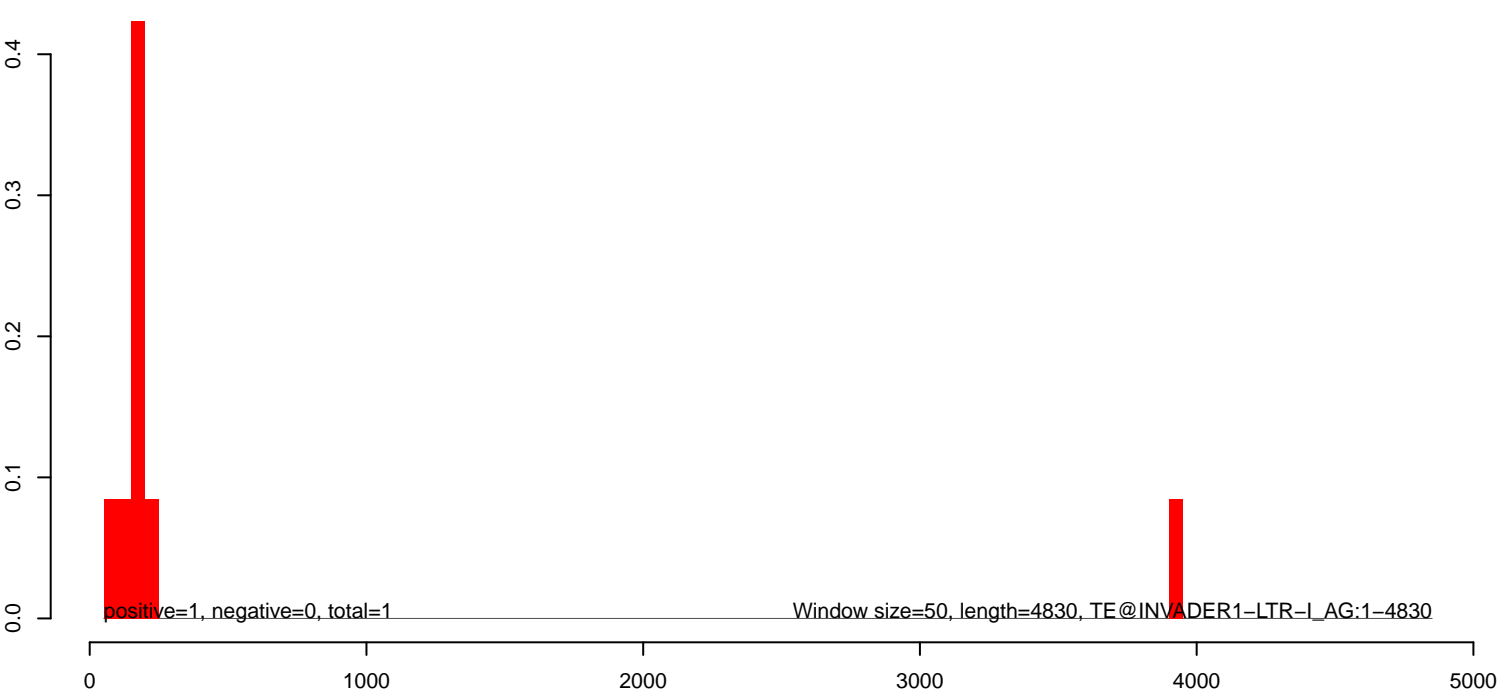
AnGam_Mos55_cells.18_23.rep



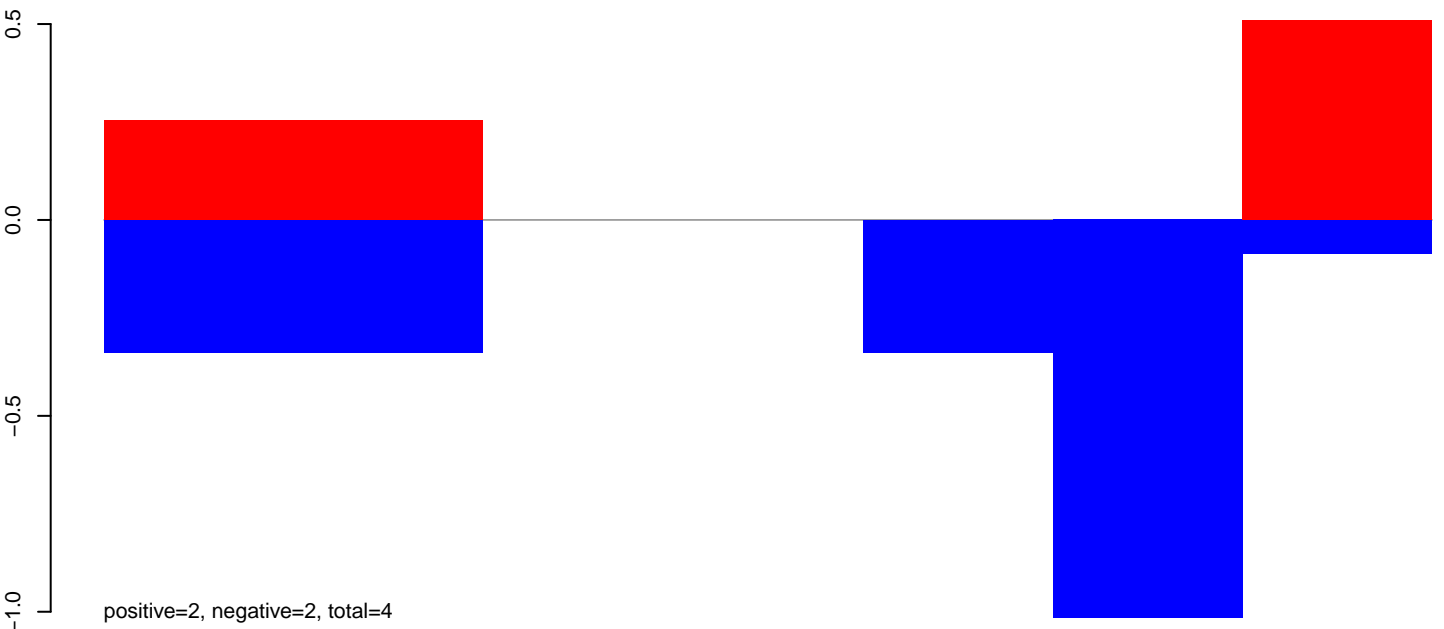
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



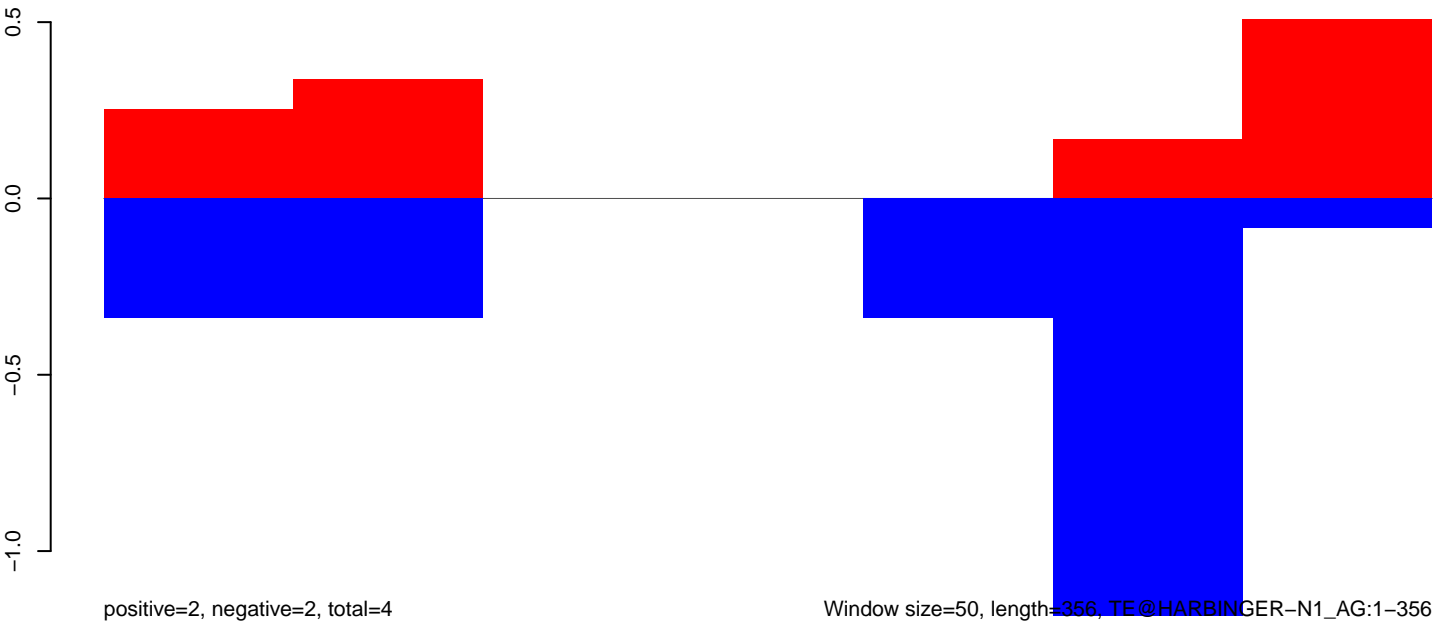
AnGam_Mos55_cells.18_23.rep



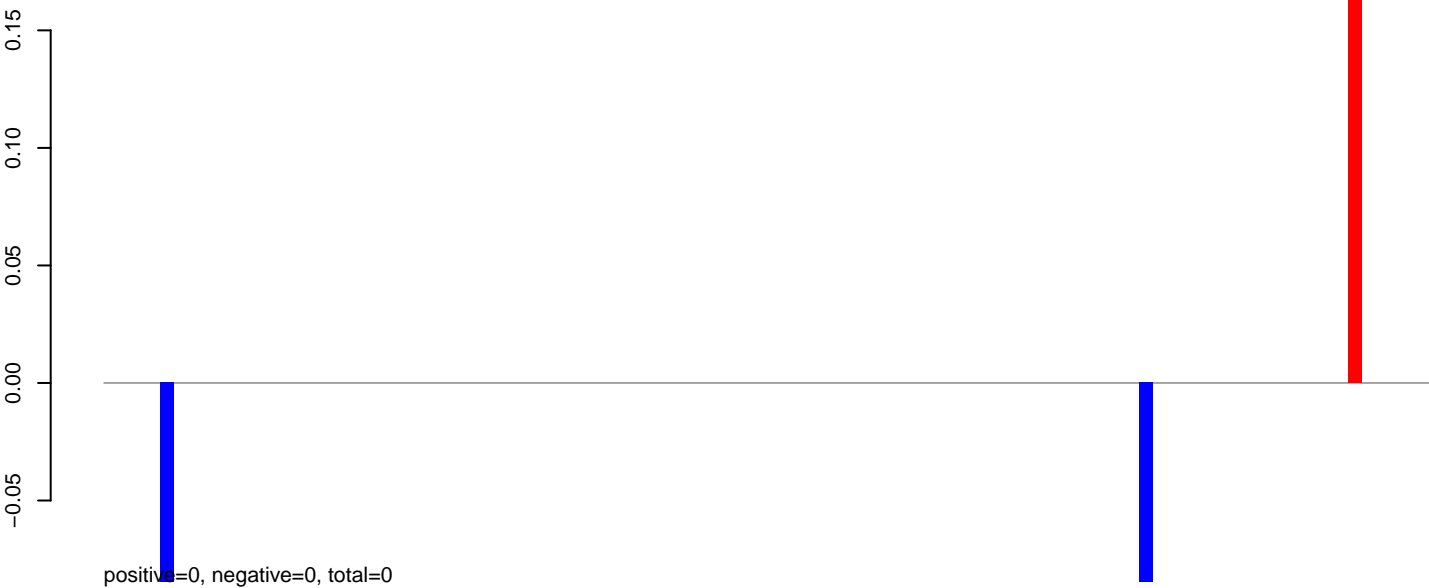
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



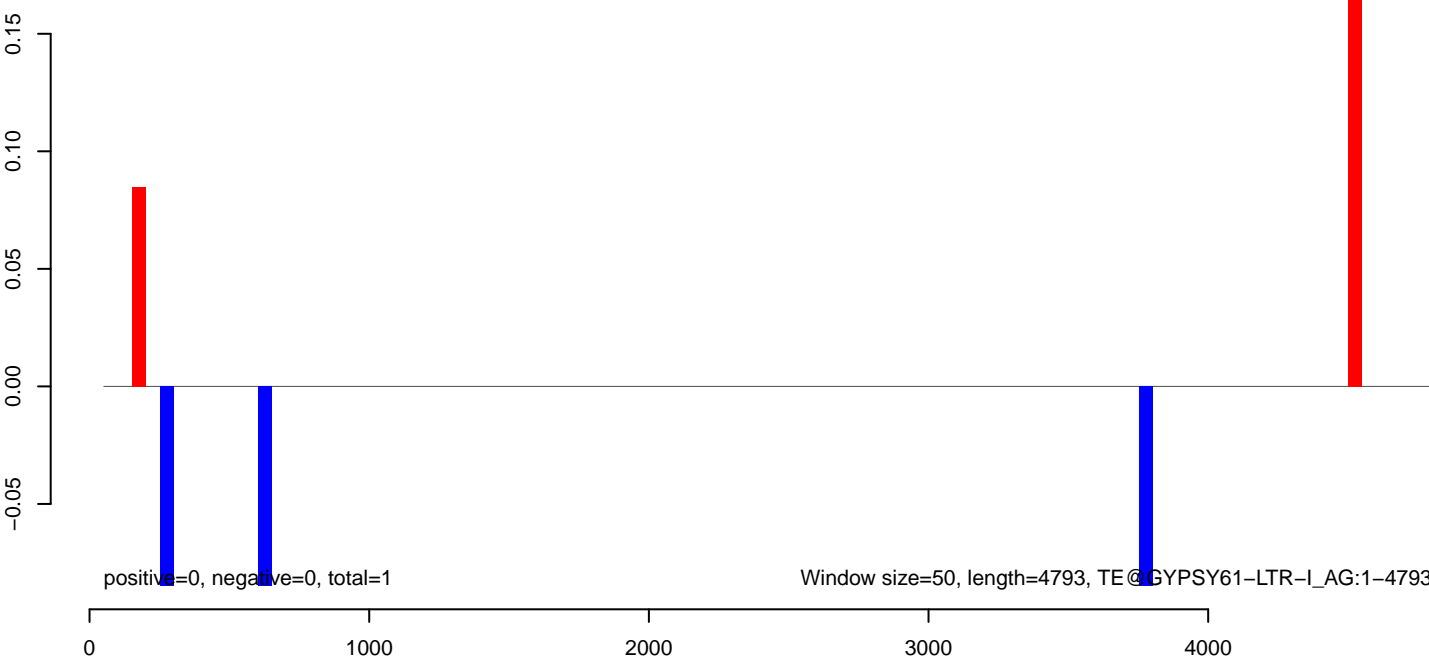
AnGam_Mos55_cells.18_23.rep



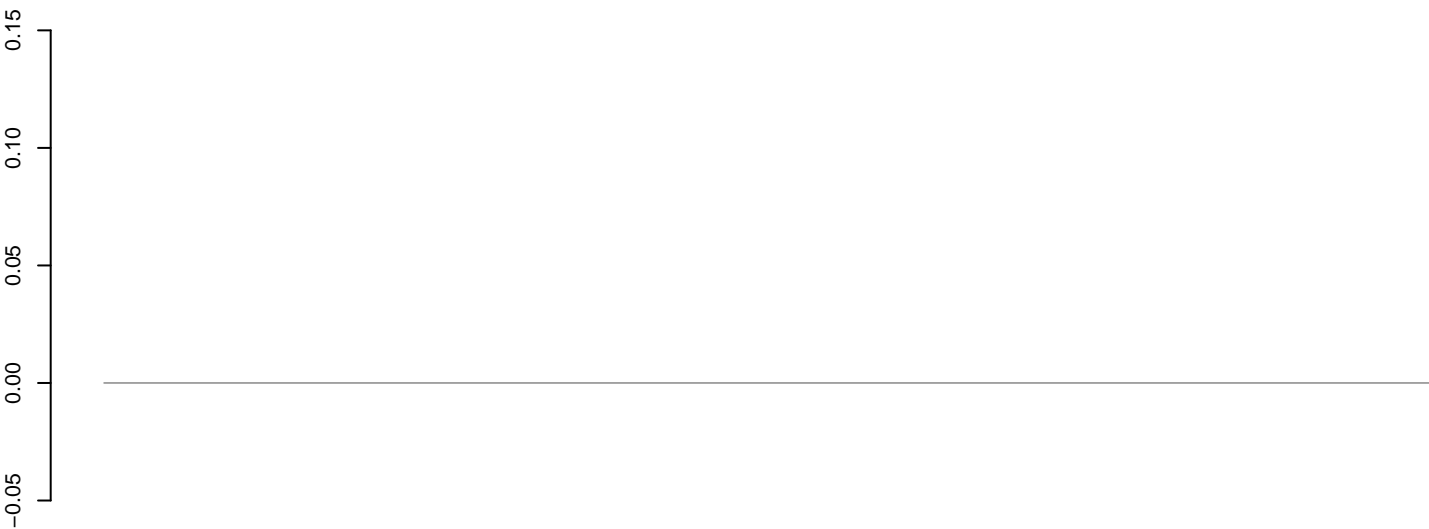
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

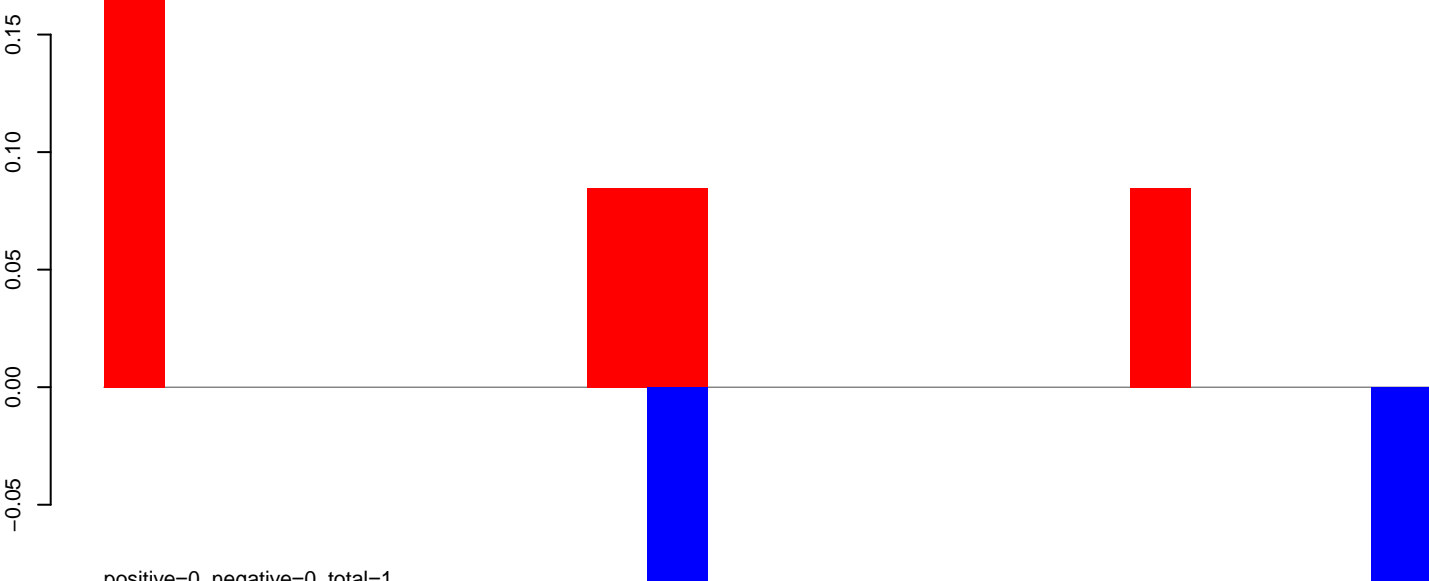


AnGam_Mos55_cells.18_23.rep



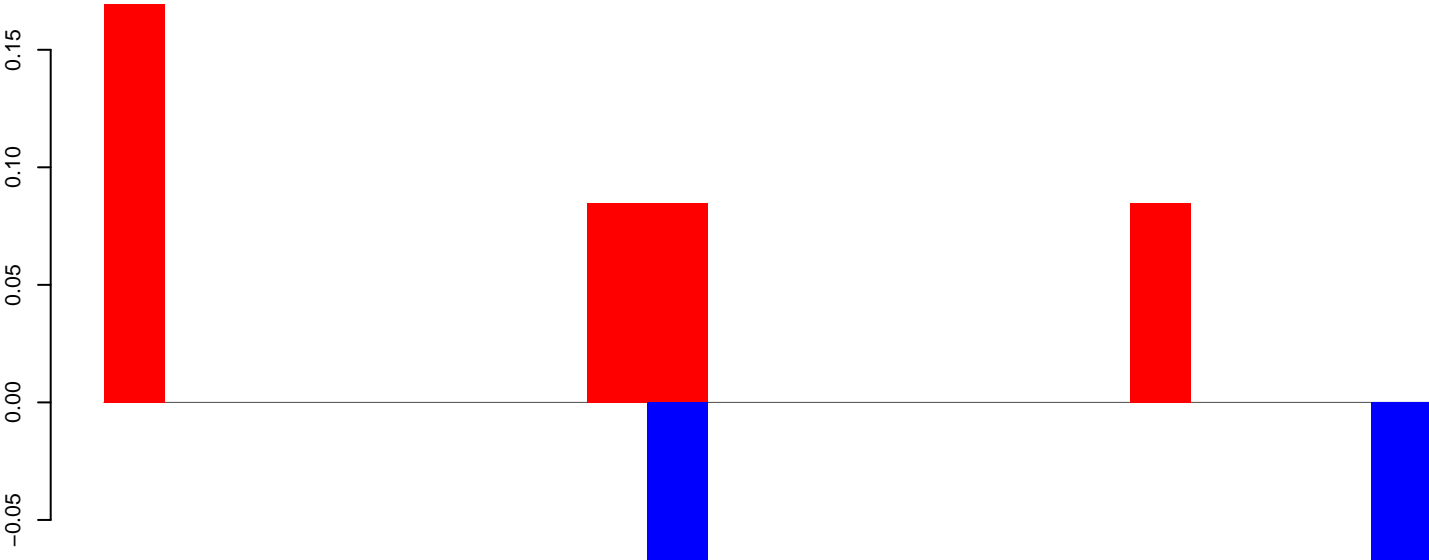
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=1

AnGam_Mos55_cells.rep

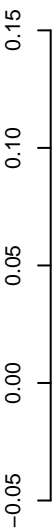


positive=0, negative=0, total=1

Window size=50, length=1110, TE@EAL40181-Unknown:1-1110

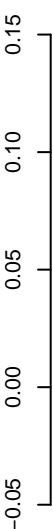
200 400 600 800 1000

AnGam_Mos55_cells.18_23.rep



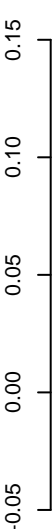
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



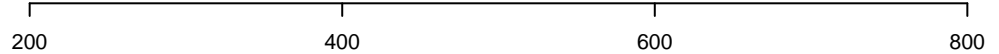
positive=1, negative=0, total=1

AnGam_Mos55_cells.rep

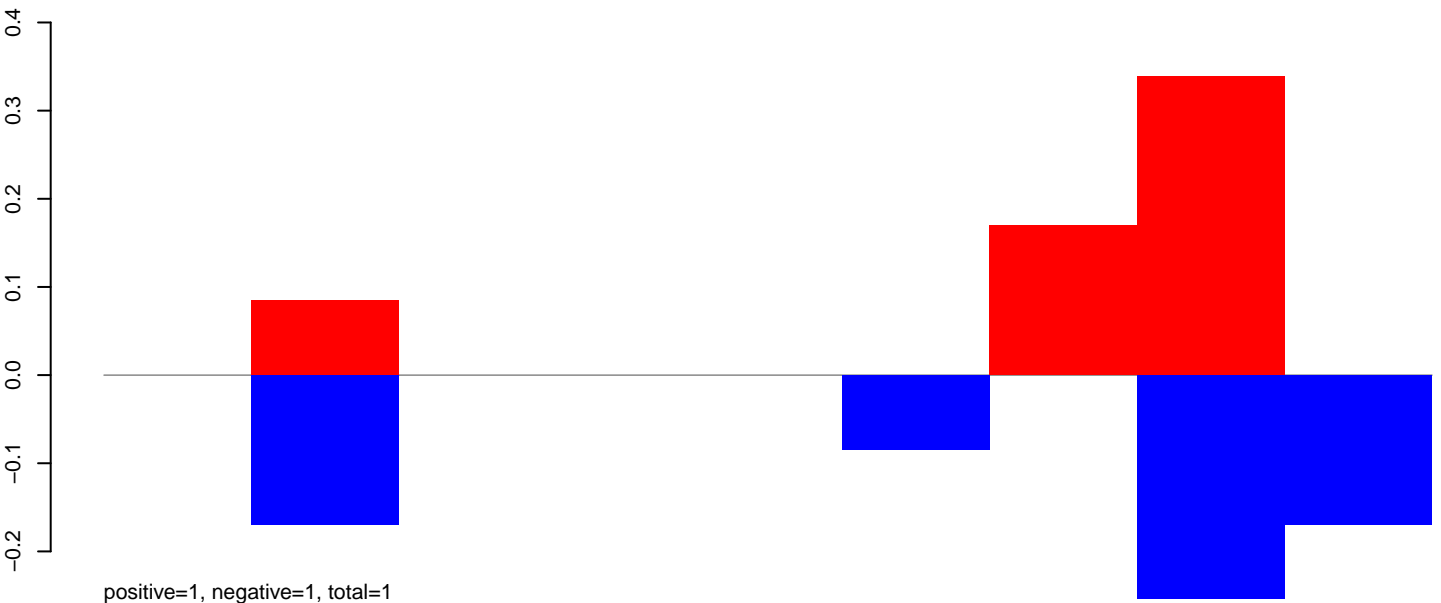


positive=1, negative=0, total=1

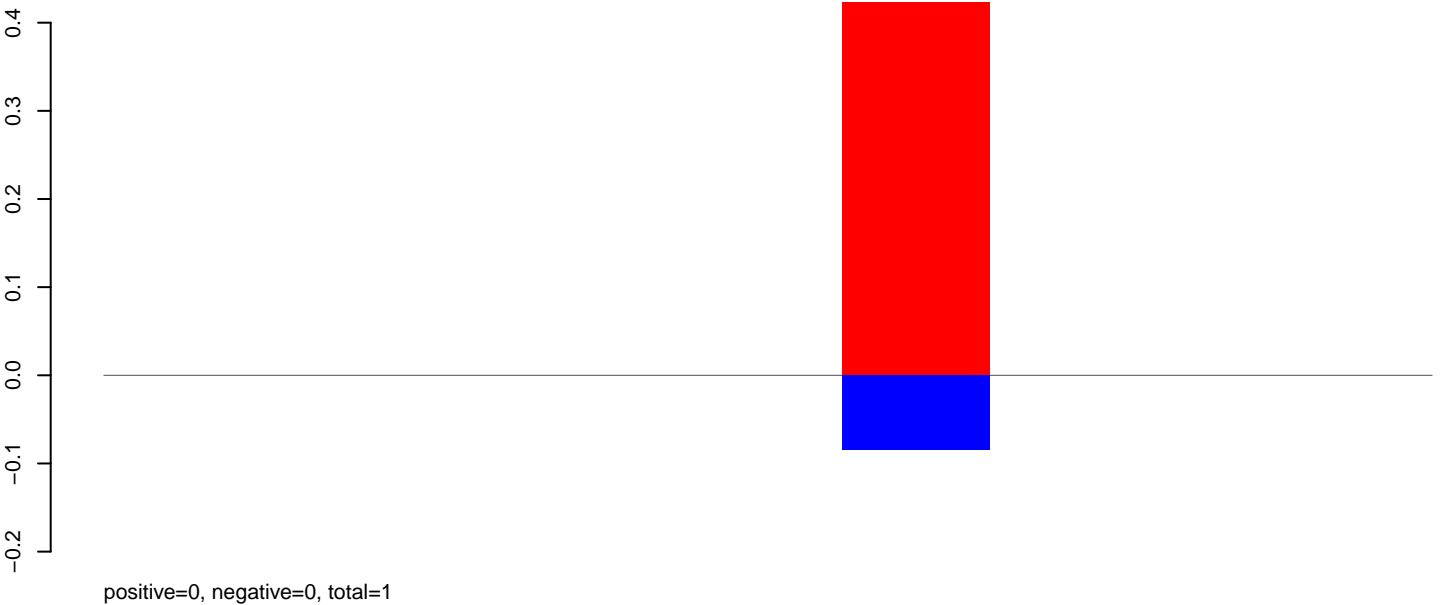
Window size=50, length=851, TE@Clu-87B_AG:1-851



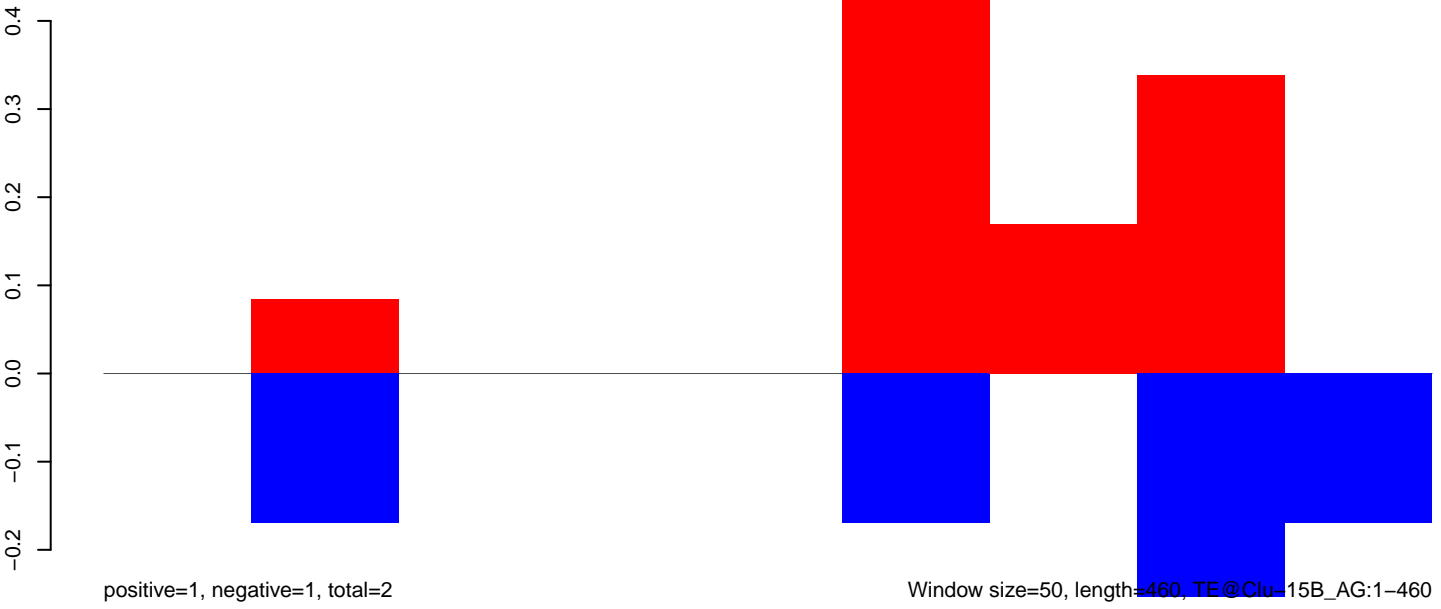
AnGam_Mos55_cells.18_23.rep



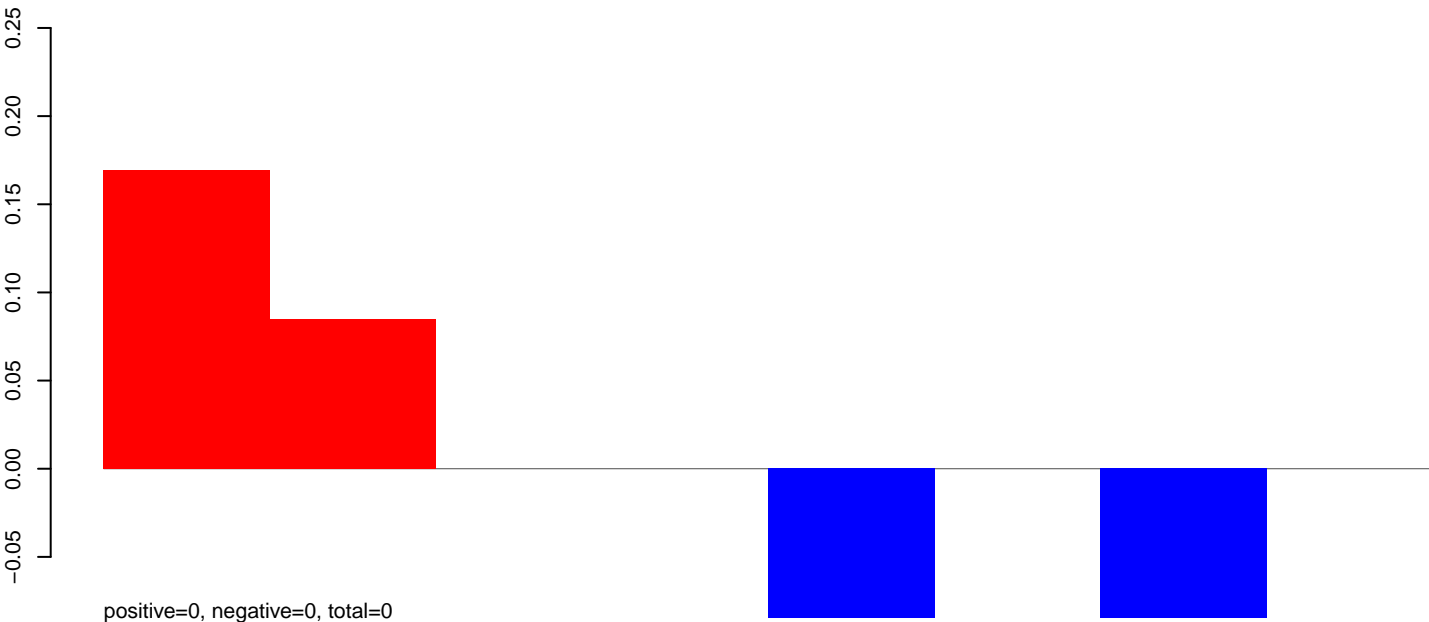
AnGam_Mos55_cells.24_35.rep



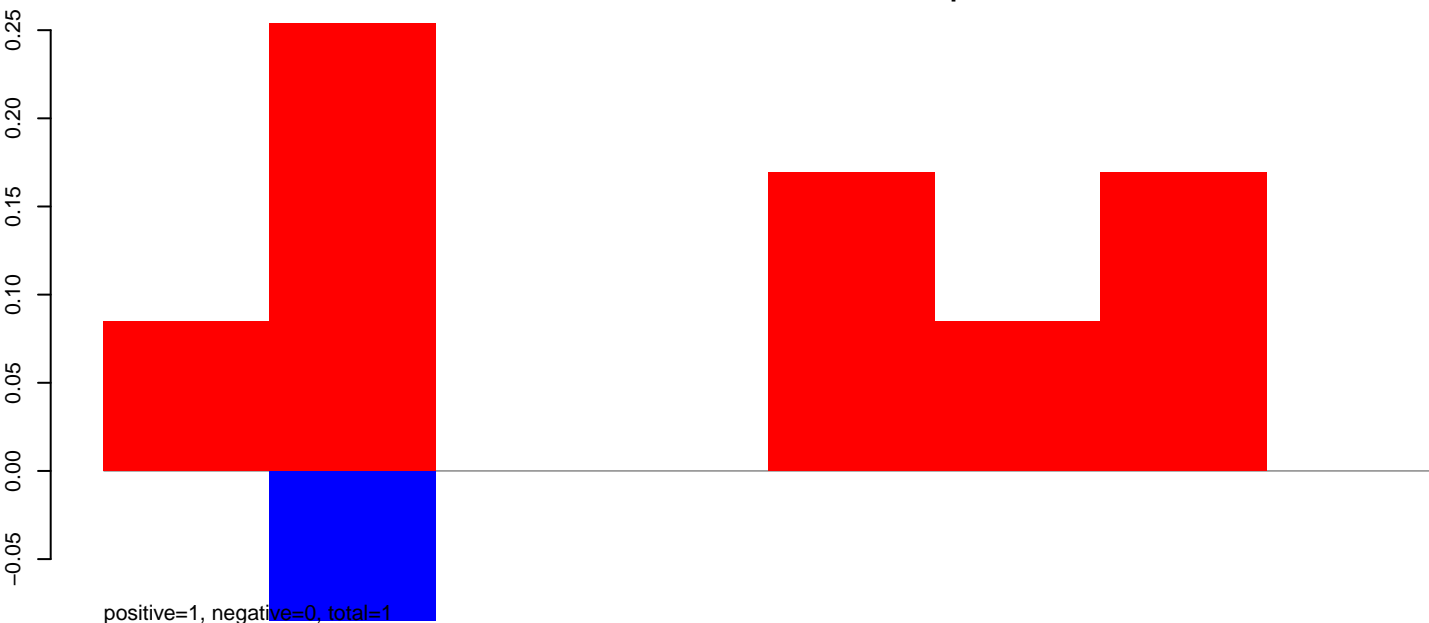
AnGam_Mos55_cells.rep



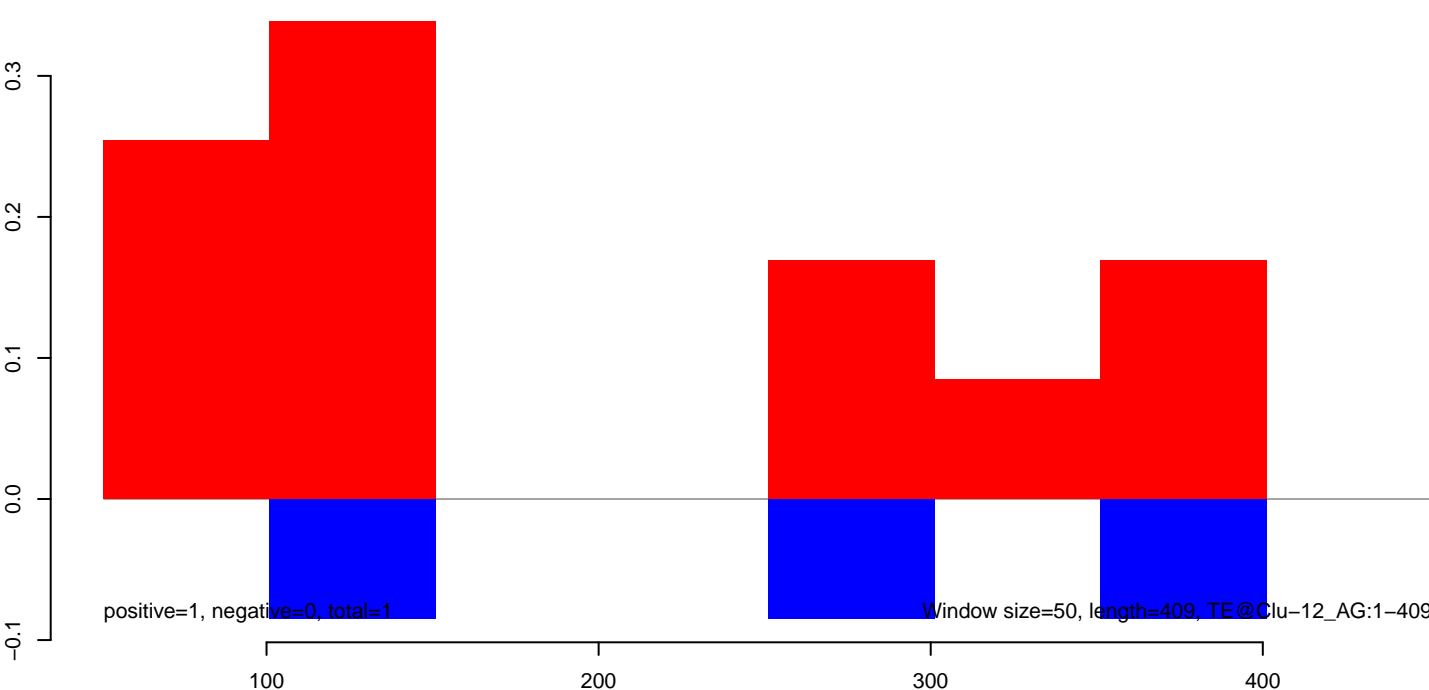
AnGam_Mos55_cells.18_23.rep

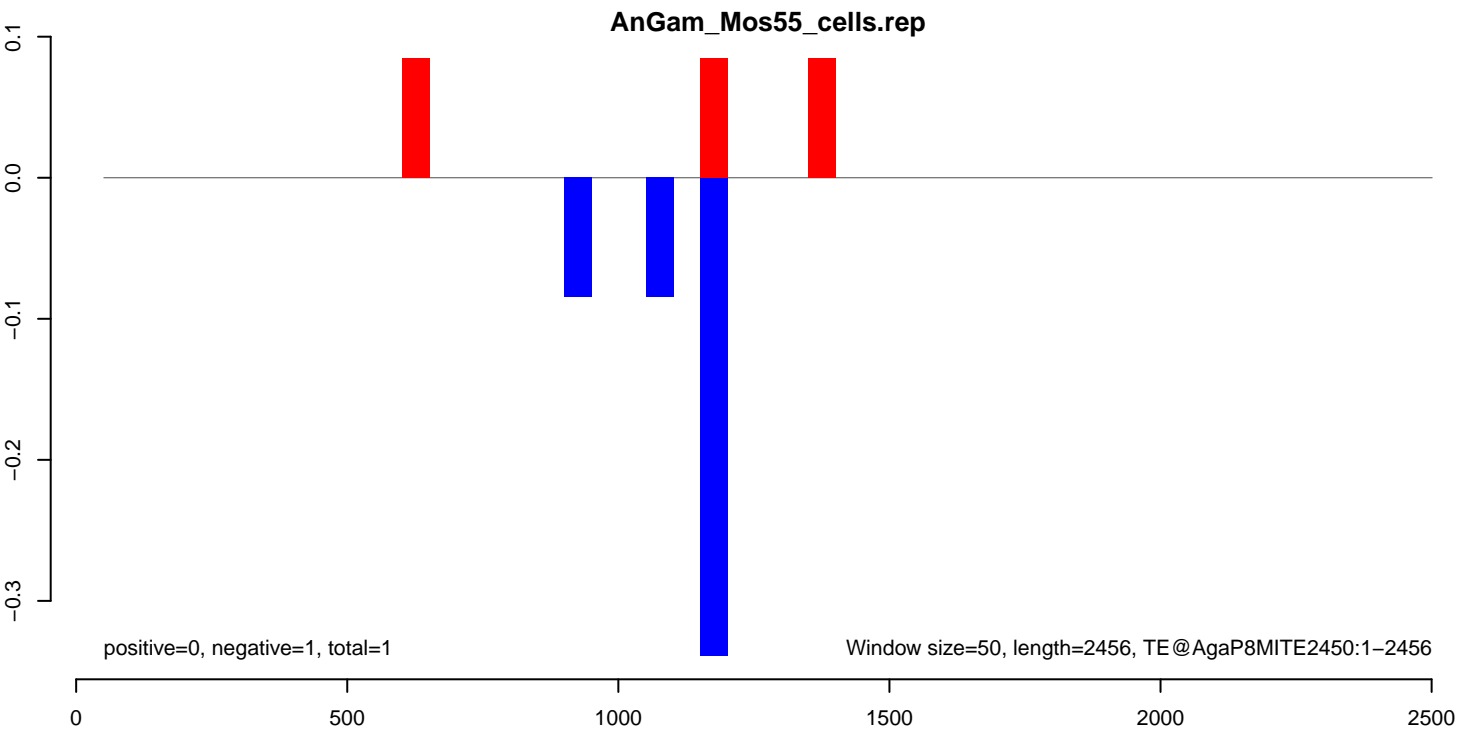
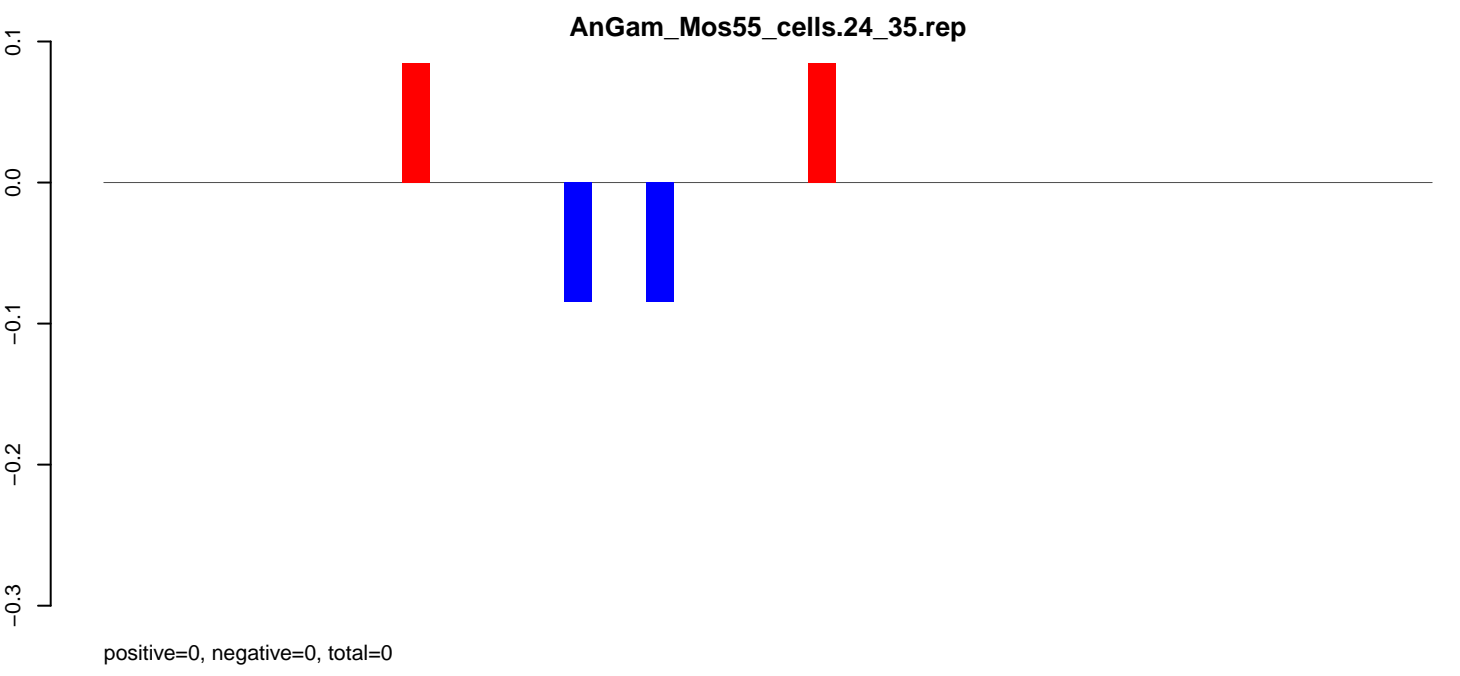
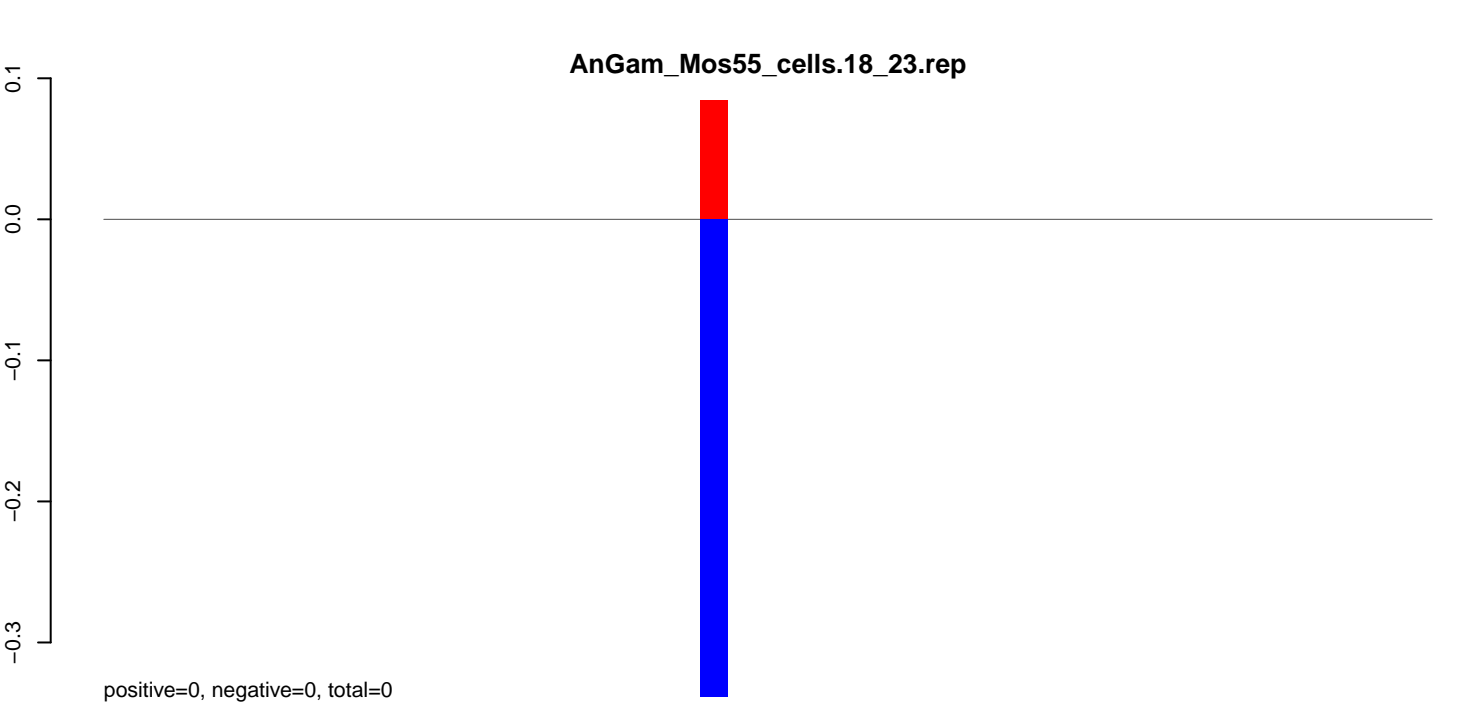


AnGam_Mos55_cells.24_35.rep

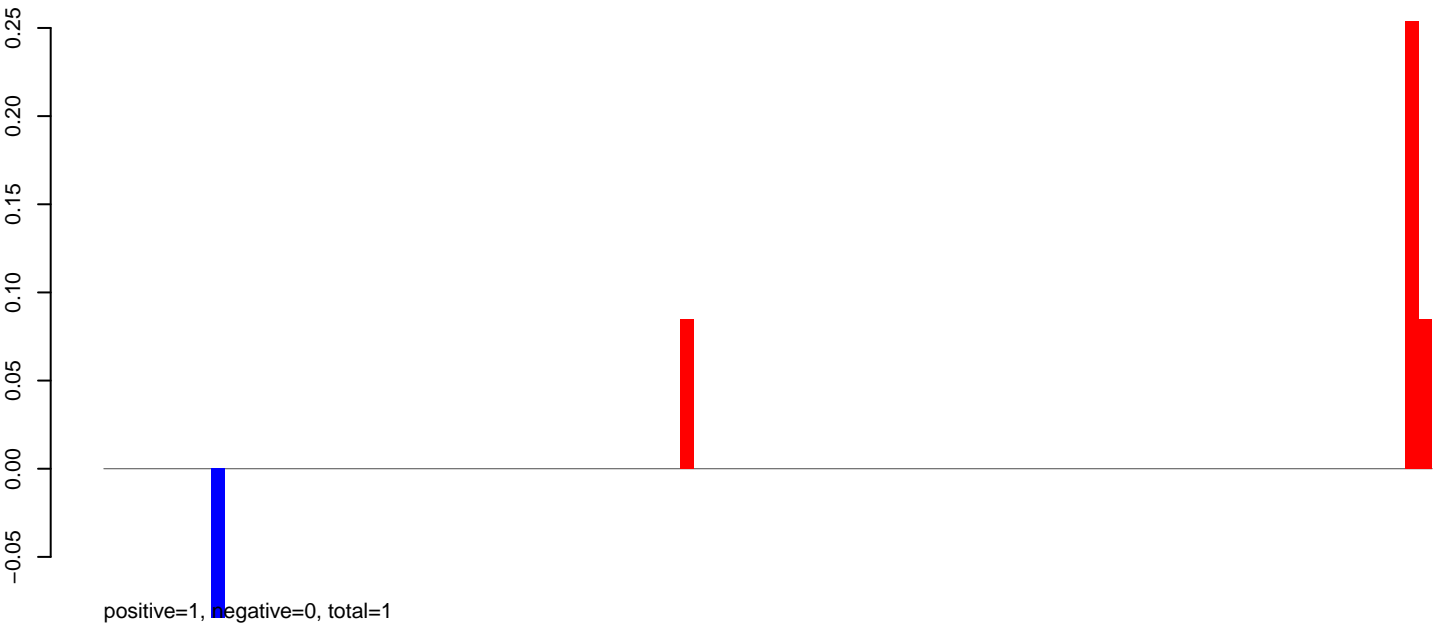


AnGam_Mos55_cells.rep





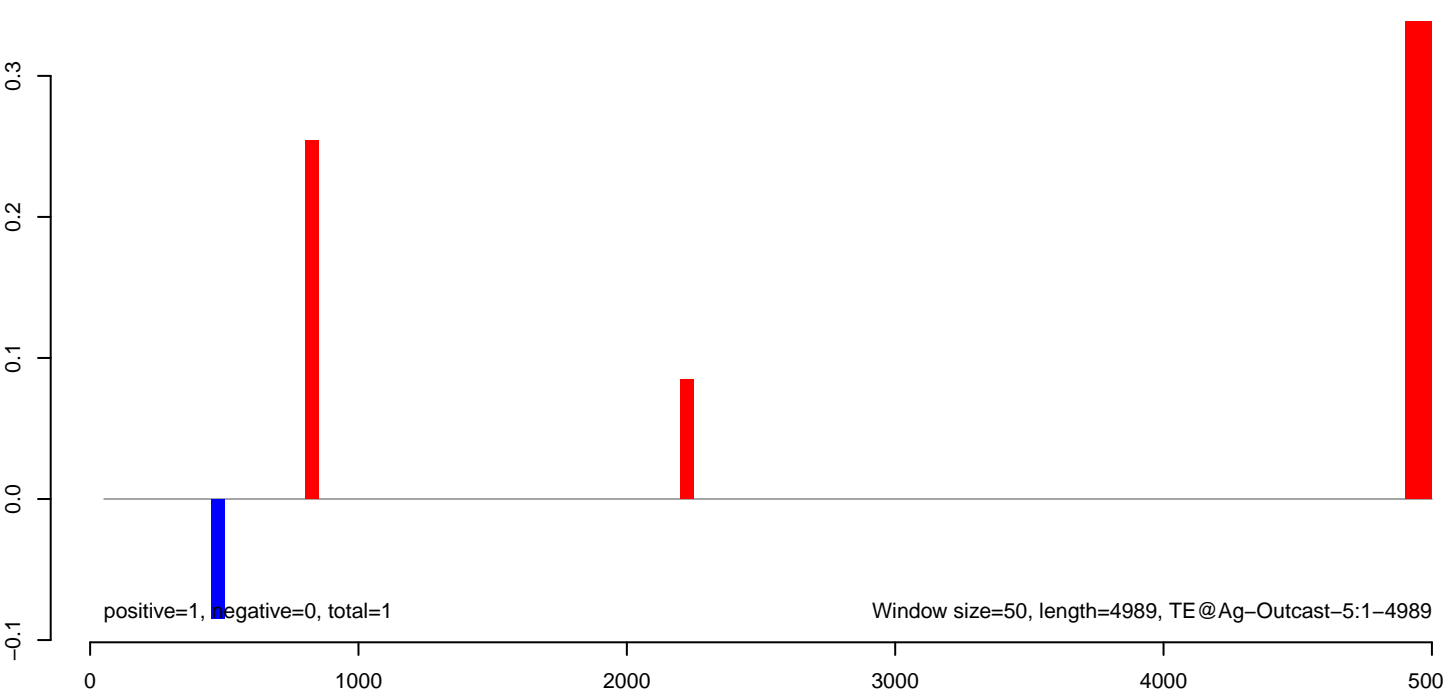
AnGam_Mos55_cells.18_23.rep



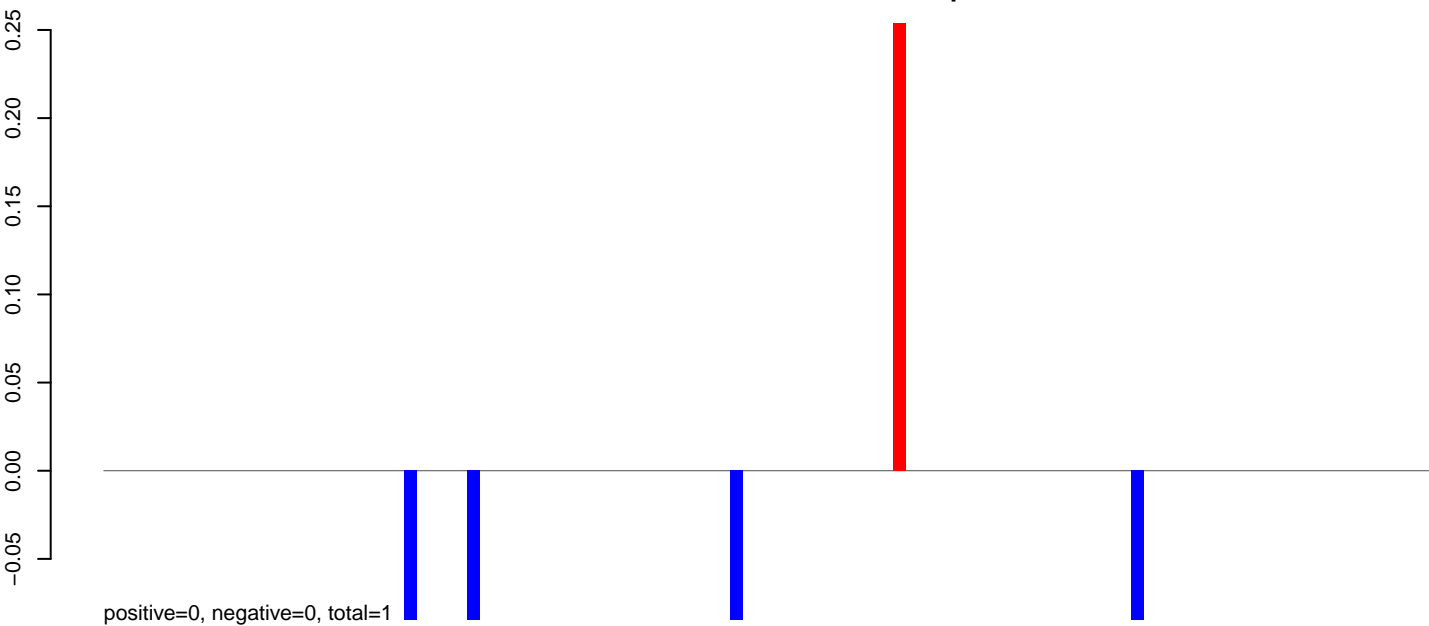
AnGam_Mos55_cells.24_35.rep



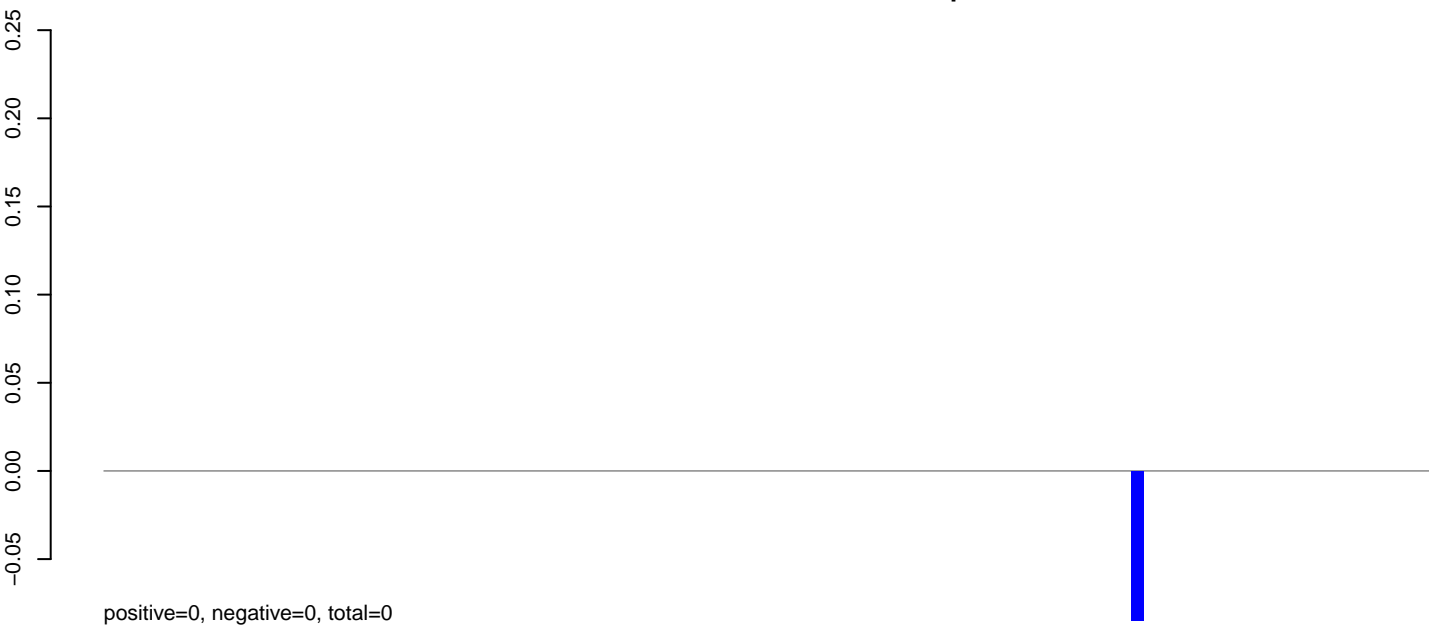
AnGam_Mos55_cells.rep



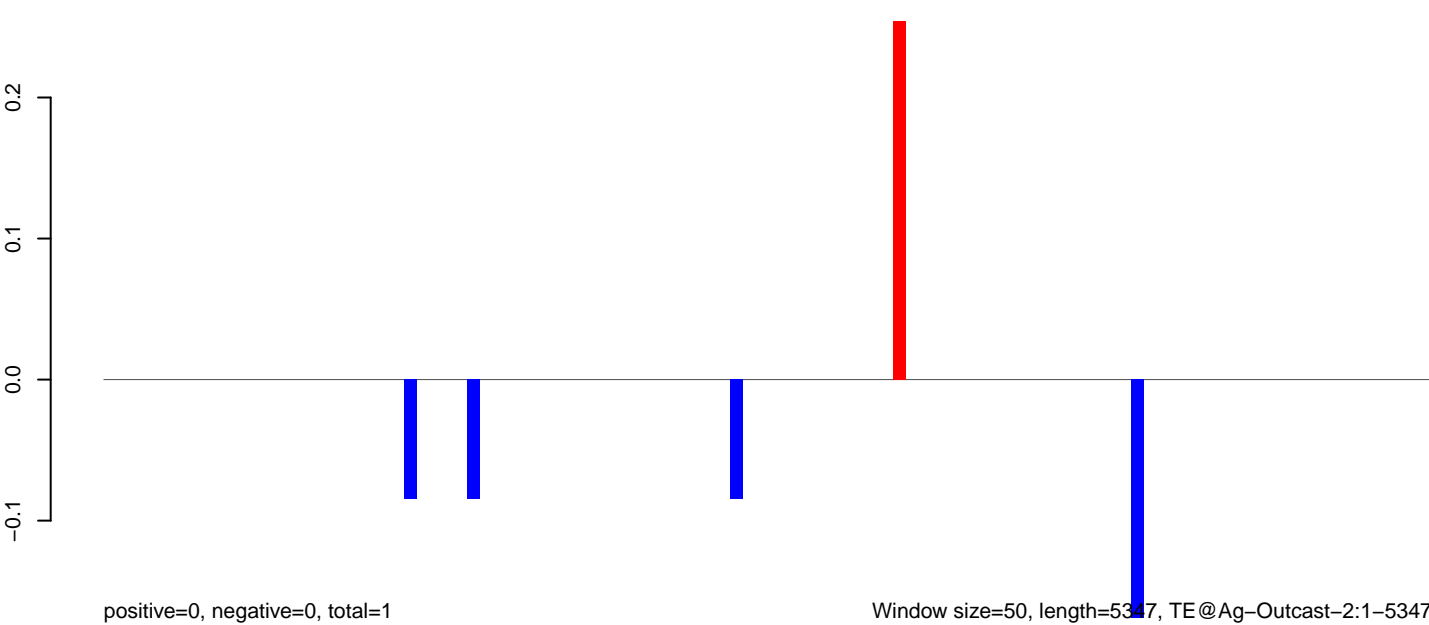
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



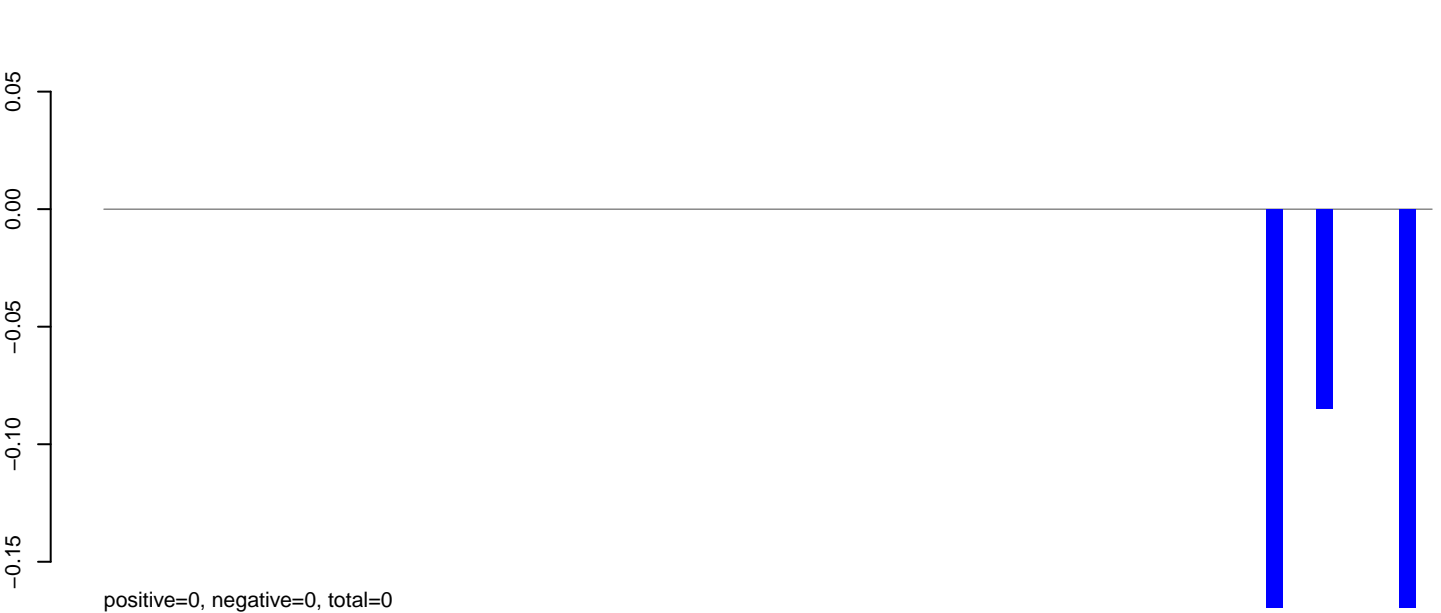
Window size=50, length=5347, TE@Ag-Outcast-2:1-5347

0 1000 2000 3000 4000 5000

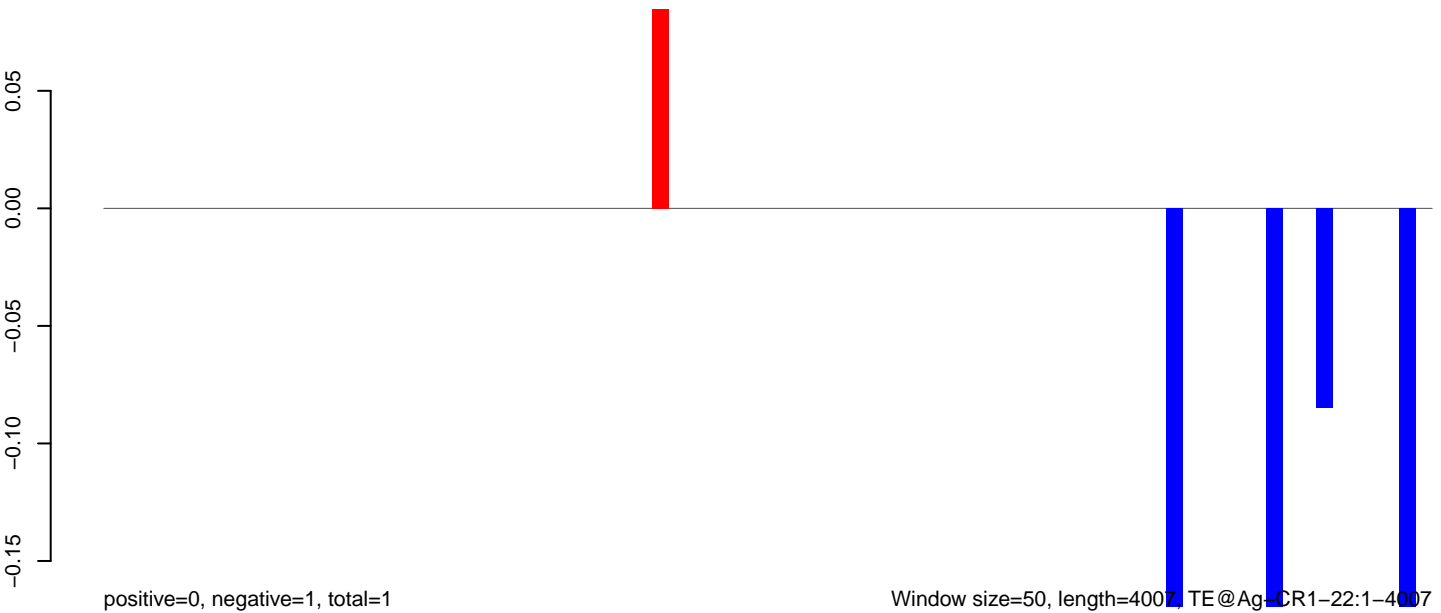
AnGam_Mos55_cells.18_23.rep



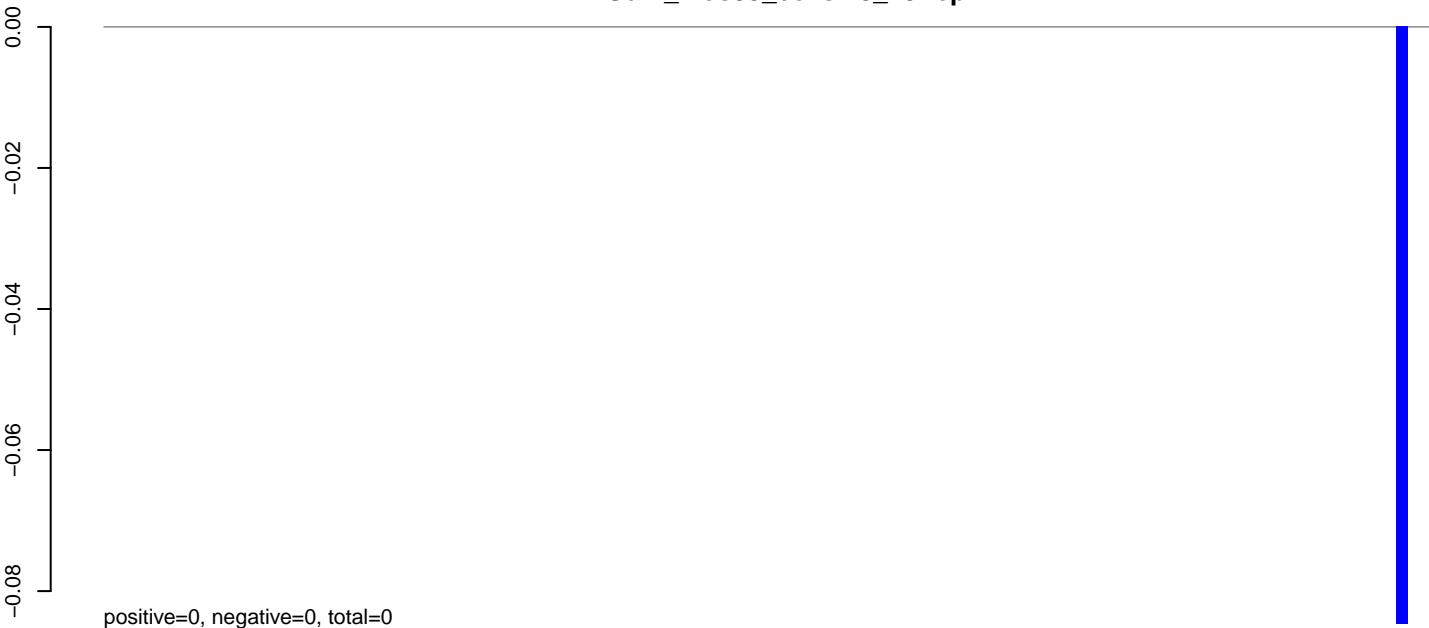
AnGam_Mos55_cells.24_35.rep



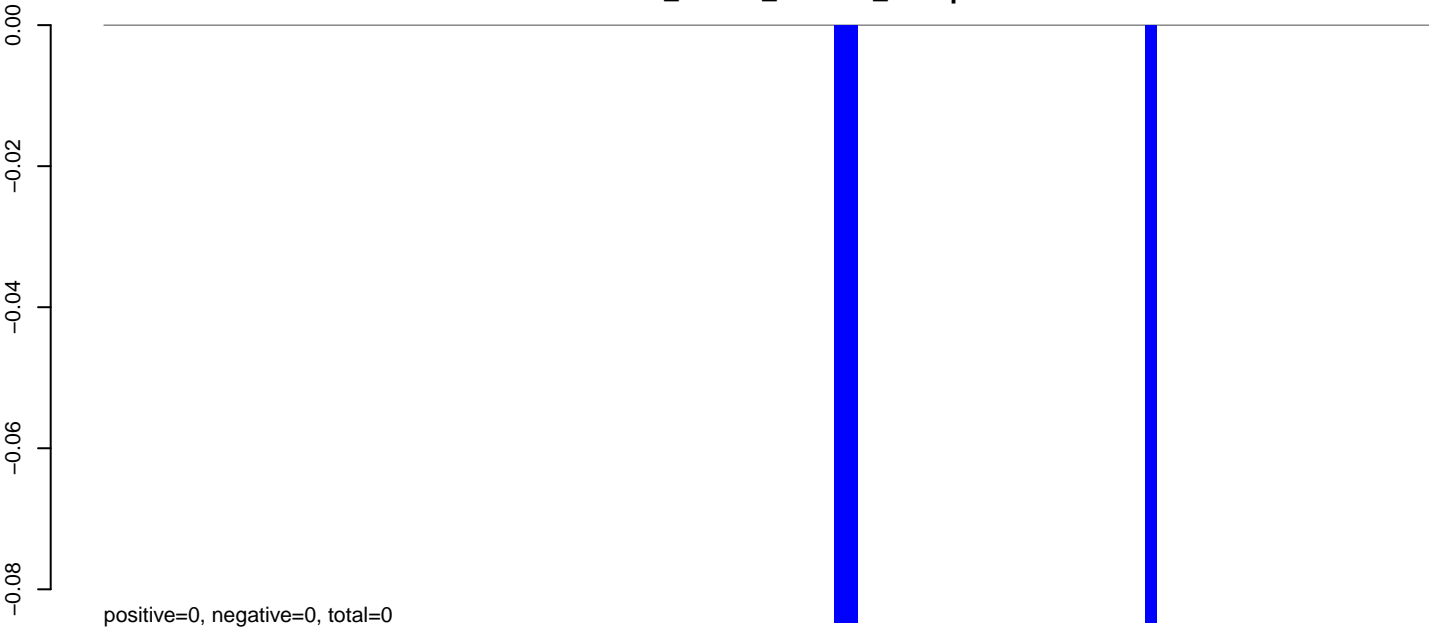
AnGam_Mos55_cells.rep



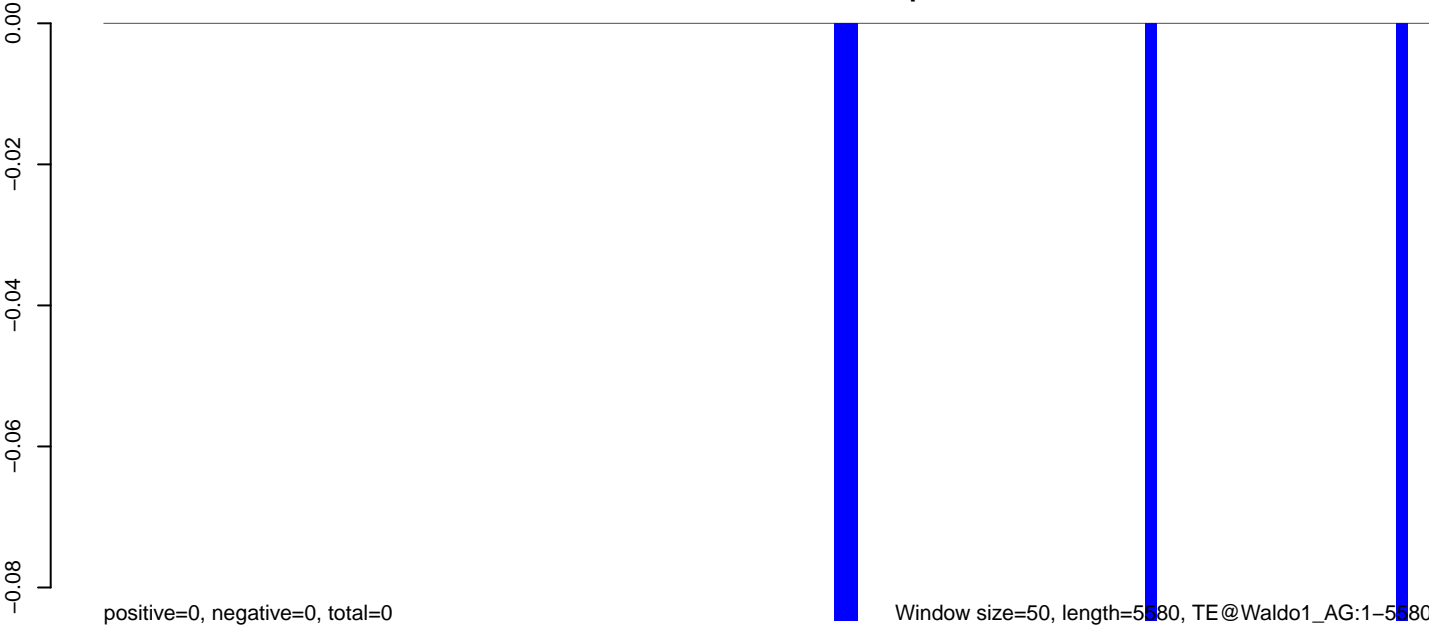
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

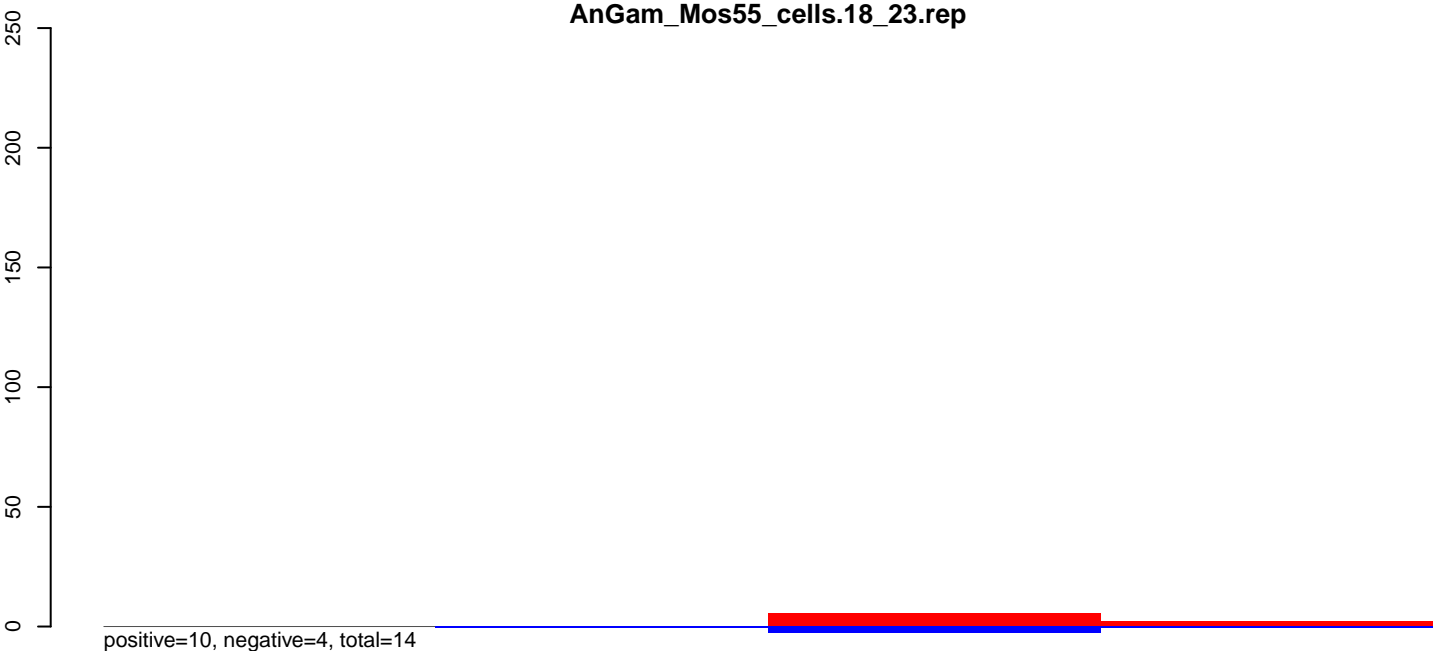


AnGam_Mos55_cells.rep

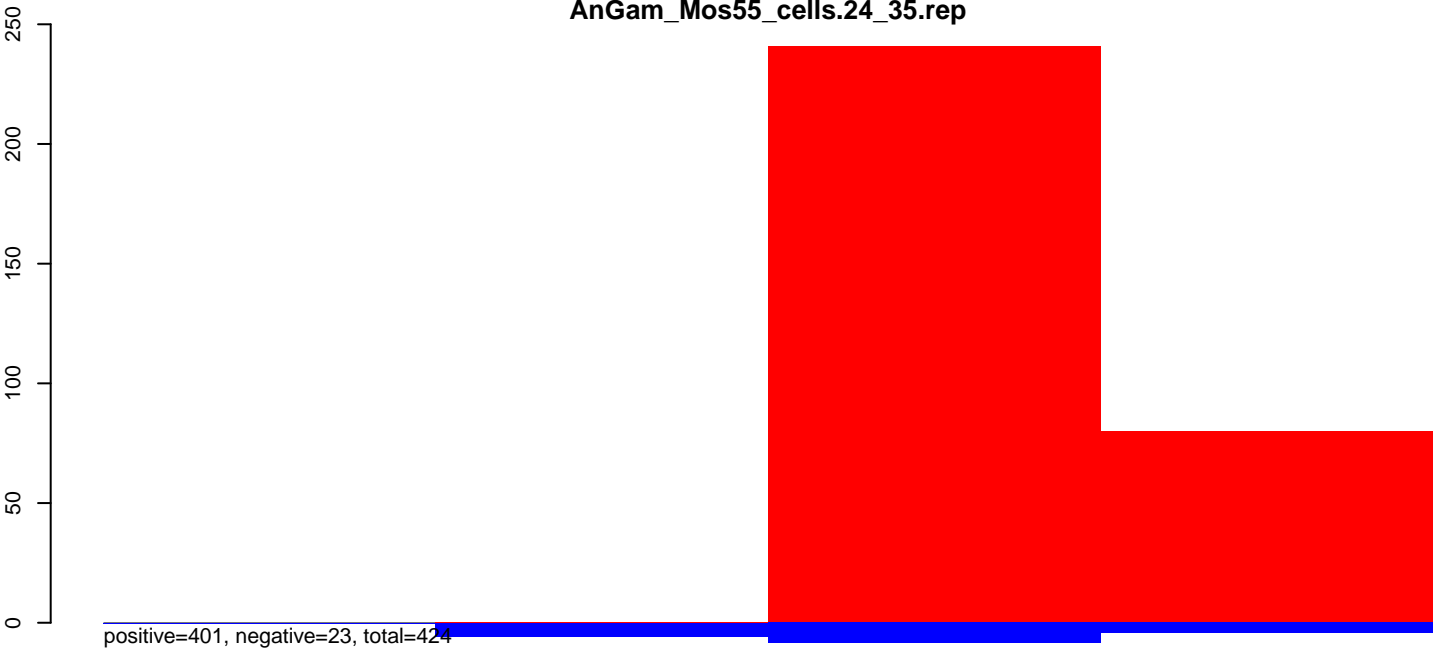


0 1000 2000 3000 4000 5000

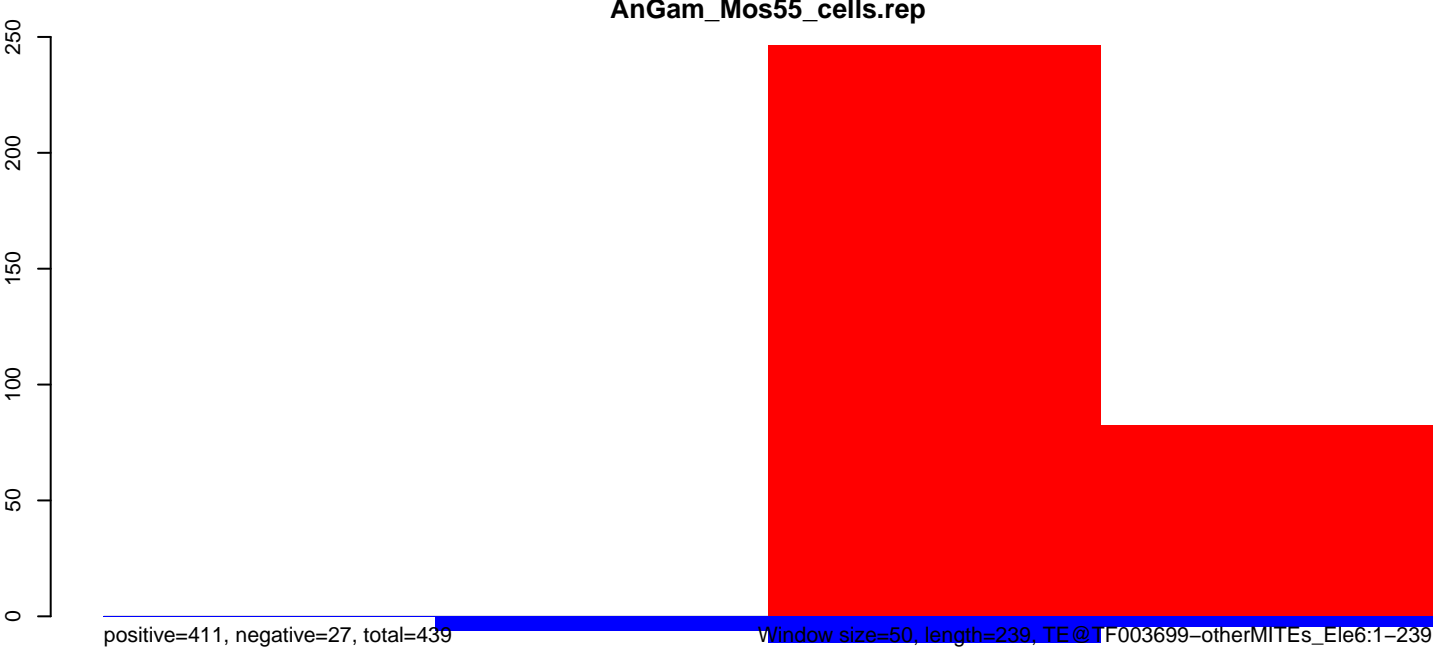
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

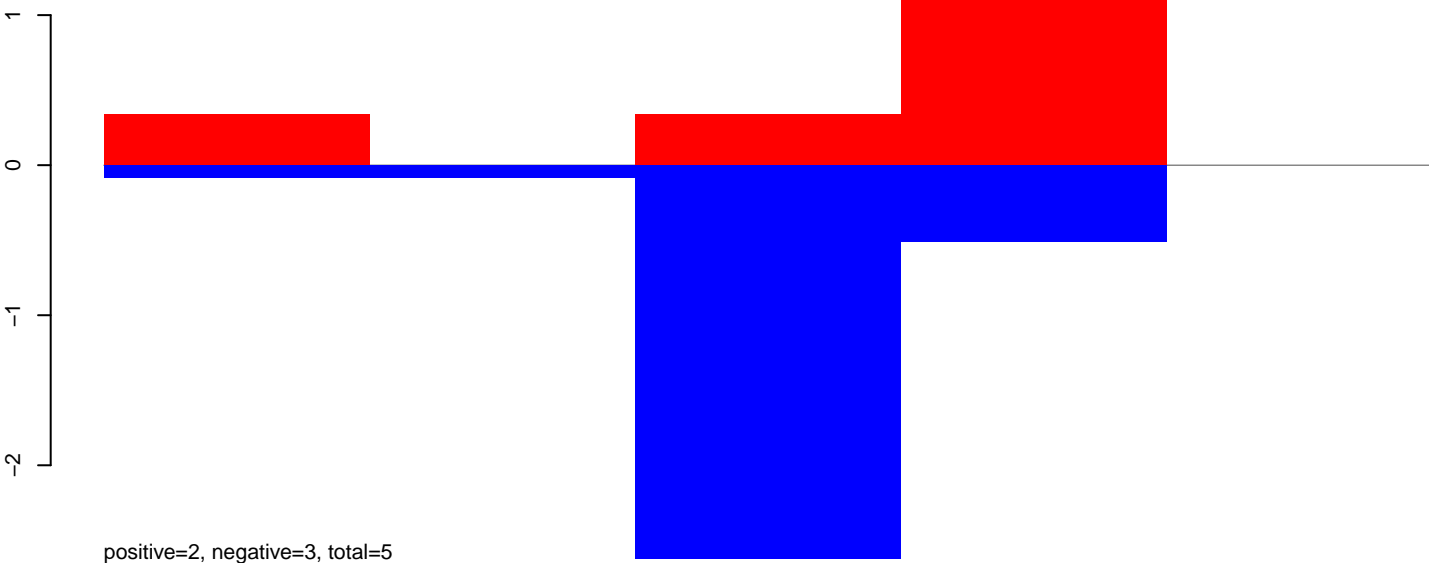


AnGam_Mos55_cells.rep

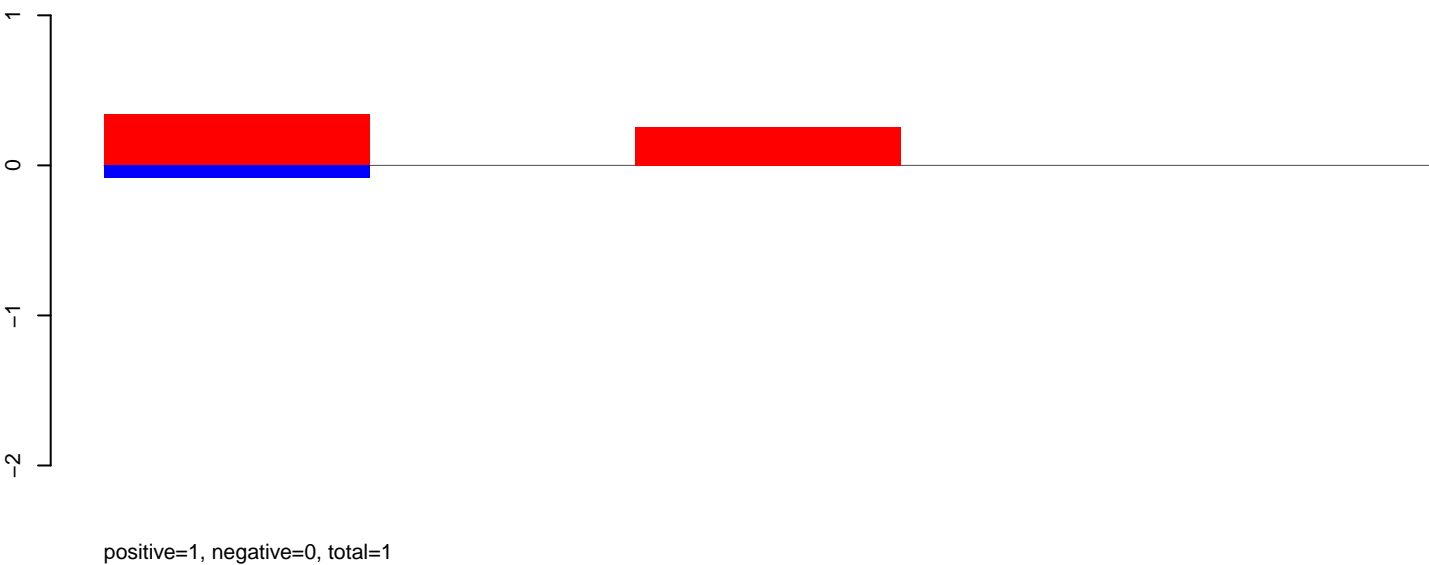


50 100 150 200 250

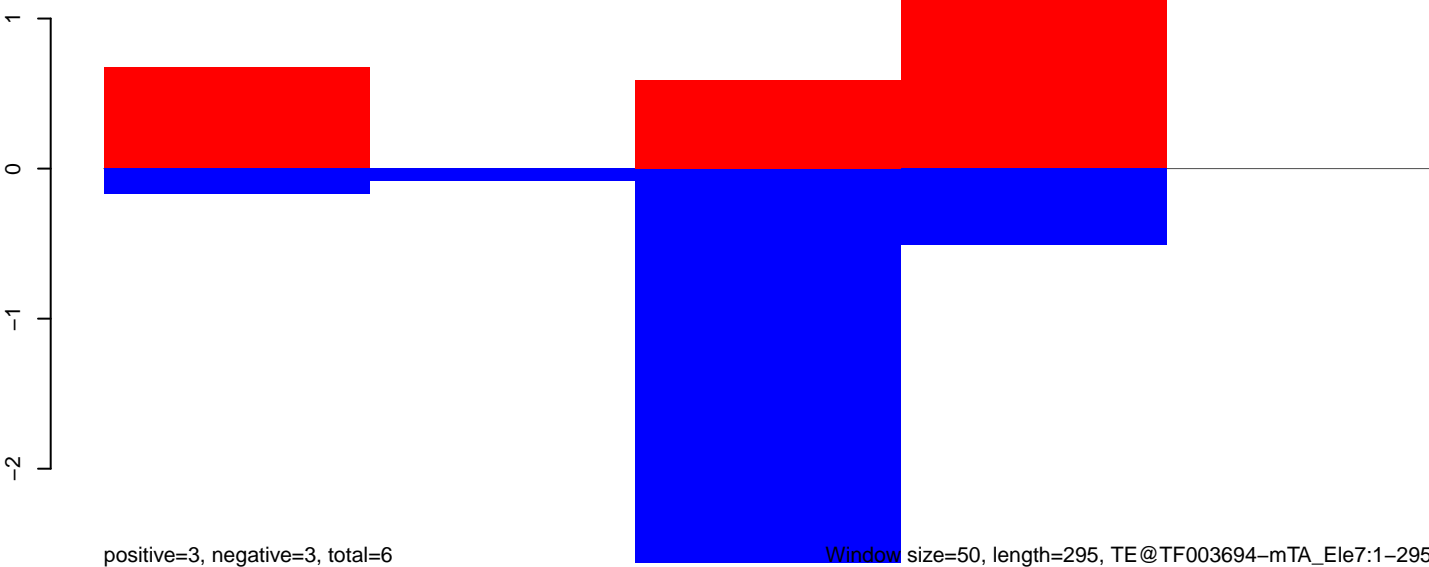
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

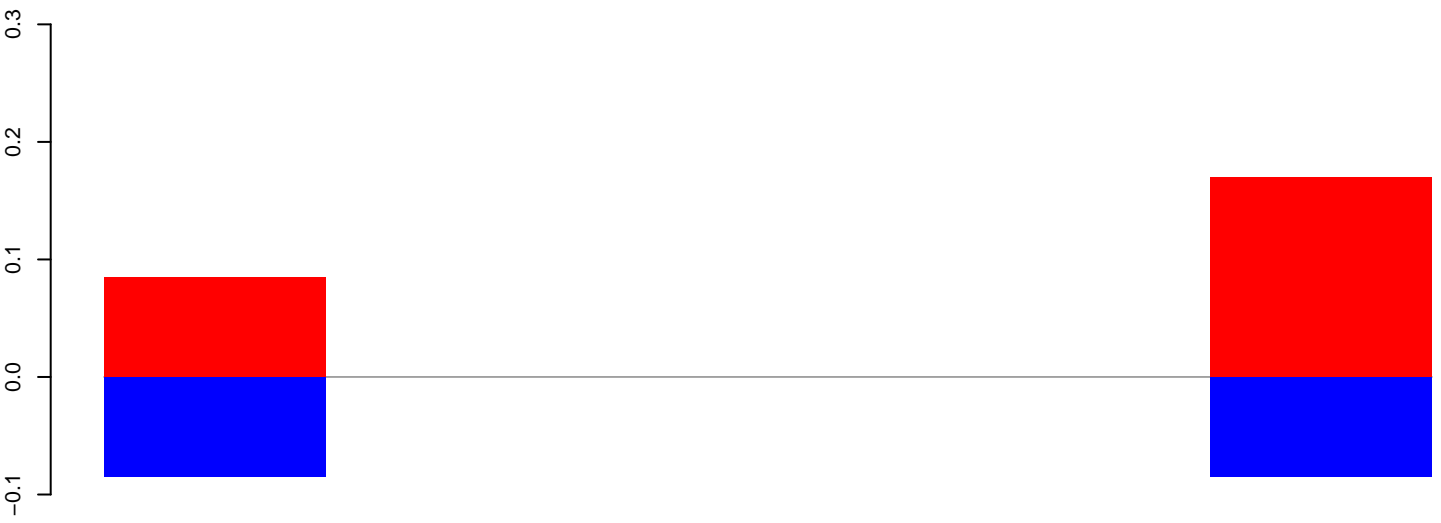


AnGam_Mos55_cells.rep



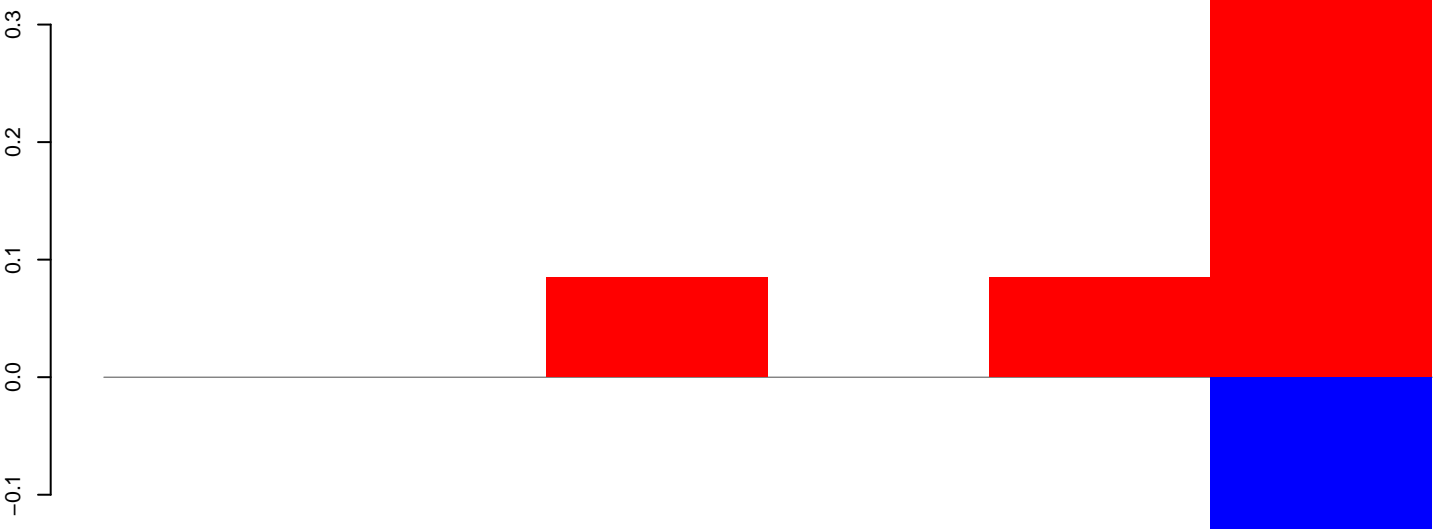
Window size=50, length=295, TE@TF003694-mTA_Ele7:1-295

AnGam_Mos55_cells.18_23.rep



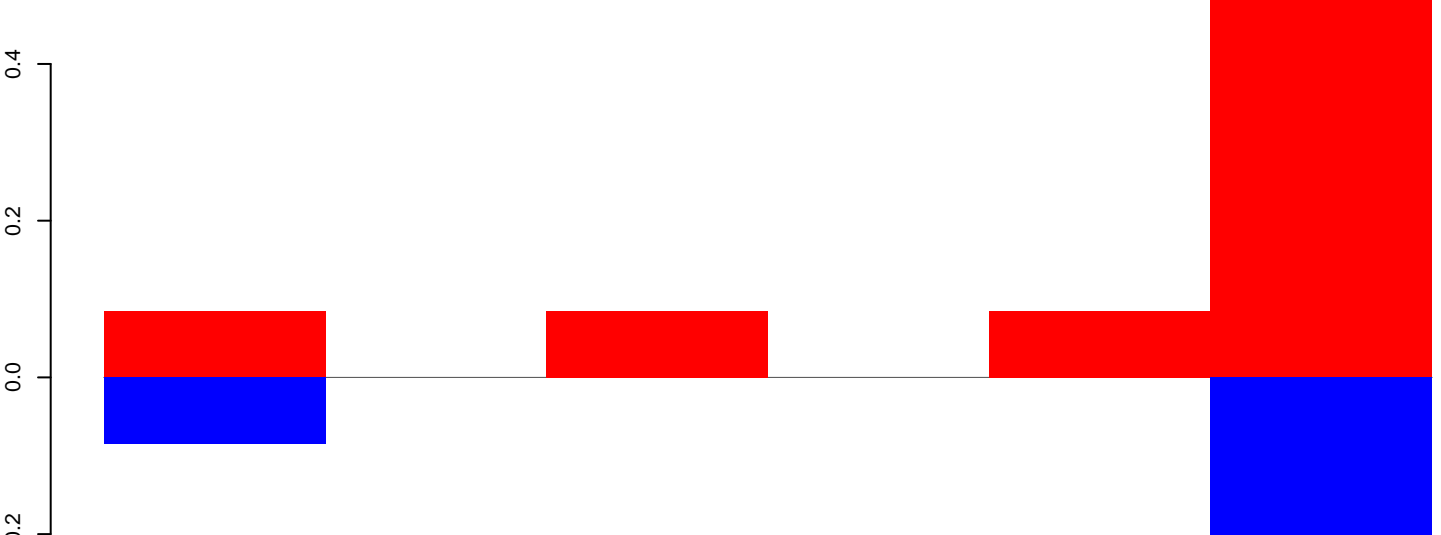
positive=0, negative=0, total=1

AnGam_Mos55_cells.24_35.rep



positive=1, negative=0, total=1

AnGam_Mos55_cells.rep

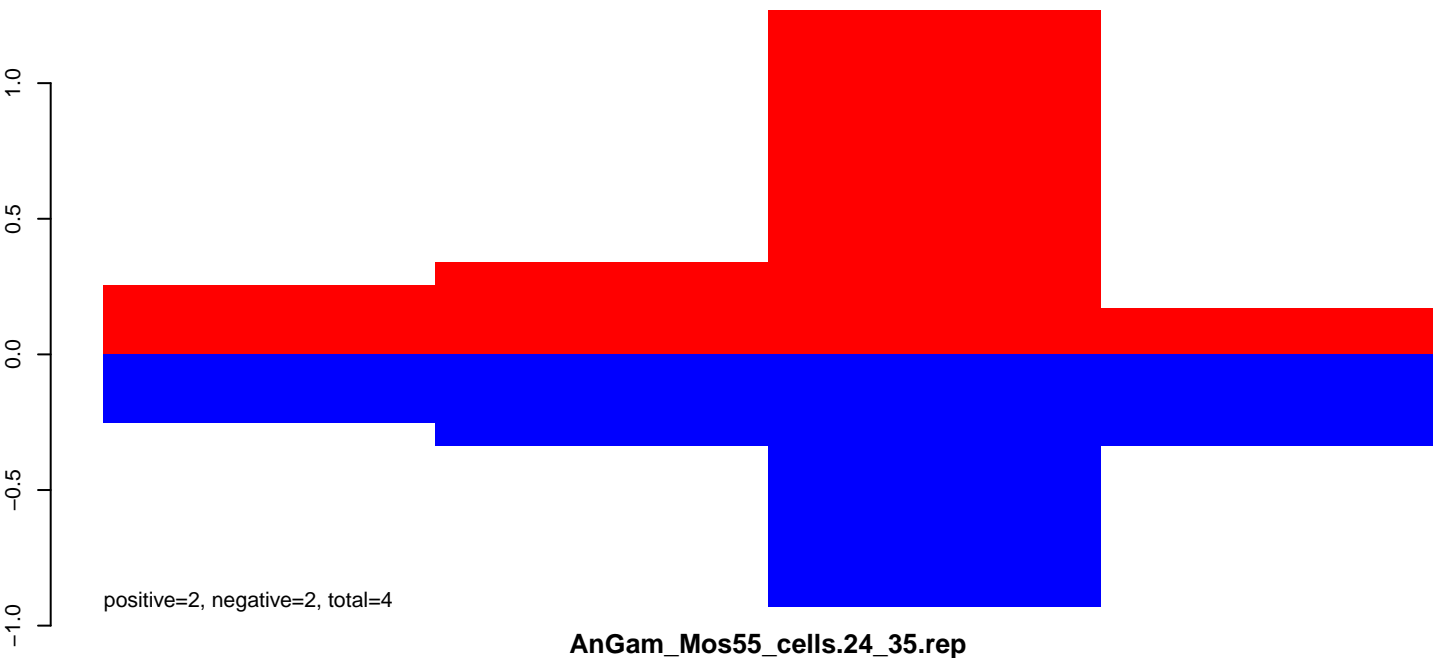


positive=1, negative=1, total=2

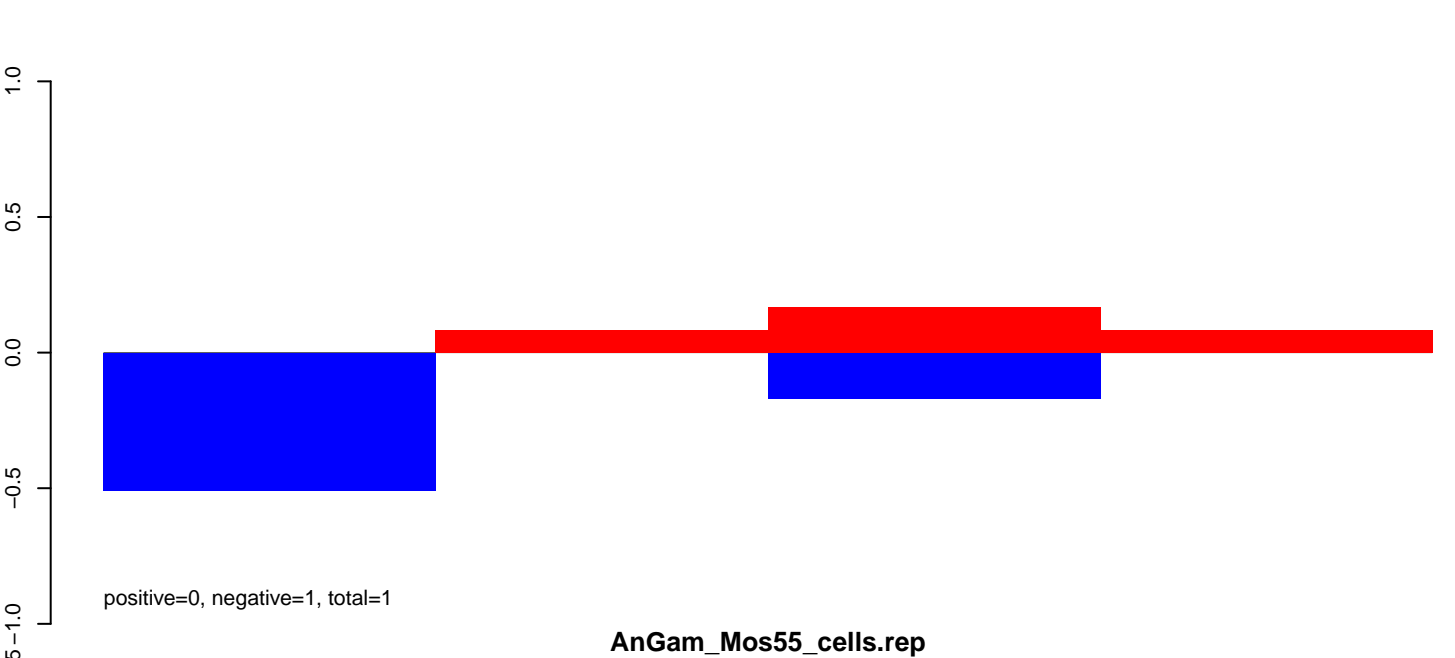
Window size=50, length=345, TE@TF003674-mTA_Ele12:1-345

50 100 150 200 250 300 350

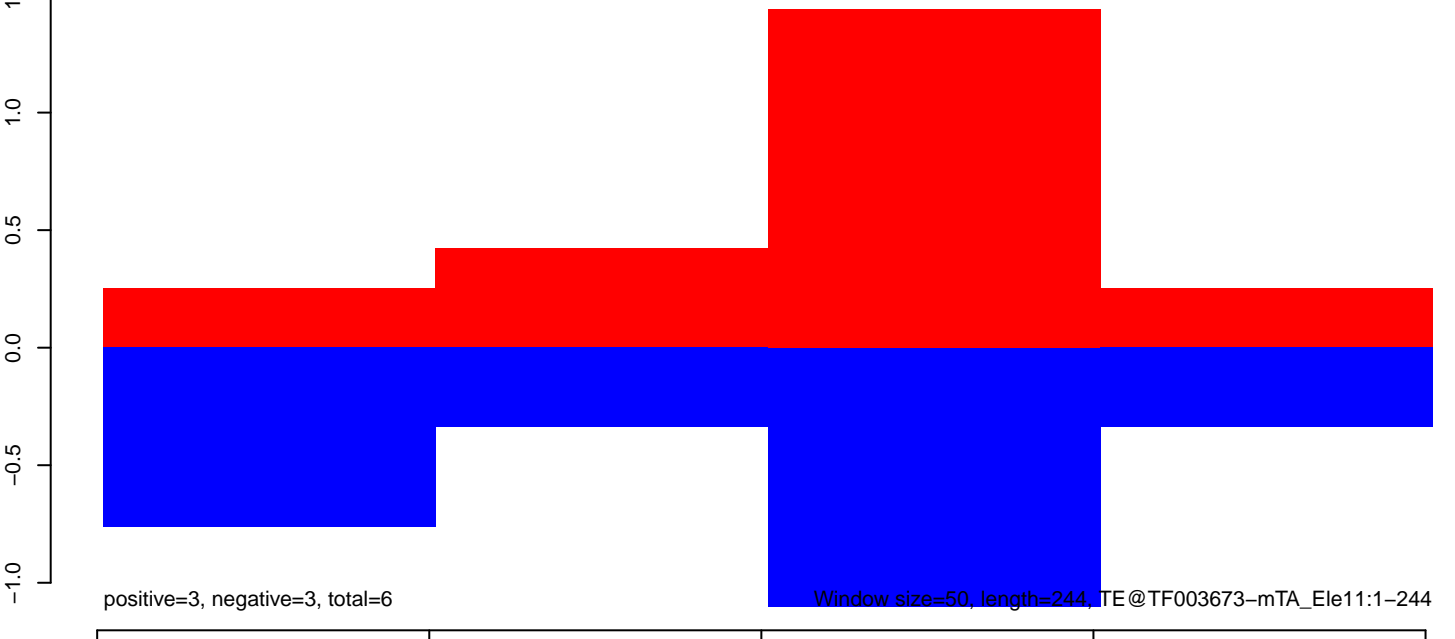
AnGam_Mos55_cells.18_23.rep



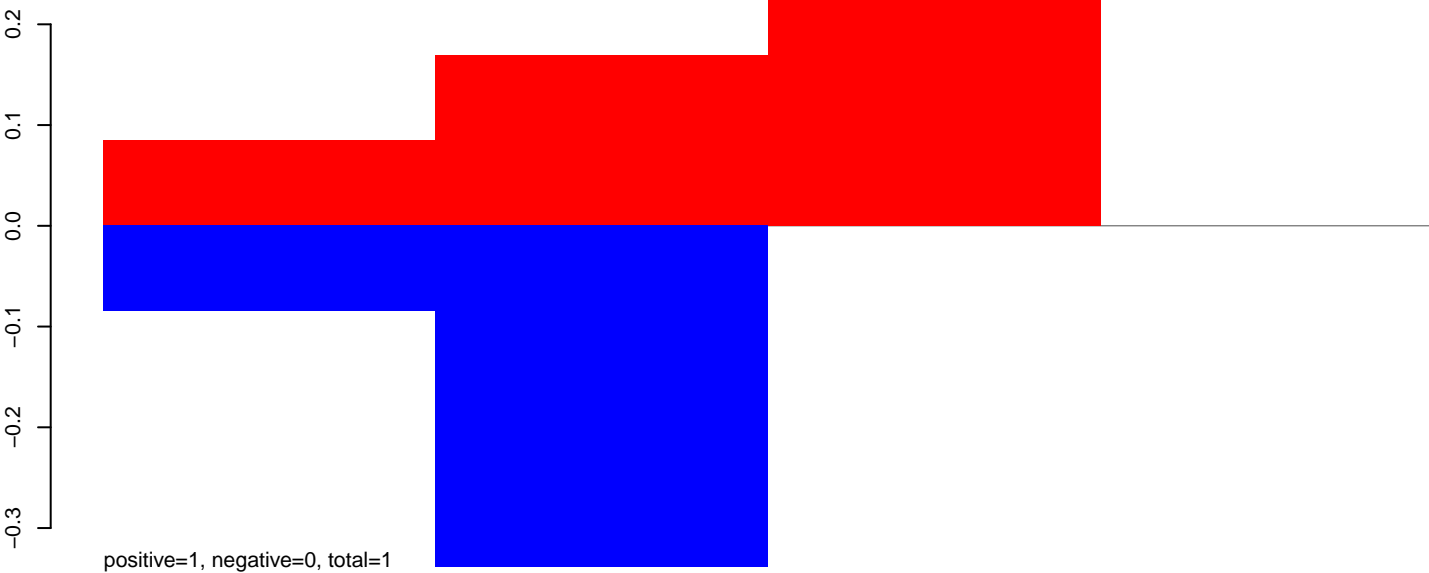
AnGam_Mos55_cells.24_35.rep



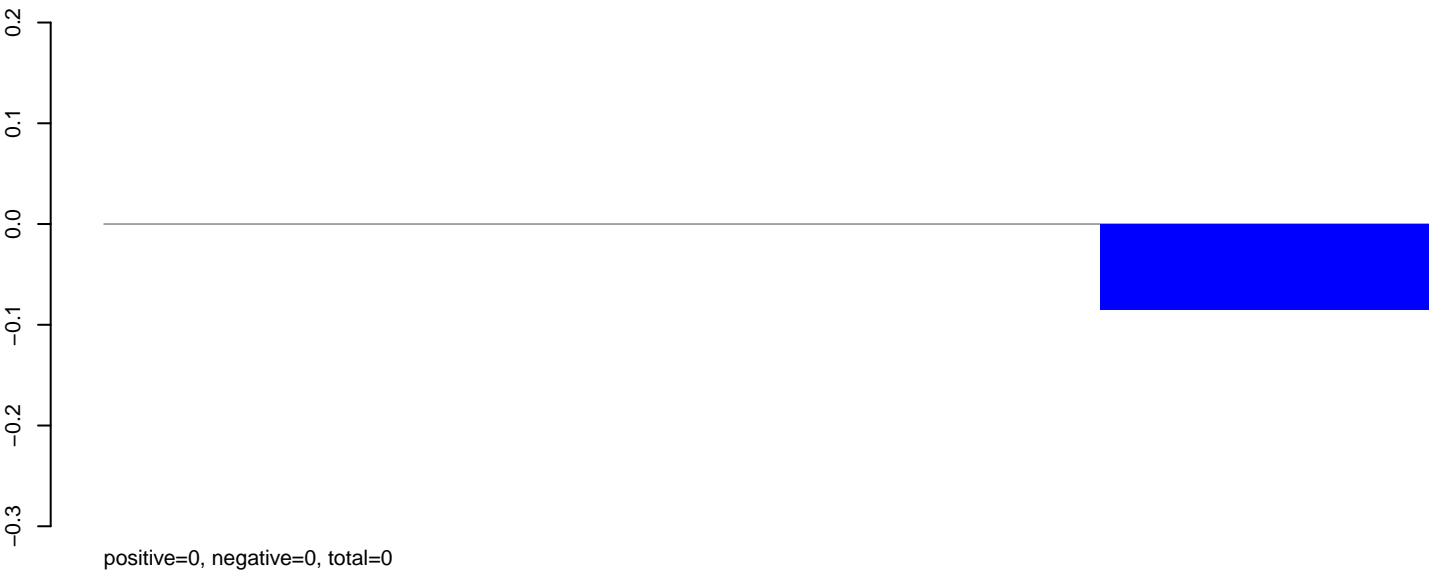
AnGam_Mos55_cells.rep



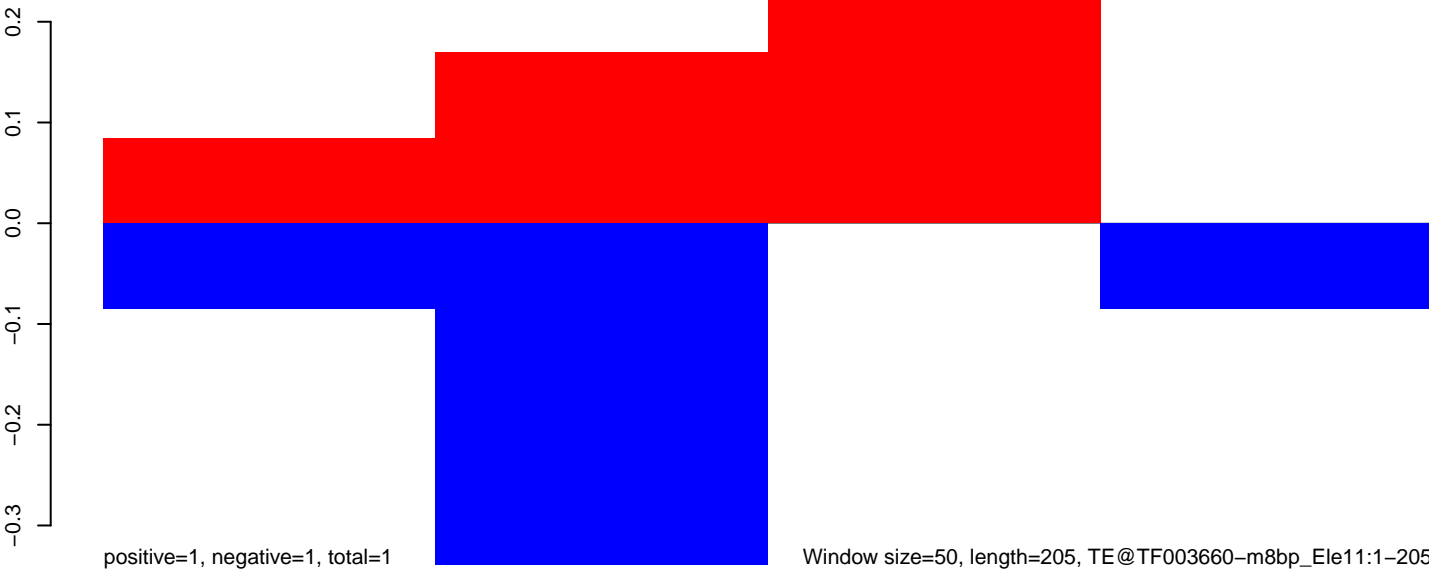
AnGam_Mos55_cells.18_23.rep



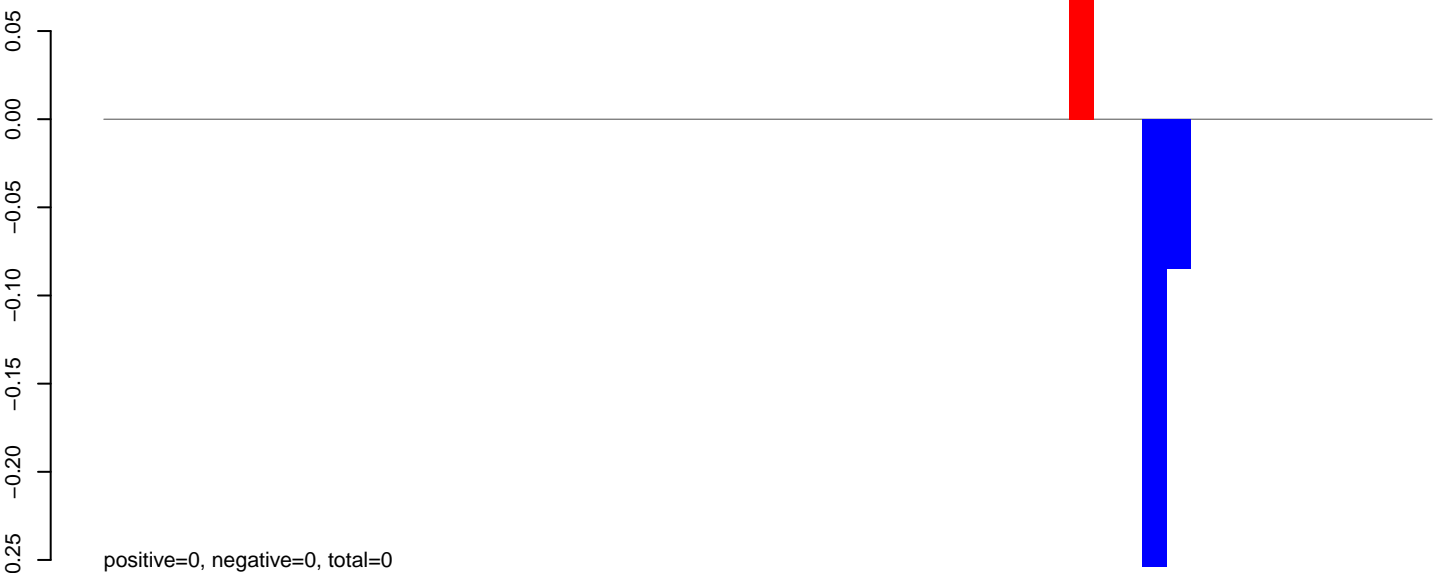
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



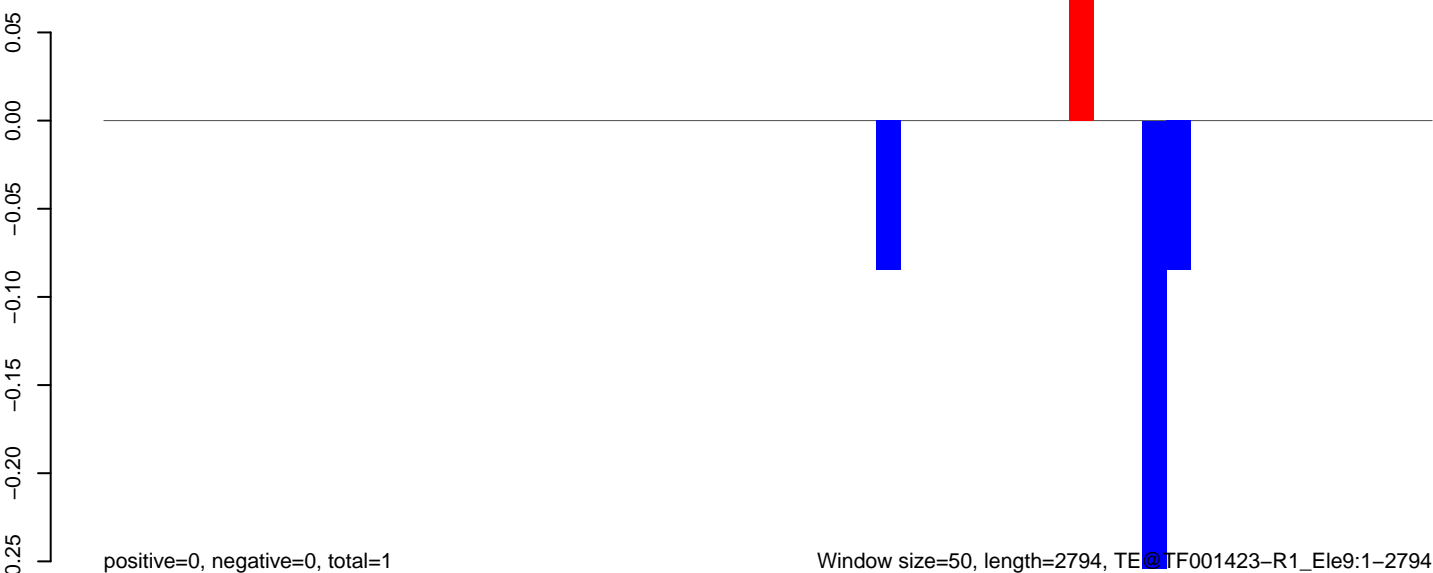
AnGam_Mos55_cells.18_23.rep



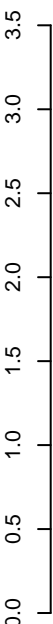
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

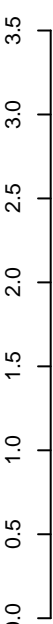


AnGam_Mos55_cells.18_23.rep



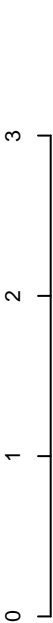
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



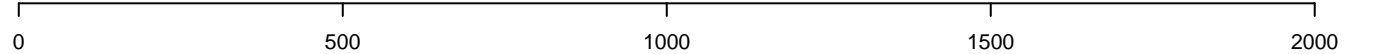
positive=13, negative=0, total=13

AnGam_Mos55_cells.rep

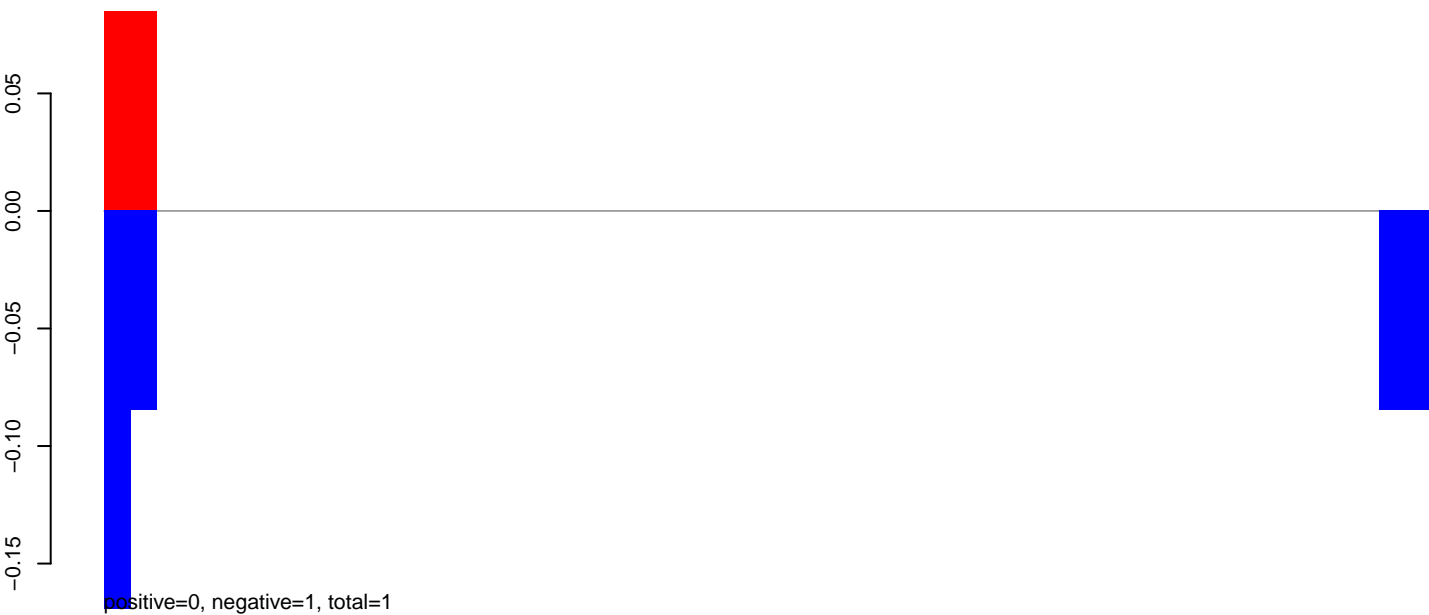


positive=14, negative=0, total=14

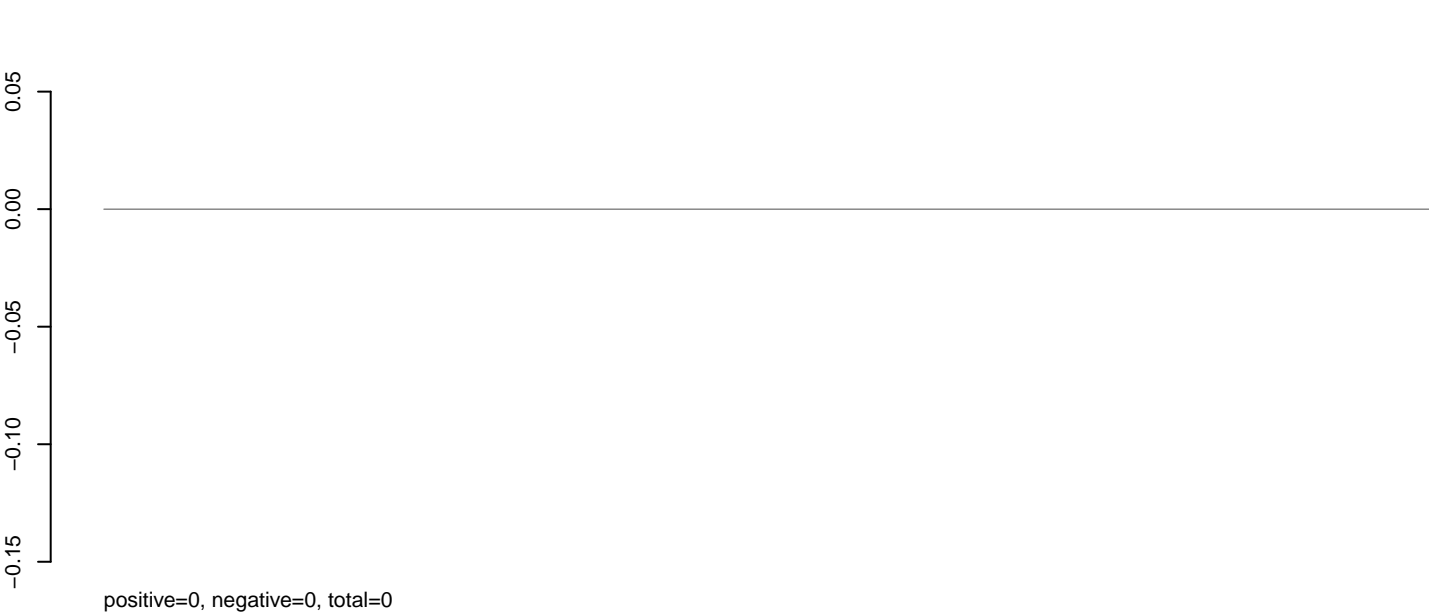
Window size=50, length=2077, TE@TF001381-CR1_Ele5:1-2077



AnGam_Mos55_cells.18_23.rep



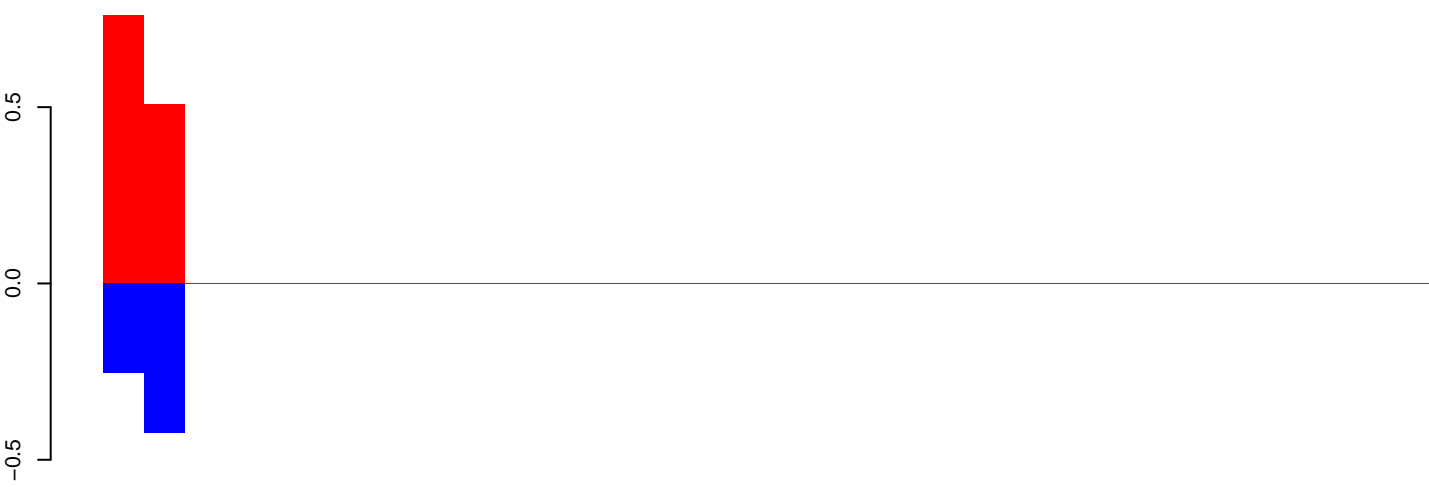
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

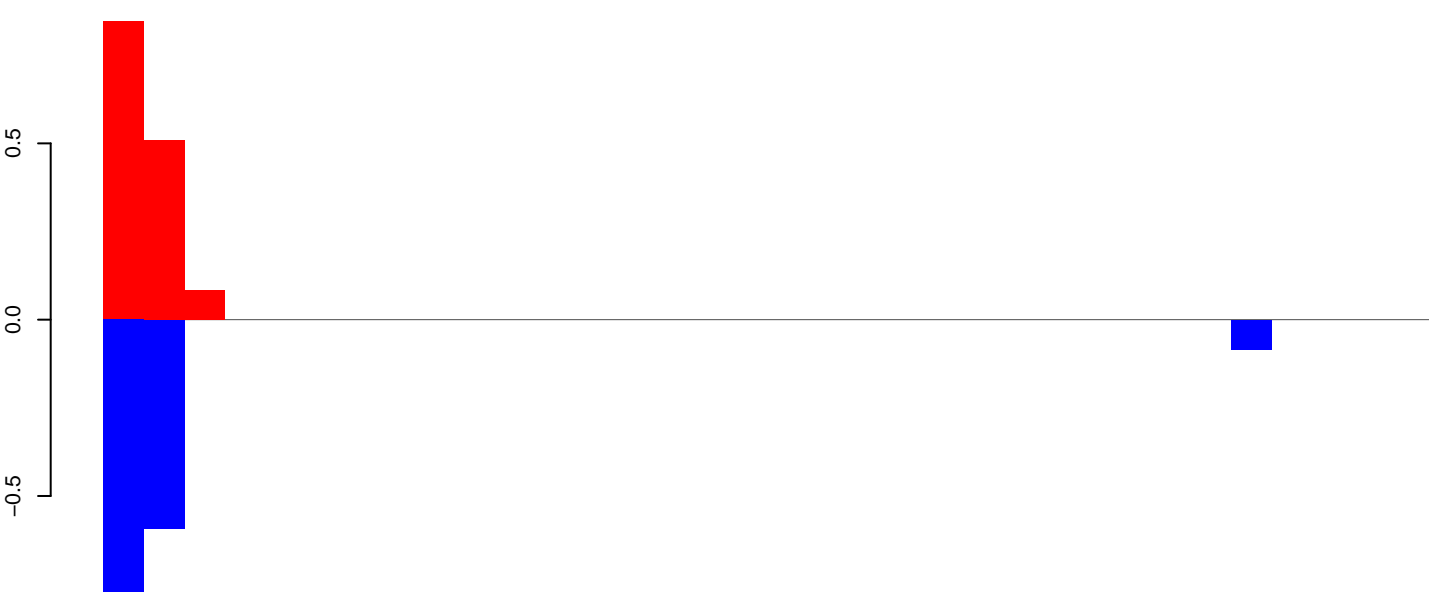


AnGam_Mos55_cells.18_23.rep



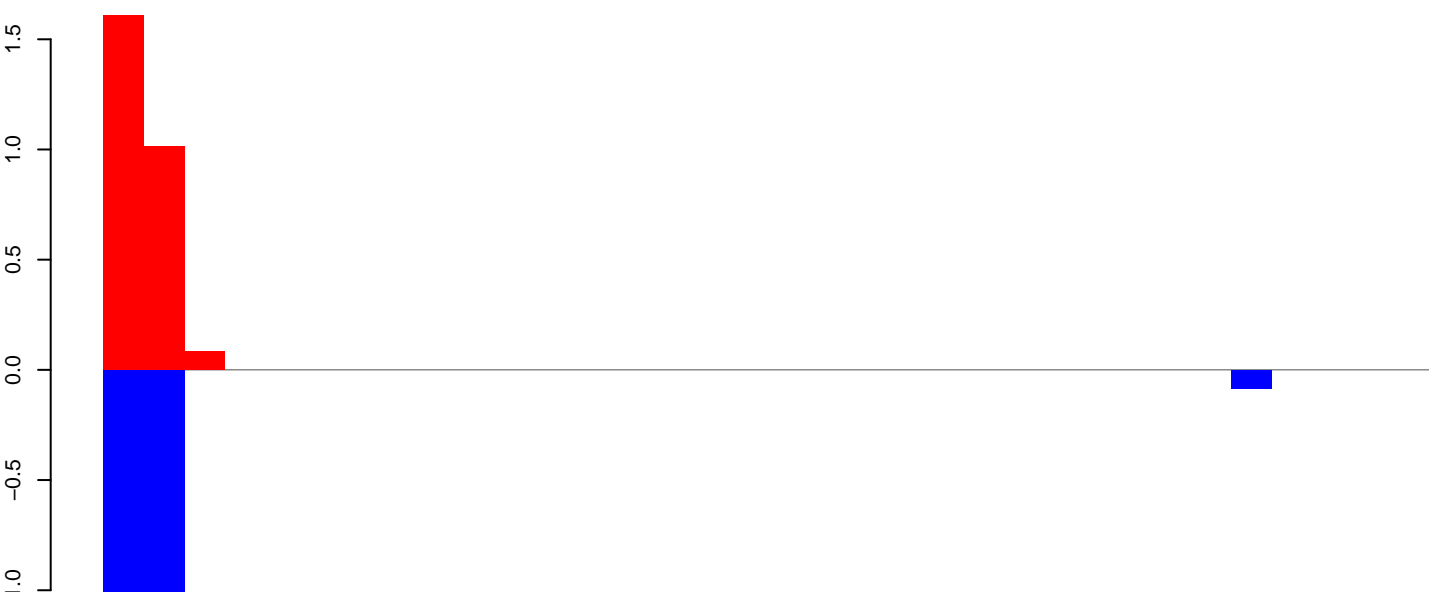
positive=1, negative=1, total=2

AnGam_Mos55_cells.24_35.rep



positive=1, negative=2, total=3

AnGam_Mos55_cells.rep



positive=3, negative=2, total=5

Window size=50, length=1679, TE@TF001208-mar_Ele31:1-1679

AnGam_Mos55_cells.18_23.rep

10
0
-10
-20

positive=9, negative=9, total=17

AnGam_Mos55_cells.24_35.rep

10
0
-10
-20

positive=42, negative=39, total=81

AnGam_Mos55_cells.rep

20
10
0
-10
-20

positive=51, negative=48, total=99

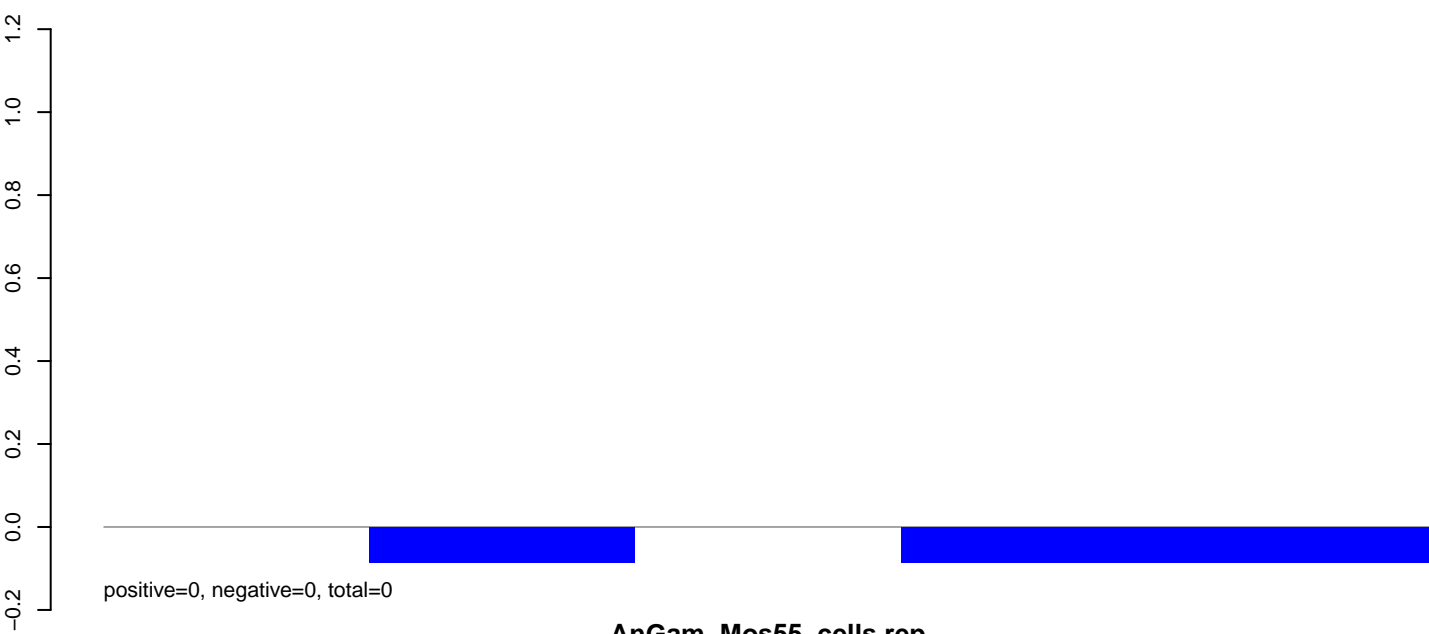
Window size=50, length=206, TE@SINEX-1_AG:1-206

50 100 150 200 250

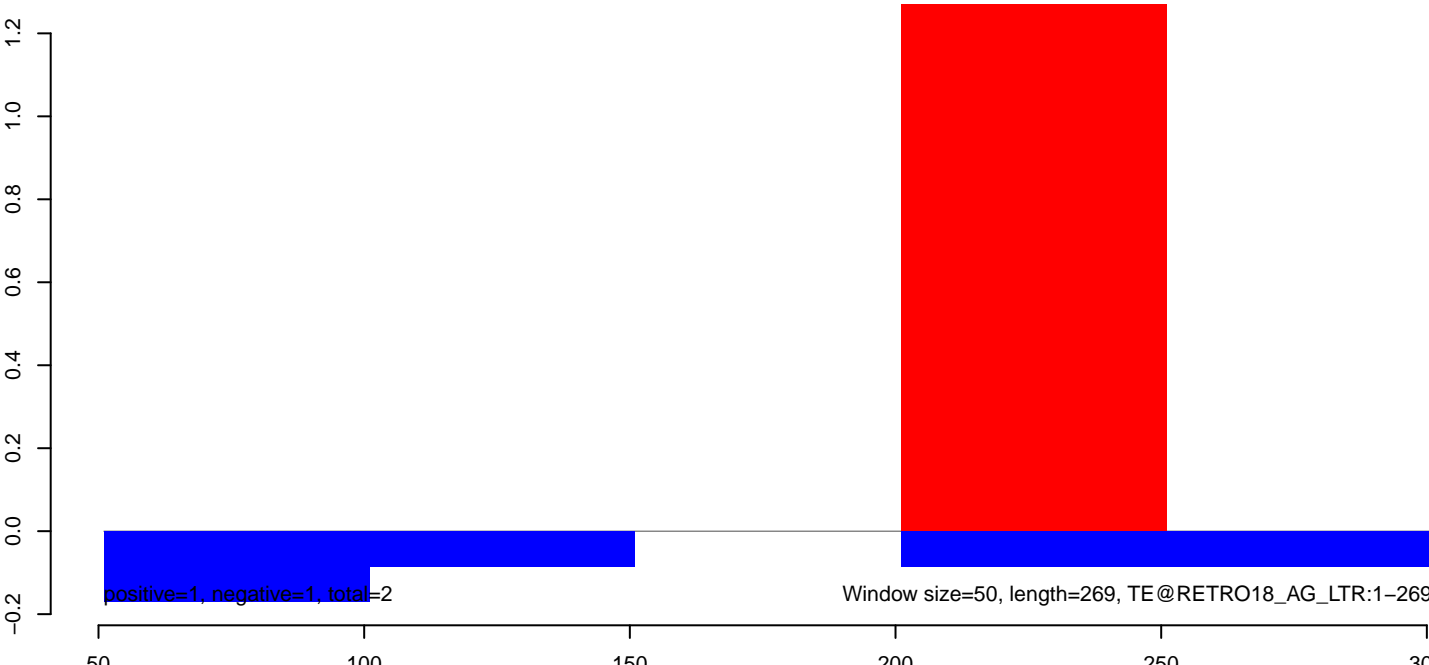
AnGam_Mos55_cells.18_23.rep



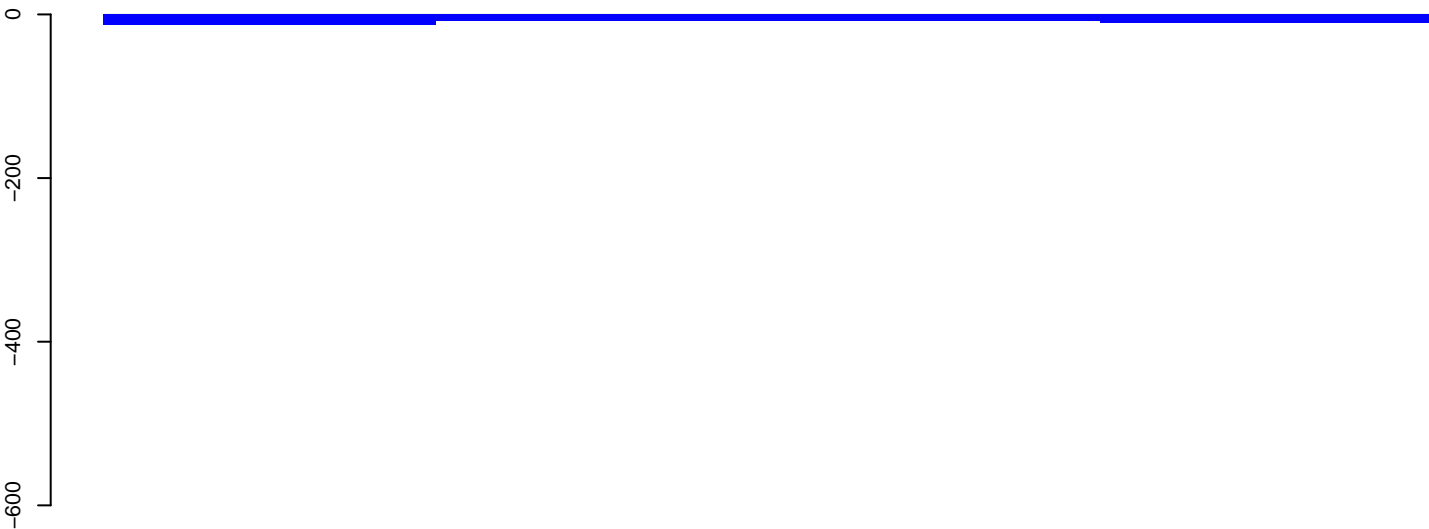
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

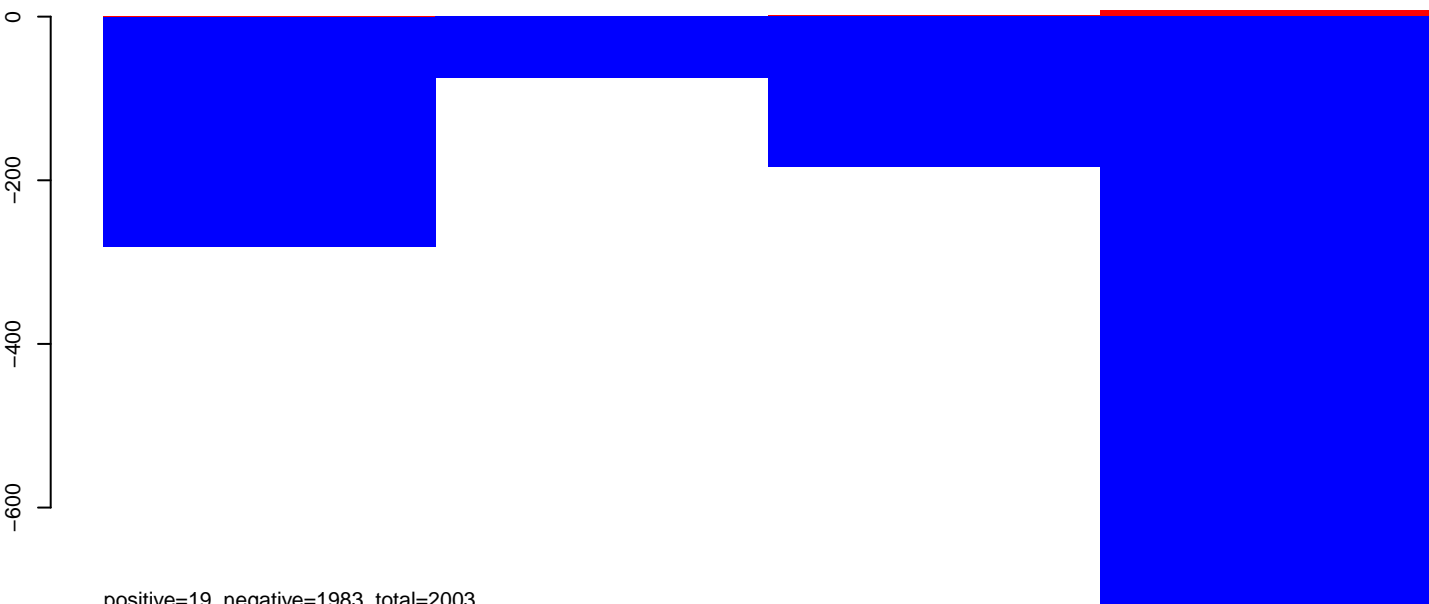


AnGam_Mos55_cells.18_23.rep



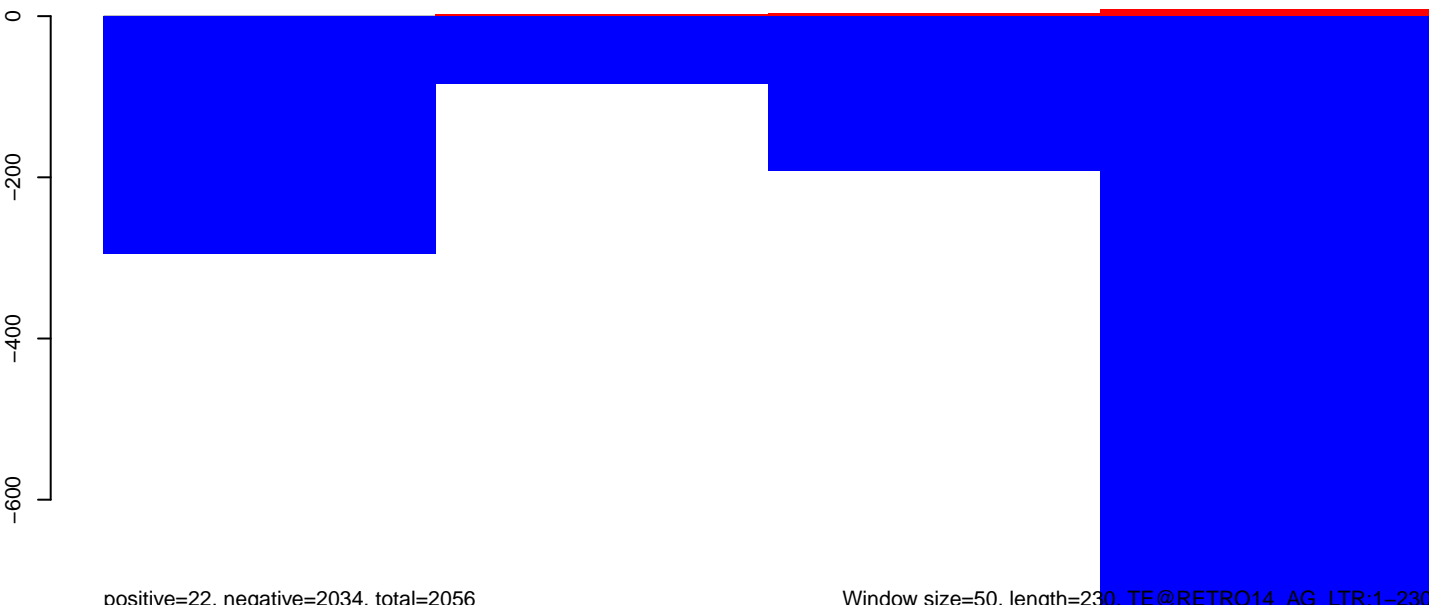
positive=3, negative=50, total=53

AnGam_Mos55_cells.24_35.rep



positive=19, negative=1983, total=2003

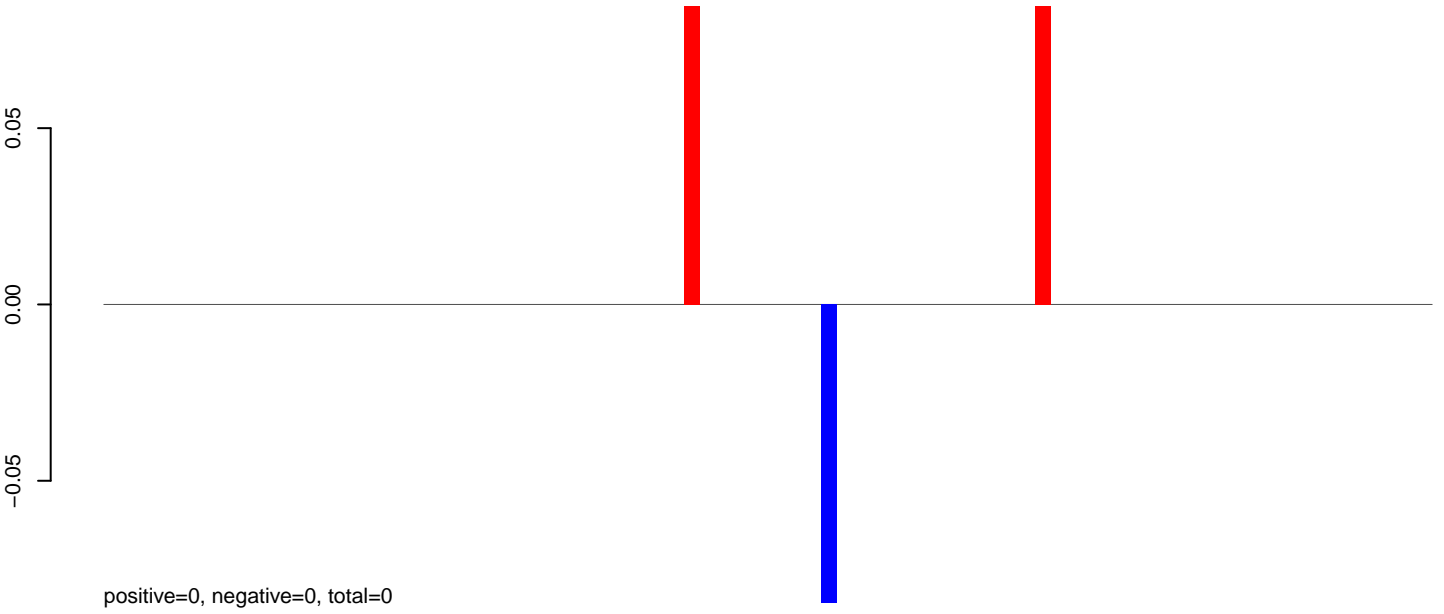
AnGam_Mos55_cells.rep



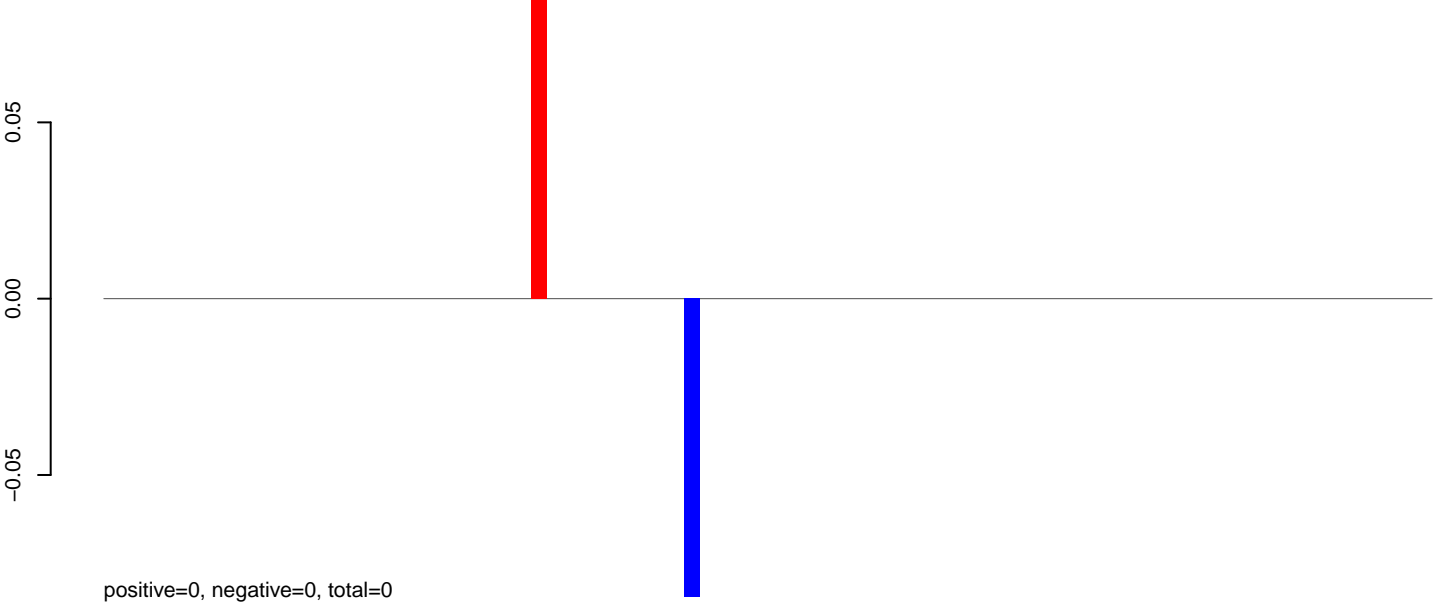
positive=22, negative=2034, total=2056

Window size=50, length=230, TE@RETRO14_AG_LTR:1-230

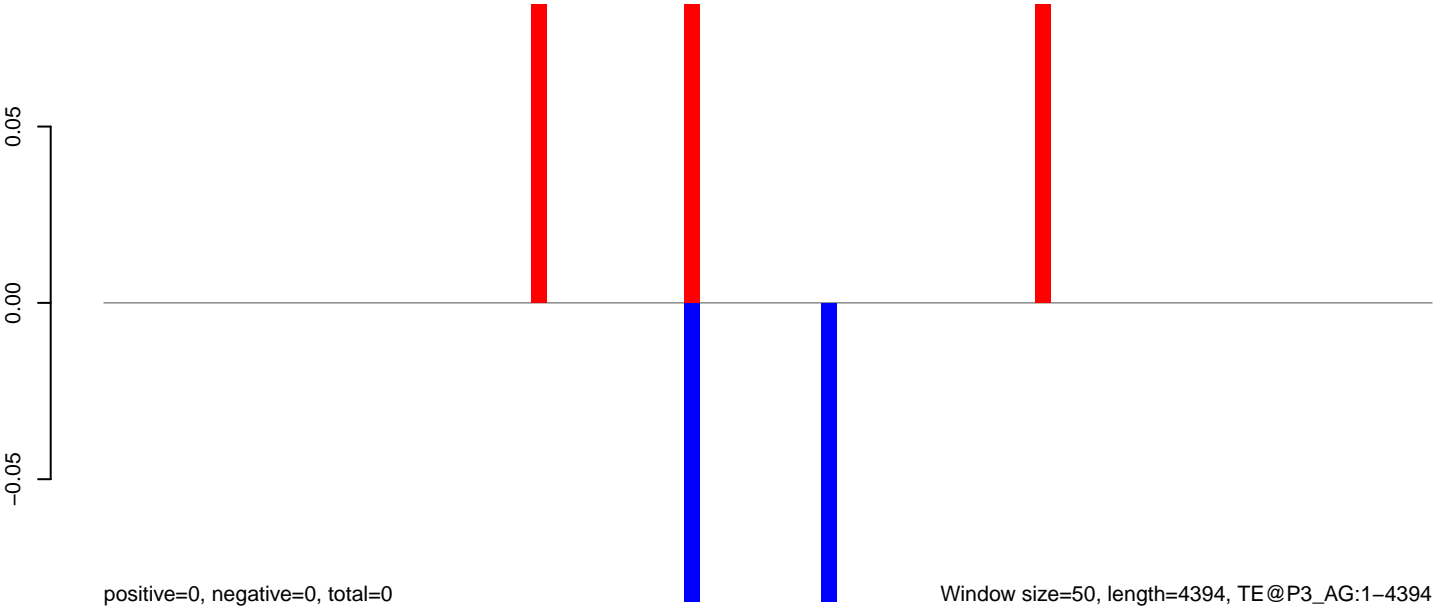
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

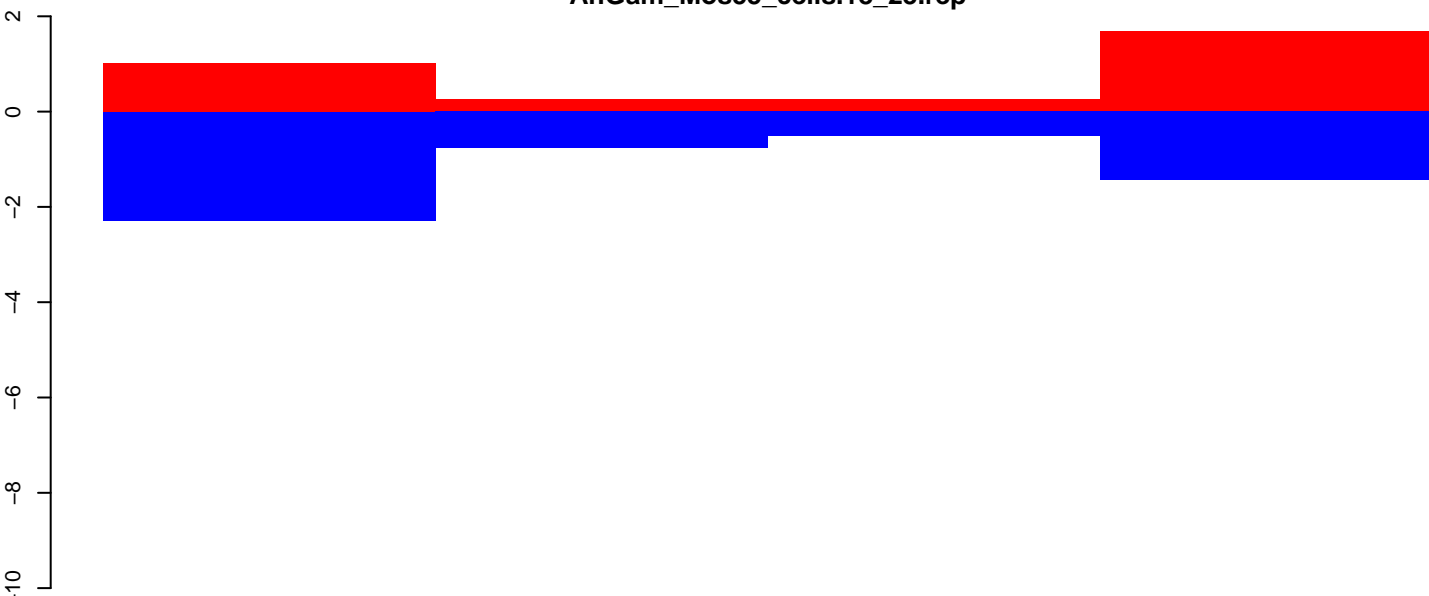


AnGam_Mos55_cells.rep

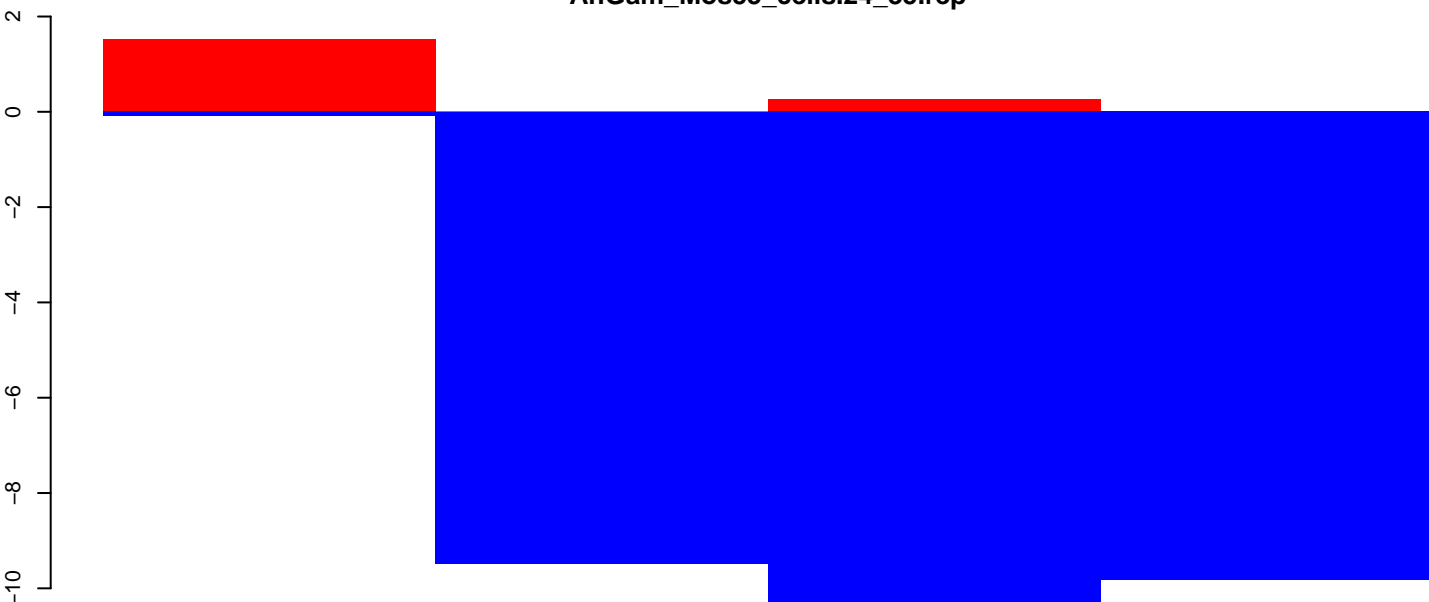


0 1000 2000 3000 4000

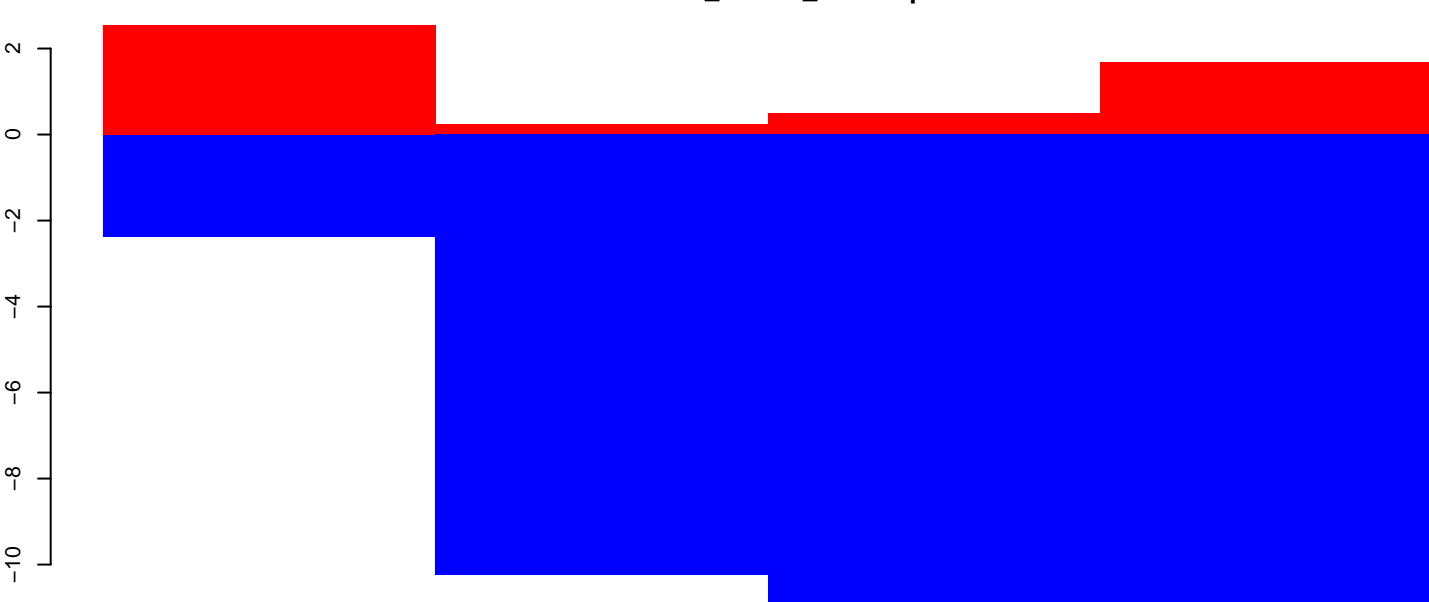
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



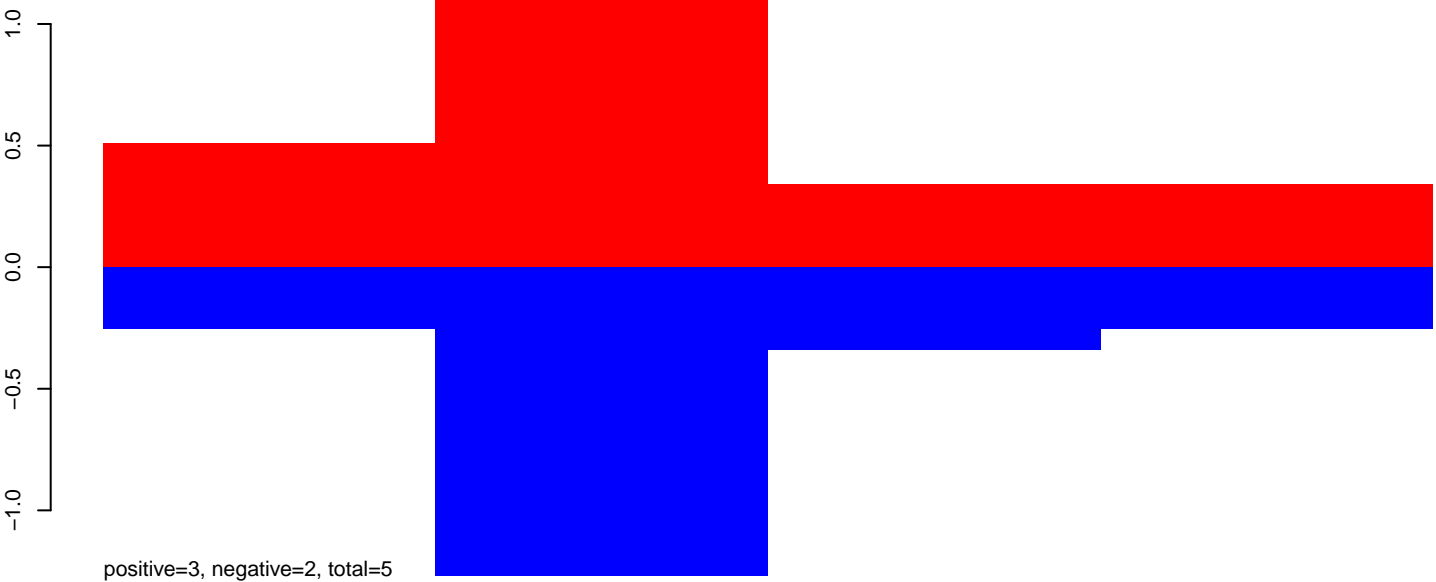
AnGam_Mos55_cells.rep



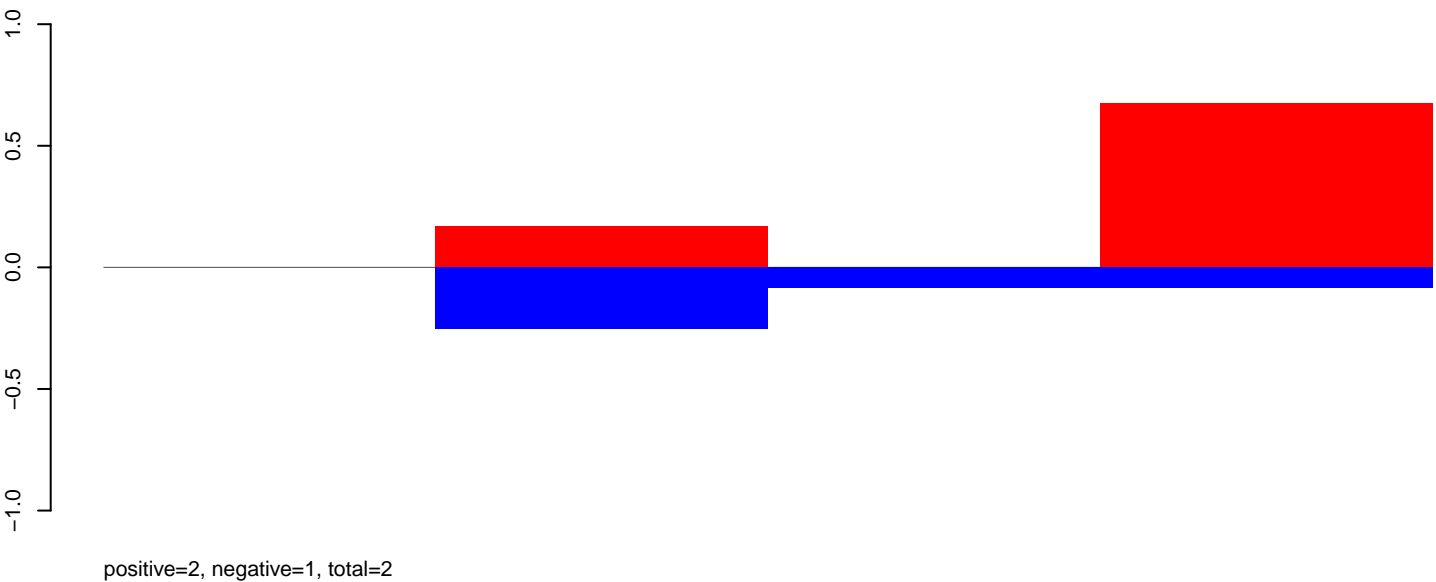
Window size=50, length=249, TE@Mariner-N15_AG:1-249

50 100 150 200 250

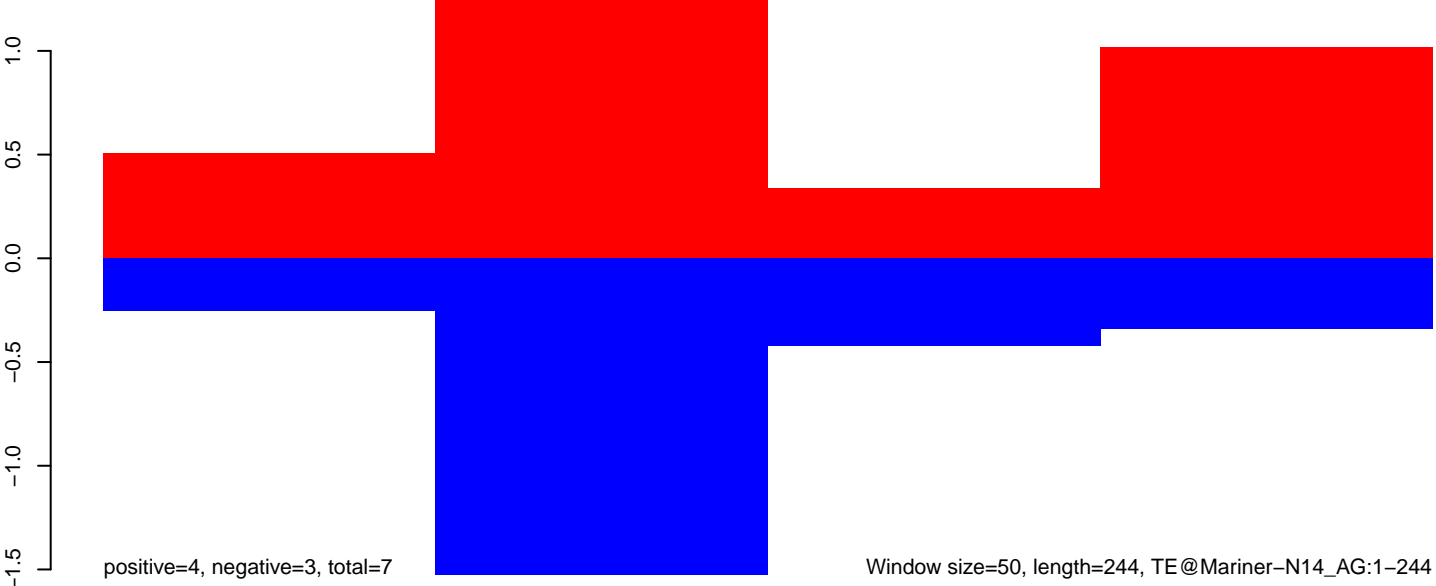
AnGam_Mos55_cells.18_23.rep



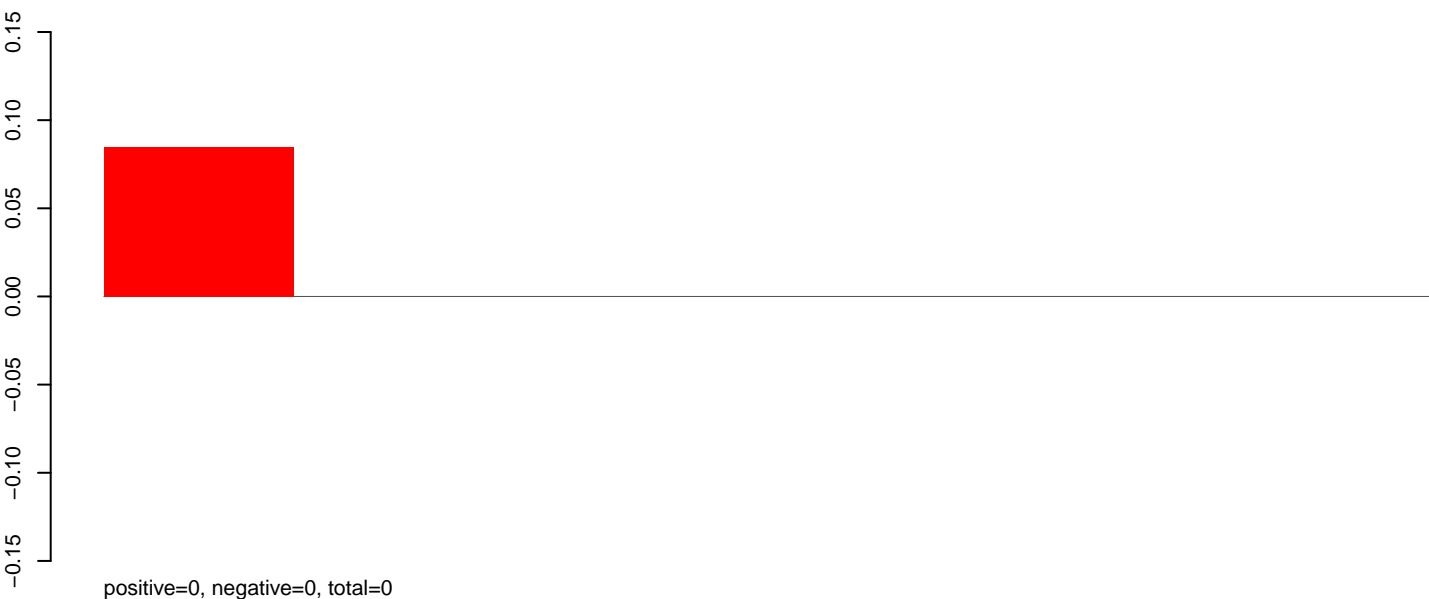
AnGam_Mos55_cells.24_35.rep



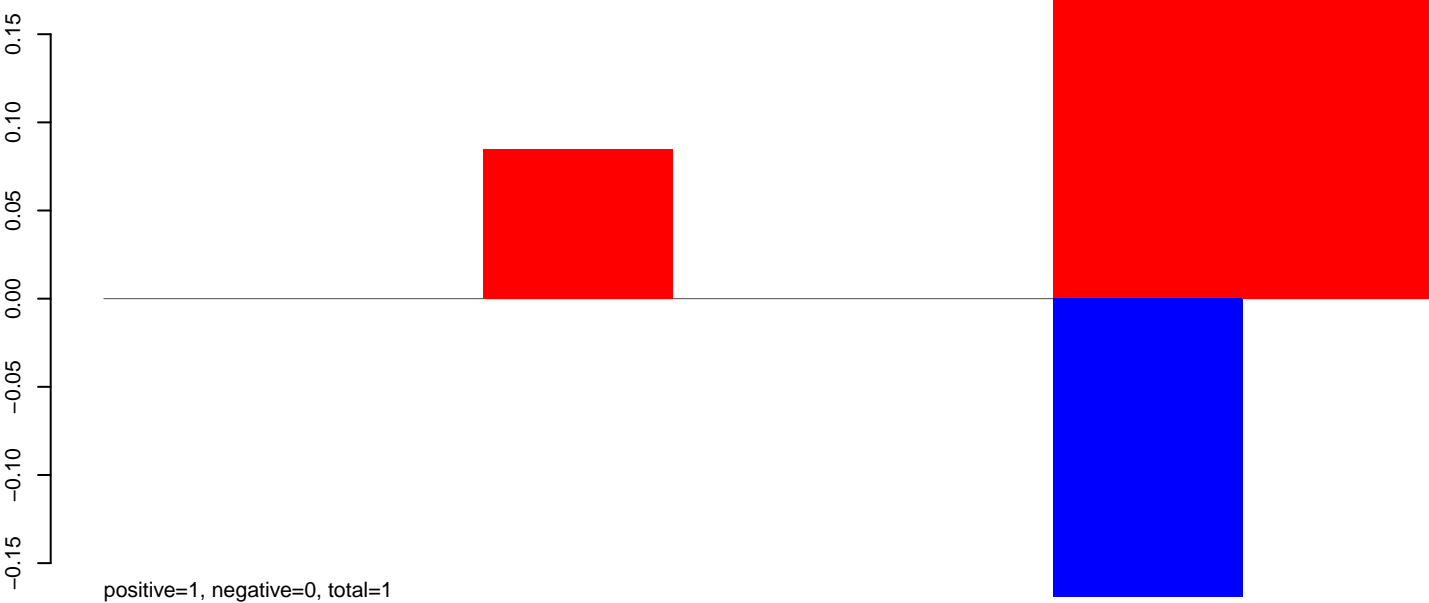
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



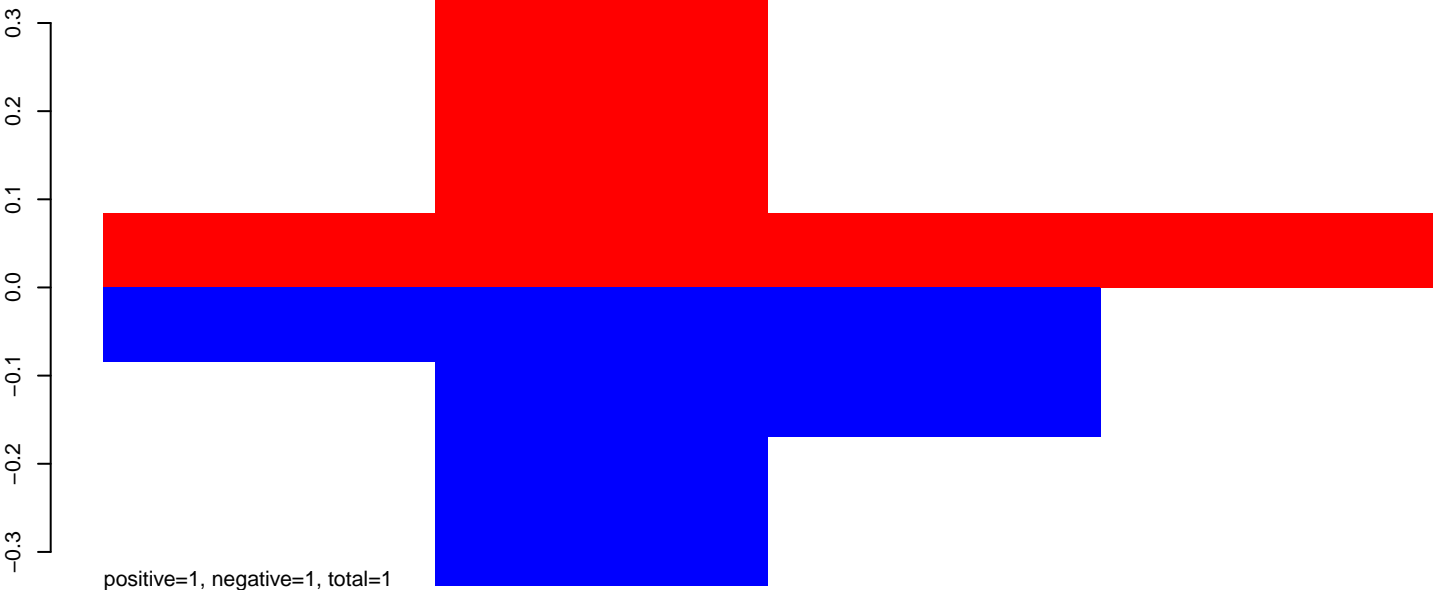
AnGam_Mos55_cells.24_35.rep



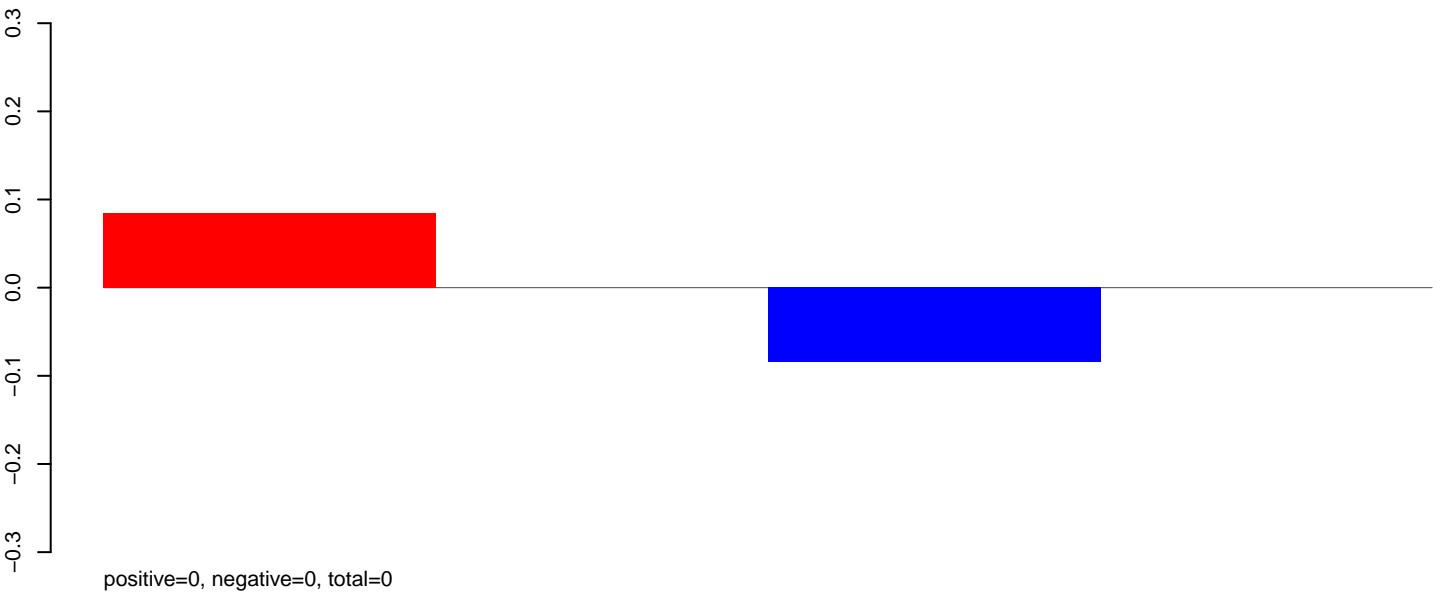
AnGam_Mos55_cells.rep



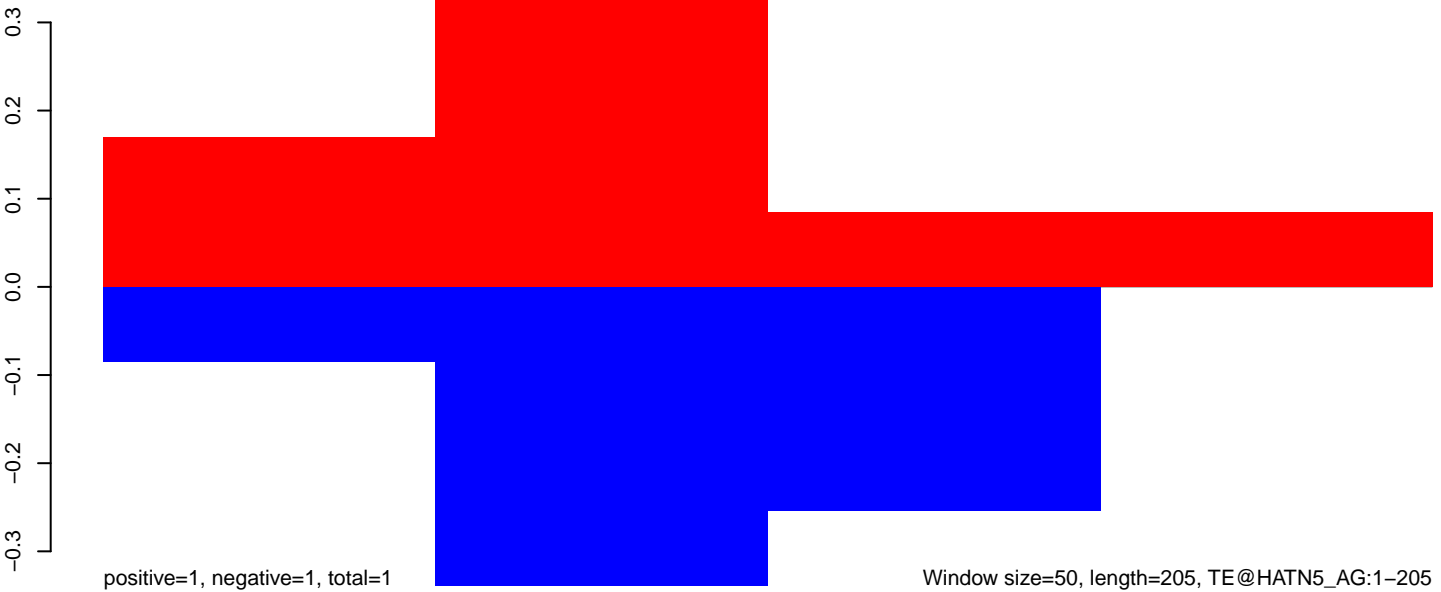
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

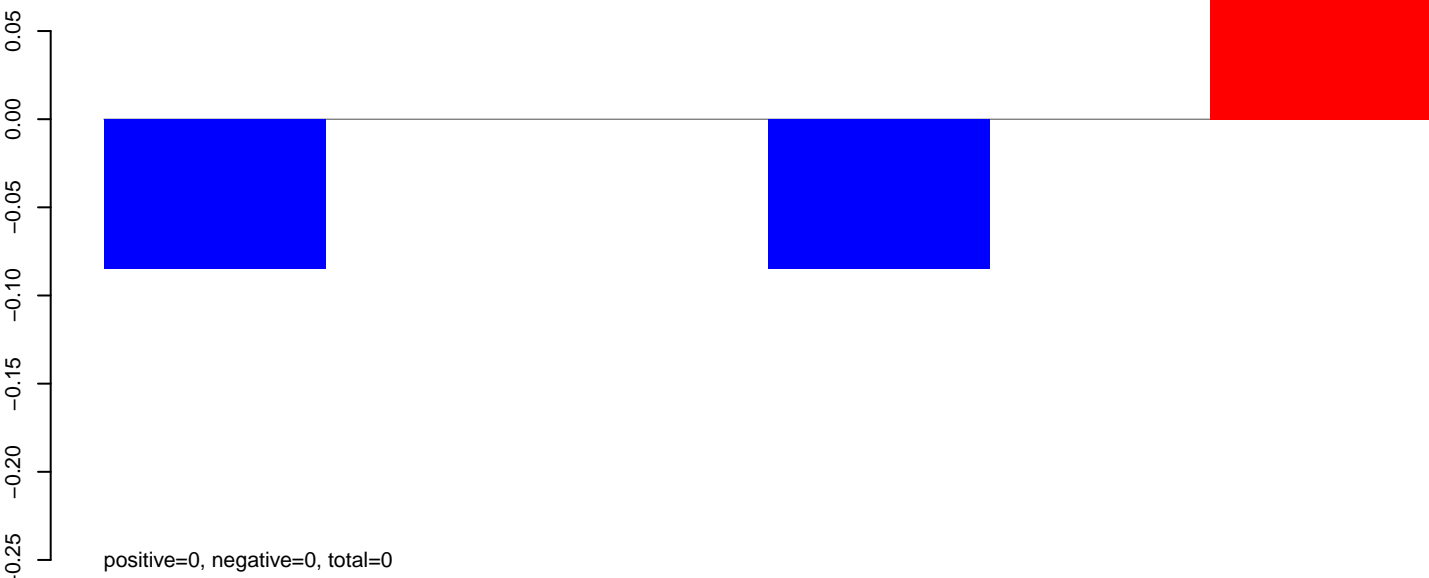


AnGam_Mos55_cells.rep



Window size=50, length=205, TE@HATN5_AG:1-205

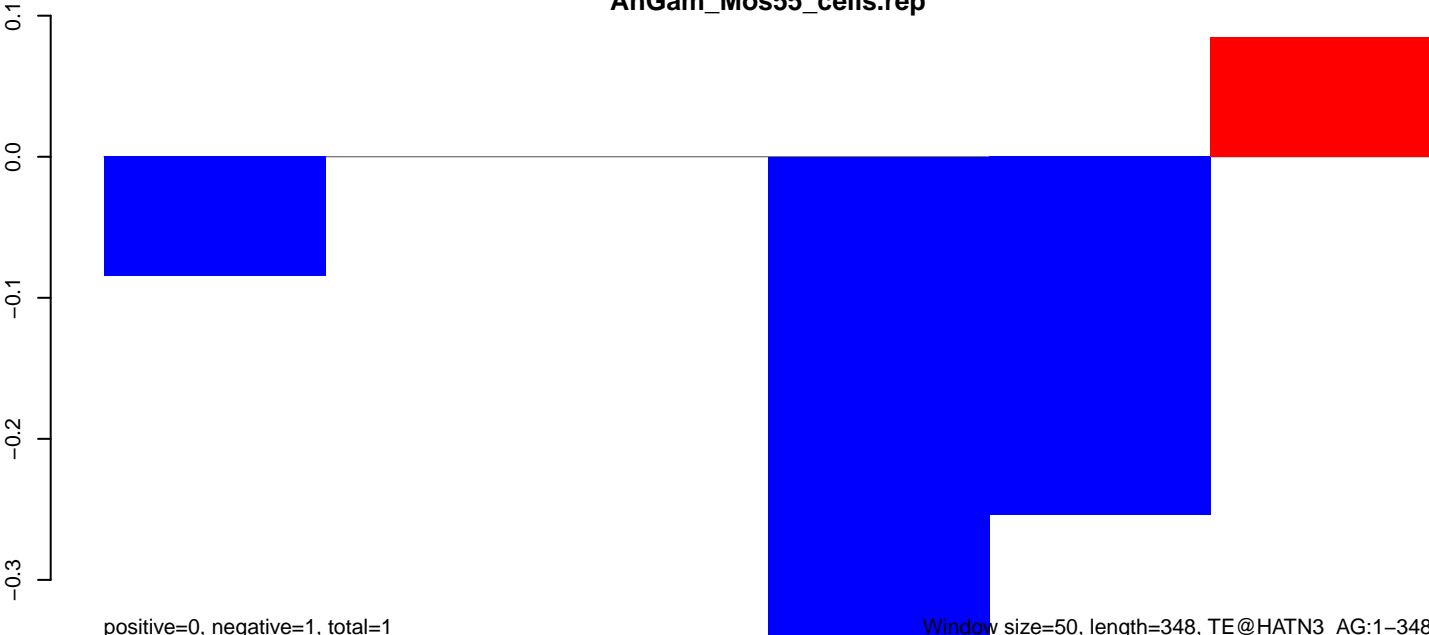
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



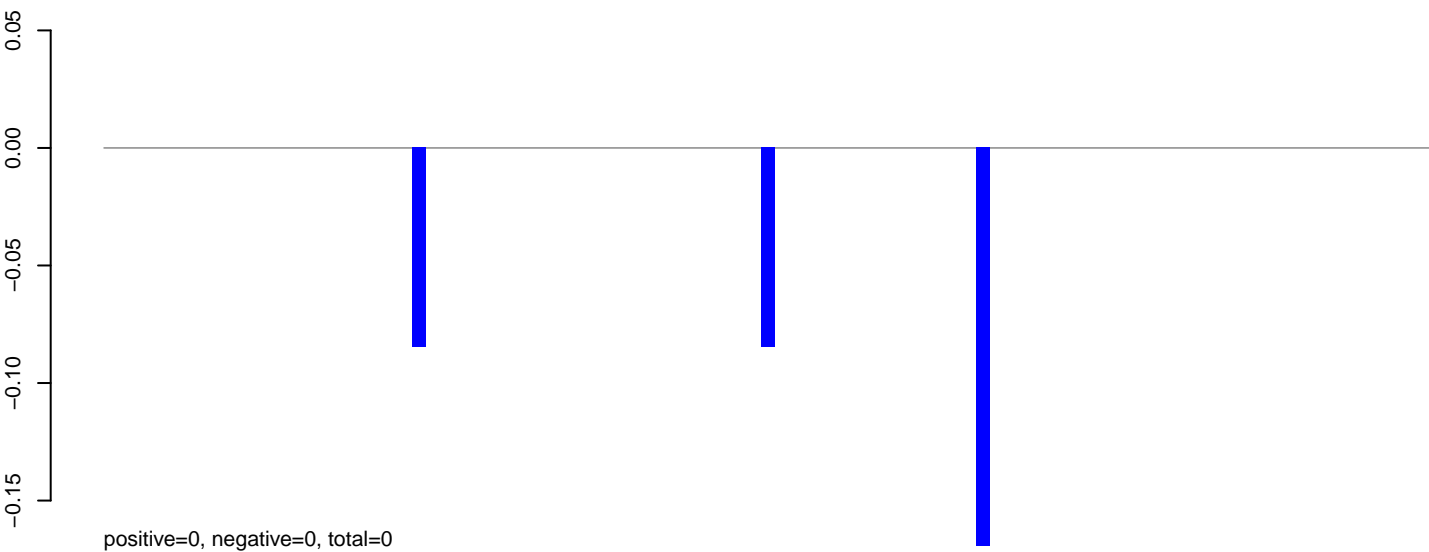
AnGam_Mos55_cells.rep



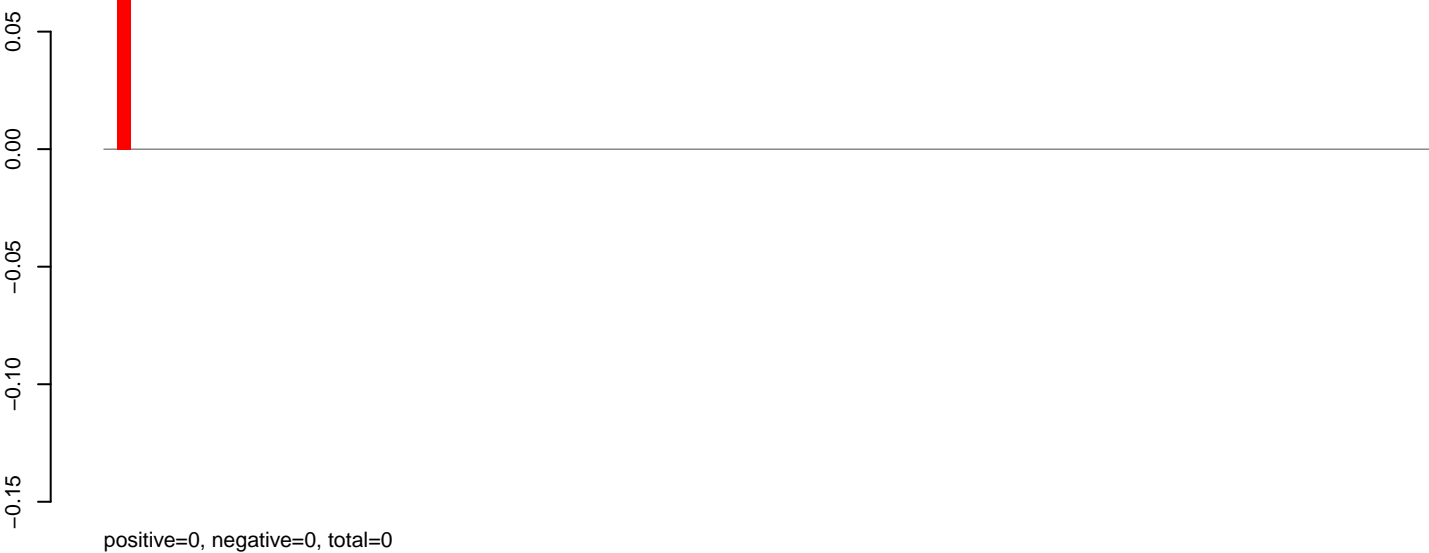
Window size=50, length=348, TE@HATN3_AG:1-348

50 100 150 200 250 300 350

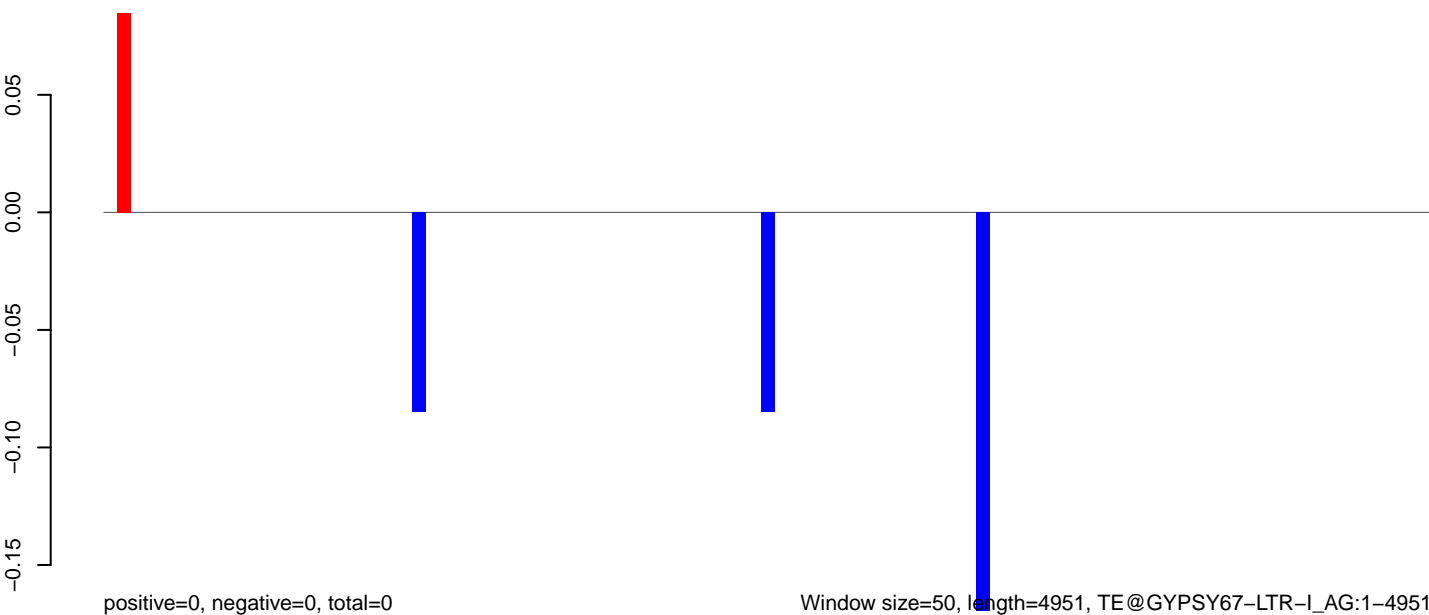
AnGam_Mos55_cells.18_23.rep



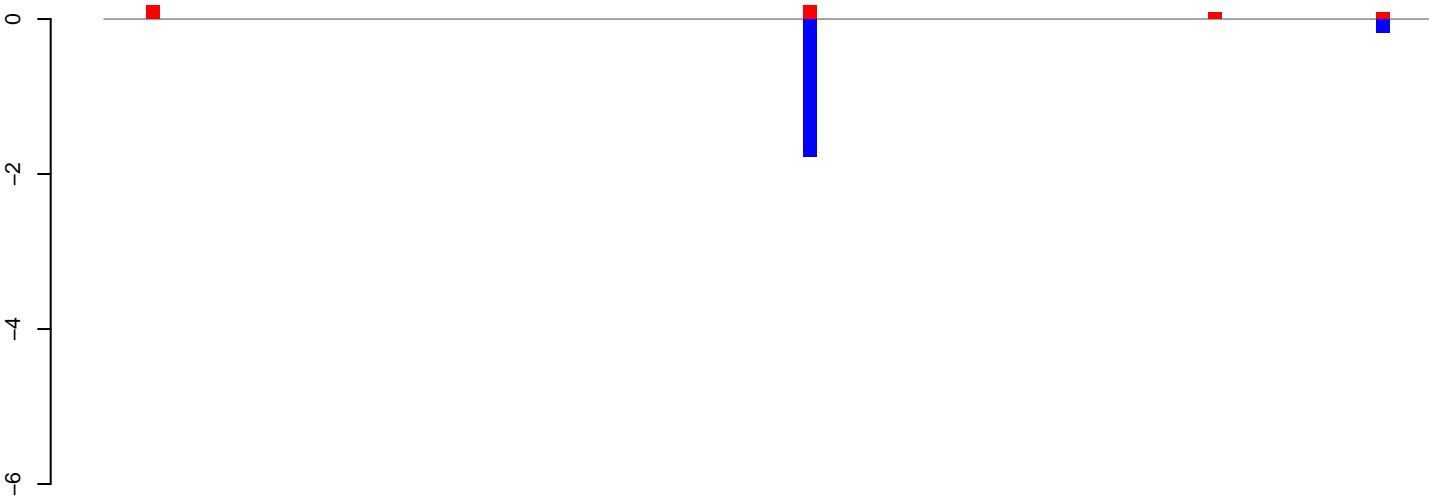
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

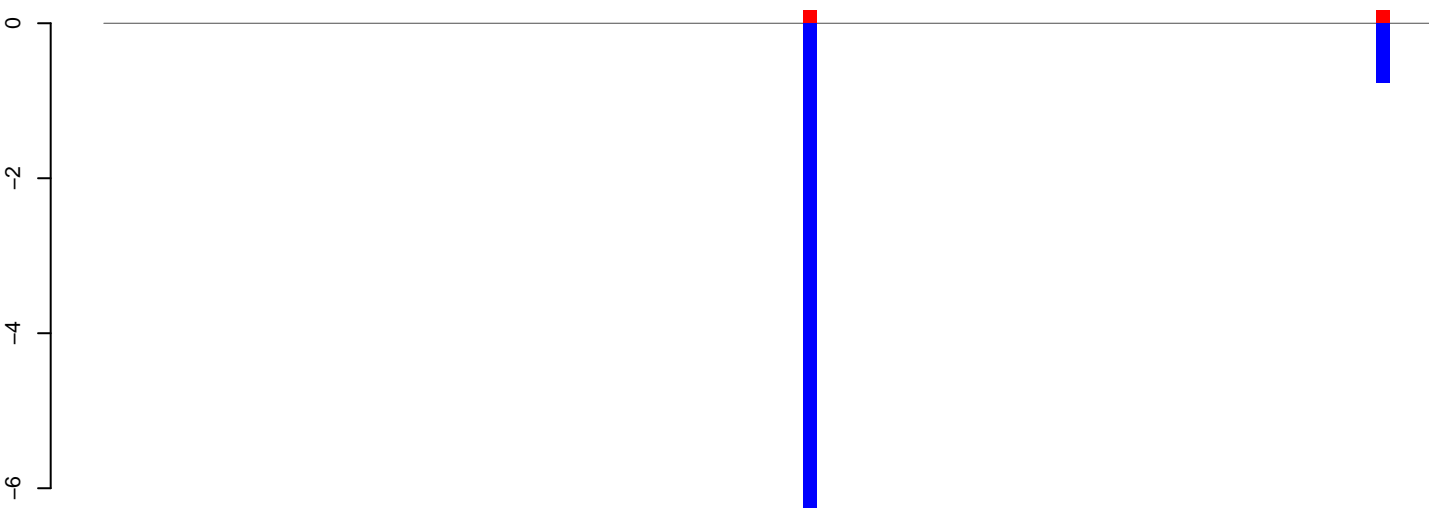


AnGam_Mos55_cells.18_23.rep



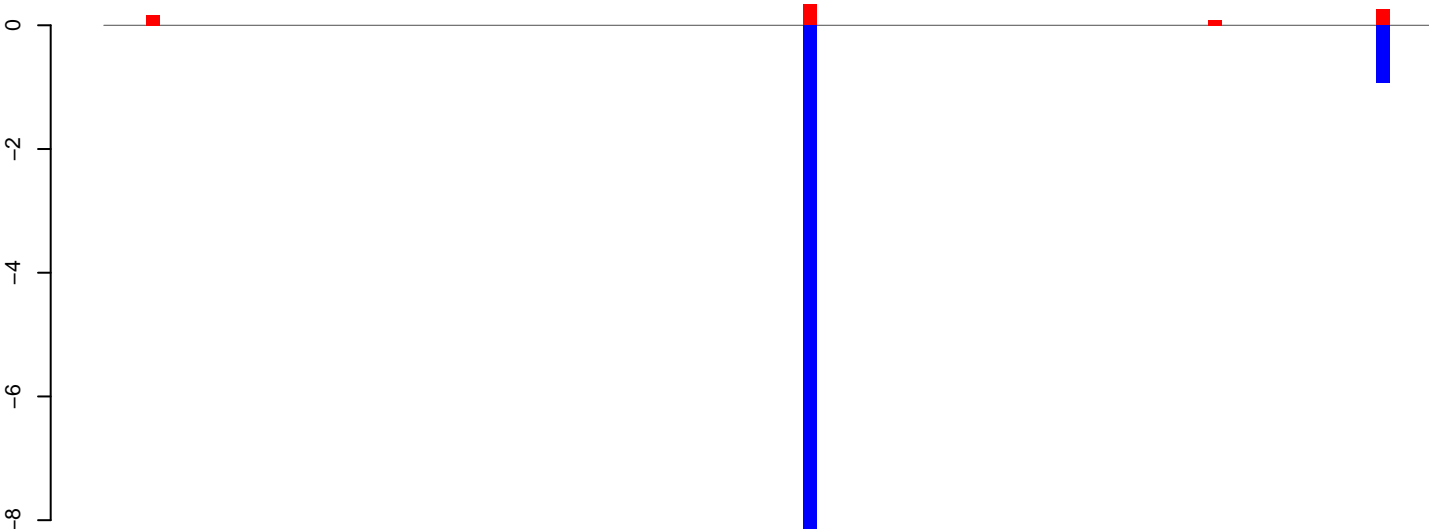
positive=1, negative=2, total=2

AnGam_Mos55_cells.24_35.rep



positive=0, negative=8, total=9

AnGam_Mos55_cells.rep

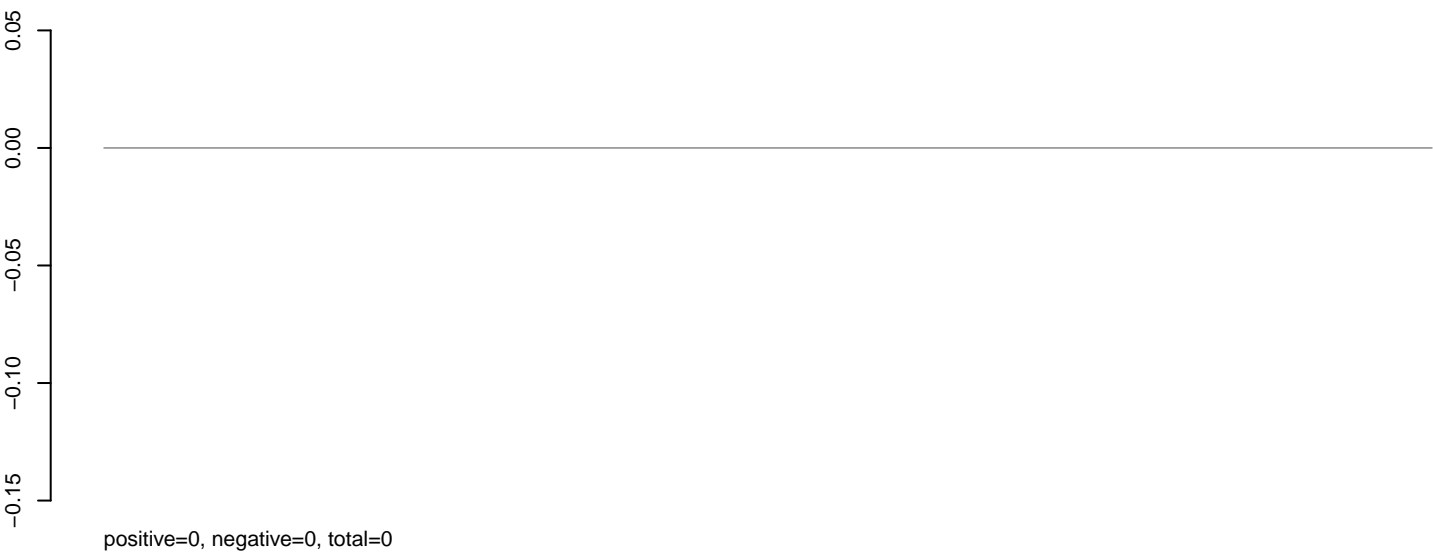


positive=1, negative=10, total=11

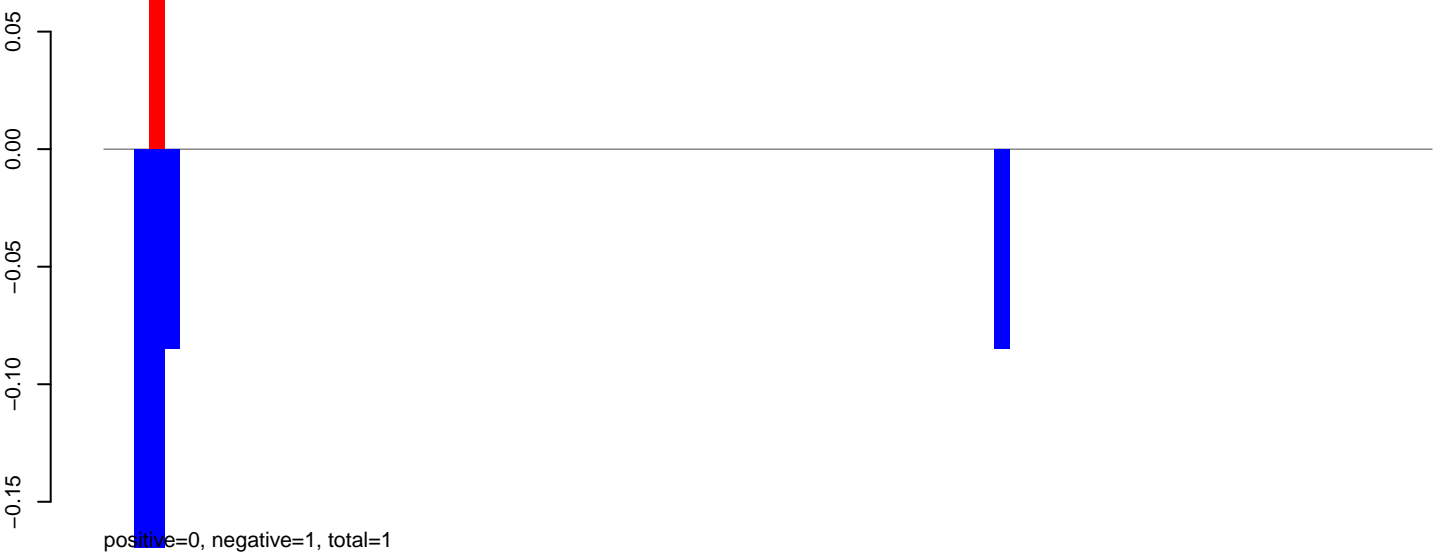
Window size=50, length=4799, TE@GYPSY6-LTR-I_AG:1-4799

0 1000 2000 3000 4000

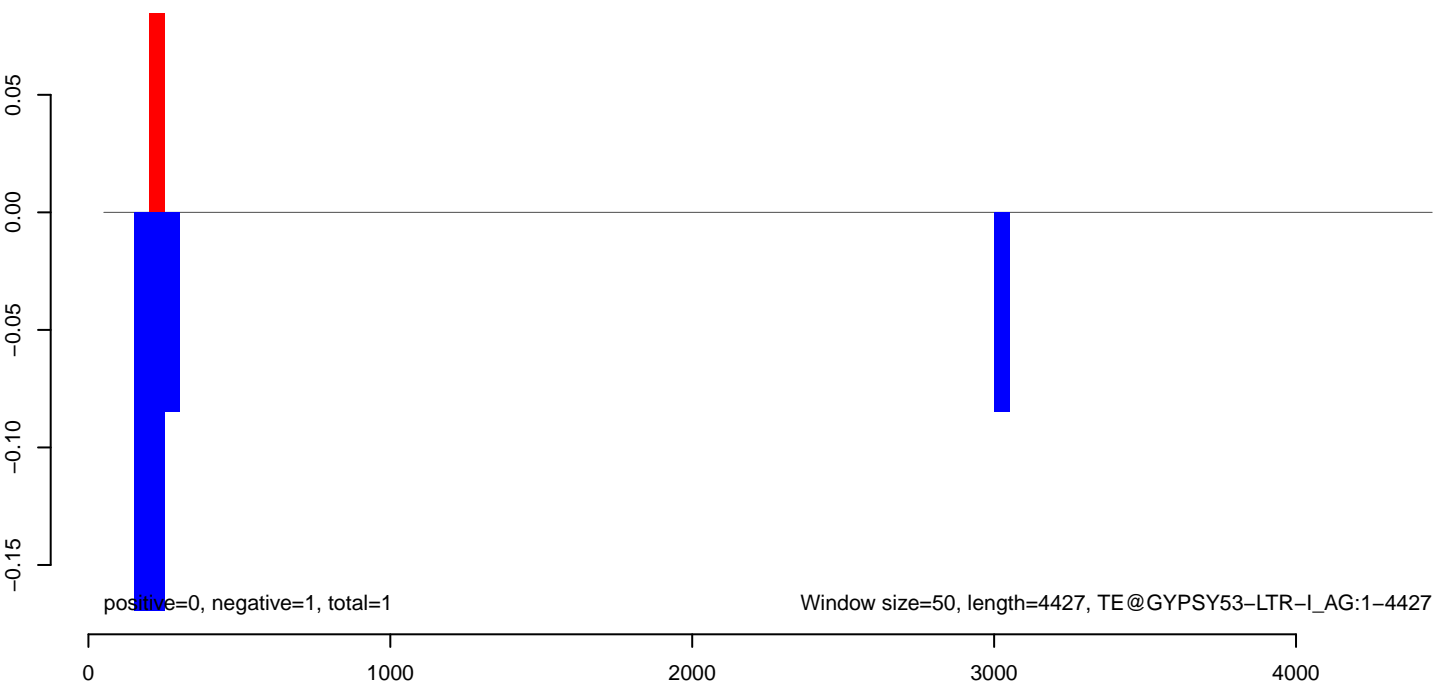
AnGam_Mos55_cells.18_23.rep



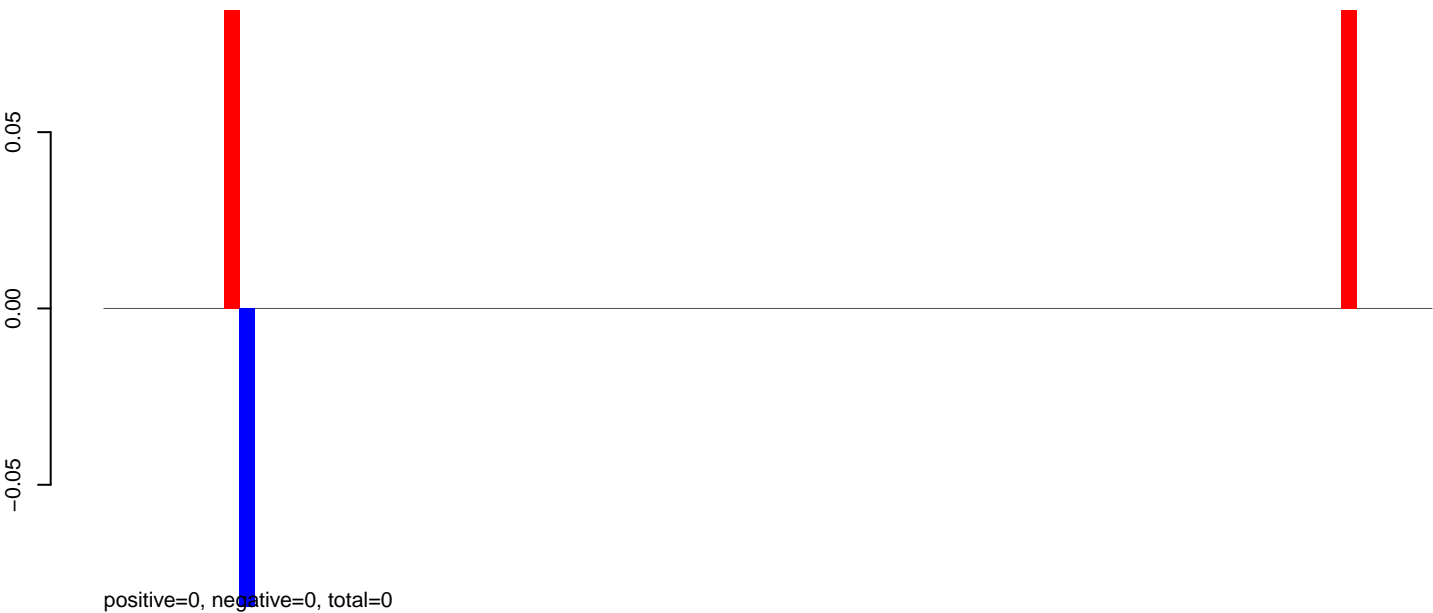
AnGam_Mos55_cells.24_35.rep



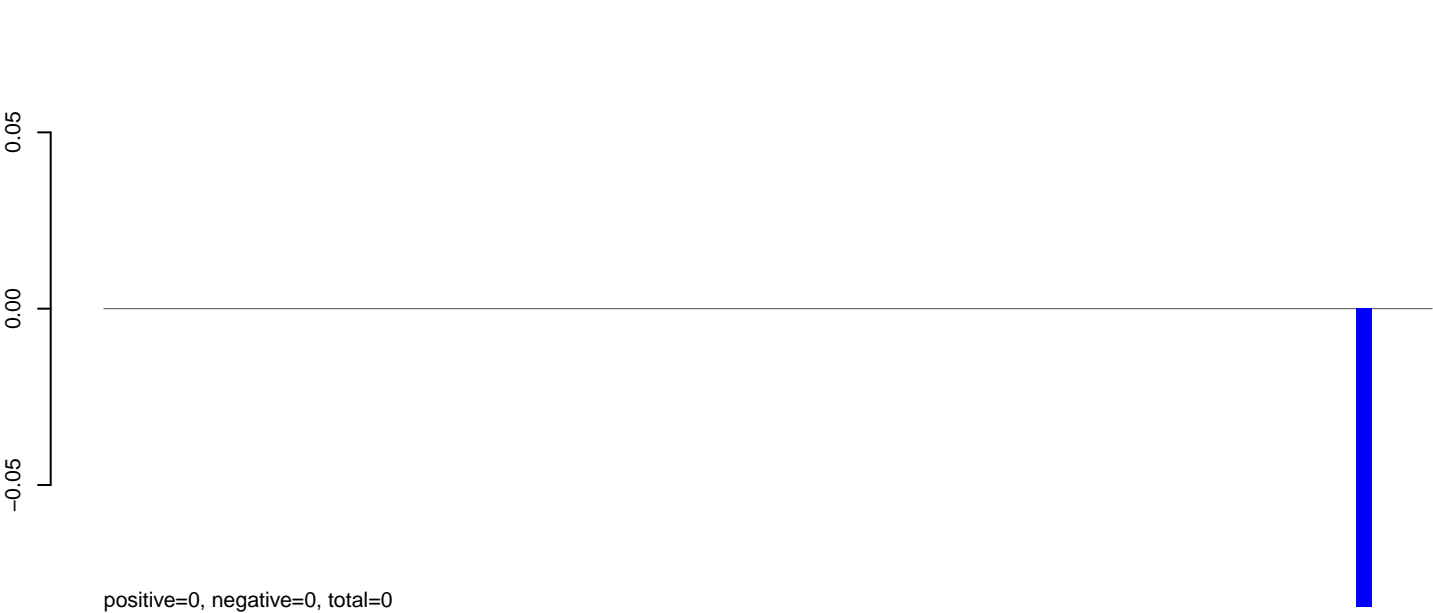
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



Window size=50, length=4435, TE@GYPSY28-LTR-I_AG 1-4435

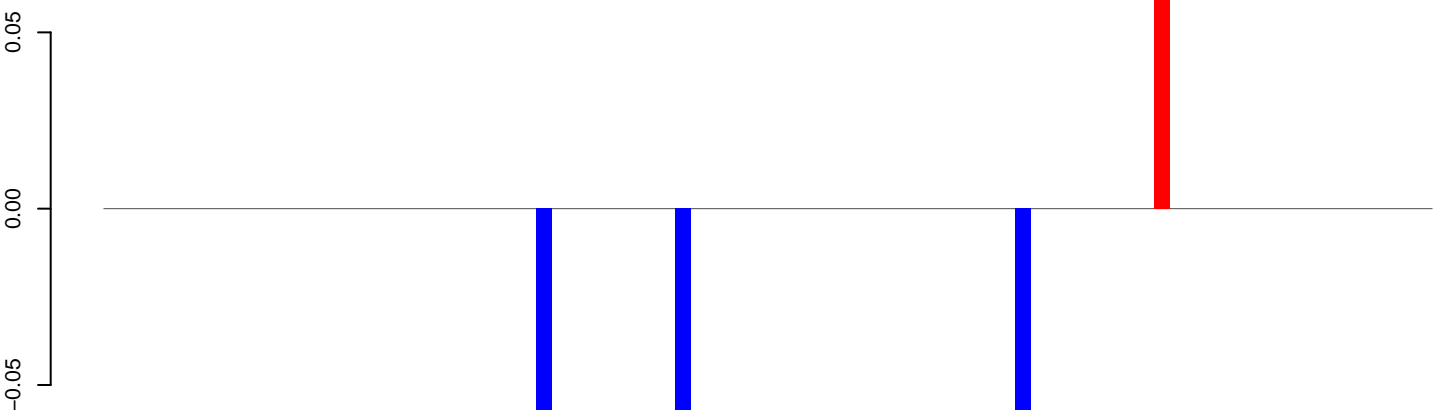
0 1000 2000 3000 4000

AnGam_Mos55_cells.18_23.rep



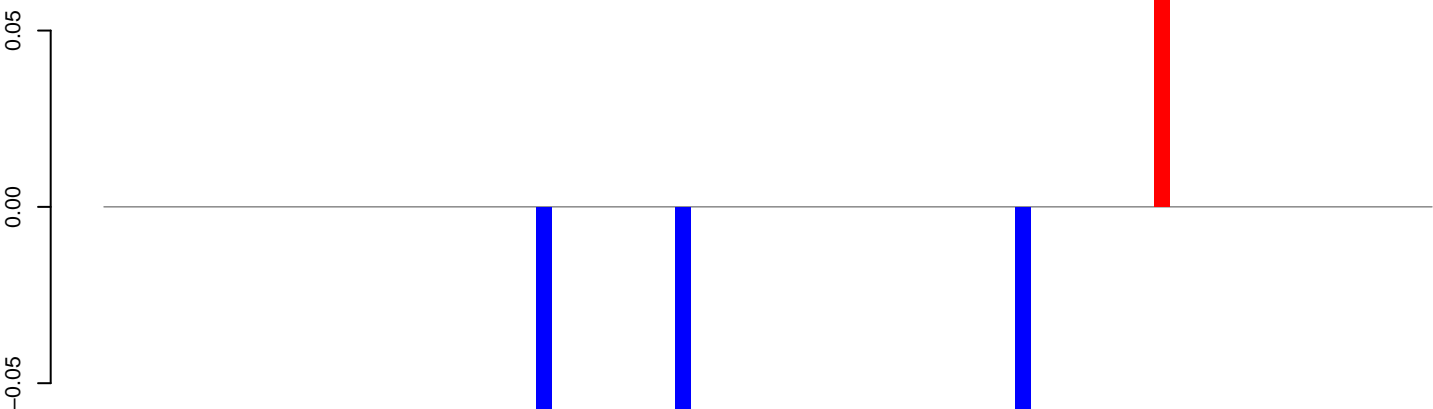
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

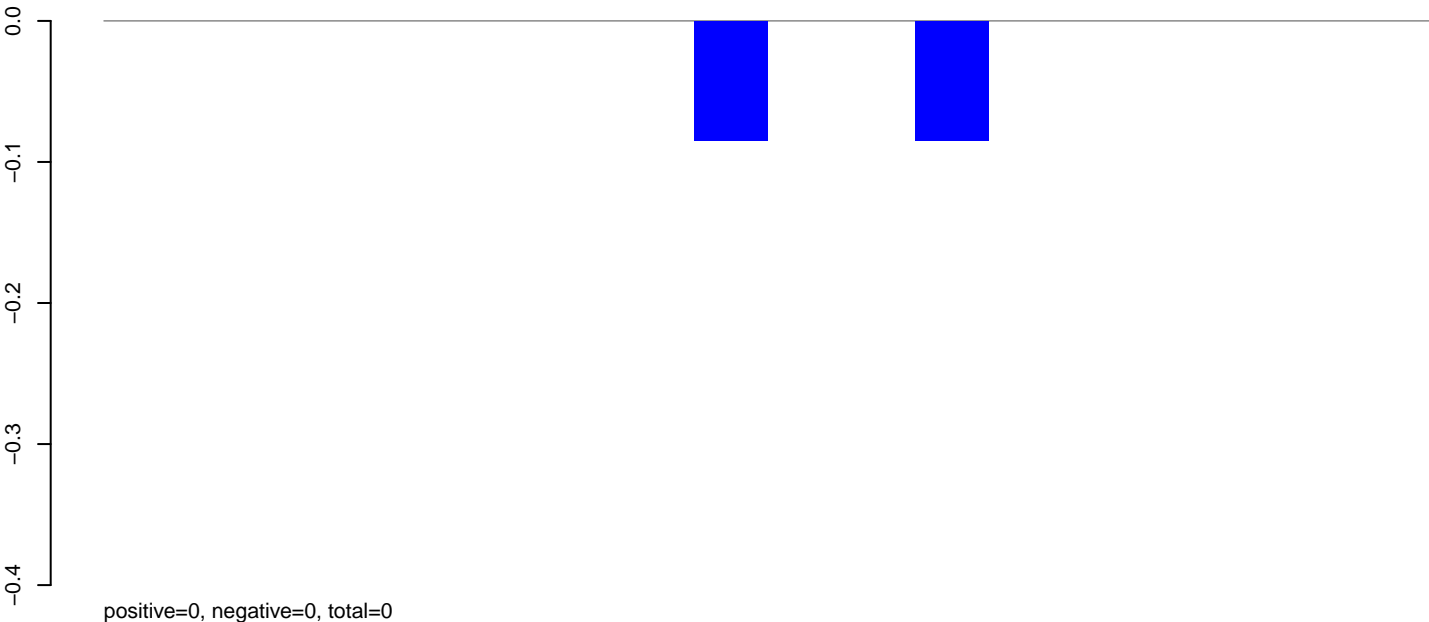


positive=0, negative=0, total=0

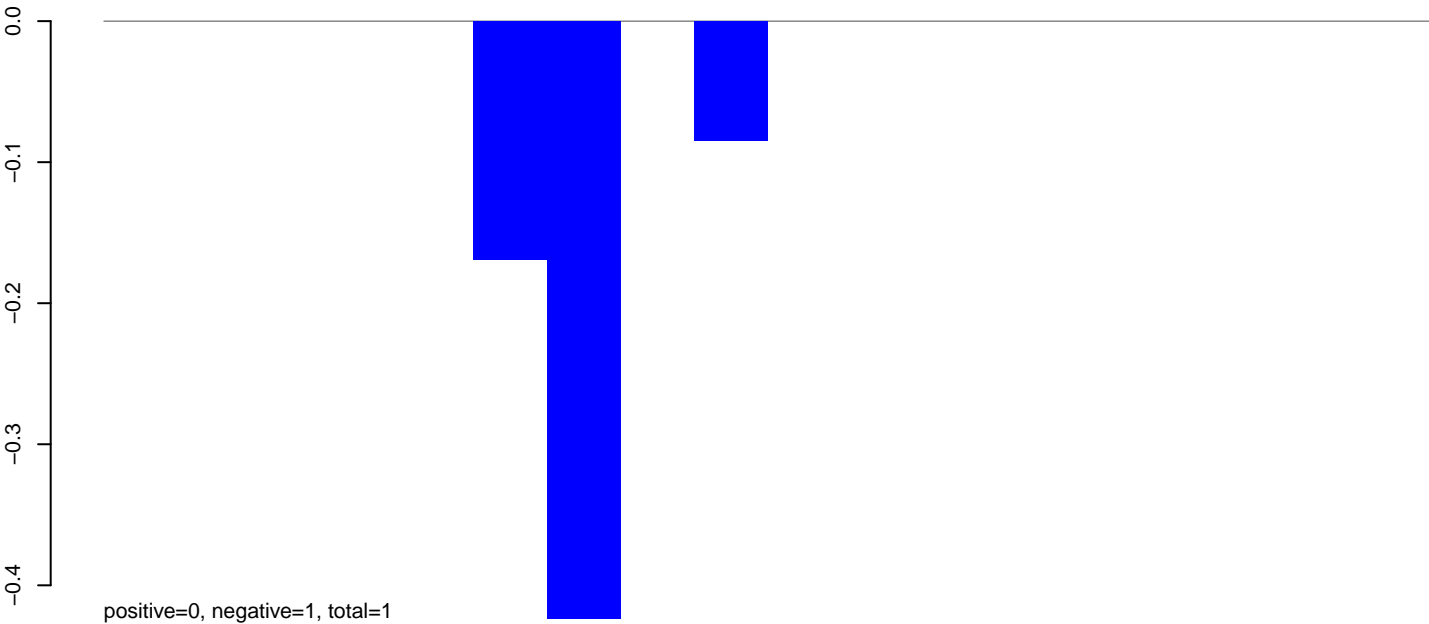
Window size=50, length=4312, TE@GYPSY19-LTR-L_AG:1-4312

0 1000 2000 3000 4000

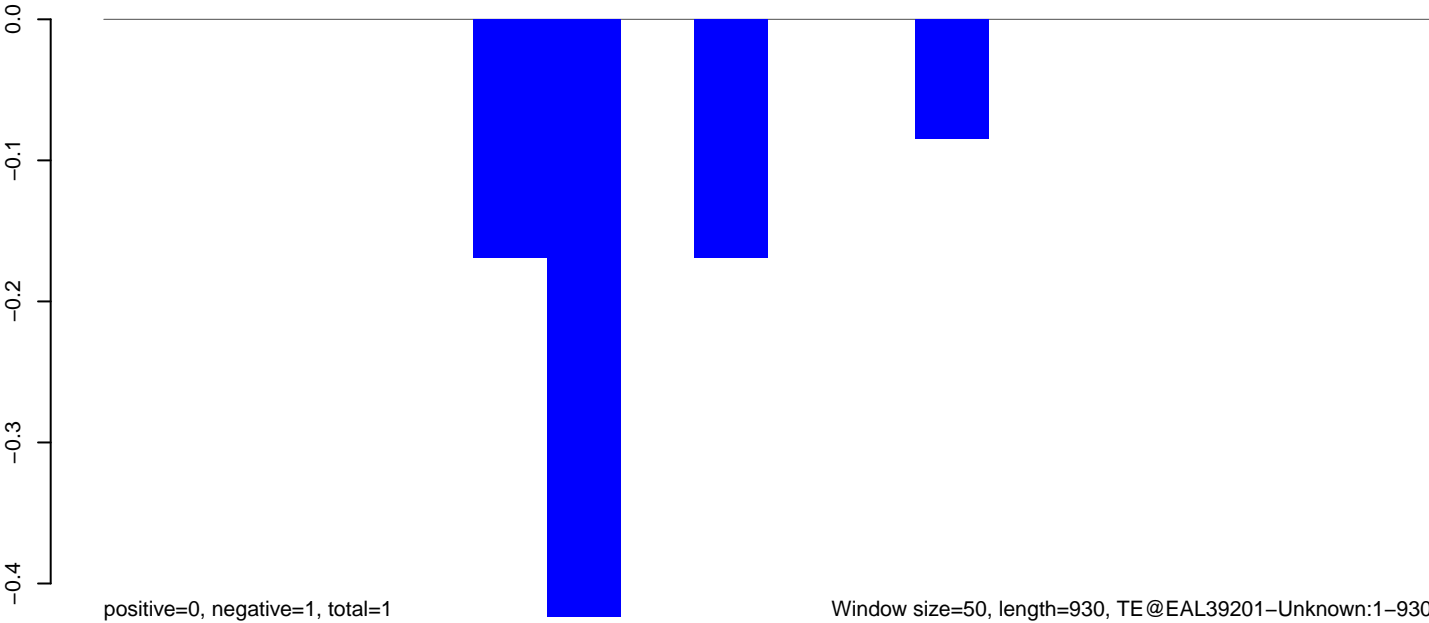
AnGam_Mos55_cells.18_23.rep



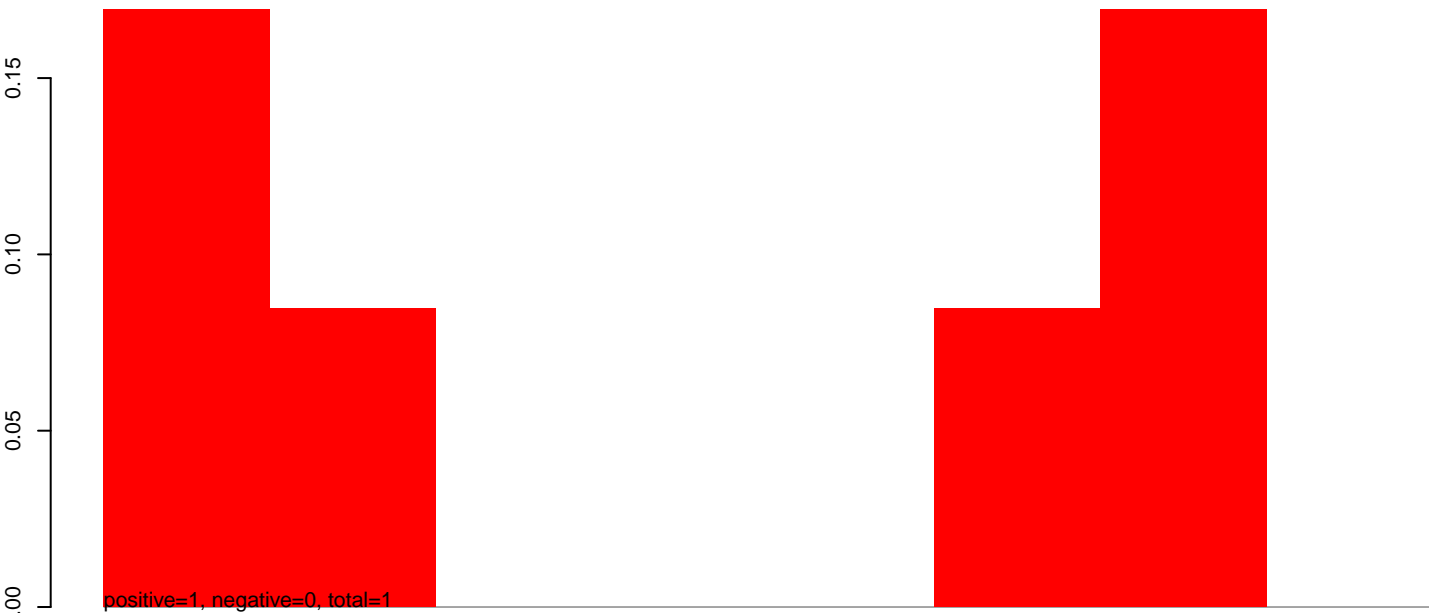
AnGam_Mos55_cells.24_35.rep



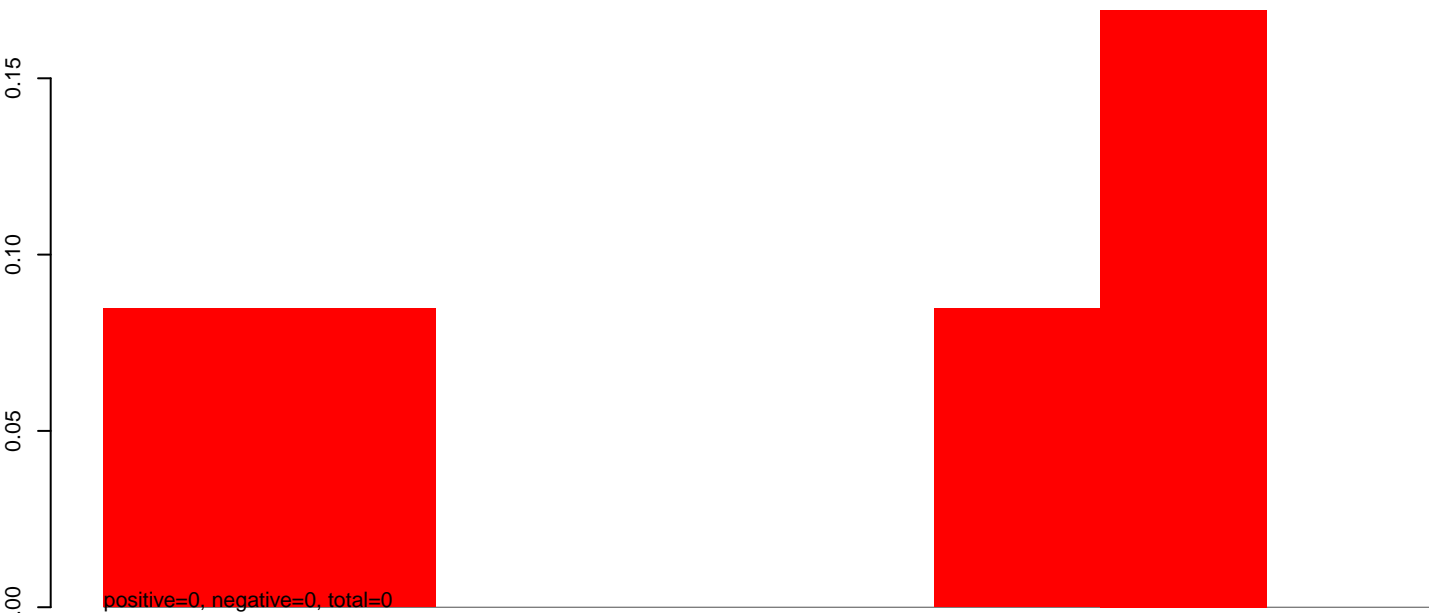
AnGam_Mos55_cells.rep



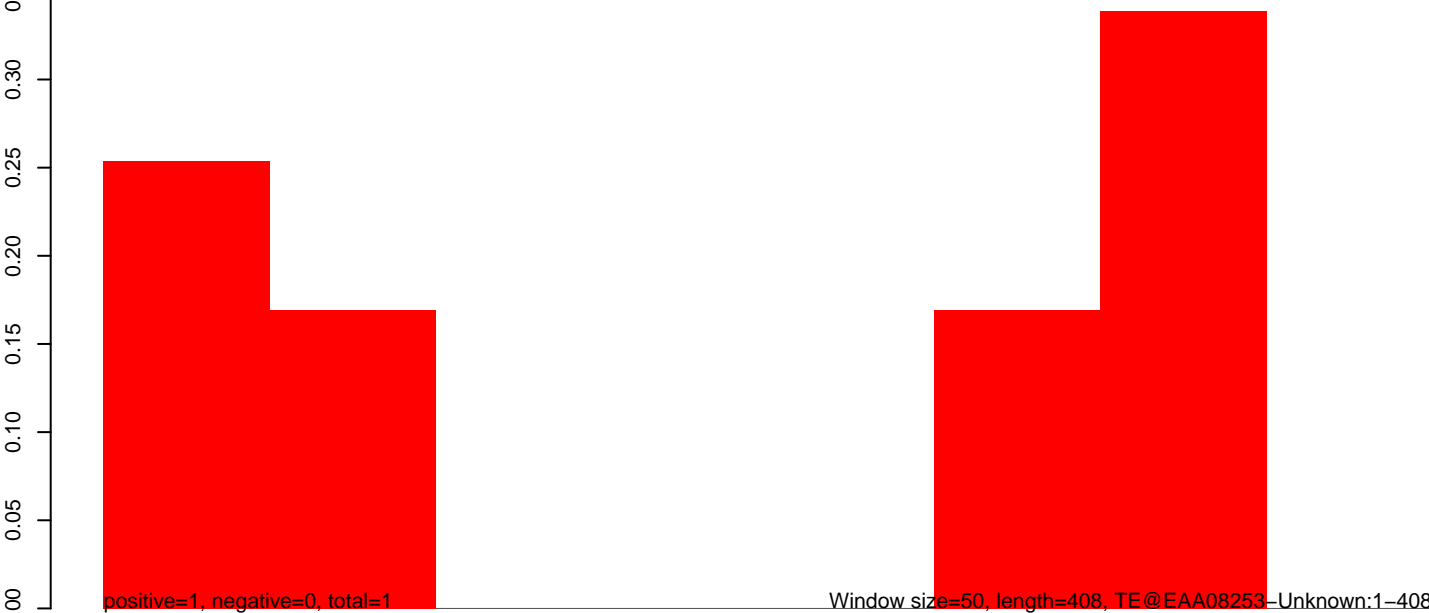
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



100

200

300

400

AnGam_Mos55_cells.18_23.rep

0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

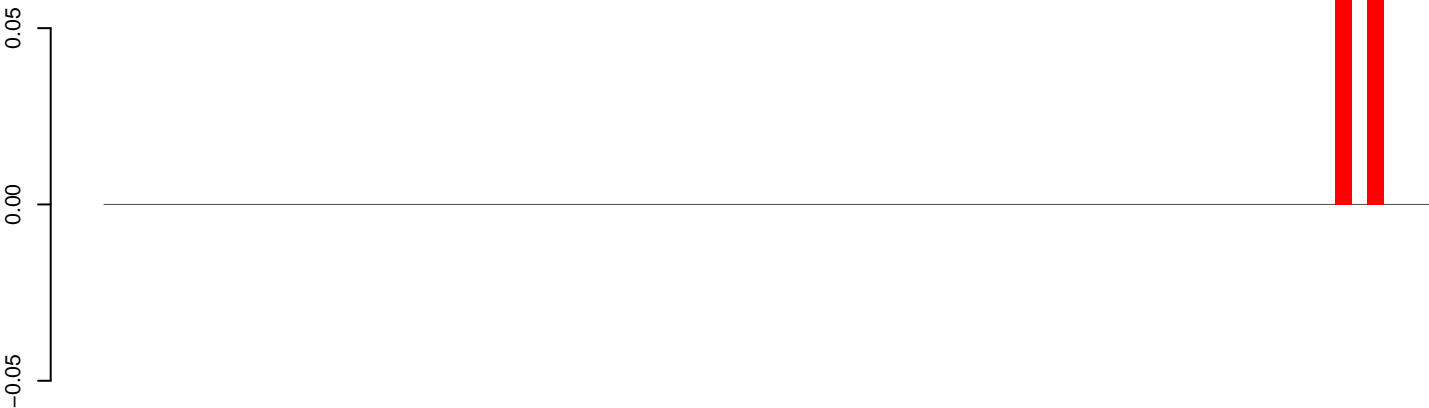
0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

Window size=50, length=1899, TE@CR1-8_AG:1-1899

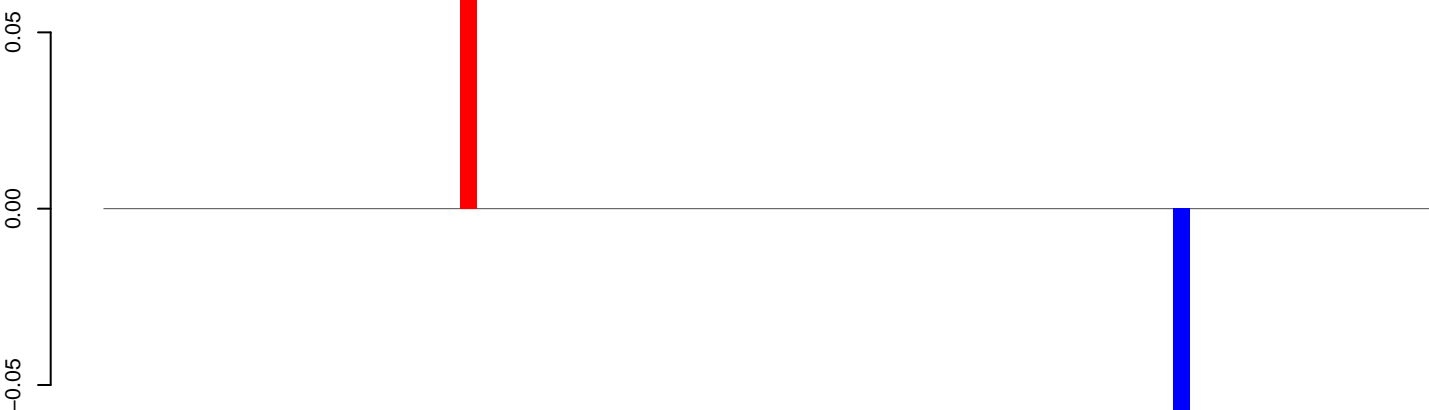
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AnGam_Mos55_cells.18_23.rep



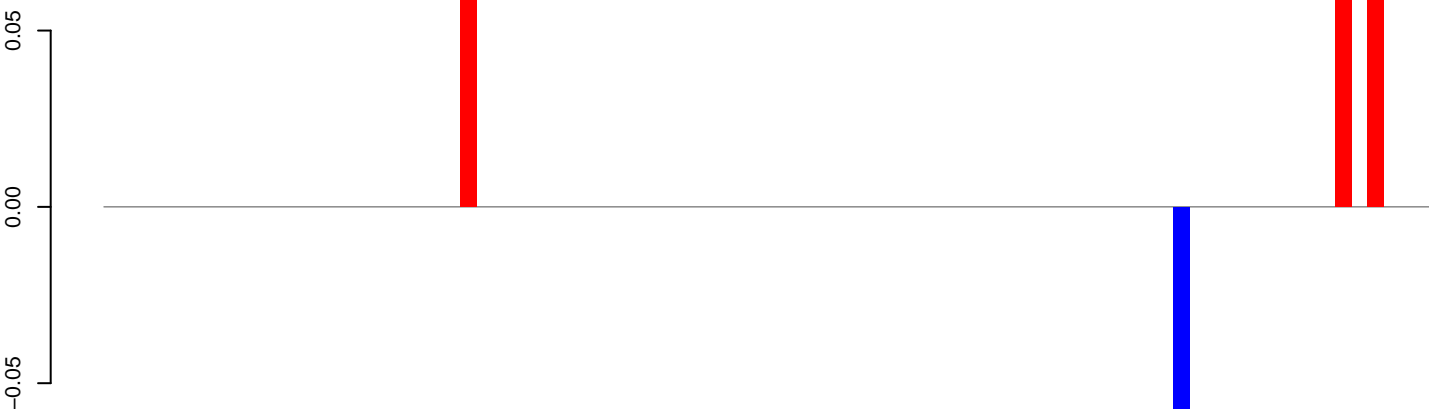
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

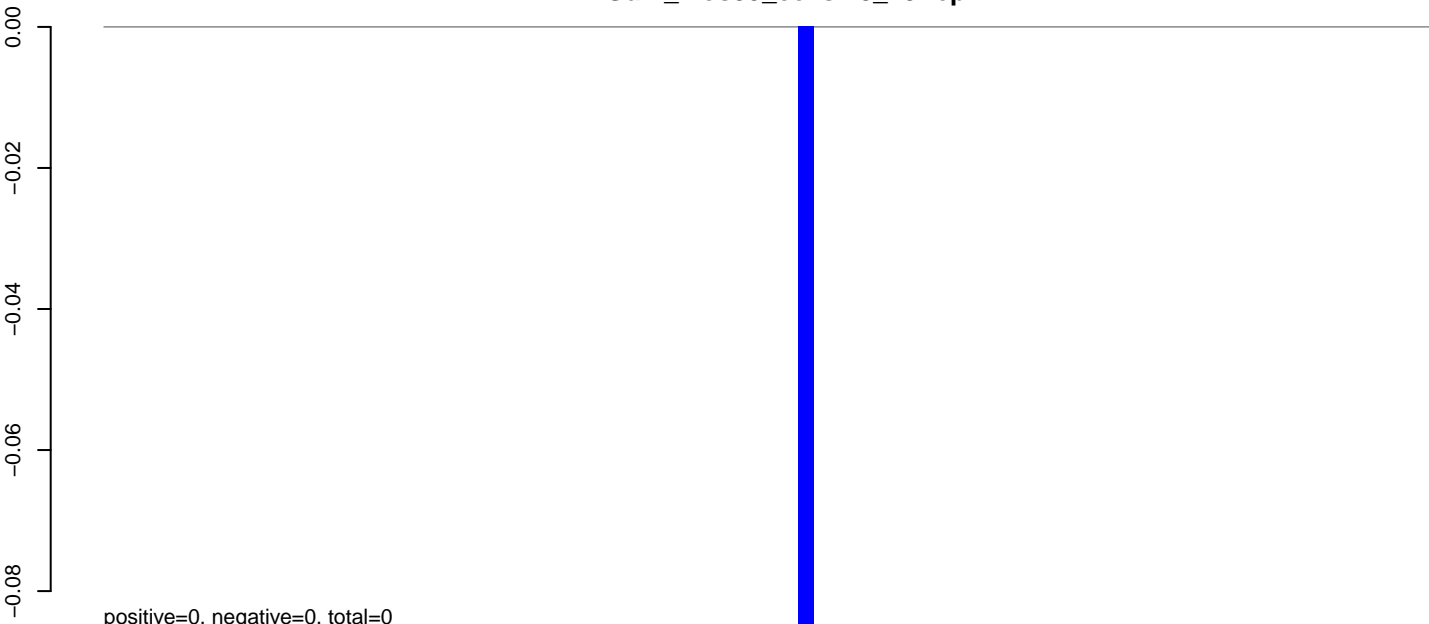


positive=0, negative=0, total=0

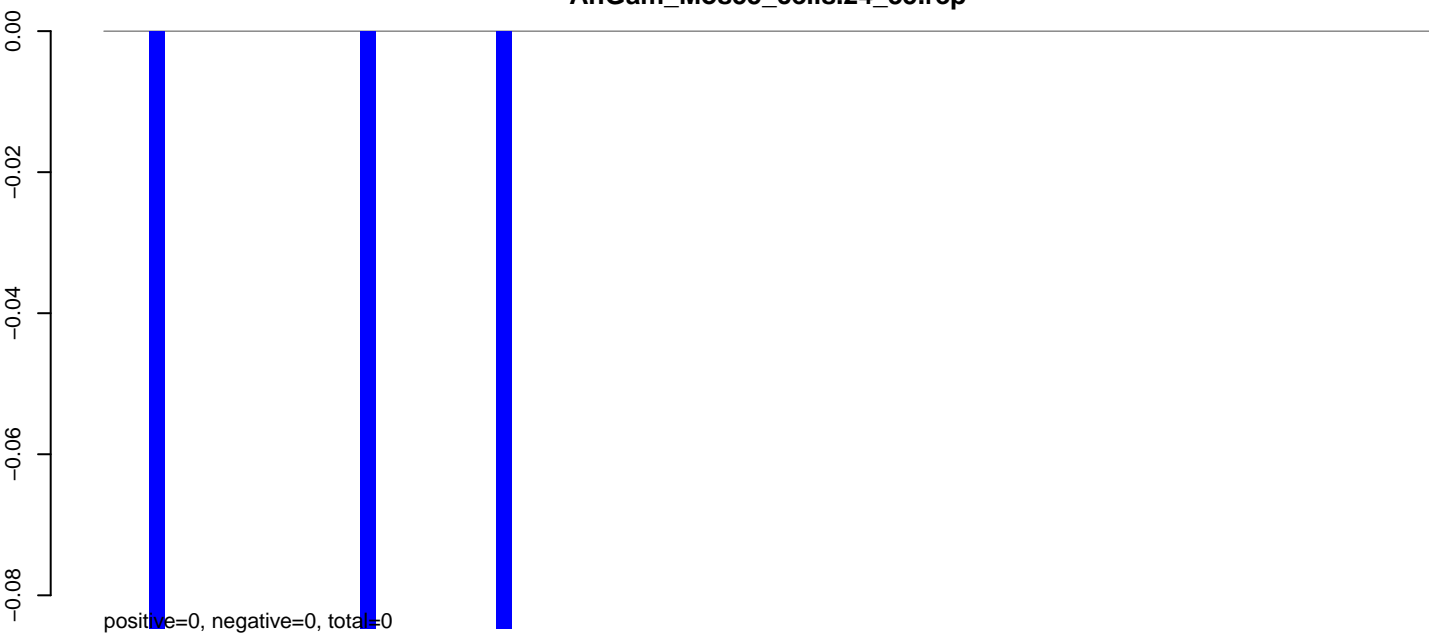
Window size=50, length=4129, TE@COPIA5-LTR-I_AG:1-4129

0 1000 2000 3000 4000

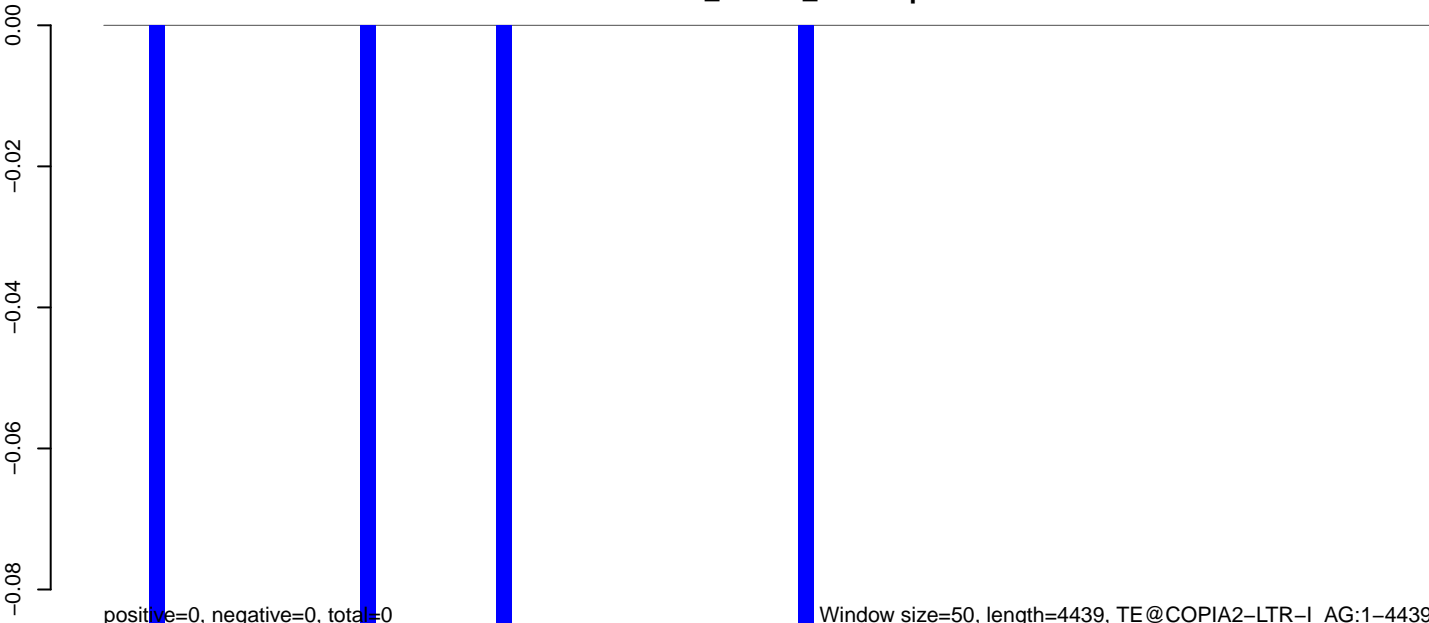
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

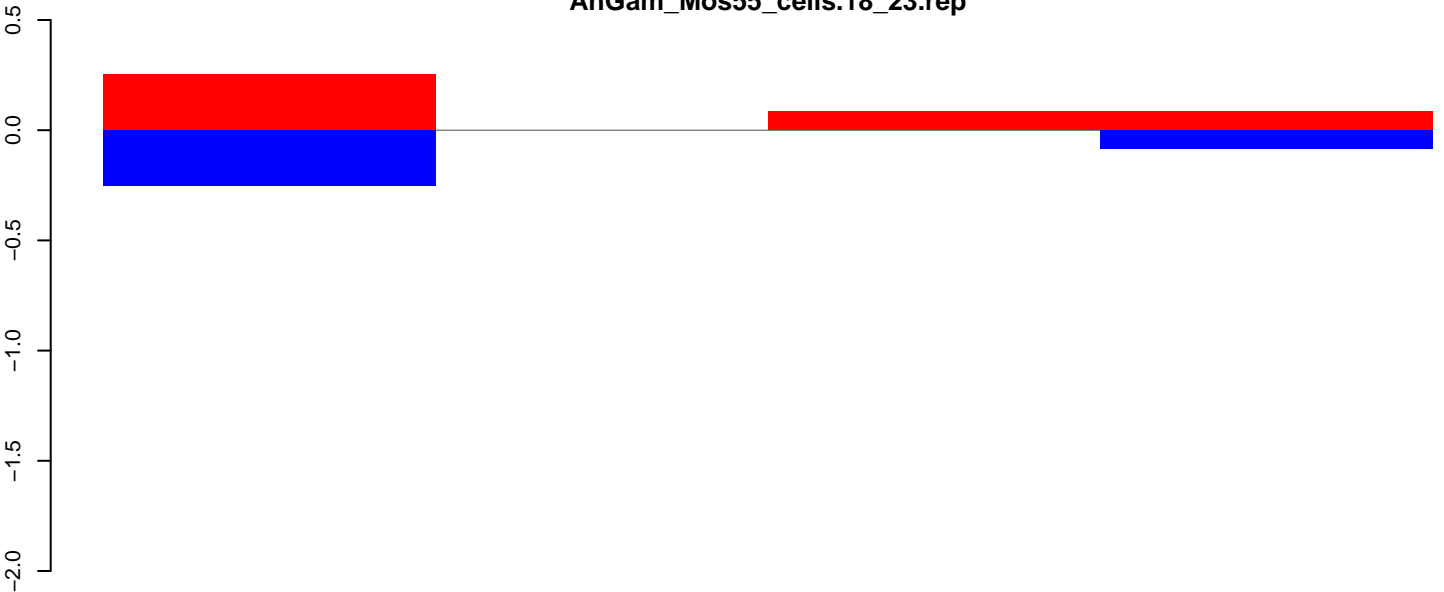


AnGam_Mos55_cells.rep



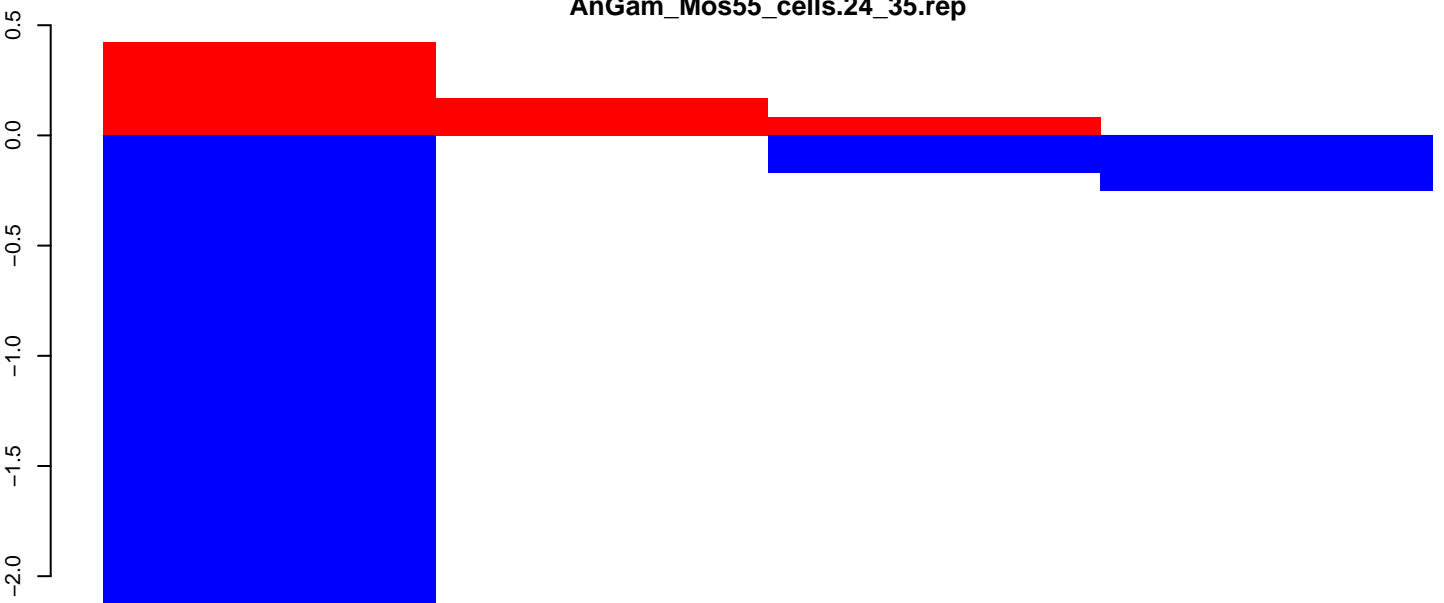
0 1000 2000 3000 4000

AnGam_Mos55_cells.18_23.rep



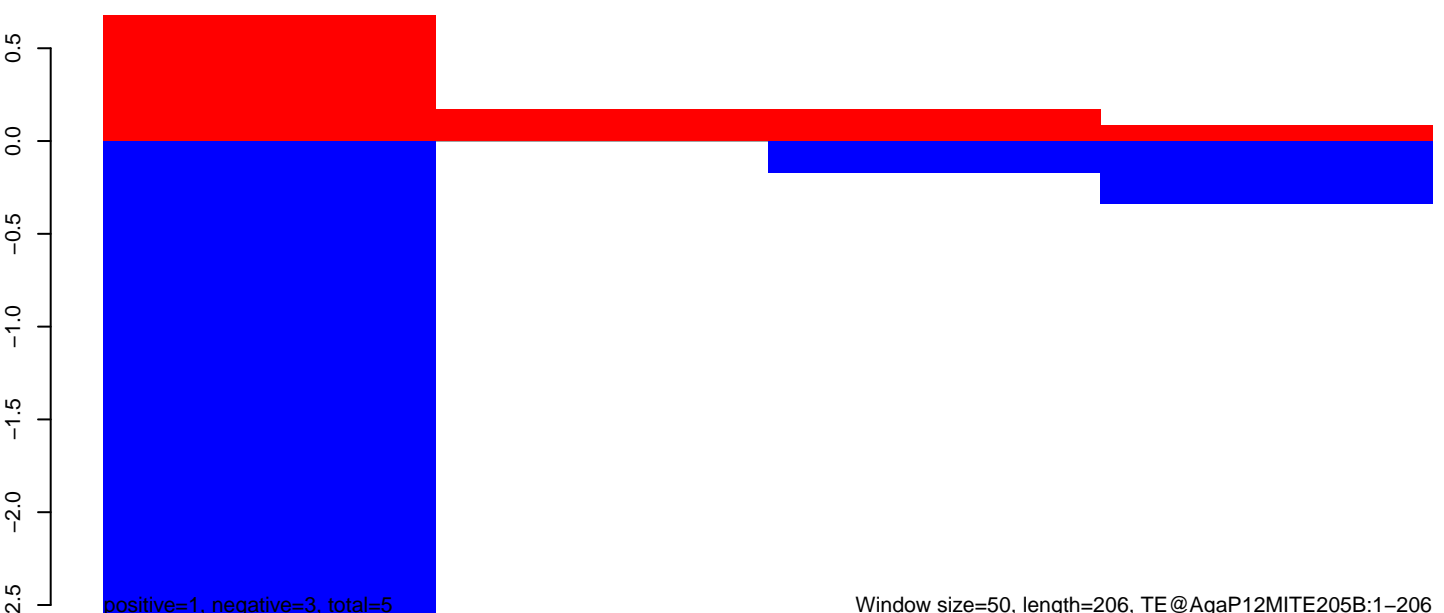
positive=1, negative=0, total=1

AnGam_Mos55_cells.24_35.rep



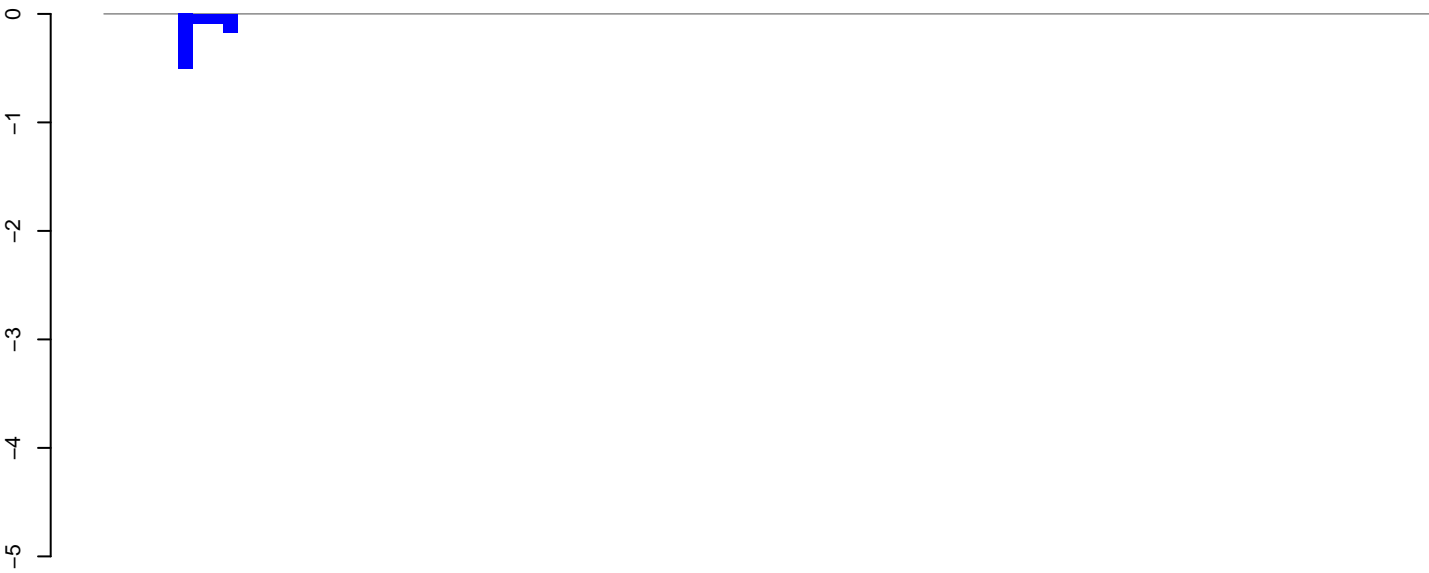
positive=1, negative=3, total=4

AnGam_Mos55_cells.rep



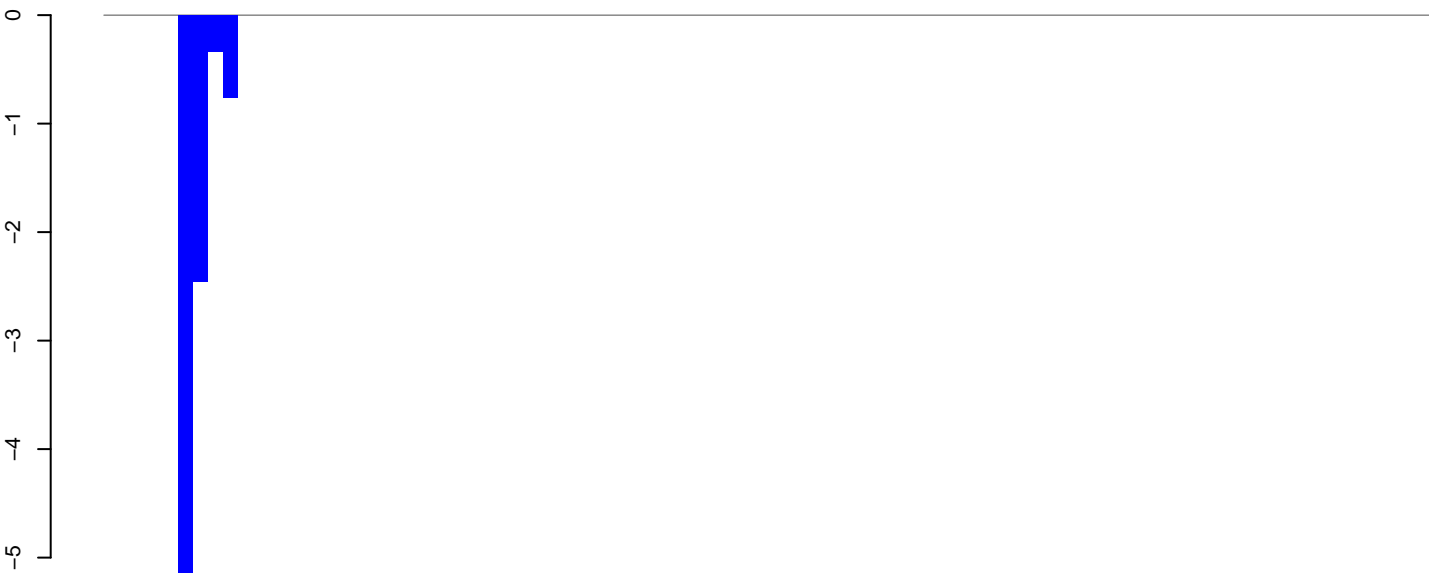
positive=1, negative=3, total=5

AnGam_Mos55_cells.18_23.rep



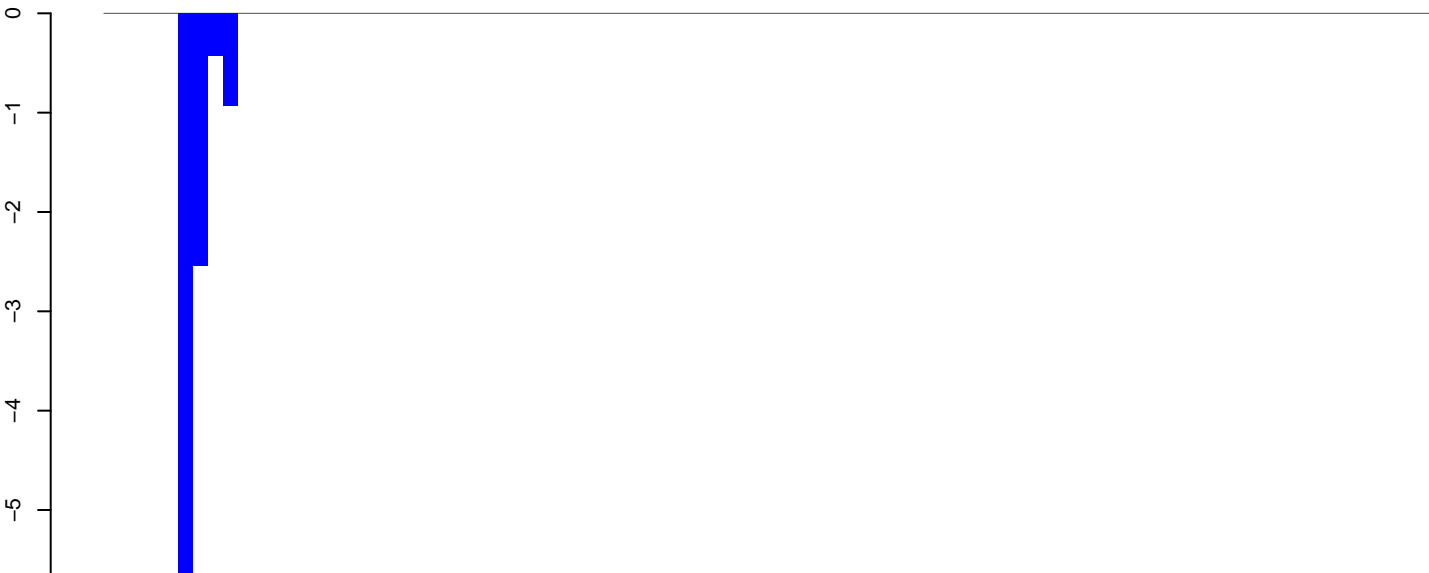
positive=0, negative=1, total=1

AnGam_Mos55_cells.24_35.rep



positive=0, negative=9, total=9

AnGam_Mos55_cells.rep

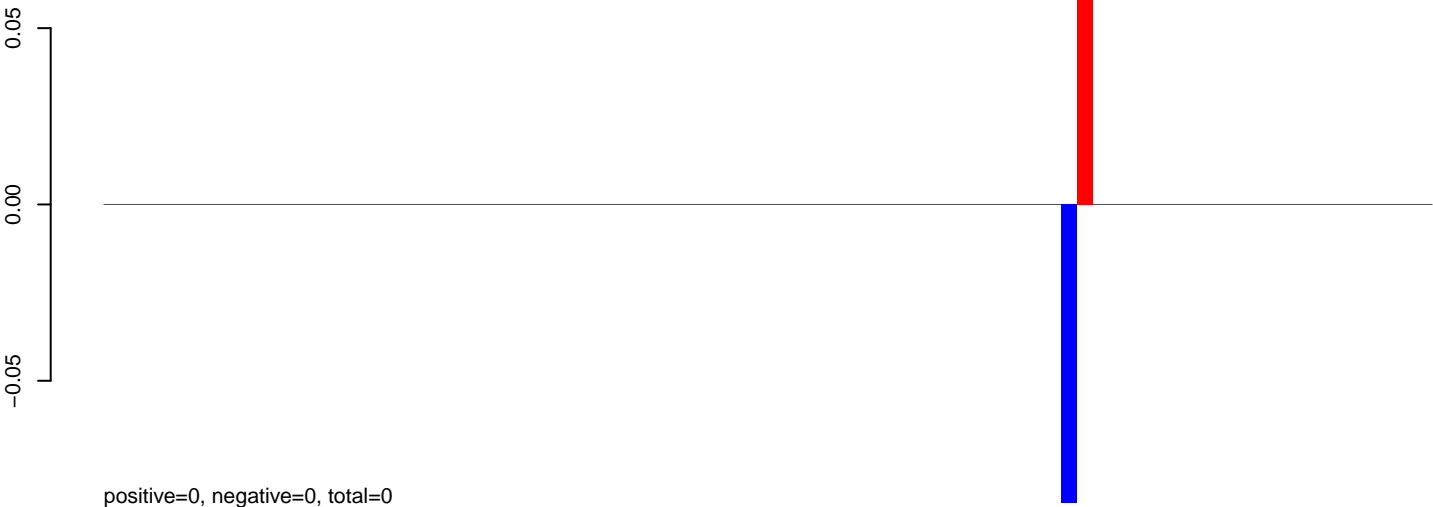


positive=0, negative=10, total=10

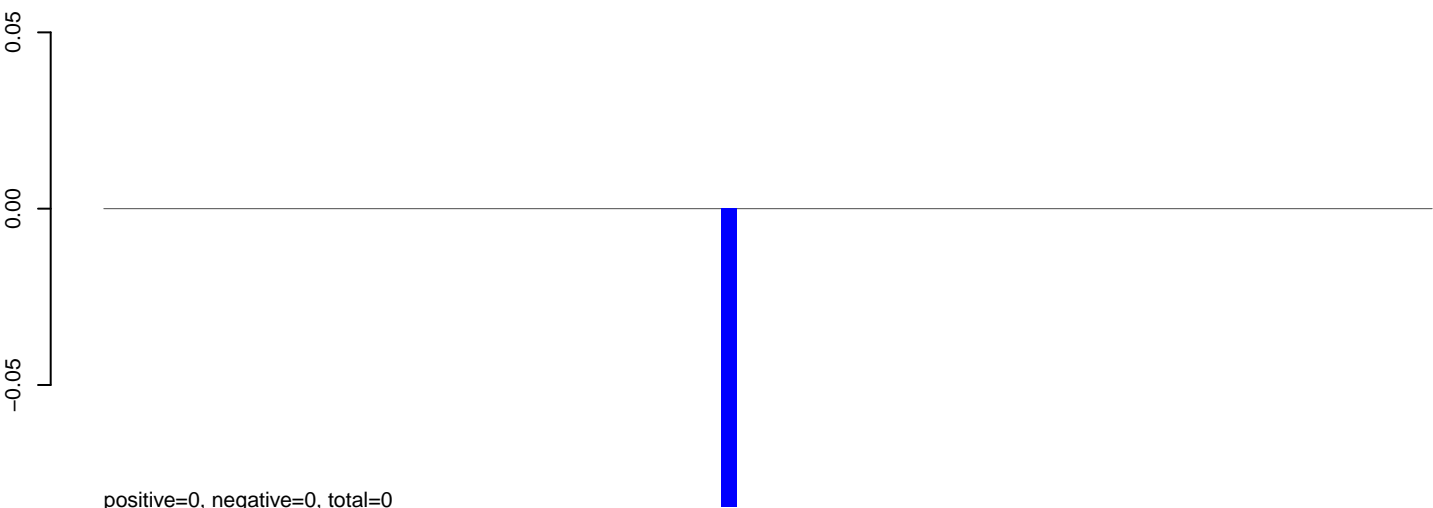
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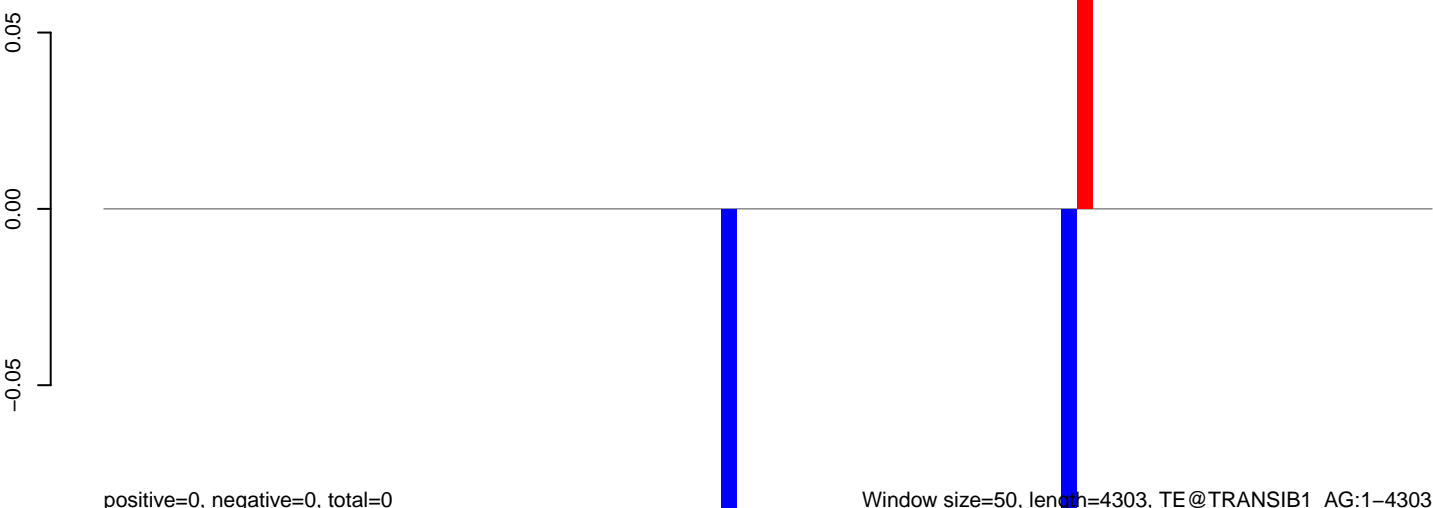
AnGam_Mos55_cells.18_23.rep



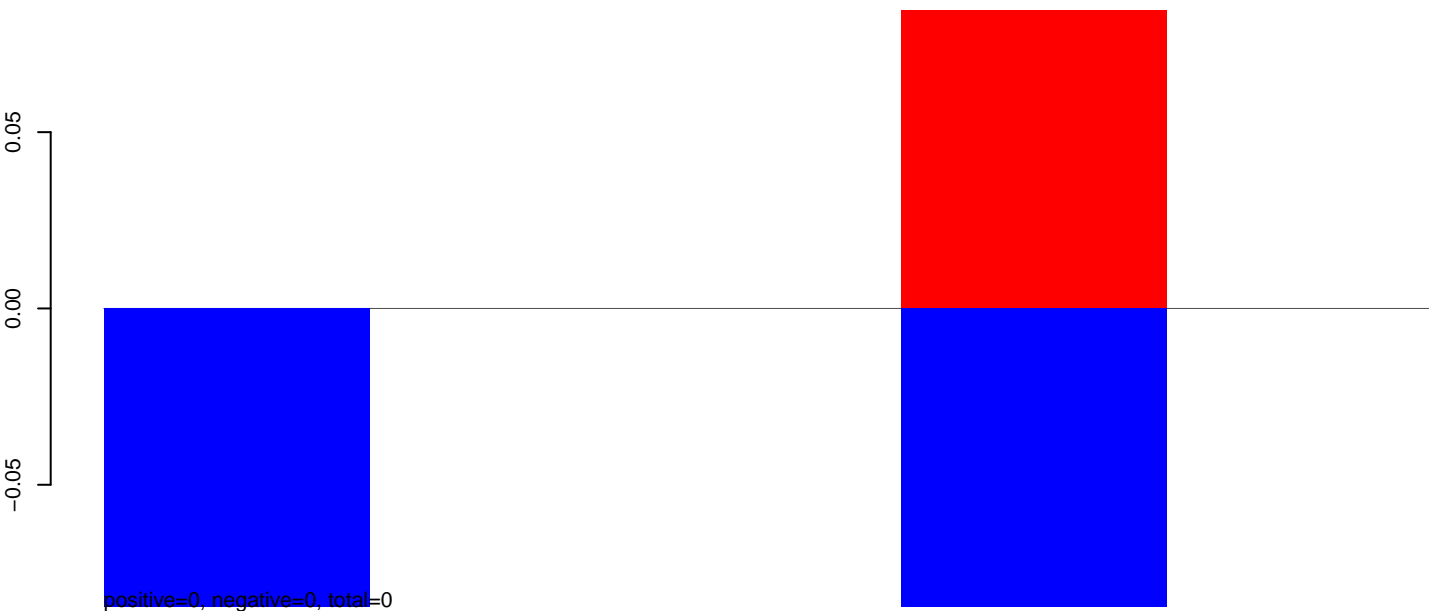
AnGam_Mos55_cells.24_35.rep



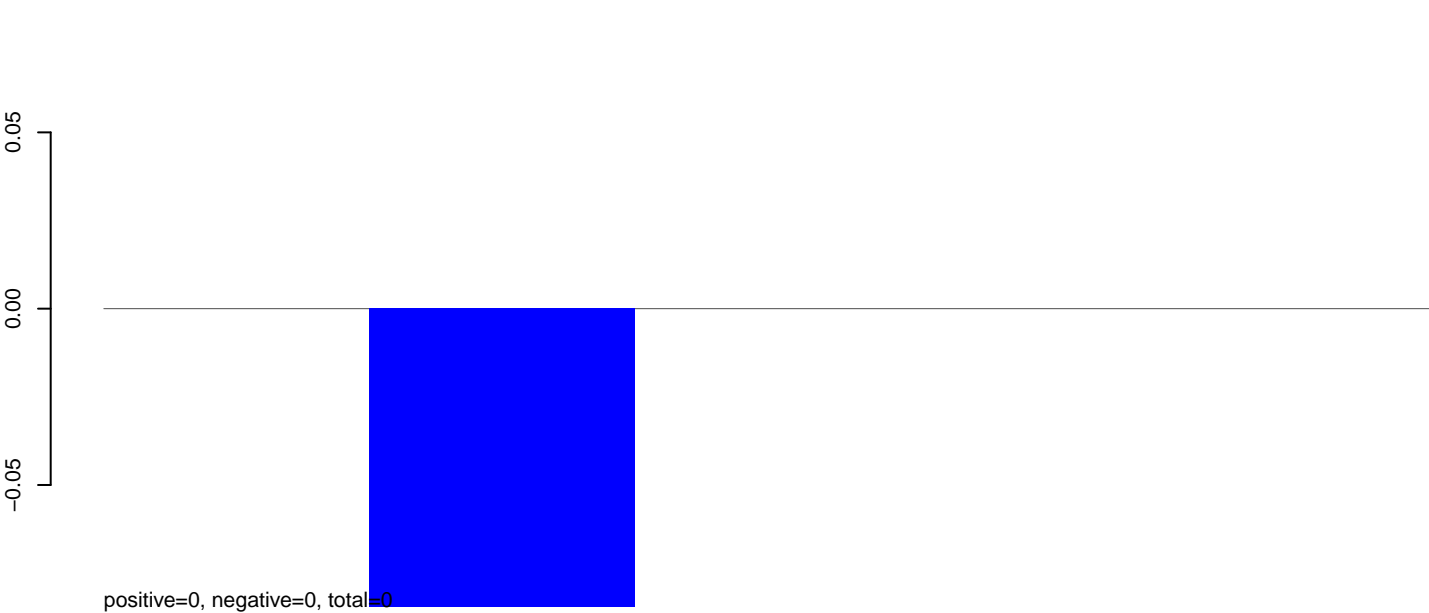
AnGam_Mos55_cells.rep



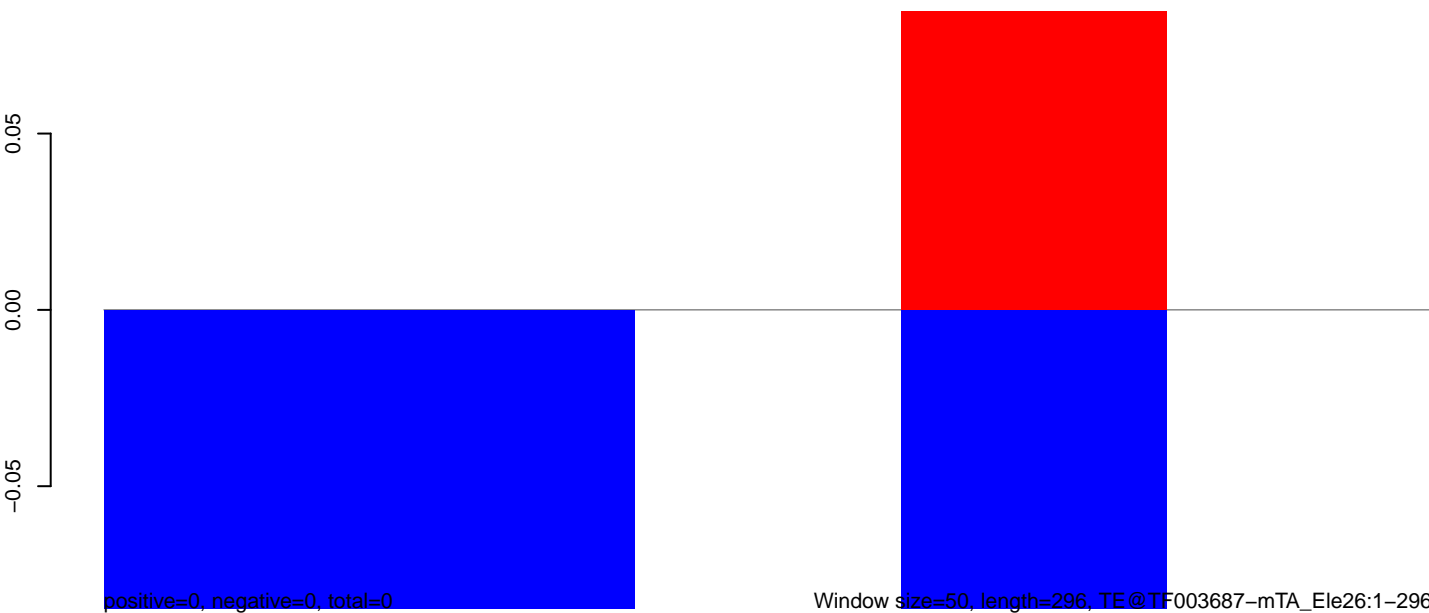
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



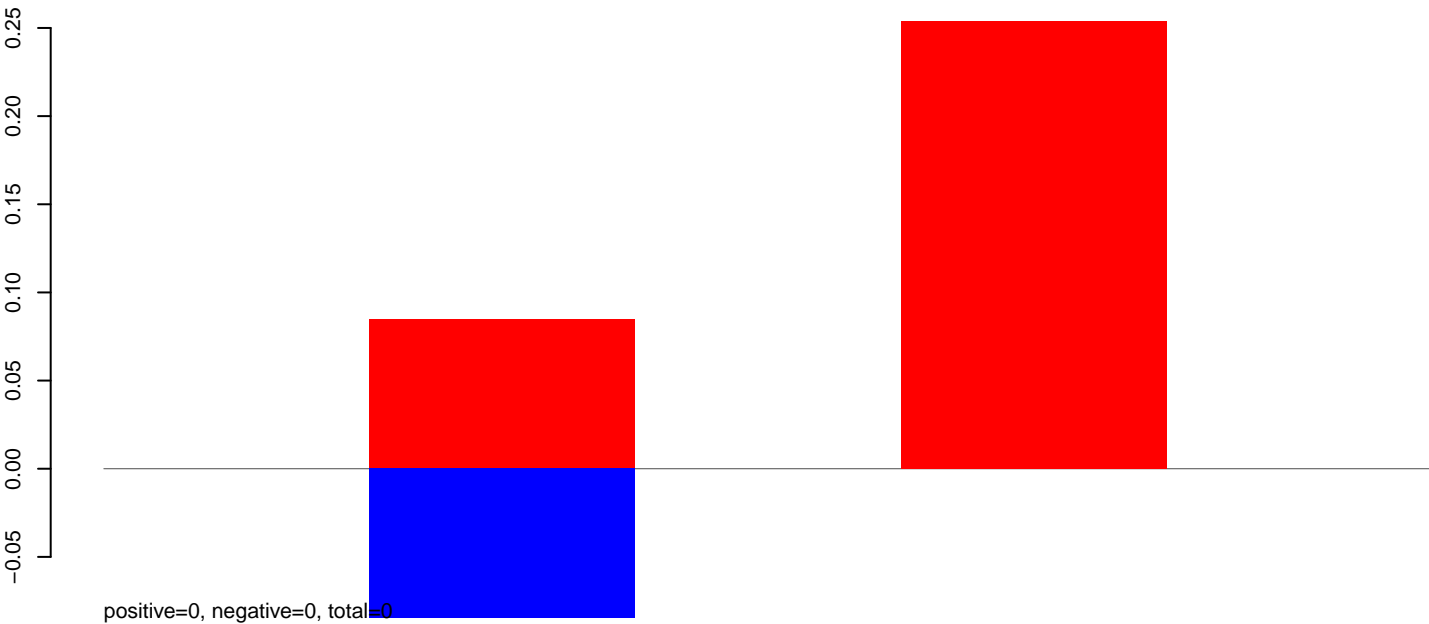
AnGam_Mos55_cells.rep



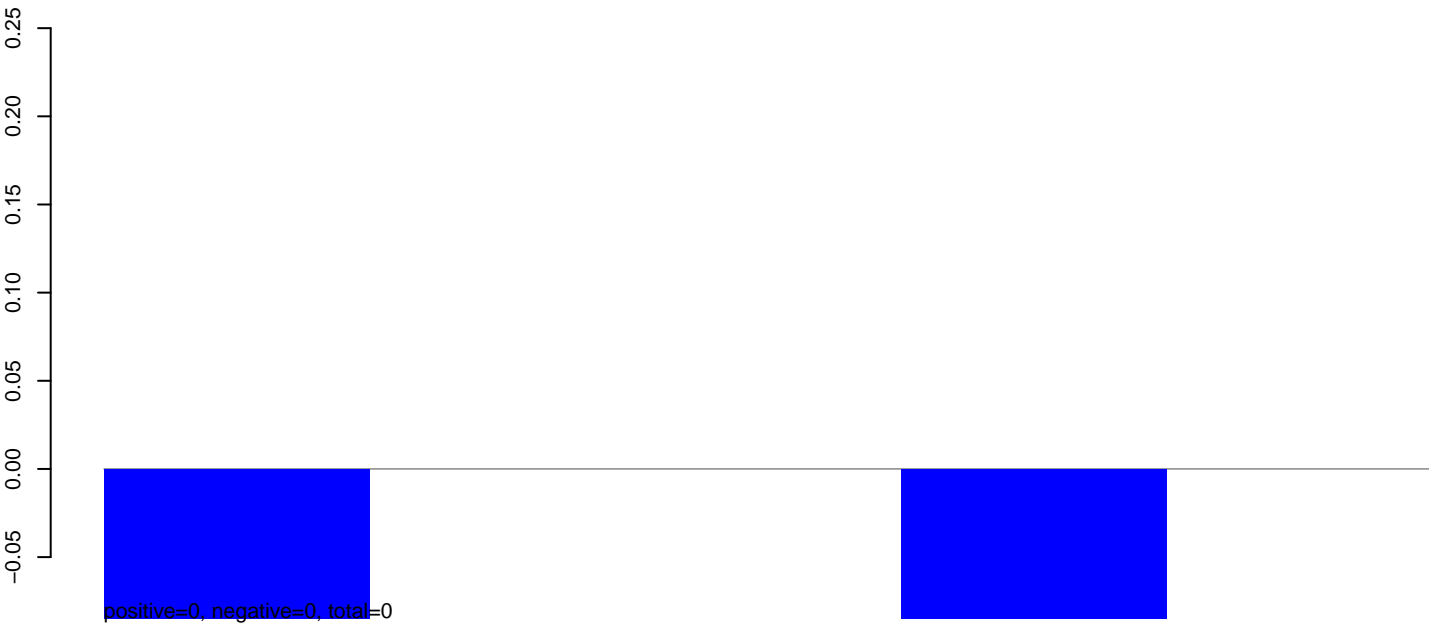
Window size=50, length=296, TE@TF003687-mTA_Ele26:1-296

50 100 150 200 250 300

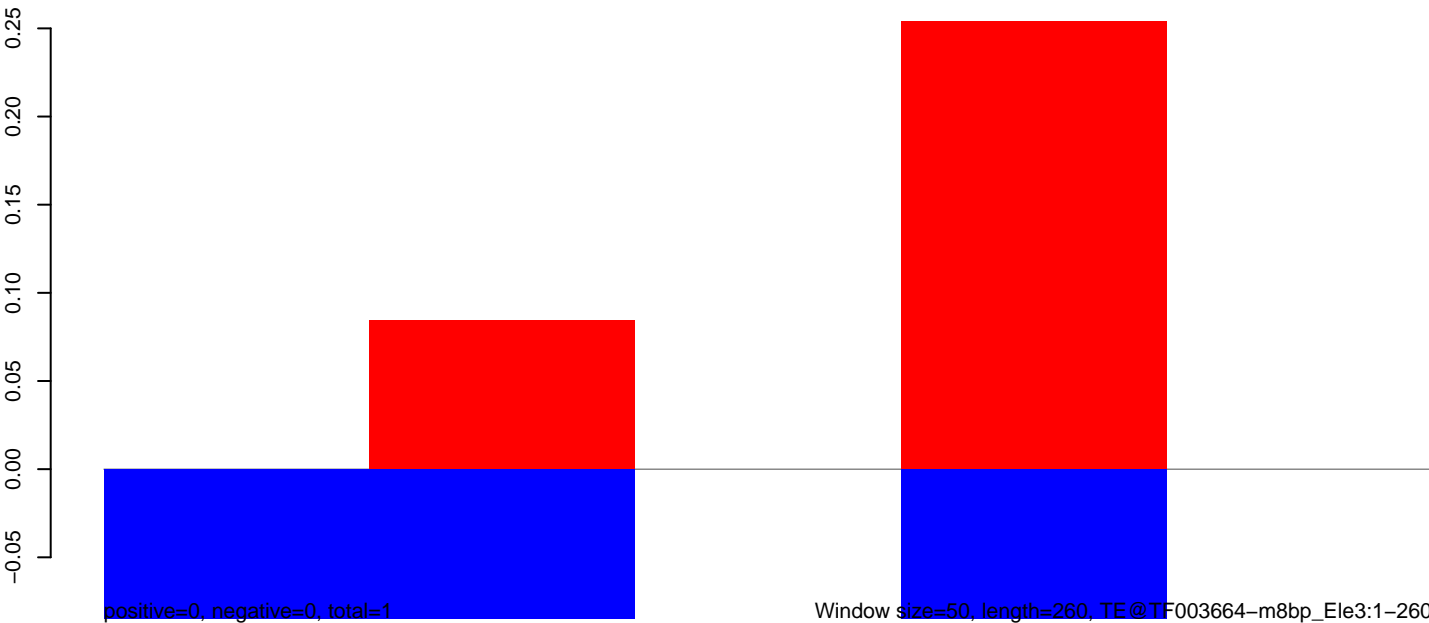
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

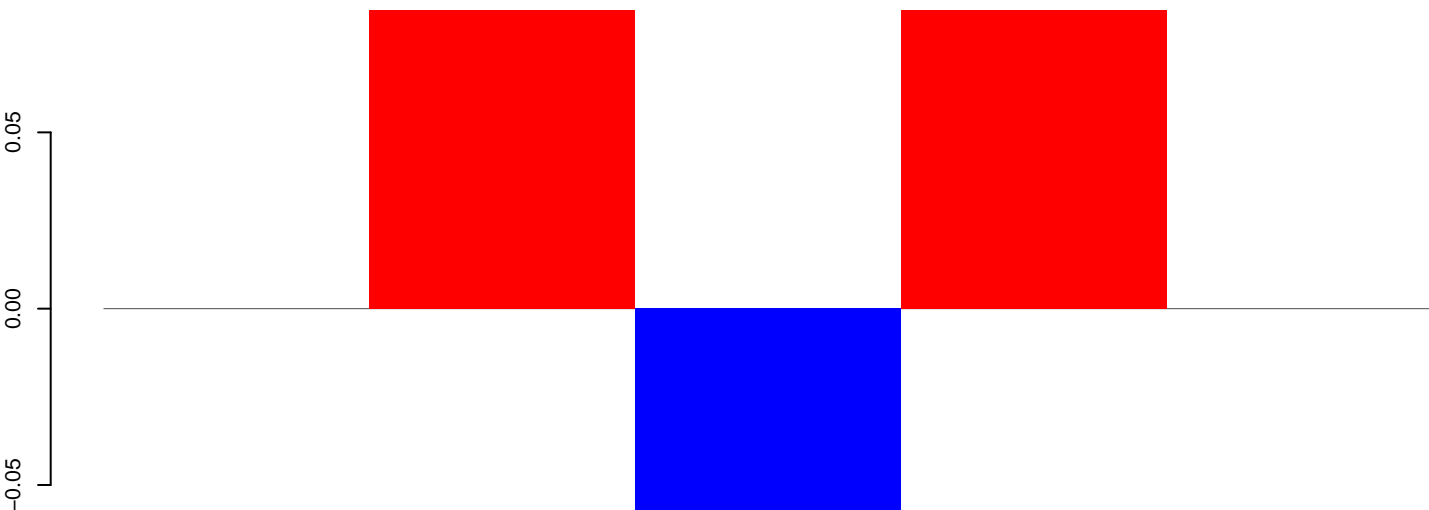


AnGam_Mos55_cells.18_23.rep



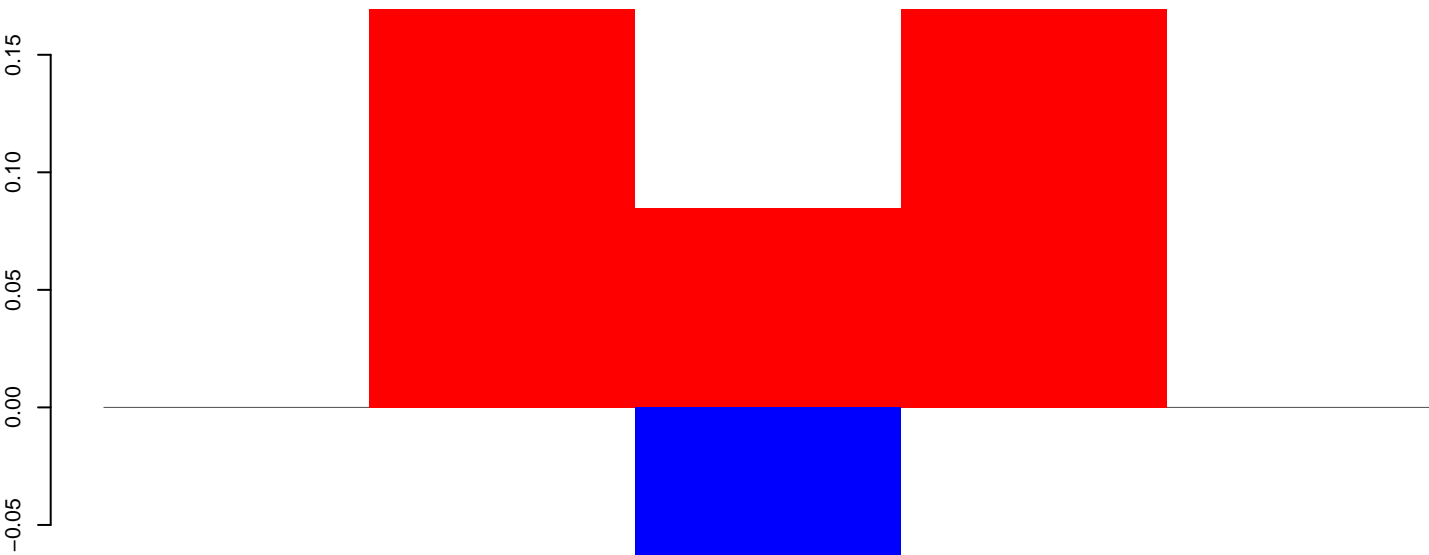
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

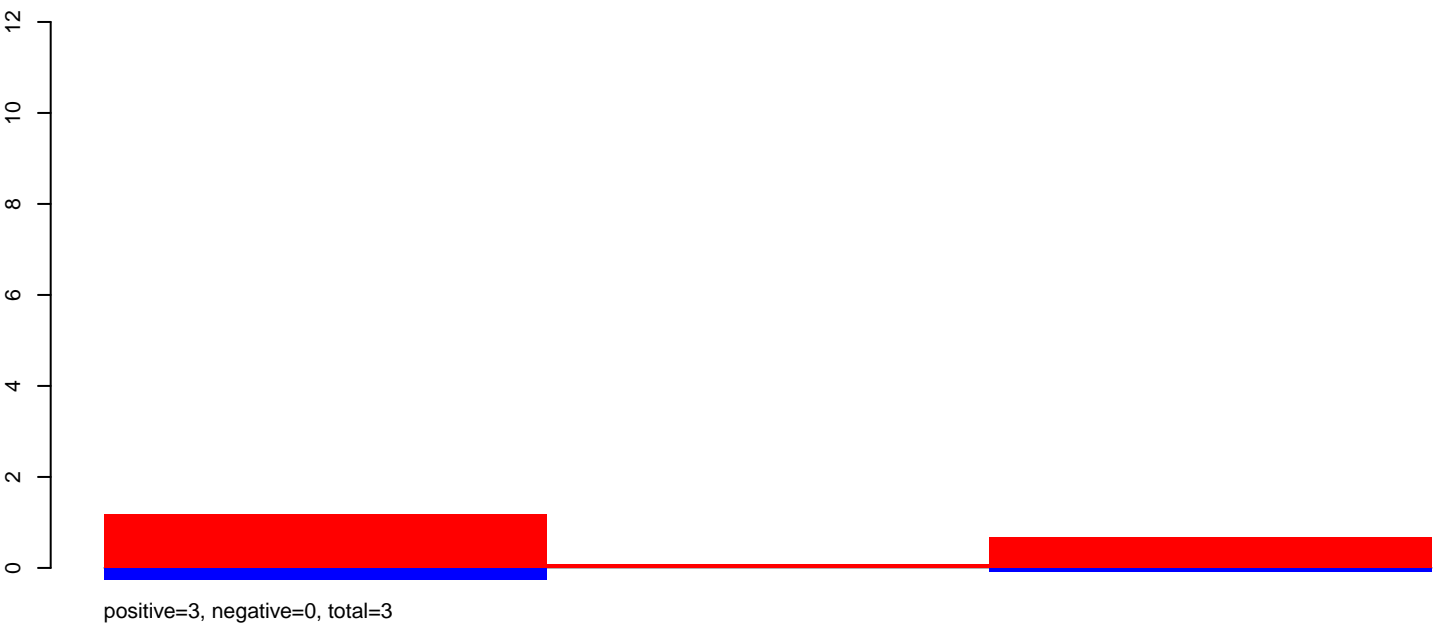


positive=0, negative=0, total=1

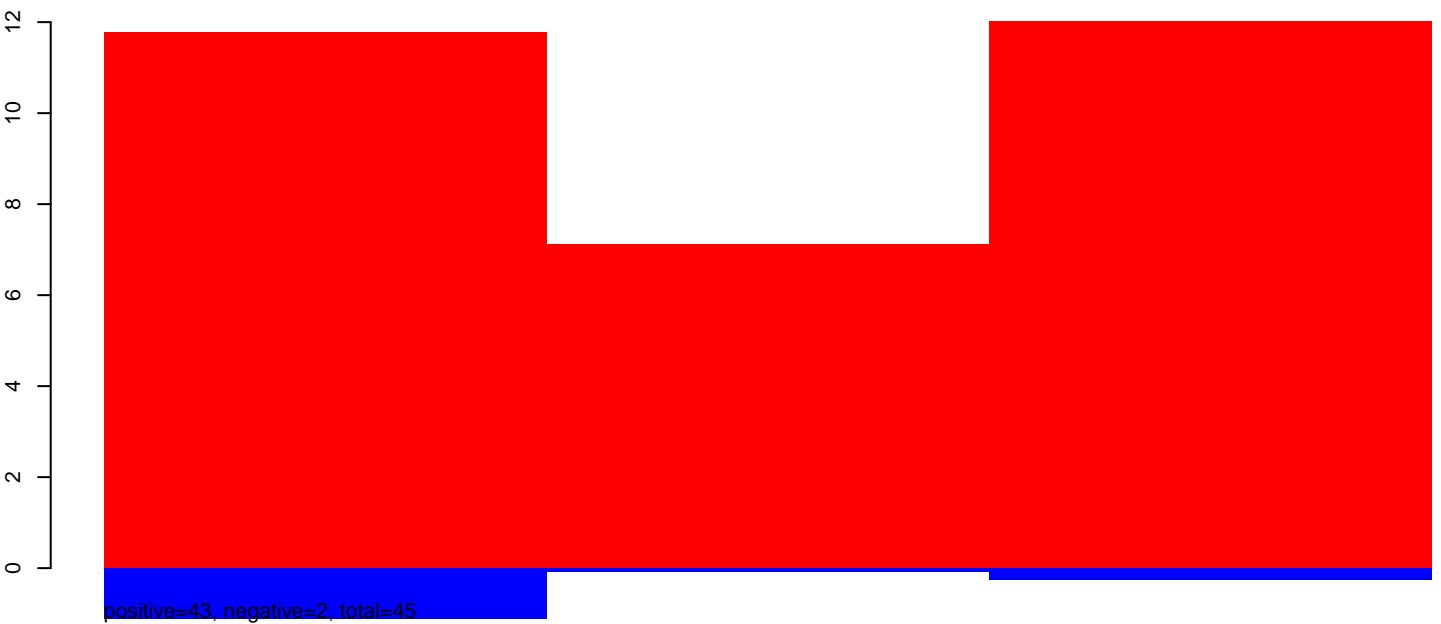
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50 100 150 200 250 300

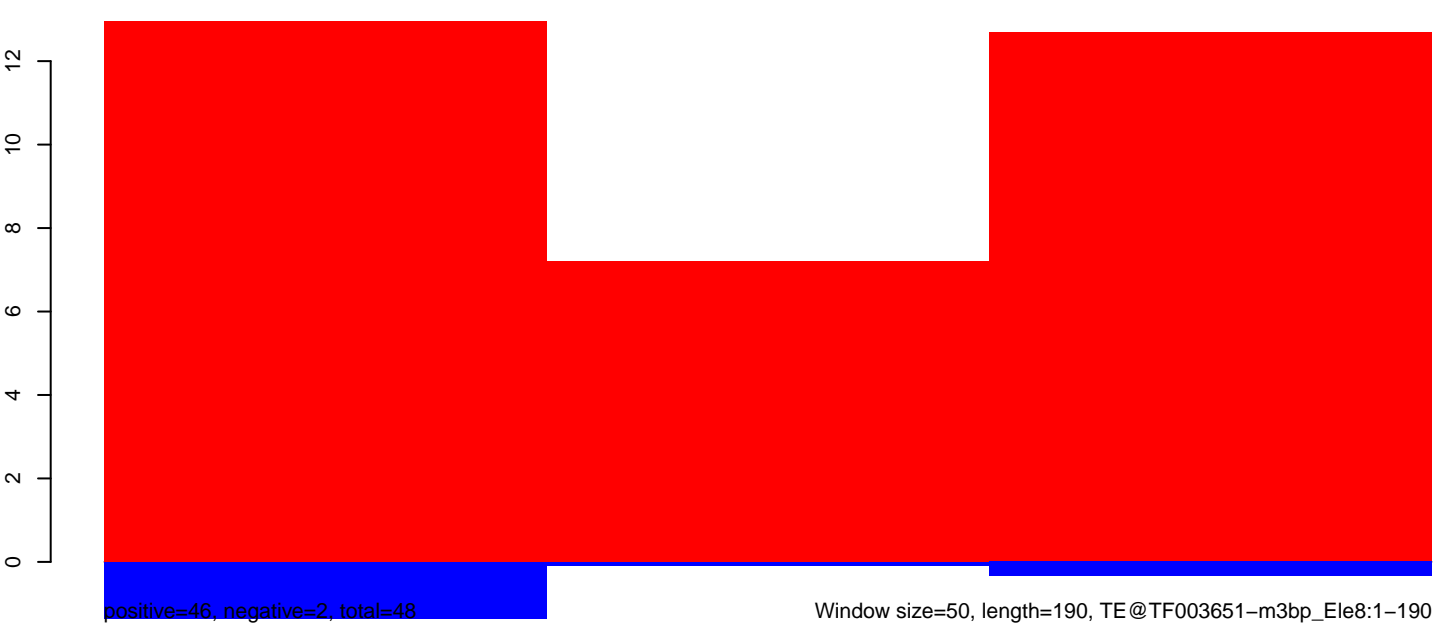
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

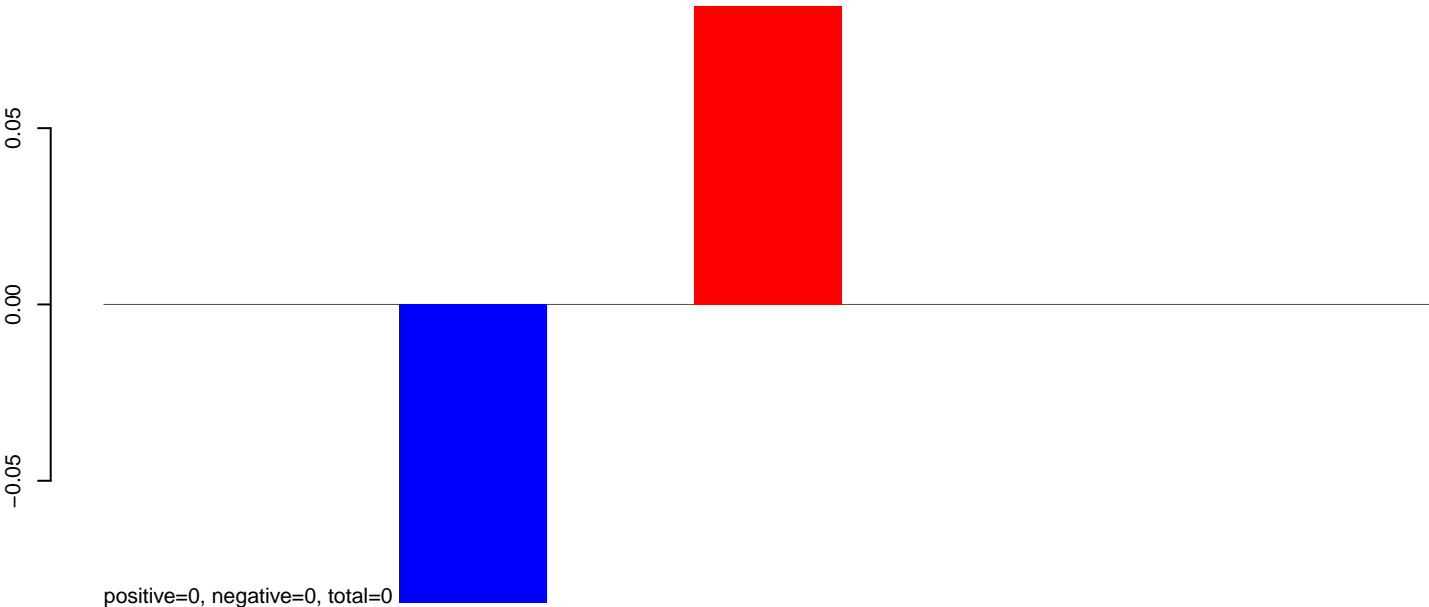


AnGam_Mos55_cells.rep



Window size=50, length=190, TE@TF003651-m3bp_Ele8:1-190

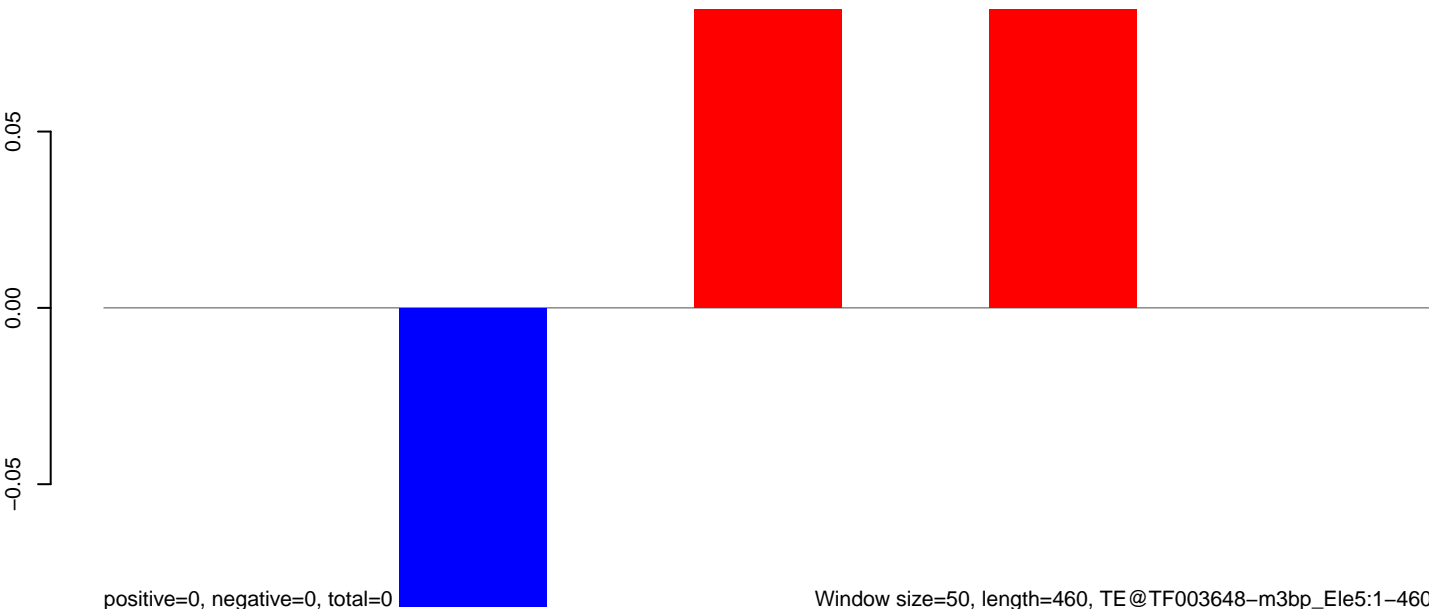
AnGam_Mos55_cells.18_23.rep



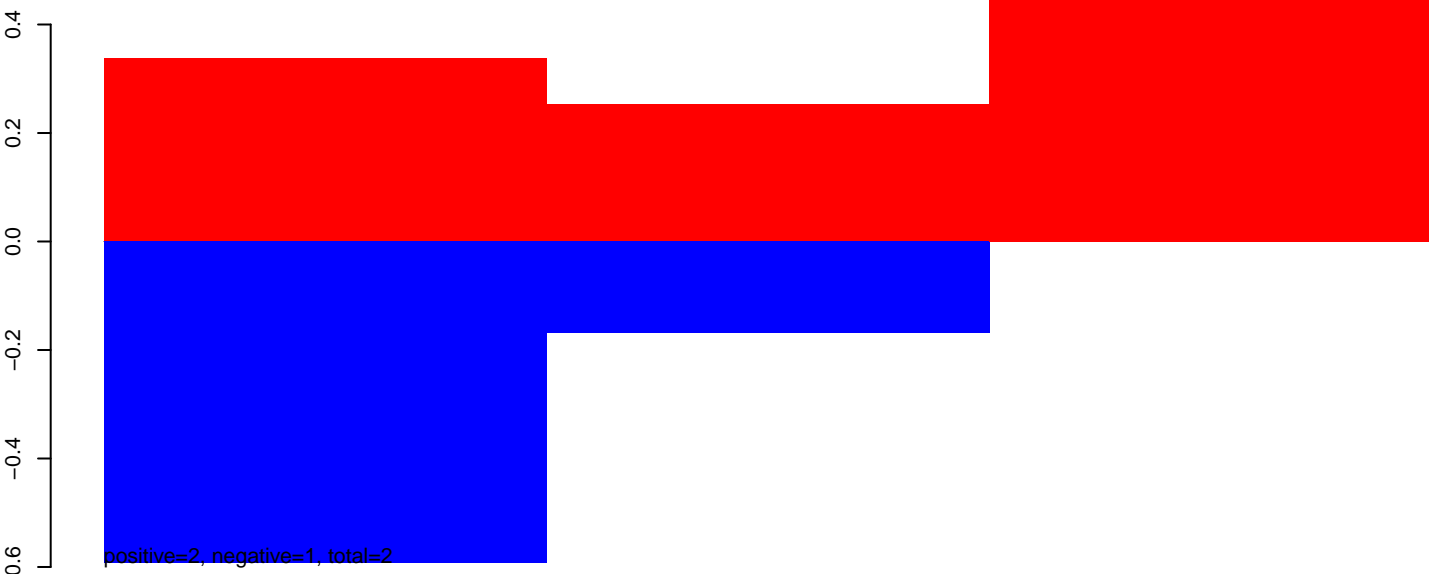
AnGam_Mos55_cells.24_35.rep



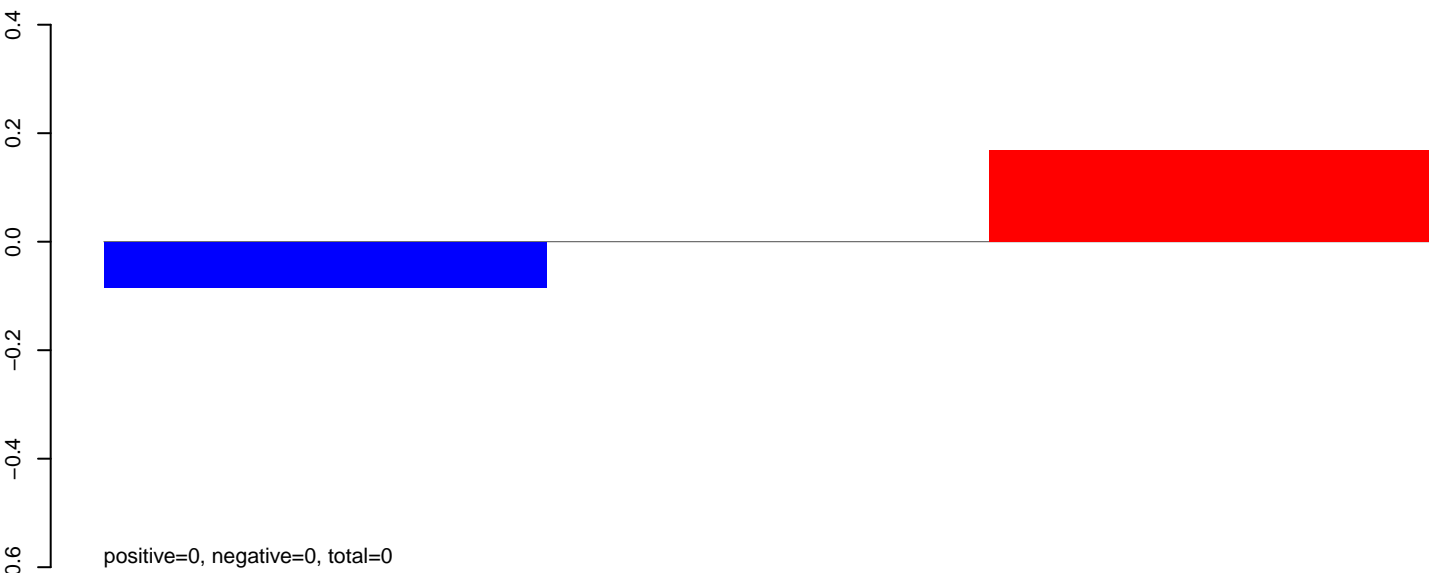
AnGam_Mos55_cells.rep



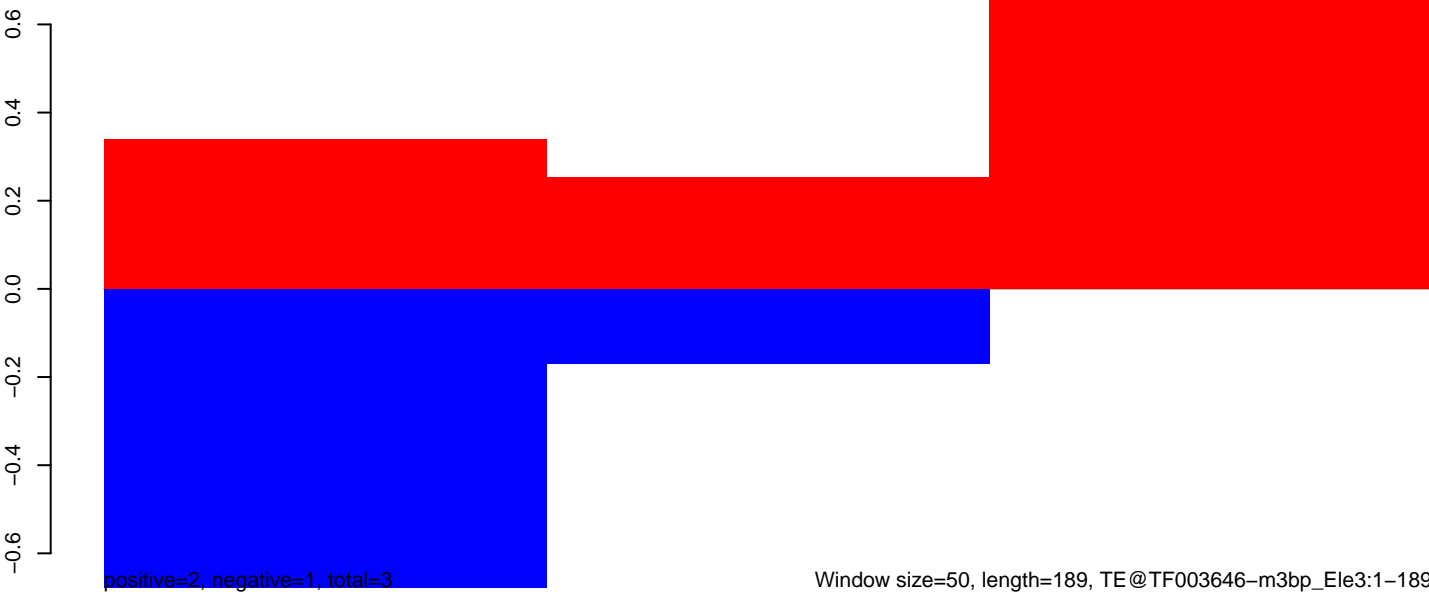
AnGam_Mos55_cells.18_23.rep



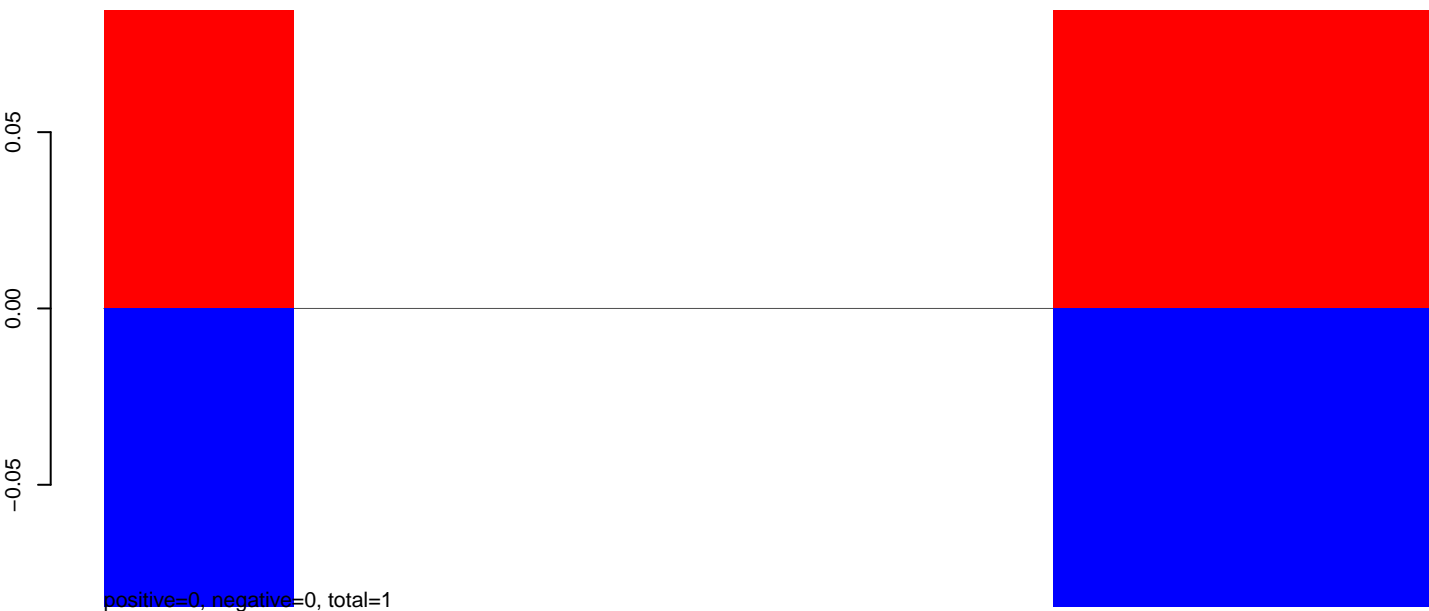
AnGam_Mos55_cells.24_35.rep



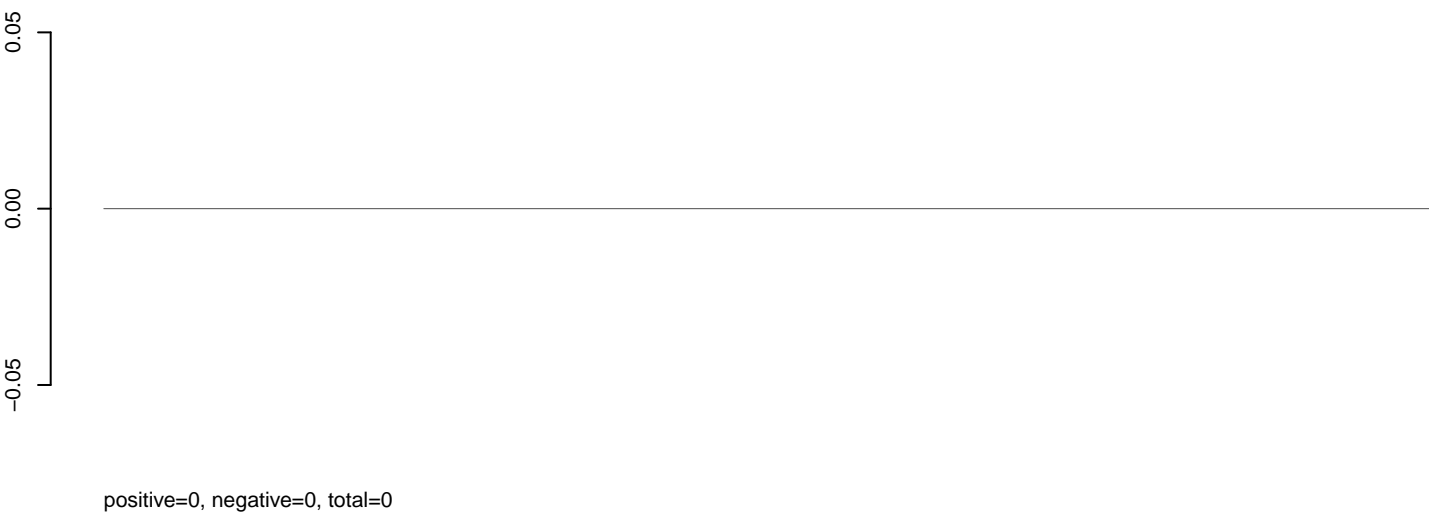
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



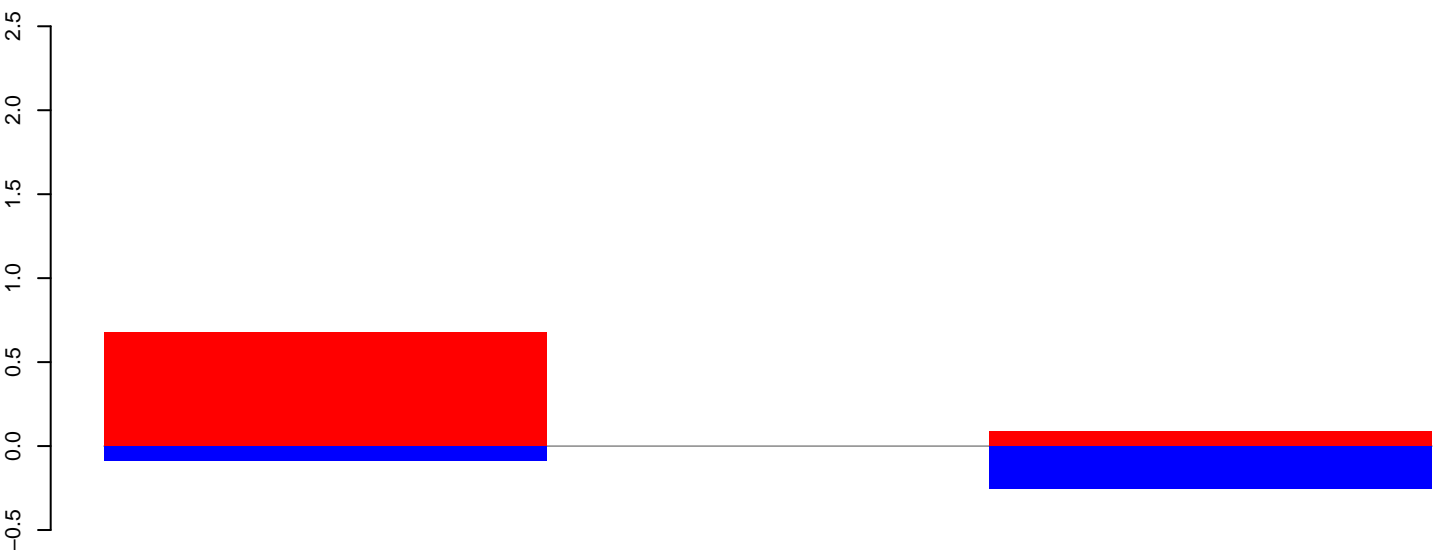
AnGam_Mos55_cells.rep



Window size=50, length=352, TE@TF003640-m3bp_Ele24:1-352

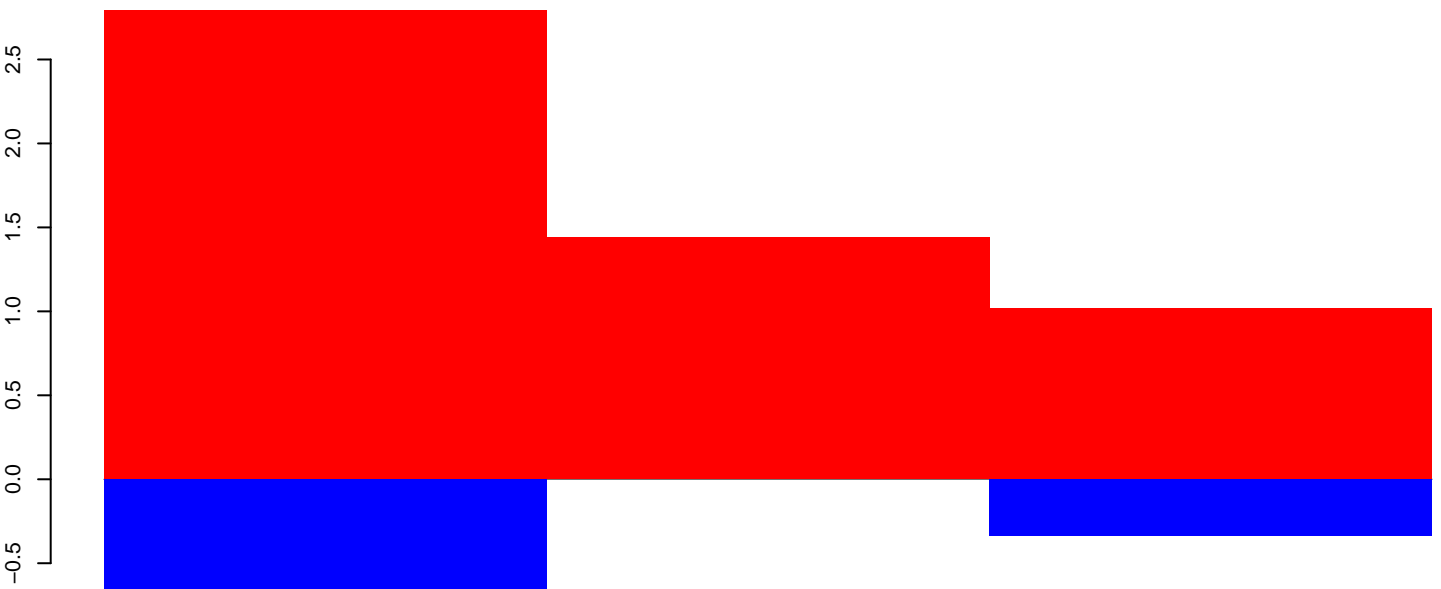
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AnGam_Mos55_cells.18_23.rep



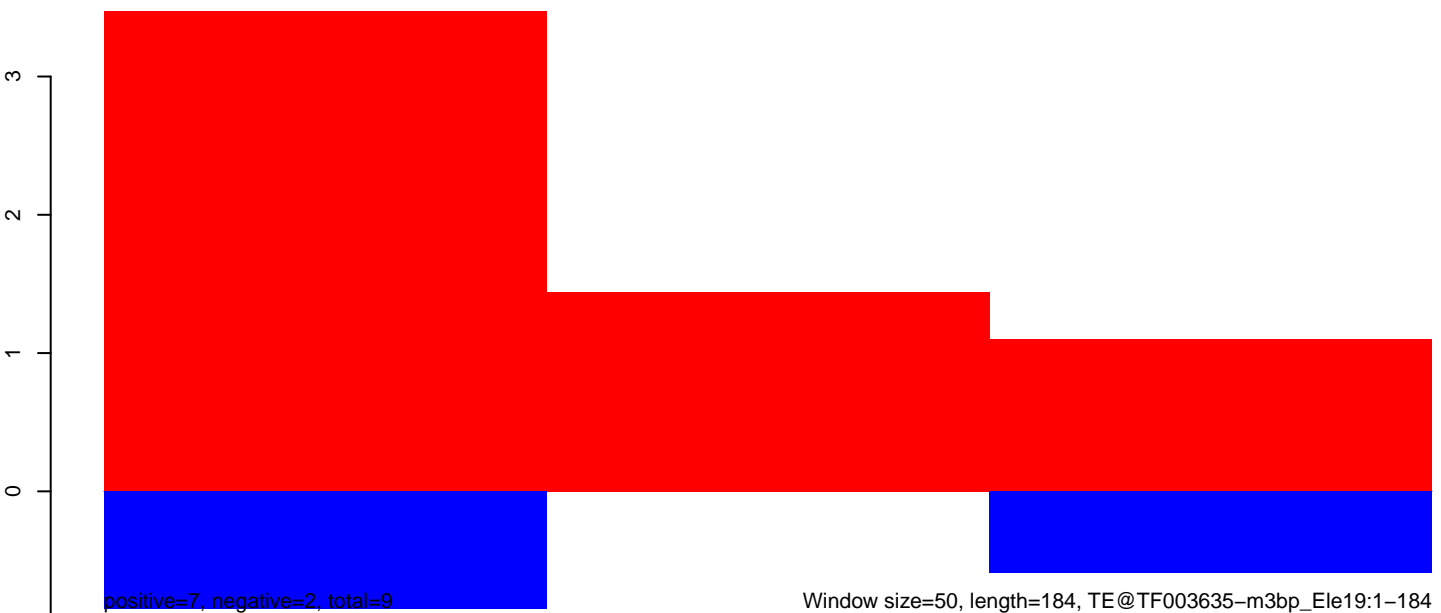
positive=1, negative=1, total=1

AnGam_Mos55_cells.24_35.rep



positive=6, negative=1, total=8

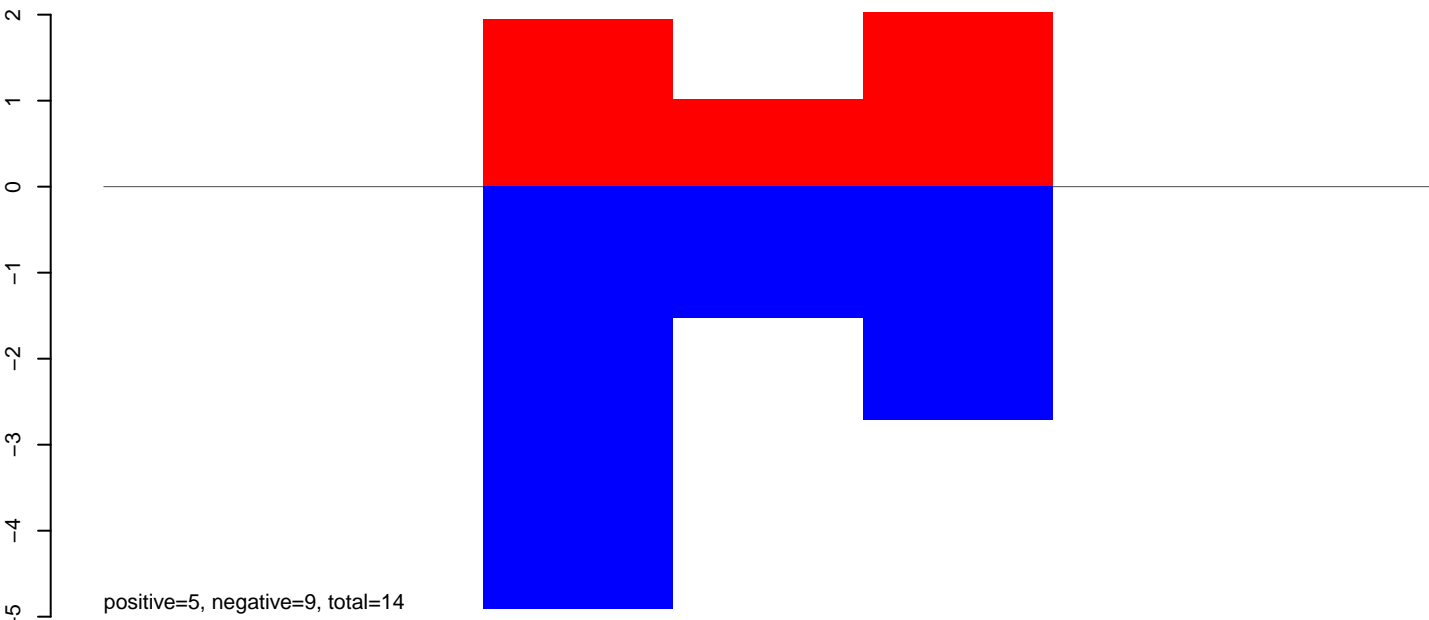
AnGam_Mos55_cells.rep



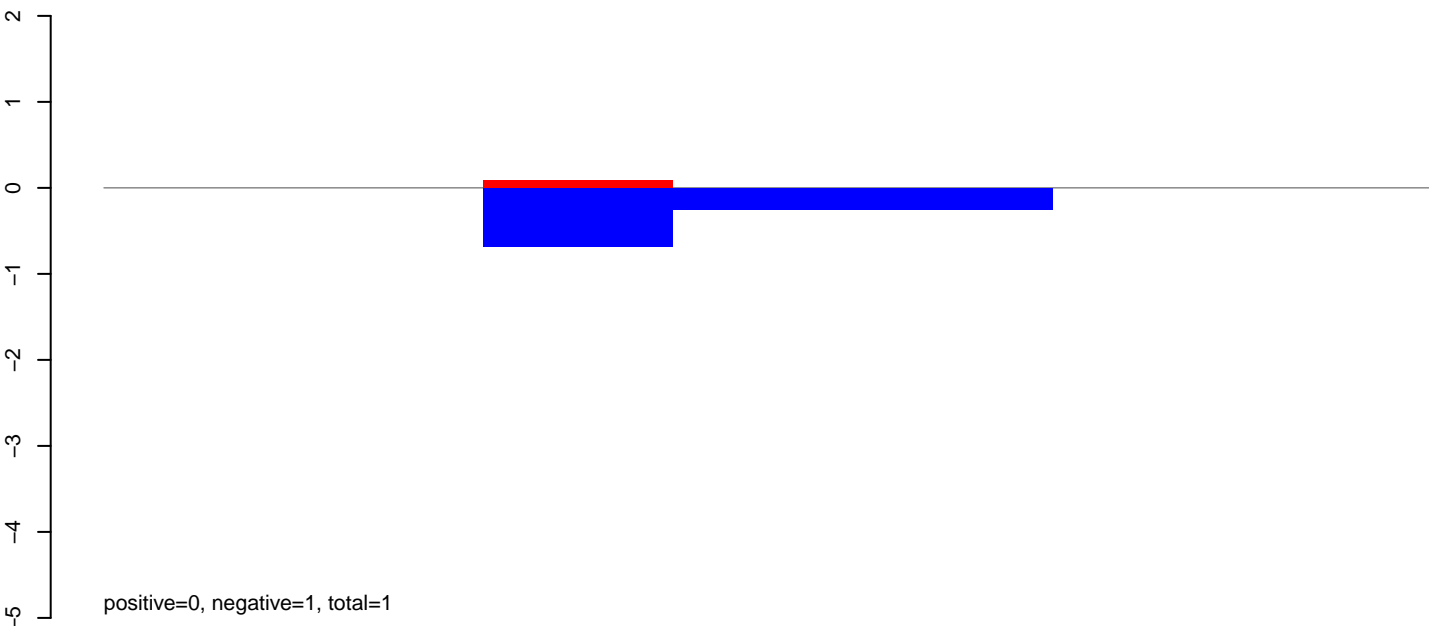
positive=7, negative=2, total=9

Window size=50, length=184, TE@TF003635-m3bp_Ele19:1-184

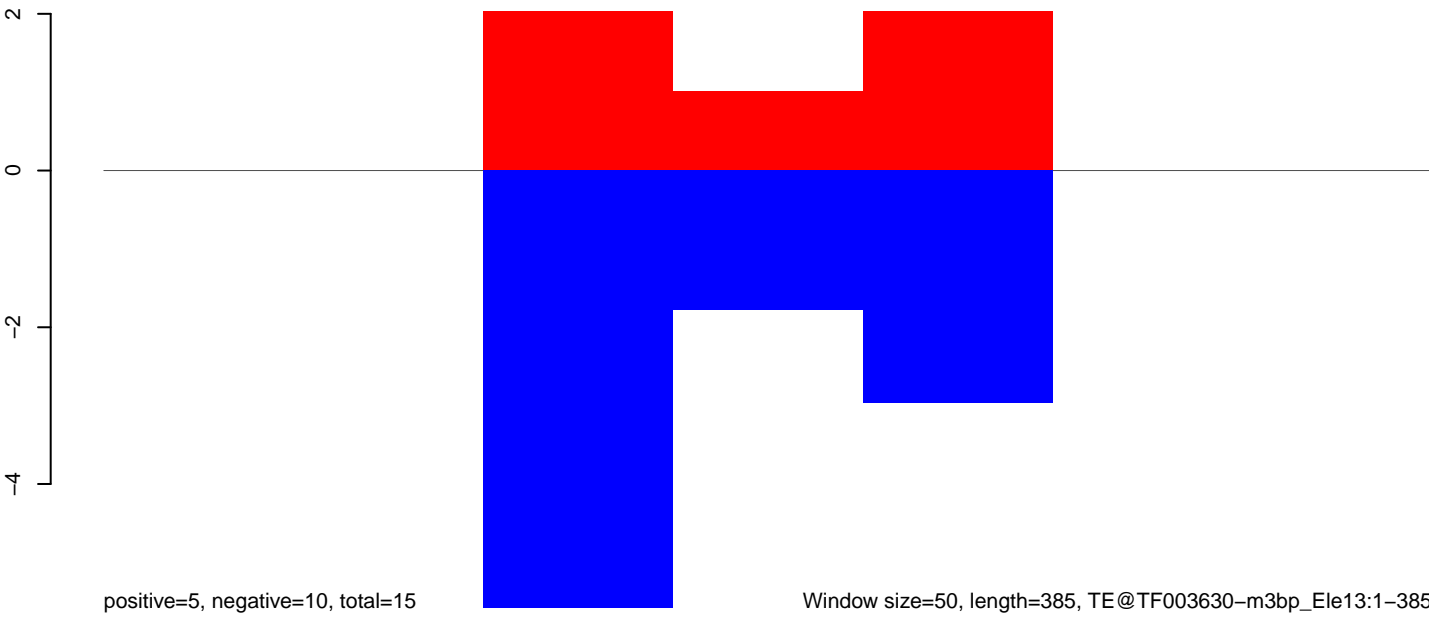
AnGam_Mos55_cells.18_23.rep



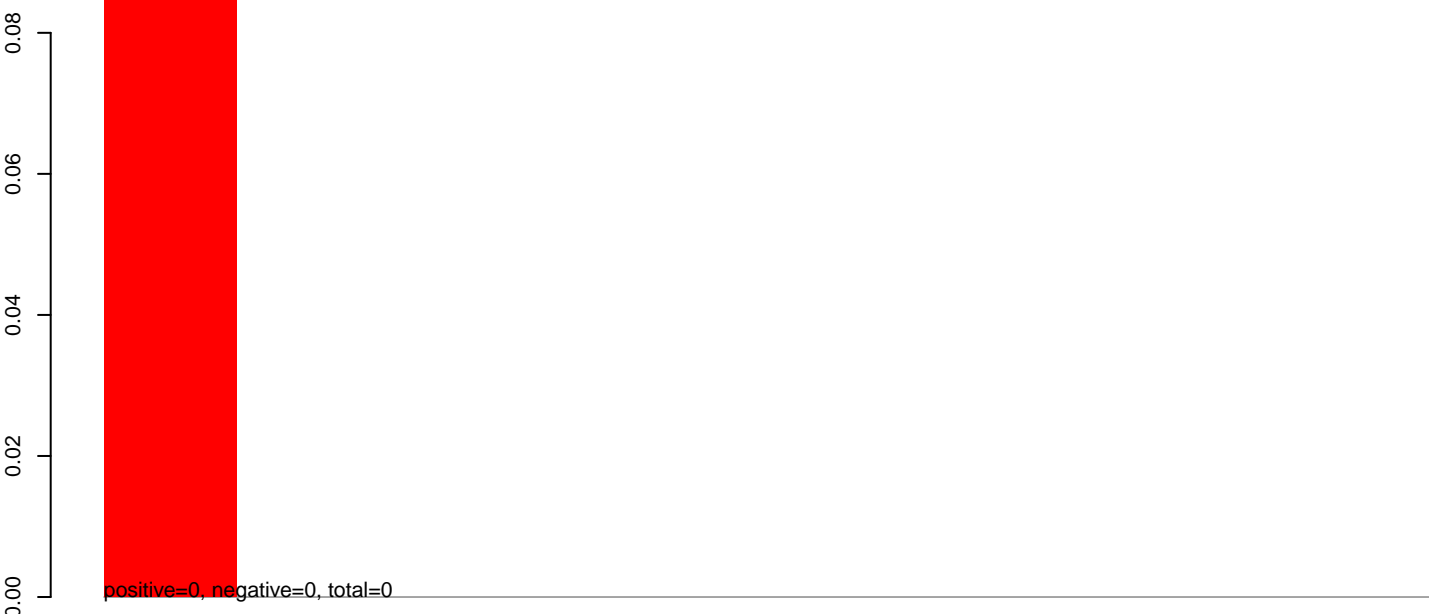
AnGam_Mos55_cells.24_35.rep



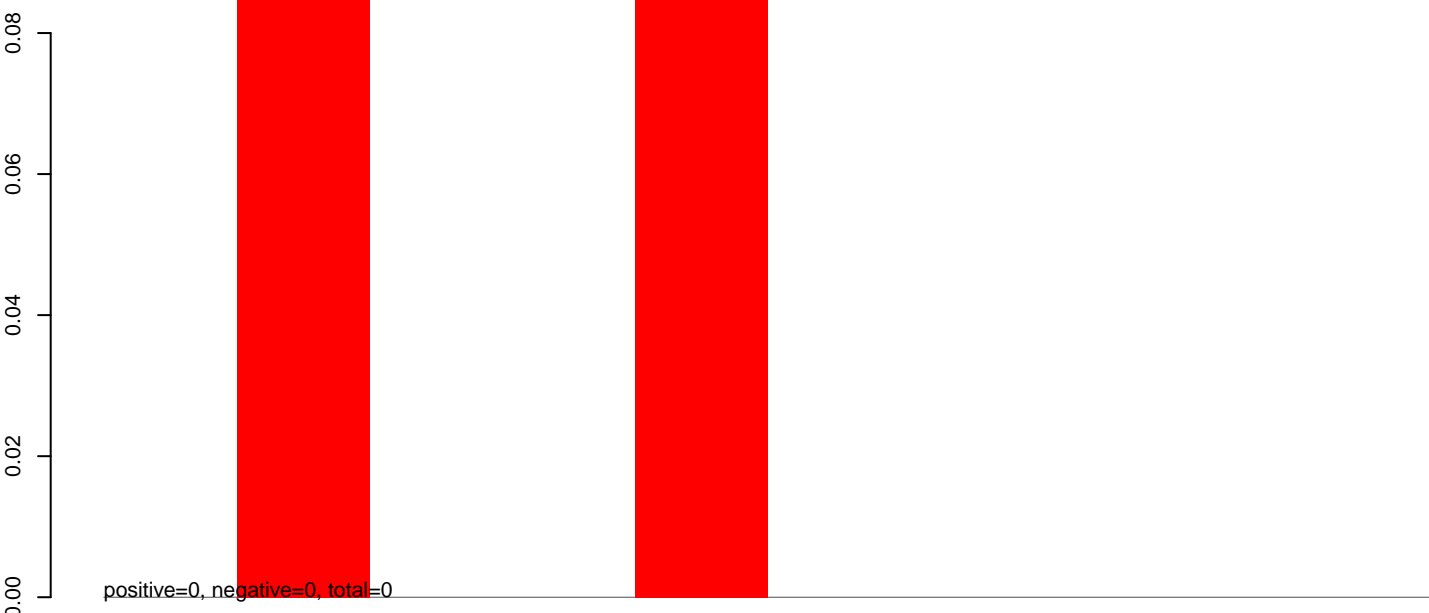
AnGam_Mos55_cells.rep



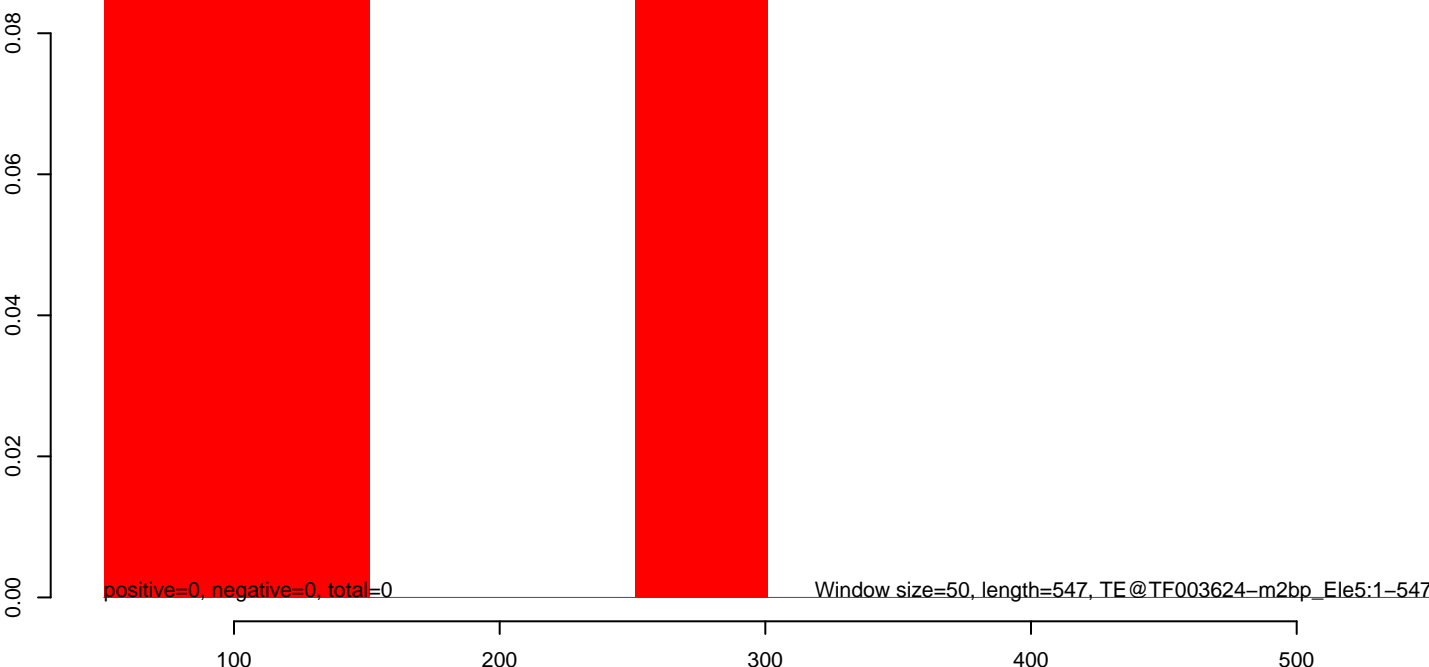
AnGam_Mos55_cells.18_23.rep



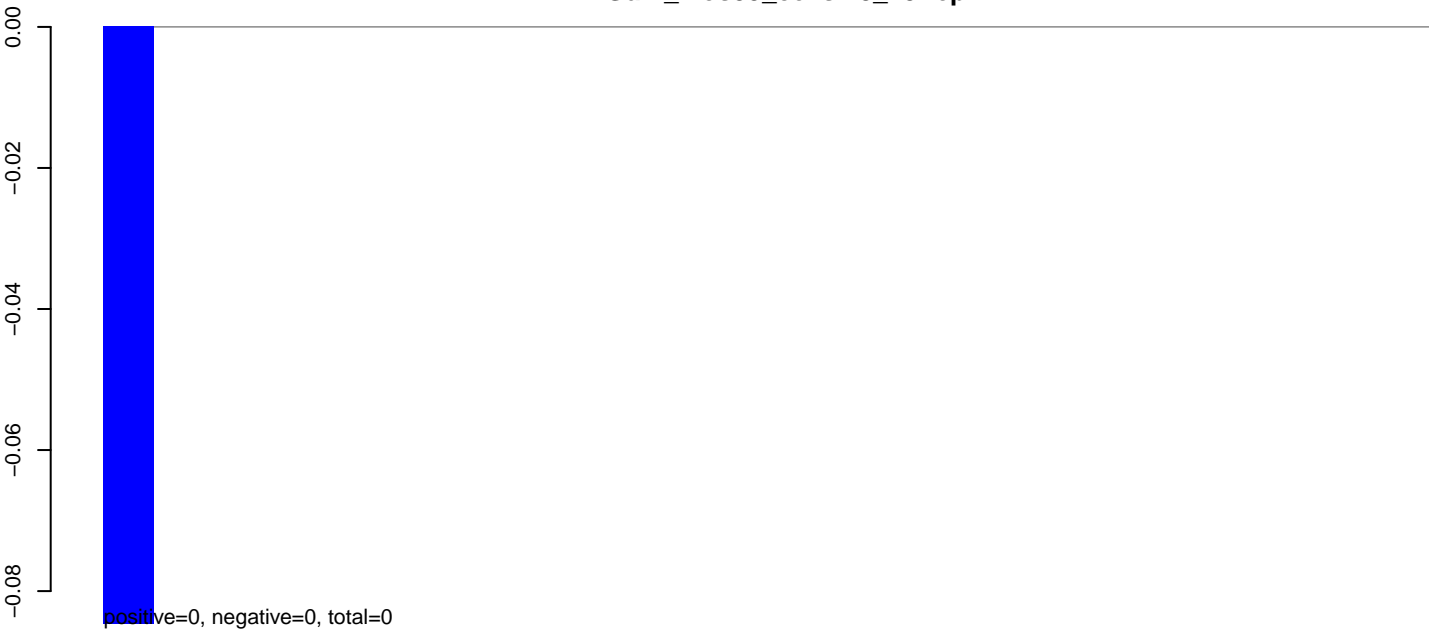
AnGam_Mos55_cells.24_35.rep



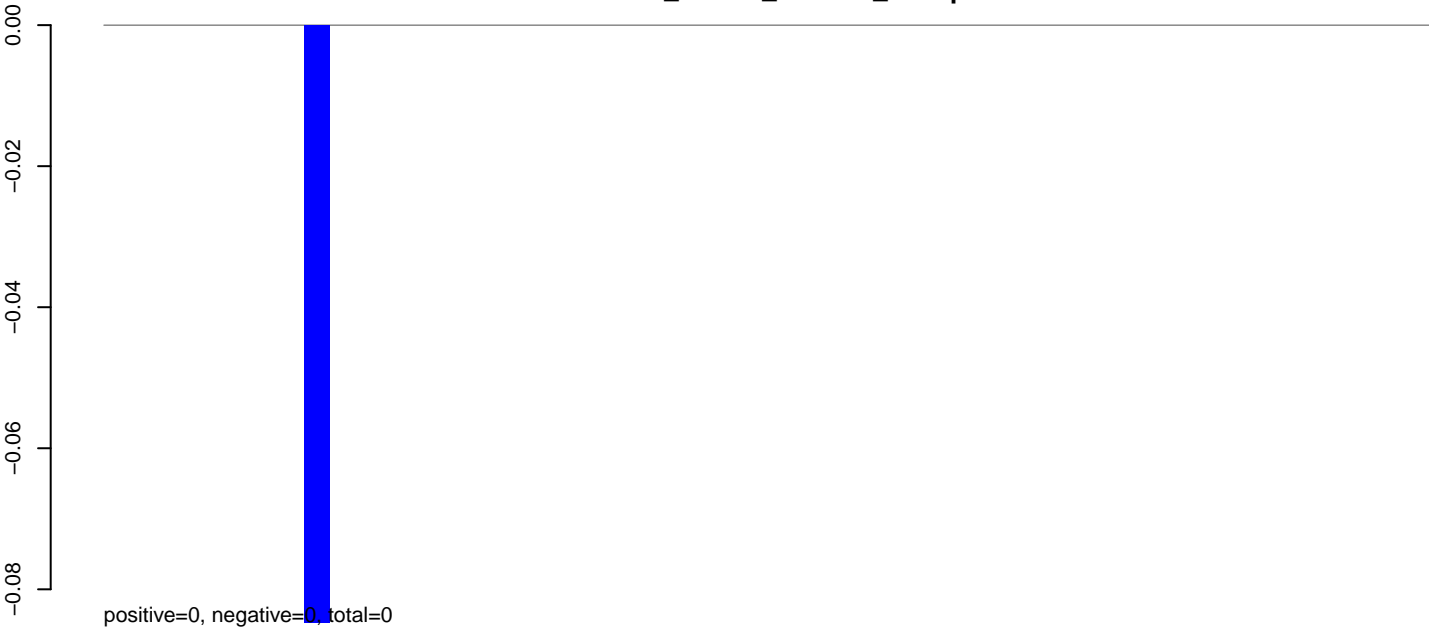
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



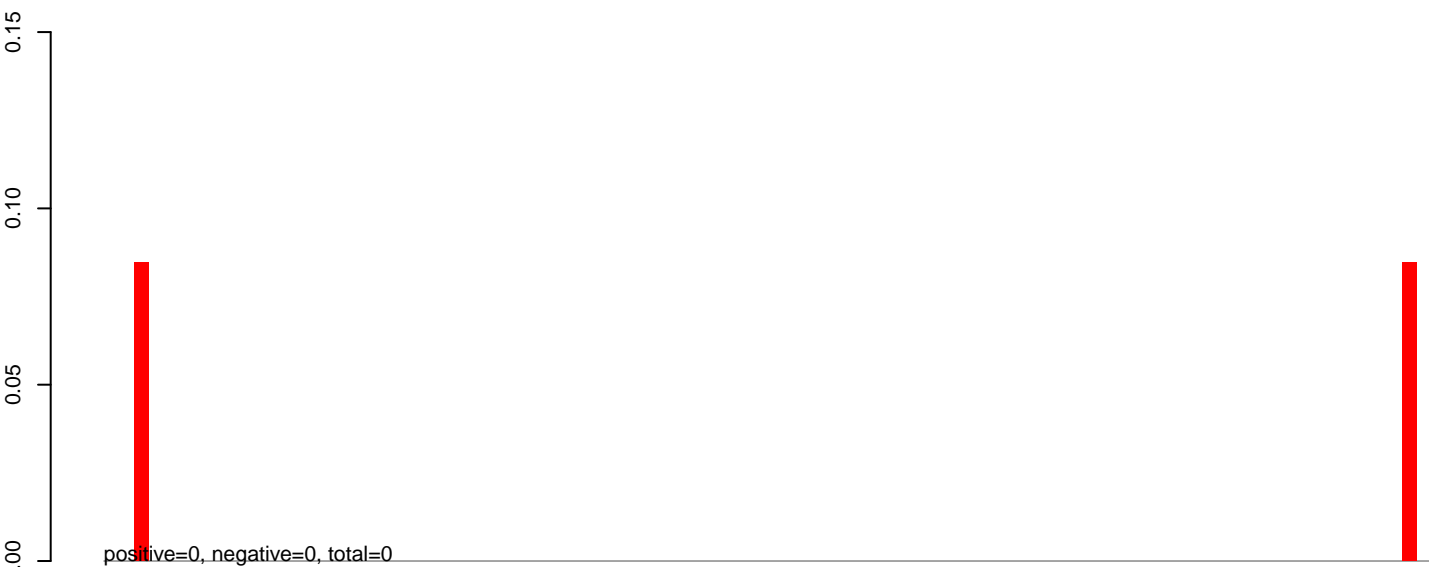
AnGam_Mos55_cells.24_35.rep



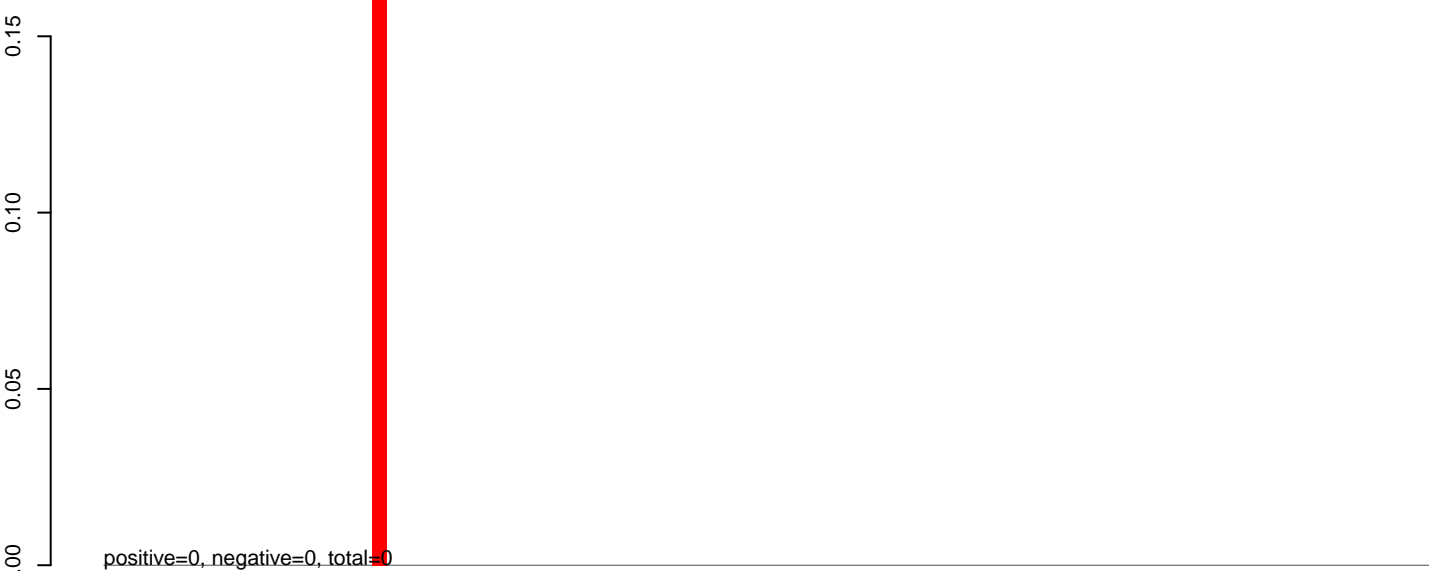
AnGam_Mos55_cells.rep



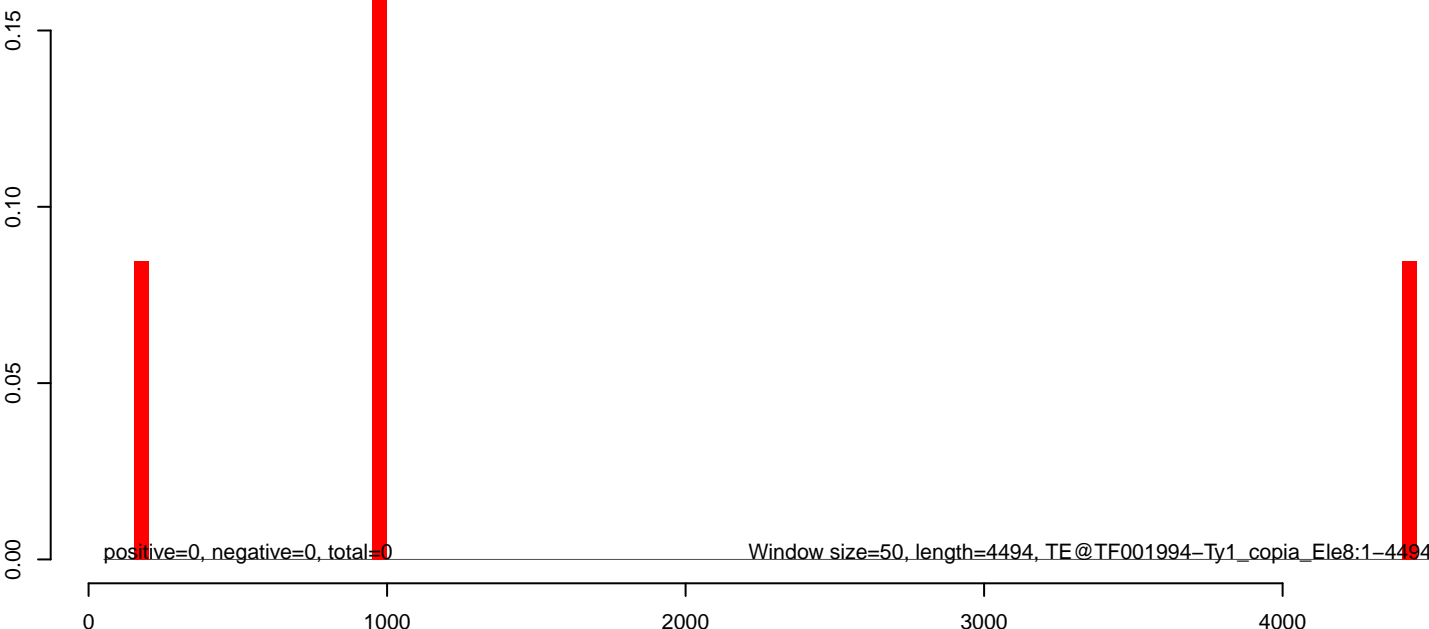
AnGam_Mos55_cells.18_23.rep



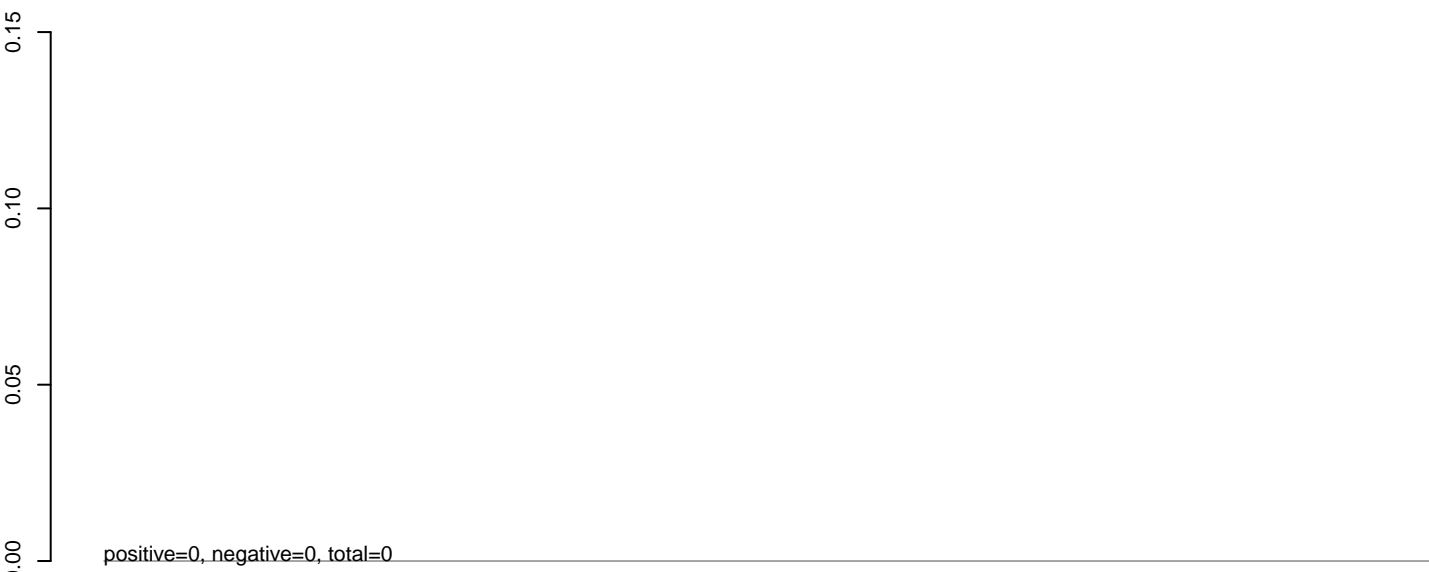
AnGam_Mos55_cells.24_35.rep



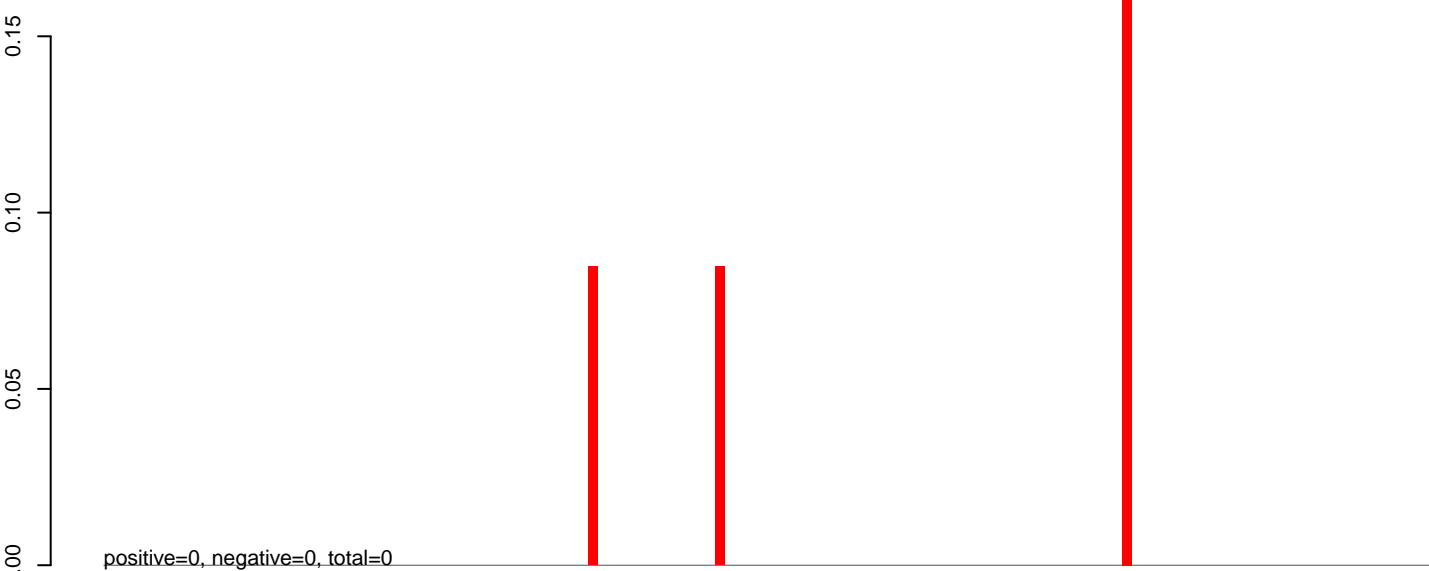
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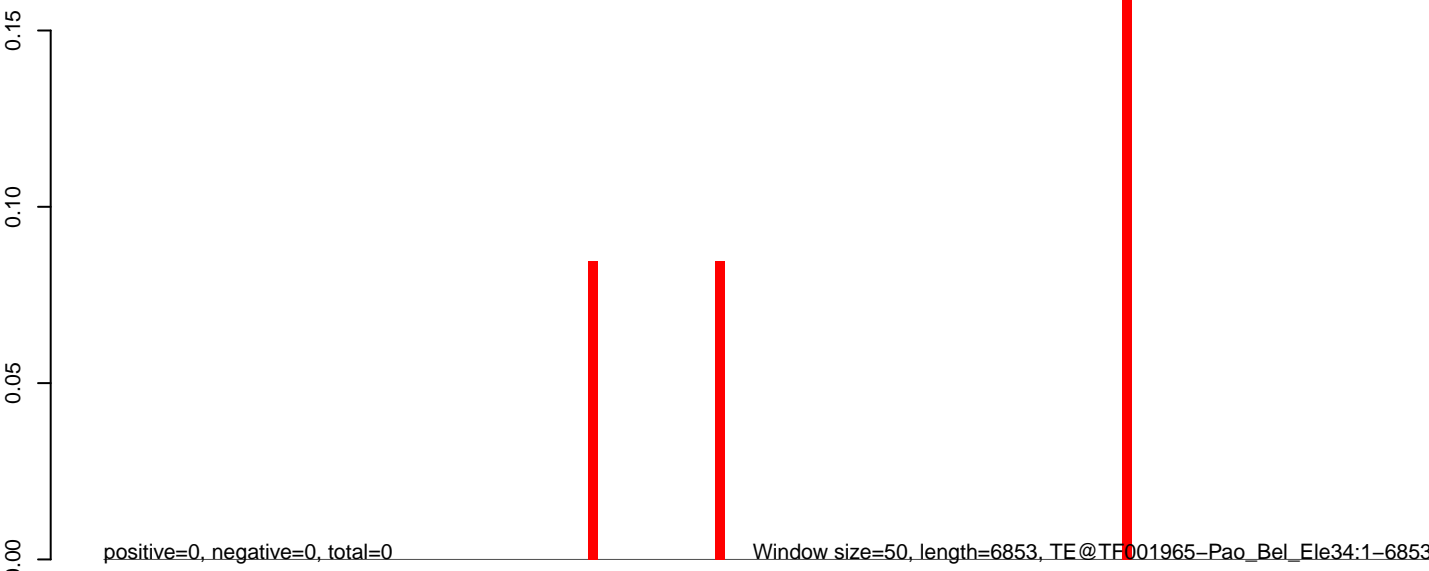
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

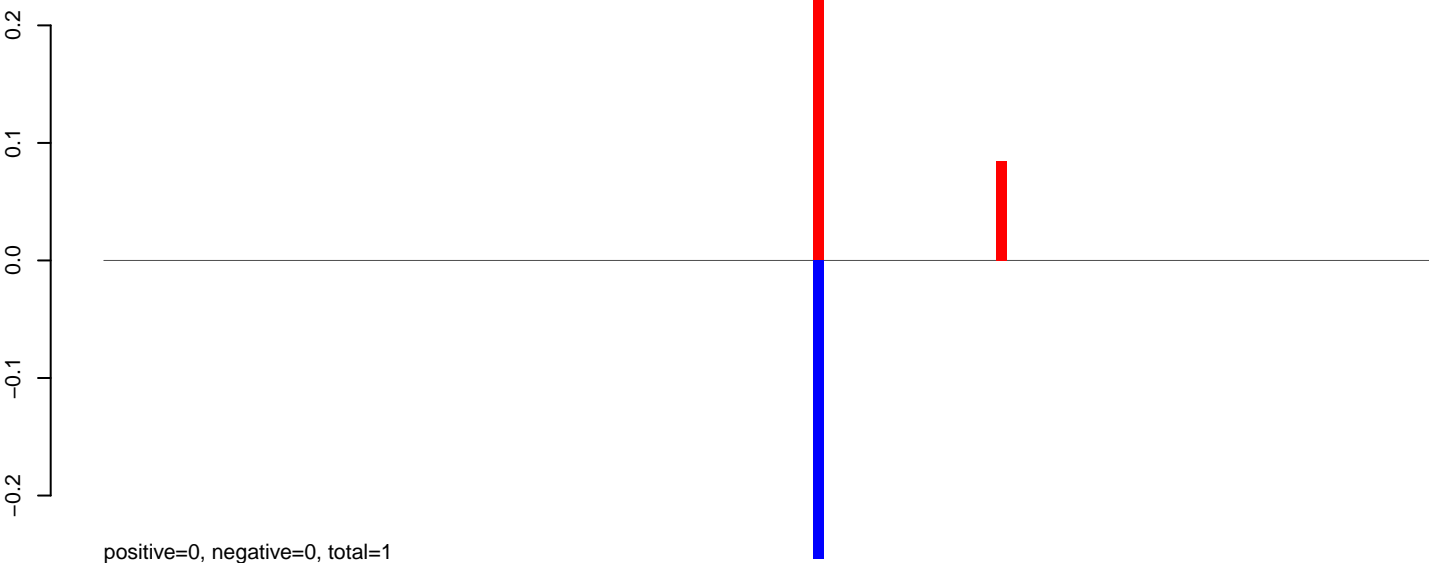


AnGam_Mos55_cells.rep

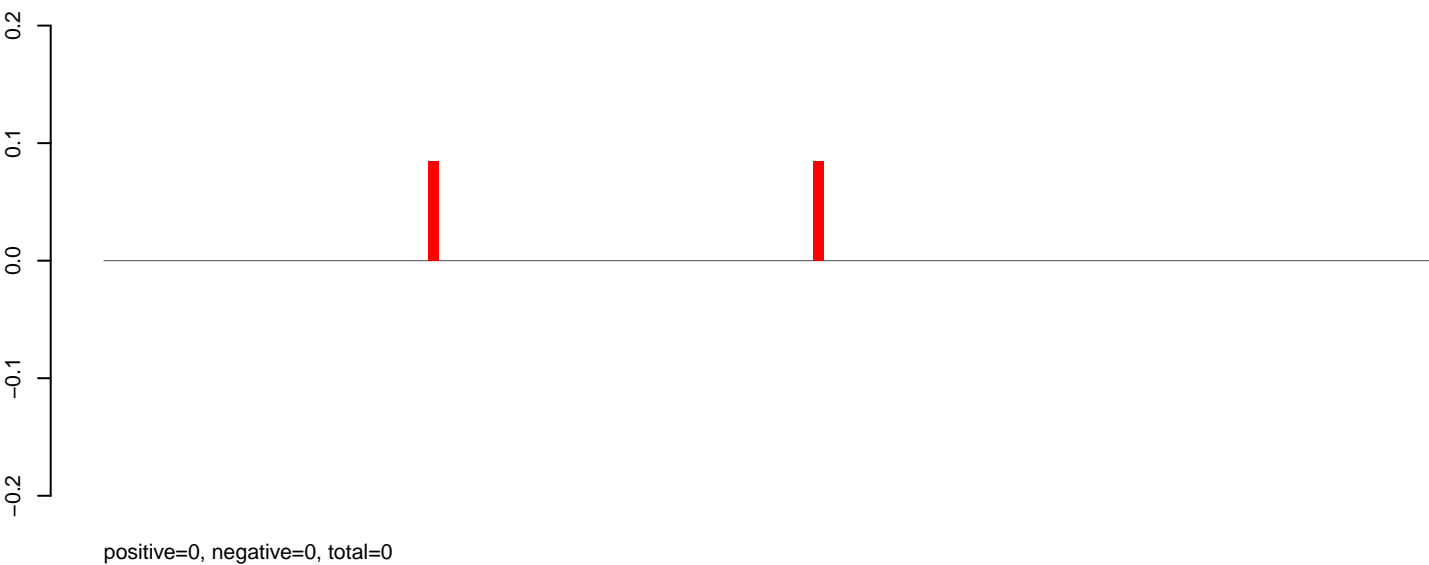


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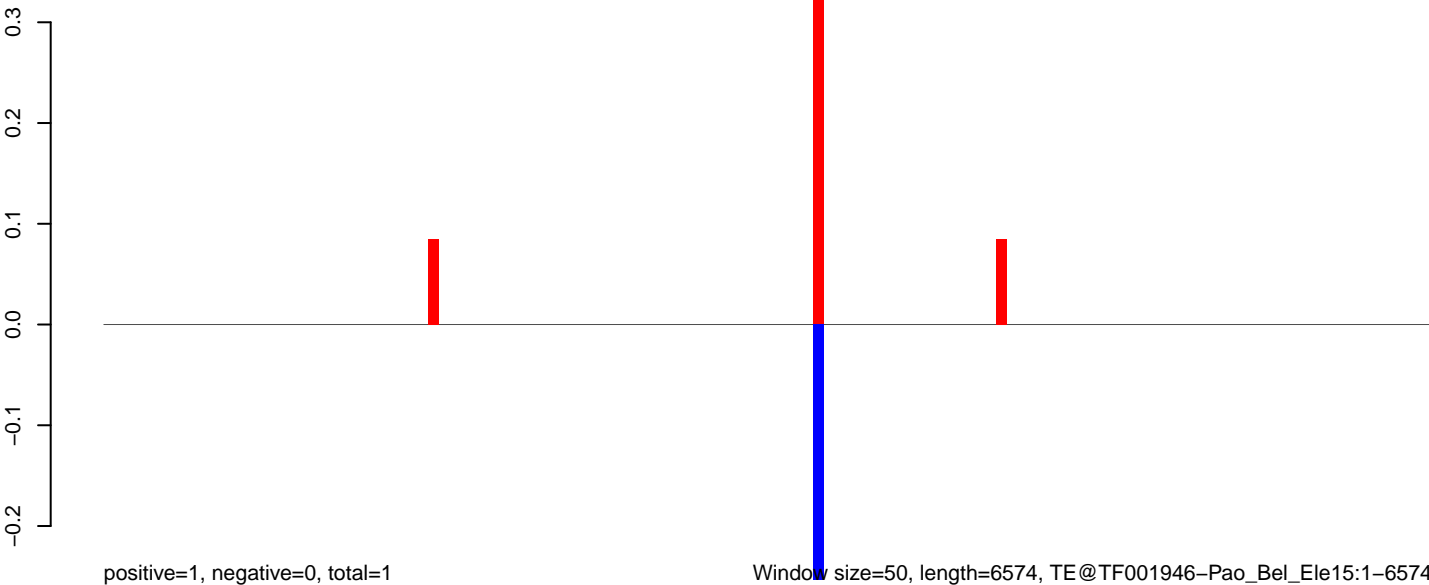
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

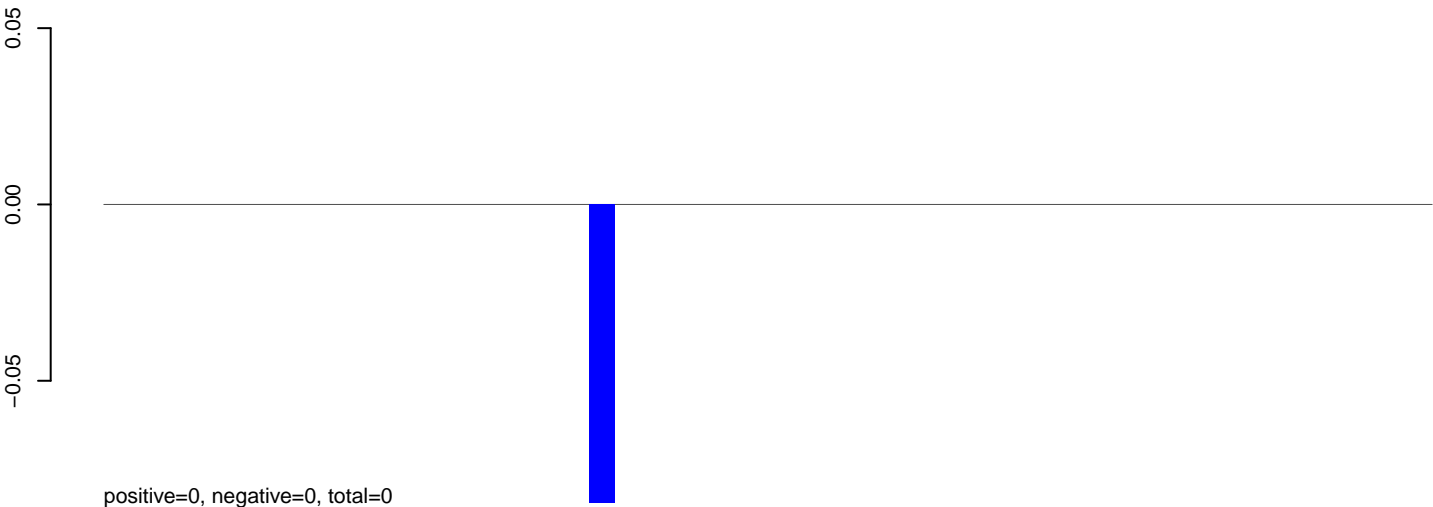


AnGam_Mos55_cells.rep

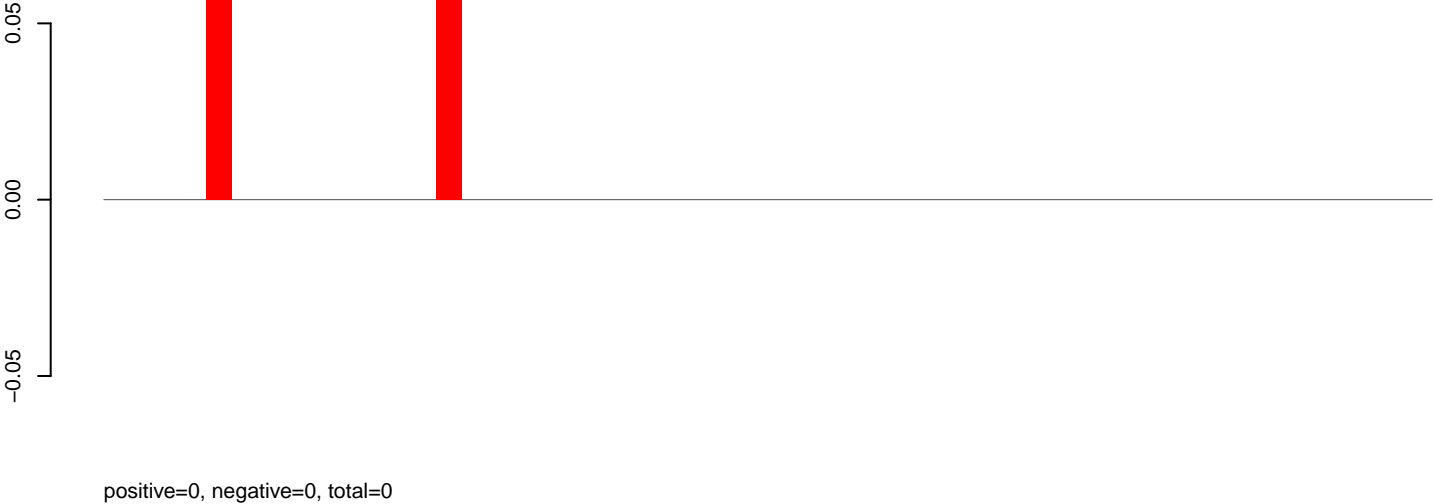


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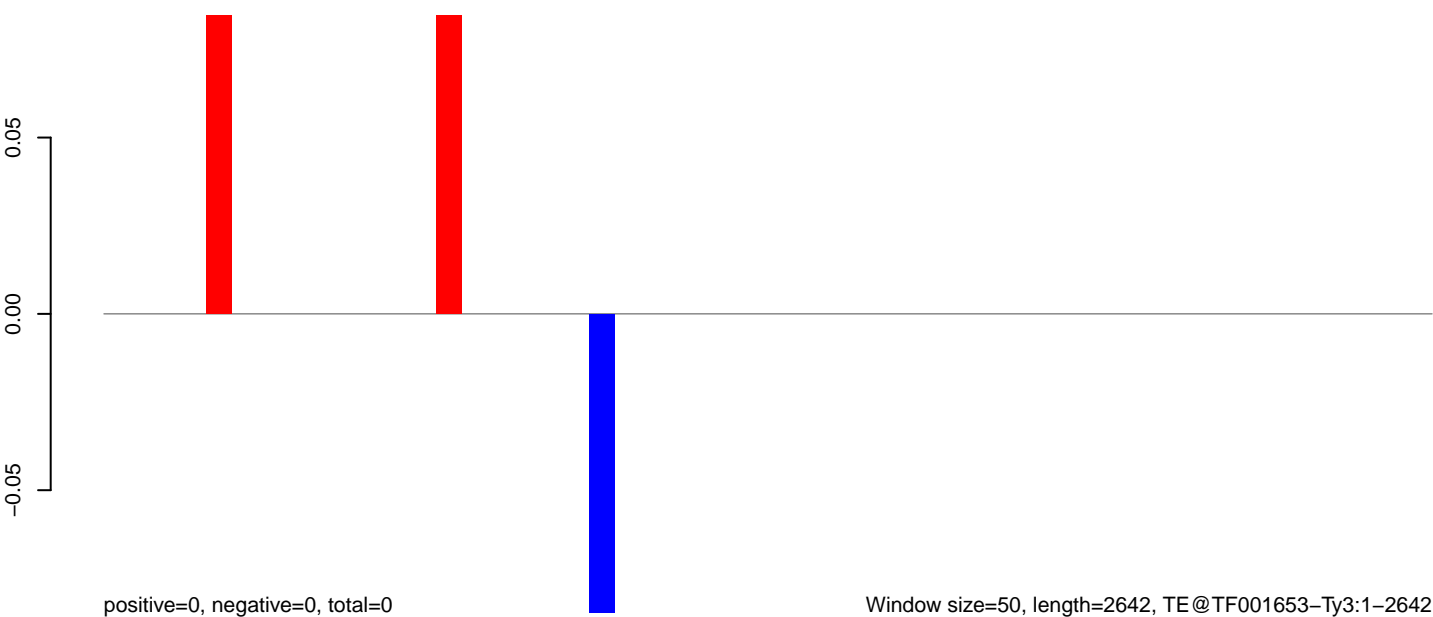
AnGam_Mos55_cells.18_23.rep



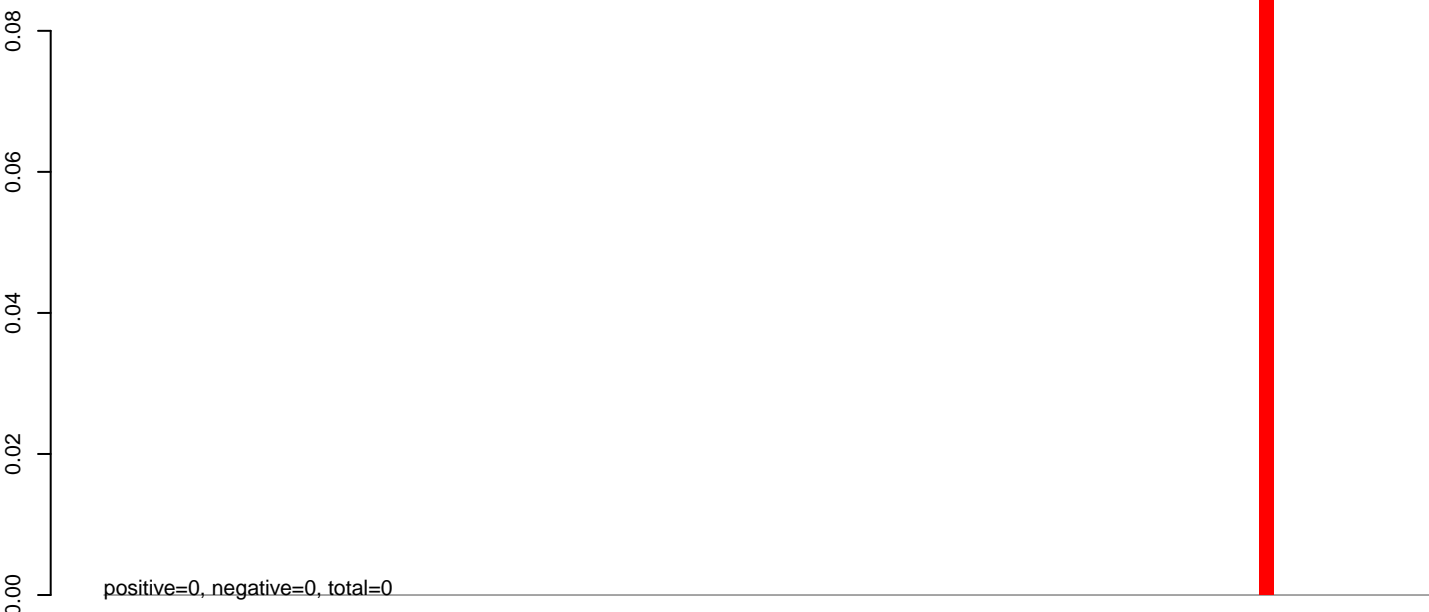
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



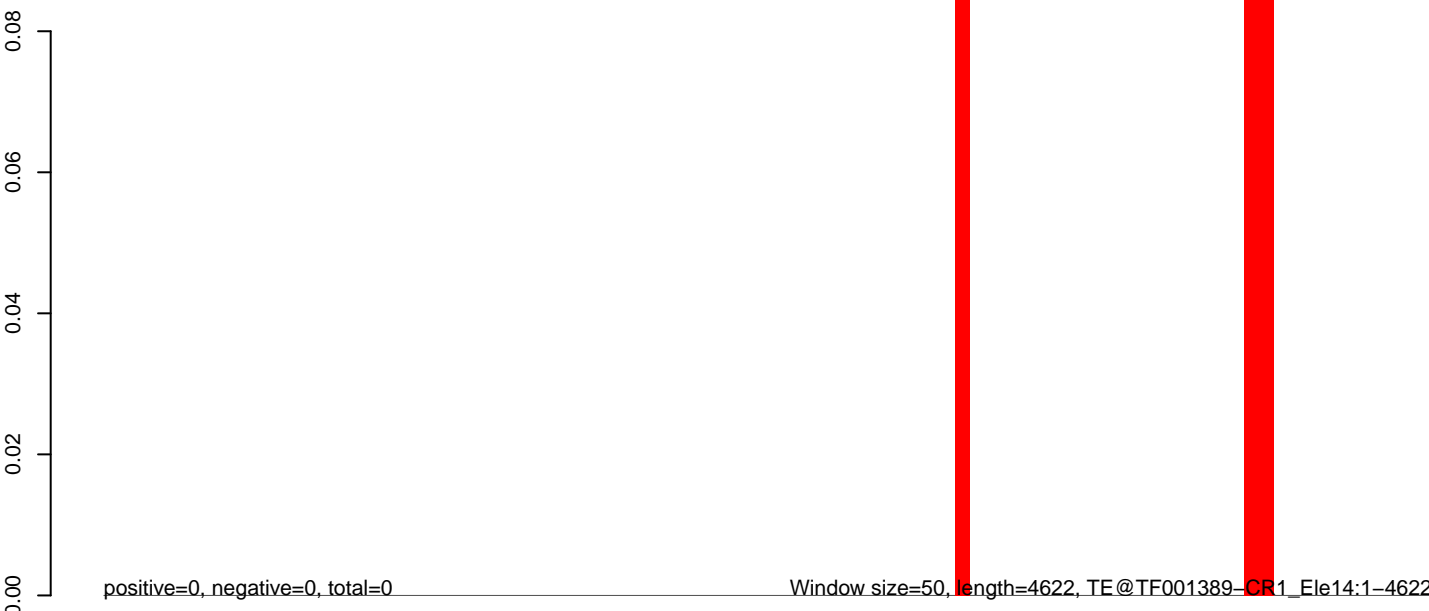
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

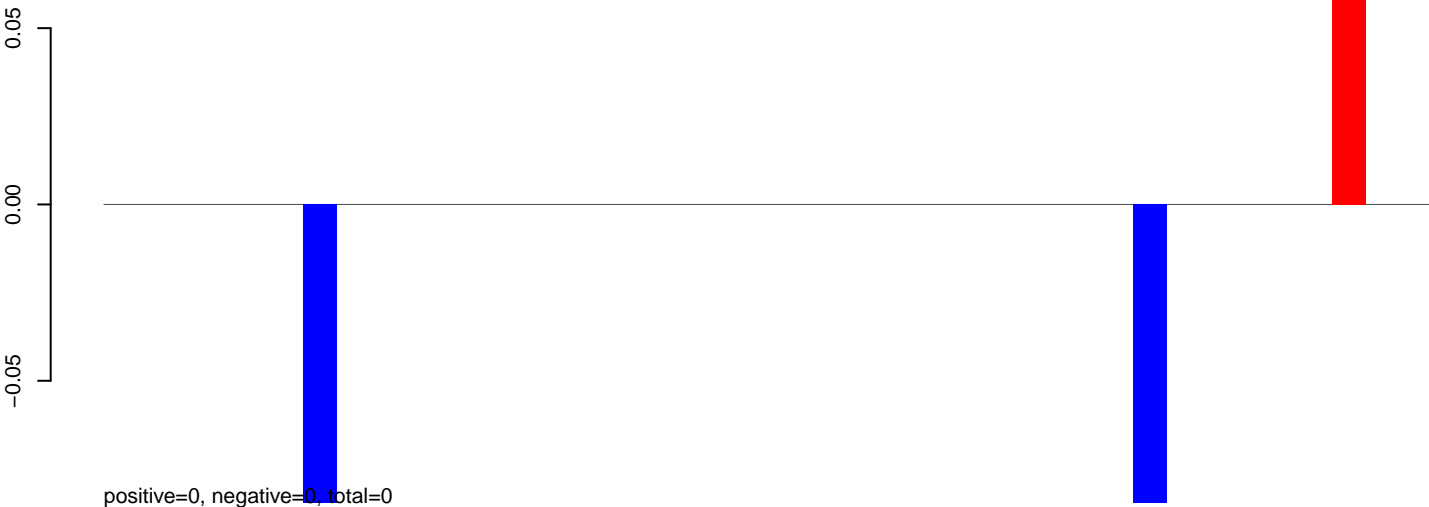


AnGam_Mos55_cells.rep

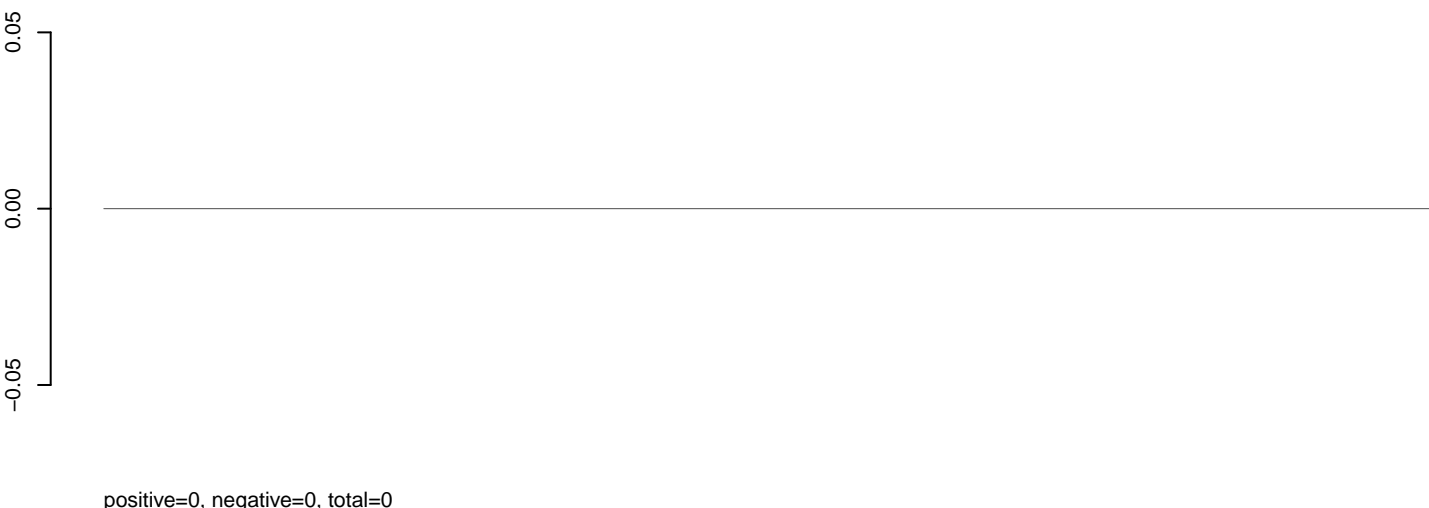


0 1000 2000 3000 4000

AnGam_Mos55_cells.18_23.rep



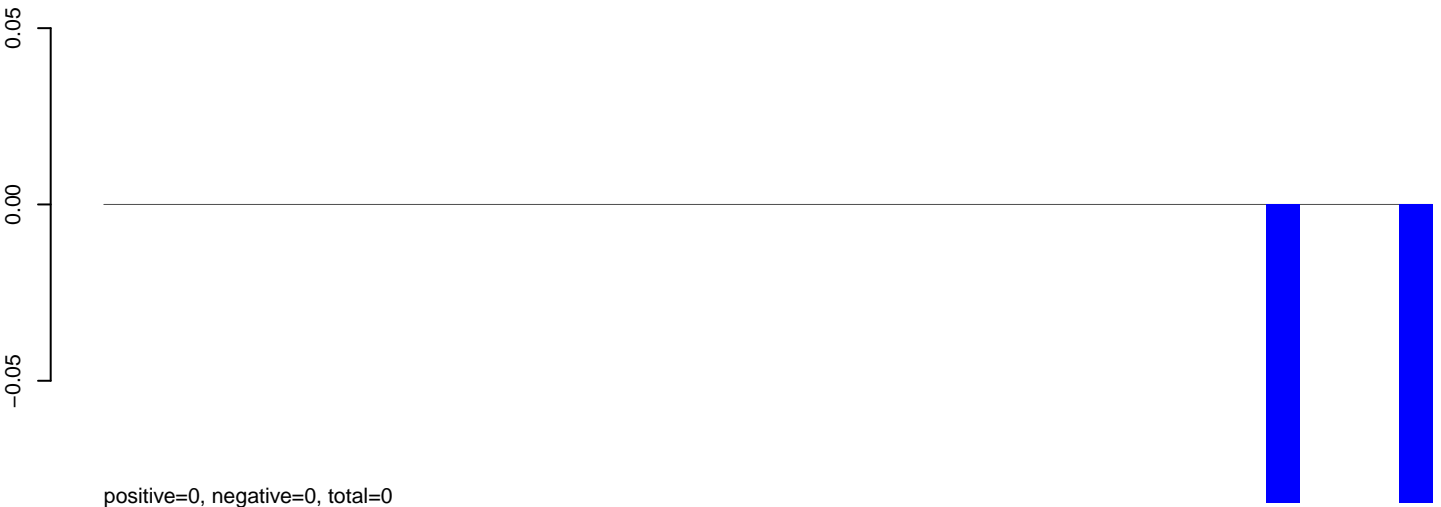
AnGam_Mos55_cells.24_35.rep



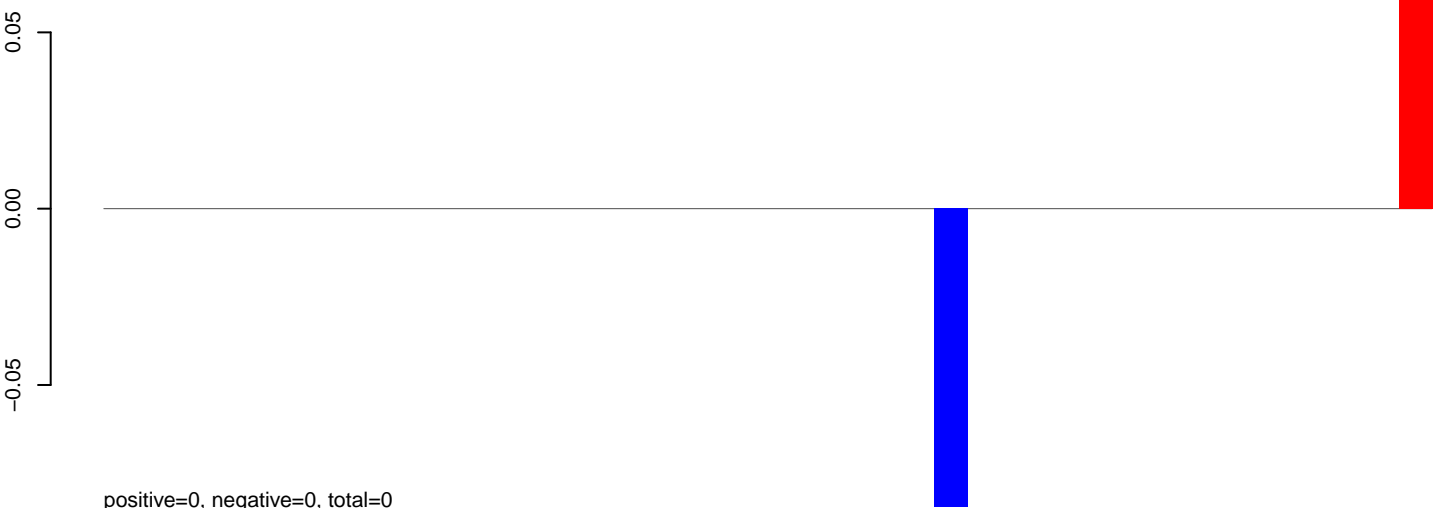
AnGam_Mos55_cells.rep



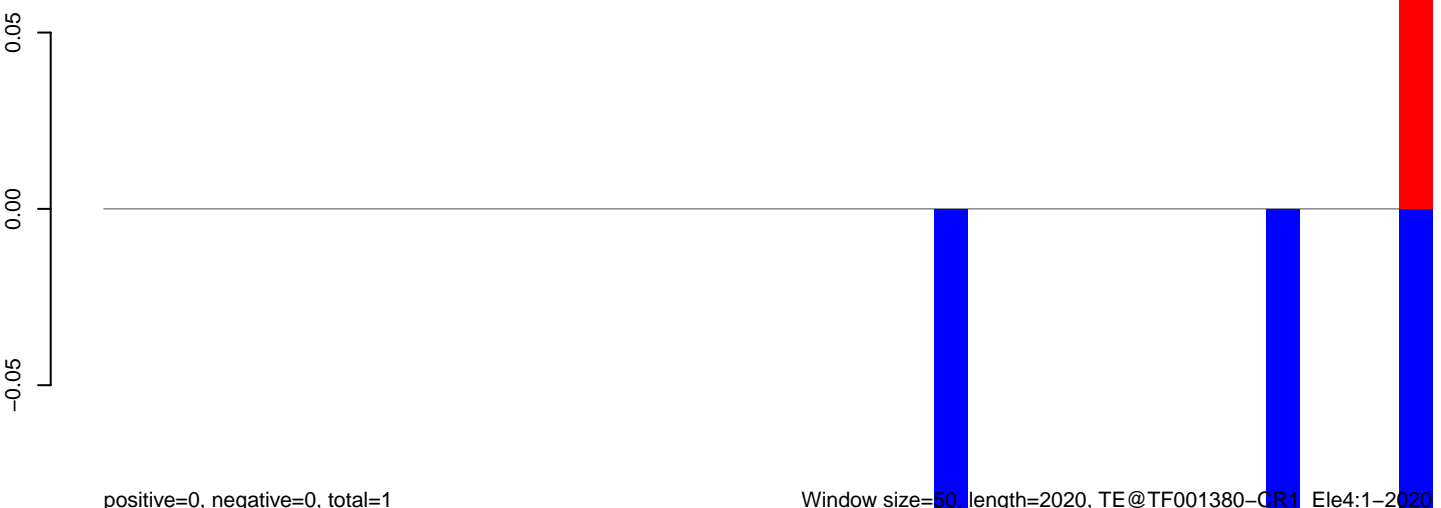
AnGam_Mos55_cells.18_23.rep



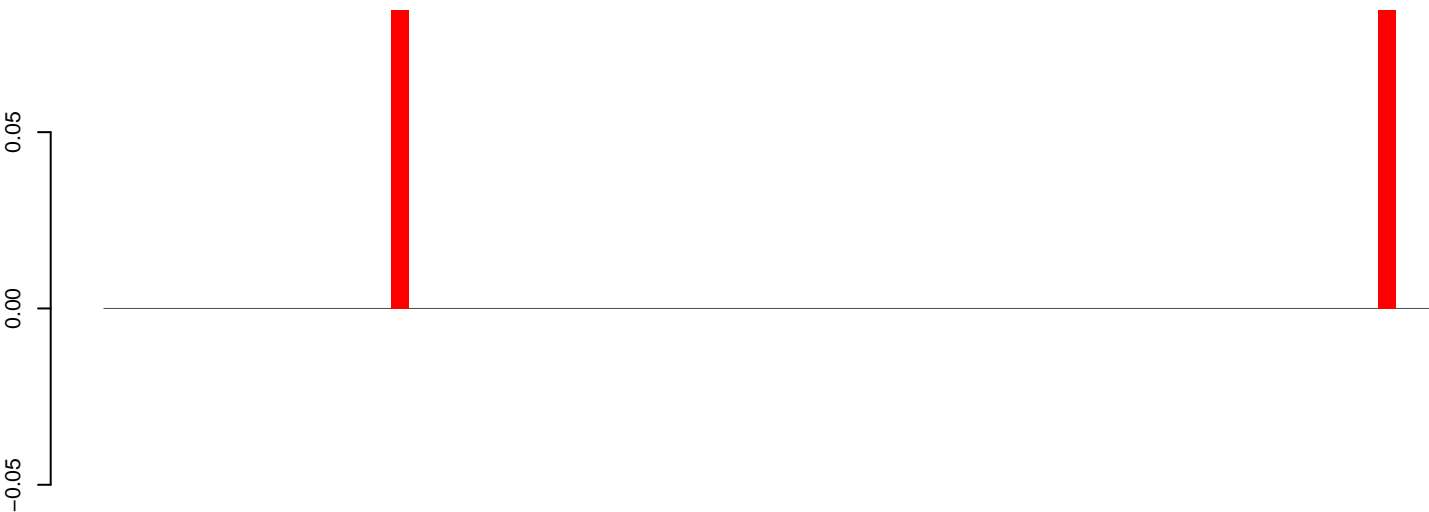
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

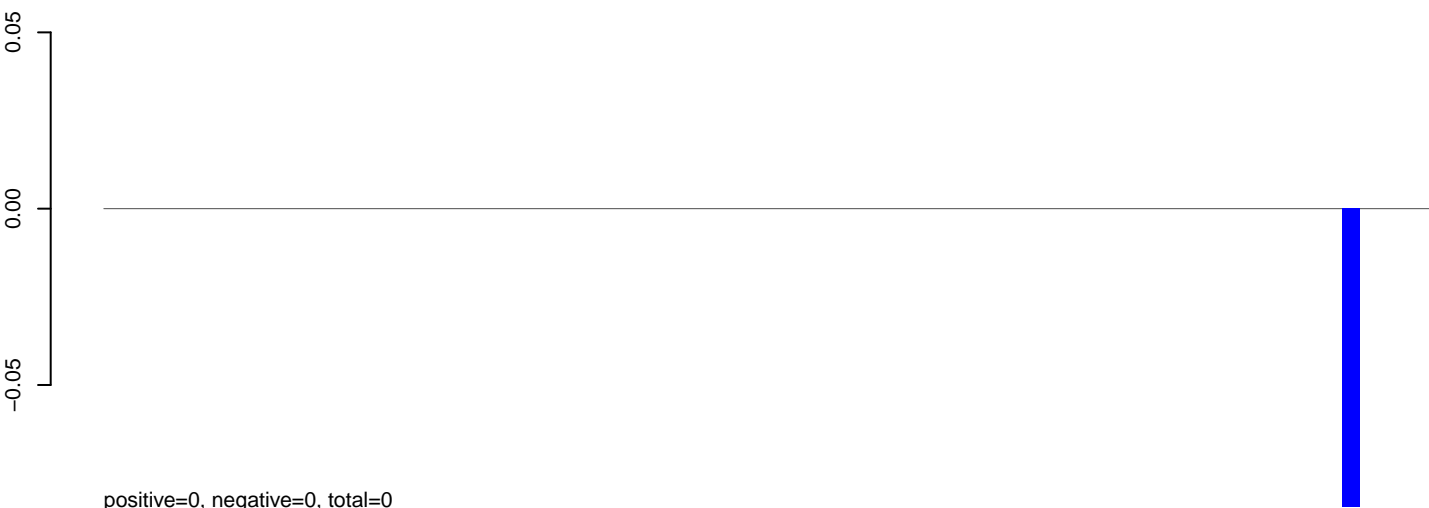


AnGam_Mos55_cells.18_23.rep



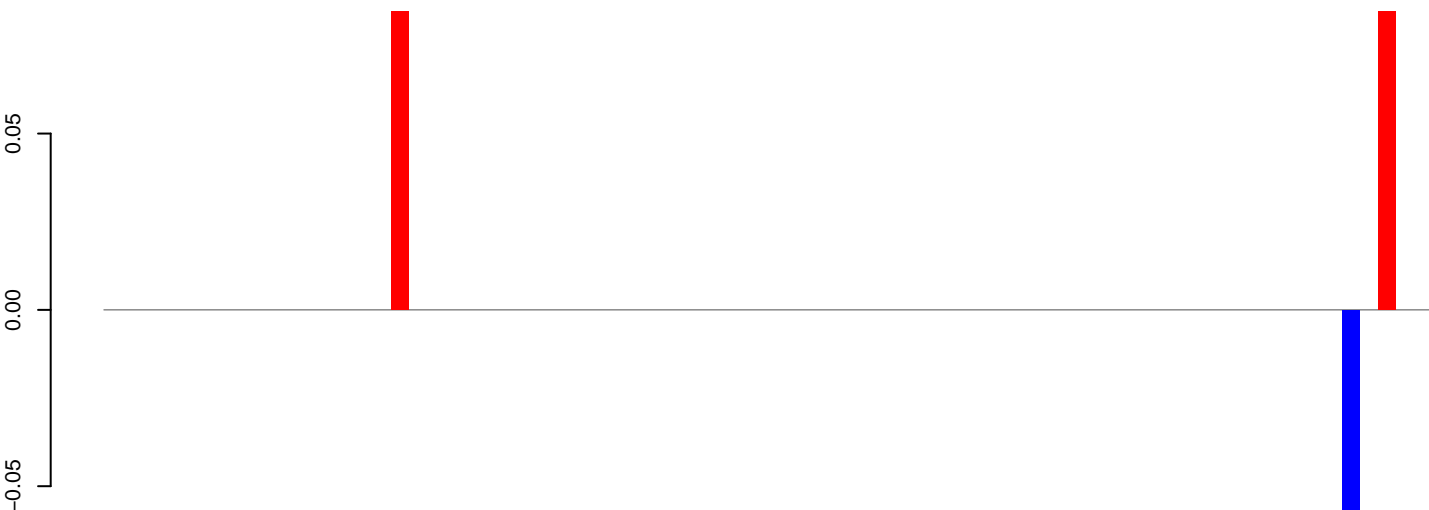
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

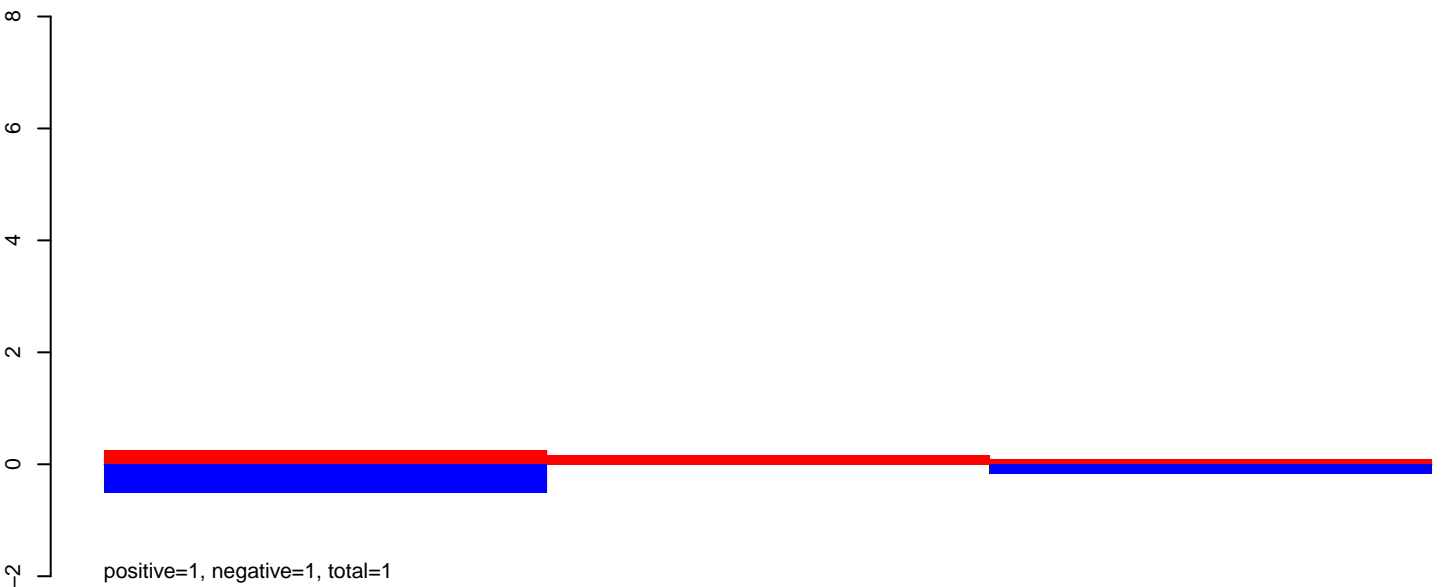


positive=0, negative=0, total=0

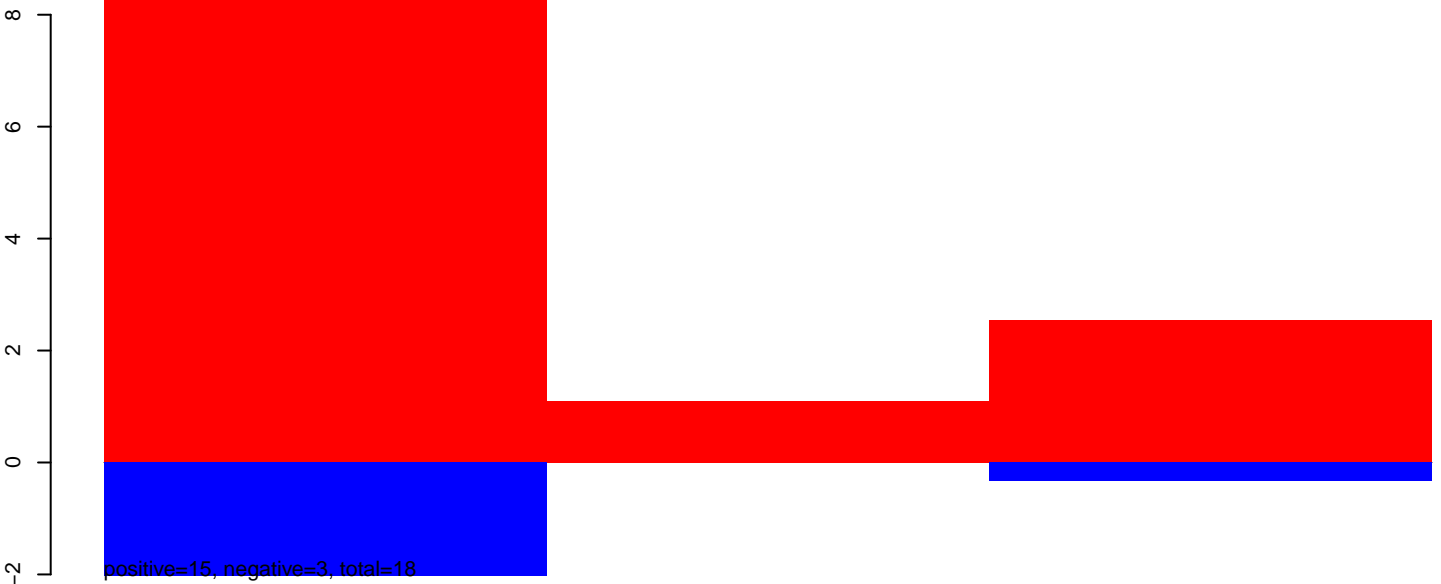
Window size=50, length=3735, TE@TF001372-Jockey_Ele18-Jen_51-3735



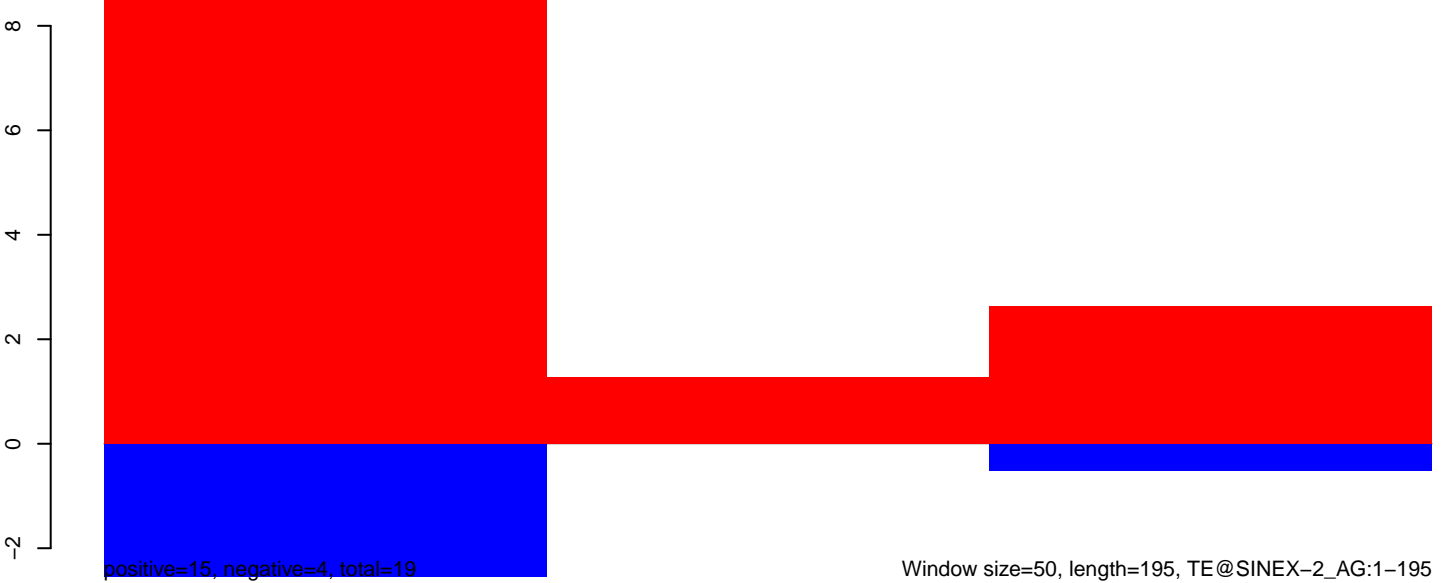
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

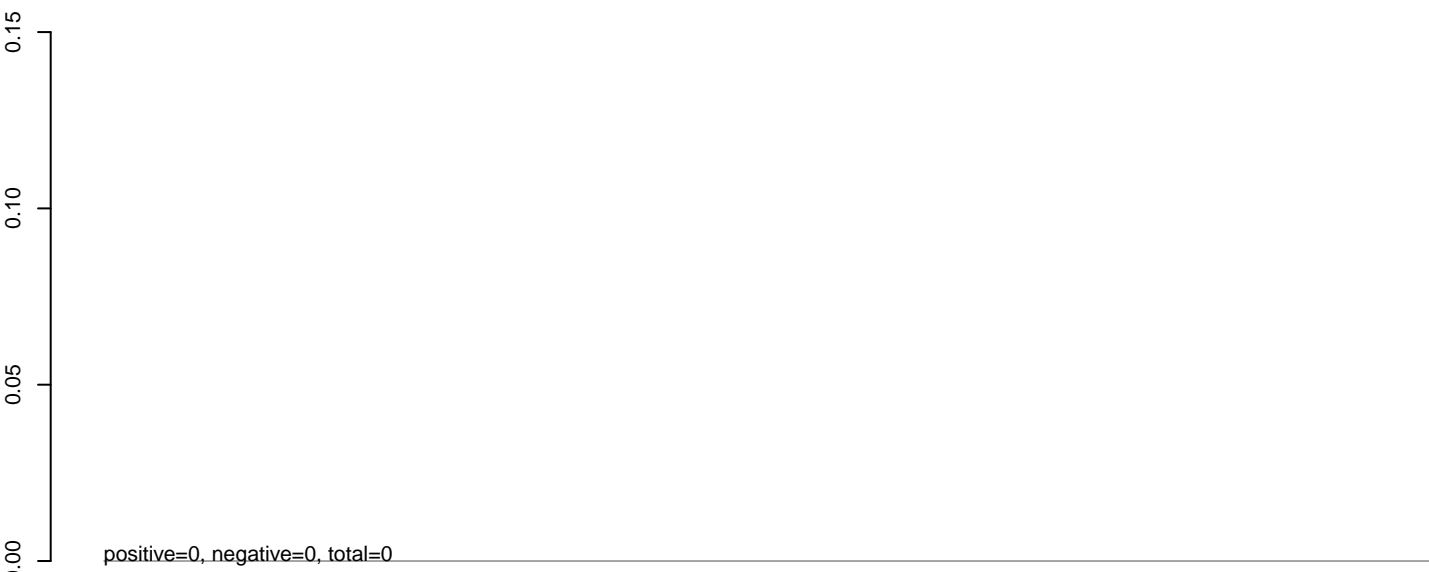


AnGam_Mos55_cells.rep

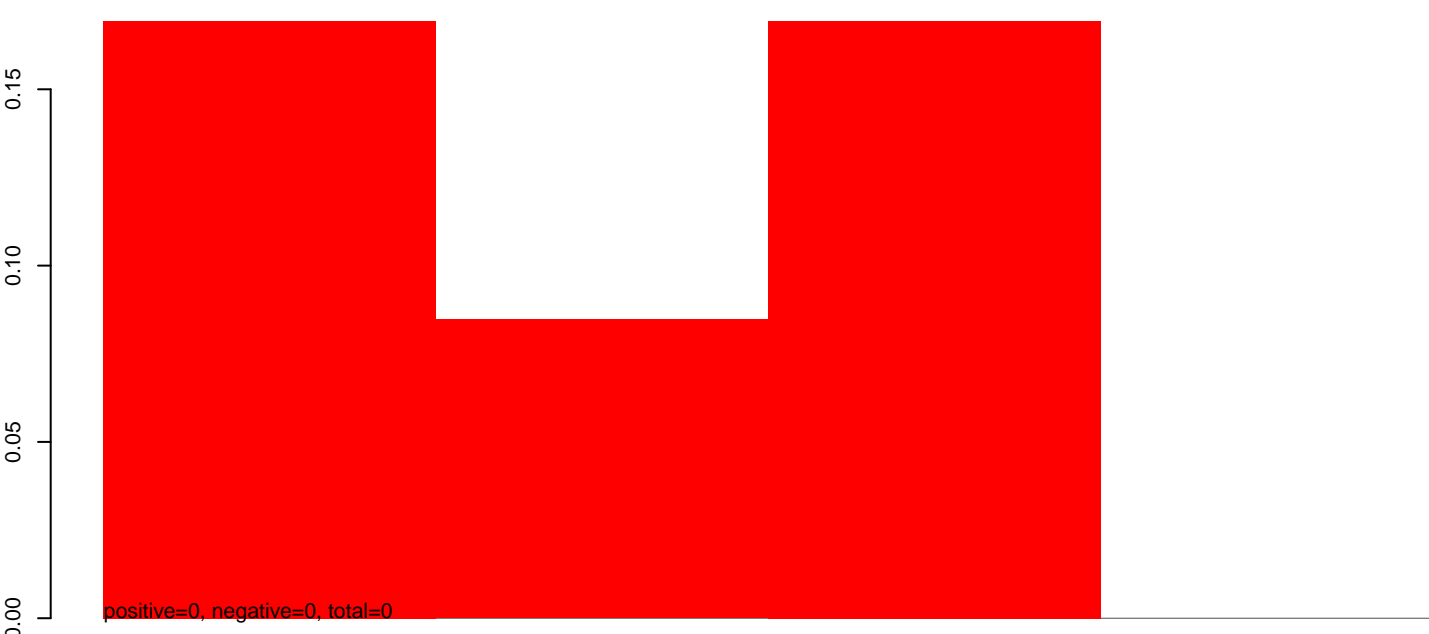


Window size=50, length=195, TE@SINEX-2_AG:1-195

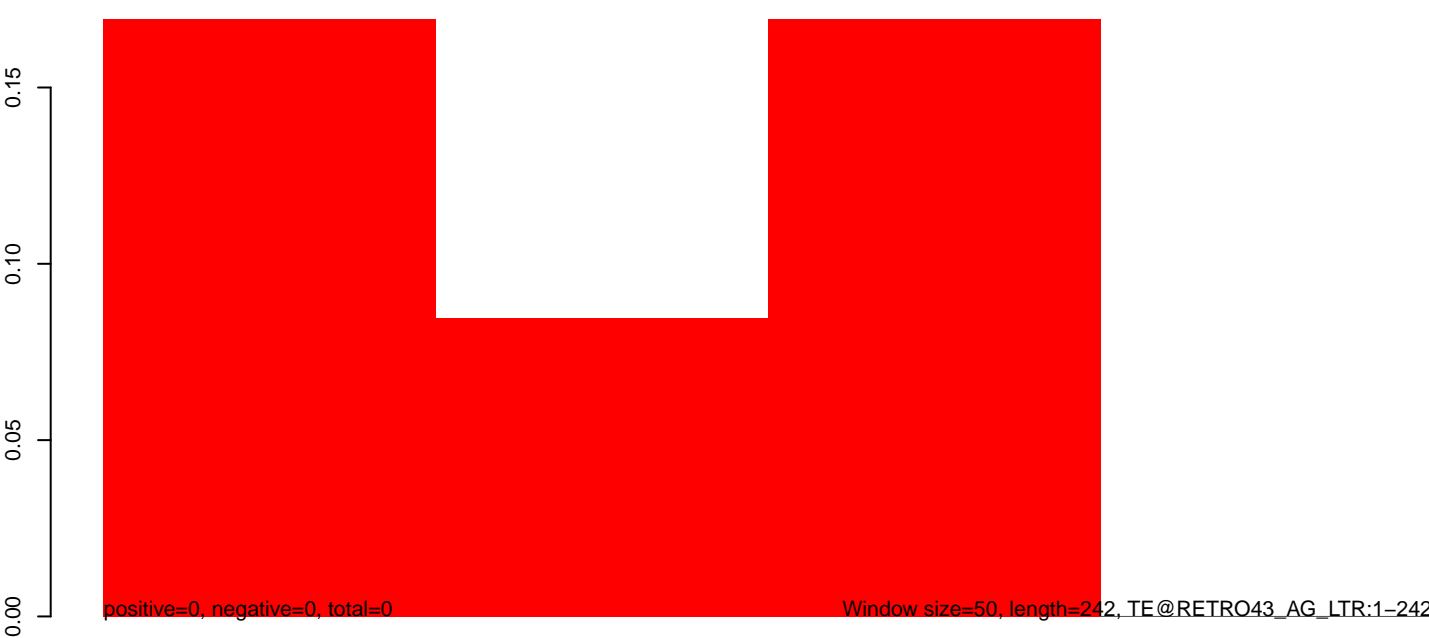
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

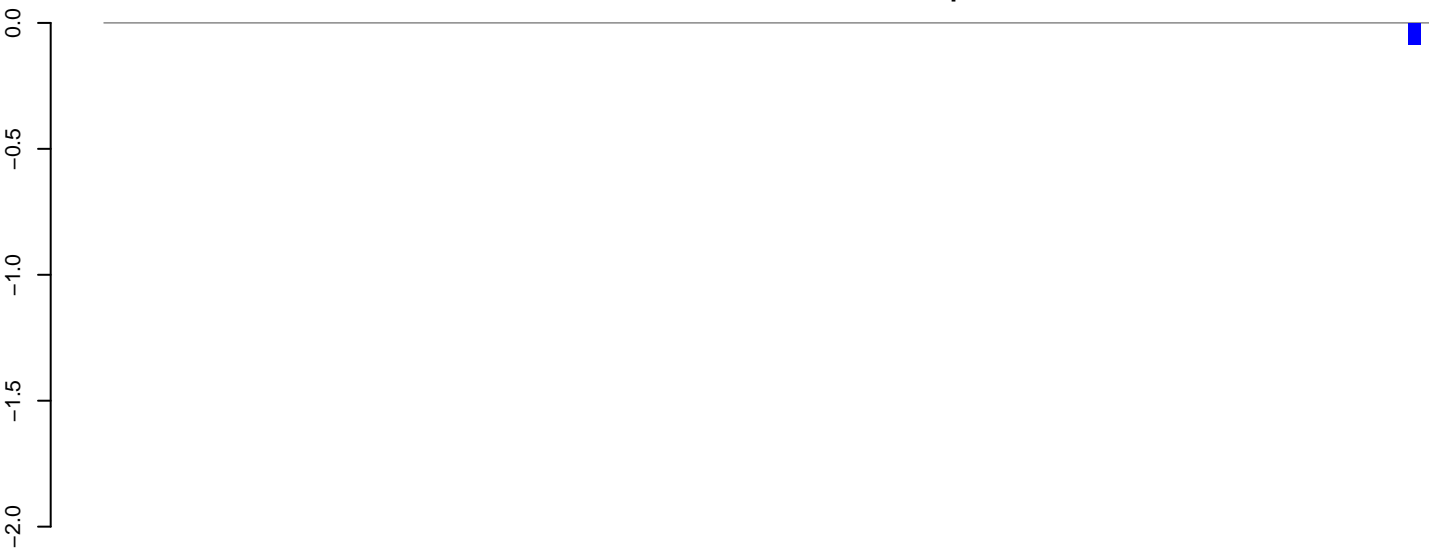


AnGam_Mos55_cells.rep



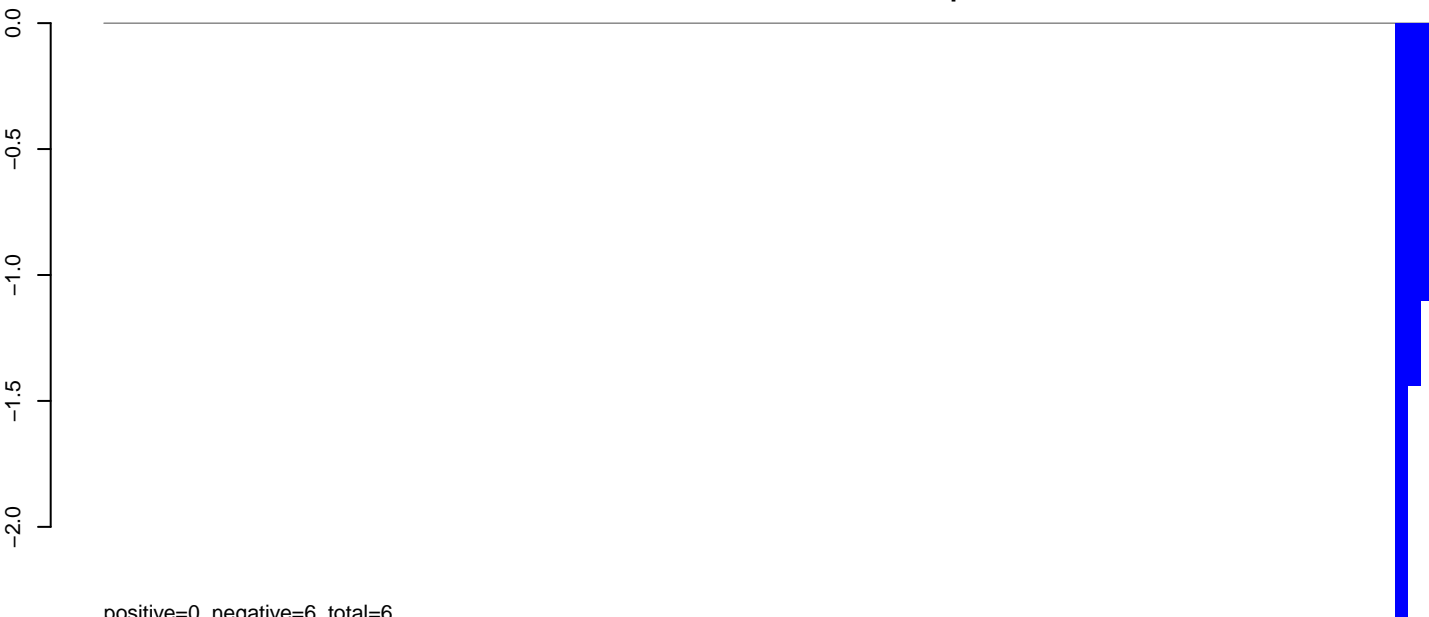
Window size=50, length=242, TE@RETRO43_AG_LTR:1-242

AnGam_Mos55_cells.18_23.rep



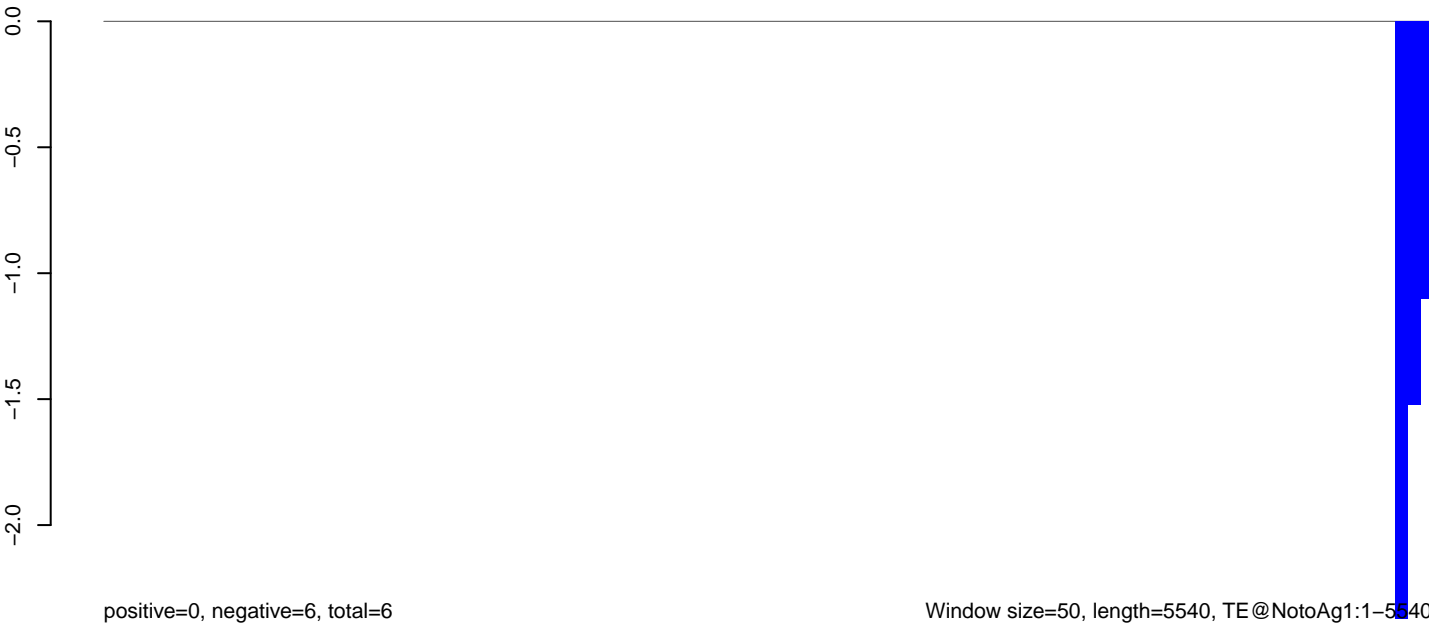
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=6, total=6

AnGam_Mos55_cells.rep

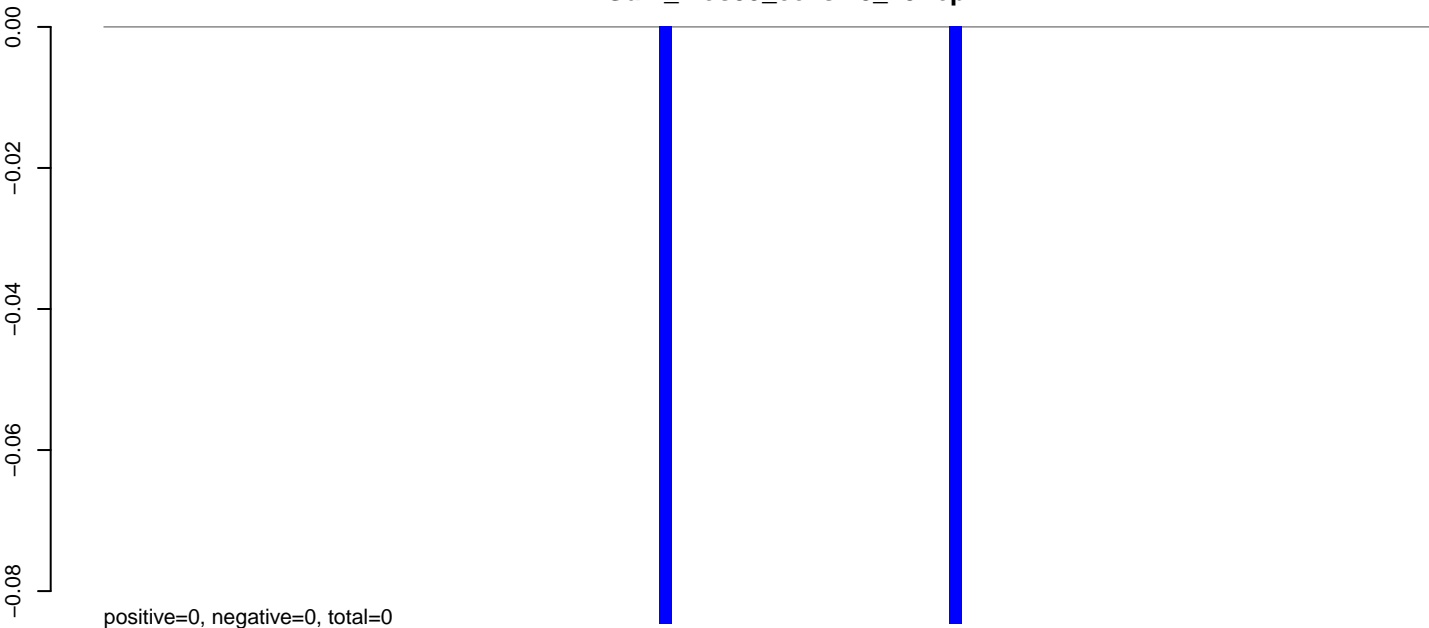


positive=0, negative=6, total=6

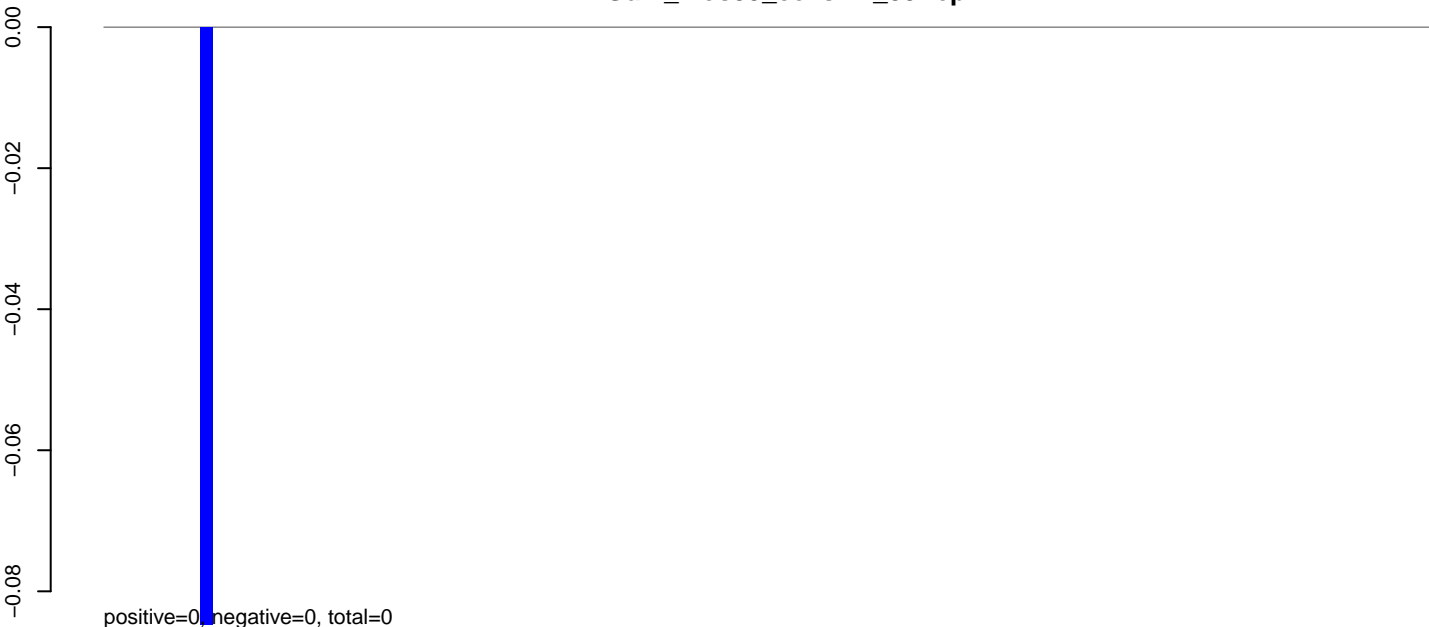
Window size=50, length=5540, TE@NotoAg1:1-5540



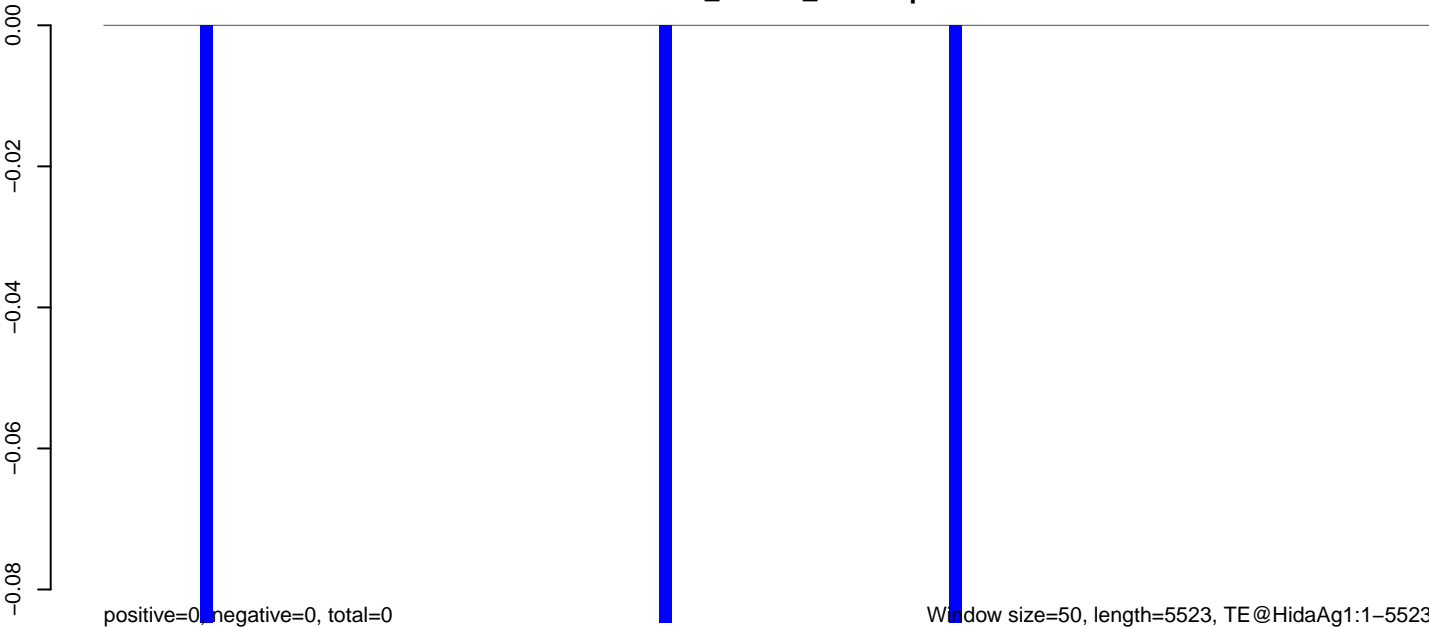
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

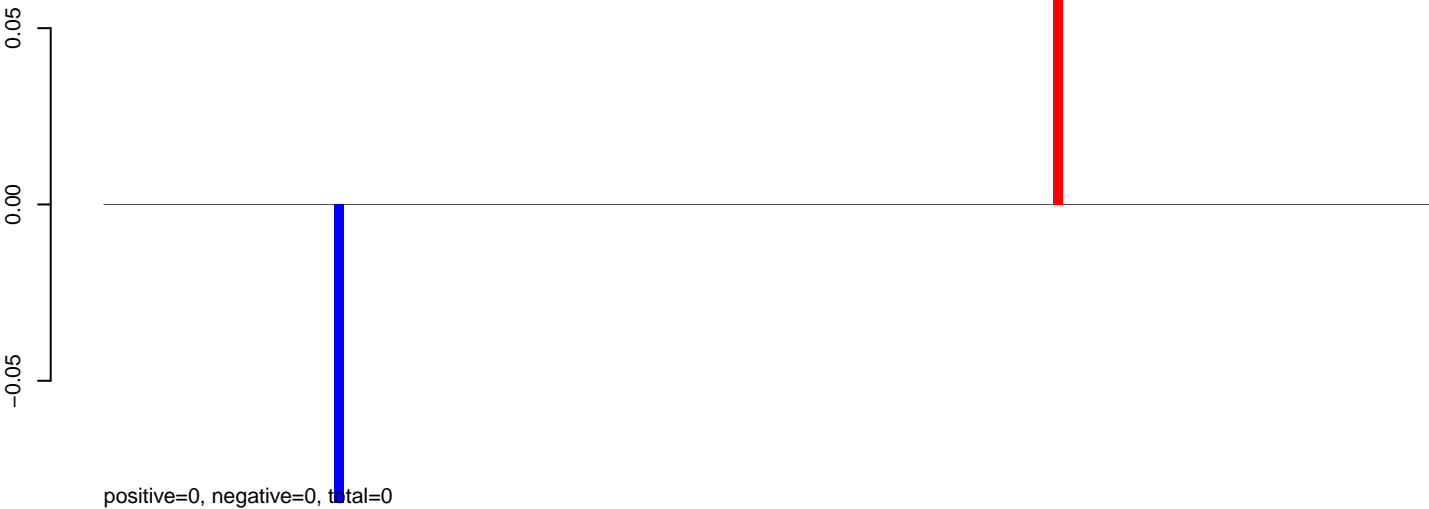


AnGam_Mos55_cells.rep

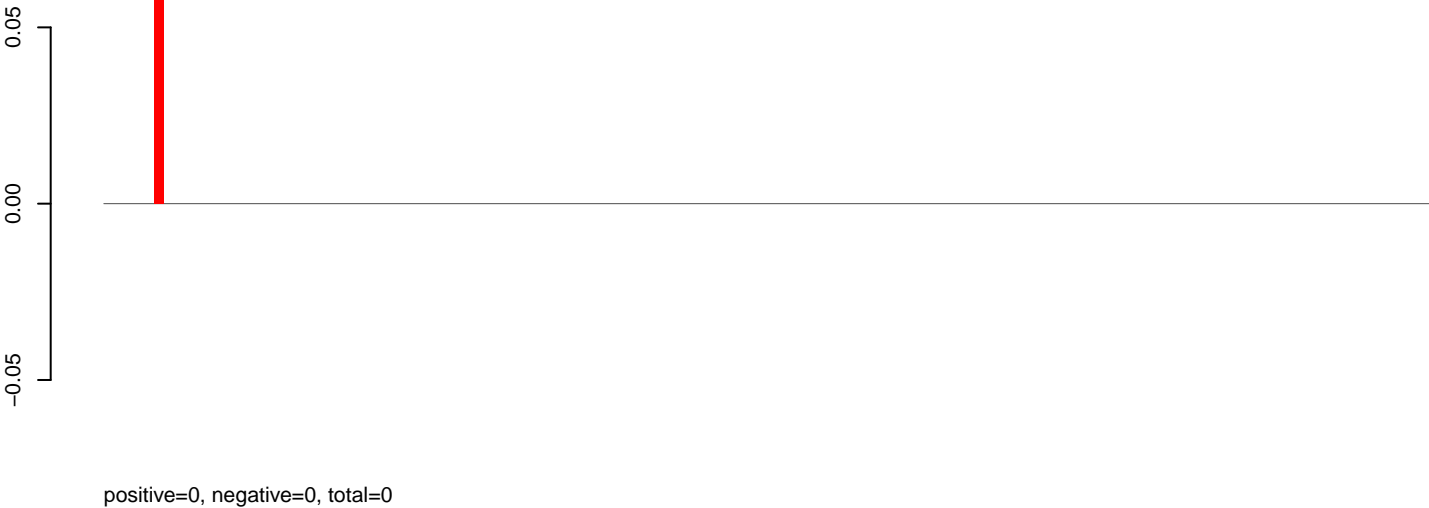


0 1000 2000 3000 4000 5000

AnGam_Mos55_cells.18_23.rep



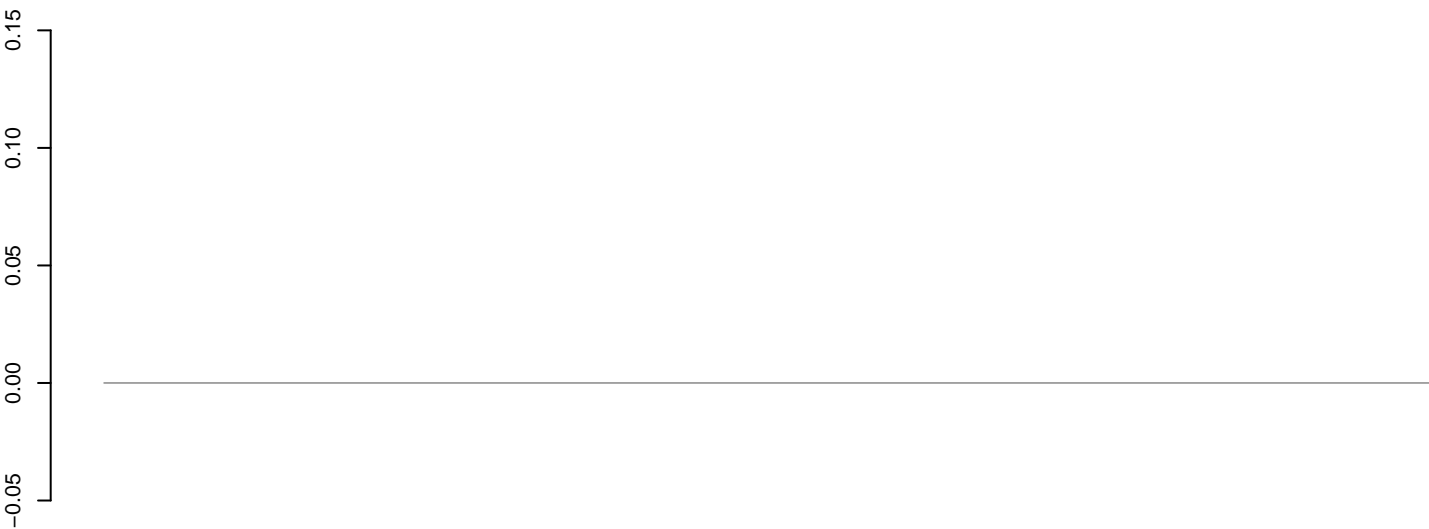
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



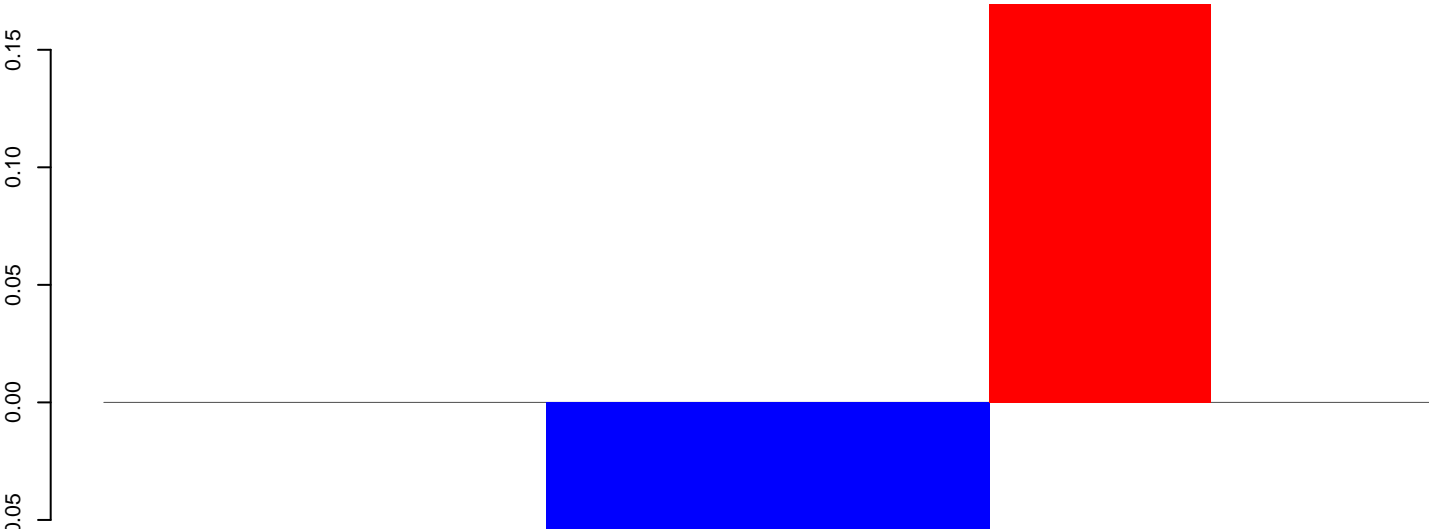
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

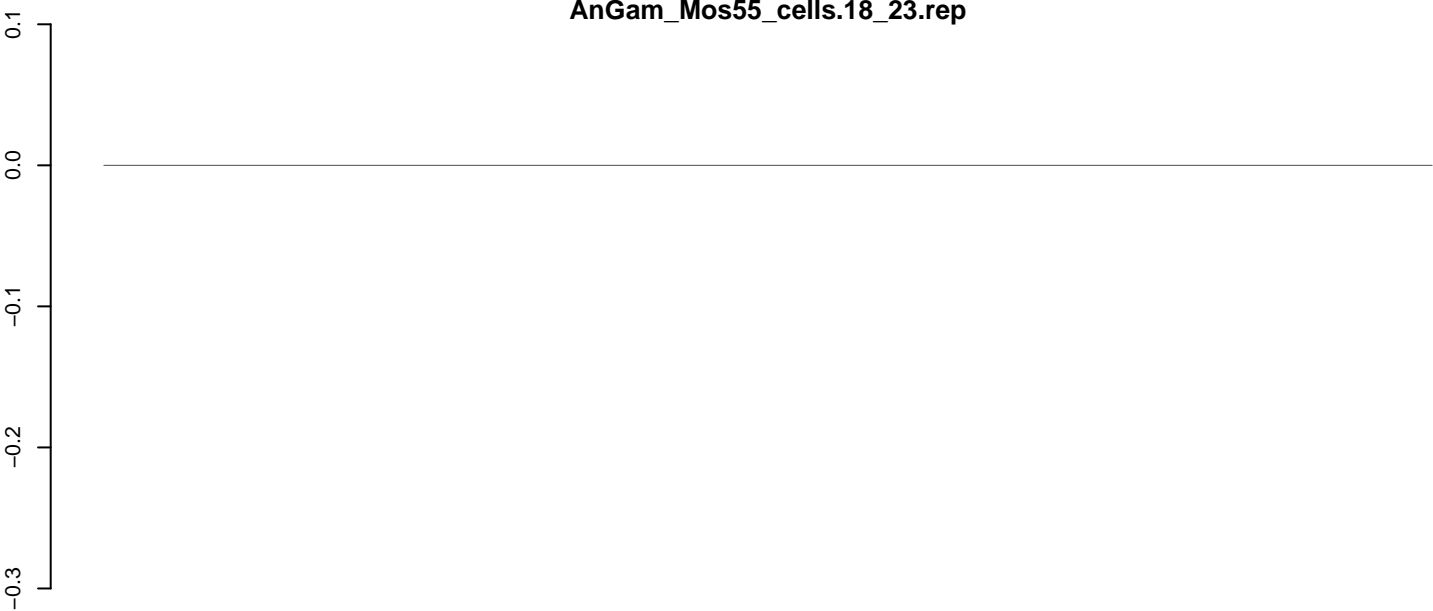


positive=0, negative=0, total=0

Window size=50, length=315, TE@HATN6_AG:1-315

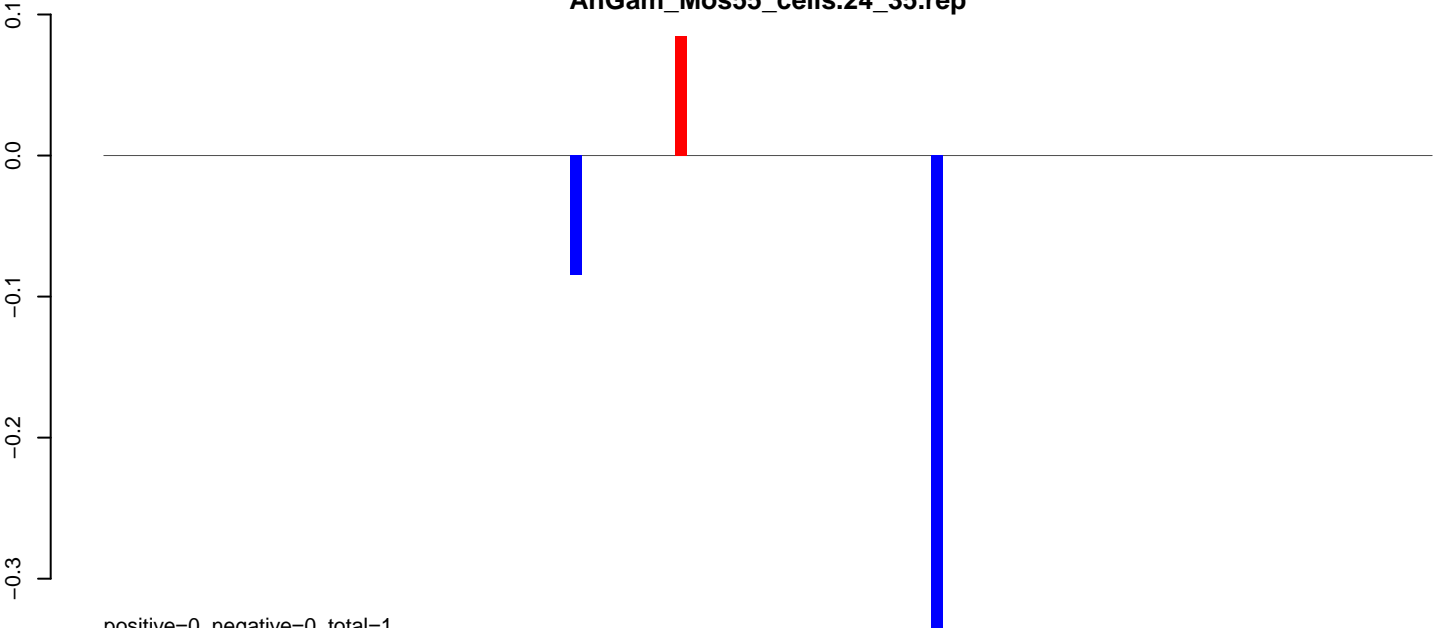
50 100 150 200 250 300 350

AnGam_Mos55_cells.18_23.rep



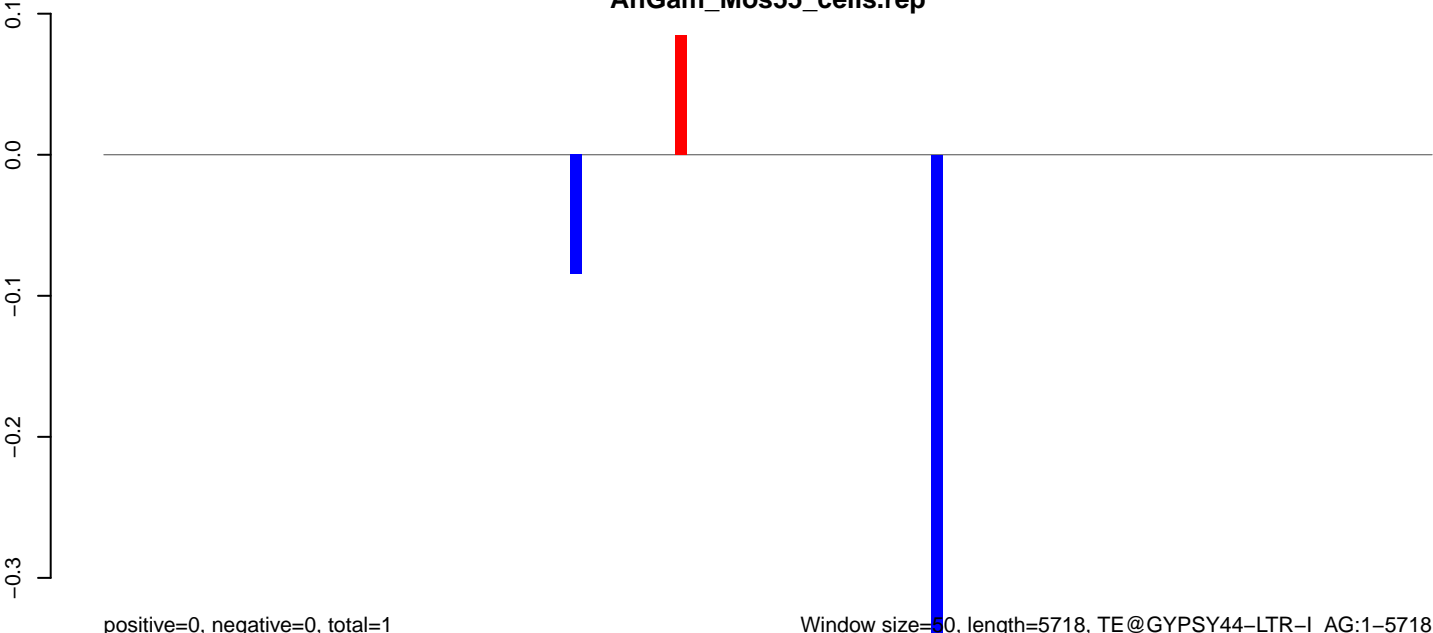
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=1

AnGam_Mos55_cells.rep

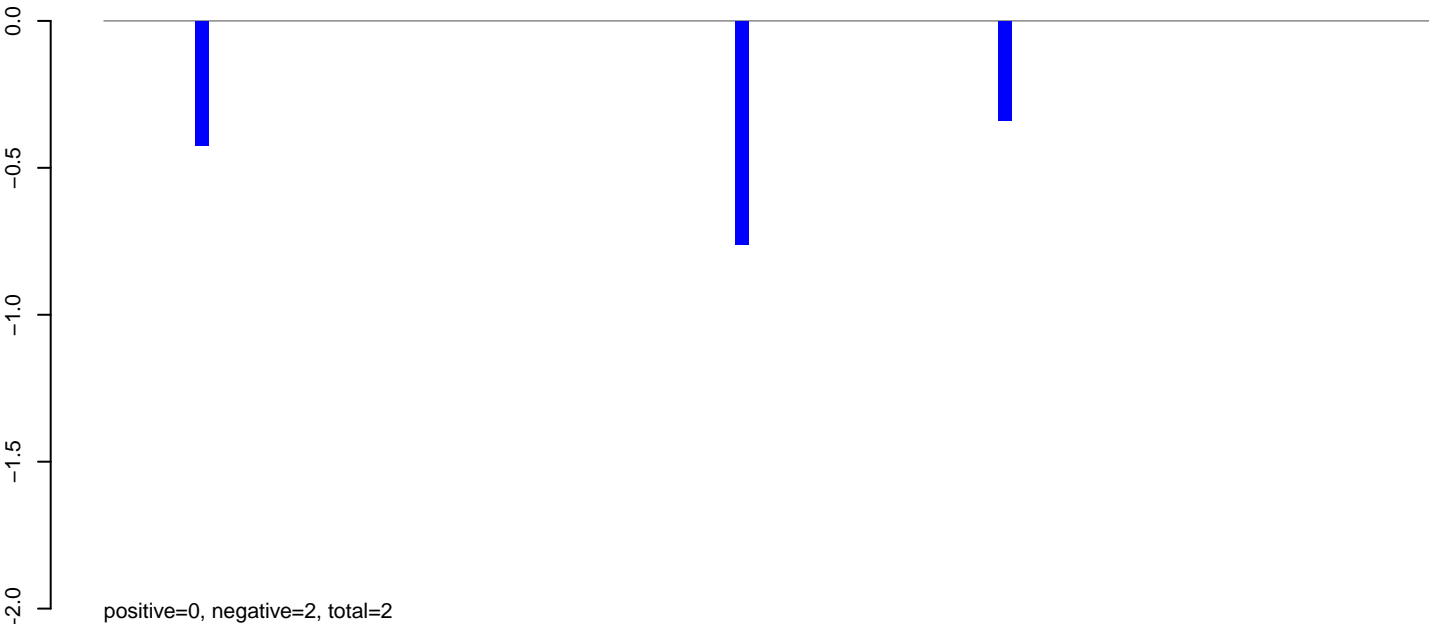


positive=0, negative=0, total=1

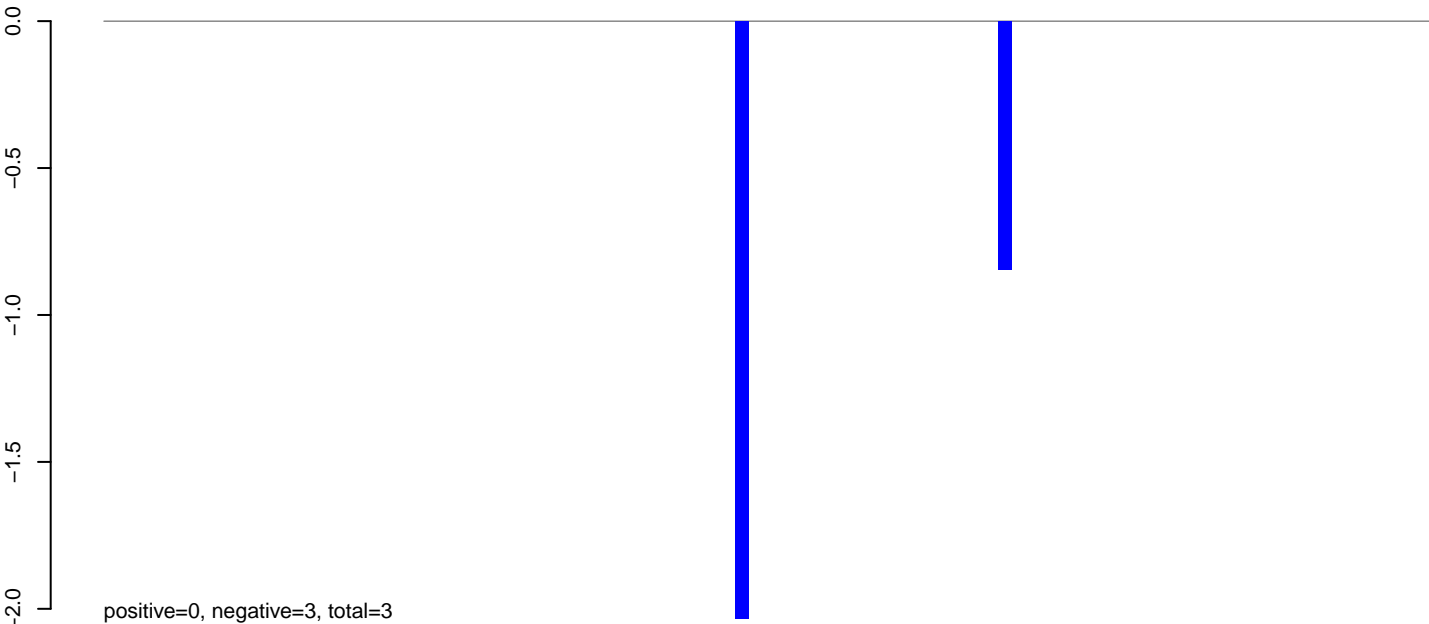
Window size=50, length=5718, TE@GYPSY44-LTR-I_AG:1-5718

0 1000 2000 3000 4000 5000

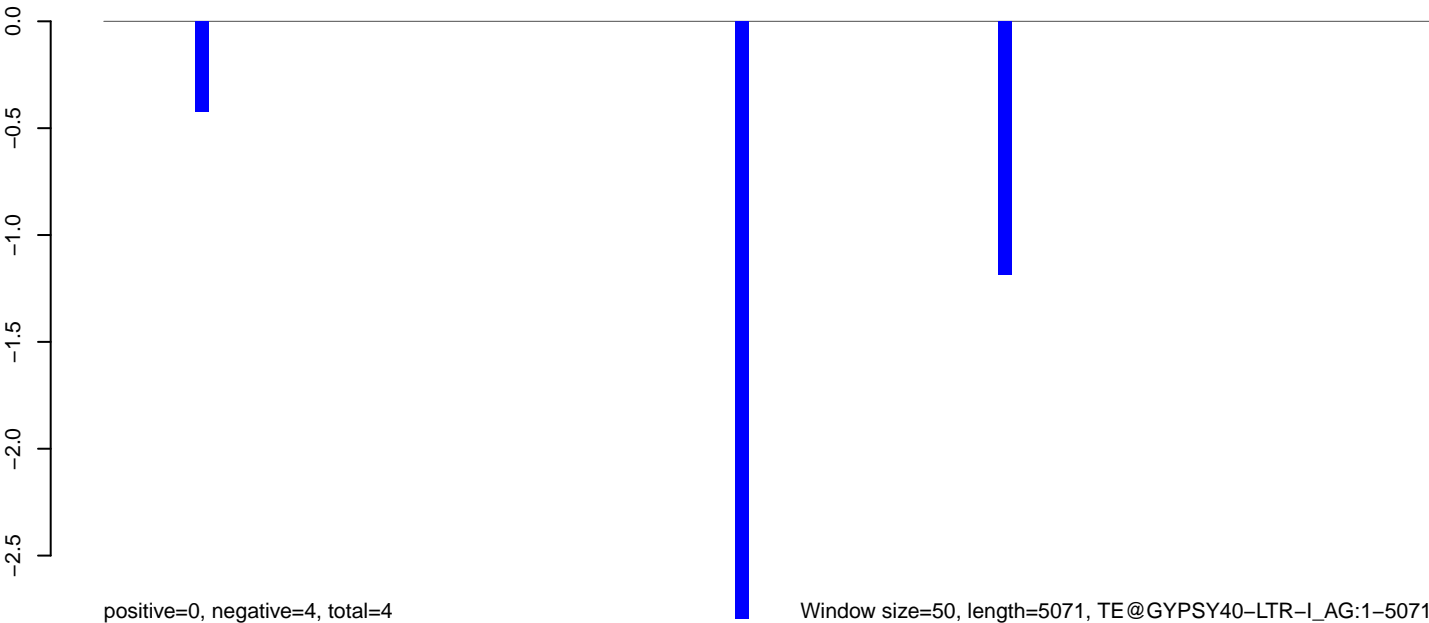
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

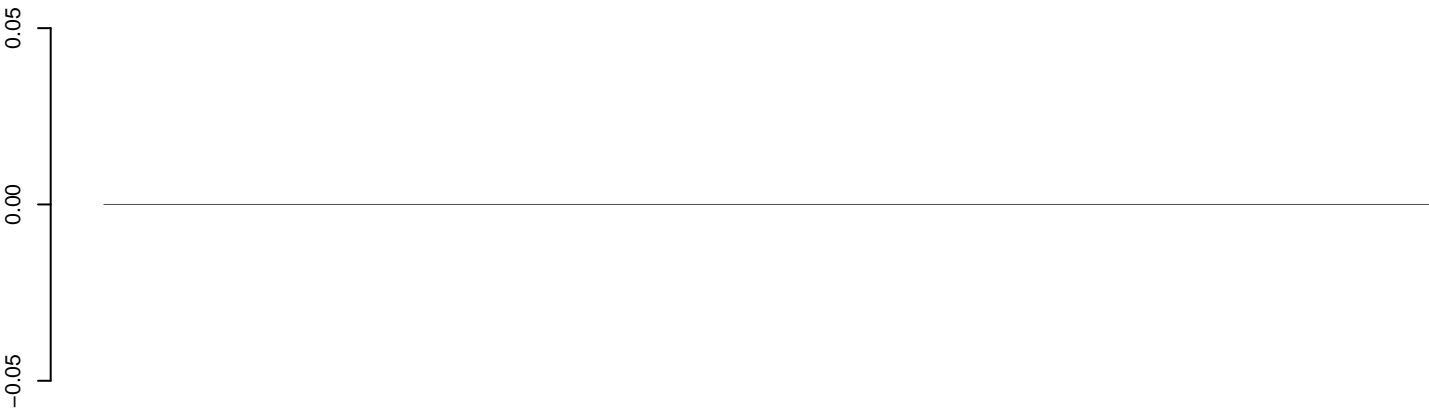


AnGam_Mos55_cells.rep



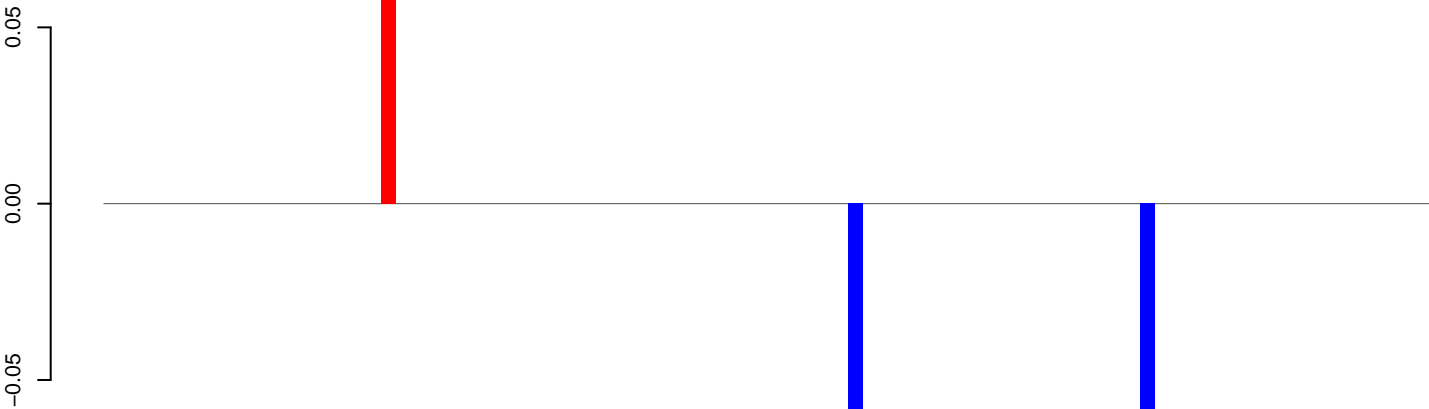
Window size=50, length=5071, TE@GYPSY40-LTR-I_AG:1-5071

AnGam_Mos55_cells.18_23.rep



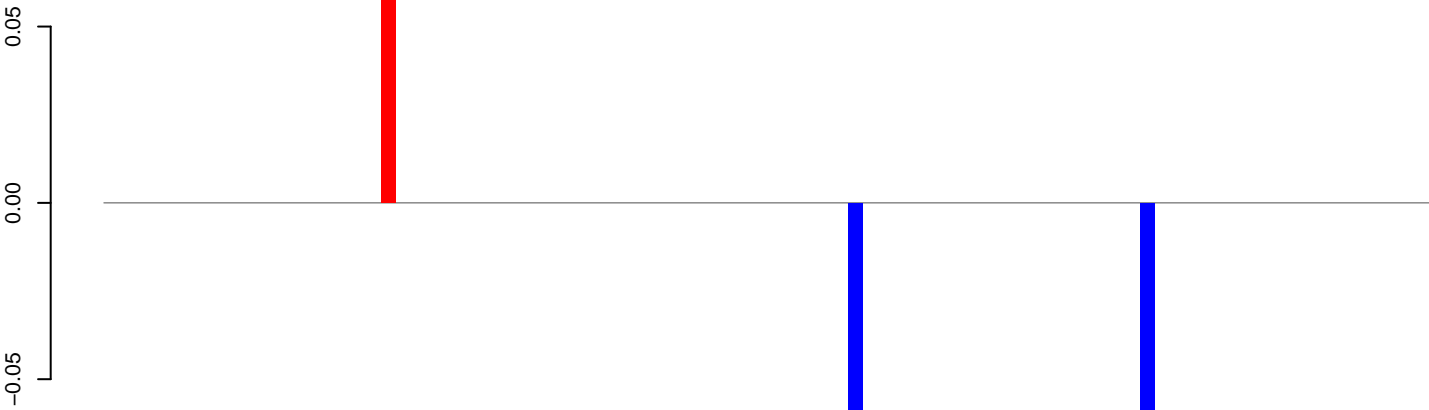
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



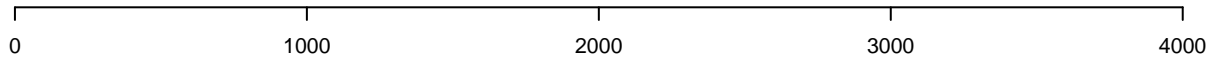
positive=0, negative=0, total=0

AnGam_Mos55_cells.rep



positive=0, negative=0, total=0

Window size=50, length=4555, TE@GYPSY22-LTR-I_AG:1-4555



AnGam_Mos55_cells.18_23.rep

0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

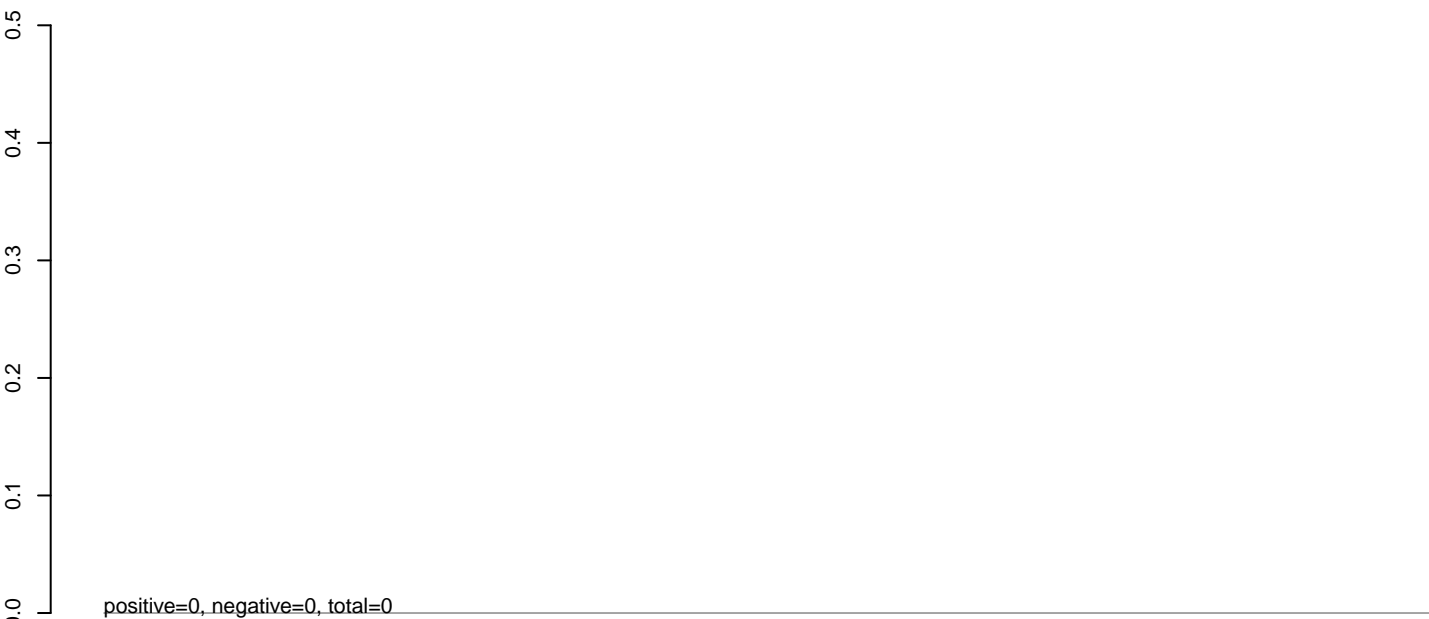
0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

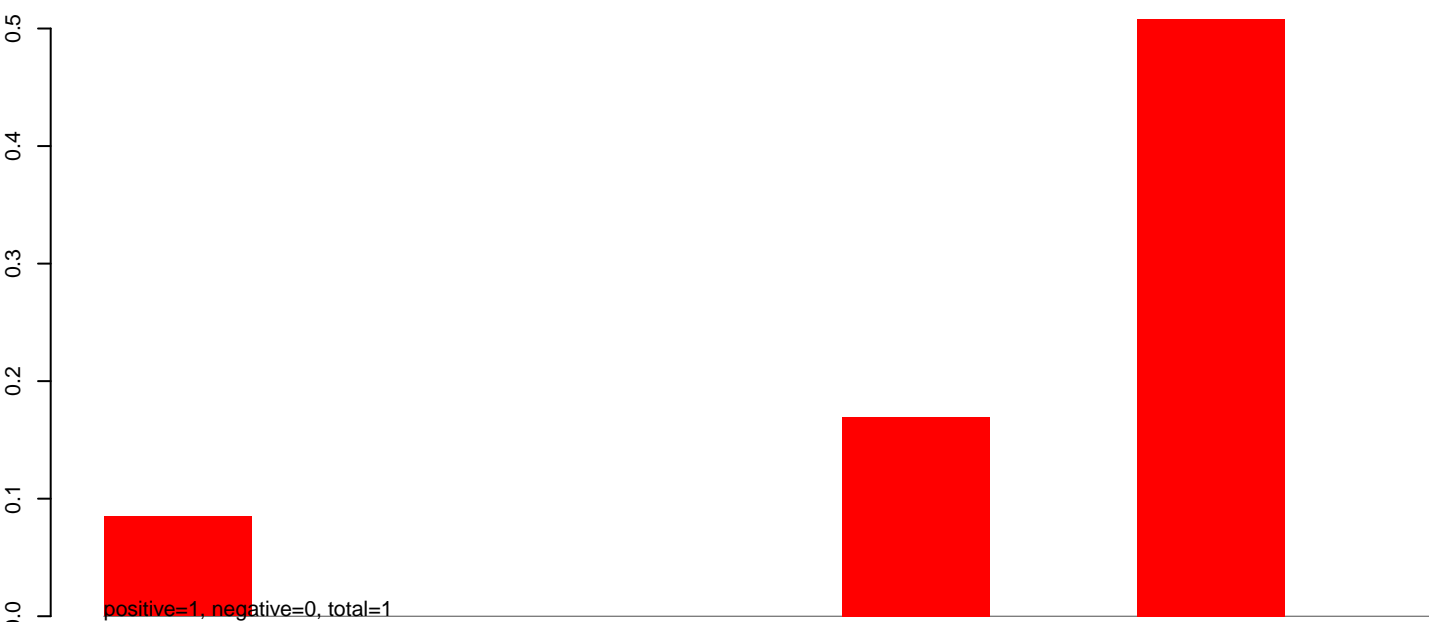
Window size=50, length=1113, TE@EAA44450-Unknown:1-1113

200 400 600 800 1000

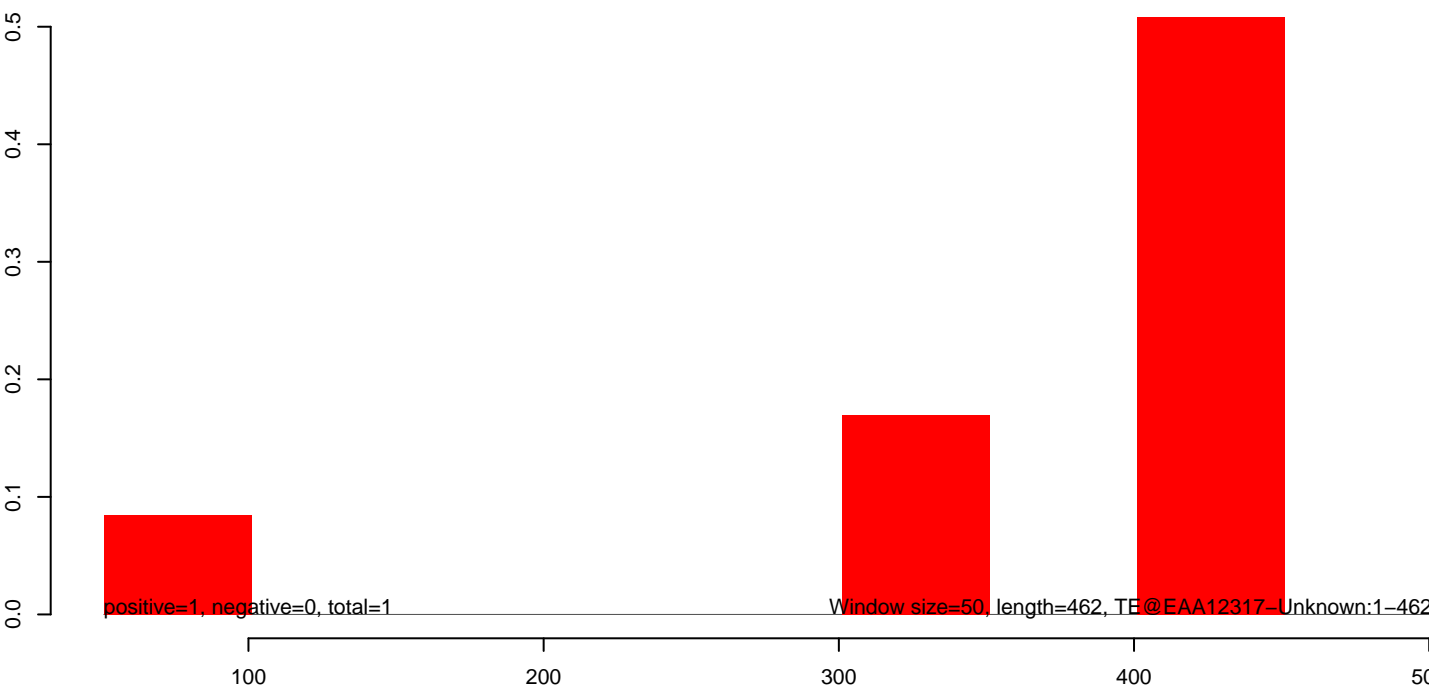
AnGam_Mos55_cells.18_23.rep



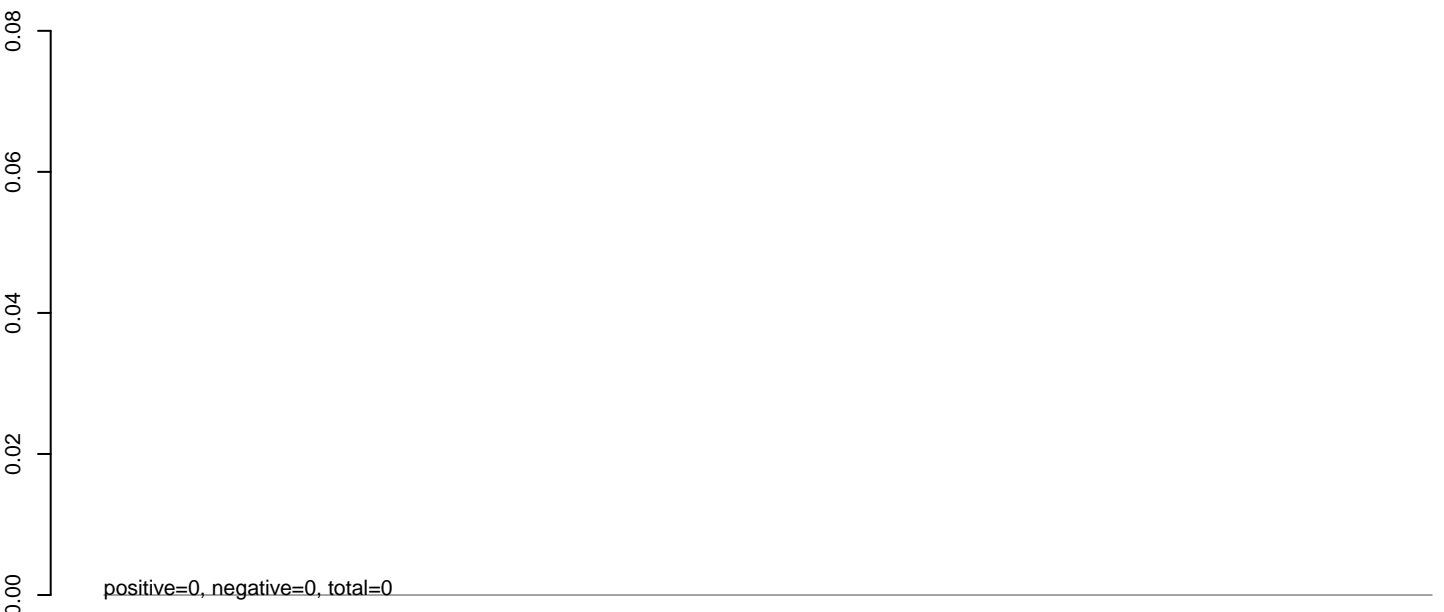
AnGam_Mos55_cells.24_35.rep



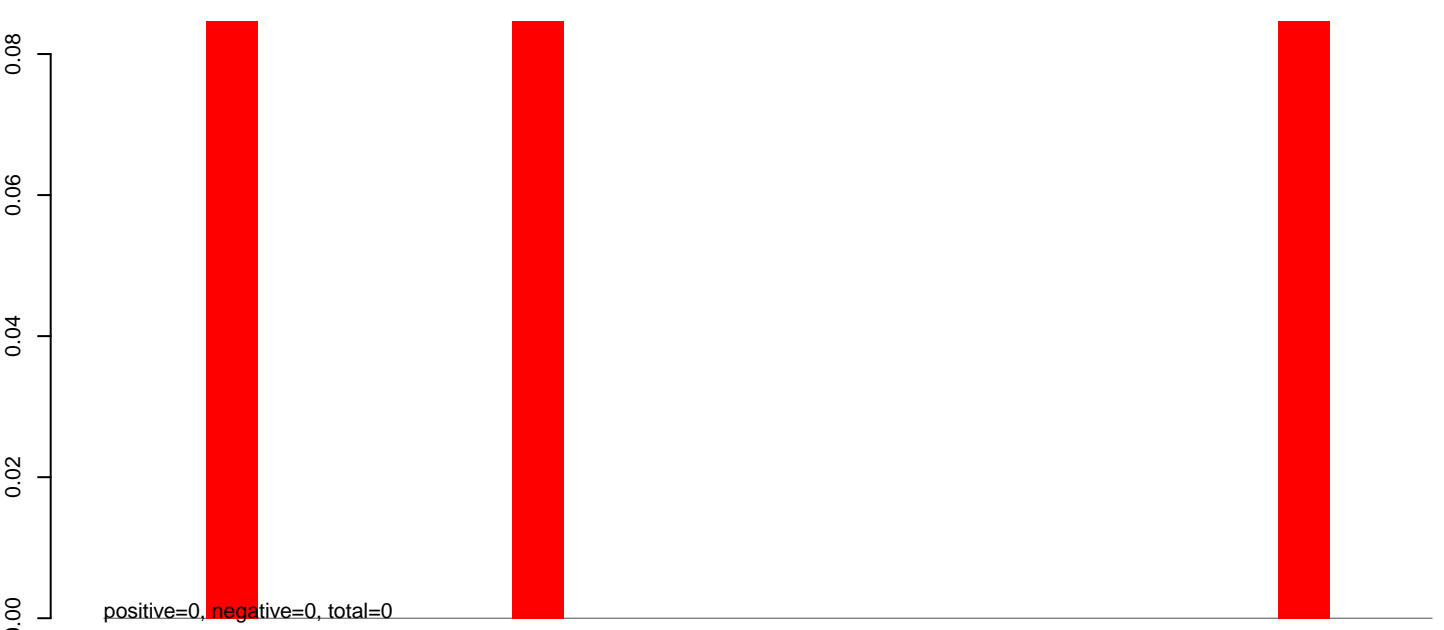
AnGam_Mos55_cells.rep



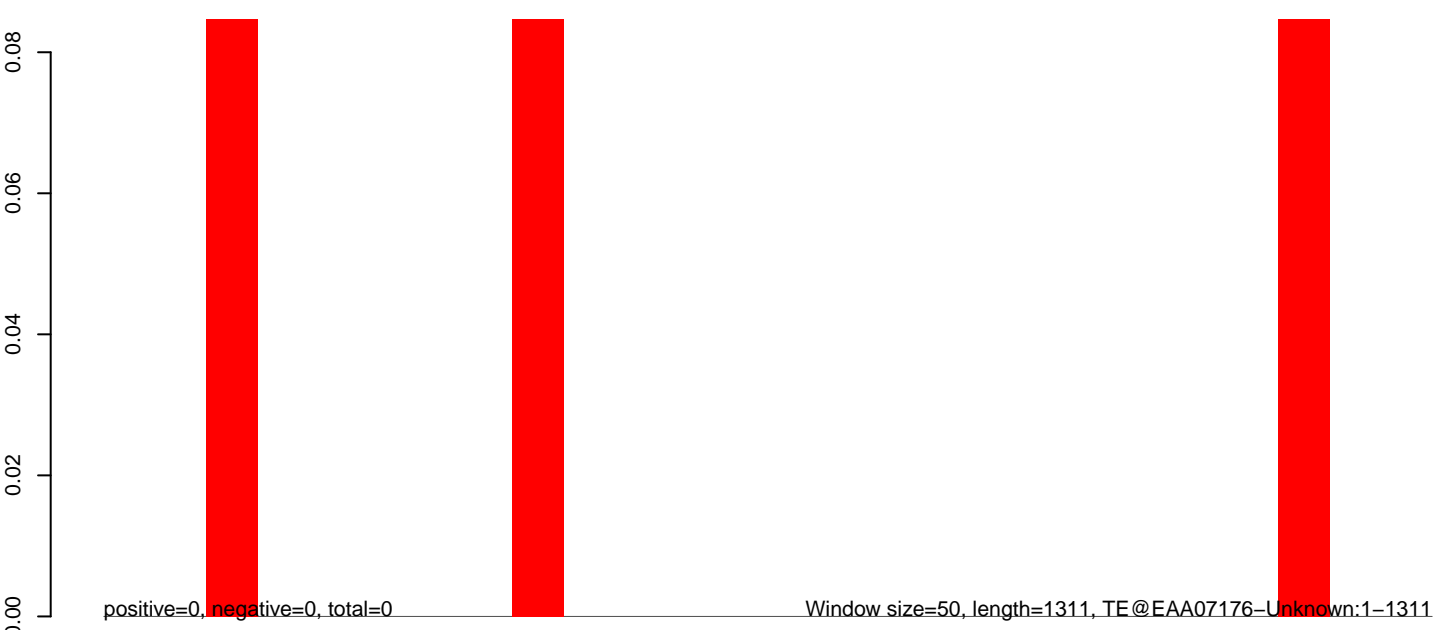
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

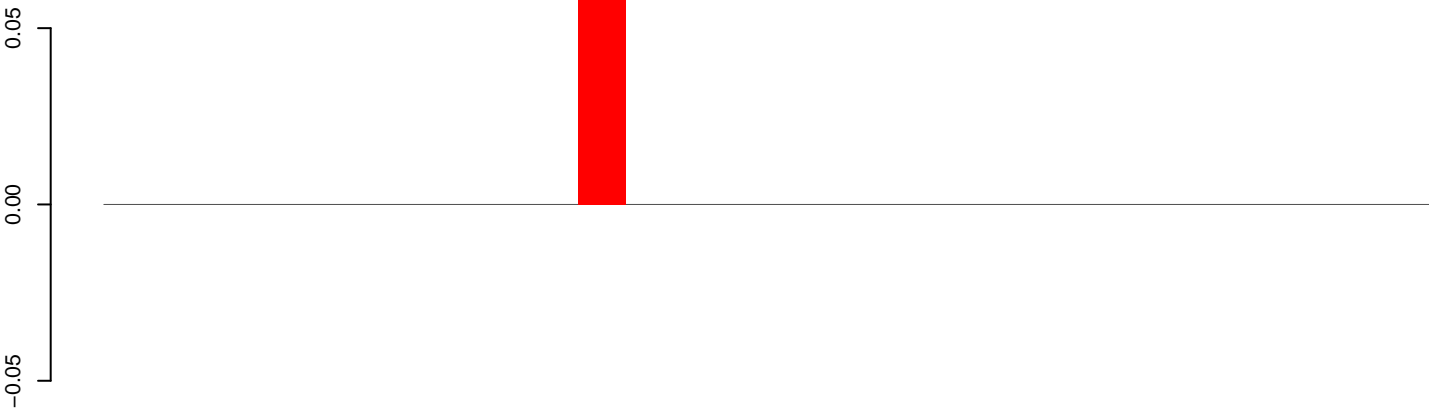


AnGam_Mos55_cells.rep



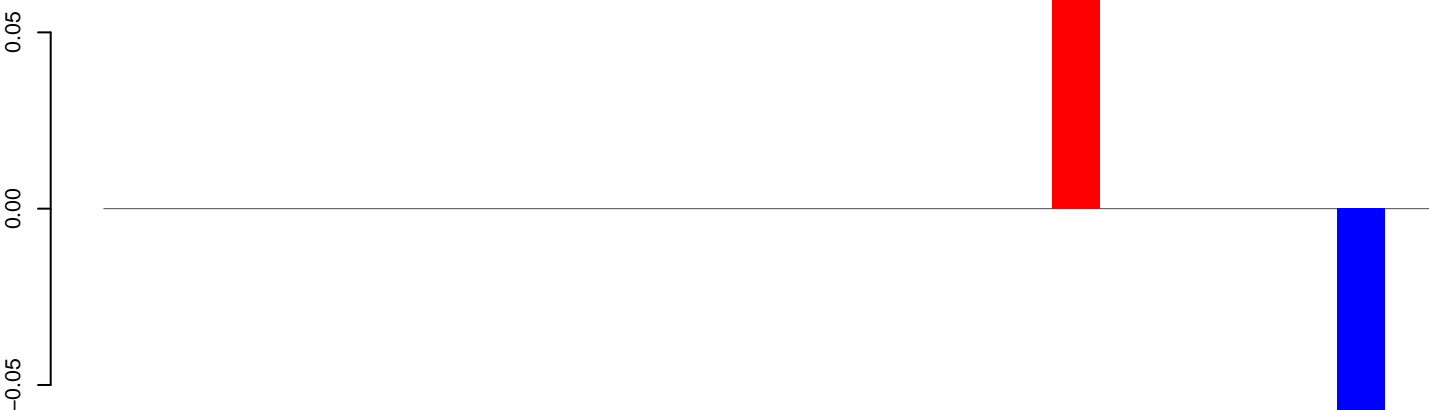
0 200 400 600 800 1000 1200 1400

AnGam_Mos55_cells.18_23.rep



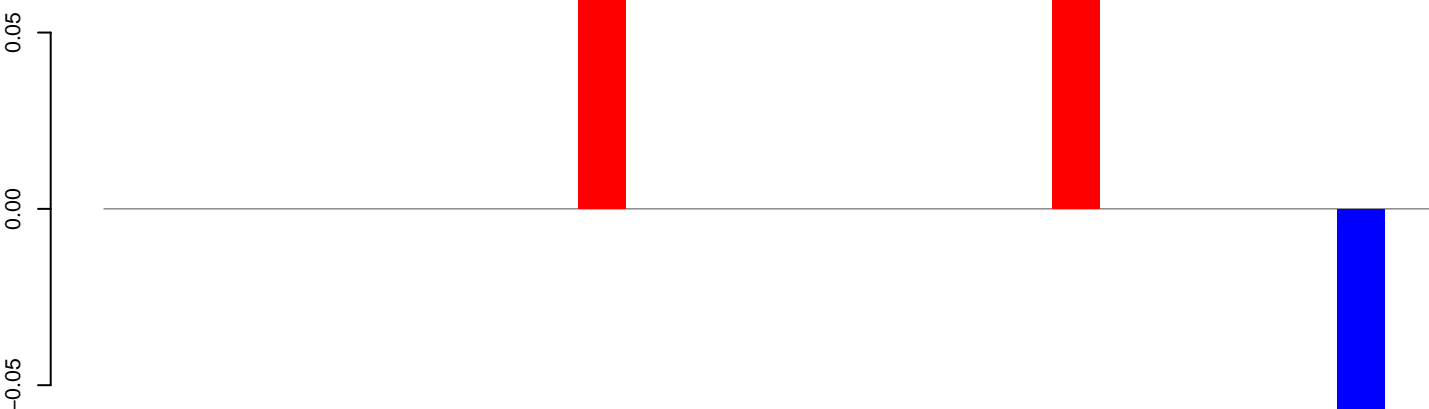
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

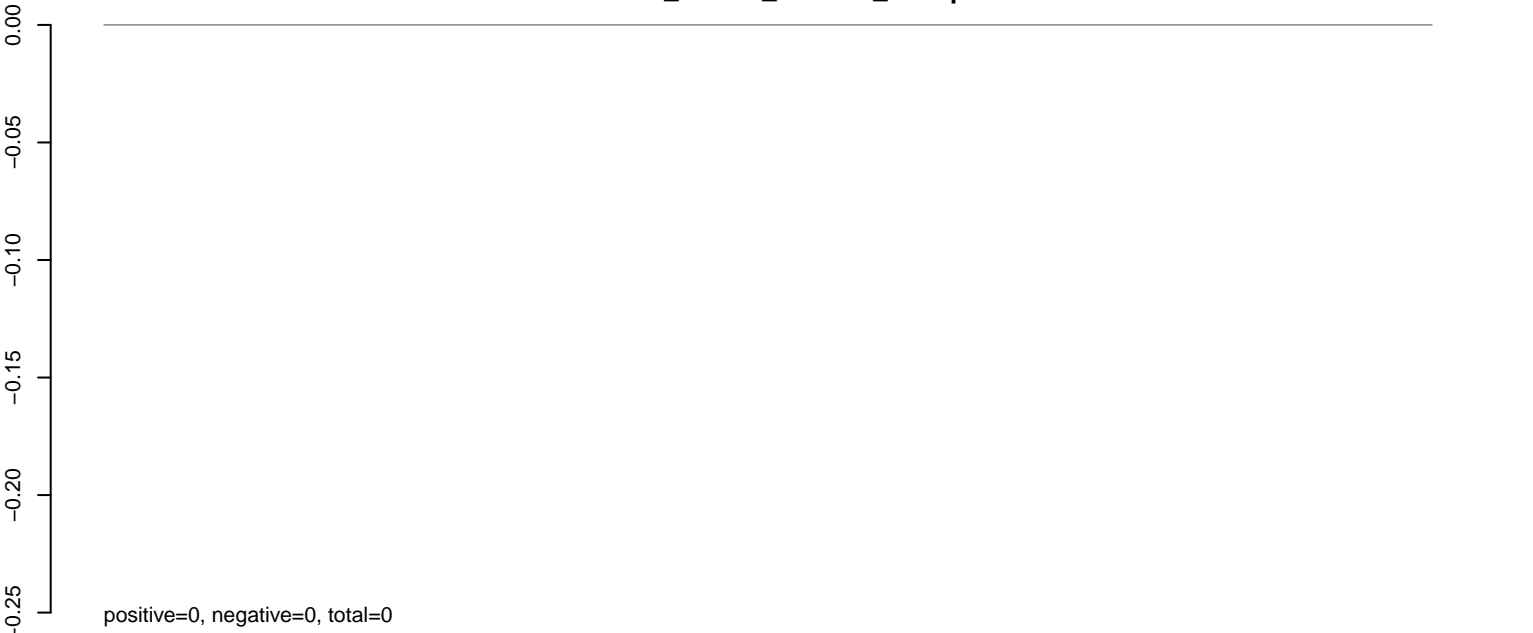


positive=0, negative=0, total=0

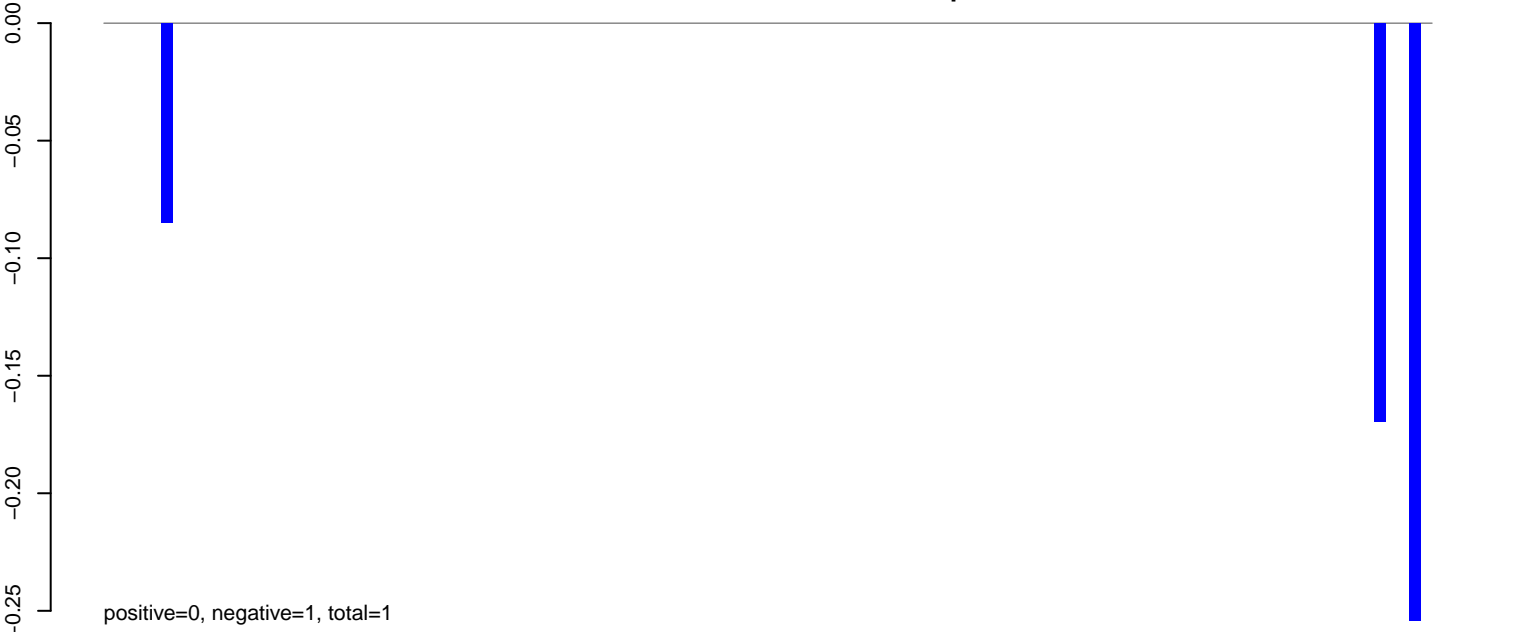
Window size=50, length=1412, TE@Clu-168_AG:1-1412

0 200 400 600 800 1000 1200 1400

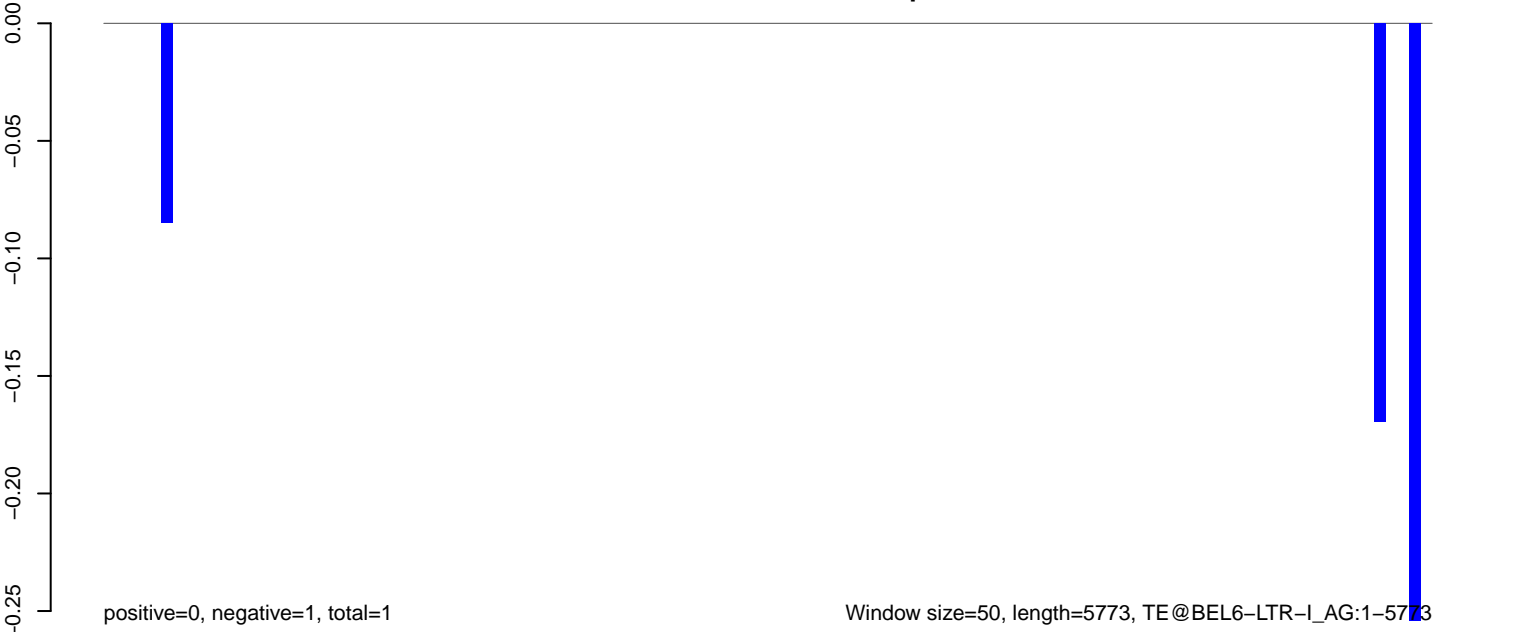
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

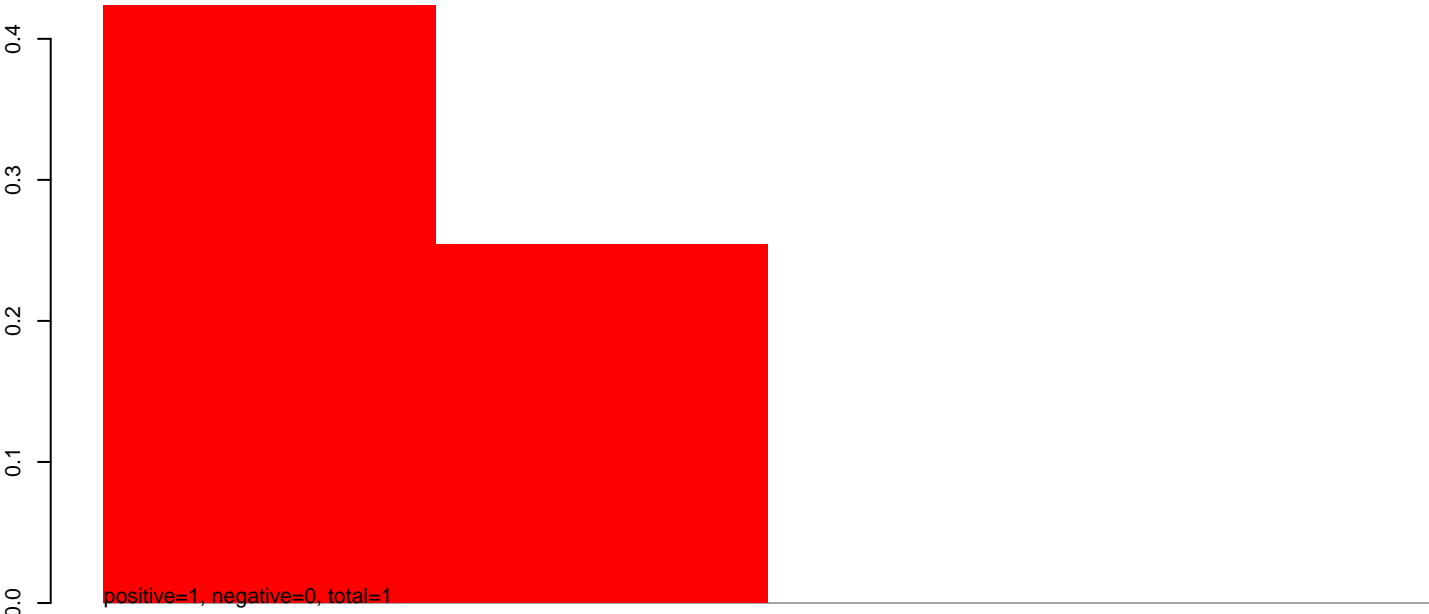


AnGam_Mos55_cells.rep

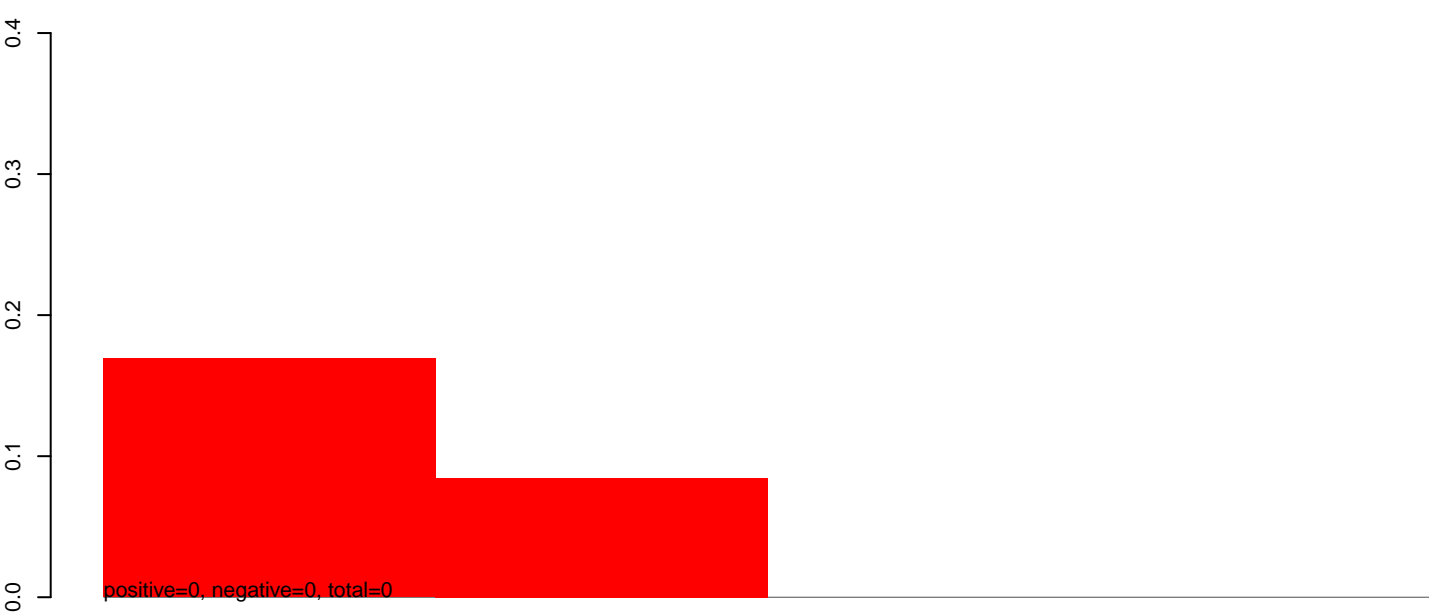


0 1000 2000 3000 4000 5000 6000

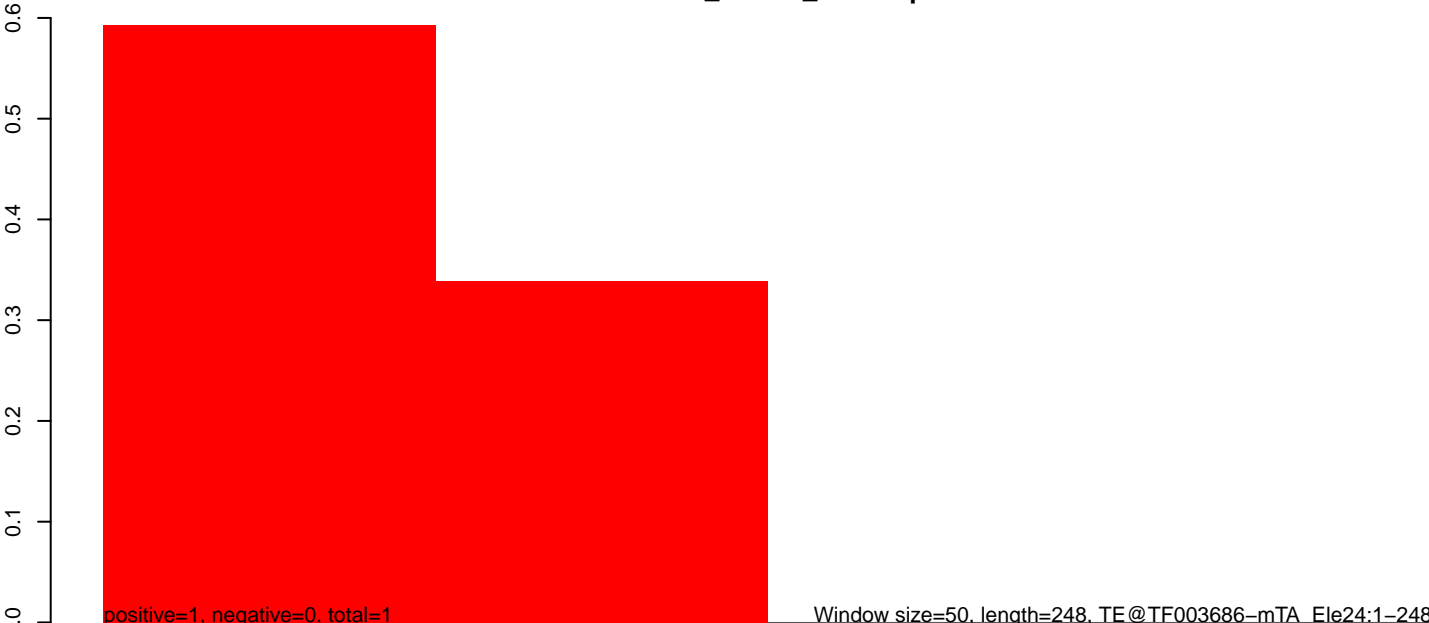
AnGam_Mos55_cells.18_23.rep



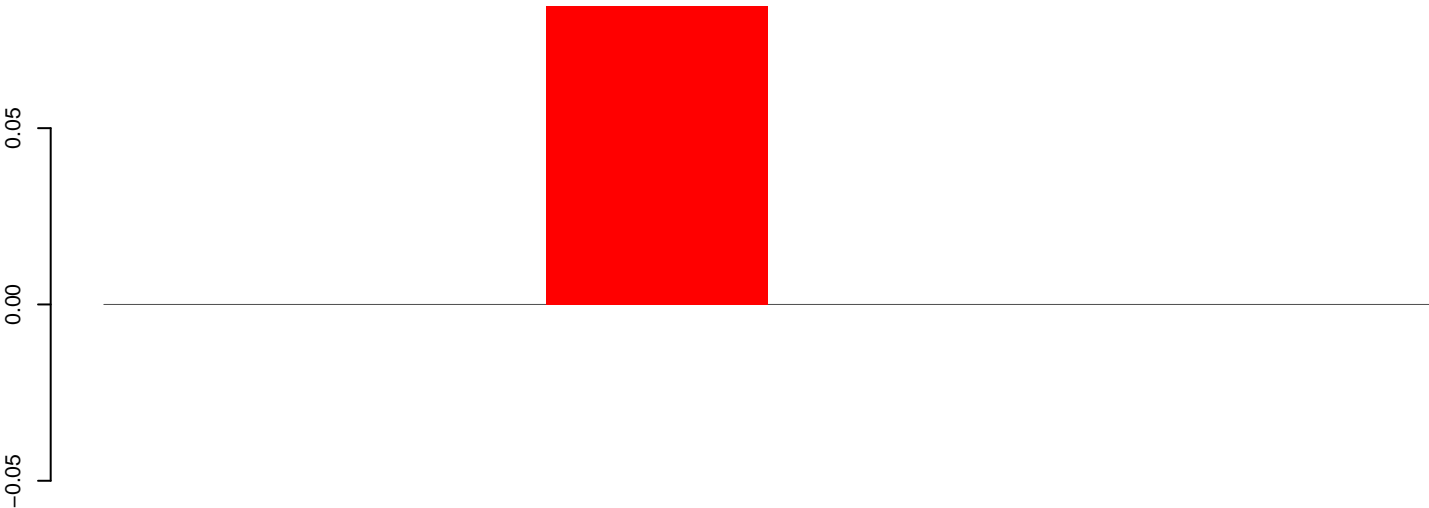
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

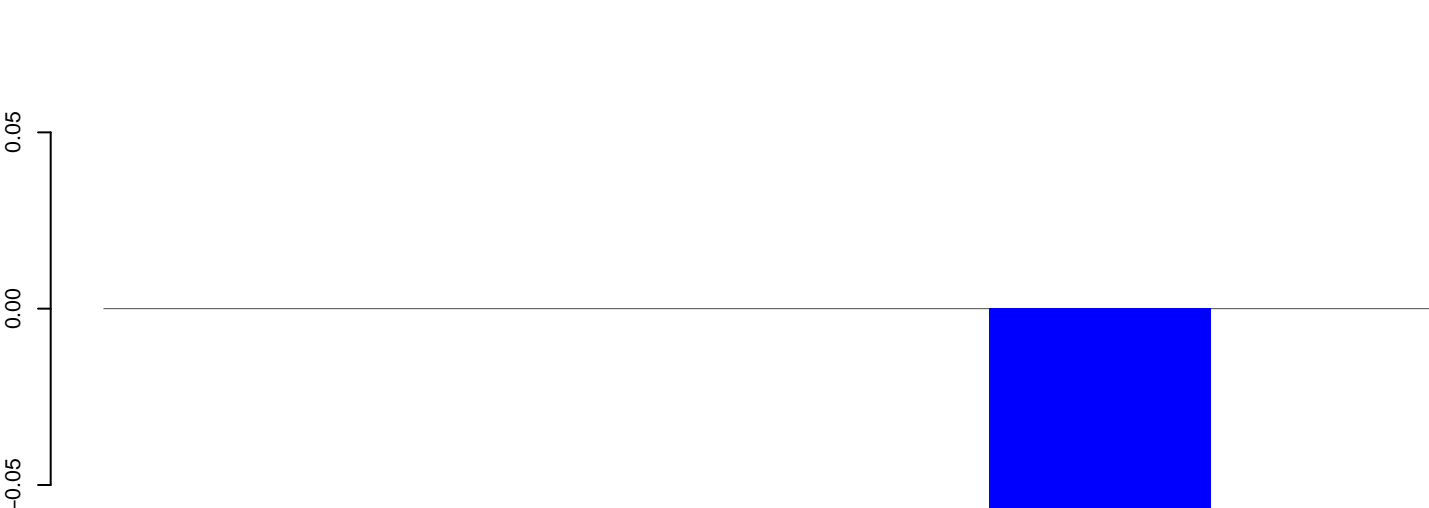


AnGam_Mos55_cells.18_23.rep



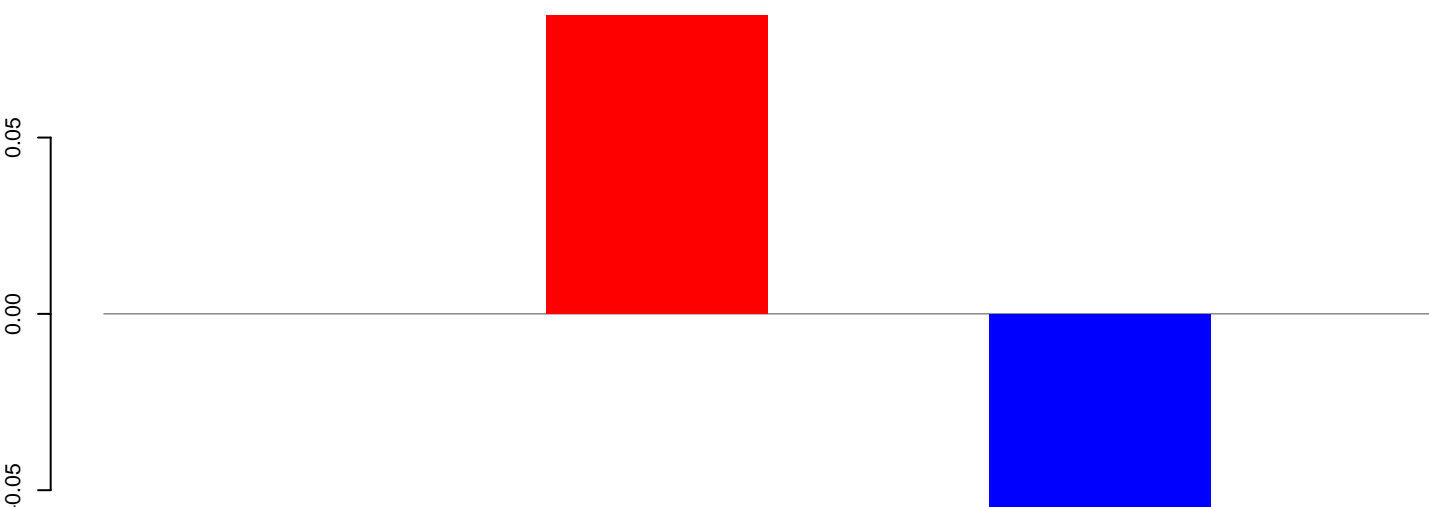
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

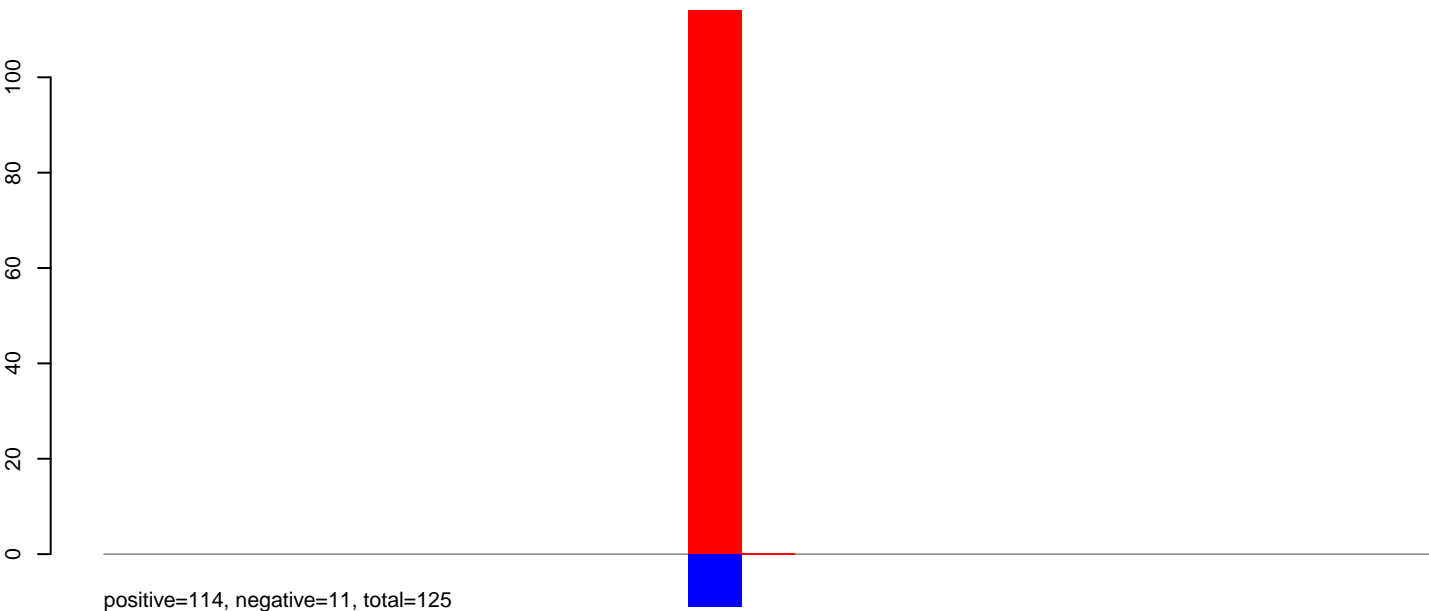


positive=0, negative=0, total=0

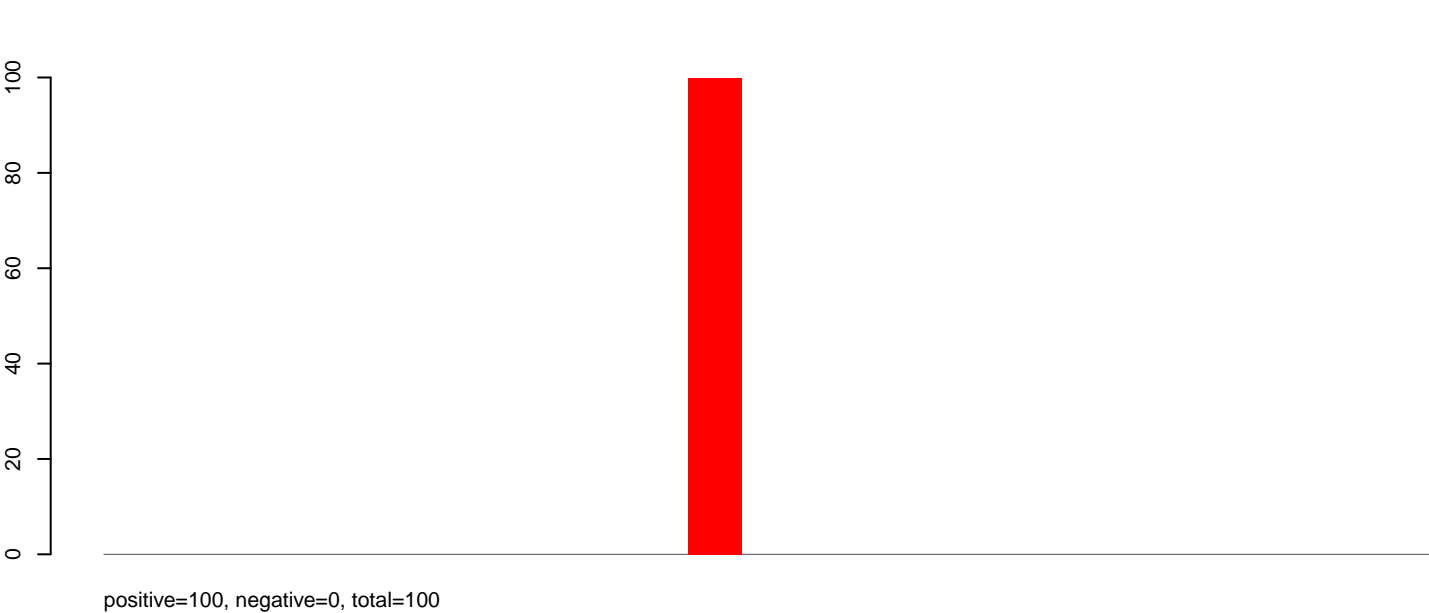
Window size=50, length=326, TE@TF003666-m8bp_Ele5:1-326

50 100 150 200 250 300 350

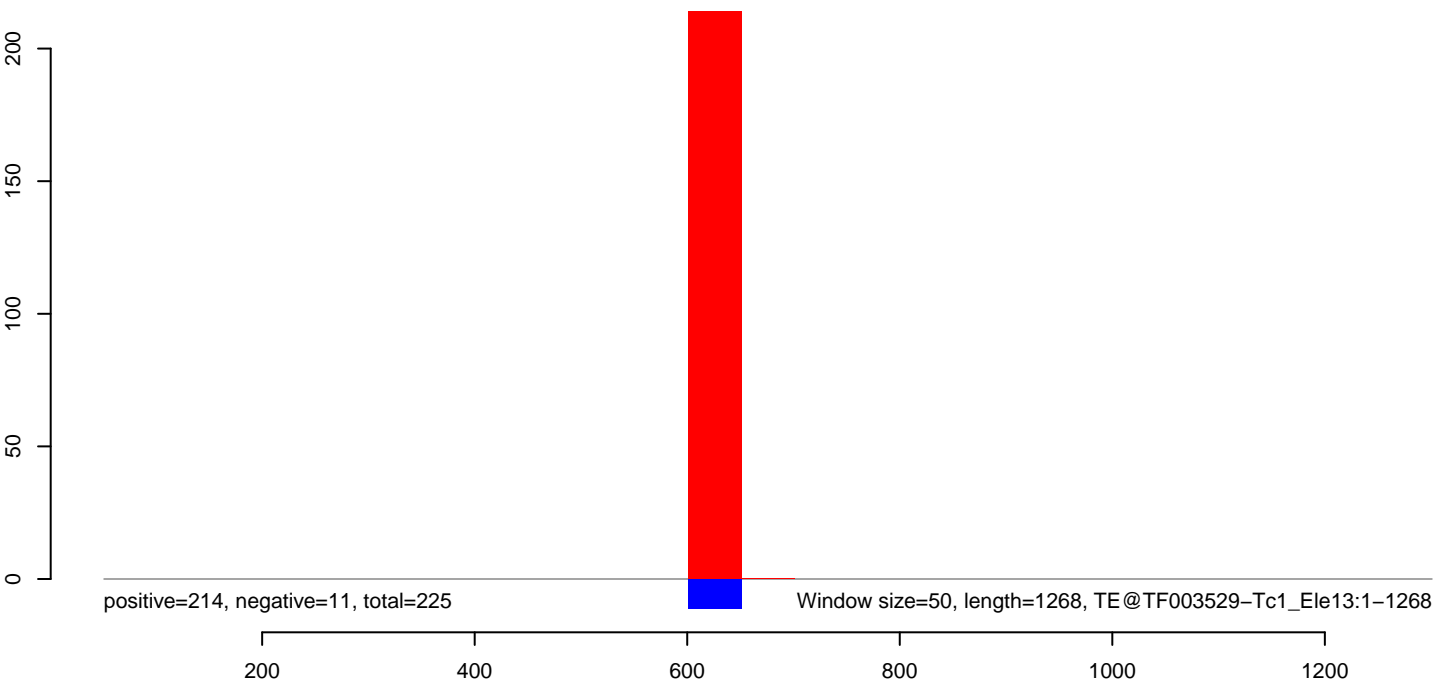
AnGam_Mos55_cells.18_23.rep



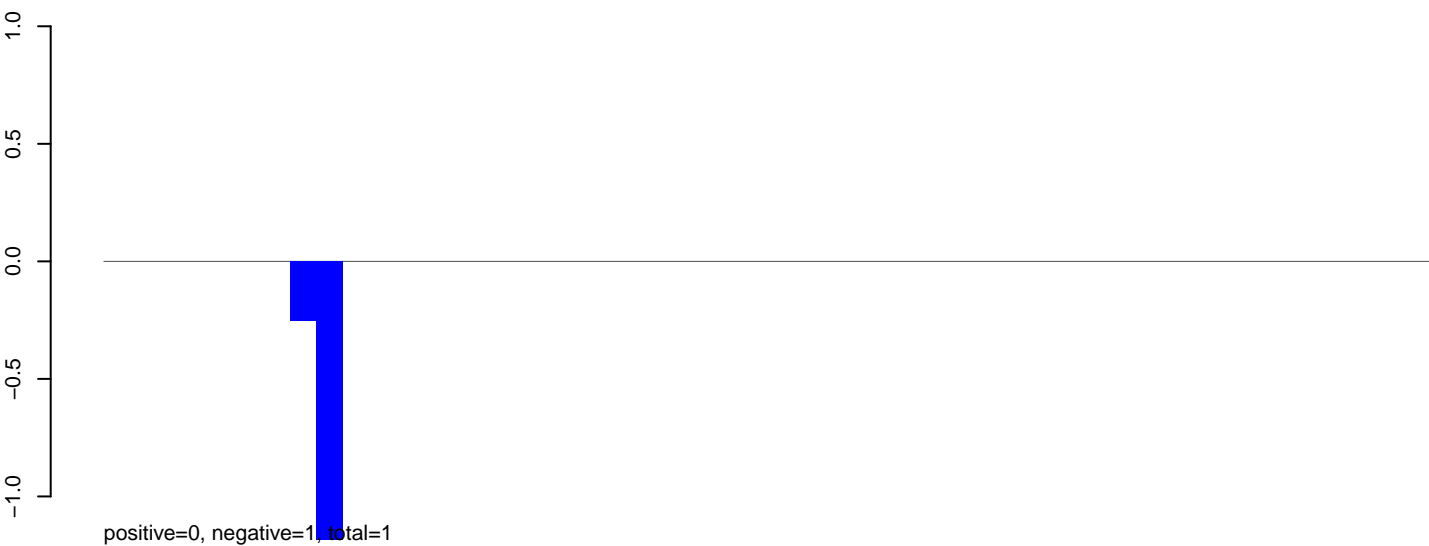
AnGam_Mos55_cells.24_35.rep



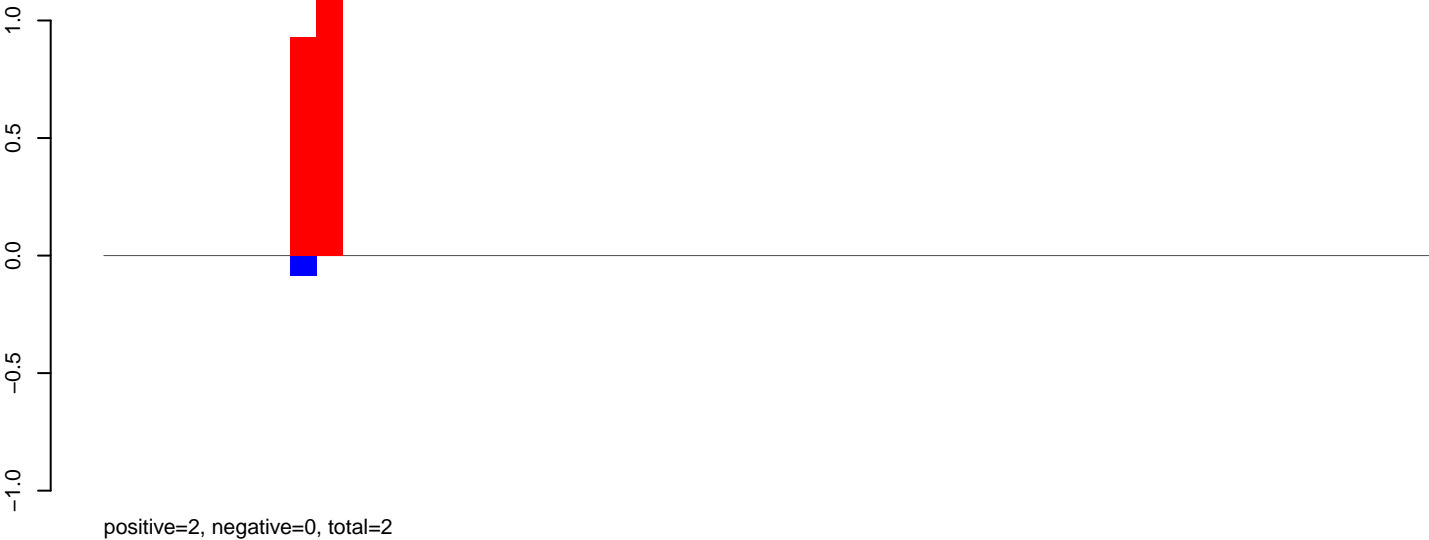
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



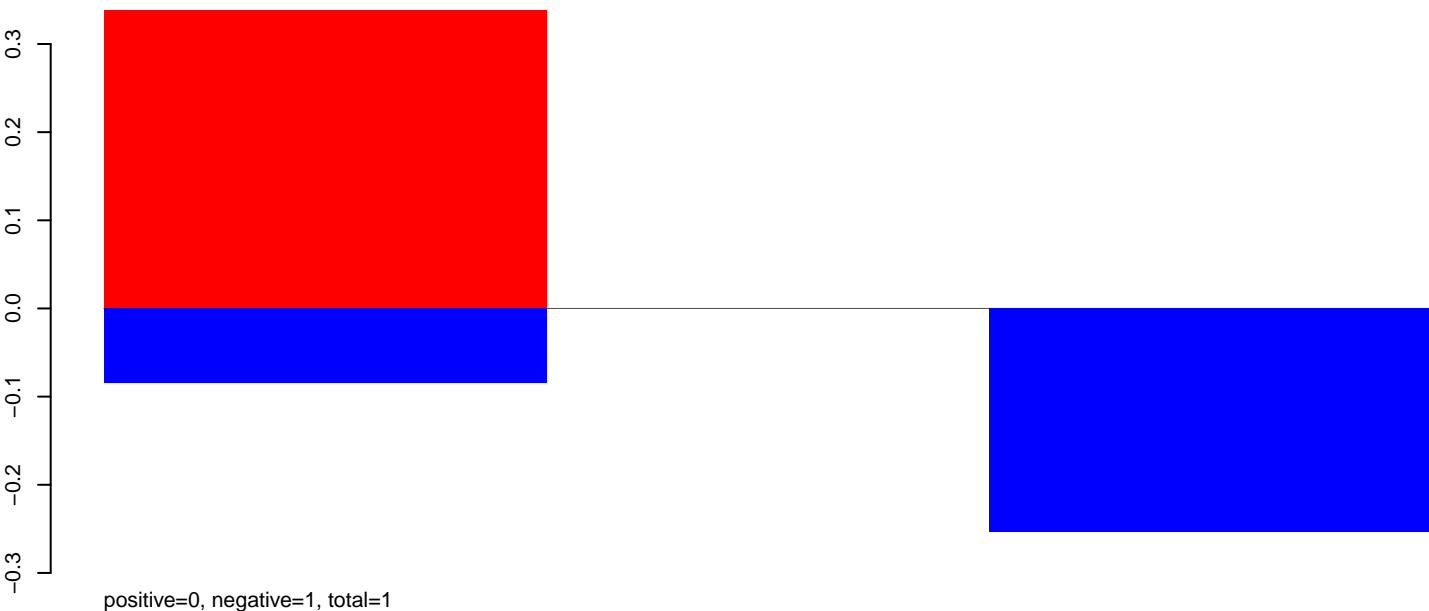
AnGam_Mos55_cells.24_35.rep



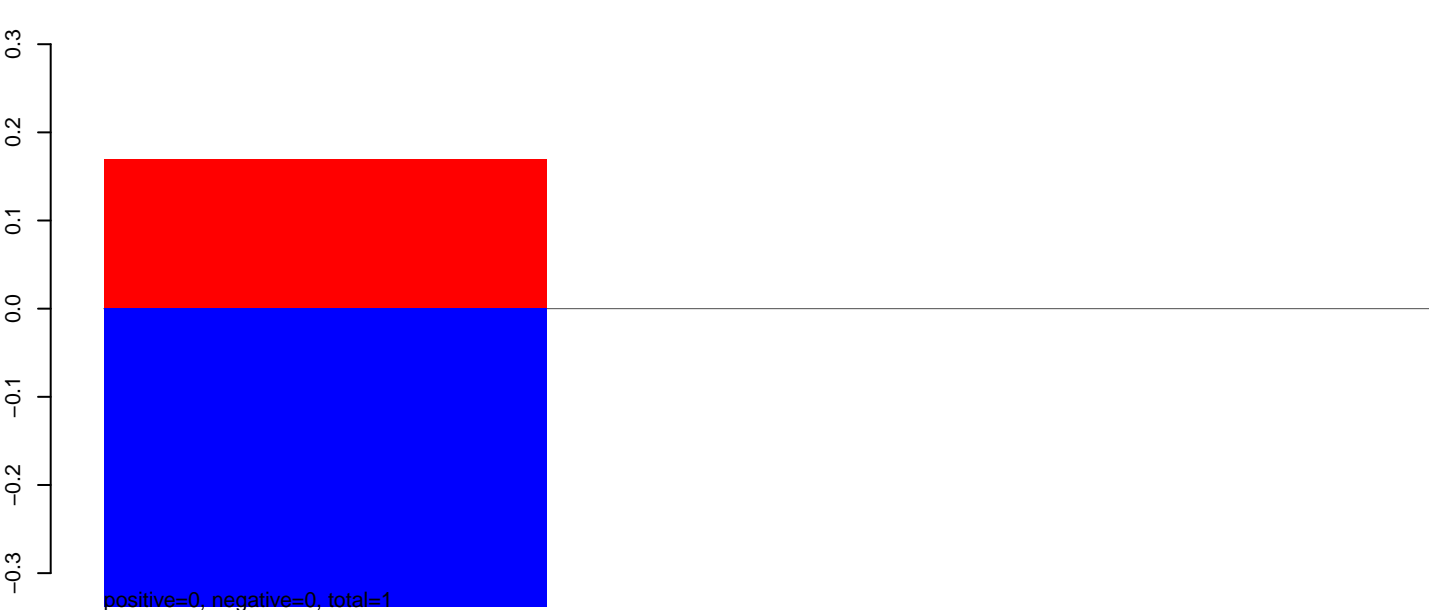
AnGam_Mos55_cells.rep



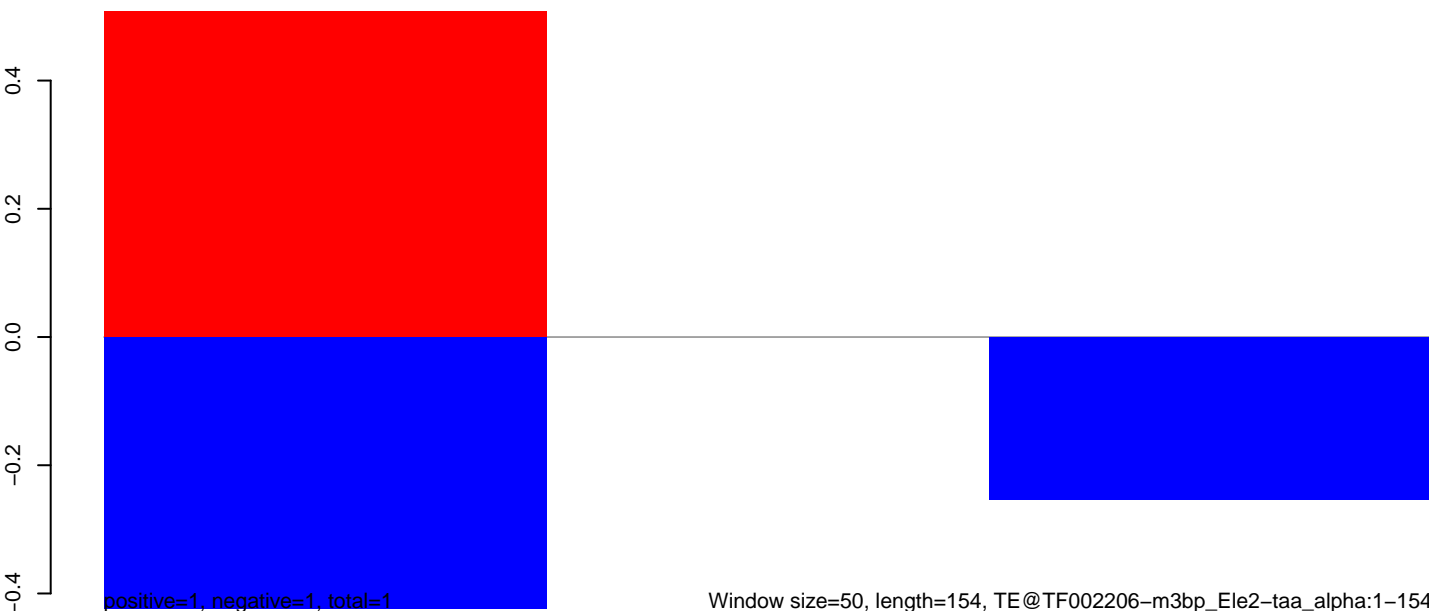
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



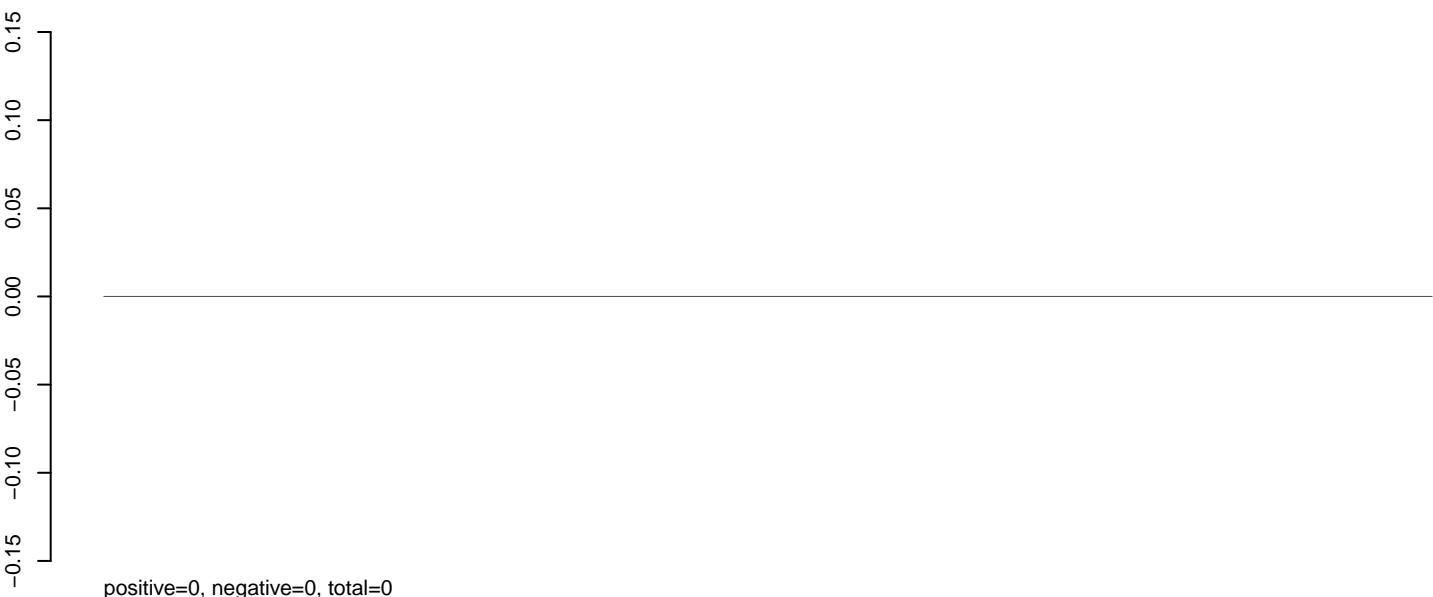
AnGam_Mos55_cells.rep



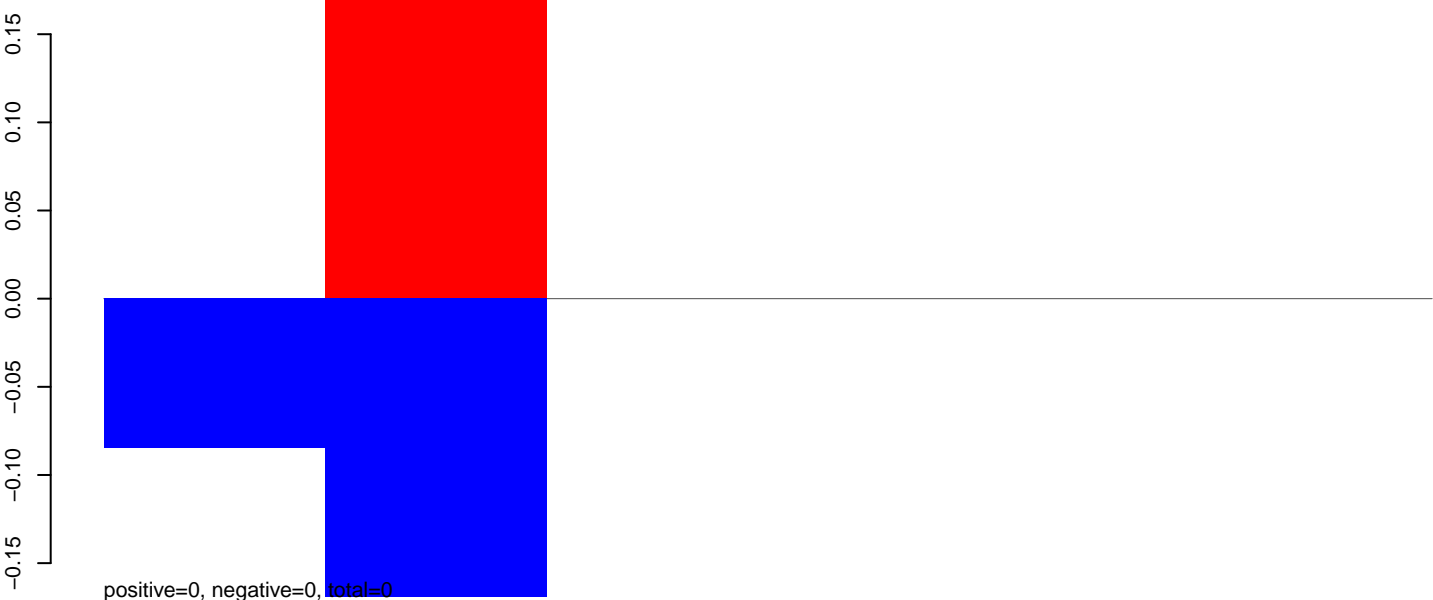
Window size=50, length=154, TE@TF002206-m3bp_Ele2-taa_alpha:1-154

50 100 150 200

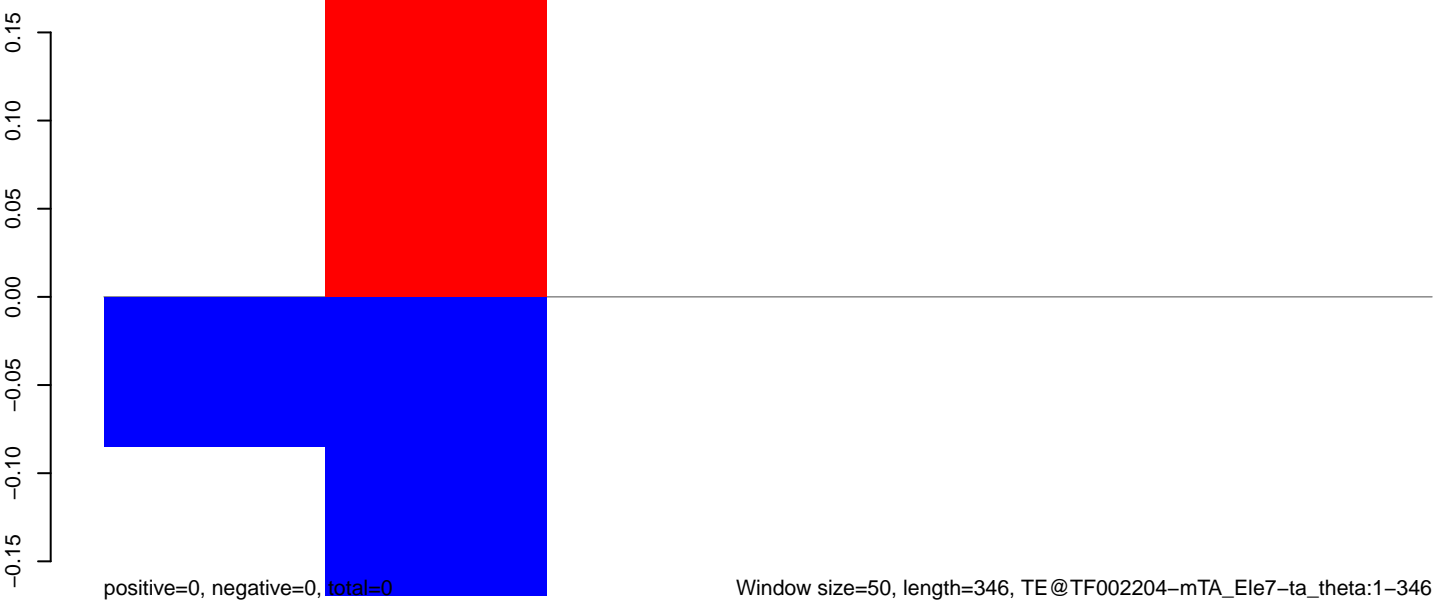
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



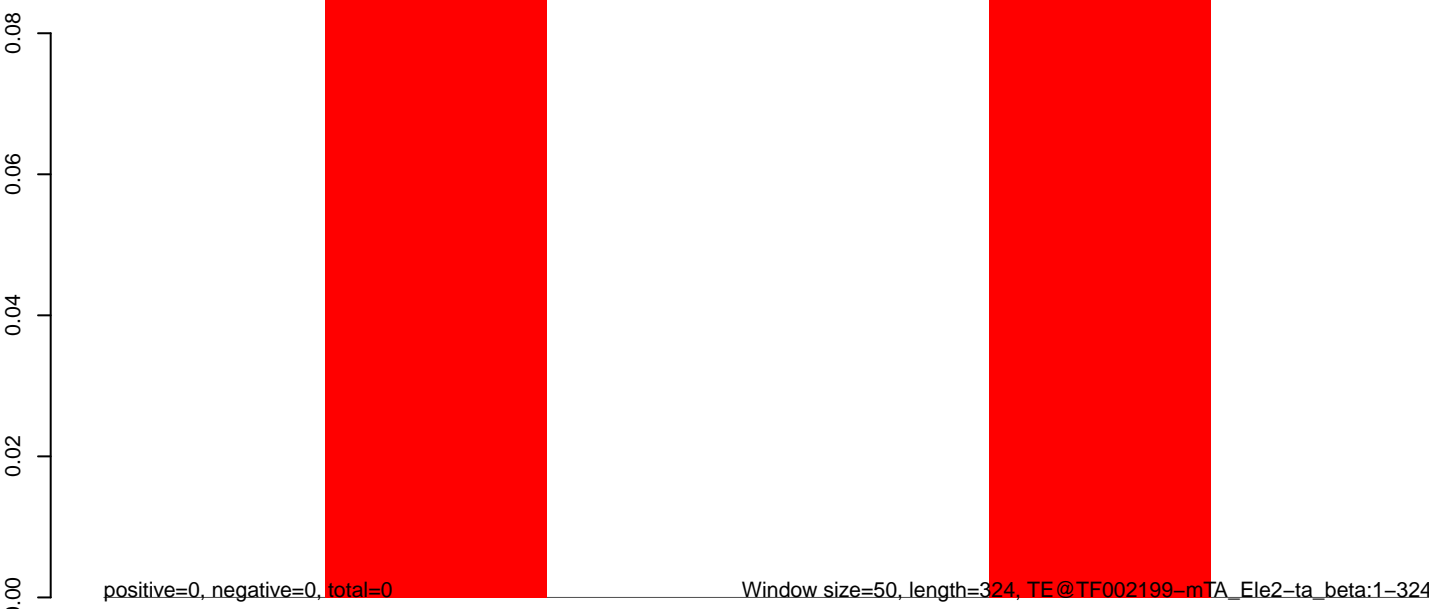
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=7126, TE@TF001970-Pao_Bel_Ele39:1-7126

0 1000 2000 3000 4000 5000 6000 7000

AnGam_Mos55_cells.18_23.rep

0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

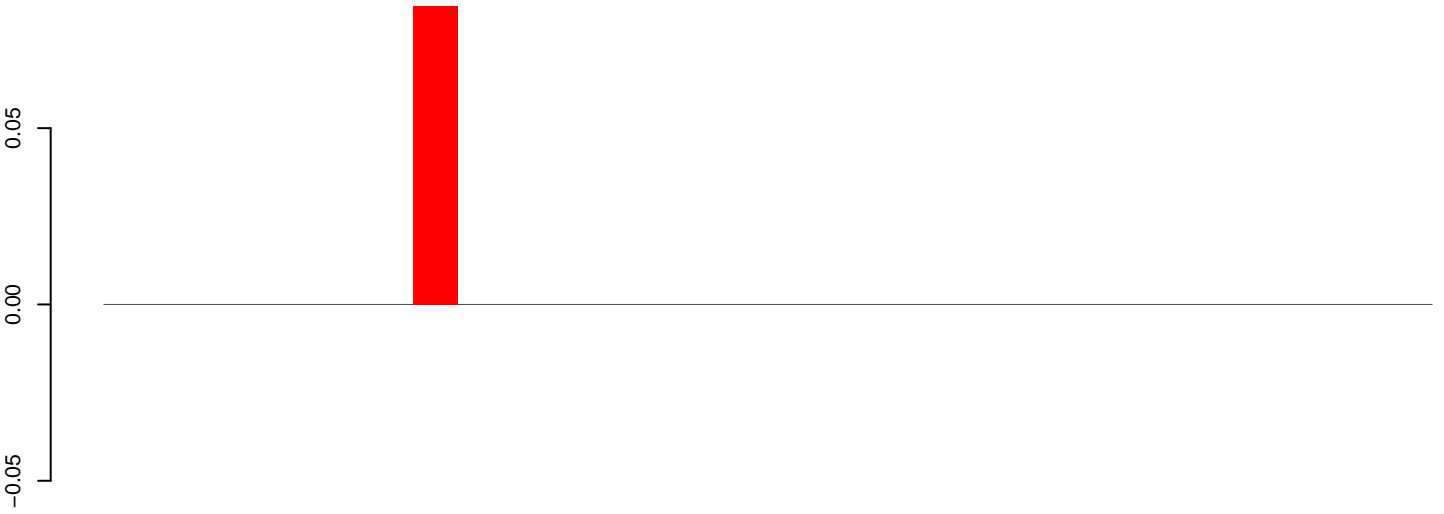
0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

Window size=50, length=2507, TE@TF001435-CR1_Ele2:1-2507

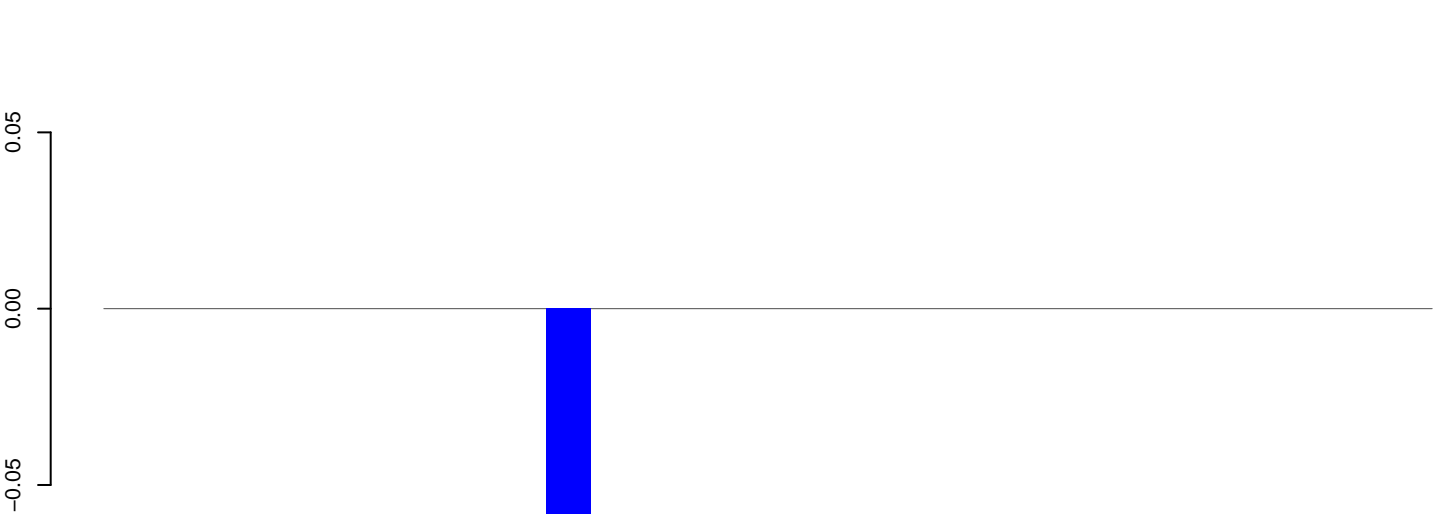
0 500 1000 1500 2000 2500

AnGam_Mos55_cells.18_23.rep



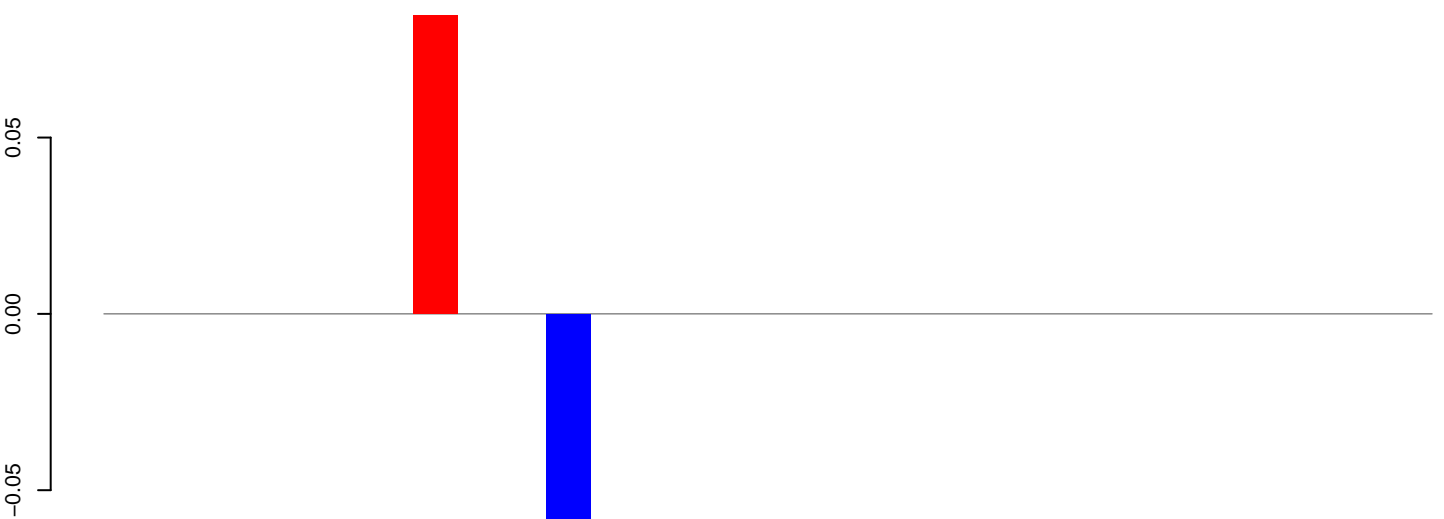
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

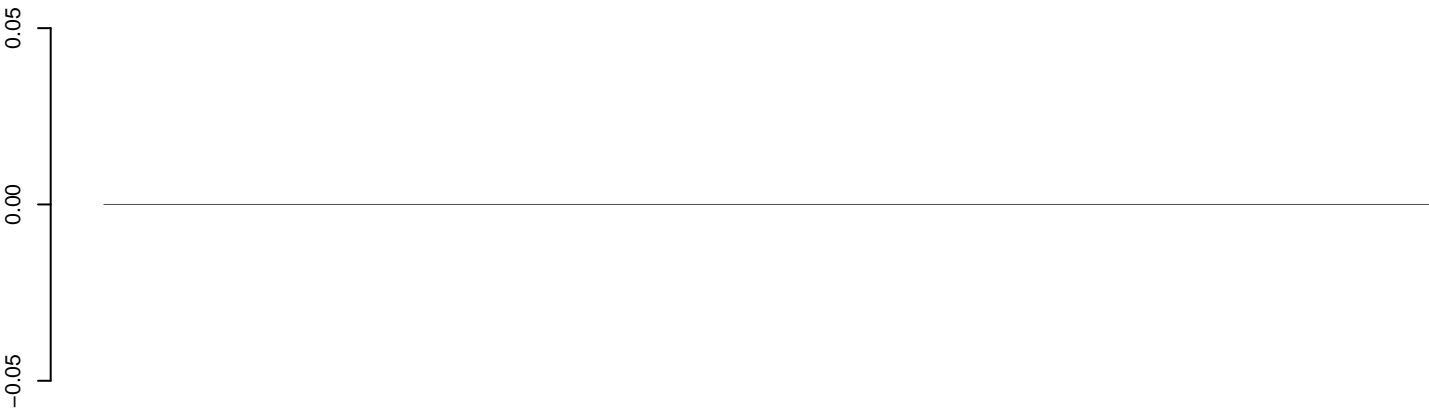


positive=0, negative=0, total=0

Window size=50, length=1512, TE@TF001403-L2_Ele2:1-1512

0 500 1000 1500

AnGam_Mos55_cells.18_23.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



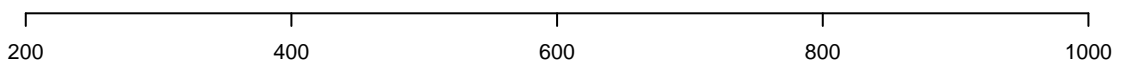
positive=0, negative=0, total=0

AnGam_Mos55_cells.rep



positive=0, negative=0, total=0

Window size=50, length=1017, TE@TF001231-gambol_Ele9-Watteau:1-1017



AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=848, TE@TF001213-mar_Ele7:1-848

200

400

600

800

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

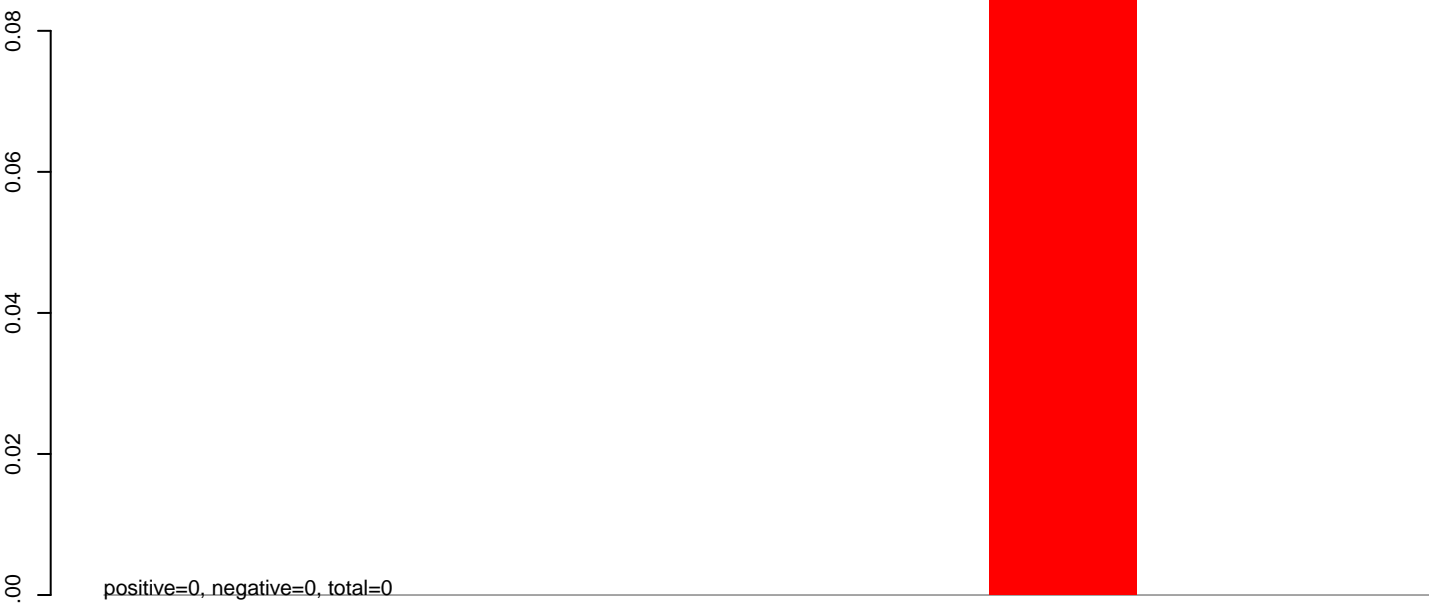
0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

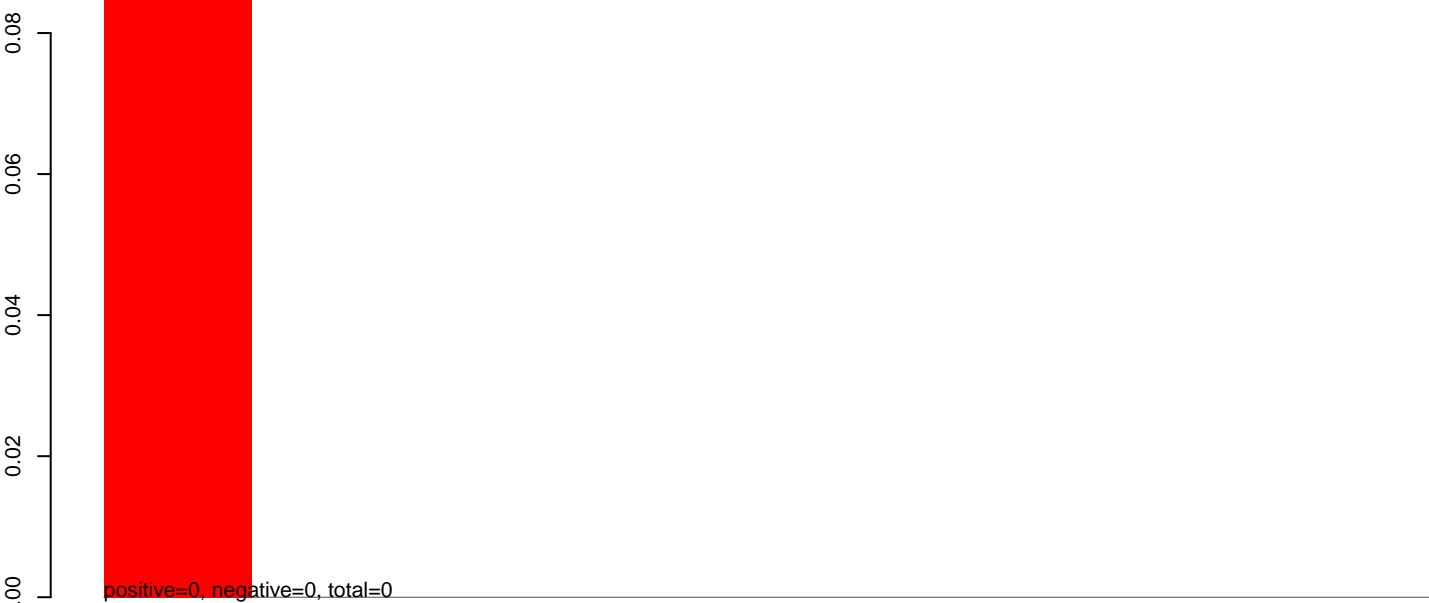
Window size=50, length=410, TE@RETRO6_AG_LTR:1-410

100 200 300 400

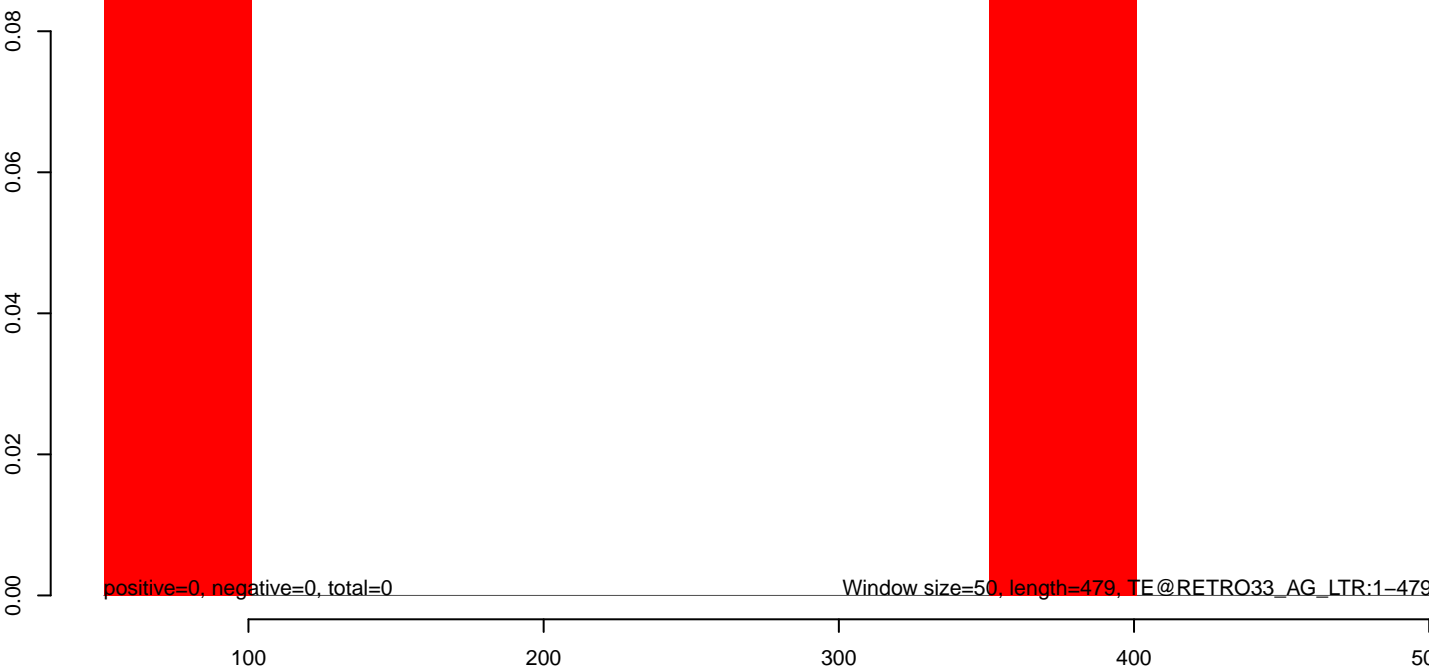
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=3696, TE@PegasusA:1-3696

0 1000 2000 3000

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

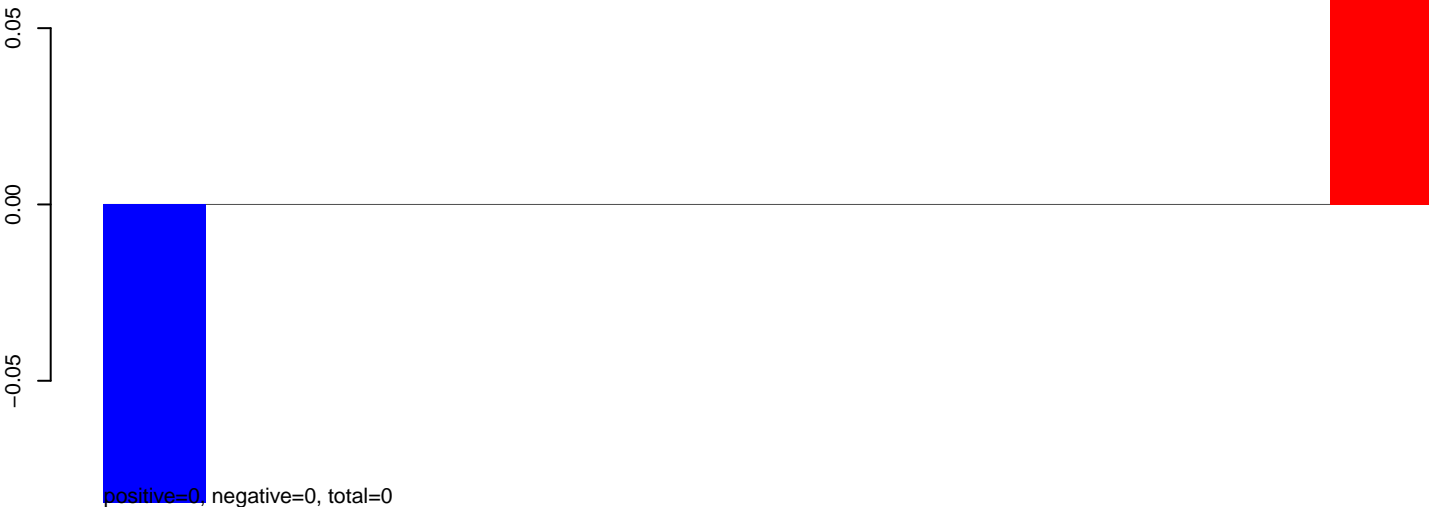
0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

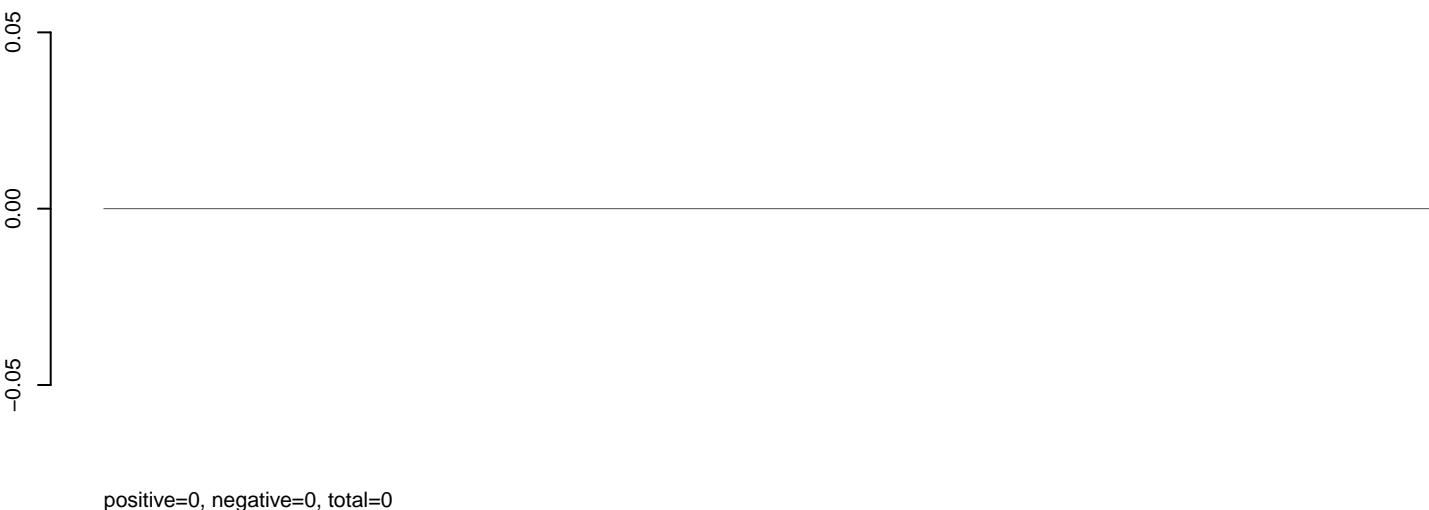
Window size=50, length=4196, TE@P2_AG1-4196

0 1000 2000 3000 4000

AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



100

200

300

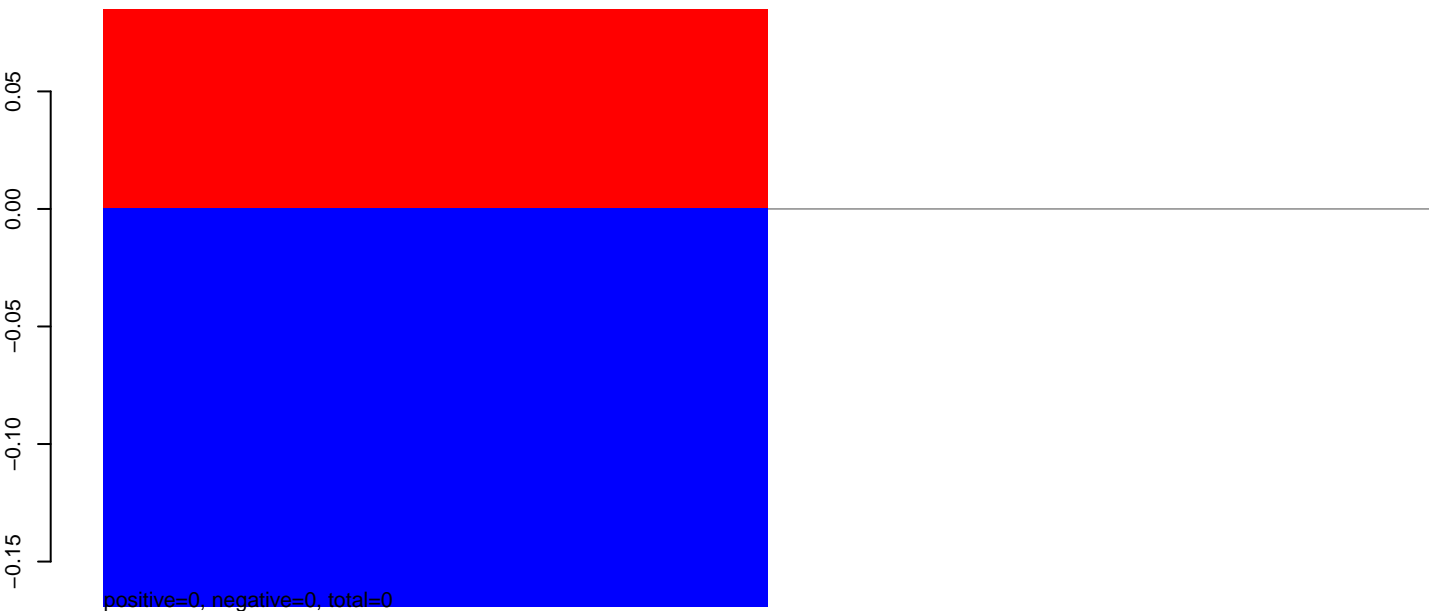
400

500

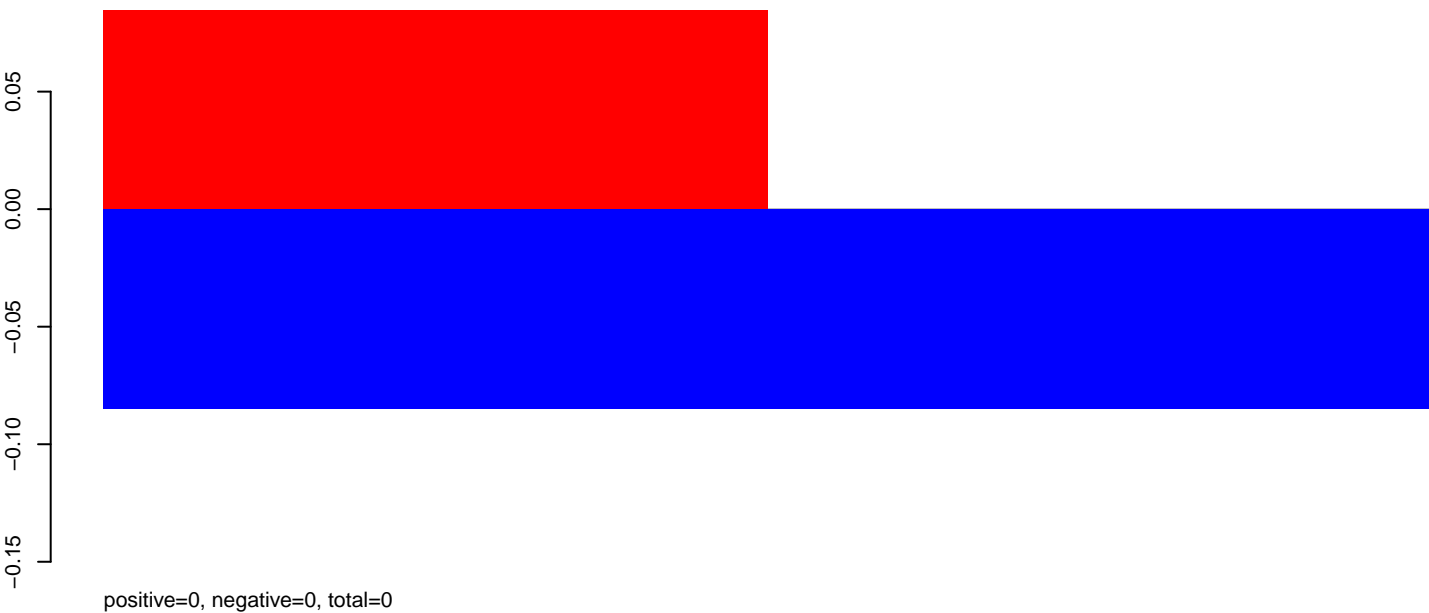
600

700

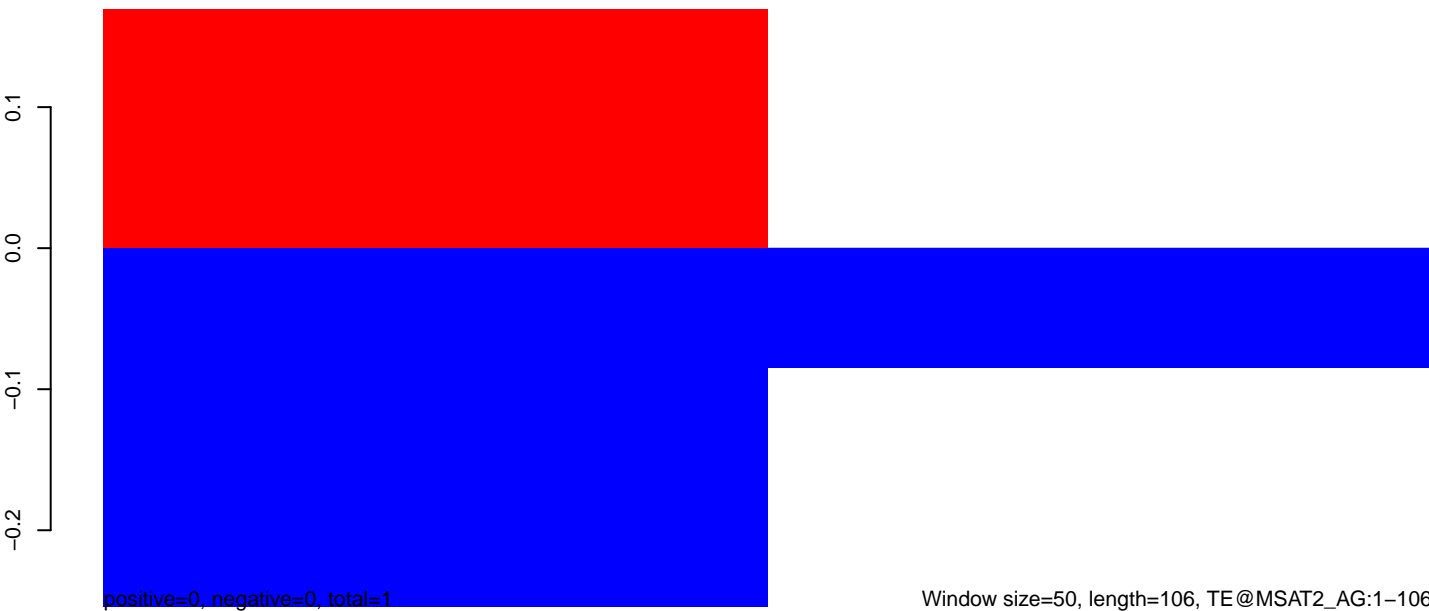
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



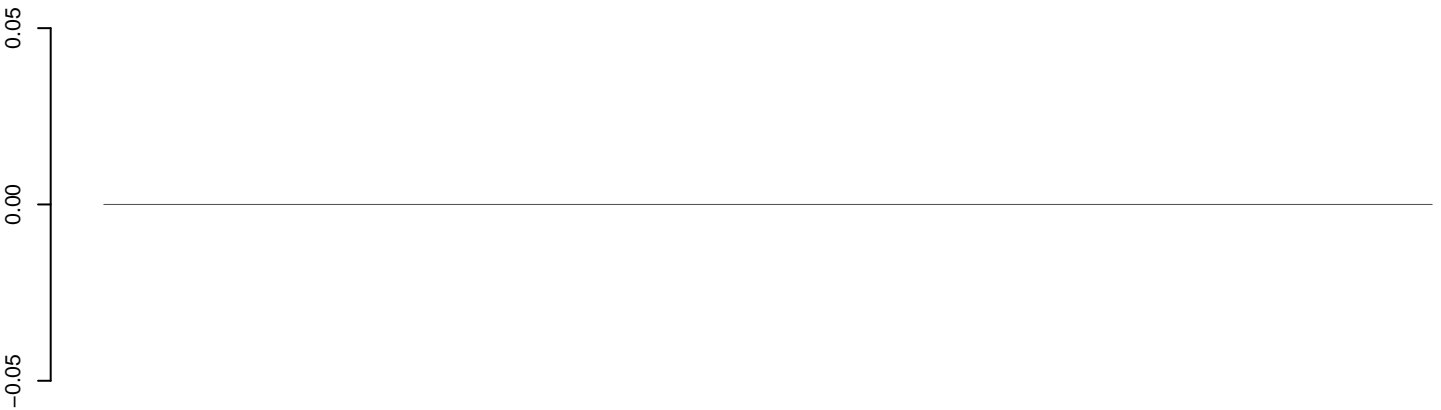
AnGam_Mos55_cells.rep



Window size=50, length=106, TE@MSAT2_AG:1-106

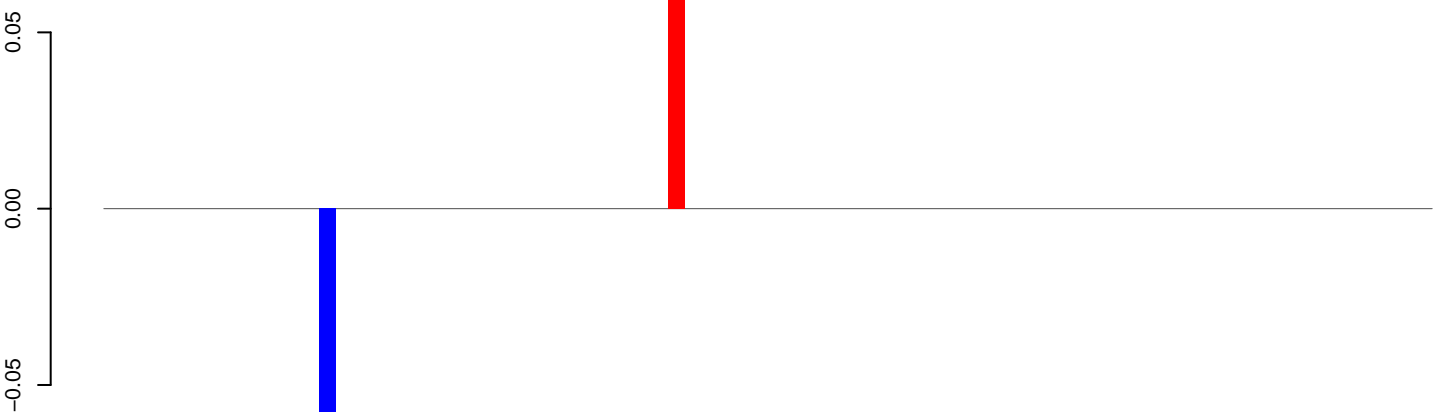
60 80 100 120 140

AnGam_Mos55_cells.18_23.rep



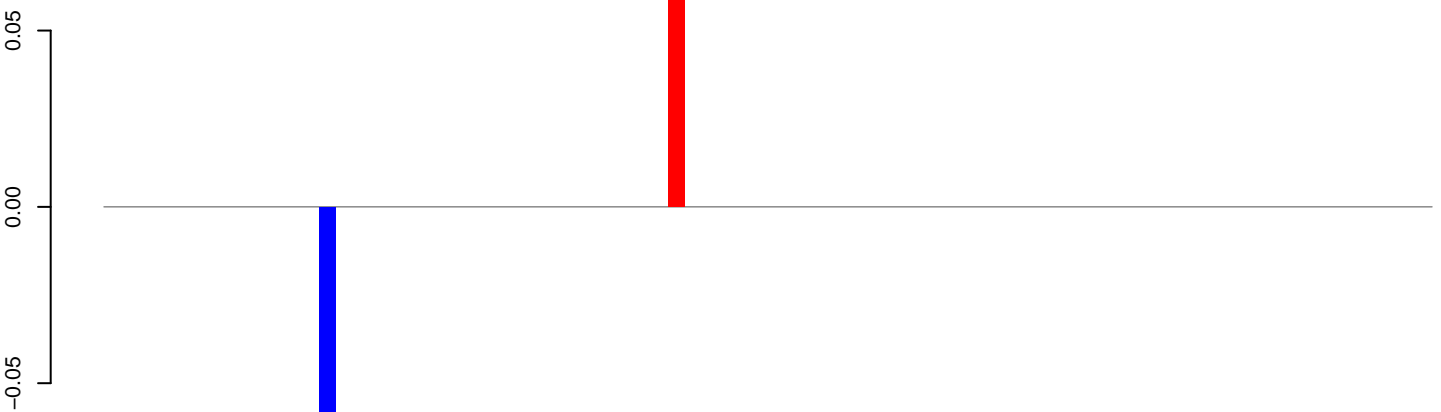
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.rep



positive=0, negative=0, total=0

Window size=50, length=4030, TE@Gypsy-30_AG-LTR-I:1-4030

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

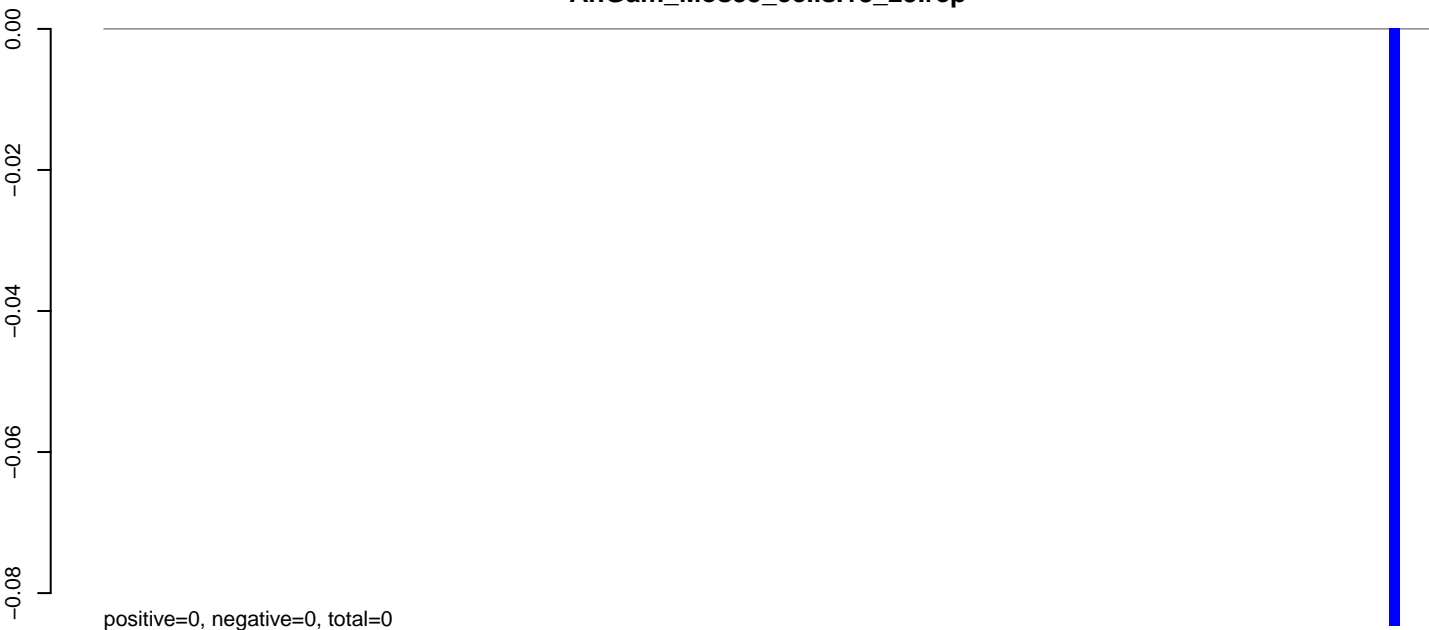
0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

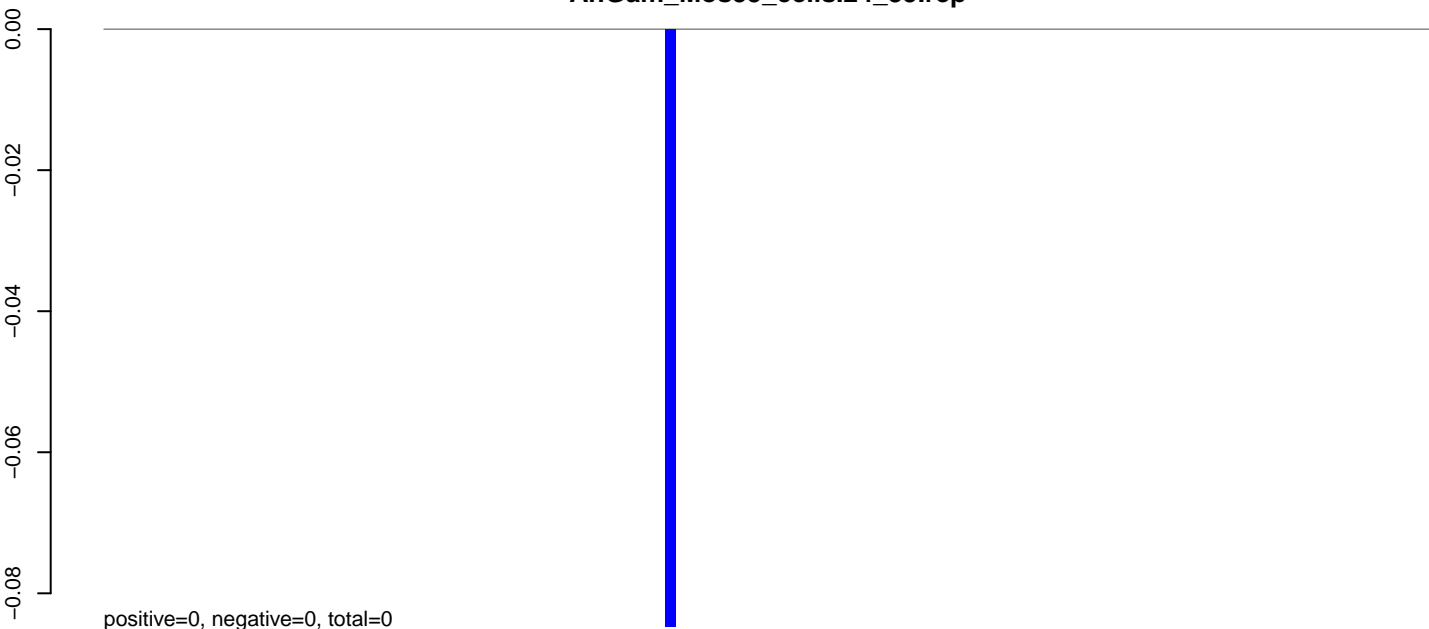
Window size=50, length=4112, TE@Gypsy-18, AG-LTR-I:1-4112

0 1000 2000 3000 4000

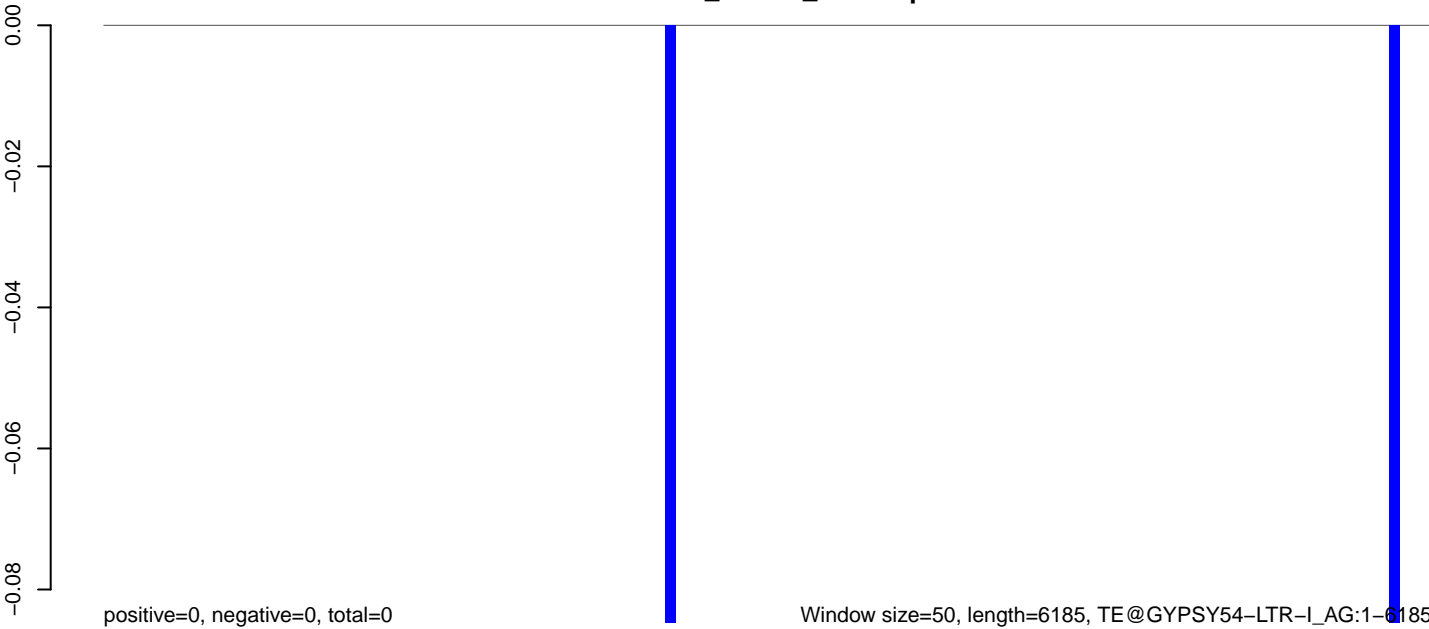
AnGam_Mos55_cells.18_23.rep



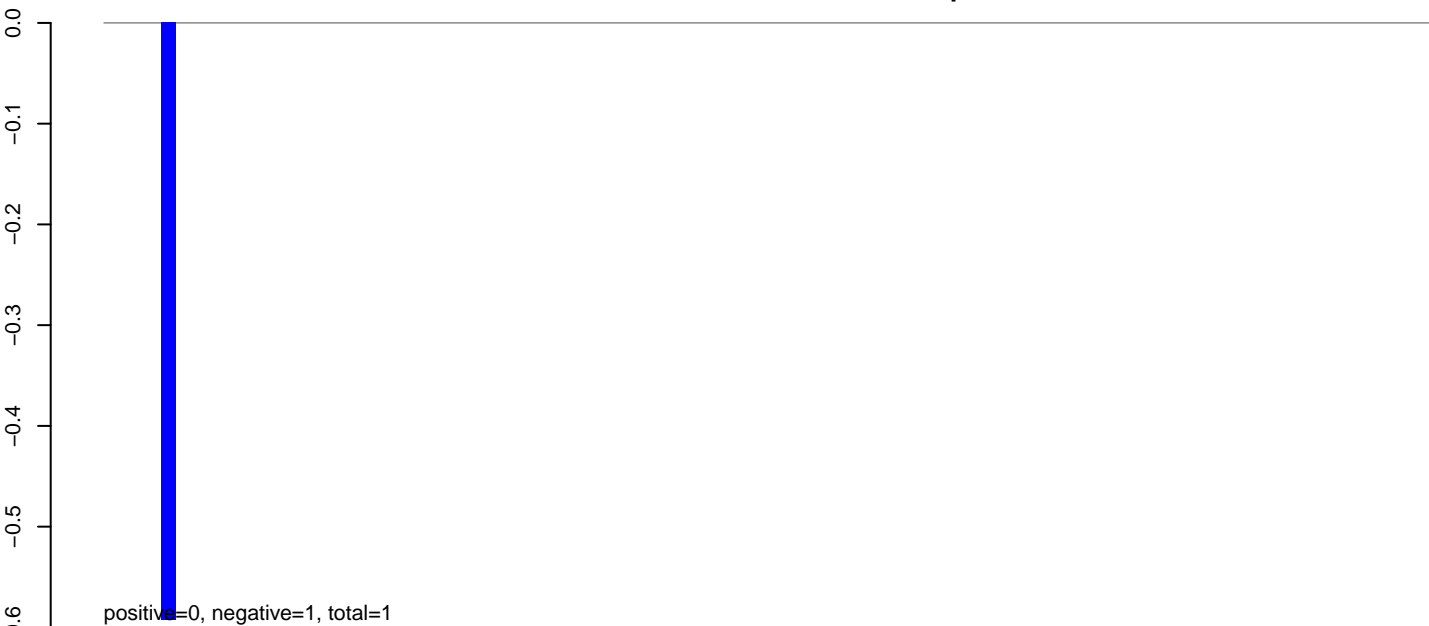
AnGam_Mos55_cells.24_35.rep



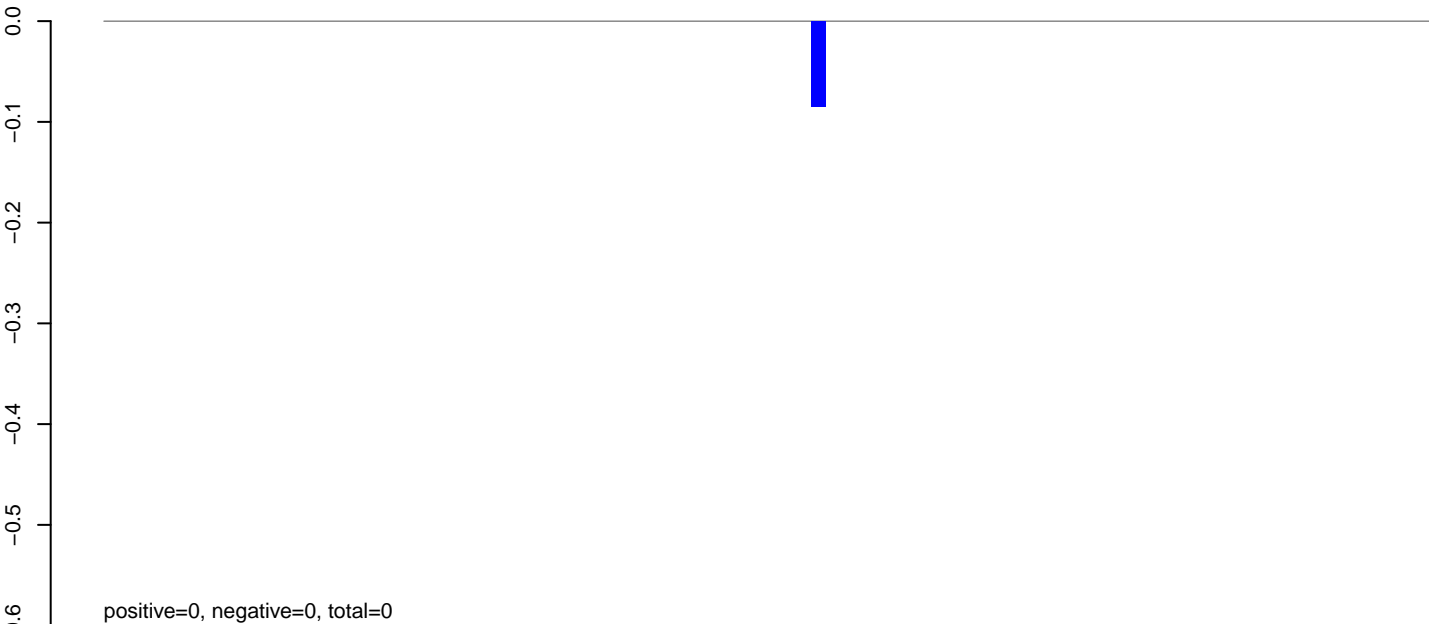
AnGam_Mos55_cells.rep



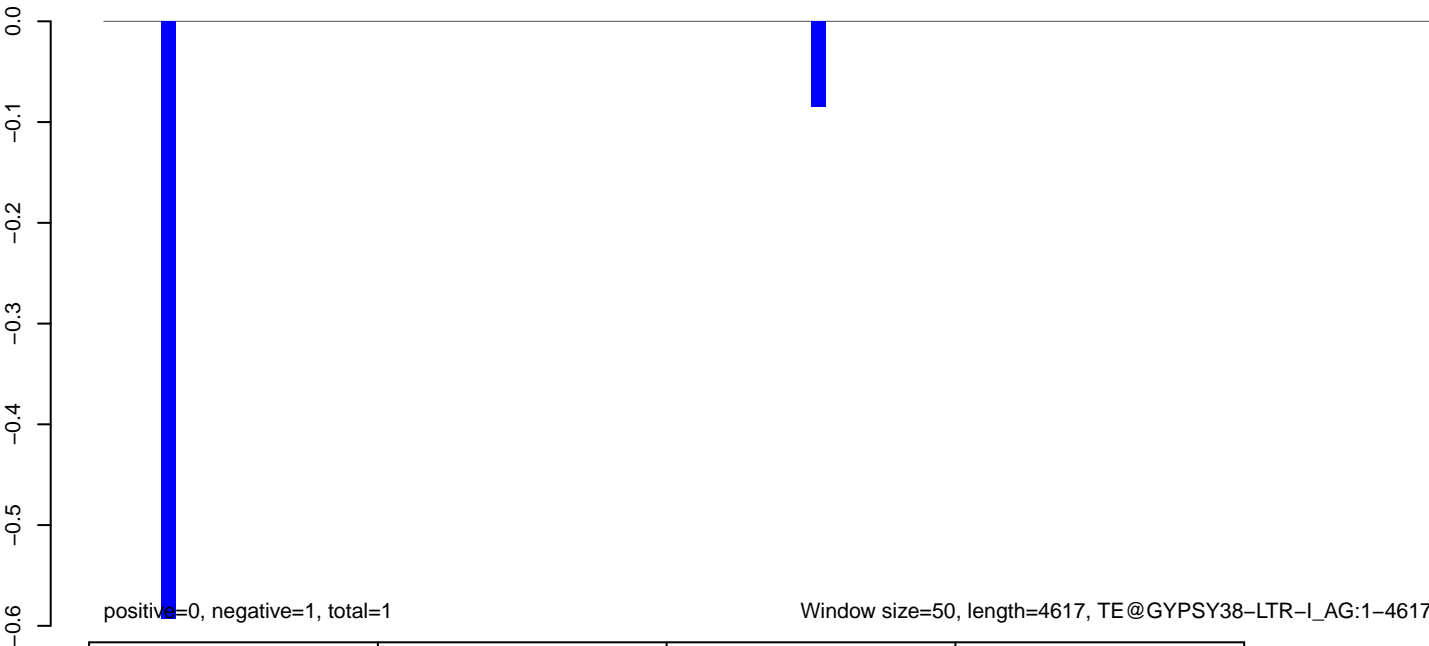
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



Window size=50, length=4617, TE@GYPSY38-LTR-L_AG:1-4617

0 1000 2000 3000 4000

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

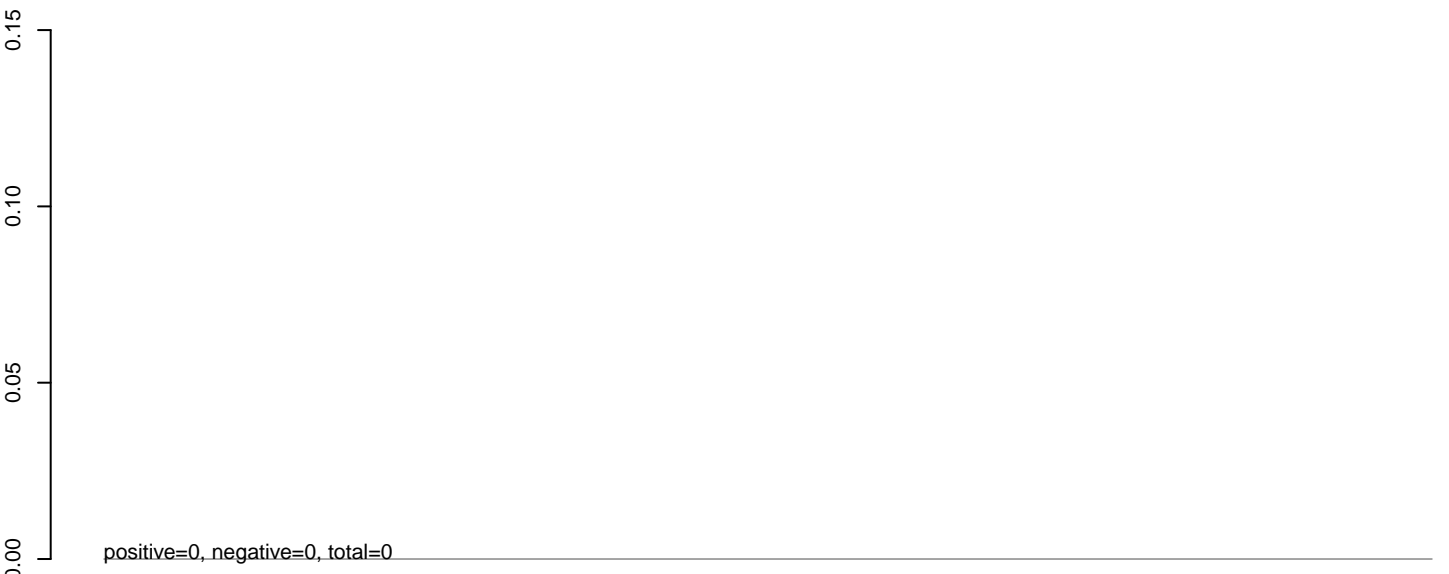
0.00
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-0.04
-0.06
-0.08

positive=0, negative=0, total=0

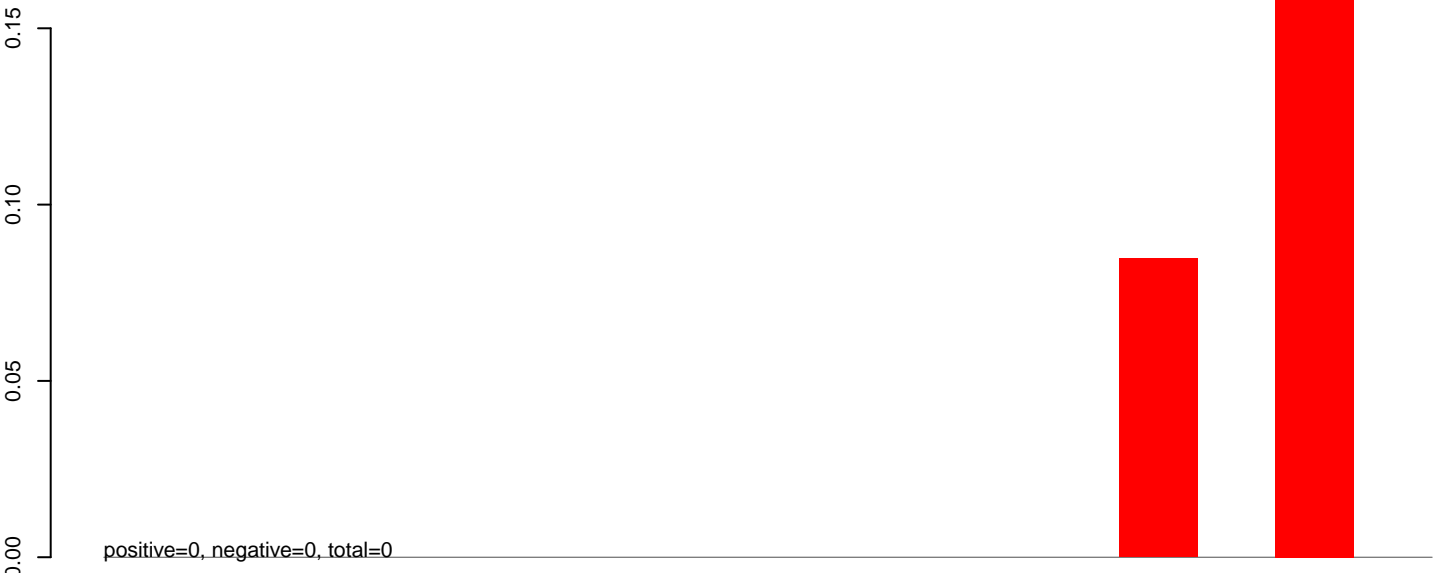
Window size=50, length=4116, TE@GYPSY26-LTR-I_AG:1-4116

0 1000 2000 3000 4000

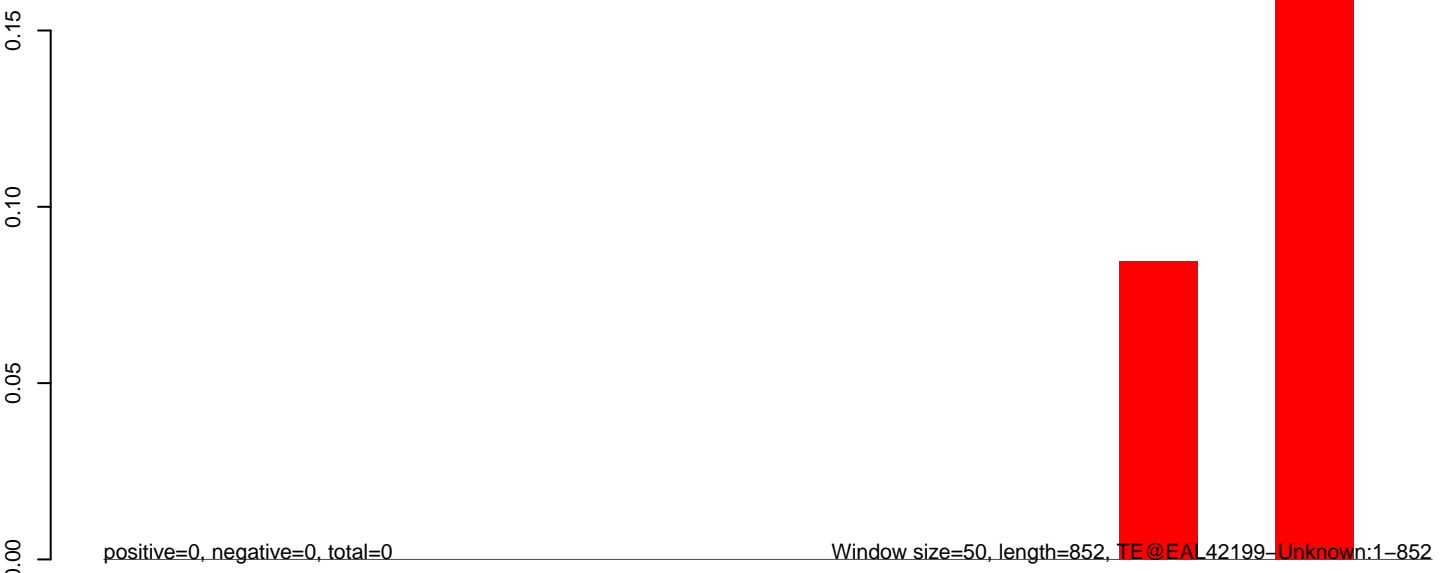
AnGam_Mos55_cells.18_23.rep



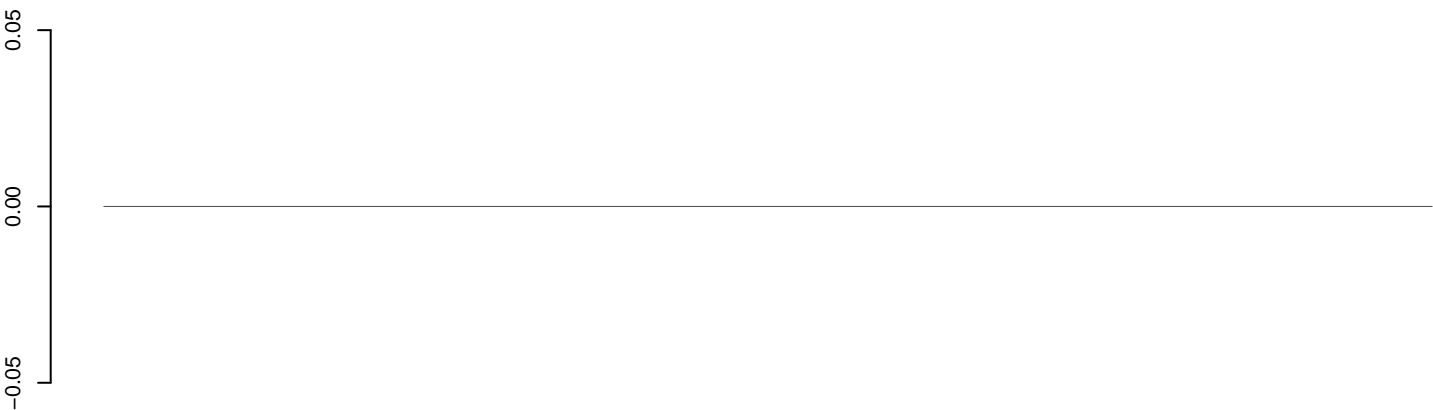
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

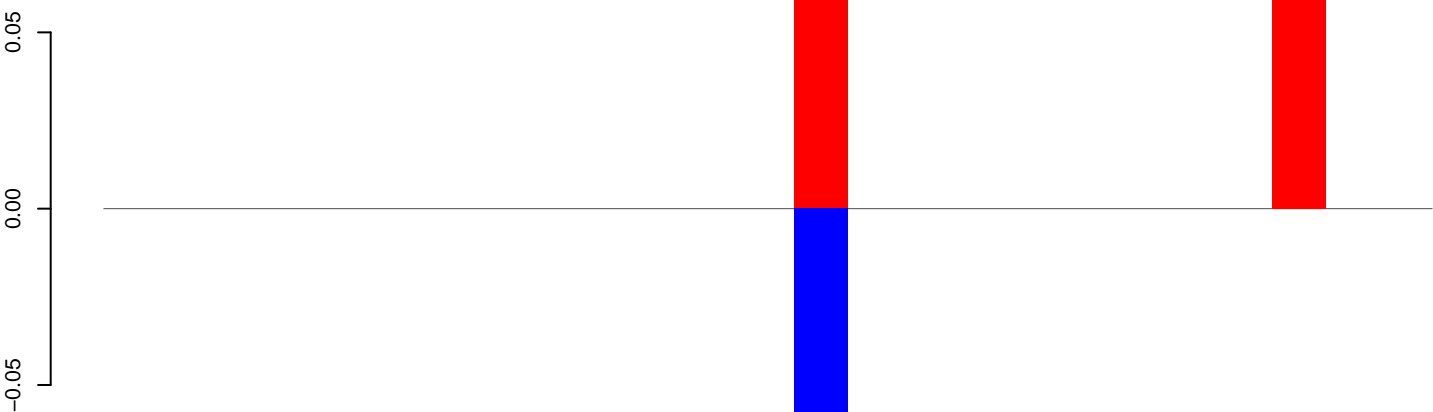


AnGam_Mos55_cells.18_23.rep



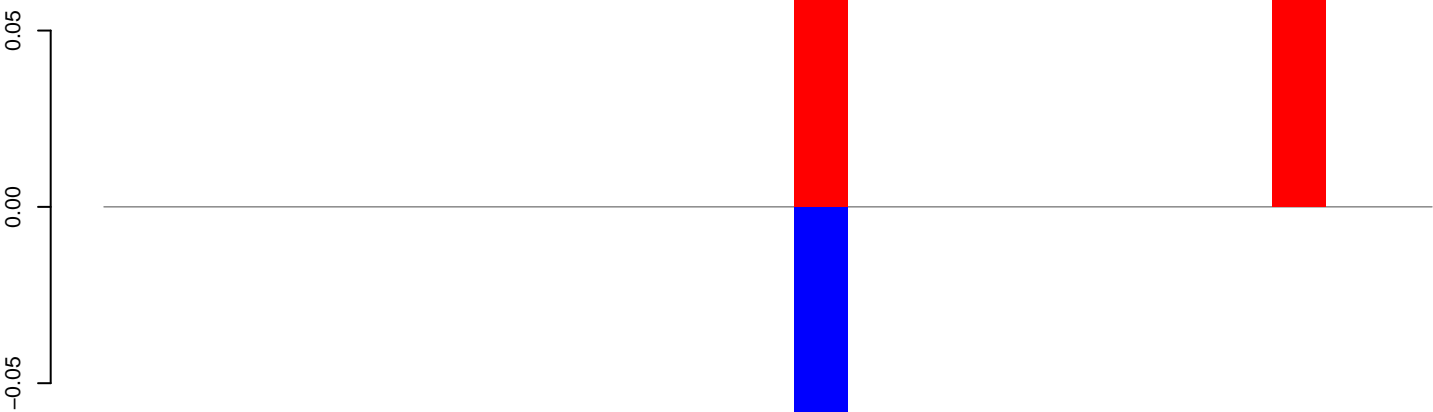
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

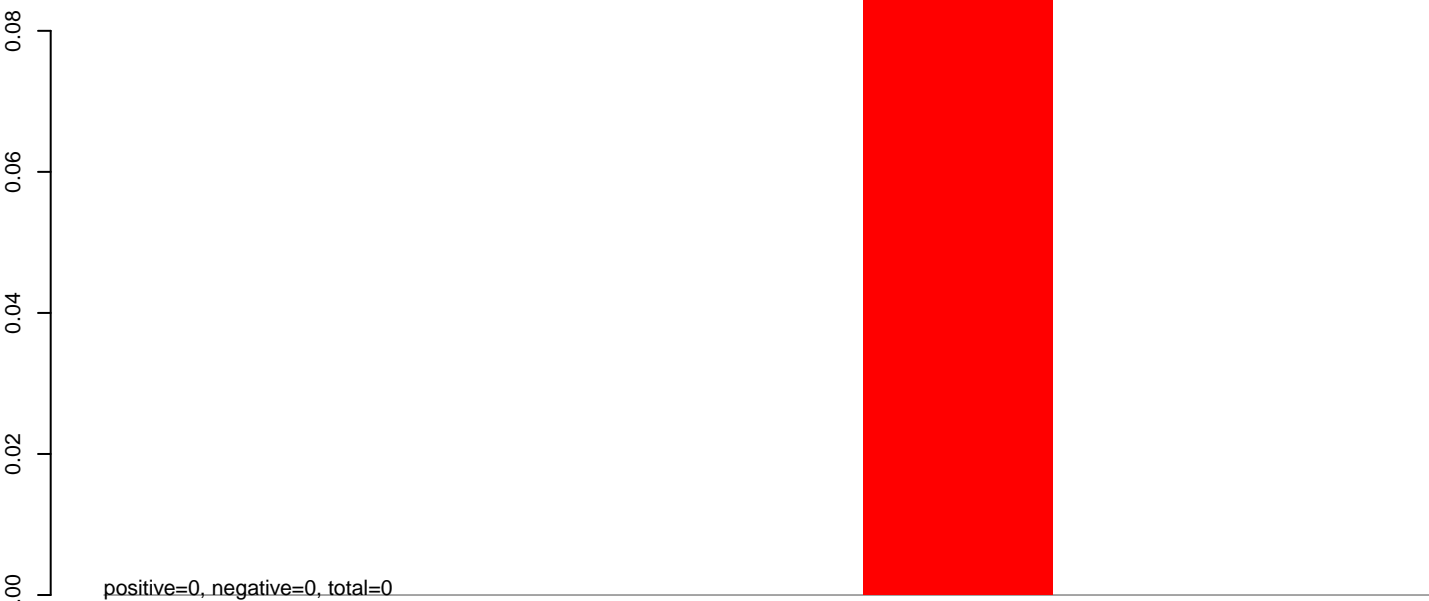


positive=0, negative=0, total=0

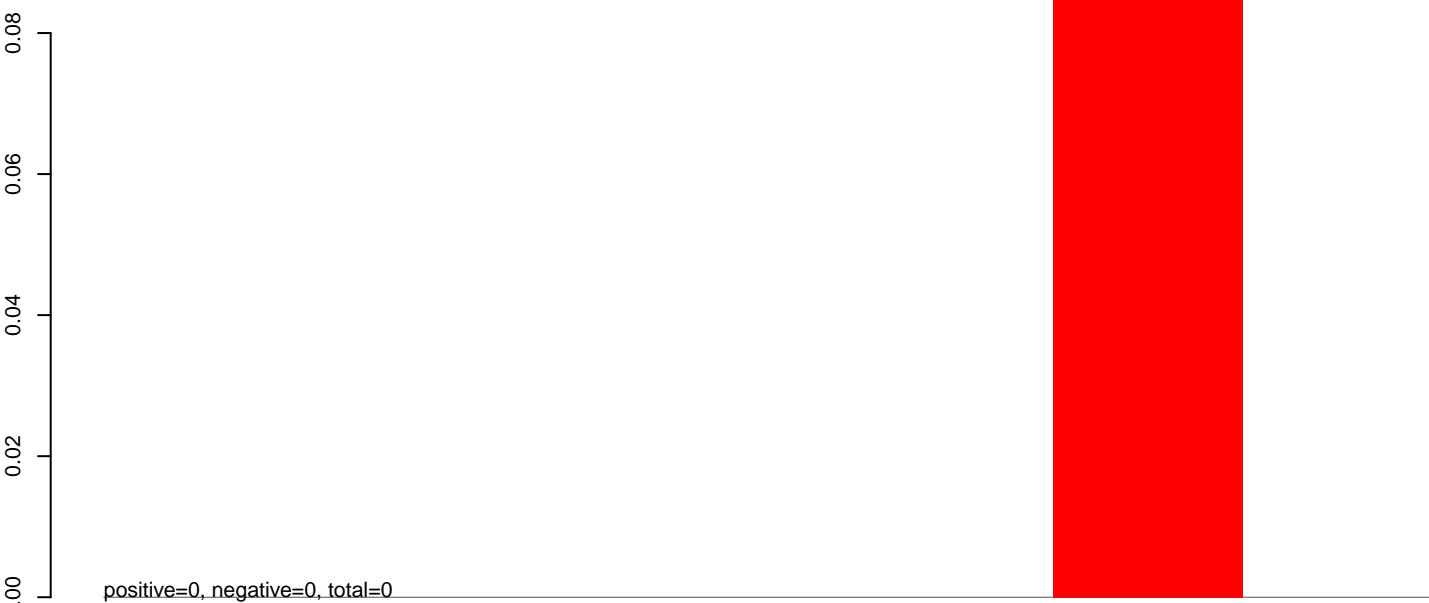
Window size=50, length=1278, TE@EAL42169-Unknown:1-1278

200 400 600 800 1000 1200

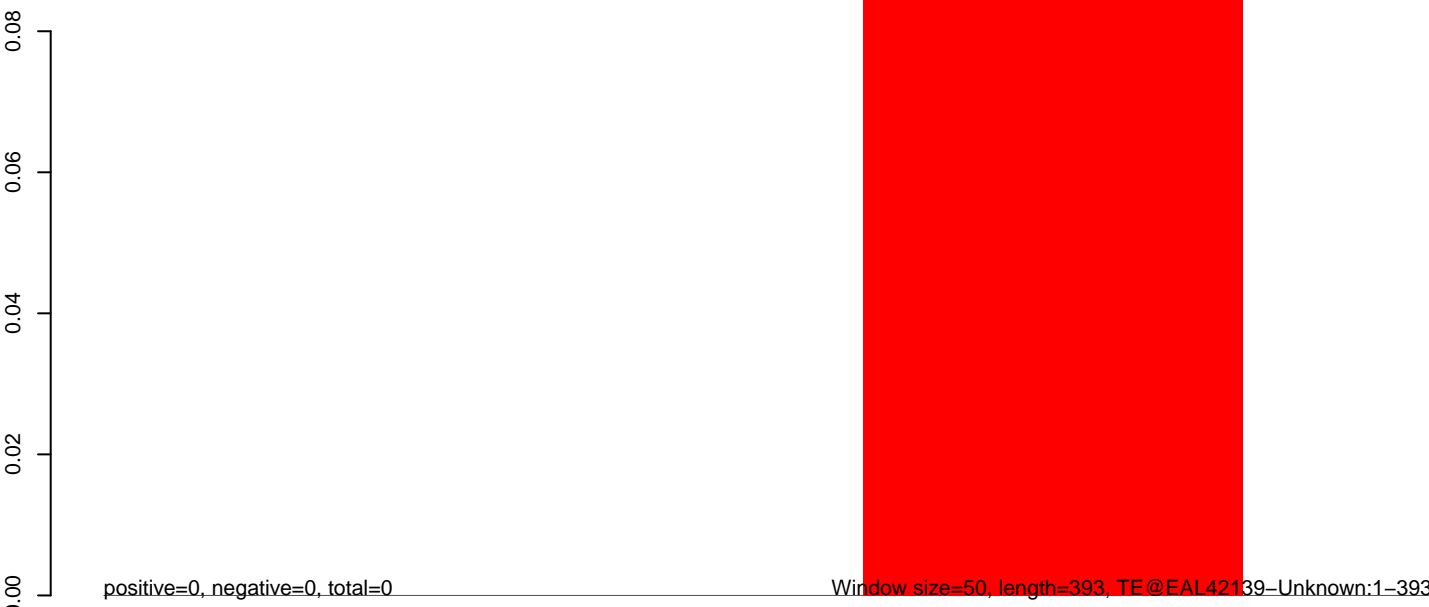
AnGam_Mos55_cells.18_23.rep



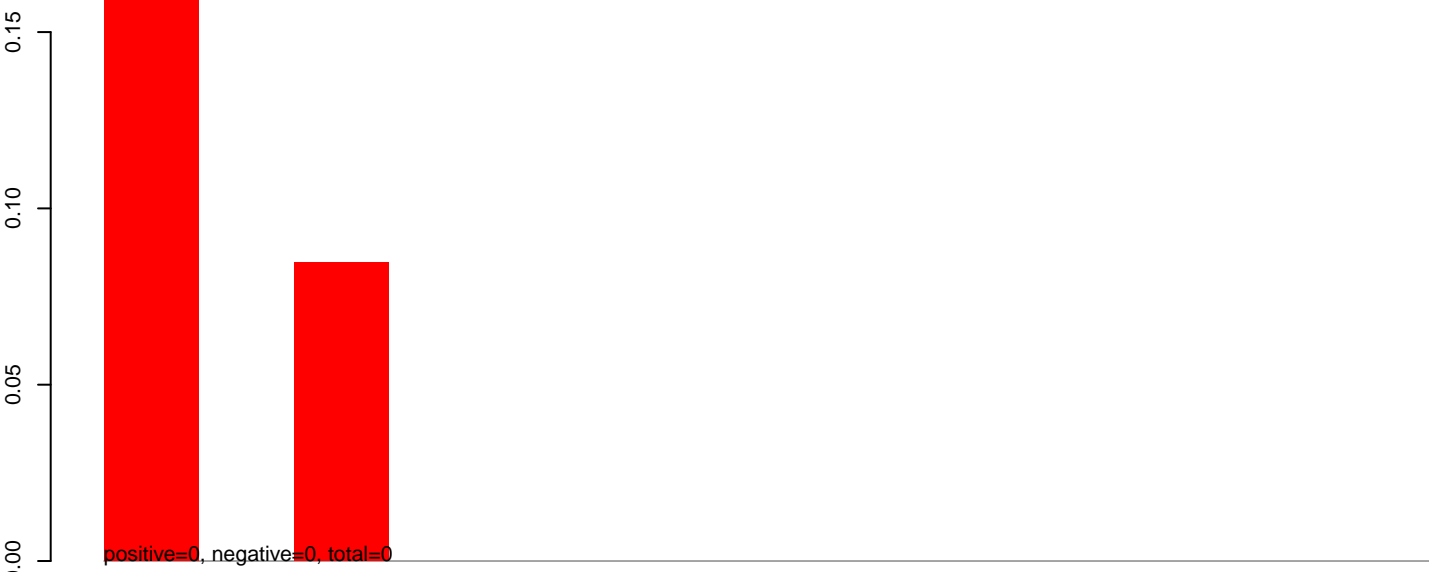
AnGam_Mos55_cells.24_35.rep



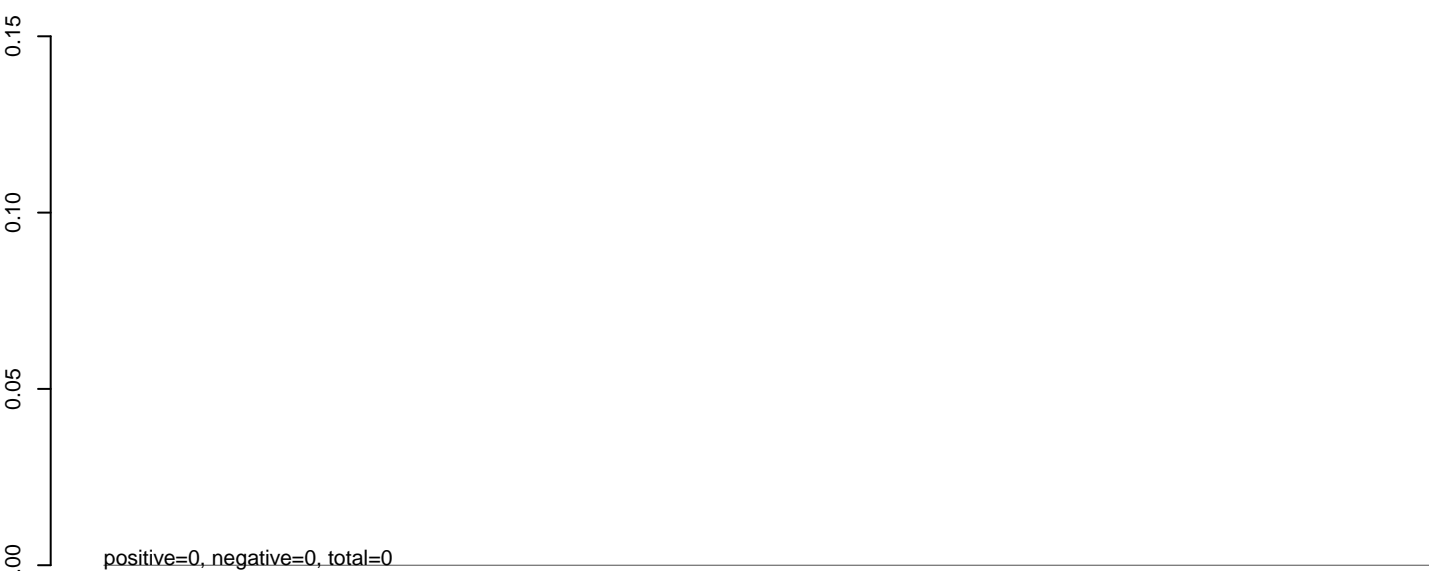
AnGam_Mos55_cells.rep



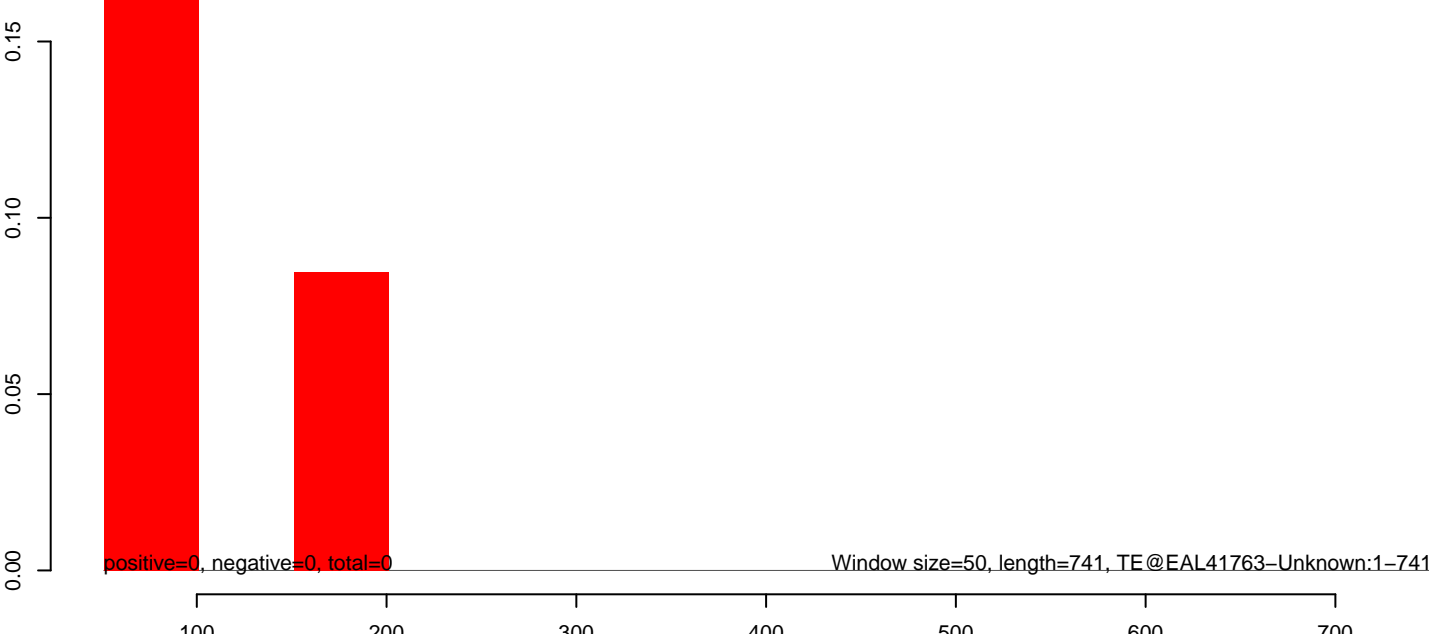
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

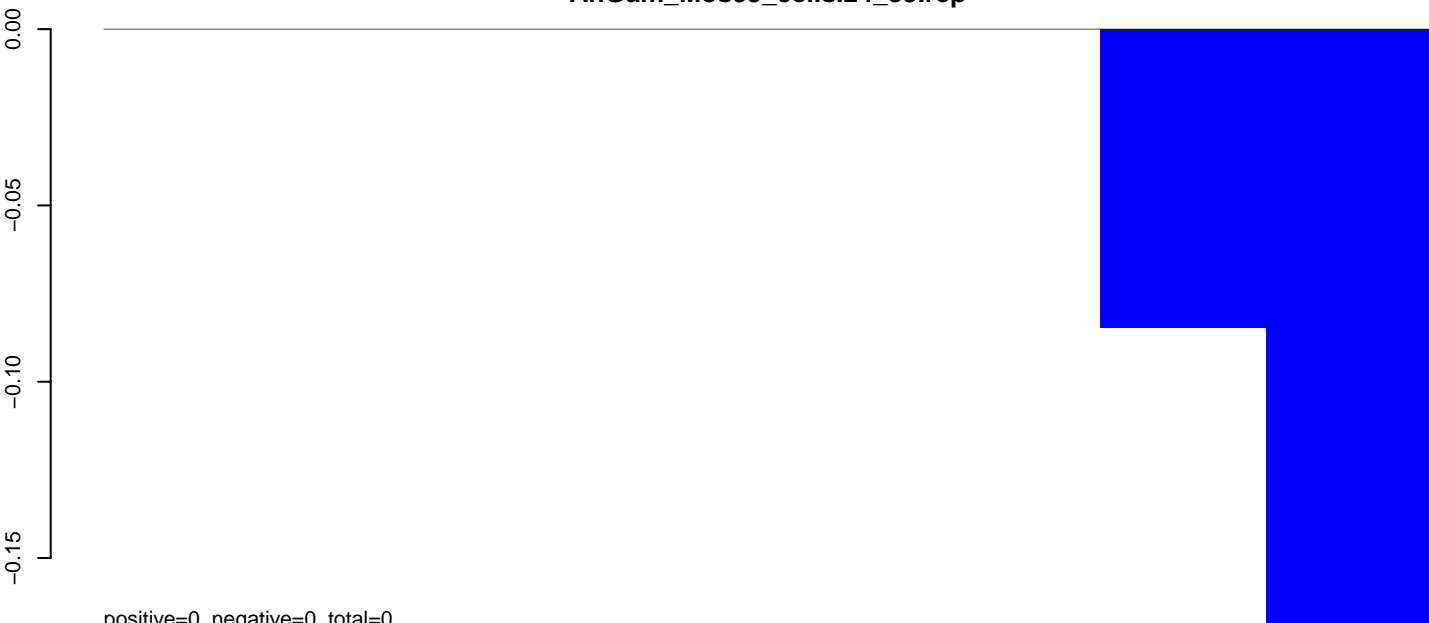


AnGam_Mos55_cells.18_23.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

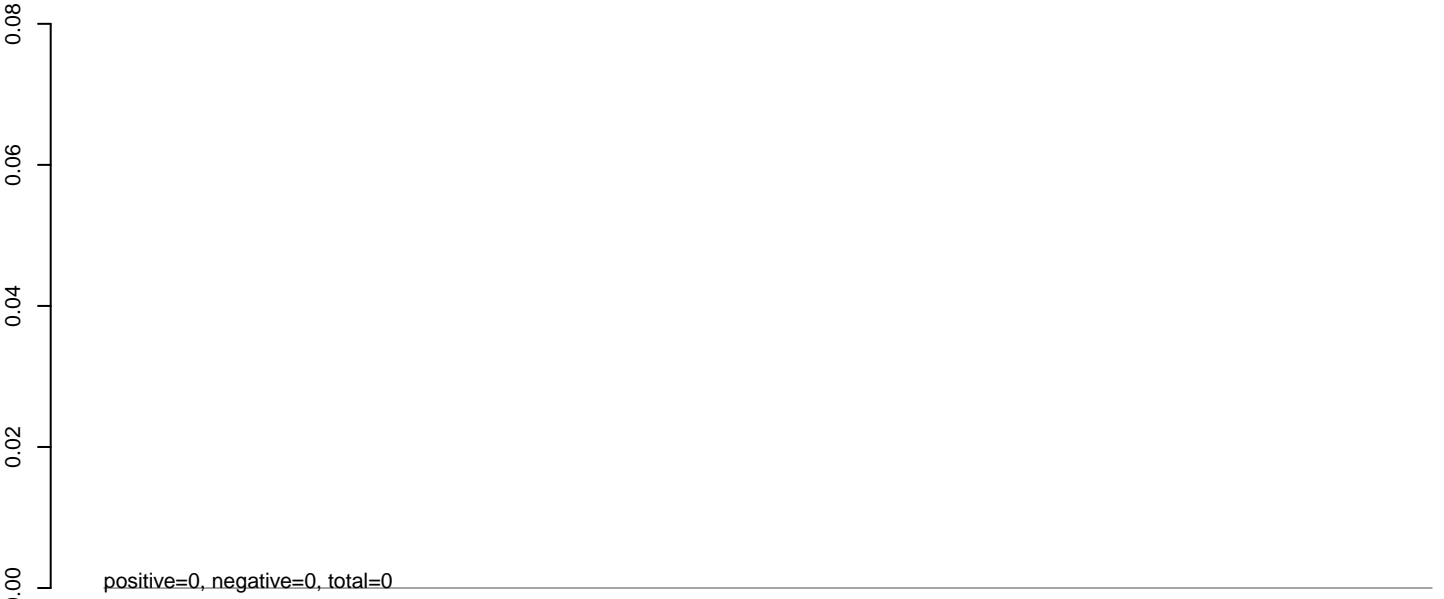


positive=0, negative=1, total=1

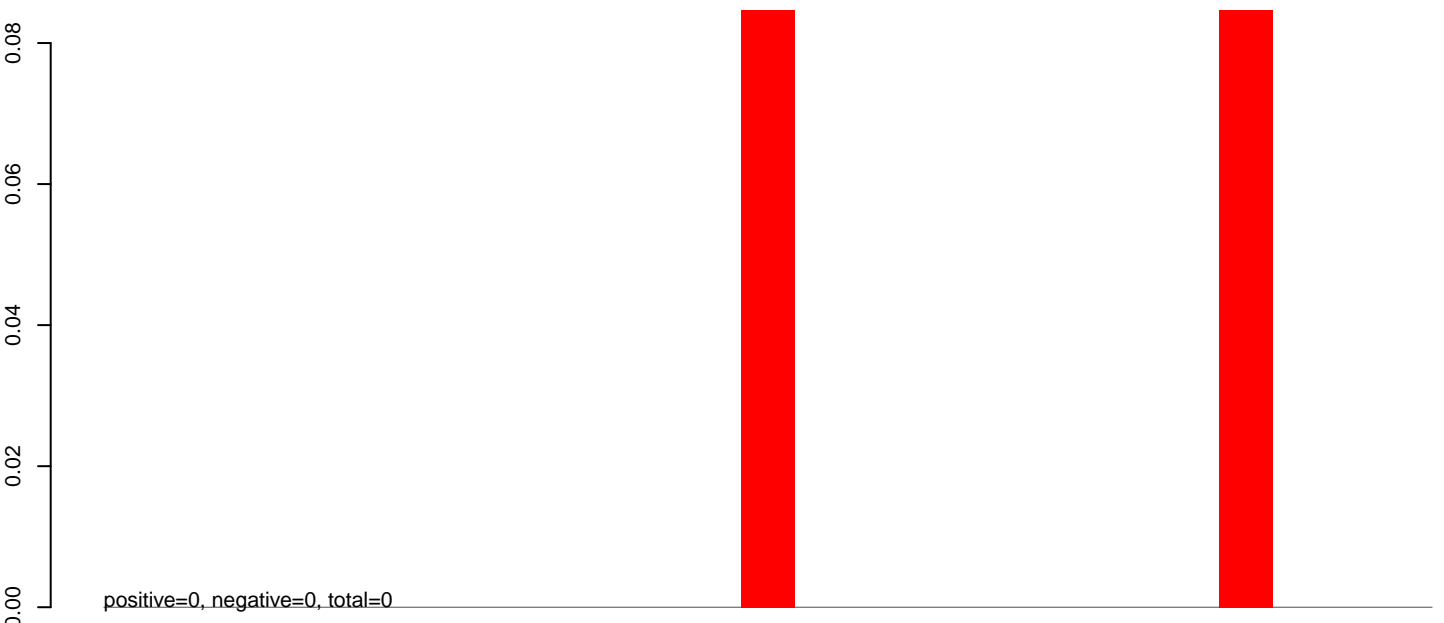
Window size=50, length=447, TE@EAL40760-Unknown:1-447

100 200 300 400

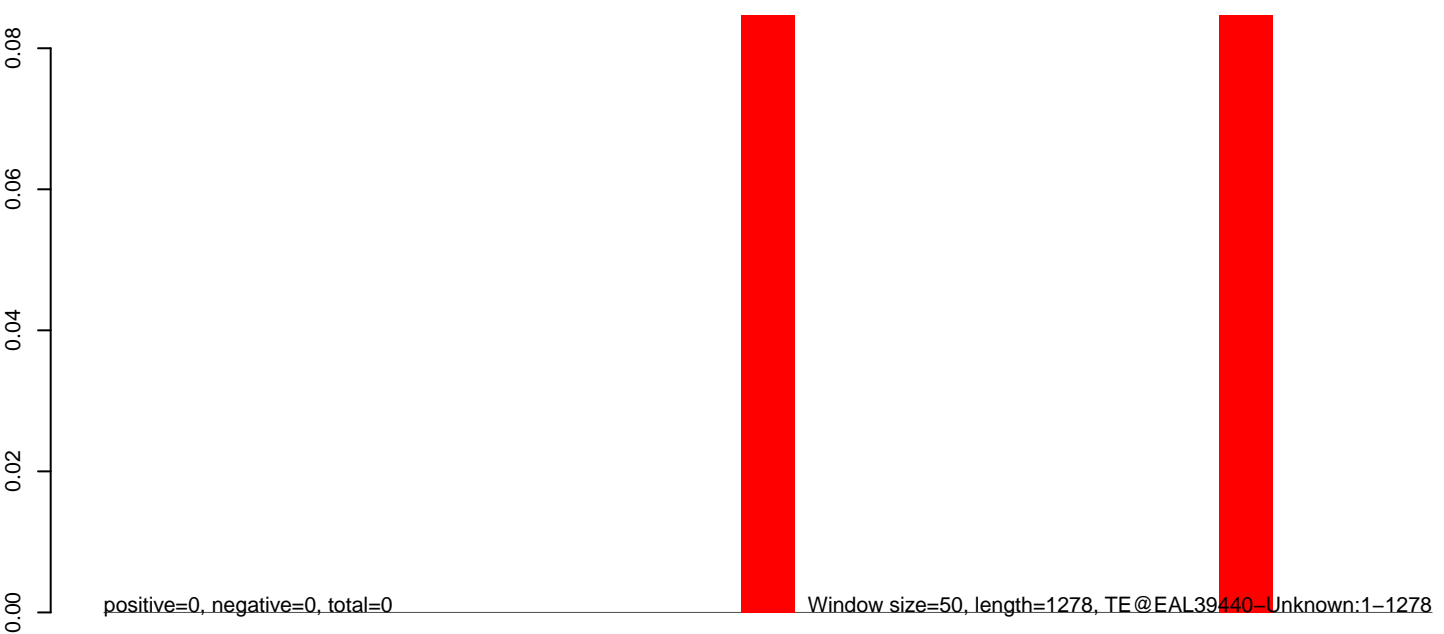
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



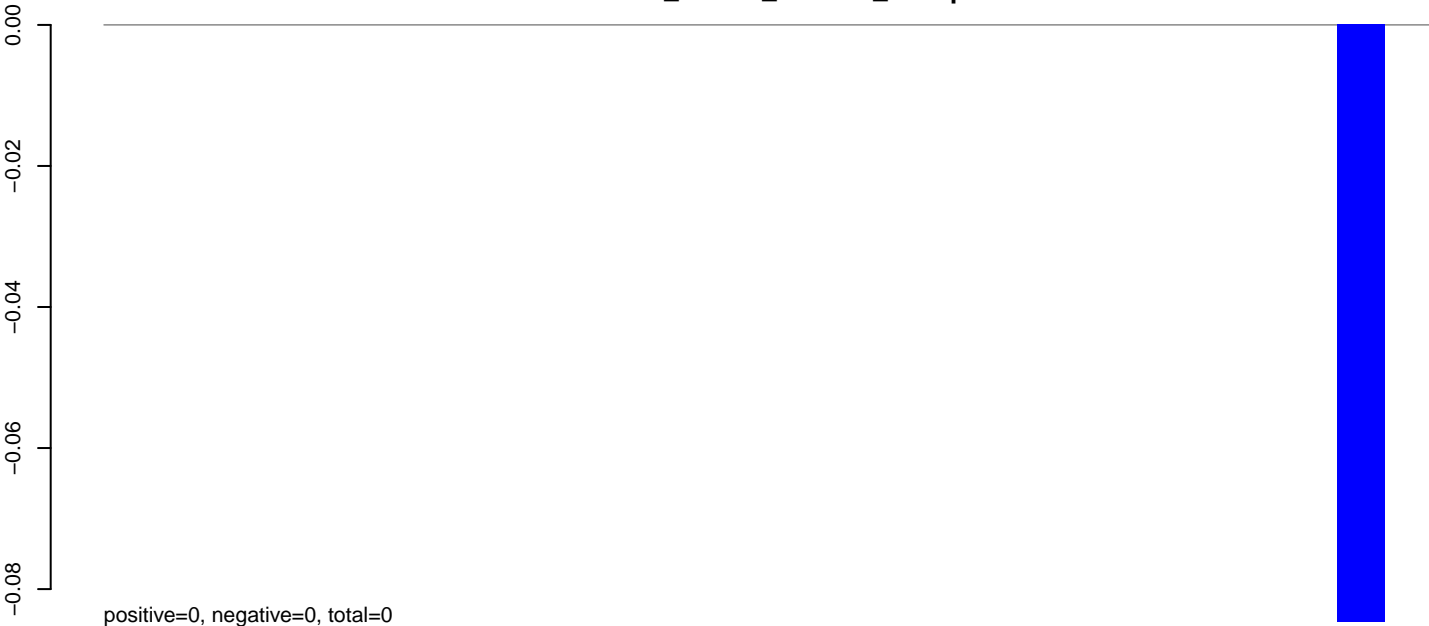
AnGam_Mos55_cells.rep



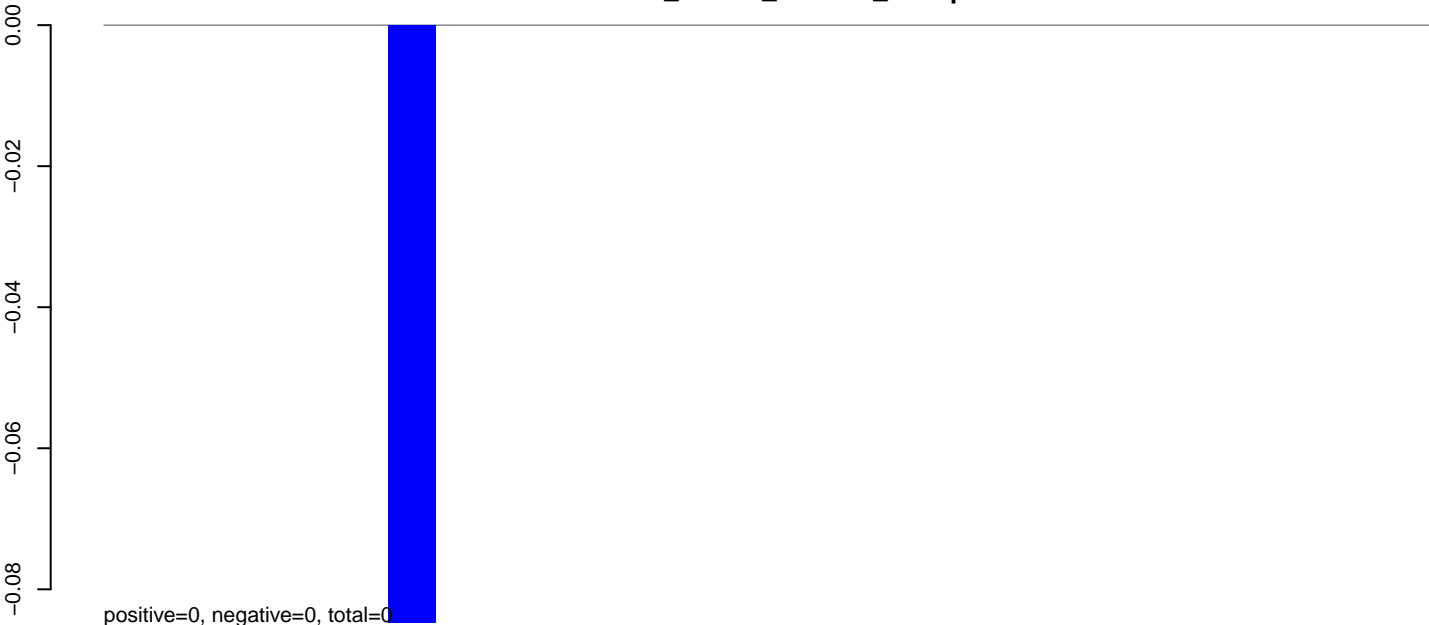
200 400 600 800 1000 1200

Window size=50, length=1278, TE@EAL39440-Unknown:1-1278

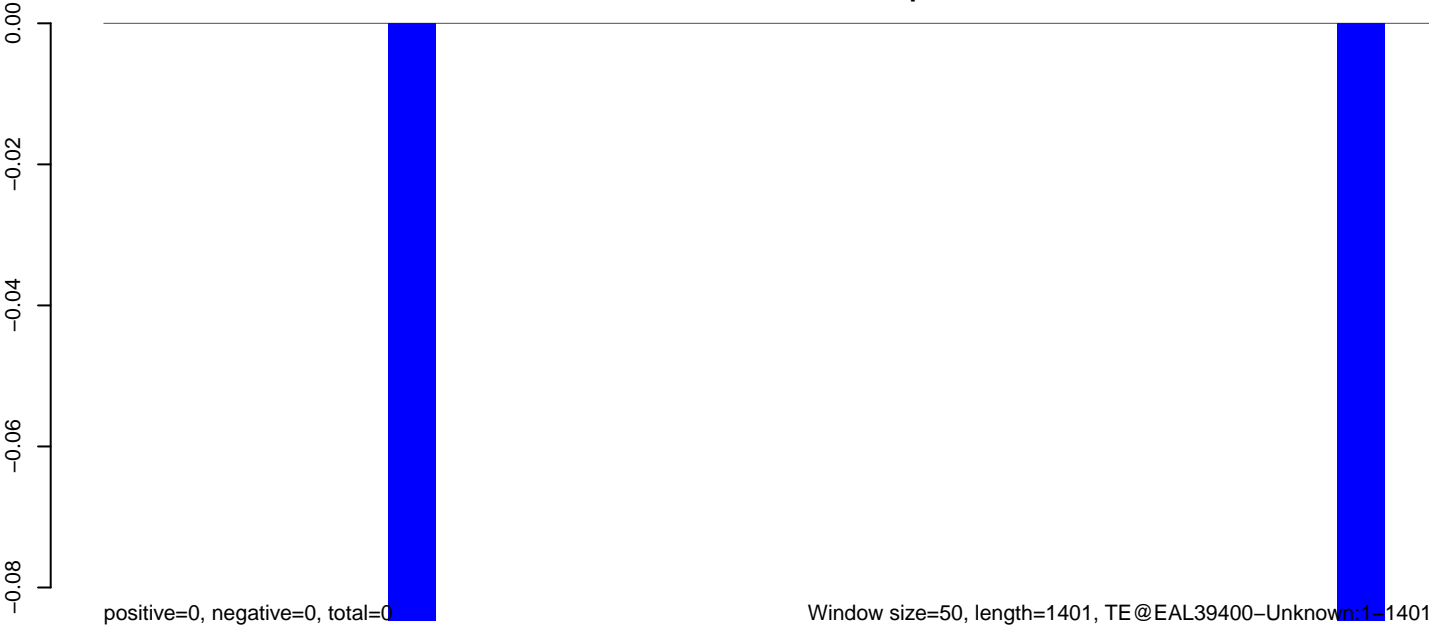
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

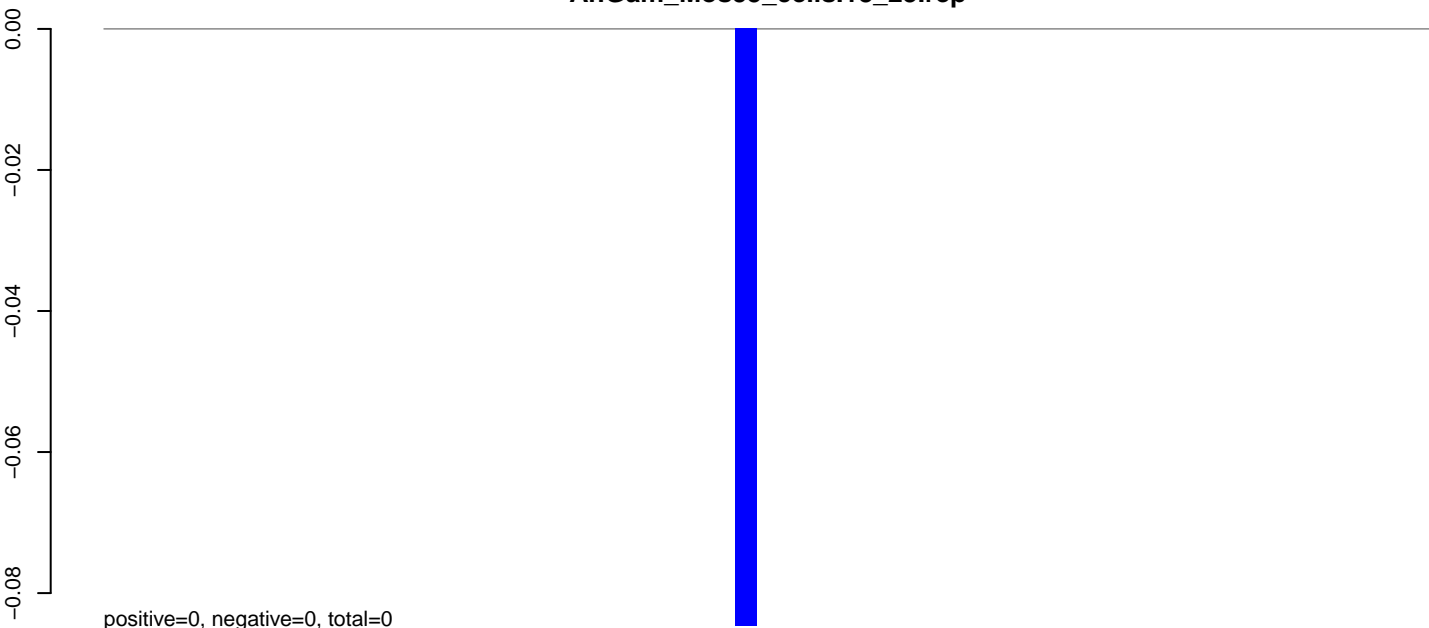


AnGam_Mos55_cells.rep

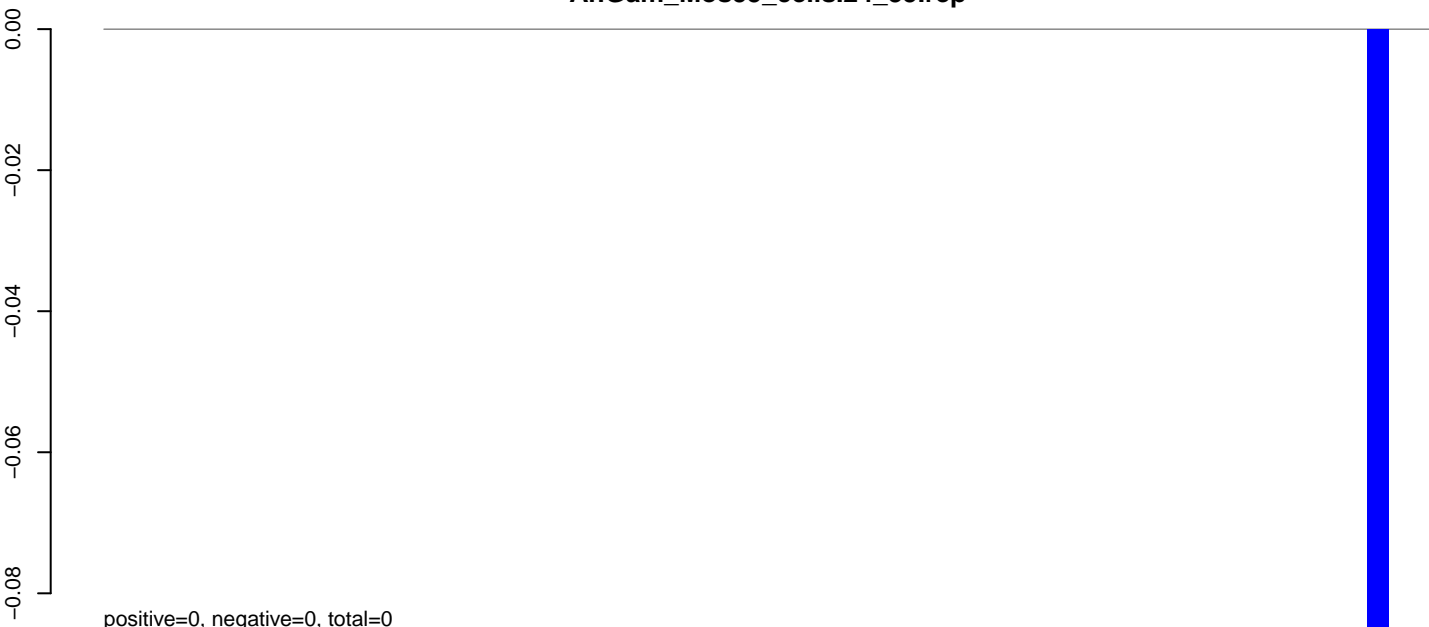


0 200 400 600 800 1000 1200 1400

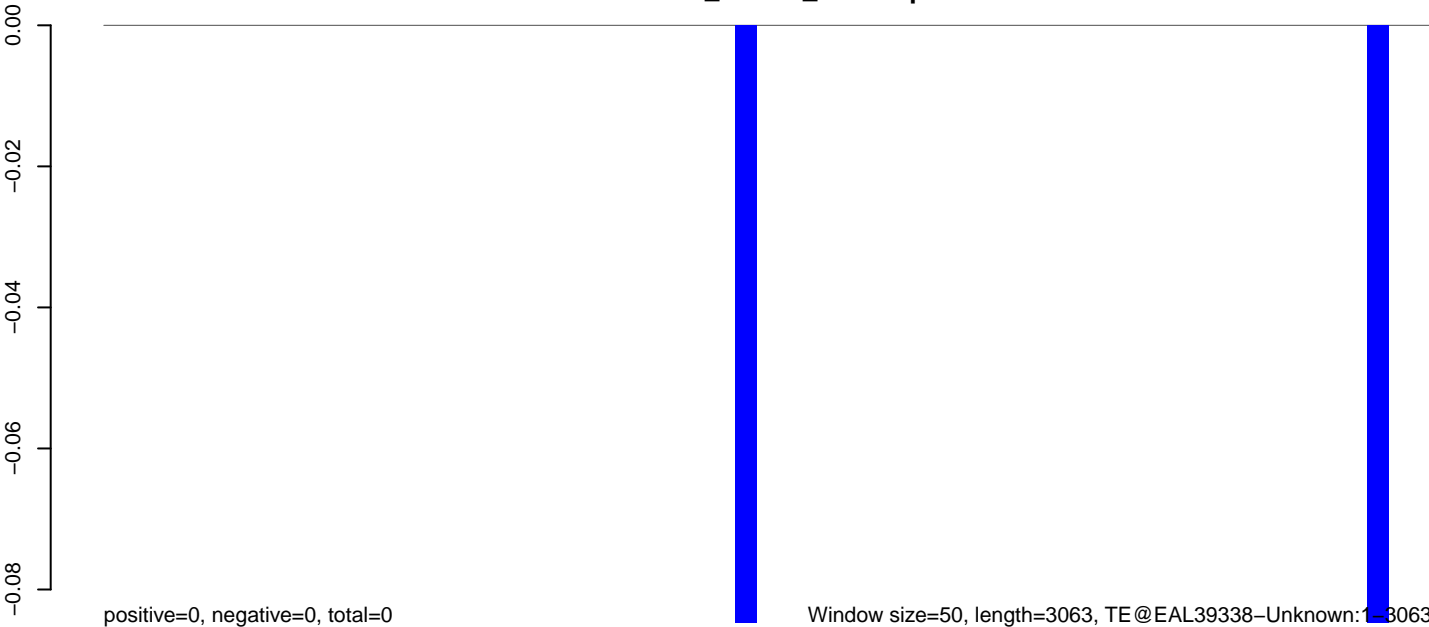
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



0 500 1000 1500 2000 2500 3000

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=897, TE@EAL39174-Unknown:1-897

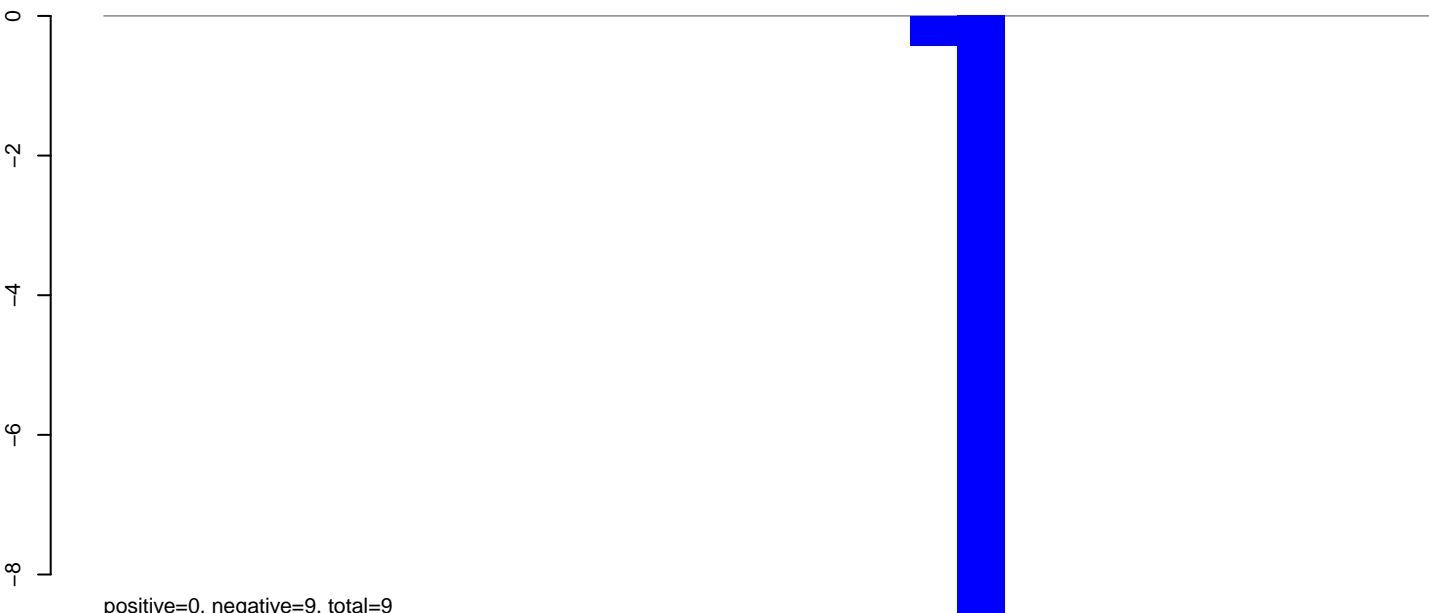
200

400

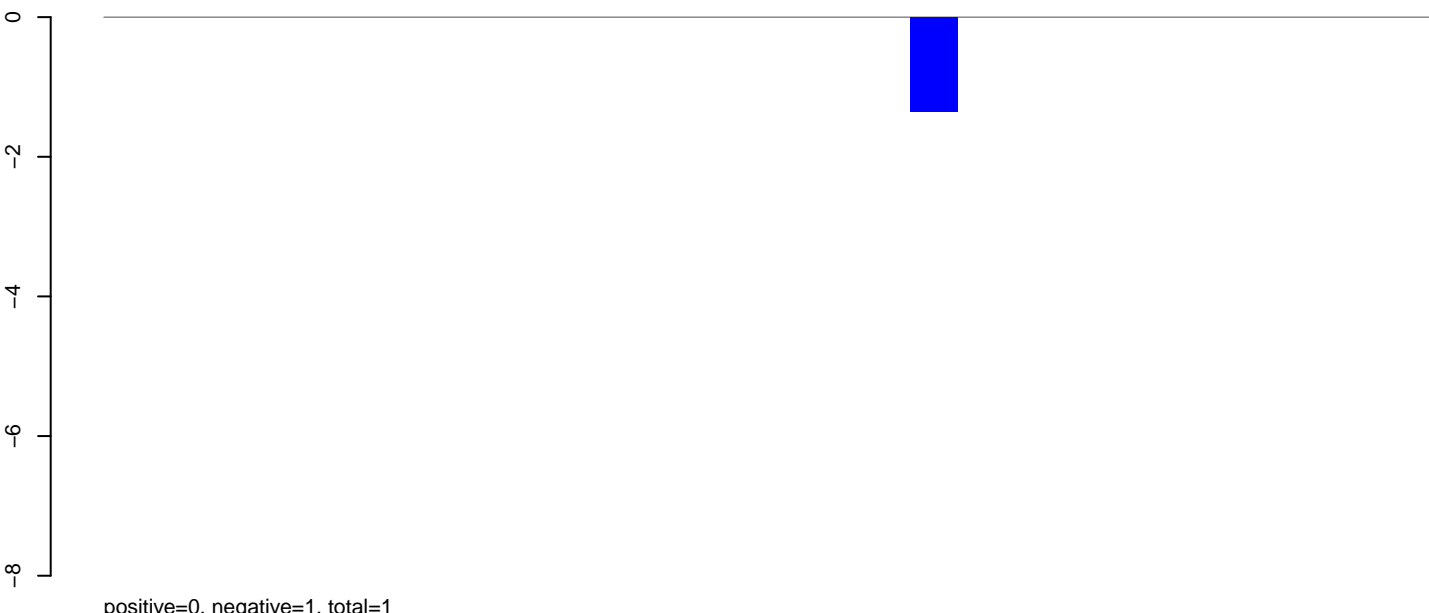
600

800

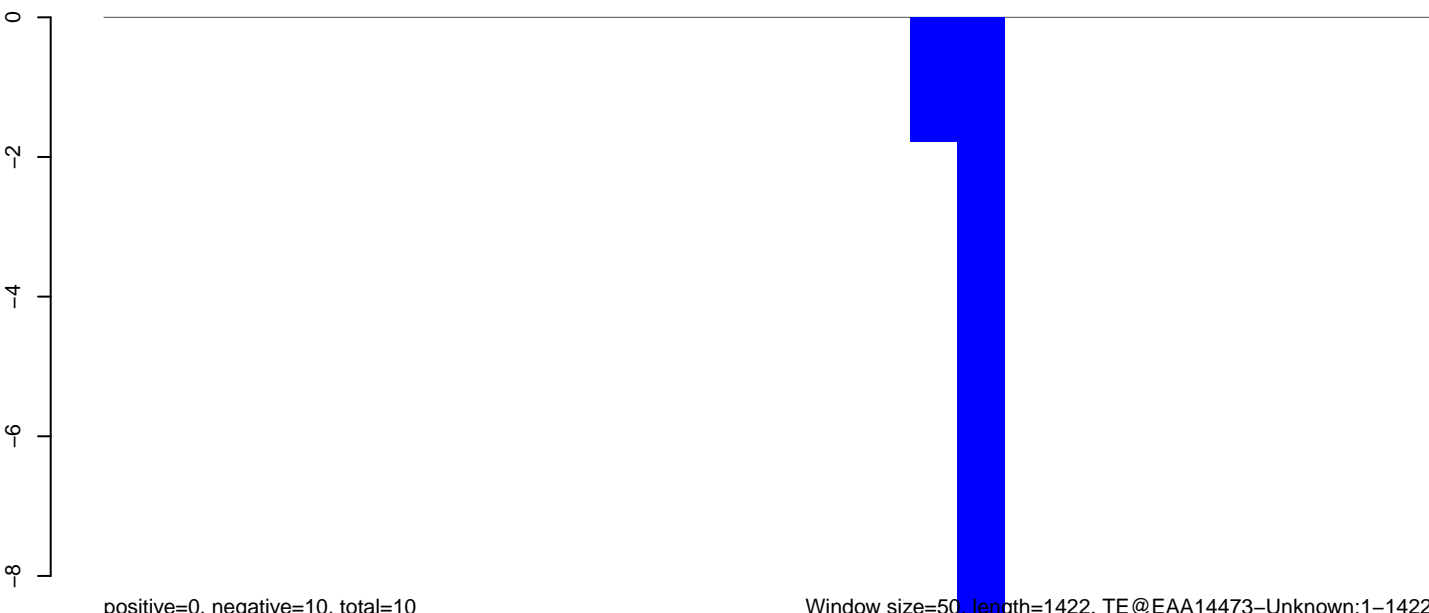
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep

0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

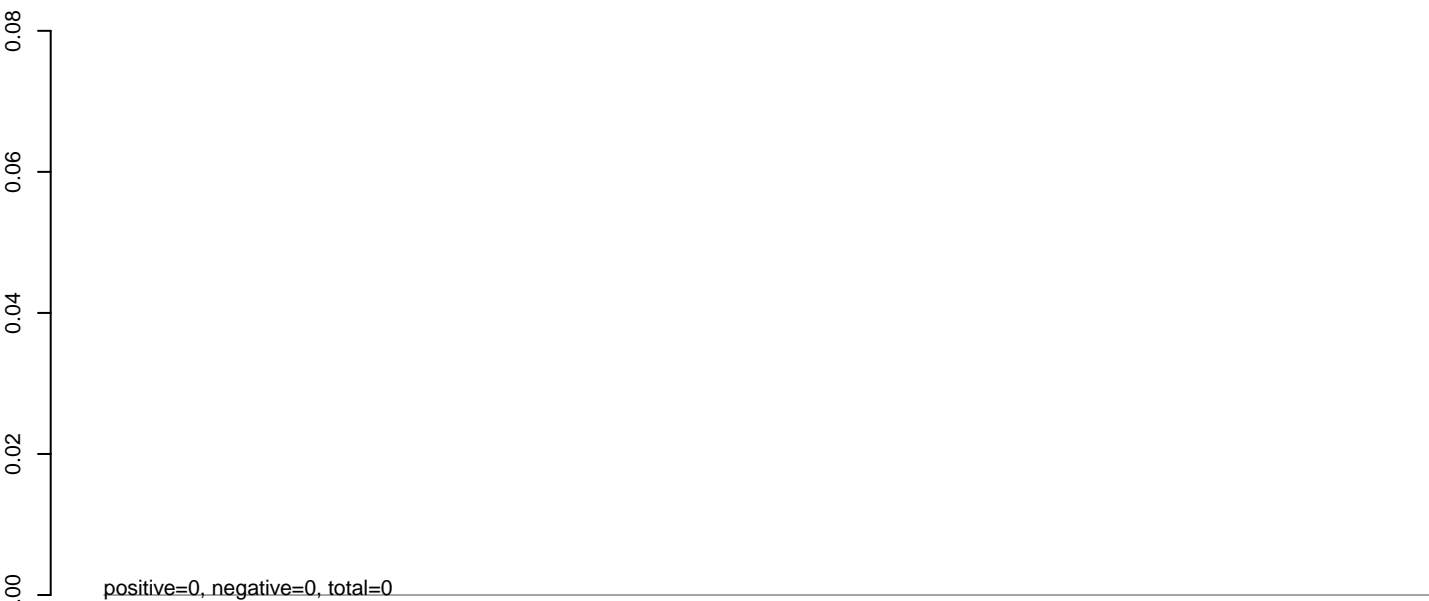
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positive=0, negative=0, total=0

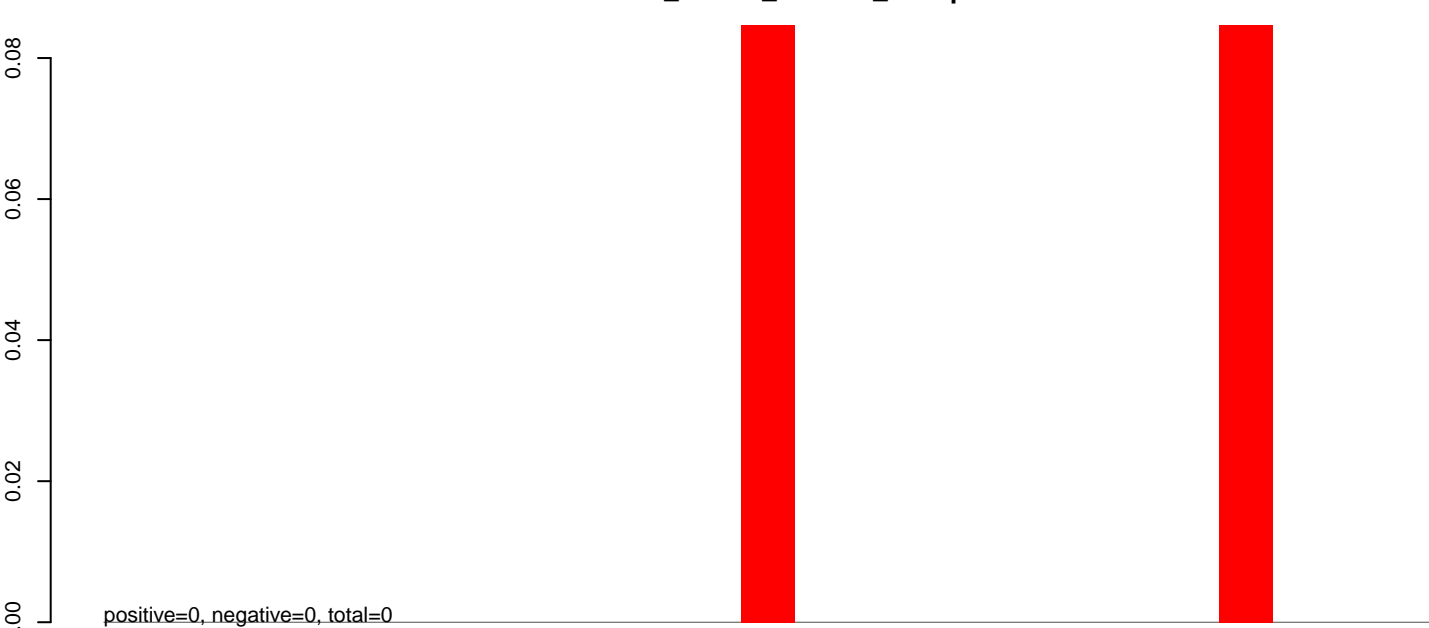
Window size=50, length=807, TE@EAA09884-Unknown:1-807

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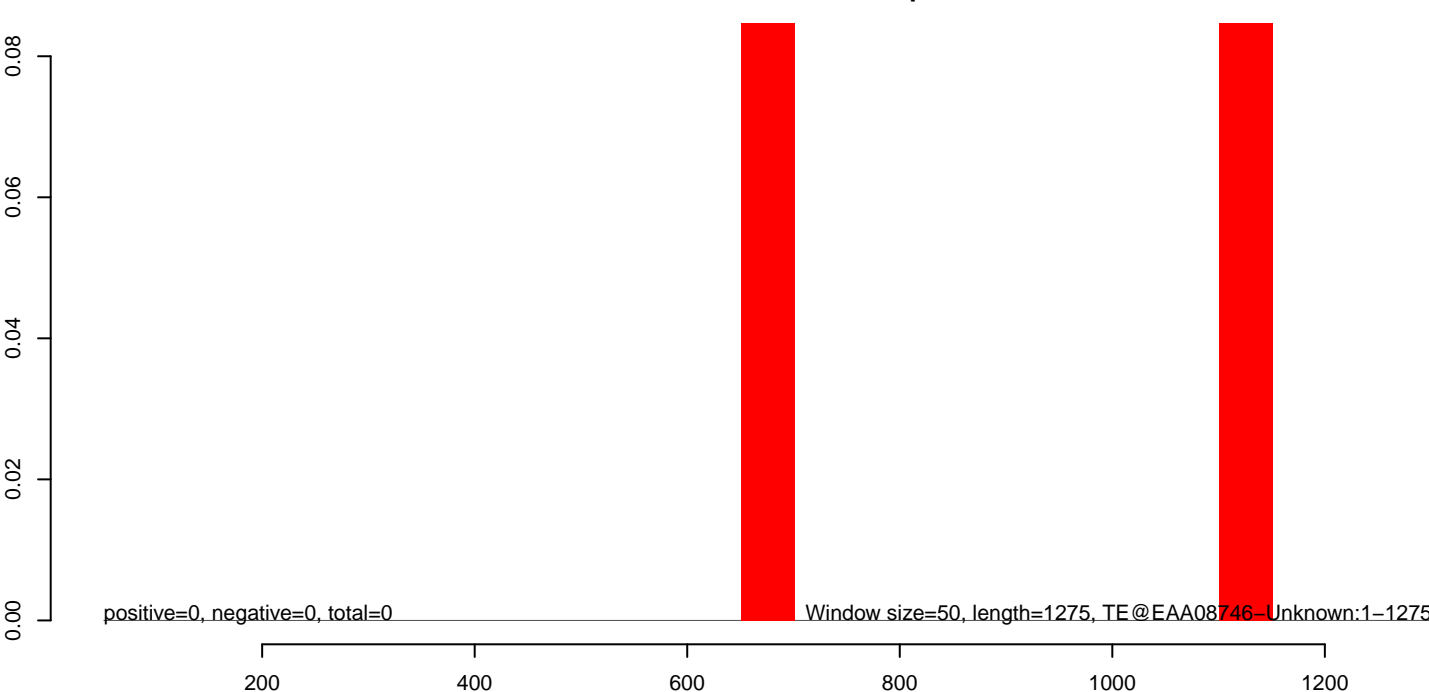
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

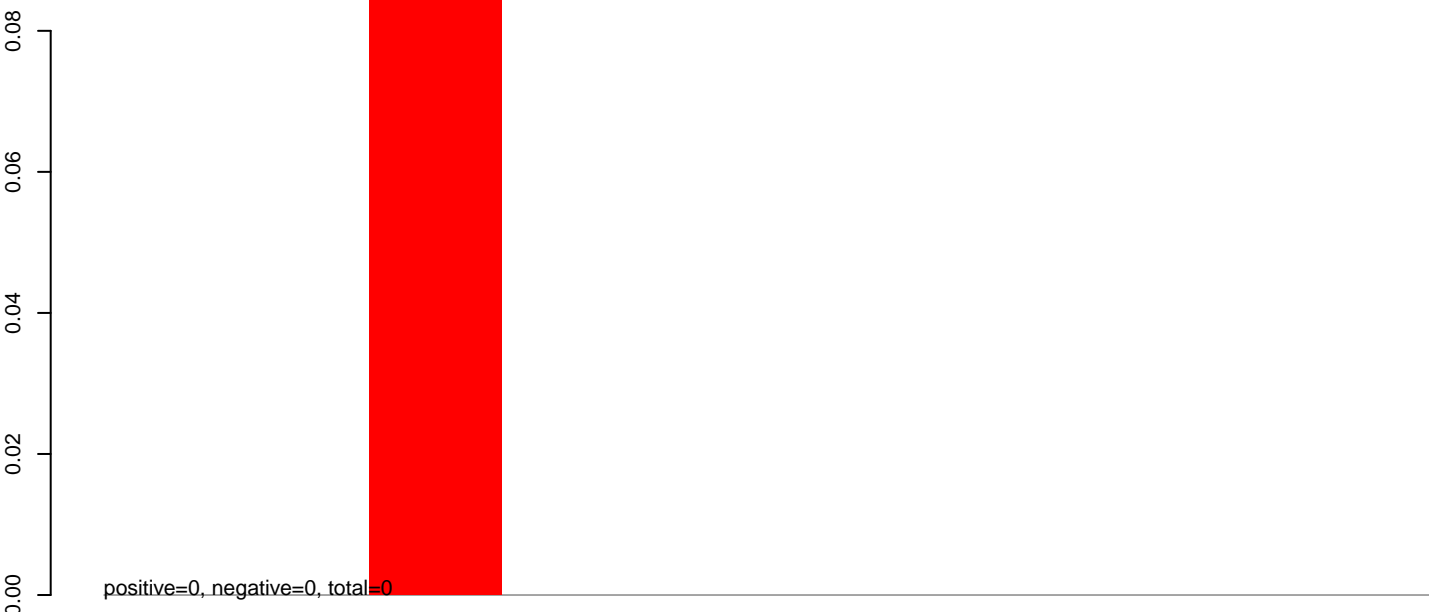
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-0.06
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positive=0, negative=0, total=0

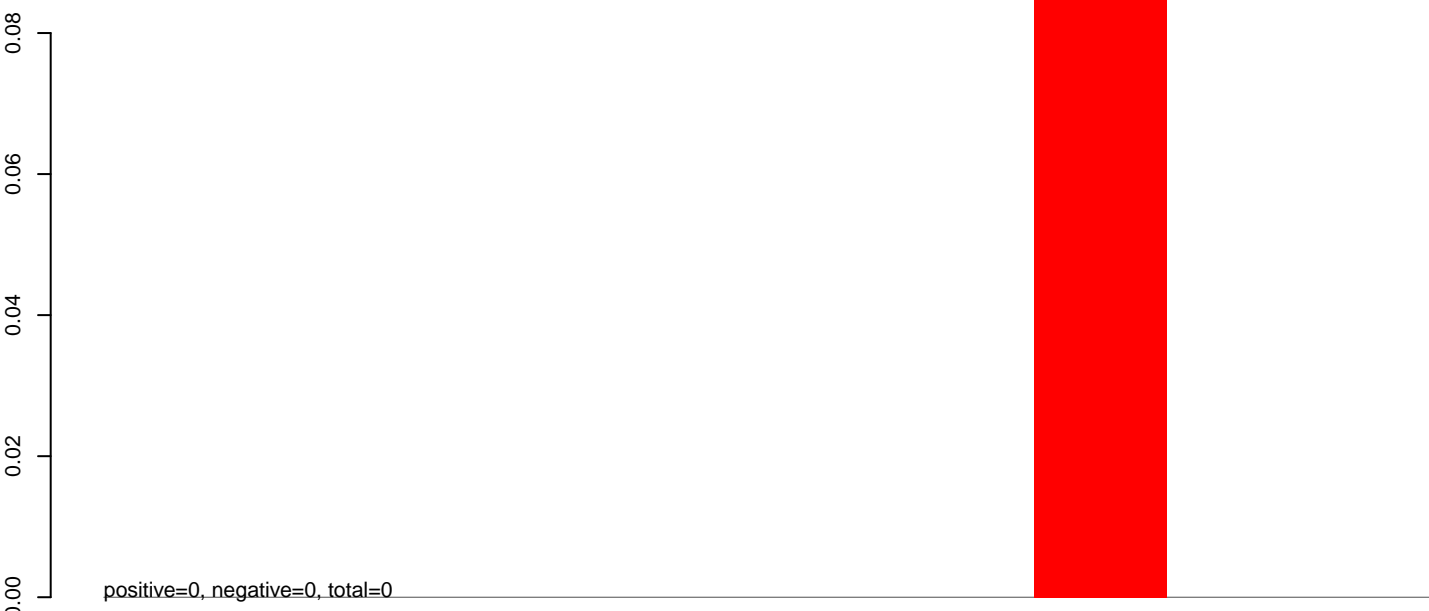
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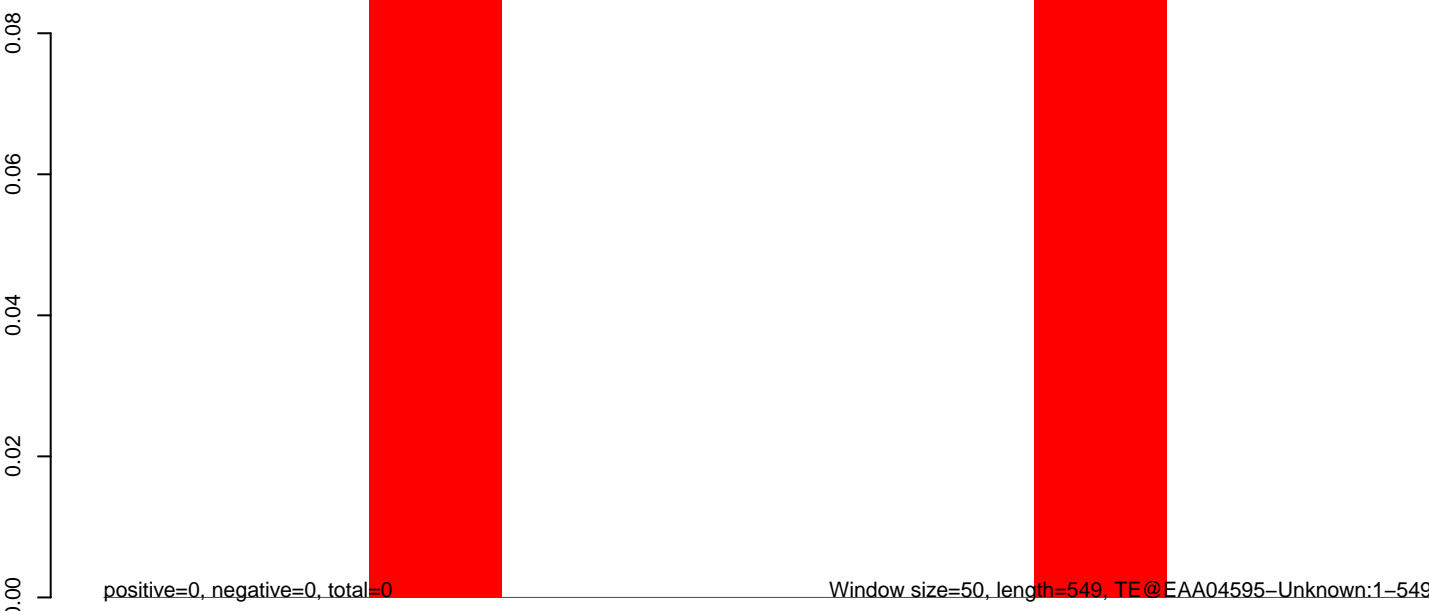
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

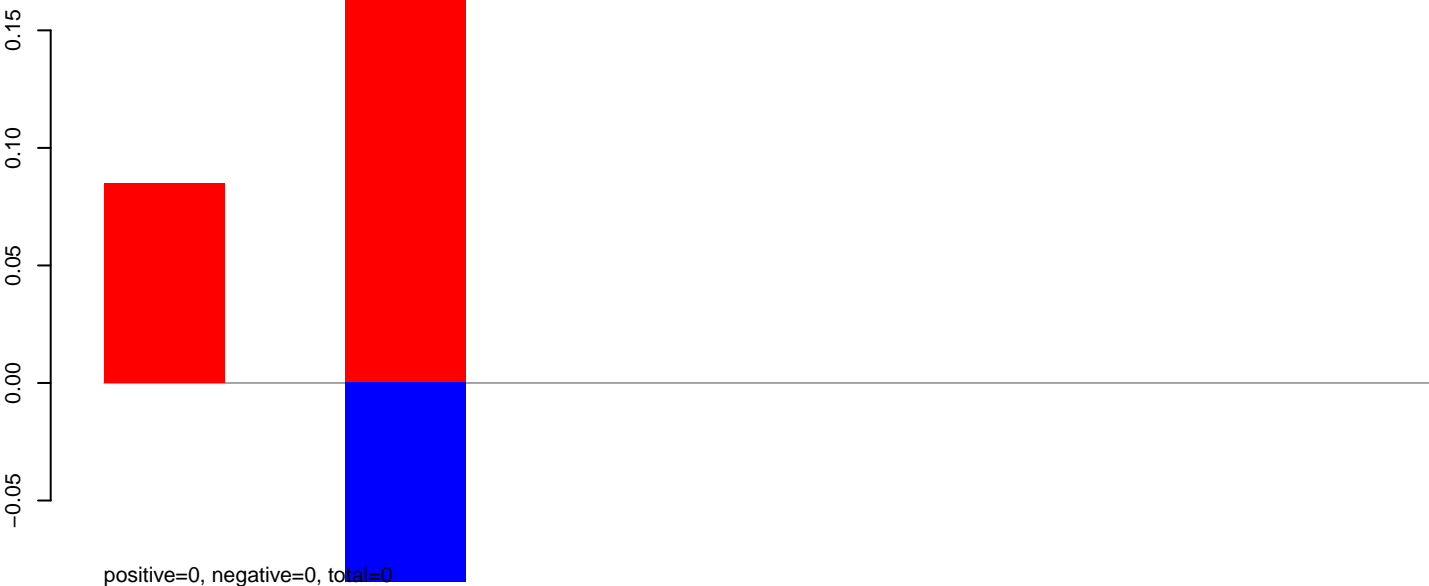


AnGam_Mos55_cells.rep

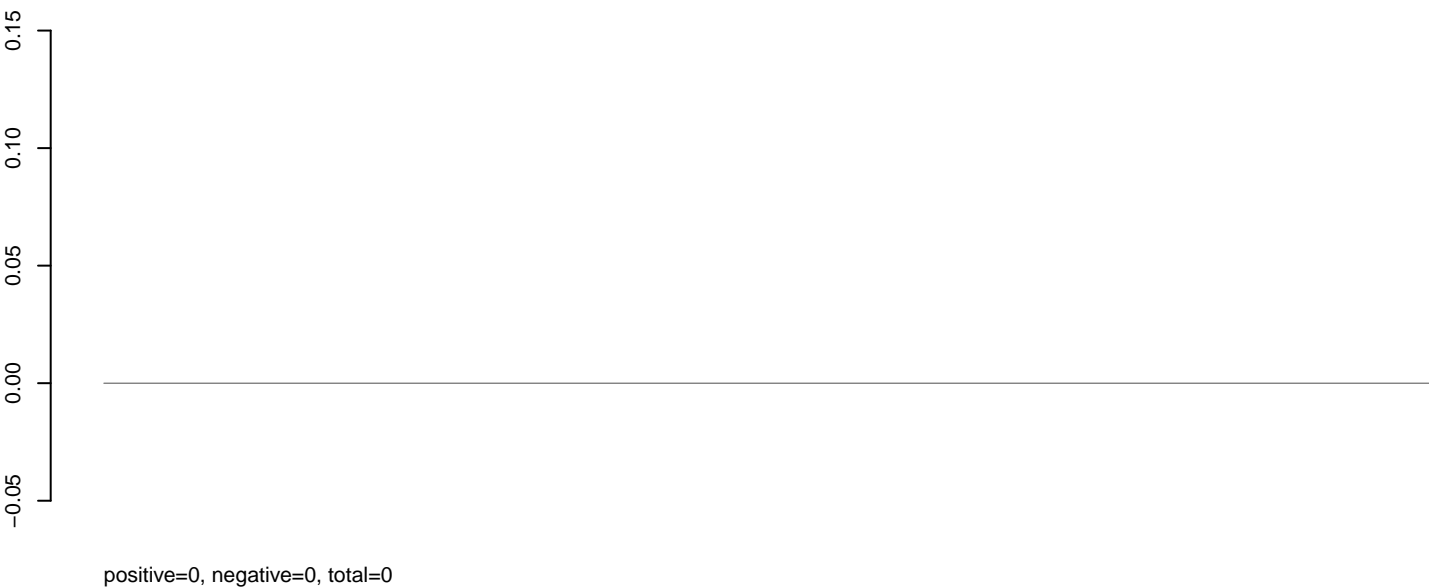


100 200 300 400 500

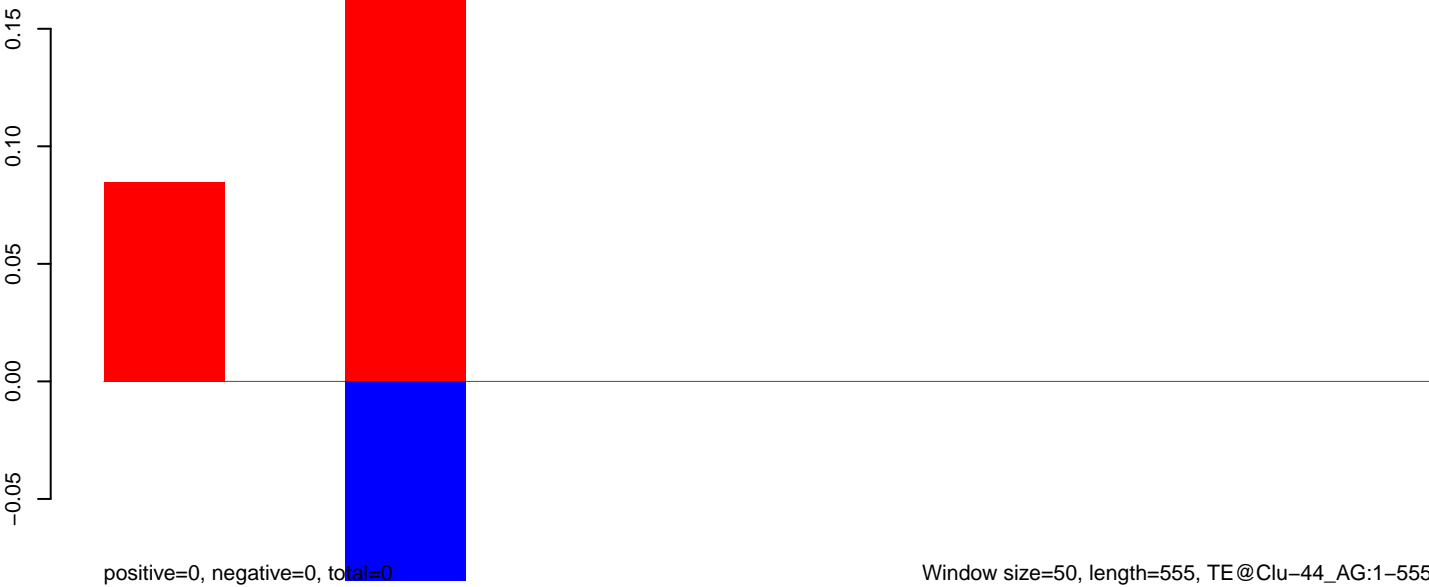
AnGam_Mos55_cells.18_23.rep



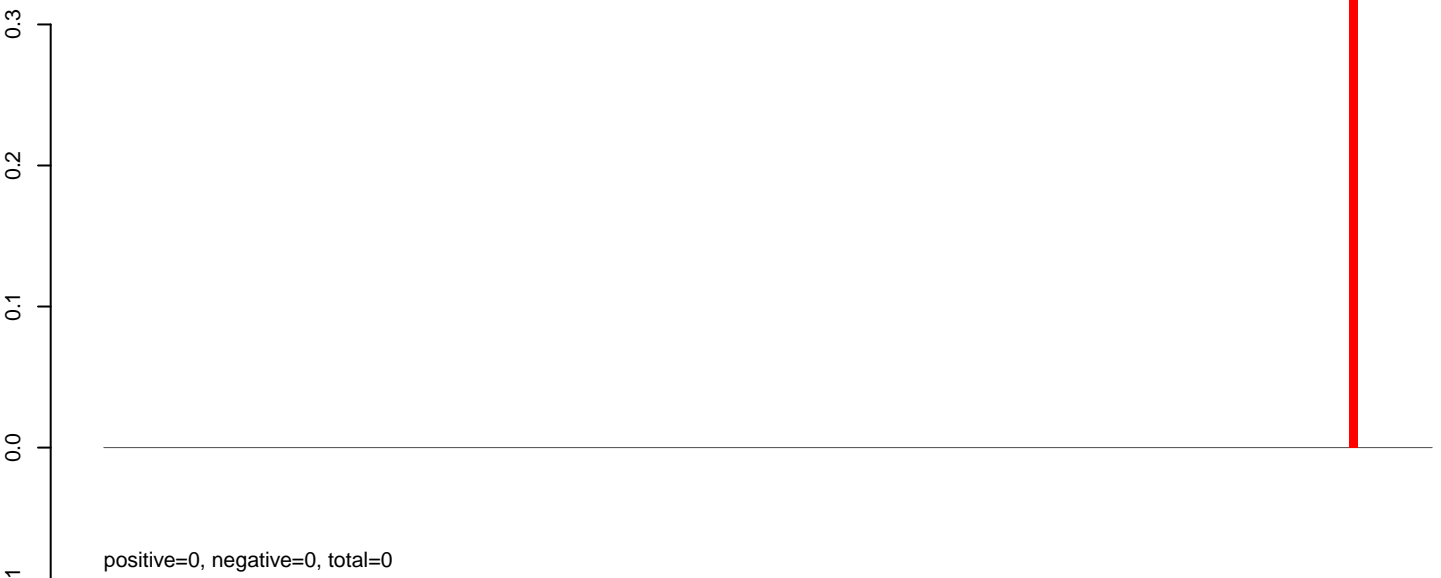
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



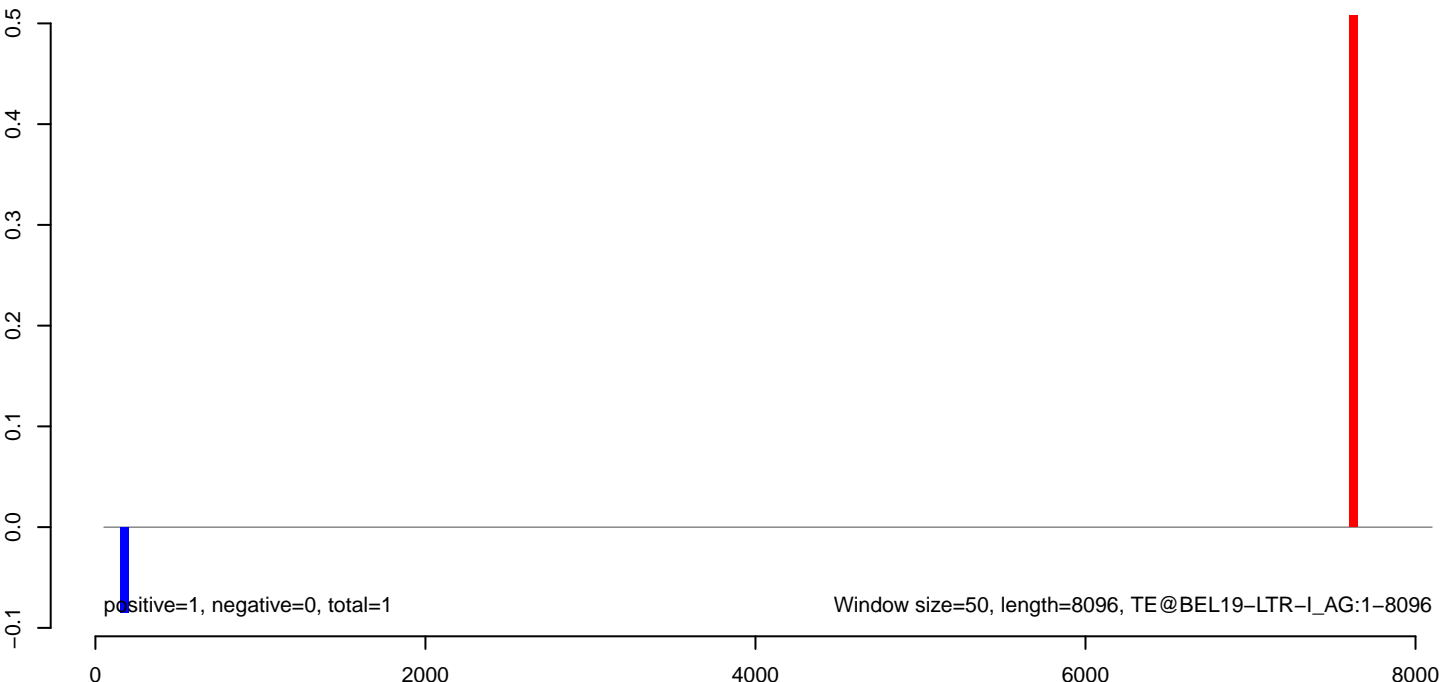
AnGam_Mos55_cells.18_23.rep



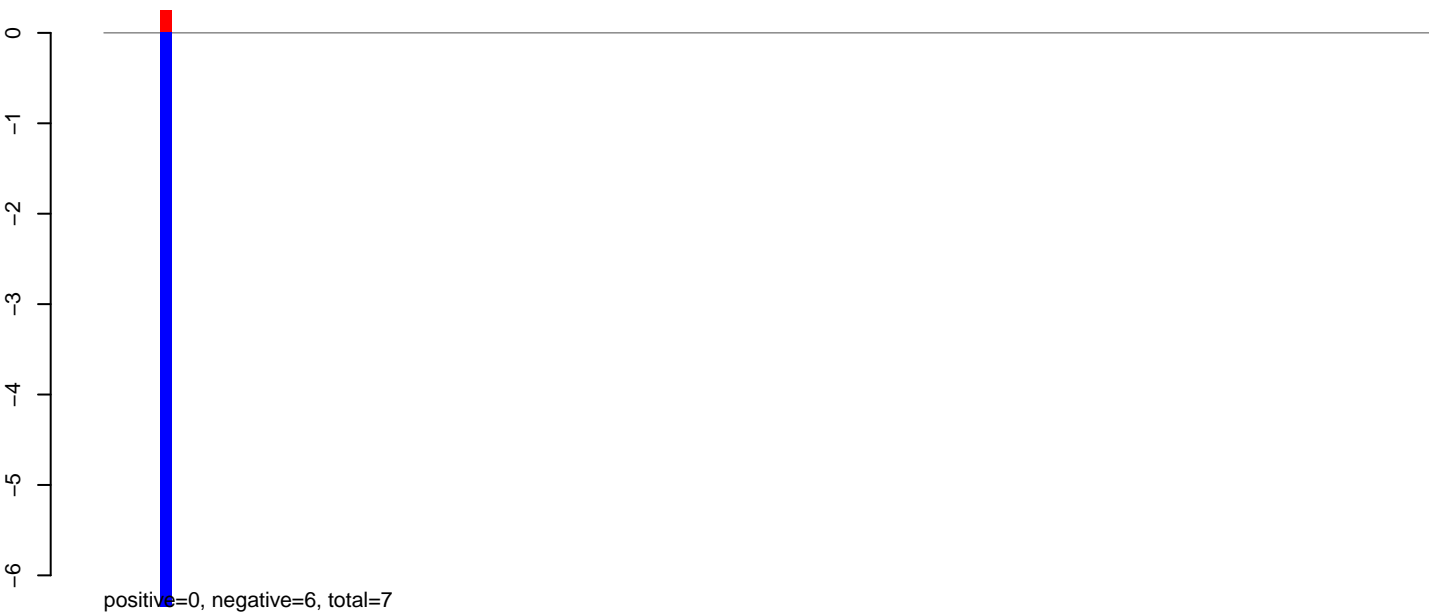
AnGam_Mos55_cells.24_35.rep



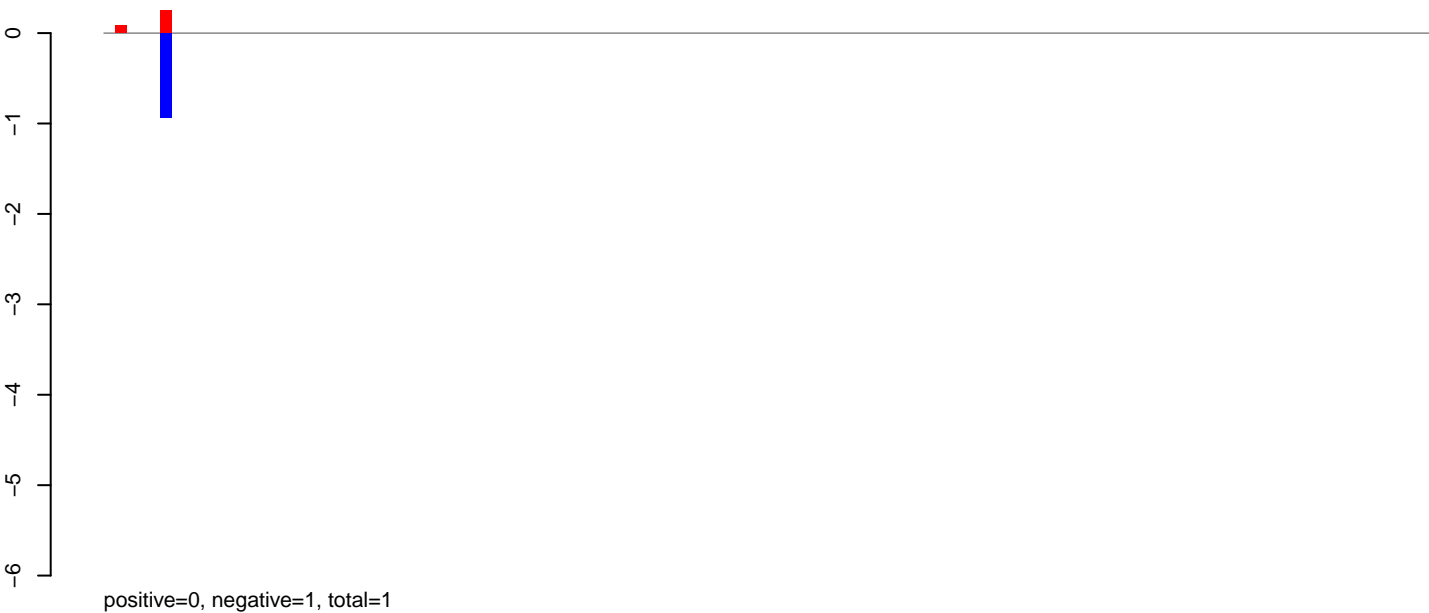
AnGam_Mos55_cells.rep



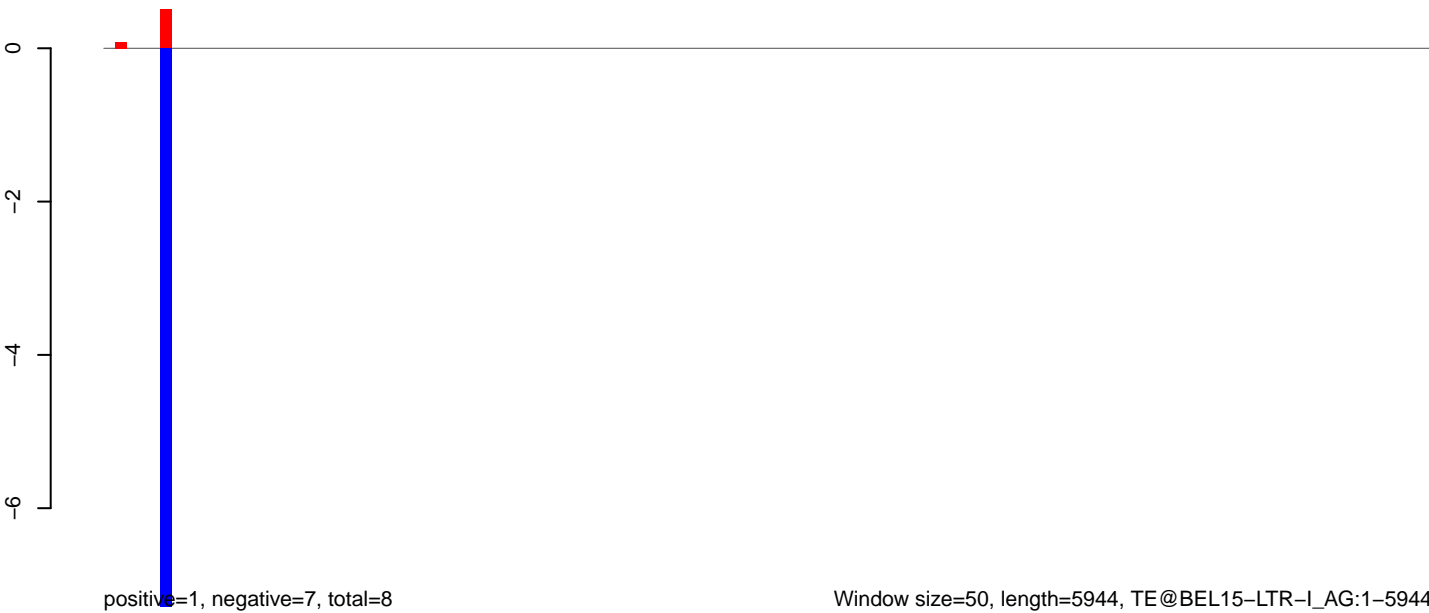
AnGam_Mos55_cells.18_23.rep



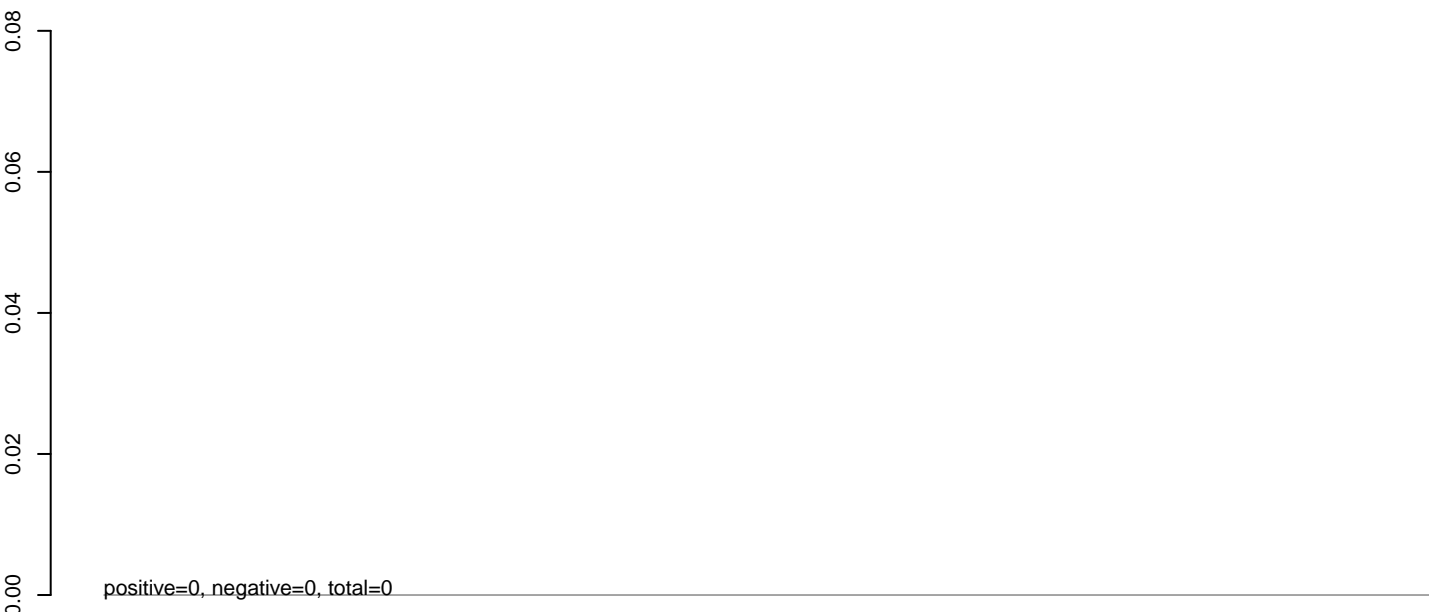
AnGam_Mos55_cells.24_35.rep



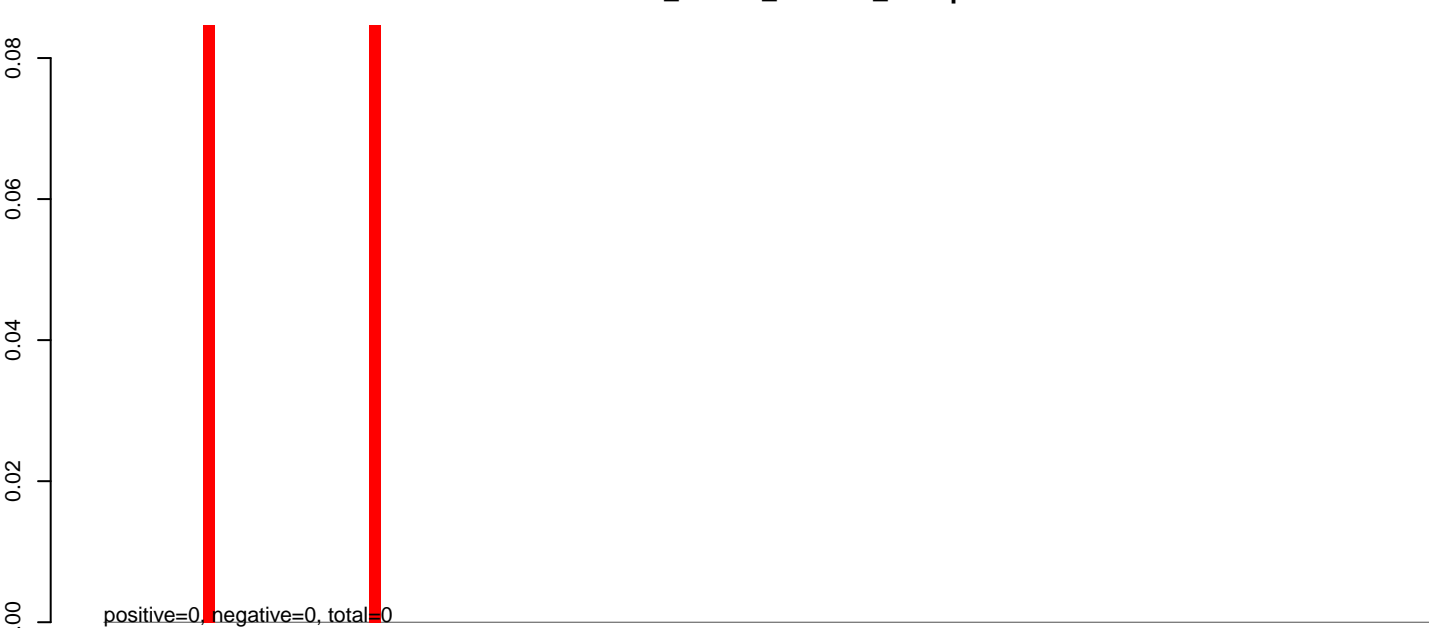
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



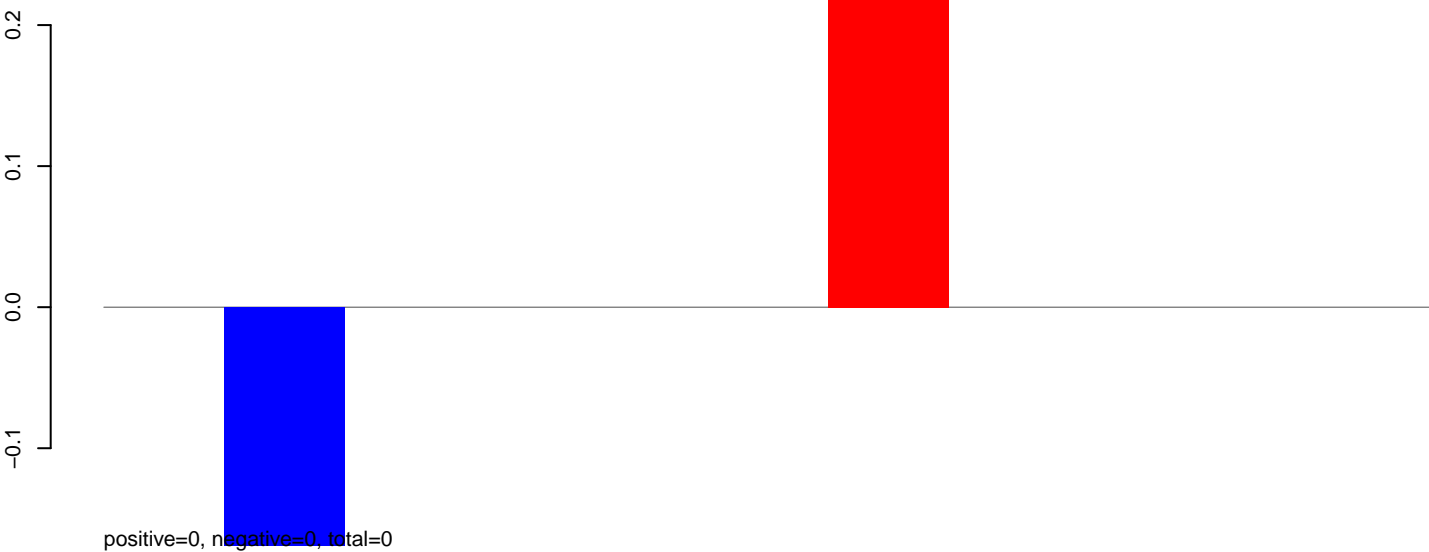
AnGam_Mos55_cells.rep



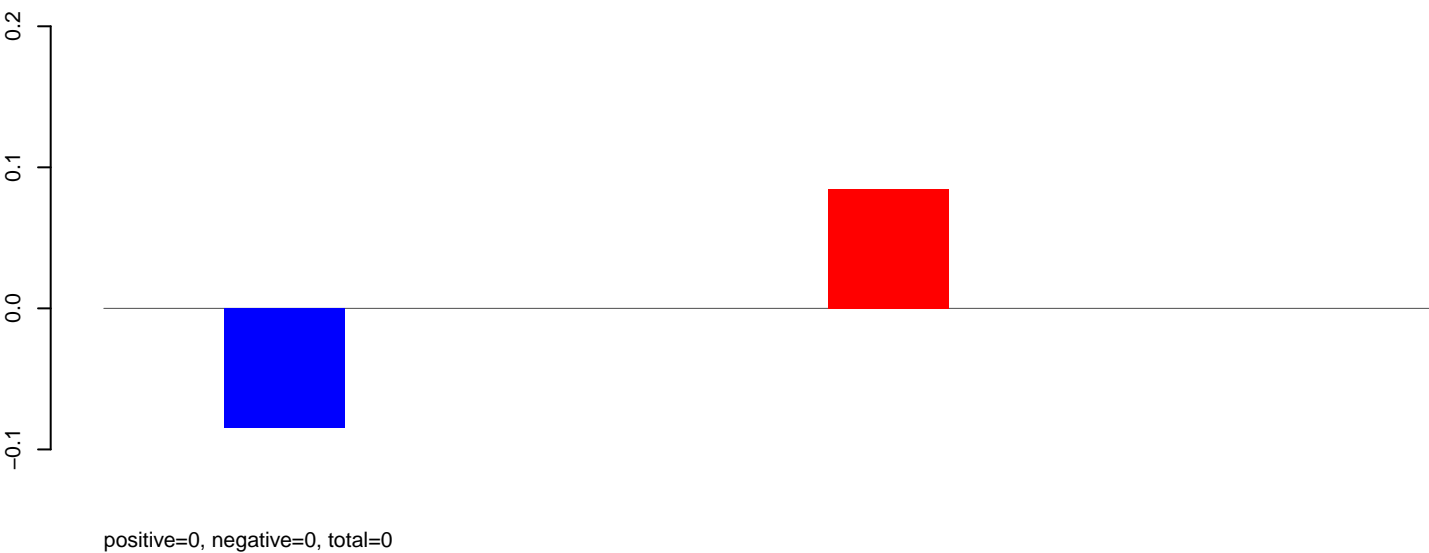
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0 1000 2000 3000 4000 5000 6000

AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



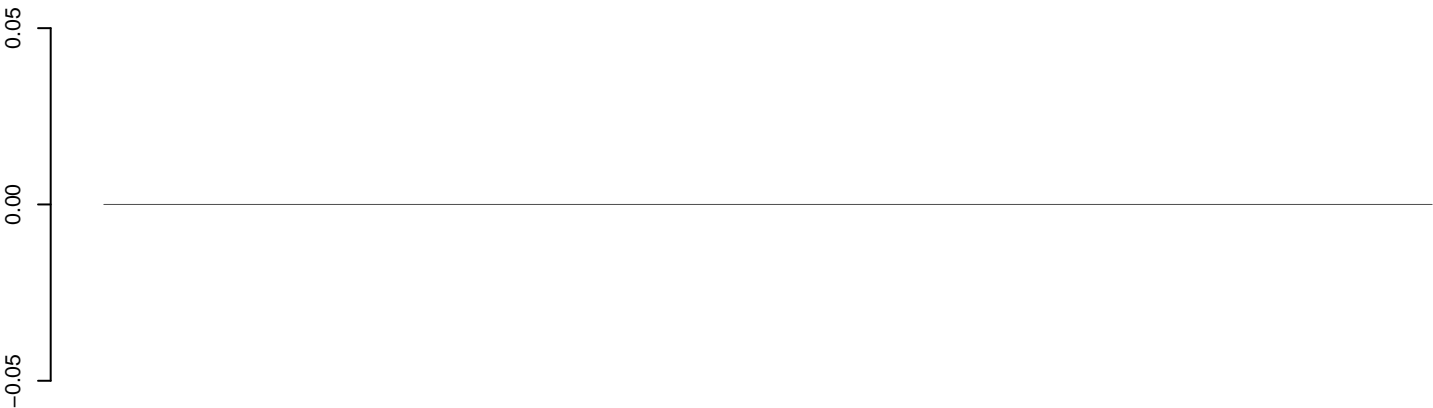
AnGam_Mos55_cells.rep



Window size=50, length=559, TE@AgaP4MITE559:1-559

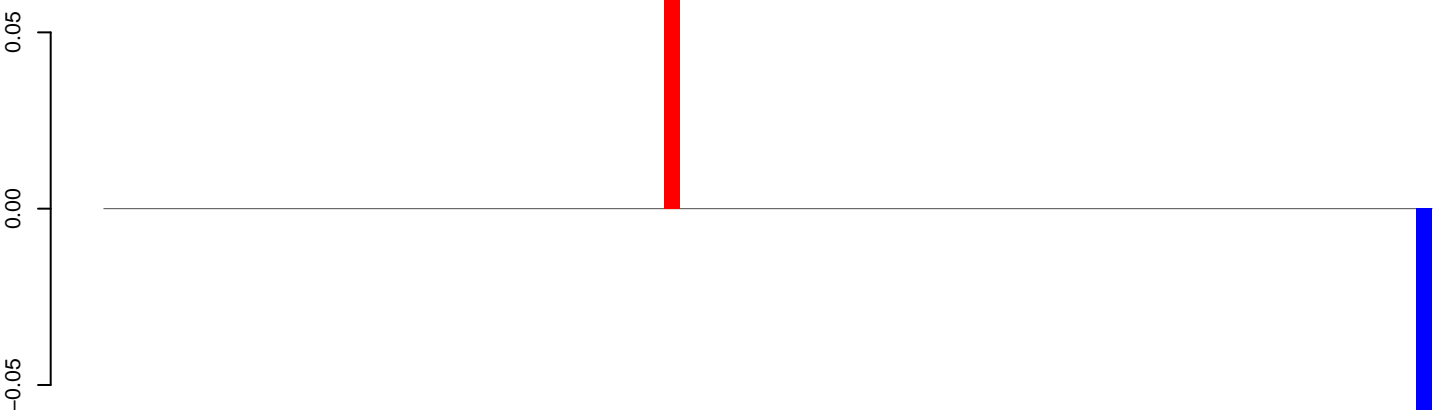
100 200 300 400 500 600

AnGam_Mos55_cells.18_23.rep



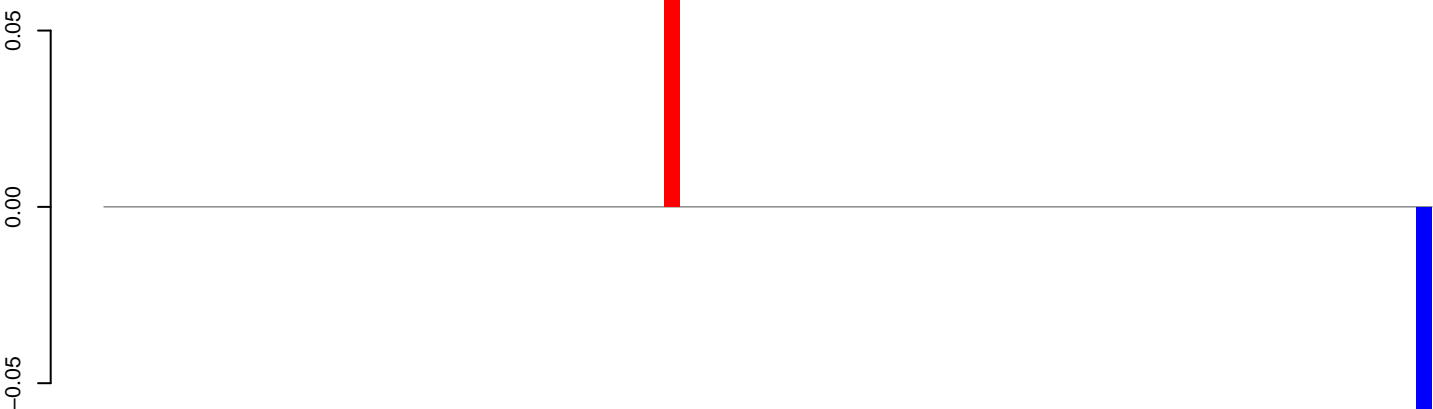
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



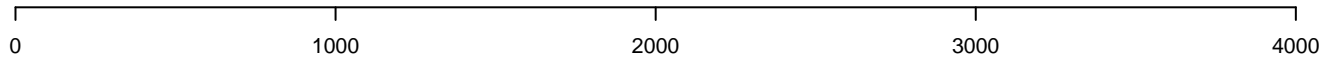
positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

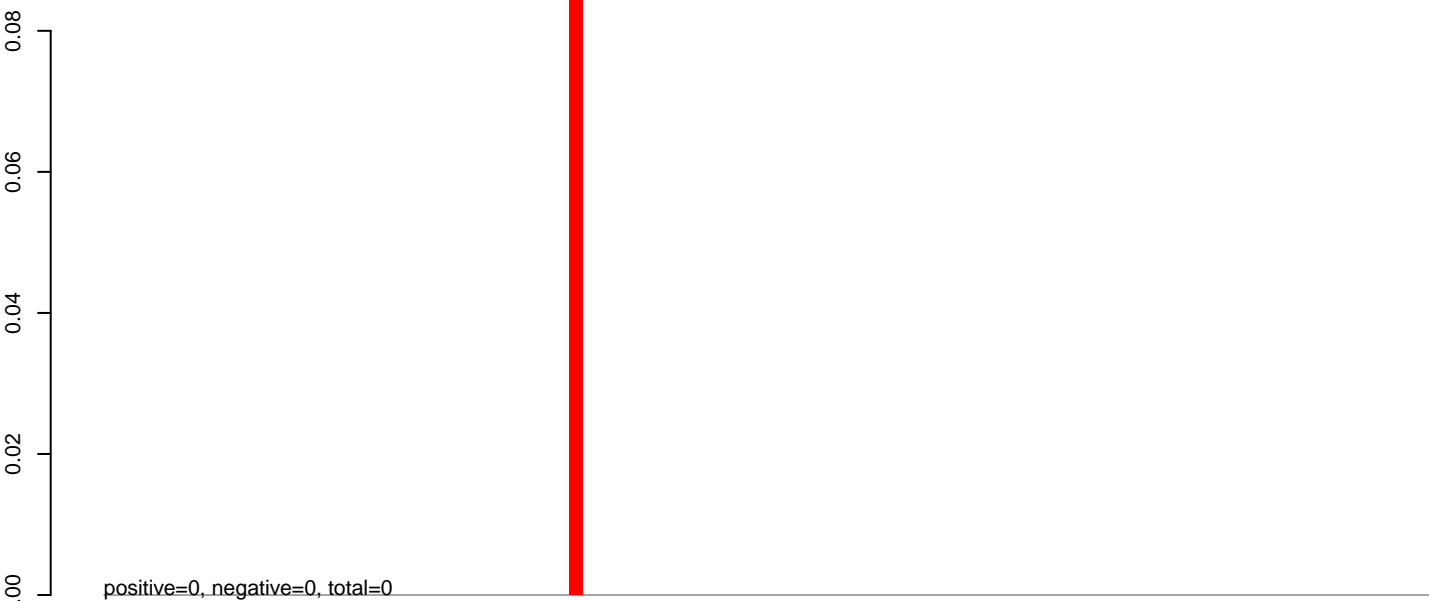


positive=0, negative=0, total=0

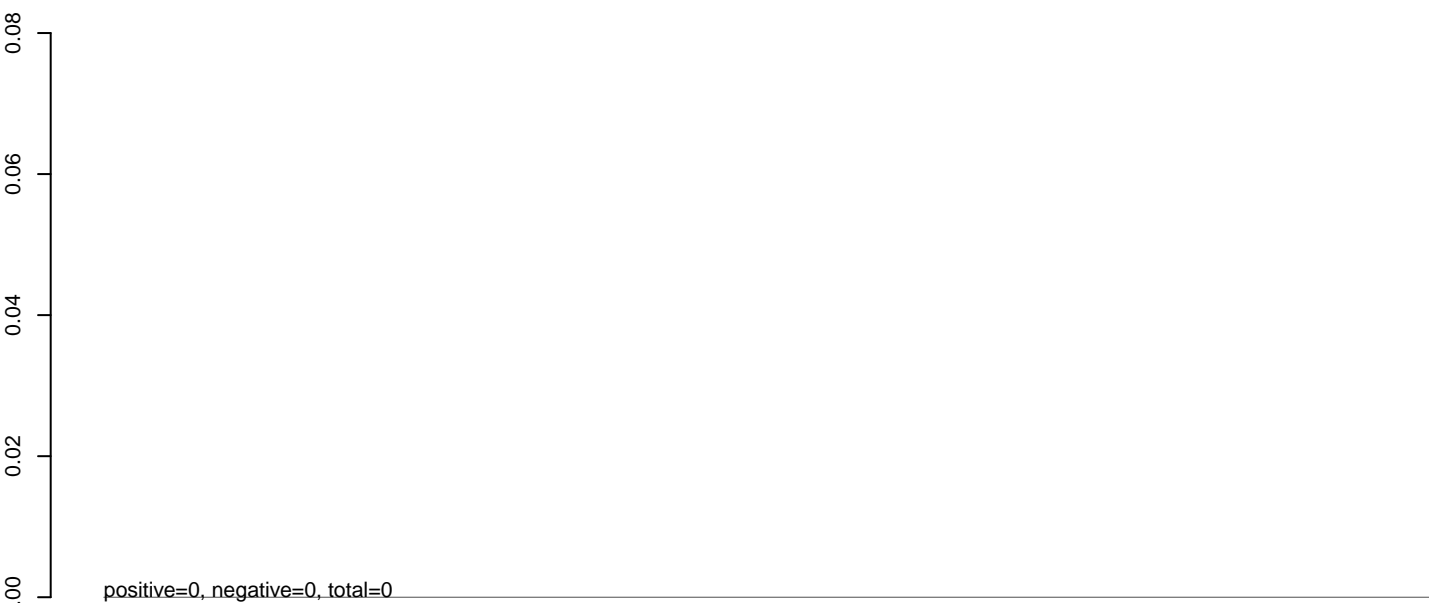
Window size=50, length=4178, TE@Ag-Jen-11:1-4178



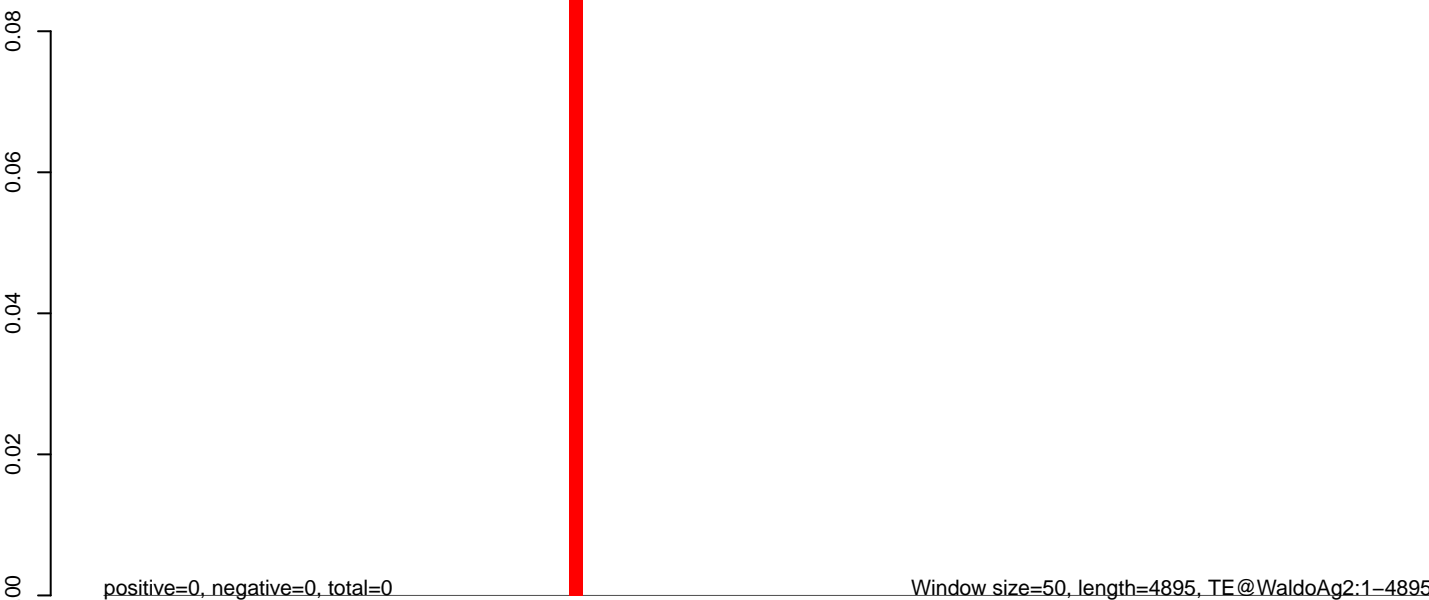
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



0 1000 2000 3000 4000 5000

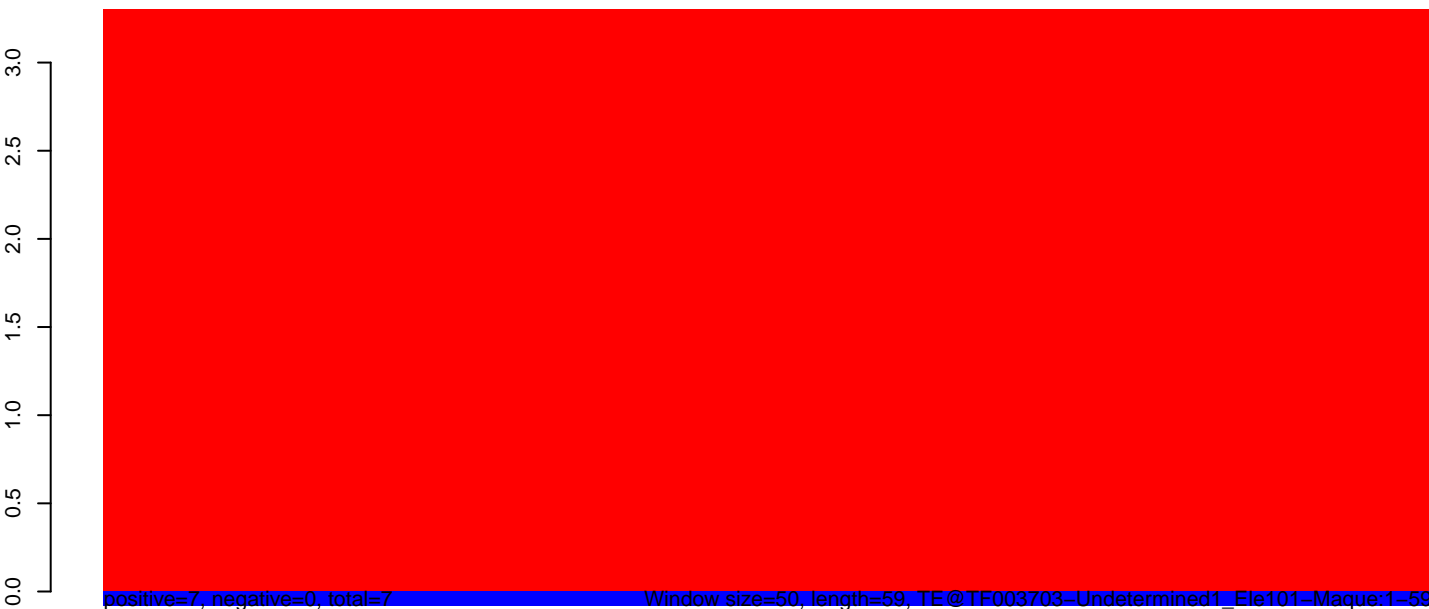
AnGam_Mos55_cells.18_23.rep



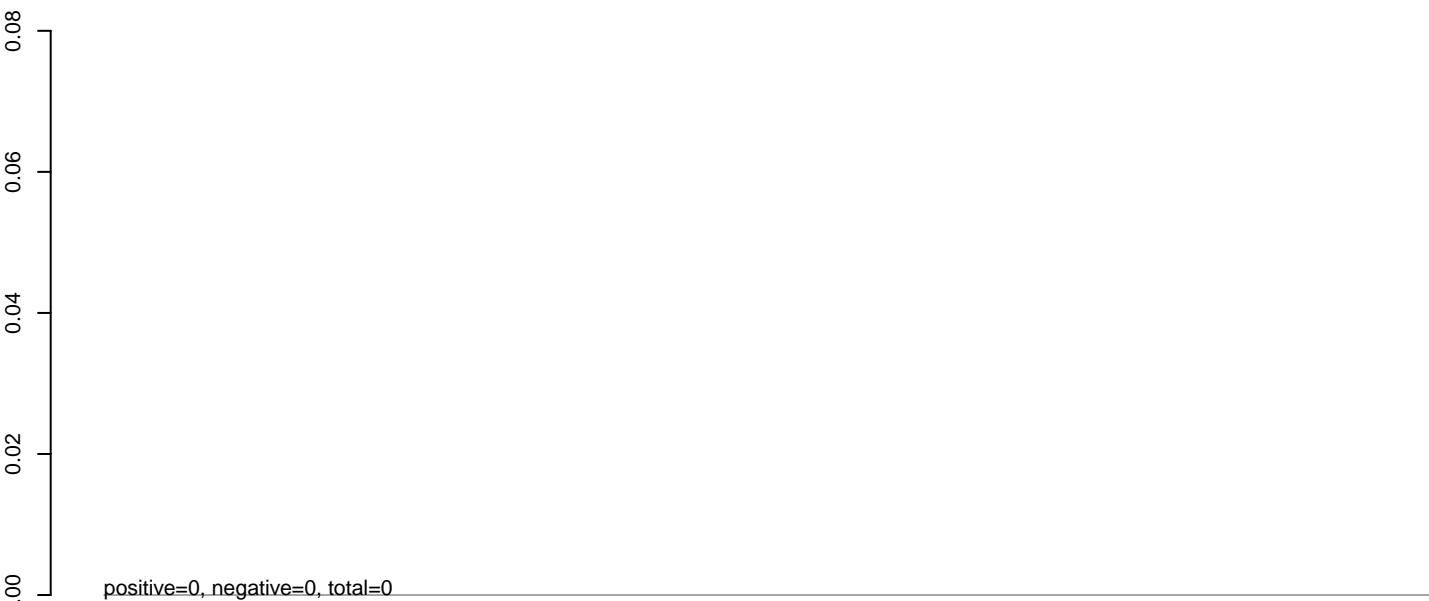
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



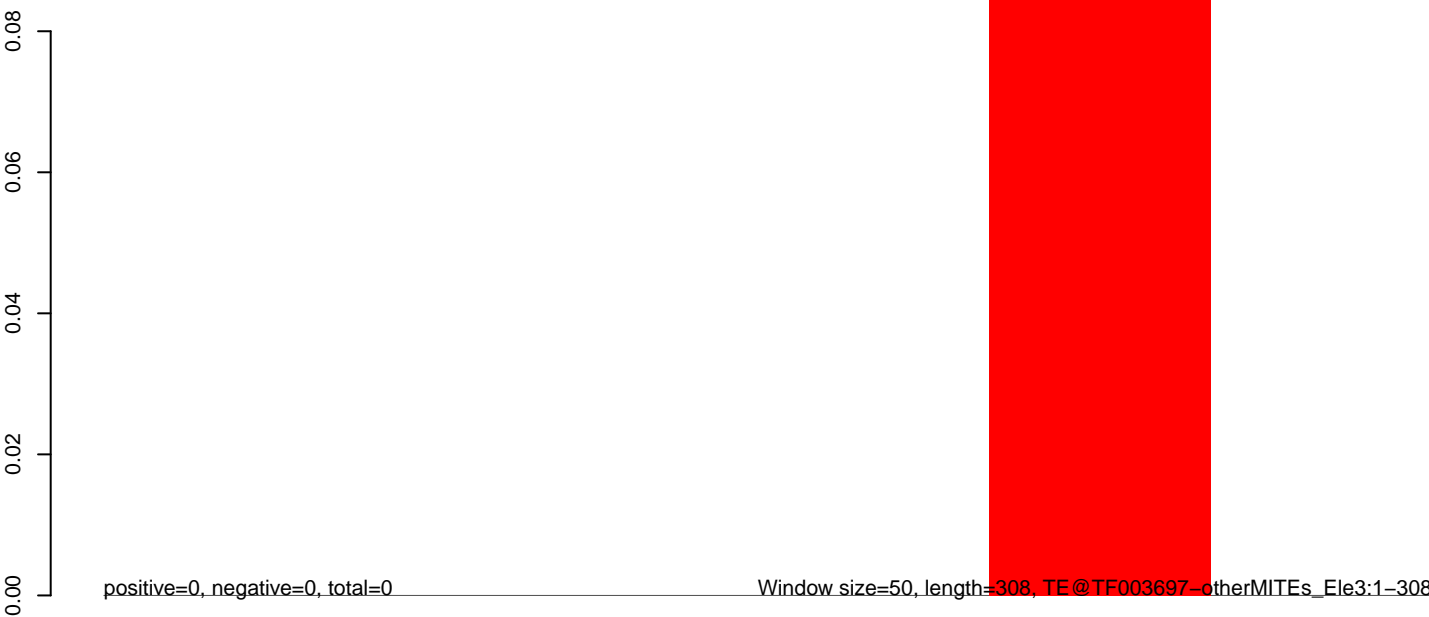
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

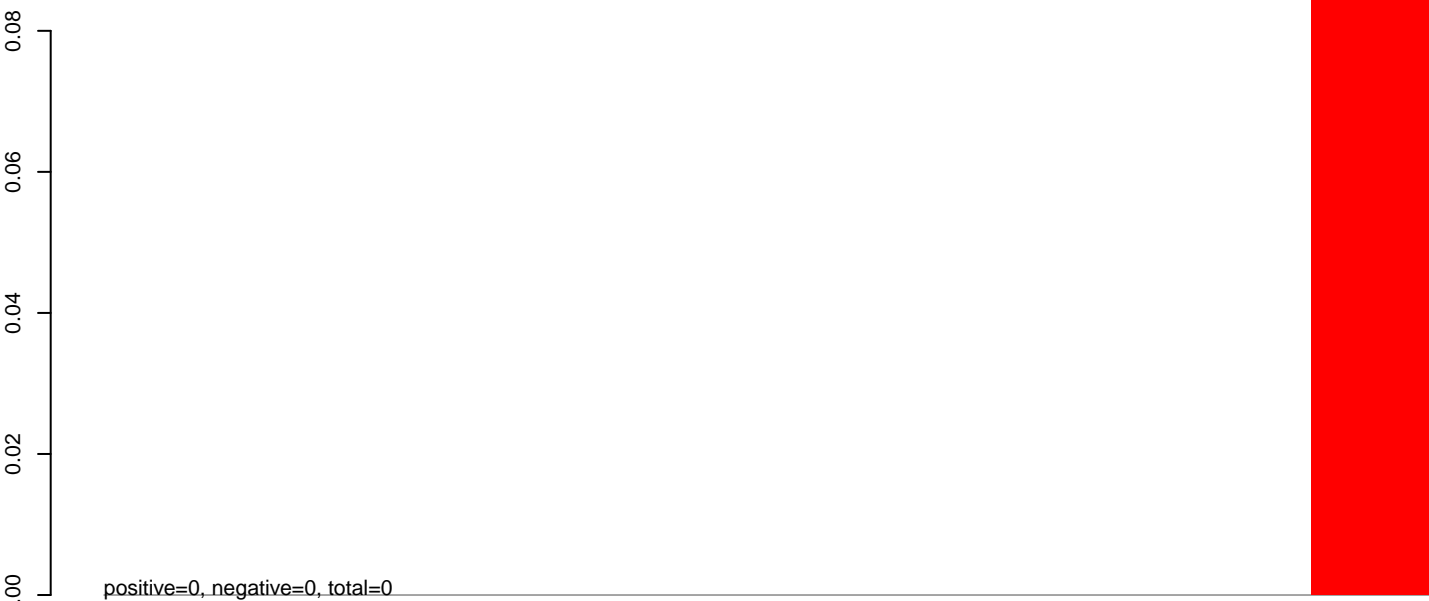


AnGam_Mos55_cells.rep

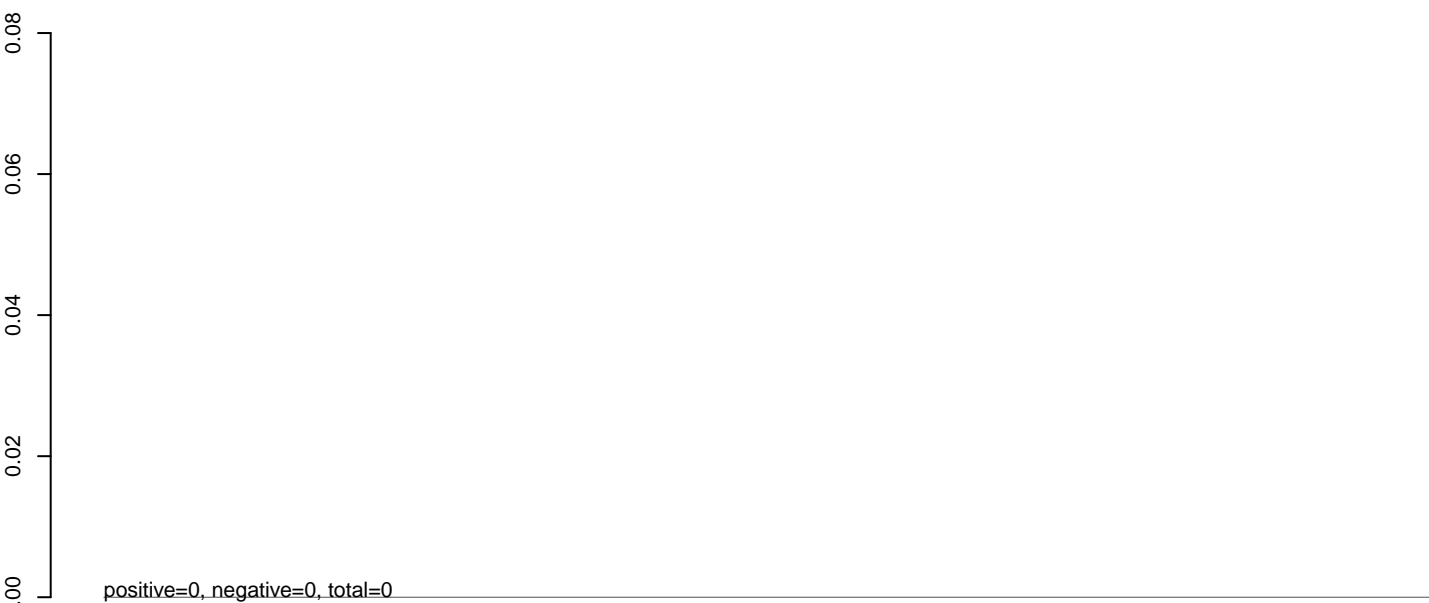


50 100 150 200 250 300 350

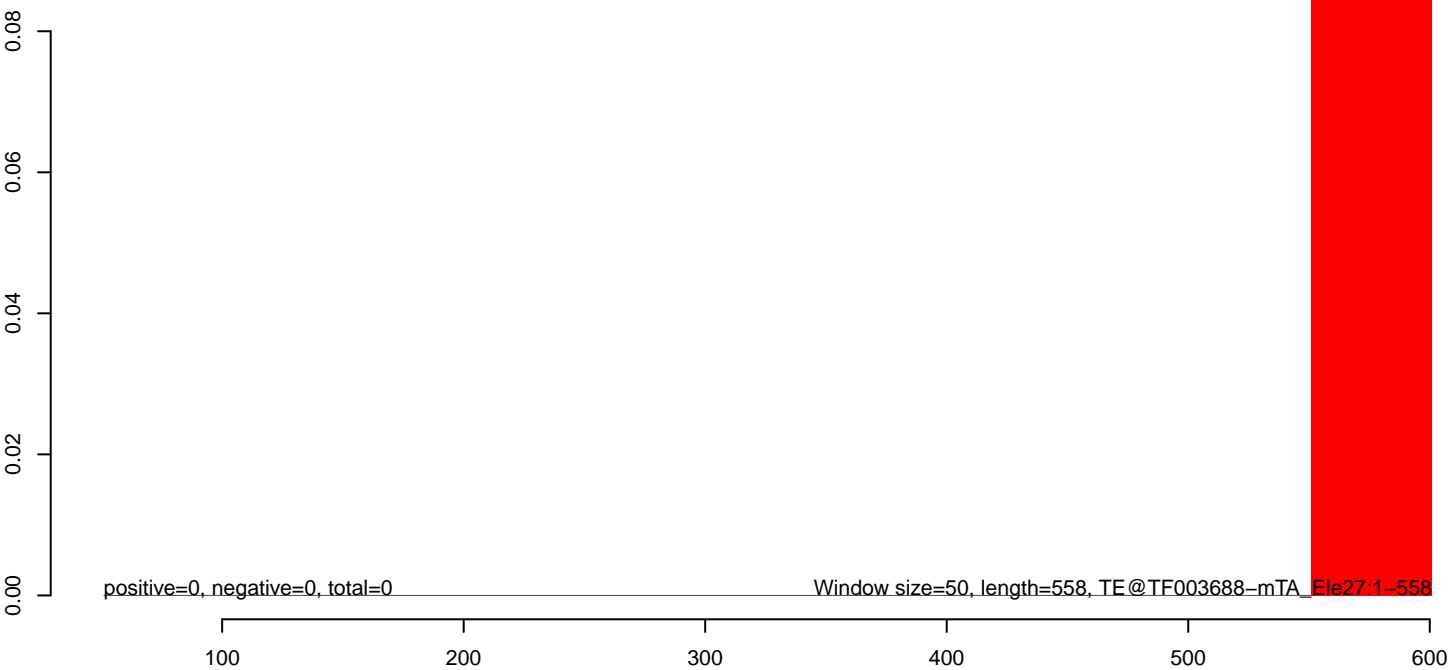
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

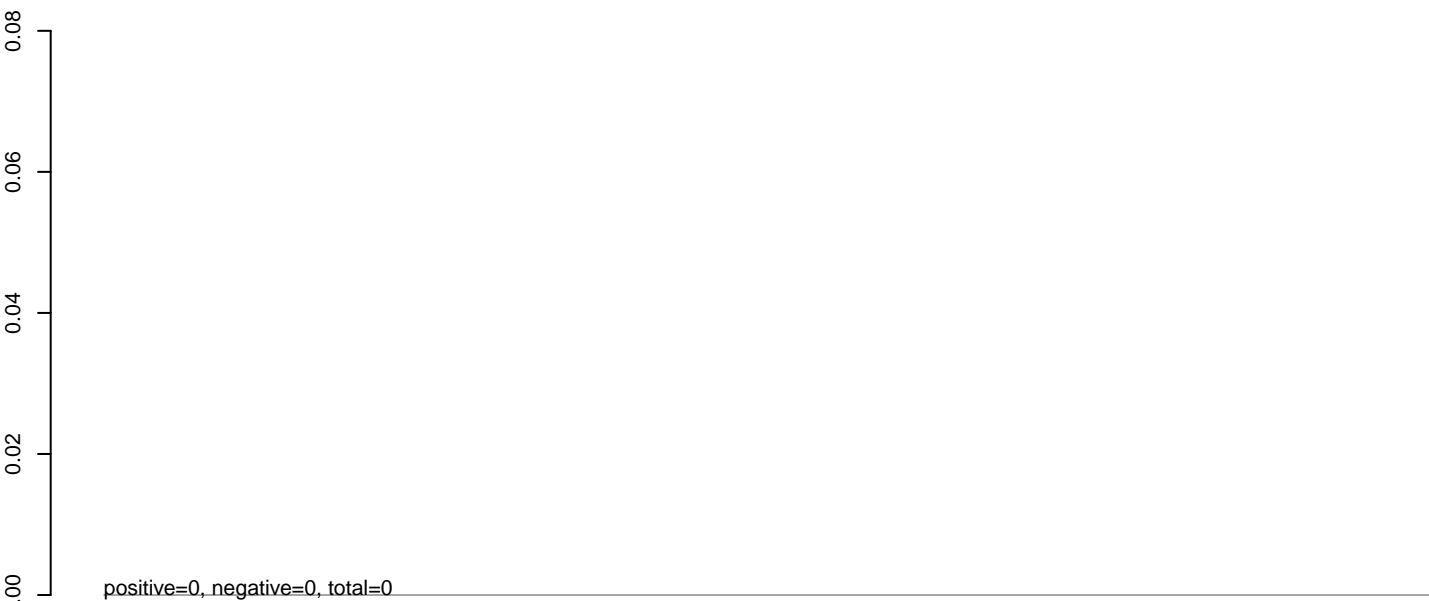
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-0.08

positive=0, negative=0, total=0

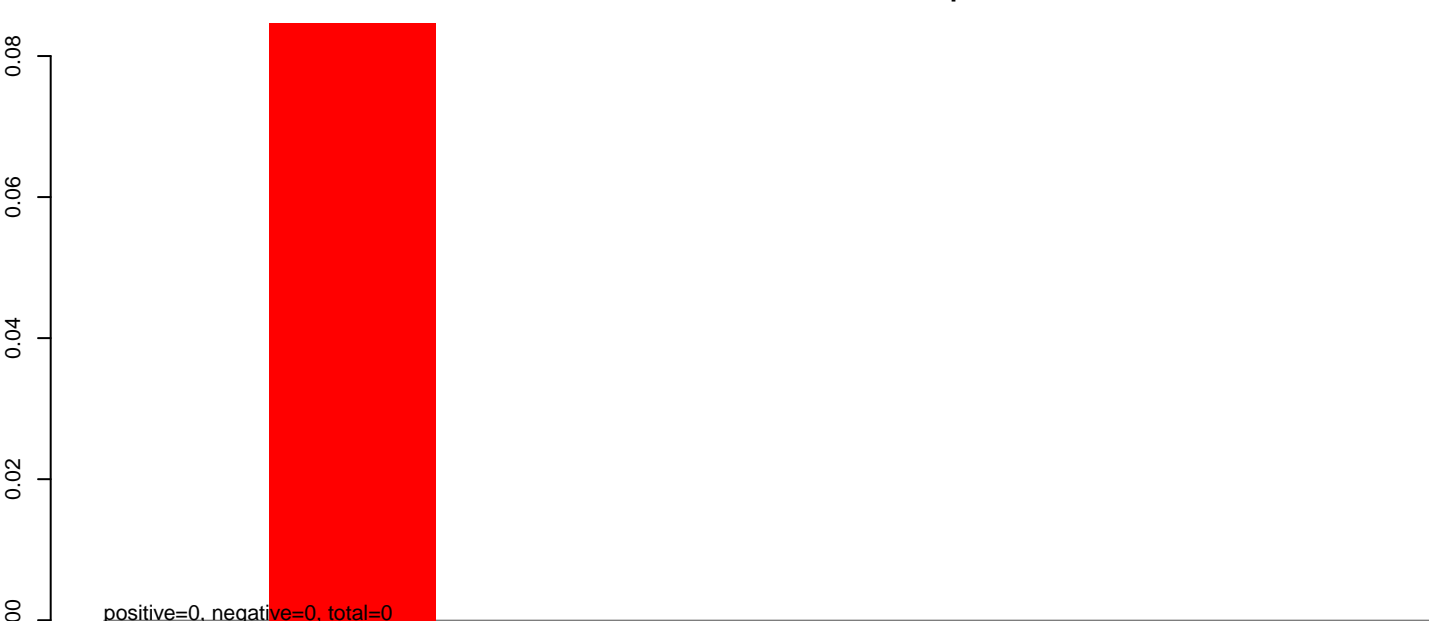
Window size=50, length=528, TE@TF003669-m8bp_Ele8:1-528

100 200 300 400 500

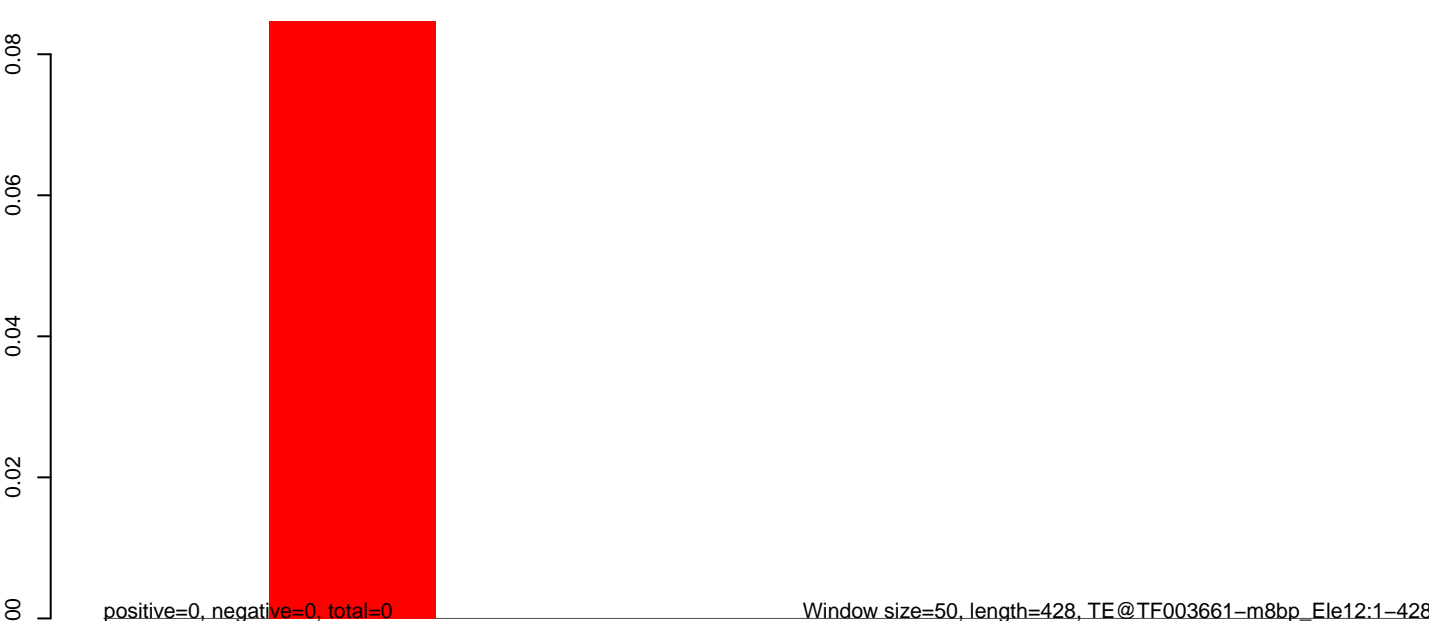
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

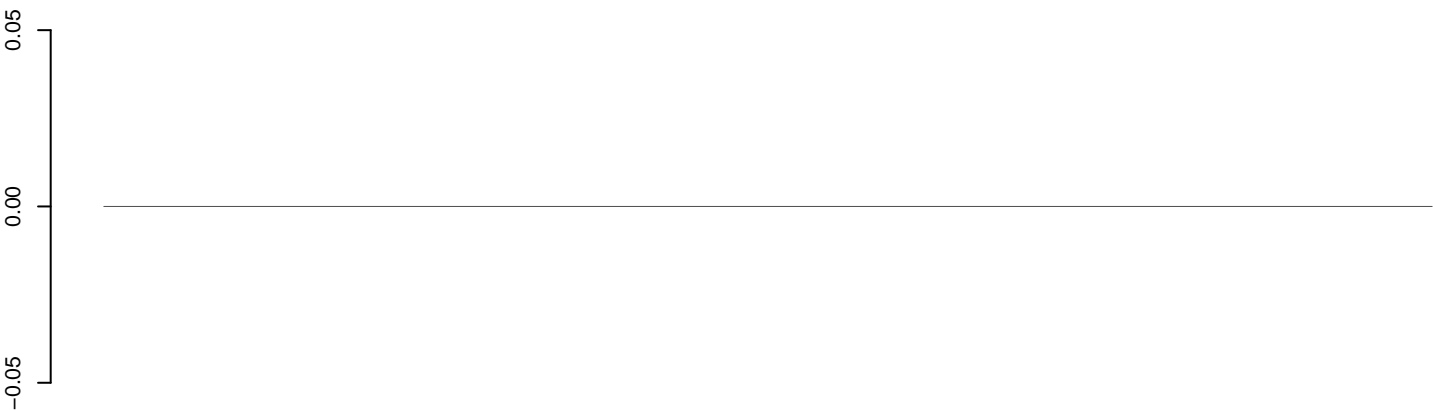


100

200

300

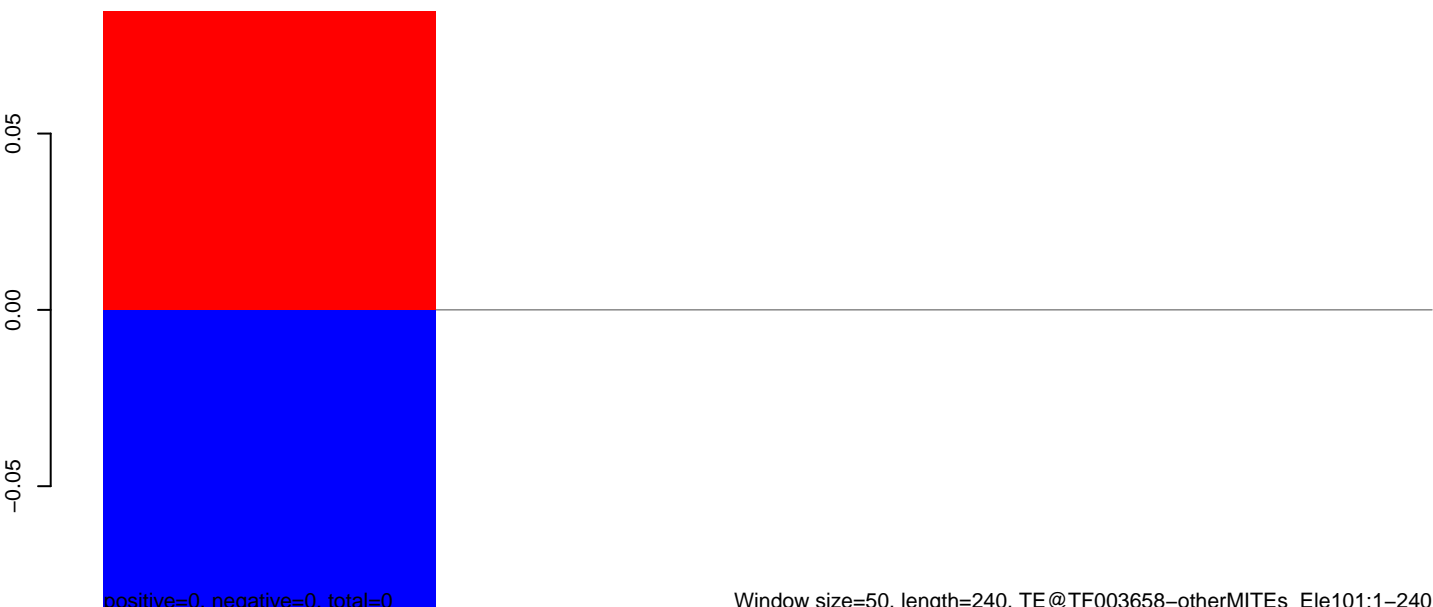
400



positive=0, negative=0, total=0

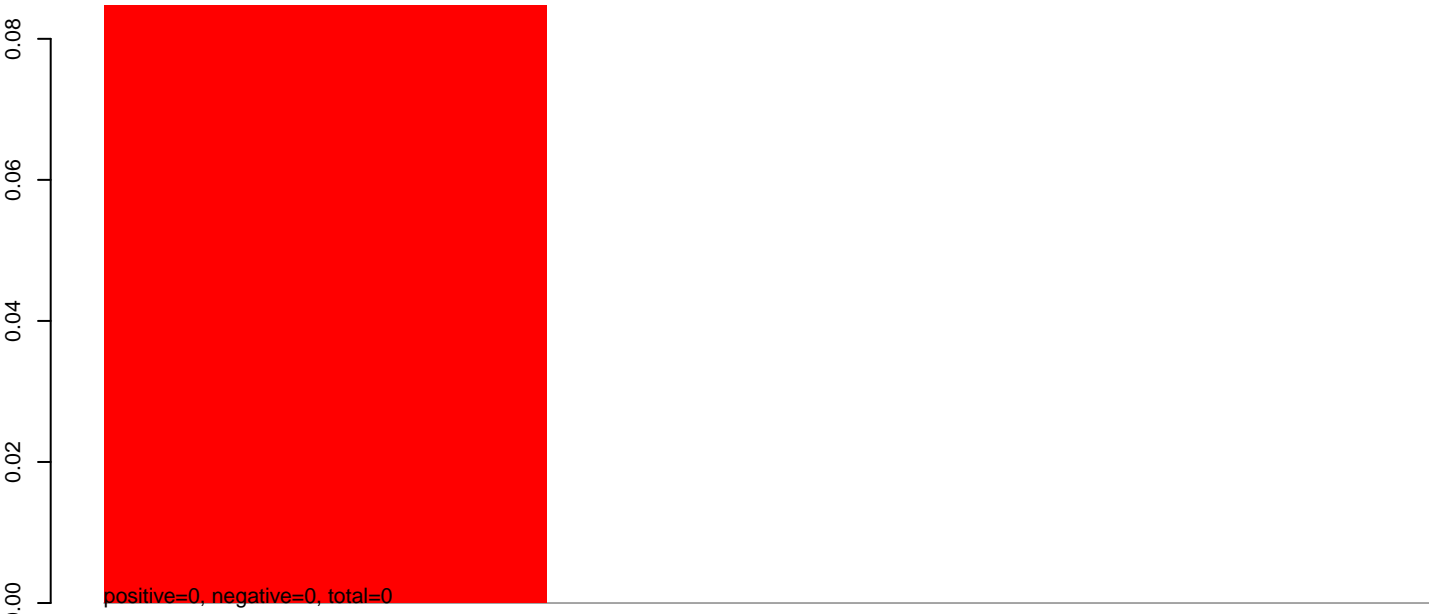


positive=0, negative=0, total=0

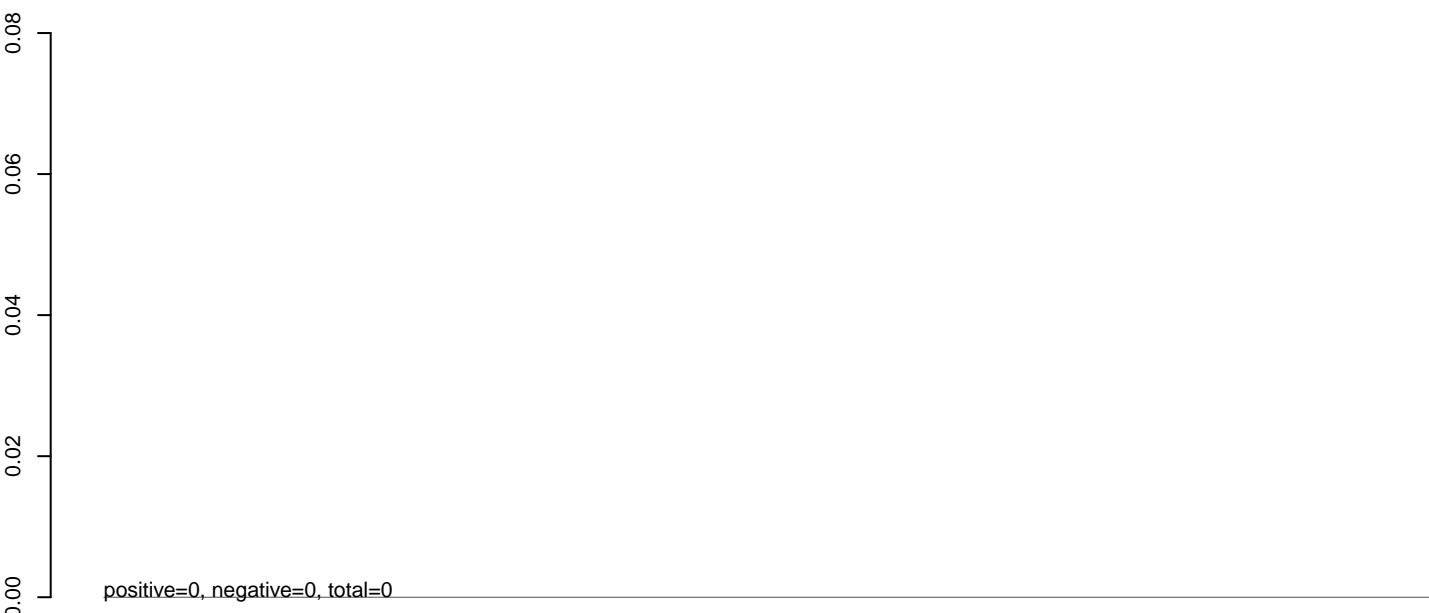


positive=0, negative=0, total=0

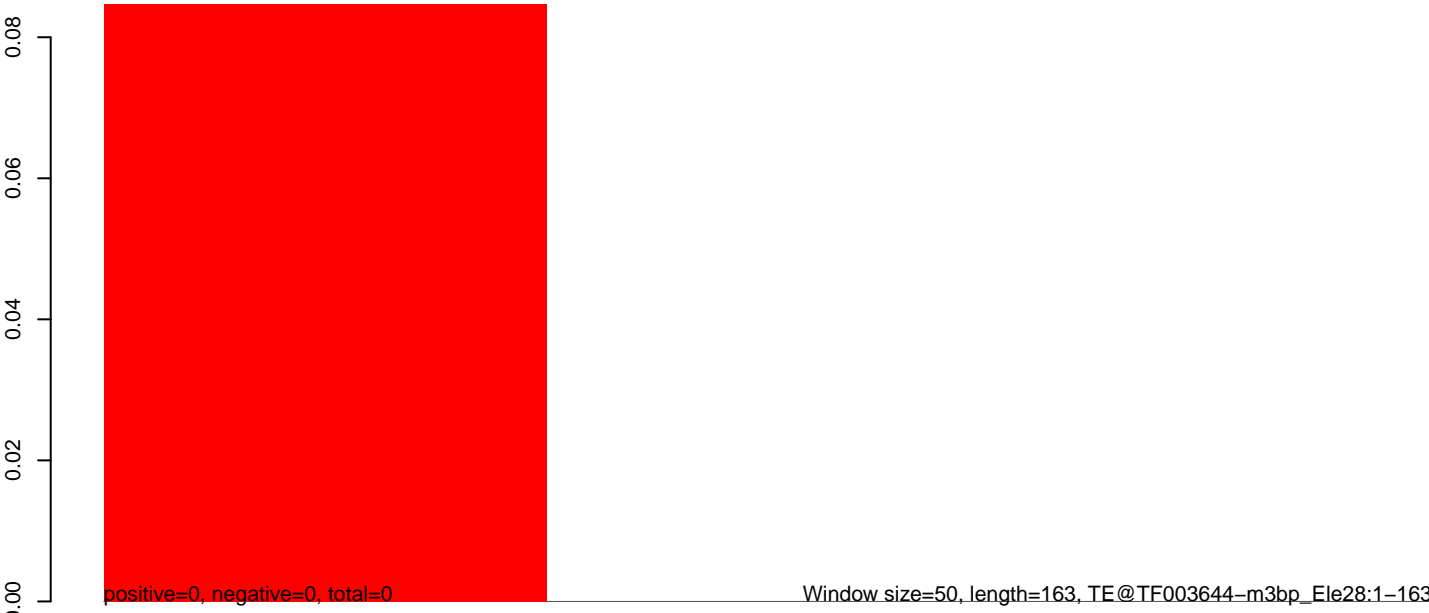
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

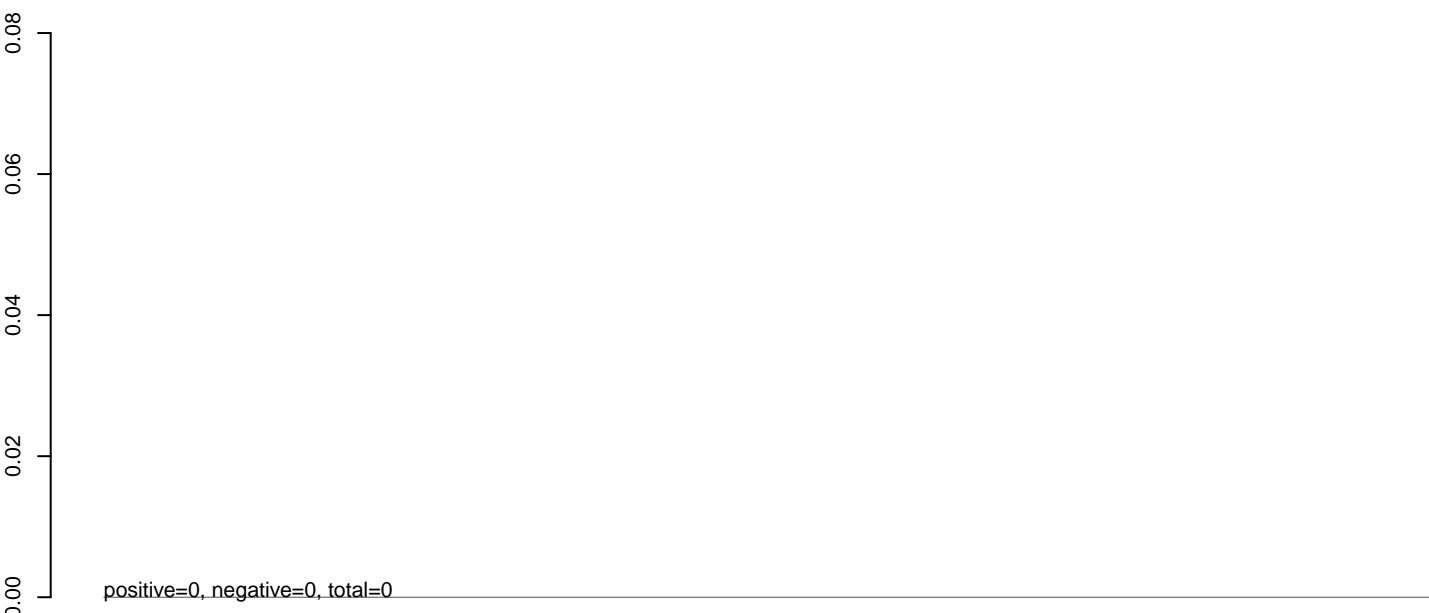


Window size=50, length=163, TE@TF003644-m3bp_Ele28:1-163

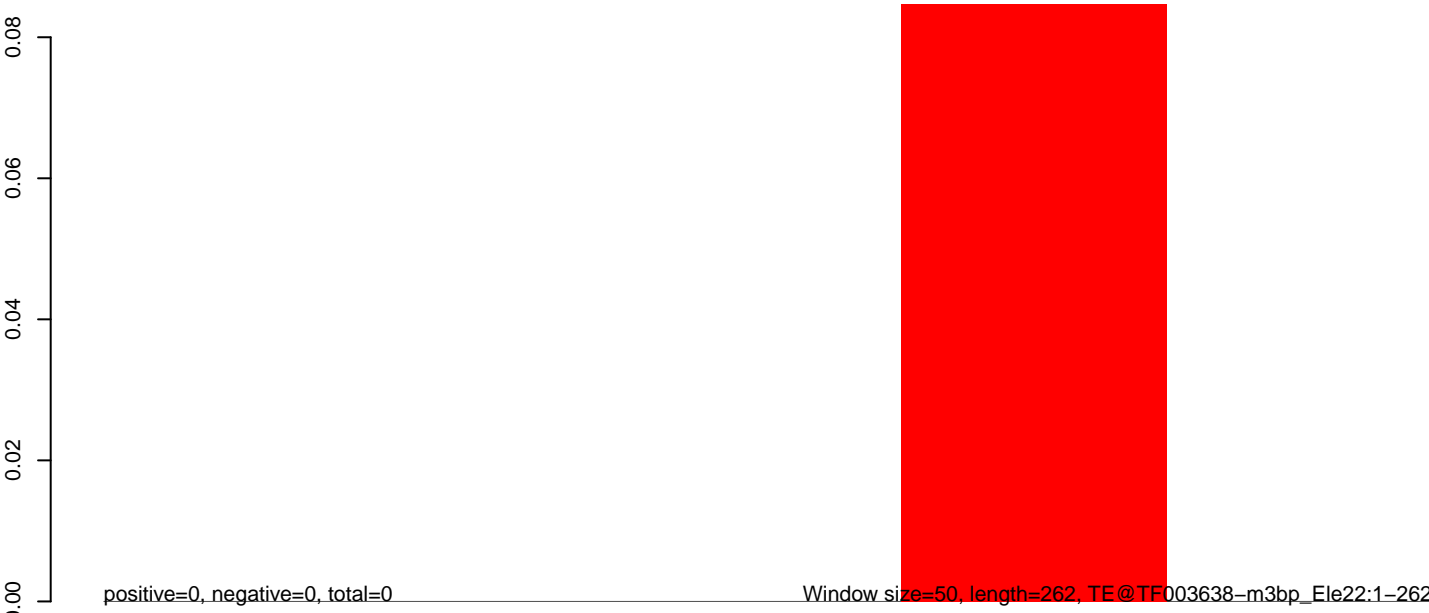
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



50 100 150 200 250 300

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=683, TE@TF003533-Tc1_Le16-Mariner1_AG:1-683

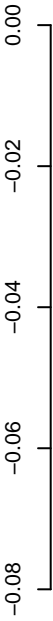
100 200 300 400 500 600 700

AnGam_Mos55_cells.18_23.rep



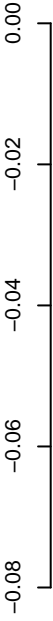
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



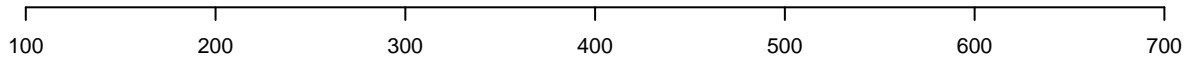
positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

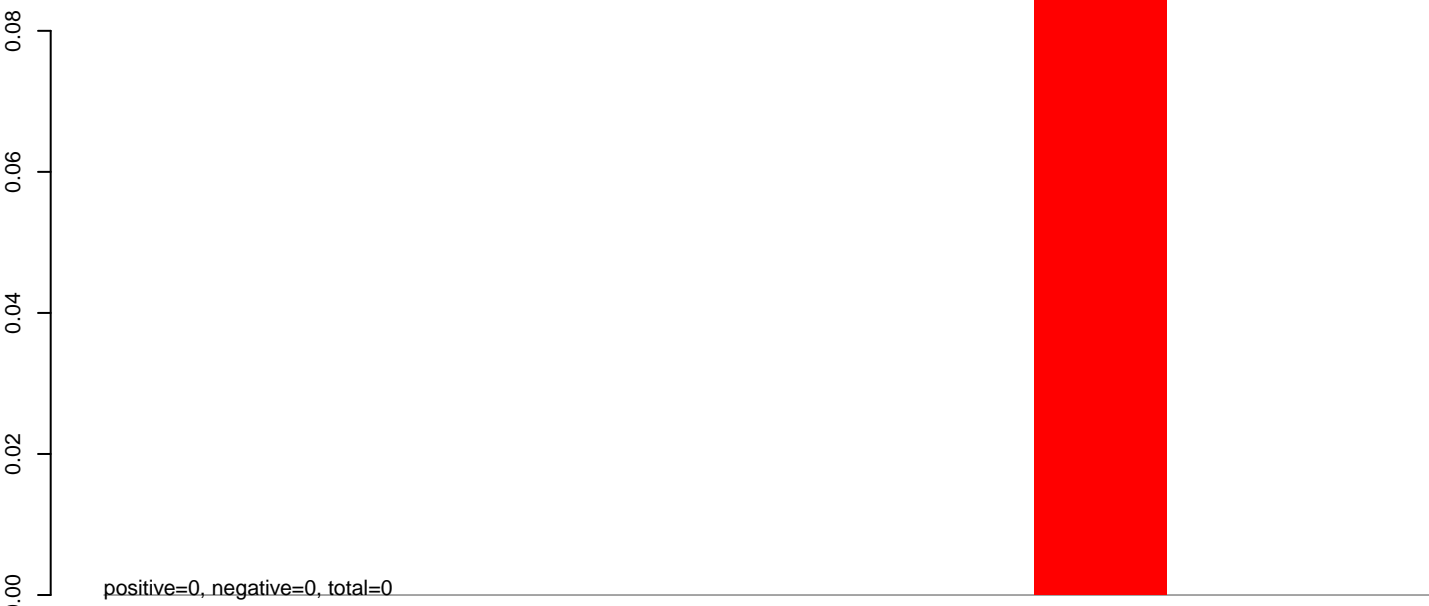


positive=0, negative=0, total=0

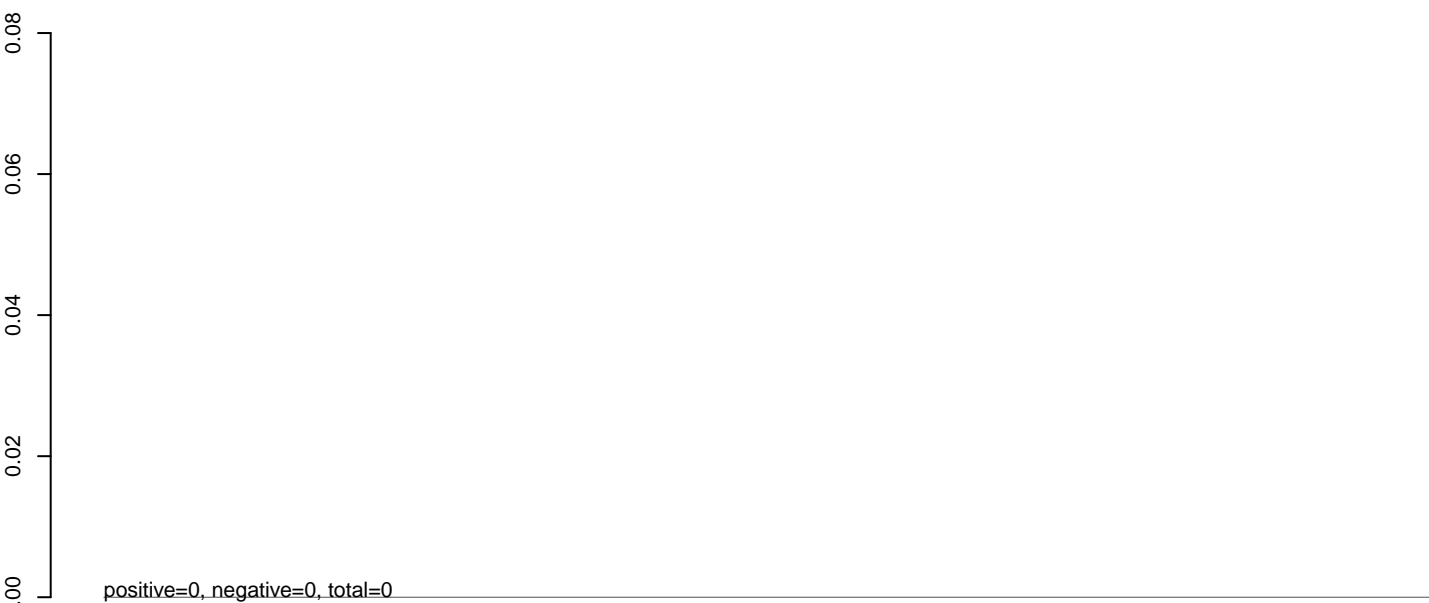
Window size=50, length=730, TE@TF003526-Tc1_Ele10:1-730



AnGam_Mos55_cells.18_23.rep



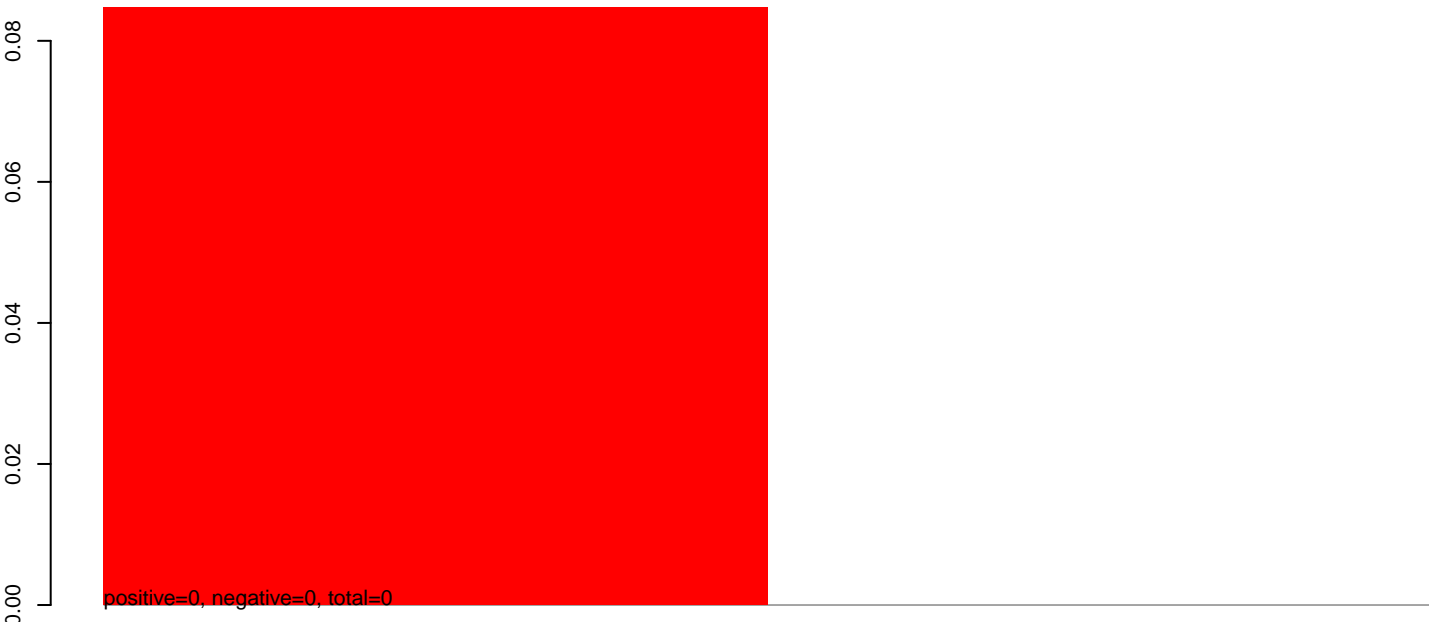
AnGam_Mos55_cells.24_35.rep



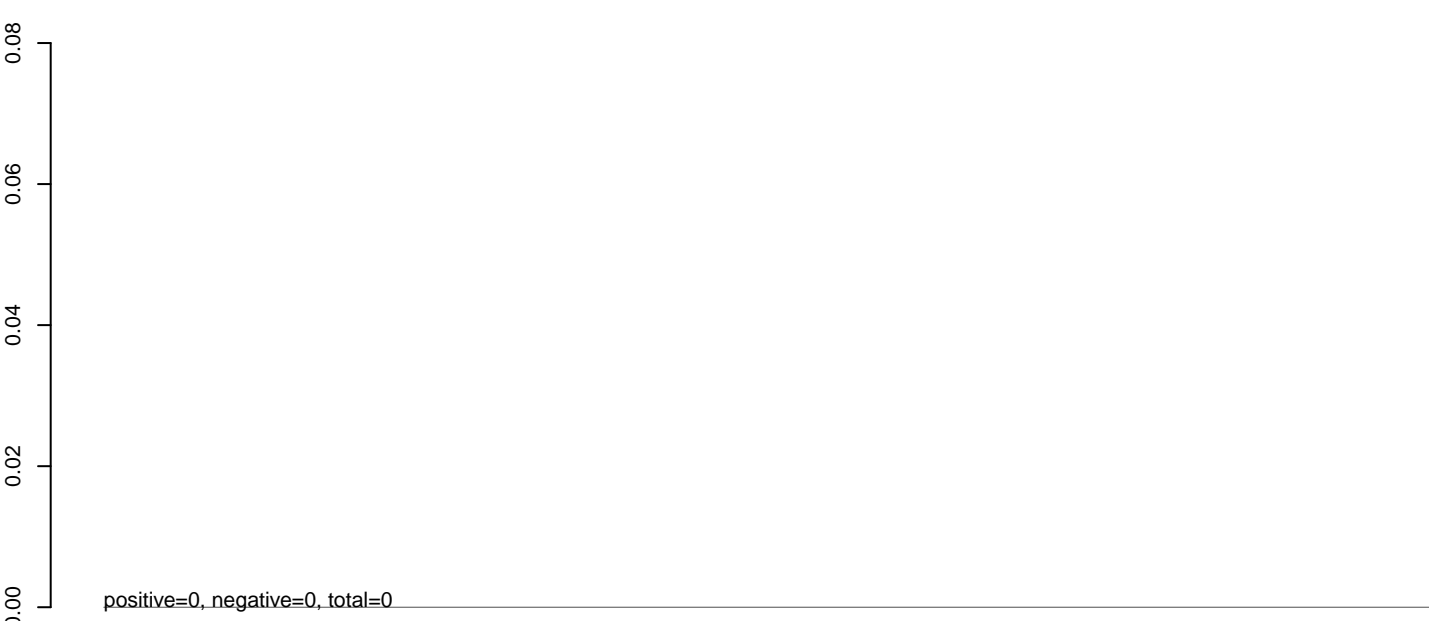
AnGam_Mos55_cells.rep



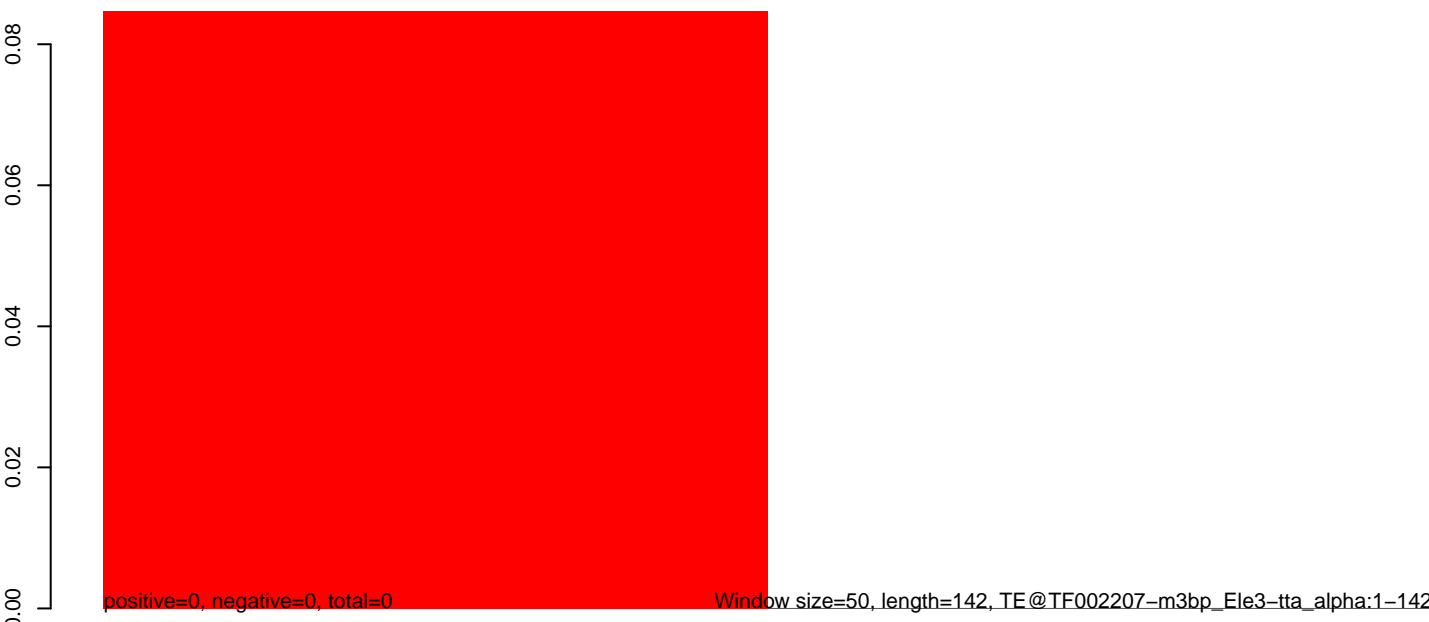
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

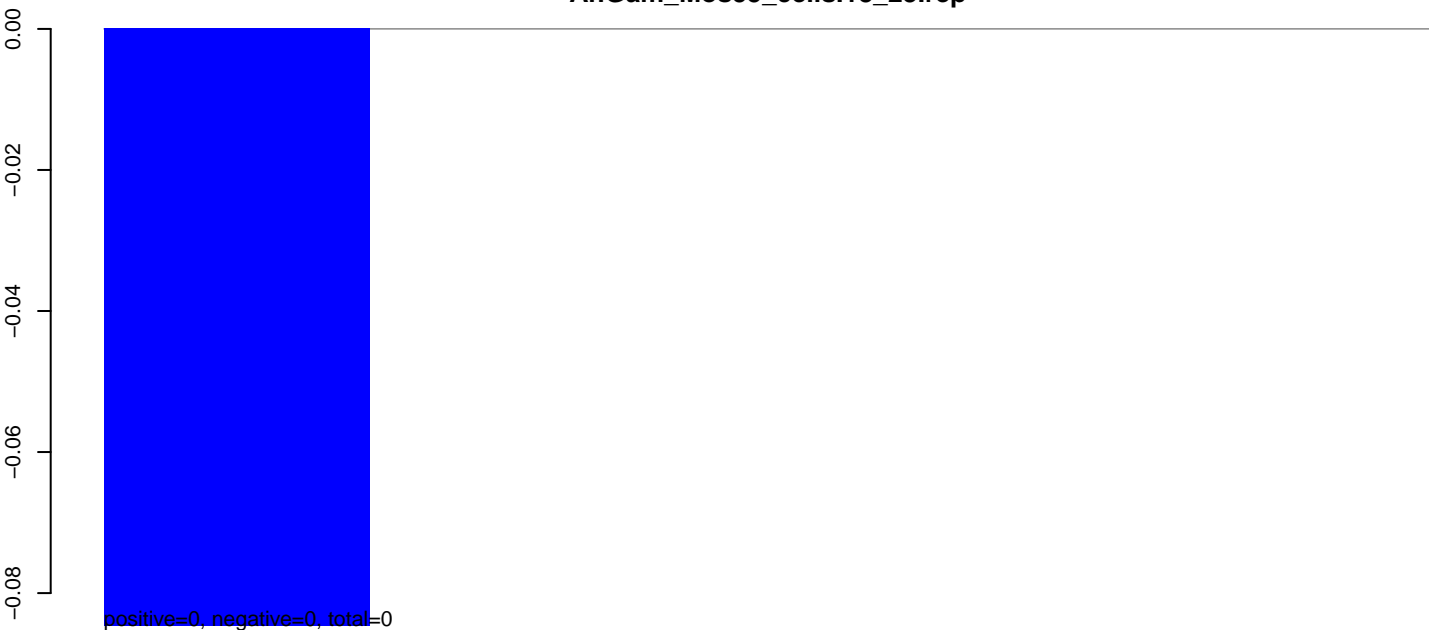


AnGam_Mos55_cells.rep

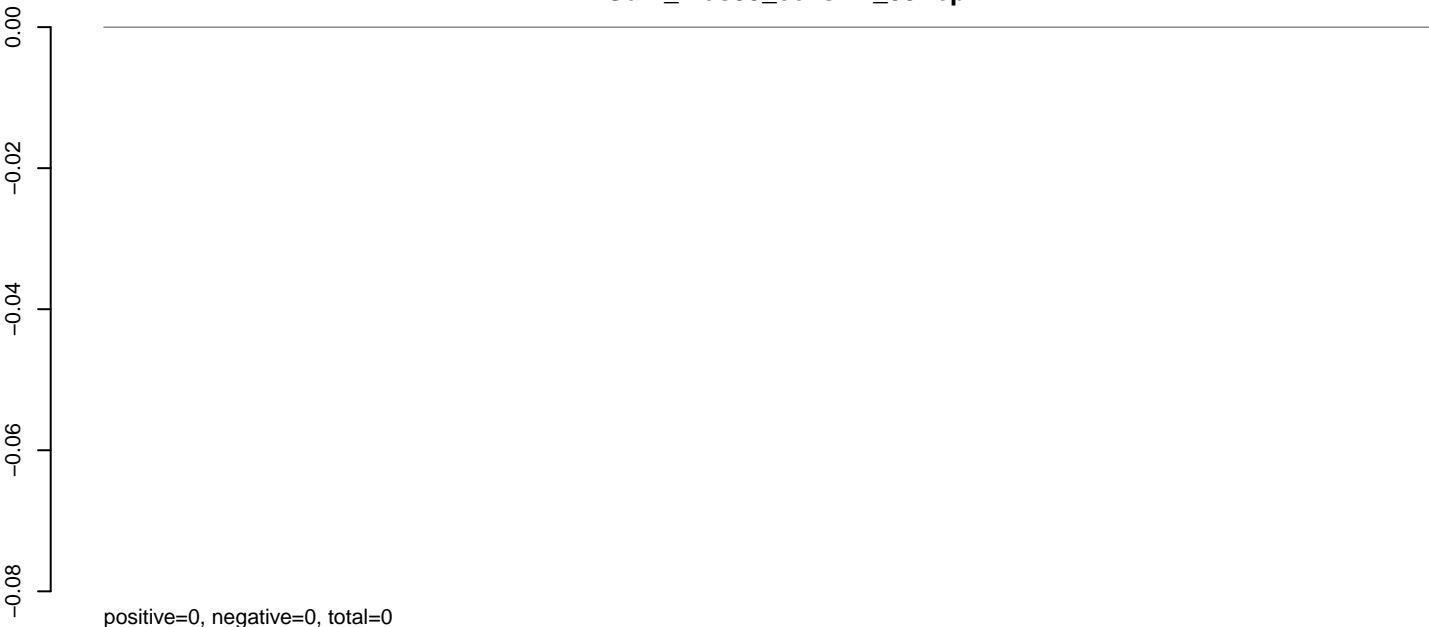


60 80 100 120 140

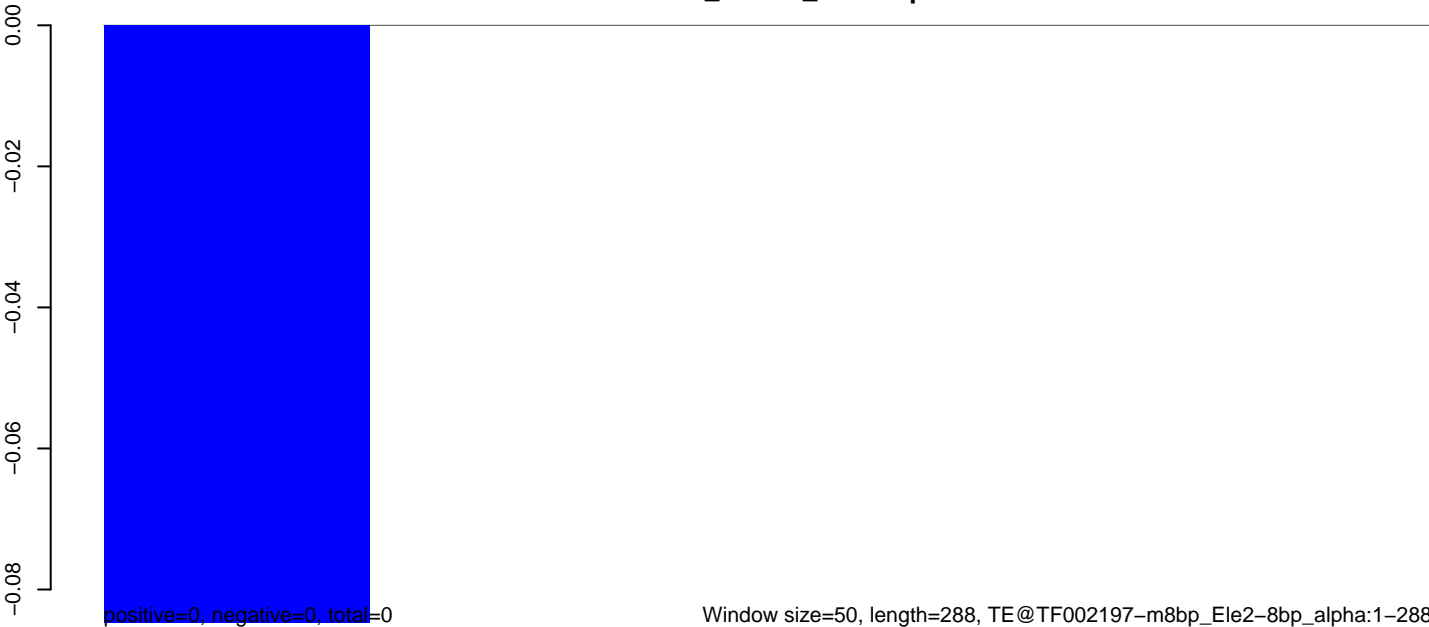
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



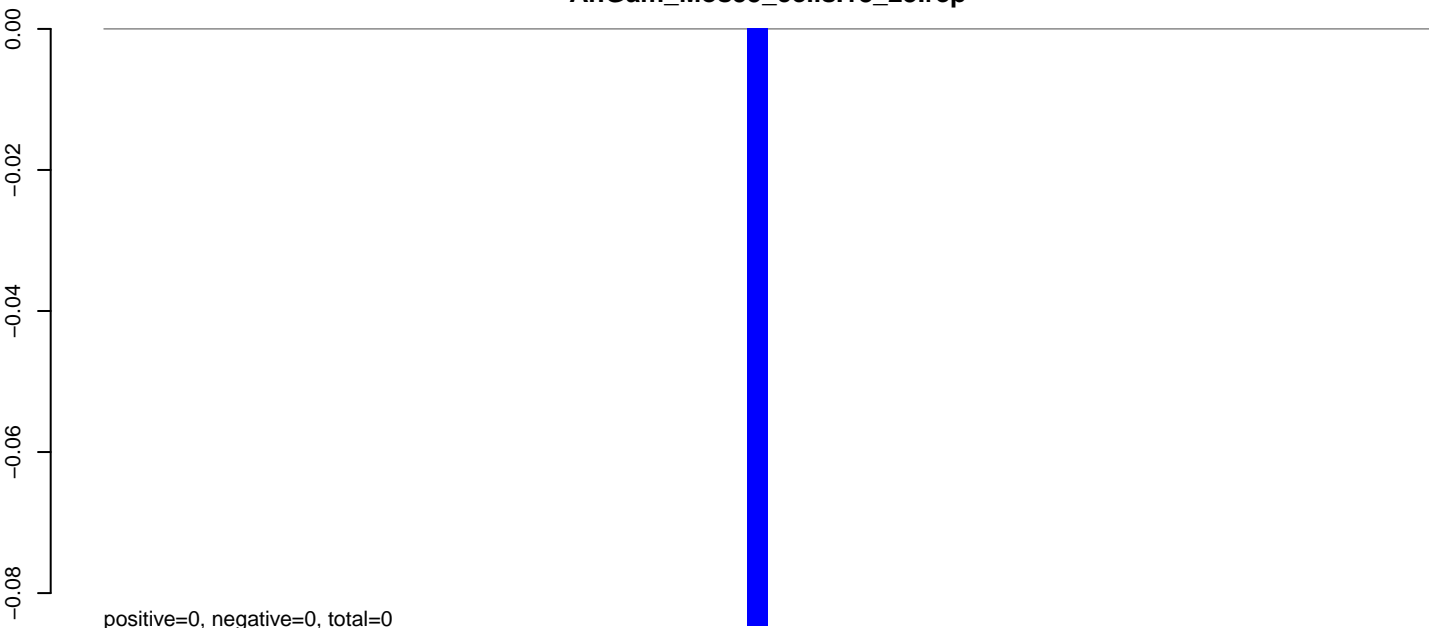
AnGam_Mos55_cells.rep



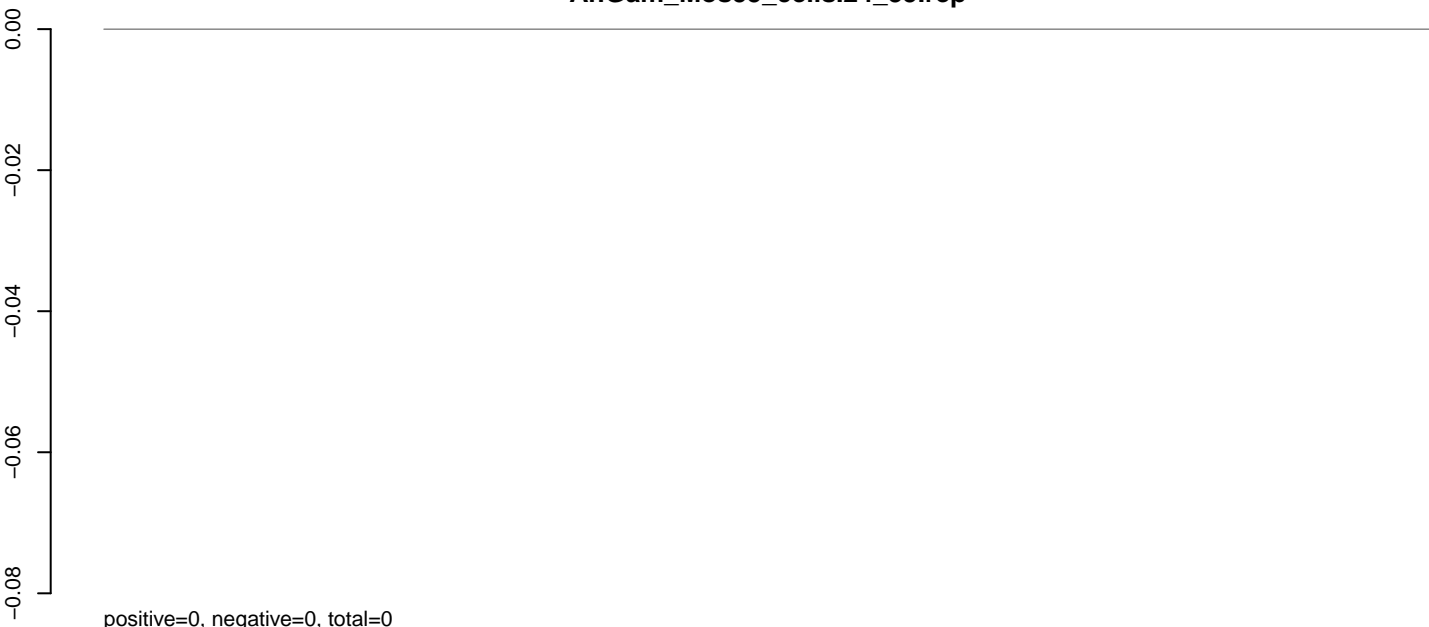
Window size=50, length=288, TE@TF002197-m8bp_Ele2-8bp_alpha:1-288

50 100 150 200 250 300

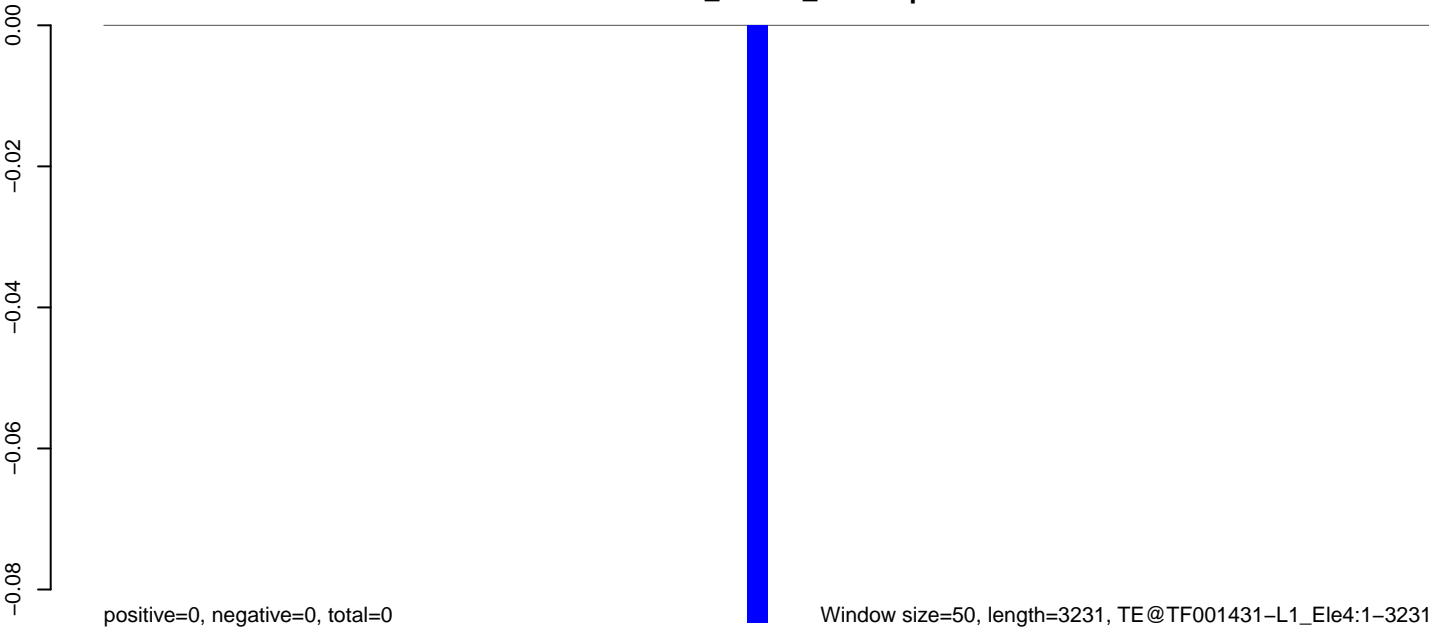
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



0 500 1000 1500 2000 2500 3000

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=740, TE@TE001411-Outcast_Ele10:1-740

100 200 300 400 500 600 700

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

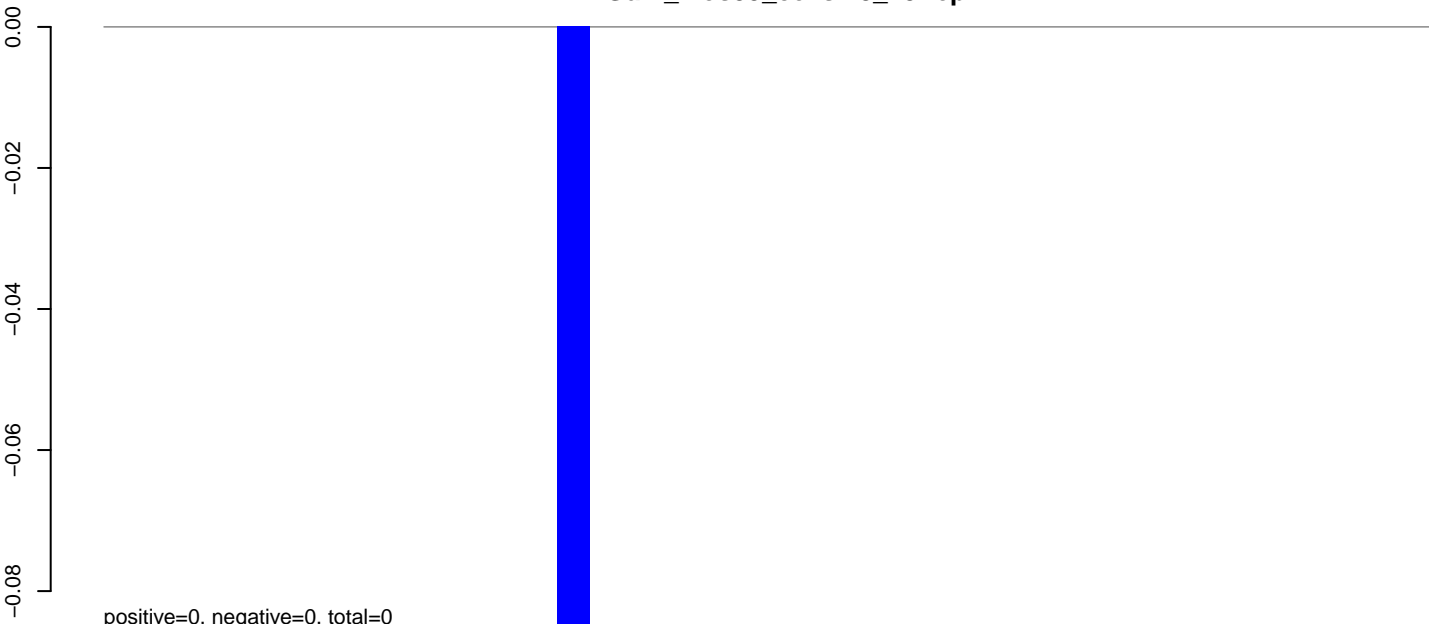
0.00
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-0.06
-0.08

positive=0, negative=0, total=0

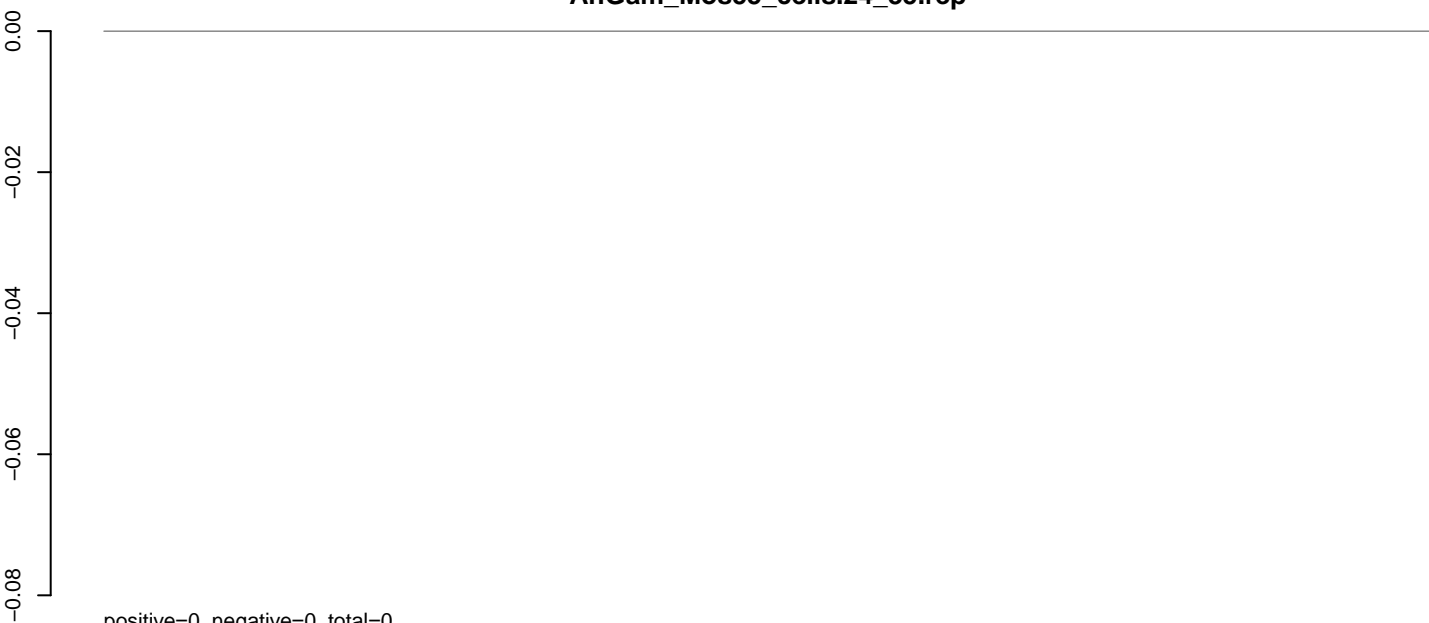
Window size=50, length=741, TE@TF001410-Outcast_Ele9:1-741

100 200 300 400 500 600 700

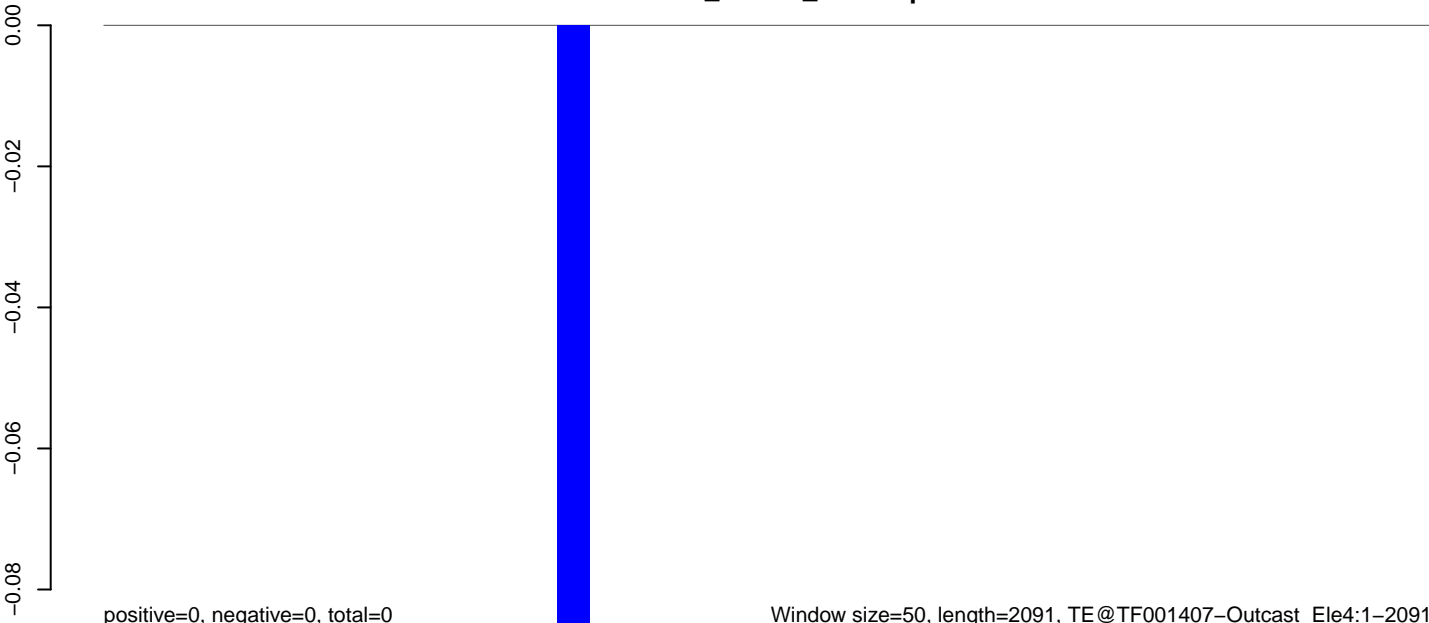
AnGam_Mos55_cells.18_23.rep



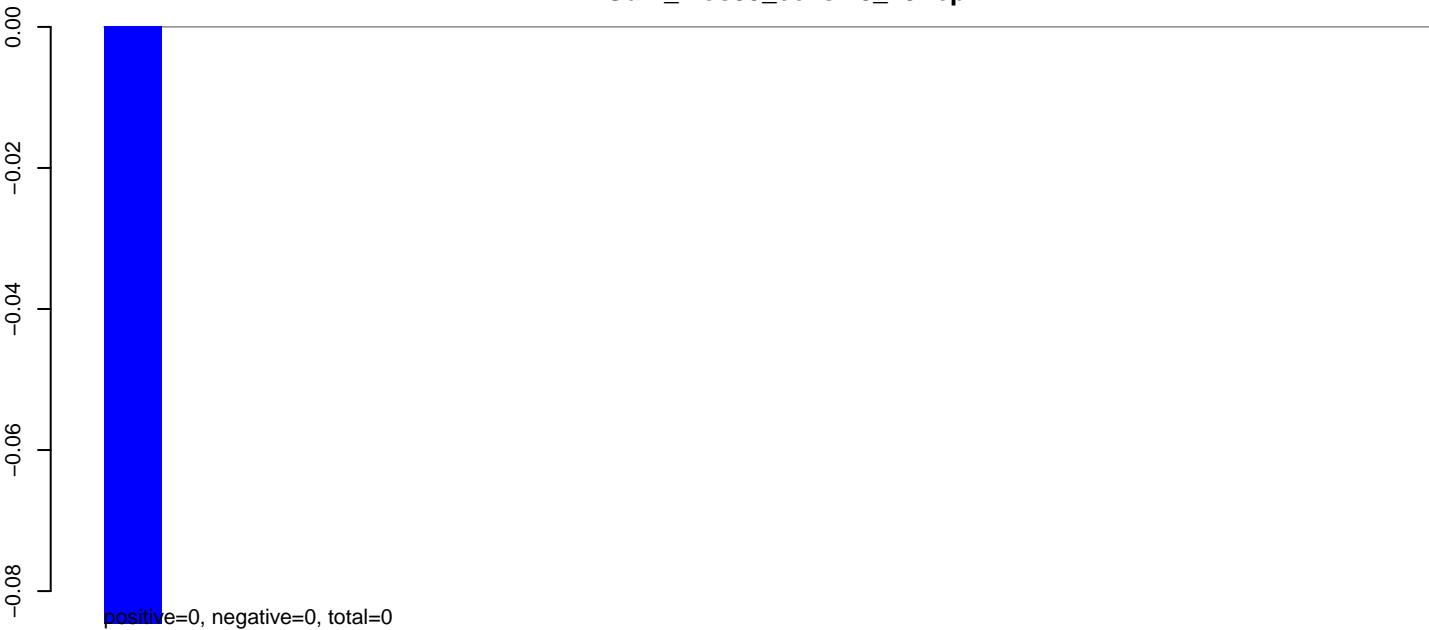
AnGam_Mos55_cells.24_35.rep



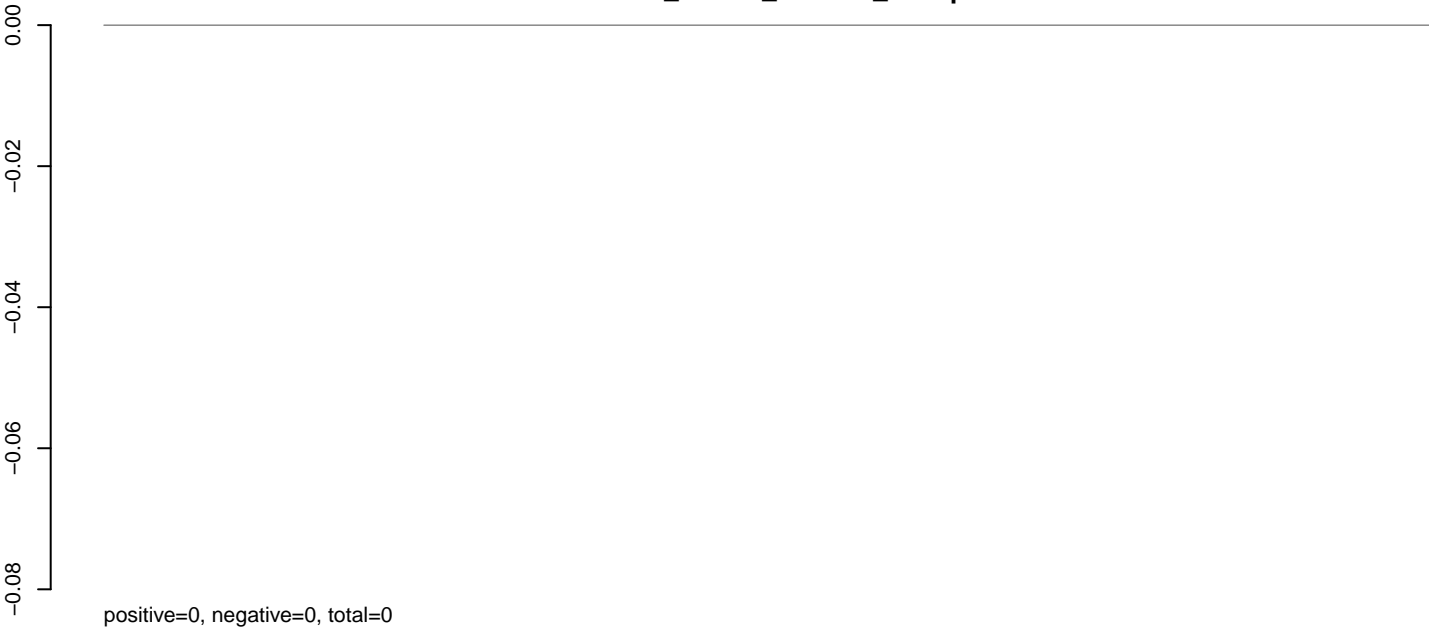
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



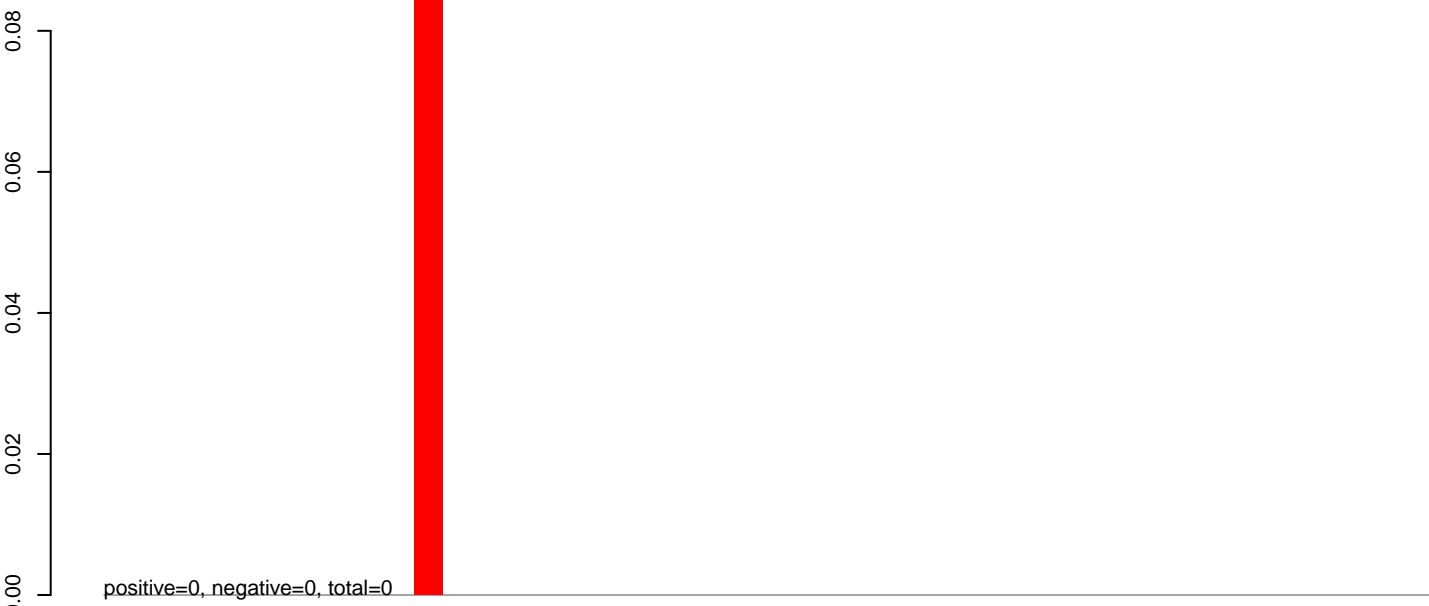
AnGam_Mos55_cells.24_35.rep



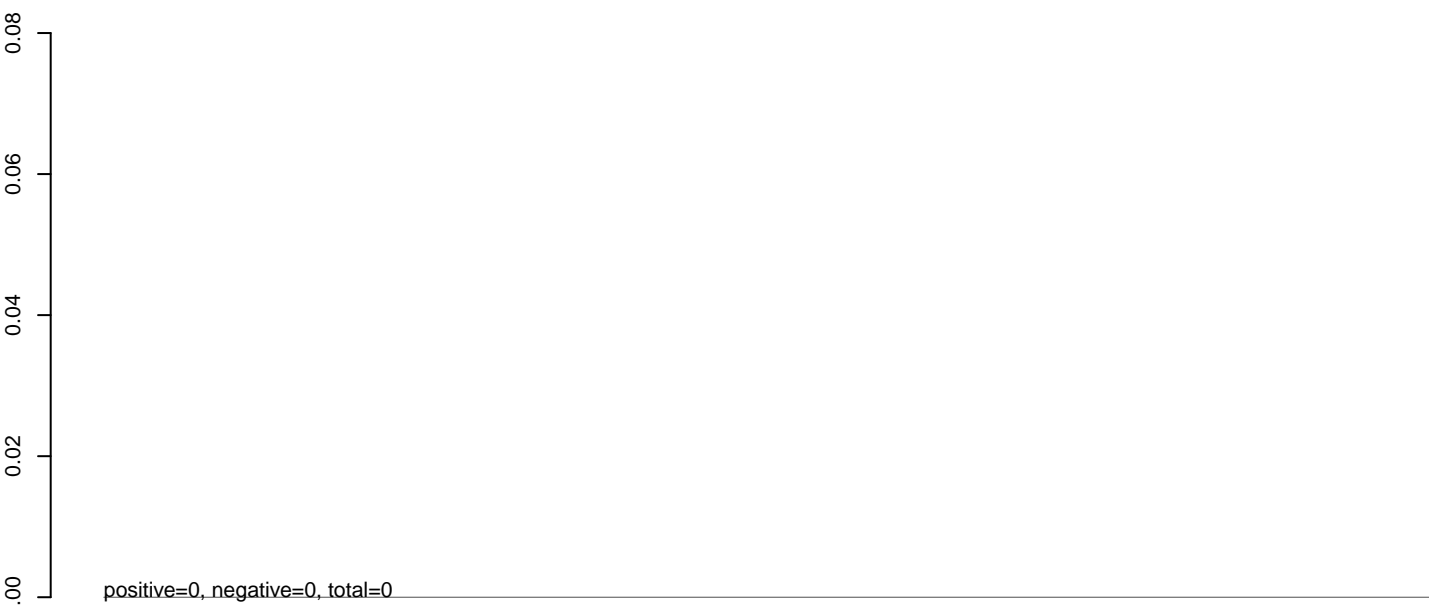
AnGam_Mos55_cells.rep



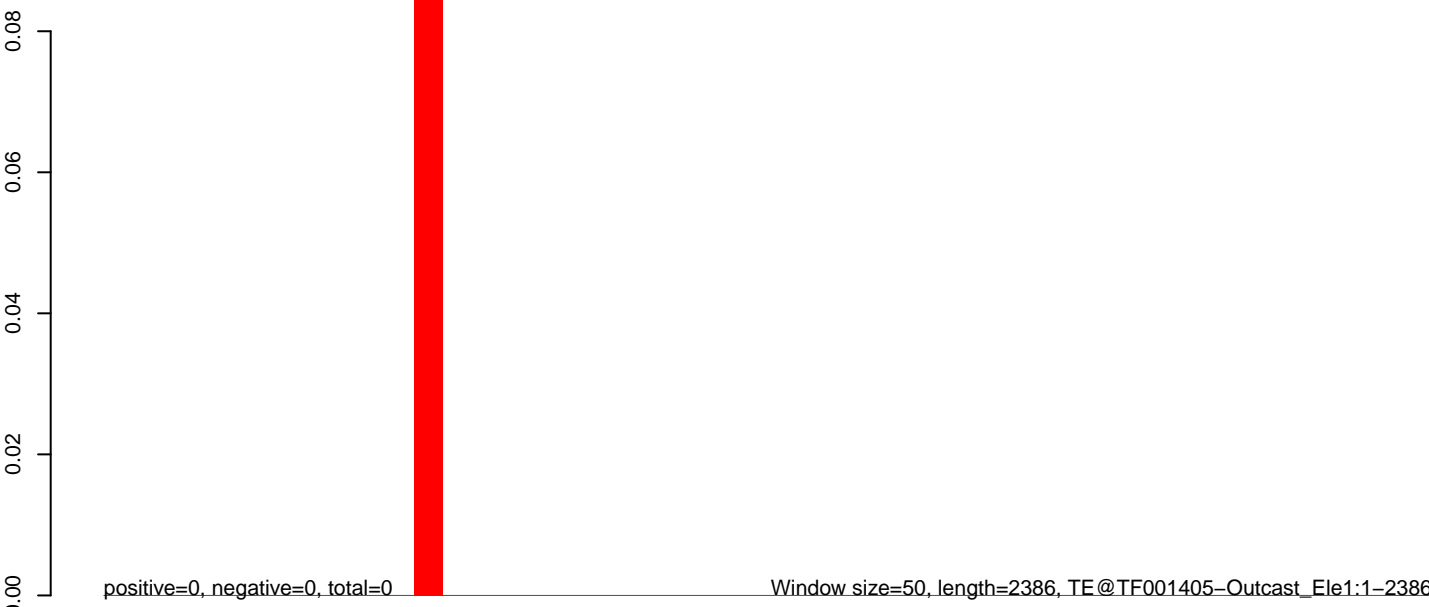
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



0 500 1000 1500 2000

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

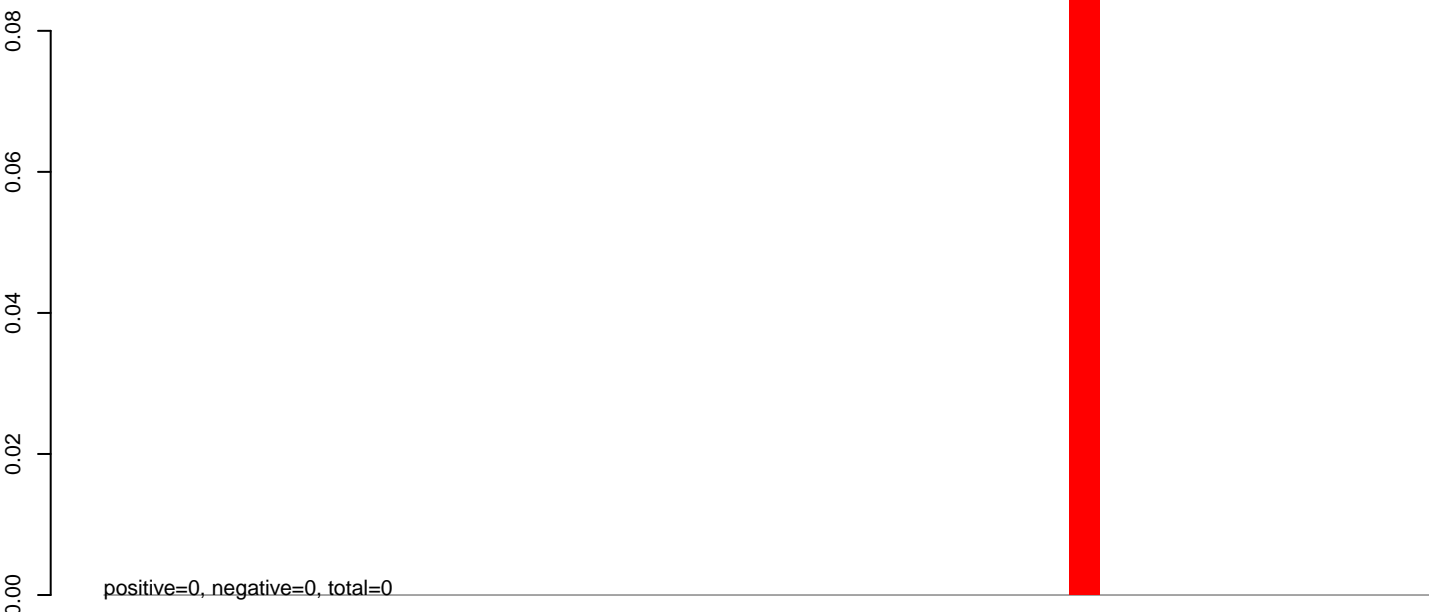
0.00
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-0.06
-0.08

positive=0, negative=0, total=0

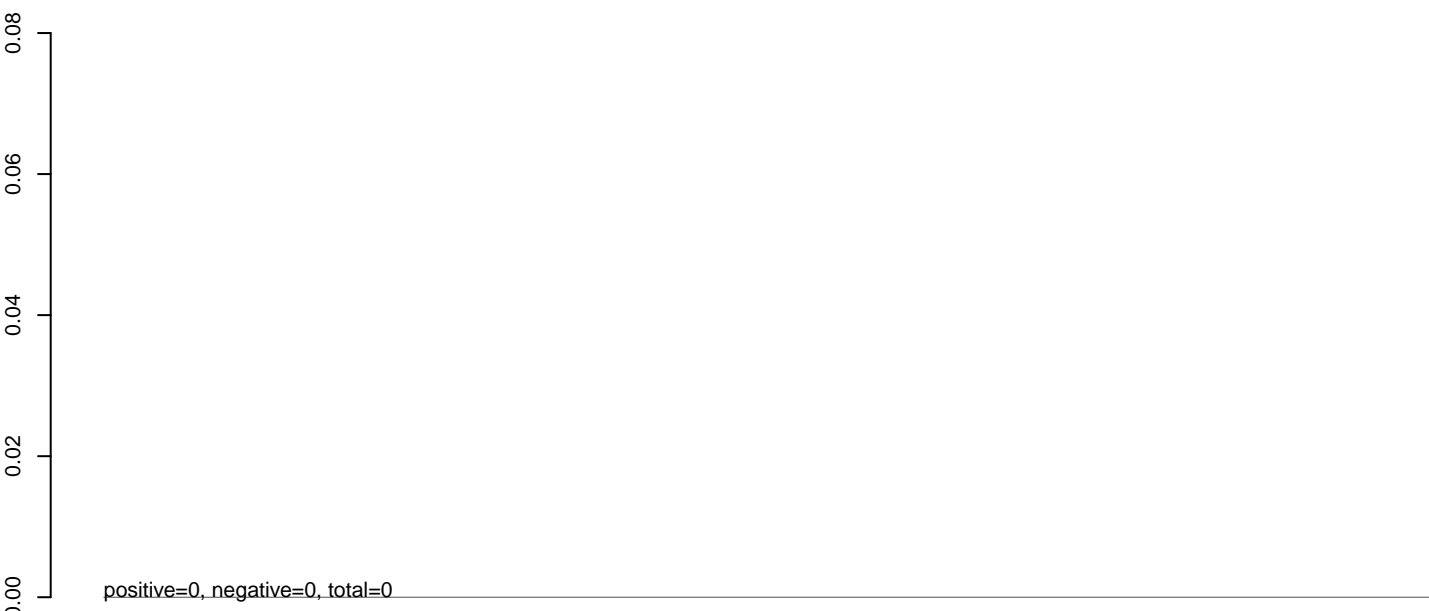
Window size=50, length=2223, TE@TF001398-CR1_Ele25:1-2223

0 500 1000 1500 2000

AnGam_Mos55_cells.18_23.rep



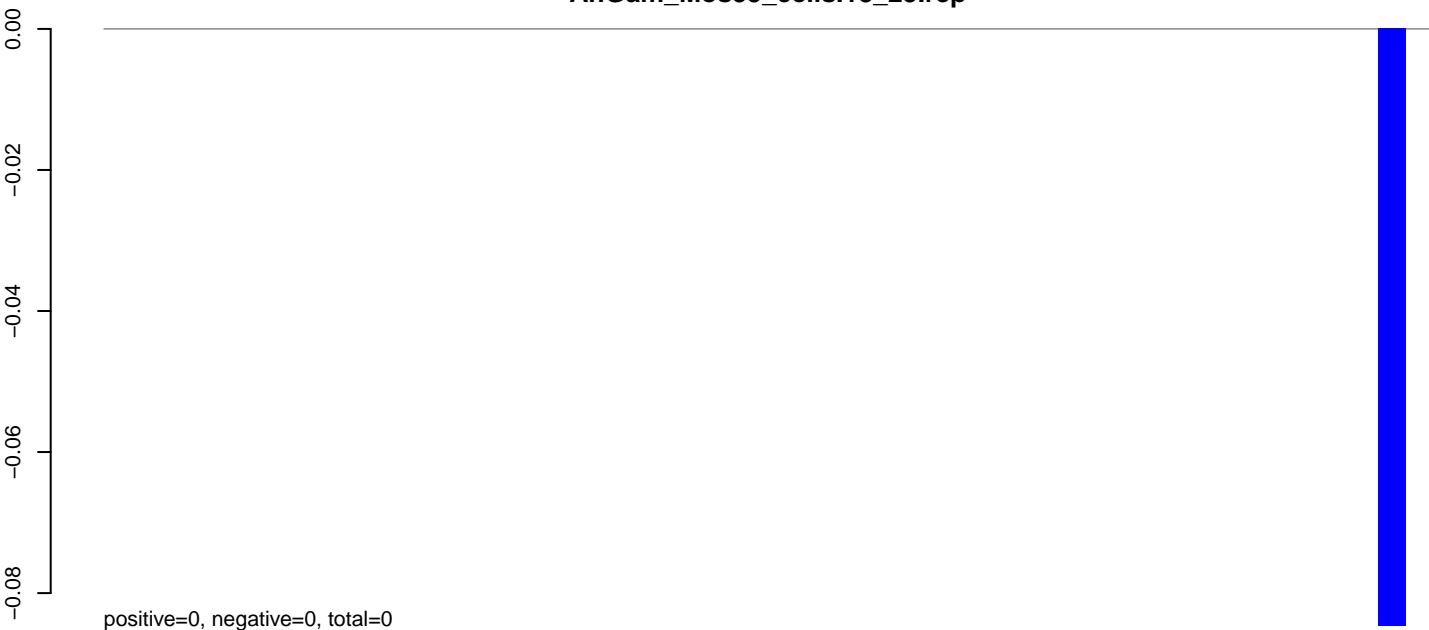
AnGam_Mos55_cells.24_35.rep



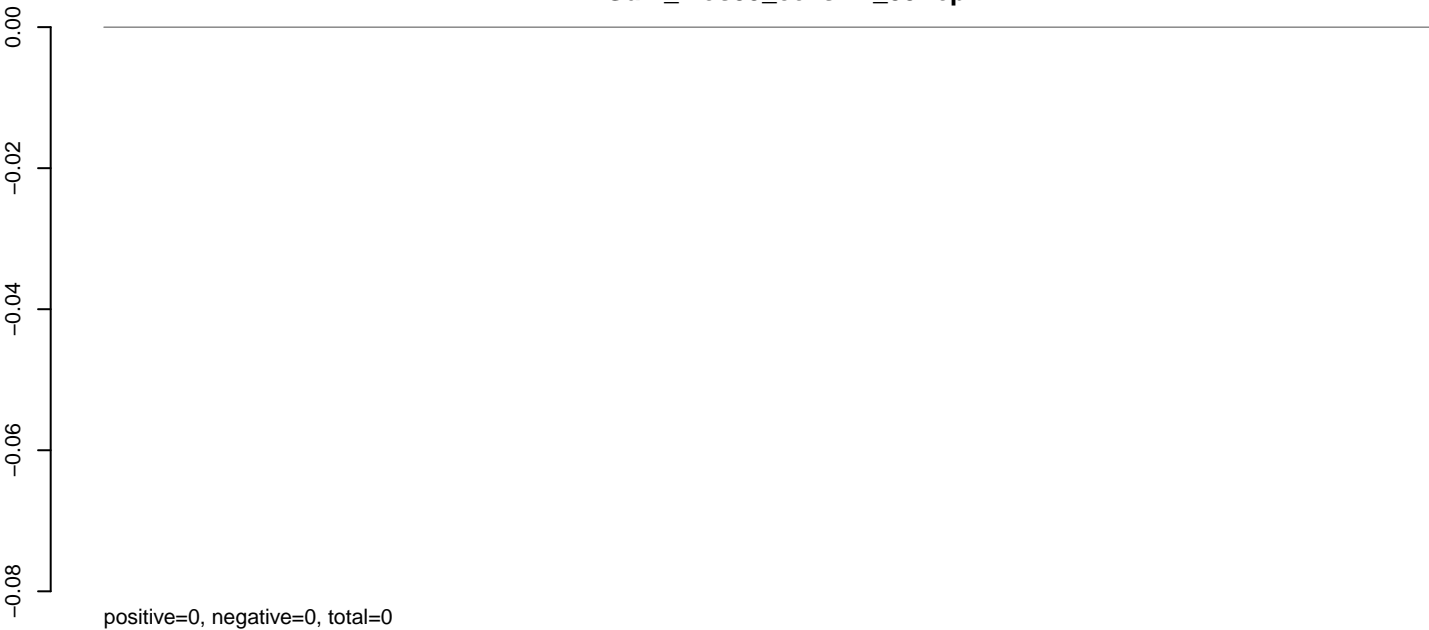
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



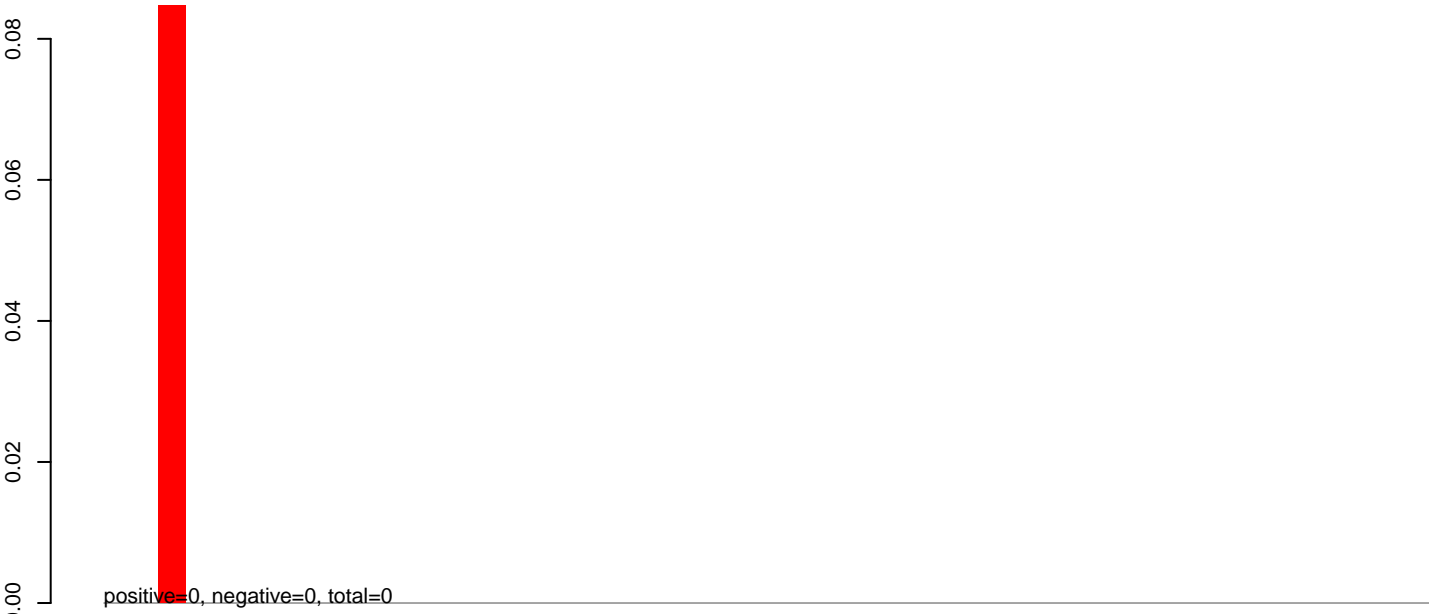
AnGam_Mos55_cells.24_35.rep



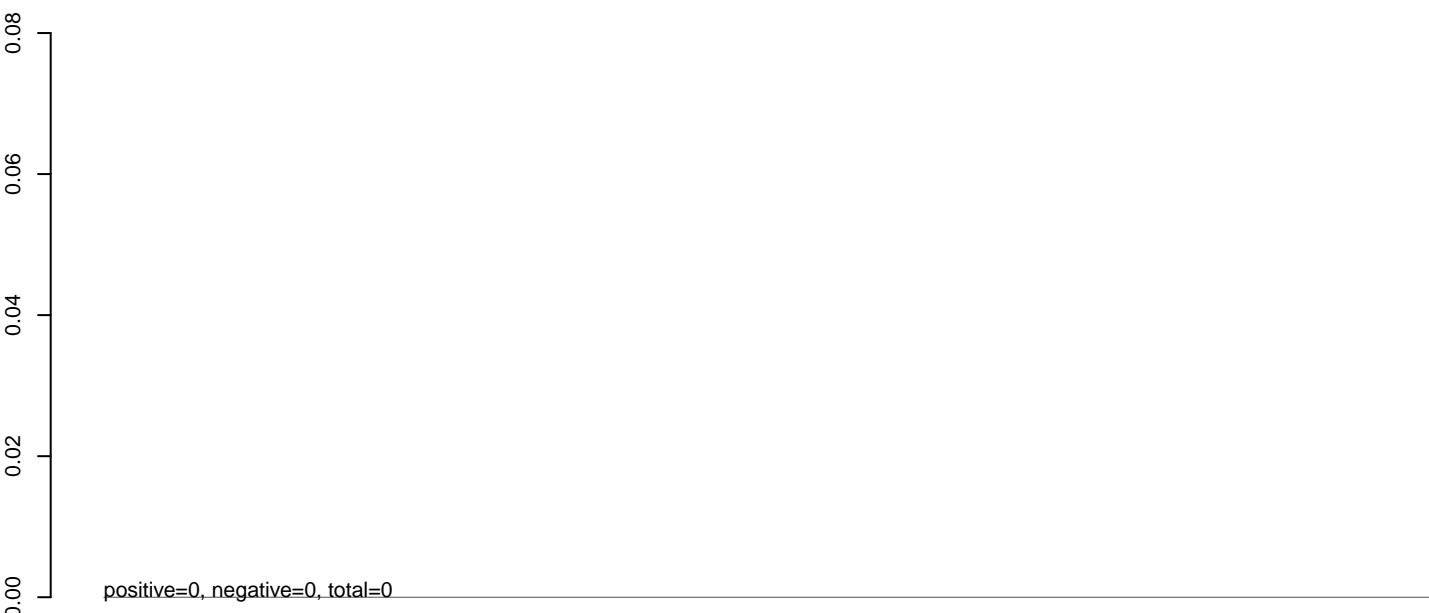
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



0 500 1000 1500 2000 2500

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

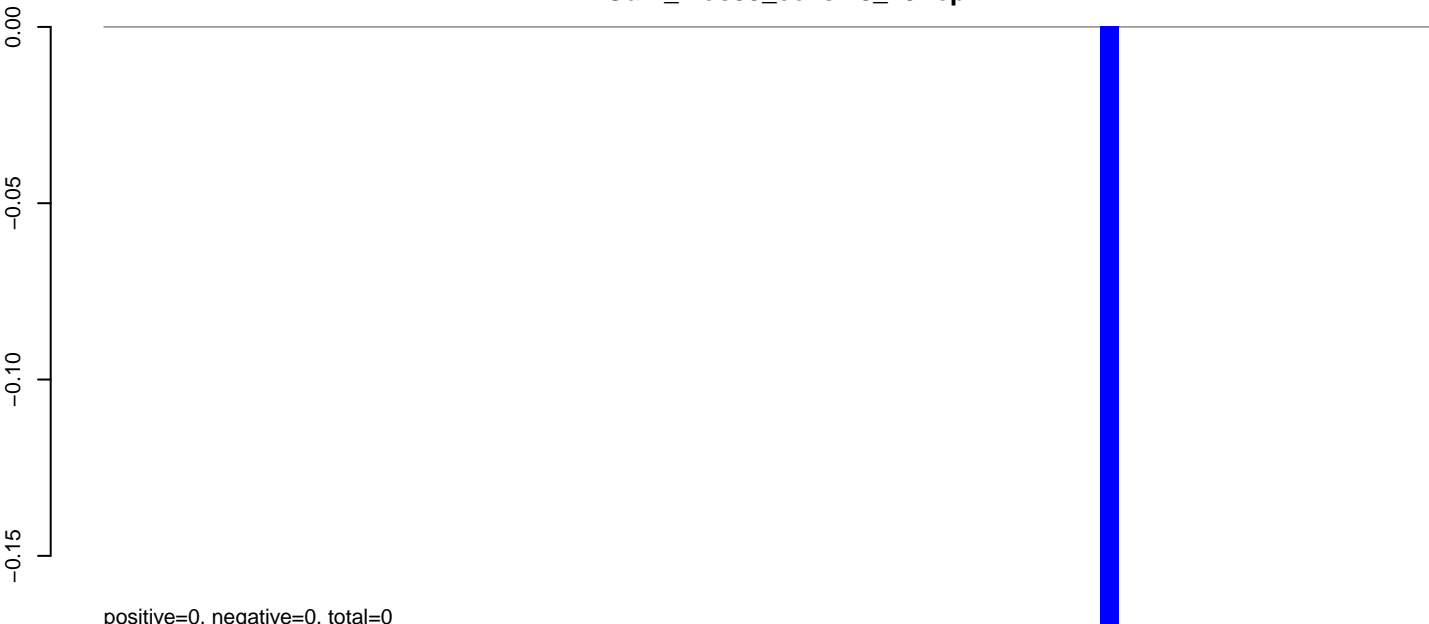
0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

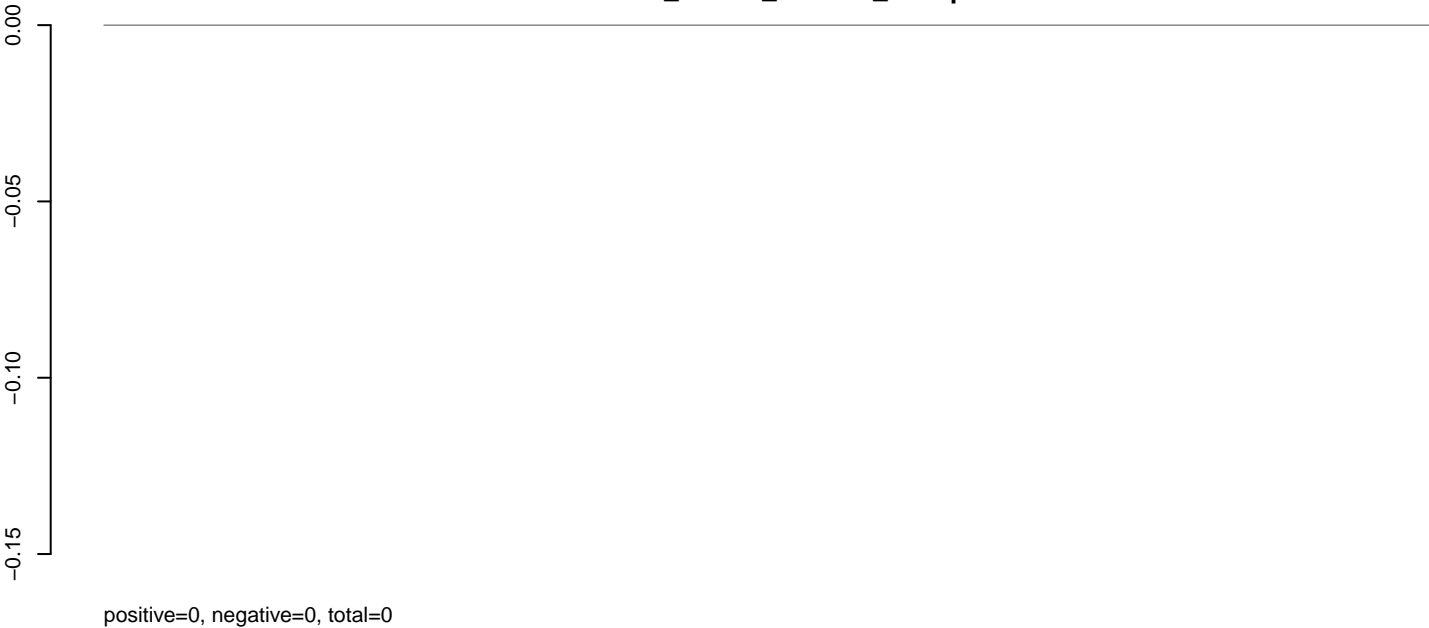
Window size=50, length=3595, TE@TF001358-I_Ele51-3595

0 500 1000 1500 2000 2500 3000 3500

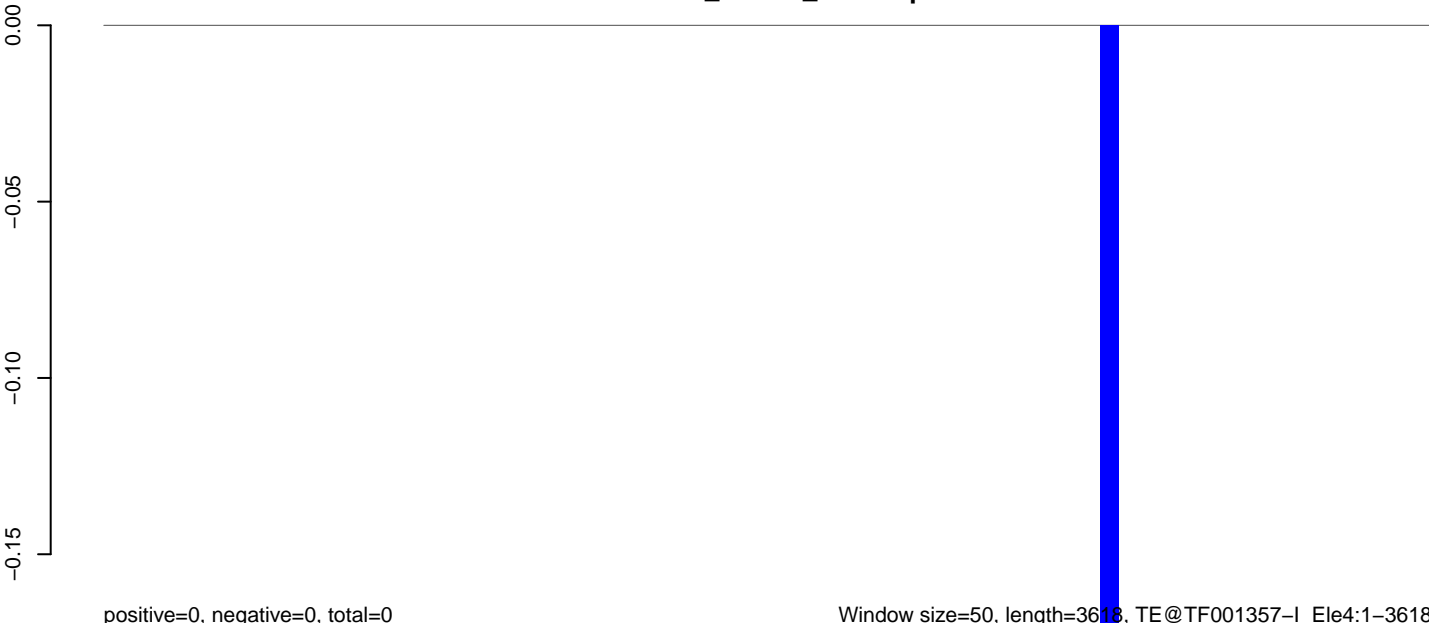
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



Window size=50, length=3618, TE@TF001357-I_Ele4:1-3618

0 500 1000 1500 2000 2500 3000 3500

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=5112, TE@TF001355-I_Element:1-5112

0 1000 2000 3000 4000 5000

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=823, TE@TE001238-Tc1_Ele7:1-823

200 400 600 800

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=631, TE@TF001220-mar_Ele26:1-631

100 200 300 400 500 600

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=768, TE@TF001216-mar_Ele27:1-768

200 400 600 800

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

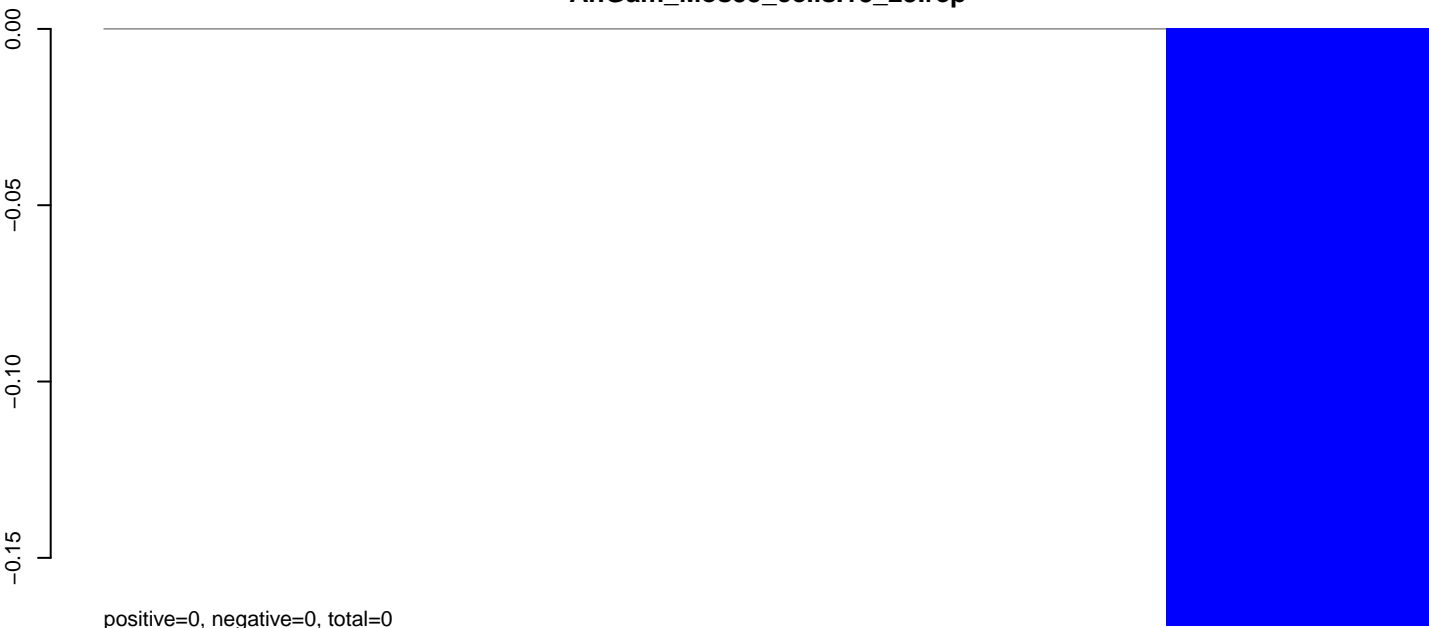
0.00
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-0.04
-0.06
-0.08

positive=0, negative=0, total=0

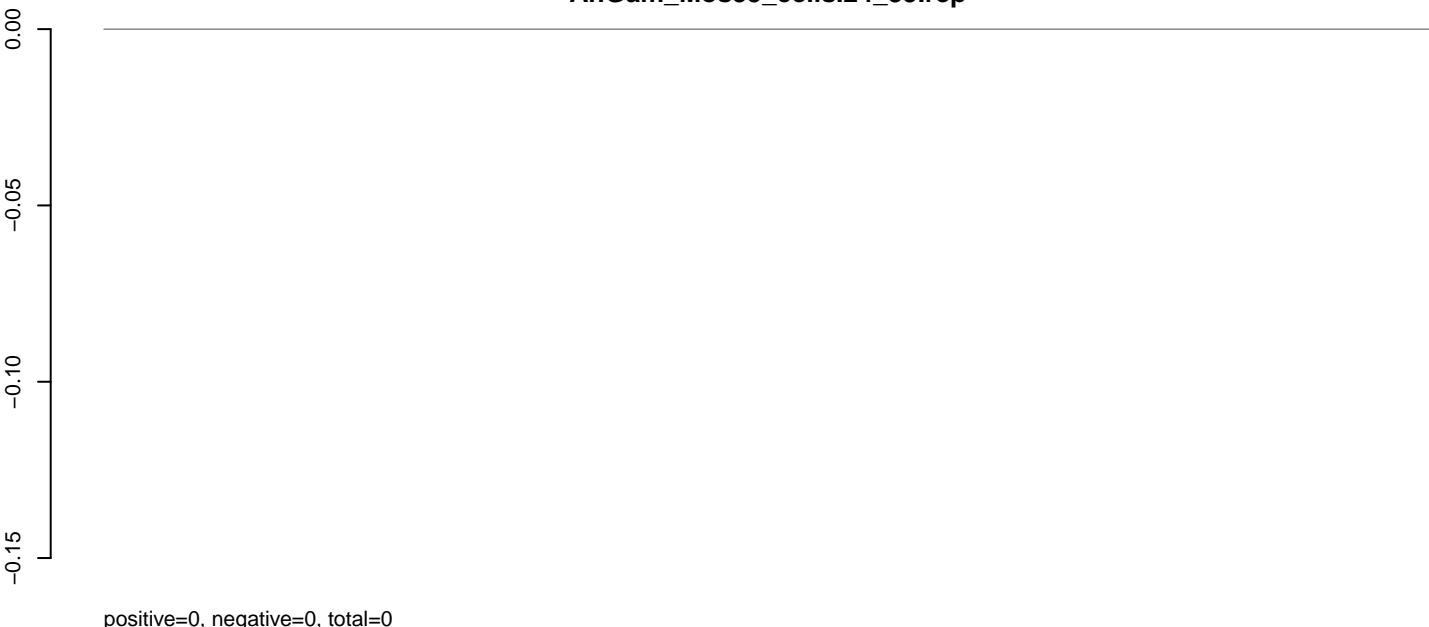
Window size=50, length=1278, TE@TF001207-mar_Ele28:1-1278

200 400 600 800 1000 1200

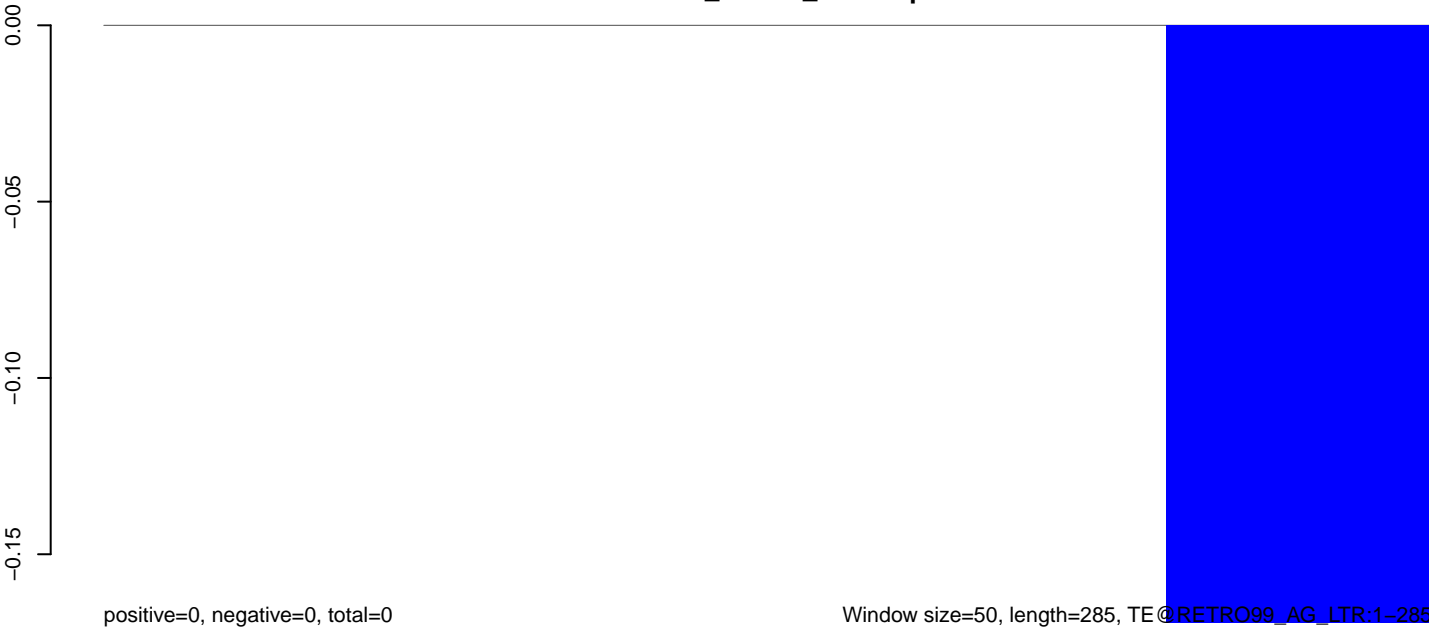
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

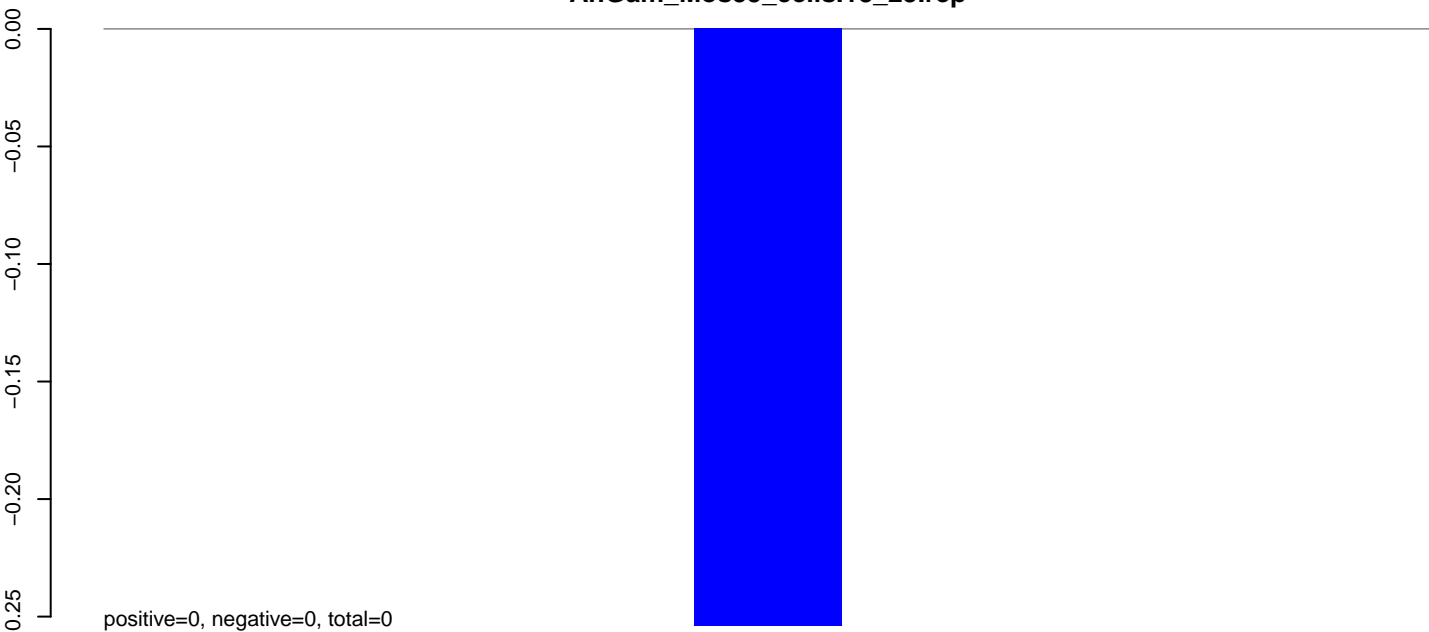


AnGam_Mos55_cells.rep

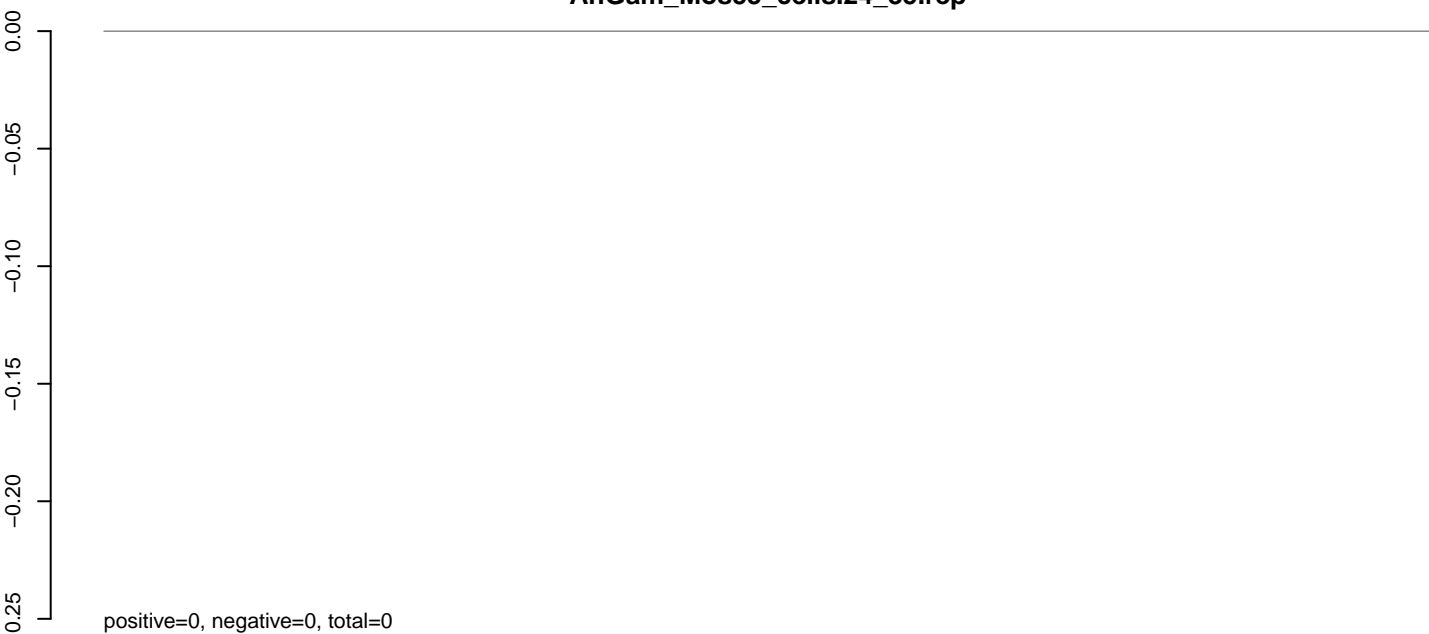


Window size=50, length=285, TE@RETRO99_AG_LTR:1-285

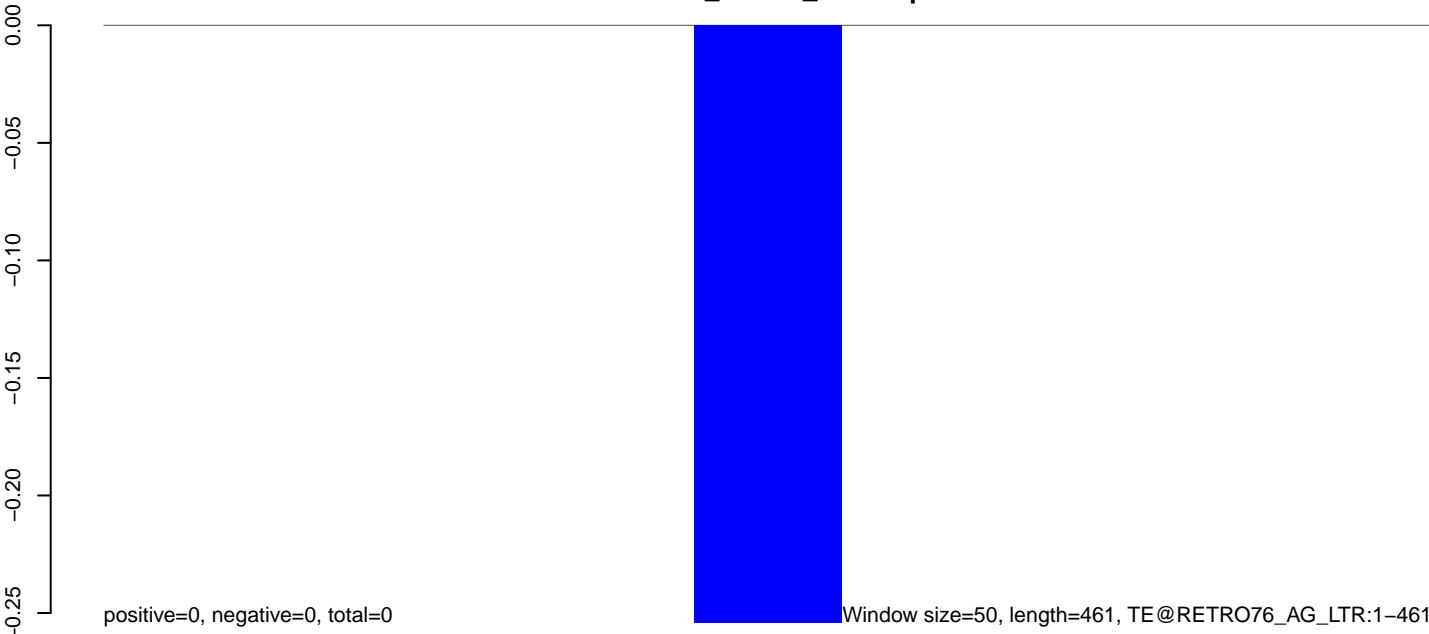
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

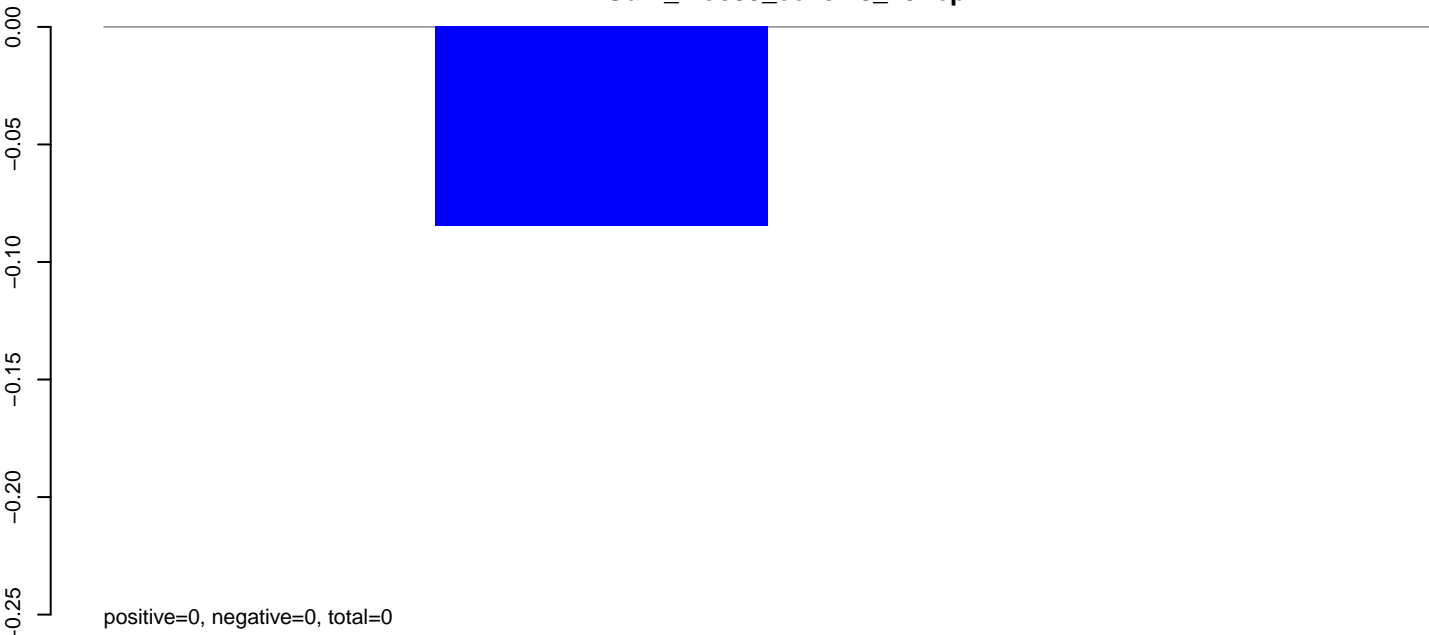


AnGam_Mos55_cells.rep

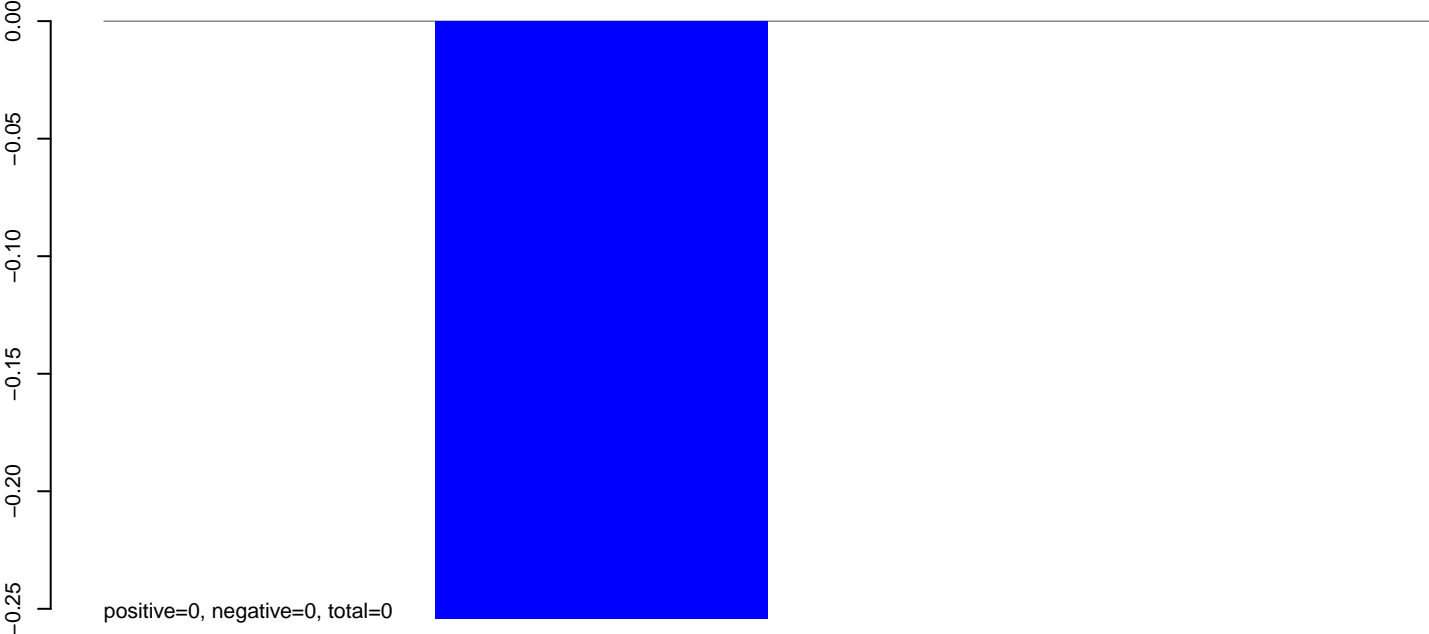


100 200 300 400 500

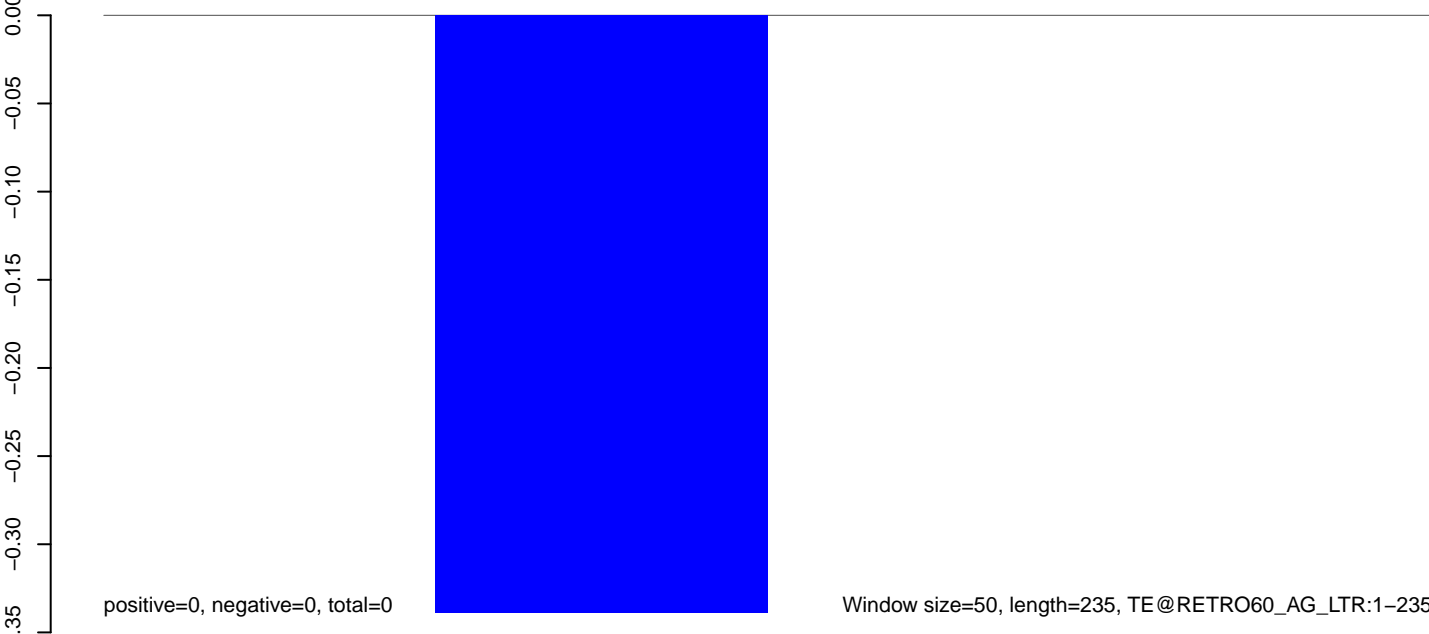
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

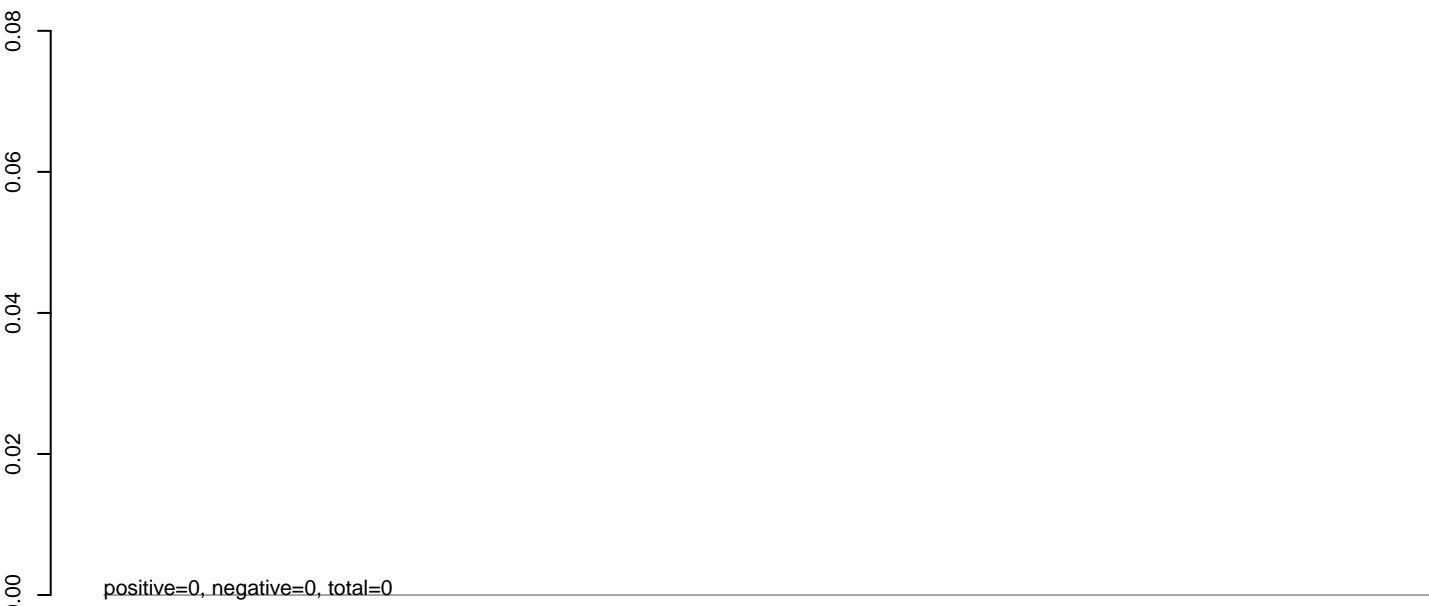


AnGam_Mos55_cells.rep

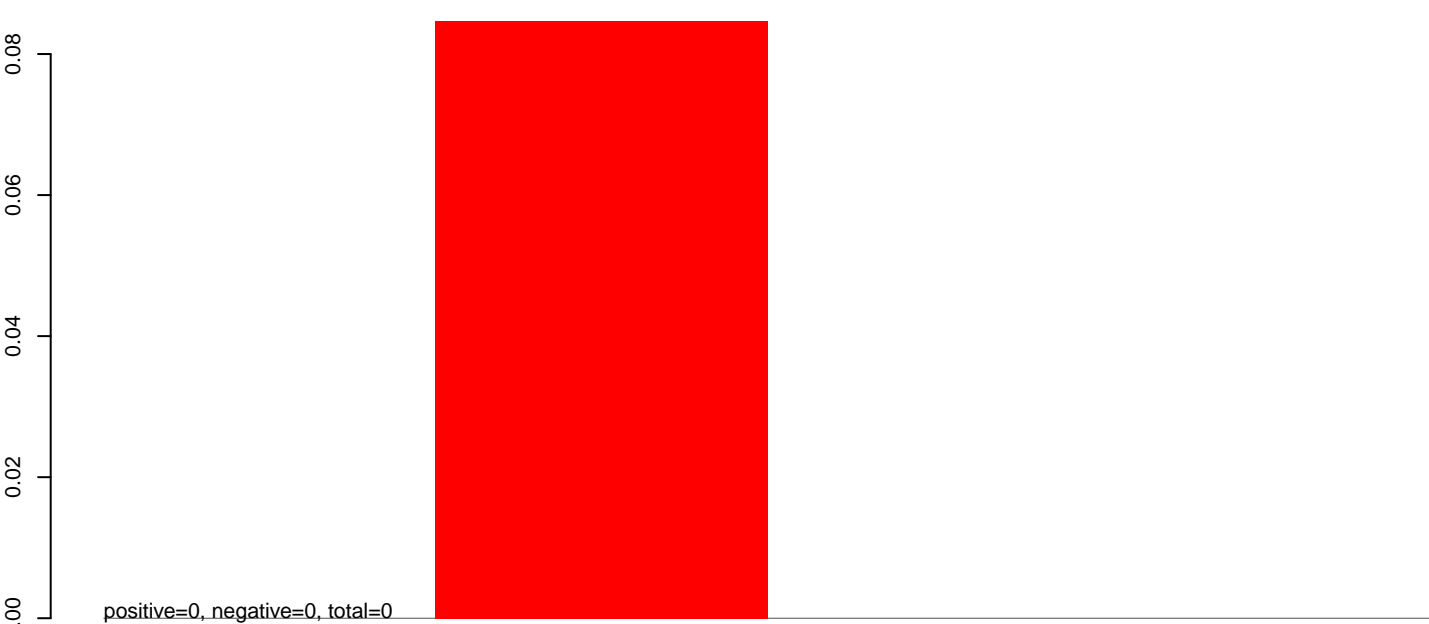


Window size=50, length=235, TE@RETRO60_AG_LTR:1-235

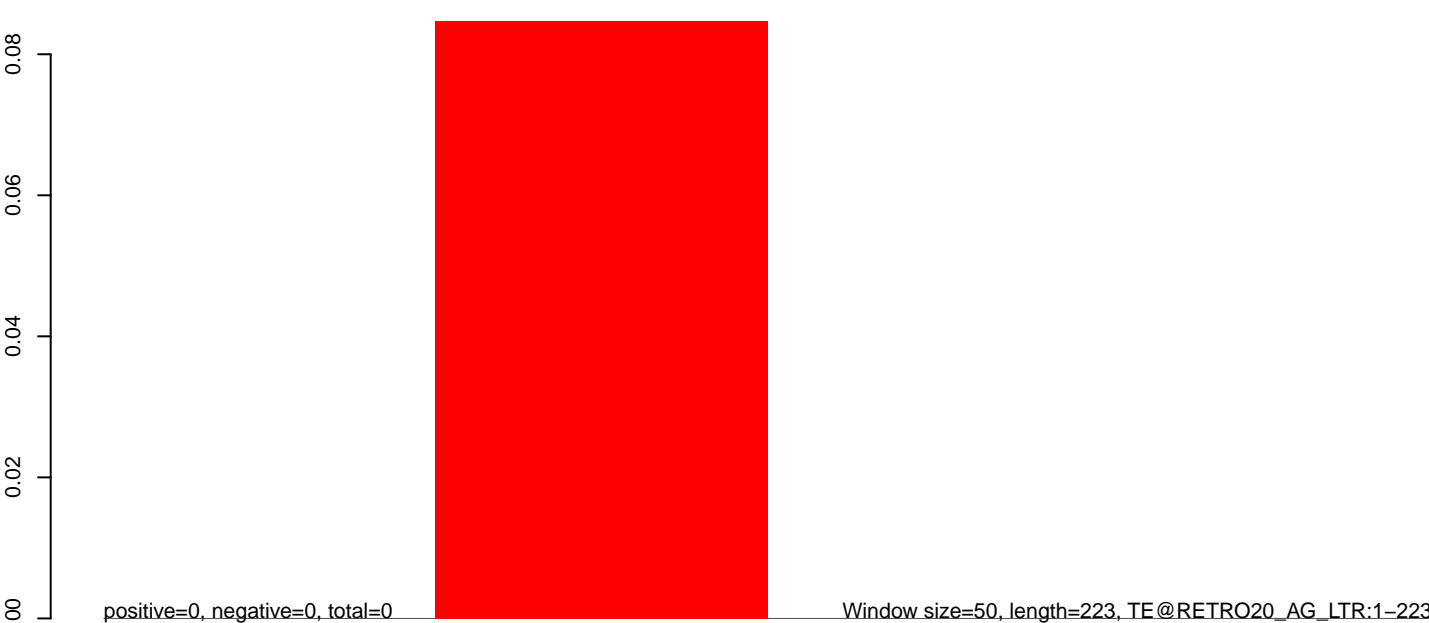
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

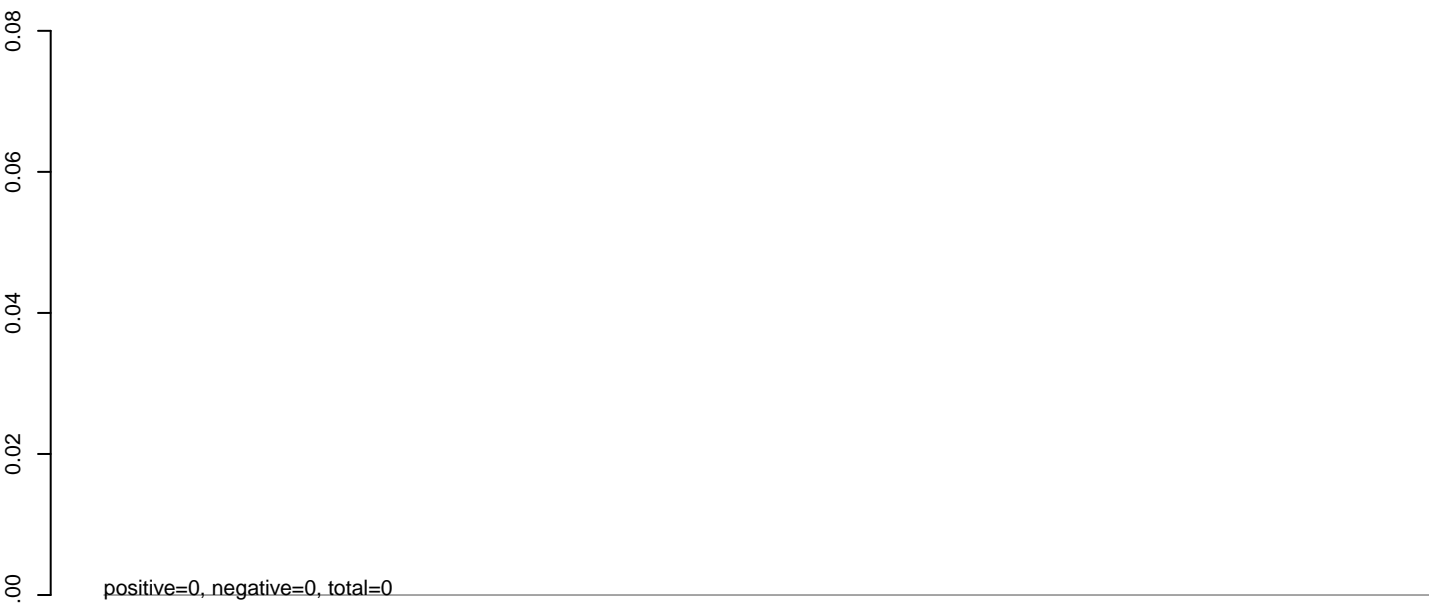
0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

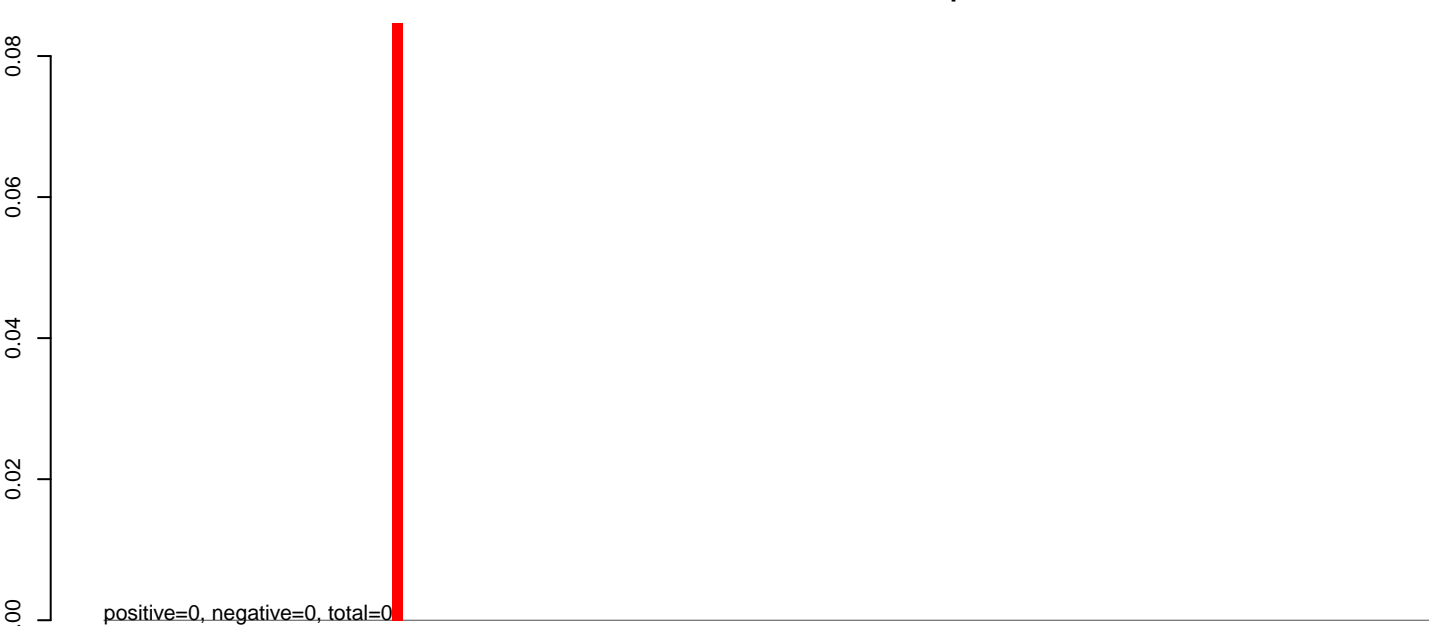
Window size=50, length=205, TE@RETRO16_AG_LTR:1-205

50 100 150 200 250

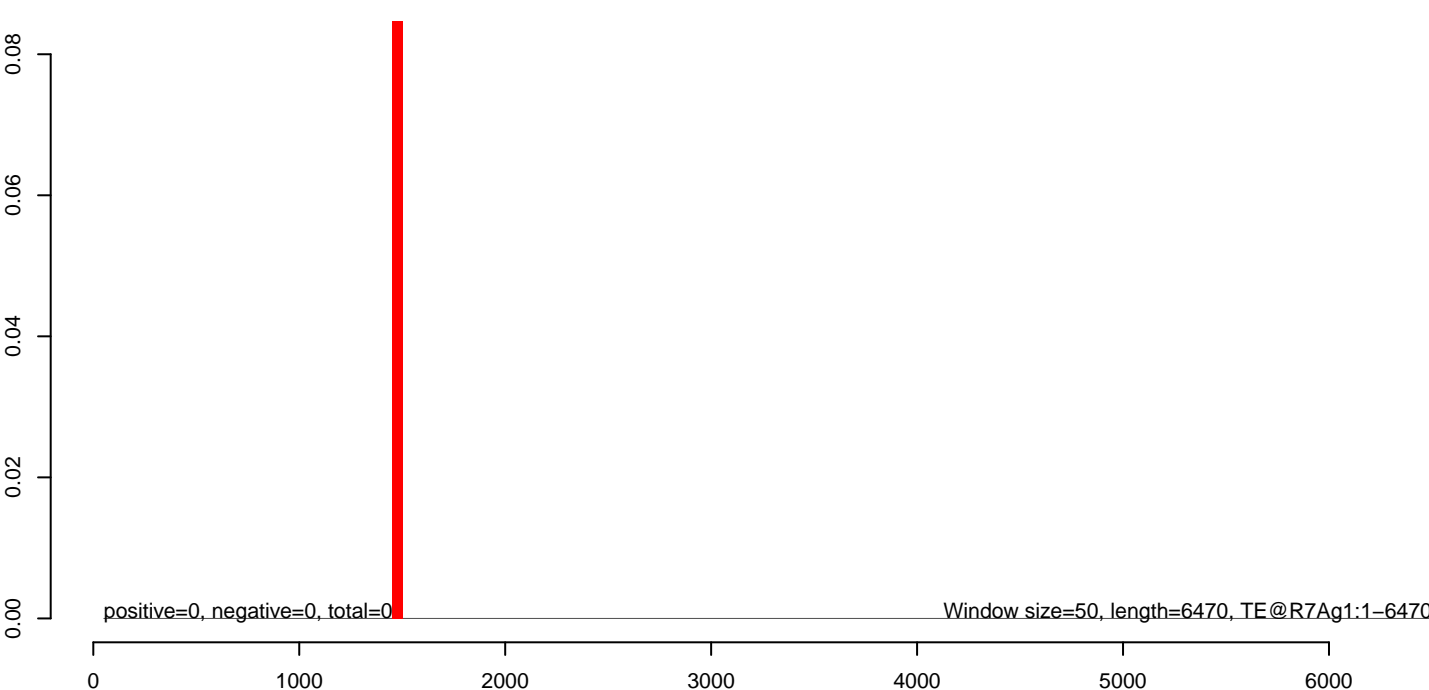
AnGam_Mos55_cells.18_23.rep



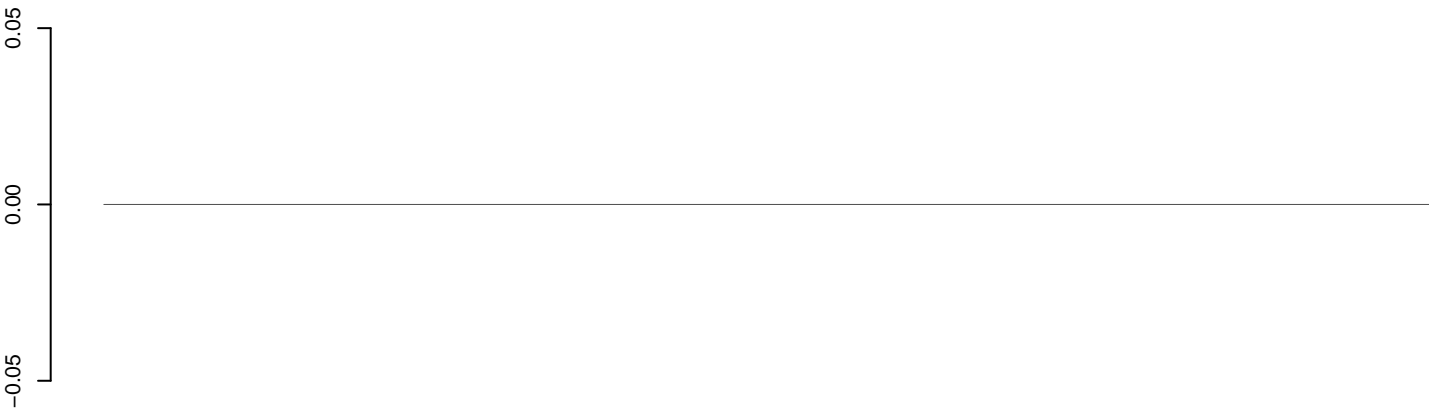
AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep

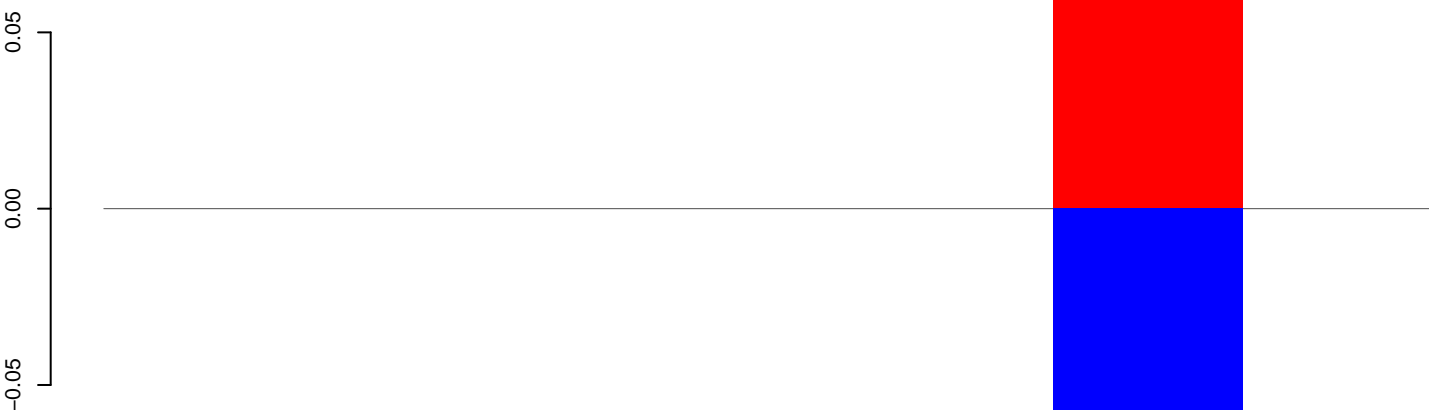


AnGam_Mos55_cells.18_23.rep



positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



positive=0, negative=0, total=0

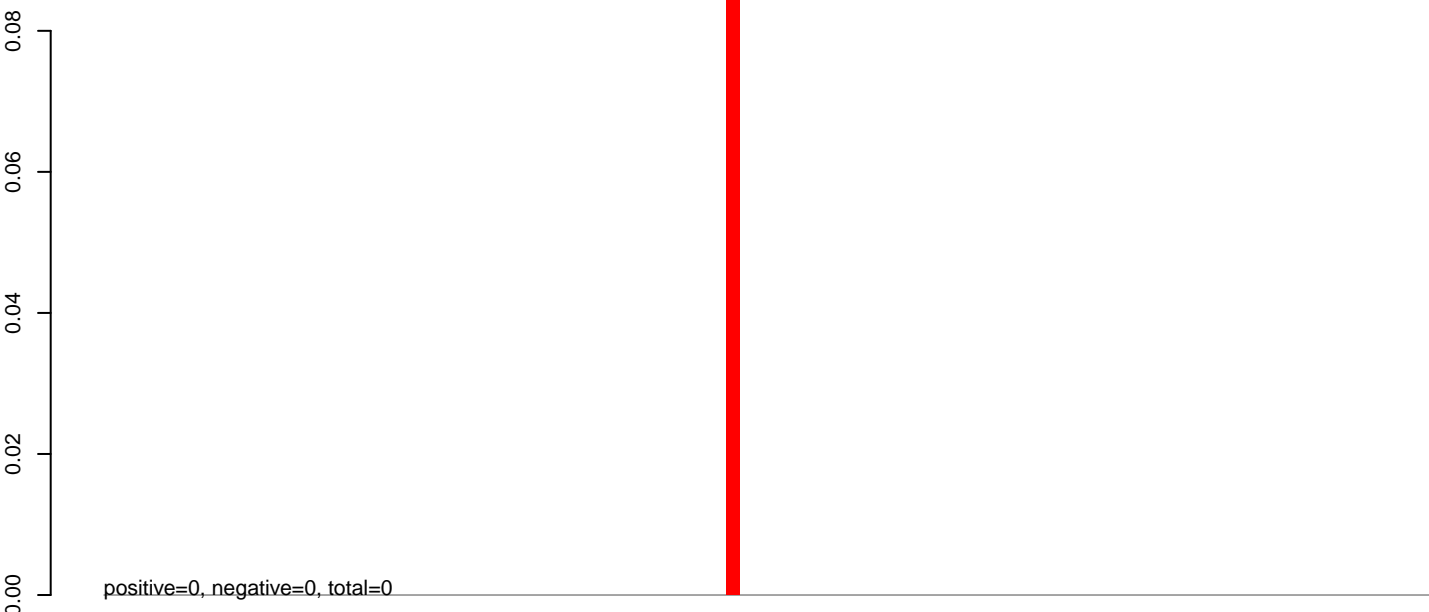
AnGam_Mos55_cells.rep



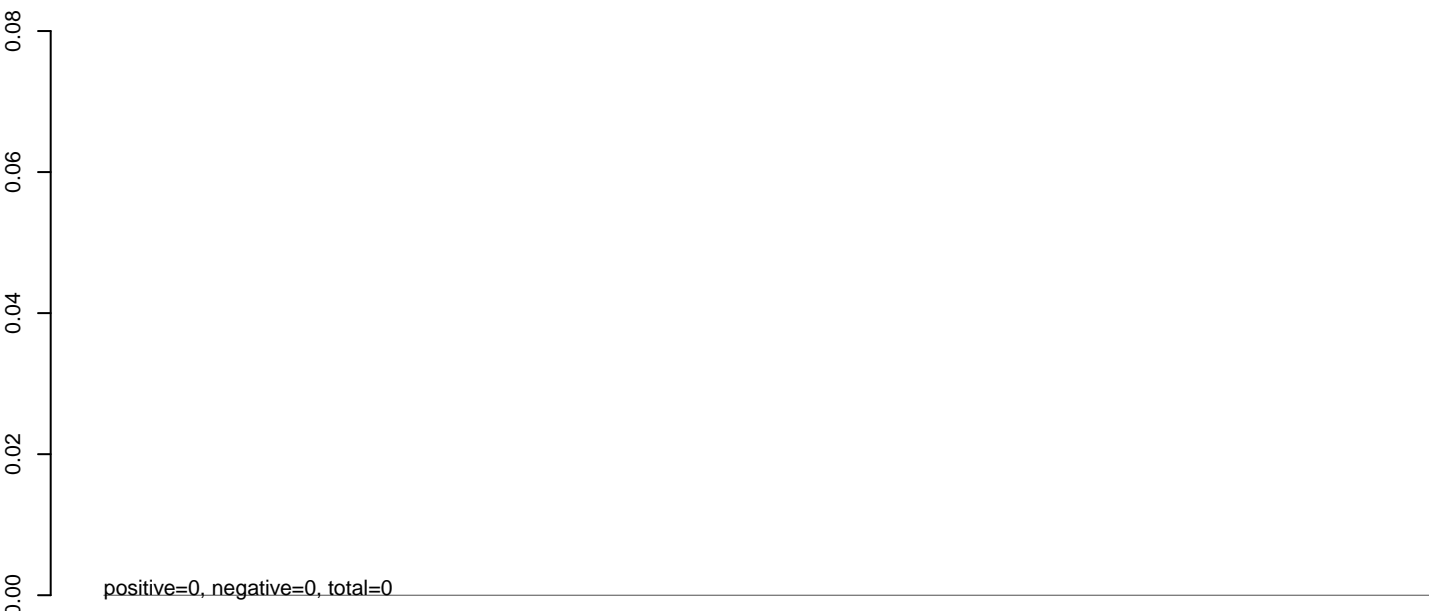
positive=0, negative=0, total=0

Window size=50, length=357, TE@MARINERN3a_AG:1-357

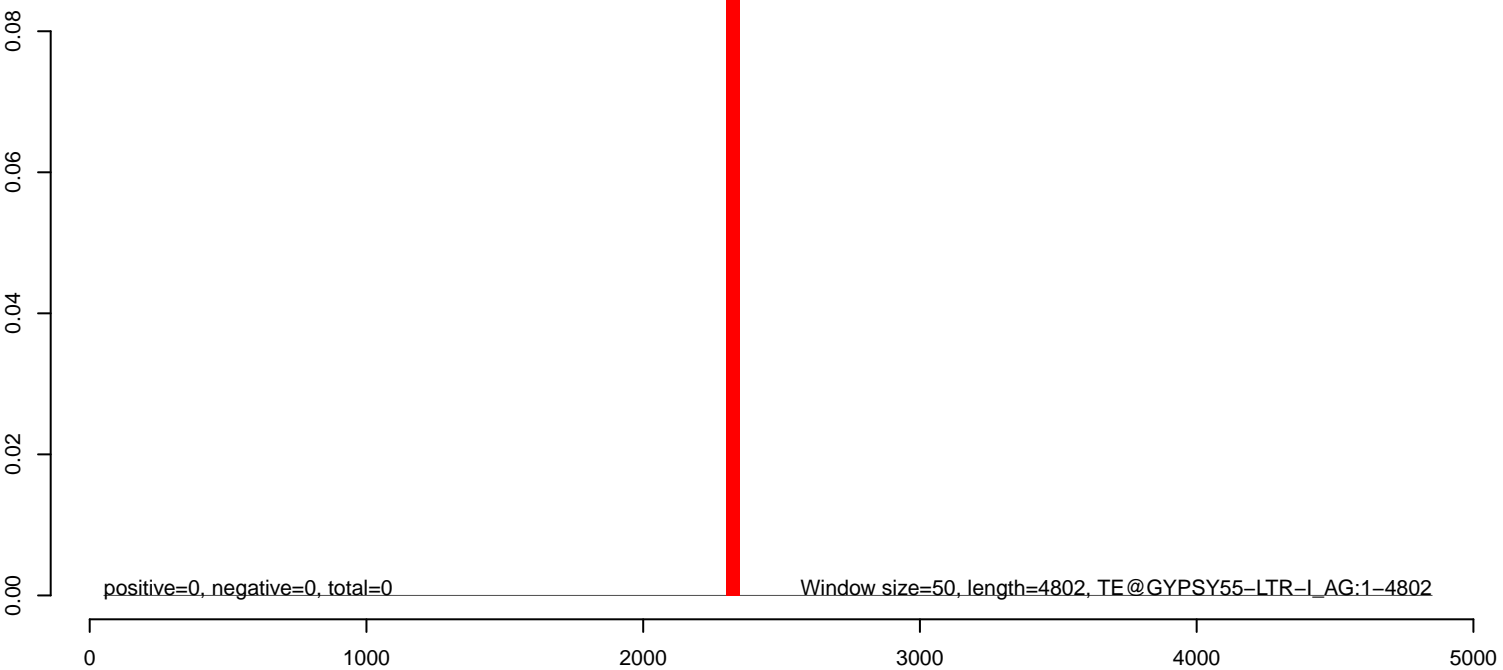
AnGam_Mos55_cells.18_23.rep



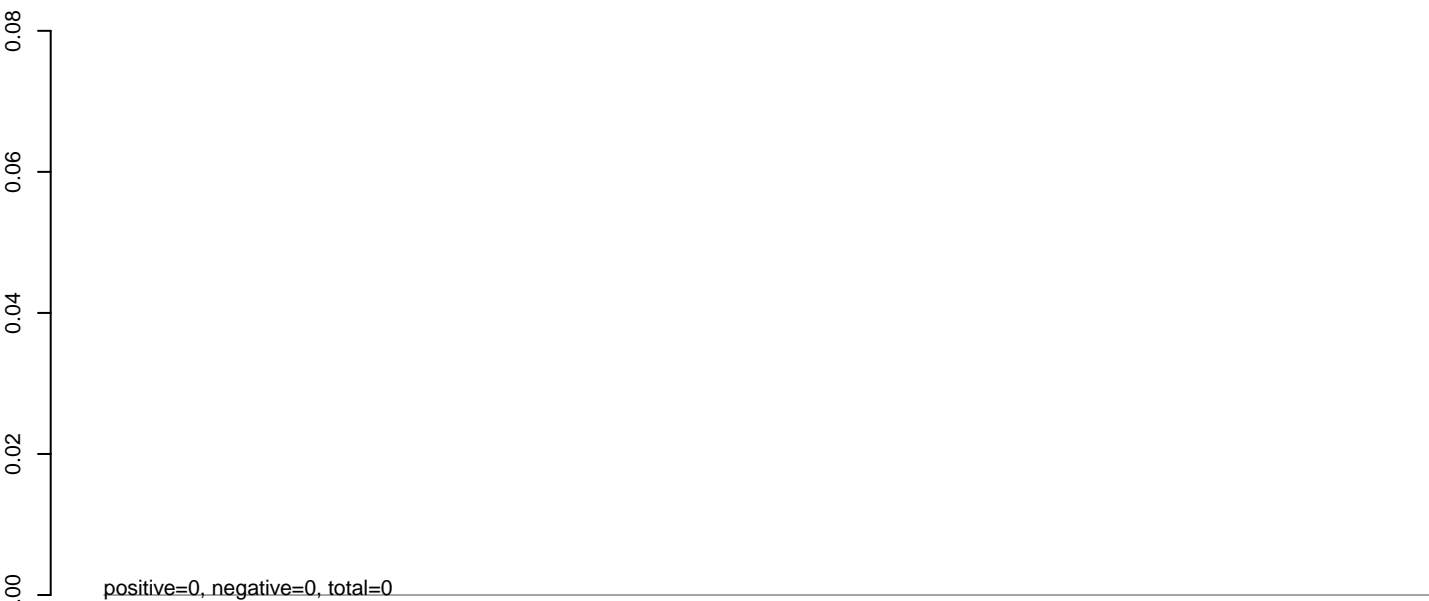
AnGam_Mos55_cells.24_35.rep



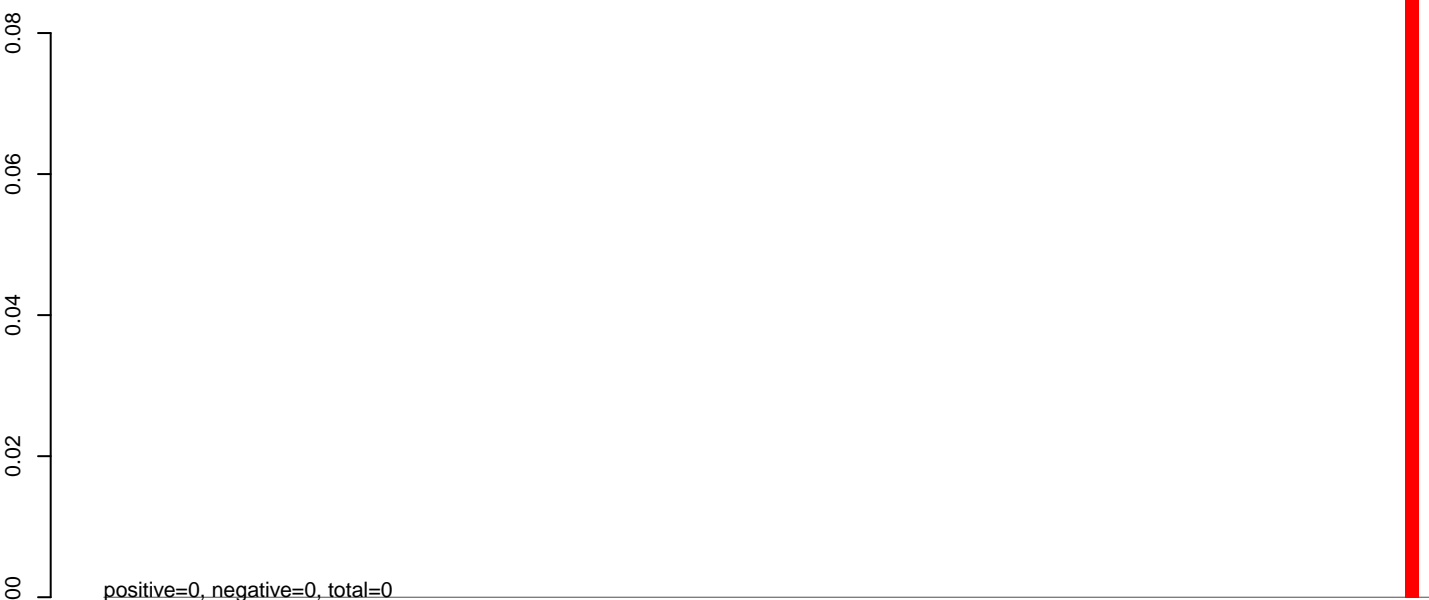
AnGam_Mos55_cells.rep



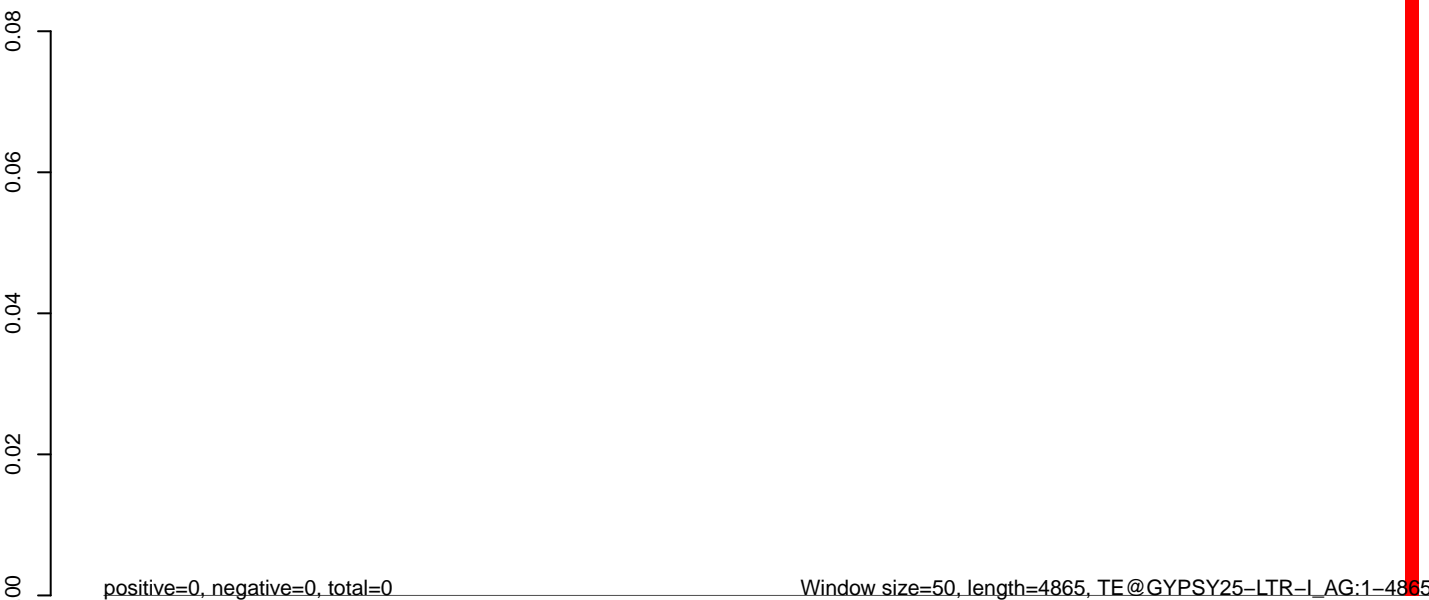
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

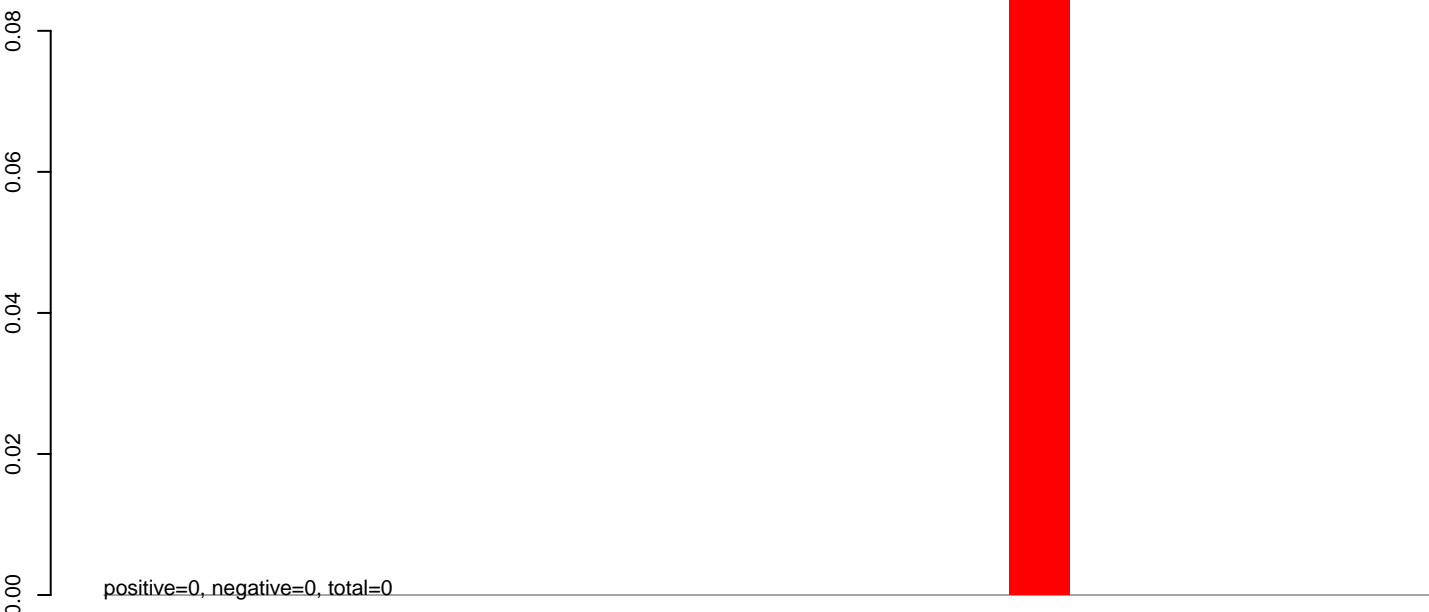


AnGam_Mos55_cells.rep

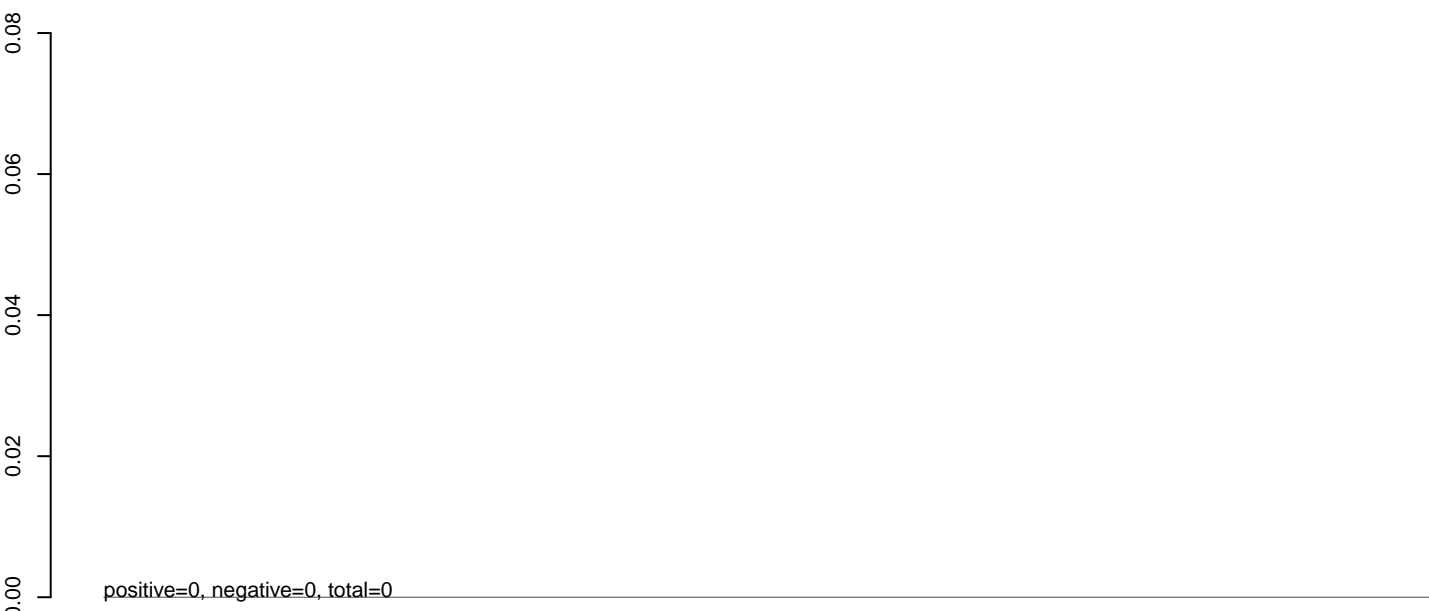


0 1000 2000 3000 4000 5000

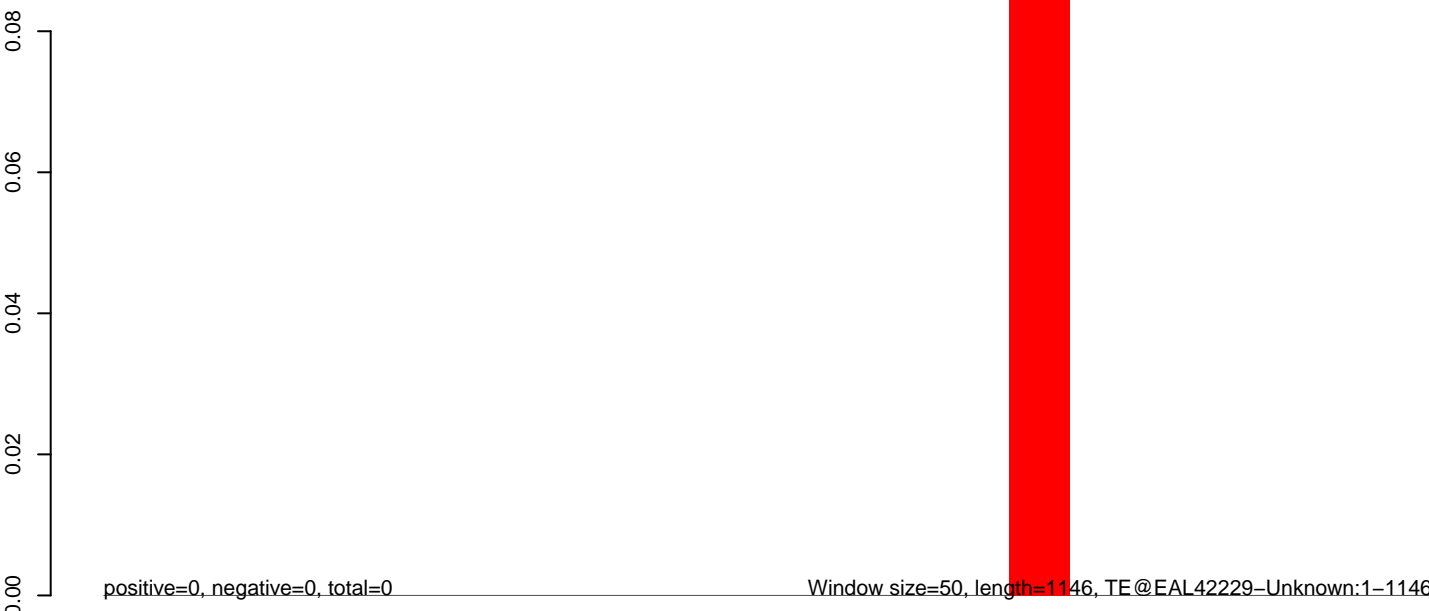
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=477, TE@EAL42162-Unknown:1-477

100 200 300 400 500

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

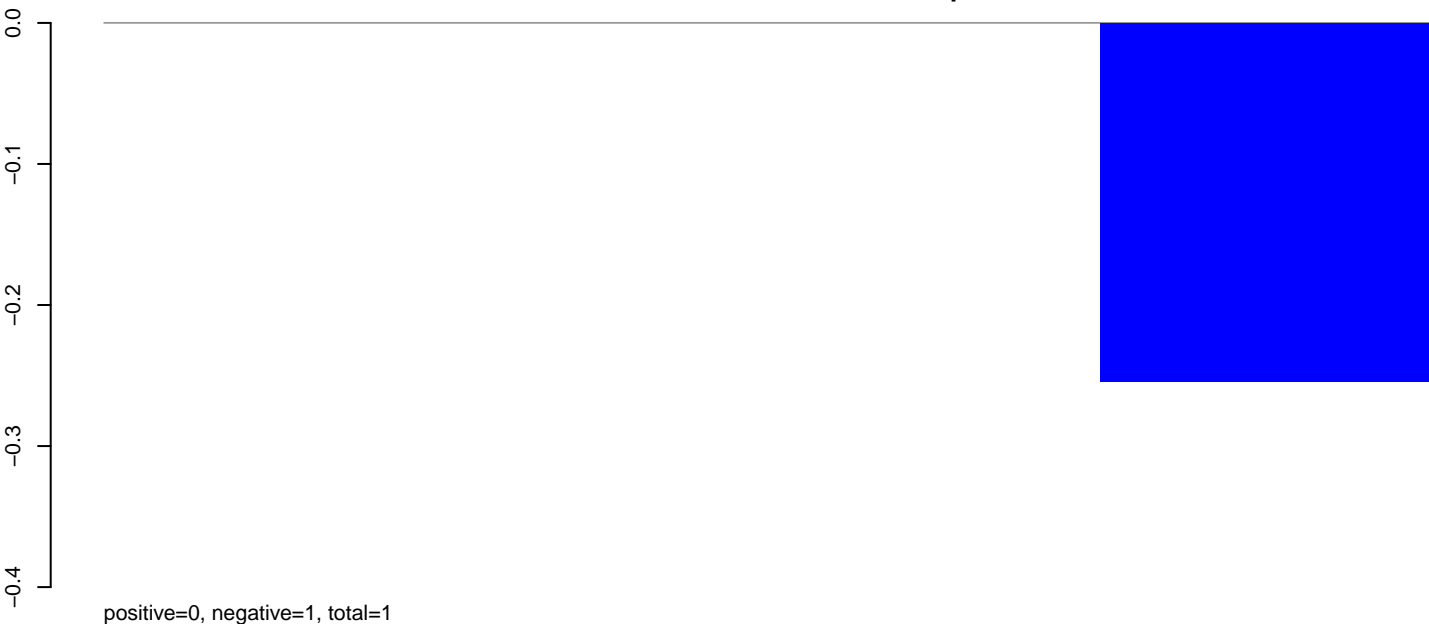
0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

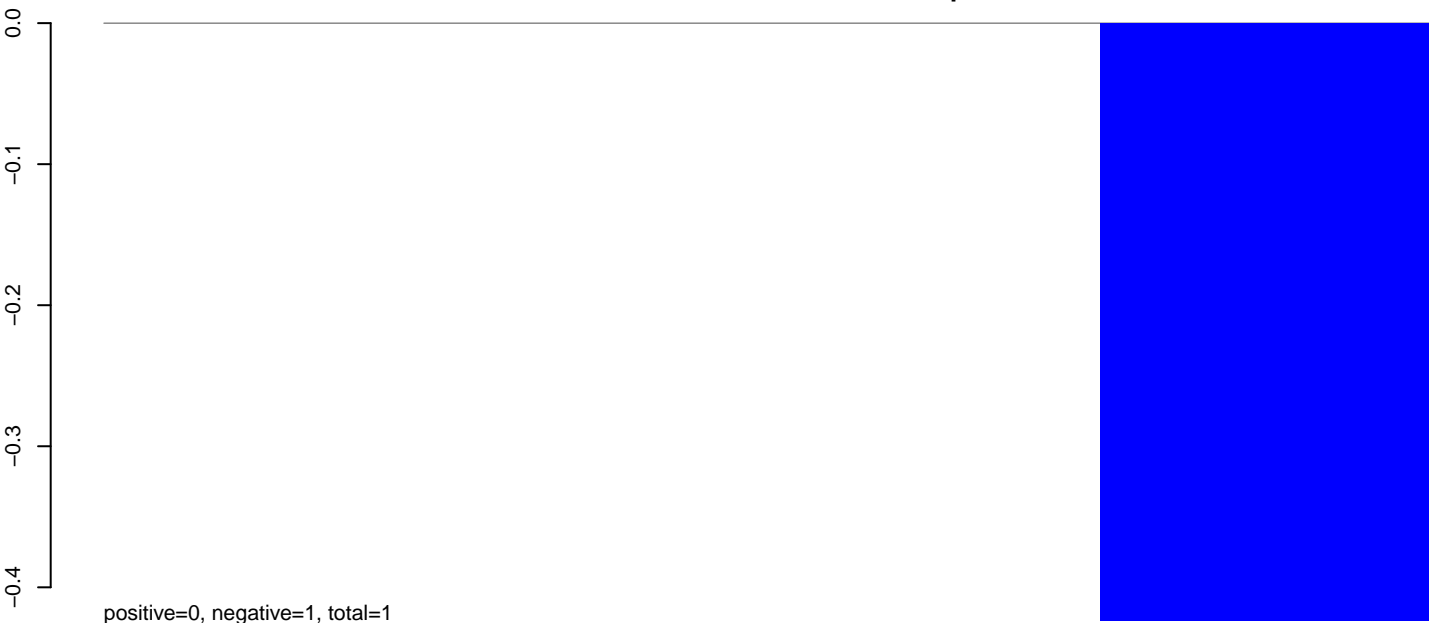
Window size=50, length=1101, TE@EAL40346-Unknown:1-1101

200 400 600 800 1000

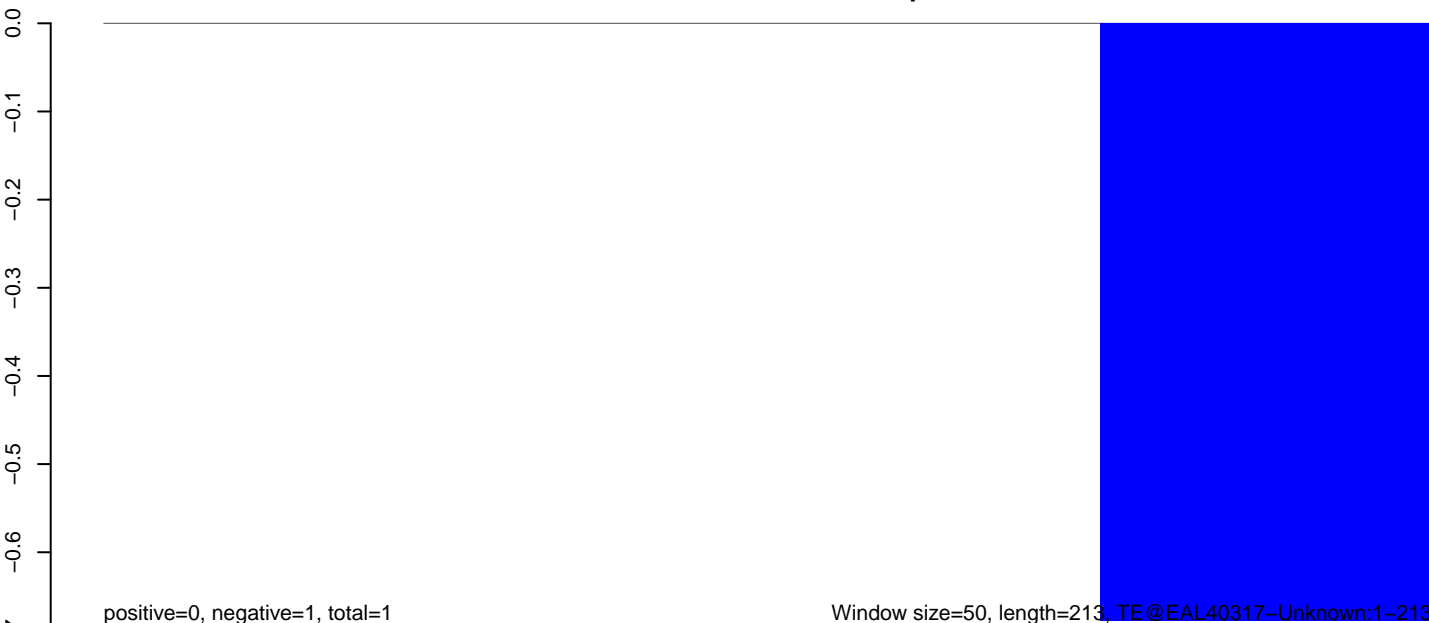
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



Window size=50, length=213, TE@EAL40317-Unknown:1-213

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=408, TE@EAL40183-Unknown:1-408

100

200

300

400



AnGam_Mos55_cells.18_23.rep

0.00
-0.05
-0.10
-0.15
-0.20
-0.25
-0.30
-0.35

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.05
-0.10
-0.15
-0.20
-0.25
-0.30
-0.35

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

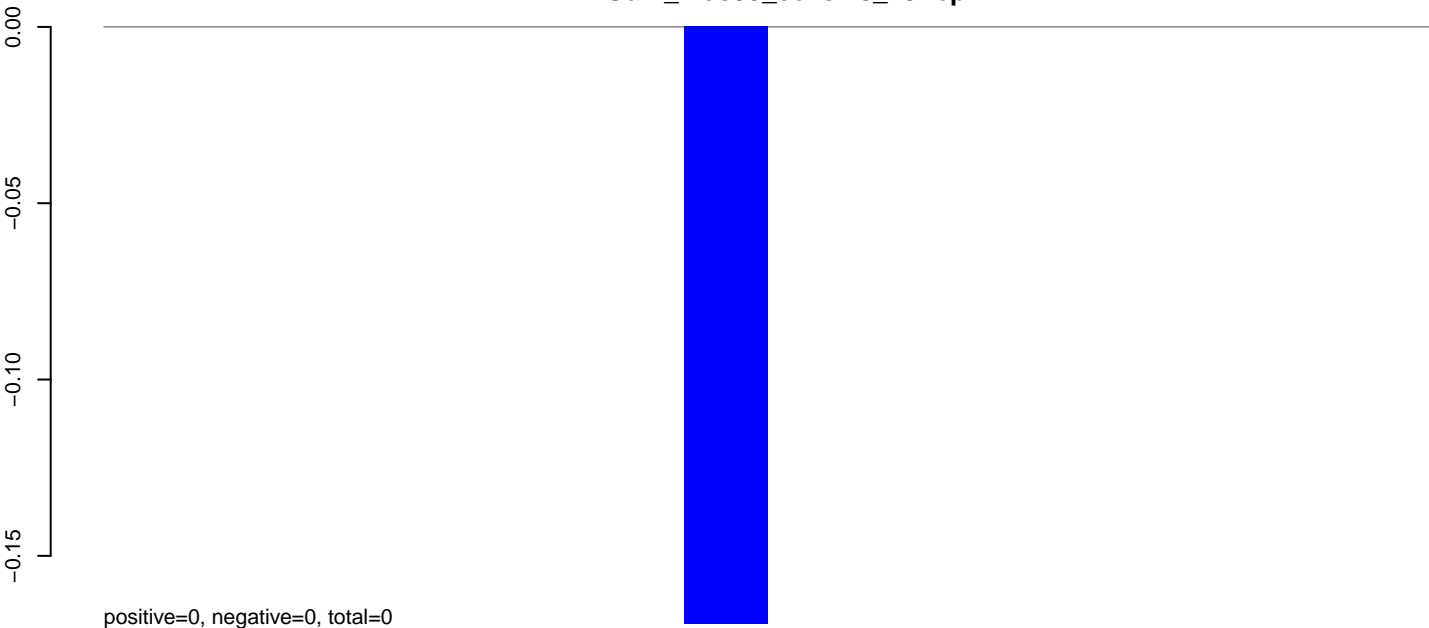
0.00
-0.05
-0.10
-0.15
-0.20
-0.25
-0.30
-0.35

positive=0, negative=0, total=0

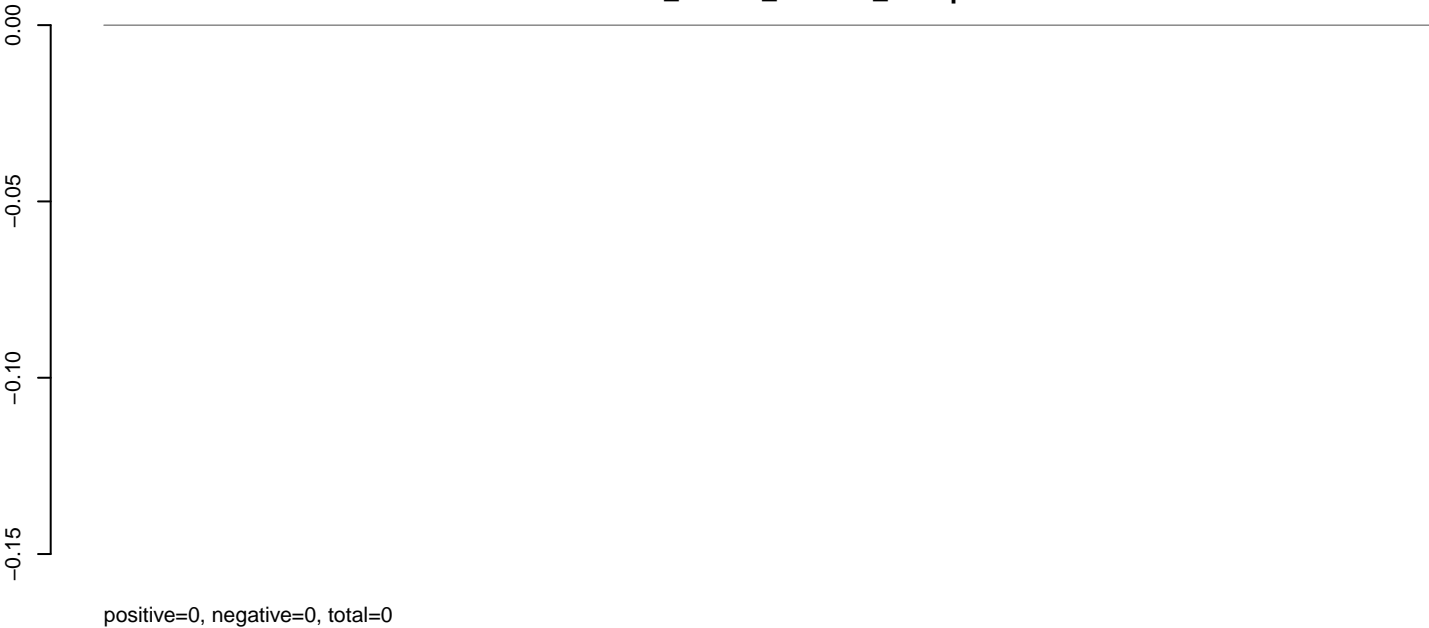
Window size=50, length=1086, TE@EAL39774-Unknown:1-1086

200 400 600 800 1000

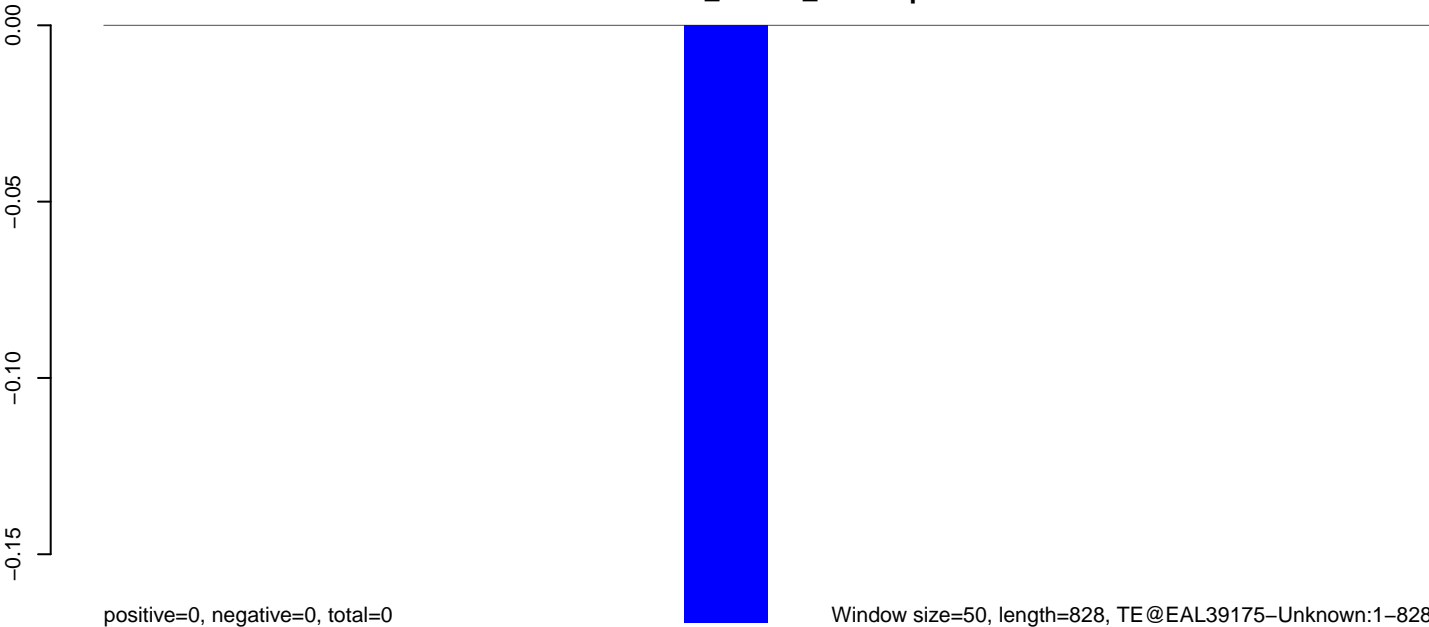
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



200 400 600 800

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

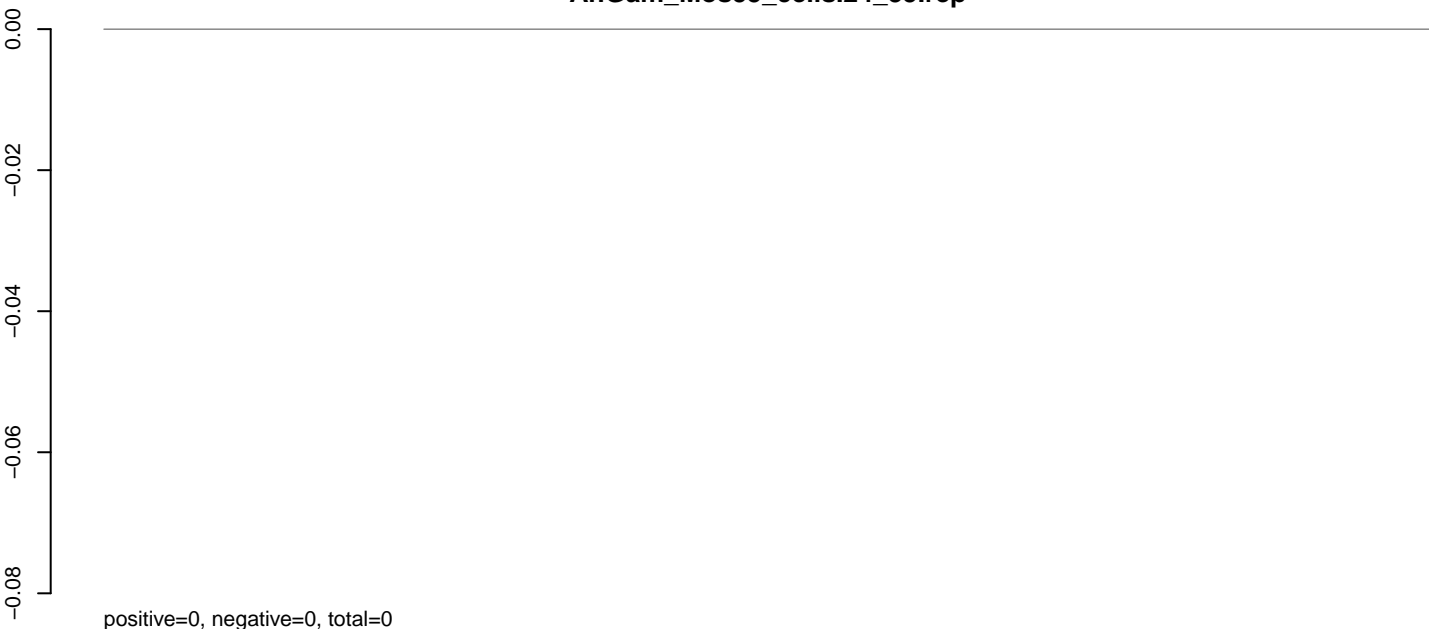
Window size=50, length=399, TE@EAA45679-Unknown:1-399

50 100 150 200 250 300 350 400

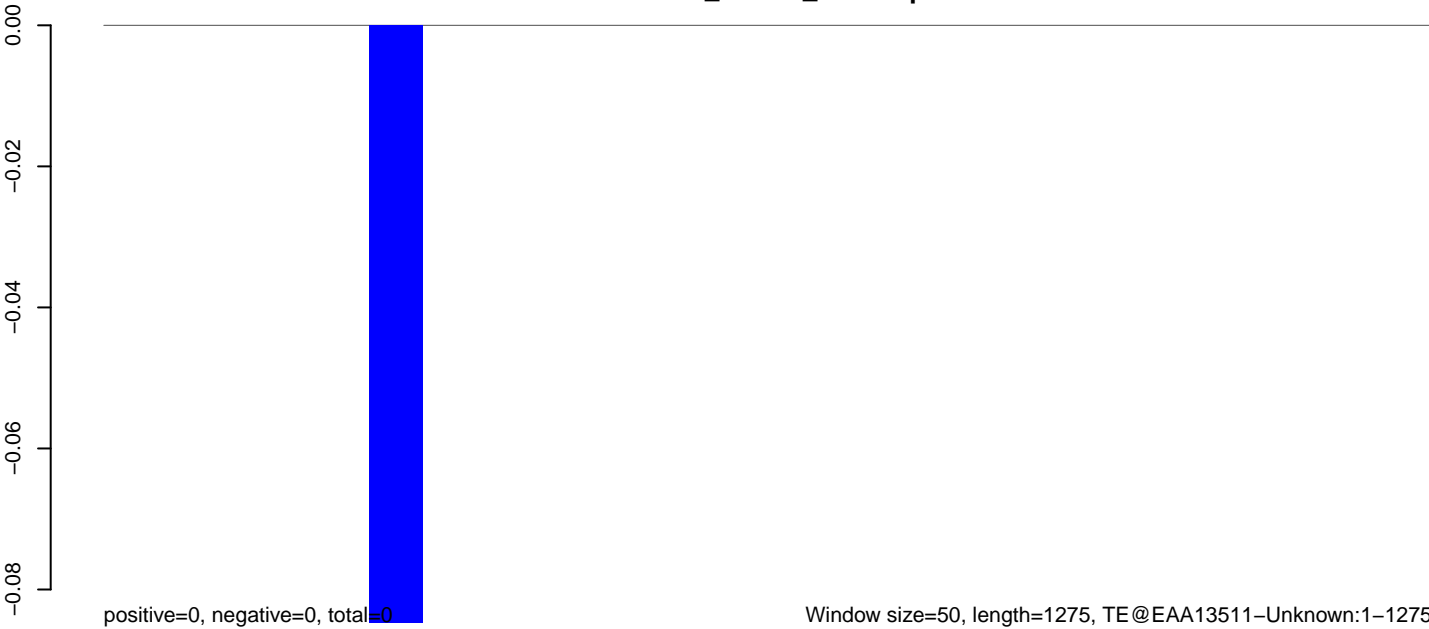
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



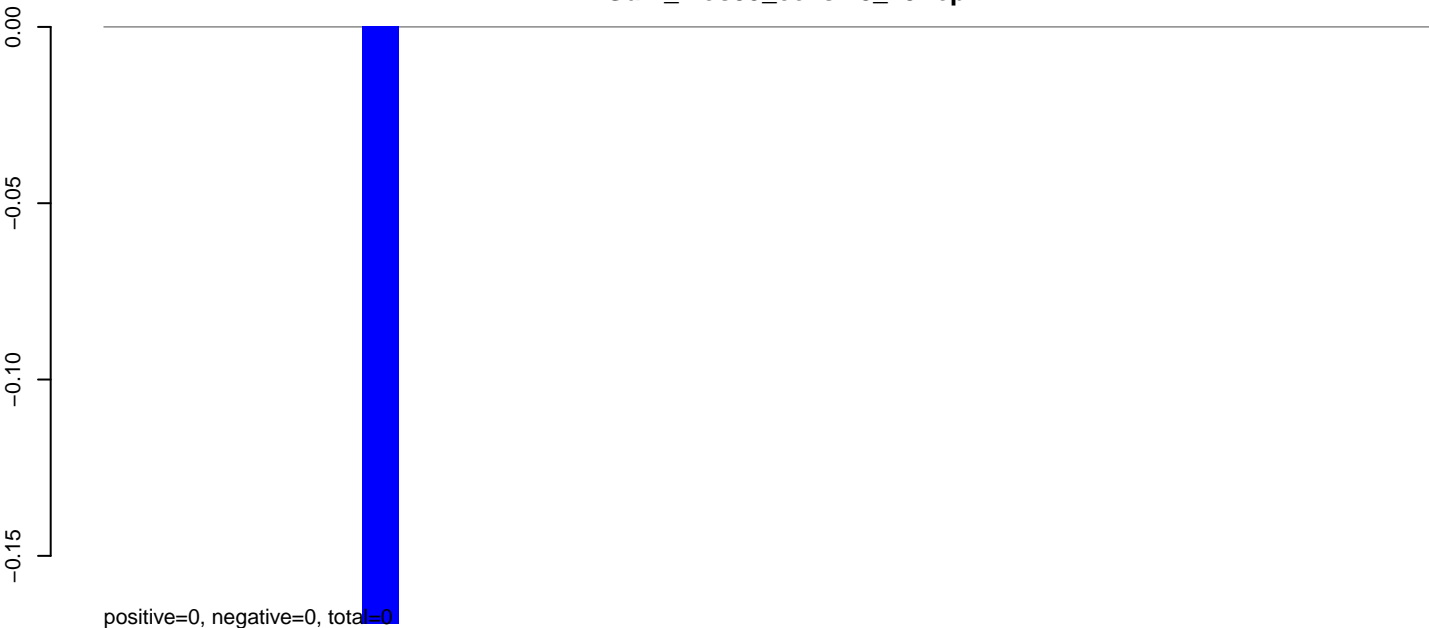
AnGam_Mos55_cells.rep



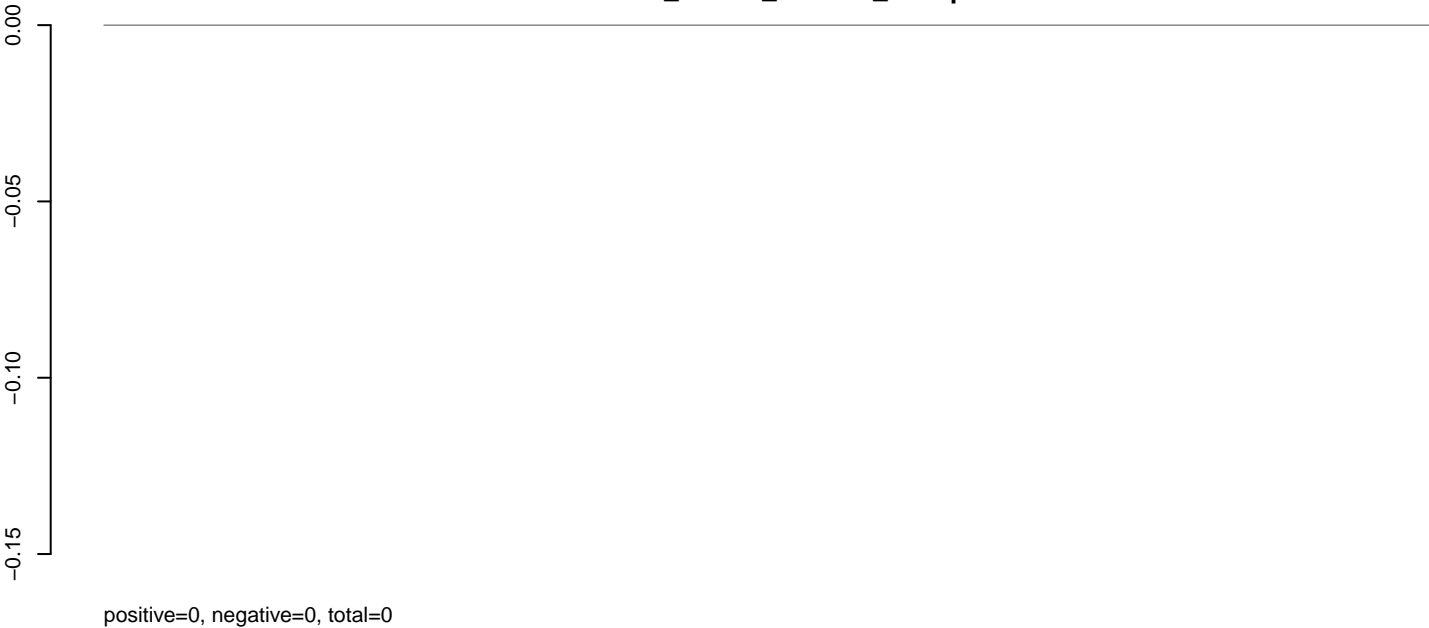
Window size=50, length=1275, TE@EAA13511-Unknown:1-1275

200 400 600 800 1000 1200

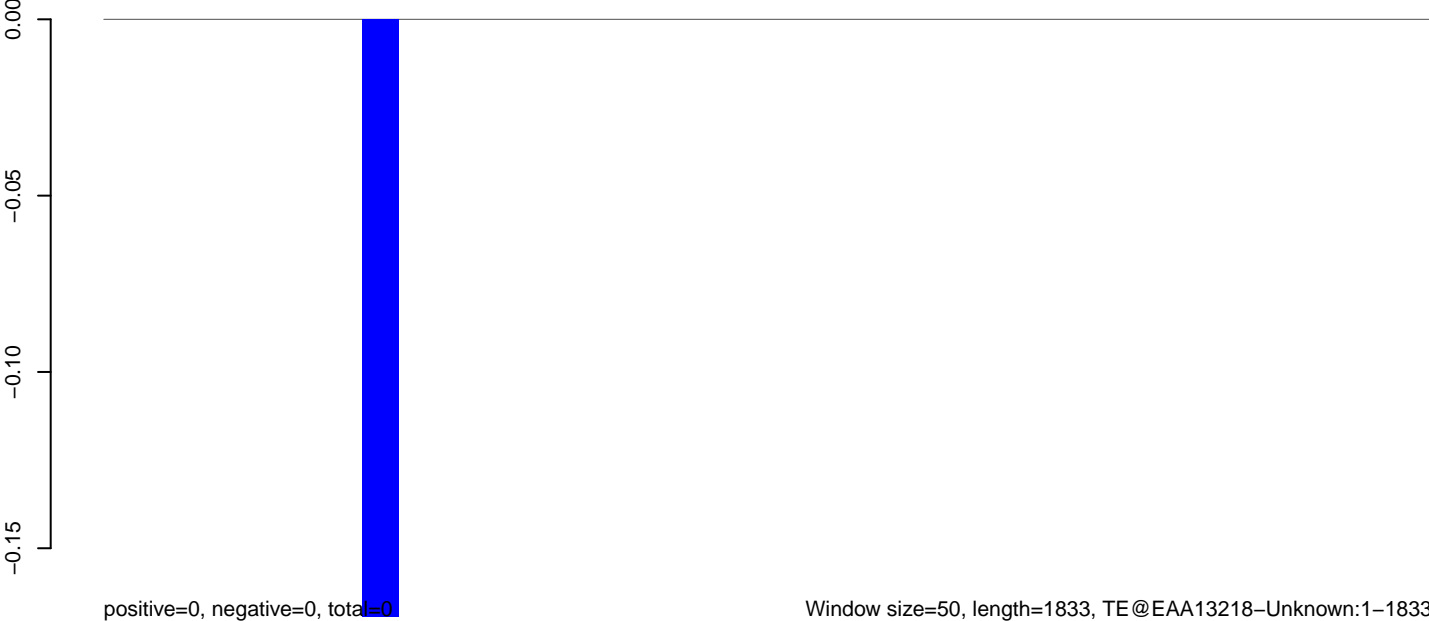
AnGam_Mos55_cells.18_23.rep



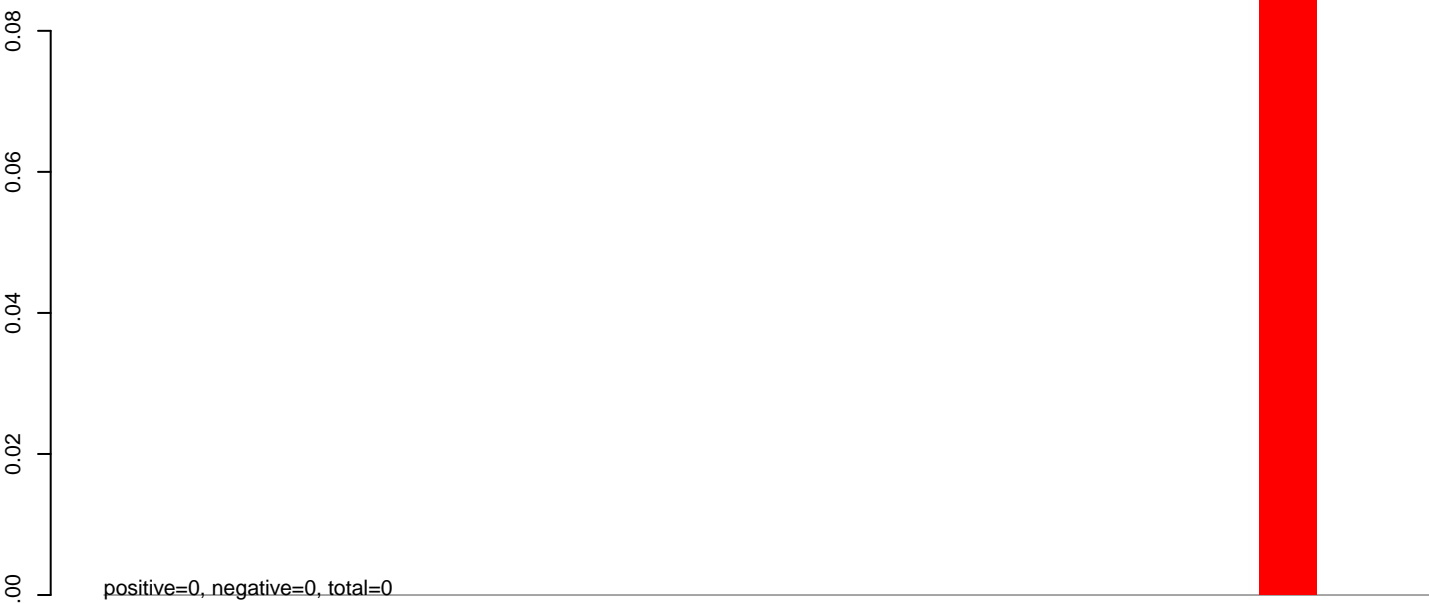
AnGam_Mos55_cells.24_35.rep



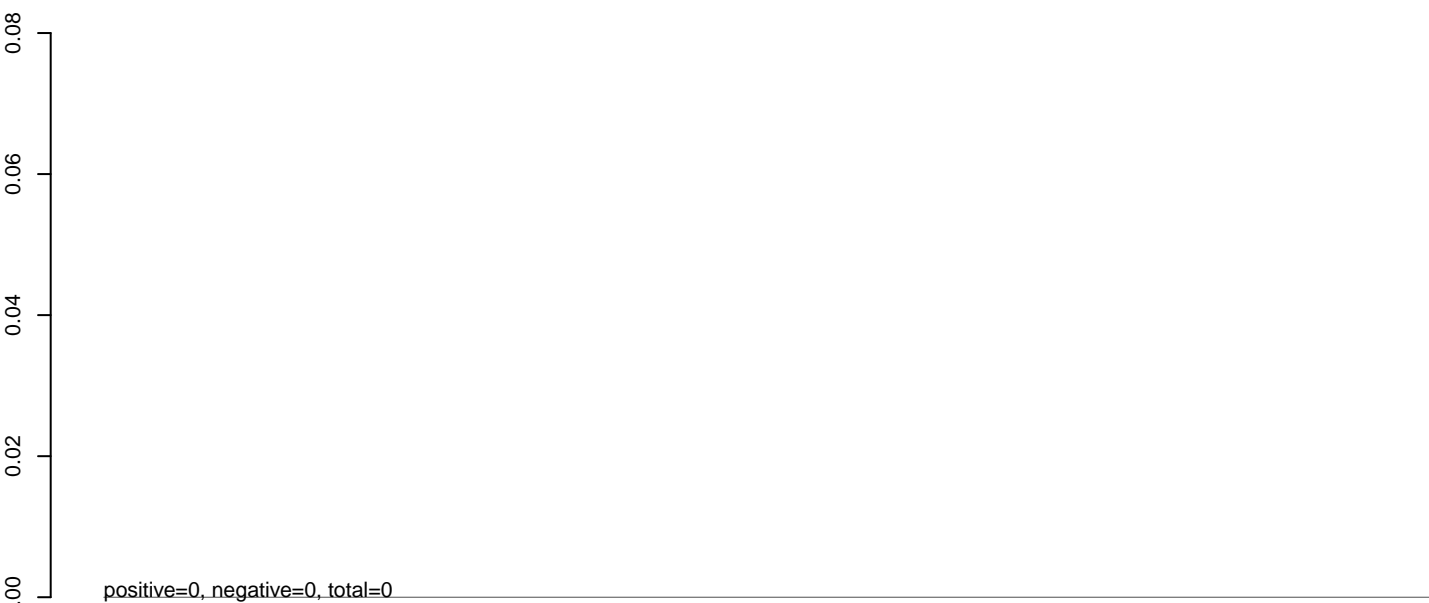
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=567, TE@EAA02779-Unknown:1-567

100 200 300 400 500 600

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=567, TE@EAA02405-Unknown:1-567

100 200 300 400 500 600

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

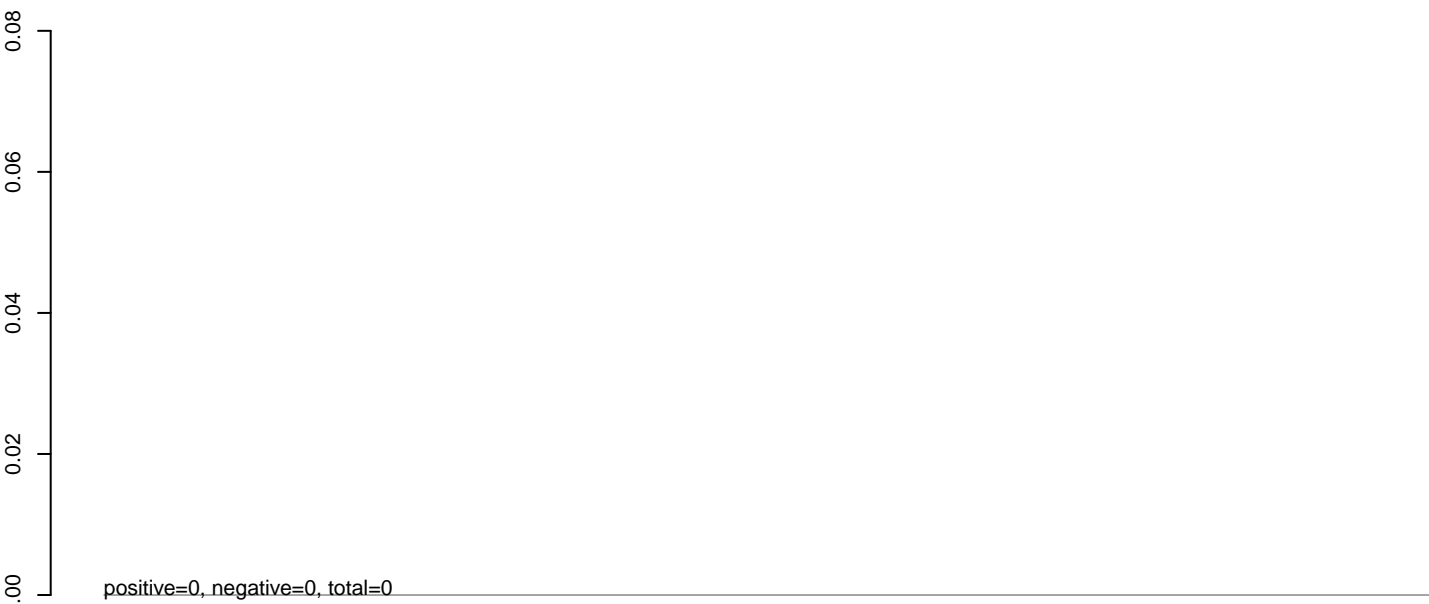
0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

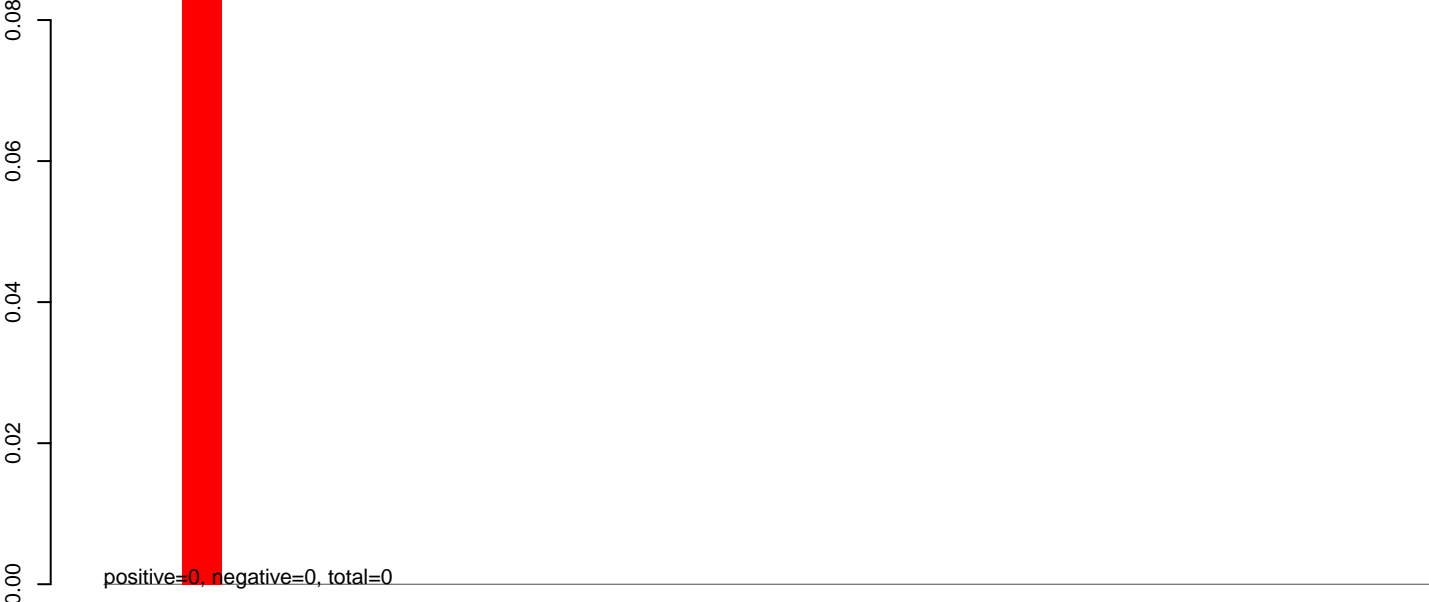
Window size=50, length=3448, TE@CR1-10_AG:1-3448

0 500 1000 1500 2000 2500 3000 3500

AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep

0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

Window size=50, length=402, TE@CAI96717-Unknown:1-402

100 200 300 400

AnGam_Mos55_cells.18_23.rep

0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

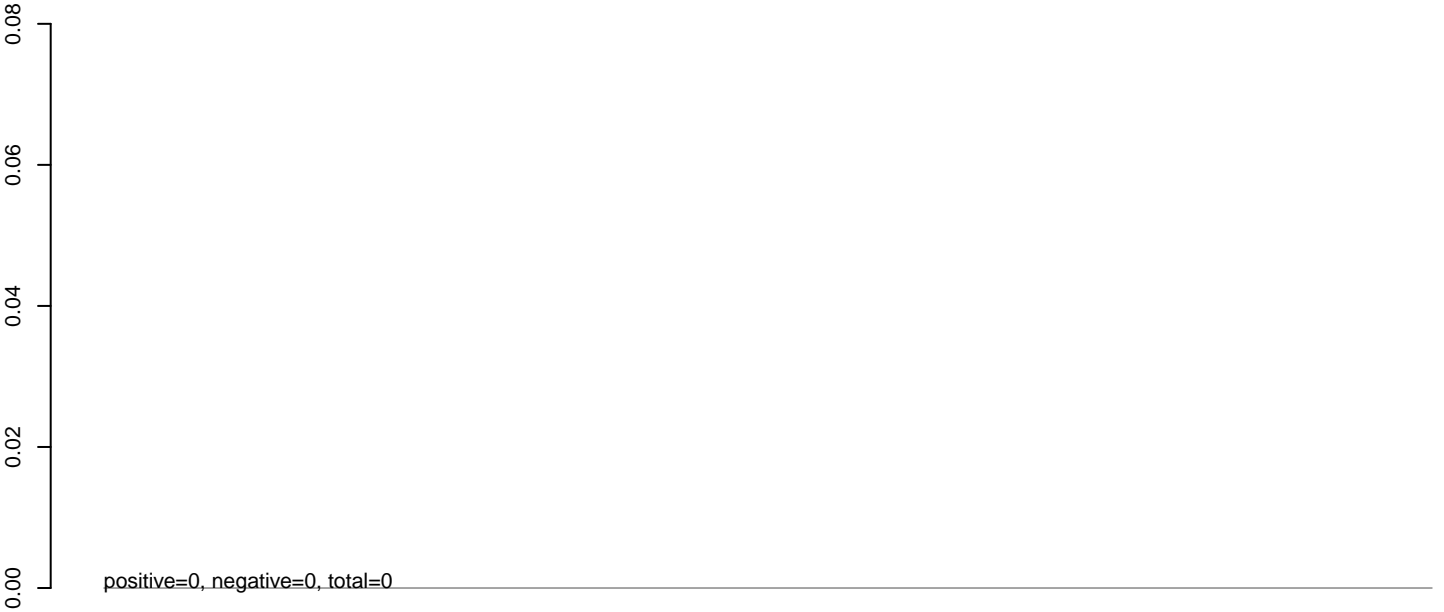
0.00
-0.05
-0.10
-0.15

positive=0, negative=0, total=0

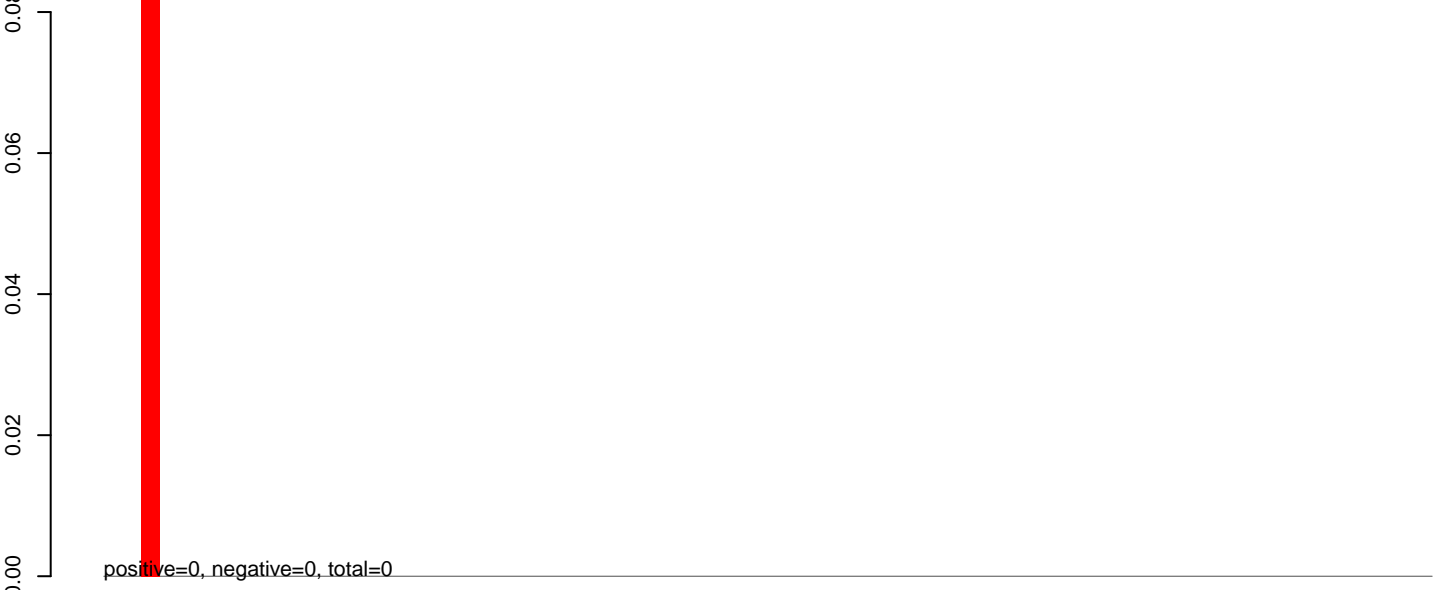
Window size=50, length=6176, TE@BEL14-LTR-I_AG:1-6176

0 1000 2000 3000 4000 5000 6000

AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep

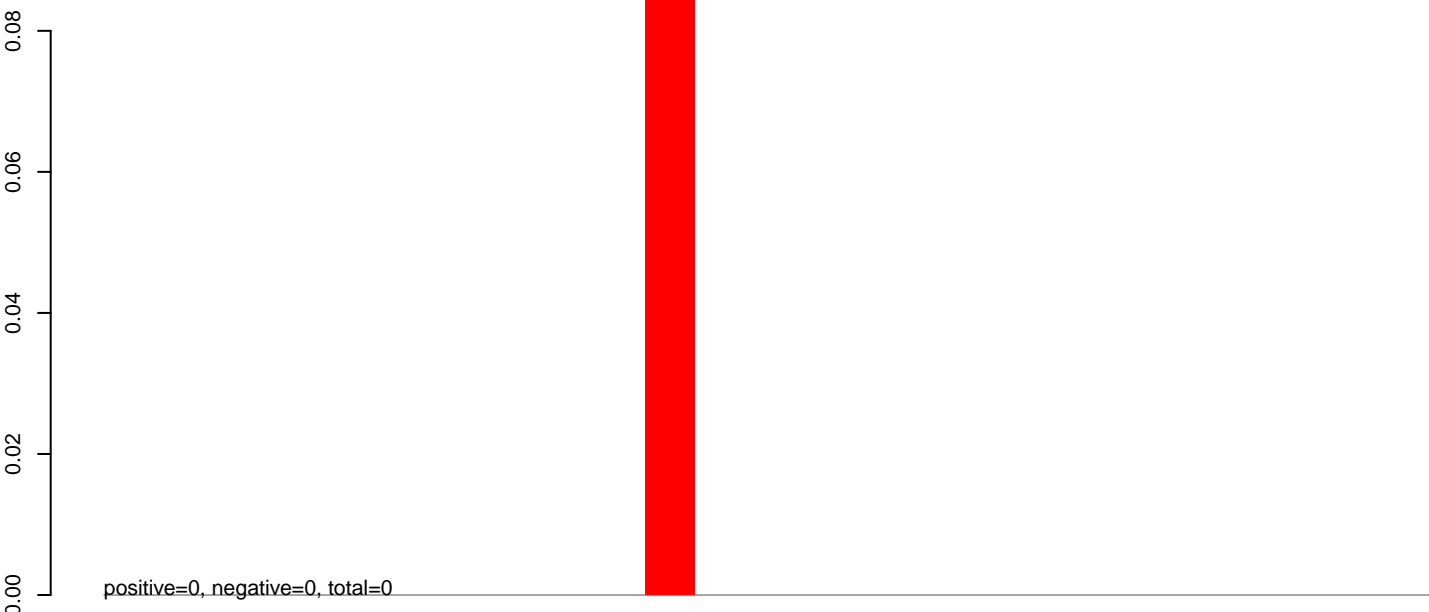


AnGam_Mos55_cells.rep

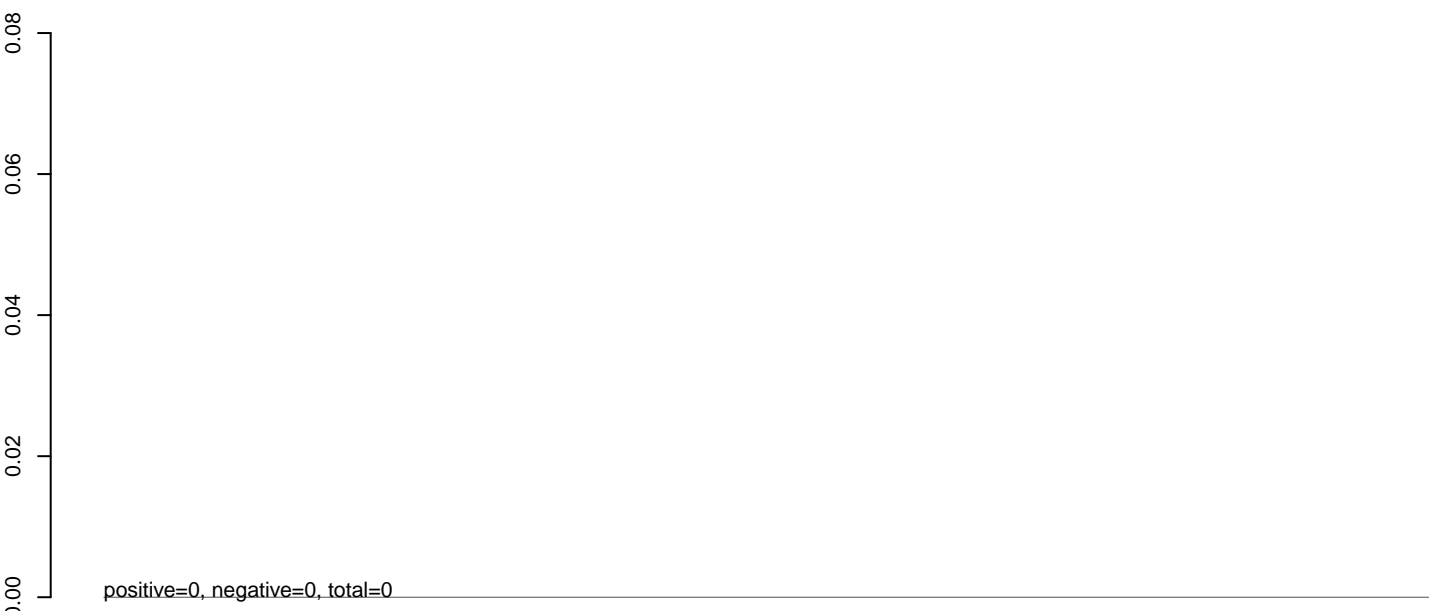


0 500 1000 1500 2000 2500 3000 3500

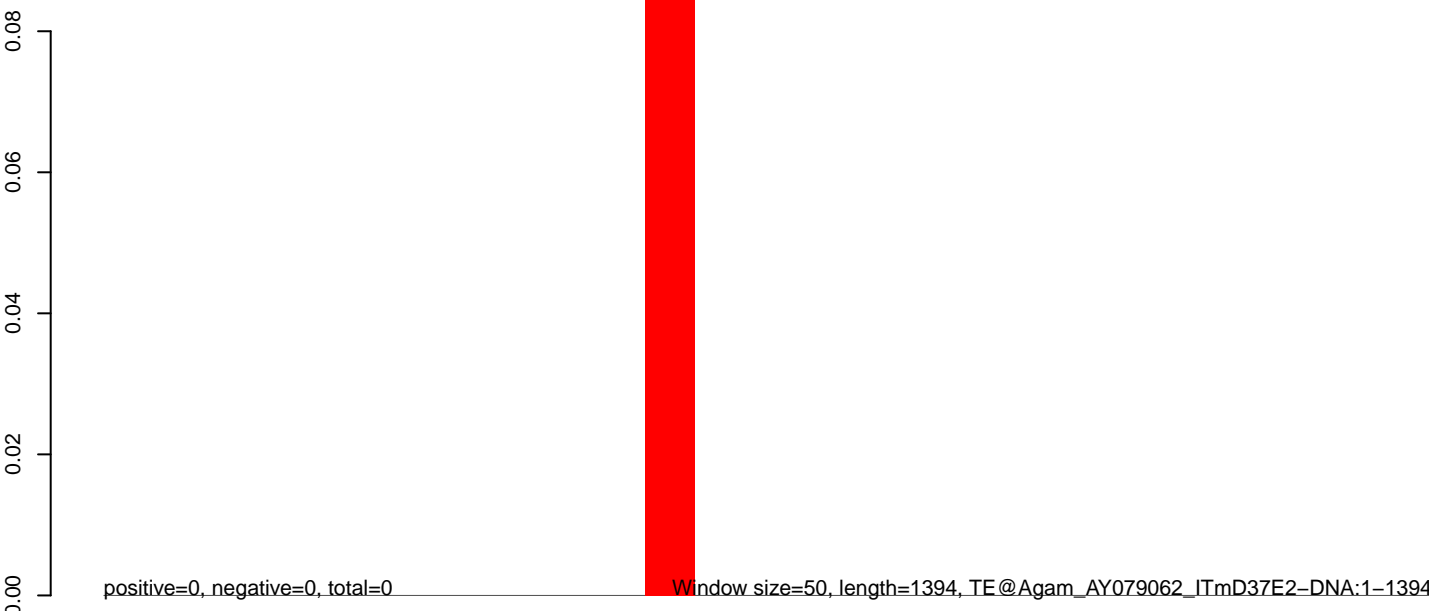
AnGam_Mos55_cells.18_23.rep



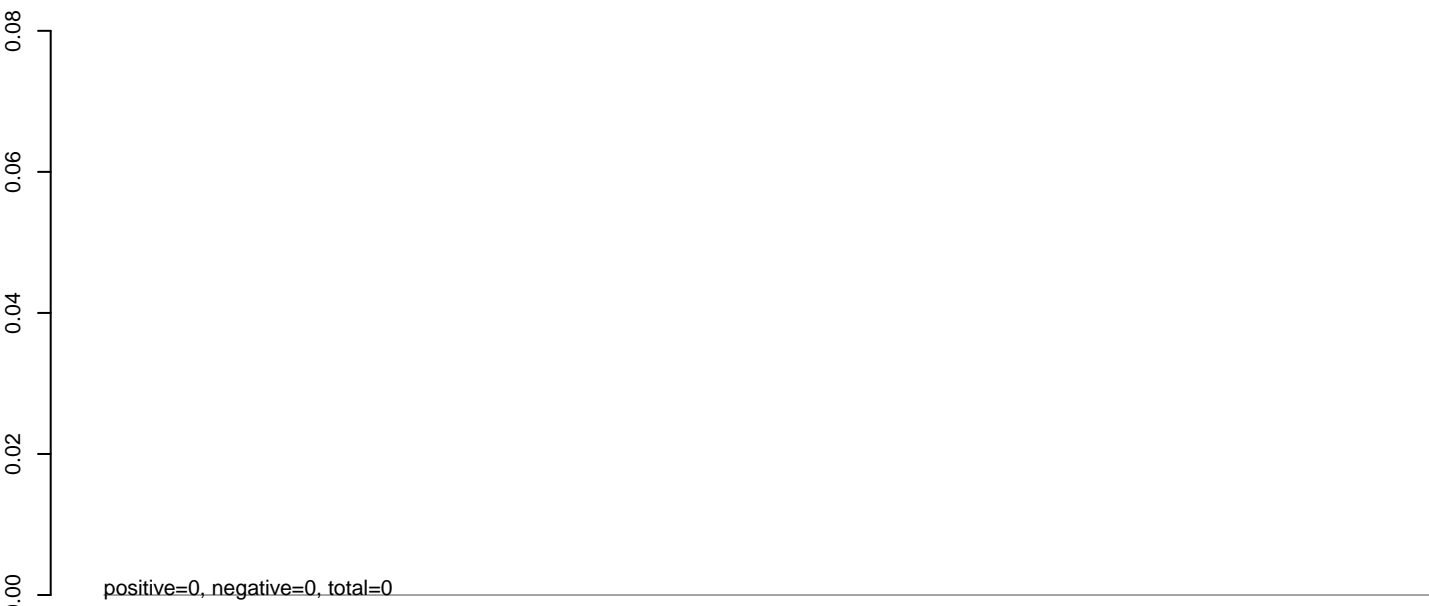
AnGam_Mos55_cells.24_35.rep



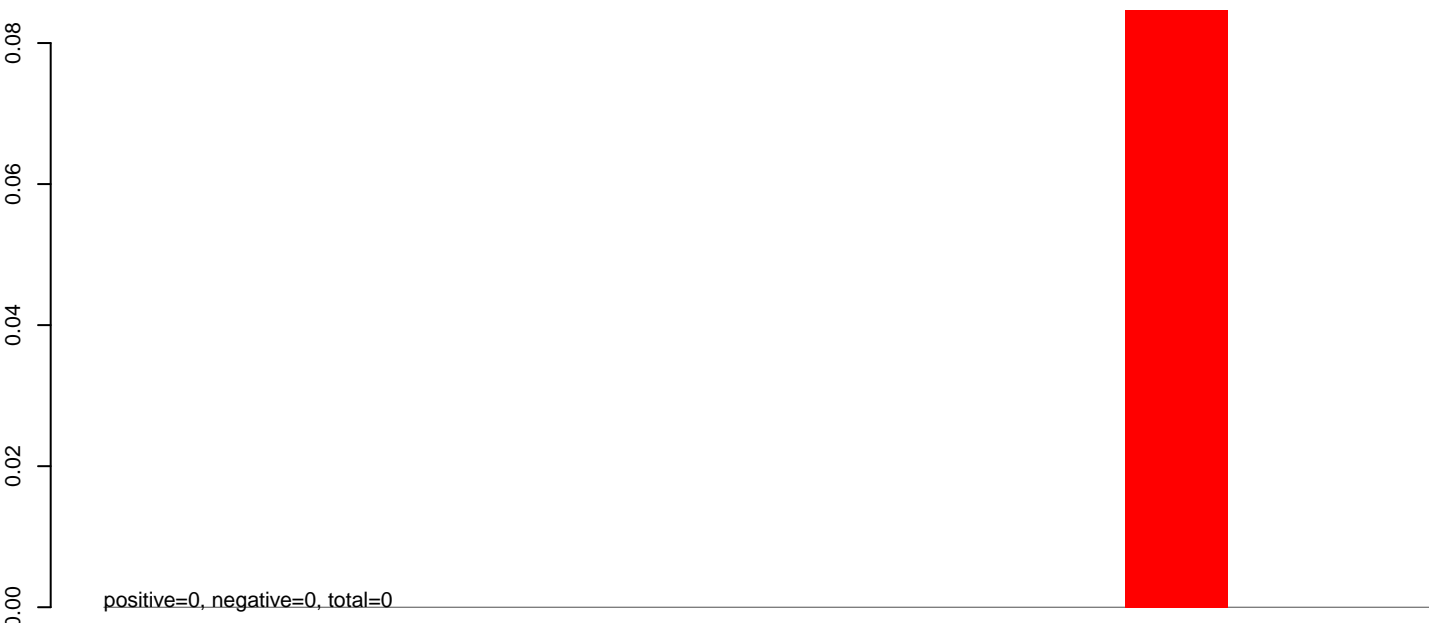
AnGam_Mos55_cells.rep



AnGam_Mos55_cells.18_23.rep



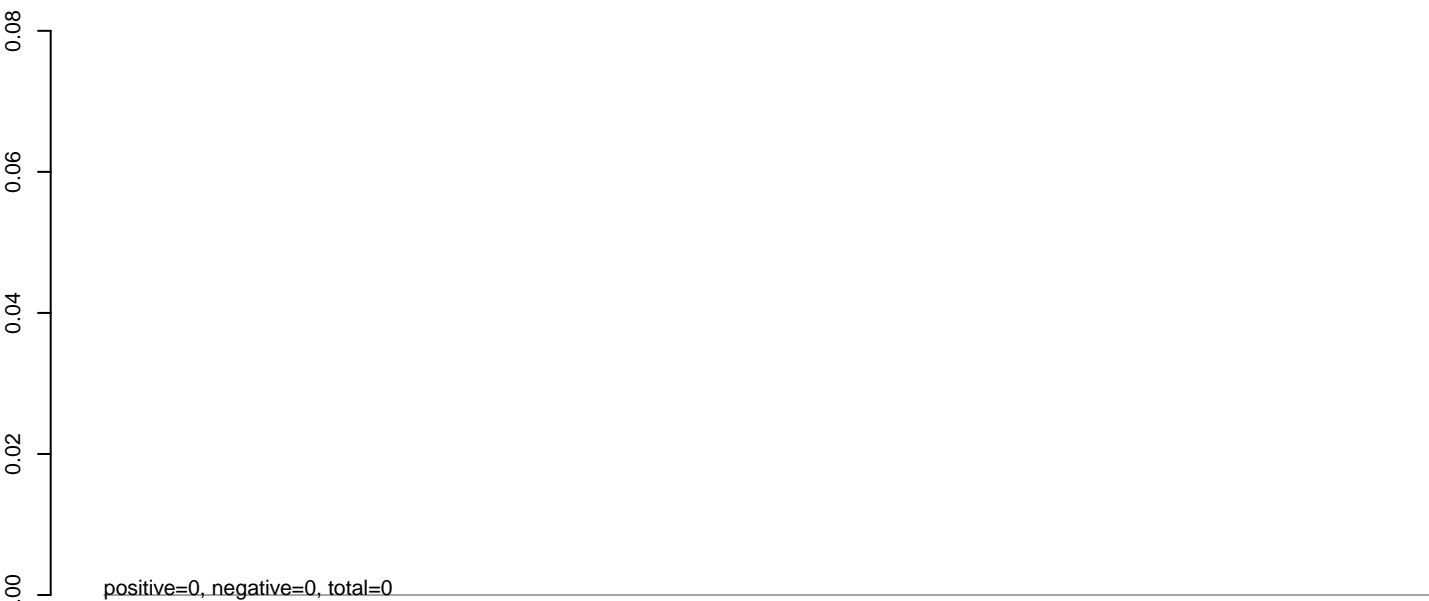
AnGam_Mos55_cells.24_35.rep



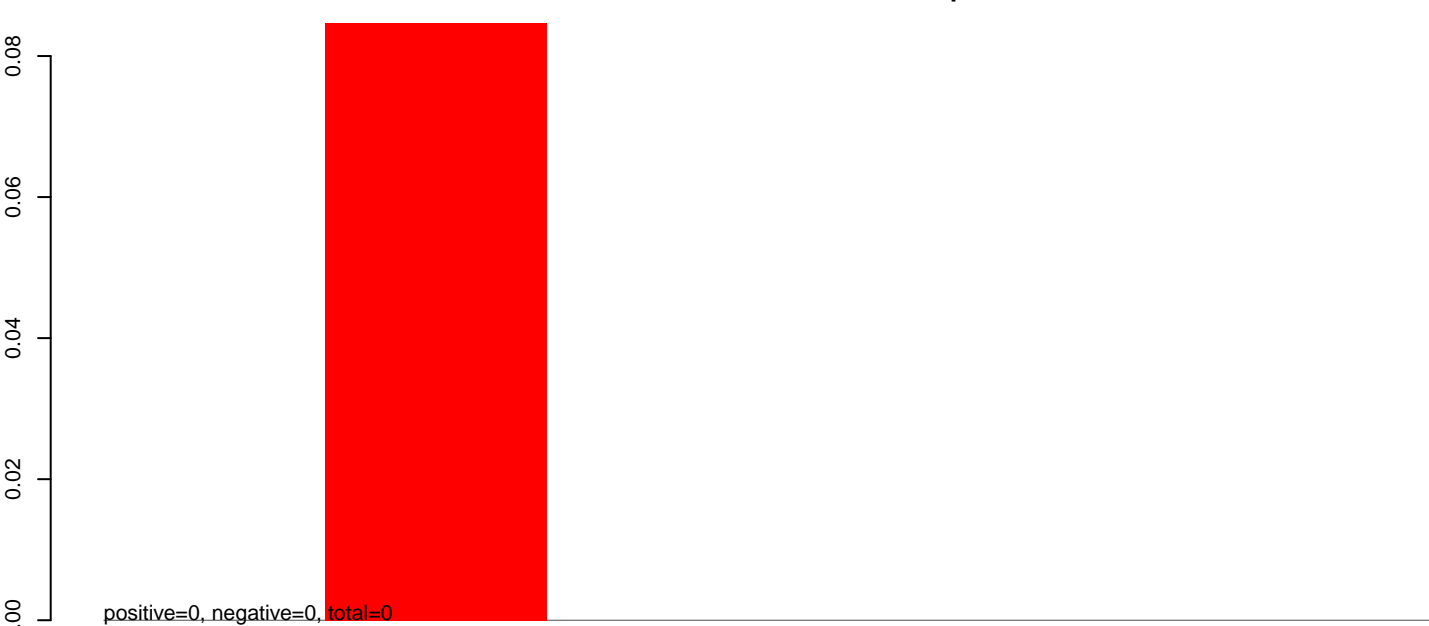
AnGam_Mos55_cells.rep



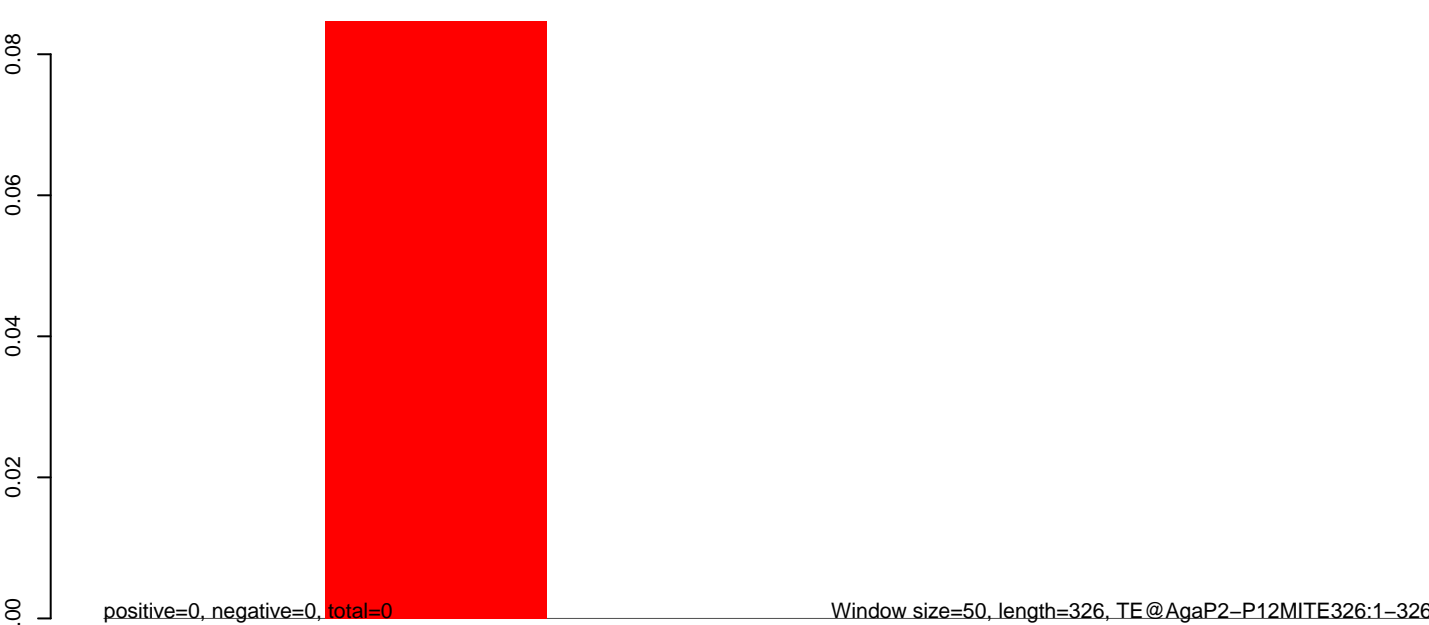
AnGam_Mos55_cells.18_23.rep



AnGam_Mos55_cells.24_35.rep



AnGam_Mos55_cells.rep



Window size=50, length=326, TE@AgaP2-P12MITE326:1-326

AnGam_Mos55_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AnGam_Mos55_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

Window size=50, length=412, TE@AgaP13MITE412:1-412

100

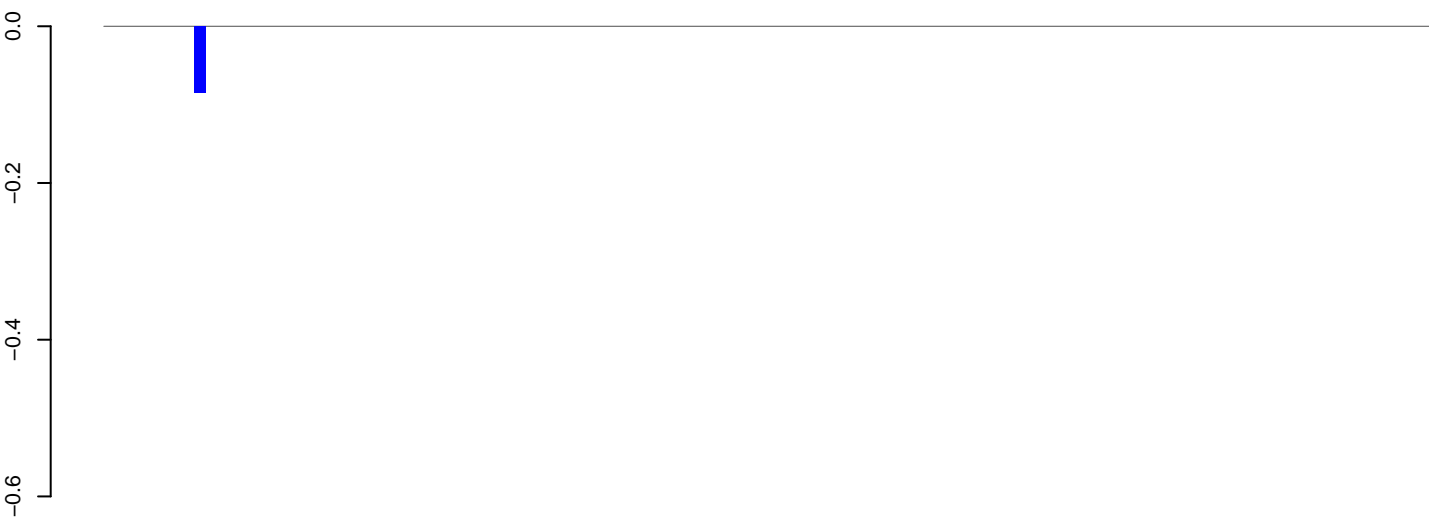
200

300

400

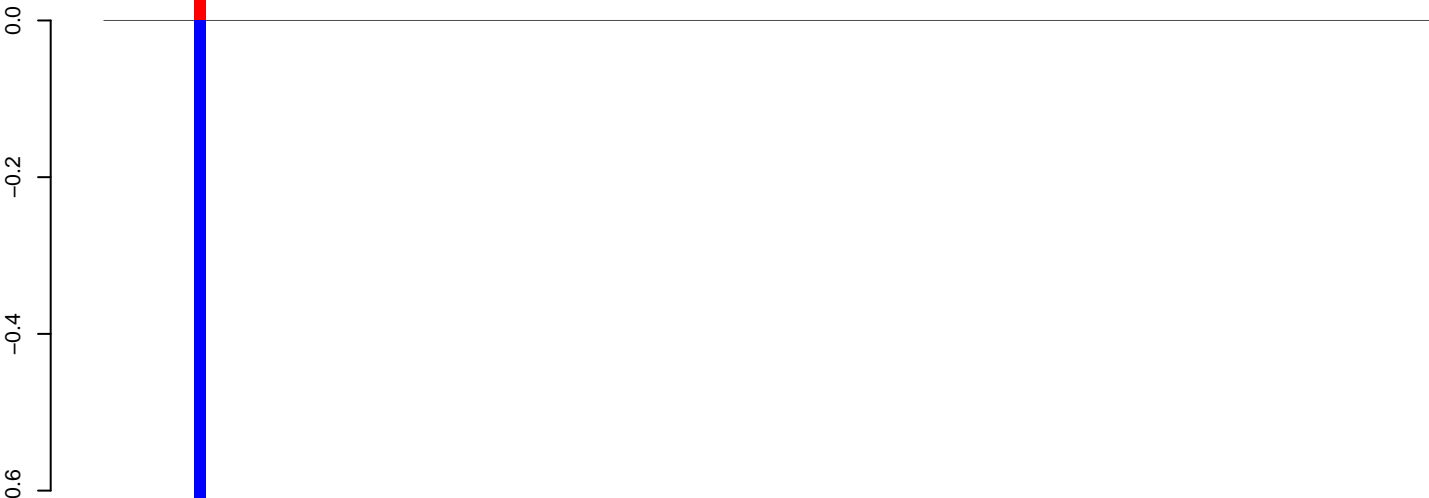


AnGam_Mos55_cells.18_23.rep



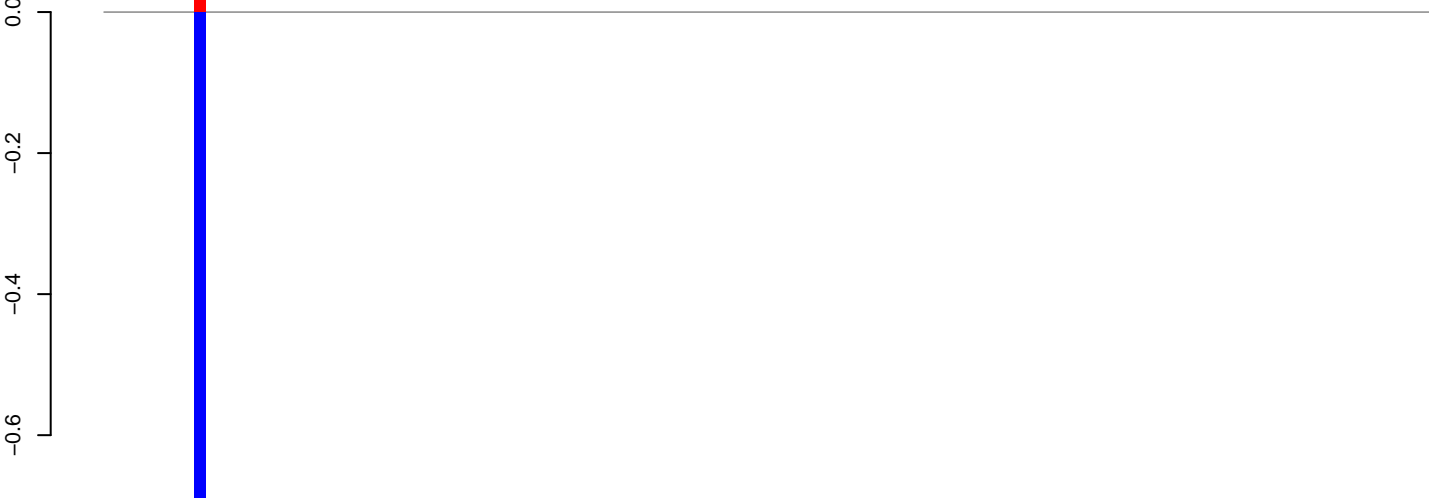
positive=0, negative=0, total=0

AnGam_Mos55_cells.24_35.rep



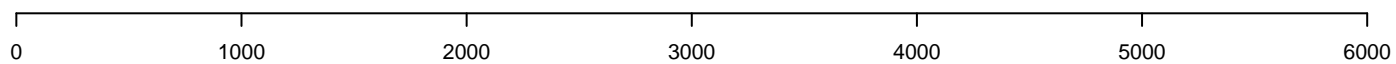
positive=0, negative=1, total=1

AnGam_Mos55_cells.rep

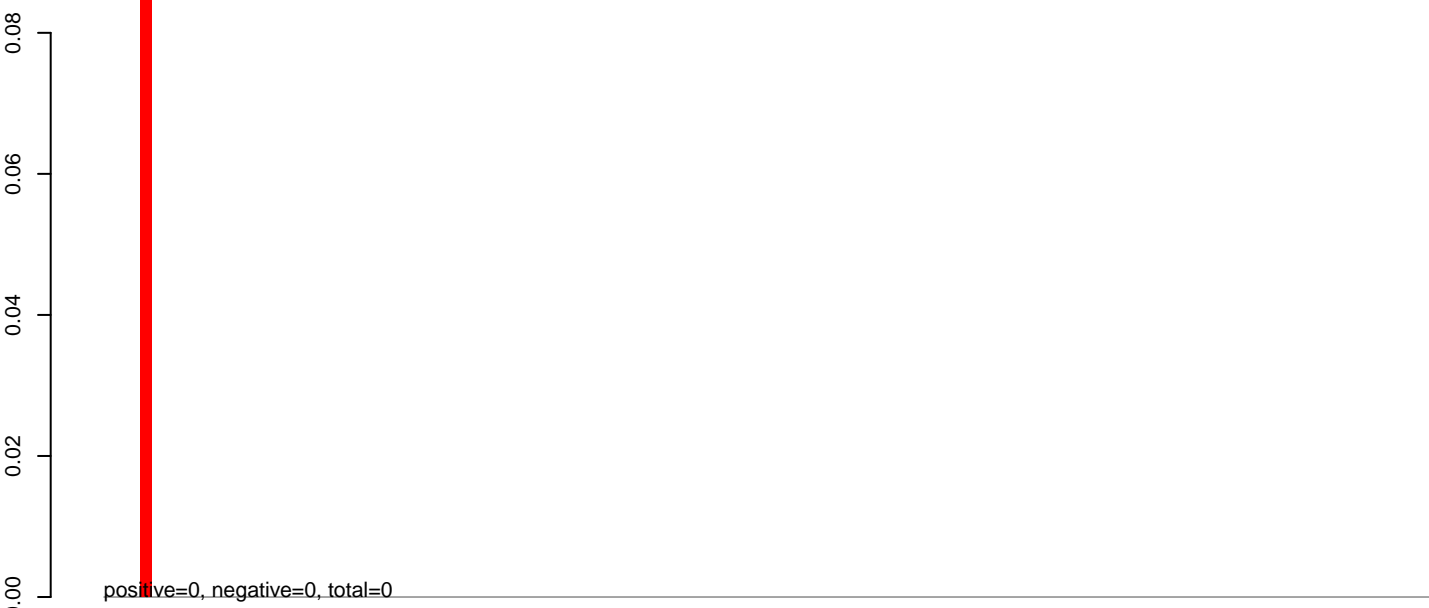


positive=0, negative=1, total=1

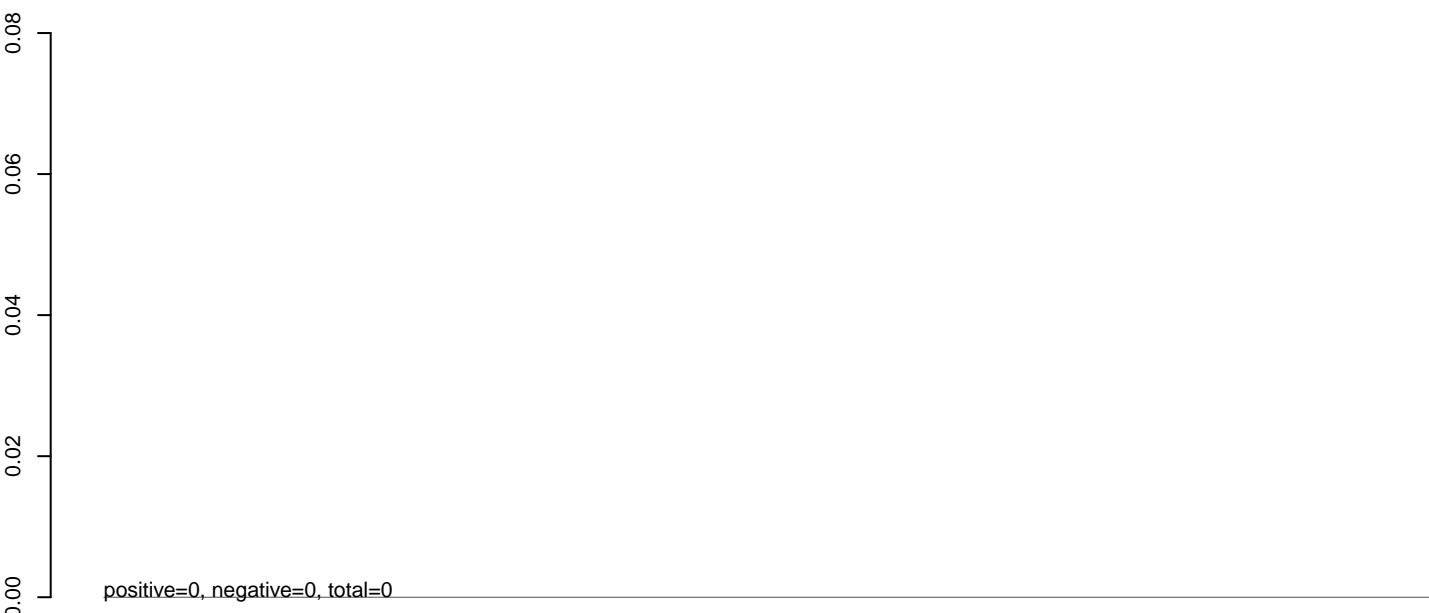
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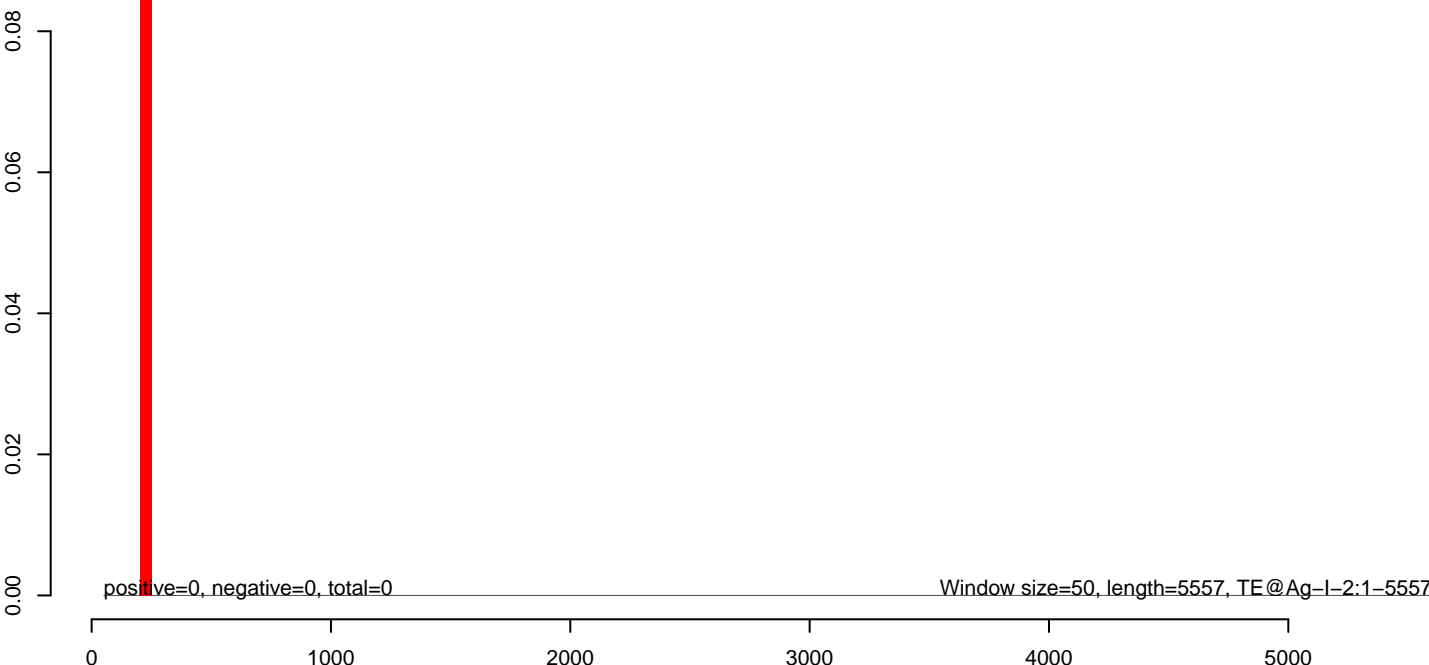
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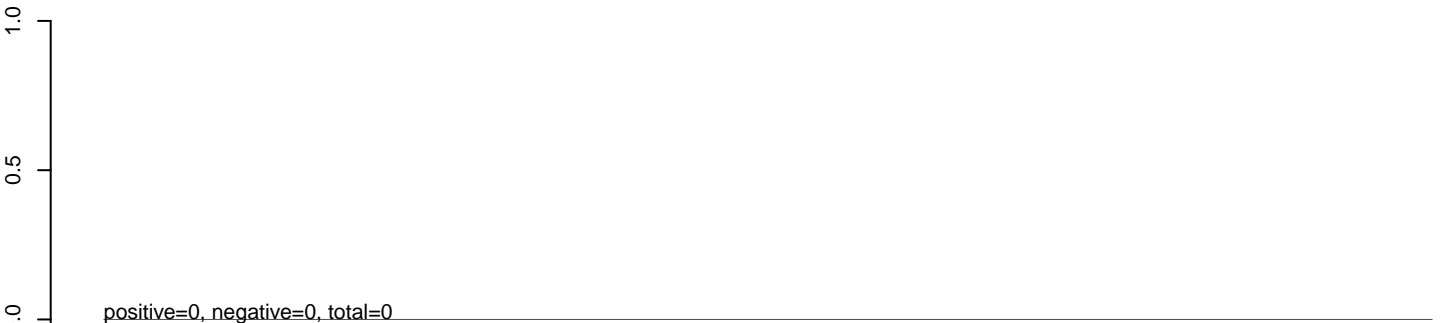
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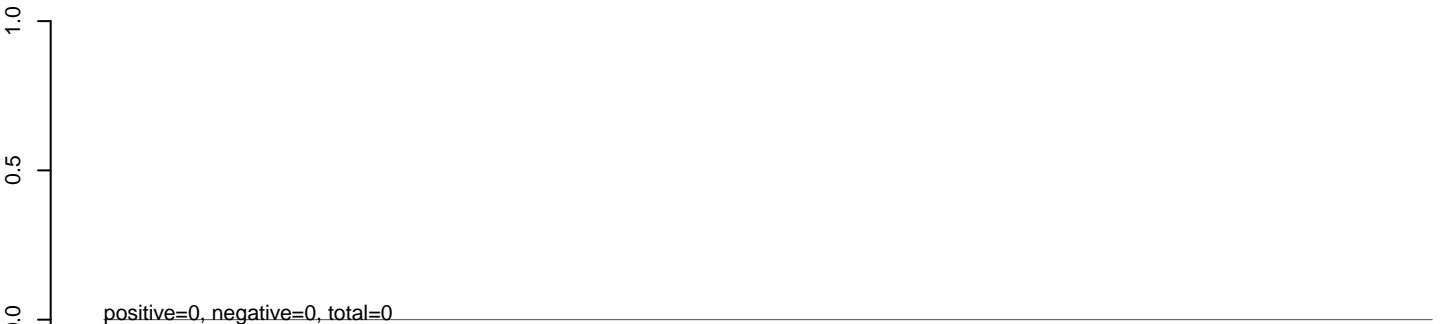
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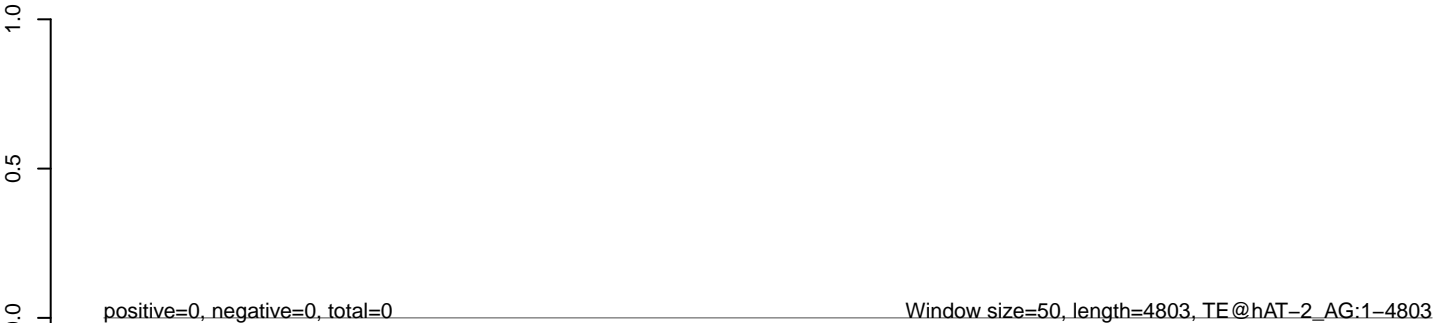
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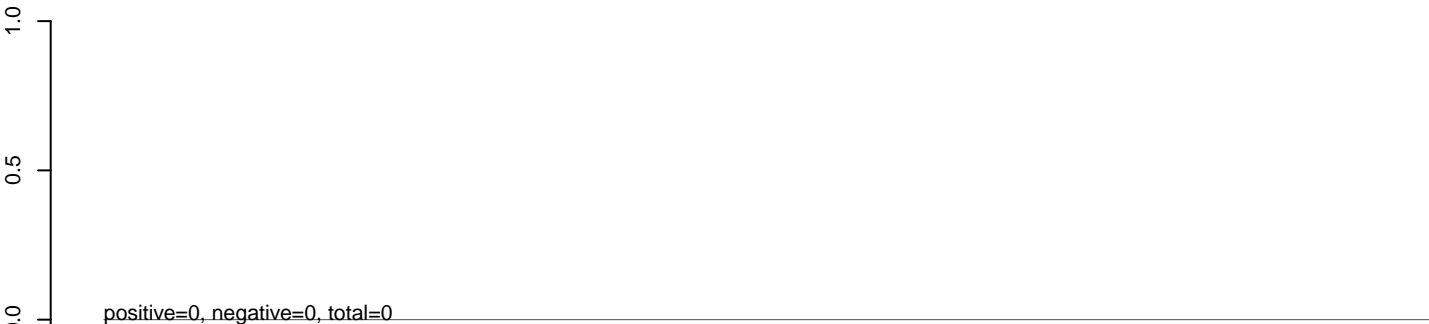
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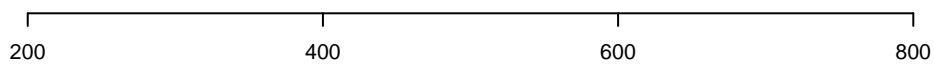
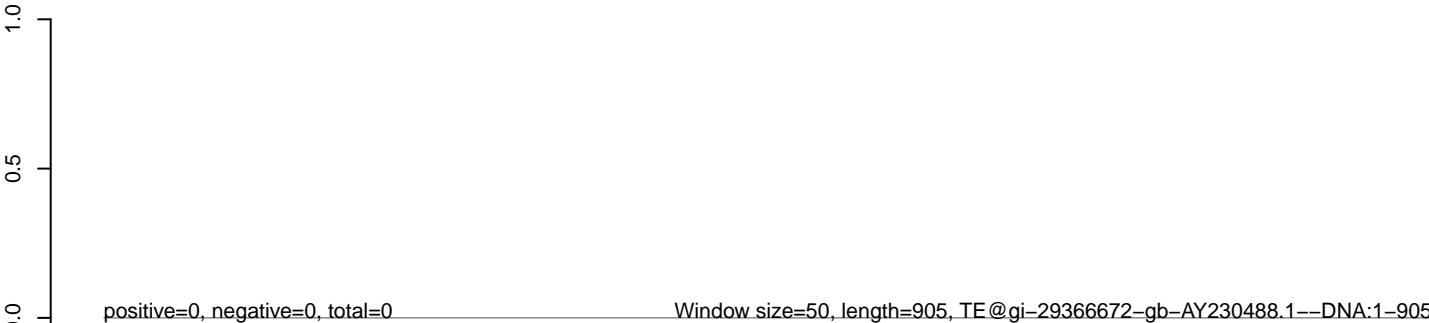
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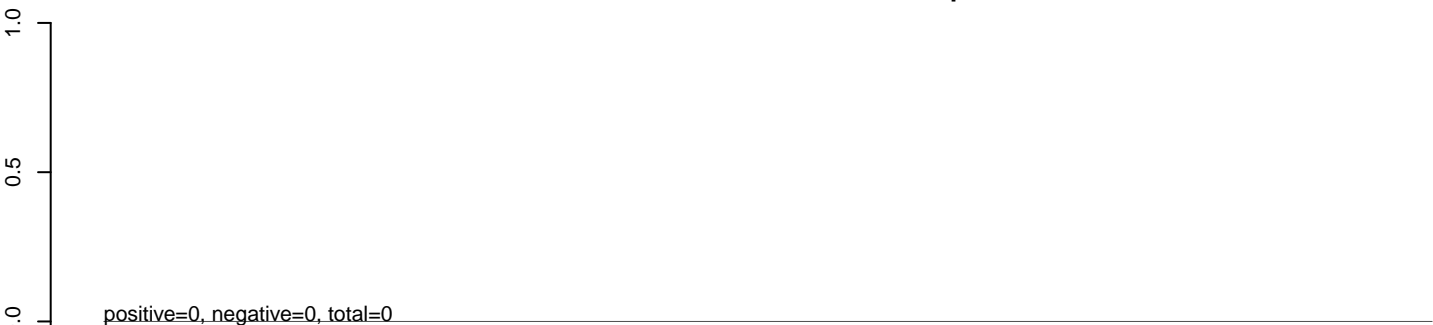
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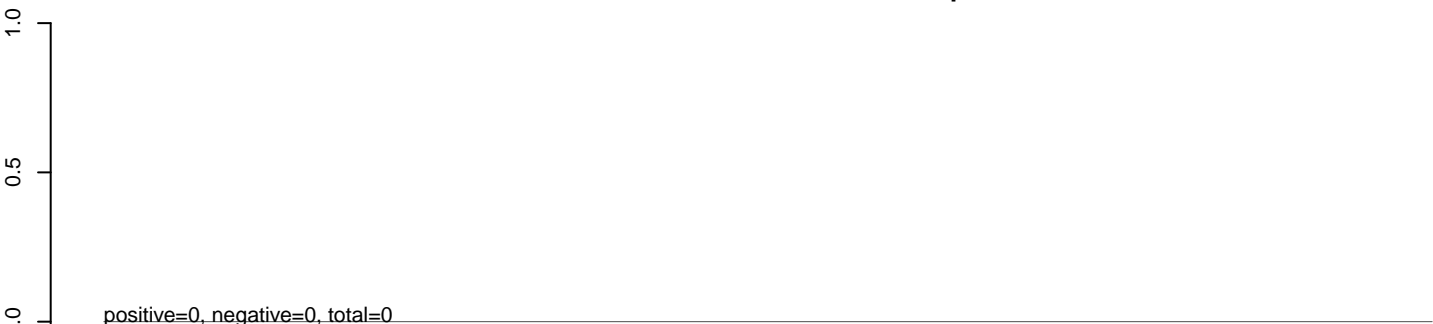
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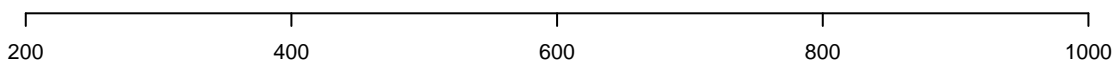
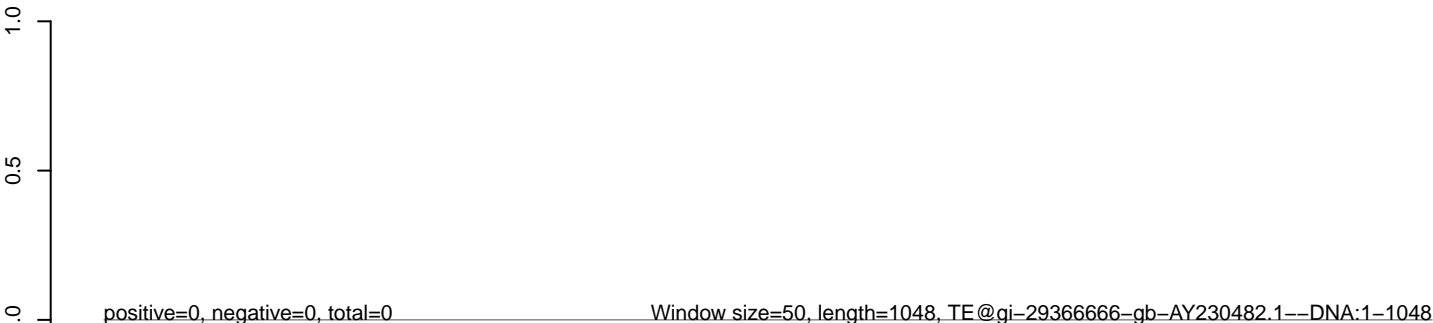
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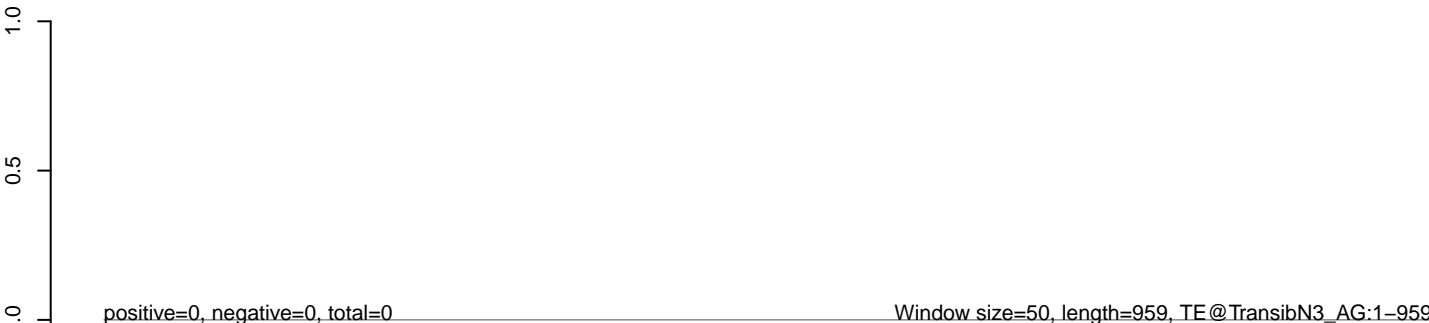
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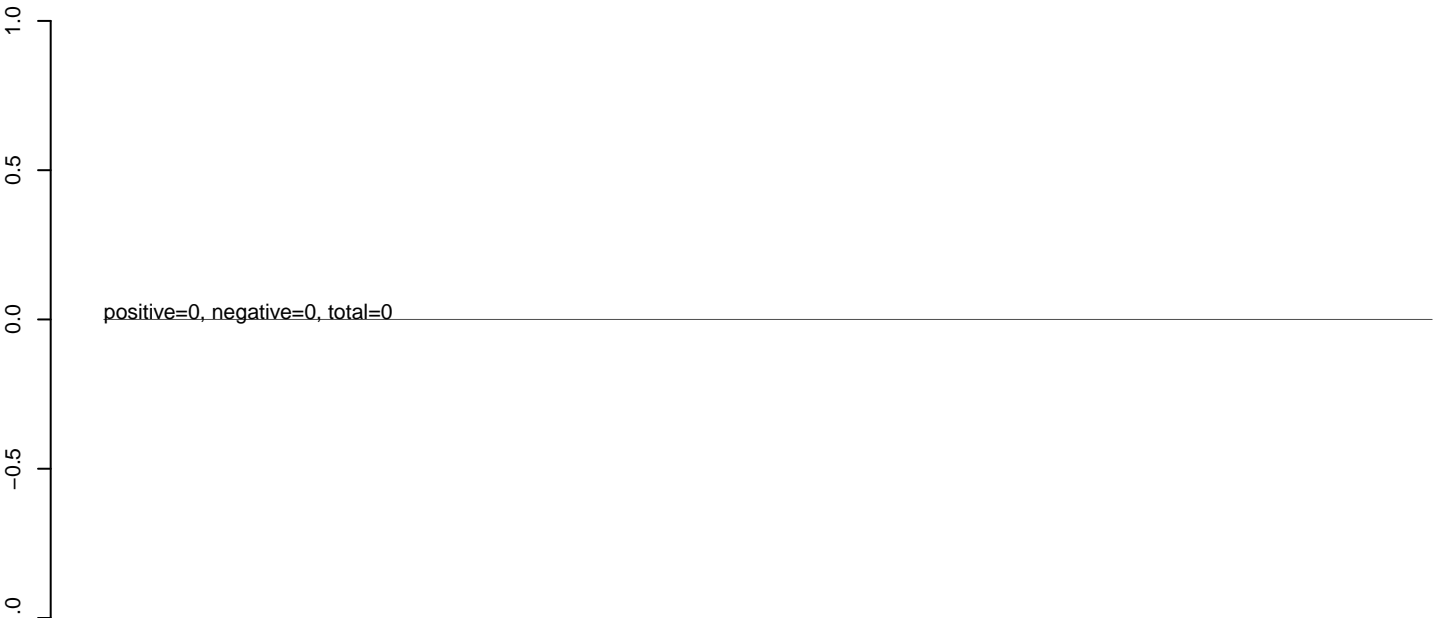
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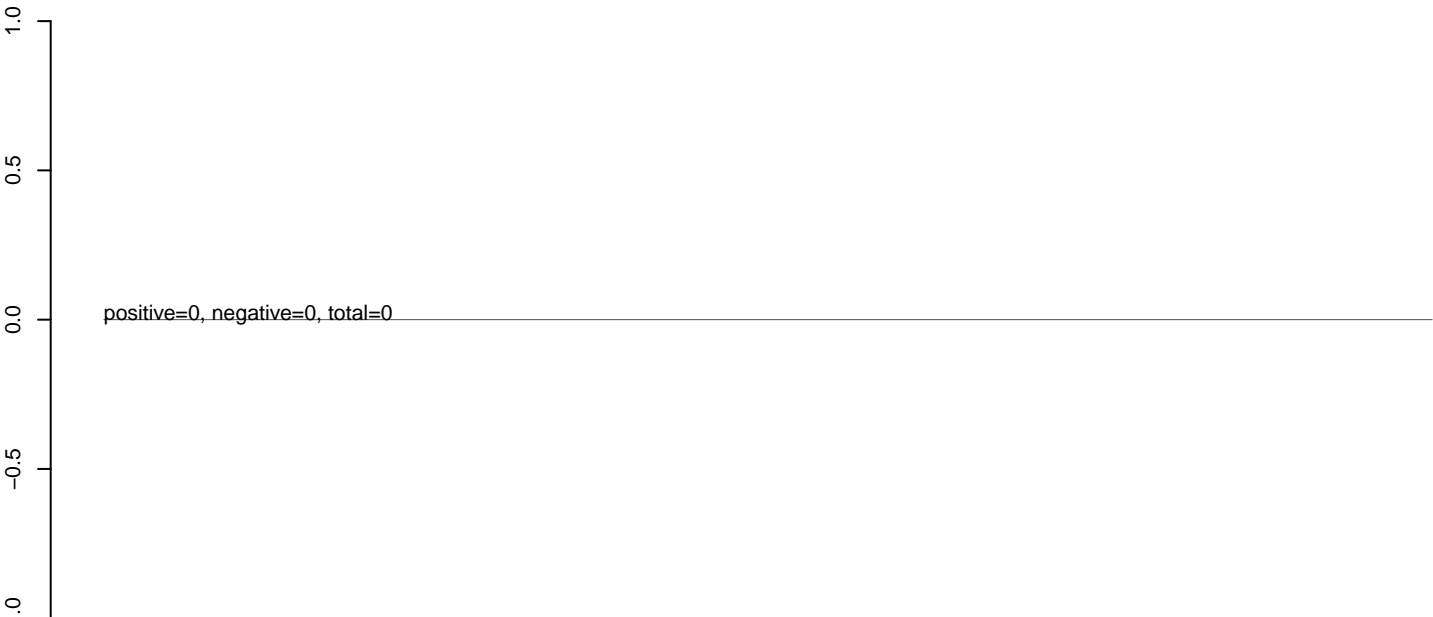
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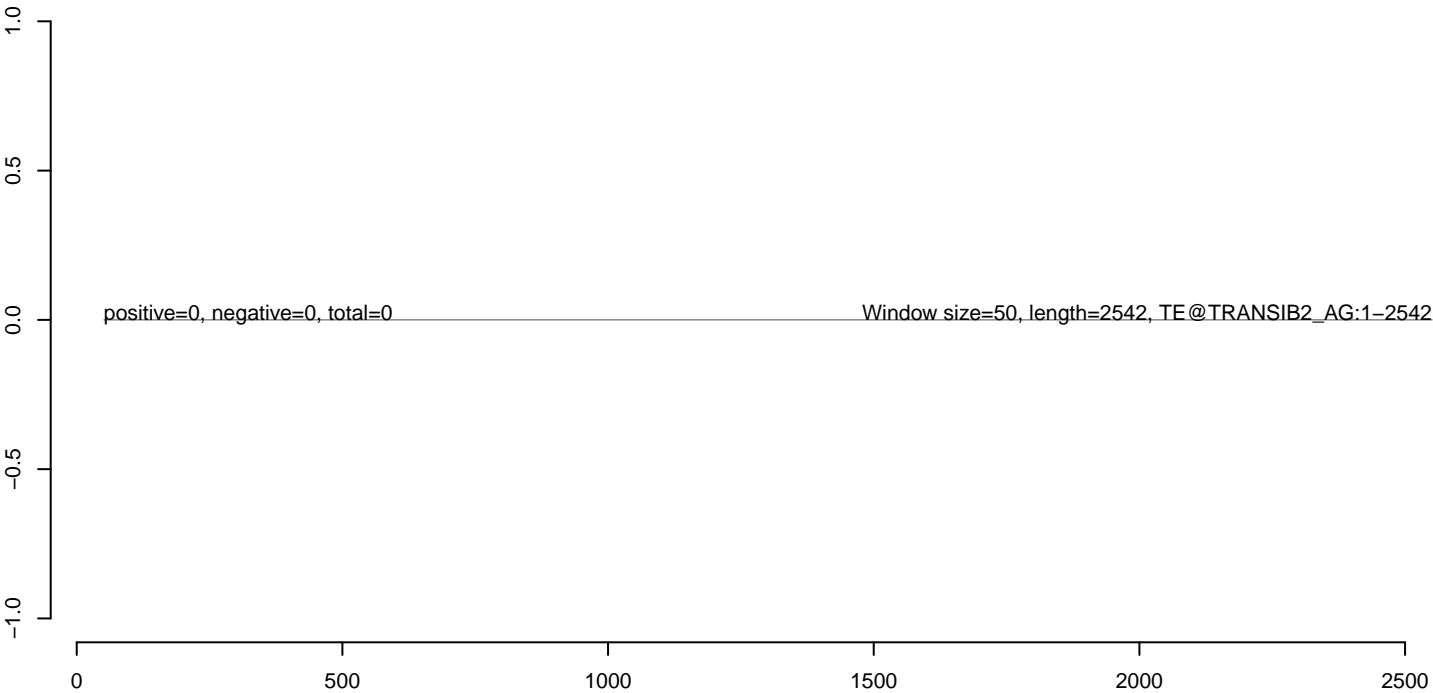
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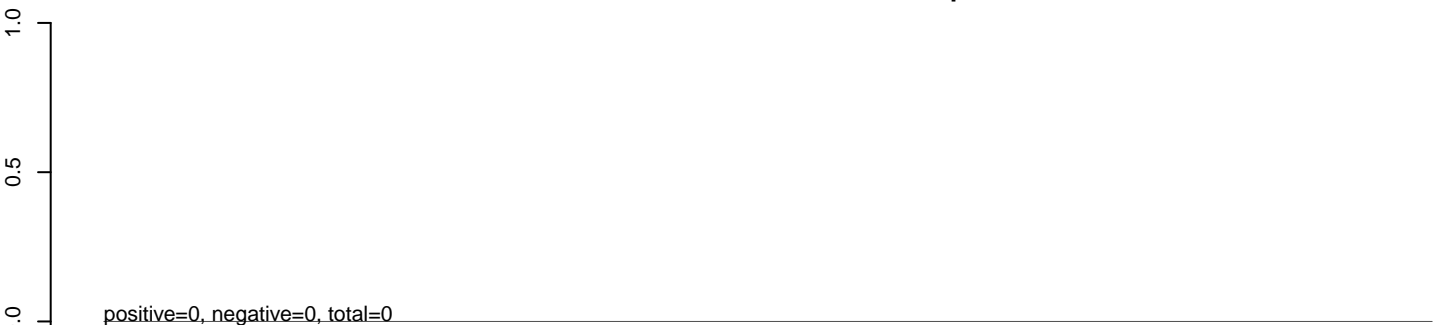
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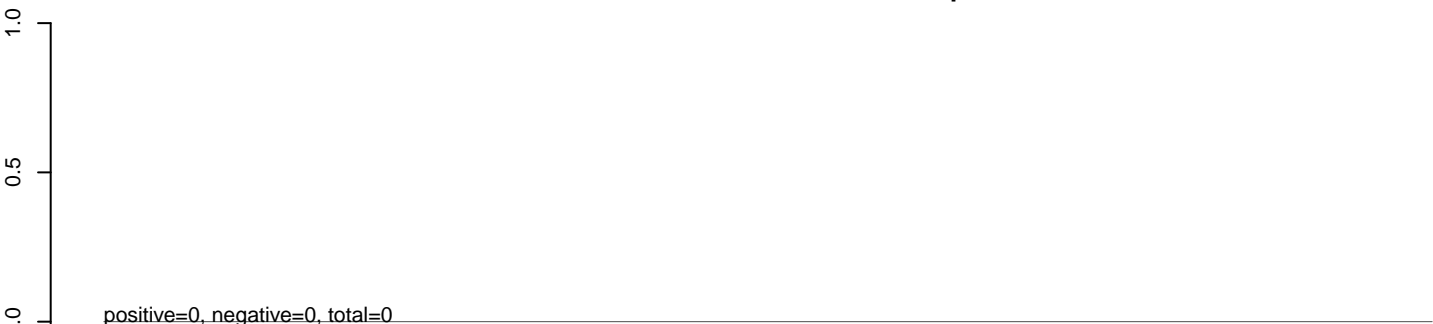
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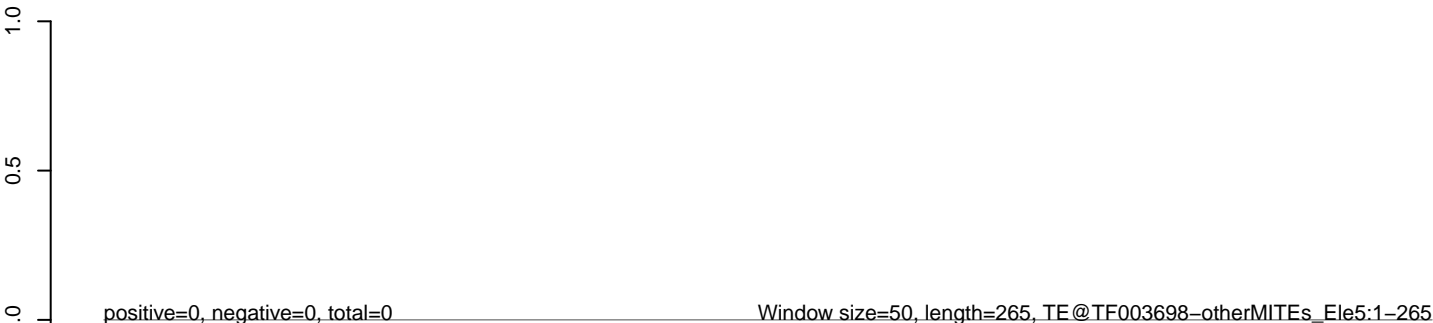
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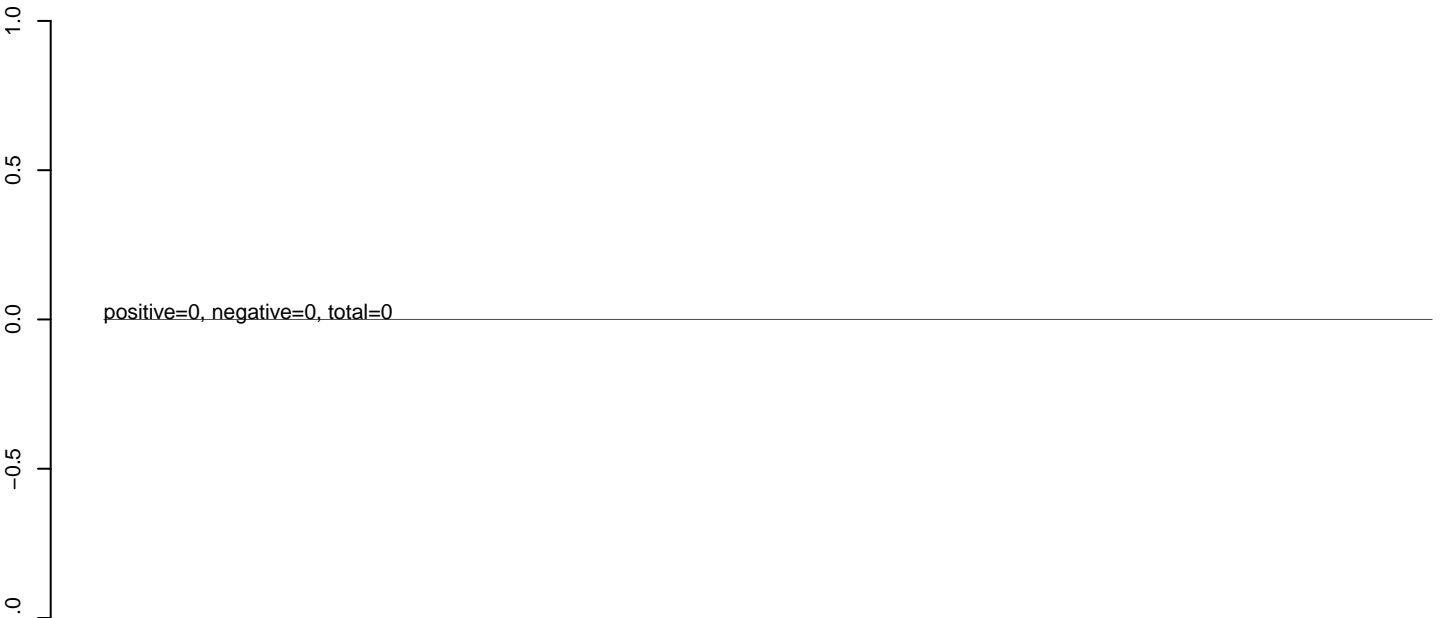
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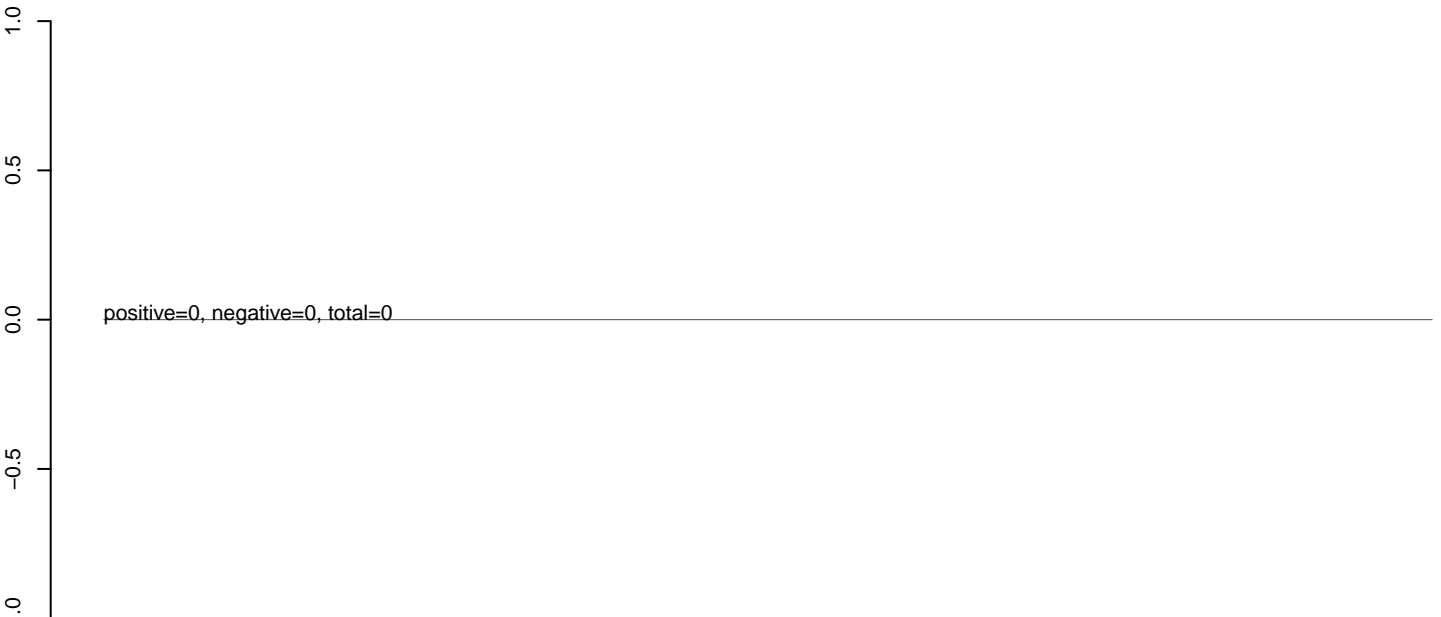
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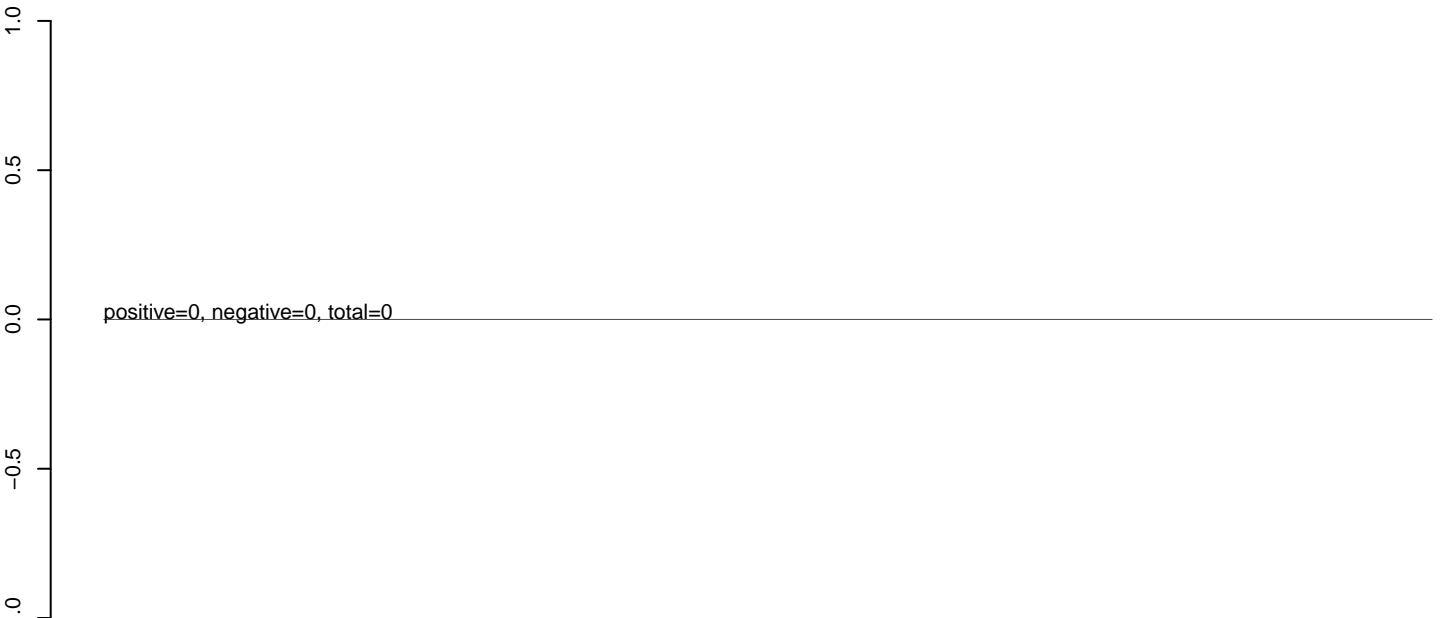
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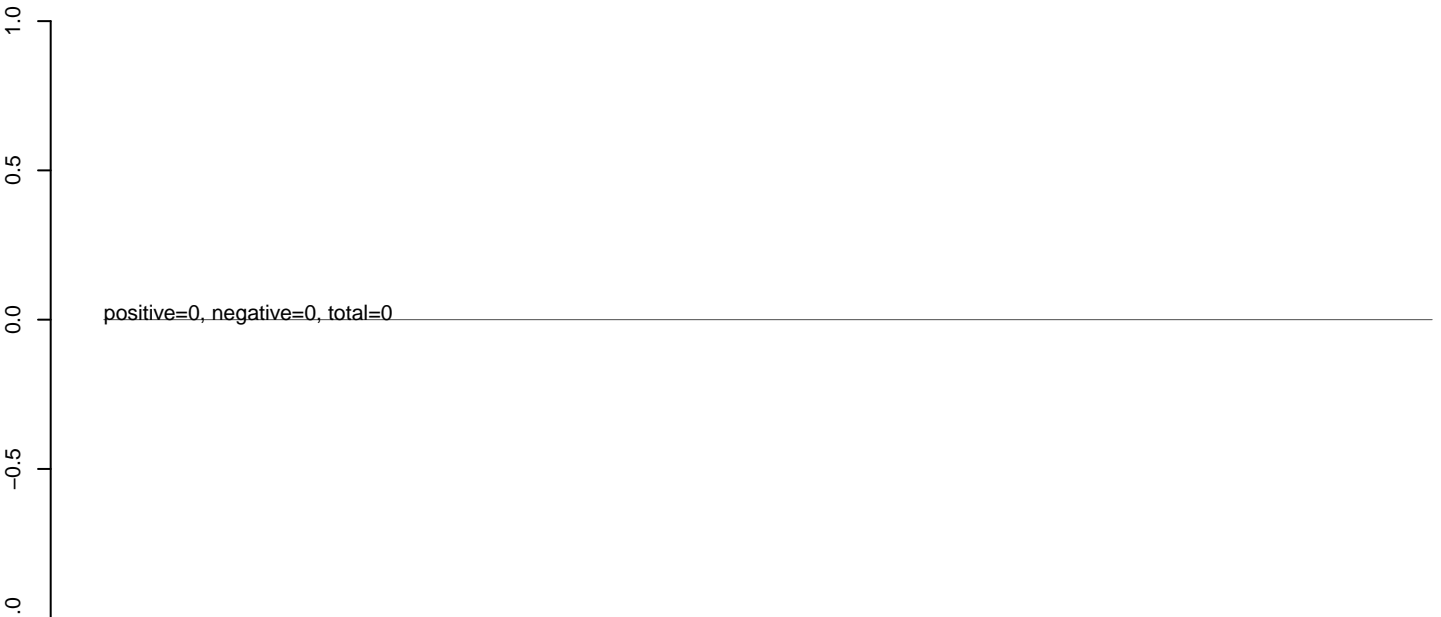
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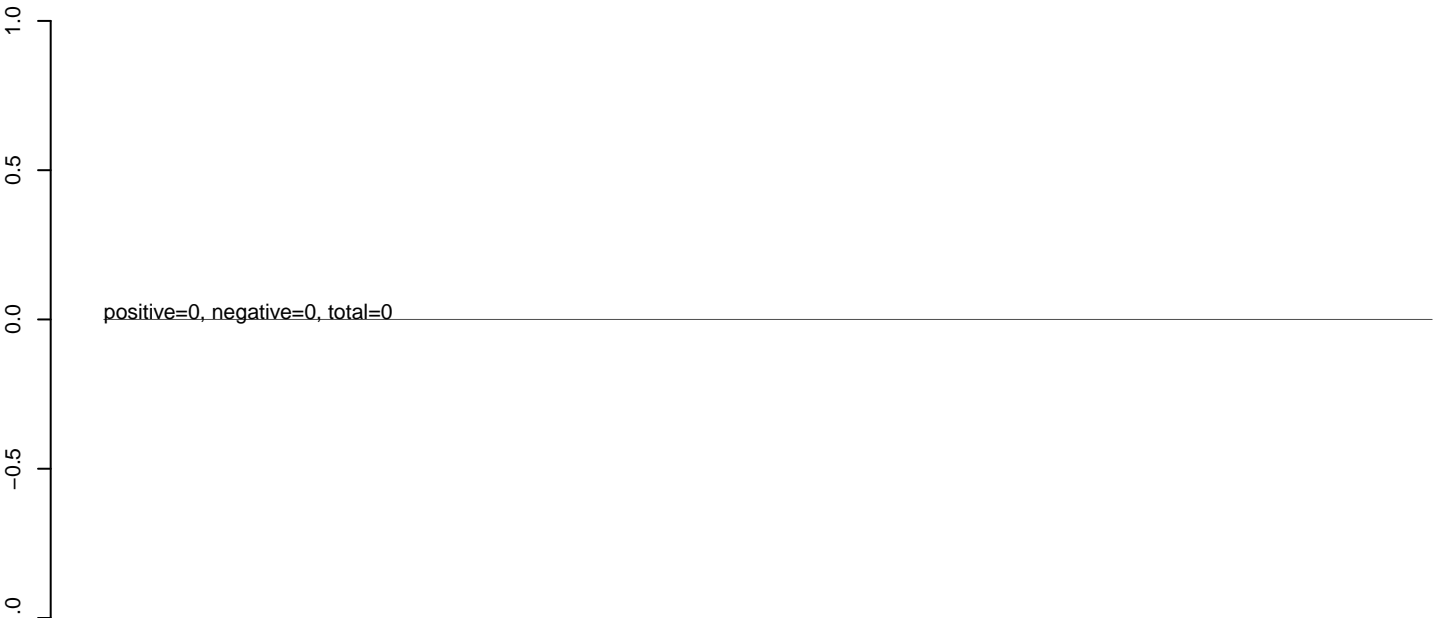
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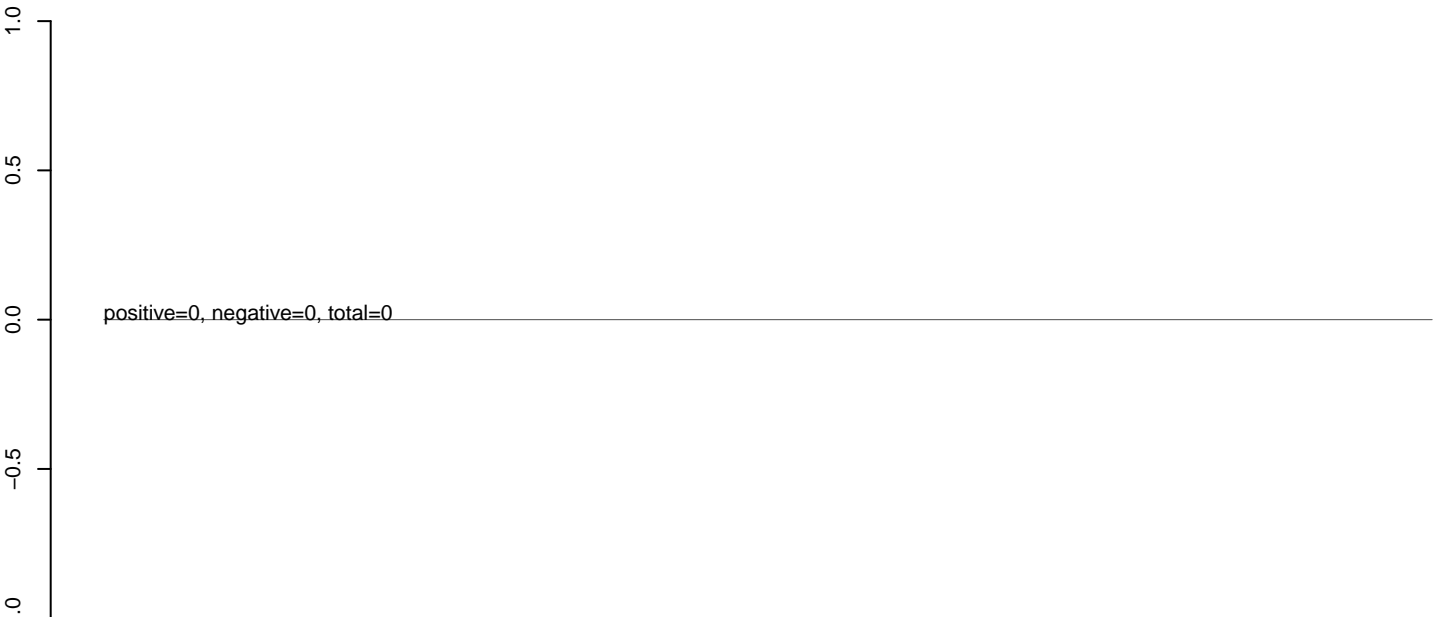
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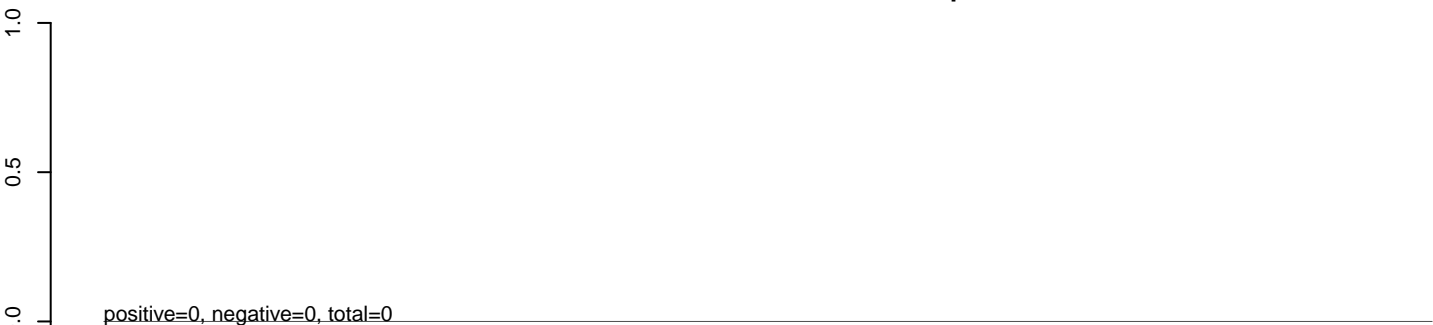
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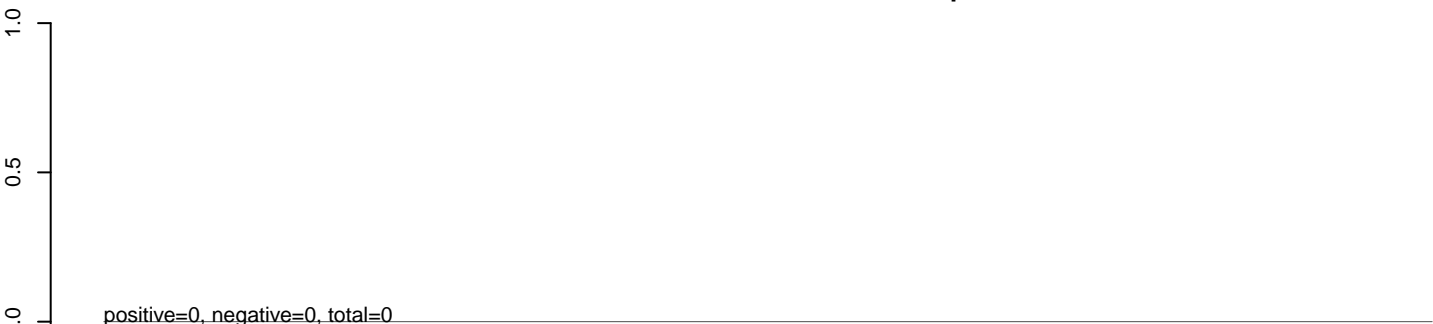
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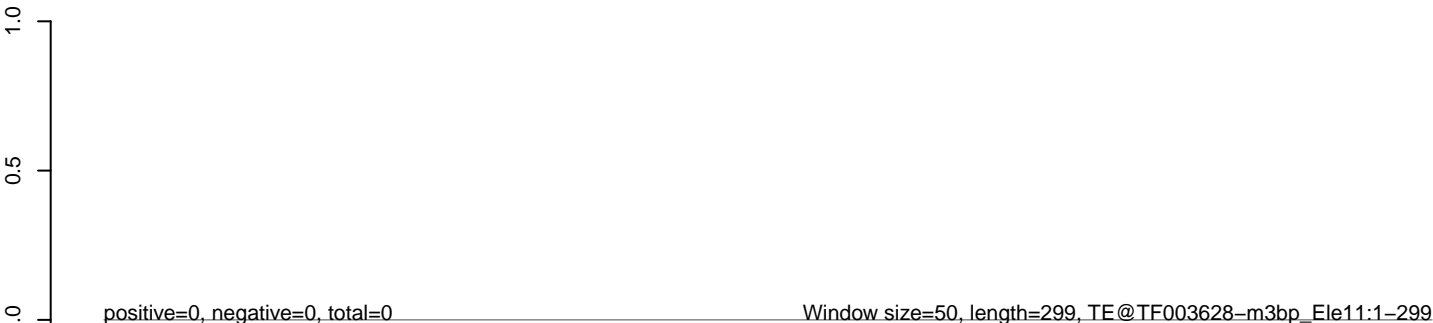
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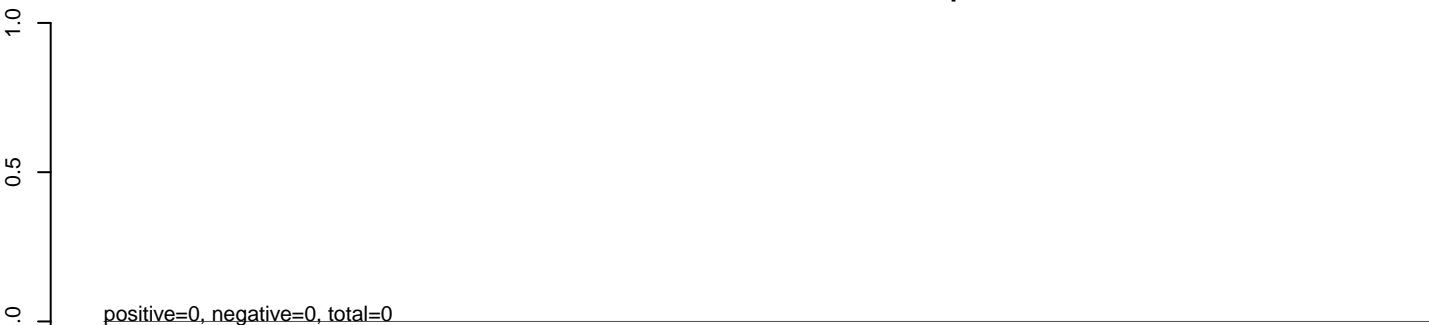
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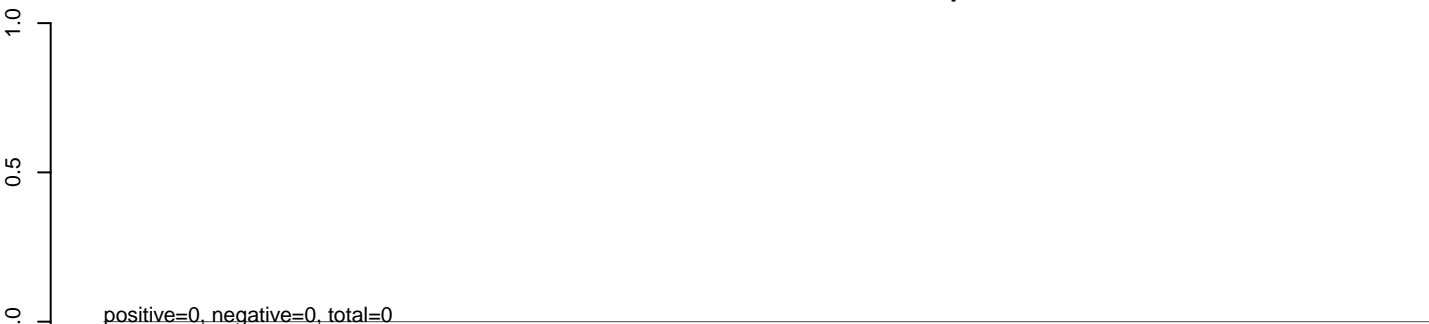
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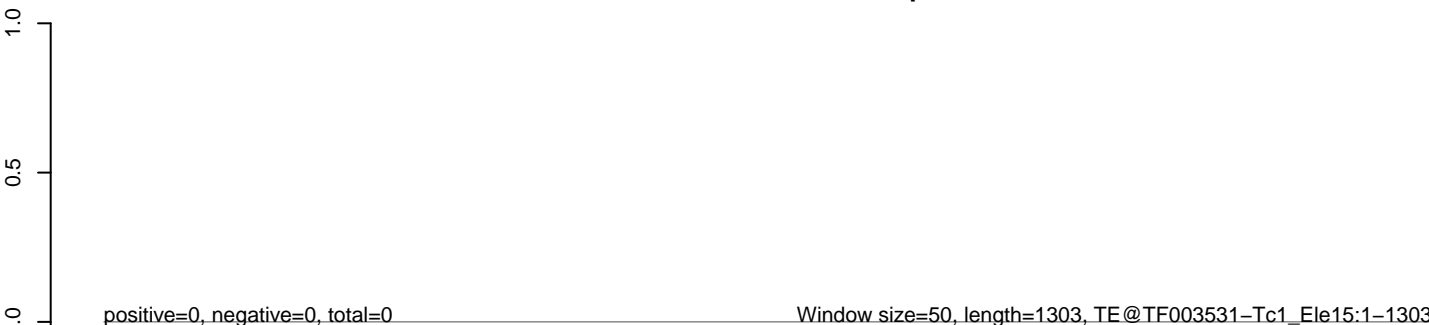
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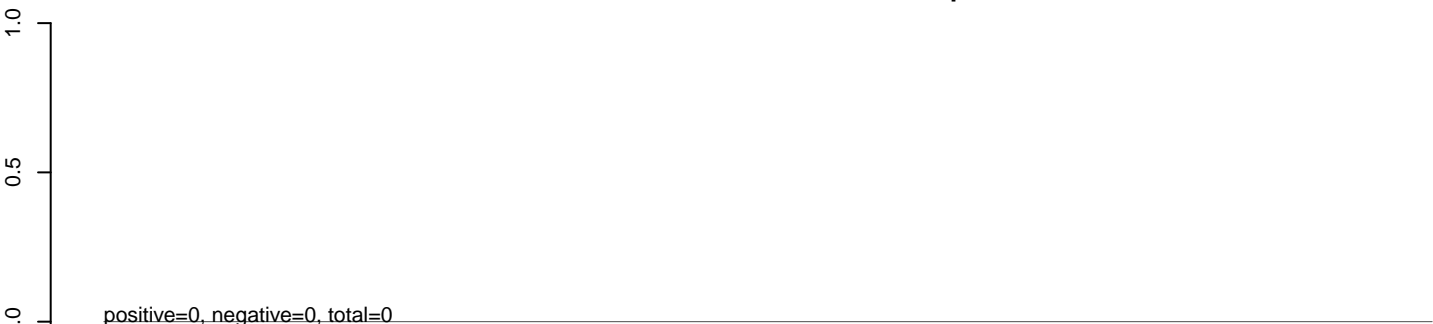
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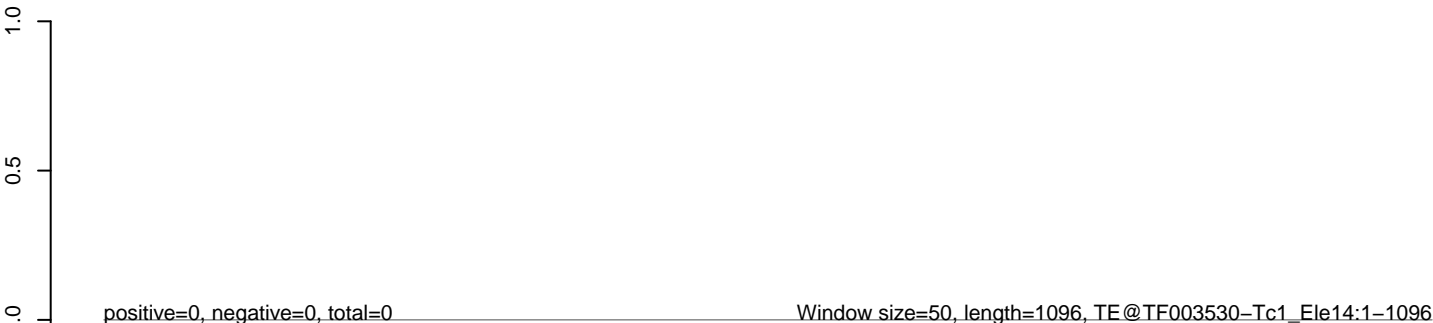
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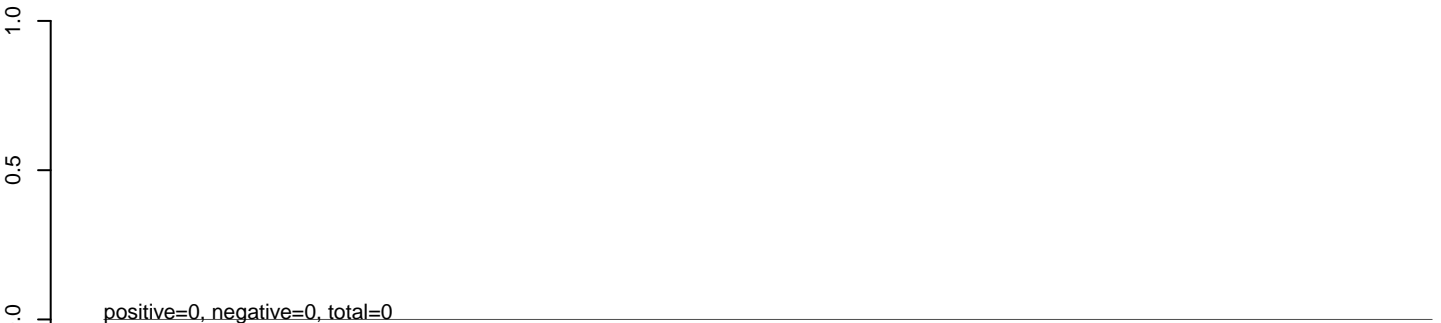


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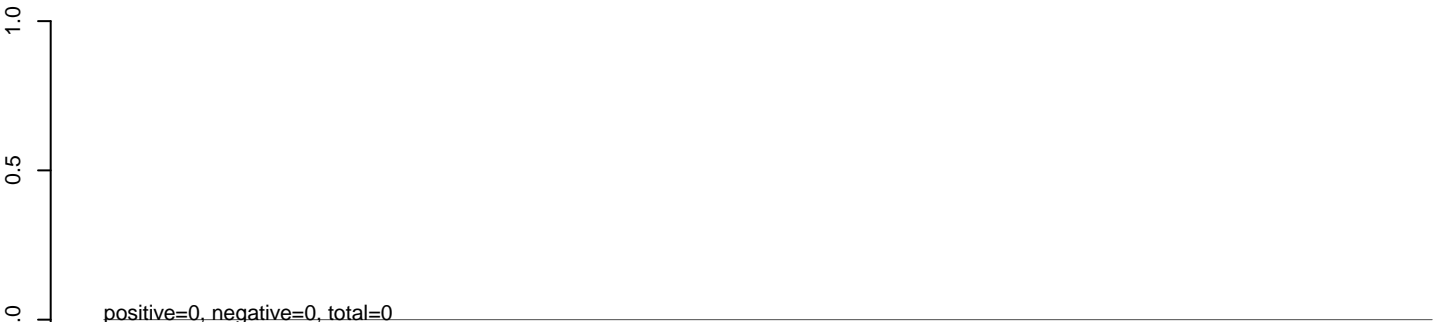


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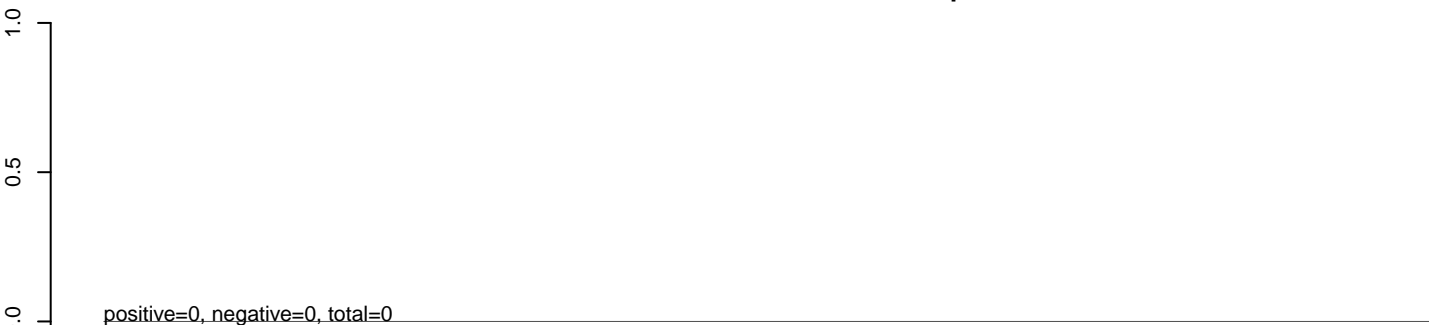
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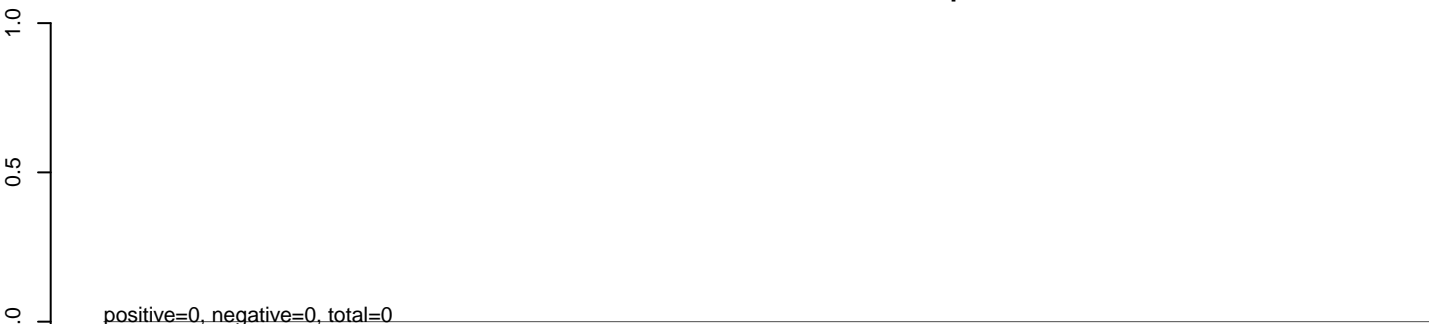
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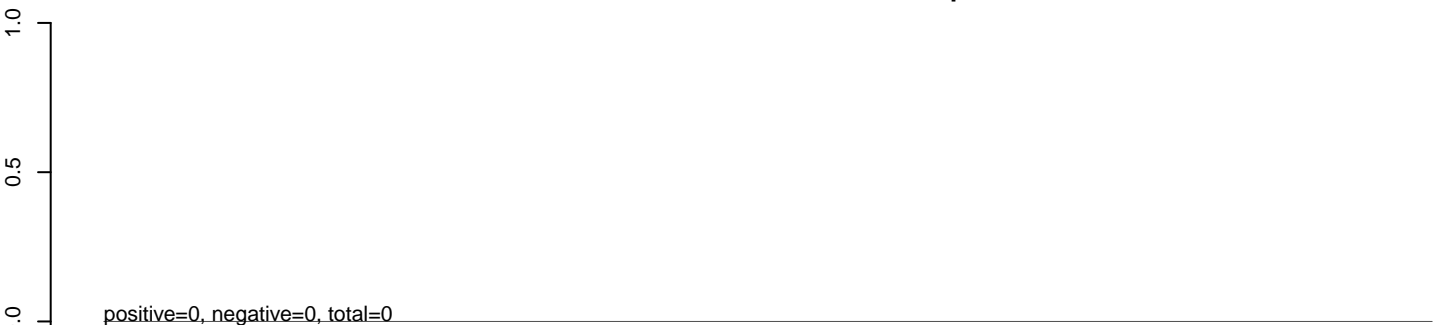
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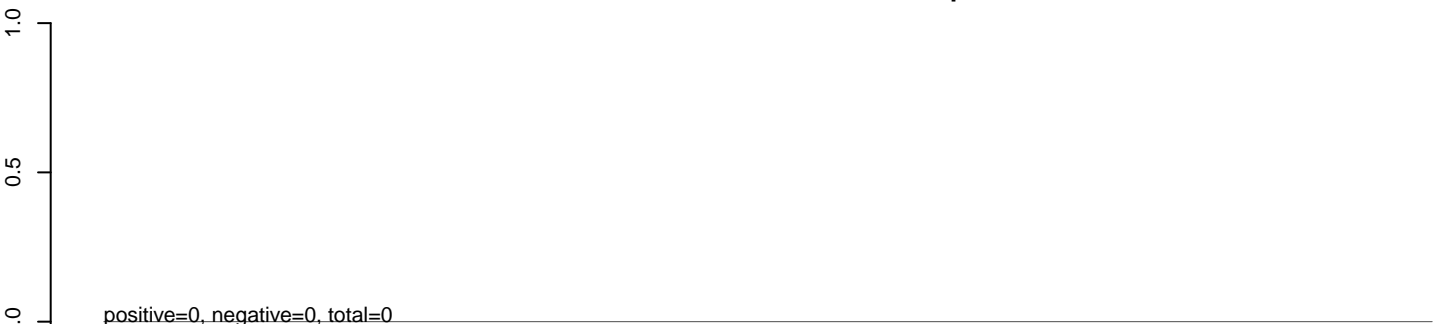
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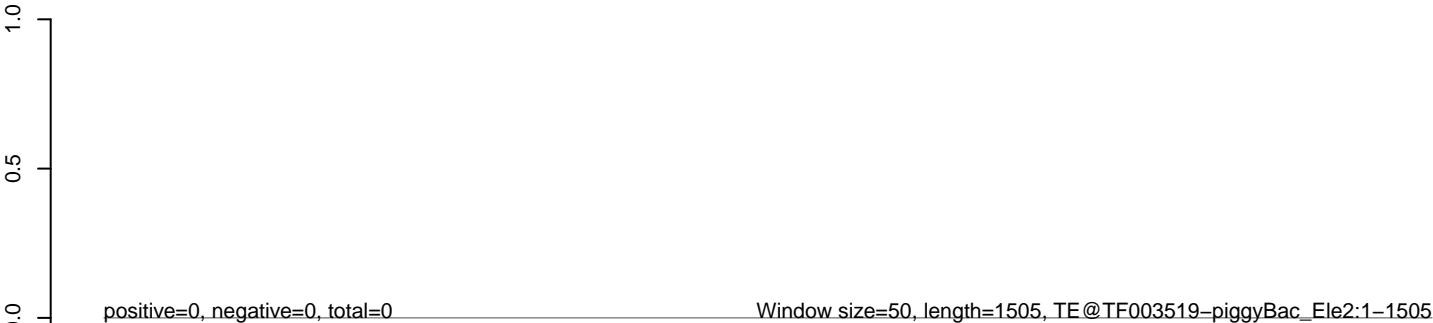
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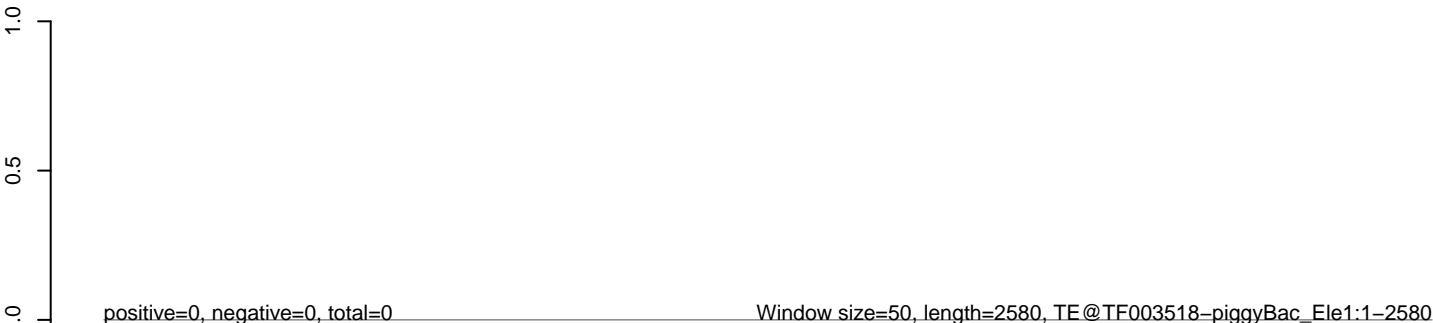
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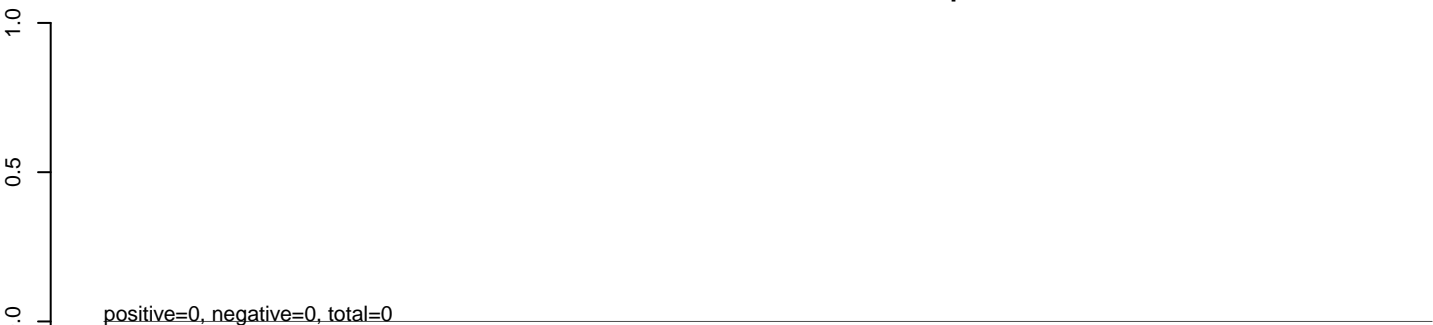
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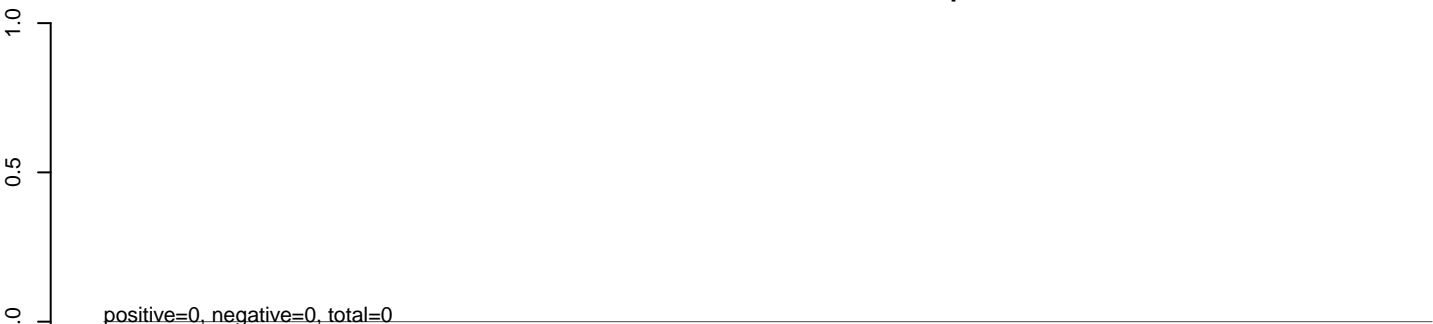
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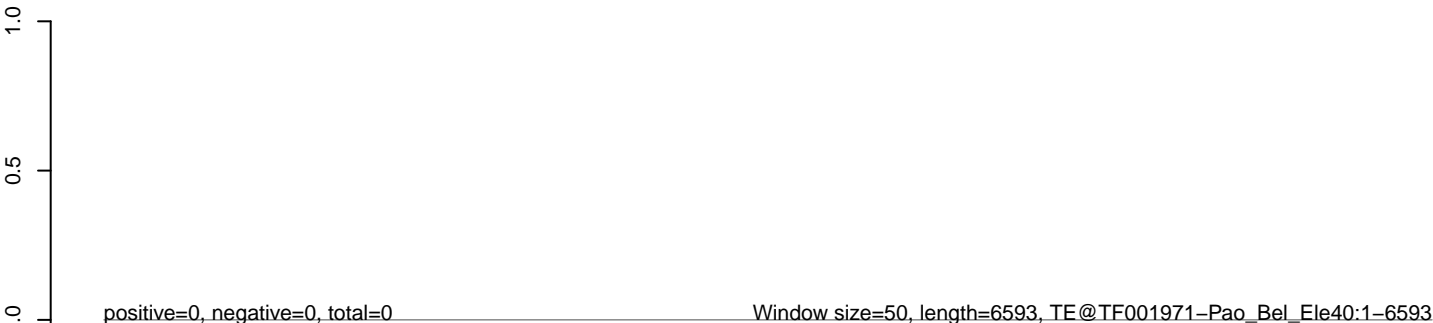
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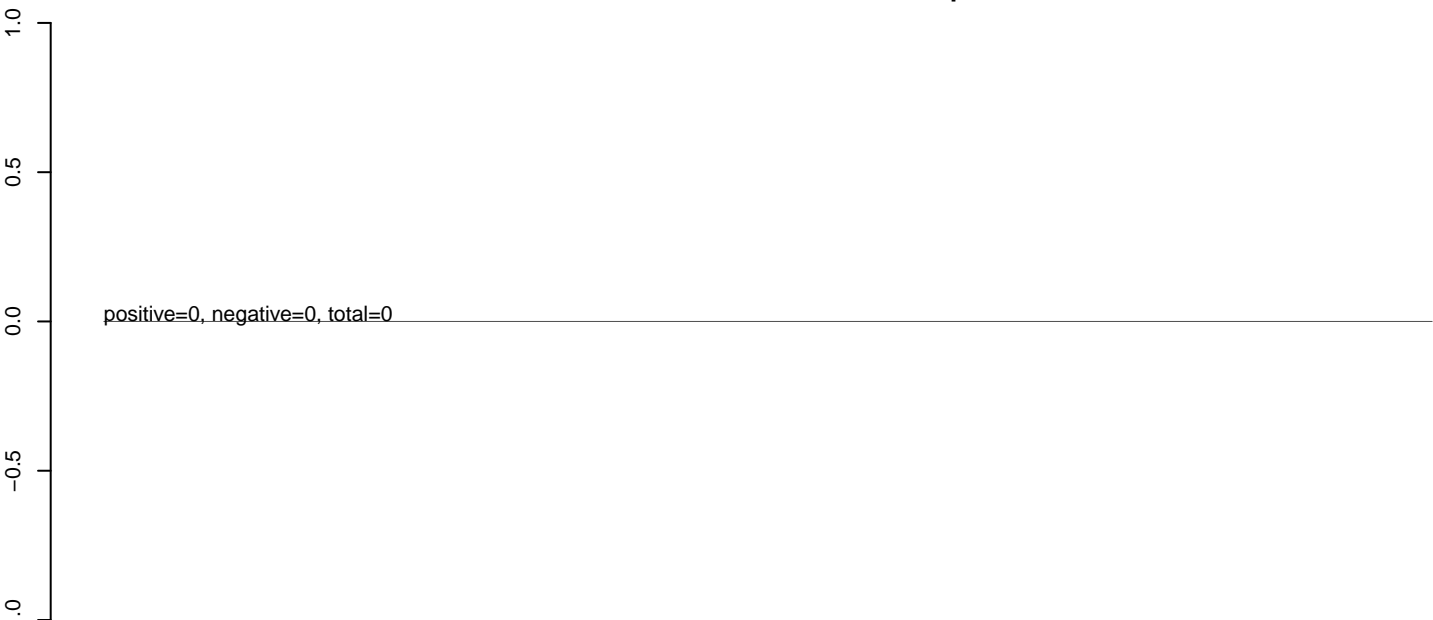


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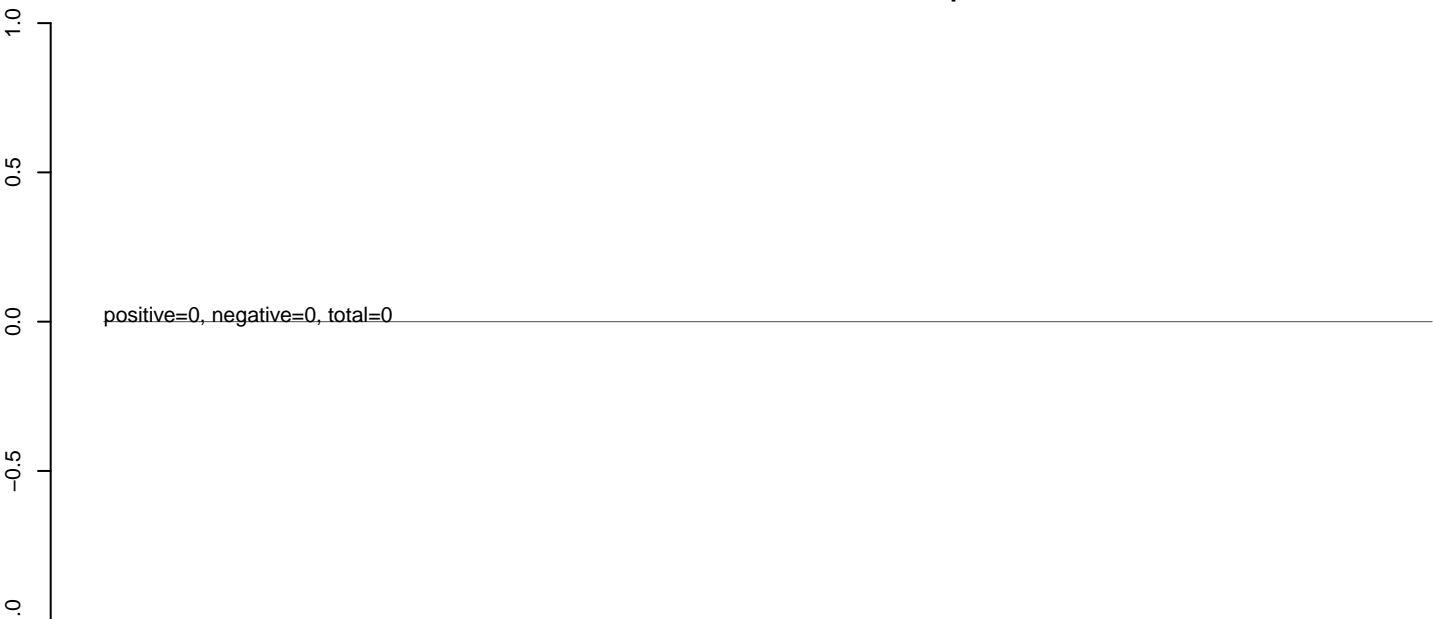


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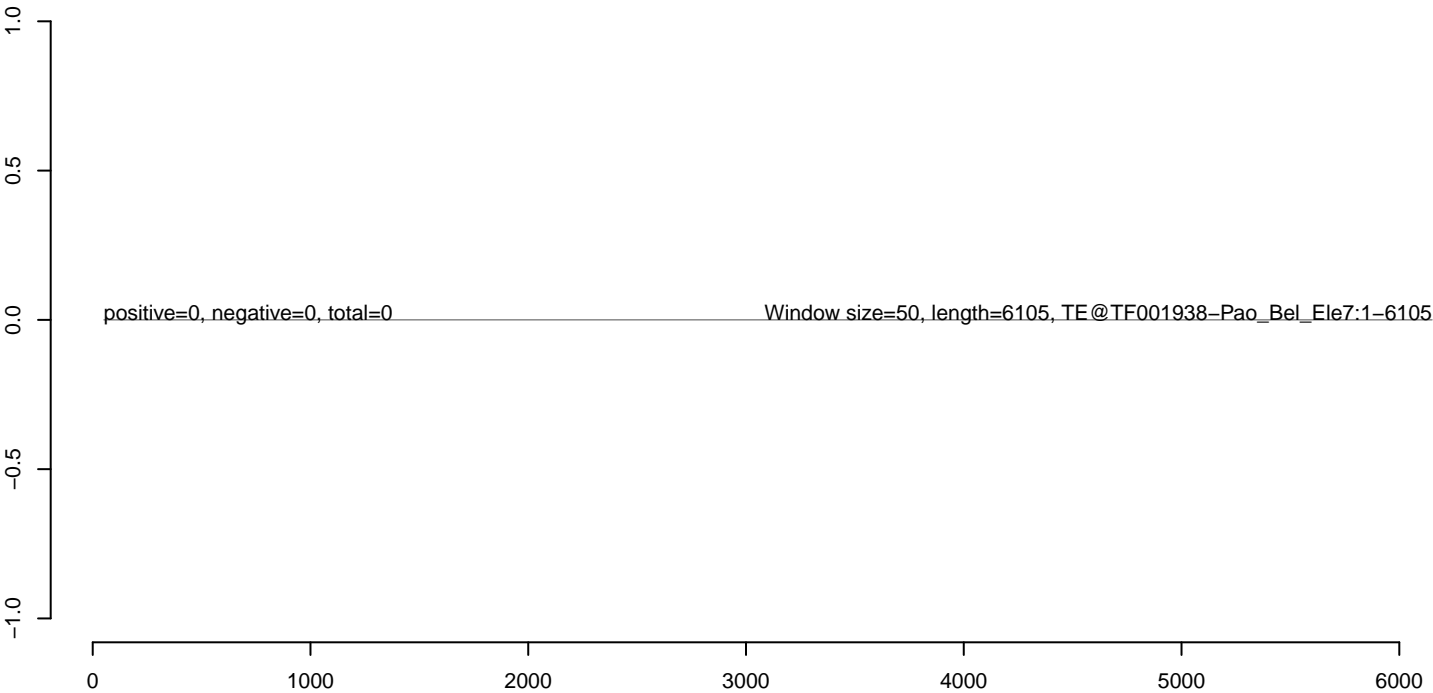
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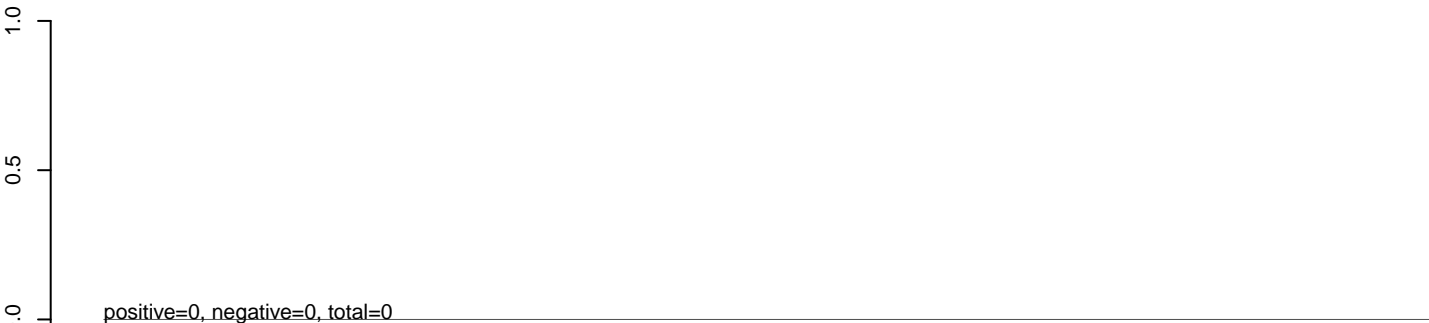
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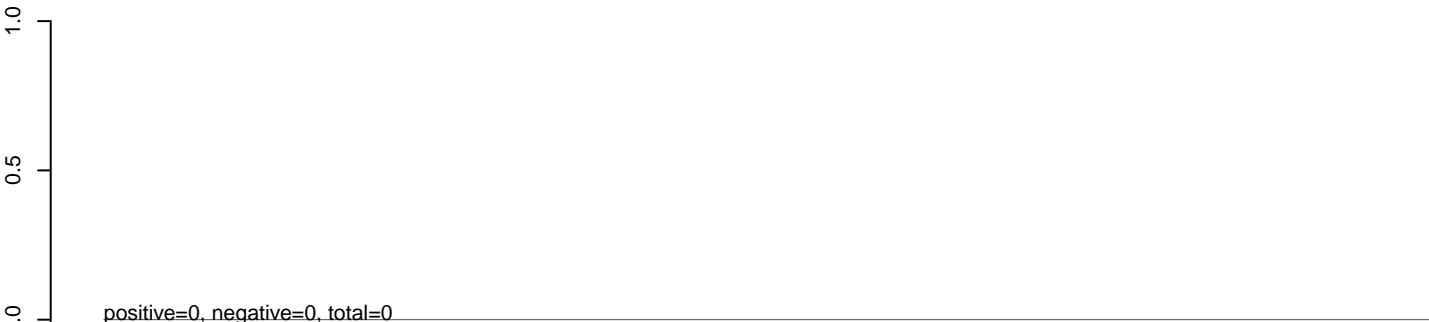
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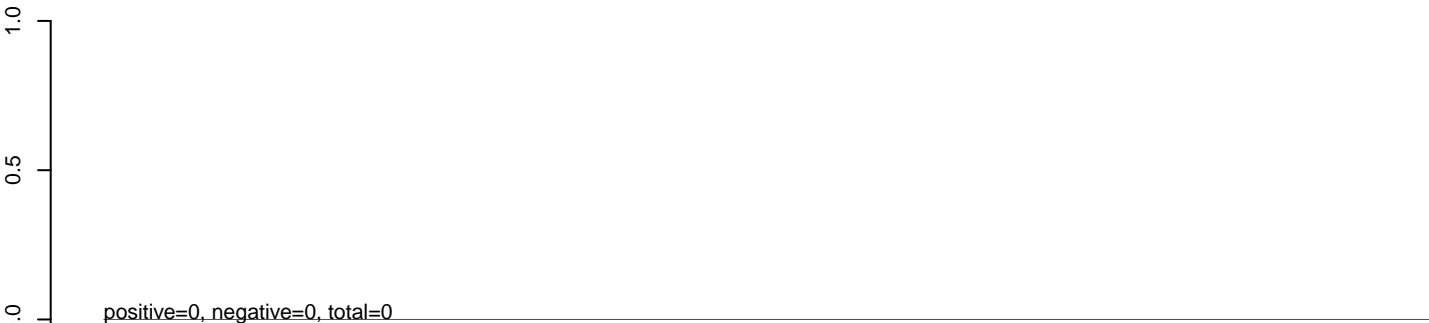


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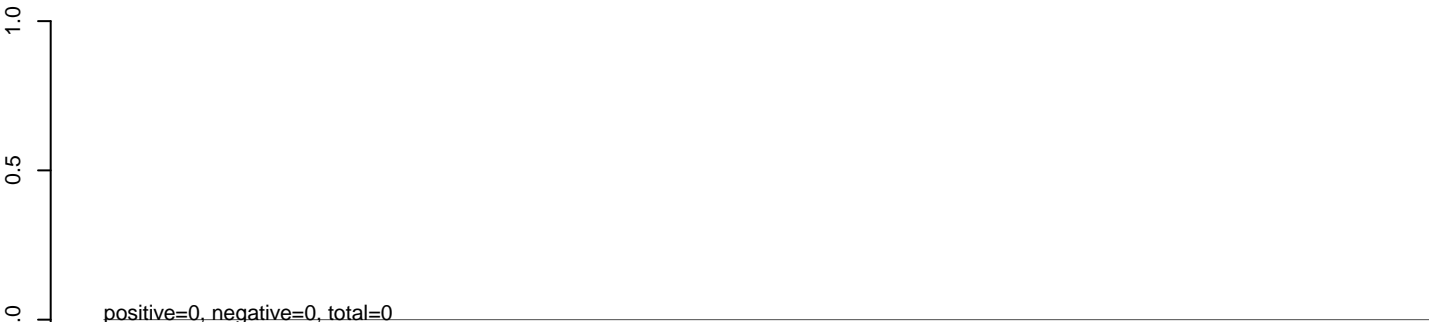


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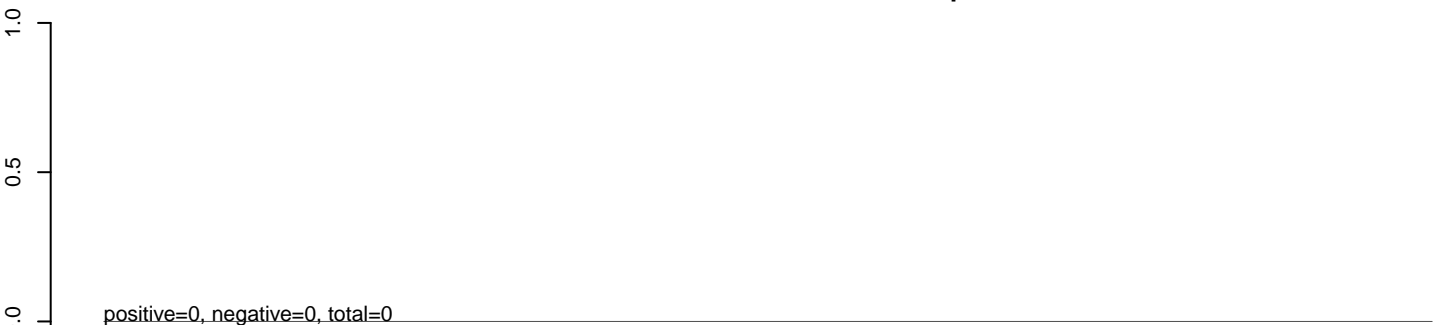
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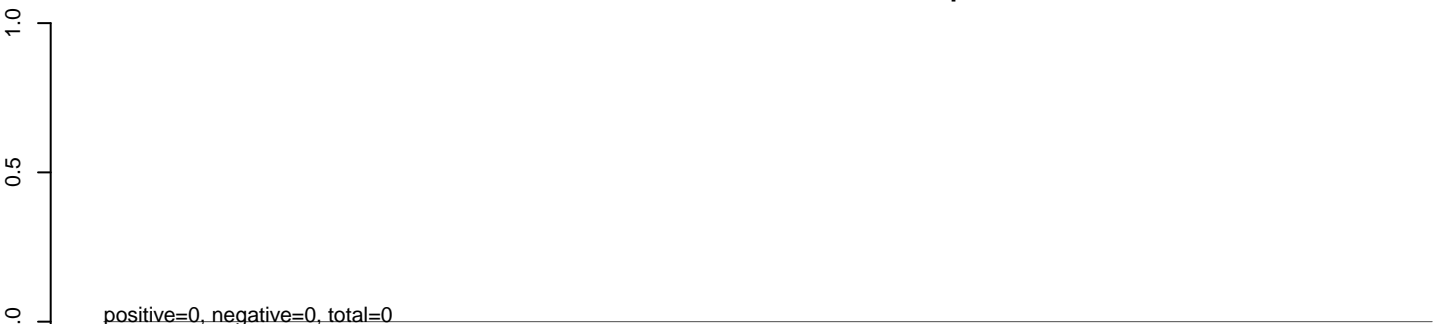
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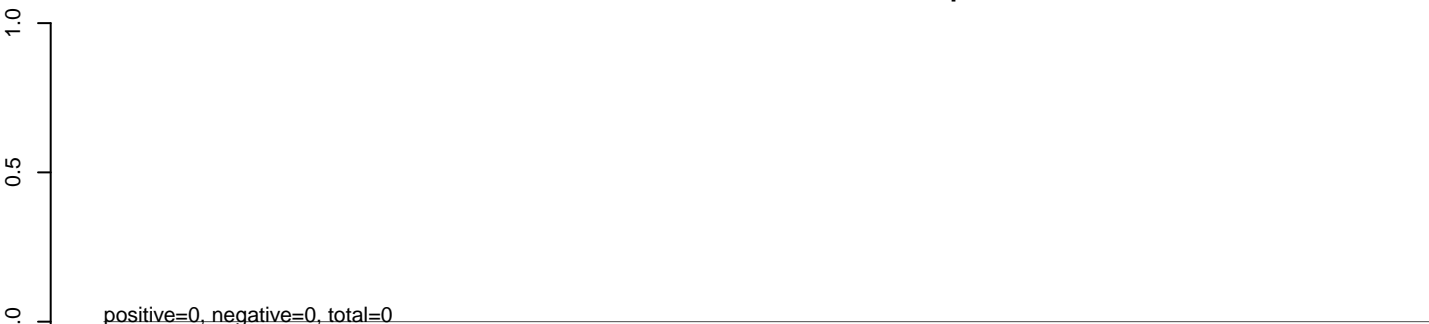


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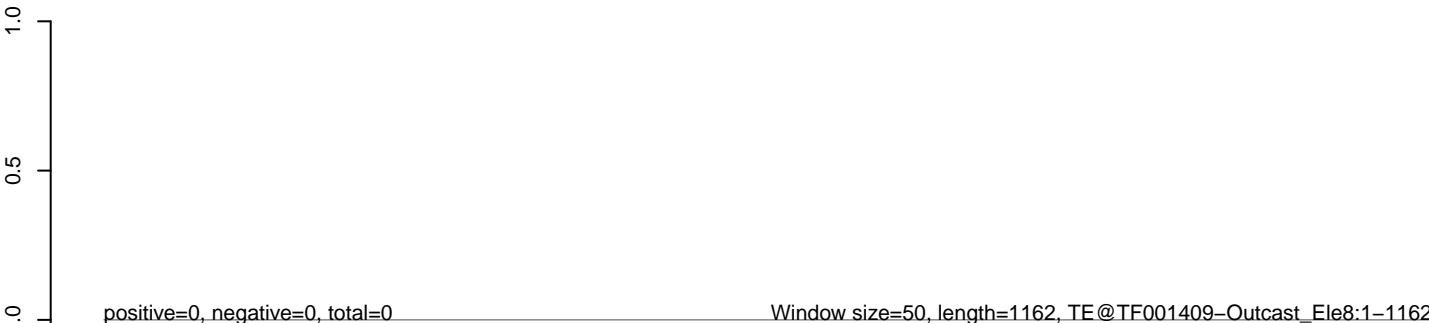
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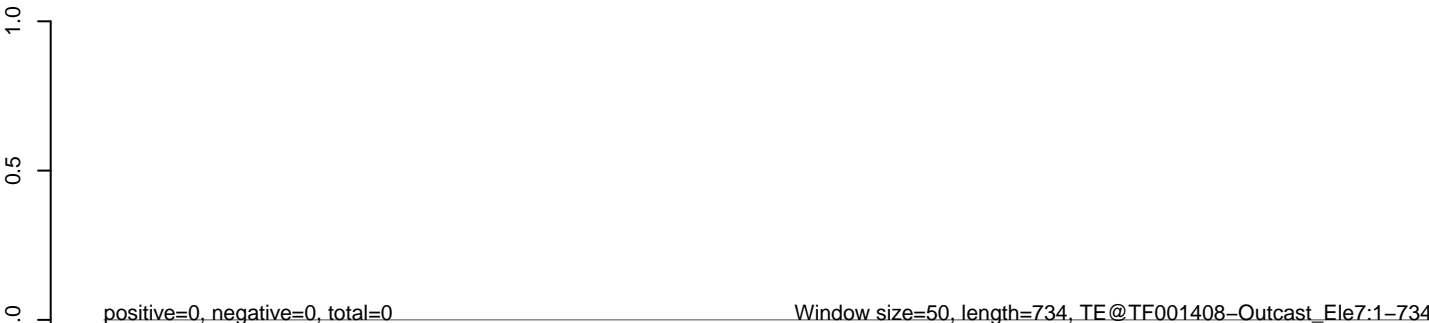
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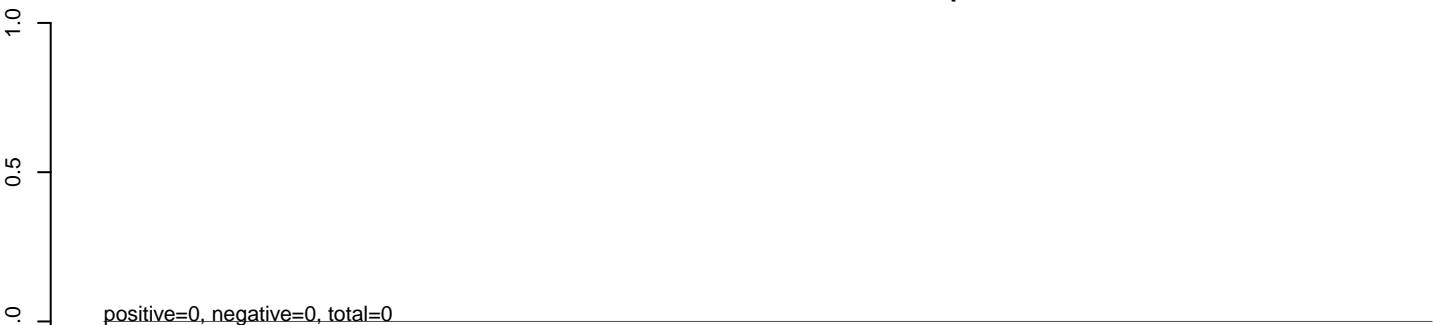
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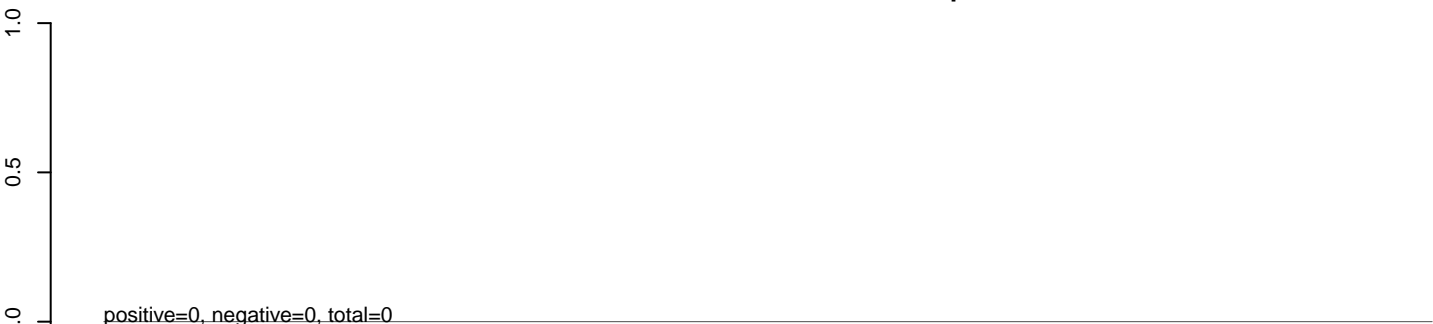
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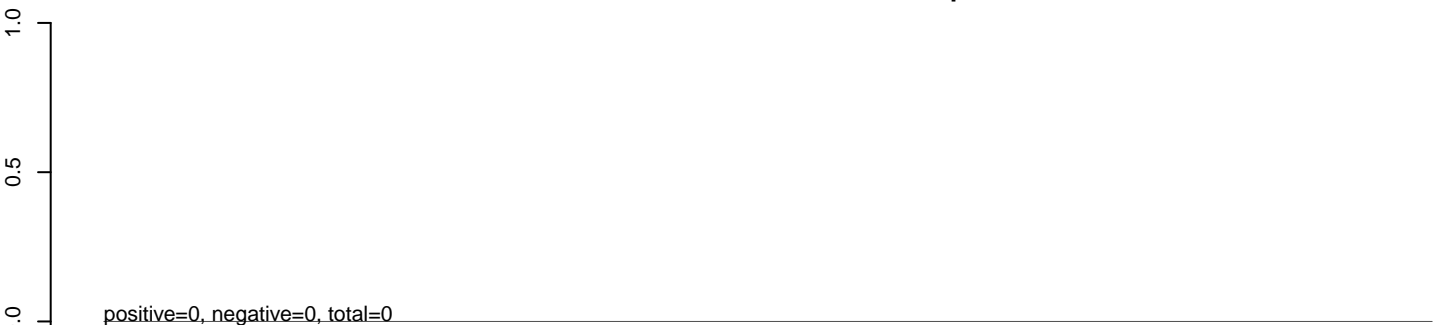


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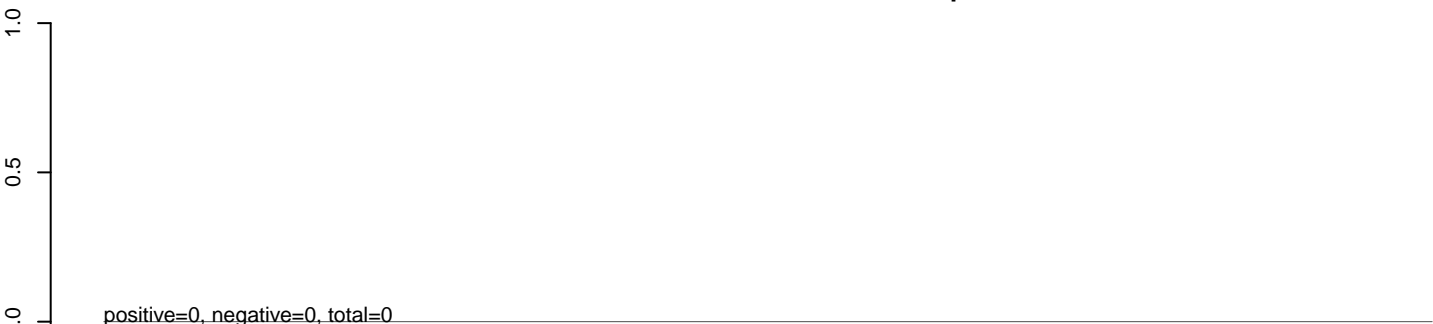


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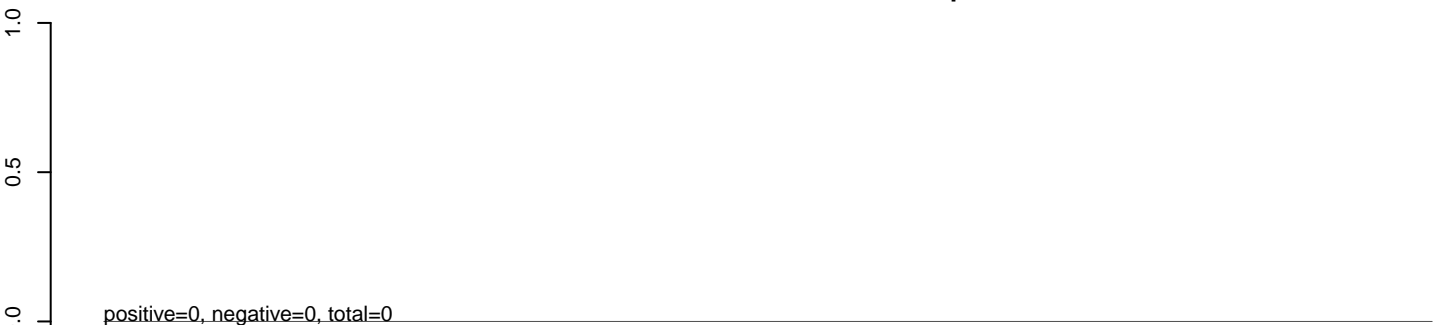
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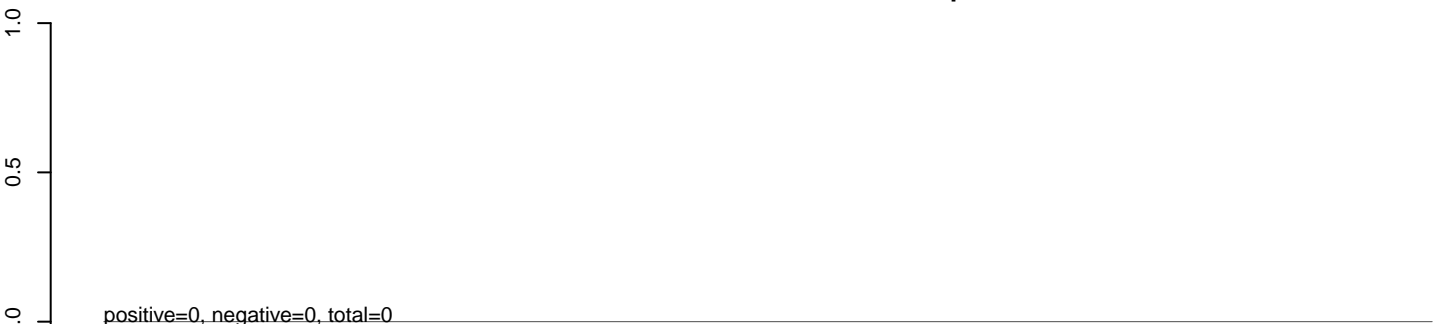
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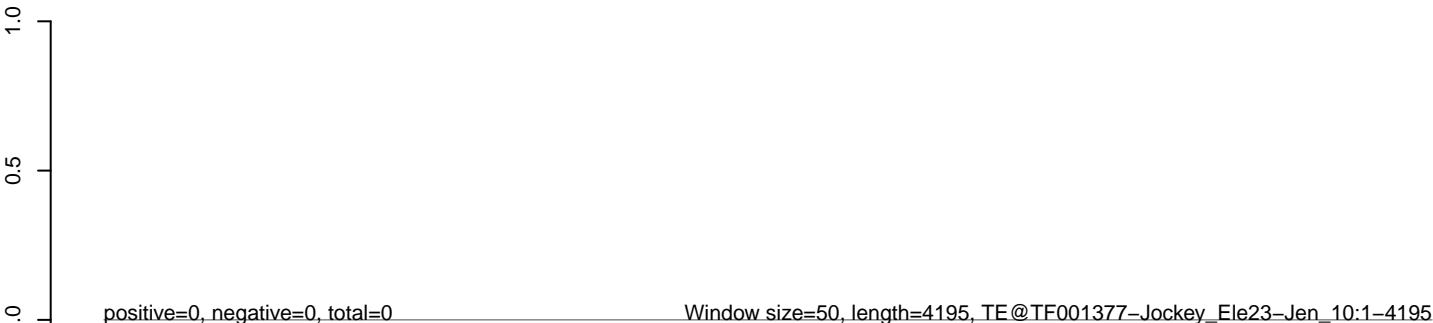
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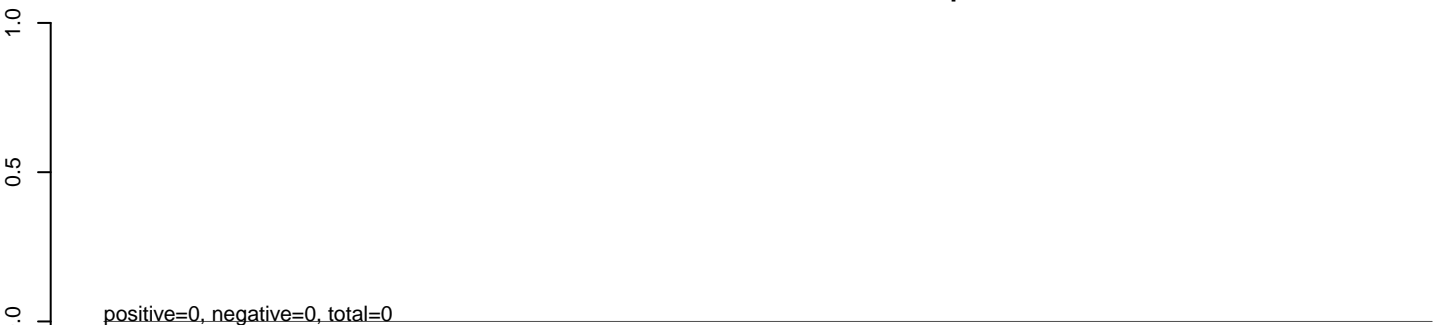
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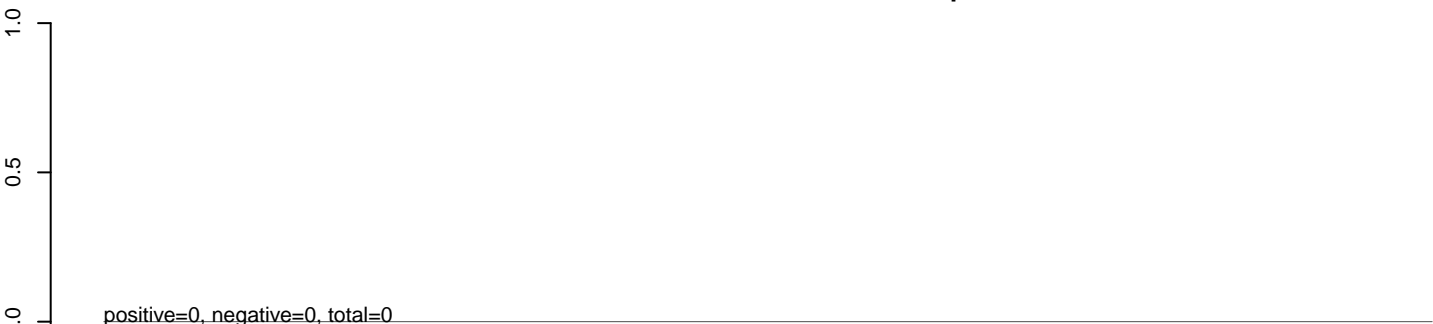
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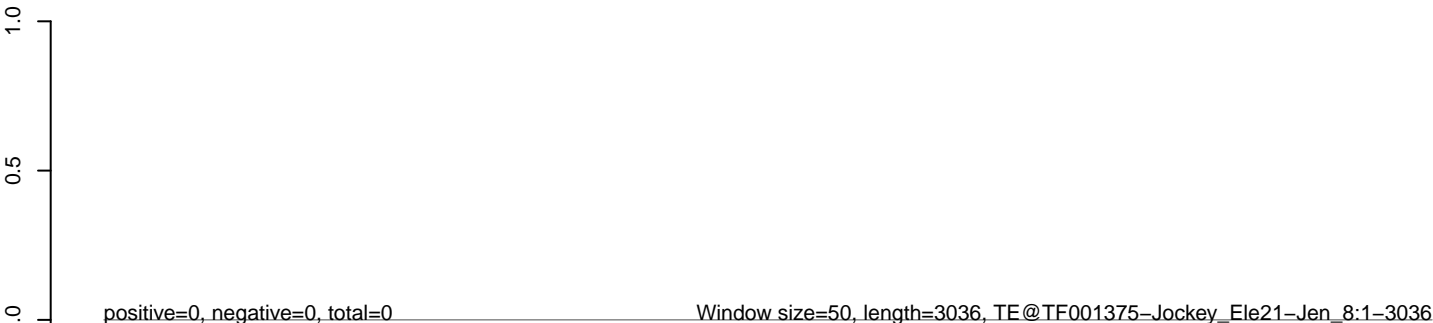
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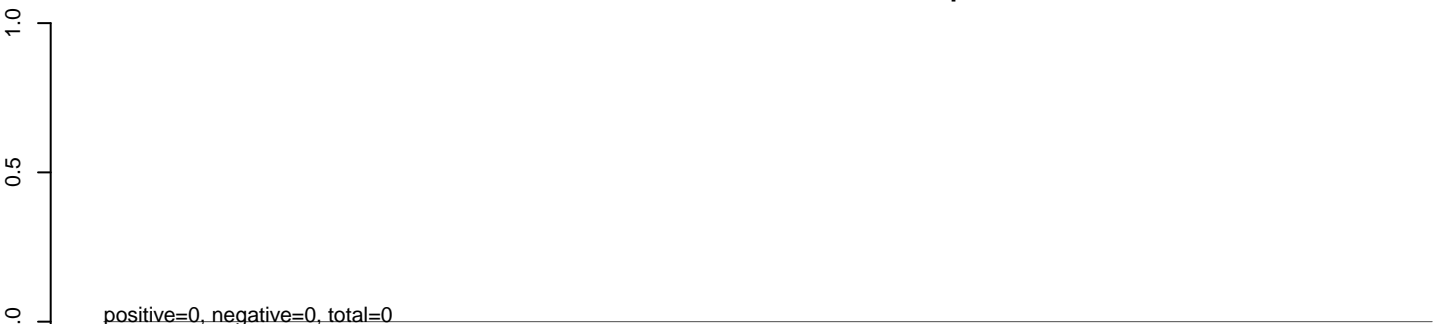
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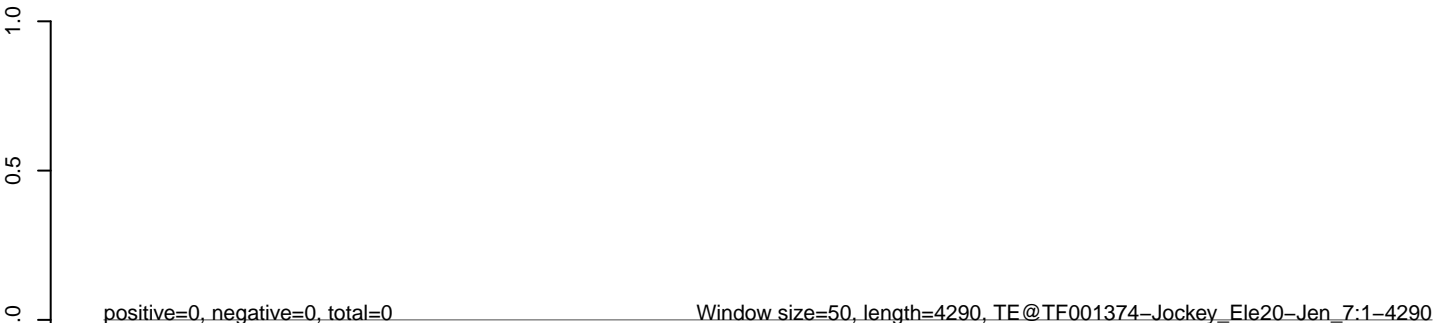
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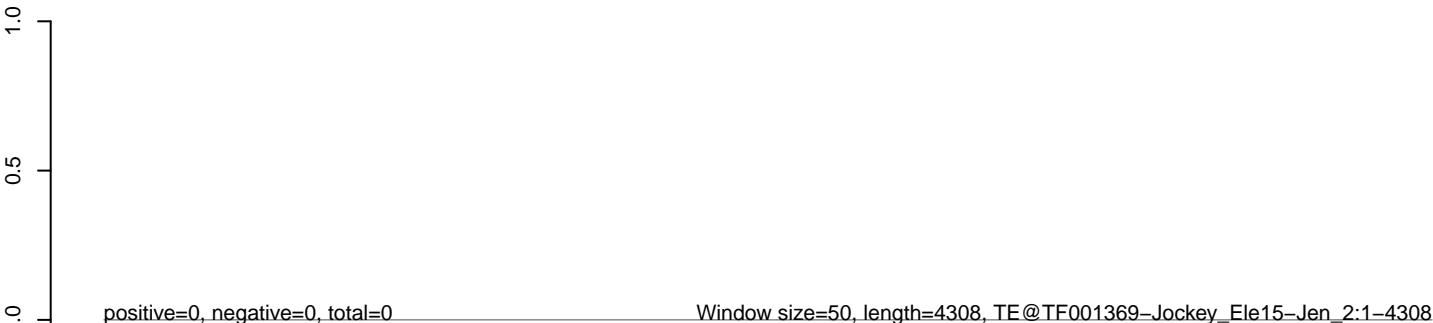
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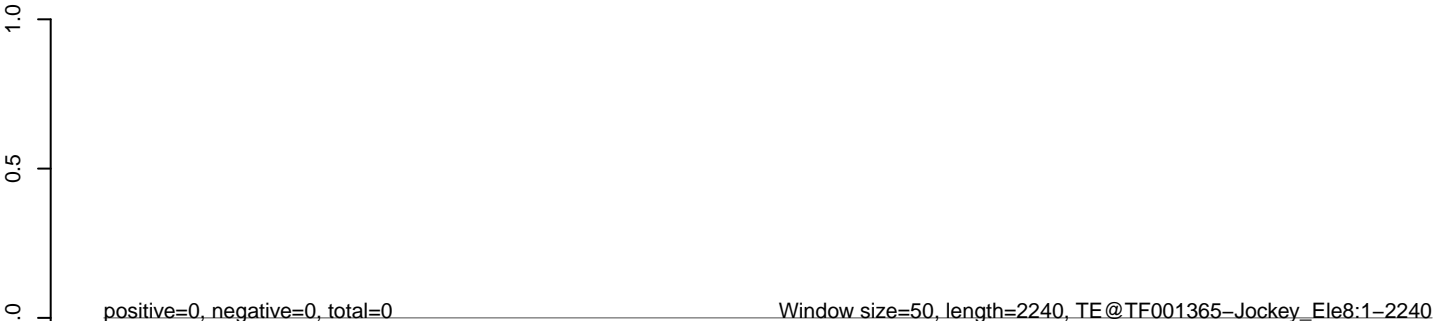
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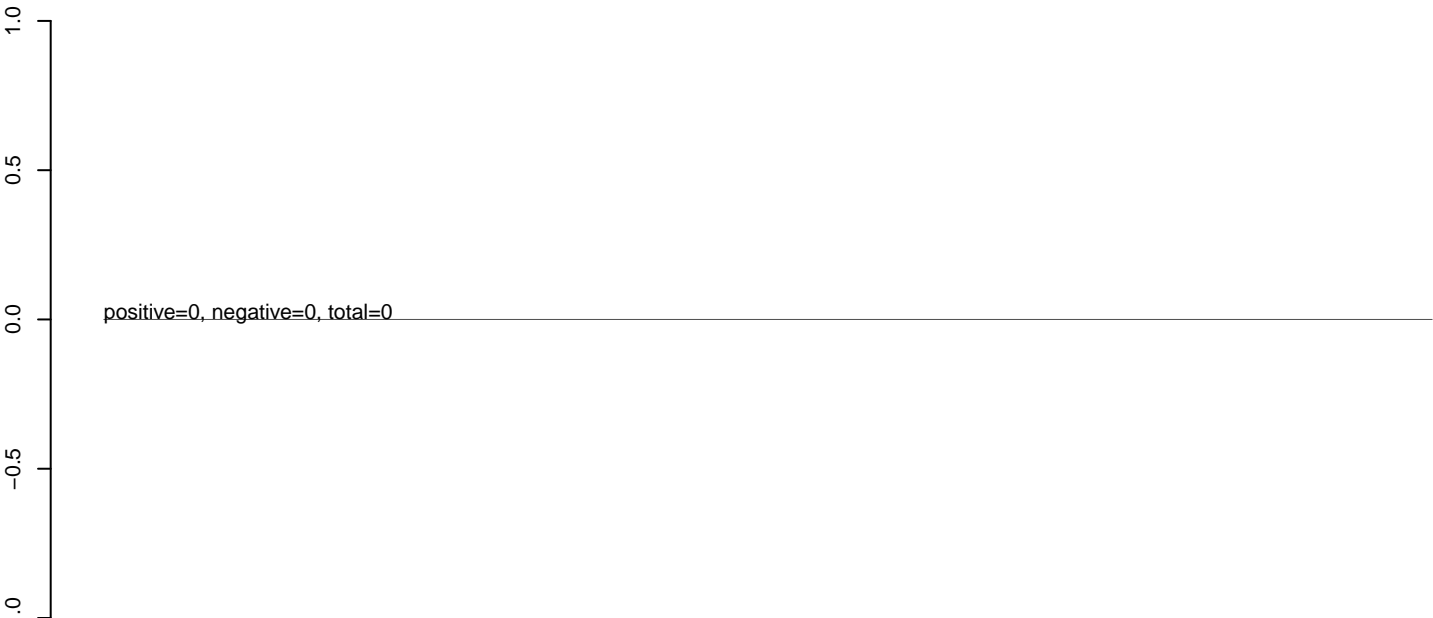
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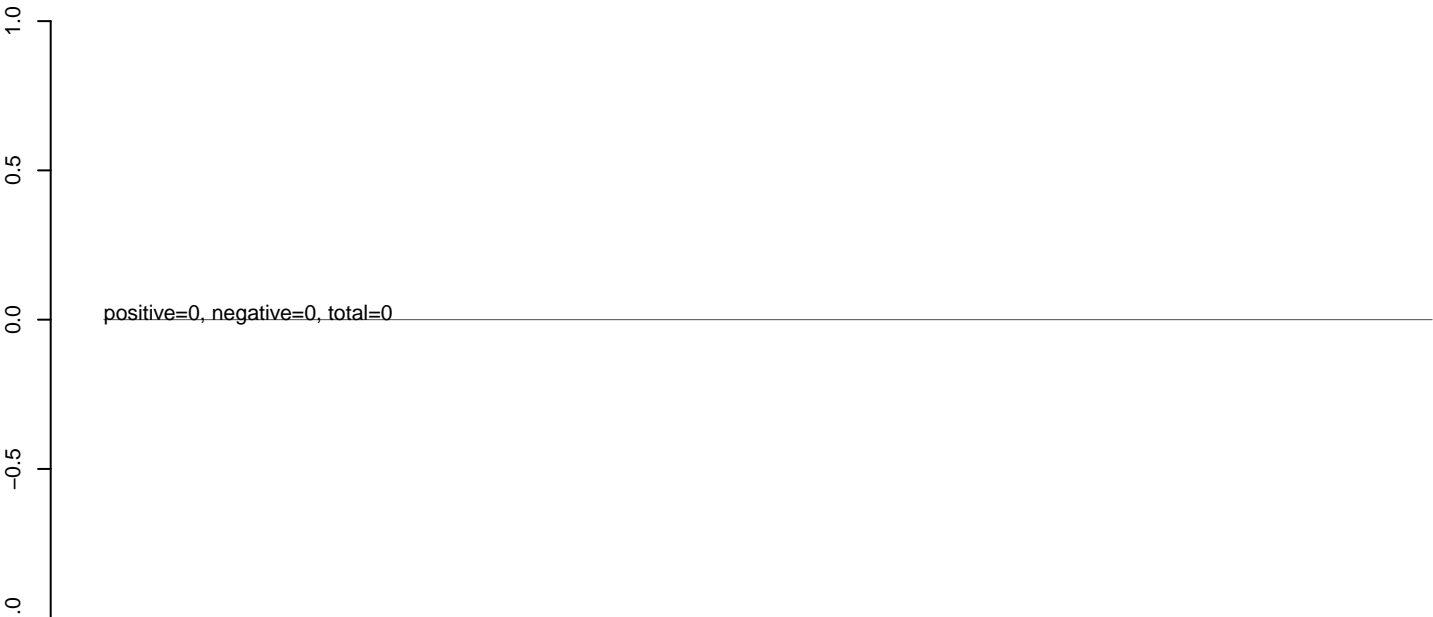
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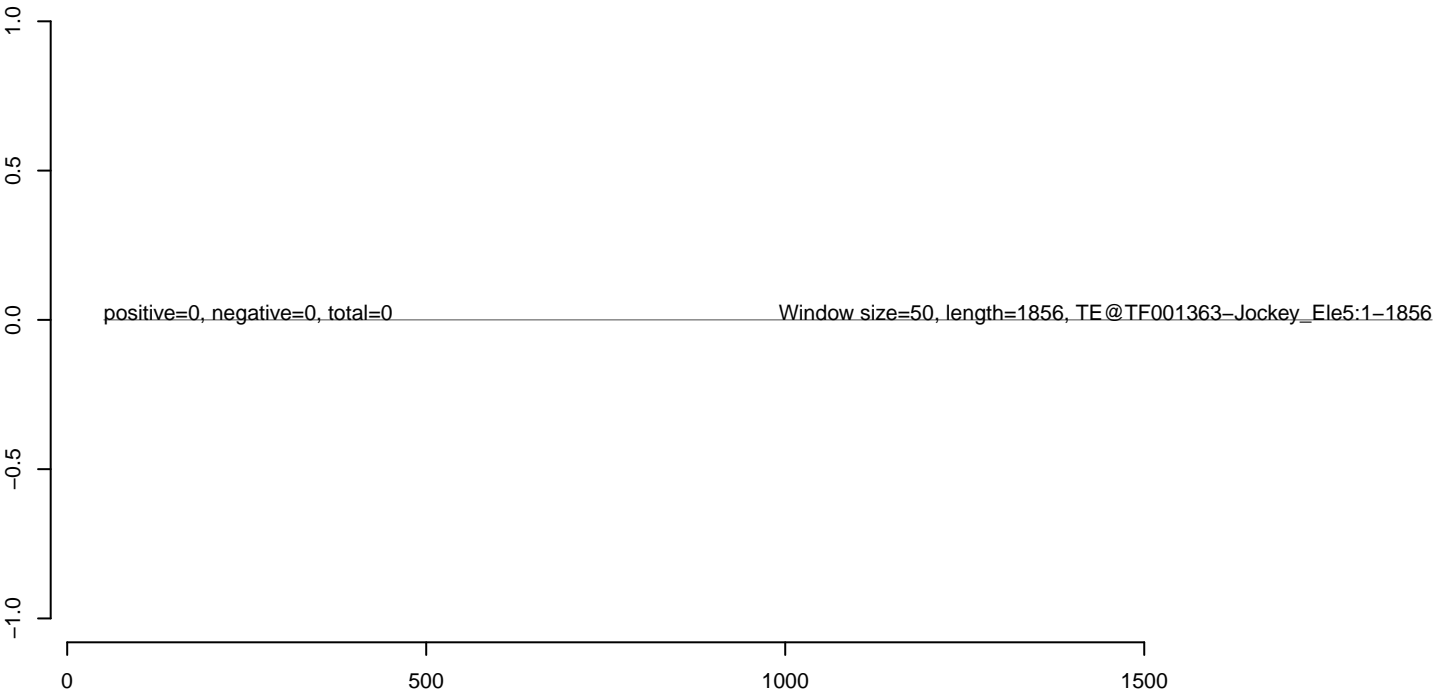
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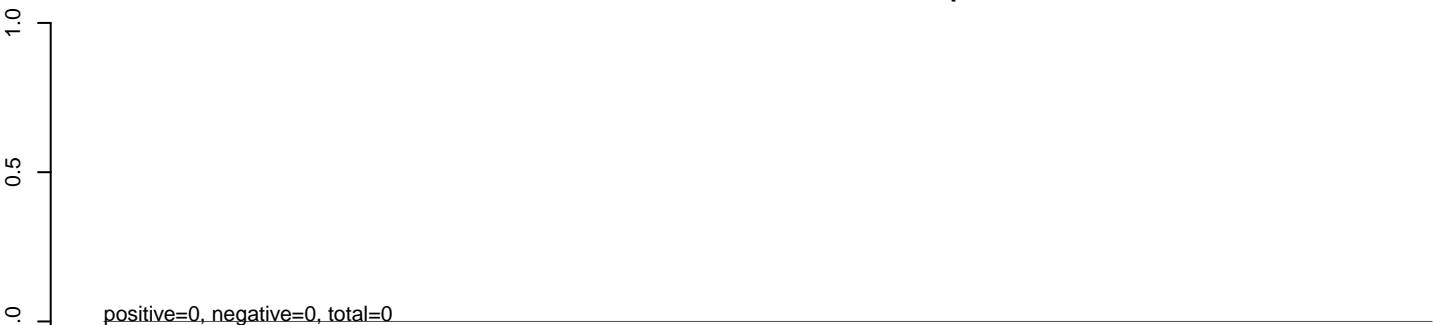
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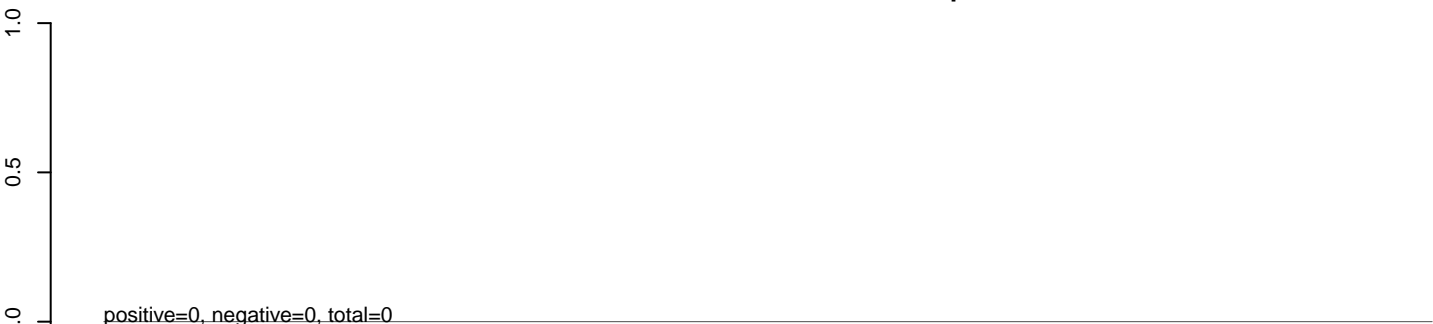
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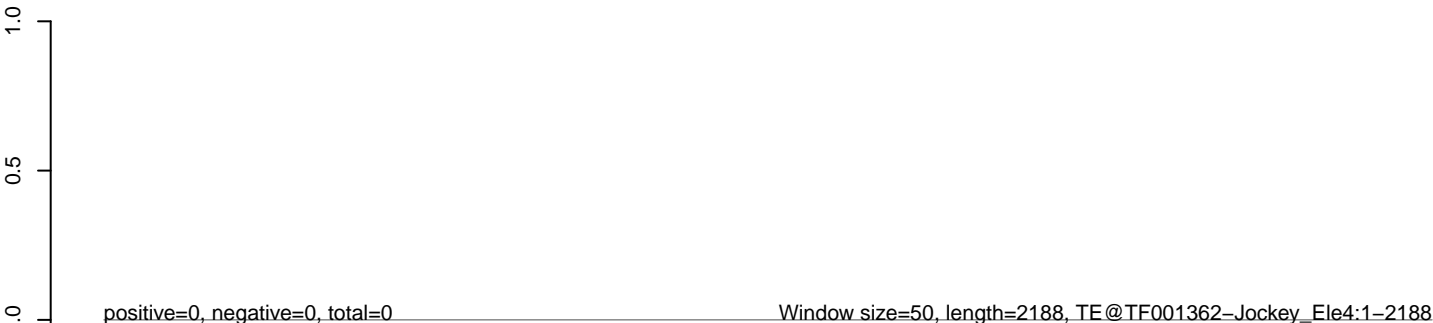
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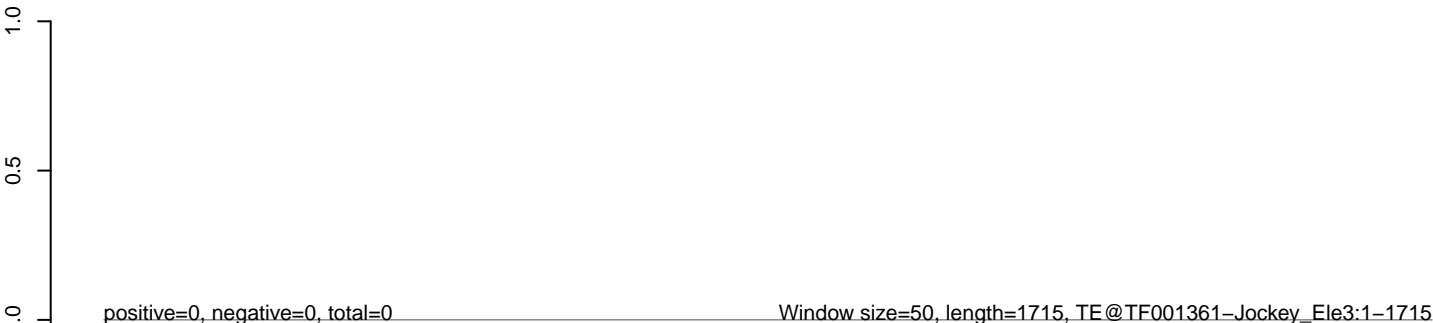
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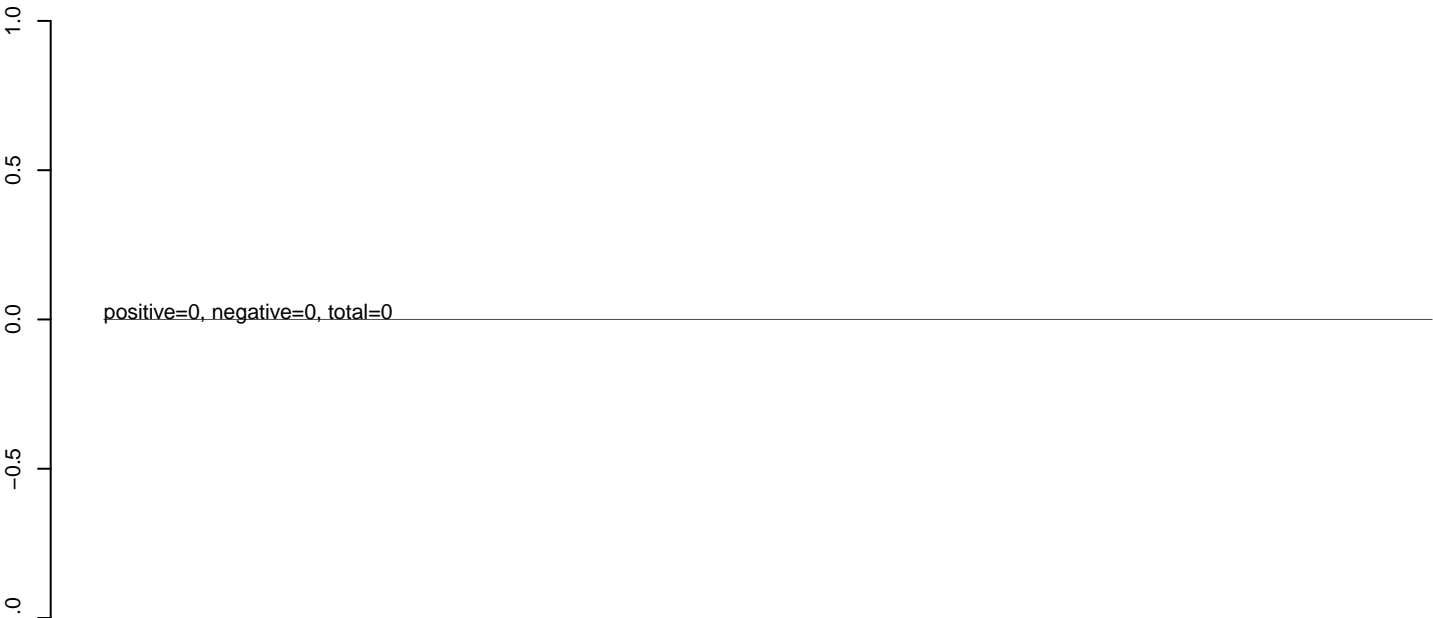
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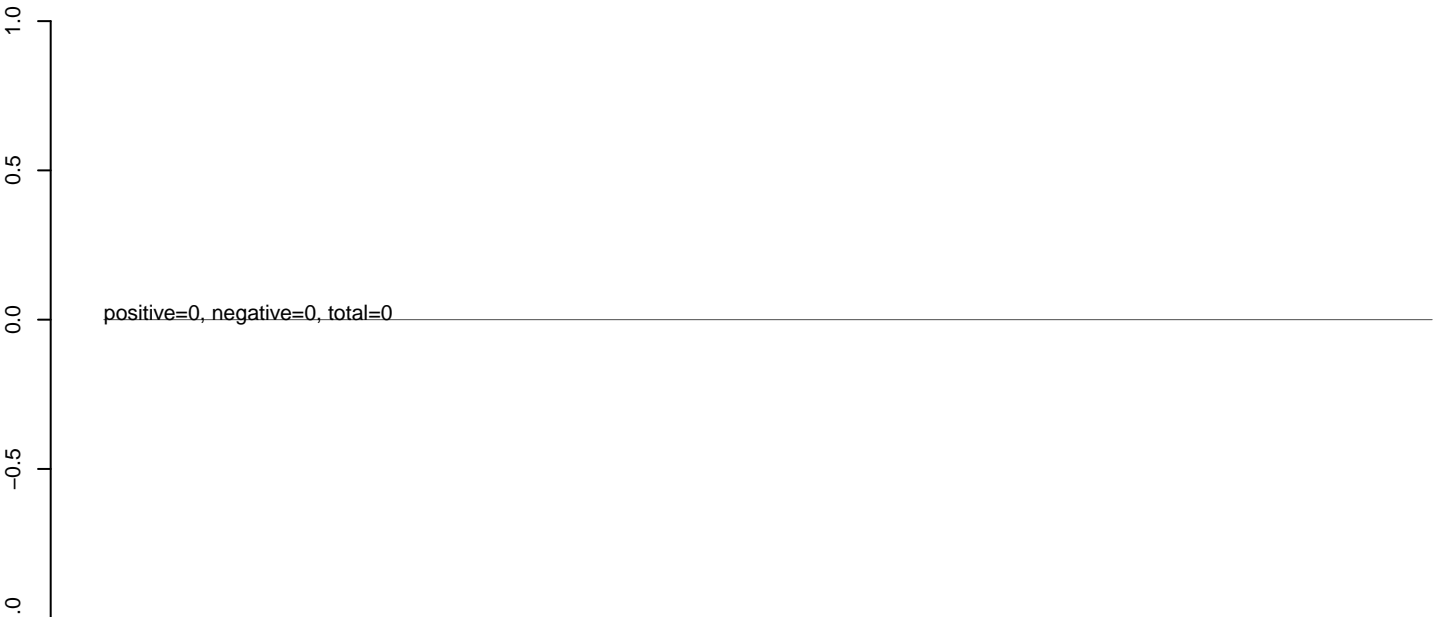
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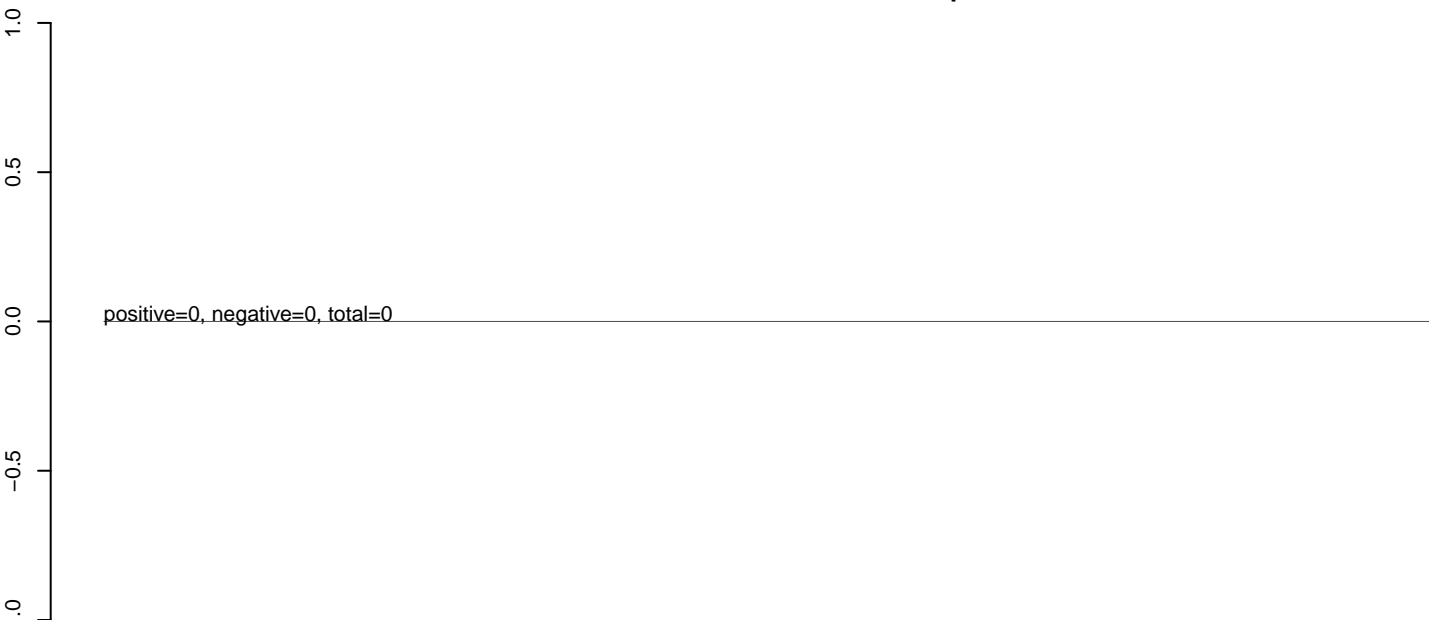
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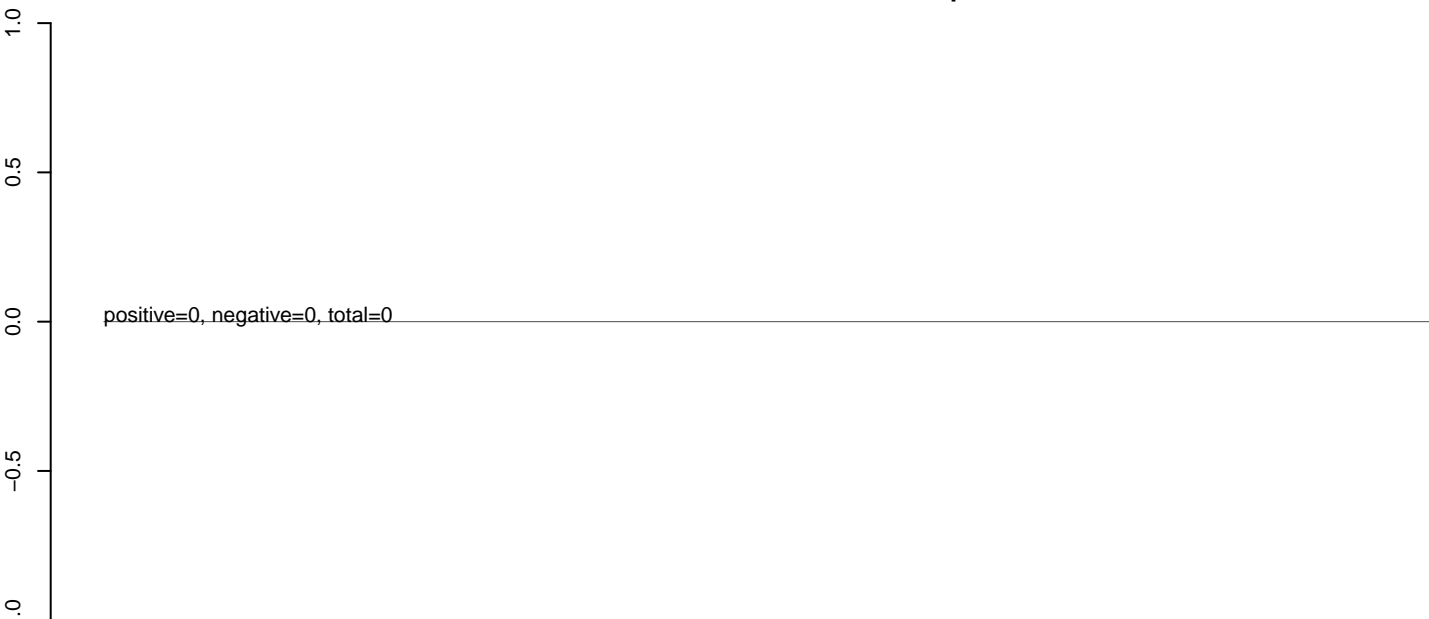
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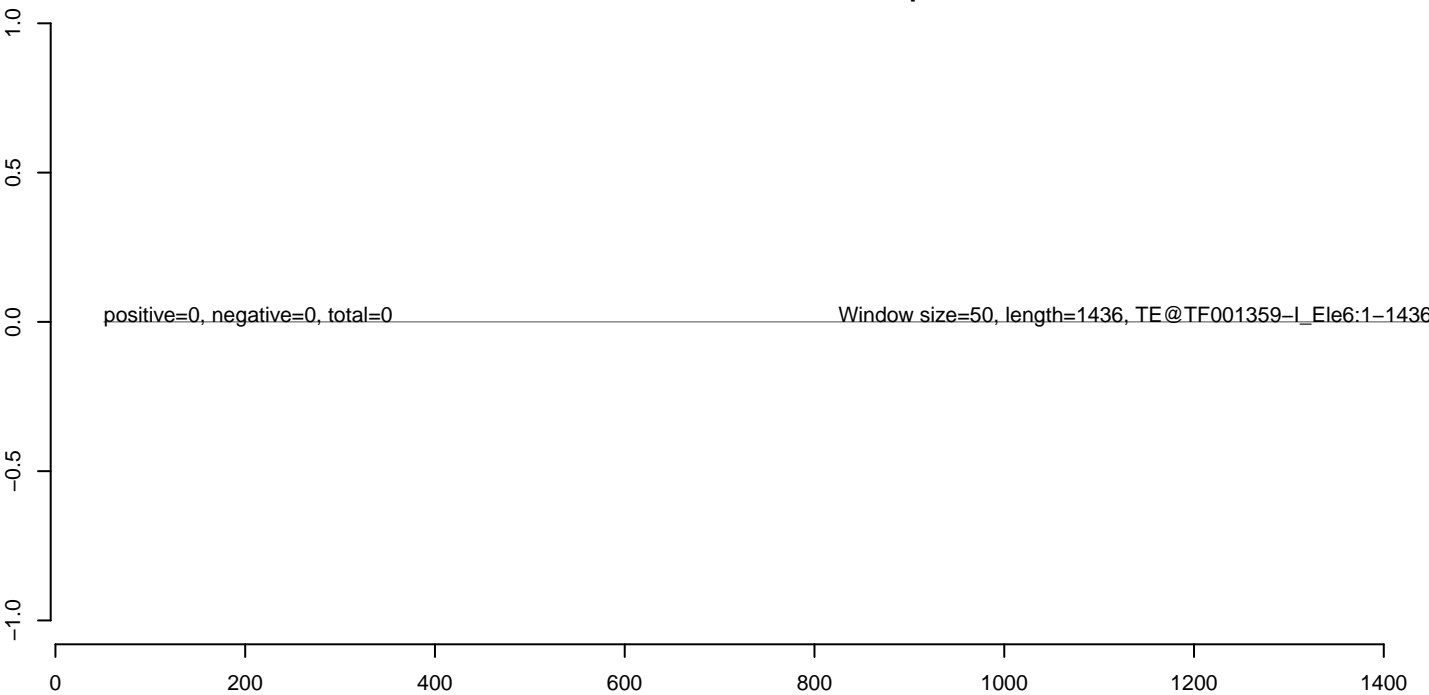
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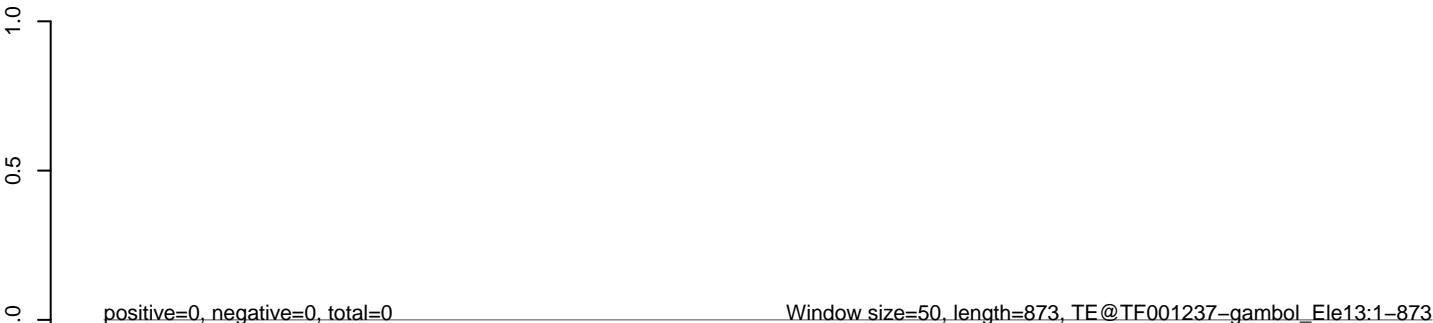
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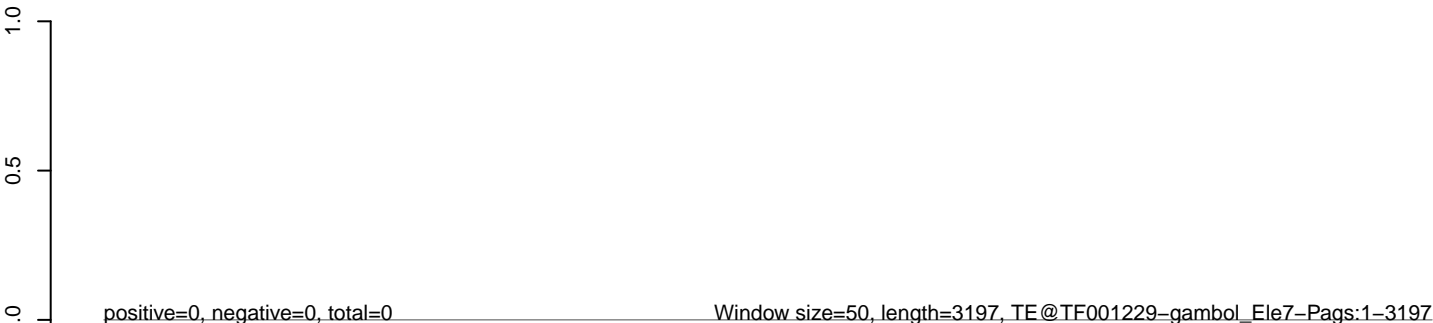
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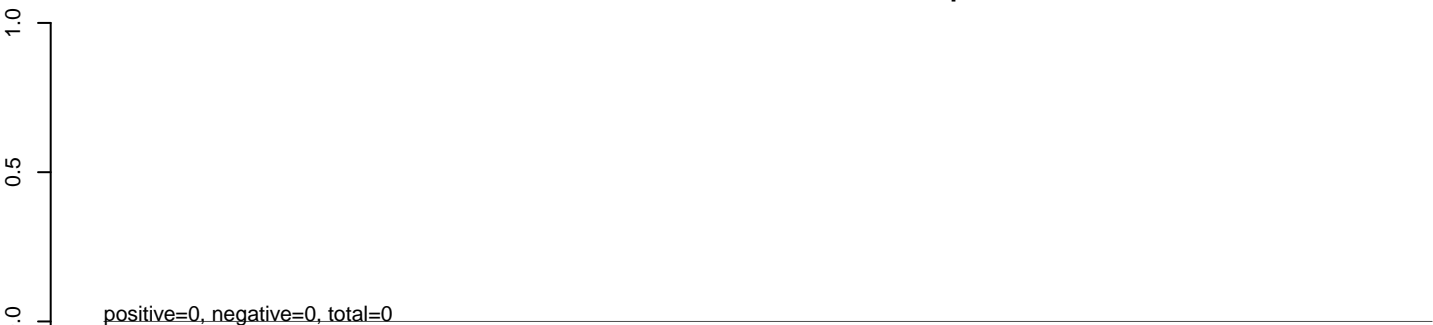
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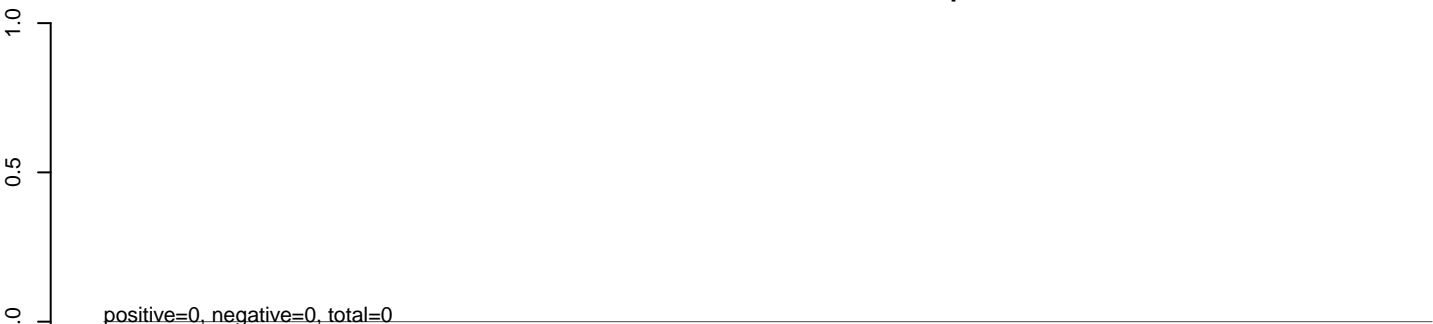
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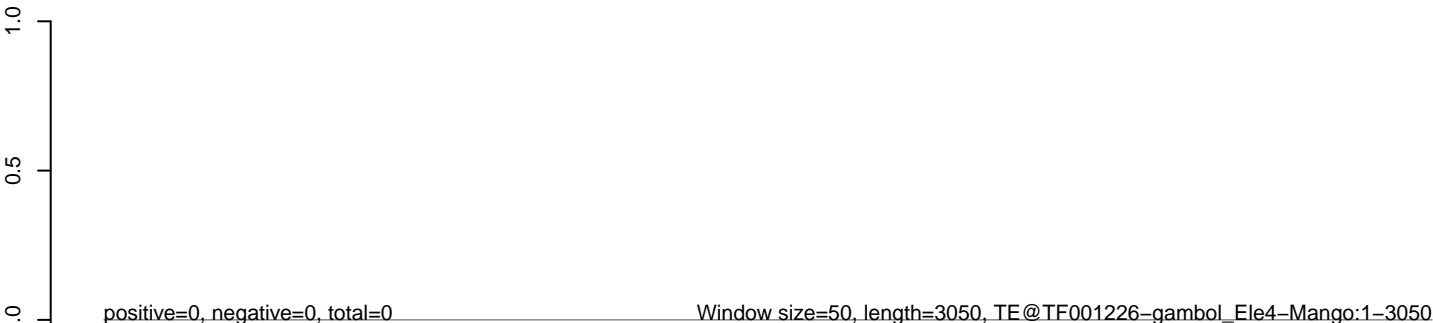
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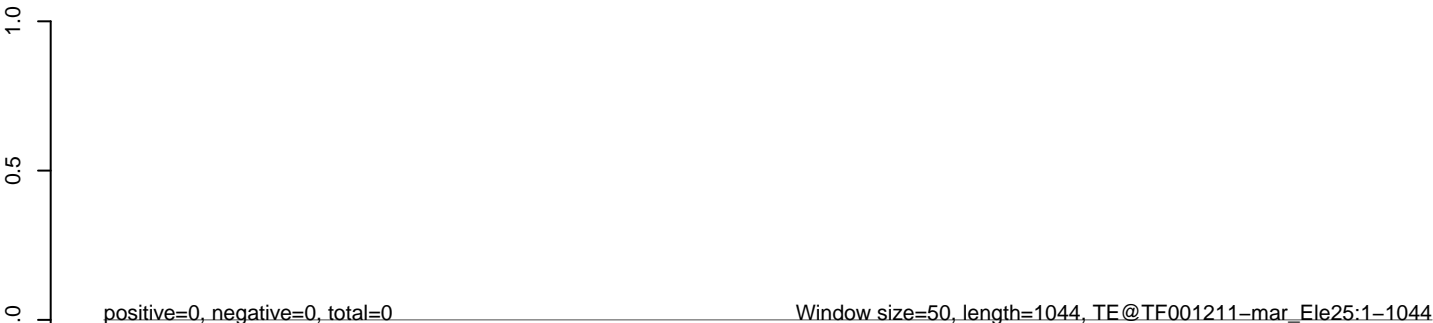
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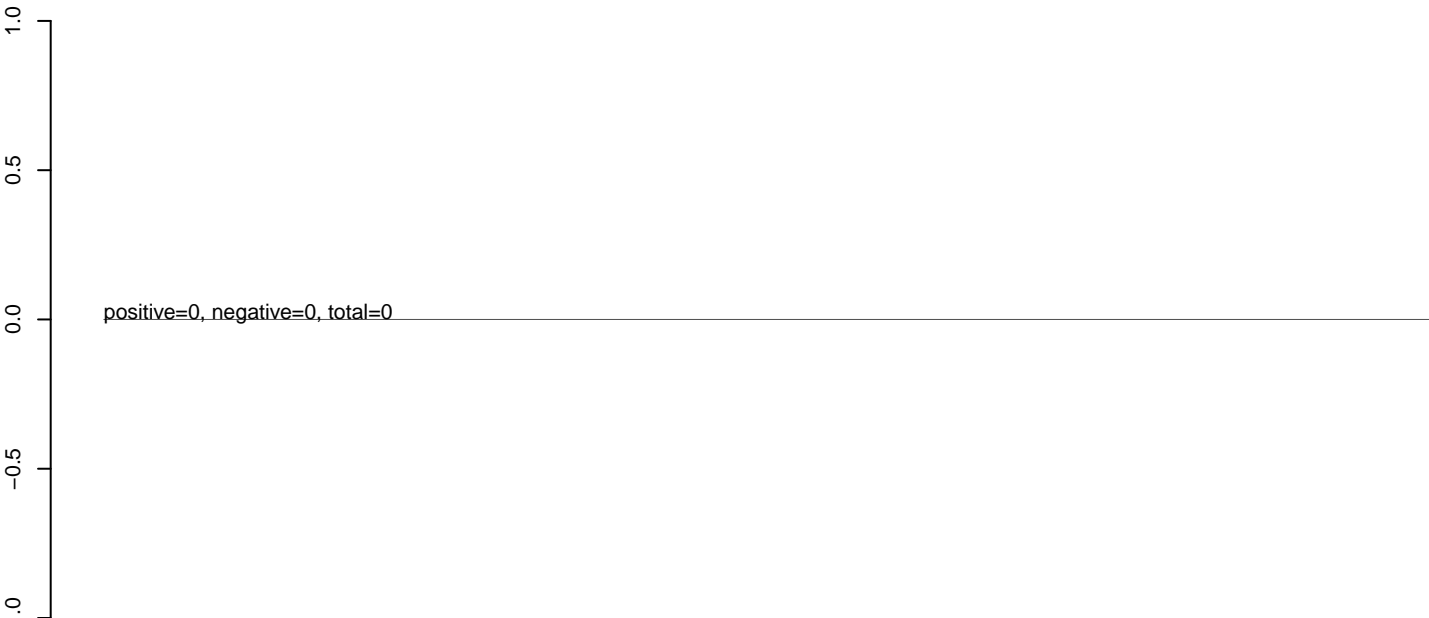


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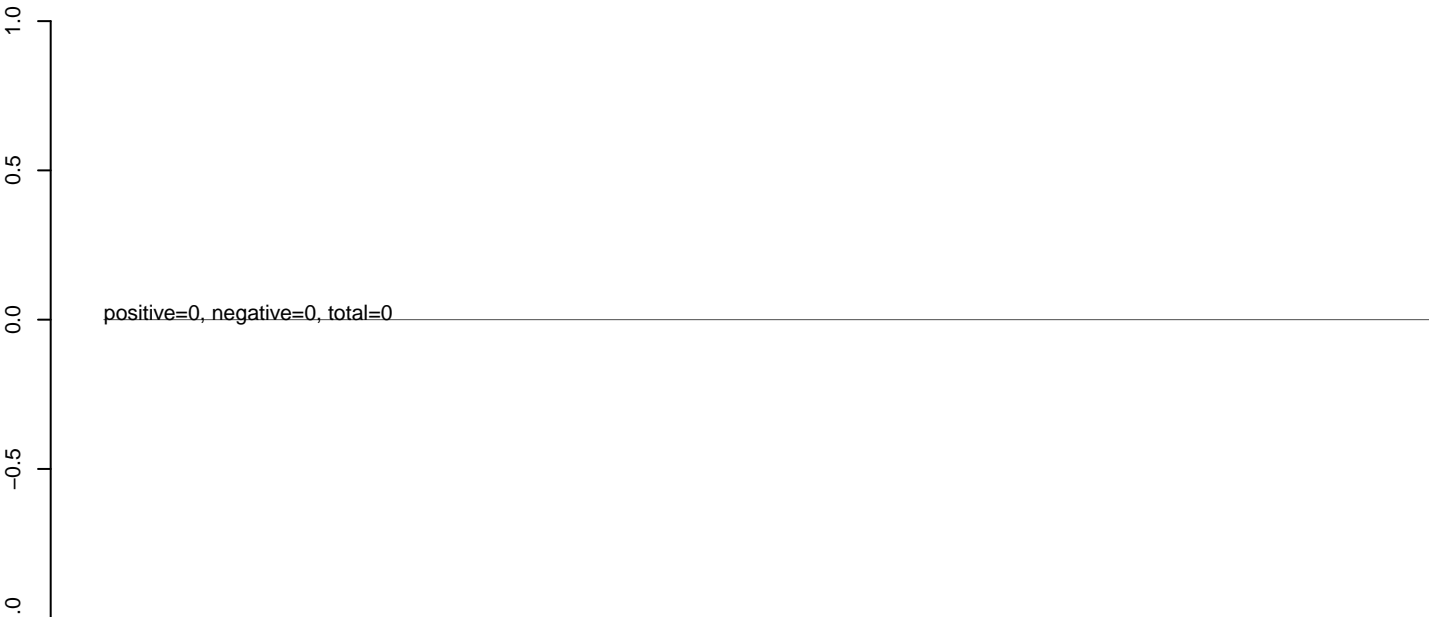


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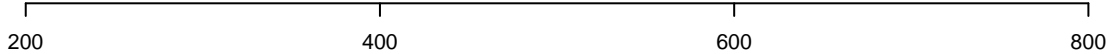
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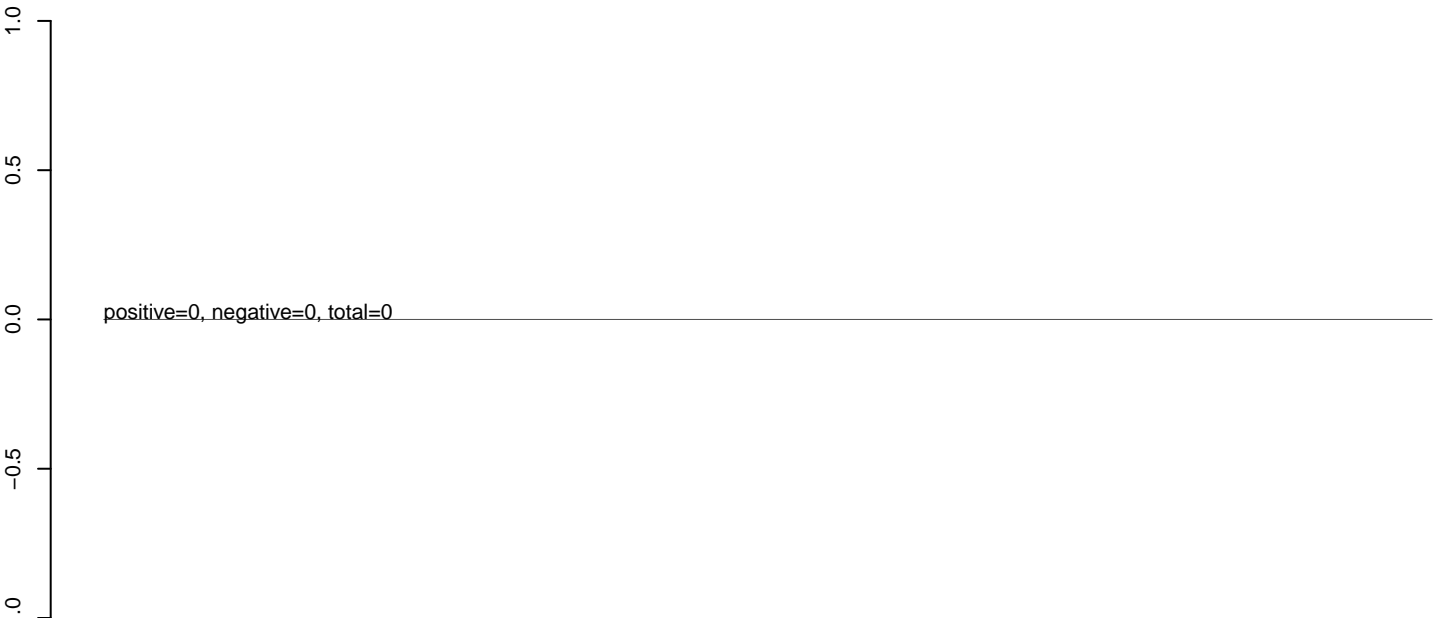
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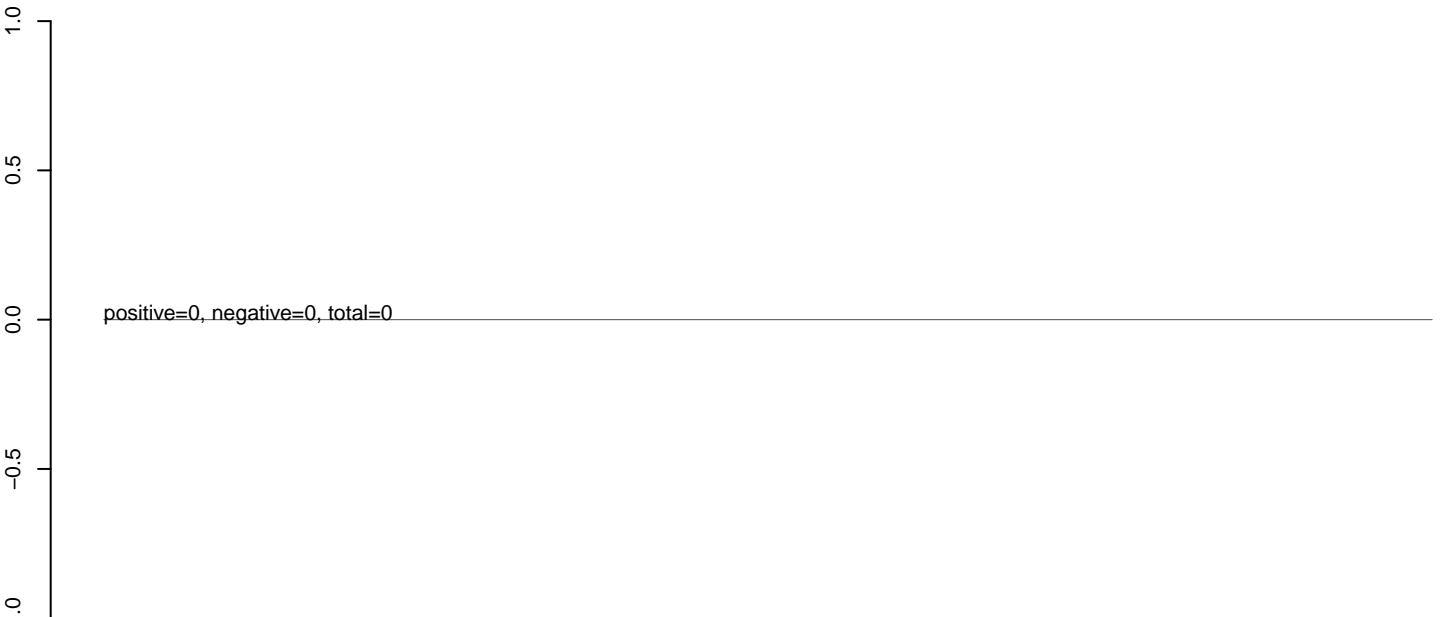
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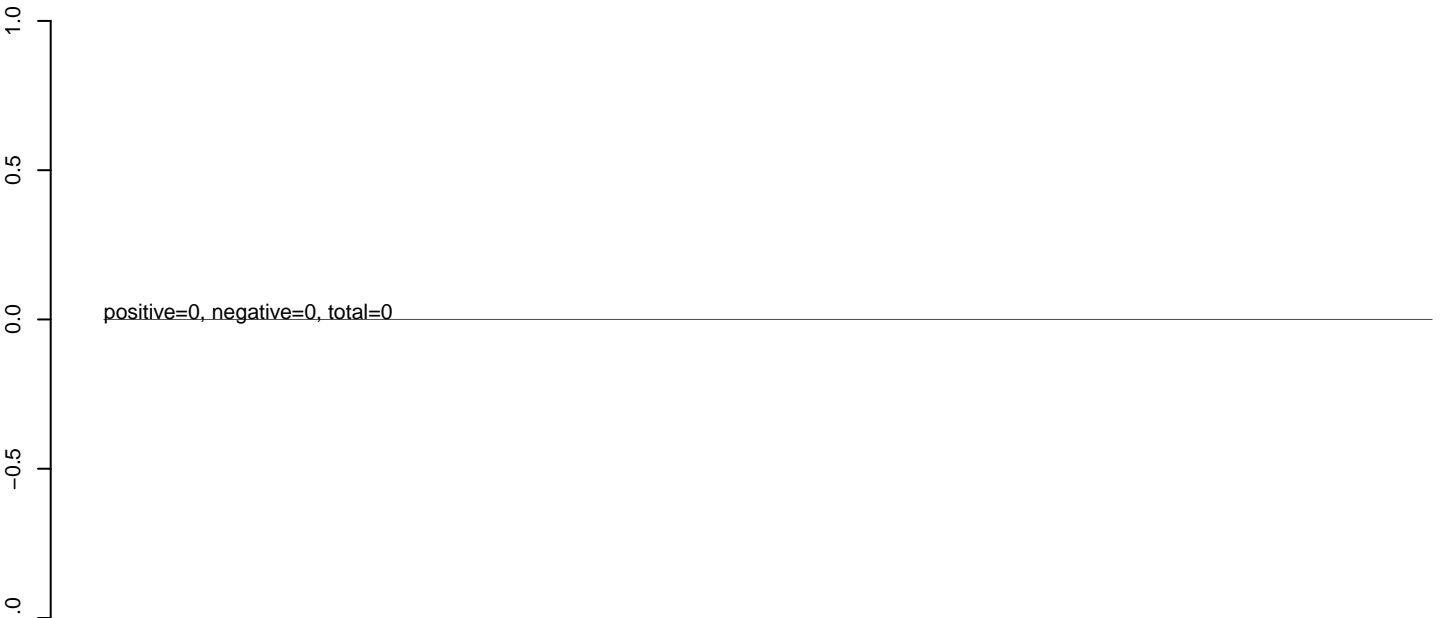
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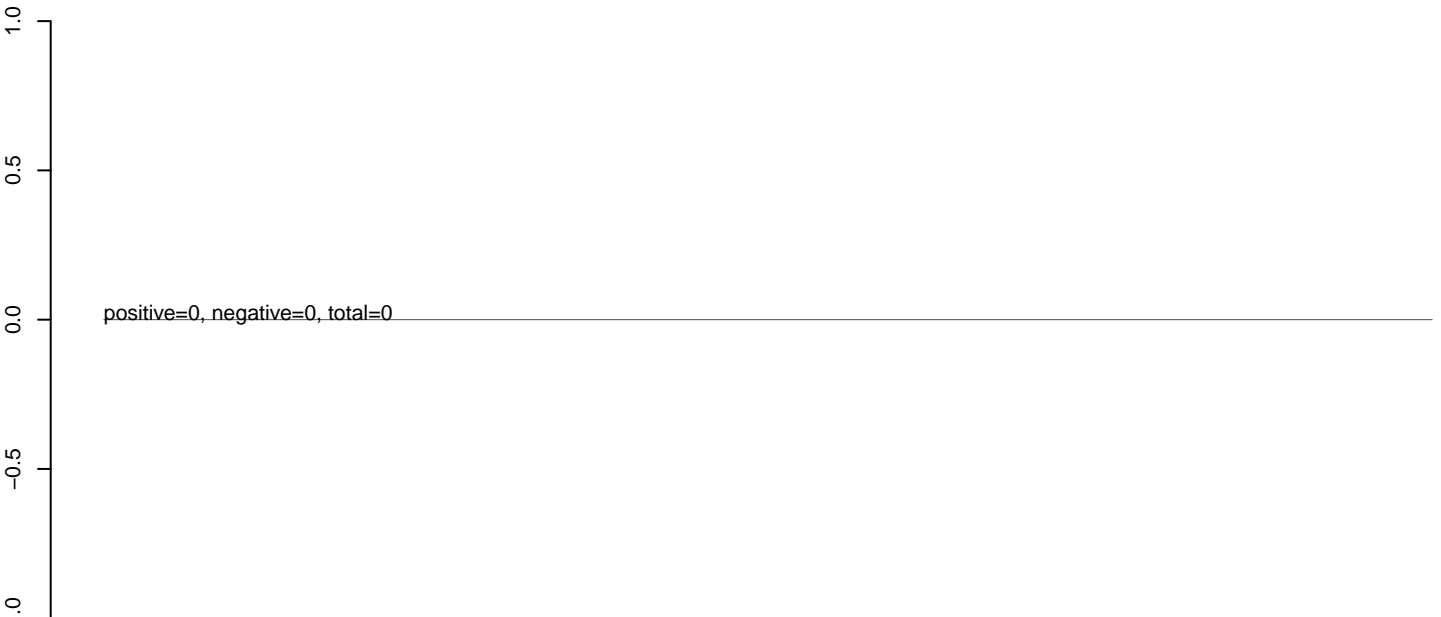
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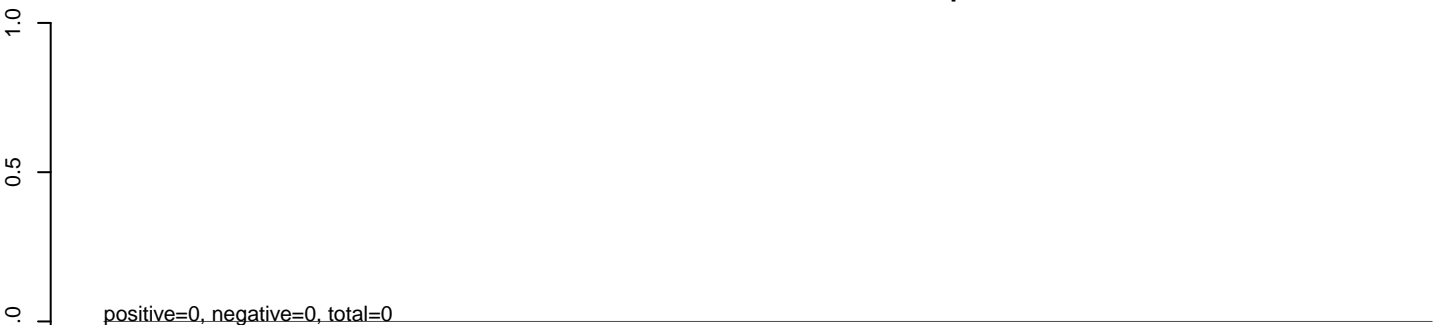
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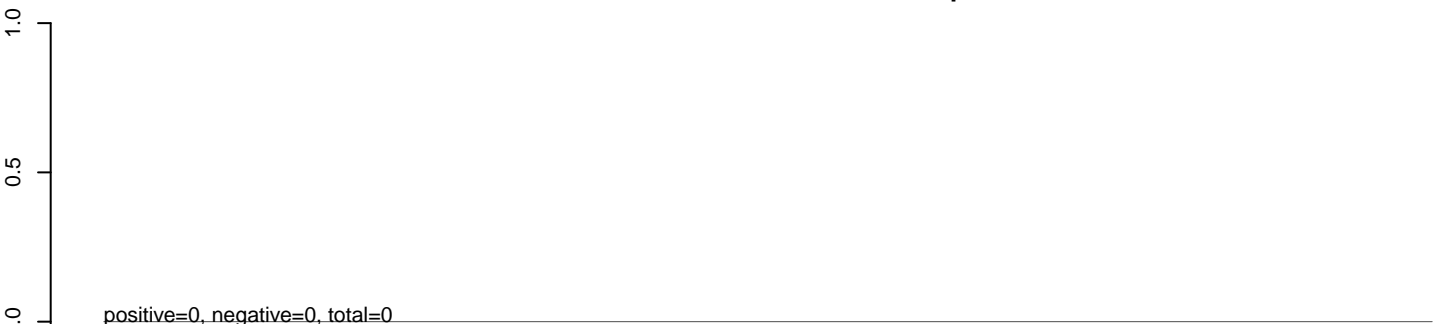
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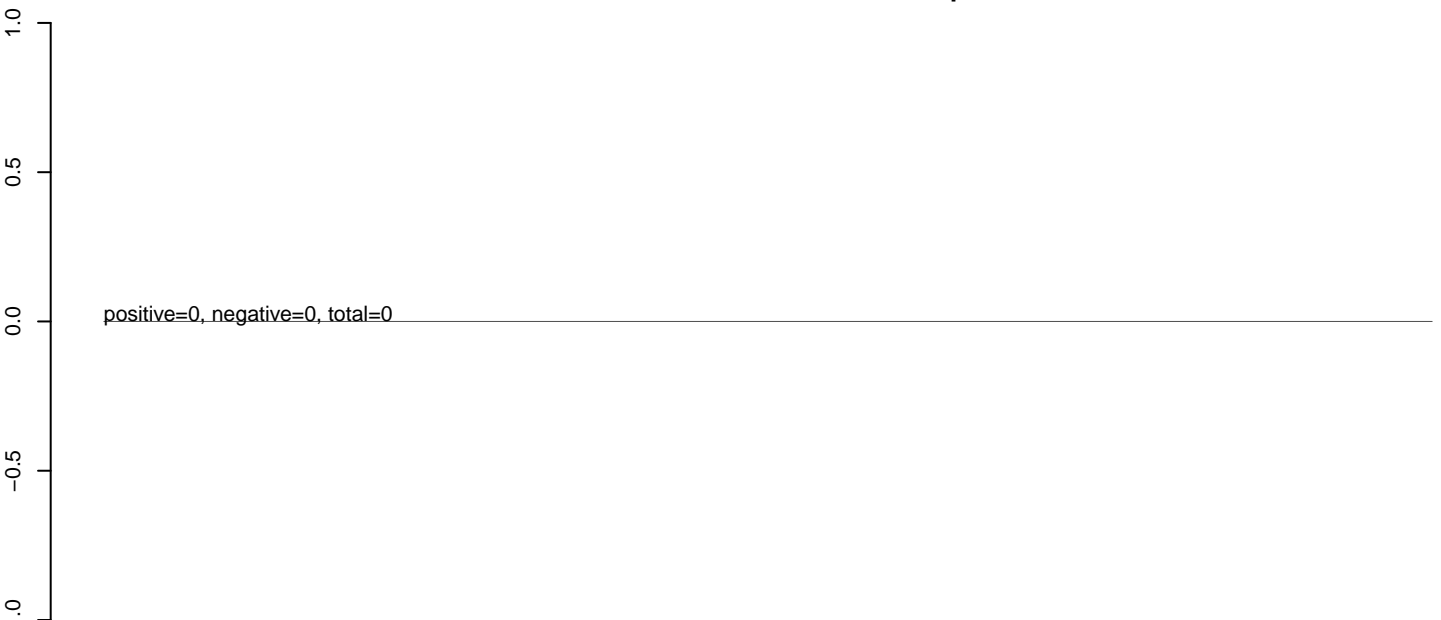
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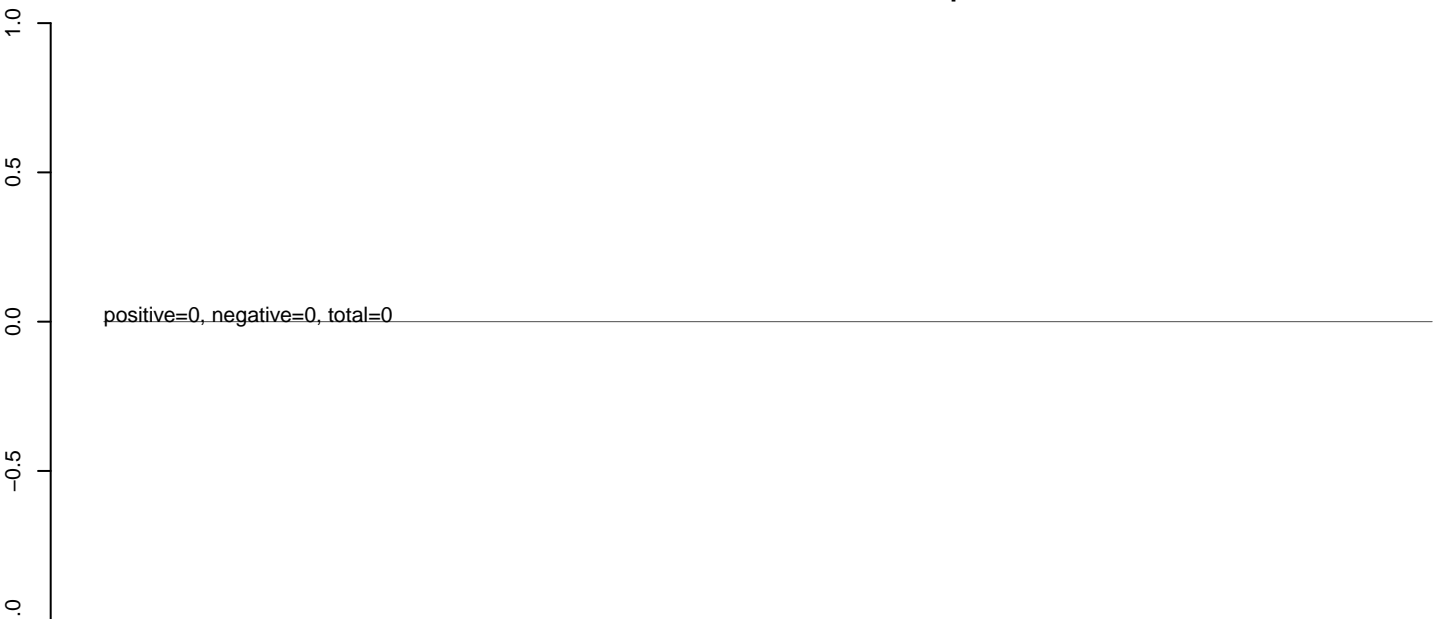
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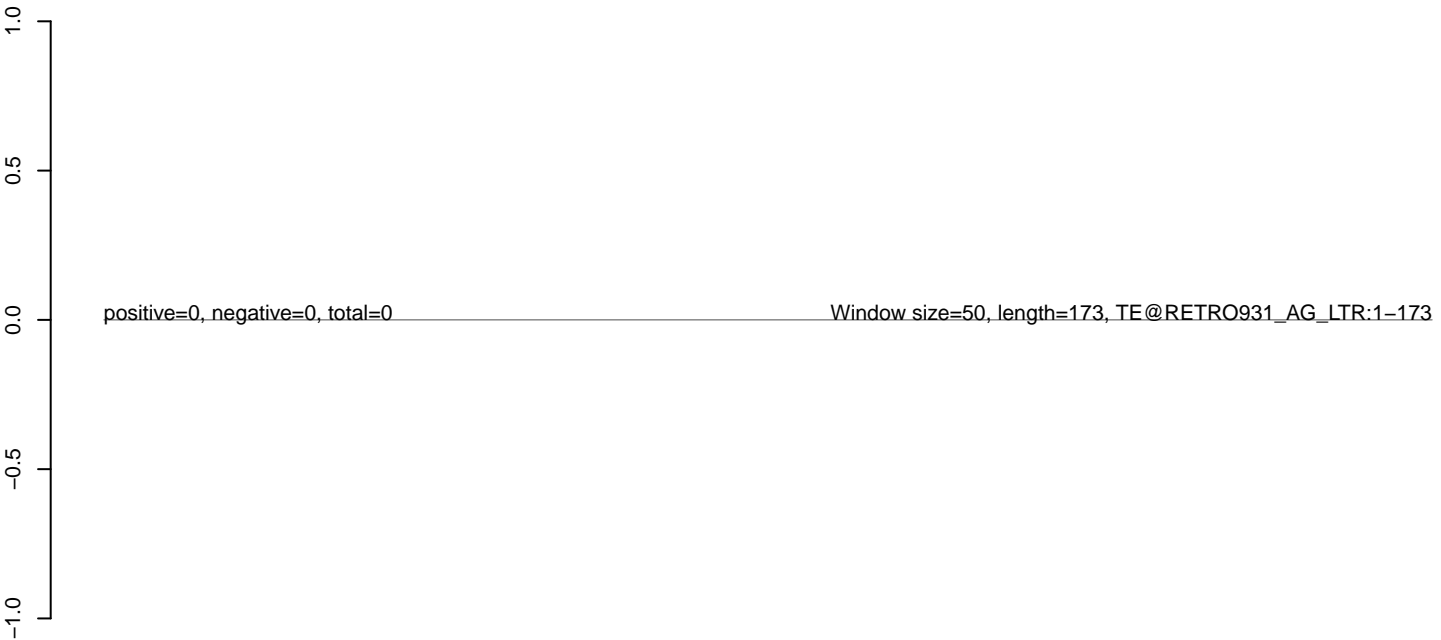
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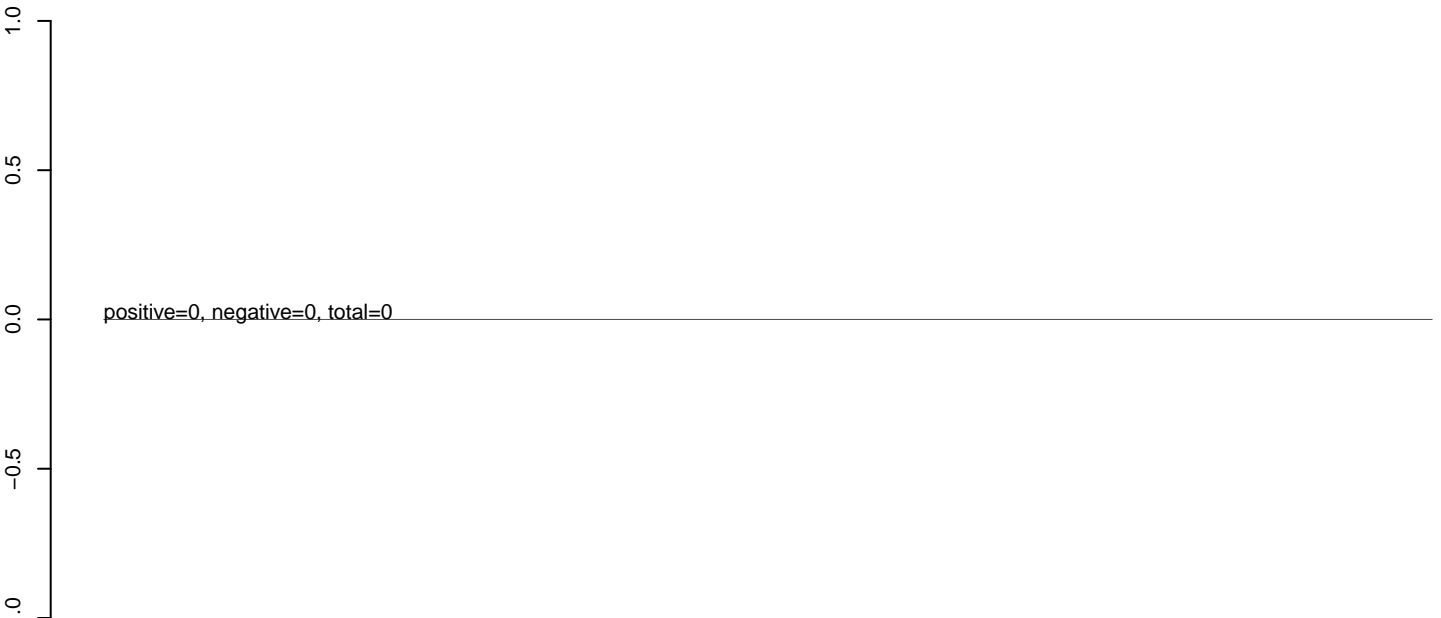
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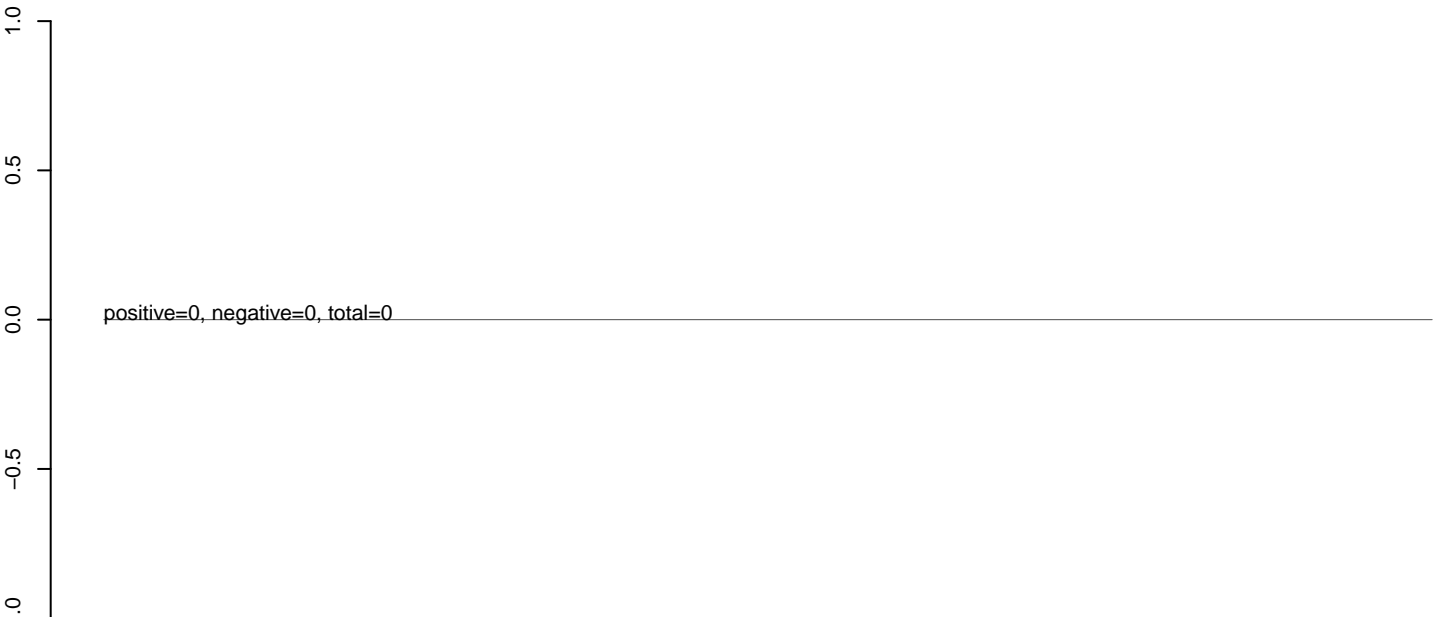
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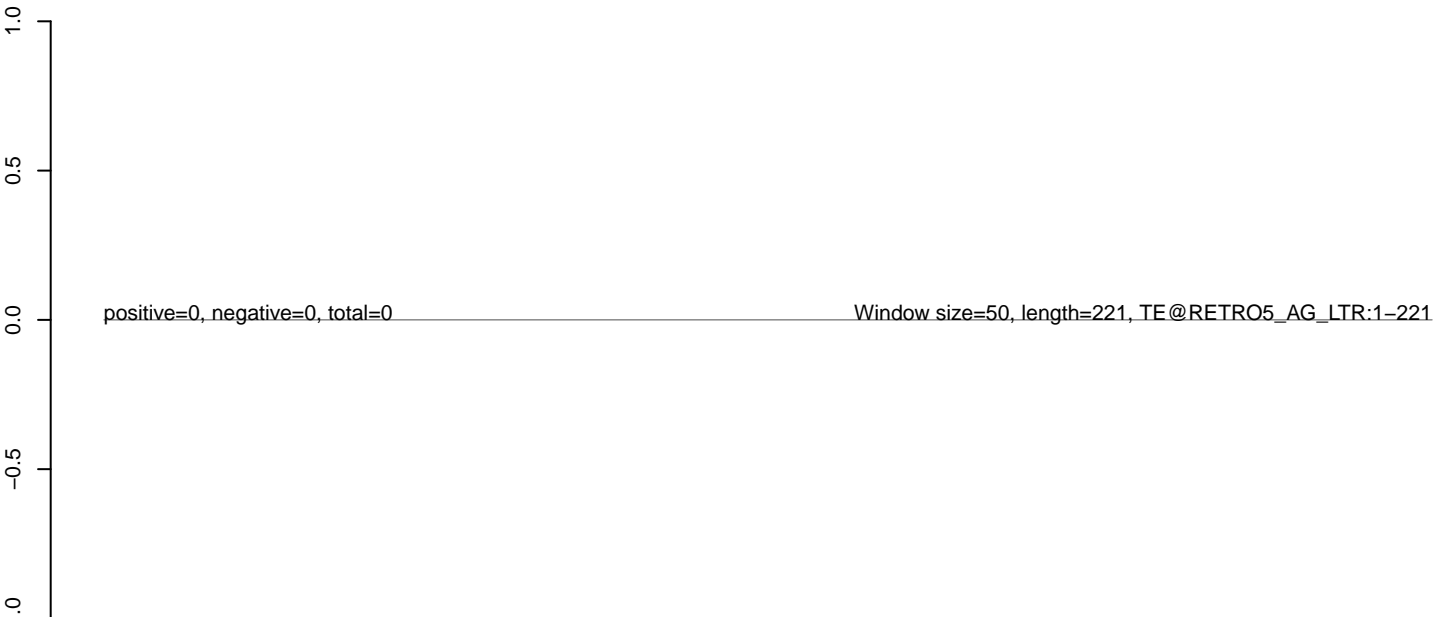
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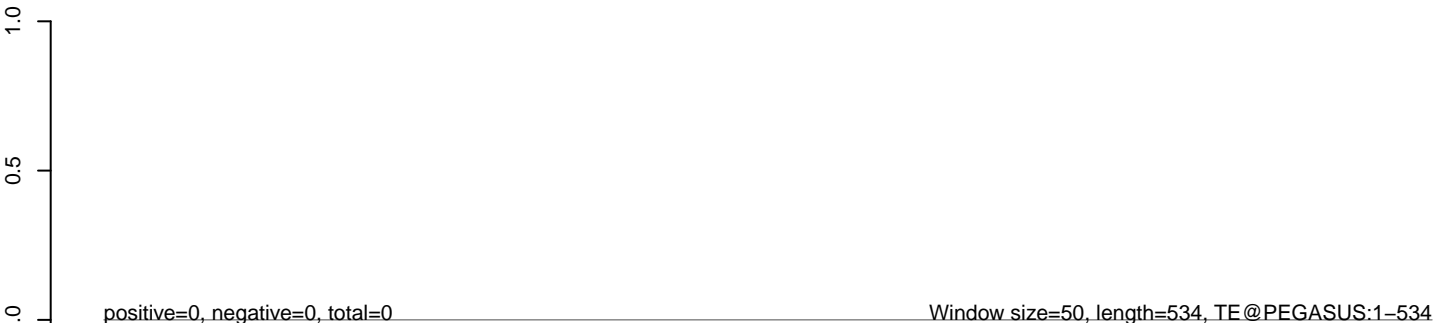
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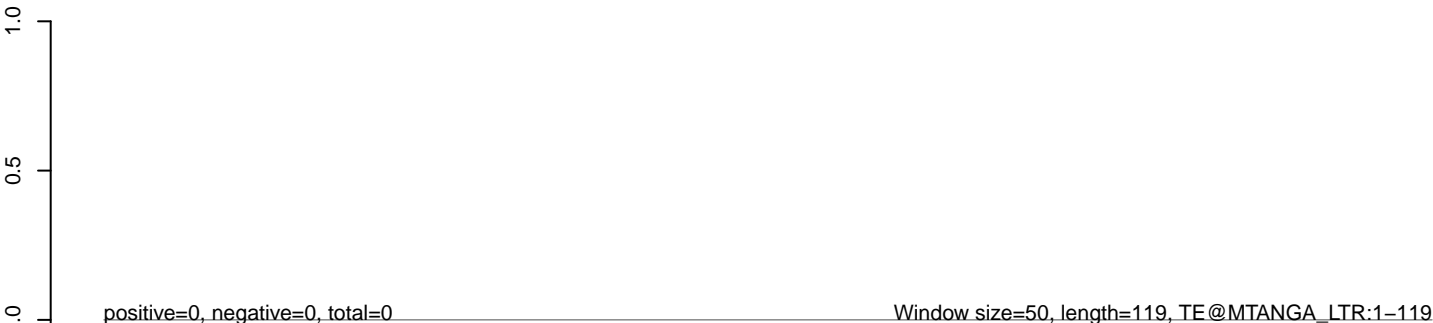
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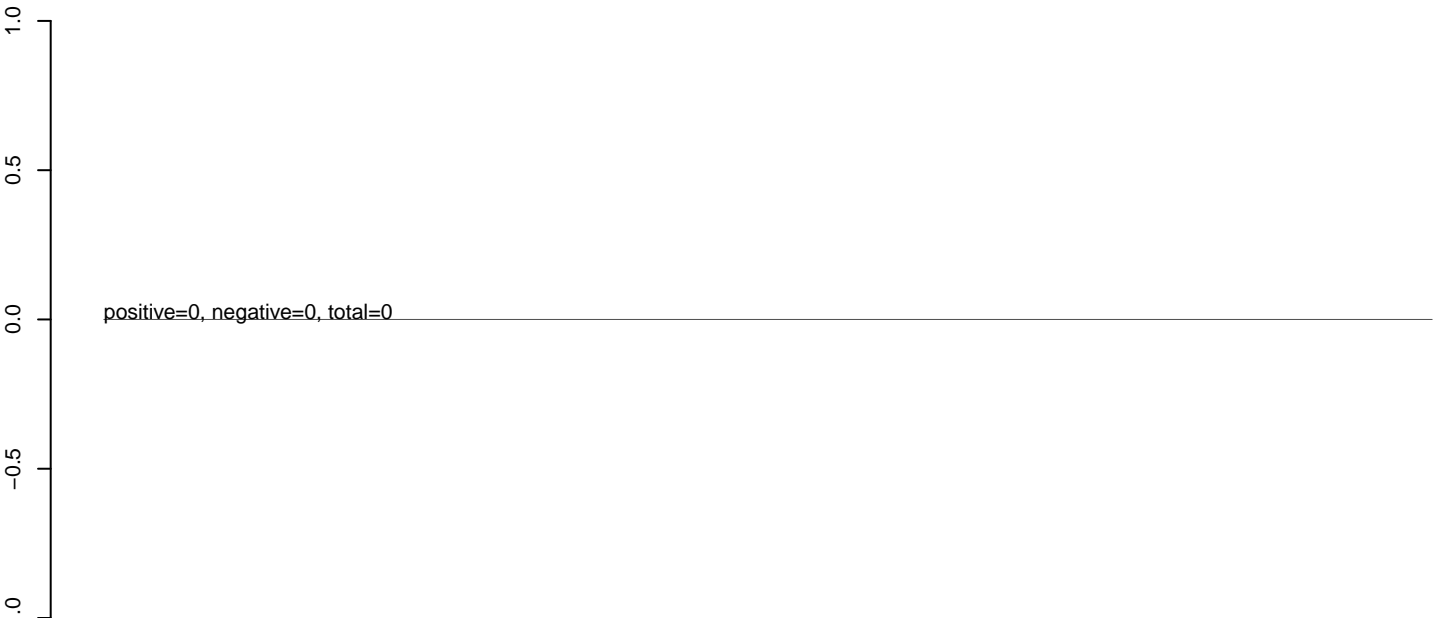
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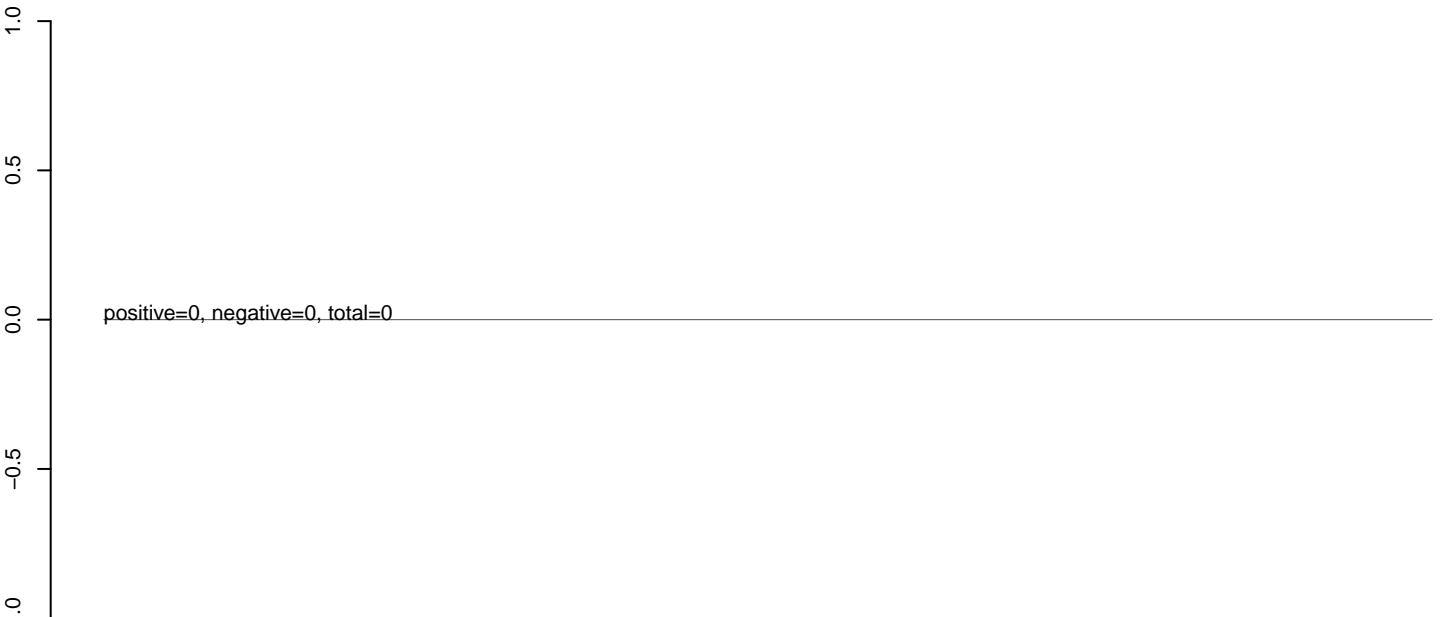
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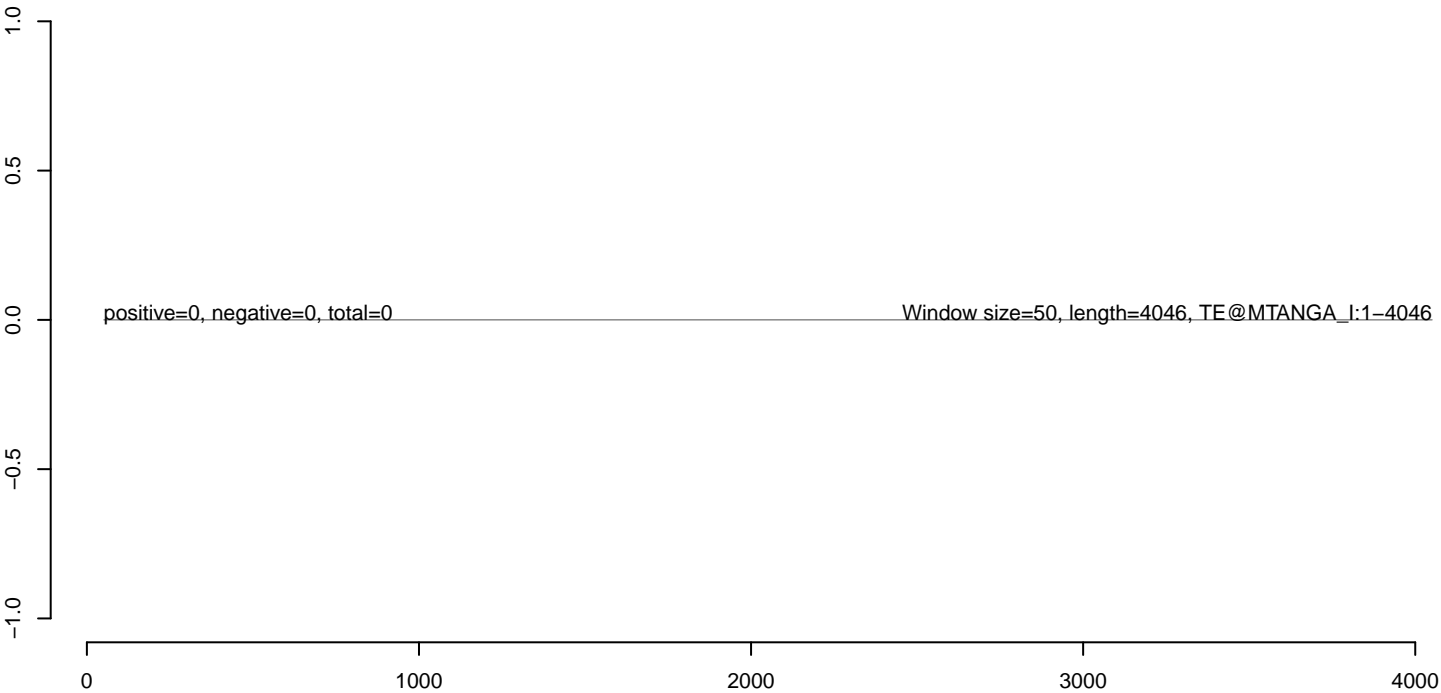
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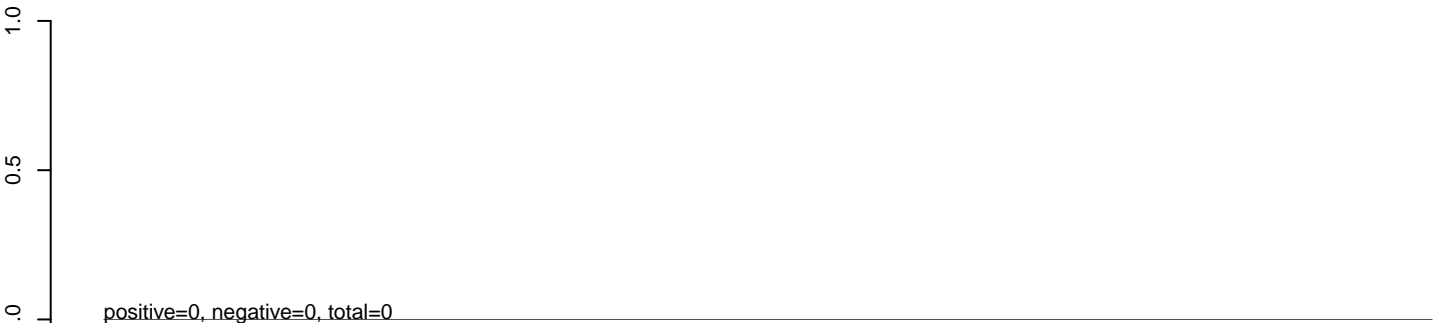
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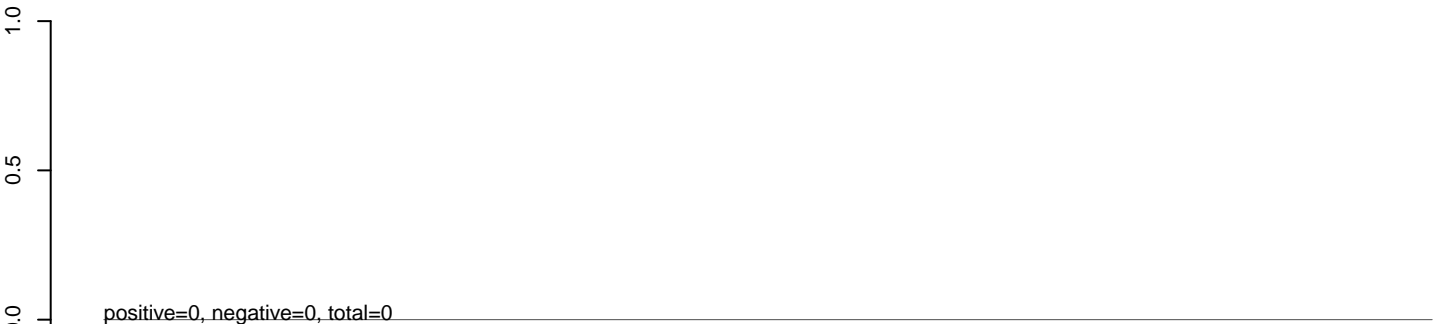
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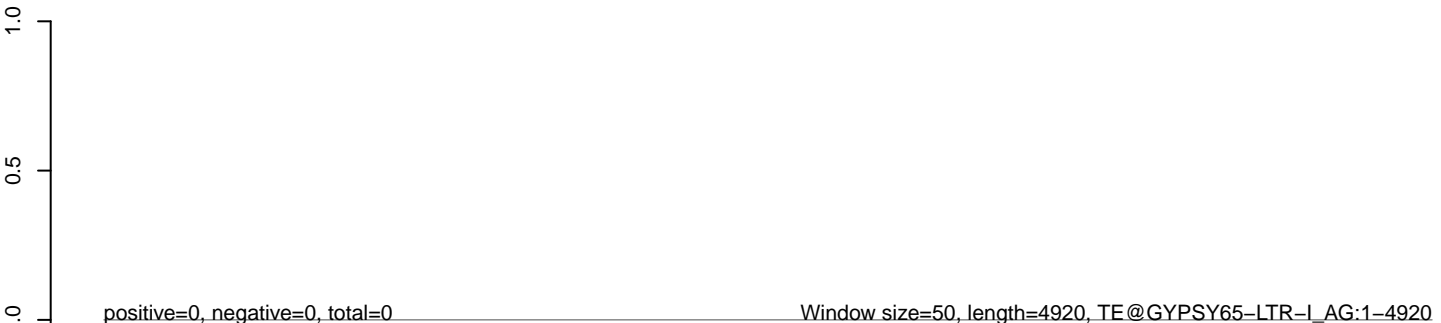
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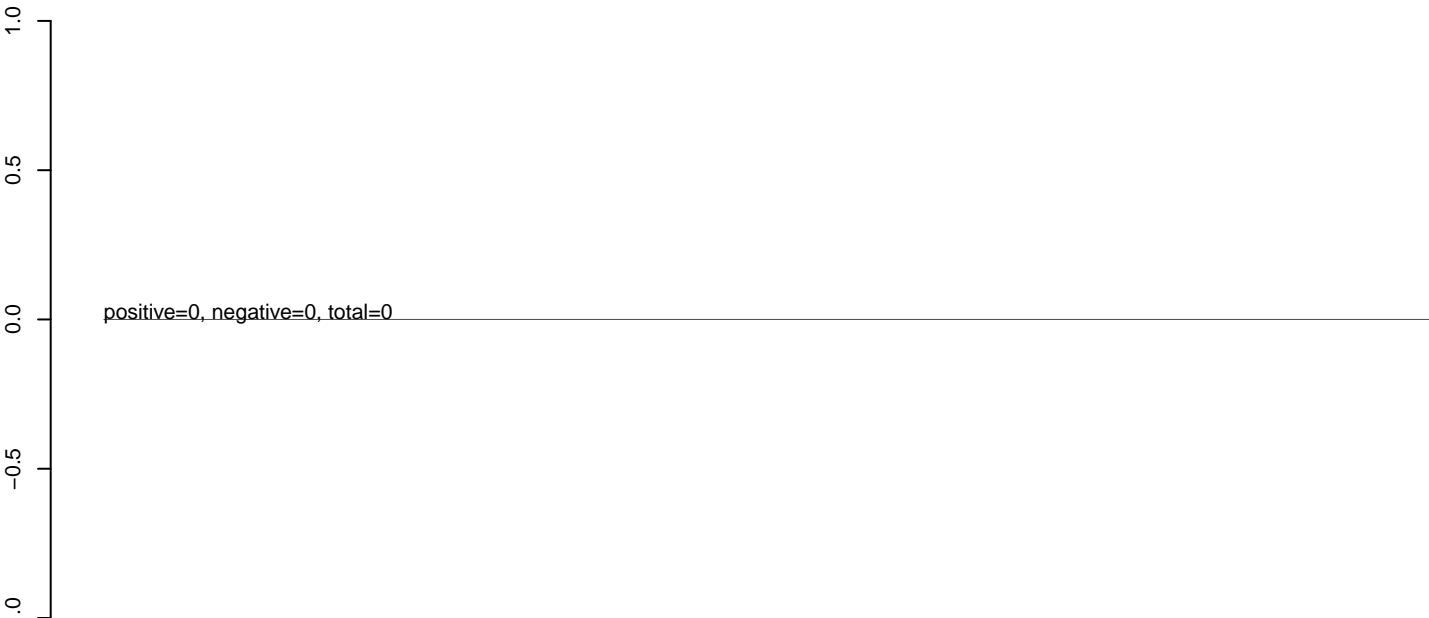


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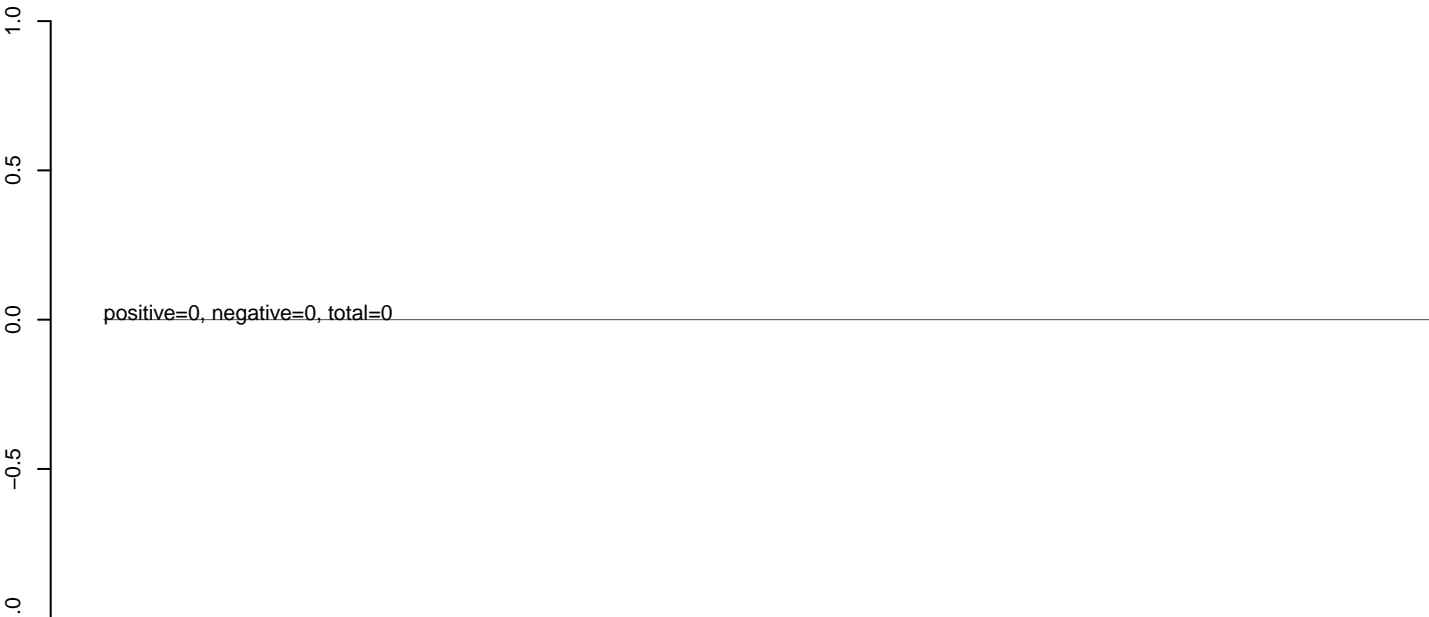


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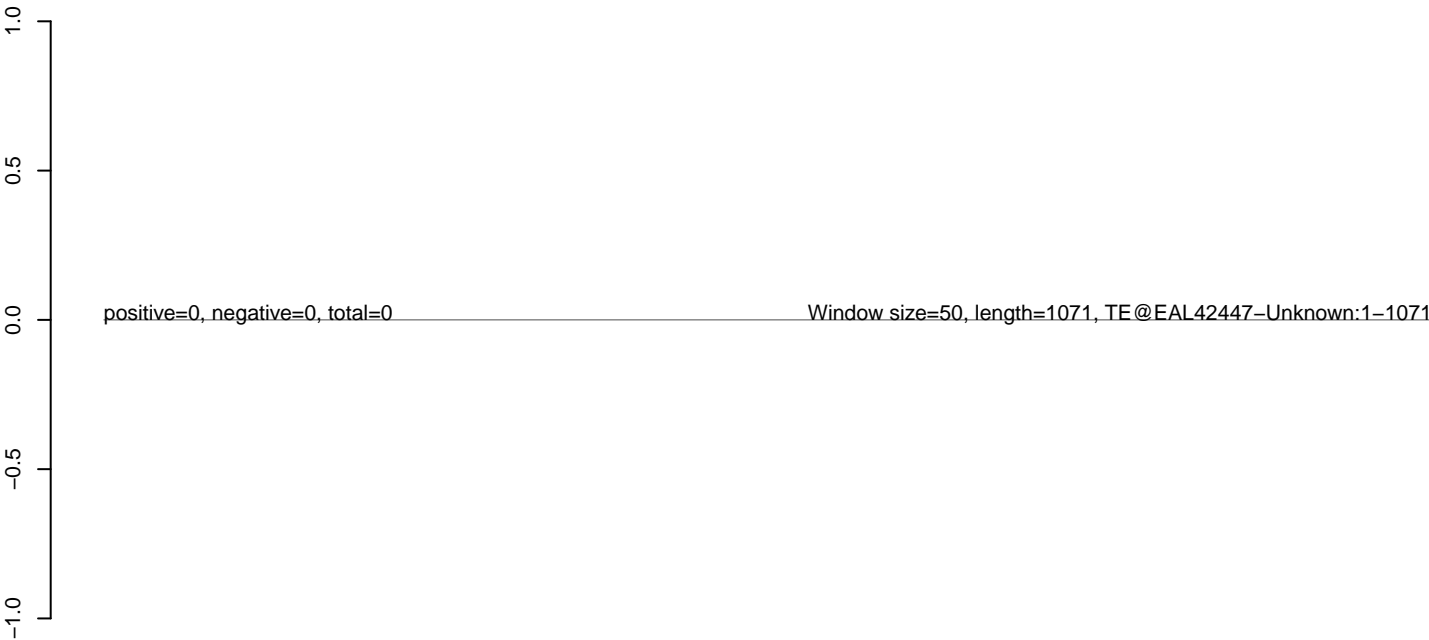
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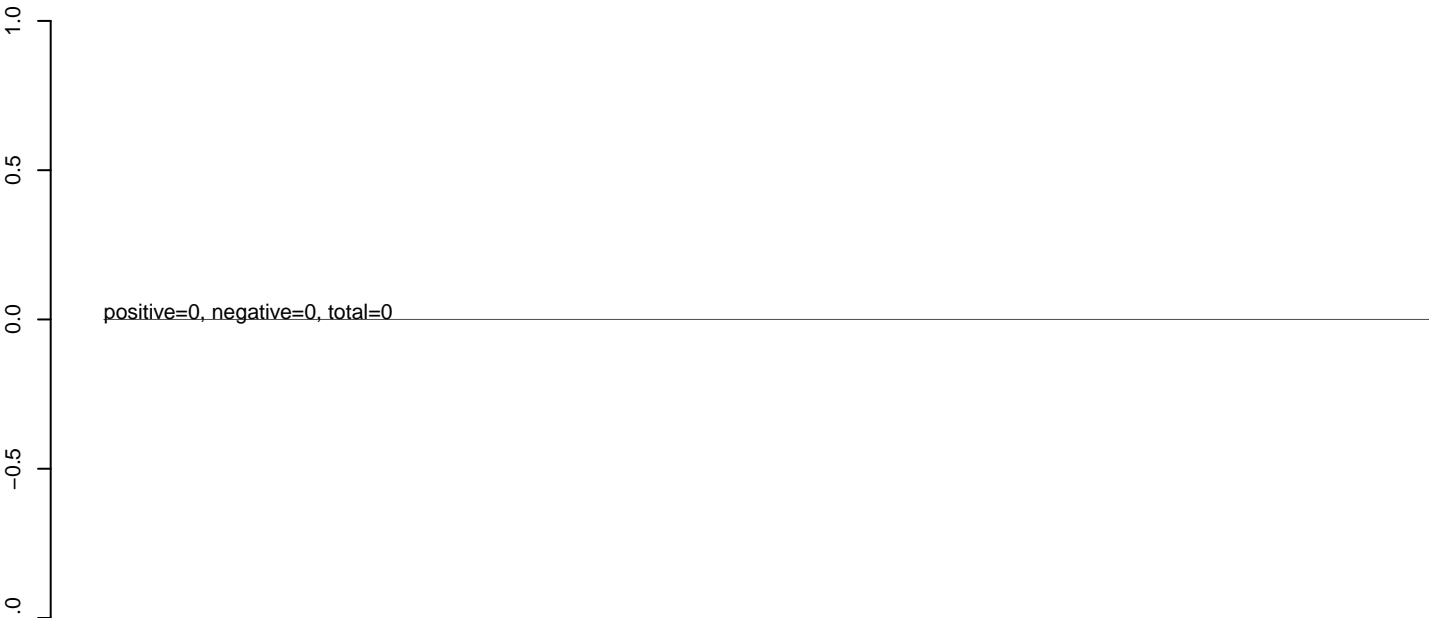
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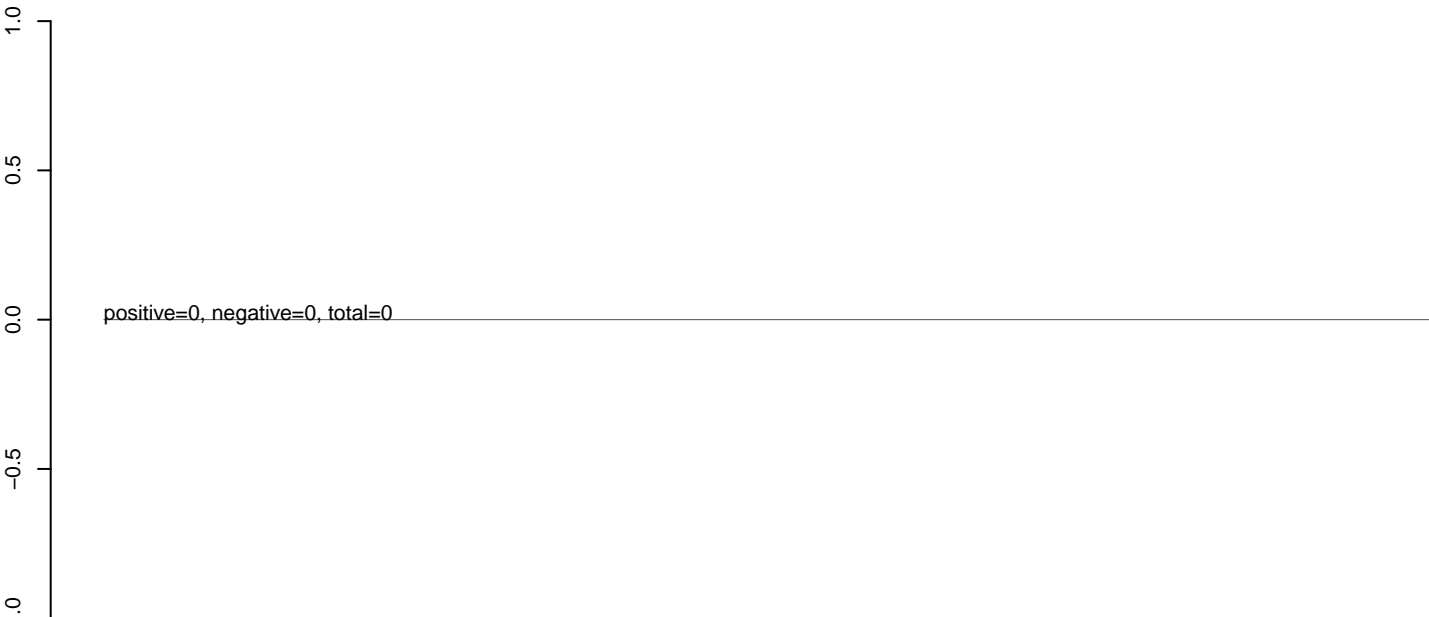
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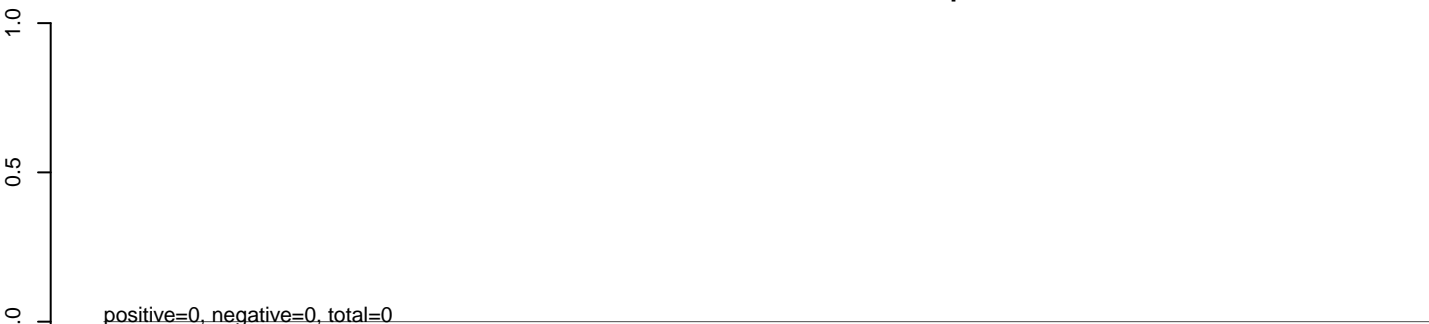


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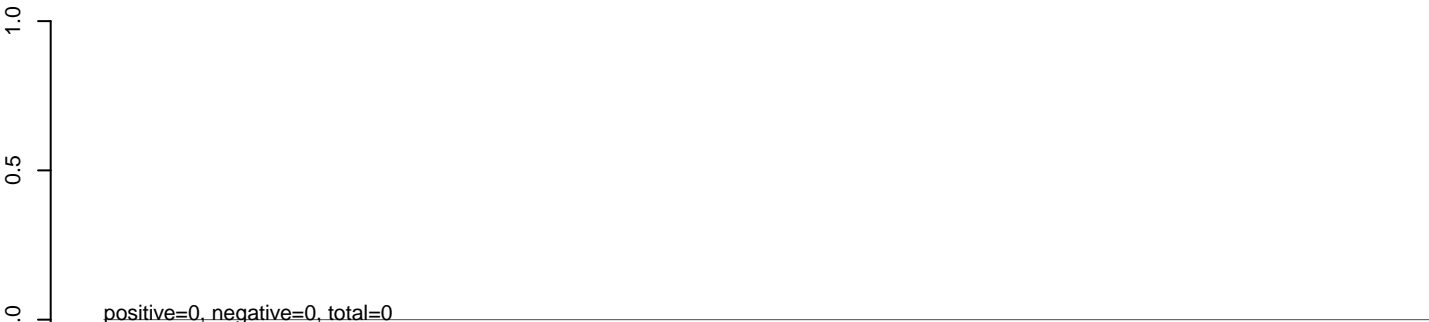
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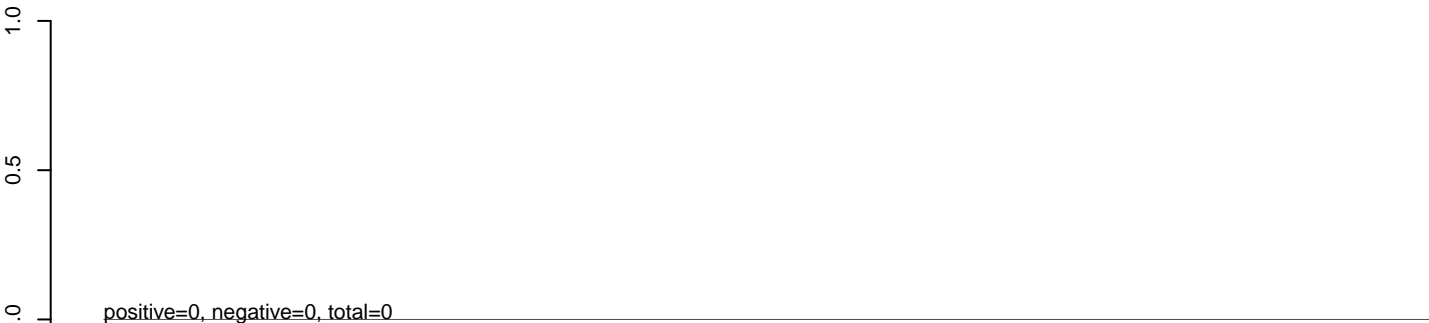
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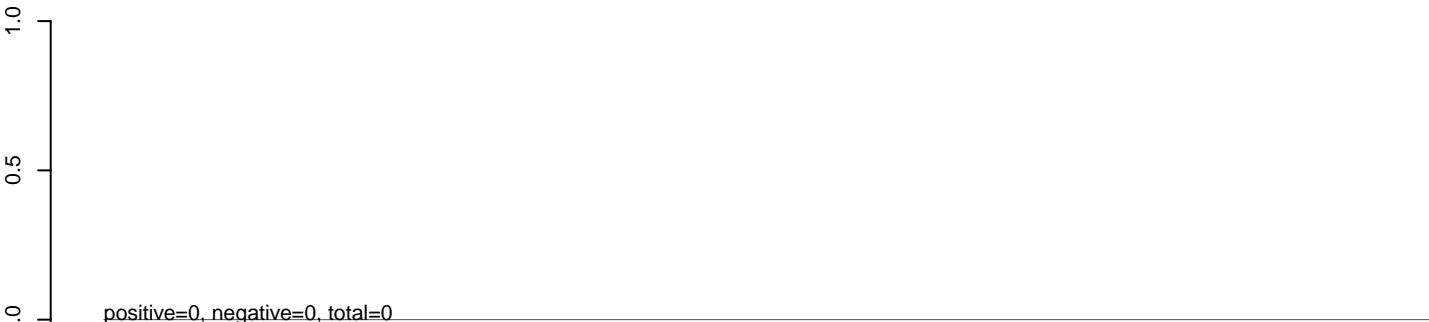
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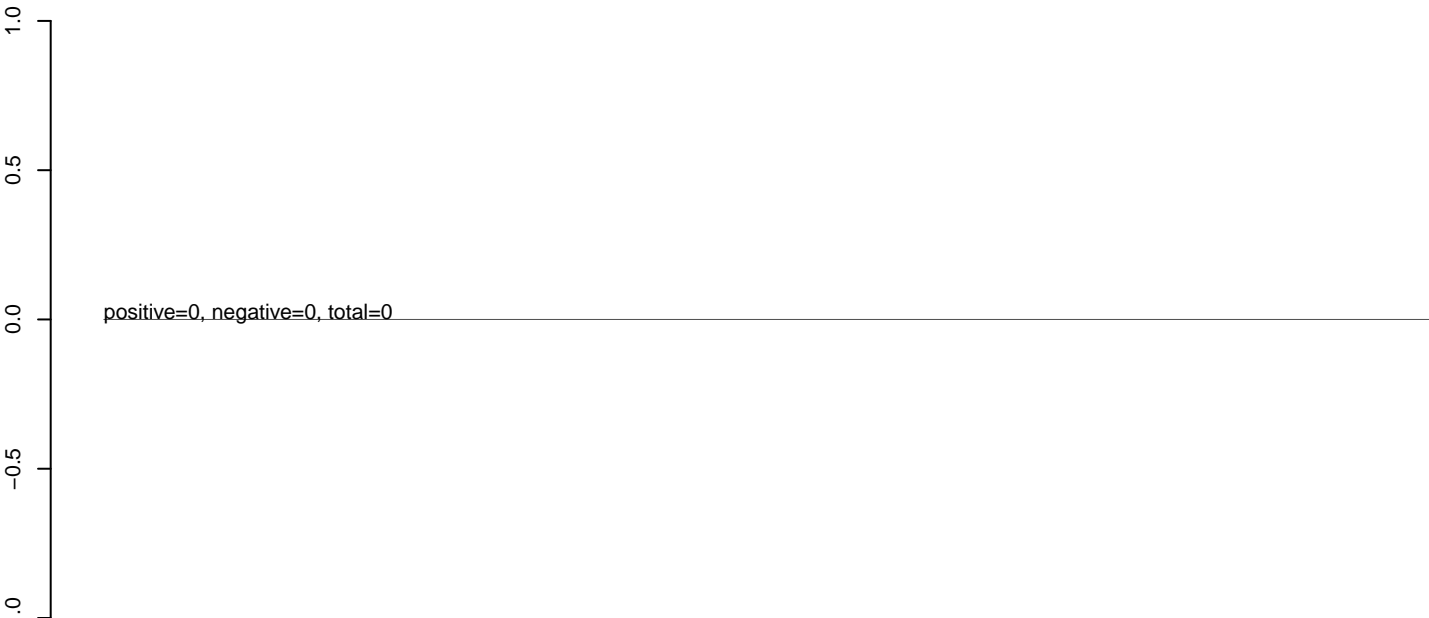
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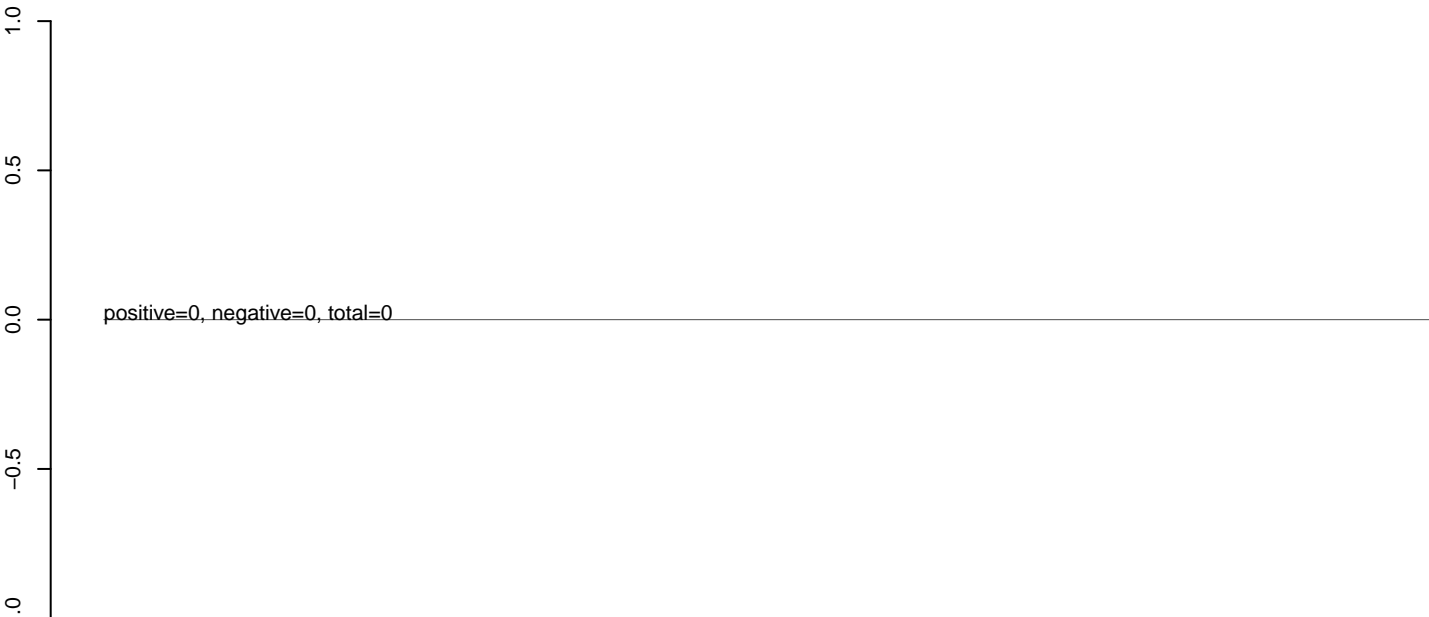
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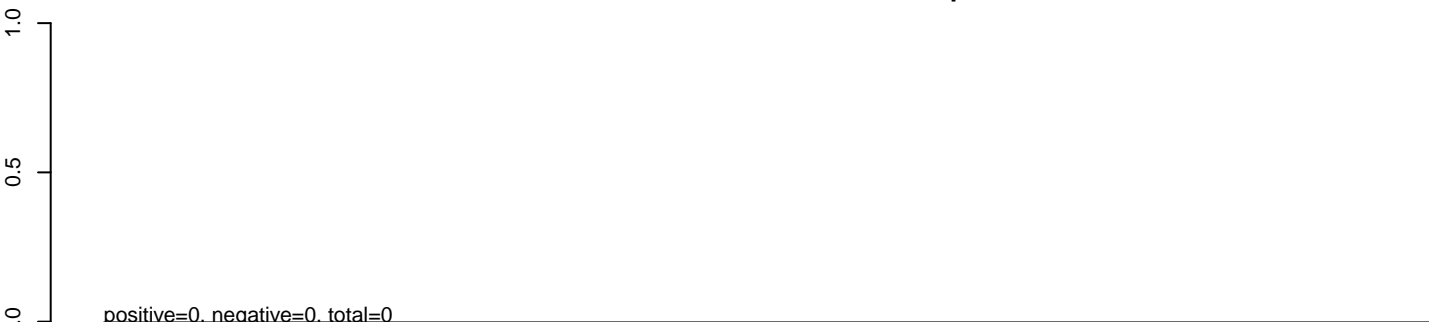
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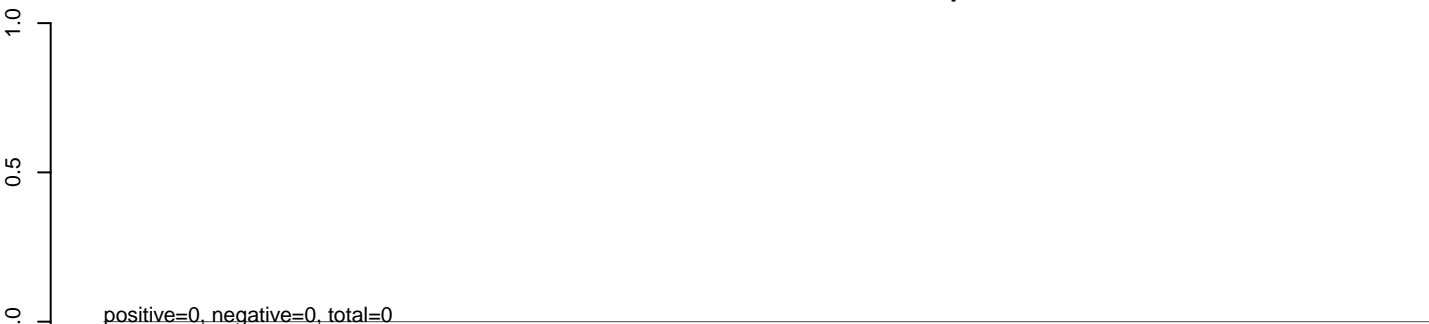
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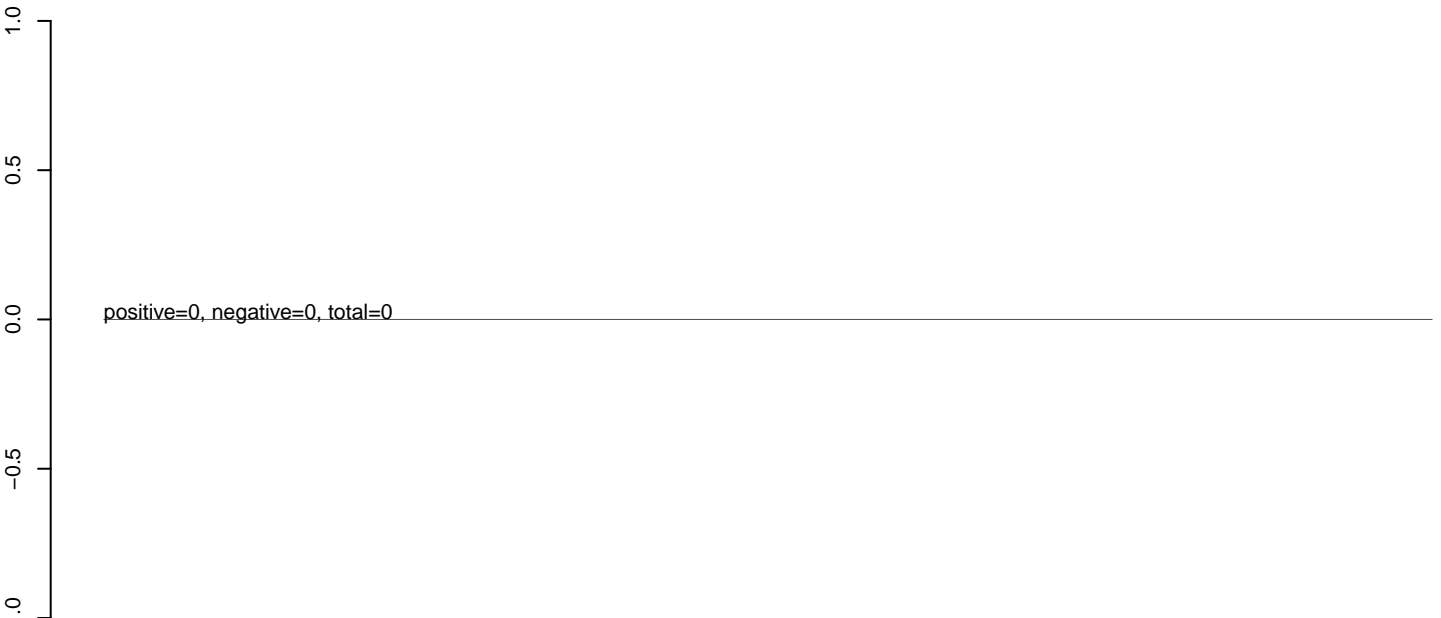
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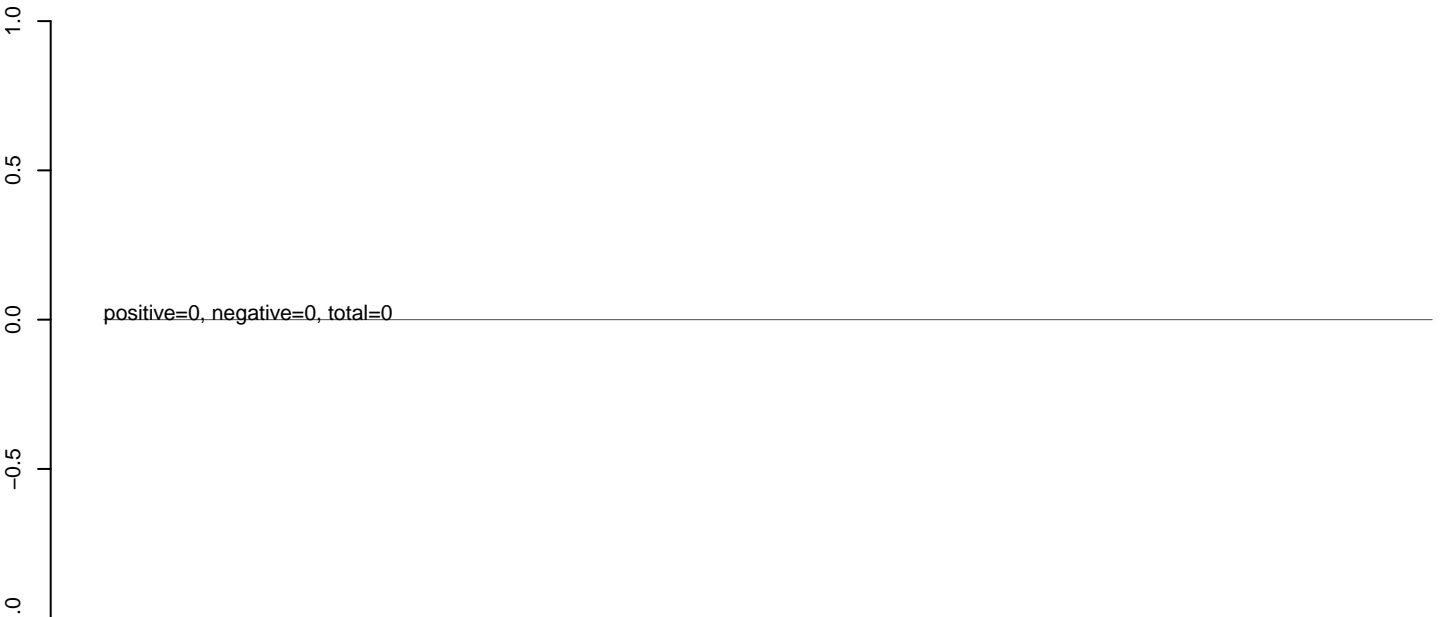
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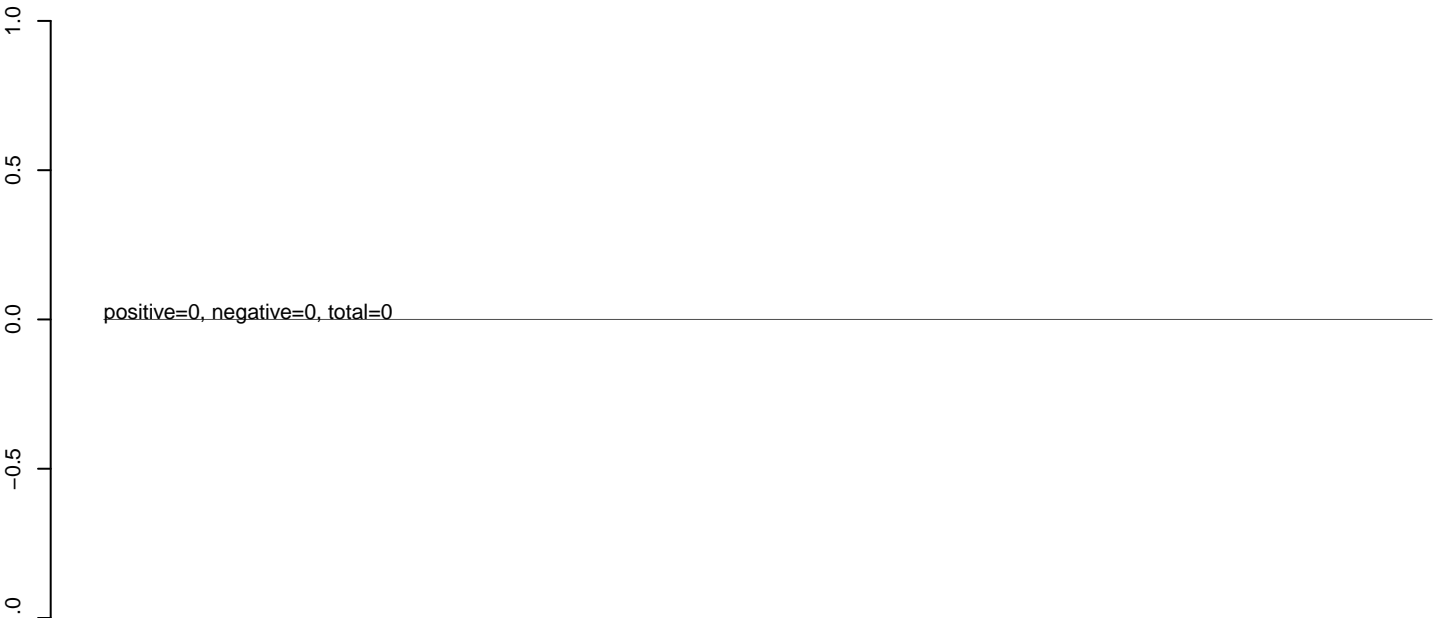
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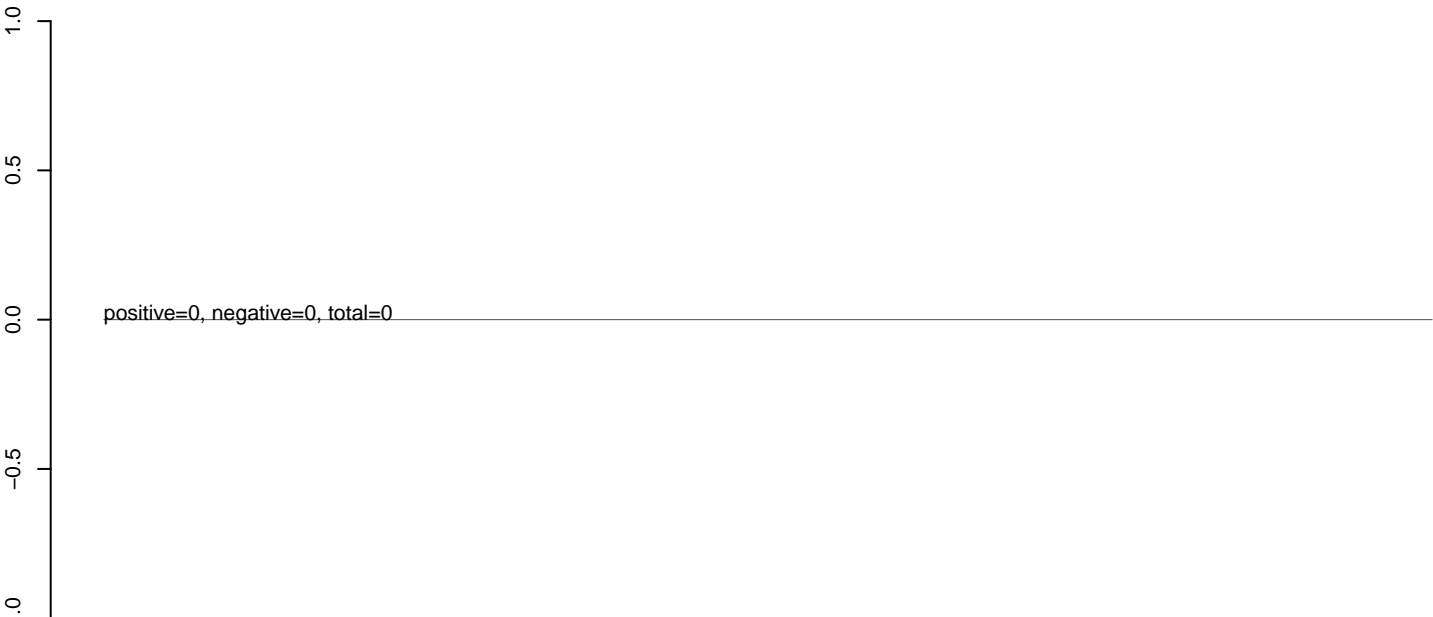
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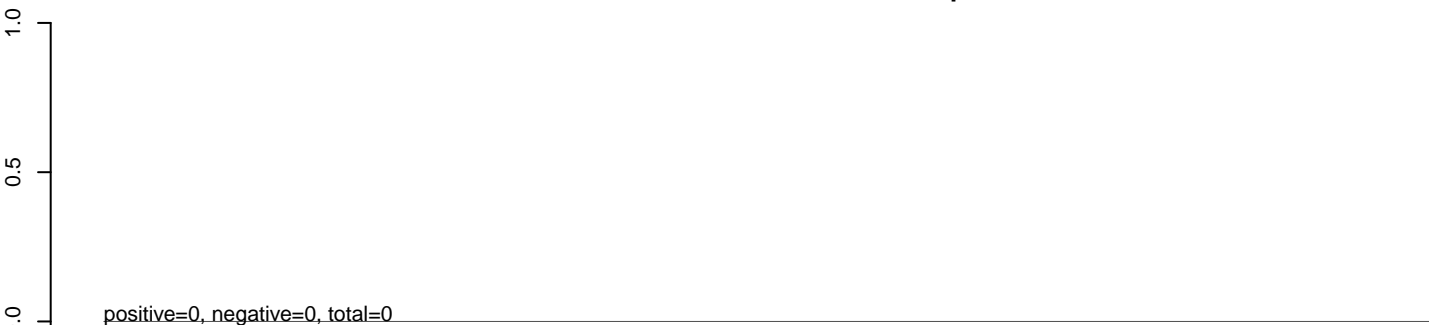
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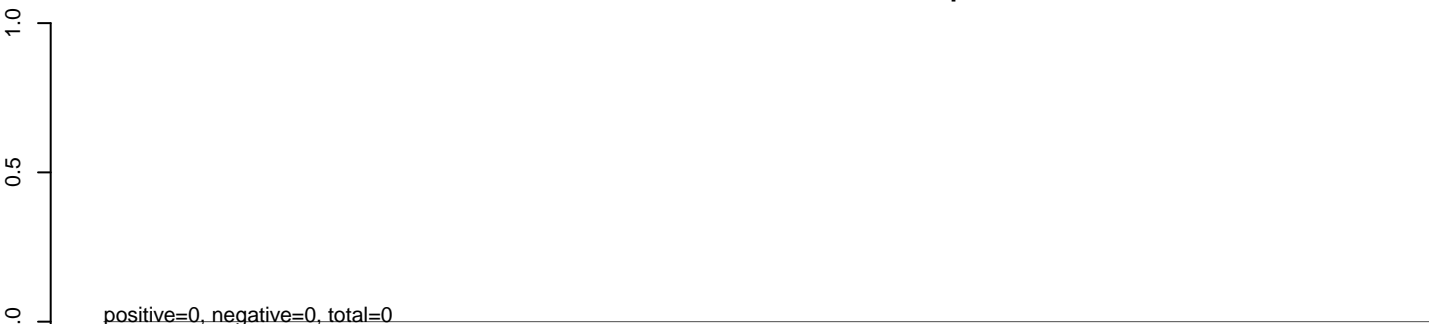
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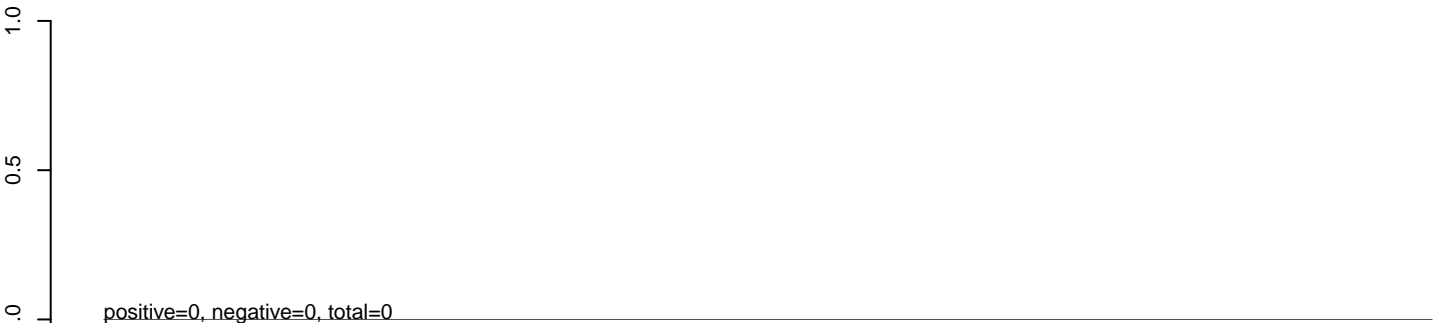
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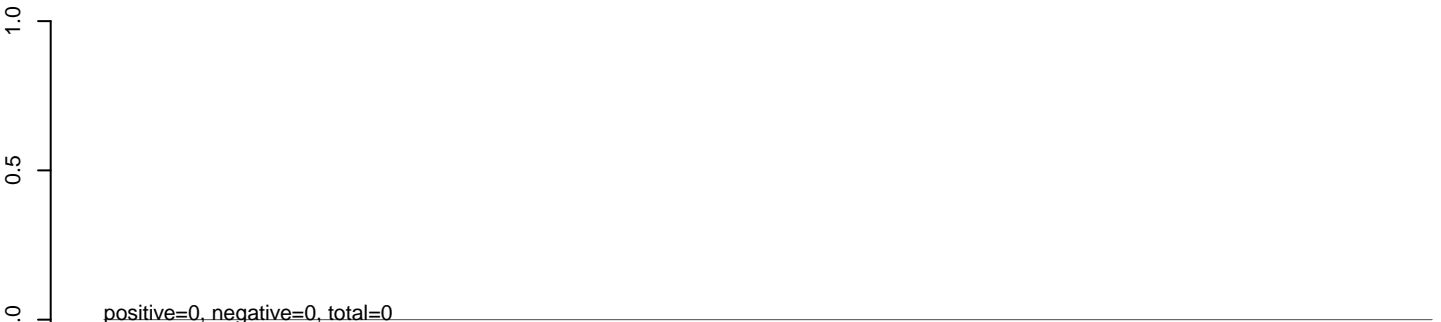
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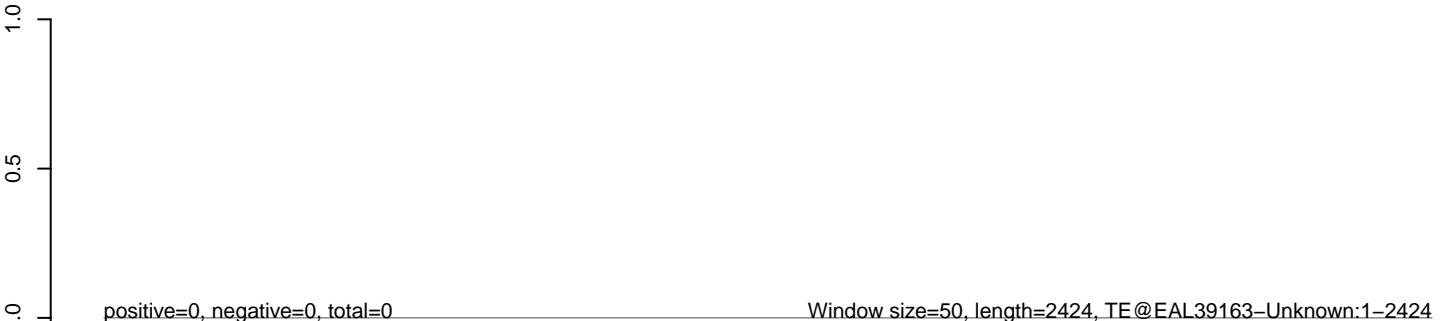
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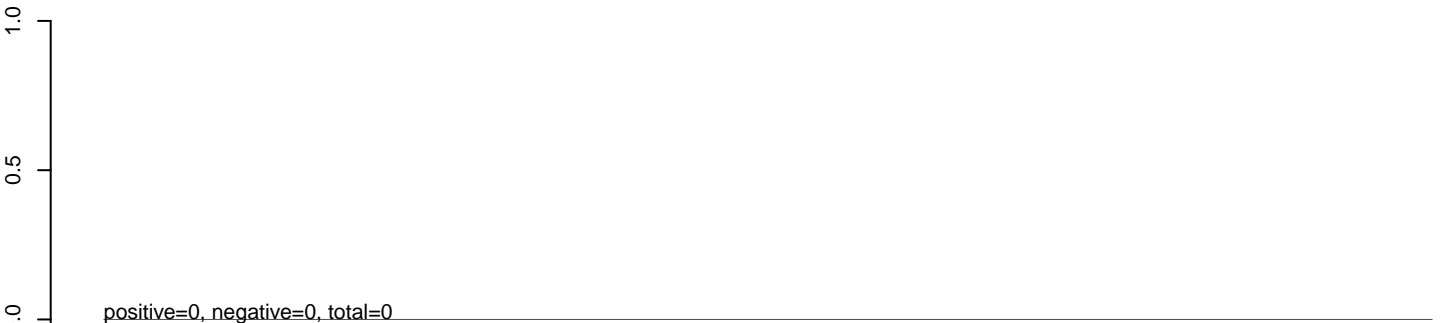


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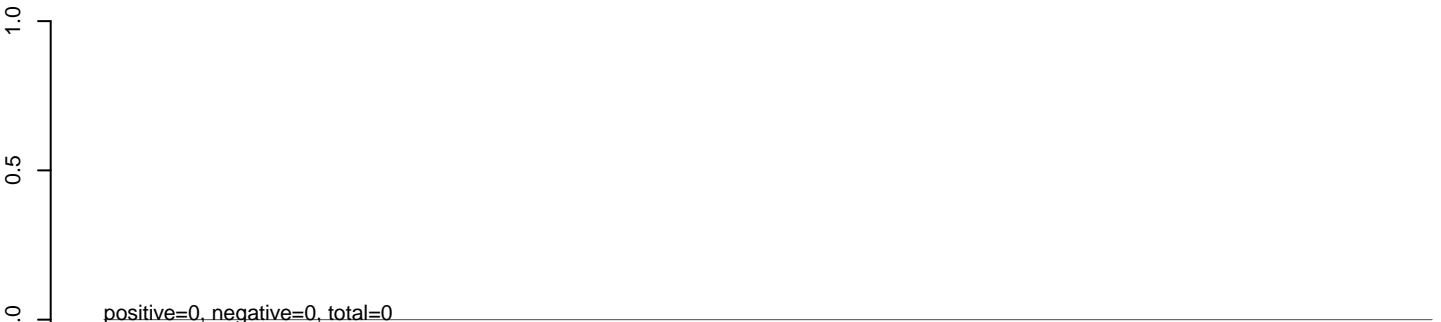


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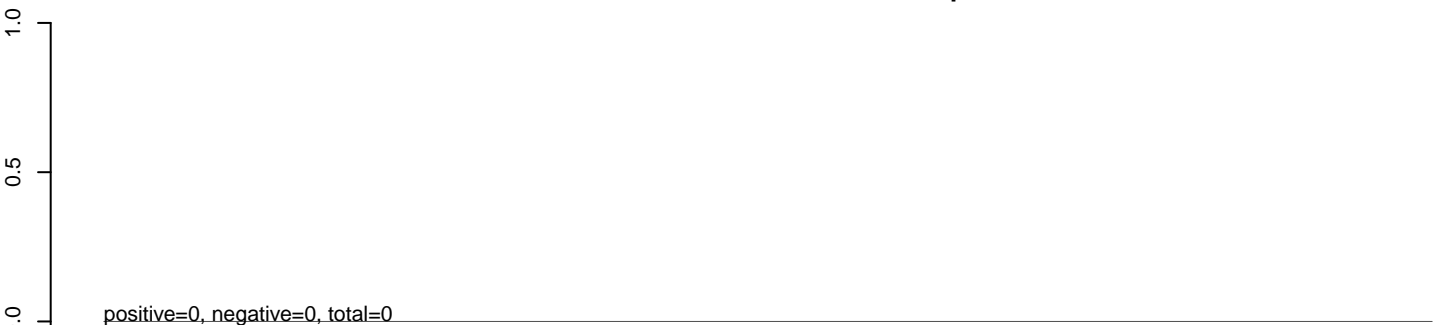
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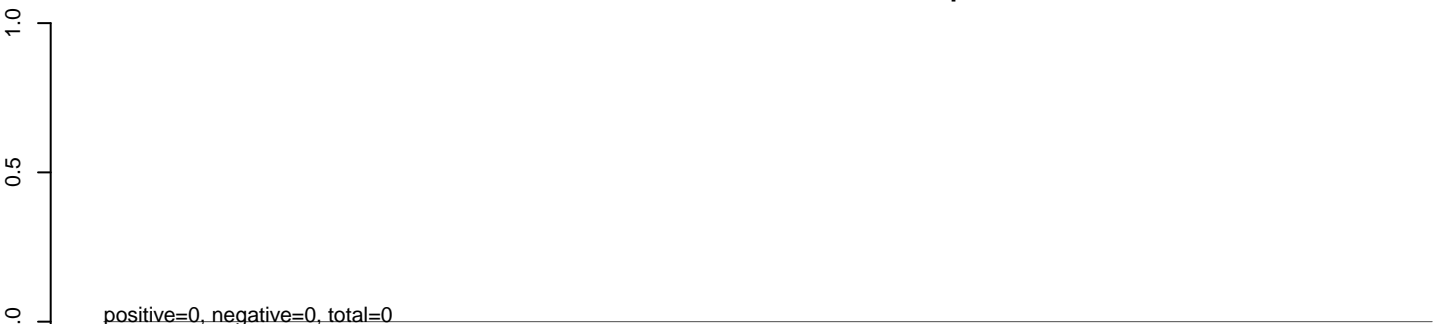
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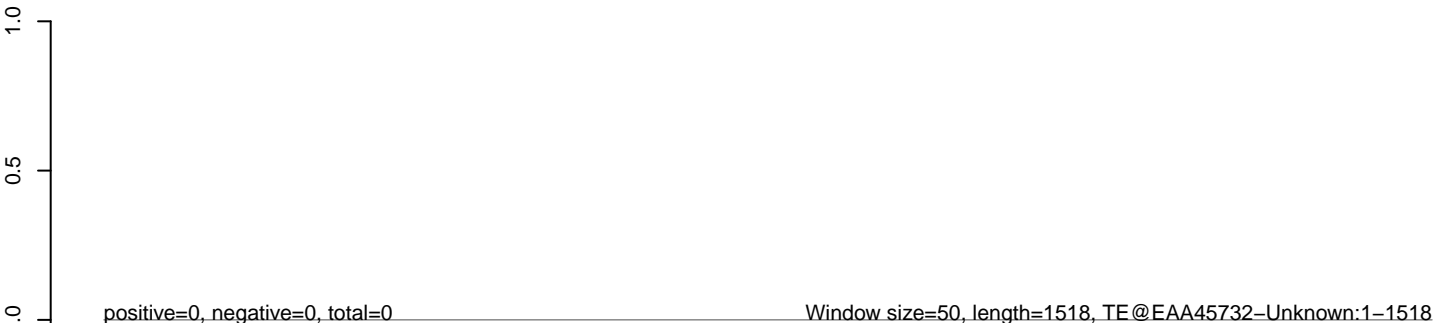
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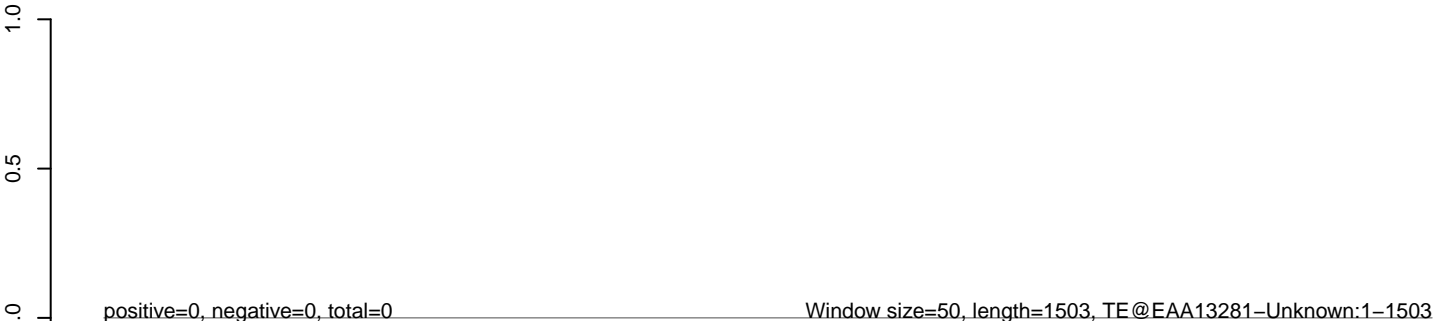
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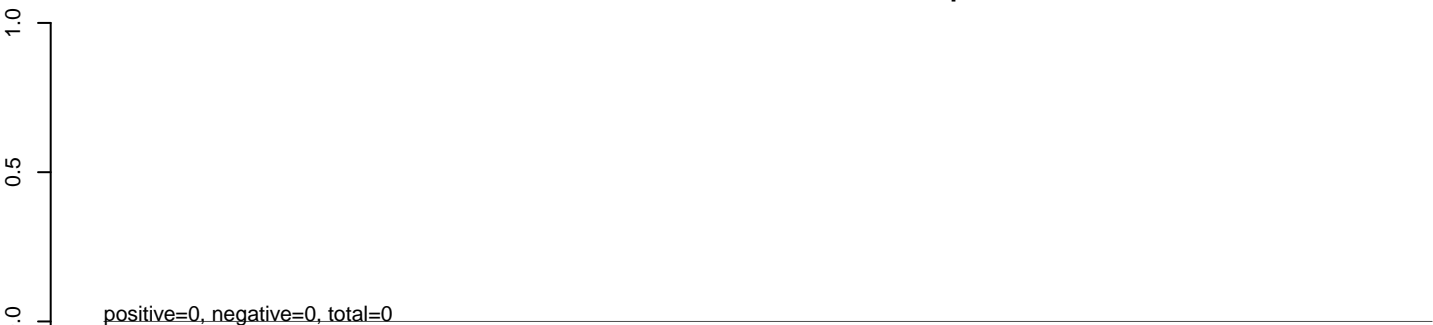
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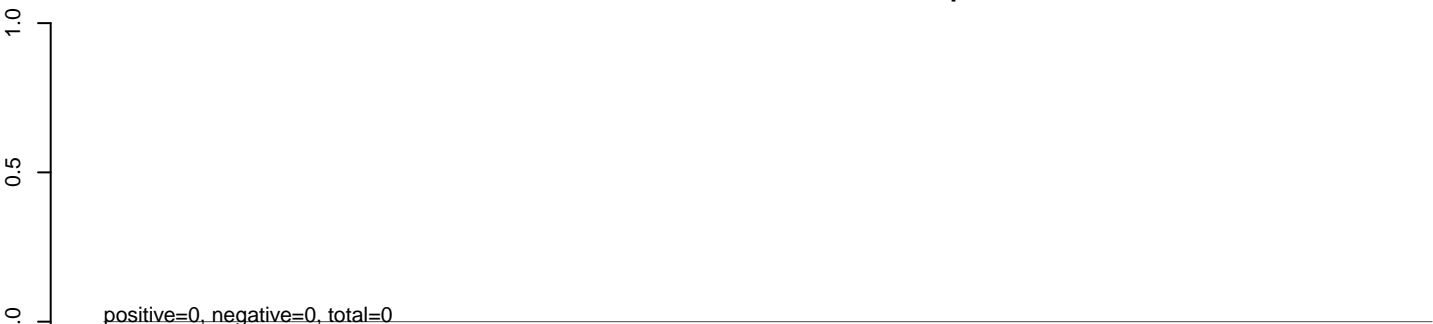
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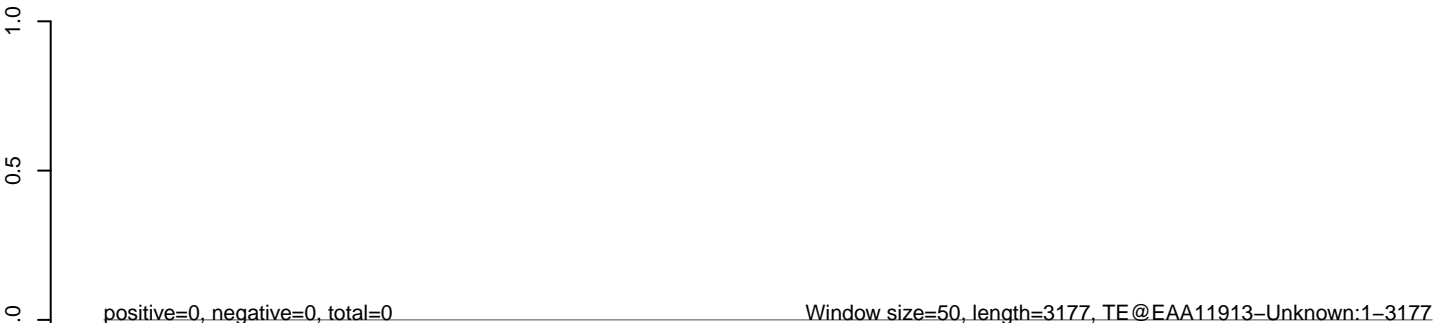
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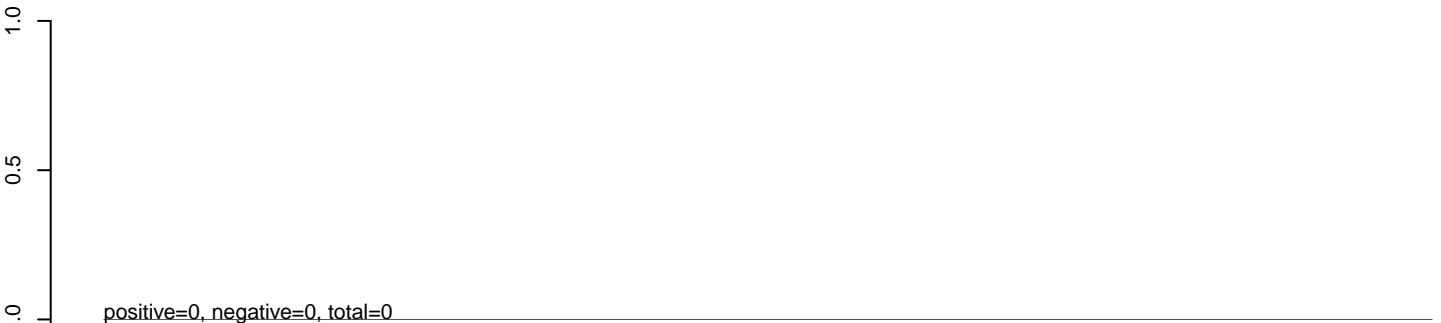


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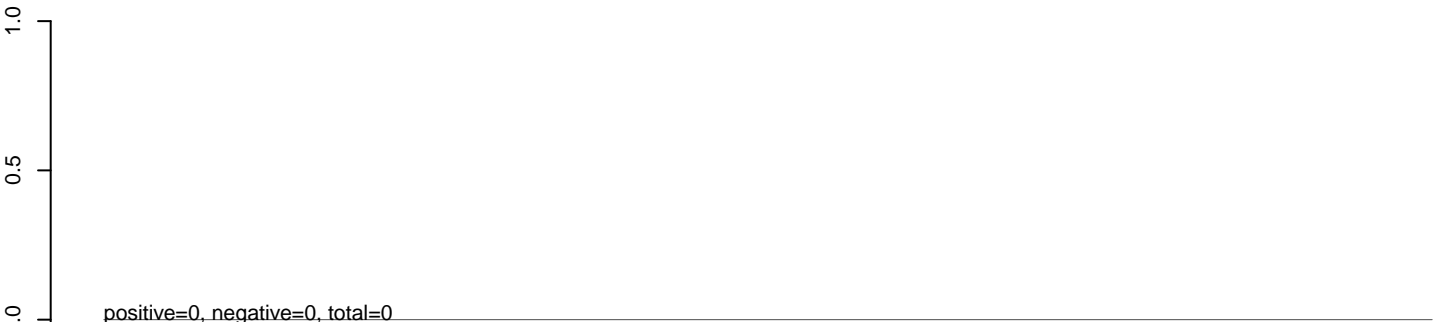


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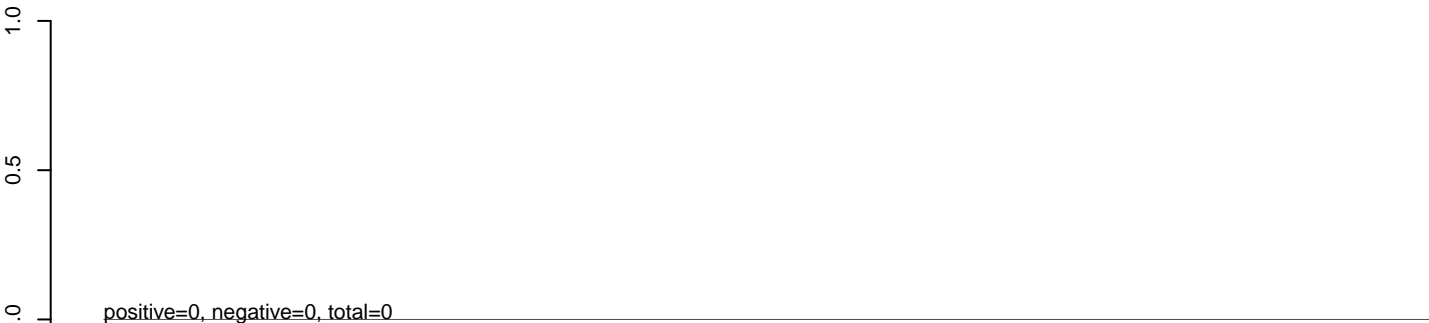
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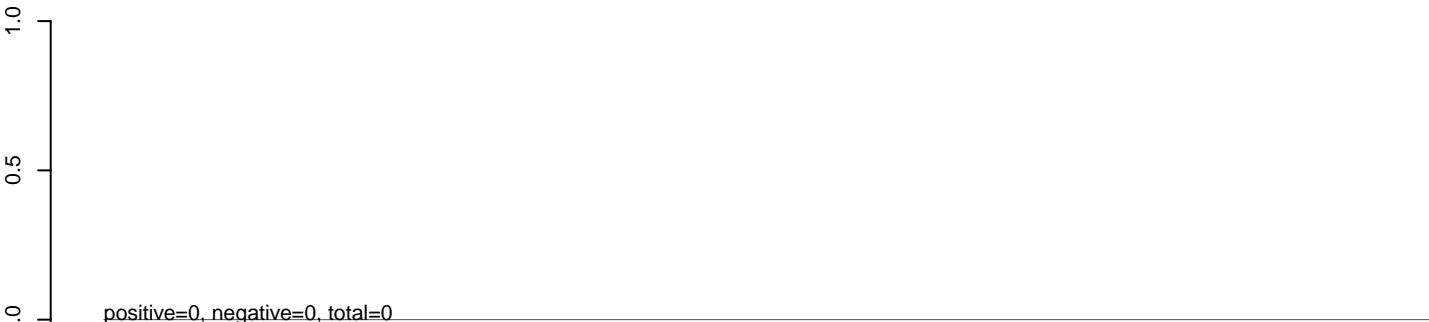
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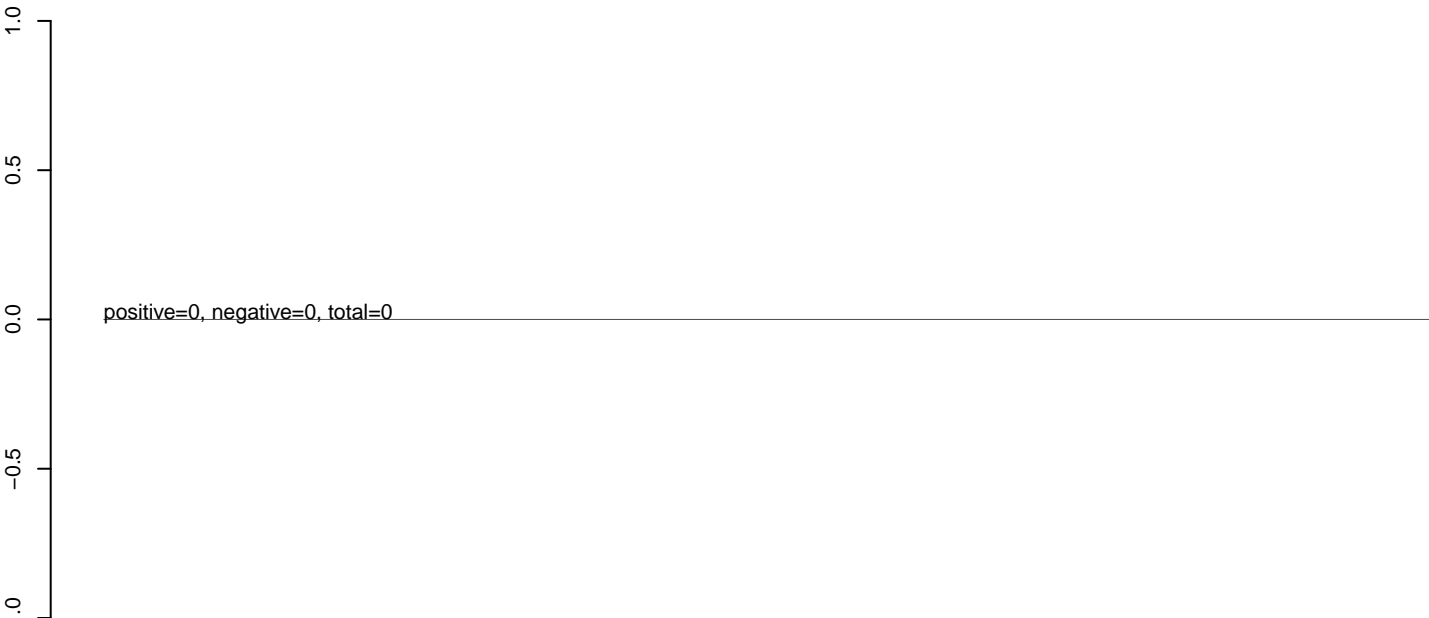
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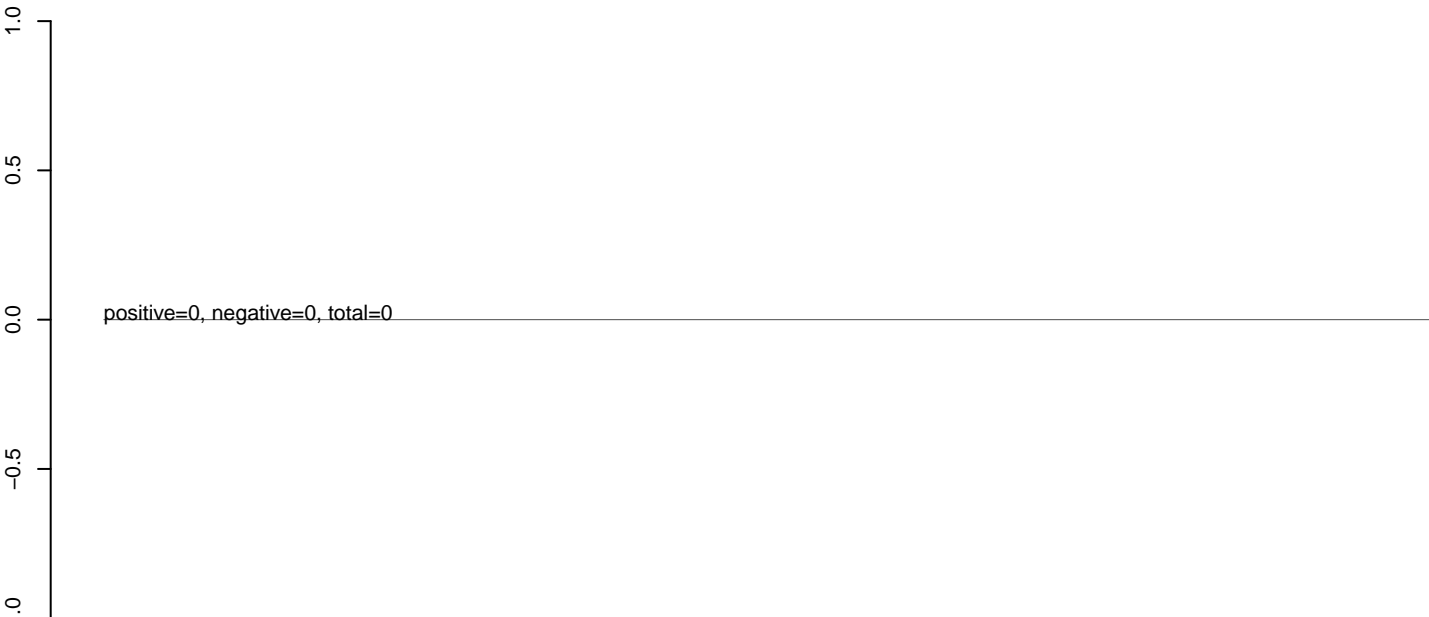
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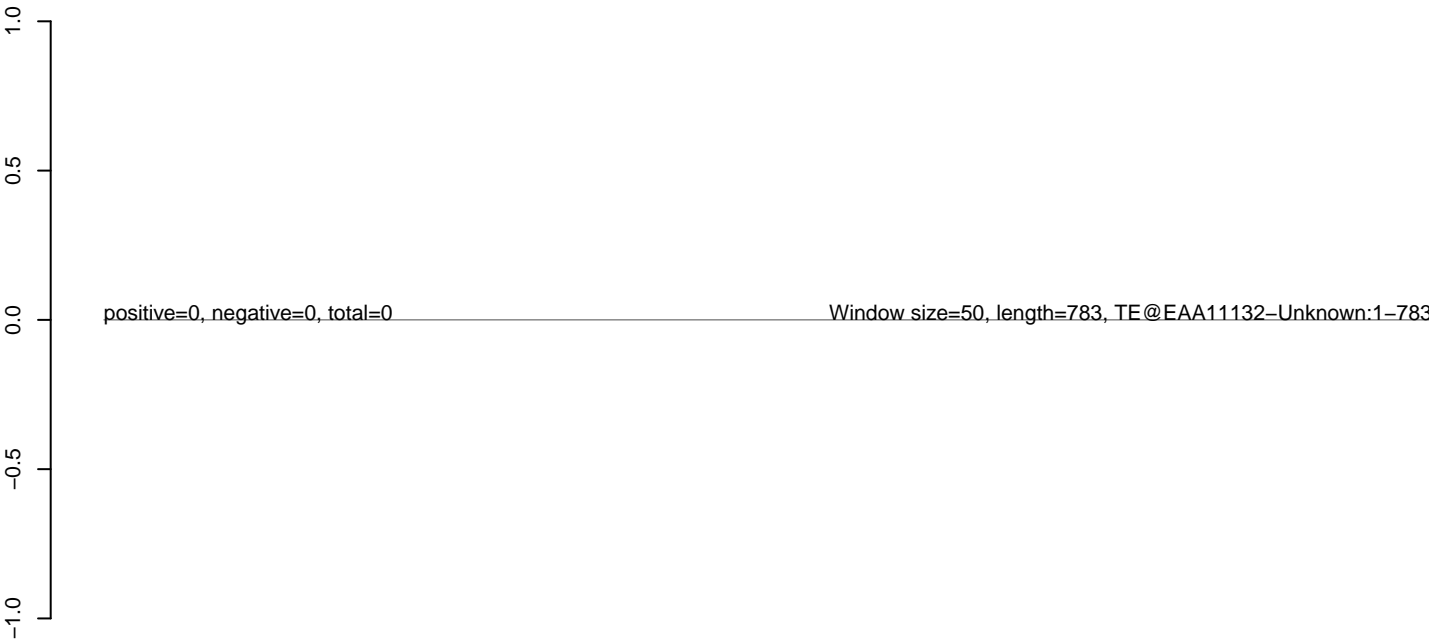
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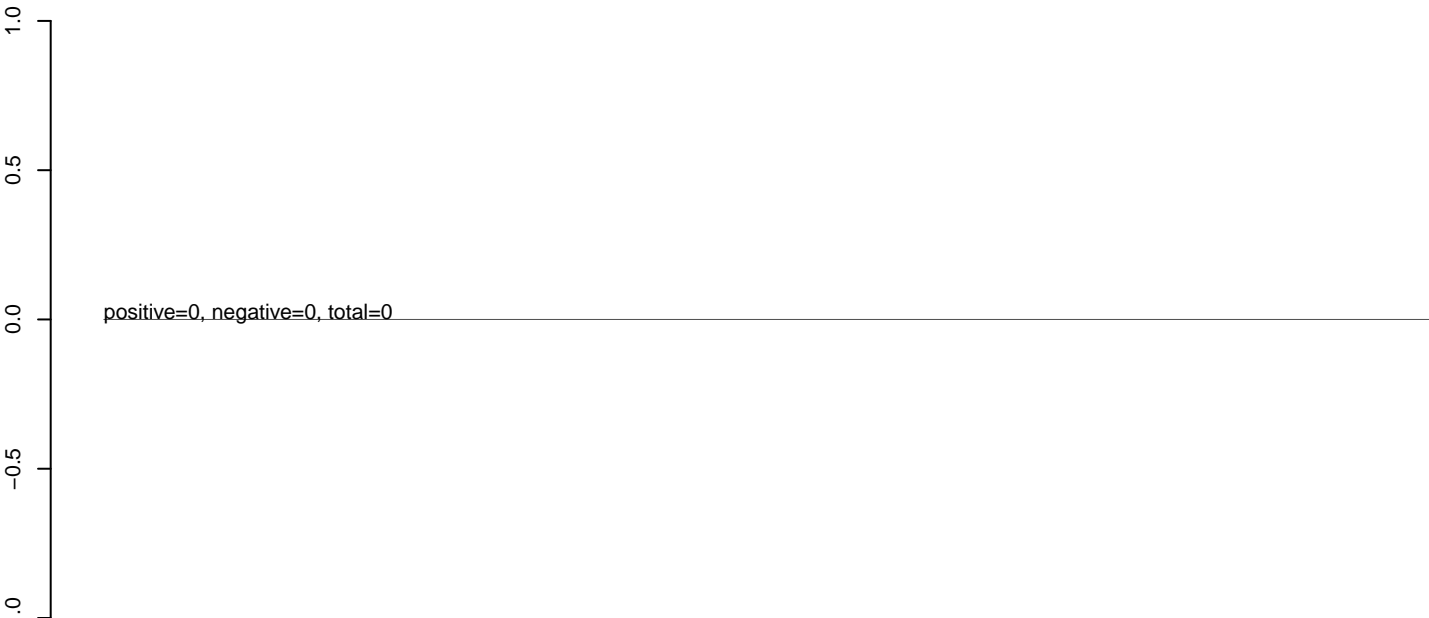
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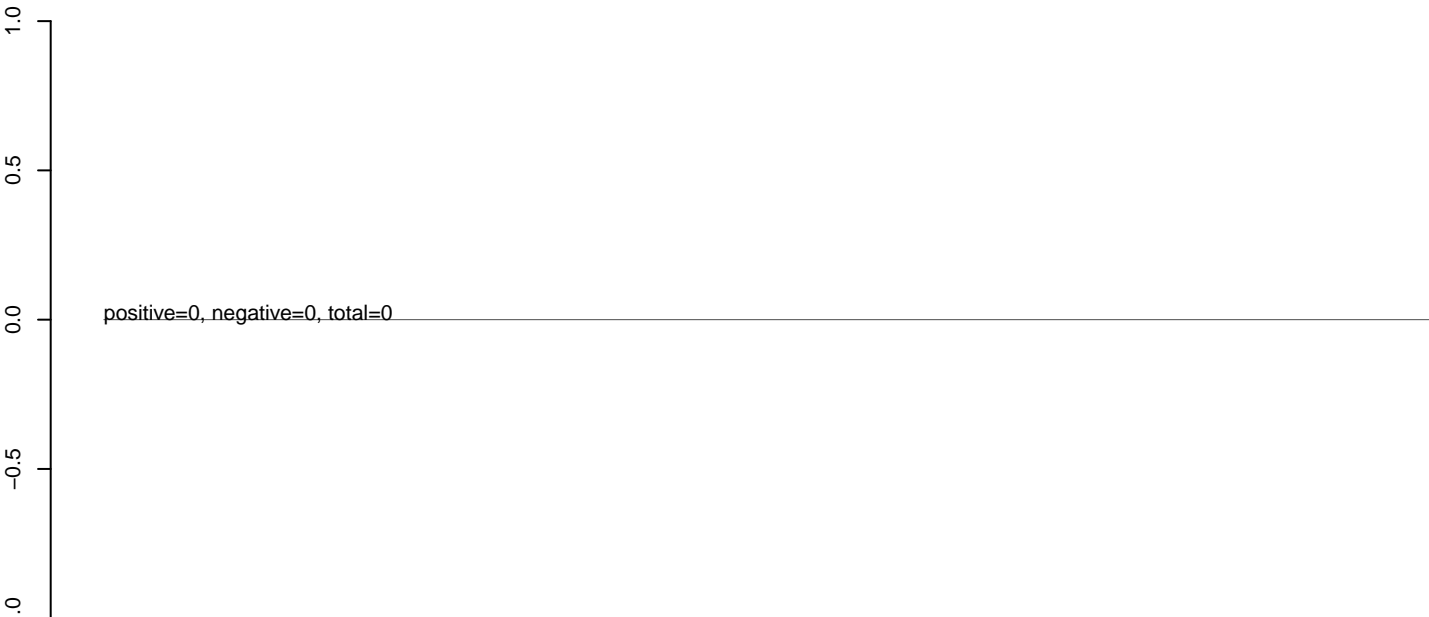
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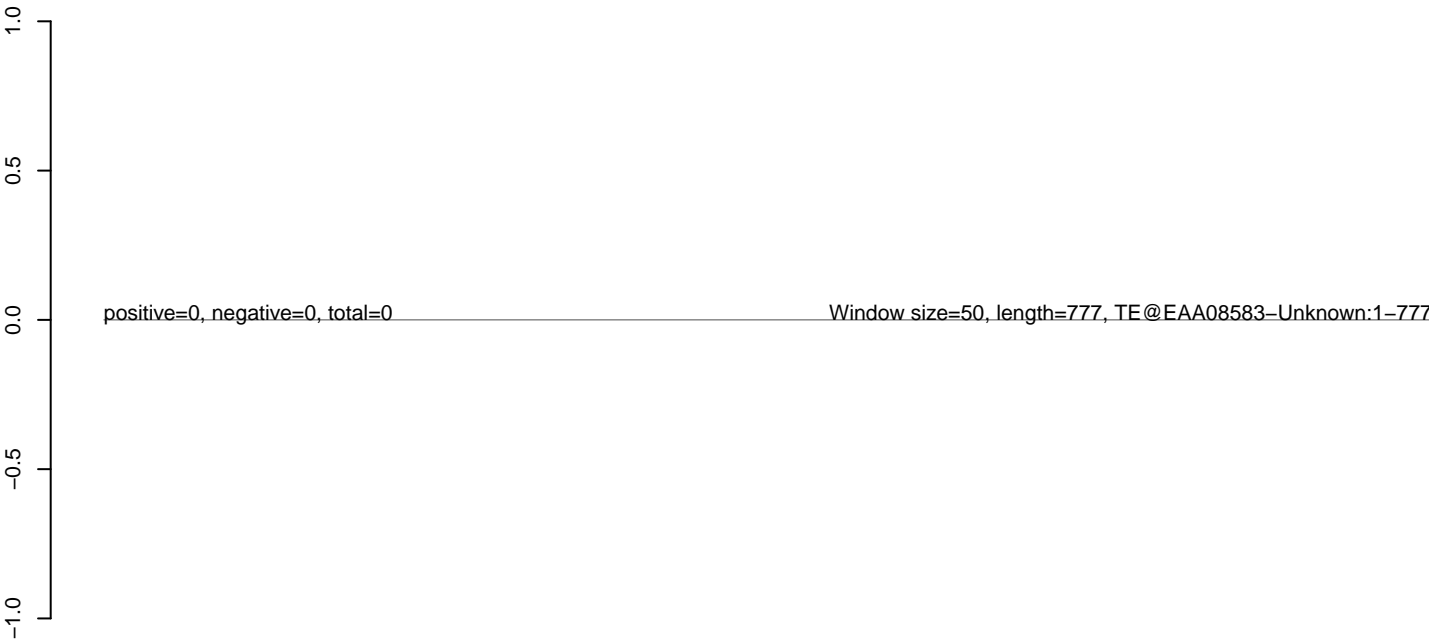
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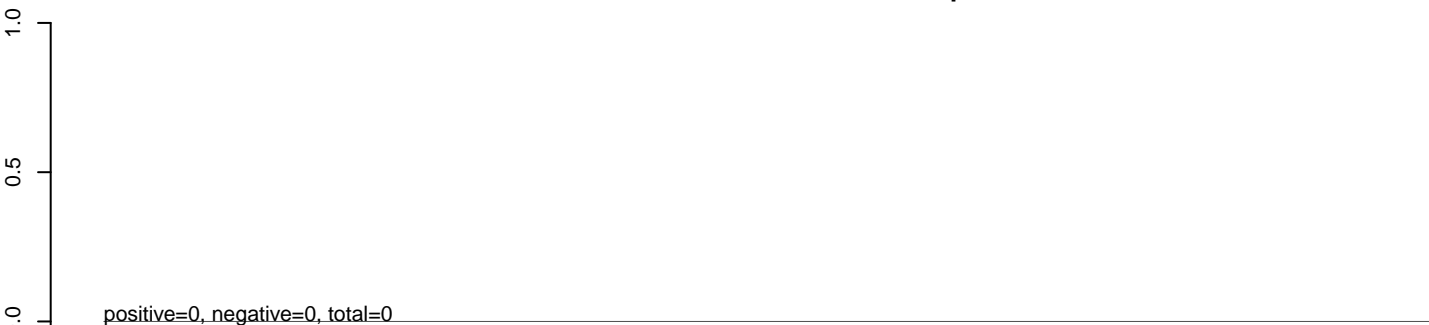
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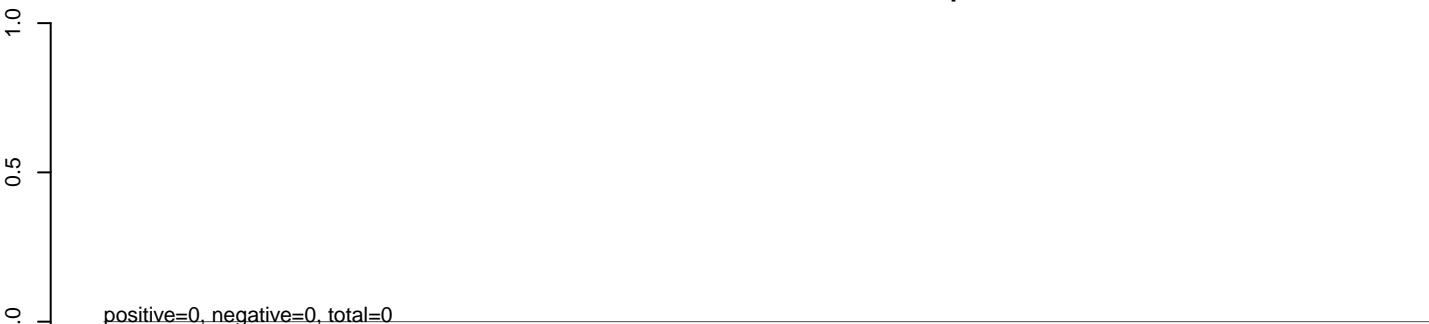
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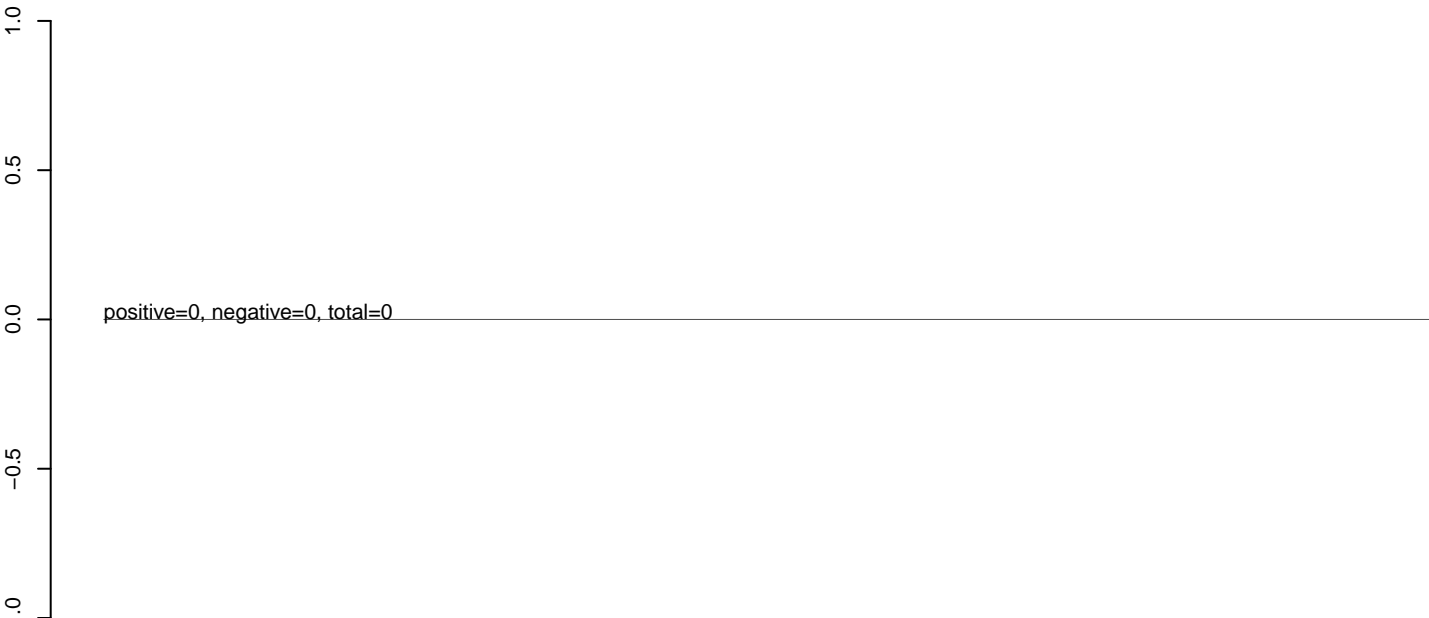
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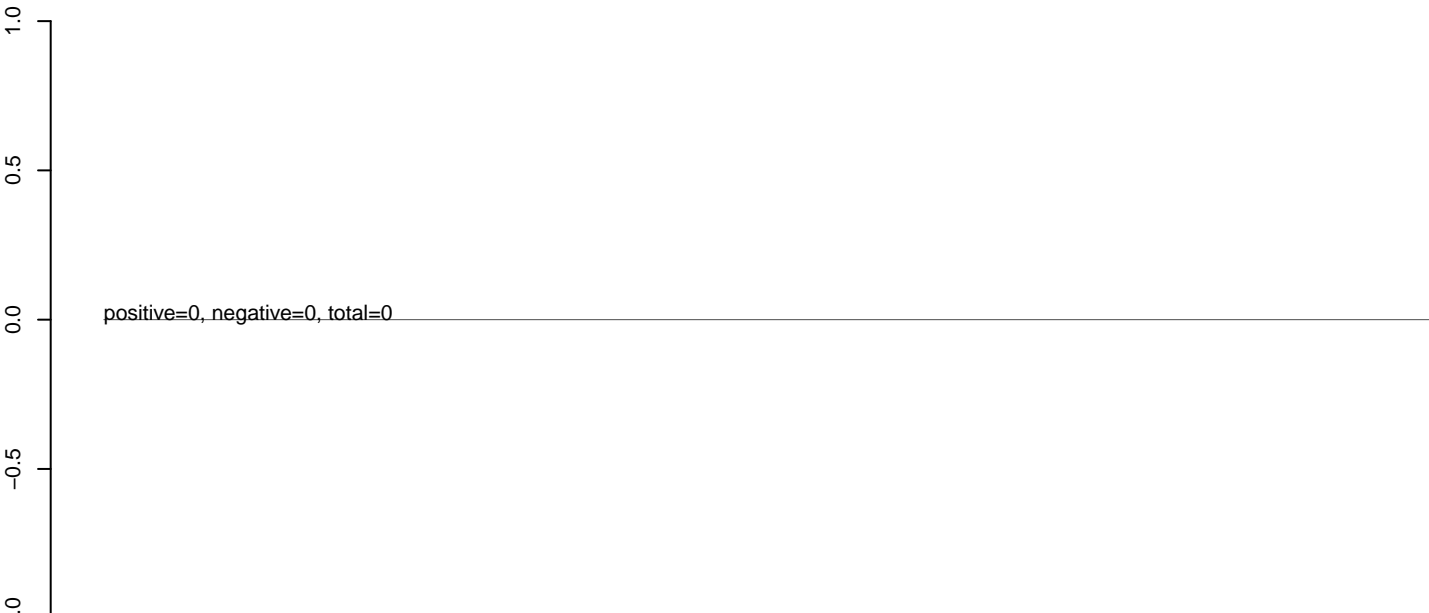
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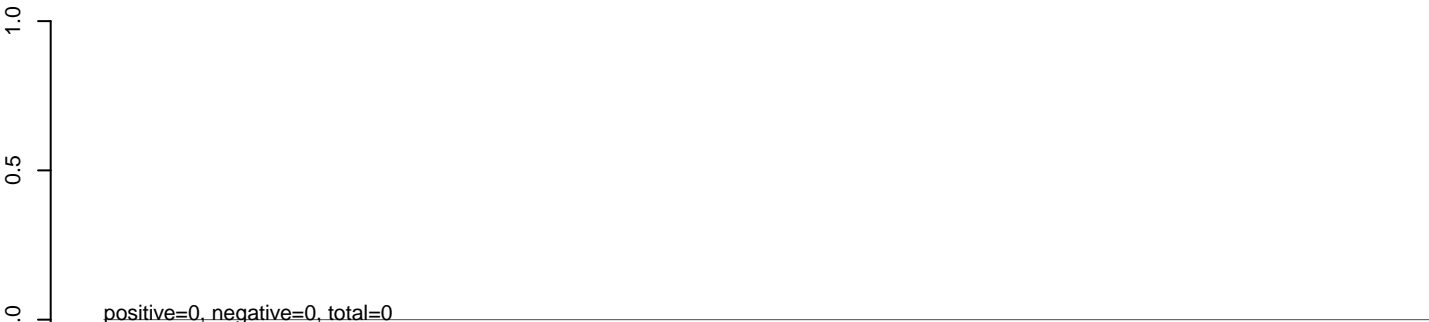
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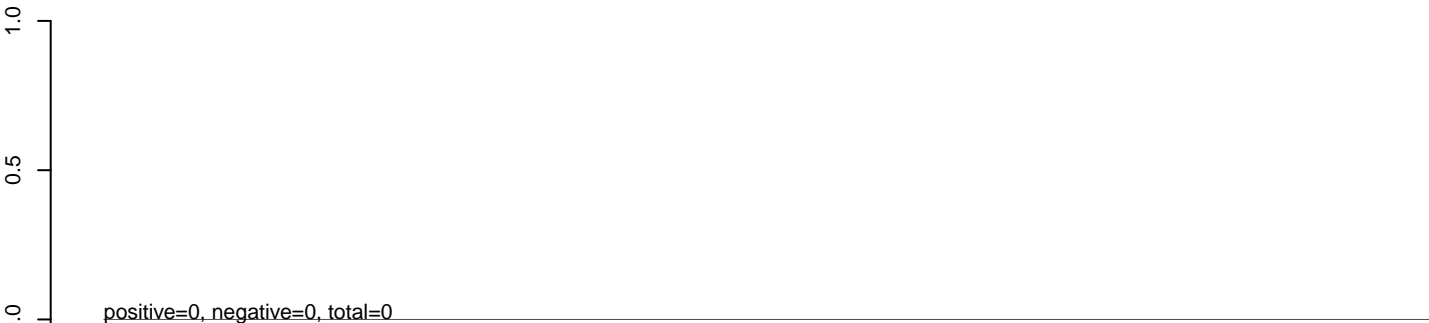
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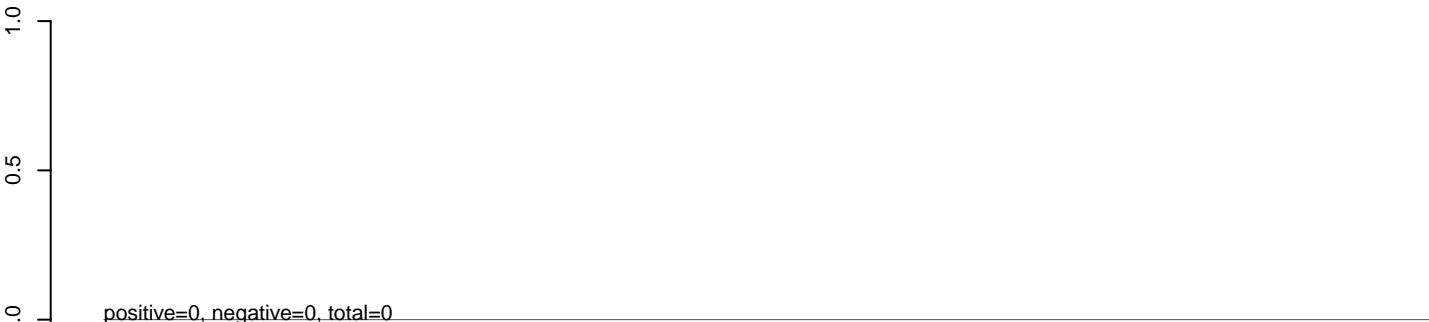
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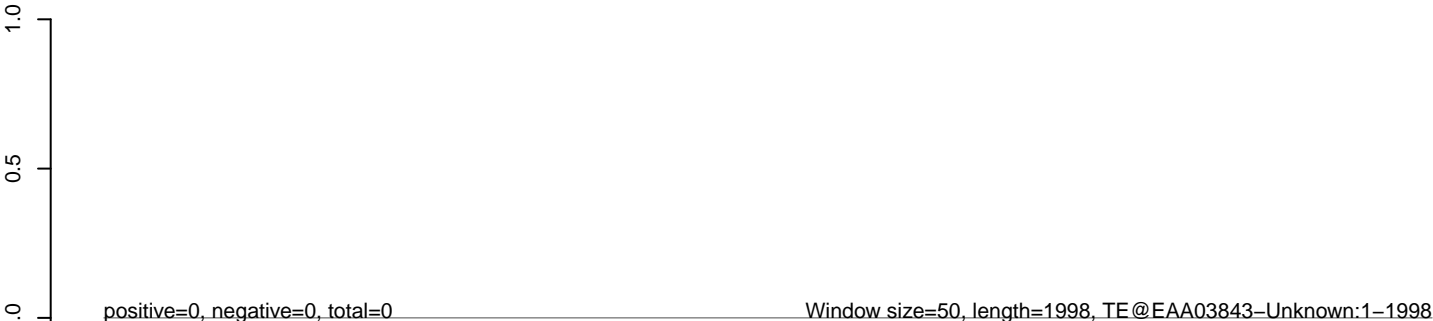
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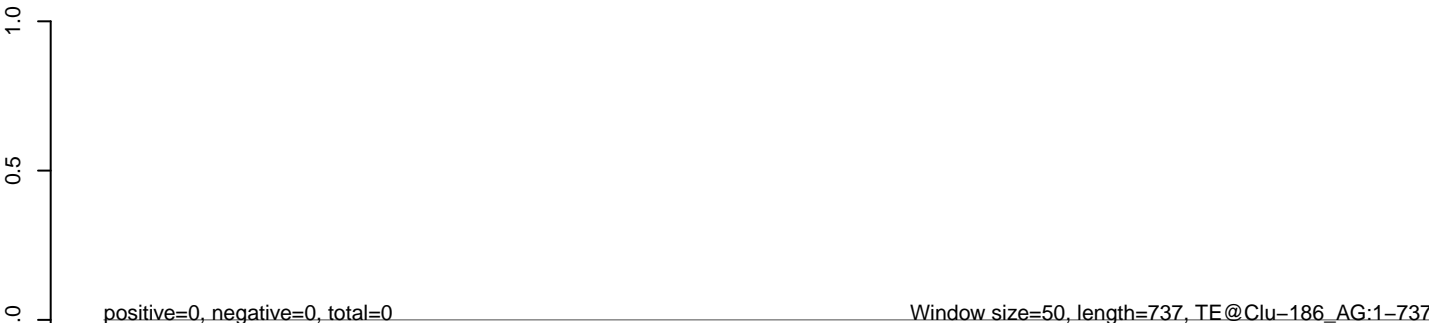
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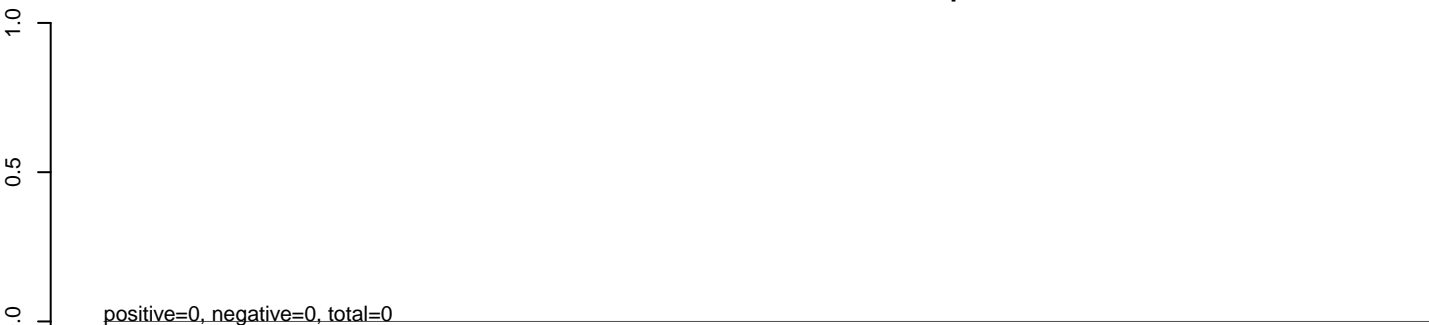


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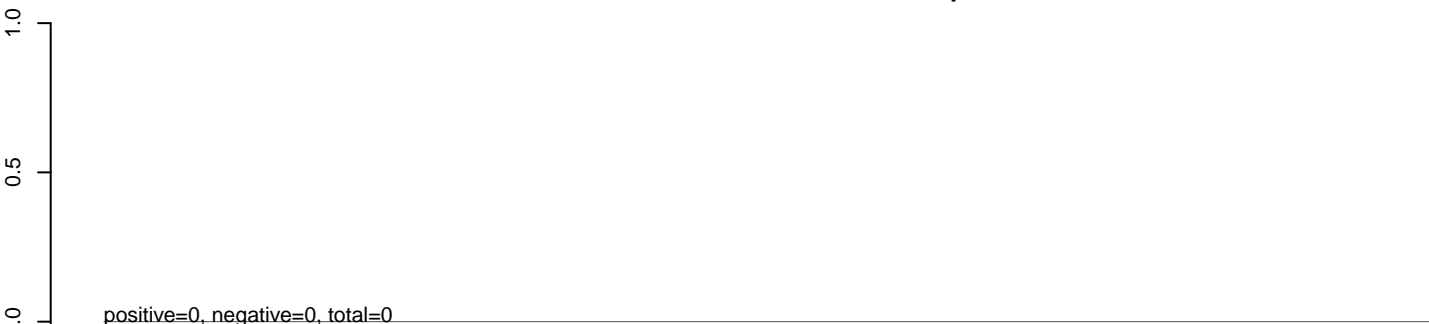


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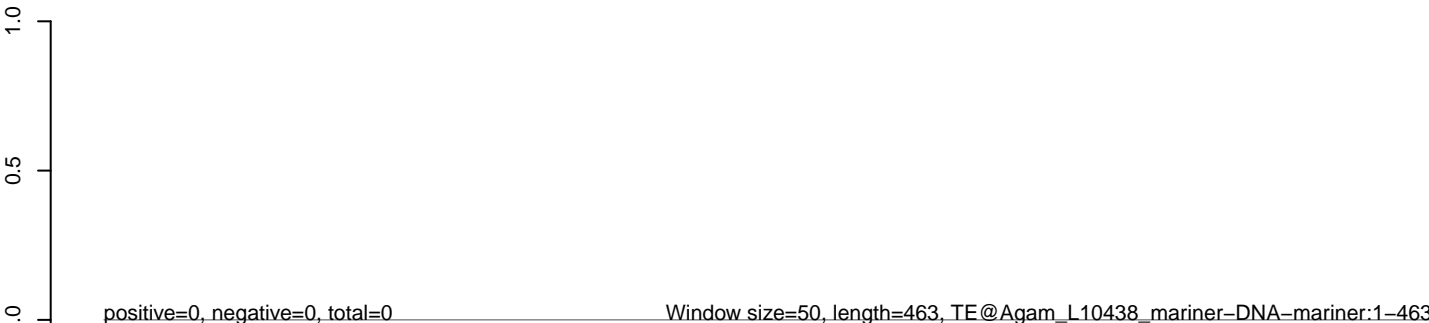
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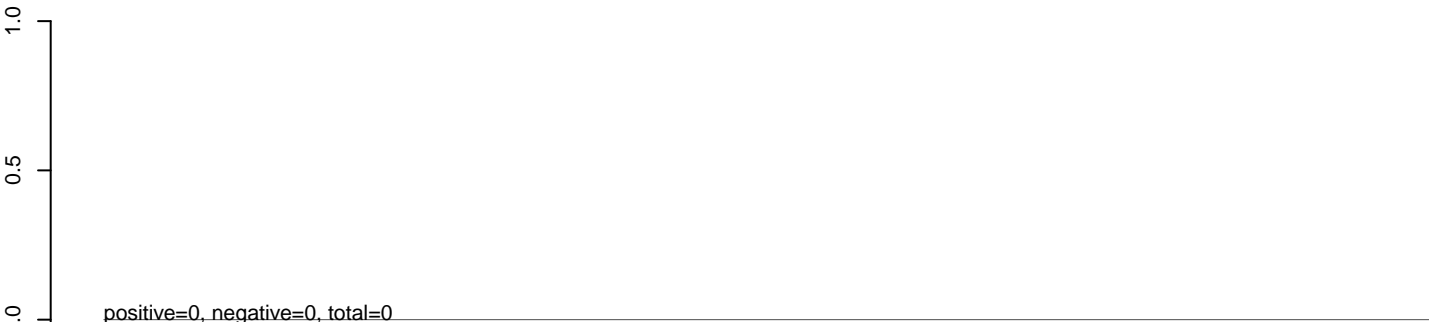
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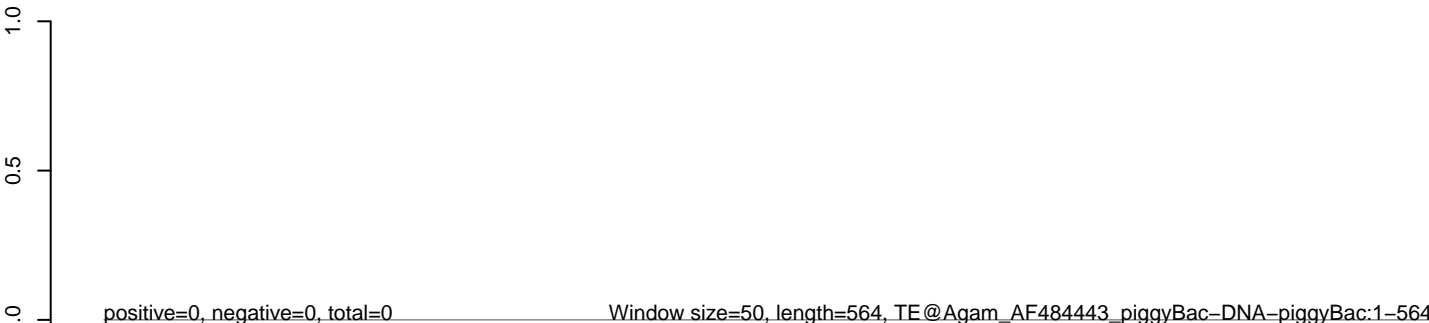
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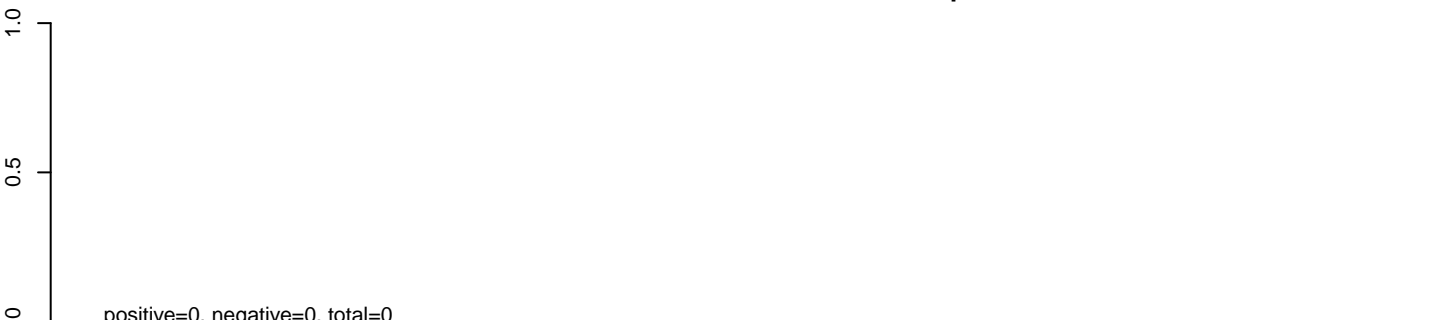
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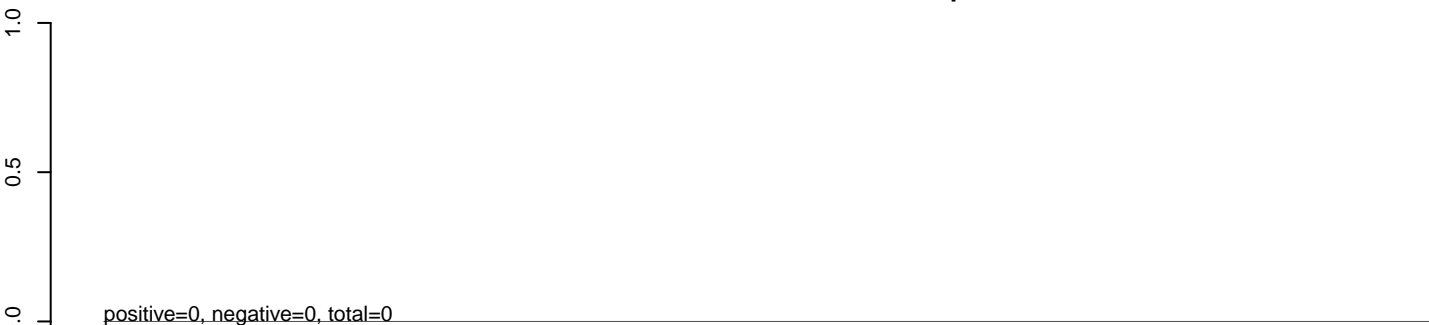
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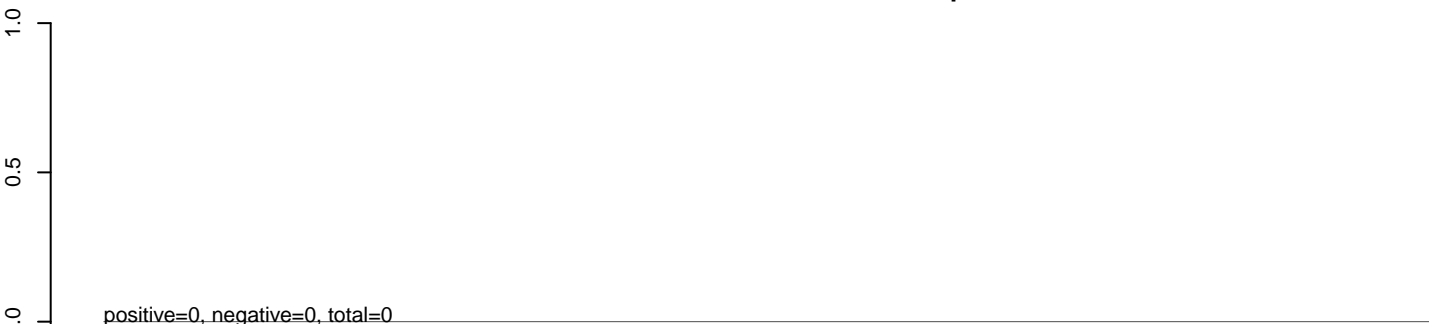
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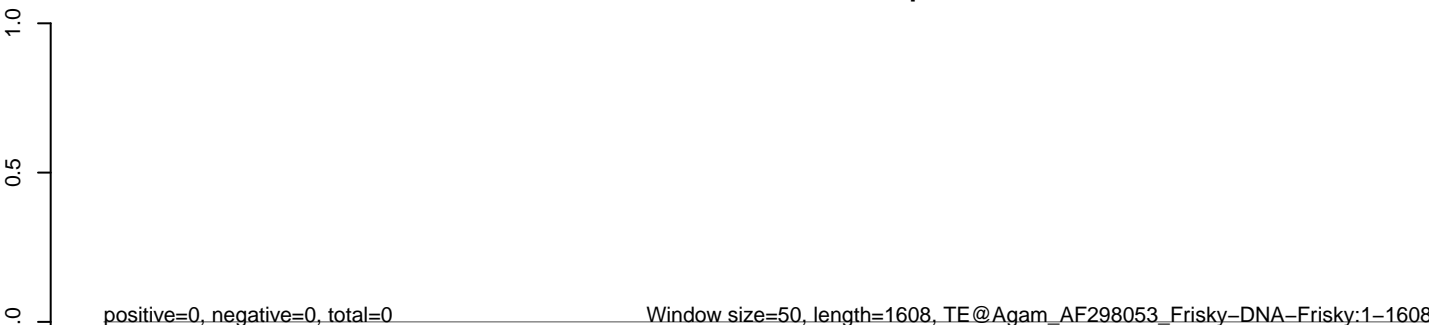
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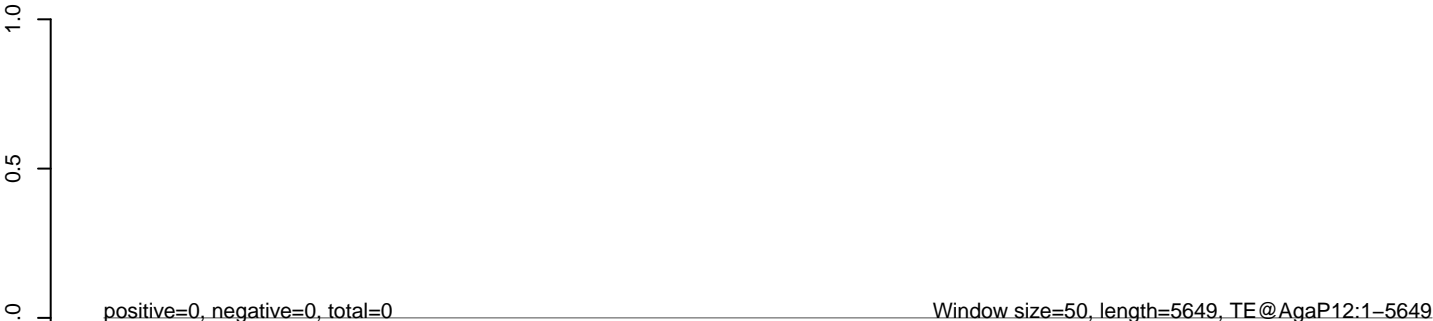
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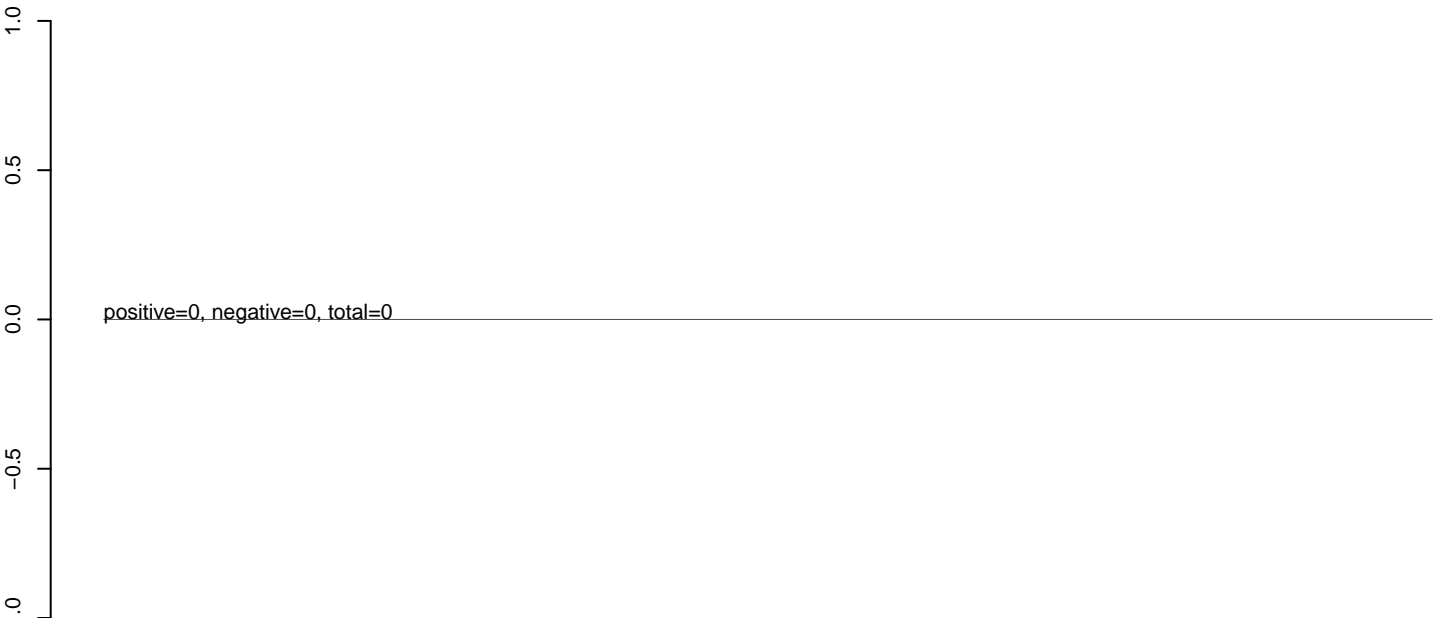


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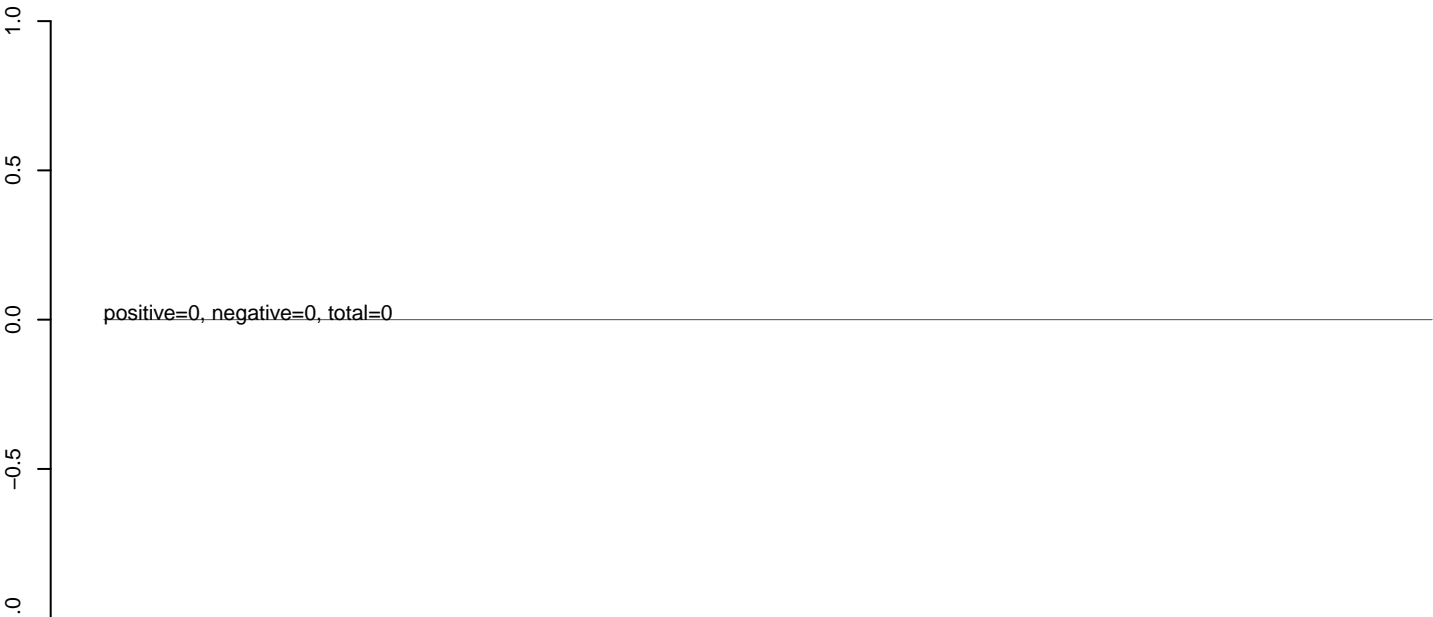


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