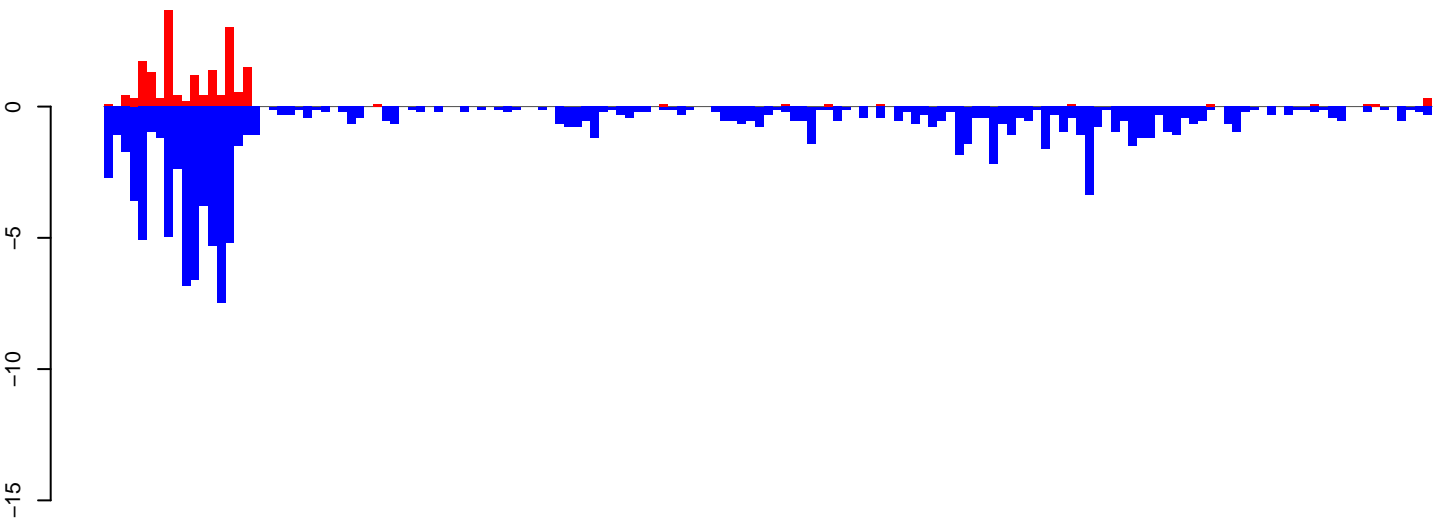
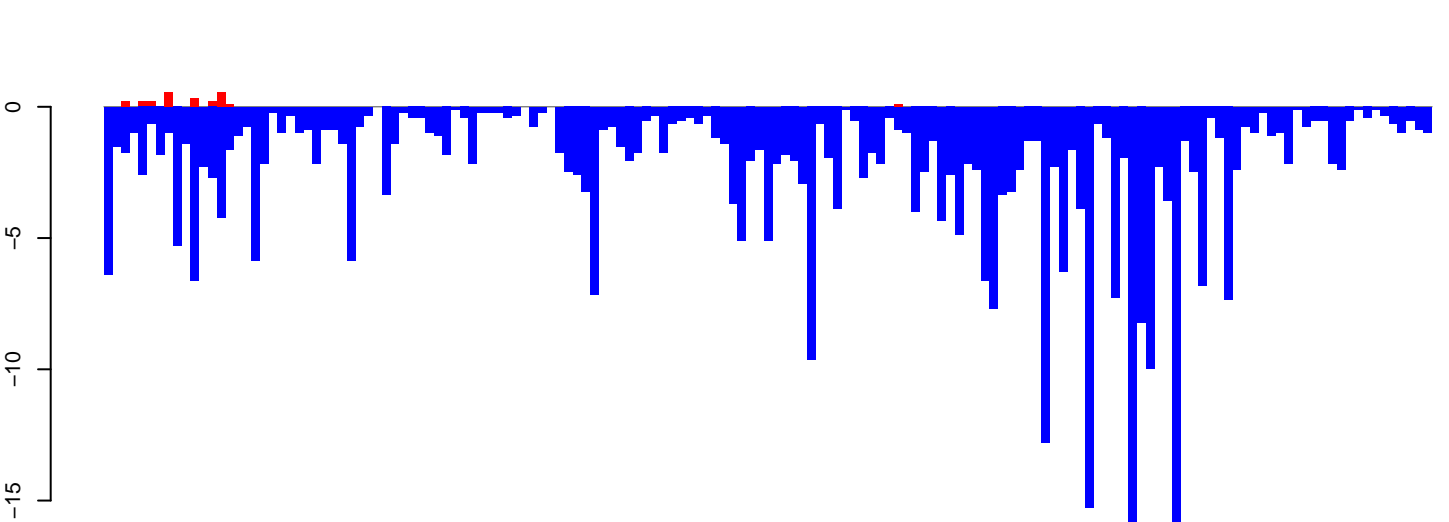


AnGam_Sua5b_cells.18_23.rep



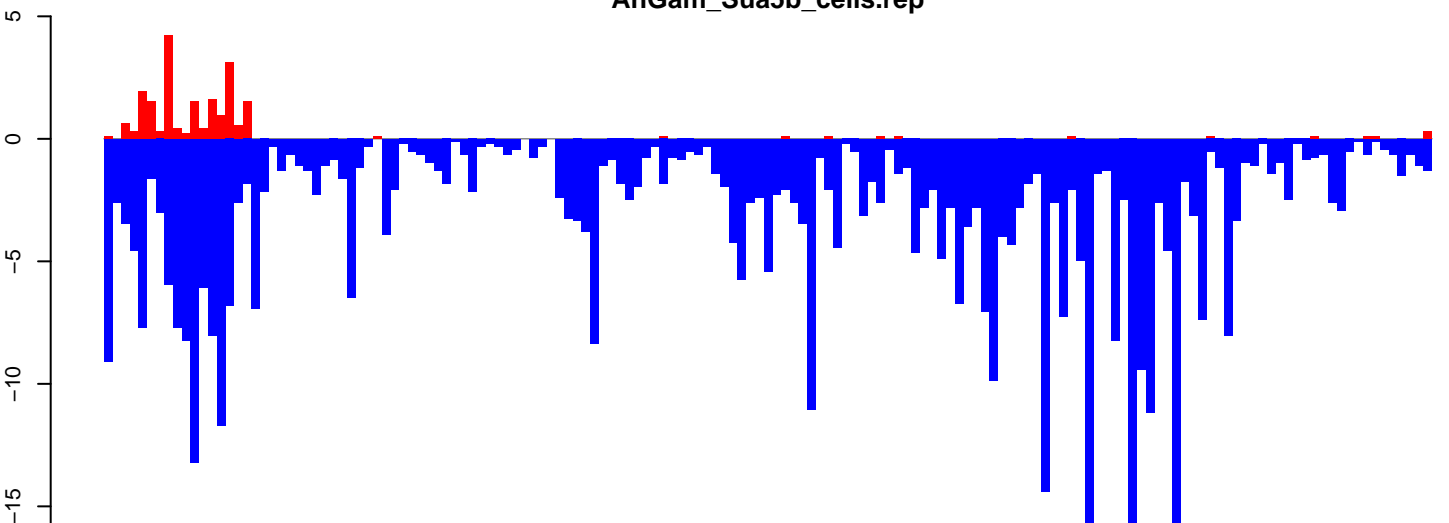
positive=19, negative=119, total=138

AnGam_Sua5b_cells.24_35.rep



positive=2, negative=369, total=371

AnGam_Sua5b_cells.rep

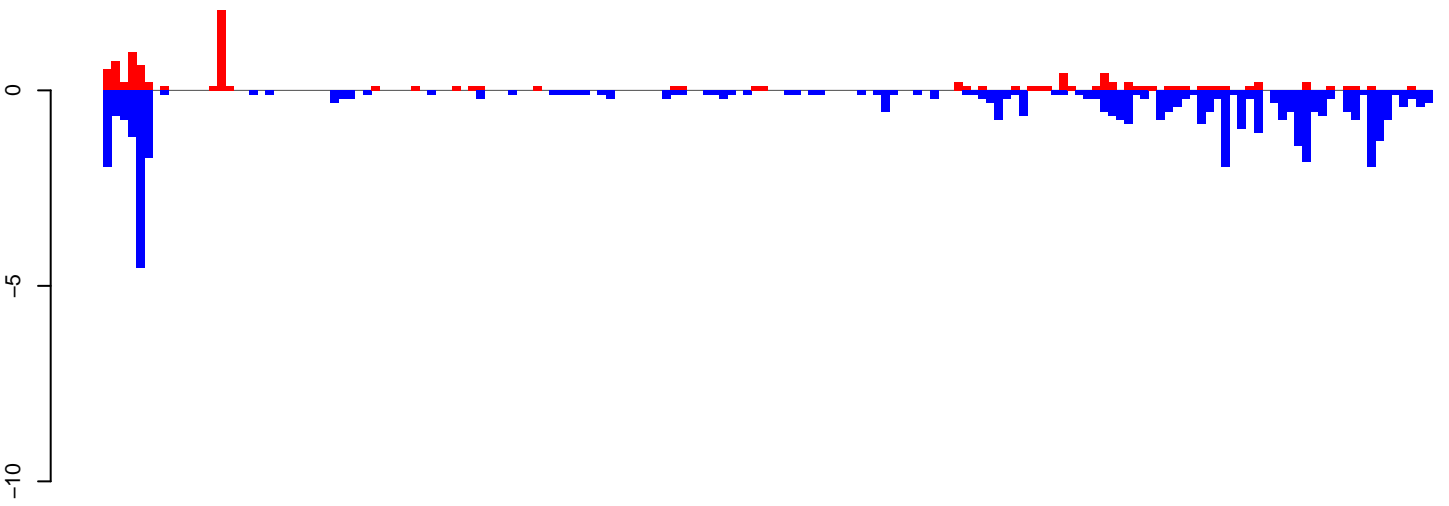


positive=21, negative=488, total=509

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

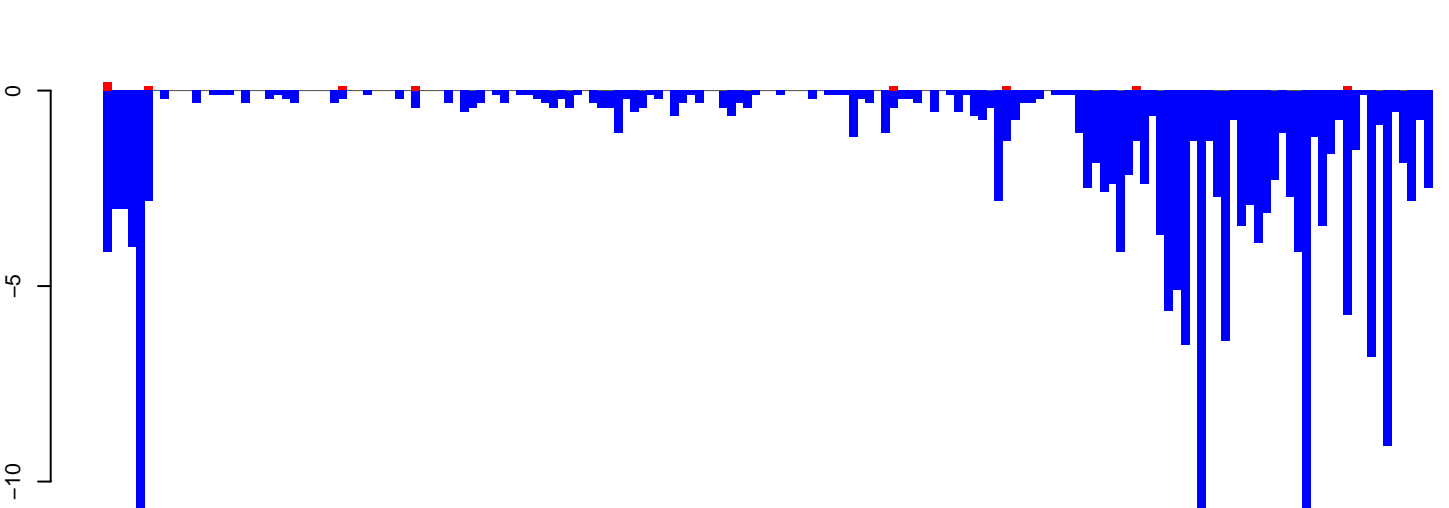
0 2000 4000 6000 8000

AnGam_Sua5b_cells.18_23.rep



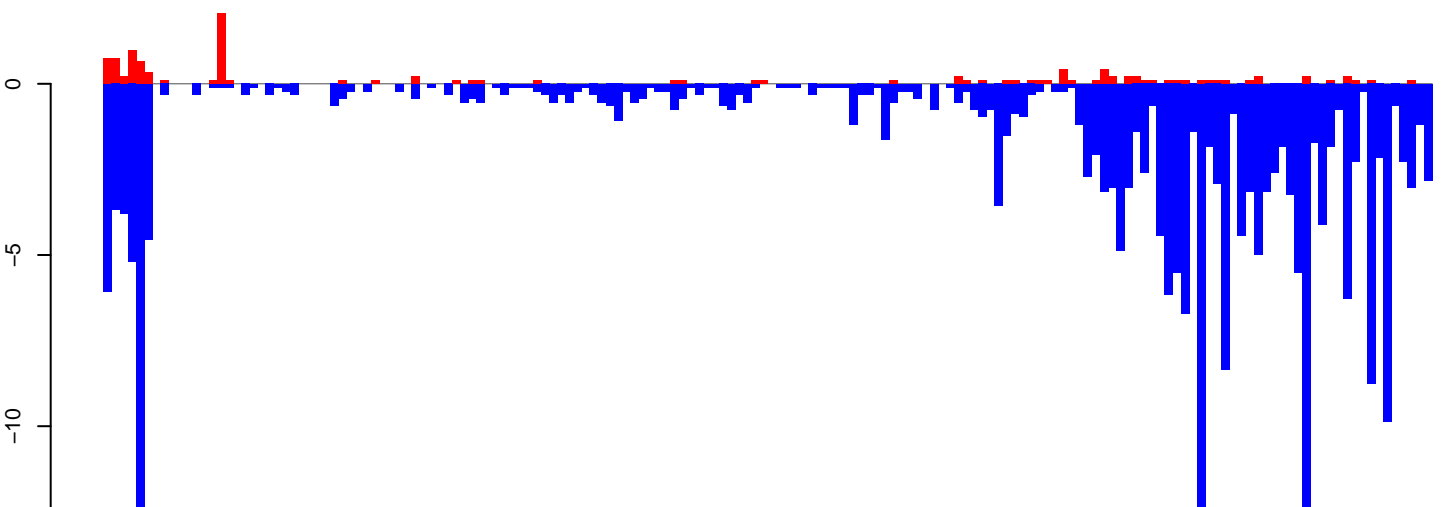
positive=11, negative=44, total=55

AnGam_Sua5b_cells.24_35.rep



positive=1, negative=202, total=203

AnGam_Sua5b_cells.rep

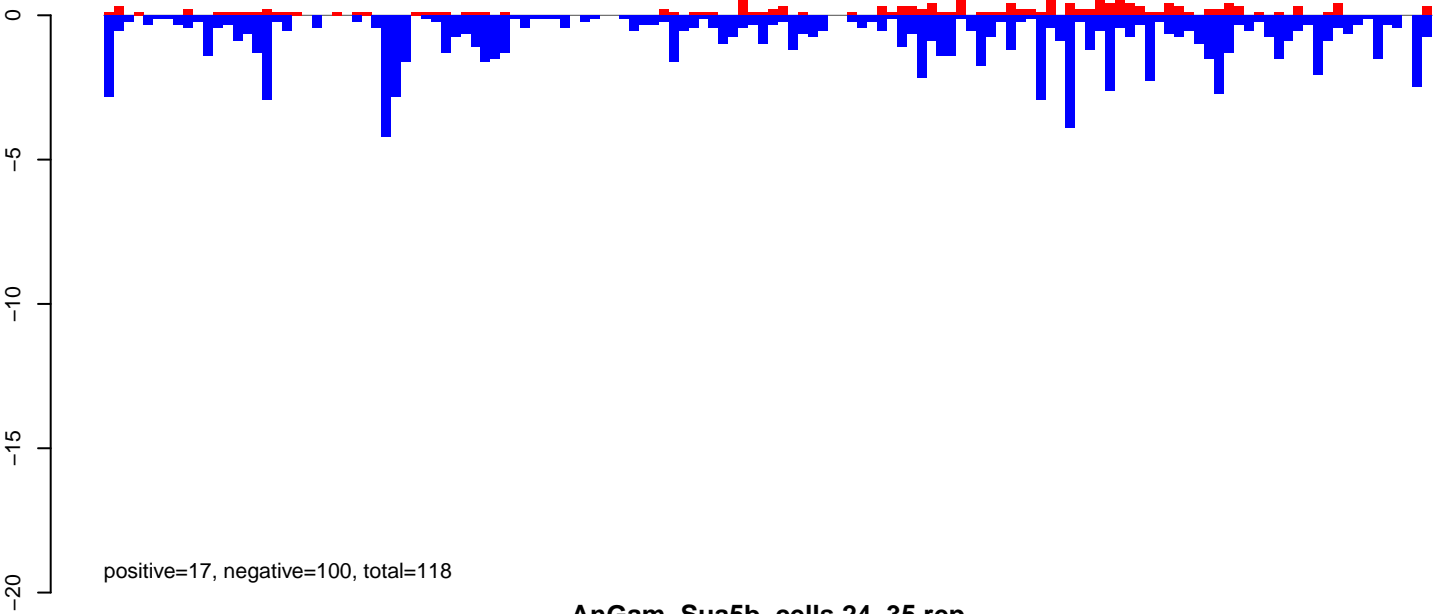


positive=12, negative=246, total=258

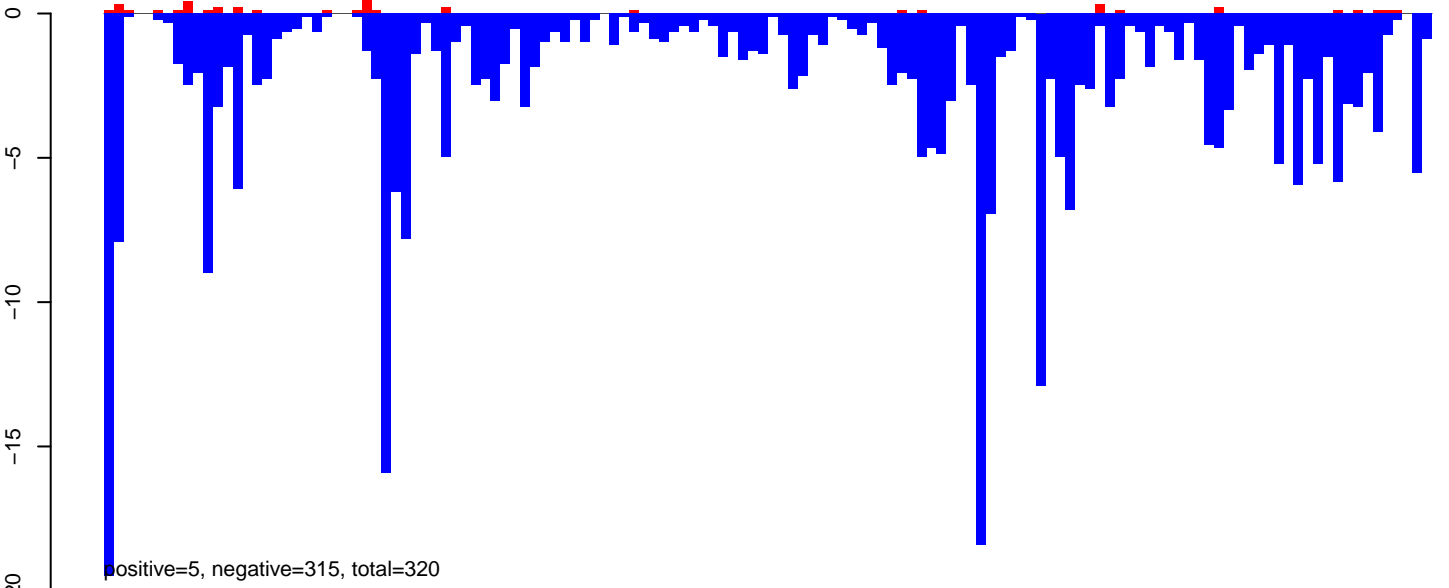
Window size=50, length=8249, TE@BEL1-LTR-1_AG:1-8249

0 2000 4000 6000 8000

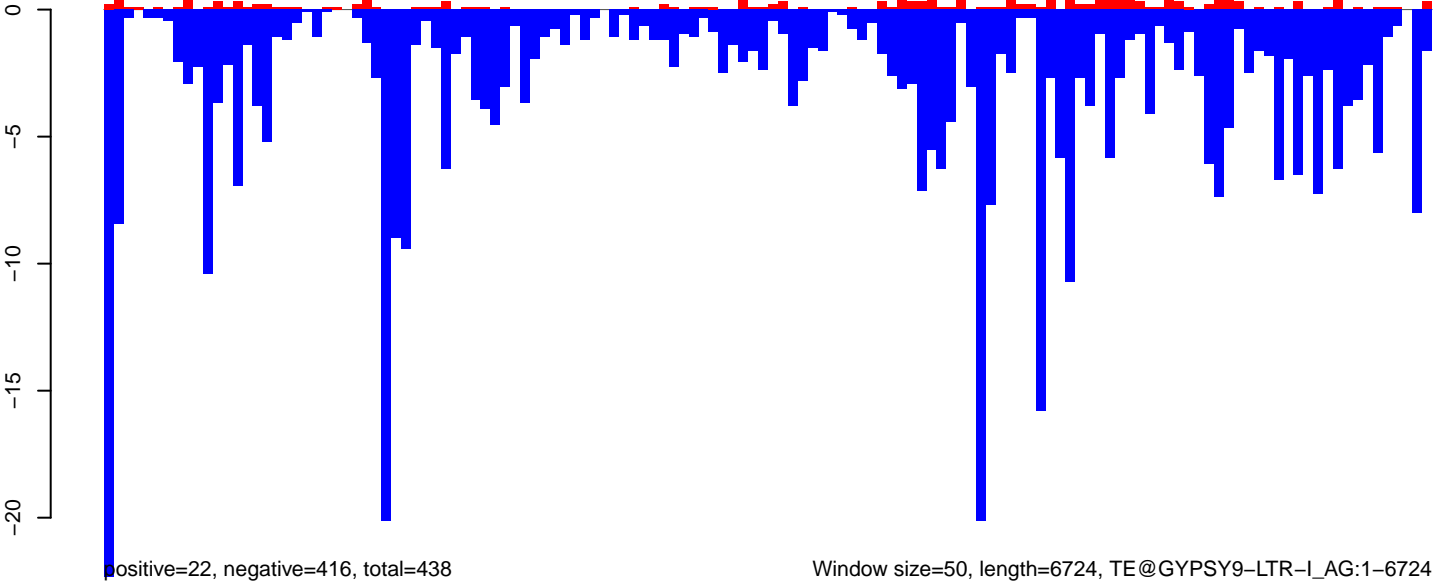
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



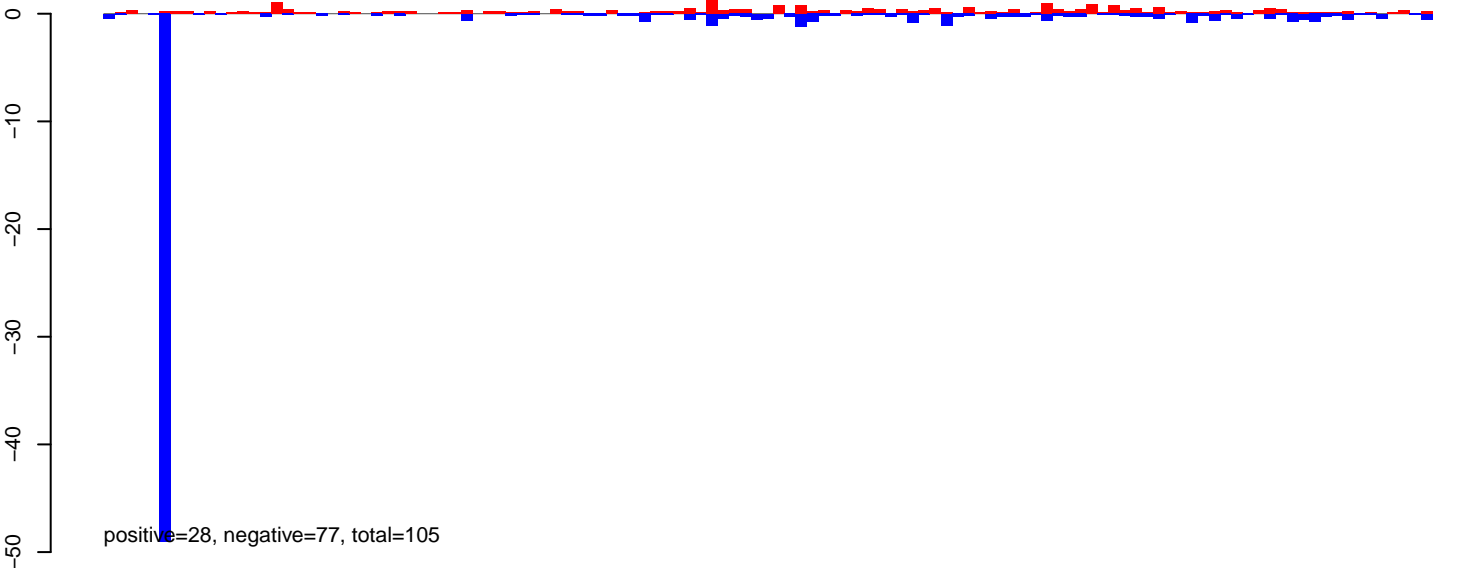
AnGam_Sua5b_cells.rep



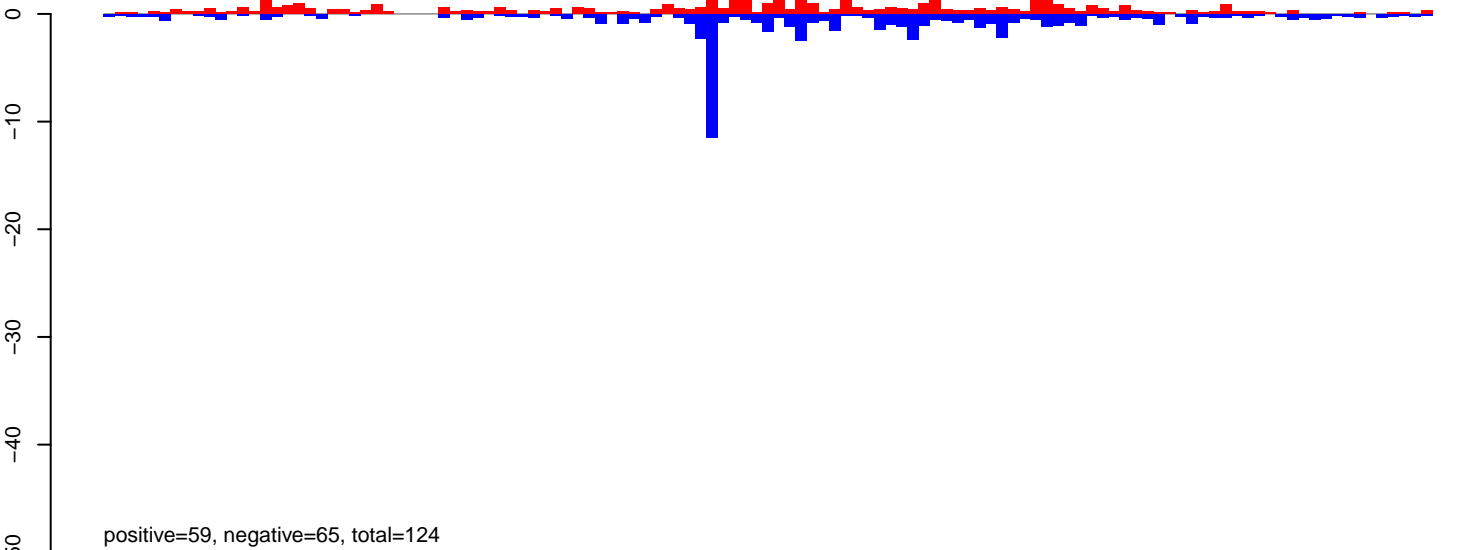
Window size=50, length=6724, TE@GYPSY9-LTR-I_AG:1-6724

0 1000 2000 3000 4000 5000 6000 7000

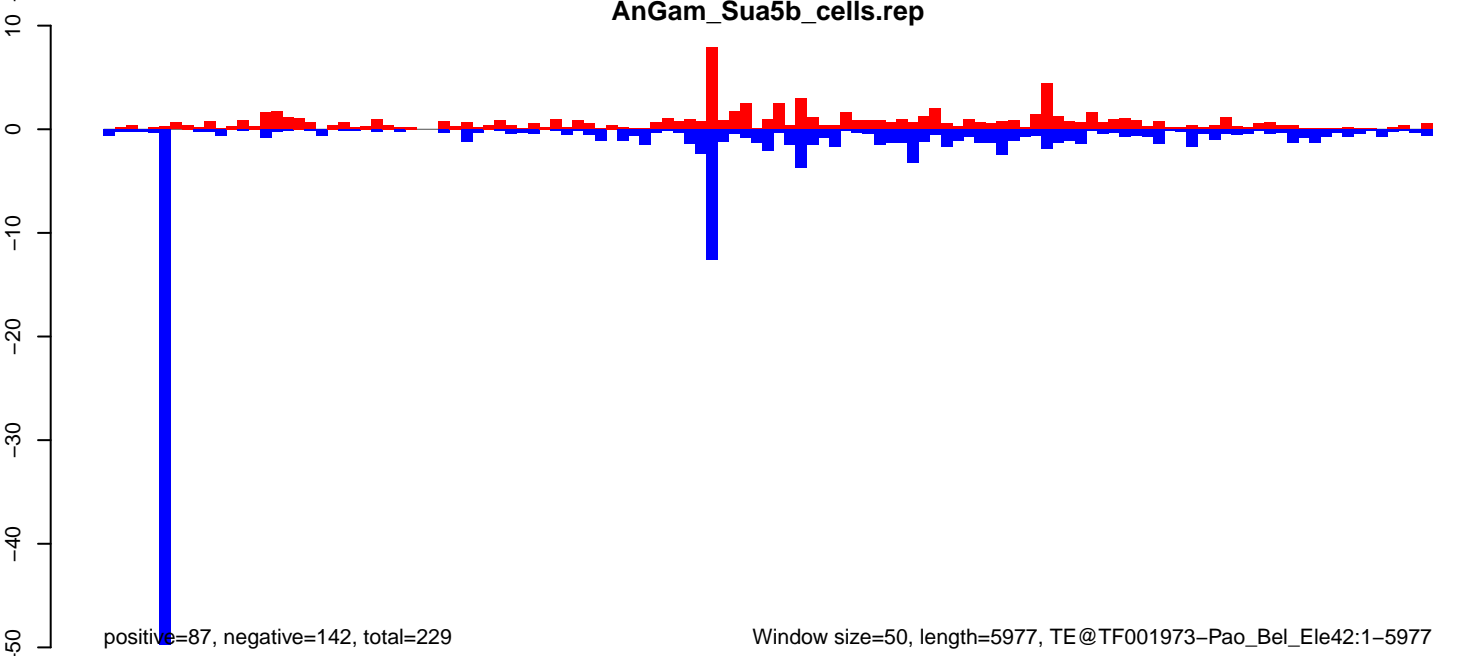
AnGam_Sua5b_cells.18_23.rep



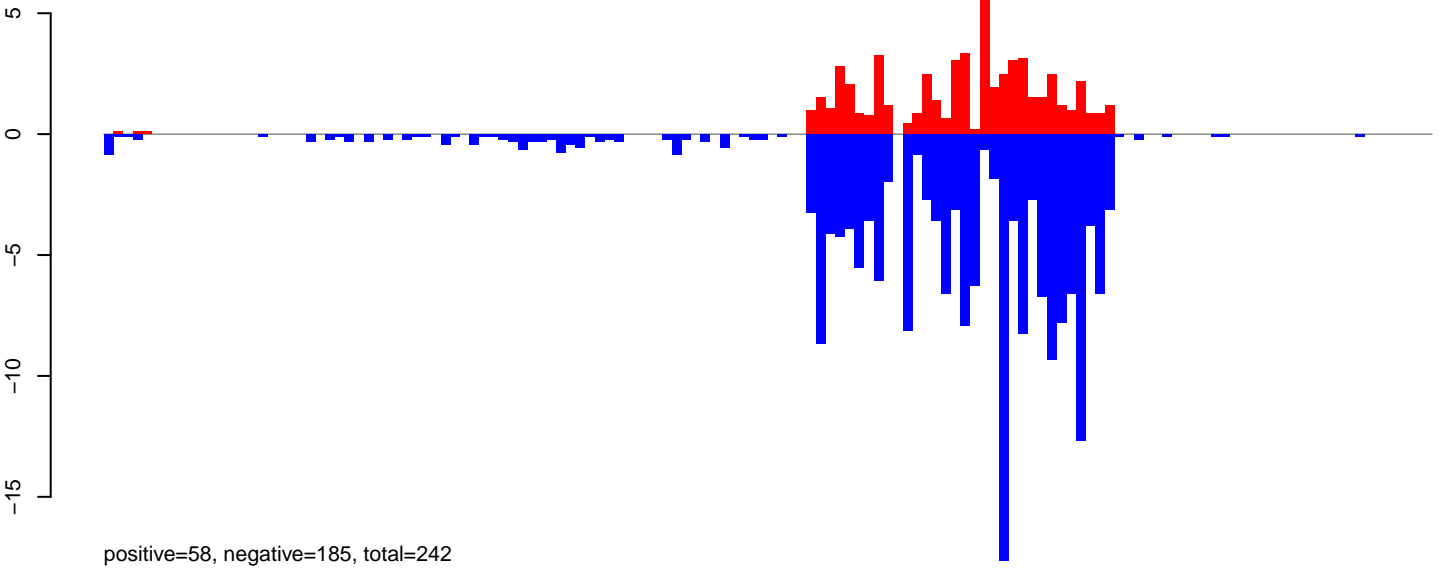
AnGam_Sua5b_cells.24_35.rep



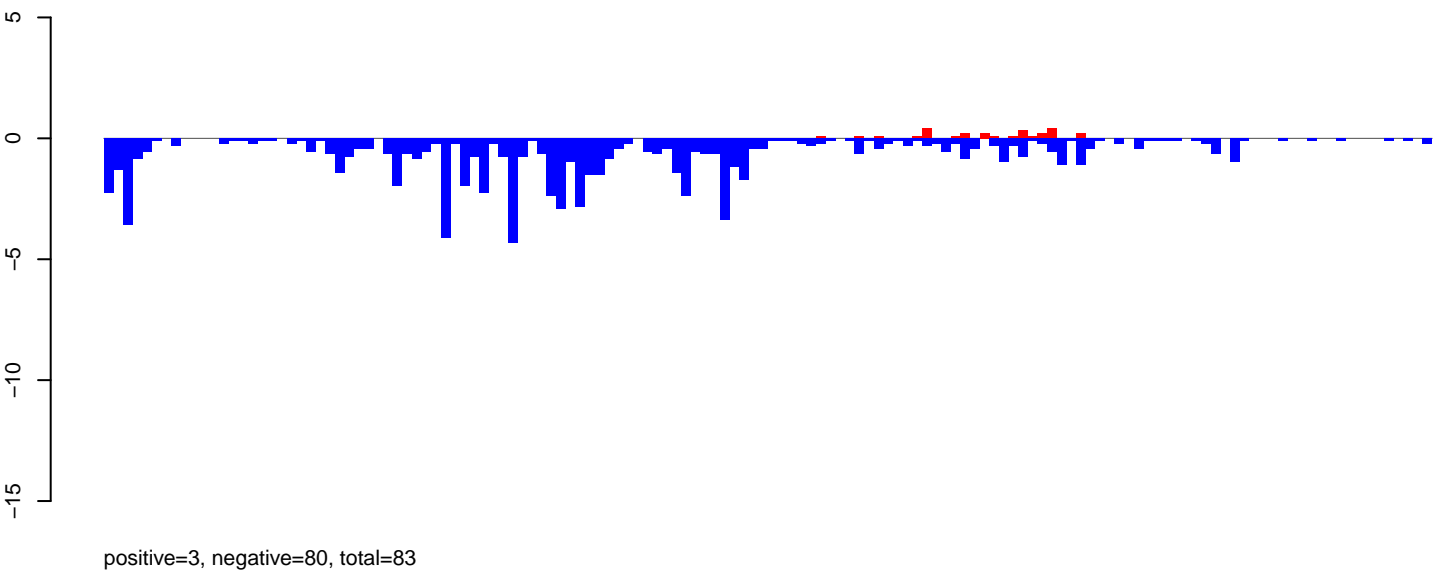
AnGam_Sua5b_cells.rep



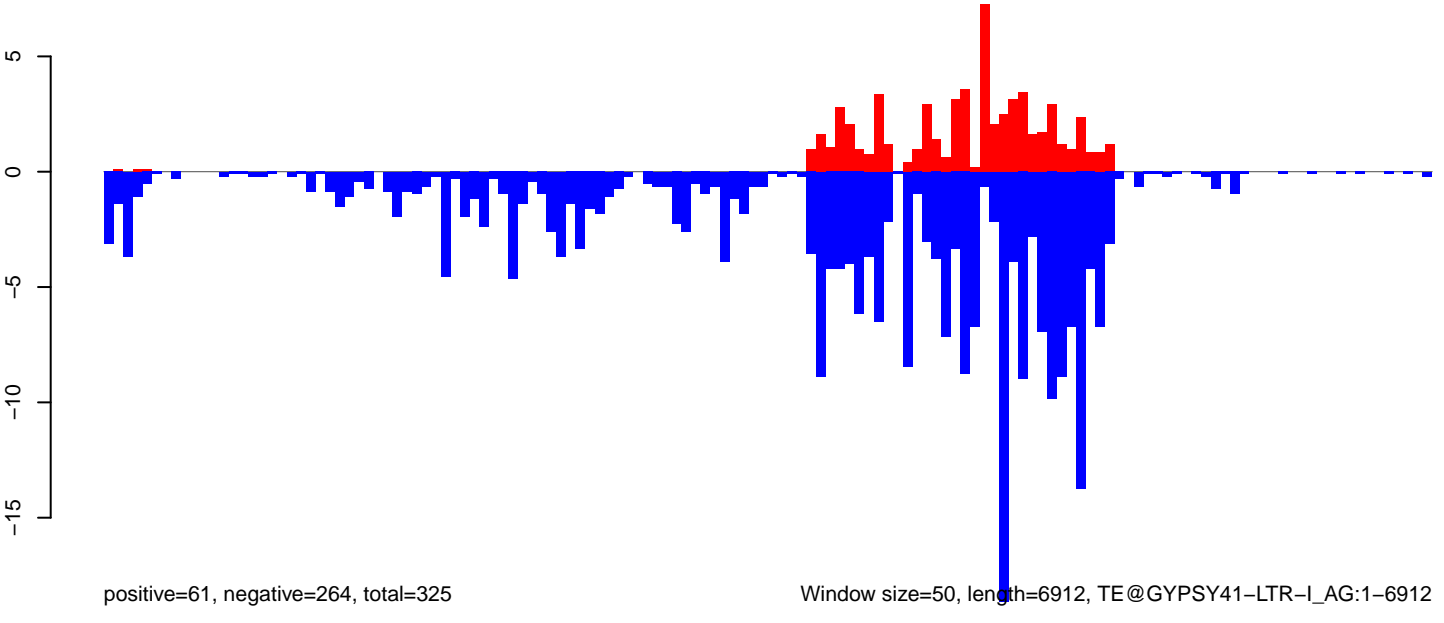
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



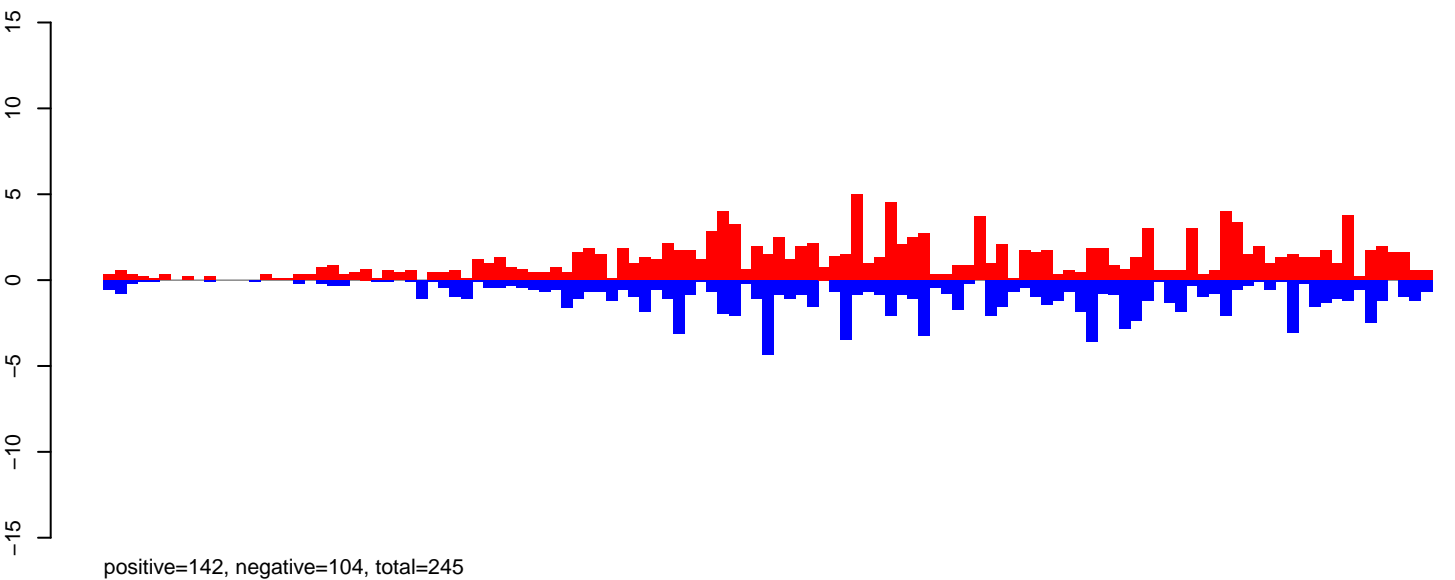
AnGam_Sua5b_cells.rep



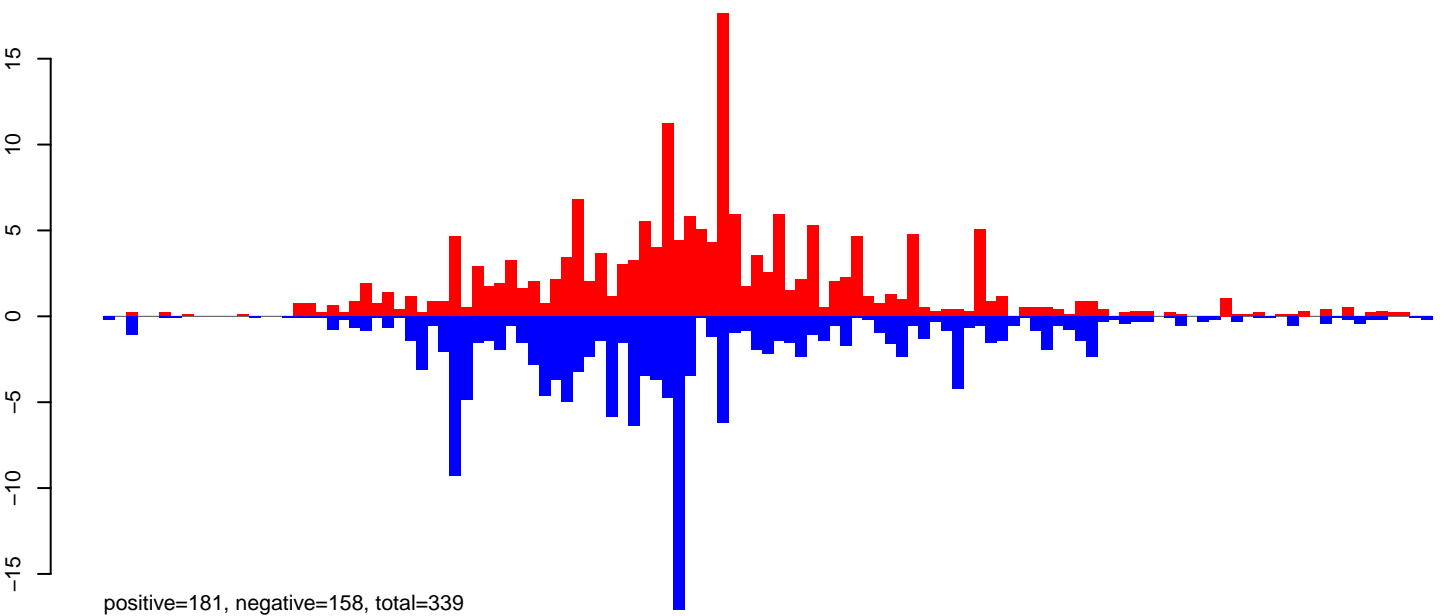
Window size=50, length=6912, TE@GYPSY41-LTR-I_AG:1-6912

0 1000 2000 3000 4000 5000 6000 7000

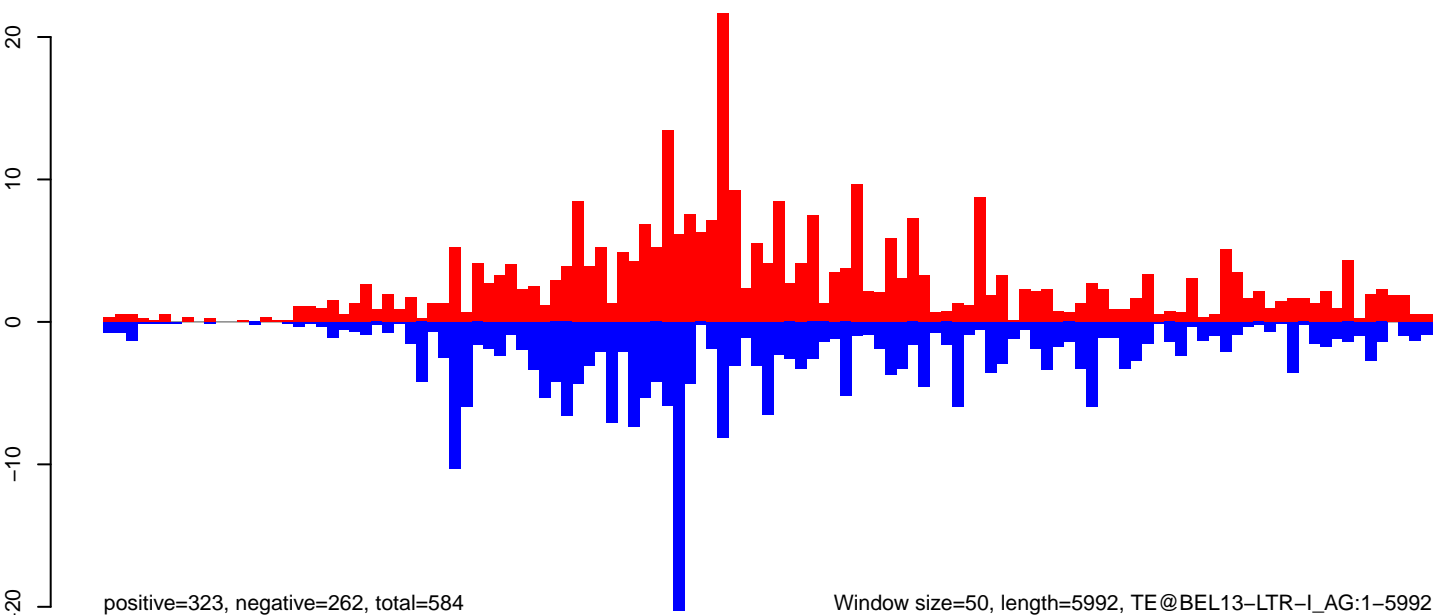
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

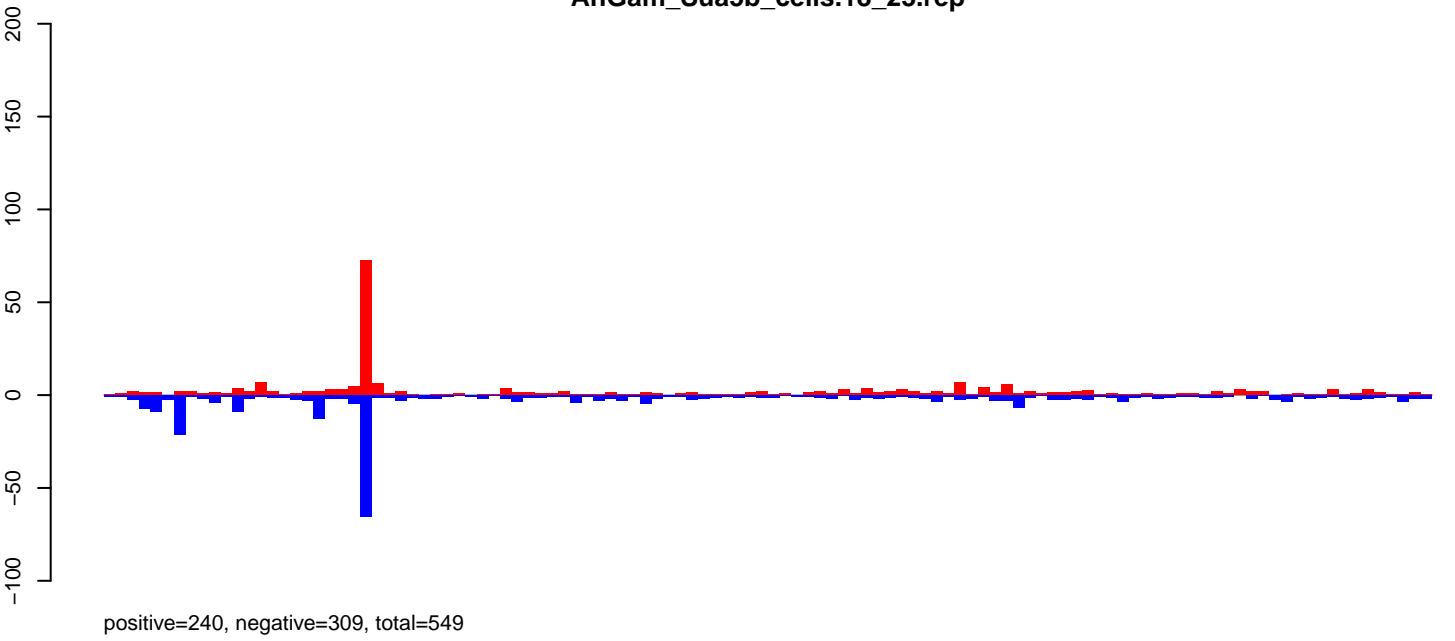


AnGam_Sua5b_cells.rep

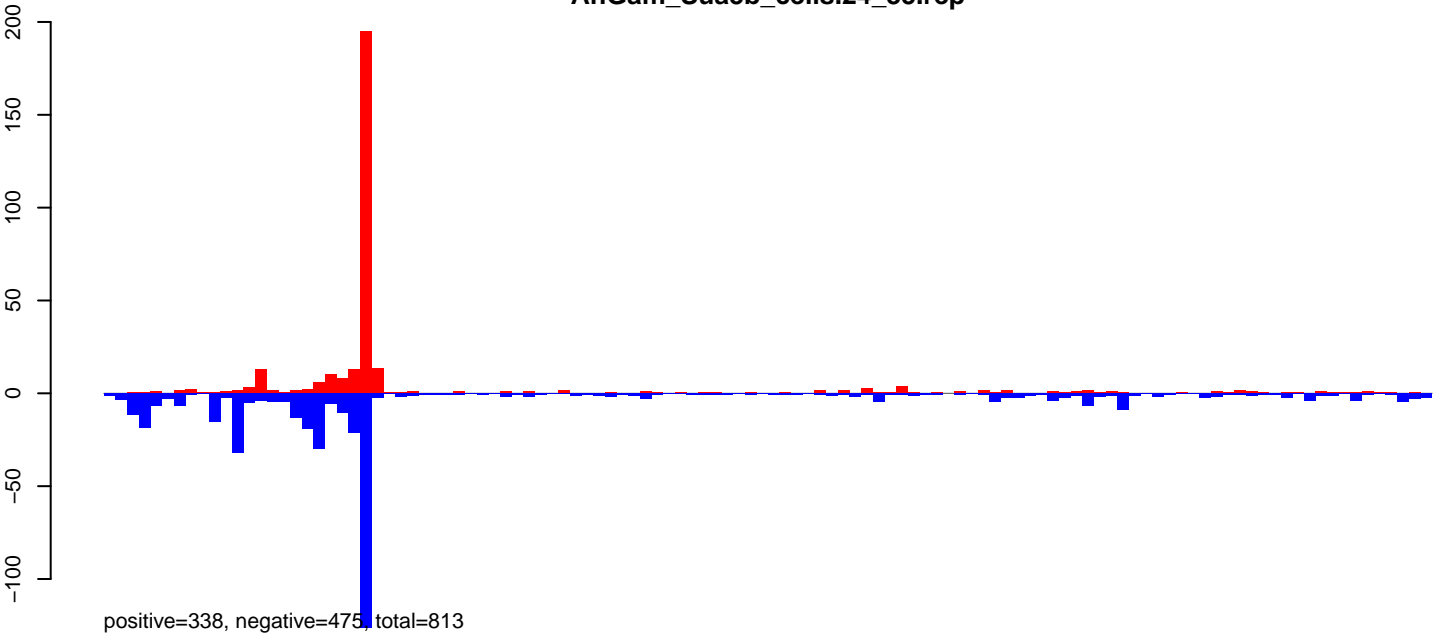


0 1000 2000 3000 4000 5000 6000

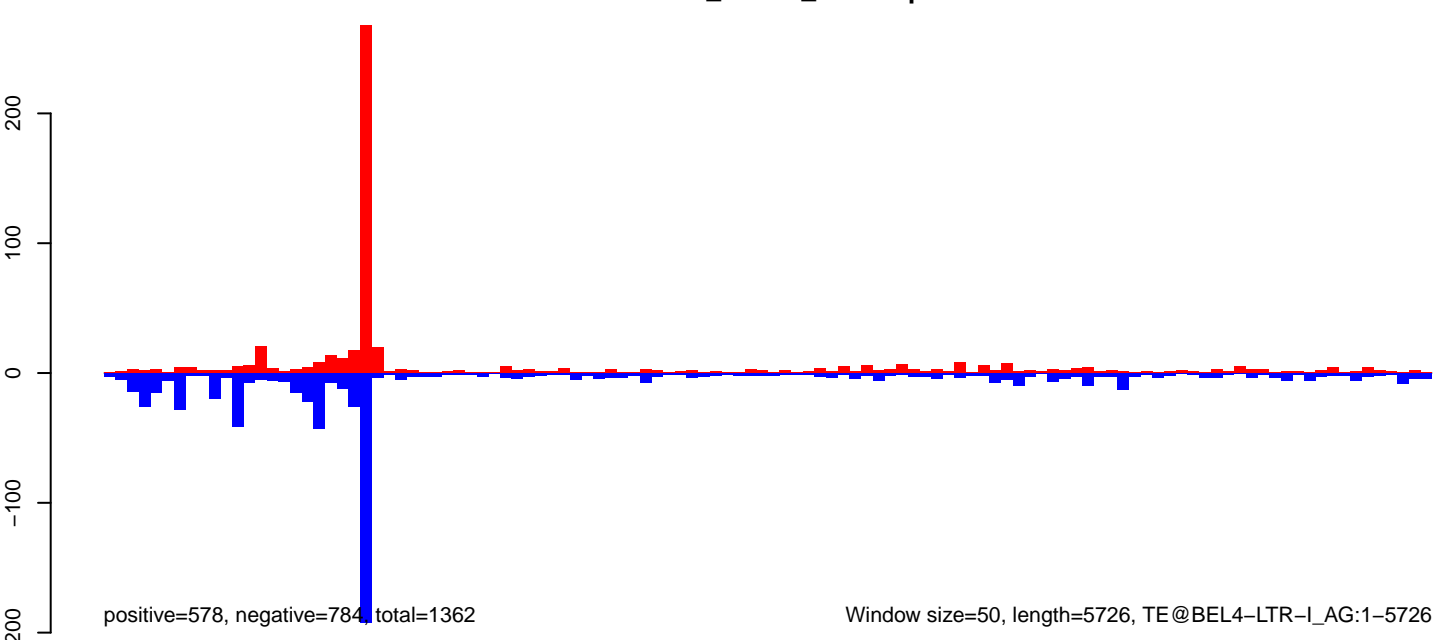
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

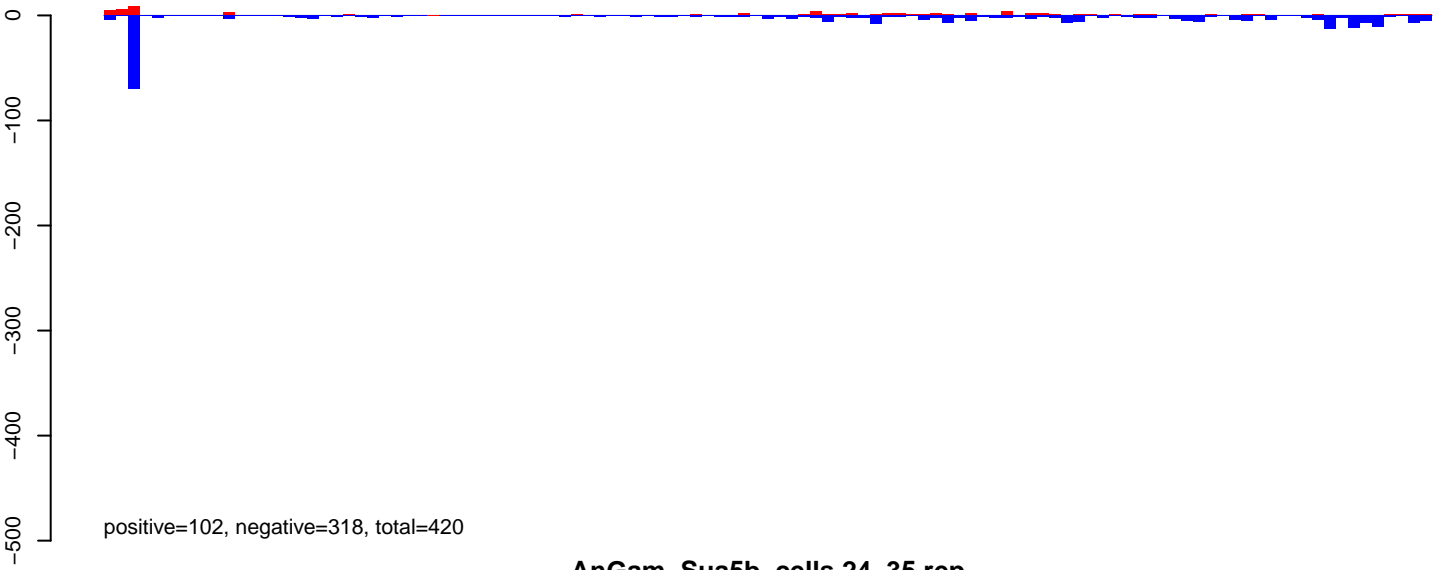


AnGam_Sua5b_cells.rep

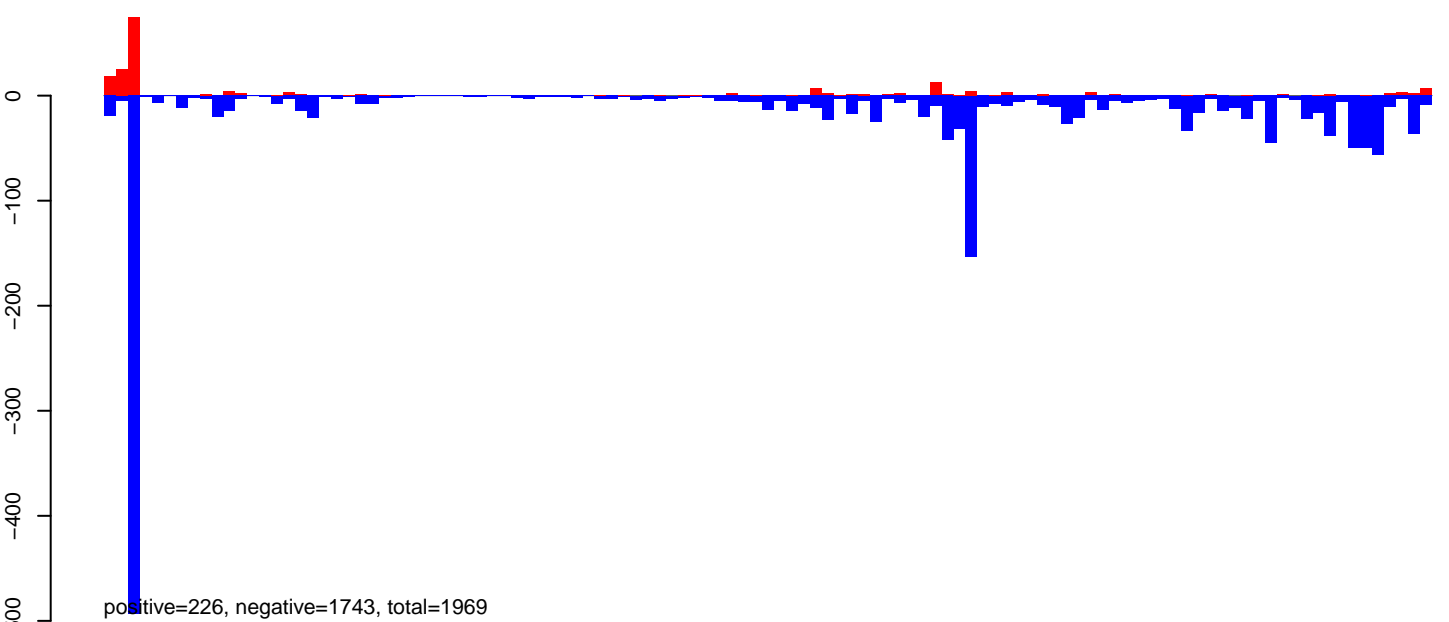


Window size=50, length=5726, TE@BEL4-LTR-I_AG:1-5726

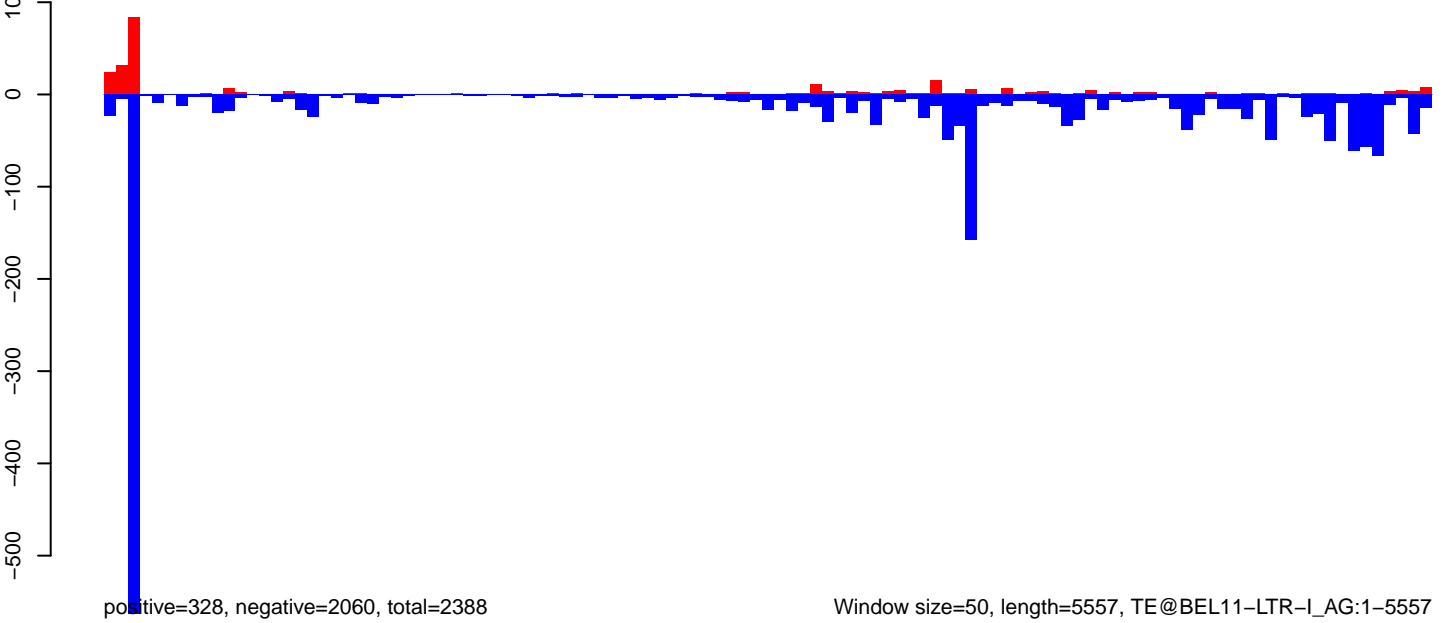
AnGam_Sua5b_cells.18_23.rep



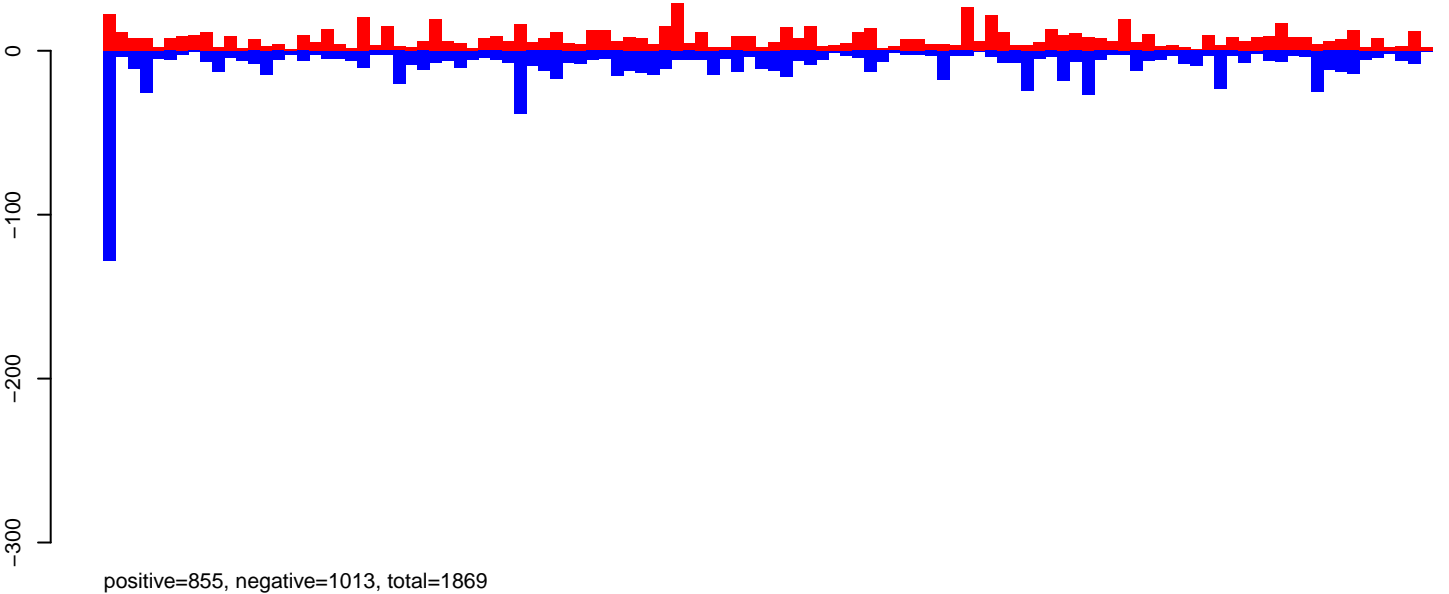
AnGam_Sua5b_cells.24_35.rep



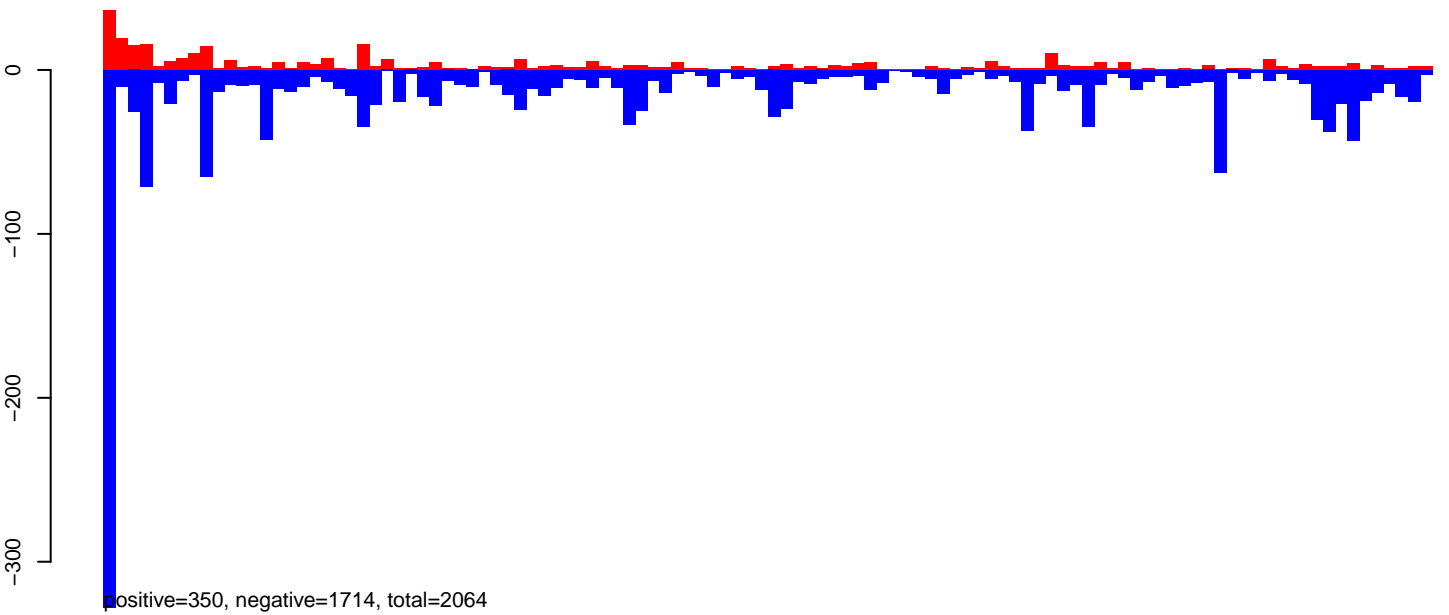
AnGam_Sua5b_cells.rep



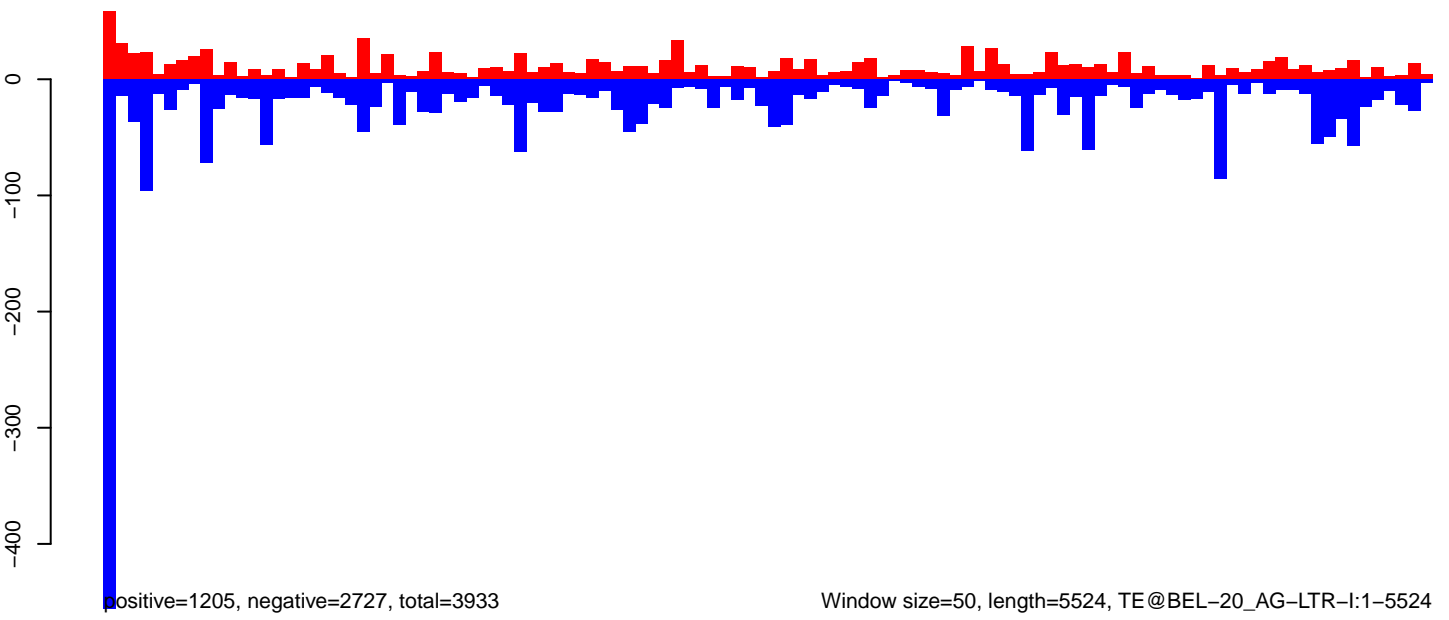
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

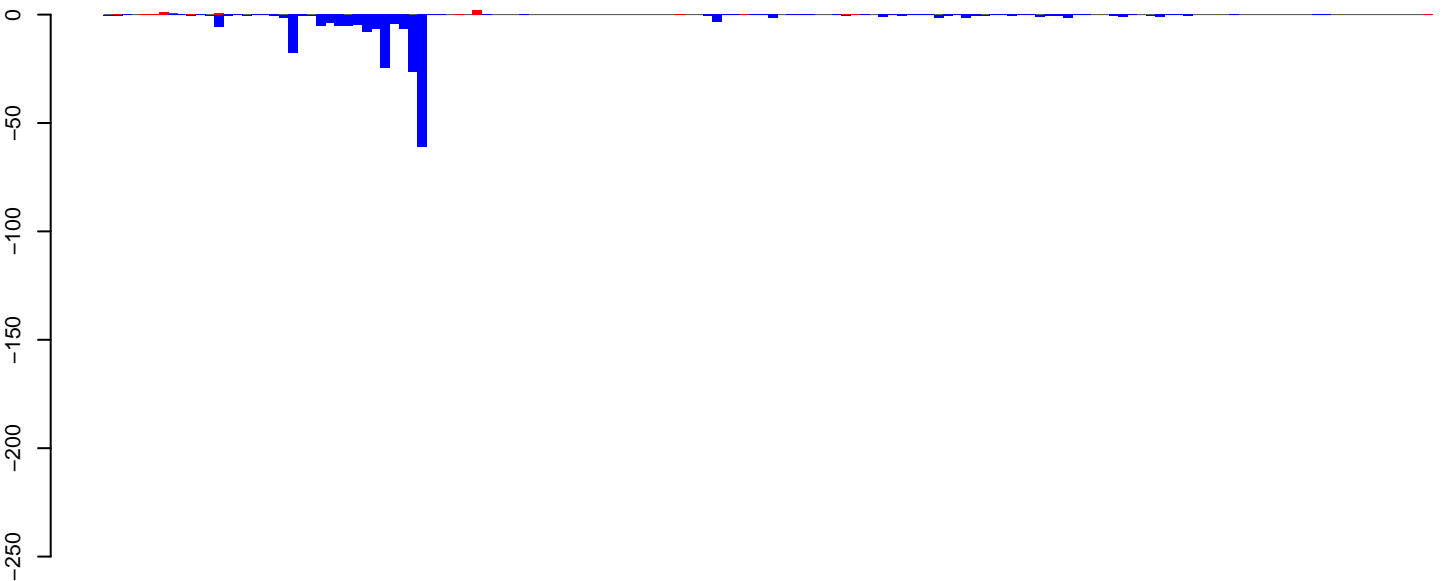


AnGam_Sua5b_cells.rep

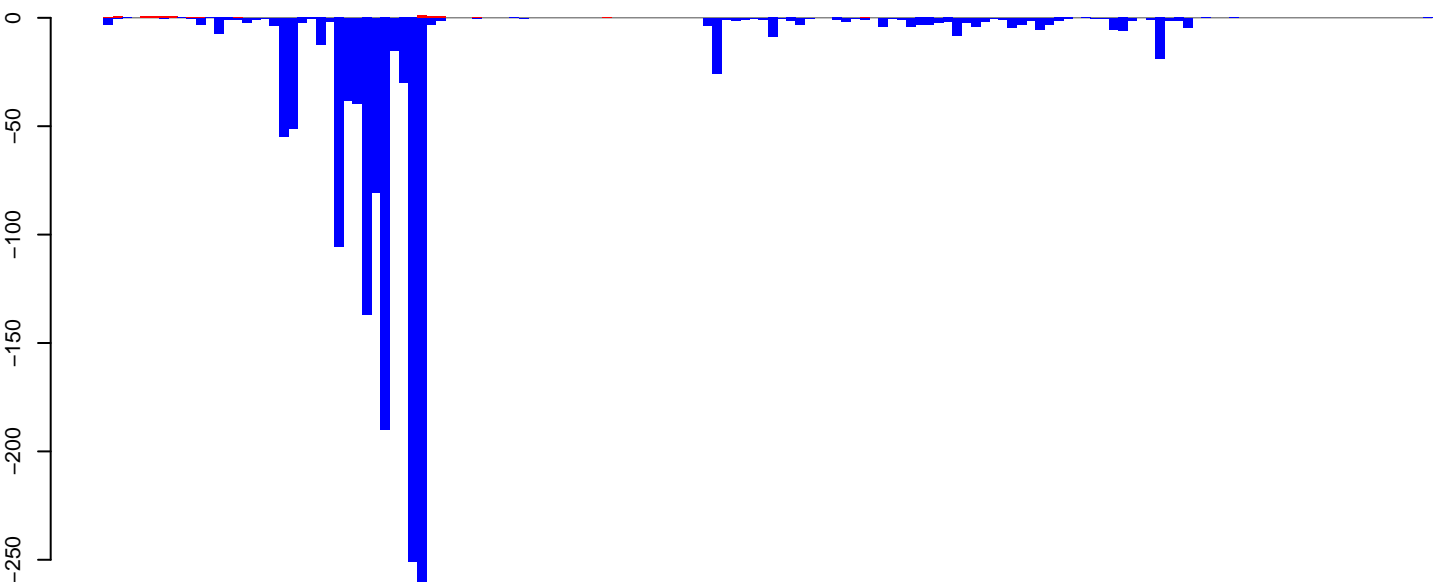


Window size=50, length=5524, TE@BEL-20_AG-LTR-I:1-5524

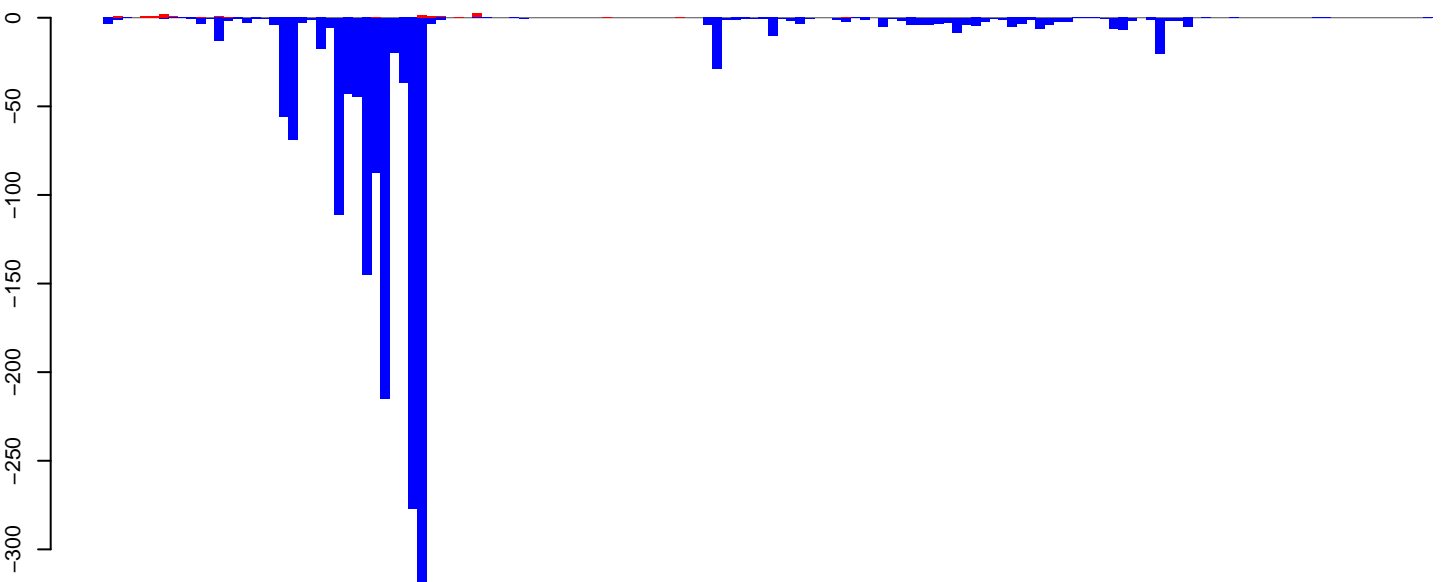
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



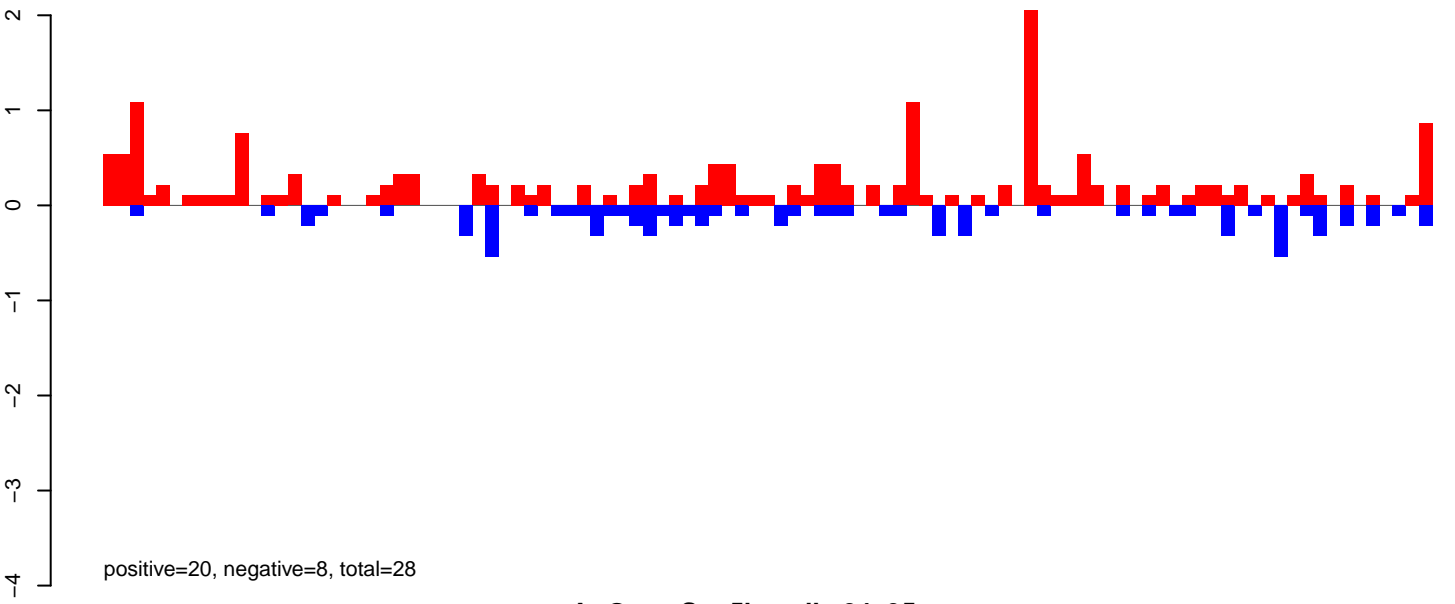
AnGam_Sua5b_cells.rep



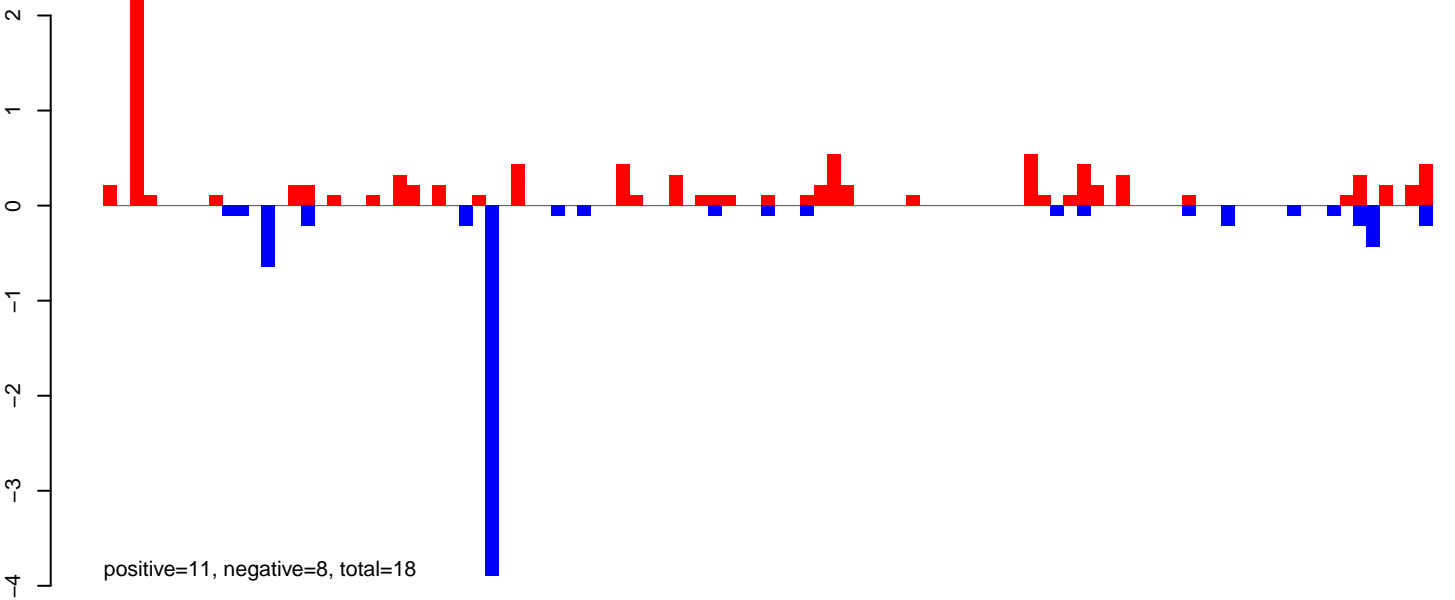
Window size=50, length=7224, TE@GYPSY10-LTR-I_AG:1-7224

0 1000 2000 3000 4000 5000 6000 7000

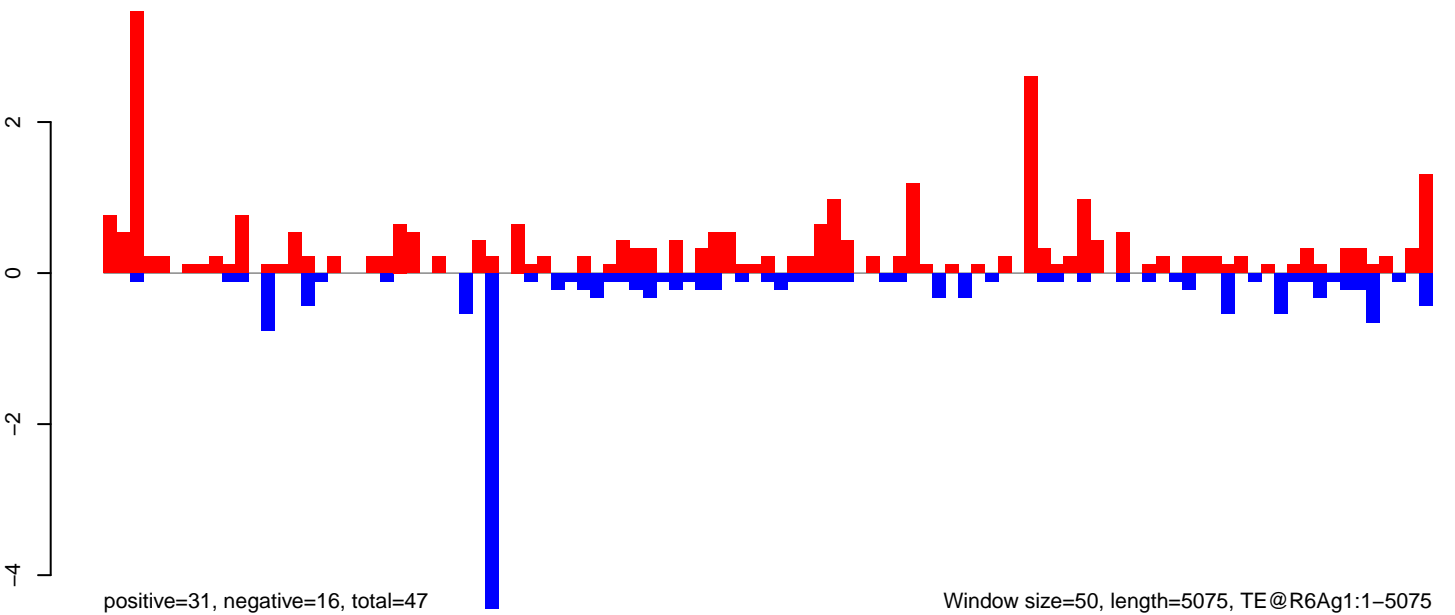
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

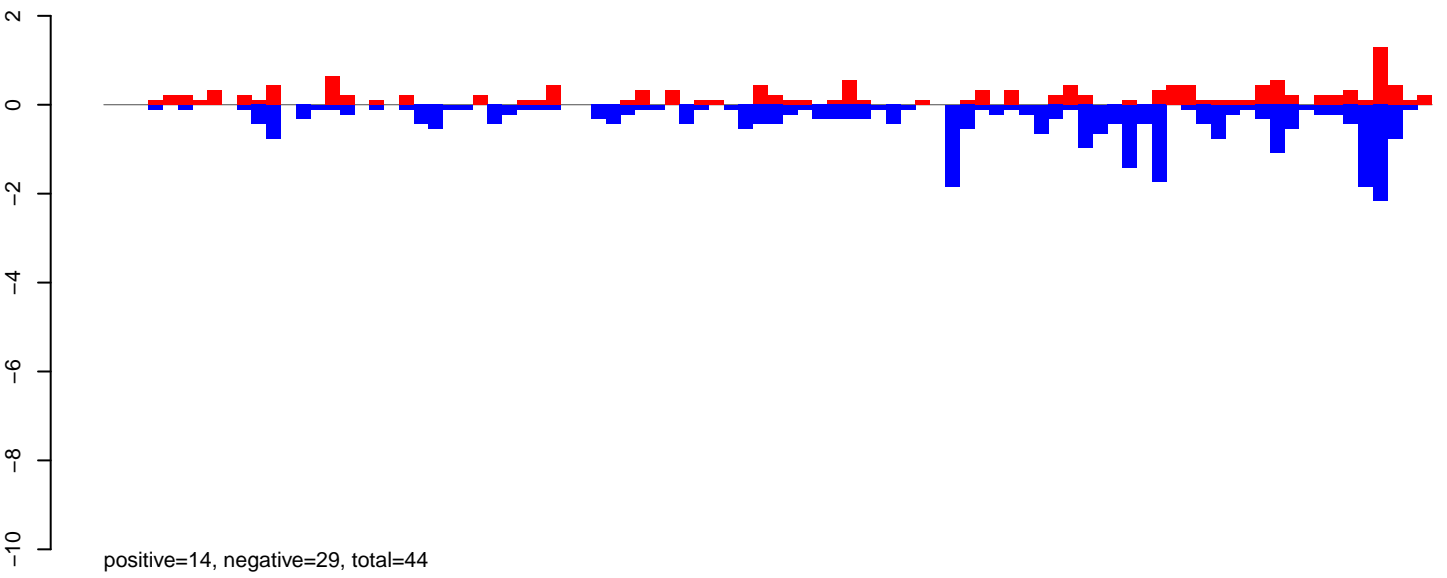


AnGam_Sua5b_cells.rep

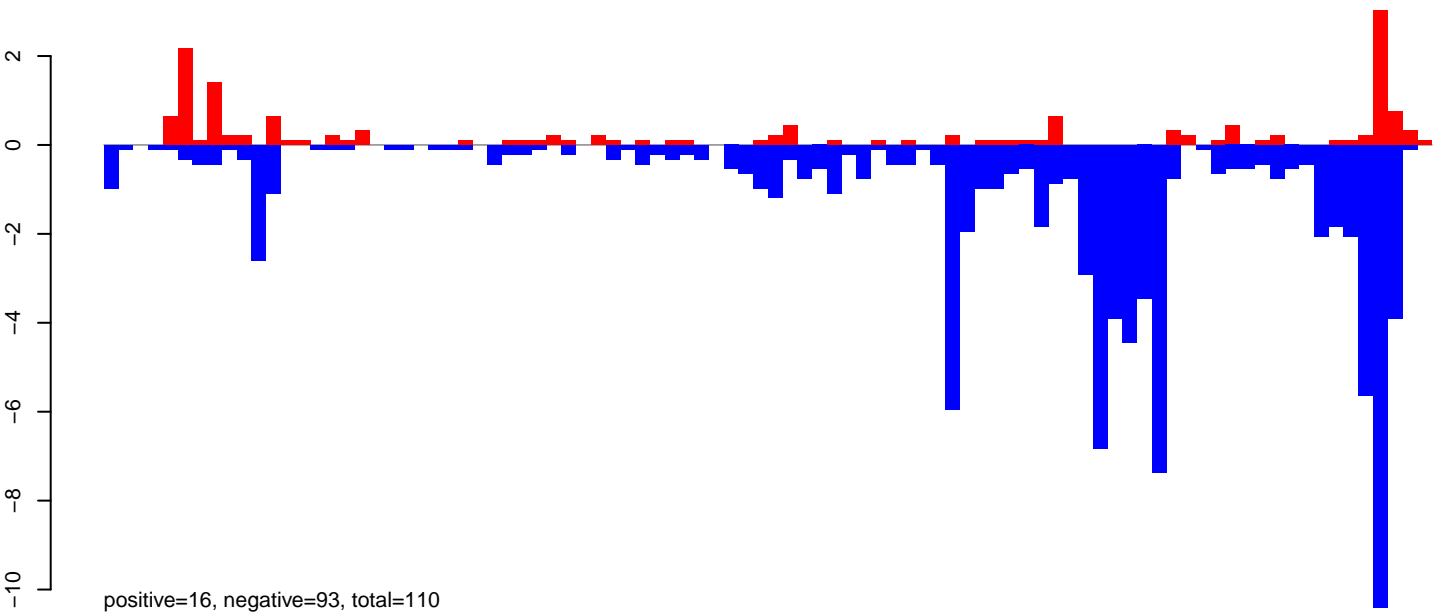


Window size=50, length=5075, TE@R6Ag1:1-5075

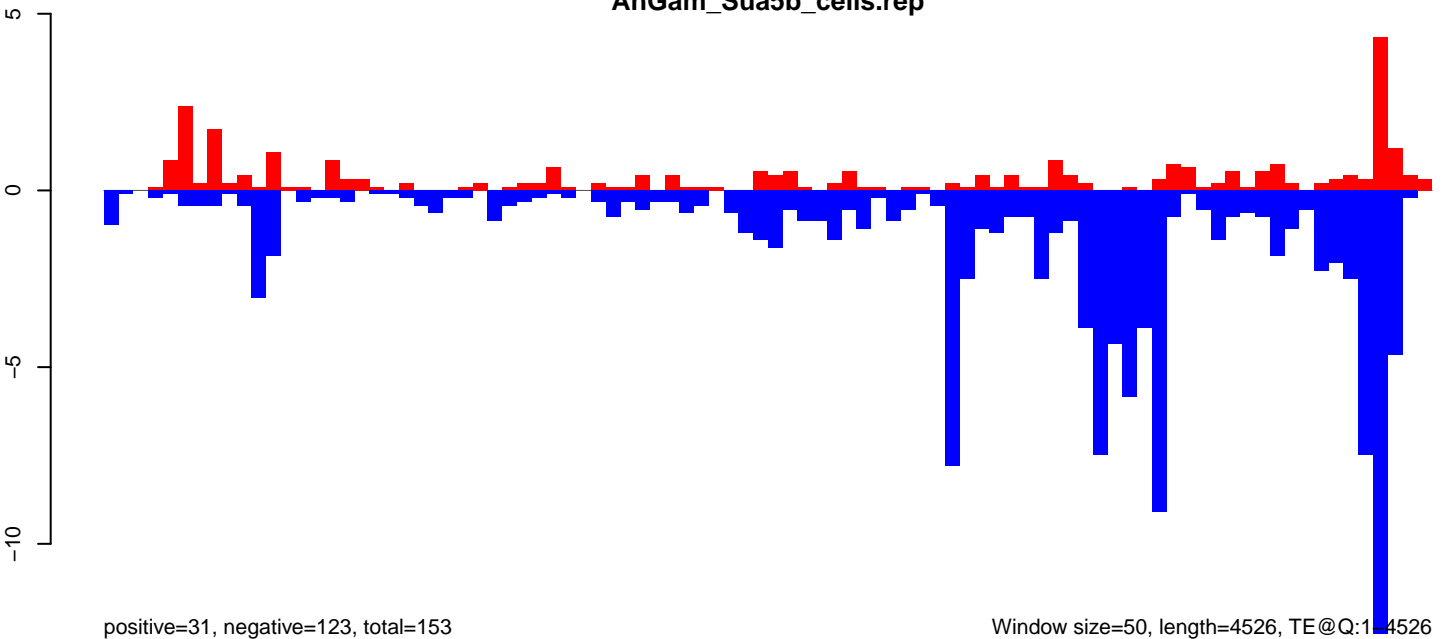
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



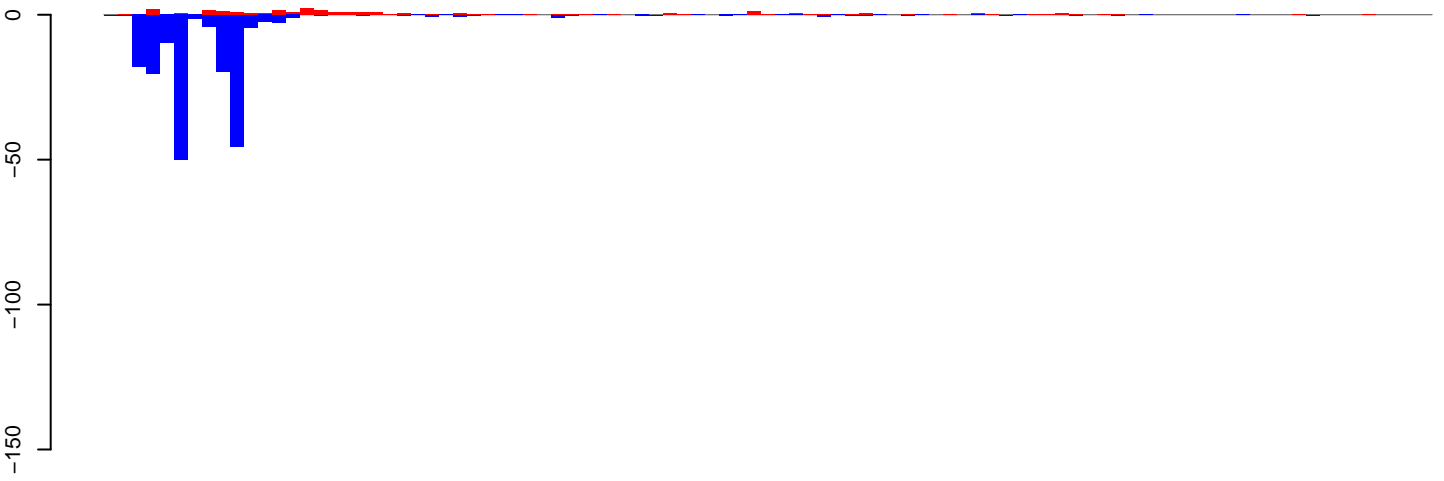
AnGam_Sua5b_cells.rep



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

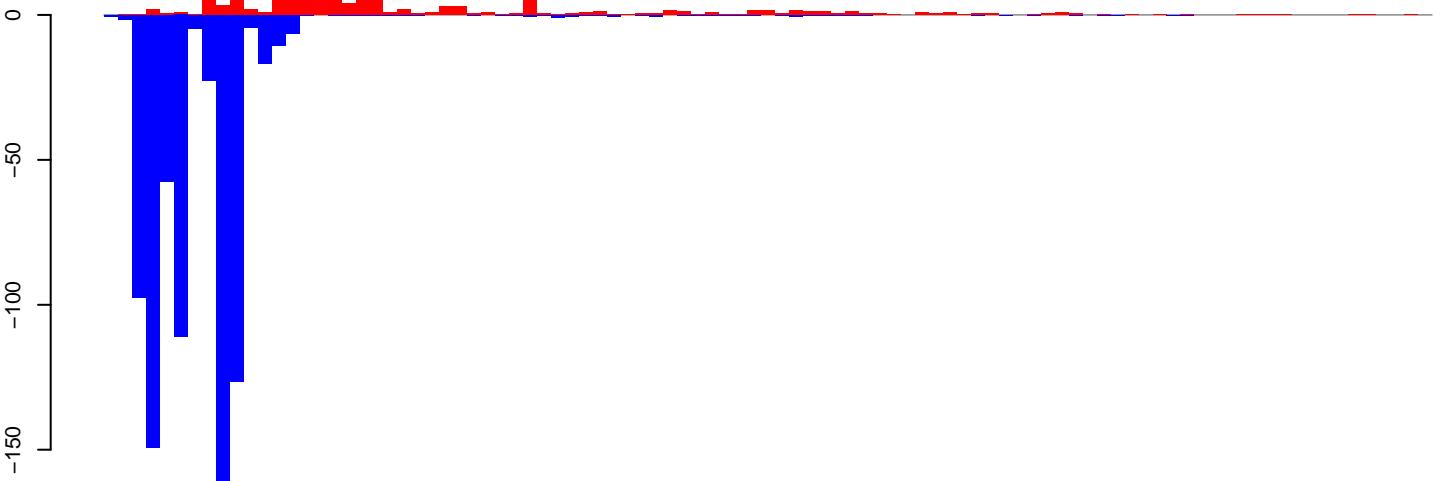
0 1000 2000 3000 4000

AnGam_Sua5b_cells.18_23.rep



positive=29, negative=188, total=217

AnGam_Sua5b_cells.24_35.rep



positive=154, negative=805, total=959

AnGam_Sua5b_cells.rep

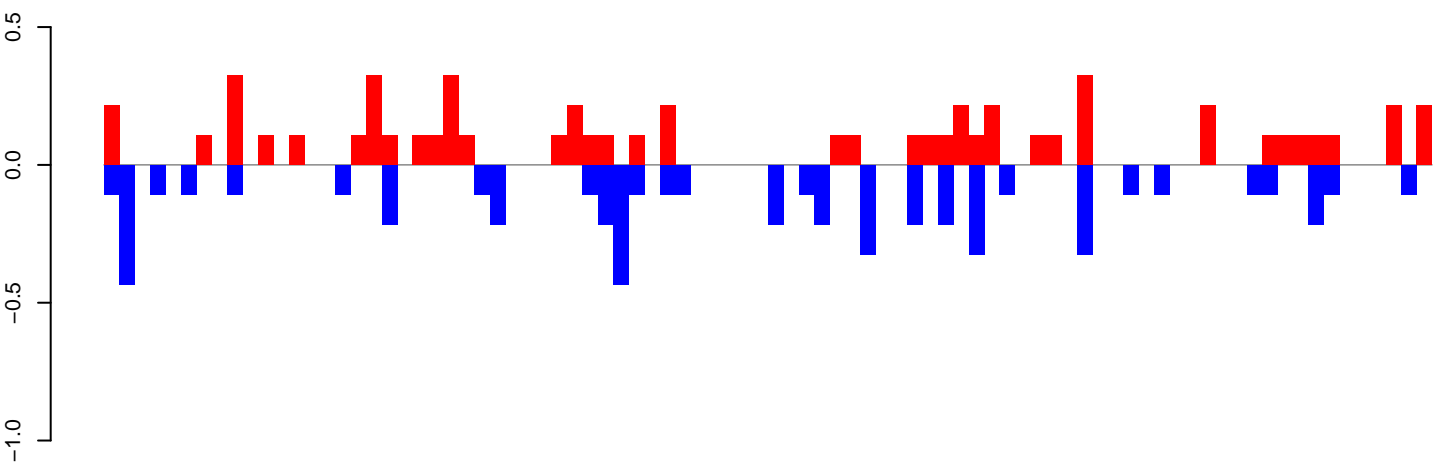


positive=183, negative=993, total=1176

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

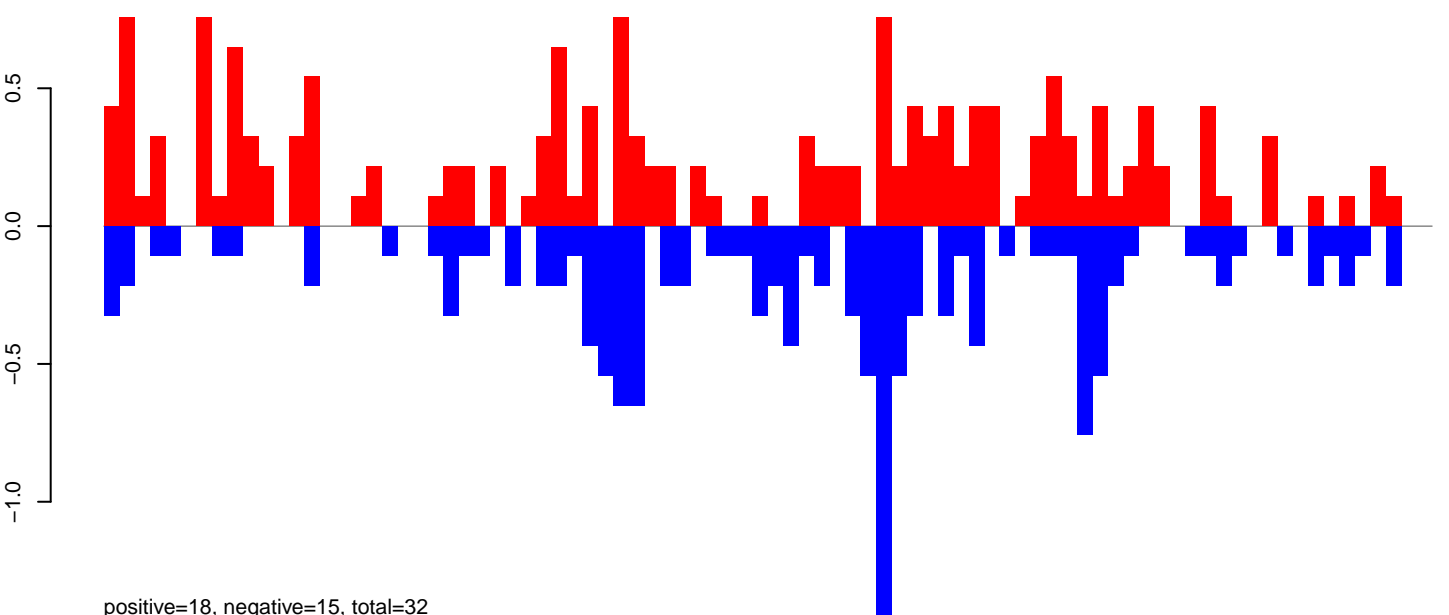
0 1000 2000 3000 4000

AnGam_Sua5b_cells.18_23.rep



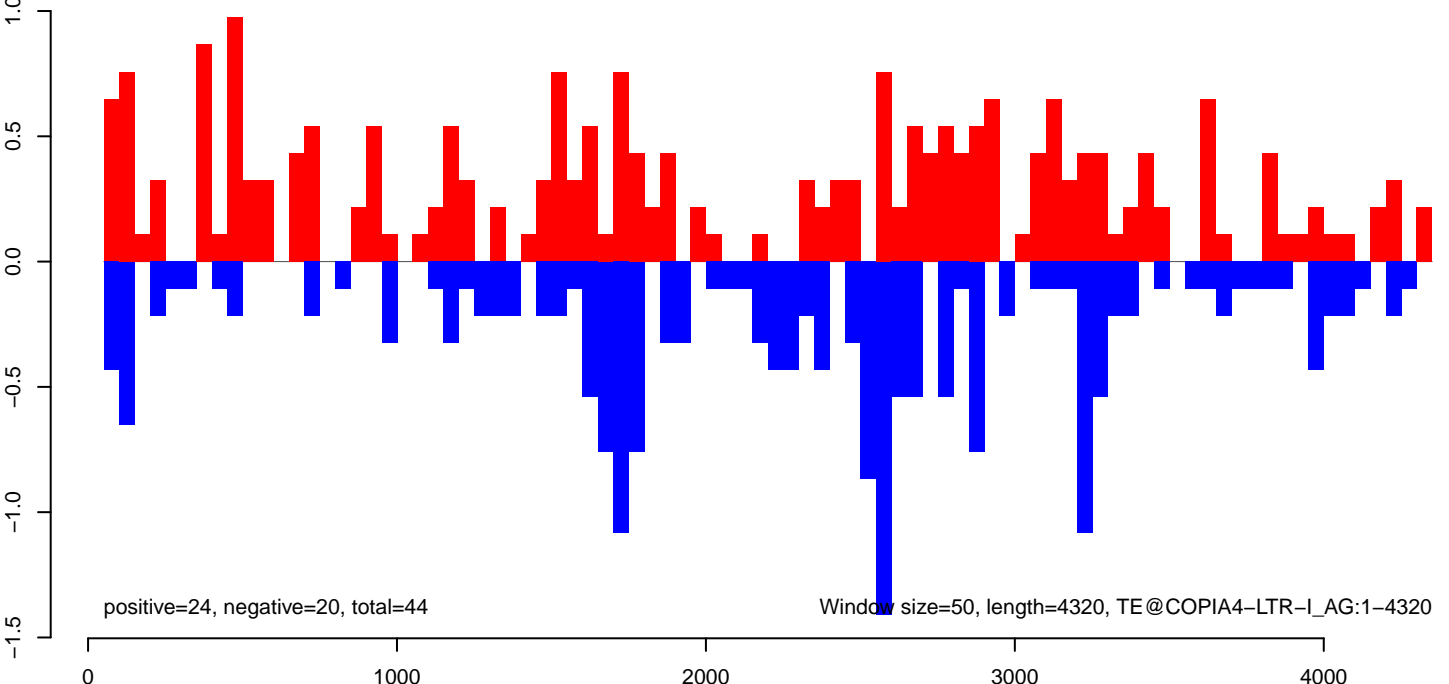
positive=6, negative=6, total=11

AnGam_Sua5b_cells.24_35.rep



positive=18, negative=15, total=32

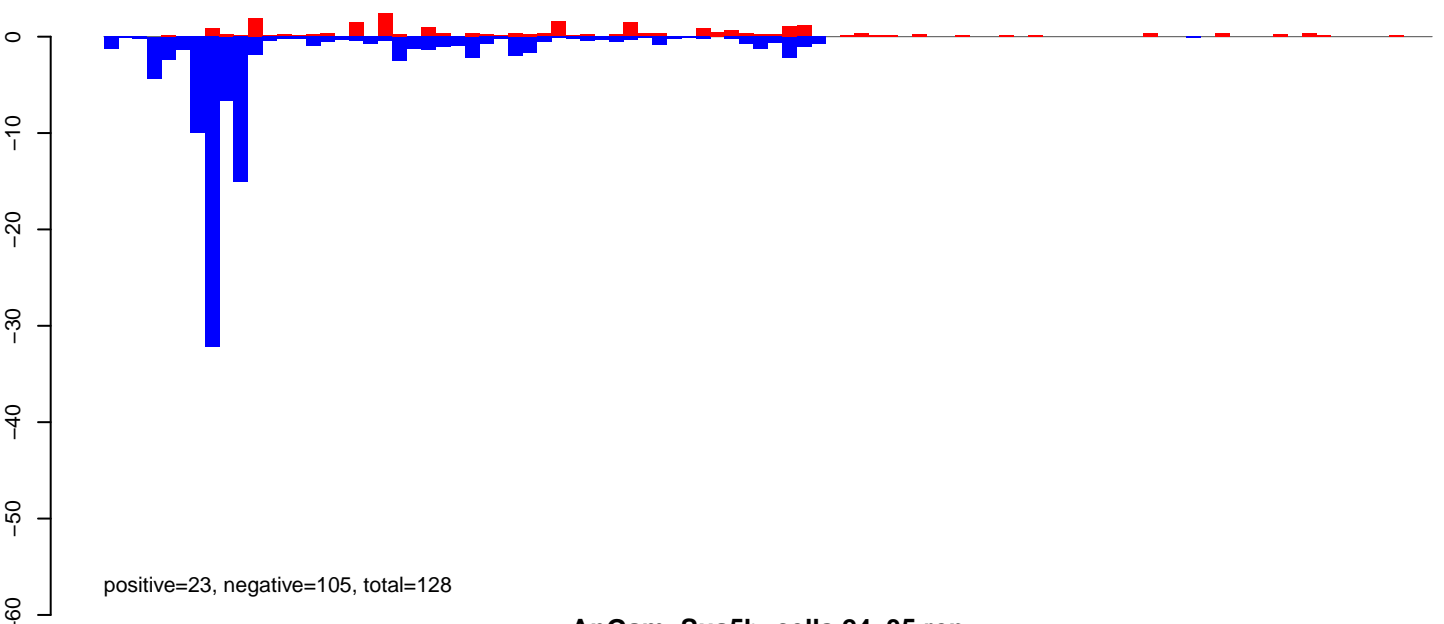
AnGam_Sua5b_cells.rep



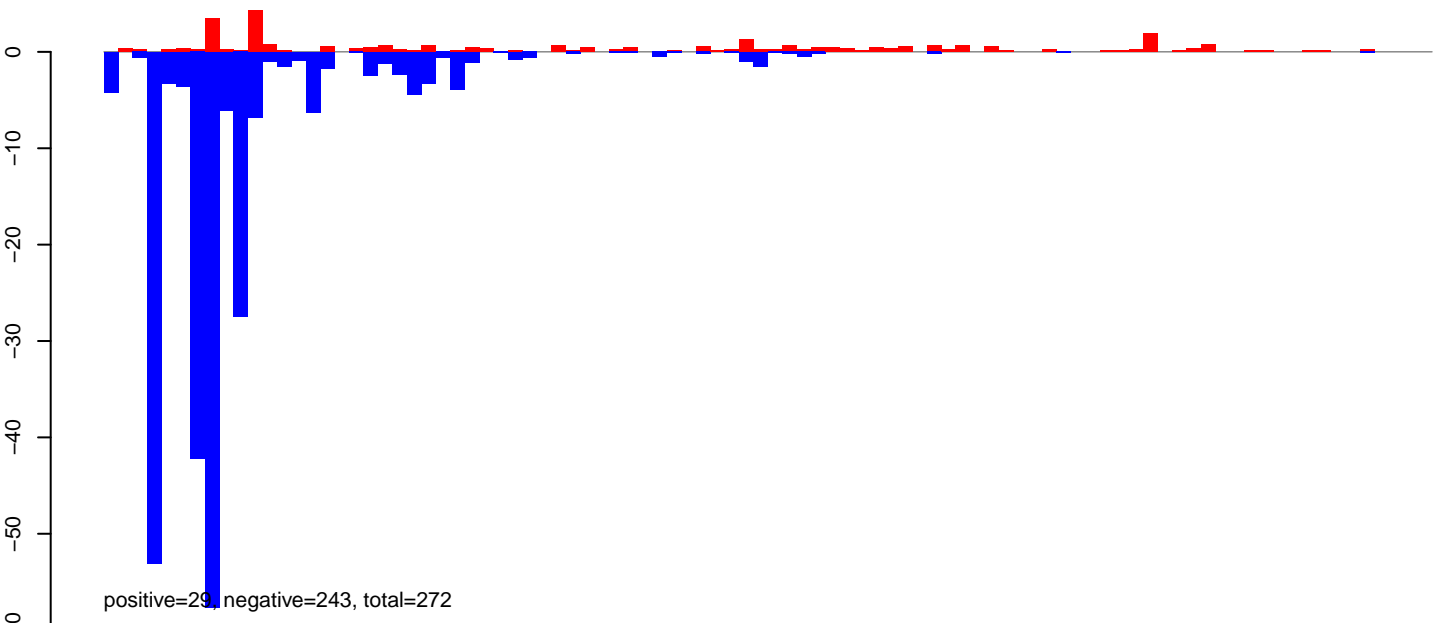
positive=24, negative=20, total=44

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

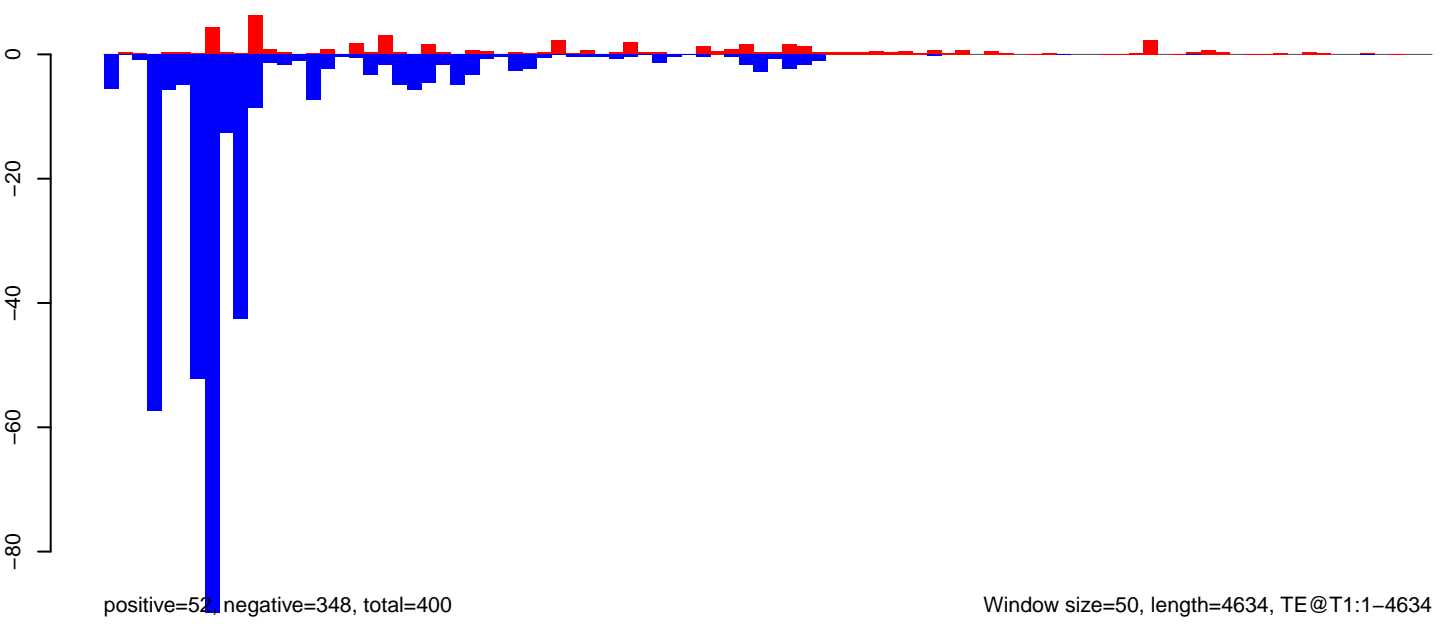
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



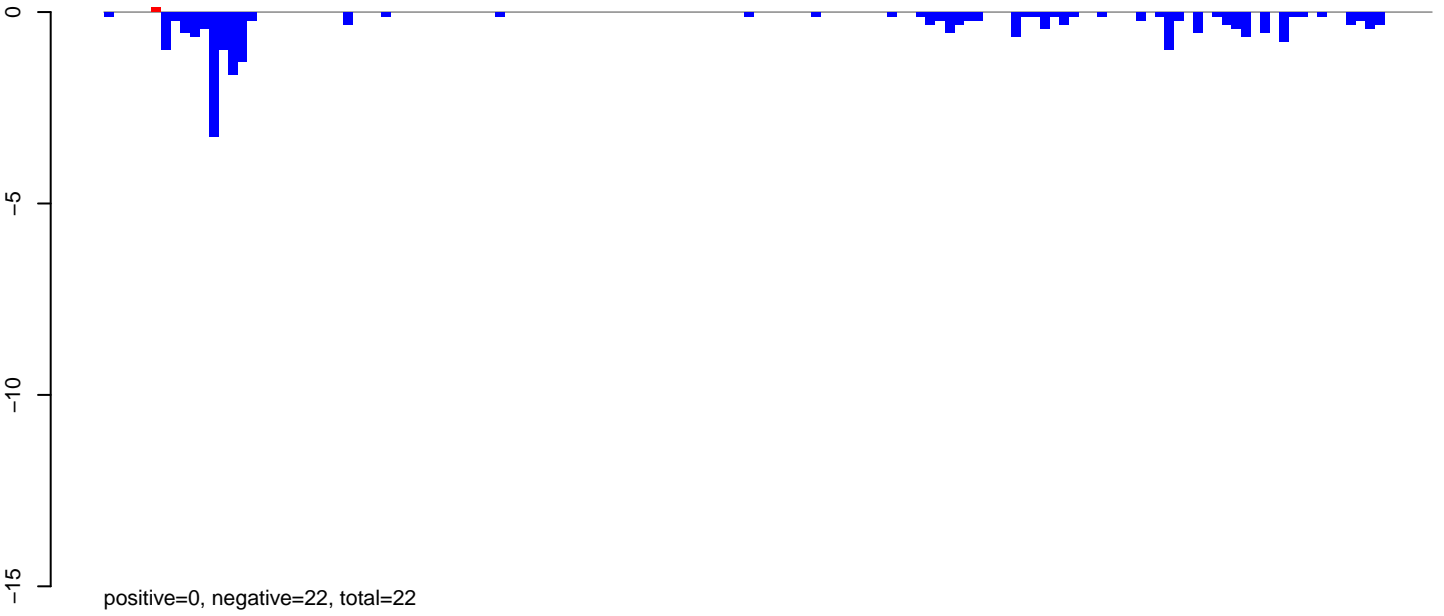
AnGam_Sua5b_cells.rep



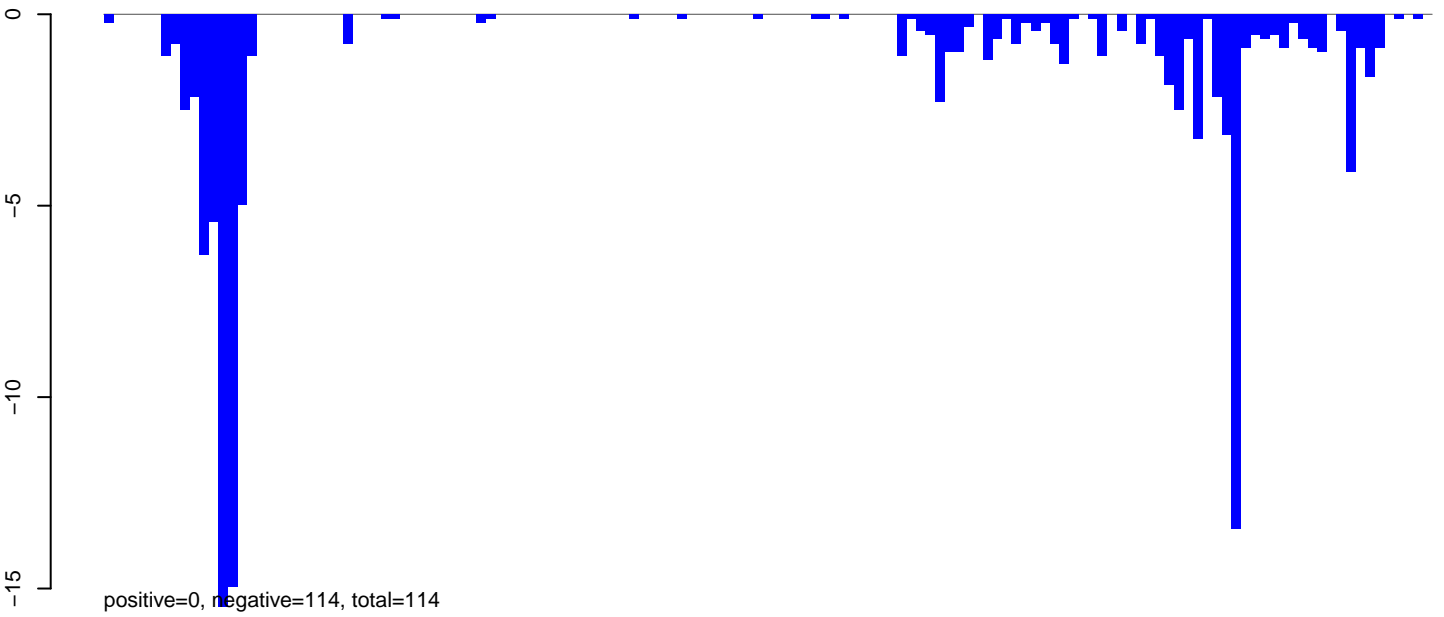
Window size=50, length=4634, TE@T1:1-4634

0 1000 2000 3000 4000

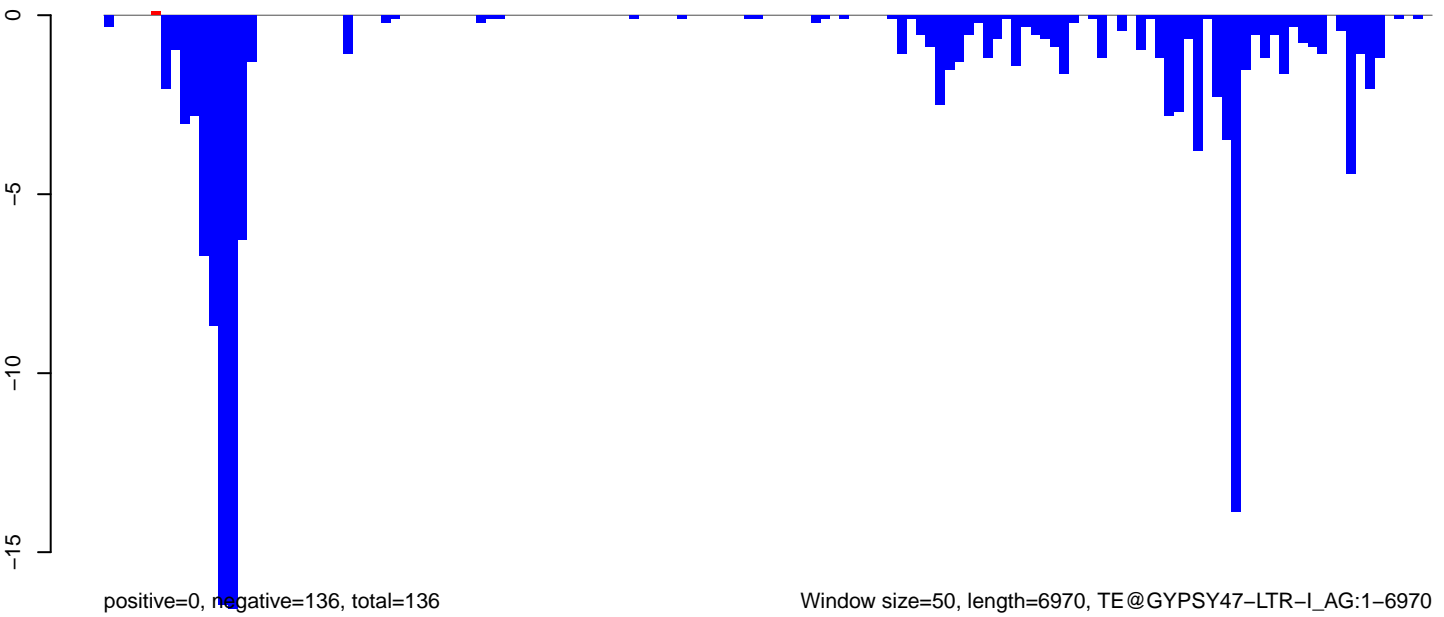
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



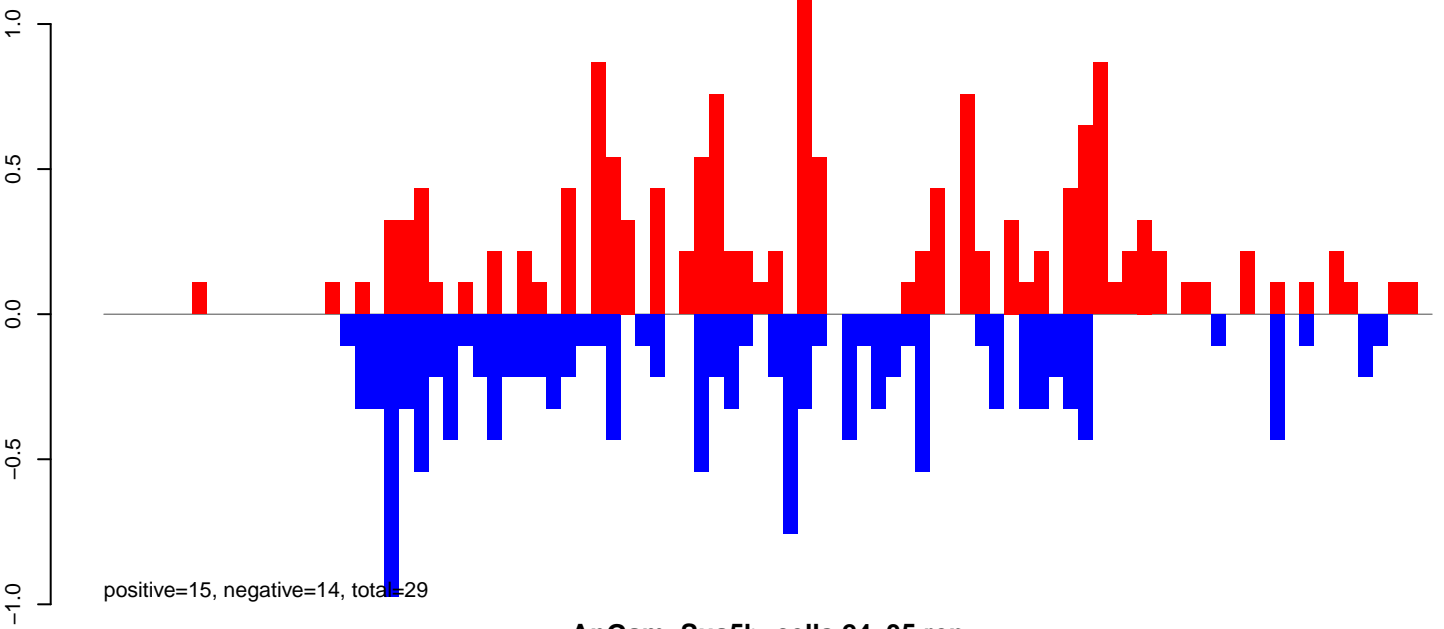
AnGam_Sua5b_cells.rep



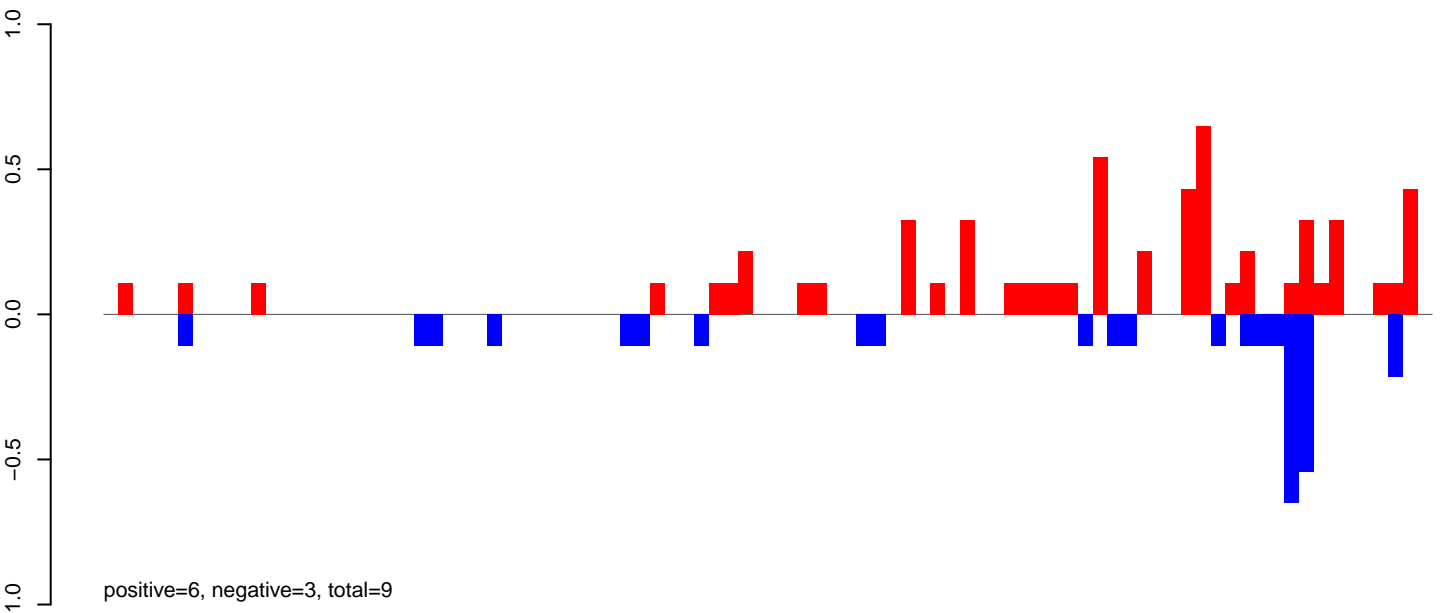
Window size=50, length=6970, TE@GYPSY47-LTR-I_AG:1-6970

0 1000 2000 3000 4000 5000 6000 7000

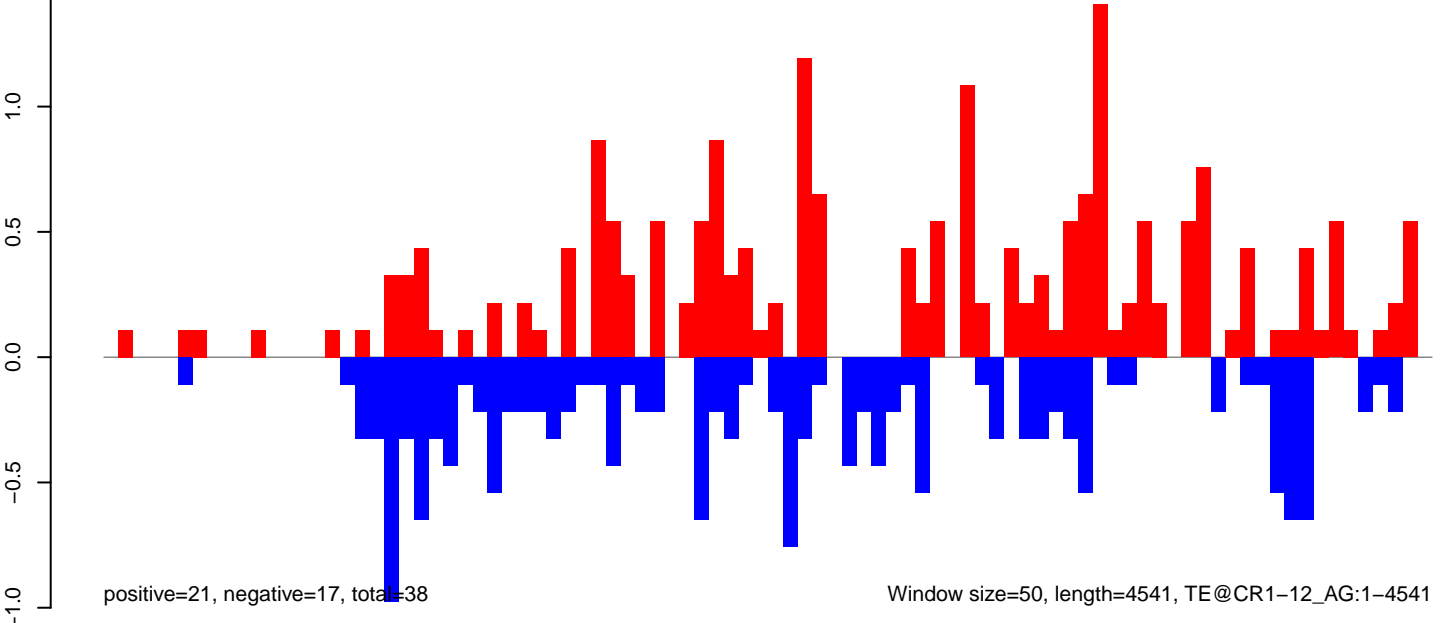
AnGam_Sua5b_cells.18_23.rep



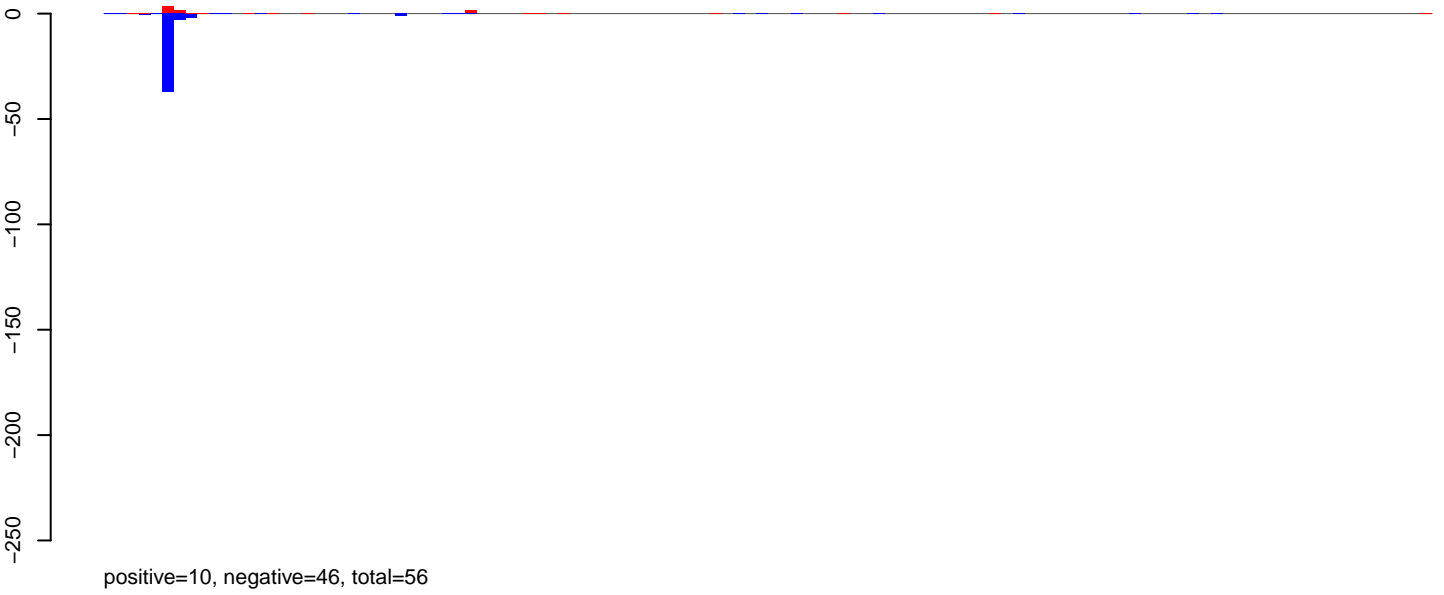
AnGam_Sua5b_cells.24_35.rep



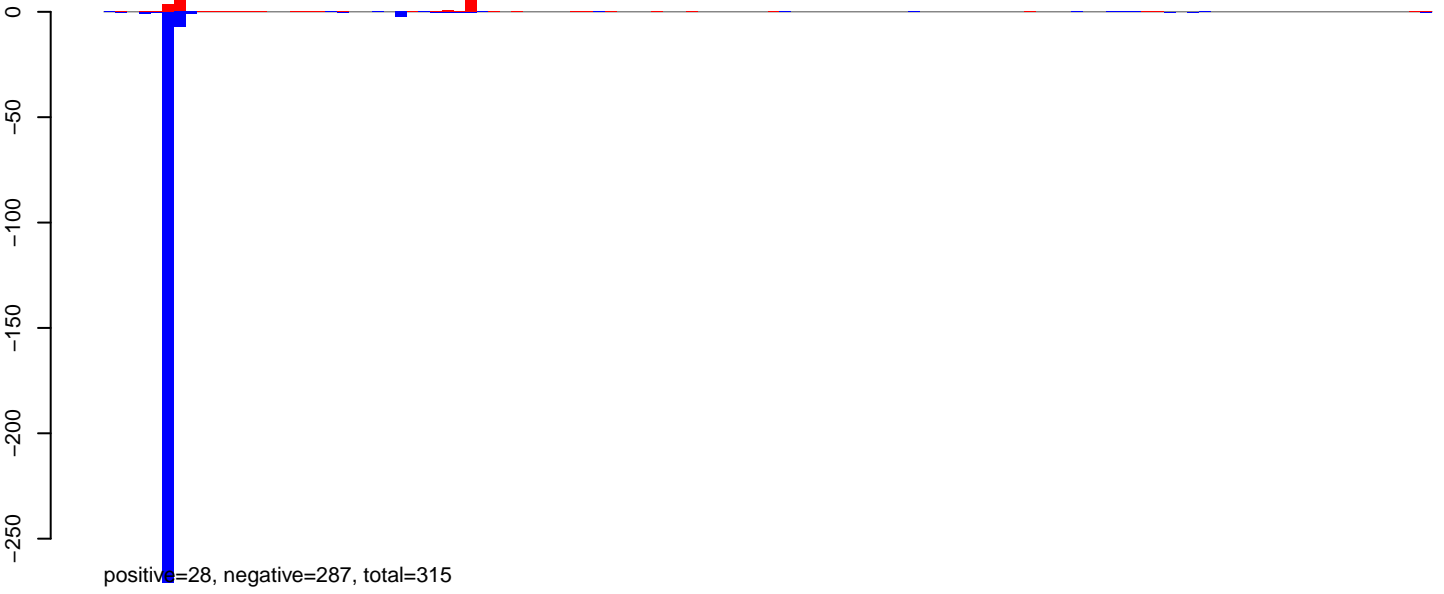
AnGam_Sua5b_cells.rep



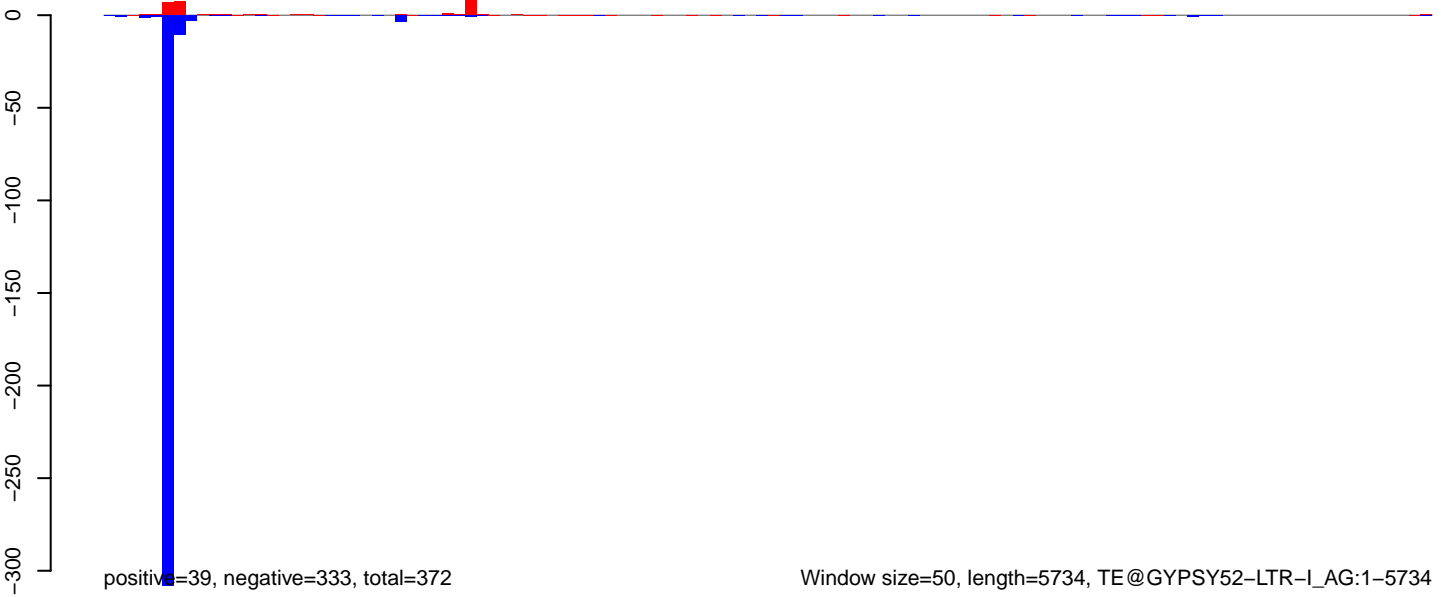
AnGam_Sua5b_cells.18_23.rep



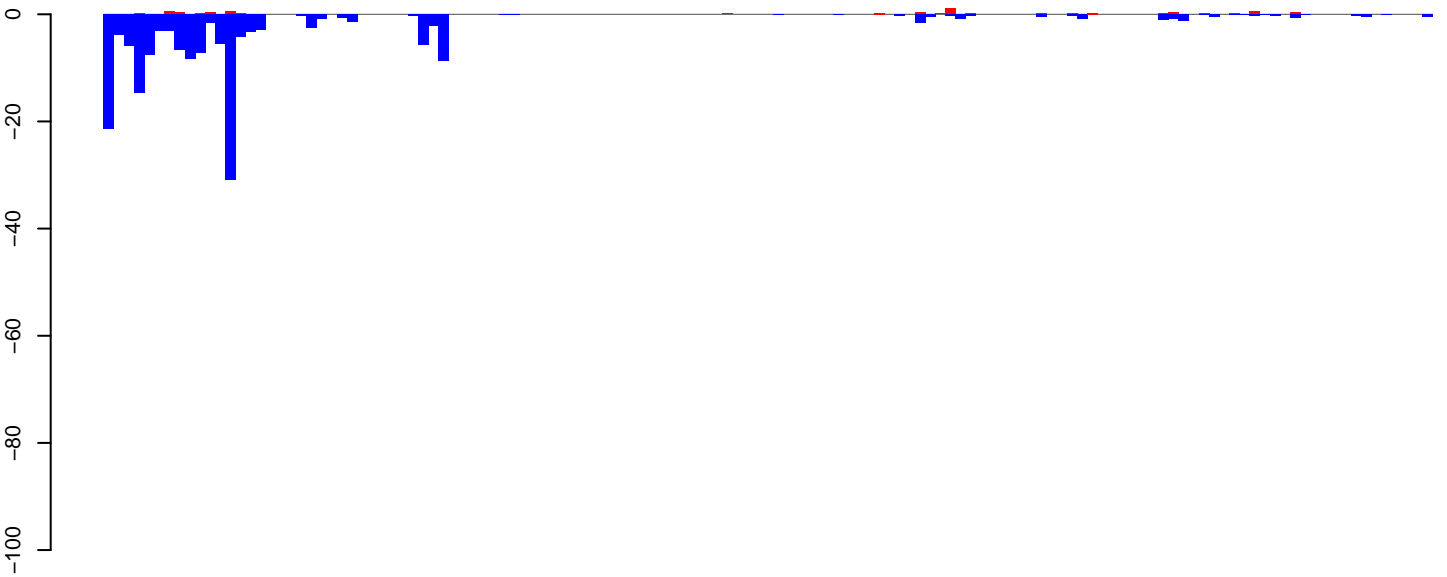
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



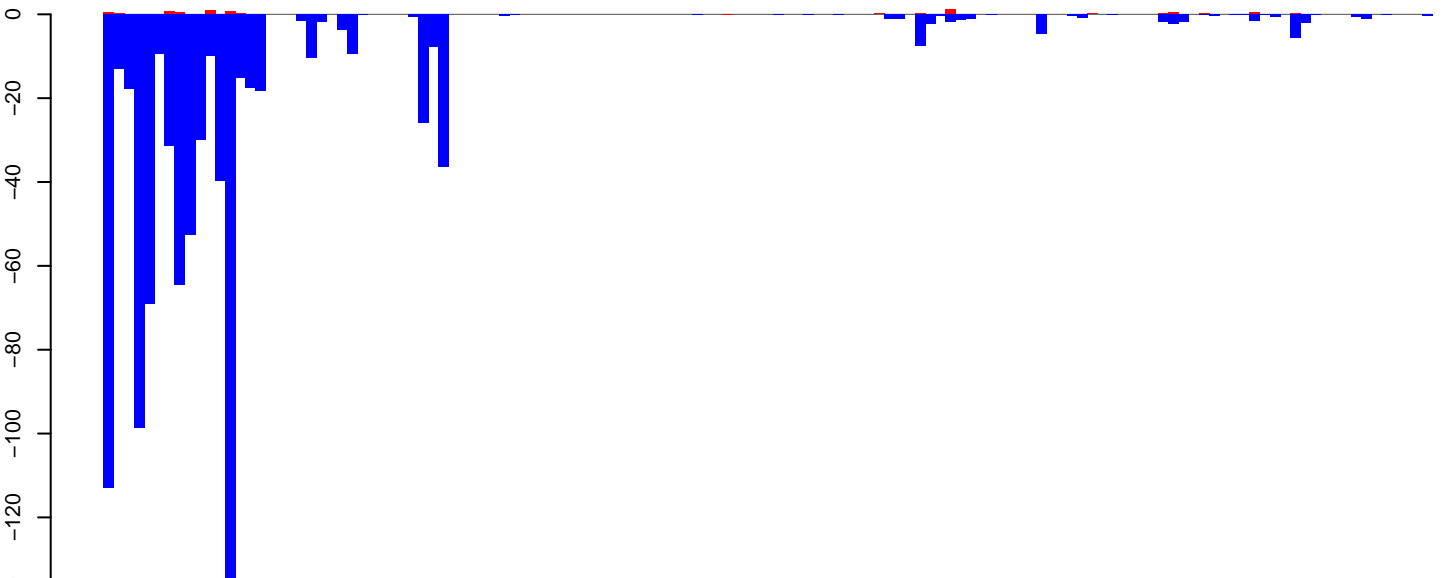
positive=7, negative=165, total=172

AnGam_Sua5b_cells.24_35.rep



positive=2, negative=718, total=720

AnGam_Sua5b_cells.rep

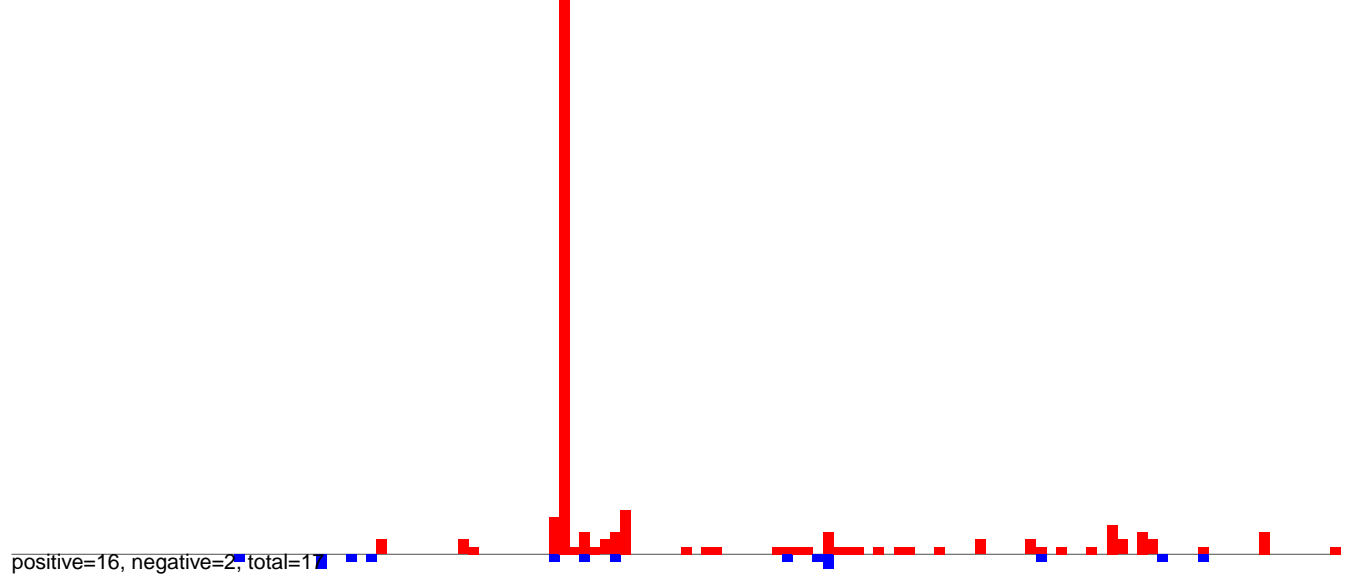


positive=9, negative=883, total=891

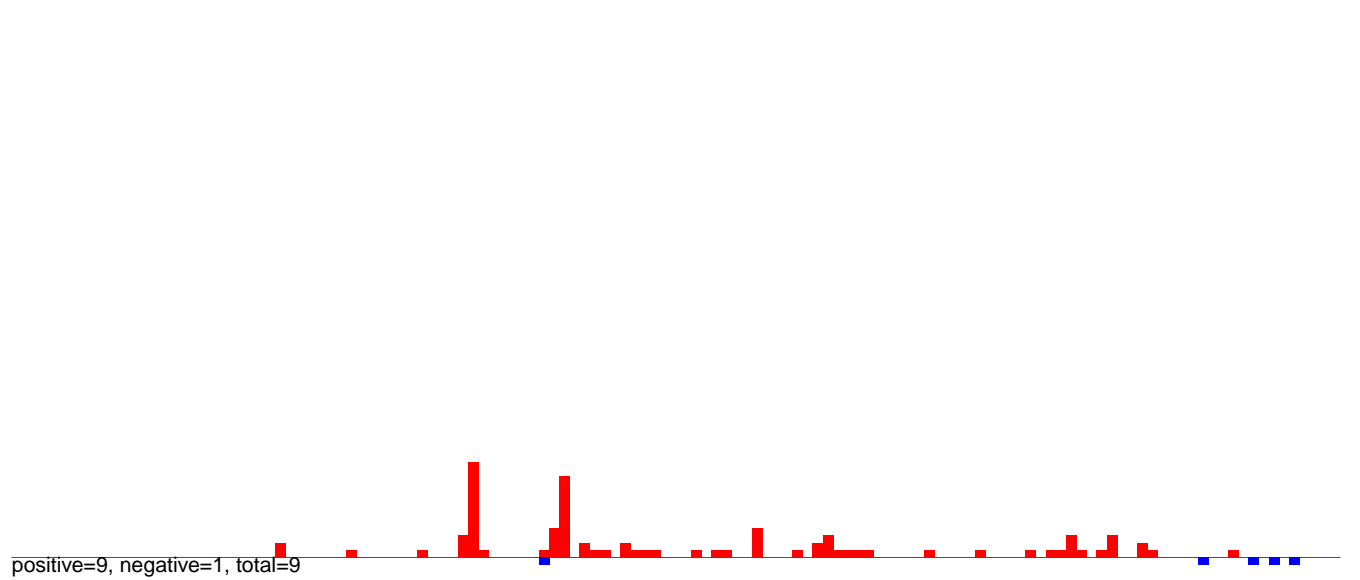
Window size=50, length=6565, TE@GYPSY12-LTR-I_AG:1-6565

0 1000 2000 3000 4000 5000 6000

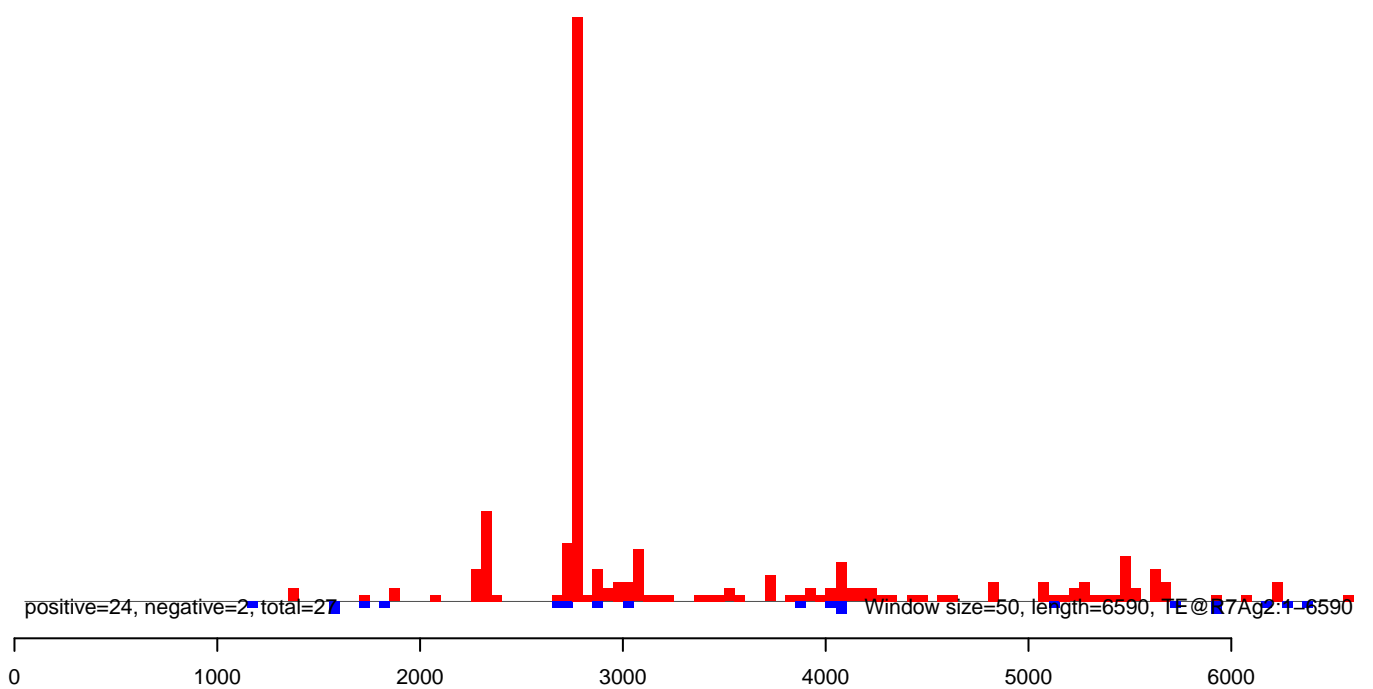
AnGam_Sua5b_cells.18_23.rep



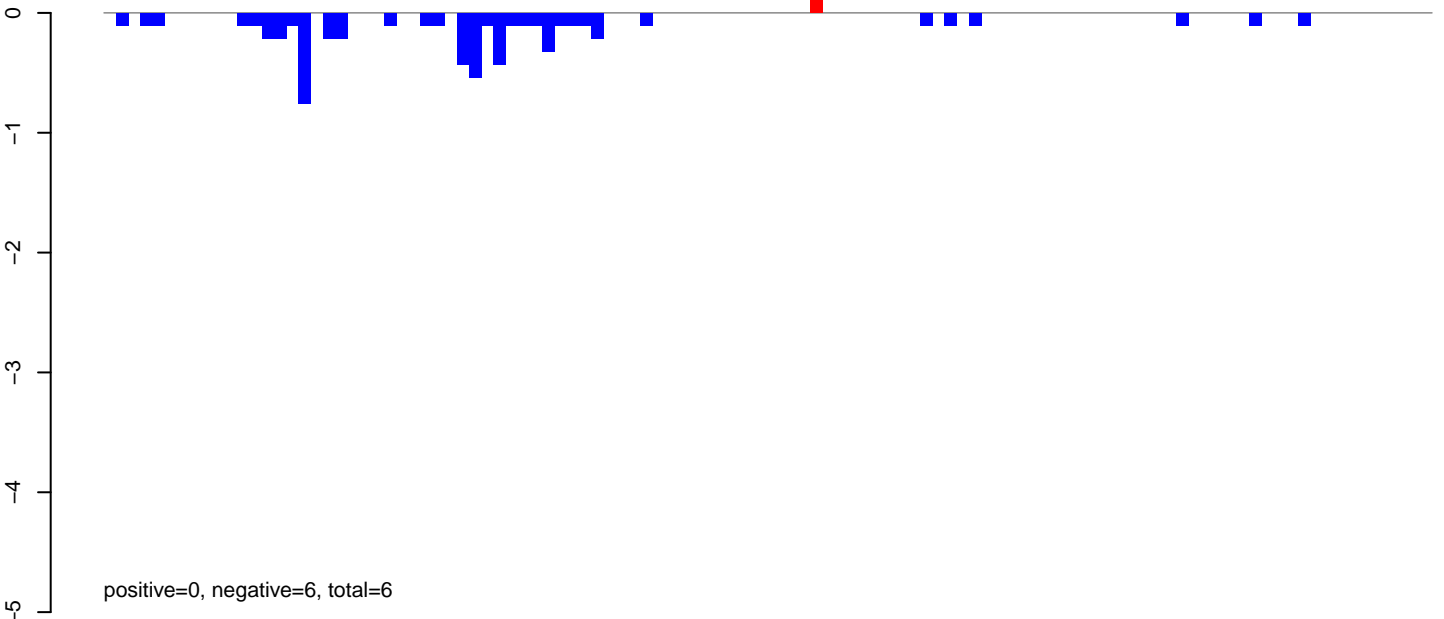
AnGam_Sua5b_cells.24_35.rep



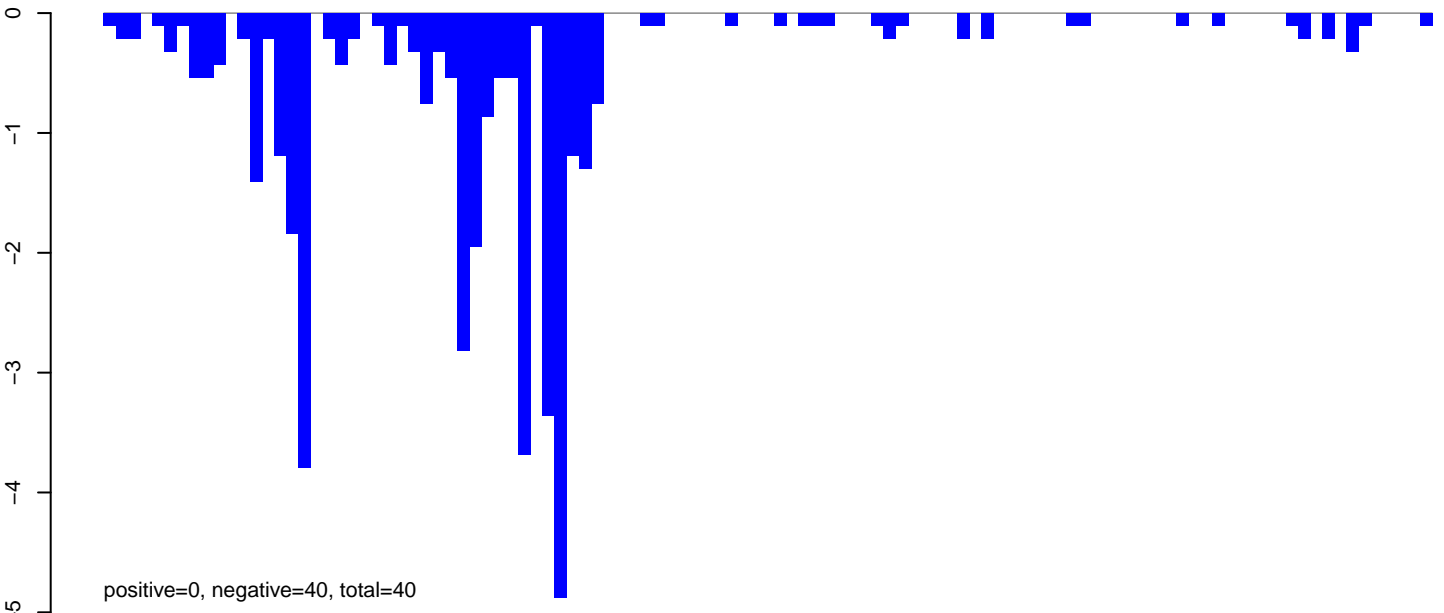
AnGam_Sua5b_cells.rep



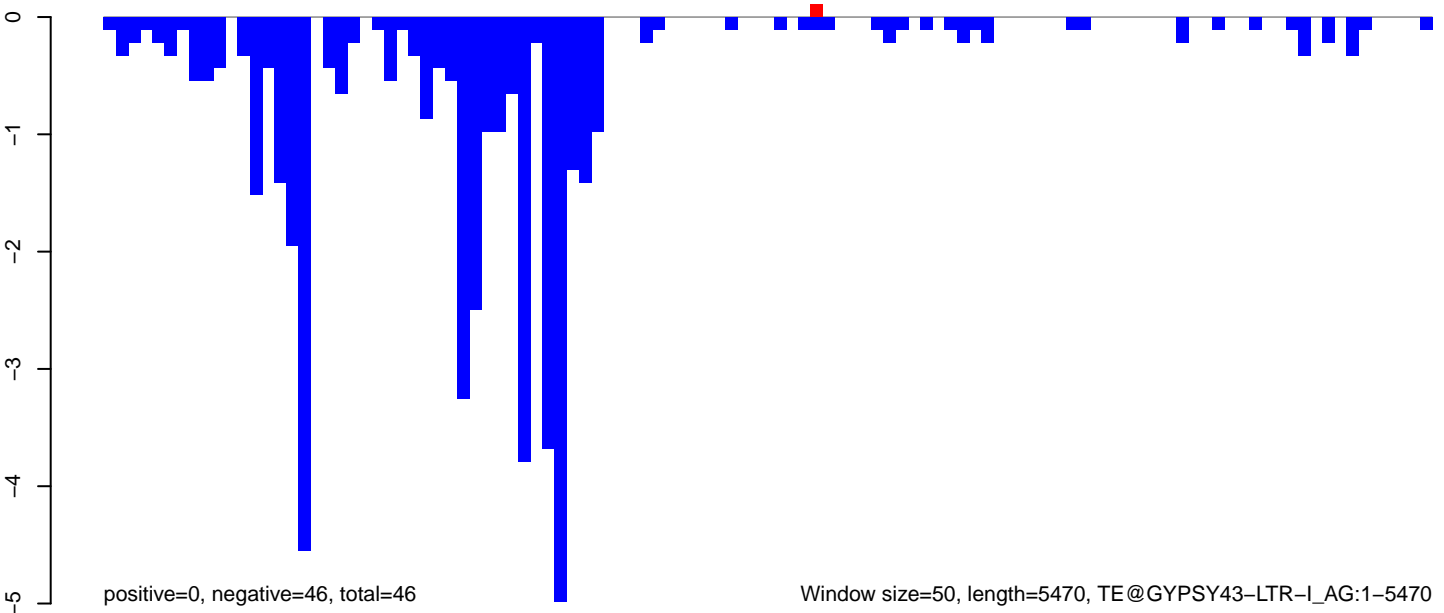
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



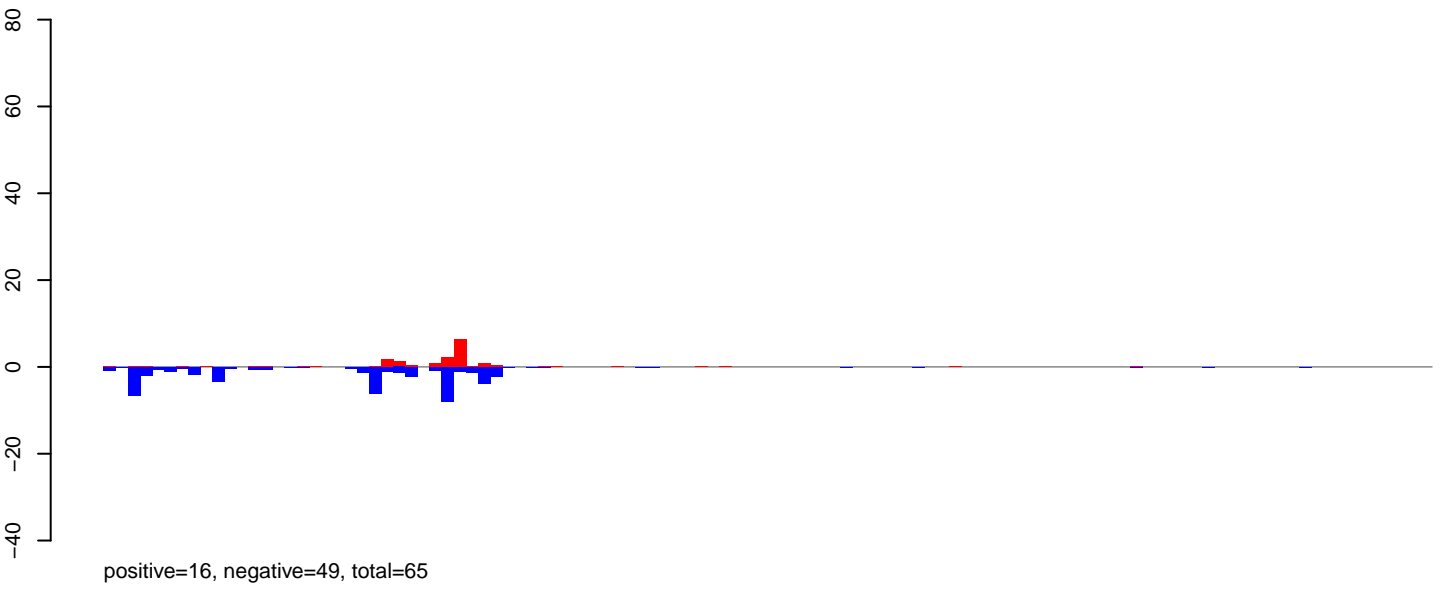
AnGam_Sua5b_cells.rep



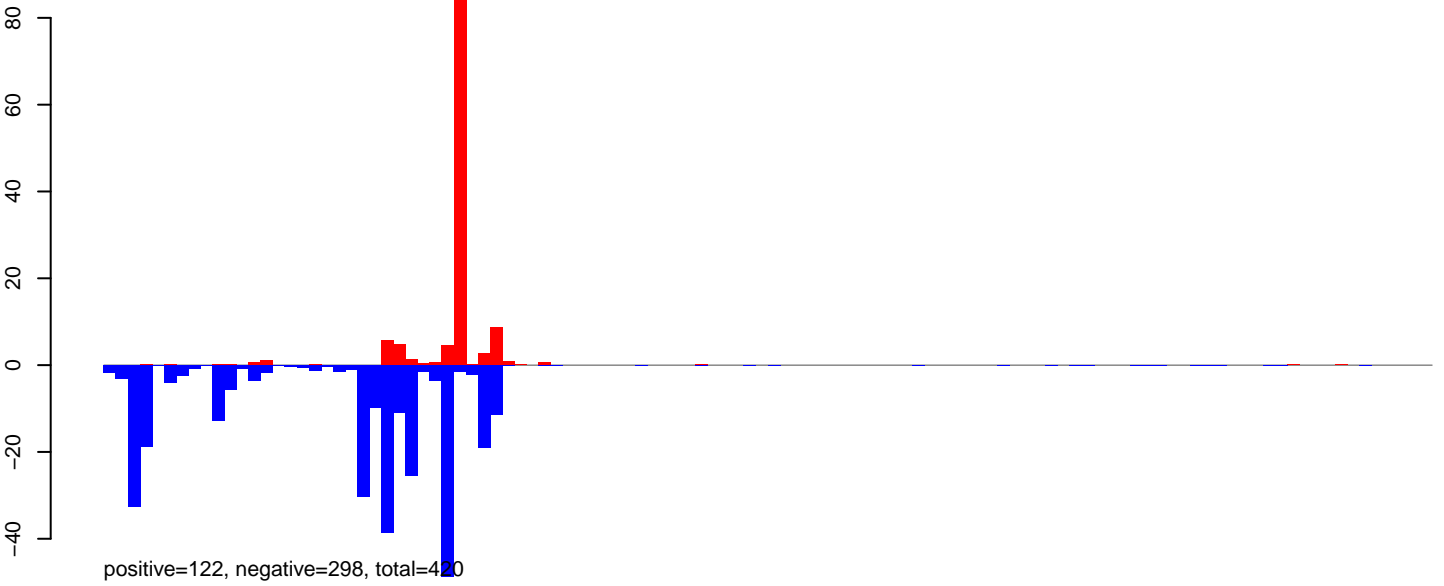
Window size=50, length=5470, TE@GYPSY43-LTR-I_AG:1-5470

0 1000 2000 3000 4000 5000

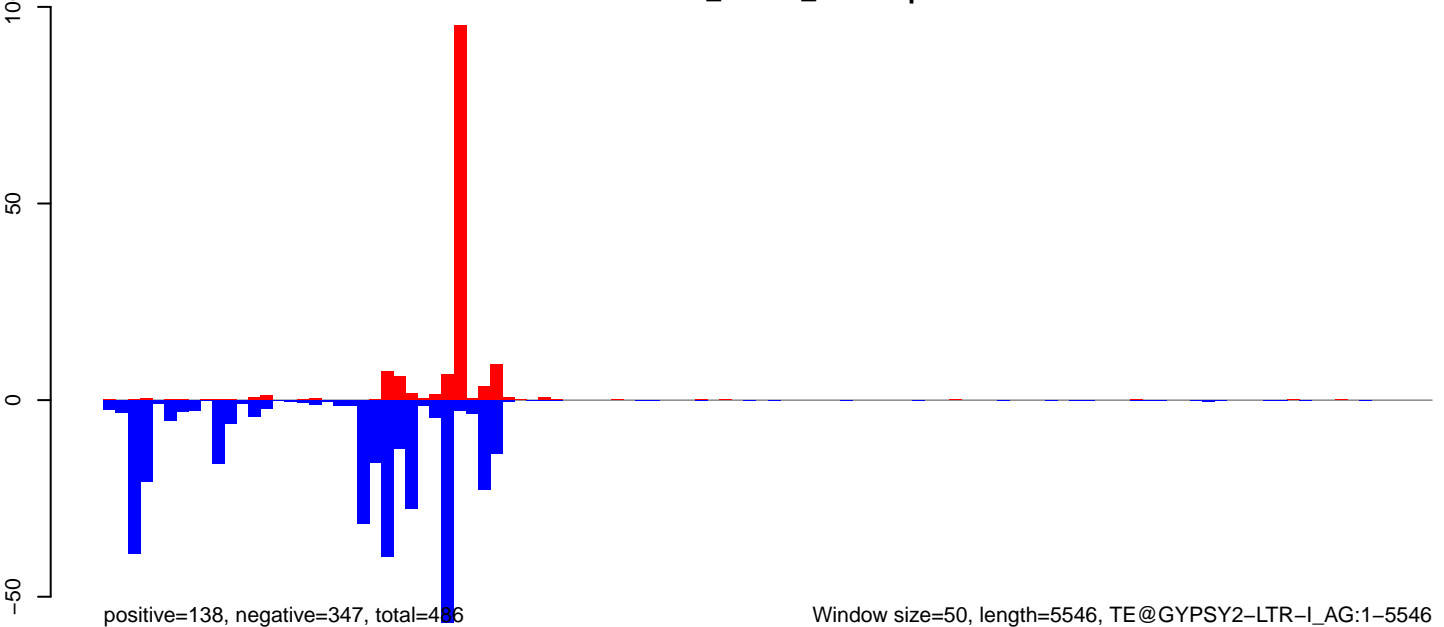
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

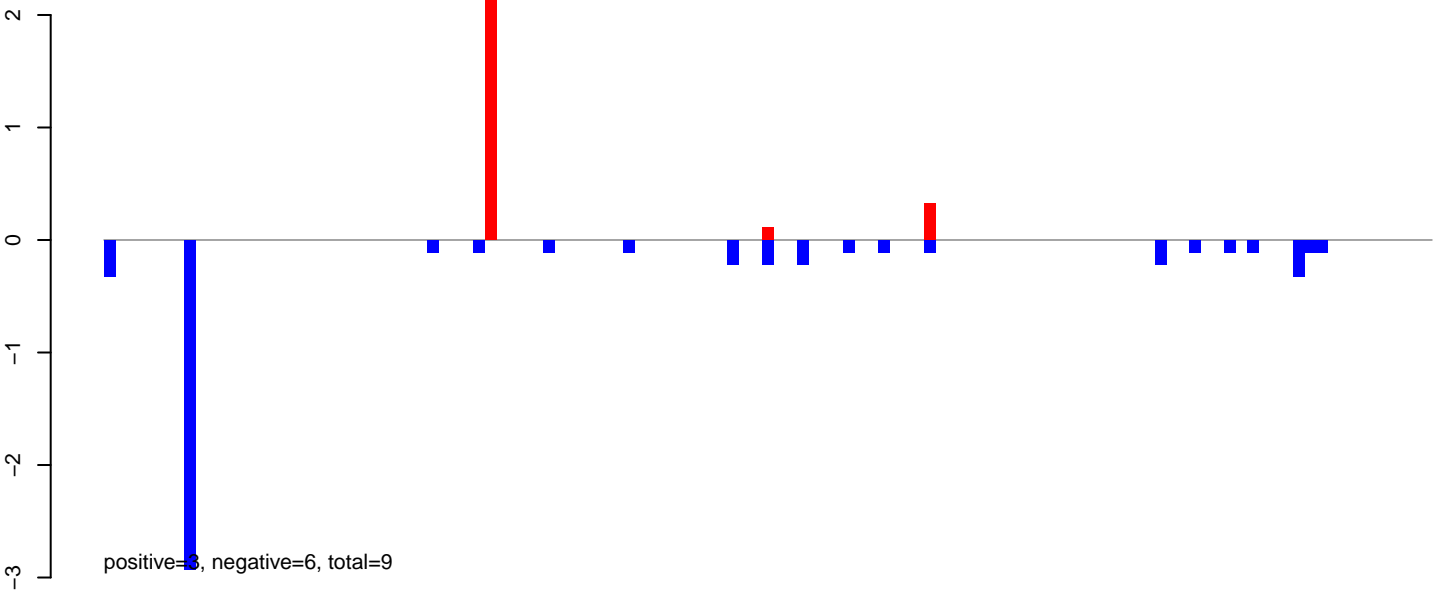


AnGam_Sua5b_cells.rep

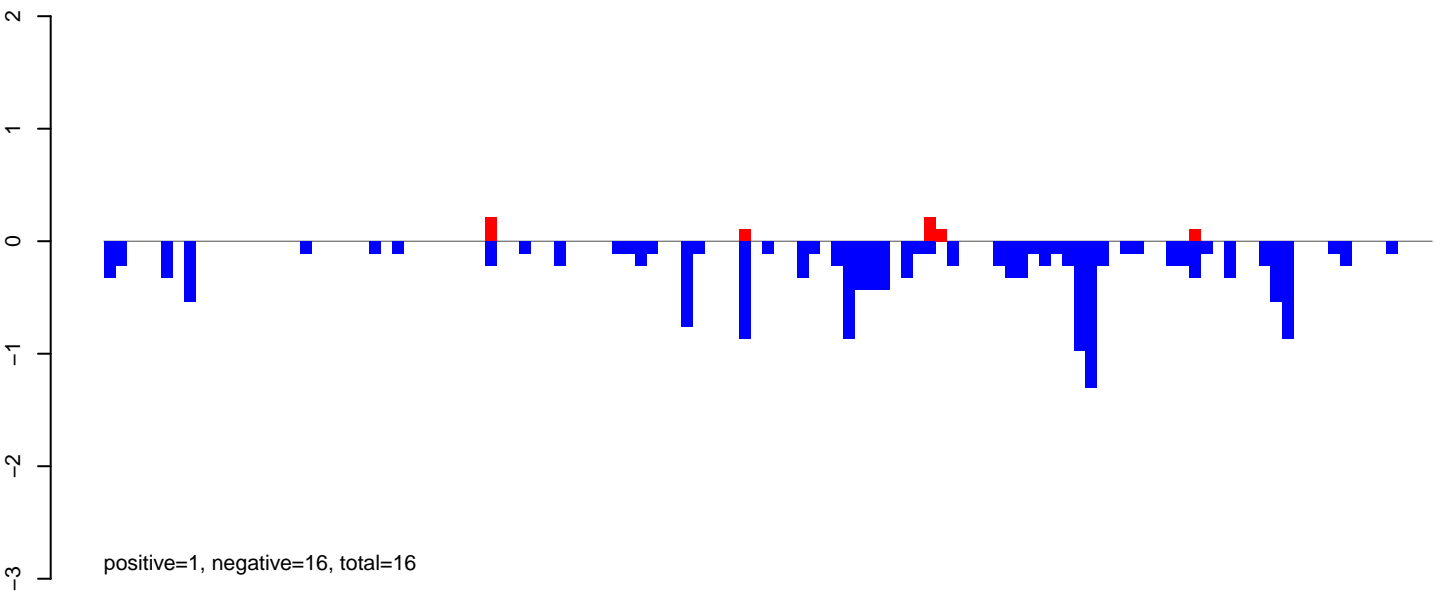


Window size=50, length=5546, TE@GYPSY2-LTR-I_AG:1-5546

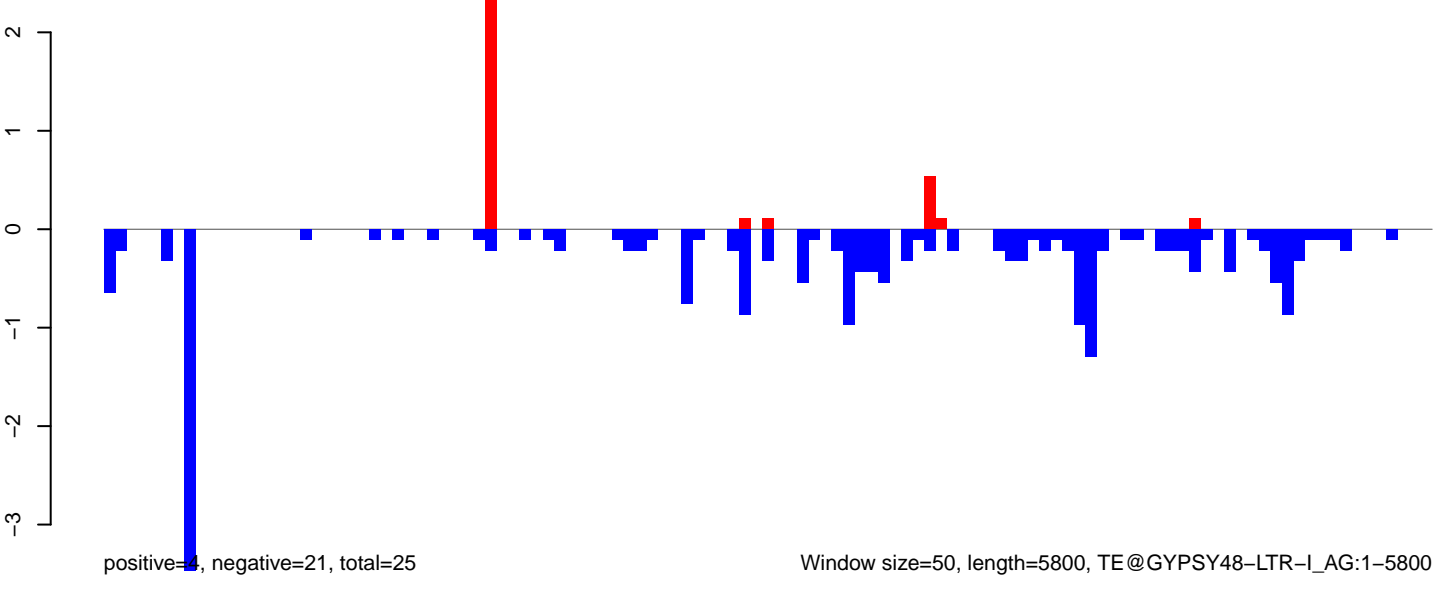
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



Window size=50, length=5800, TE@GYPSY48-LTR-I_AG:1-5800

AnGam_Sua5b_cells.18_23.rep

0
-5
-10
-15

positive=1, negative=25, total=26

AnGam_Sua5b_cells.24_35.rep

0
-5
-10
-15

positive=0, negative=80, total=81

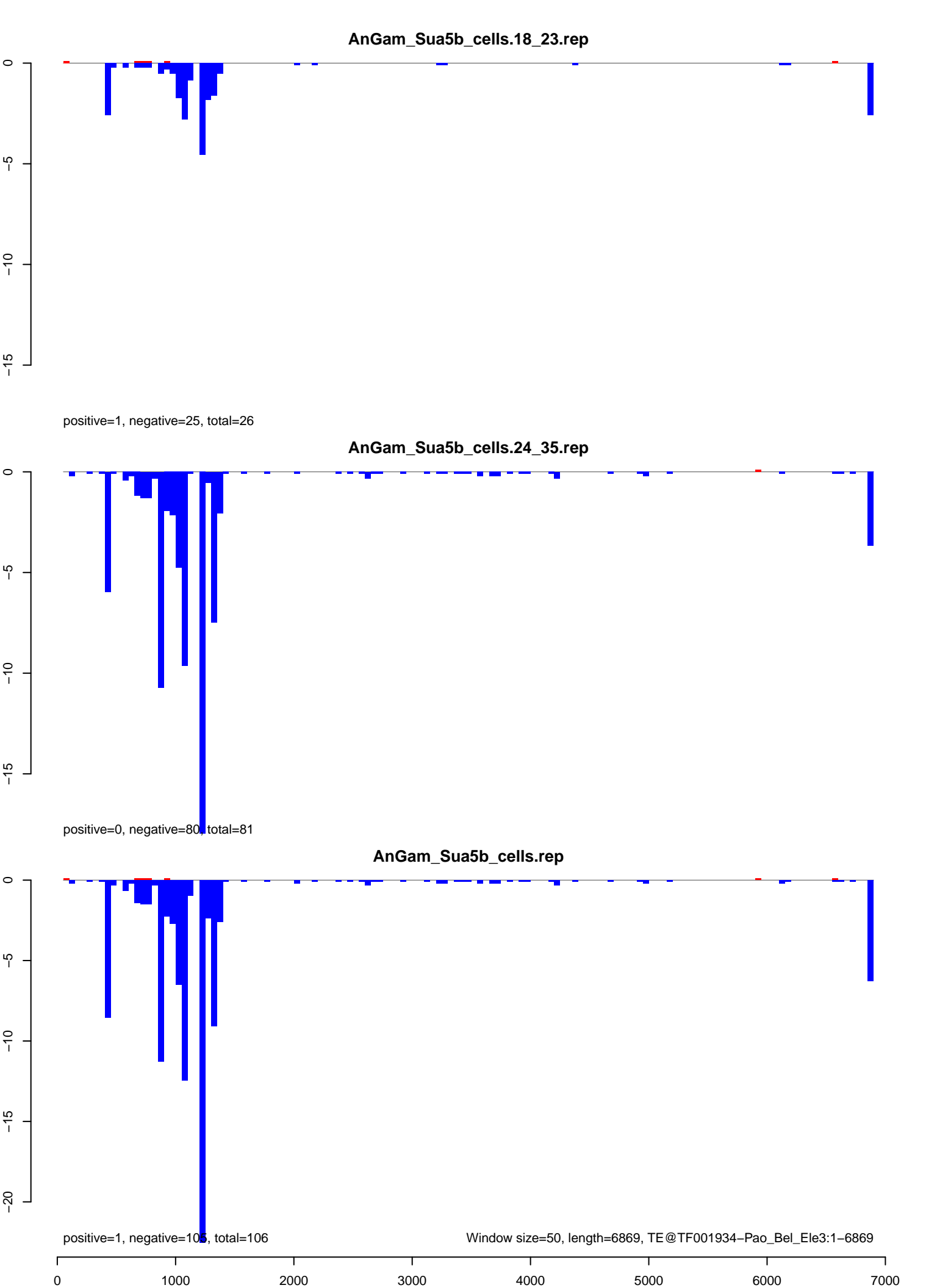
AnGam_Sua5b_cells.rep

0
-5
-10
-15
-20

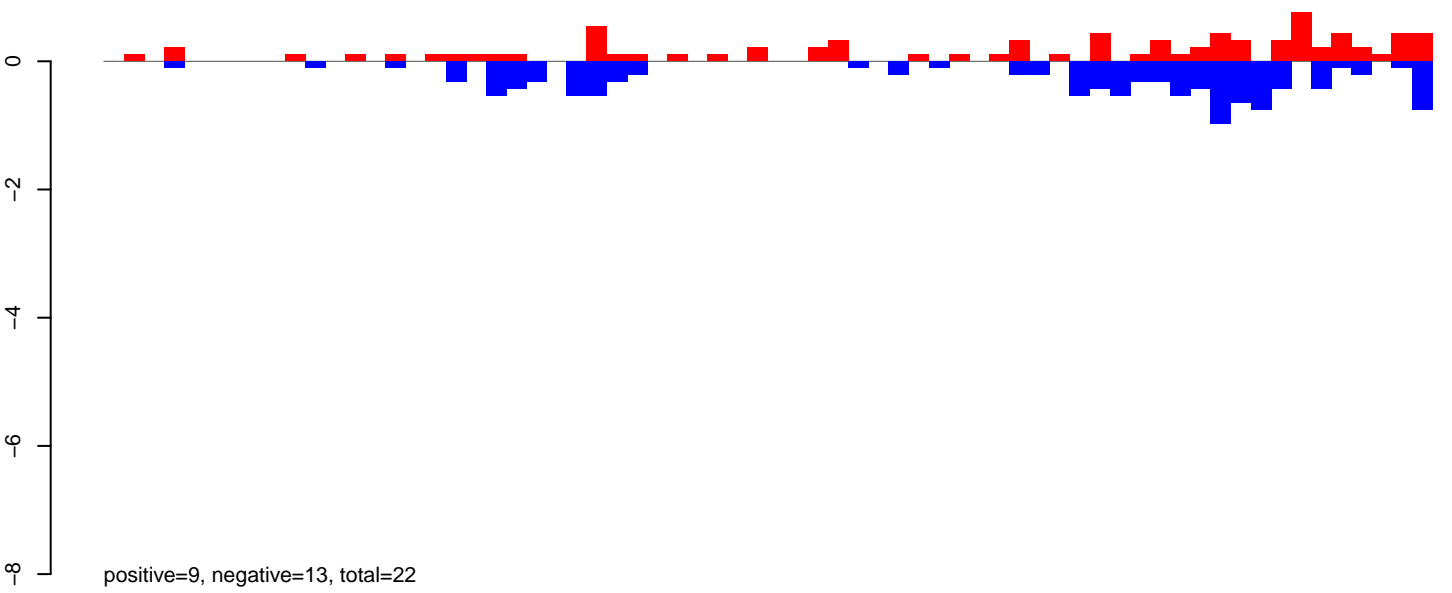
positive=1, negative=105, total=106

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

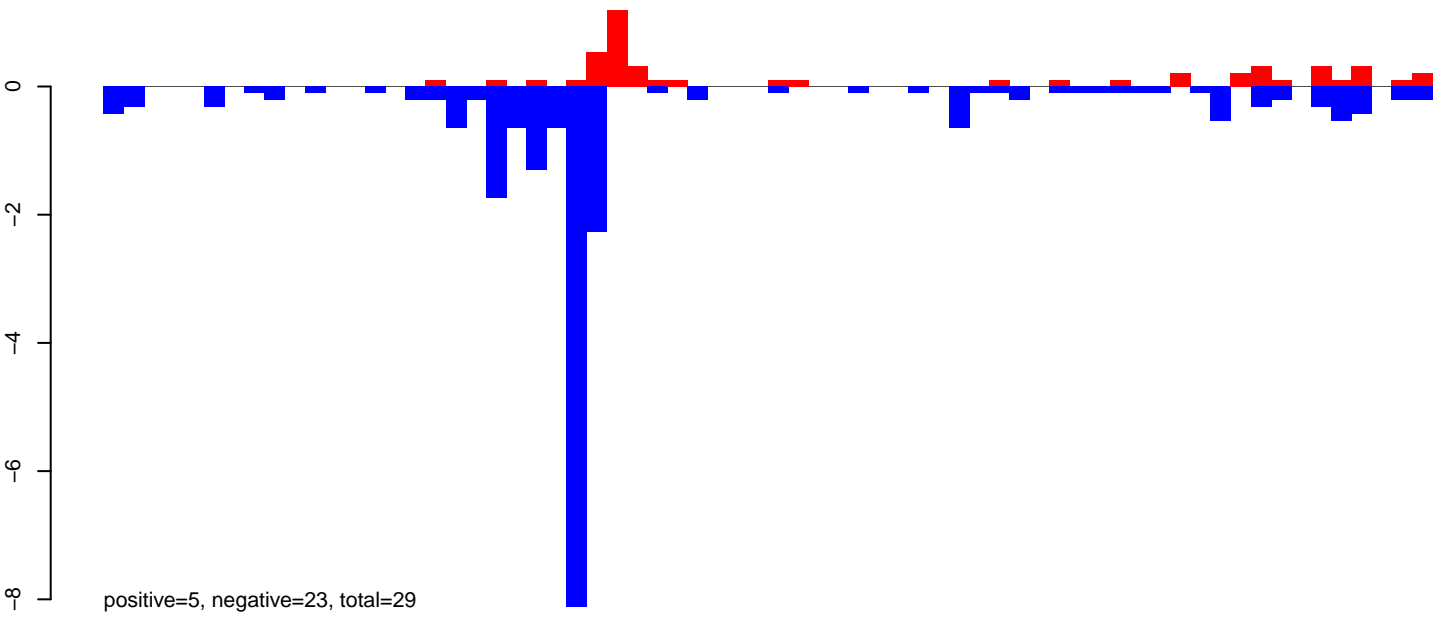
0 1000 2000 3000 4000 5000 6000 7000



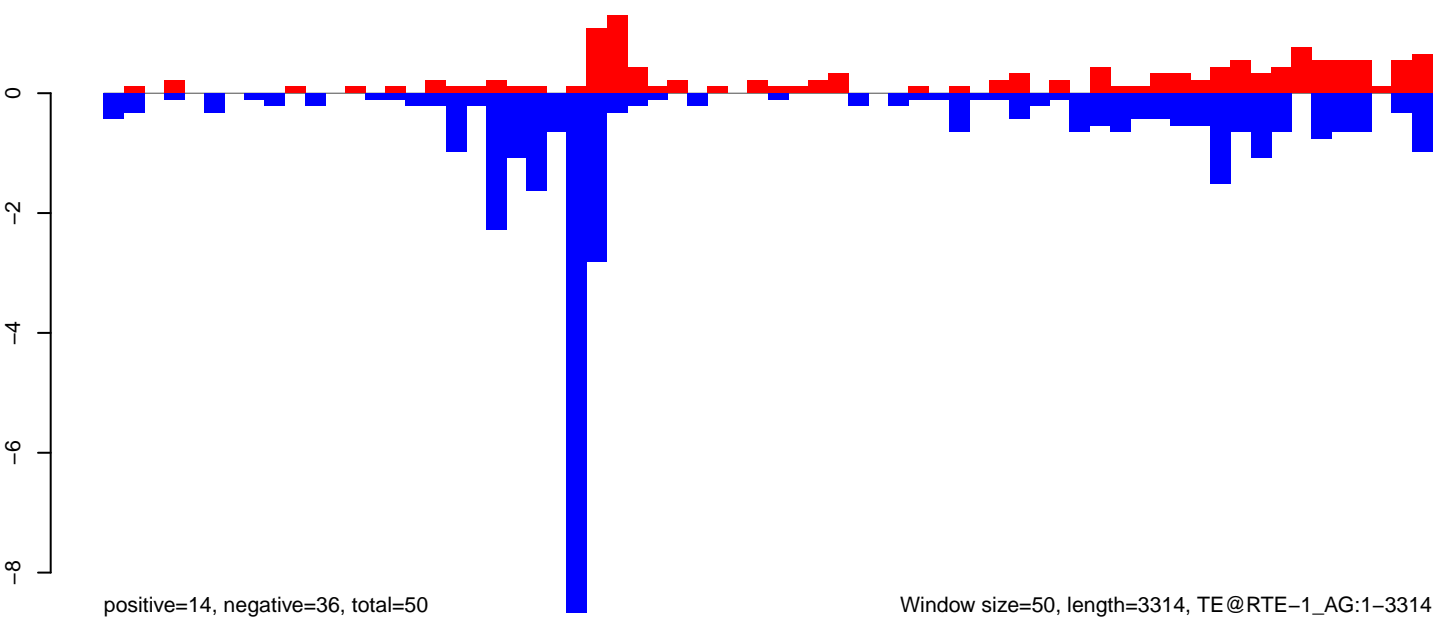
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

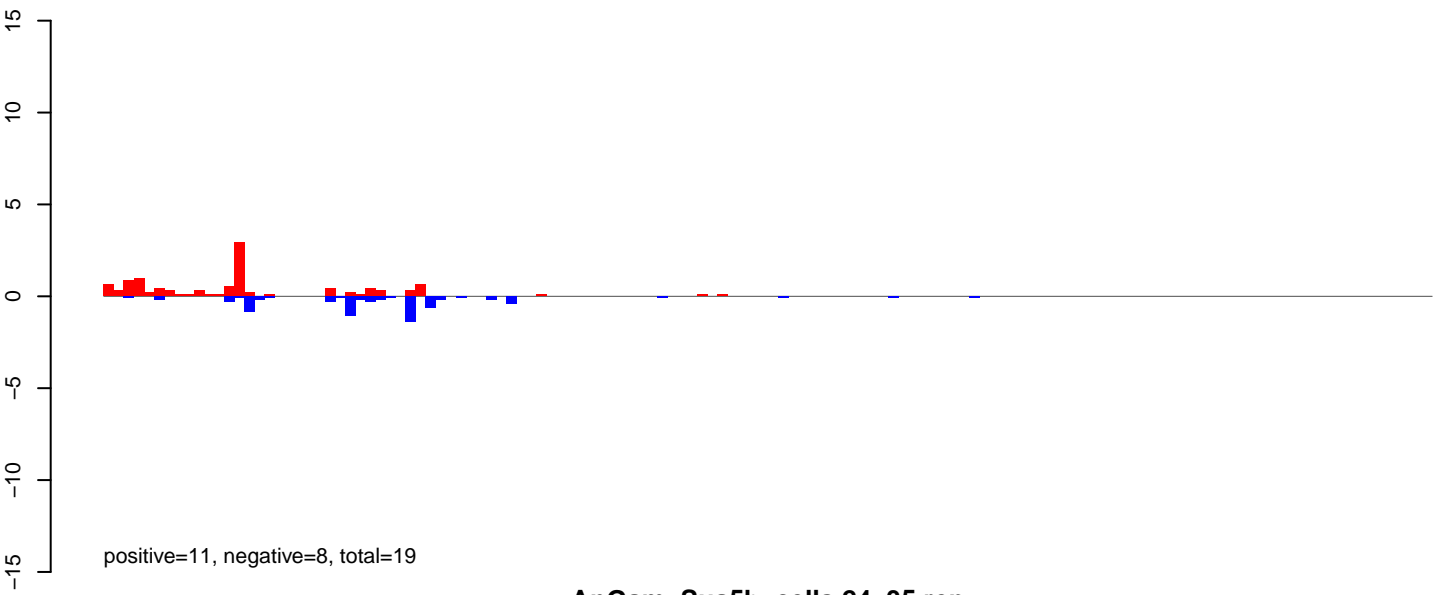


AnGam_Sua5b_cells.rep

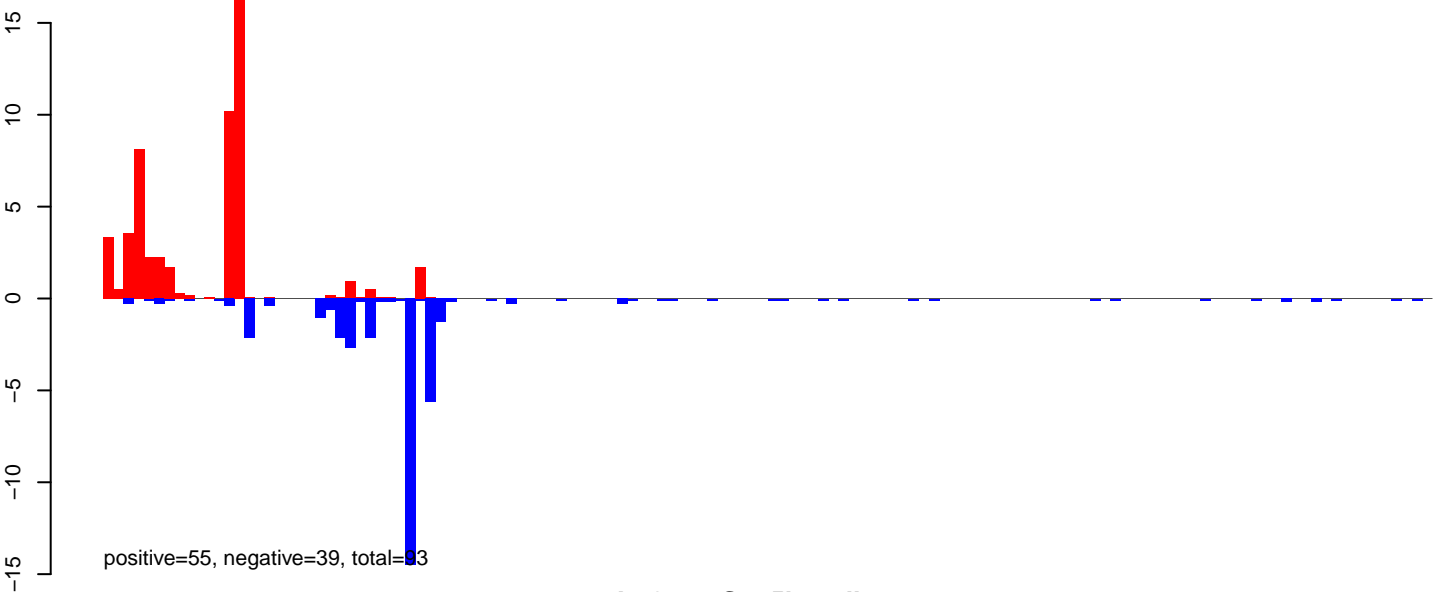


Window size=50, length=3314, TE@RTE-1_AG:1-3314

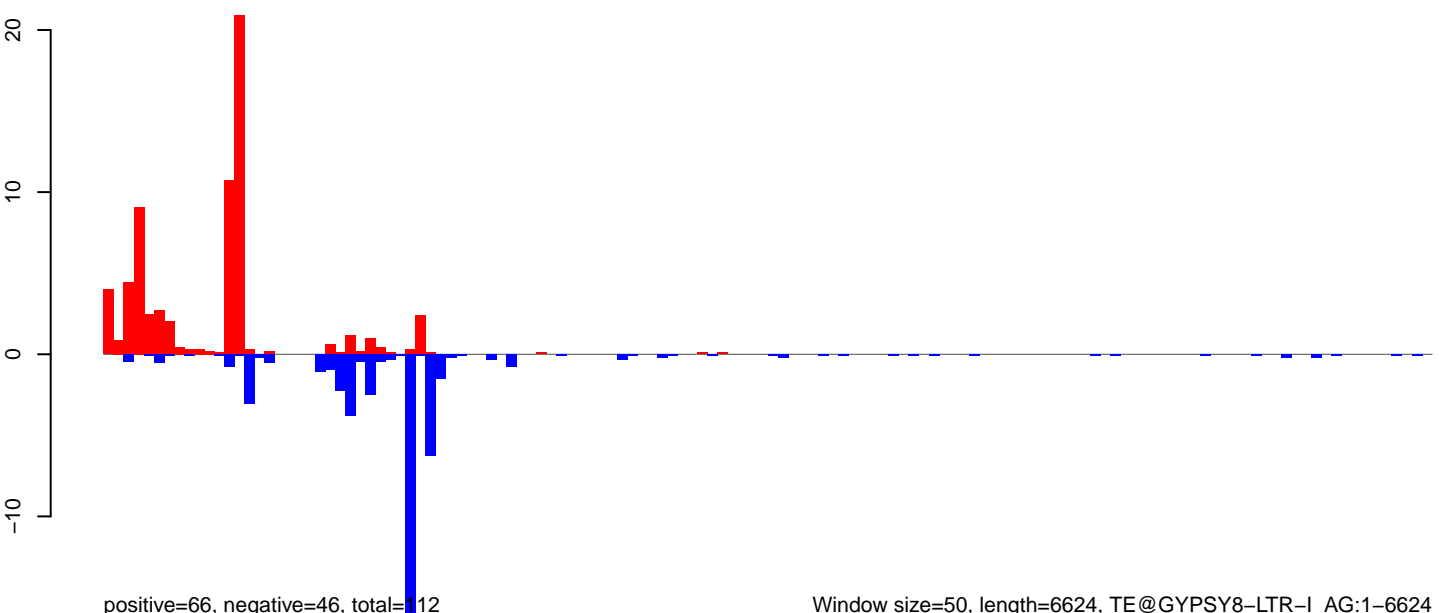
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

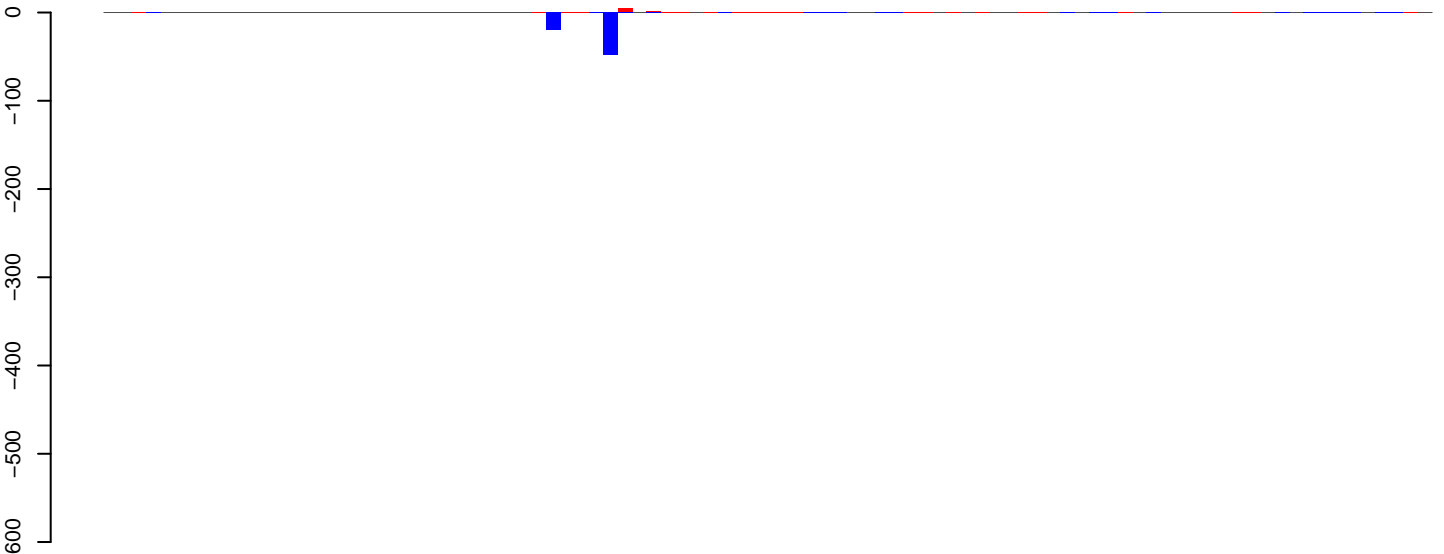


AnGam_Sua5b_cells.rep



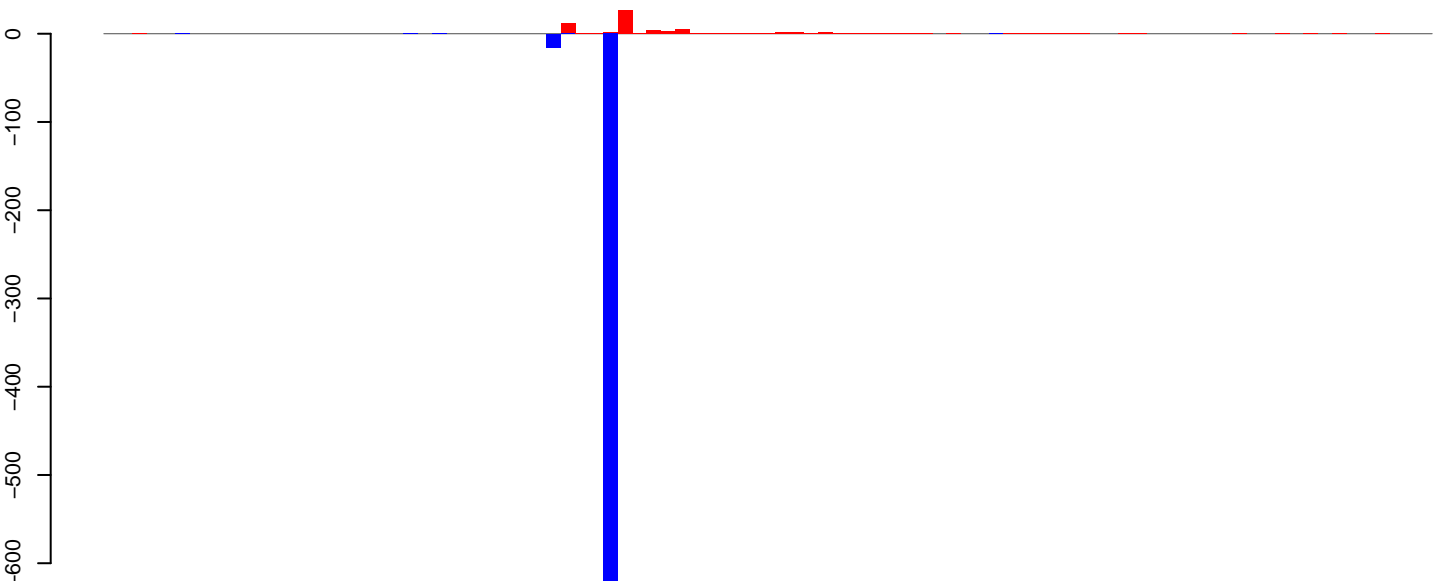
Window size=50, length=6624, TE@GYPSY8-LTR-L_AG:1-6624

AnGam_Sua5b_cells.18_23.rep



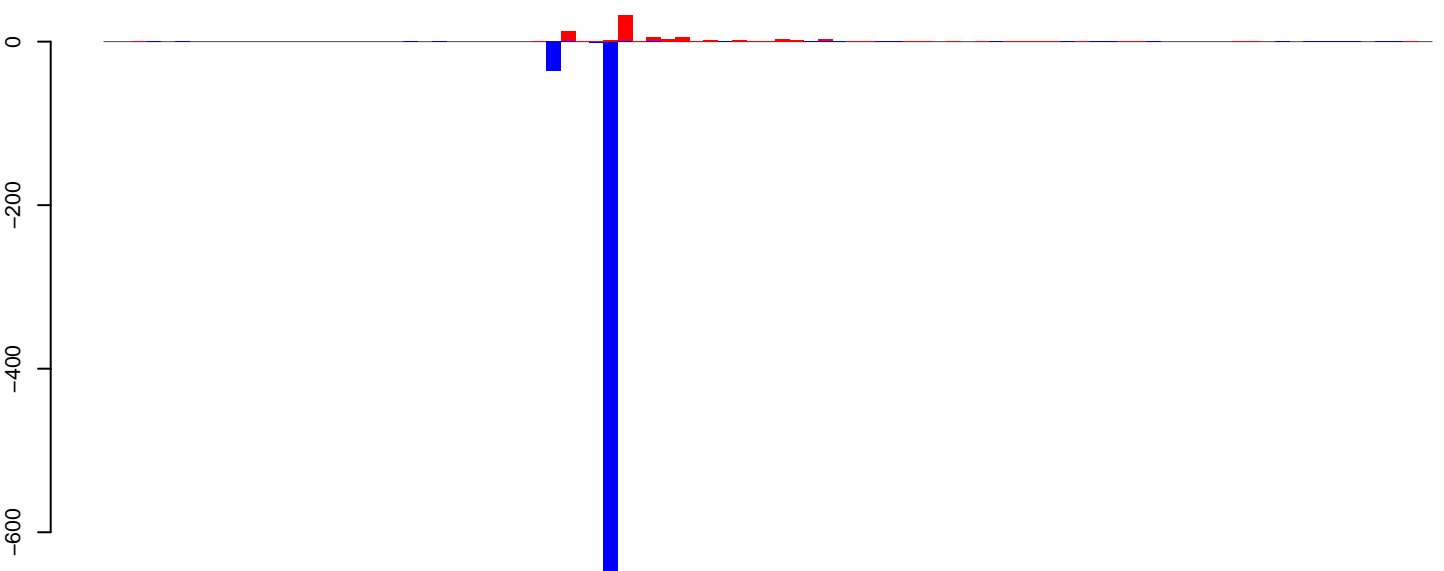
positive=15, negative=72, total=87

AnGam_Sua5b_cells.24_35.rep



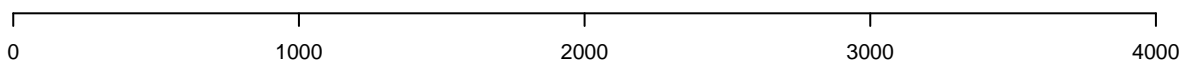
positive=70, negative=666, total=736

AnGam_Sua5b_cells.rep



positive=84, negative=738, total=823

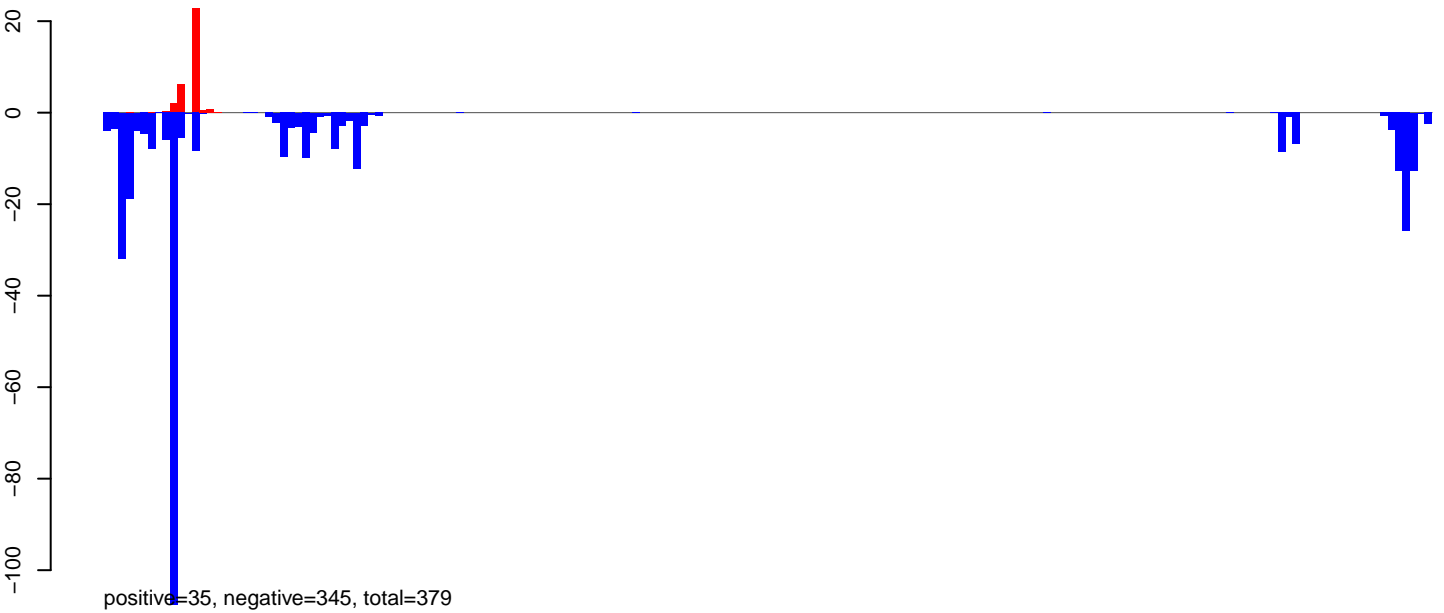
Window size=50, length=4697, TE@GYPSY4-LTR-I_AG:1-4697



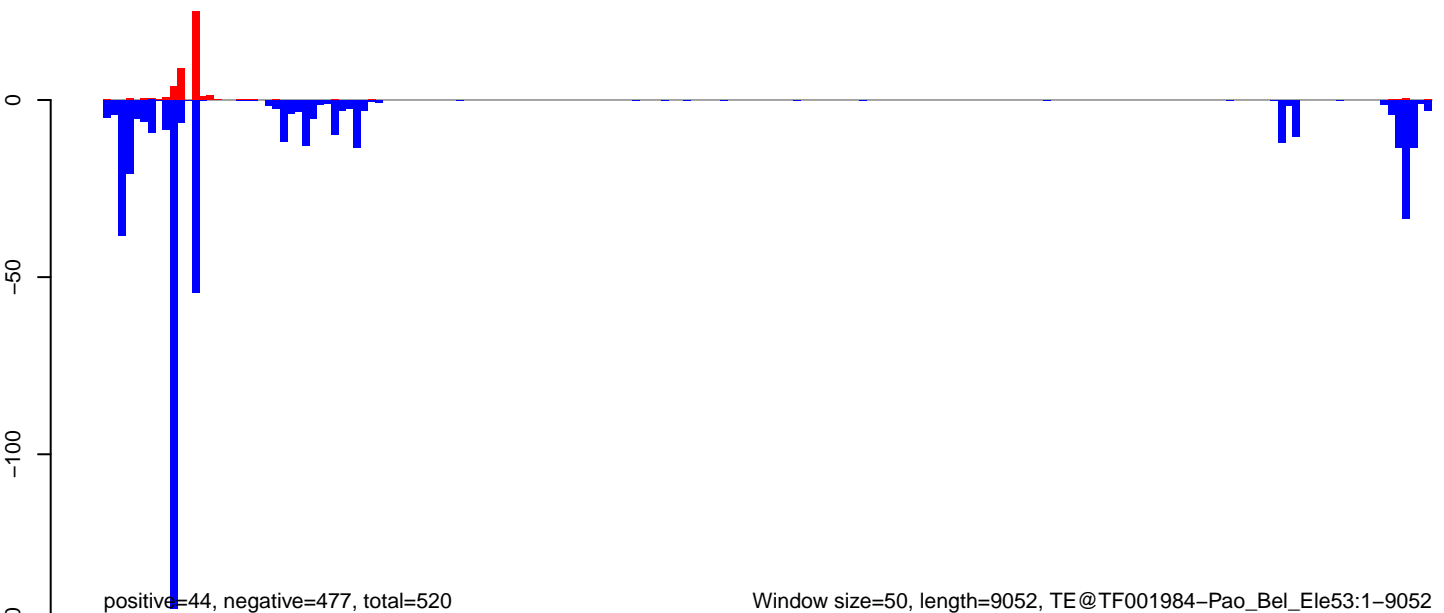
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

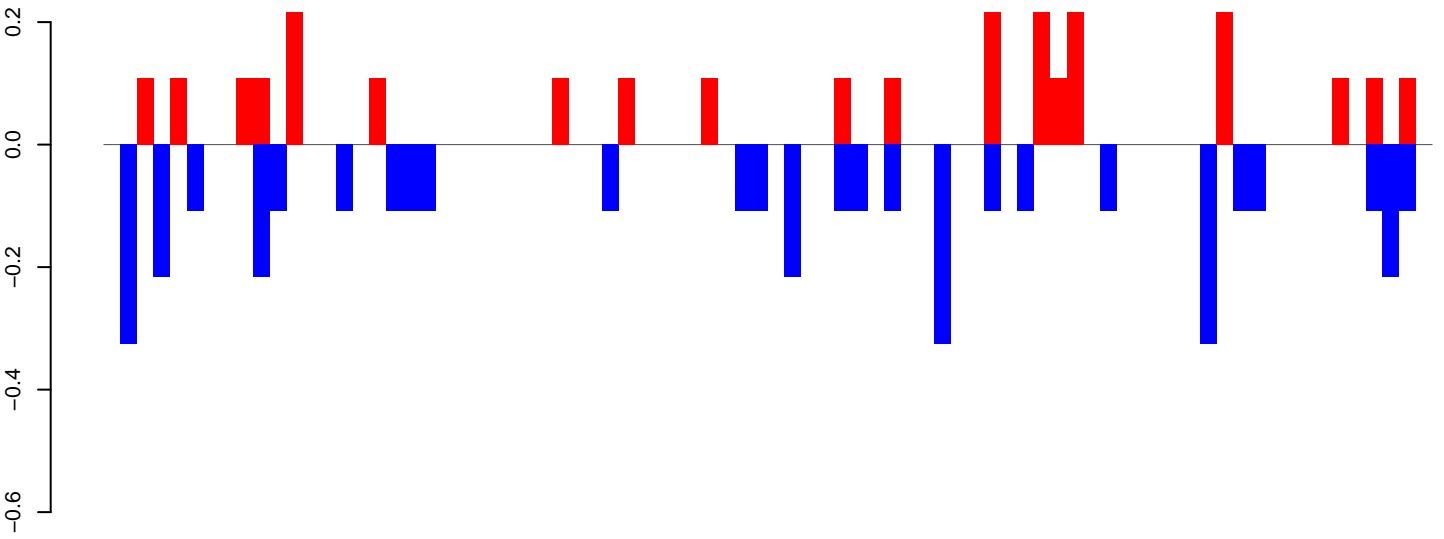


AnGam_Sua5b_cells.rep



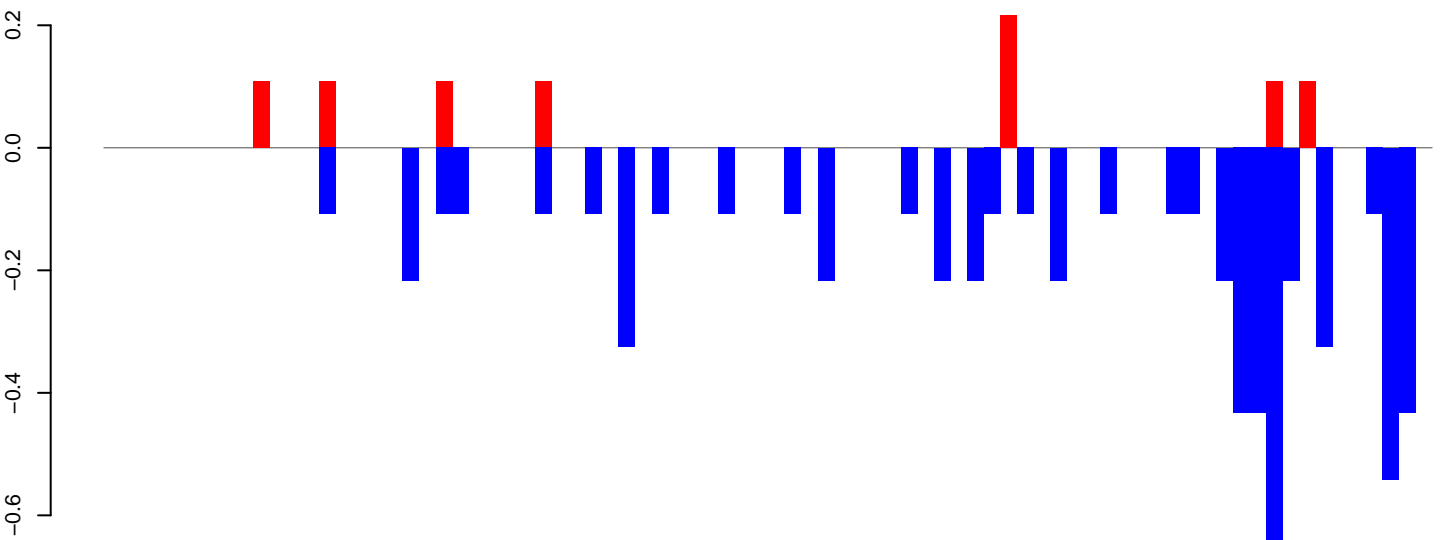
Window size=50, length=9052, TE@TF001984-Pao_Bel_Ele53:1-9052

AnGam_Sua5b_cells.18_23.rep



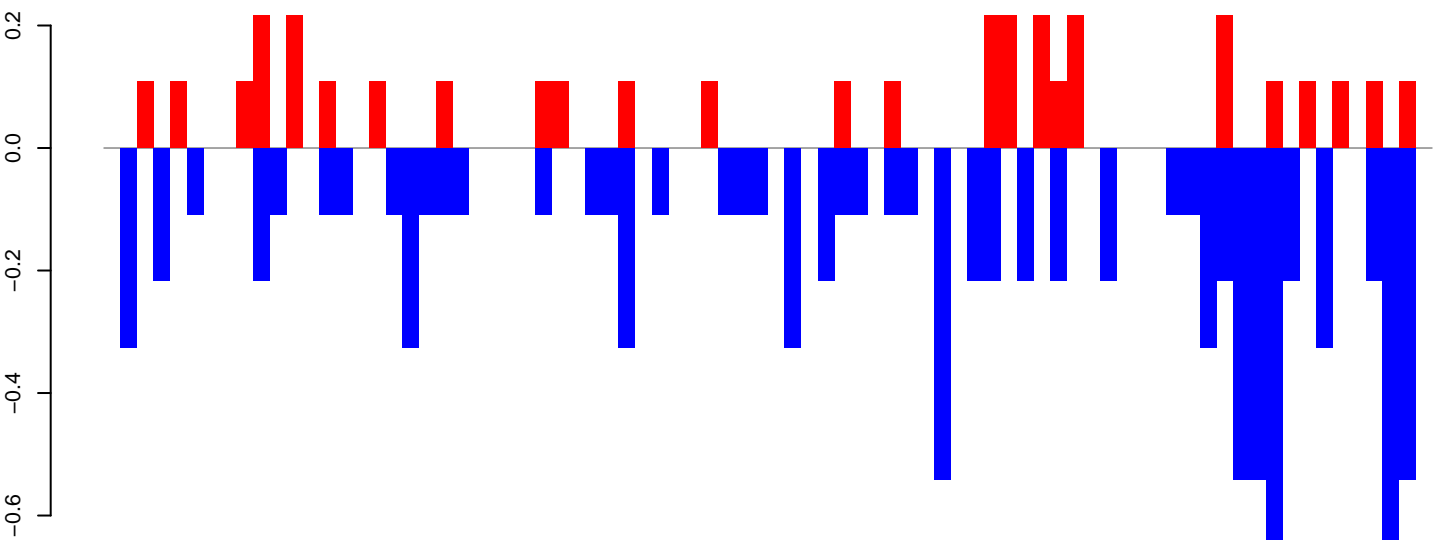
positive=3, negative=4, total=6

AnGam_Sua5b_cells.24_35.rep



positive=1, negative=6, total=7

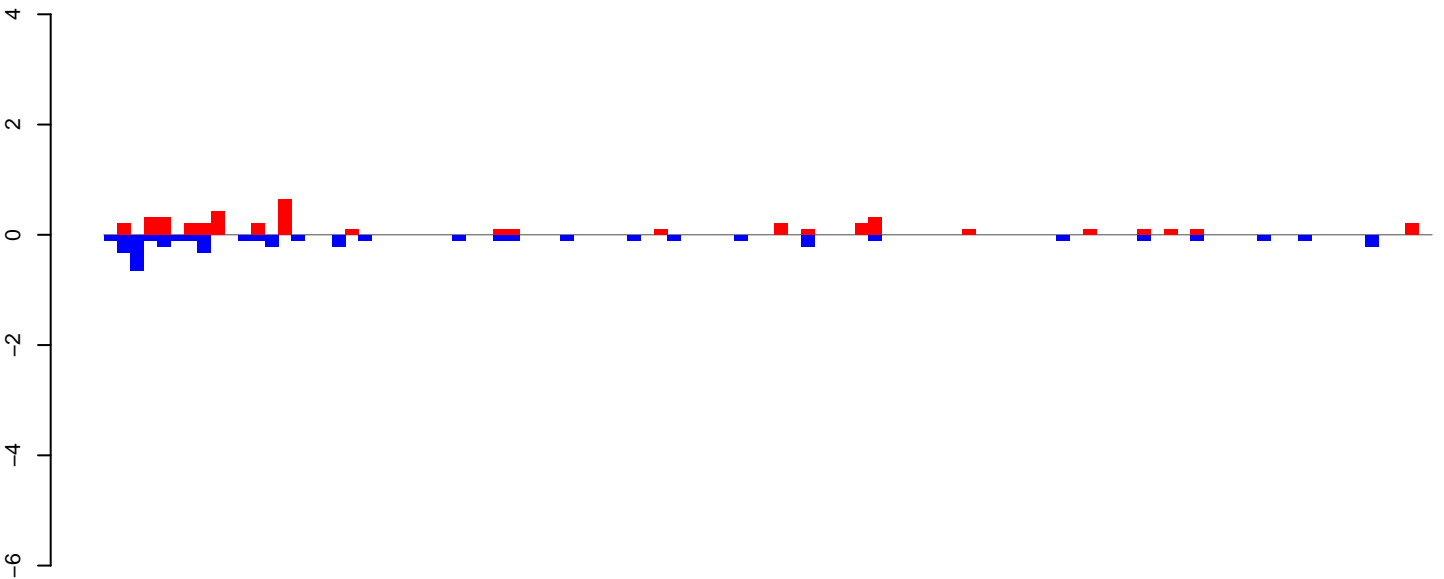
AnGam_Sua5b_cells.rep



positive=3, negative=10, total=14

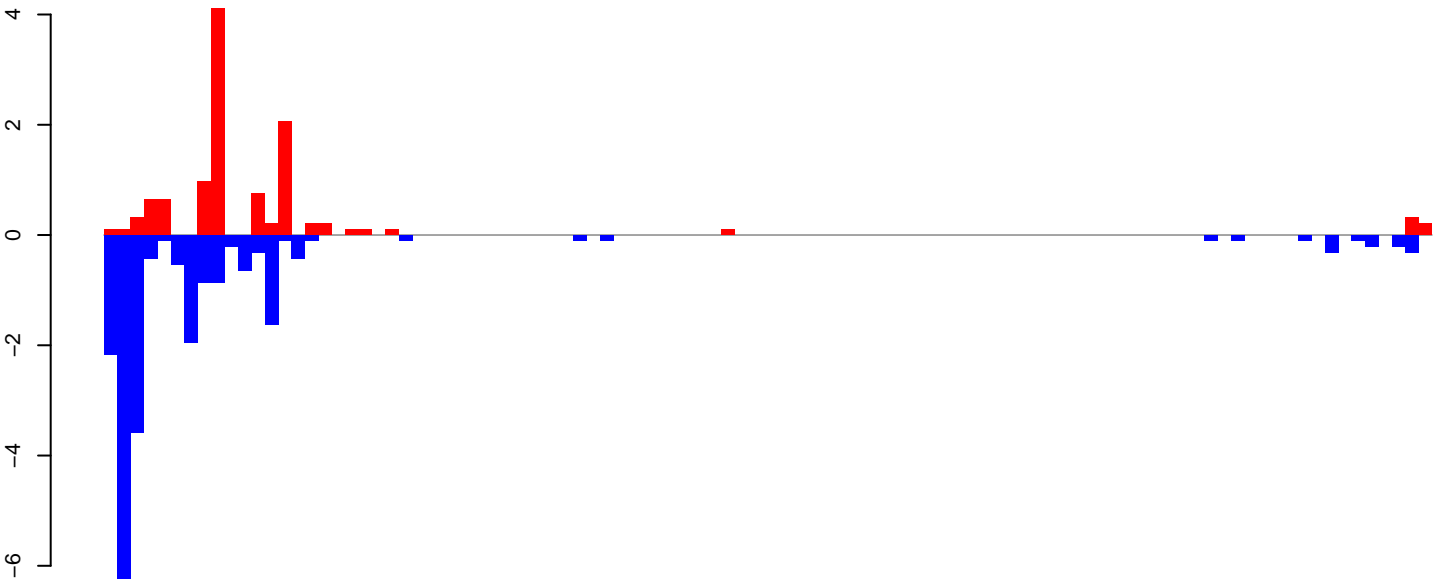
Window size=50, length=4046, TE@MTANGA_I:1-4046

AnGam_Sua5b_cells.18_23.rep



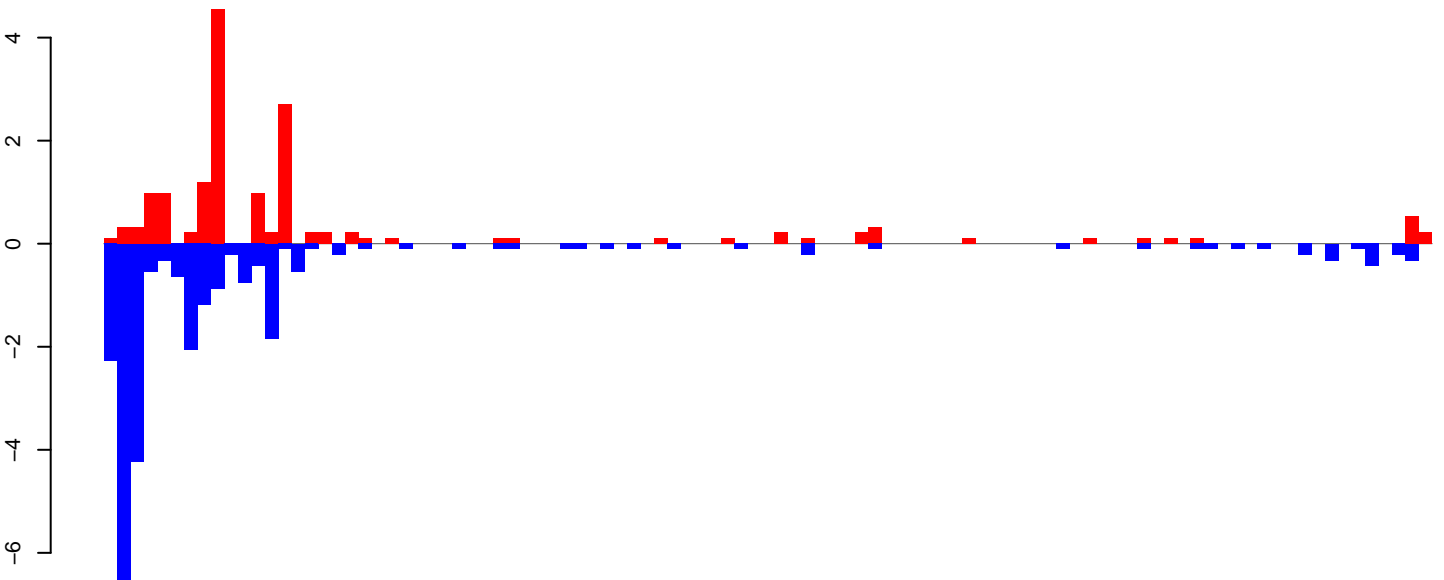
positive=5, negative=5, total=9

AnGam_Sua5b_cells.24_35.rep



positive=12, negative=23, total=34

AnGam_Sua5b_cells.rep

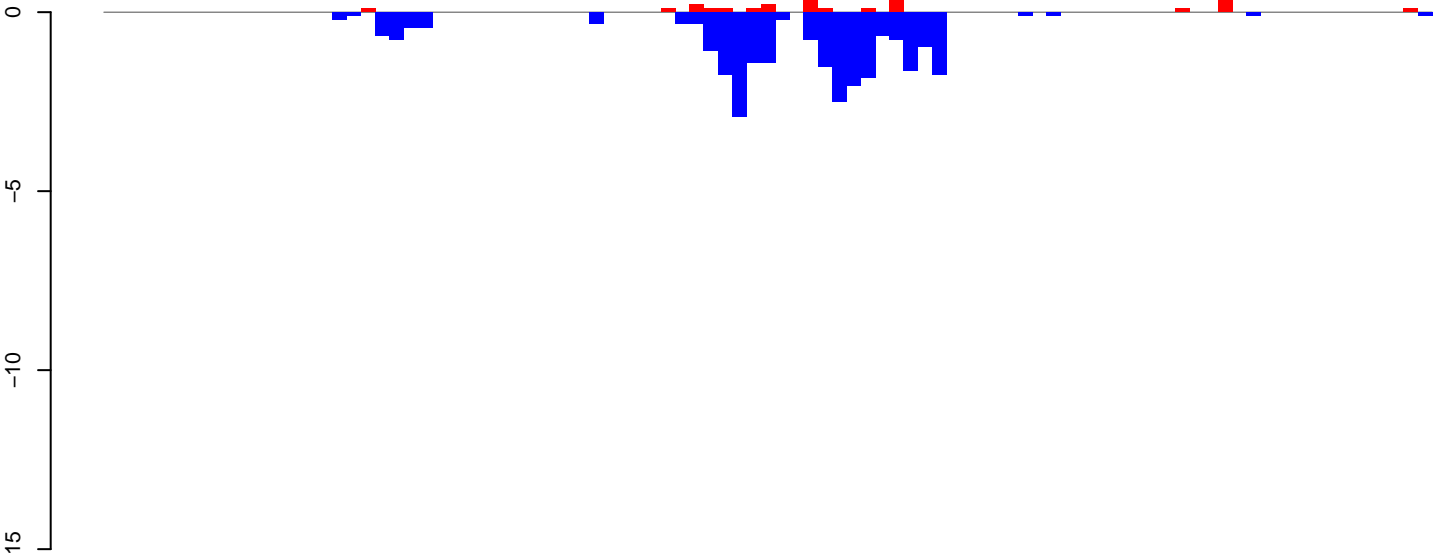


positive=16, negative=27, total=43

Window size=50, length=4989, TE@Ag-Outcast-5:1-4989

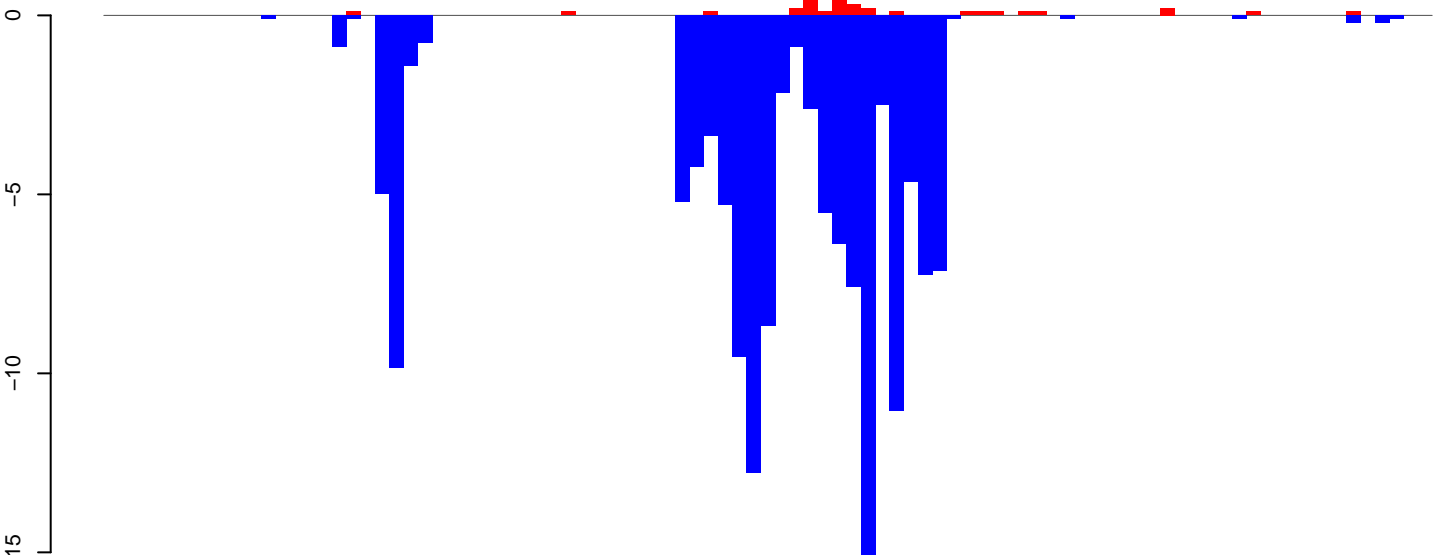
0 1000 2000 3000 4000 5000

AnGam_Sua5b_cells.18_23.rep



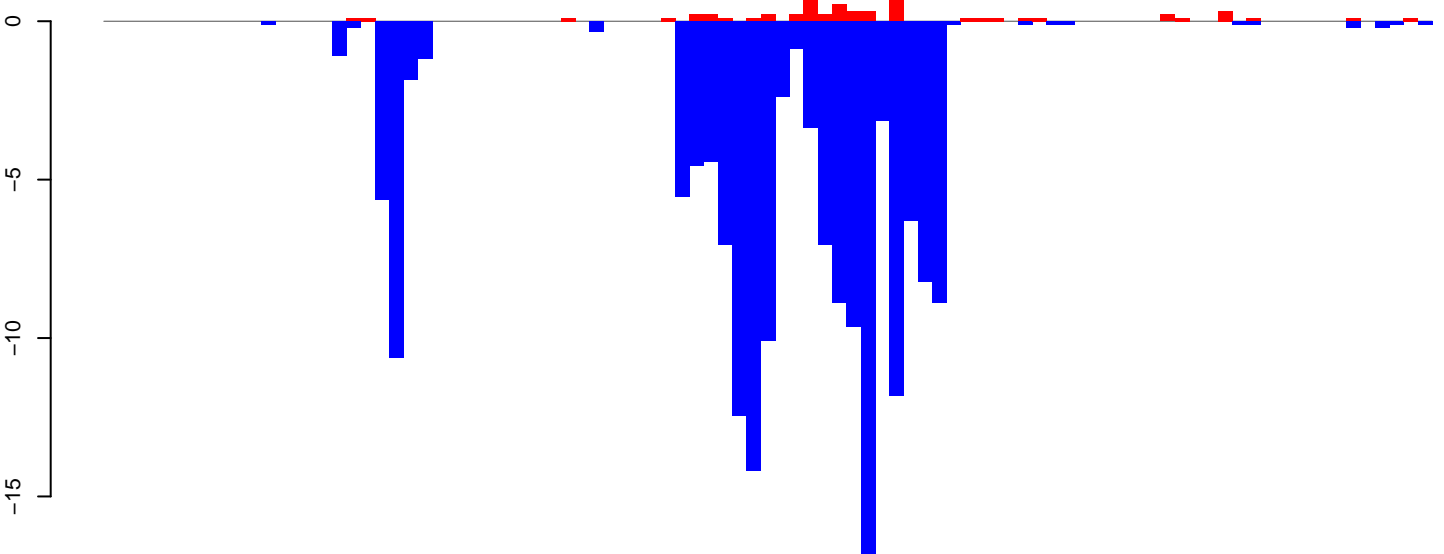
positive=3, negative=27, total=30

AnGam_Sua5b_cells.24_35.rep



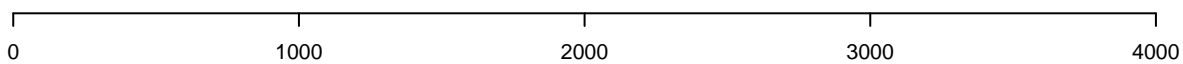
positive=3, negative=142, total=145

AnGam_Sua5b_cells.rep

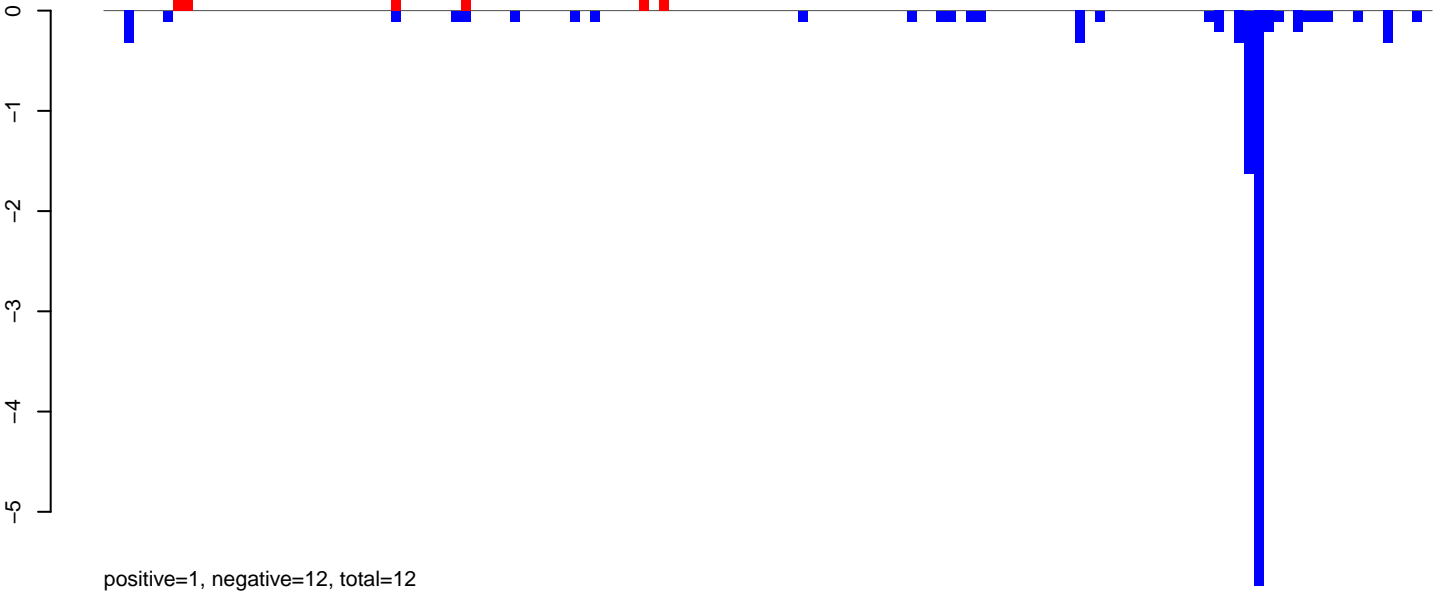


positive=6, negative=169, total=175

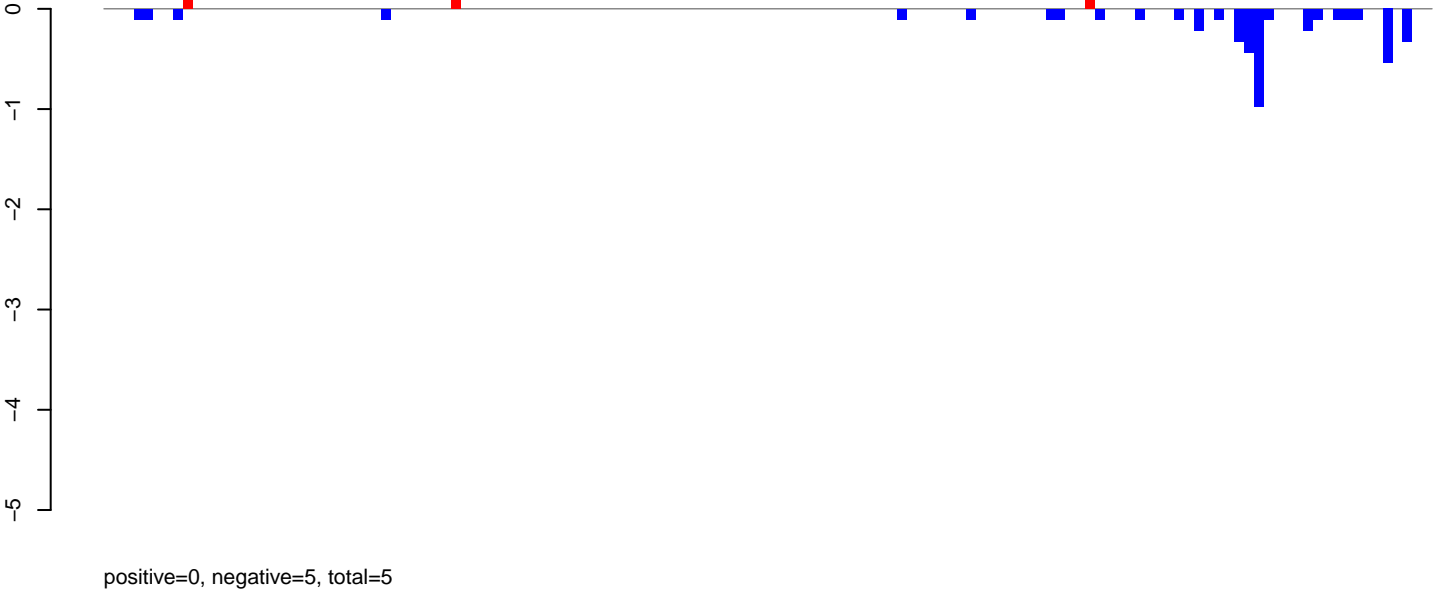
Window size=50, length=4693, TE@CR1-3_AG:1-4693



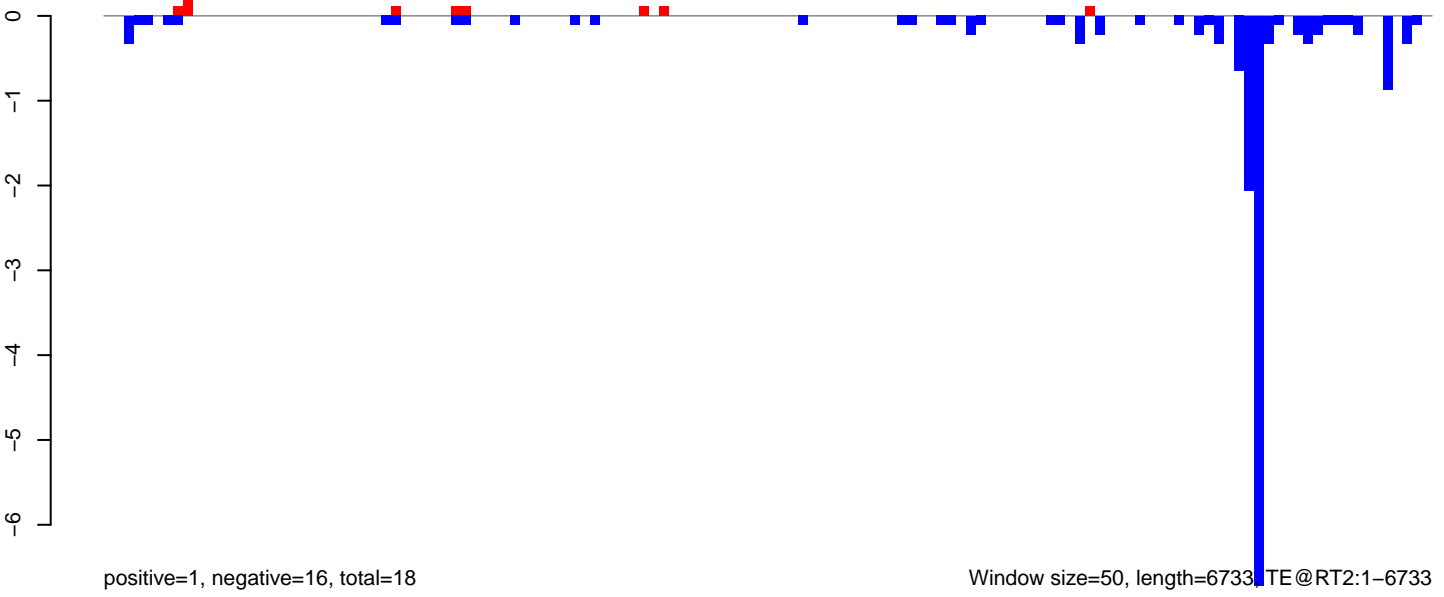
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



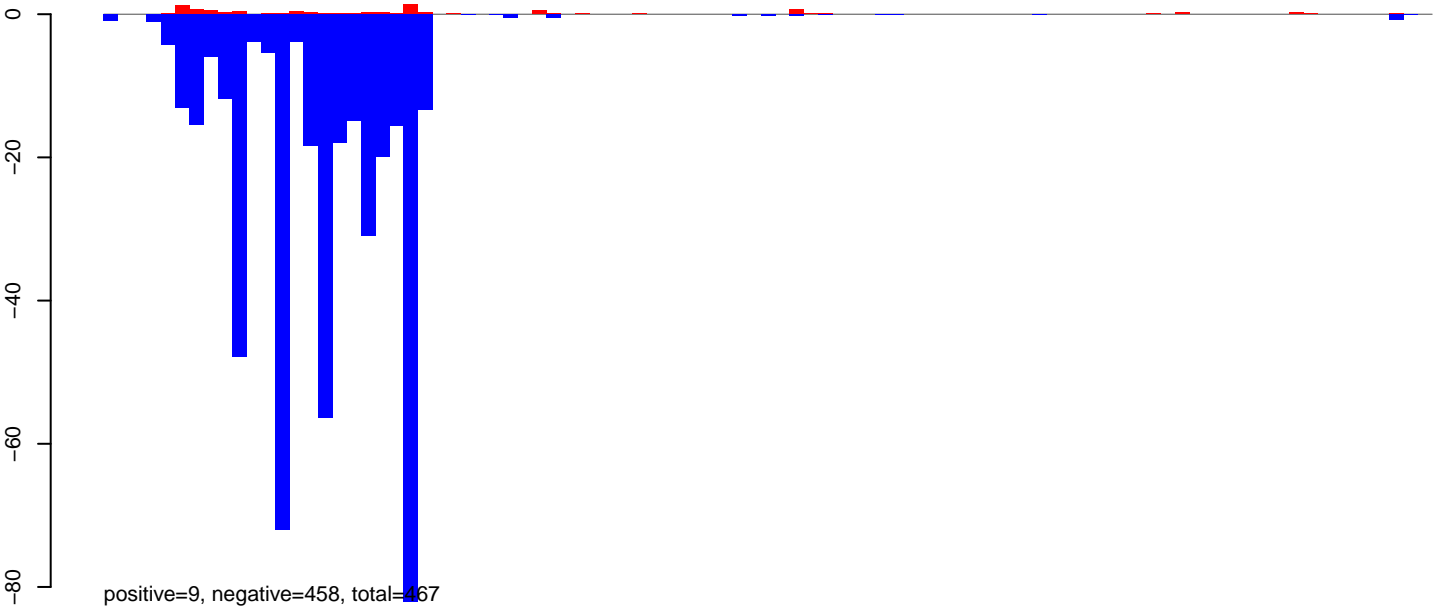
Window size=50, length=6733, TE@RT2:1-6733

0 1000 2000 3000 4000 5000 6000 7000

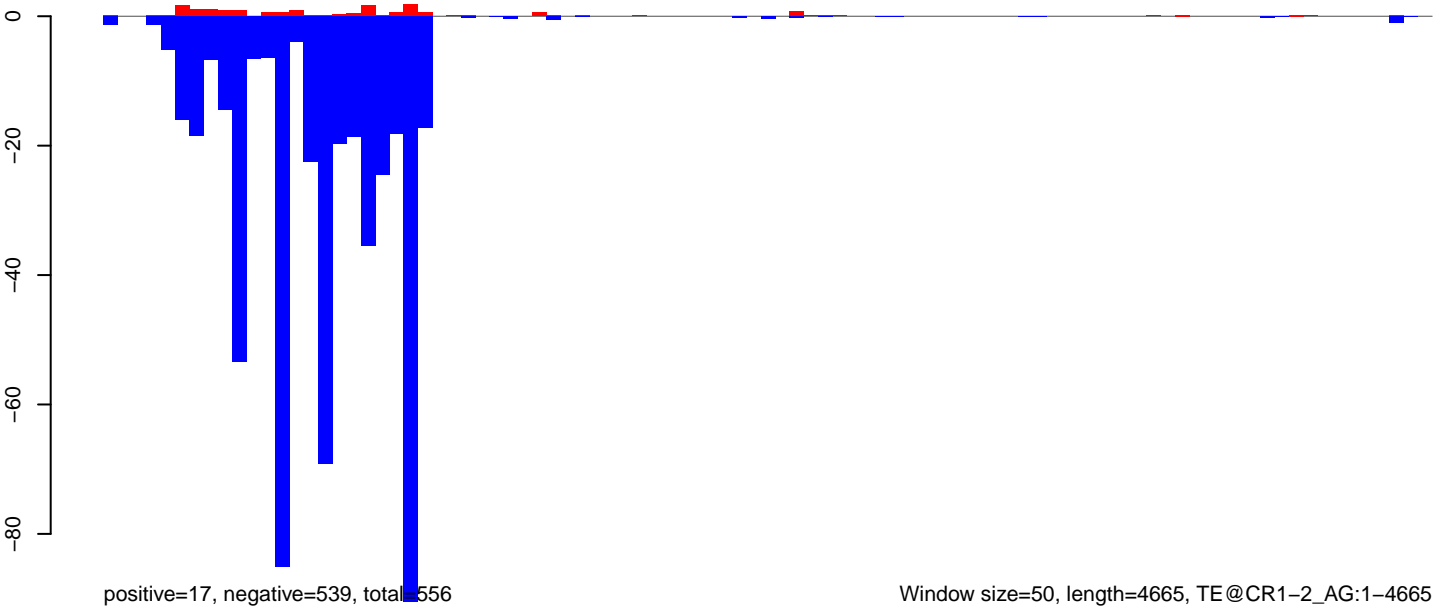
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



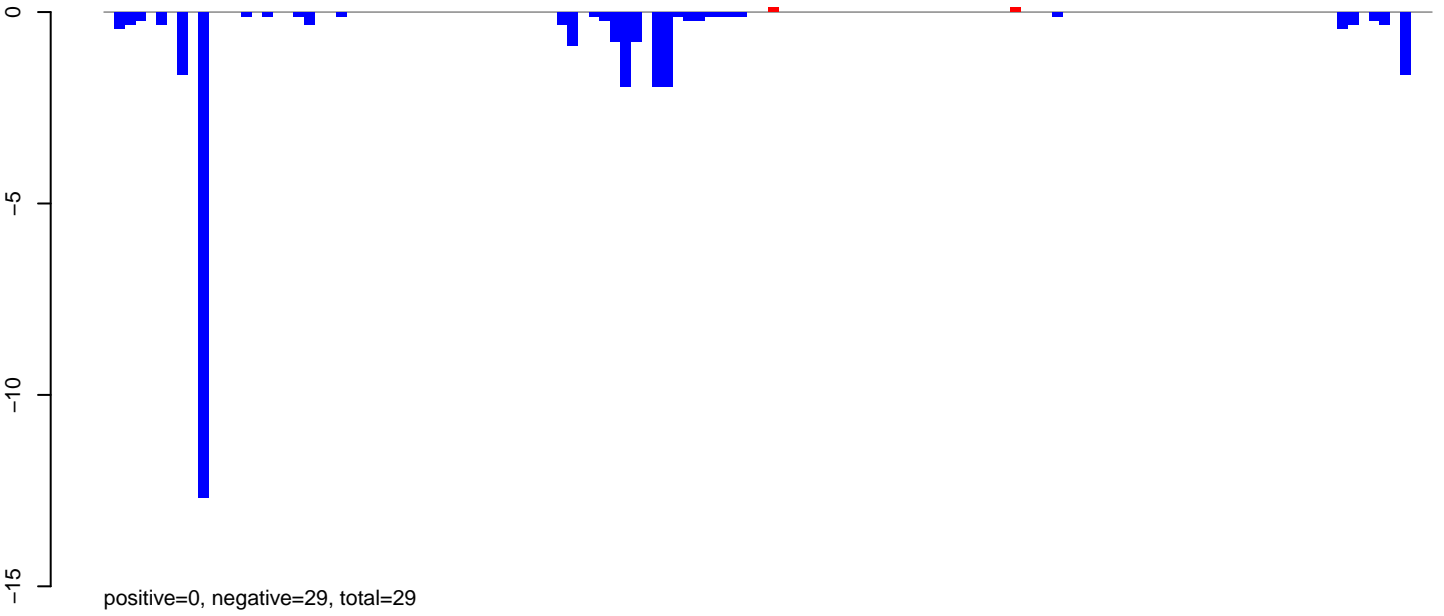
AnGam_Sua5b_cells.rep



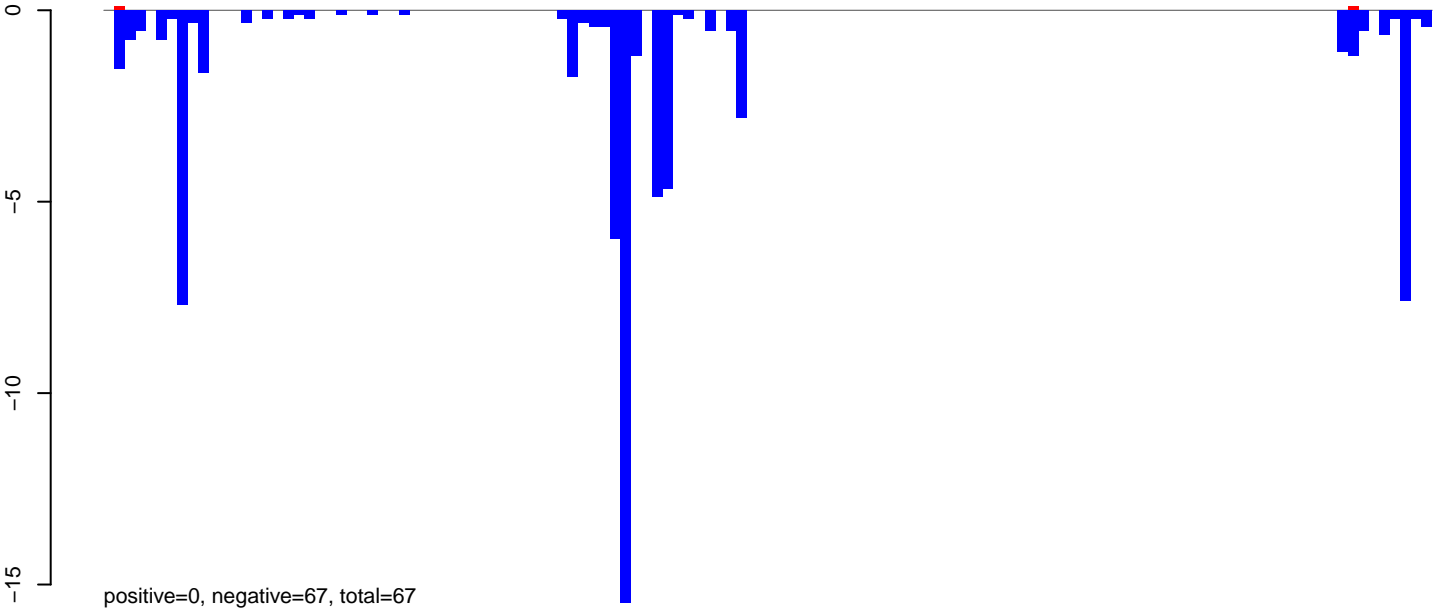
Window size=50, length=4665, TE@CR1-2_AG:1-4665

0 1000 2000 3000 4000

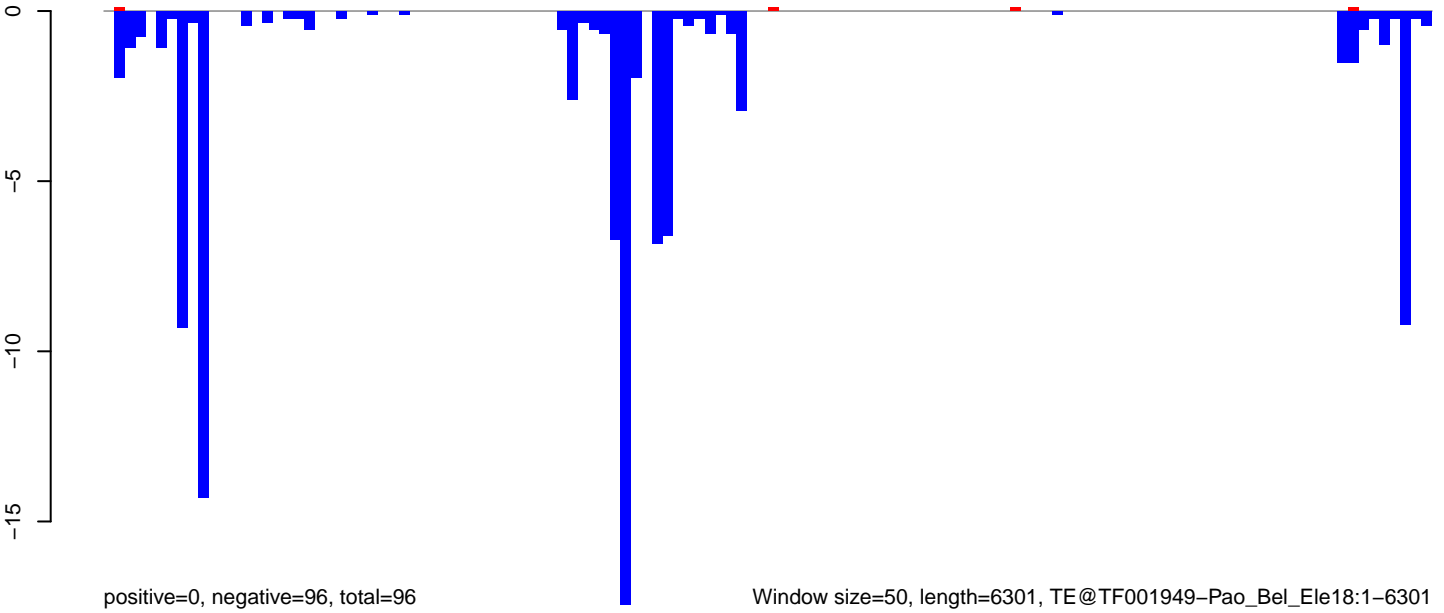
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

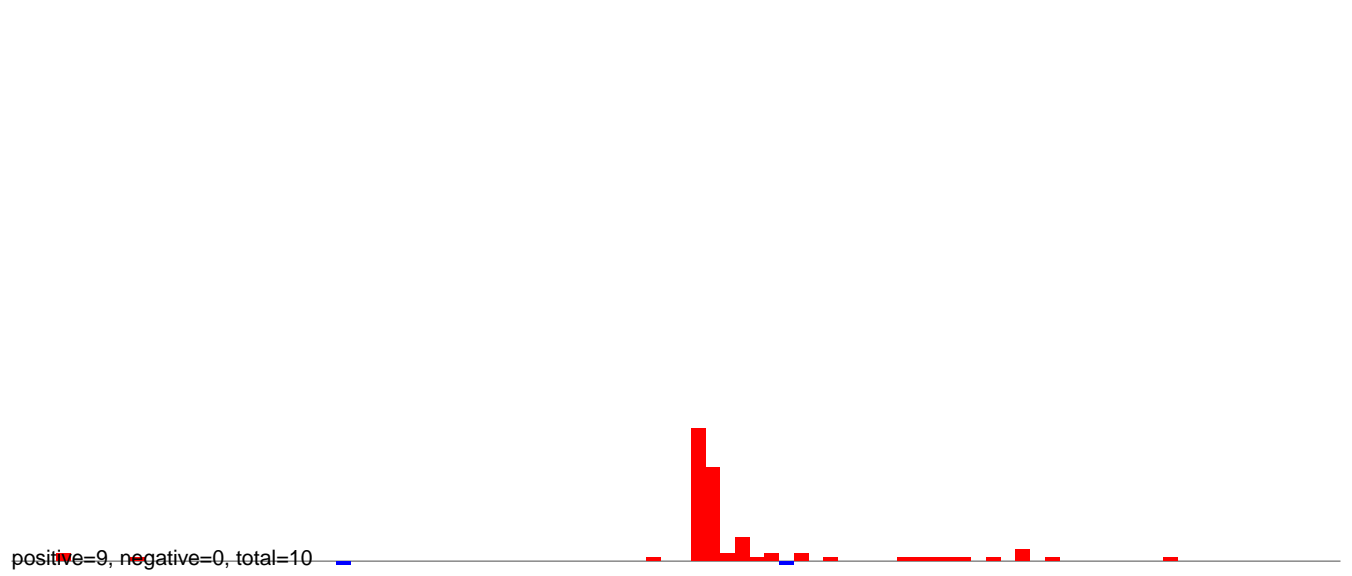


AnGam_Sua5b_cells.rep

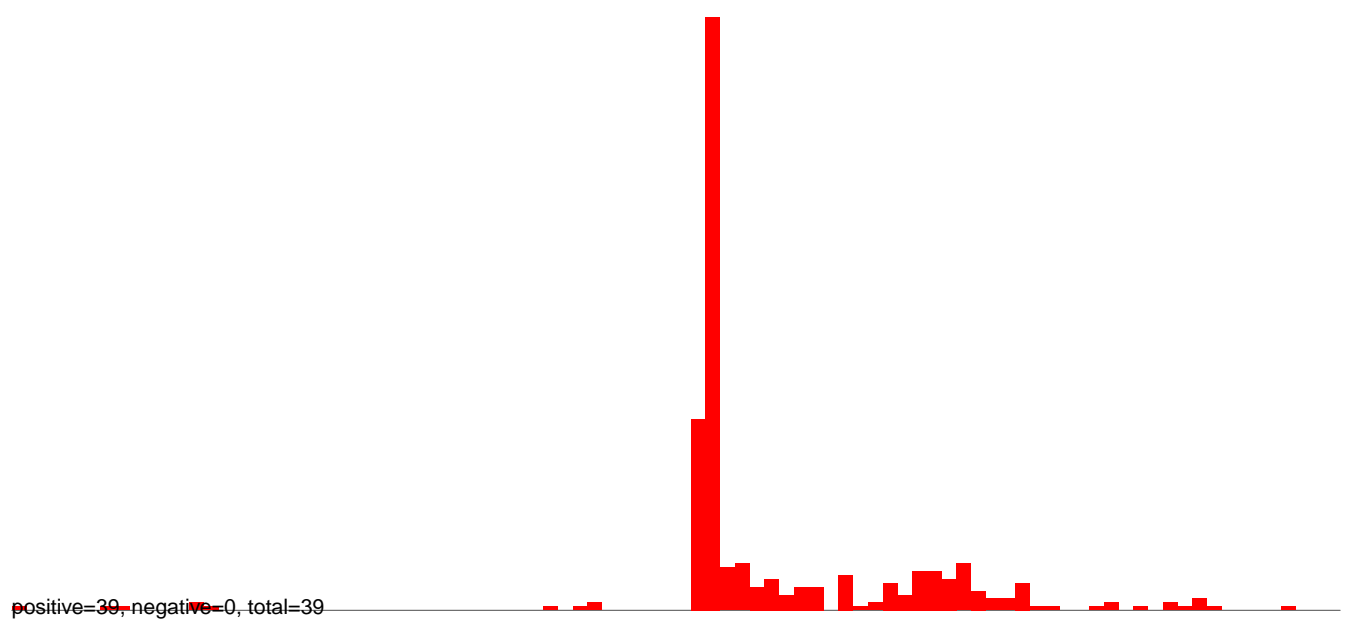


Window size=50, length=6301, TE@TF001949-Pao_Bel_Ele18:1-6301

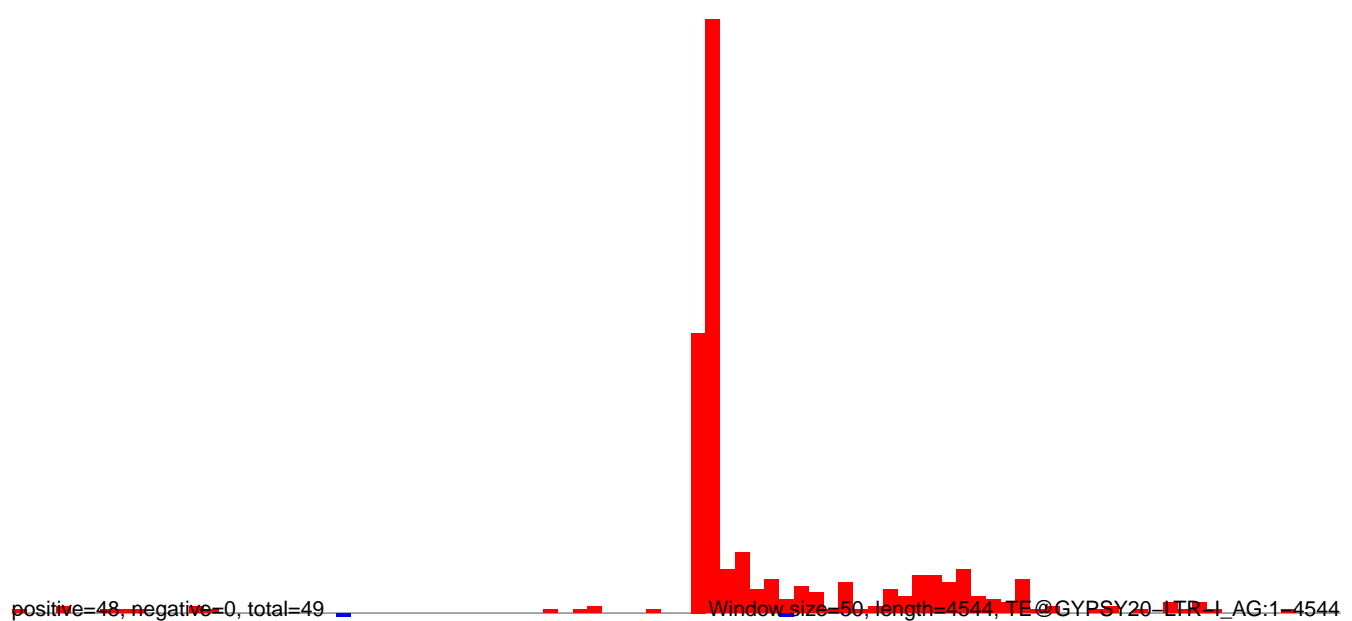
AnGam_Sua5b_cells.18_23.rep



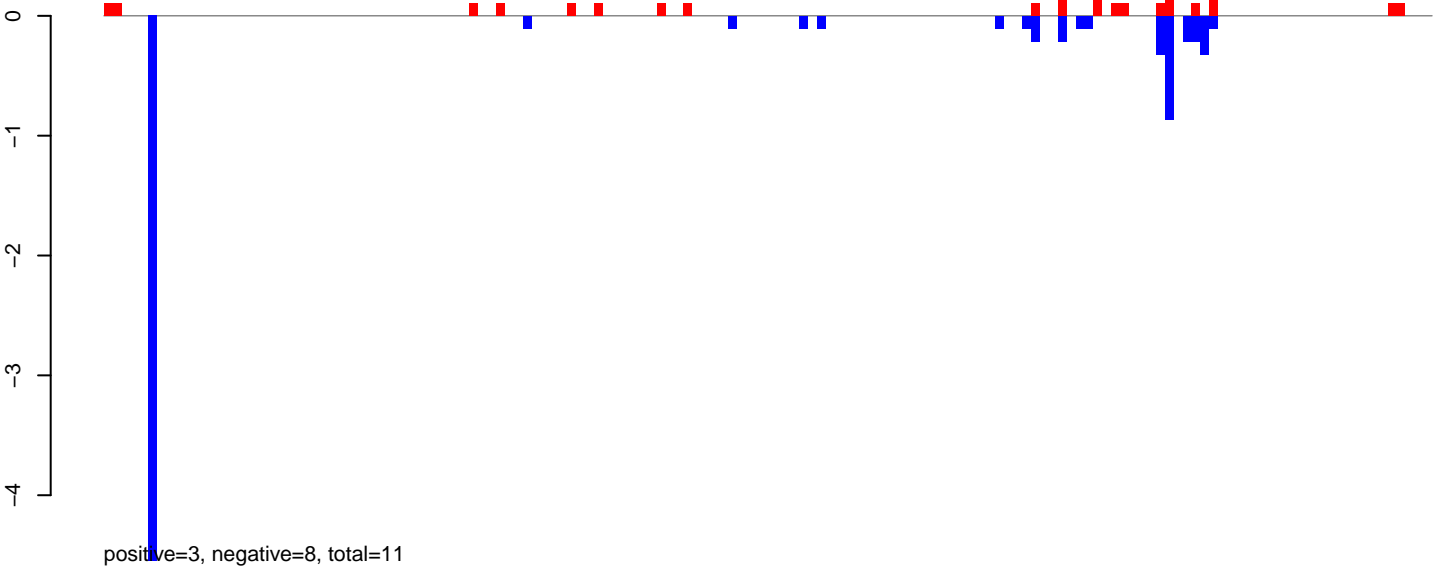
AnGam_Sua5b_cells.24_35.rep



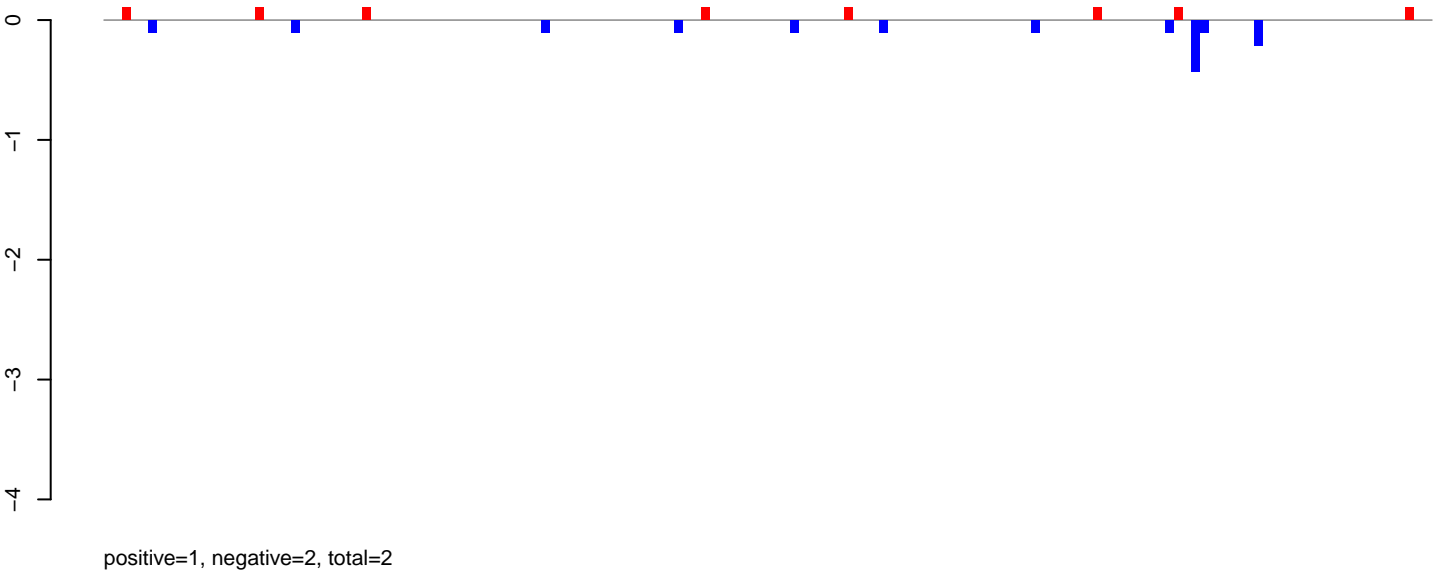
AnGam_Sua5b_cells.rep



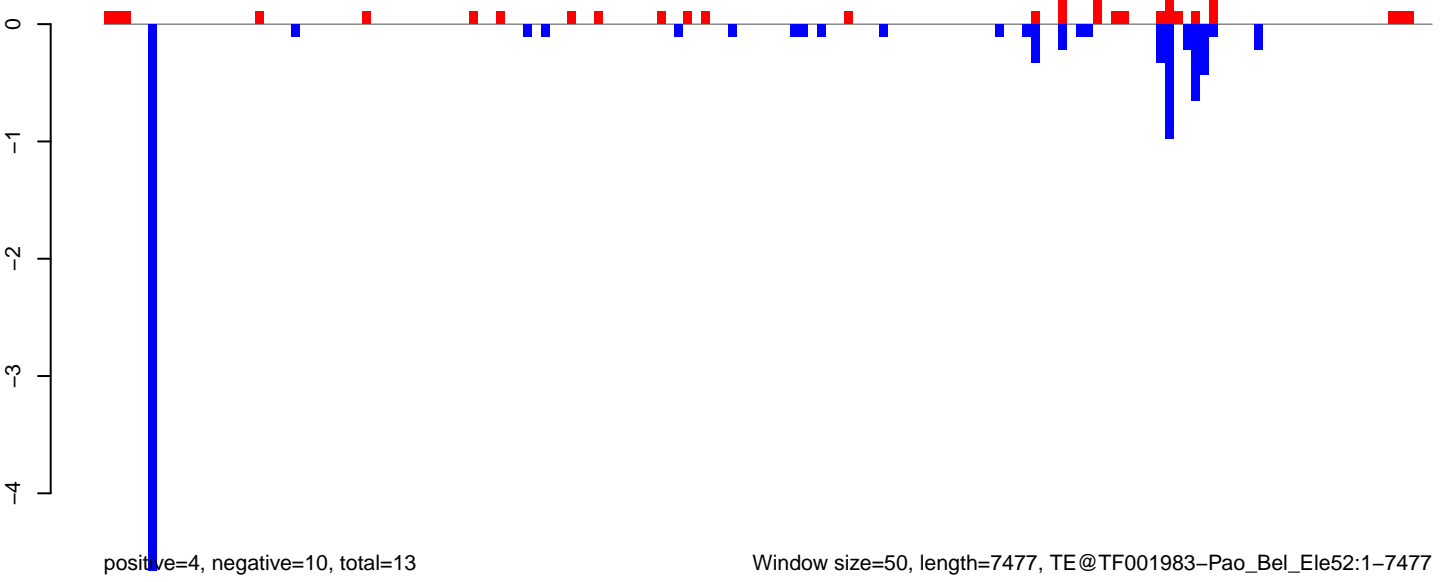
AnGam_Sua5b_cells.18_23.rep



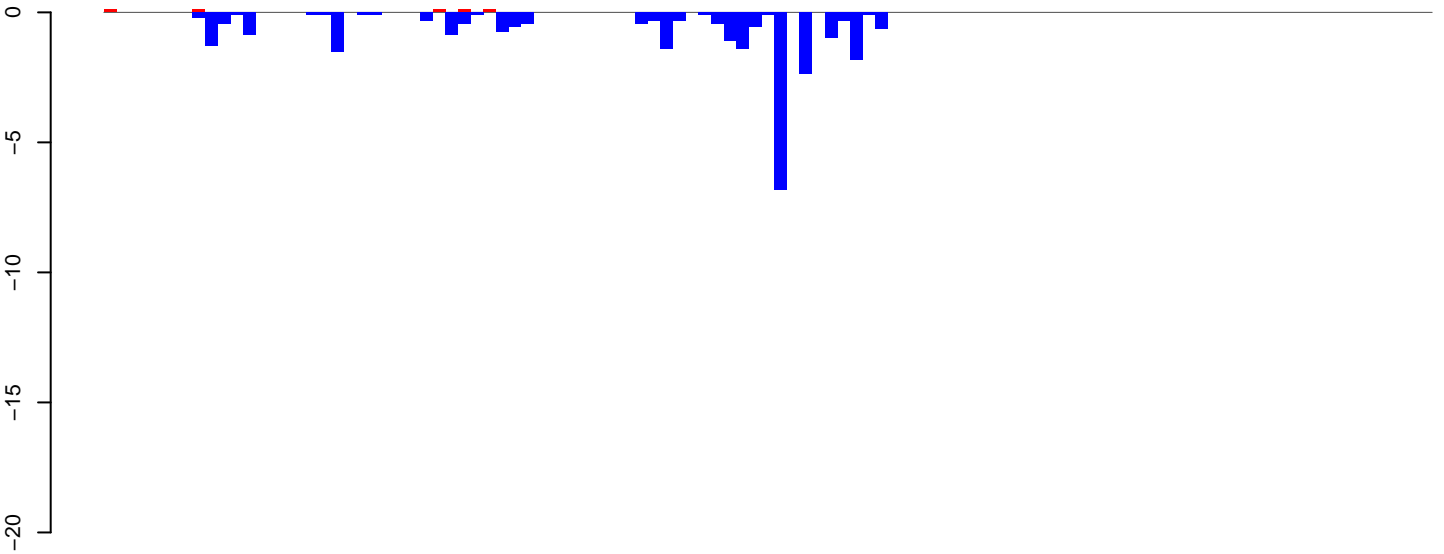
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

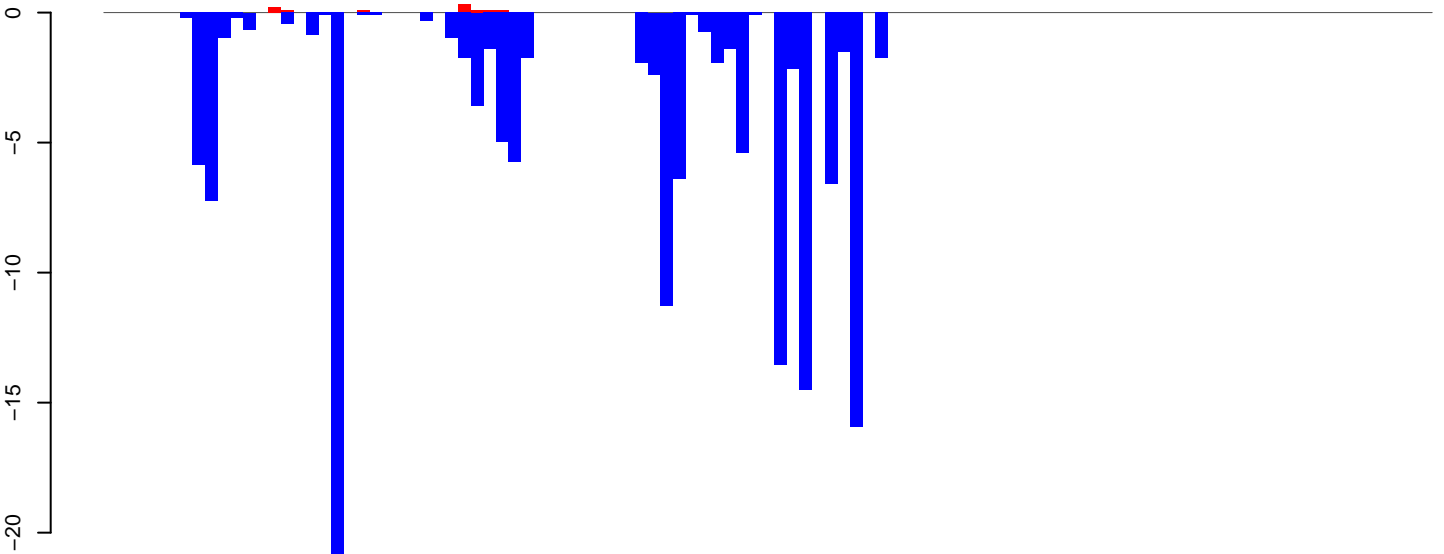


AnGam_Sua5b_cells.18_23.rep



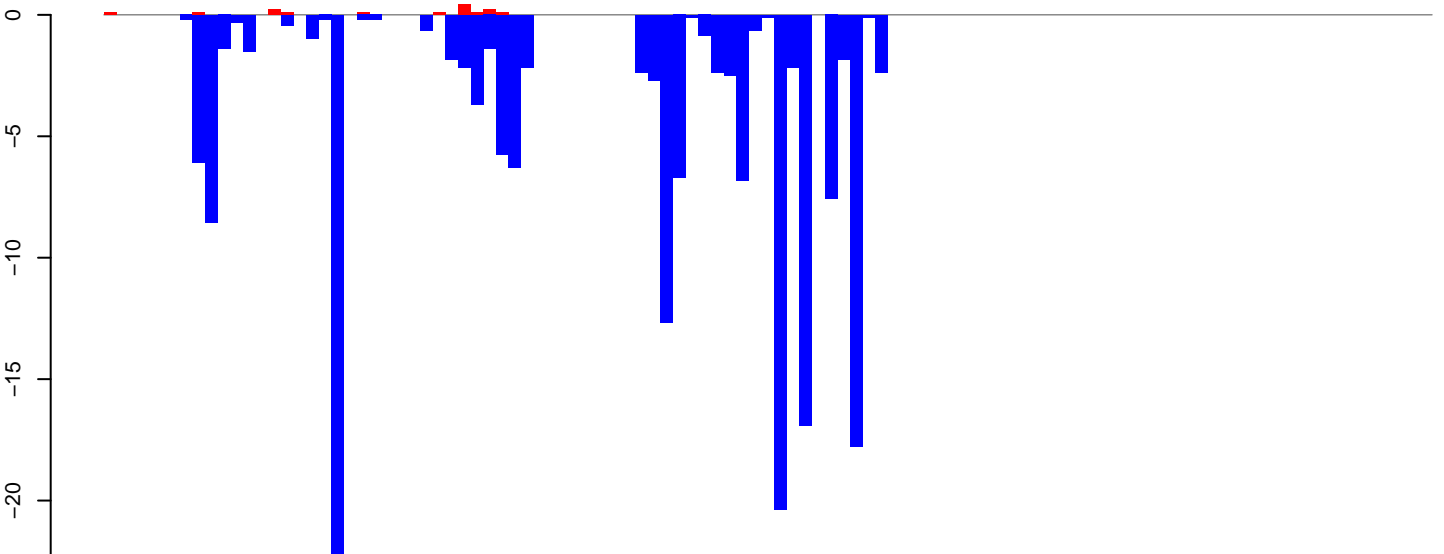
positive=1, negative=28, total=28

AnGam_Sua5b_cells.24_35.rep



positive=1, negative=148, total=149

AnGam_Sua5b_cells.rep

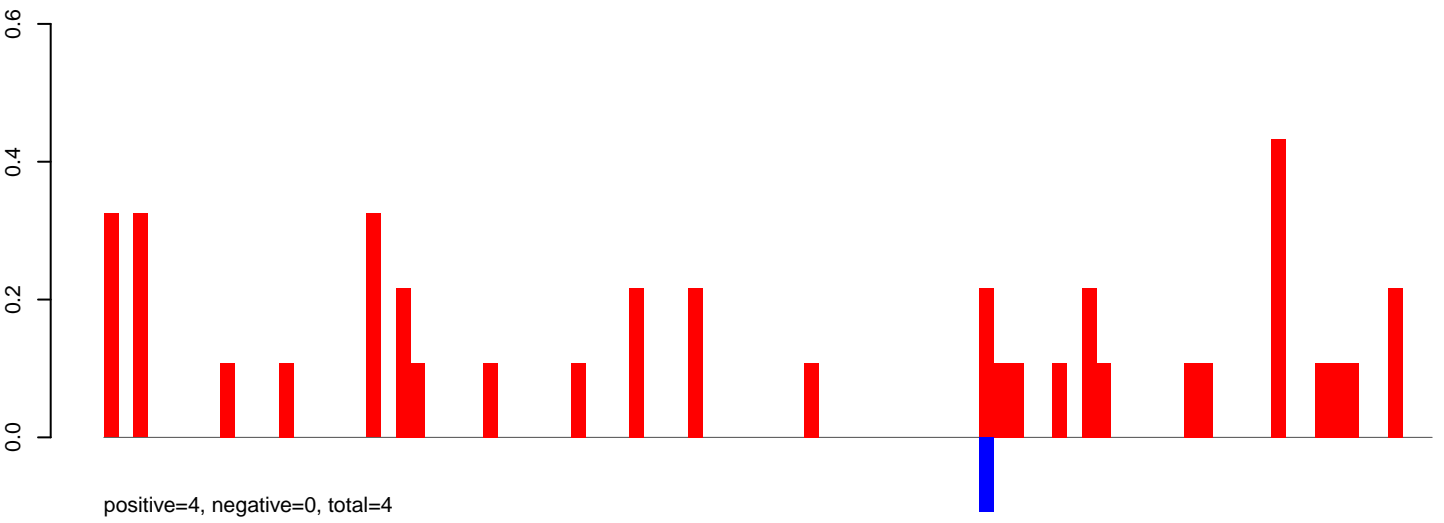


positive=2, negative=175, total=177

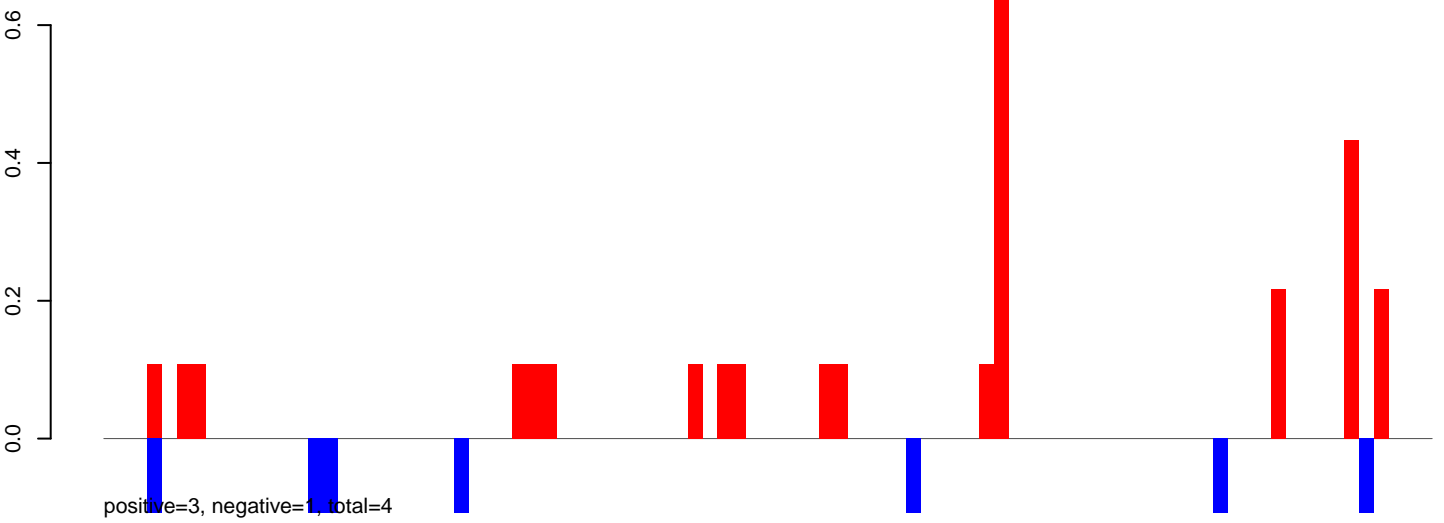
Window size=50, length=5289, TE@R6Ag3:1-5289

0 1000 2000 3000 4000 5000

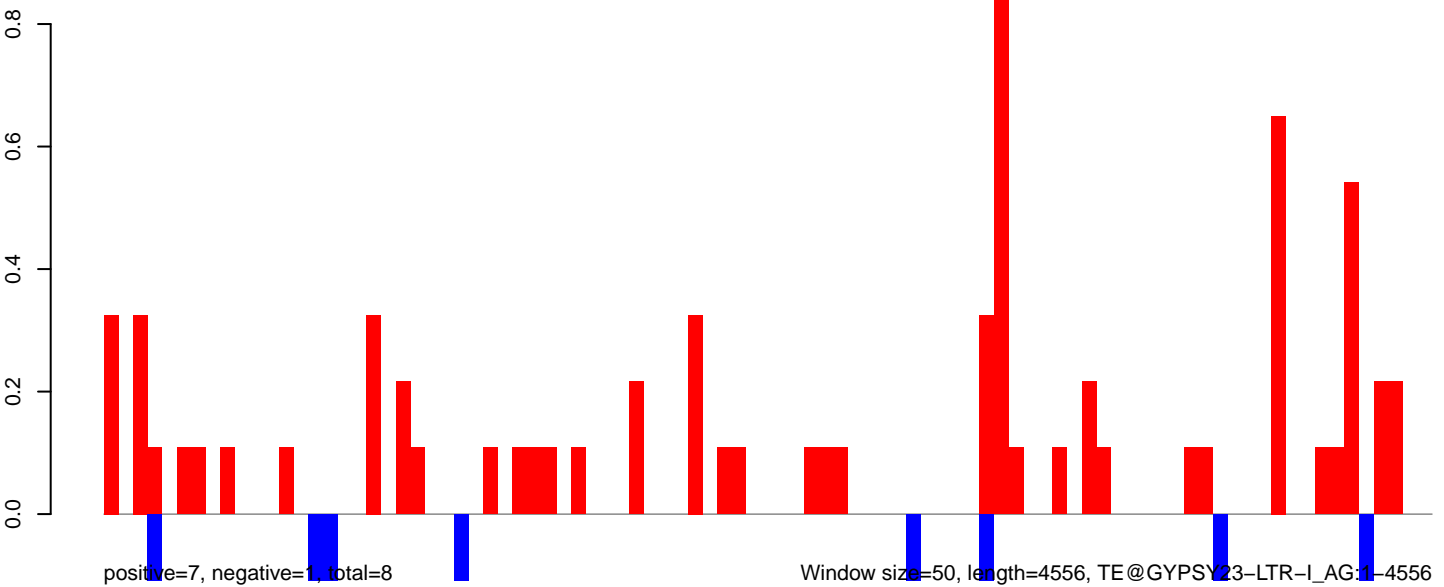
AnGam_Sua5b_cells.18_23.rep



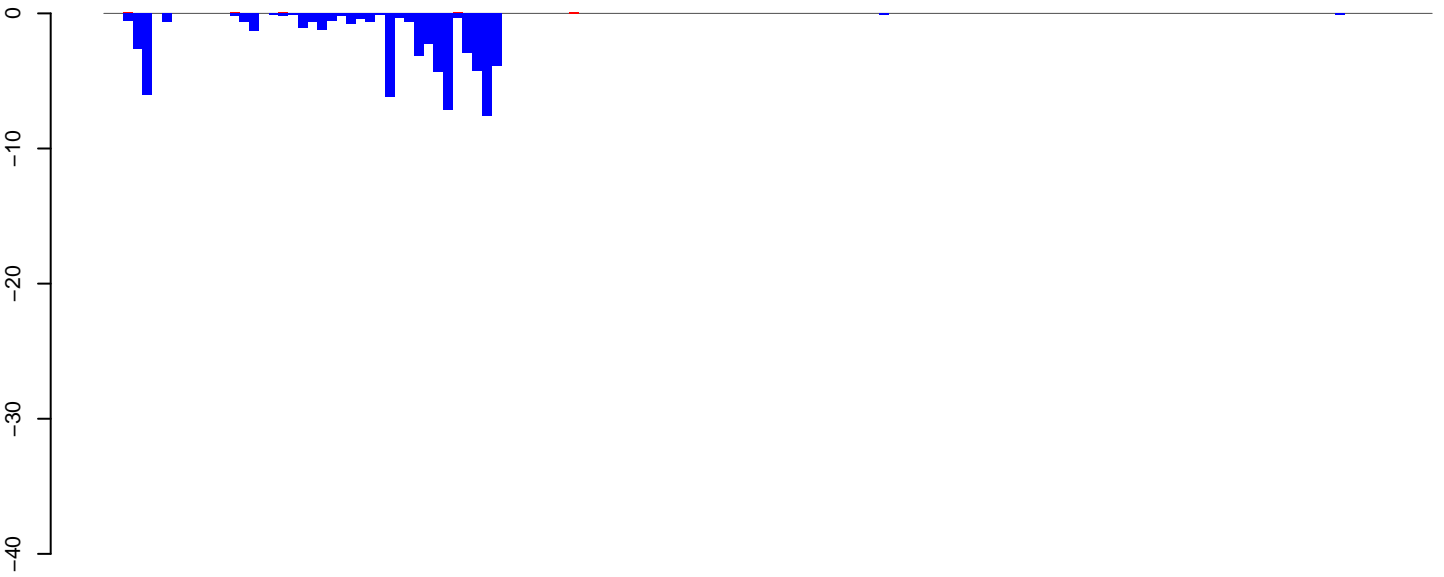
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

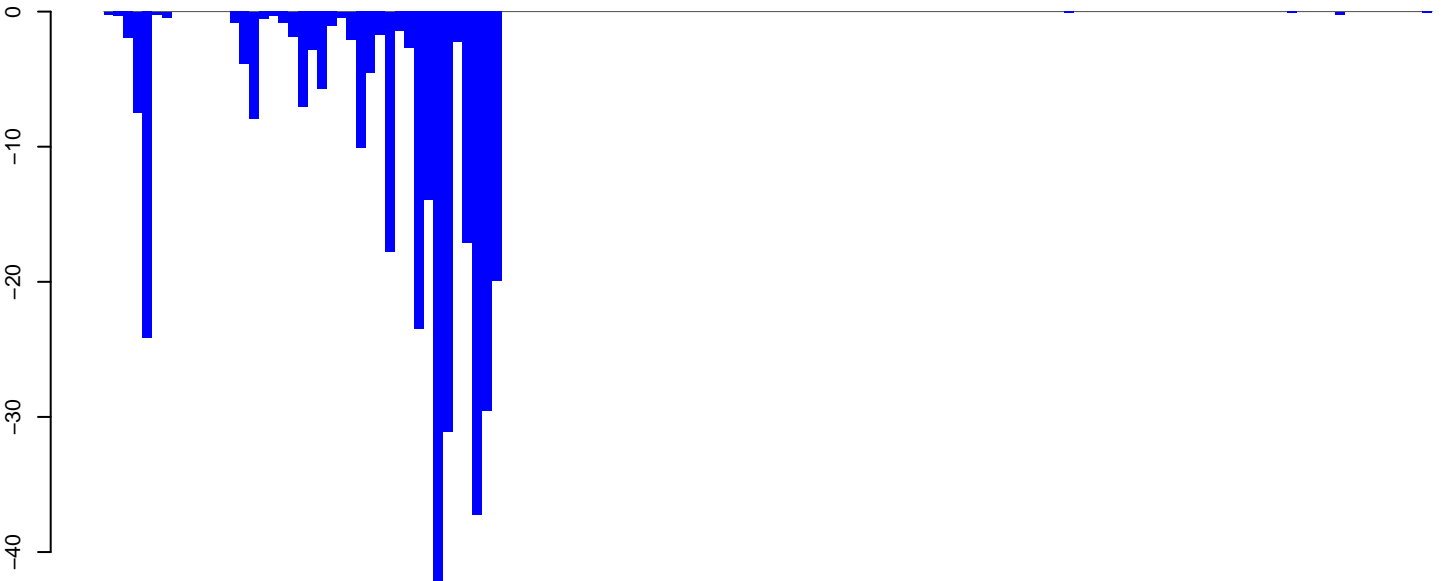


AnGam_Sua5b_cells.18_23.rep



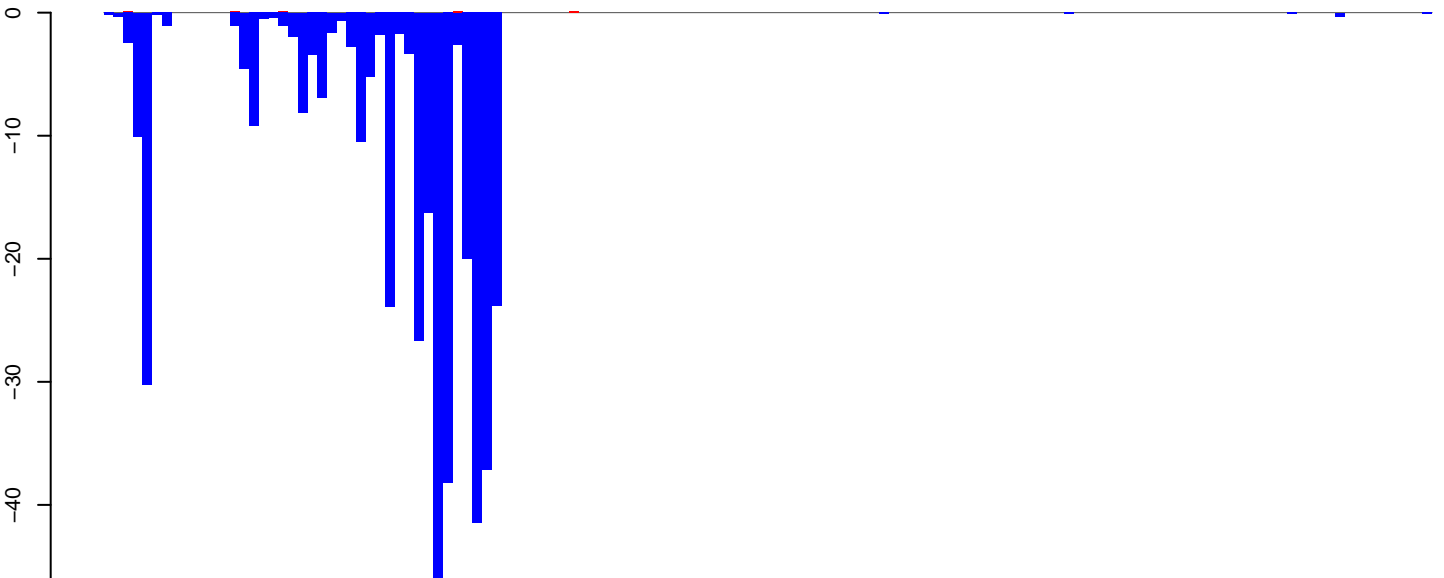
positive=1, negative=61, total=62

AnGam_Sua5b_cells.24_35.rep



positive=0, negative=328, total=328

AnGam_Sua5b_cells.rep

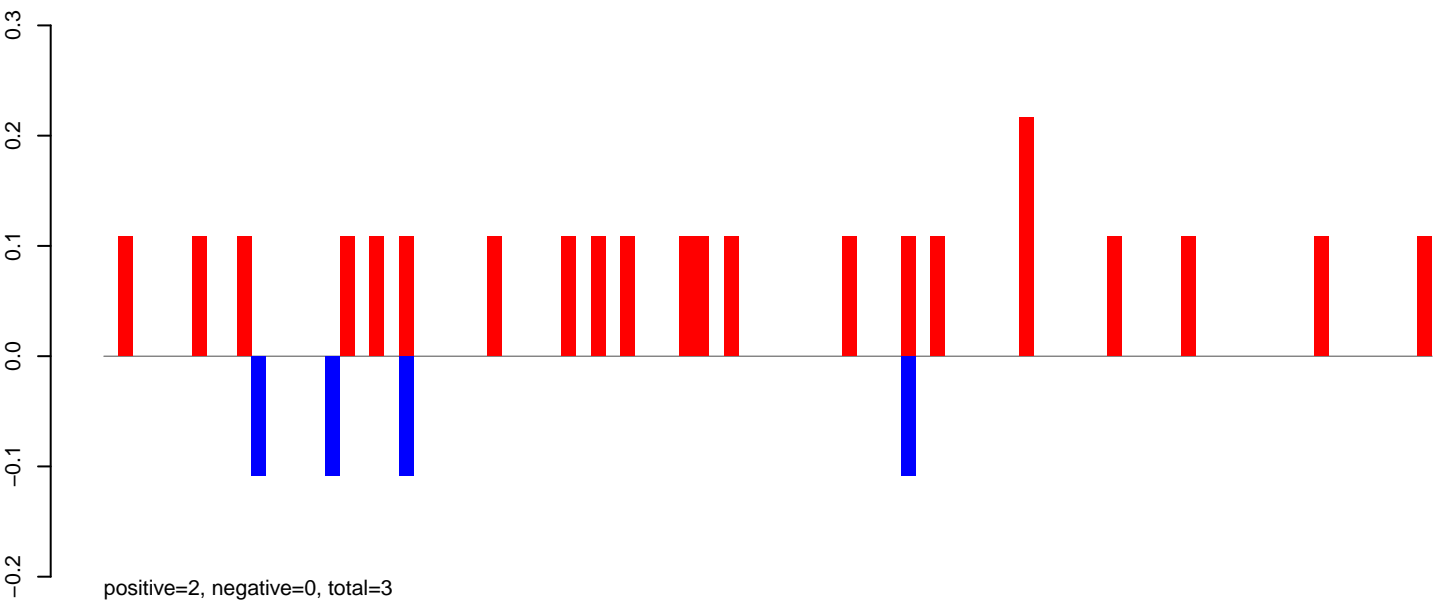


positive=1, negative=389, total=390

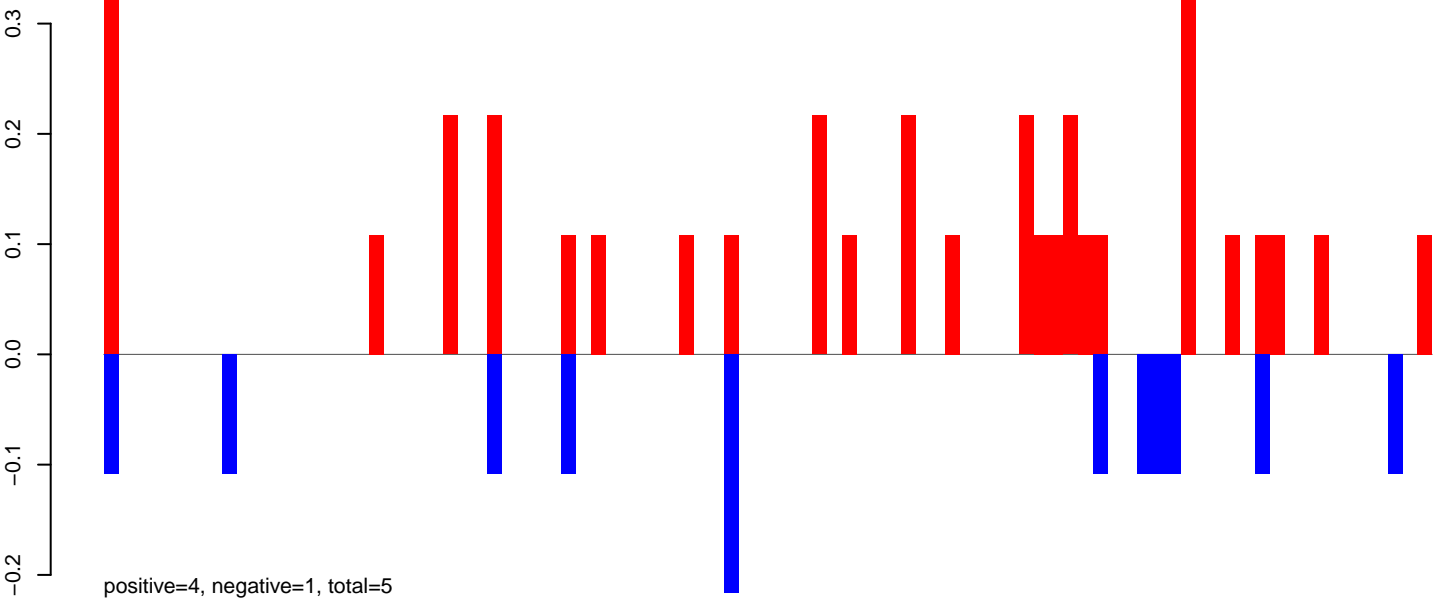
Window size=50, length=6861, TE@GYPSY7-LTR-I_AG:1-6861

0 1000 2000 3000 4000 5000 6000 7000

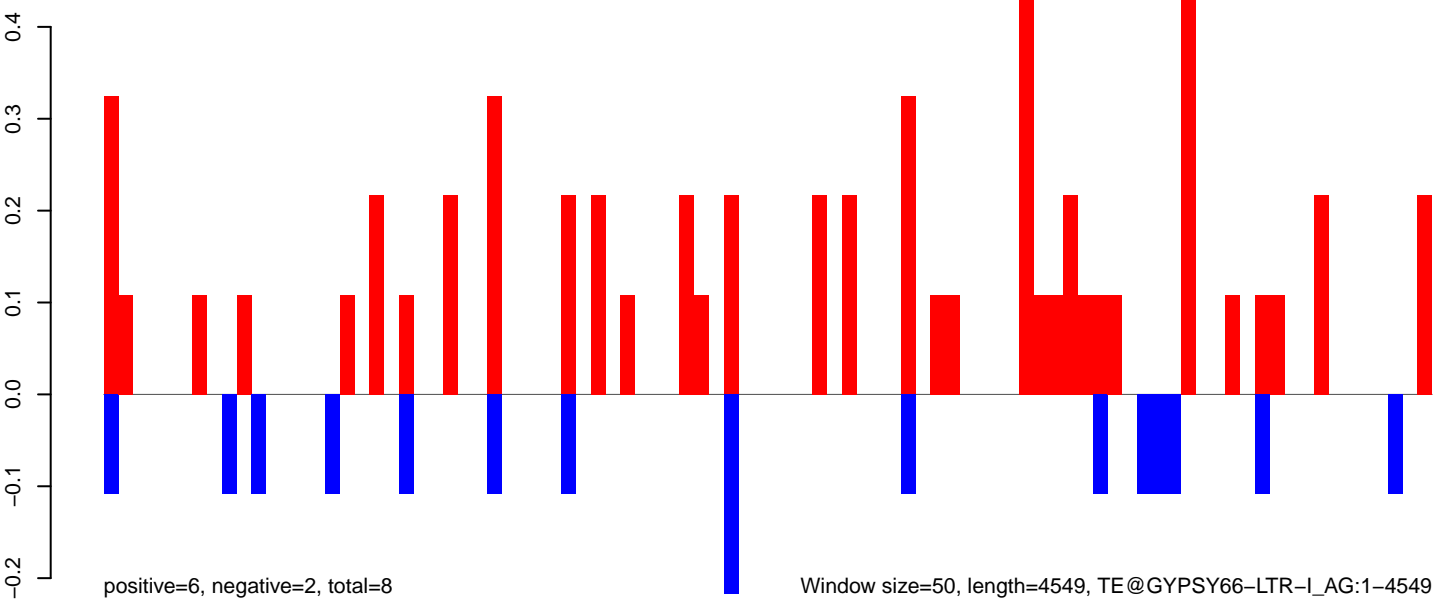
AnGam_Sua5b_cells.18_23.rep



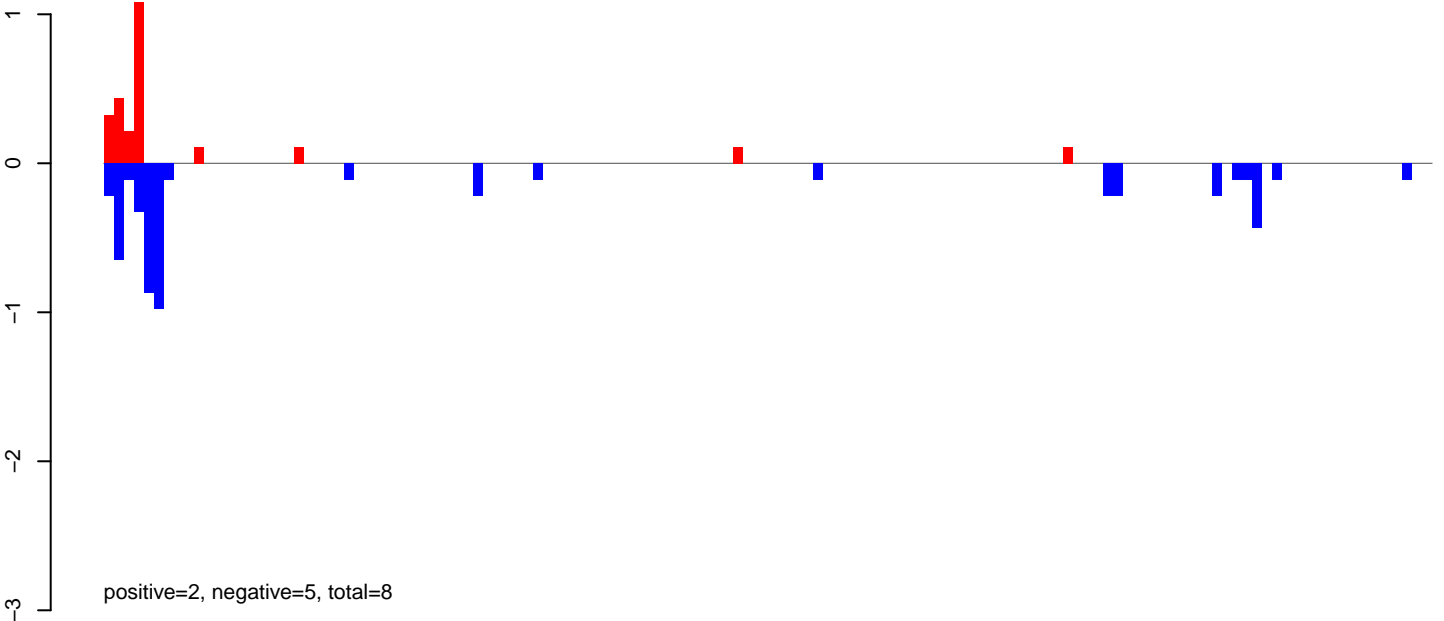
AnGam_Sua5b_cells.24_35.rep



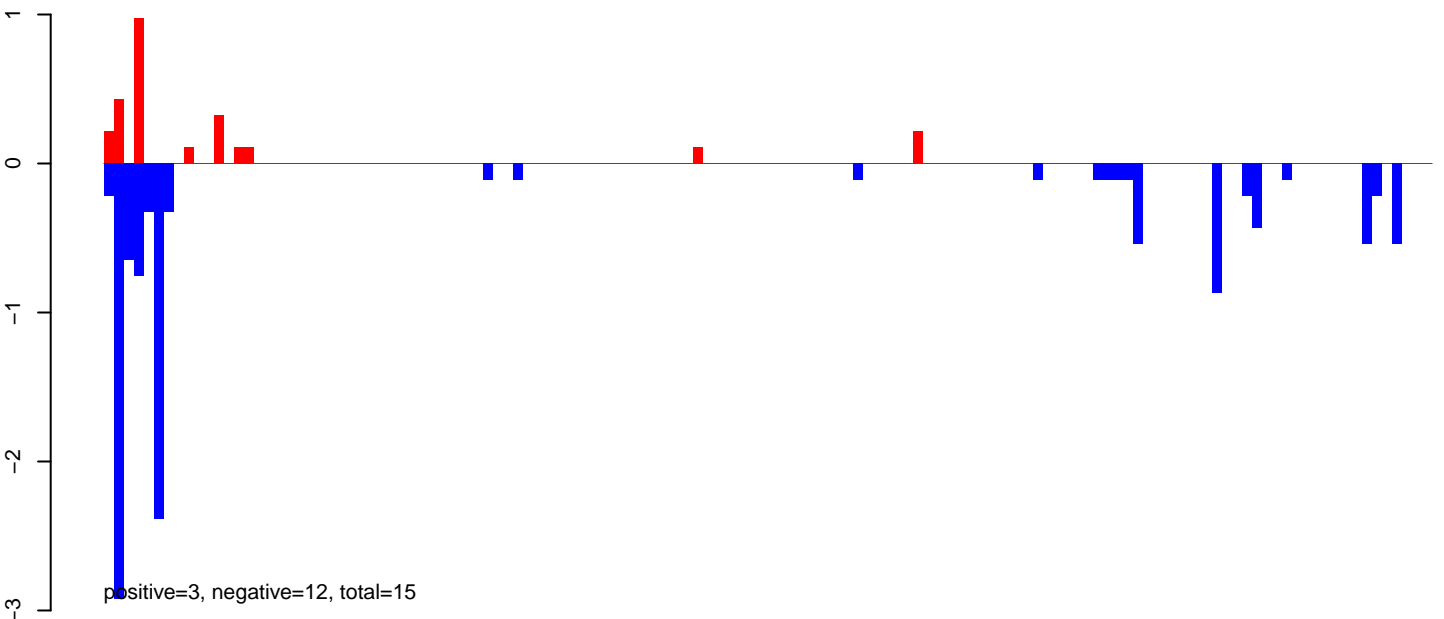
AnGam_Sua5b_cells.rep



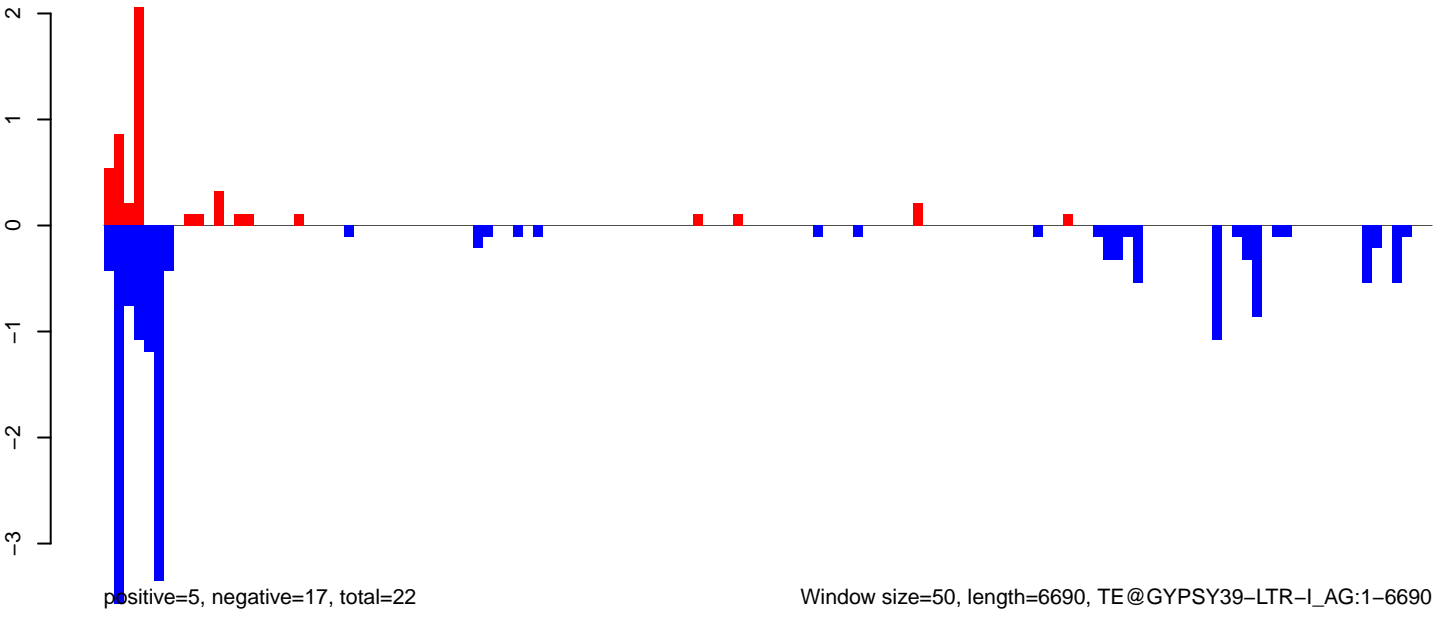
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

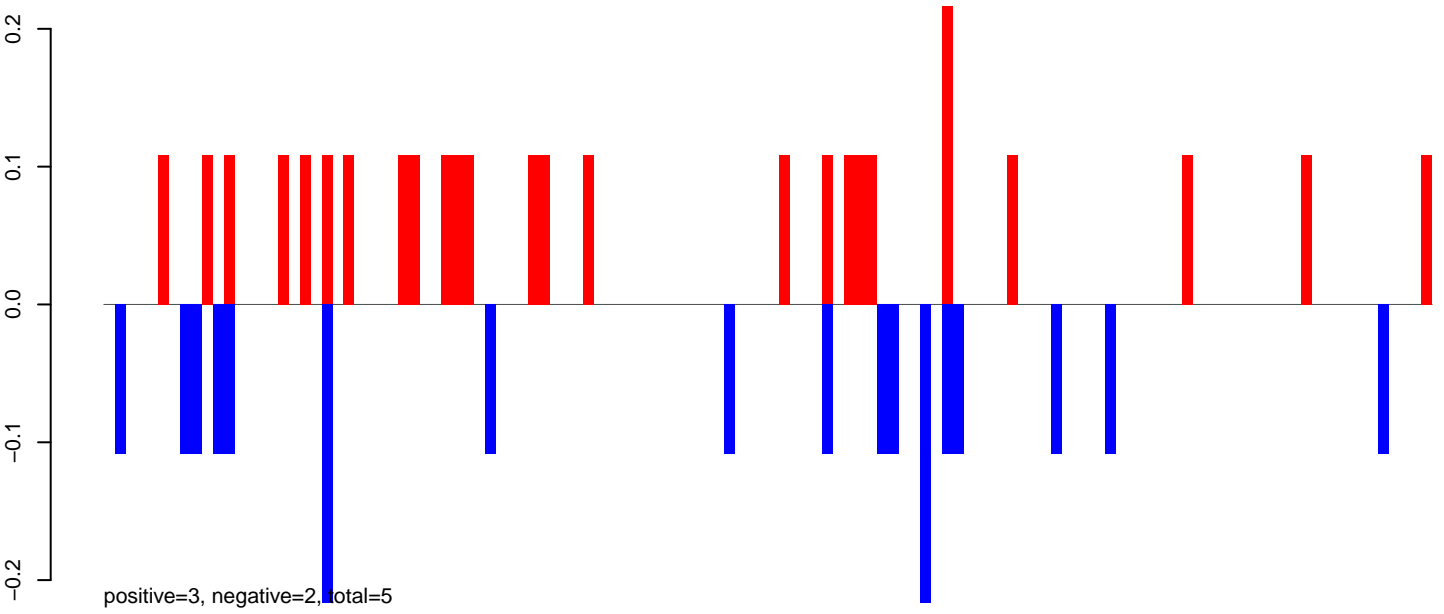


AnGam_Sua5b_cells.rep



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

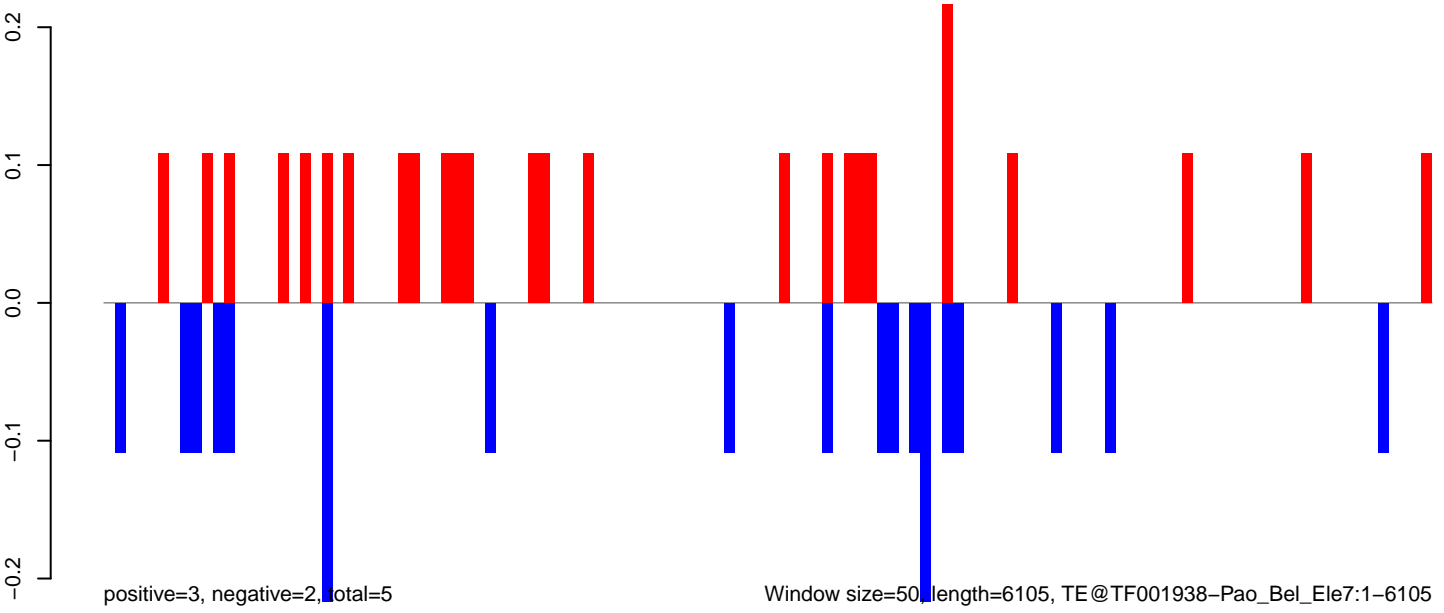
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

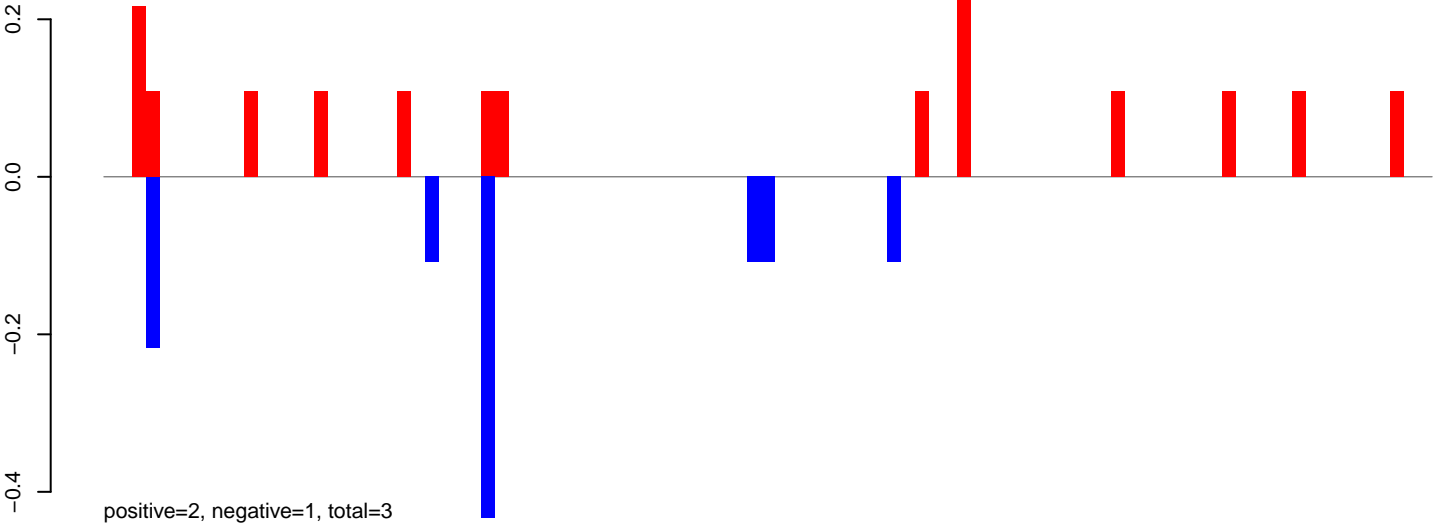


AnGam_Sua5b_cells.rep

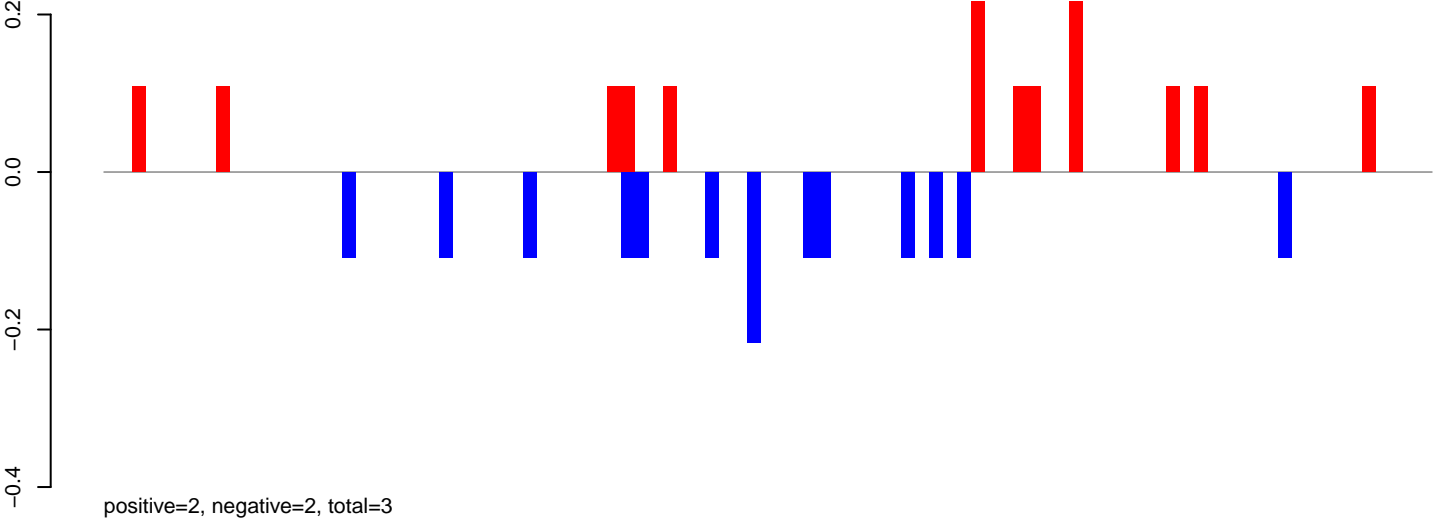


Window size=50, length=6105, TE@TF001938-Pao_Bel_Ele7:1-6105

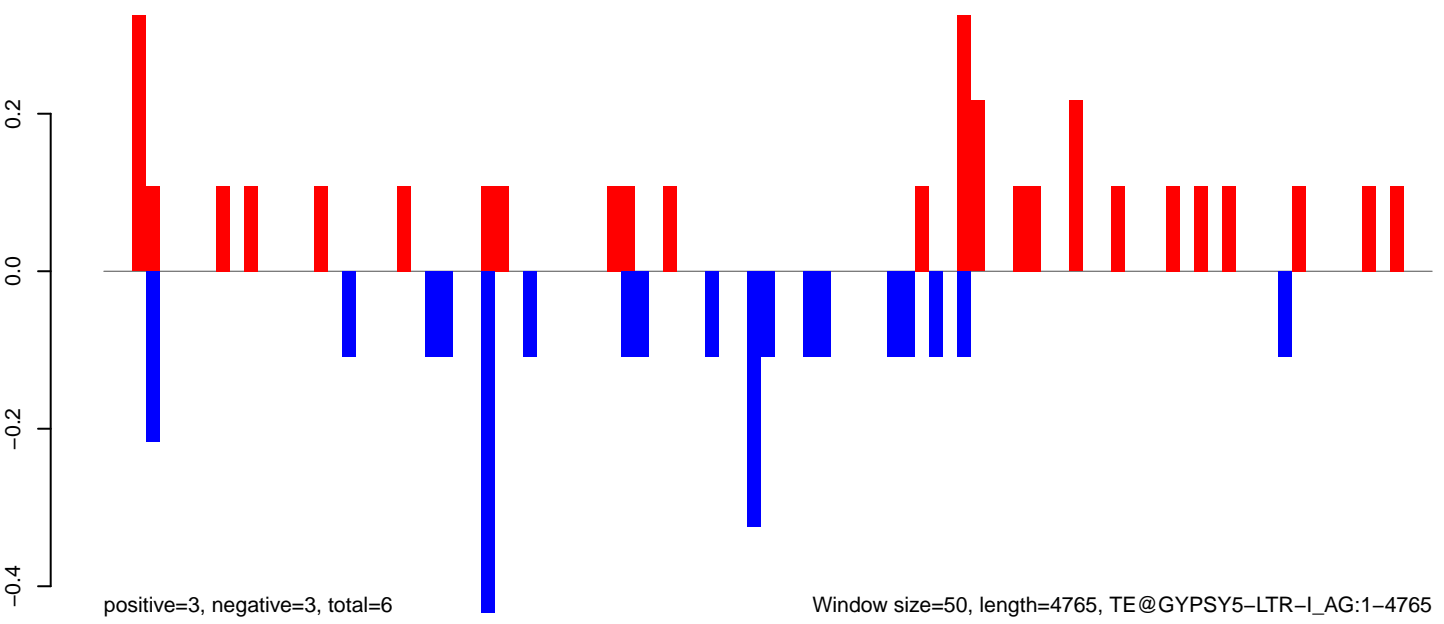
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

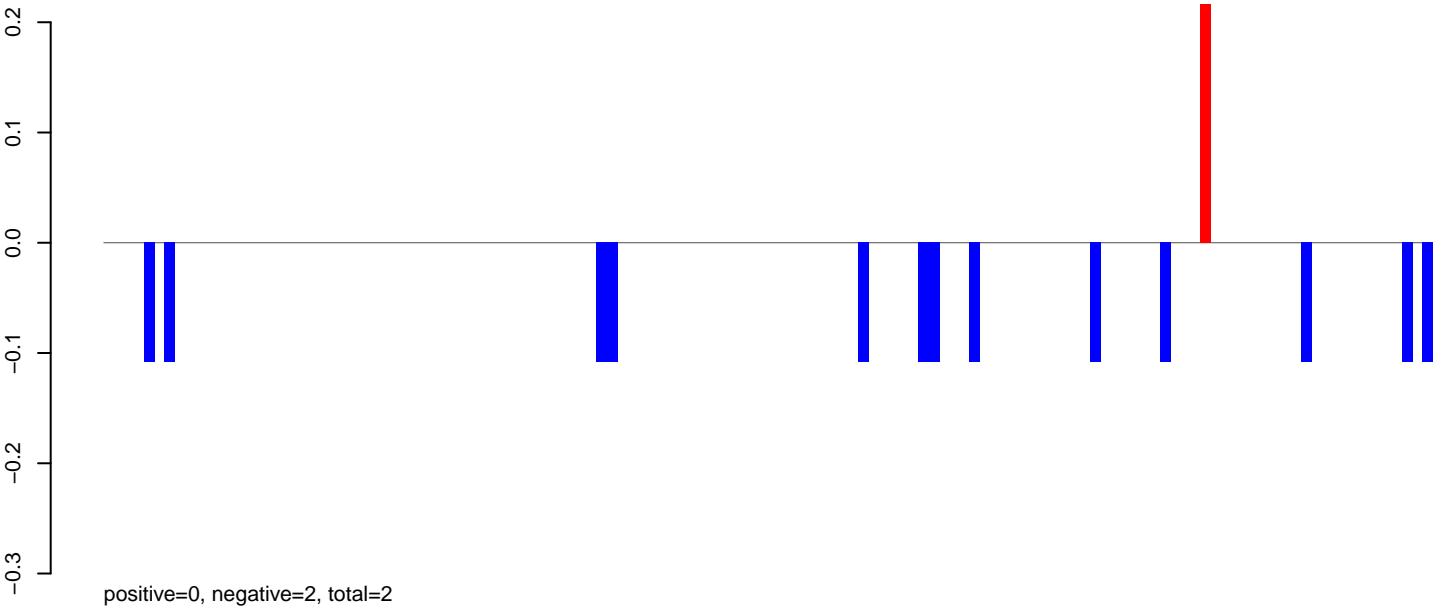


AnGam_Sua5b_cells.rep

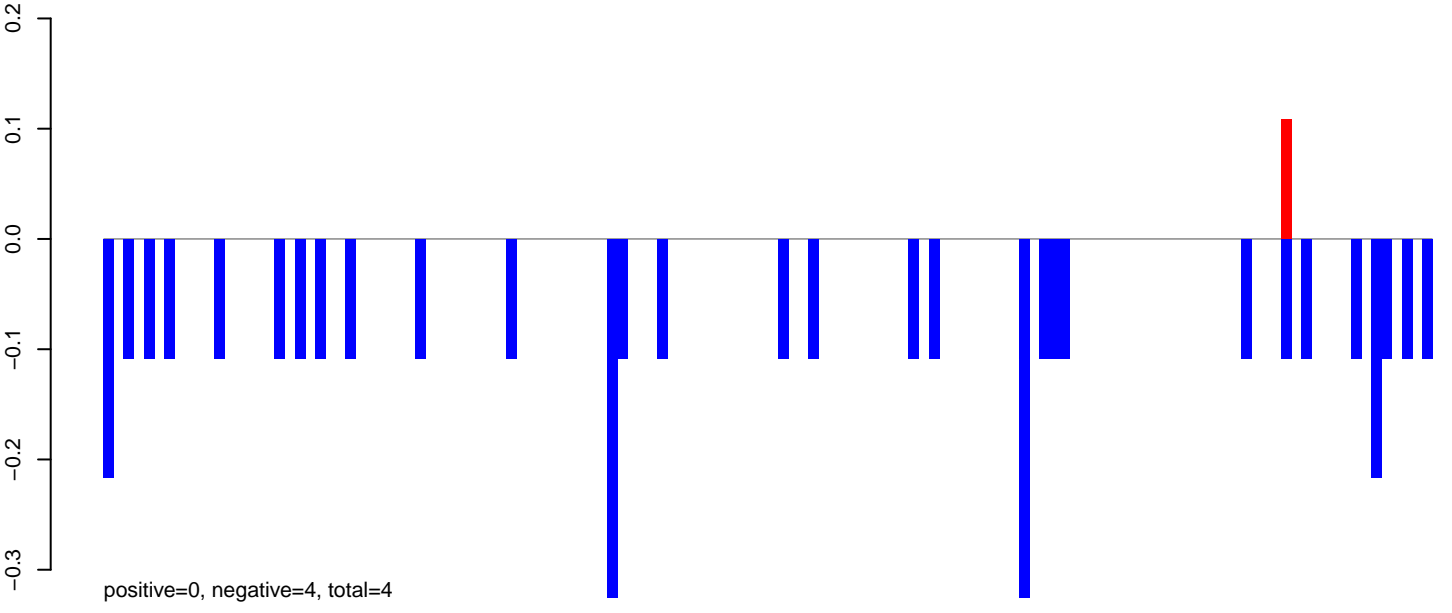


Window size=50, length=4765, TE@GYPSY5-LTR-I_AG:1-4765

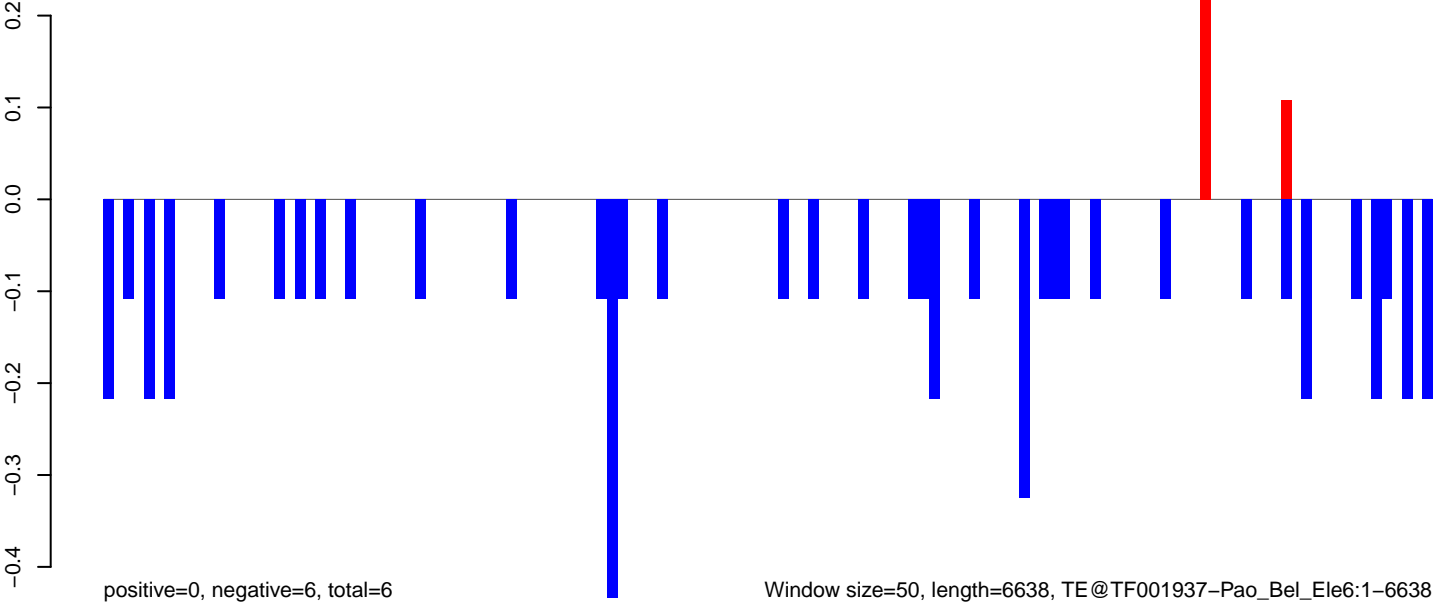
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

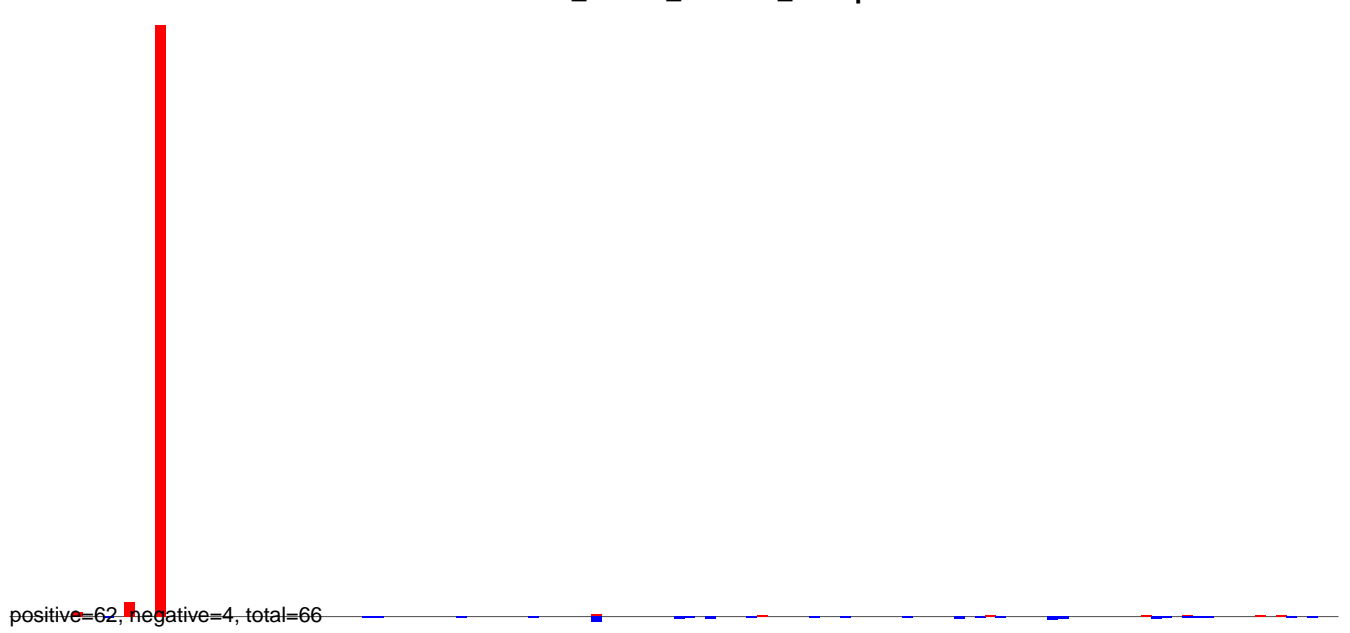


AnGam_Sua5b_cells.rep

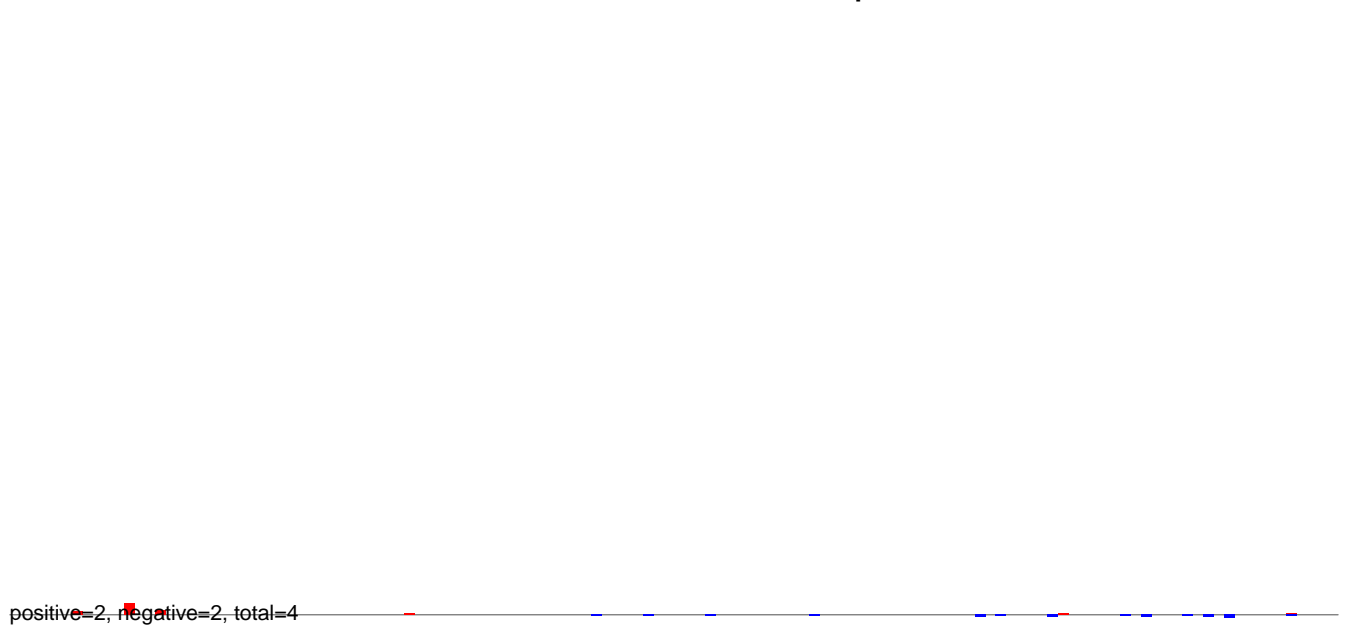


Window size=50, length=6638, TE@TF001937-Pao_Bel_Ele6:1-6638

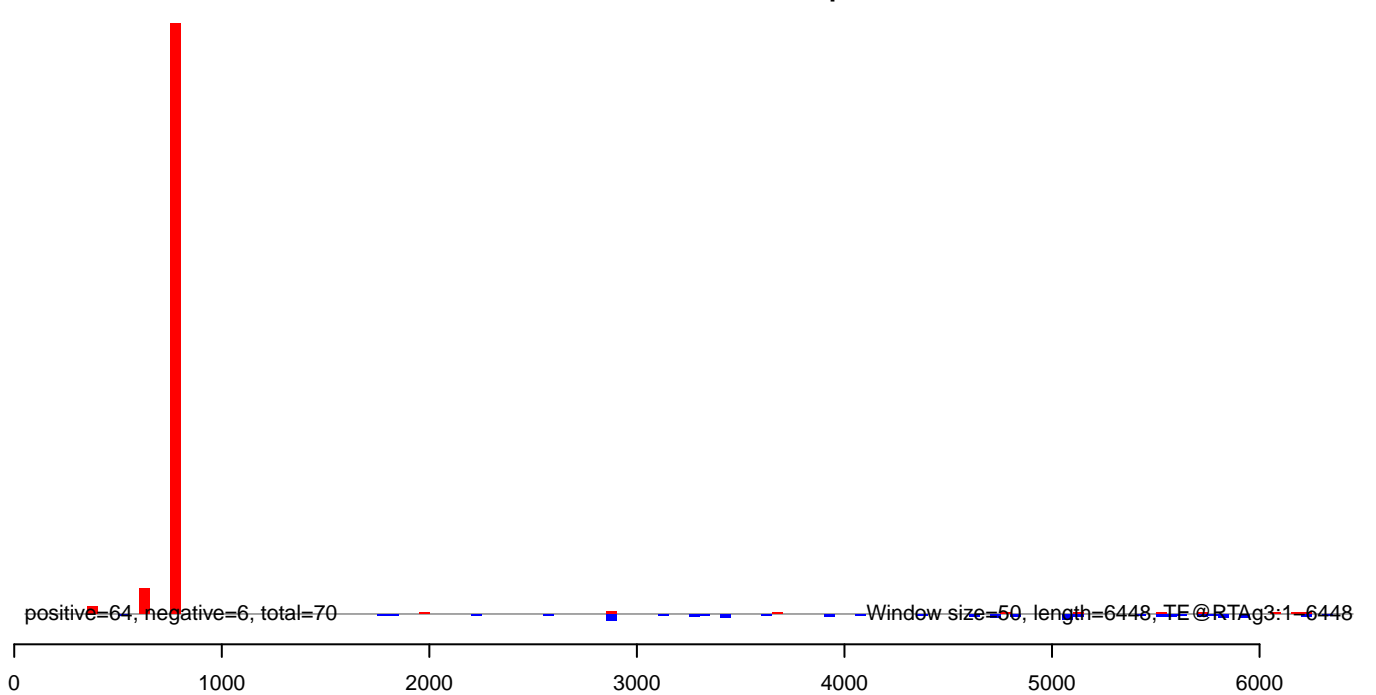
AnGam_Sua5b_cells.18_23.rep



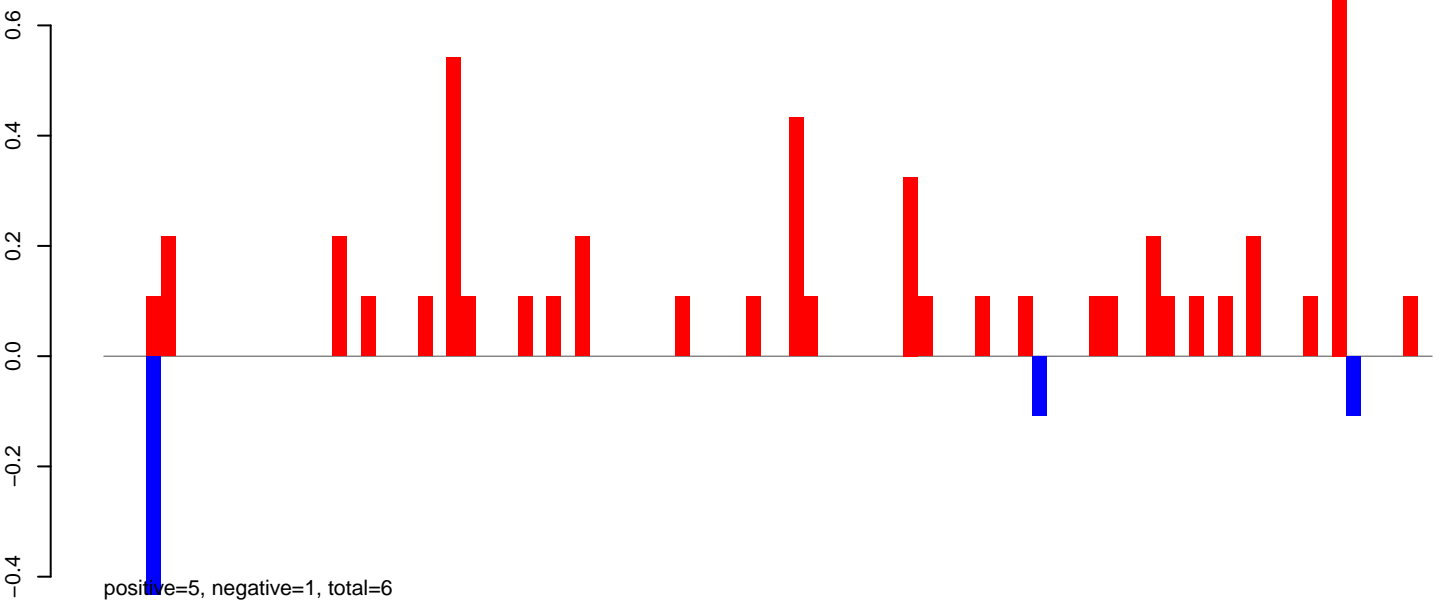
AnGam_Sua5b_cells.24_35.rep



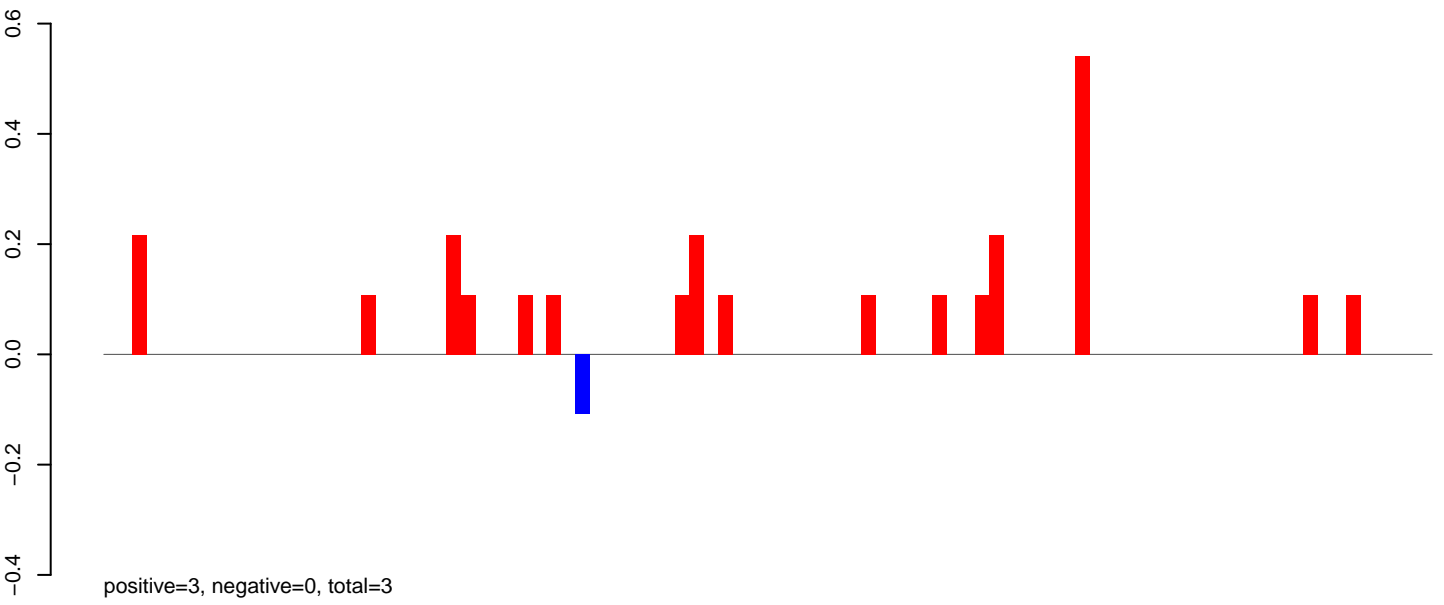
AnGam_Sua5b_cells.rep



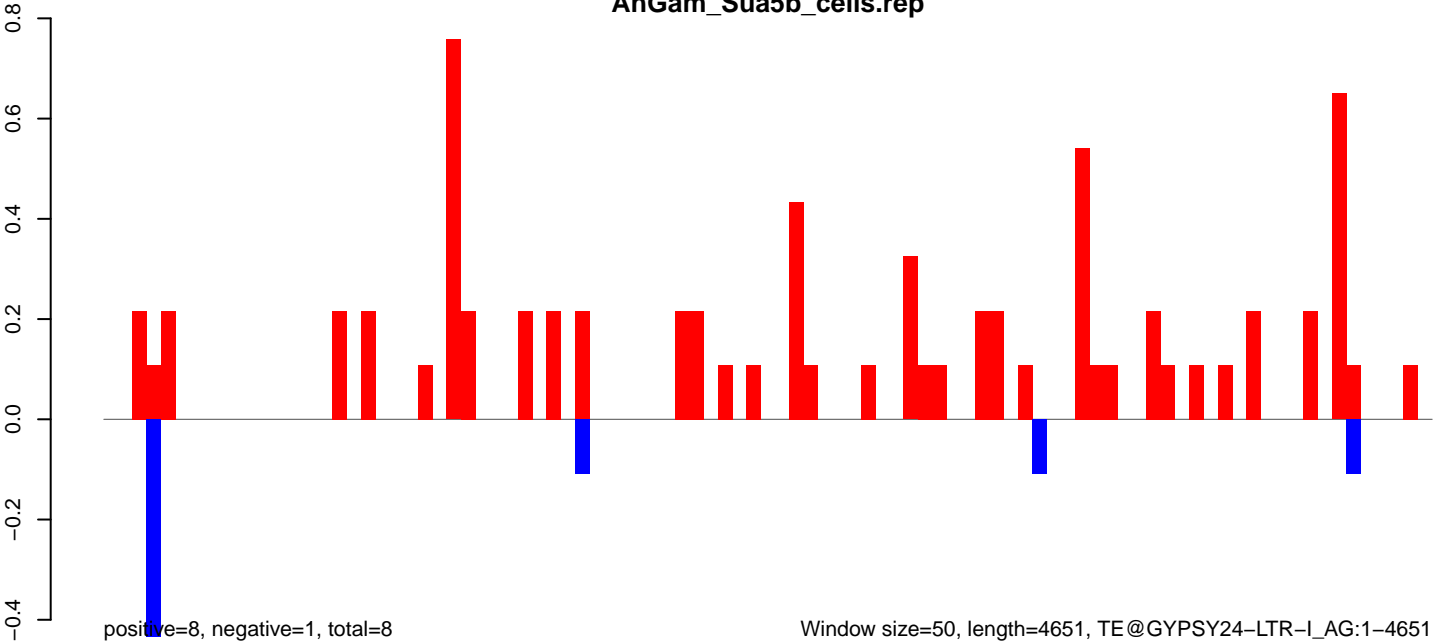
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

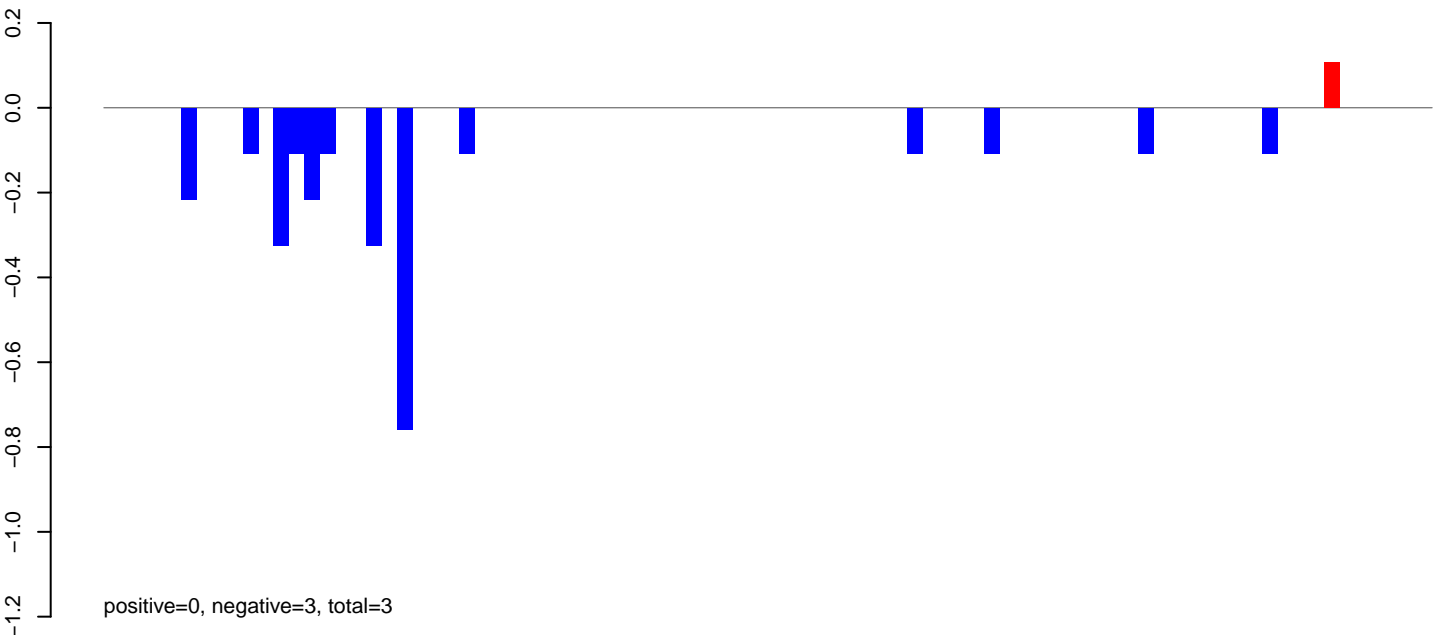


AnGam_Sua5b_cells.rep

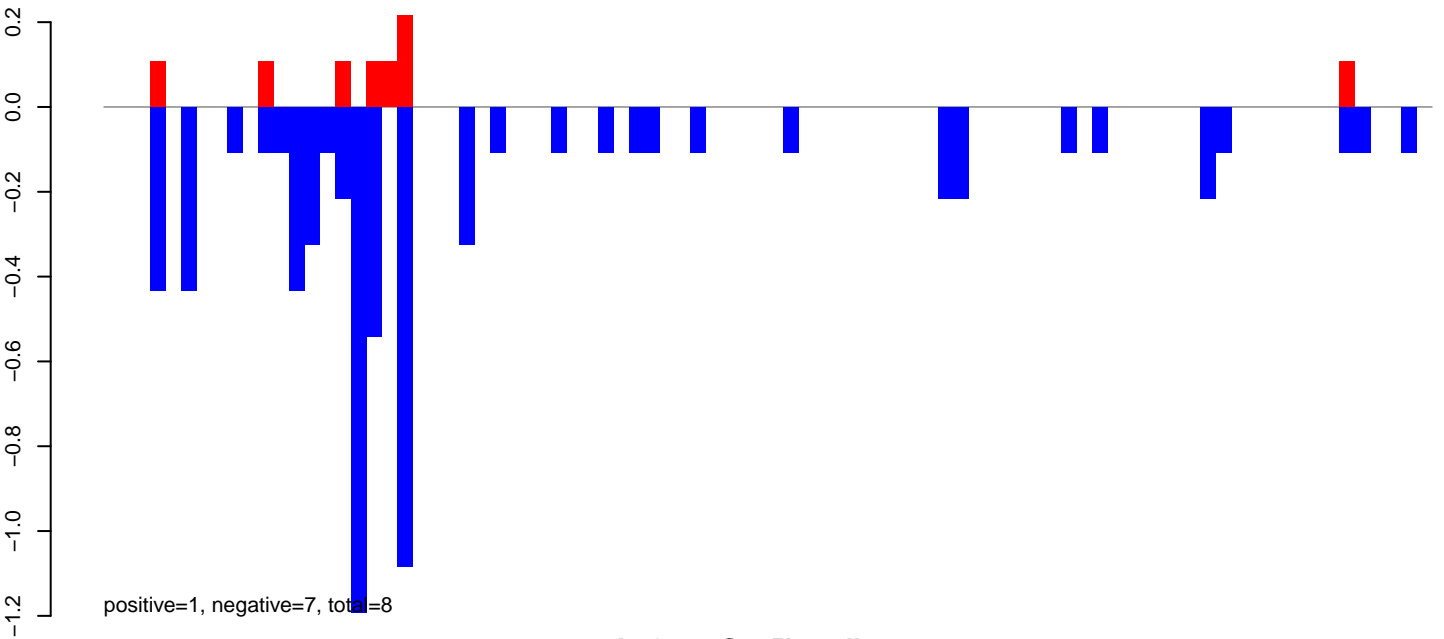


0 1000 2000 3000 4000

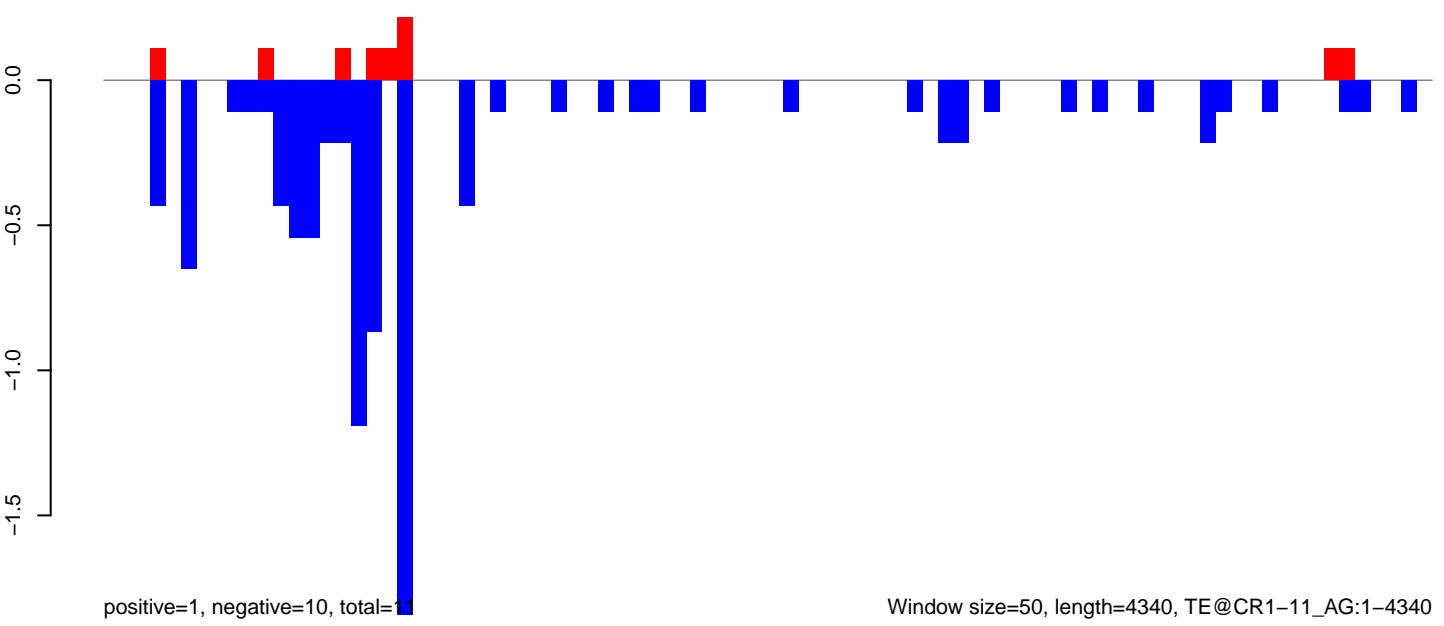
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

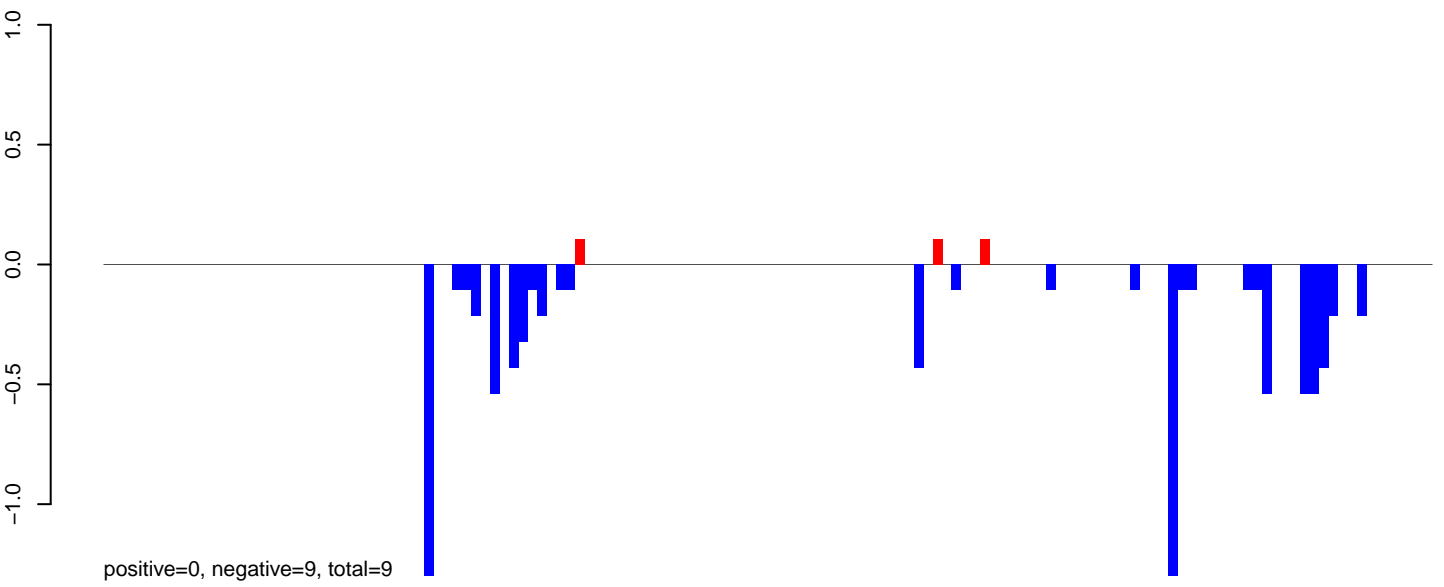


AnGam_Sua5b_cells.rep

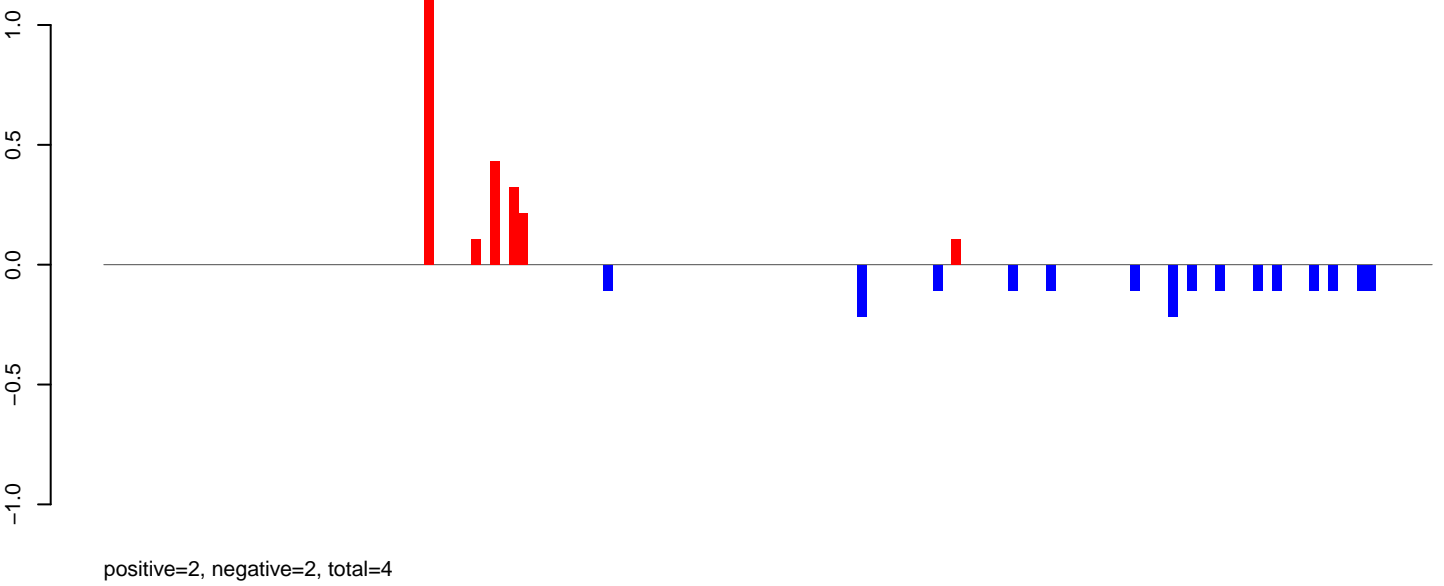


Window size=50, length=4340, TE@CR1-11_AG:1-4340

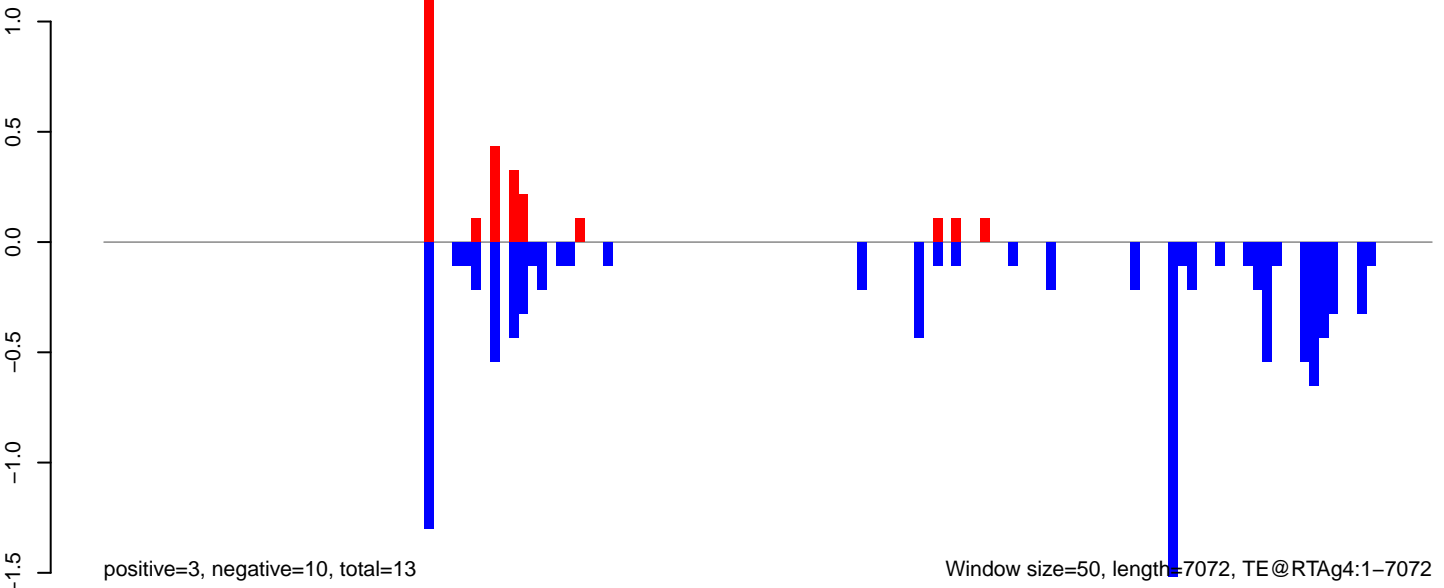
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



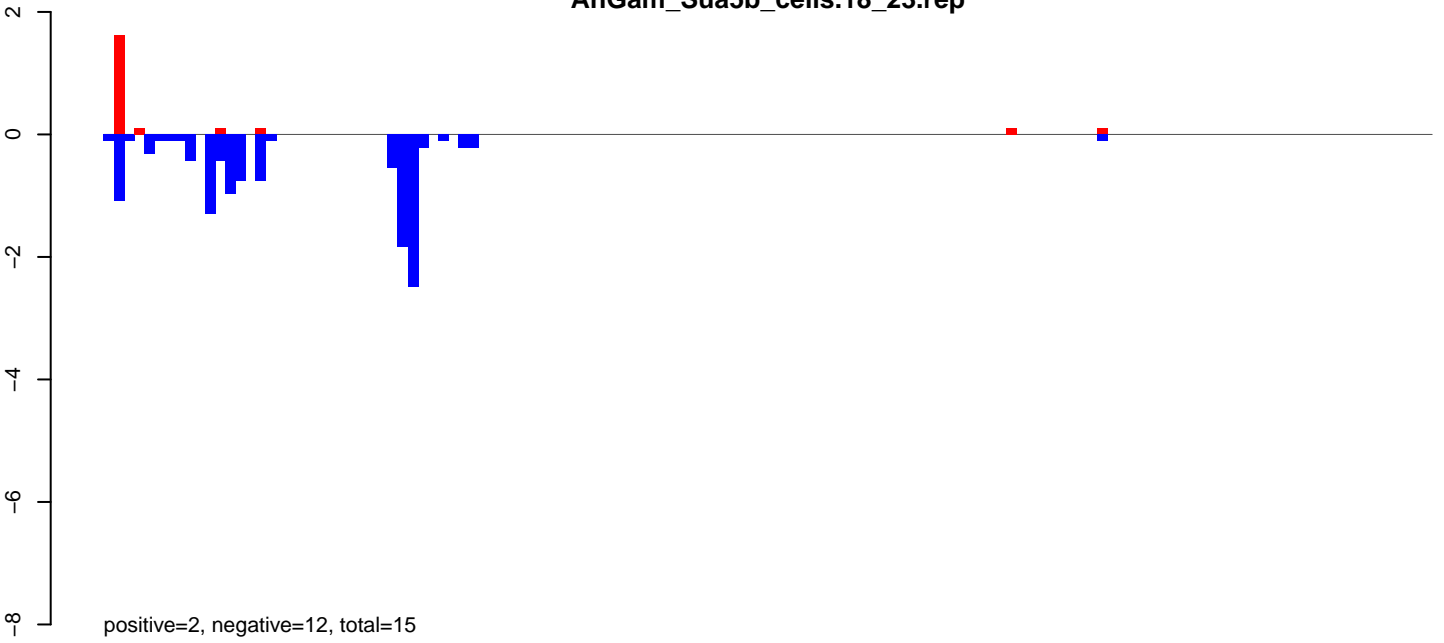
AnGam_Sua5b_cells.rep



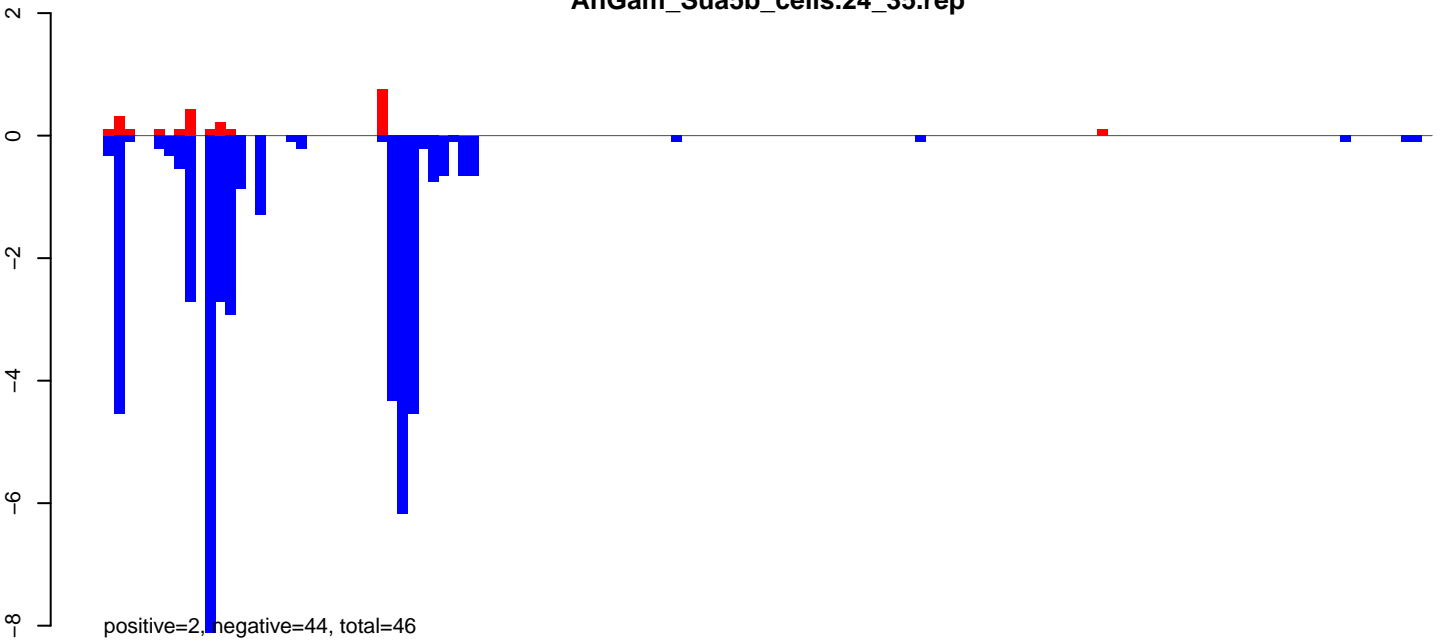
Window size=50, length=7072, TE@RTAg4:1-7072

0 1000 2000 3000 4000 5000 6000 7000

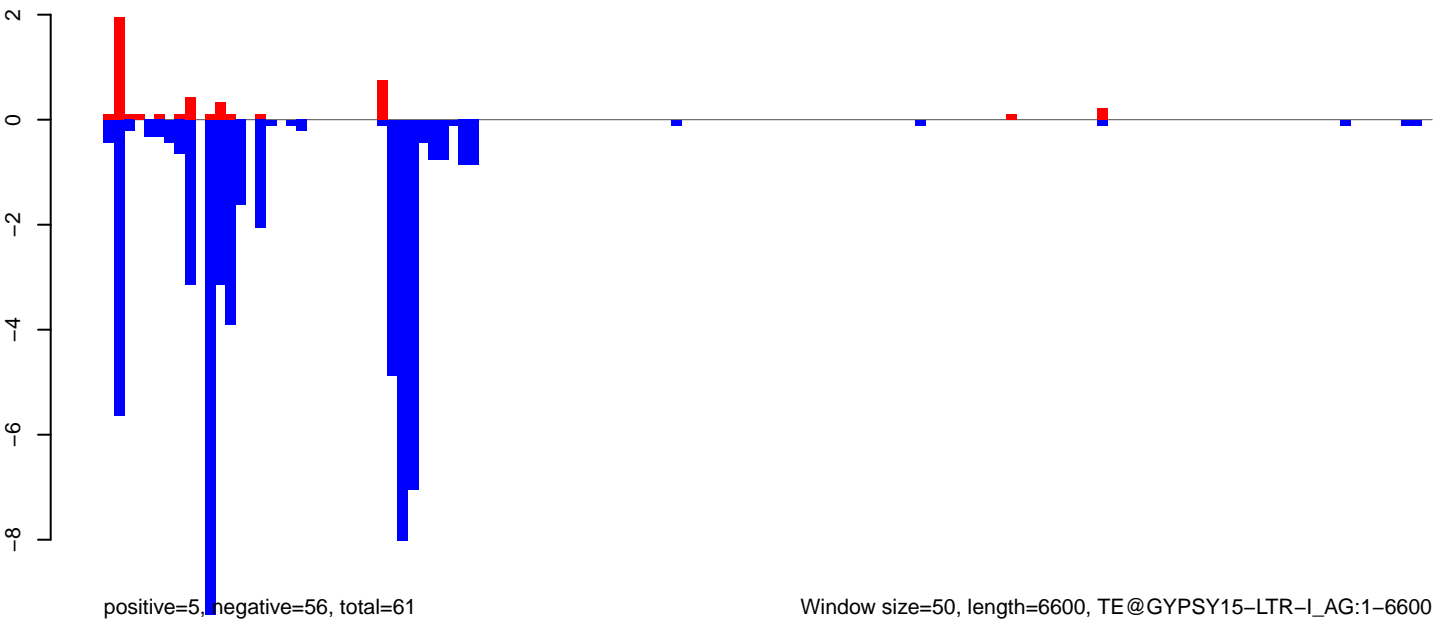
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

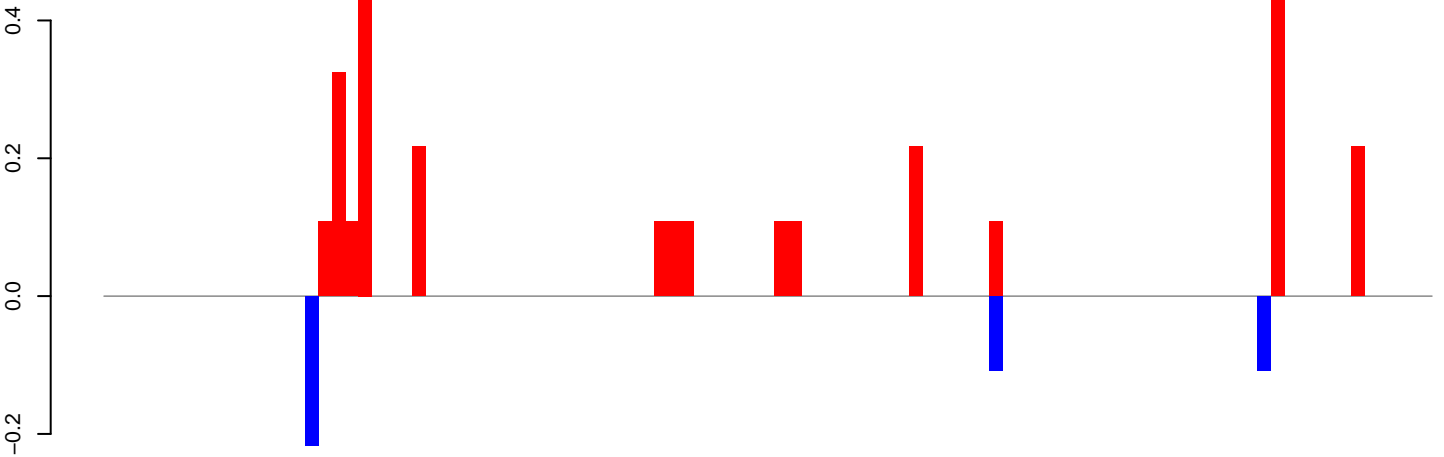


AnGam_Sua5b_cells.rep



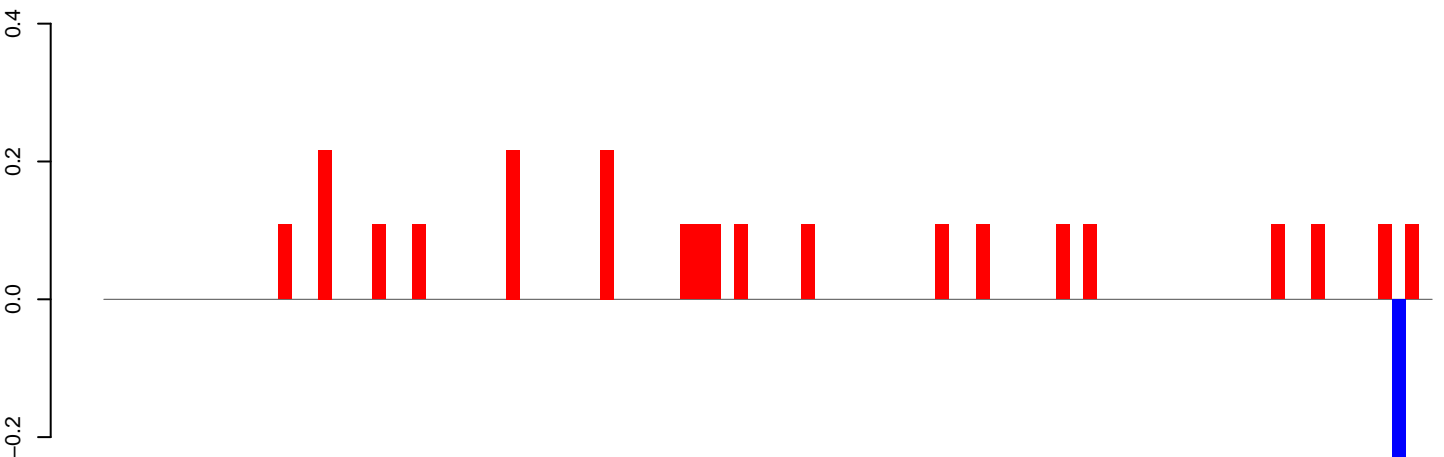
0 1000 2000 3000 4000 5000 6000

AnGam_Sua5b_cells.18_23.rep



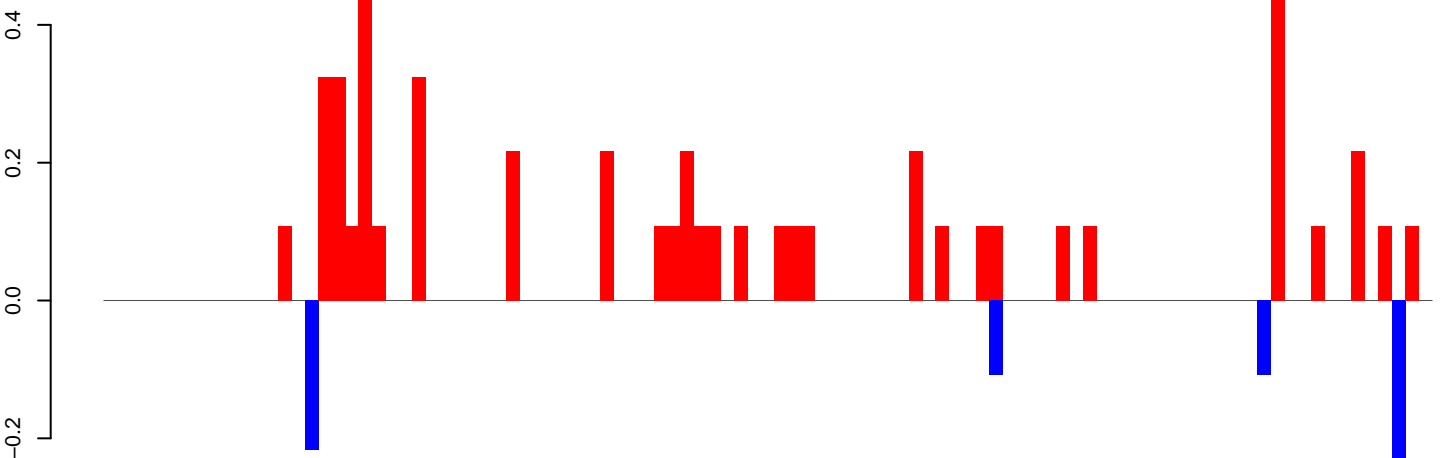
positive=3, negative=0, total=3

AnGam_Sua5b_cells.24_35.rep



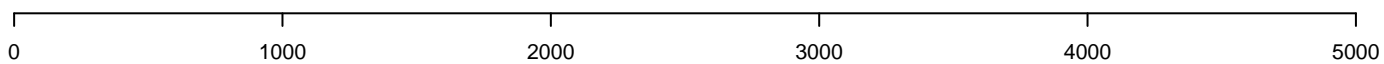
positive=2, negative=0, total=3

AnGam_Sua5b_cells.rep

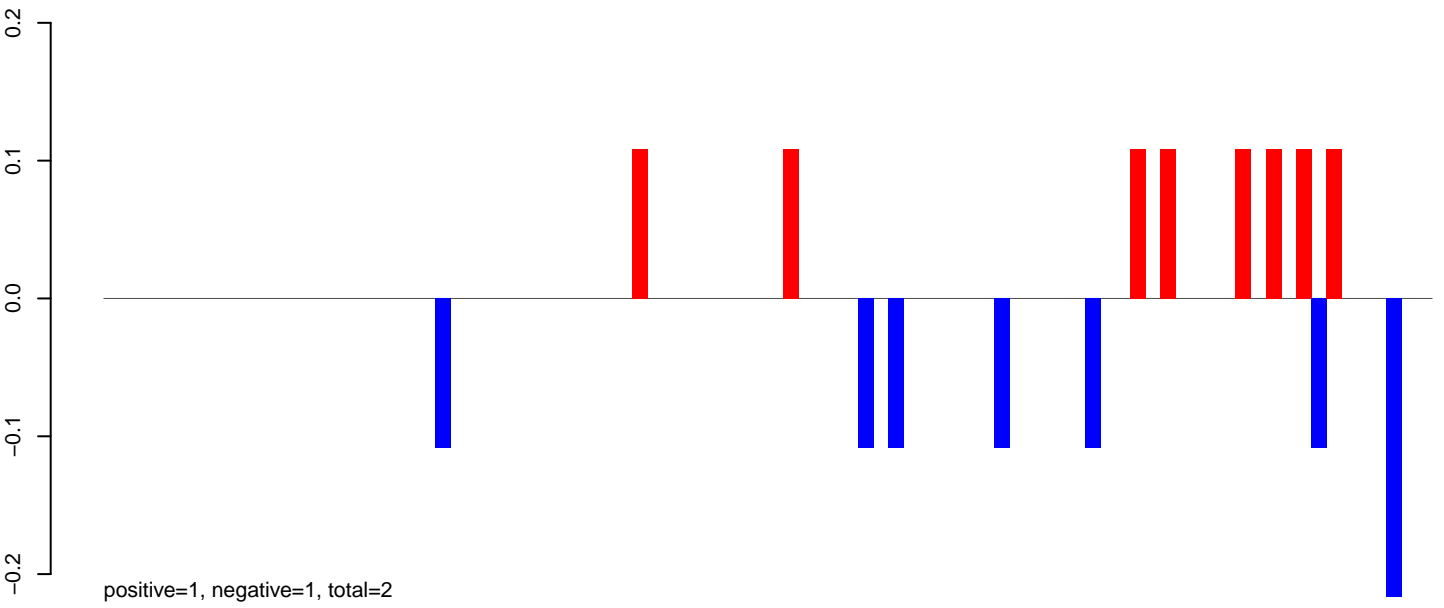


positive=5, negative=1, total=6

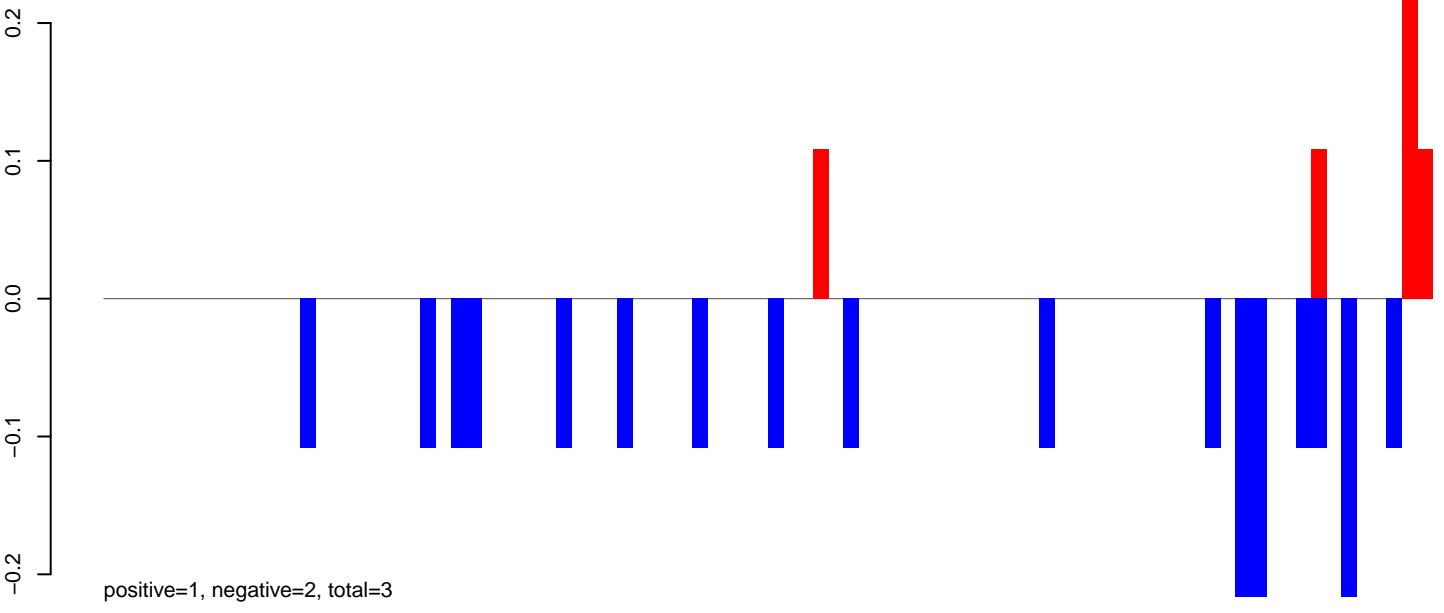
Window size=50, length=4991, TE@Agam_AB090818_R6Ag2-LINE-R6Ag2:1-4991



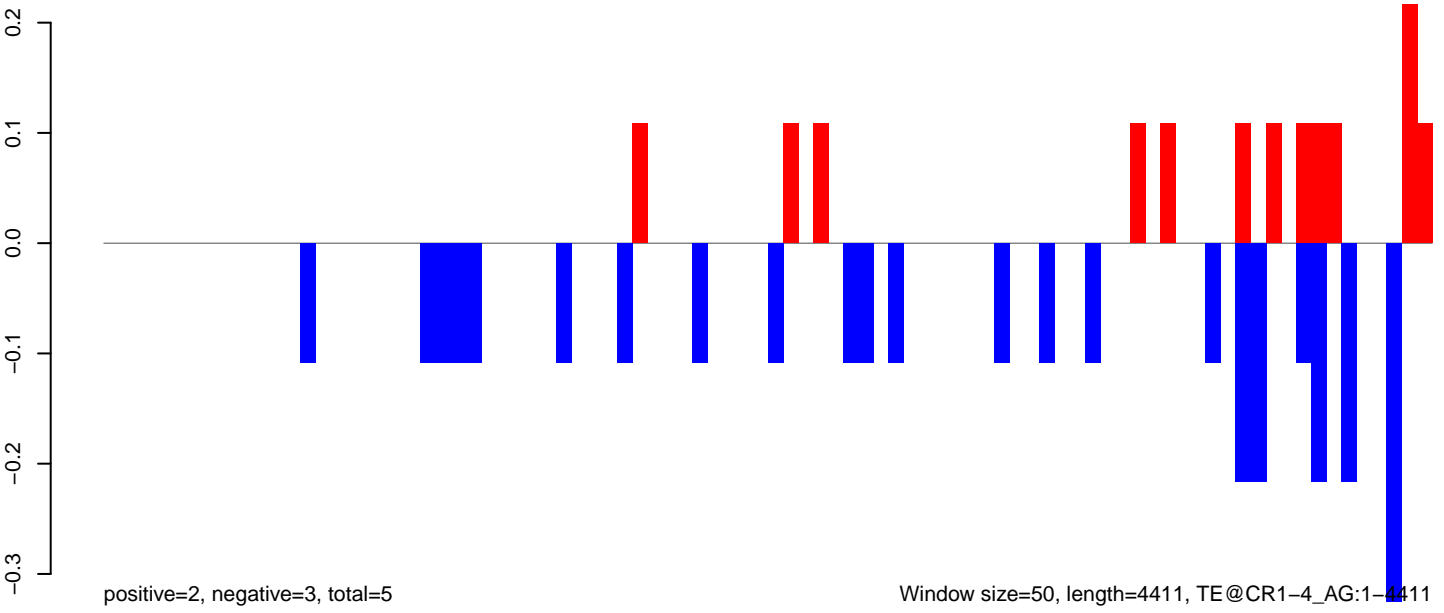
AnGam_Sua5b_cells.18_23.rep



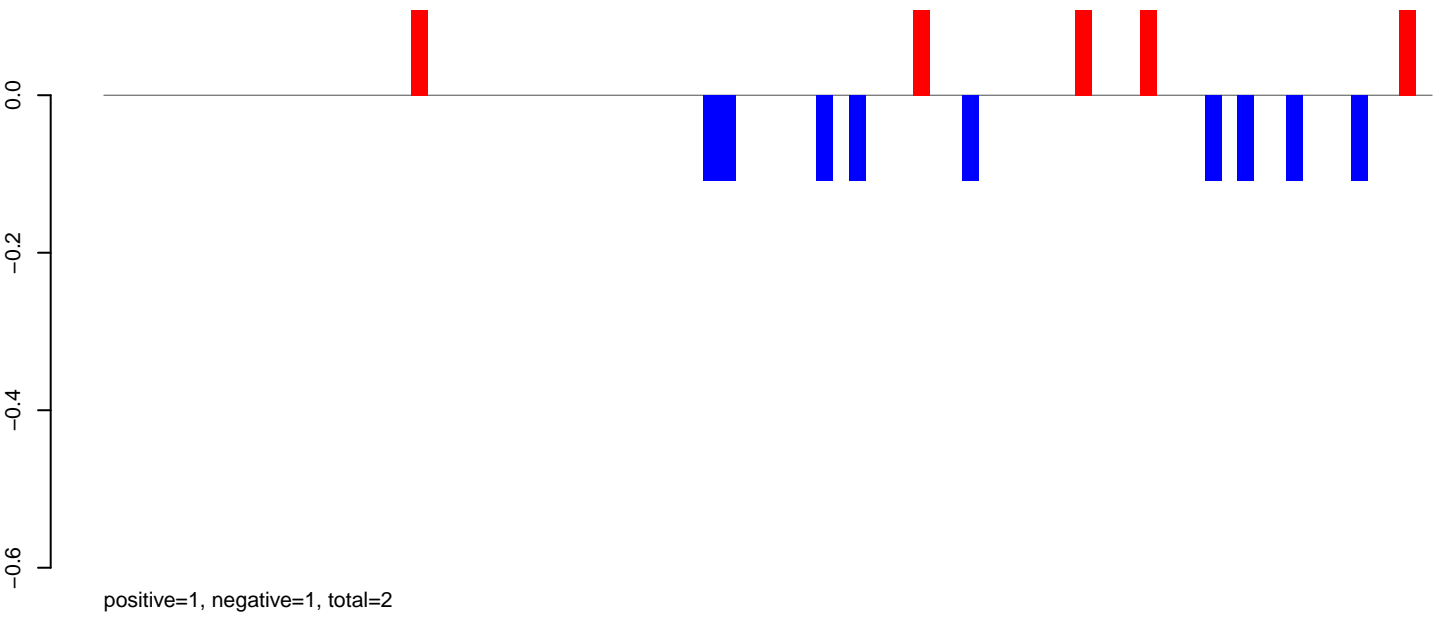
AnGam_Sua5b_cells.24_35.rep



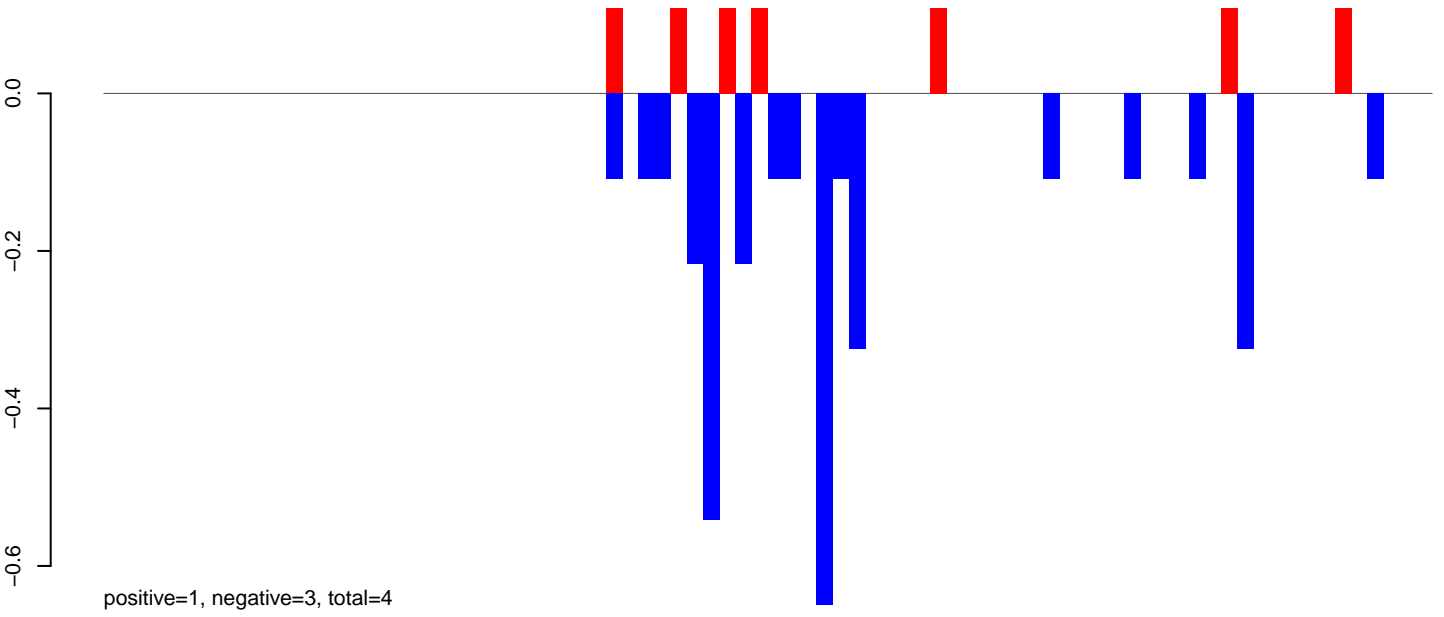
AnGam_Sua5b_cells.rep



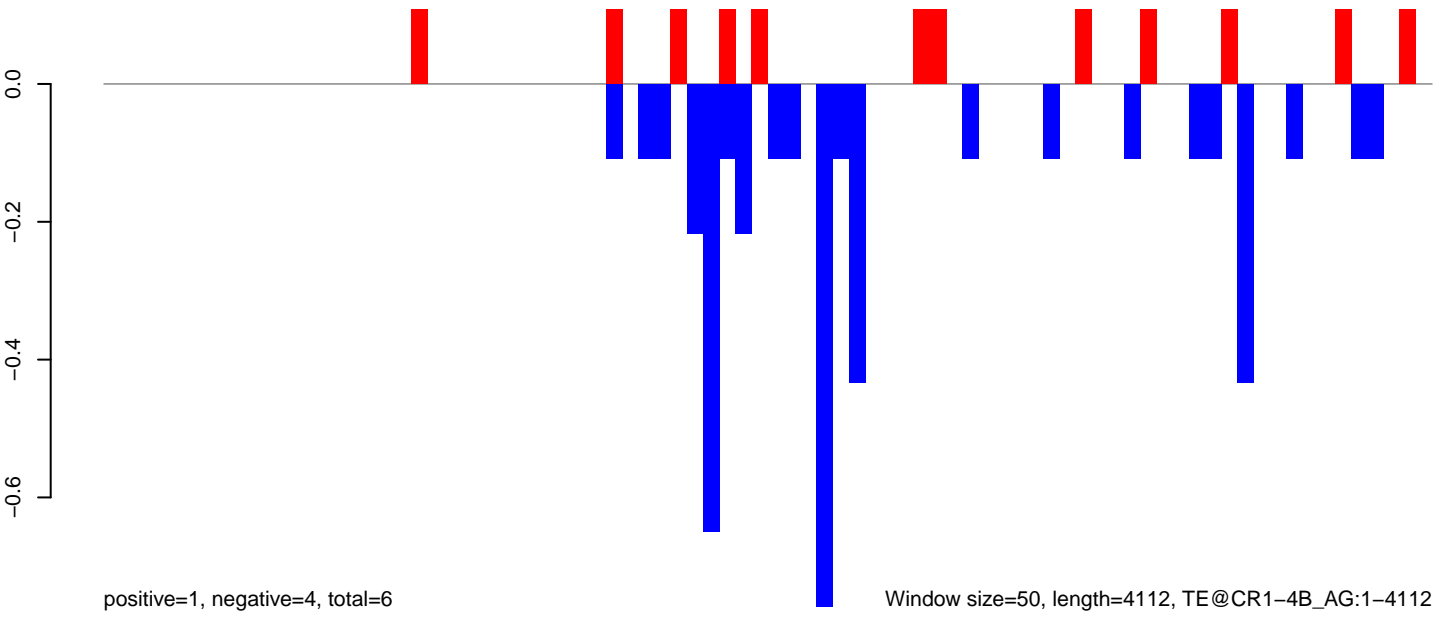
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

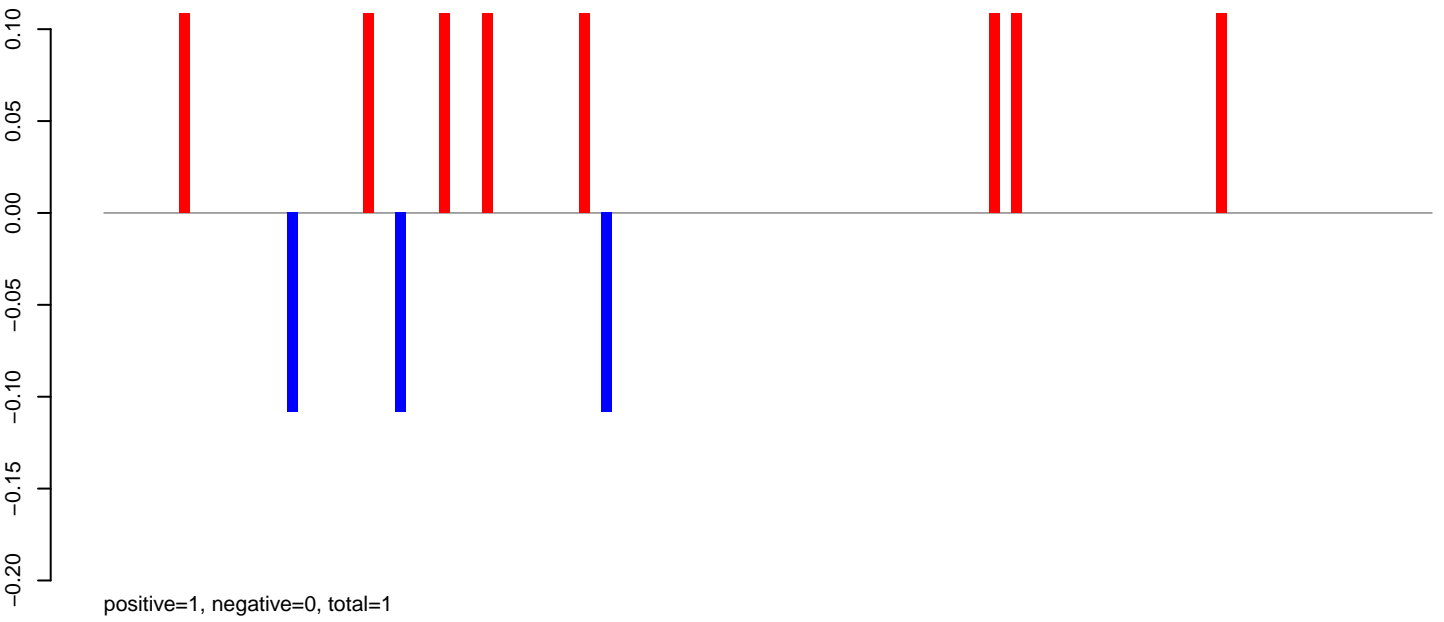


AnGam_Sua5b_cells.rep

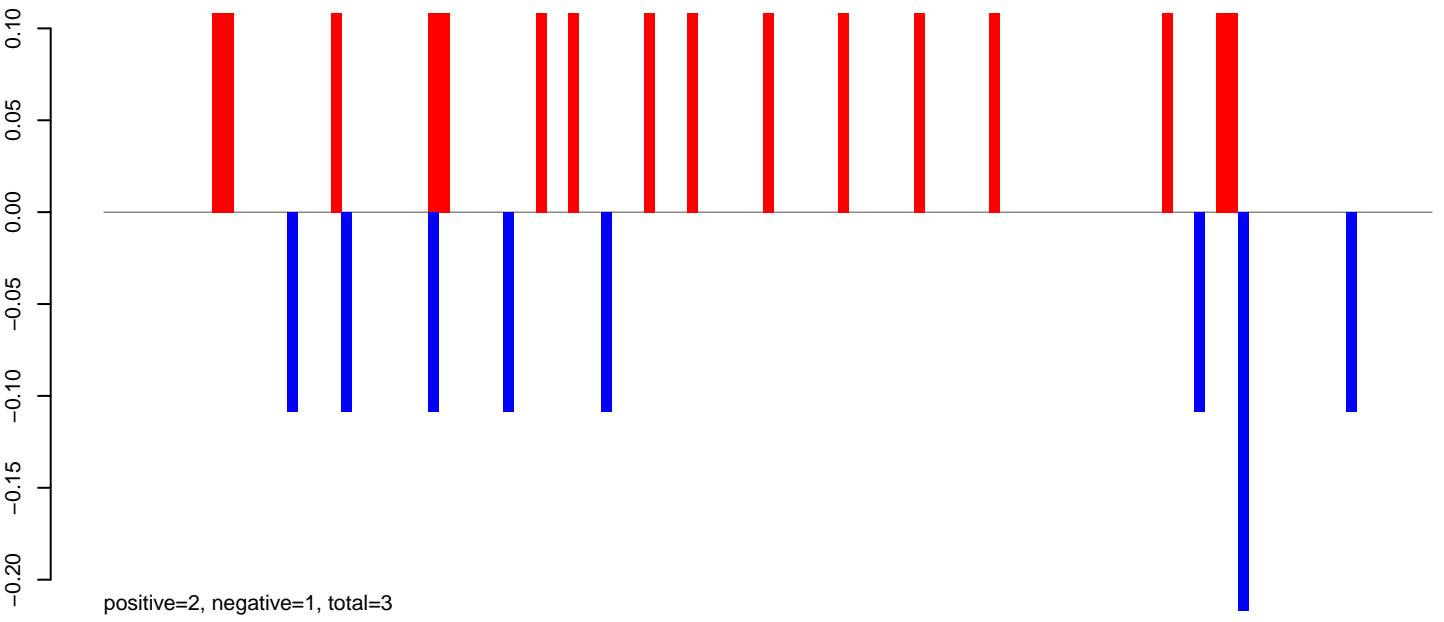


Window size=50, length=4112, TE@CR1-4B_AG:1-4112

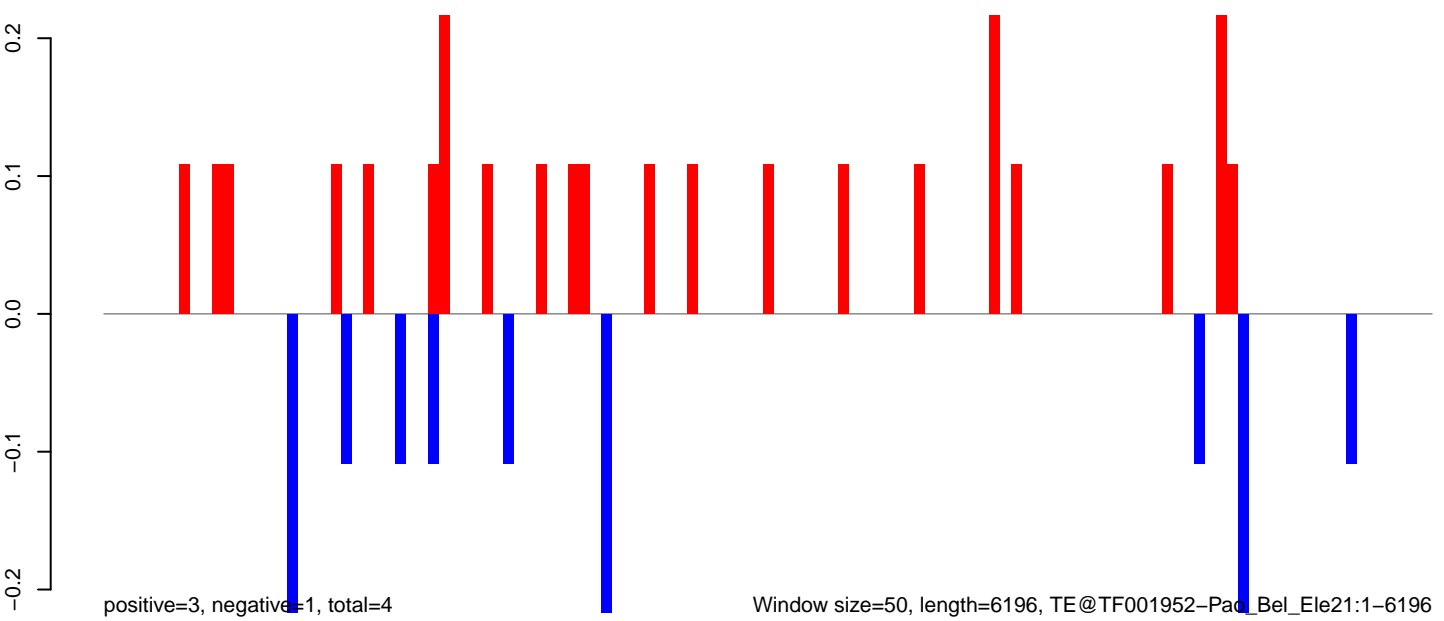
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

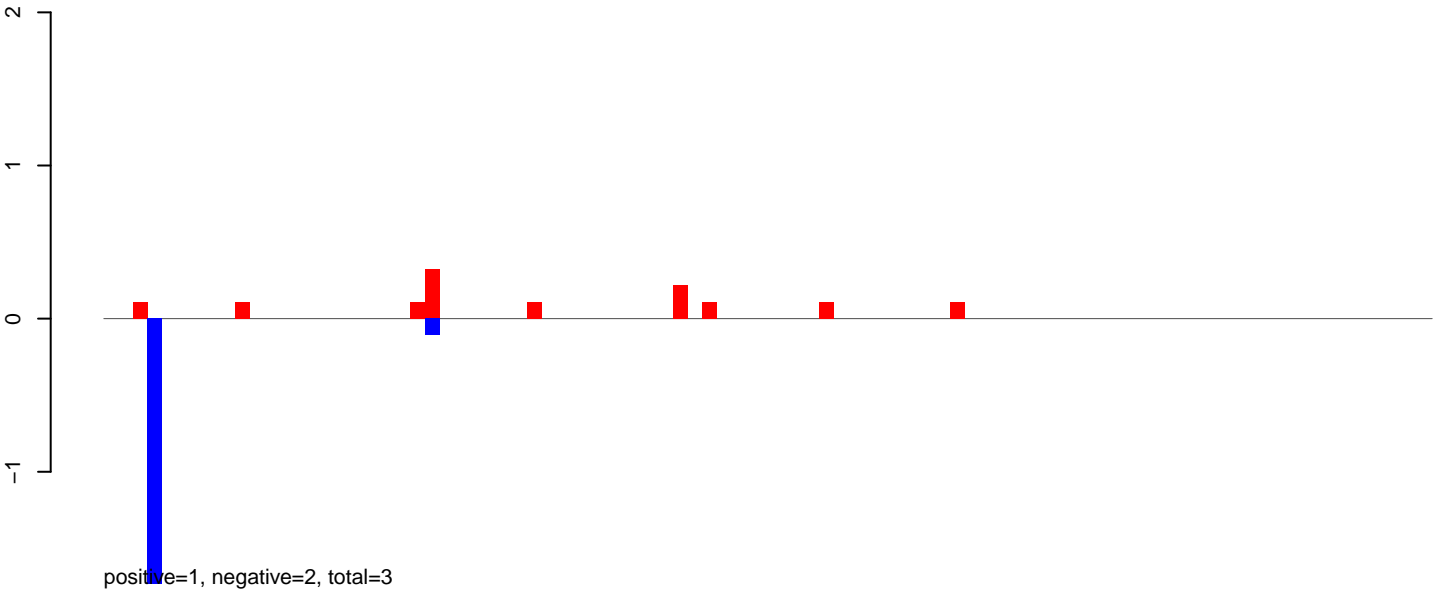


AnGam_Sua5b_cells.rep

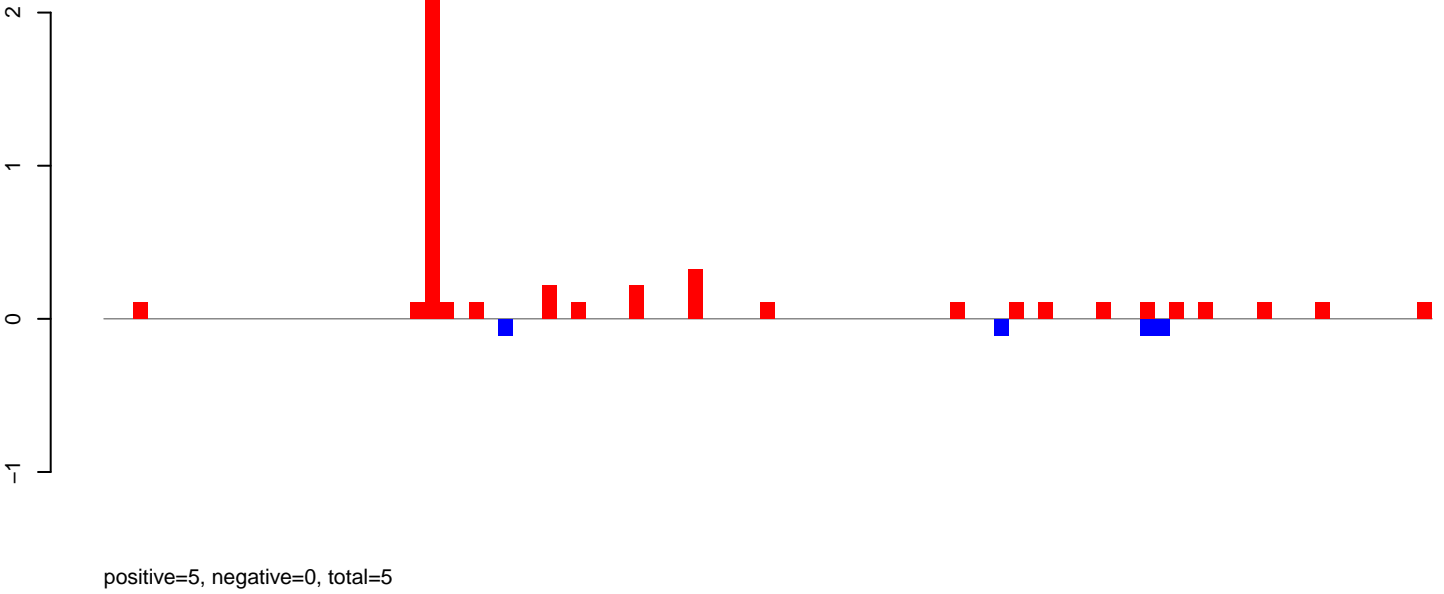


Window size=50, length=6196, TE@TF001952-Pac_Bel_Ele21:1-6196

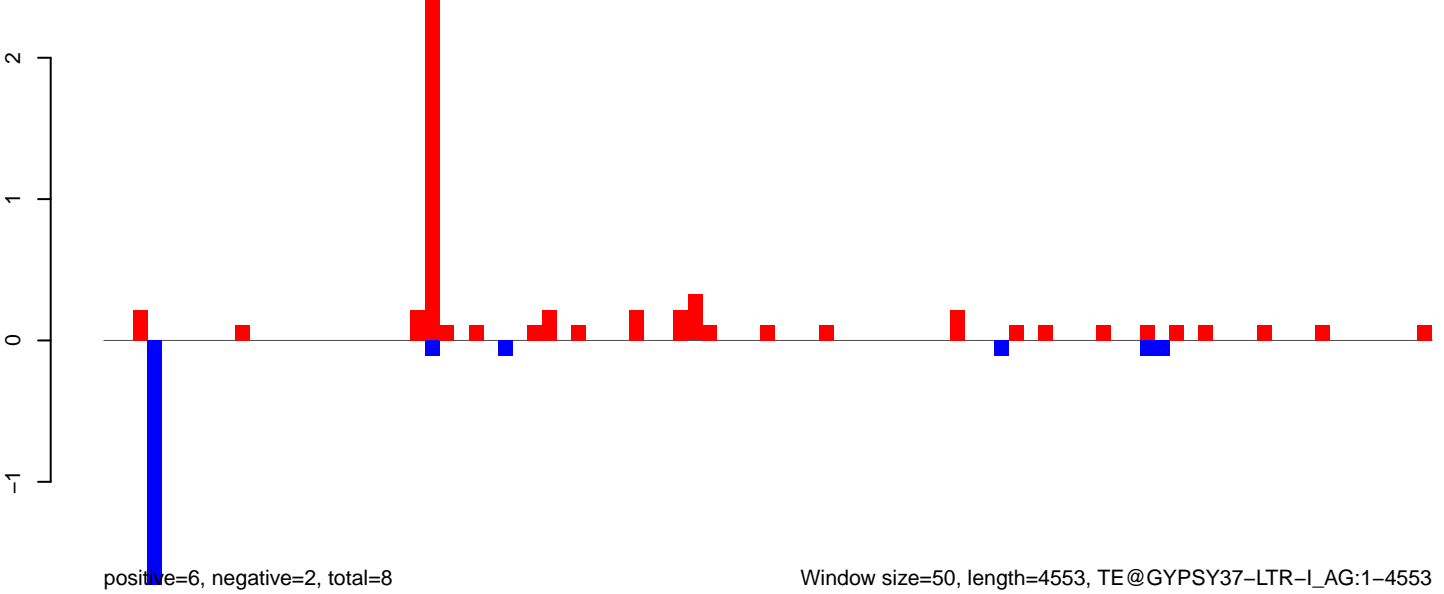
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

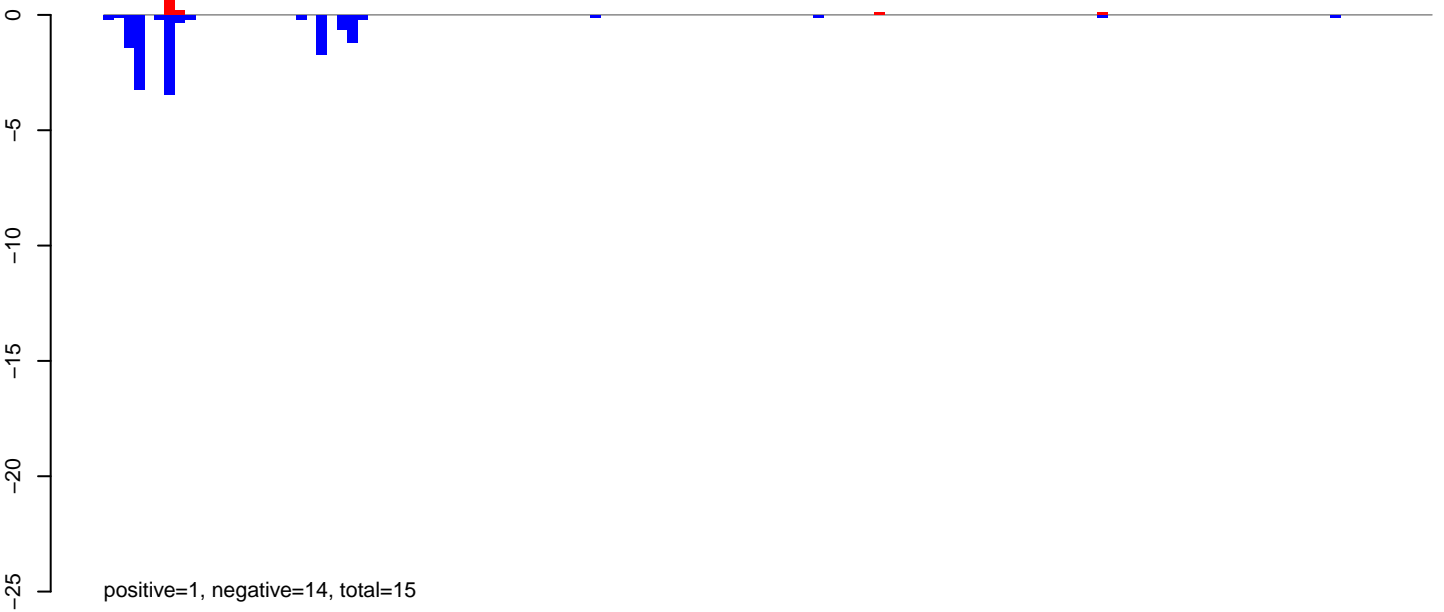


AnGam_Sua5b_cells.rep



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

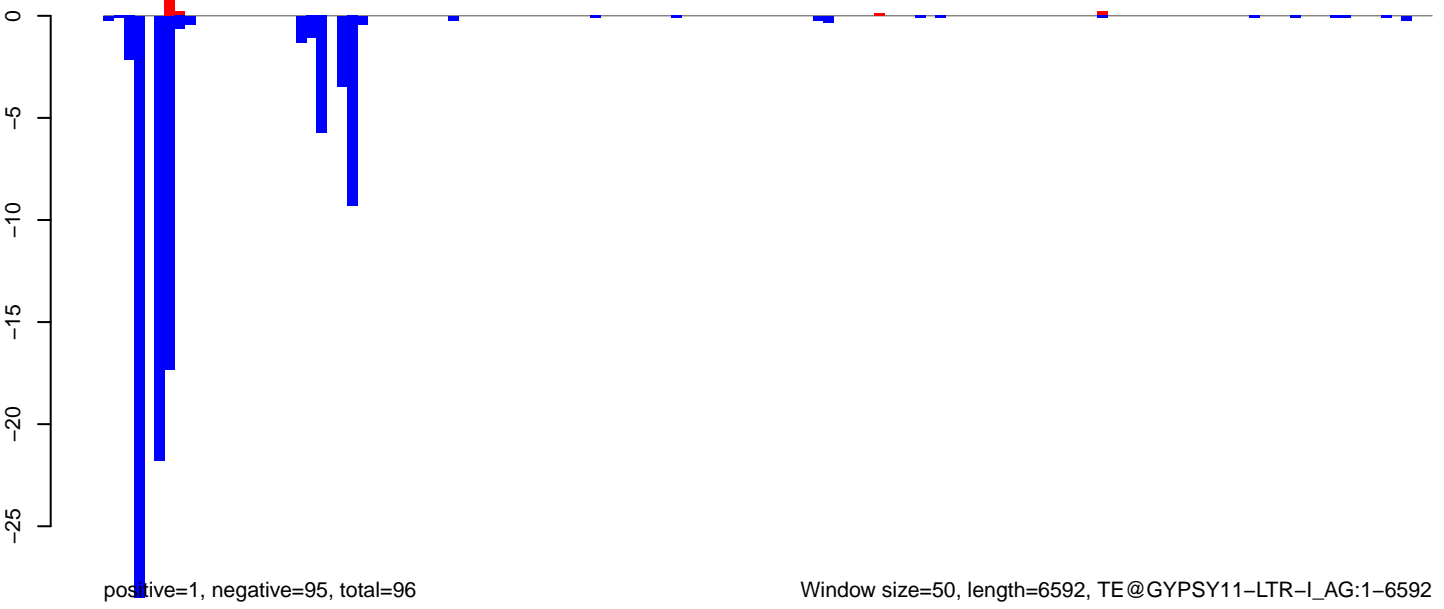
AnGam_Sua5b_cells.18_23.rep



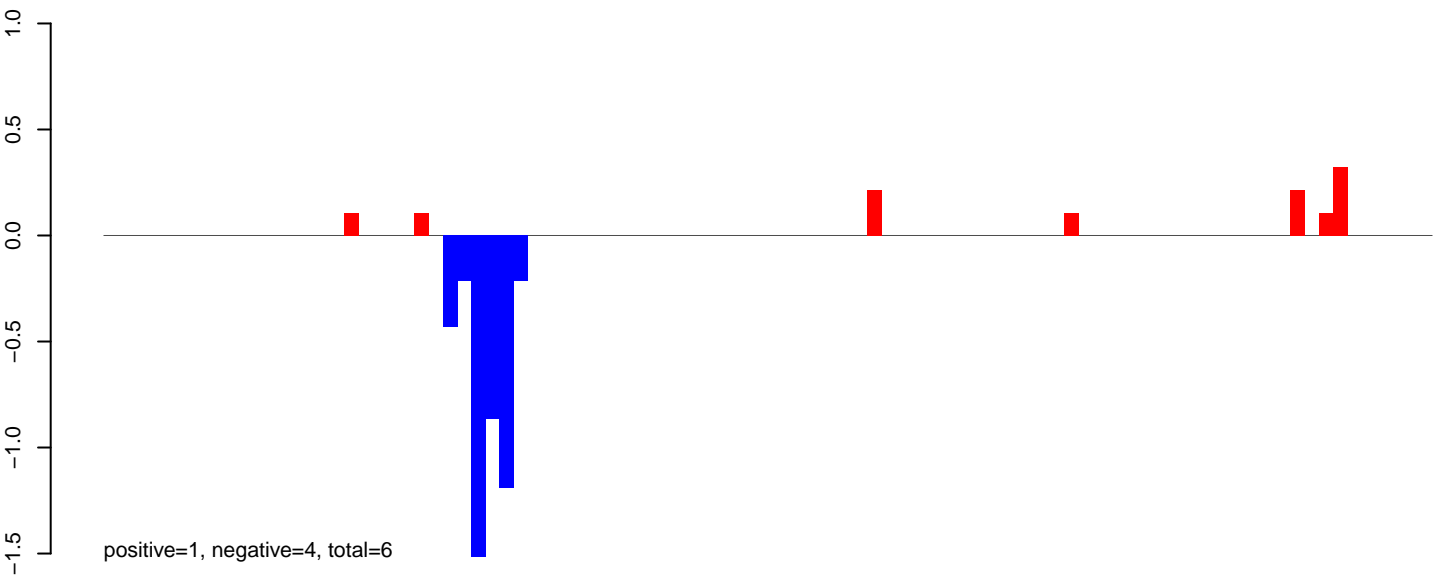
AnGam_Sua5b_cells.24_35.rep



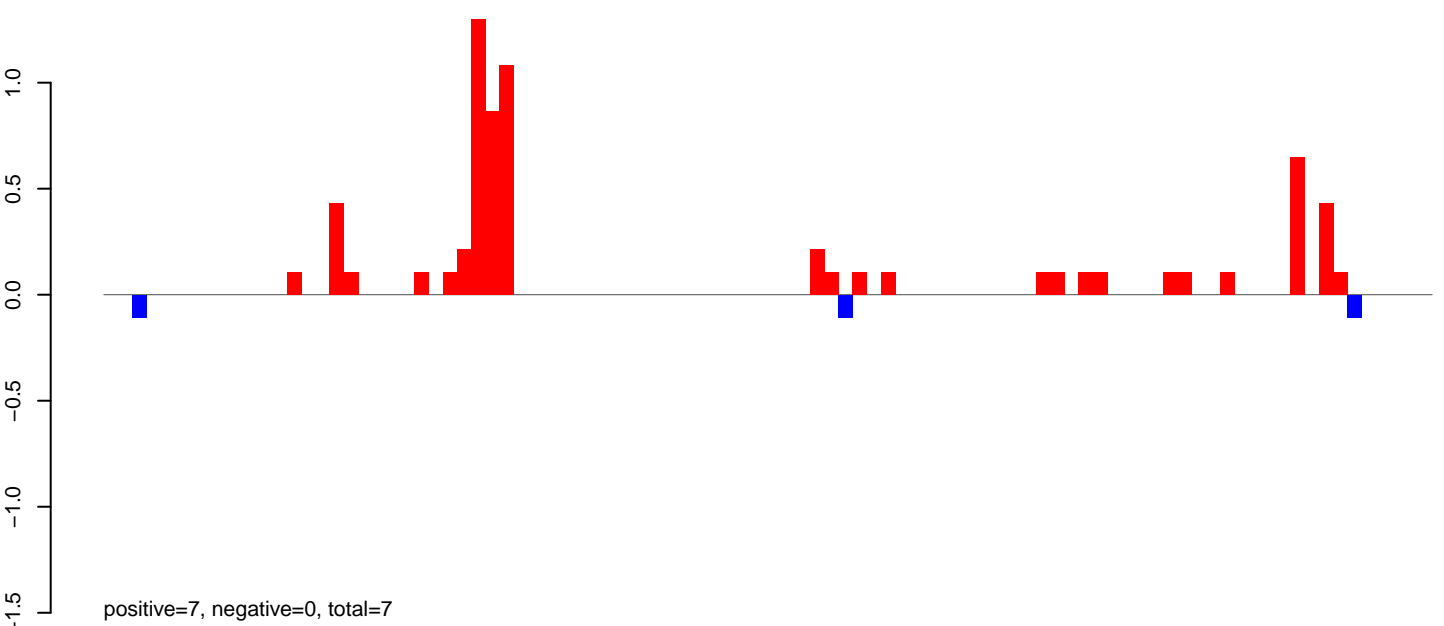
AnGam_Sua5b_cells.rep



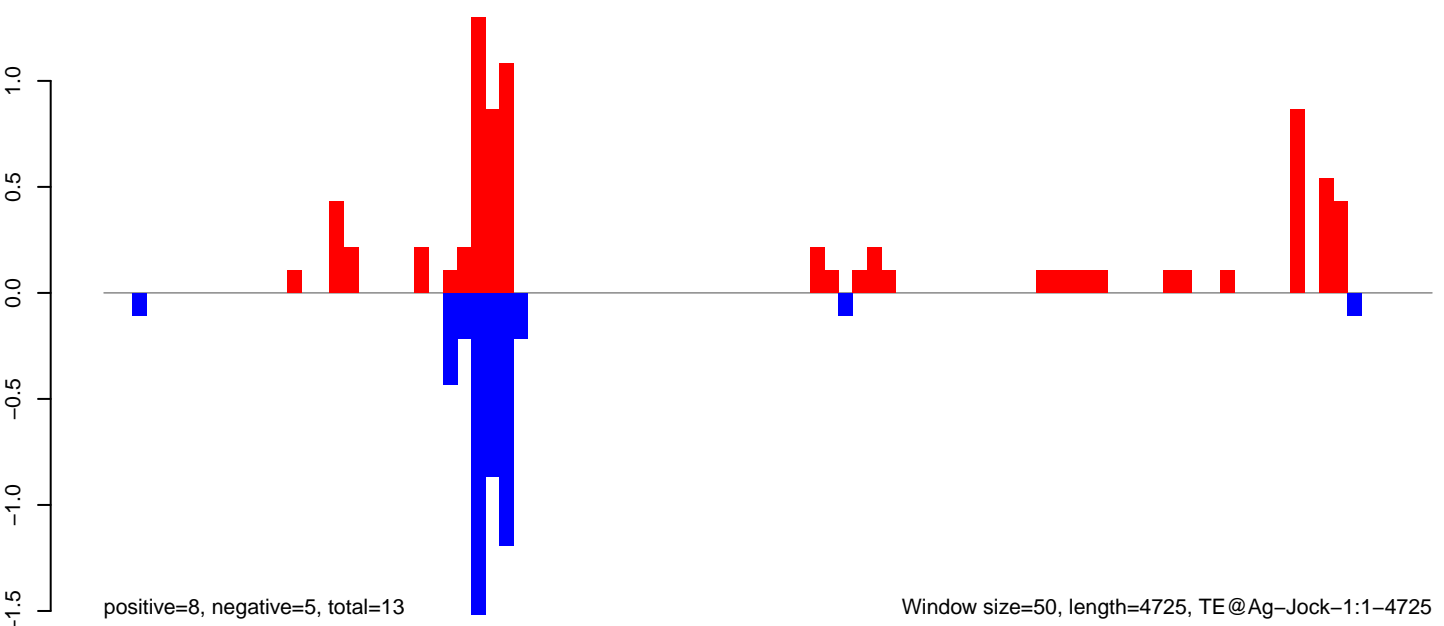
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



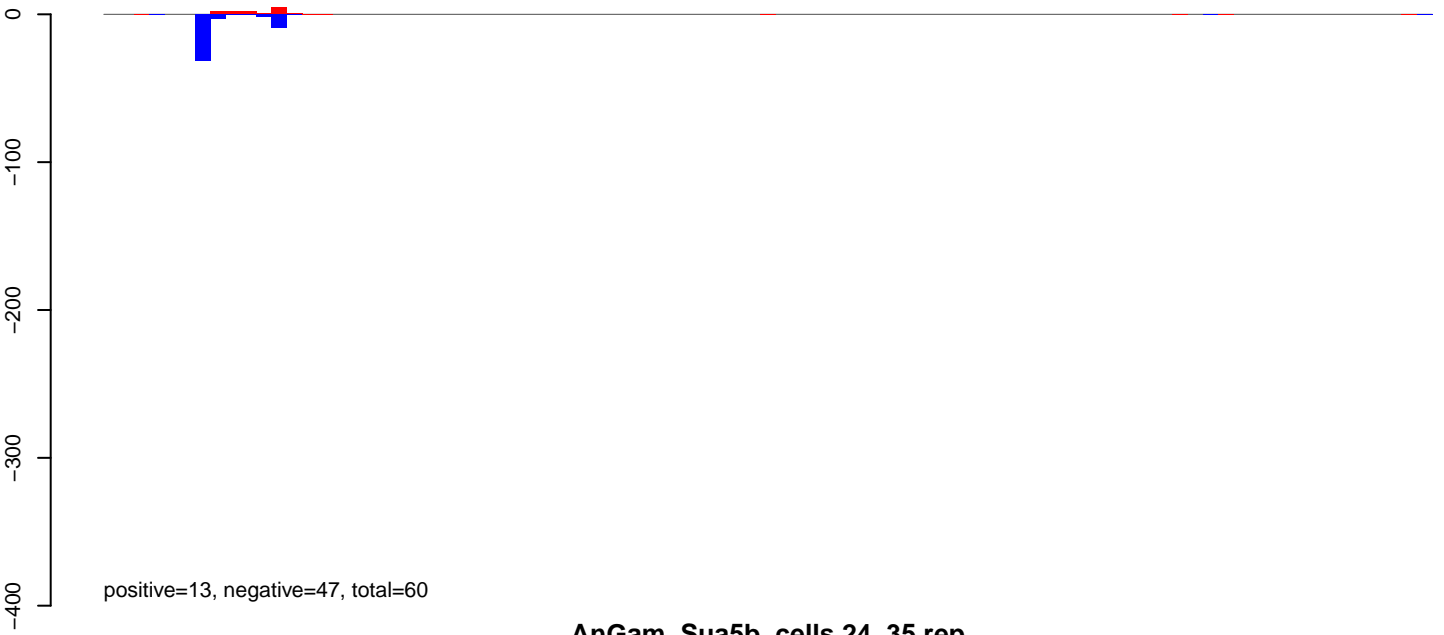
AnGam_Sua5b_cells.rep



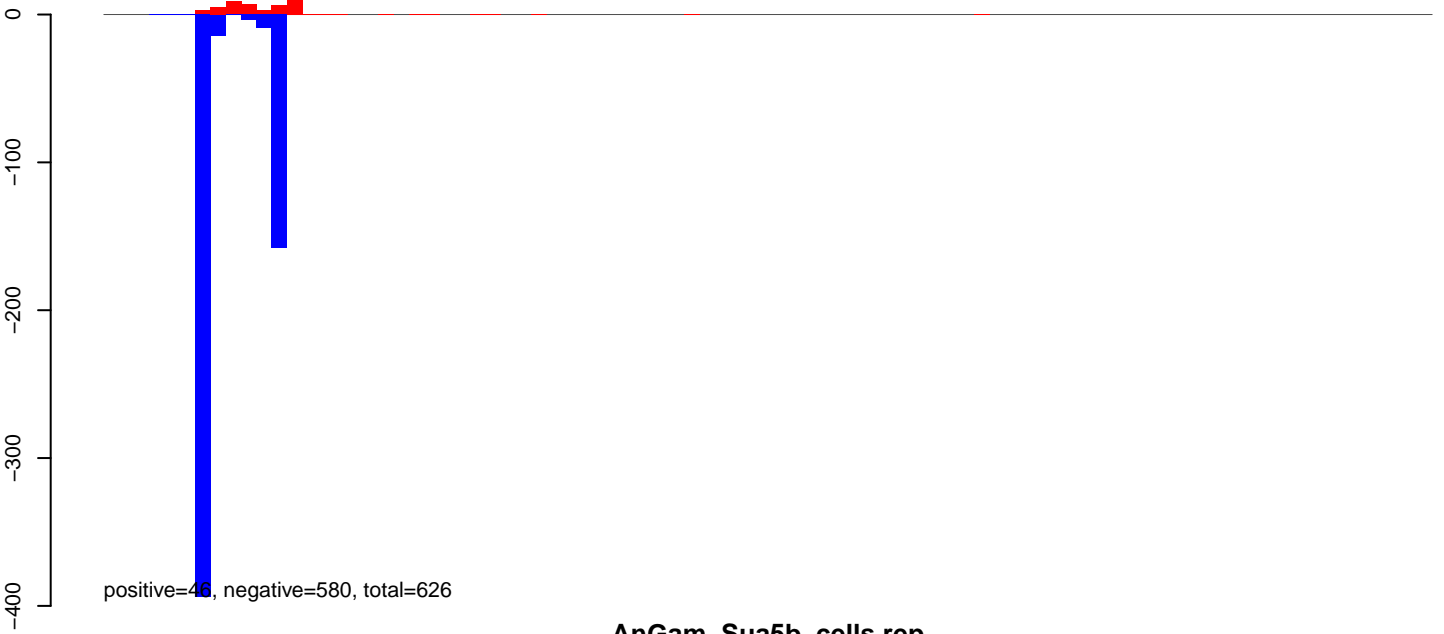
Window size=50, length=4725, TE@Ag-Jock-1:1-4725

0 1000 2000 3000 4000

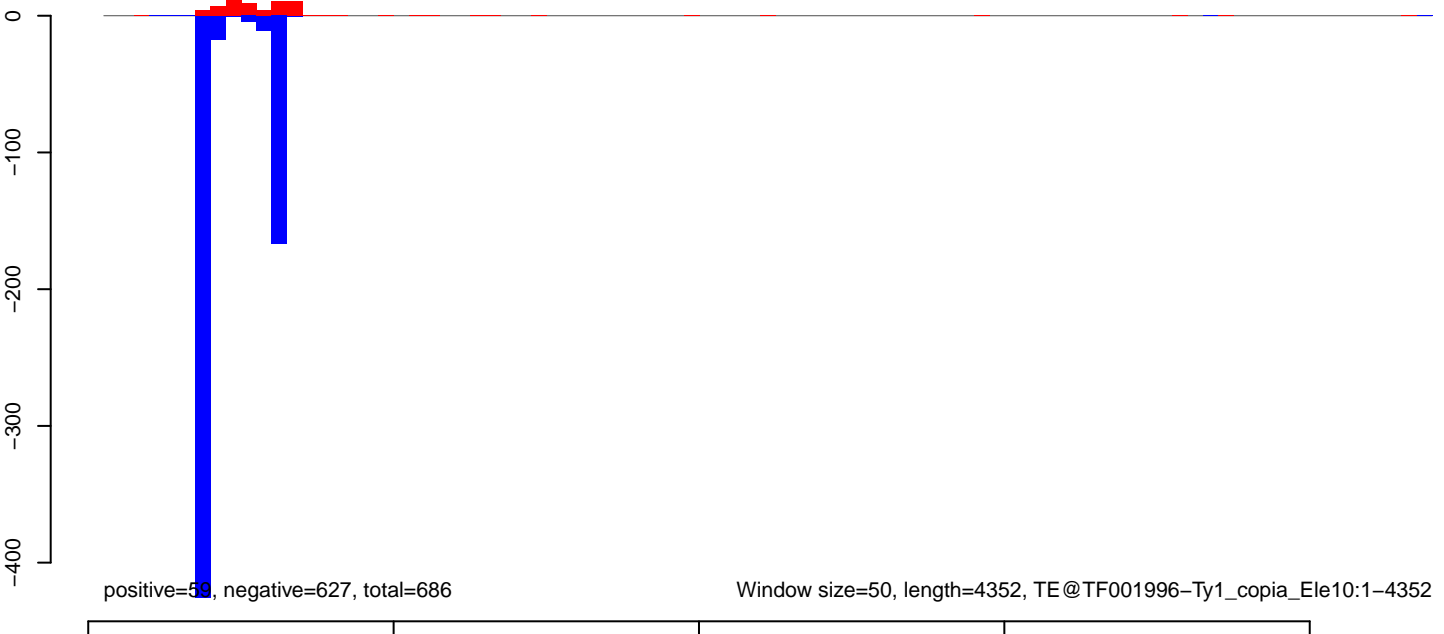
AnGam_Sua5b_cells.18_23.rep



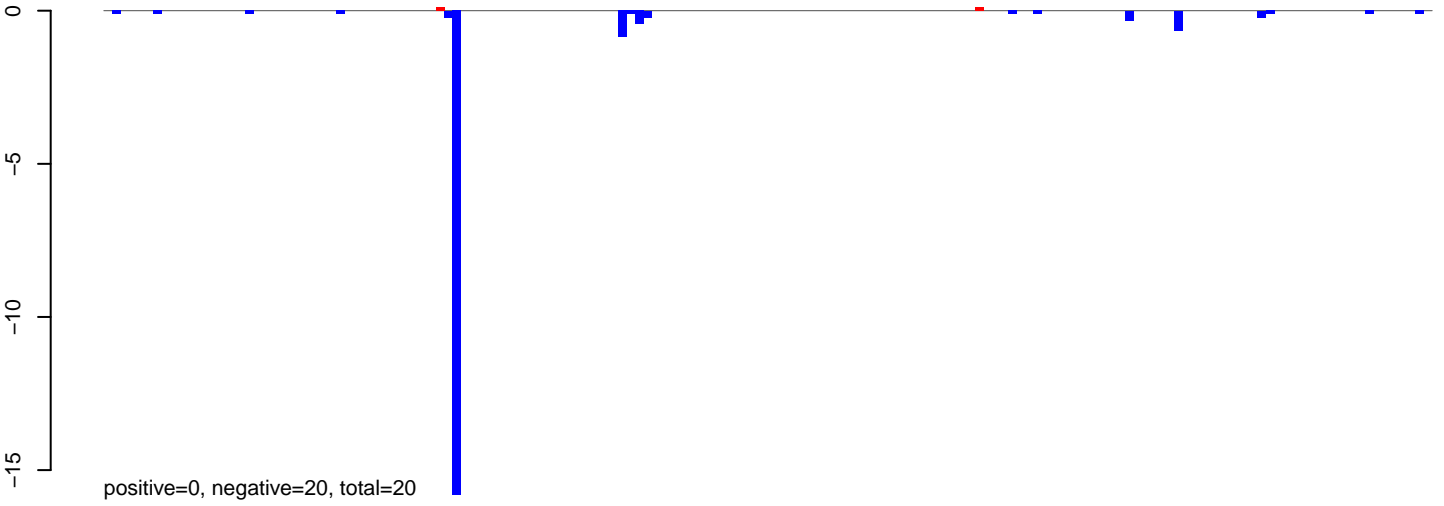
AnGam_Sua5b_cells.24_35.rep



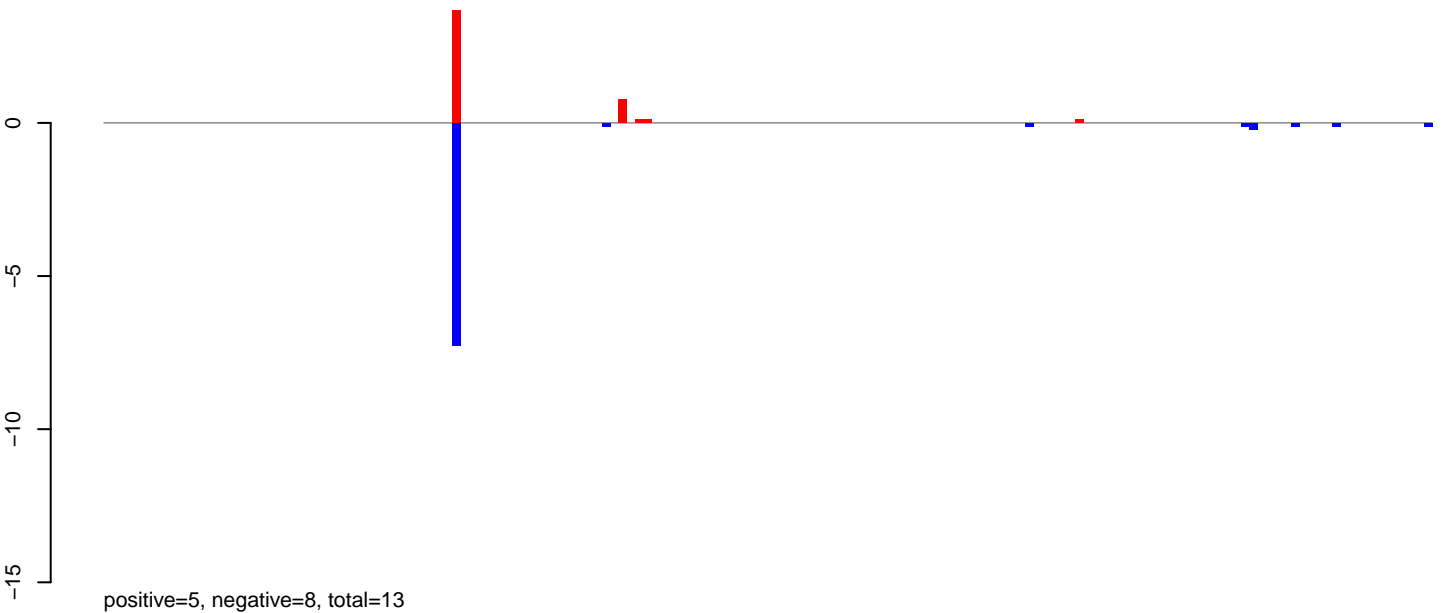
AnGam_Sua5b_cells.rep



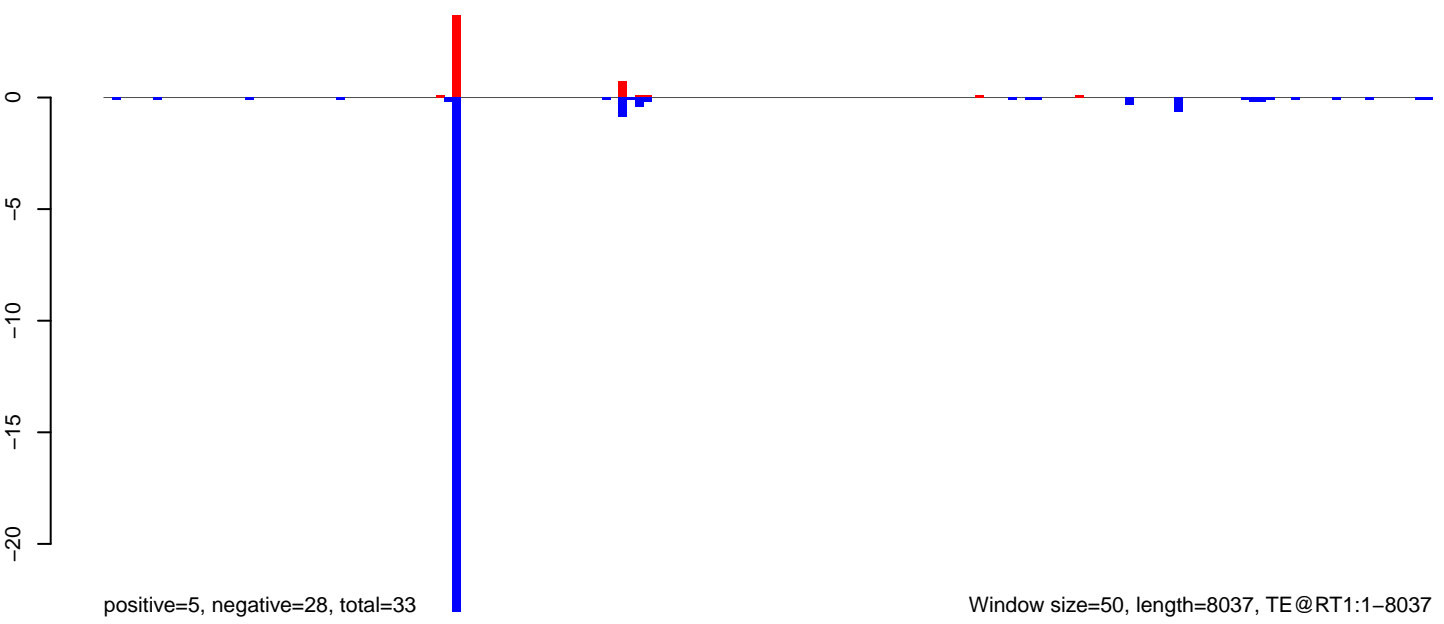
AnGam_Sua5b_cells.18_23.rep



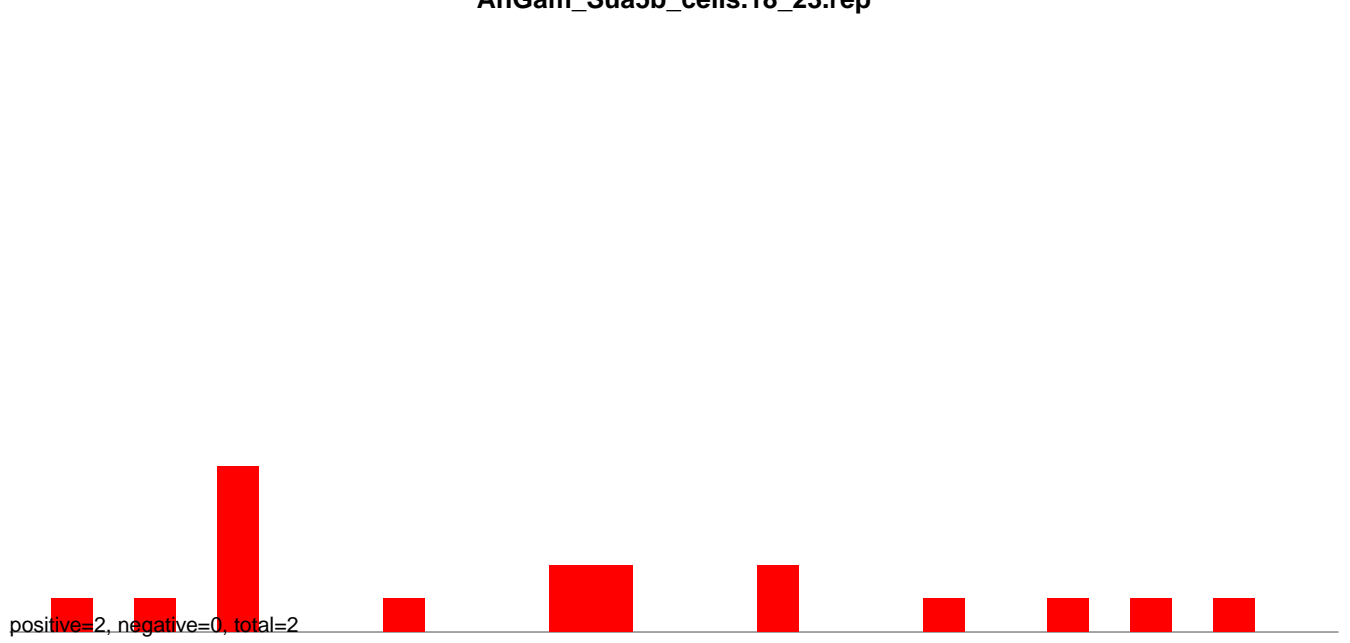
AnGam_Sua5b_cells.24_35.rep



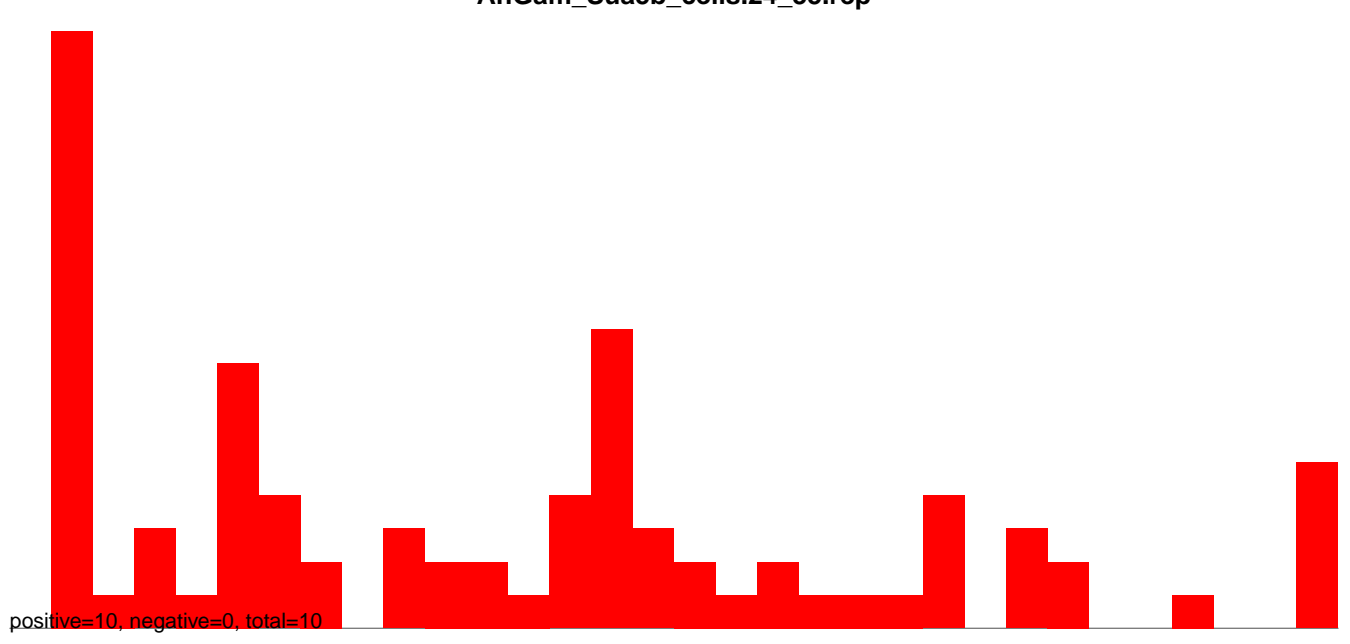
AnGam_Sua5b_cells.rep



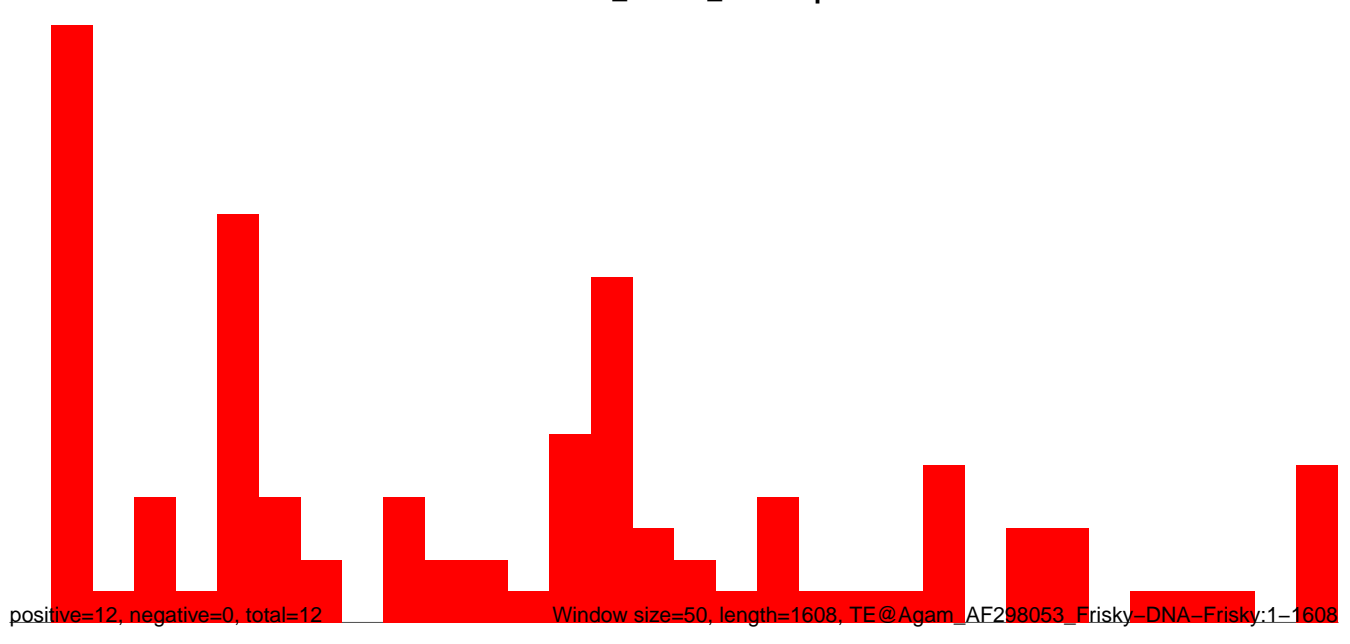
AnGam_Sua5b_cells.18_23.rep



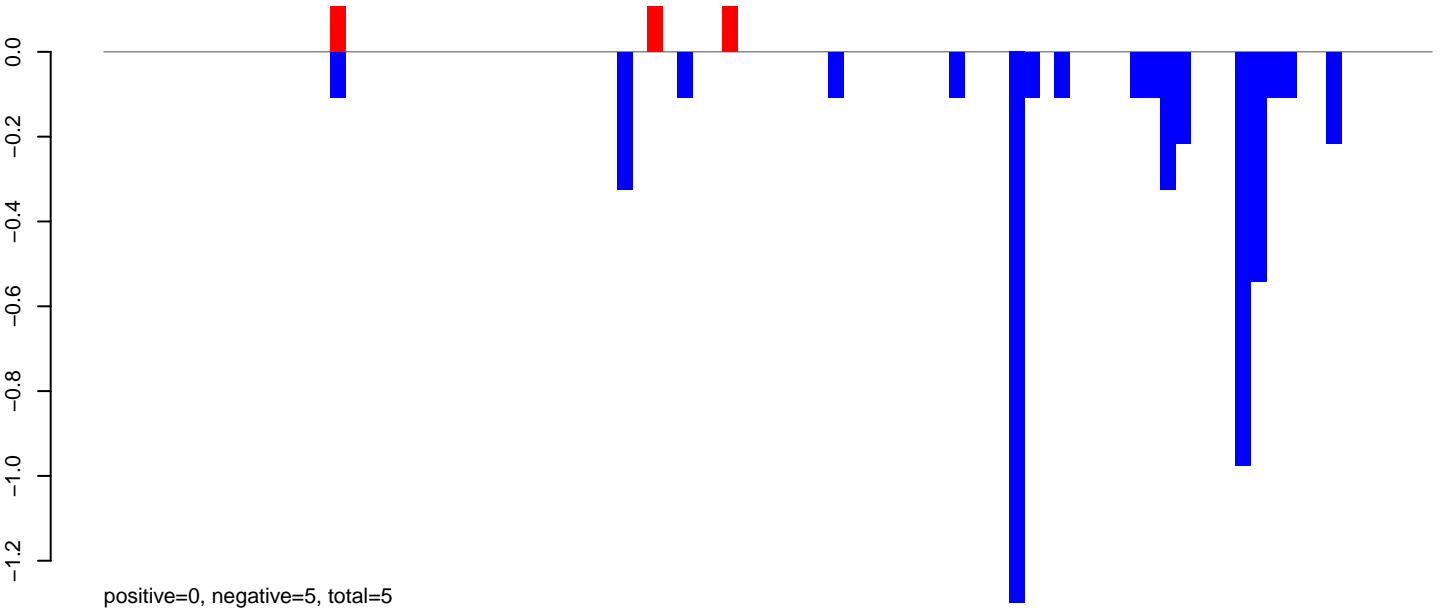
AnGam_Sua5b_cells.24_35.rep



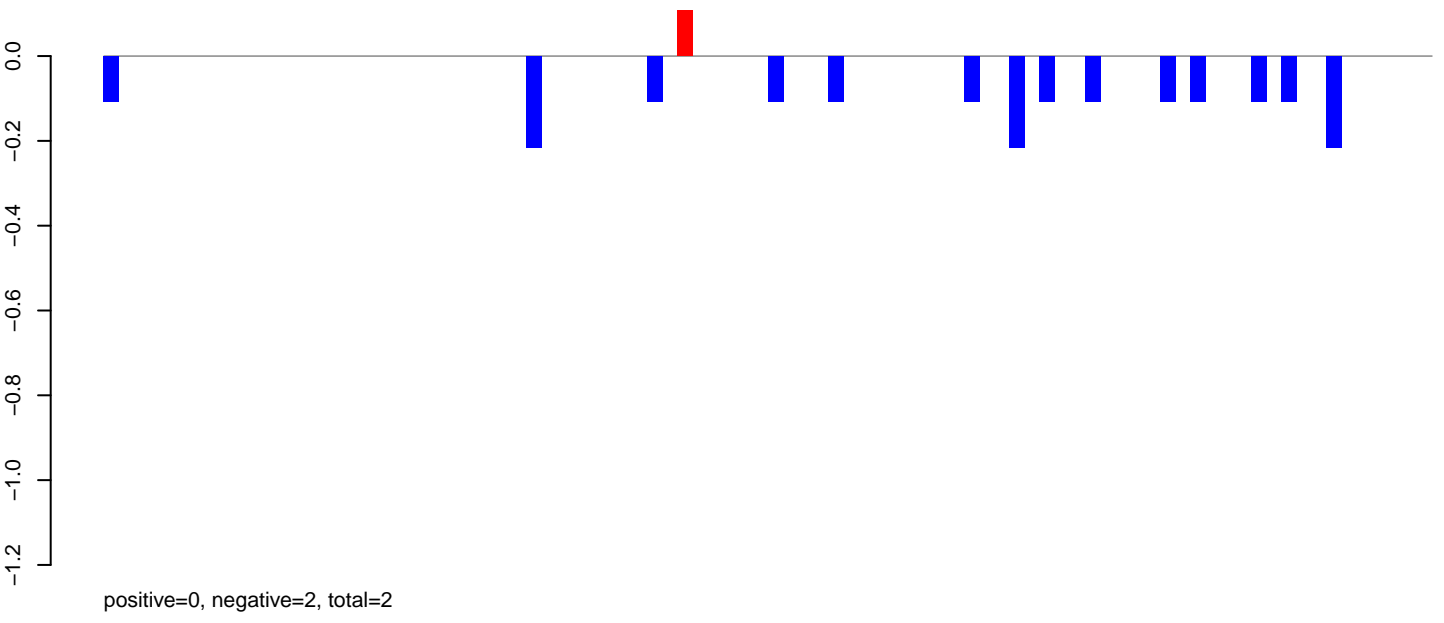
AnGam_Sua5b_cells.rep



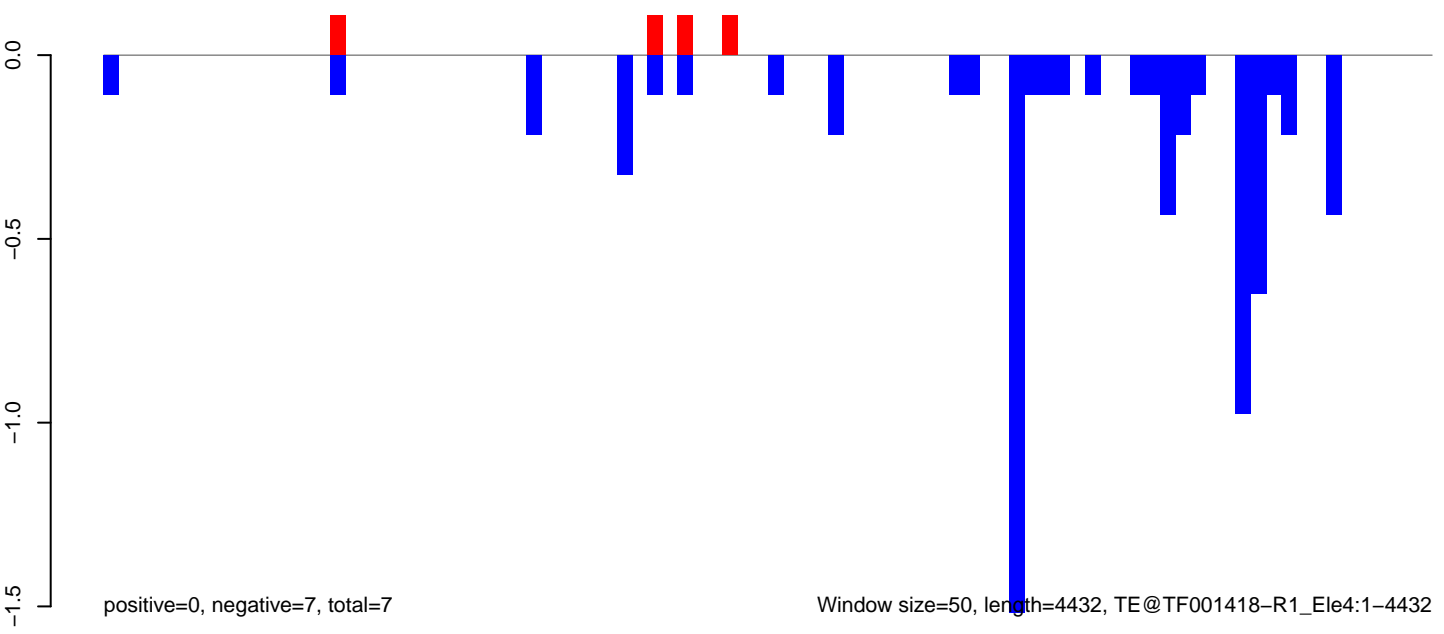
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

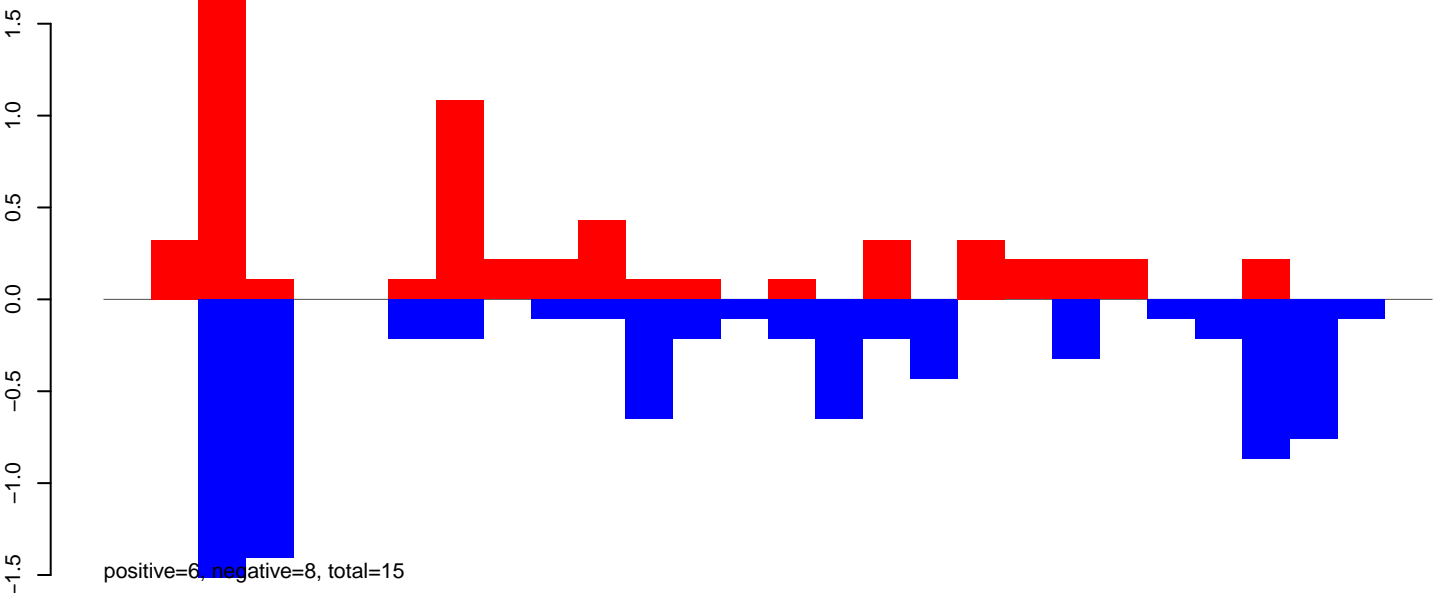


AnGam_Sua5b_cells.rep

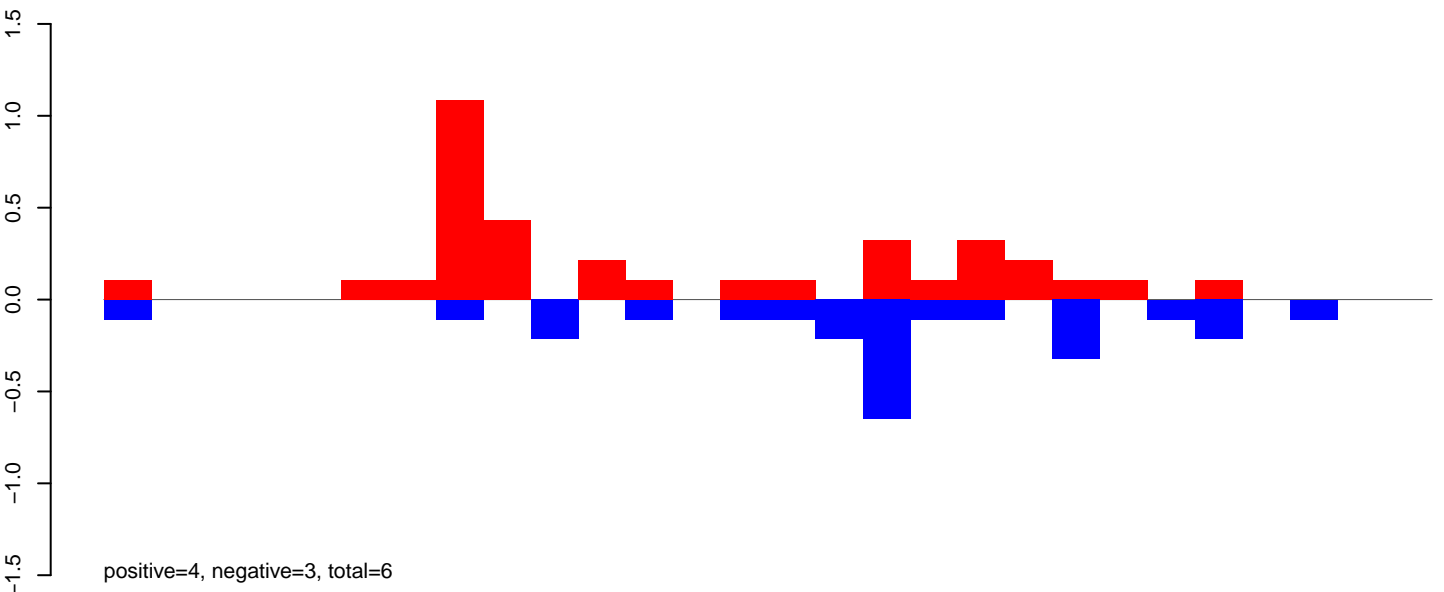


Window size=50, length=4432, TE@TF001418-R1_Ele4:1-4432

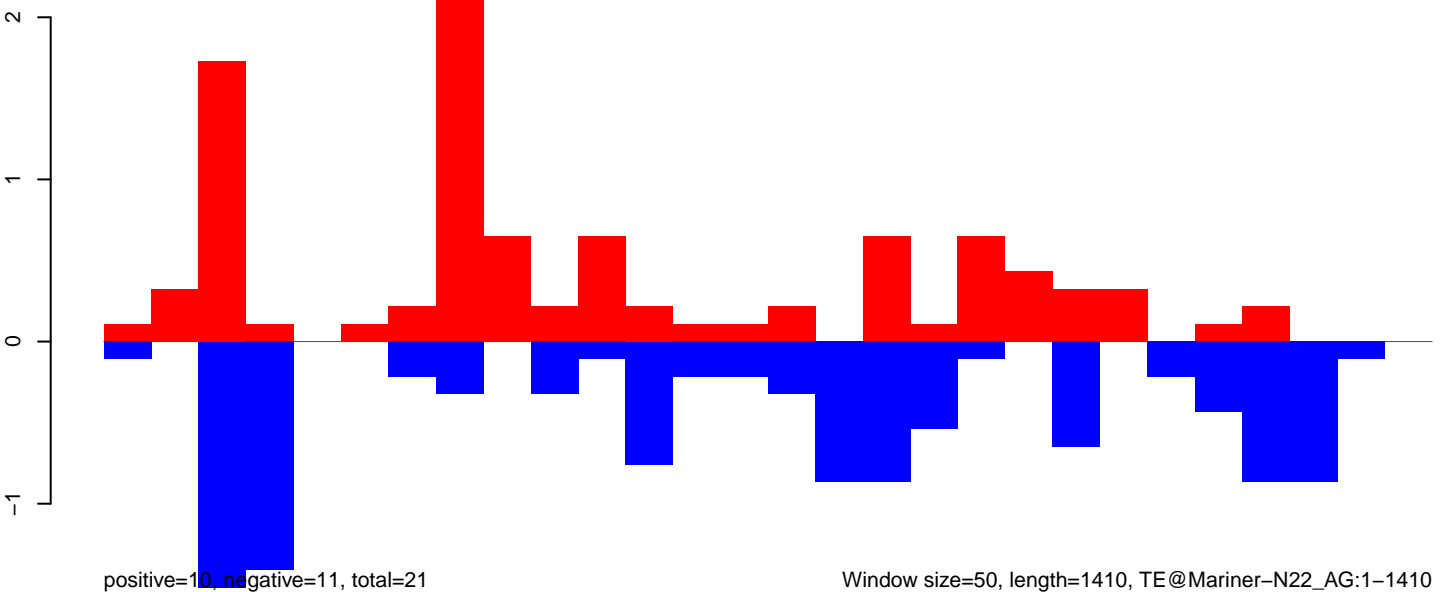
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

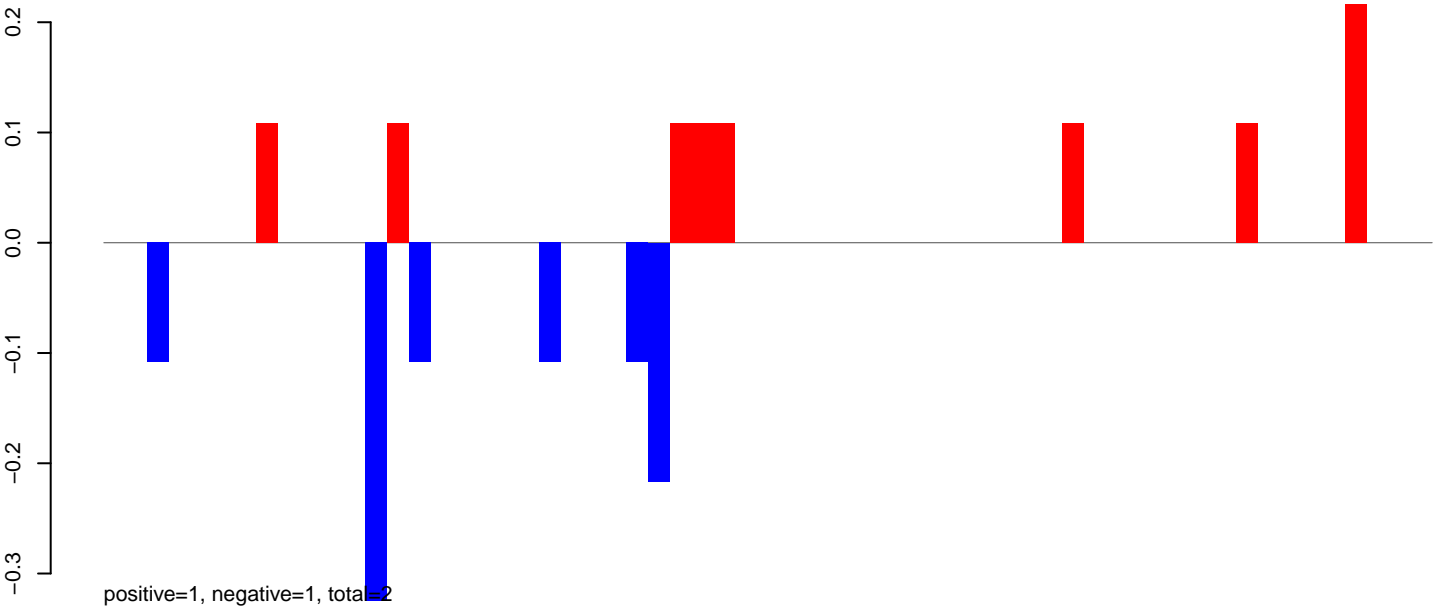


AnGam_Sua5b_cells.rep

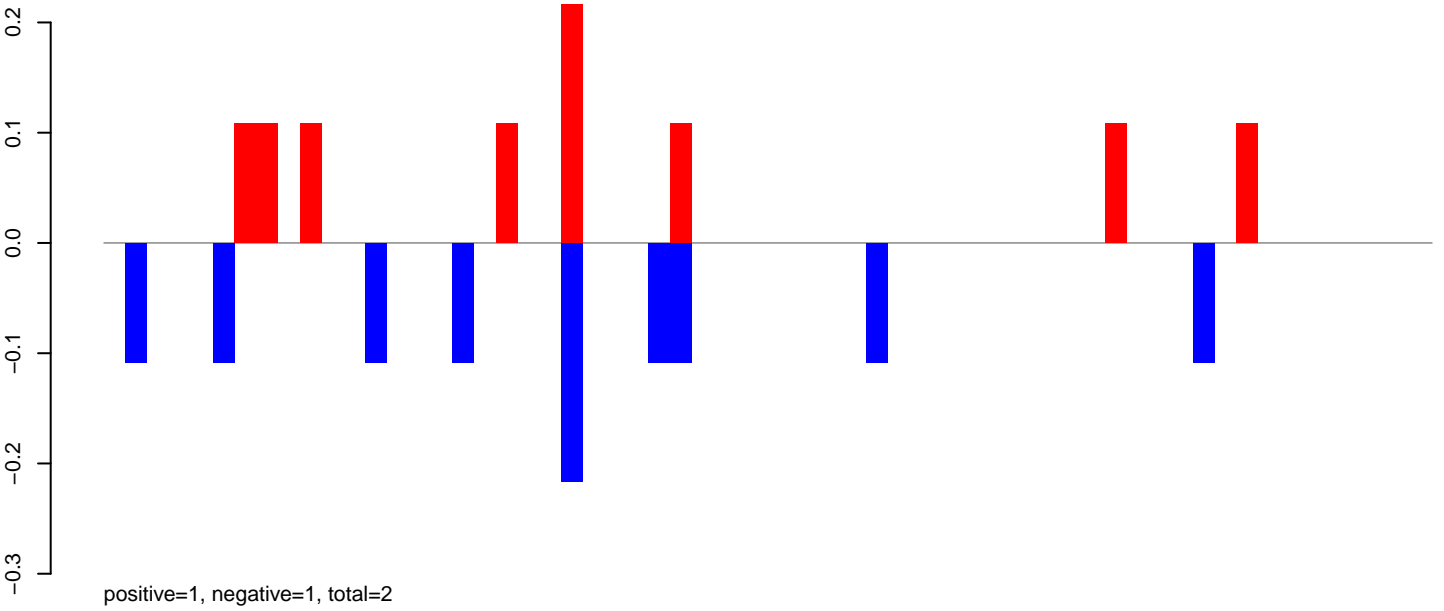


Window size=50, length=1410, TE@Mariner-N22_AG:1-1410

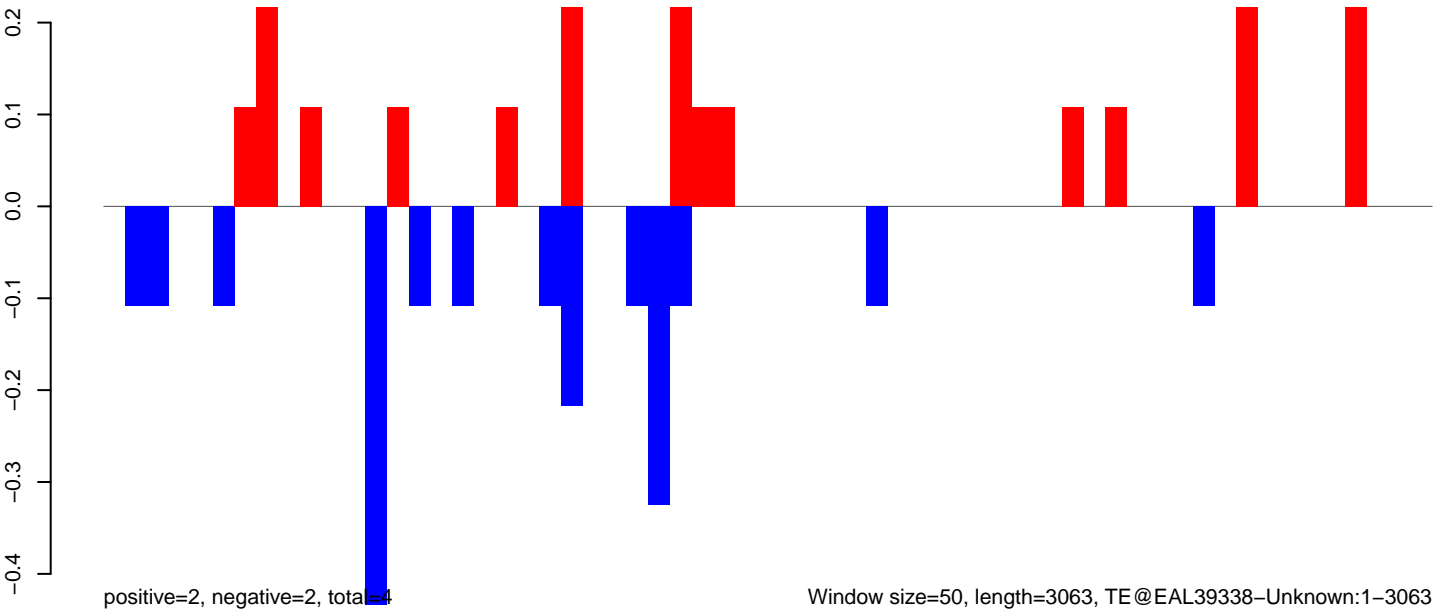
AnGam_Sua5b_cells.18_23.rep



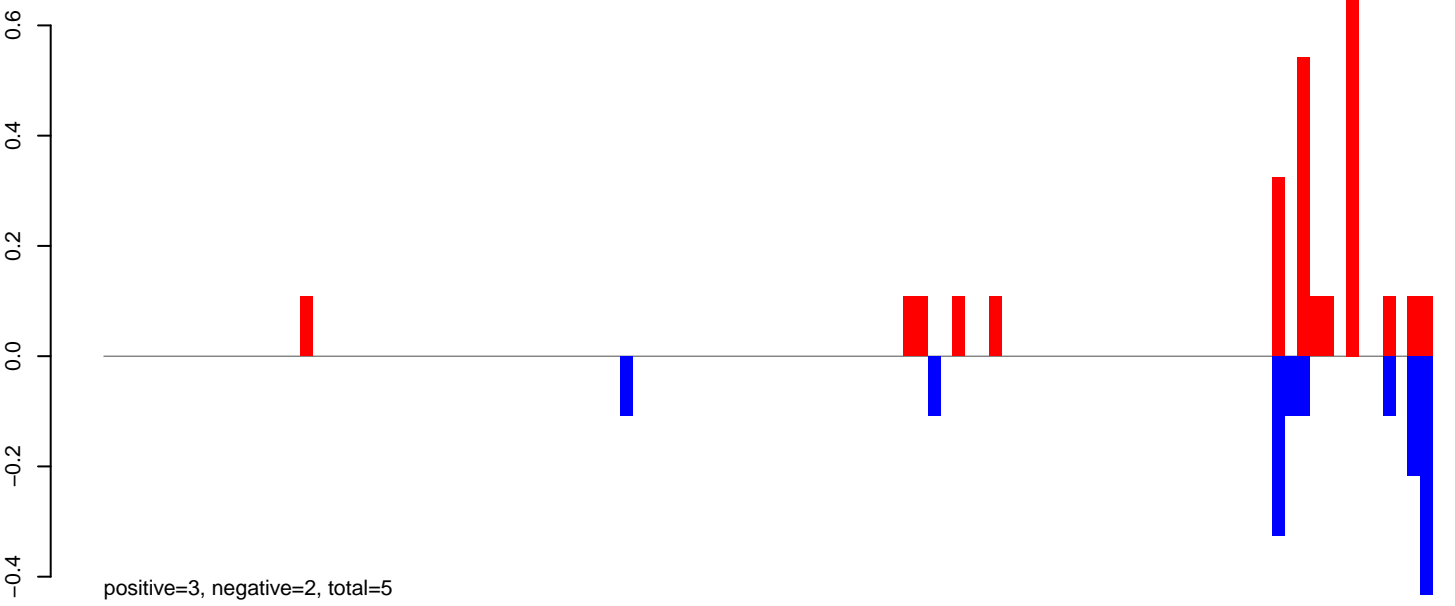
AnGam_Sua5b_cells.24_35.rep



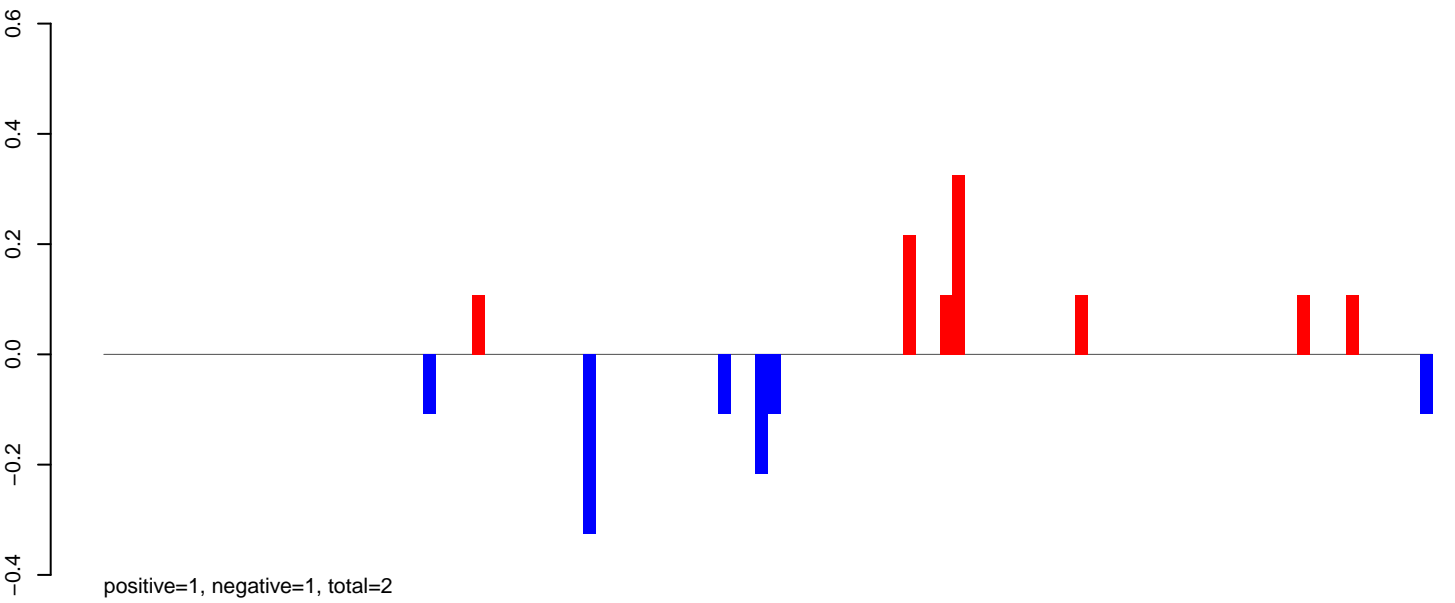
AnGam_Sua5b_cells.rep



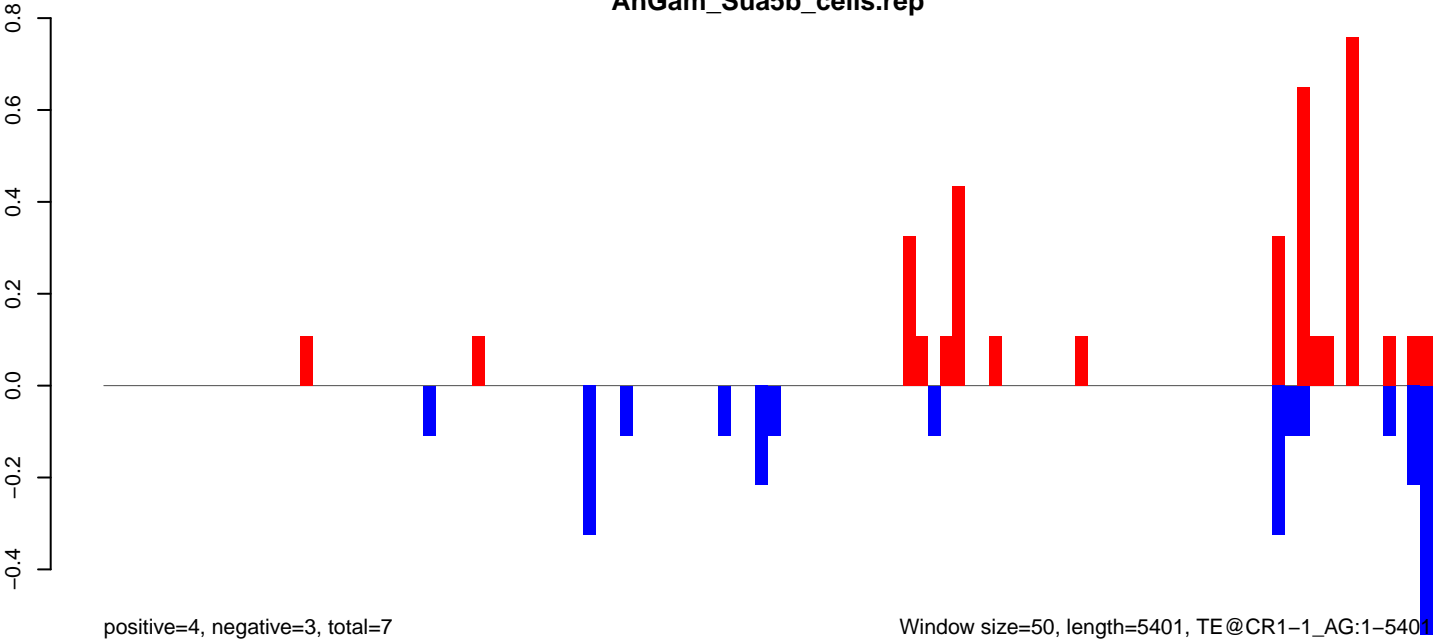
AnGam_Sua5b_cells.18_23.rep



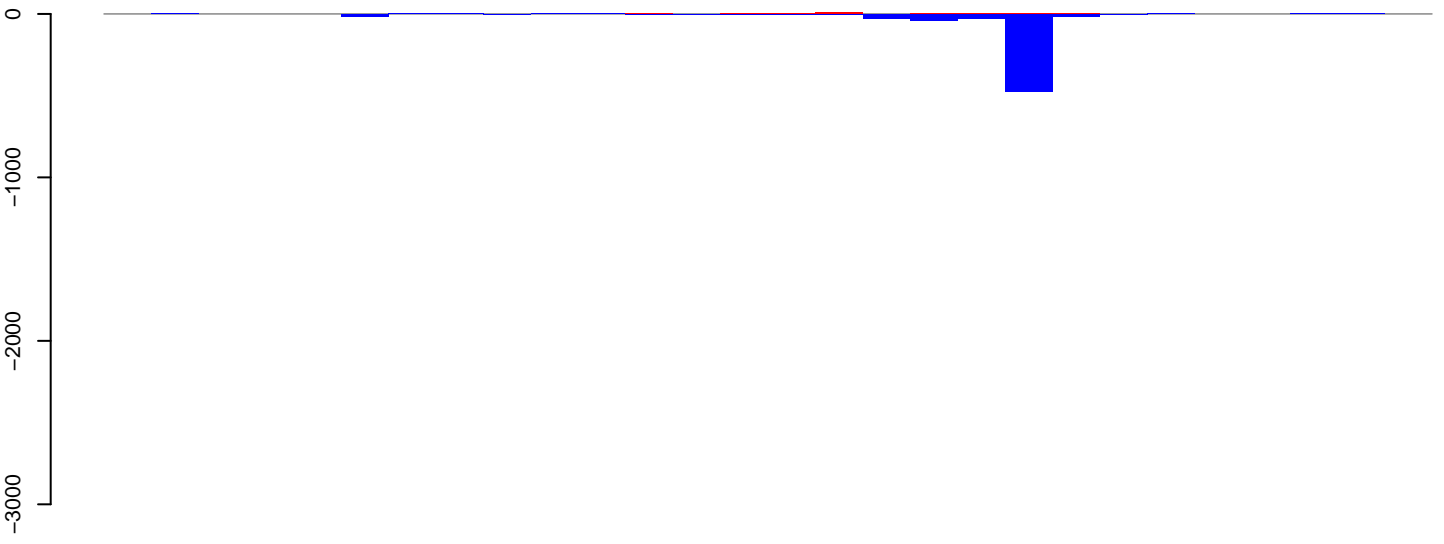
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



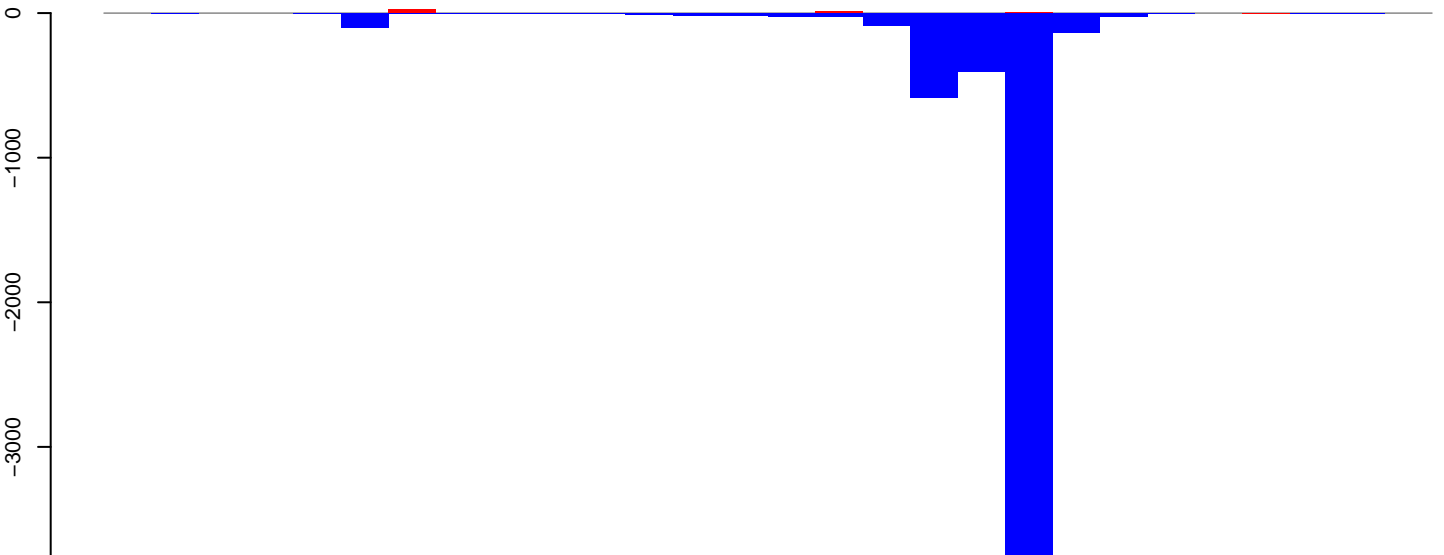
positive=13, negative=618, total=631

AnGam_Sua5b_cells.24_35.rep



positive=35, negative=4914, total=4949

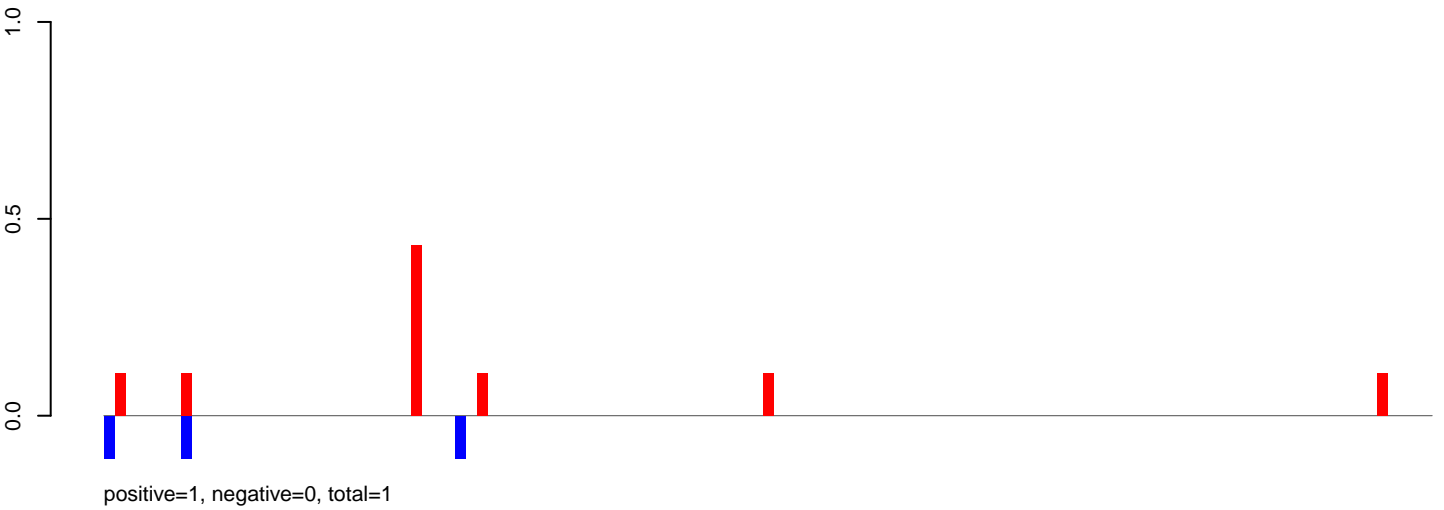
AnGam_Sua5b_cells.rep



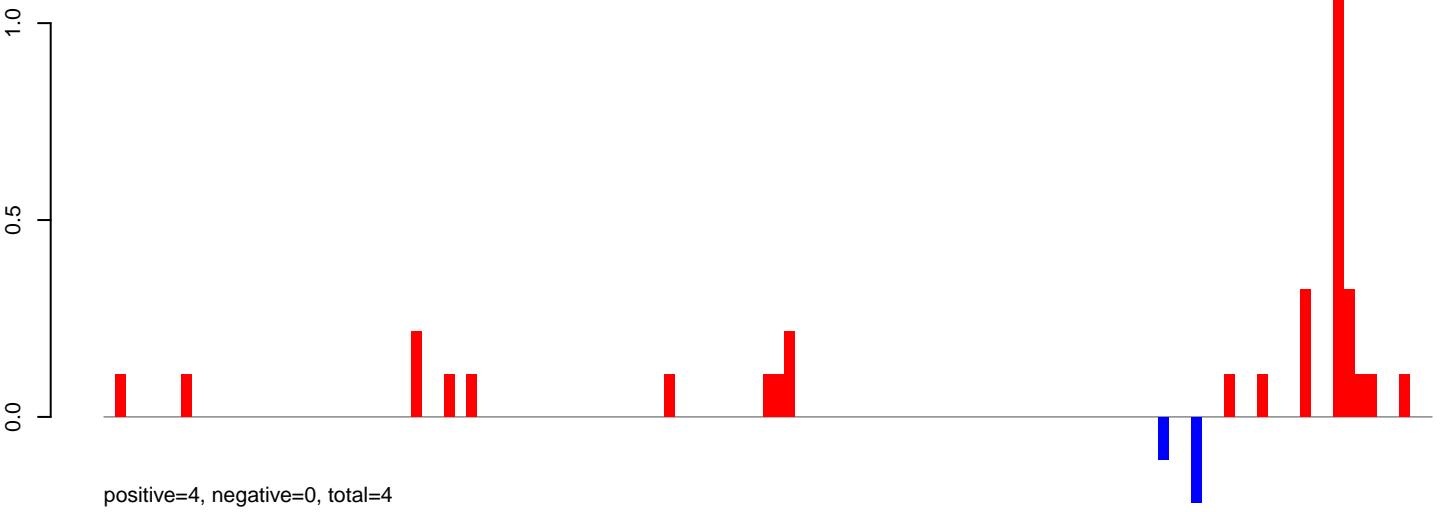
positive=48, negative=5532, total=5580

Window size=50, length=1413, TE@Tc1-1_AG:1-1413

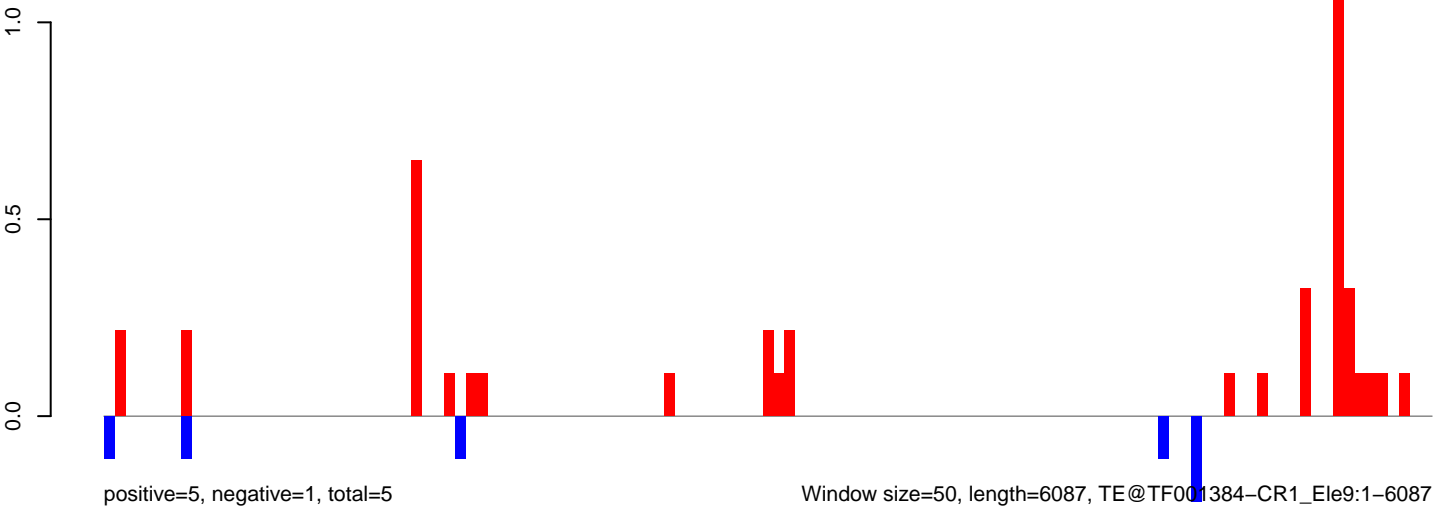
AnGam_Sua5b_cells.18_23.rep



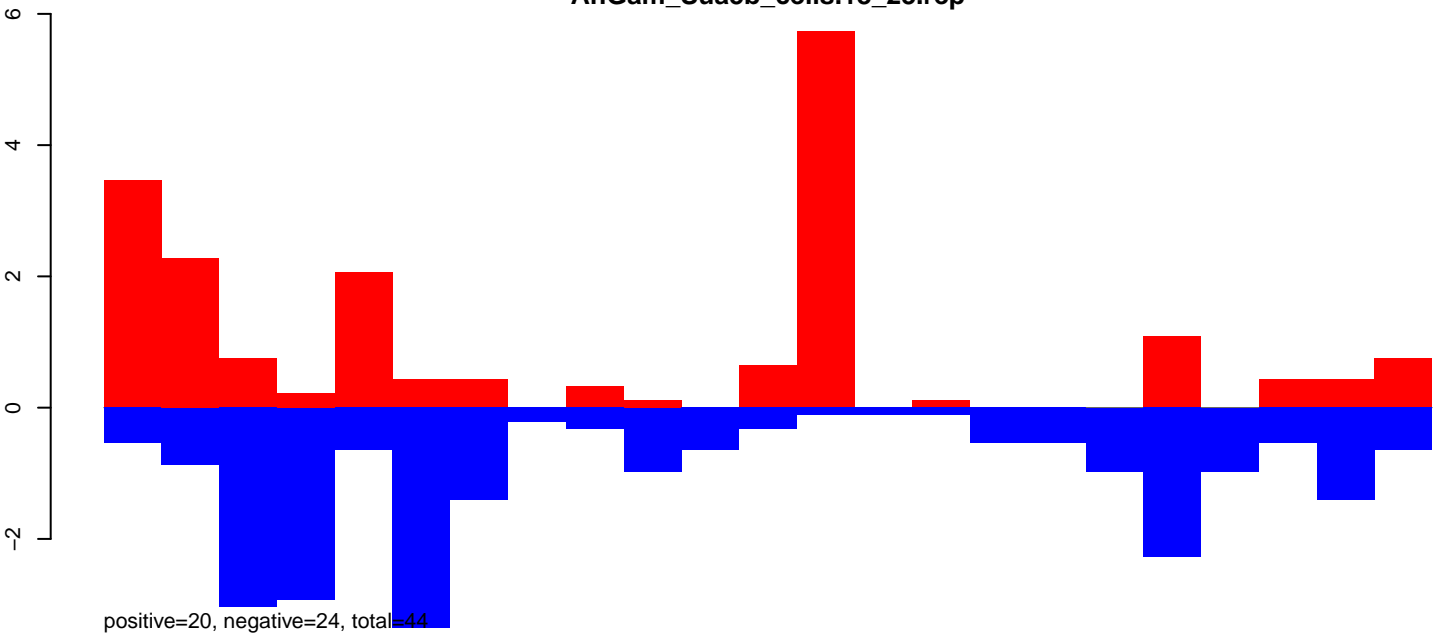
AnGam_Sua5b_cells.24_35.rep



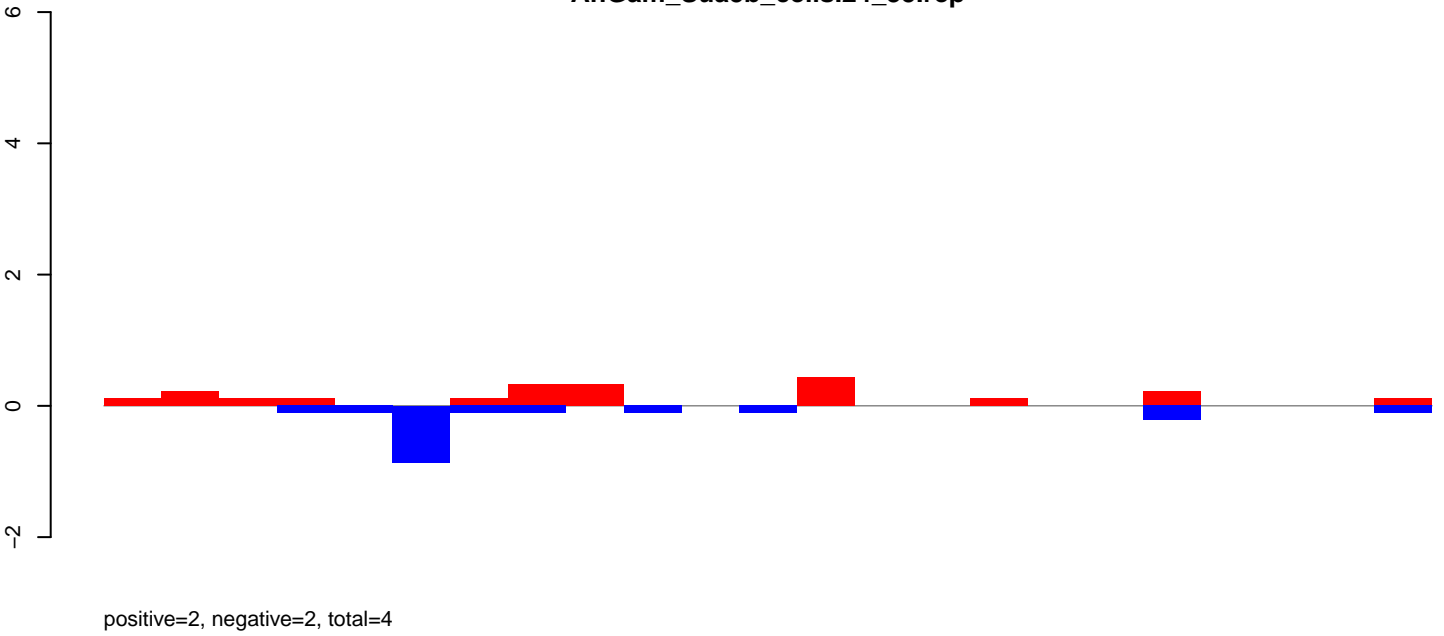
AnGam_Sua5b_cells.rep



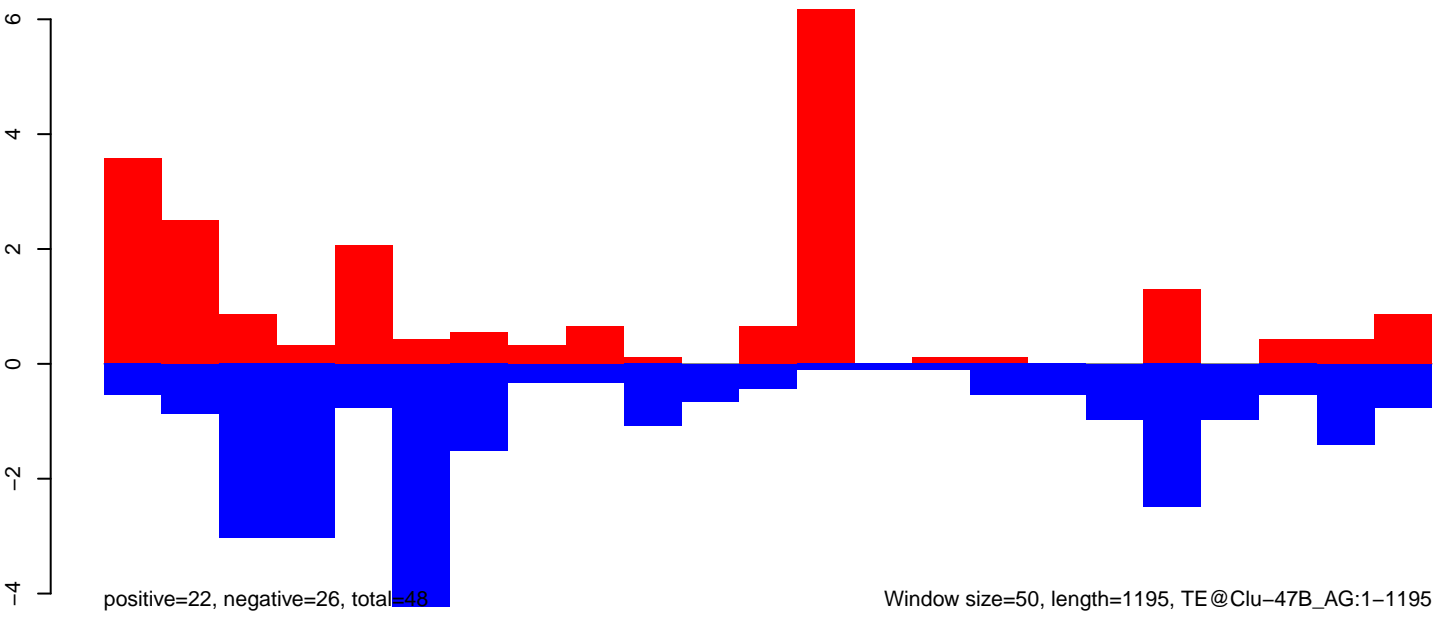
AnGam_Sua5b_cells.18_23.rep



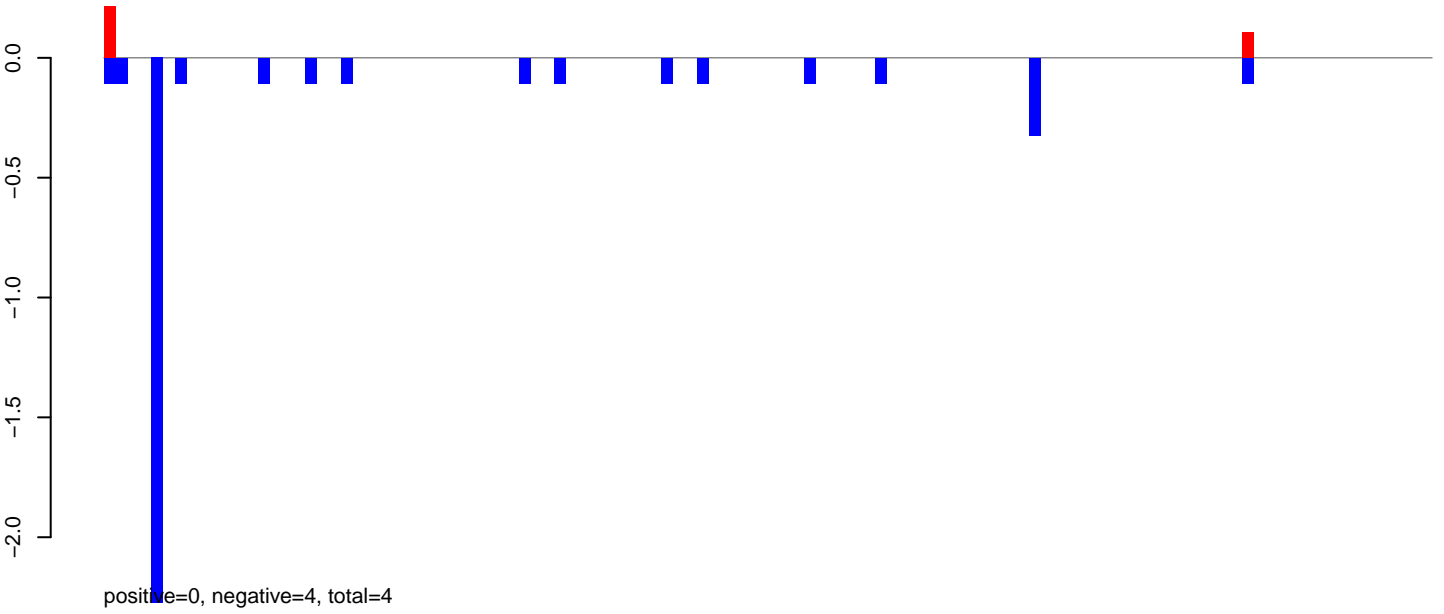
AnGam_Sua5b_cells.24_35.rep



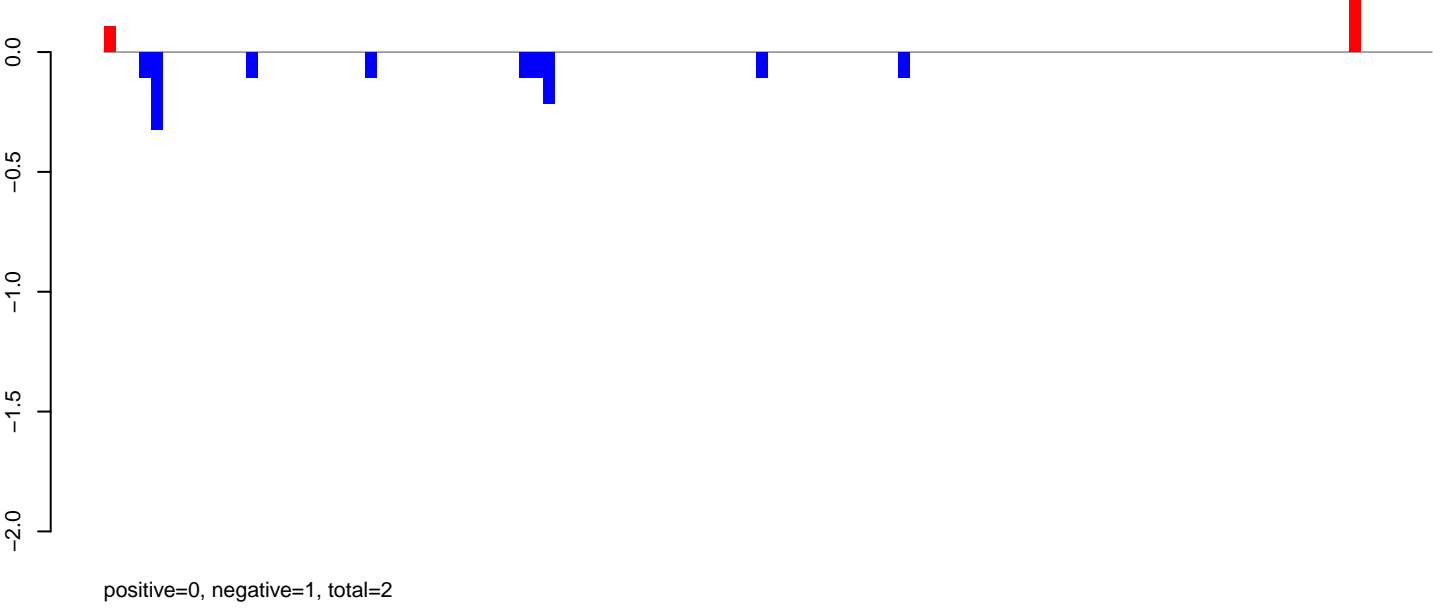
AnGam_Sua5b_cells.rep



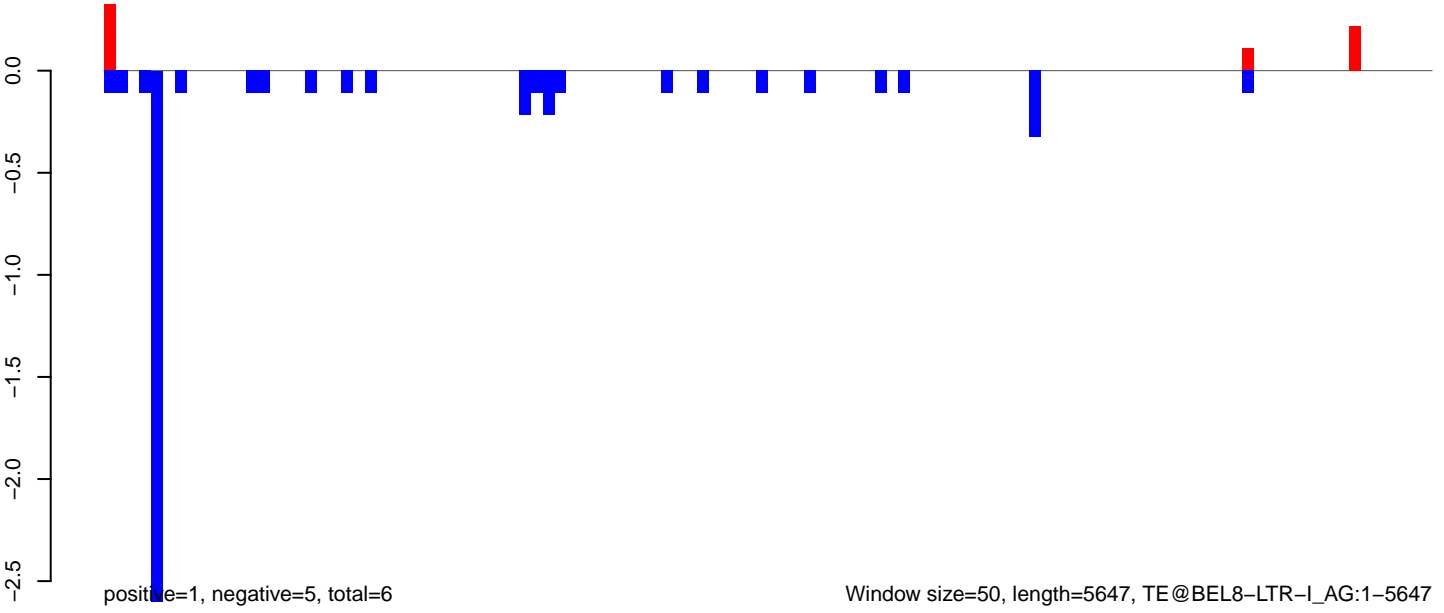
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



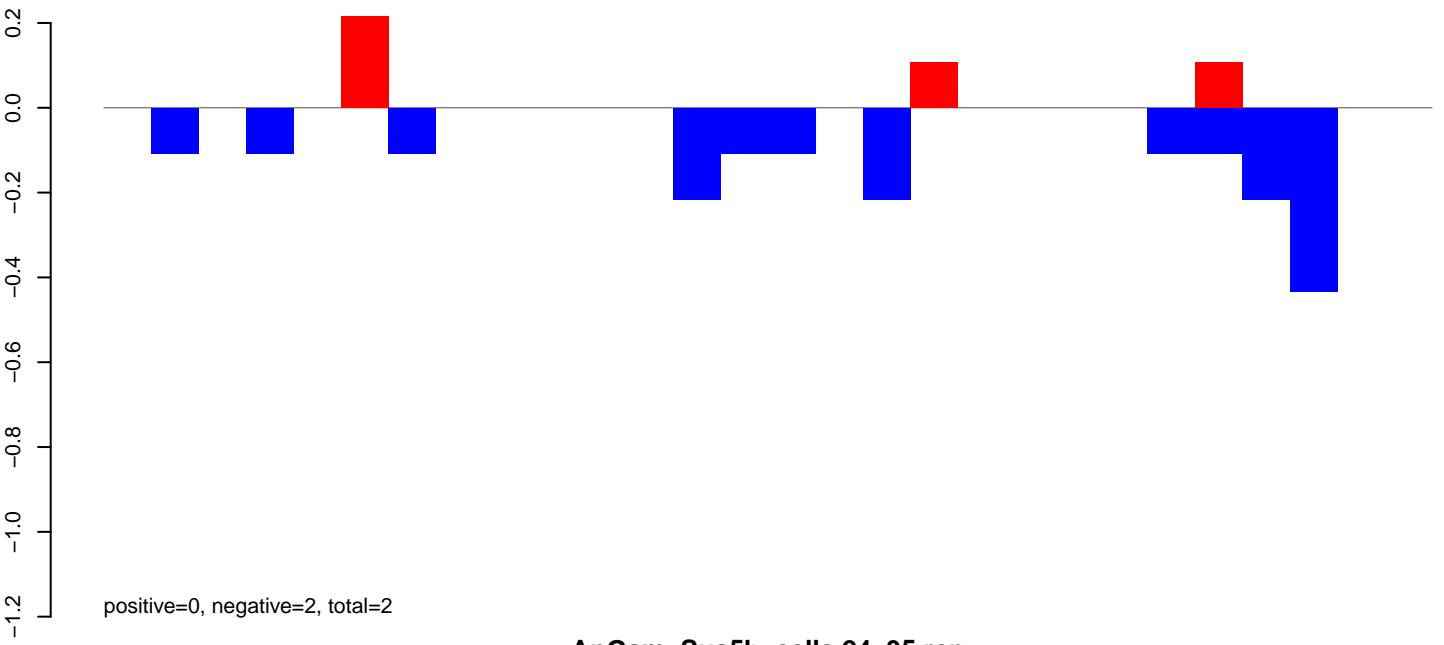
AnGam_Sua5b_cells.rep



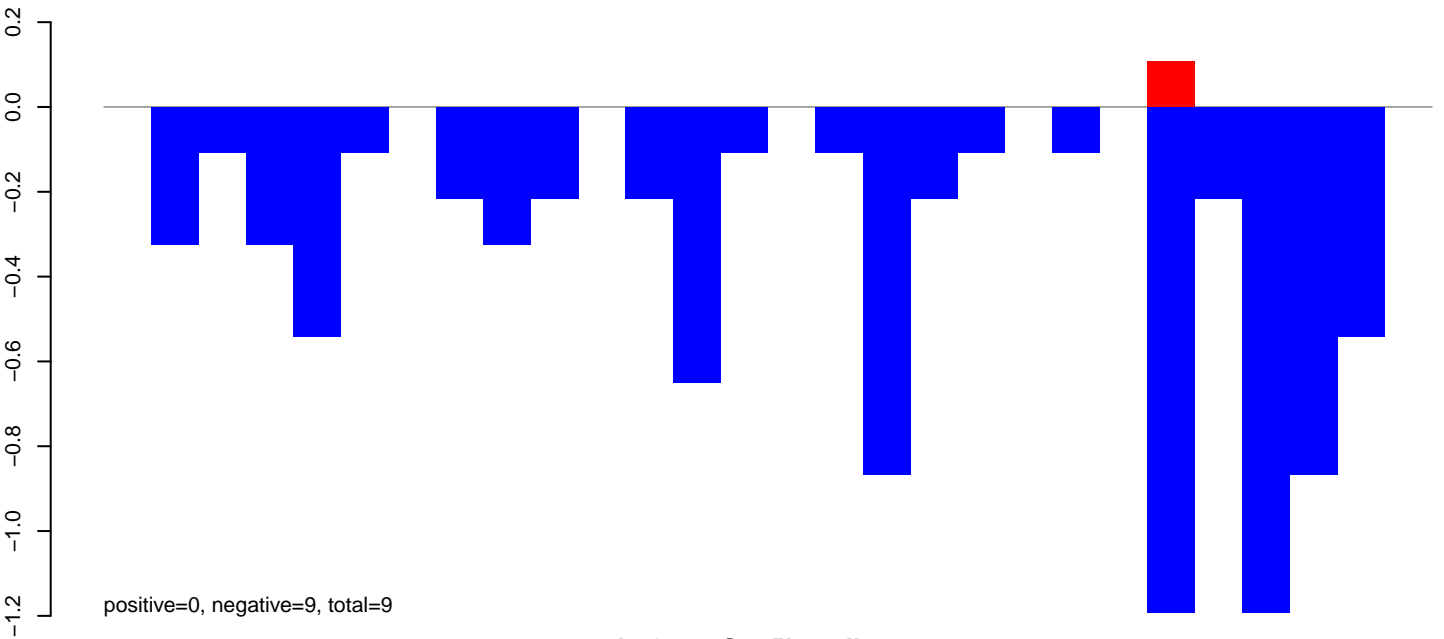
Window size=50, length=5647, TE@BEL8-LTR-I_AG:1-5647

0 1000 2000 3000 4000 5000

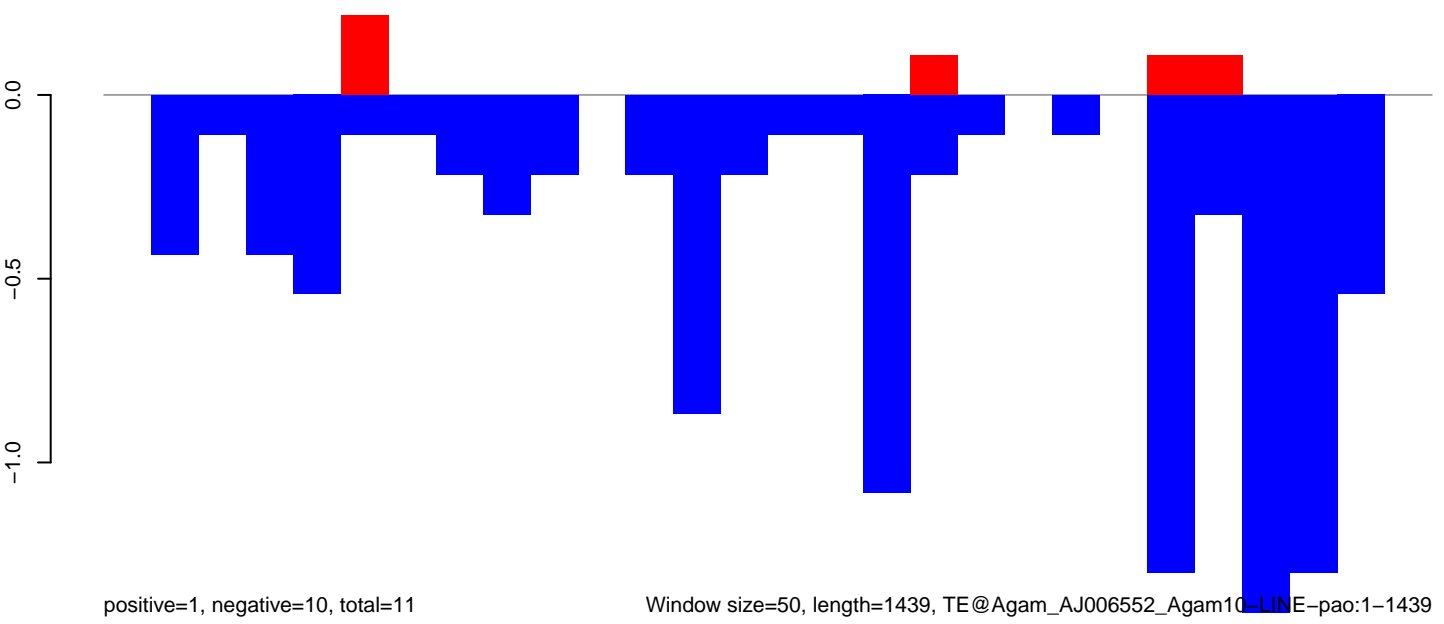
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



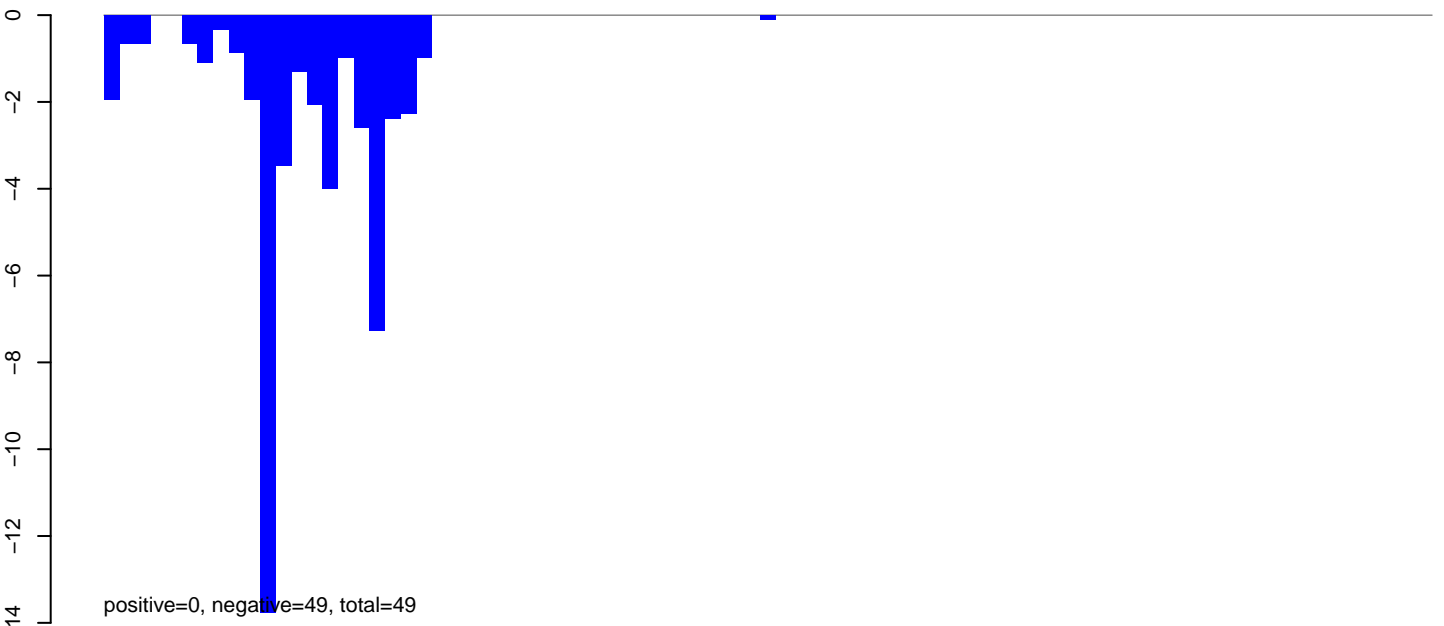
AnGam_Sua5b_cells.rep



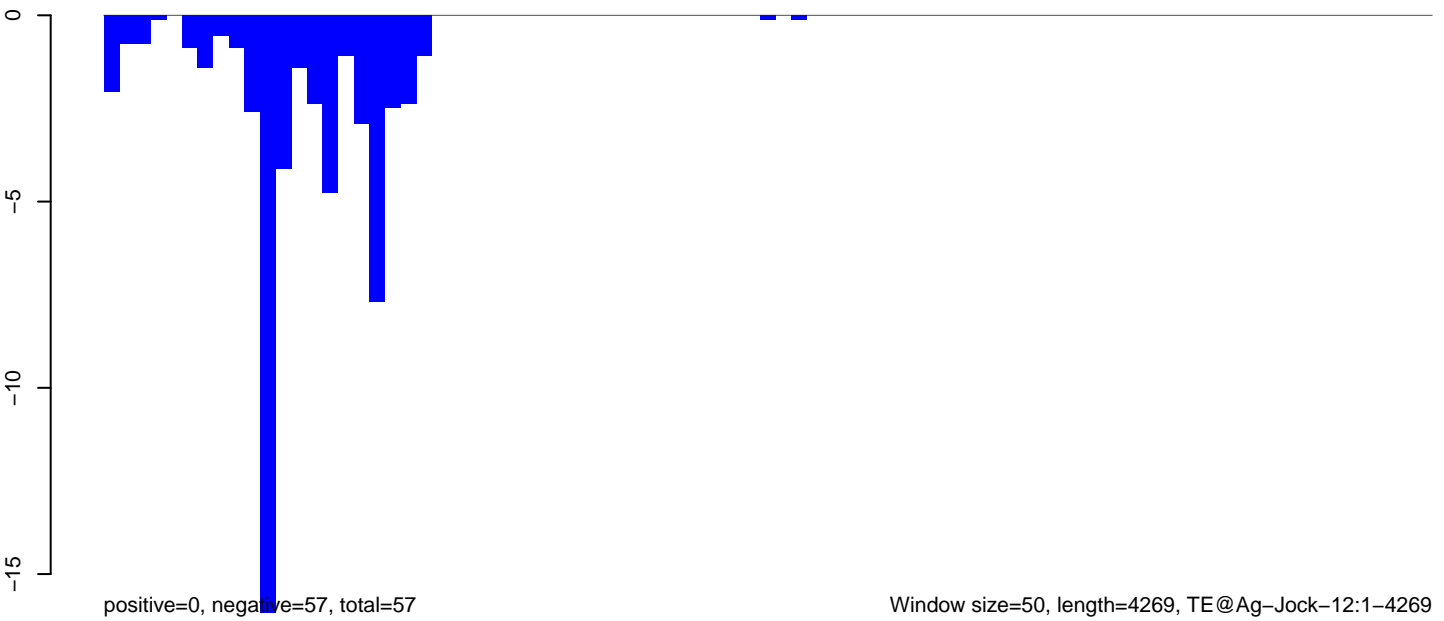
AnGam_Sua5b_cells.18_23.rep



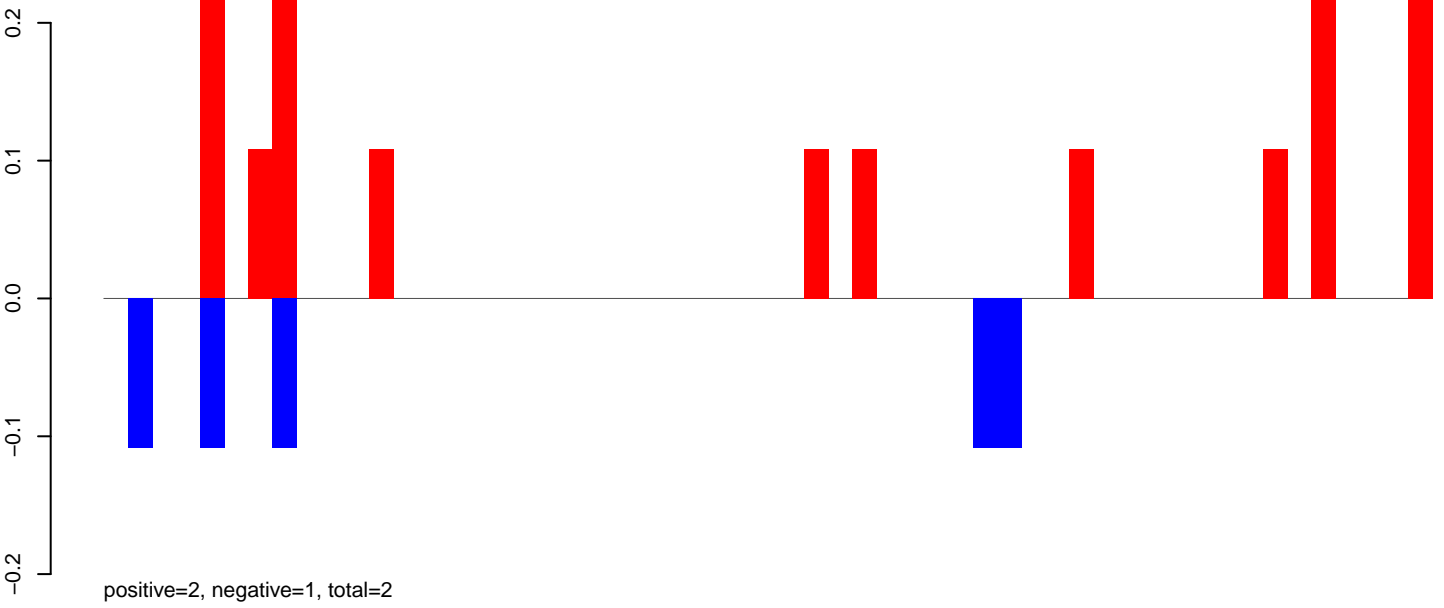
AnGam_Sua5b_cells.24_35.rep



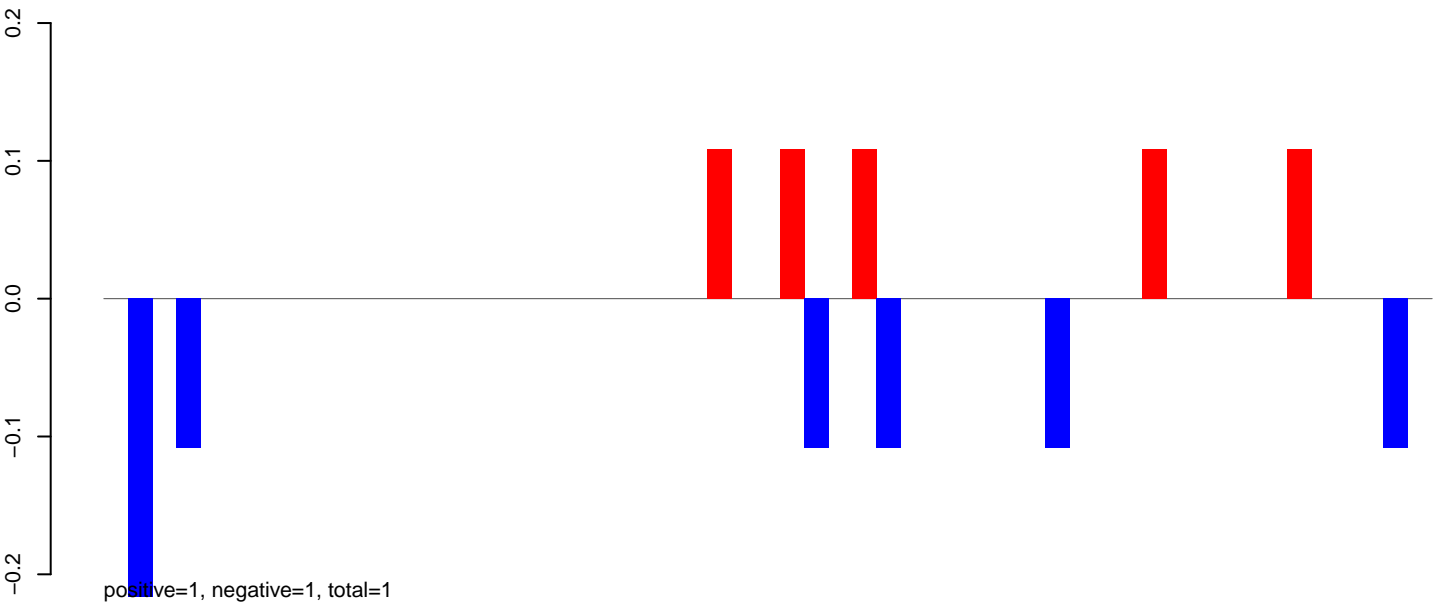
AnGam_Sua5b_cells.rep



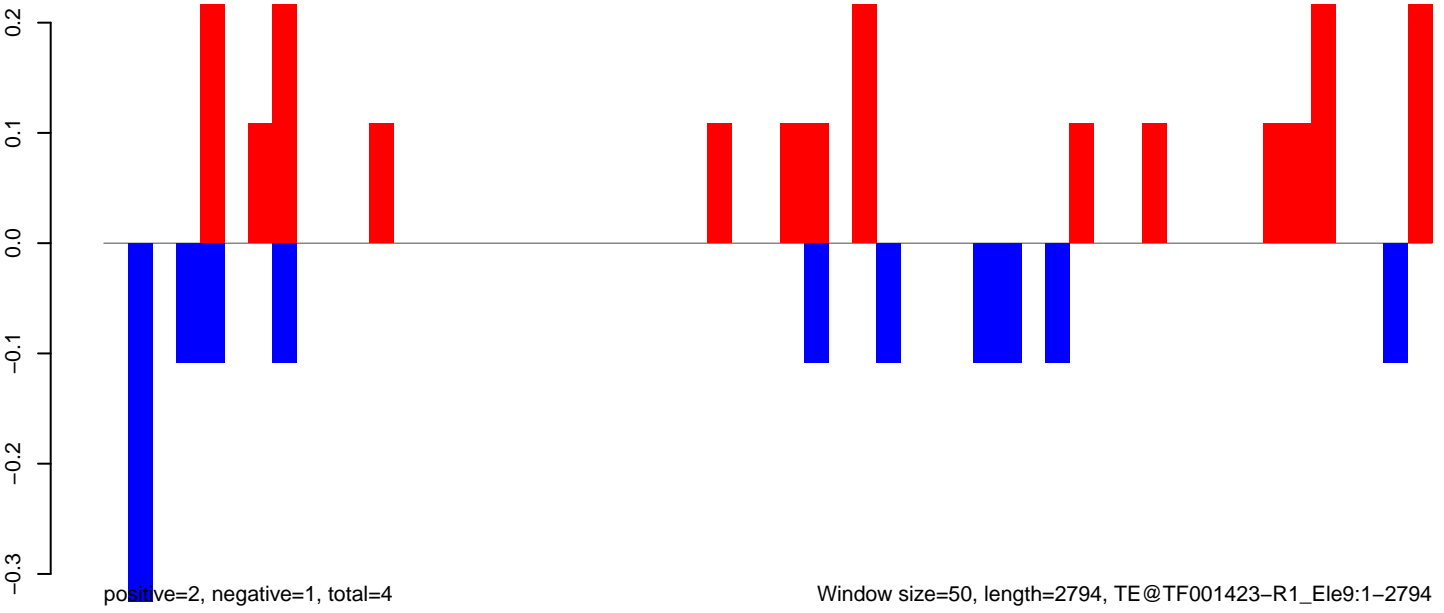
AnGam_Sua5b_cells.18_23.rep



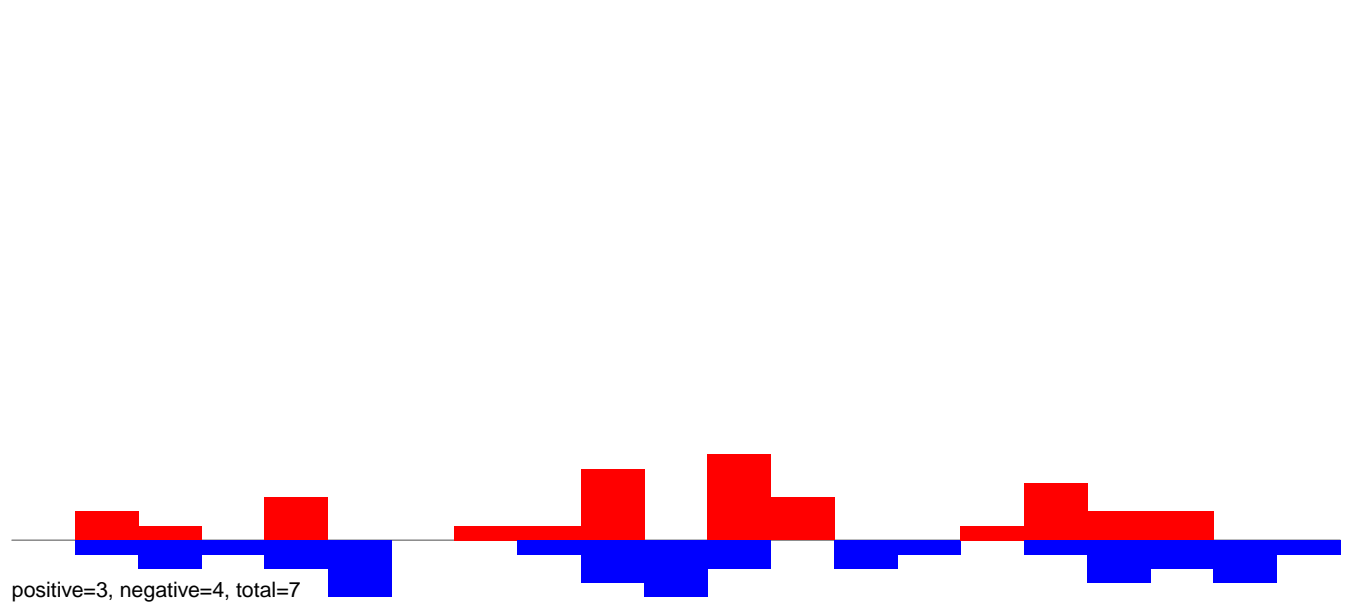
AnGam_Sua5b_cells.24_35.rep



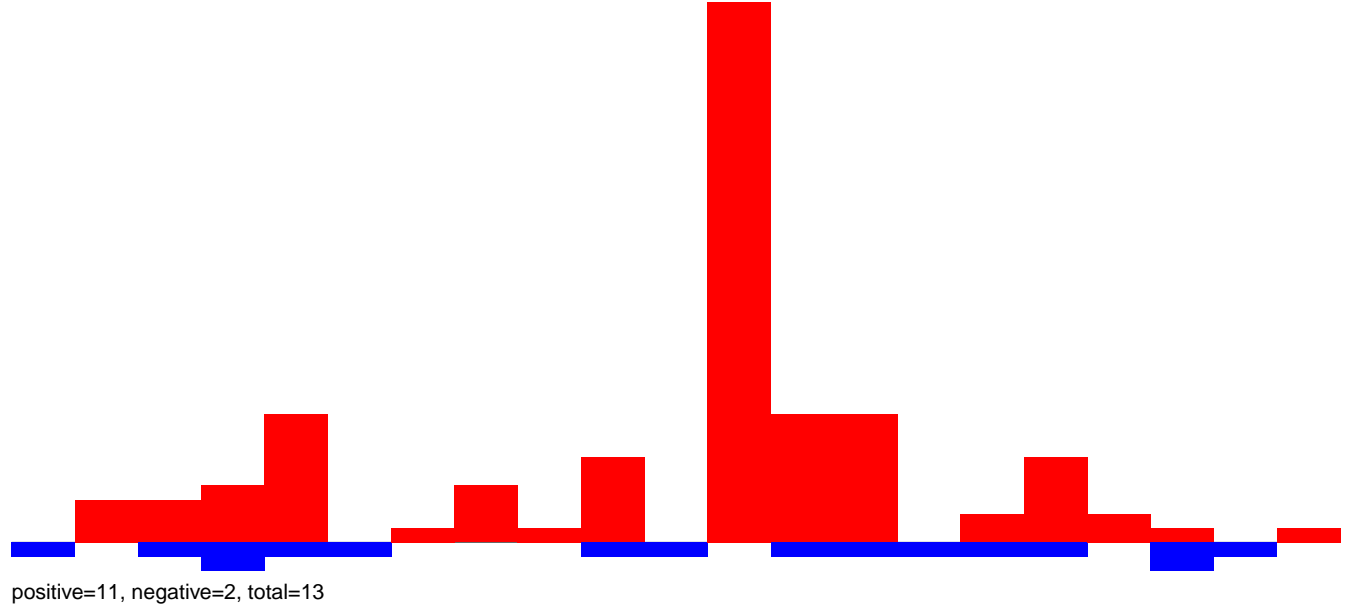
AnGam_Sua5b_cells.rep



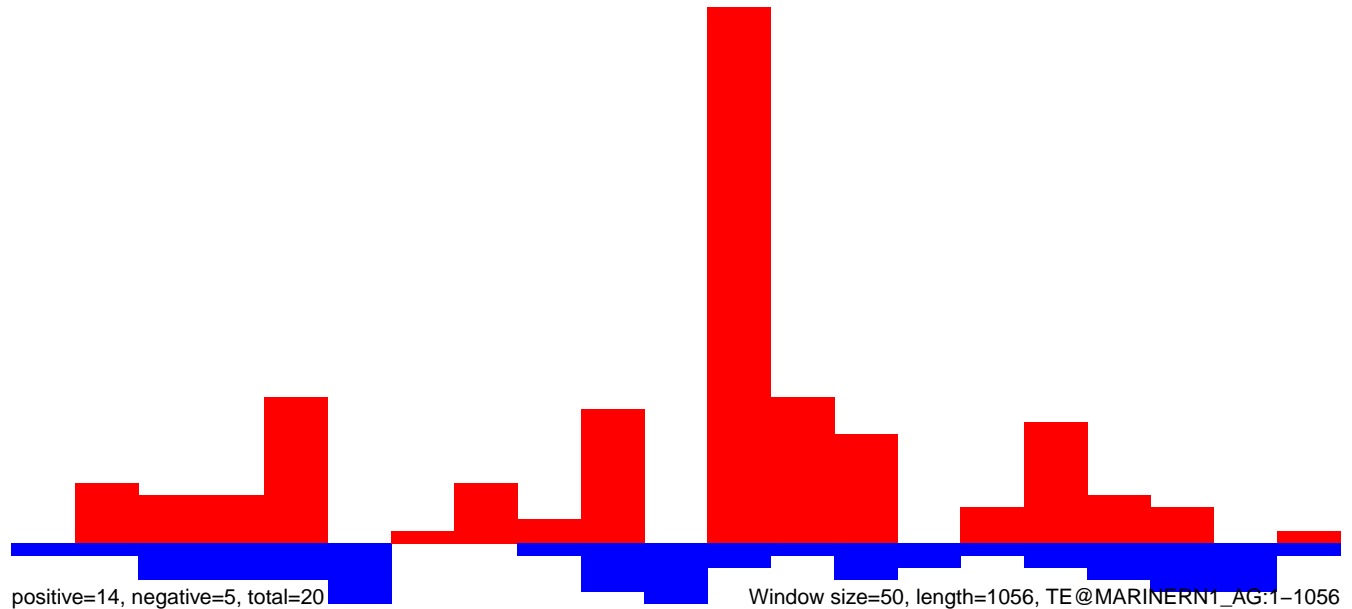
AnGam_Sua5b_cells.18_23.rep



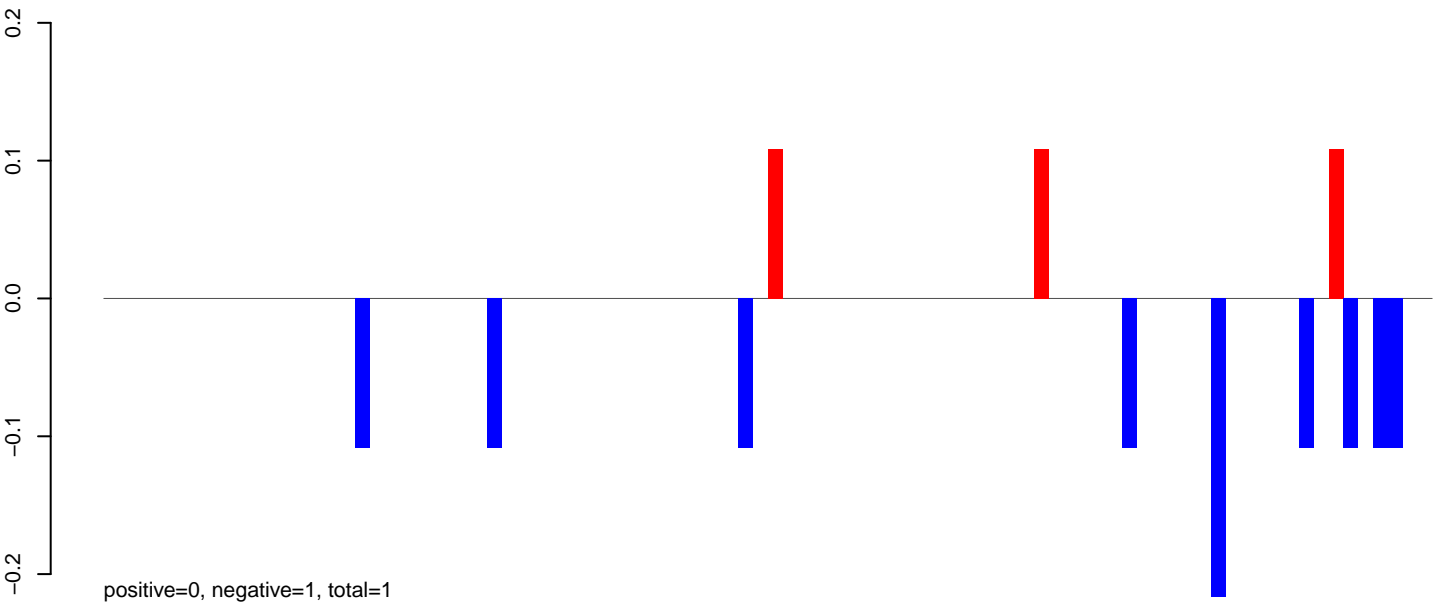
AnGam_Sua5b_cells.24_35.rep



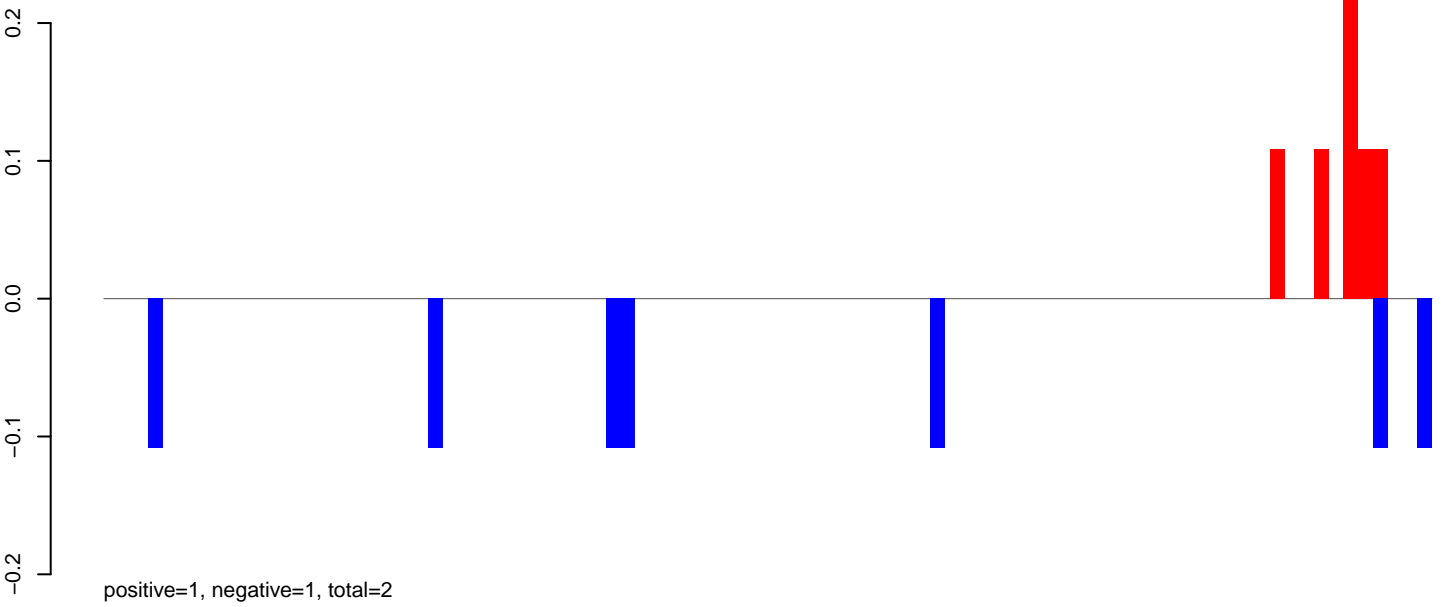
AnGam_Sua5b_cells.rep



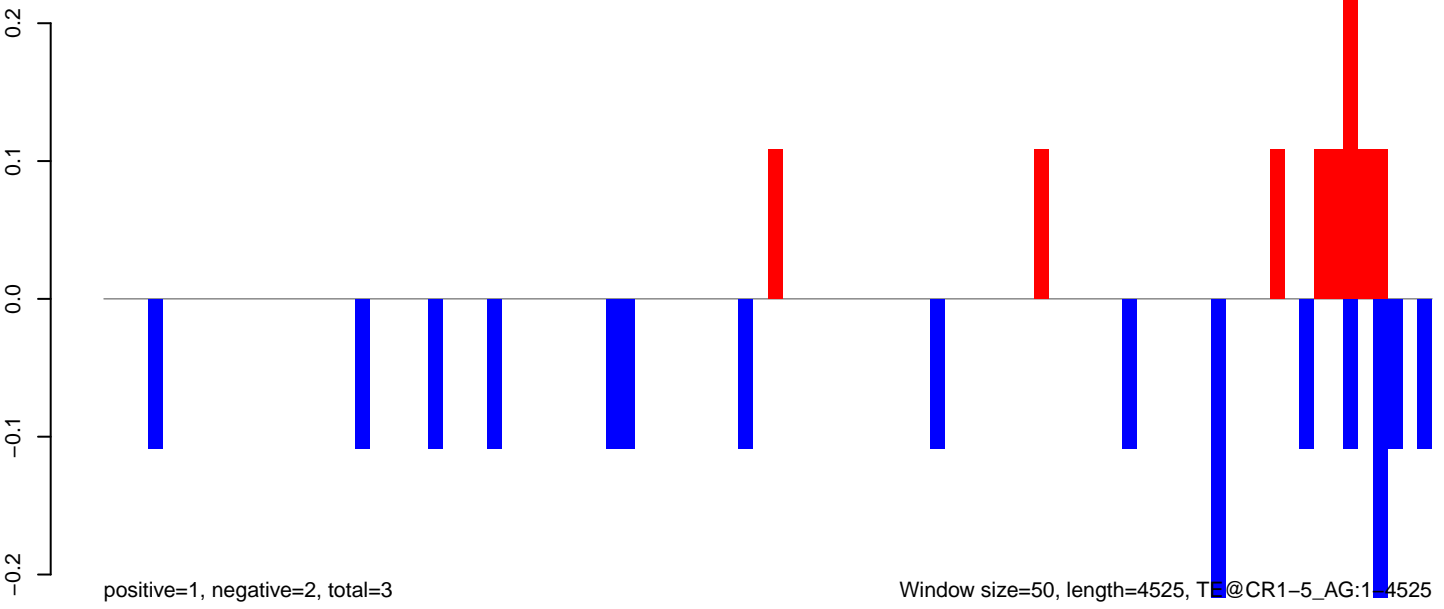
AnGam_Sua5b_cells.18_23.rep



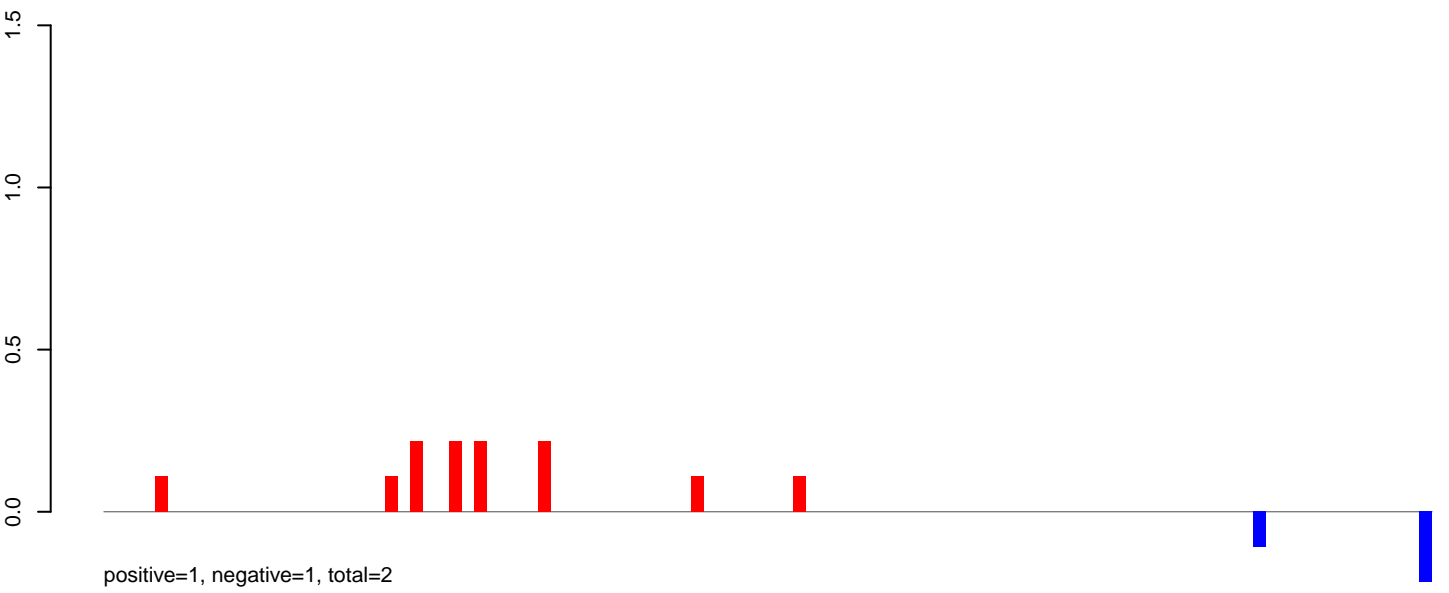
AnGam_Sua5b_cells.24_35.rep



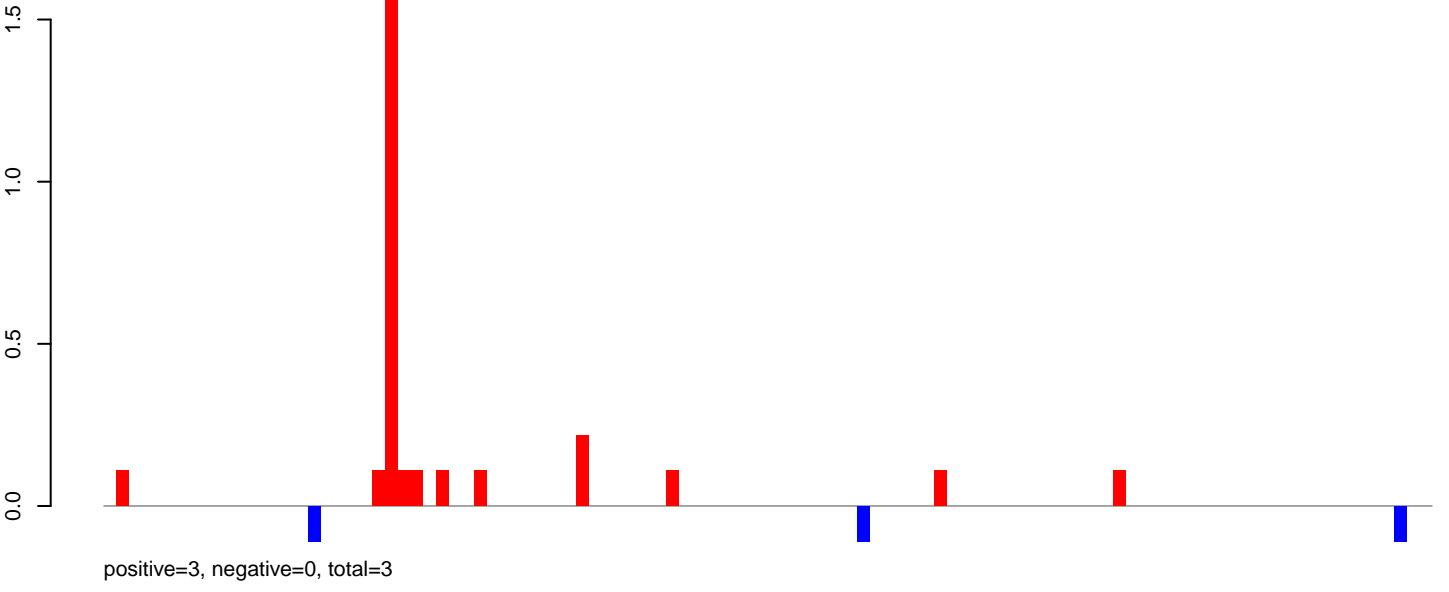
AnGam_Sua5b_cells.rep



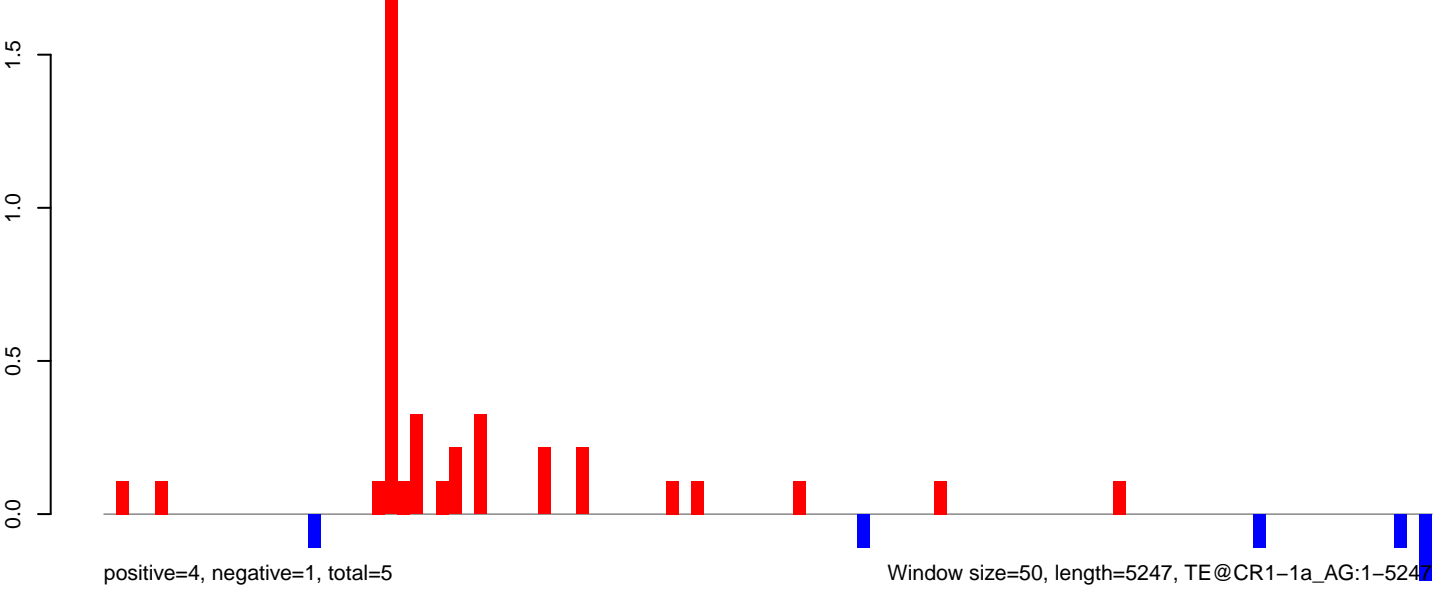
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

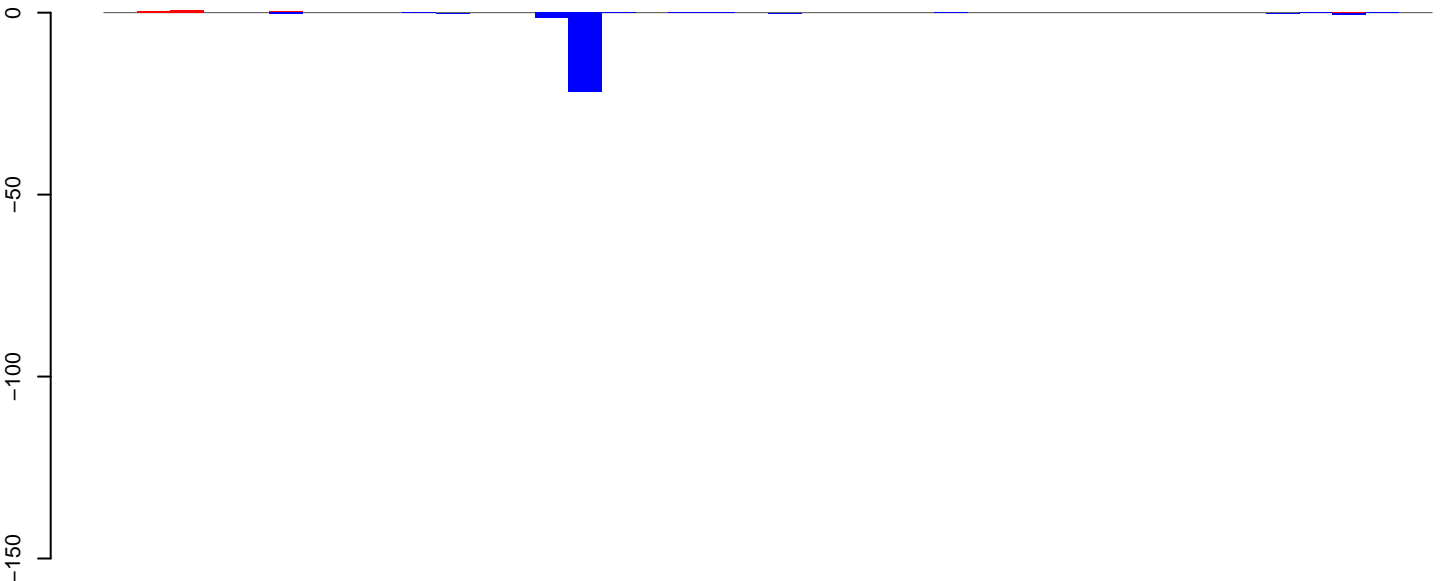


AnGam_Sua5b_cells.rep



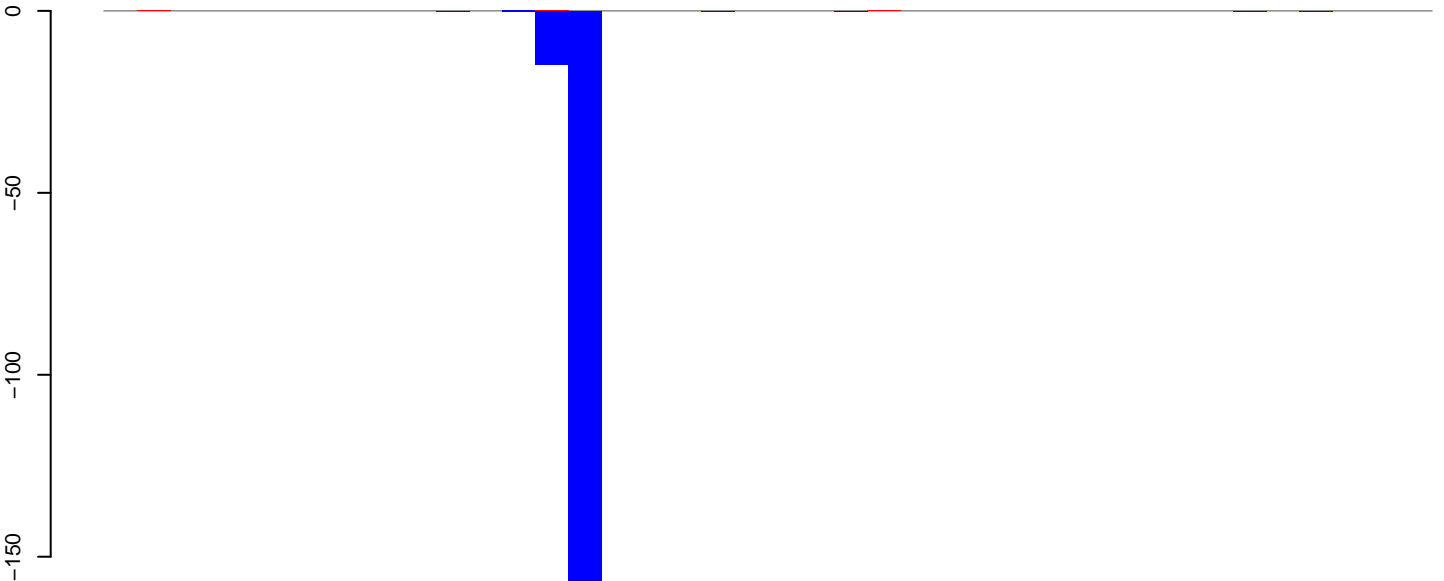
Window size=50, length=5247, TE@CR1-1a_AG:1-5247

AnGam_Sua5b_cells.18_23.rep



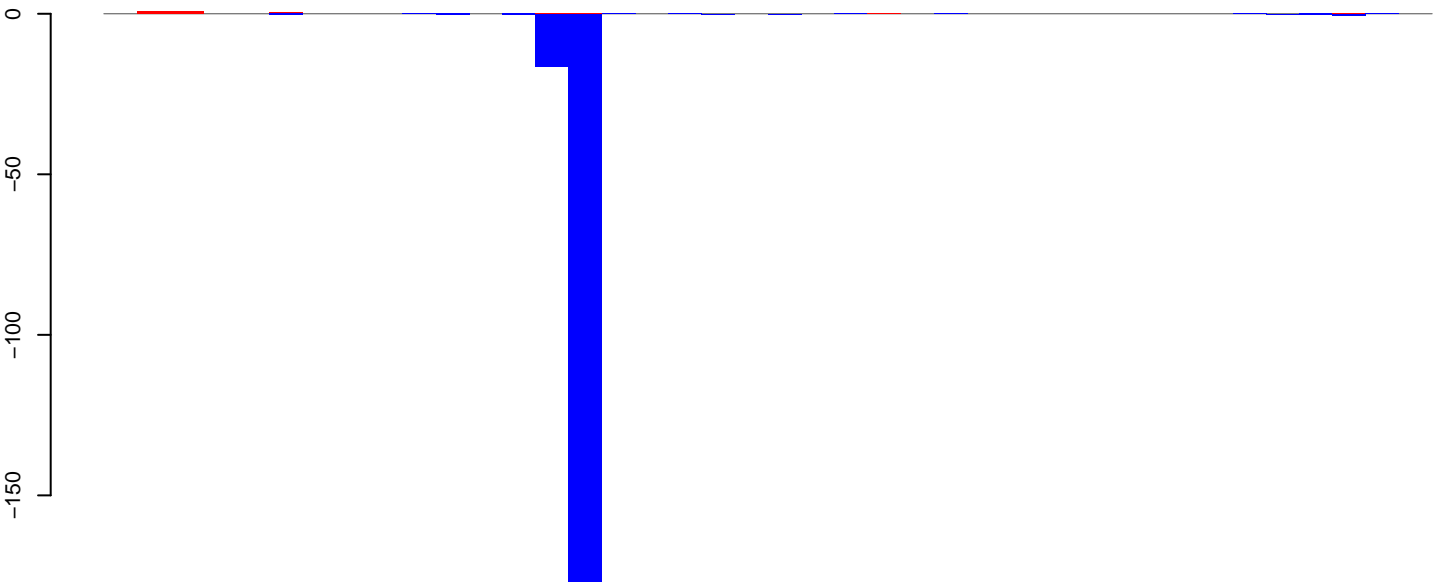
positive=2, negative=26, total=28

AnGam_Sua5b_cells.24_35.rep



positive=0, negative=179, total=180

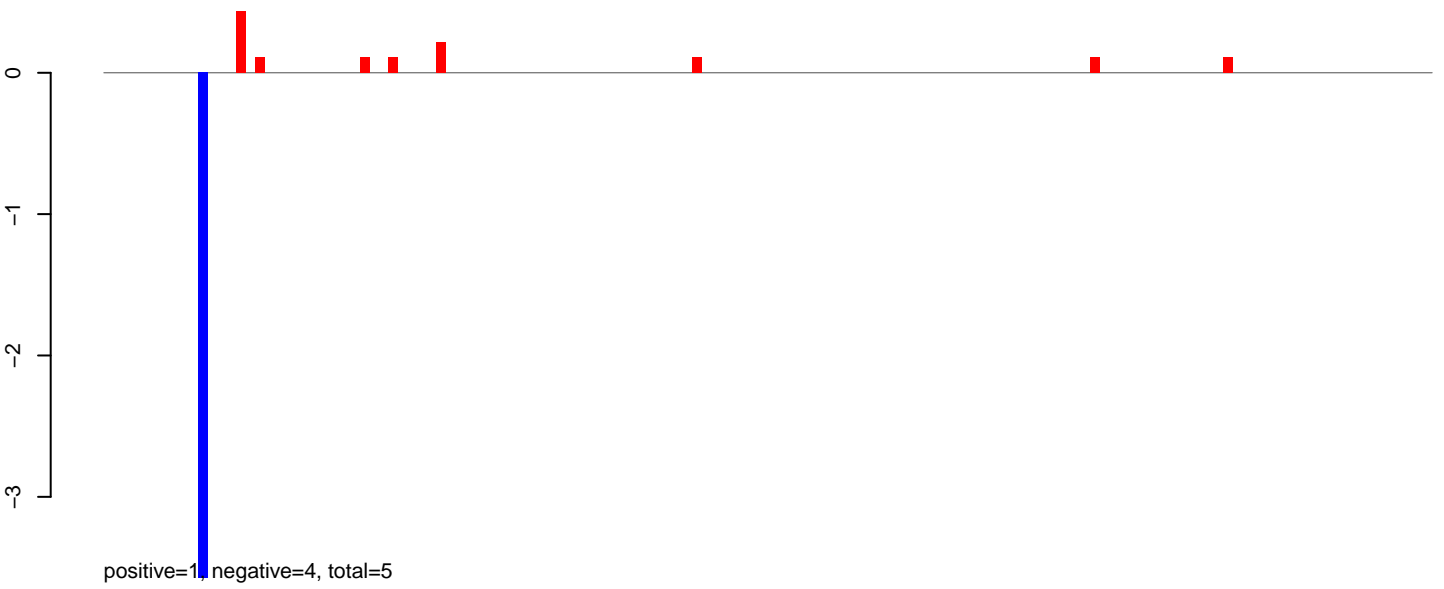
AnGam_Sua5b_cells.rep



positive=3, negative=205, total=208

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

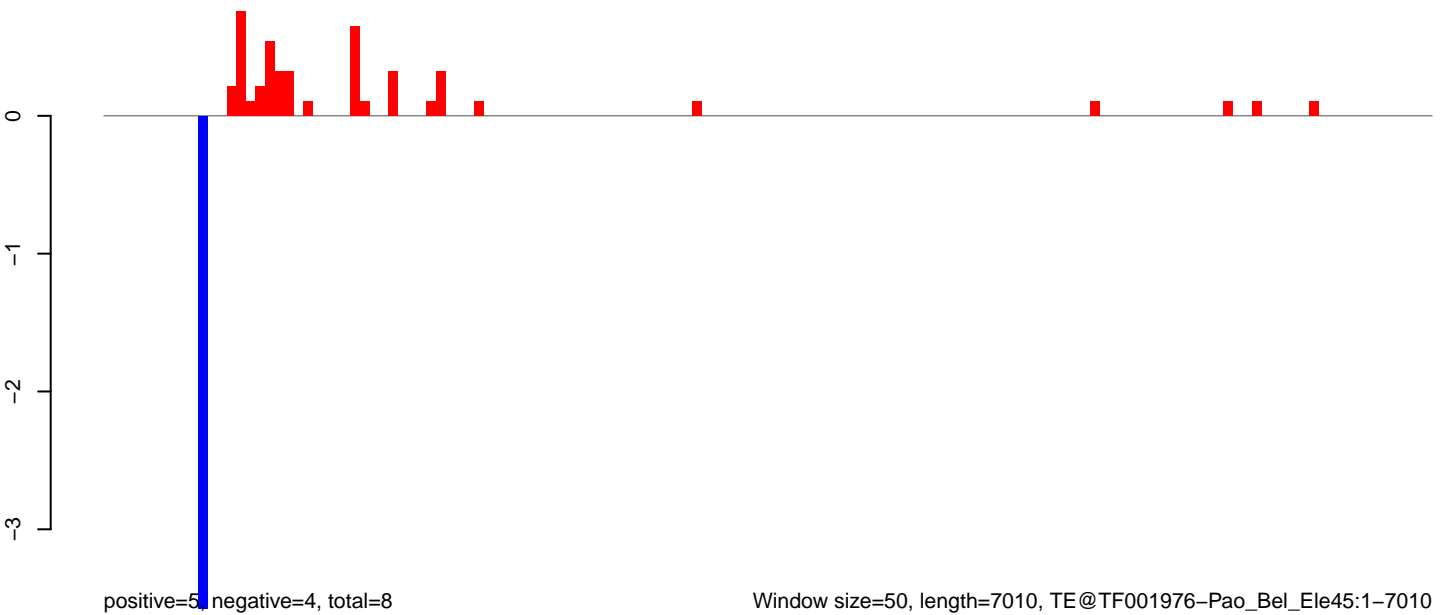
AnGam_Sua5b_cells.18_23.rep



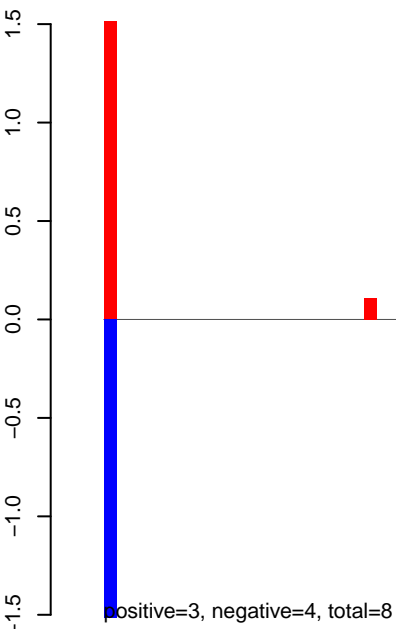
AnGam_Sua5b_cells.24_35.rep



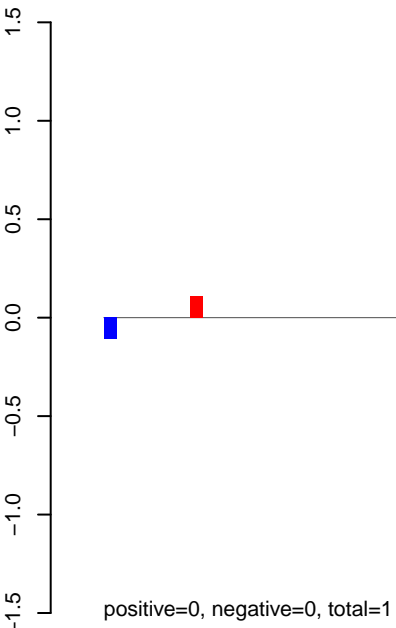
AnGam_Sua5b_cells.rep



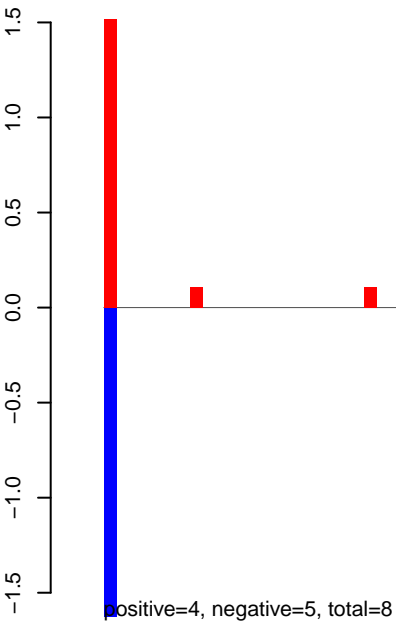
AnGam_Sua5b_cells.18_23.rep



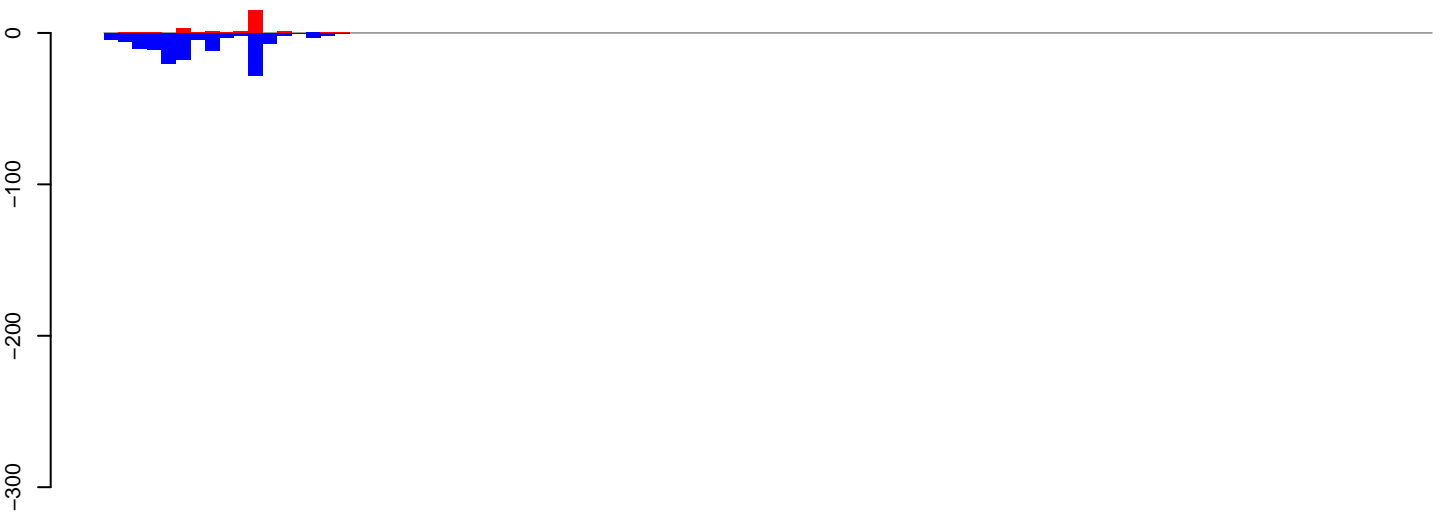
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

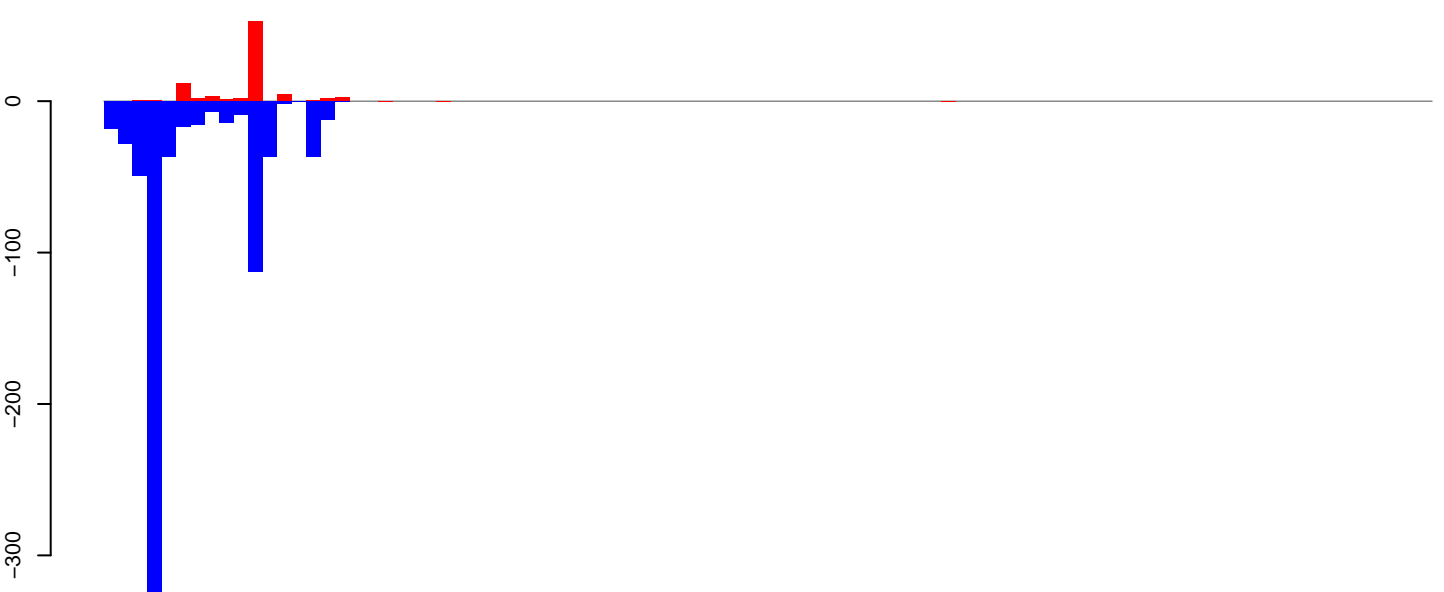


AnGam_Sua5b_cells.18_23.rep



positive=22, negative=133, total=155

AnGam_Sua5b_cells.24_35.rep



positive=83, negative=737, total=820

AnGam_Sua5b_cells.rep

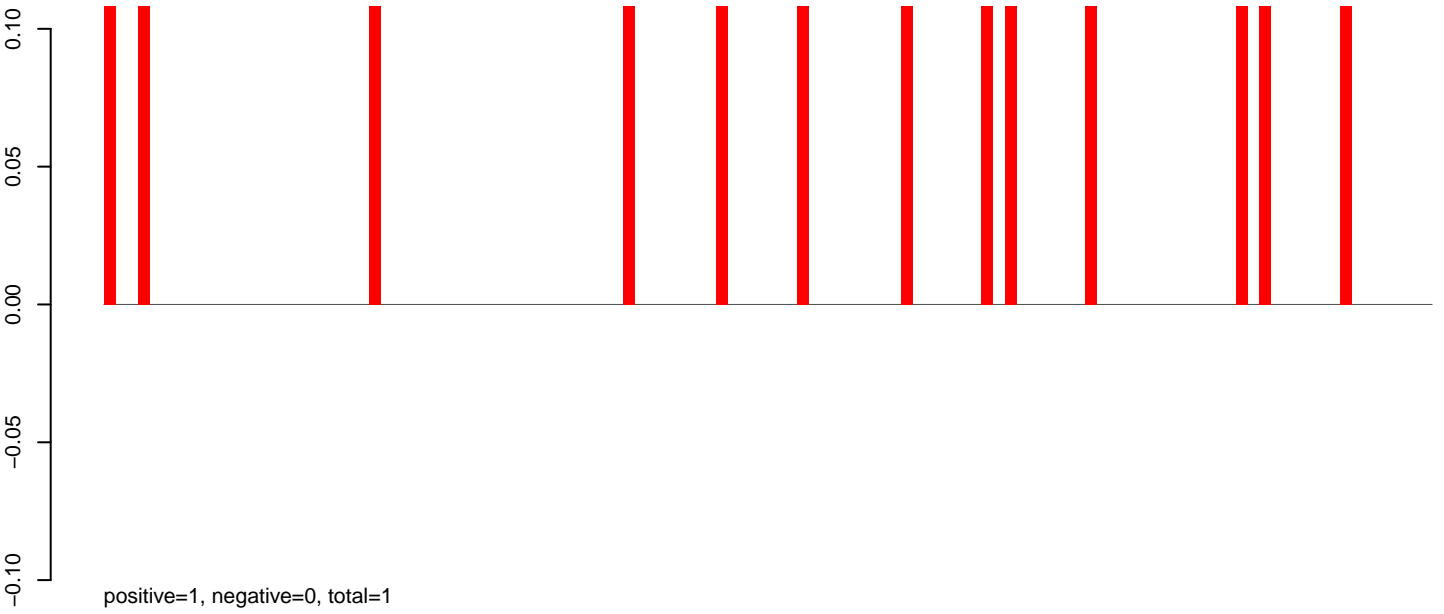


positive=106, negative=870, total=975

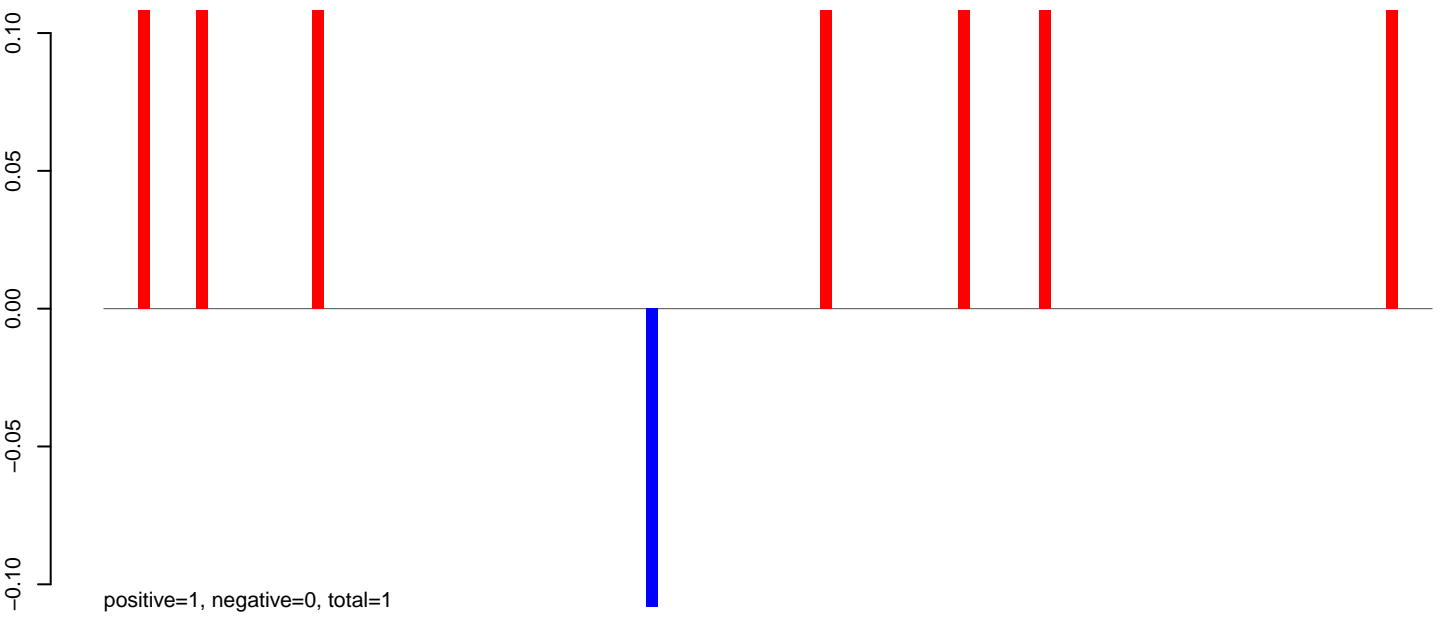
Window size=50, length=4641, TE@GYPSY35-LTR-I_AG:1-4641

0 1000 2000 3000 4000

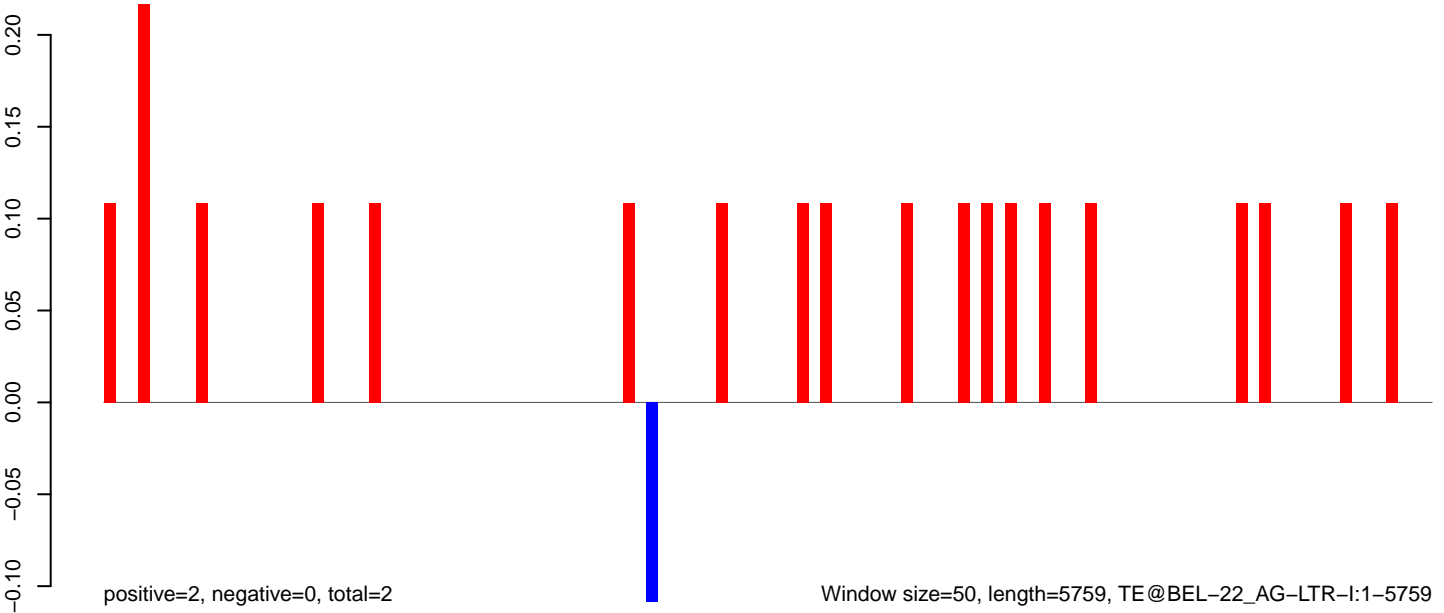
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



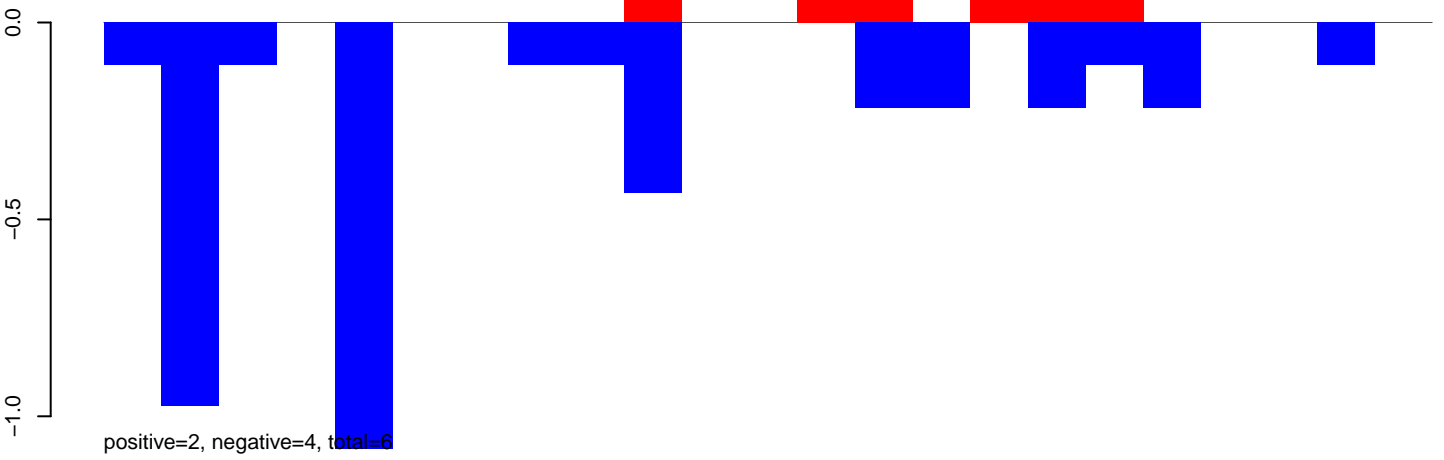
AnGam_Sua5b_cells.rep



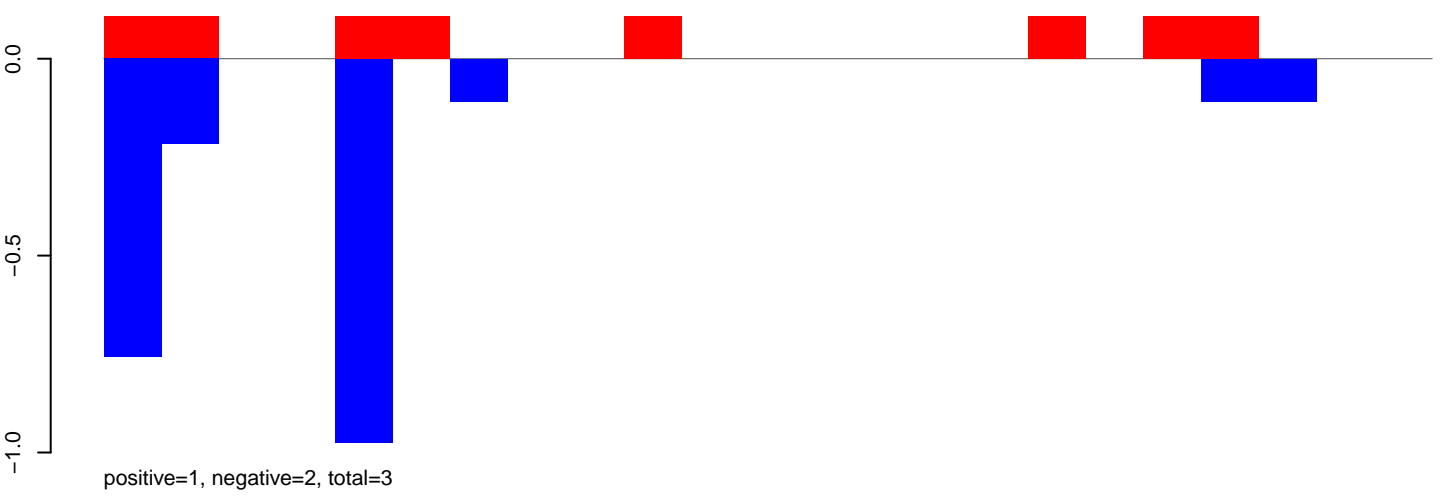
Window size=50, length=5759, TE@BEL-22_AG-LTR-I:1-5759

0 1000 2000 3000 4000 5000 6000

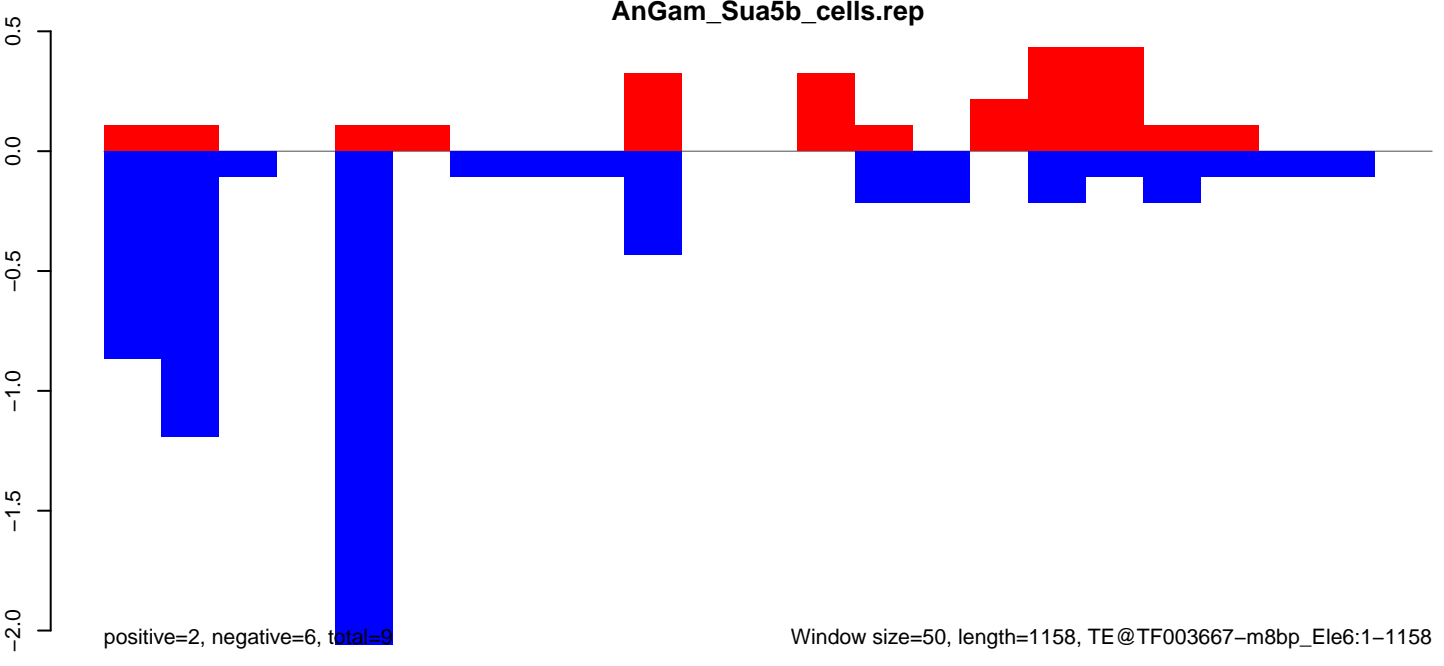
AnGam_Sua5b_cells.18_23.rep



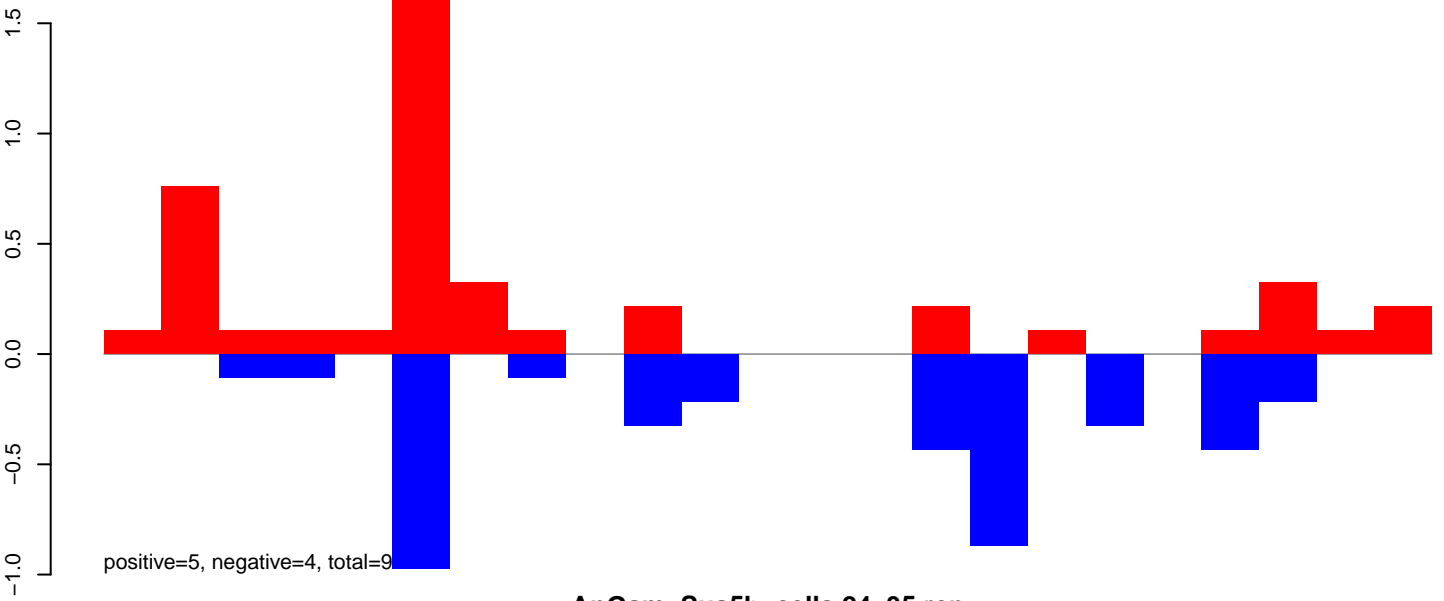
AnGam_Sua5b_cells.24_35.rep



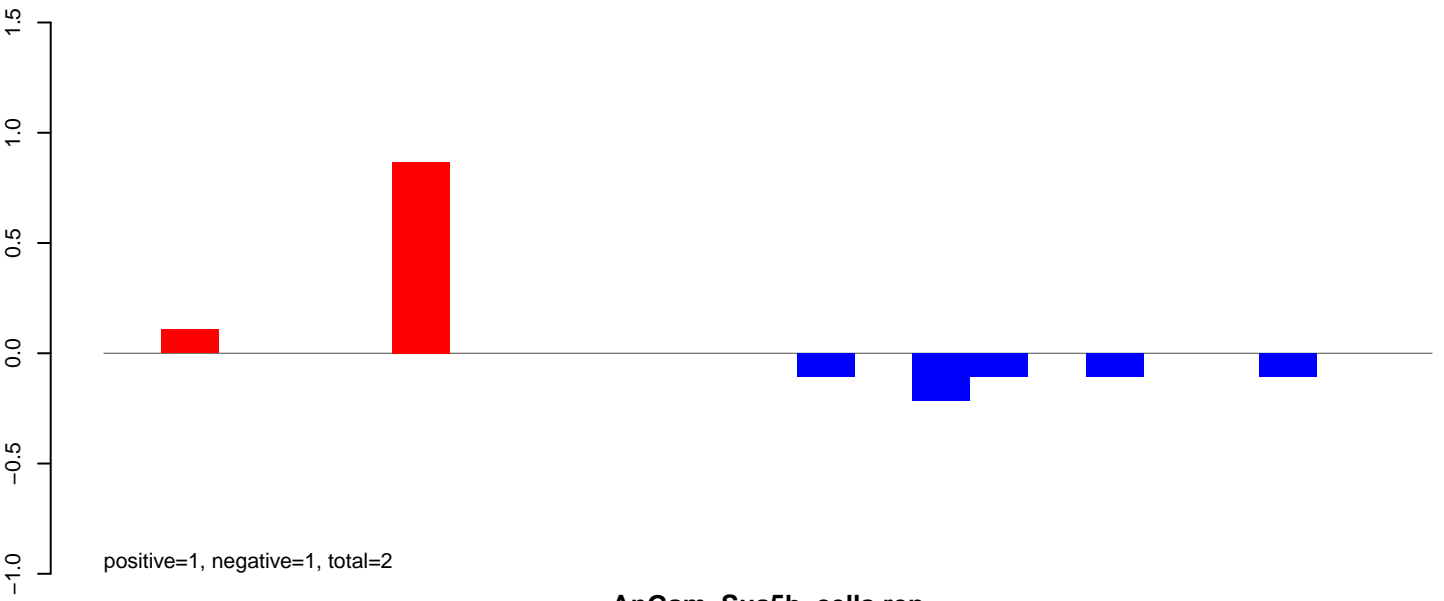
AnGam_Sua5b_cells.rep



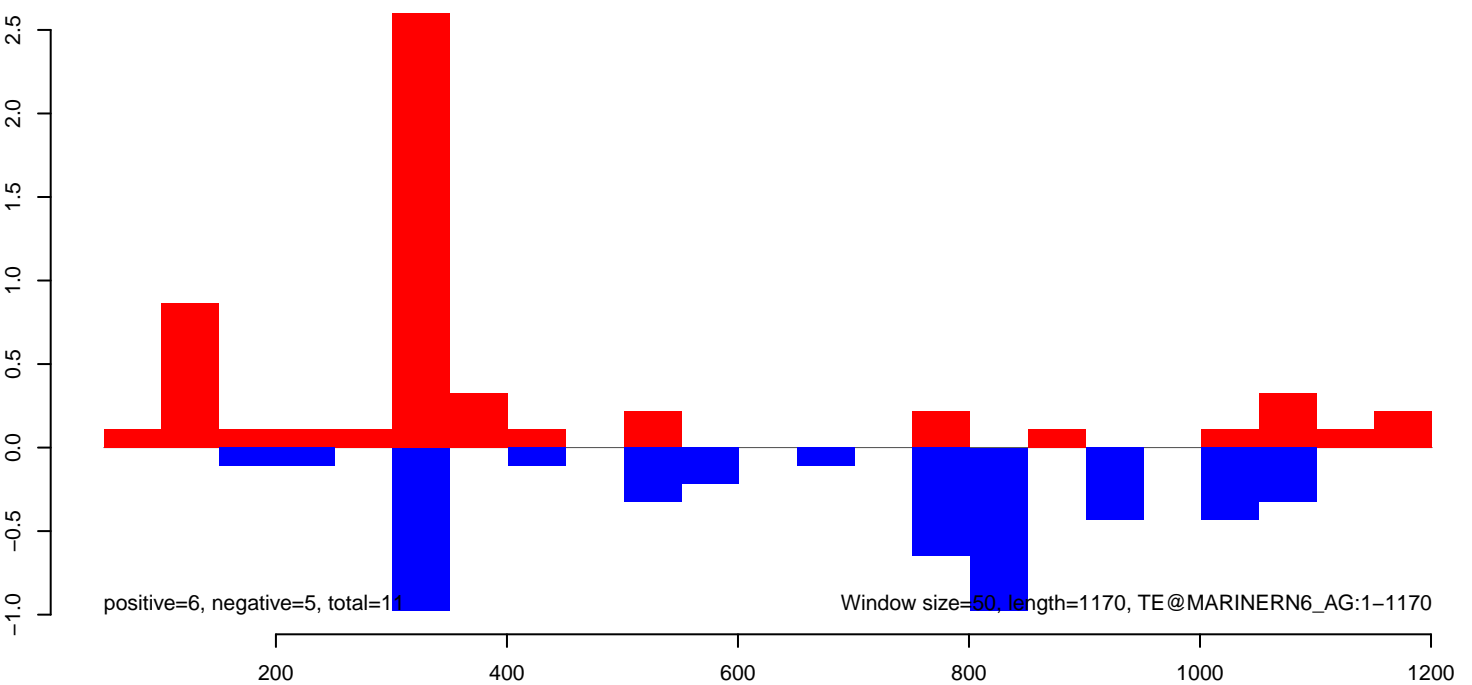
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

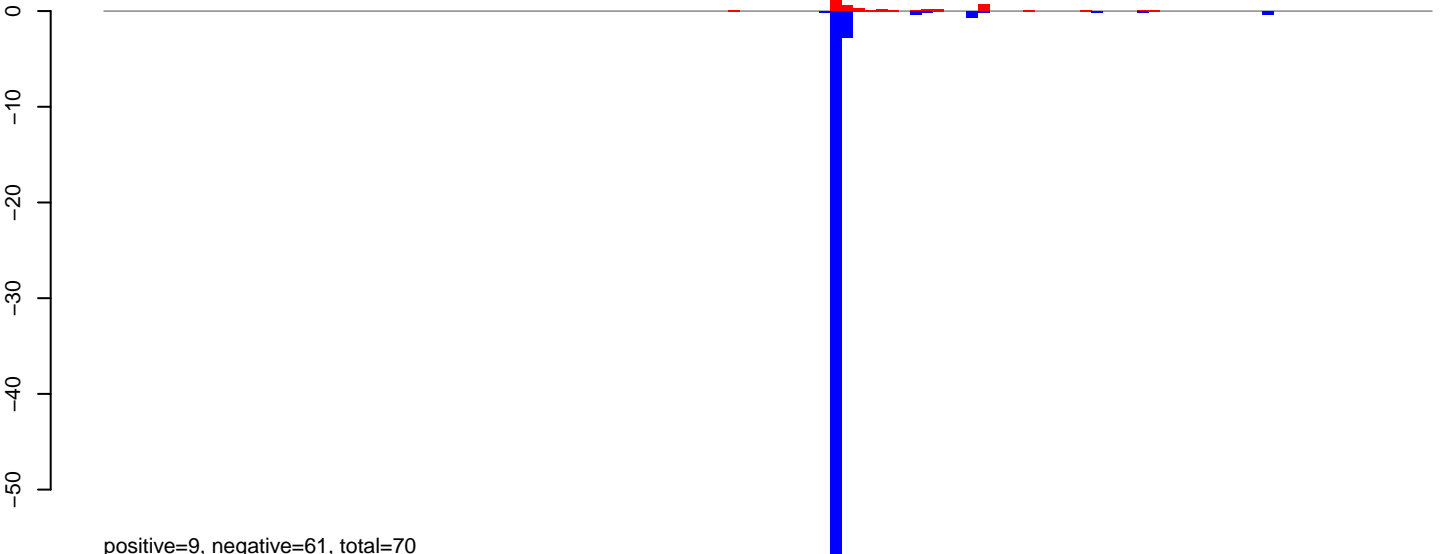


AnGam_Sua5b_cells.18_23.rep



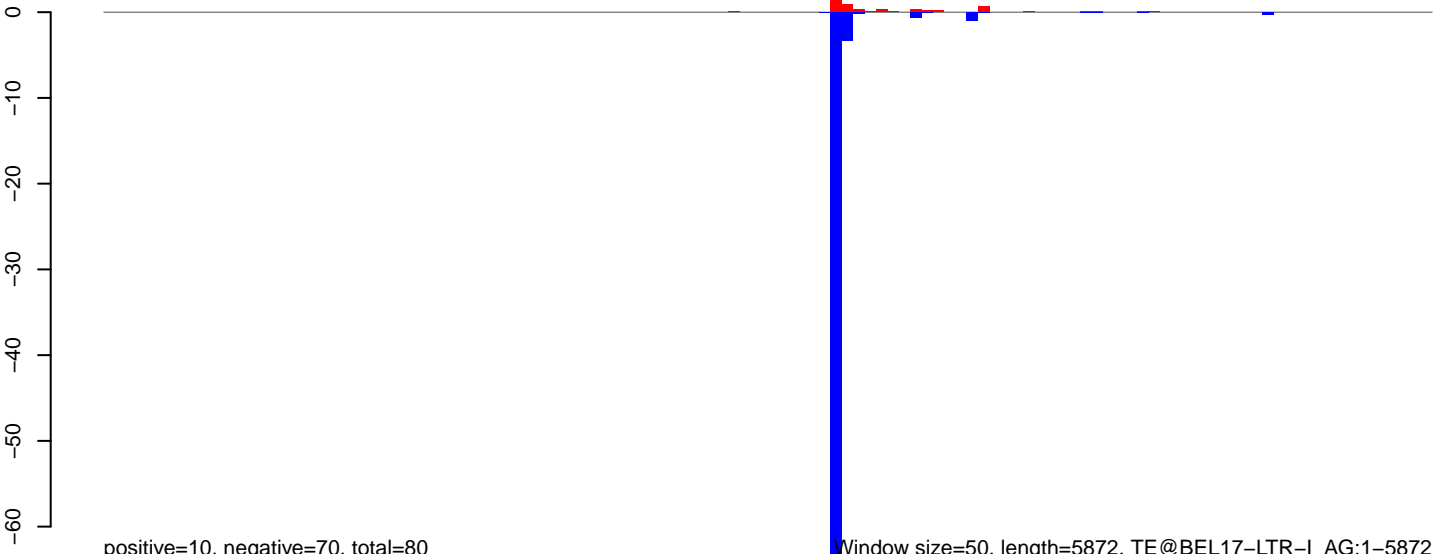
positive=1, negative=8, total=10

AnGam_Sua5b_cells.24_35.rep



positive=9, negative=61, total=70

AnGam_Sua5b_cells.rep

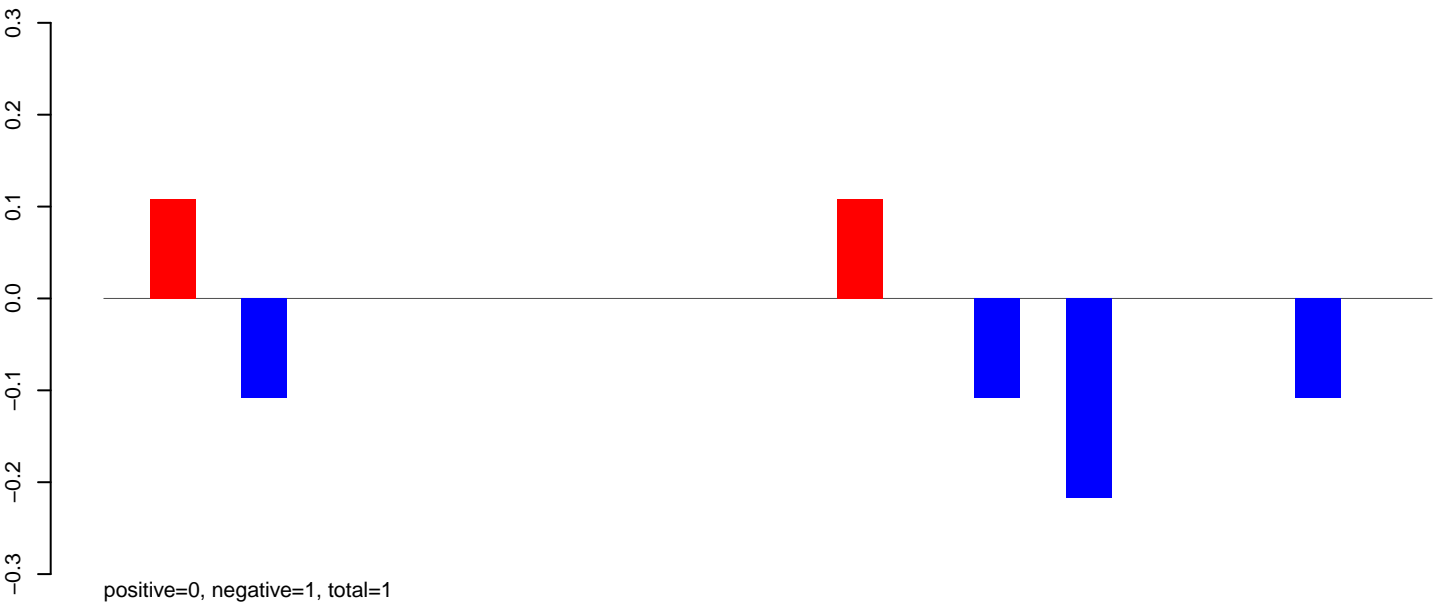


positive=10, negative=70, total=80

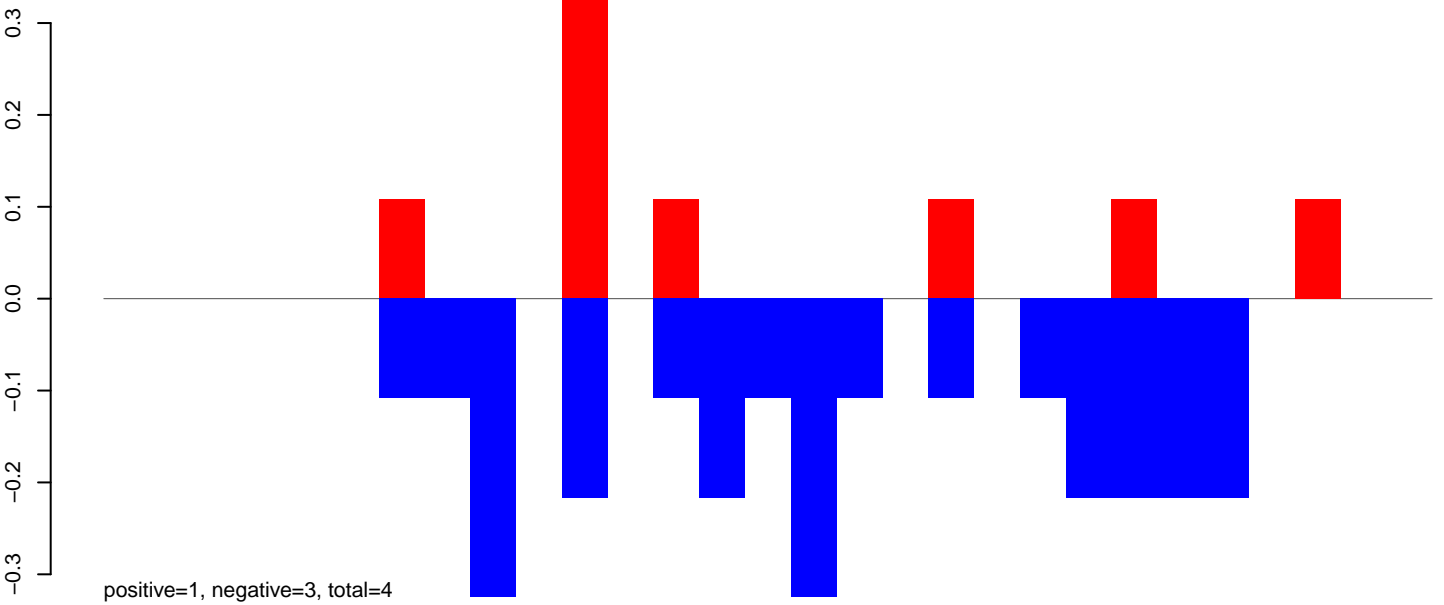
Window size=50, length=5872, TE@BEL17-LTR-I_AG:1-5872

0 1000 2000 3000 4000 5000 6000

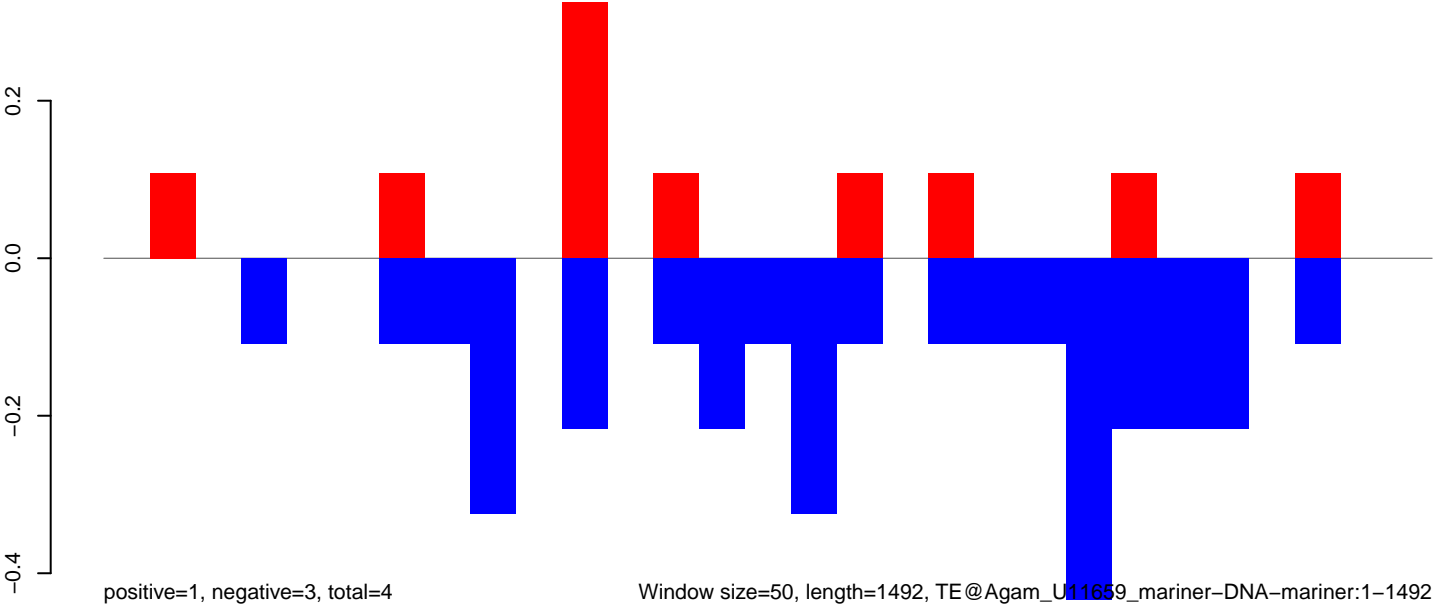
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



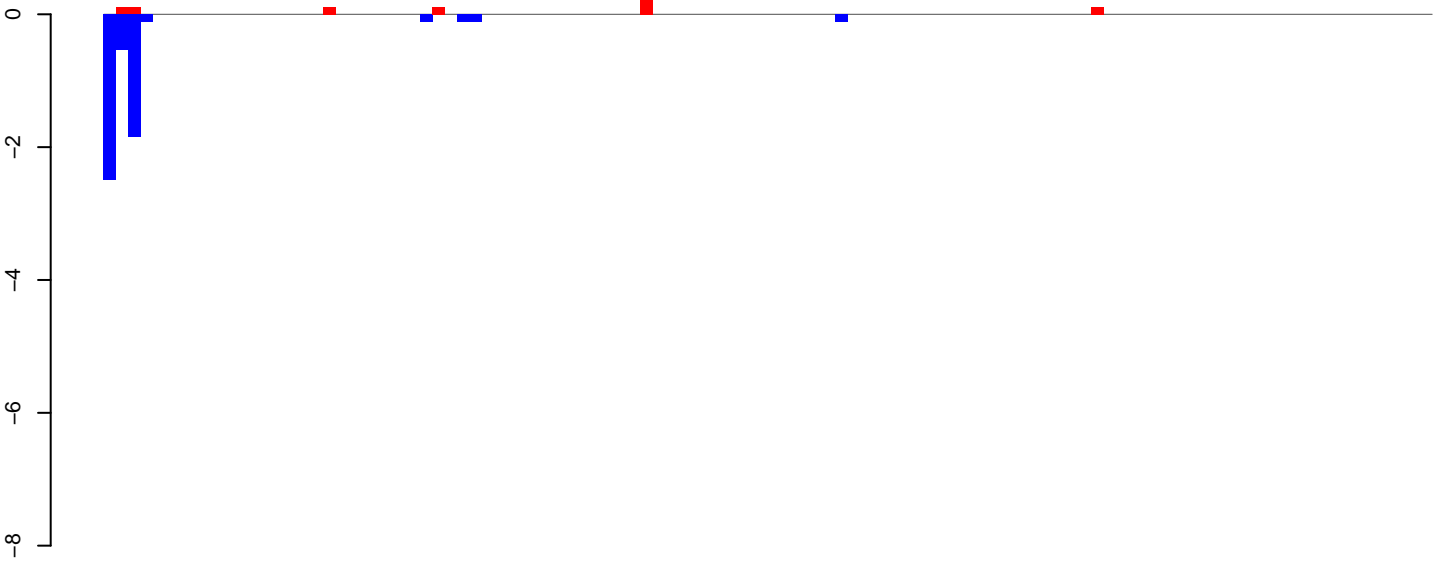
AnGam_Sua5b_cells.rep



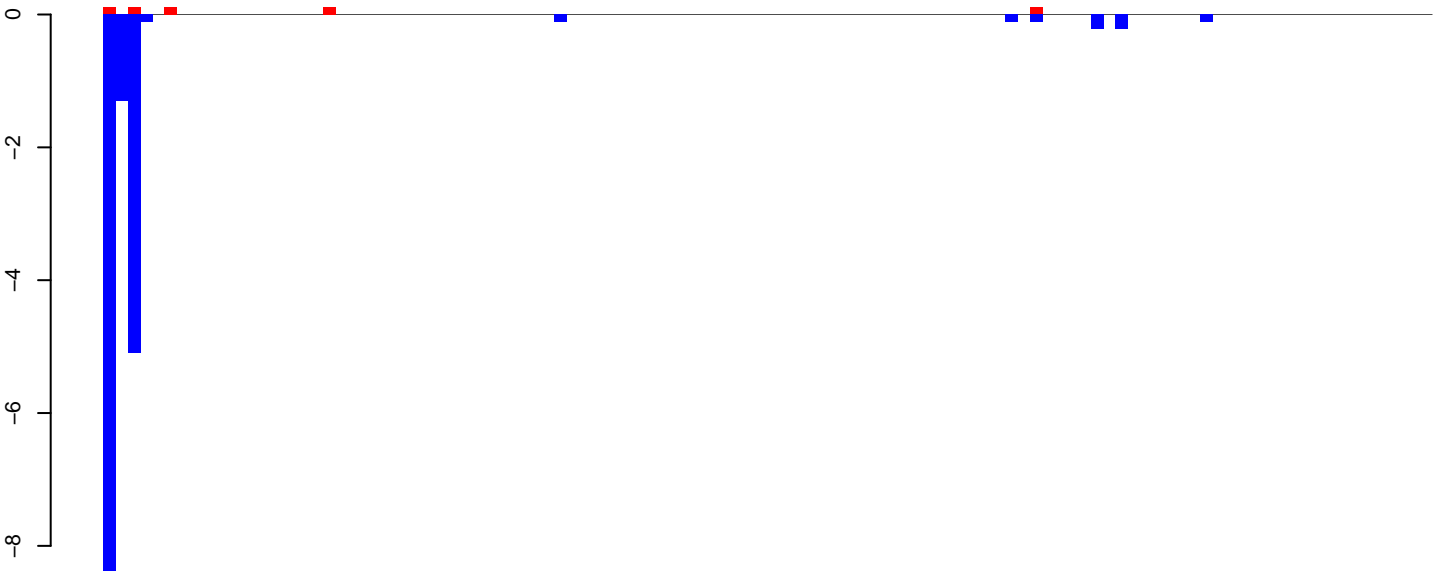
Window size=50, length=1492, TE@Agam_U11659_mariner-DNA-mariner:1-1492

0 500 1000 1500

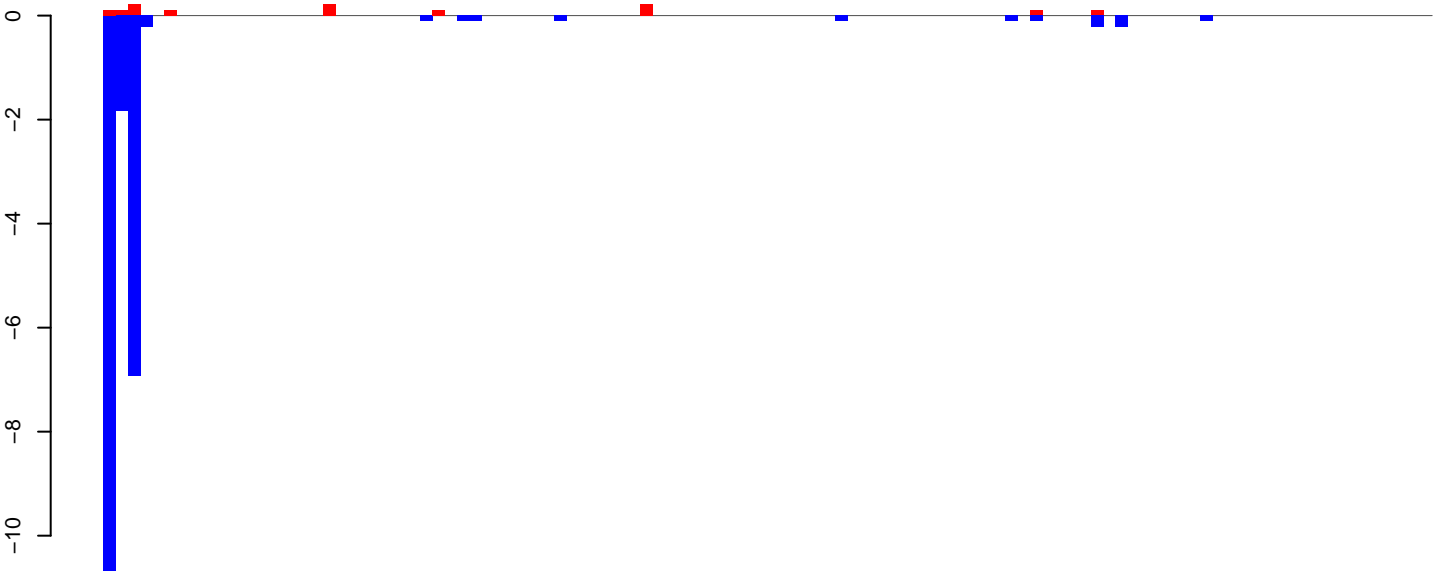
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



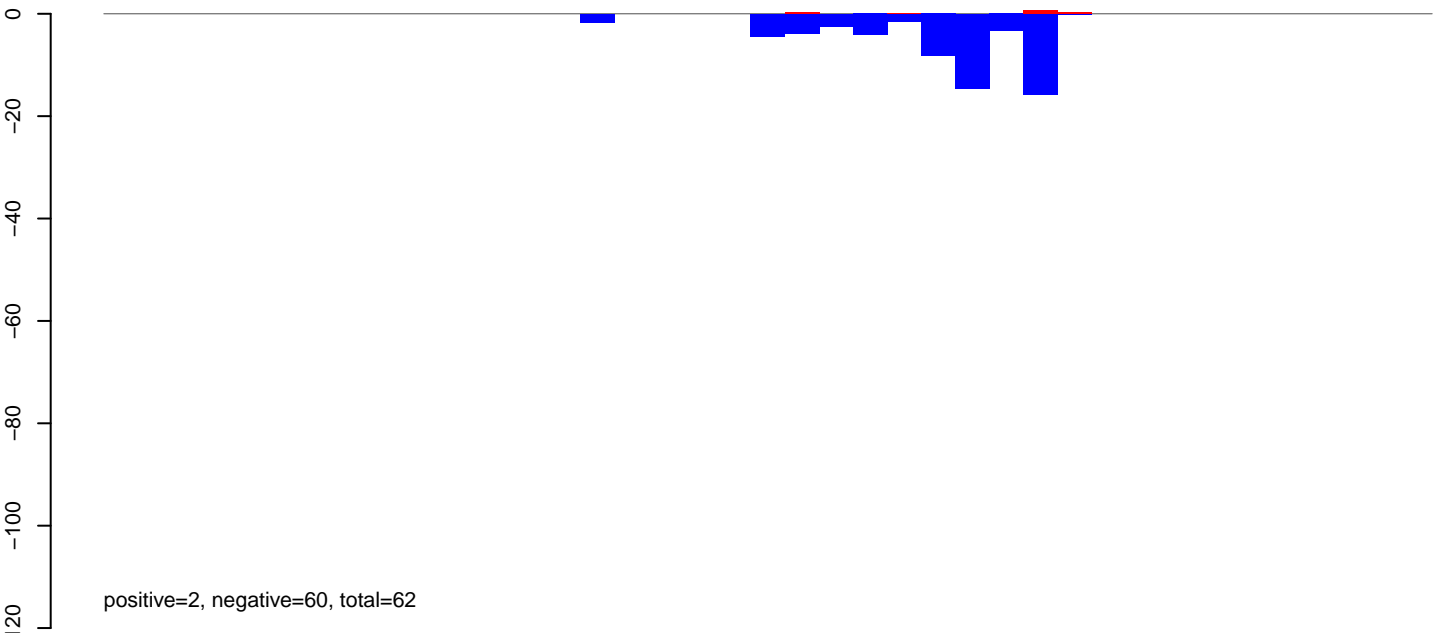
AnGam_Sua5b_cells.rep



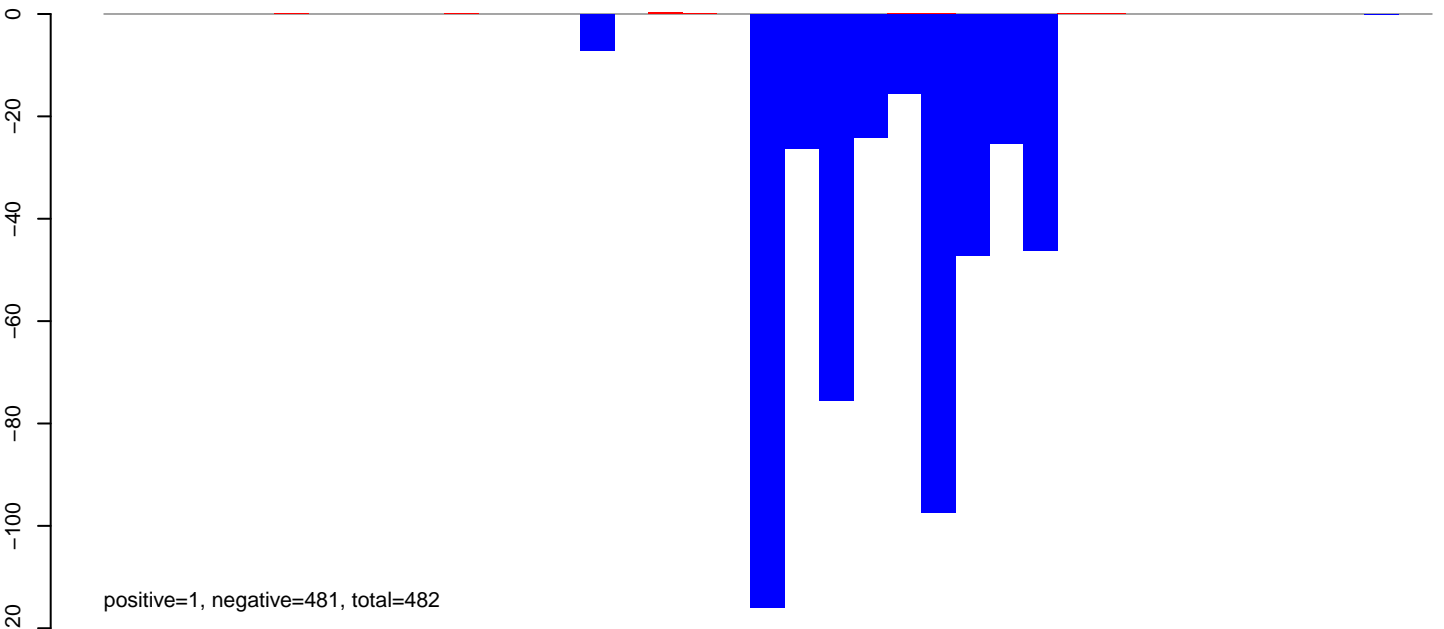
Window size=50, length=5492, TE@BEL5-LTR-I_AG:1-5492

0 1000 2000 3000 4000 5000

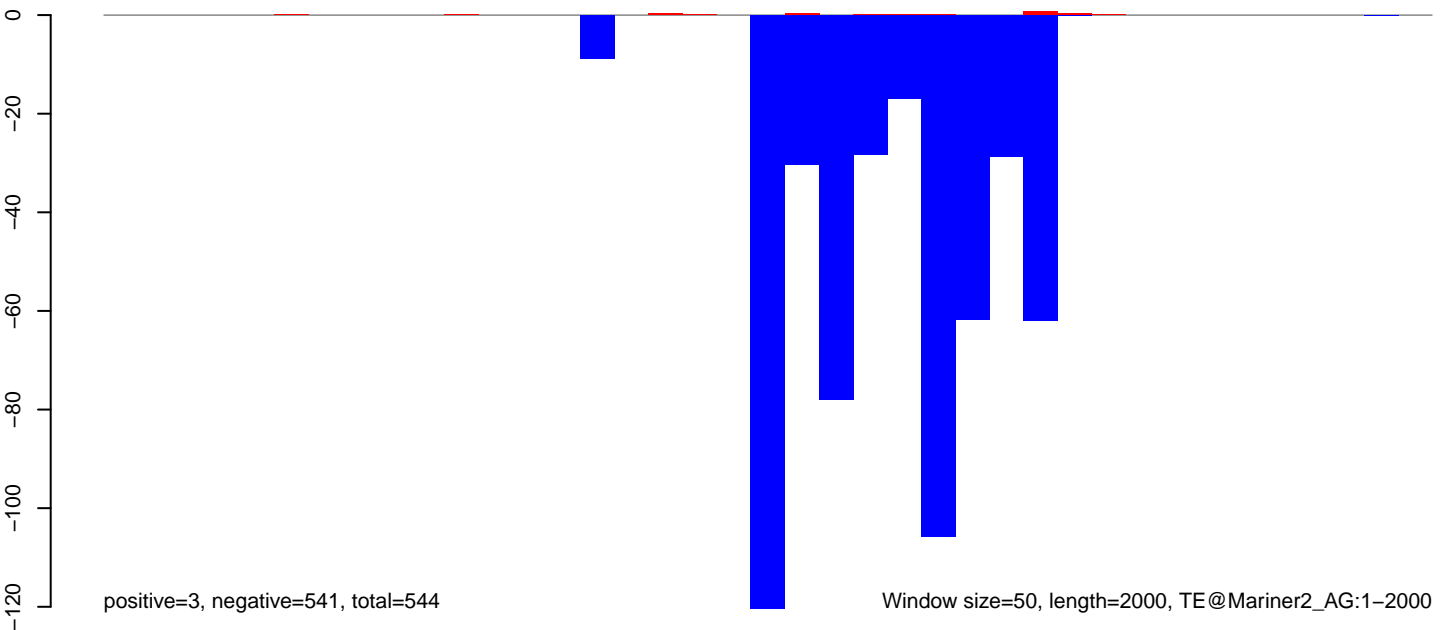
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

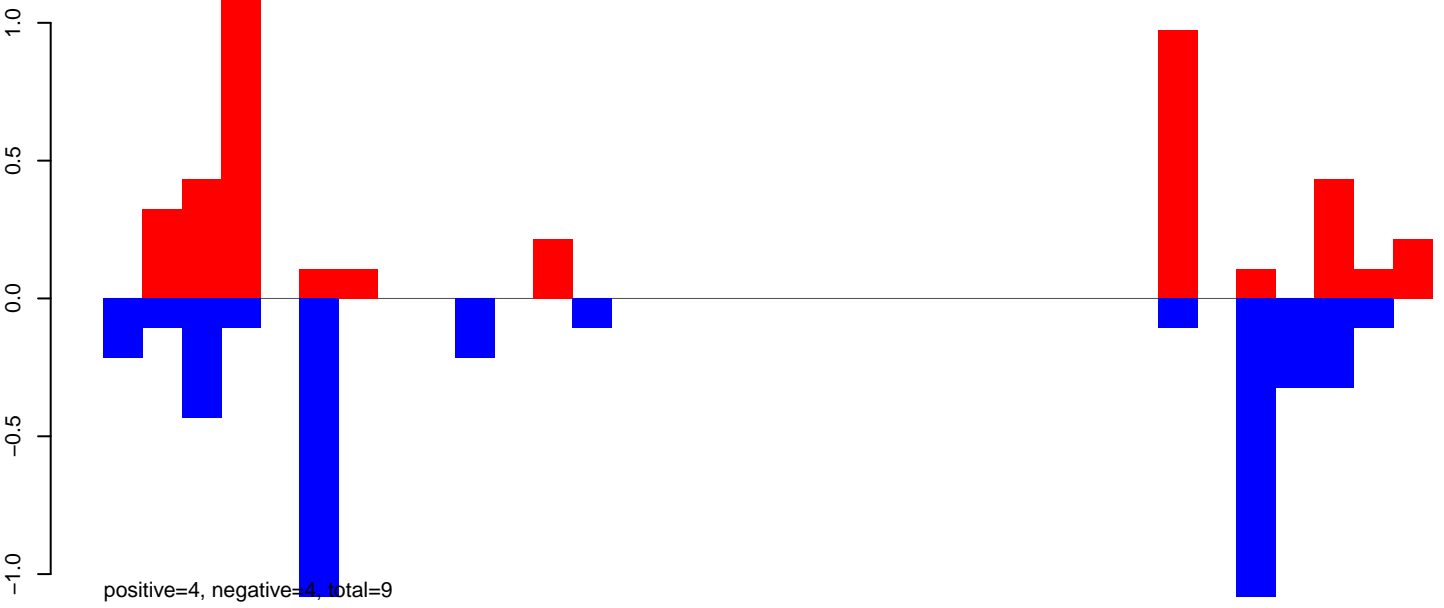


AnGam_Sua5b_cells.rep

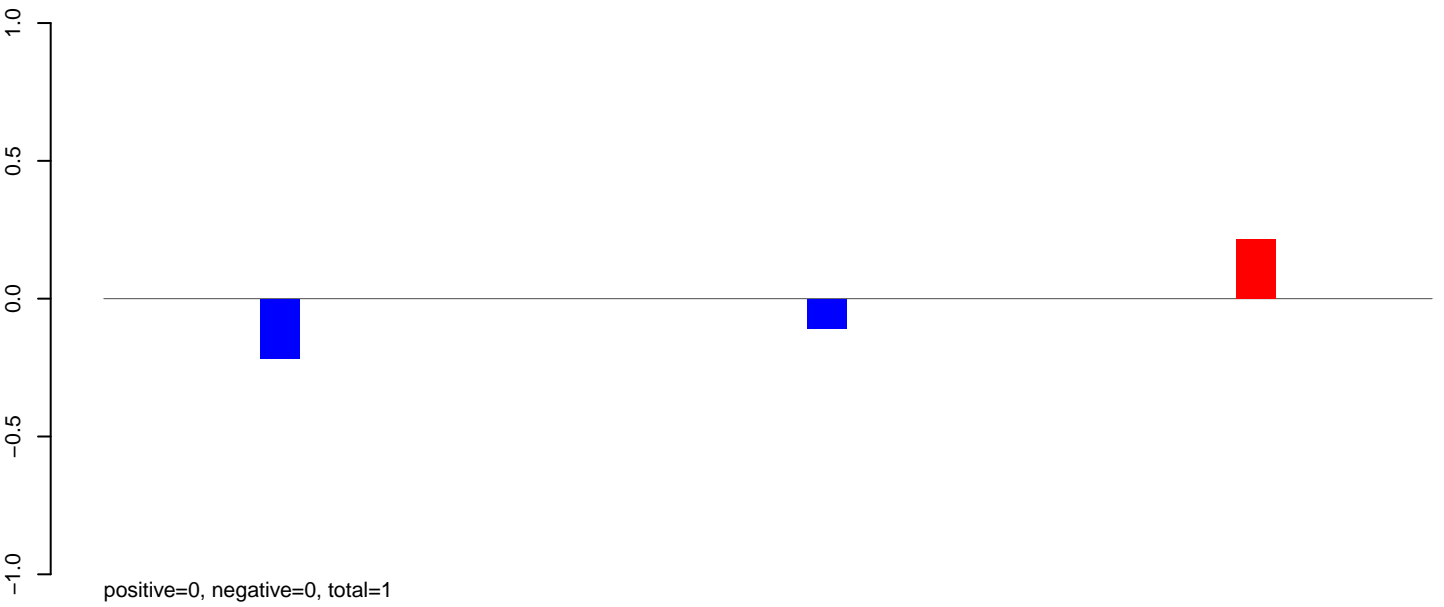


0 500 1000 1500 2000

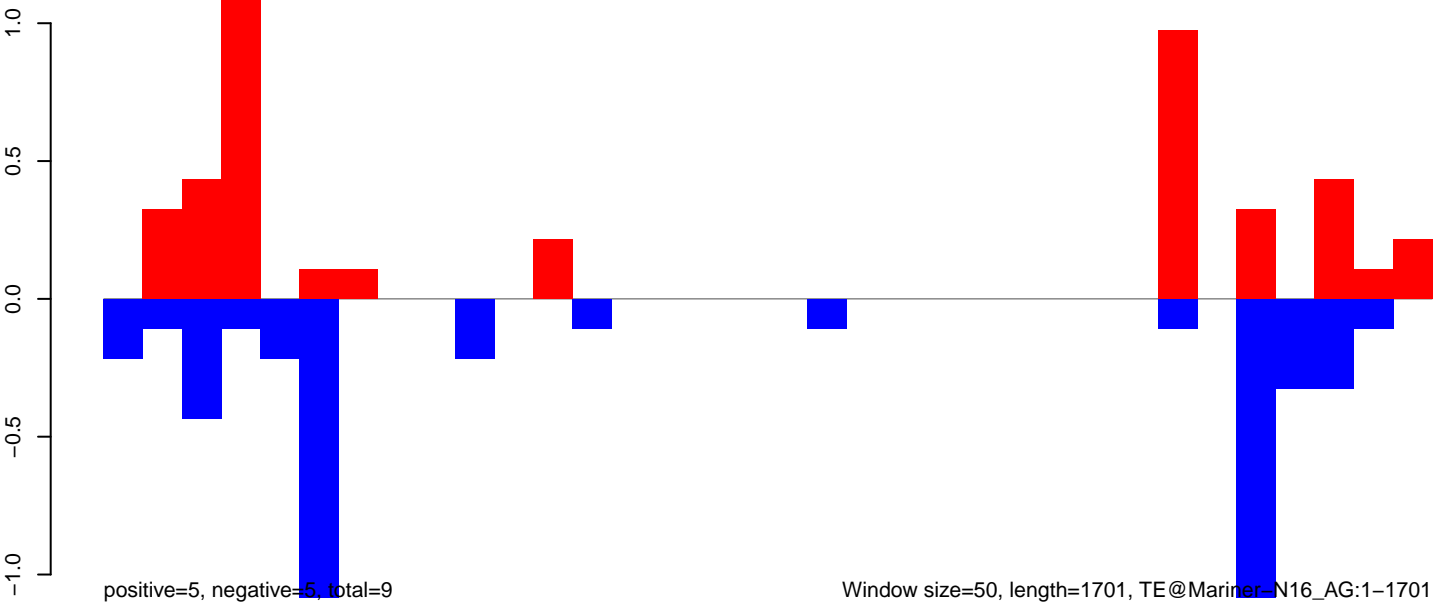
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

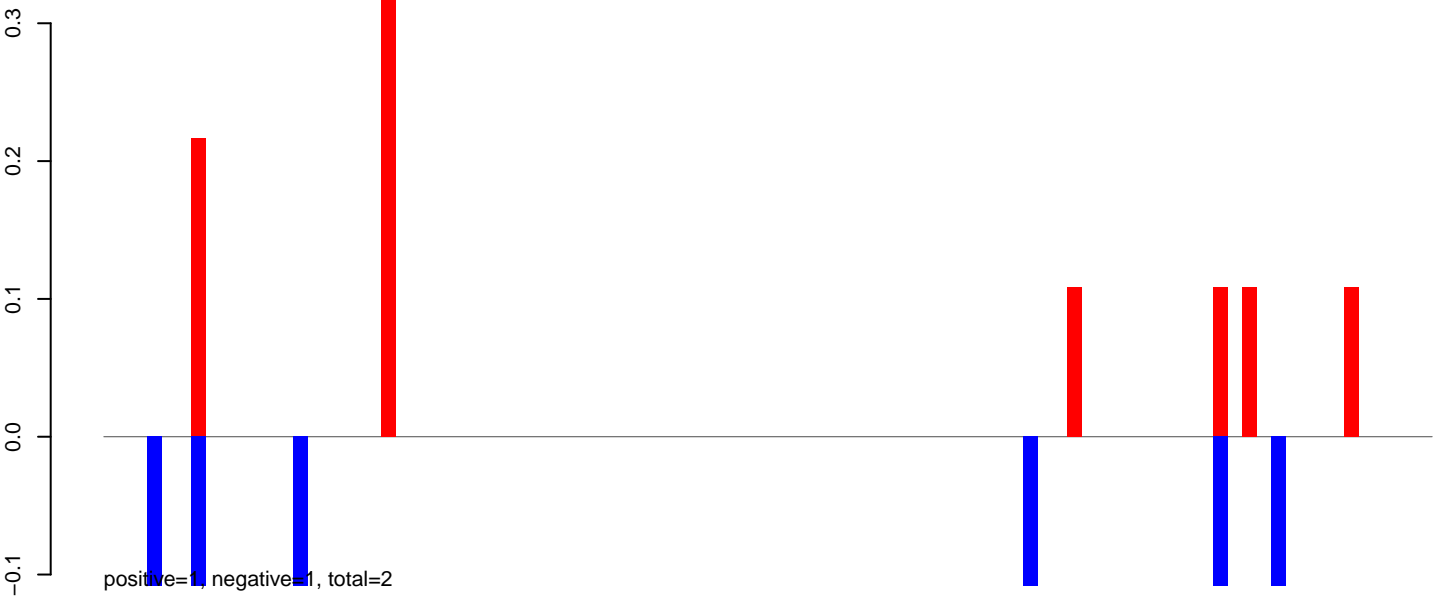


AnGam_Sua5b_cells.rep

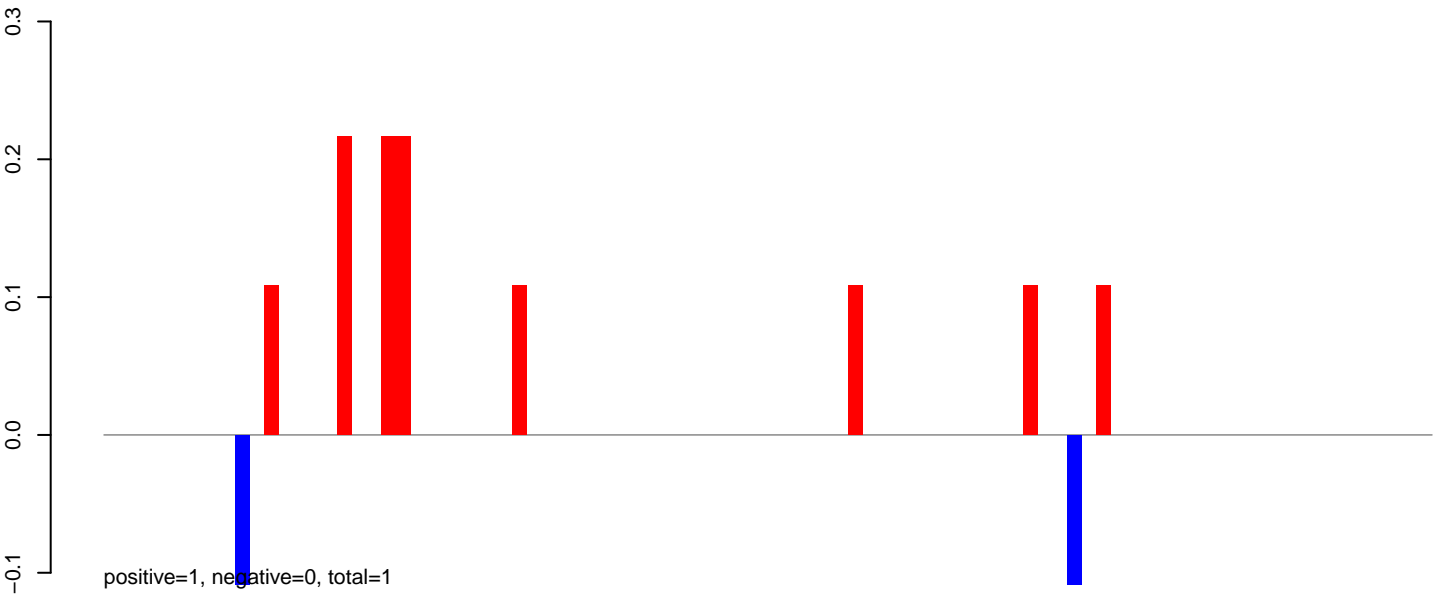


Window size=50, length=1701, TE@Mariner-N16_AG:1-1701

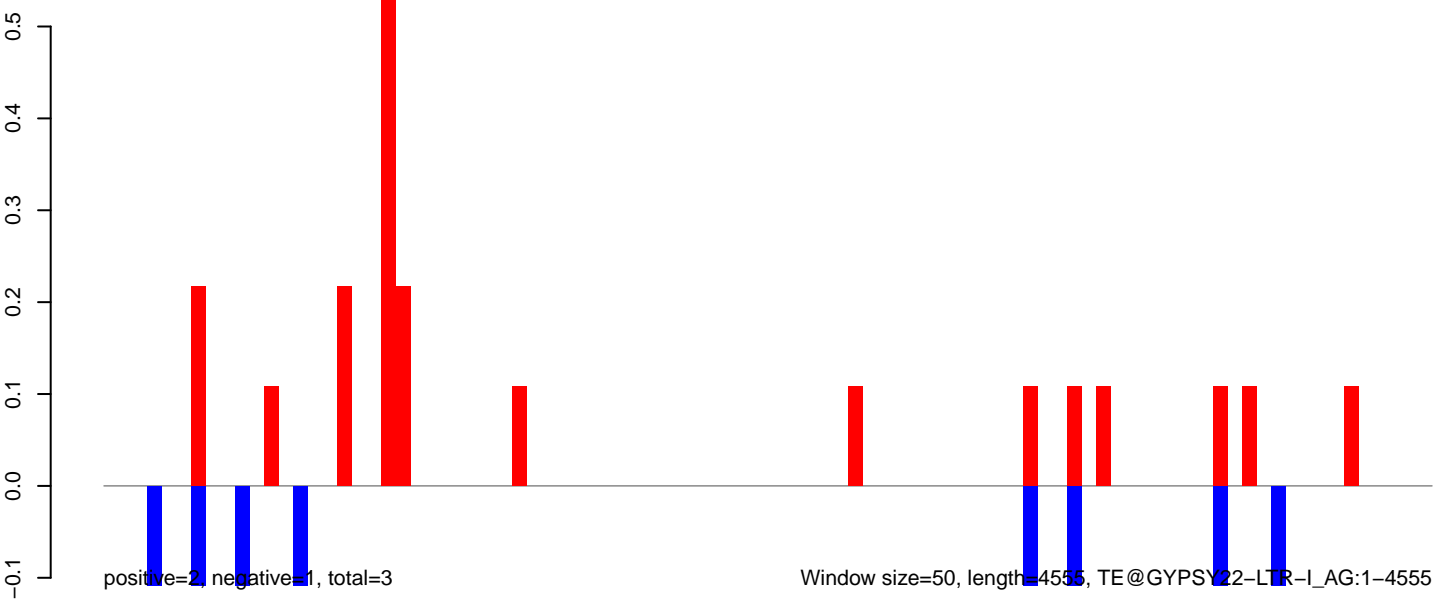
AnGam_Sua5b_cells.18_23.rep



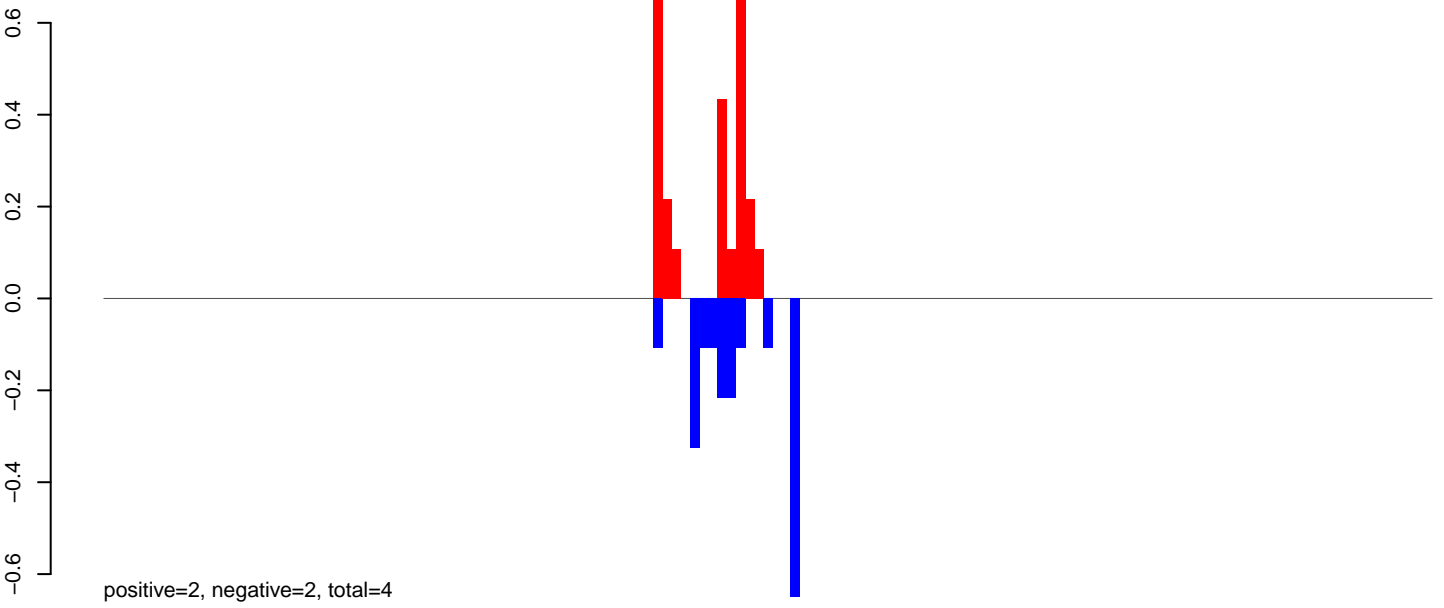
AnGam_Sua5b_cells.24_35.rep



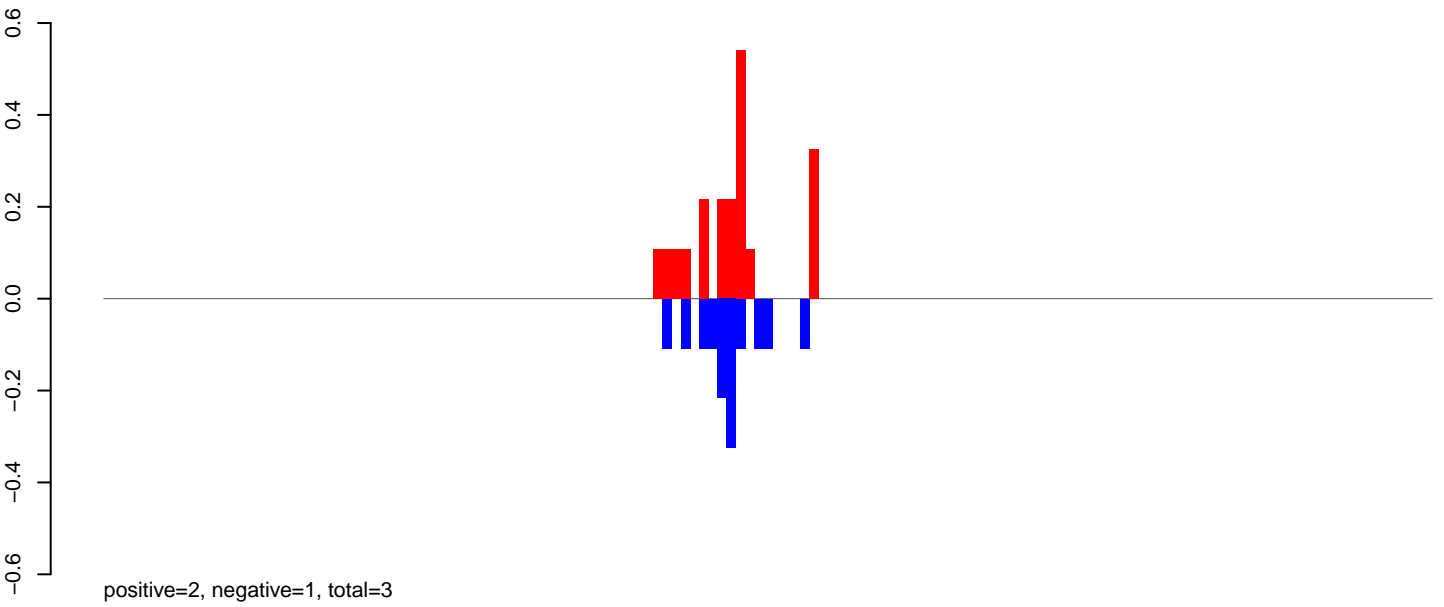
AnGam_Sua5b_cells.rep



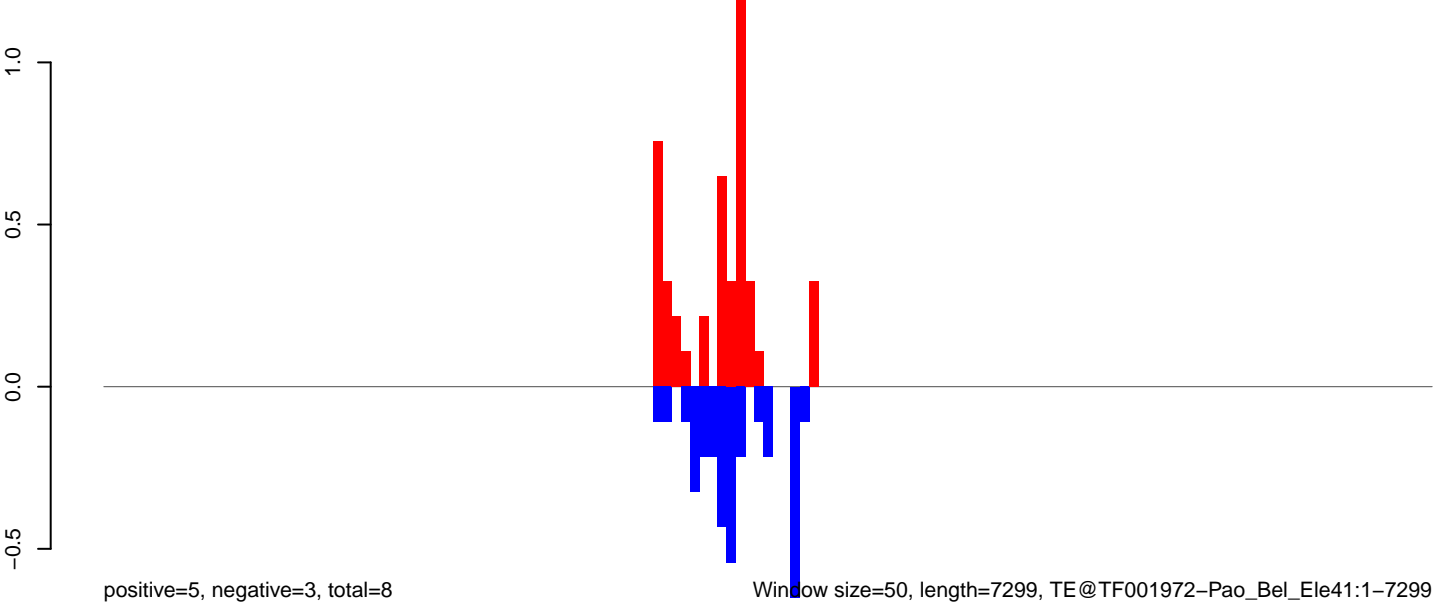
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



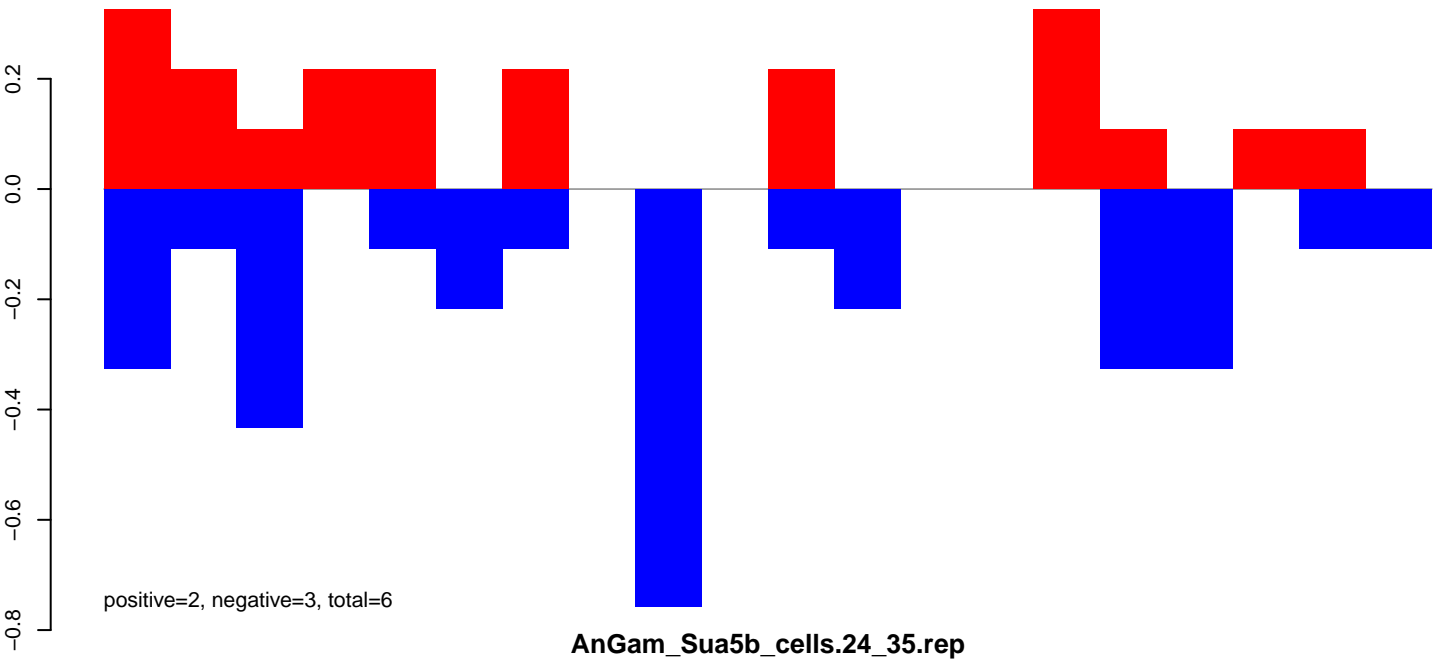
AnGam_Sua5b_cells.rep



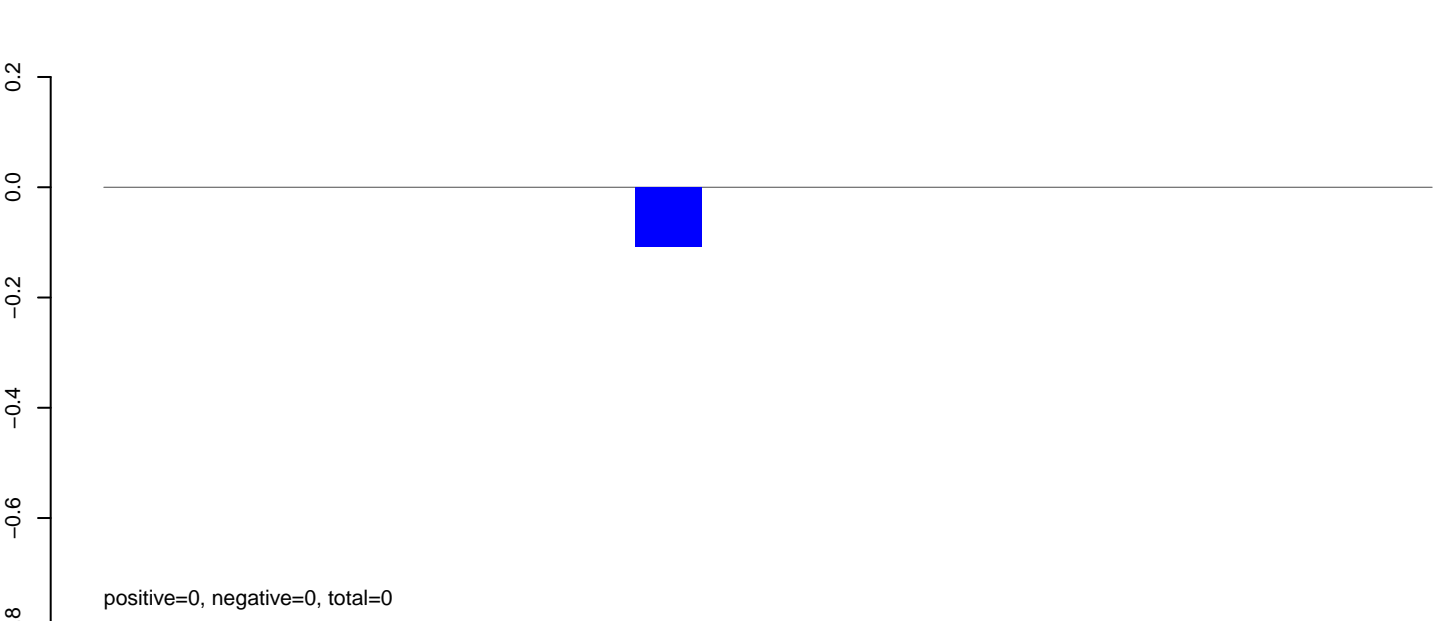
Window size=50, length=7299, TE@TF001972-Pao_Bel_Ele41:1-7299

0 2000 4000 6000

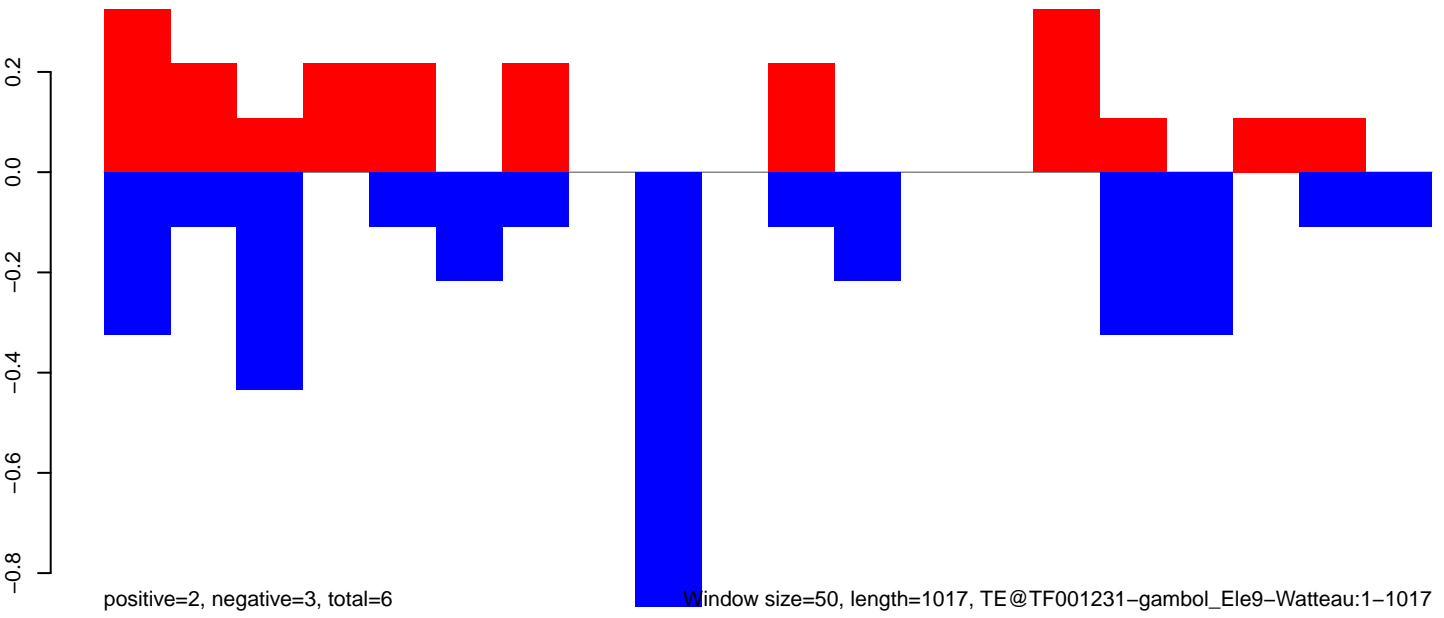
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

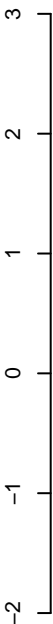


AnGam_Sua5b_cells.rep



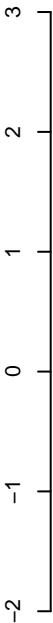
Window size=50, length=1017, TE@TF001231-gambol_Ele9-Watteau:1-1017

AnGam_Sua5b_cells.18_23.rep



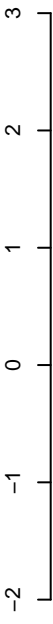
positive=8, negative=7, total=15

AnGam_Sua5b_cells.24_35.rep



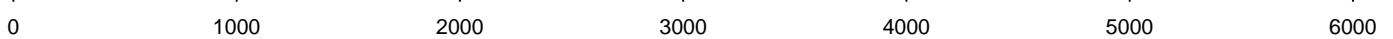
positive=0, negative=1, total=1

AnGam_Sua5b_cells.rep



positive=8, negative=8, total=16

Window size=50, length=5956, TE@TF001224-gambol_Ele2-Ozzie:1-5956

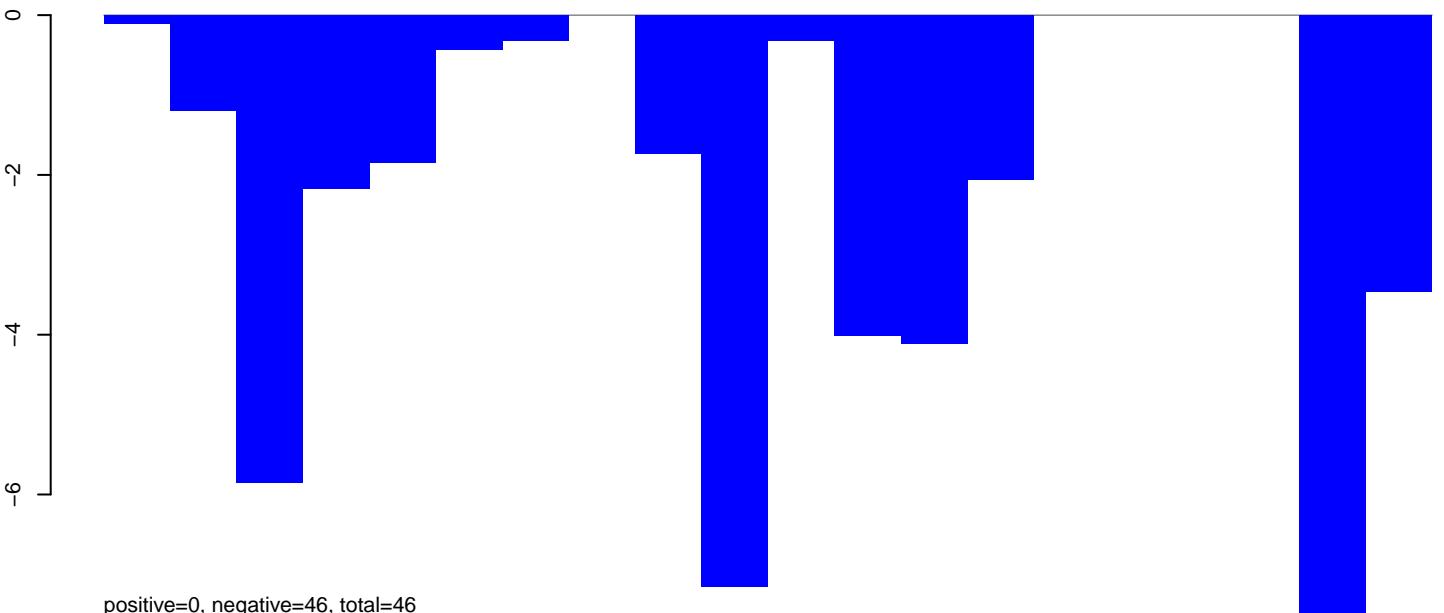


AnGam_Sua5b_cells.18_23.rep



positive=0, negative=8, total=8

AnGam_Sua5b_cells.24_35.rep



positive=0, negative=46, total=46

AnGam_Sua5b_cells.rep

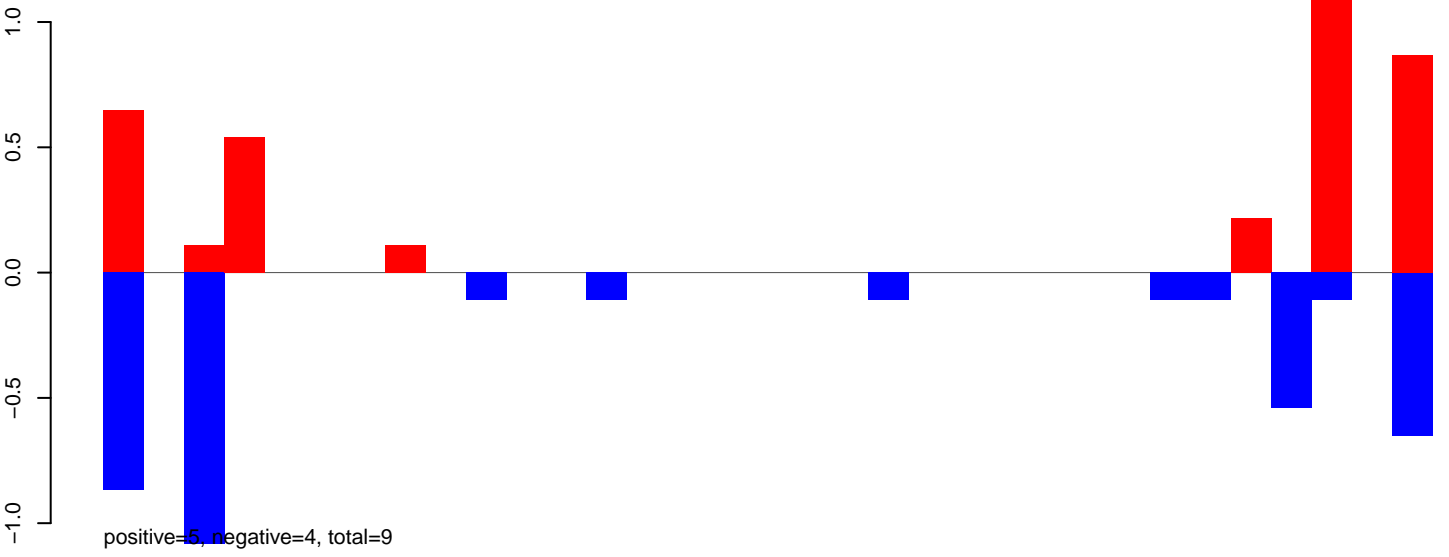


positive=0, negative=53, total=53

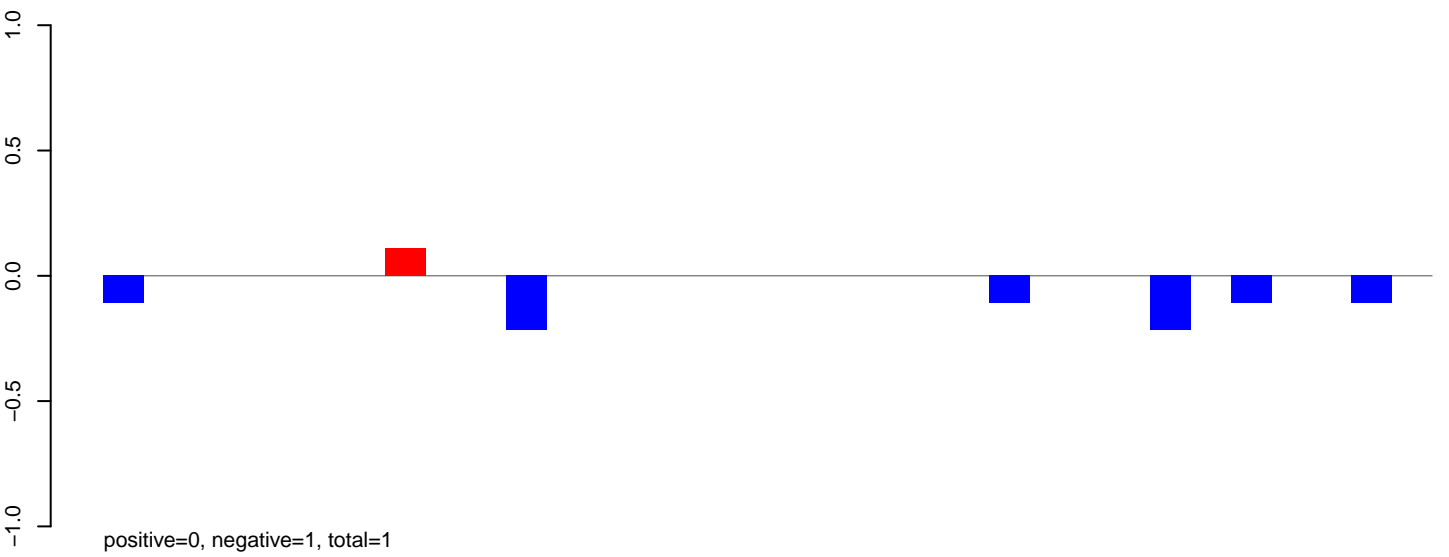
Window size=50, length=1014, TE@EAA10695-Unknown:1-1014

200 400 600 800 1000

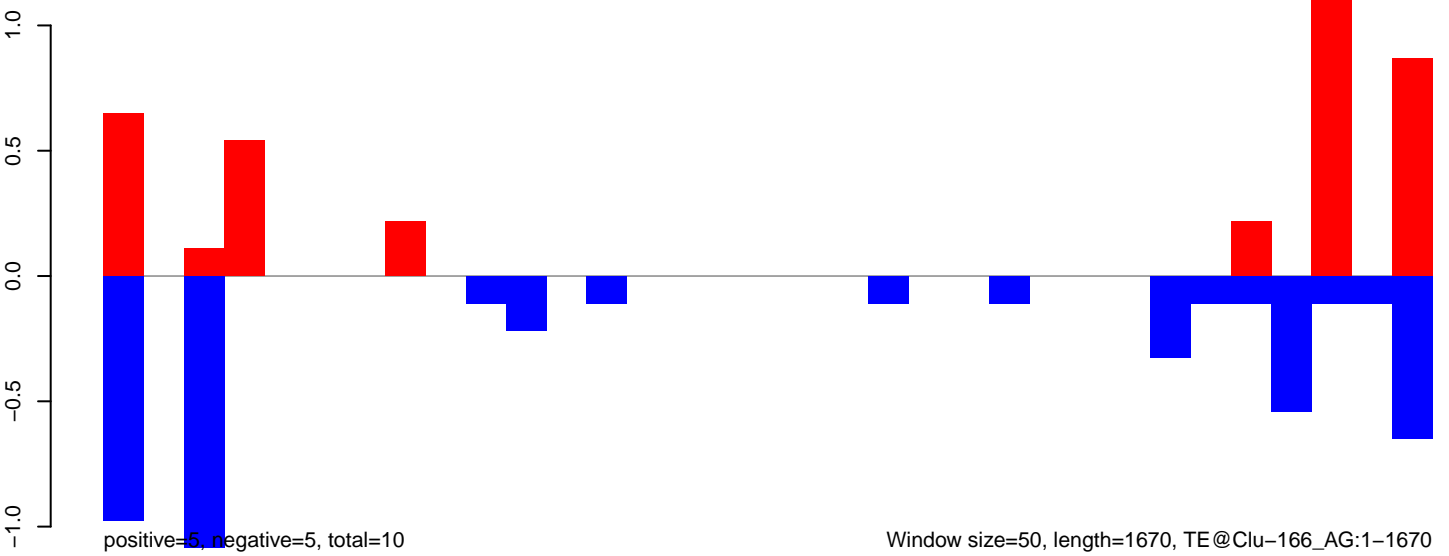
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

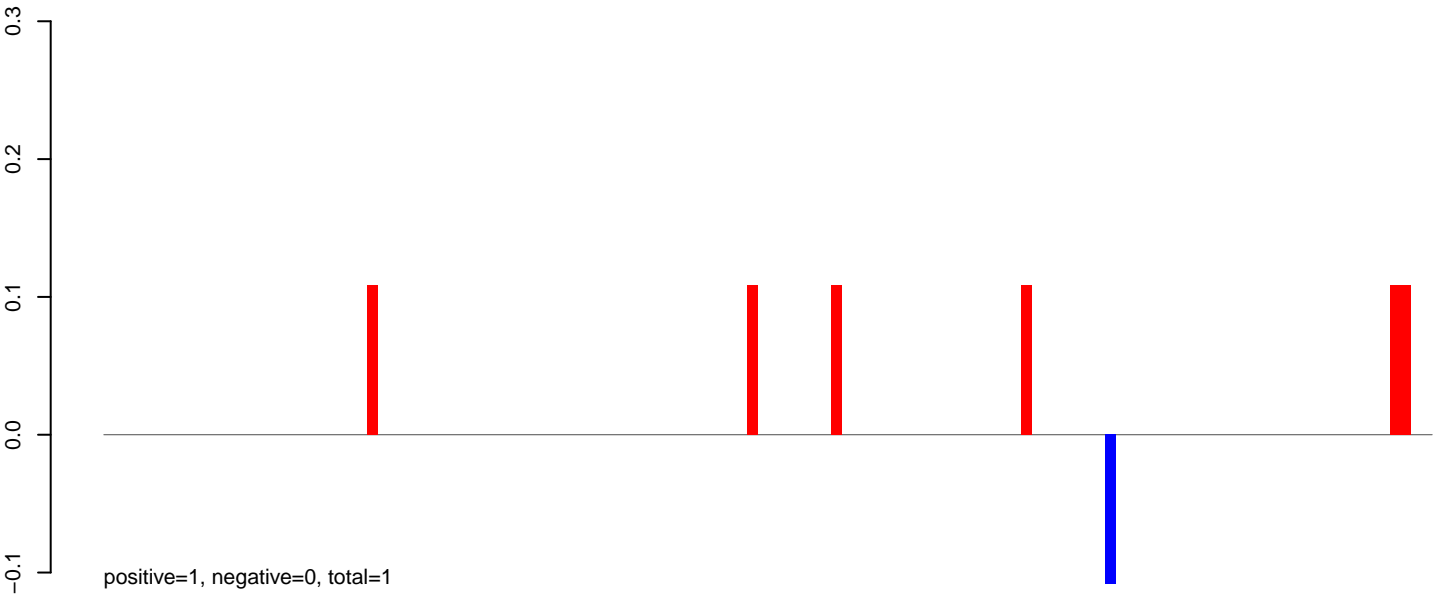


AnGam_Sua5b_cells.rep

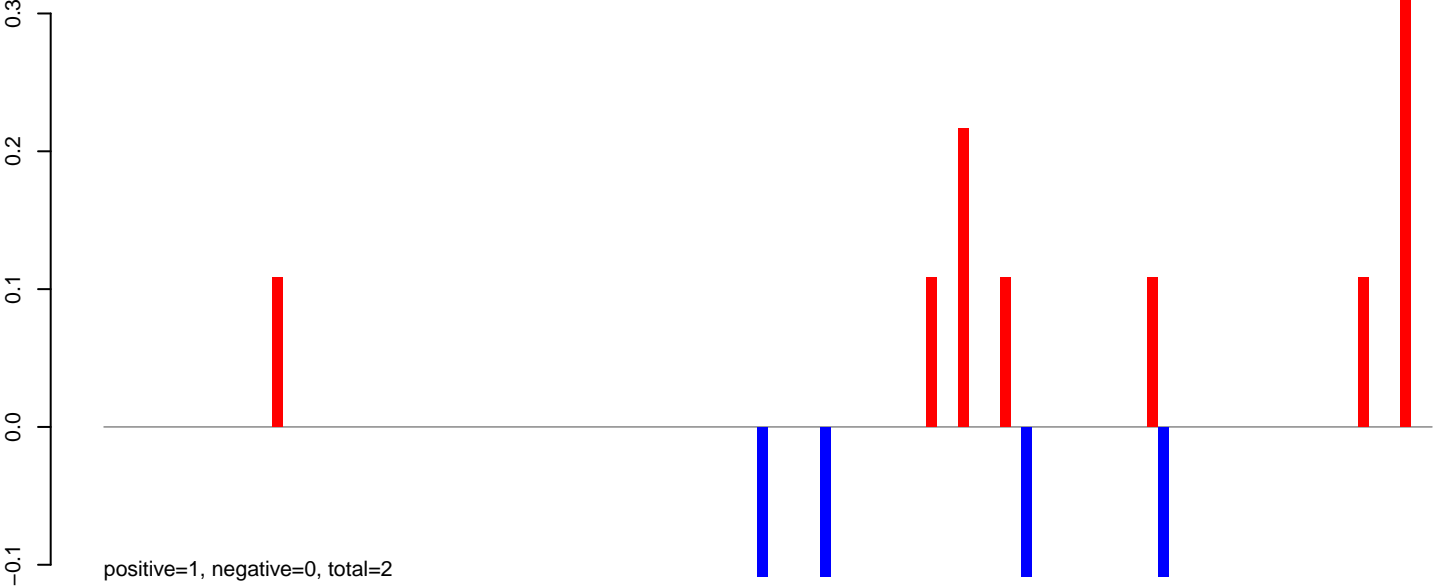


Window size=50, length=1670, TE@Clu-166_AG:1-1670

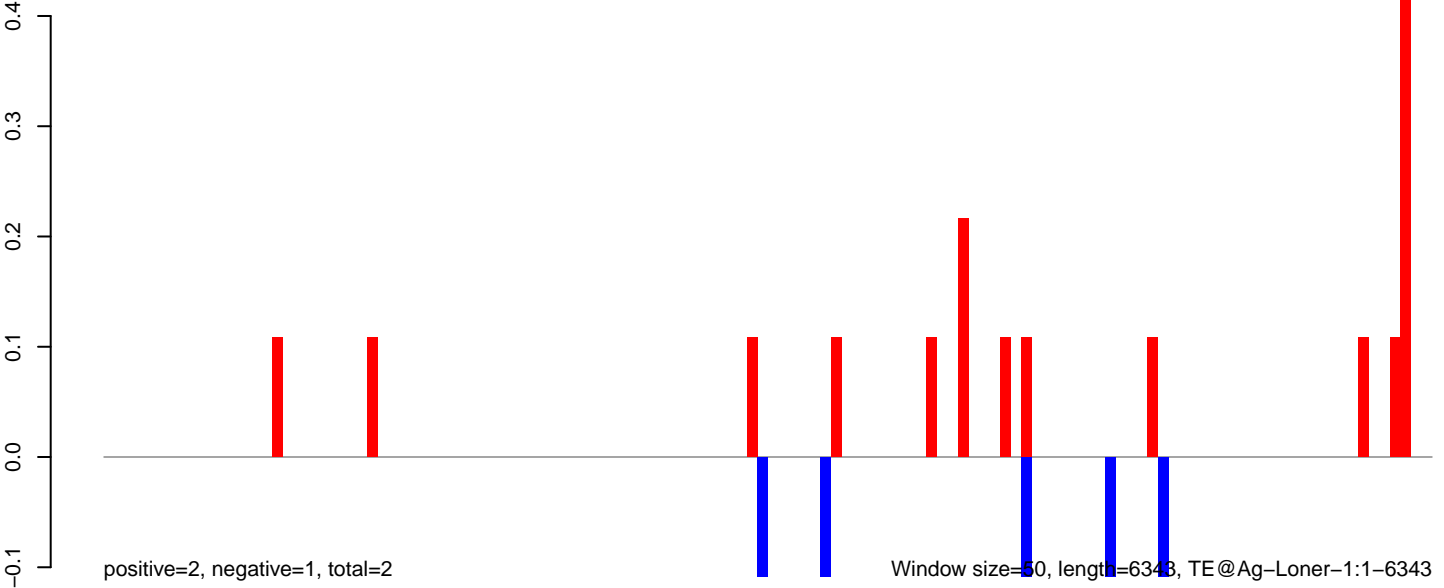
AnGam_Sua5b_cells.18_23.rep



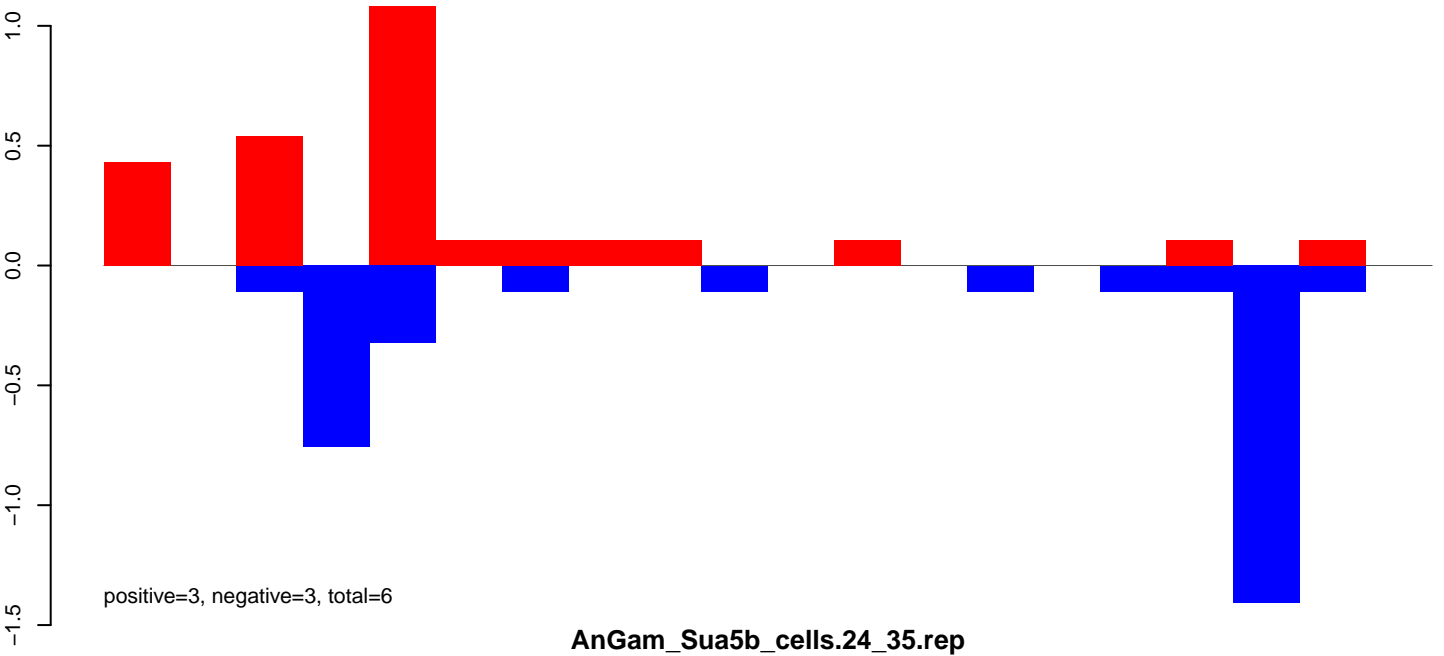
AnGam_Sua5b_cells.24_35.rep



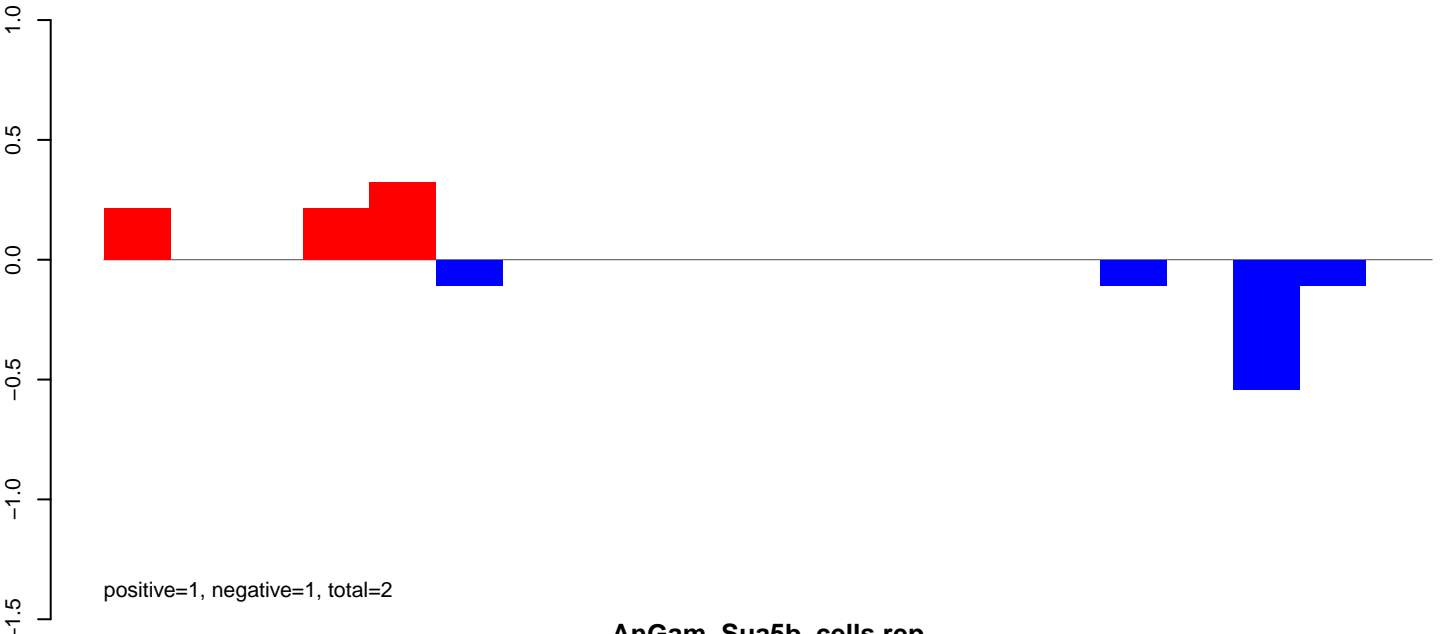
AnGam_Sua5b_cells.rep



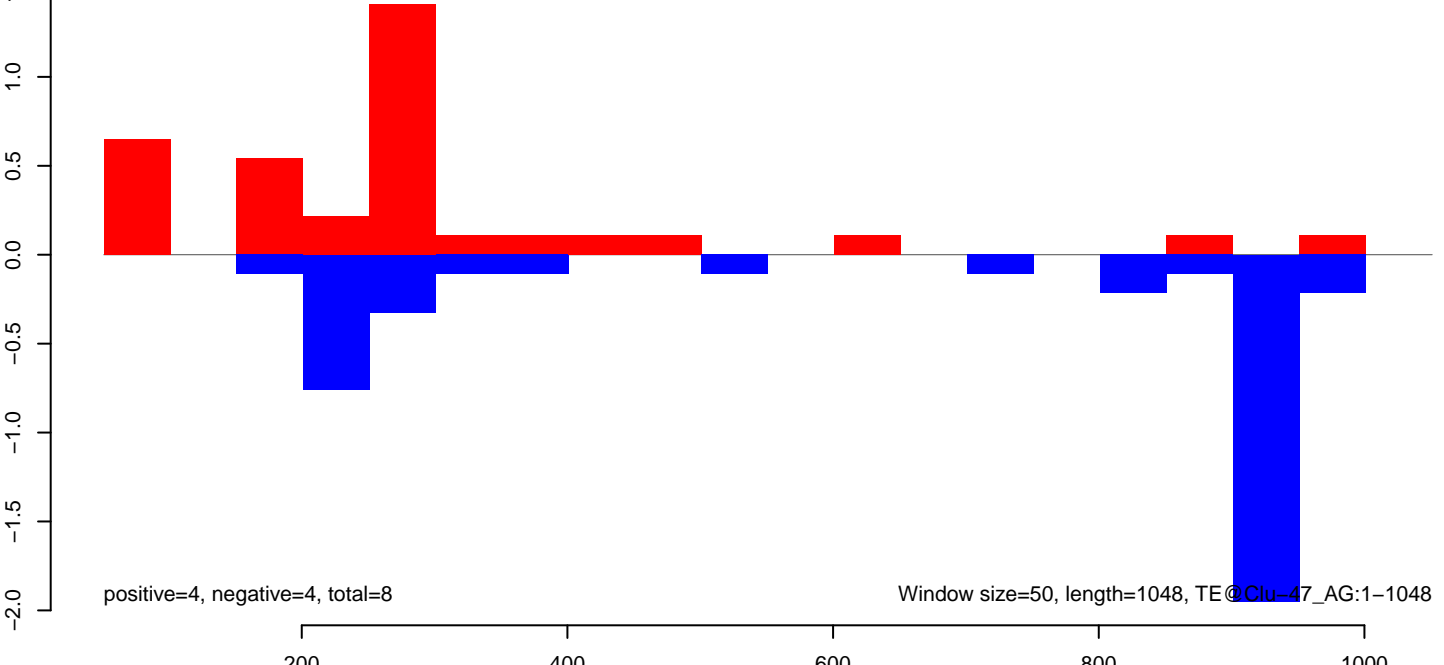
AnGam_Sua5b_cells.18_23.rep



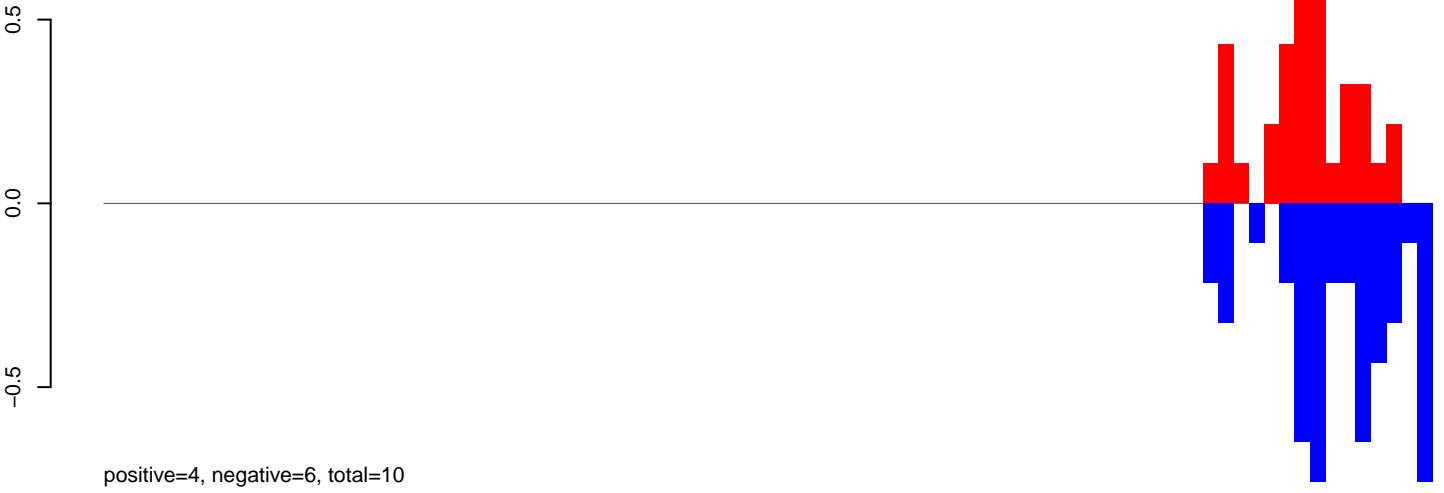
AnGam_Sua5b_cells.24_35.rep



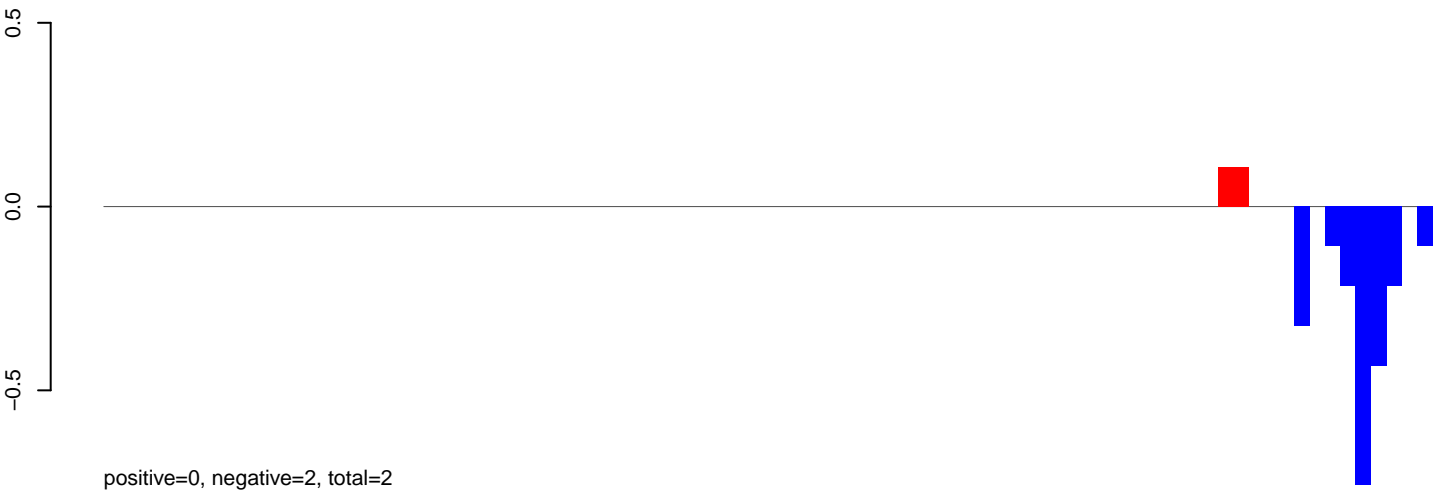
AnGam_Sua5b_cells.rep



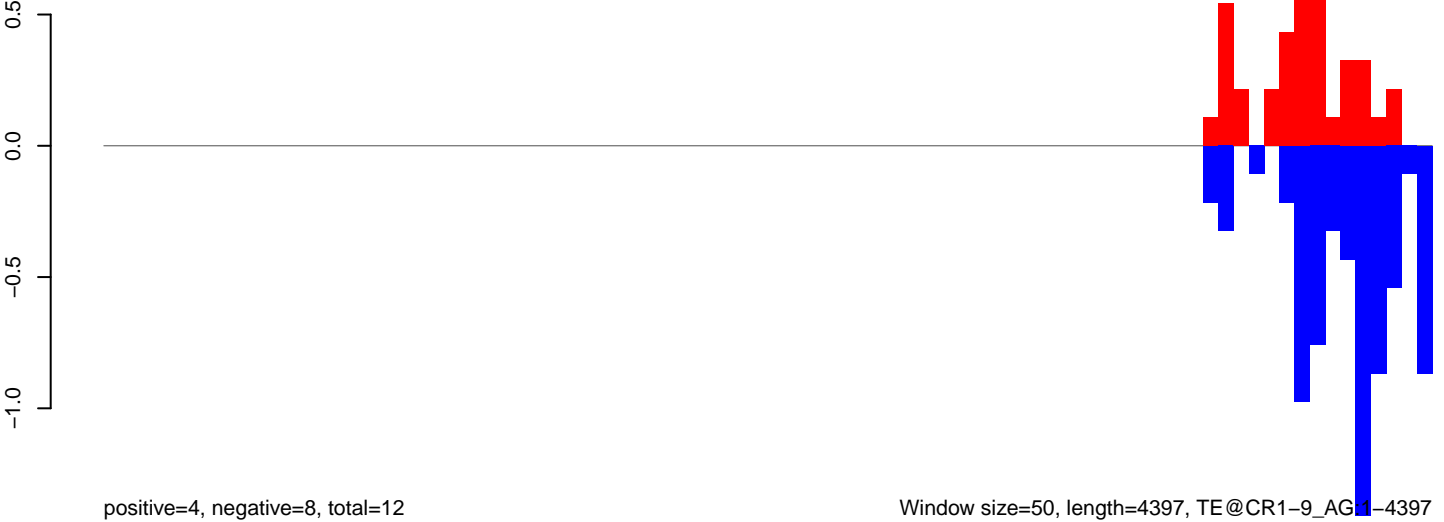
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



Window size=50, length=4397, TE@CR1-9_AG1-4397

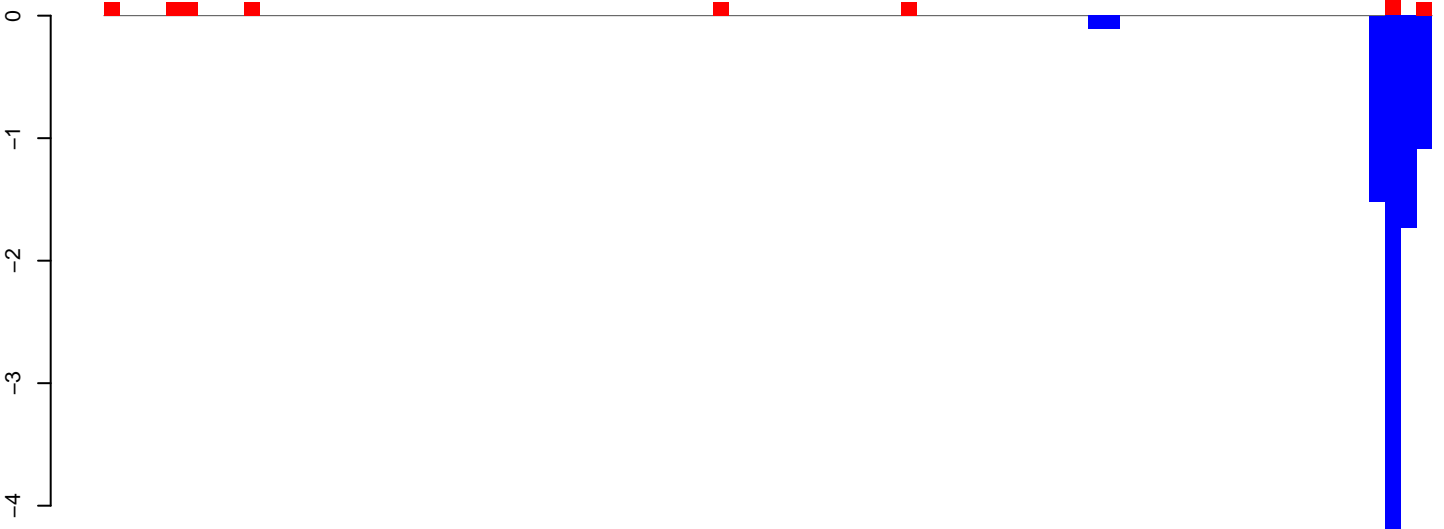
0 1000 2000 3000 4000

AnGam_Sua5b_cells.18_23.rep



positive=1, negative=2, total=2

AnGam_Sua5b_cells.24_35.rep



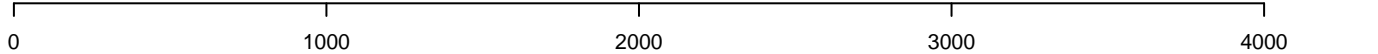
positive=1, negative=10, total=11

AnGam_Sua5b_cells.rep

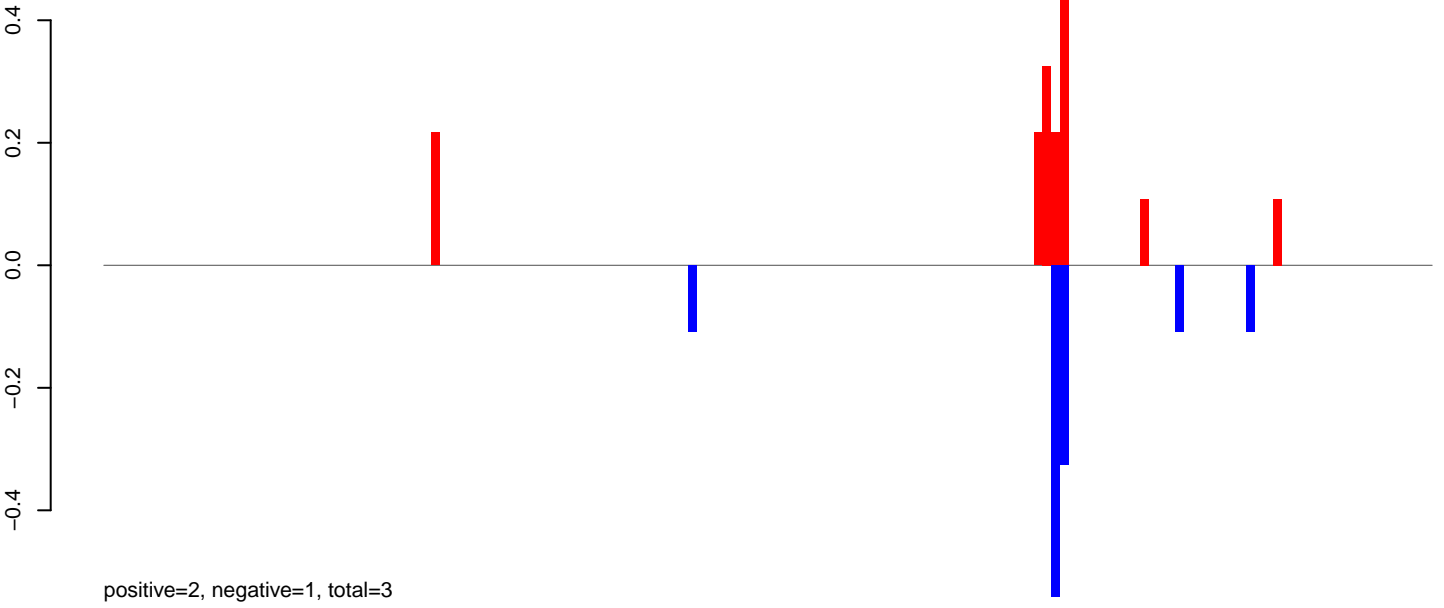


positive=2, negative=12, total=14

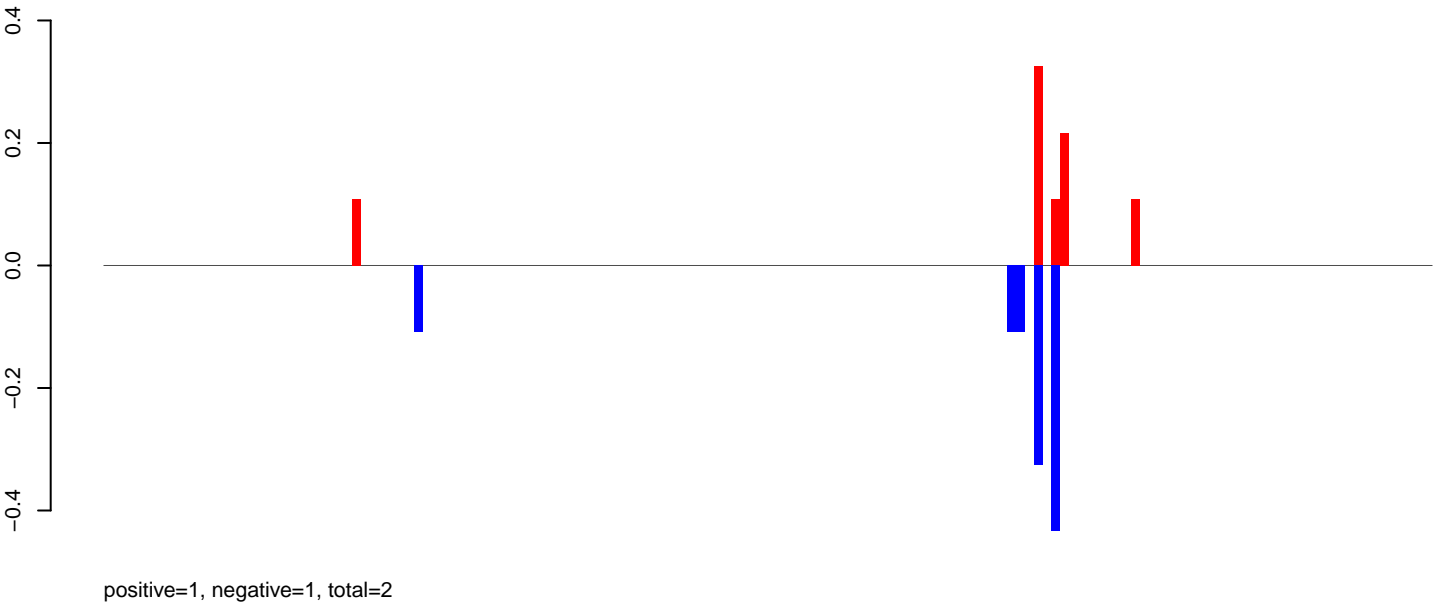
Window size=50, length=4293, TE@CR1-6_AG:1-4293



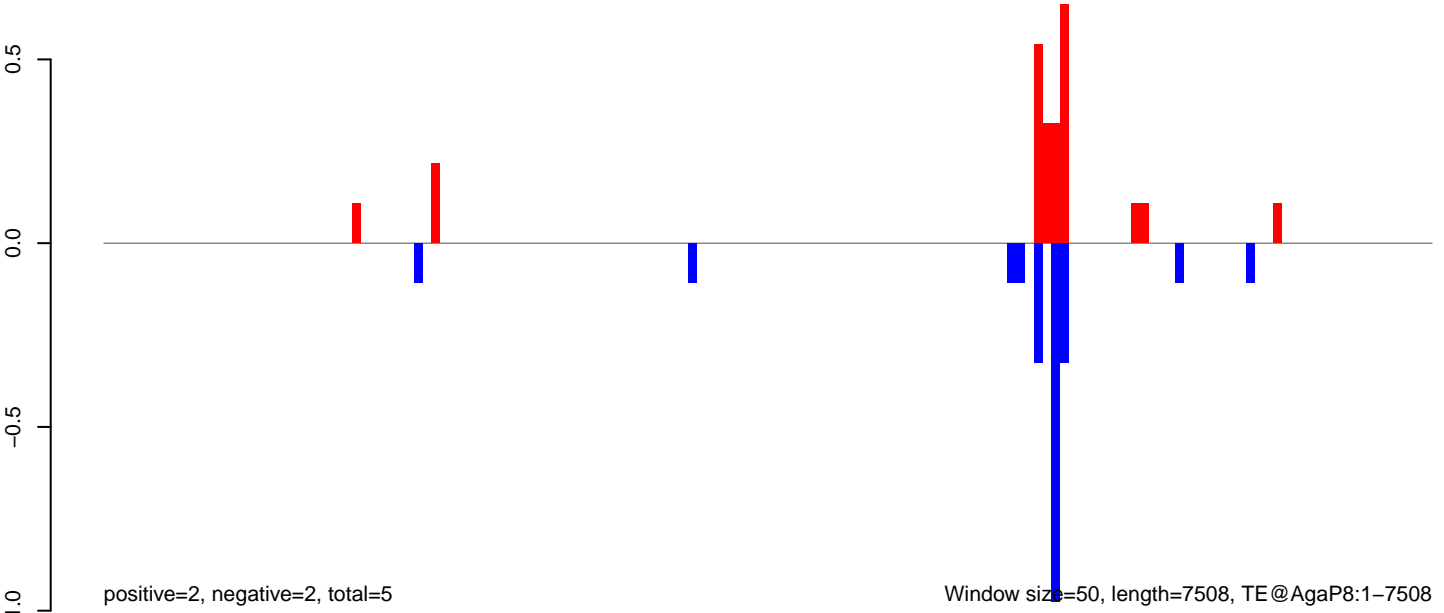
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

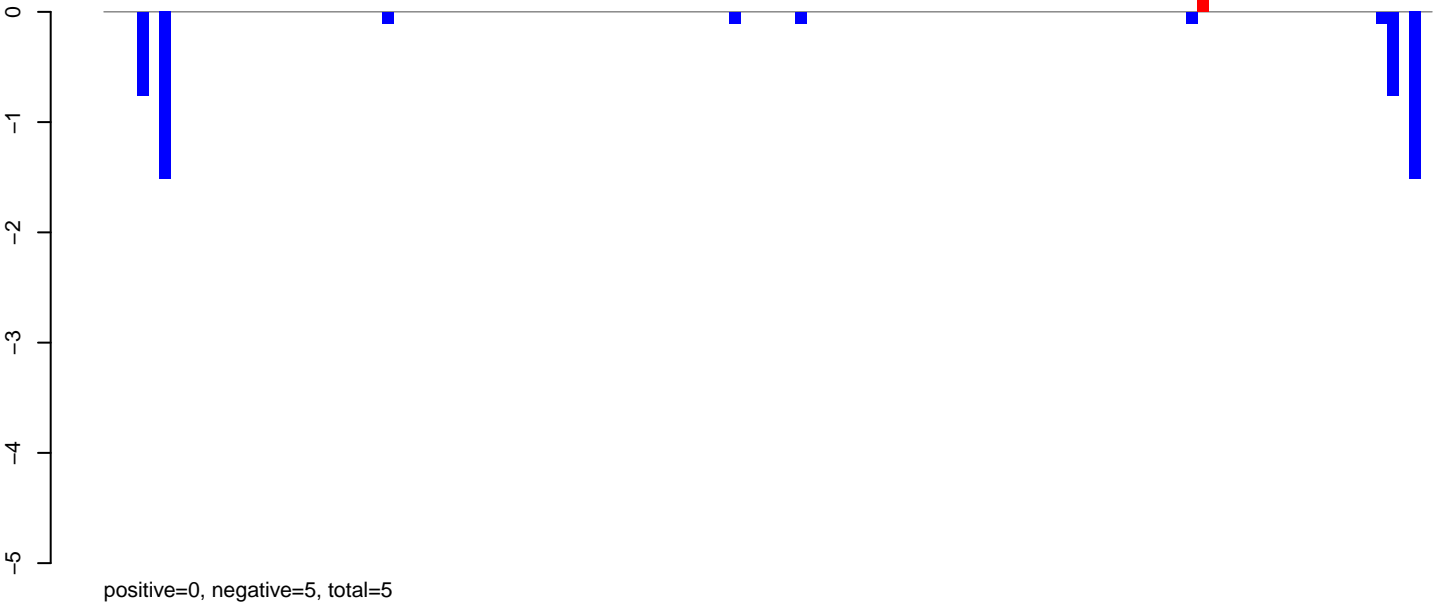


AnGam_Sua5b_cells.rep

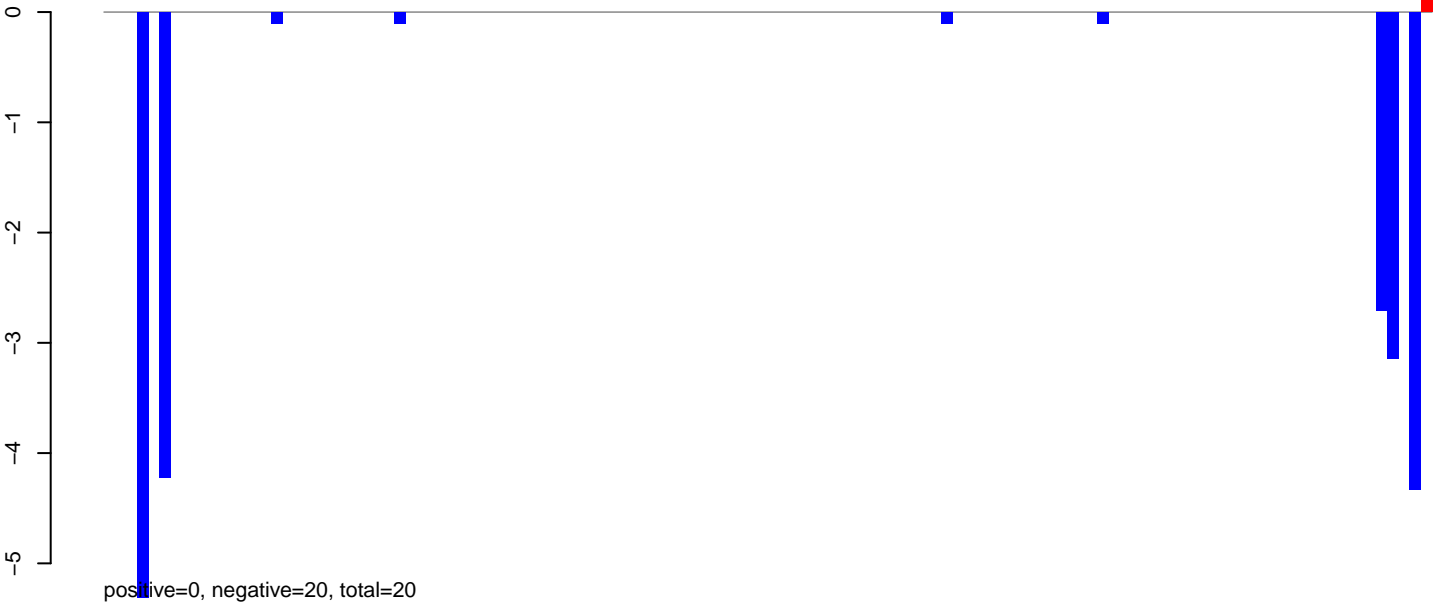


Window size=50, length=7508, TE@AgaP8:1-7508

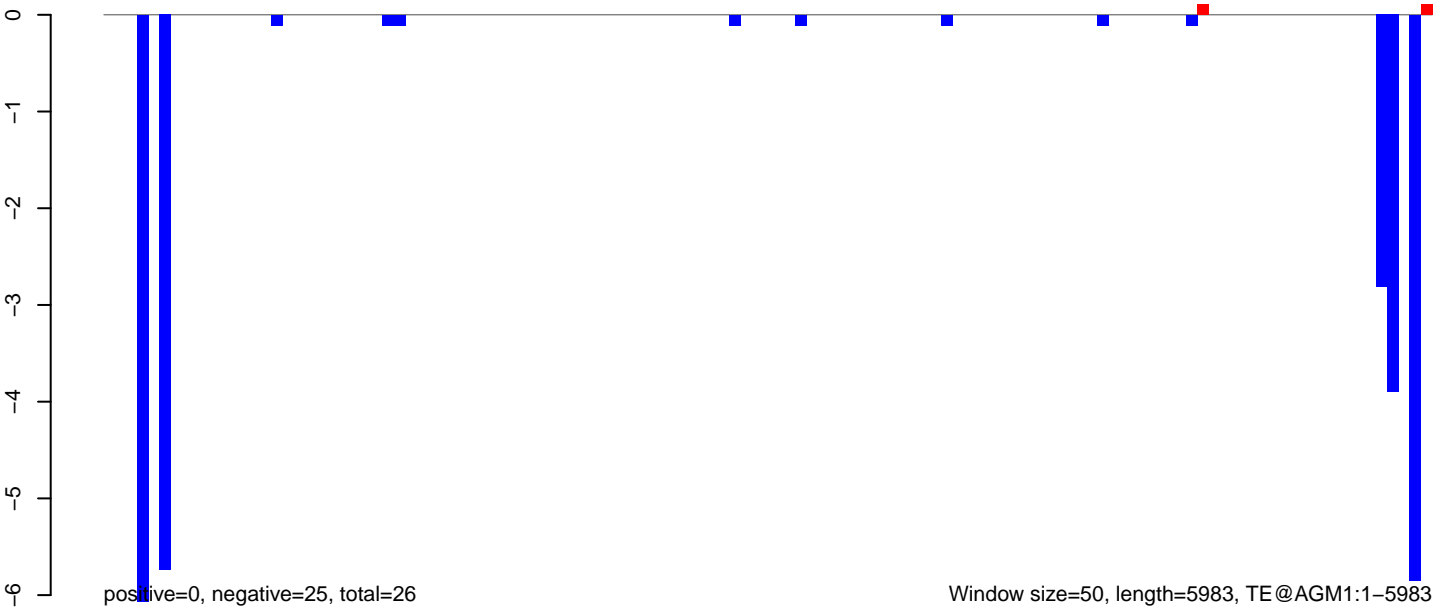
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

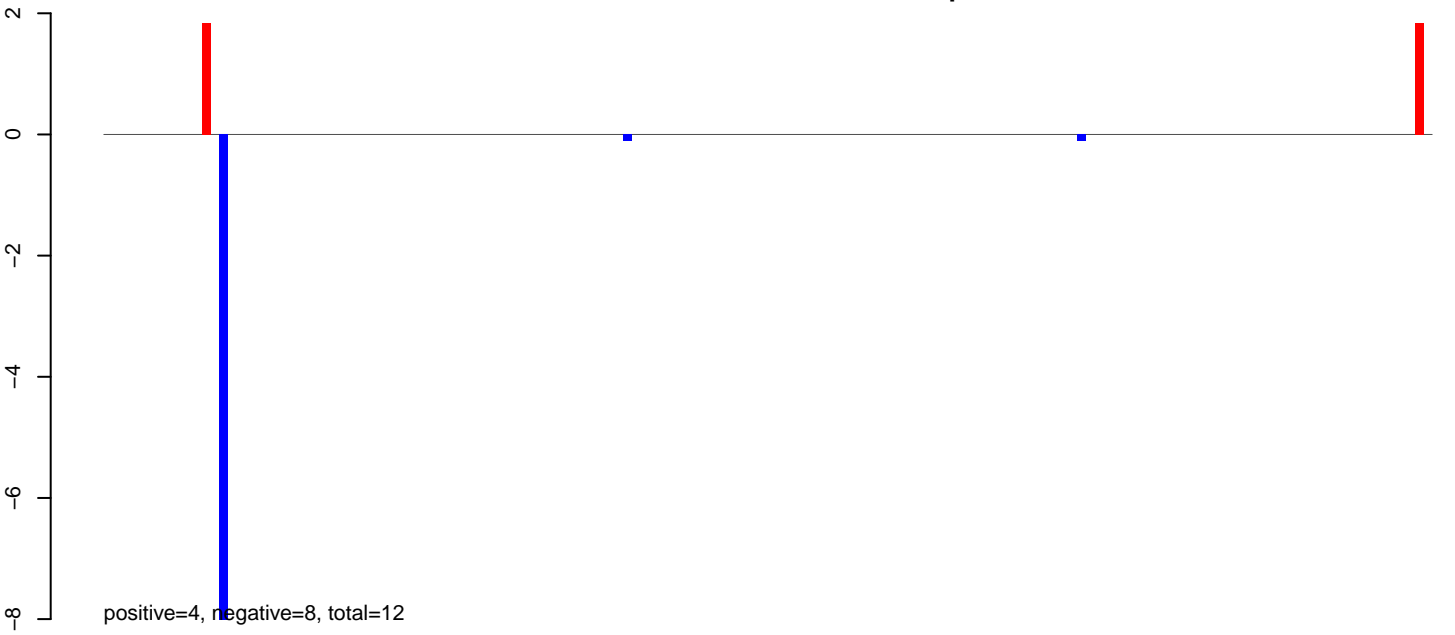


AnGam_Sua5b_cells.rep

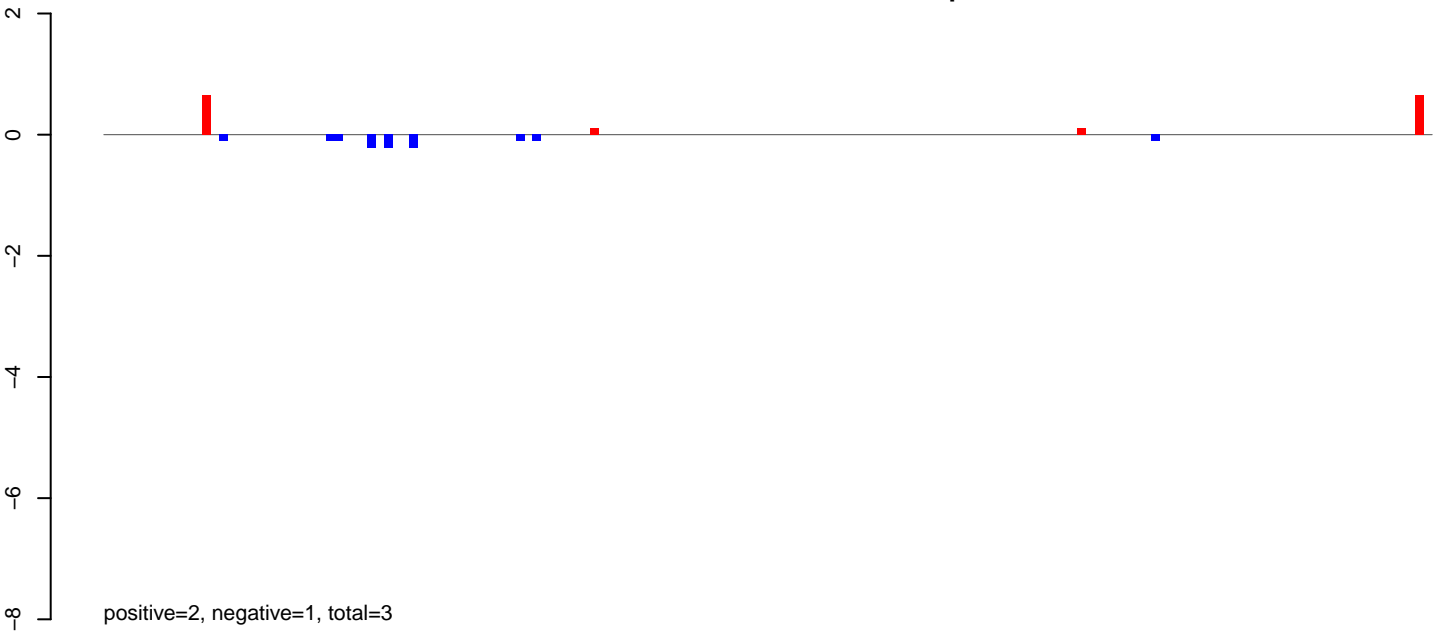


Window size=50, length=5983, TE@AGM1:1-5983

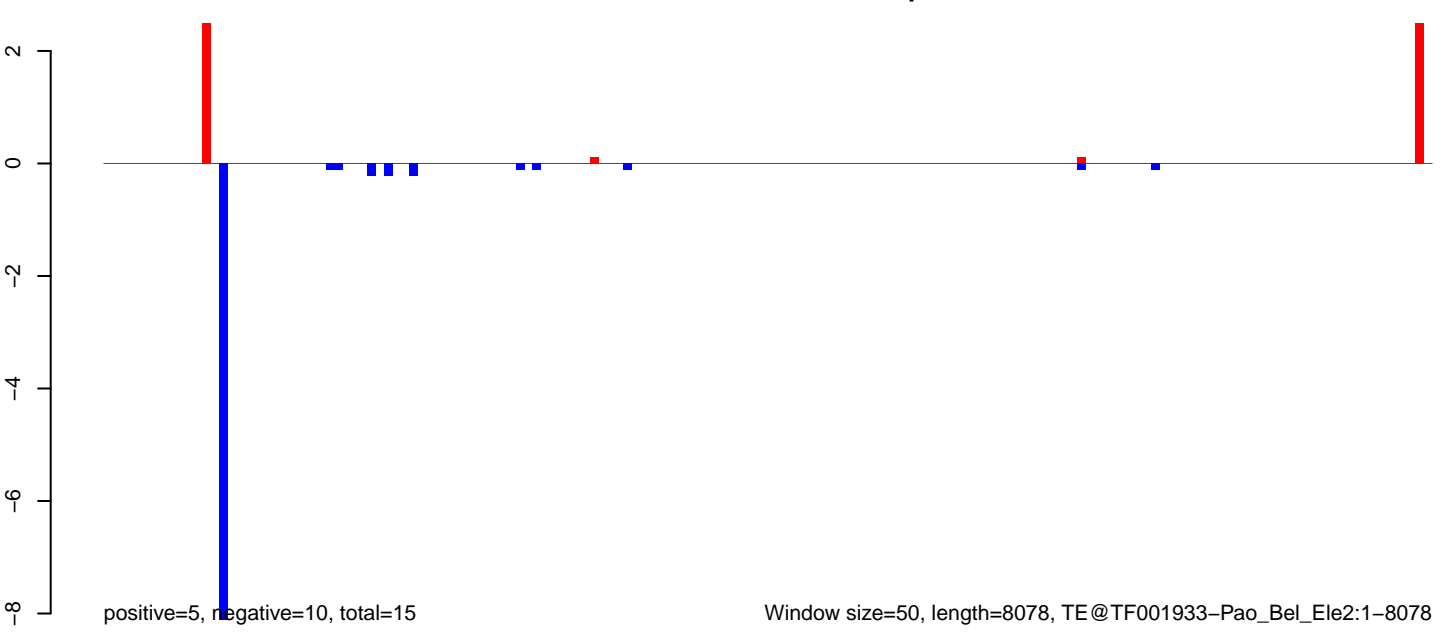
AnGam_Sua5b_cells.18_23.rep



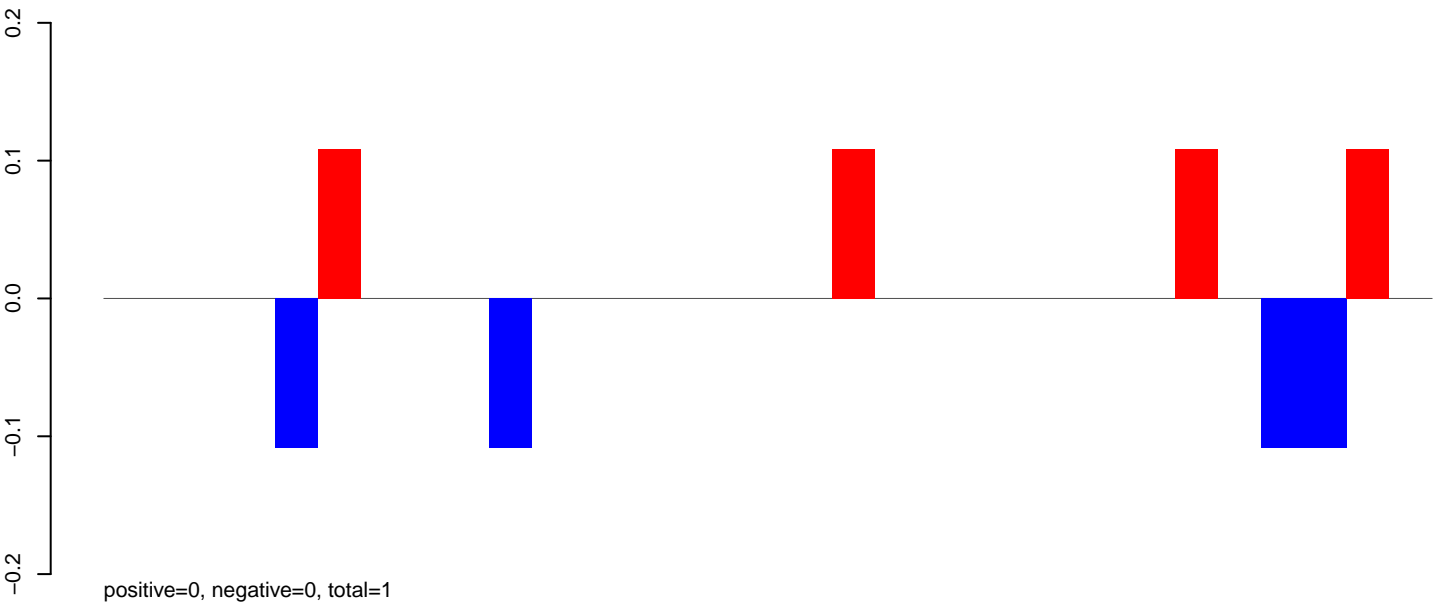
AnGam_Sua5b_cells.24_35.rep



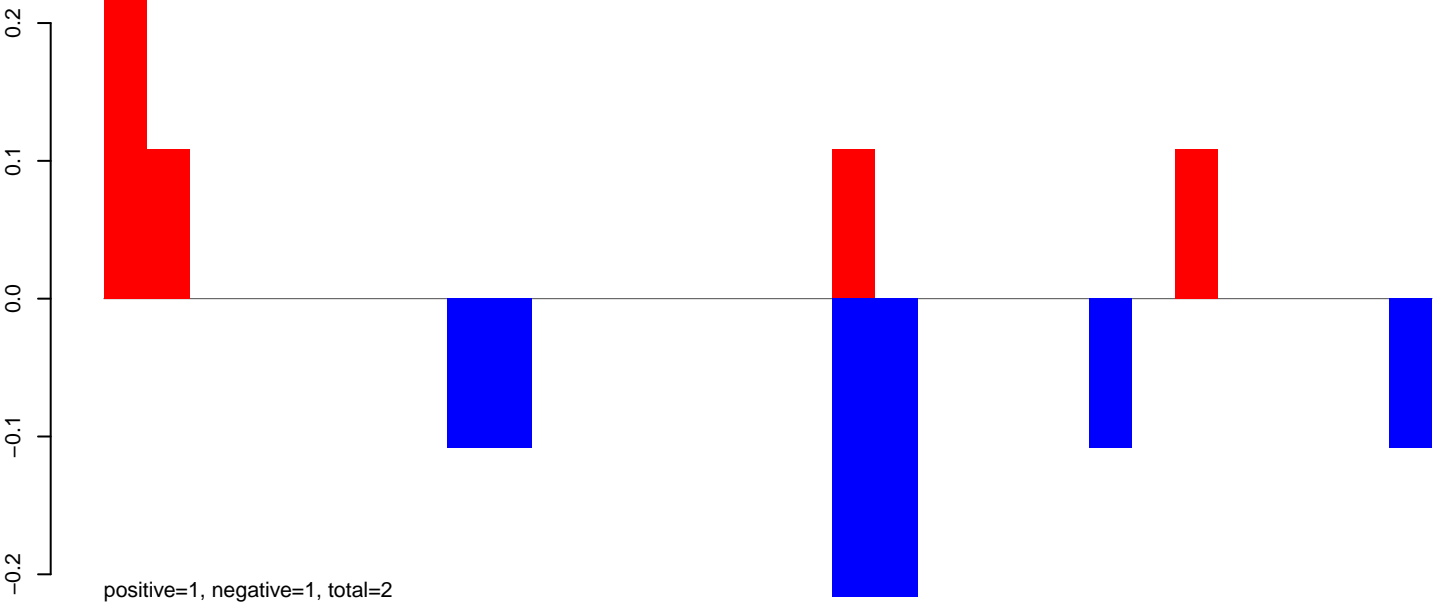
AnGam_Sua5b_cells.rep



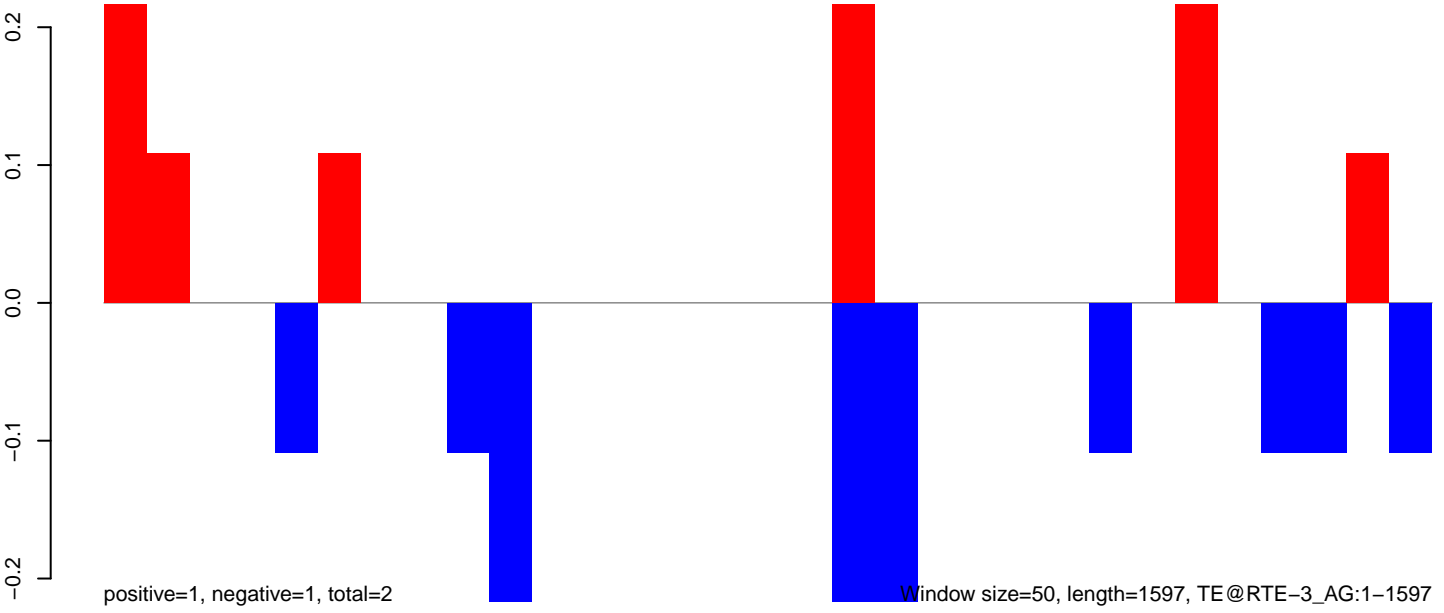
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

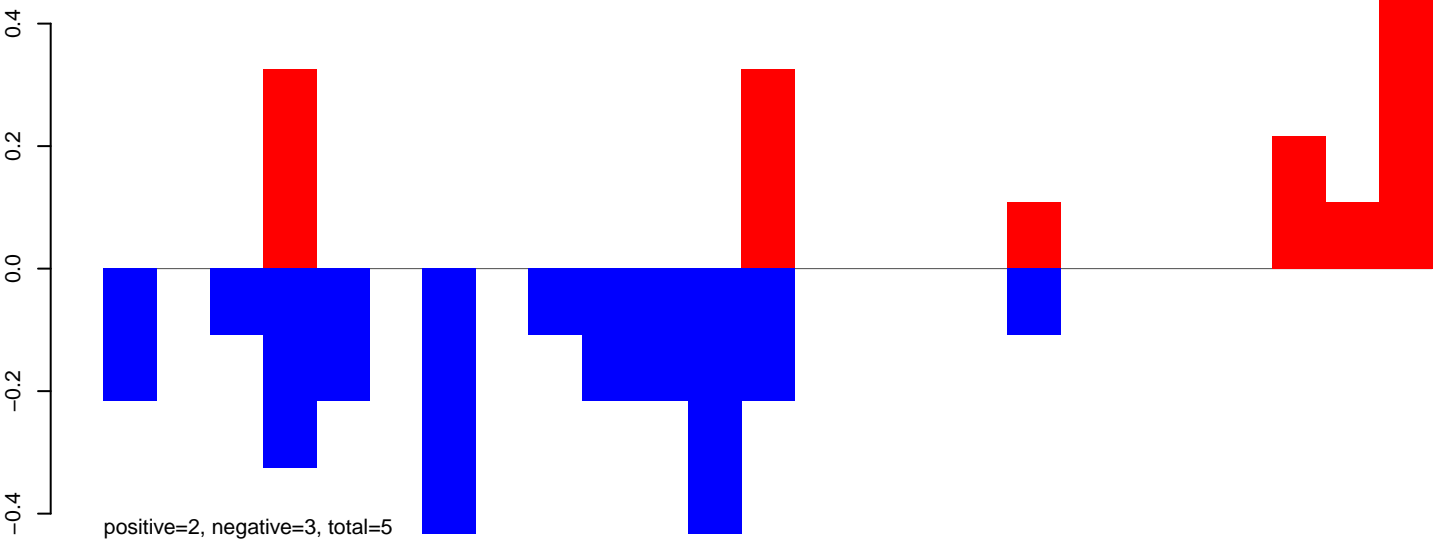


AnGam_Sua5b_cells.rep

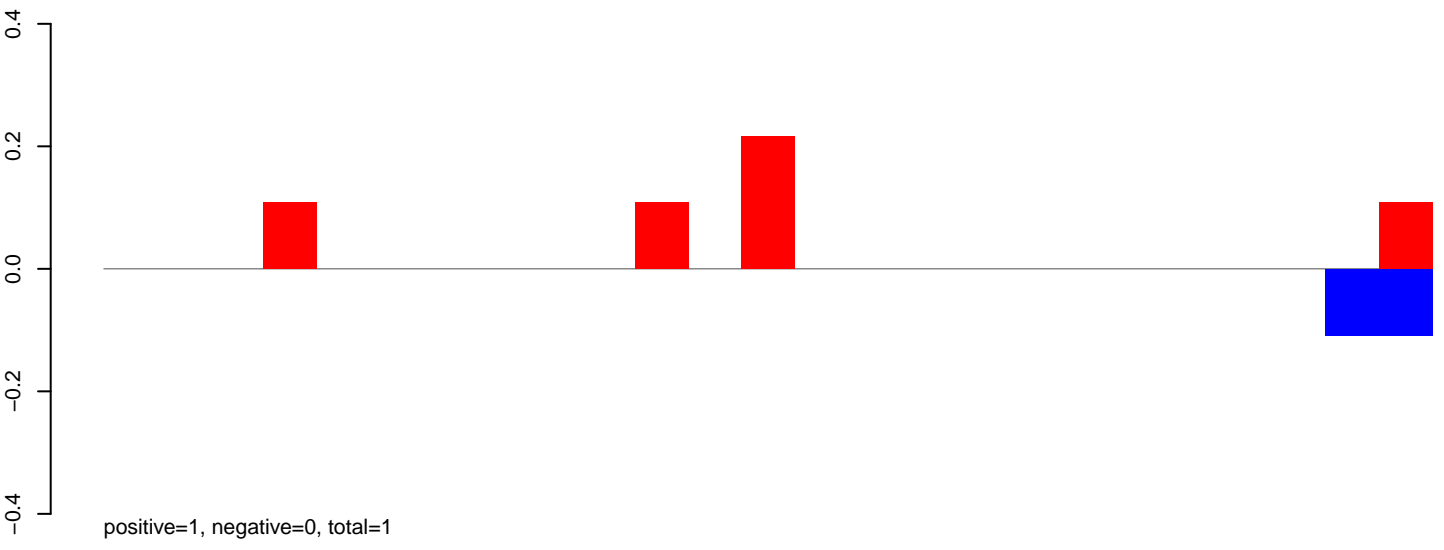


Window size=50, length=1597, TE@RTE-3_AG:1-1597

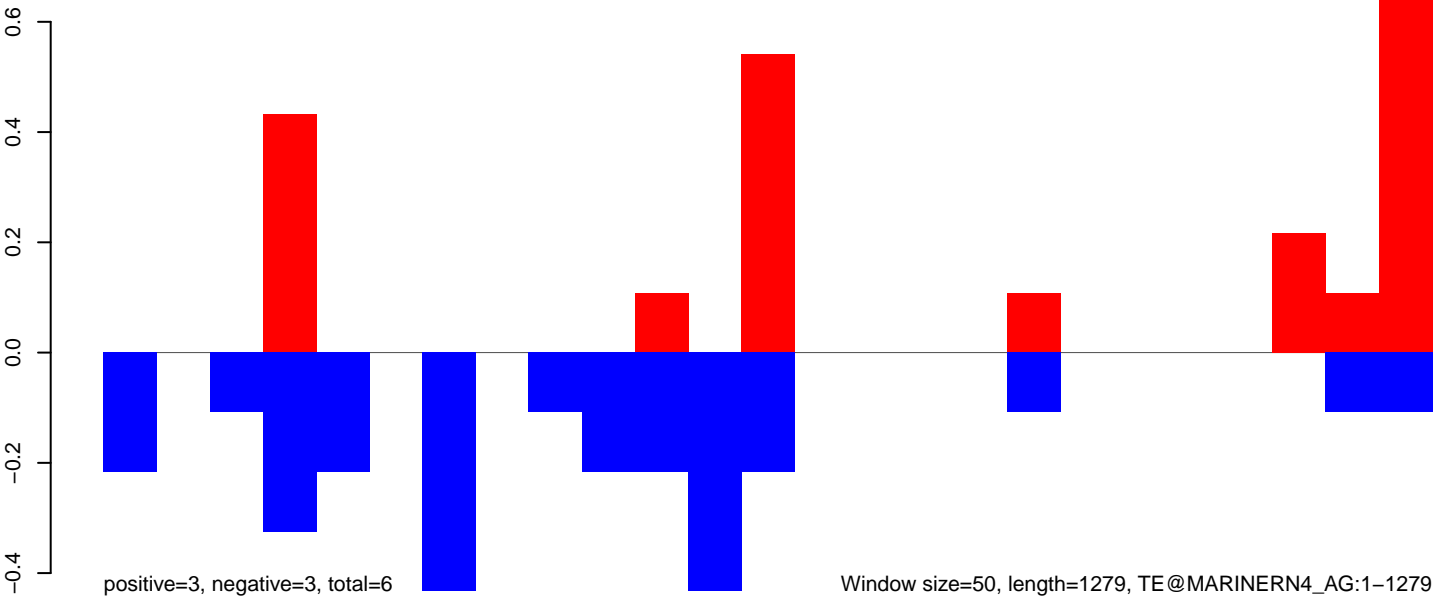
AnGam_Sua5b_cells.18_23.rep



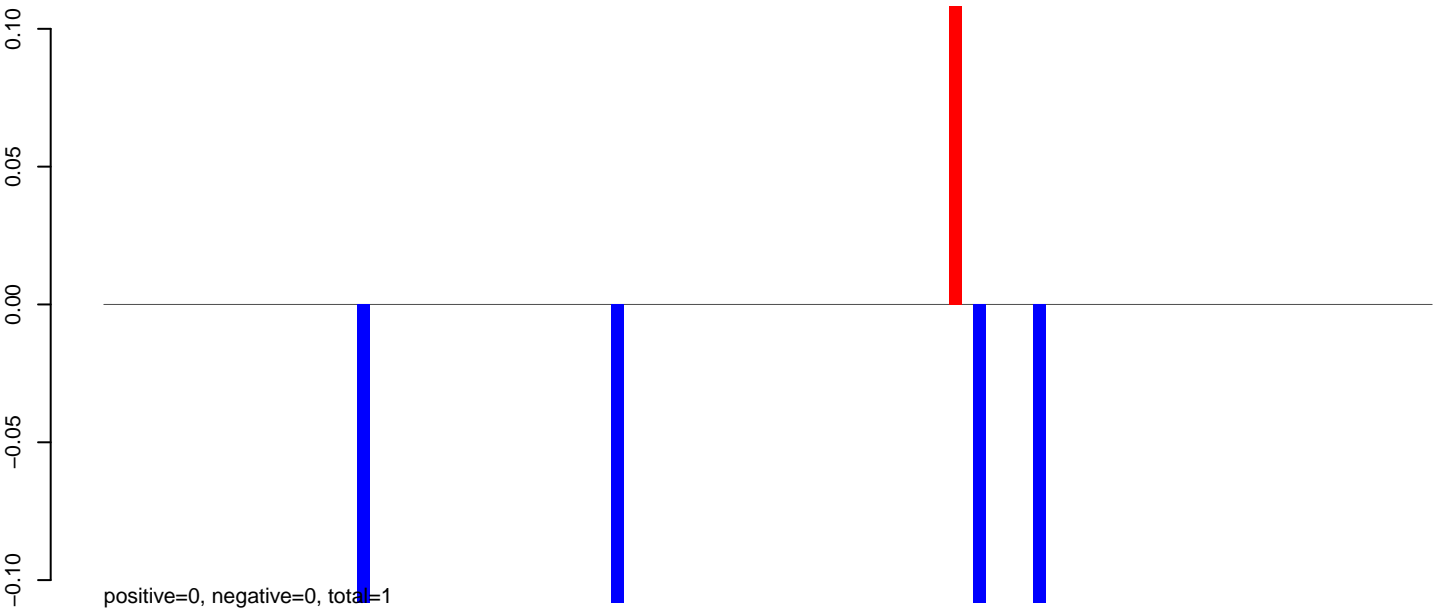
AnGam_Sua5b_cells.24_35.rep



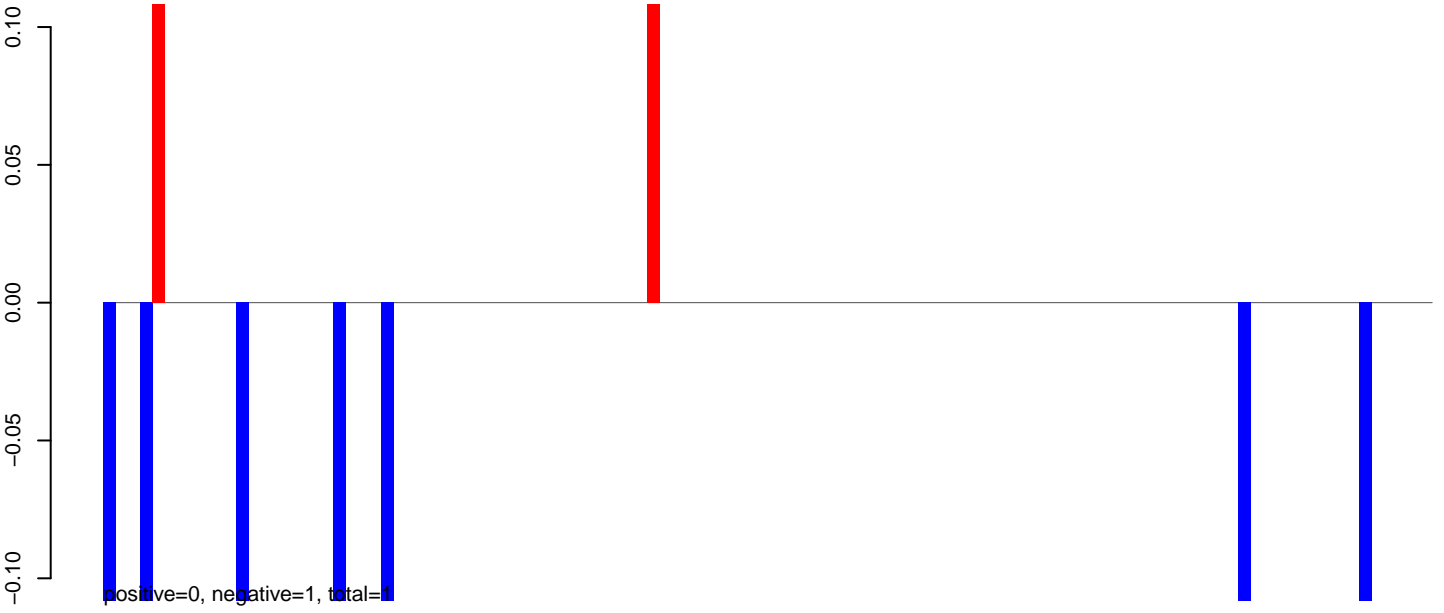
AnGam_Sua5b_cells.rep



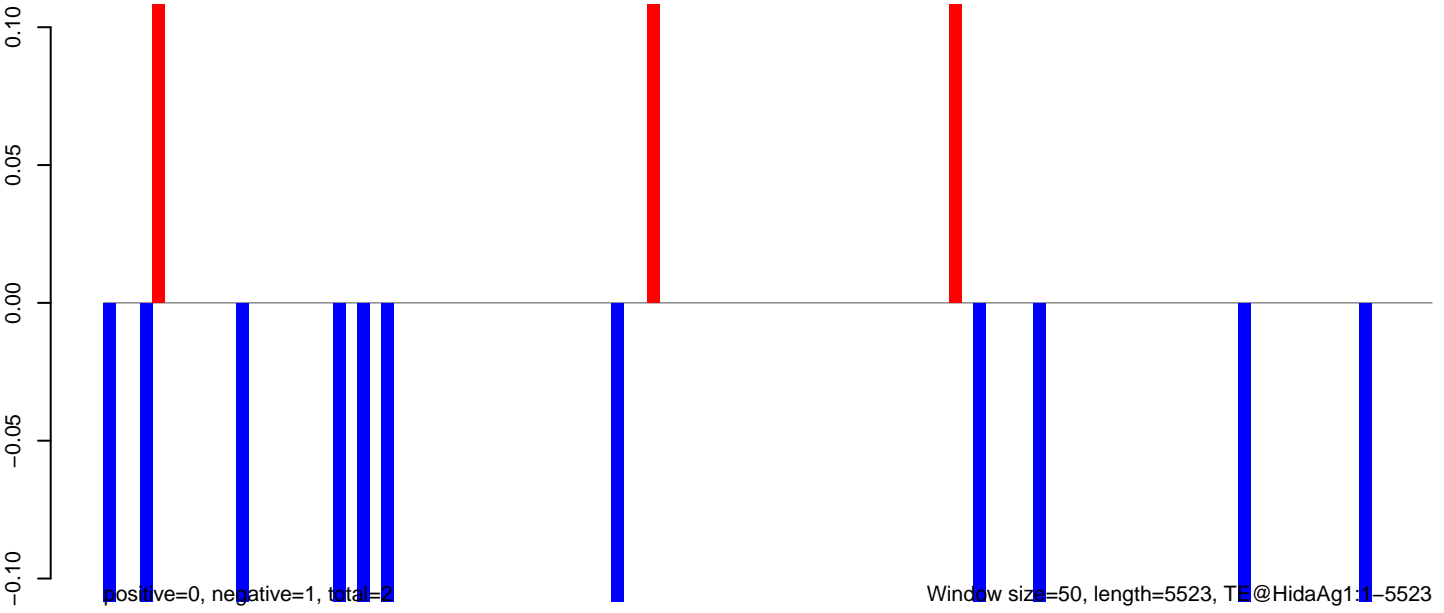
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

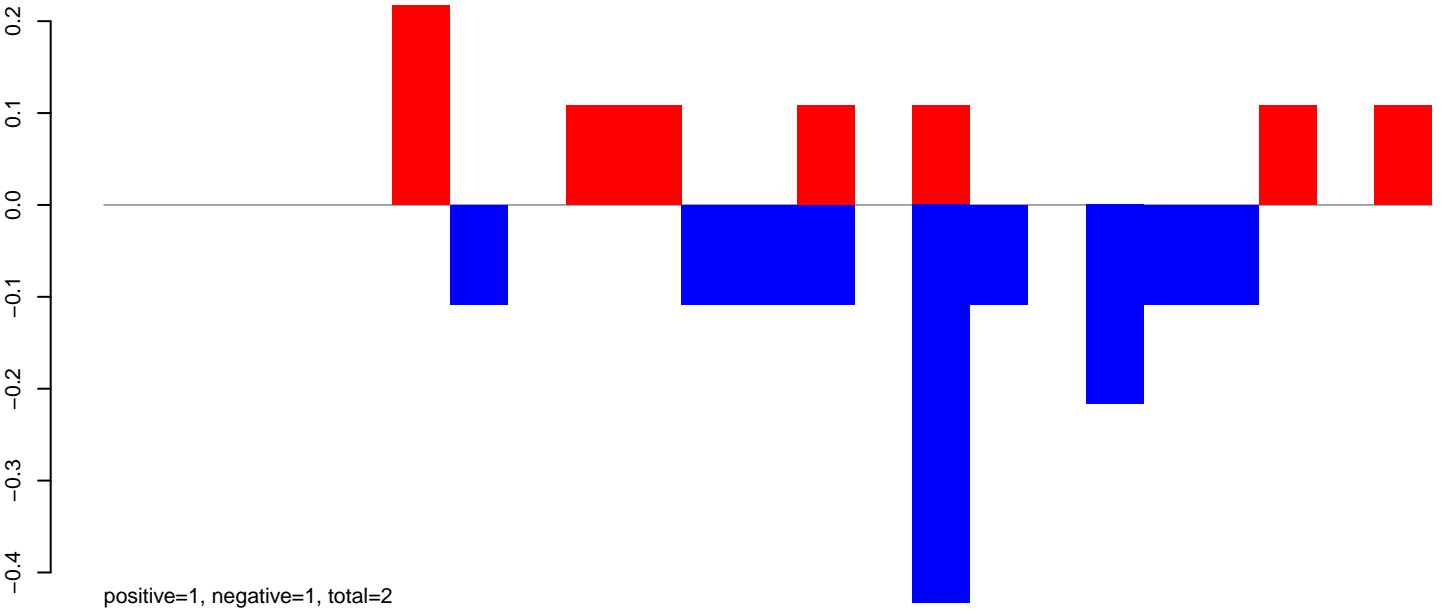


AnGam_Sua5b_cells.rep

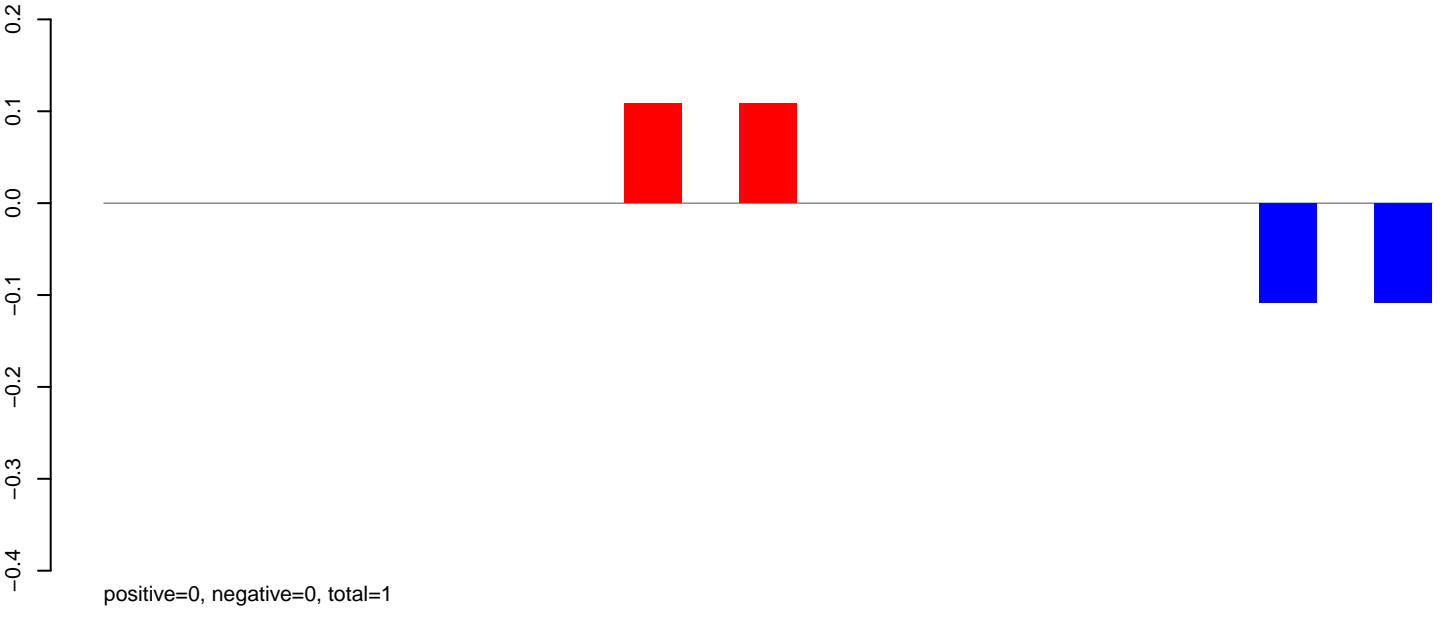


0 1000 2000 3000 4000 5000

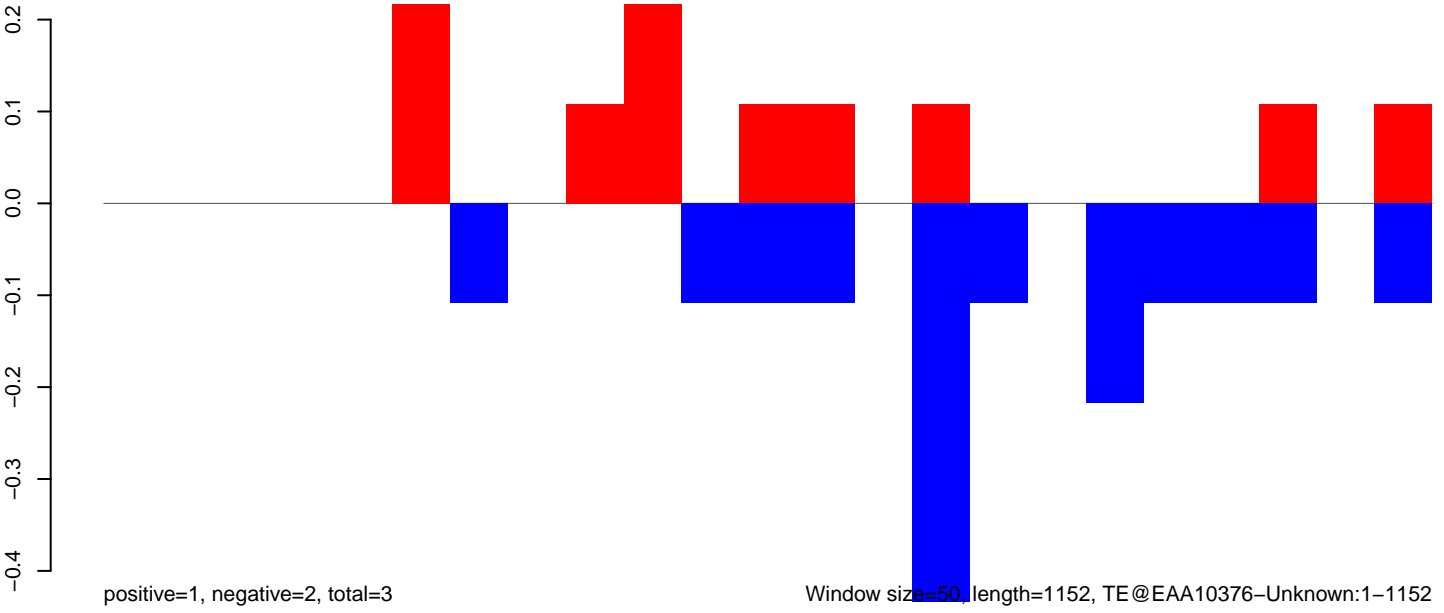
AnGam_Sua5b_cells.18_23.rep



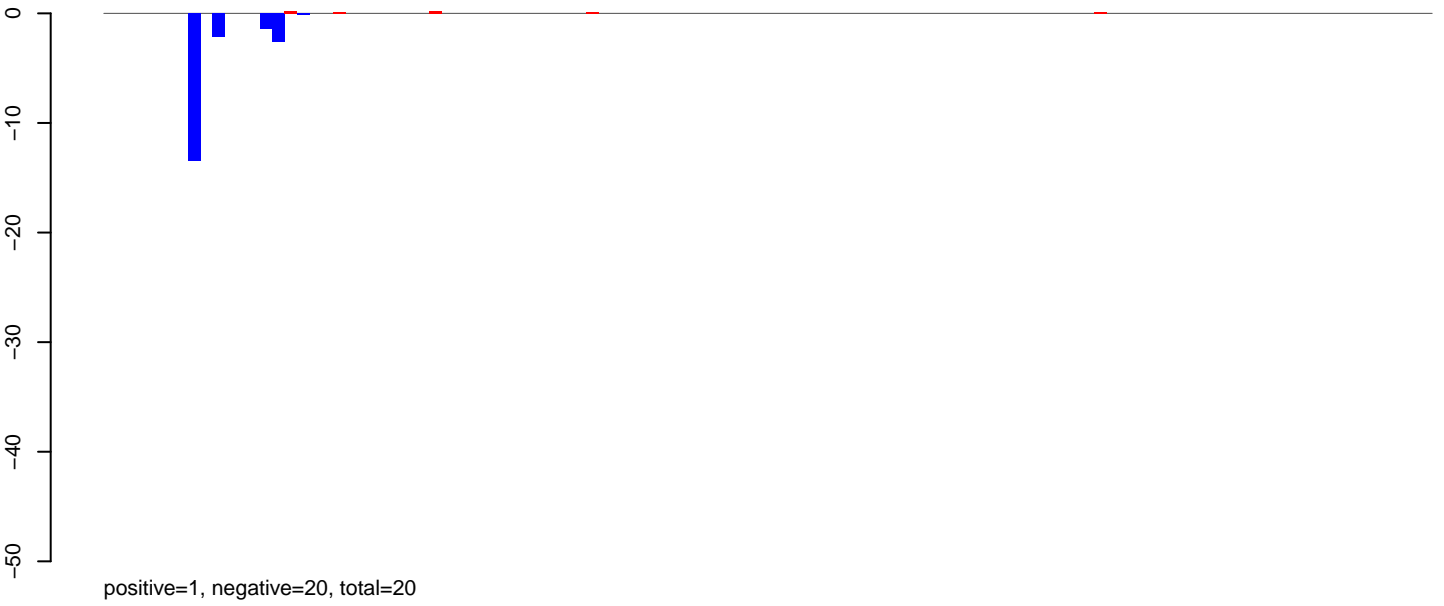
AnGam_Sua5b_cells.24_35.rep



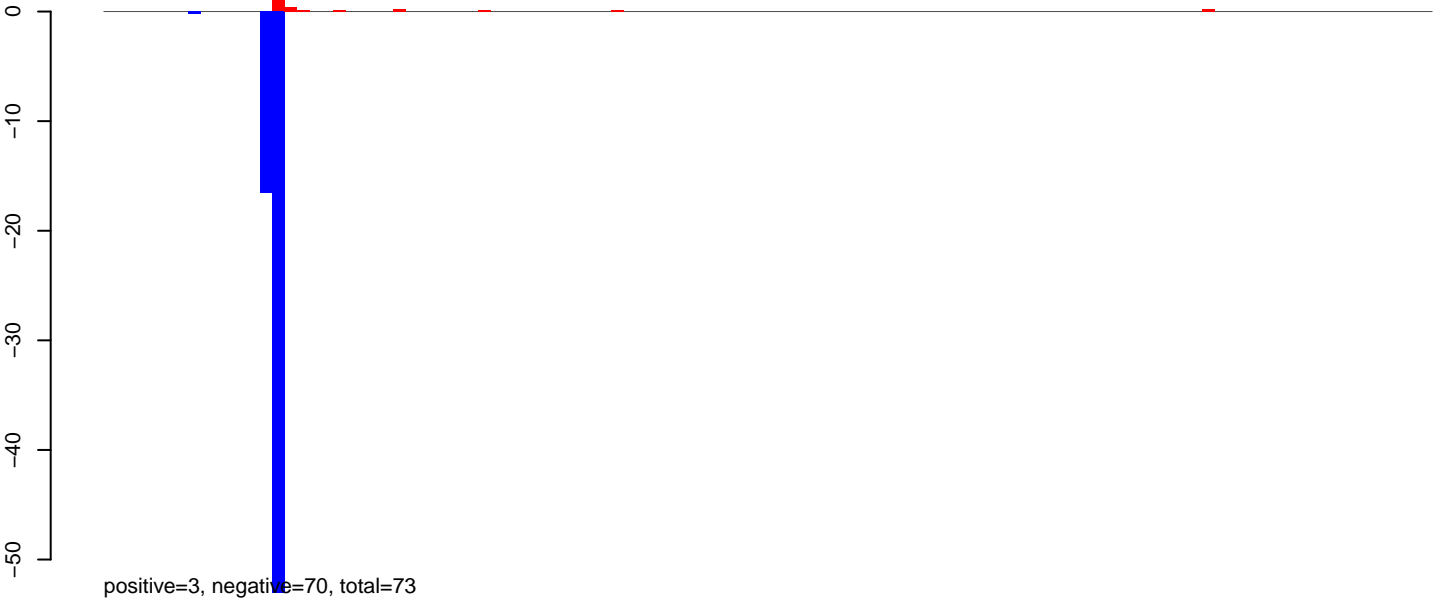
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



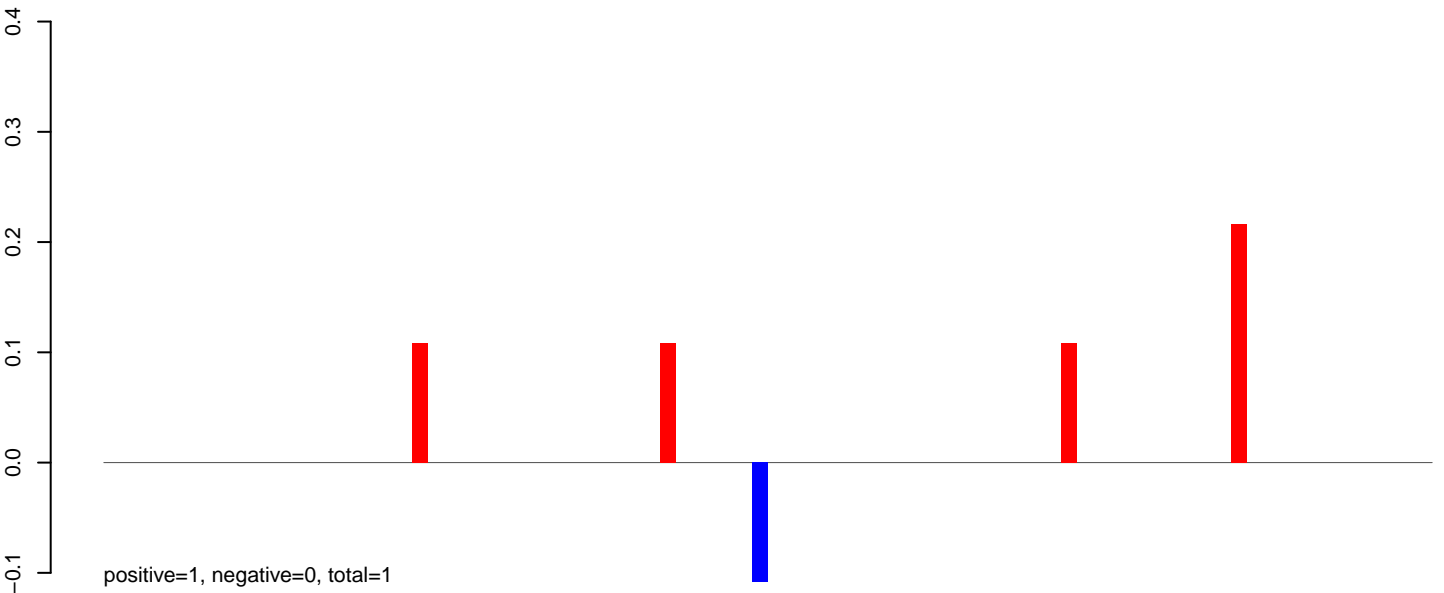
AnGam_Sua5b_cells.24_35.rep



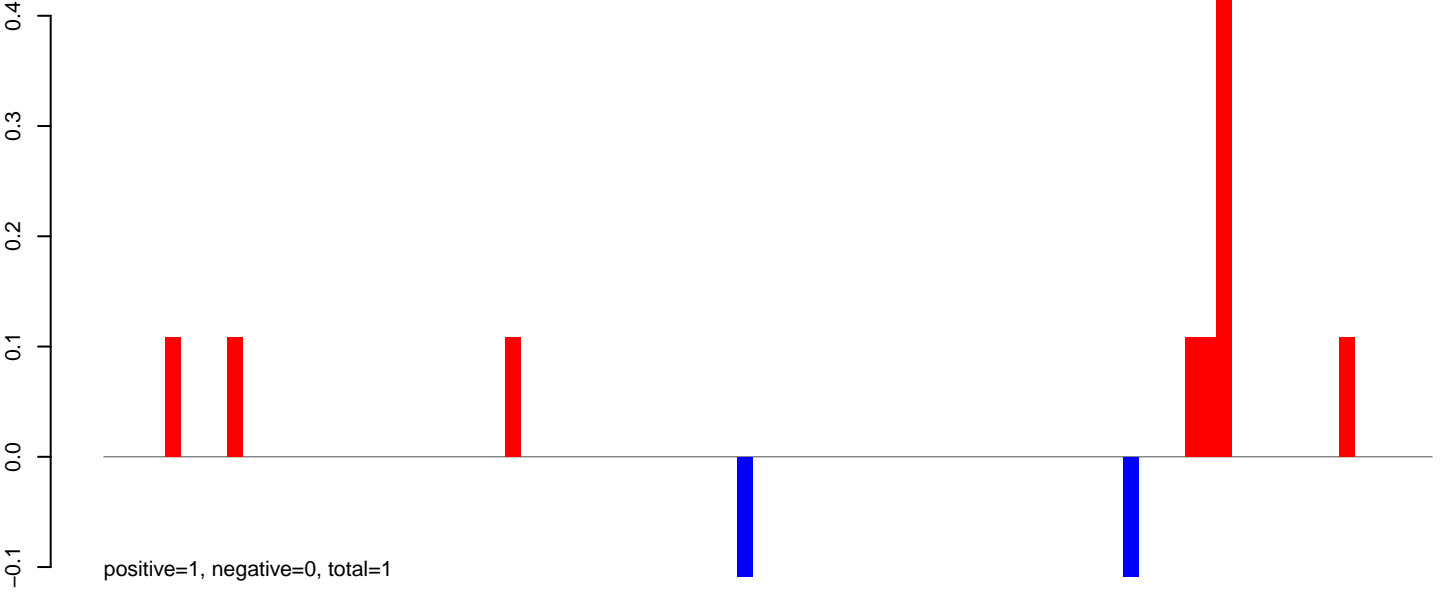
AnGam_Sua5b_cells.rep



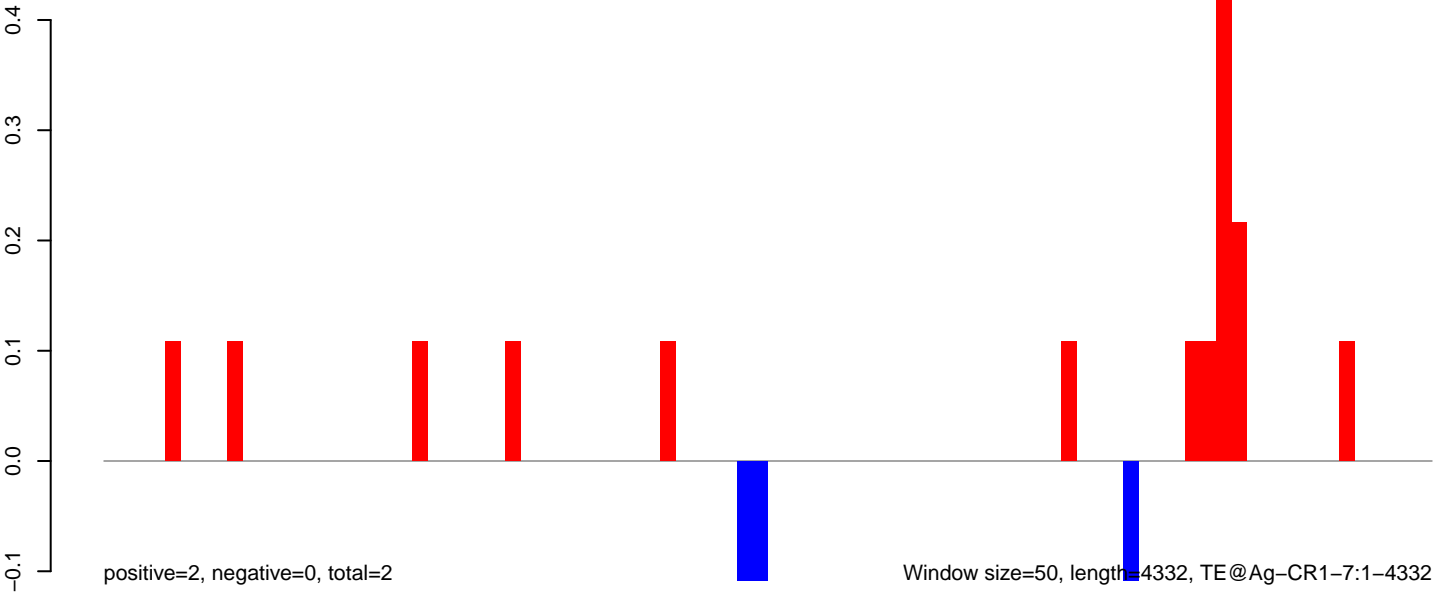
AnGam_Sua5b_cells.18_23.rep



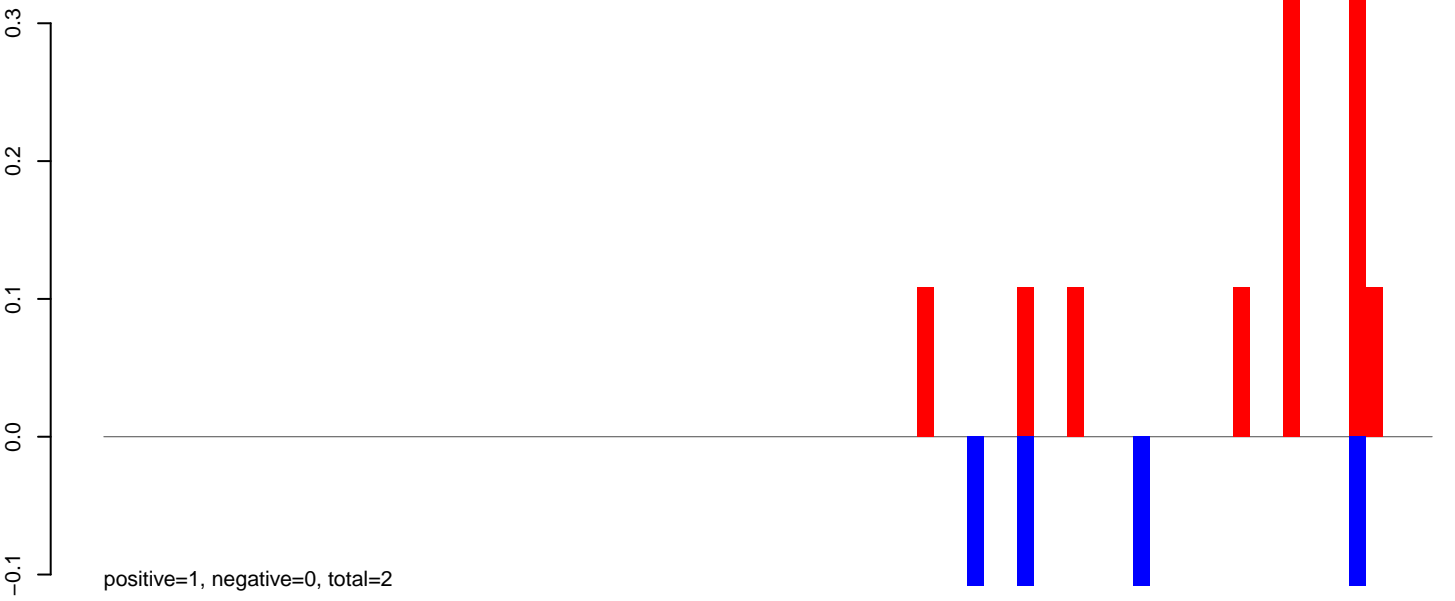
AnGam_Sua5b_cells.24_35.rep



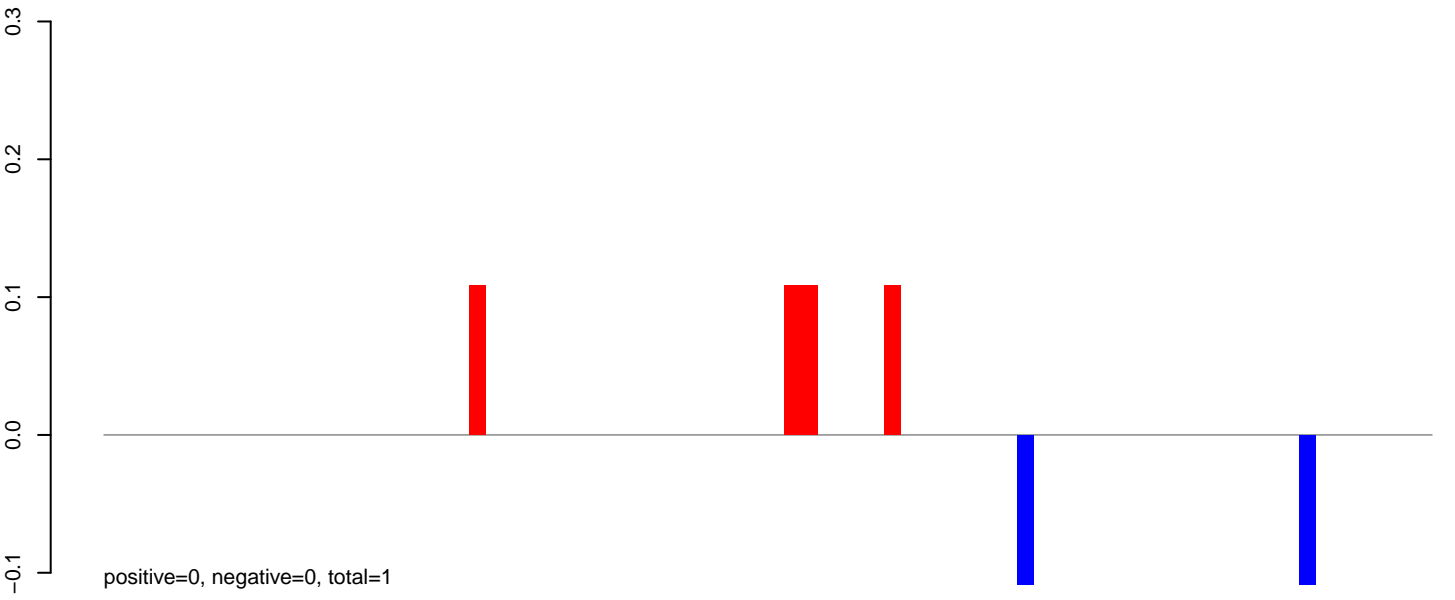
AnGam_Sua5b_cells.rep



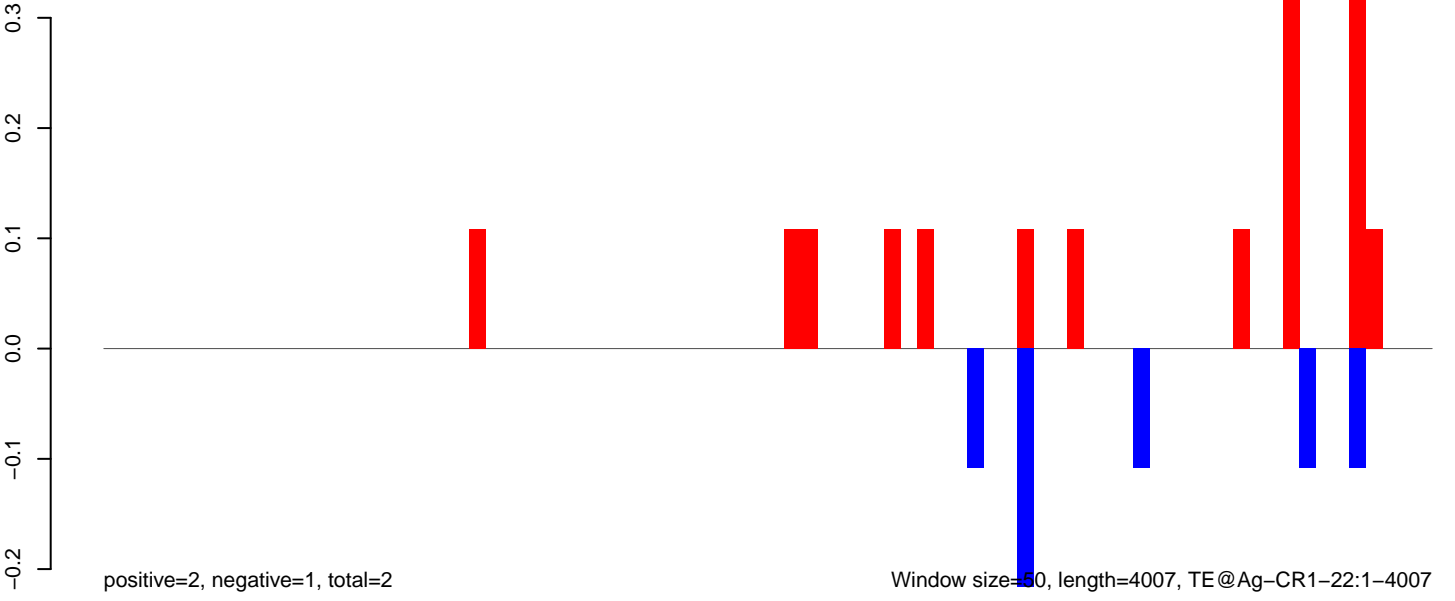
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



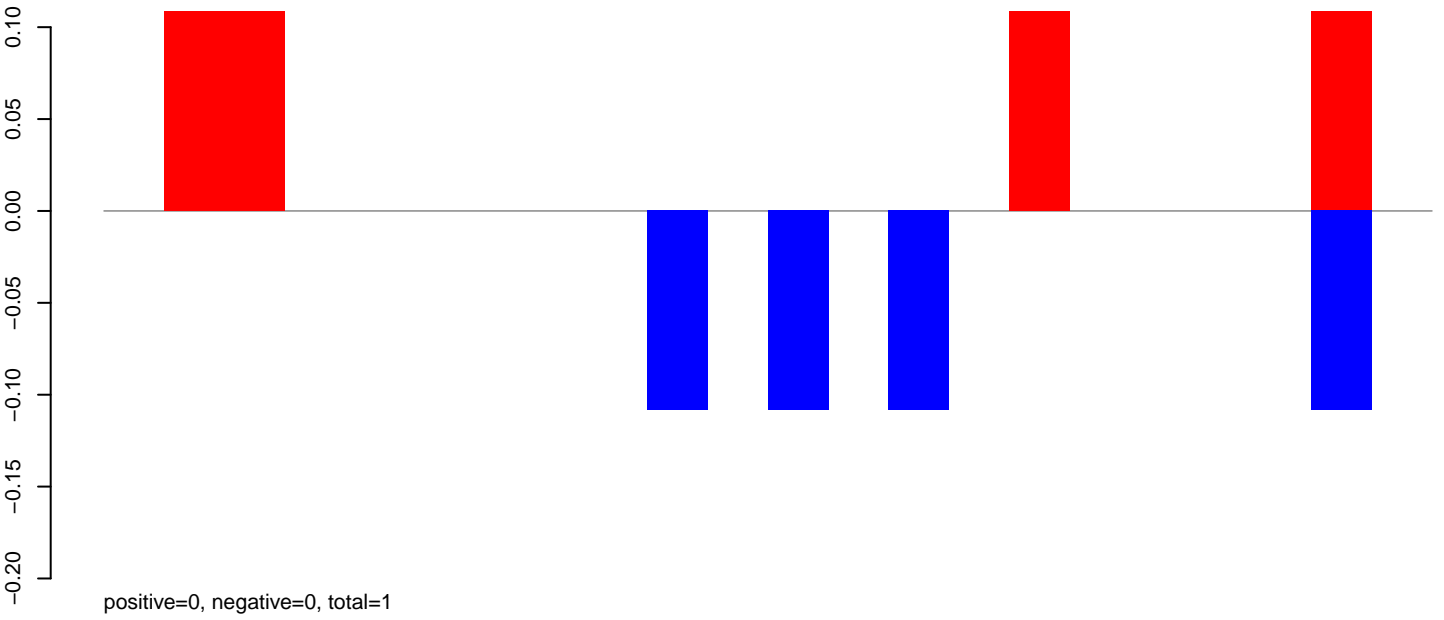
AnGam_Sua5b_cells.rep



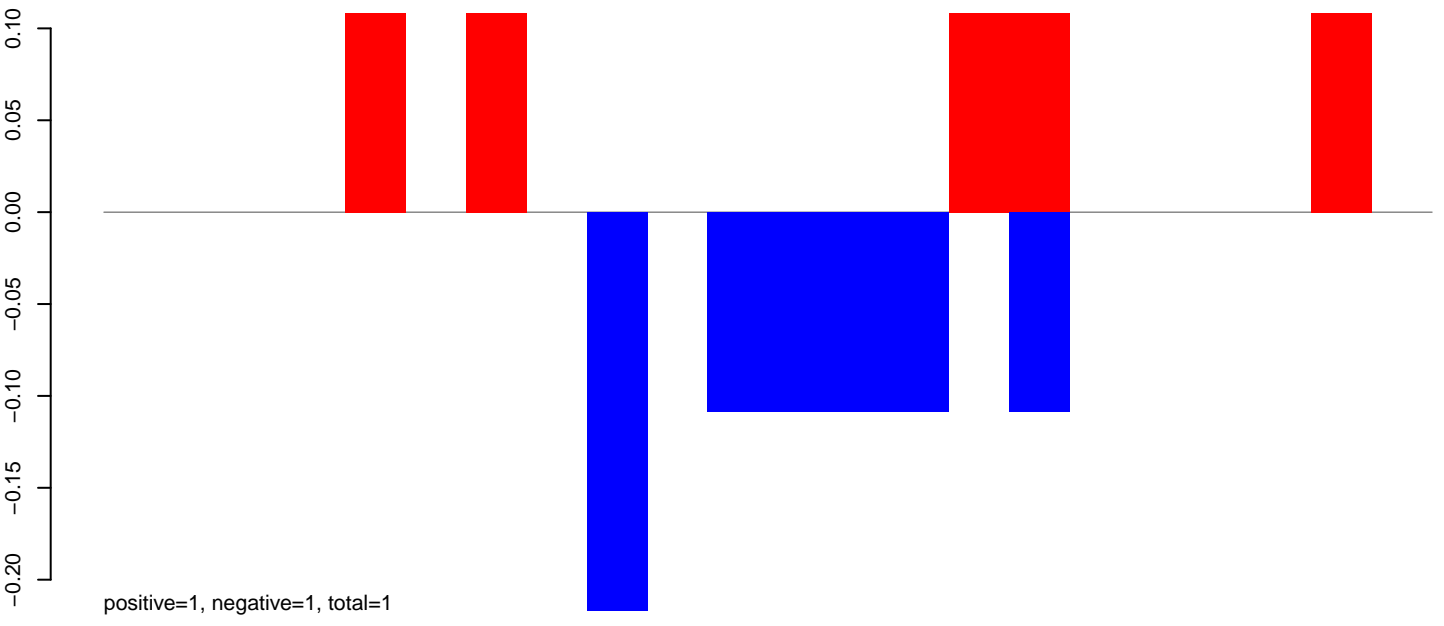
Window size=50, length=4007, TE@Ag-CR1-22:1-4007

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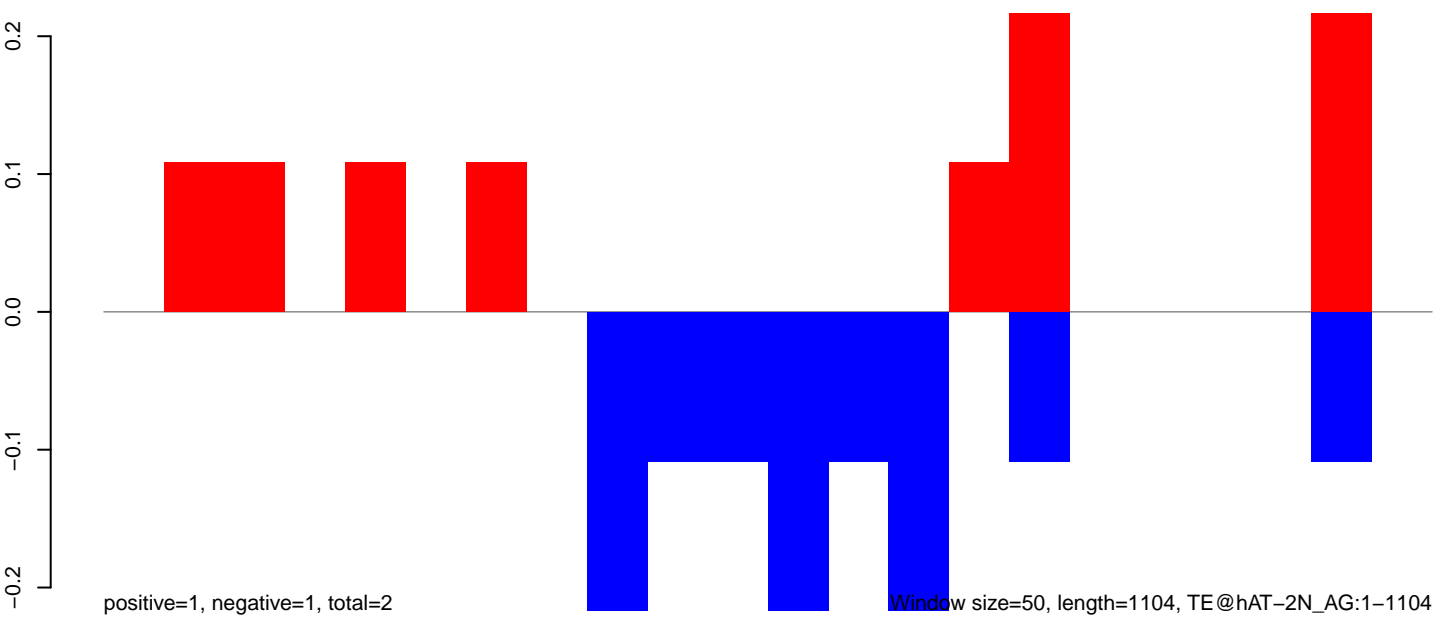
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

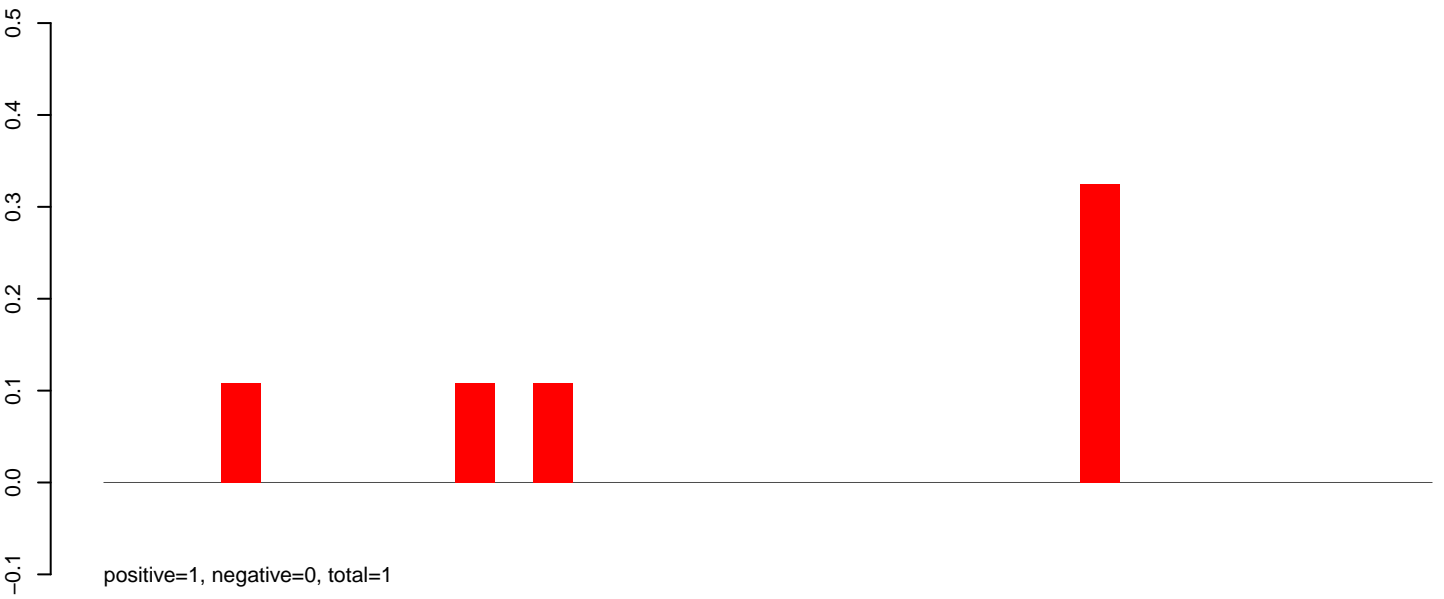


AnGam_Sua5b_cells.rep

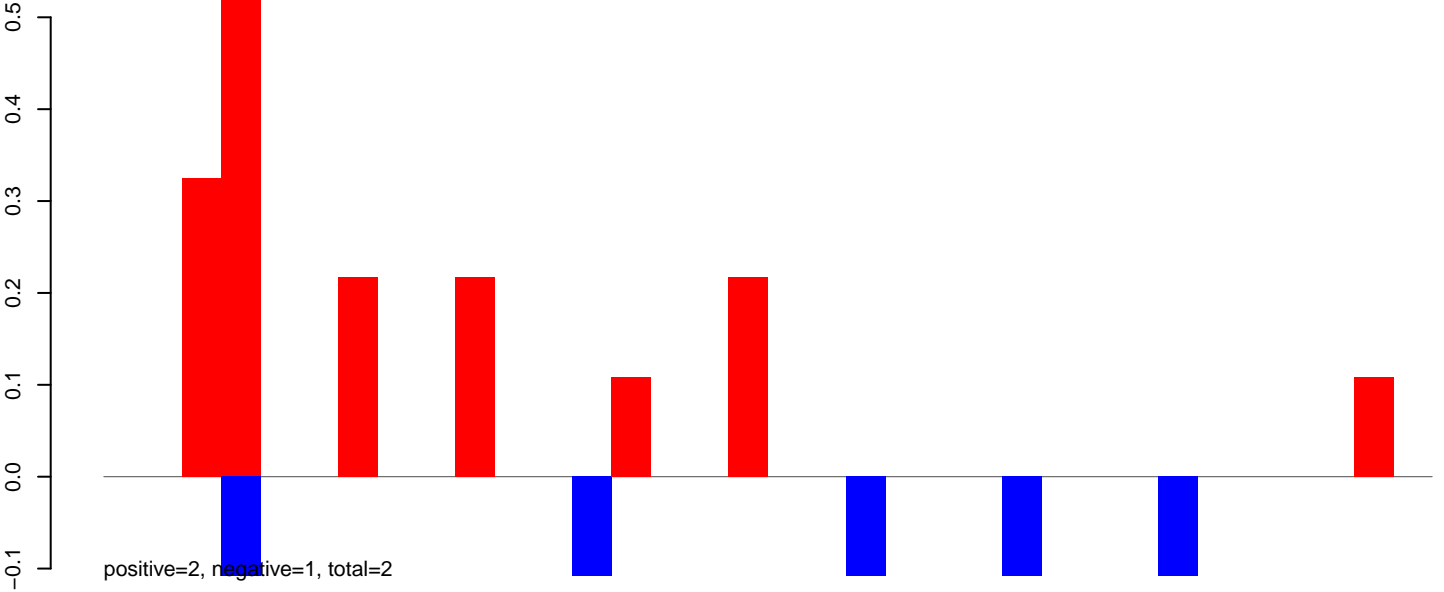


Window size=50, length=1104, TE@hAT-2N_AG:1-1104

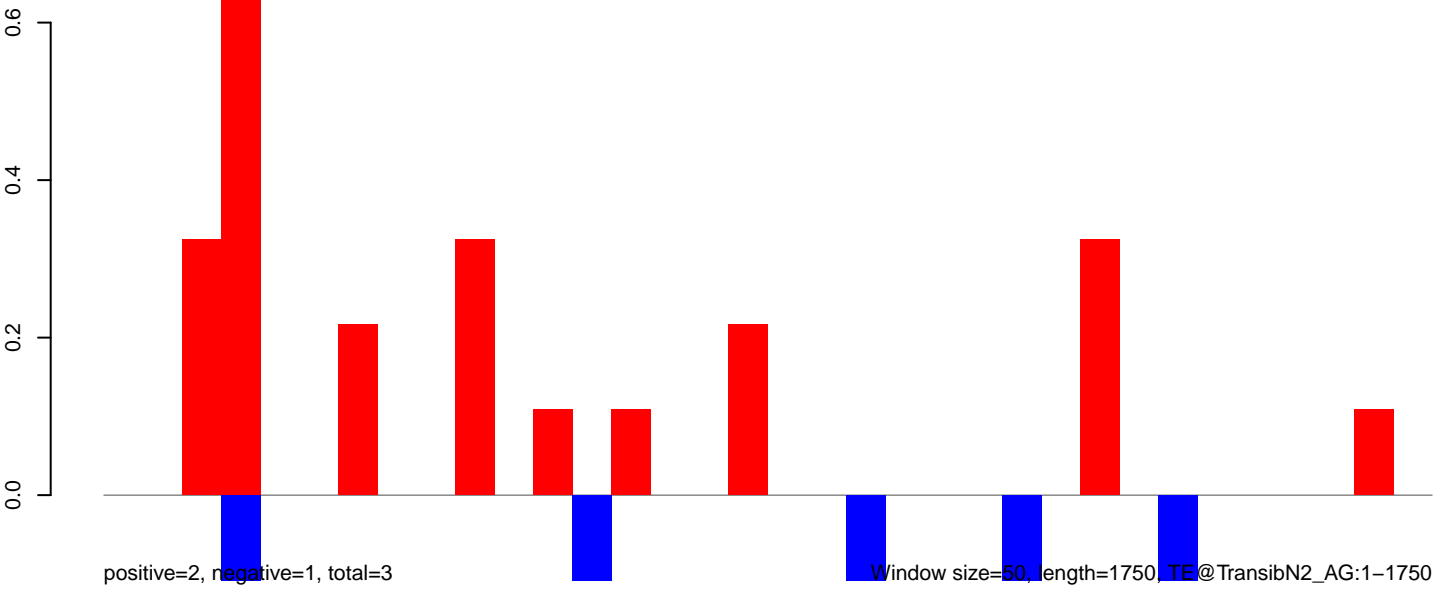
AnGam_Sua5b_cells.18_23.rep



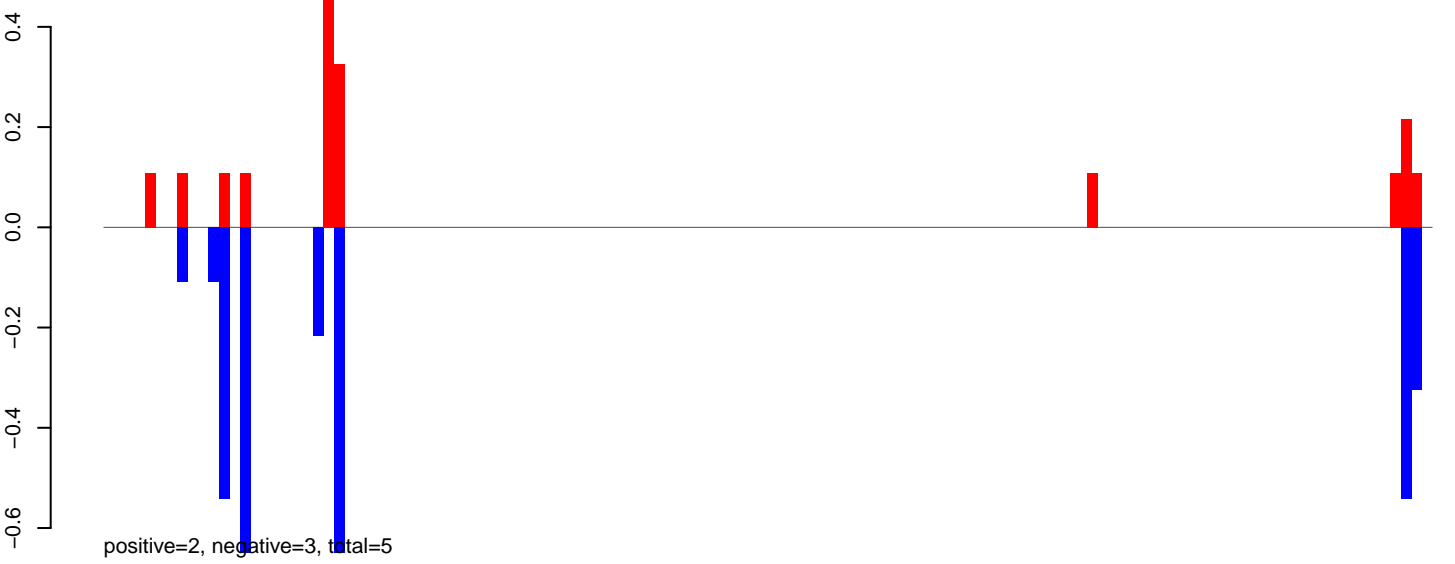
AnGam_Sua5b_cells.24_35.rep



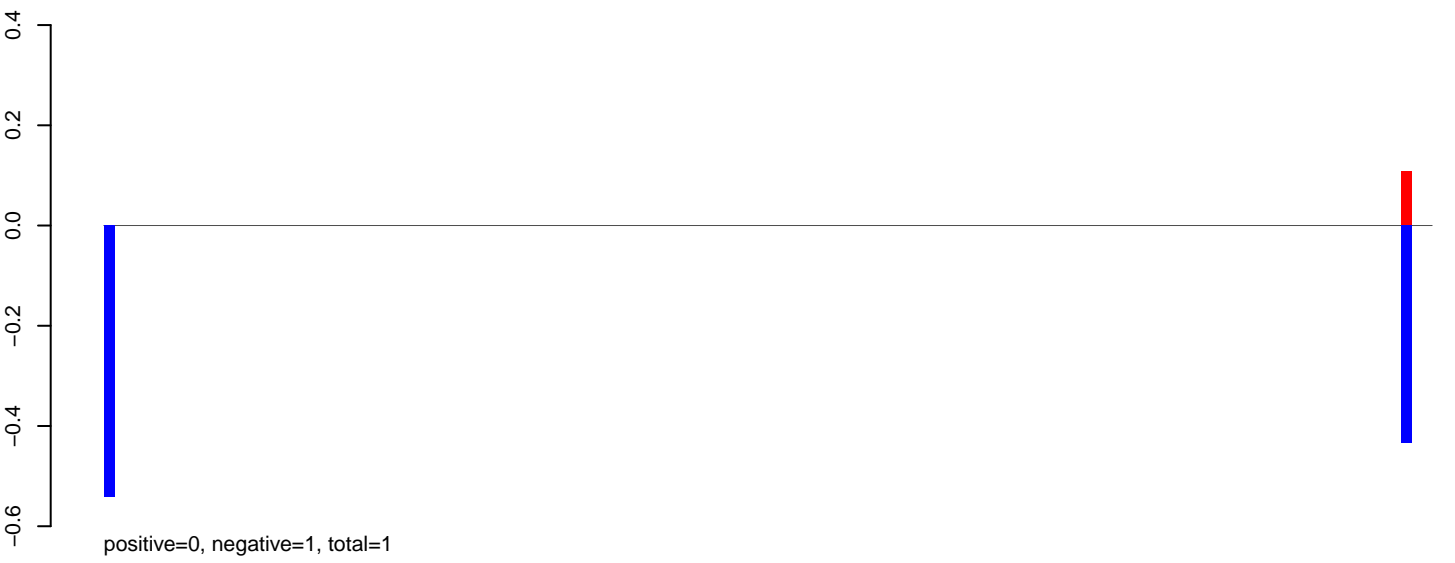
AnGam_Sua5b_cells.rep



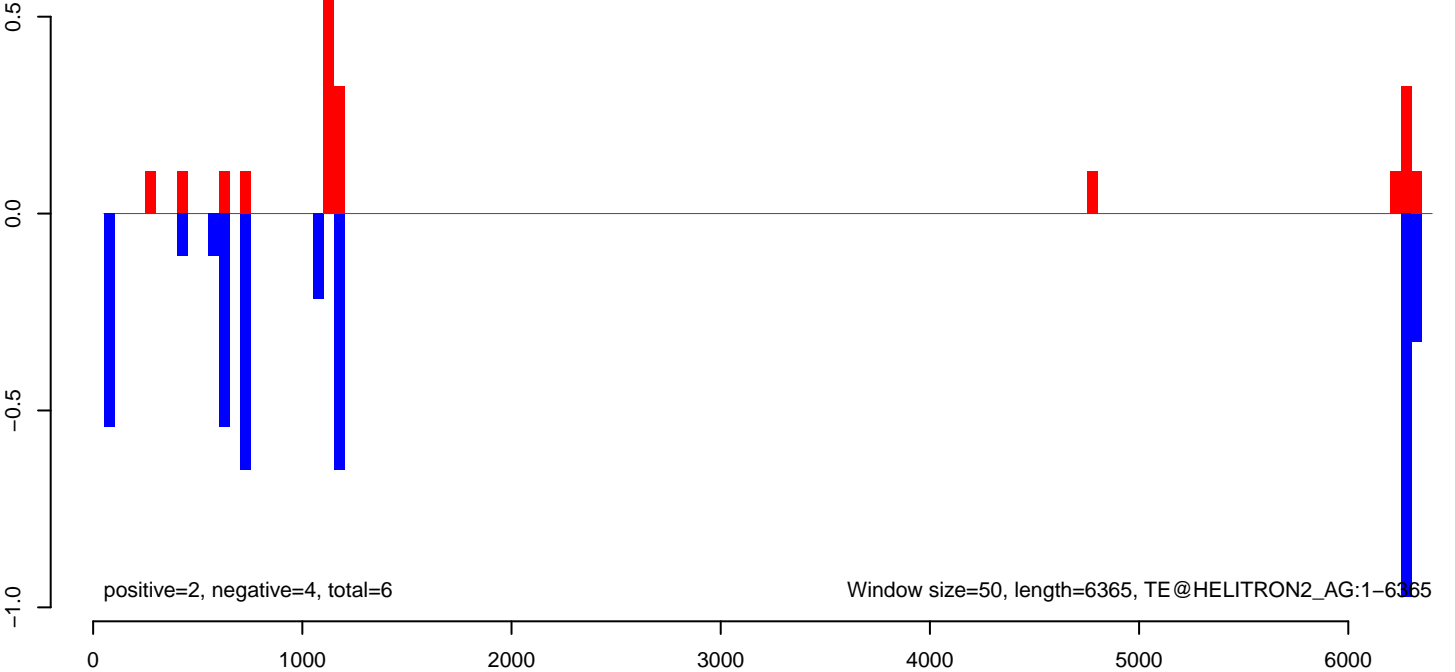
AnGam_Sua5b_cells.18_23.rep



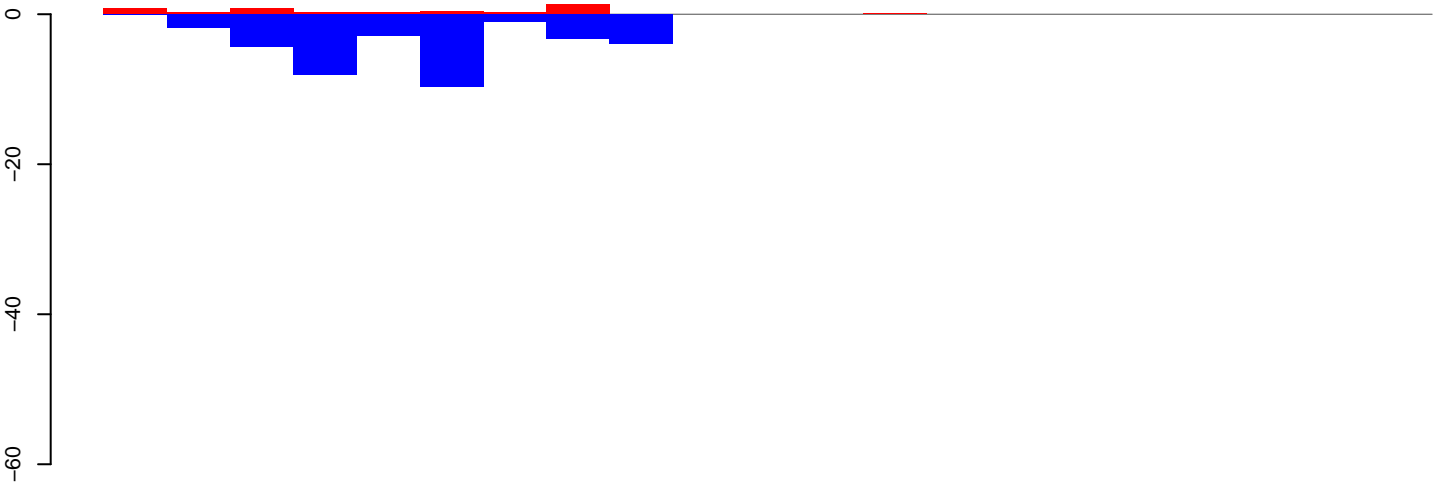
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

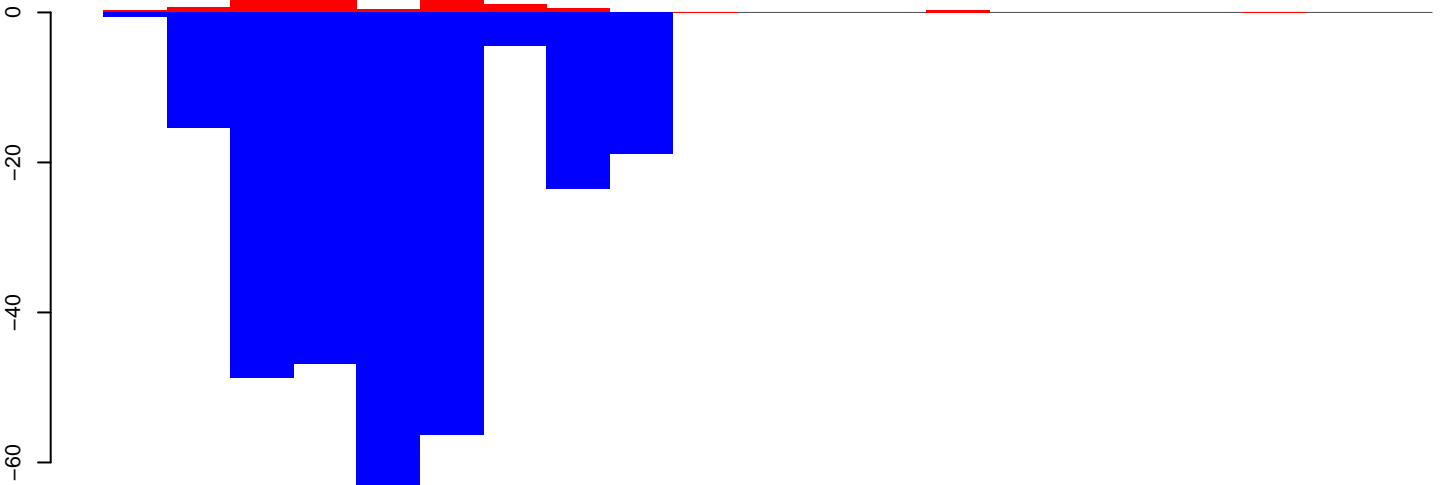


AnGam_Sua5b_cells.18_23.rep



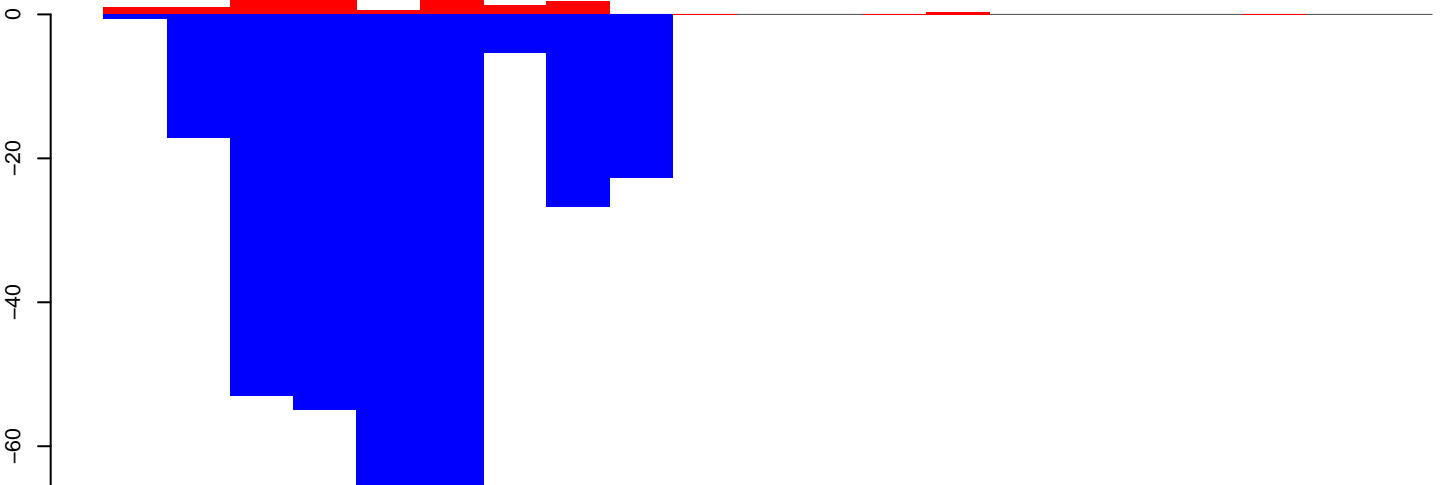
positive=4, negative=35, total=39

AnGam_Sua5b_cells.24_35.rep



positive=12, negative=290, total=301

AnGam_Sua5b_cells.rep



positive=16, negative=325, total=341

Window size=50, length=1086, TE@EAA02555-Unknown:1-1086

200

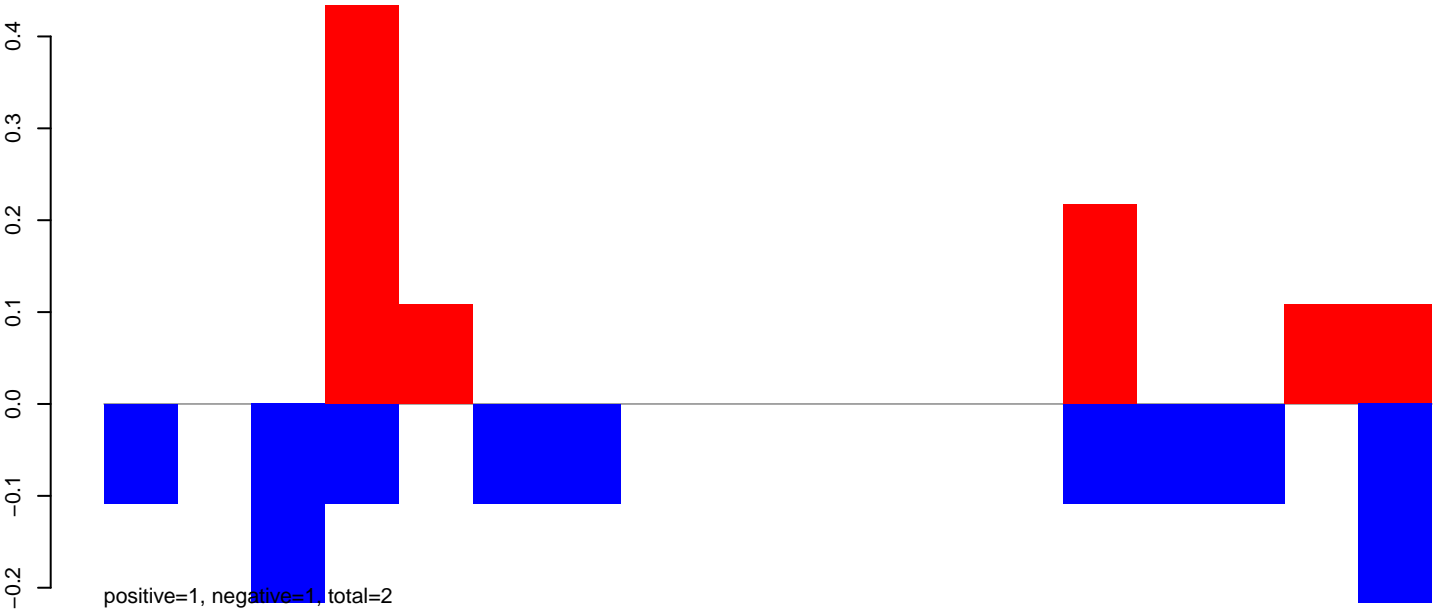
400

600

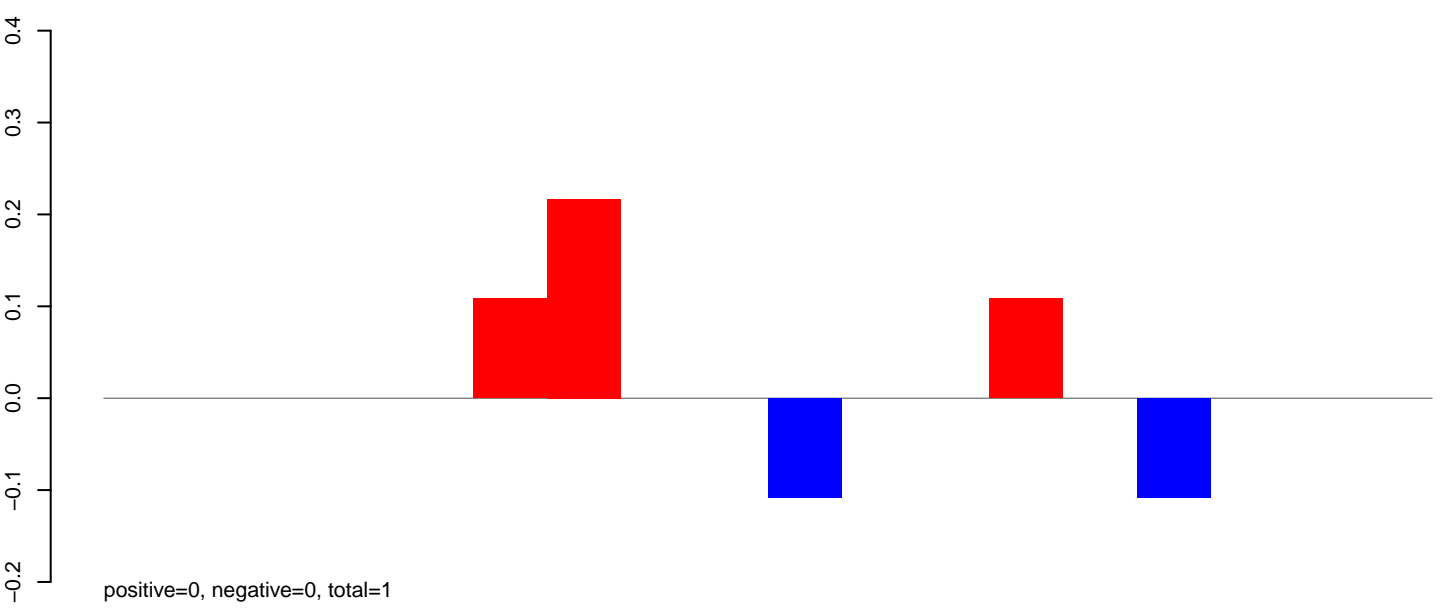
800

1000

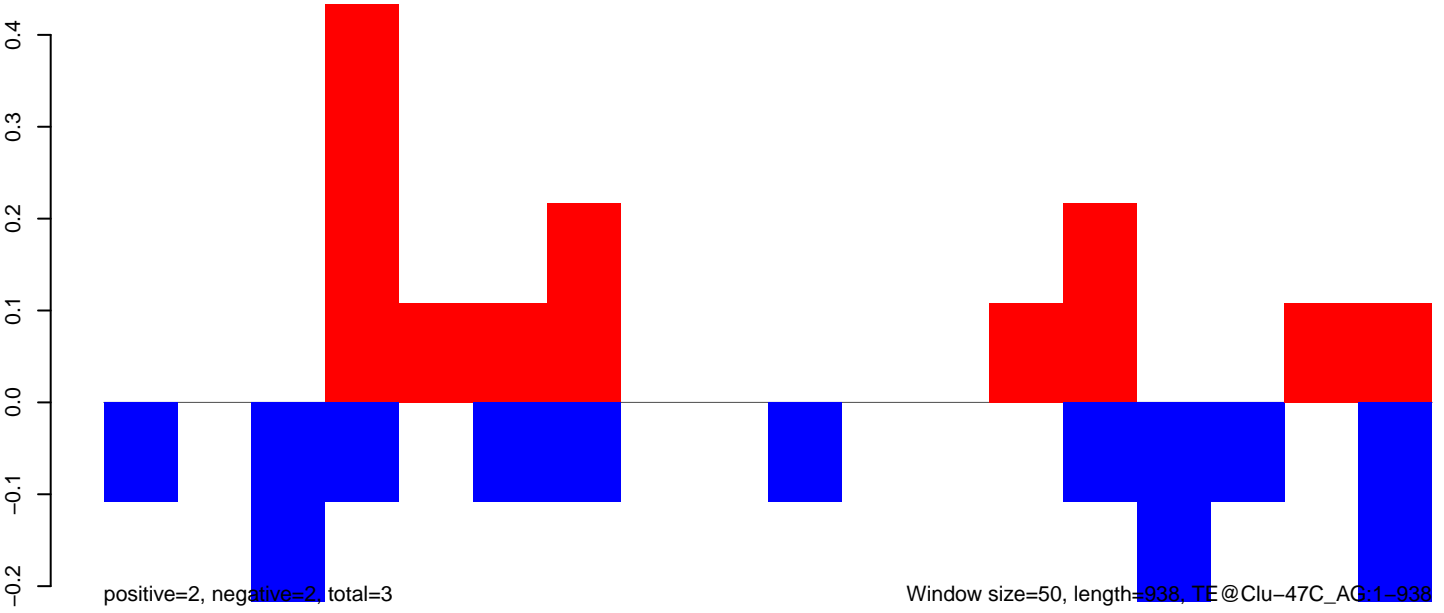
AnGam_Sua5b_cells.18_23.rep



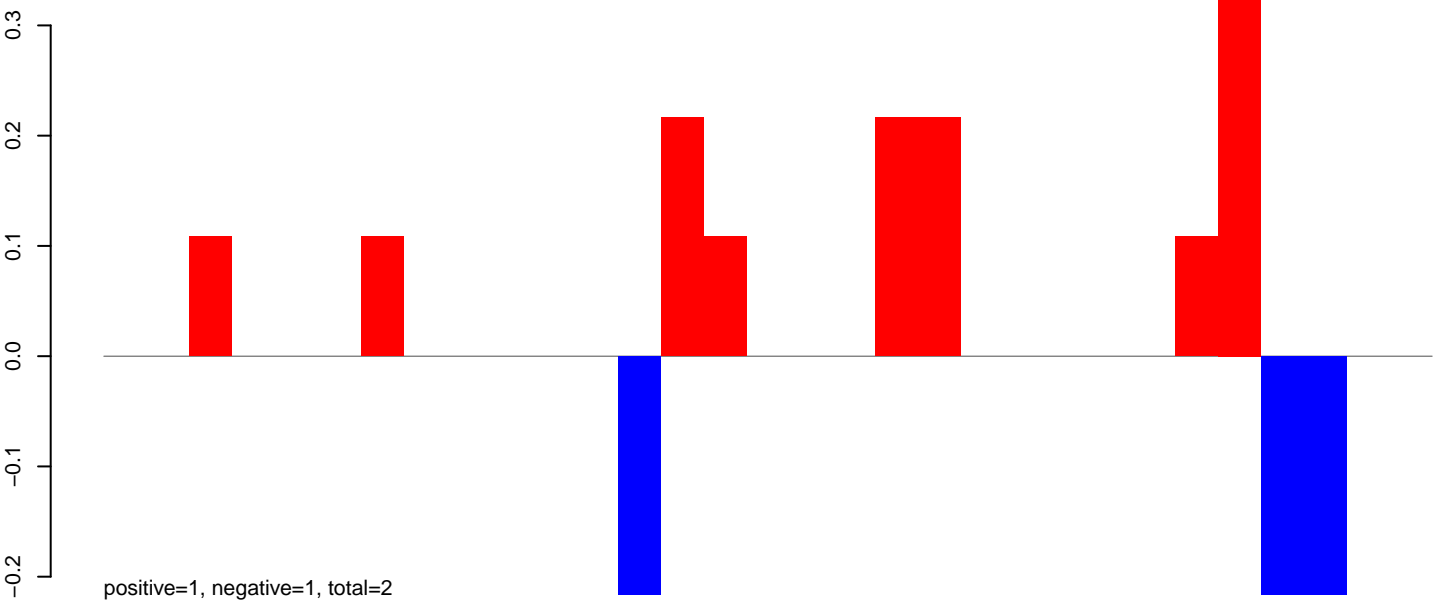
AnGam_Sua5b_cells.24_35.rep



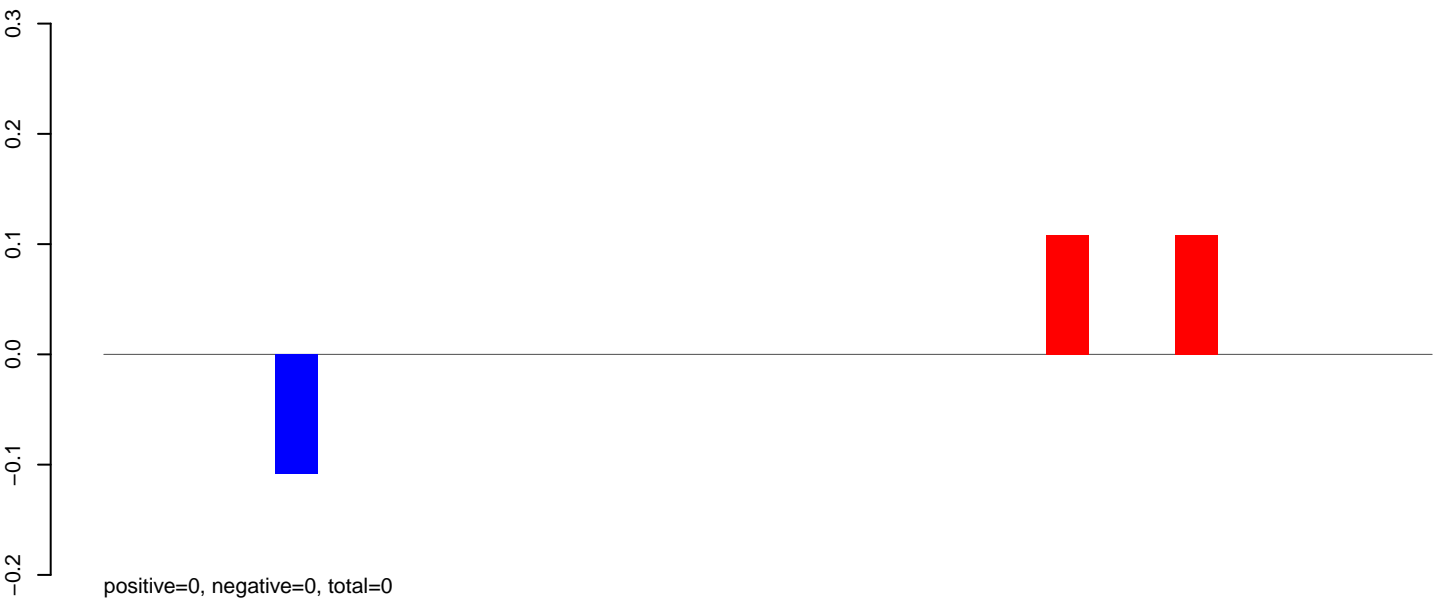
AnGam_Sua5b_cells.rep



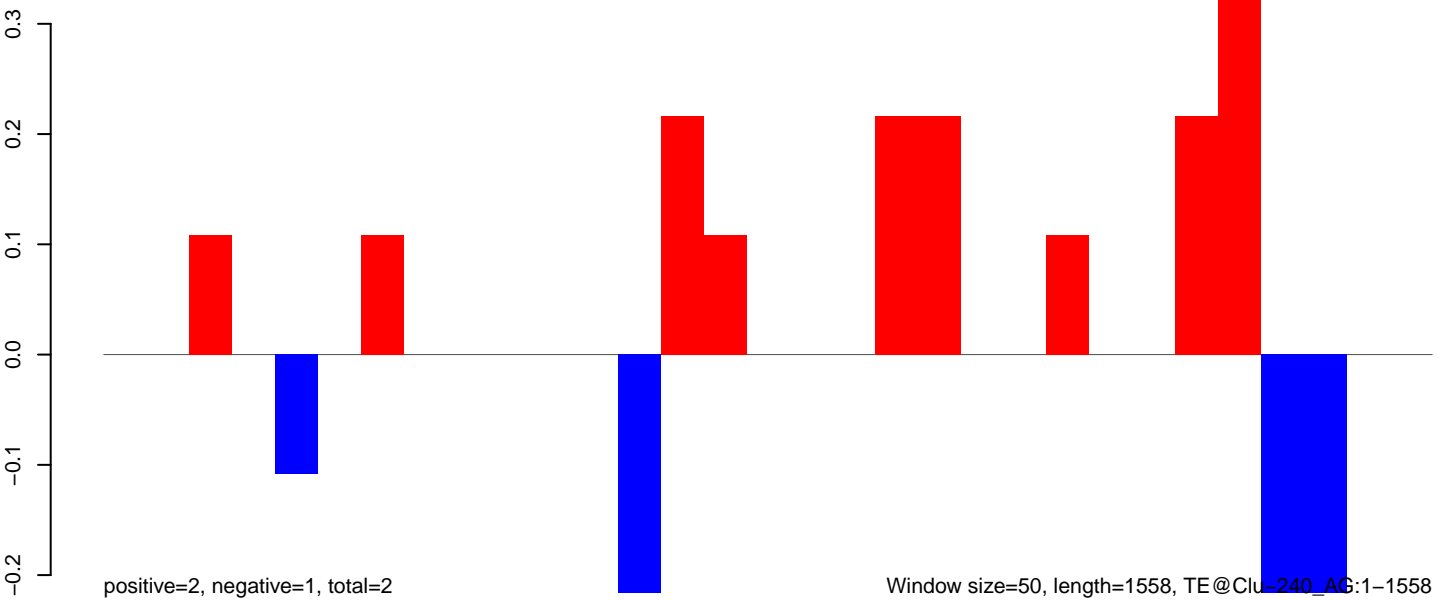
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

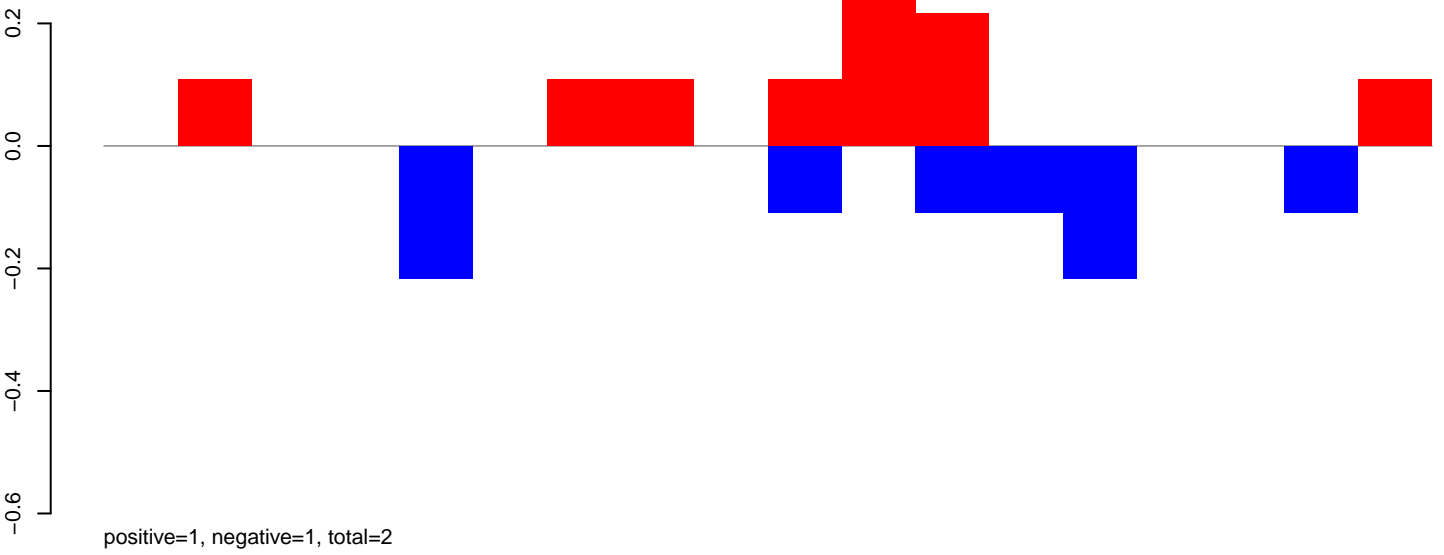


AnGam_Sua5b_cells.rep

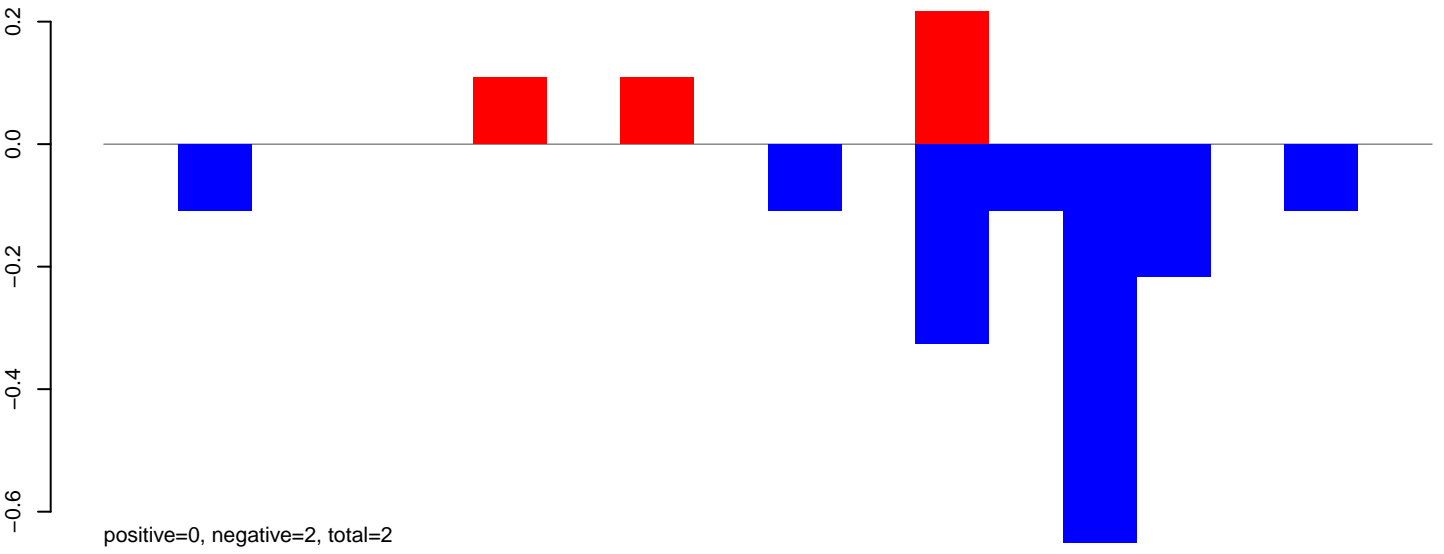


Window size=50, length=1558, TE@Clu_240_AG:1-1558

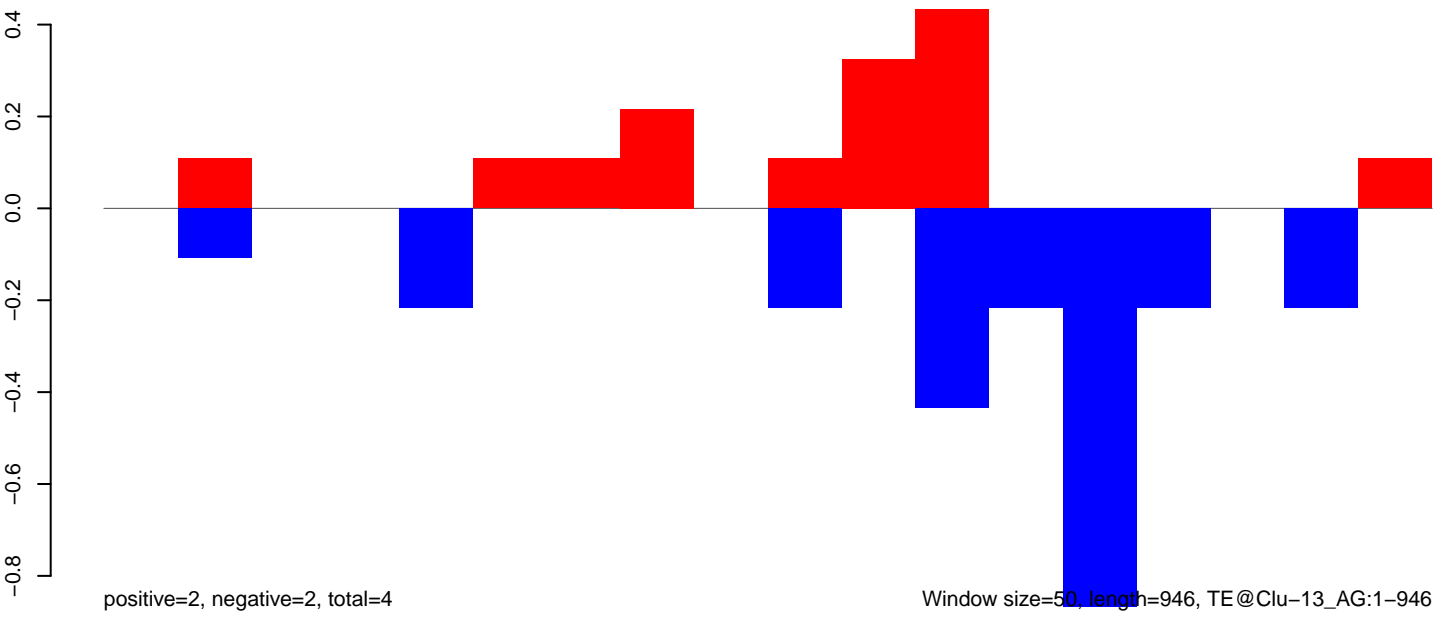
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

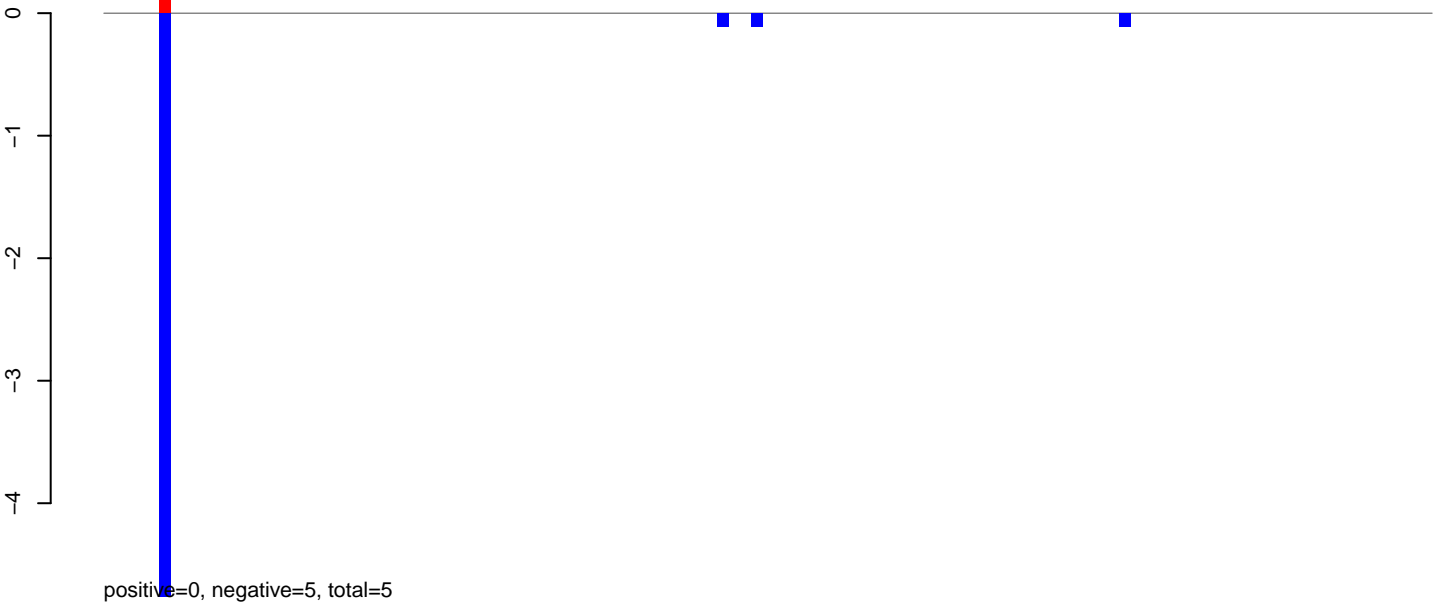


AnGam_Sua5b_cells.rep

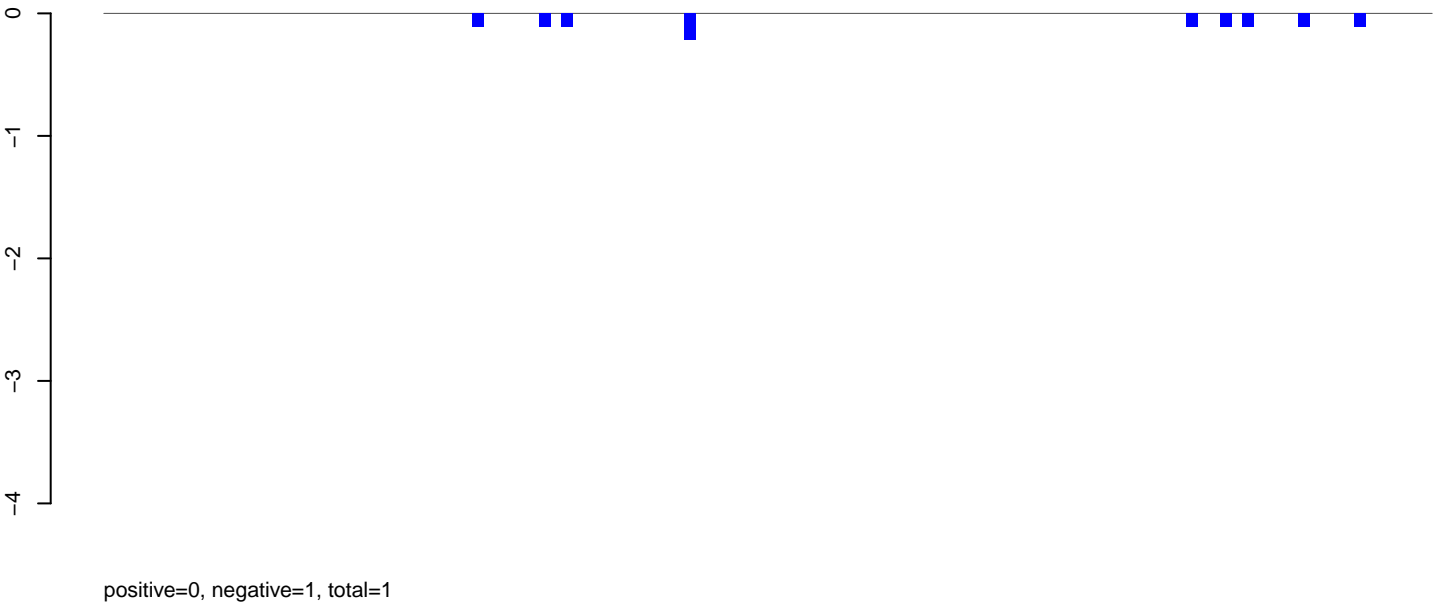


Window size=50, length=946, TE@Clu-13_AG:1-946

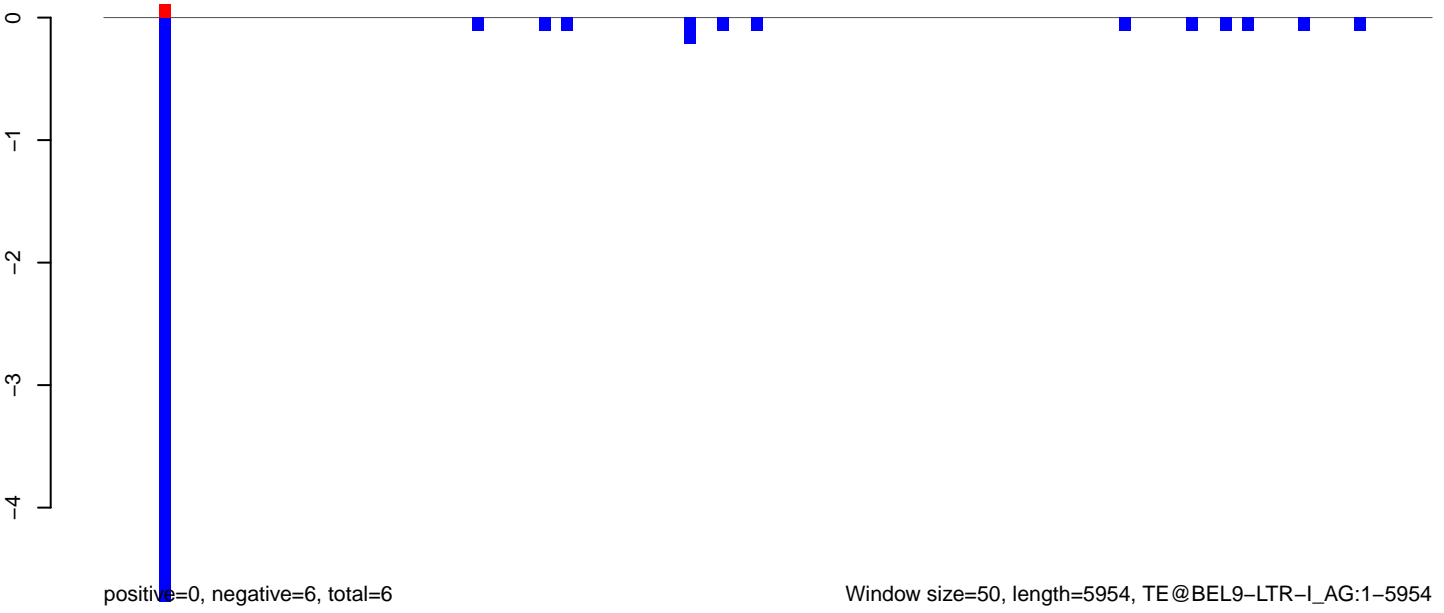
AnGam_Sua5b_cells.18_23.rep



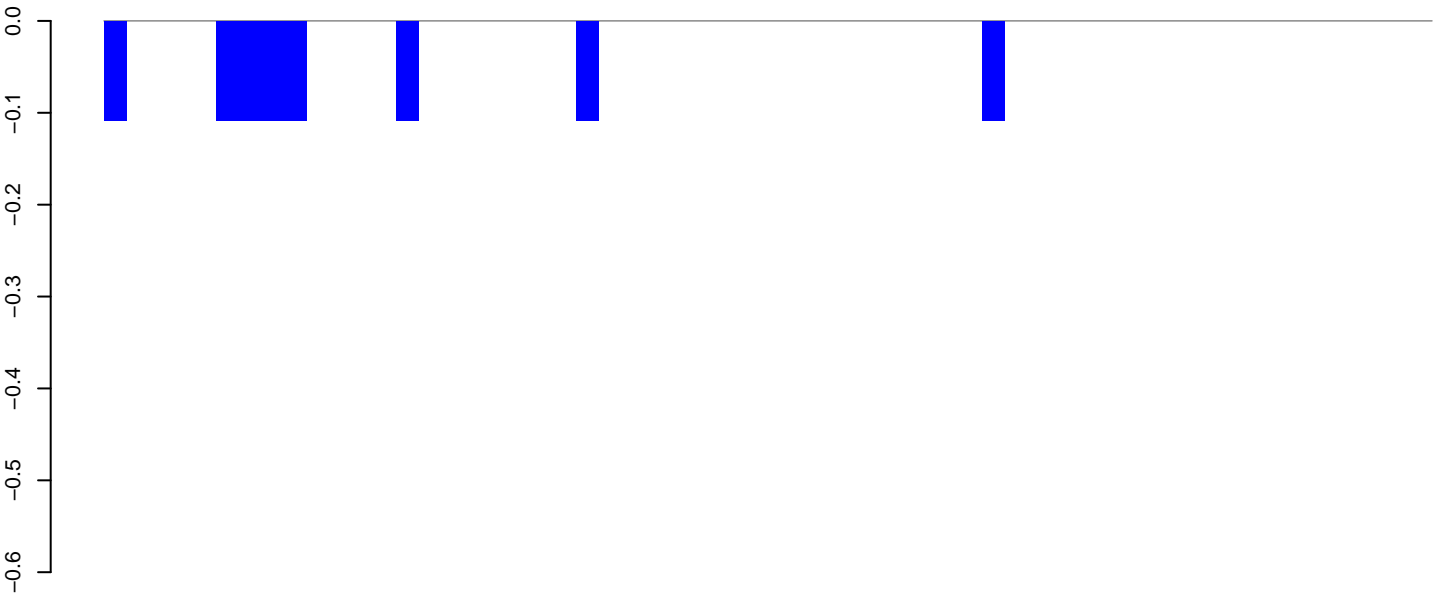
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

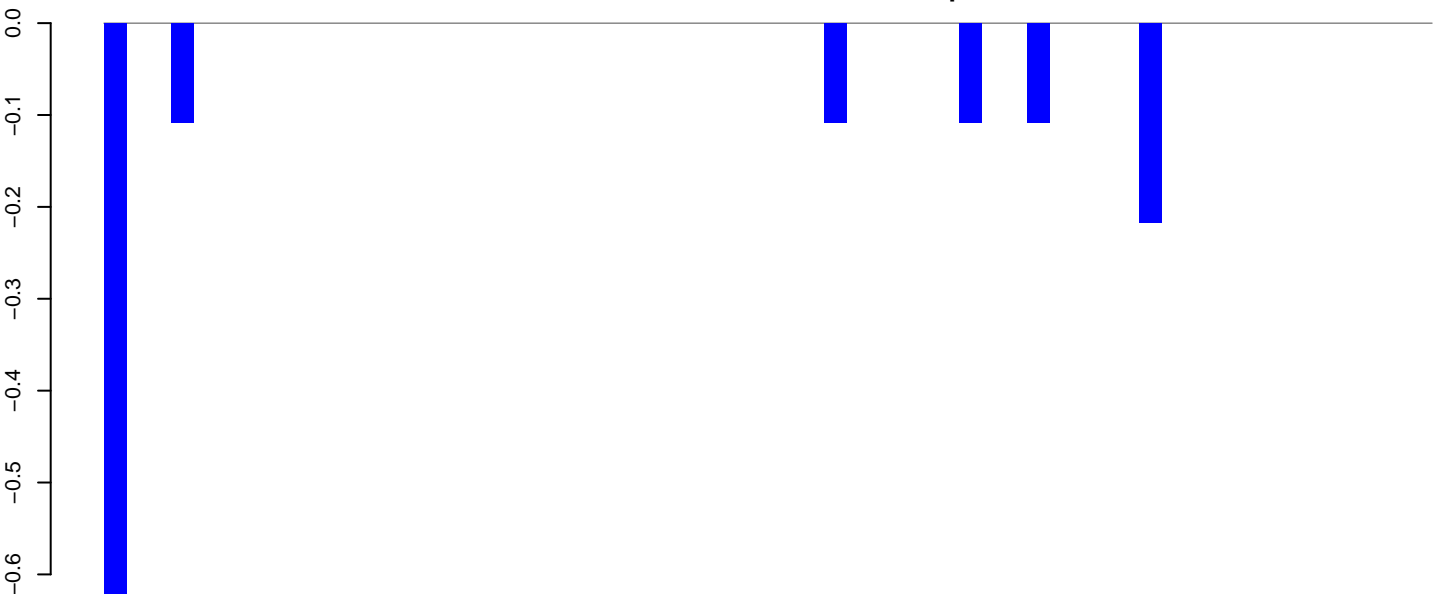


AnGam_Sua5b_cells.18_23.rep



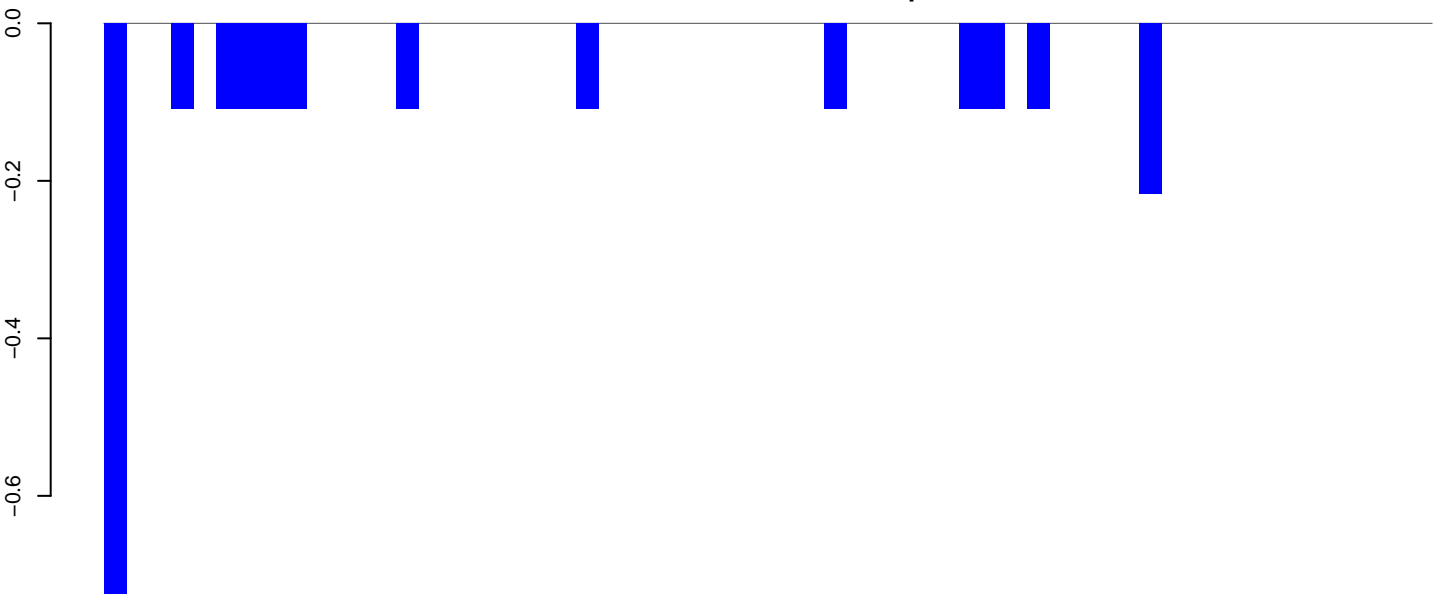
positive=0, negative=1, total=1

AnGam_Sua5b_cells.24_35.rep



positive=0, negative=1, total=1

AnGam_Sua5b_cells.rep

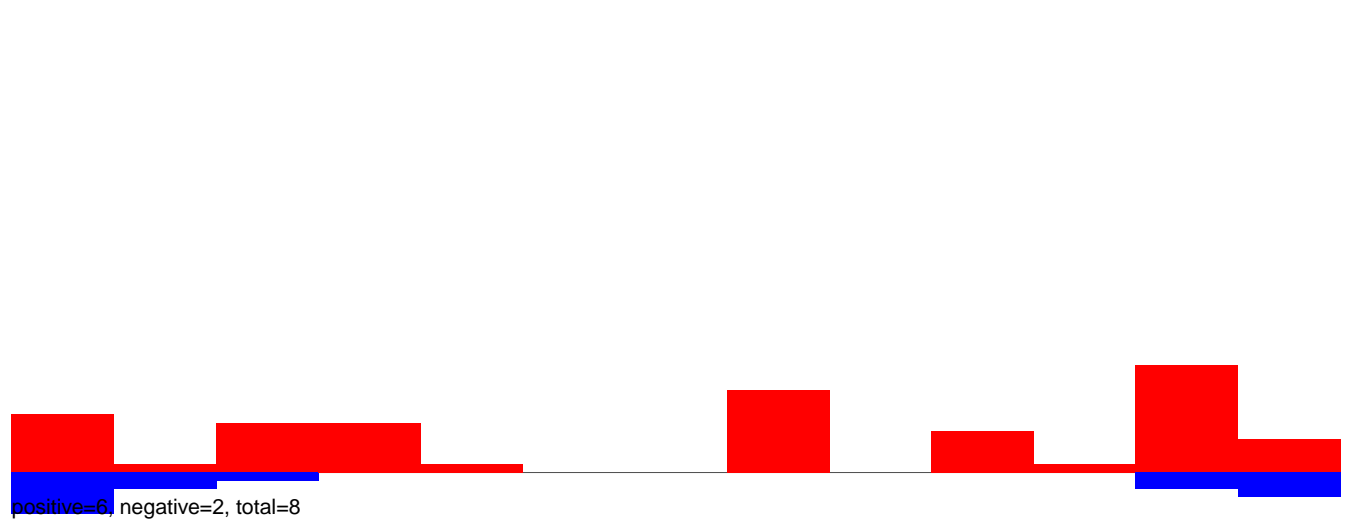


positive=0, negative=2, total=2

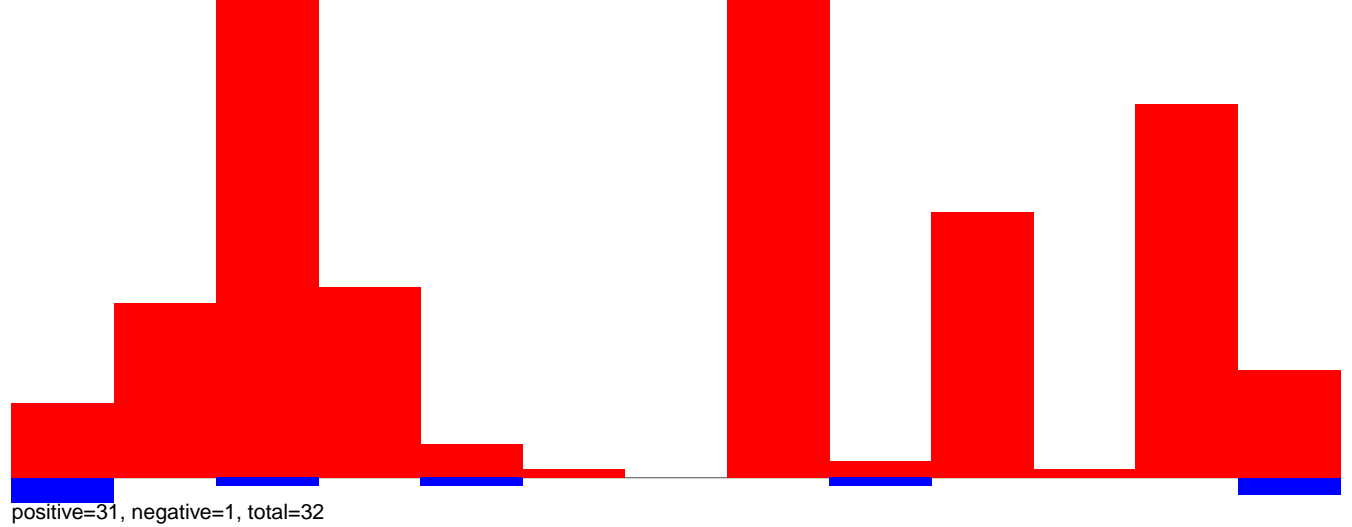
Window size=50, length=2969, TE@BEL-25_AG-LTR-I:1-2969

0 500 1000 1500 2000 2500 3000

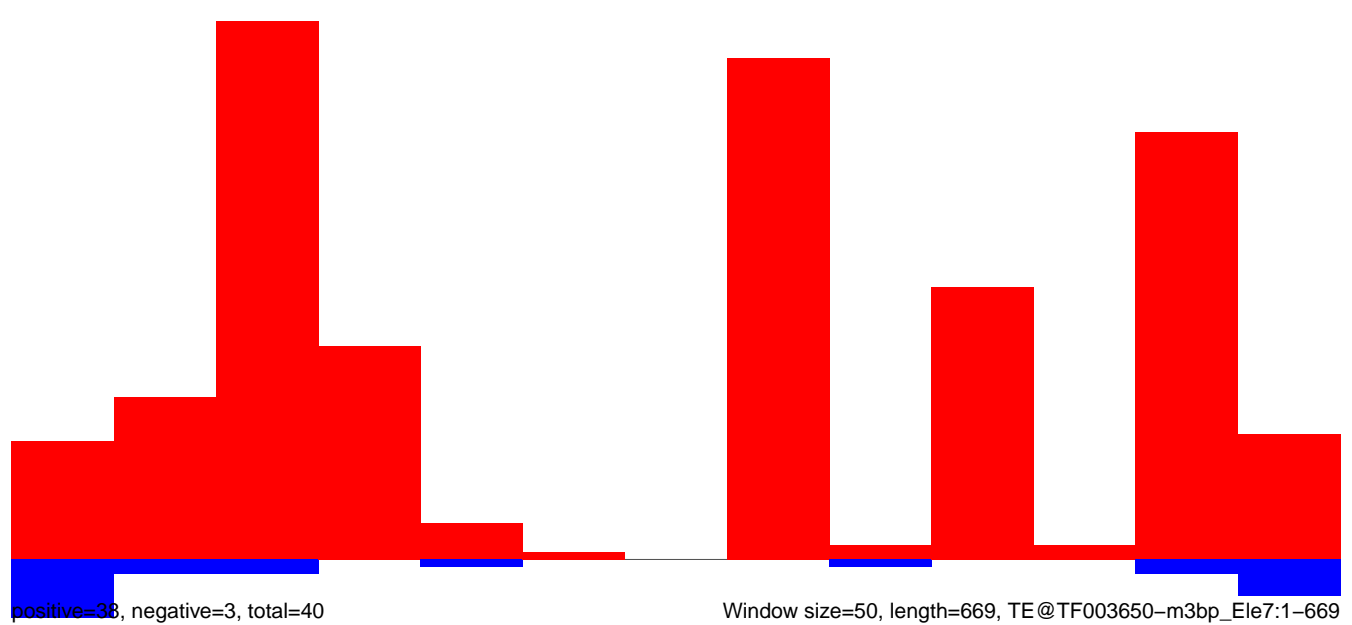
AnGam_Sua5b_cells.18_23.rep



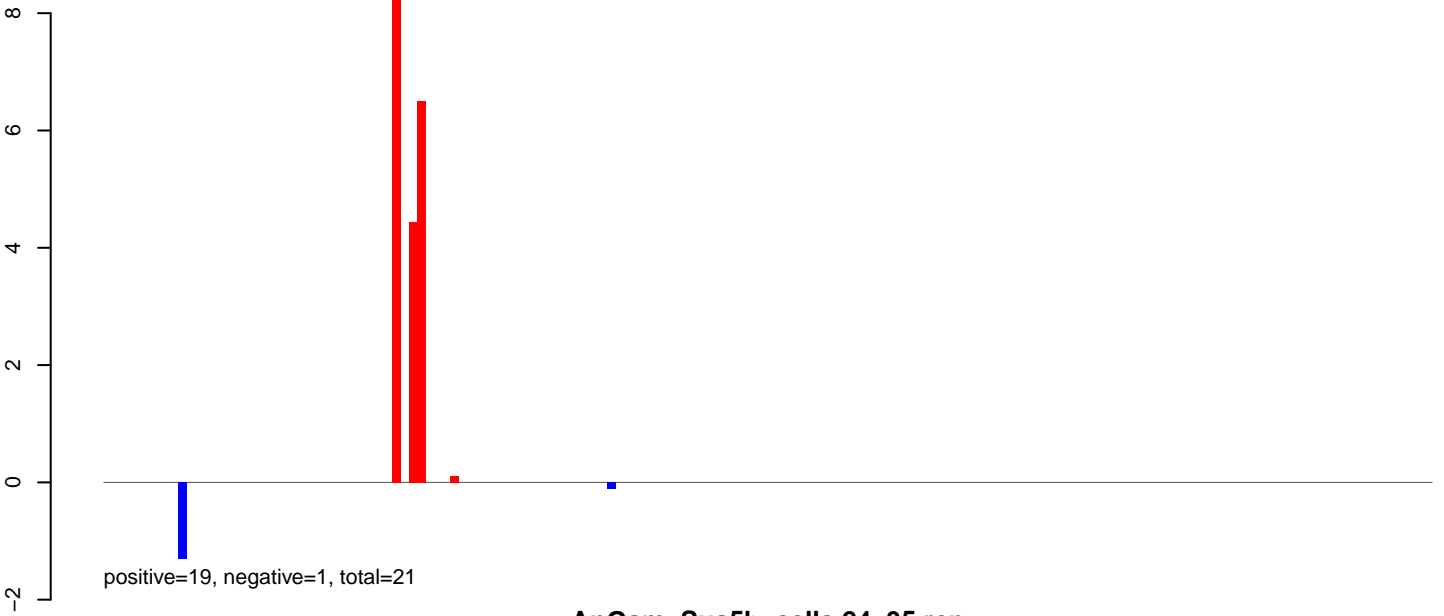
AnGam_Sua5b_cells.24_35.rep



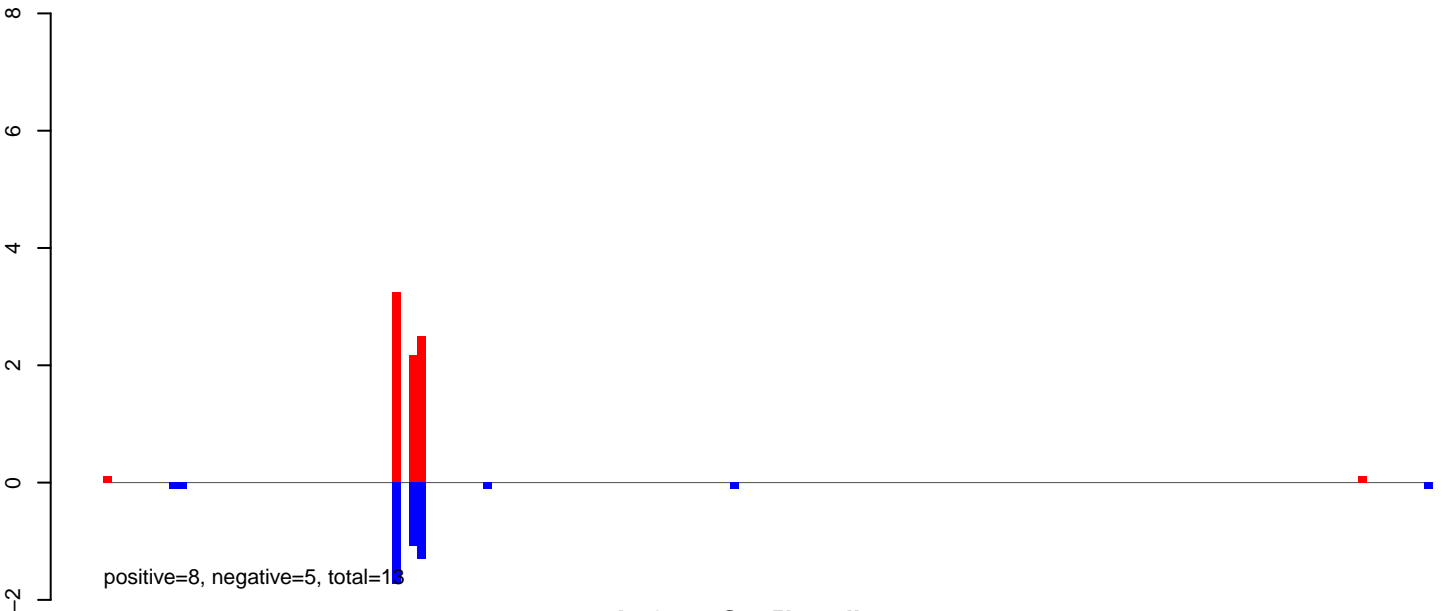
AnGam_Sua5b_cells.rep



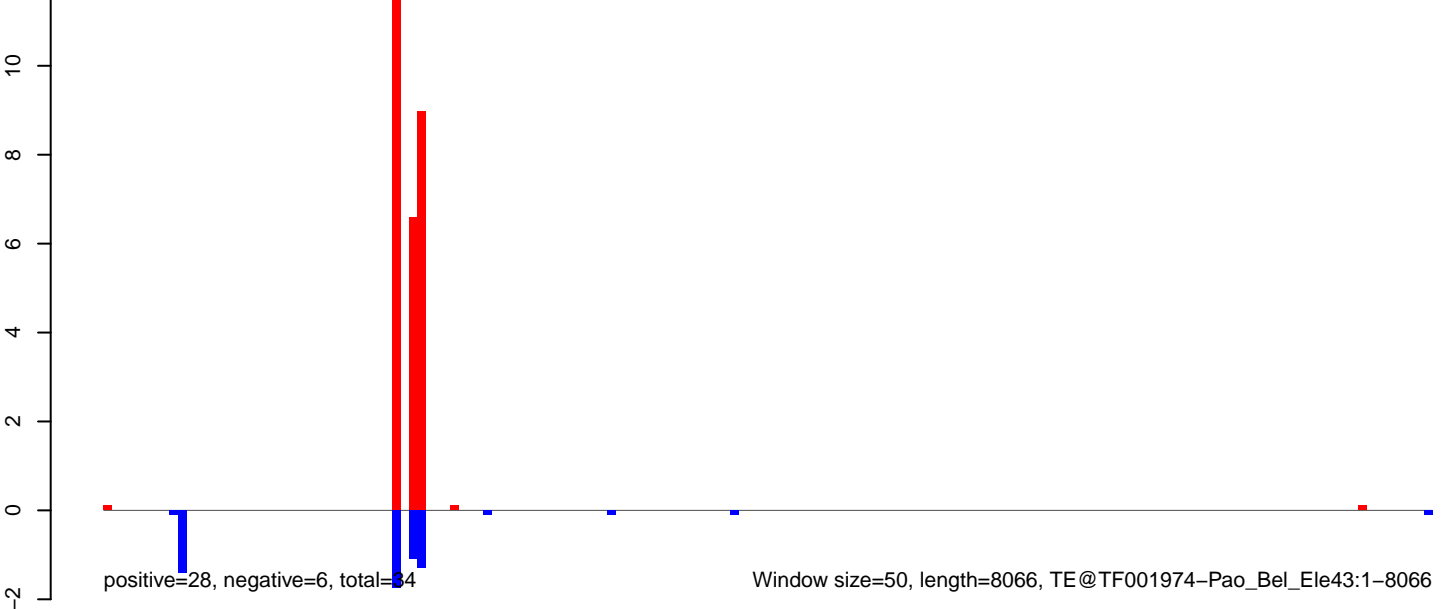
AnGam_Sua5b_cells.18_23.rep



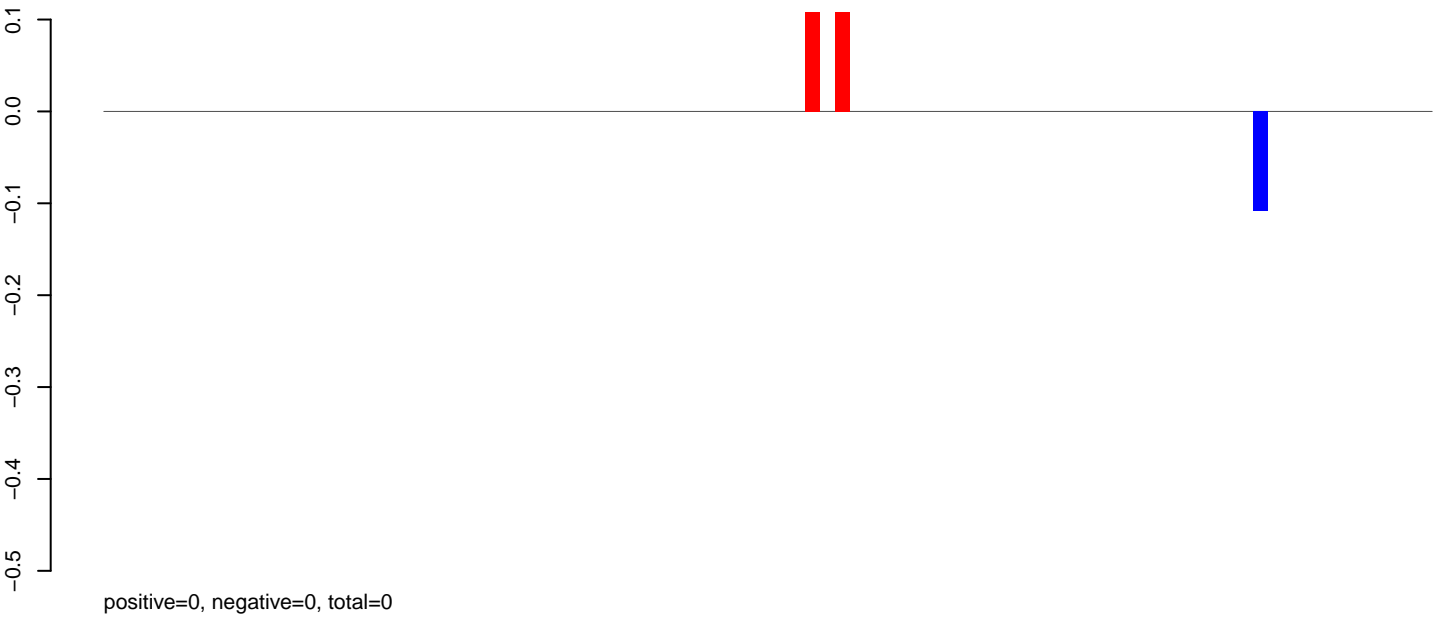
AnGam_Sua5b_cells.24_35.rep



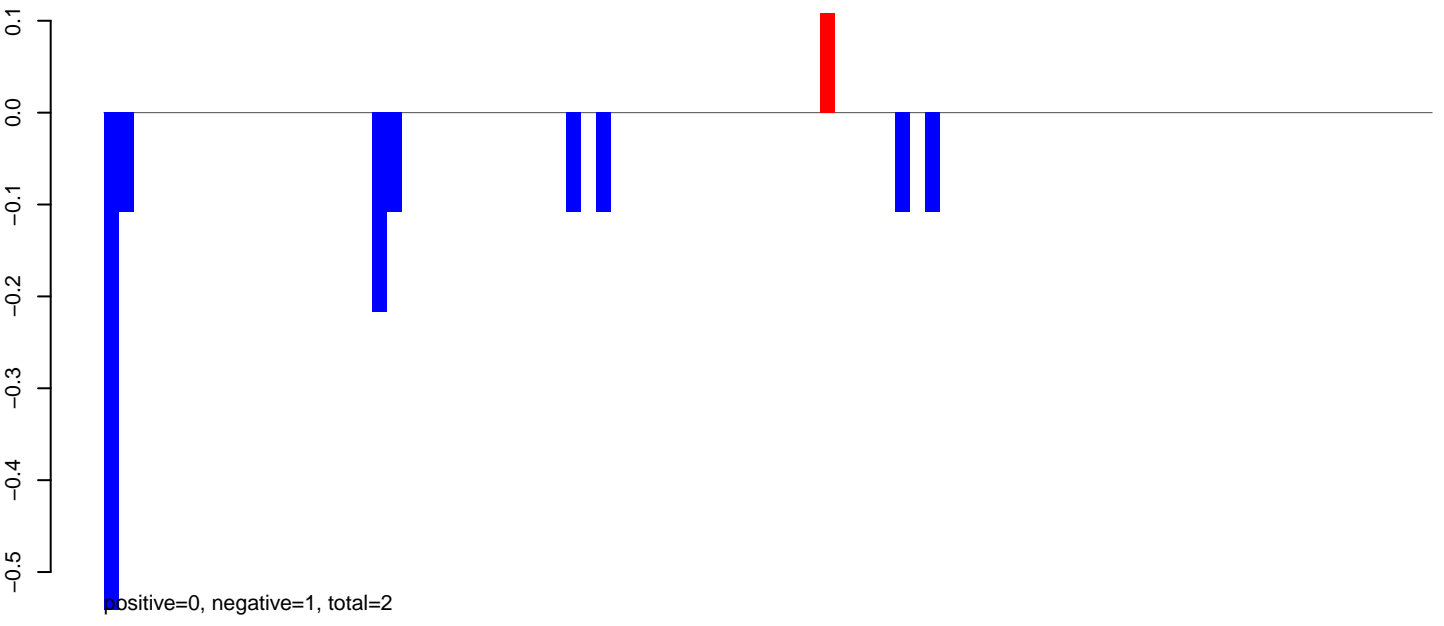
AnGam_Sua5b_cells.rep



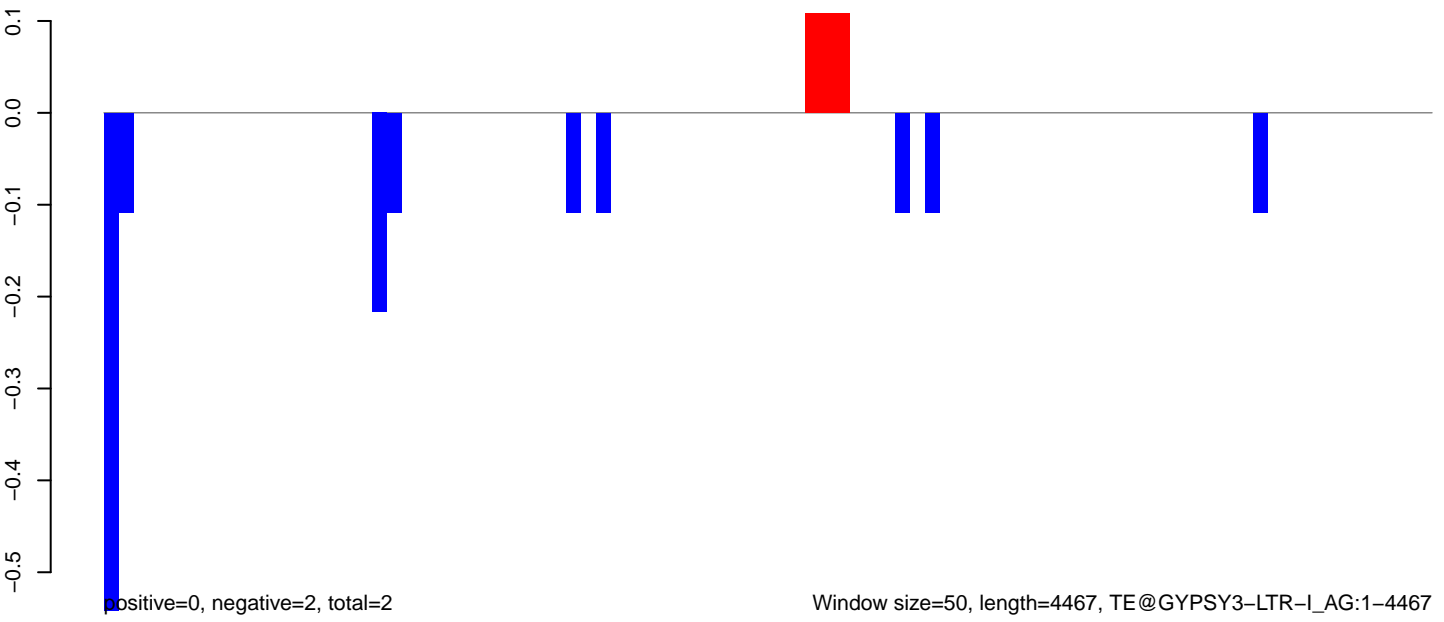
AnGam_Sua5b_cells.18_23.rep



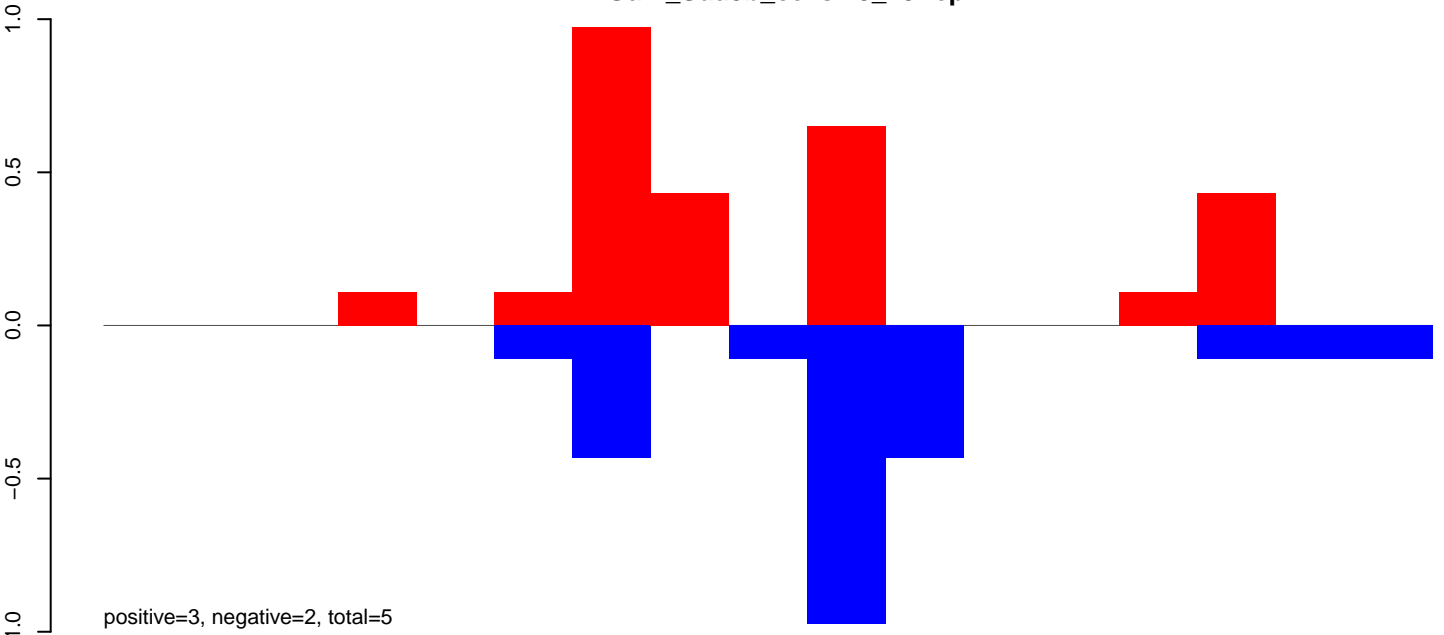
AnGam_Sua5b_cells.24_35.rep



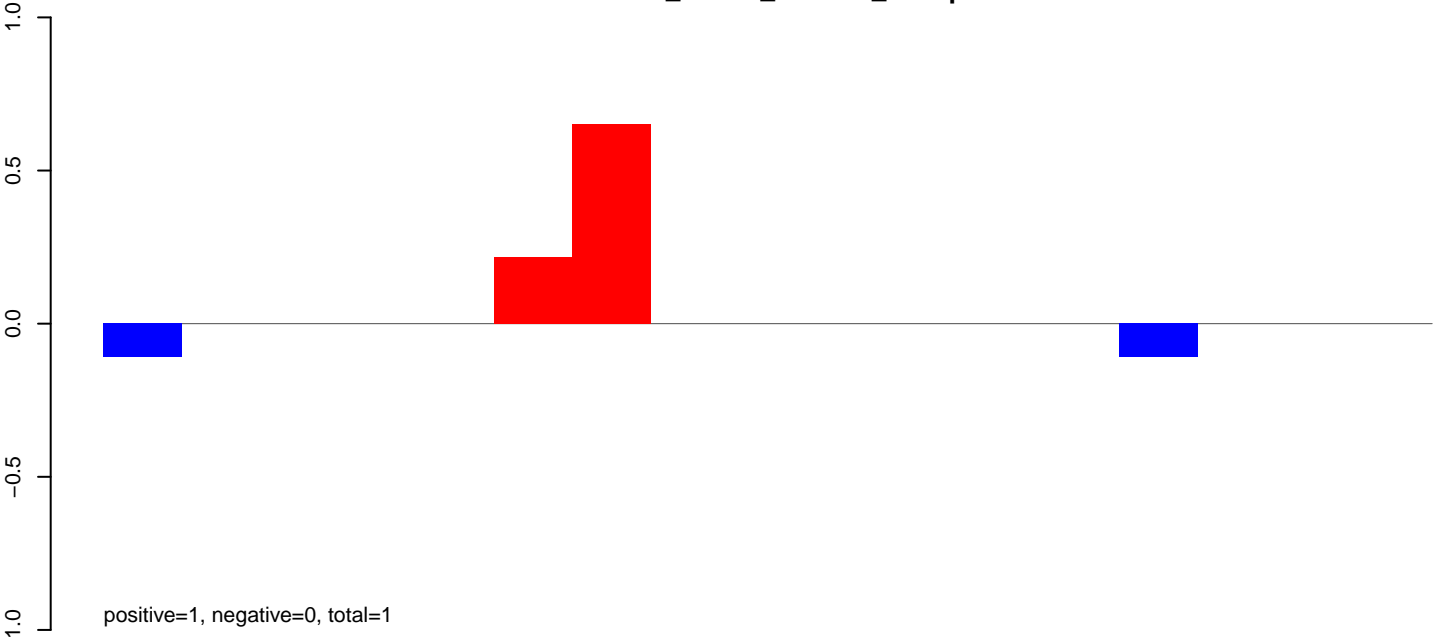
AnGam_Sua5b_cells.rep



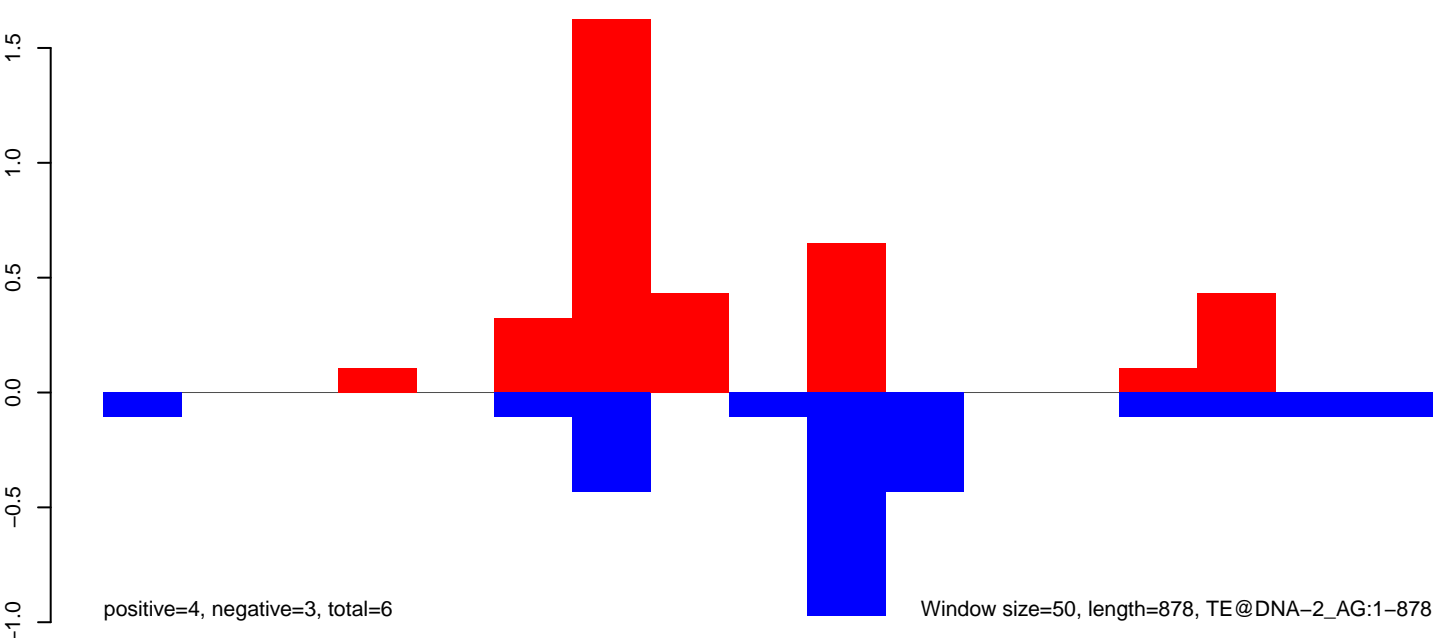
AnGam_Sua5b_cells.18_23.rep



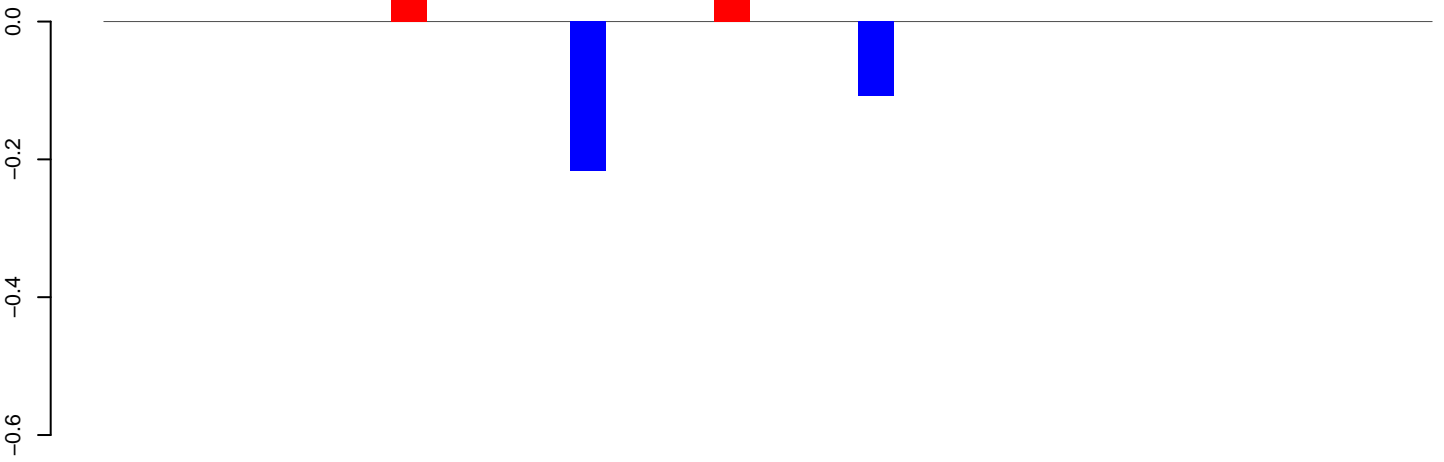
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

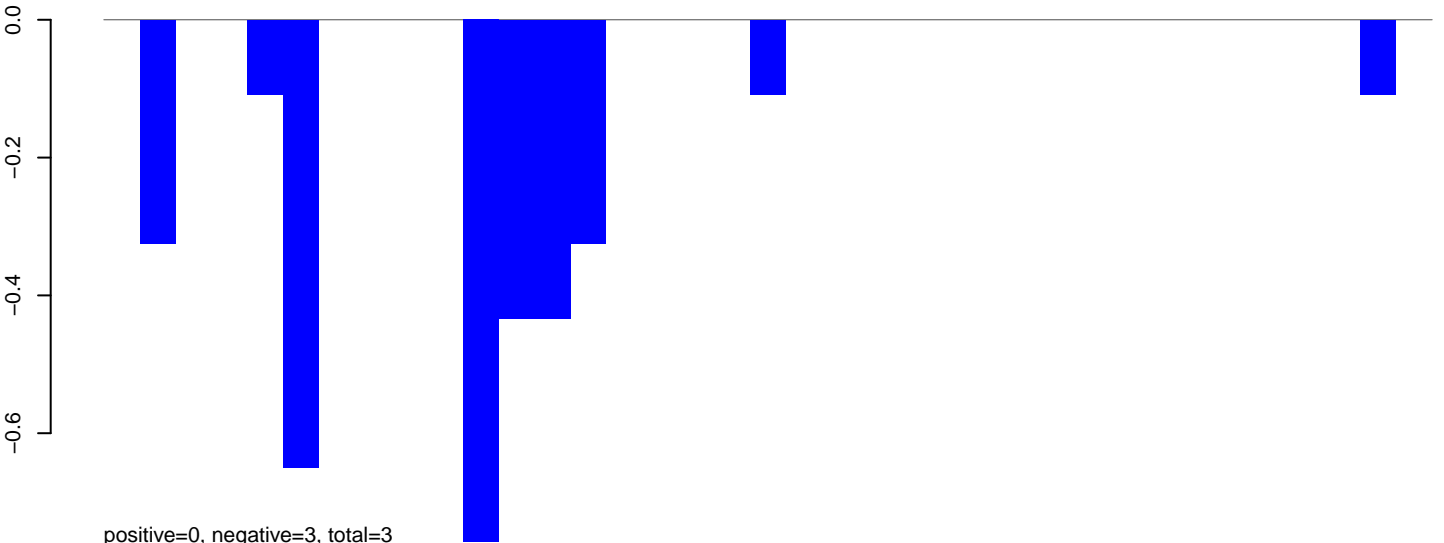


AnGam_Sua5b_cells.18_23.rep



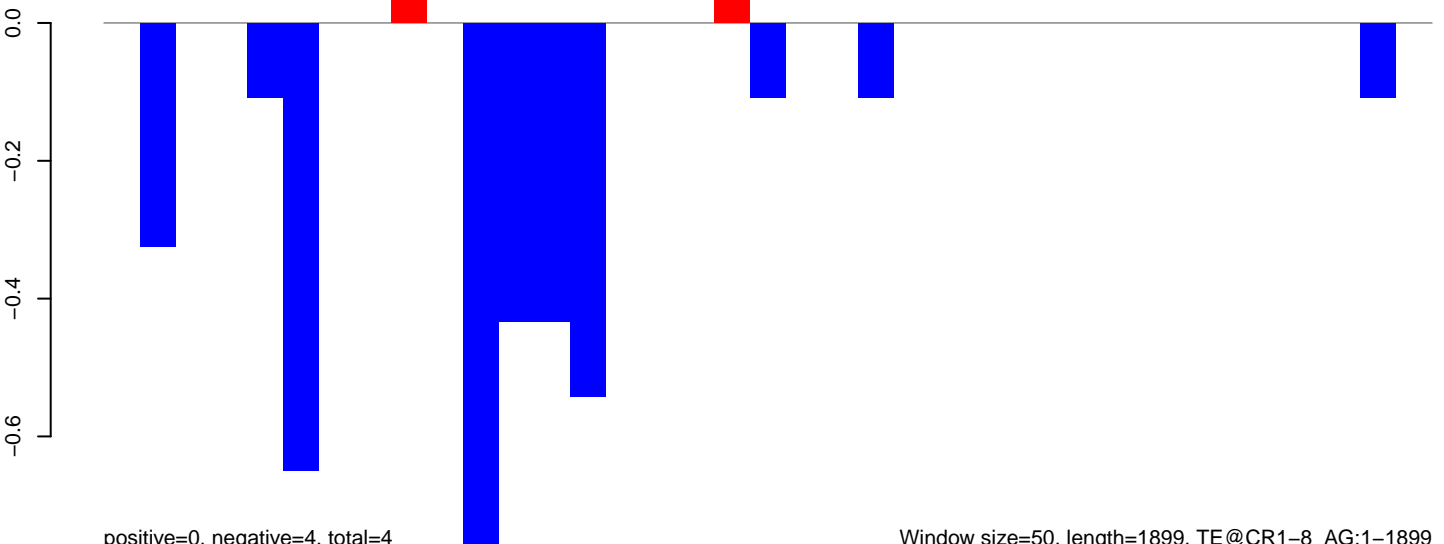
positive=0, negative=0, total=1

AnGam_Sua5b_cells.24_35.rep



positive=0, negative=3, total=3

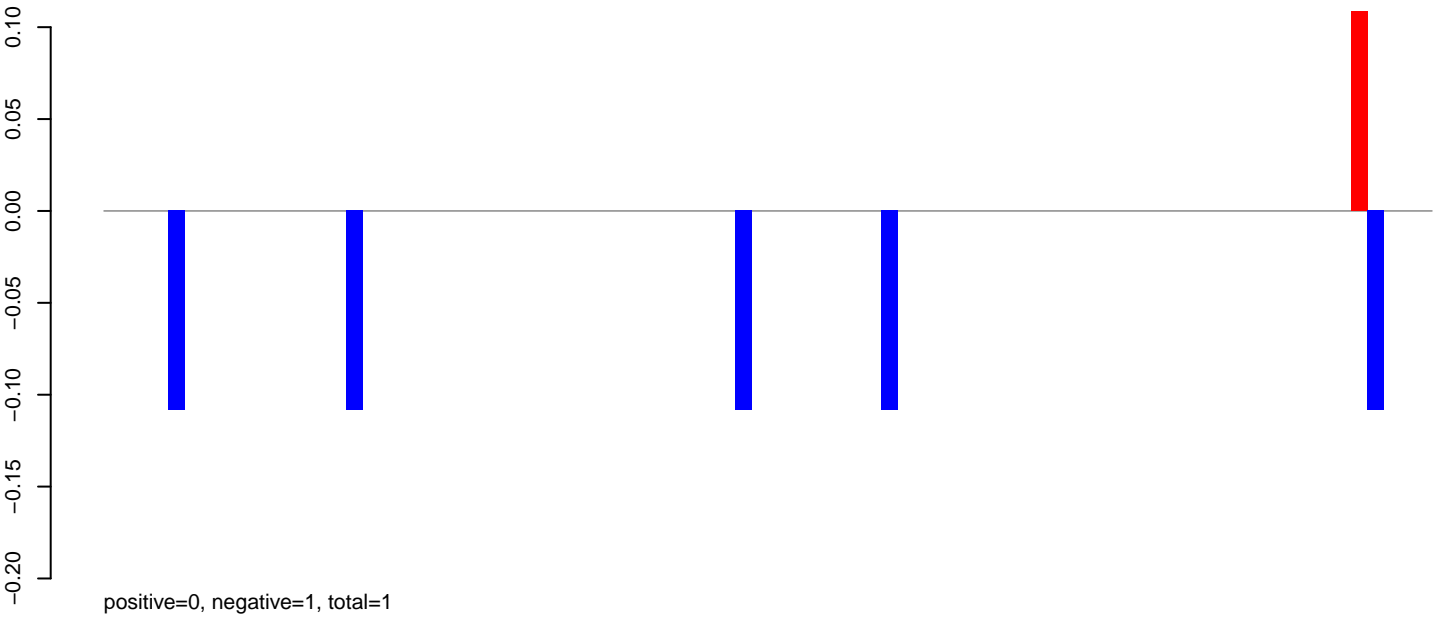
AnGam_Sua5b_cells.rep



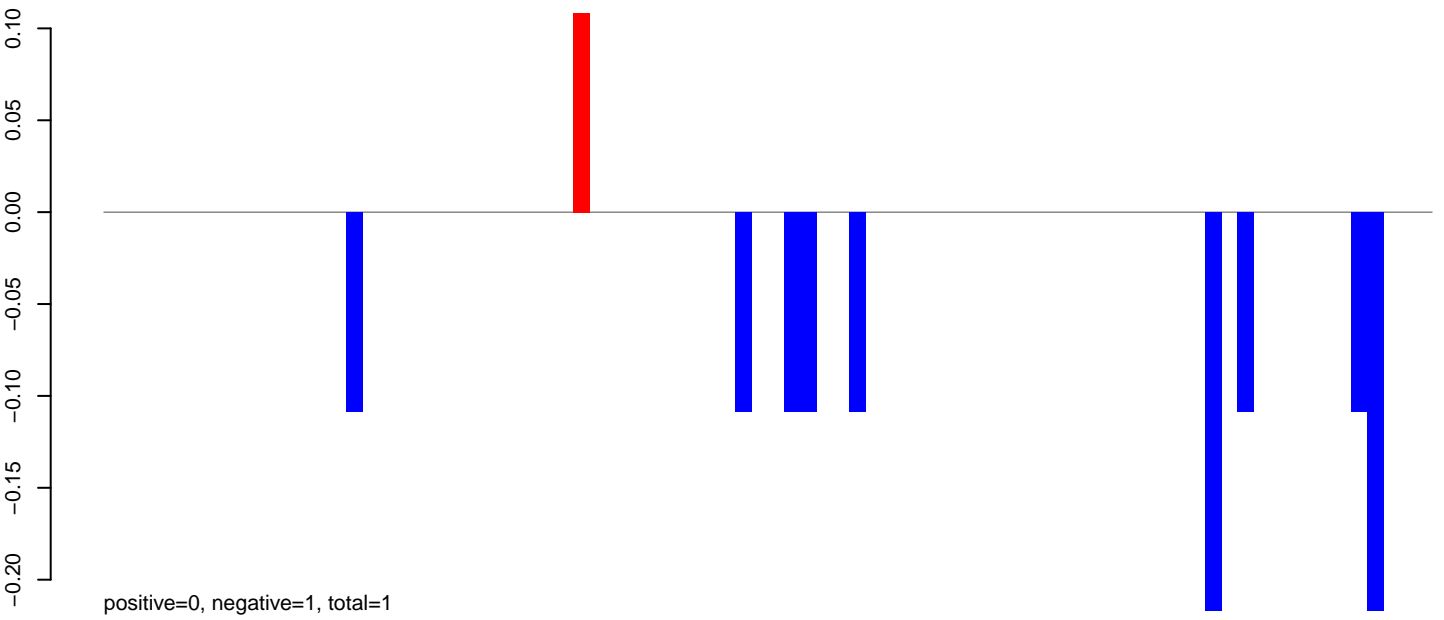
positive=0, negative=4, total=4

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

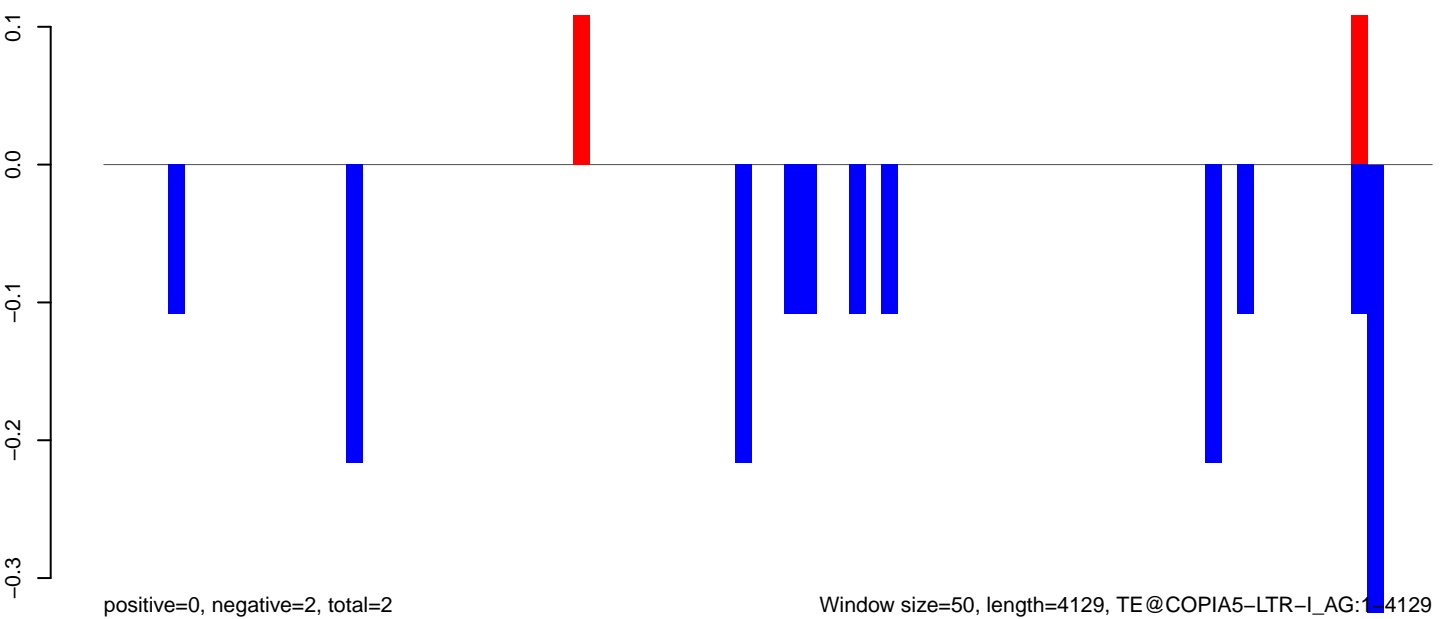
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



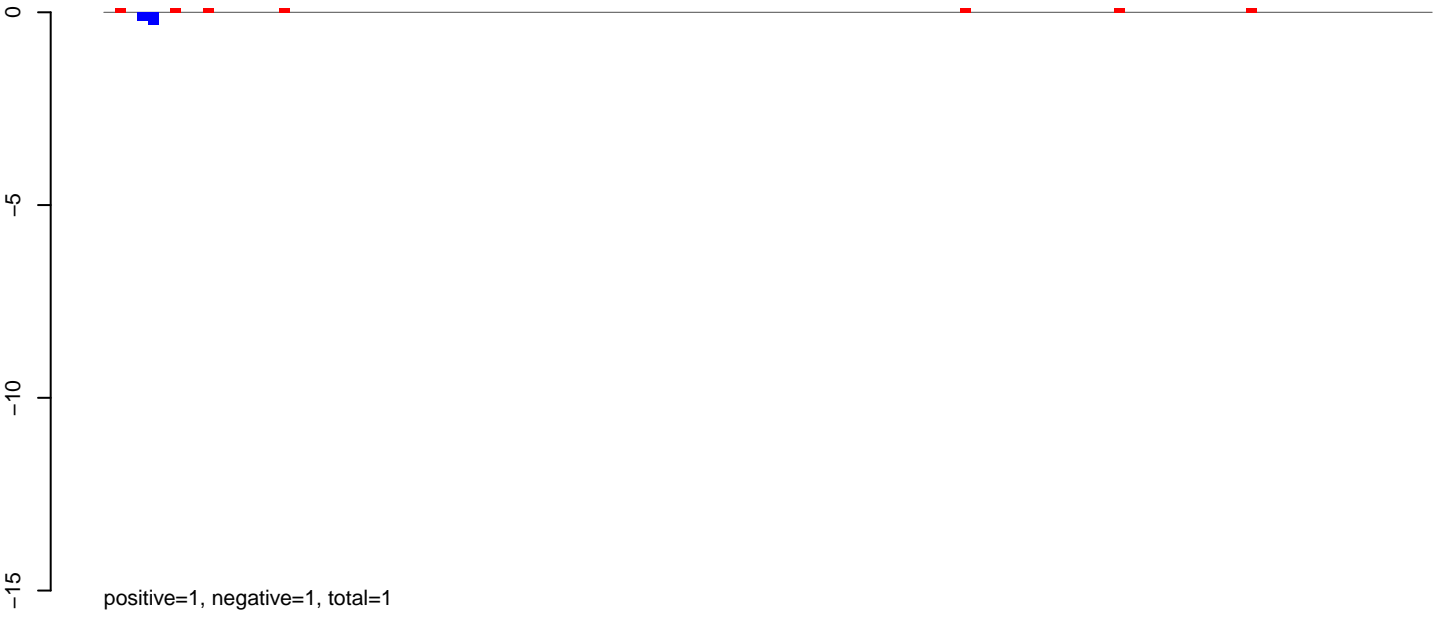
AnGam_Sua5b_cells.rep



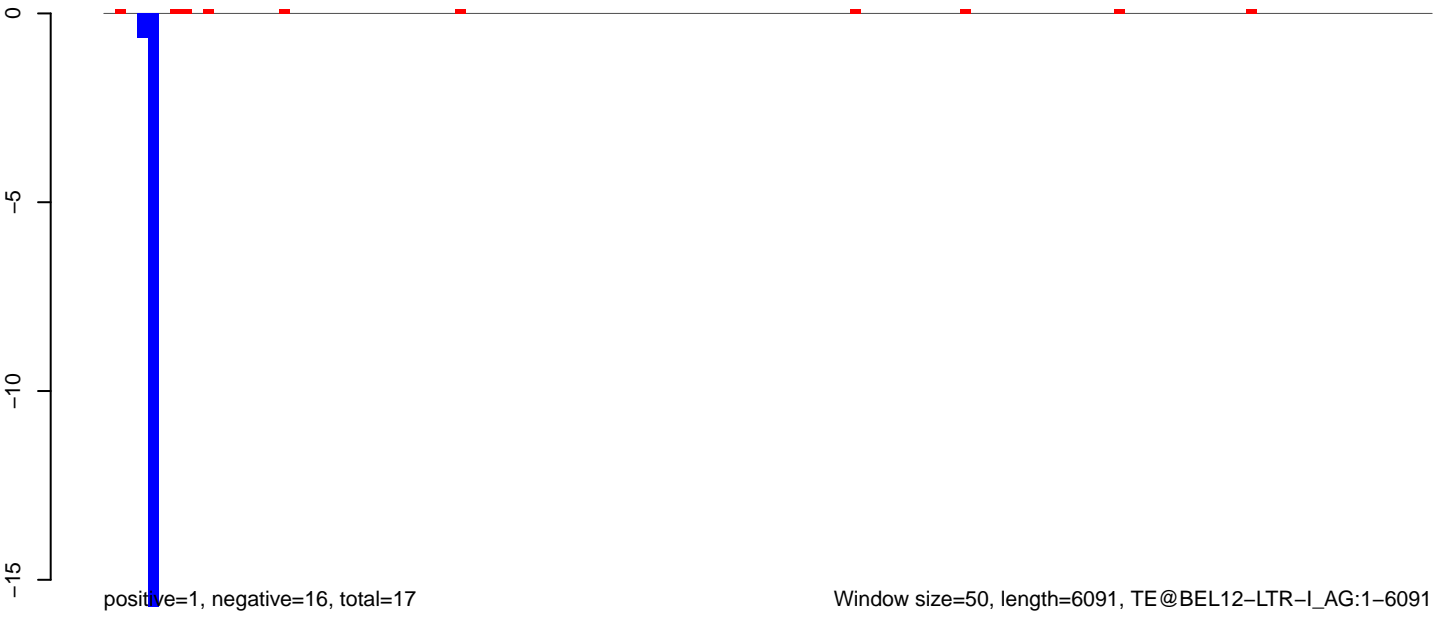
AnGam_Sua5b_cells.18_23.rep



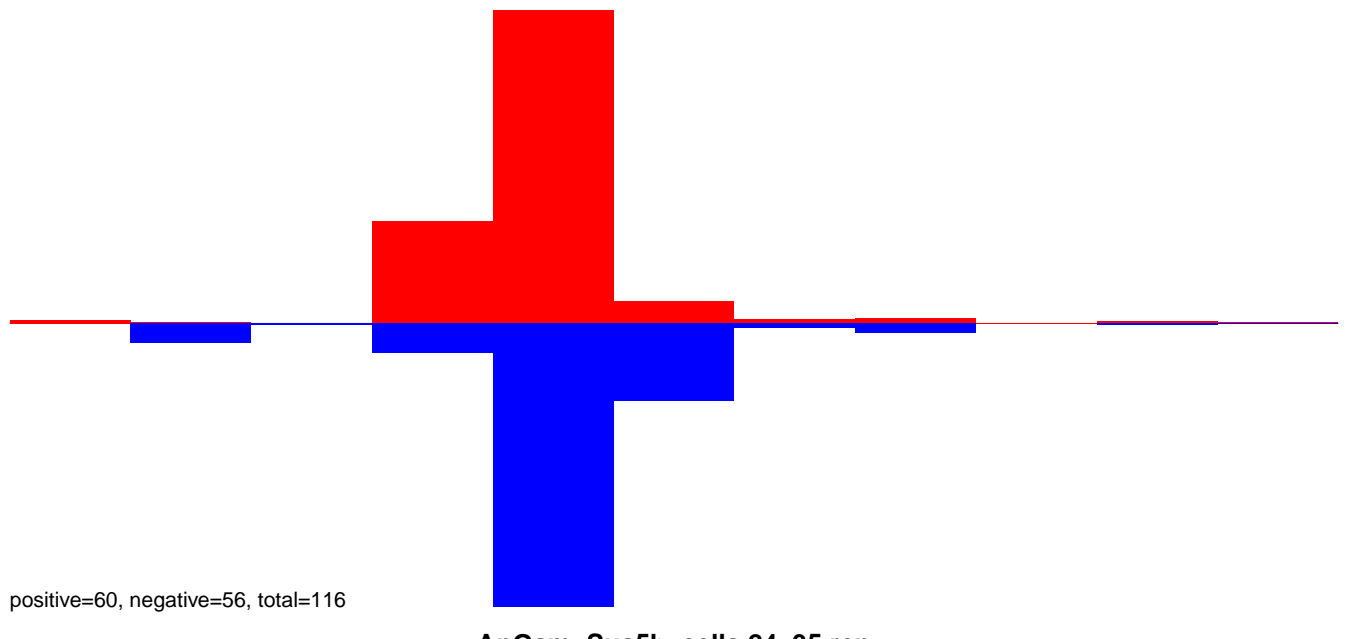
AnGam_Sua5b_cells.24_35.rep



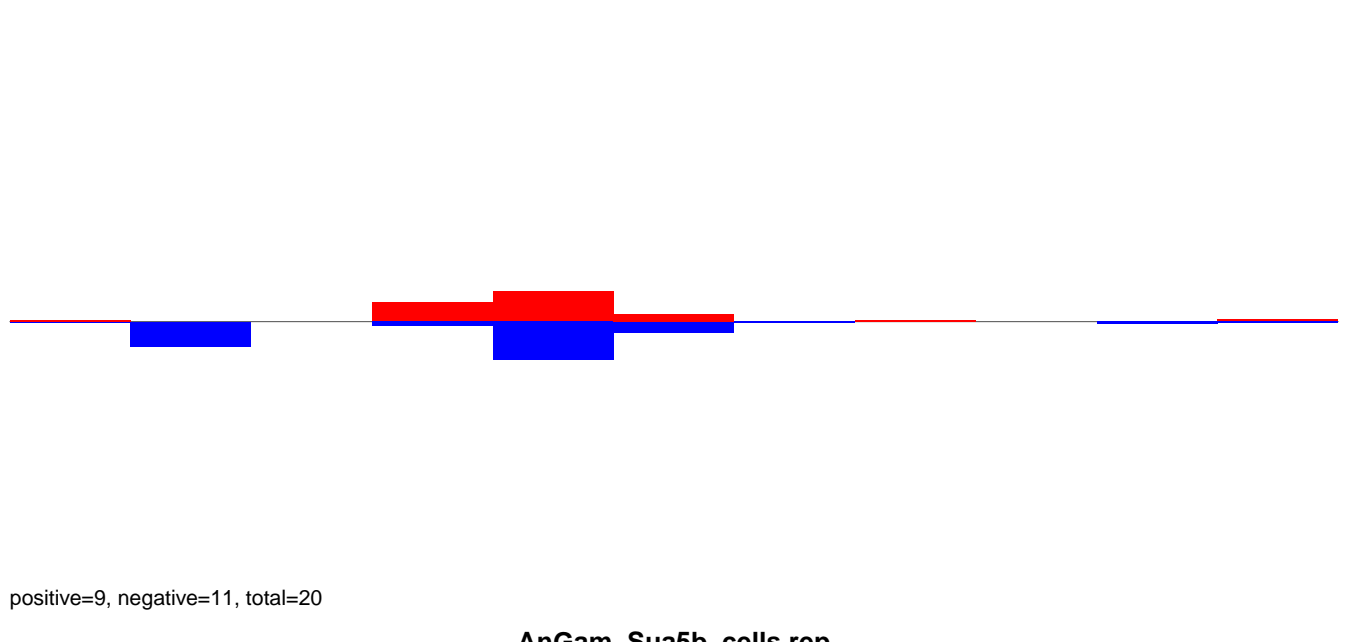
AnGam_Sua5b_cells.rep



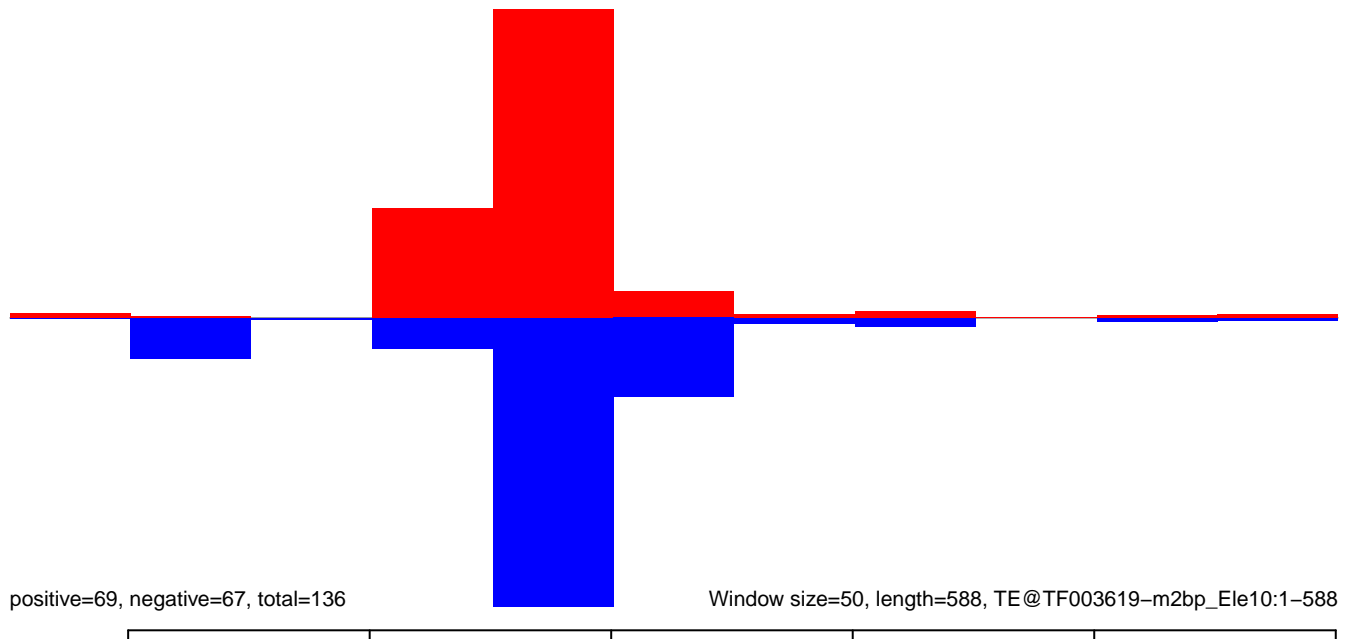
AnGam_Sua5b_cells.18_23.rep



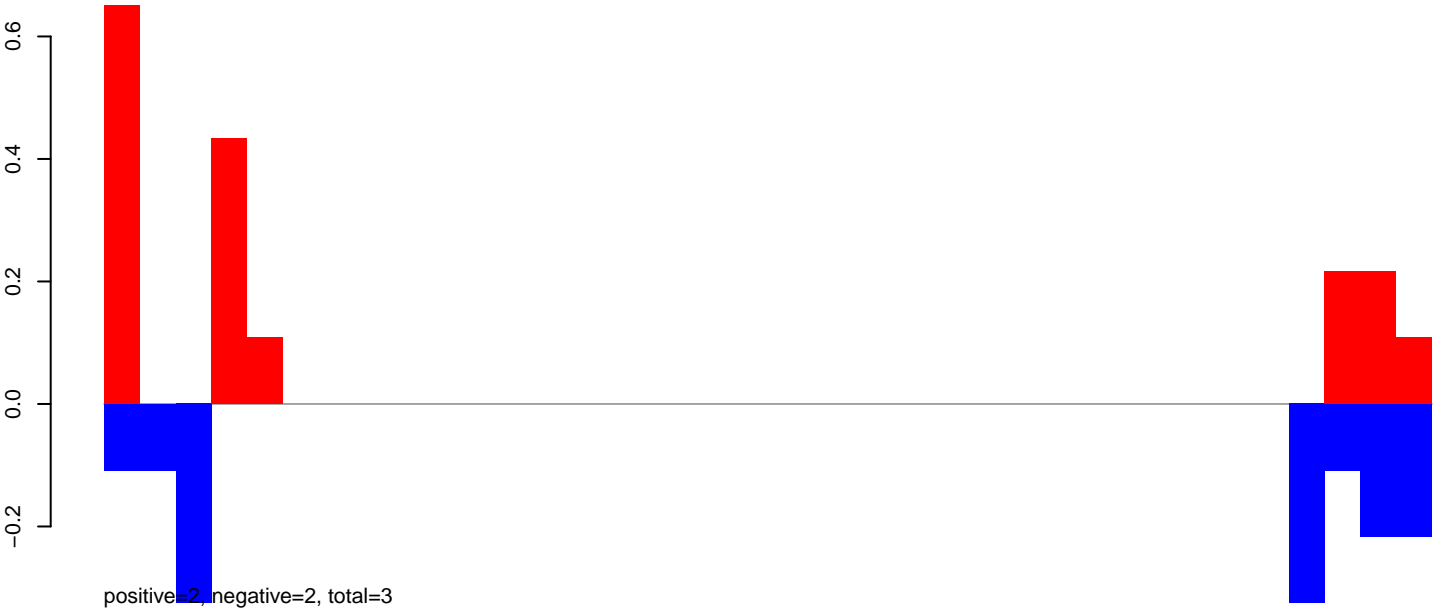
AnGam_Sua5b_cells.24_35.rep



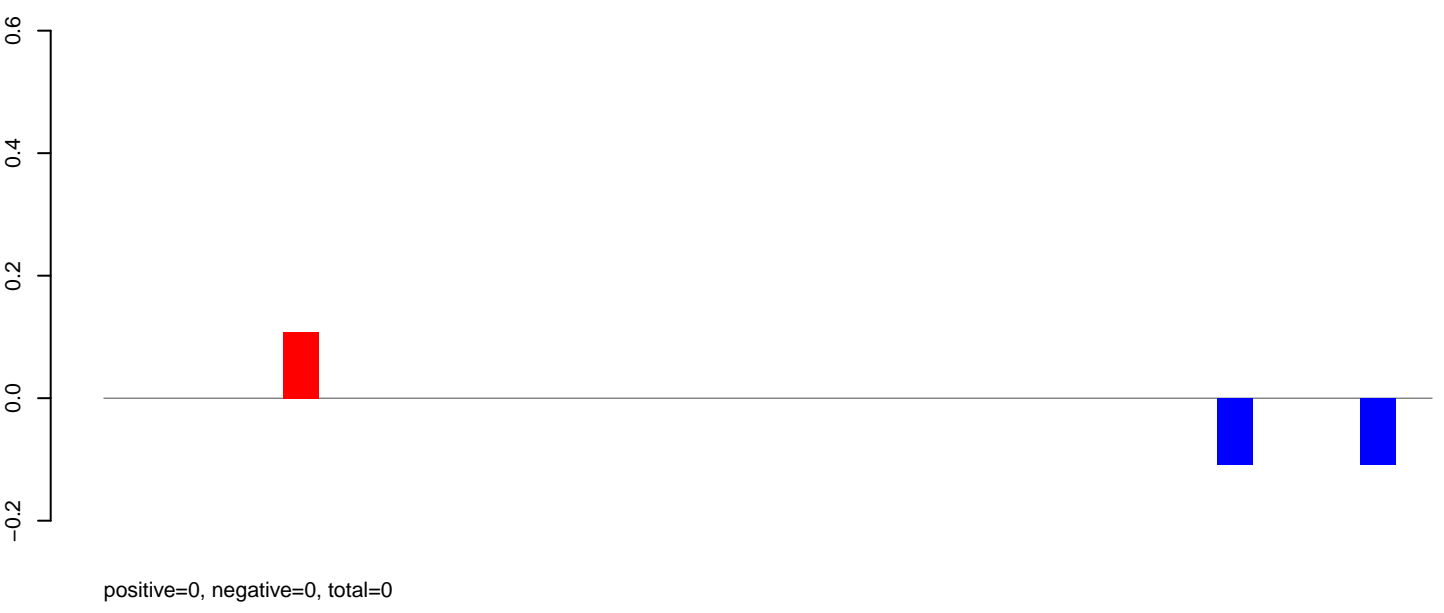
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

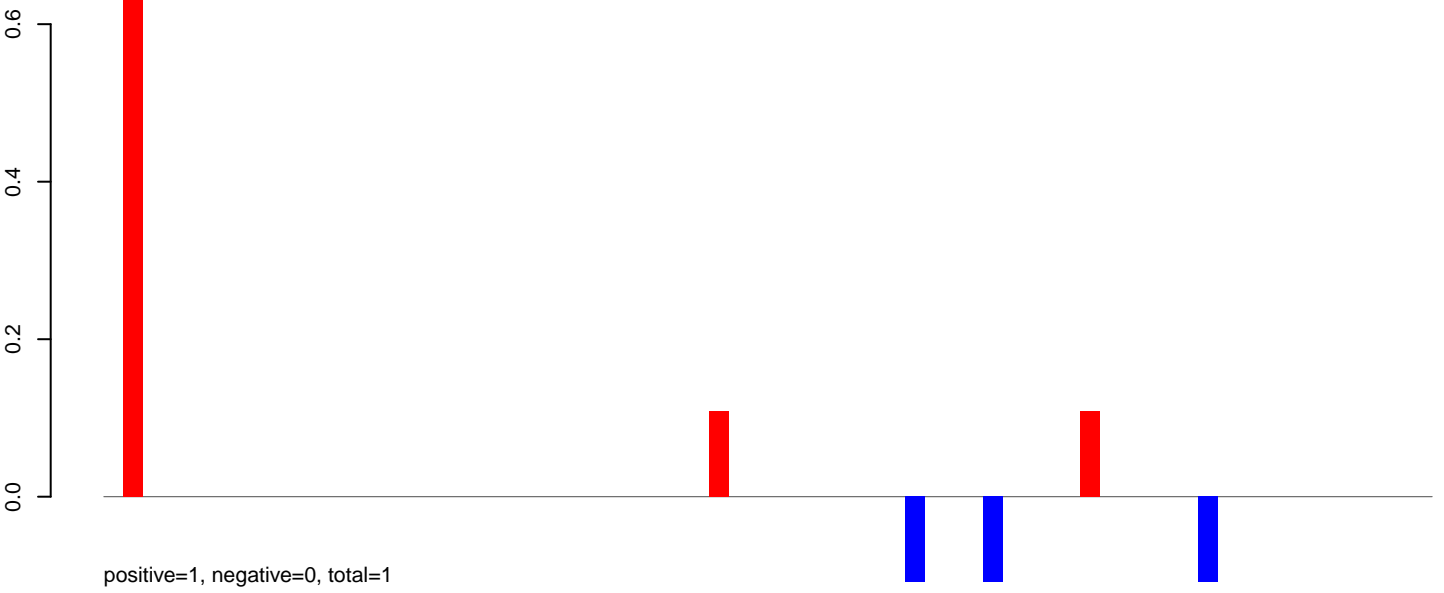


AnGam_Sua5b_cells.rep

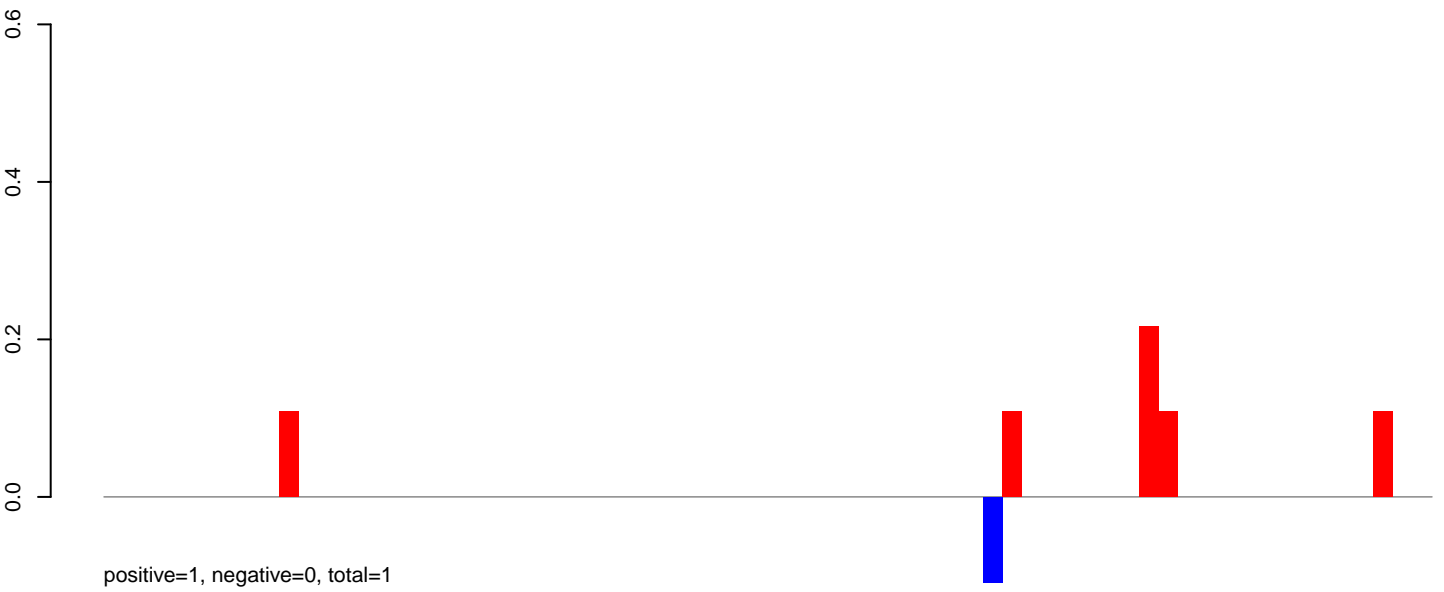


0 500 1000 1500

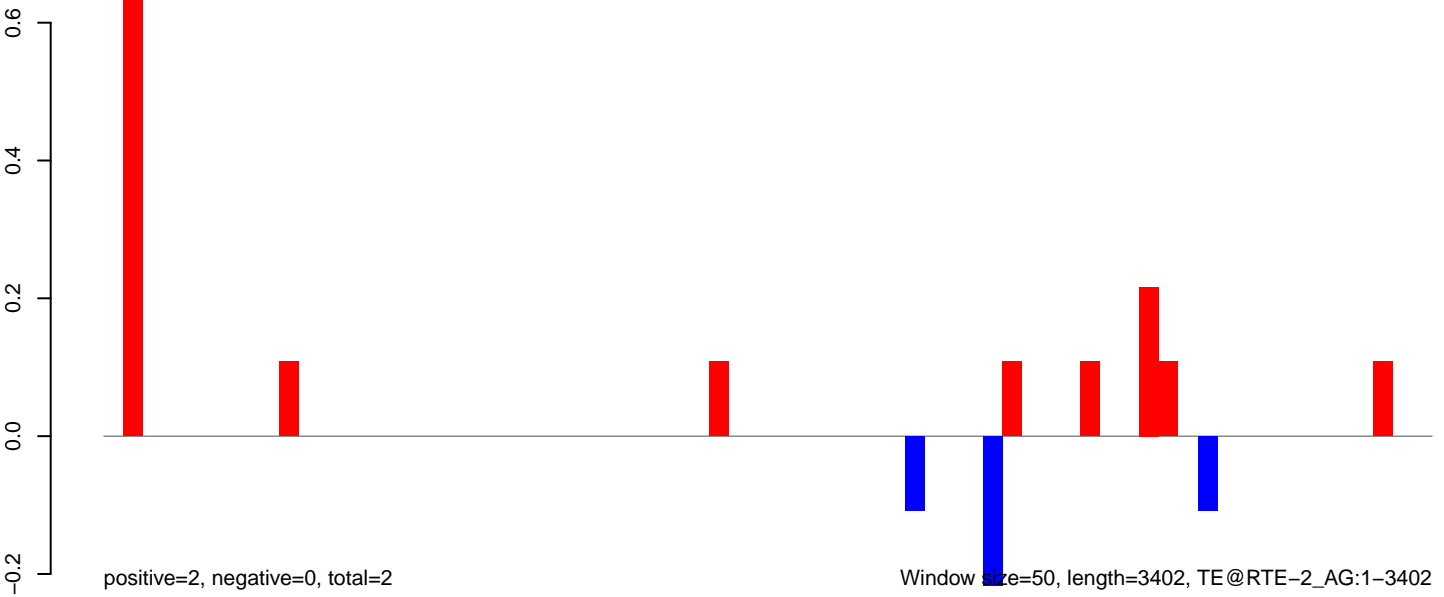
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



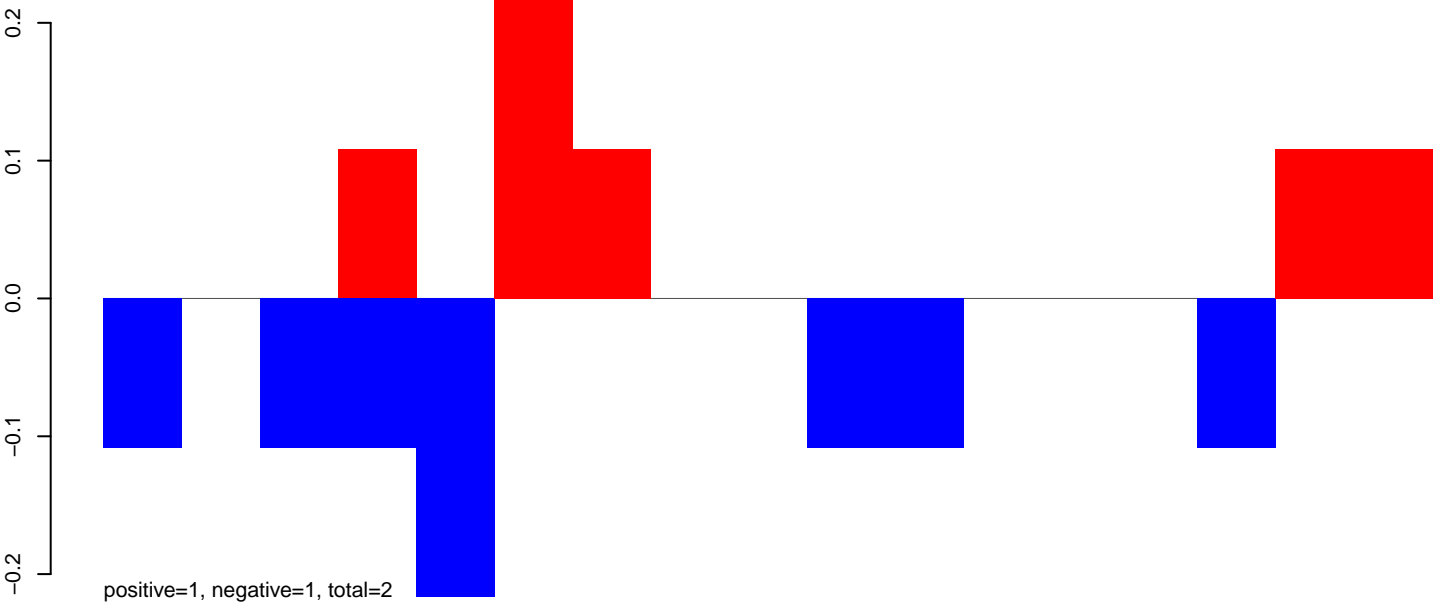
AnGam_Sua5b_cells.rep



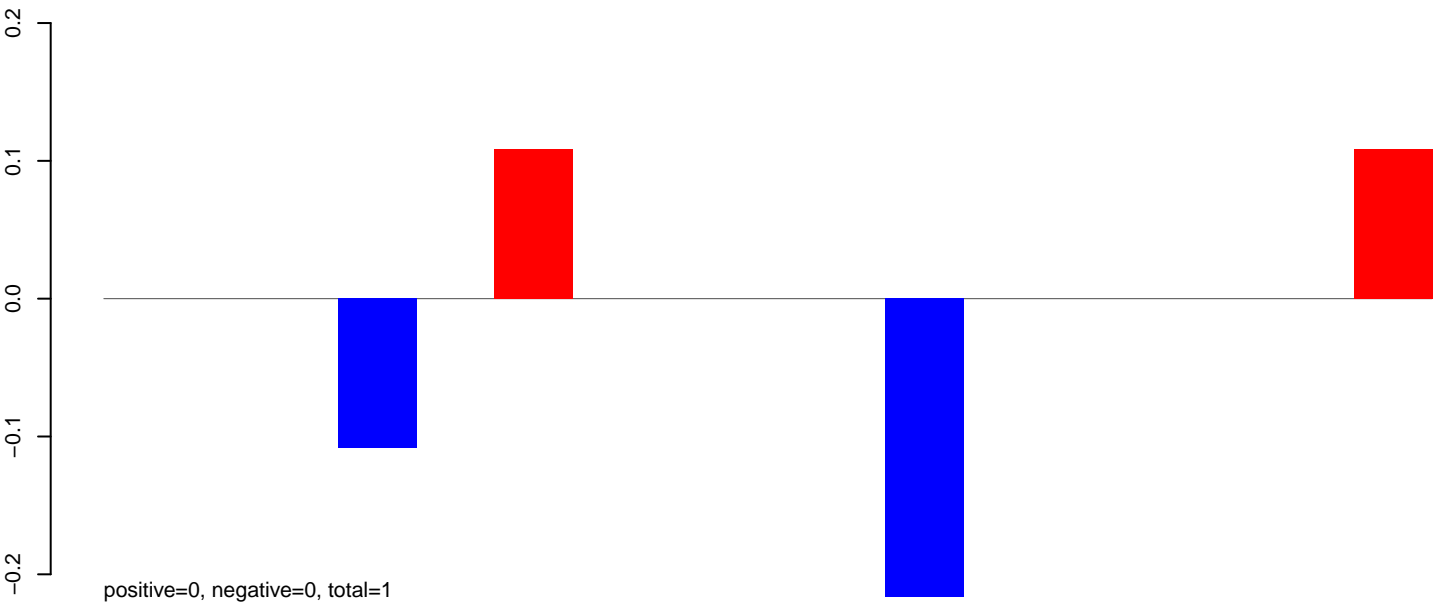
Window size=50, length=3402, TE@RTE-2_AG:1-3402

0 500 1000 1500 2000 2500 3000 3500

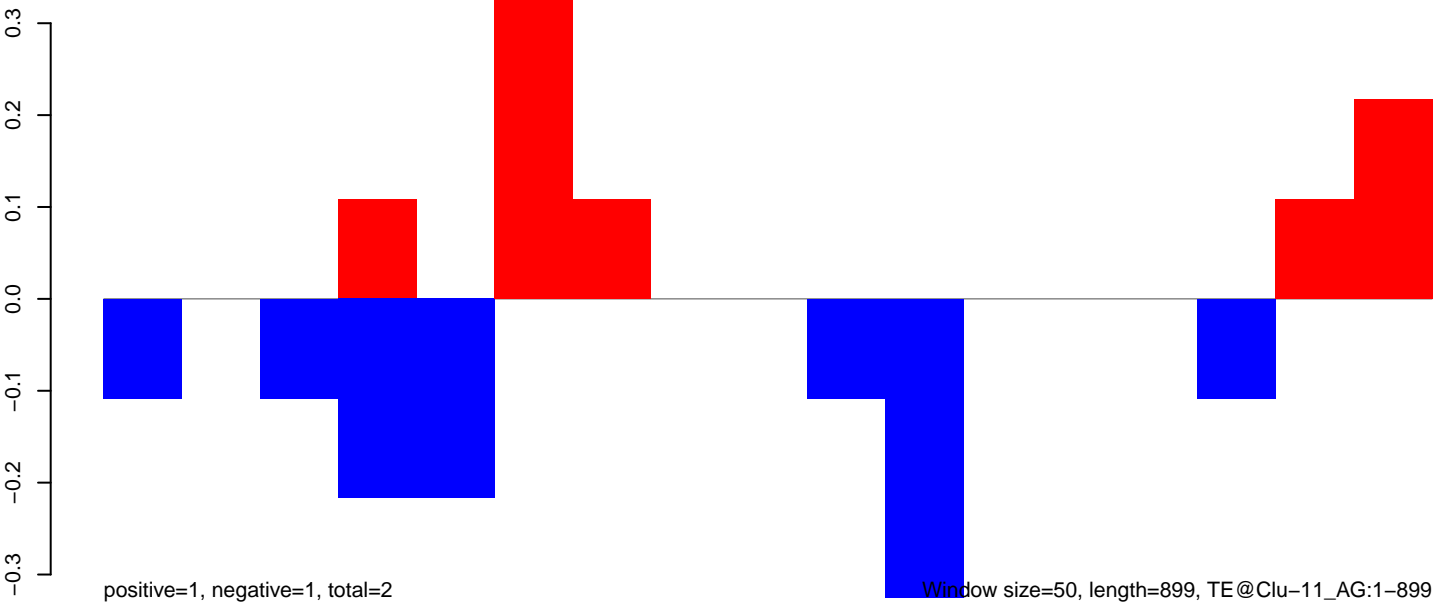
AnGam_Sua5b_cells.18_23.rep



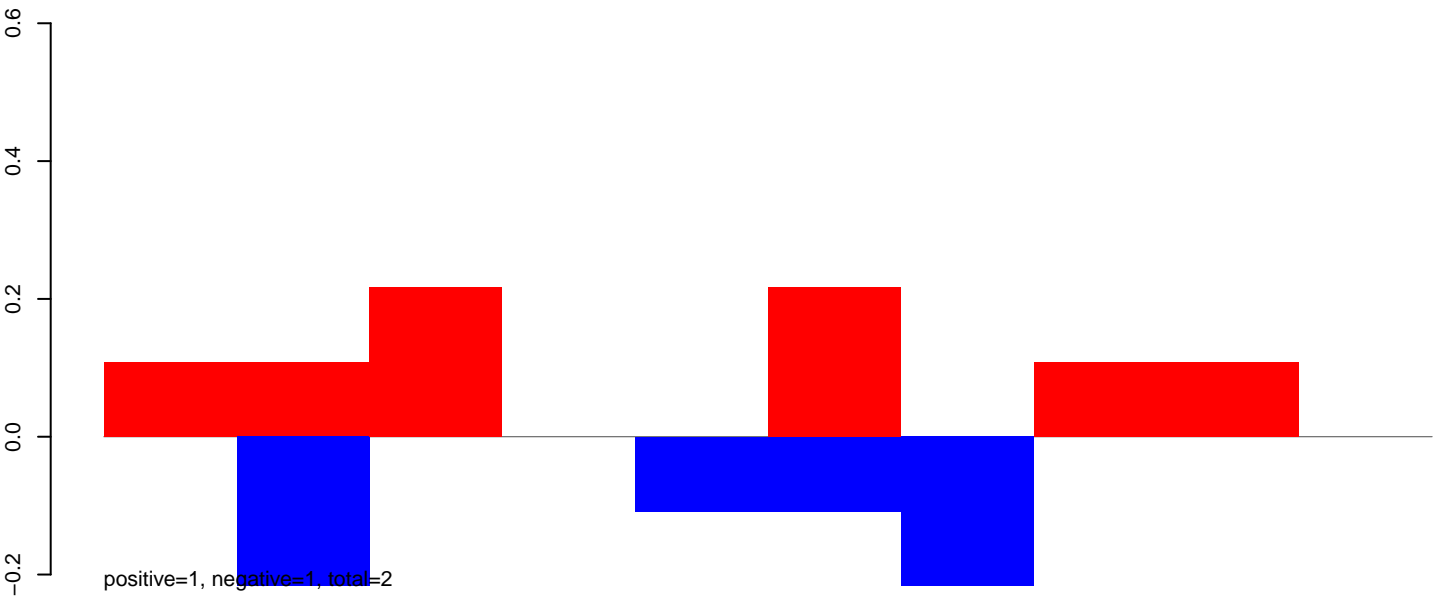
AnGam_Sua5b_cells.24_35.rep



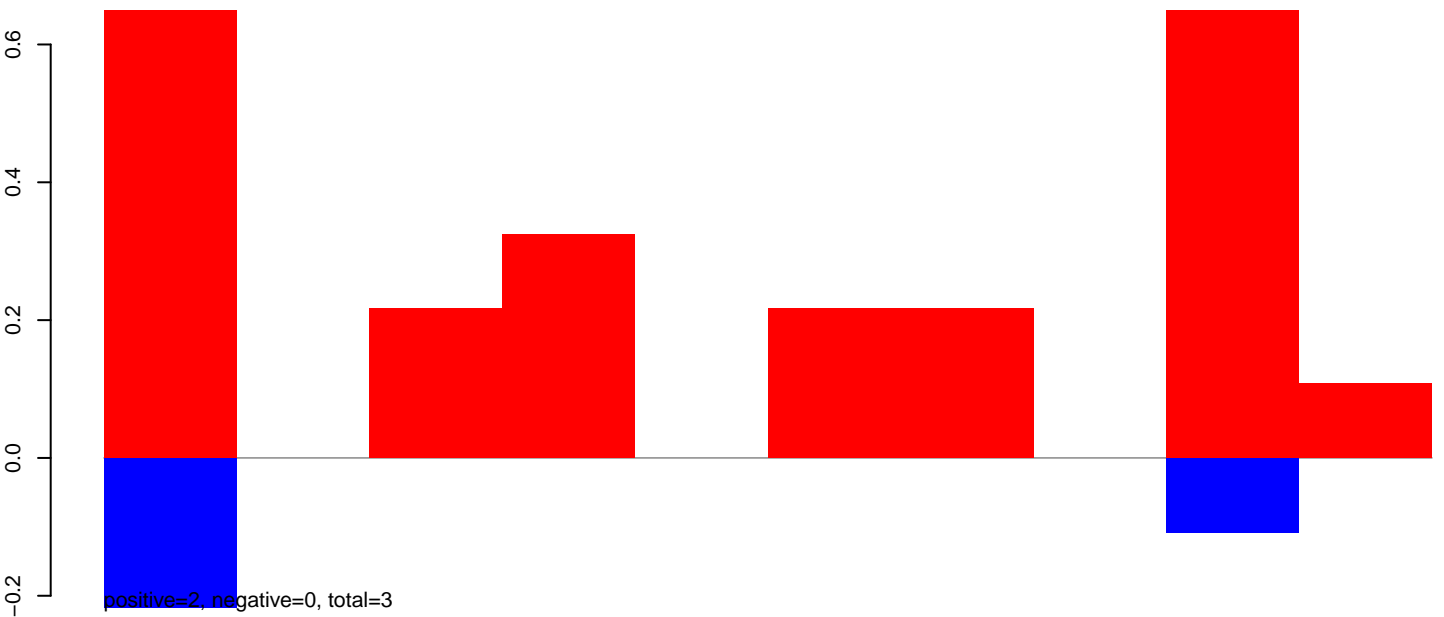
AnGam_Sua5b_cells.rep



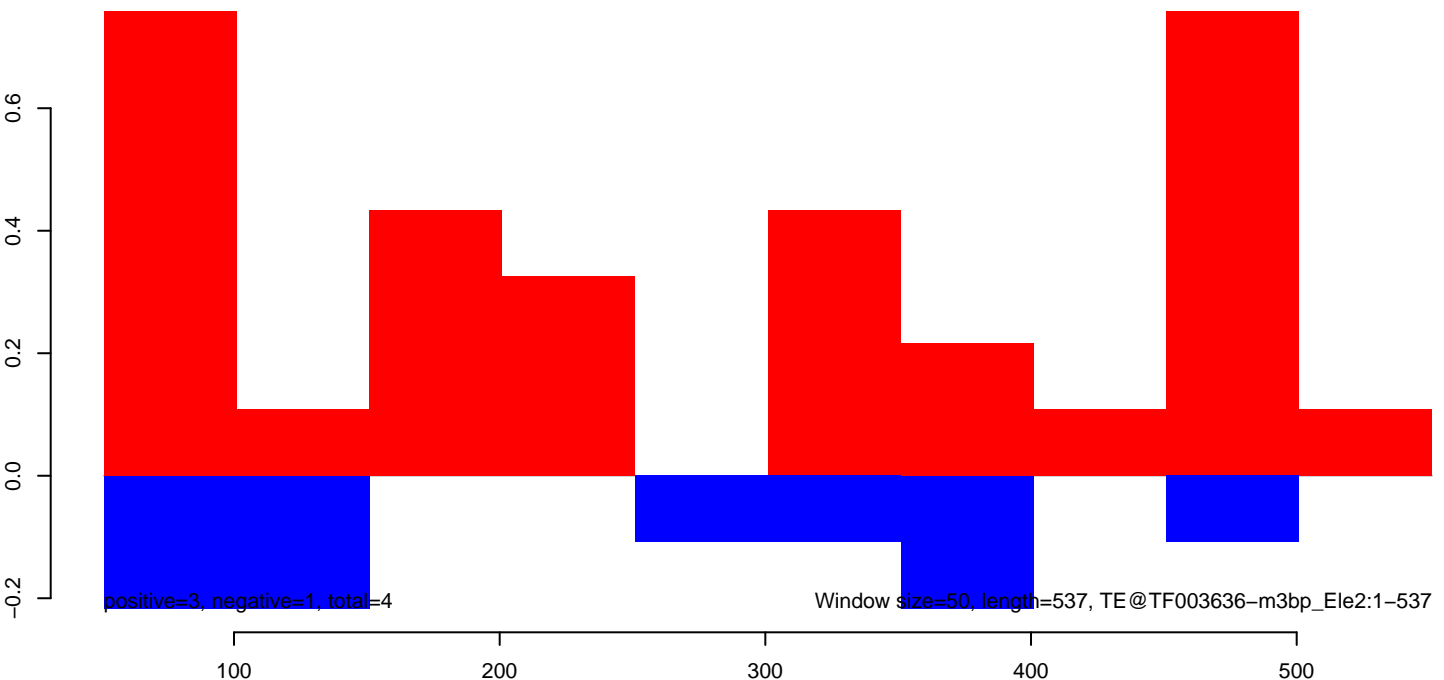
AnGam_Sua5b_cells.18_23.rep



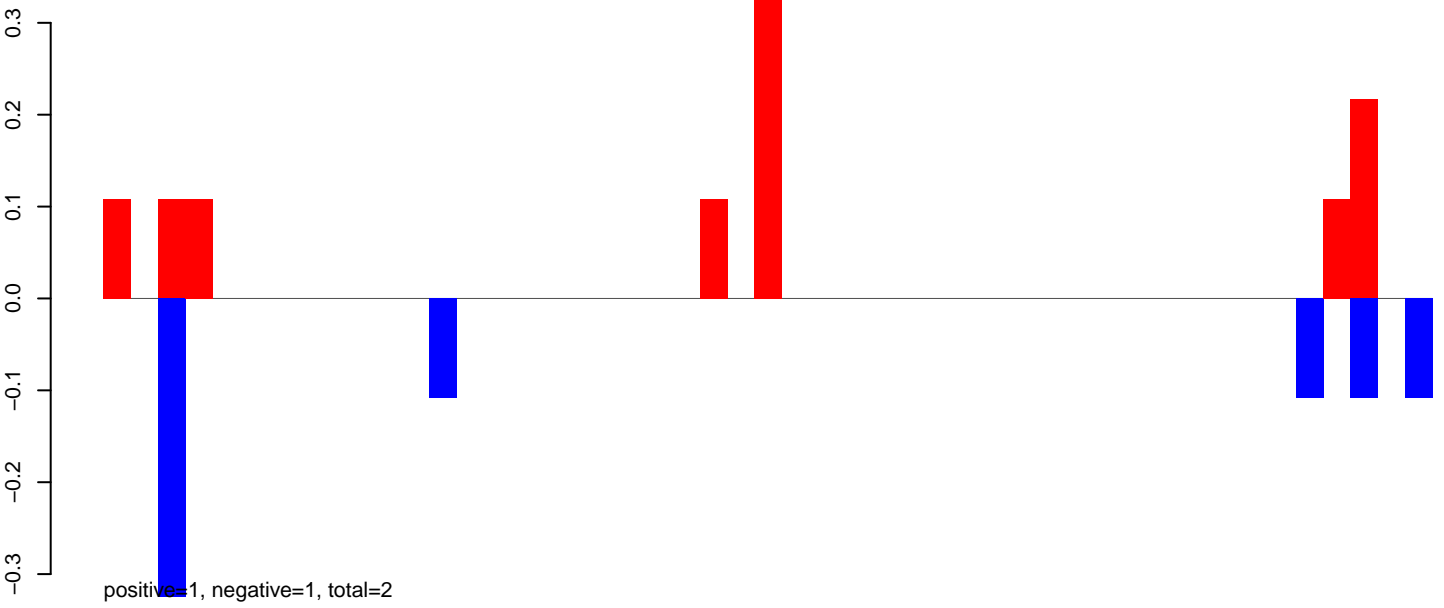
AnGam_Sua5b_cells.24_35.rep



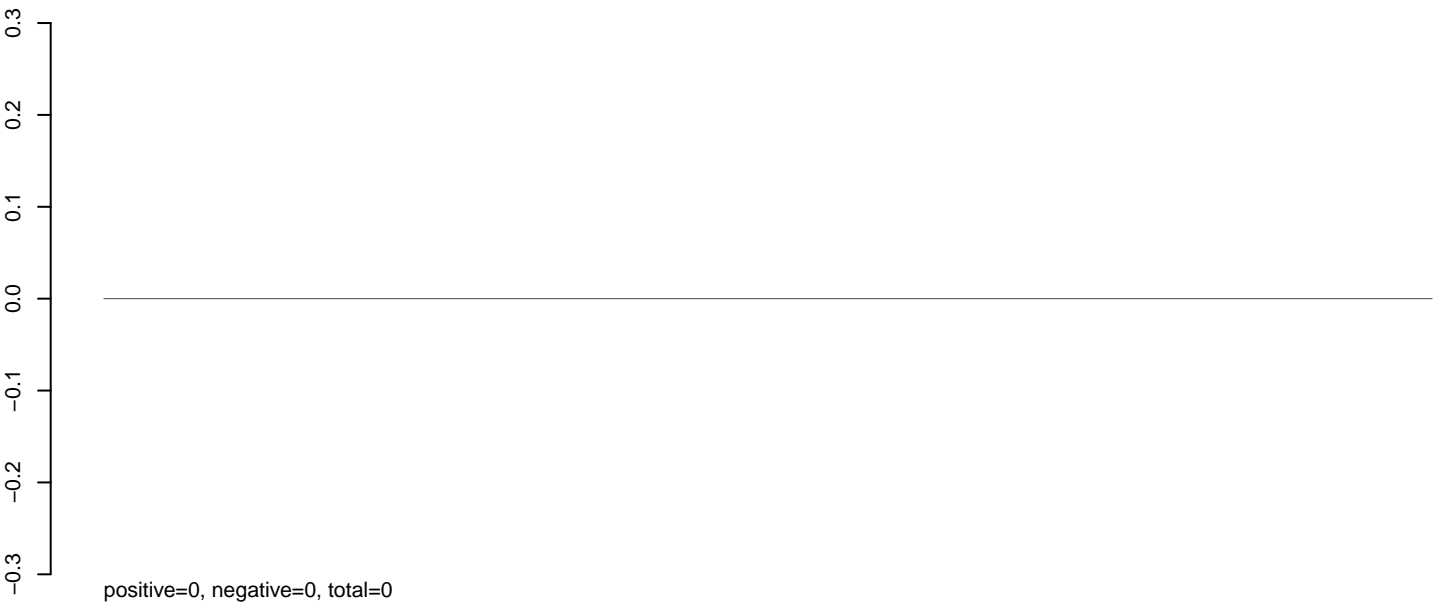
AnGam_Sua5b_cells.rep



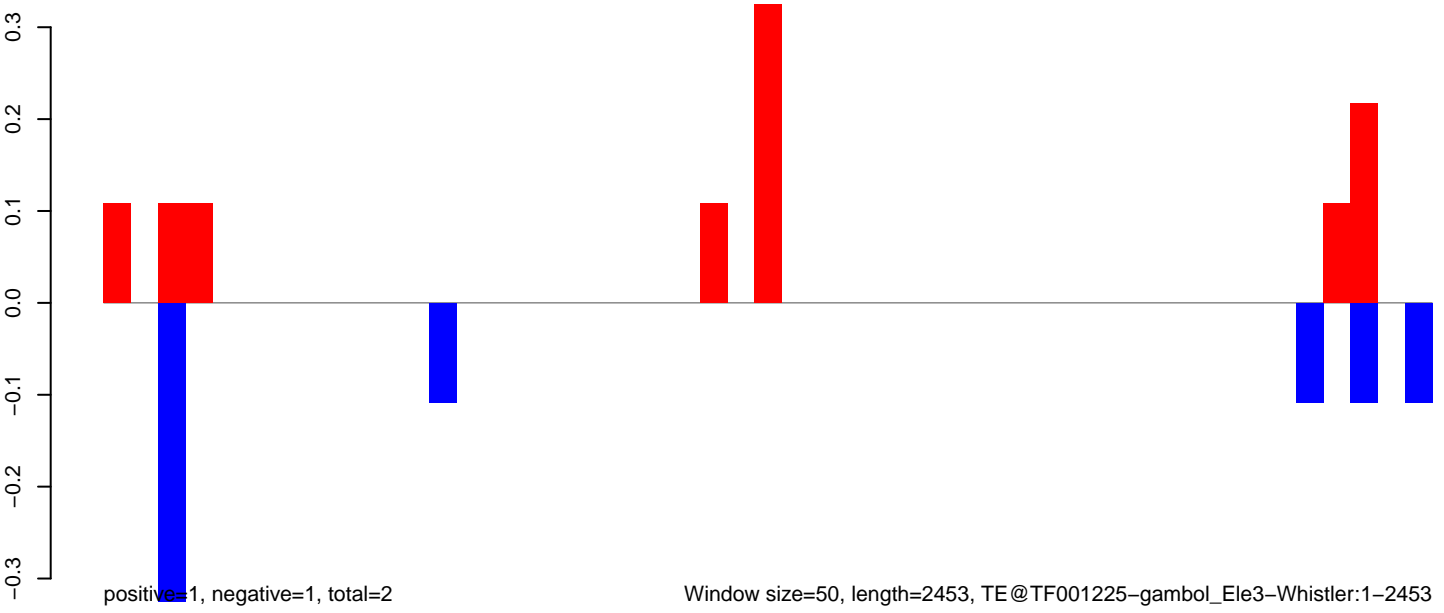
AnGam_Sua5b_cells.18_23.rep



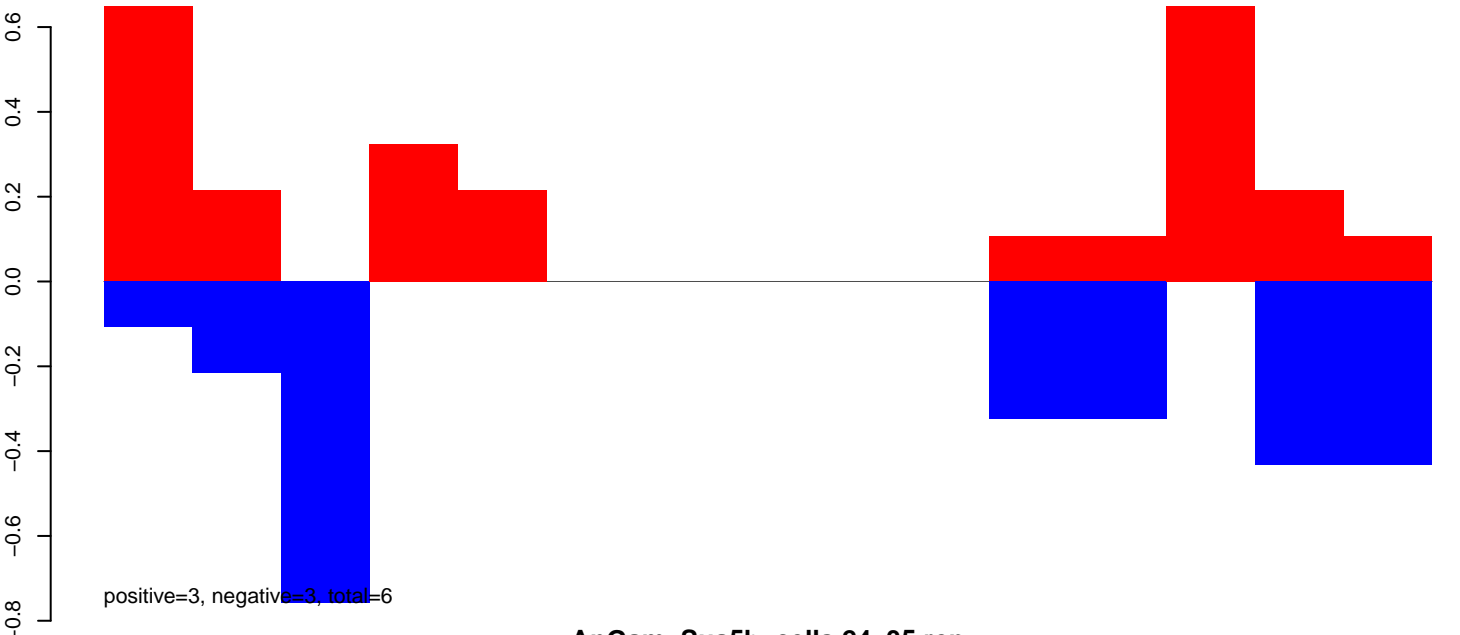
AnGam_Sua5b_cells.24_35.rep



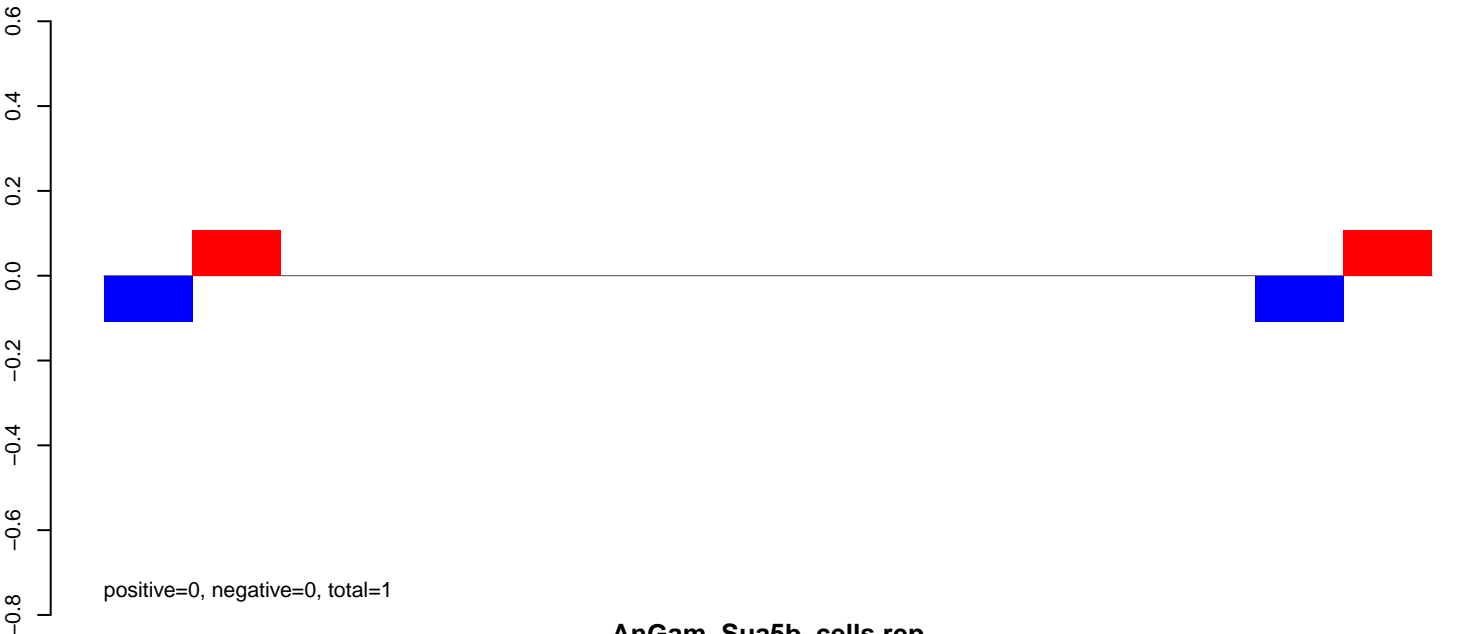
AnGam_Sua5b_cells.rep



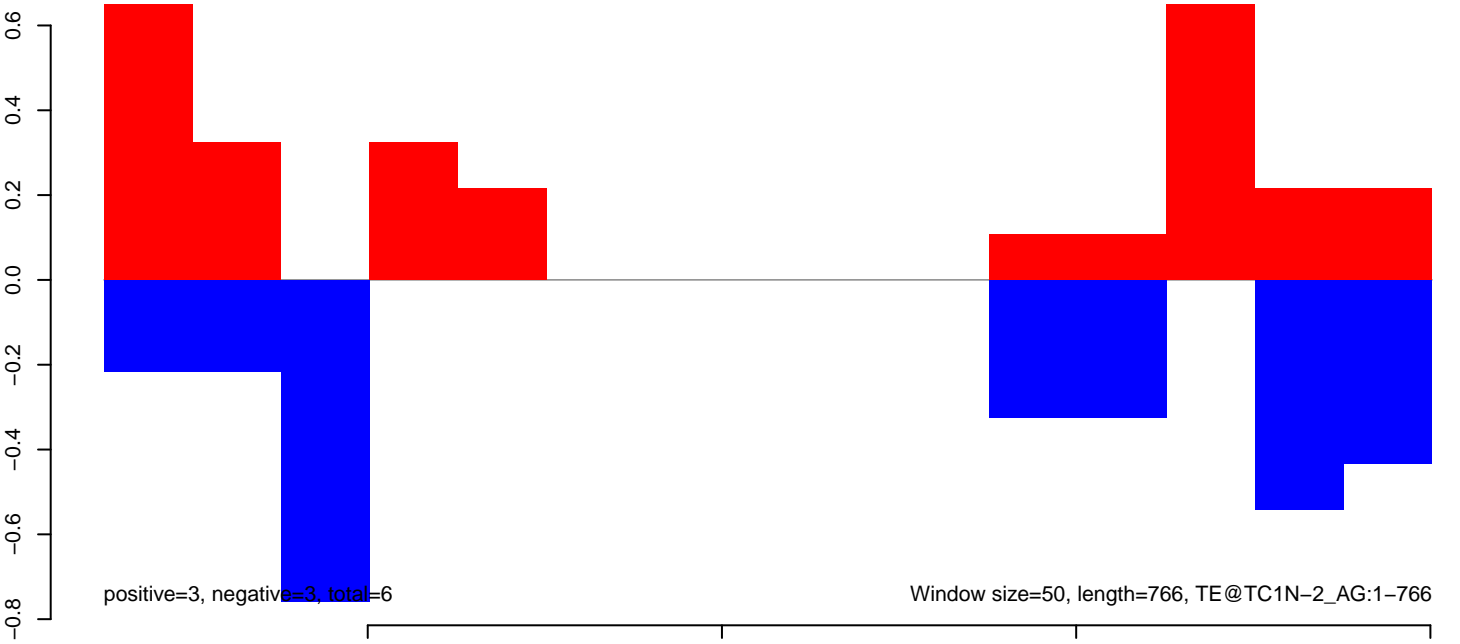
AnGam_Sua5b_cells.18_23.rep



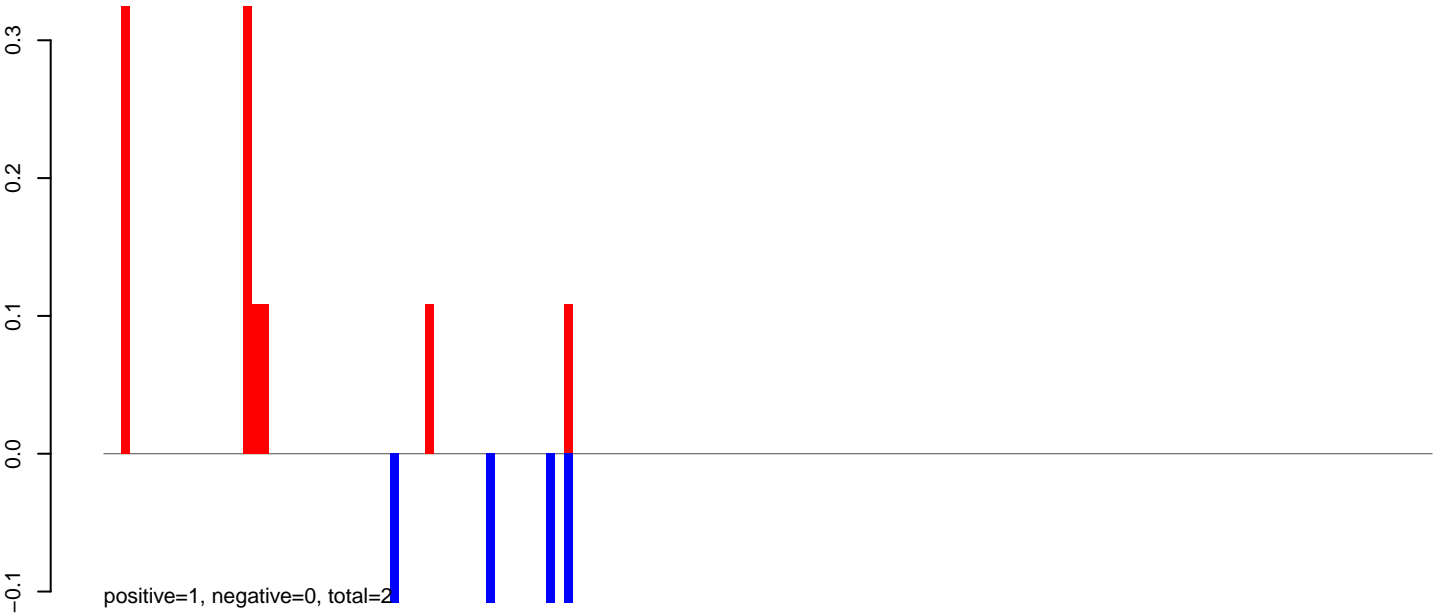
AnGam_Sua5b_cells.24_35.rep



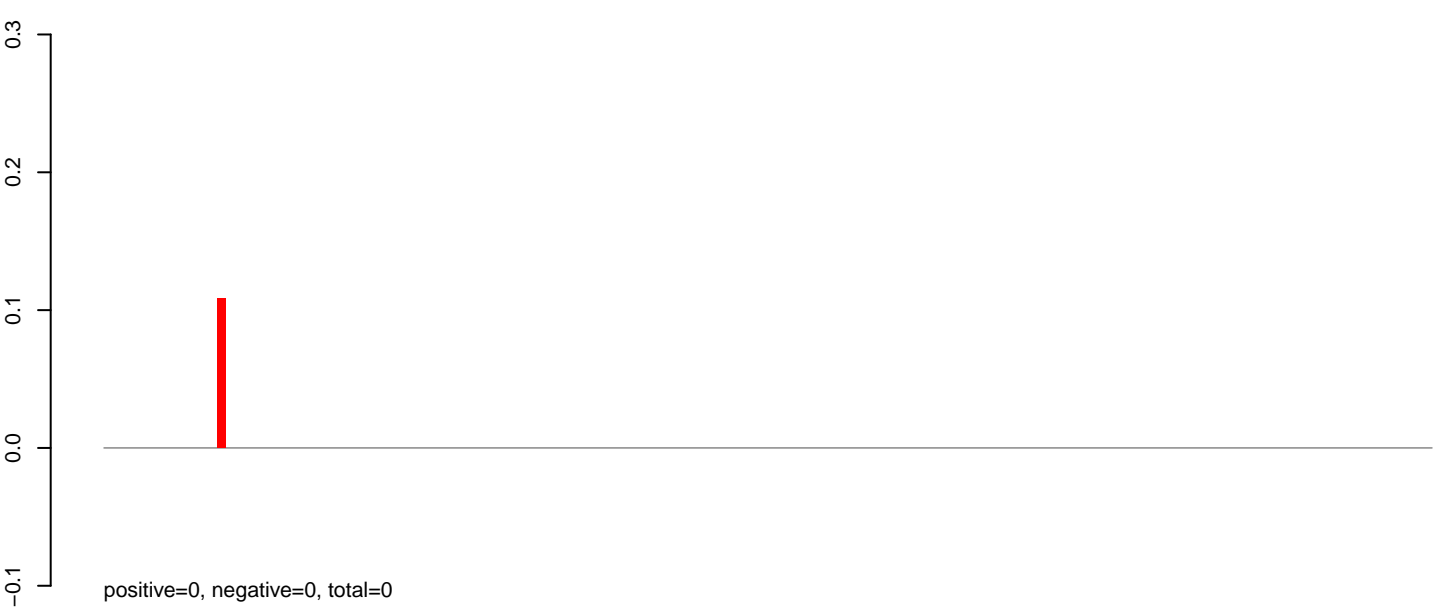
AnGam_Sua5b_cells.rep



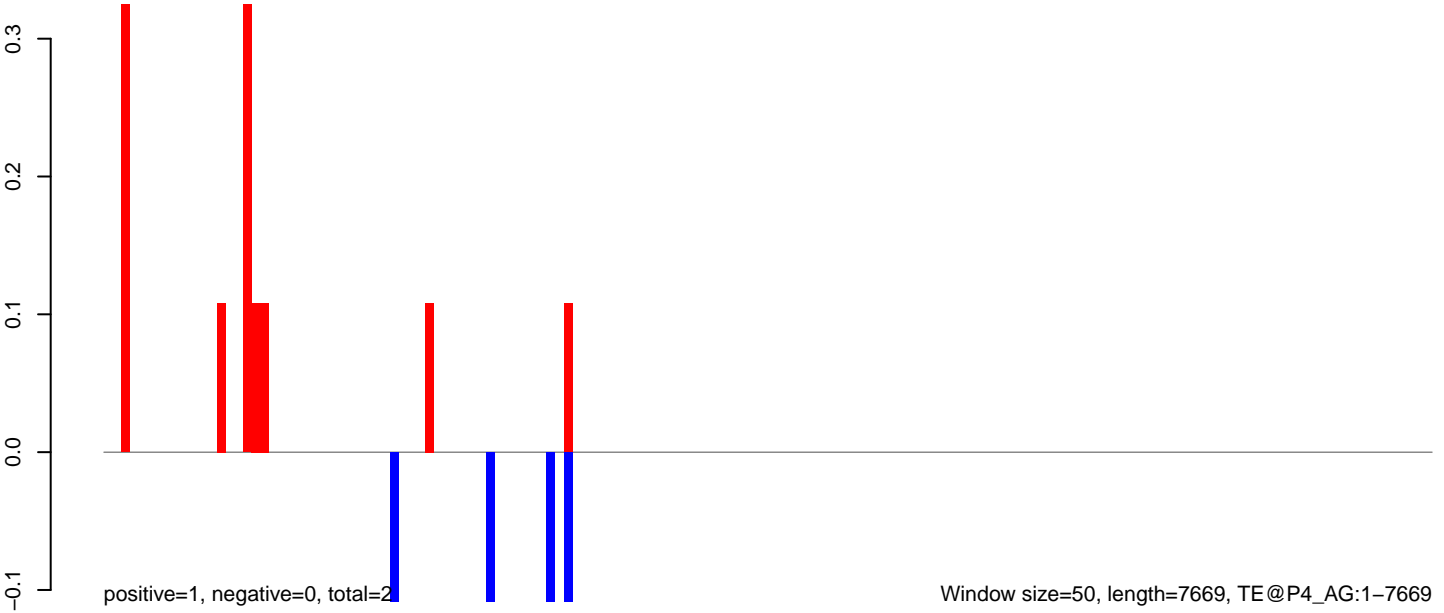
AnGam_Sua5b_cells.18_23.rep



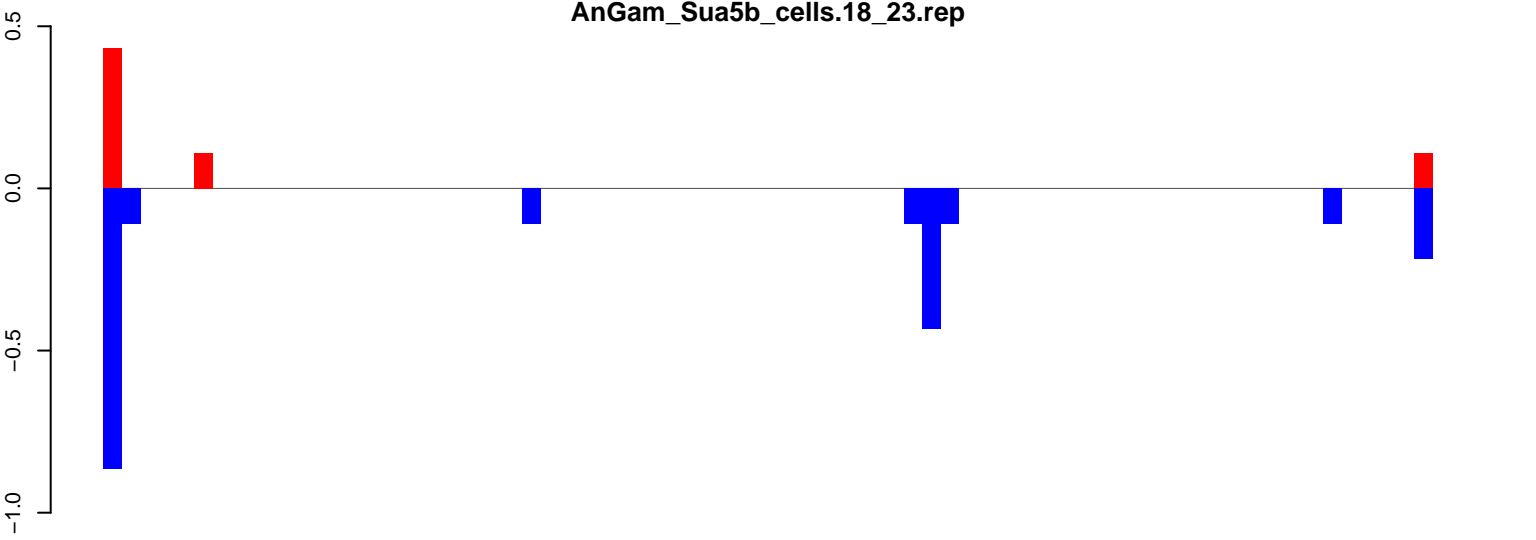
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

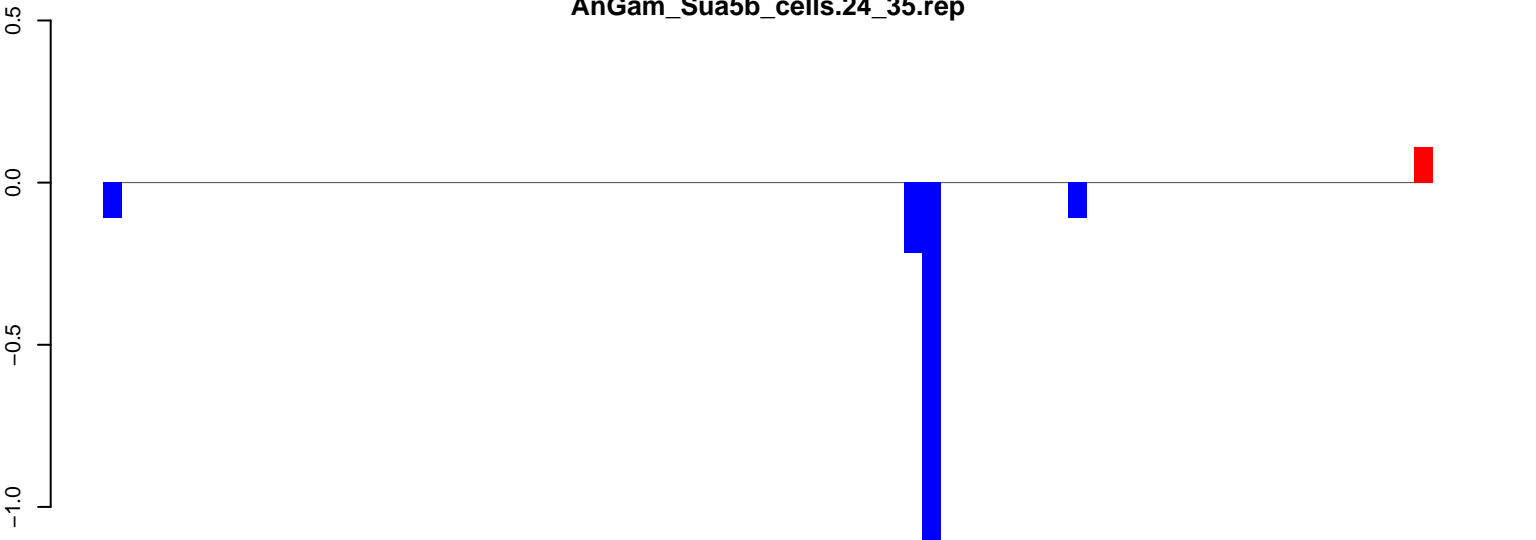


AnGam_Sua5b_cells.18_23.rep



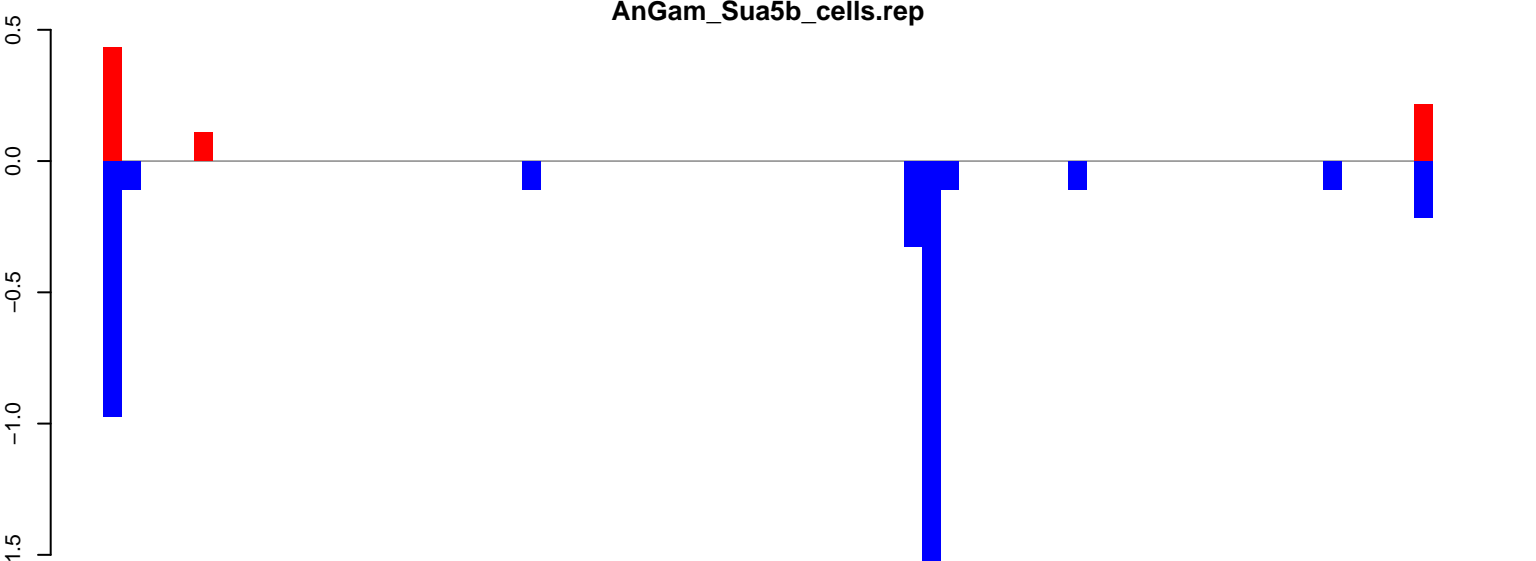
positive=1, negative=2, total=3

AnGam_Sua5b_cells.24_35.rep



positive=0, negative=2, total=2

AnGam_Sua5b_cells.rep



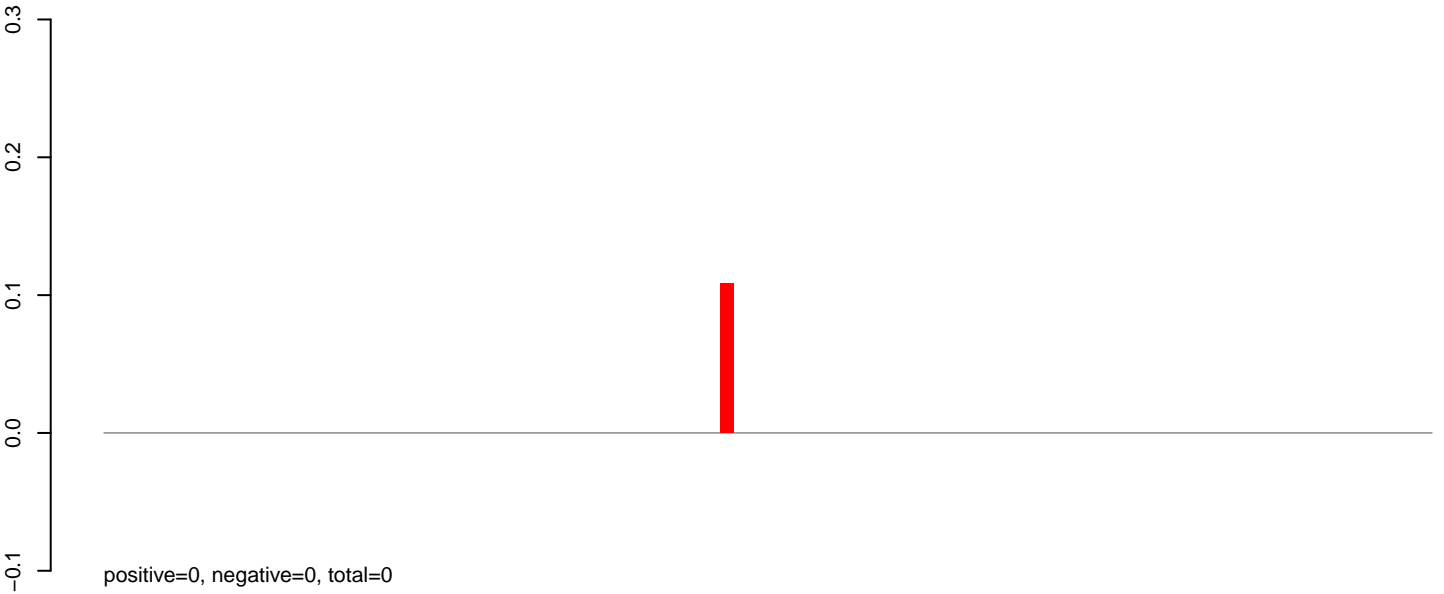
positive=1, negative=4, total=5

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

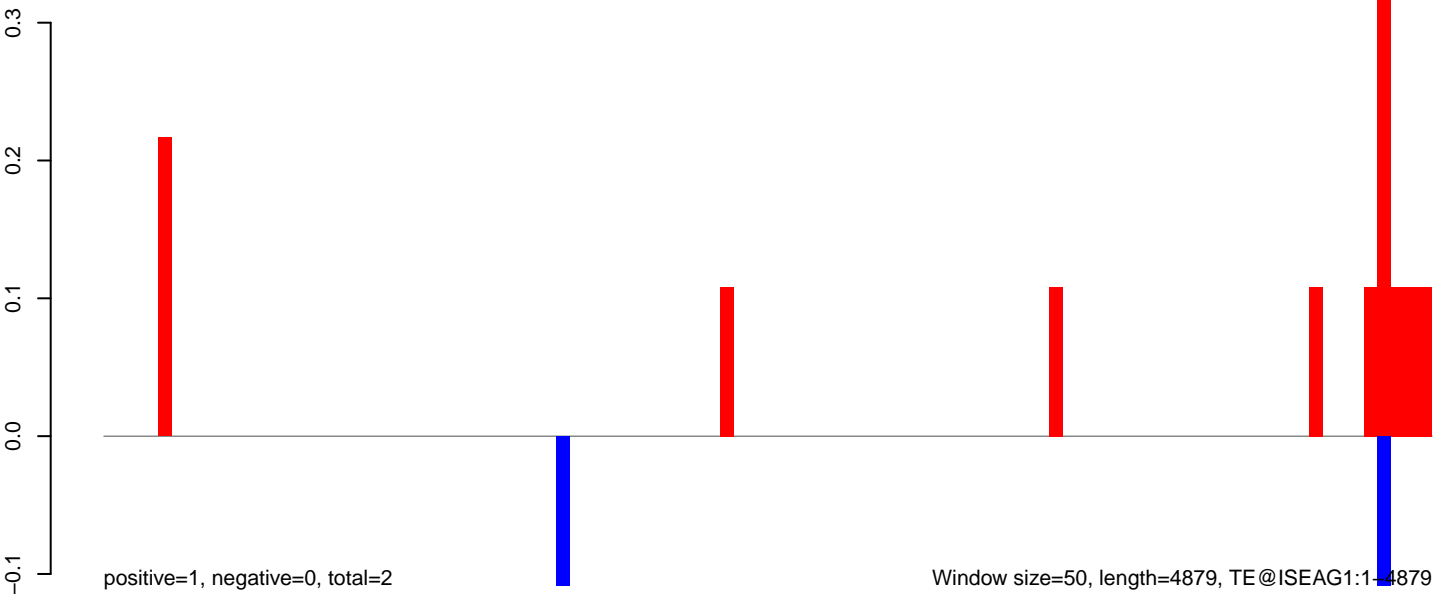
AnGam_Sua5b_cells.18_23.rep



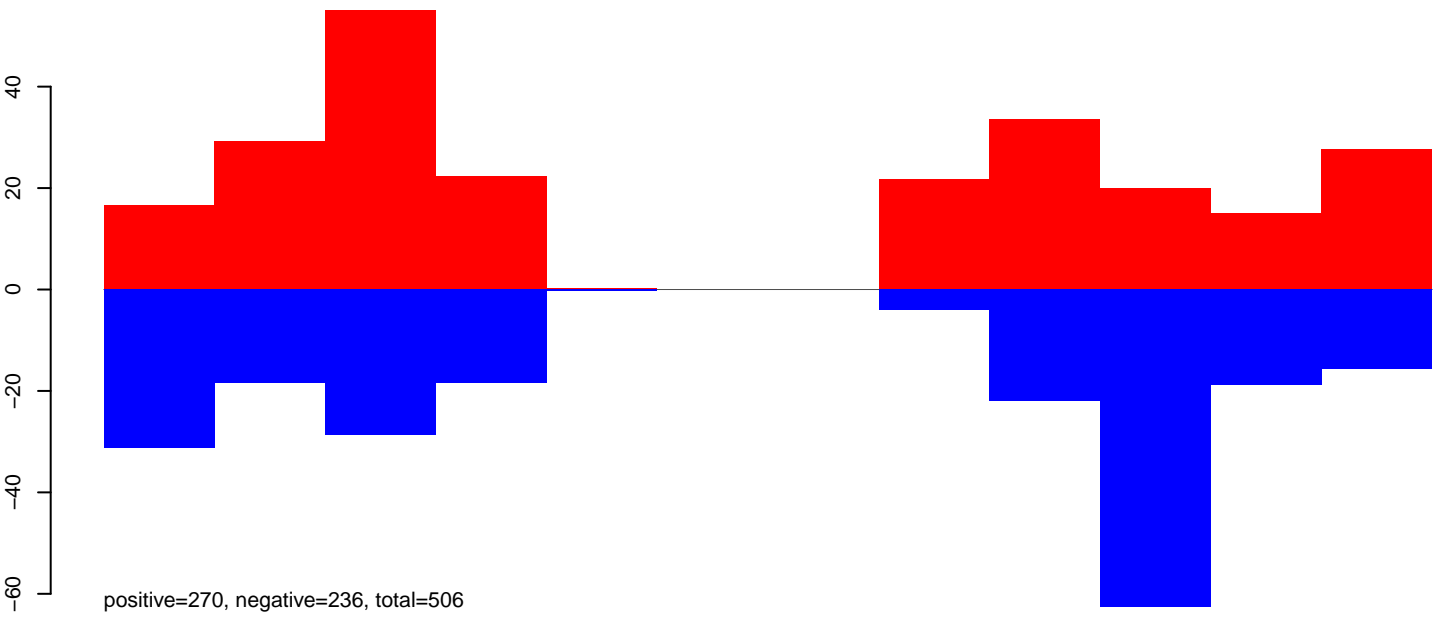
AnGam_Sua5b_cells.24_35.rep



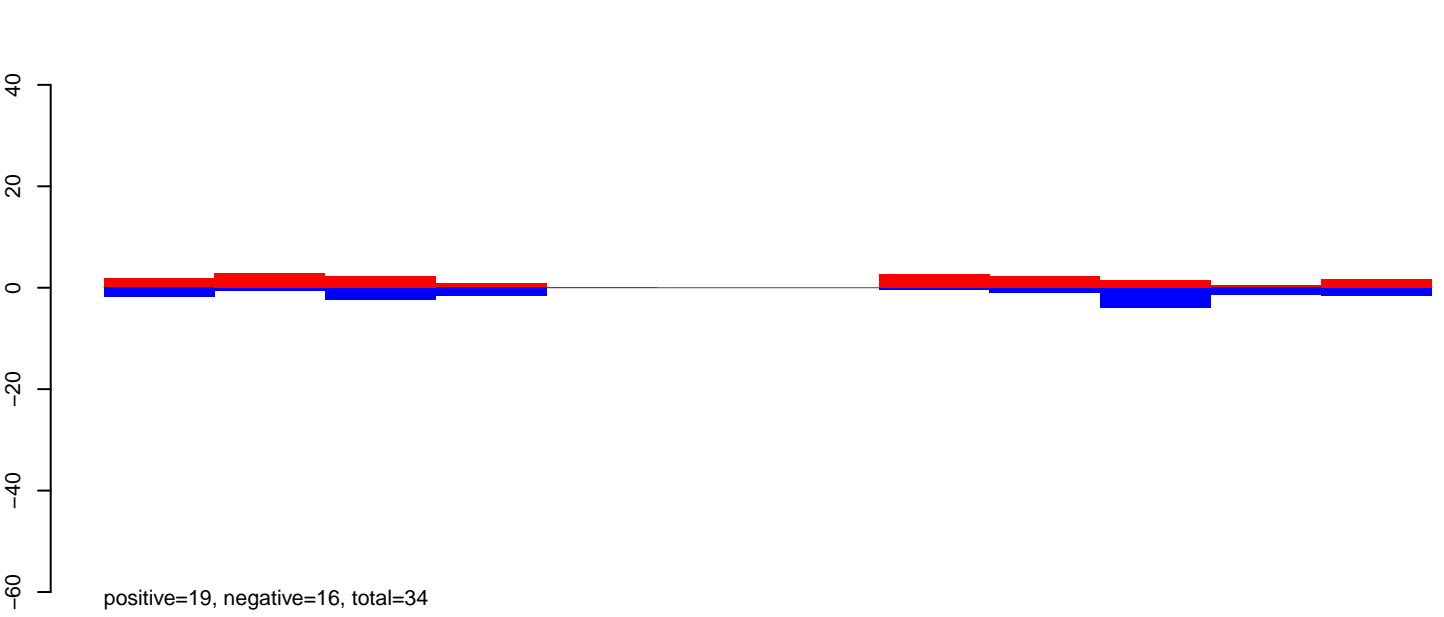
AnGam_Sua5b_cells.rep



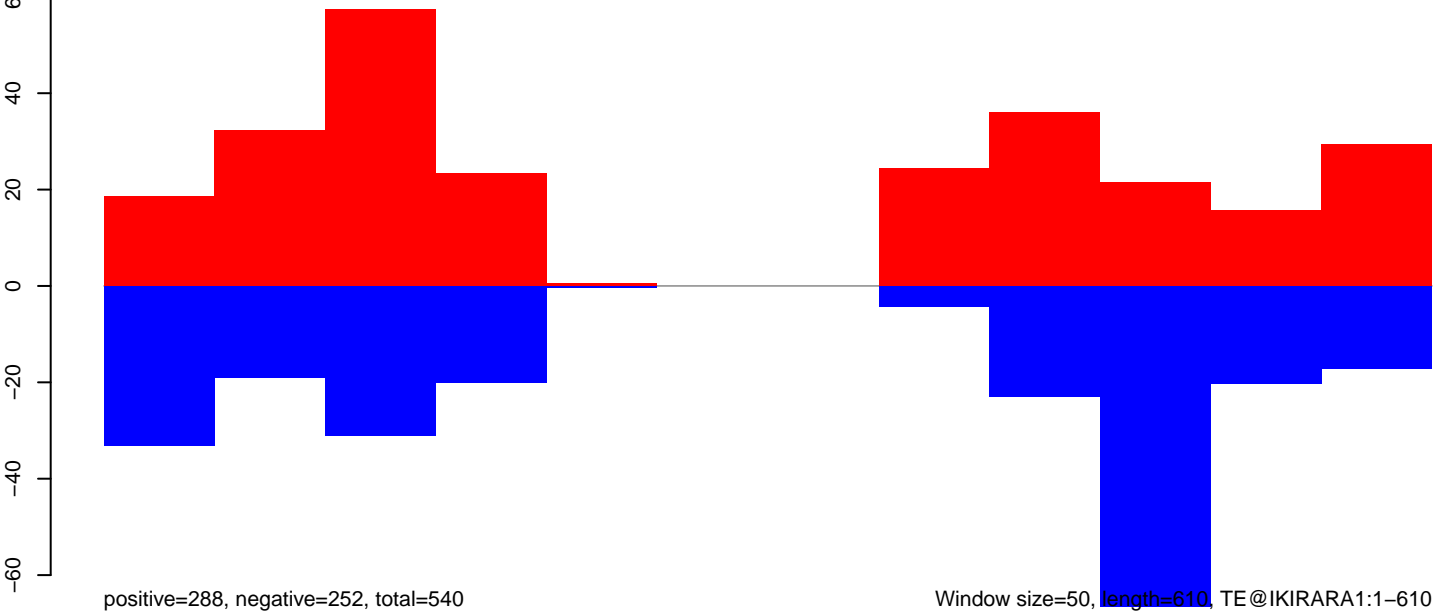
AnGam_Sua5b_cells.18_23.rep



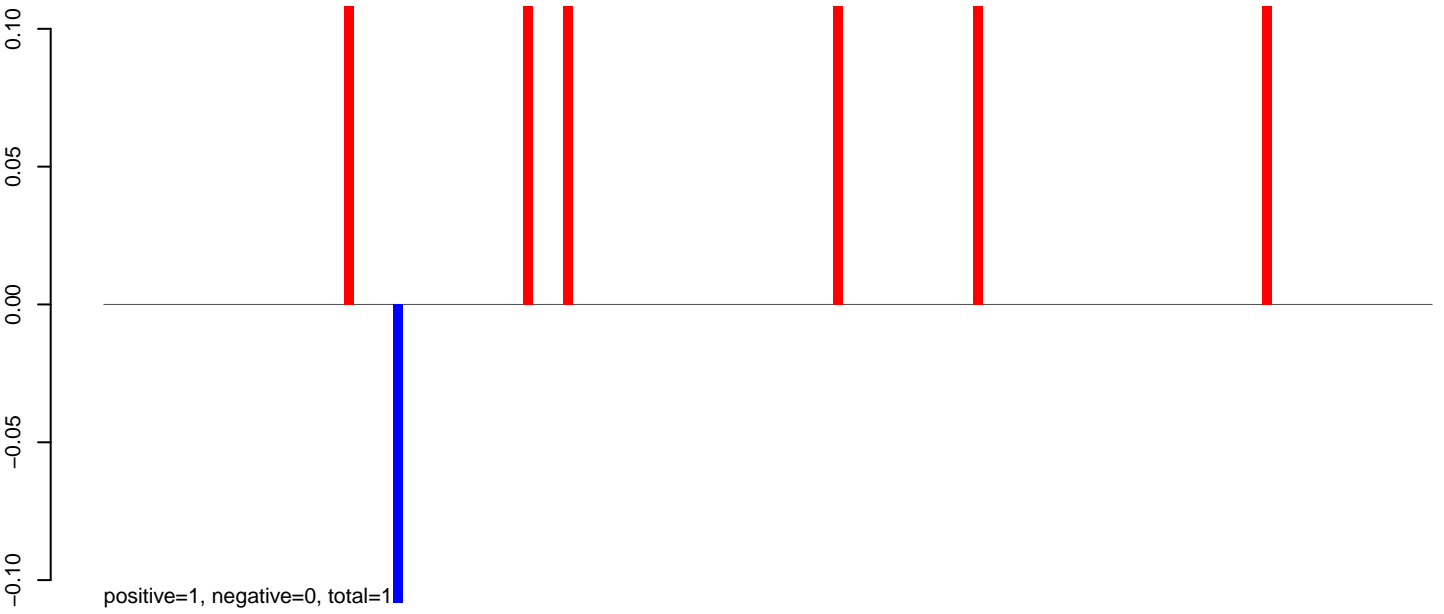
AnGam_Sua5b_cells.24_35.rep



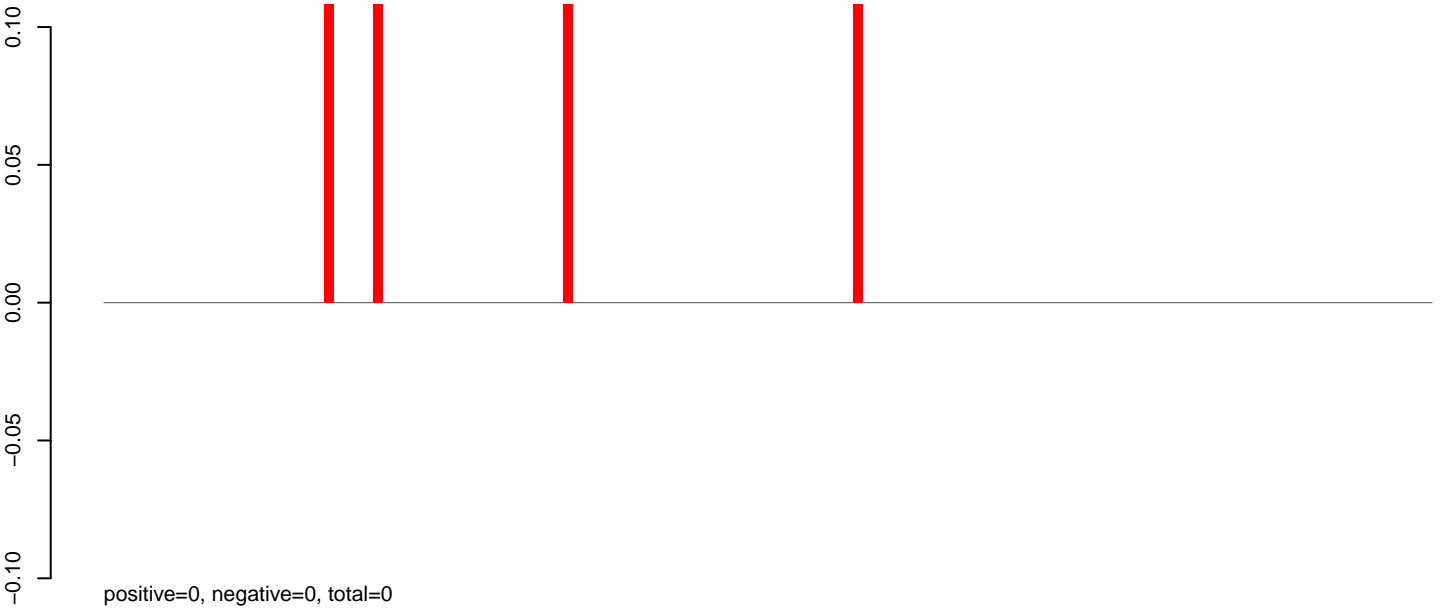
AnGam_Sua5b_cells.rep



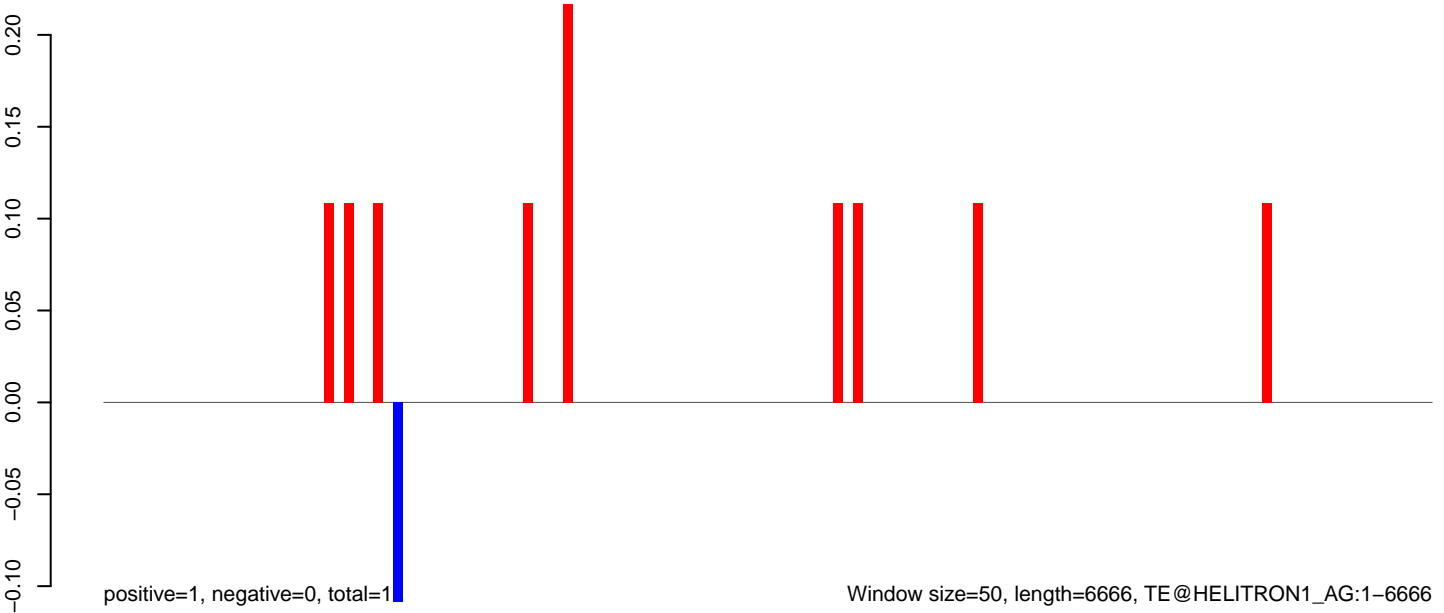
AnGam_Sua5b_cells.18_23.rep



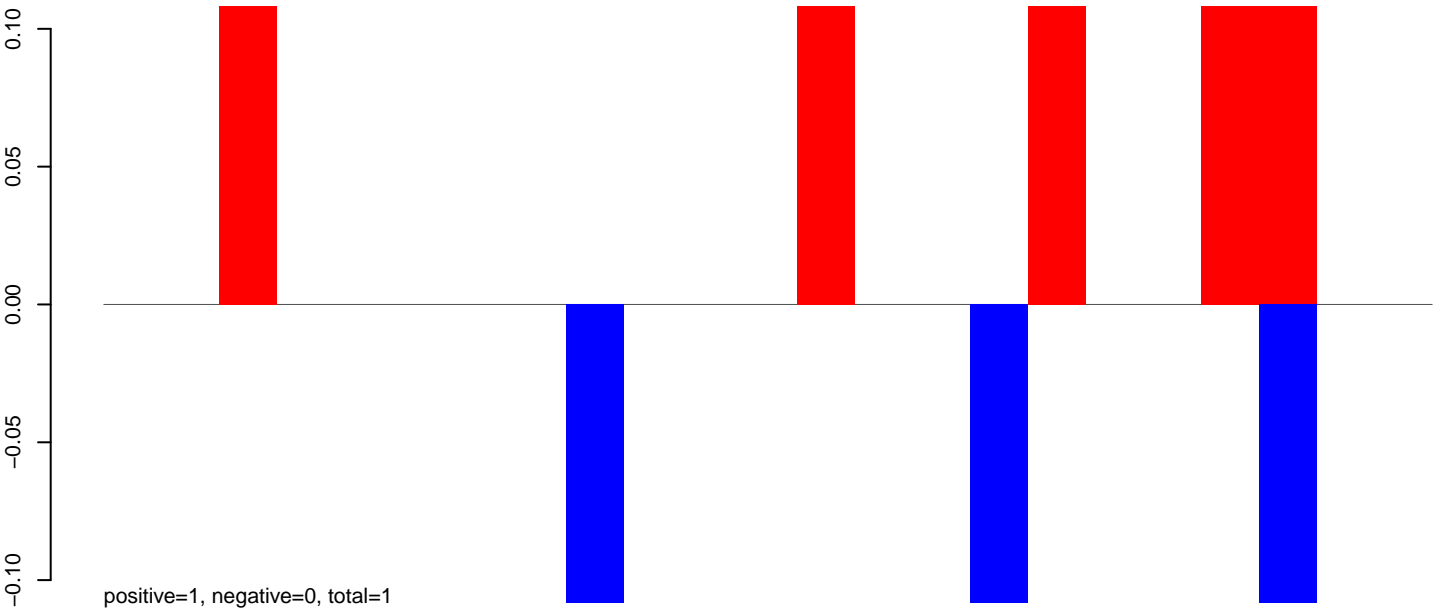
AnGam_Sua5b_cells.24_35.rep



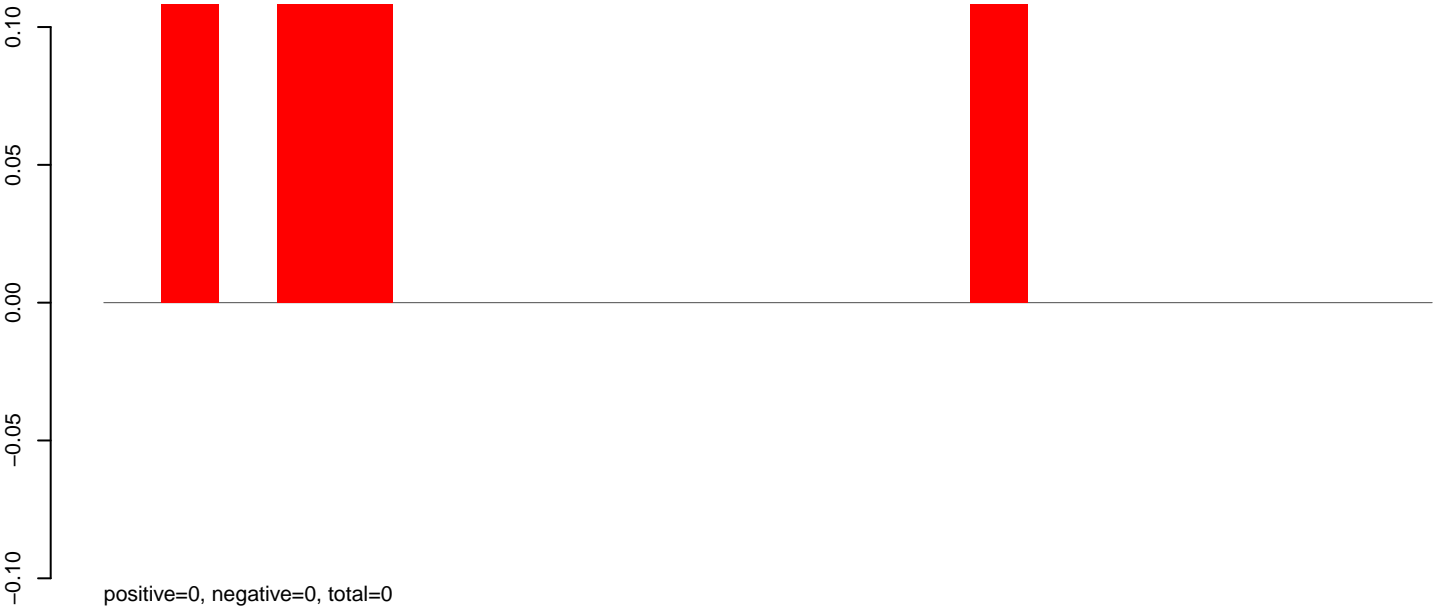
AnGam_Sua5b_cells.rep



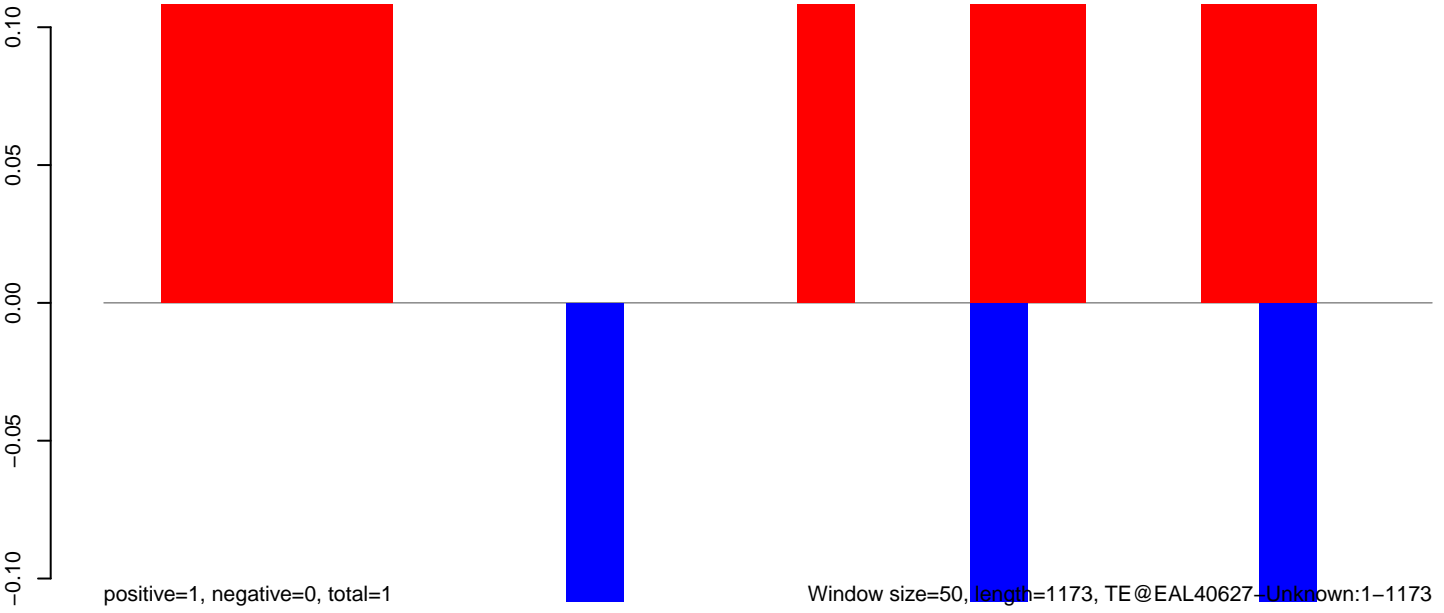
AnGam_Sua5b_cells.18_23.rep



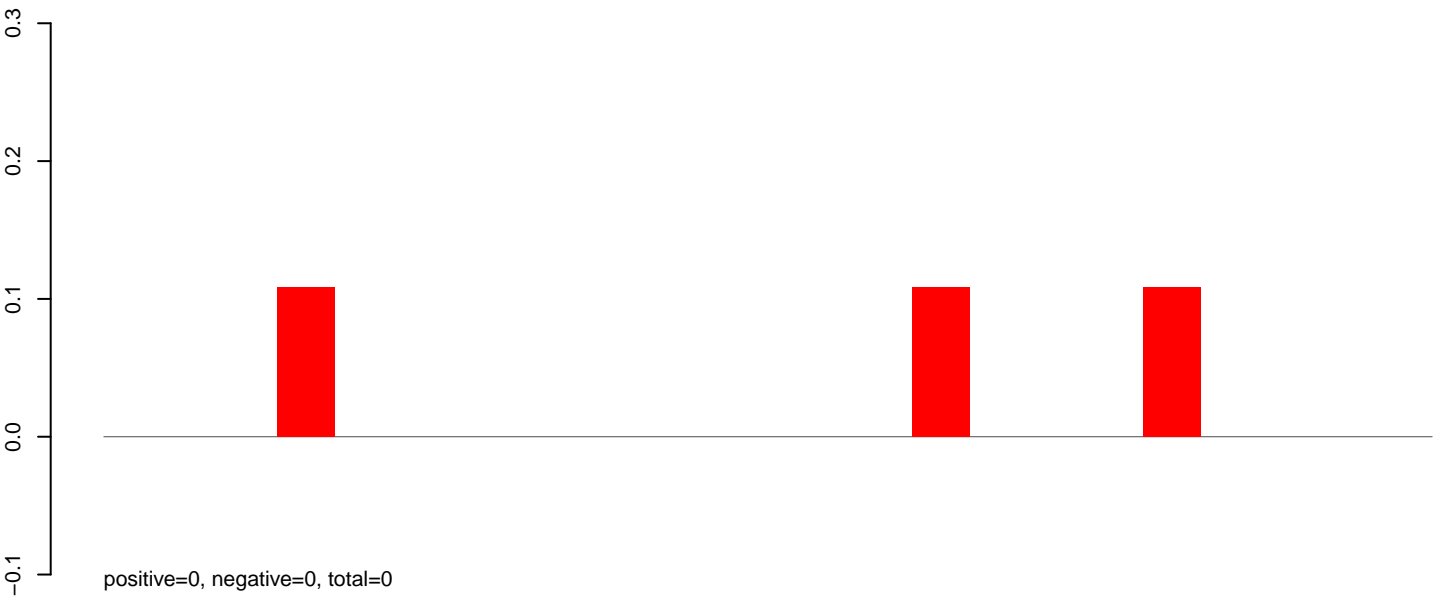
AnGam_Sua5b_cells.24_35.rep



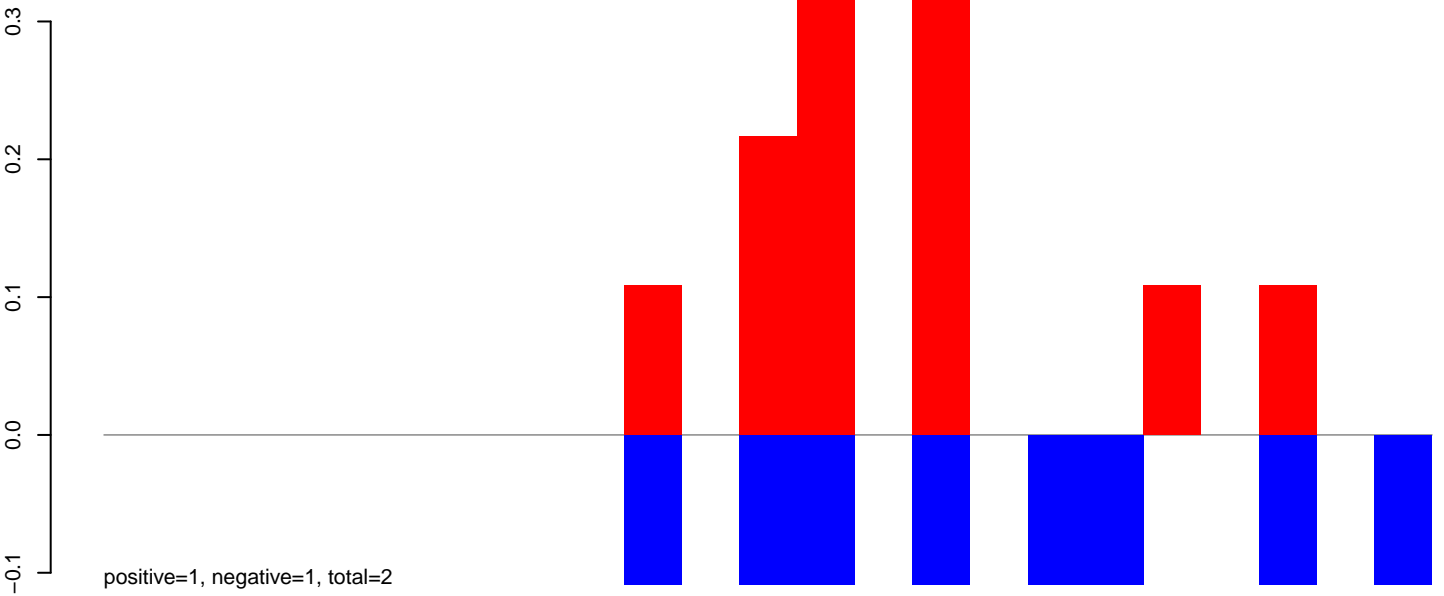
AnGam_Sua5b_cells.rep



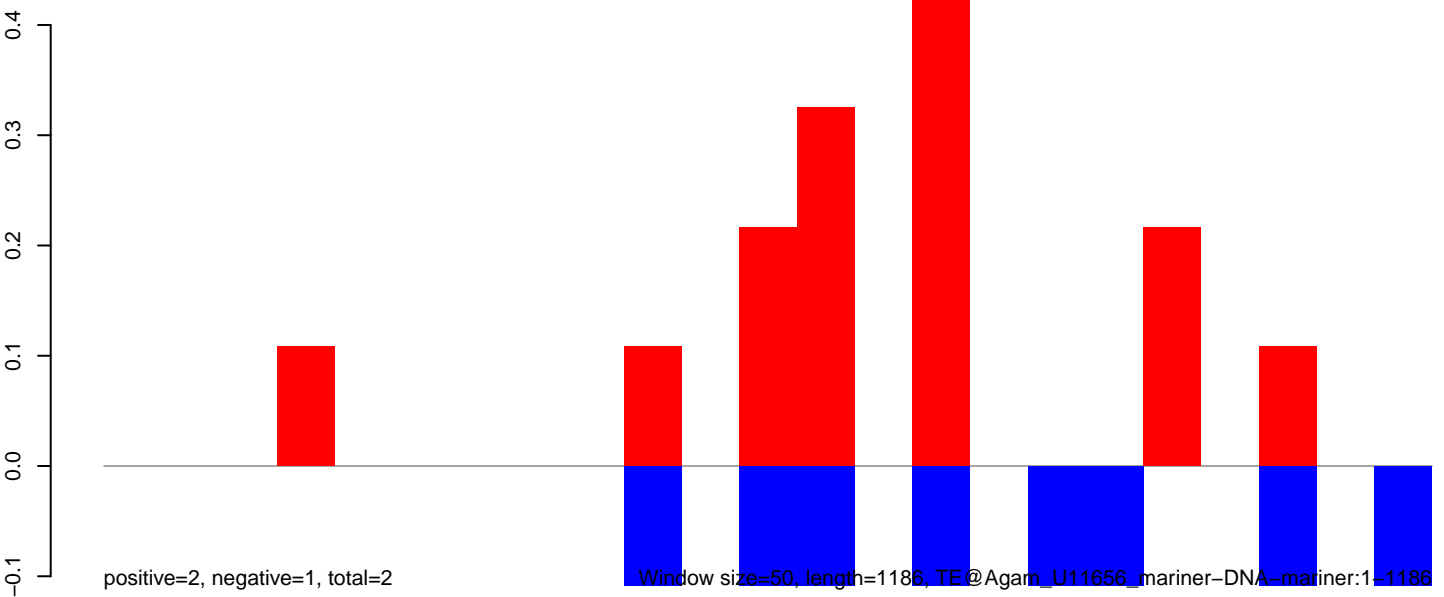
AnGam_Sua5b_cells.18_23.rep



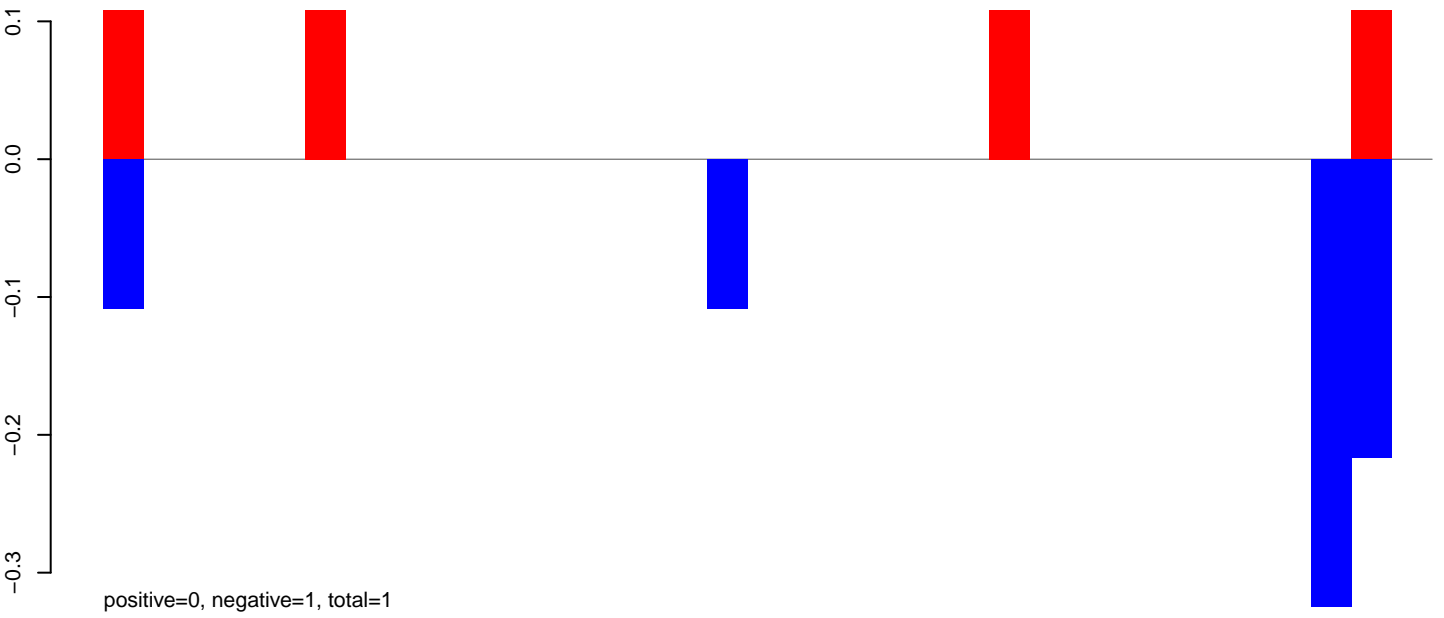
AnGam_Sua5b_cells.24_35.rep



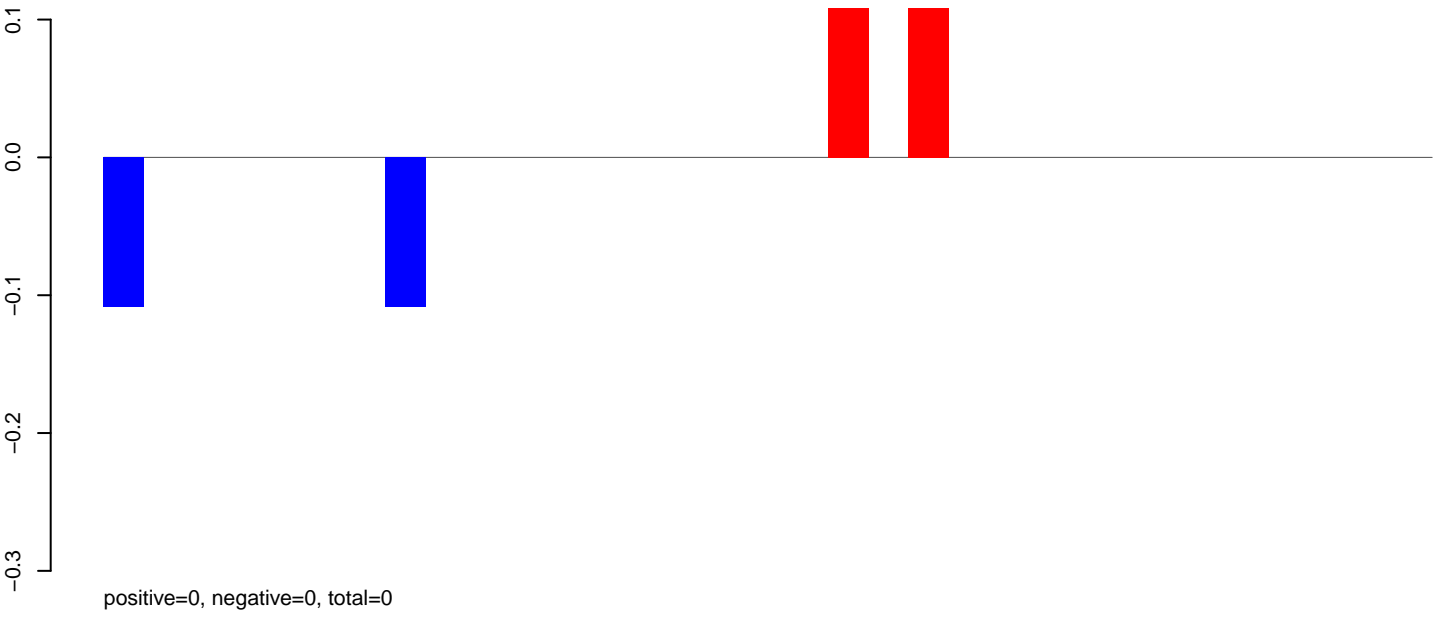
AnGam_Sua5b_cells.rep



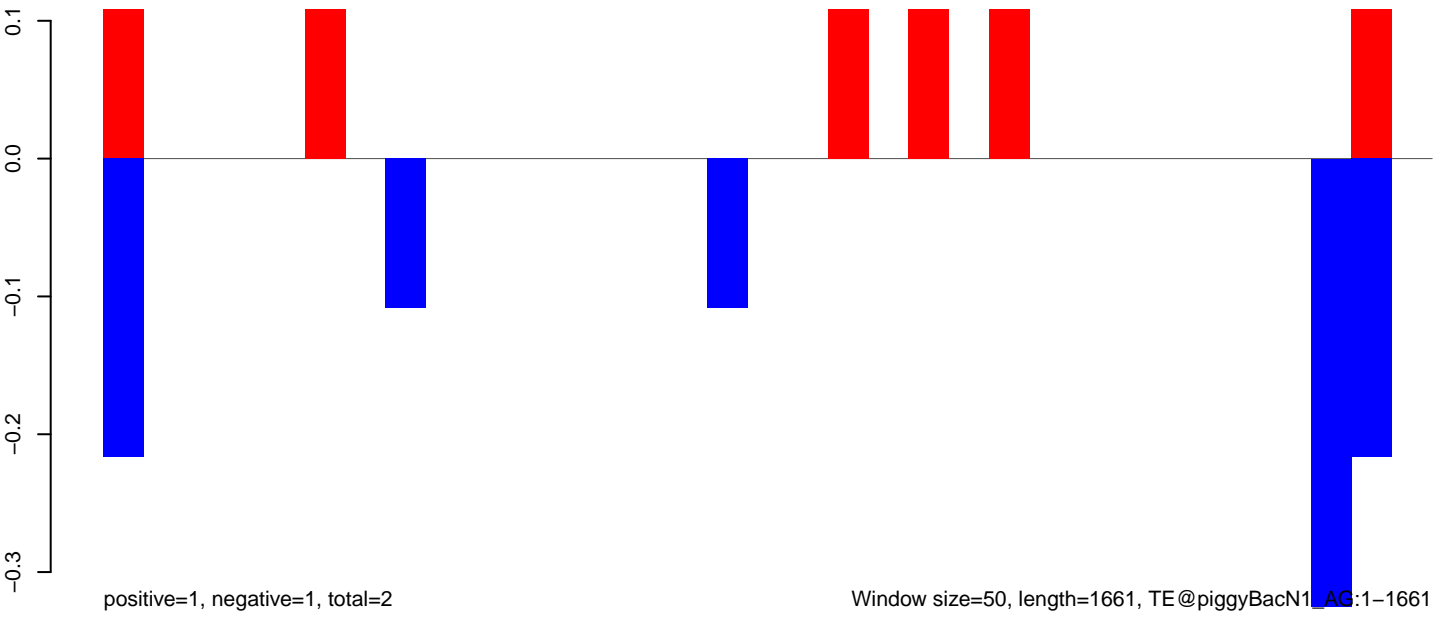
AnGam_Sua5b_cells.18_23.rep



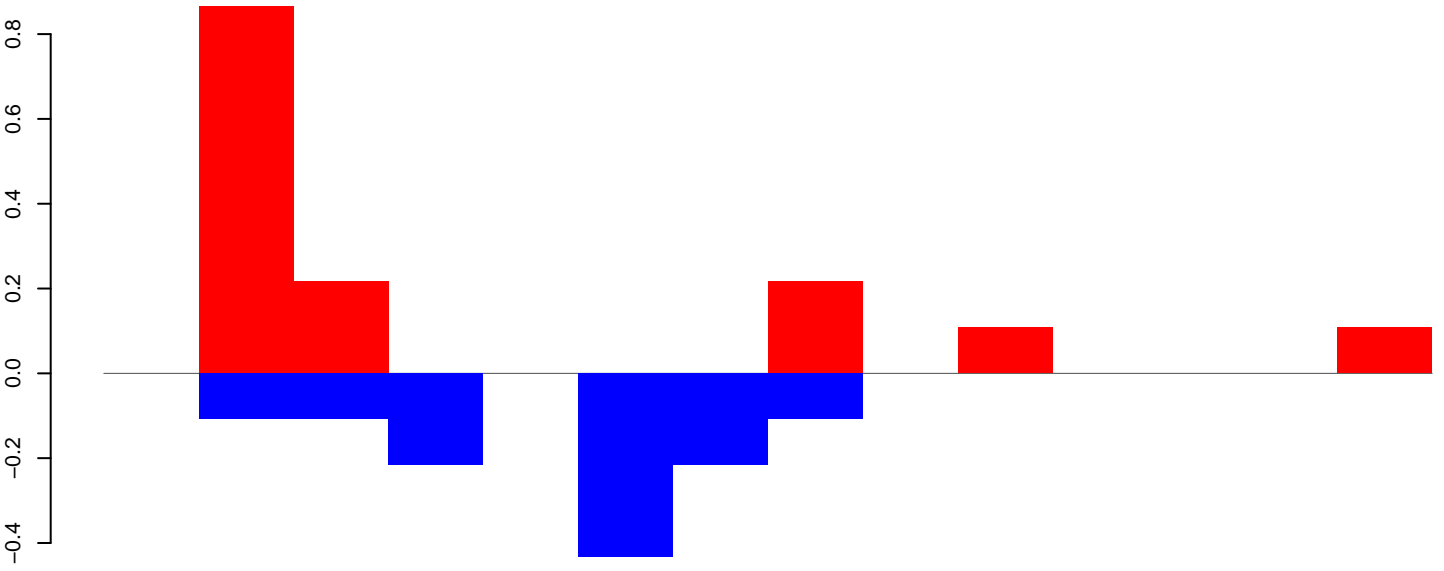
AnGam_Sua5b_cells.24_35.rep



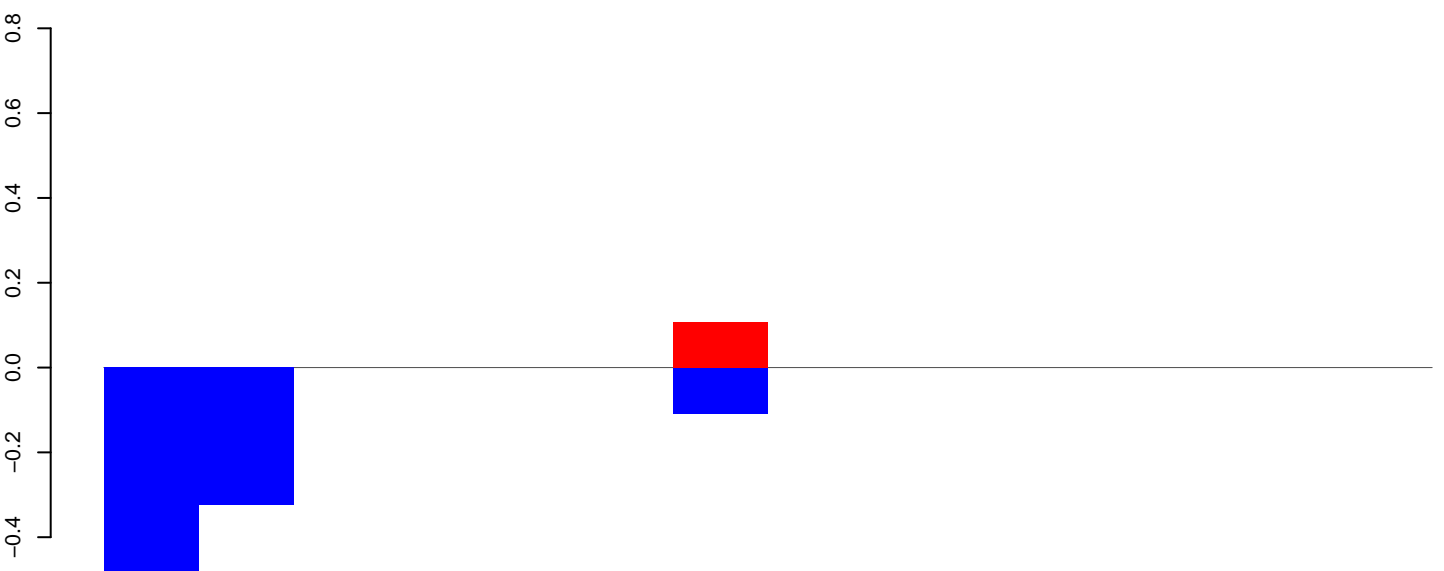
AnGam_Sua5b_cells.rep



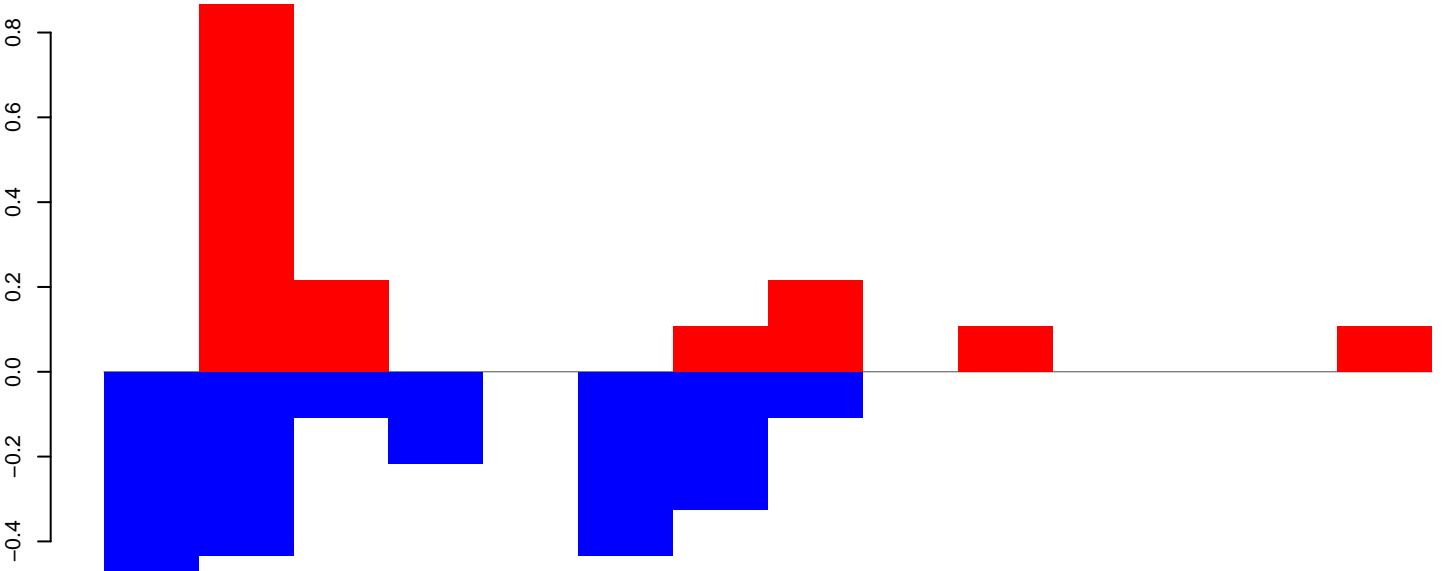
AnGam_Sua5b_cells.18_23.rep



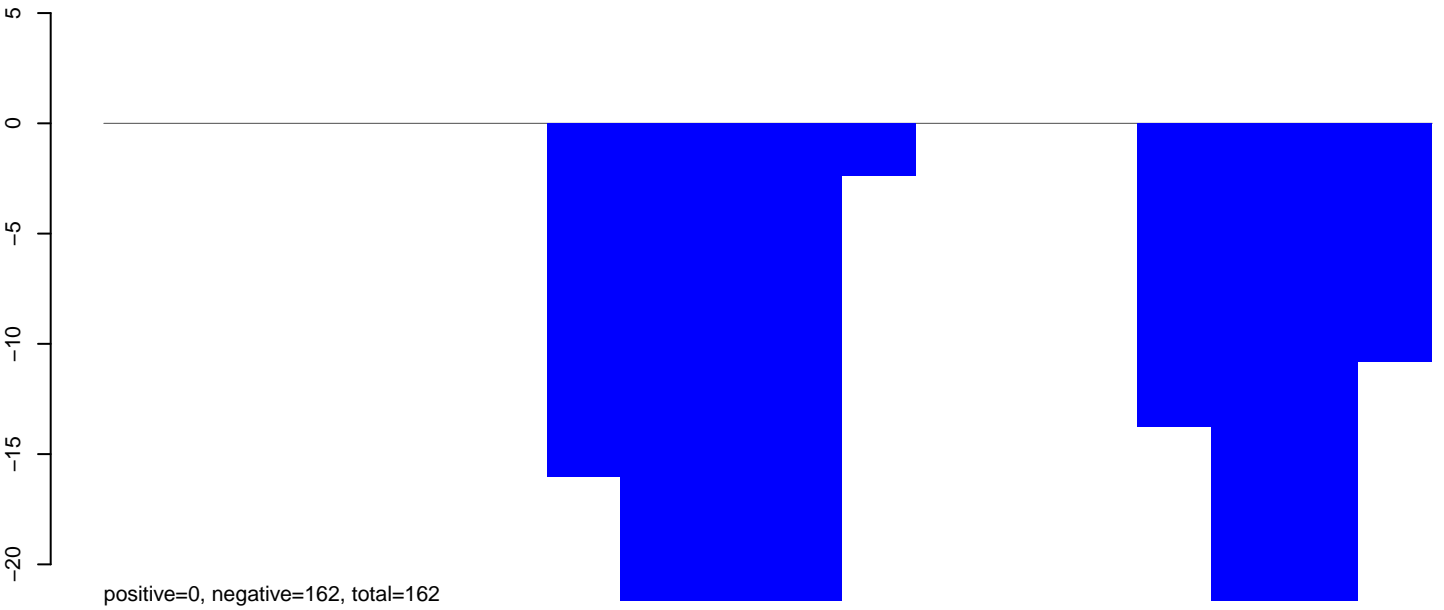
AnGam_Sua5b_cells.24_35.rep



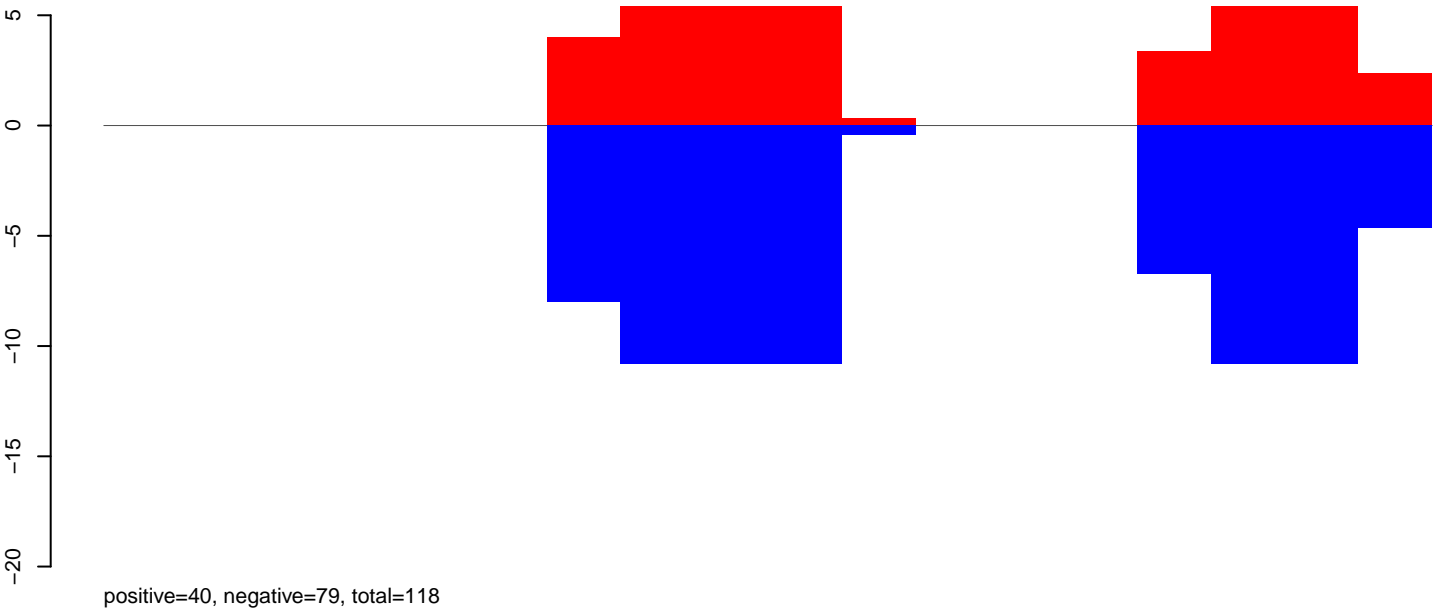
AnGam_Sua5b_cells.rep



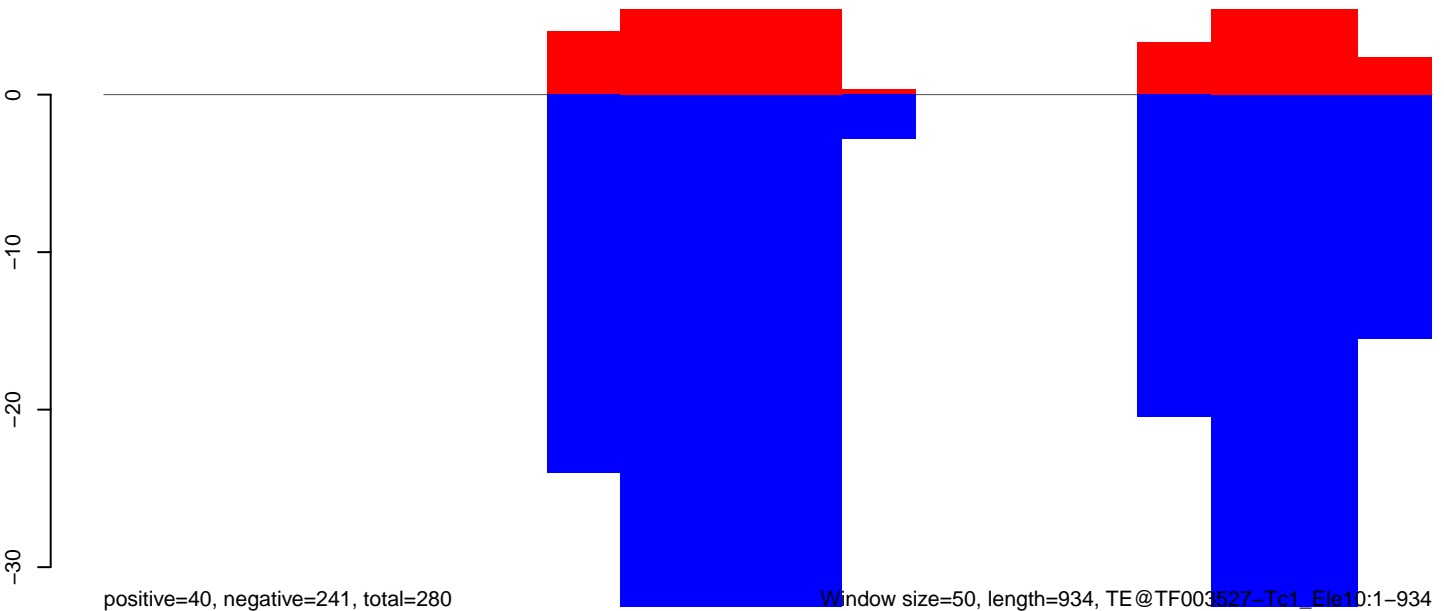
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



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

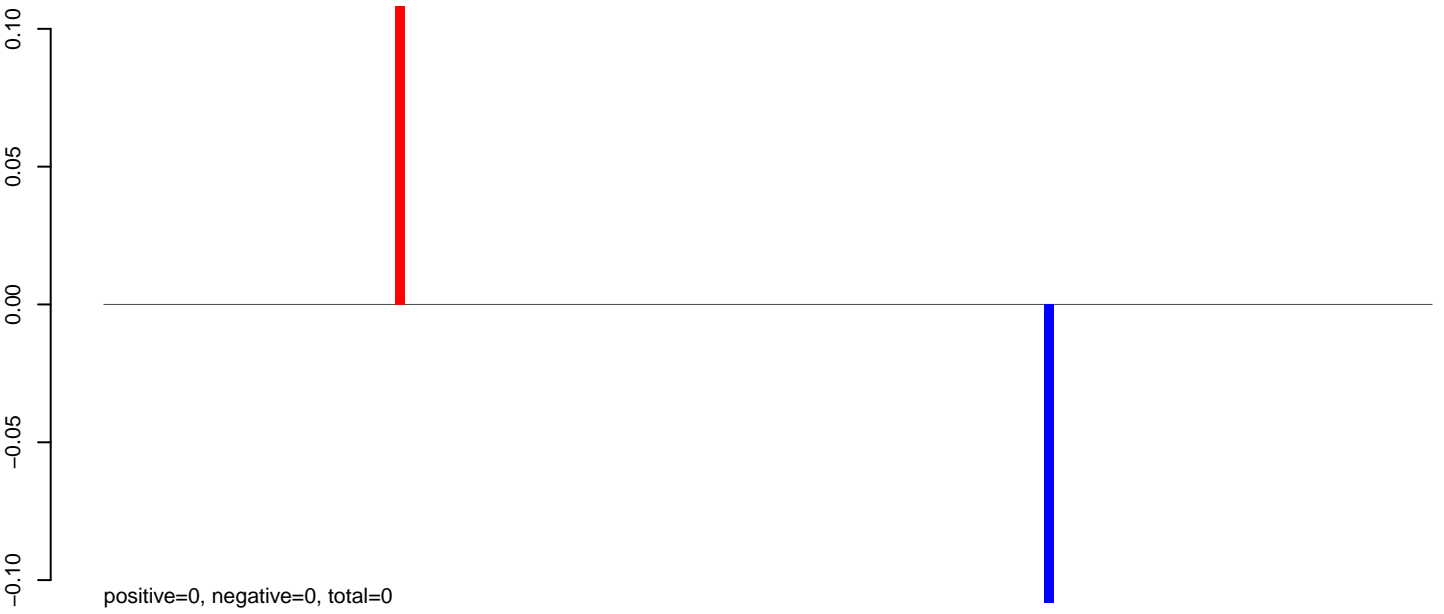
200

400

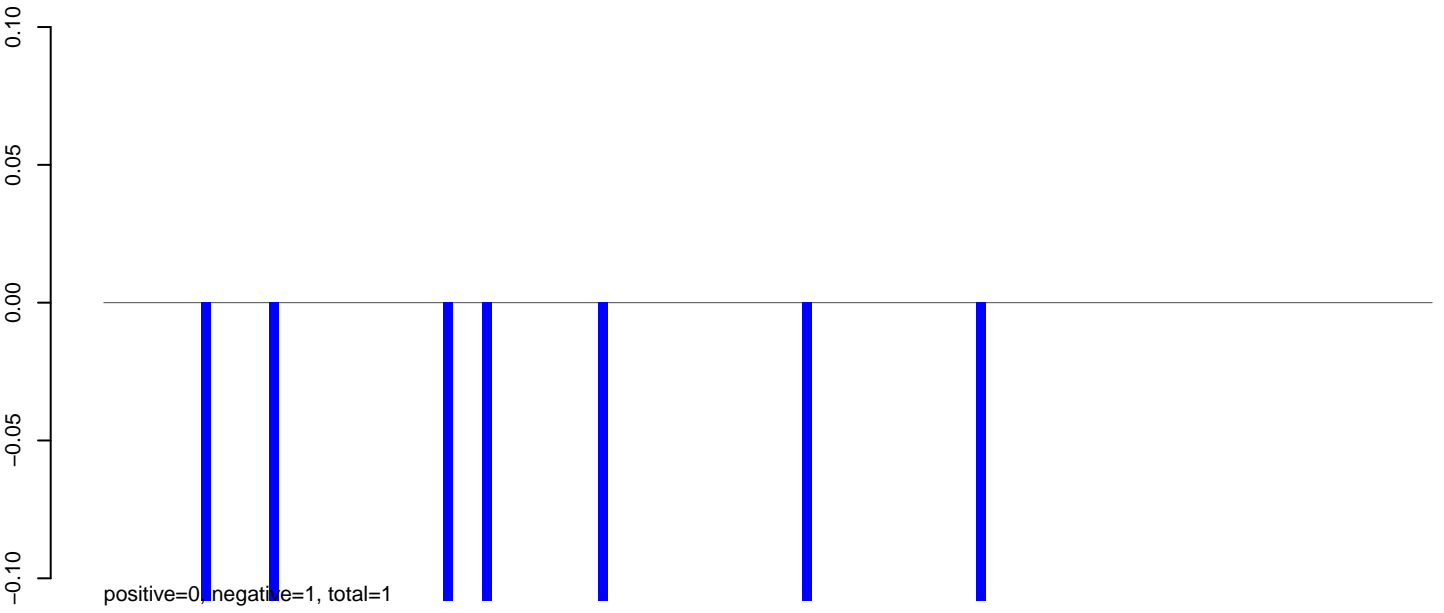
600

800

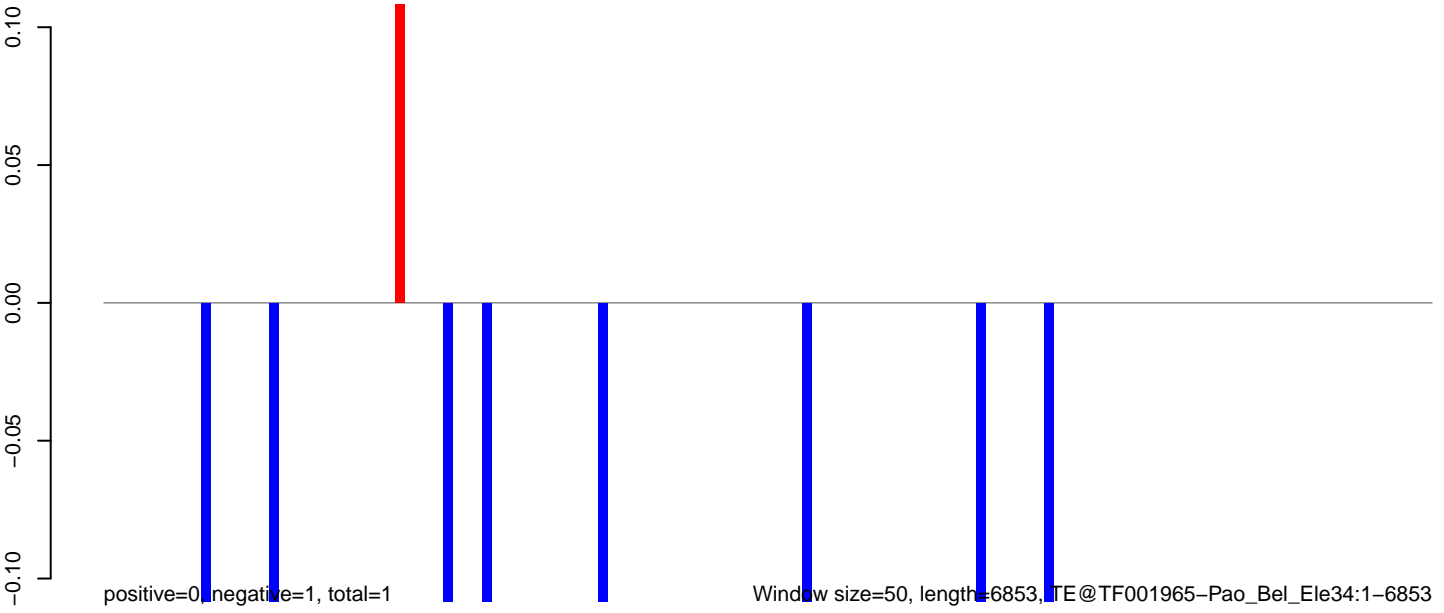
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

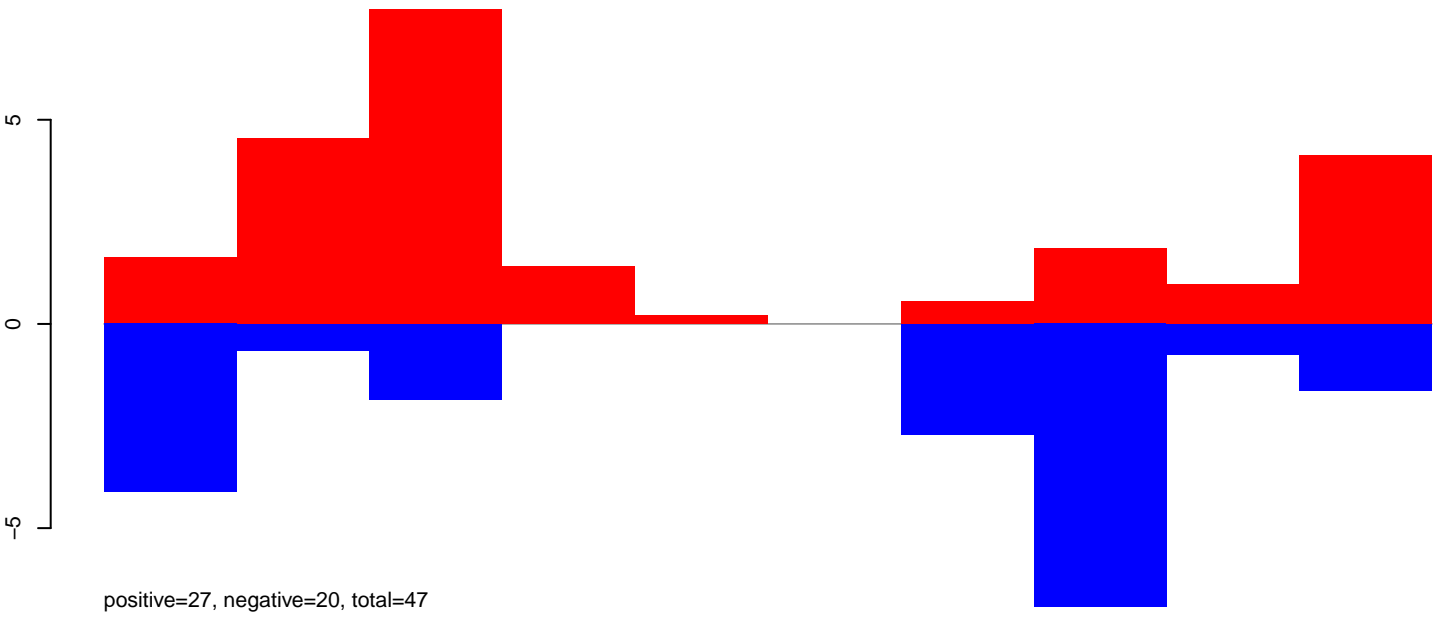


AnGam_Sua5b_cells.rep

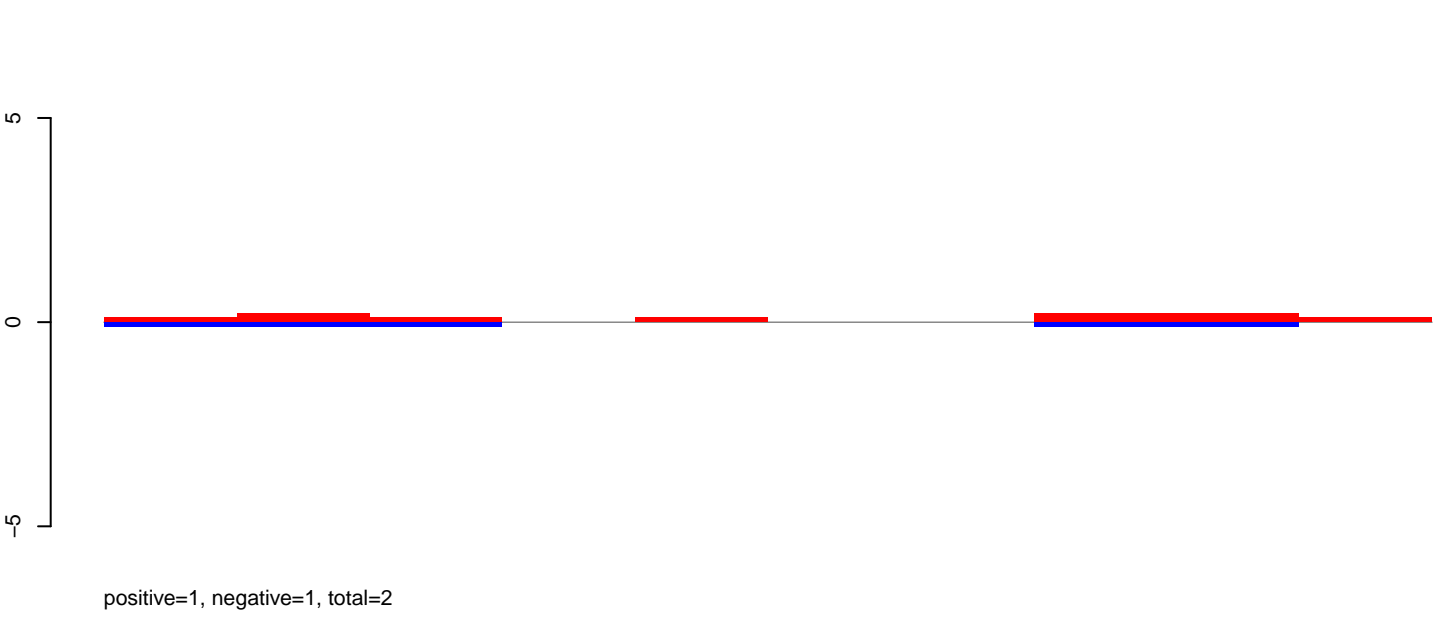


0 1000 2000 3000 4000 5000 6000 7000

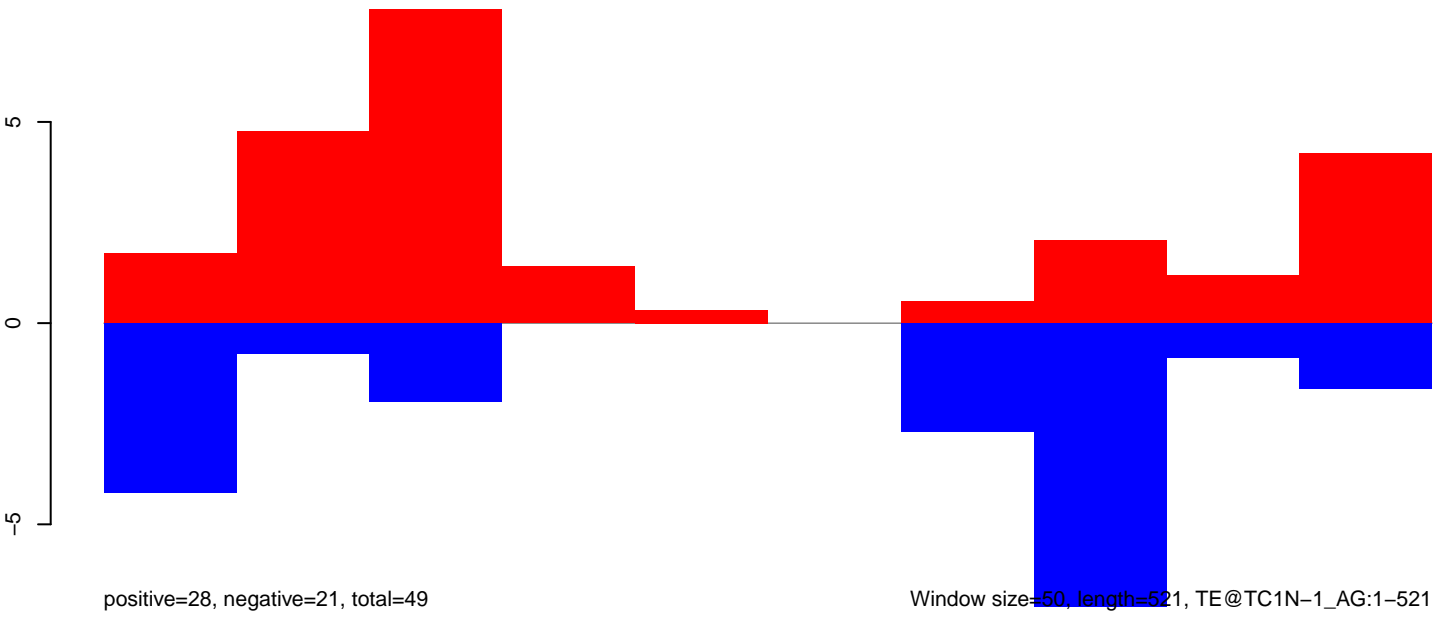
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



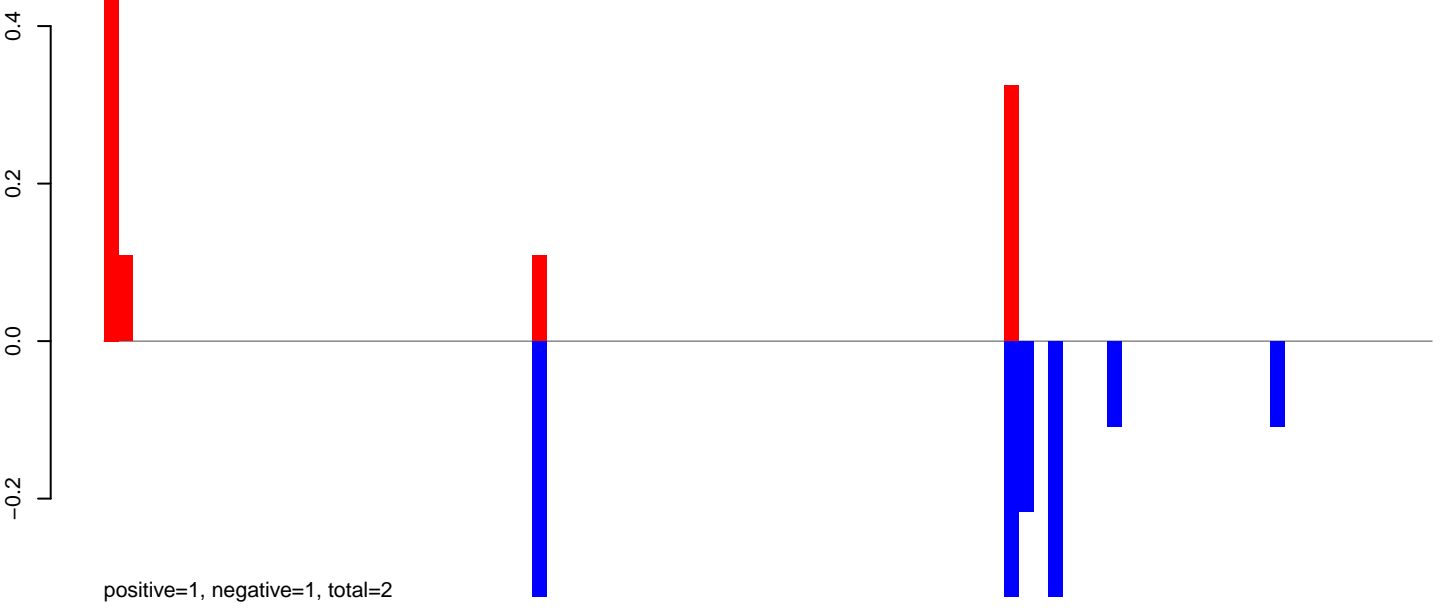
AnGam_Sua5b_cells.rep



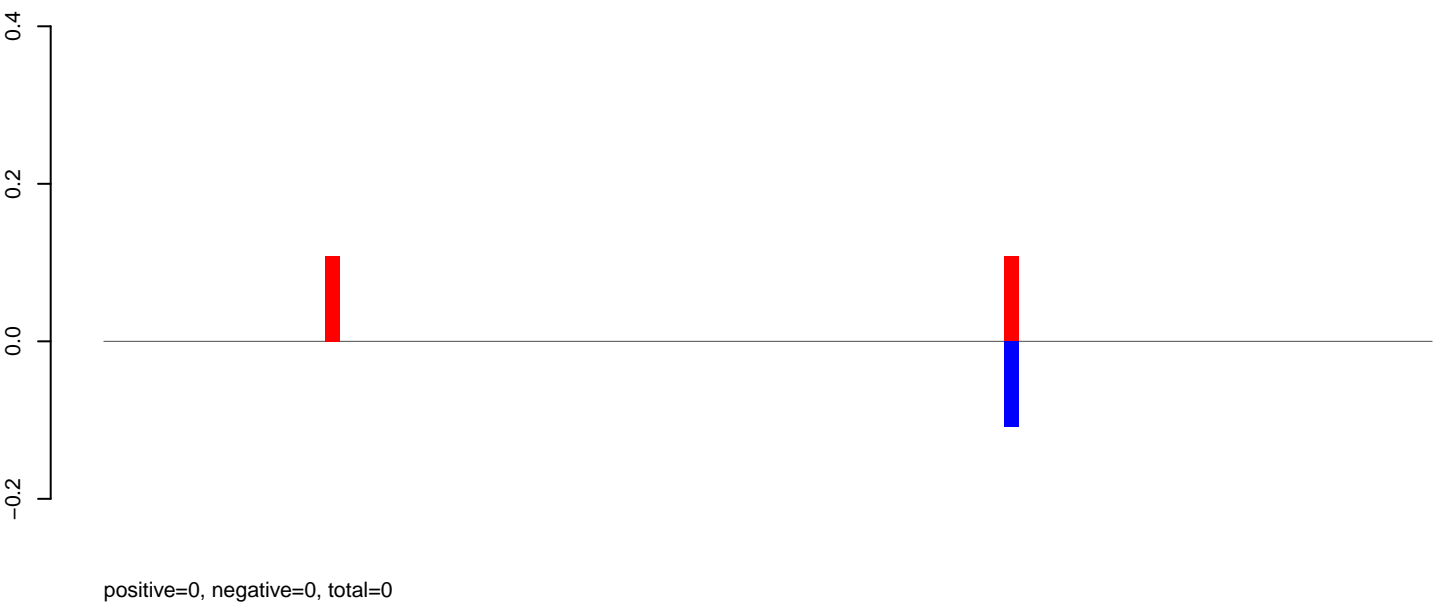
Window size=50, length=521, TE@TC1N-1_AG:1-521

100 200 300 400 500

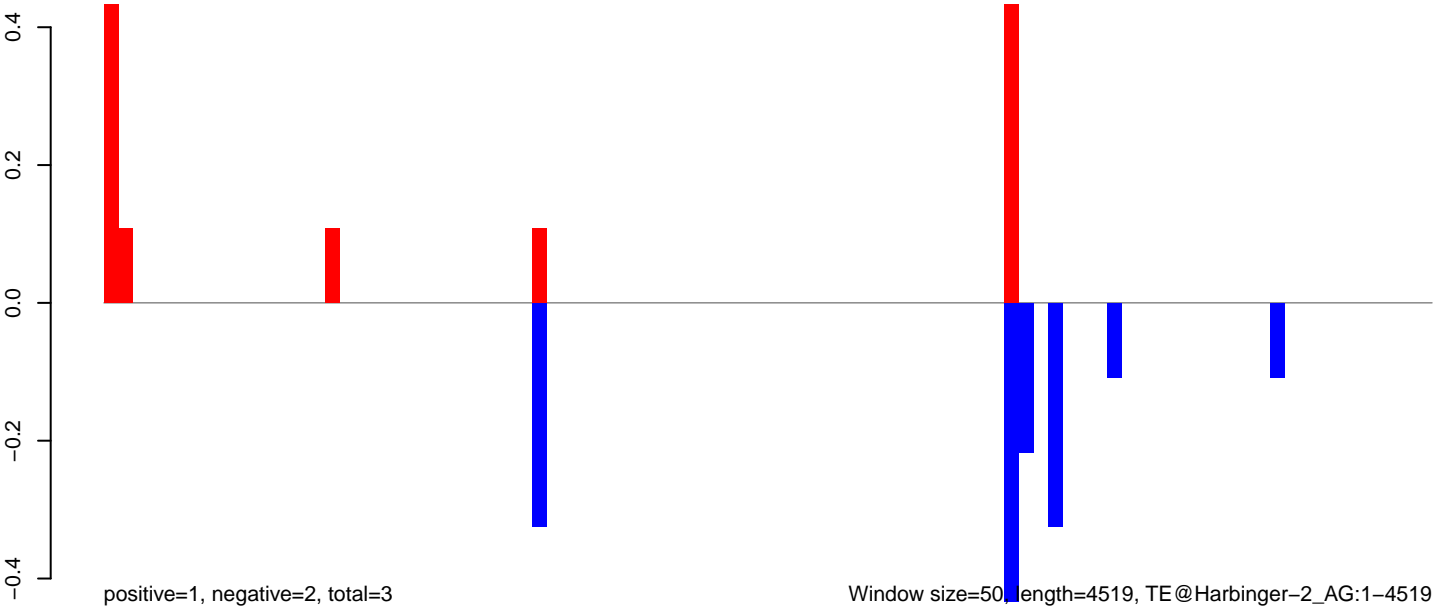
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



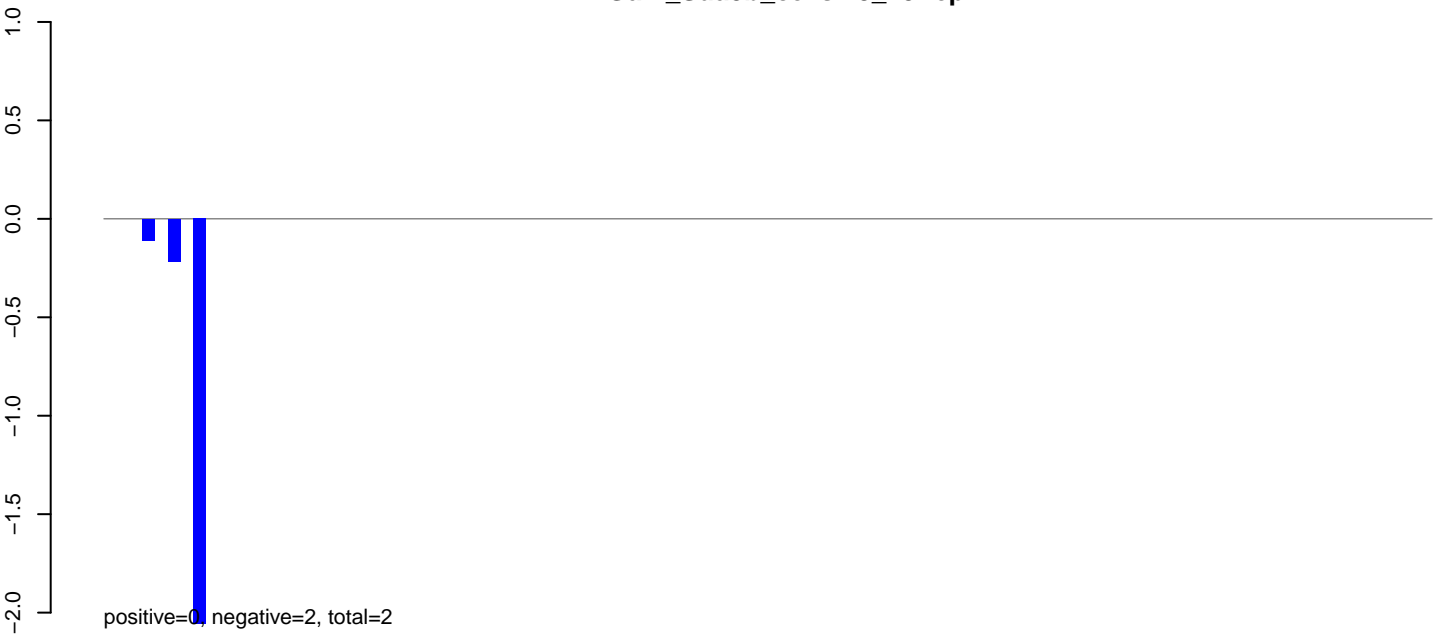
AnGam_Sua5b_cells.rep



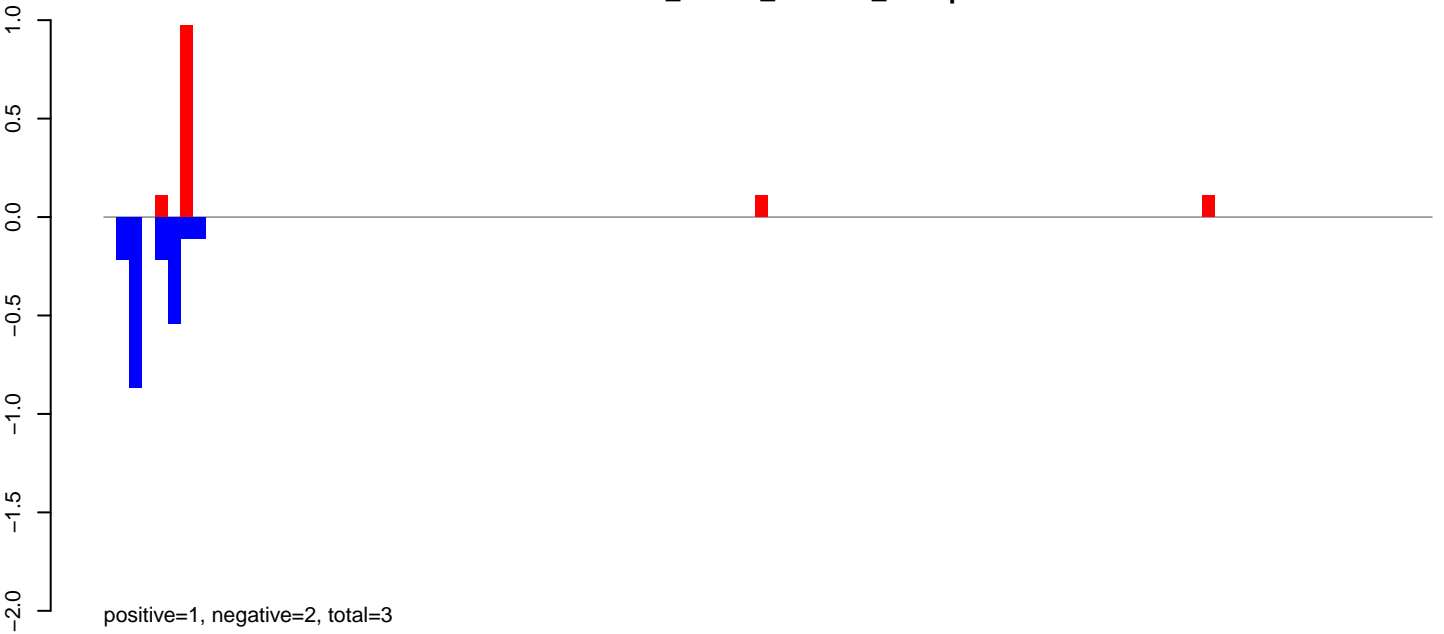
Window size=50, length=4519, TE@Harbinger-2_AG:1-4519

0 1000 2000 3000 4000

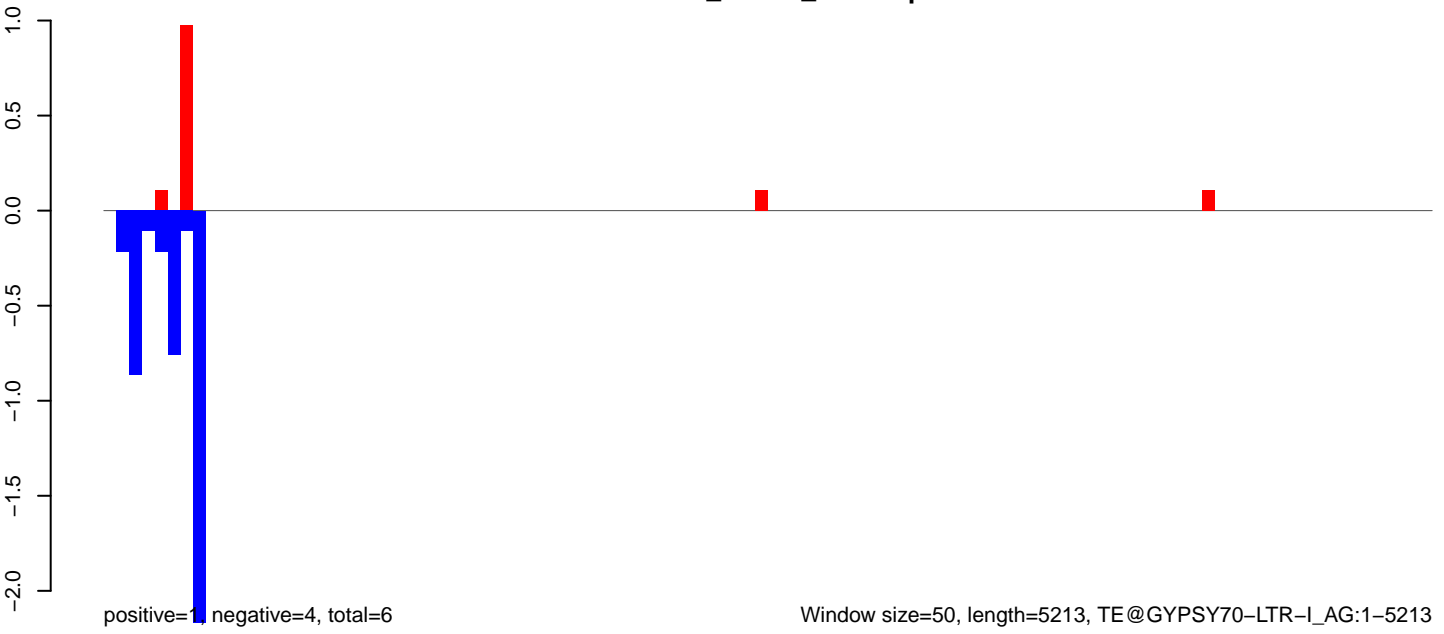
AnGam_Sua5b_cells.18_23.rep



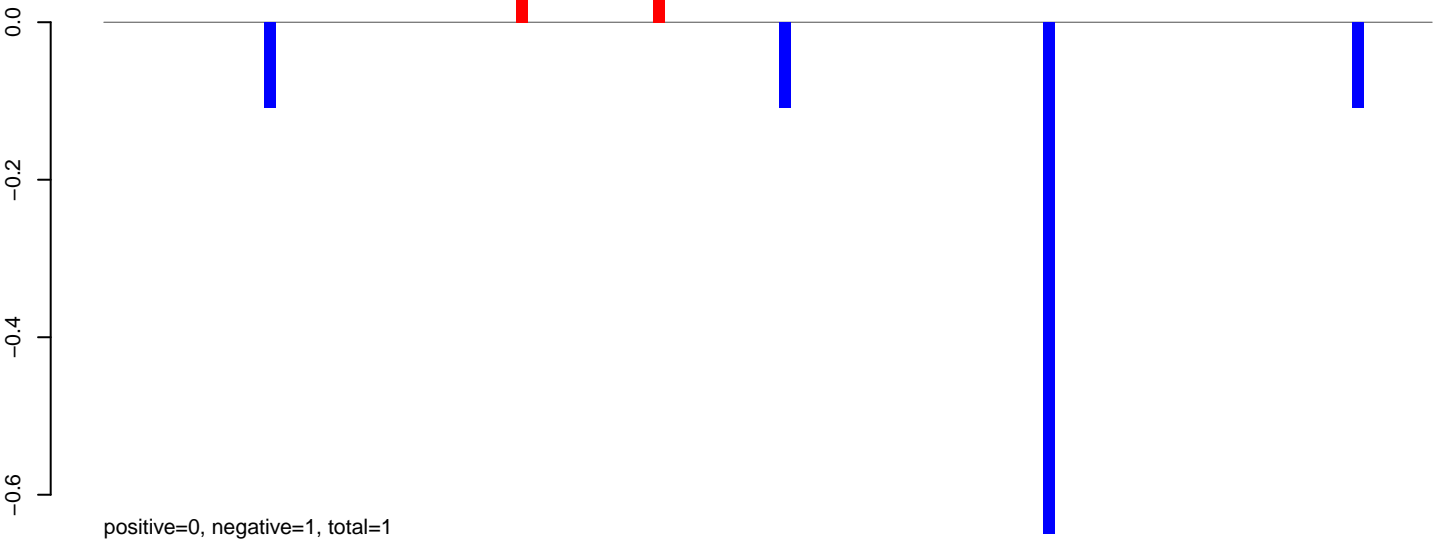
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



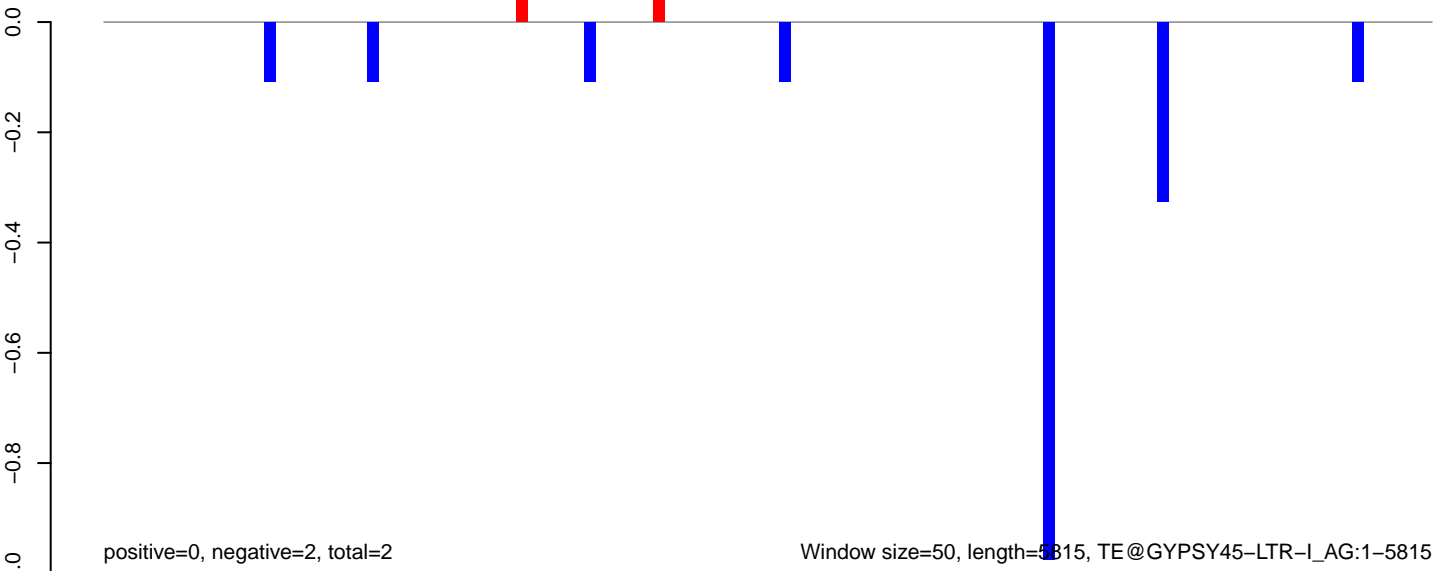
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

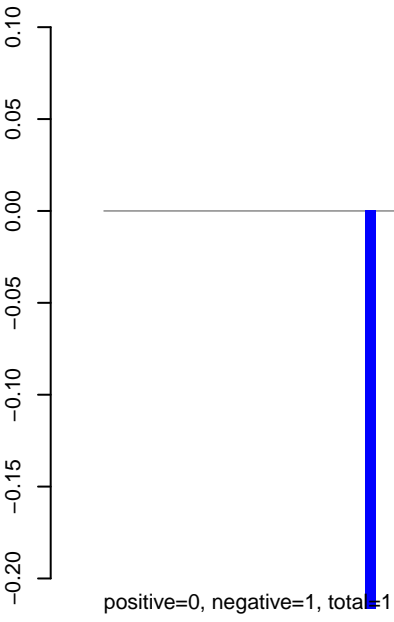


AnGam_Sua5b_cells.rep

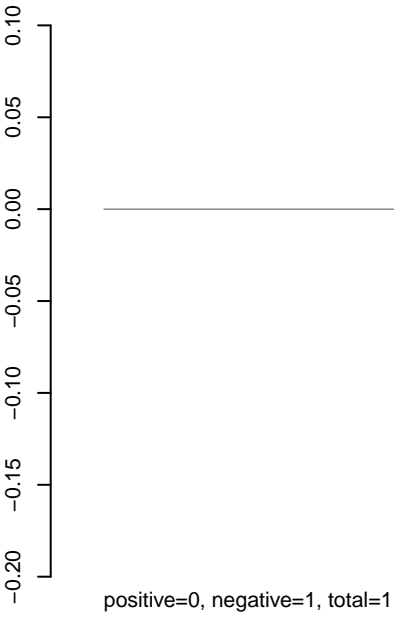


Window size=50, length=5815, TE@GYPSY45-LTR-I_AG:1-5815

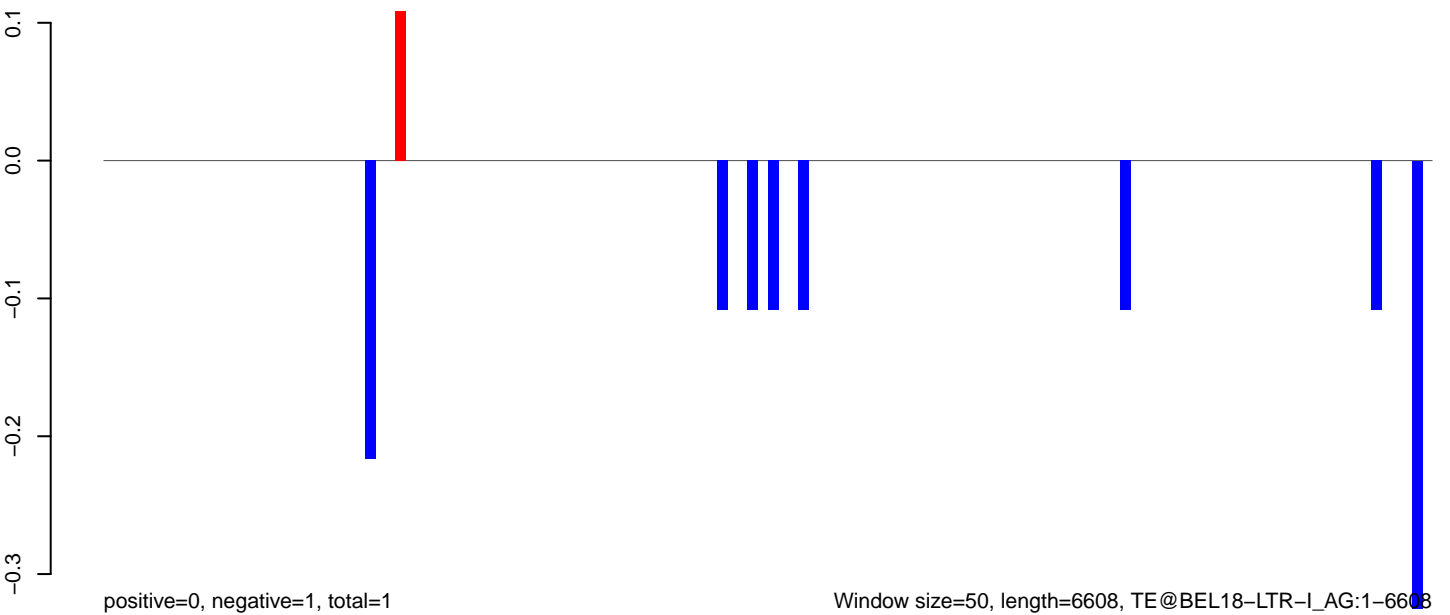
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

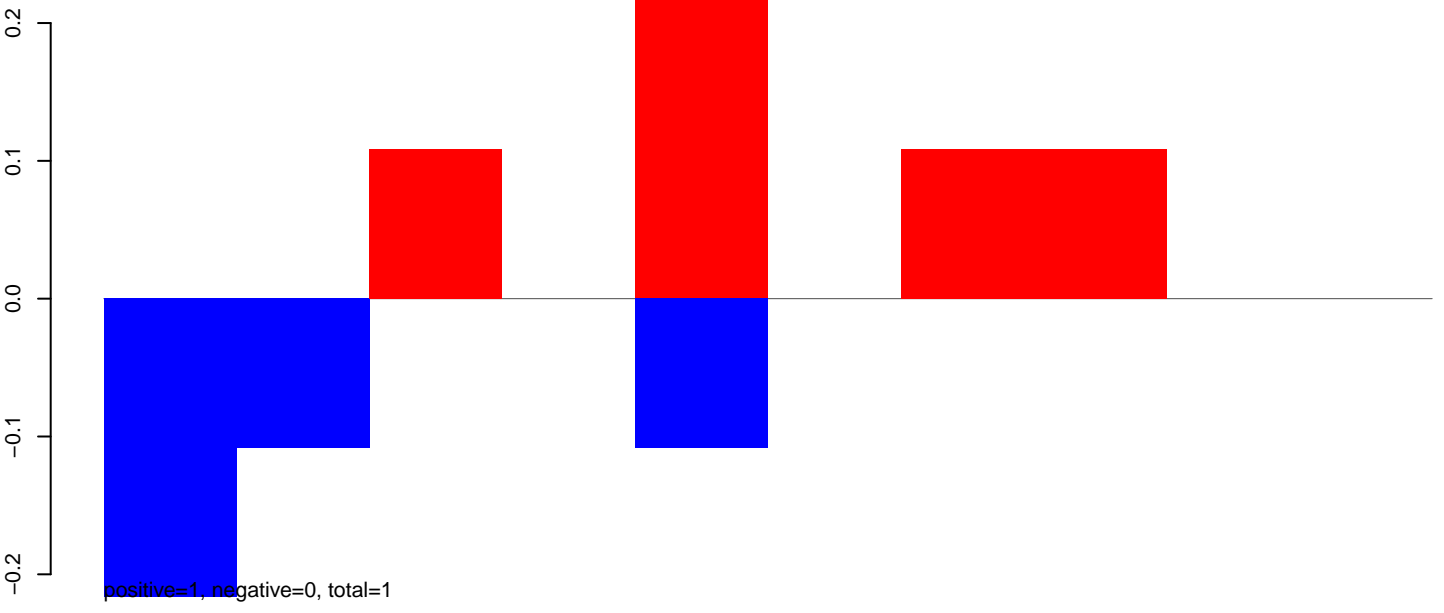


Window size=50, length=6608, TE@BEL18-LTR-I_AG:1-6608

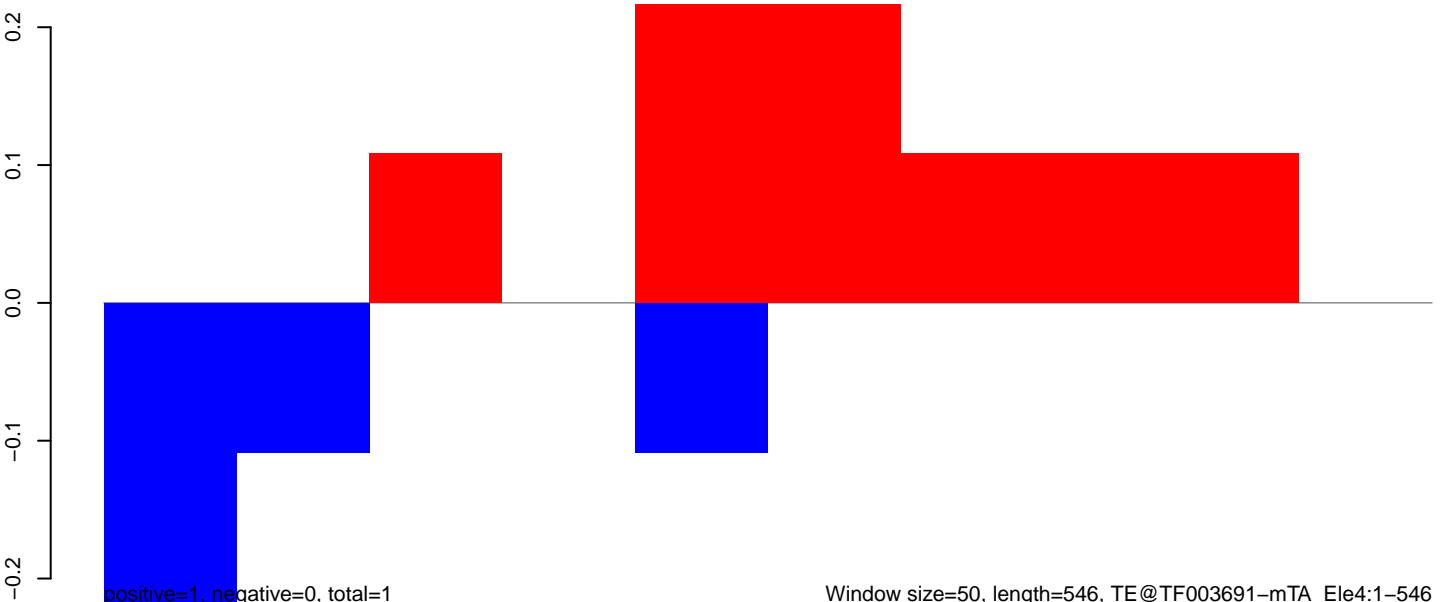
AnGam_Sua5b_cells.18_23.rep



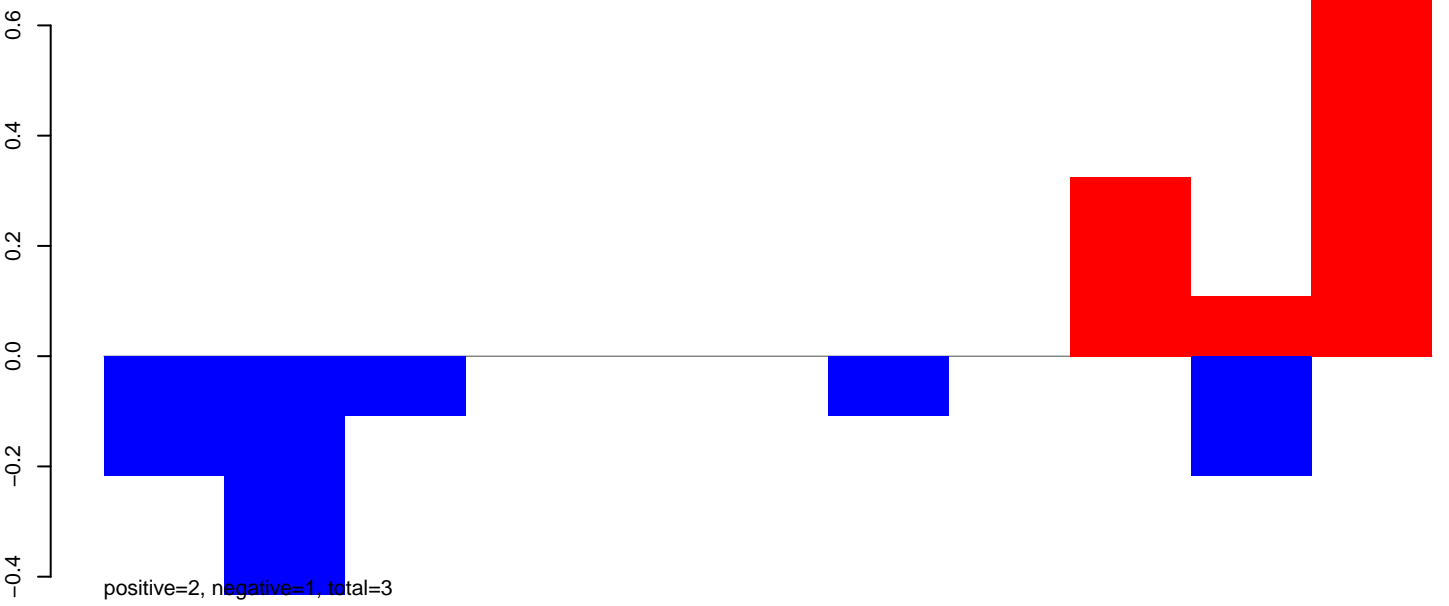
AnGam_Sua5b_cells.24_35.rep



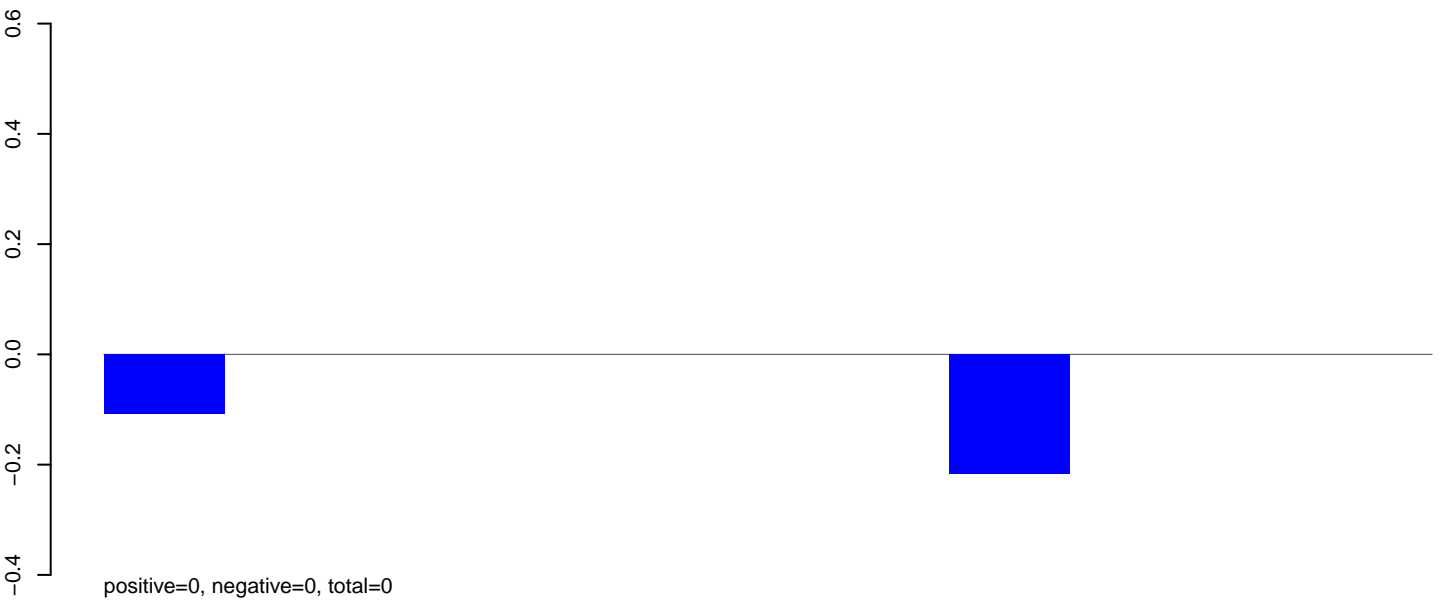
AnGam_Sua5b_cells.rep



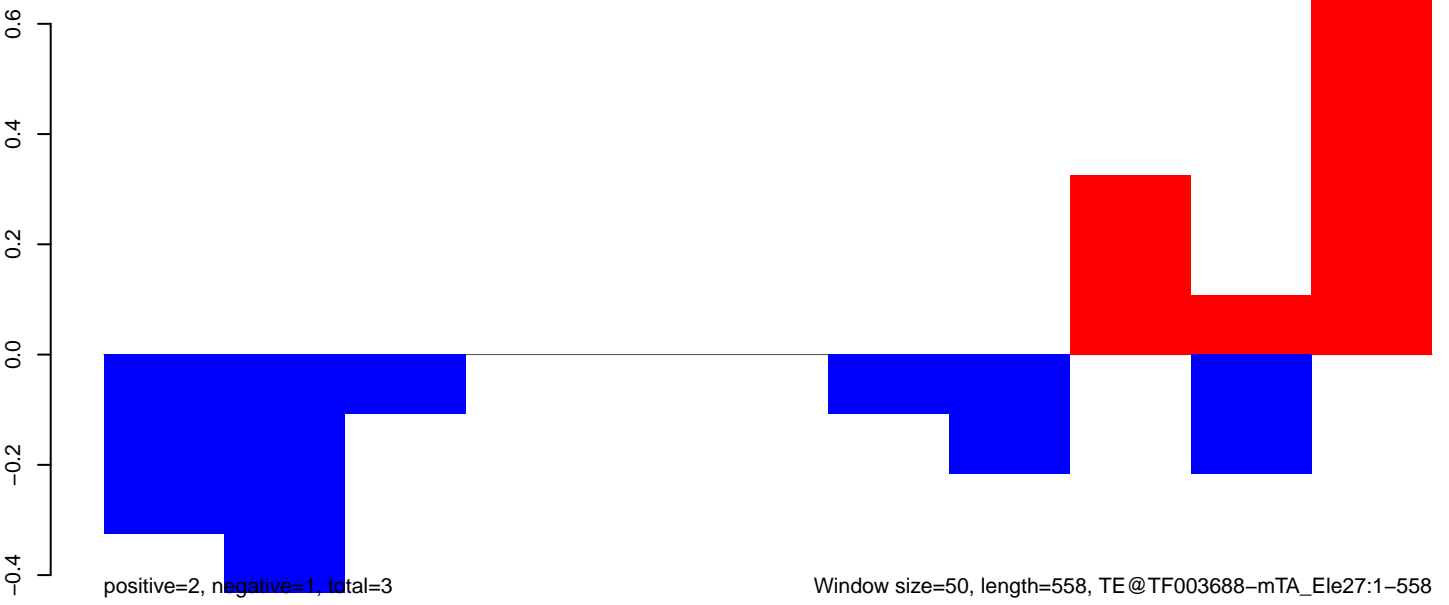
AnGam_Sua5b_cells.18_23.rep



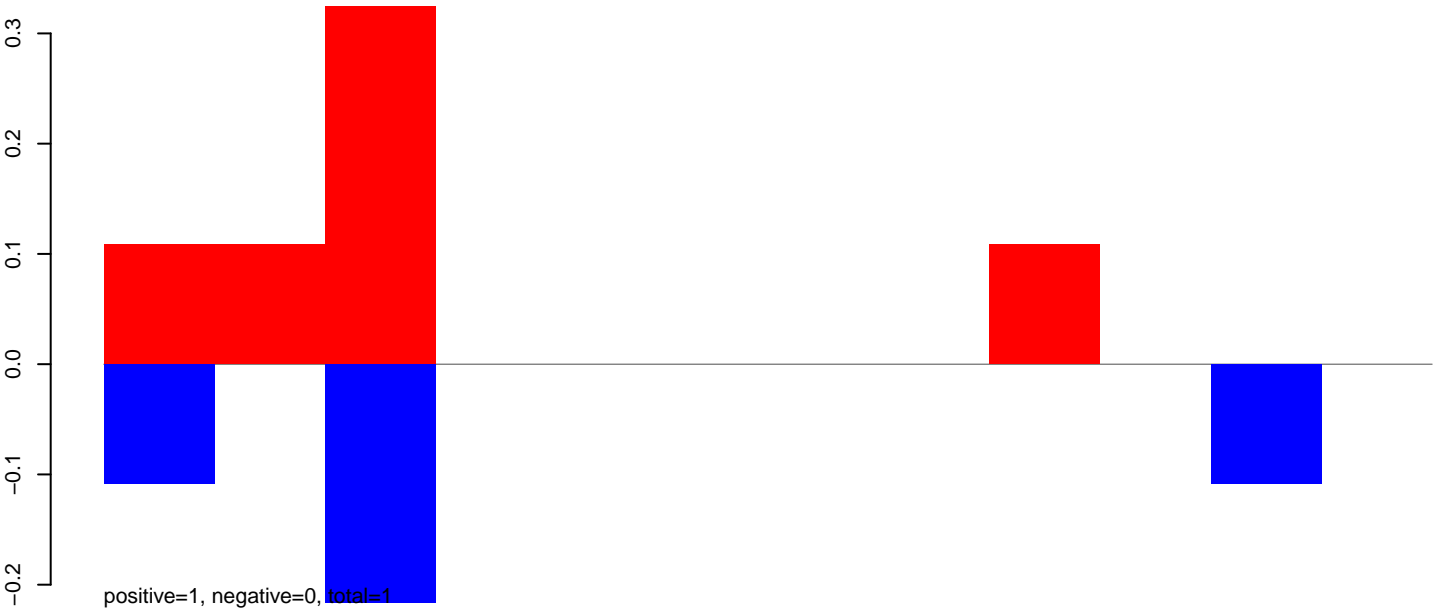
AnGam_Sua5b_cells.24_35.rep



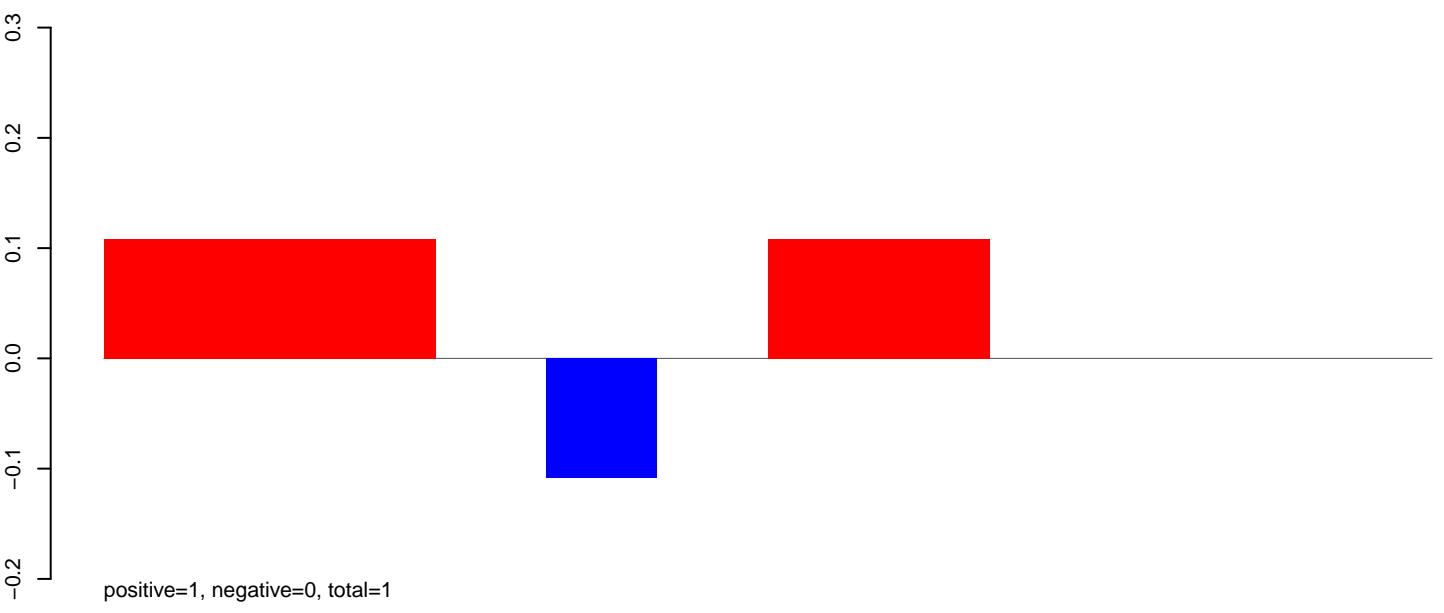
AnGam_Sua5b_cells.rep



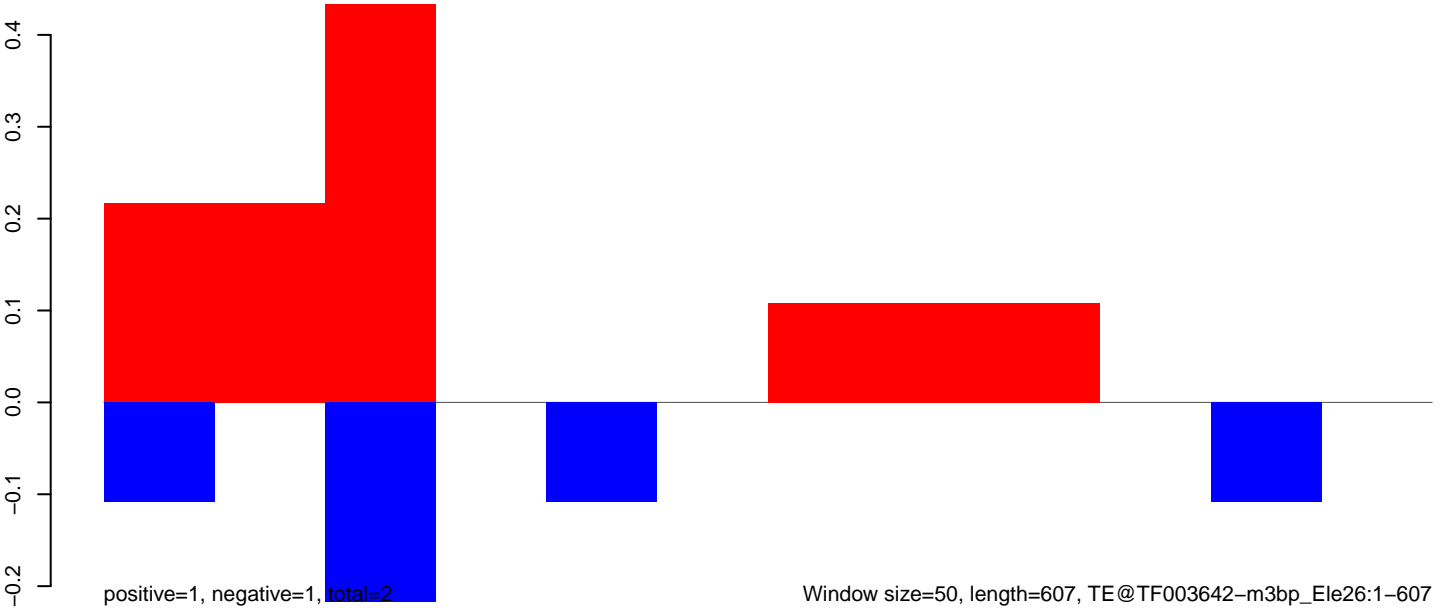
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

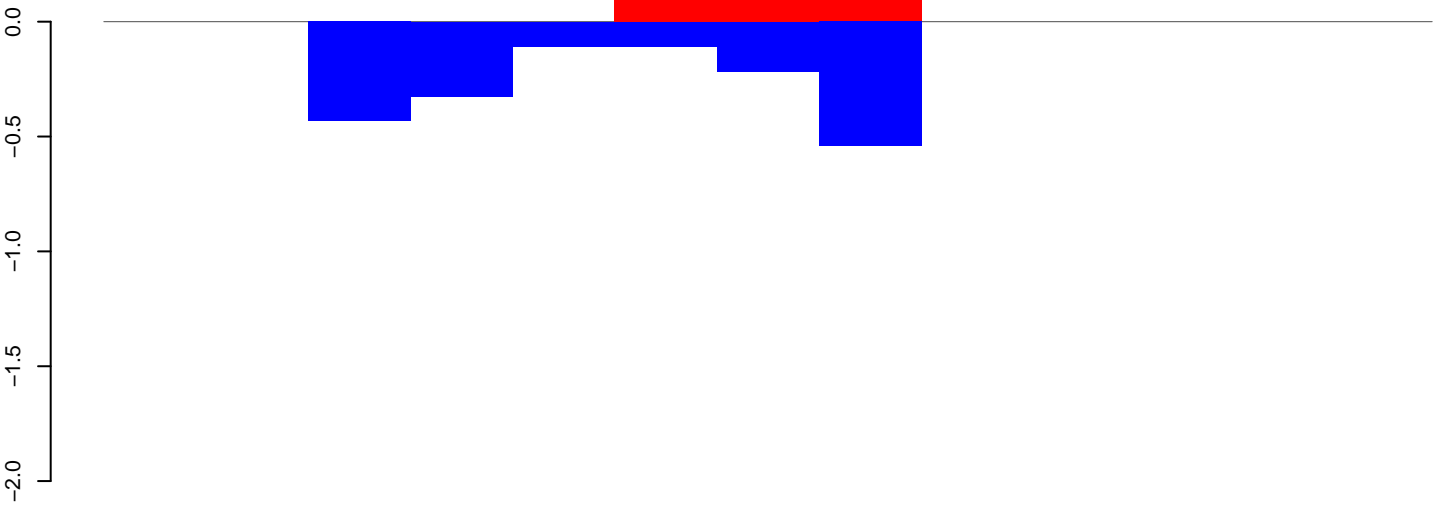


AnGam_Sua5b_cells.rep



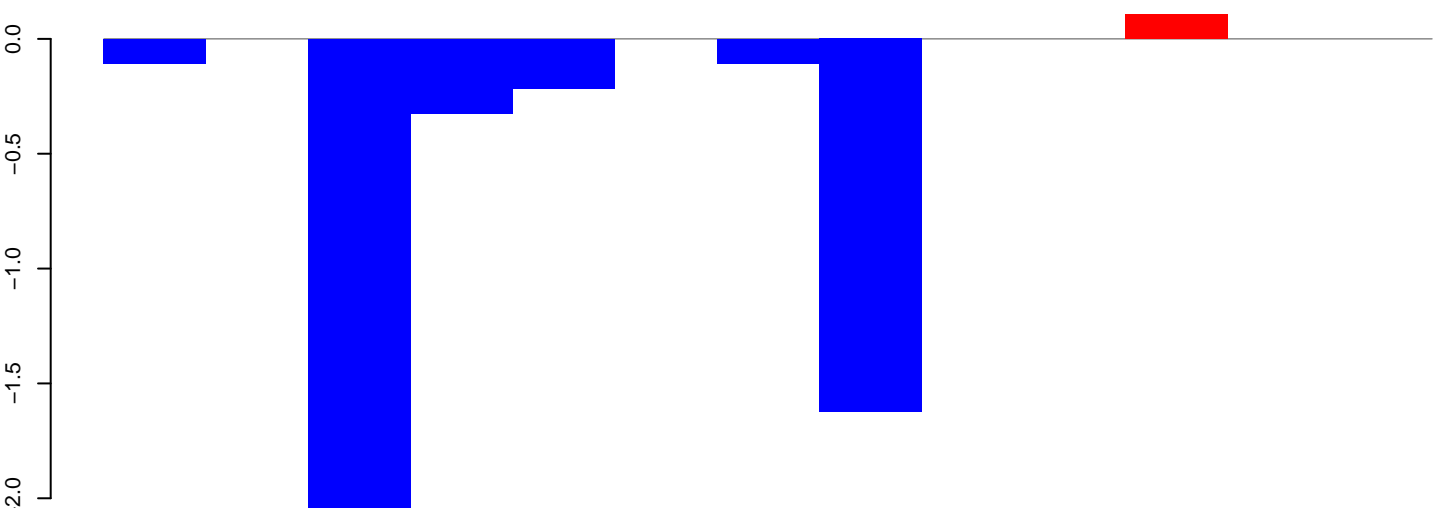
Window size=50, length=607, TE@TF003642-m3bp_Ele26:1-607

AnGam_Sua5b_cells.18_23.rep



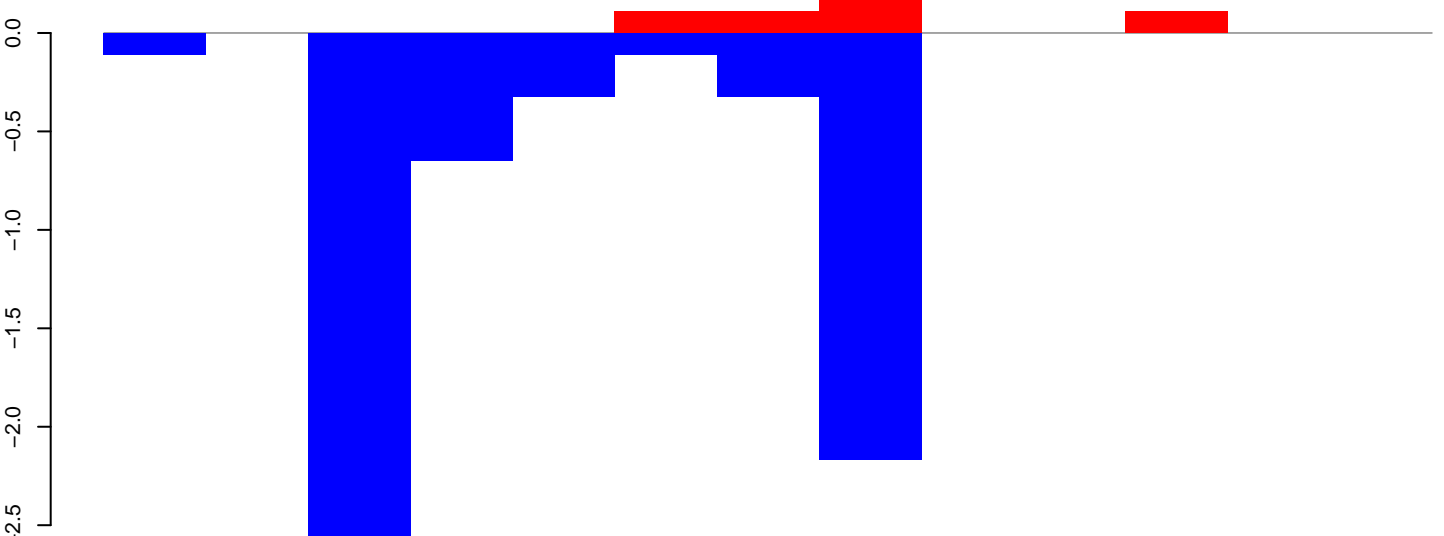
positive=0, negative=2, total=2

AnGam_Sua5b_cells.24_35.rep



positive=0, negative=5, total=5

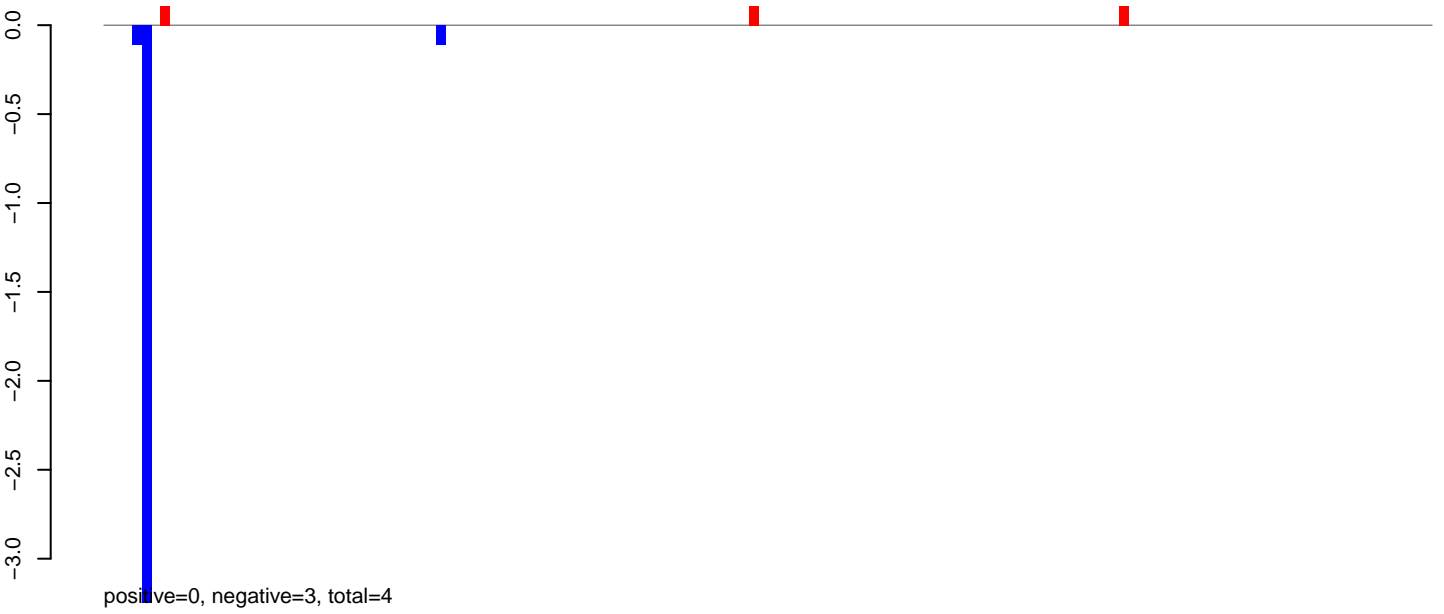
AnGam_Sua5b_cells.rep



positive=1, negative=6, total=7

Window size=50, length=678, TE@TF003625-m2bp_Ele8:1-678

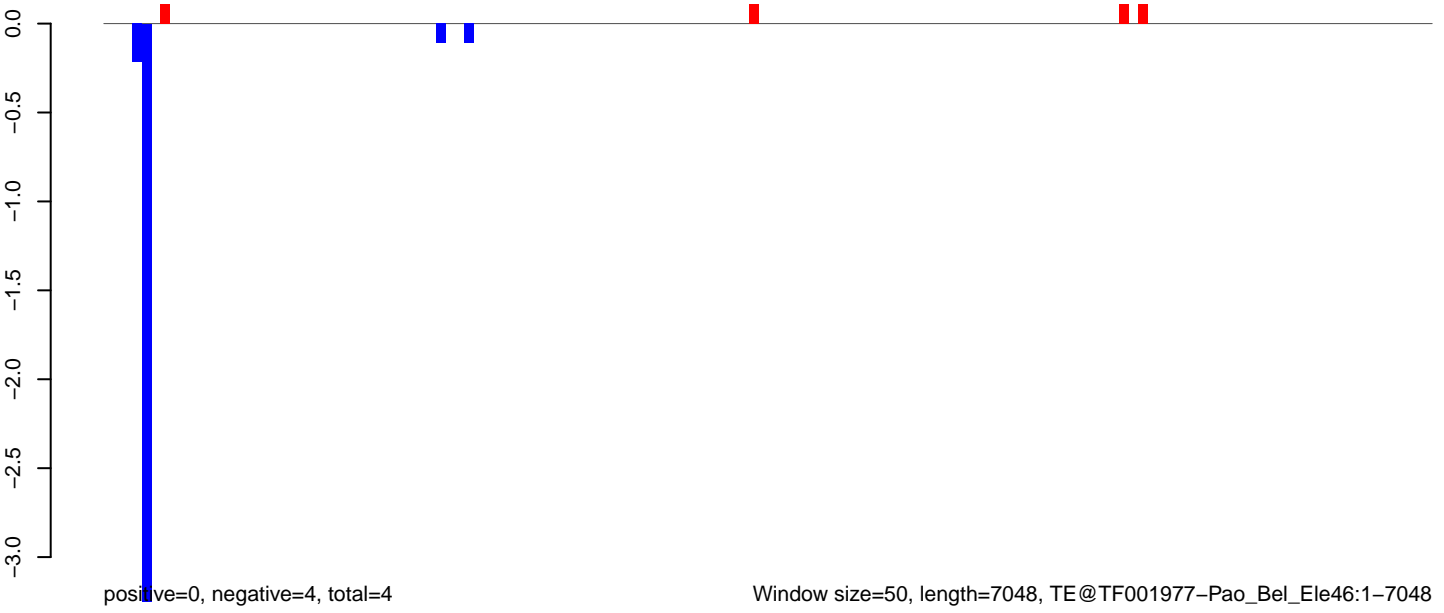
AnGam_Sua5b_cells.18_23.rep



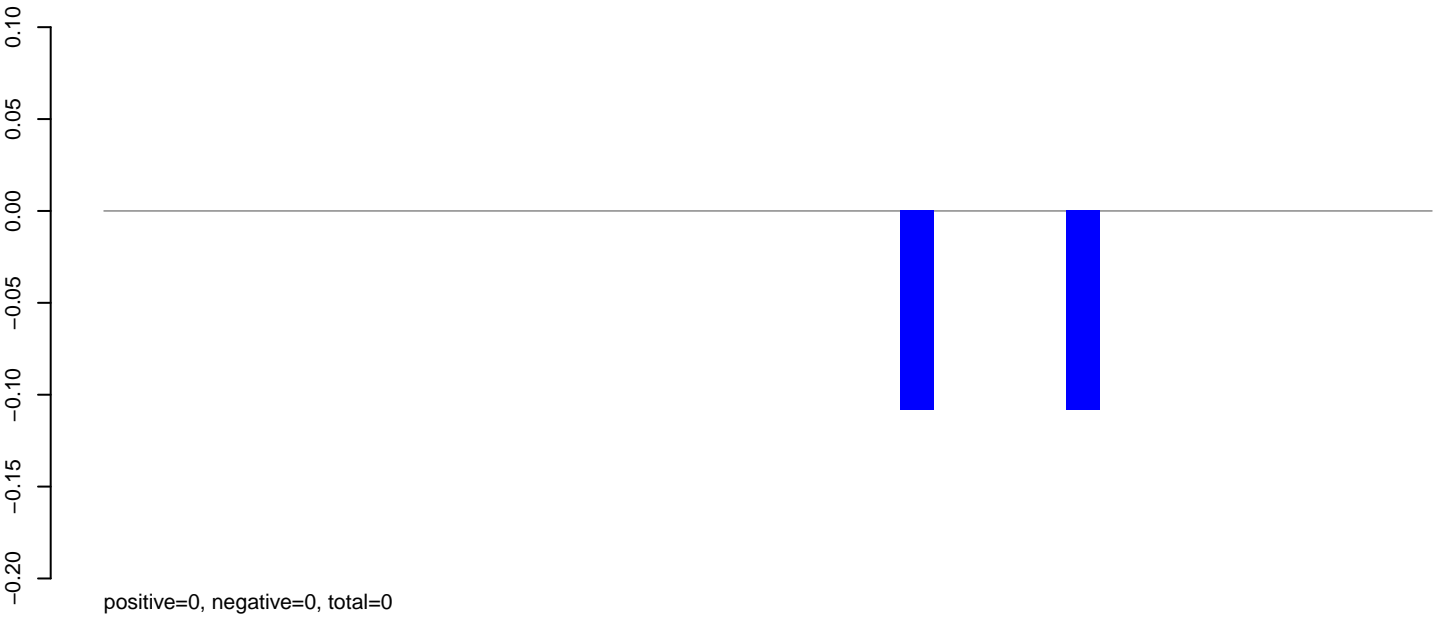
AnGam_Sua5b_cells.24_35.rep



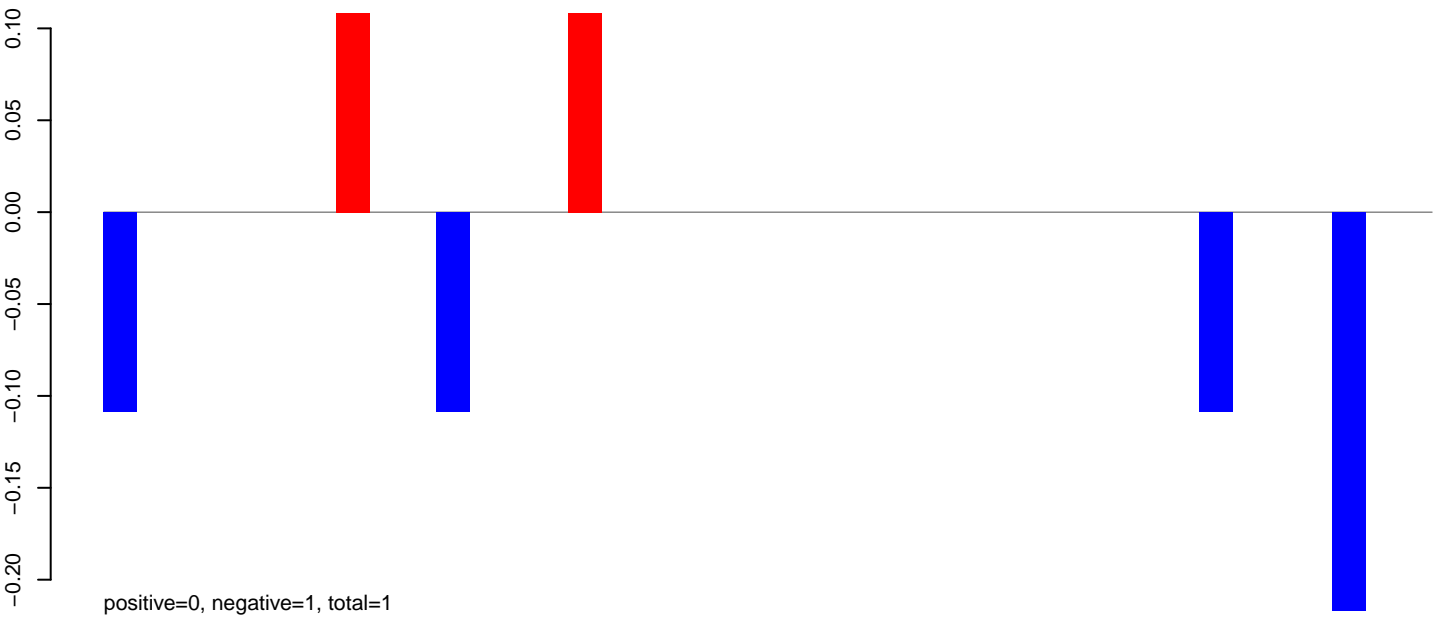
AnGam_Sua5b_cells.rep



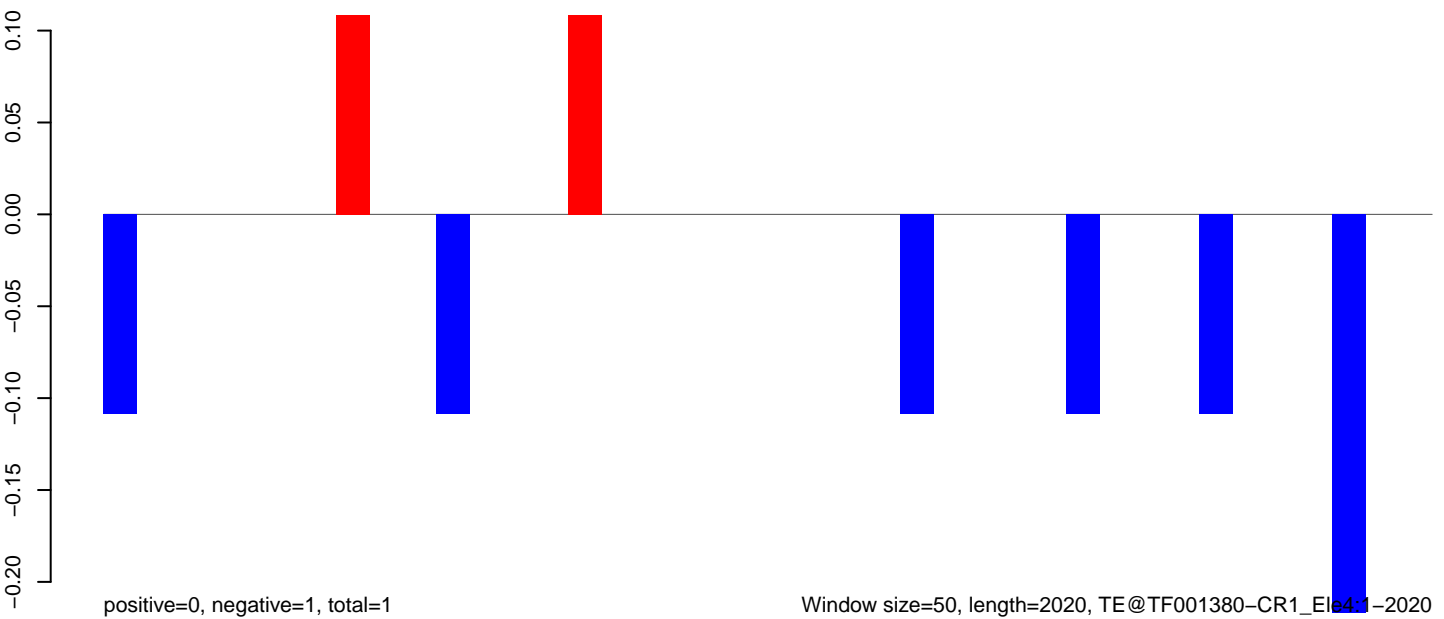
AnGam_Sua5b_cells.18_23.rep



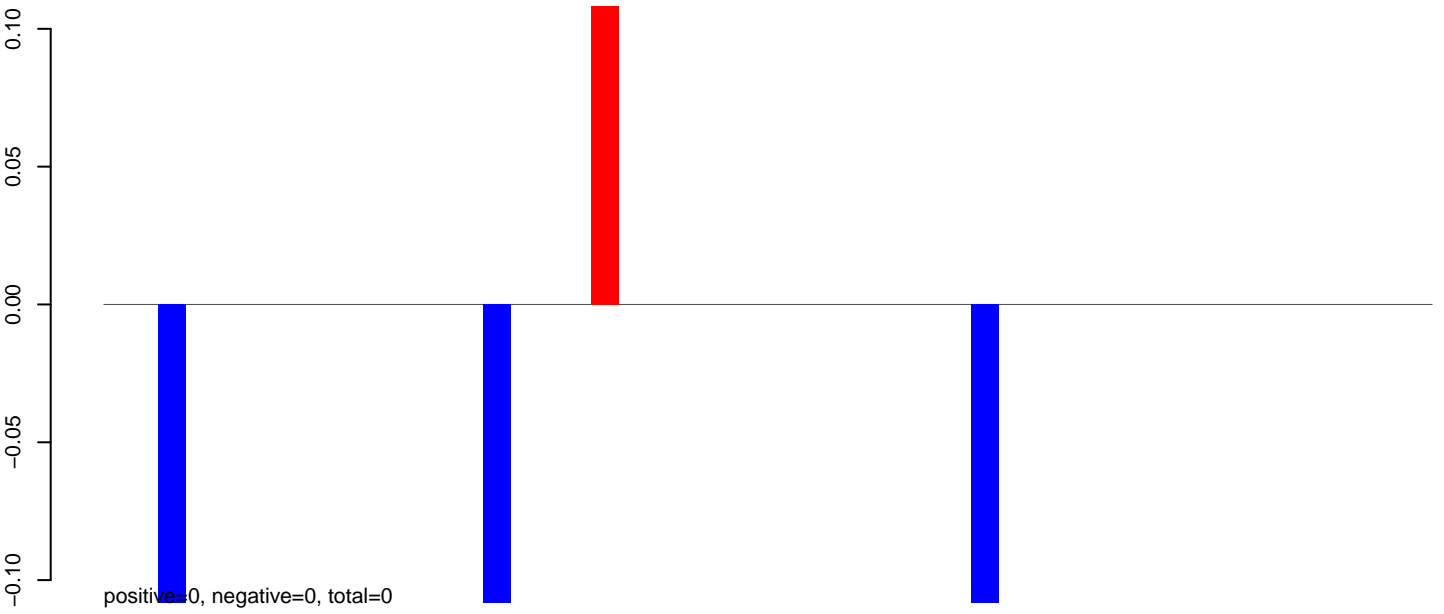
AnGam_Sua5b_cells.24_35.rep



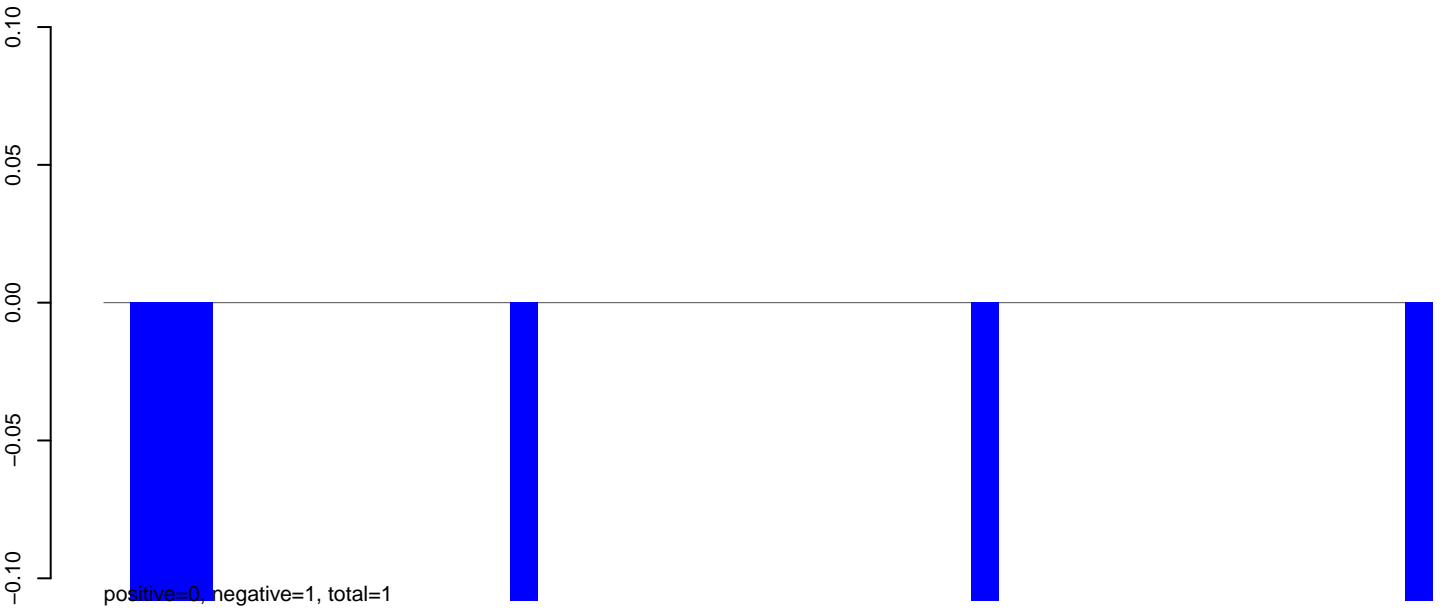
AnGam_Sua5b_cells.rep



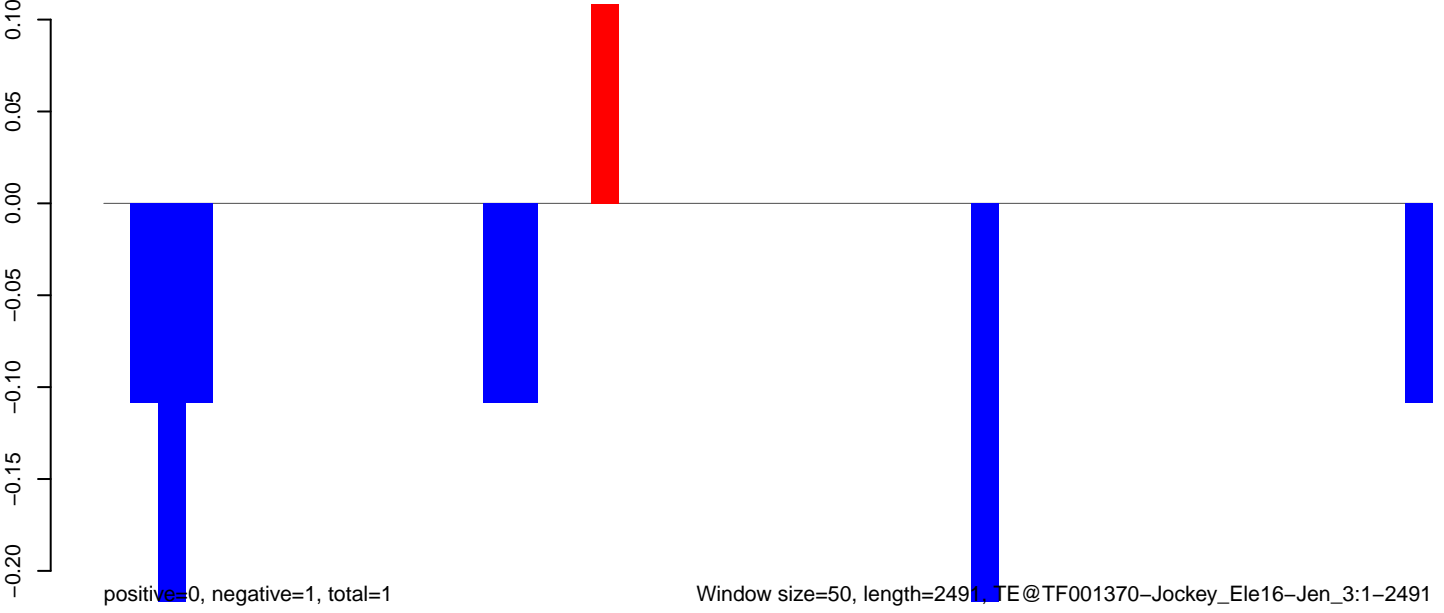
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

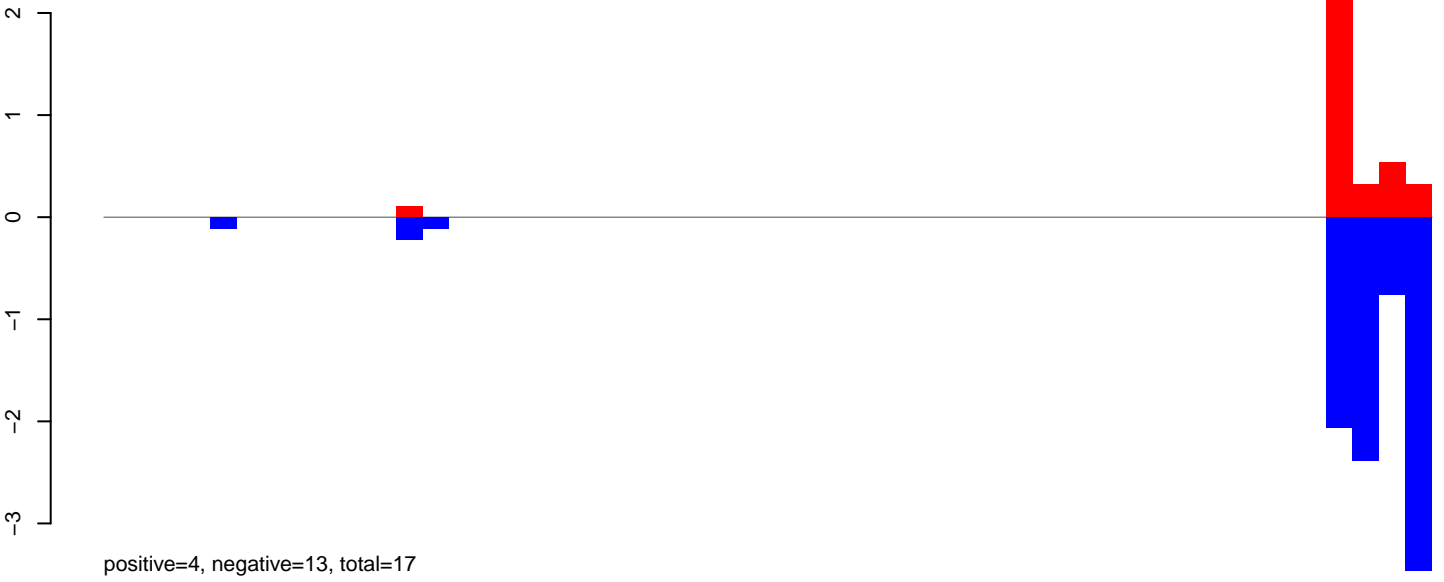


AnGam_Sua5b_cells.rep

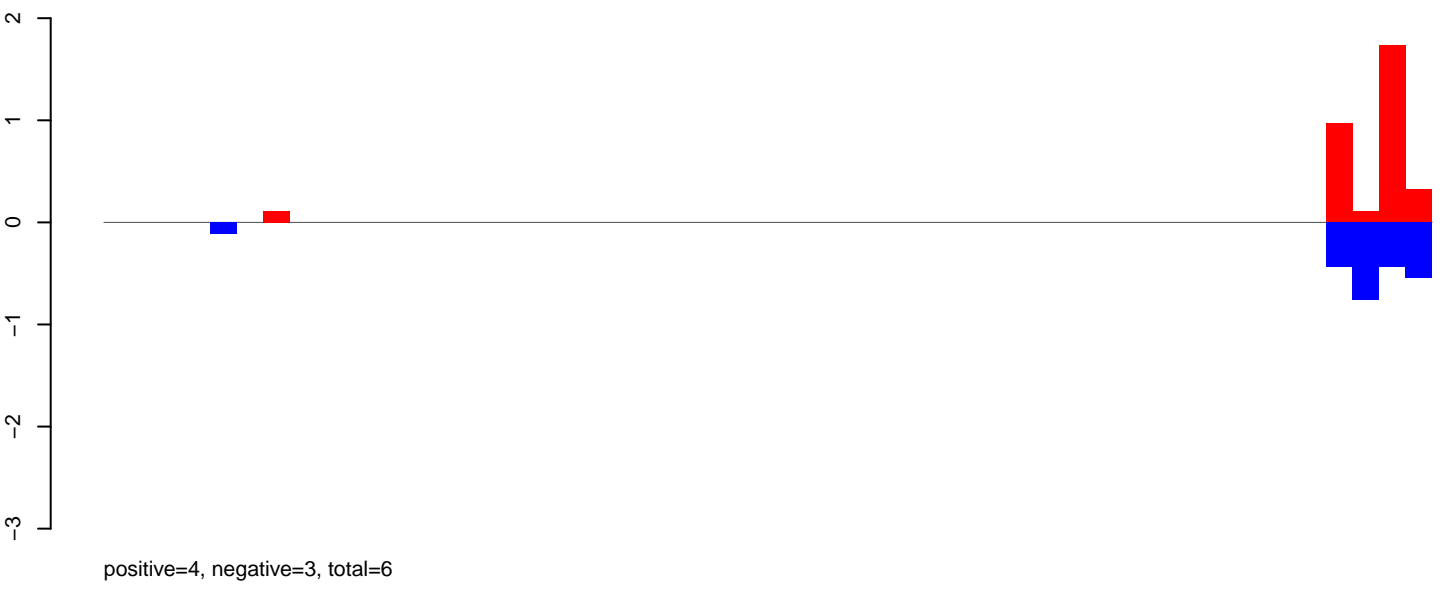


0 500 1000 1500 2000 2500

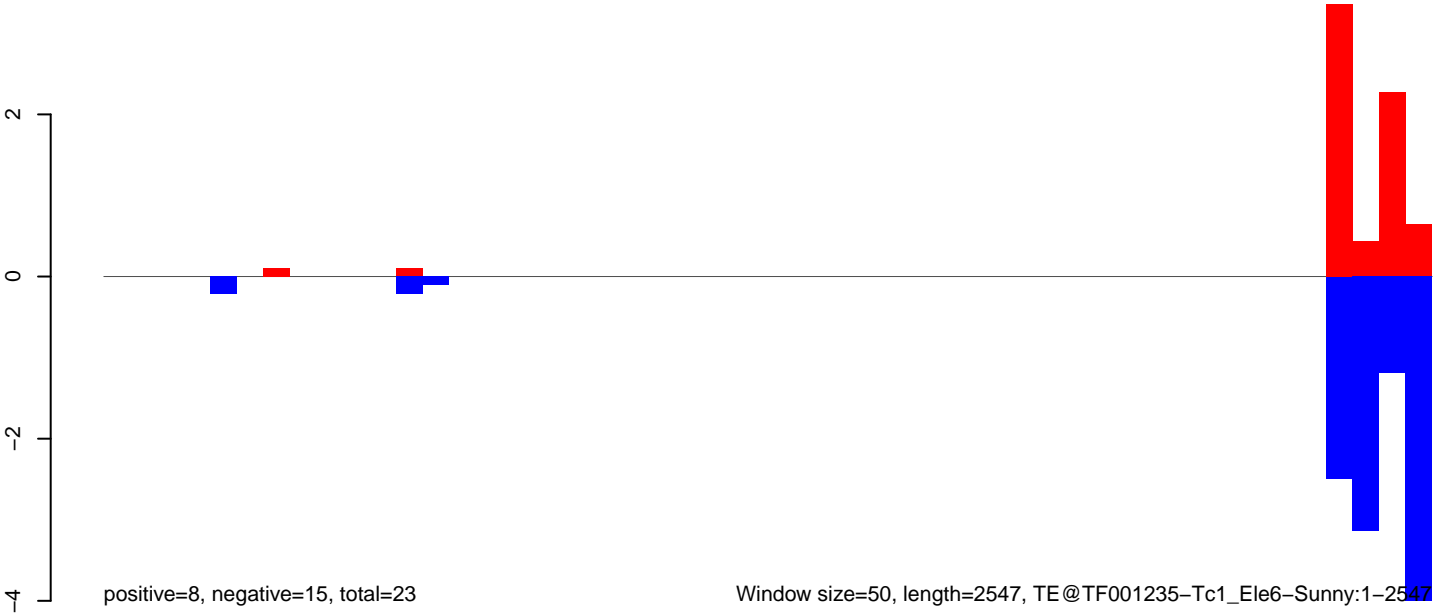
AnGam_Sua5b_cells.18_23.rep



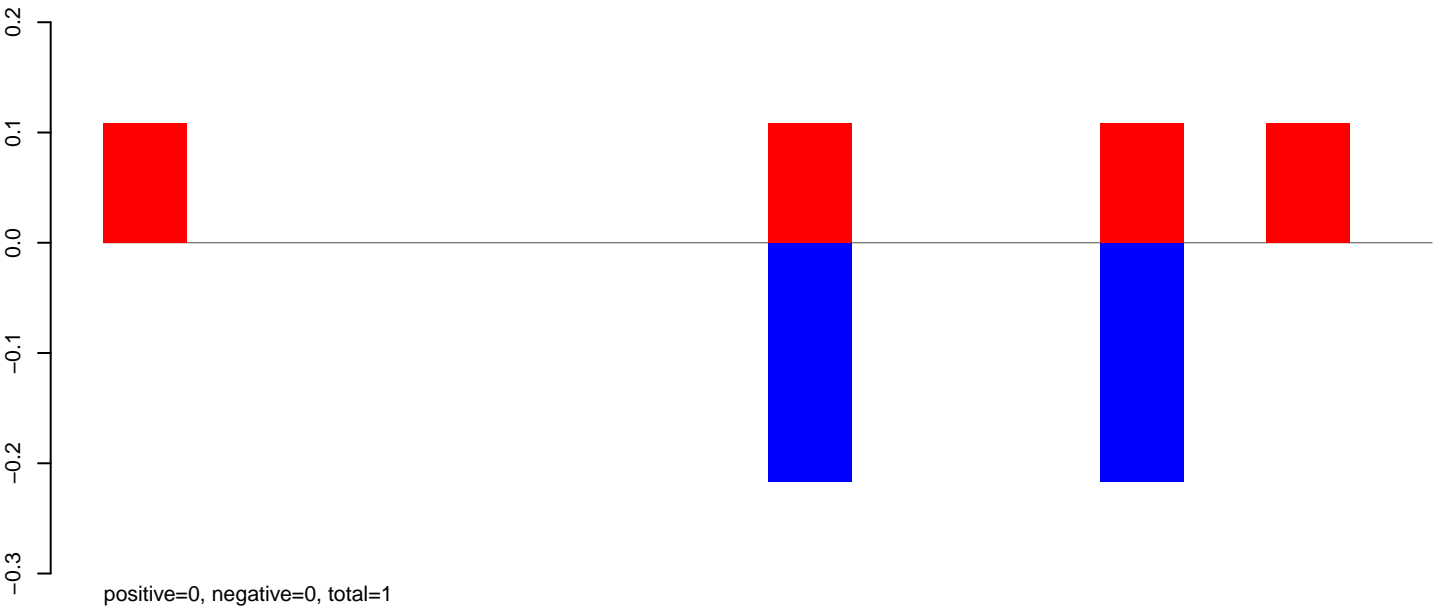
AnGam_Sua5b_cells.24_35.rep



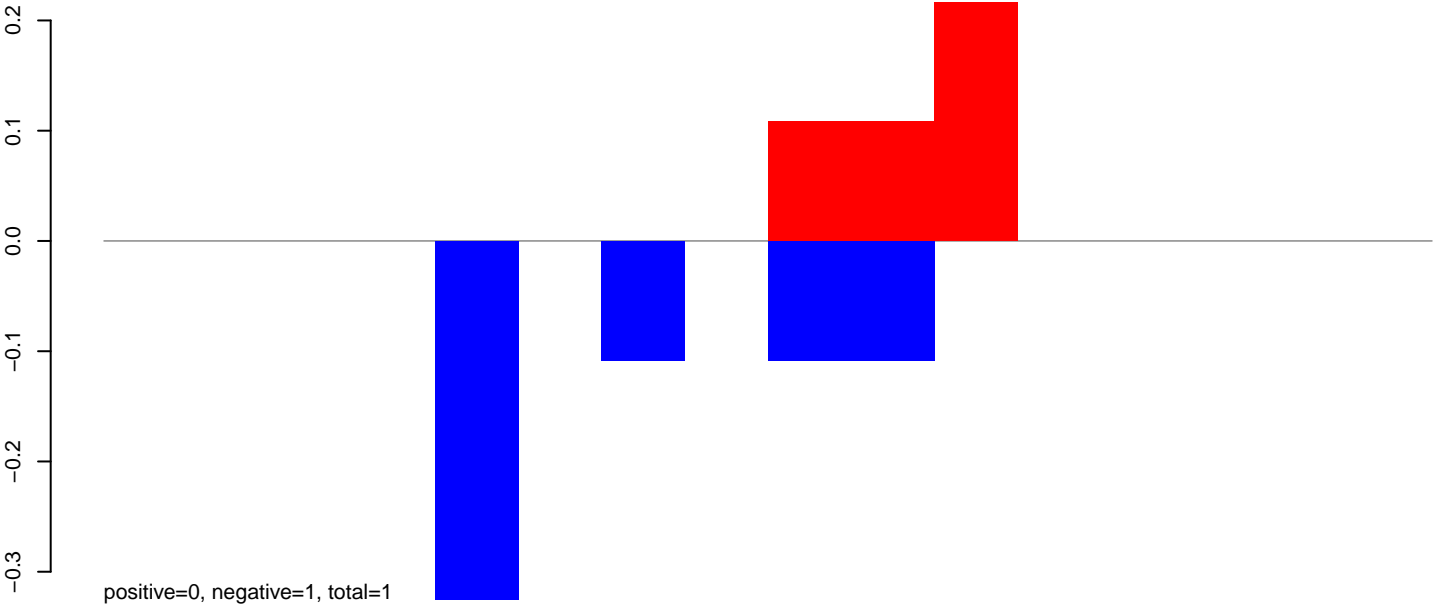
AnGam_Sua5b_cells.rep



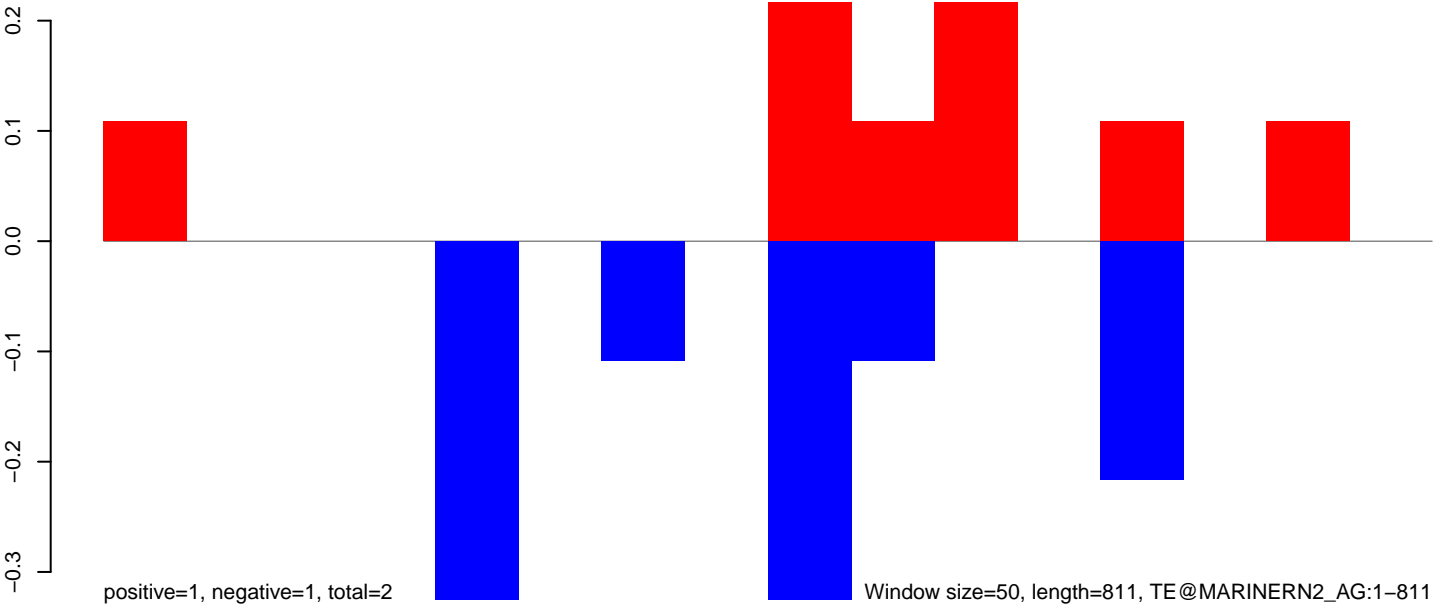
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

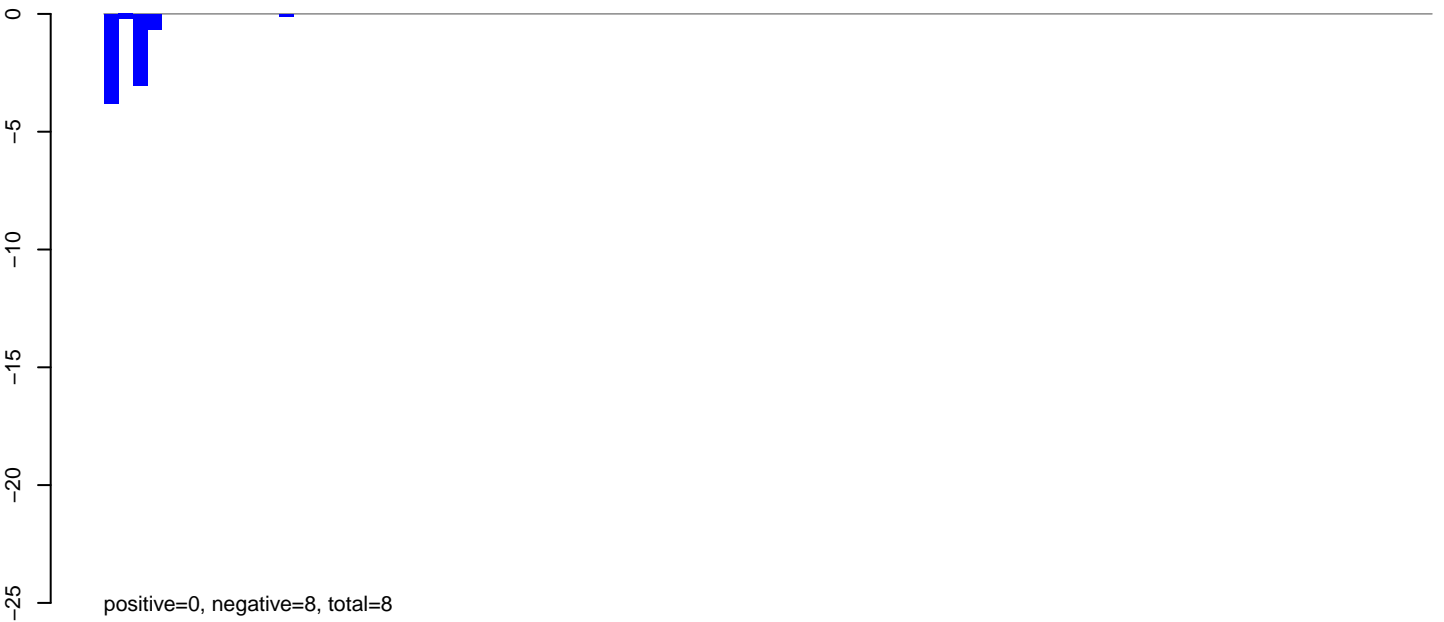


AnGam_Sua5b_cells.rep



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

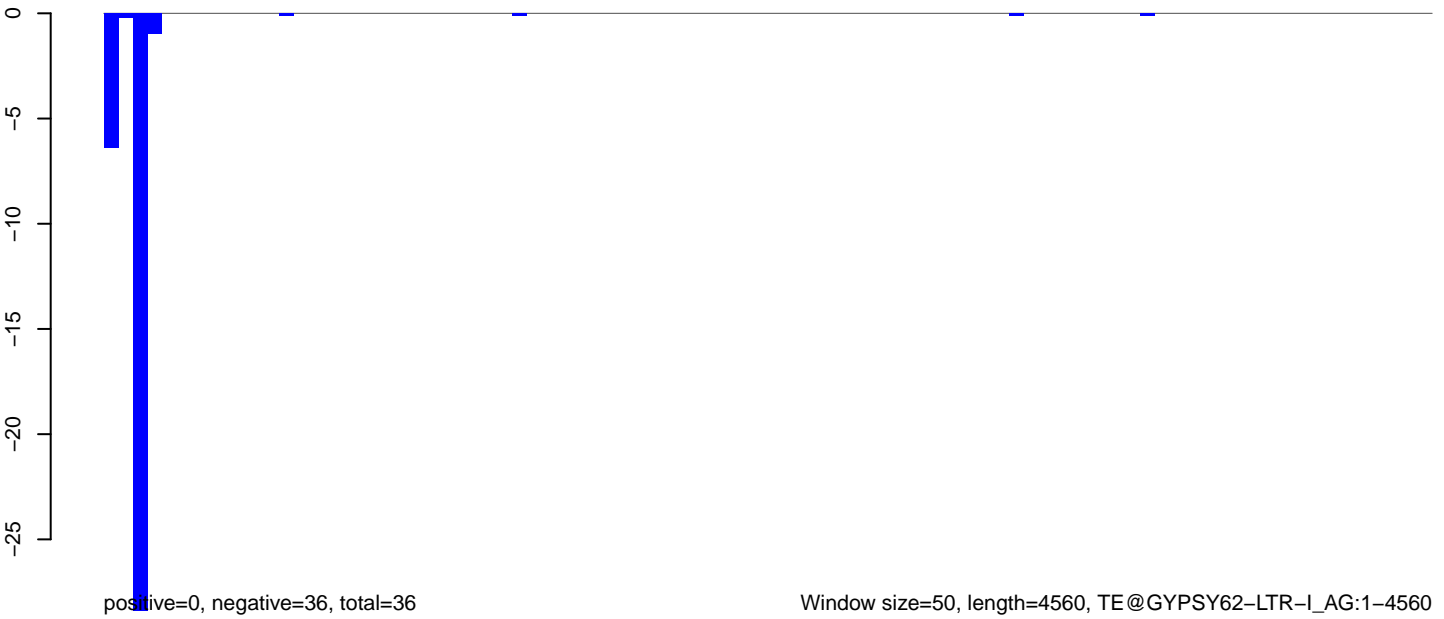
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

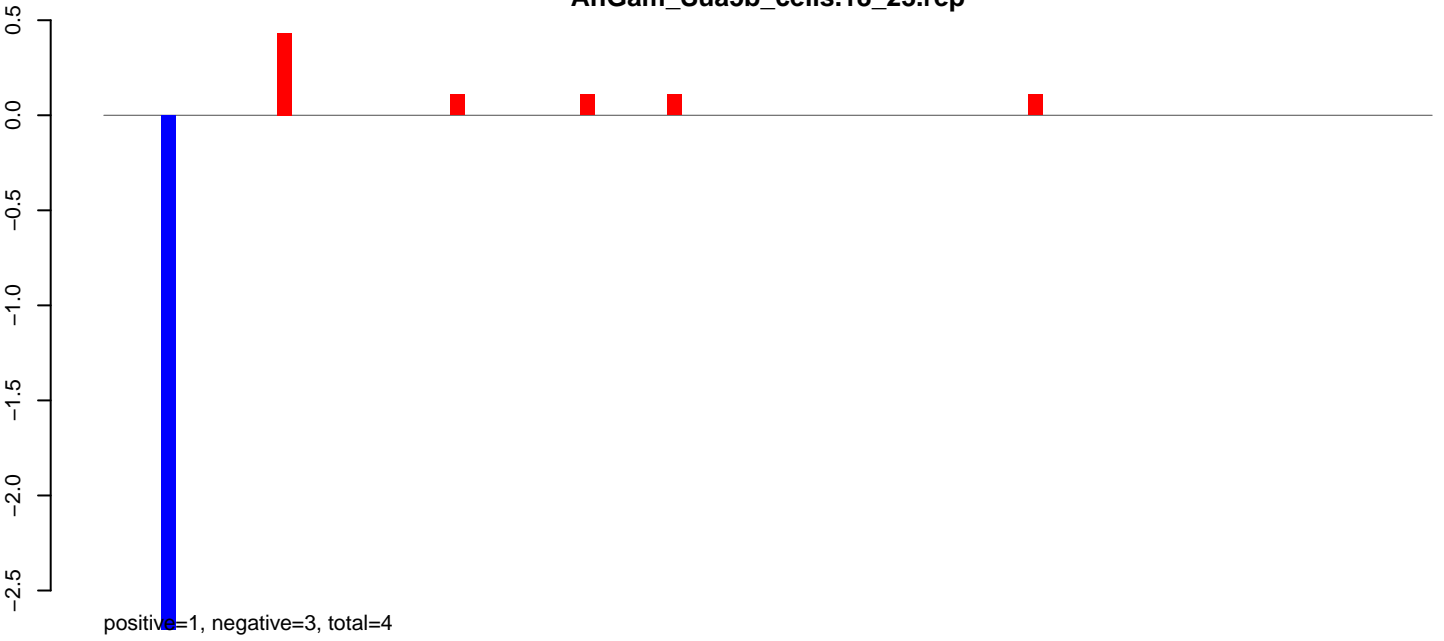


AnGam_Sua5b_cells.rep

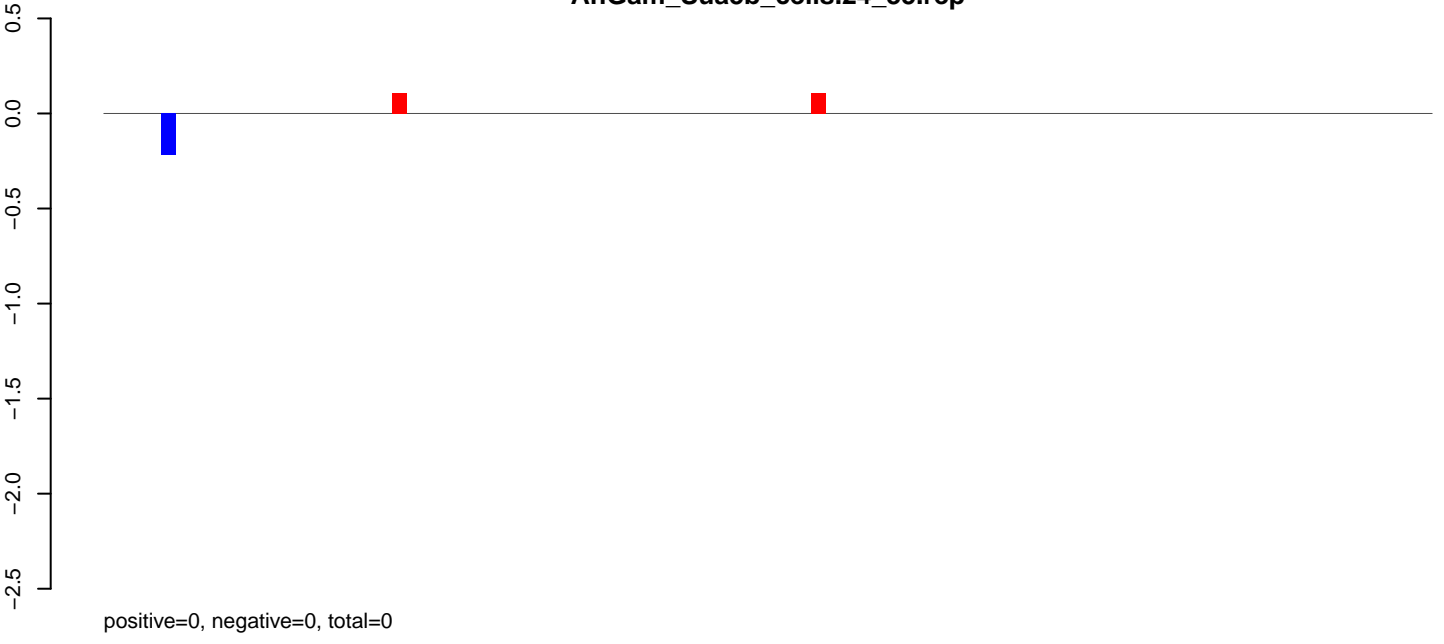


Window size=50, length=4560, TE@GYPSY62-LTR-I_AG:1-4560

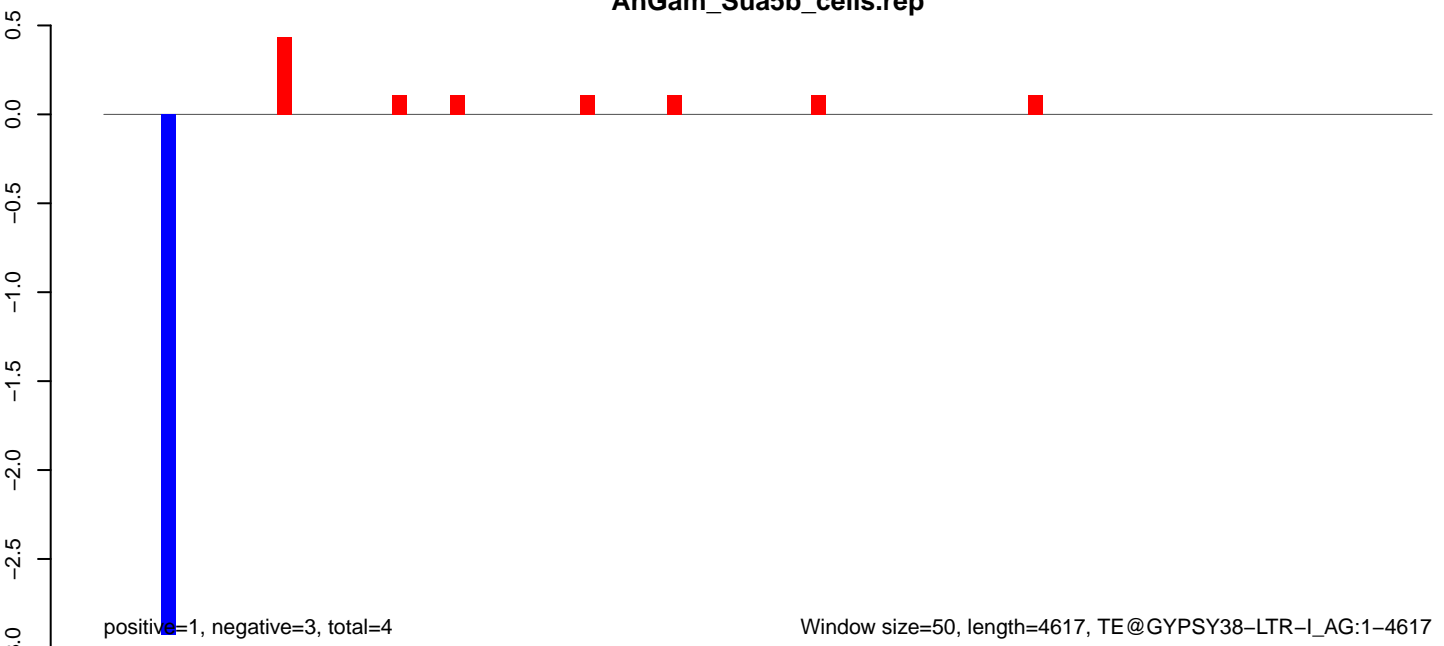
AnGam_Sua5b_cells.18_23.rep



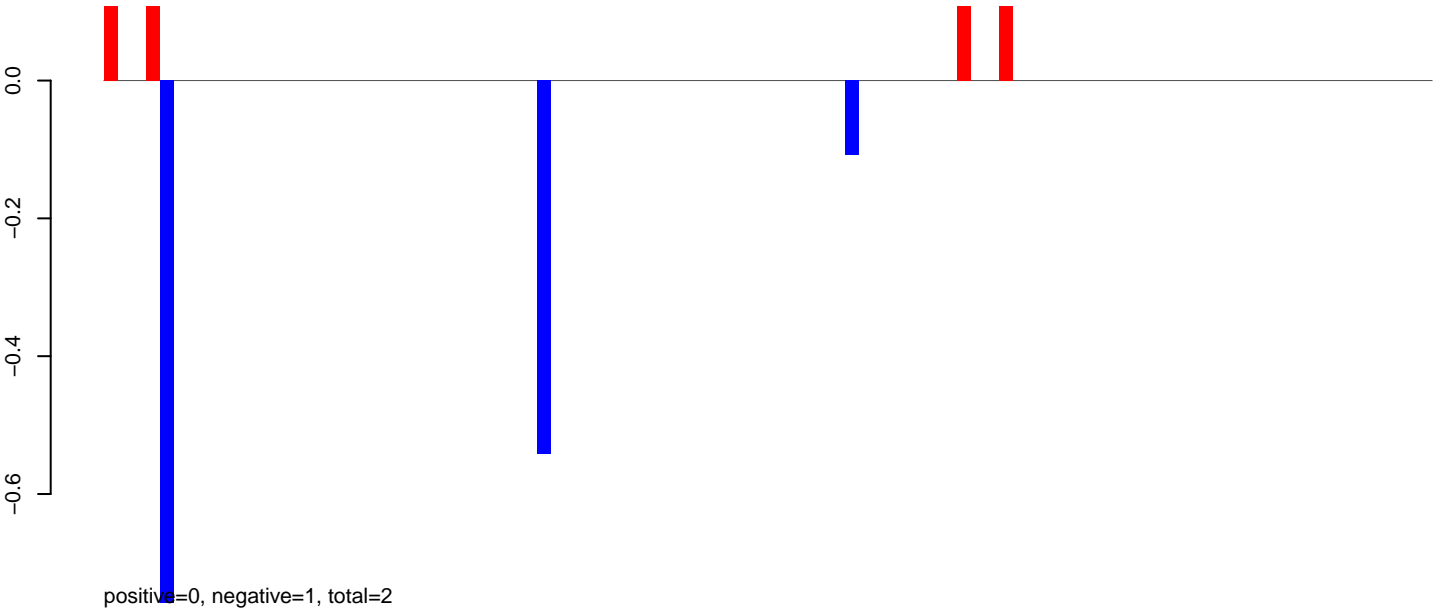
AnGam_Sua5b_cells.24_35.rep



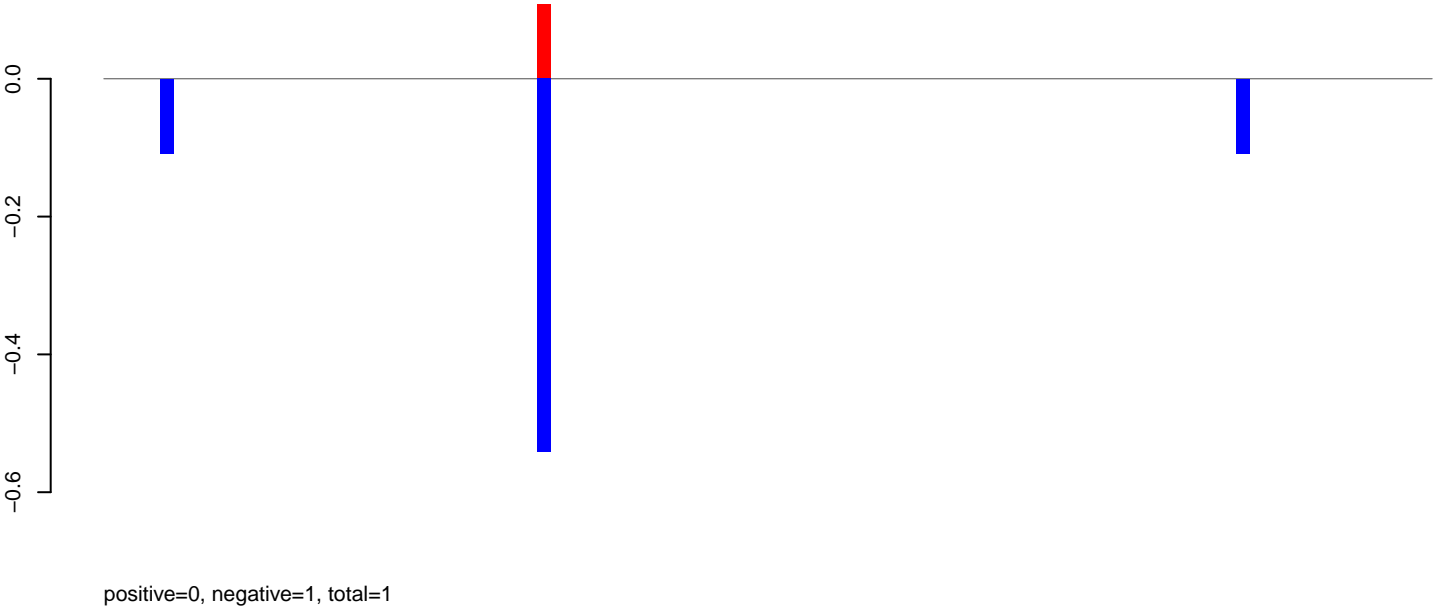
AnGam_Sua5b_cells.rep



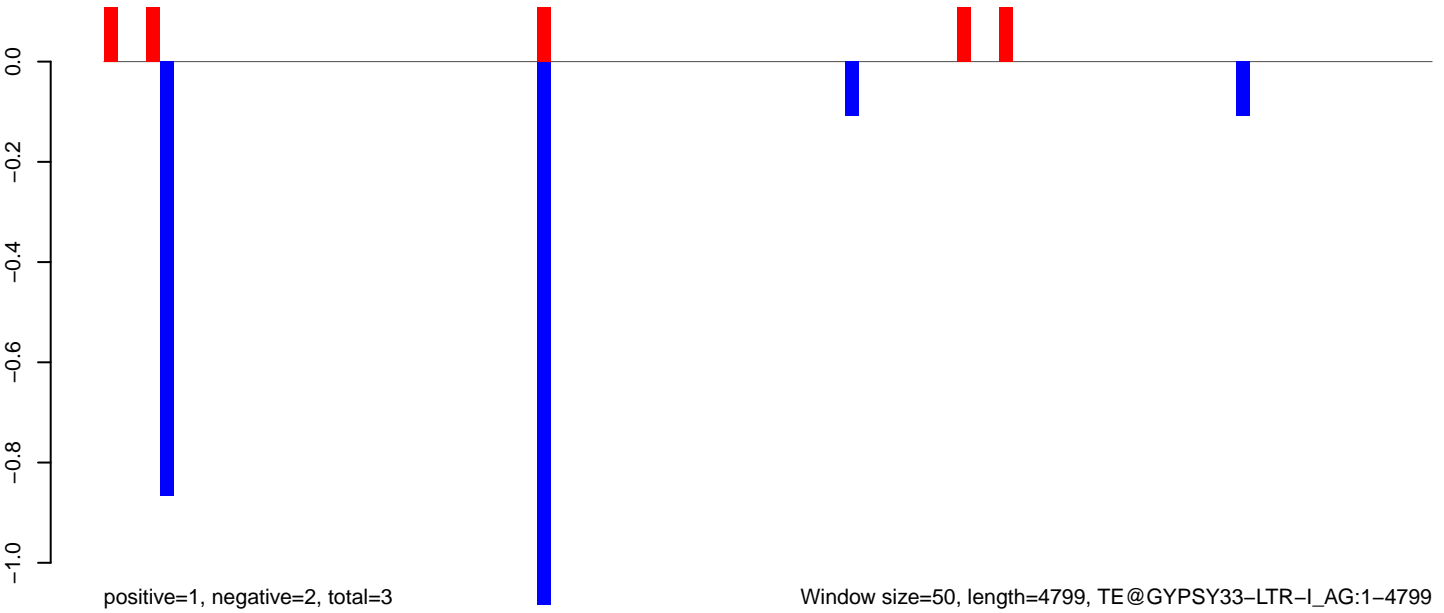
AnGam_Sua5b_cells.18_23.rep



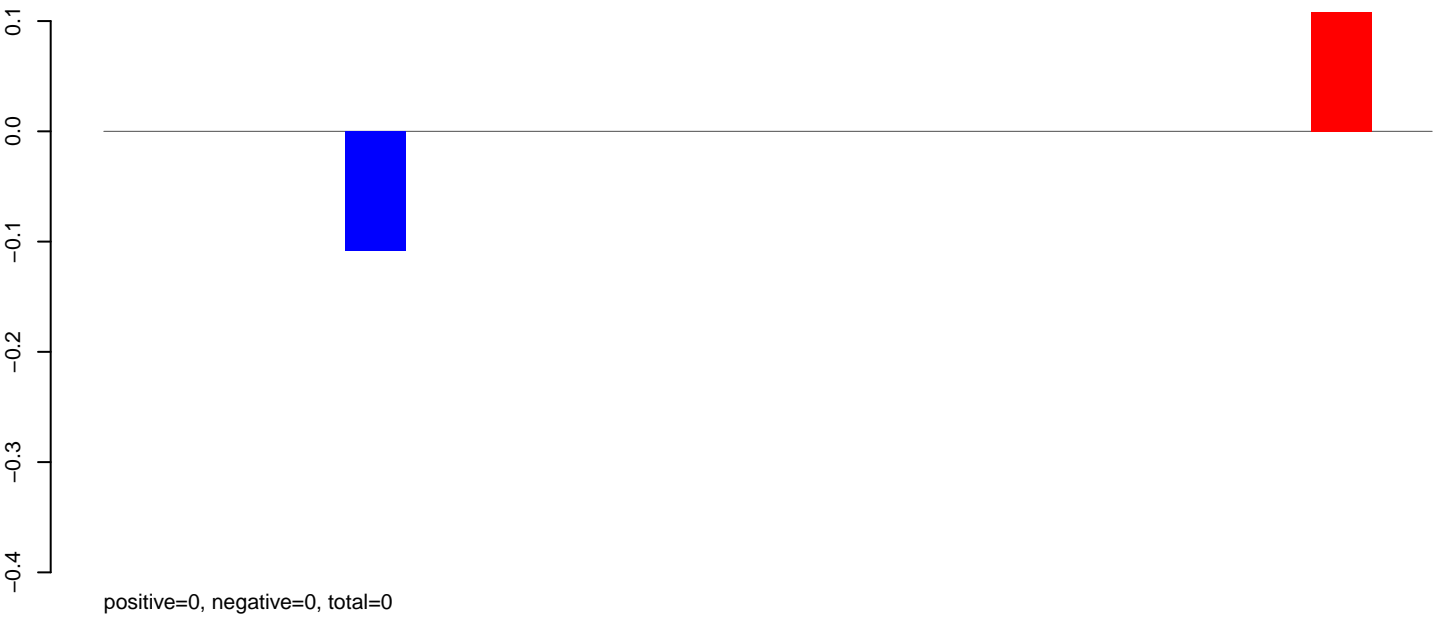
AnGam_Sua5b_cells.24_35.rep



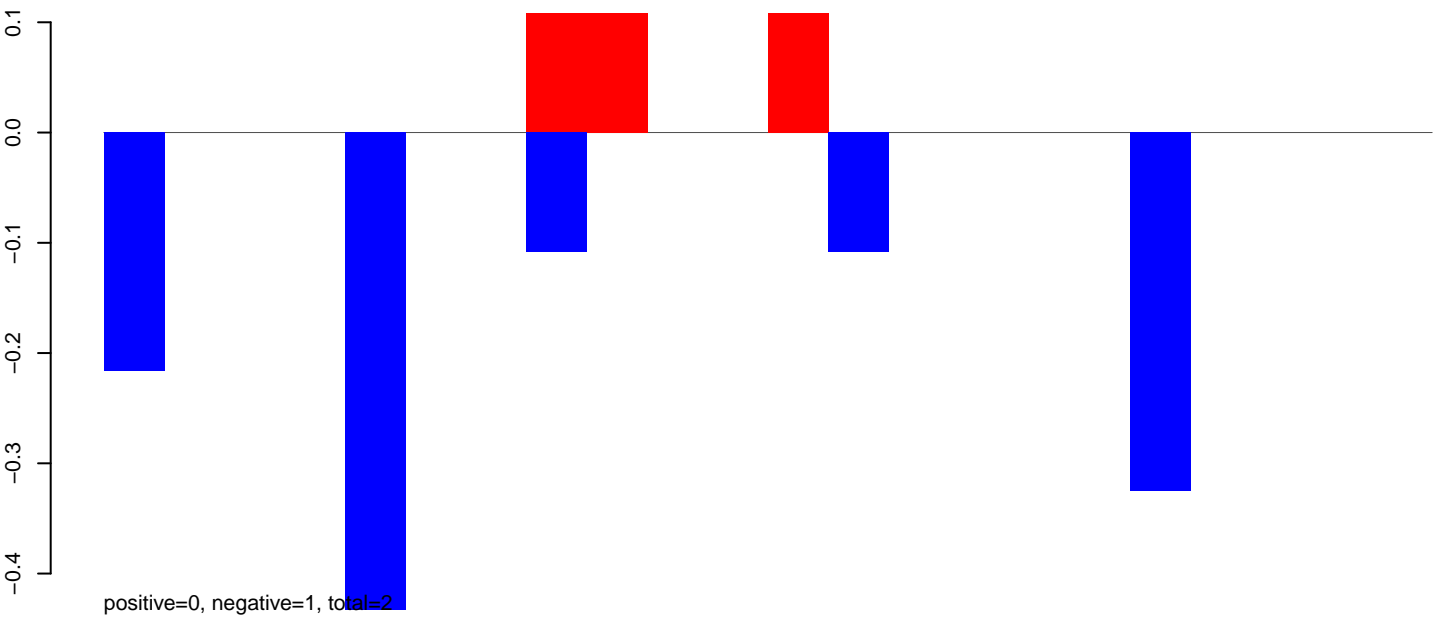
AnGam_Sua5b_cells.rep



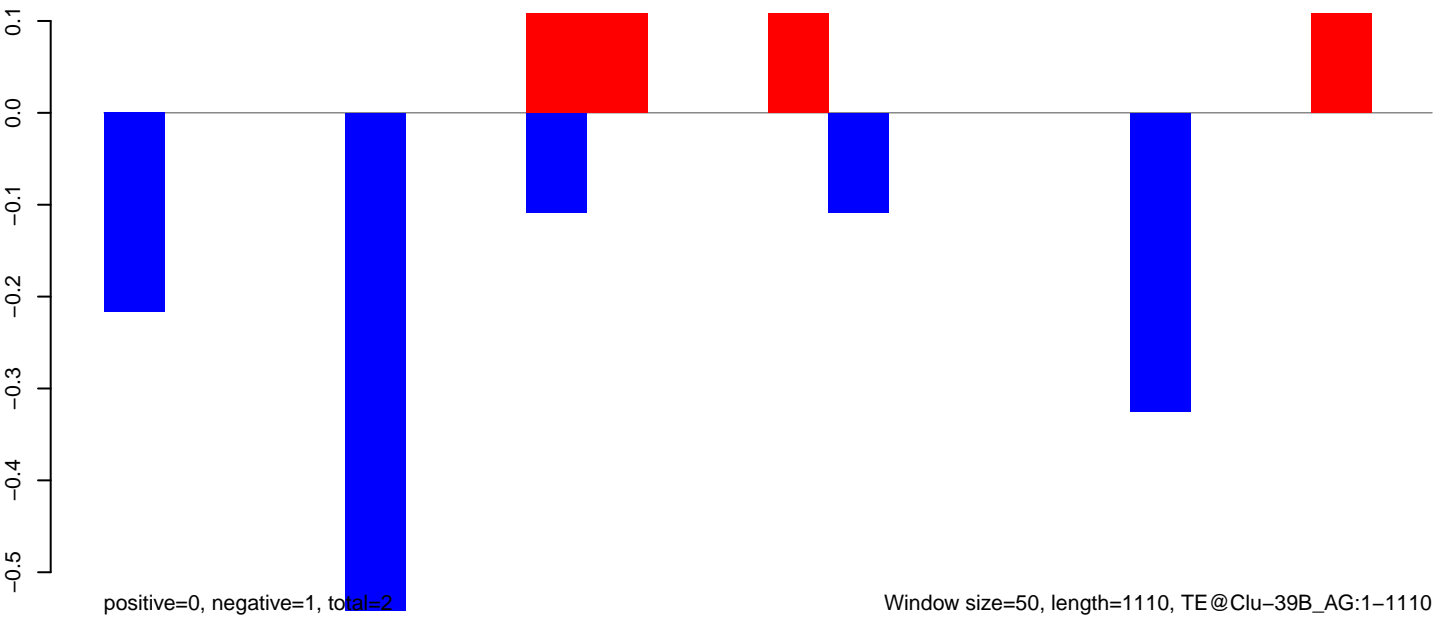
AnGam_Sua5b_cells.18_23.rep



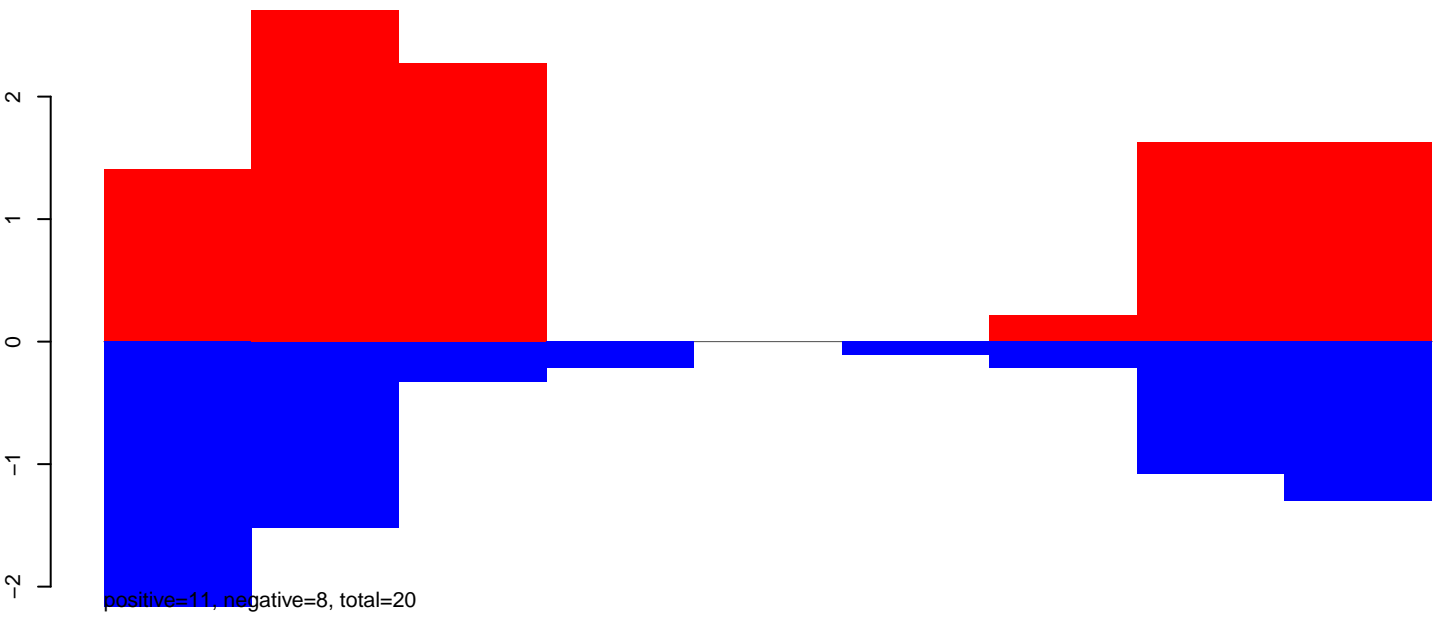
AnGam_Sua5b_cells.24_35.rep



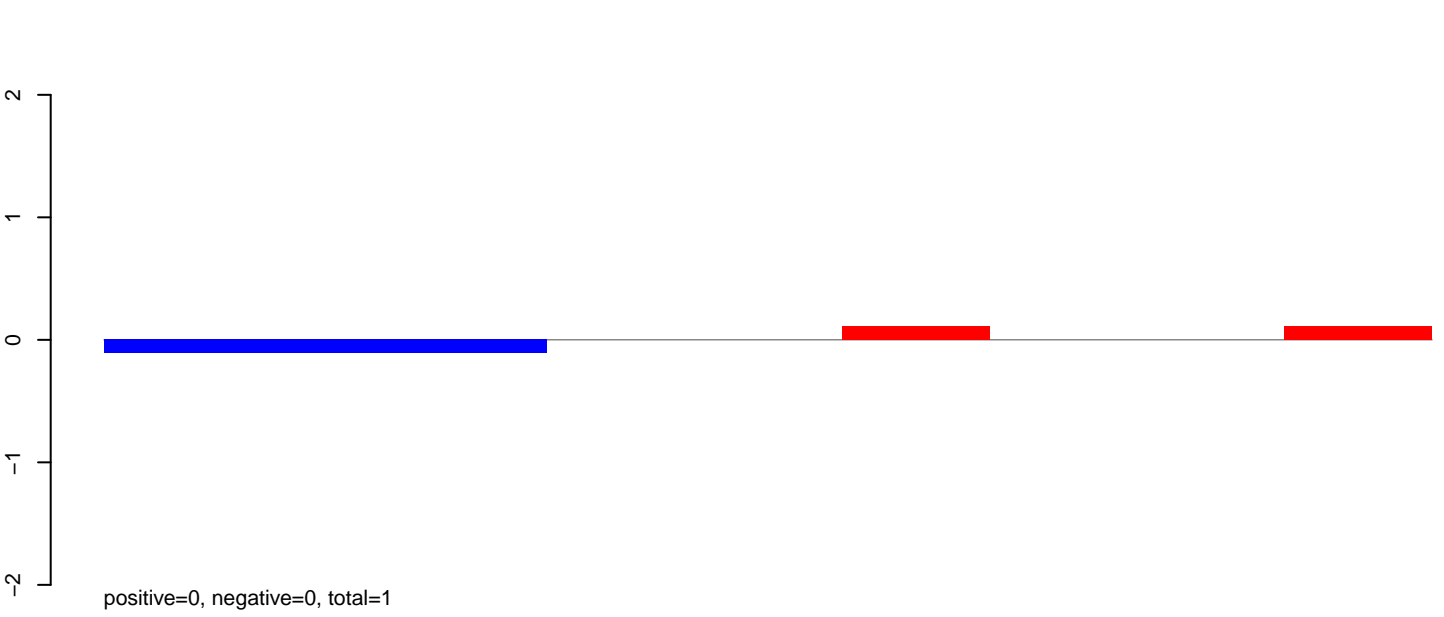
AnGam_Sua5b_cells.rep



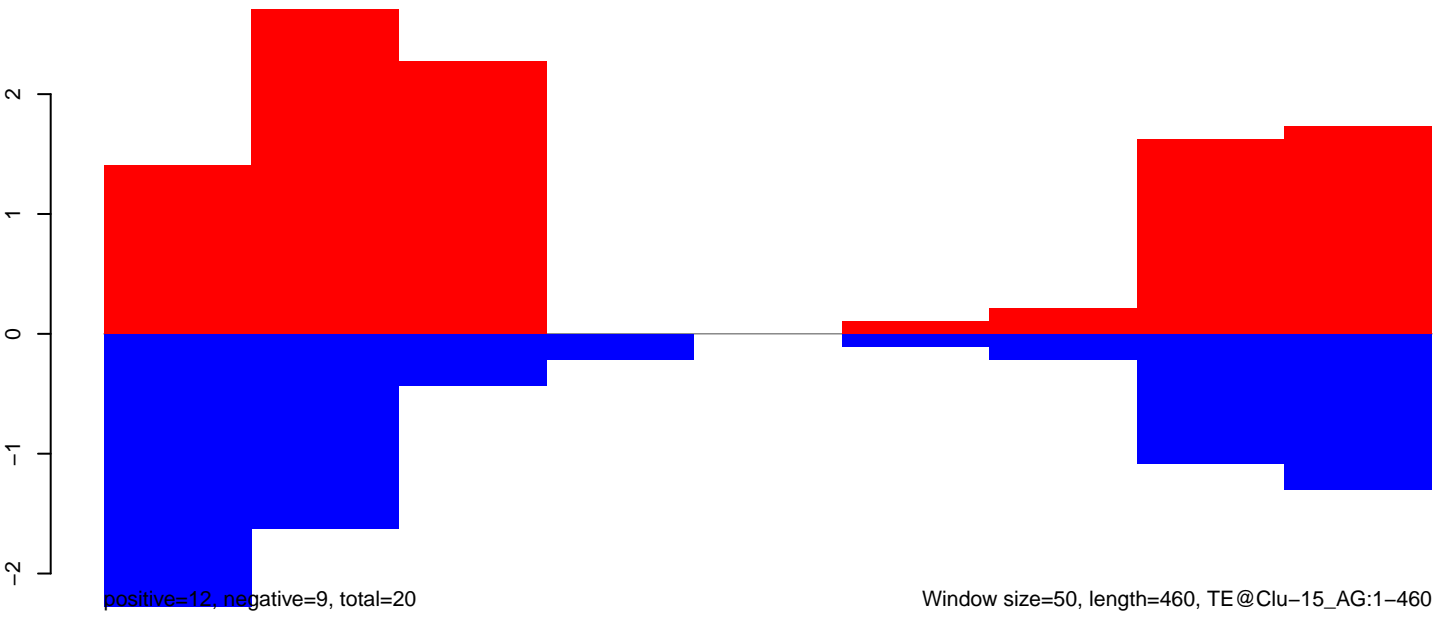
AnGam_Sua5b_cells.18_23.rep



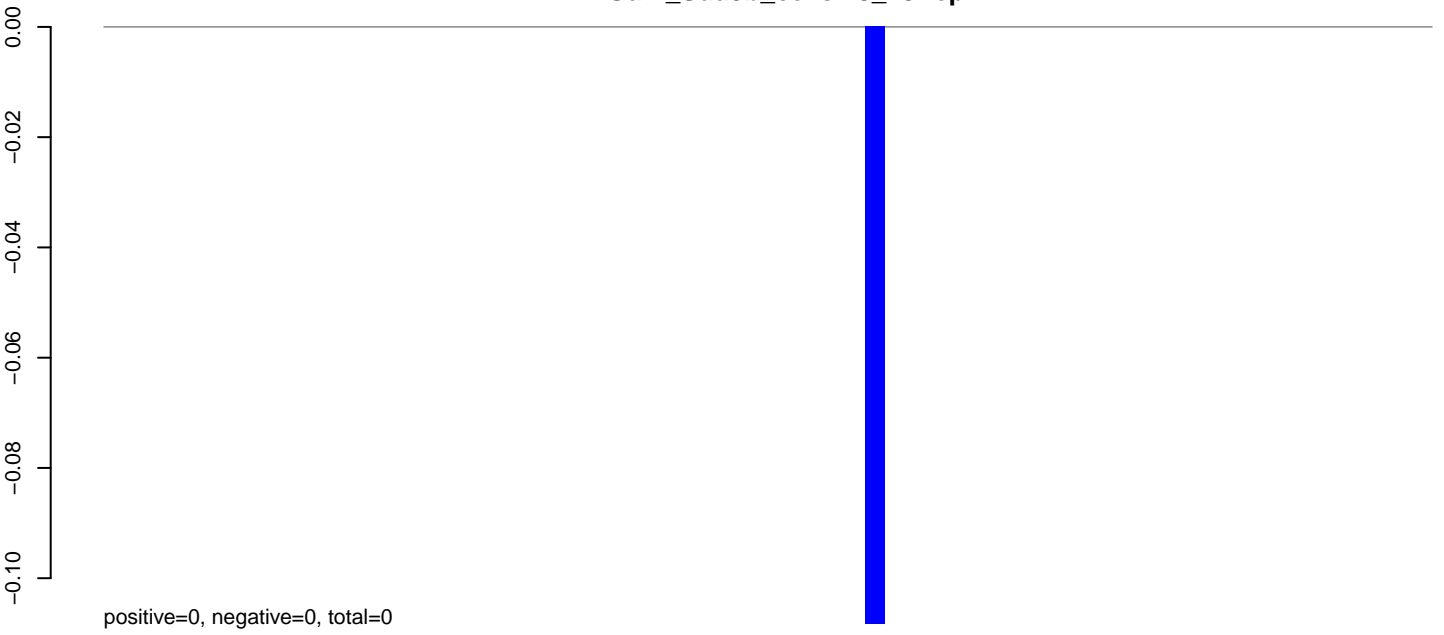
AnGam_Sua5b_cells.24_35.rep



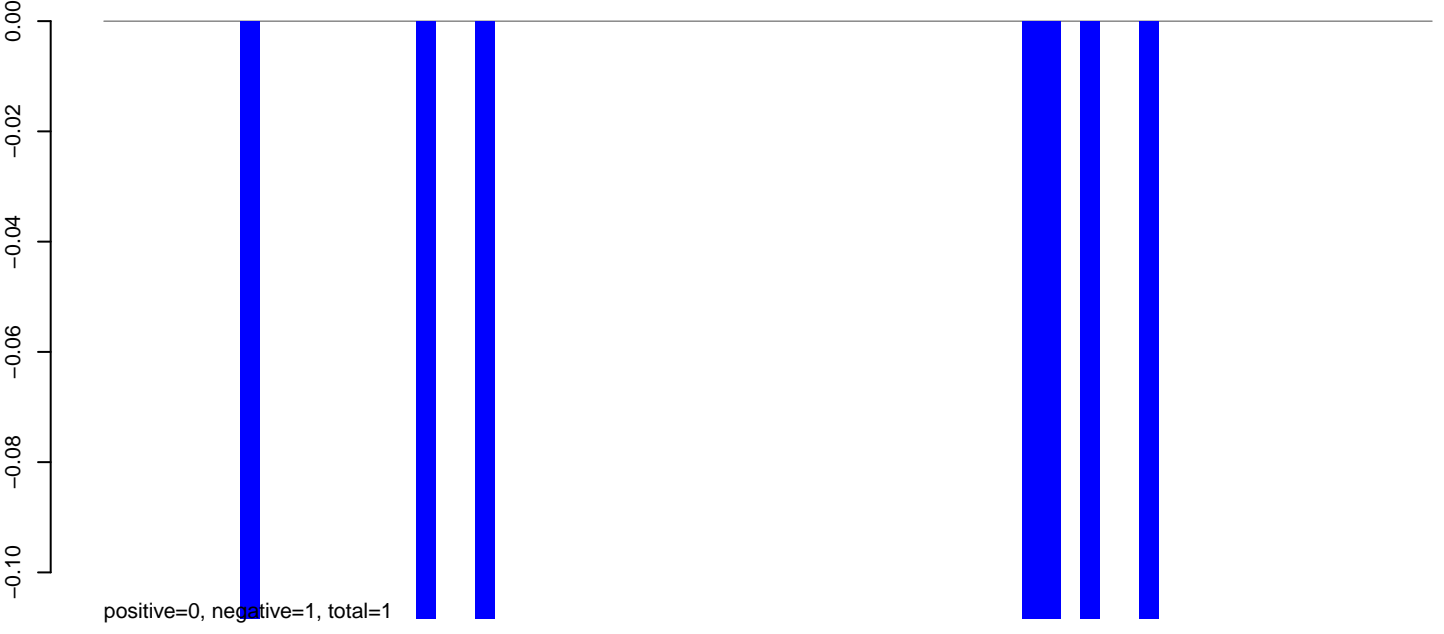
AnGam_Sua5b_cells.rep



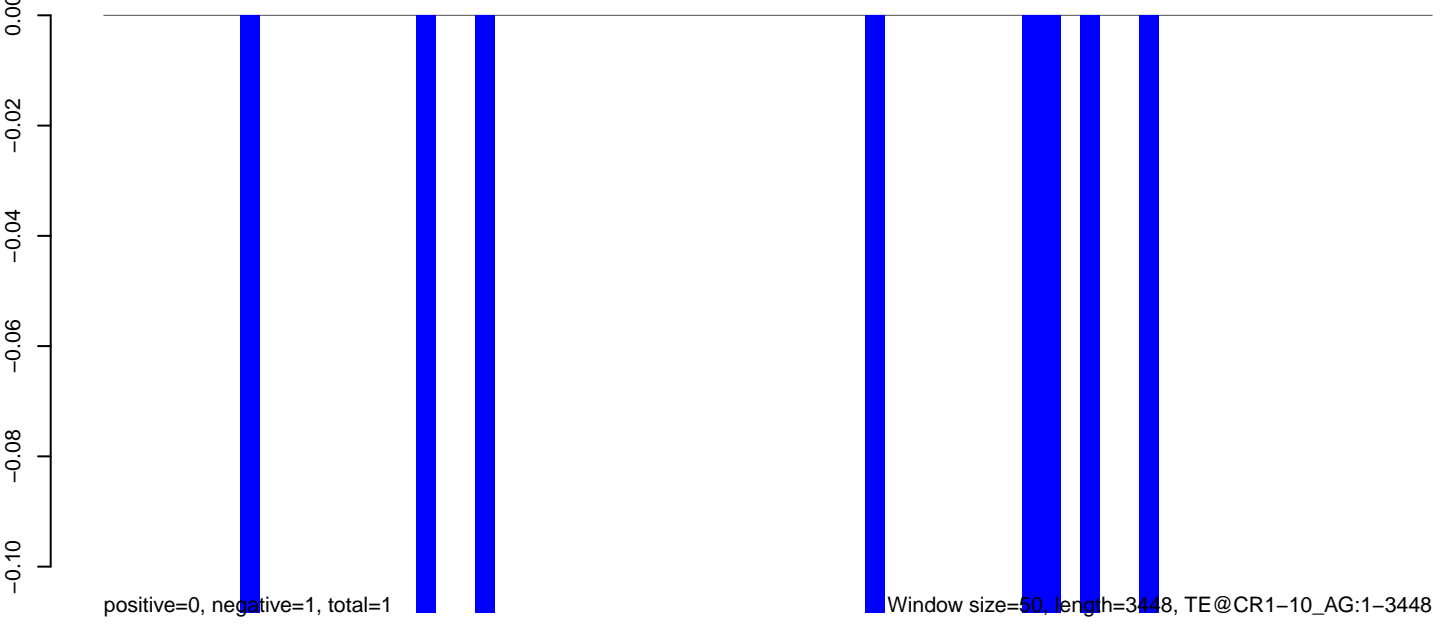
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

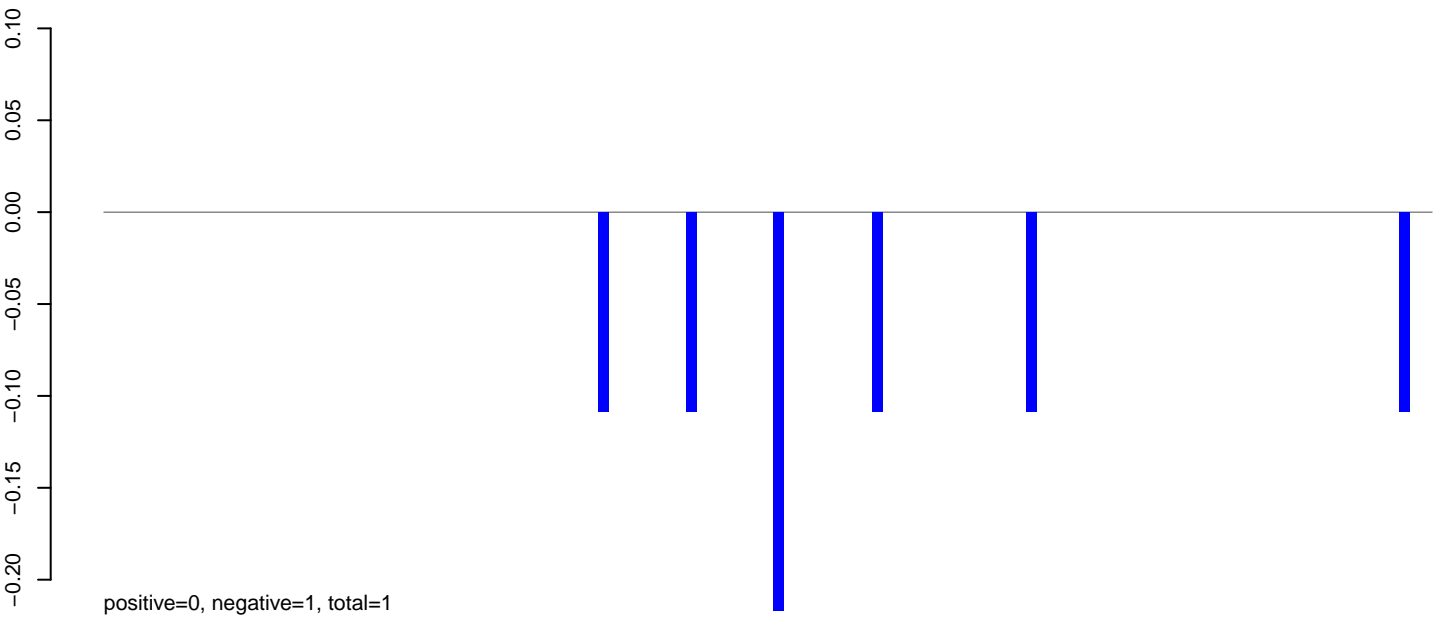


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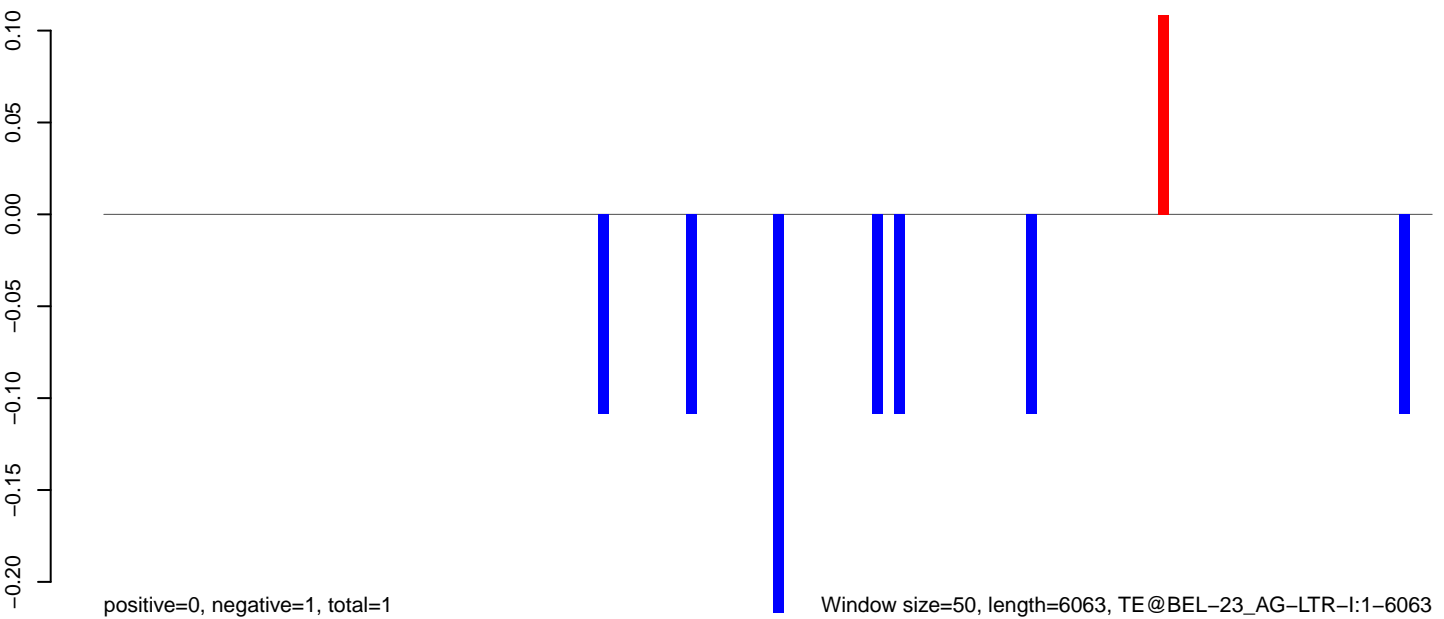
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



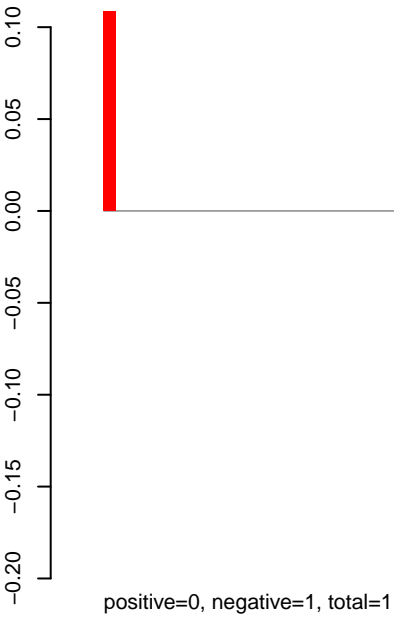
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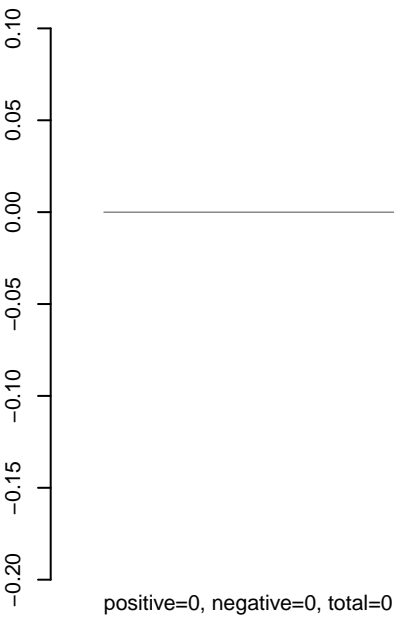
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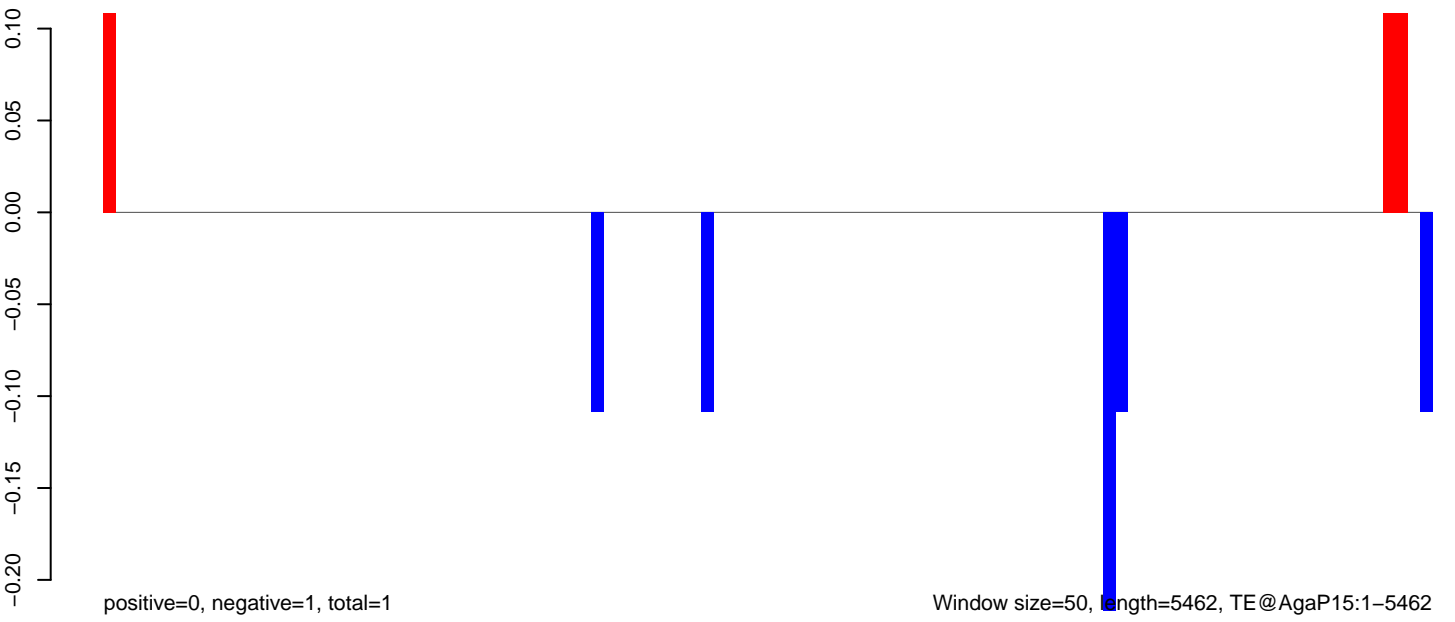
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

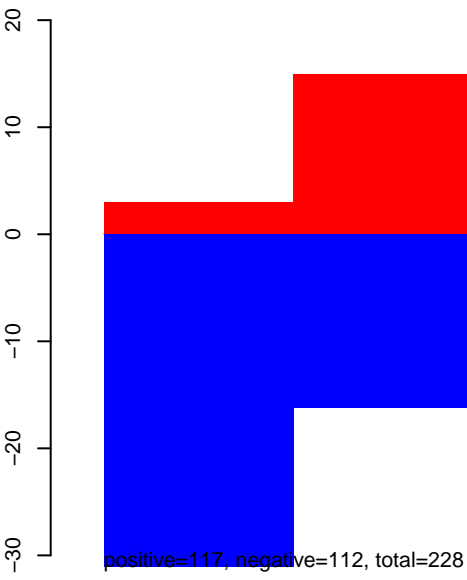


AnGam_Sua5b_cells.rep

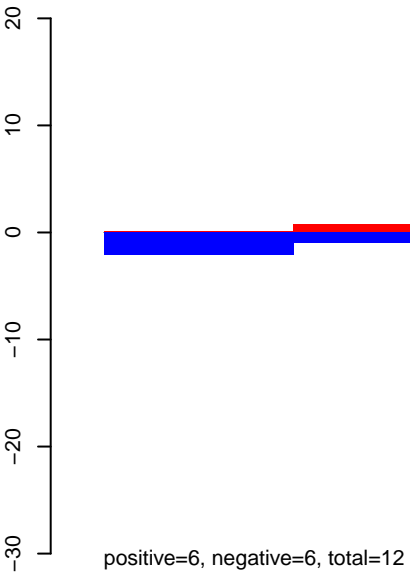


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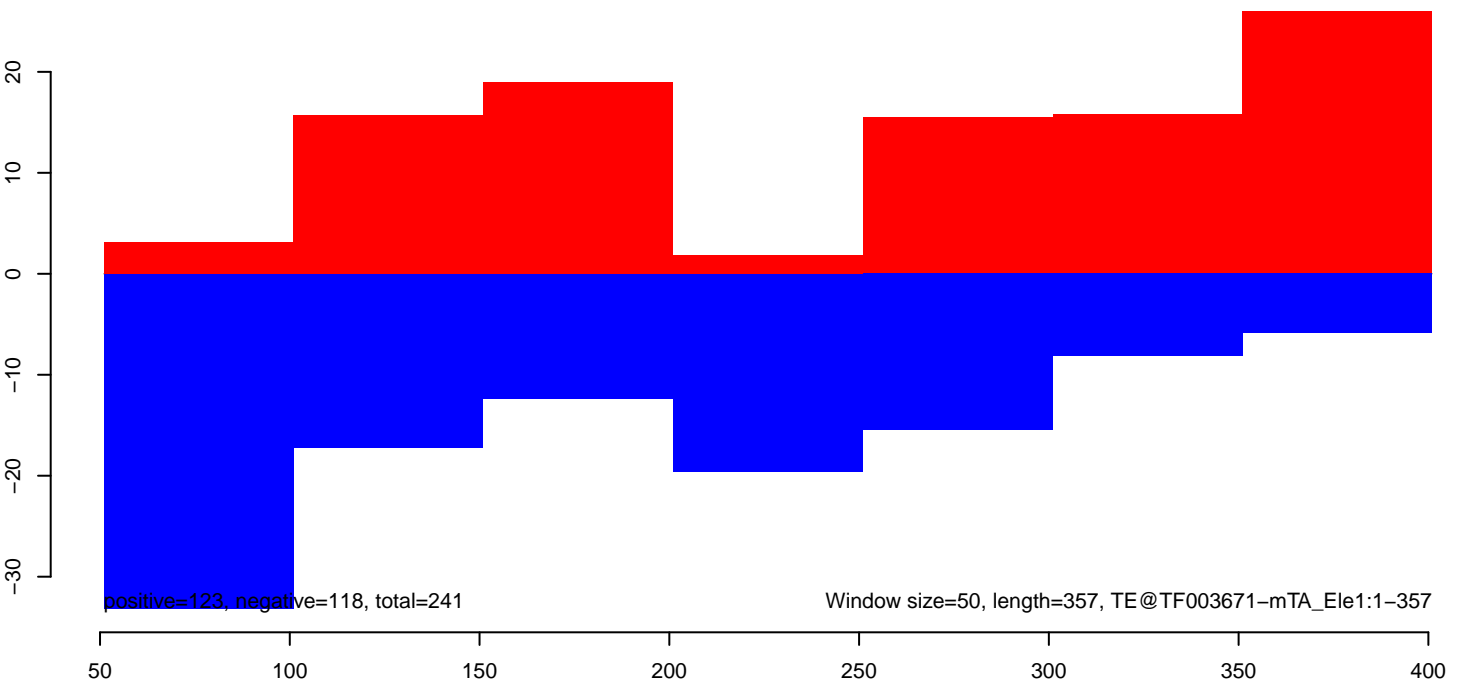
AnGam_Sua5b_cells.18_23.rep



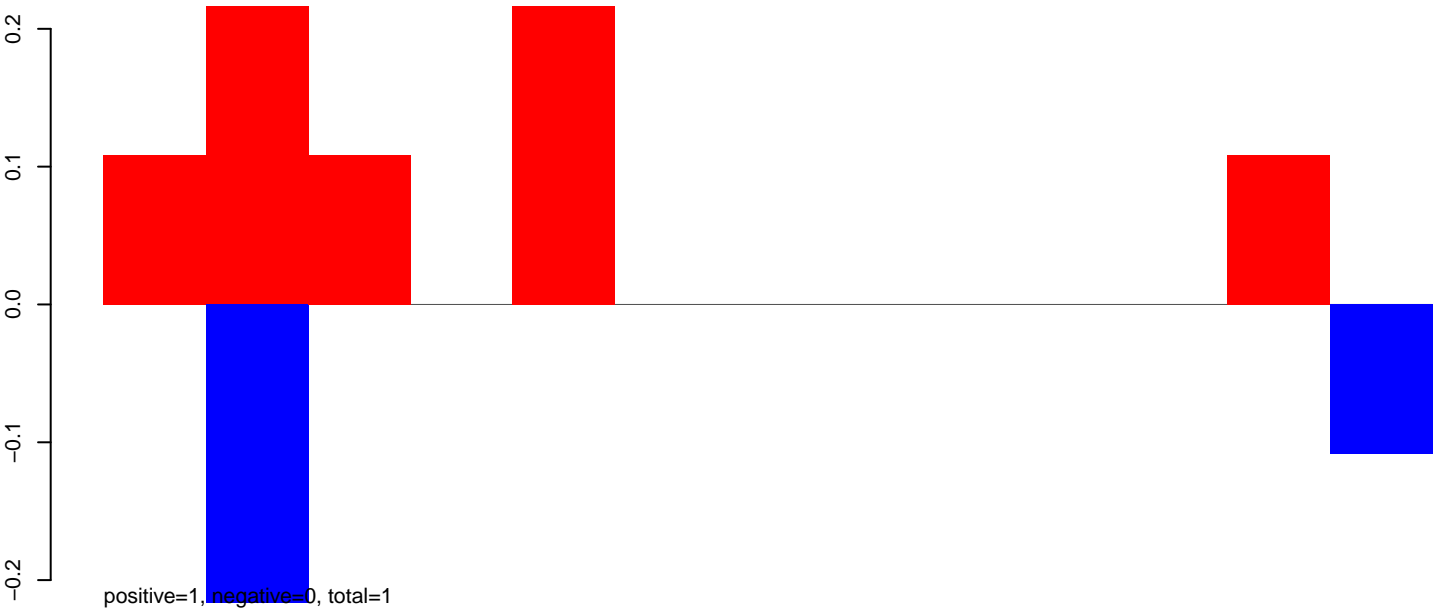
AnGam_Sua5b_cells.24_35.rep



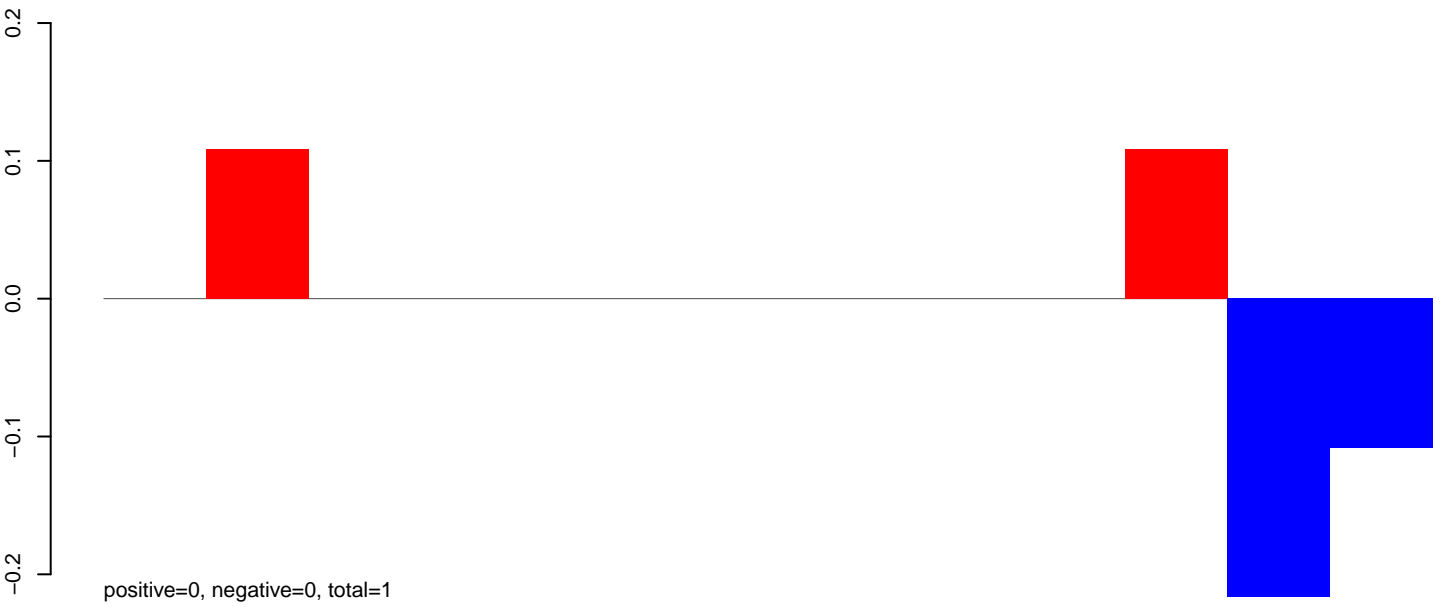
AnGam_Sua5b_cells.rep



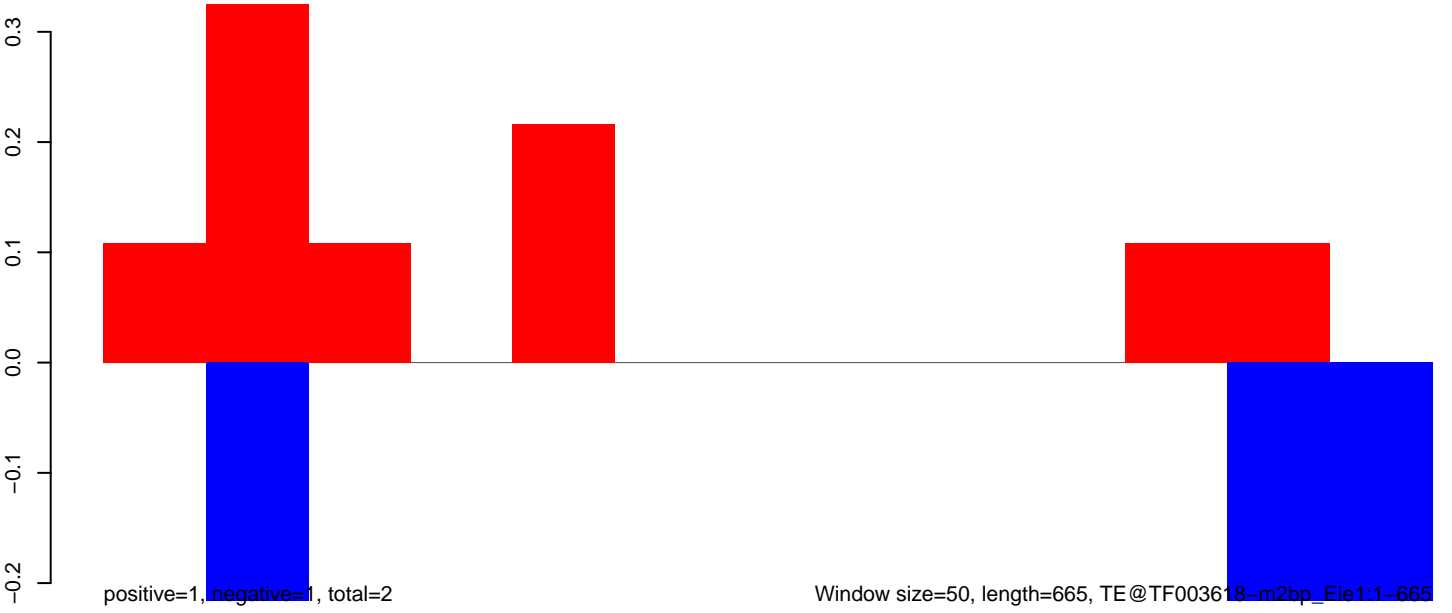
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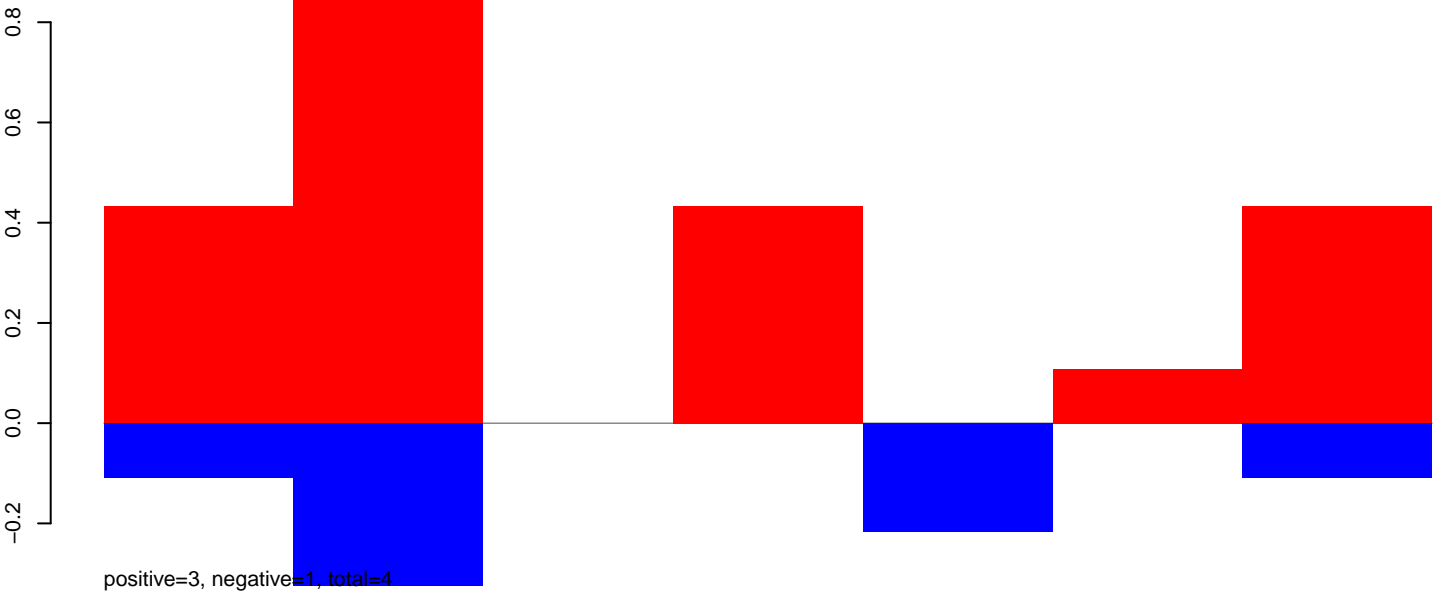
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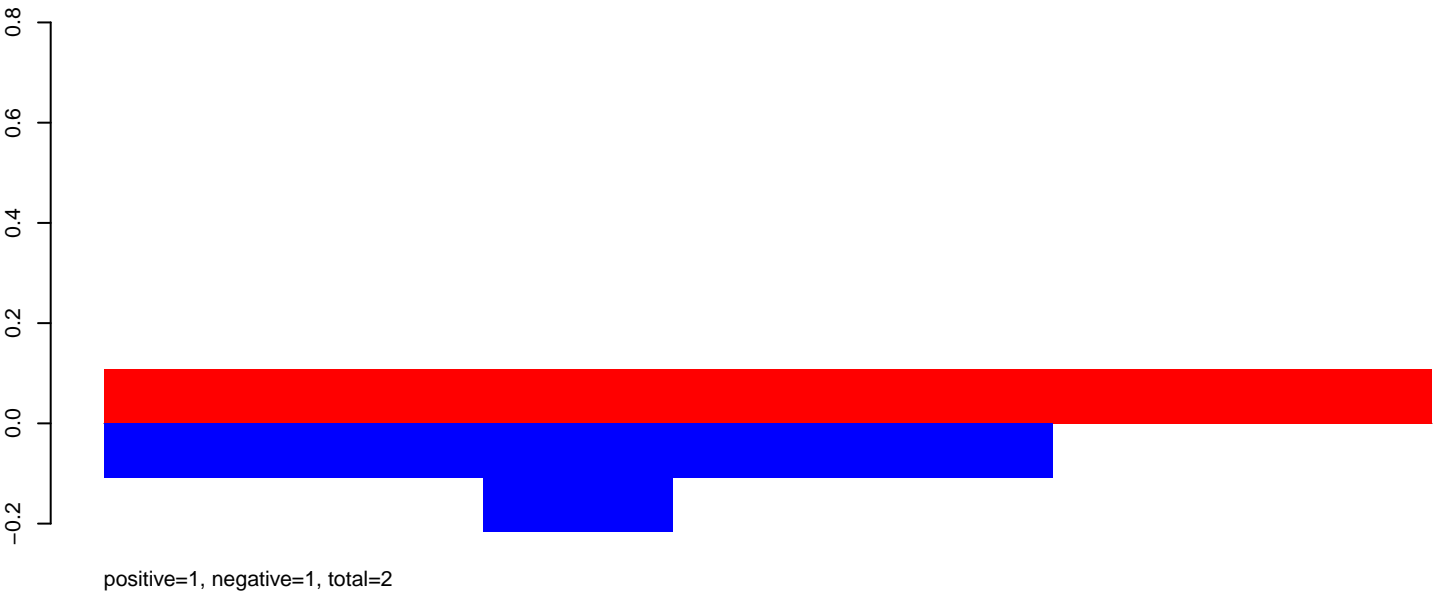
AnGam_Sua5b_cells.rep



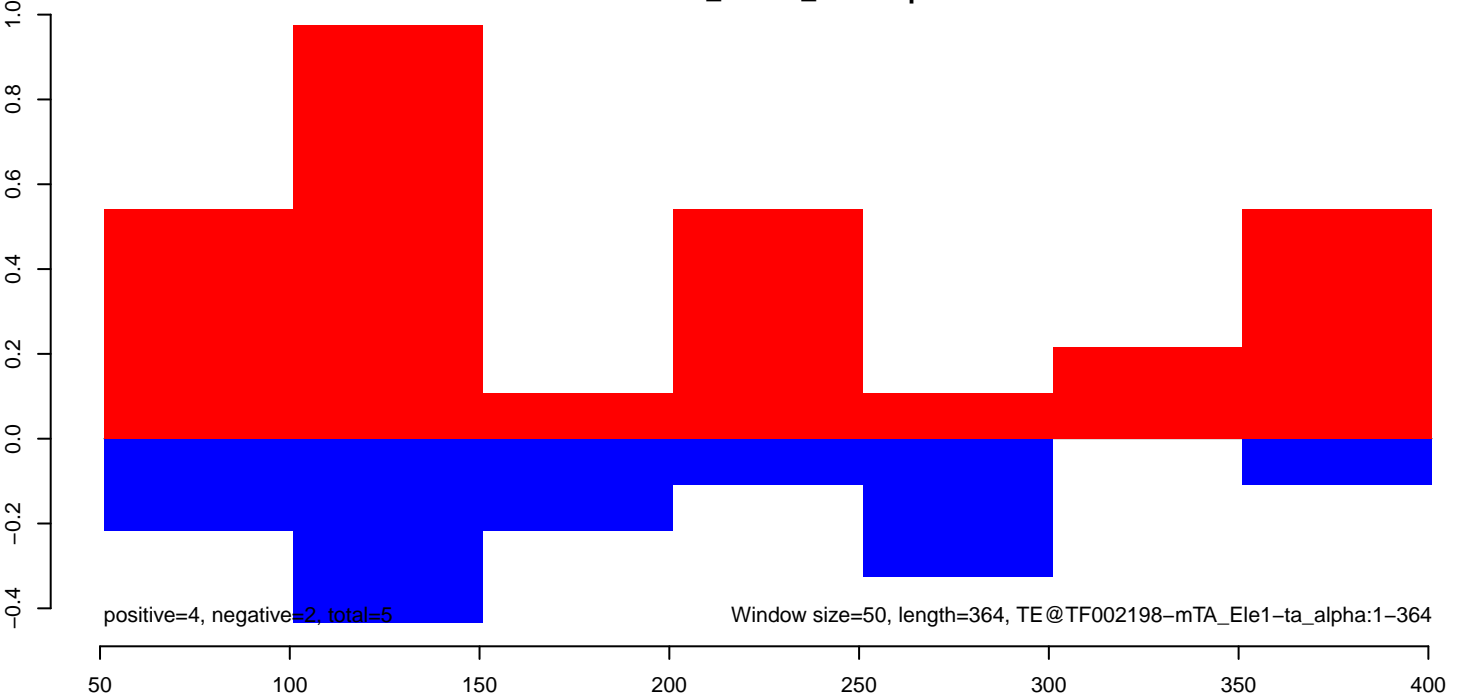
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



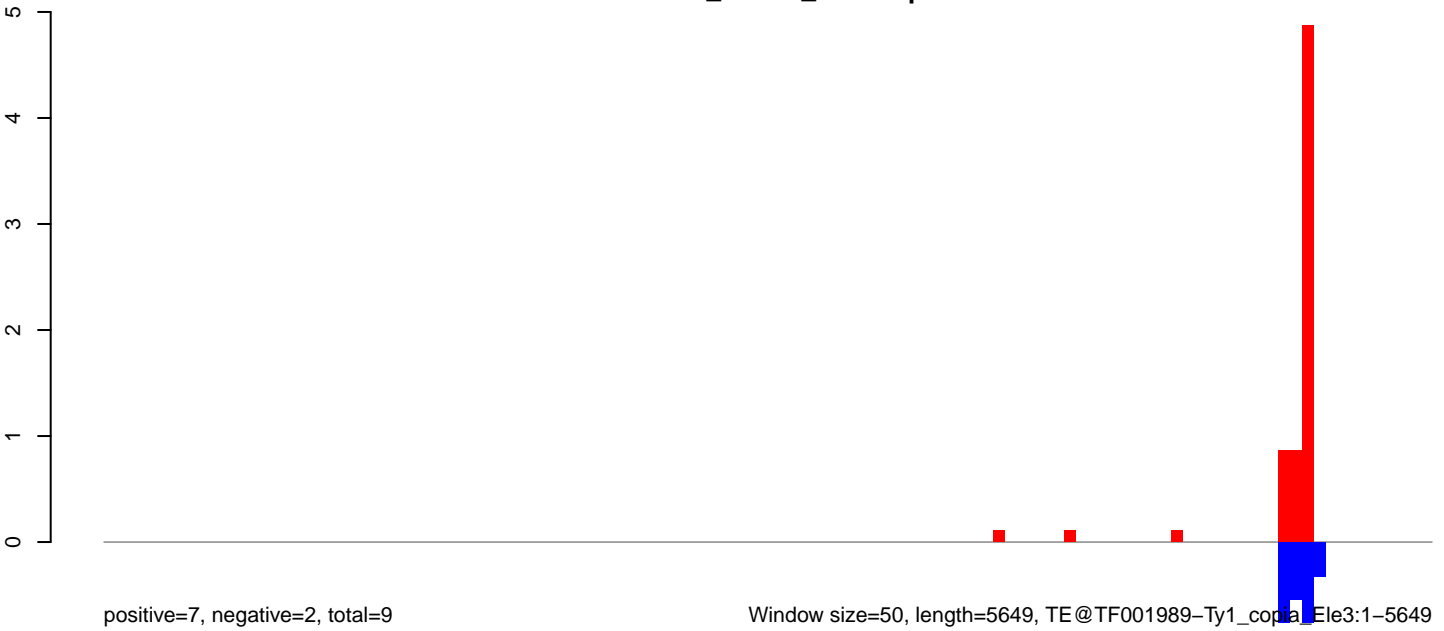
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



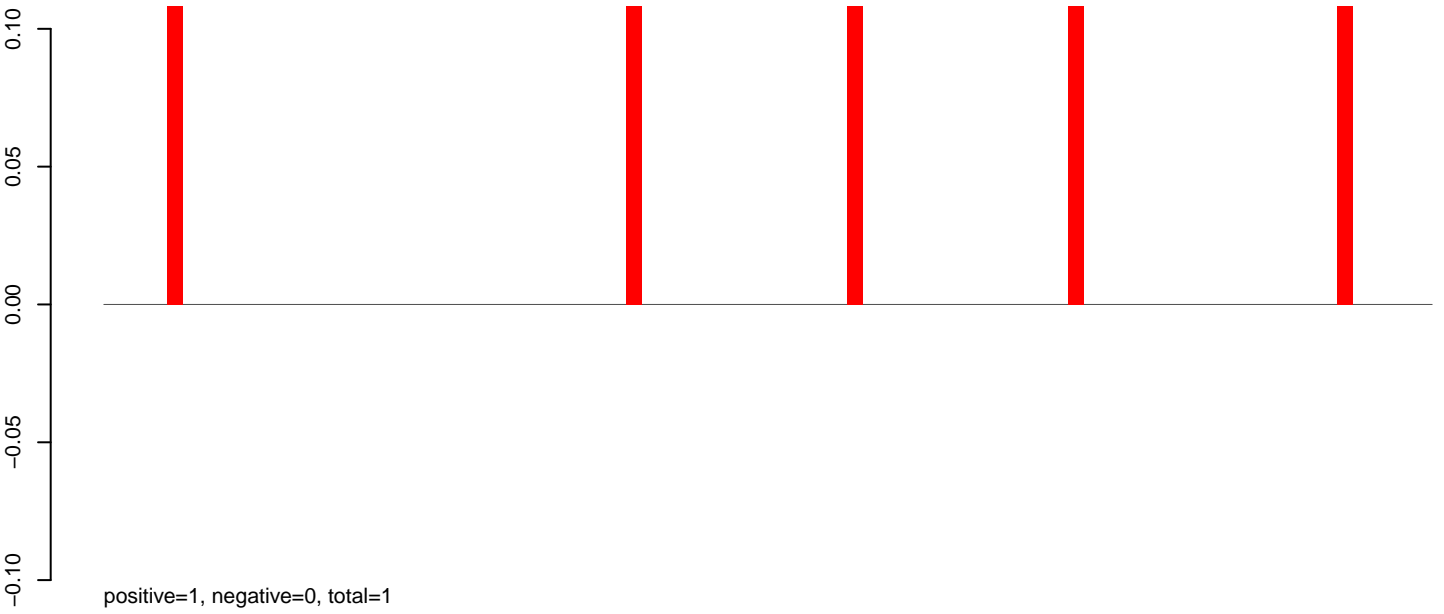
AnGam_Sua5b_cells.rep



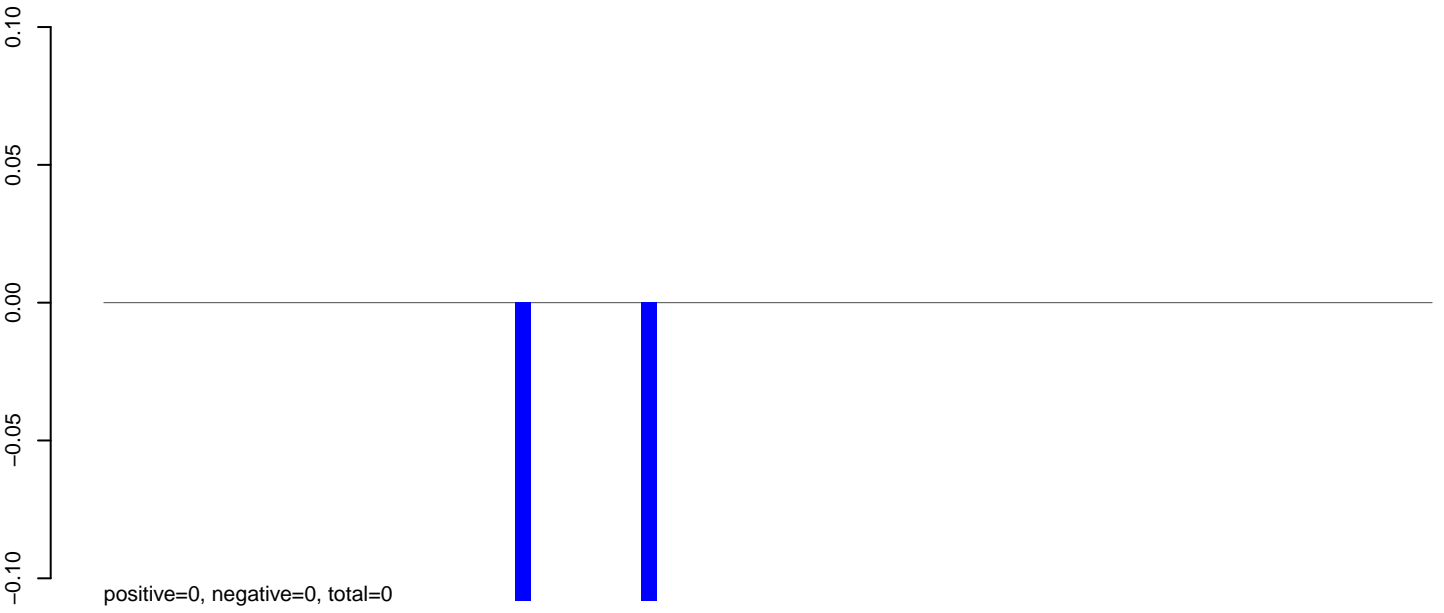
Window size=50, length=5649, TE@TF001989-Ty1_copia_Ele3:1-5649

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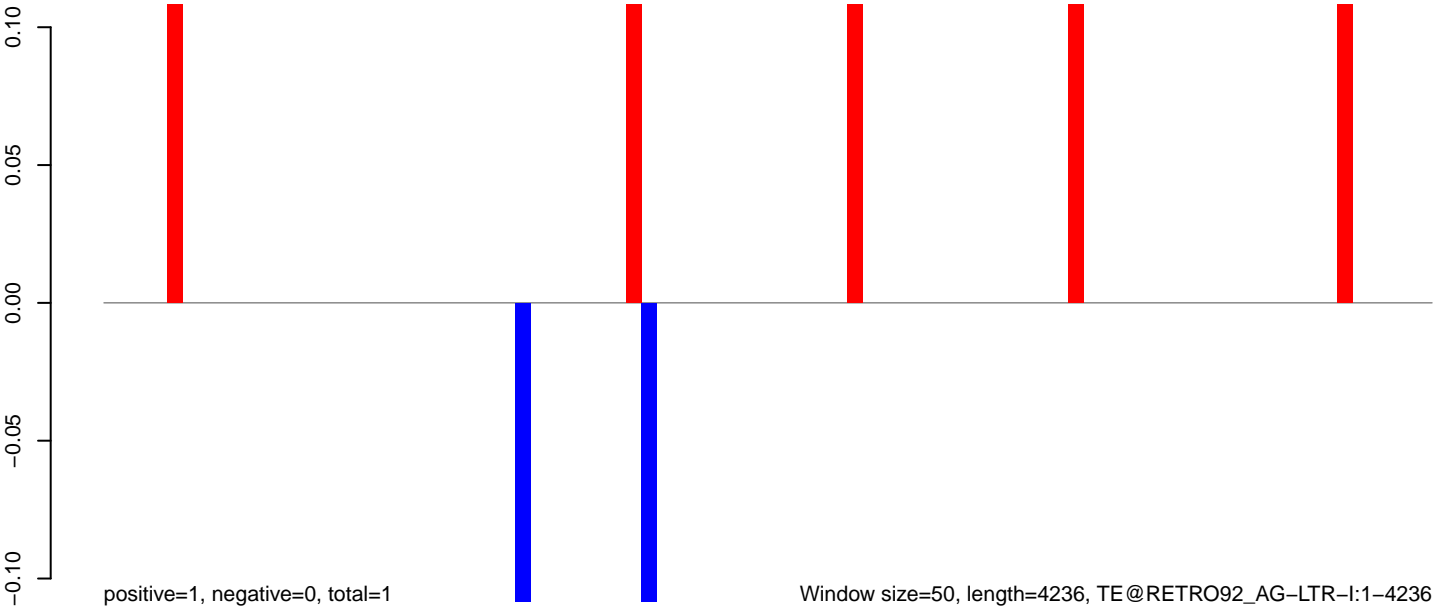
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



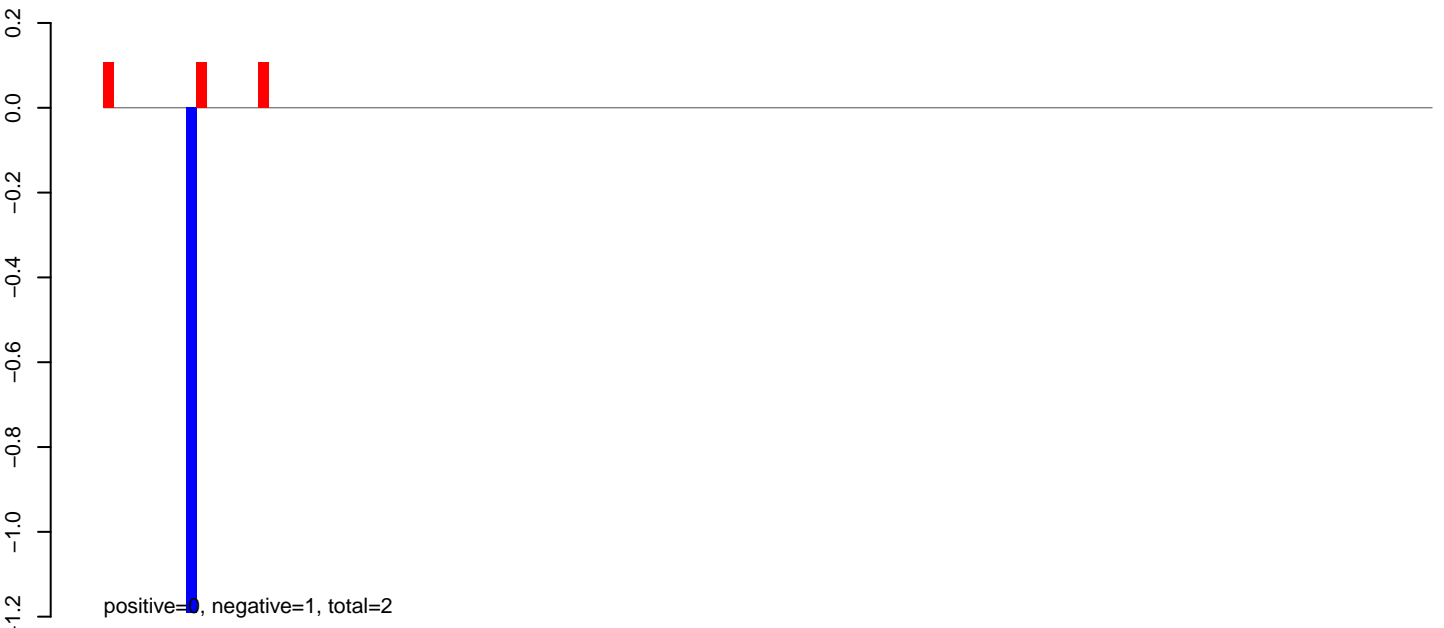
AnGam_Sua5b_cells.rep



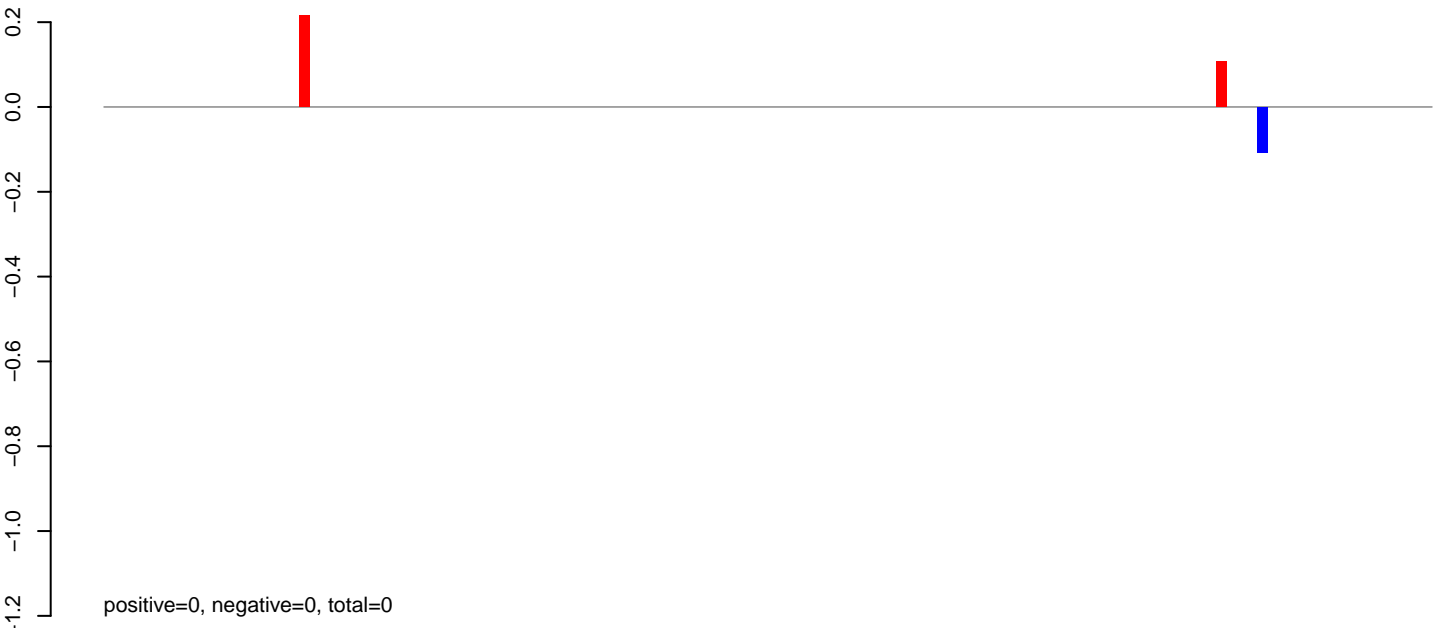
Window size=50, length=4236, TE@RETRO92_AG-LTR-I:1-4236

0 1000 2000 3000 4000

AnGam_Sua5b_cells.18_23.rep



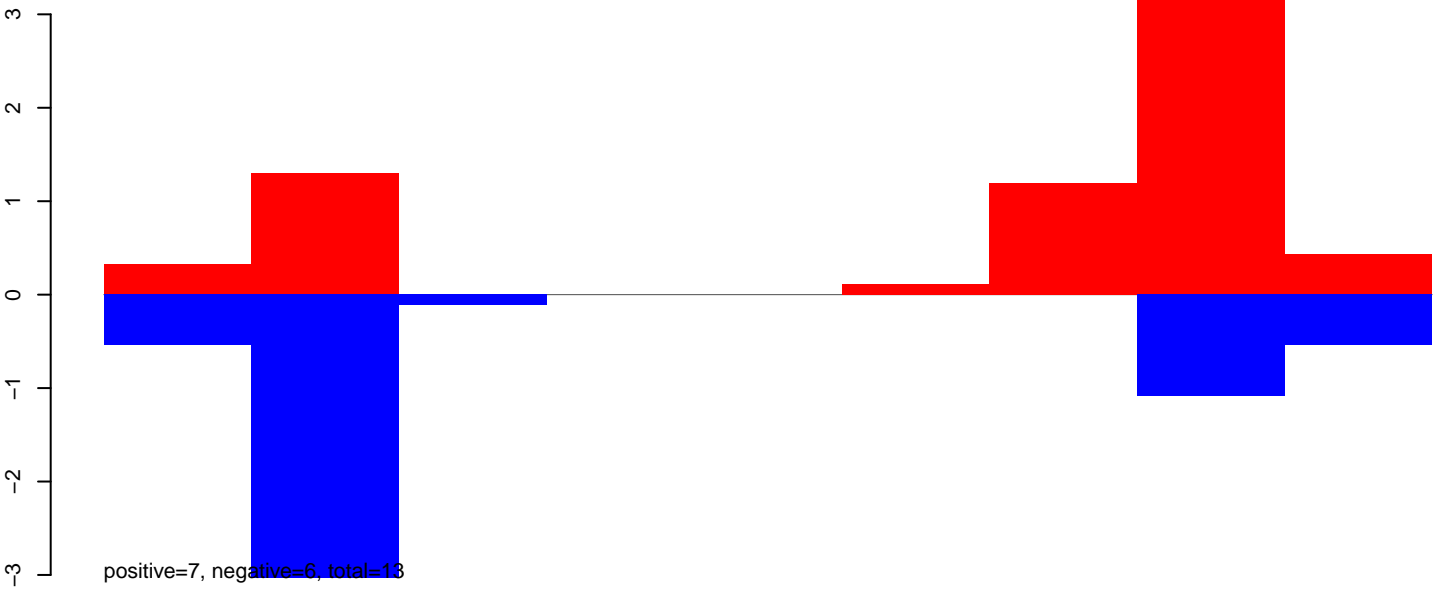
AnGam_Sua5b_cells.24_35.rep



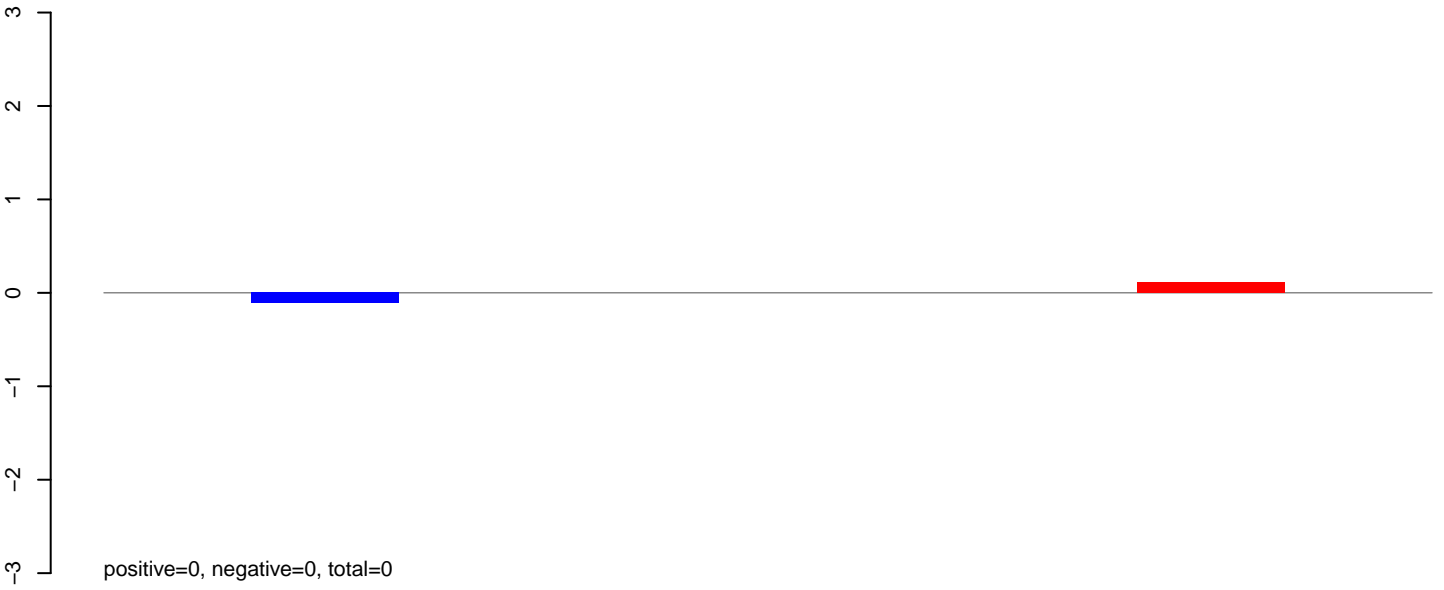
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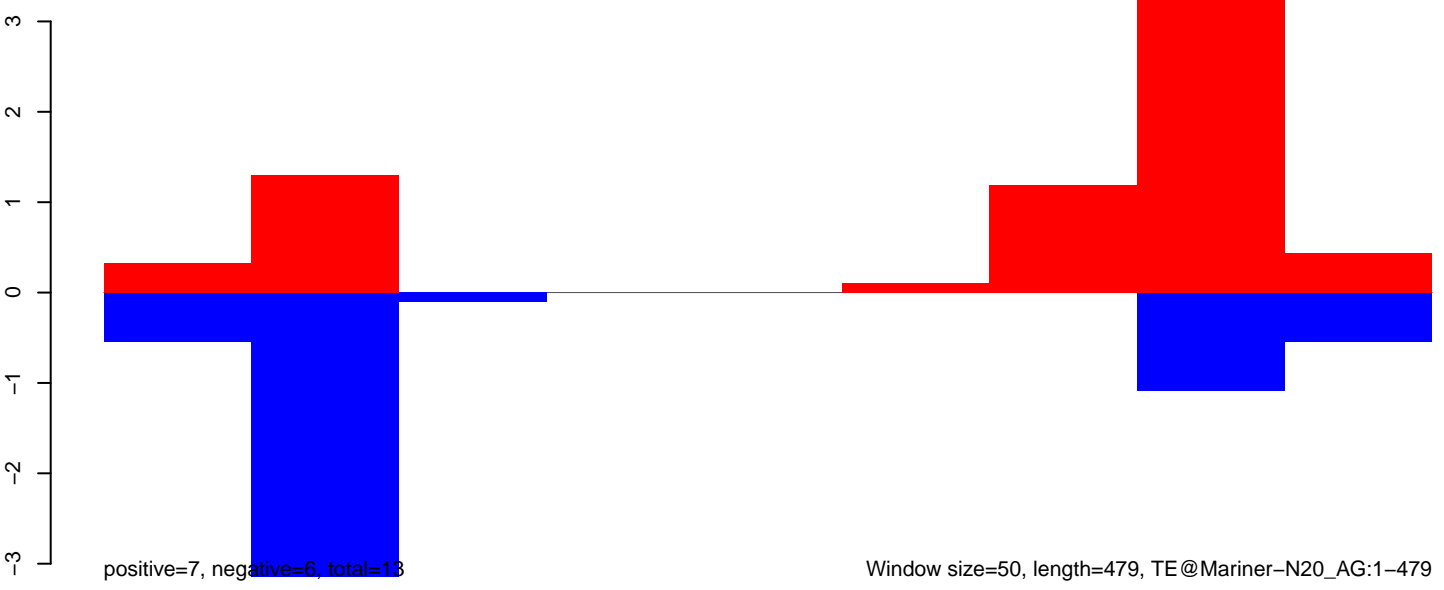
AnGam_Sua5b_cells.18_23.rep



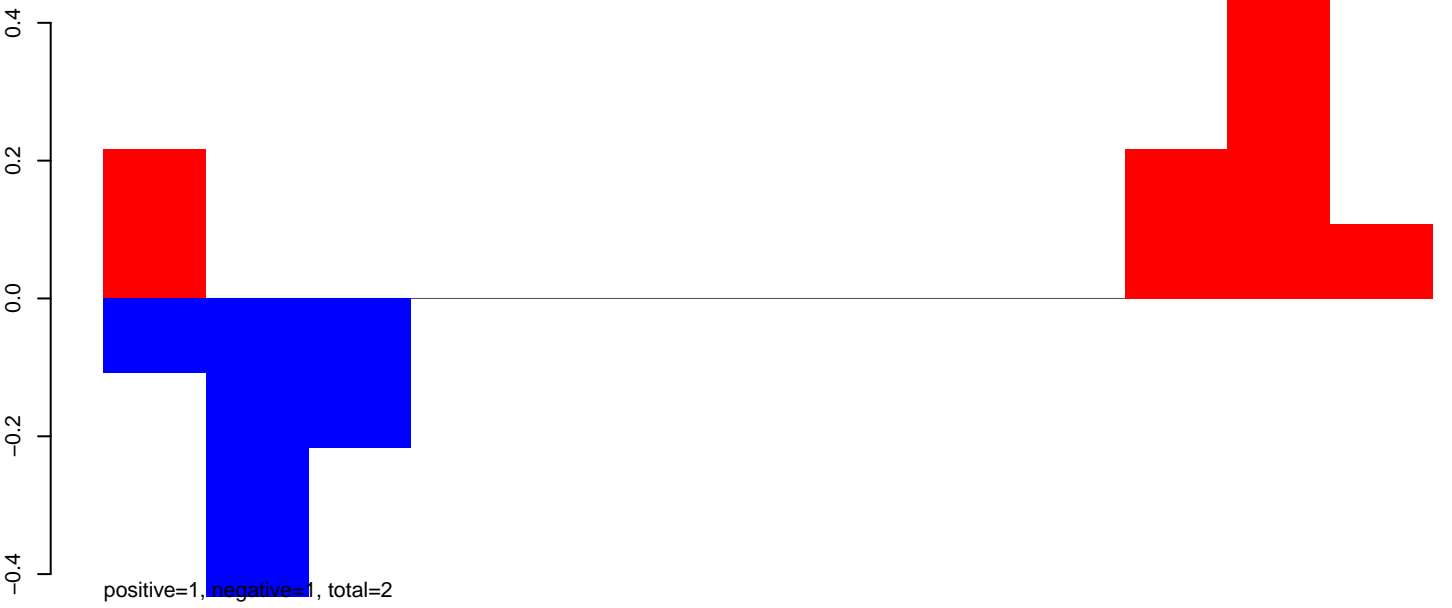
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



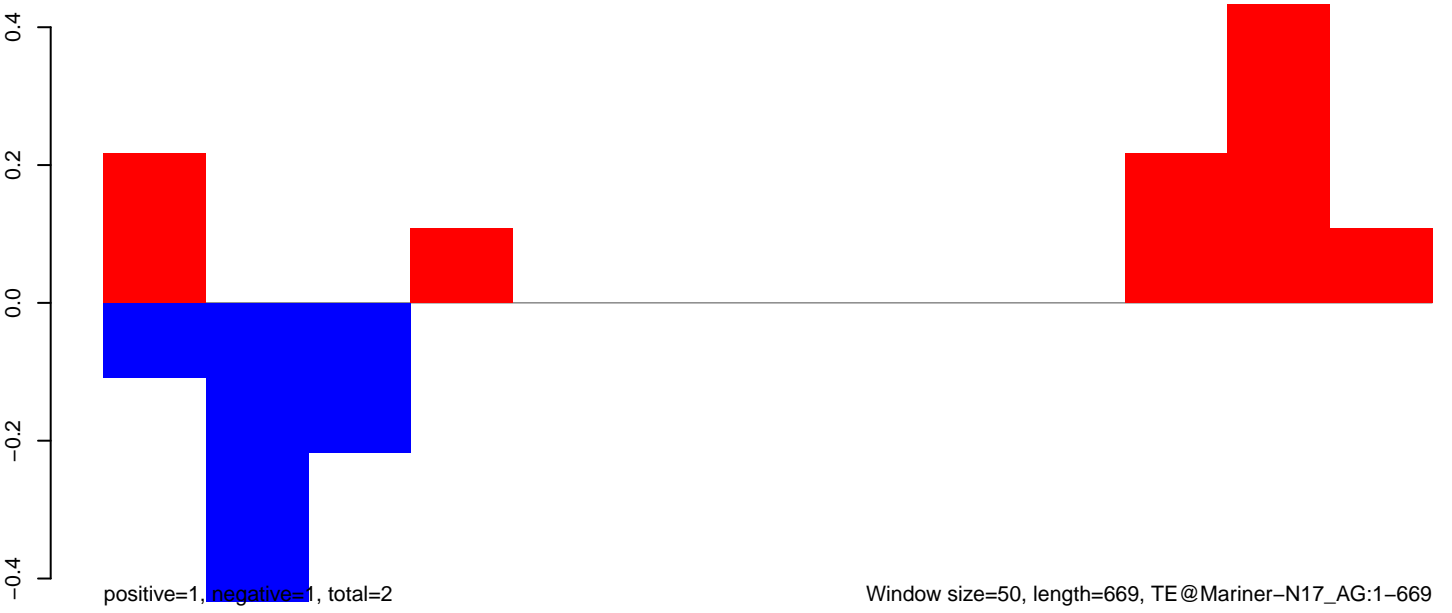
AnGam_Sua5b_cells.18_23.rep



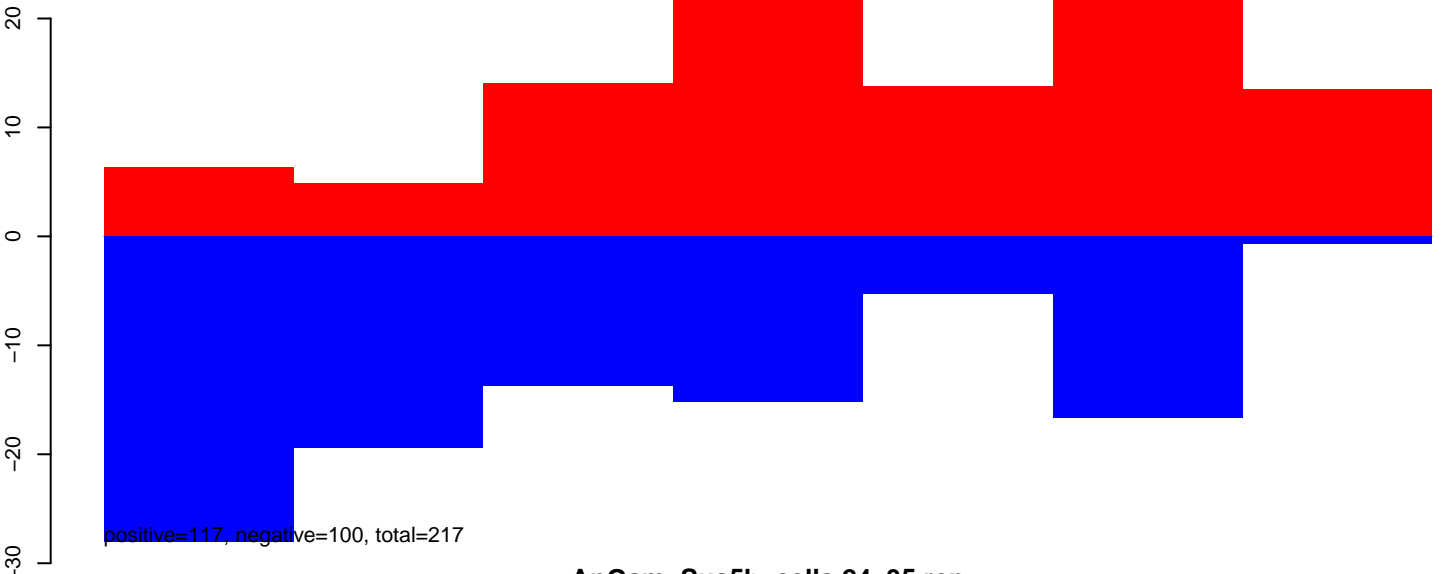
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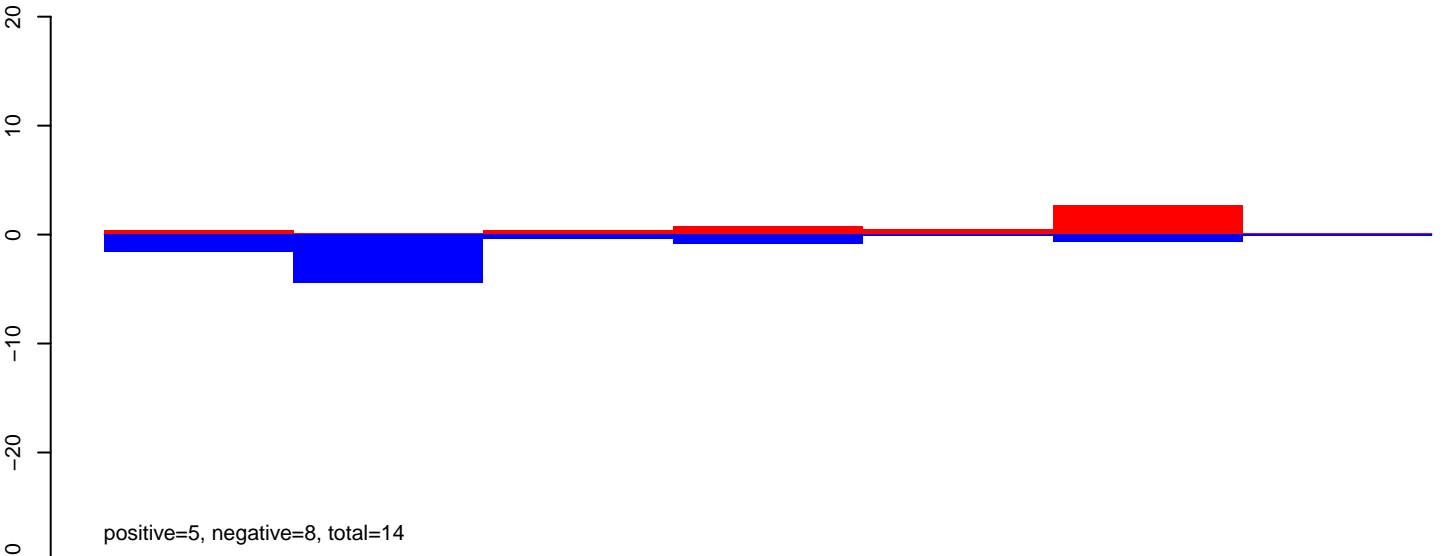
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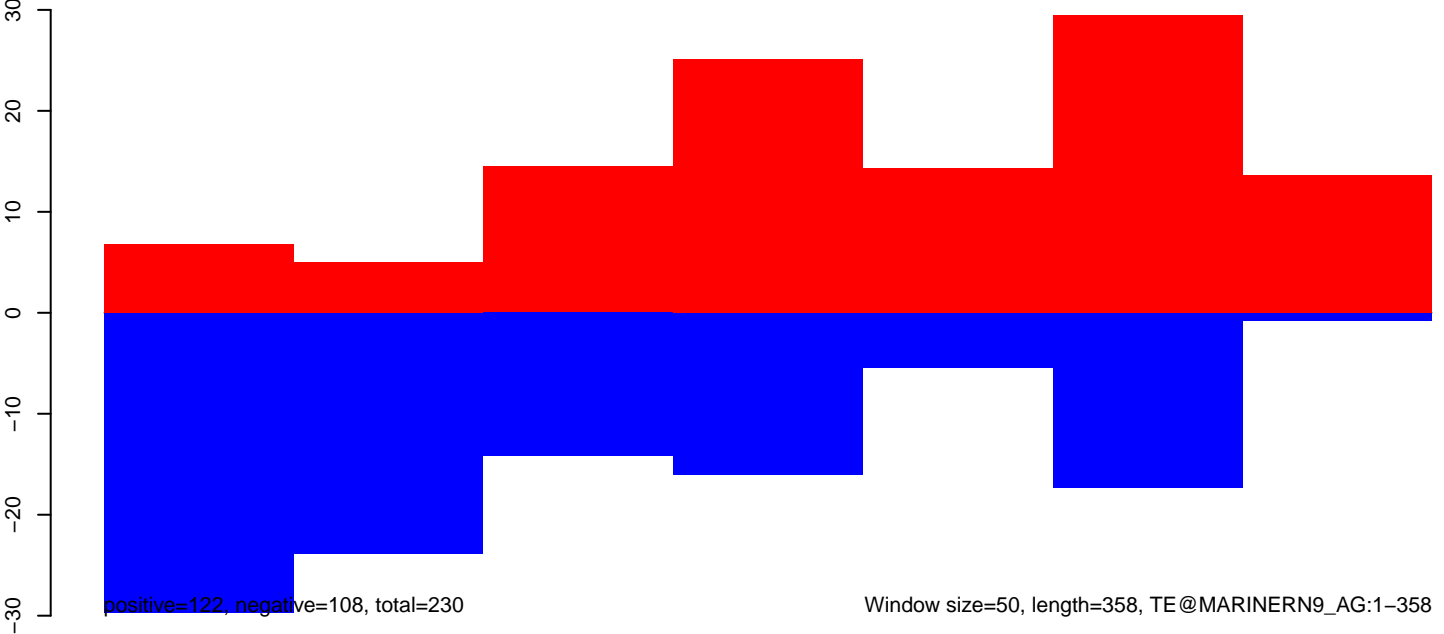
AnGam_Sua5b_cells.18_23.rep



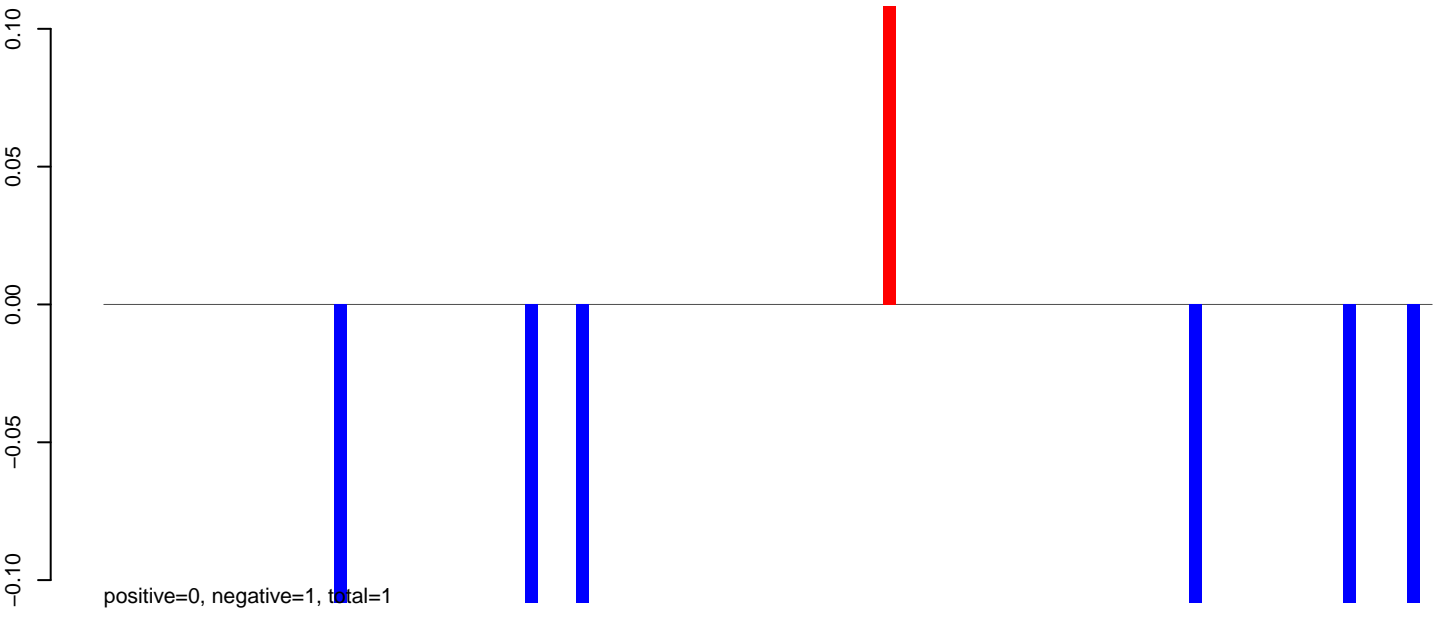
AnGam_Sua5b_cells.24_35.rep



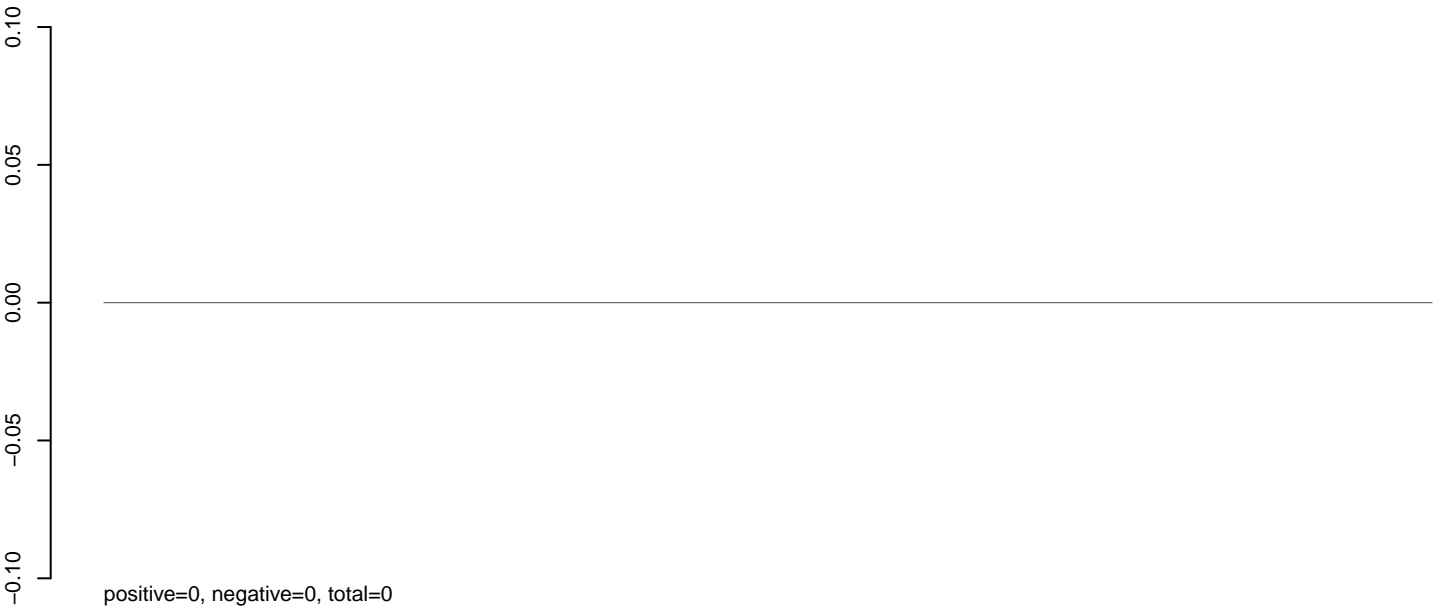
AnGam_Sua5b_cells.rep



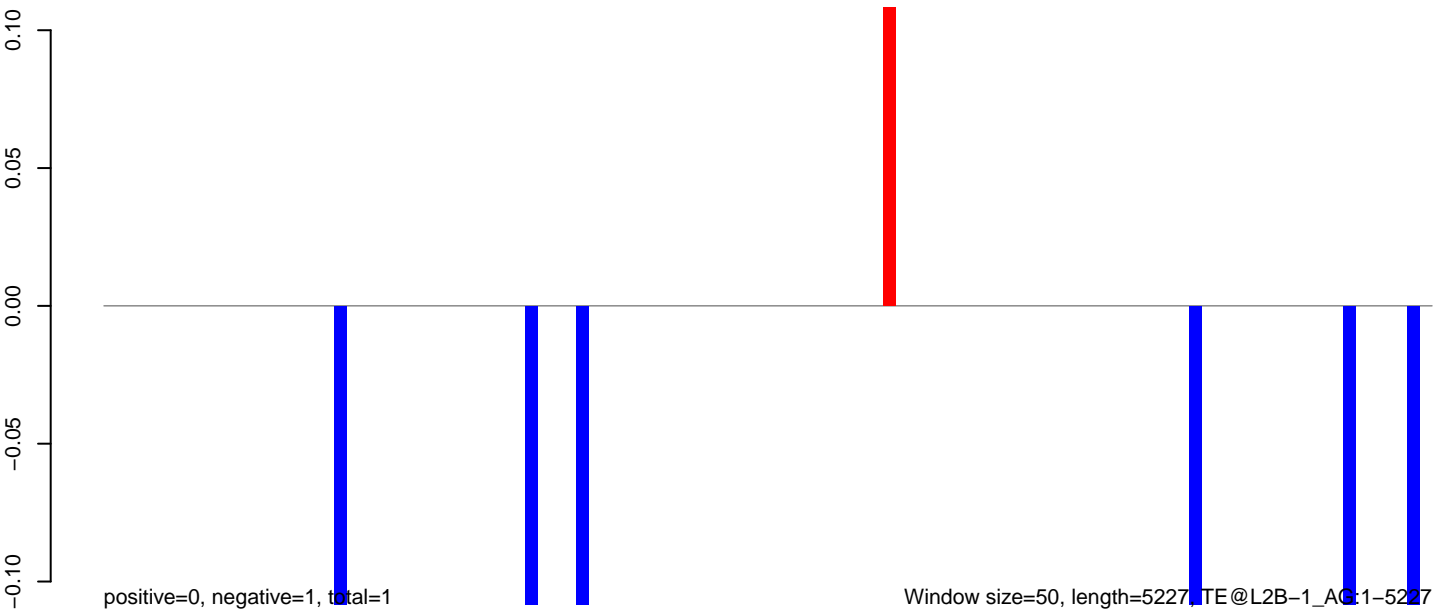
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

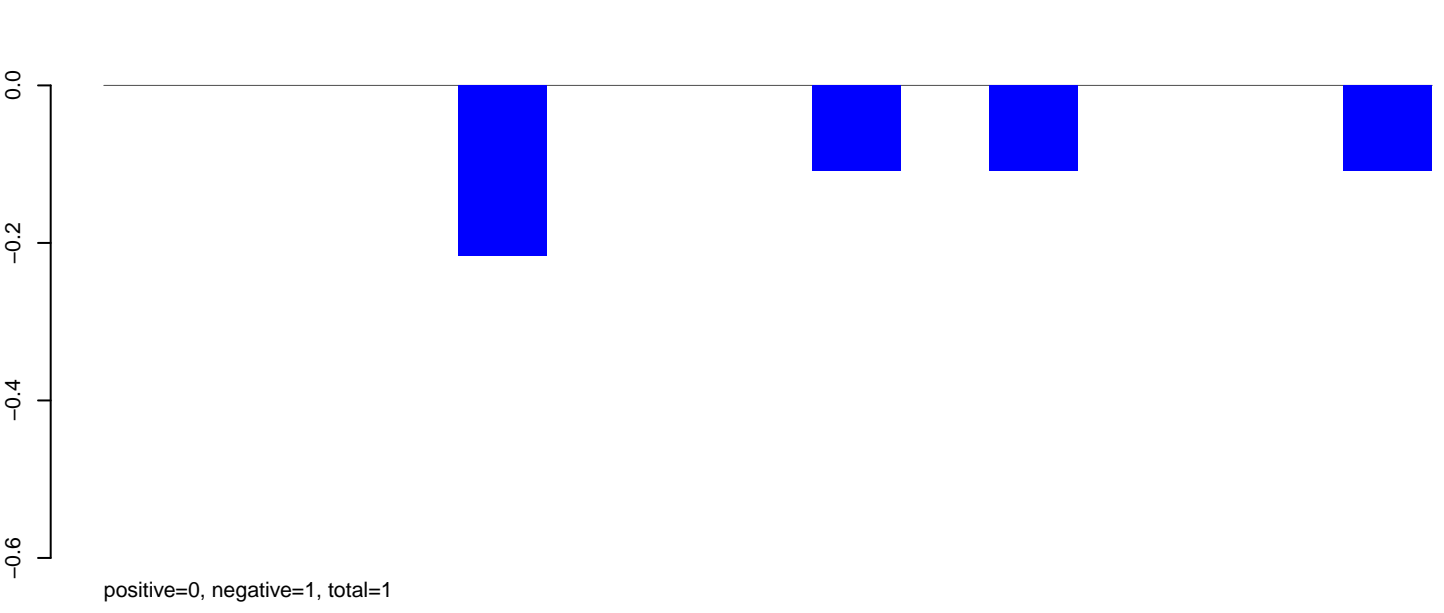


Window size=50, length=5227, TE@L2B-1_AG:1-5227

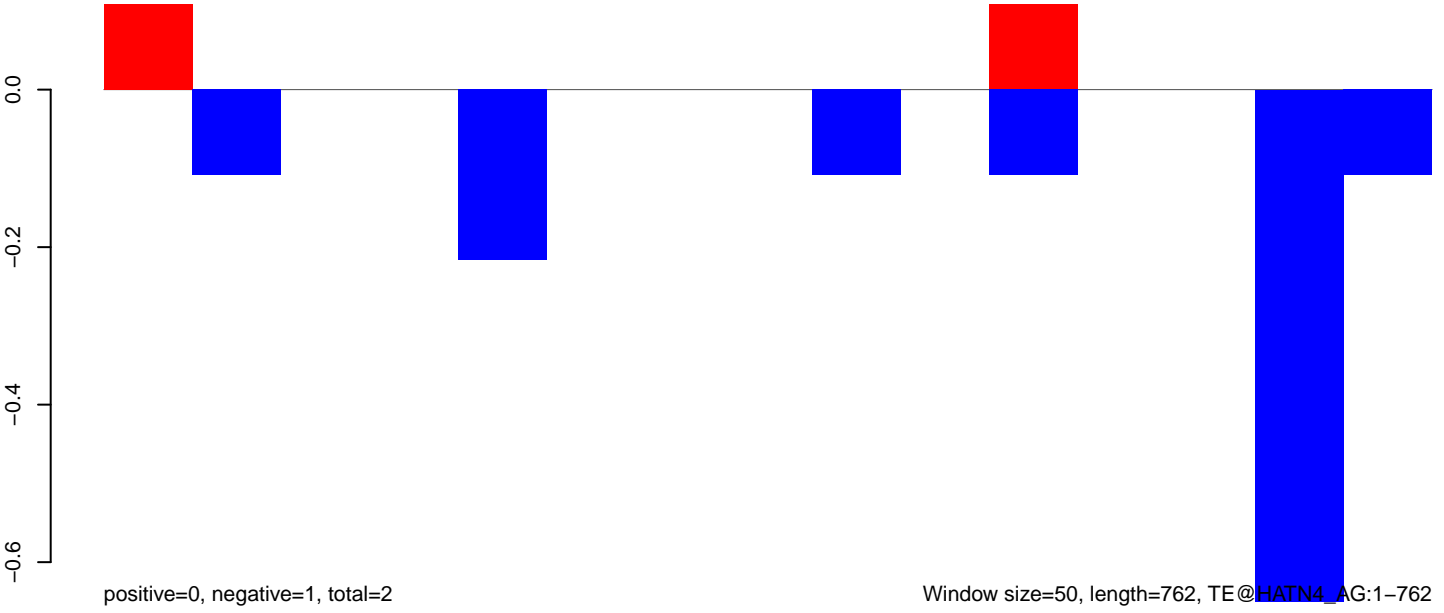
AnGam_Sua5b_cells.18_23.rep



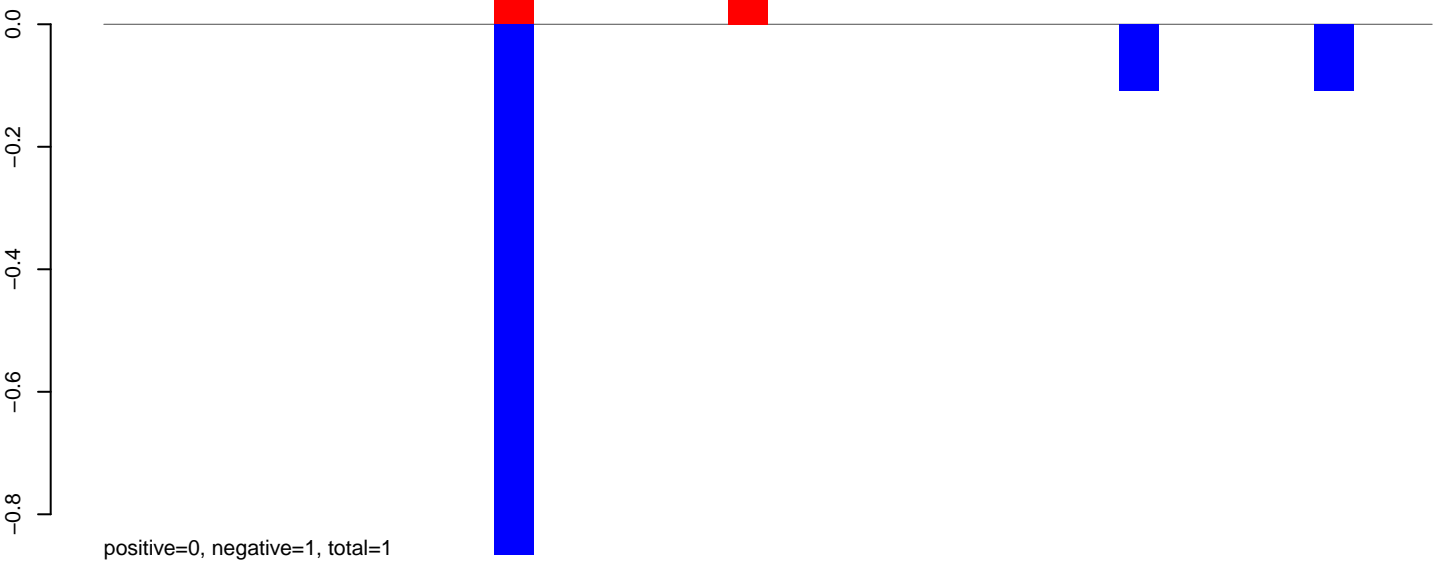
AnGam_Sua5b_cells.24_35.rep



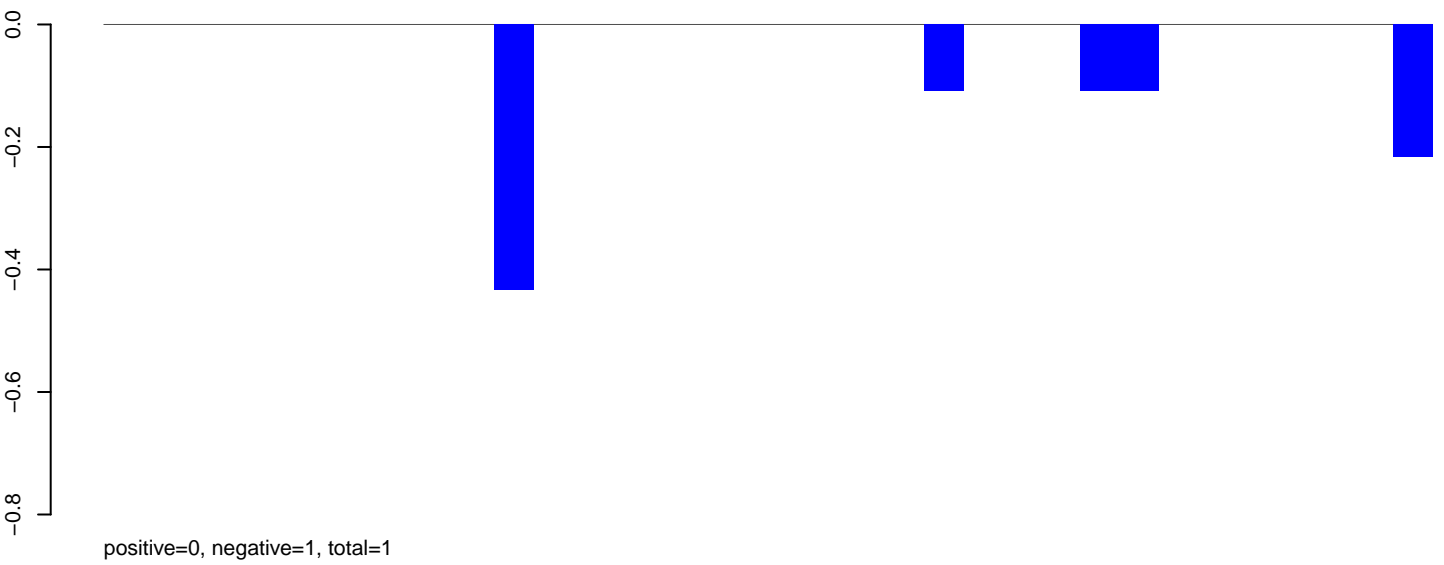
AnGam_Sua5b_cells.rep



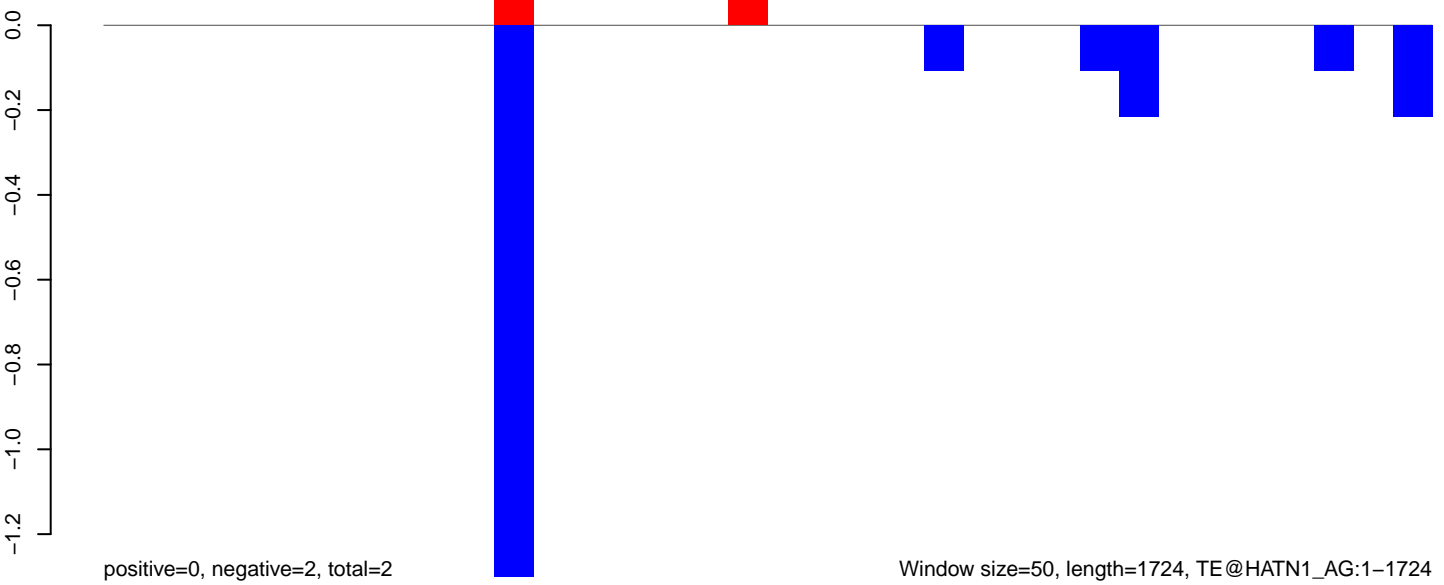
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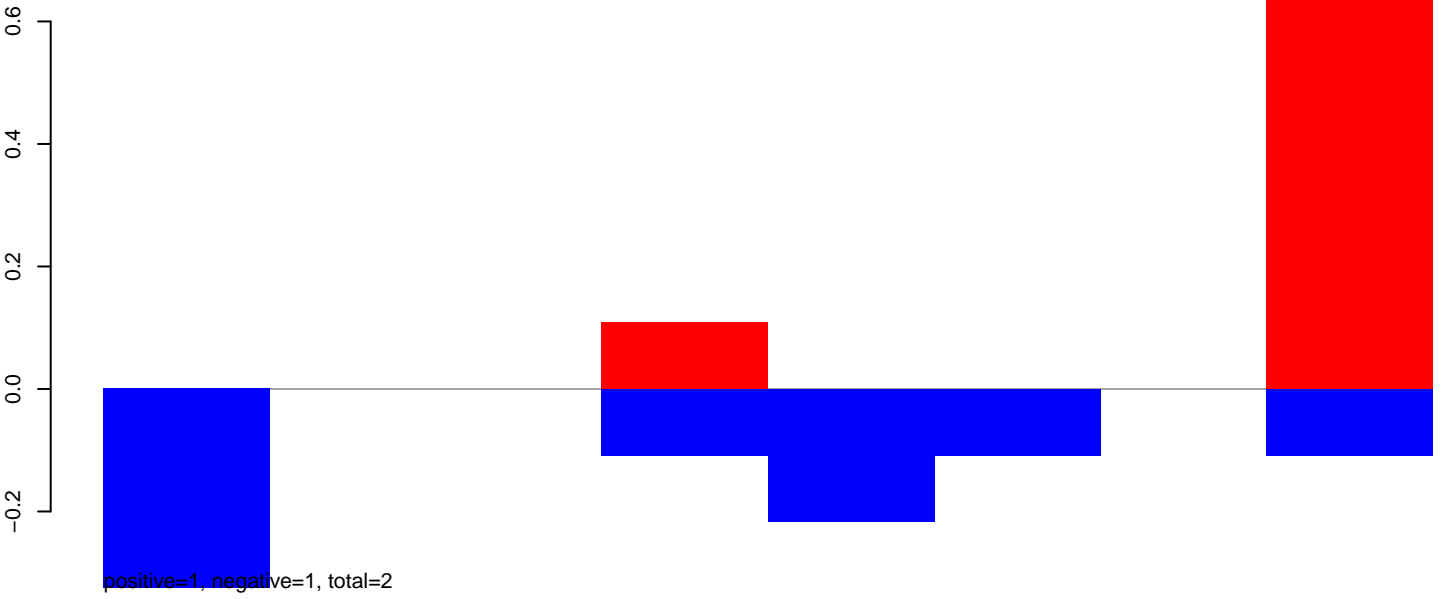
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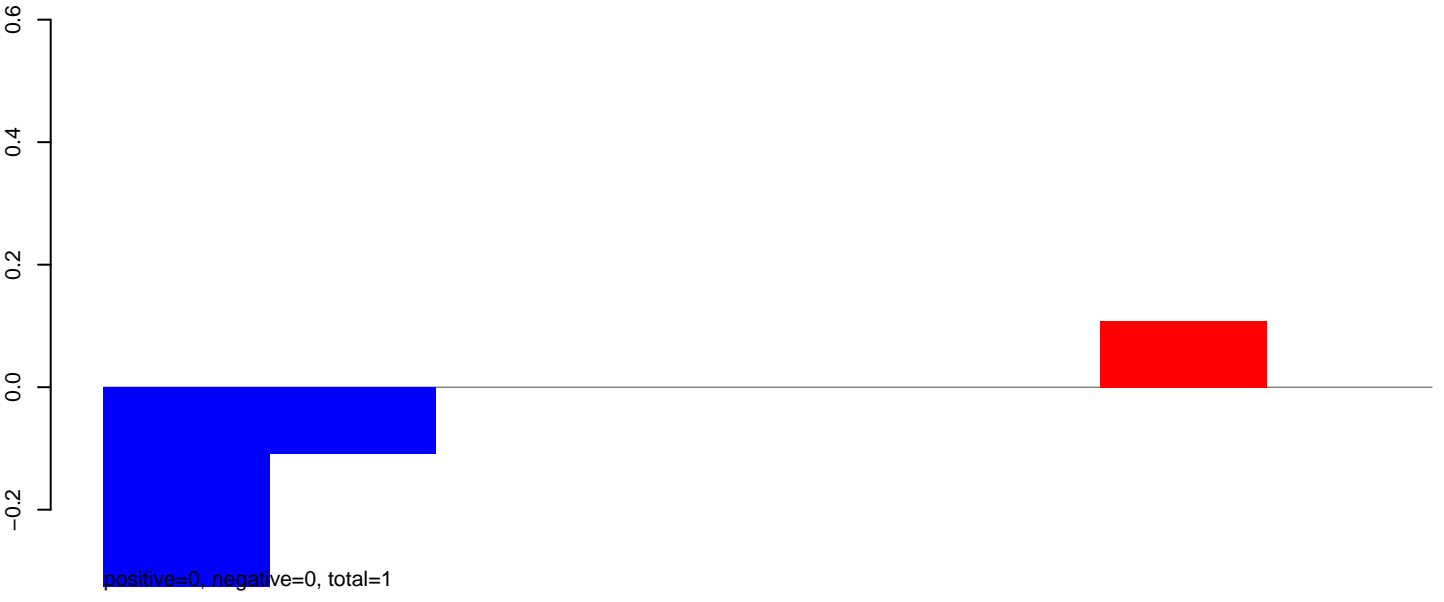
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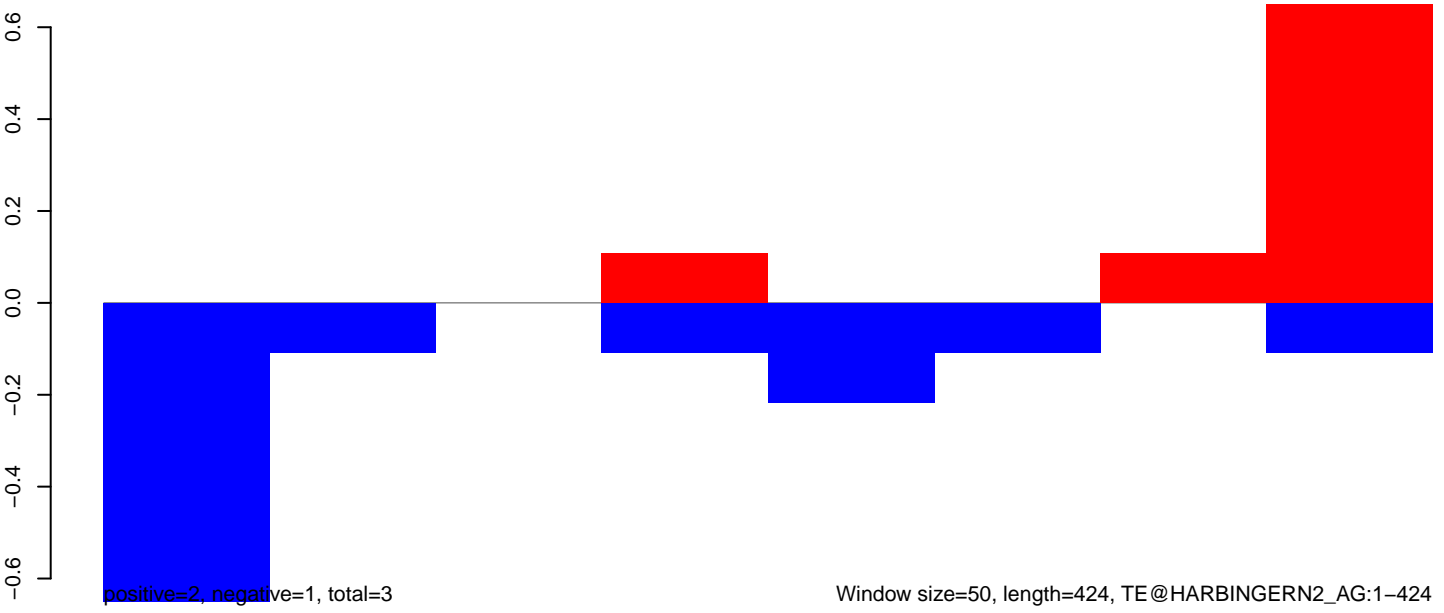
AnGam_Sua5b_cells.18_23.rep



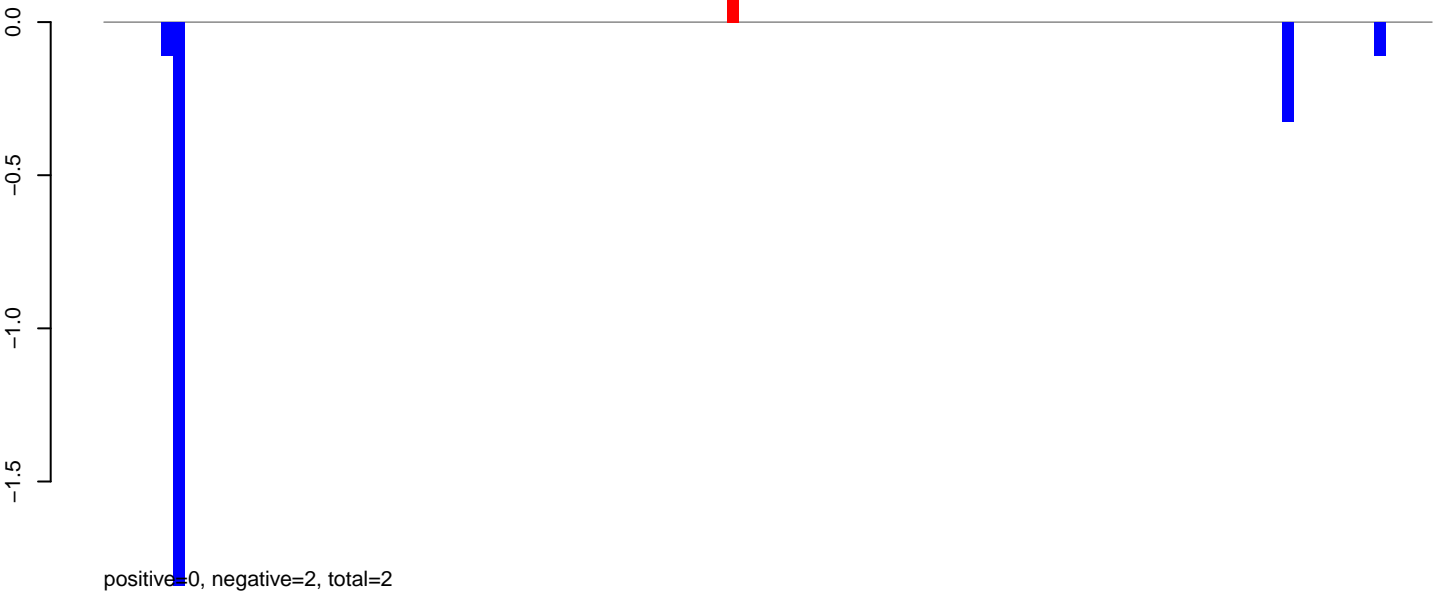
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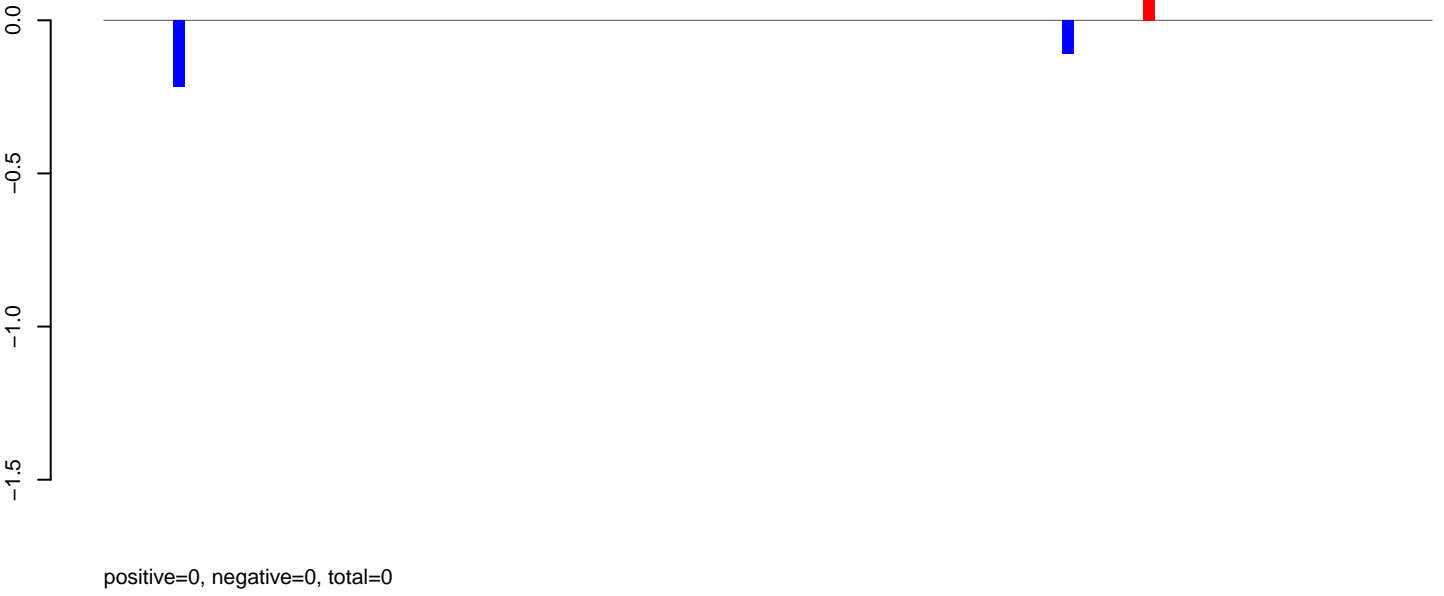
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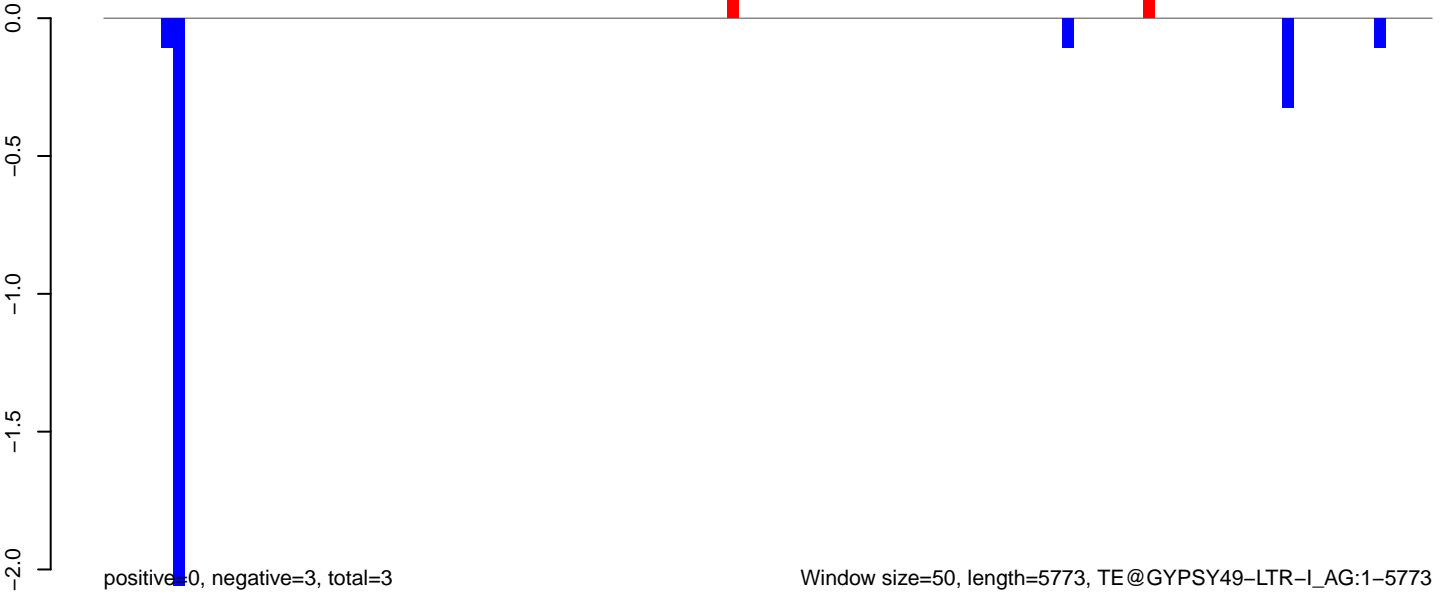
AnGam_Sua5b_cells.18_23.rep



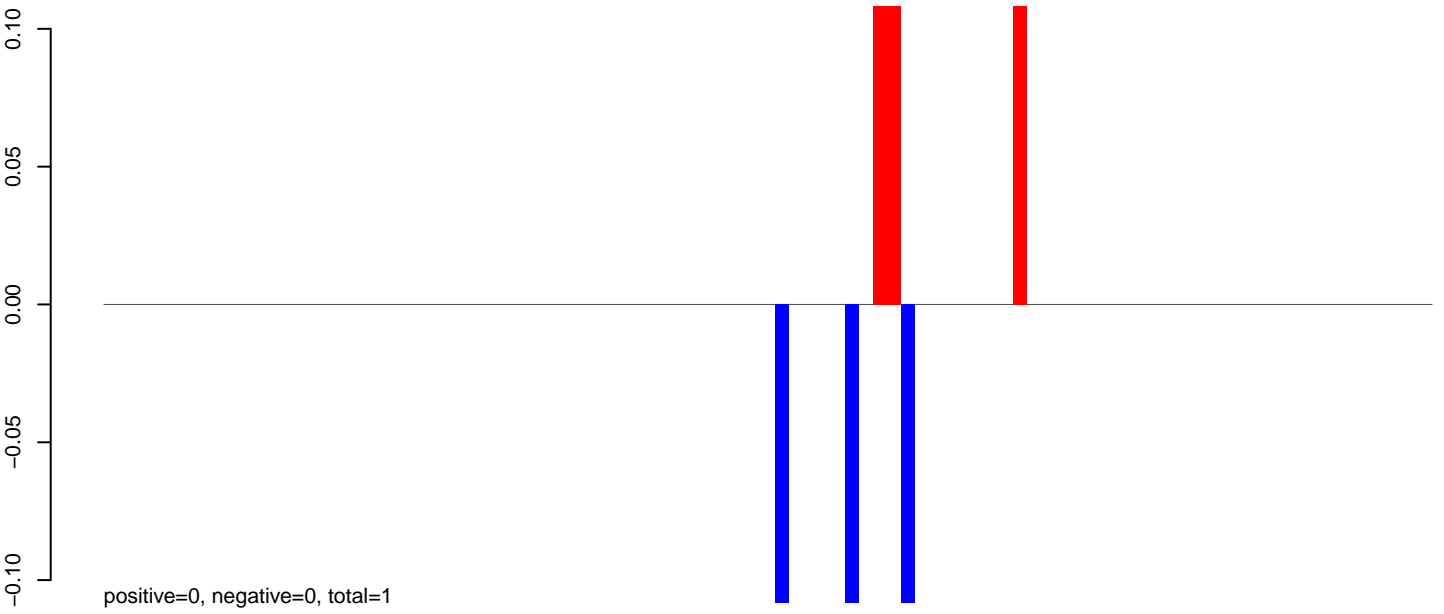
AnGam_Sua5b_cells.24_35.rep



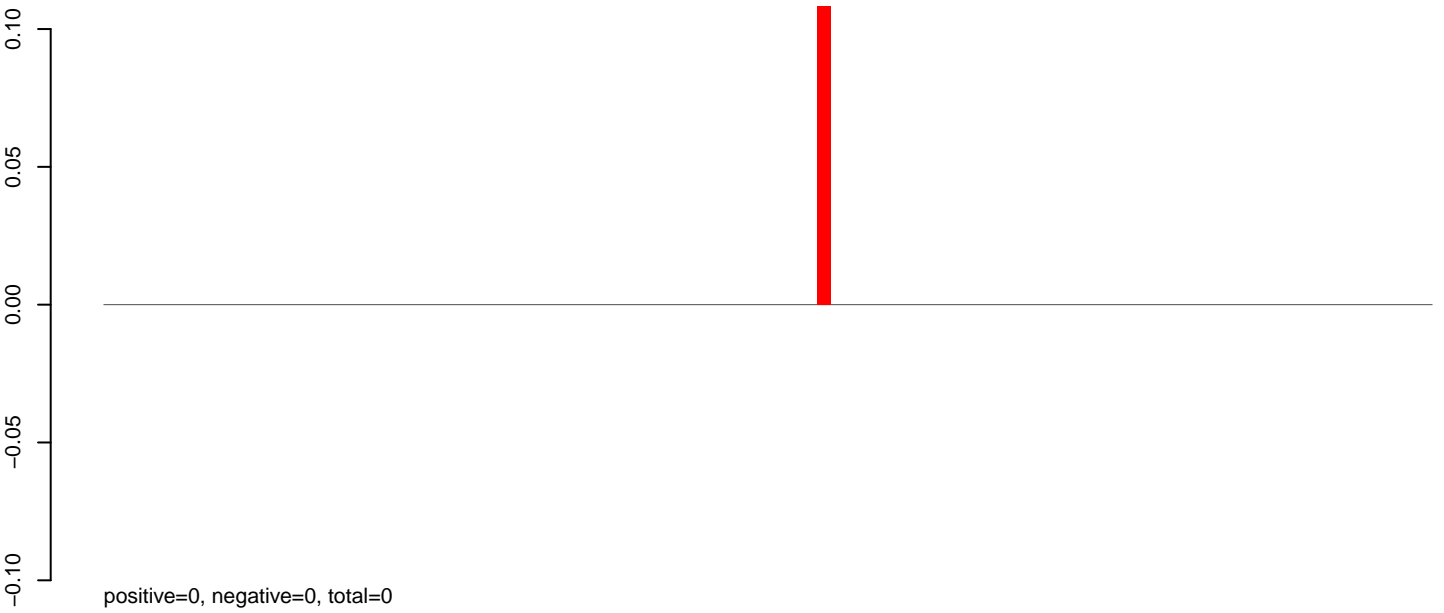
AnGam_Sua5b_cells.rep



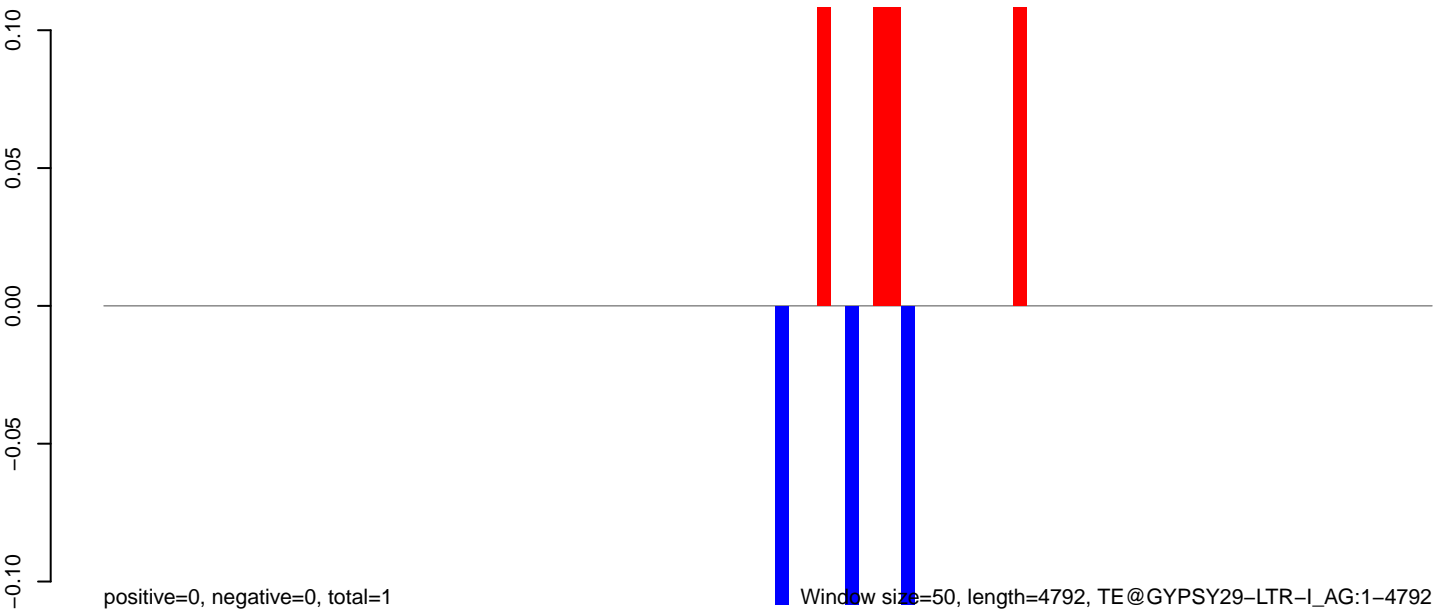
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



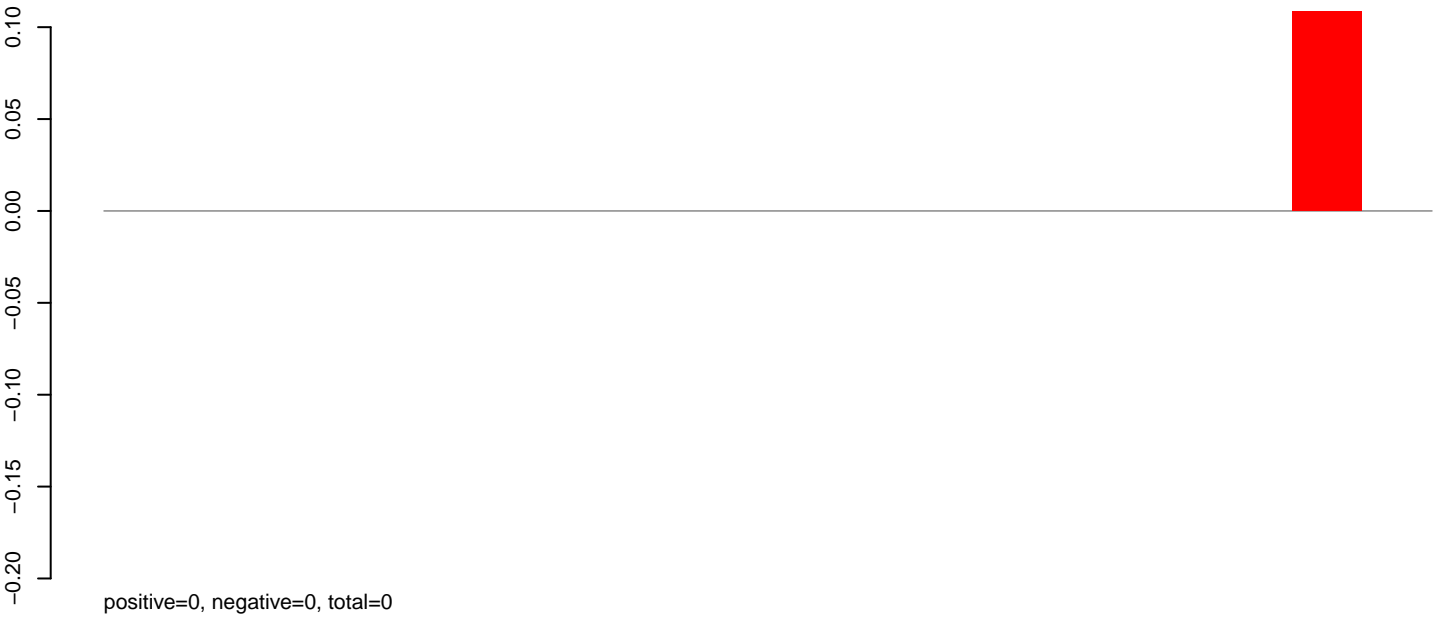
AnGam_Sua5b_cells.rep



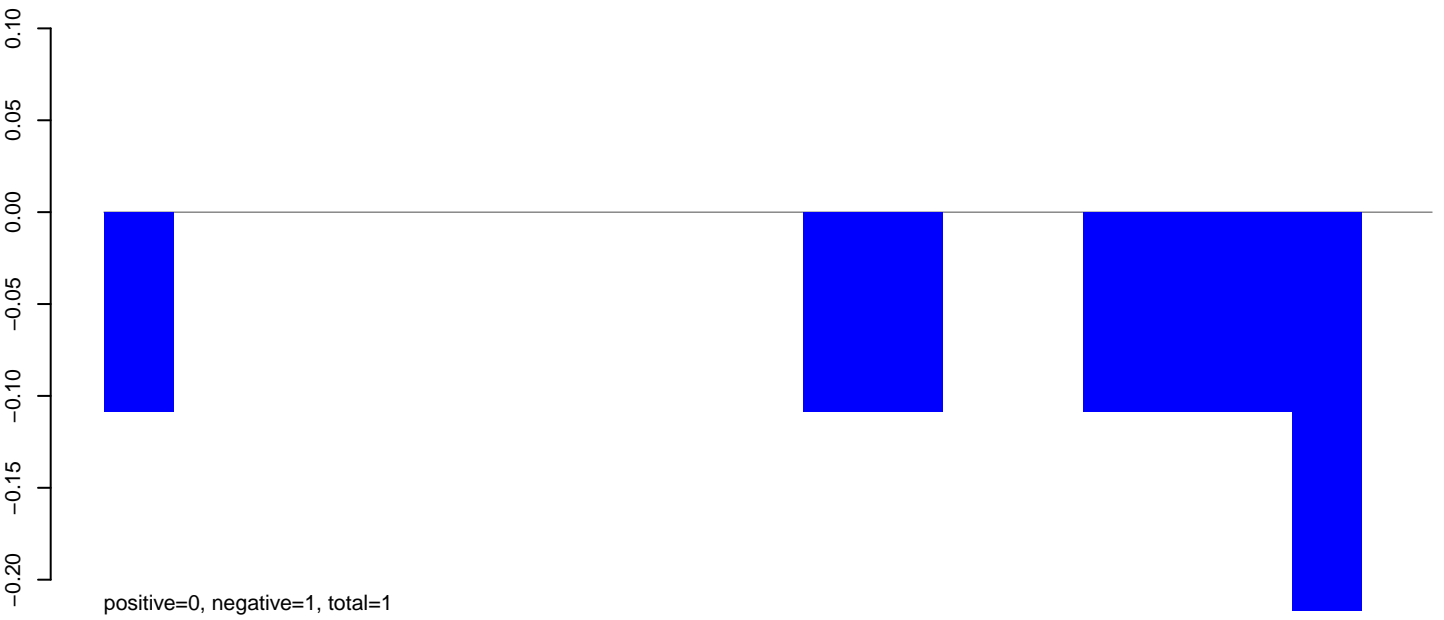
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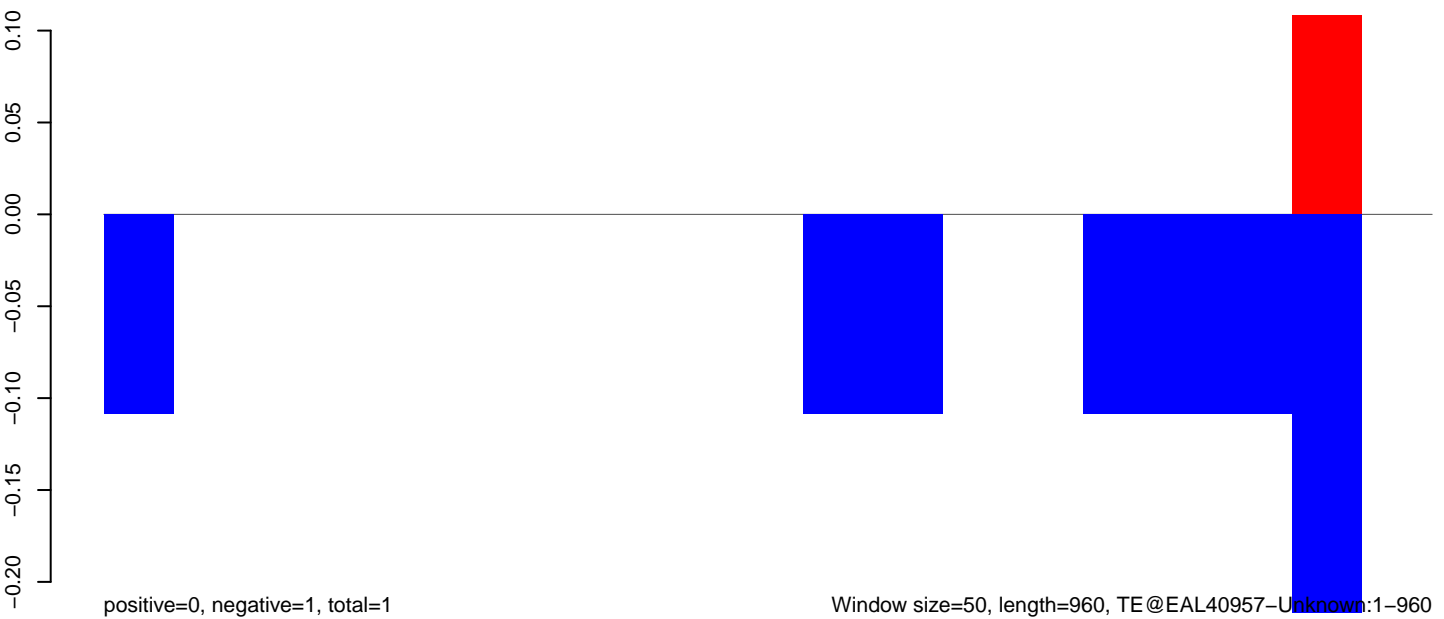
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



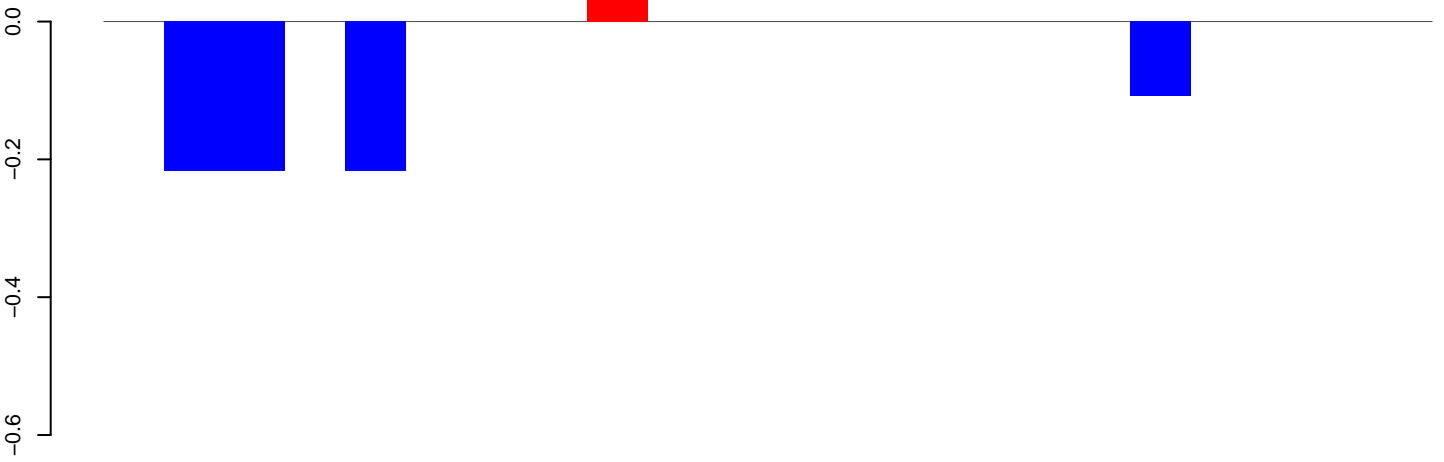
AnGam_Sua5b_cells.rep



Window size=50, length=960, TE@EAL40957-Unknown:1-960

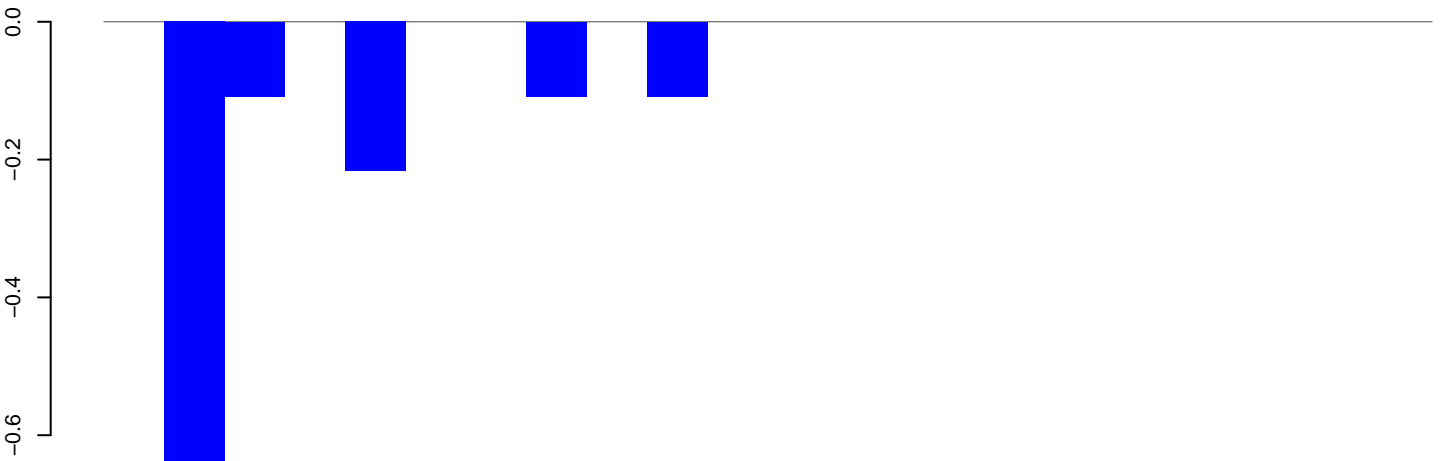
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AnGam_Sua5b_cells.18_23.rep



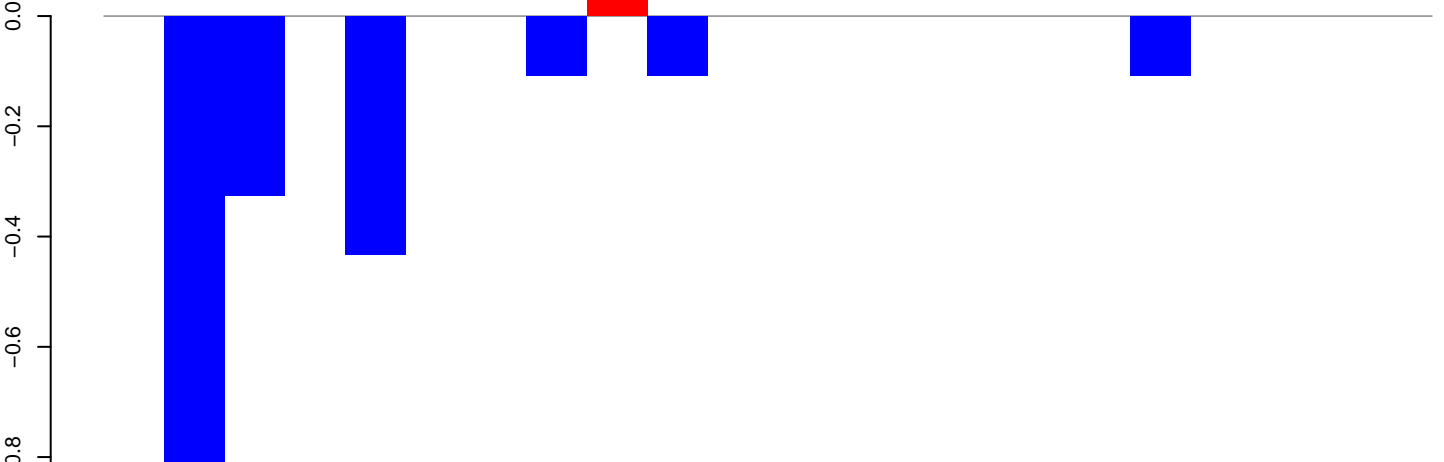
positive=0, negative=1, total=1

AnGam_Sua5b_cells.24_35.rep



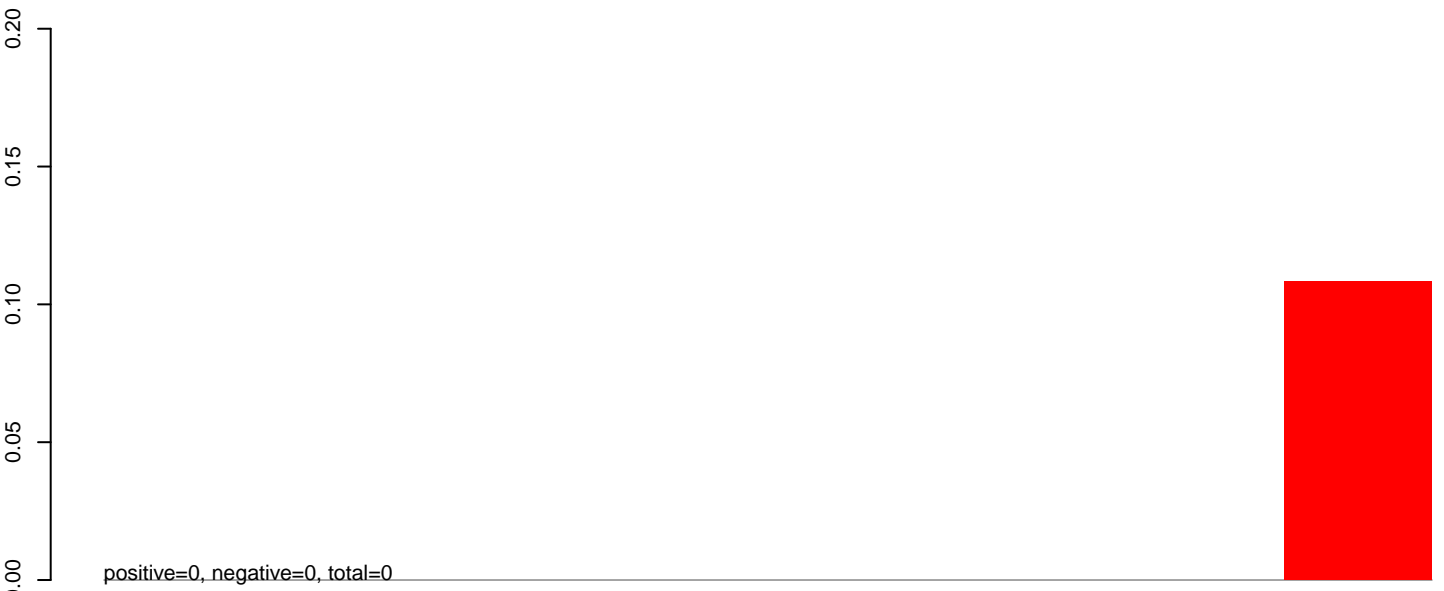
positive=0, negative=1, total=1

AnGam_Sua5b_cells.rep

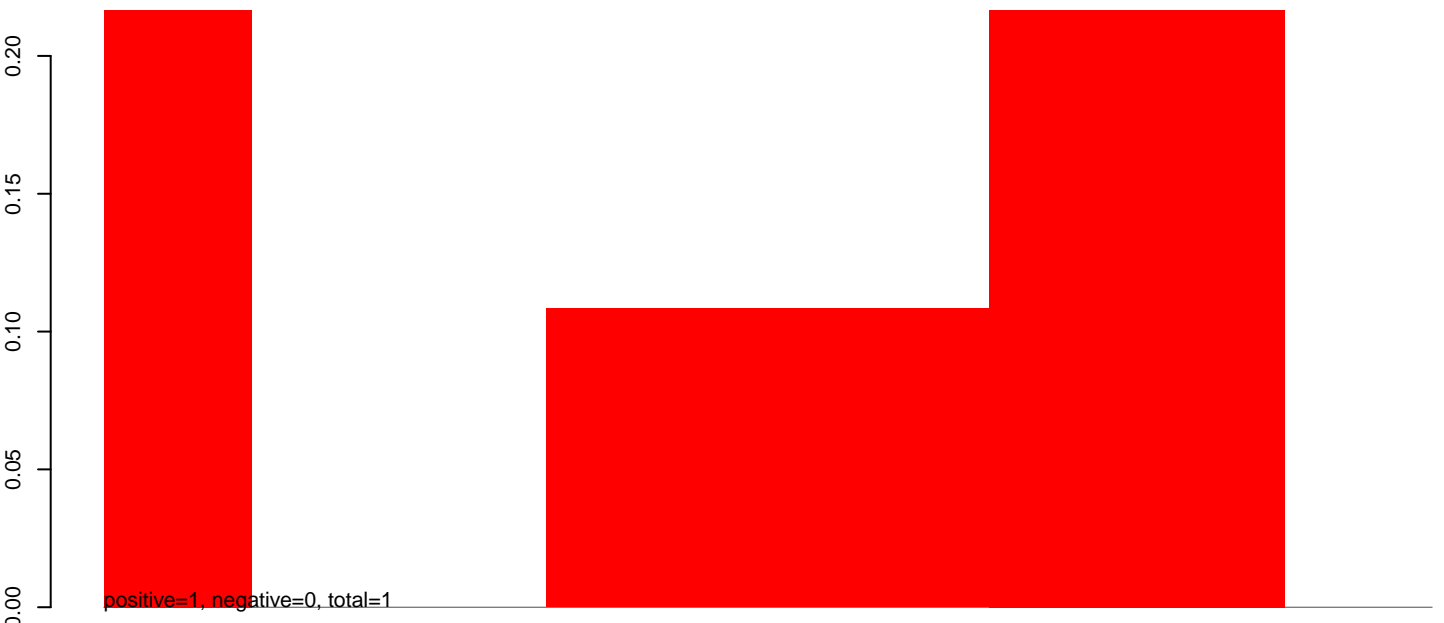


positive=0, negative=2, total=2

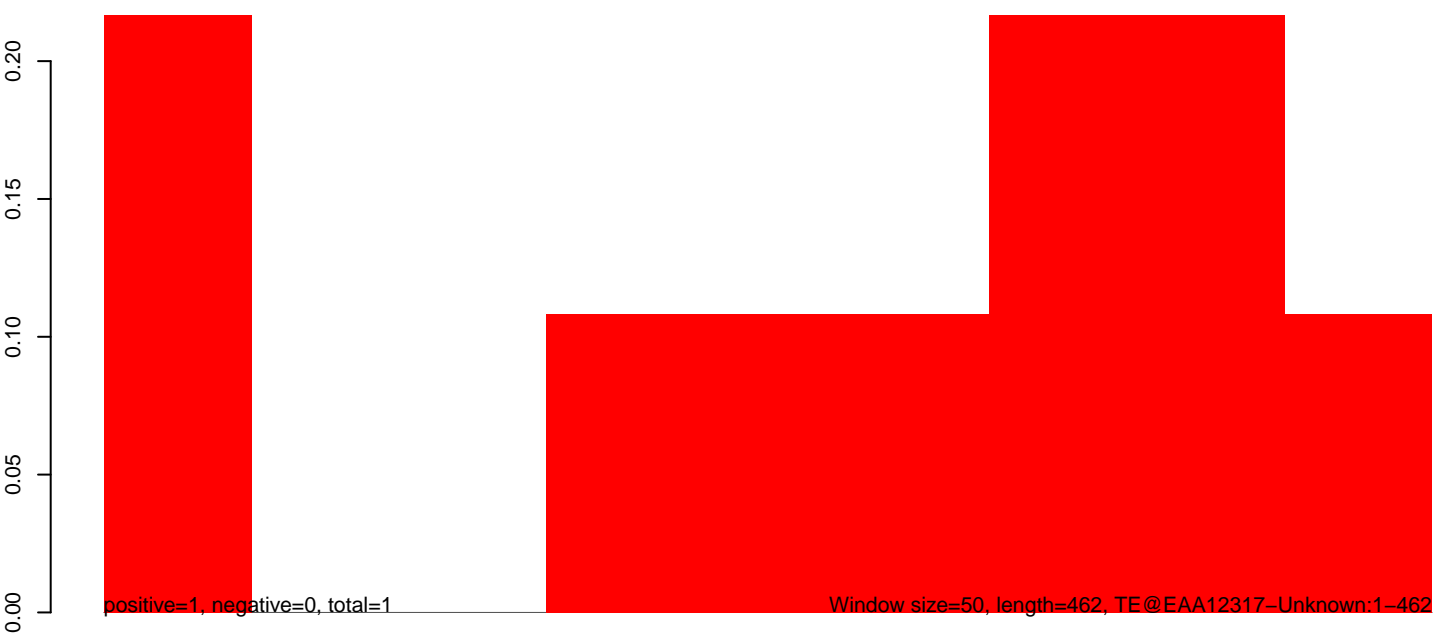
AnGam_Sua5b_cells.18_23.rep



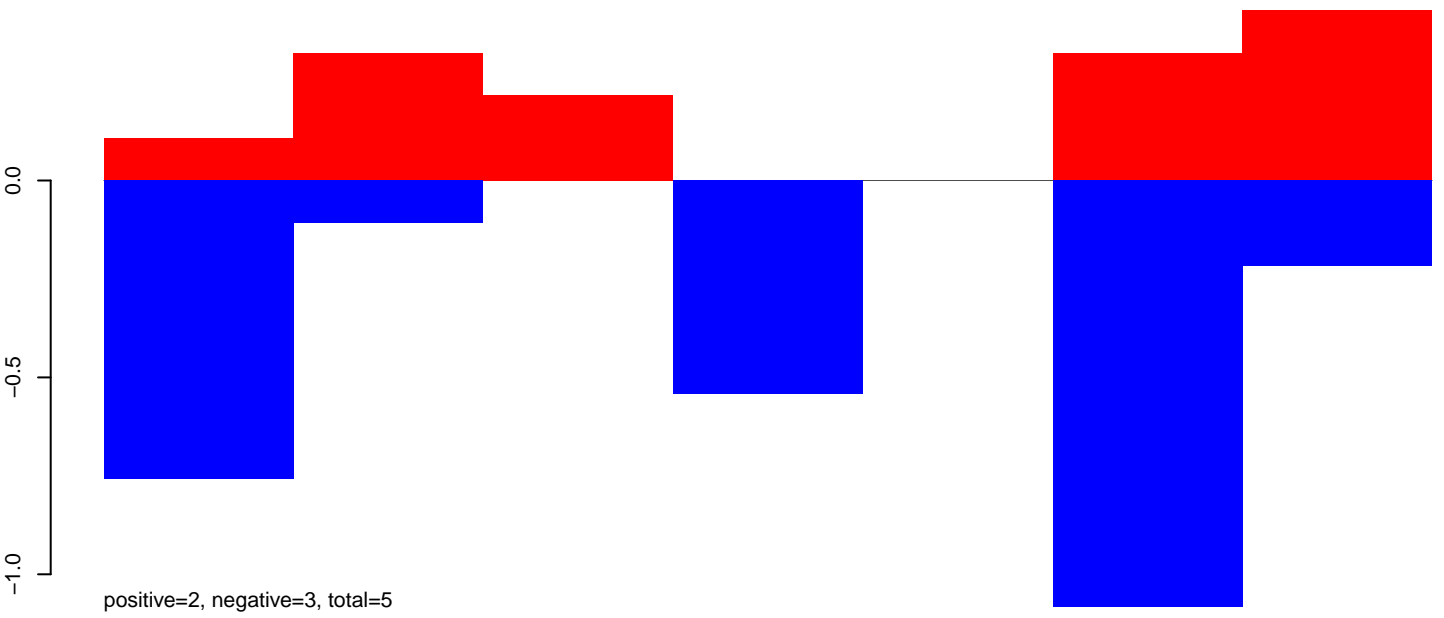
AnGam_Sua5b_cells.24_35.rep



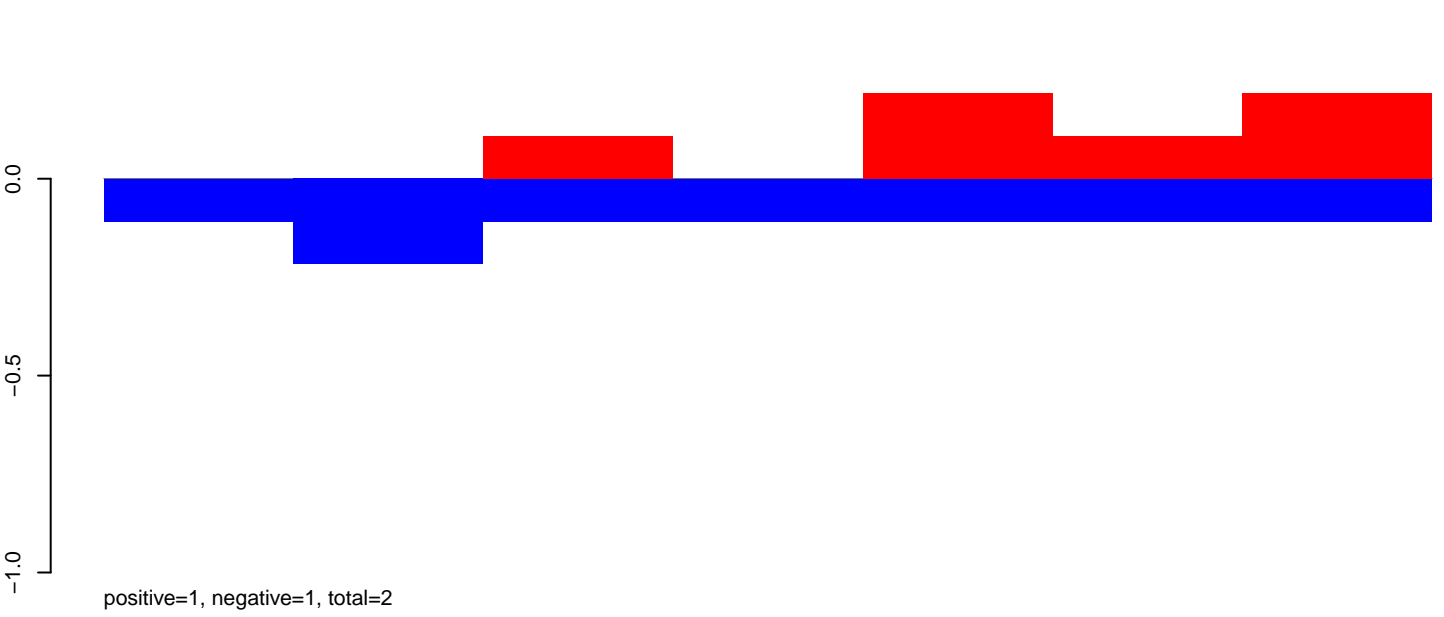
AnGam_Sua5b_cells.rep



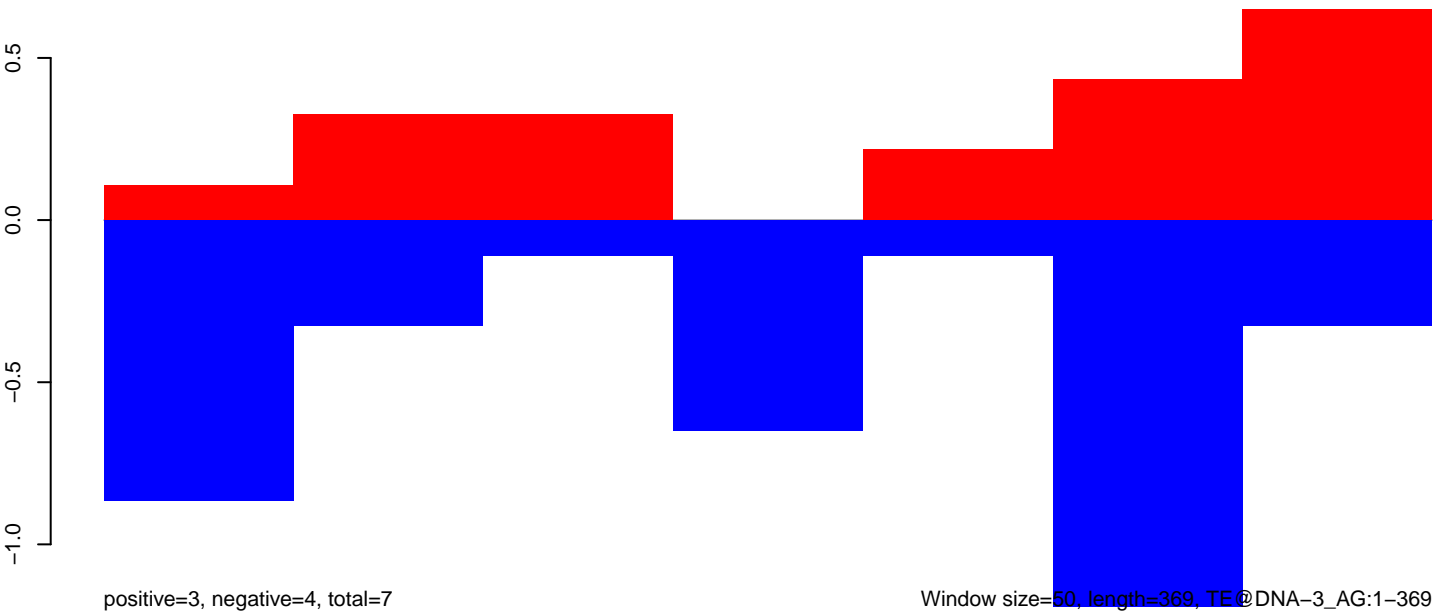
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

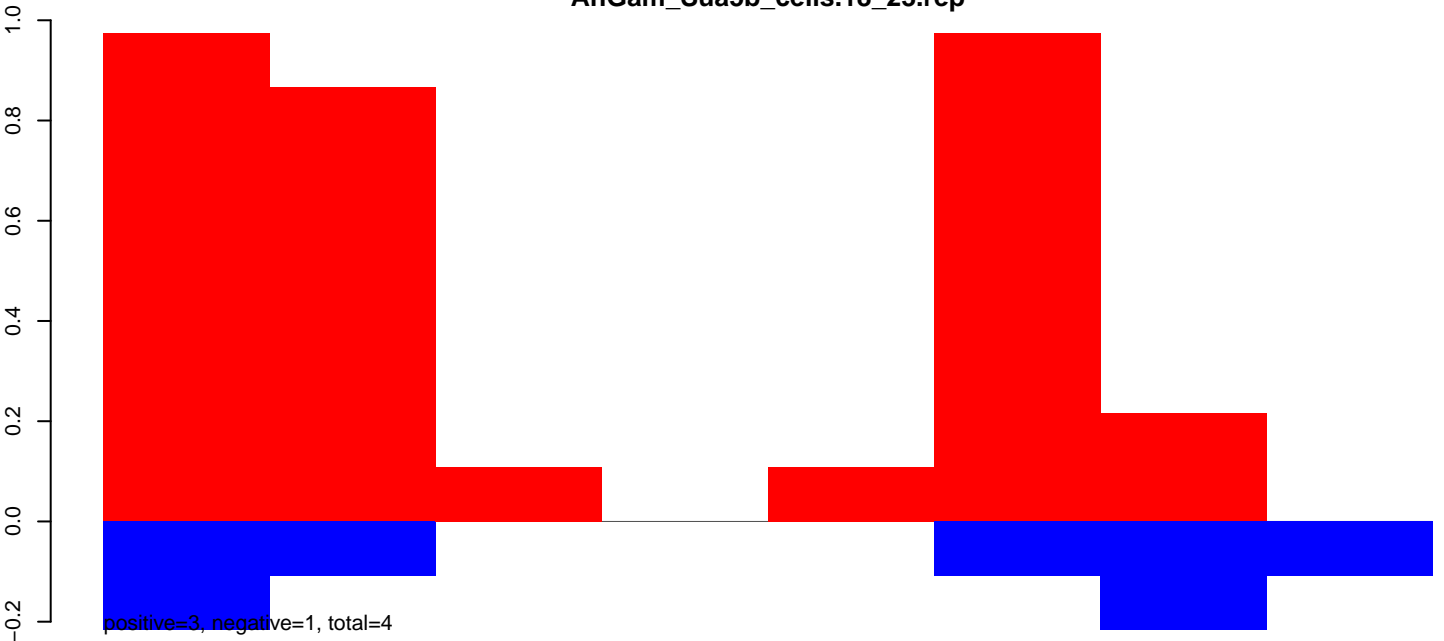


AnGam_Sua5b_cells.rep

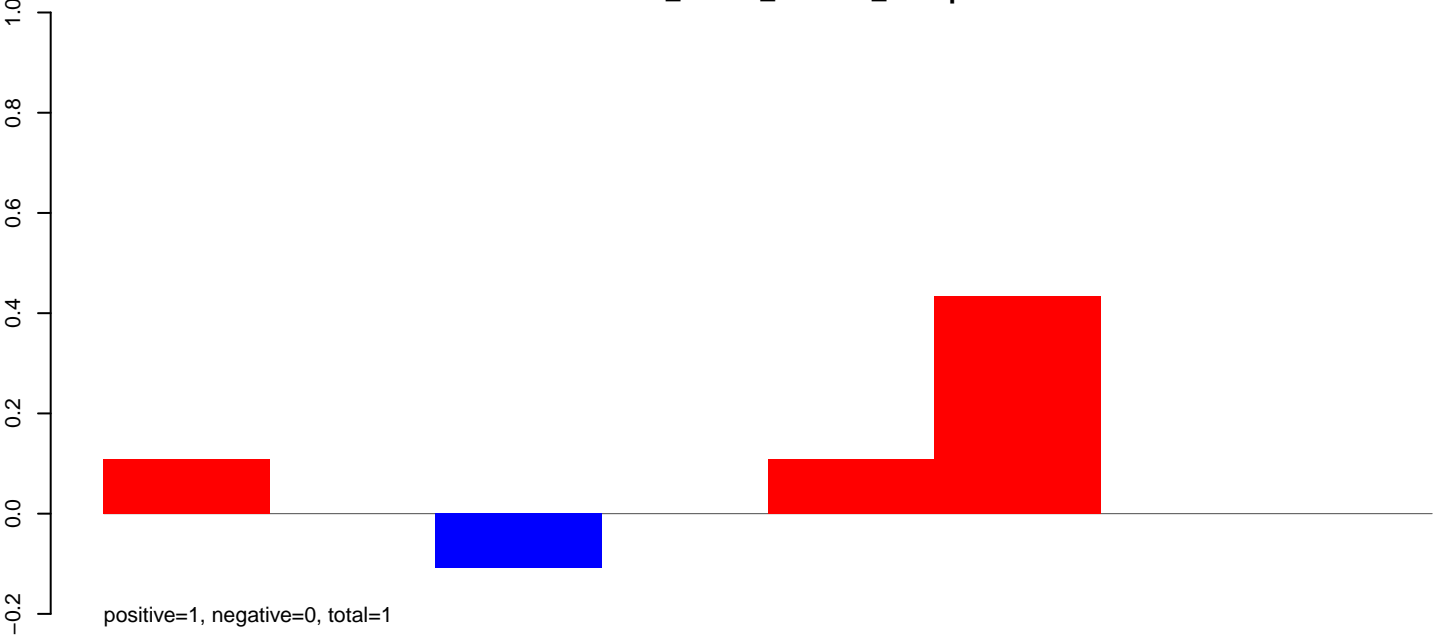


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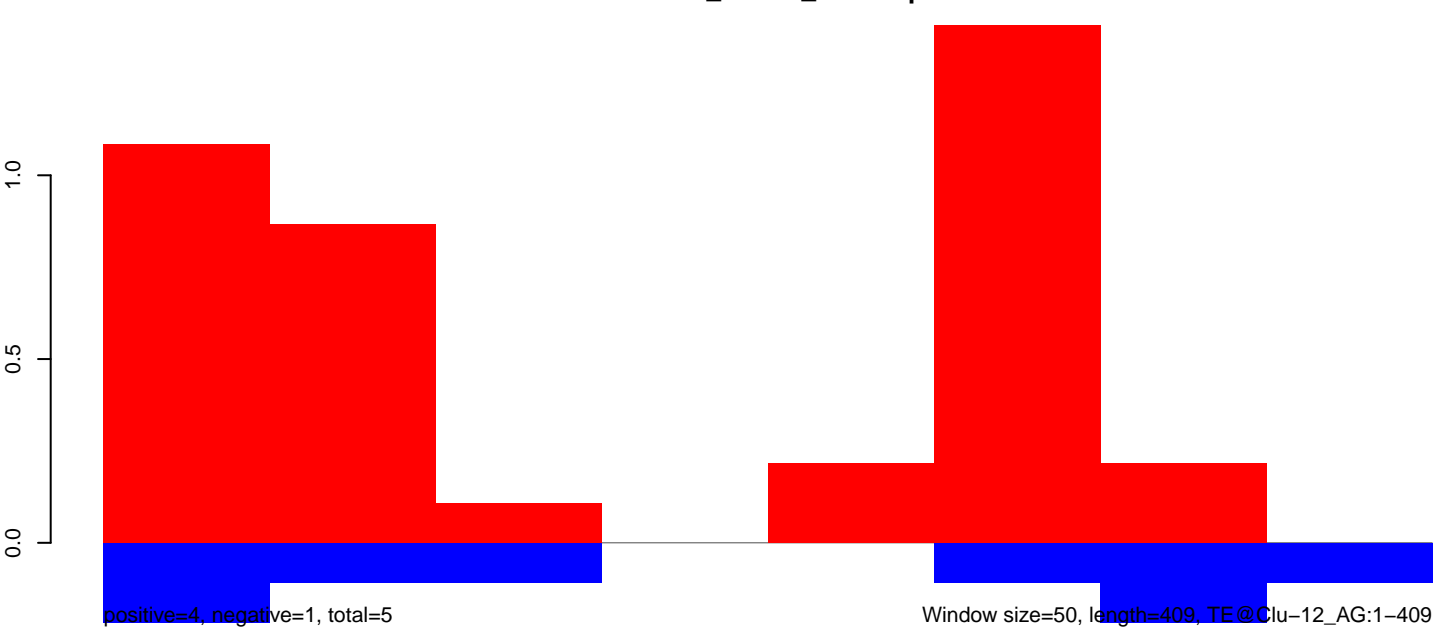
AnGam_Sua5b_cells.18_23.rep



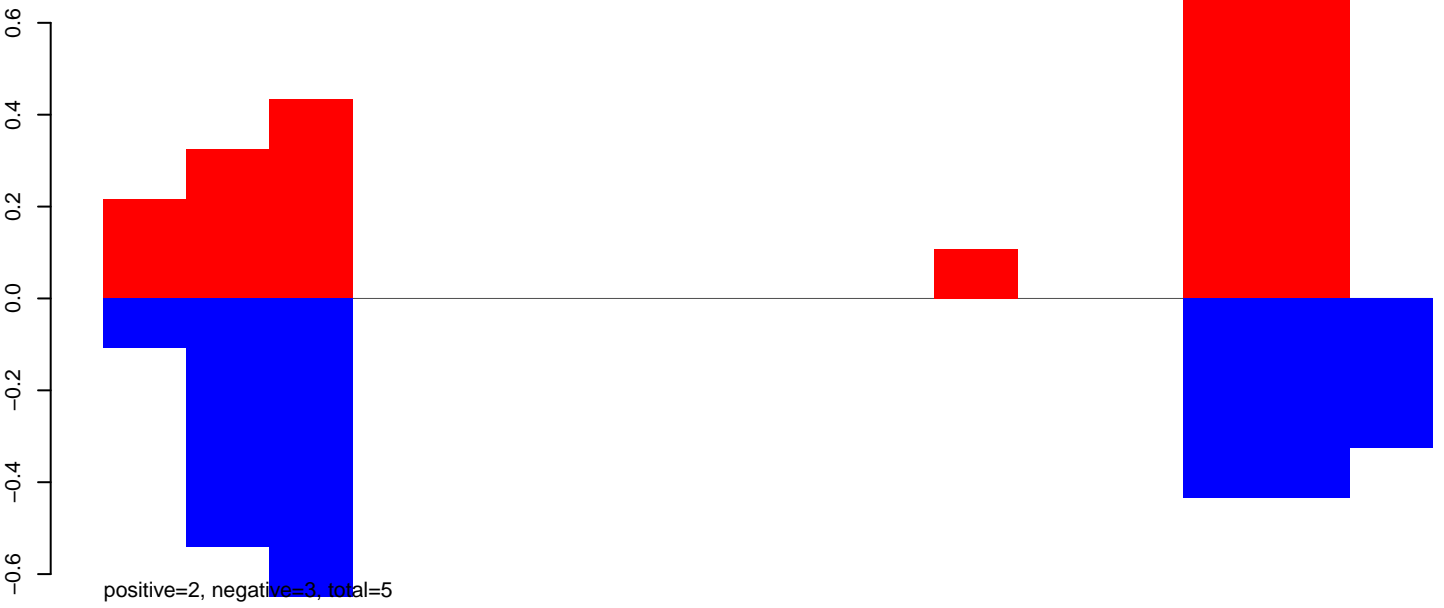
AnGam_Sua5b_cells.24_35.rep



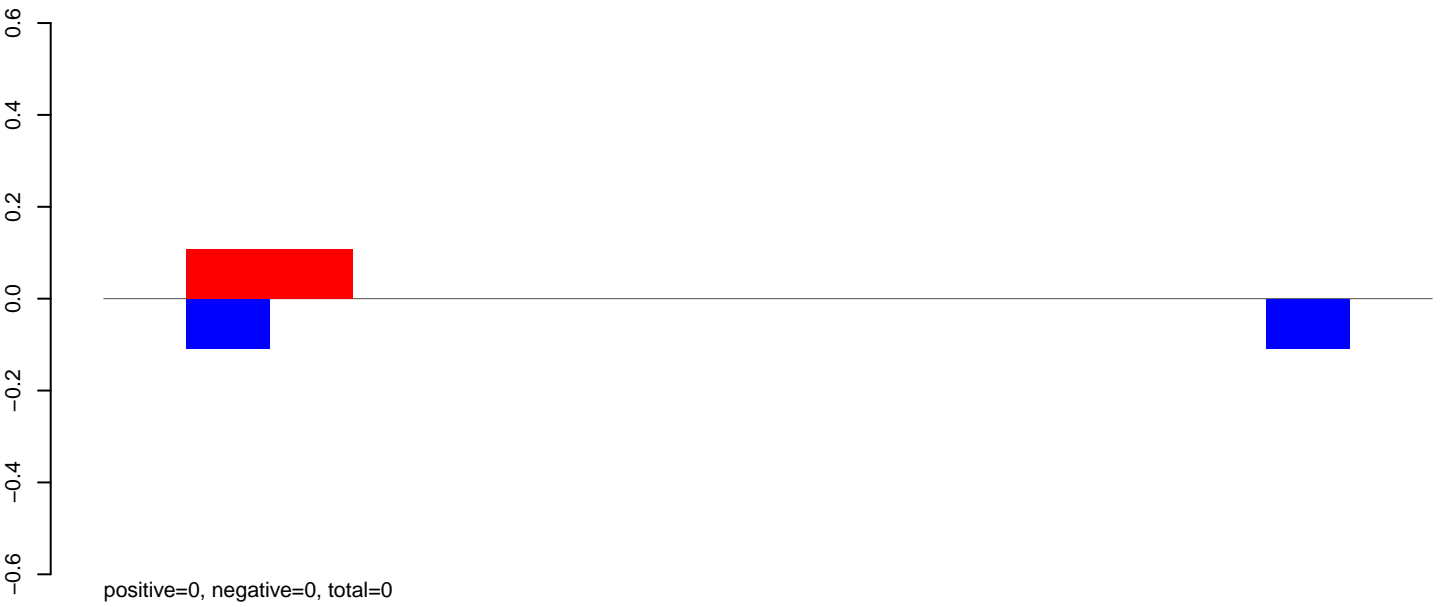
AnGam_Sua5b_cells.rep



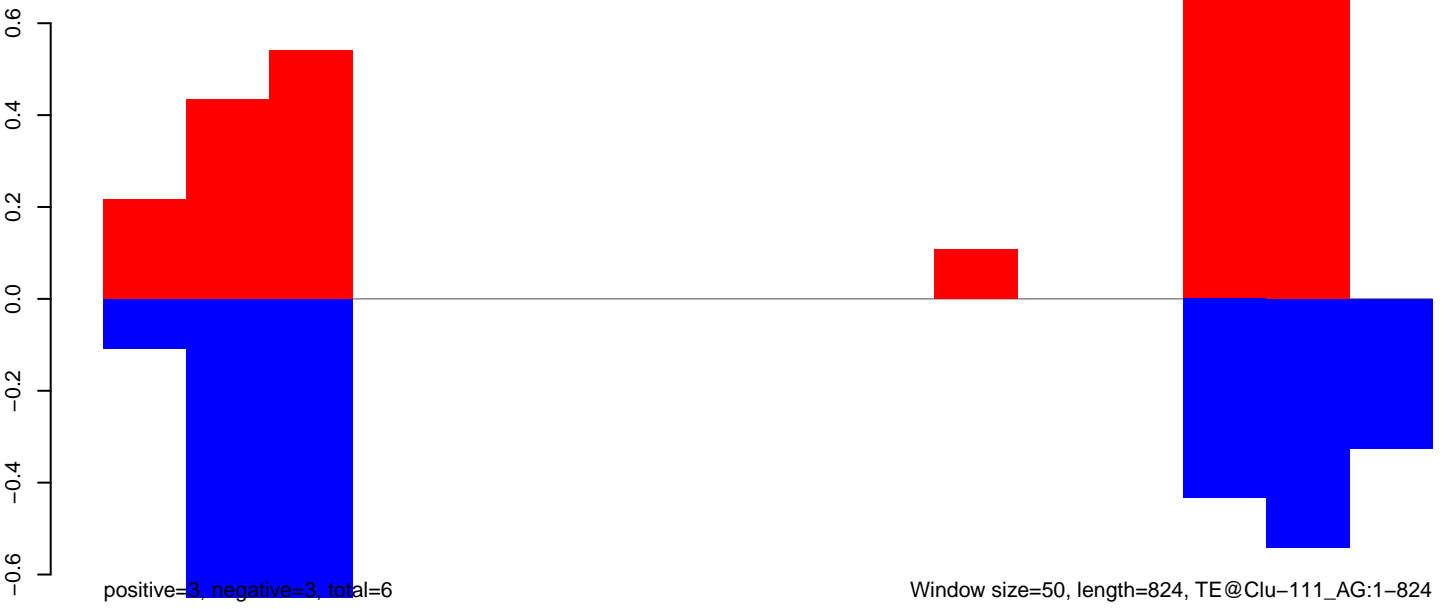
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



Window size=50, length=824, TE@Clu-111_AG:1-824

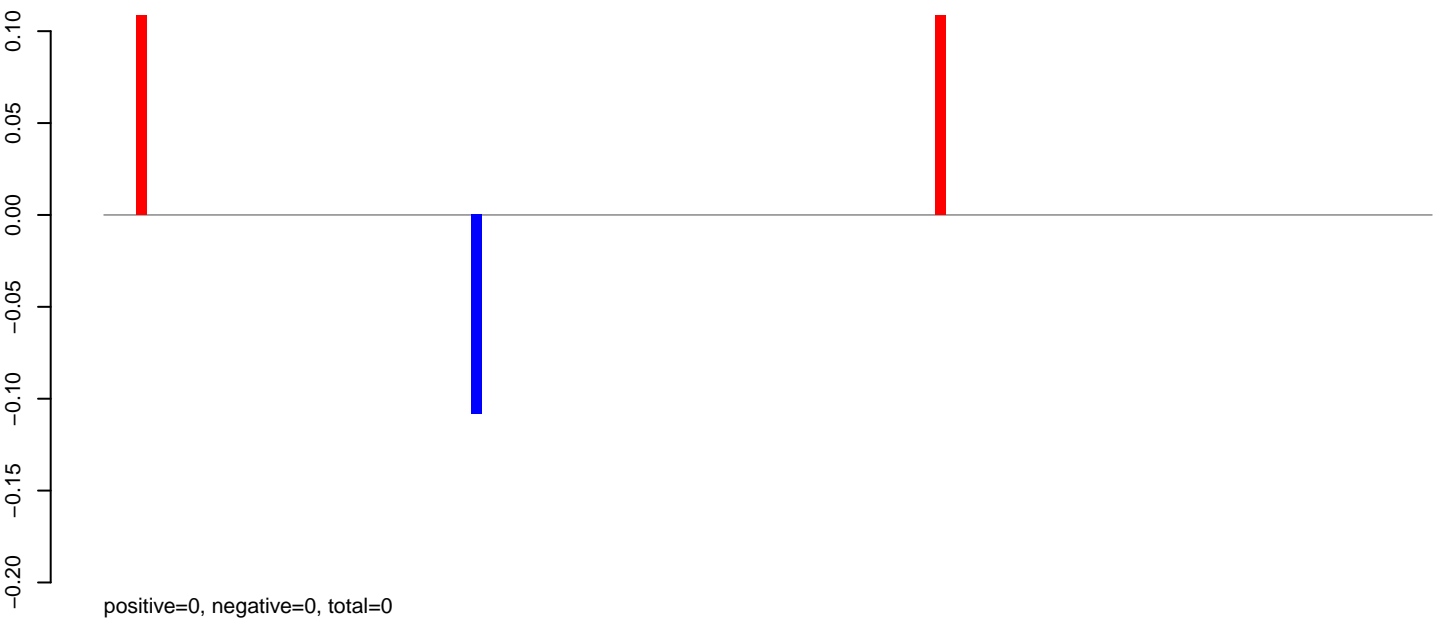
200

400

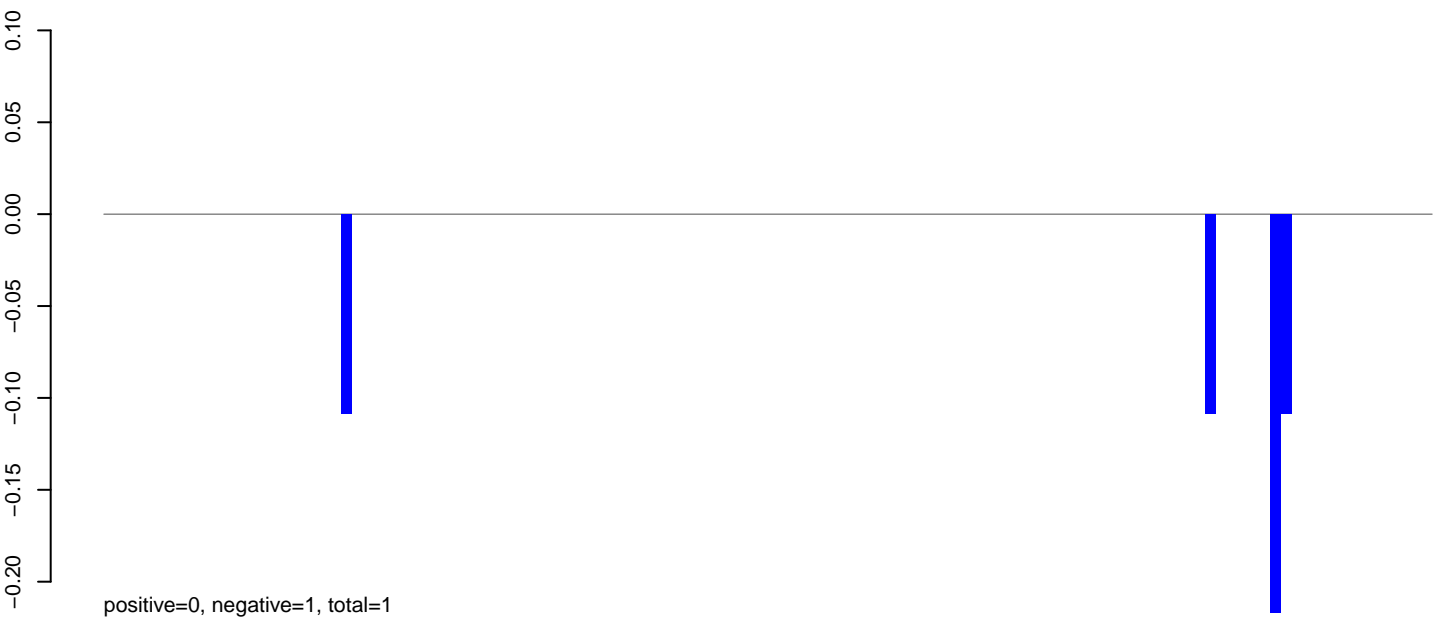
600

800

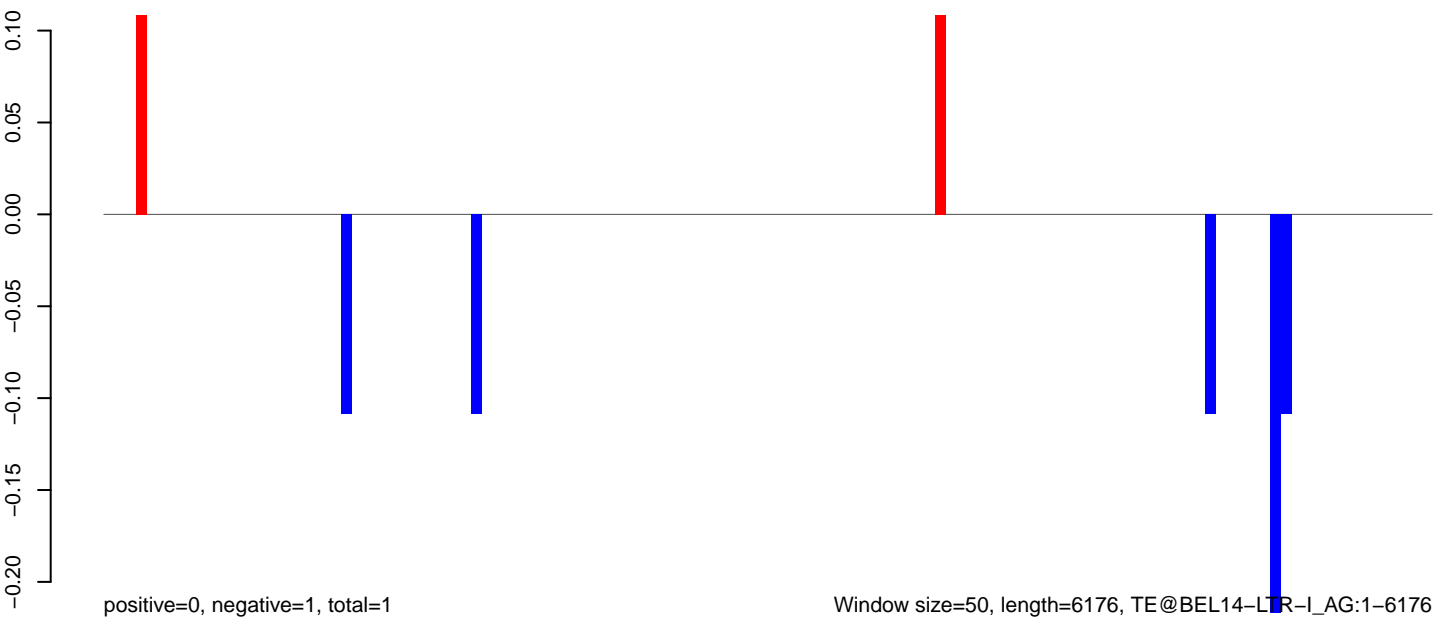
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

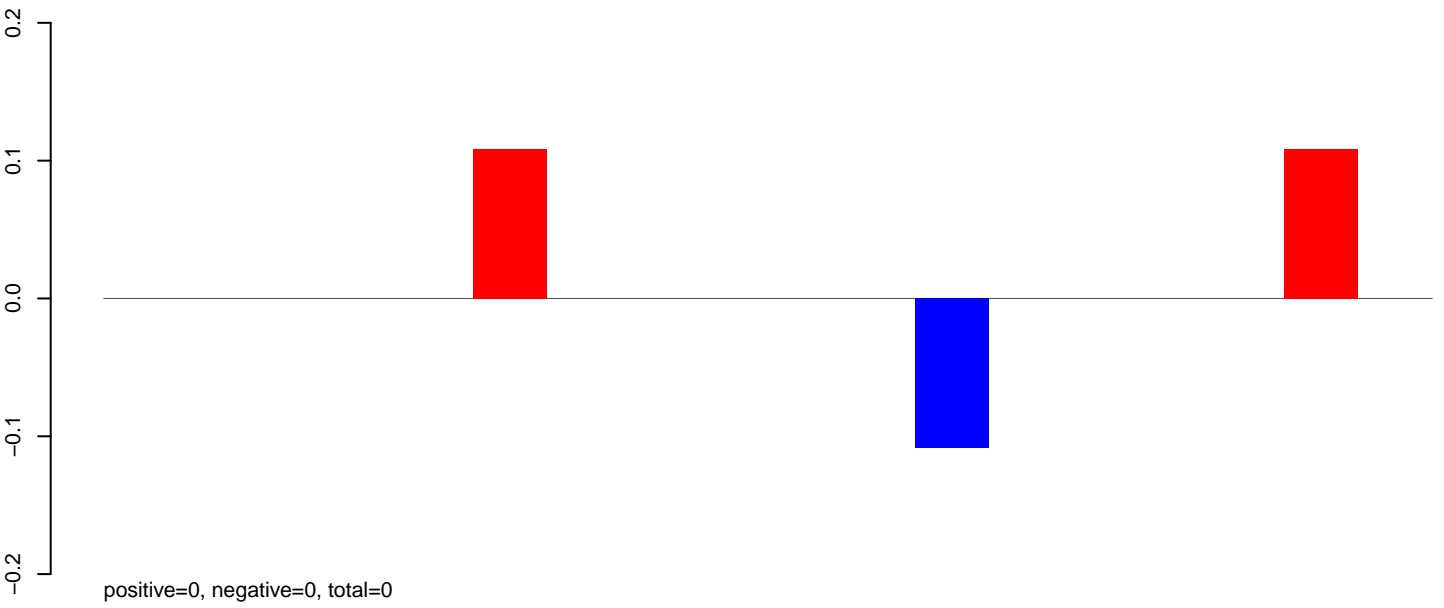


AnGam_Sua5b_cells.rep

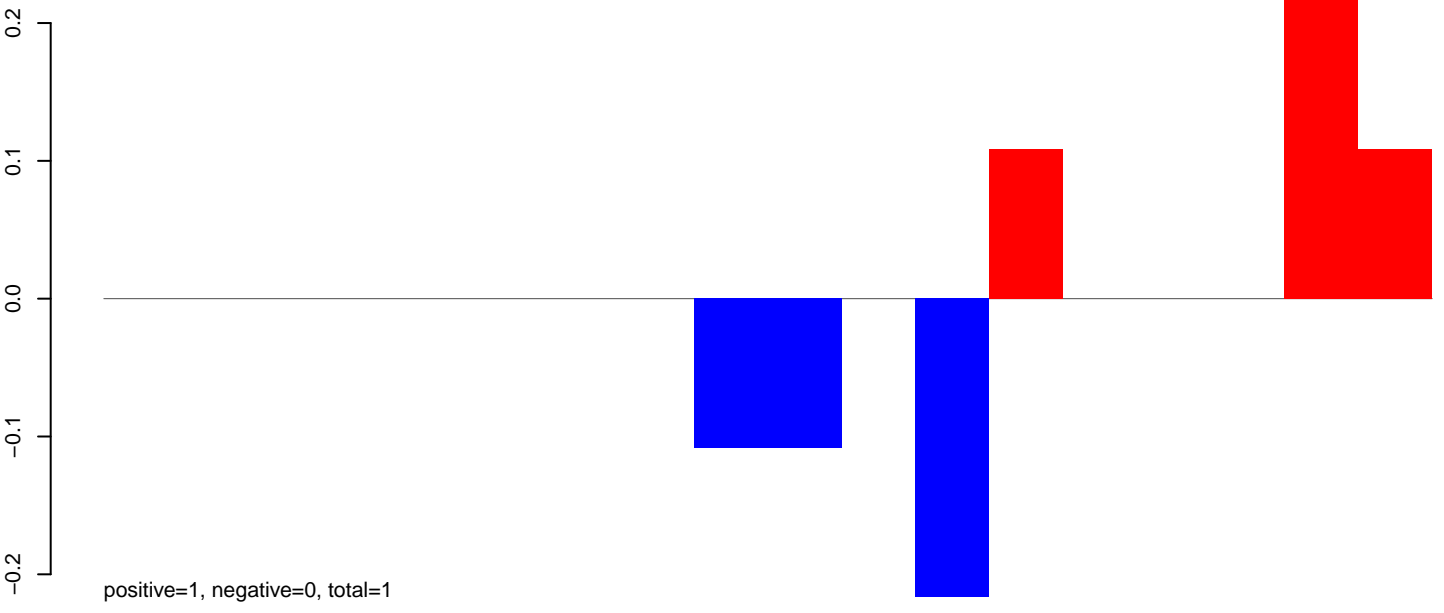


Window size=50, length=6176, TE@BEL14-LTR-I_AG:1-6176

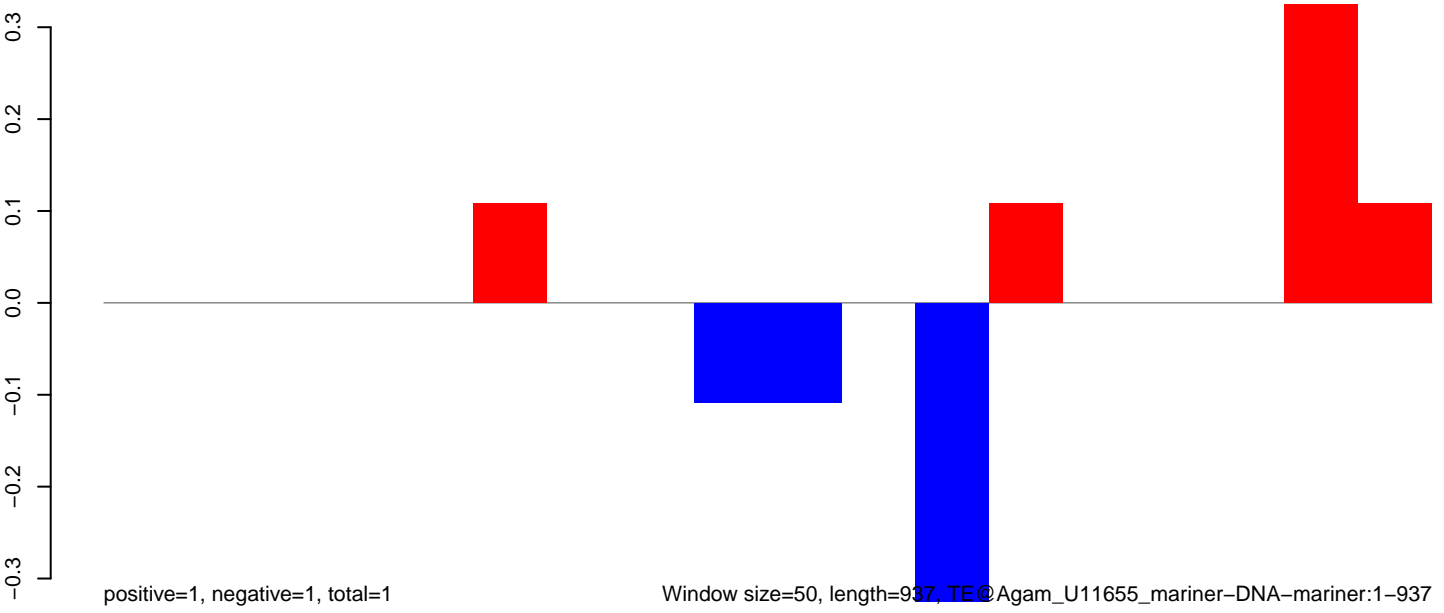
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



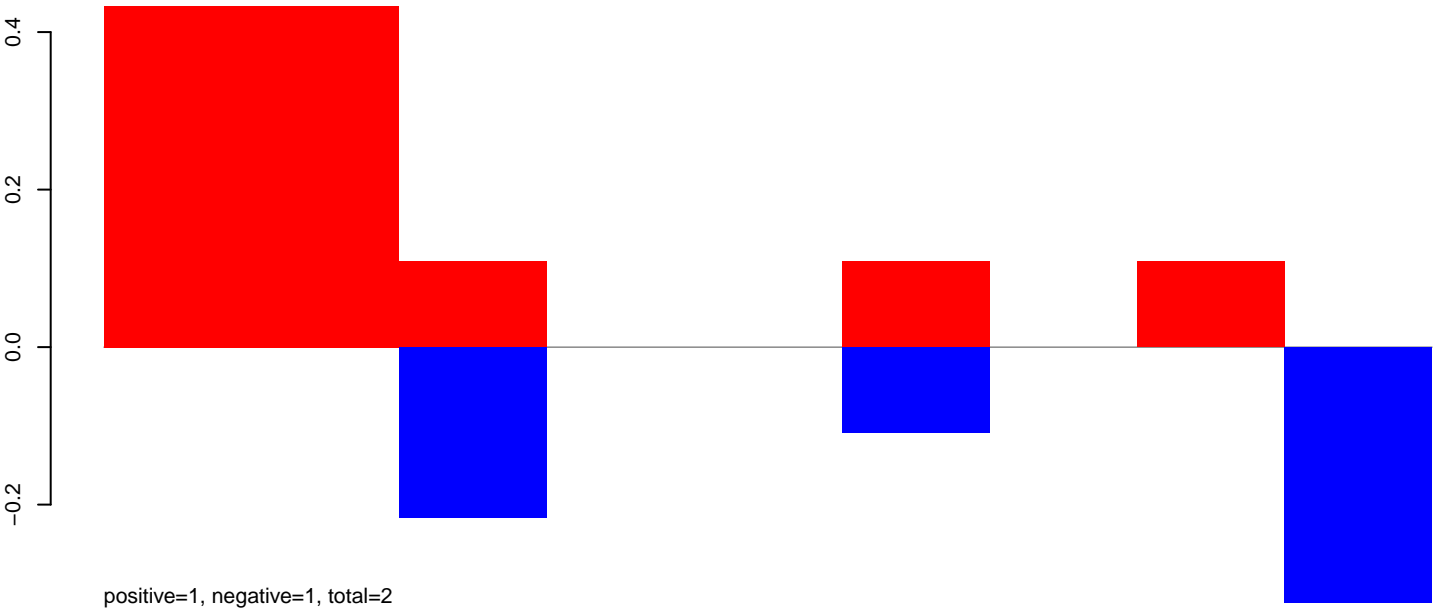
AnGam_Sua5b_cells.rep



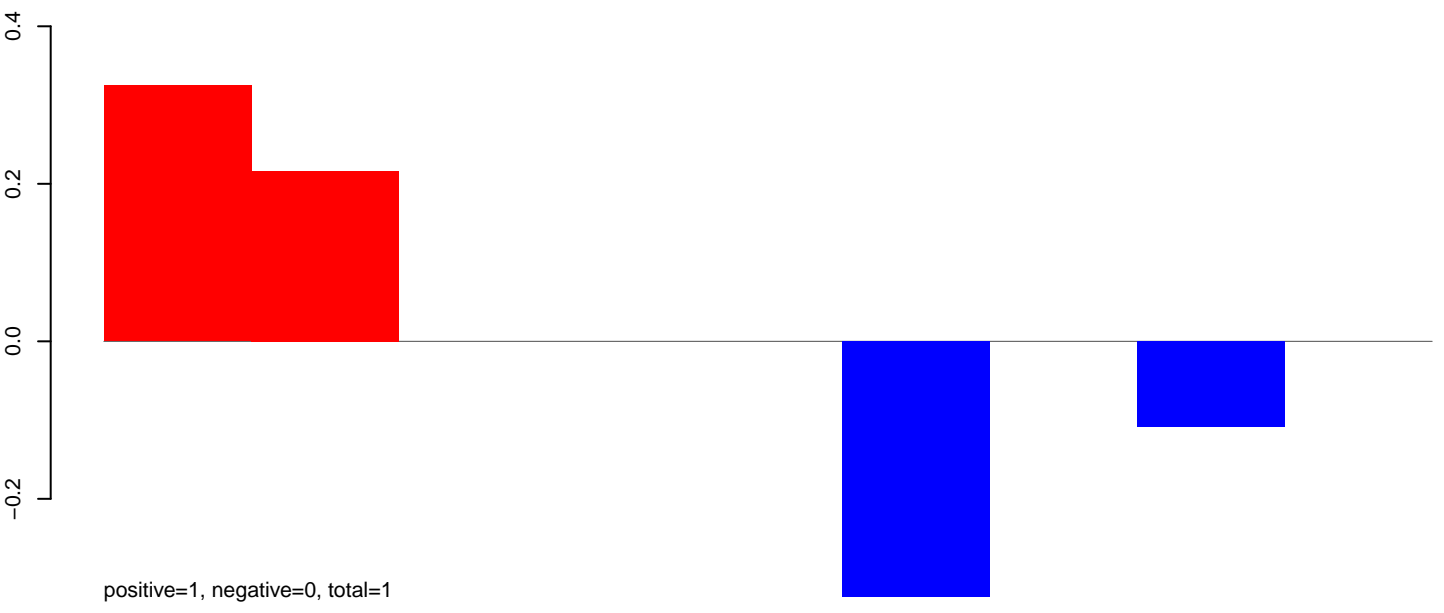
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200 400 600 800

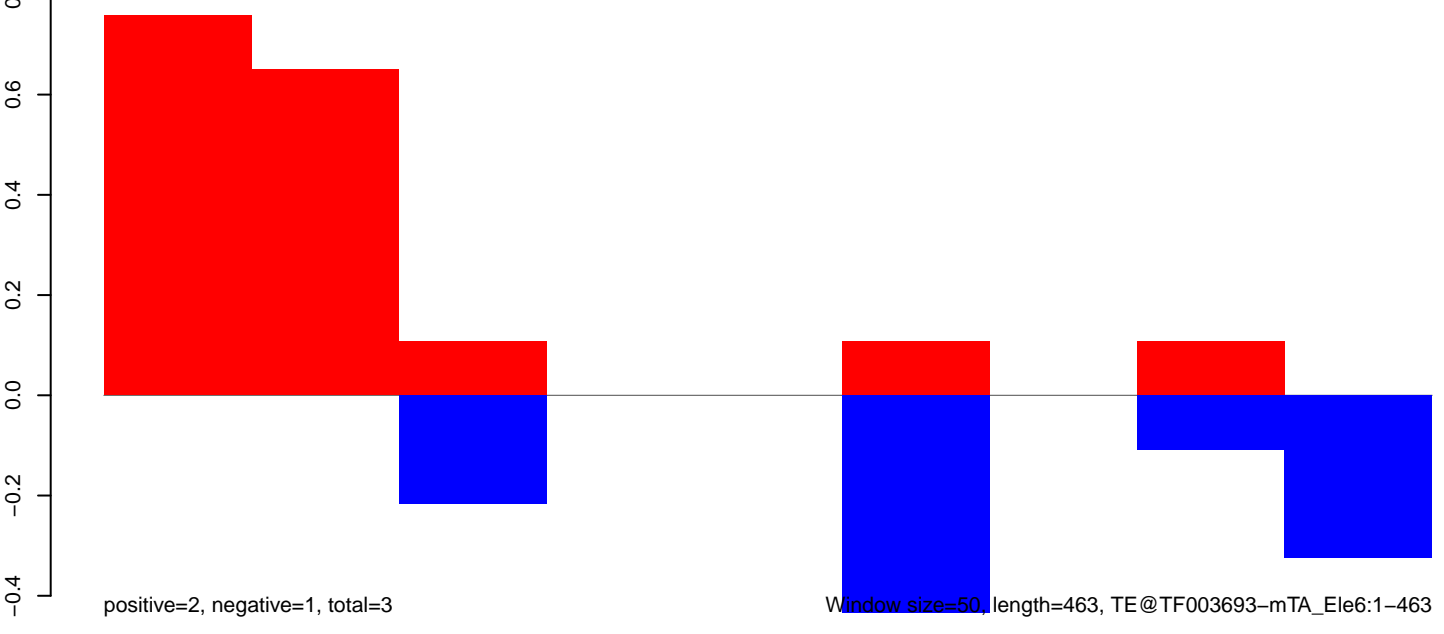
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

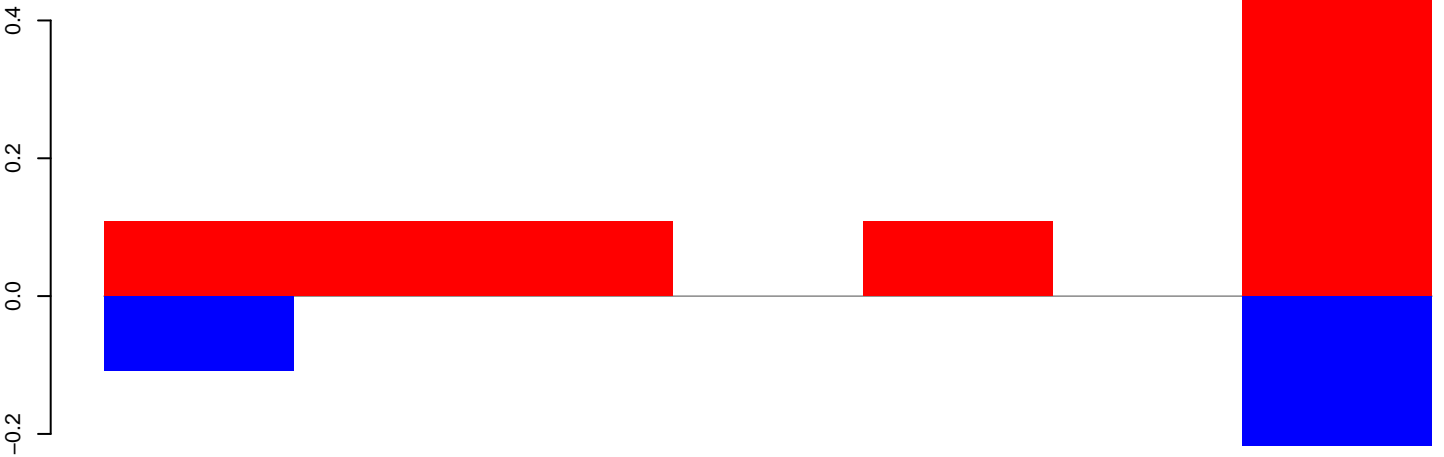


AnGam_Sua5b_cells.rep



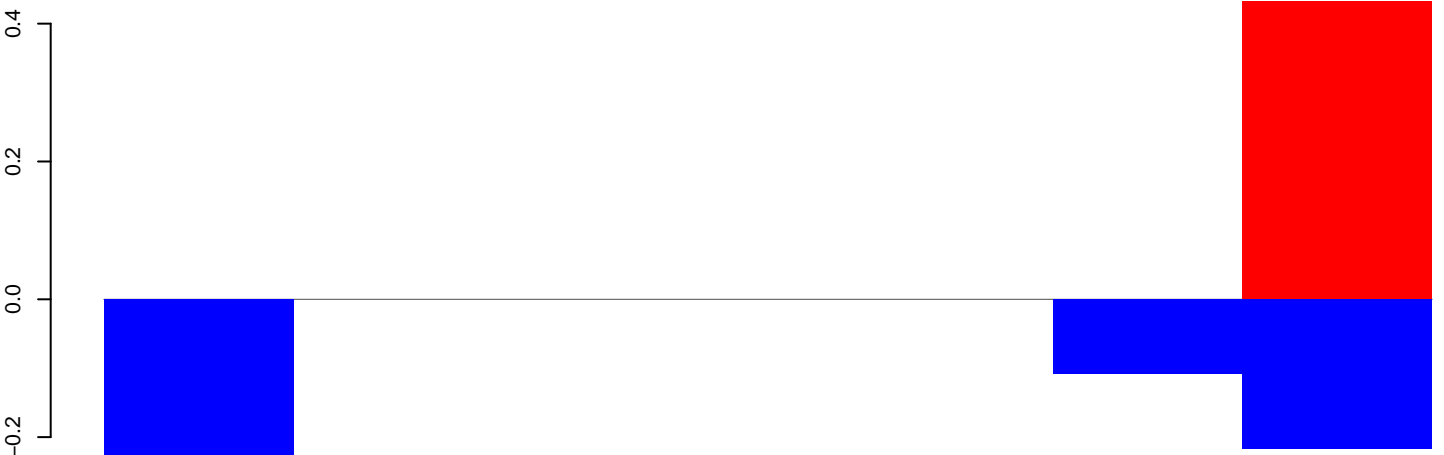
Window size=50, length=463, TE@TF003693-mTA_Ele6:1-463

AnGam_Sua5b_cells.18_23.rep



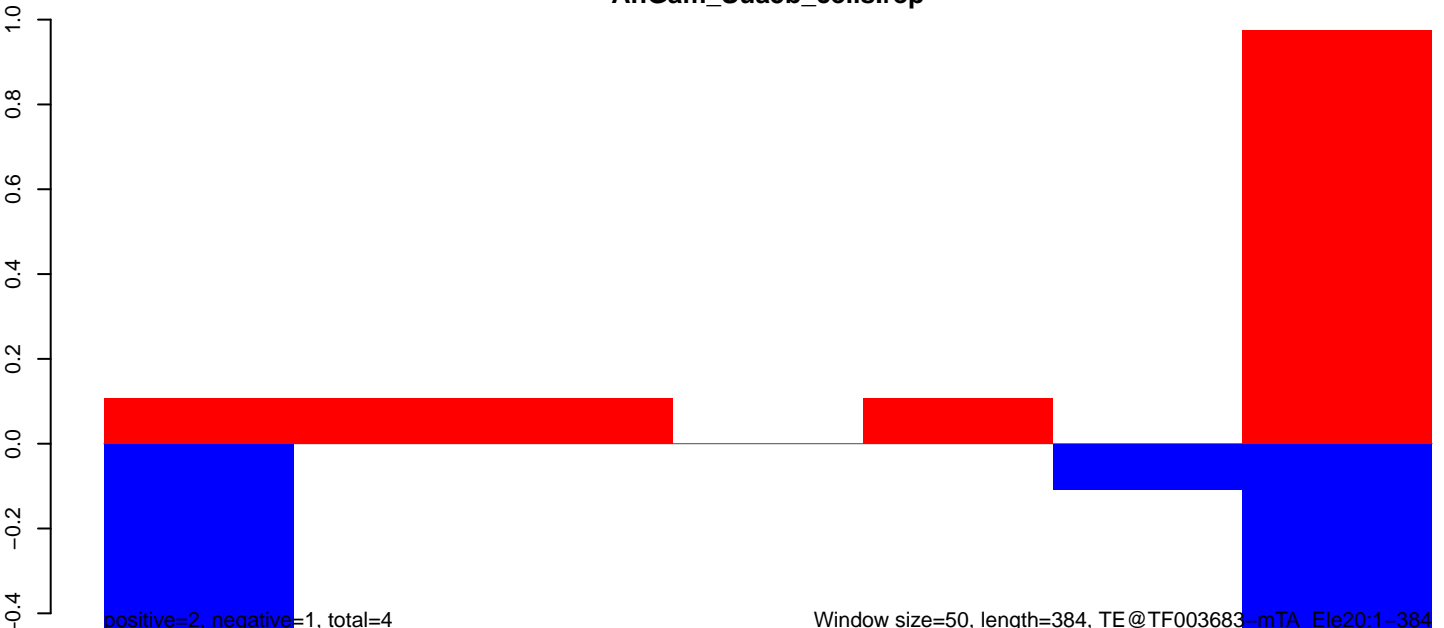
positive=2, negative=1, total=2

AnGam_Sua5b_cells.24_35.rep



positive=1, negative=1, total=2

AnGam_Sua5b_cells.rep

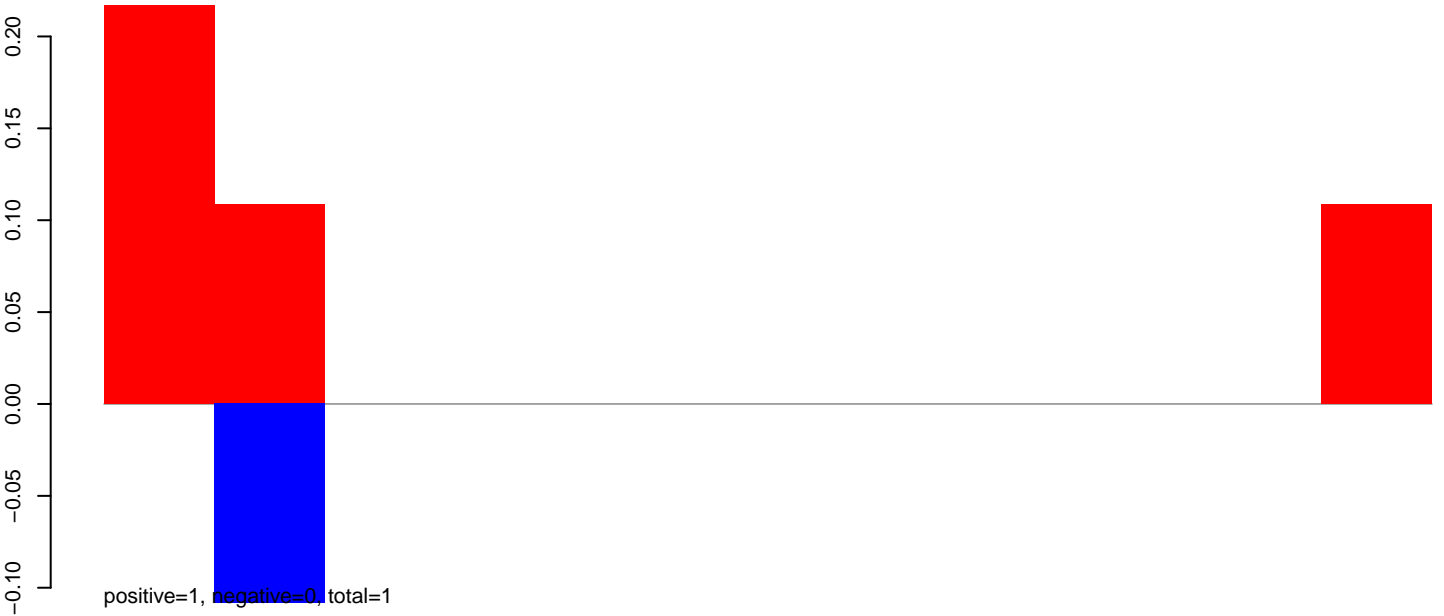


positive=2, negative=1, total=4

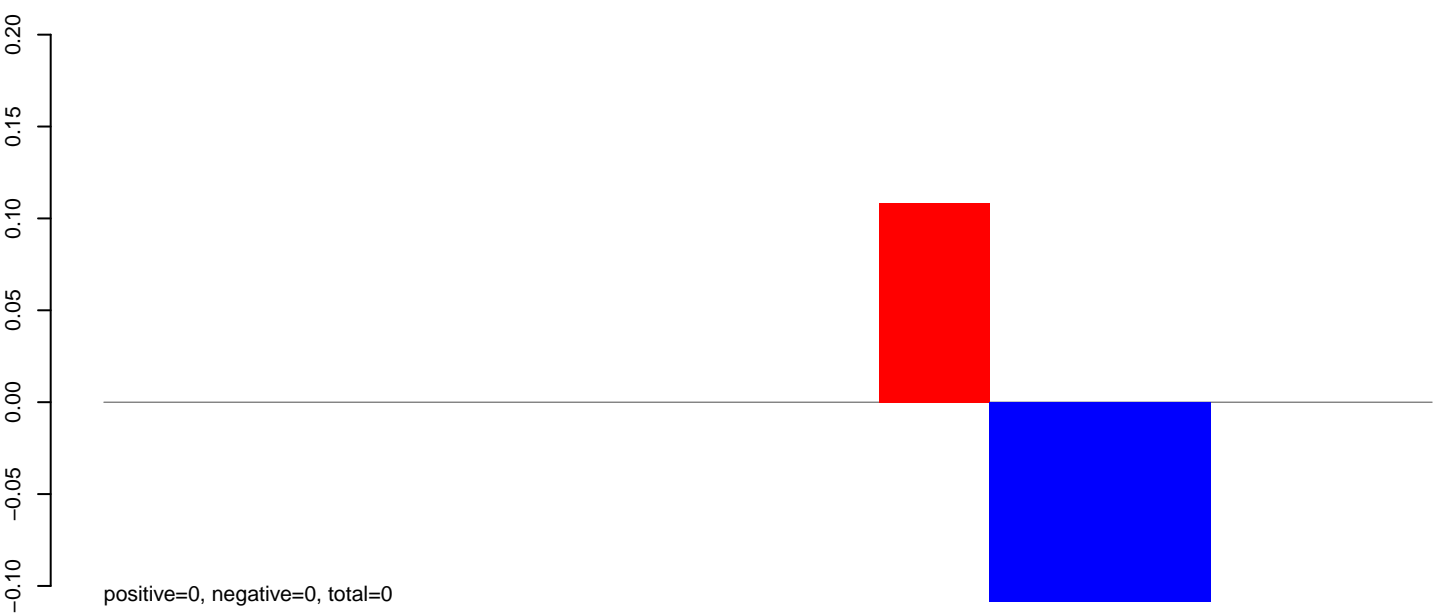
Window size=50, length=384, TE@TF003683-mTA_Ele20:1-384

50 100 150 200 250 300 350 400

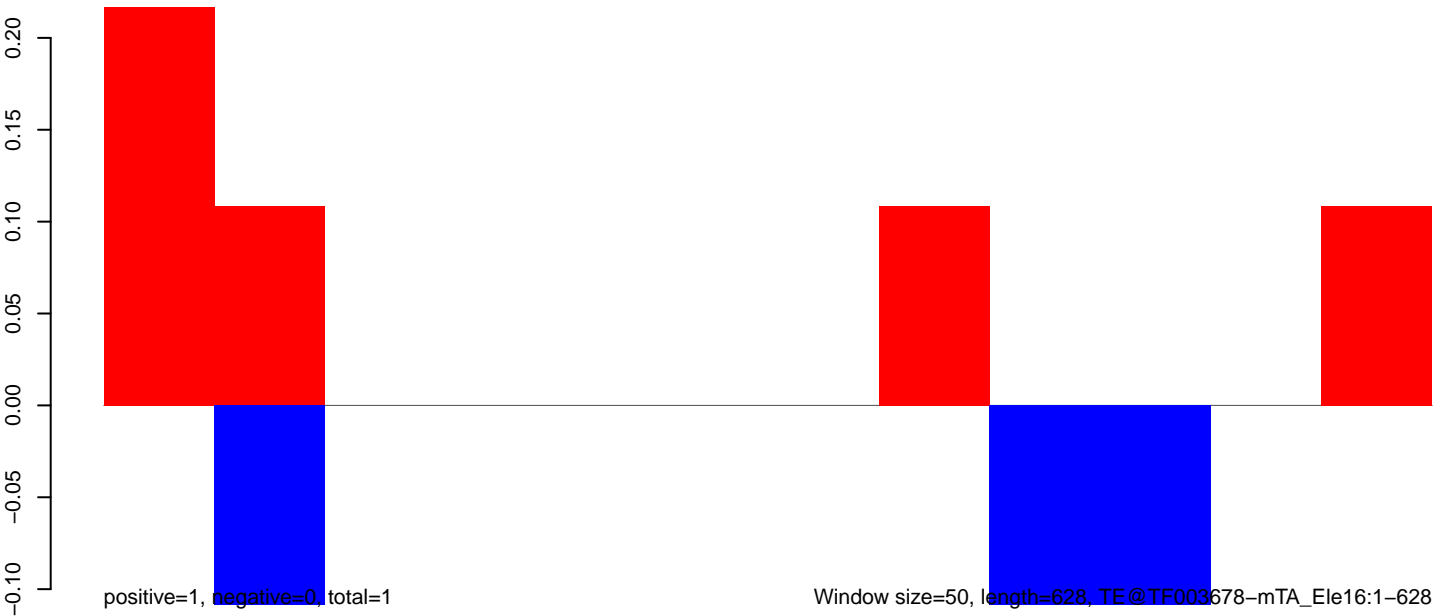
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



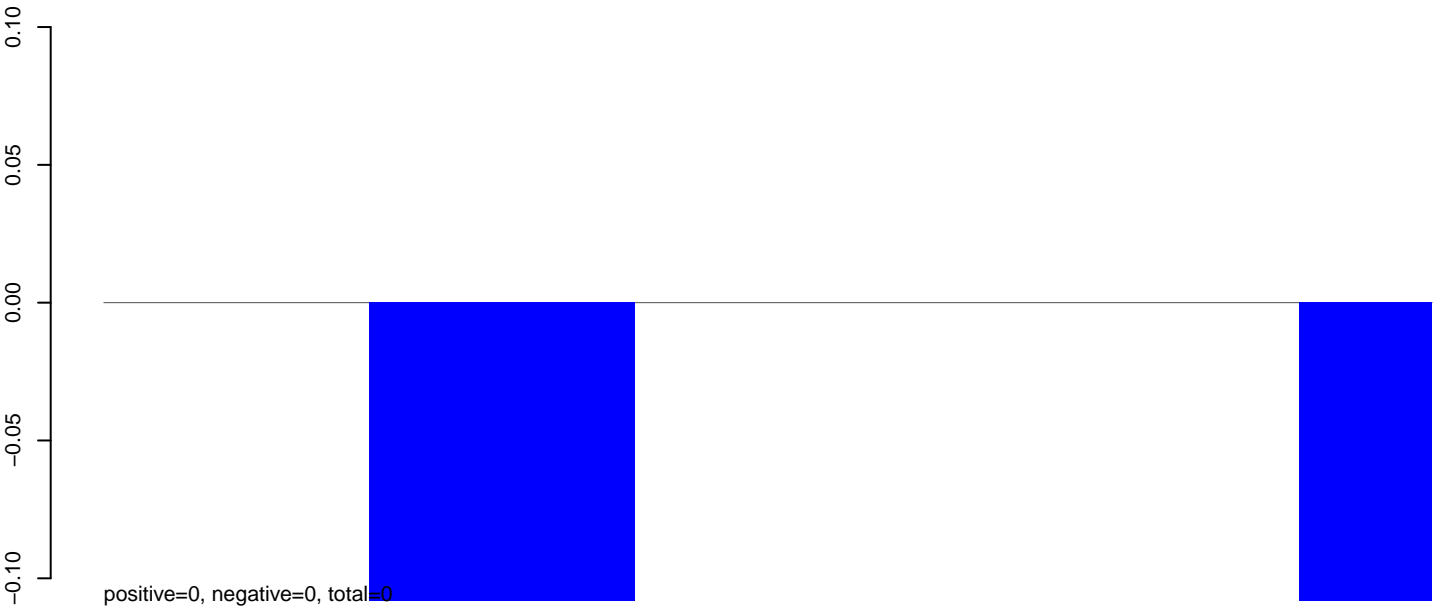
AnGam_Sua5b_cells.rep



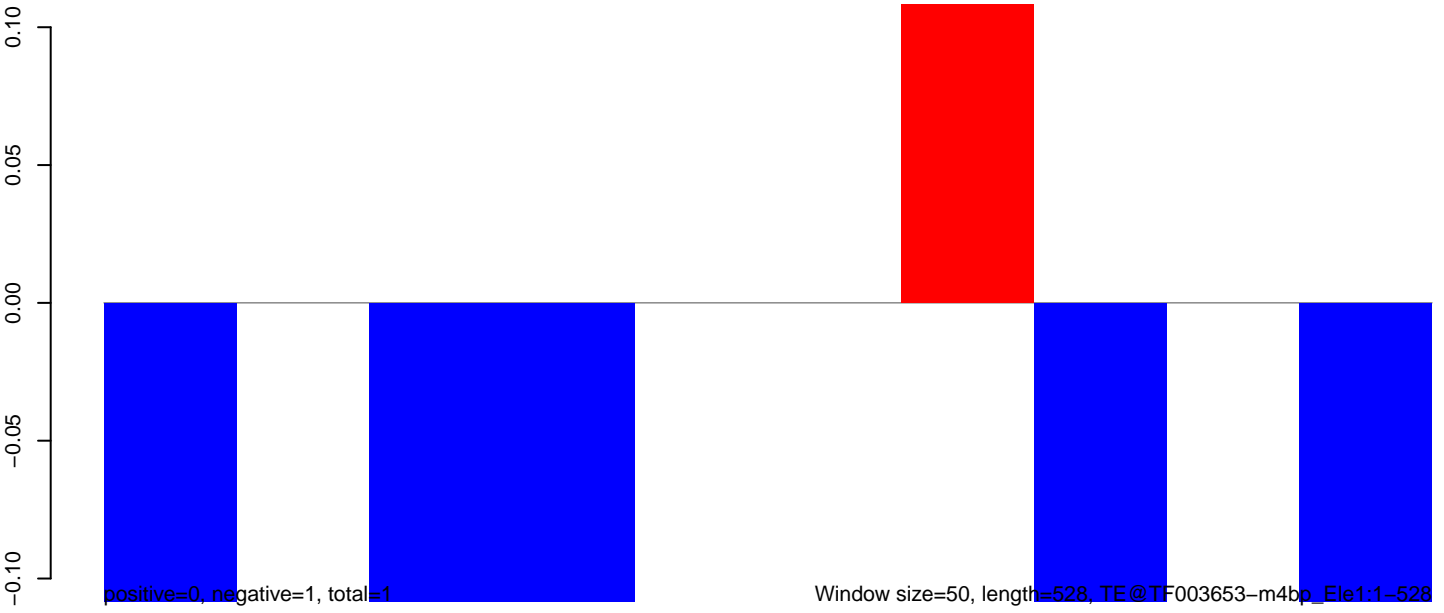
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

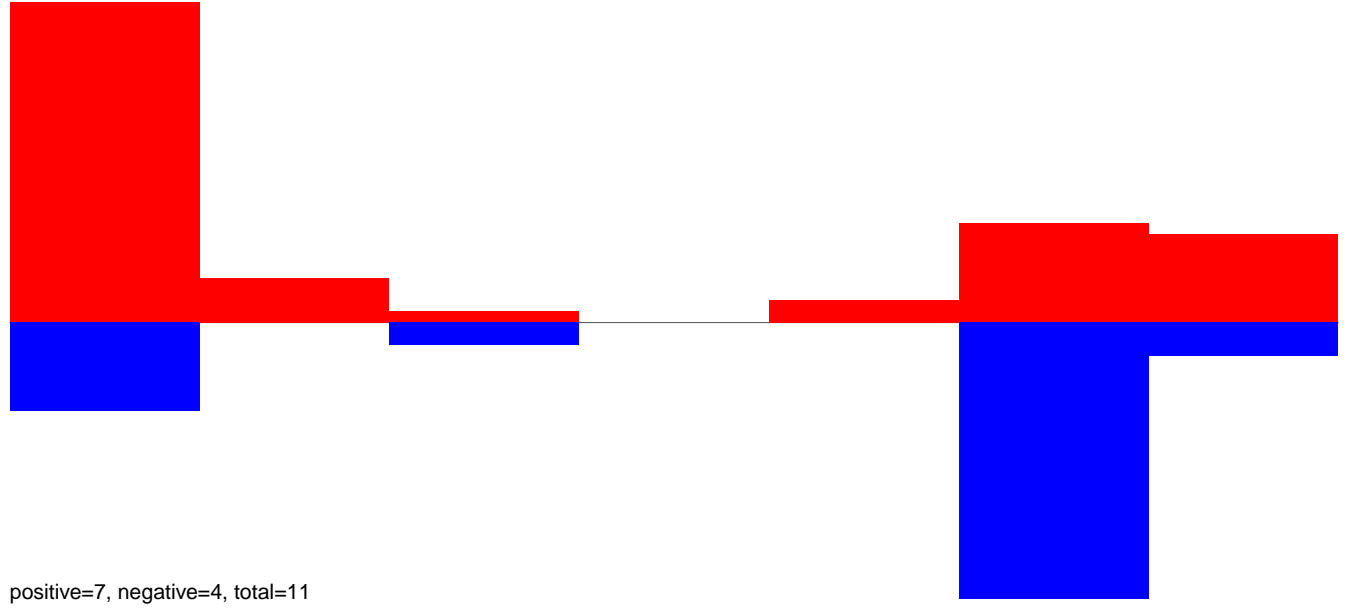
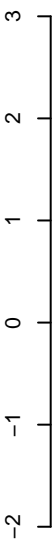


AnGam_Sua5b_cells.rep



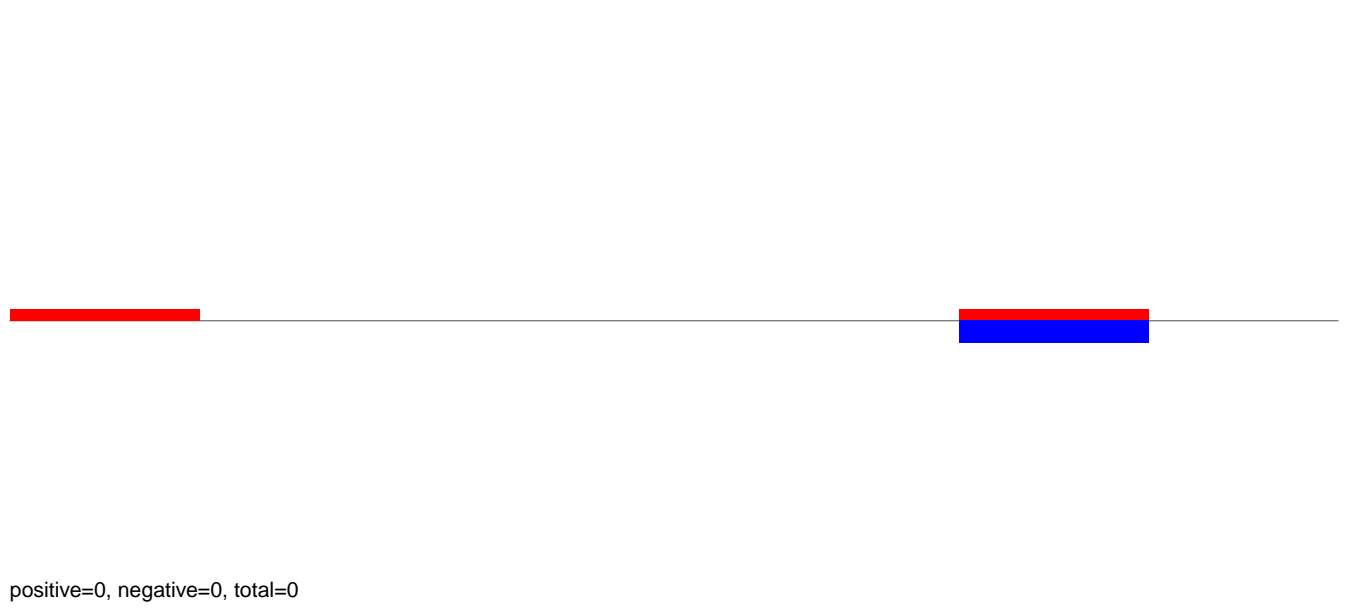
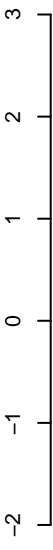
100 200 300 400 500

AnGam_Sua5b_cells.18_23.rep



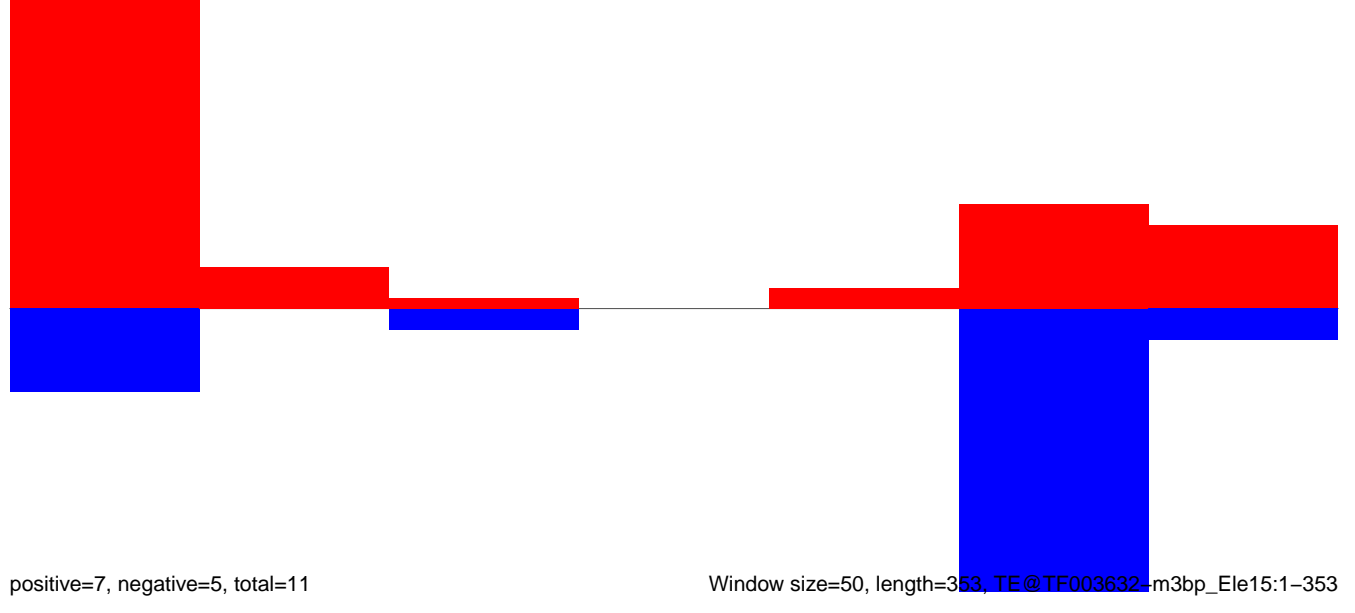
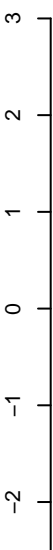
positive=7, negative=4, total=11

AnGam_Sua5b_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Sua5b_cells.rep

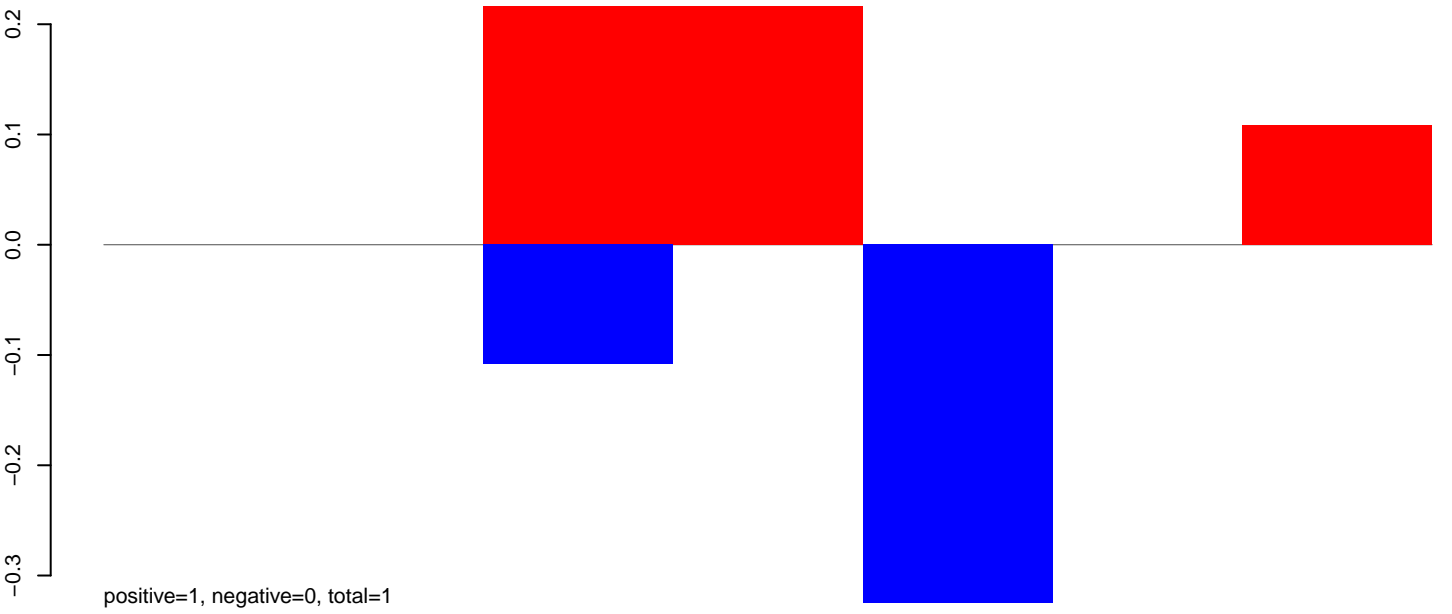


positive=7, negative=5, total=11

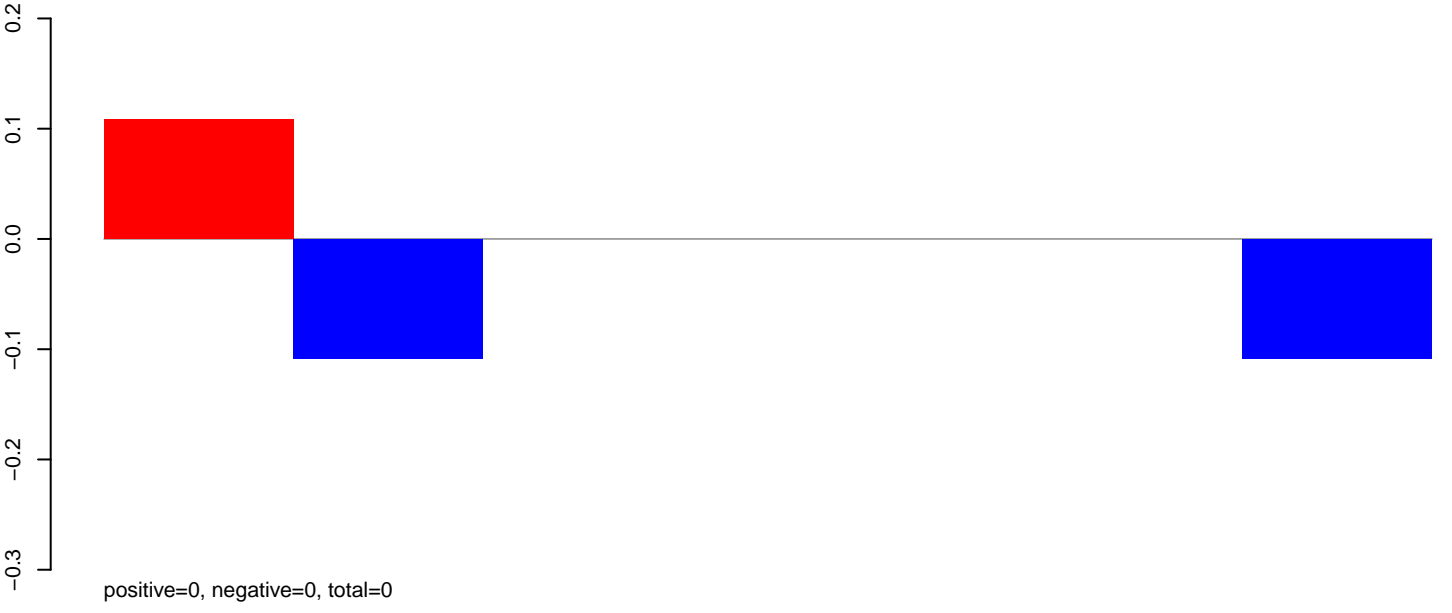
Window size=50, length=353, TE@TF003632-m3bp_Ele15:1-353

50 100 150 200 250 300 350 400

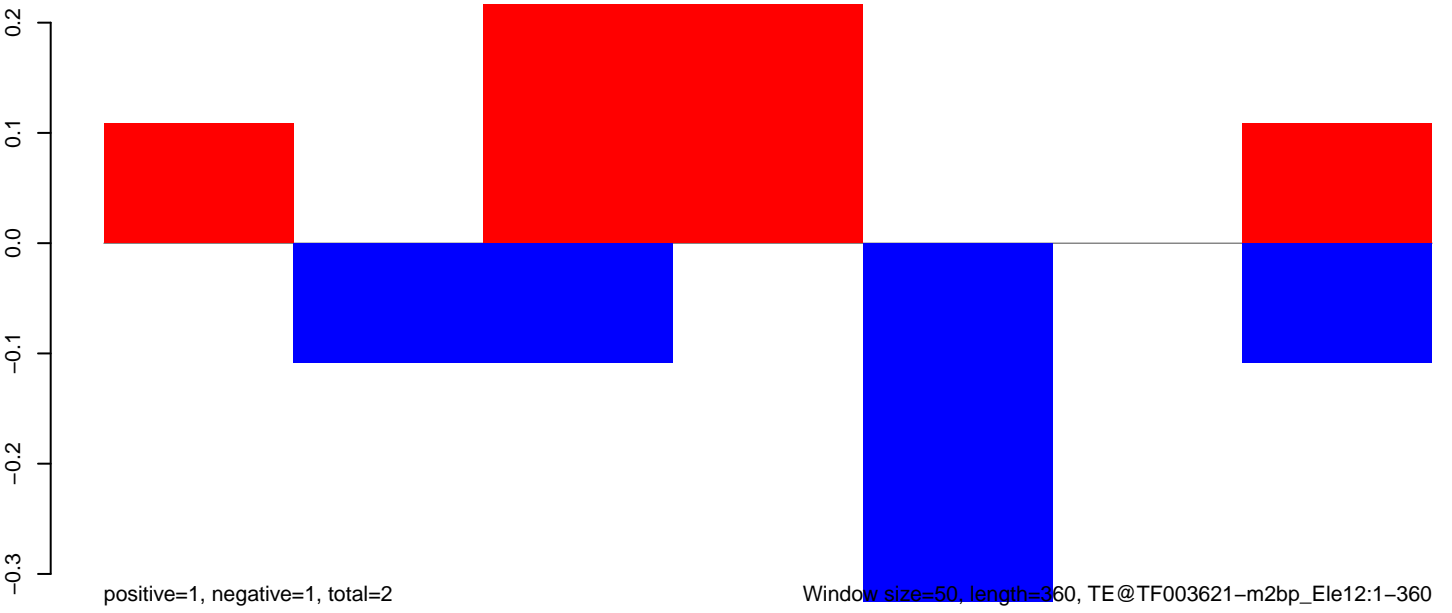
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



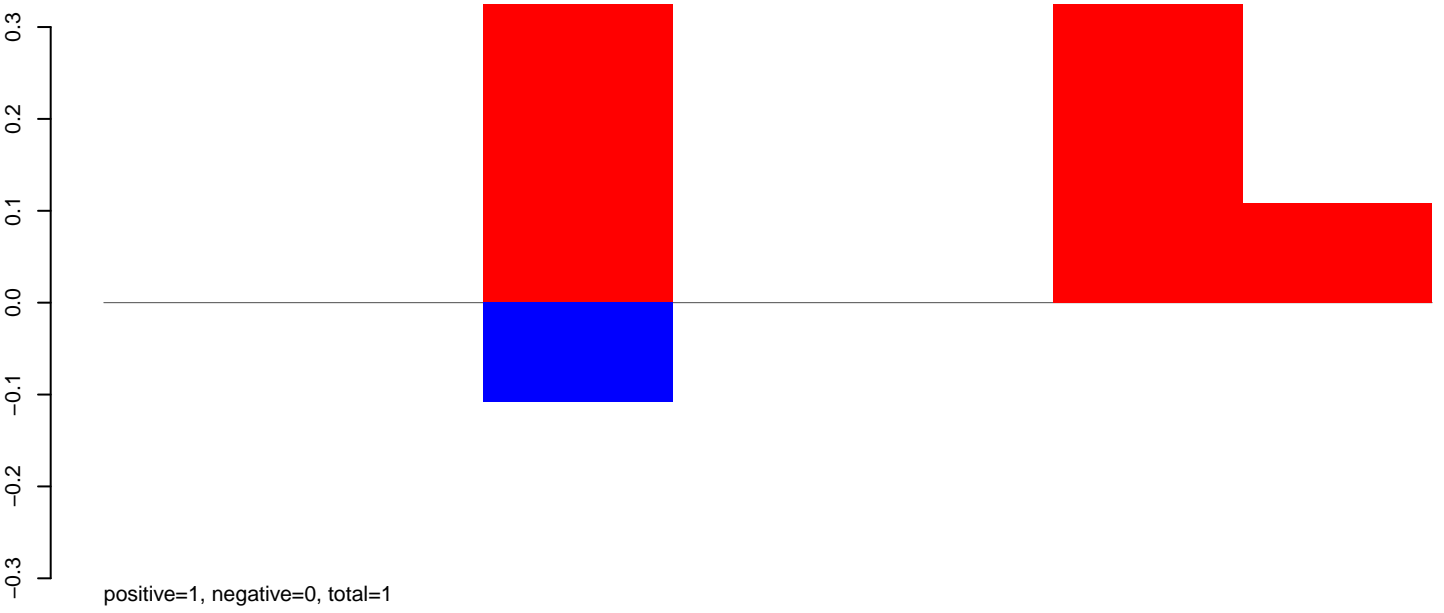
Window size=50, length=360, TE@TF003621-m2bp_Ele12:1-360

50 100 150 200 250 300 350 400

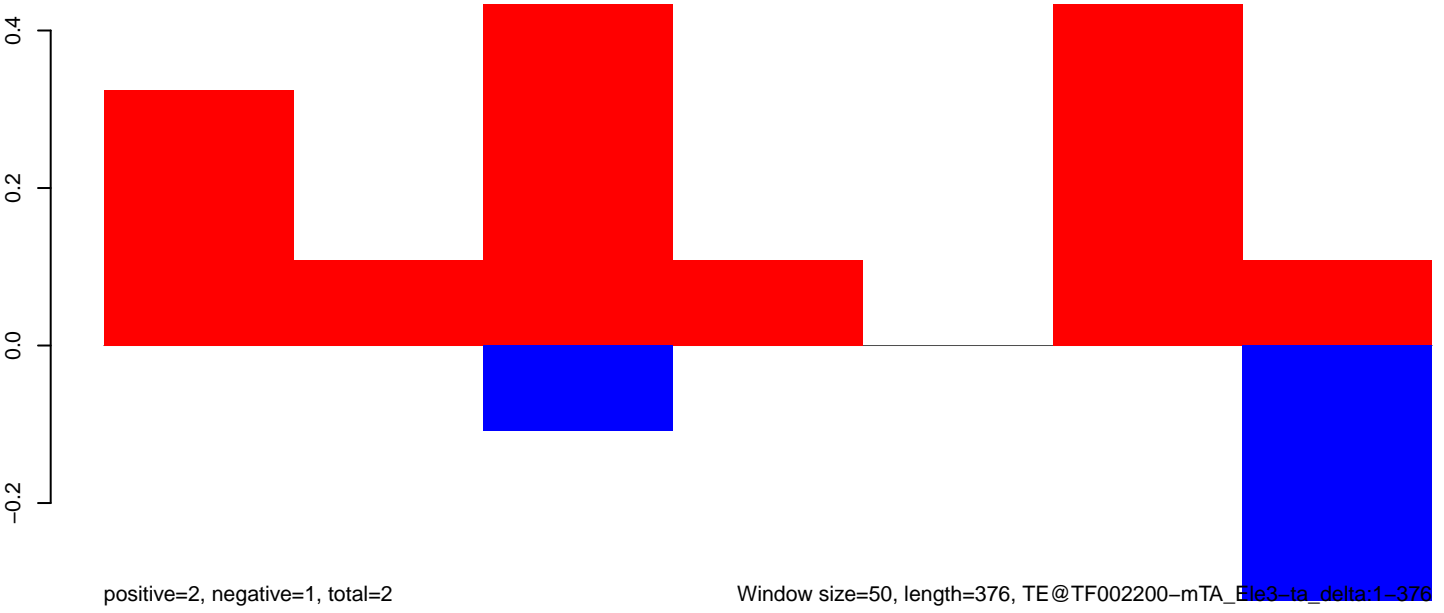
AnGam_Sua5b_cells.18_23.rep



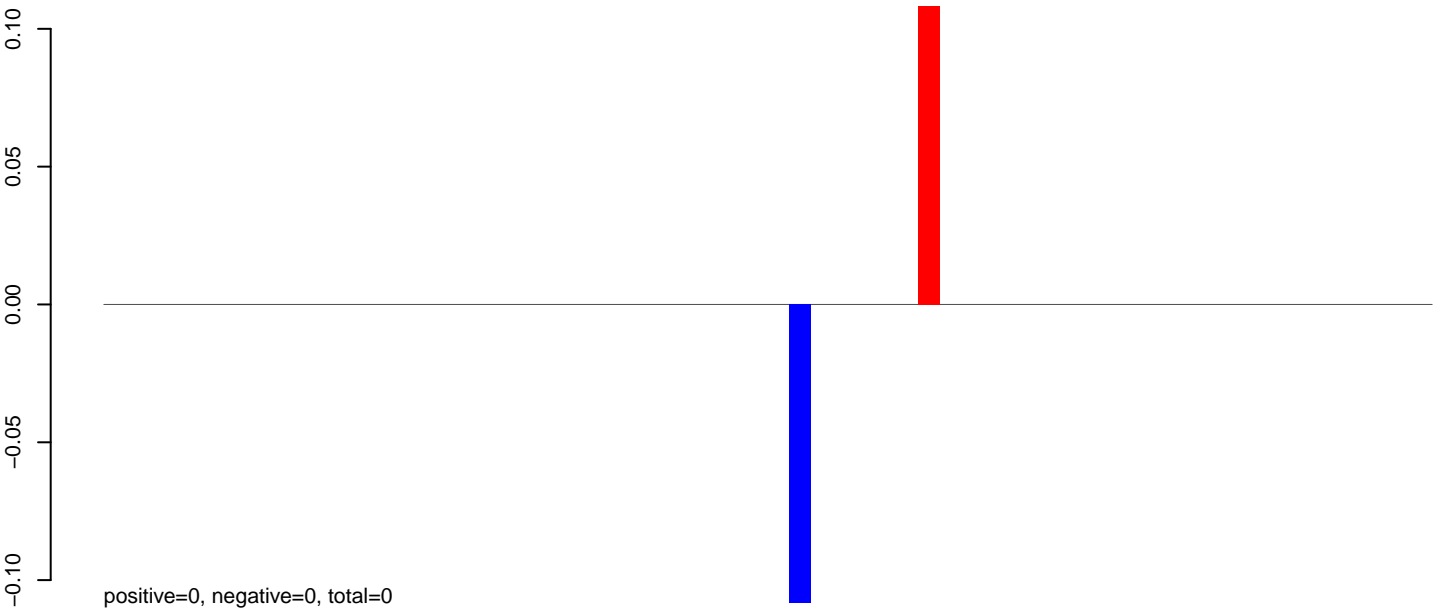
AnGam_Sua5b_cells.24_35.rep



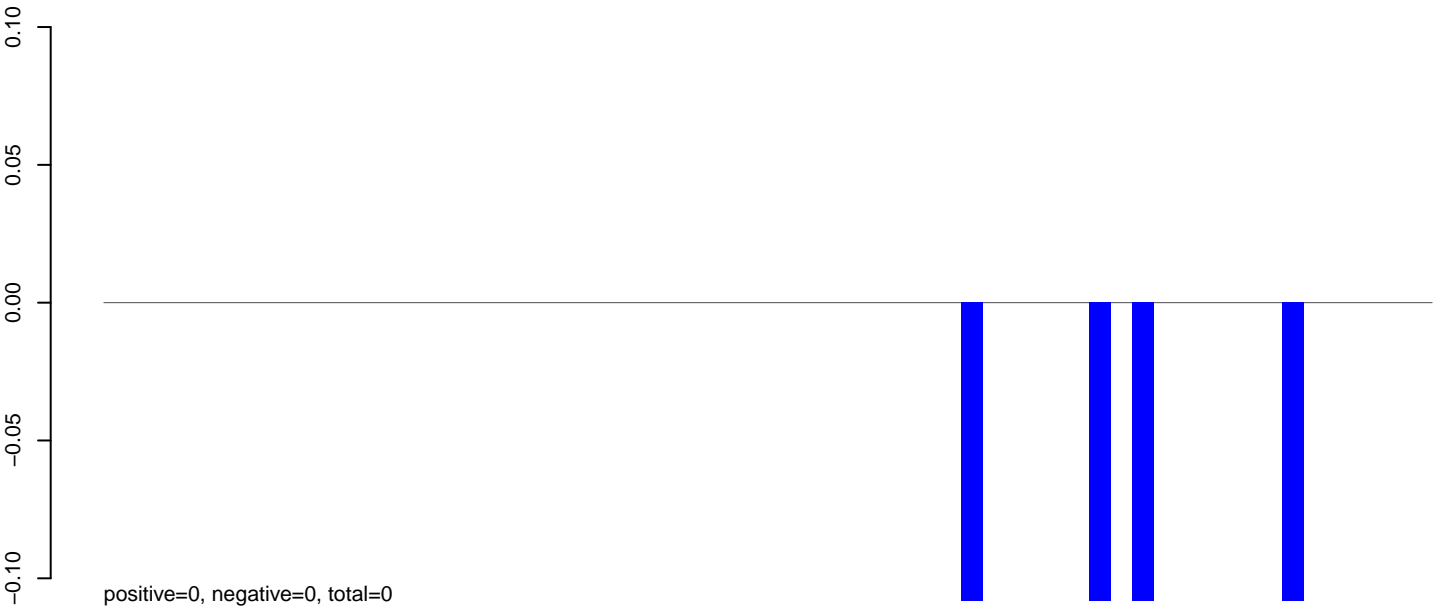
AnGam_Sua5b_cells.rep



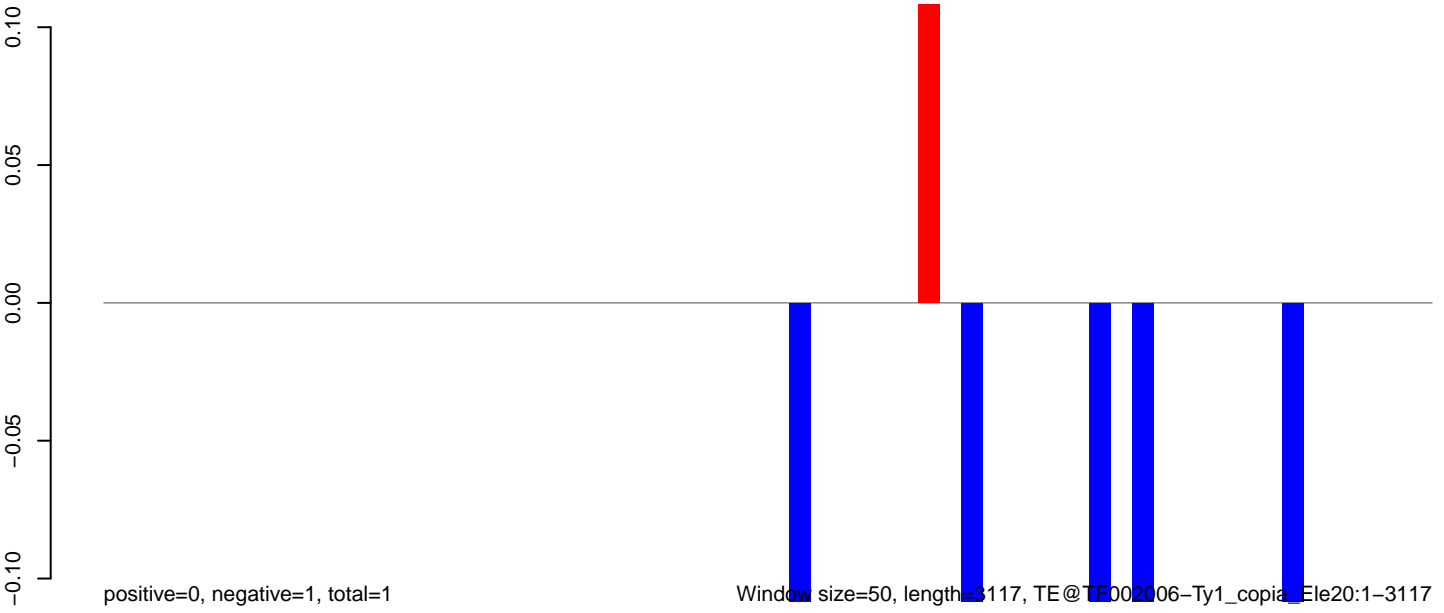
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

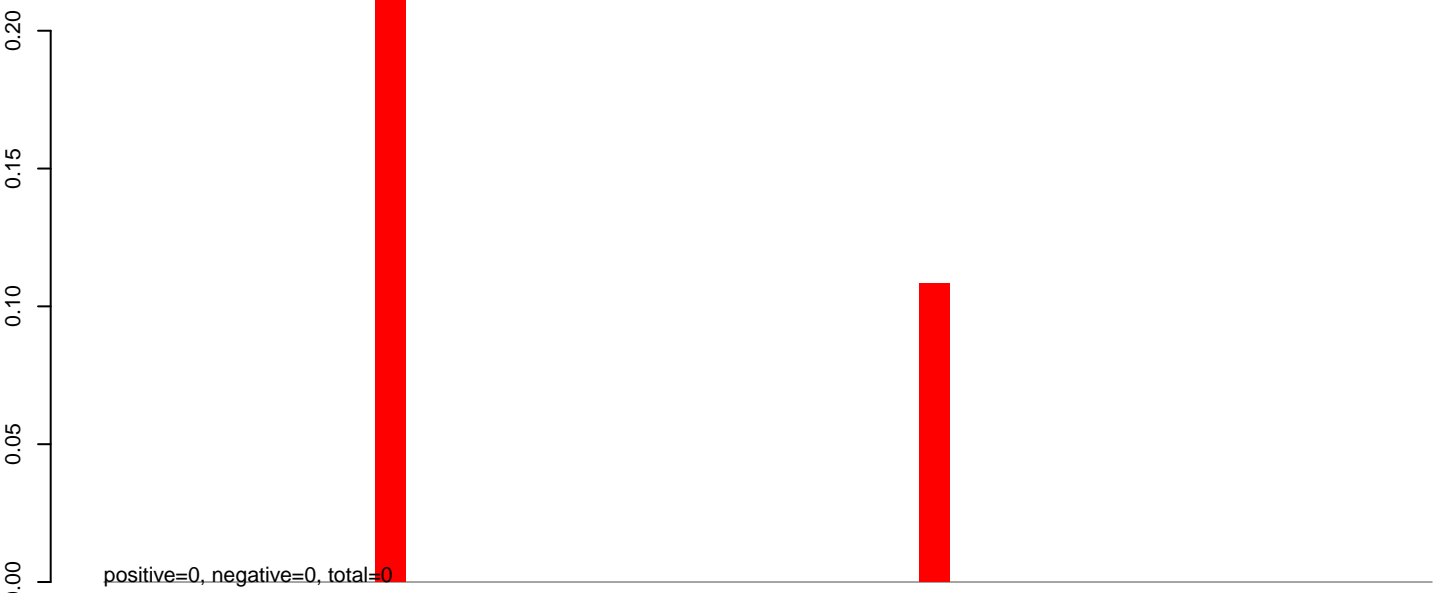


AnGam_Sua5b_cells.rep

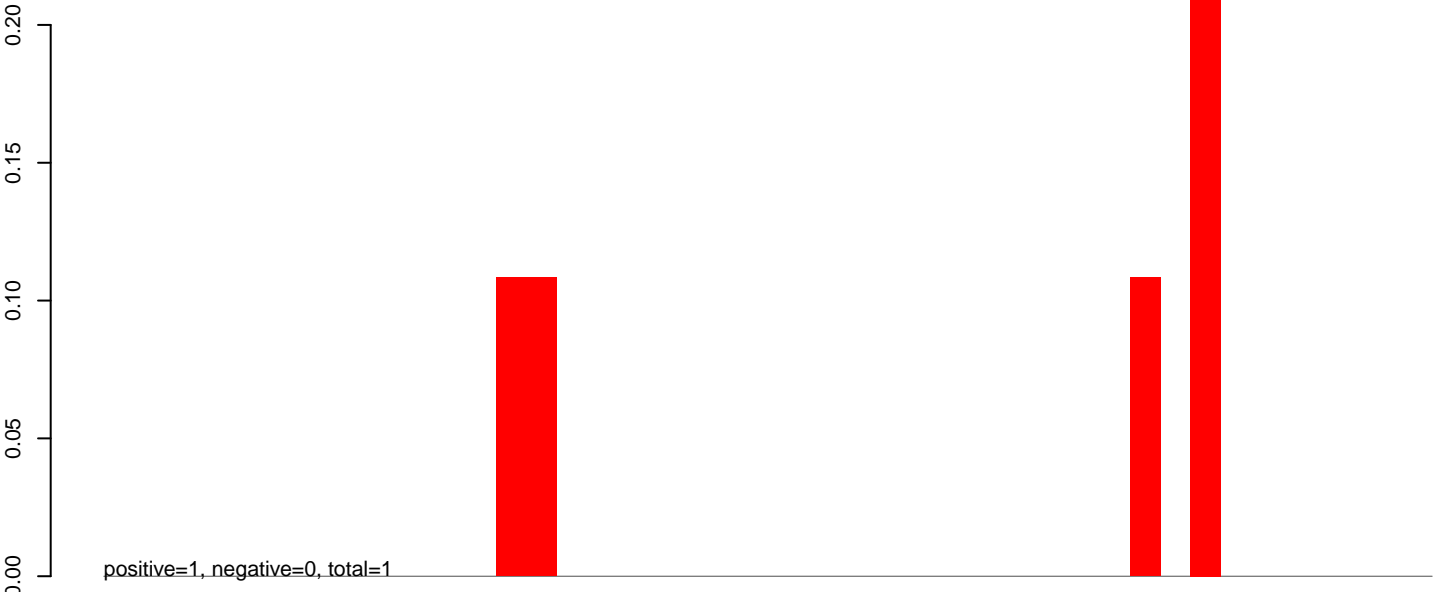


0 500 1000 1500 2000 2500 3000

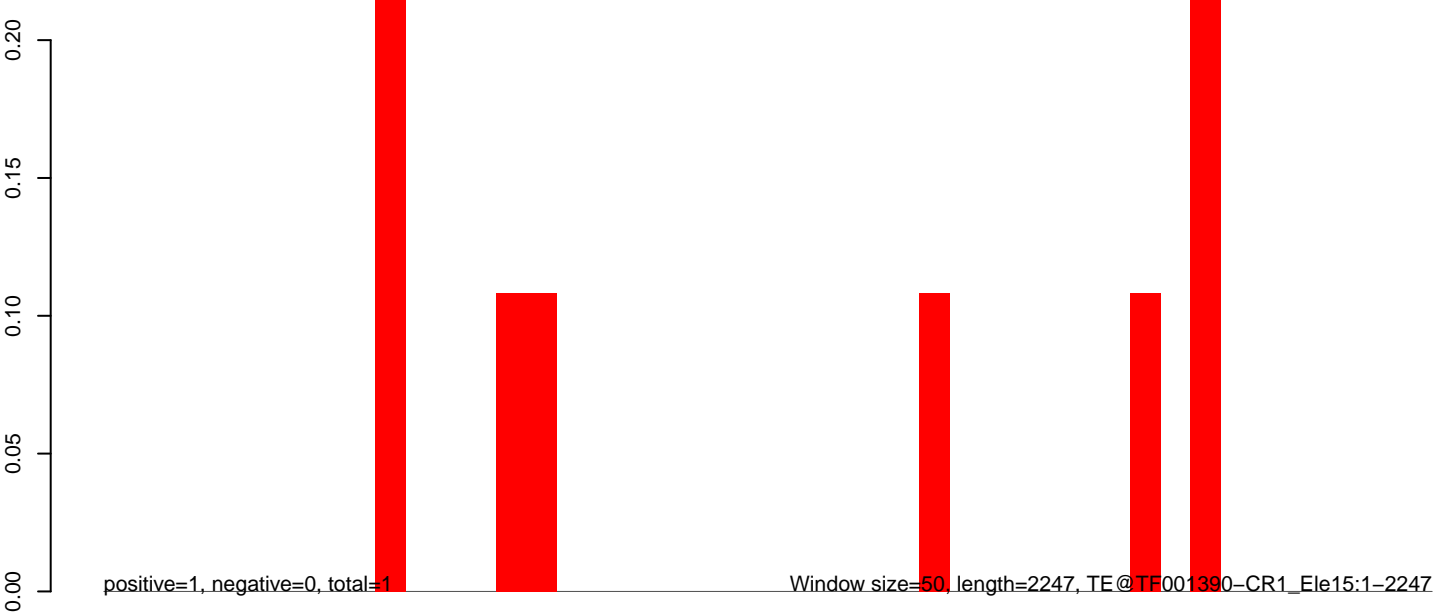
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

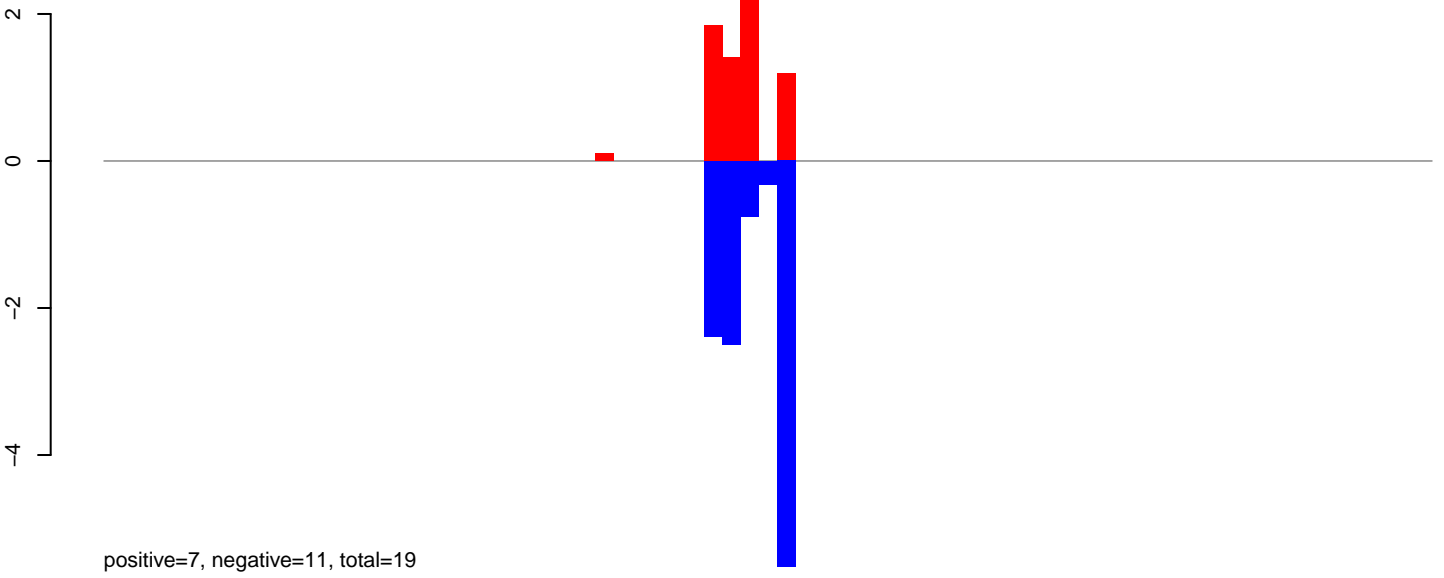


AnGam_Sua5b_cells.rep

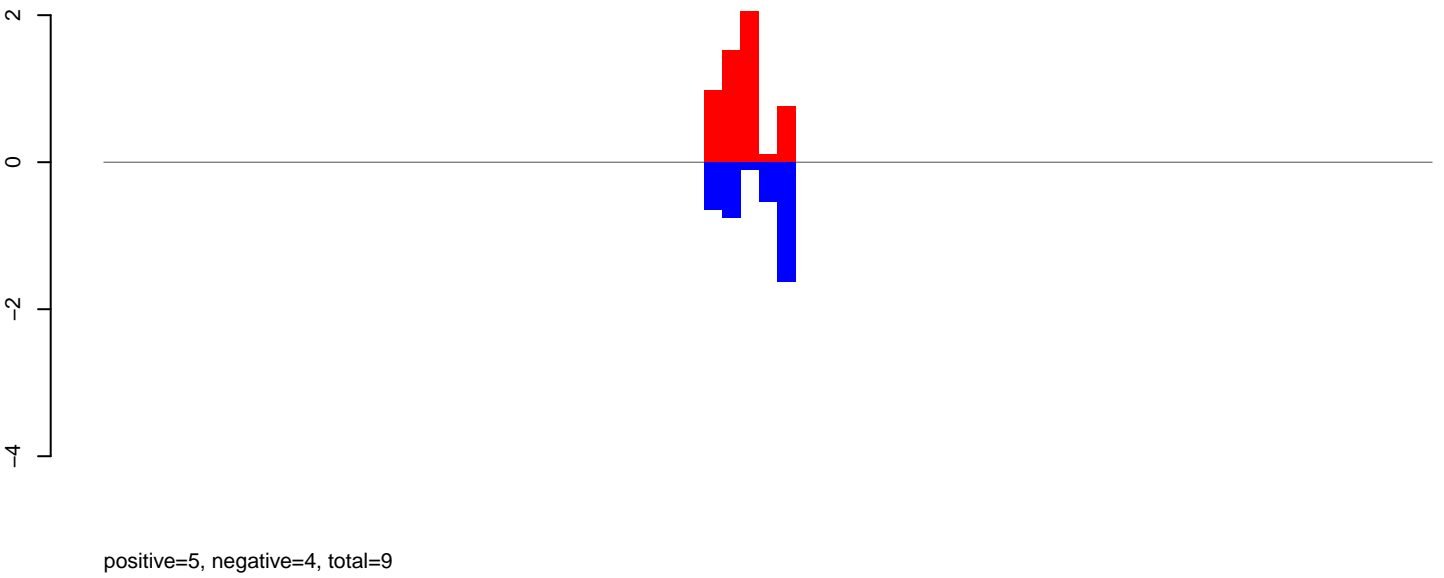


0 500 1000 1500 2000

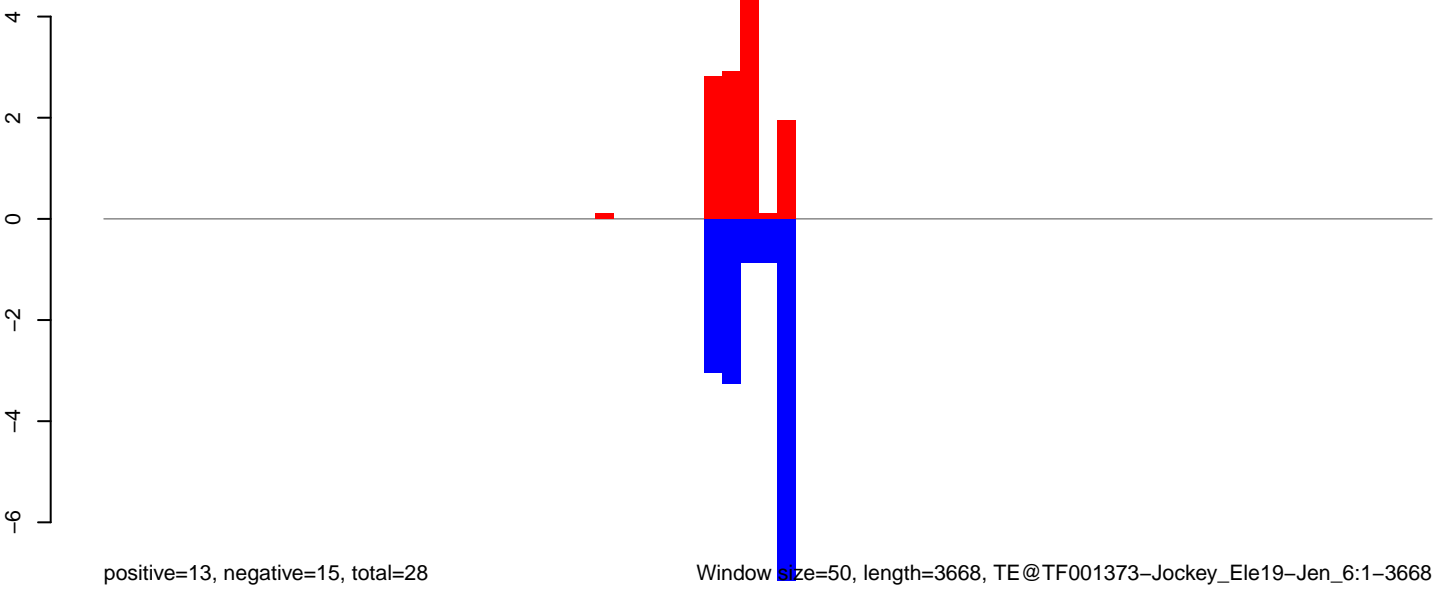
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



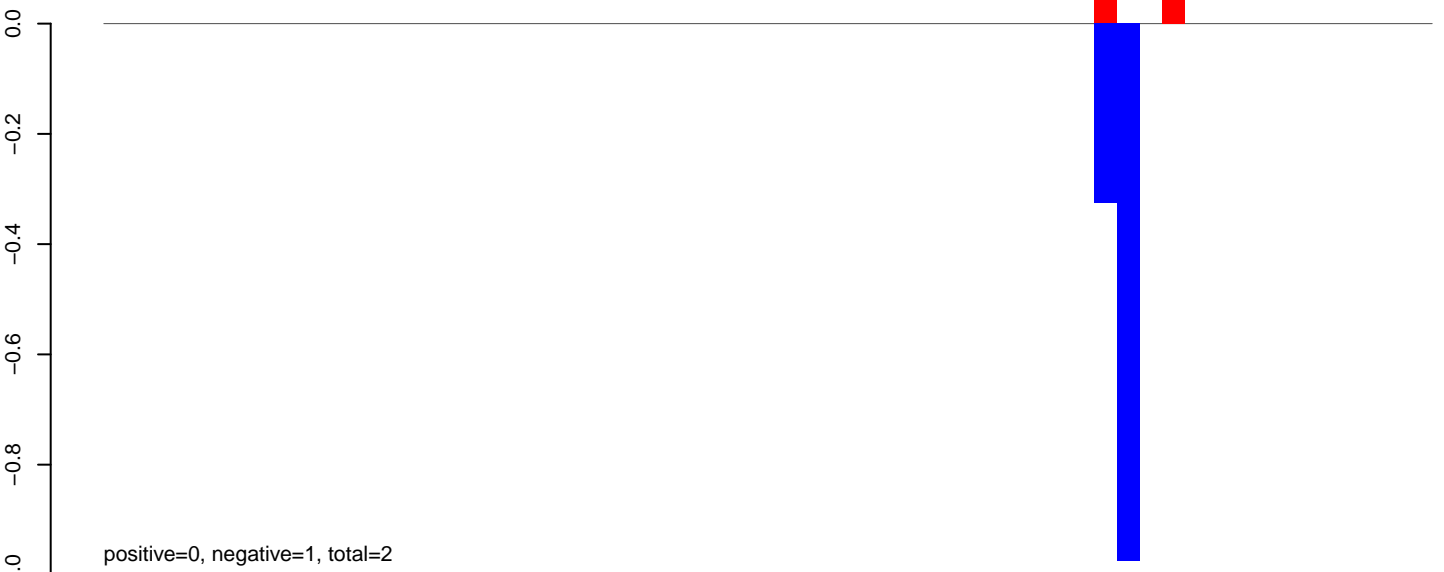
AnGam_Sua5b_cells.rep



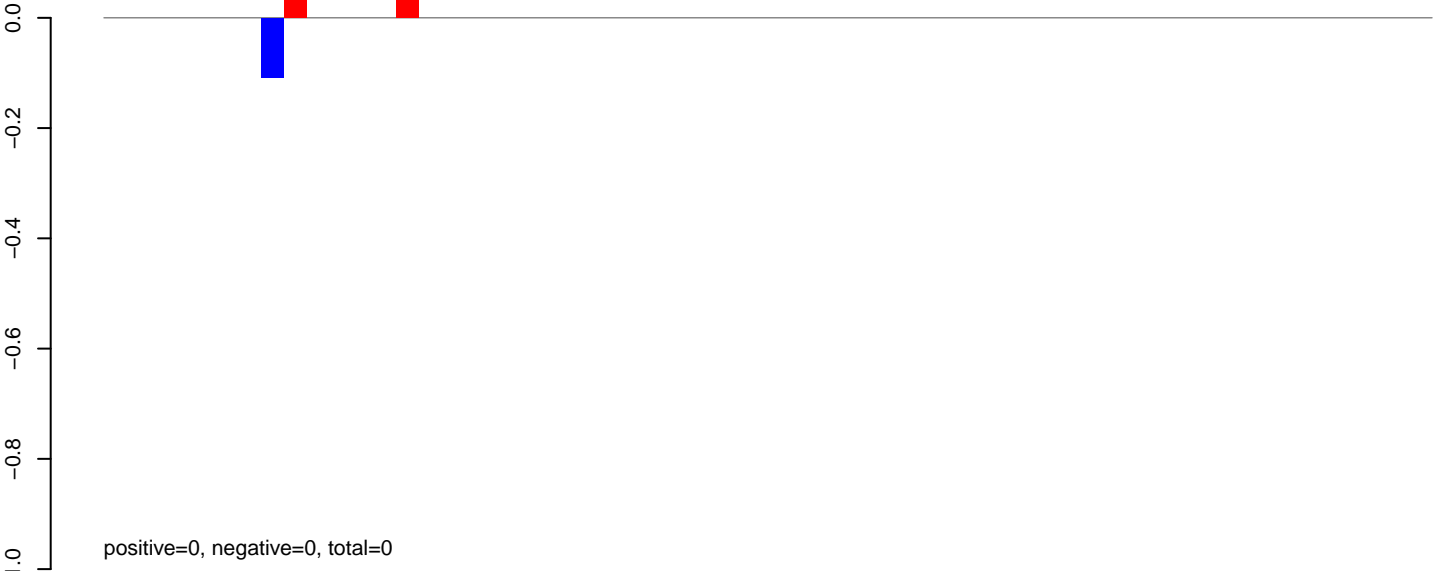
Window size=50, length=3668, TE@TF001373-Jockey_Ele19-Jen_6:1-3668

0 1000 2000 3000

AnGam_Sua5b_cells.18_23.rep



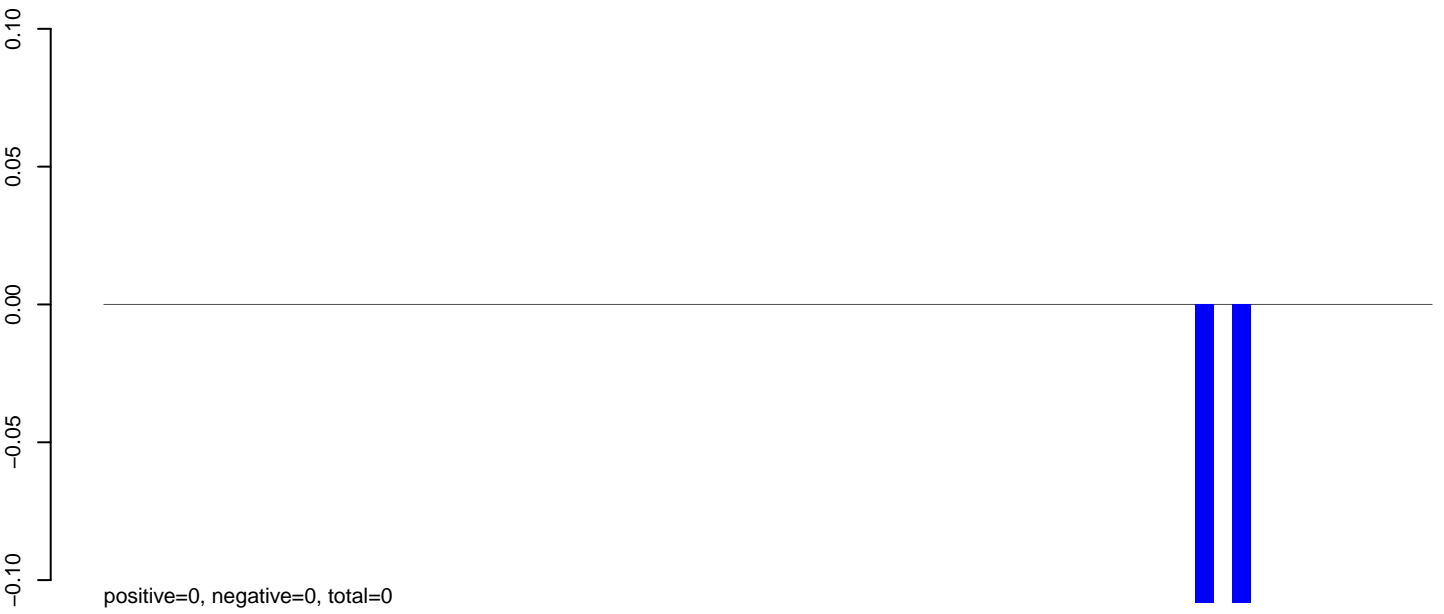
AnGam_Sua5b_cells.24_35.rep



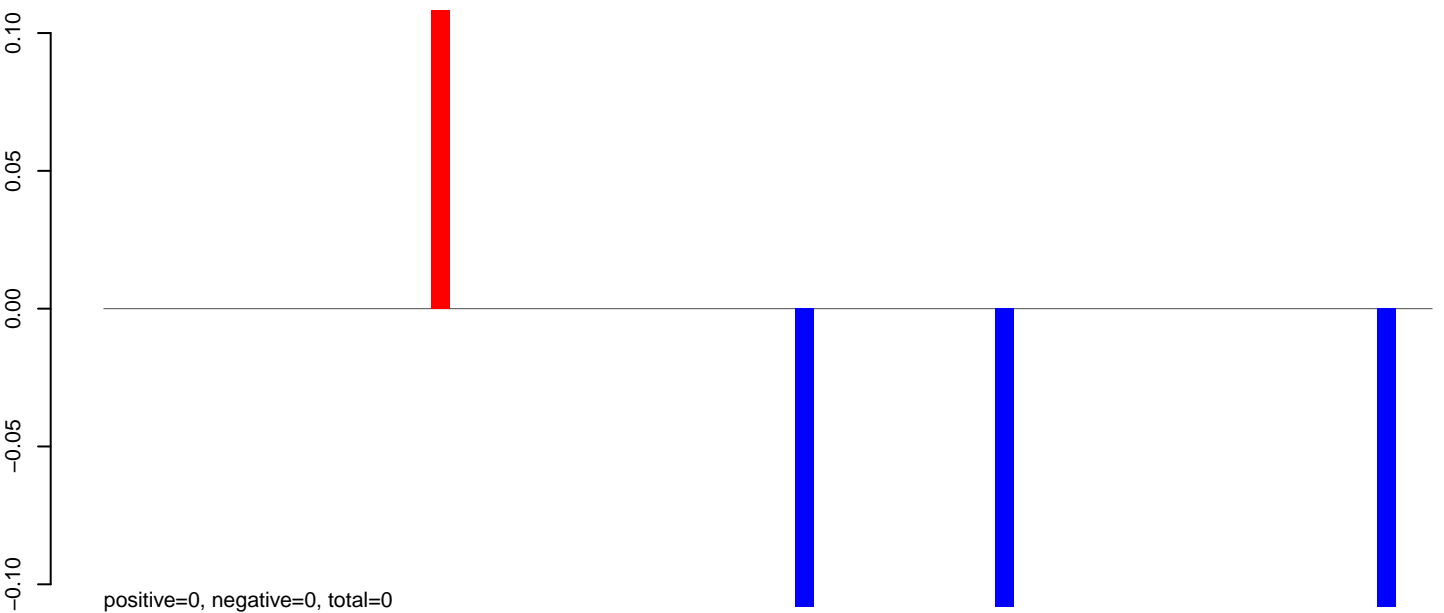
AnGam_Sua5b_cells.rep



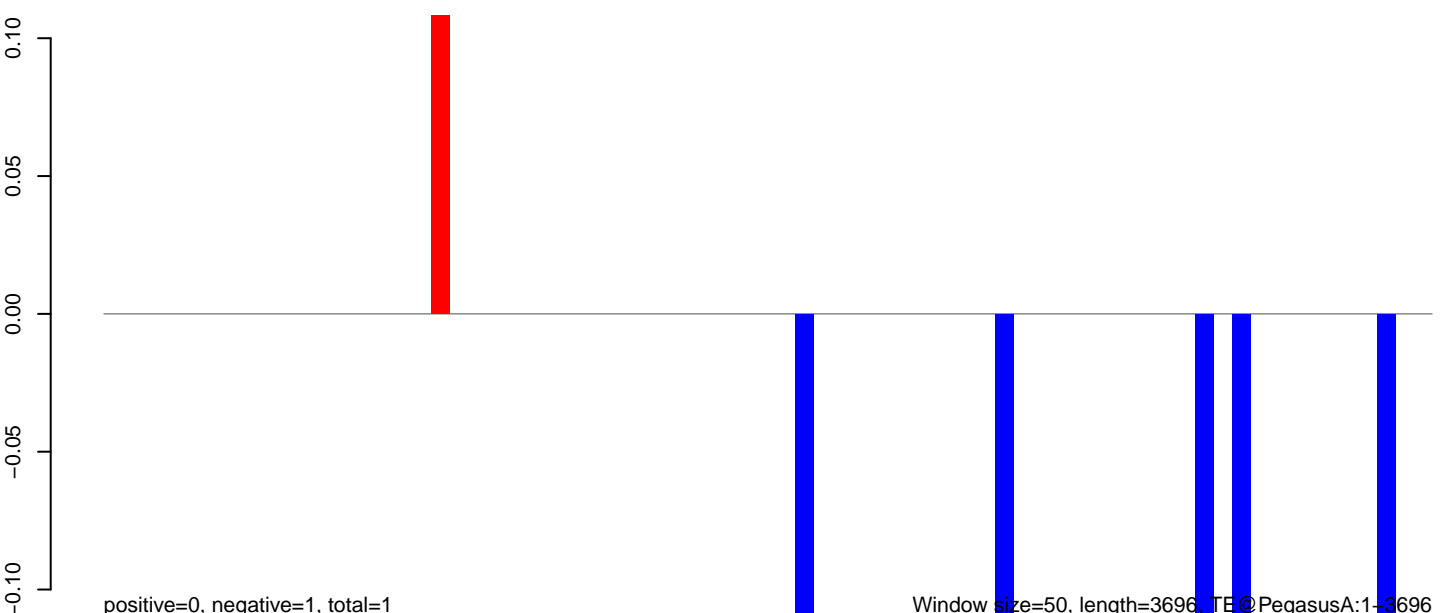
AnGam_Sua5b_cells.18_23.rep



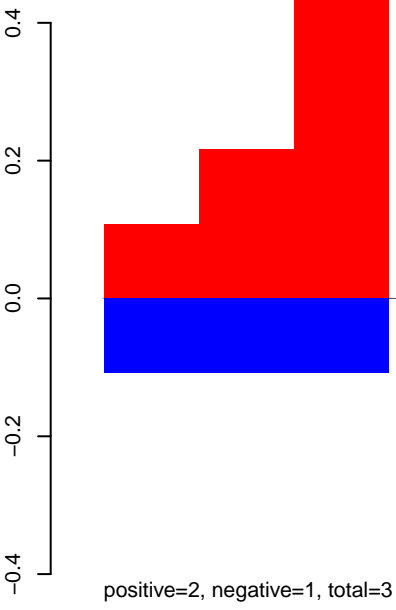
AnGam_Sua5b_cells.24_35.rep



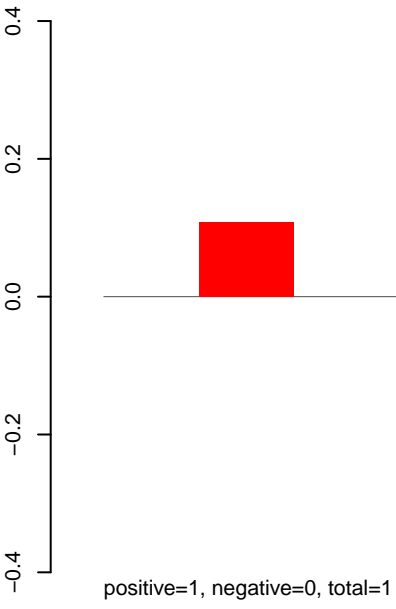
AnGam_Sua5b_cells.rep



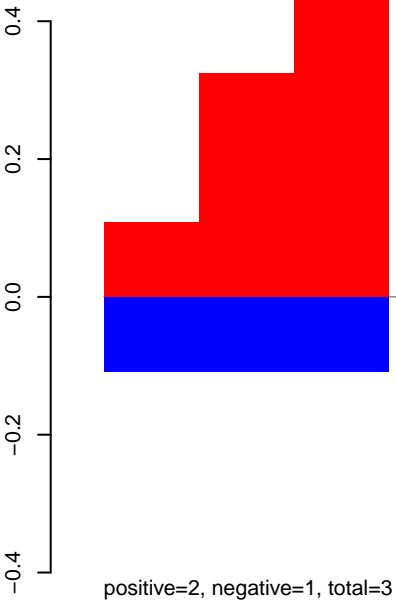
AnGam_Sua5b_cells.18_23.rep



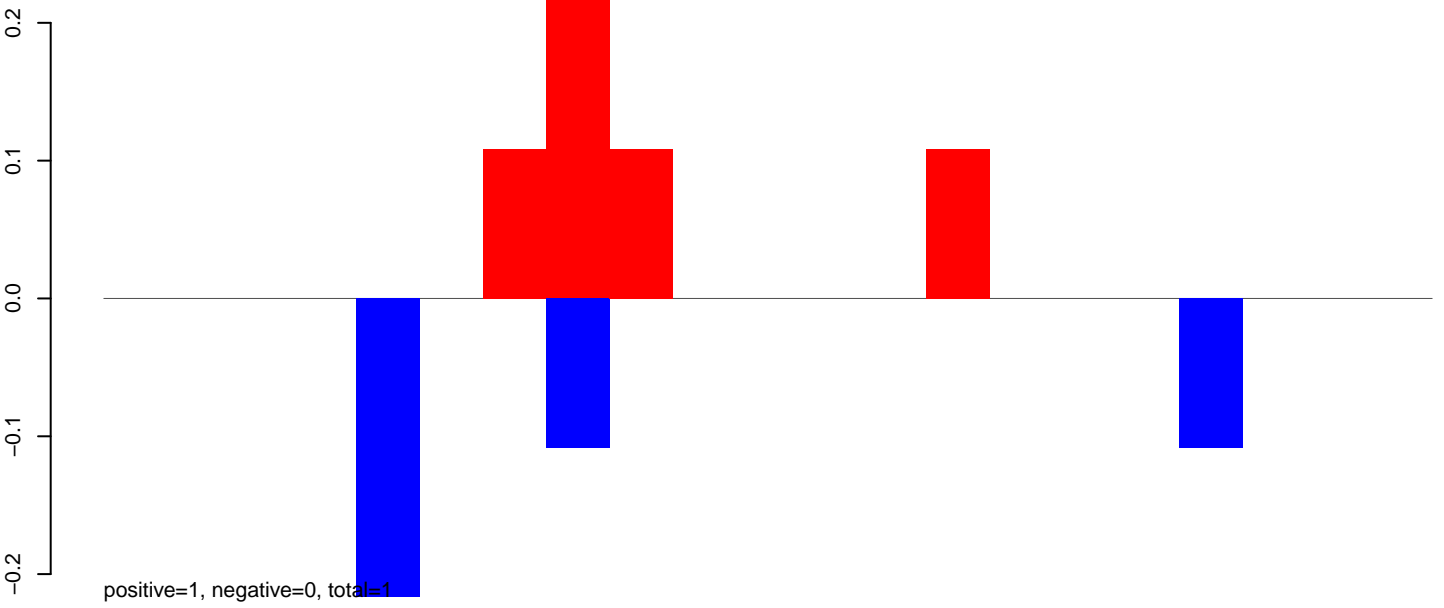
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



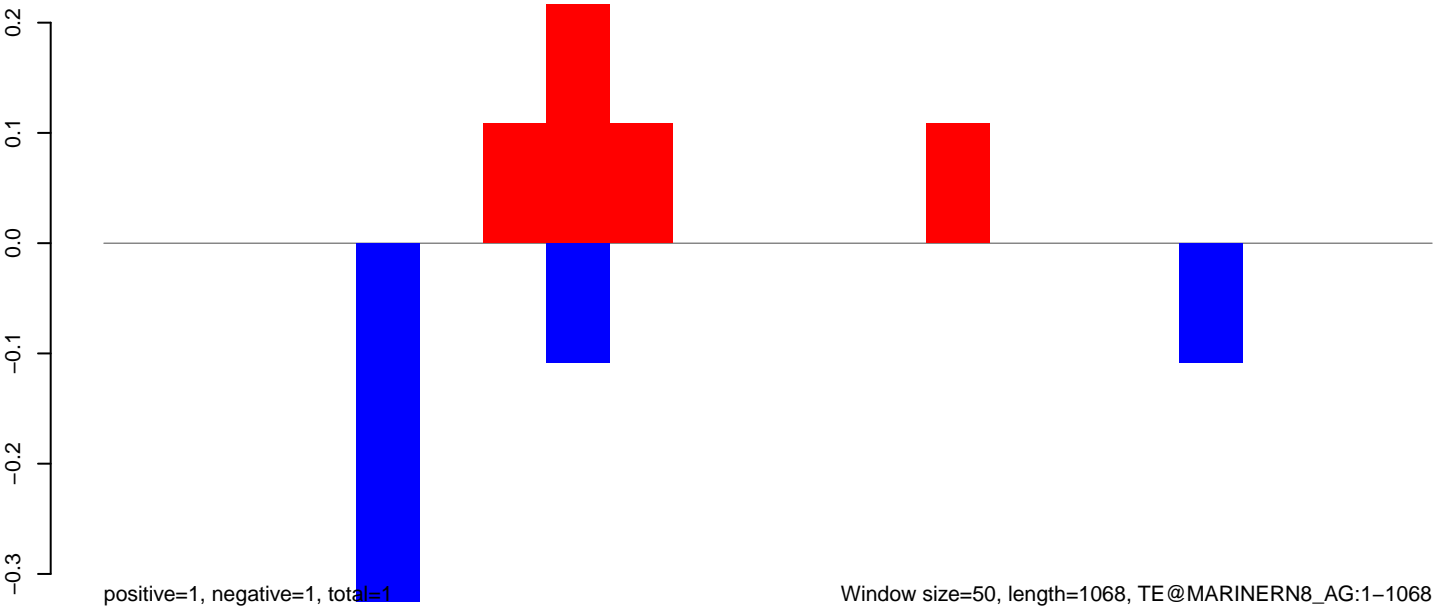
AnGam_Sua5b_cells.18_23.rep



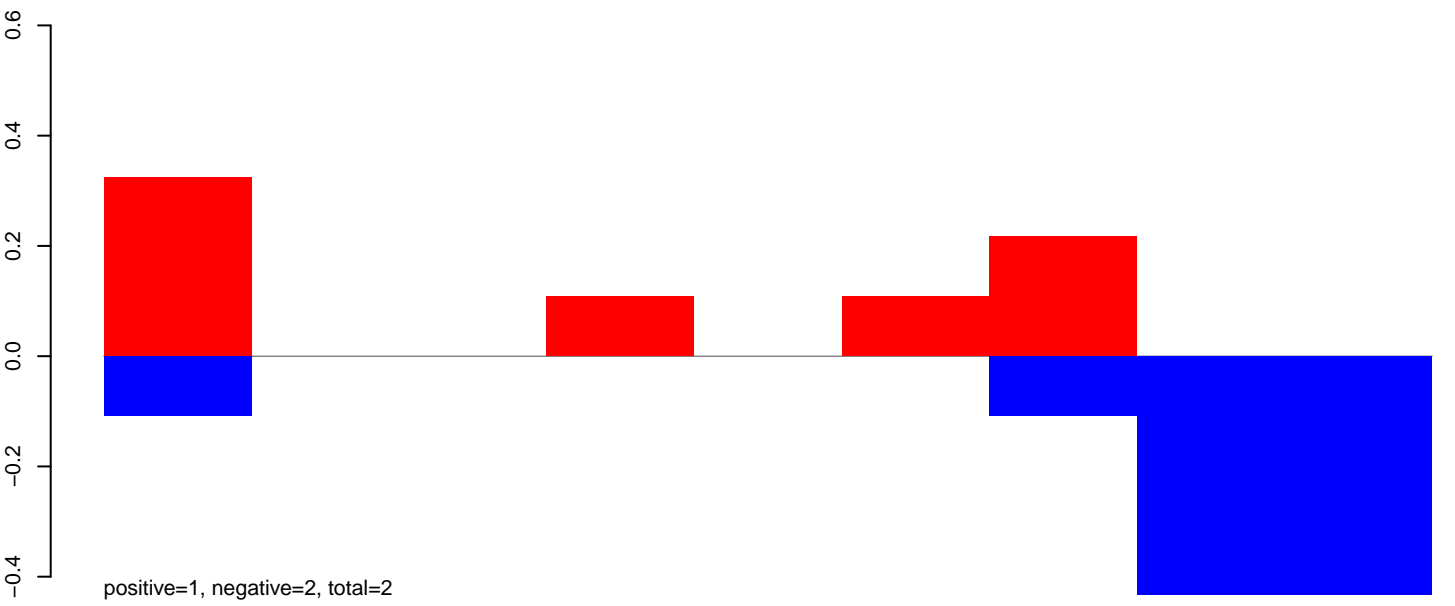
AnGam_Sua5b_cells.24_35.rep



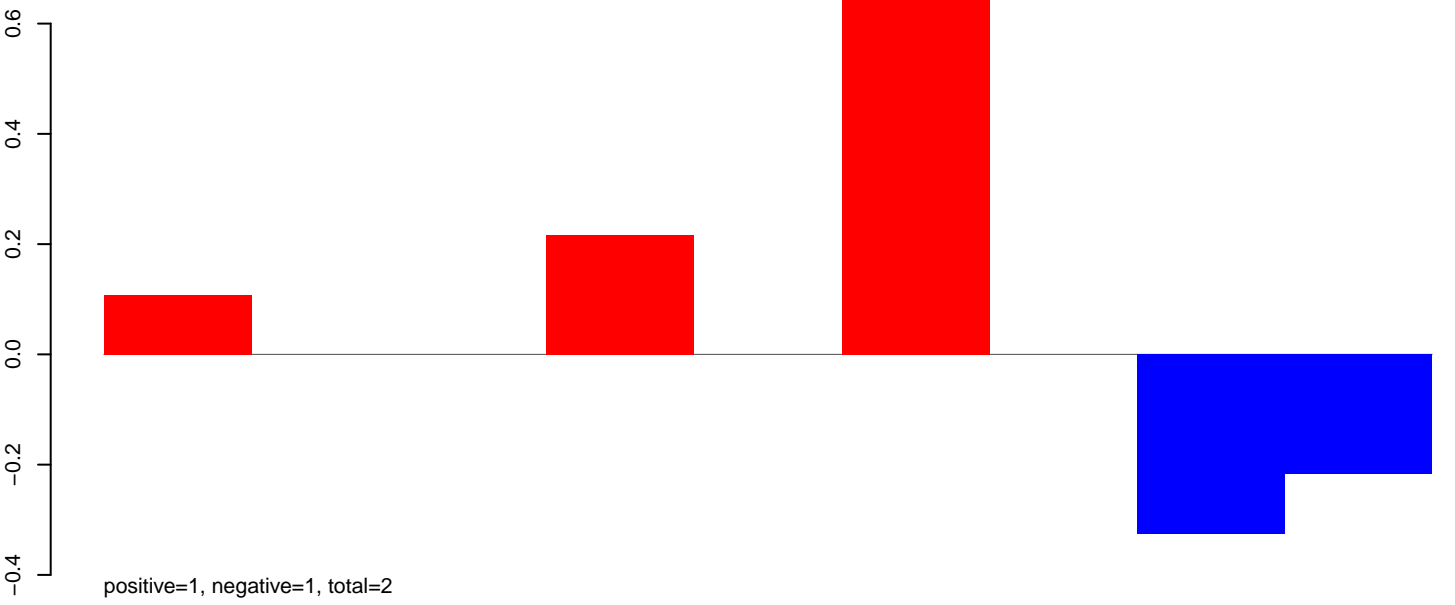
AnGam_Sua5b_cells.rep



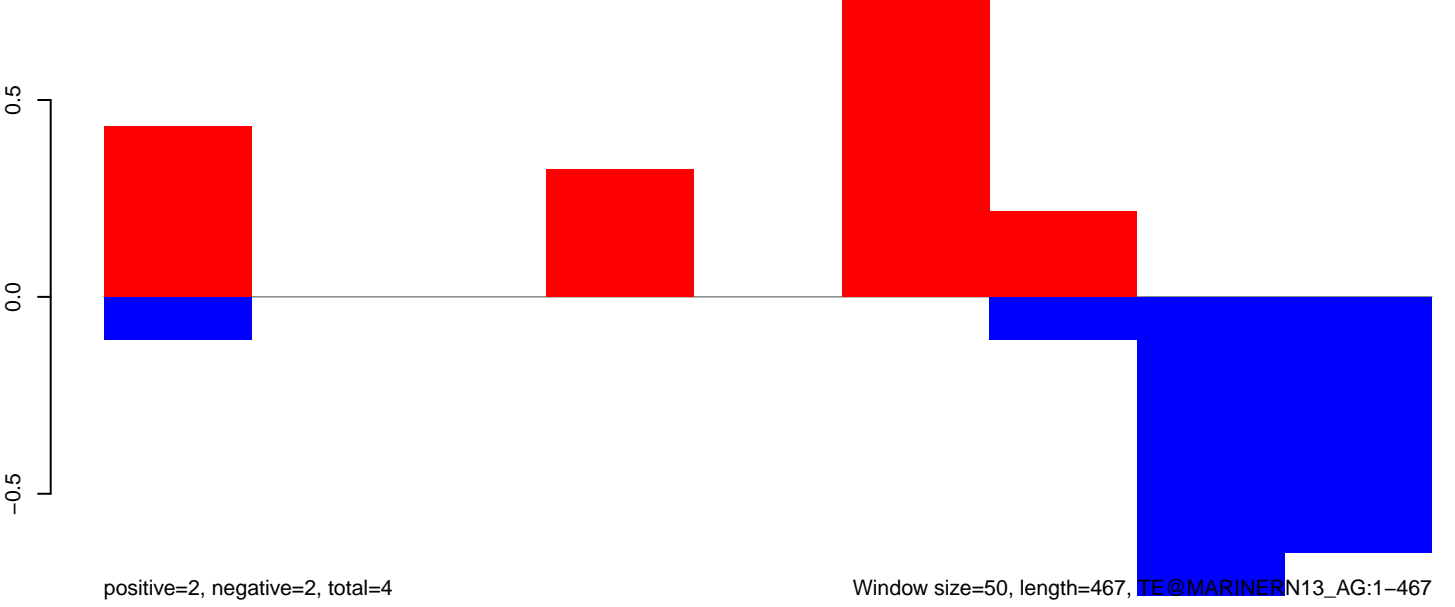
AnGam_Sua5b_cells.18_23.rep



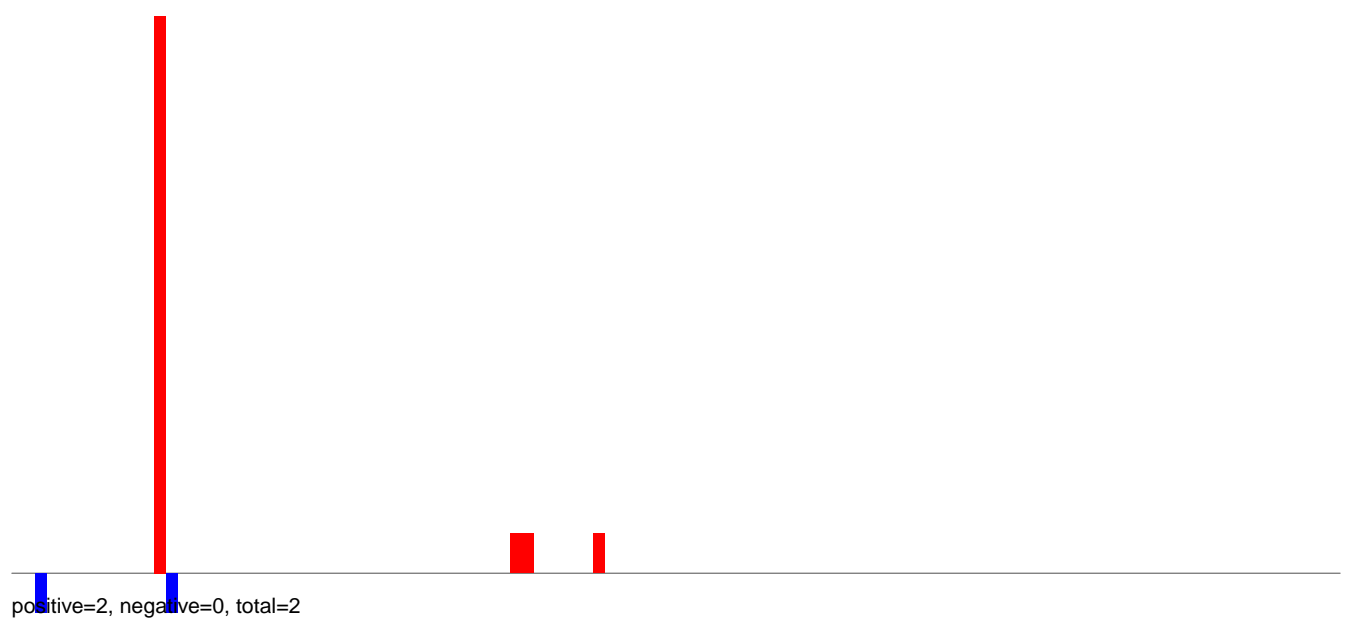
AnGam_Sua5b_cells.24_35.rep



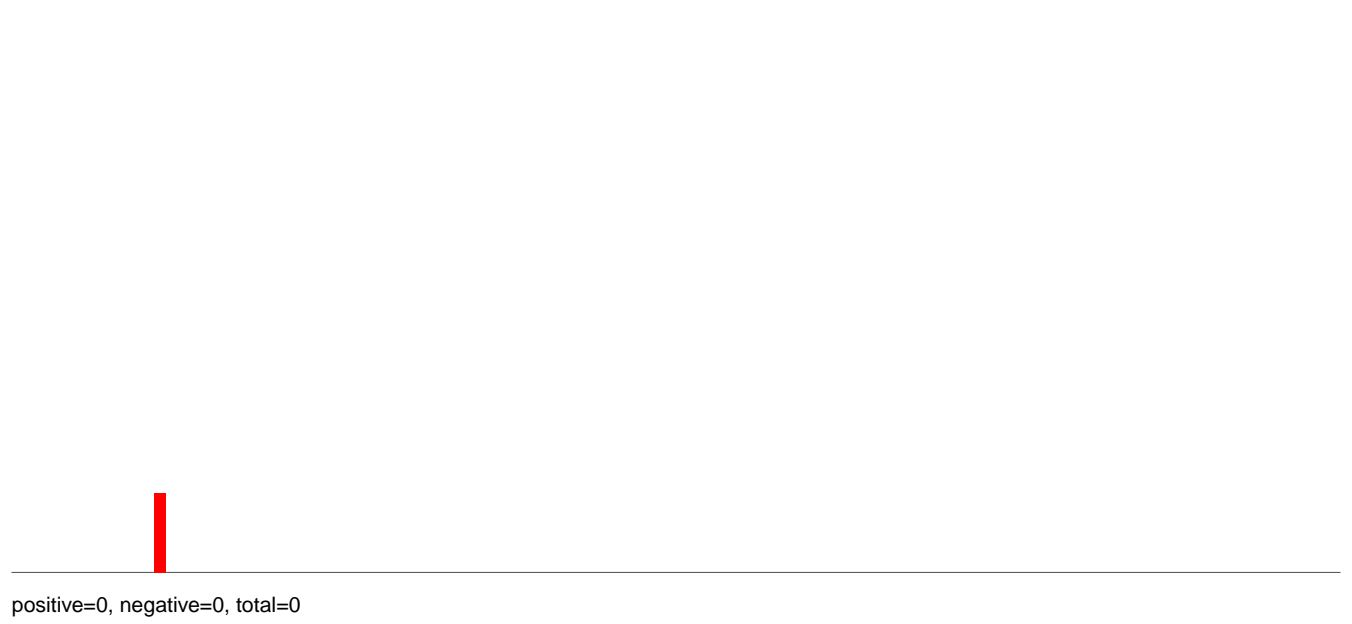
AnGam_Sua5b_cells.rep



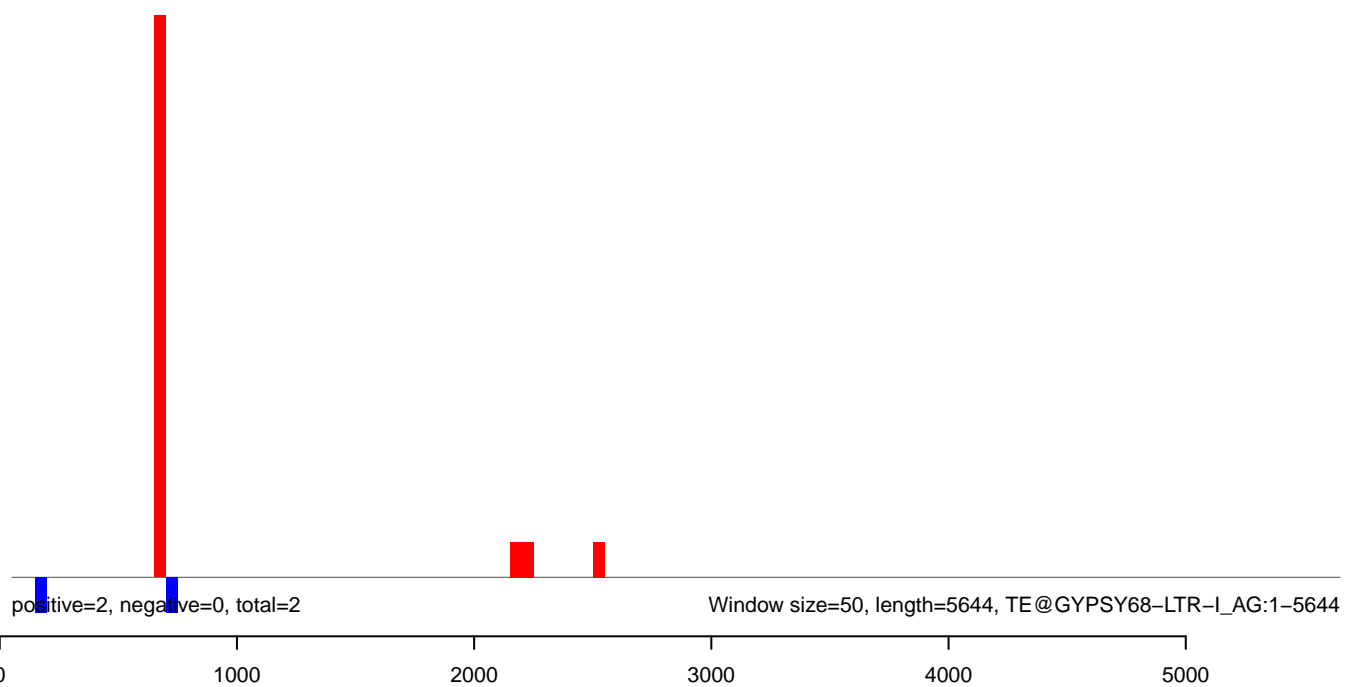
AnGam_Sua5b_cells.18_23.rep



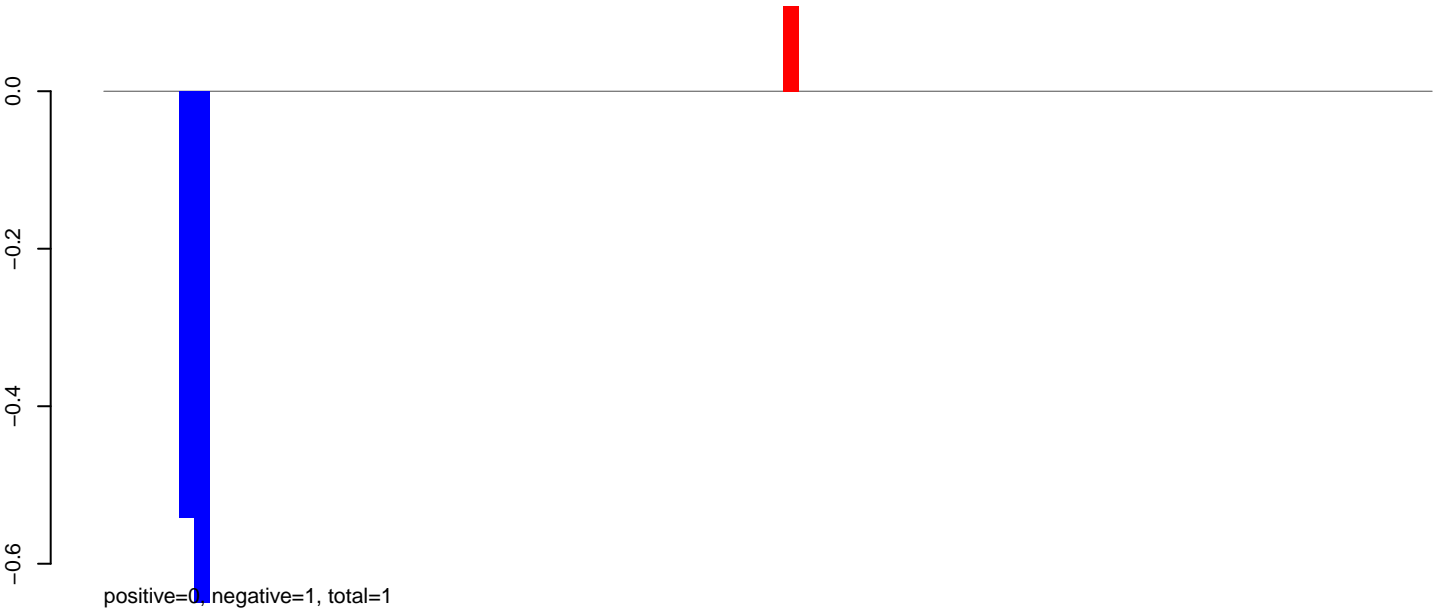
AnGam_Sua5b_cells.24_35.rep



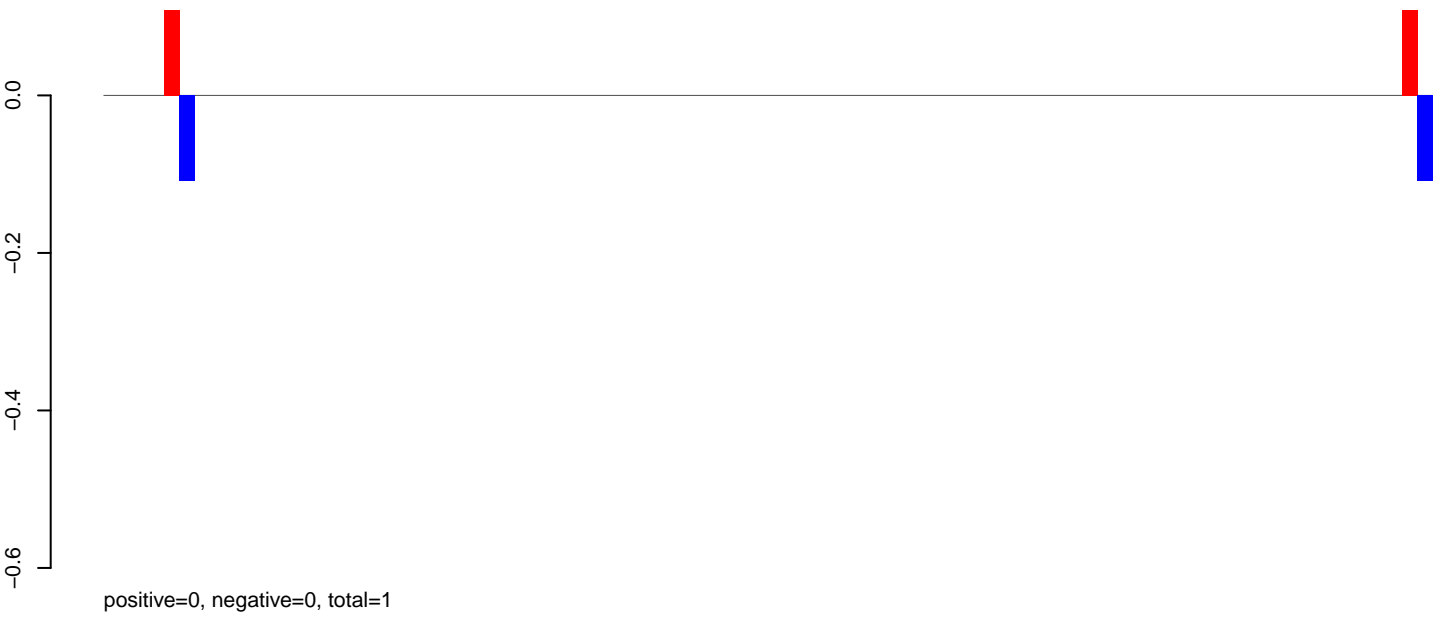
AnGam_Sua5b_cells.rep



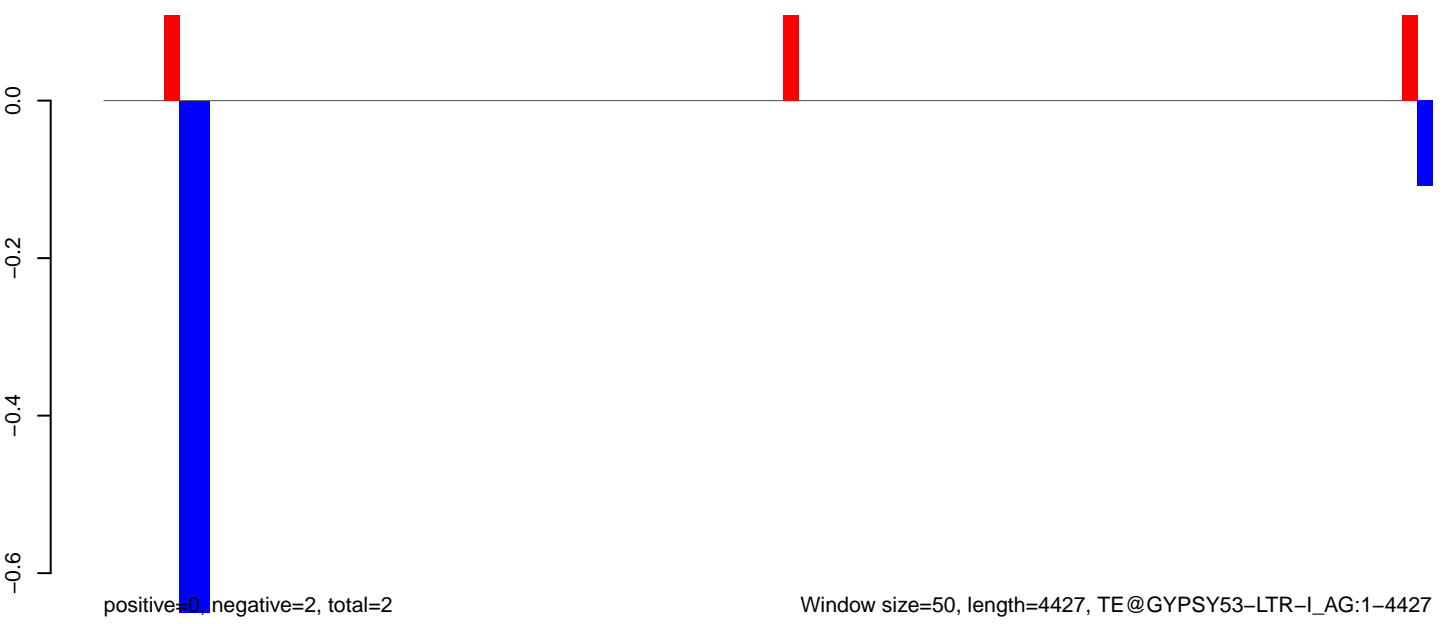
AnGam_Sua5b_cells.18_23.rep



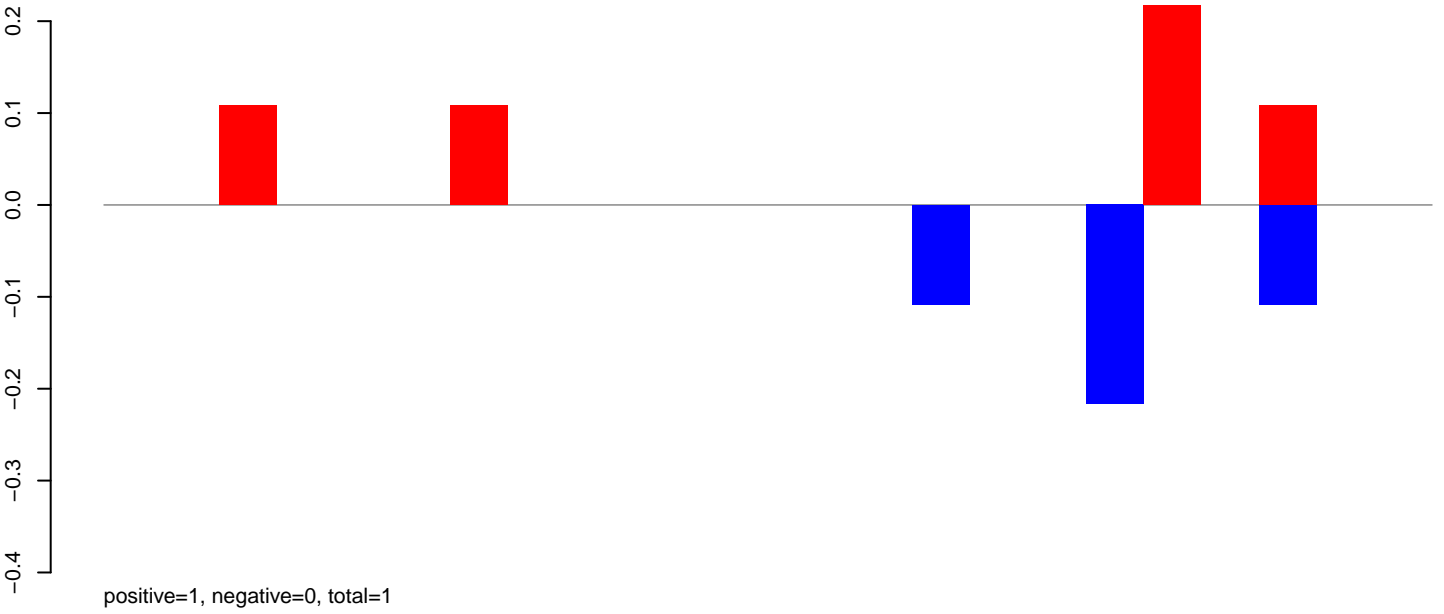
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



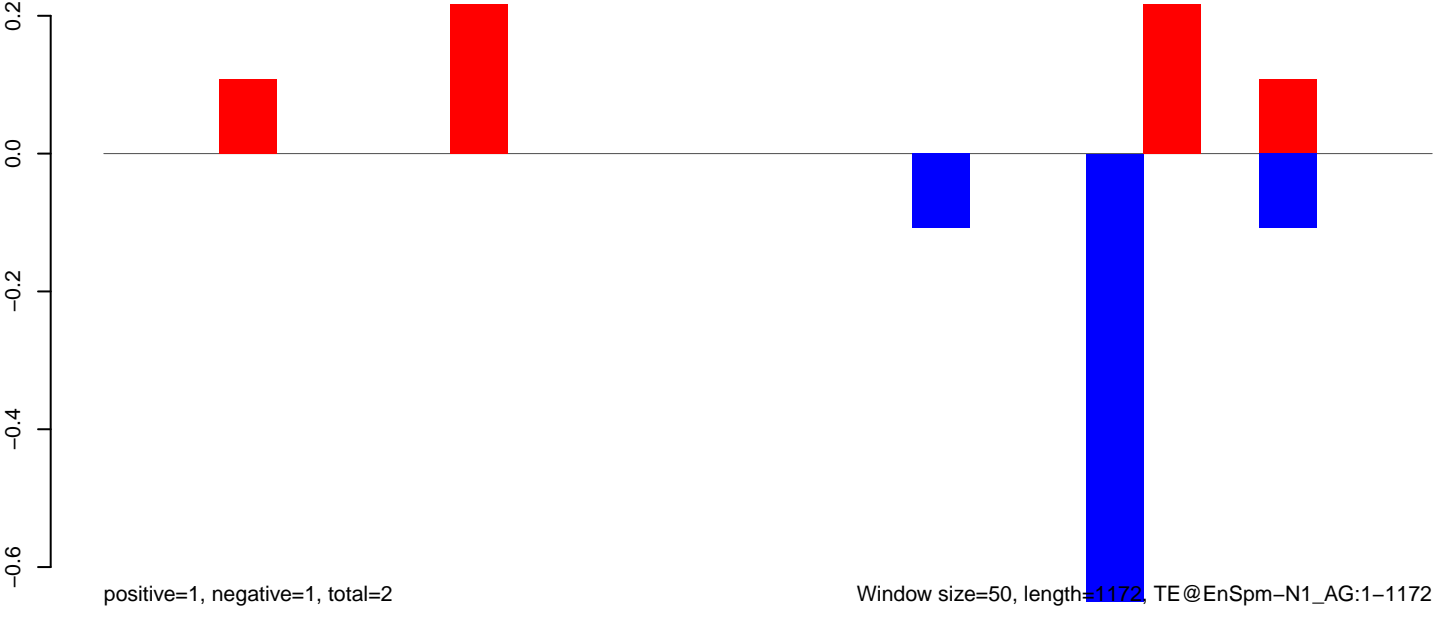
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

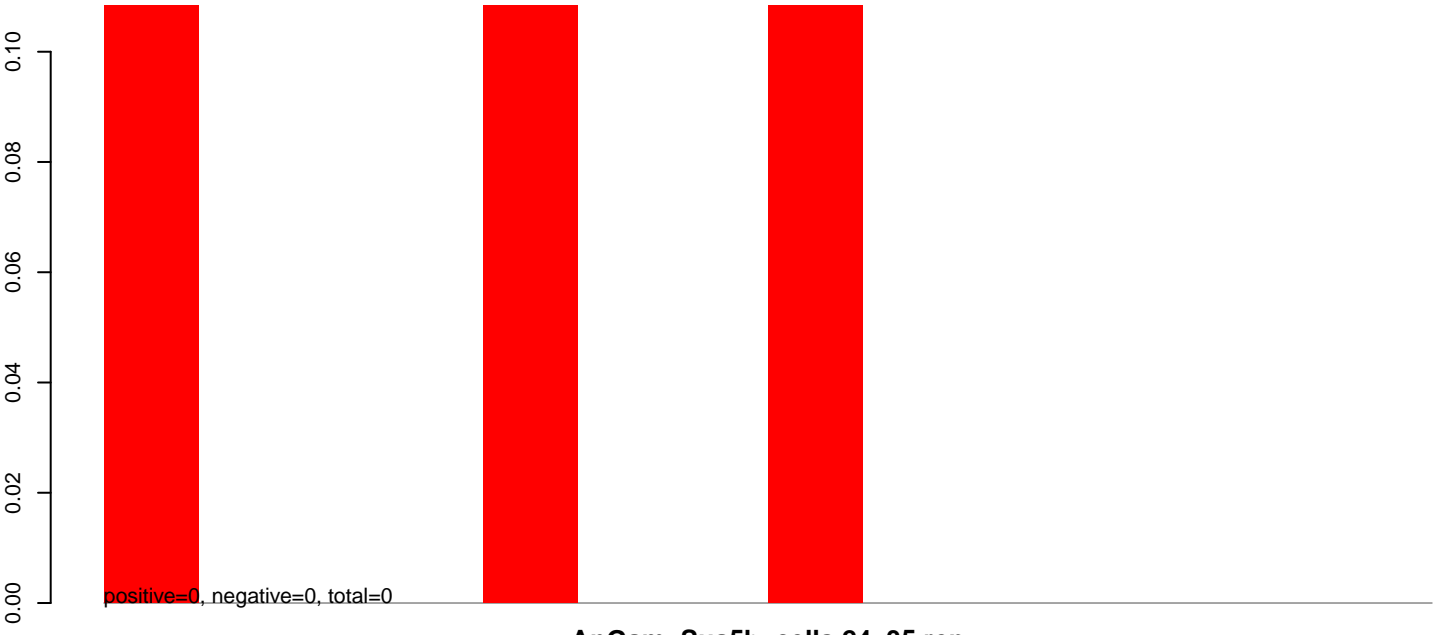


AnGam_Sua5b_cells.rep

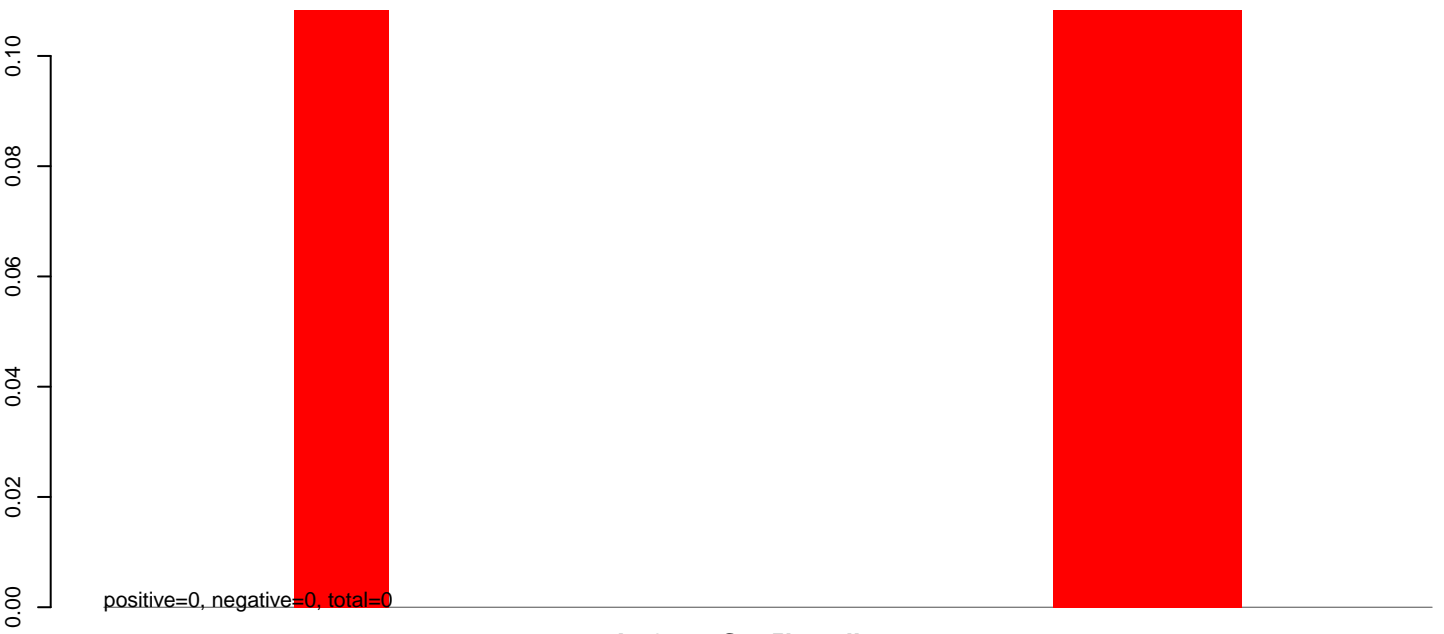


Window size=50, length=1172, TE@EnSpm-N1_AG:1-1172

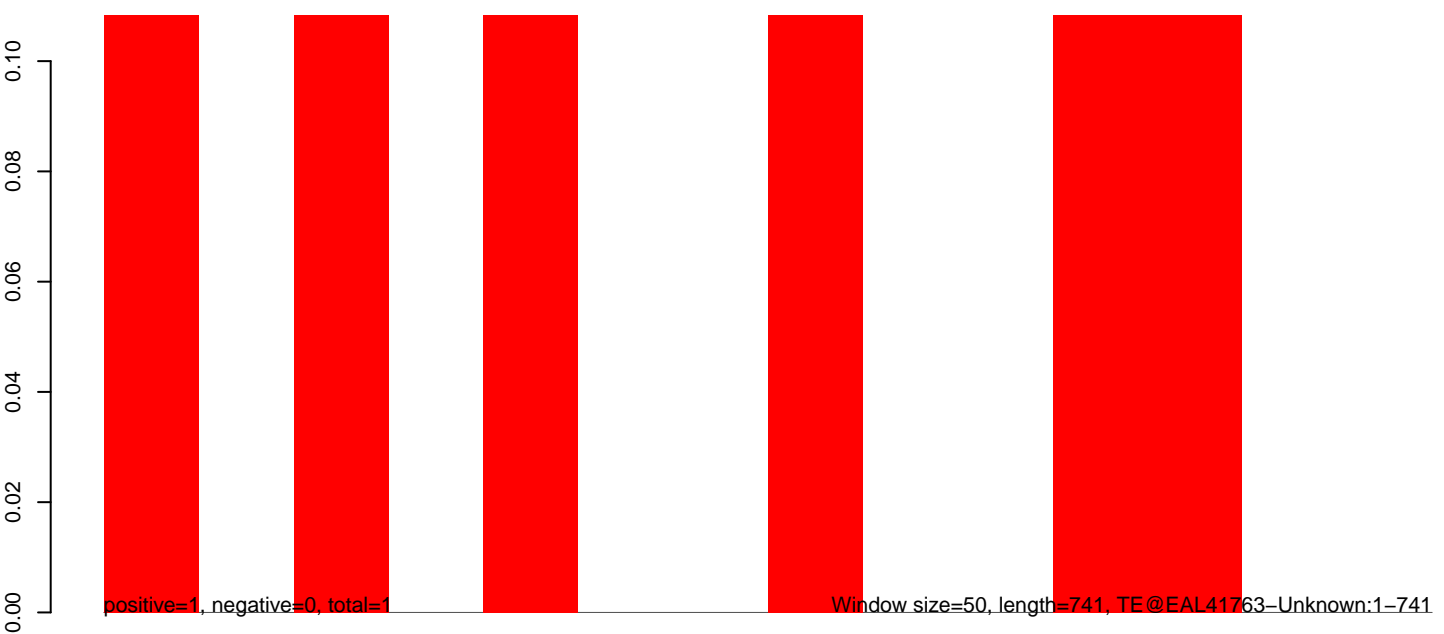
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

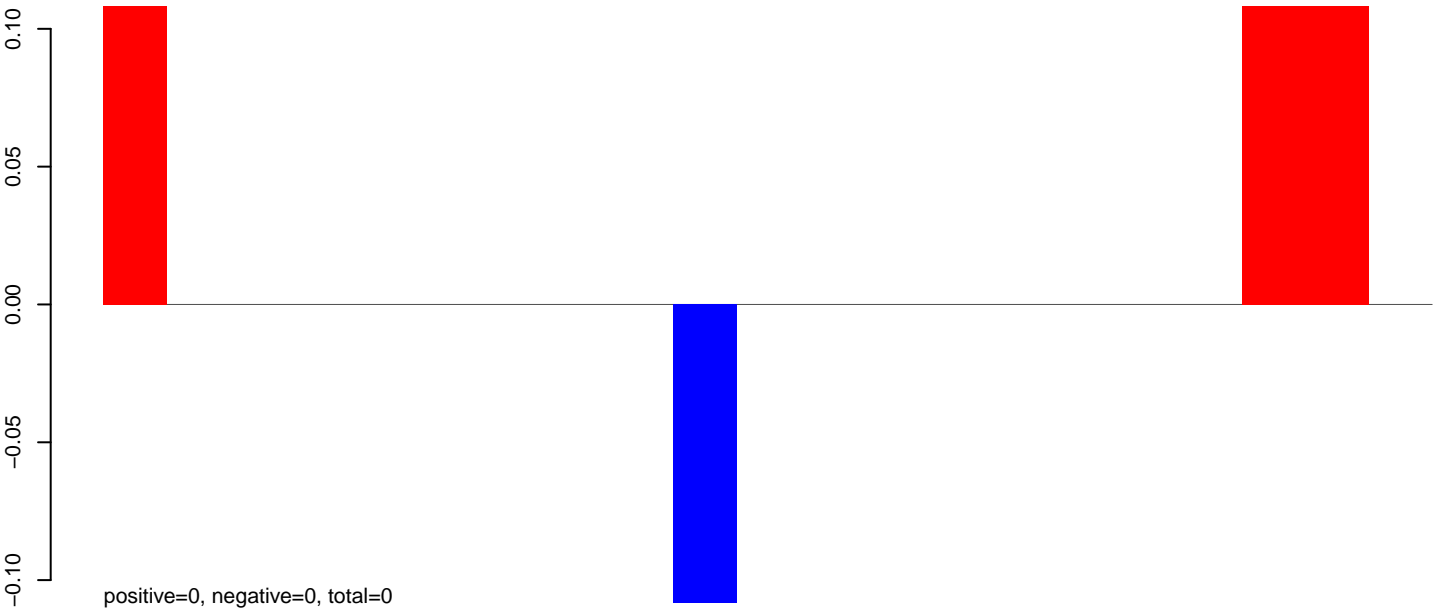


AnGam_Sua5b_cells.rep

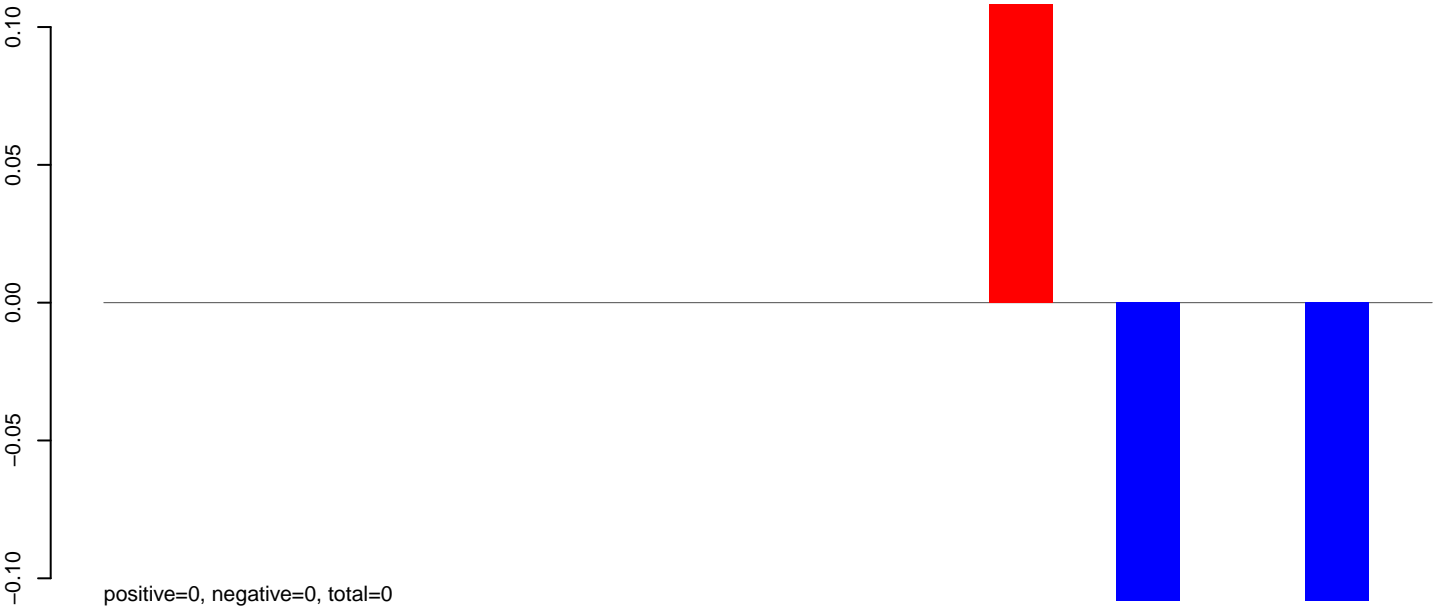


100 200 300 400 500 600 700

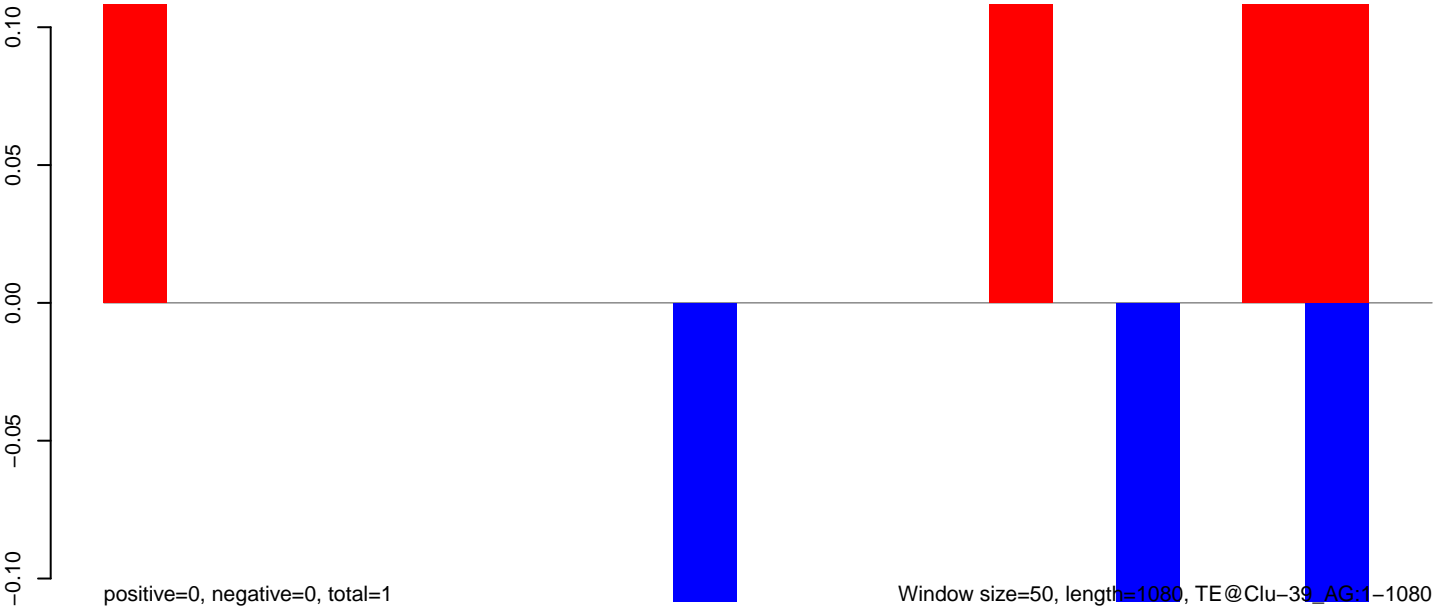
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

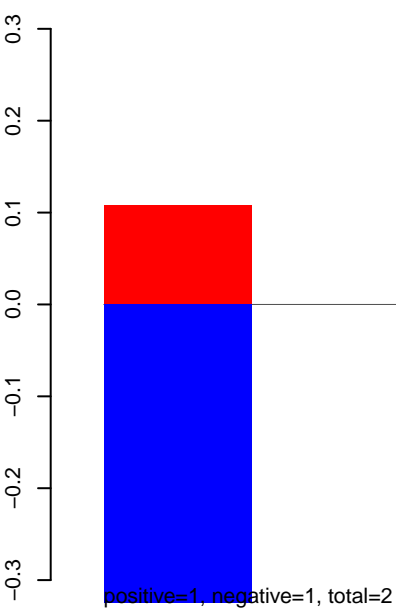


AnGam_Sua5b_cells.rep

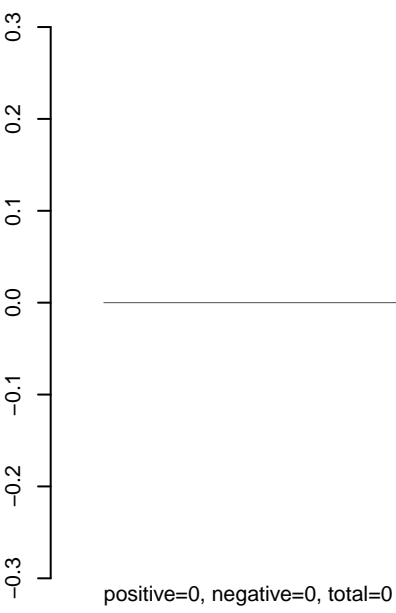


200 400 600 800 1000

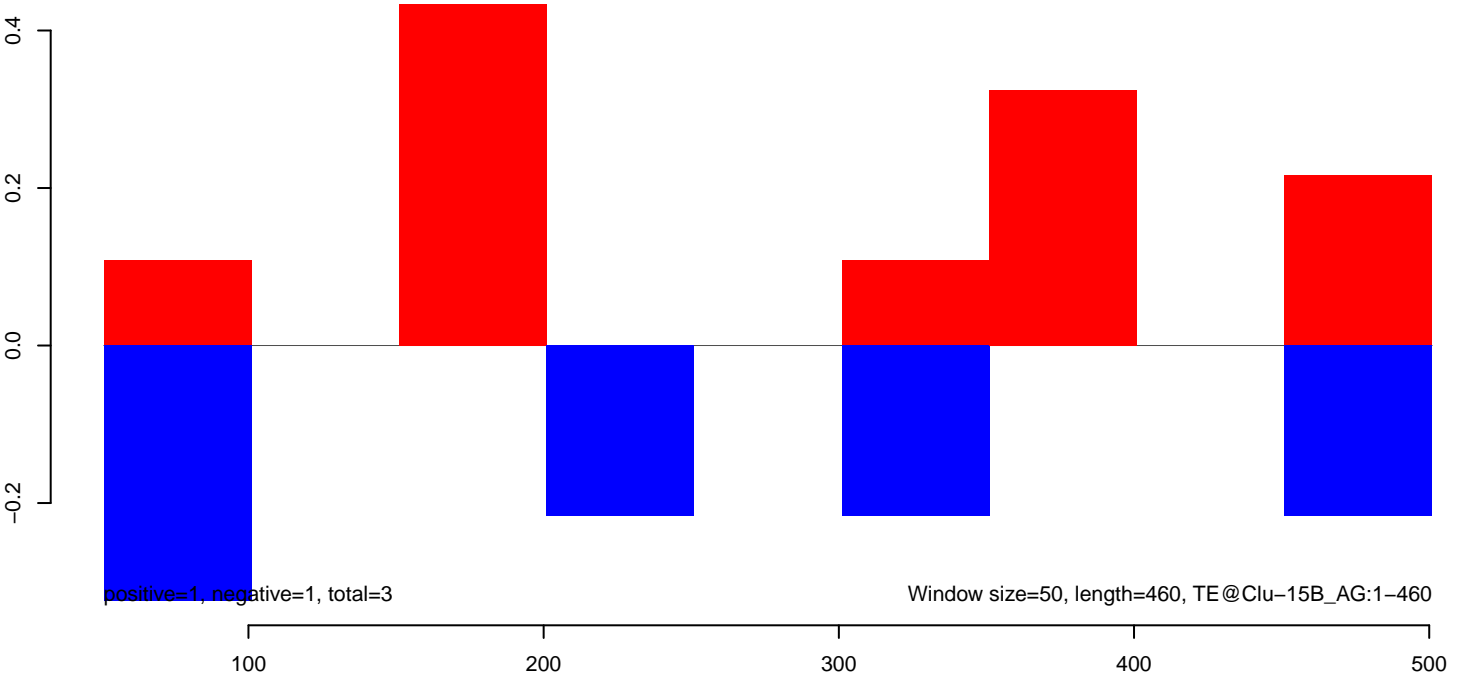
AnGam_Sua5b_cells.18_23.rep



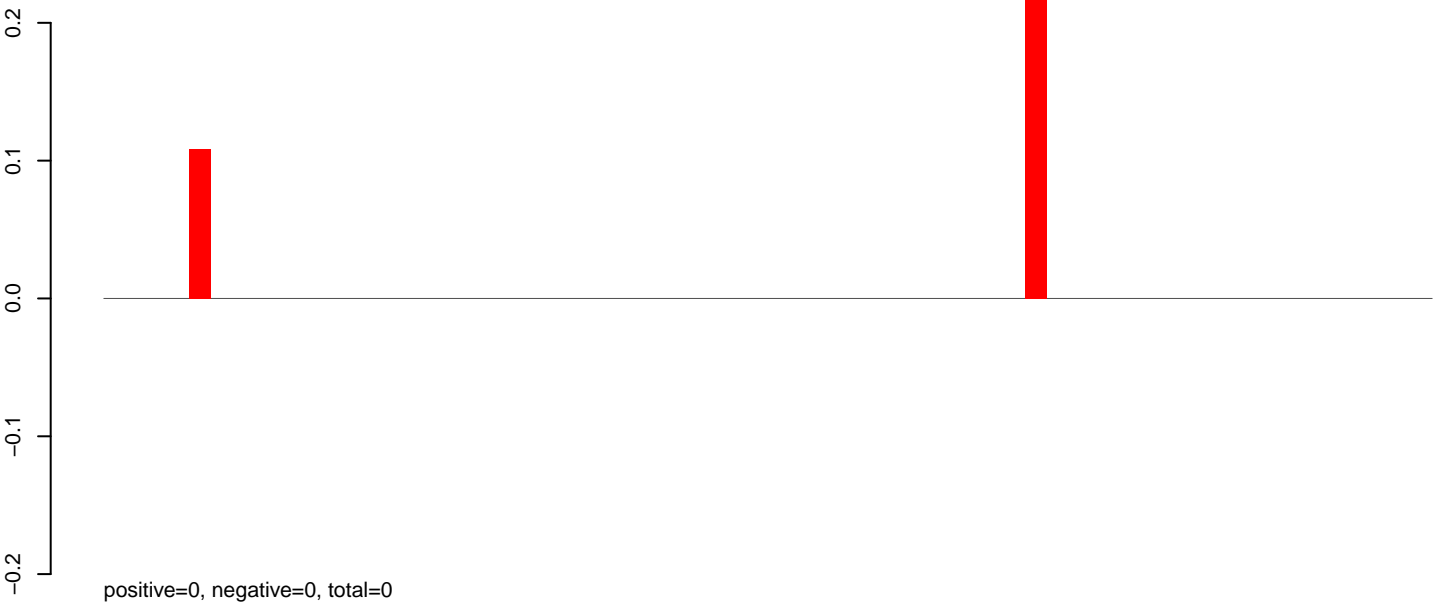
AnGam_Sua5b_cells.24_35.rep



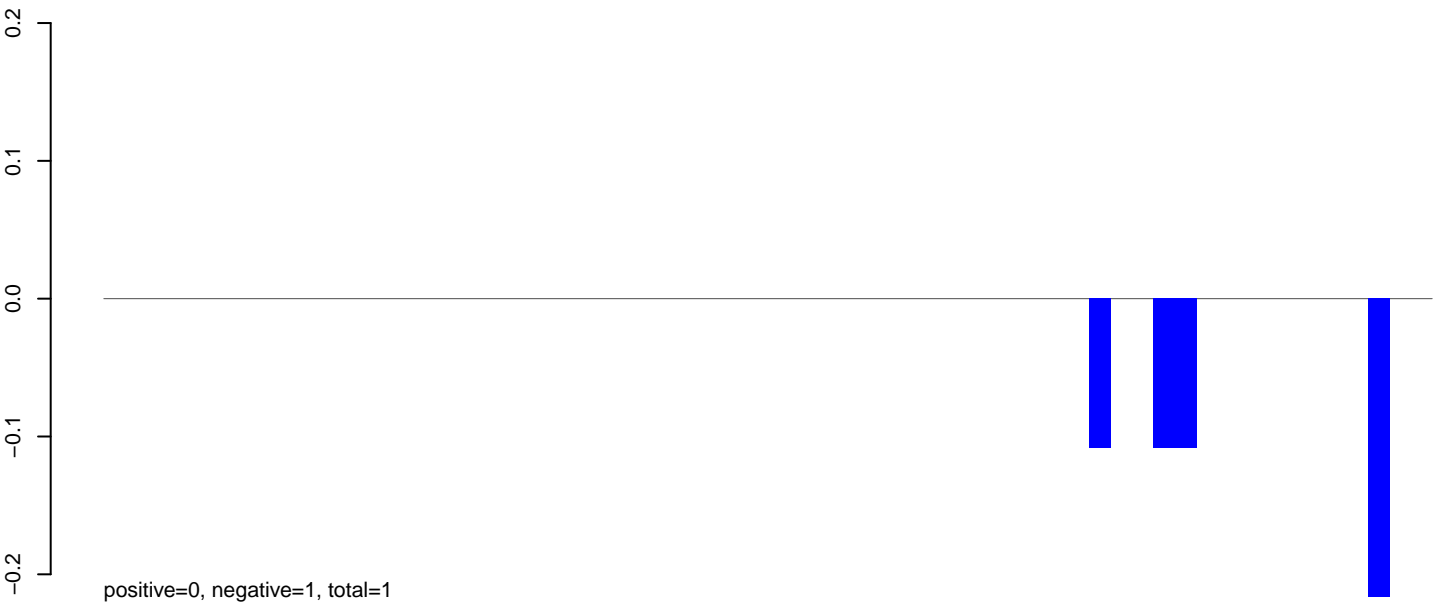
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



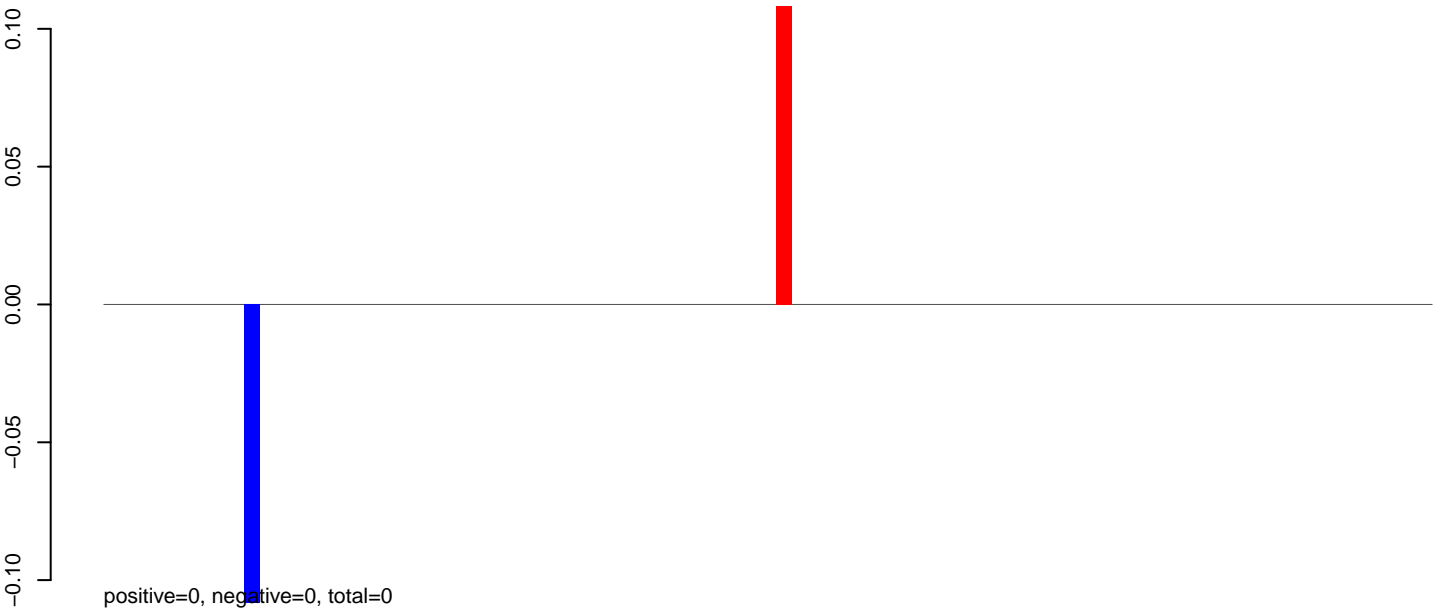
AnGam_Sua5b_cells.24_35.rep



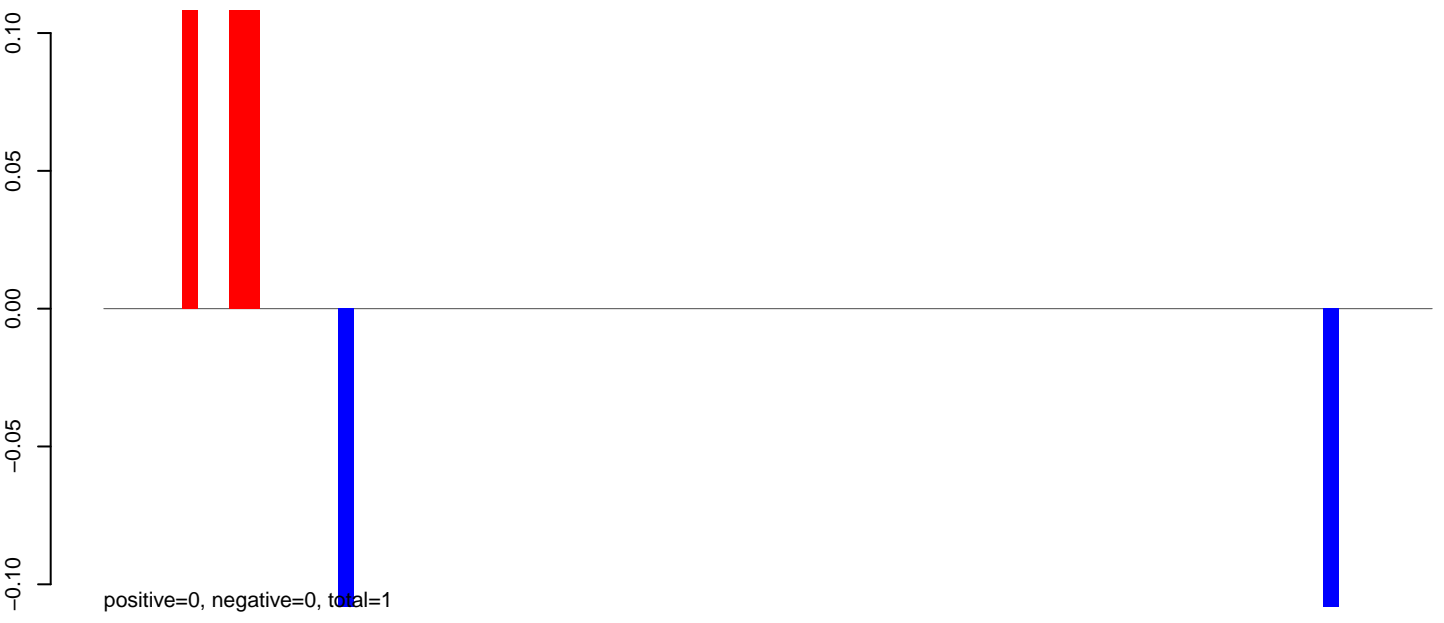
AnGam_Sua5b_cells.rep



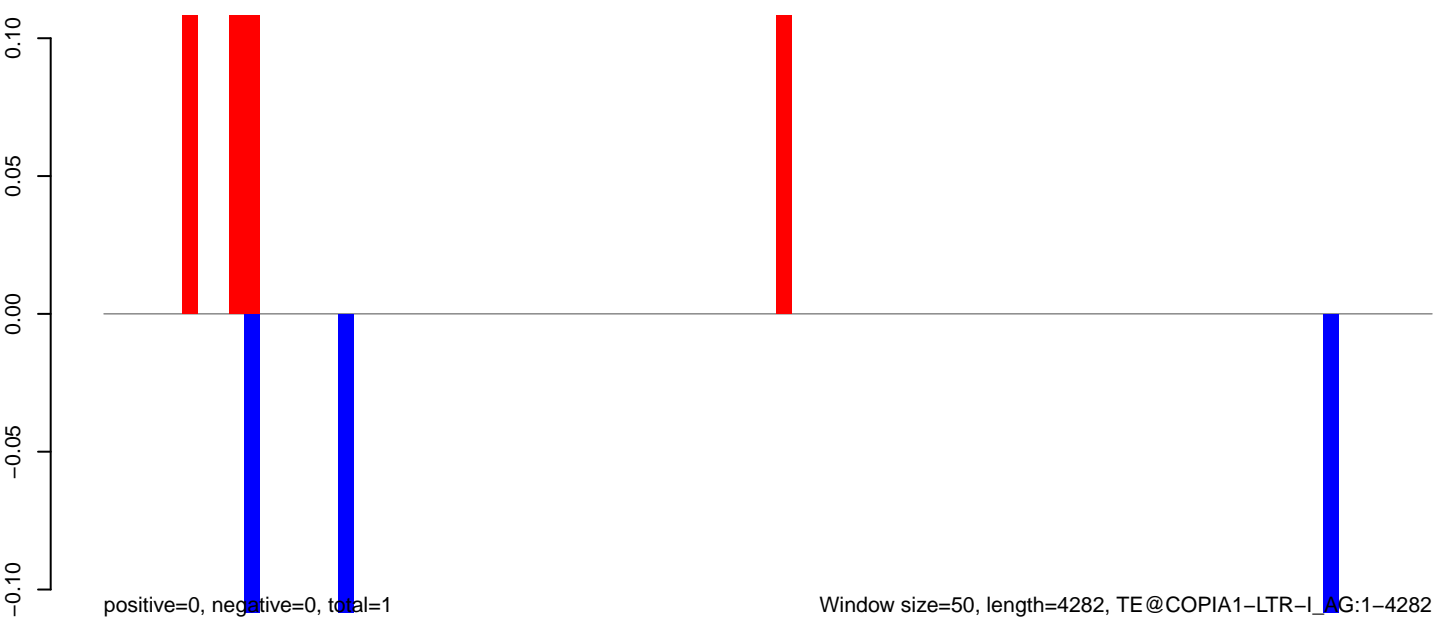
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

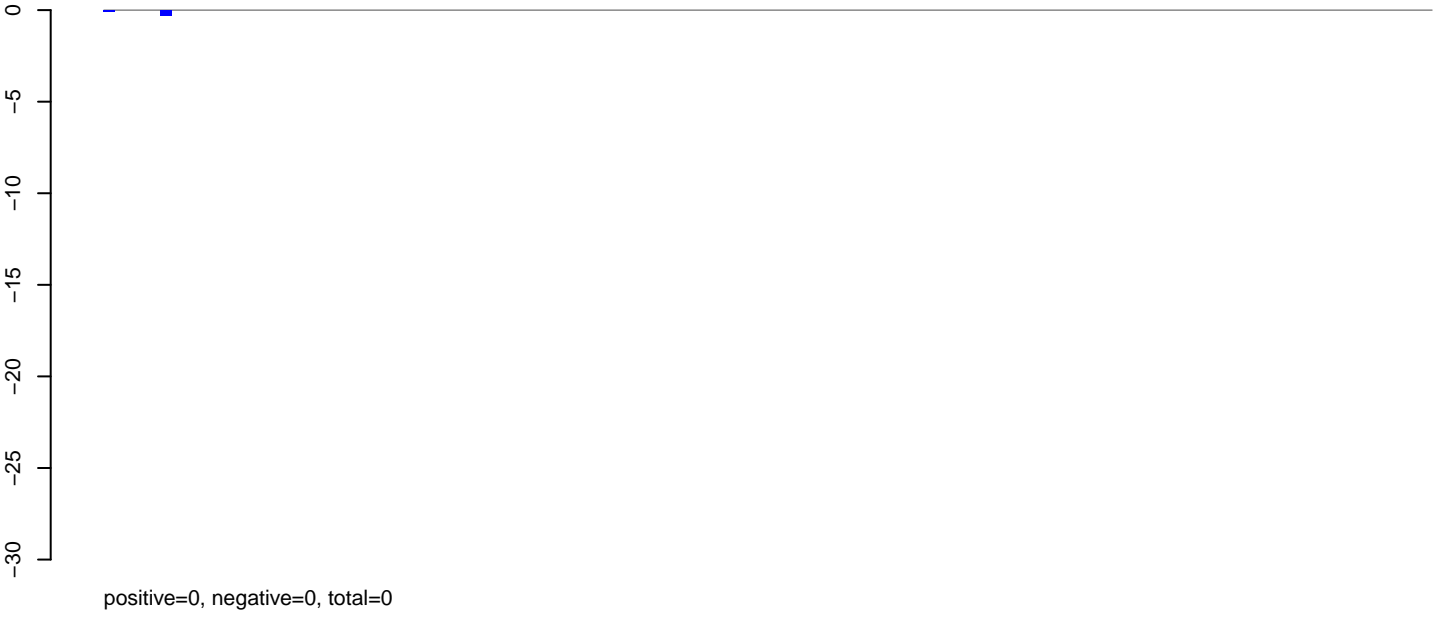


Window size=50, length=4282, TE@COPIA1-LTR-1_AG:1-4282

AnGam_Sua5b_cells.18_23.rep



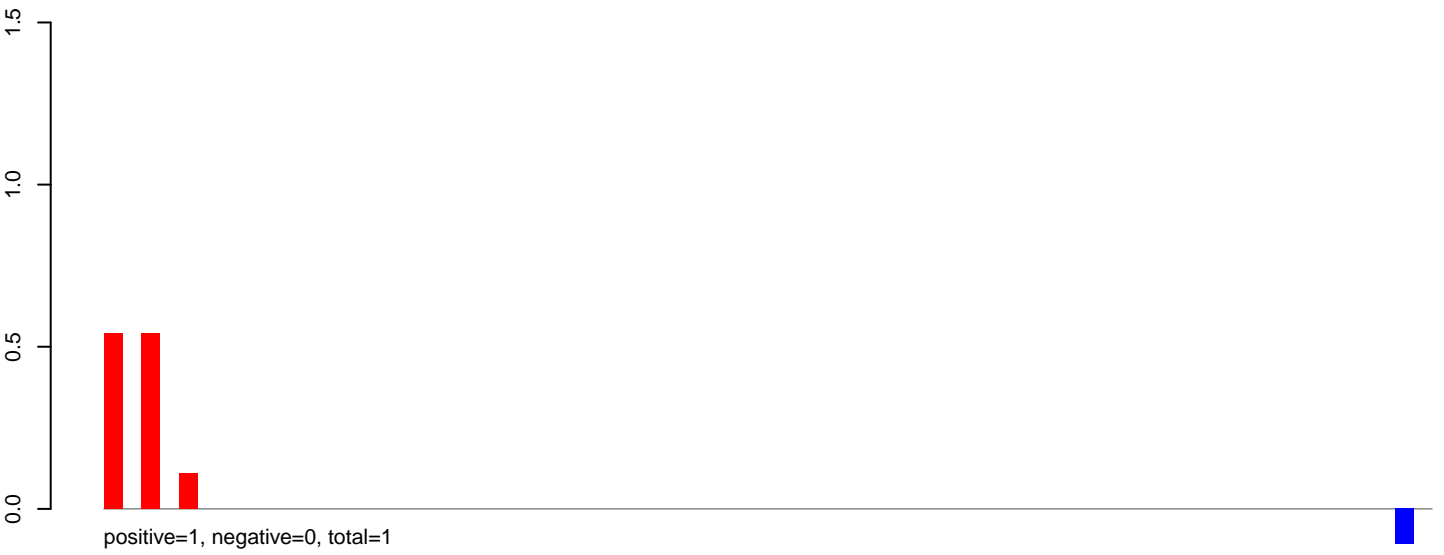
AnGam_Sua5b_cells.24_35.rep



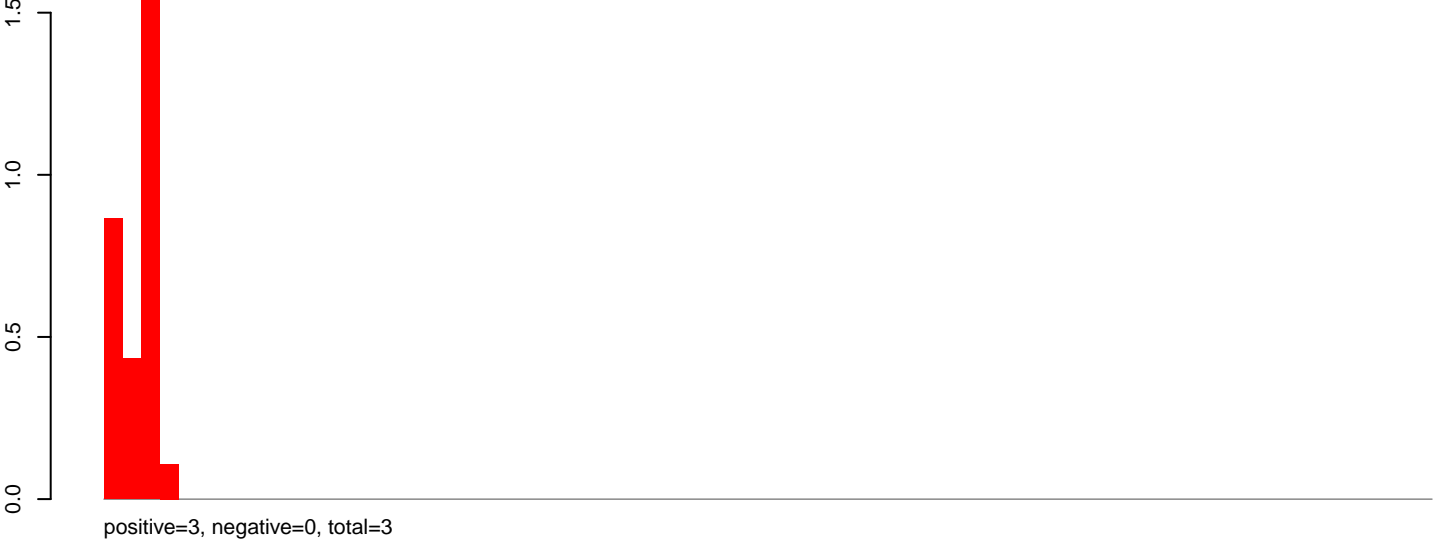
AnGam_Sua5b_cells.rep



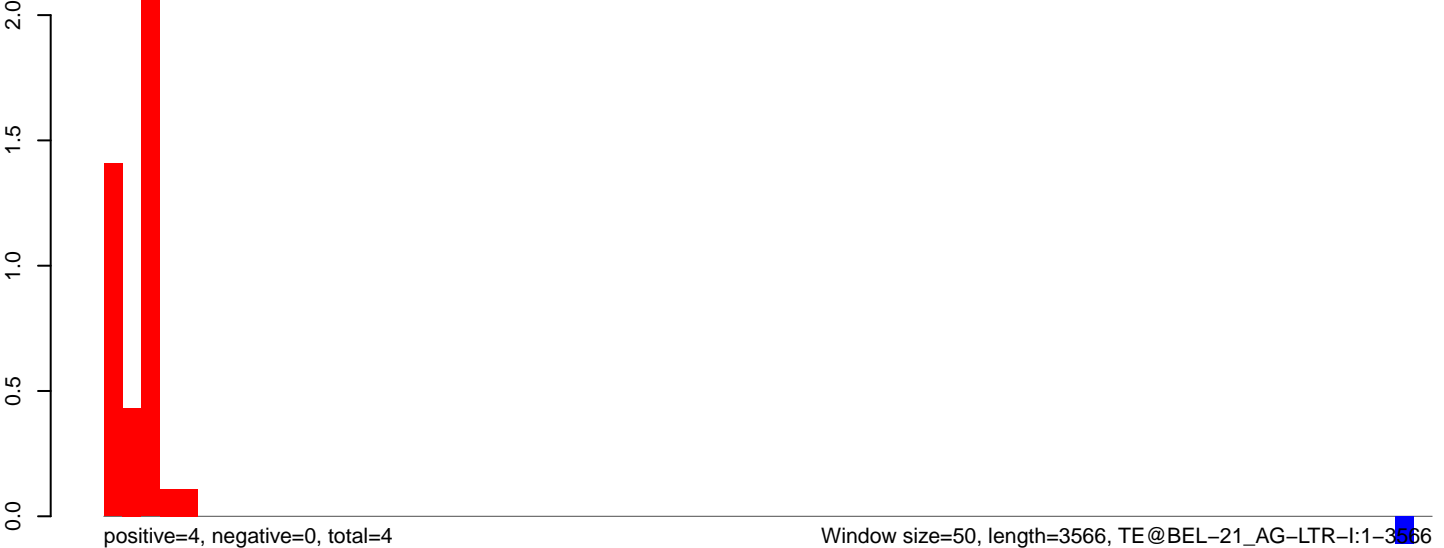
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



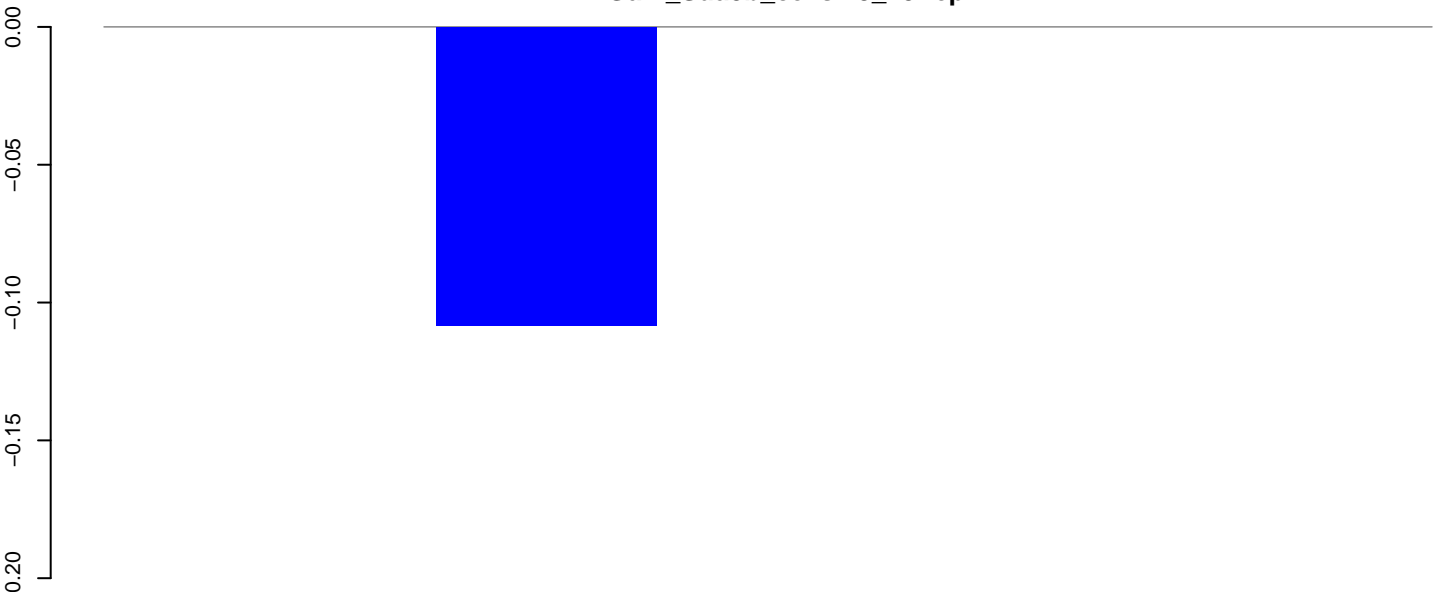
AnGam_Sua5b_cells.rep



Window size=50, length=3566, TE@BEL-21_AG-LTR-I:1-3566

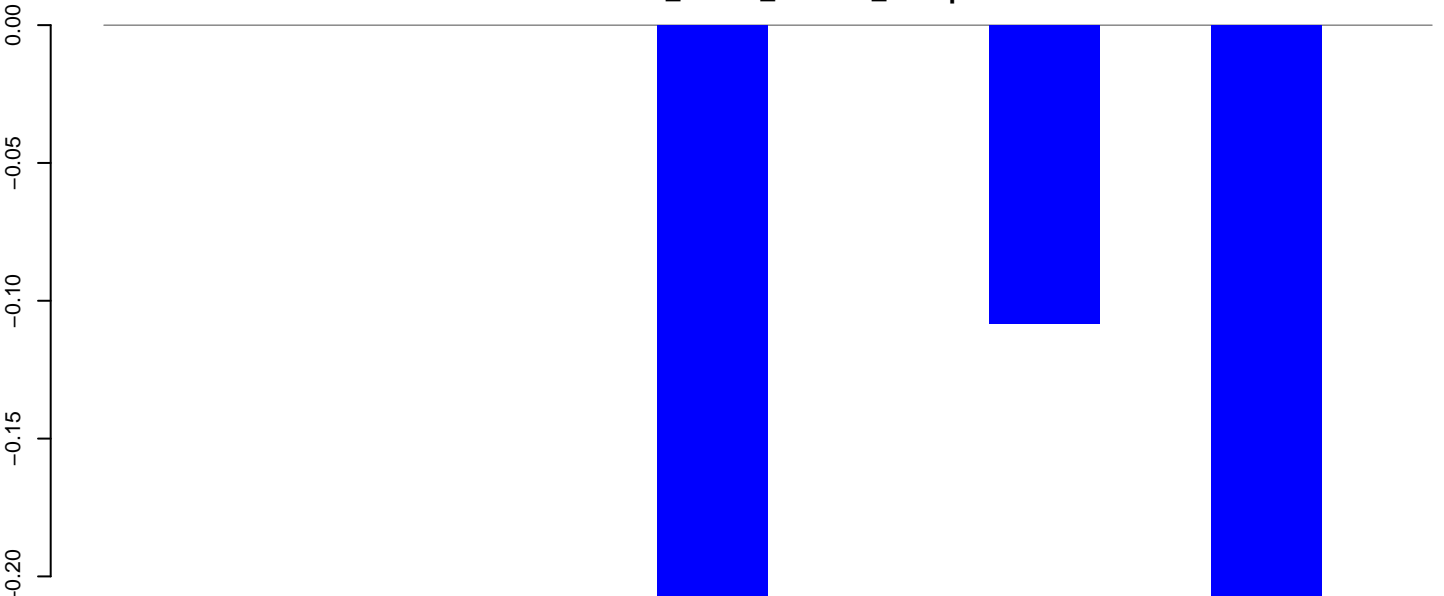
0 500 1000 1500 2000 2500 3000 3500

AnGam_Sua5b_cells.18_23.rep



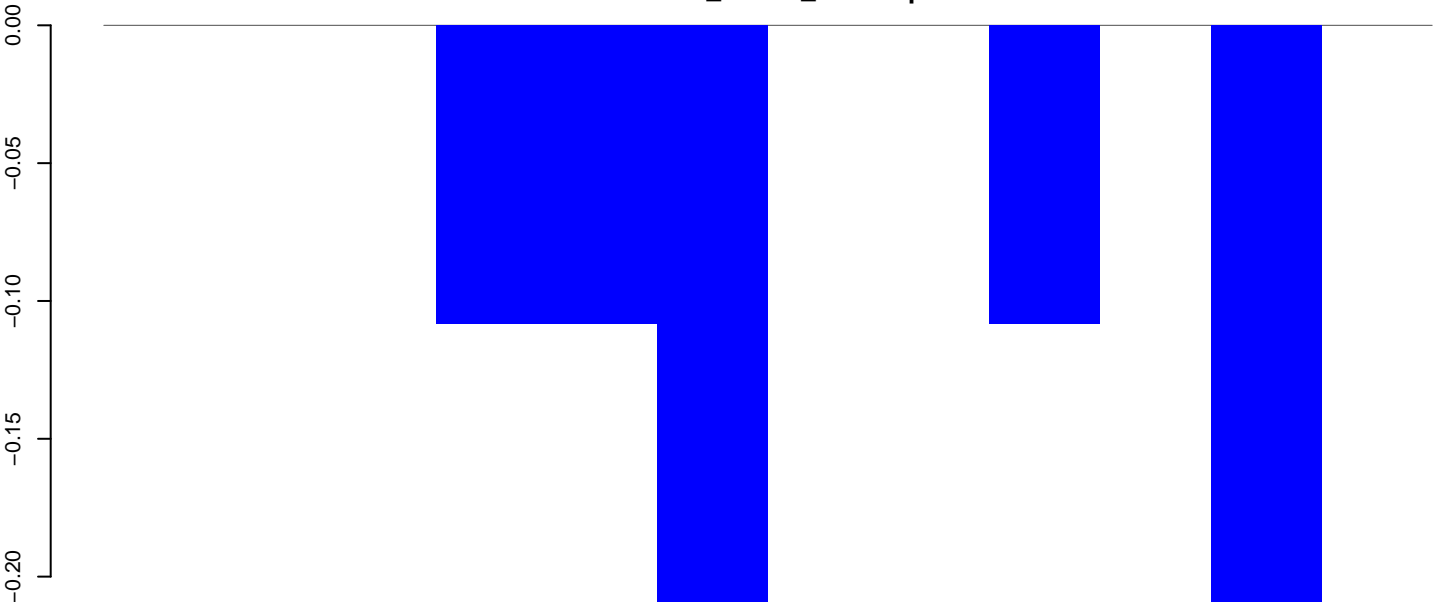
positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep



positive=0, negative=1, total=1

AnGam_Sua5b_cells.rep

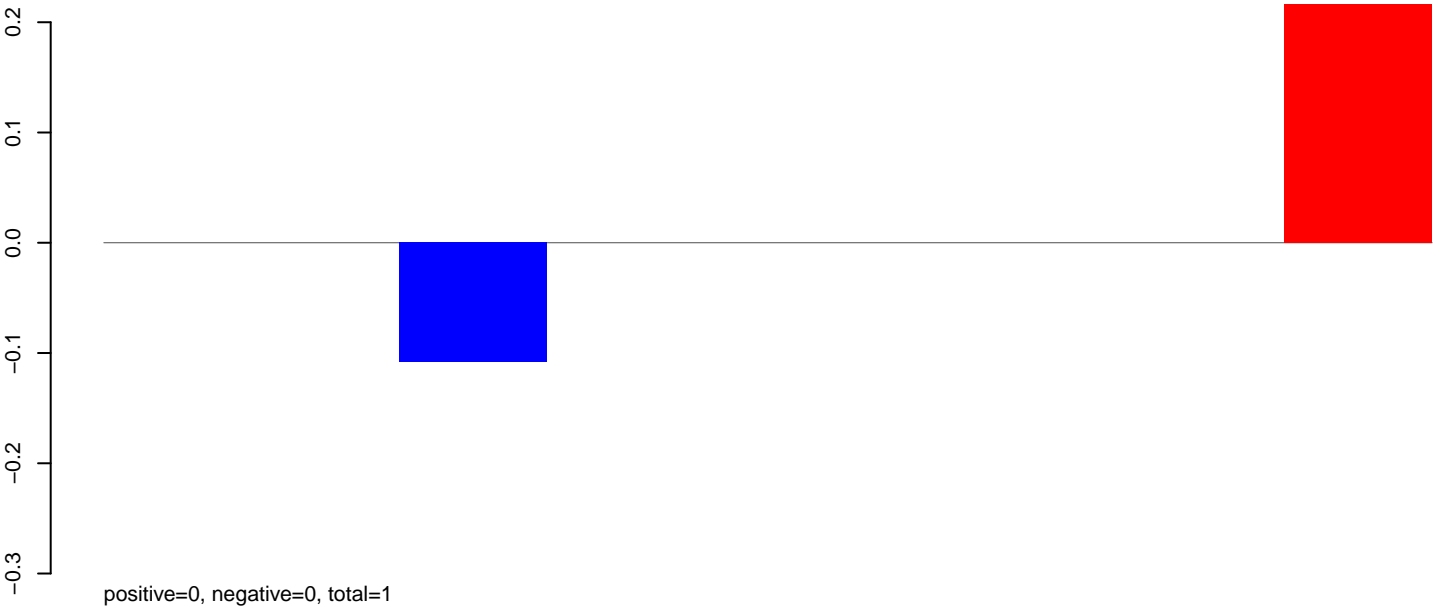


positive=0, negative=1, total=1

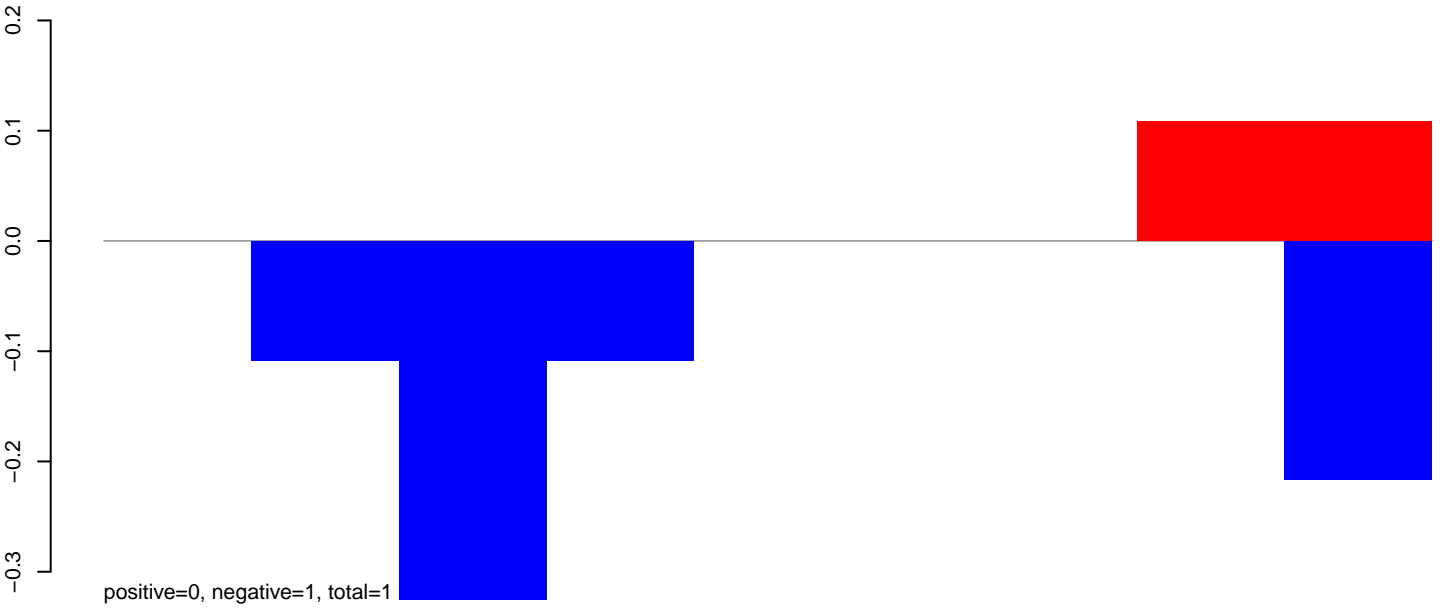
Window size=50, length=649, TE@TF003655-m4bp_Ele3:1-649

100 200 300 400 500 600

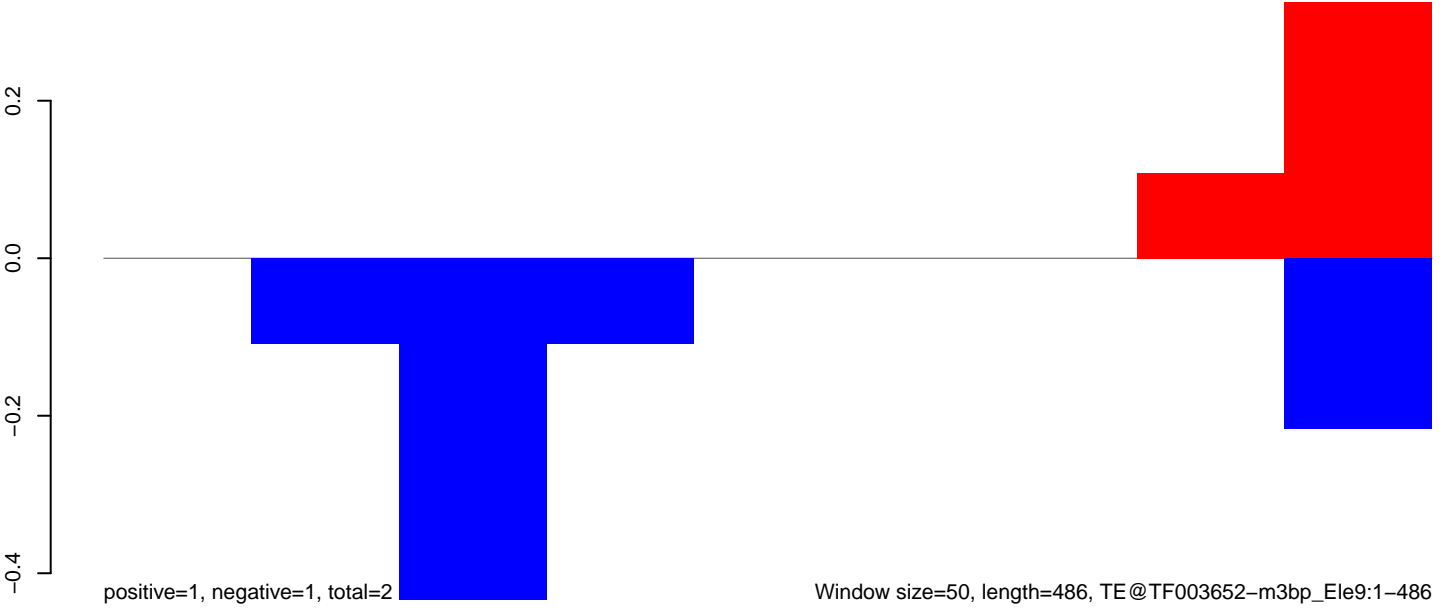
AnGam_Sua5b_cells.18_23.rep



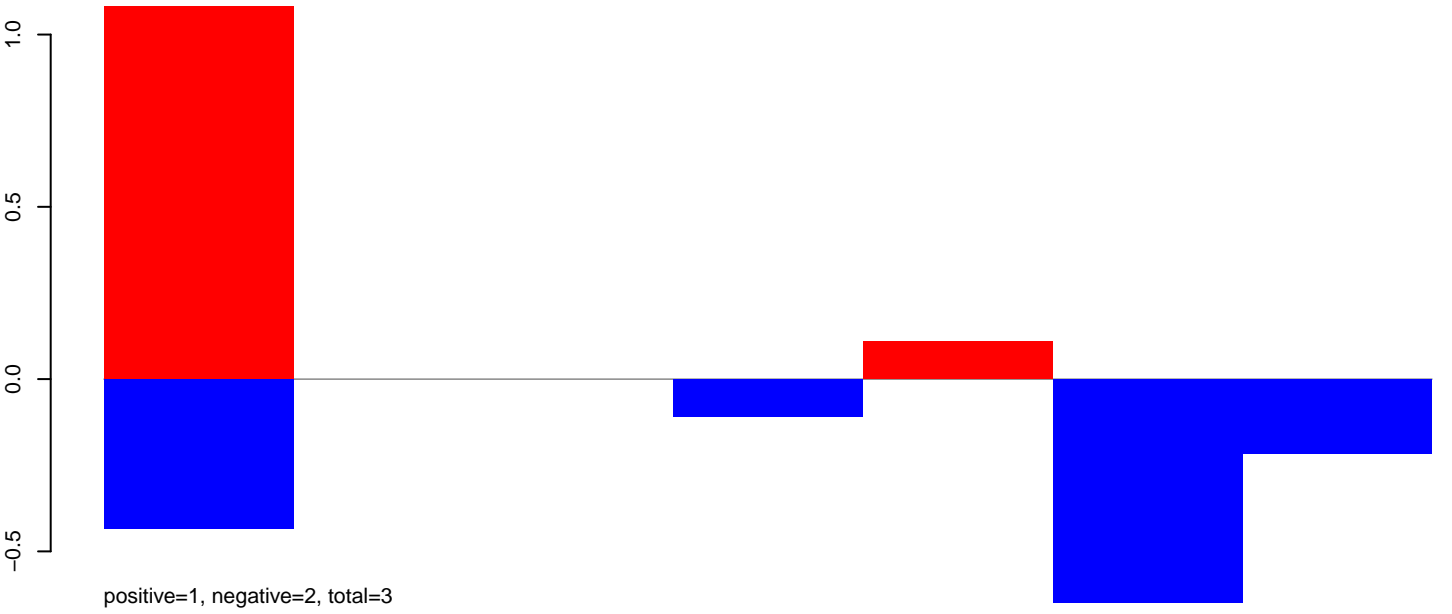
AnGam_Sua5b_cells.24_35.rep



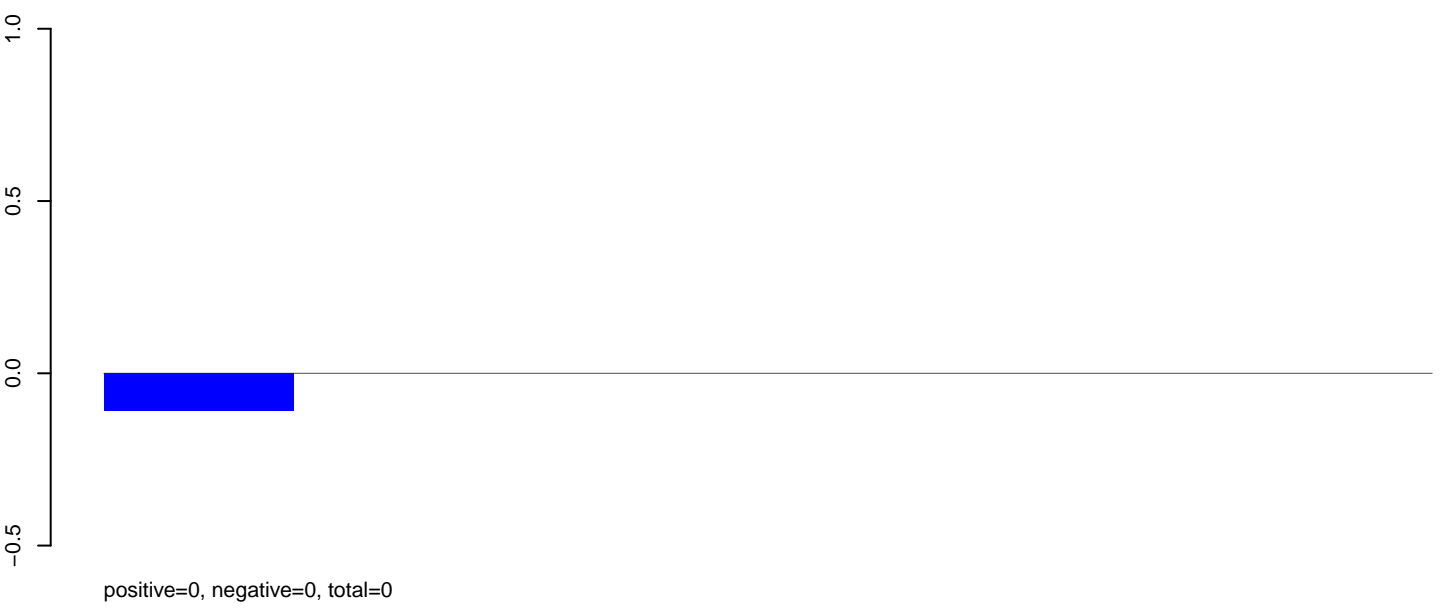
AnGam_Sua5b_cells.rep



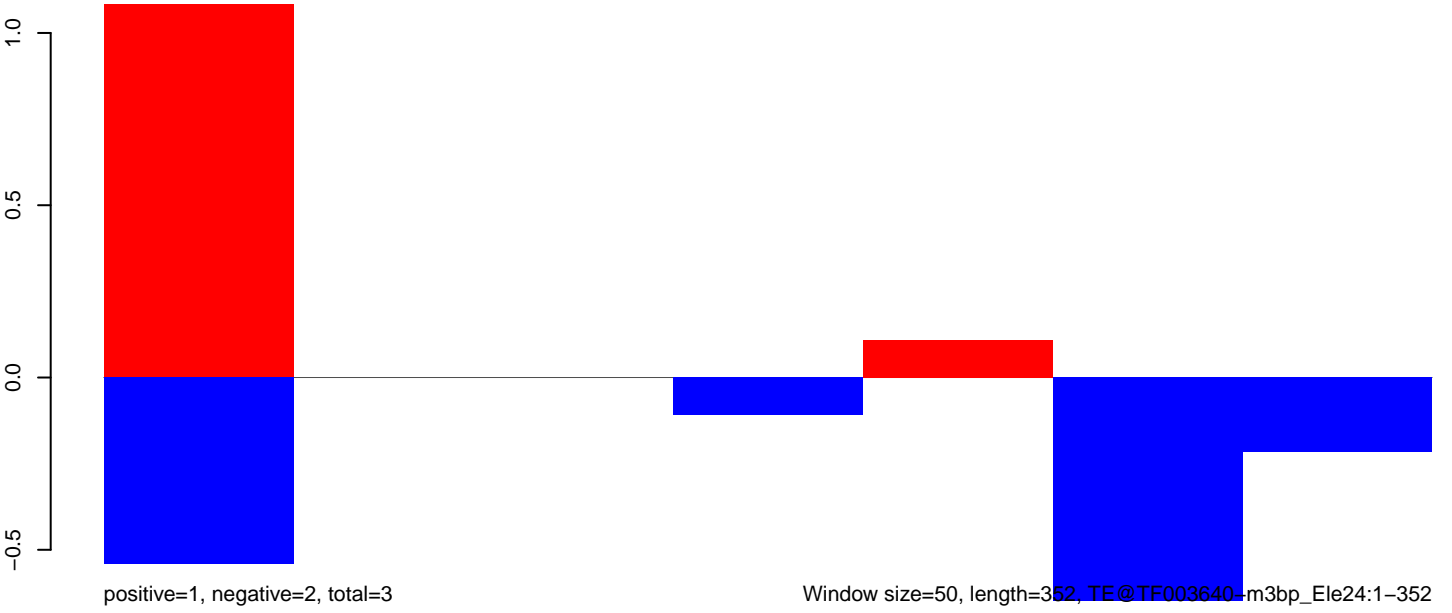
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



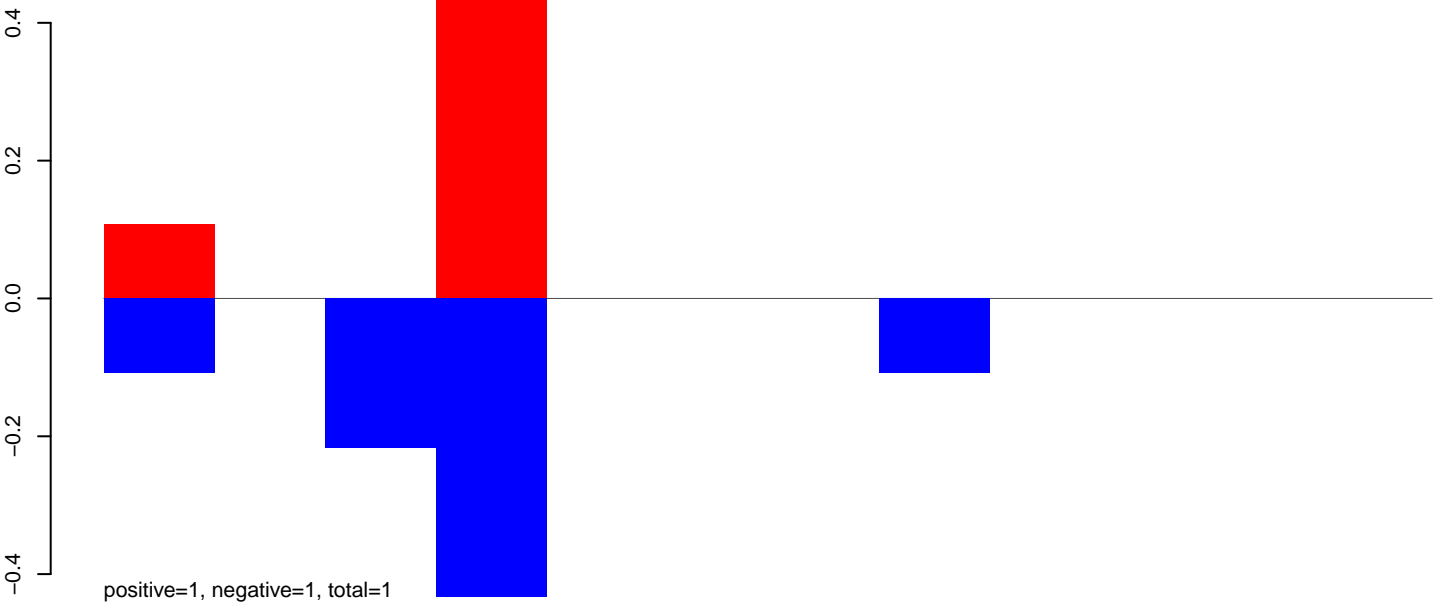
AnGam_Sua5b_cells.rep



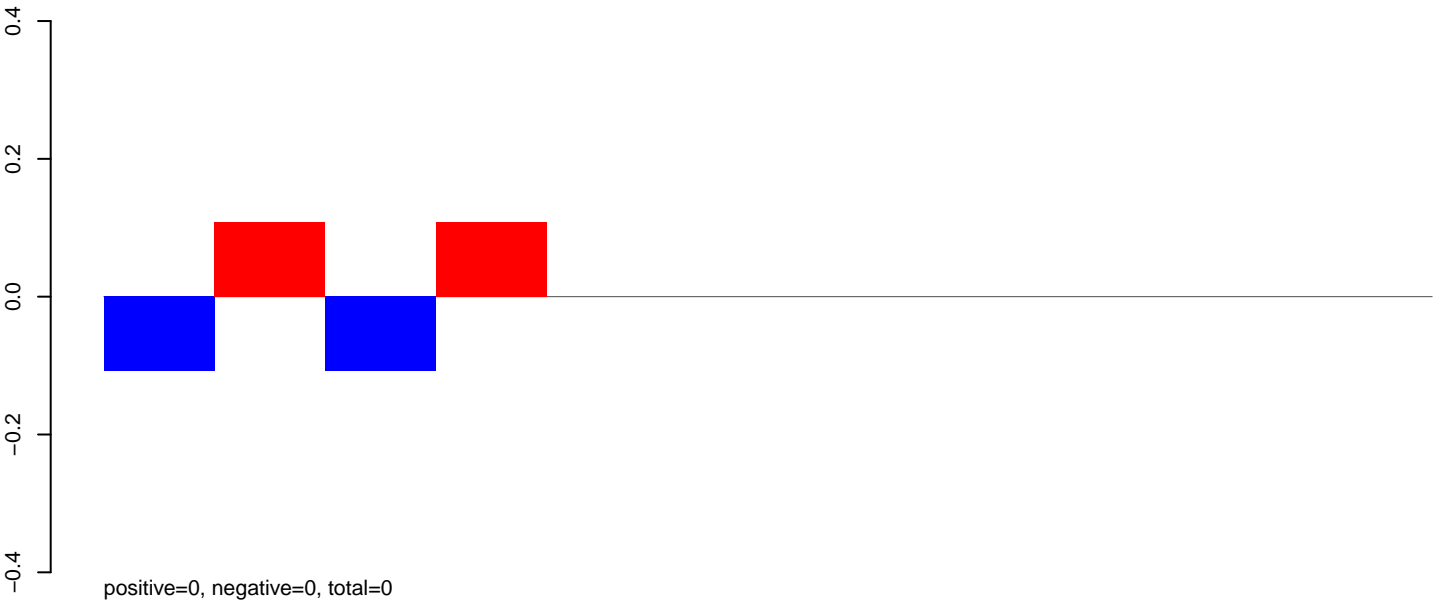
Window size=50, length=352, TE@TF003640-m3bp_Ele24:1-352

50 100 150 200 250 300 350 400

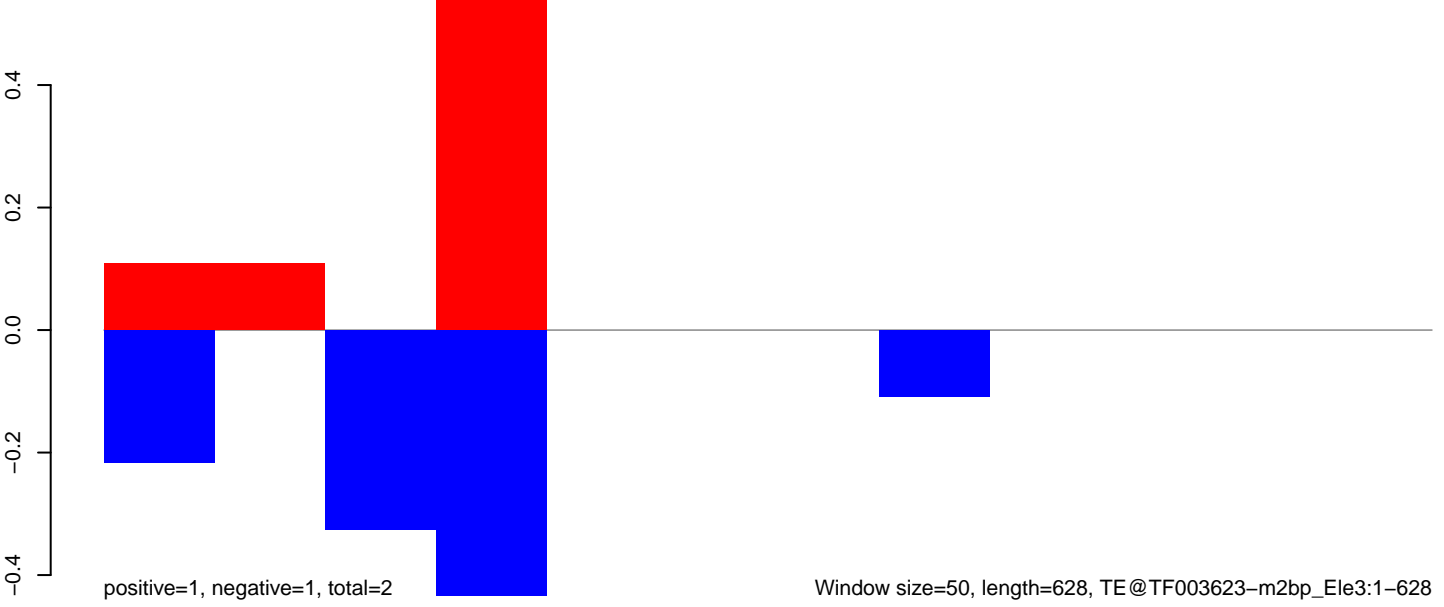
AnGam_Sua5b_cells.18_23.rep



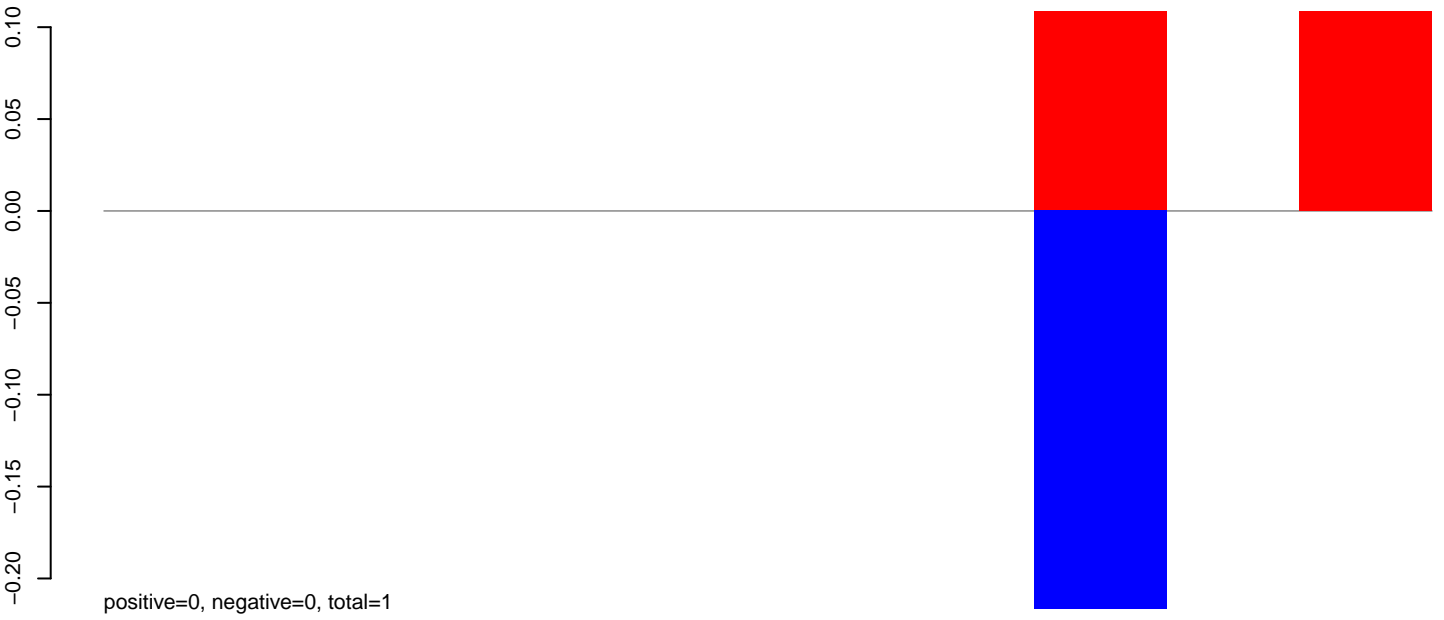
AnGam_Sua5b_cells.24_35.rep



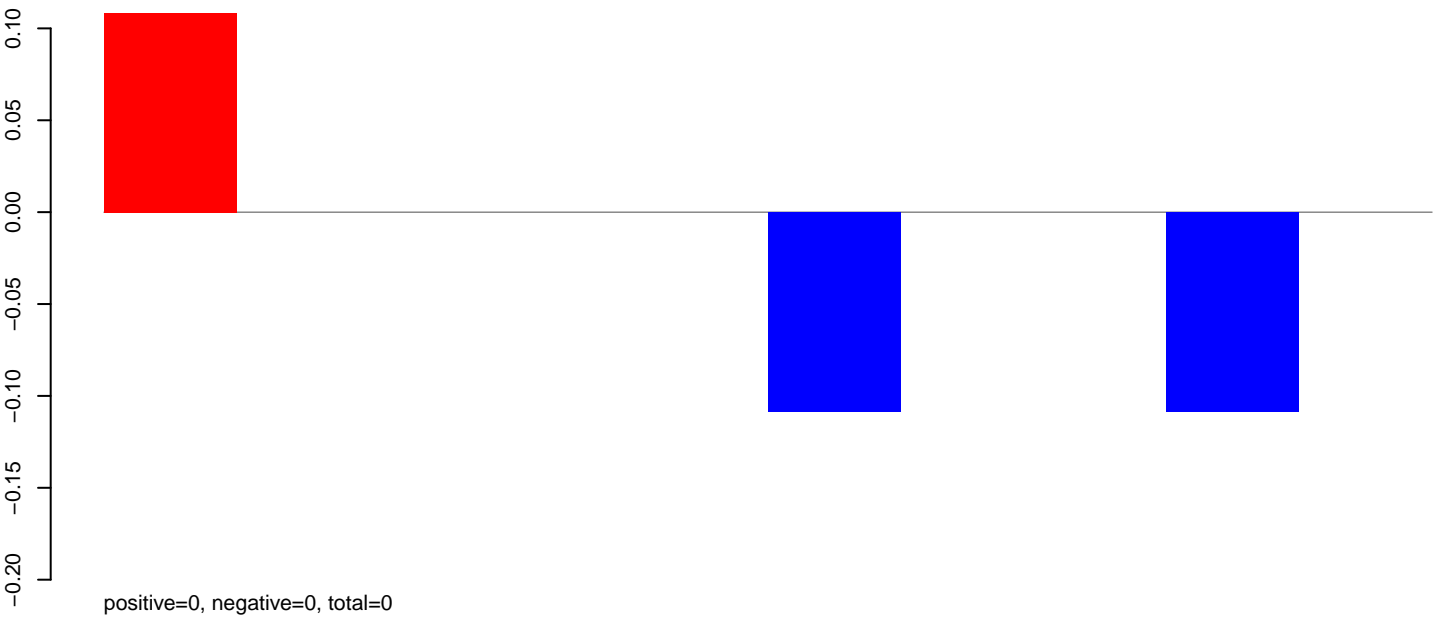
AnGam_Sua5b_cells.rep



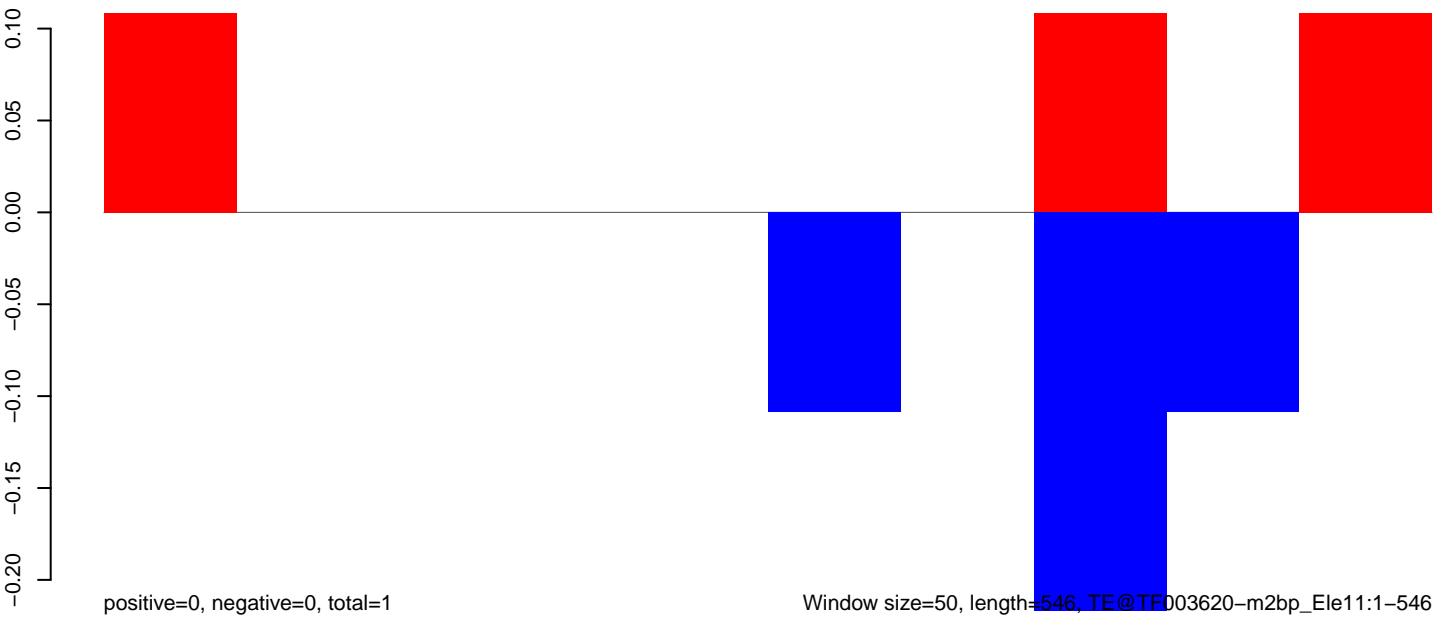
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



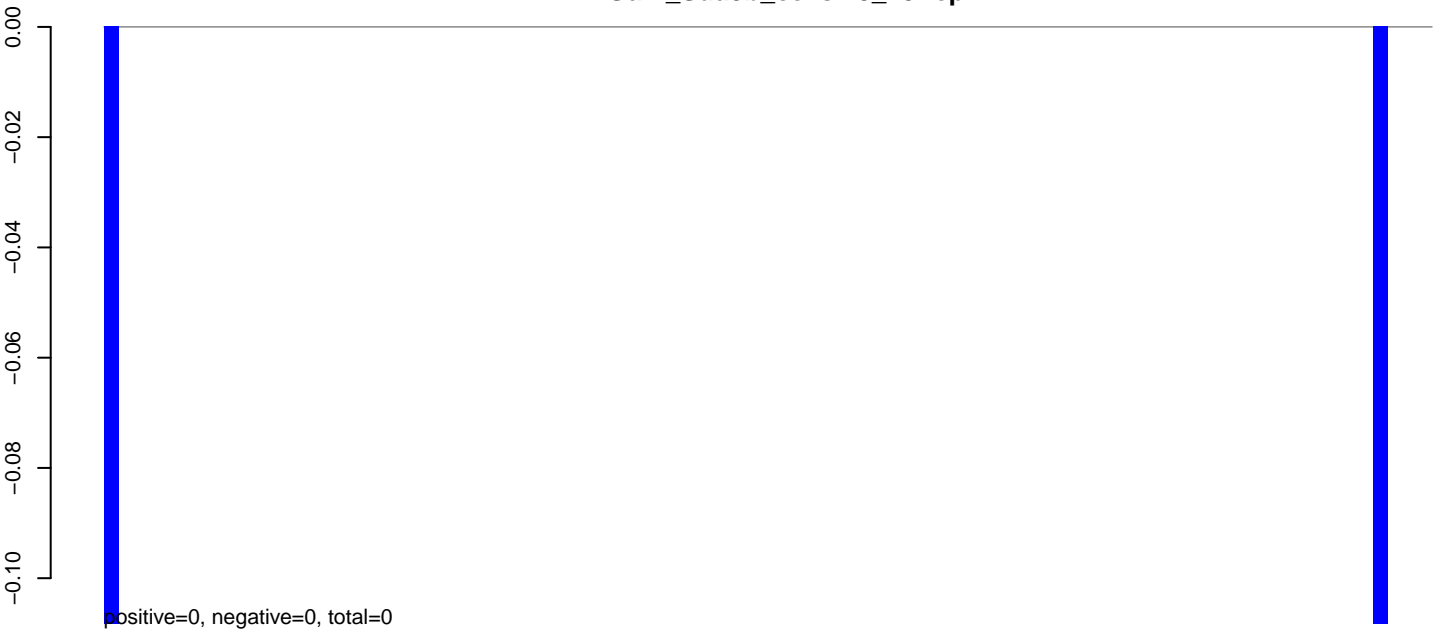
AnGam_Sua5b_cells.rep



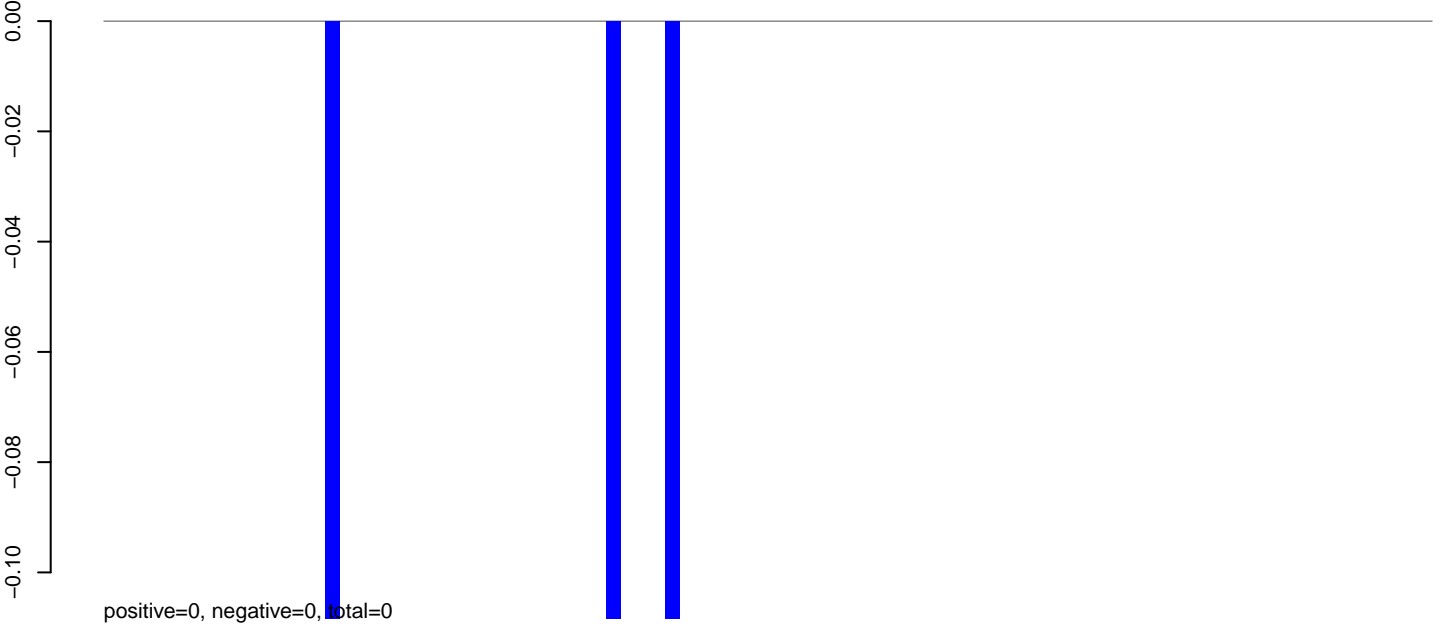
Window size=50, length=546, TE@TE003620-m2bp_Ele11:1-546

100 200 300 400 500

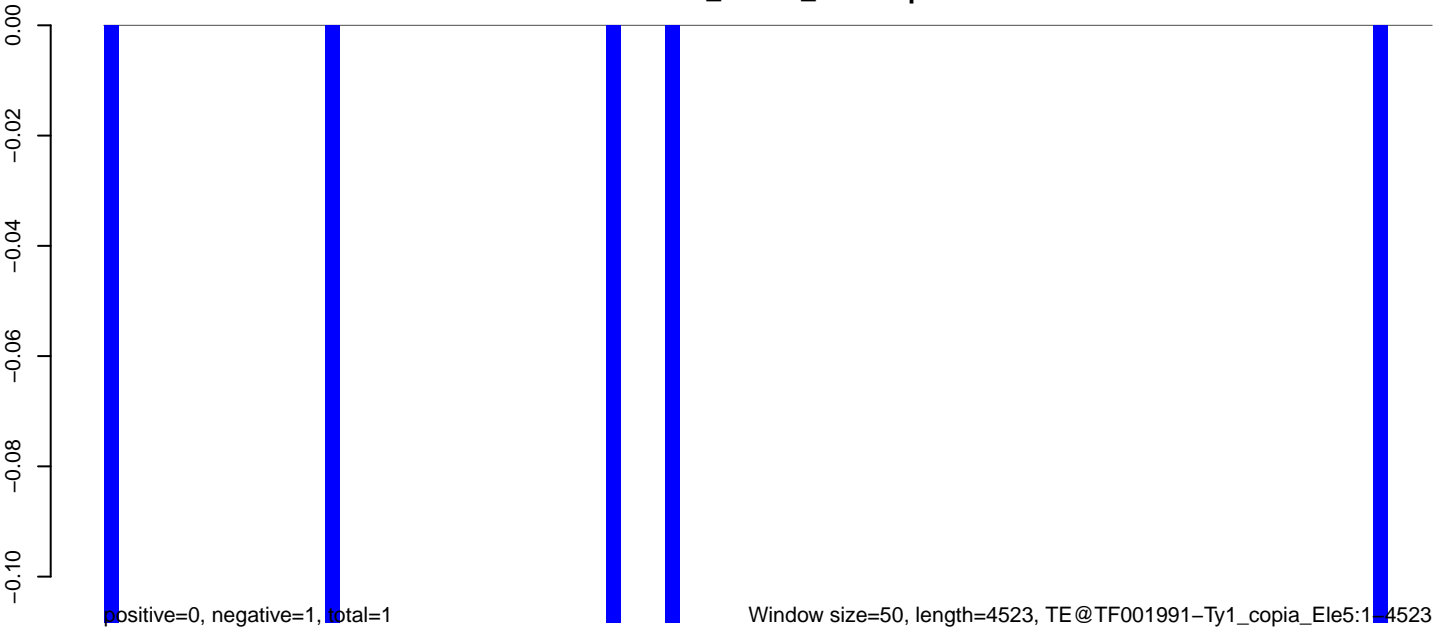
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

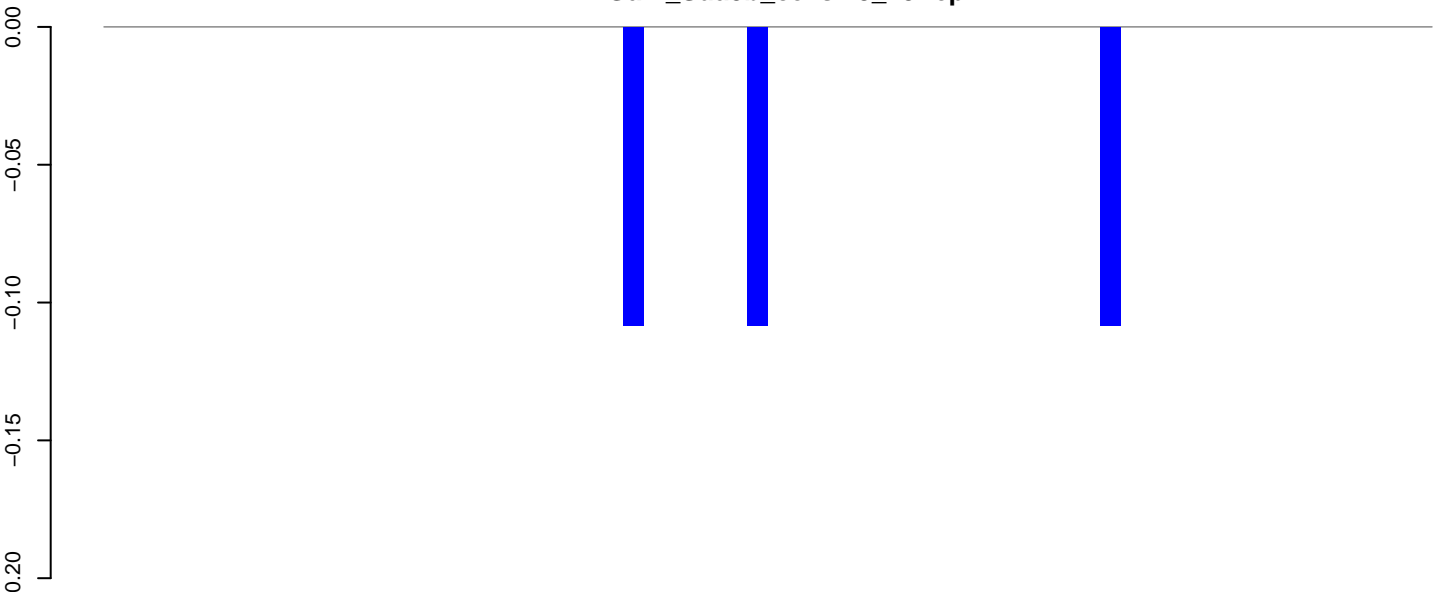


AnGam_Sua5b_cells.rep



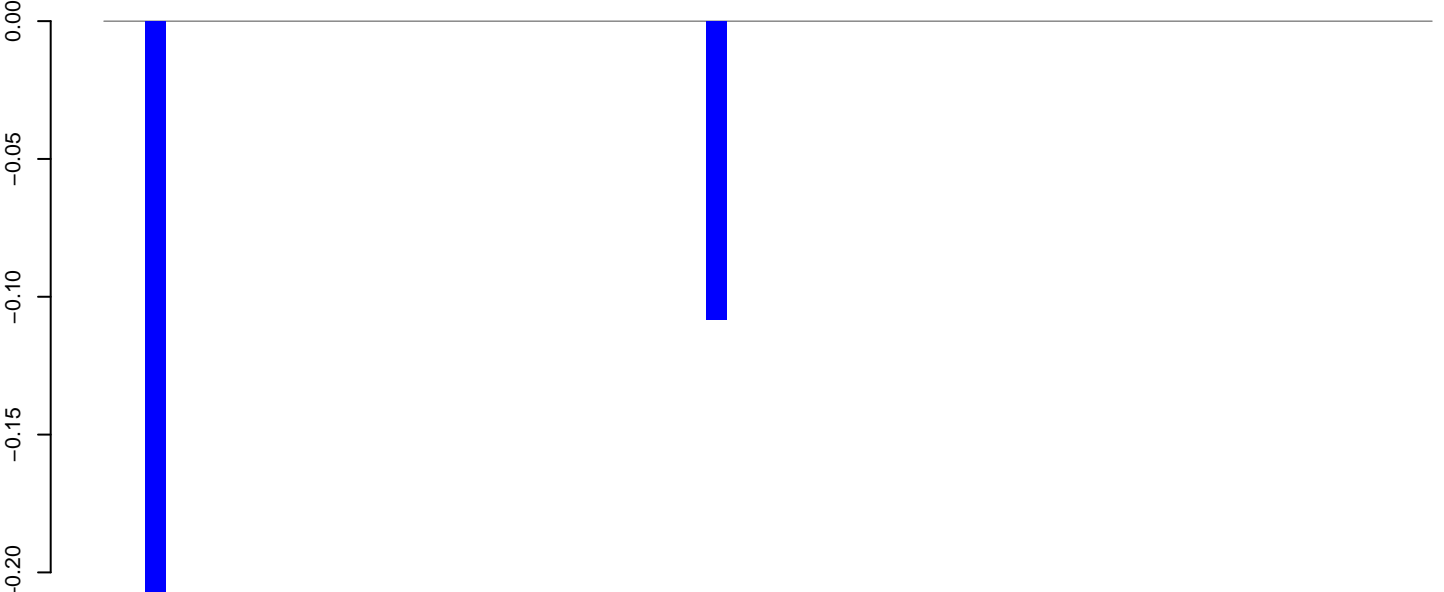
0 1000 2000 3000 4000

AnGam_Sua5b_cells.18_23.rep



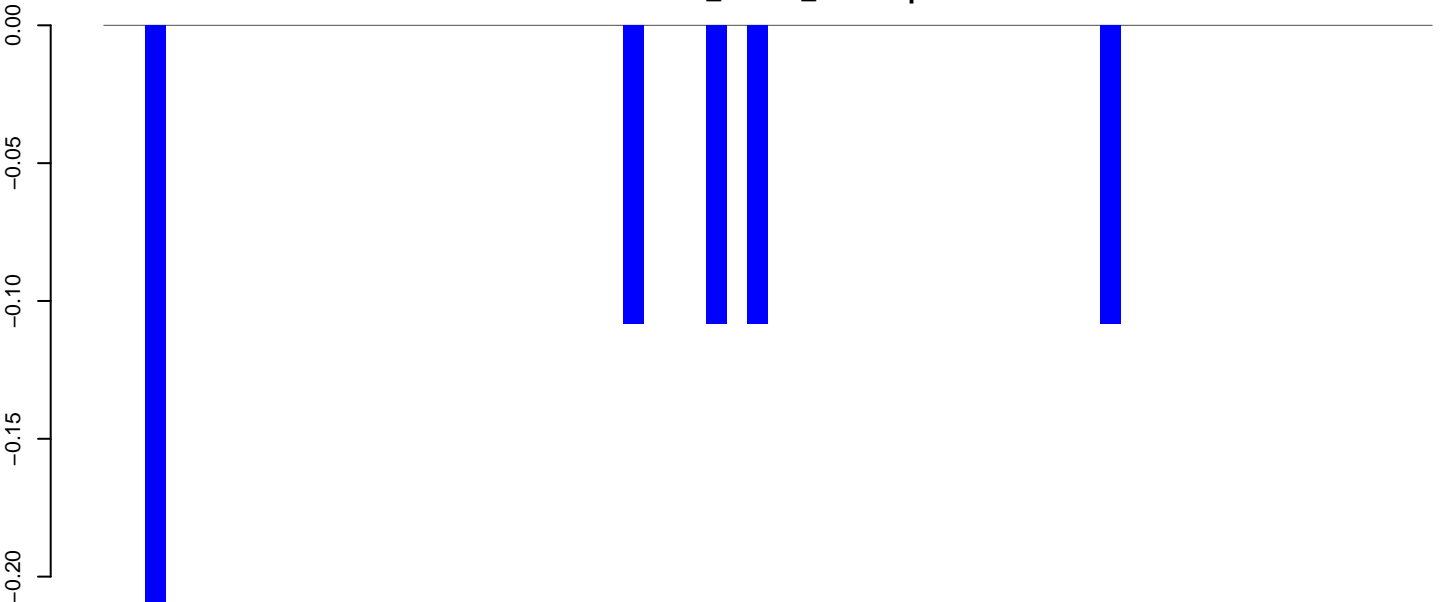
positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep



positive=0, negative=0, total=0

AnGam_Sua5b_cells.rep

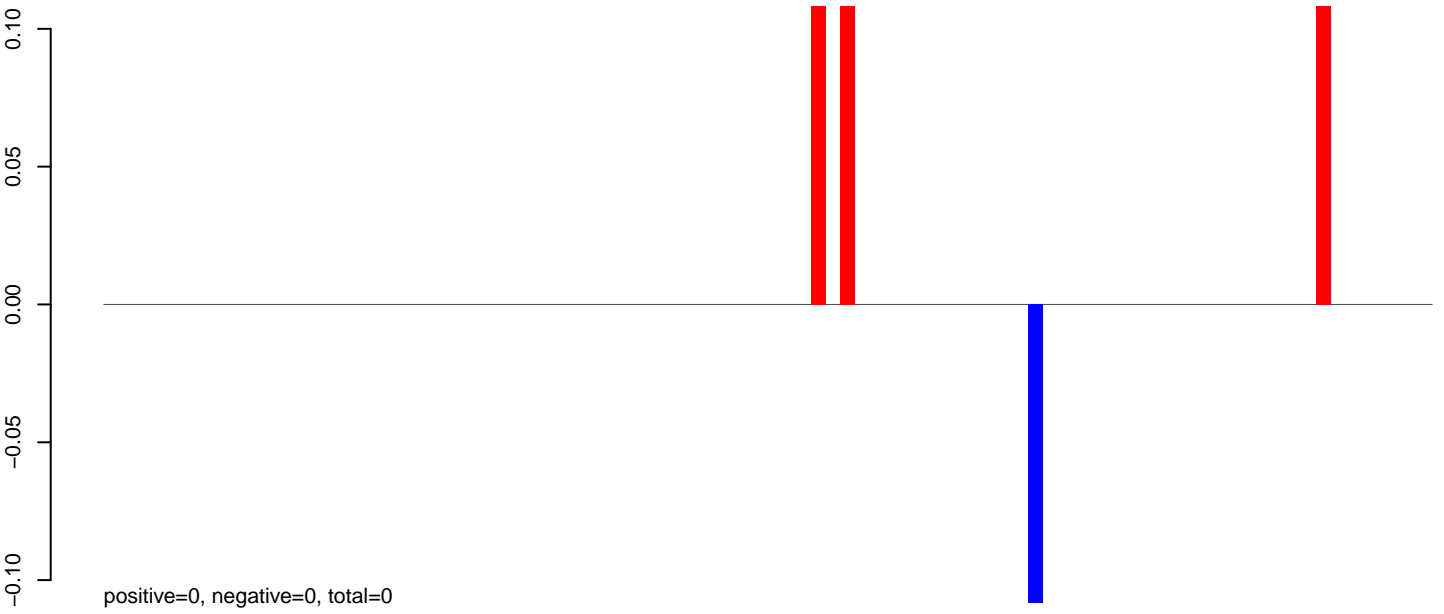


positive=0, negative=1, total=1

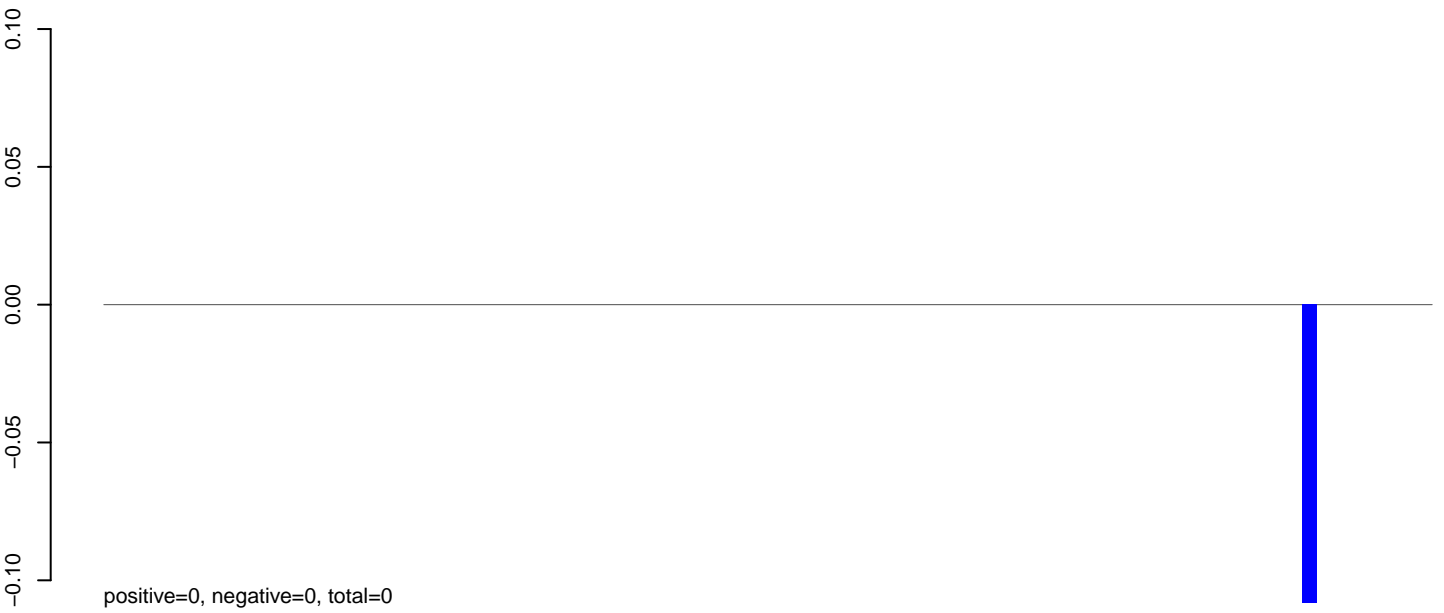
Window size=50, length=3231, TE@TF001431-L1_Ele4:1-3231

0 500 1000 1500 2000 2500 3000

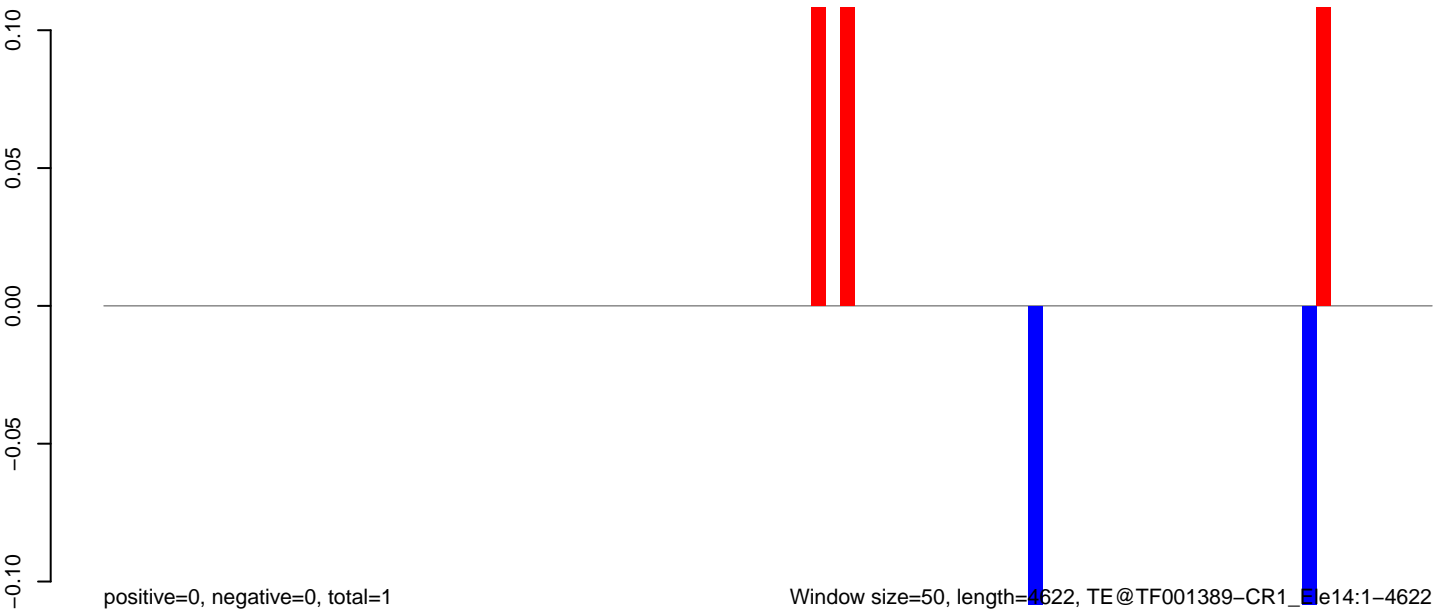
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

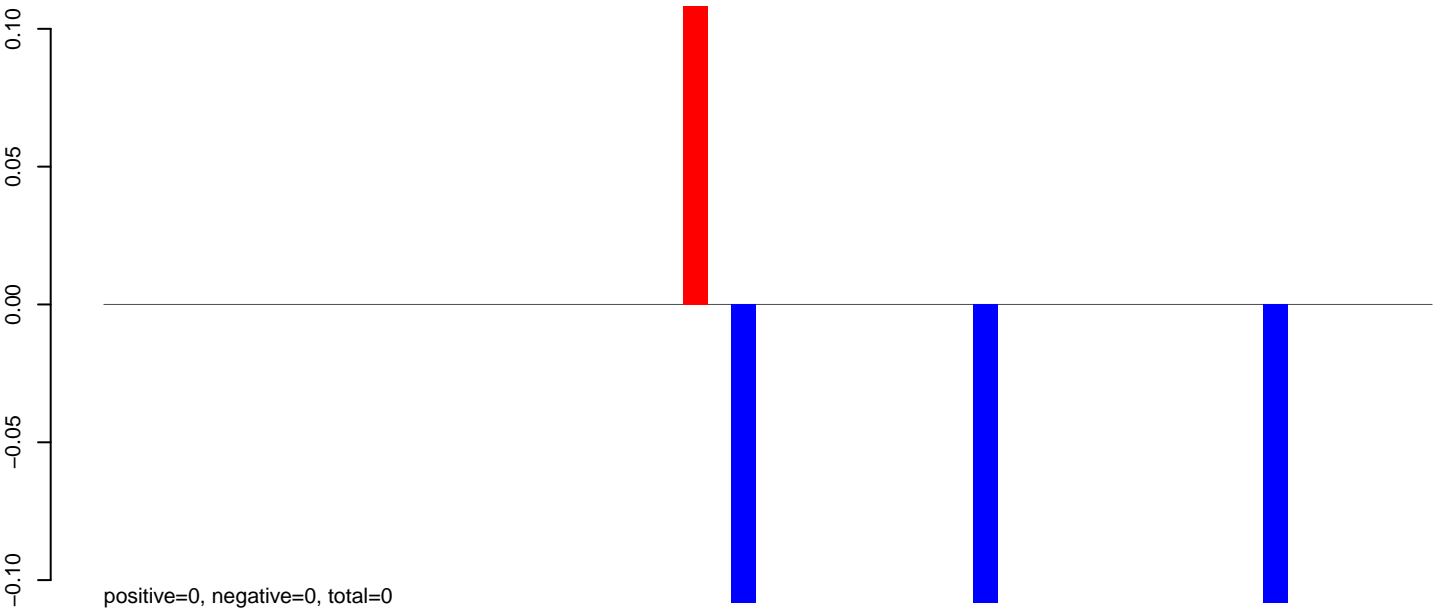


AnGam_Sua5b_cells.rep

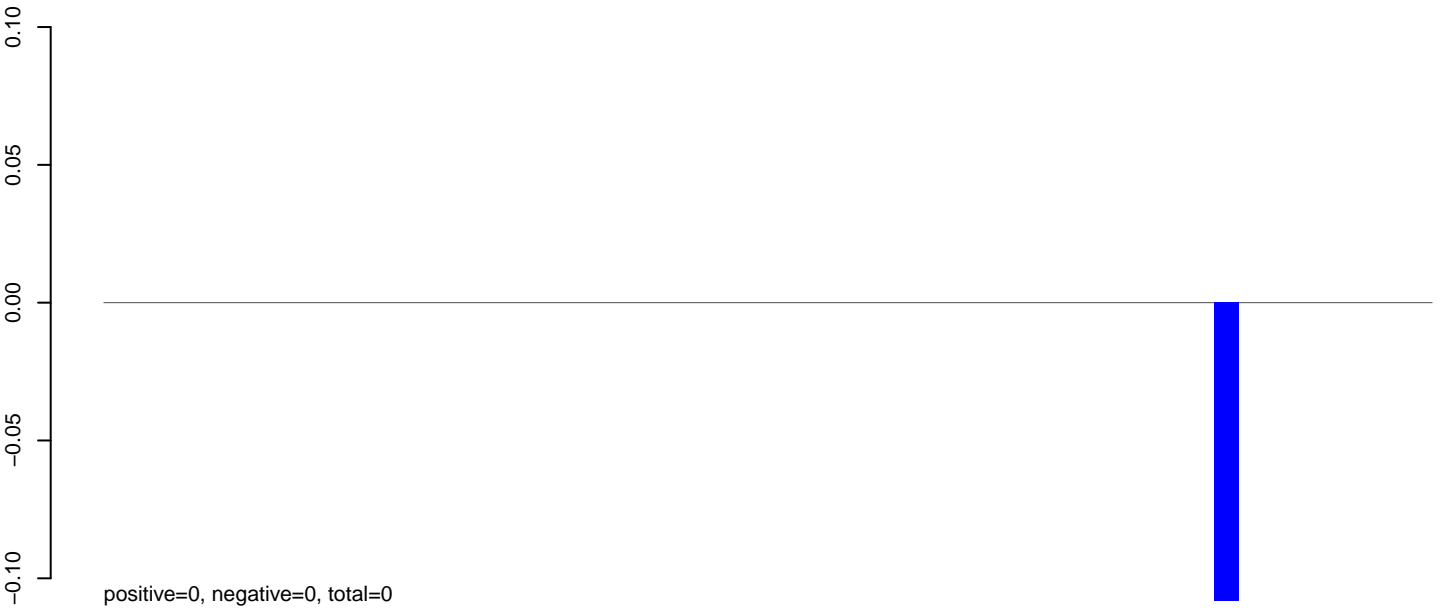


0 1000 2000 3000 4000

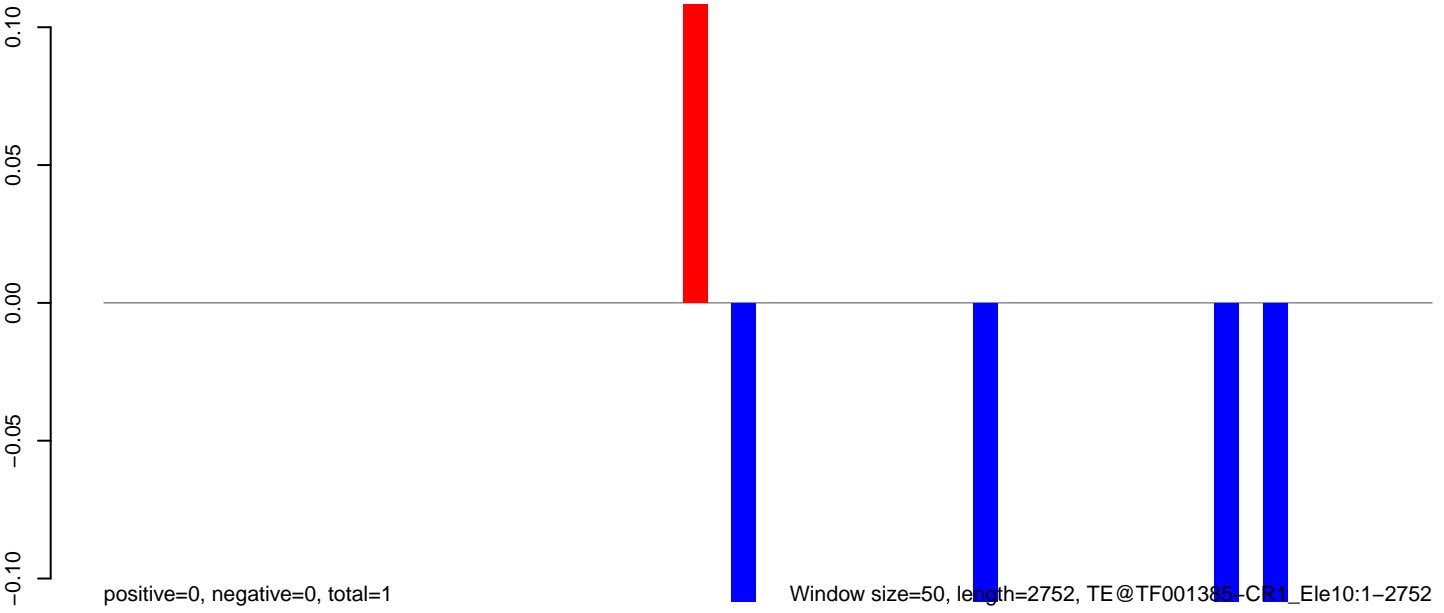
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

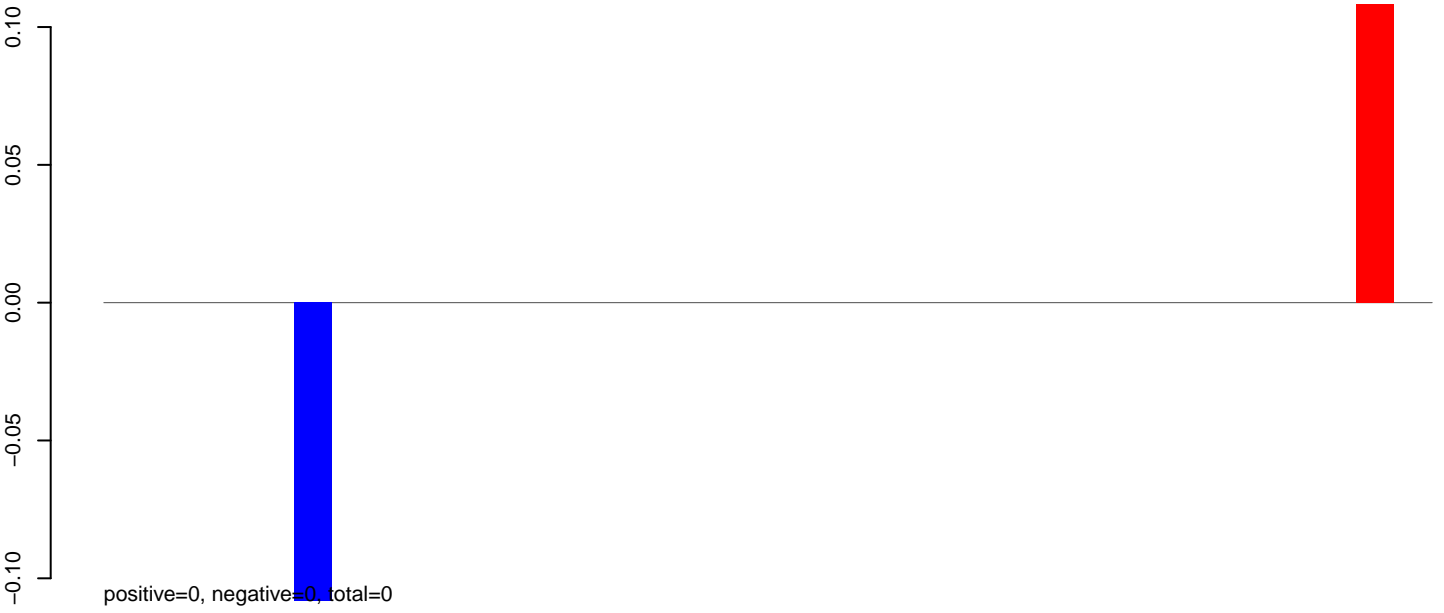


0 500 1000 1500 2000 2500

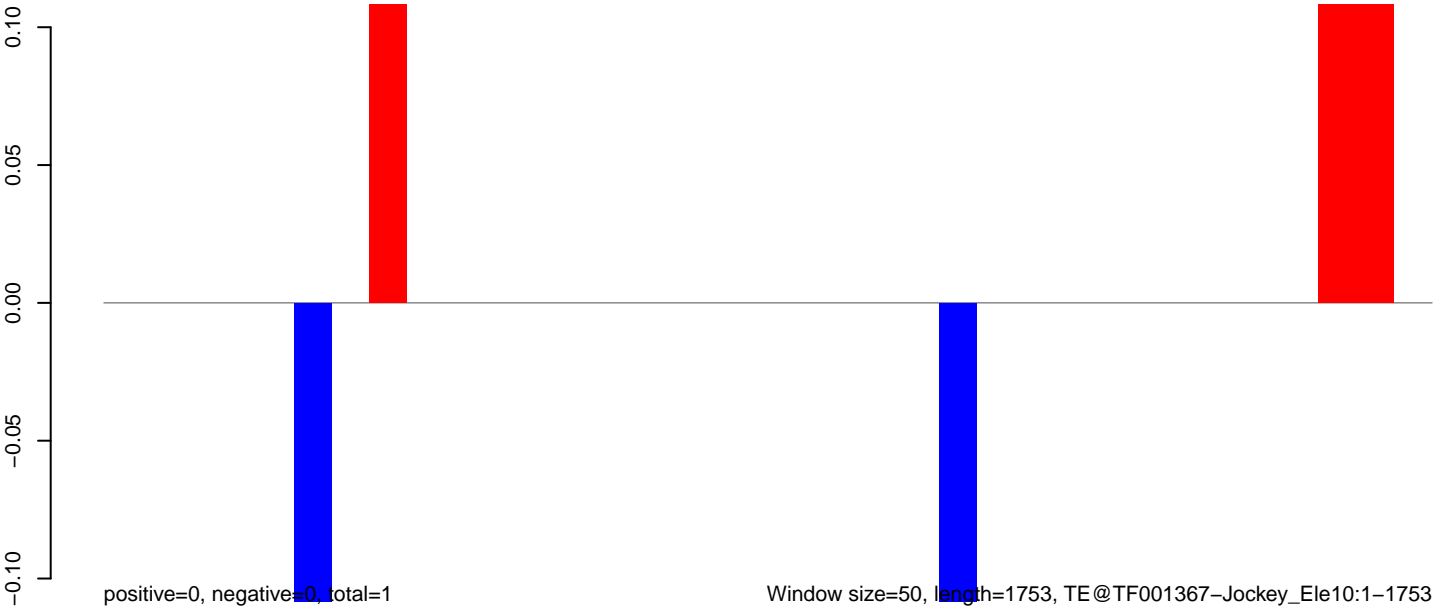
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

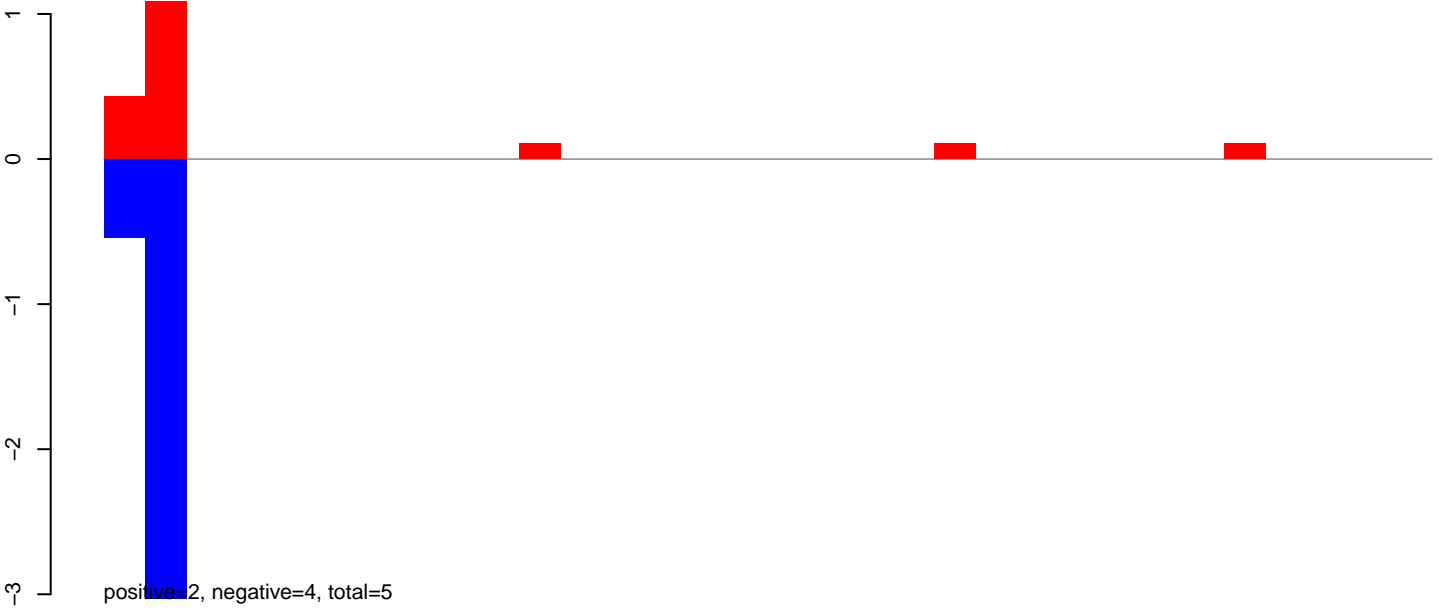


AnGam_Sua5b_cells.rep

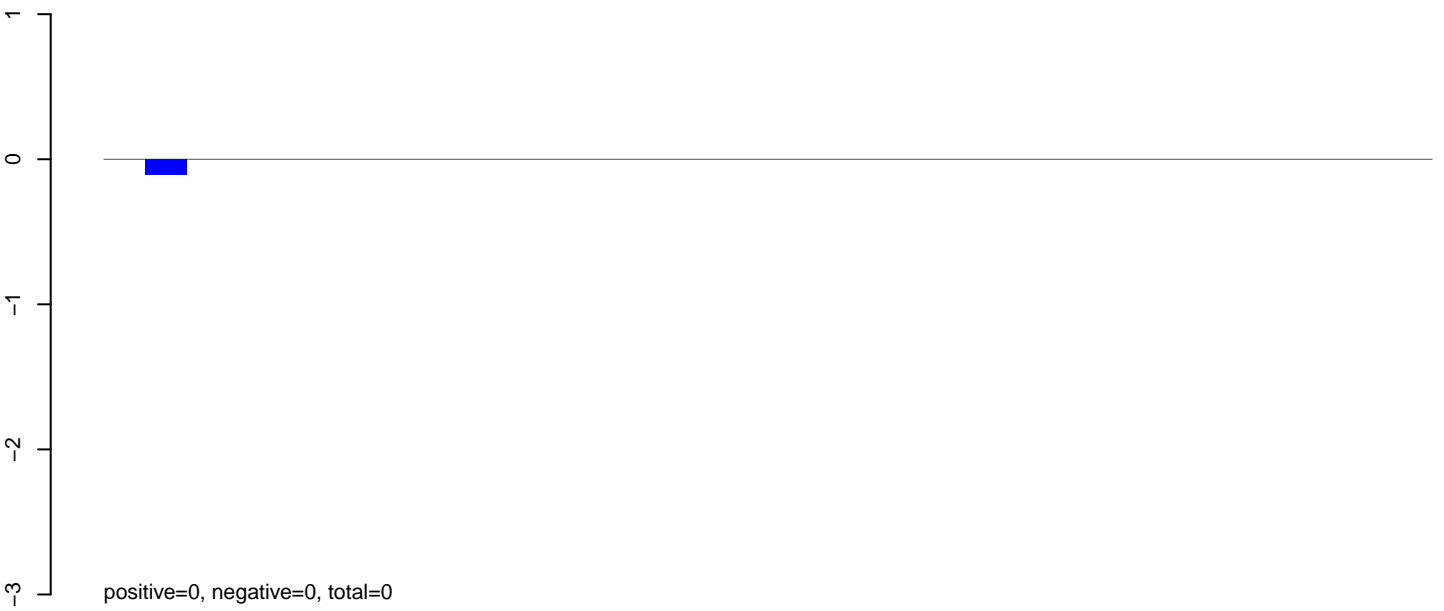


0 500 1000 1500

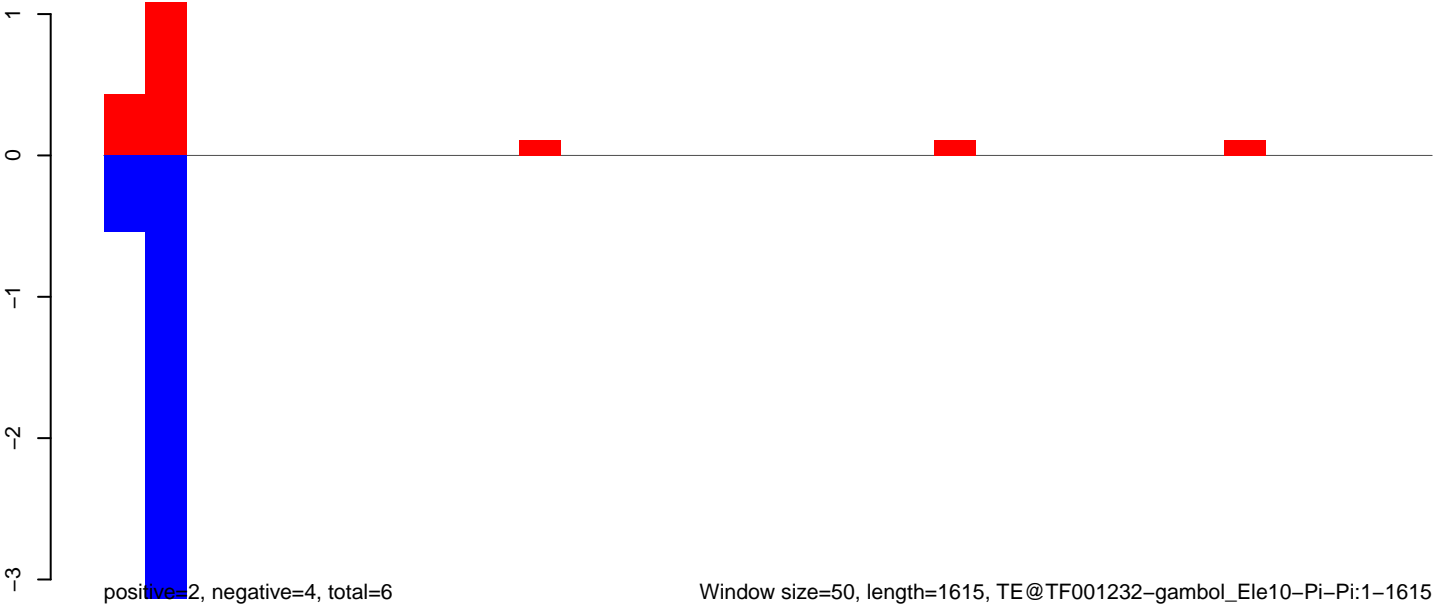
AnGam_Sua5b_cells.18_23.rep



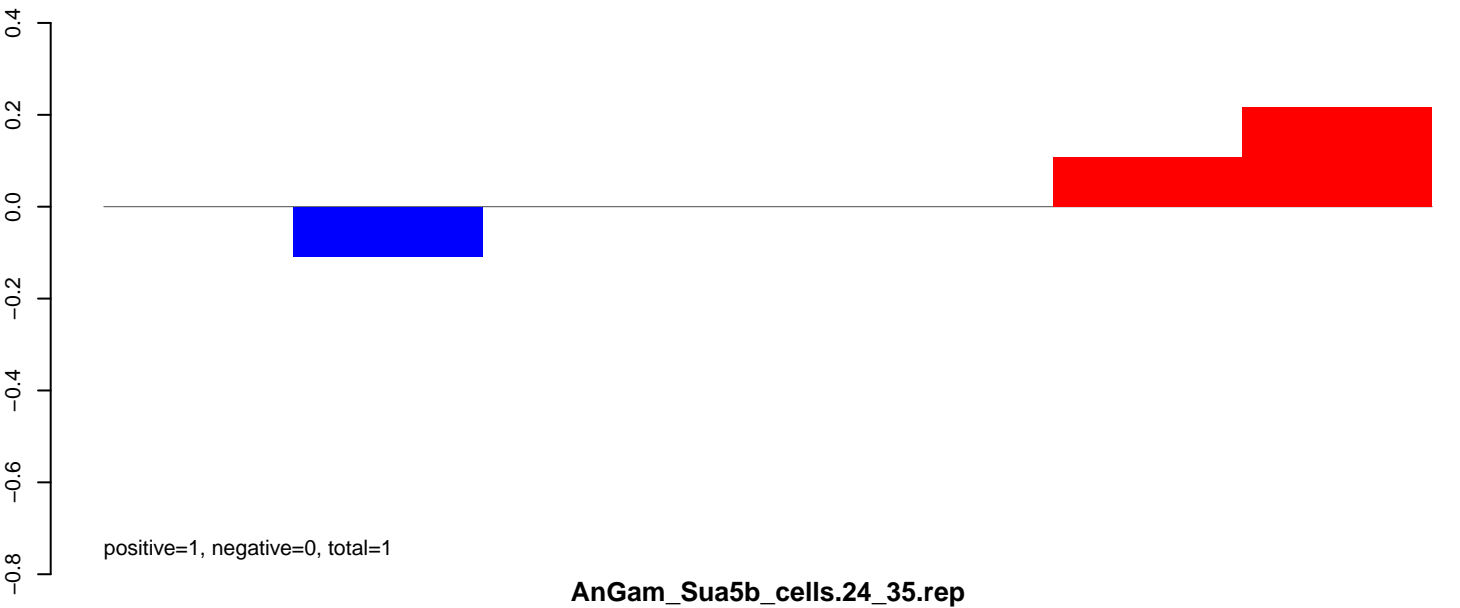
AnGam_Sua5b_cells.24_35.rep



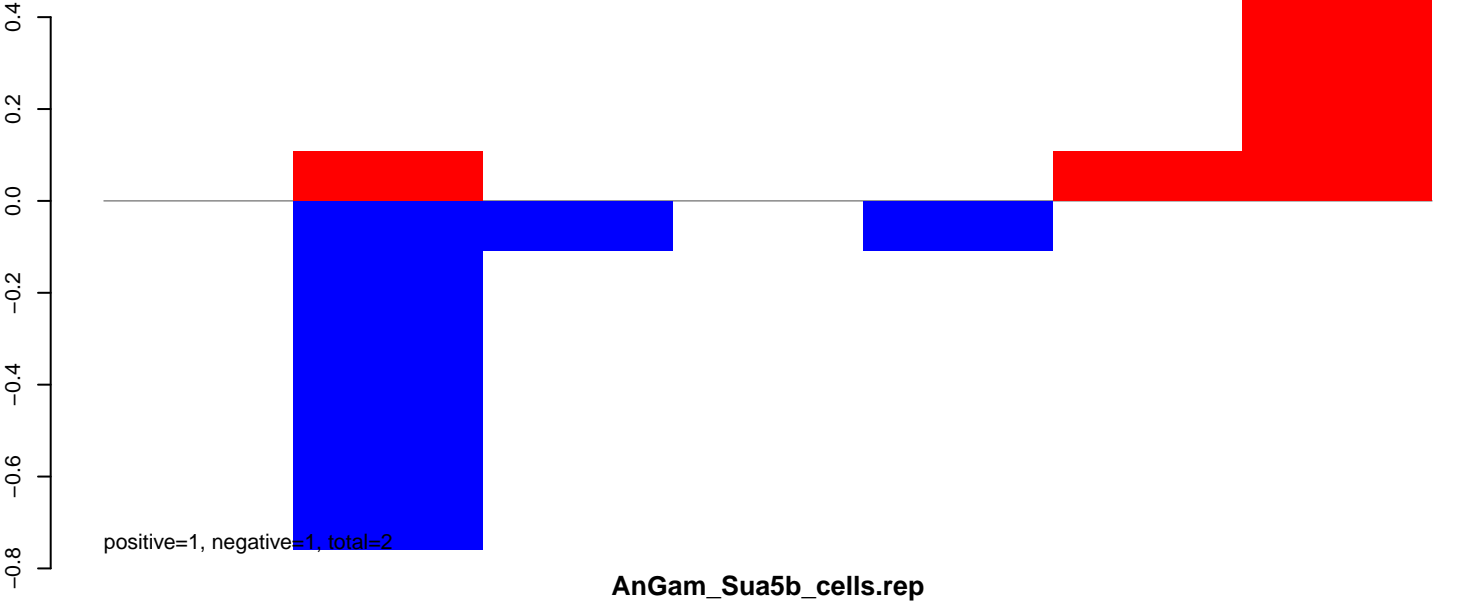
AnGam_Sua5b_cells.rep



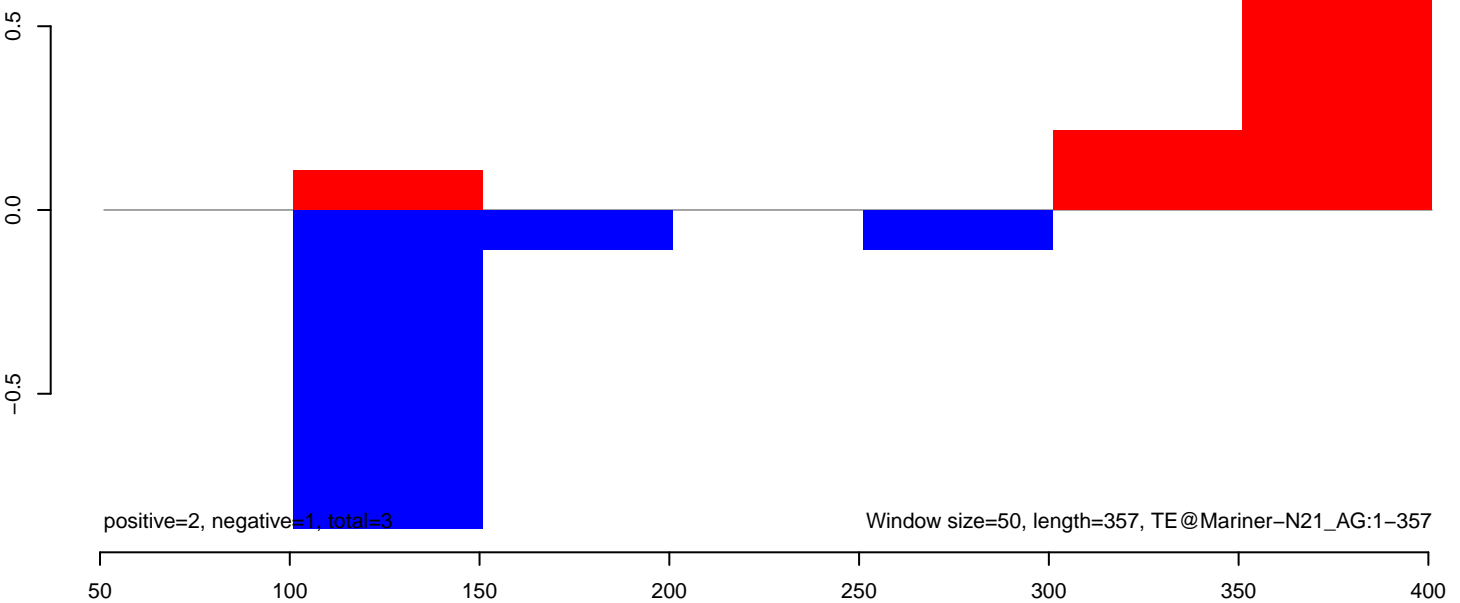
AnGam_Sua5b_cells.18_23.rep



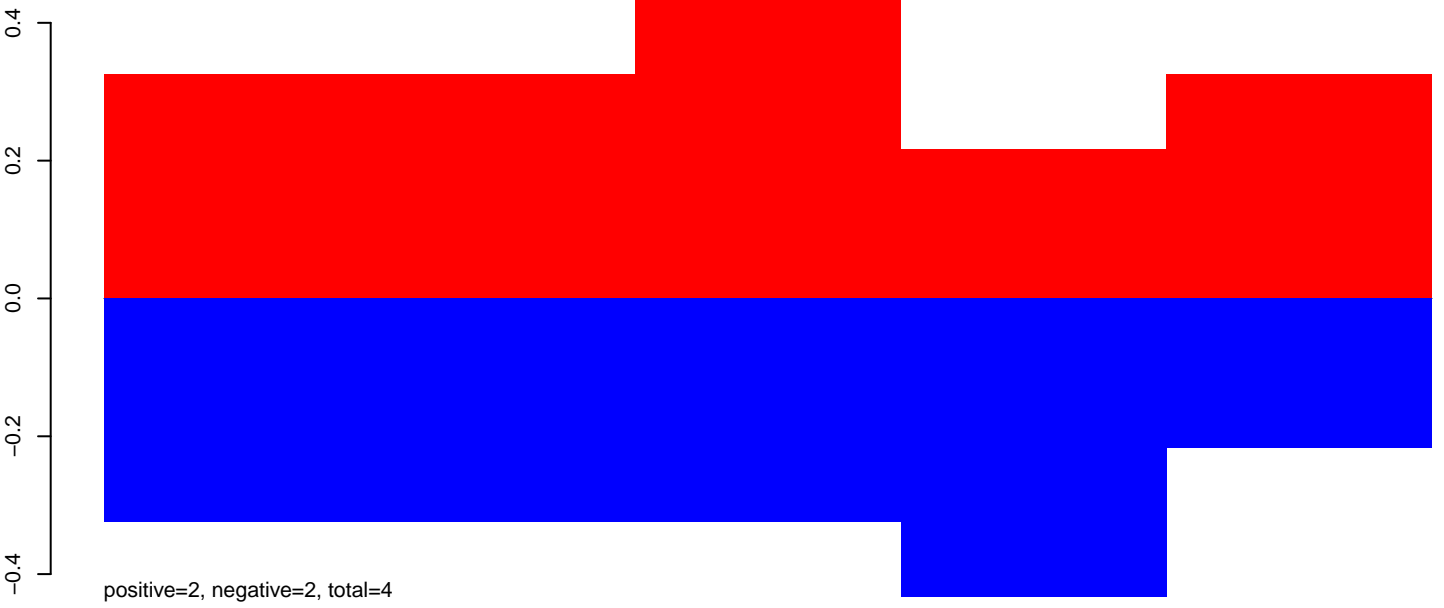
AnGam_Sua5b_cells.24_35.rep



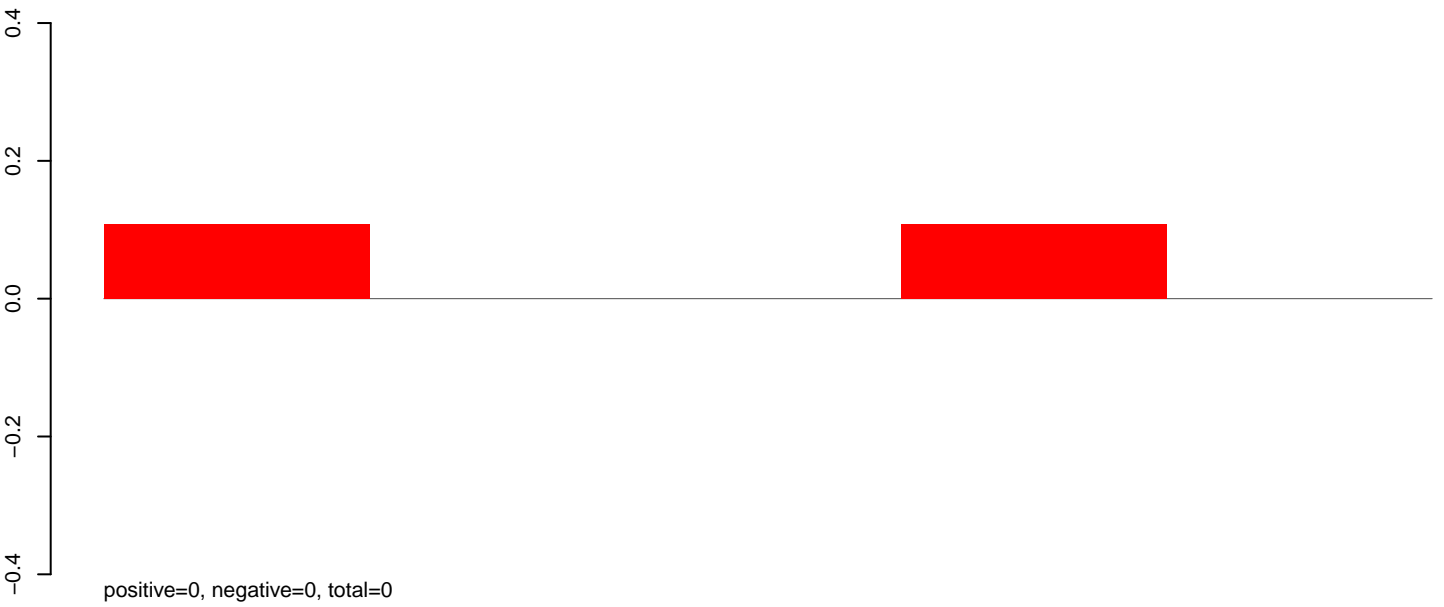
AnGam_Sua5b_cells.rep



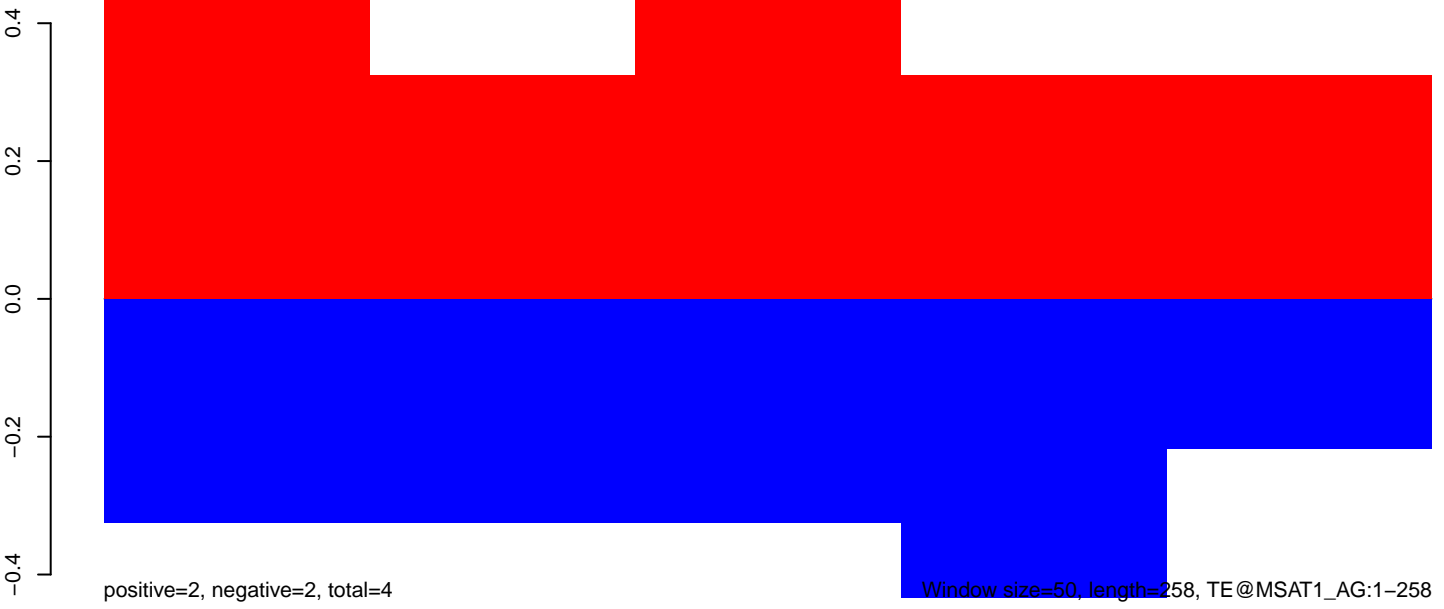
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

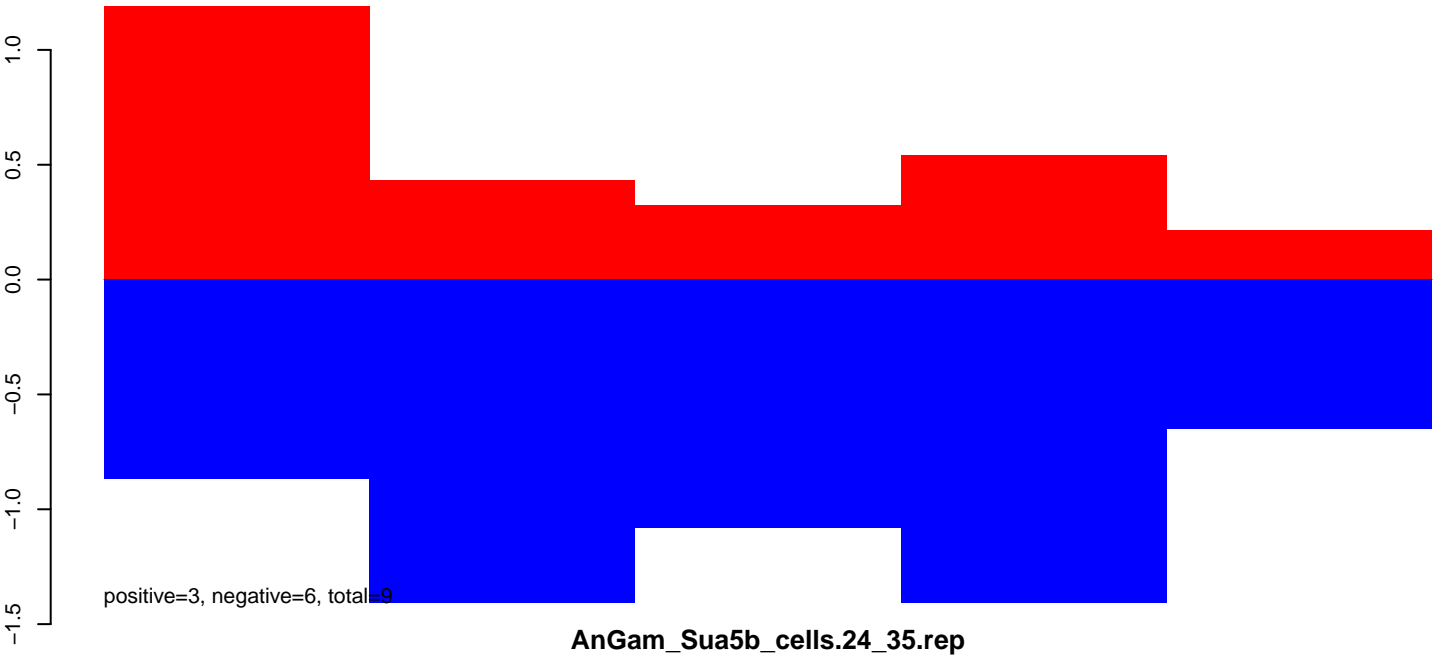


AnGam_Sua5b_cells.rep

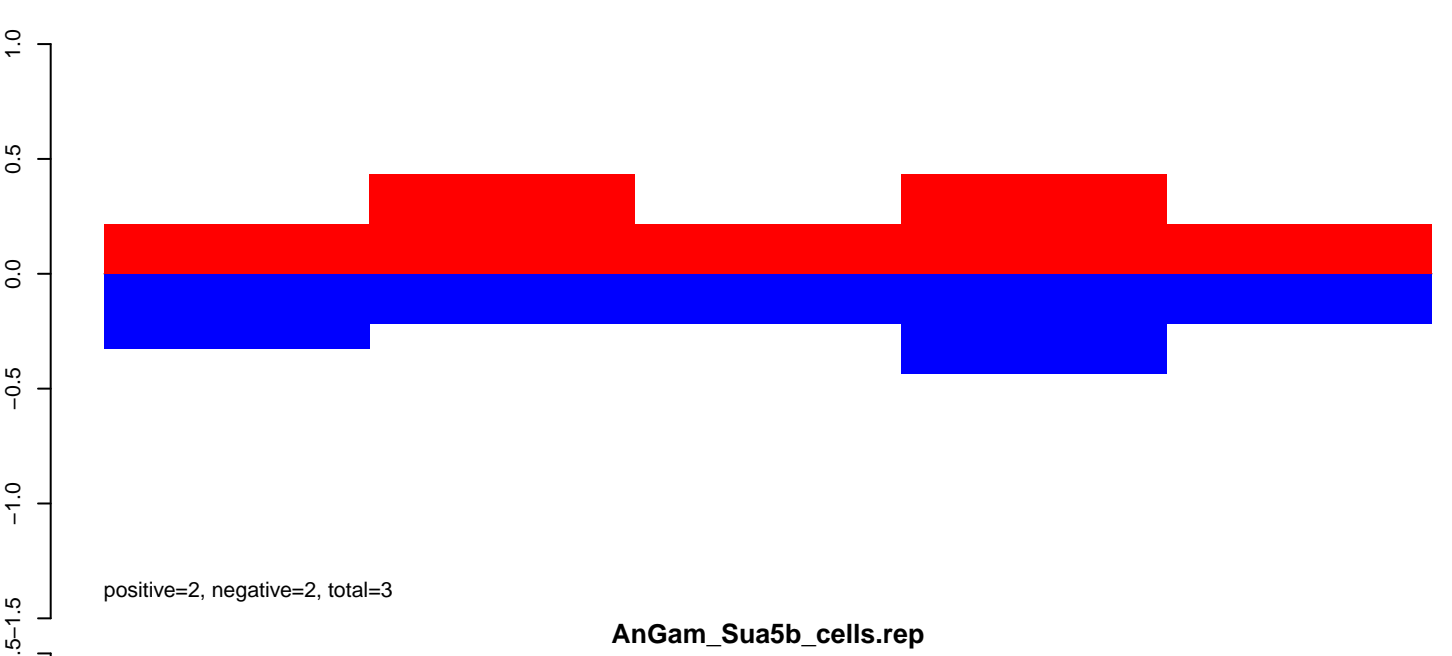


Window size=50, length=258, TE@MSAT1_AG:1-258

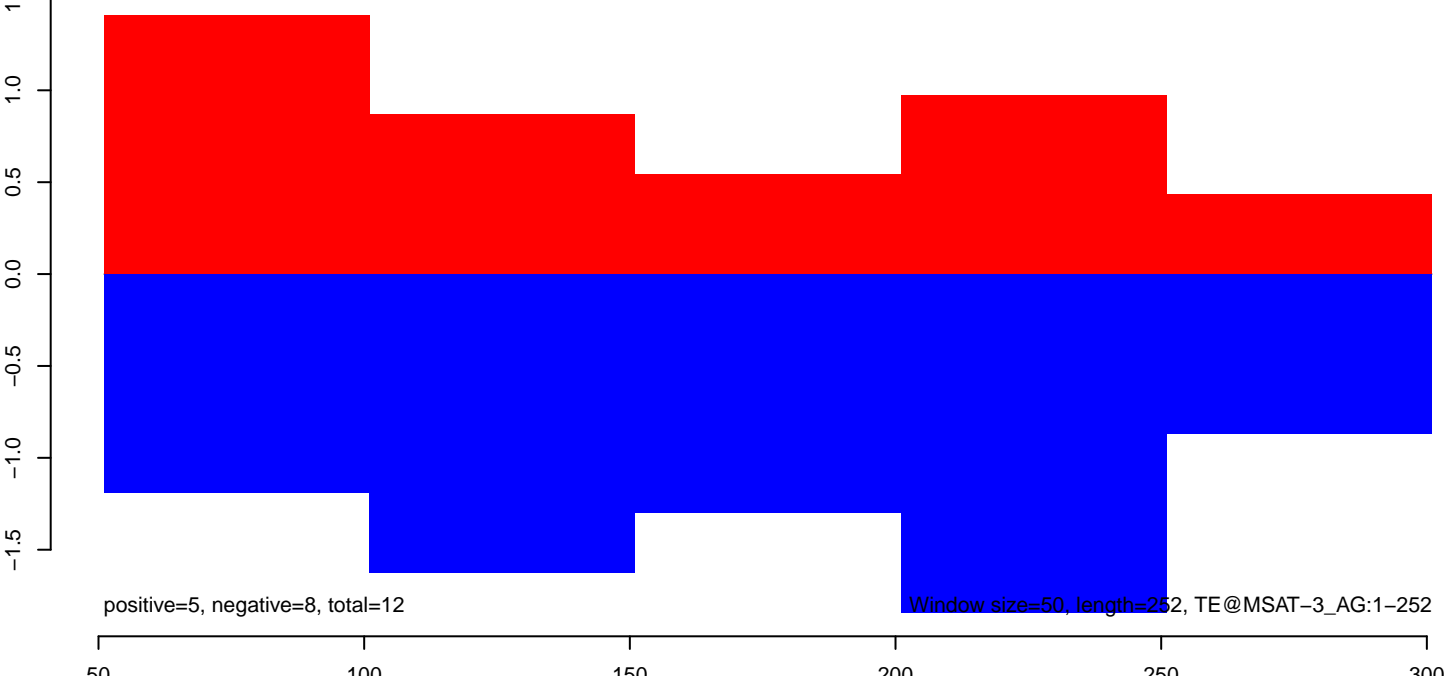
AnGam_Sua5b_cells.18_23.rep



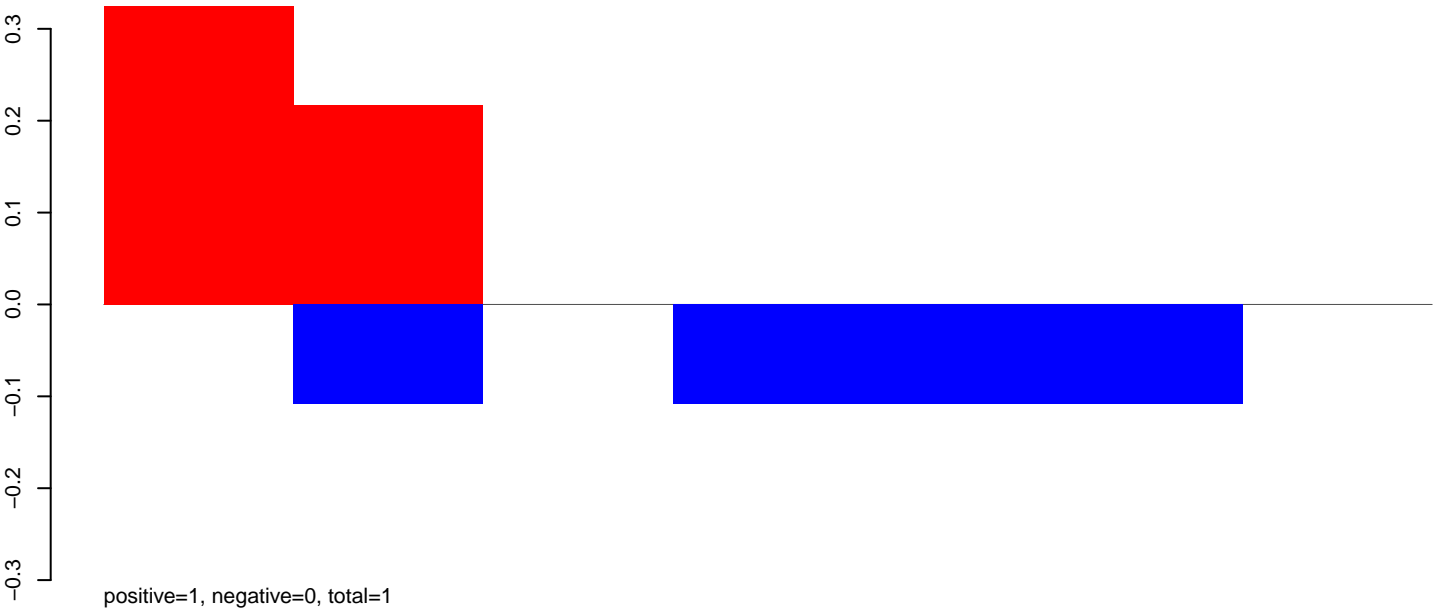
AnGam_Sua5b_cells.24_35.rep



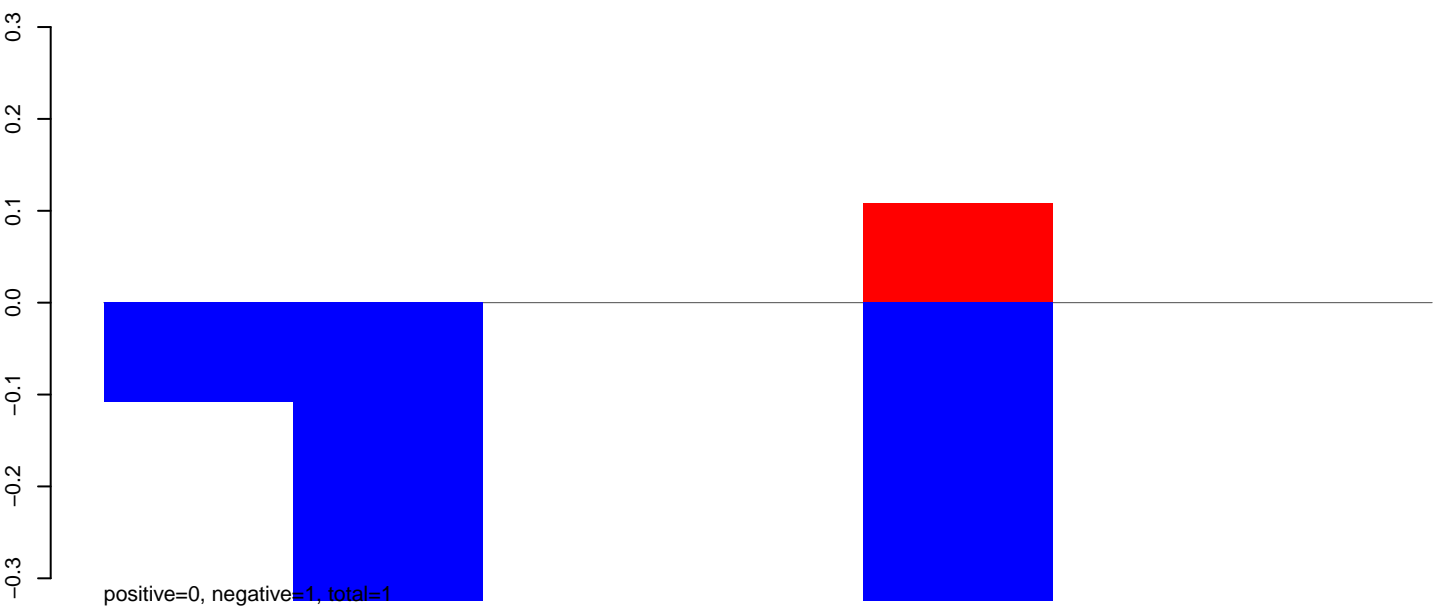
AnGam_Sua5b_cells.rep



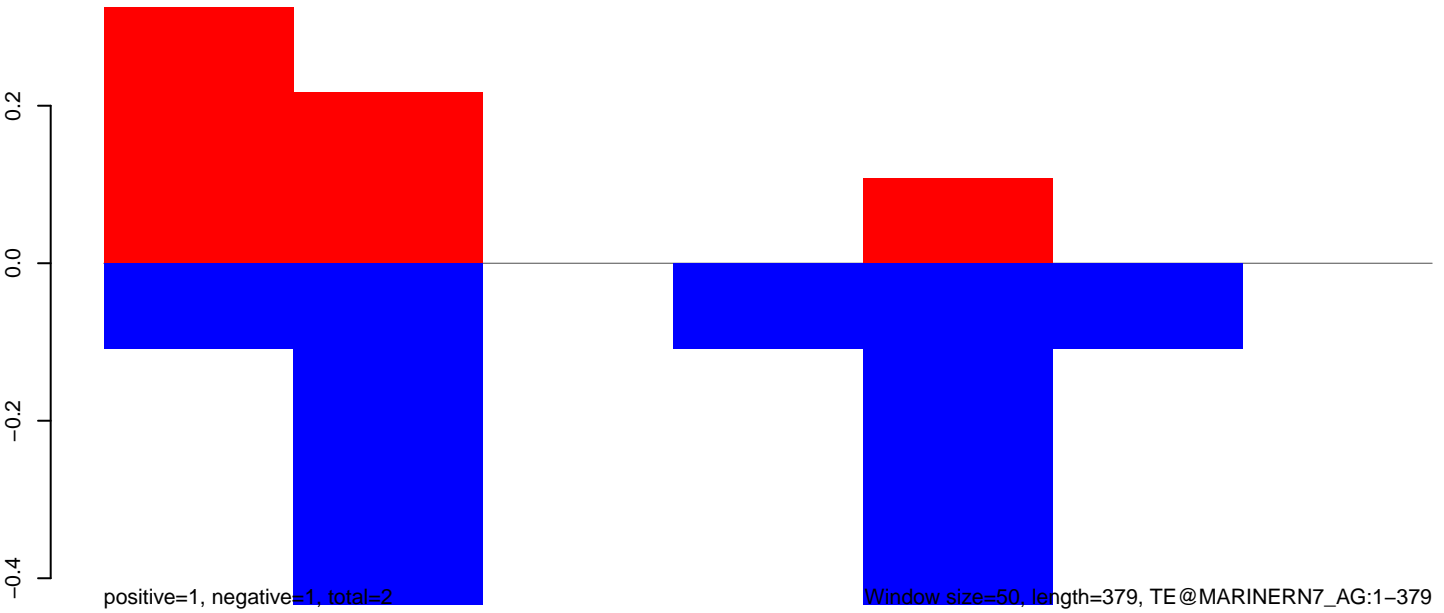
AnGam_Sua5b_cells.18_23.rep



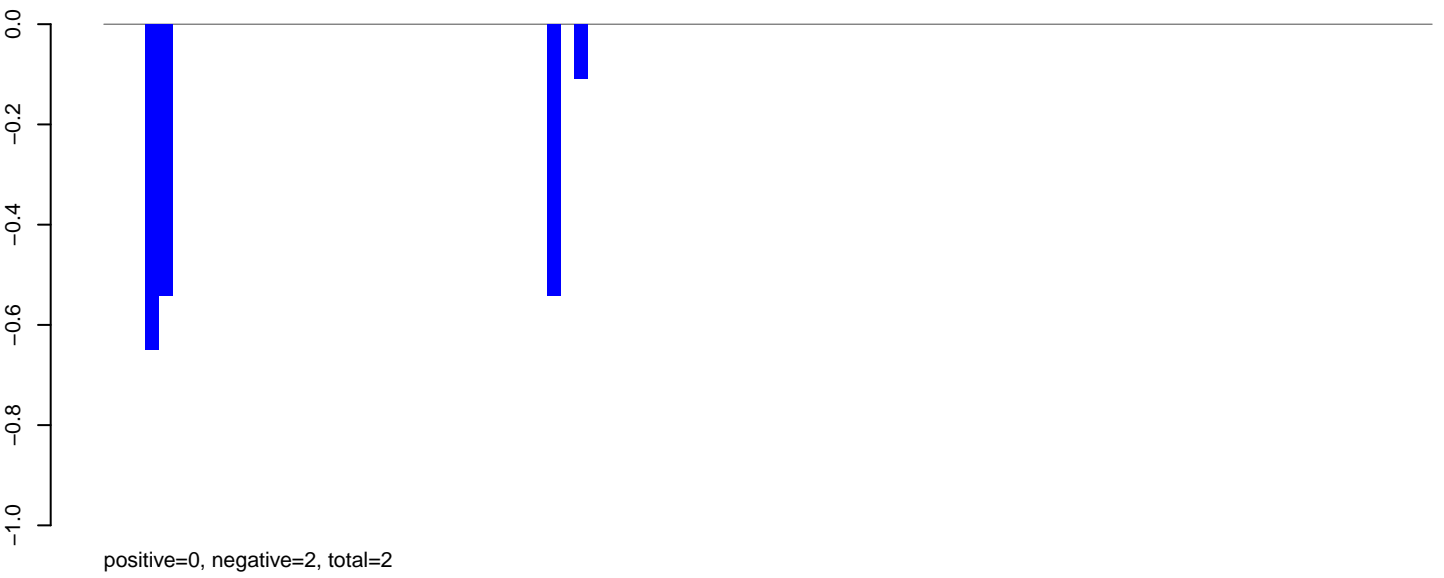
AnGam_Sua5b_cells.24_35.rep



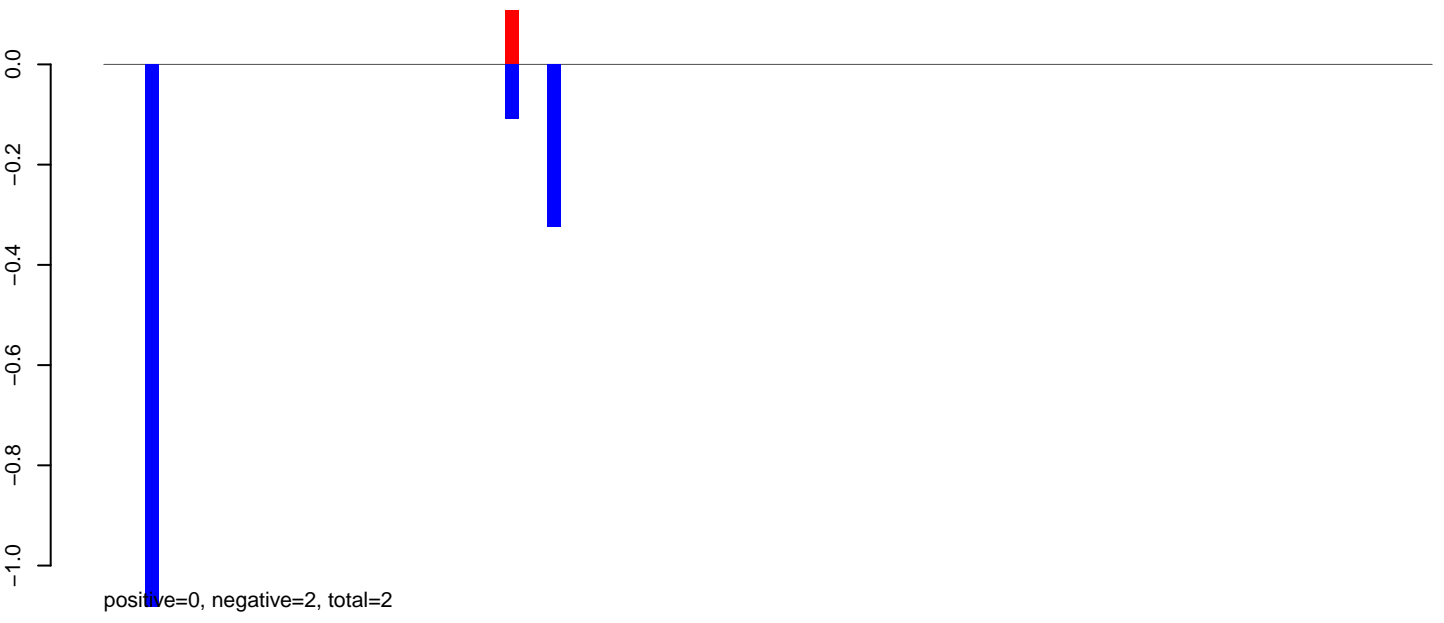
AnGam_Sua5b_cells.rep



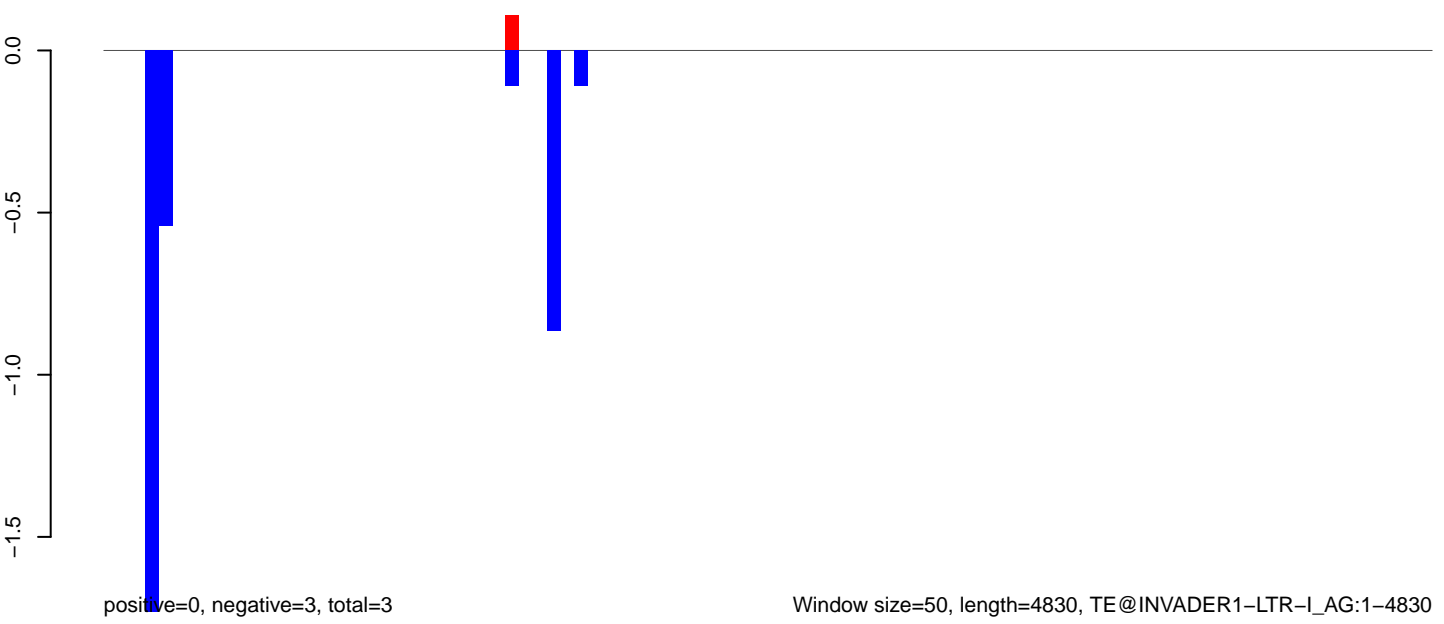
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



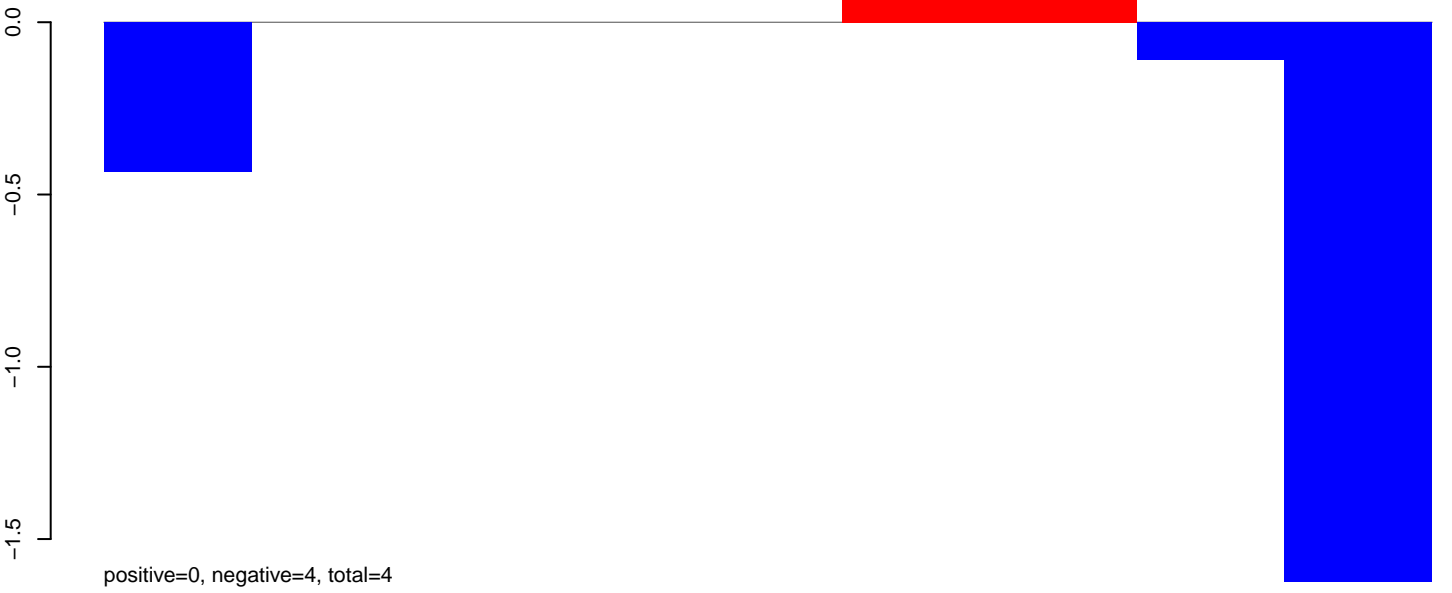
AnGam_Sua5b_cells.rep



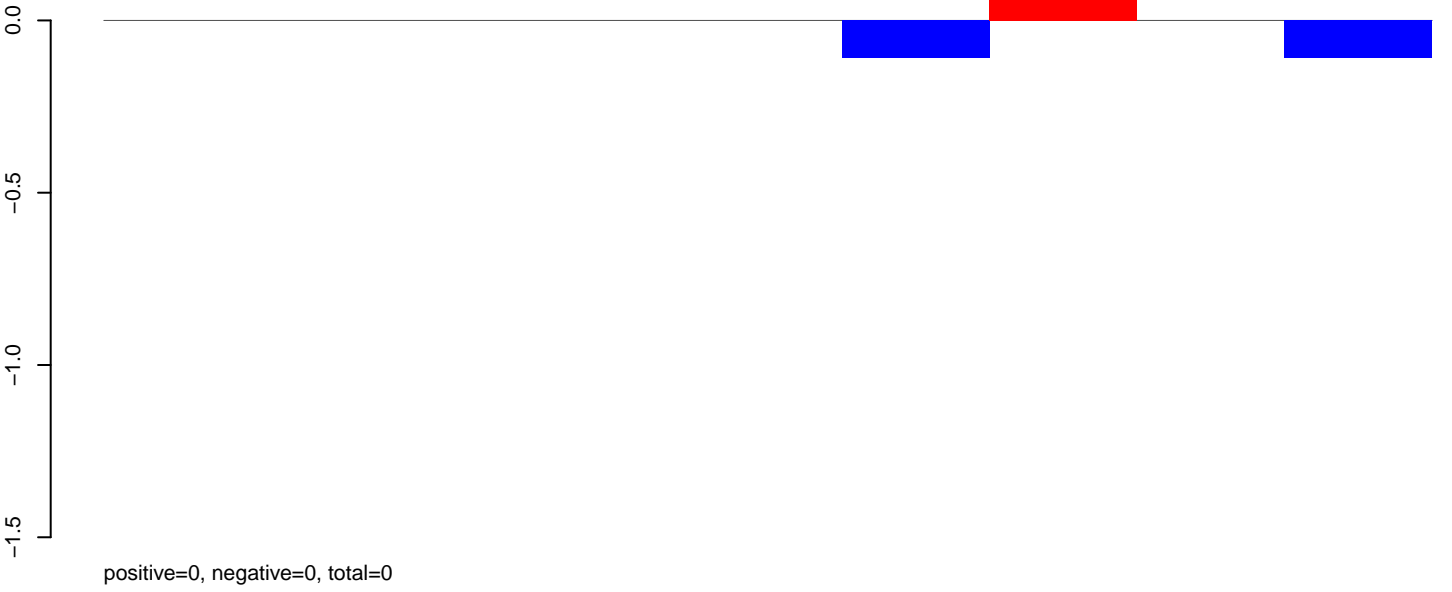
Window size=50, length=4830, TE@INVADER1-LTR-I_AG:1-4830

0 1000 2000 3000 4000 5000

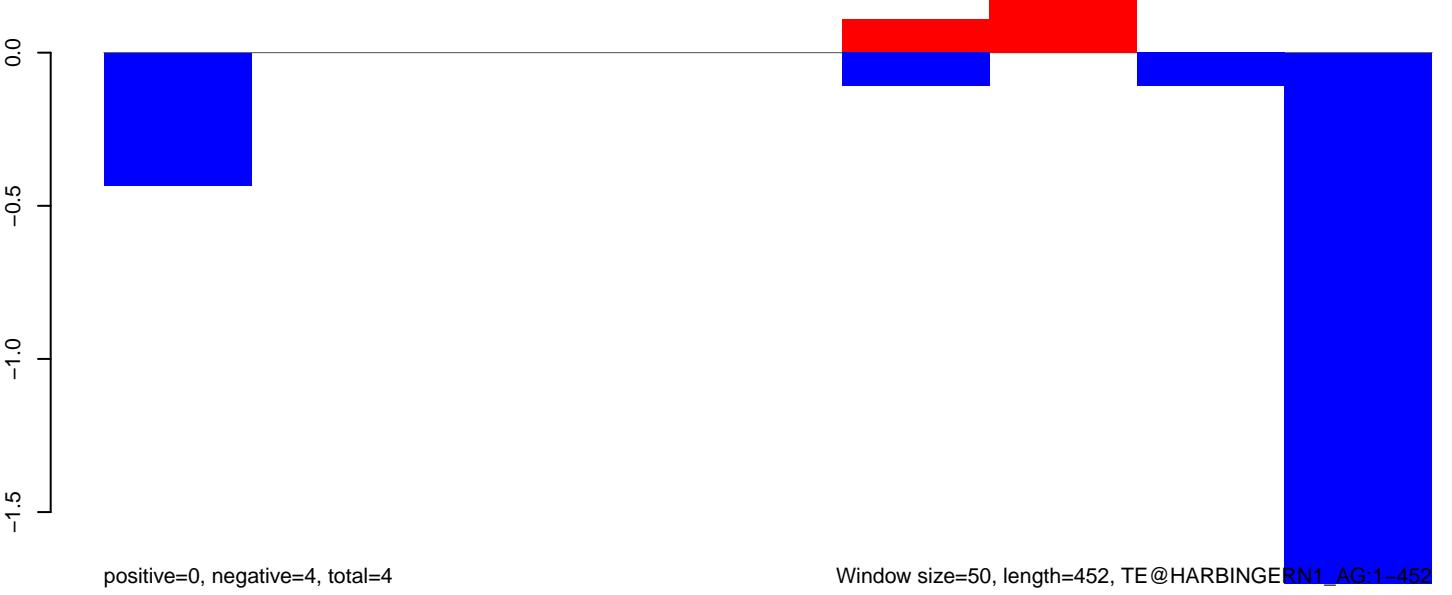
AnGam_Sua5b_cells.18_23.rep



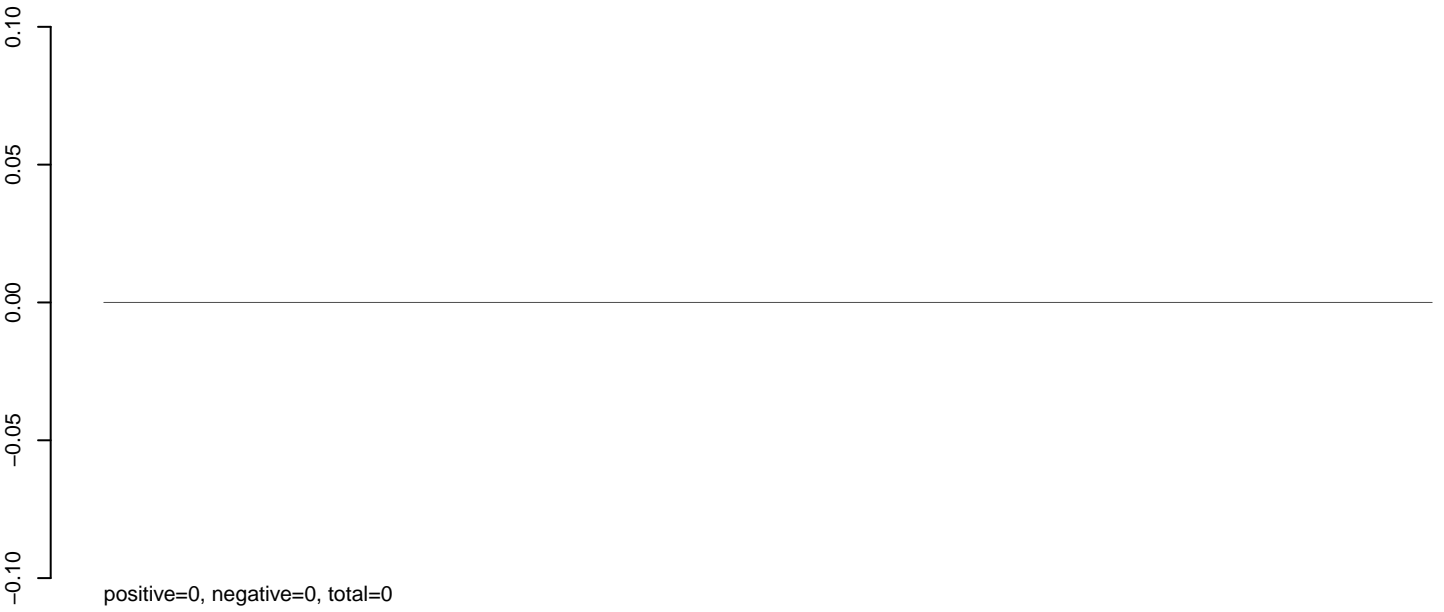
AnGam_Sua5b_cells.24_35.rep



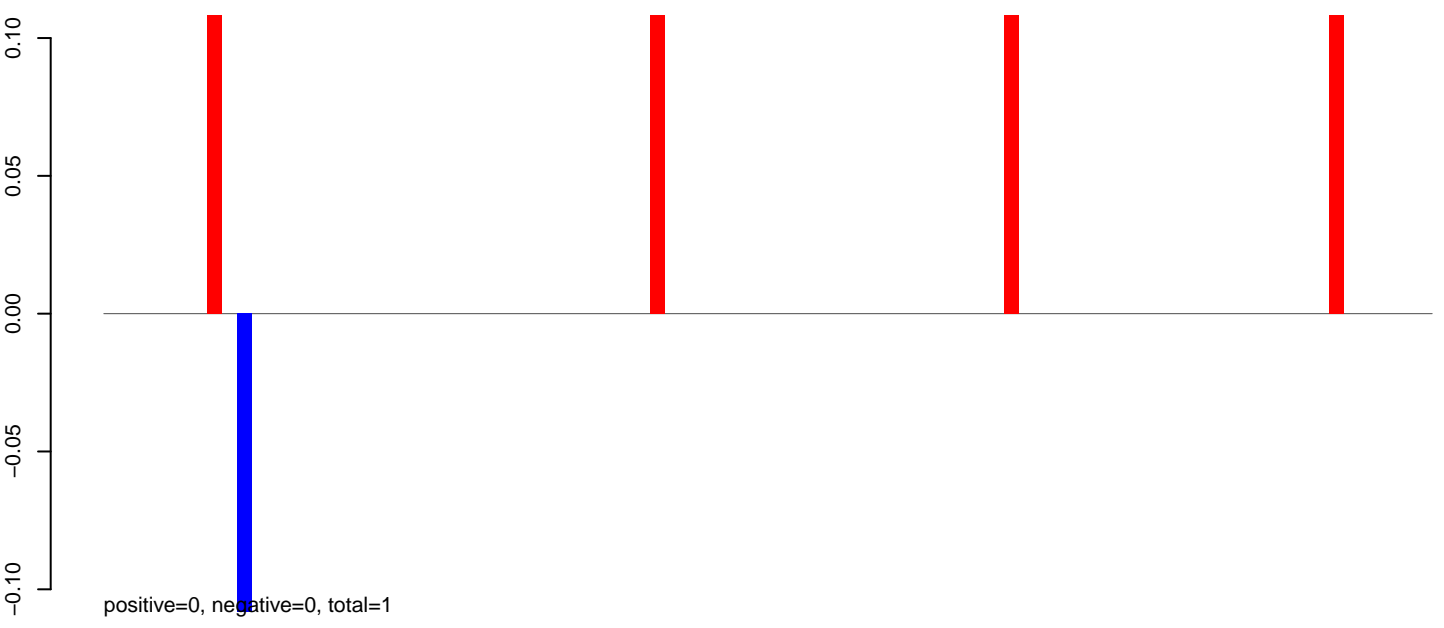
AnGam_Sua5b_cells.rep



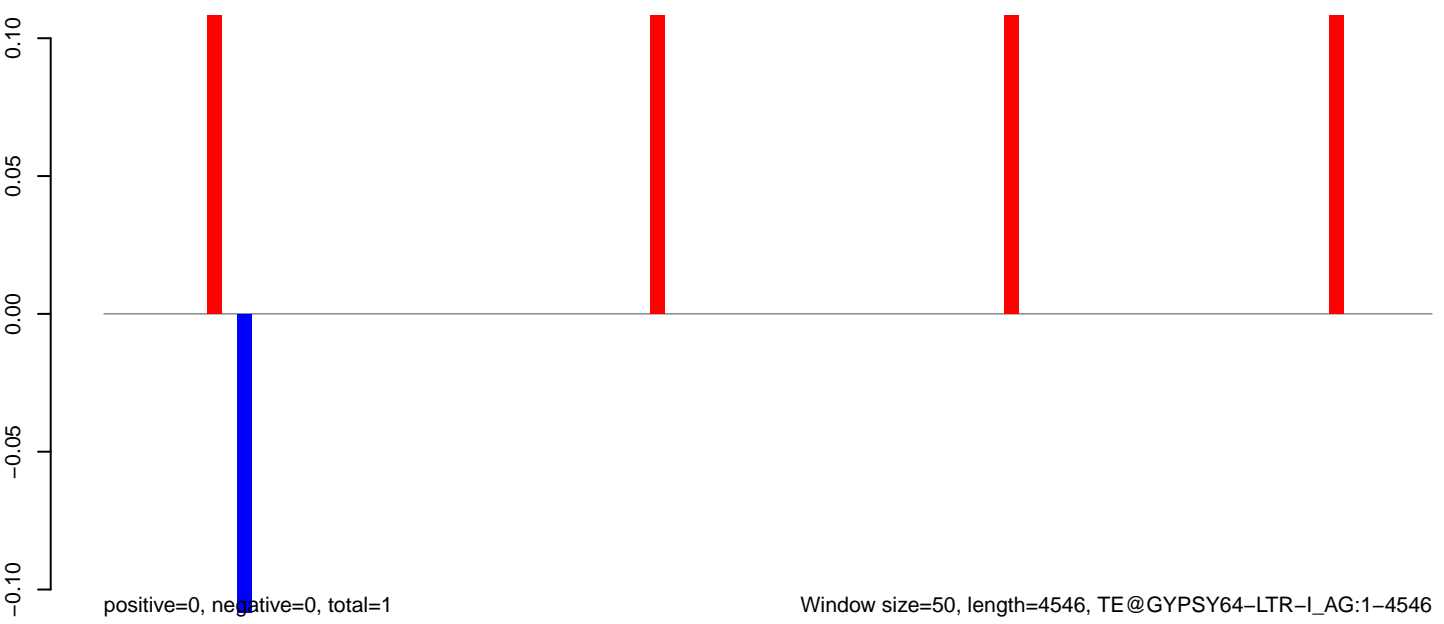
AnGam_Sua5b_cells.18_23.rep



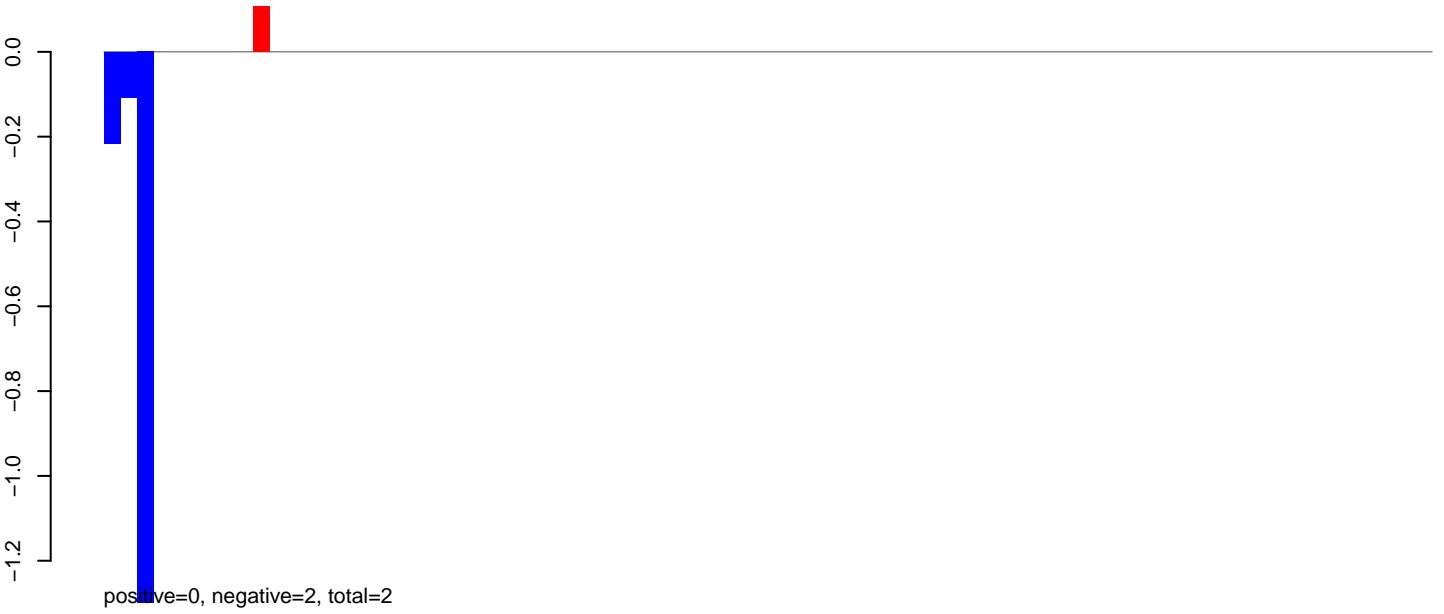
AnGam_Sua5b_cells.24_35.rep



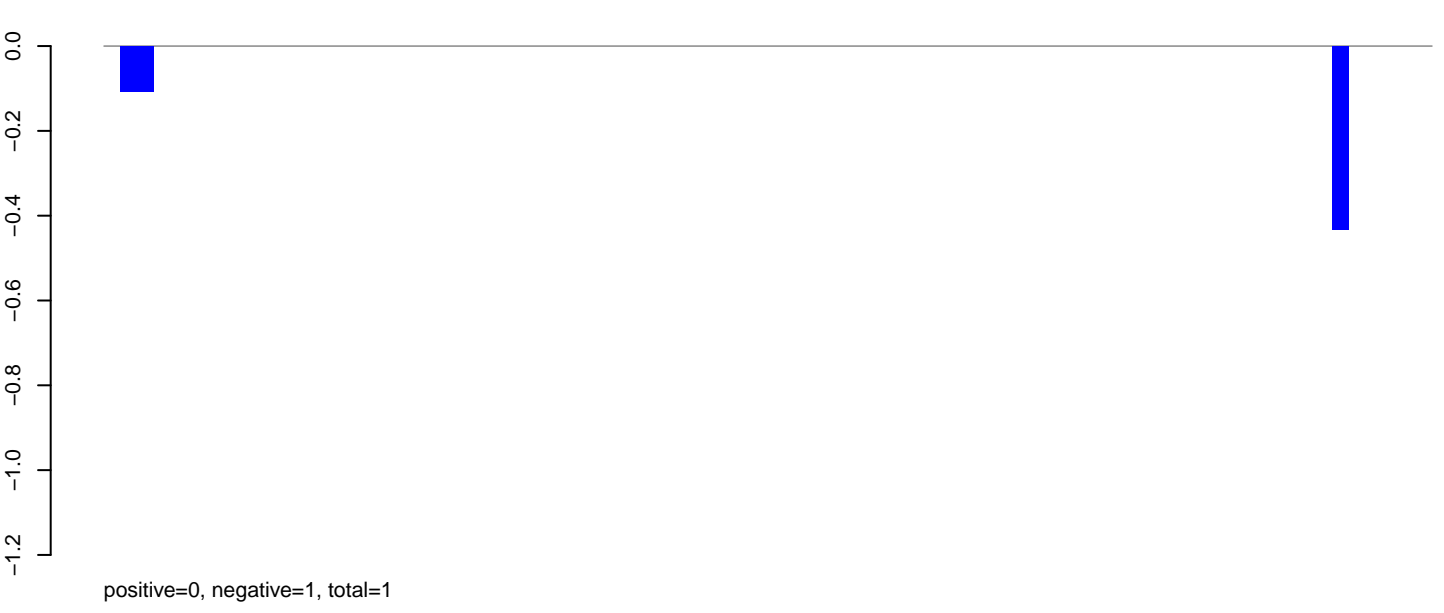
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



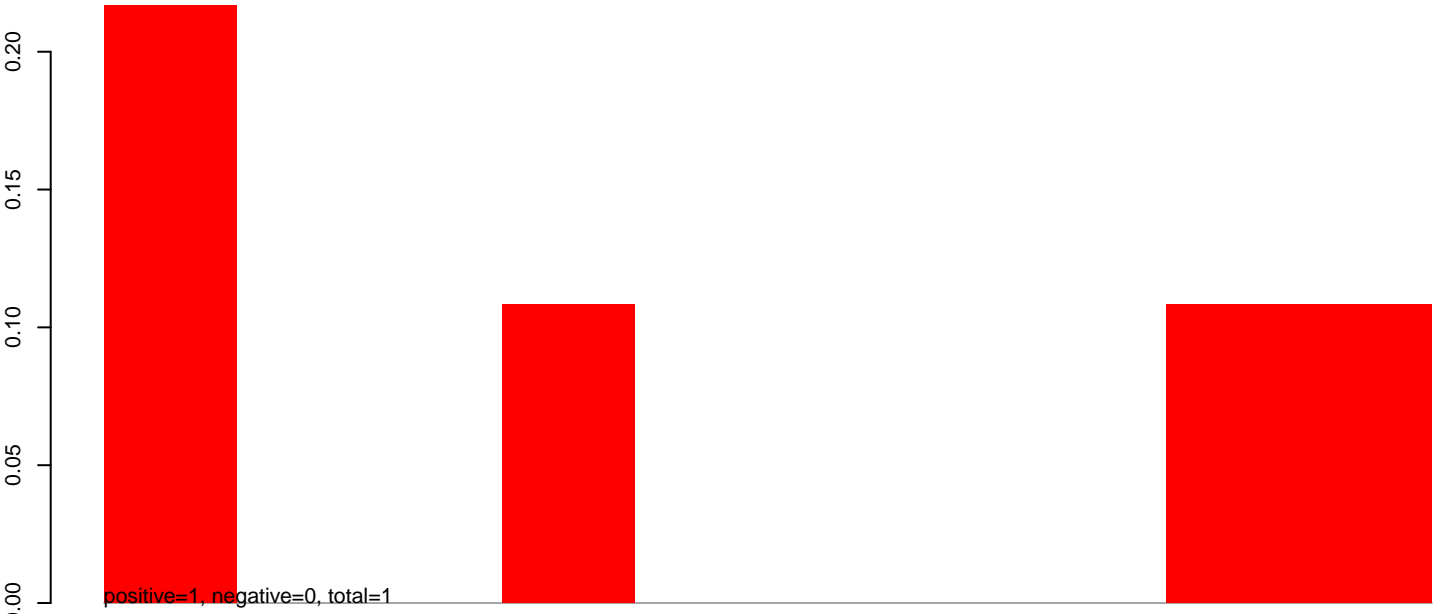
AnGam_Sua5b_cells.24_35.rep



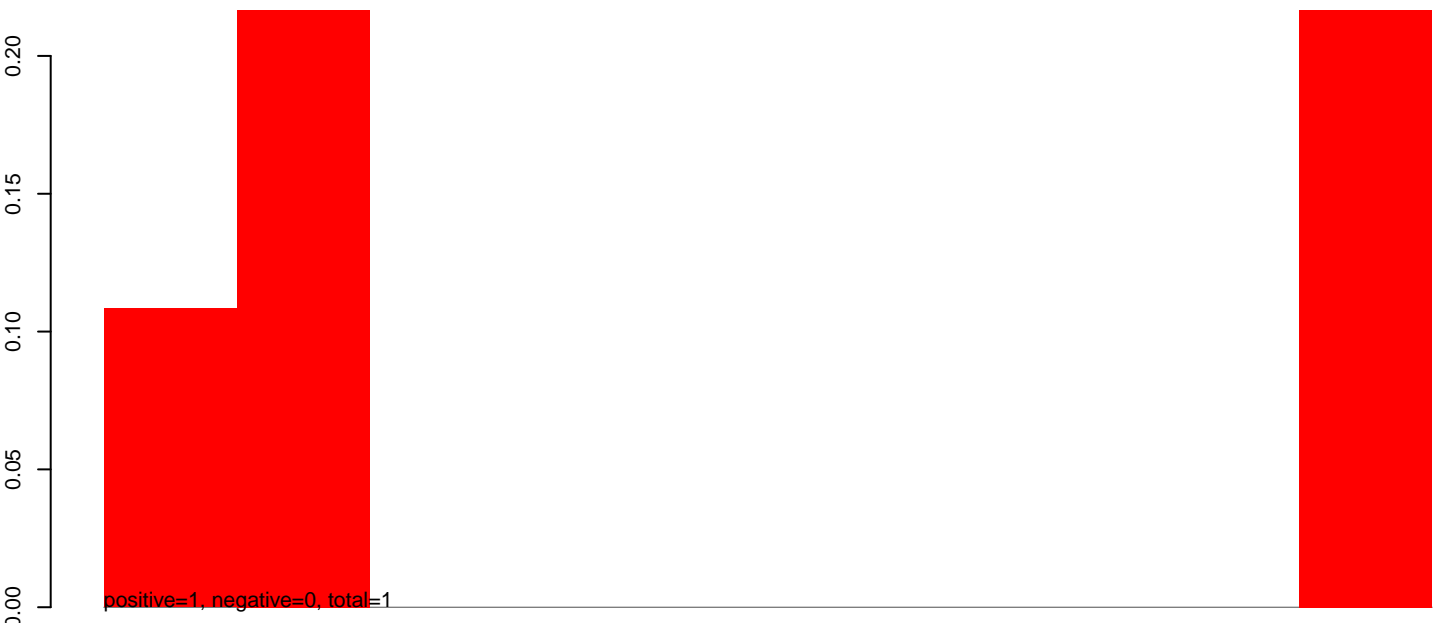
AnGam_Sua5b_cells.rep



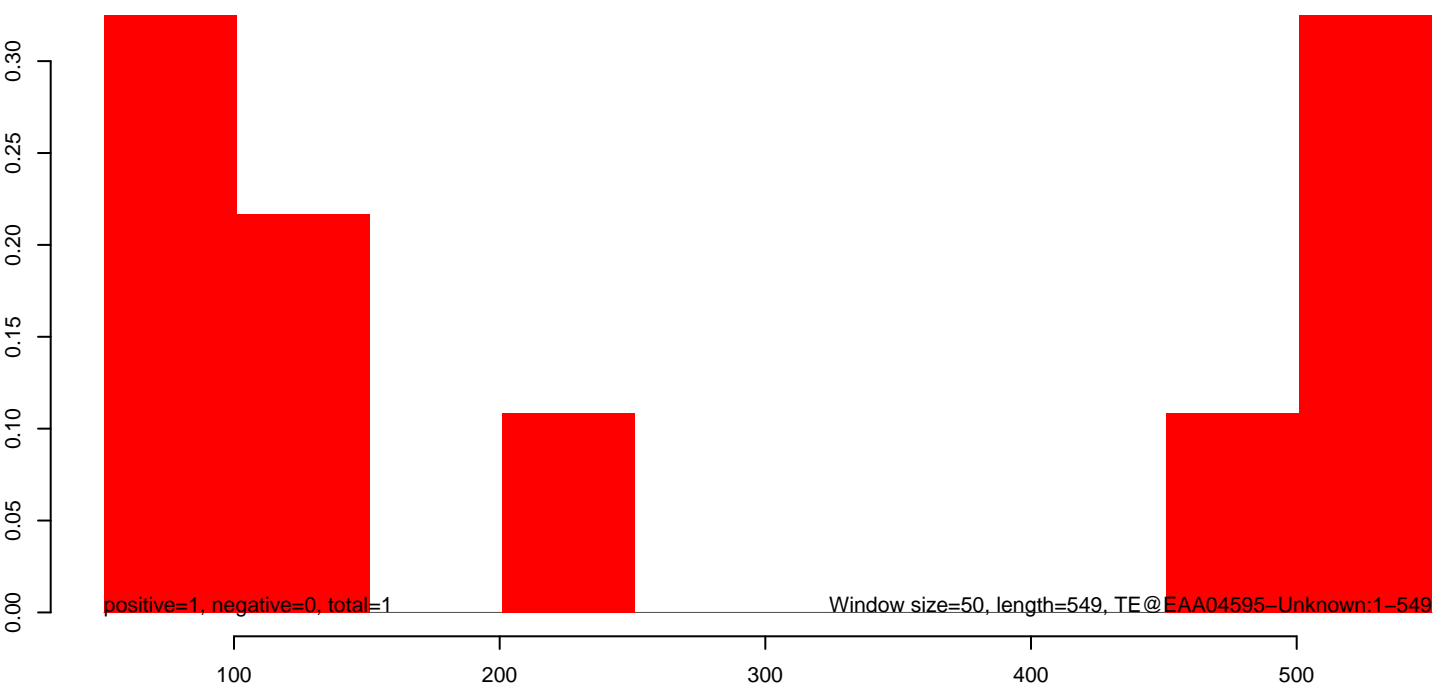
AnGam_Sua5b_cells.18_23.rep



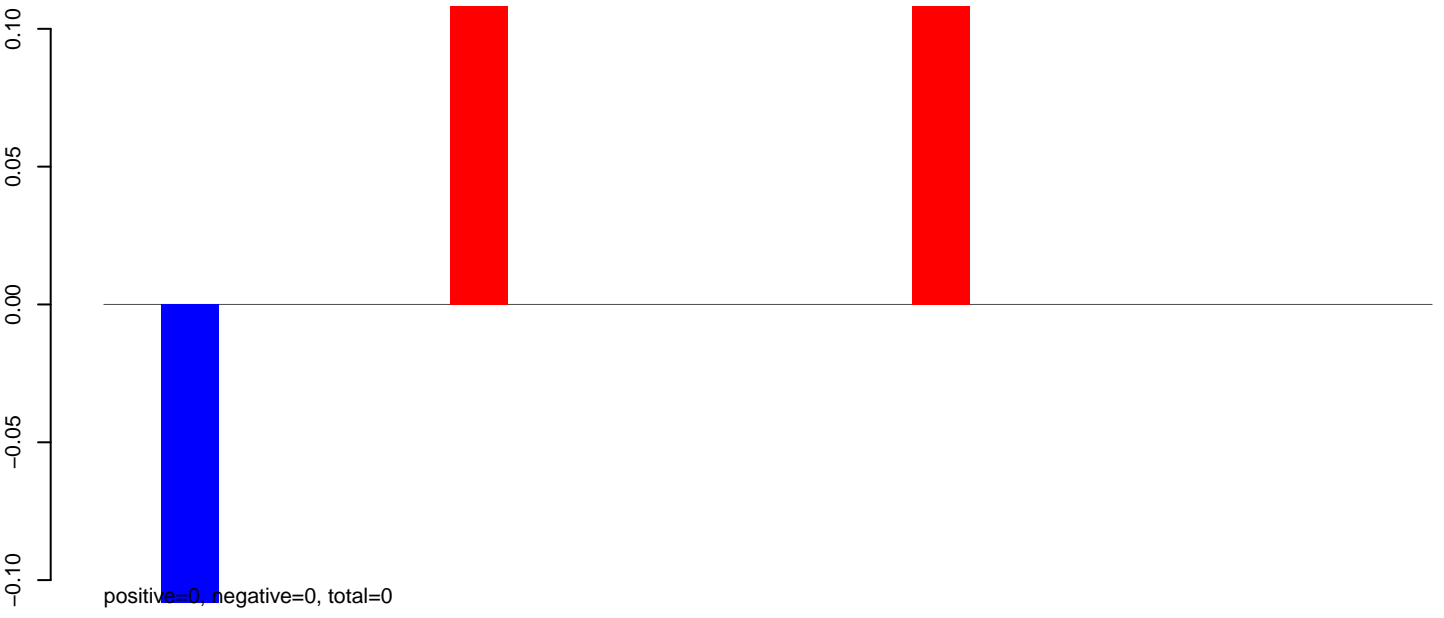
AnGam_Sua5b_cells.24_35.rep



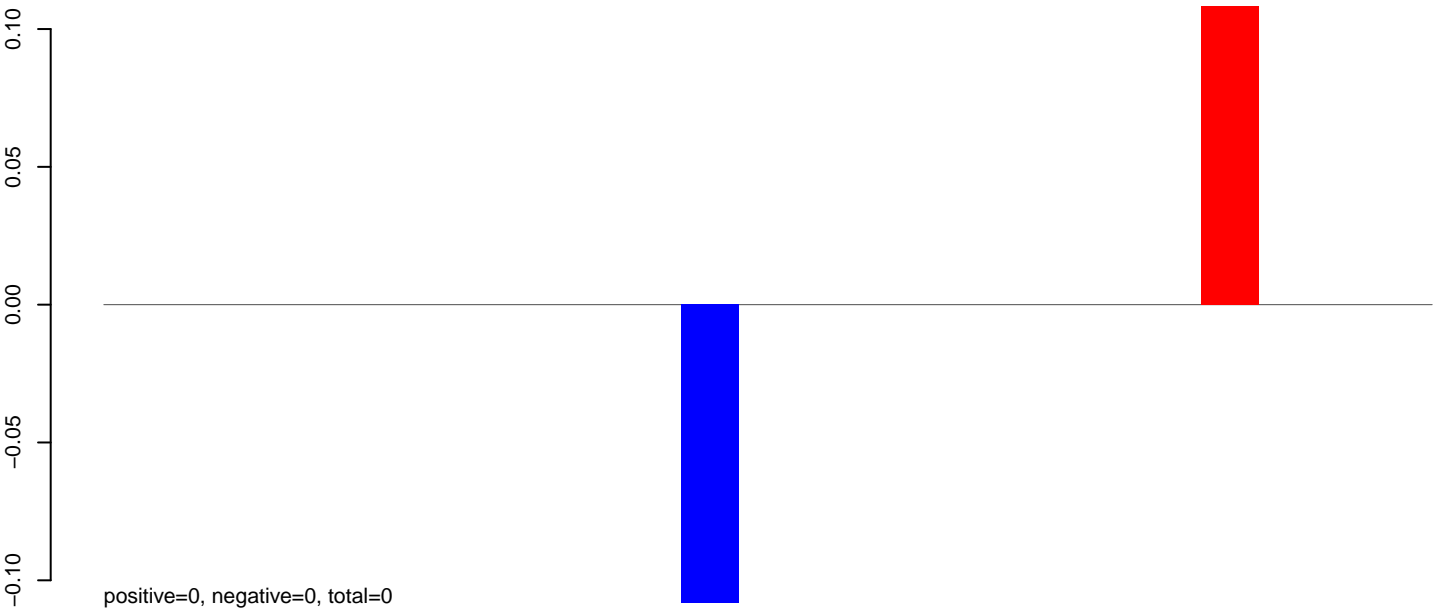
AnGam_Sua5b_cells.rep



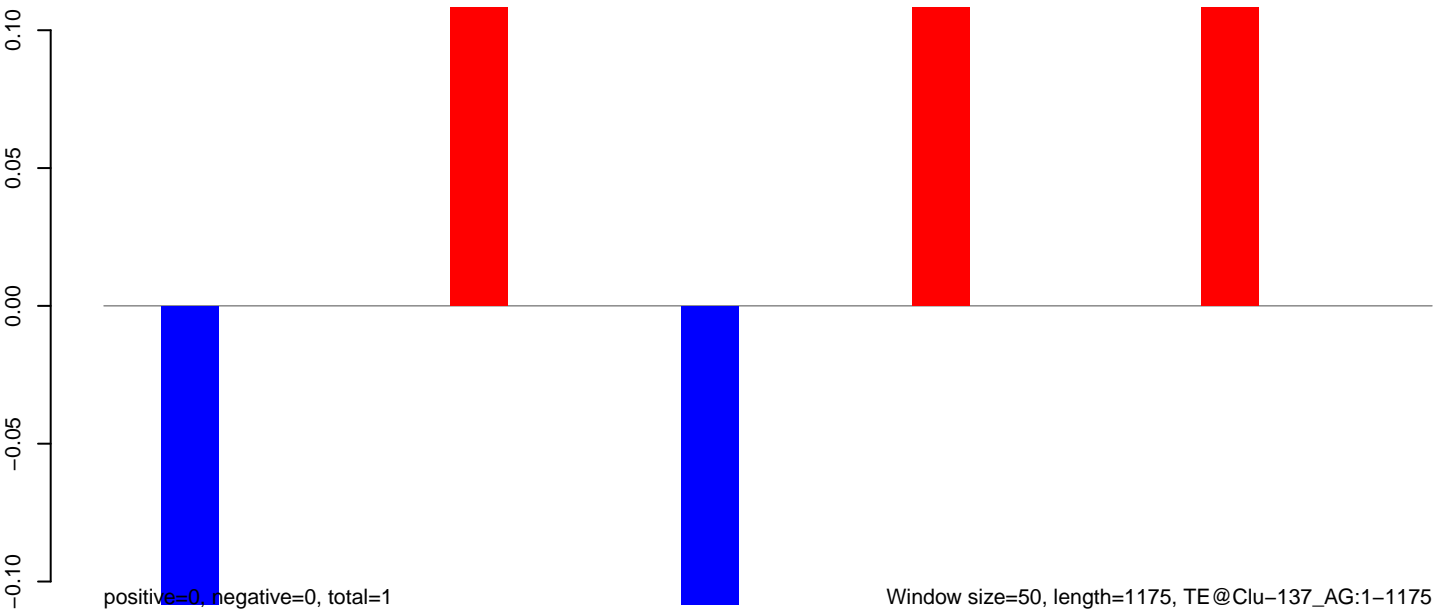
AnGam_Sua5b_cells.18_23.rep



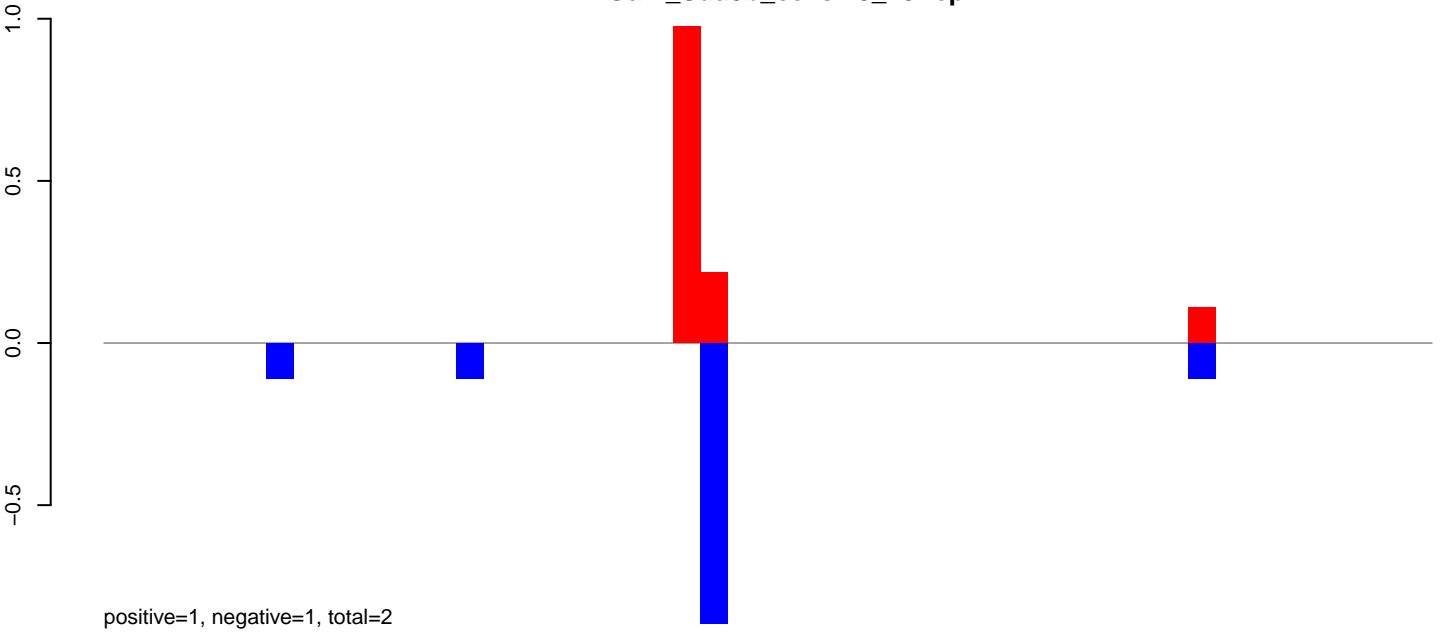
AnGam_Sua5b_cells.24_35.rep



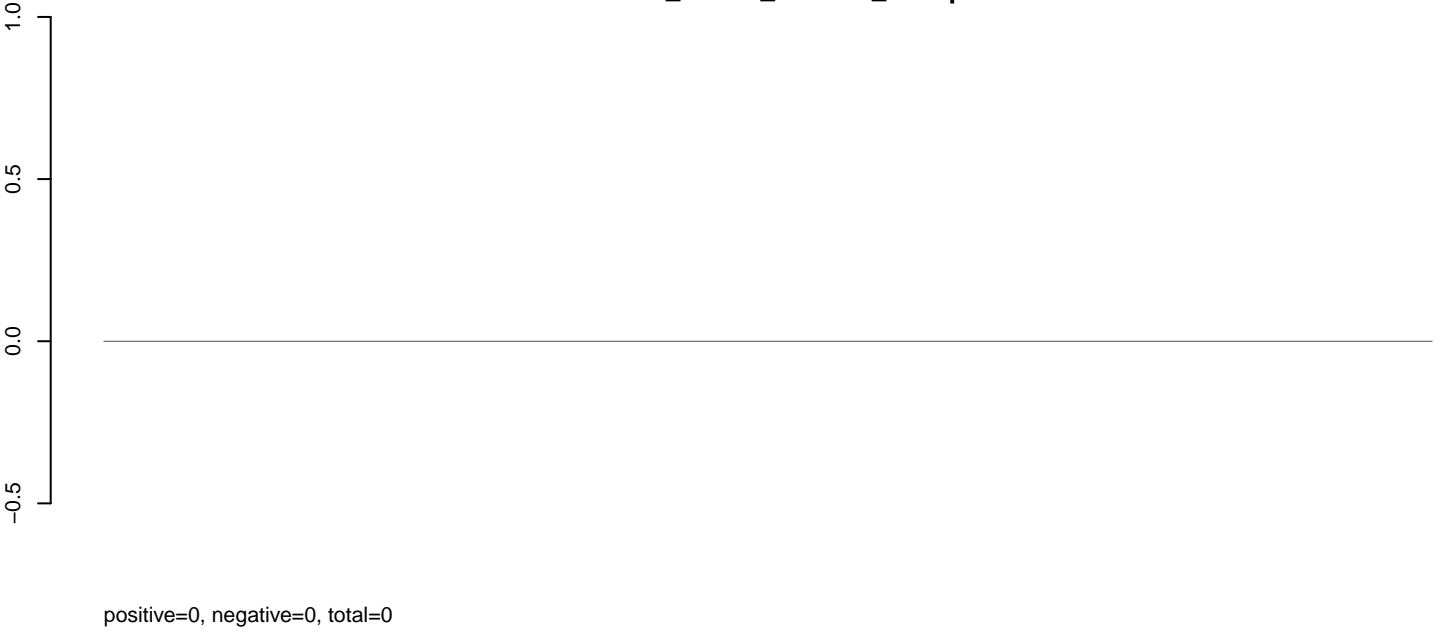
AnGam_Sua5b_cells.rep



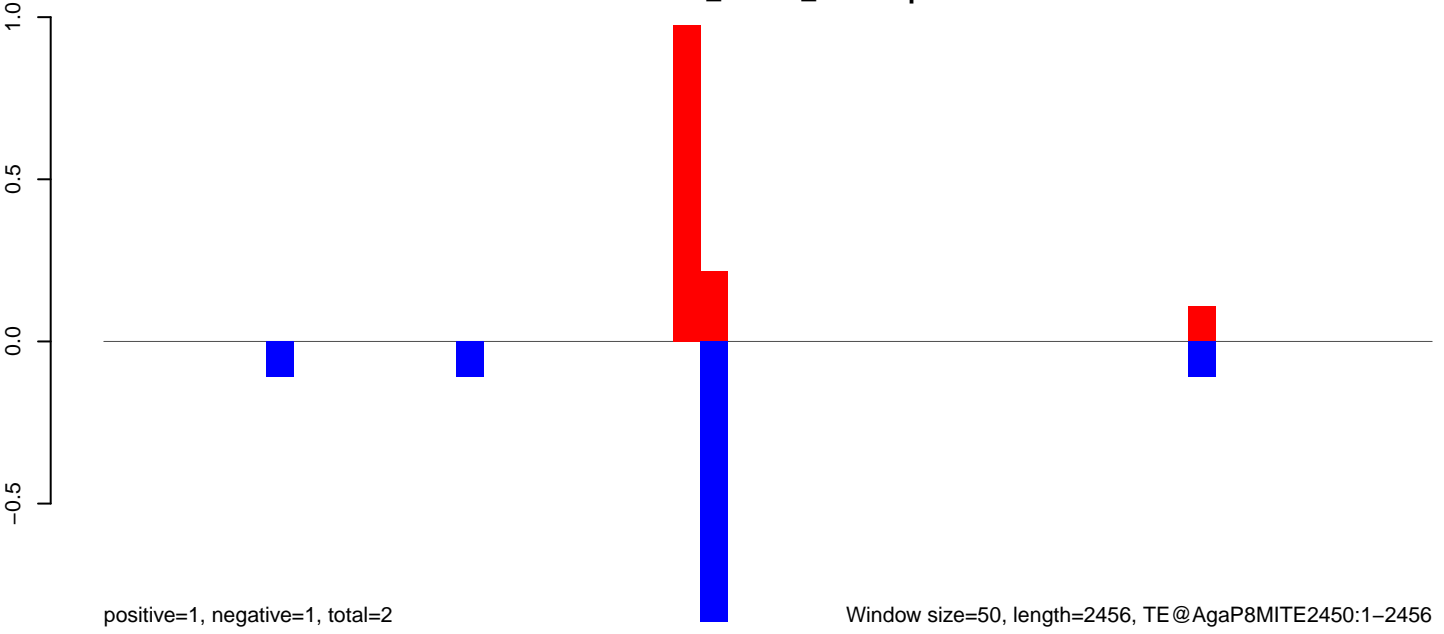
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

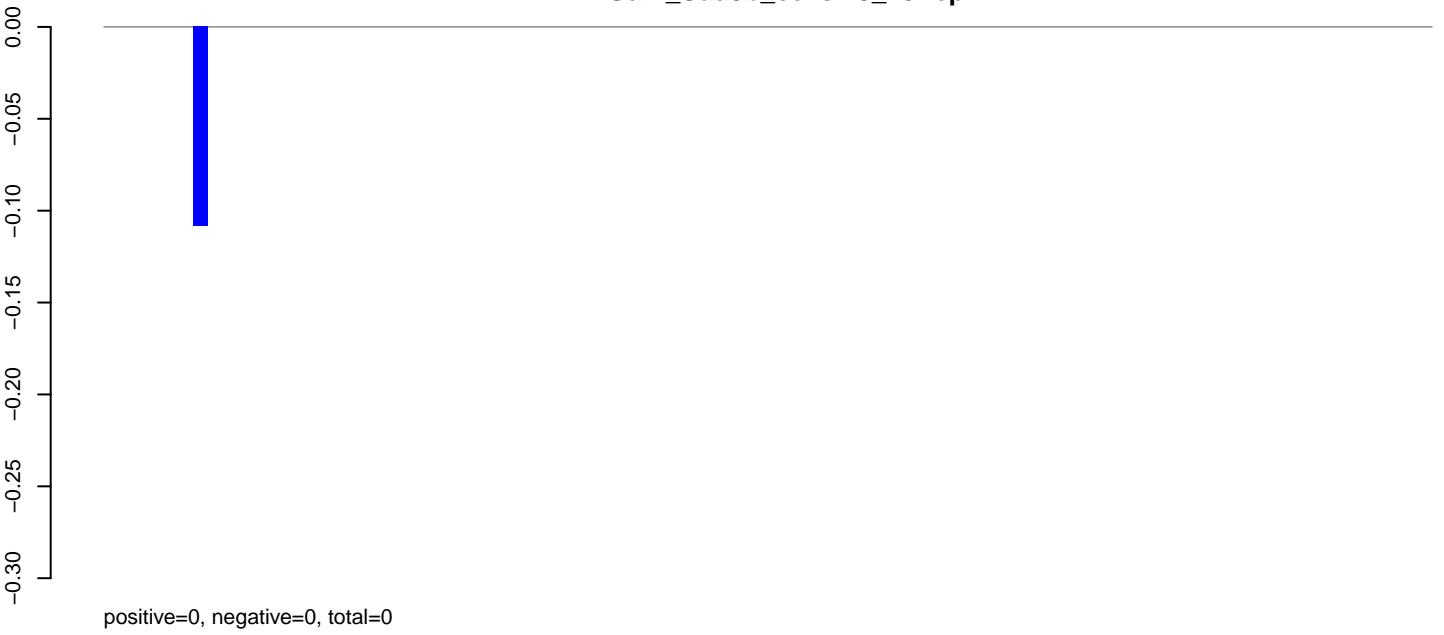


AnGam_Sua5b_cells.rep

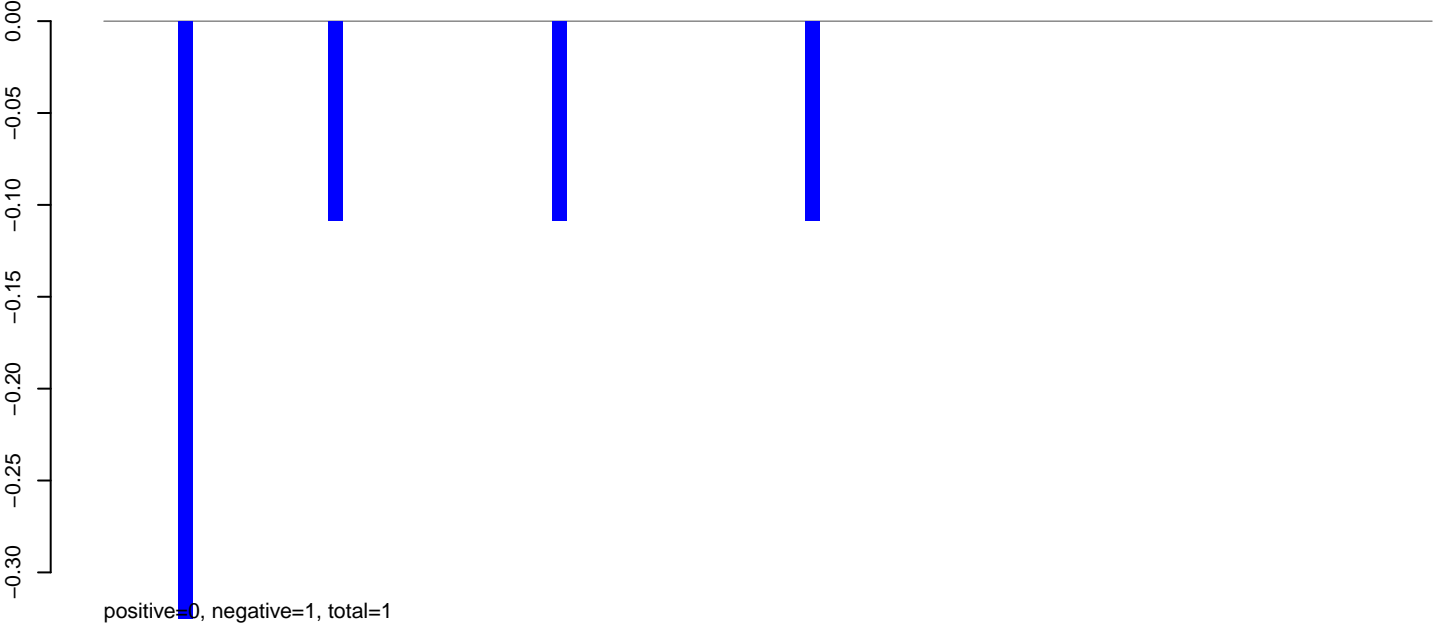


Window size=50, length=2456, TE@AgaP8MITE2450:1-2456

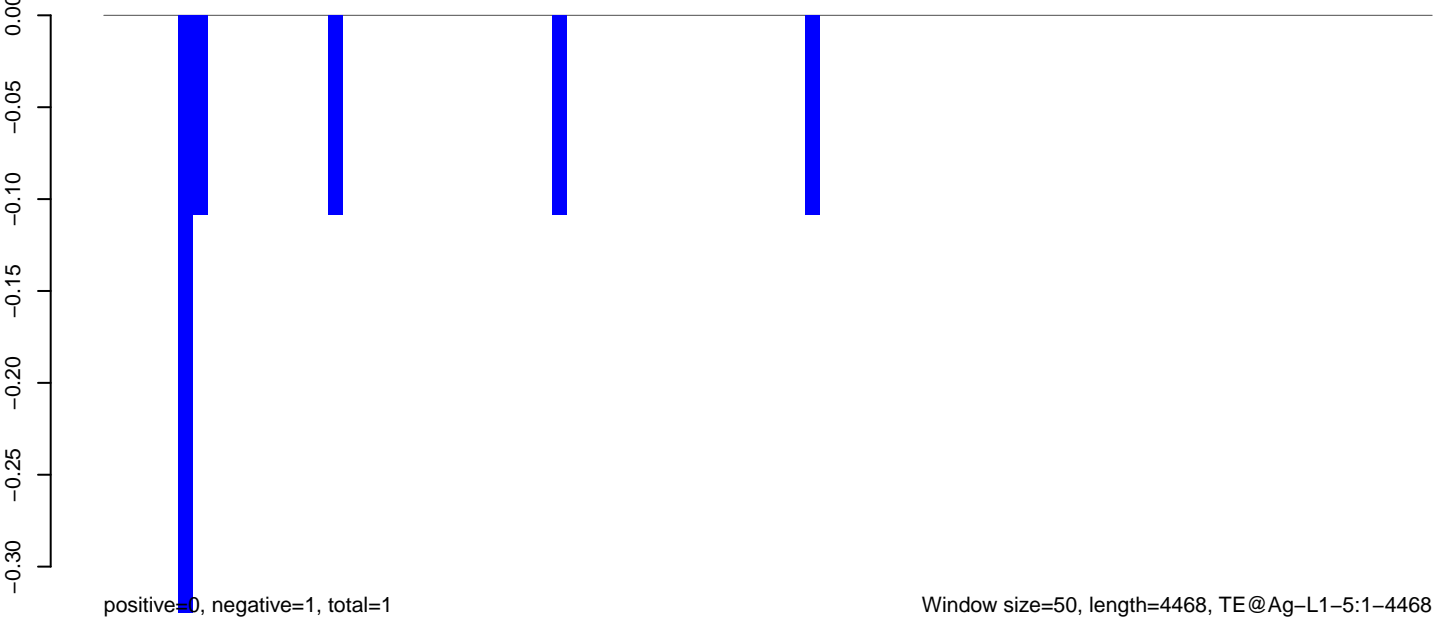
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



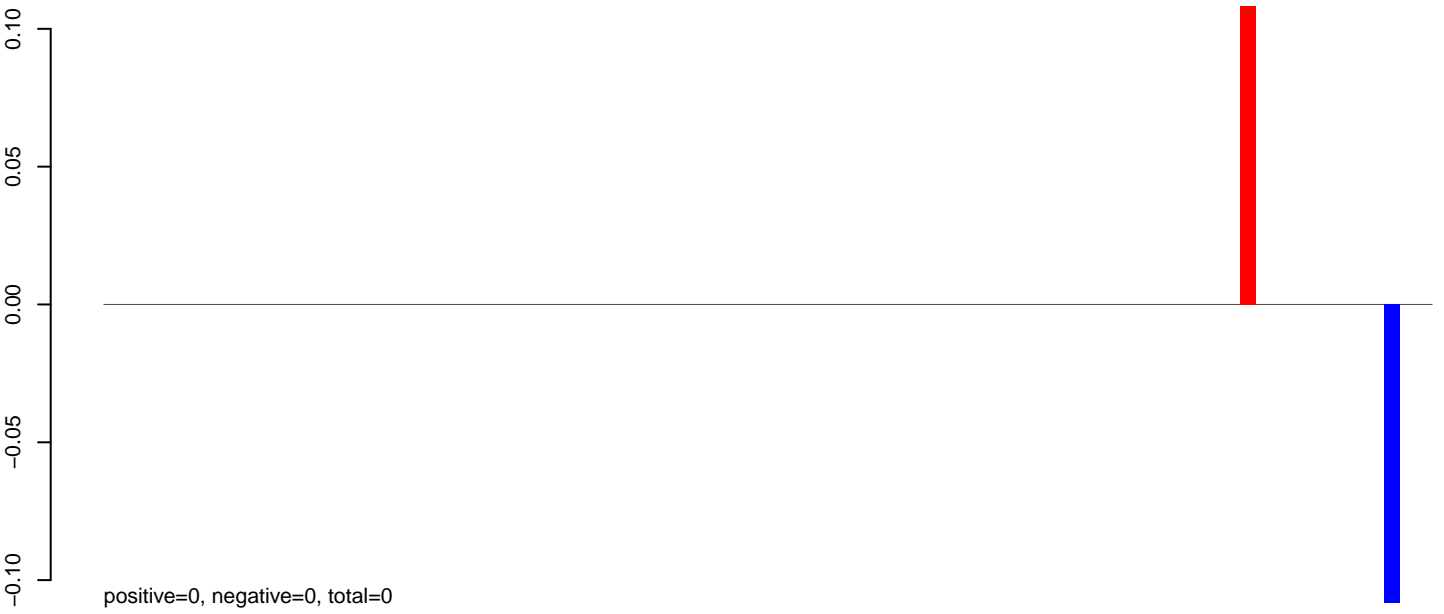
AnGam_Sua5b_cells.rep



Window size=50, length=4468, TE@Ag-L1-5:1-4468

0 1000 2000 3000 4000

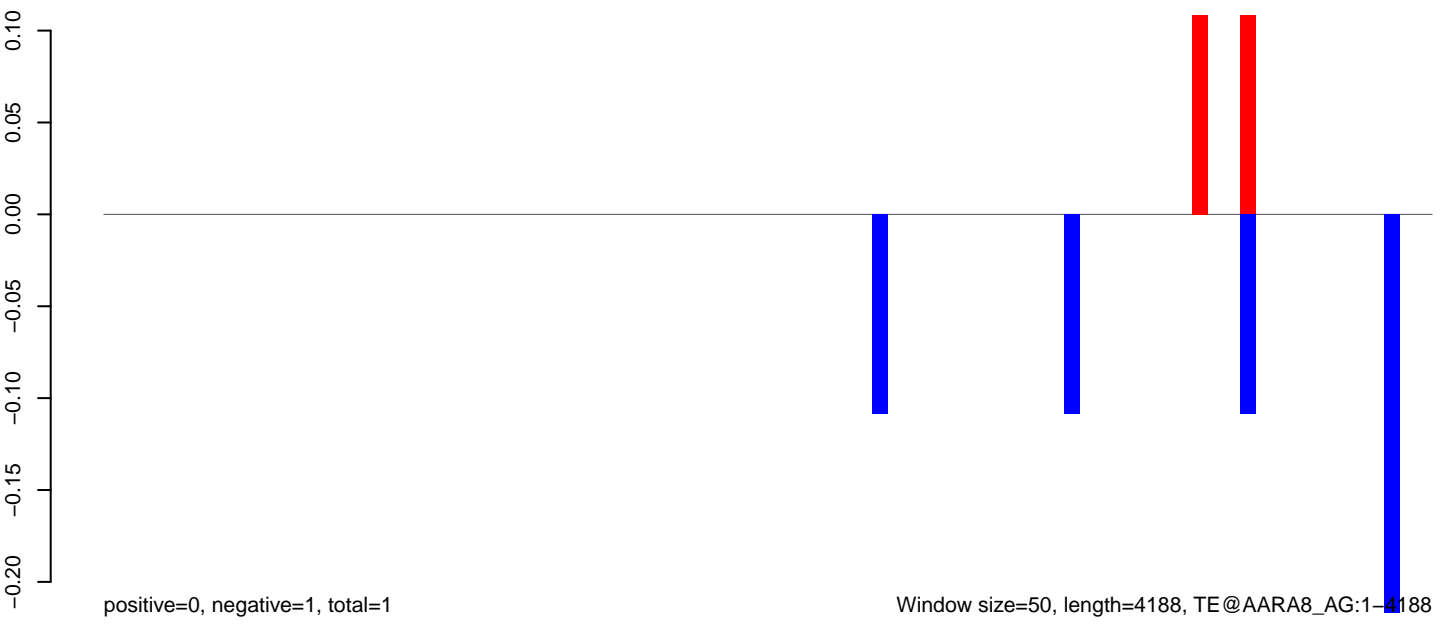
AnGam_Sua5b_cells.18_23.rep



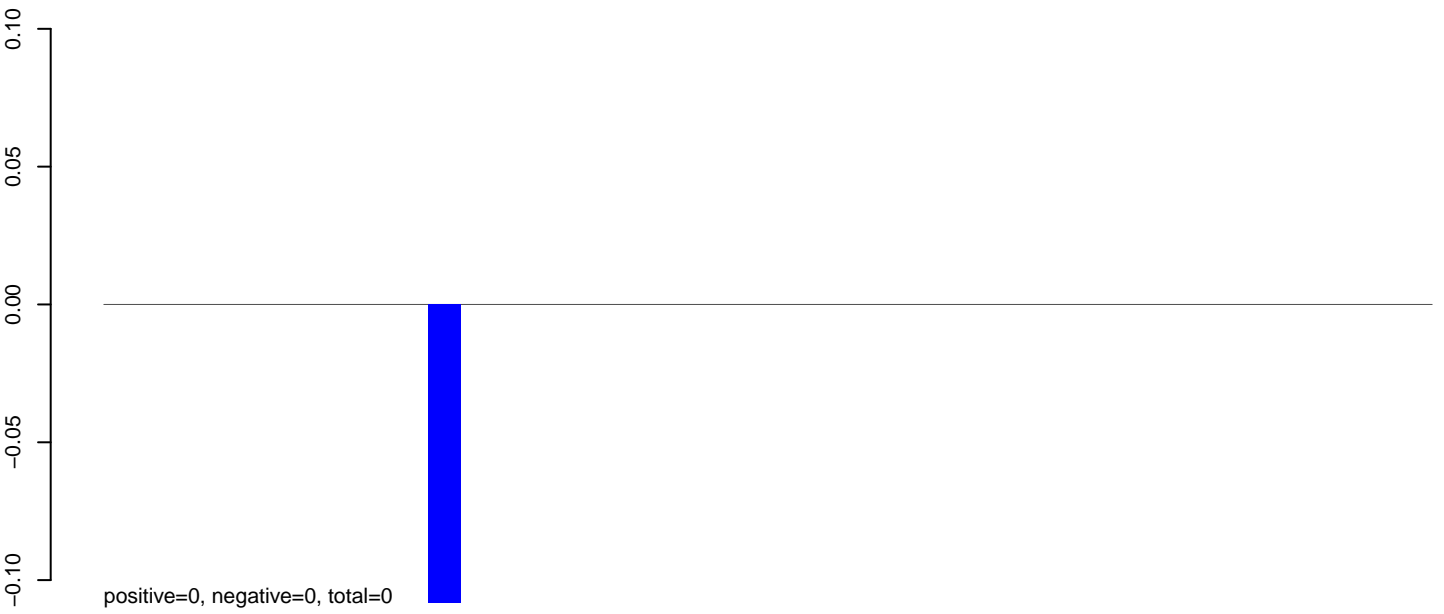
AnGam_Sua5b_cells.24_35.rep



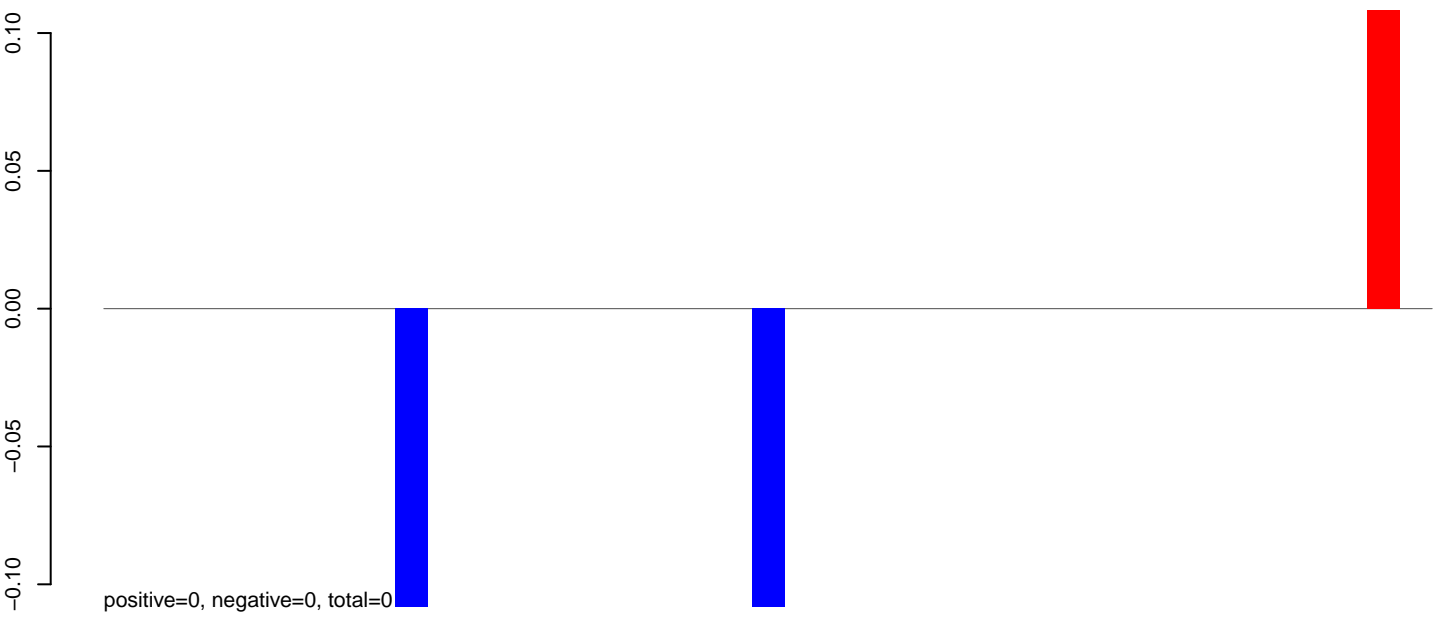
AnGam_Sua5b_cells.rep



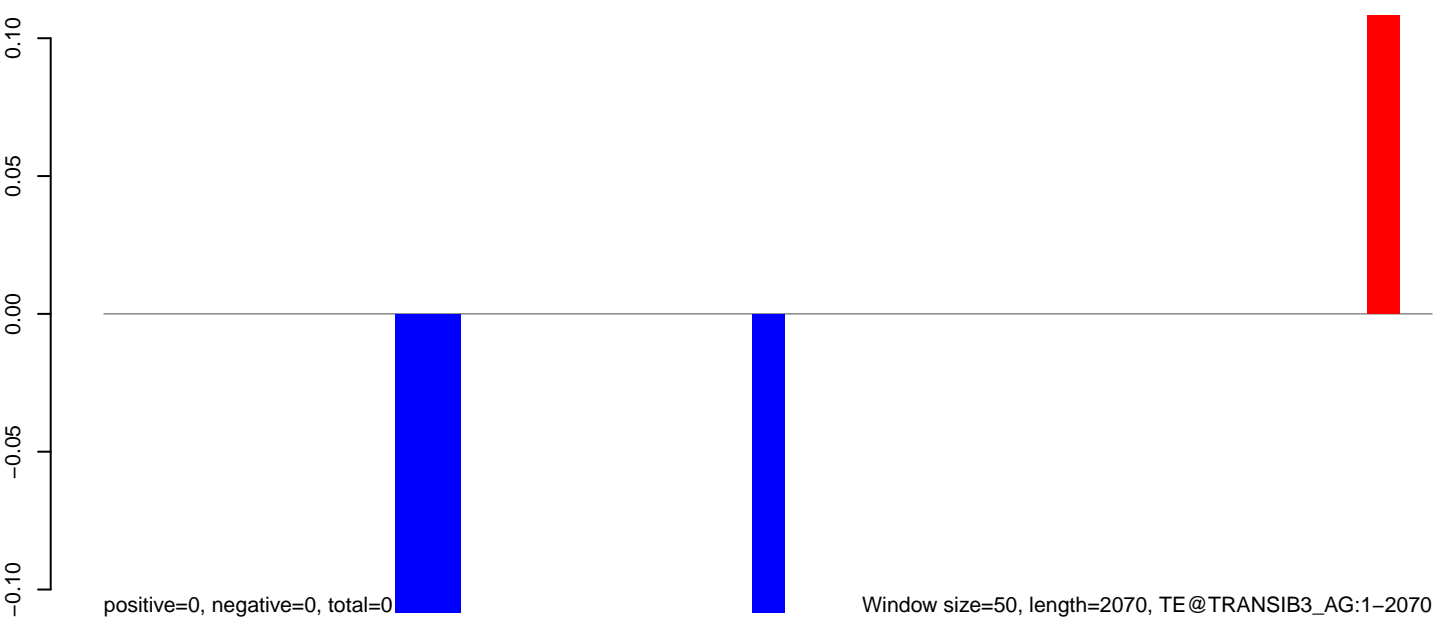
AnGam_Sua5b_cells.18_23.rep



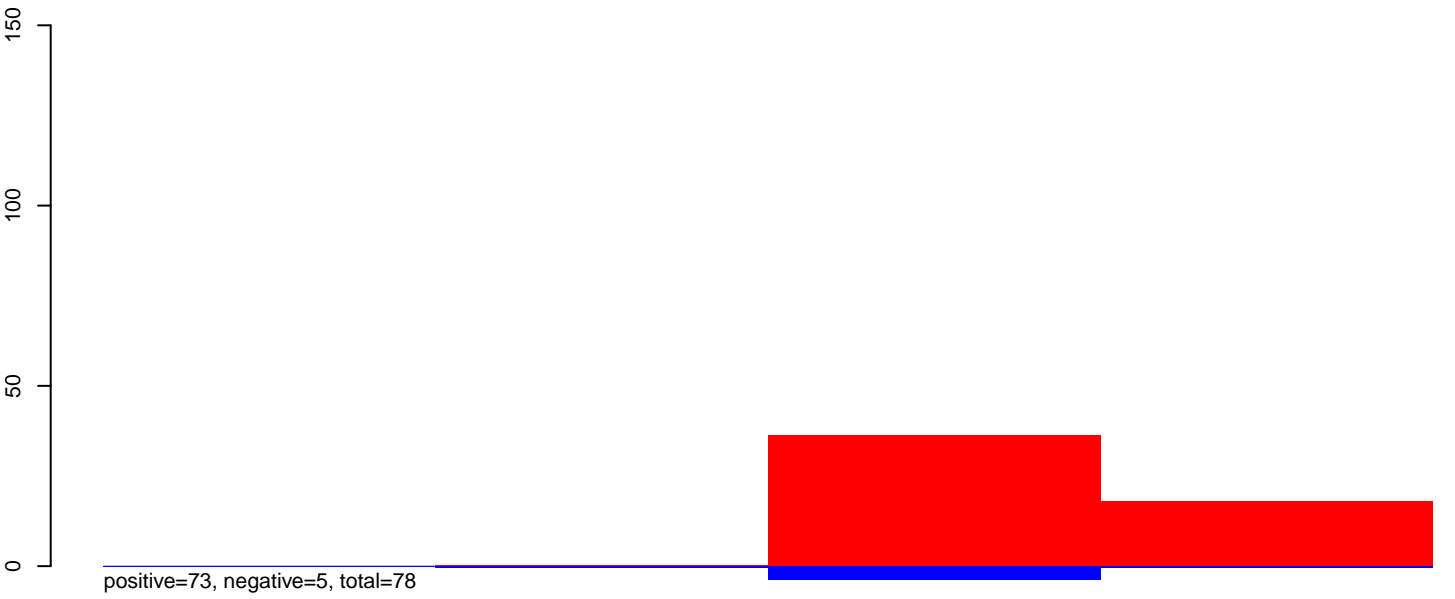
AnGam_Sua5b_cells.24_35.rep



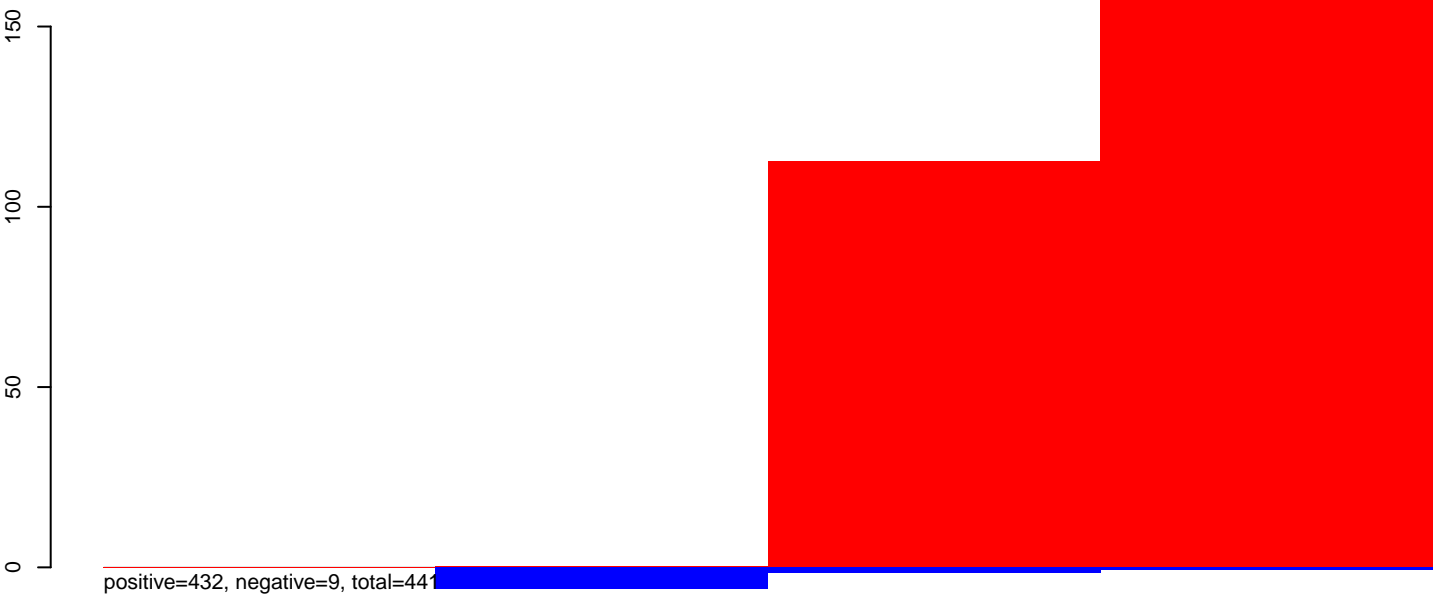
AnGam_Sua5b_cells.rep



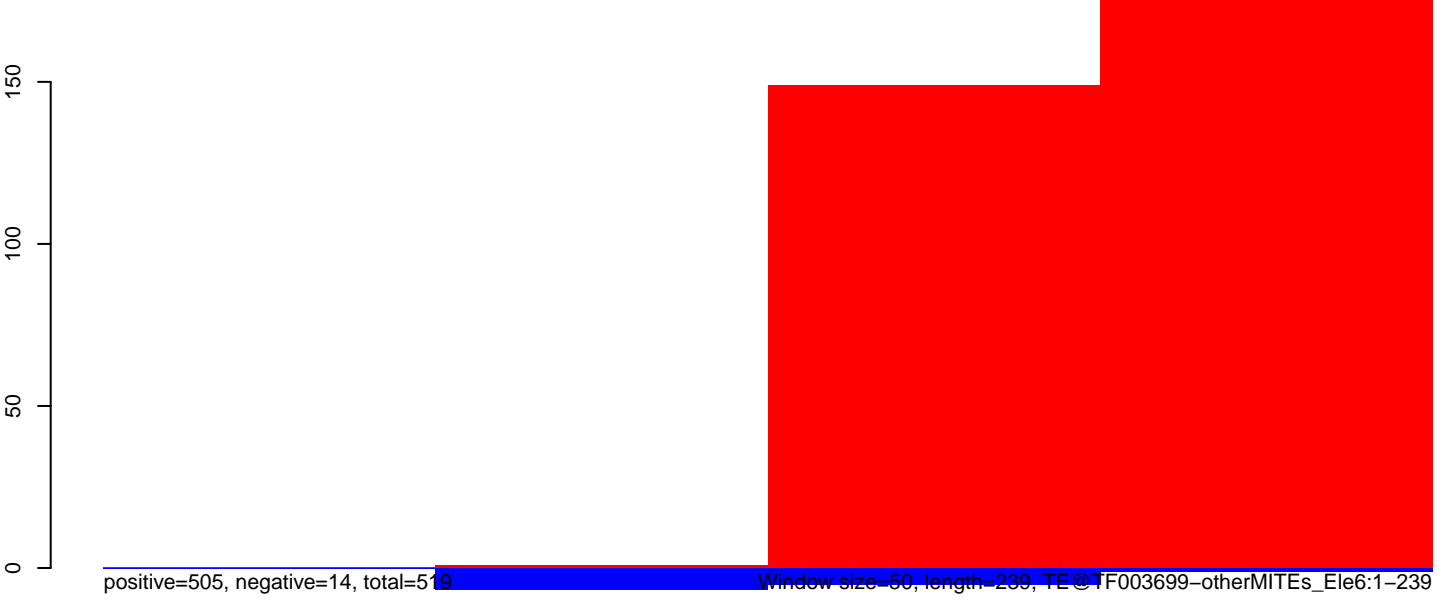
AnGam_Sua5b_cells.18_23.rep



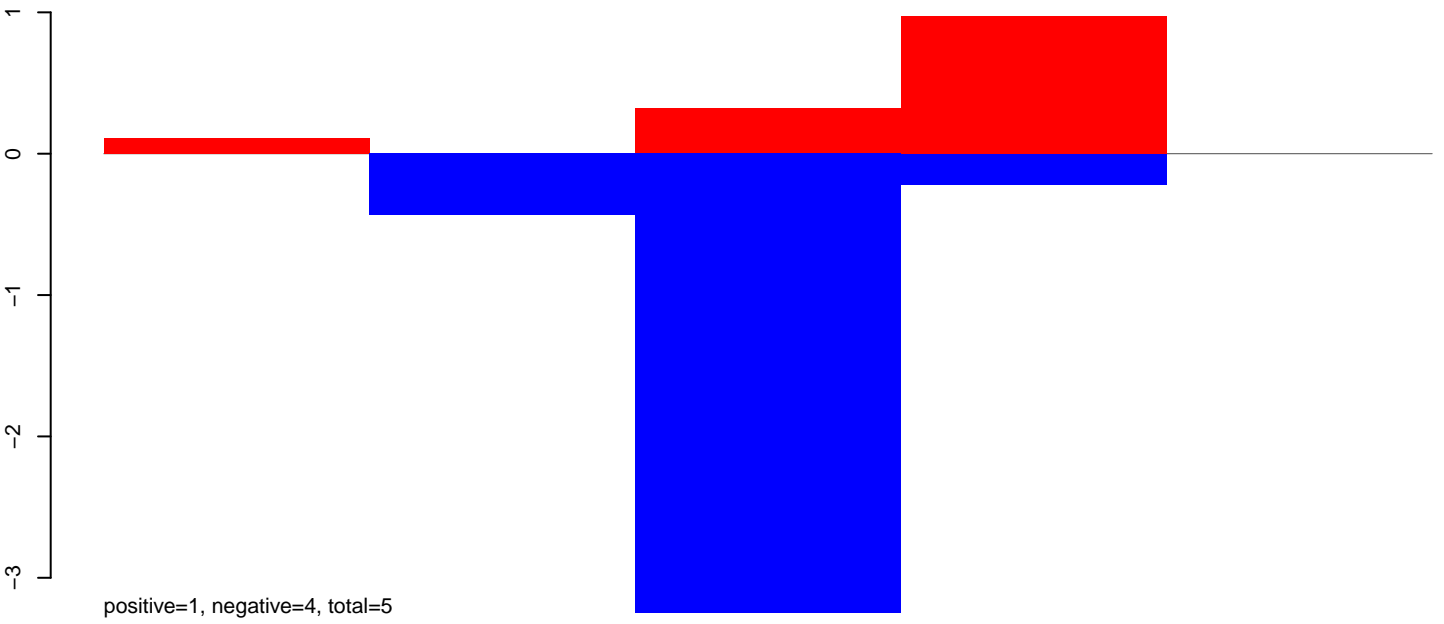
AnGam_Sua5b_cells.24_35.rep



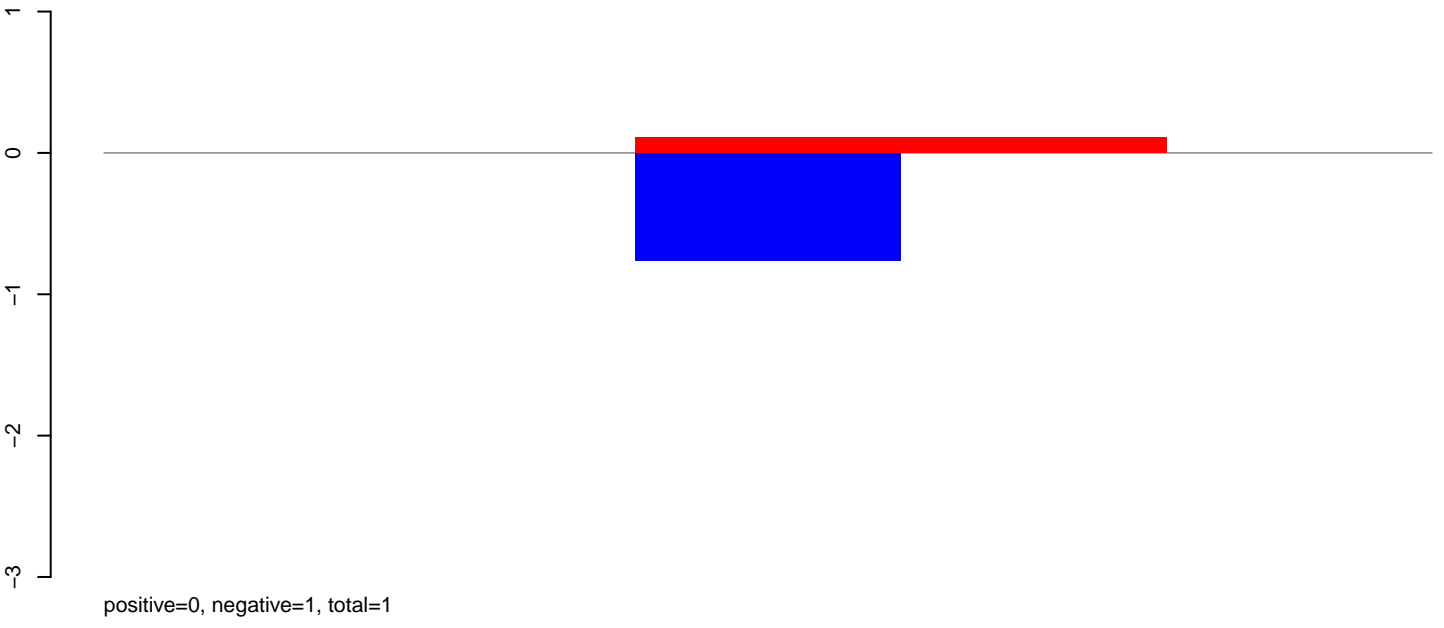
AnGam_Sua5b_cells.rep



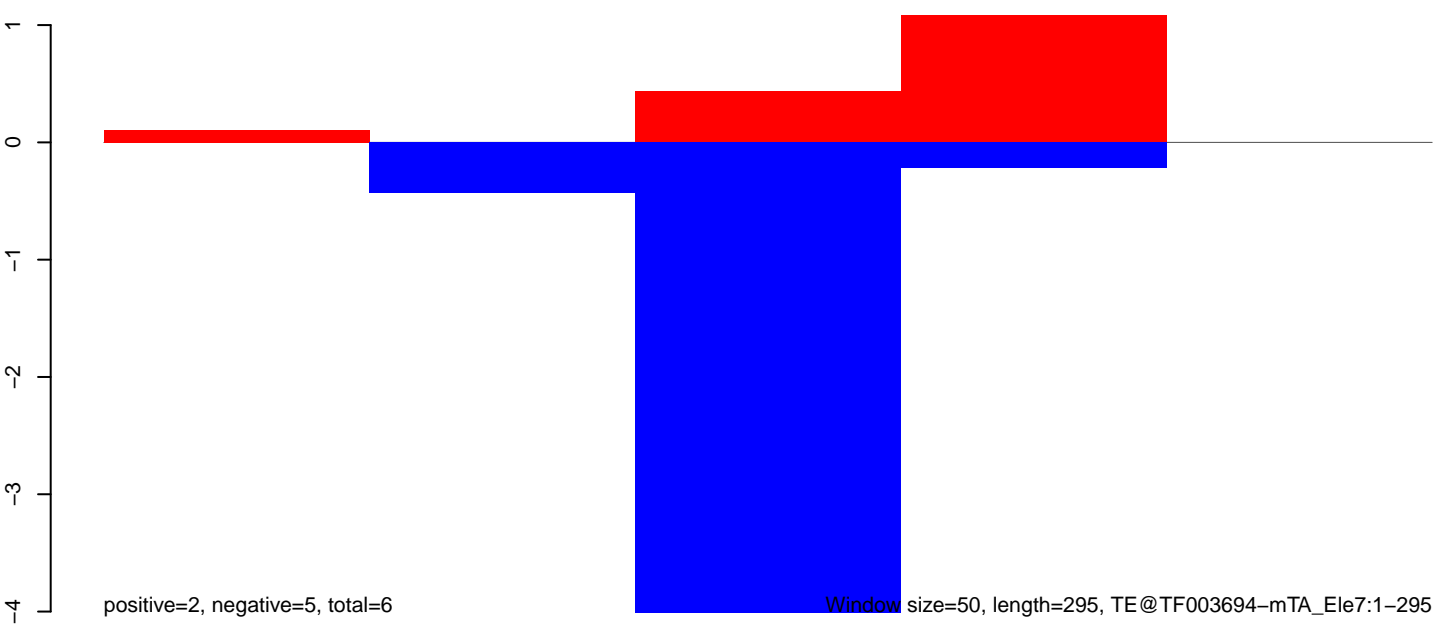
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

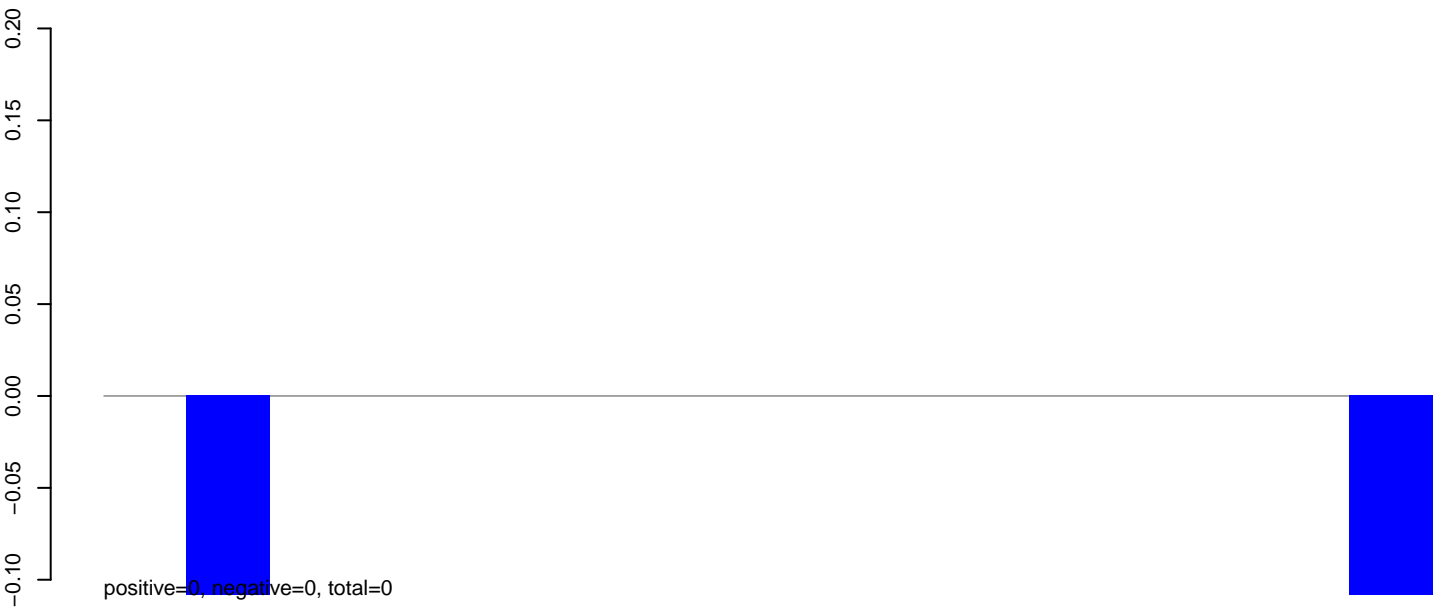


AnGam_Sua5b_cells.rep

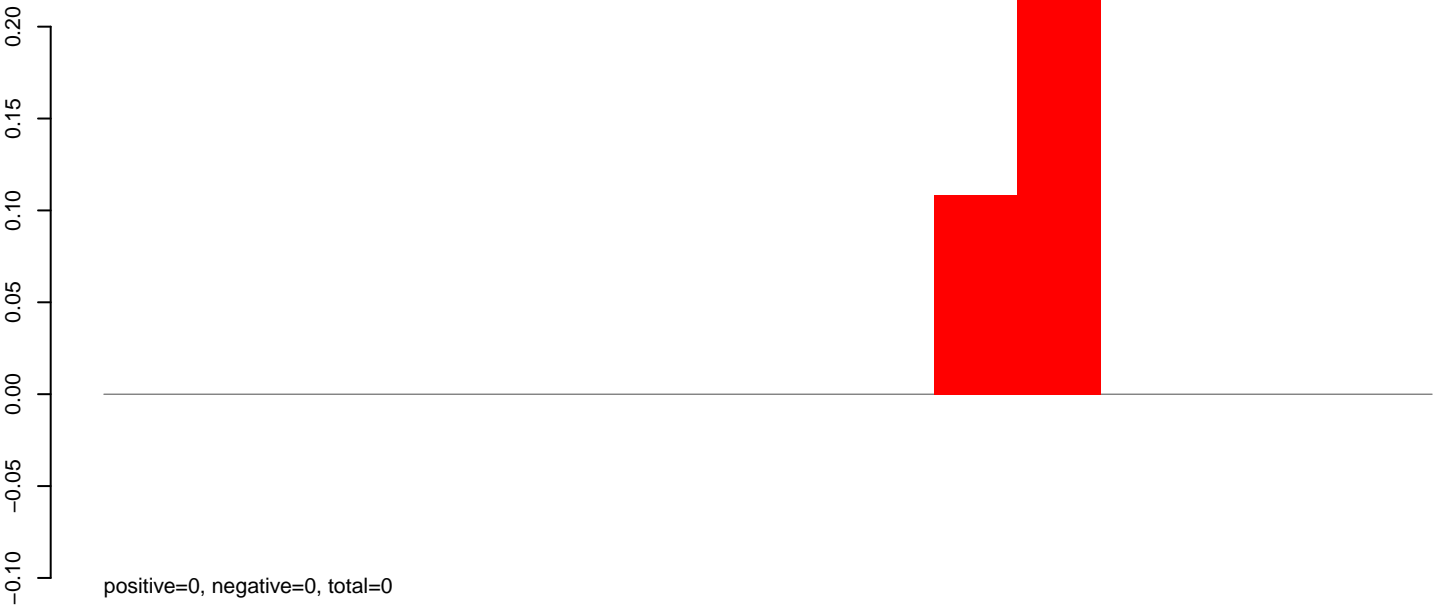


Window size=50, length=295, TE@TF003694-mTA_Ele7:1-295

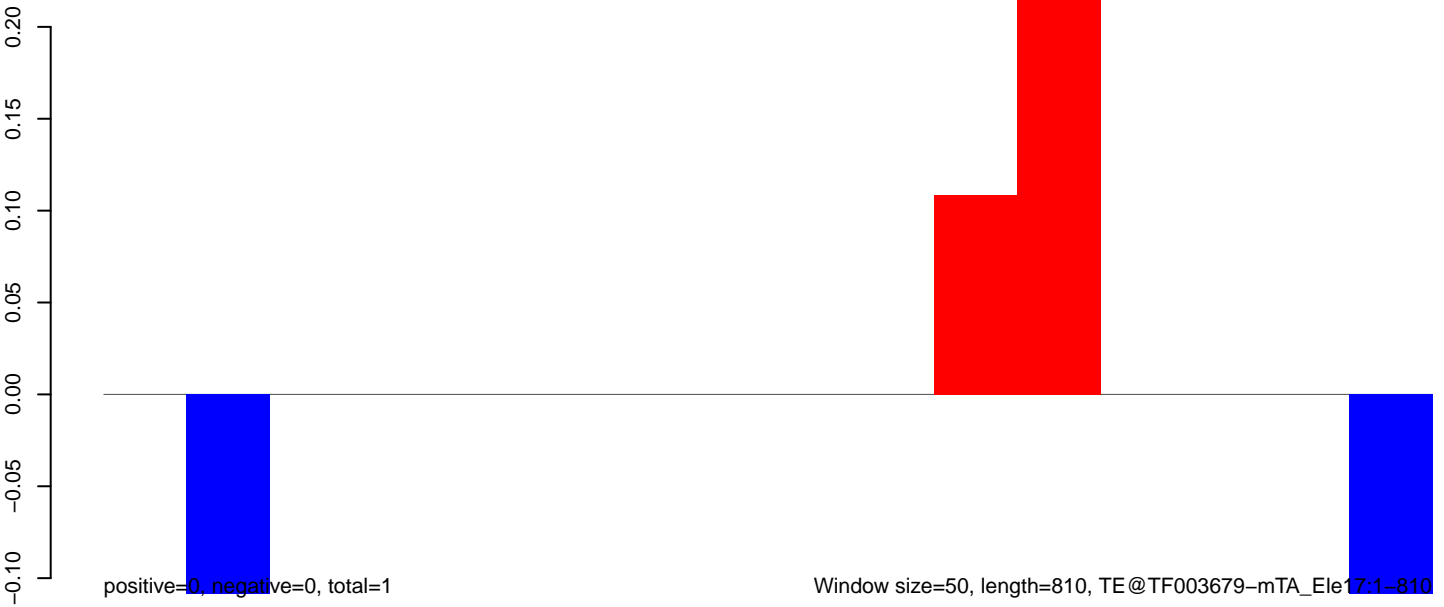
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



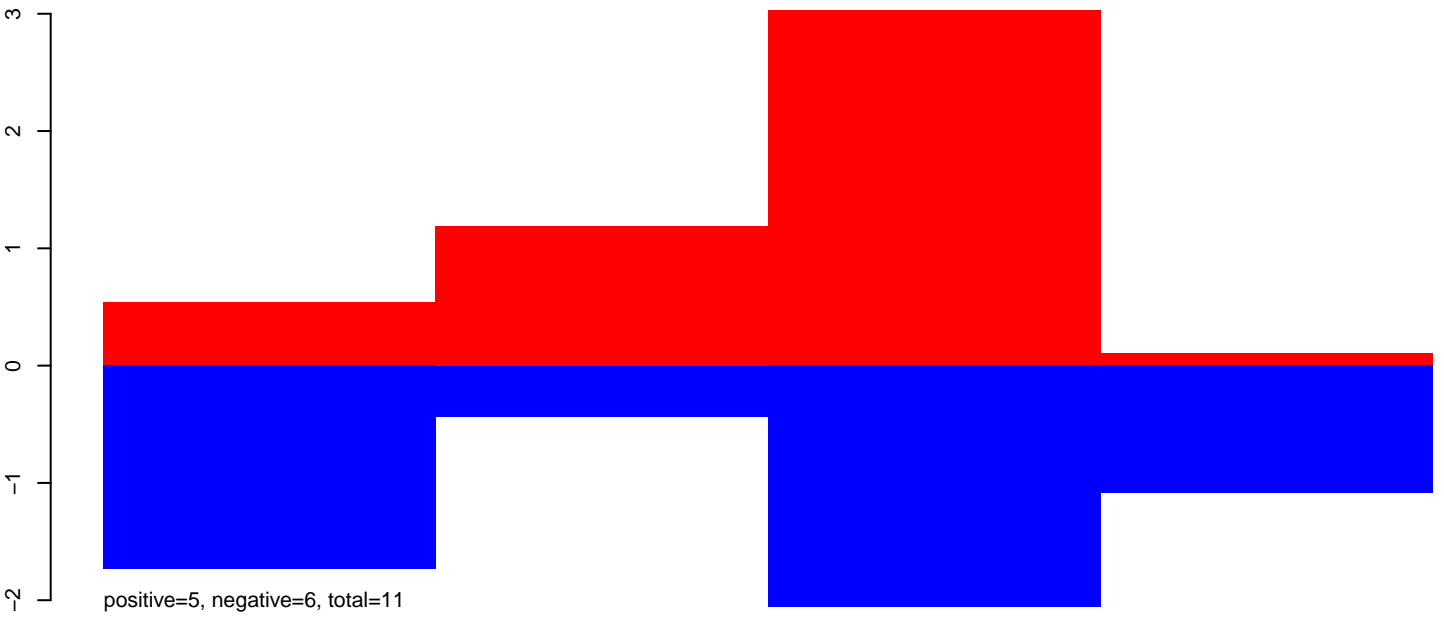
200

400

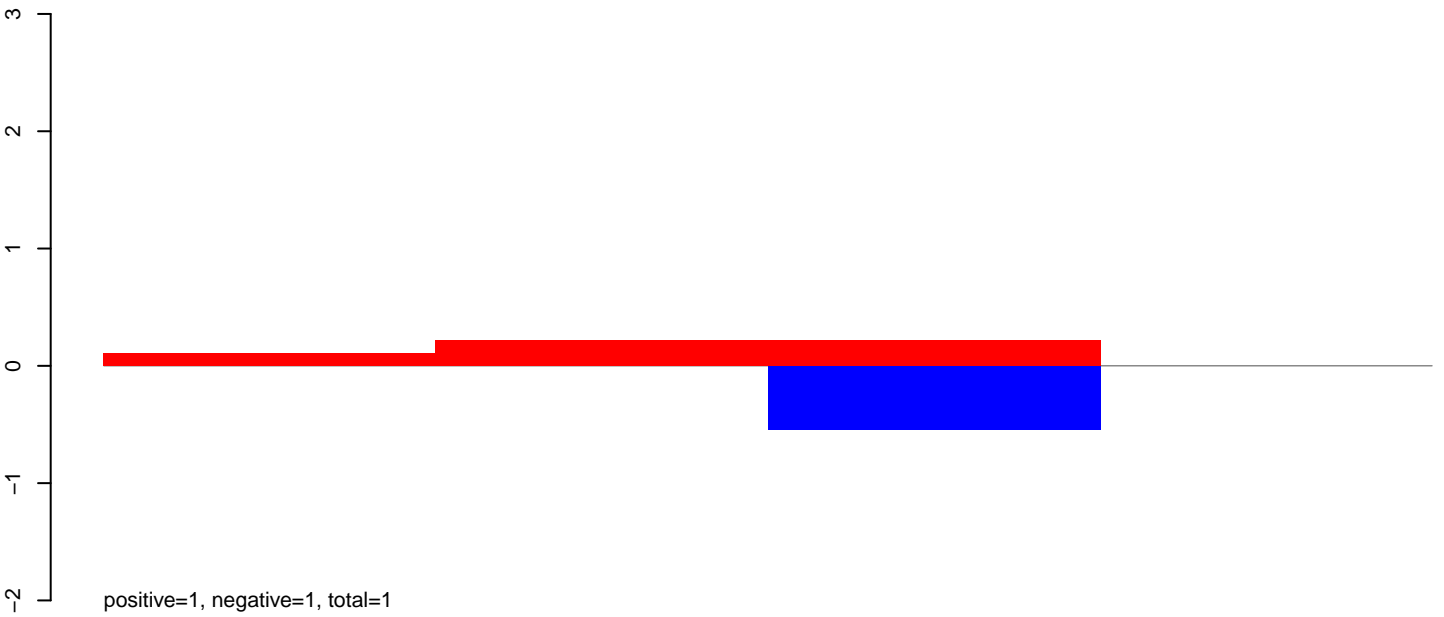
600

800

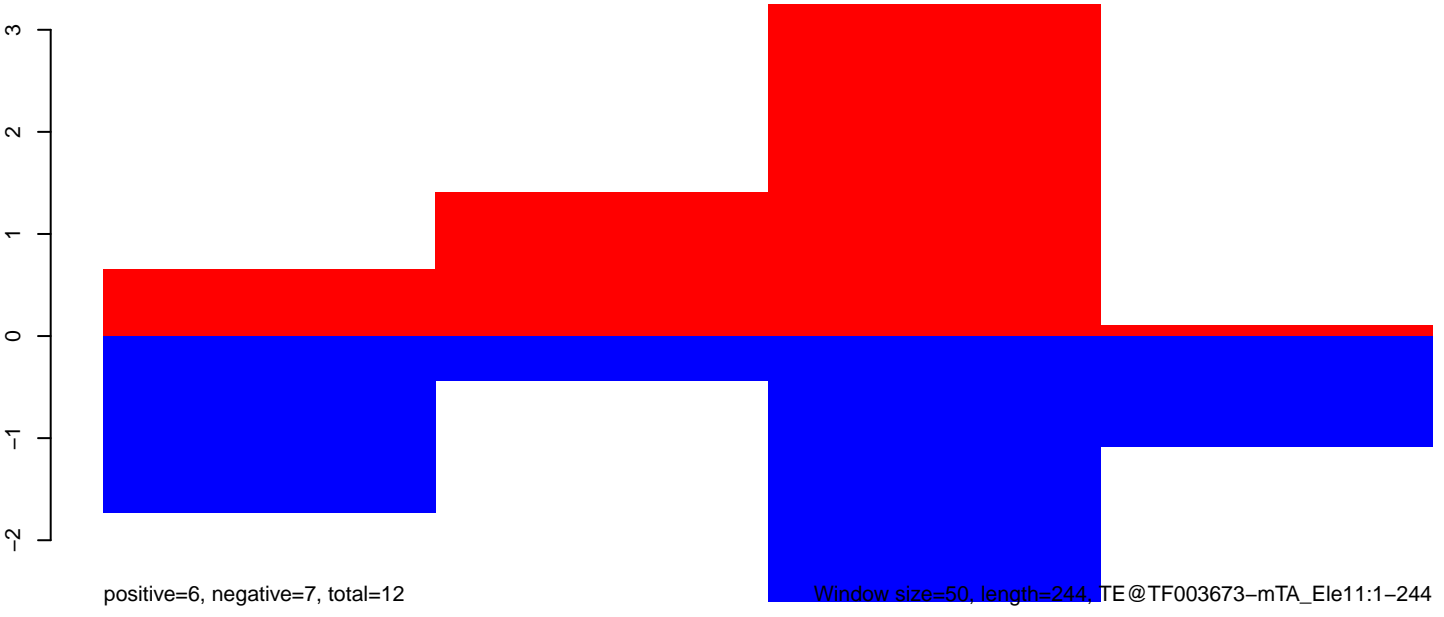
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



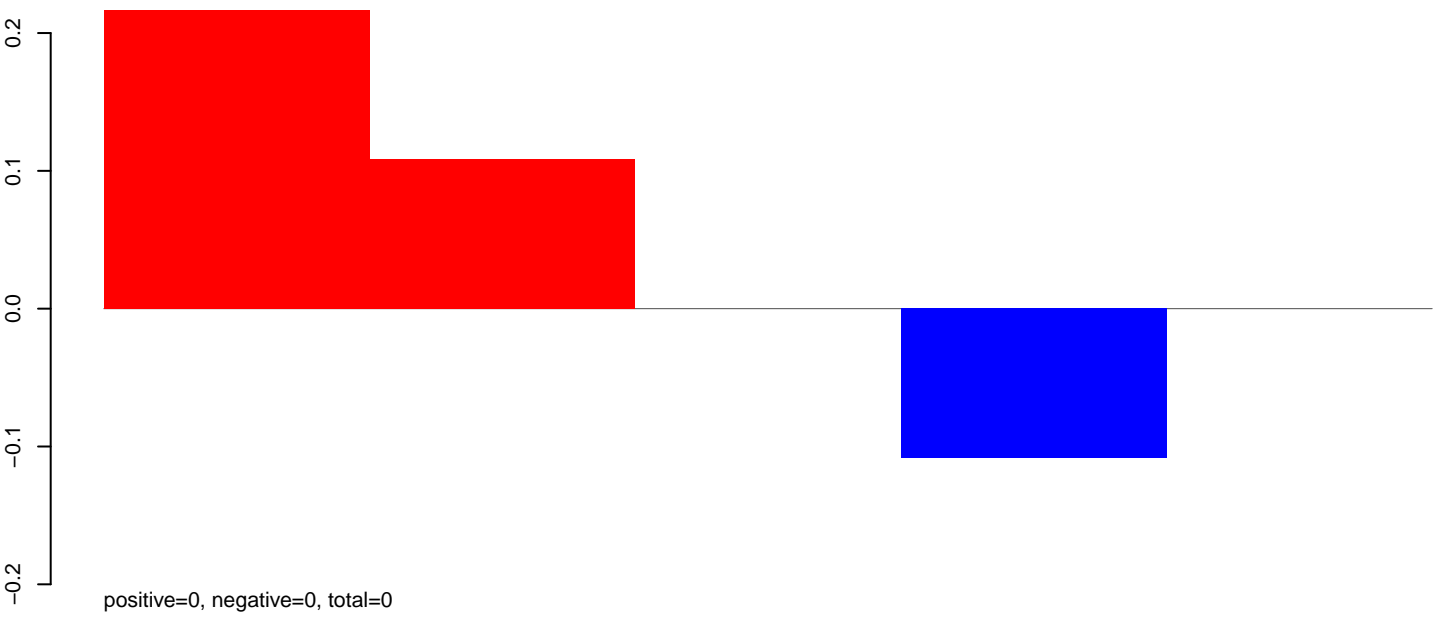
AnGam_Sua5b_cells.rep



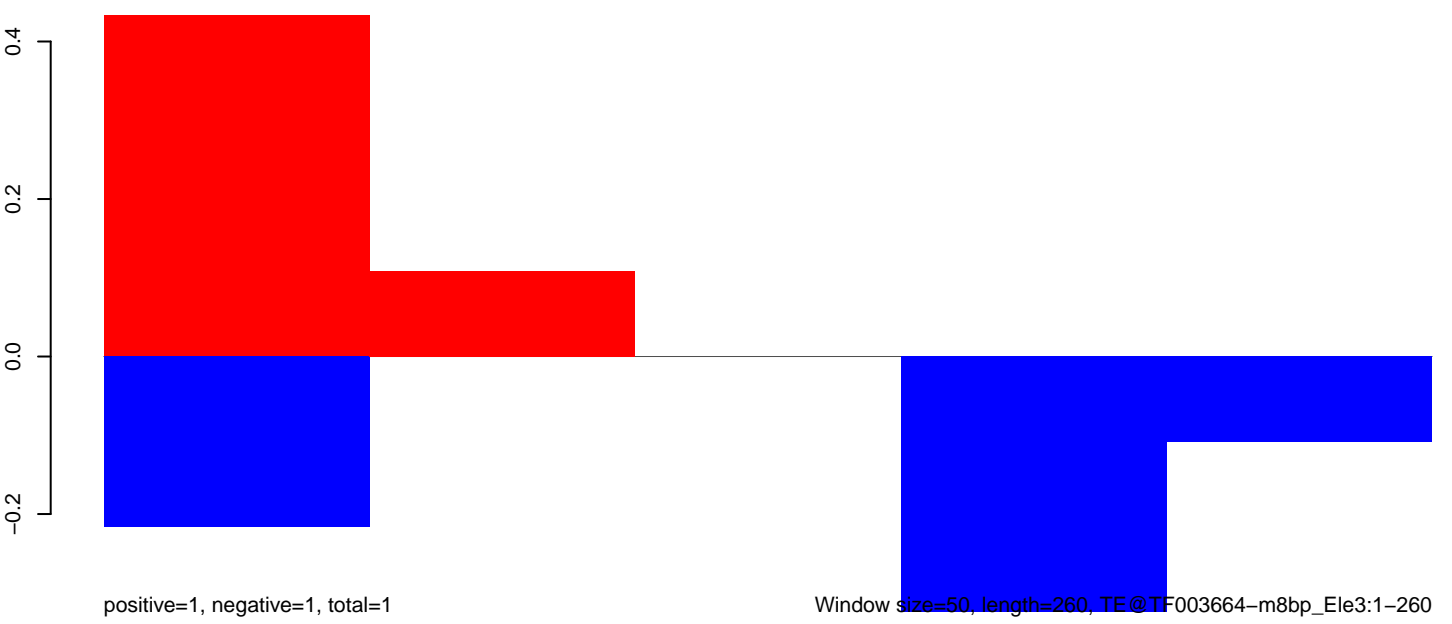
AnGam_Sua5b_cells.18_23.rep



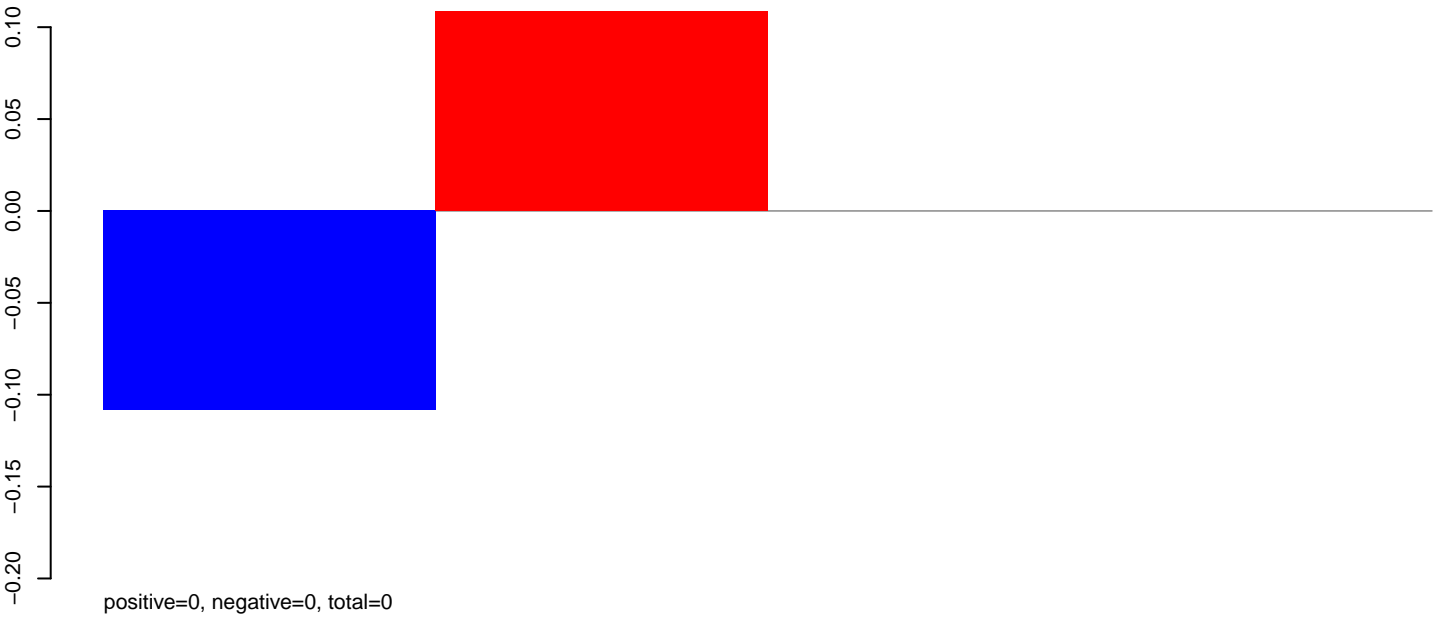
AnGam_Sua5b_cells.24_35.rep



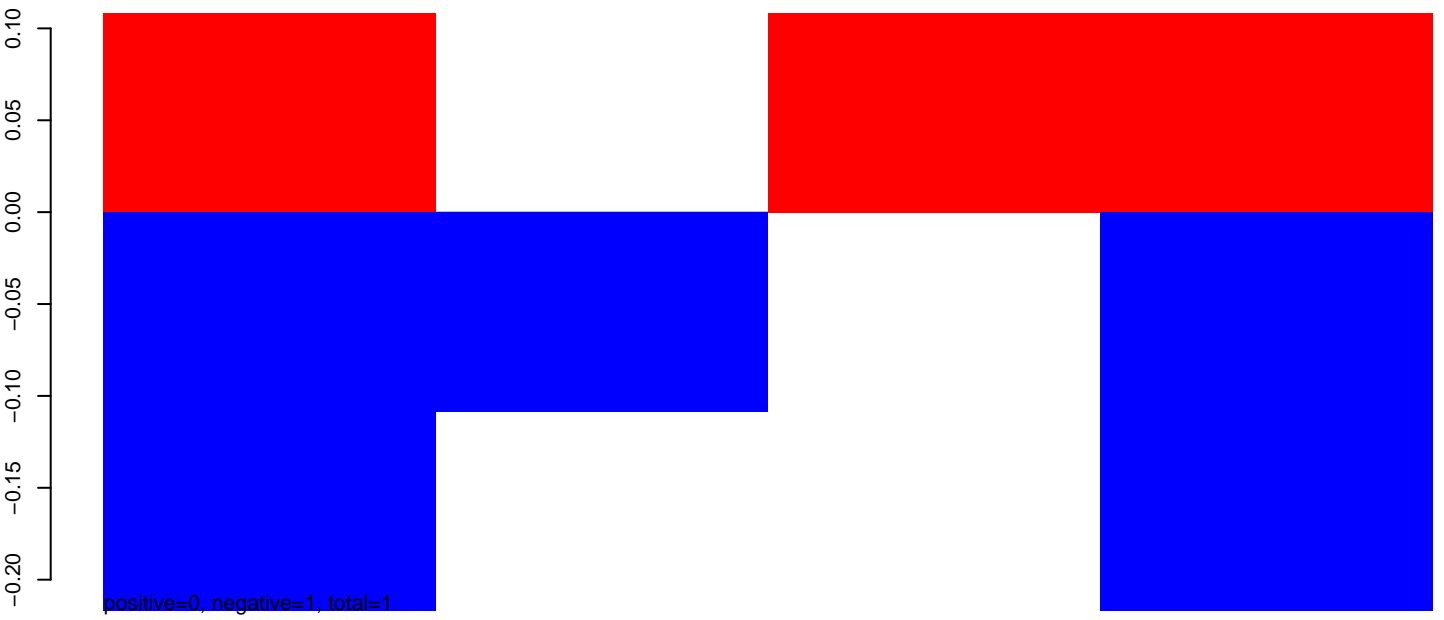
AnGam_Sua5b_cells.rep



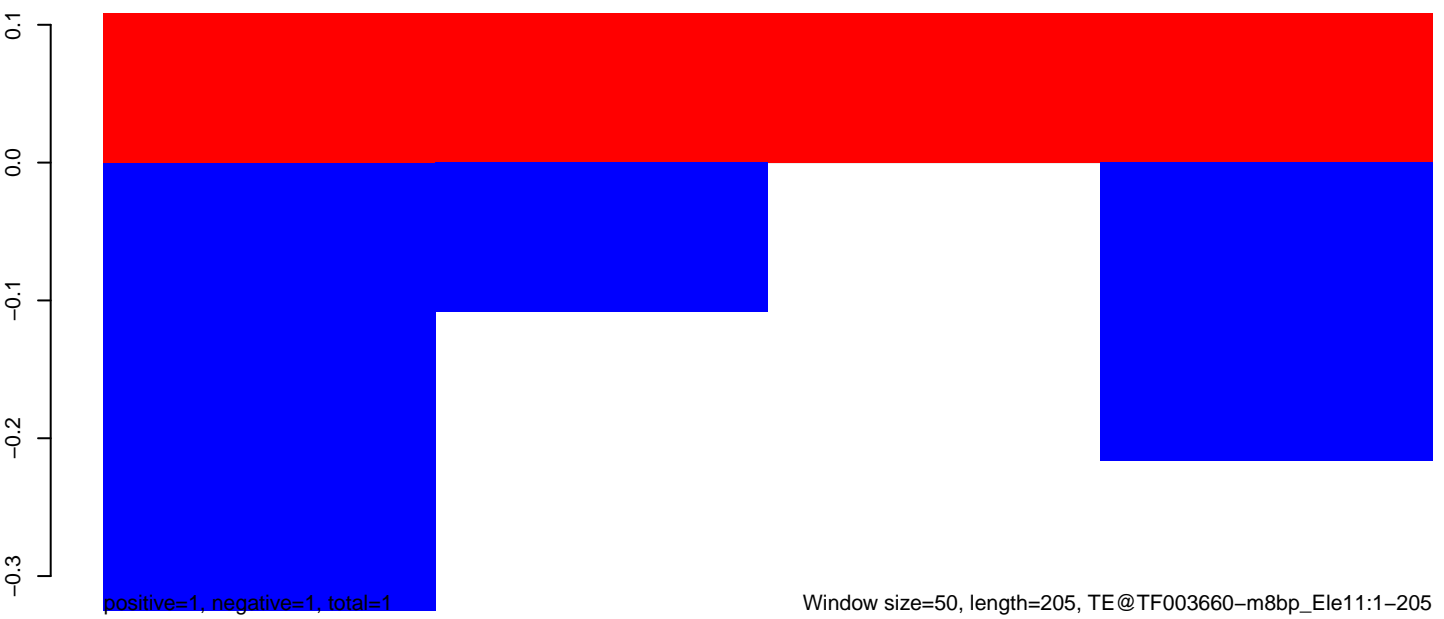
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AnGam_Sua5b_cells.24_35.rep

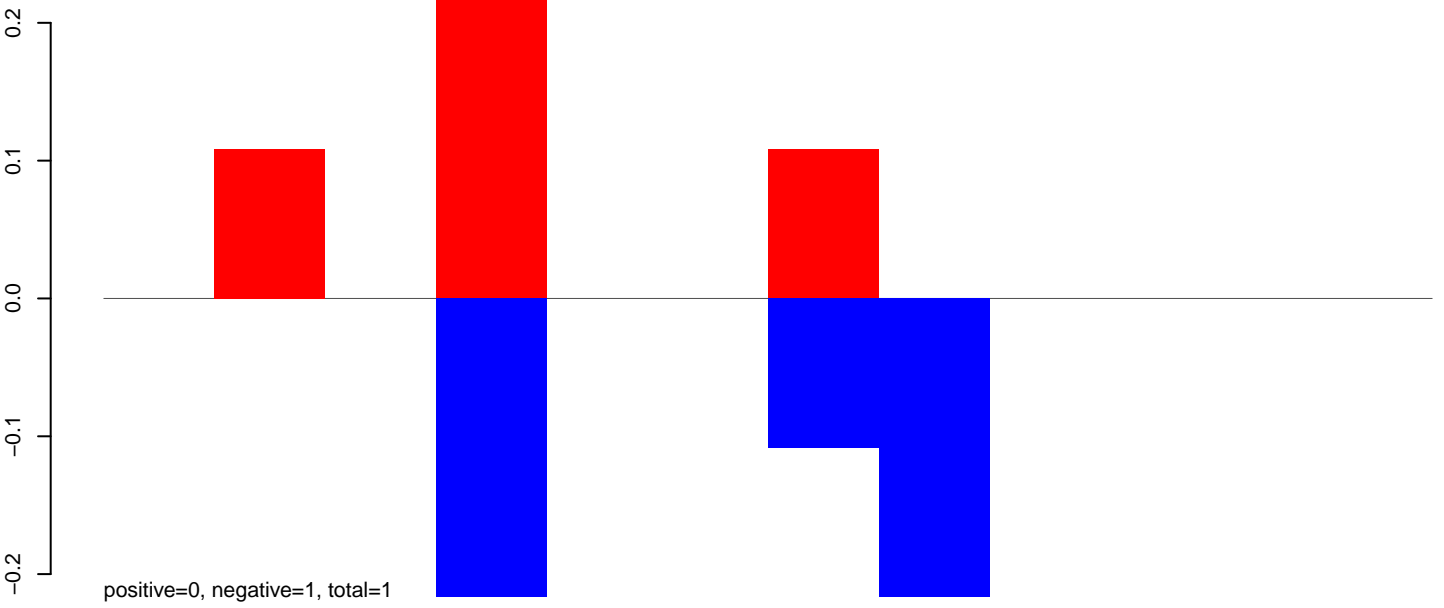


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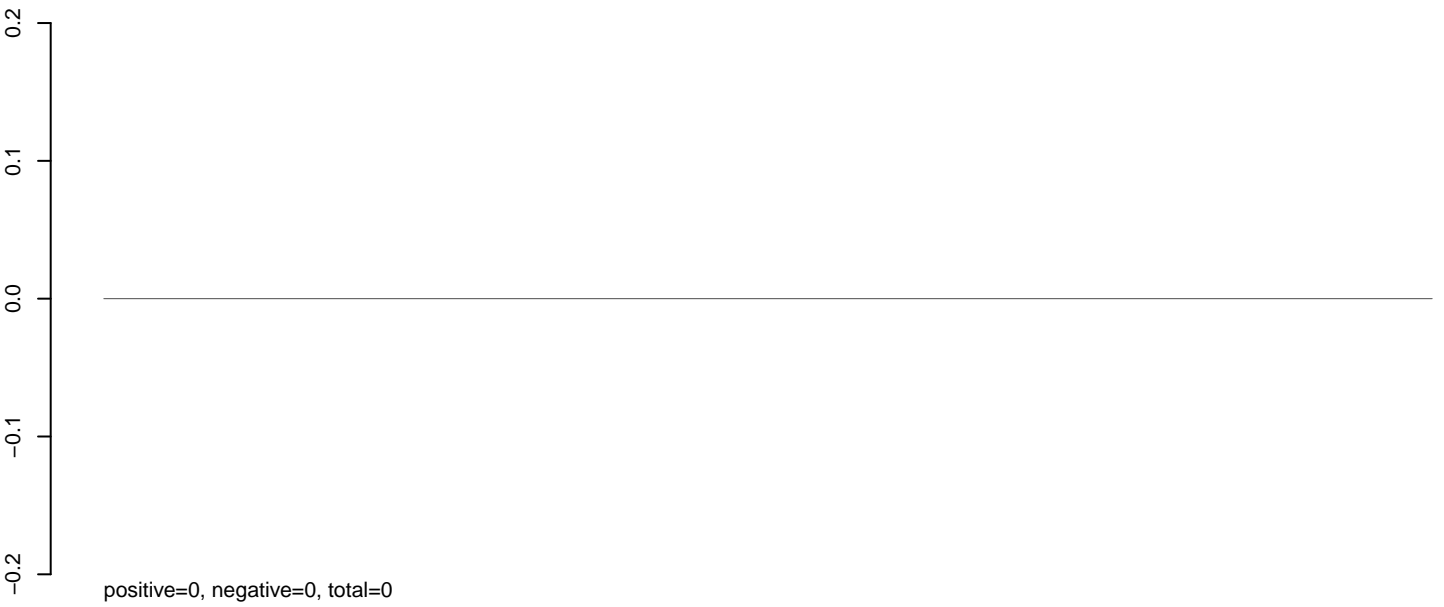


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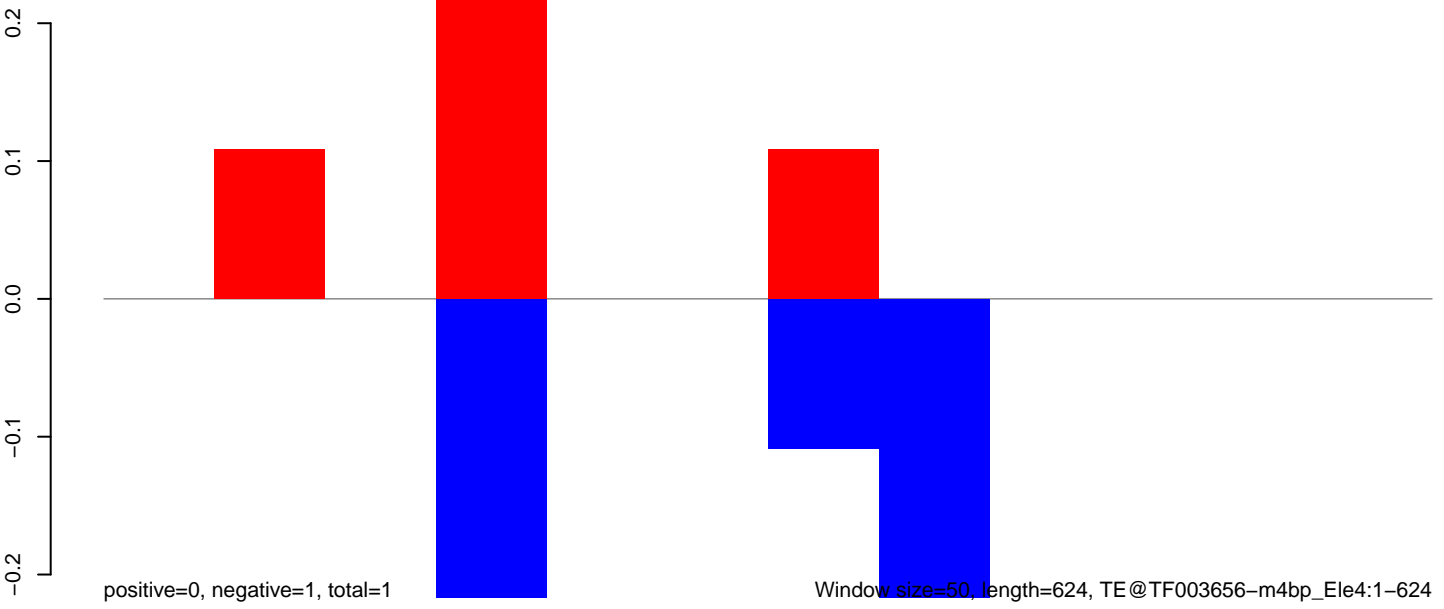
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AnGam_Sua5b_cells.24_35.rep

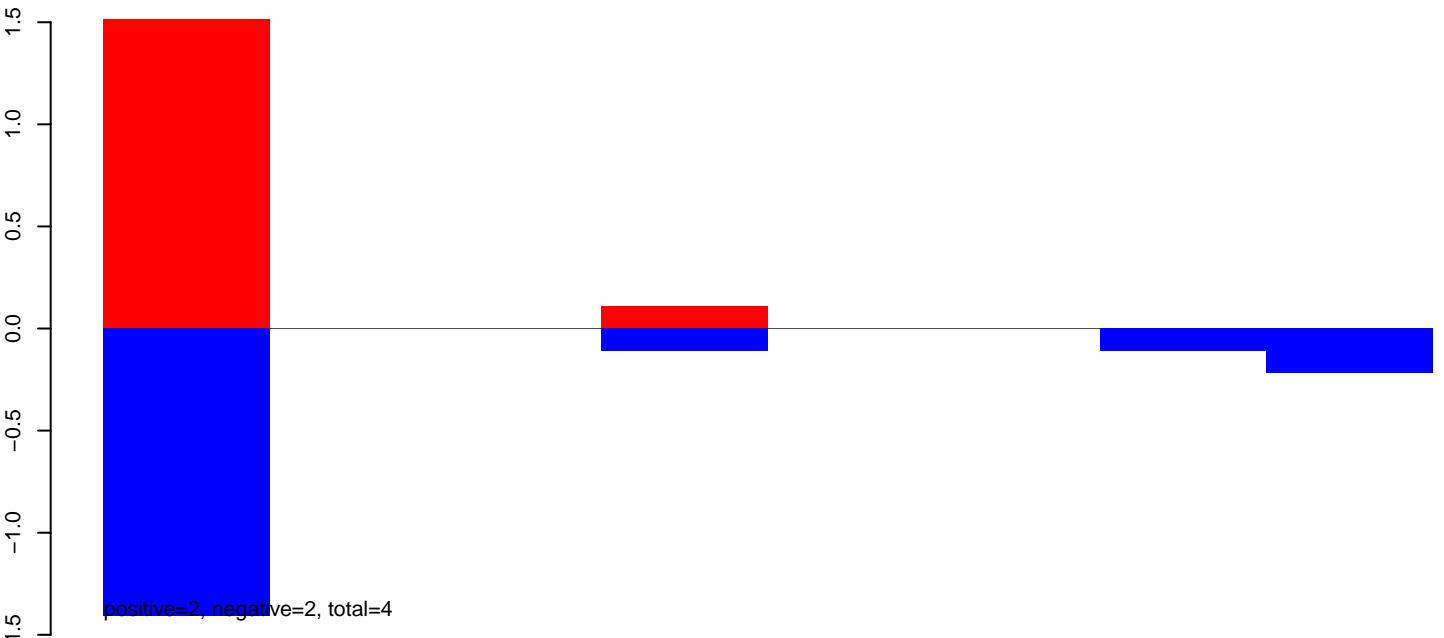


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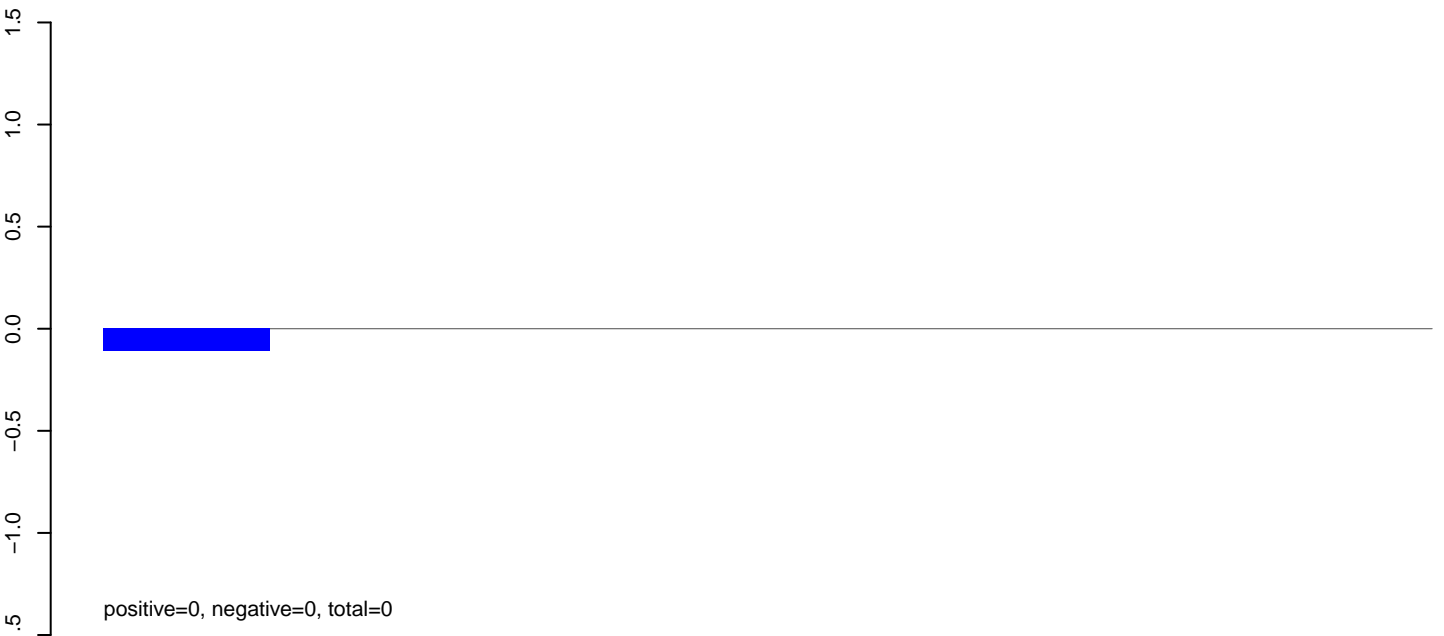


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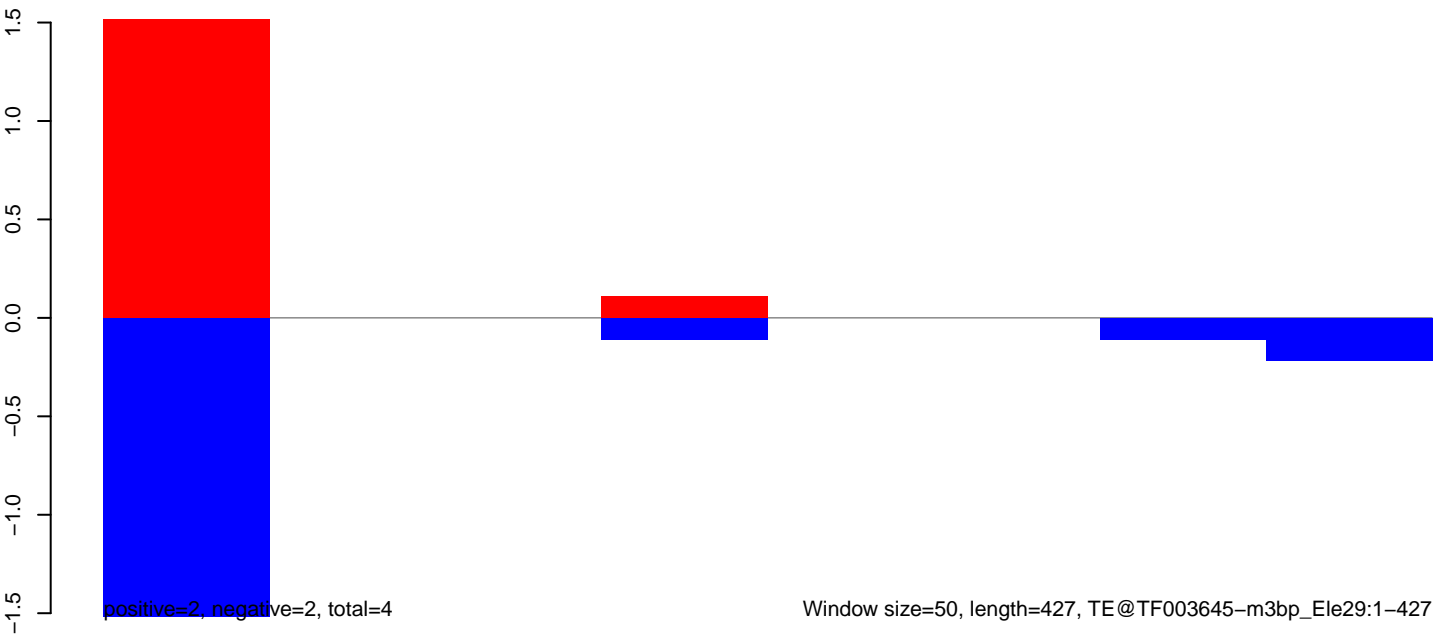
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AnGam_Sua5b_cells.24_35.rep



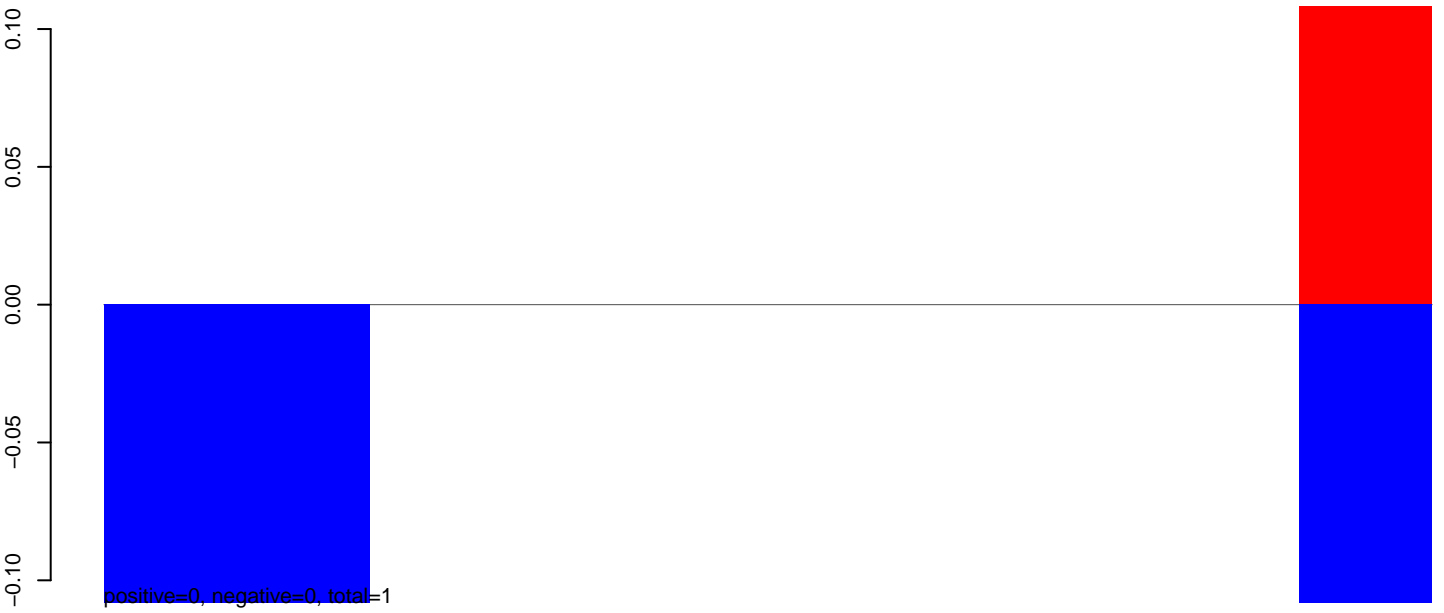
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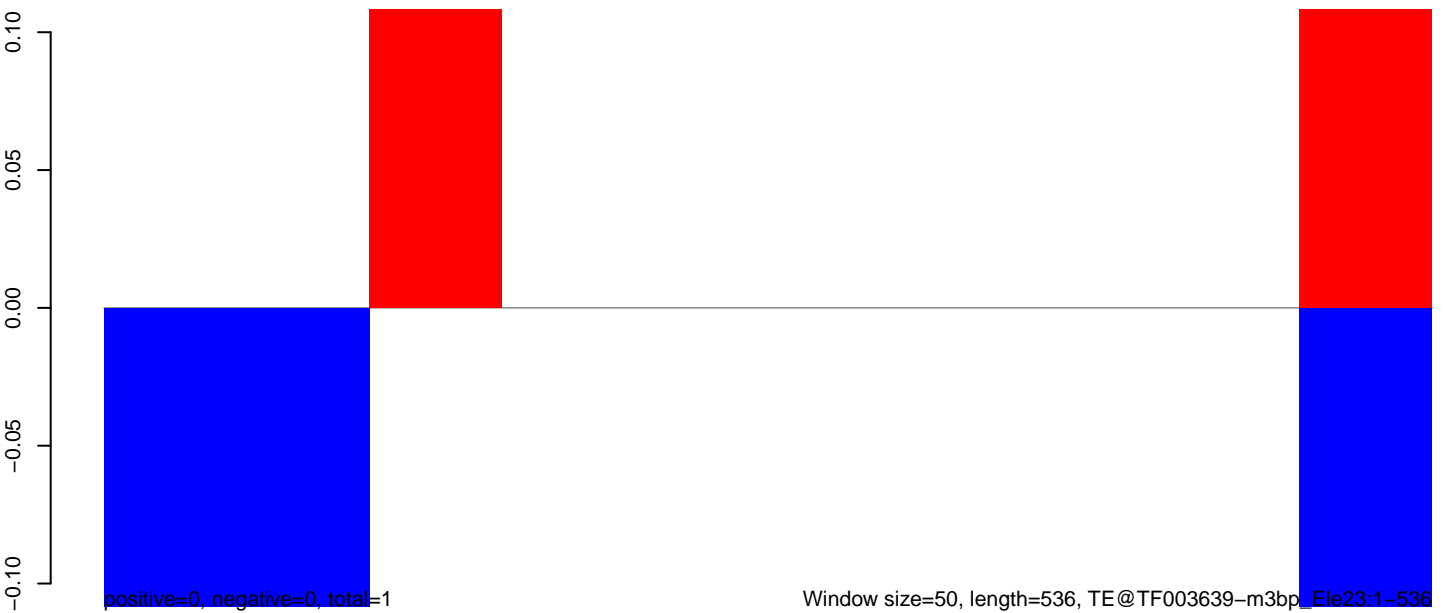
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AnGam_Sua5b_cells.24_35.rep

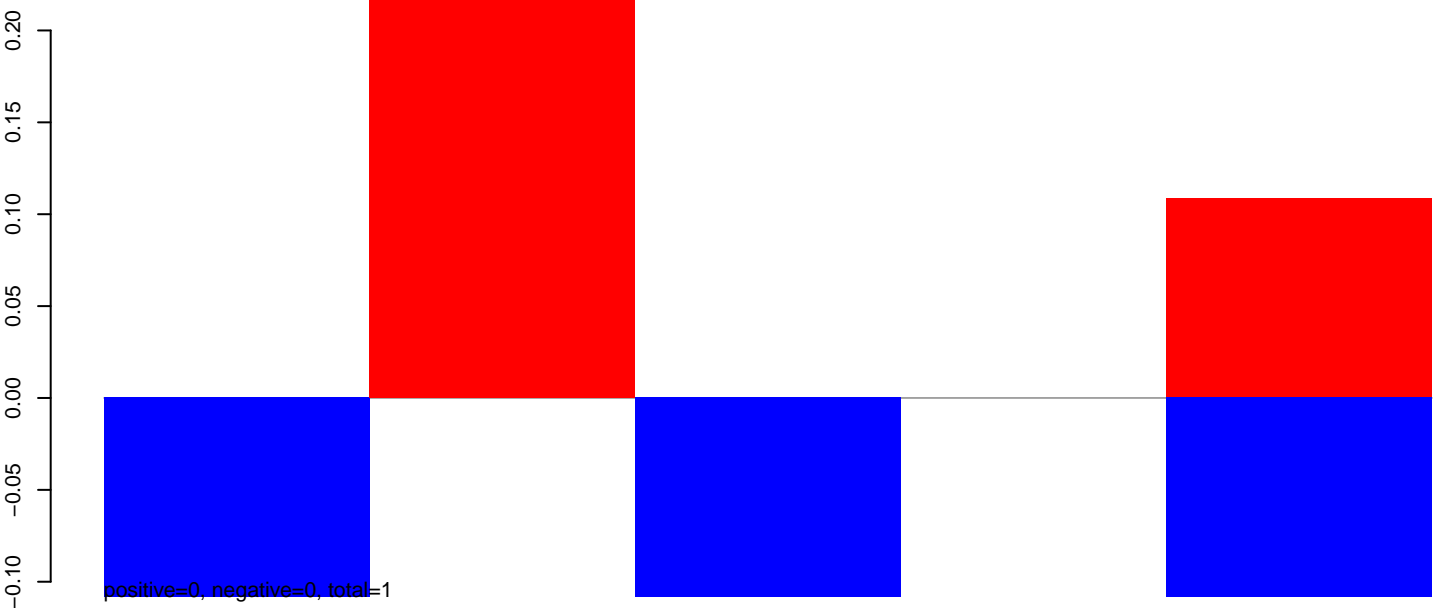


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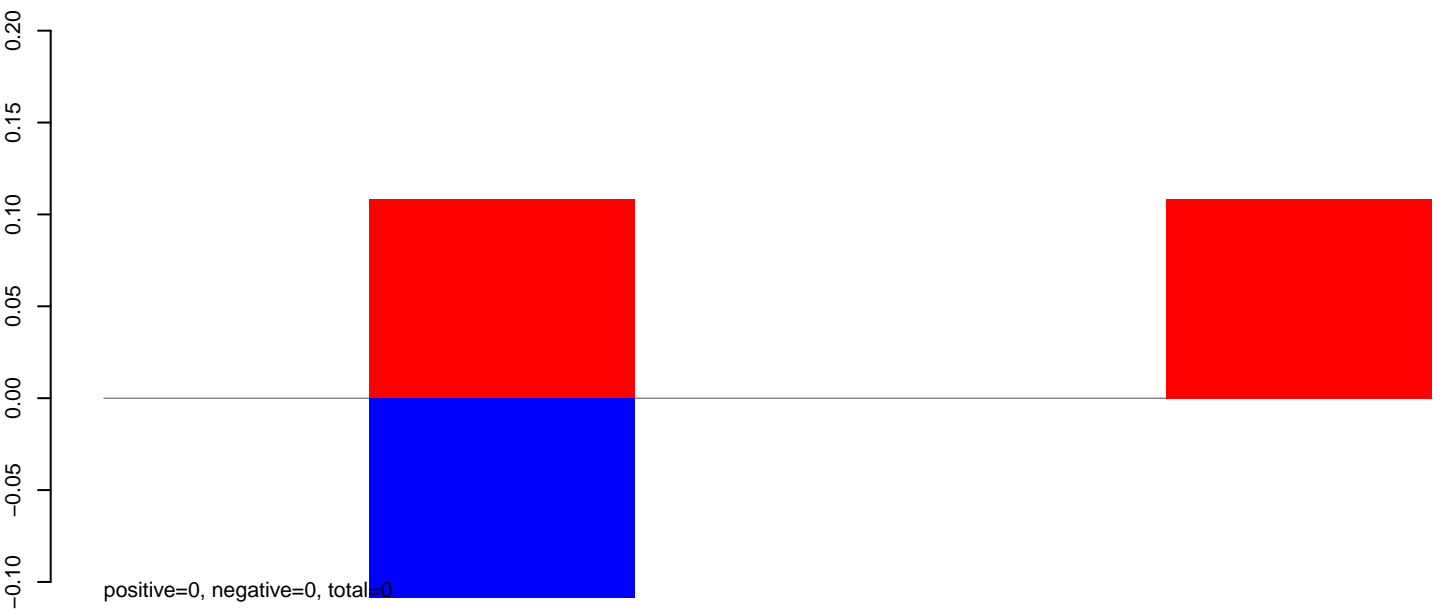


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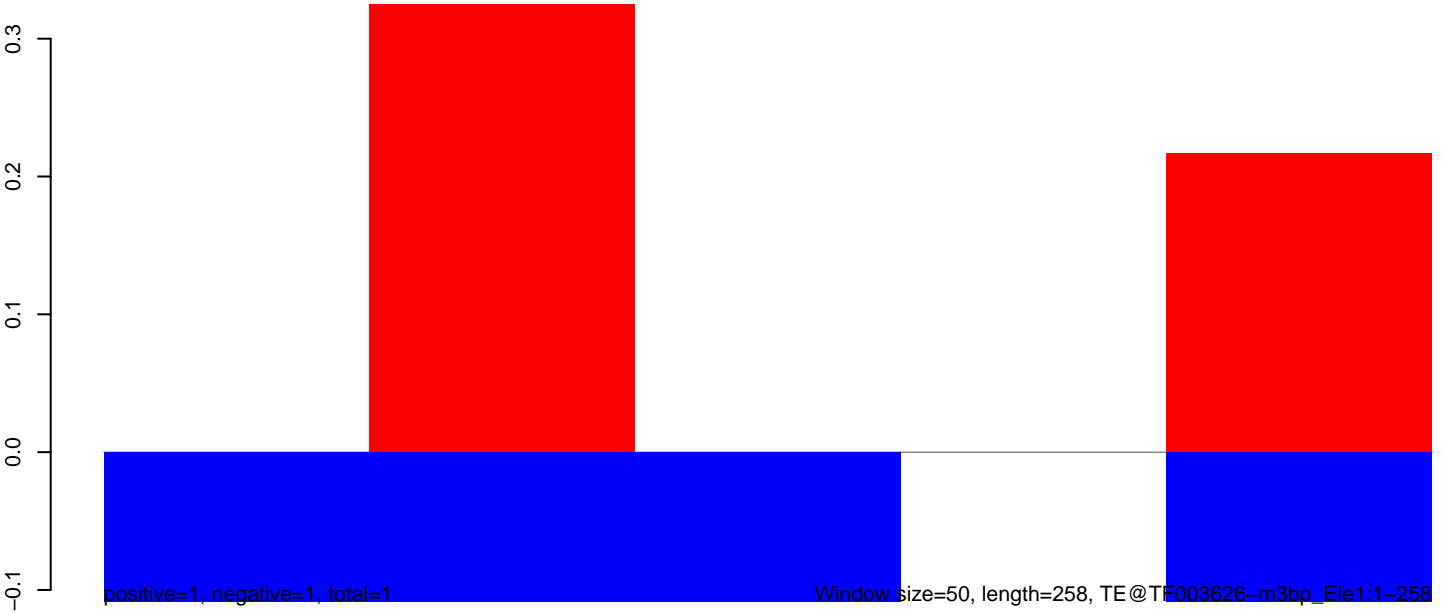
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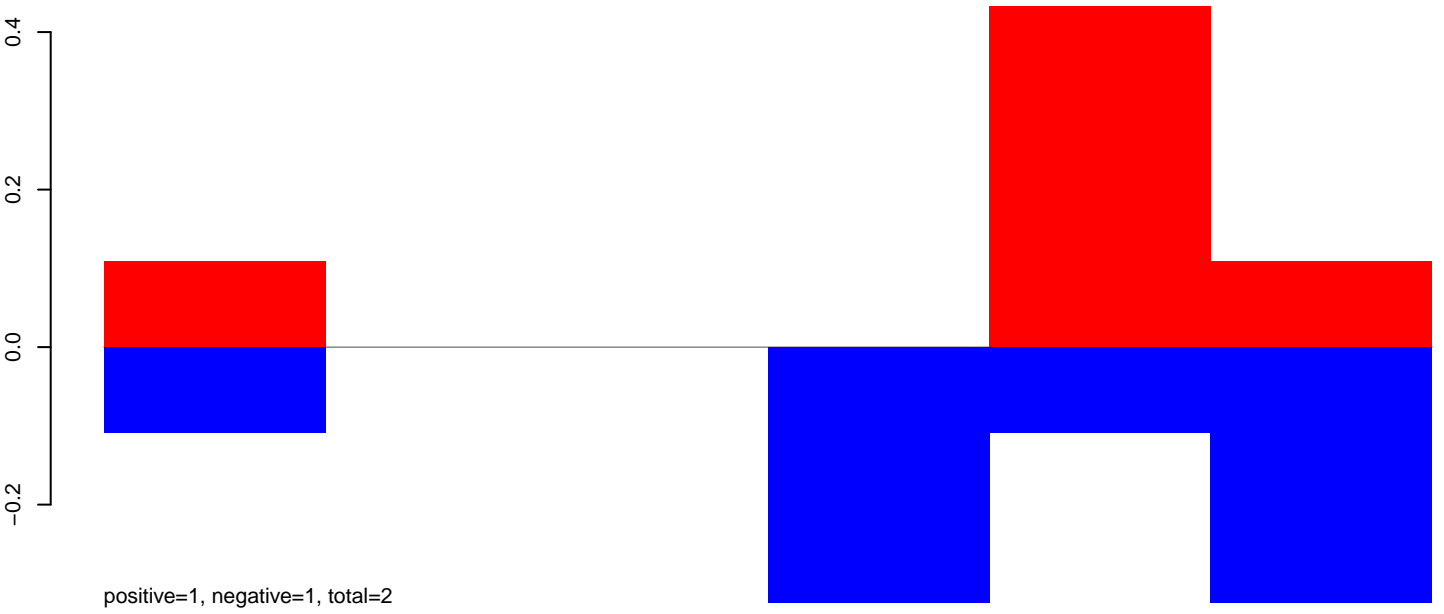
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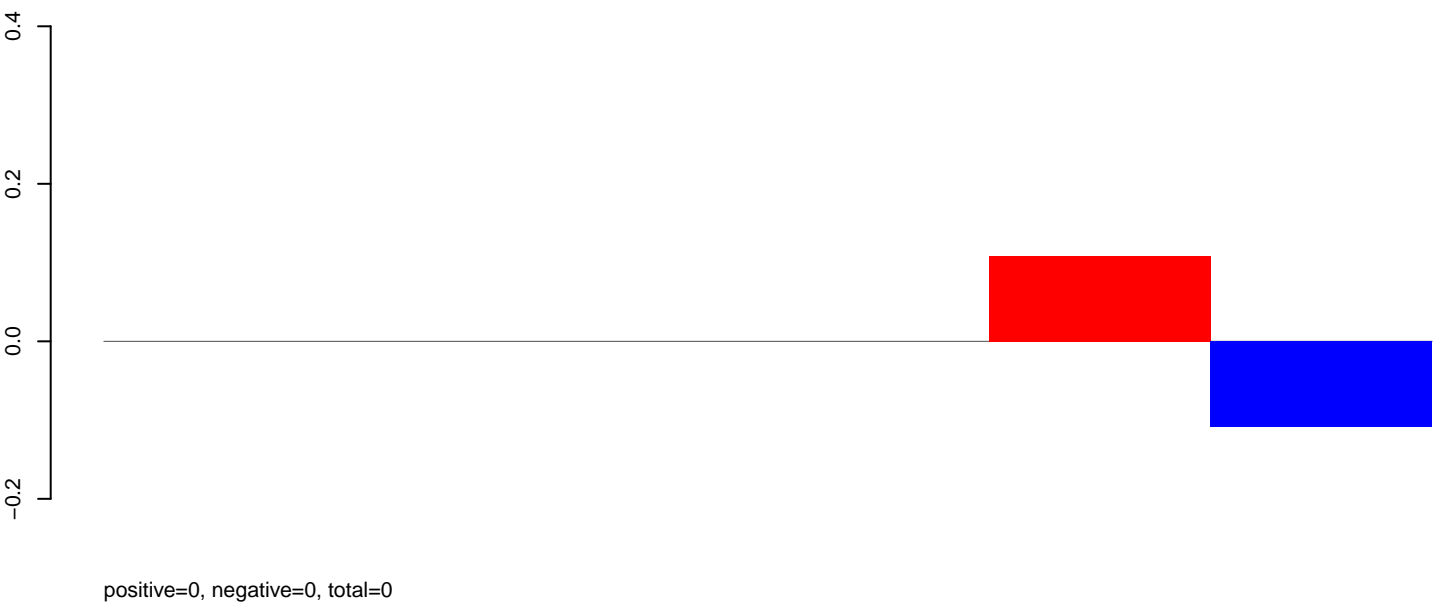
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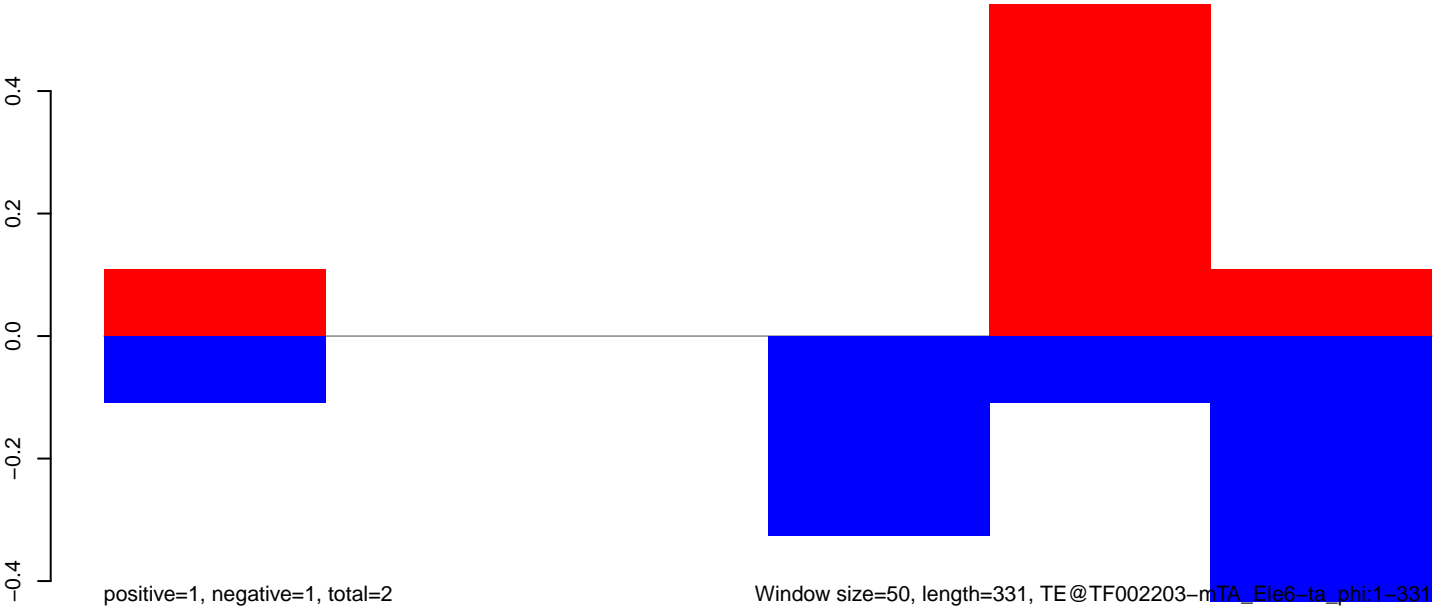
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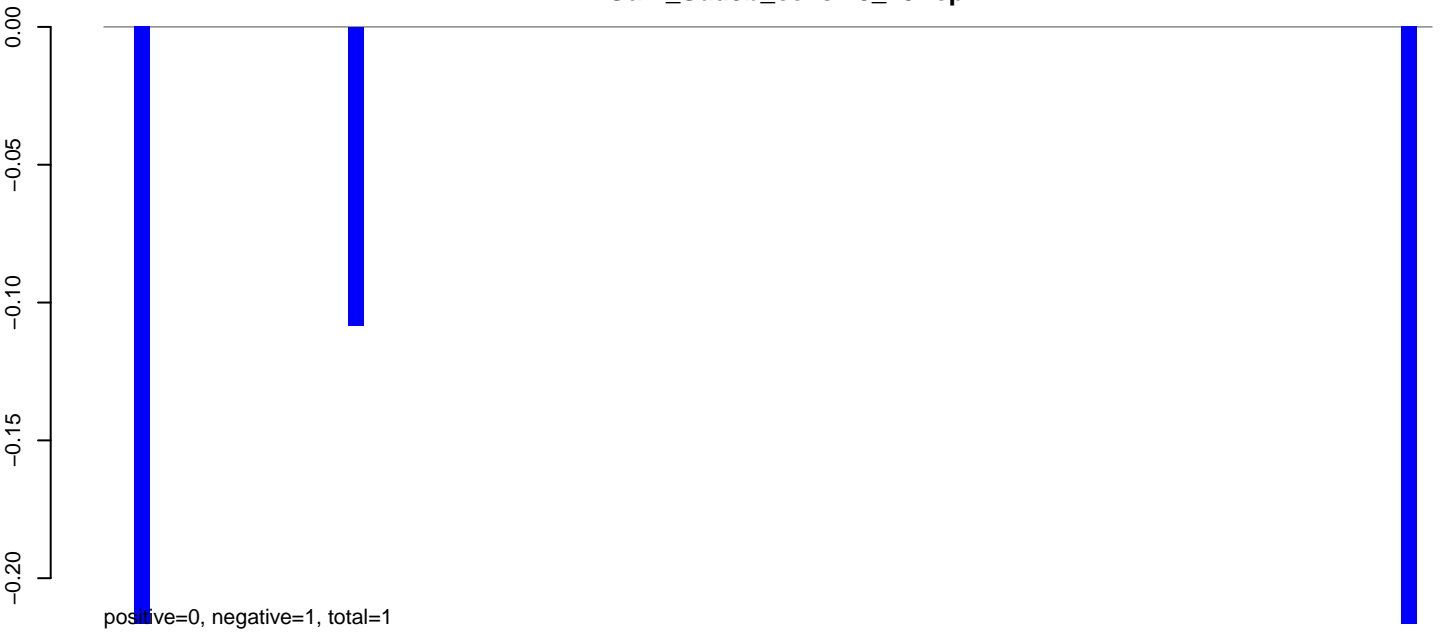
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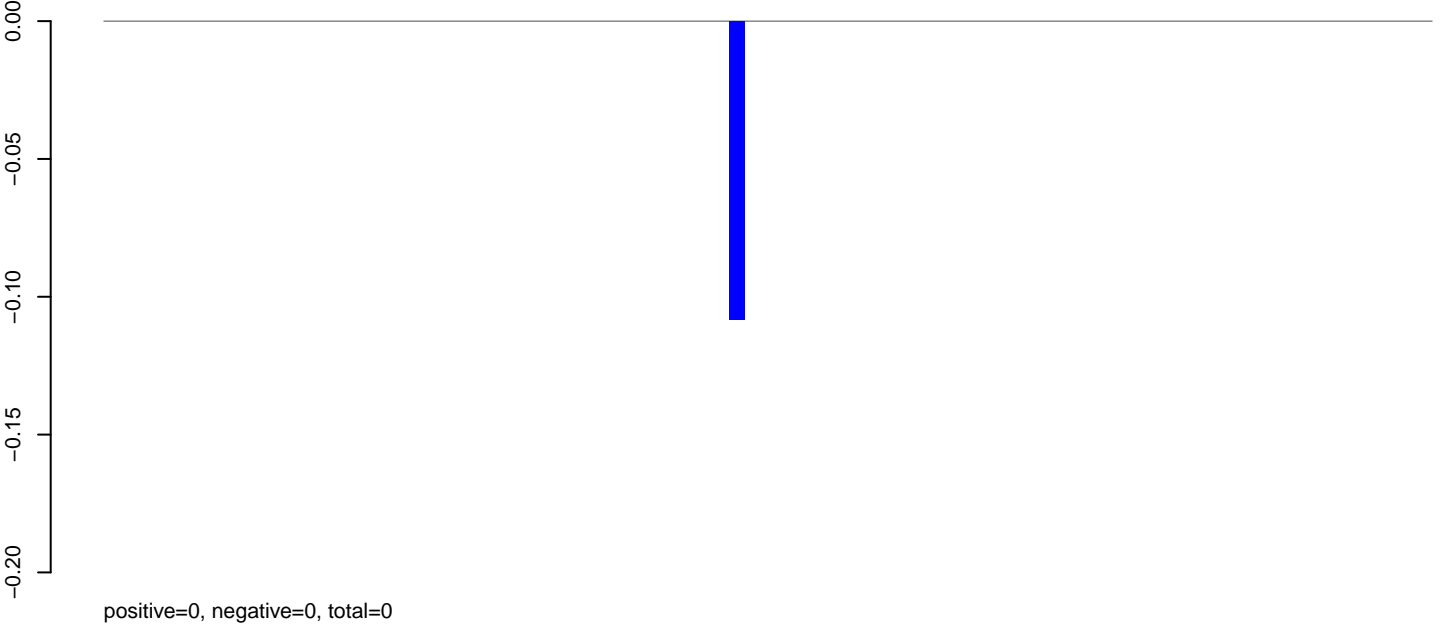
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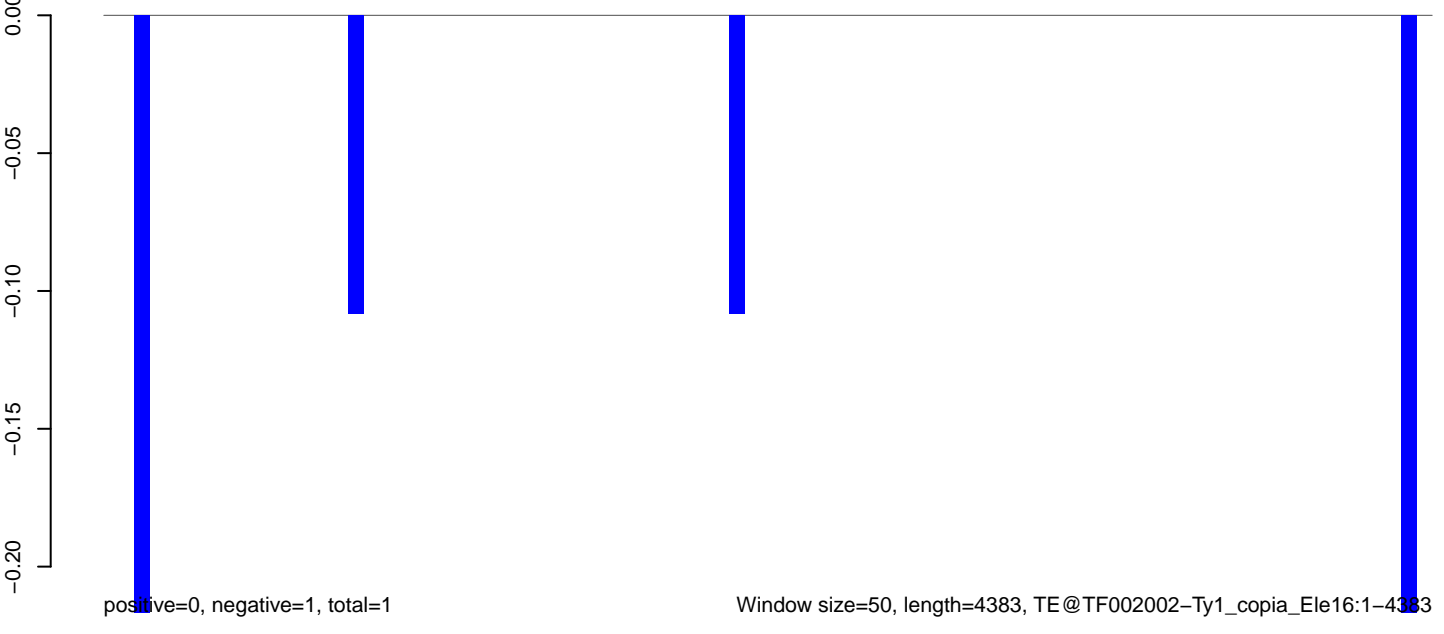
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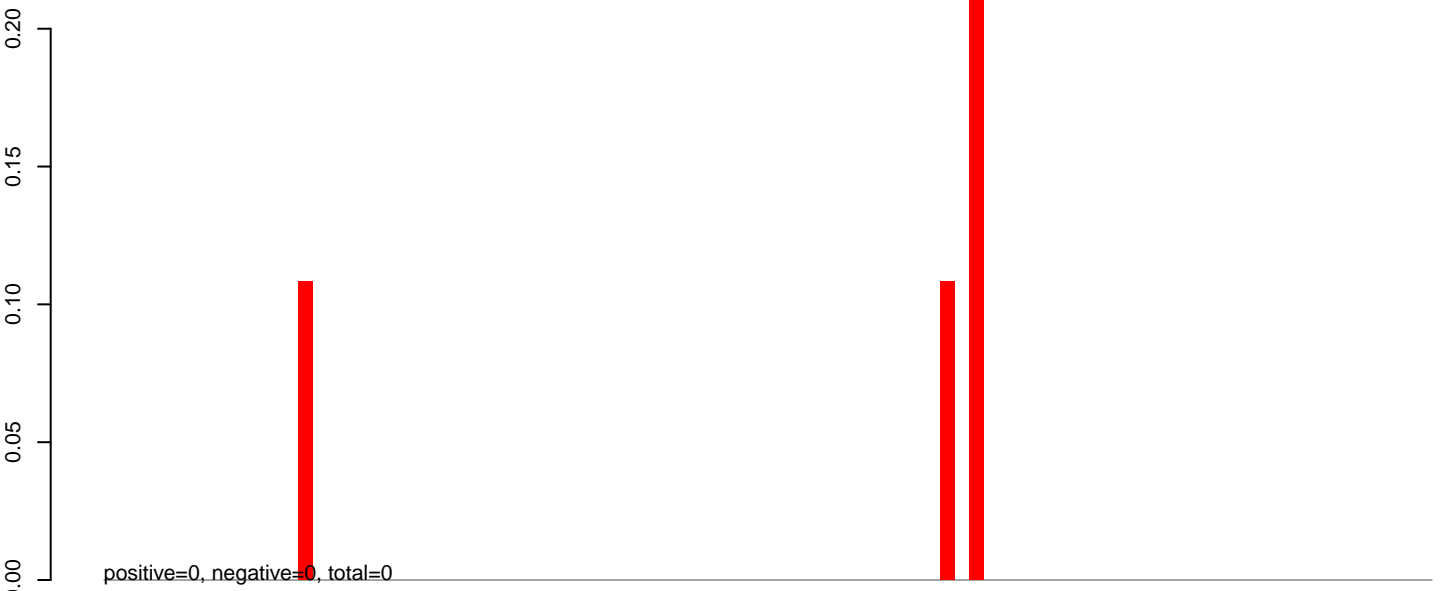
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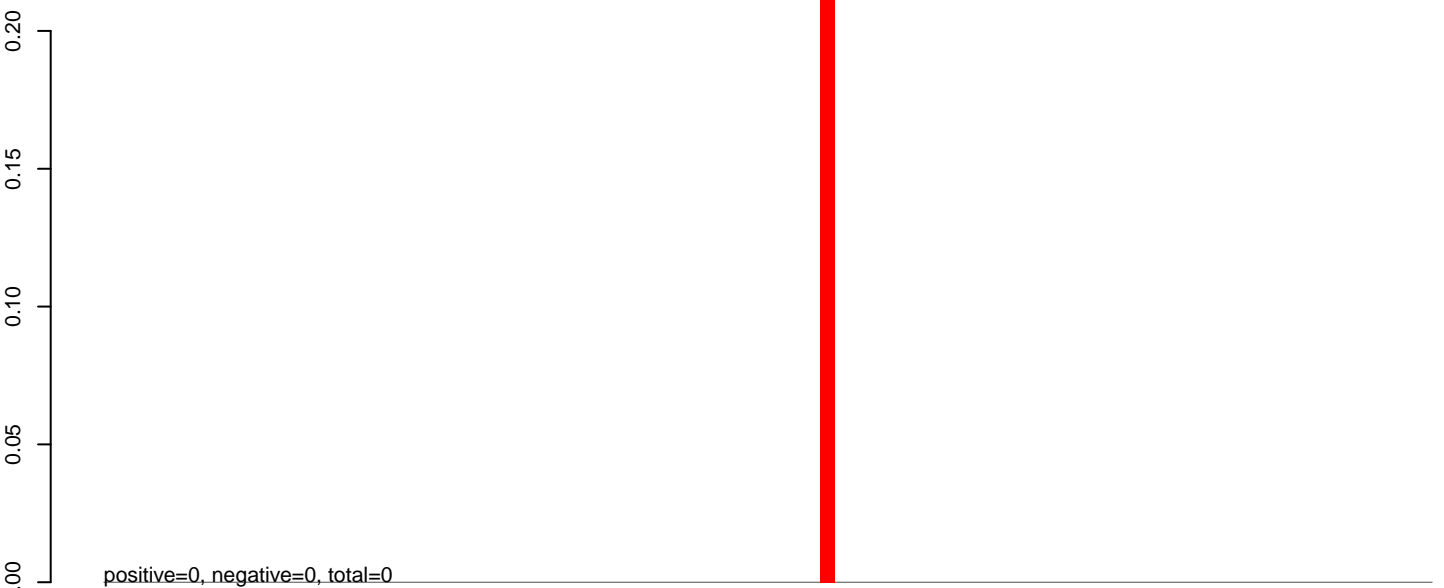
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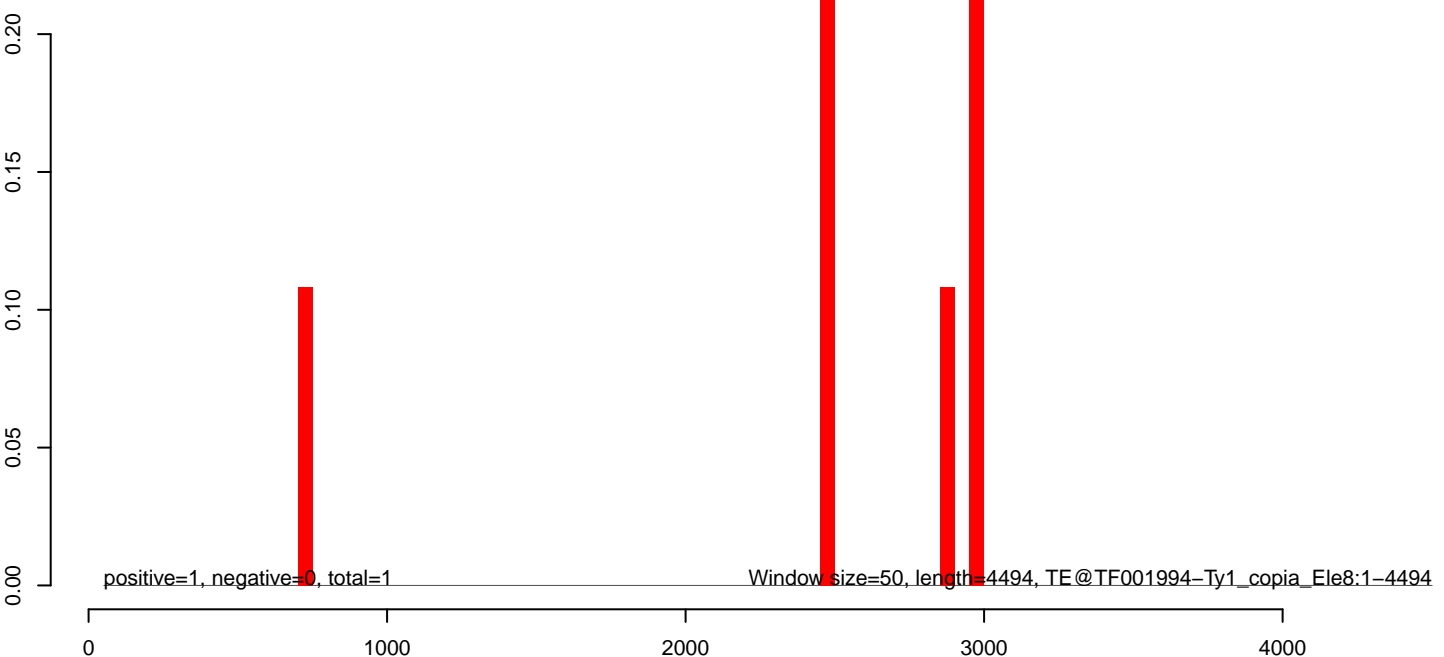
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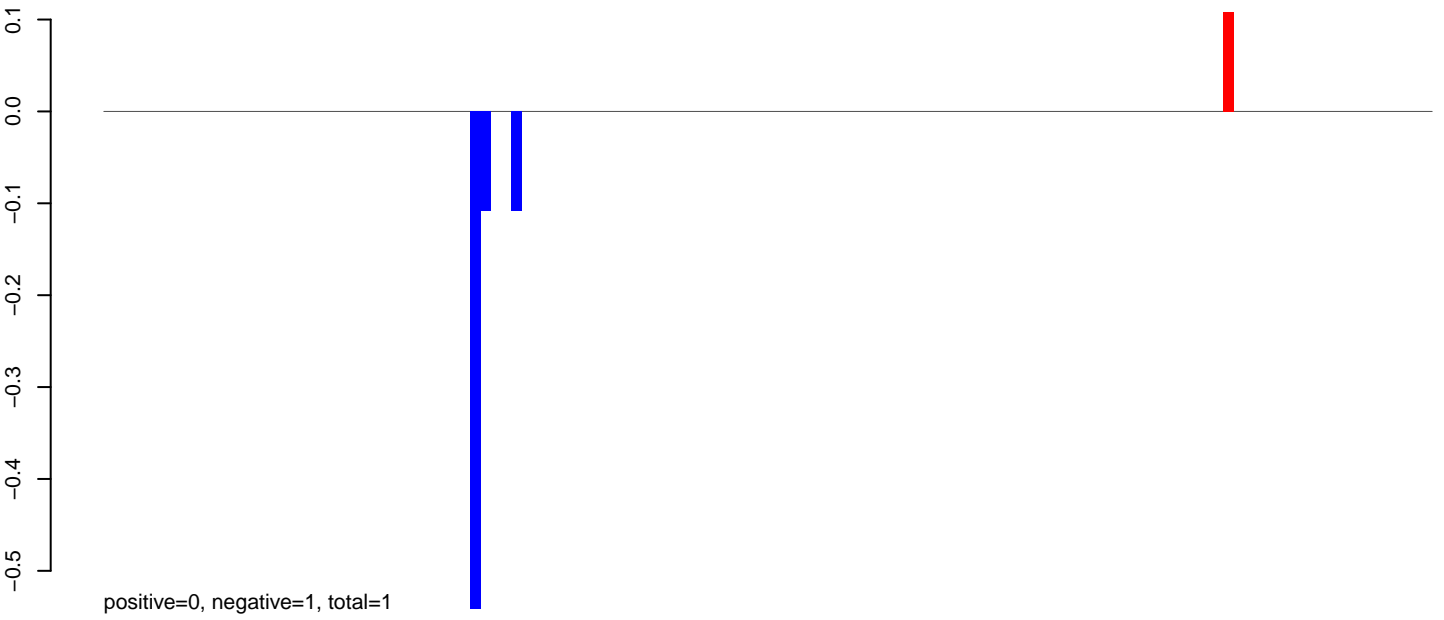
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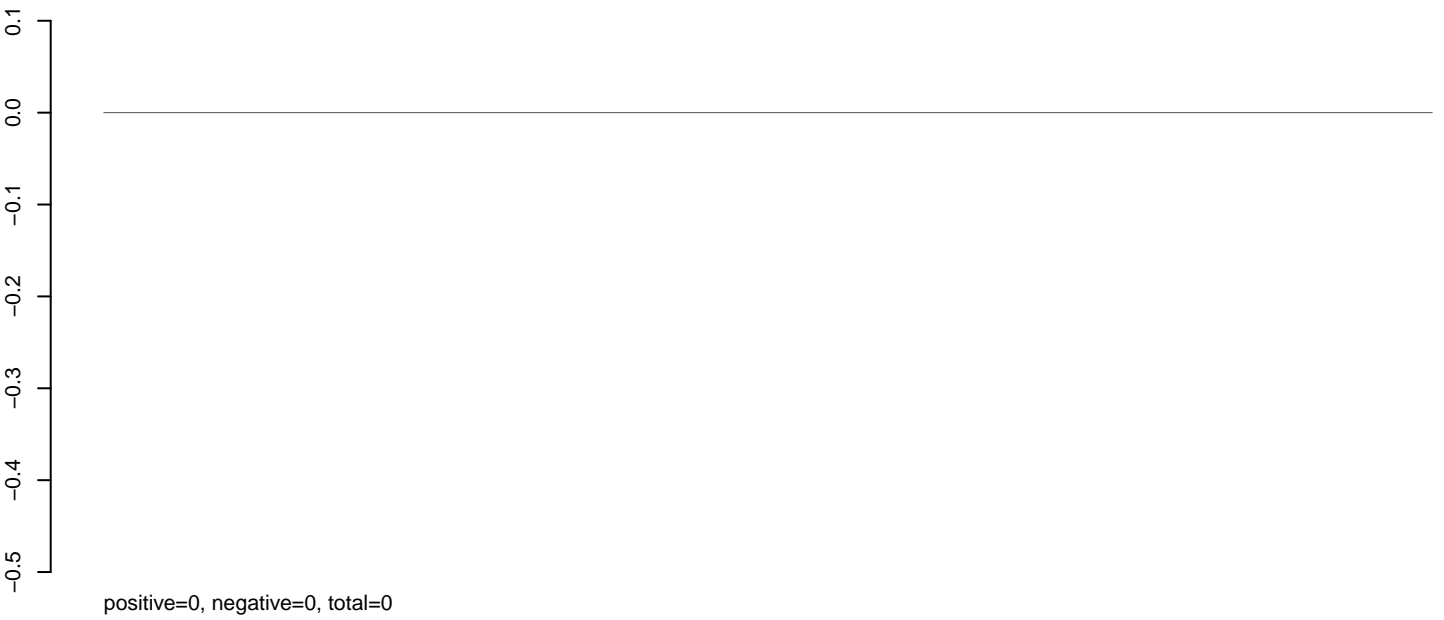
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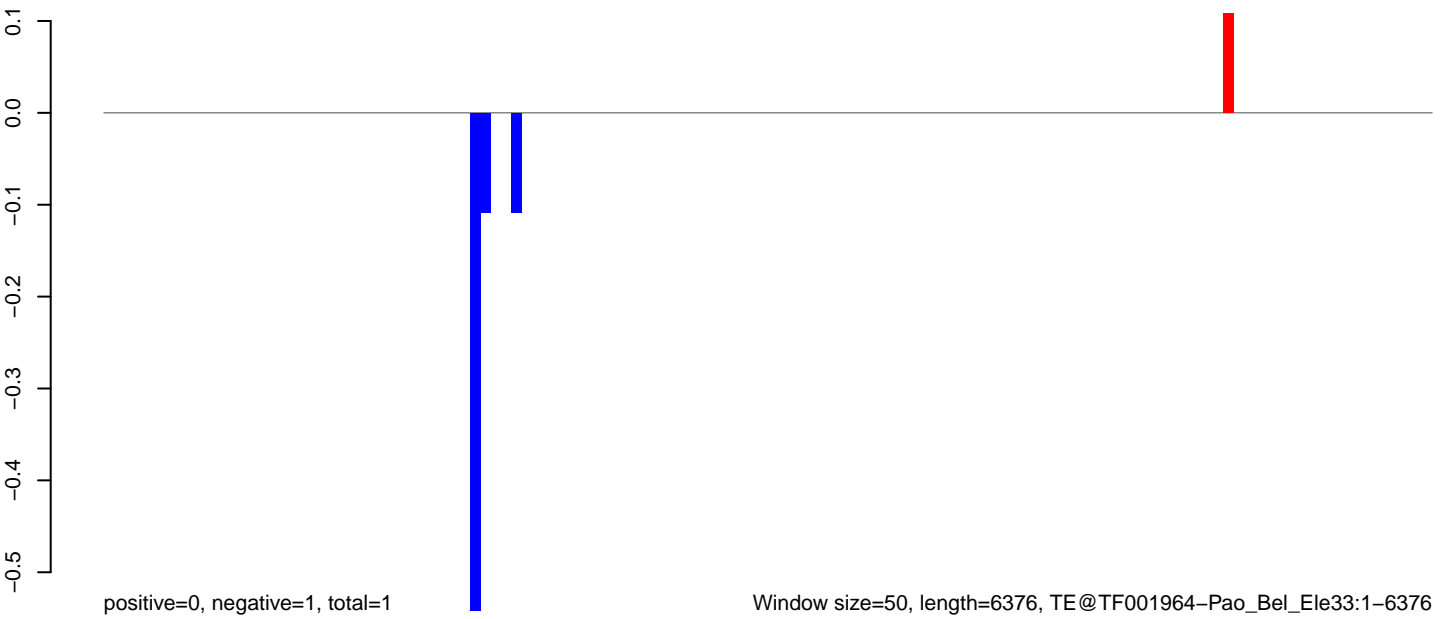
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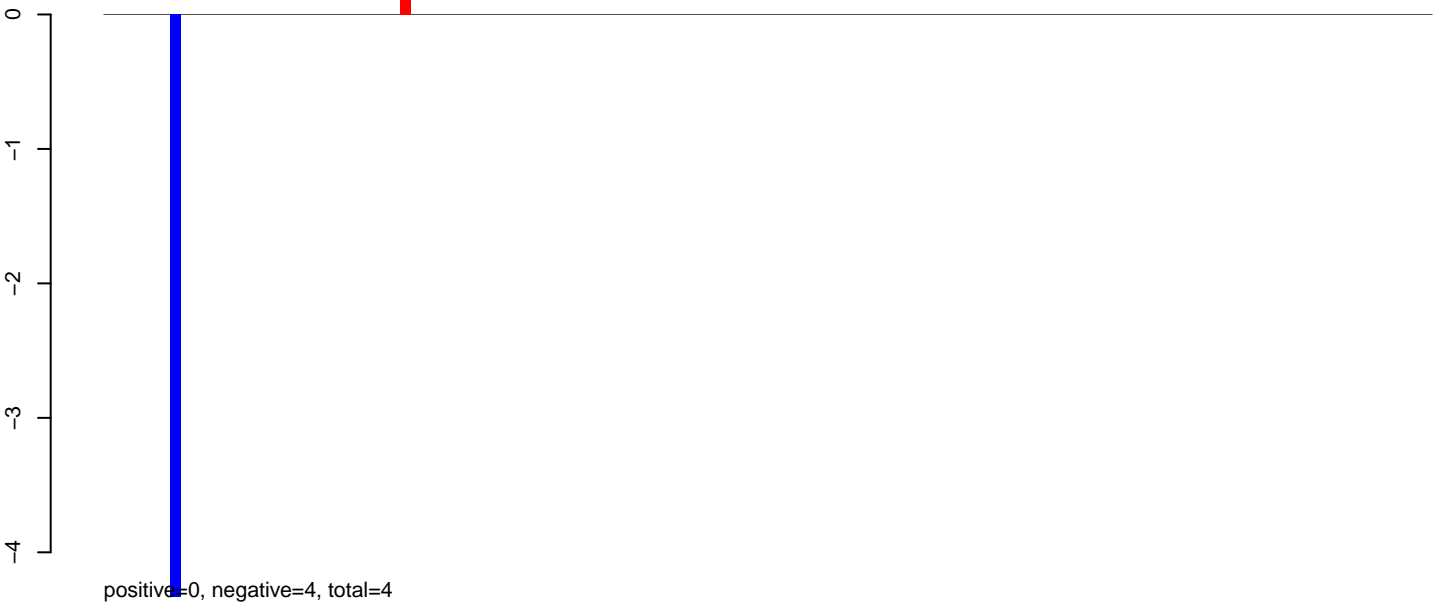
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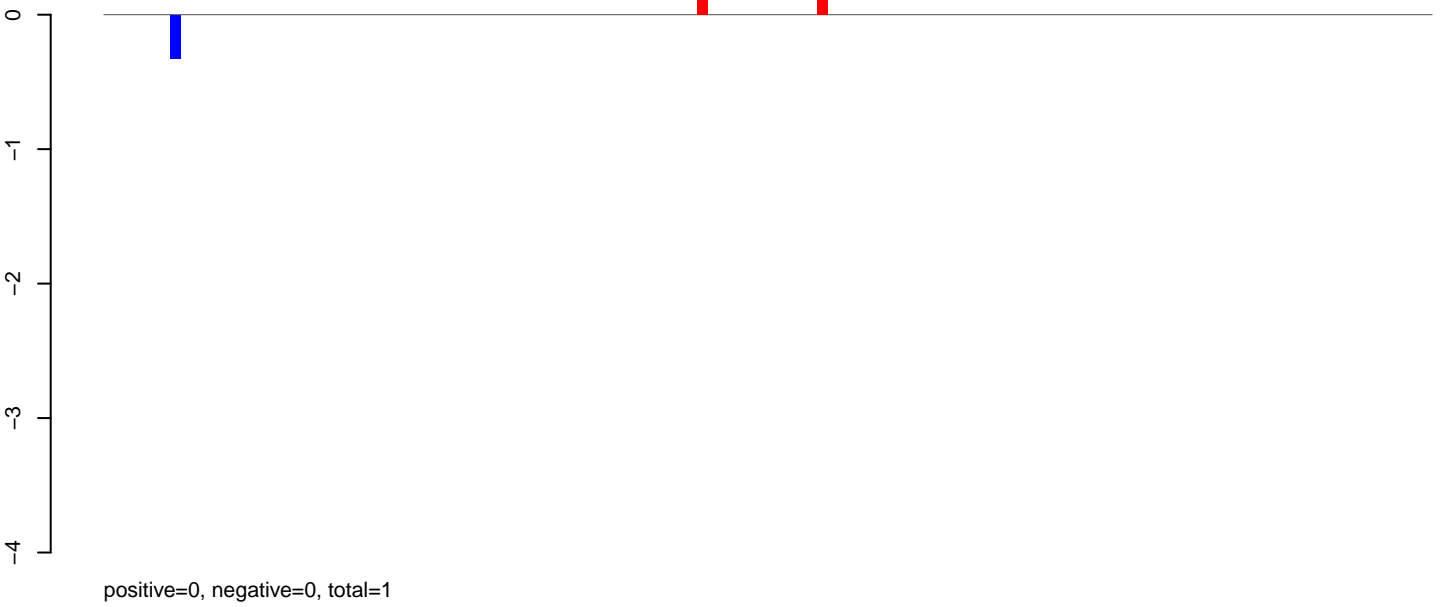
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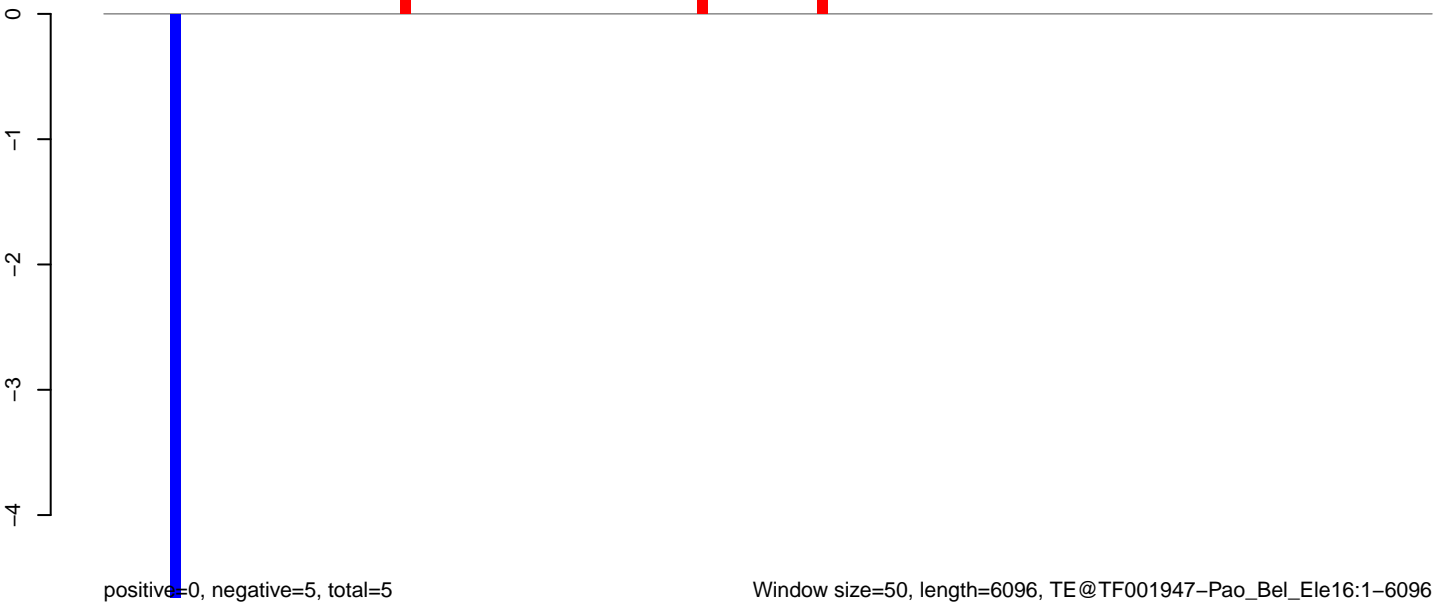
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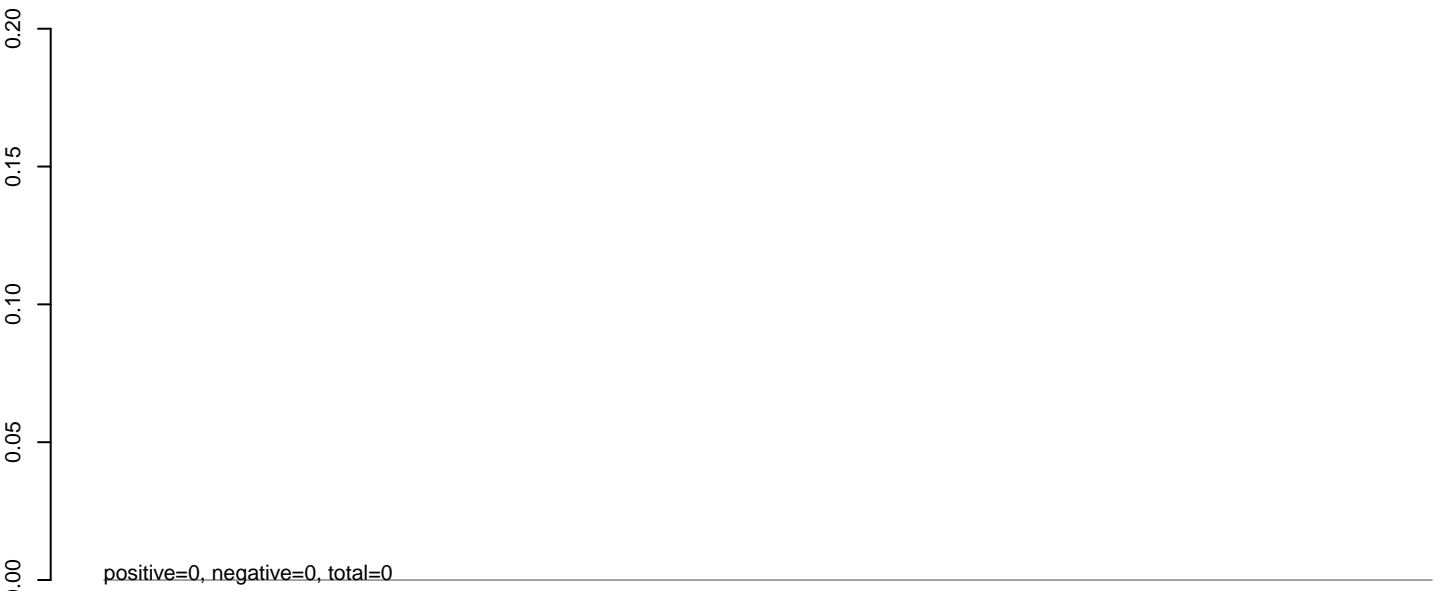
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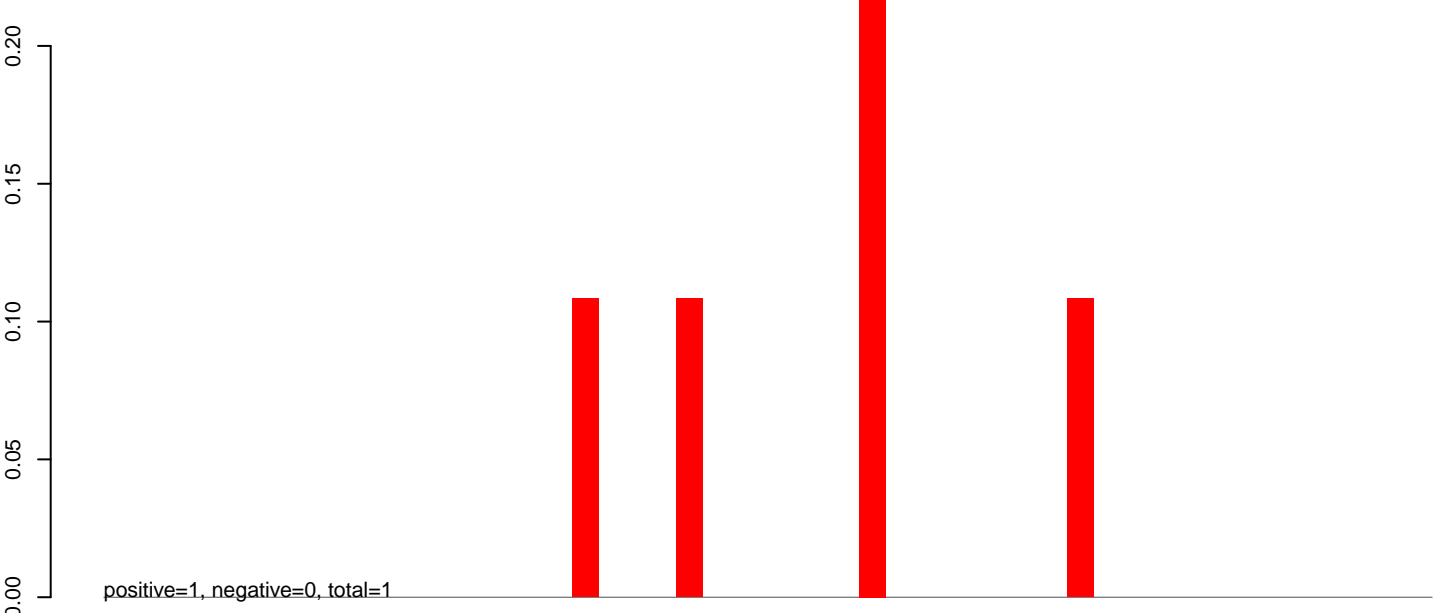
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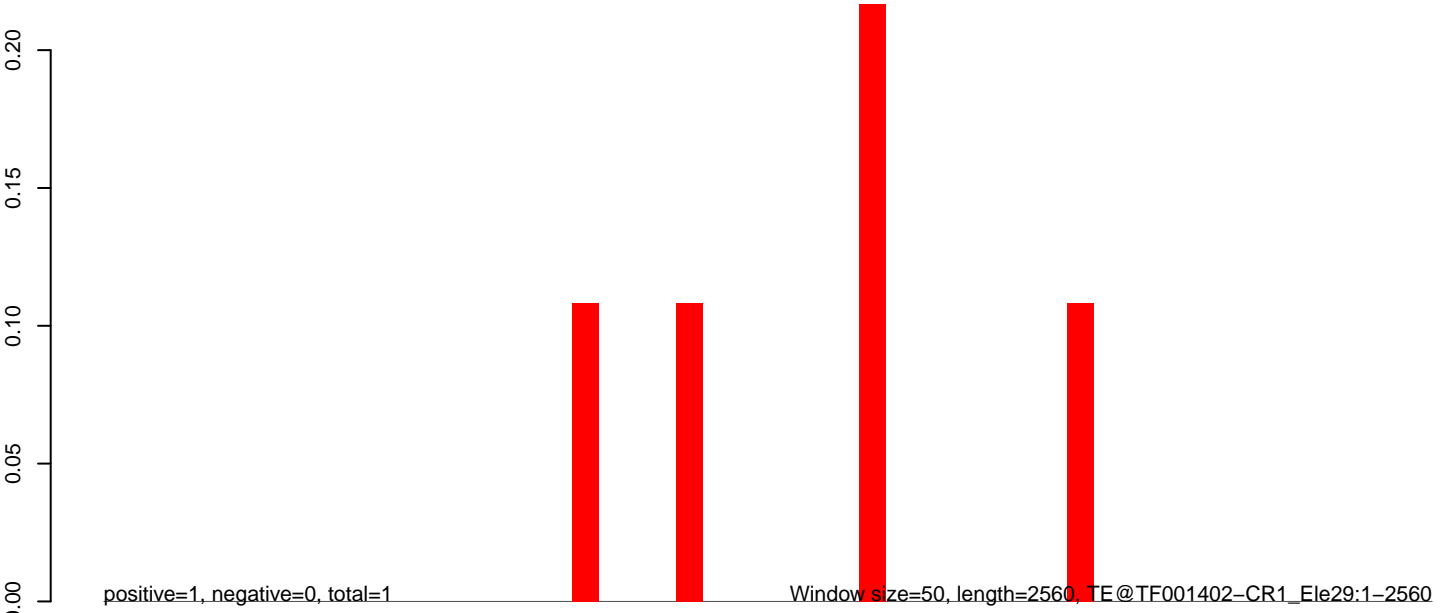
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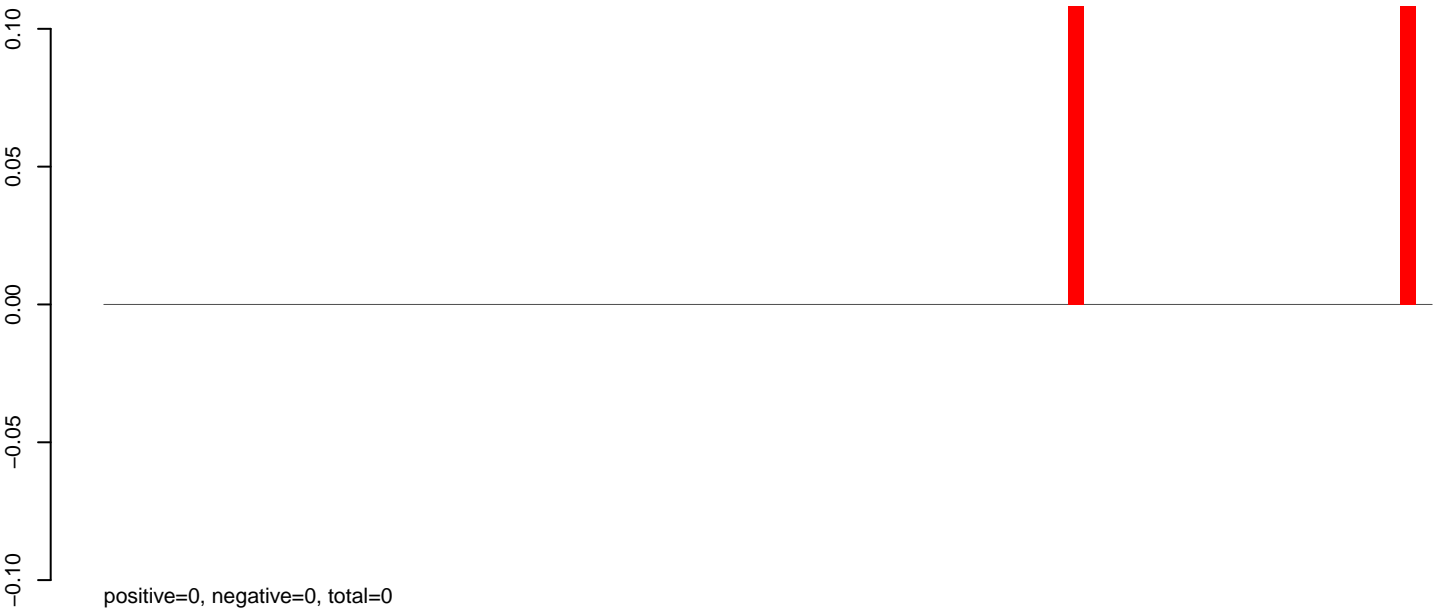


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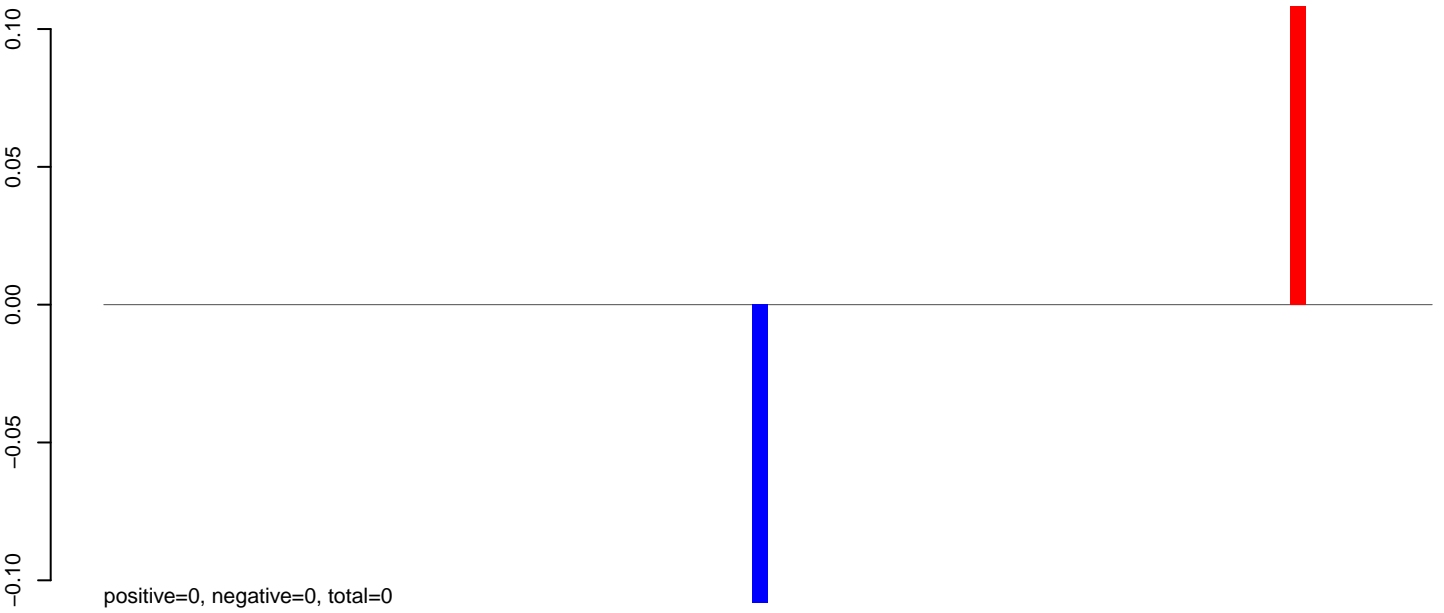


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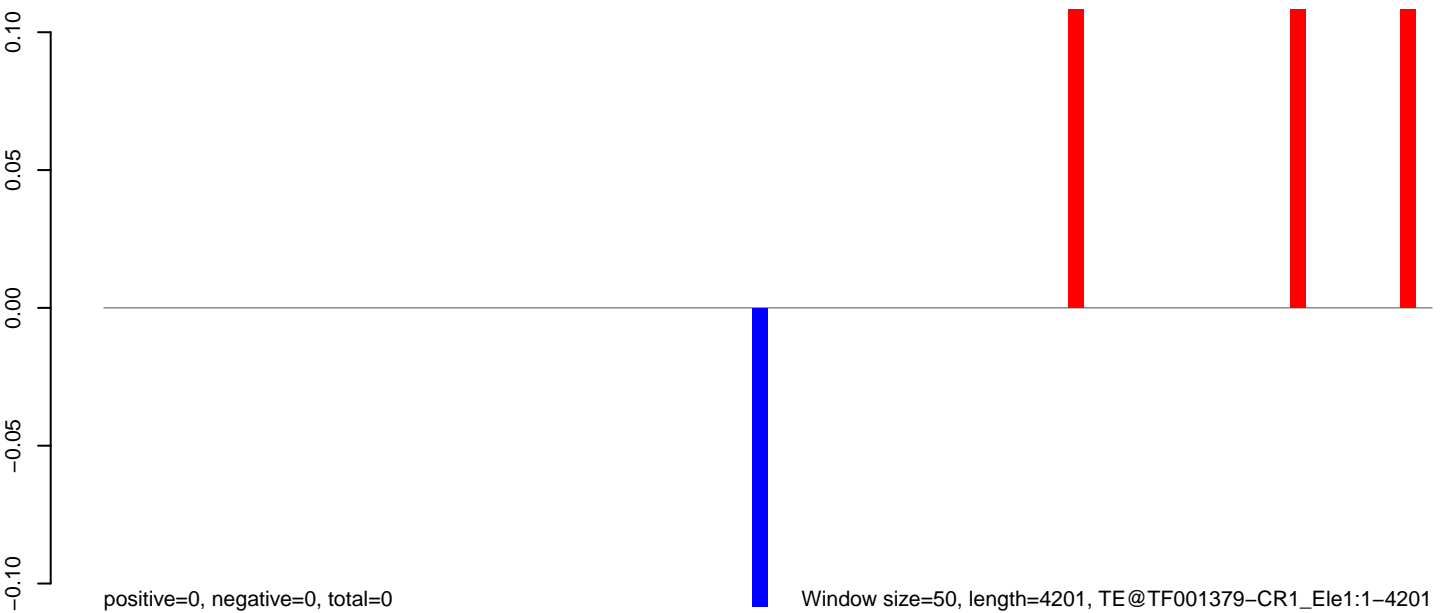
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AnGam_Sua5b_cells.24_35.rep

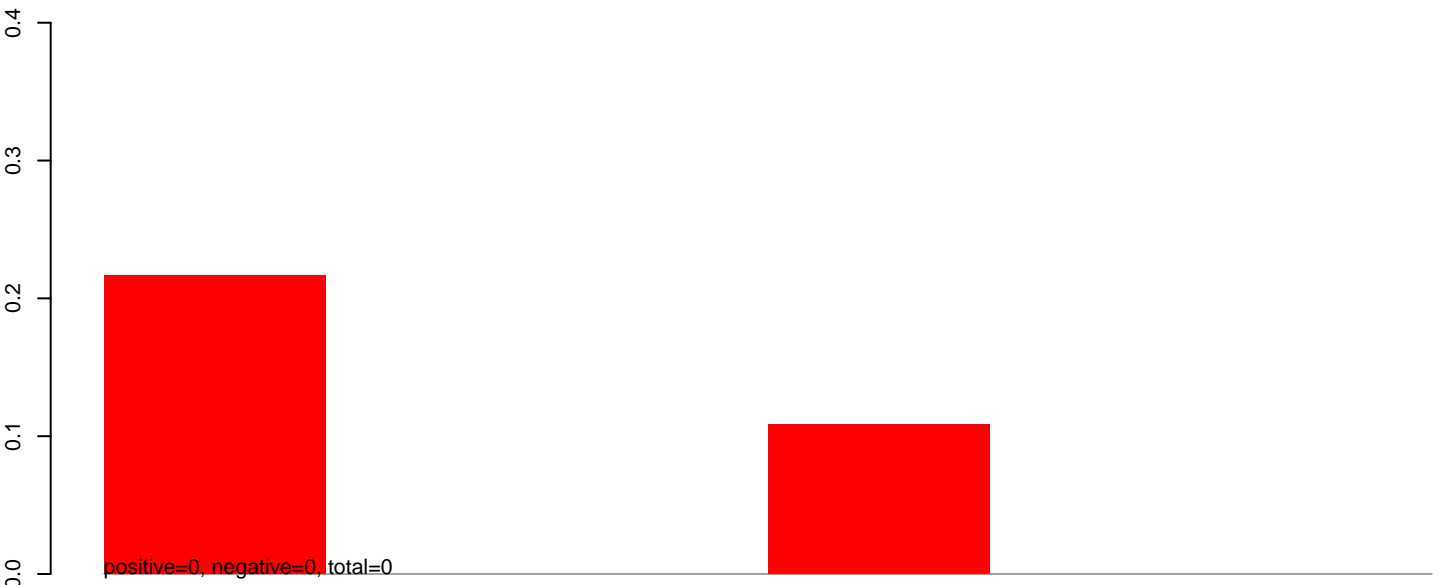


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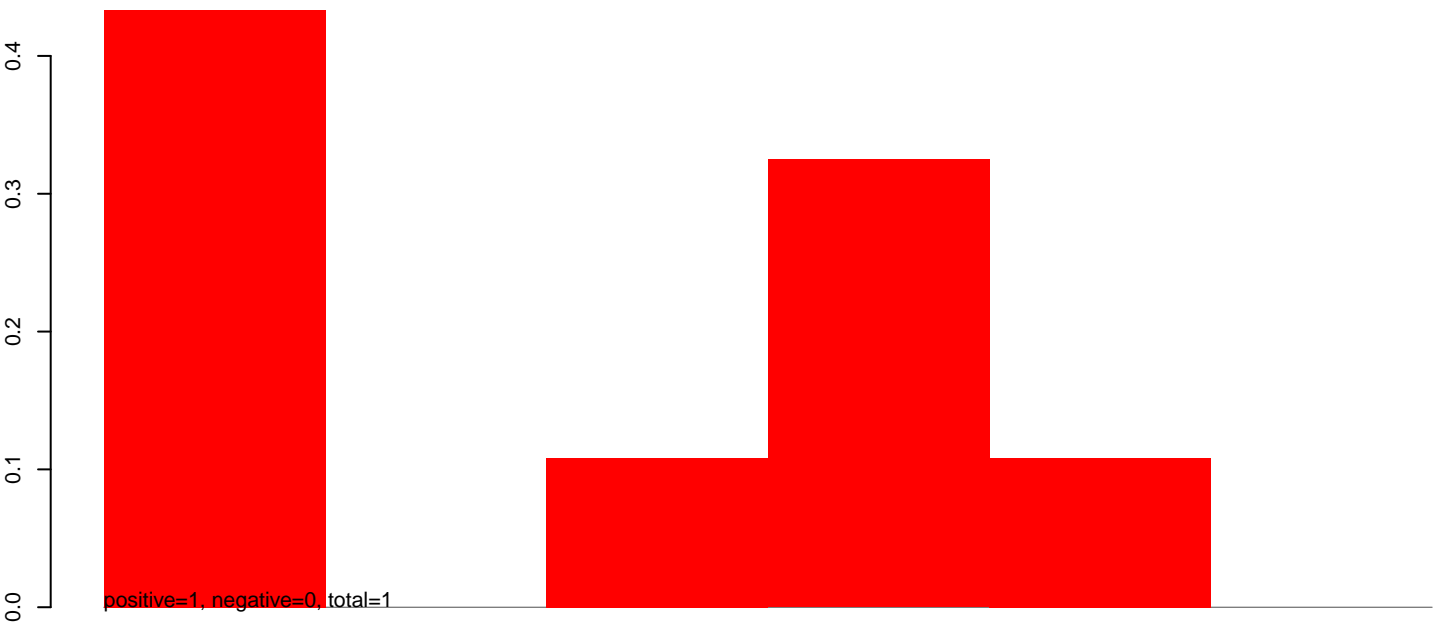


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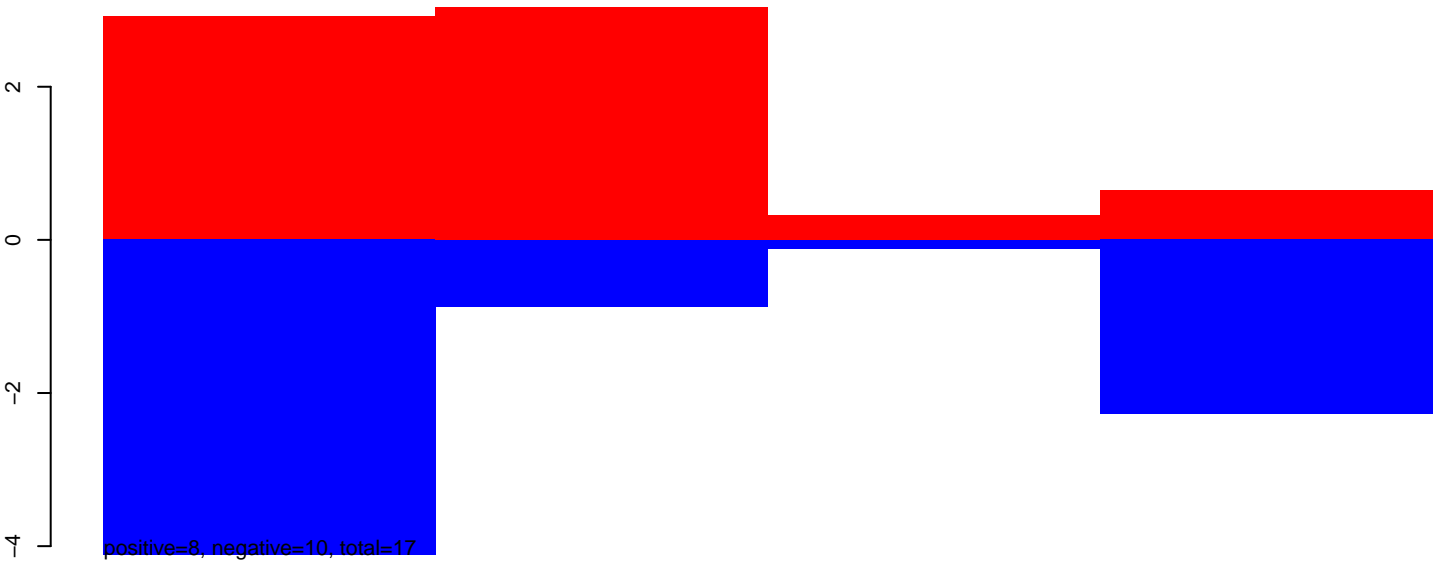
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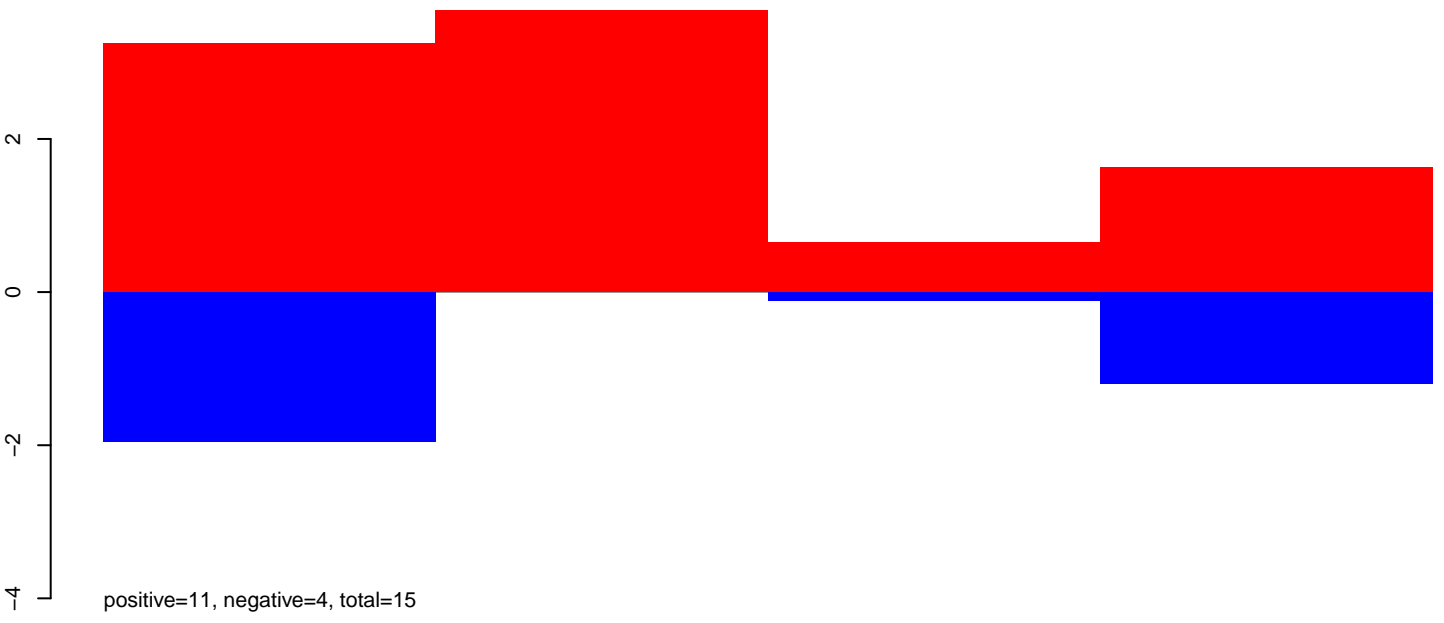
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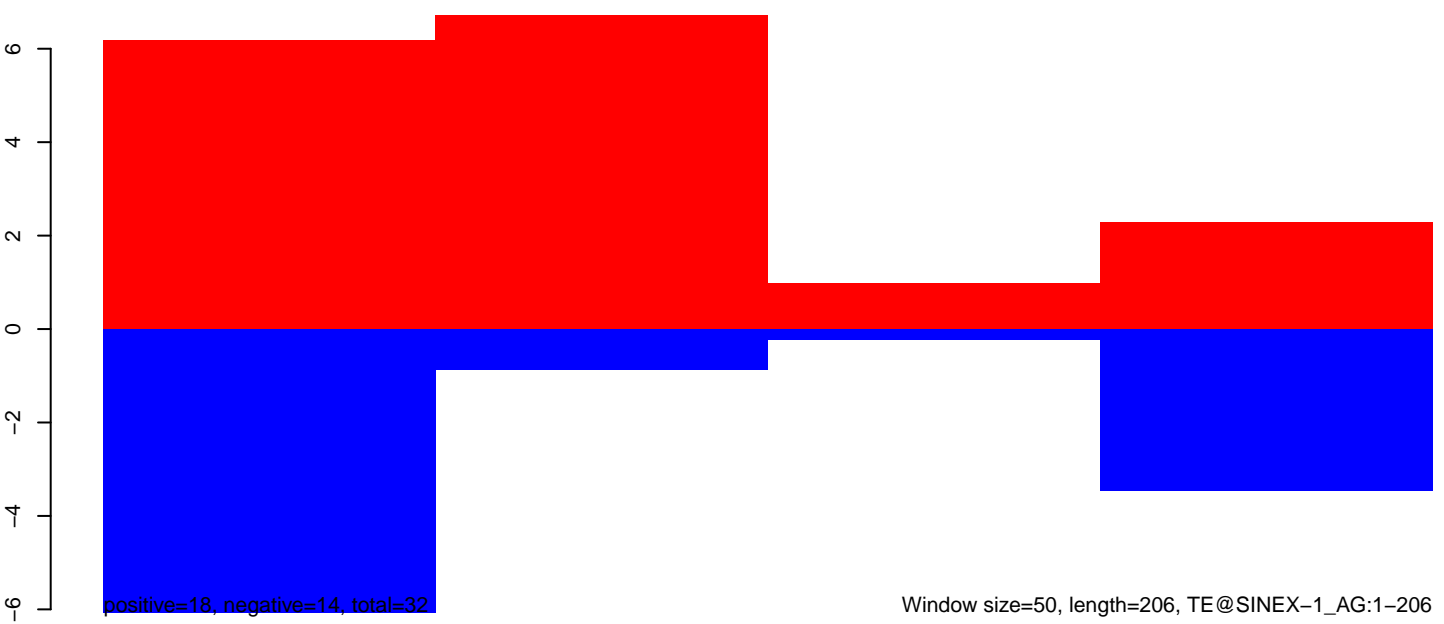
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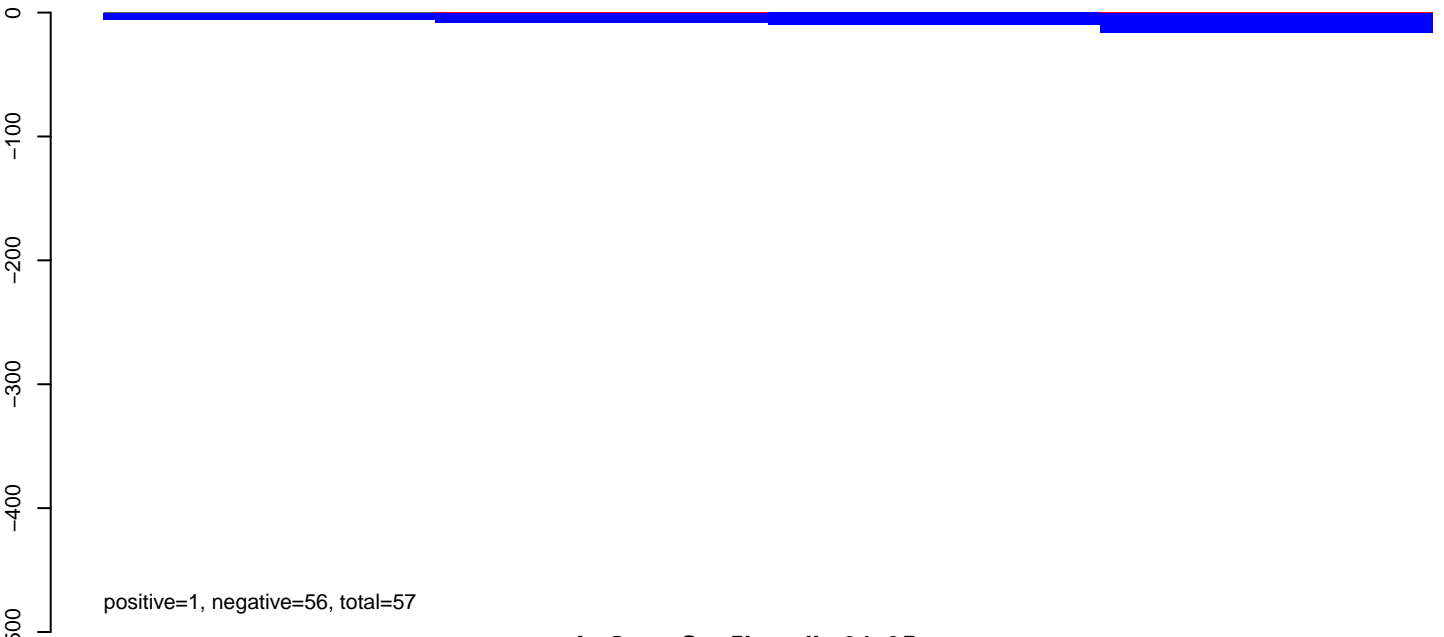
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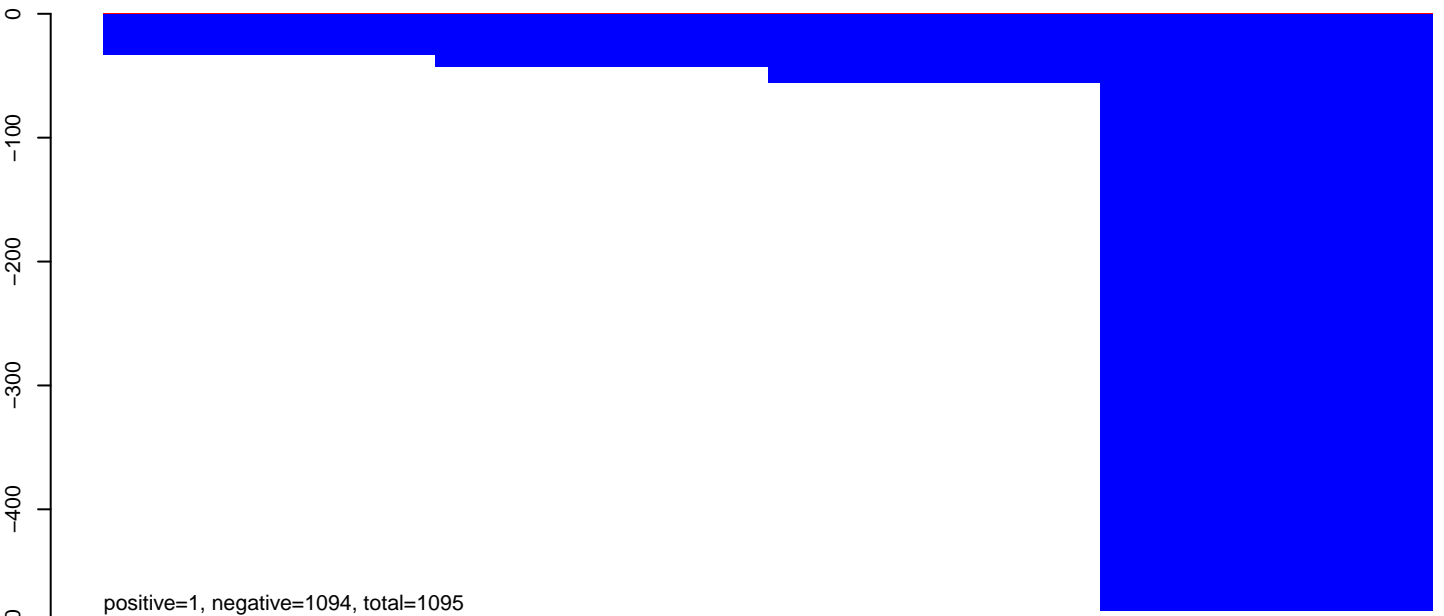
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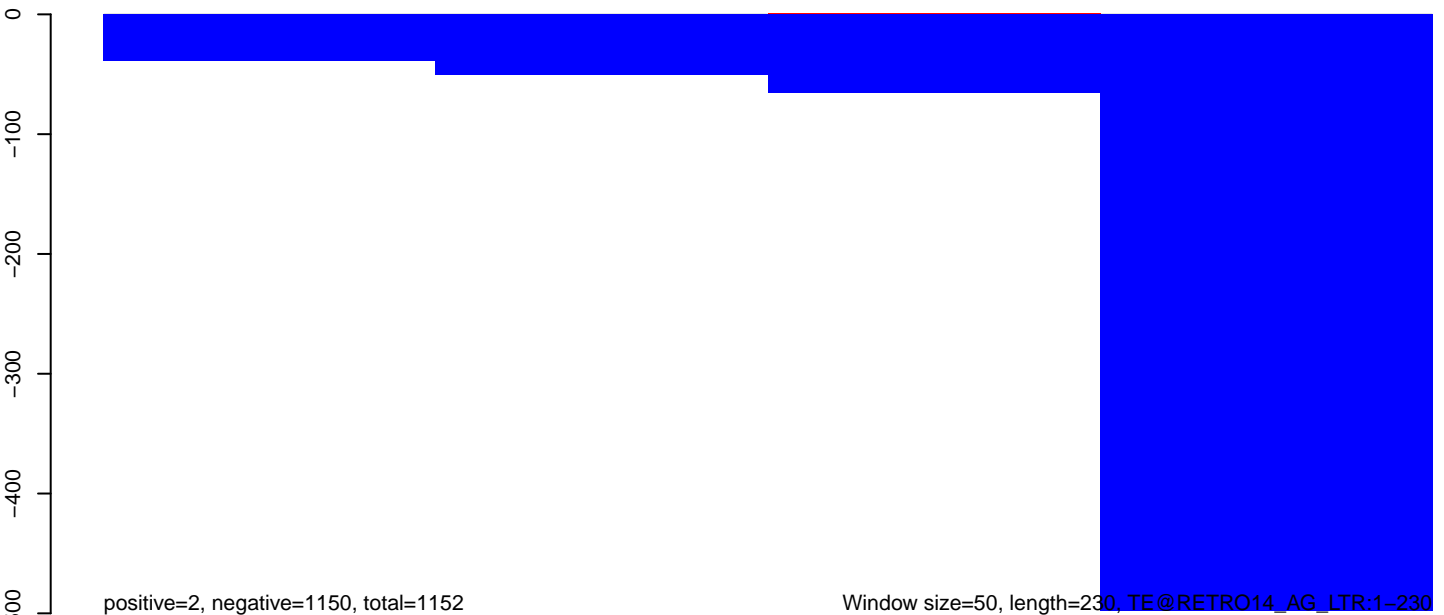
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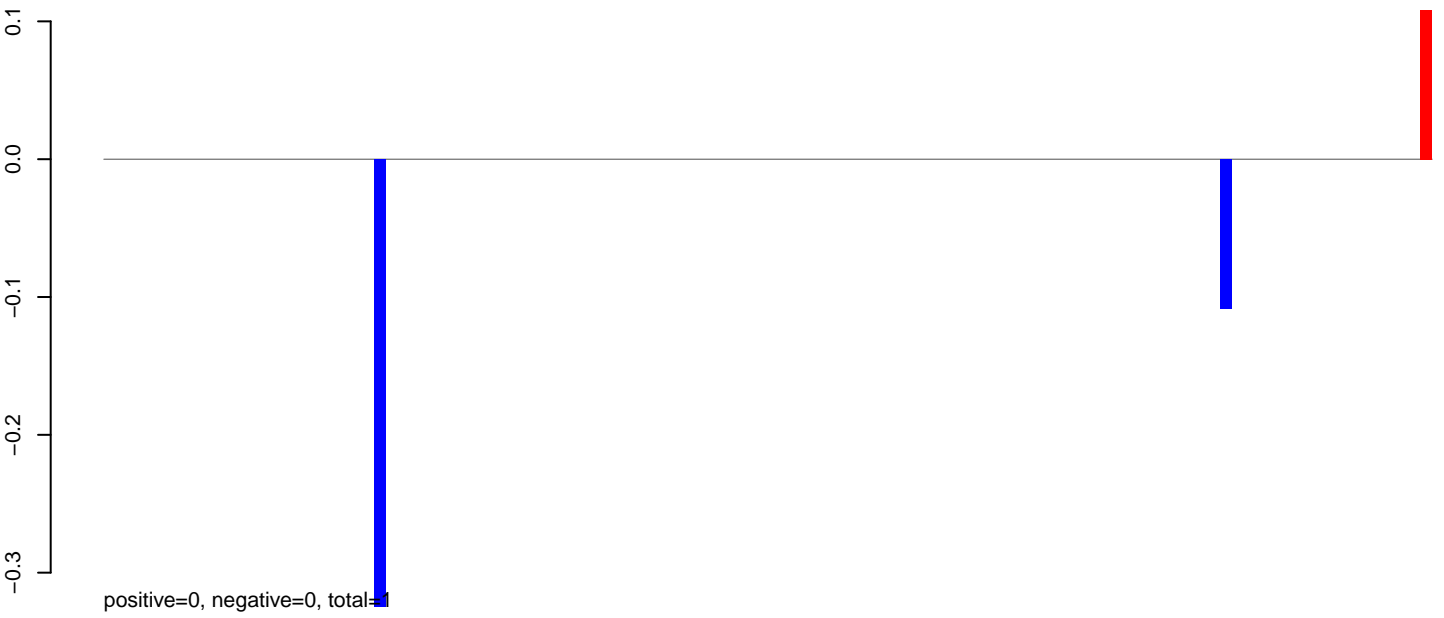


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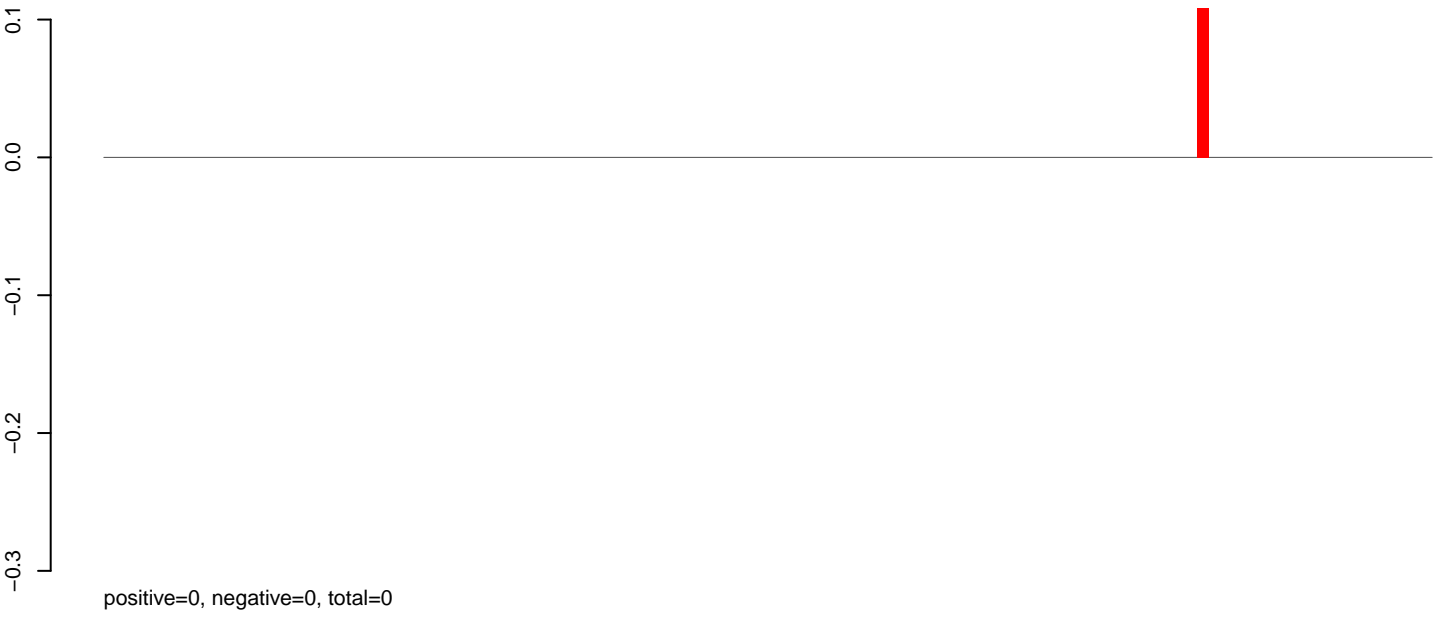


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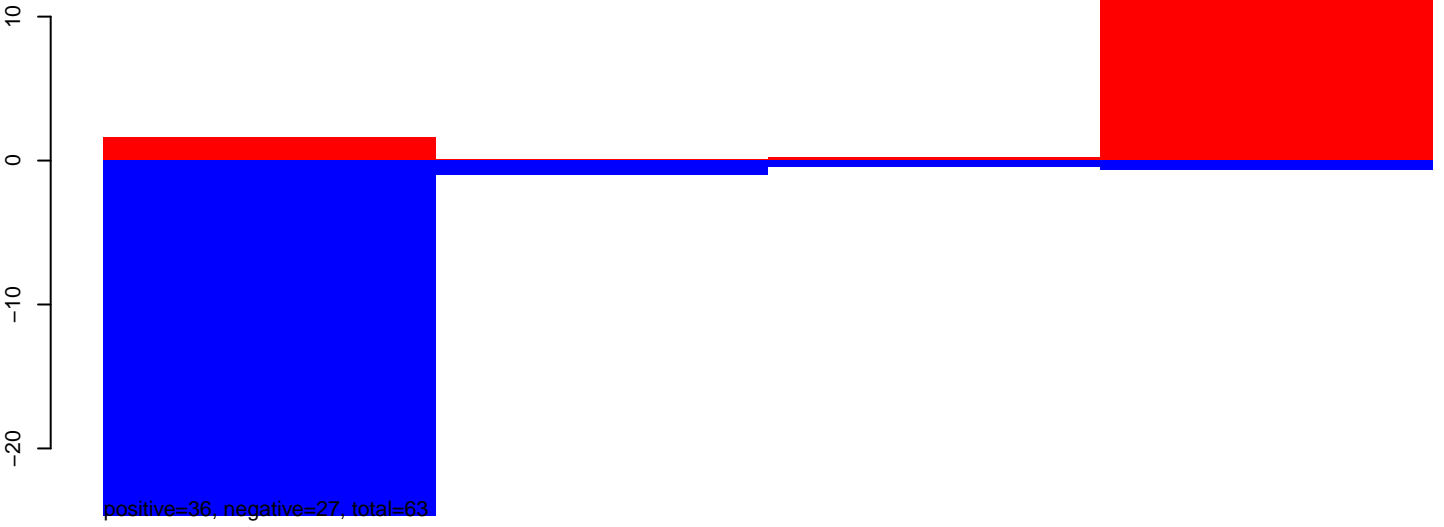
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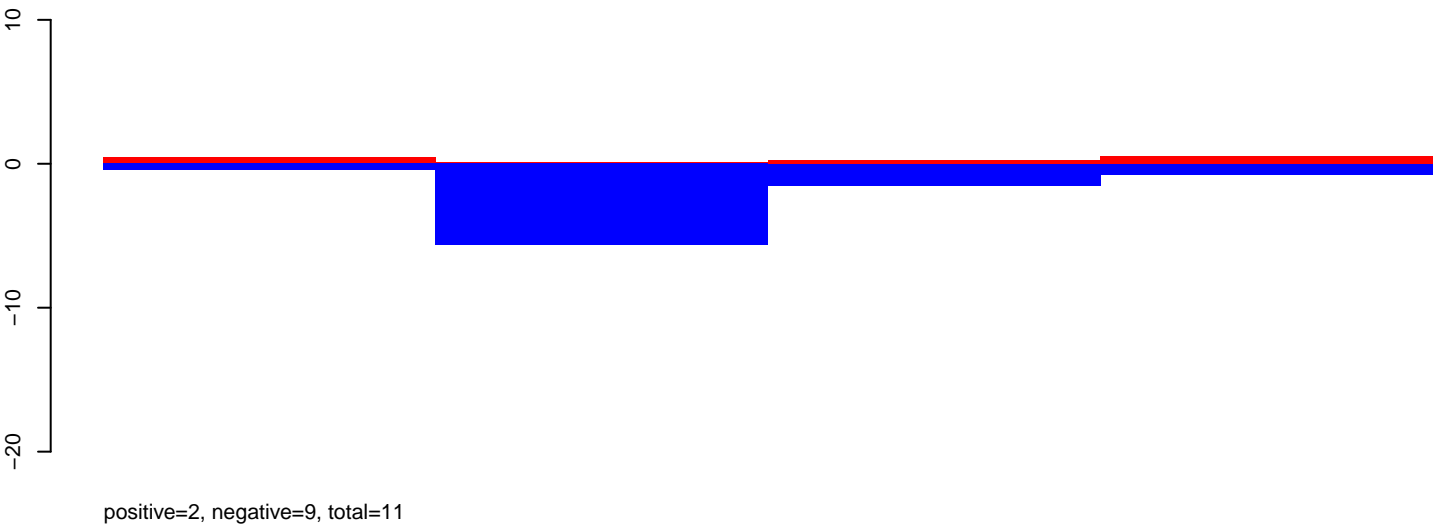
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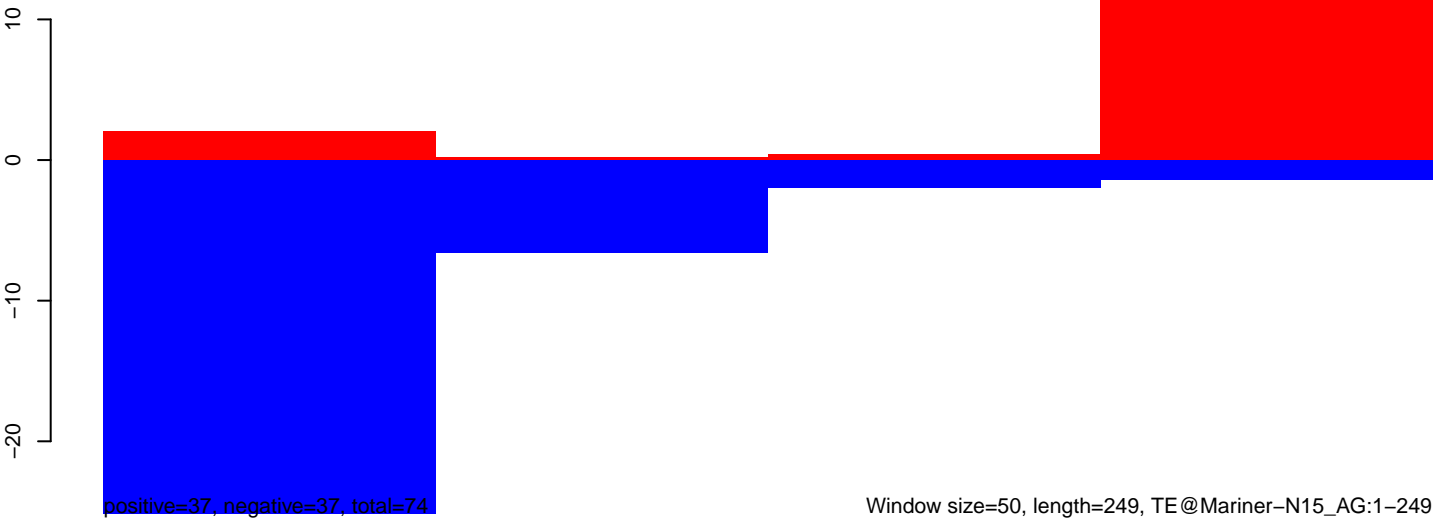
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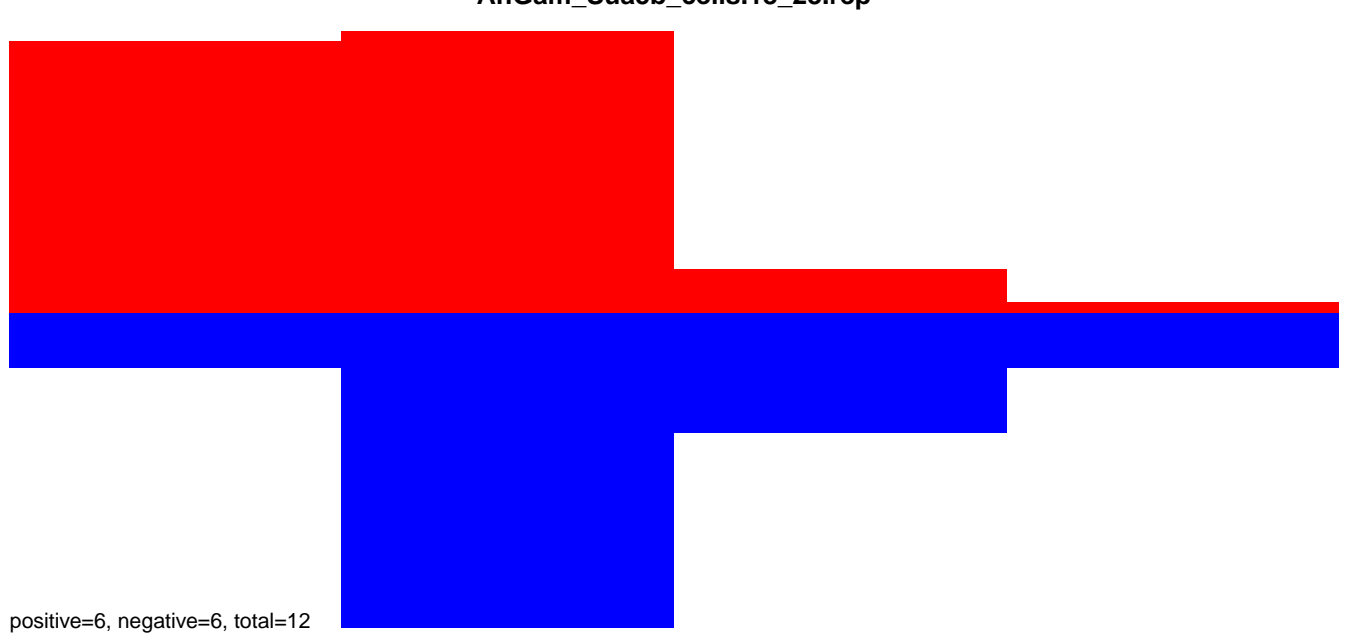
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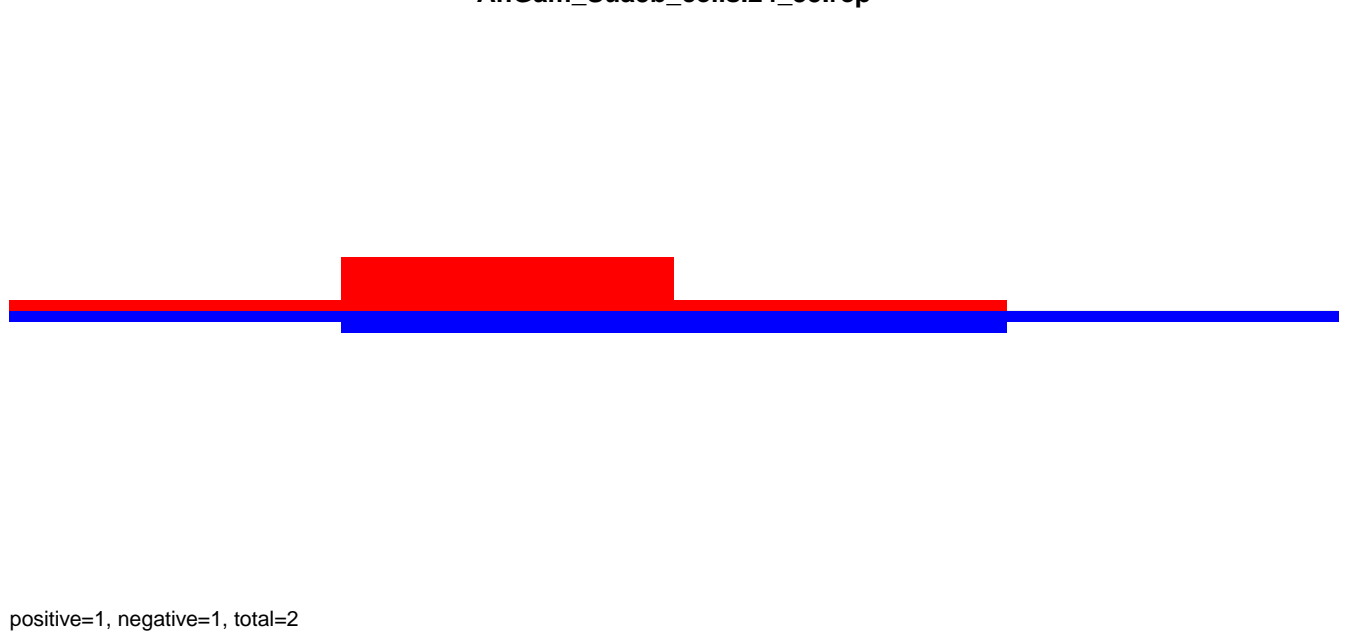
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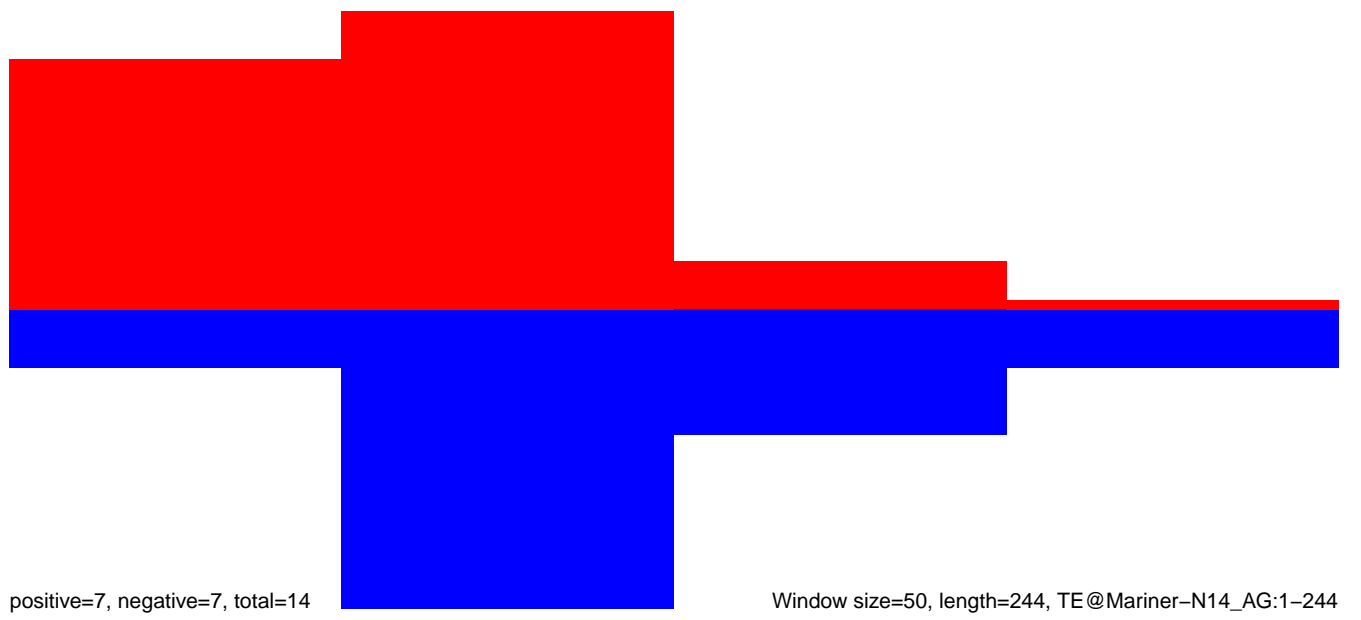
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AnGam_Sua5b_cells.24_35.rep



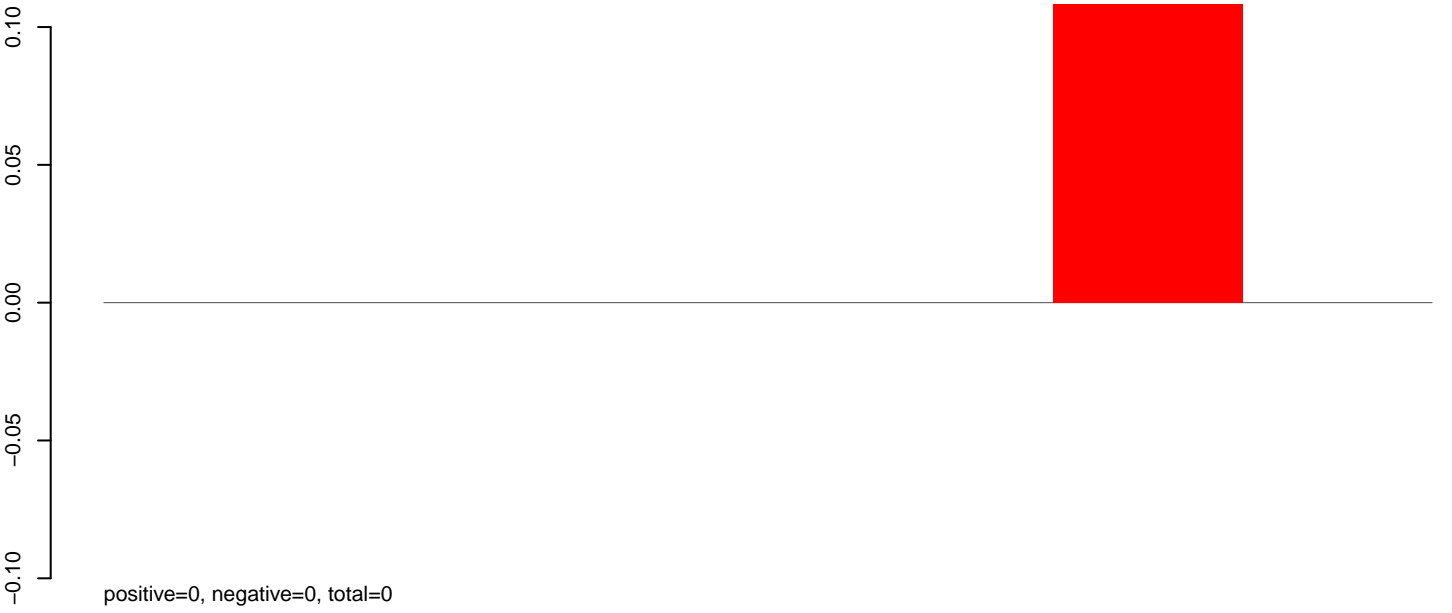
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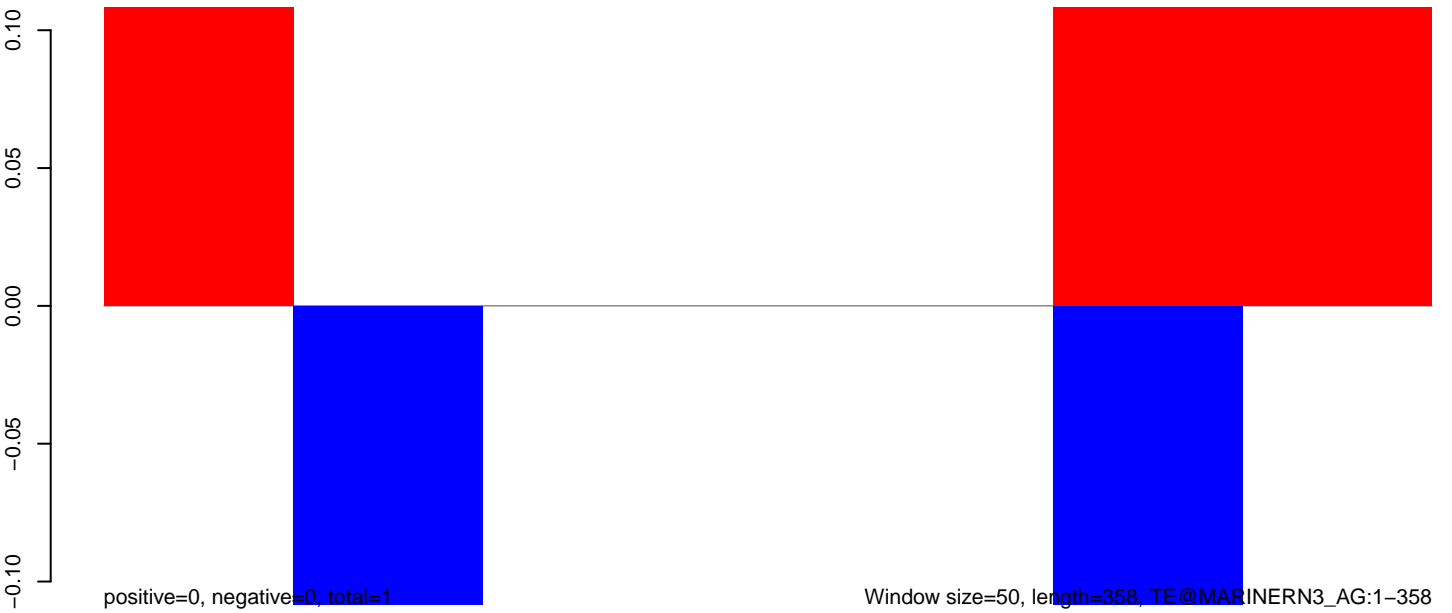
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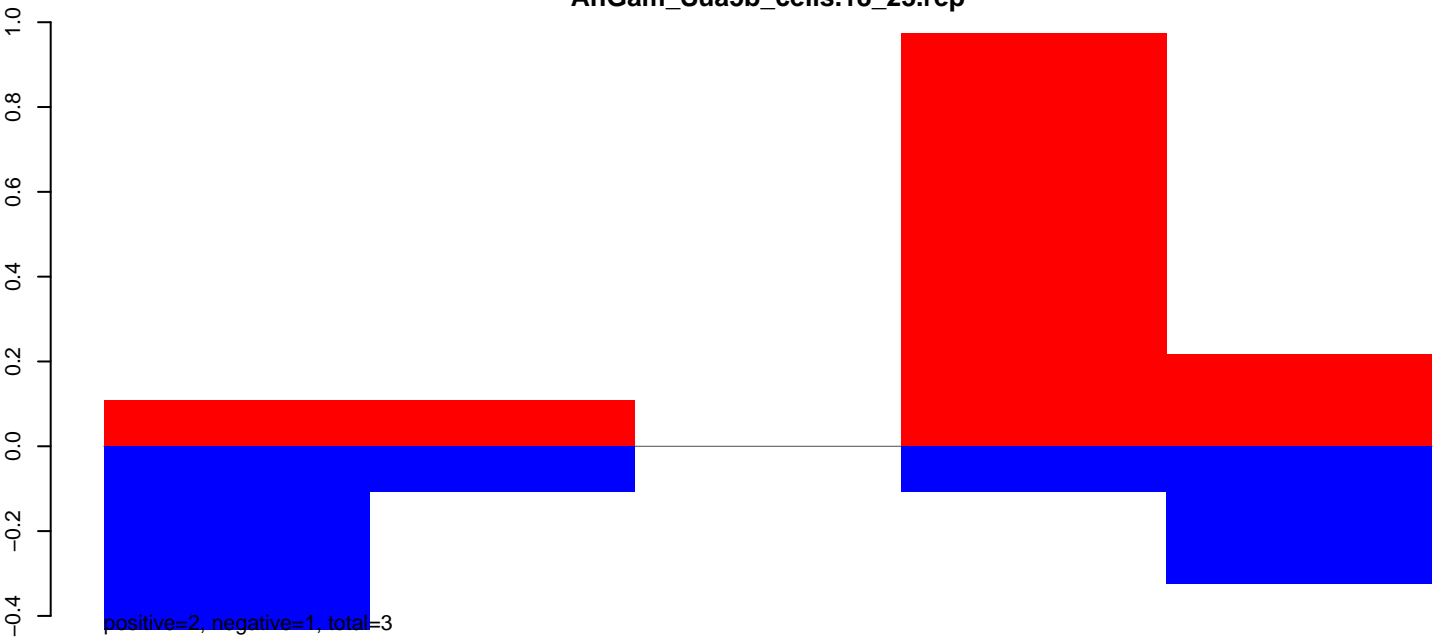
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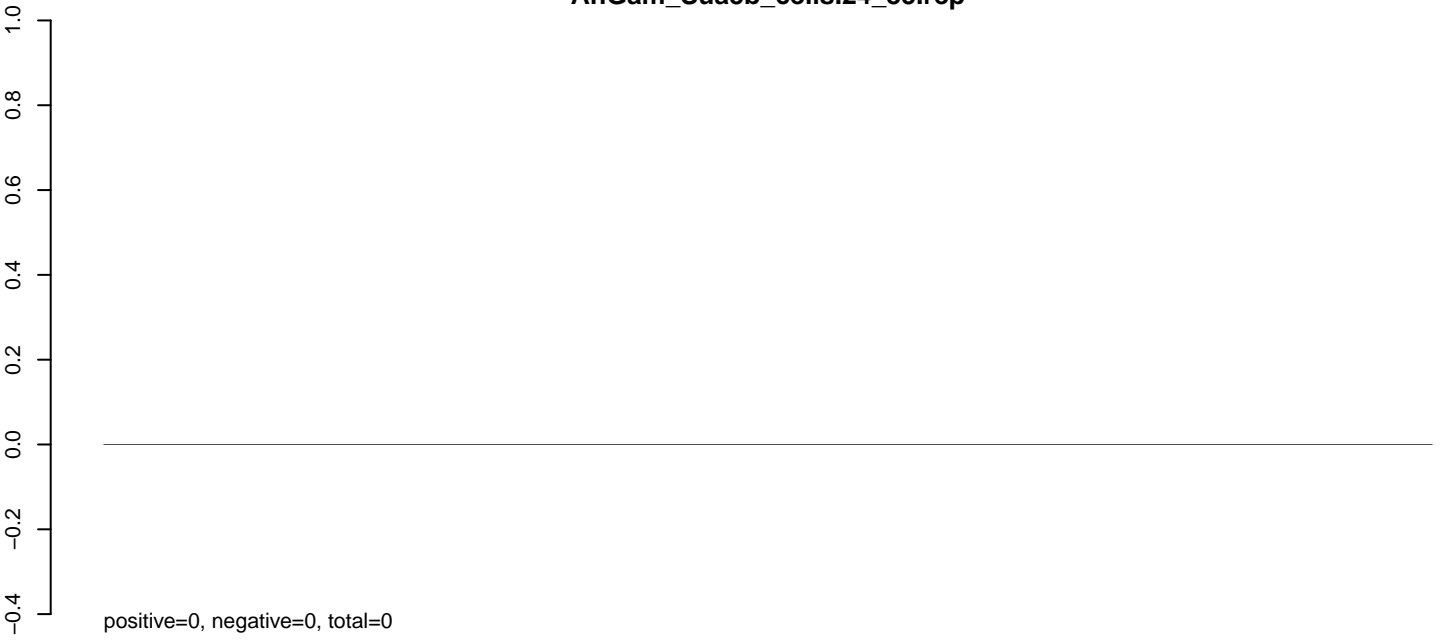
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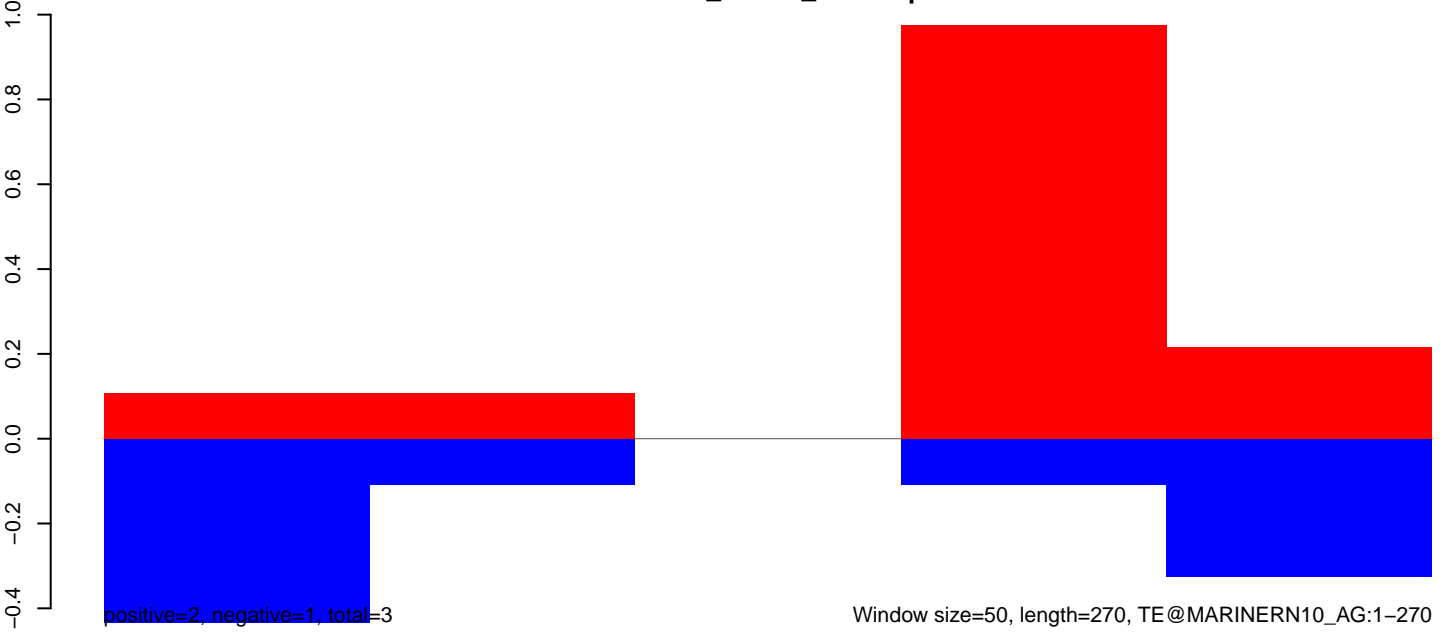
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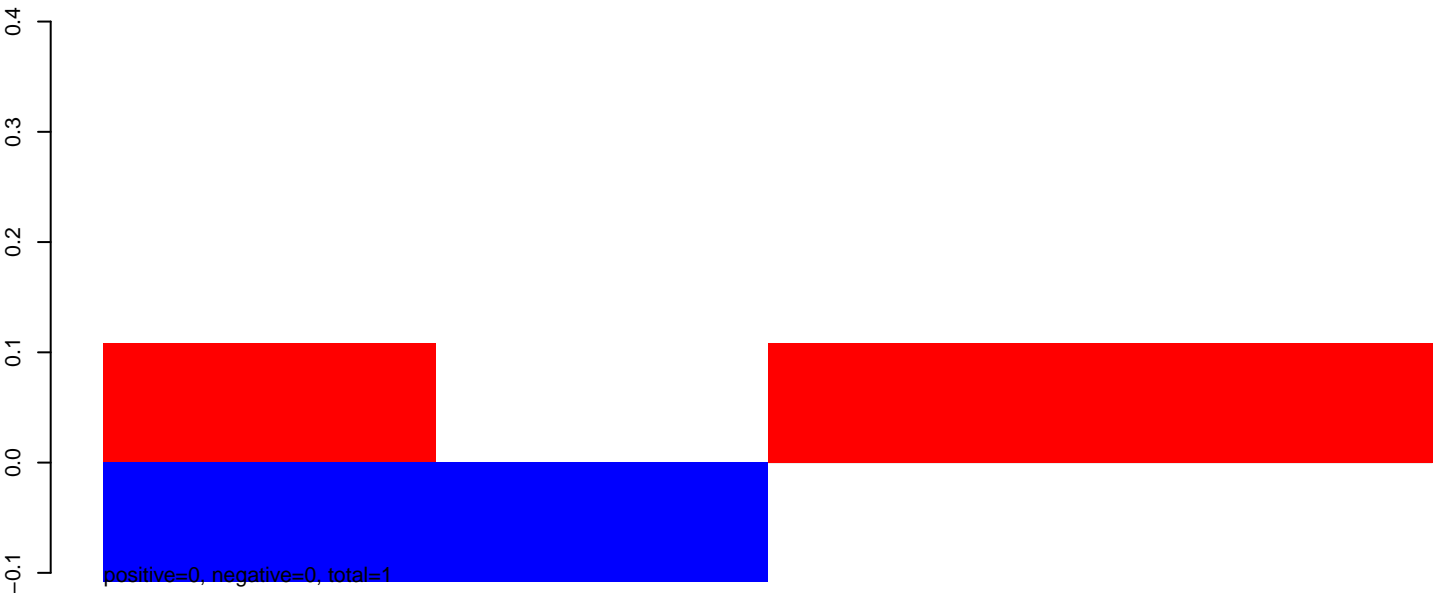
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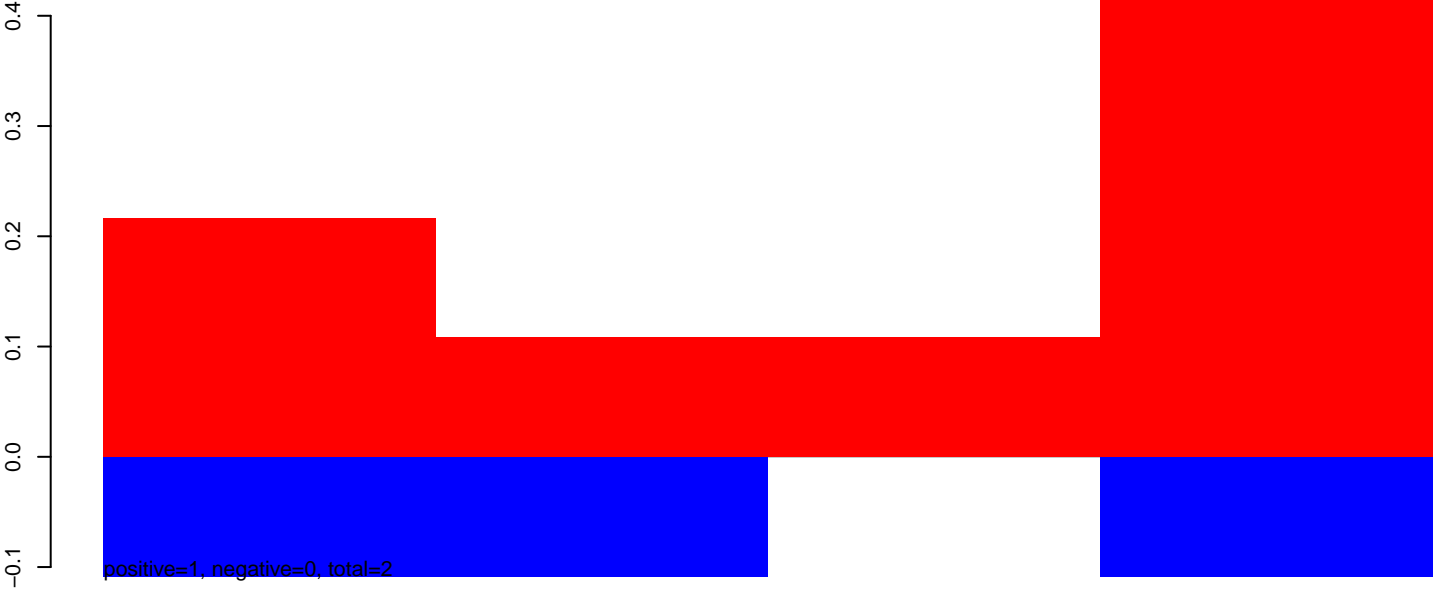
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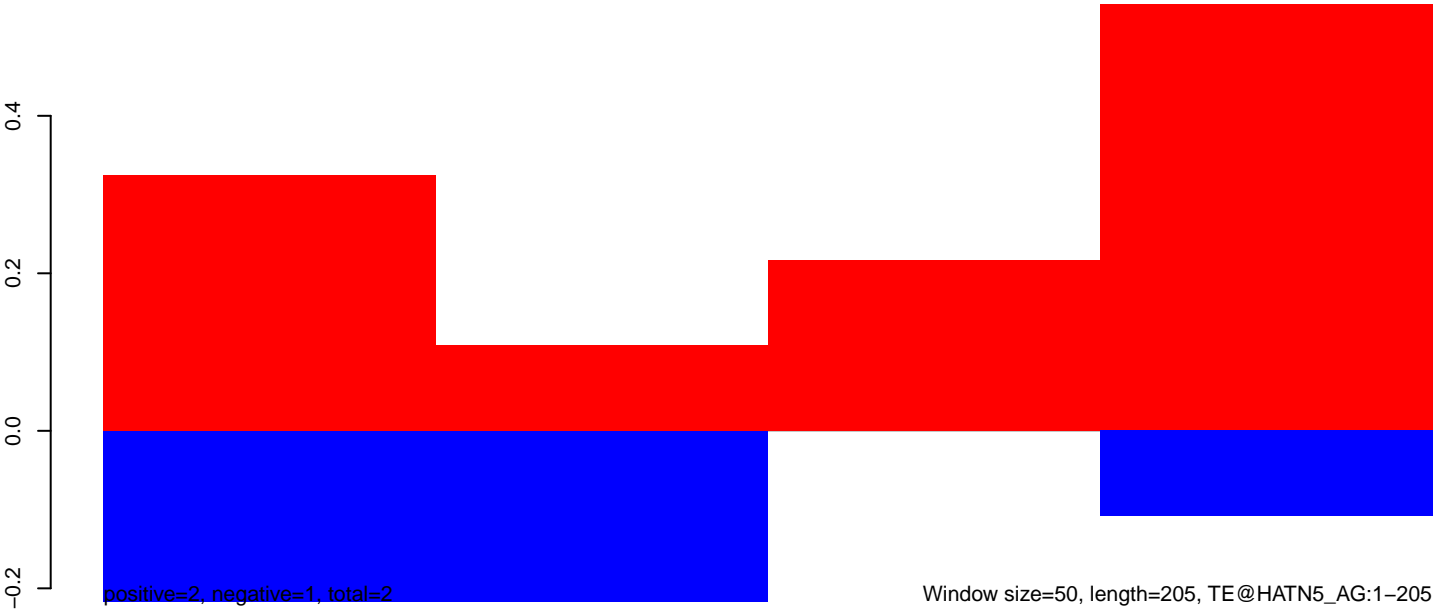
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AnGam_Sua5b_cells.24_35.rep

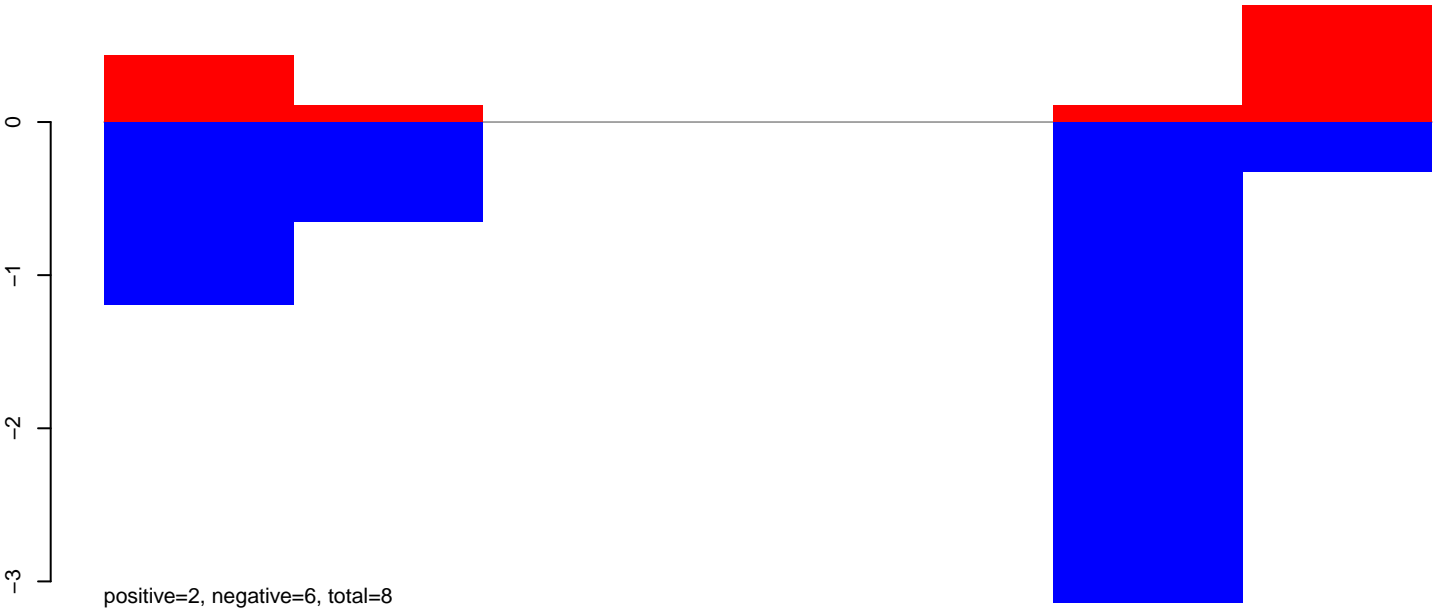


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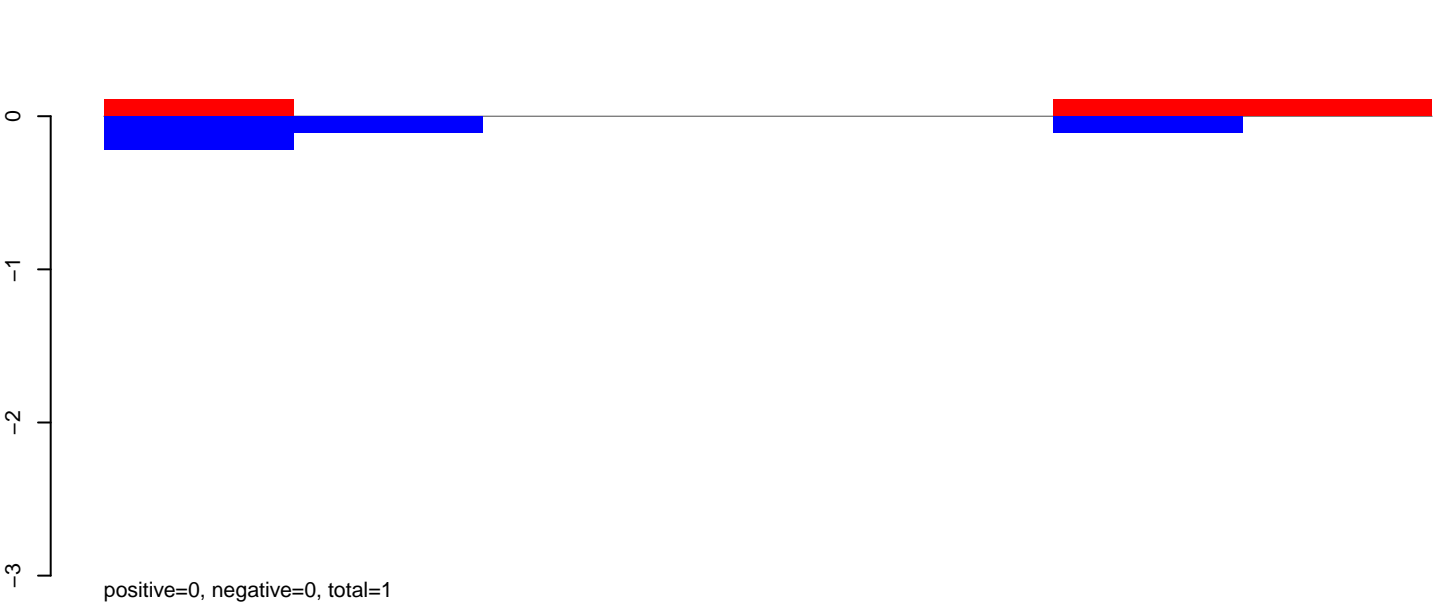


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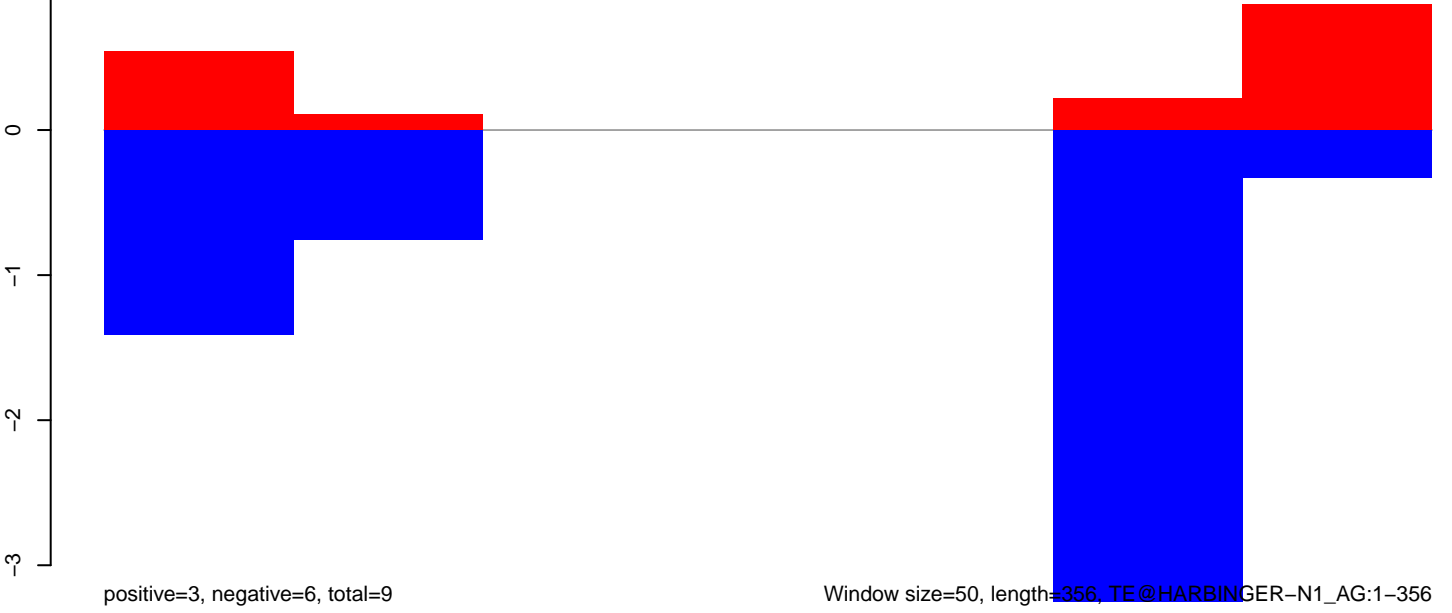
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AnGam_Sua5b_cells.24_35.rep



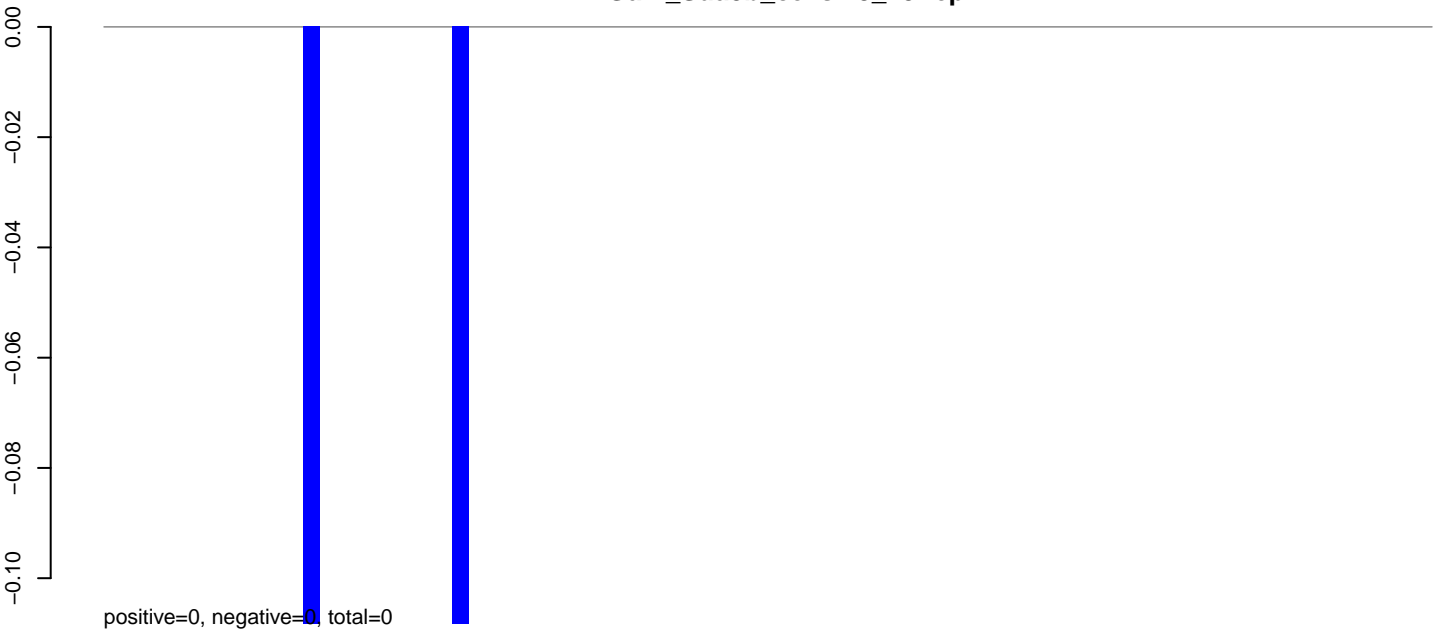
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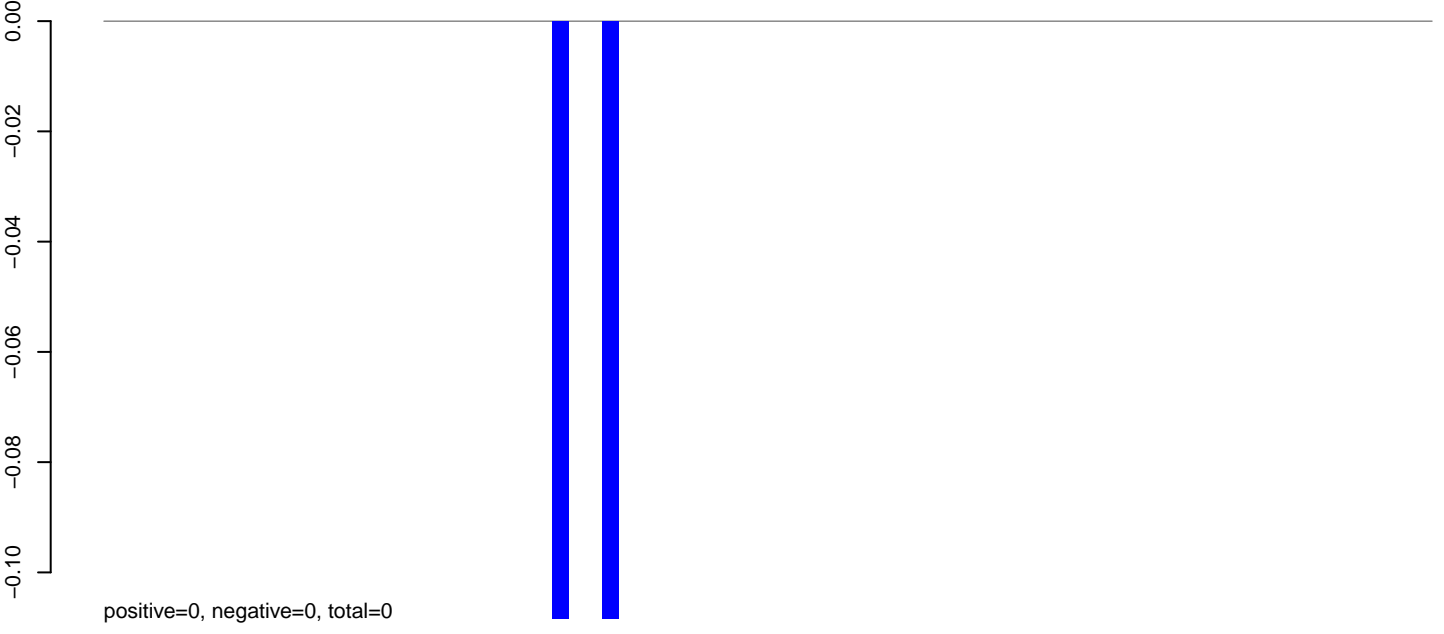
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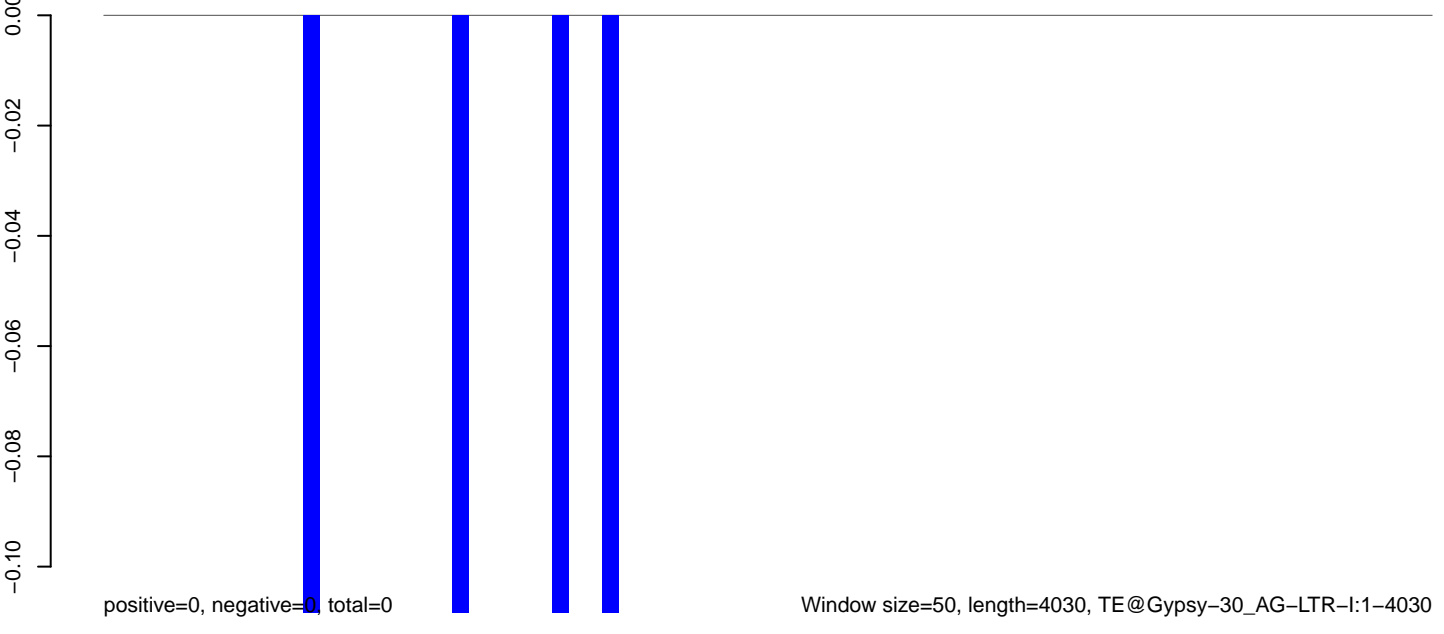
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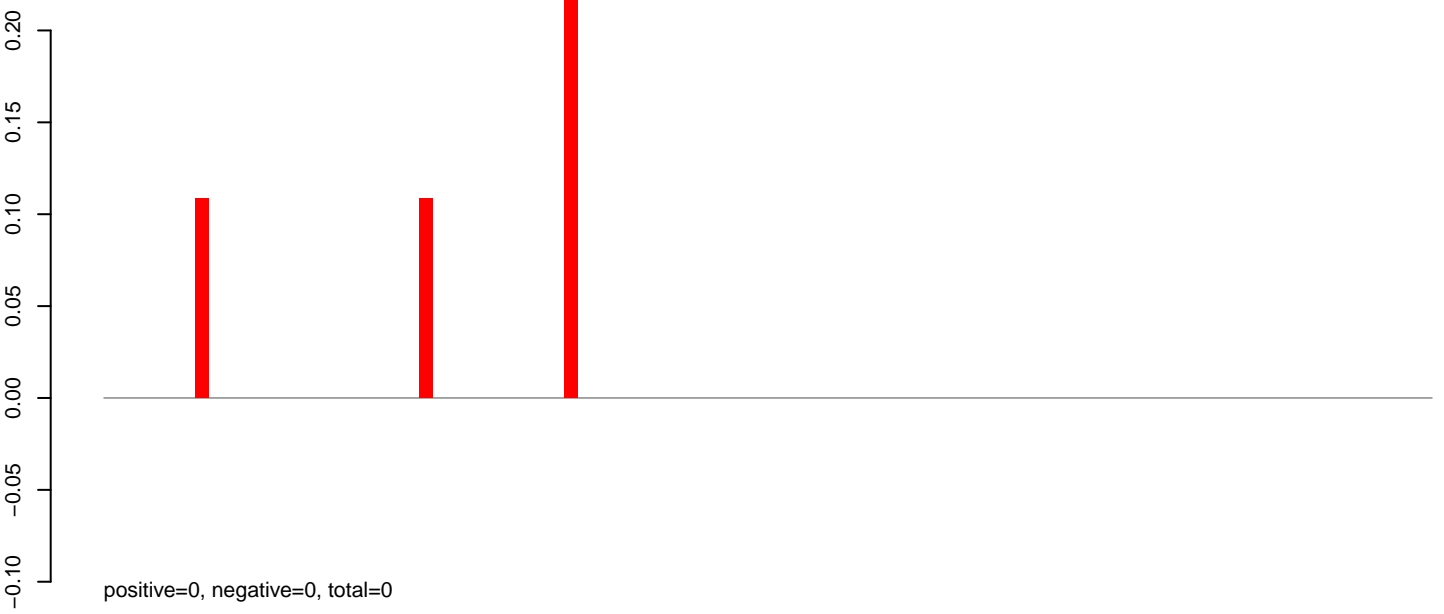
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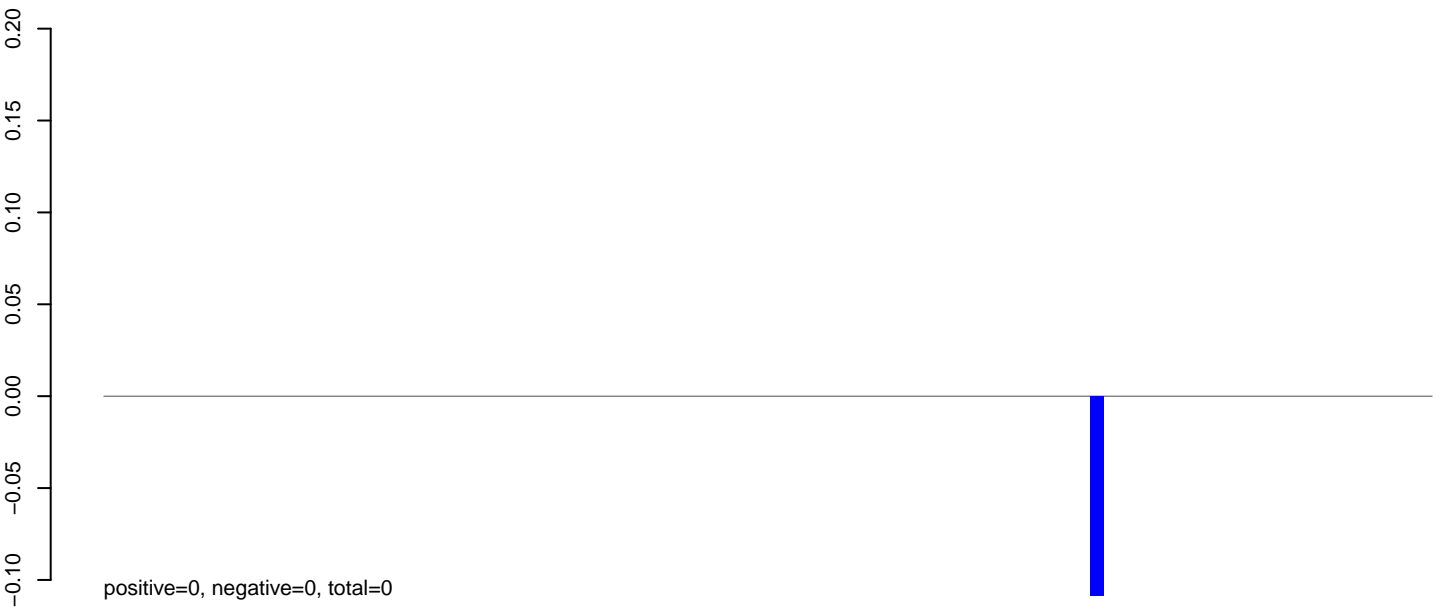
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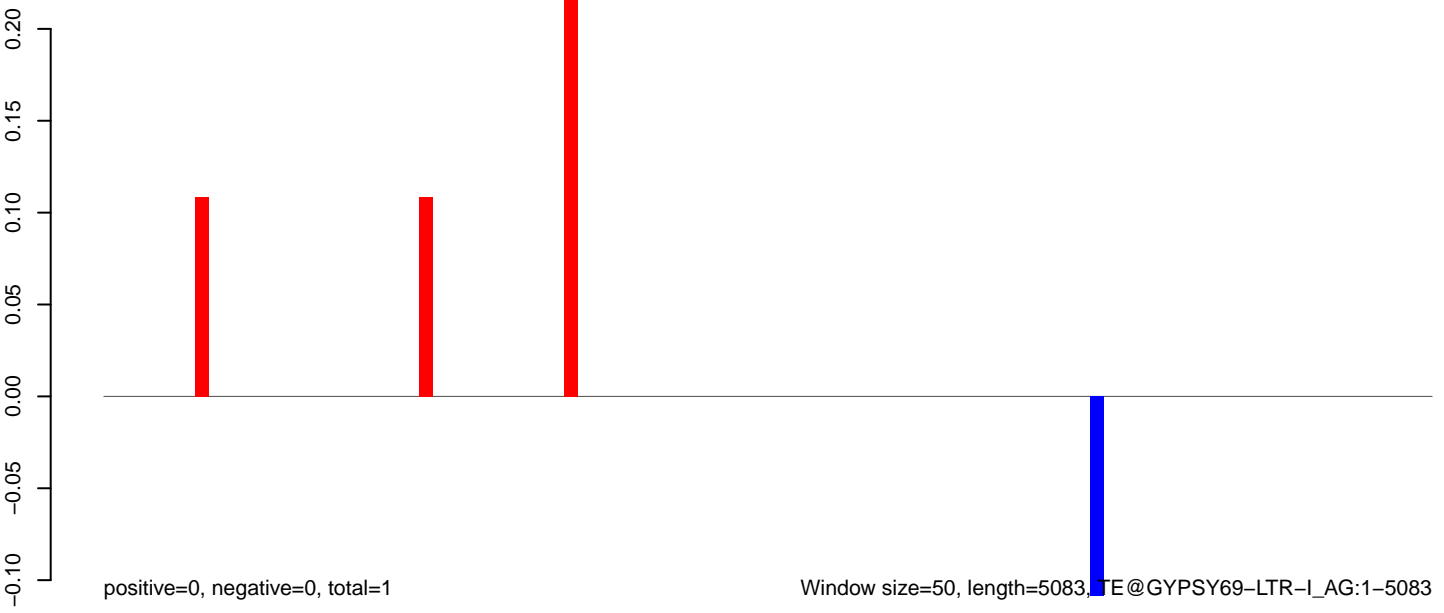
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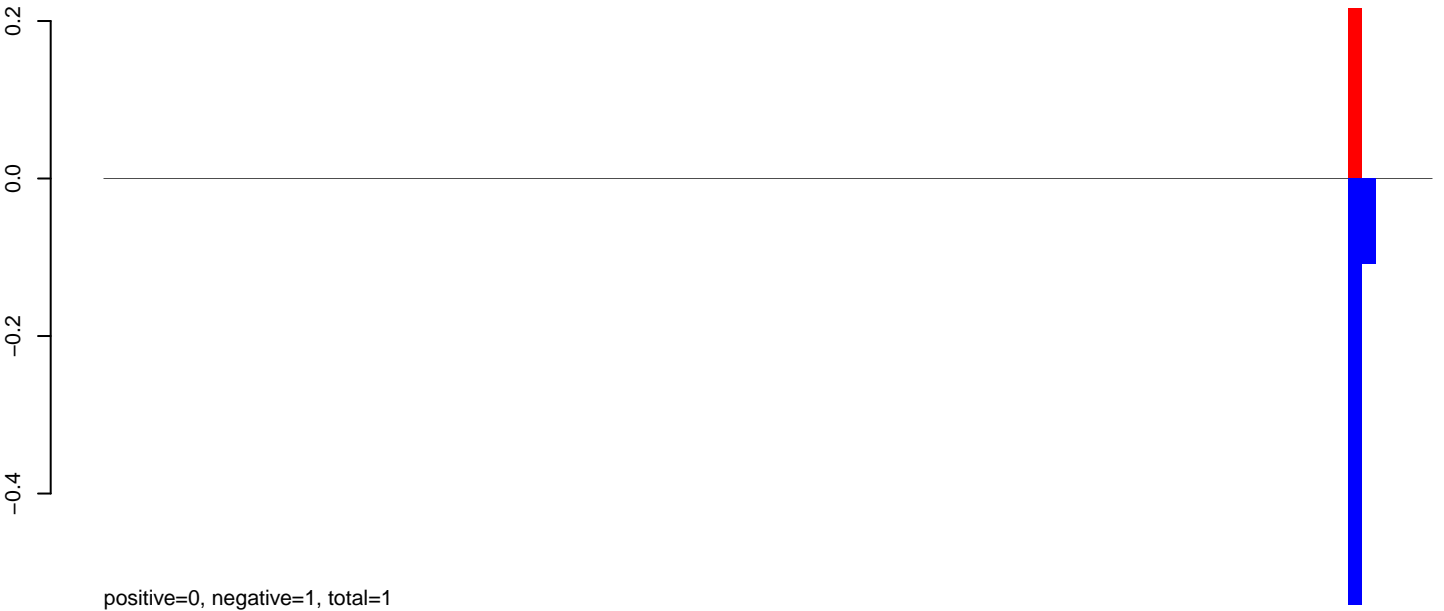
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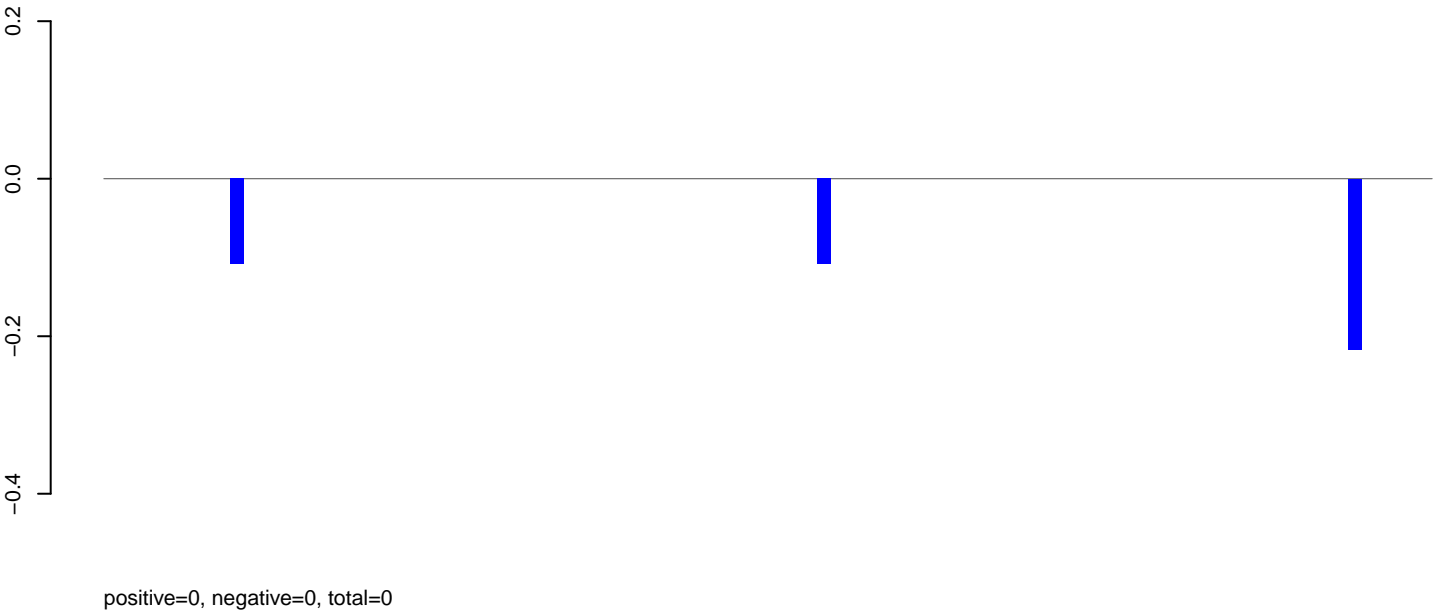
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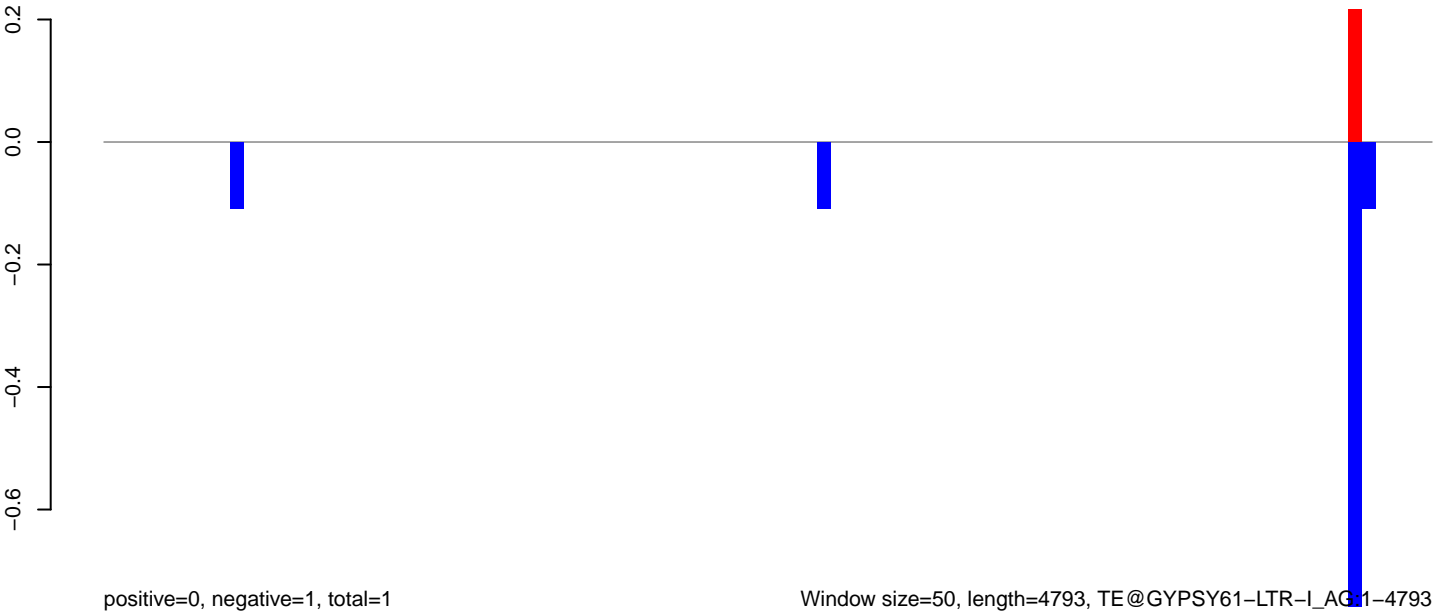
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AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



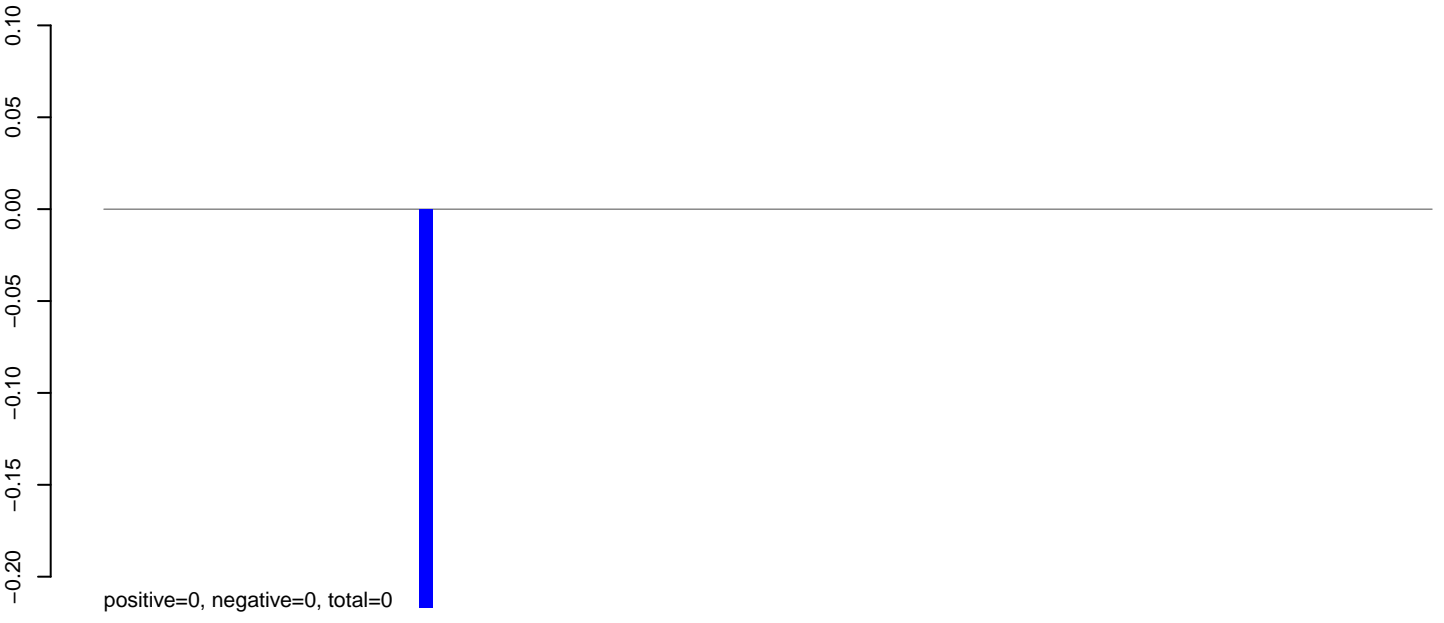
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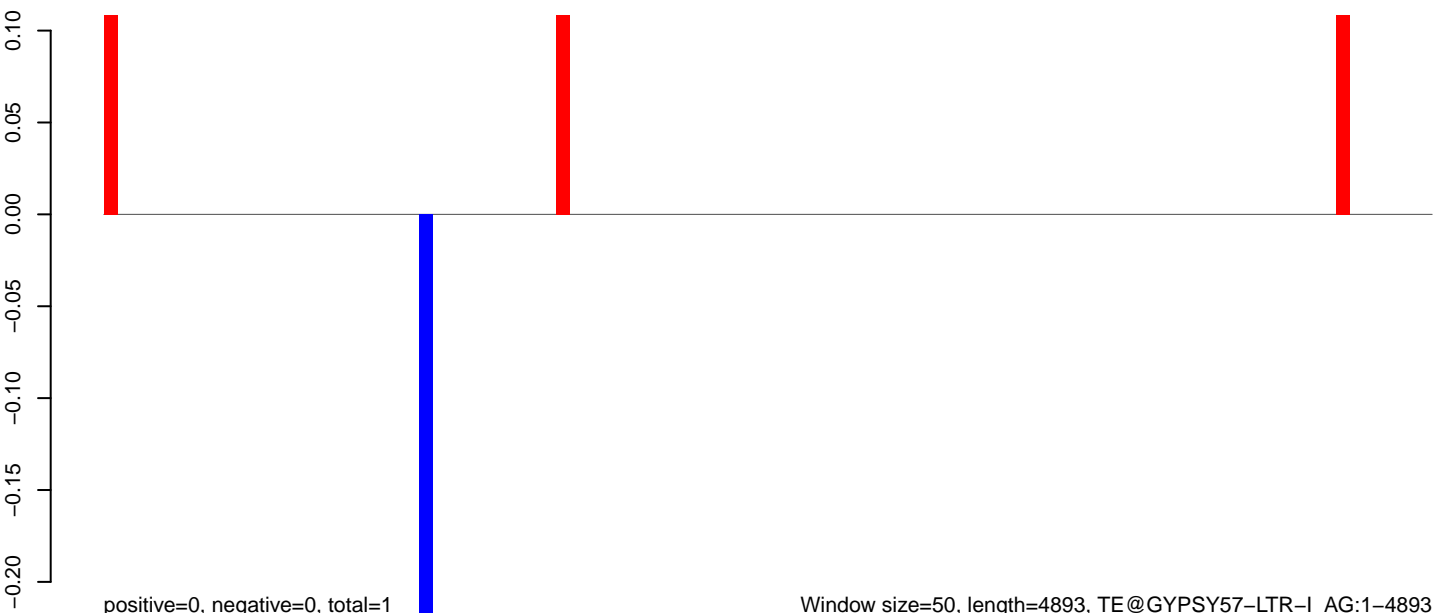
AnGam_Sua5b_cells.18_23.rep



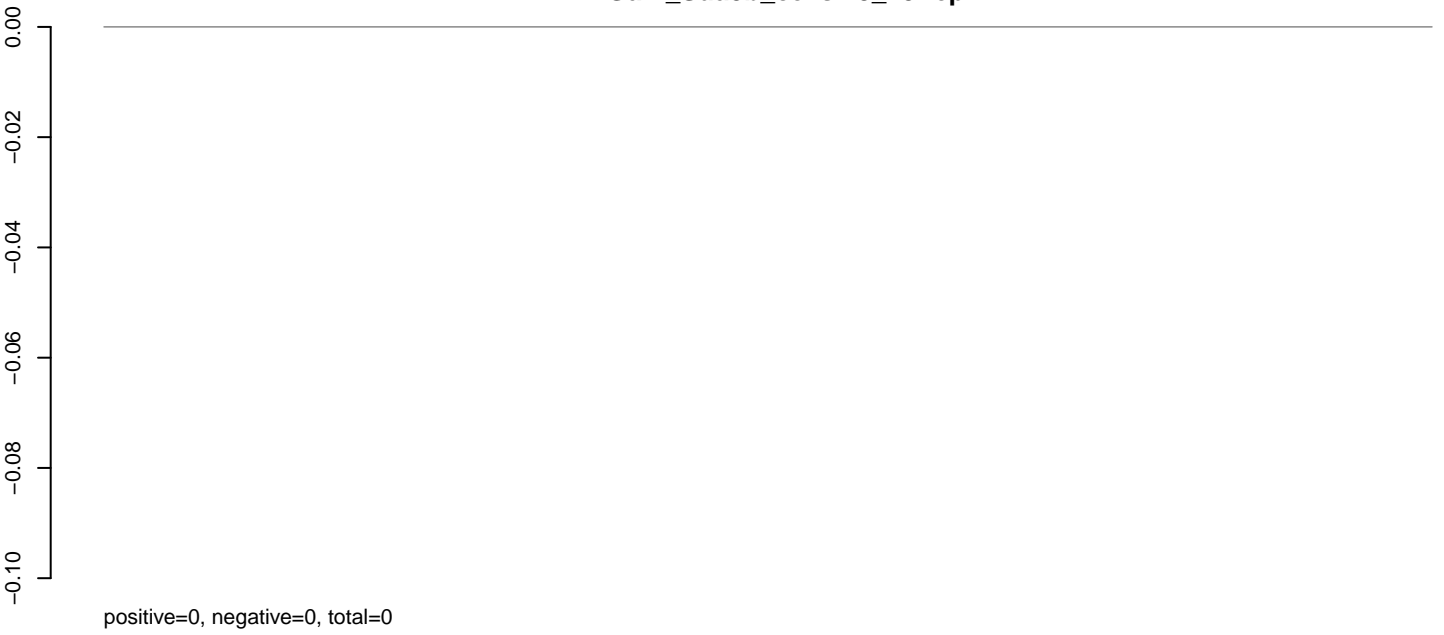
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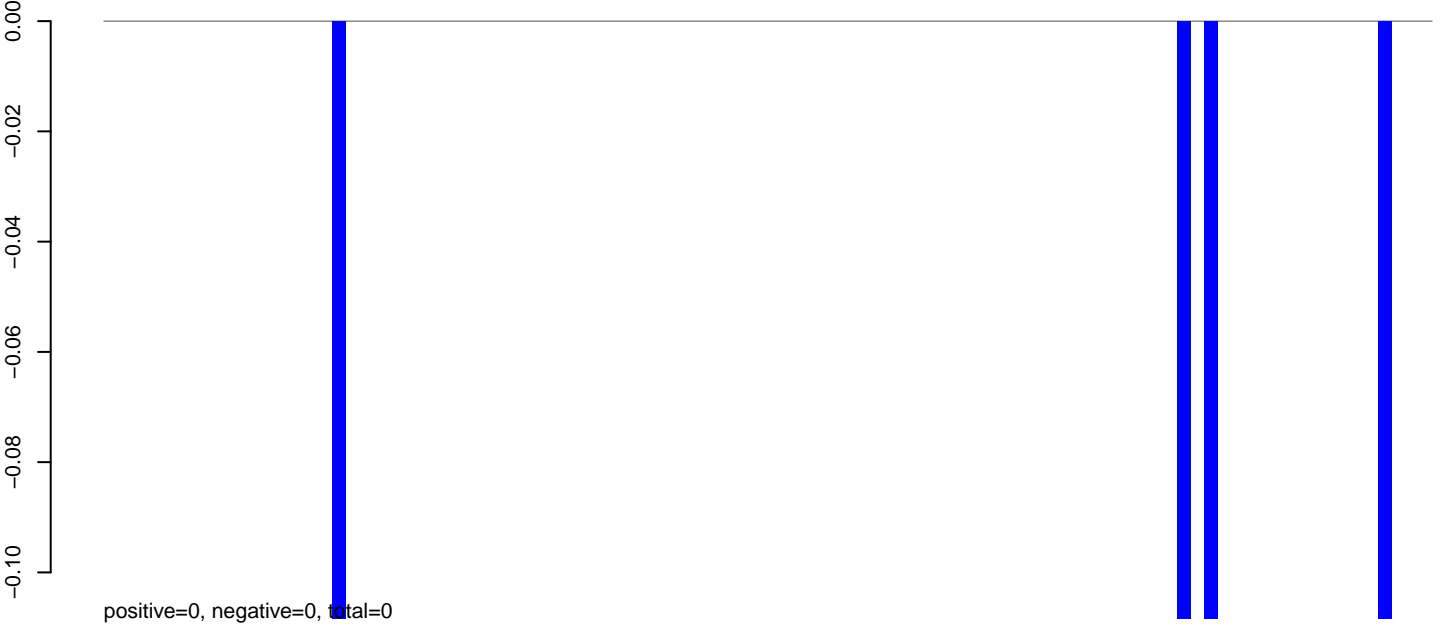
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

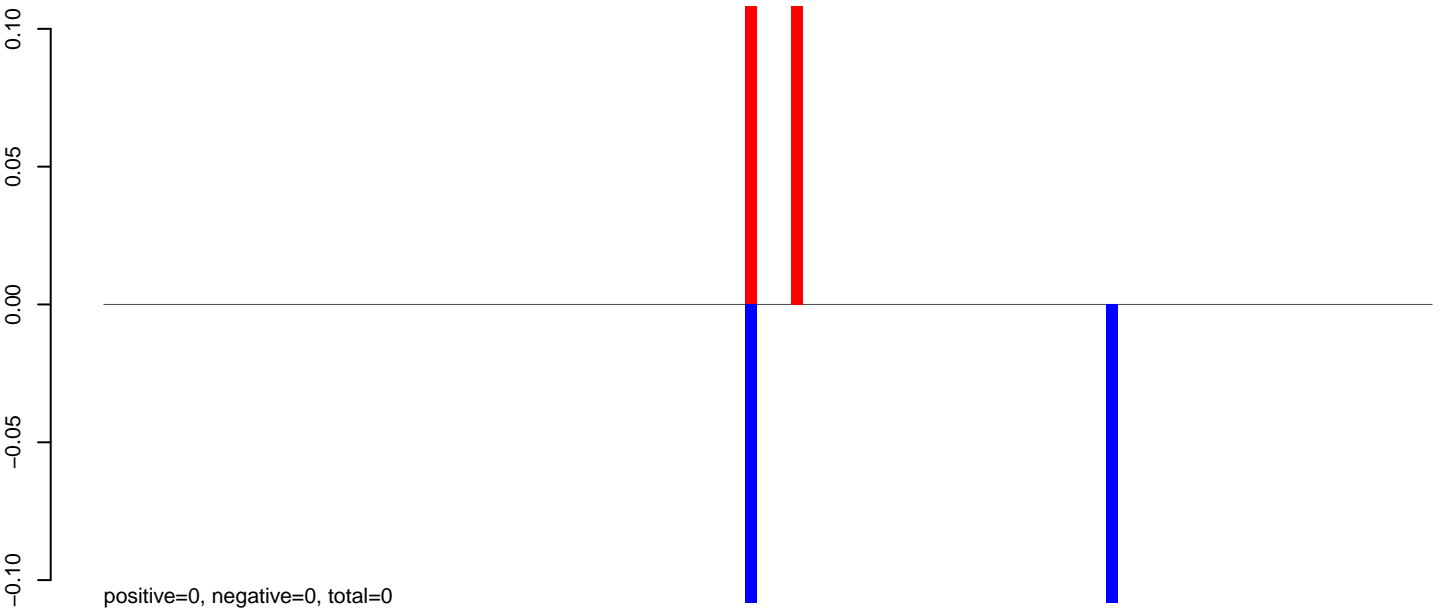


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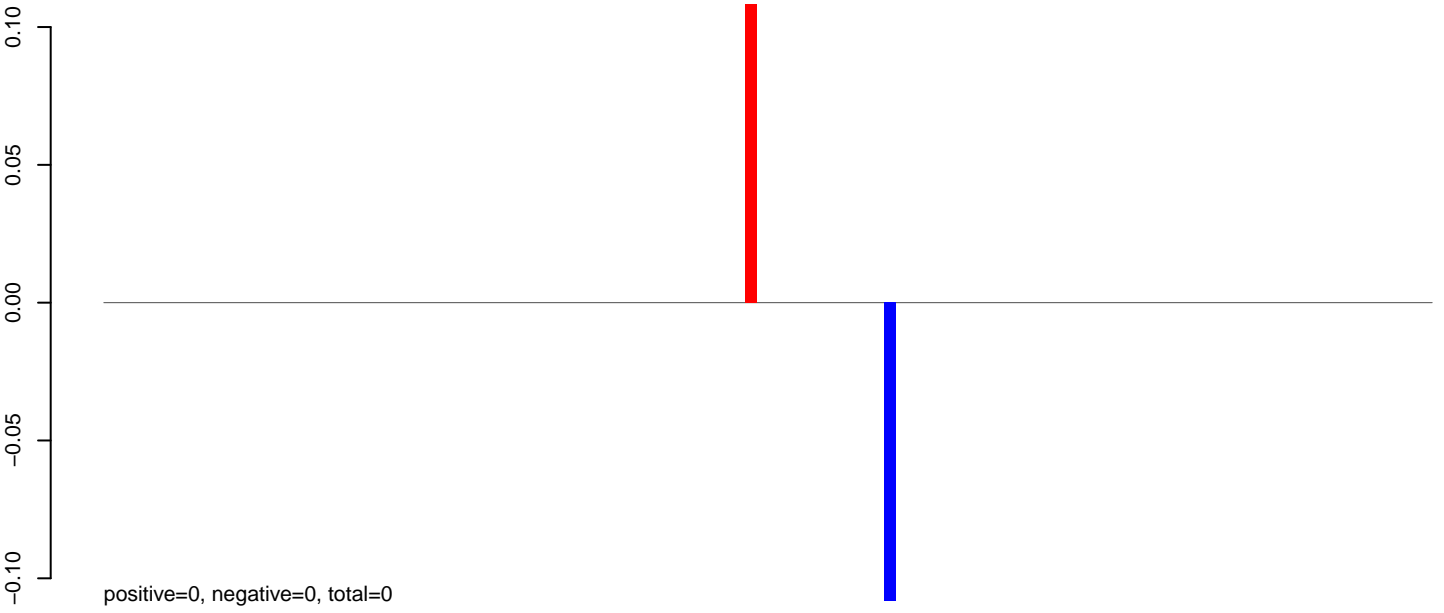


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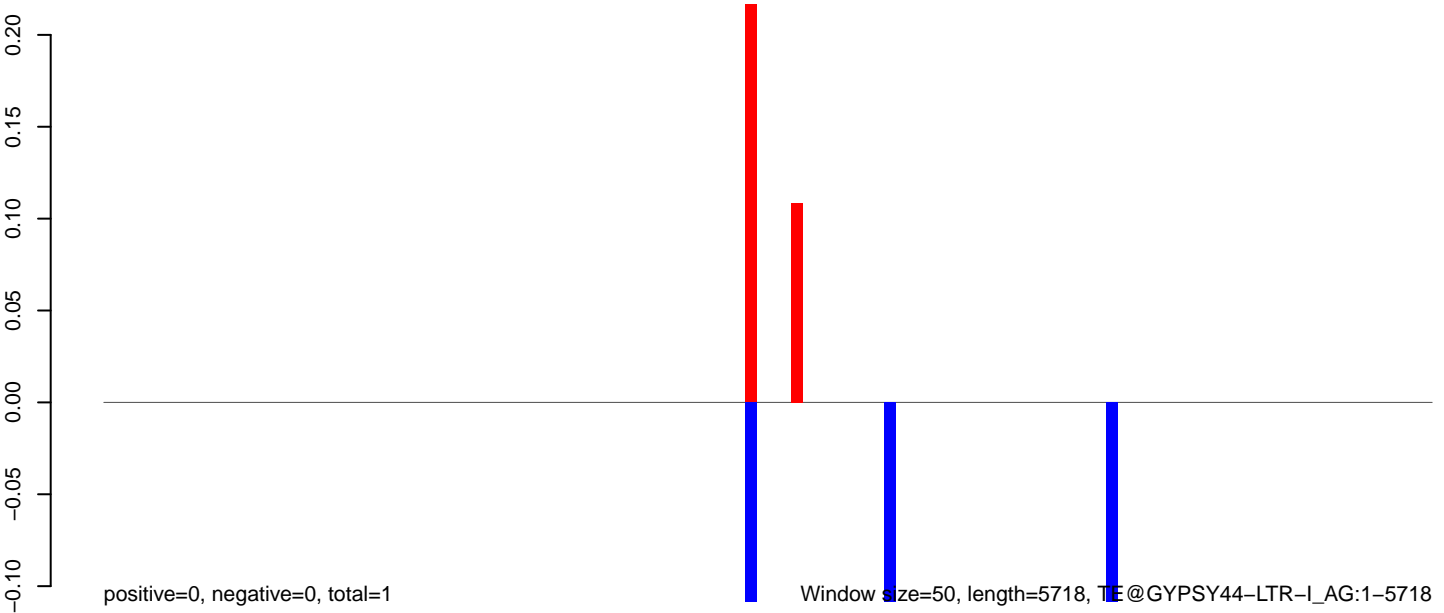
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AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

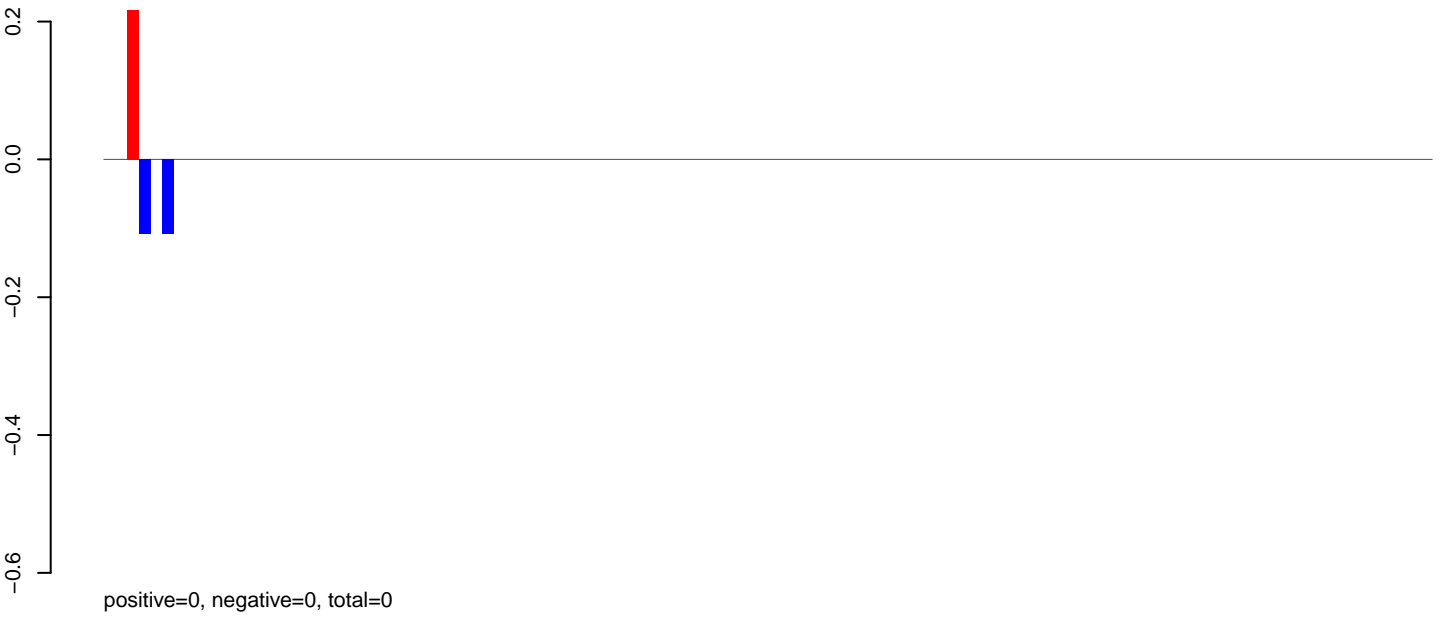


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AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

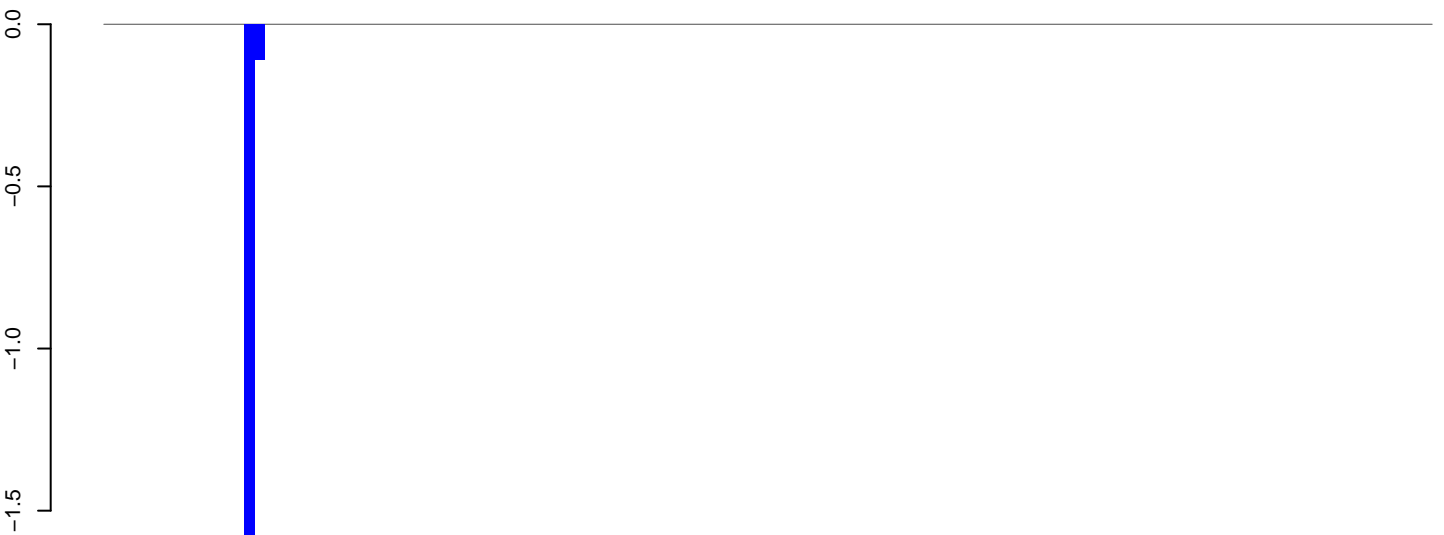


AnGam_Sua5b_cells.18_23.rep



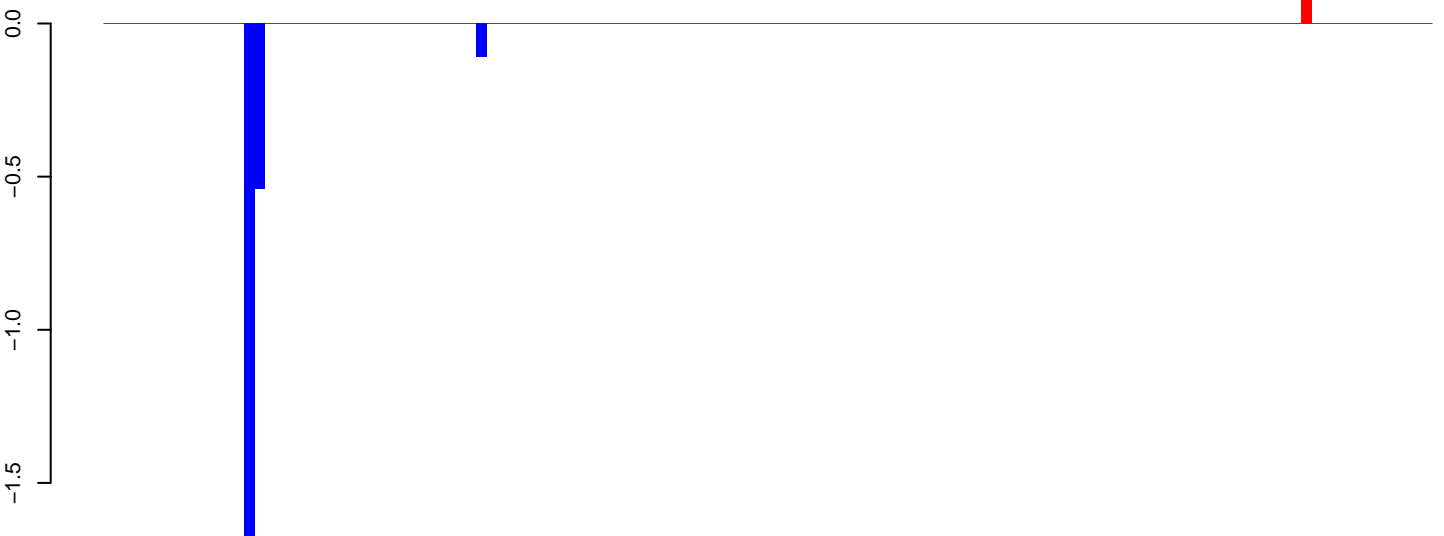
positive=0, negative=1, total=1

AnGam_Sua5b_cells.24_35.rep



positive=0, negative=2, total=2

AnGam_Sua5b_cells.rep

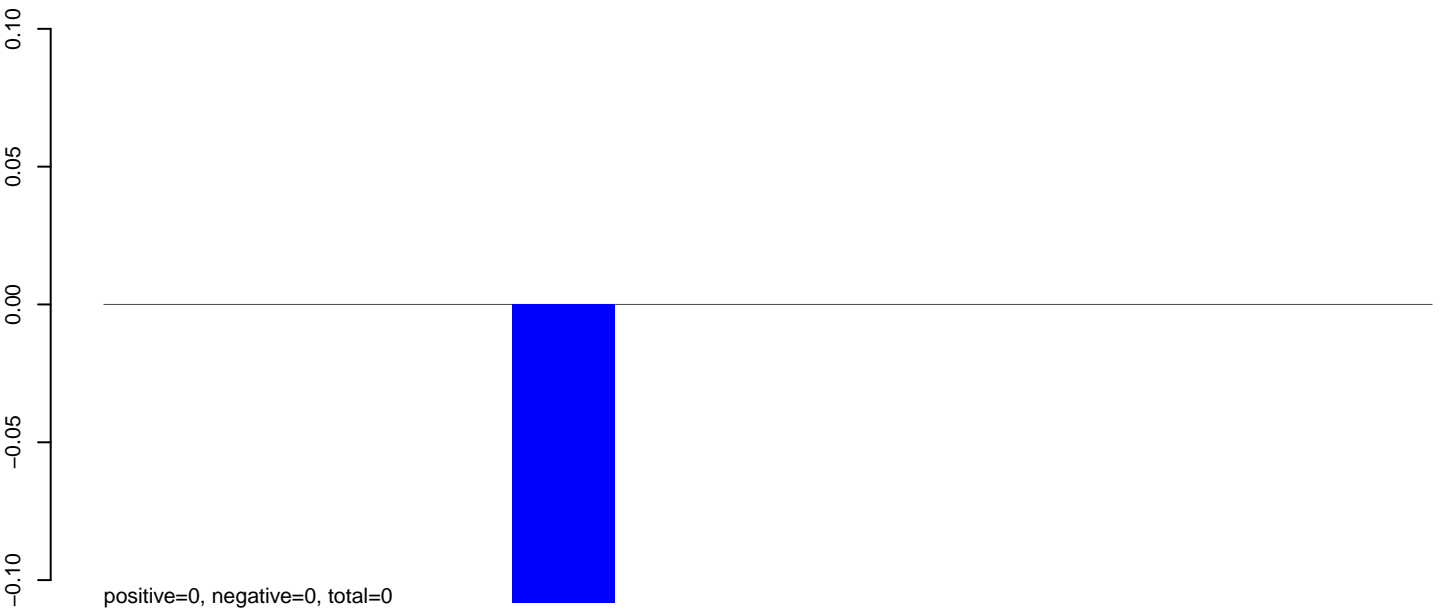


positive=0, negative=2, total=3

Window size=50, length=6641, TE@GYPSY16-LTR-L_AG:1-6641

0 1000 2000 3000 4000 5000 6000

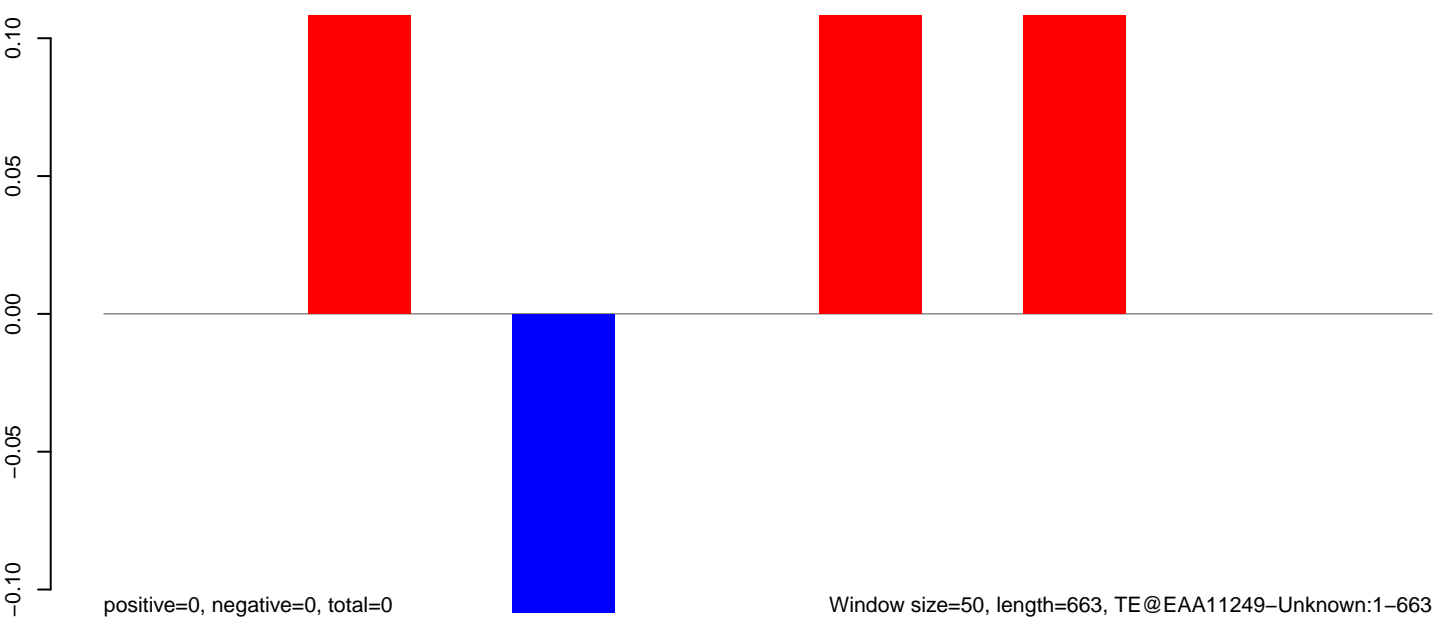
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

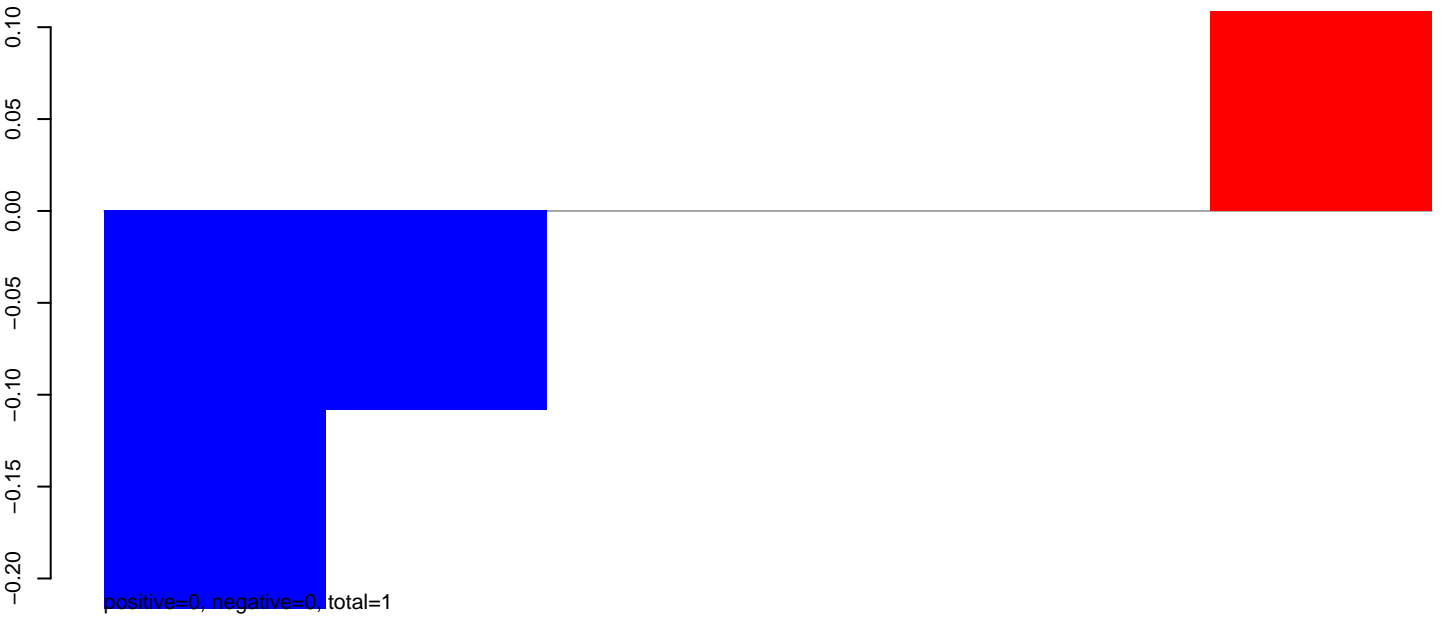


AnGam_Sua5b_cells.rep

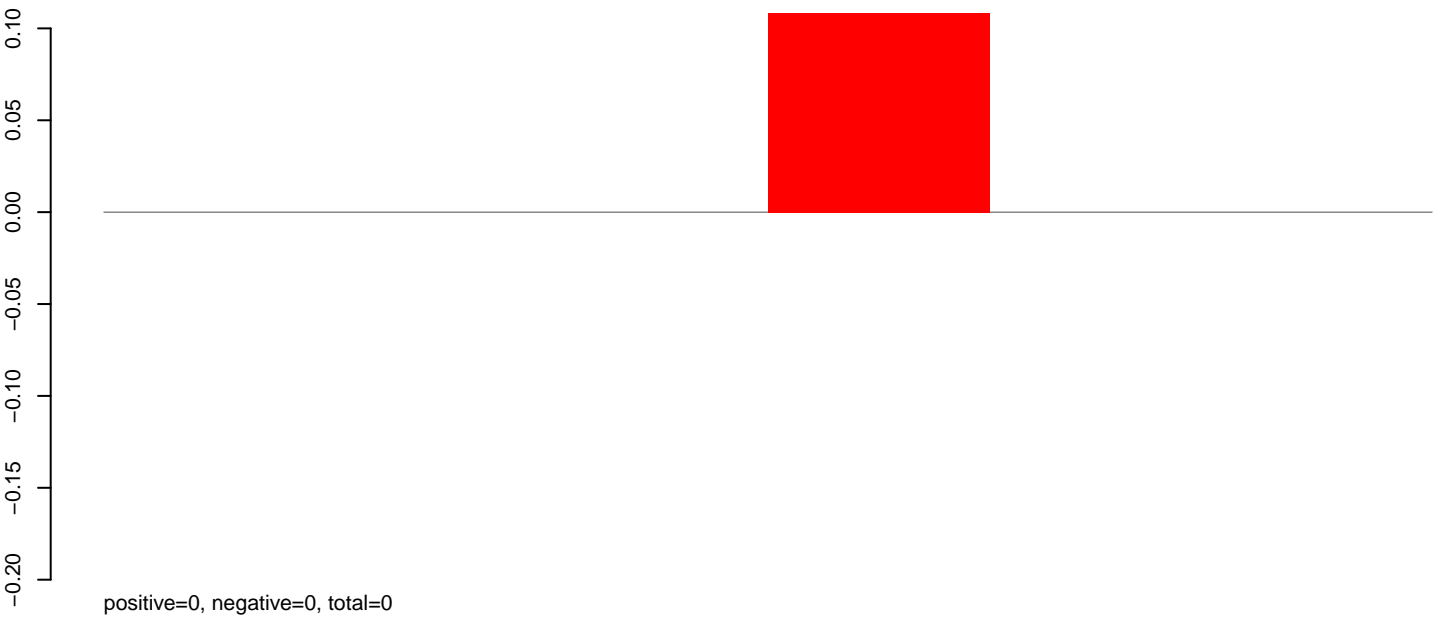


100 200 300 400 500 600 700

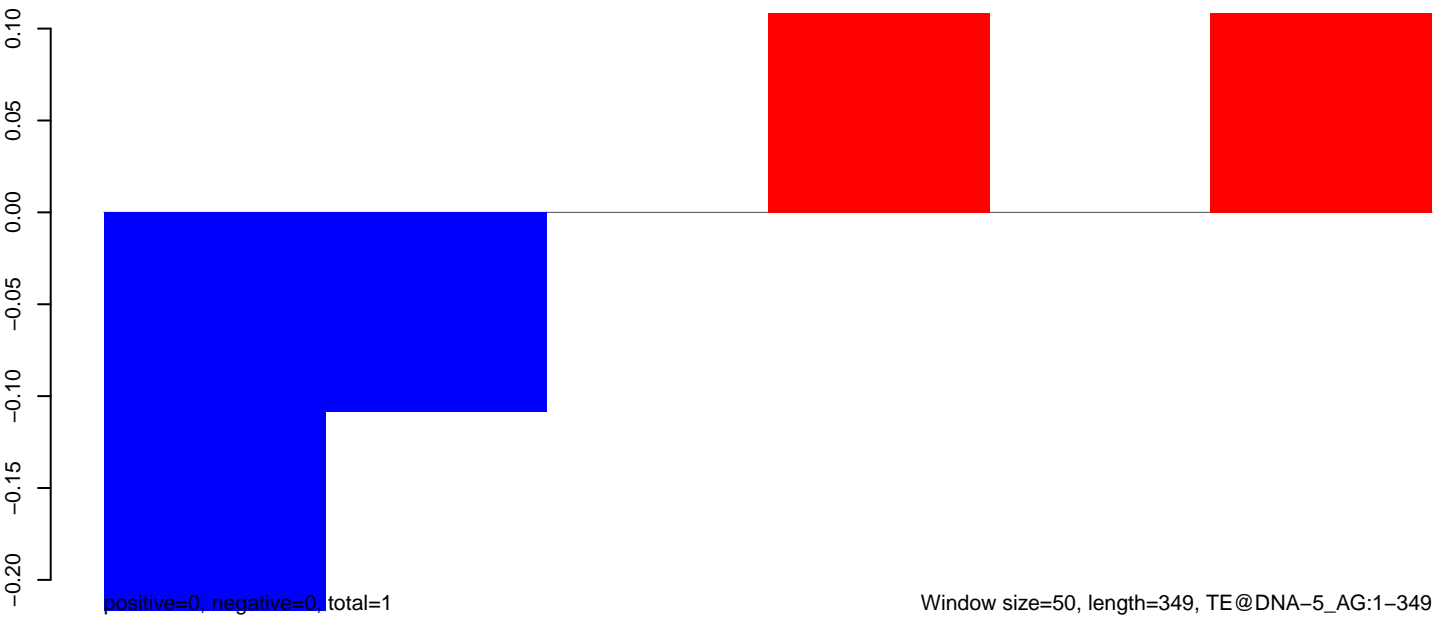
AnGam_Sua5b_cells.18_23.rep



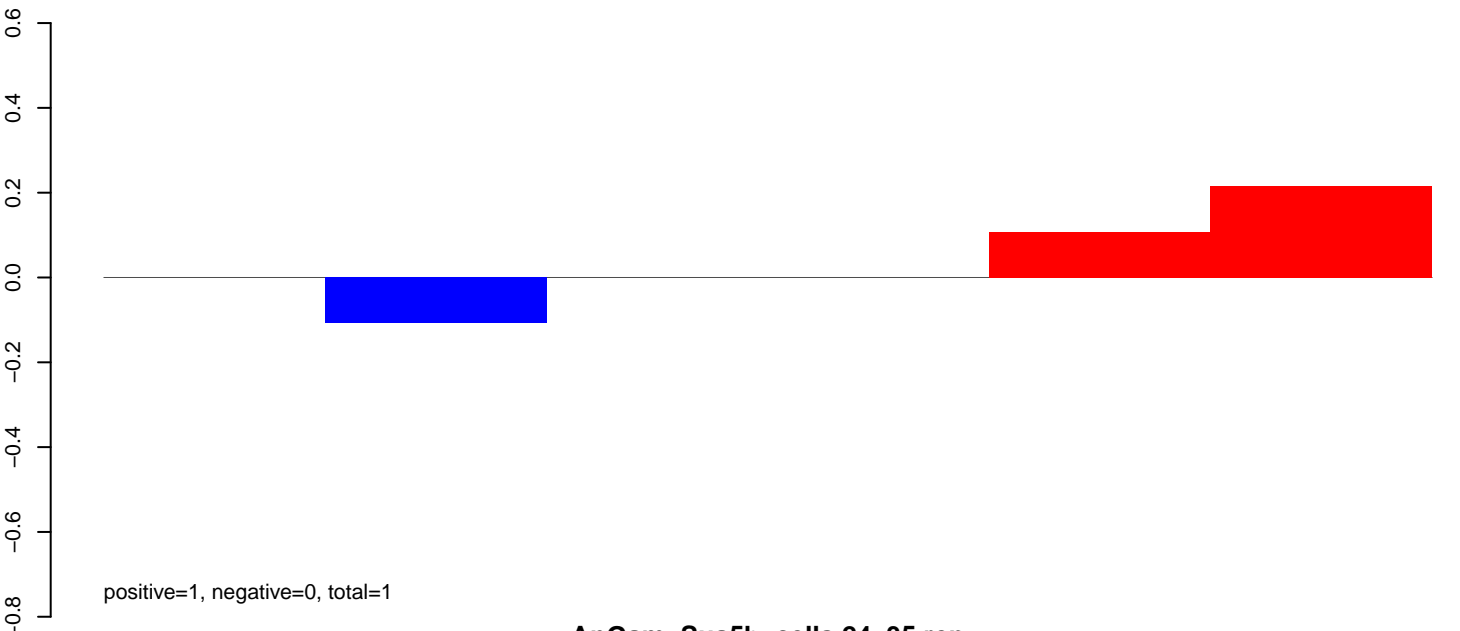
AnGam_Sua5b_cells.24_35.rep



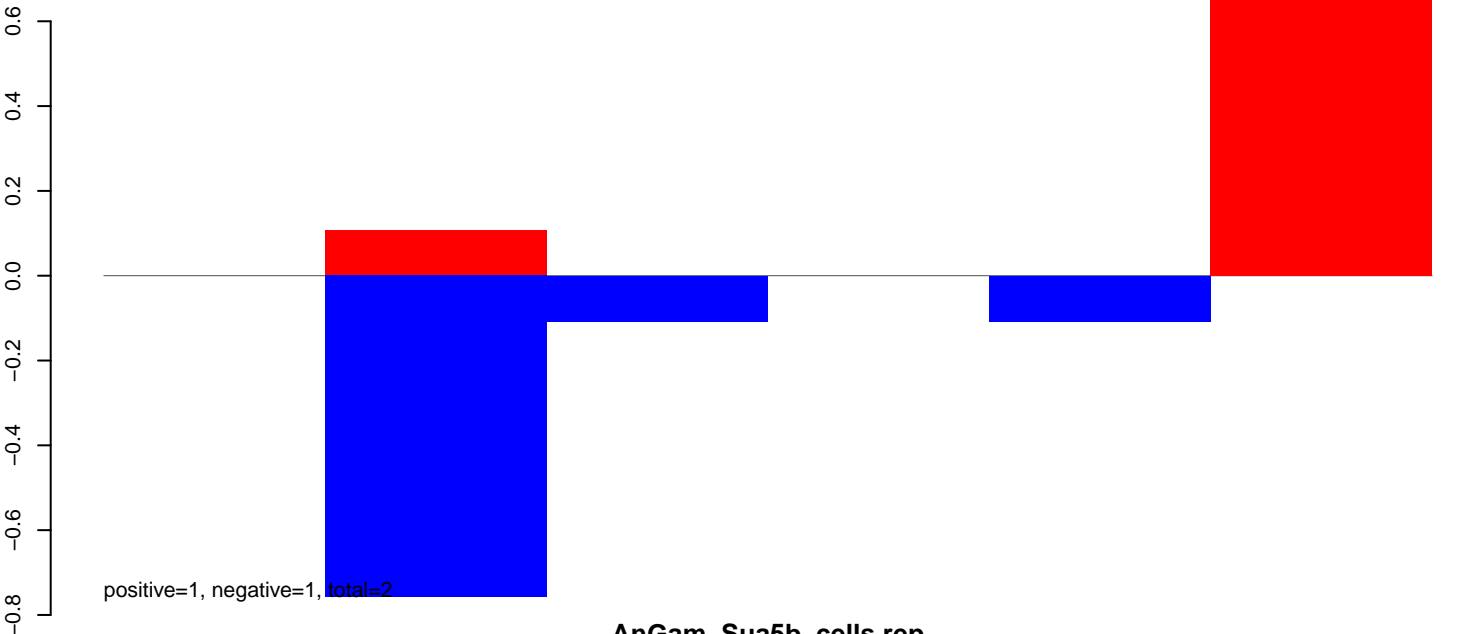
AnGam_Sua5b_cells.rep



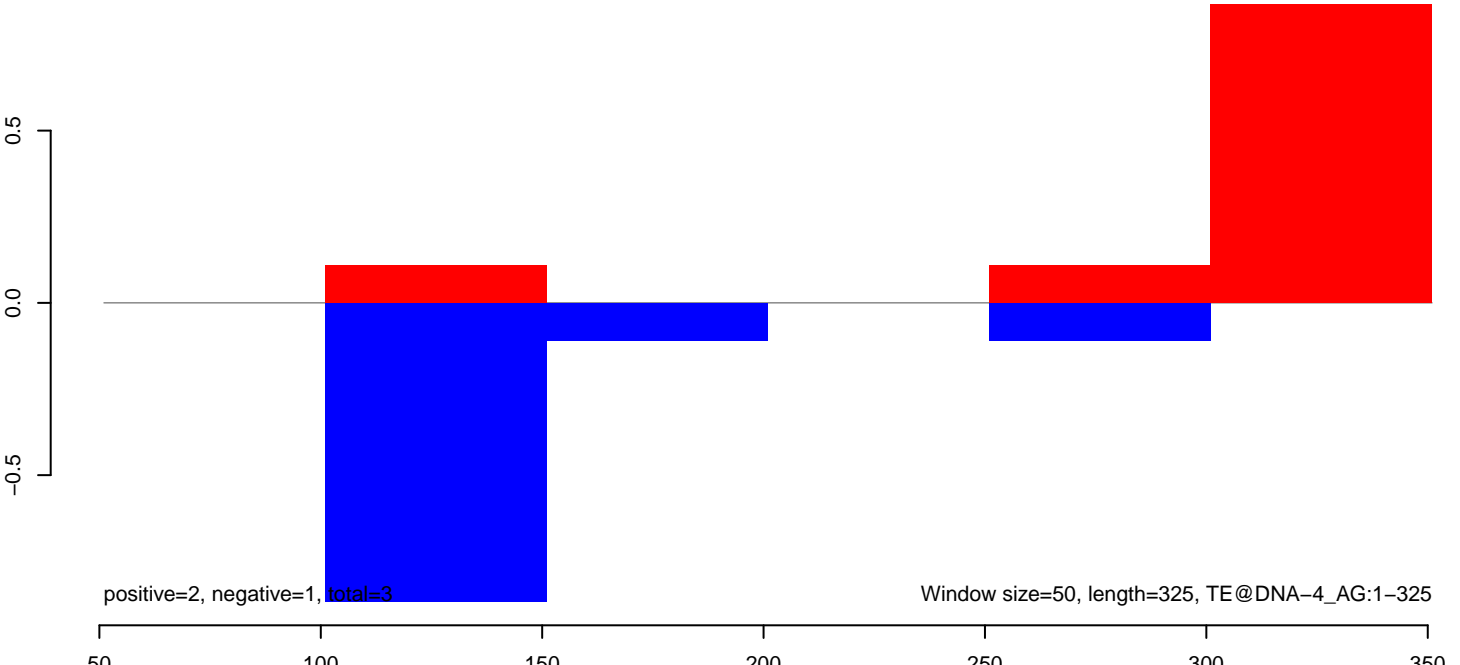
AnGam_Sua5b_cells.18_23.rep



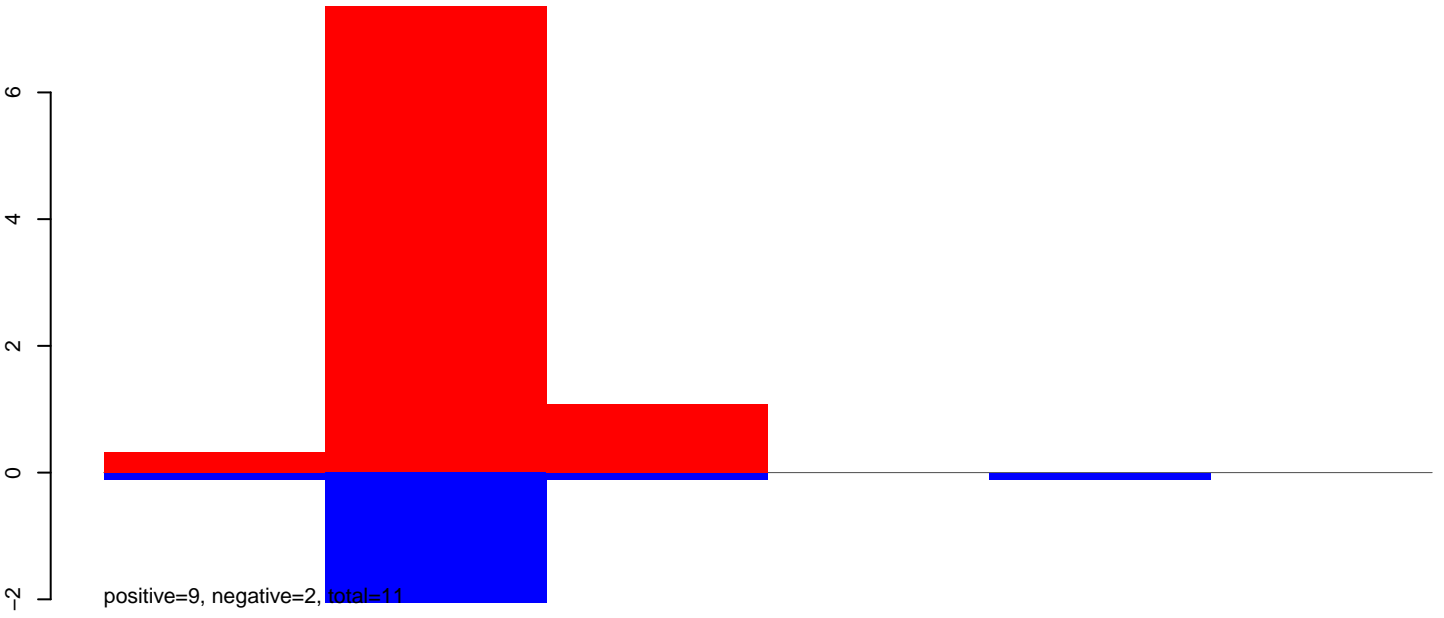
AnGam_Sua5b_cells.24_35.rep



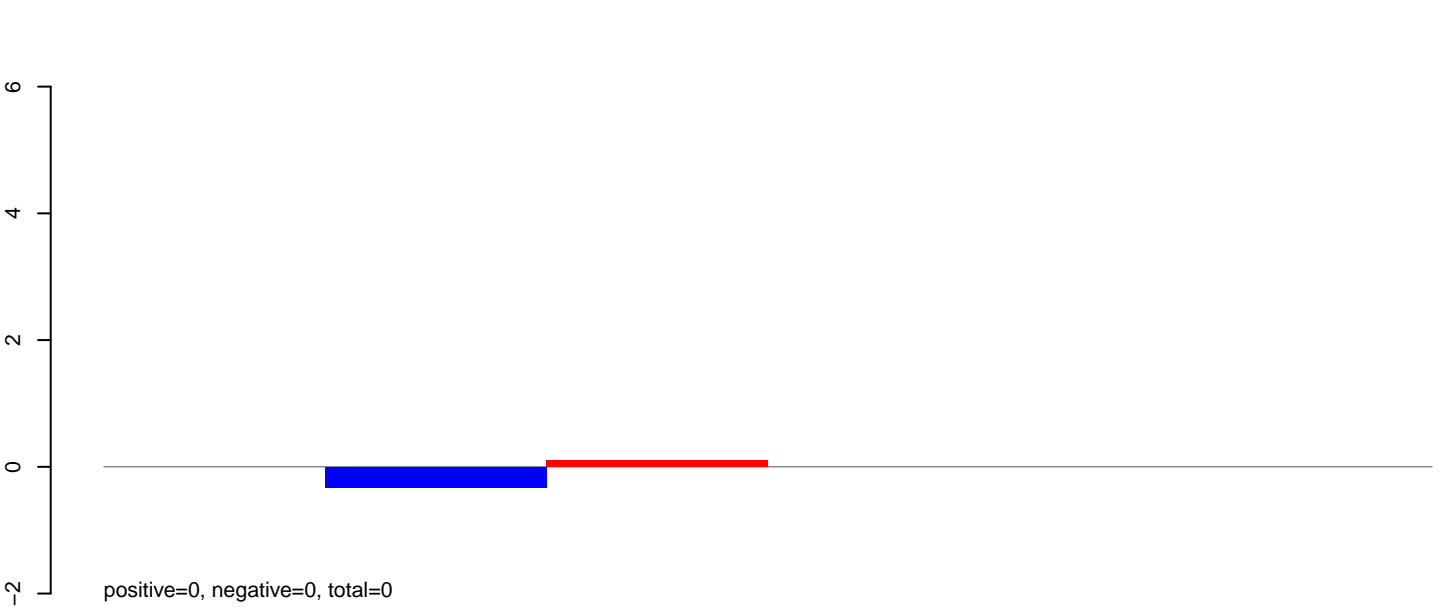
AnGam_Sua5b_cells.rep



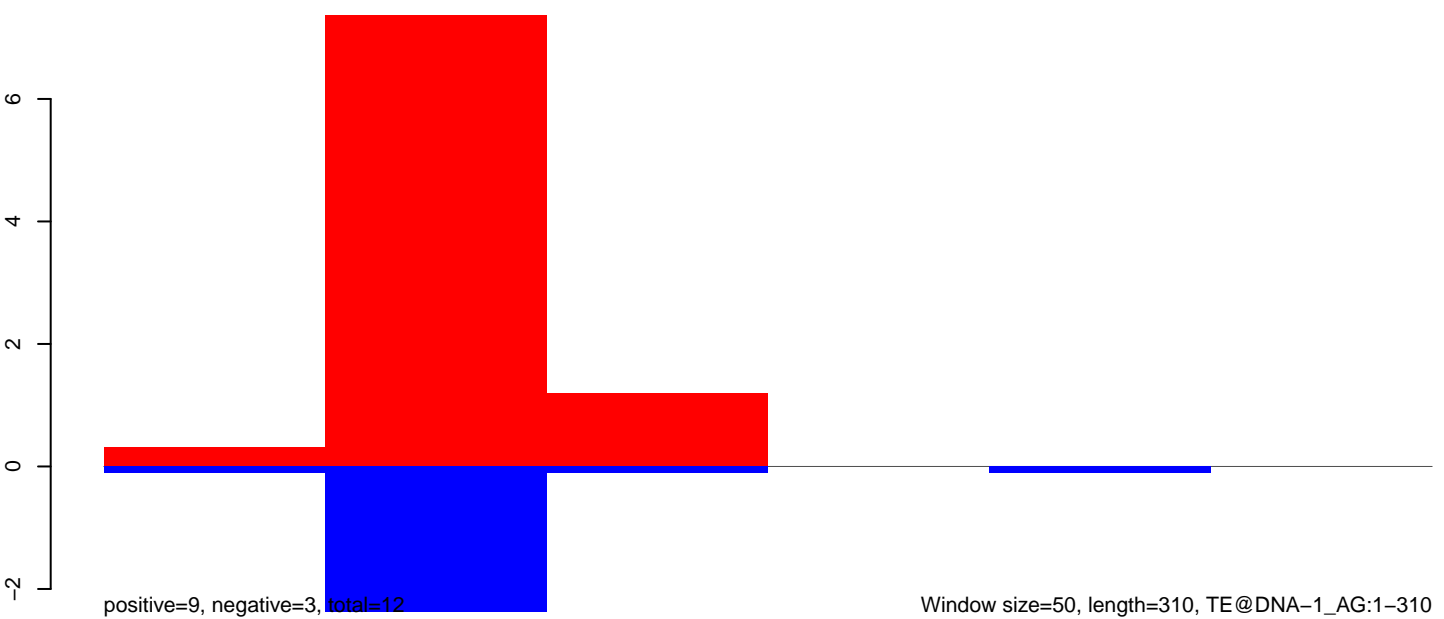
AnGam_Sua5b_cells.18_23.rep



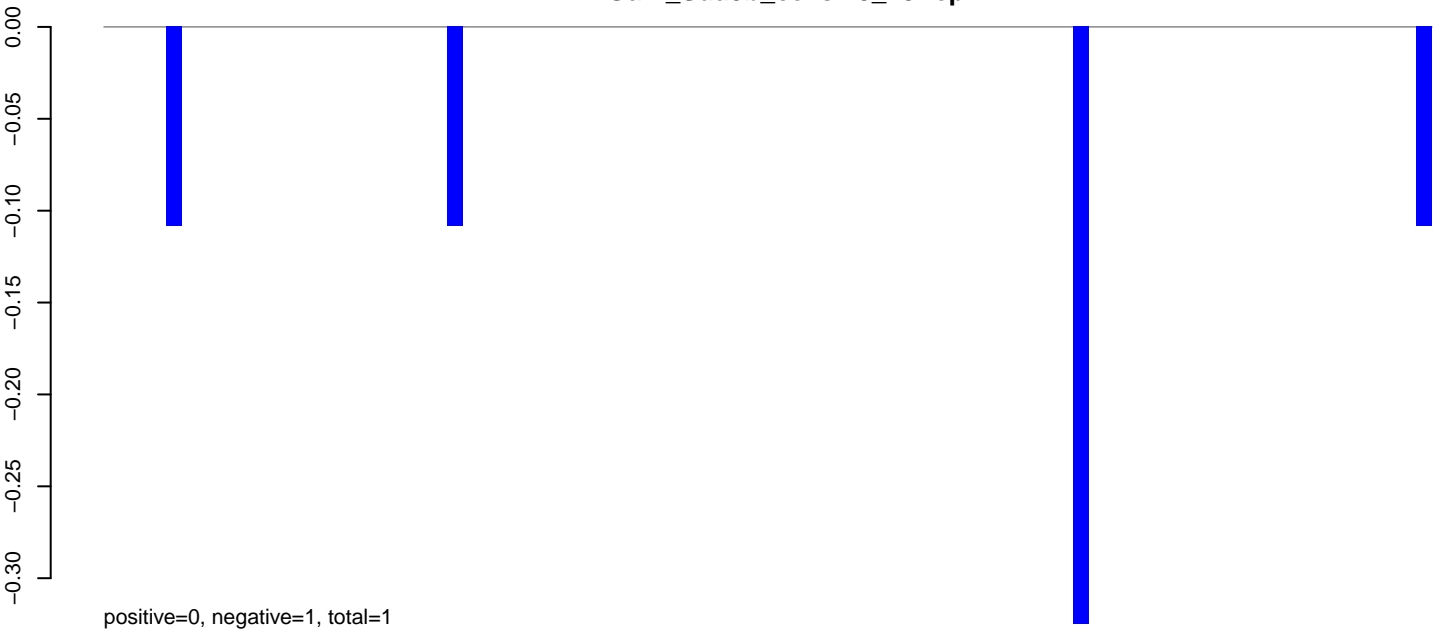
AnGam_Sua5b_cells.24_35.rep



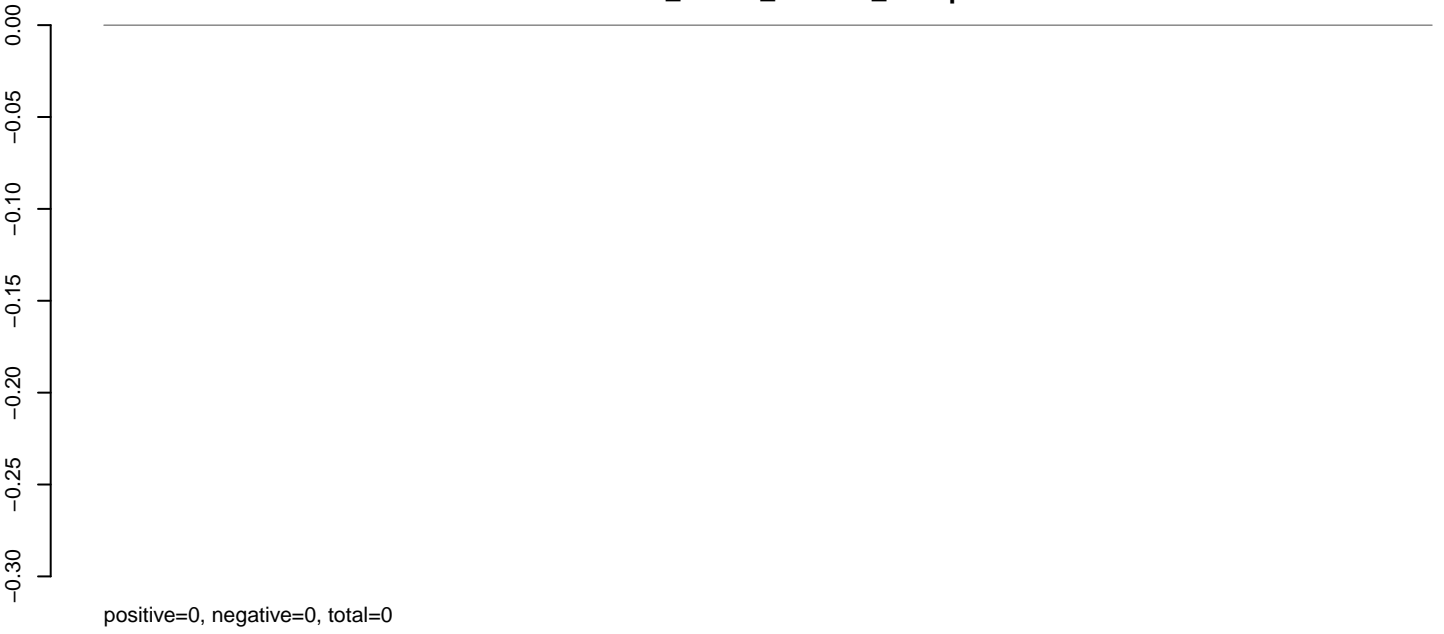
AnGam_Sua5b_cells.rep



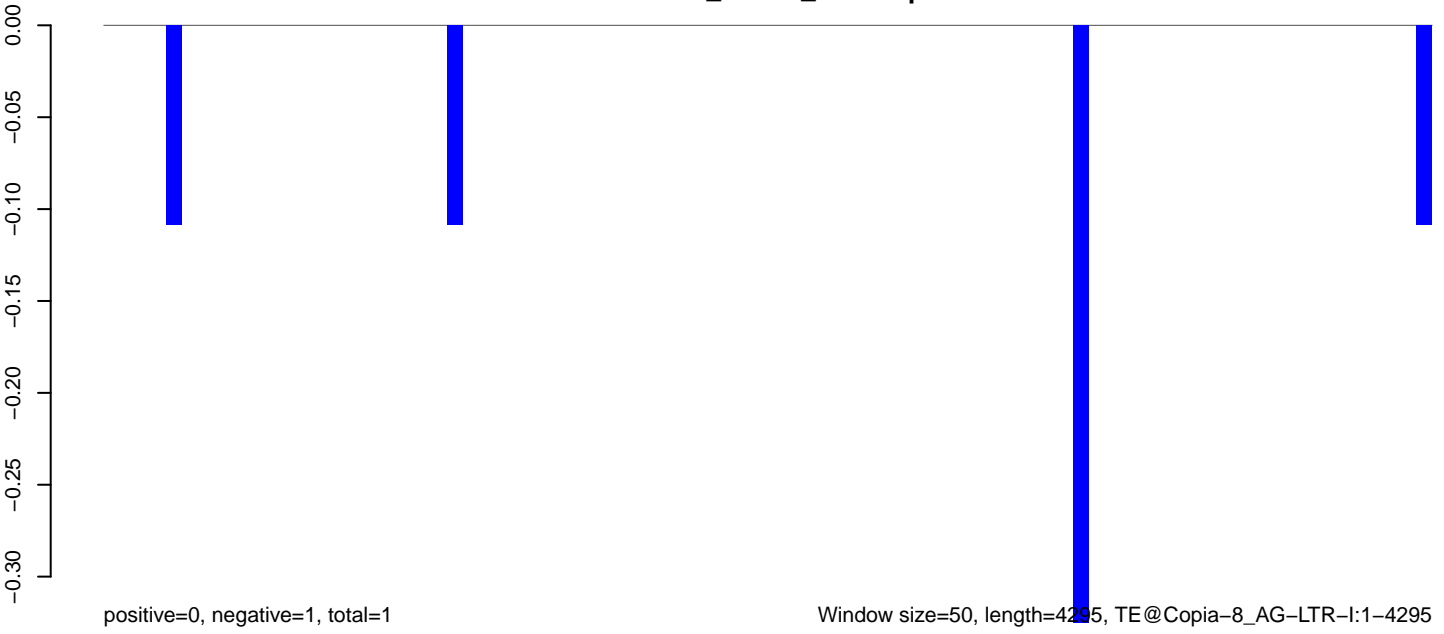
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

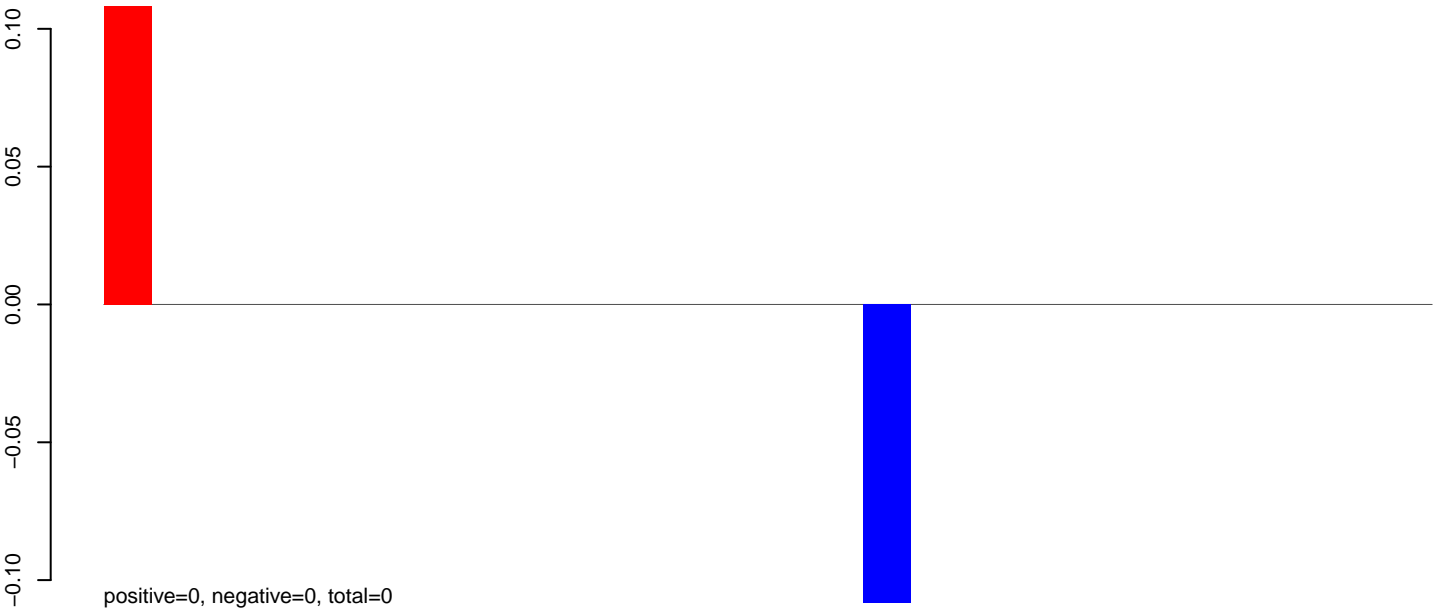


AnGam_Sua5b_cells.rep

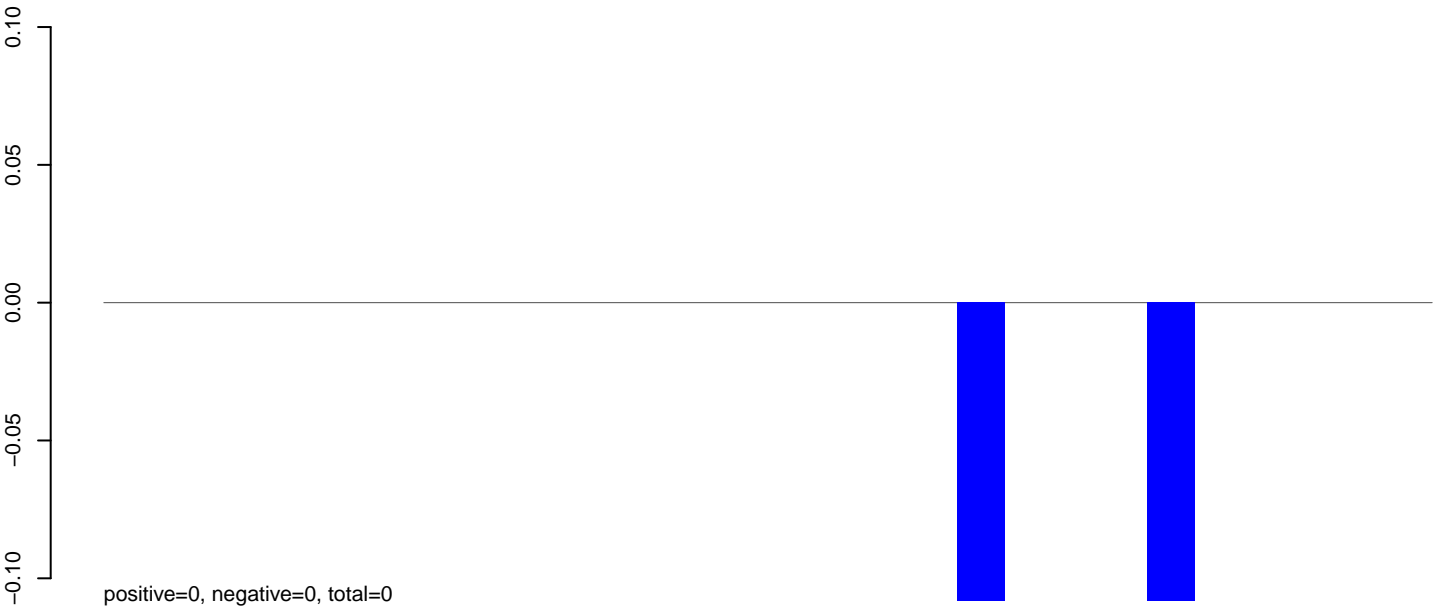


0 1000 2000 3000 4000

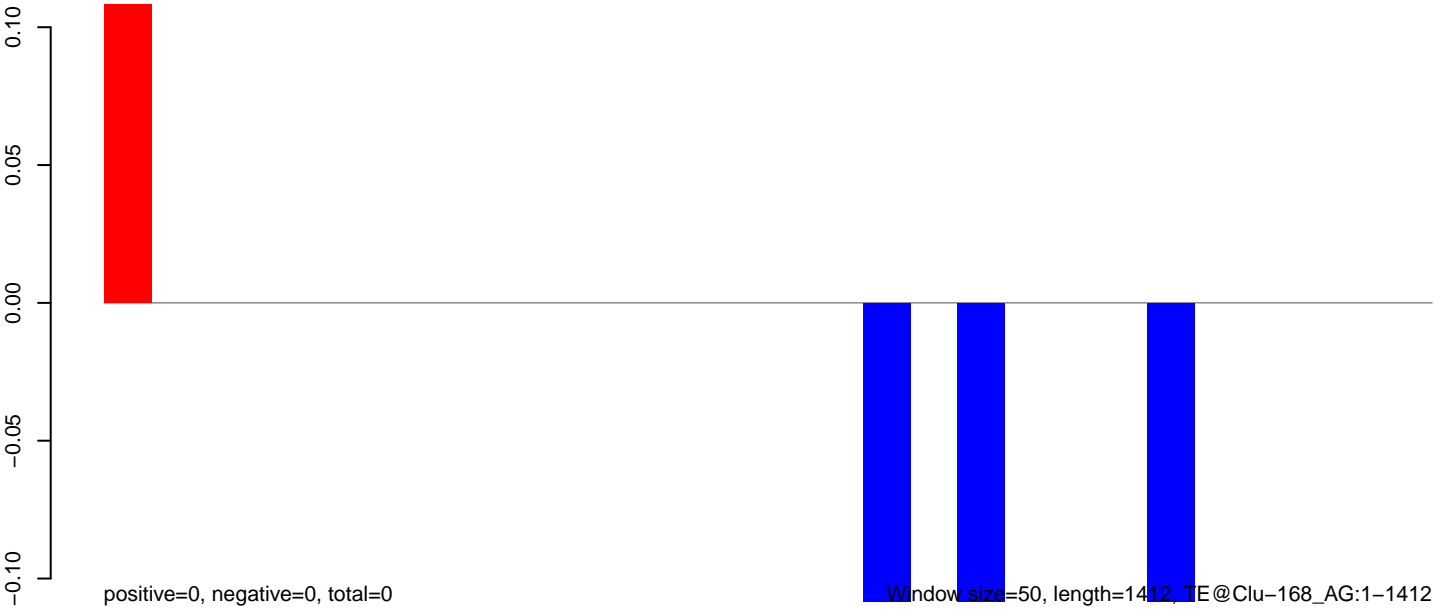
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



0 200 400 600 800 1000 1200 1400

AnGam_Sua5b_cells.18_23.rep

0.00
-0.05
-0.10
-0.15
-0.20

positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep

0.00
-0.05
-0.10
-0.15
-0.20

positive=0, negative=1, total=1

AnGam_Sua5b_cells.rep

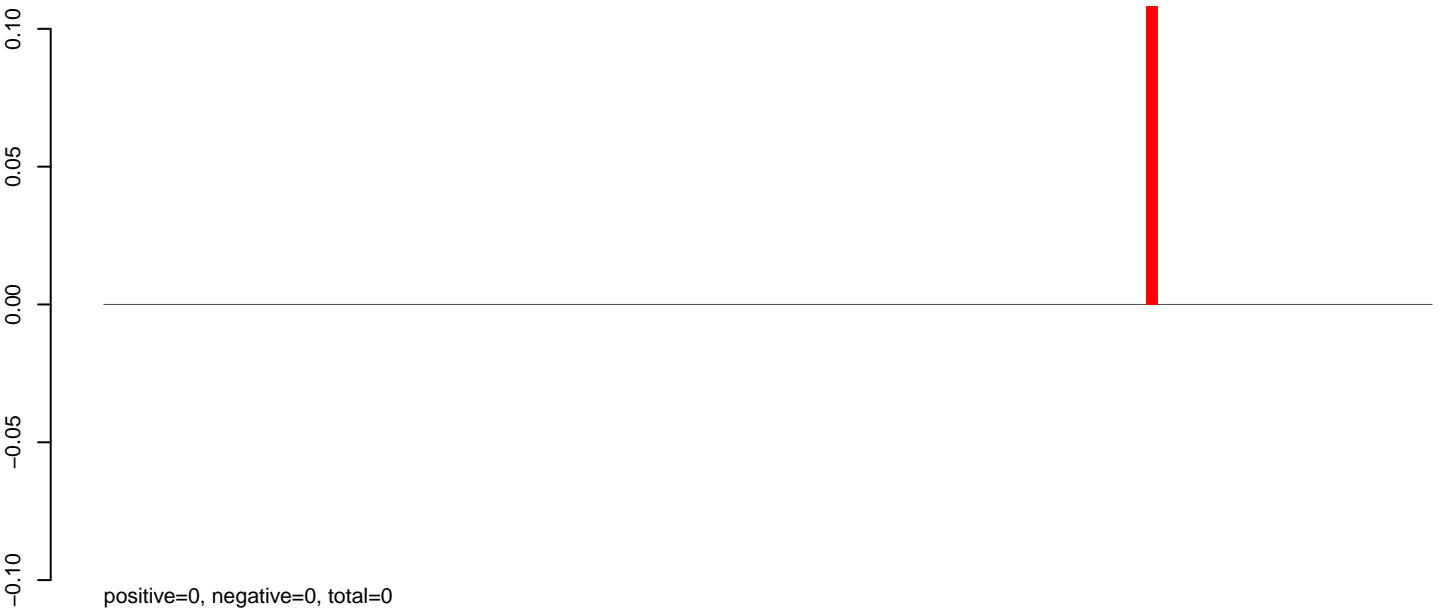
0.00
-0.05
-0.10
-0.15
-0.20

positive=0, negative=1, total=1

Window size=50, length=1717, TE@COPIA3-LTR-L_AG:1-1717

0 500 1000 1500

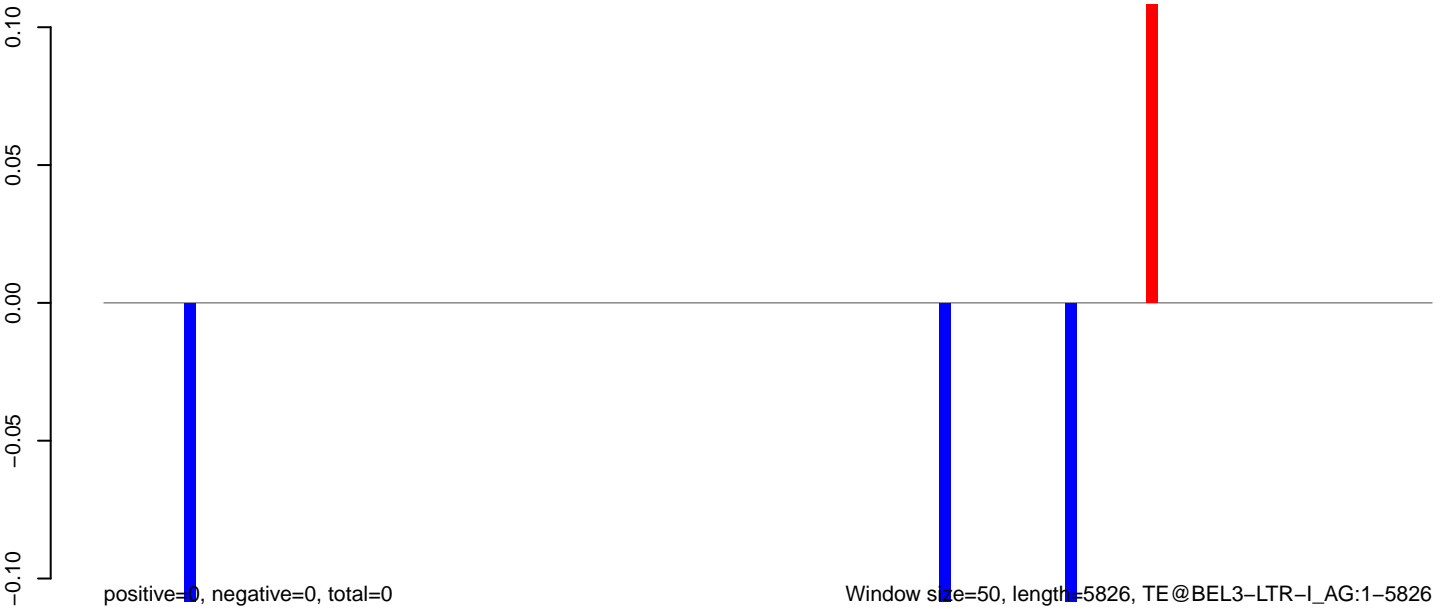
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

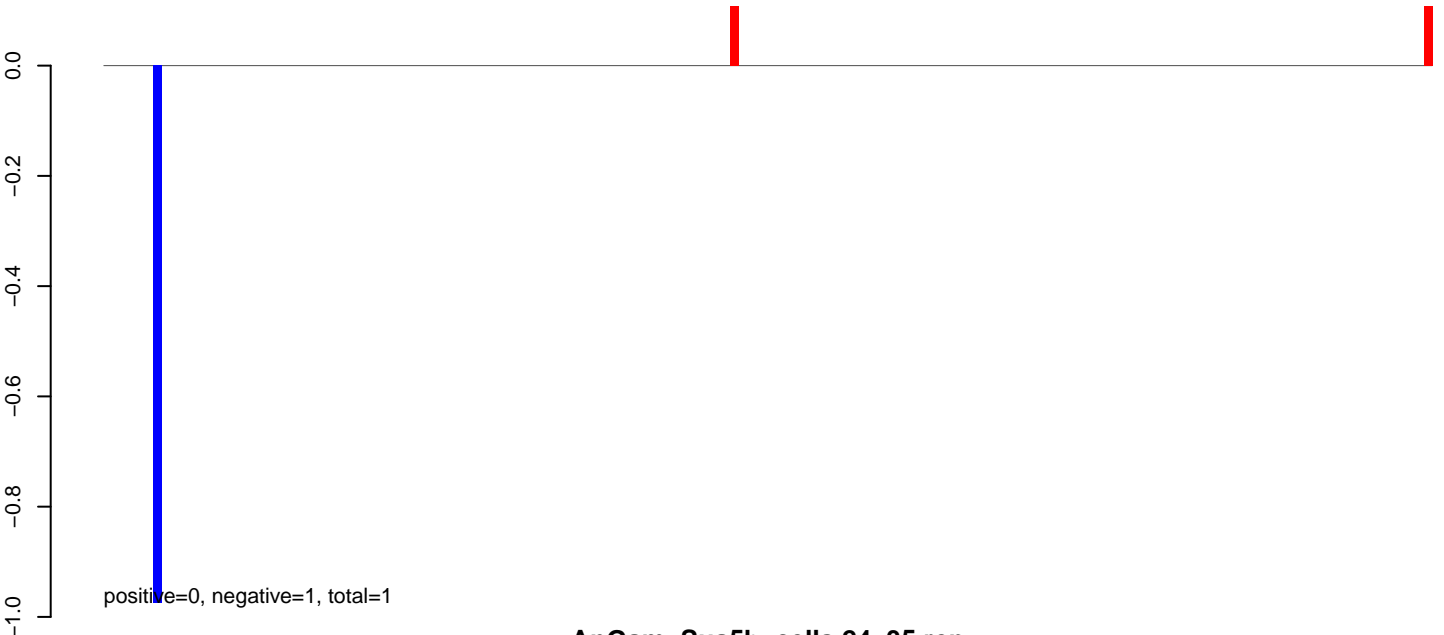


AnGam_Sua5b_cells.rep

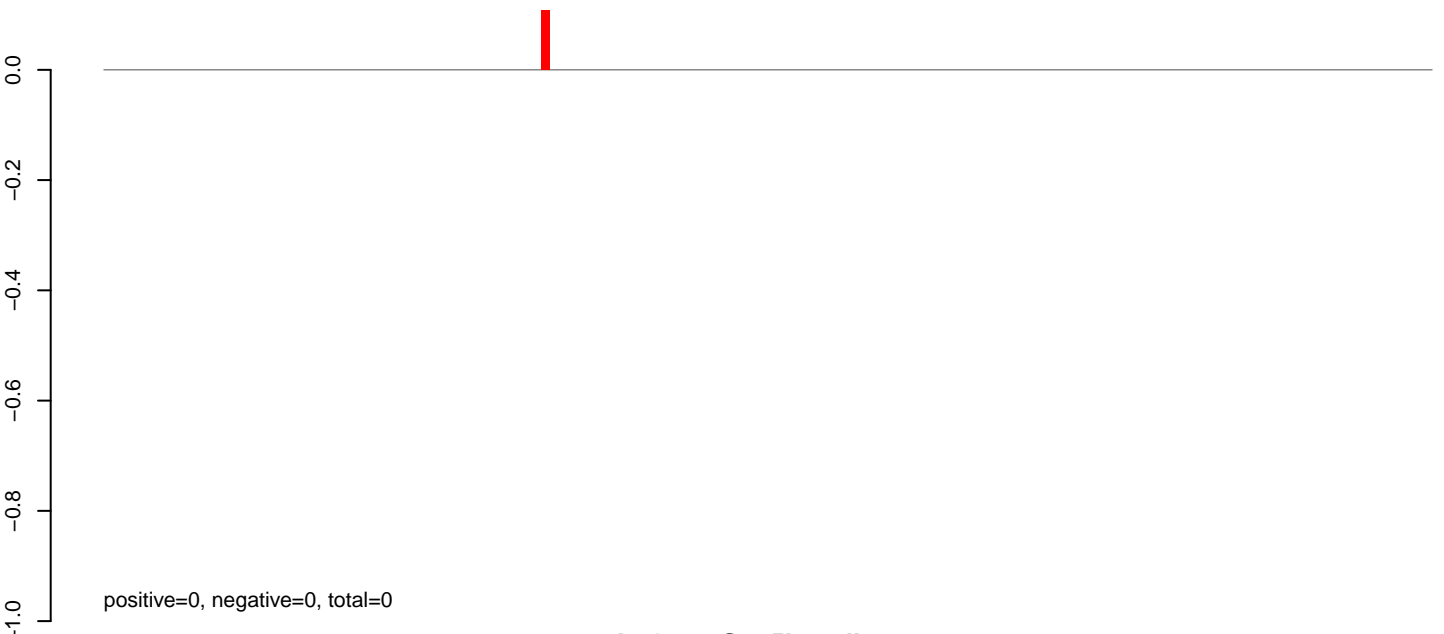


0 1000 2000 3000 4000 5000 6000

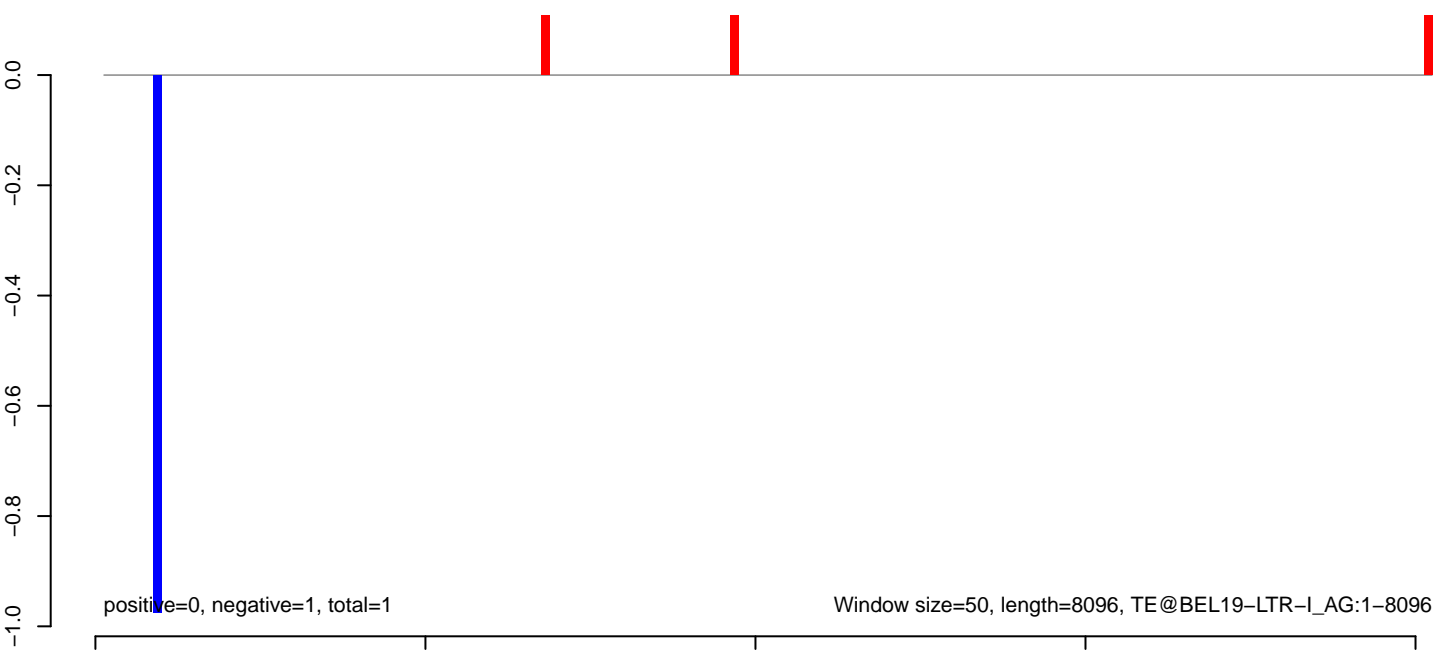
AnGam_Sua5b_cells.18_23.rep



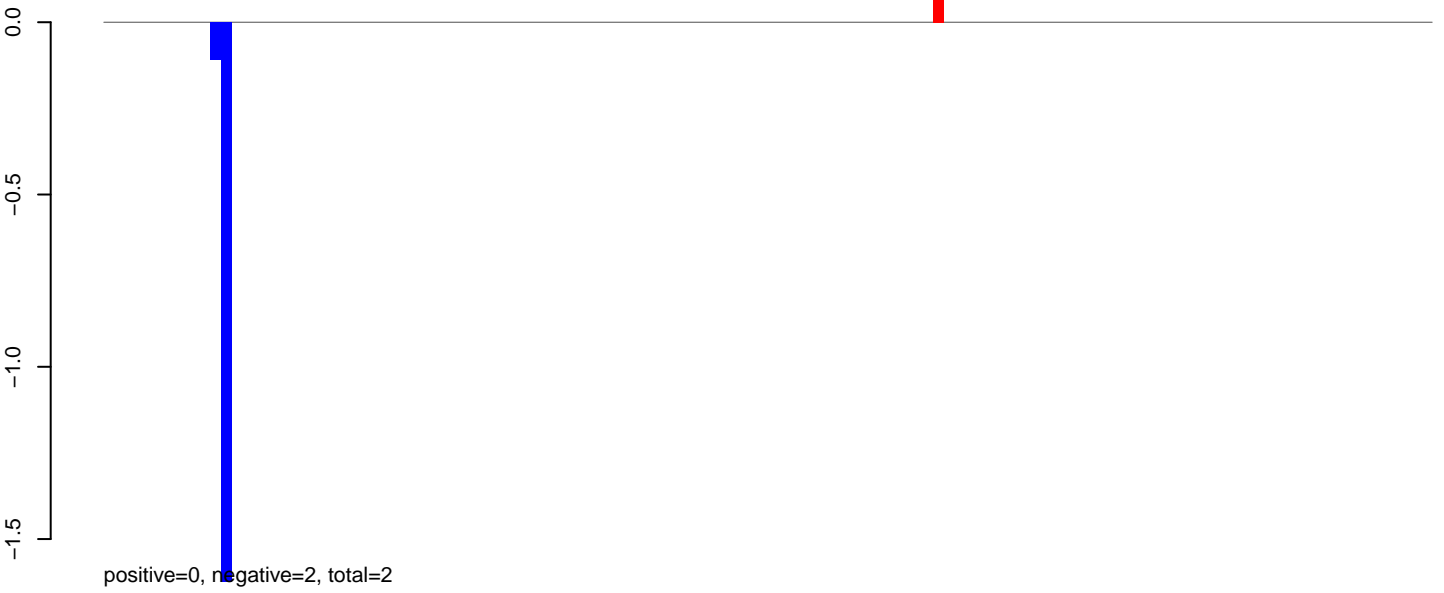
AnGam_Sua5b_cells.24_35.rep



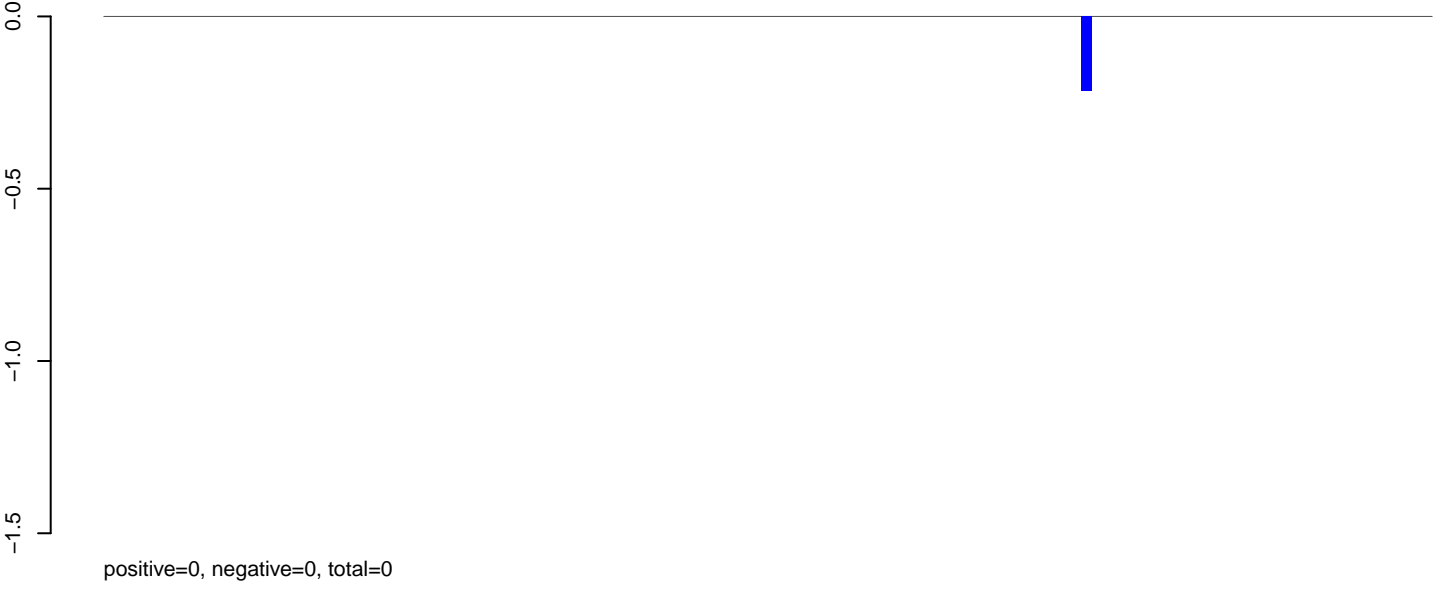
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



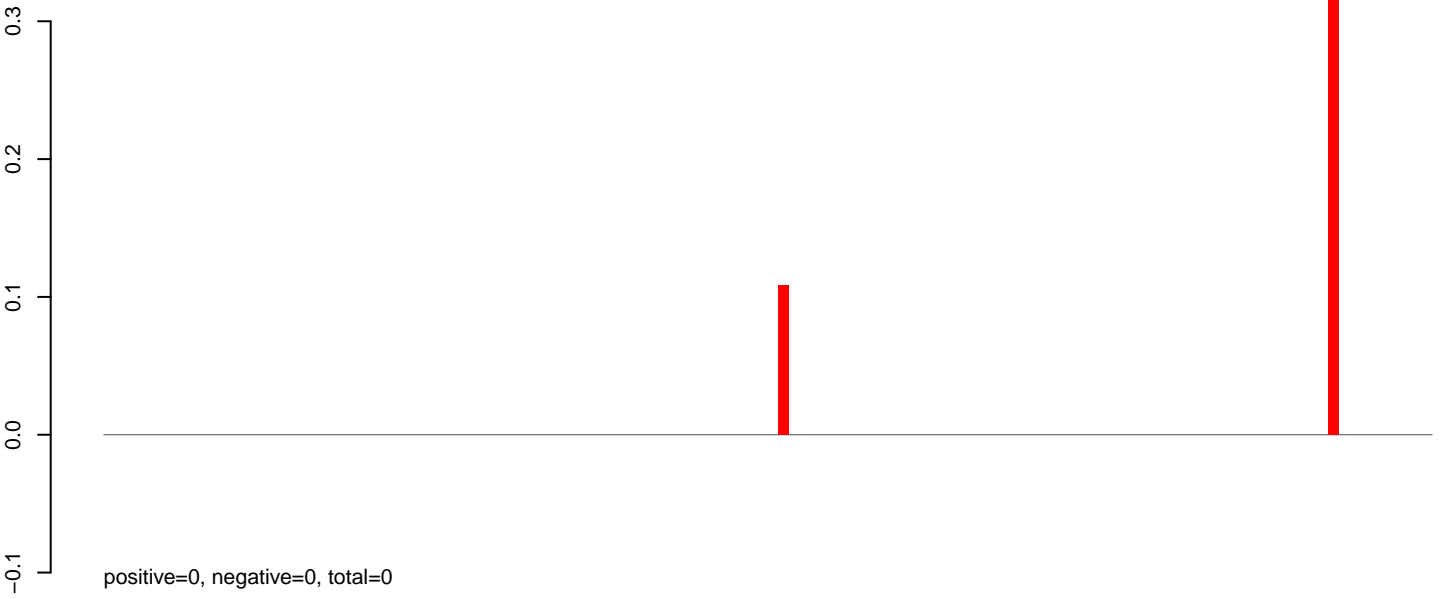
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



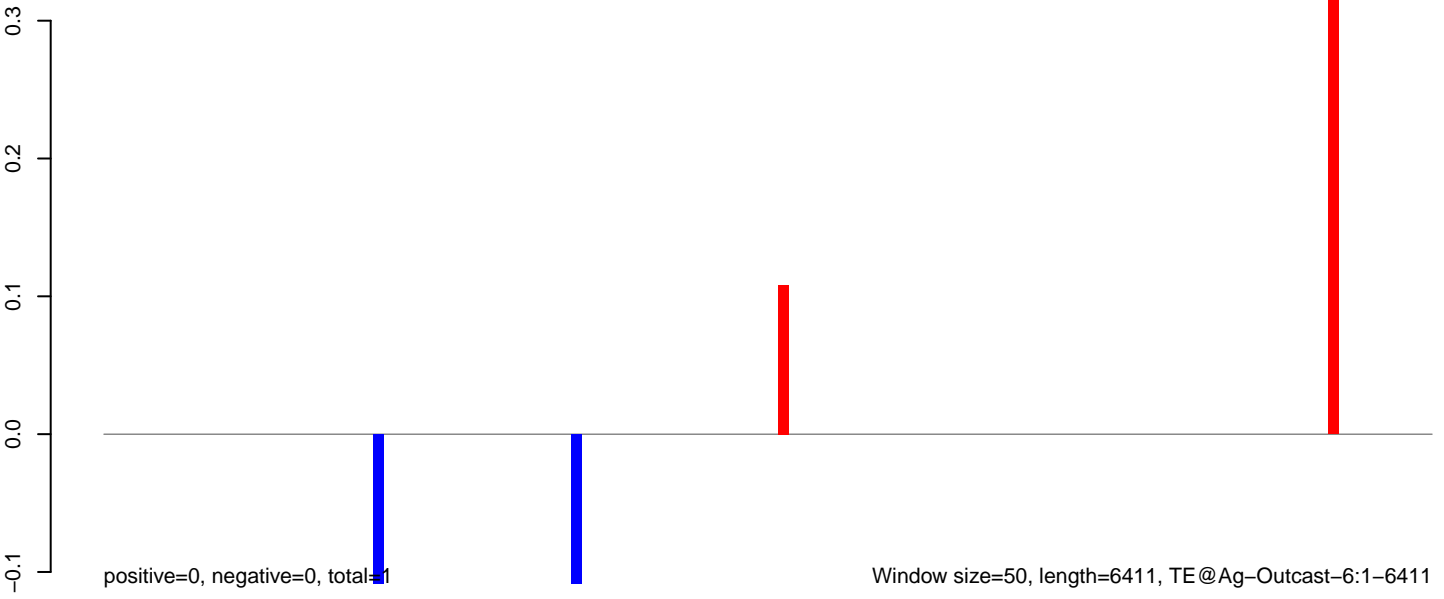
AnGam_Sua5b_cells.18_23.rep



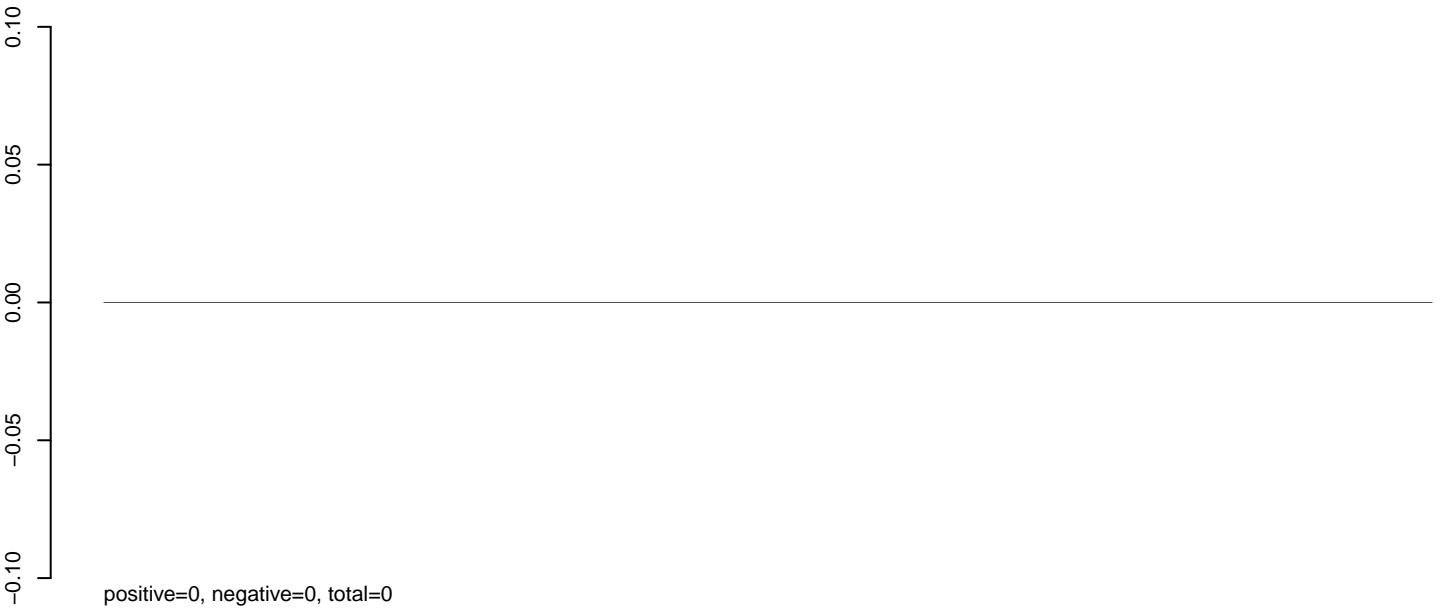
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



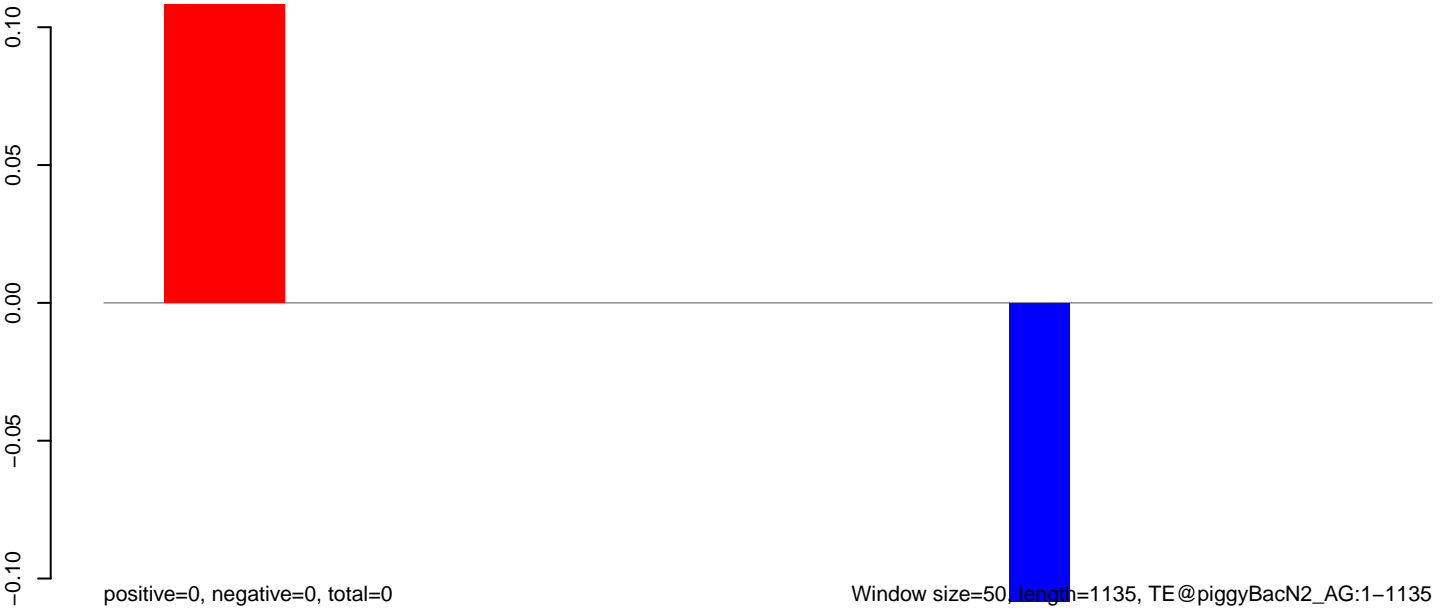
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



200

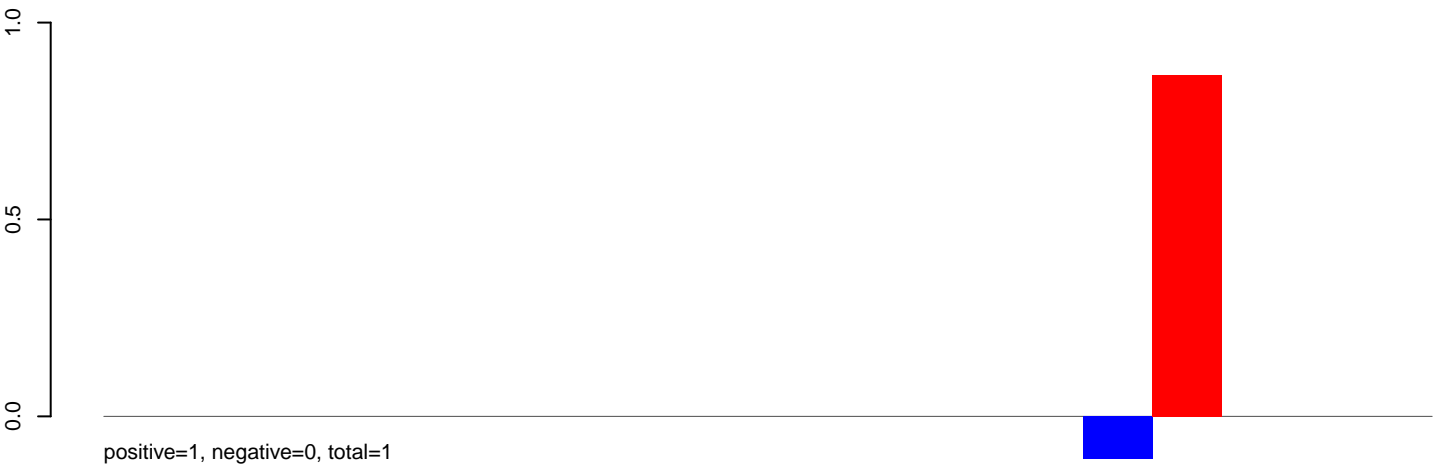
400

600

800

1000

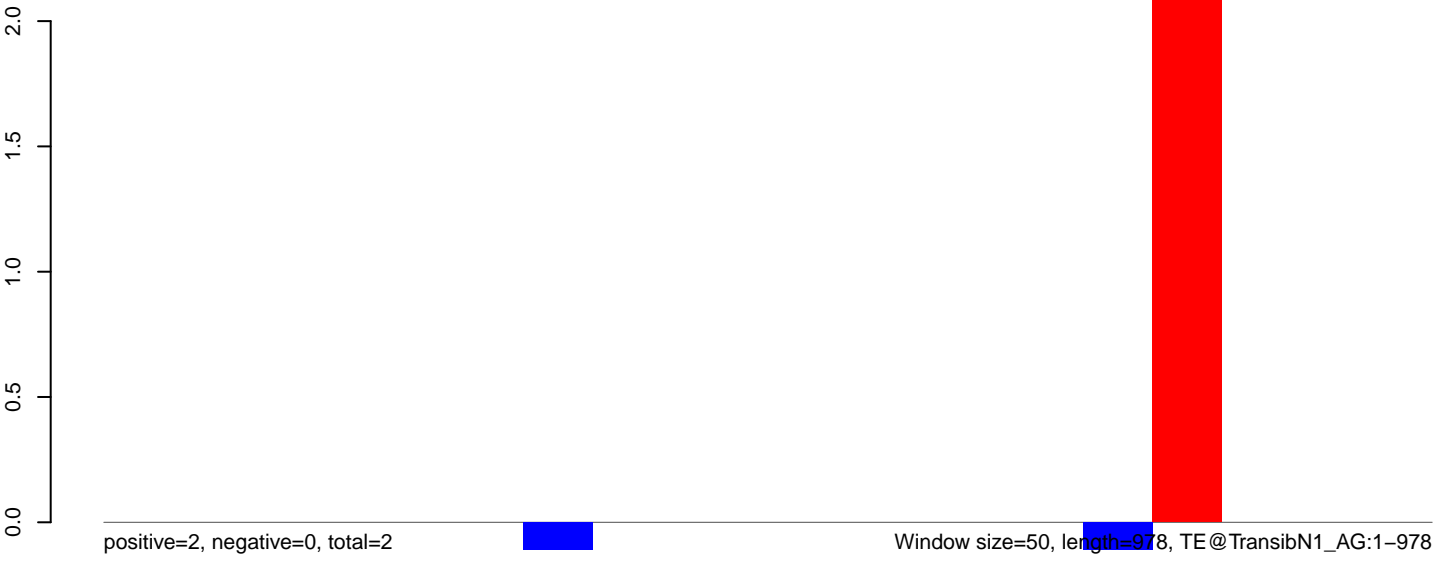
AnGam_Sua5b_cells.18_23.rep



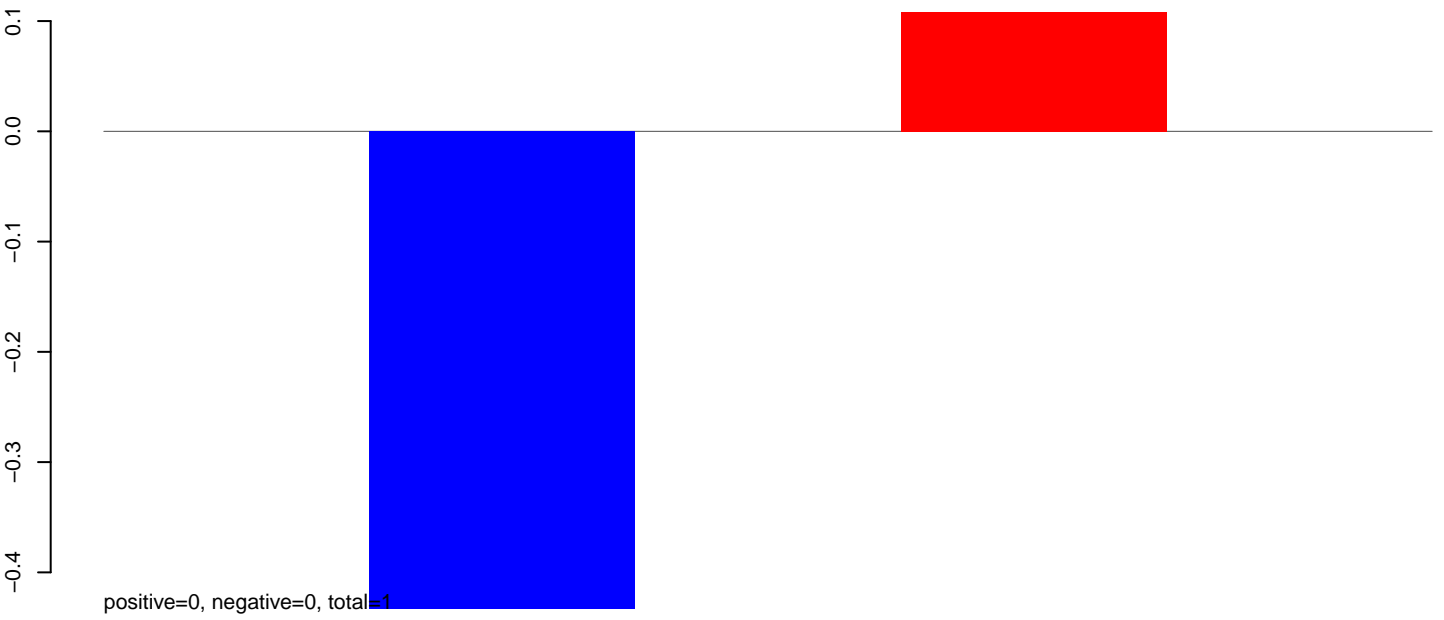
AnGam_Sua5b_cells.24_35.rep



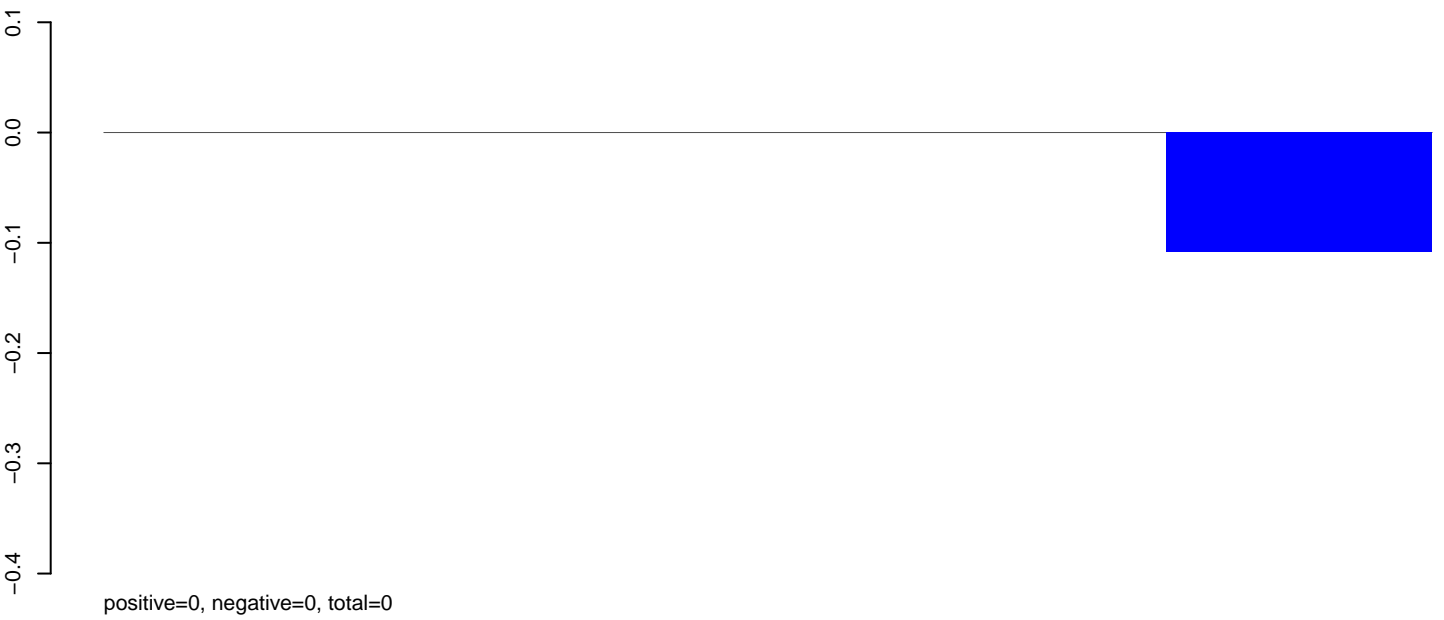
AnGam_Sua5b_cells.rep



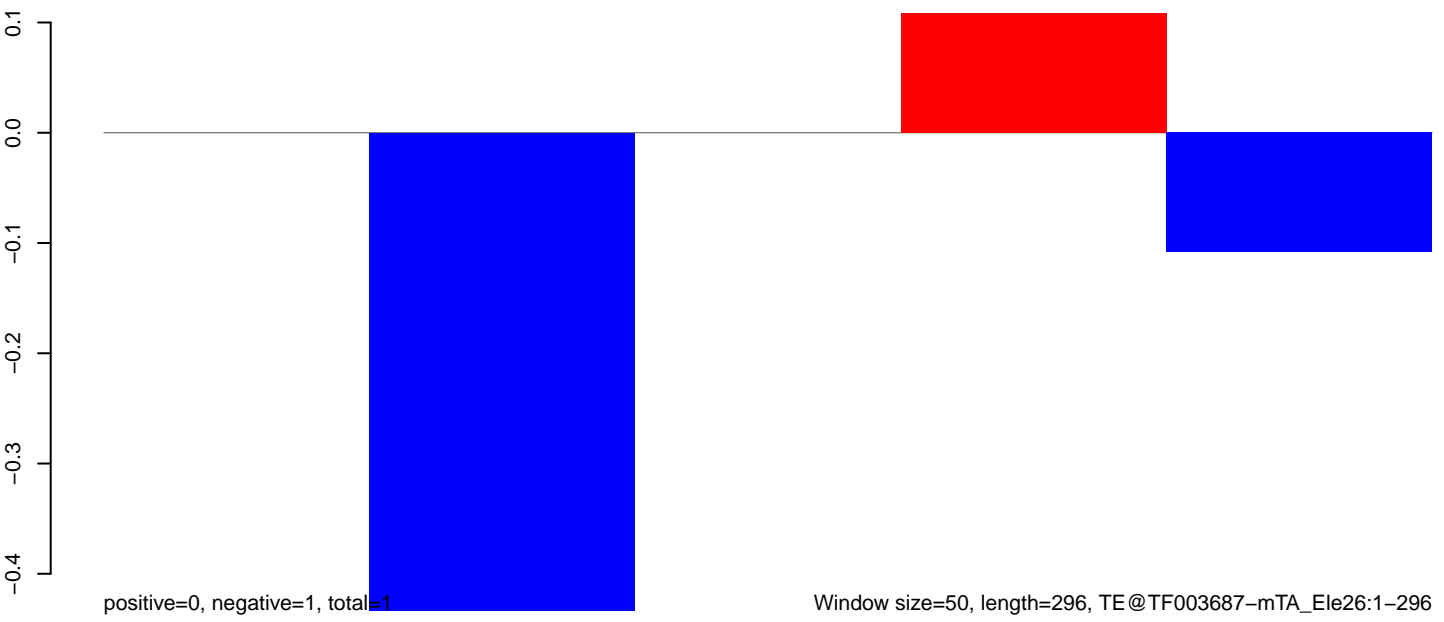
AnGam_Sua5b_cells.18_23.rep



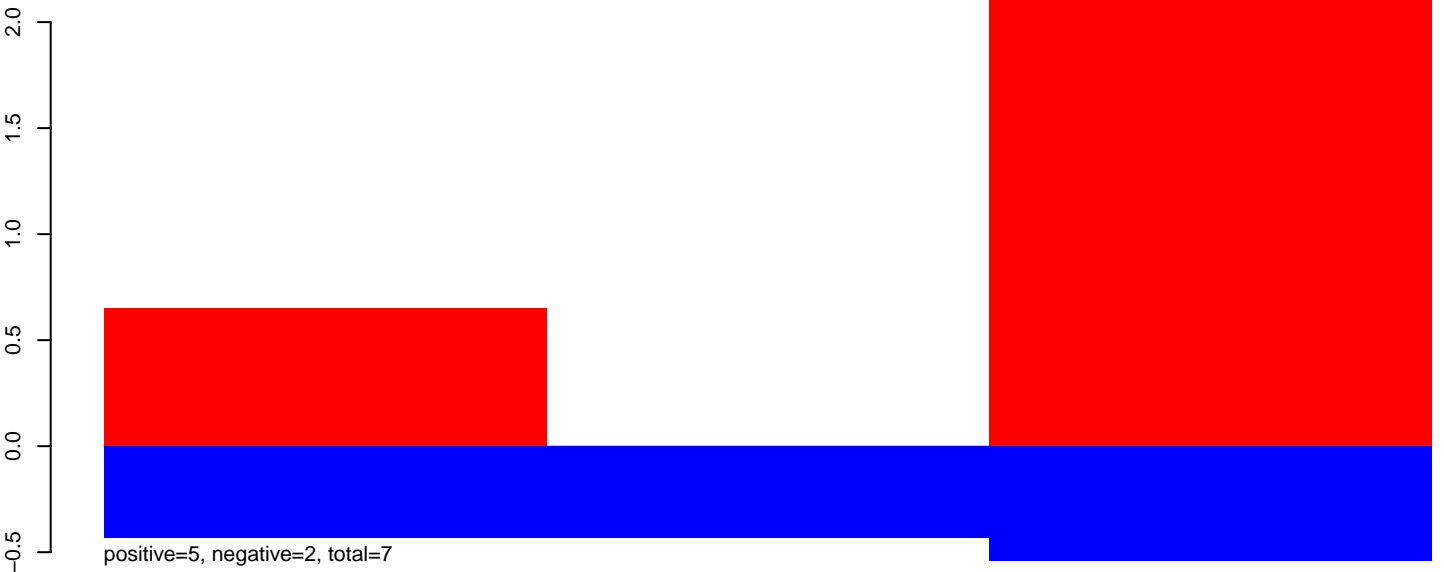
AnGam_Sua5b_cells.24_35.rep



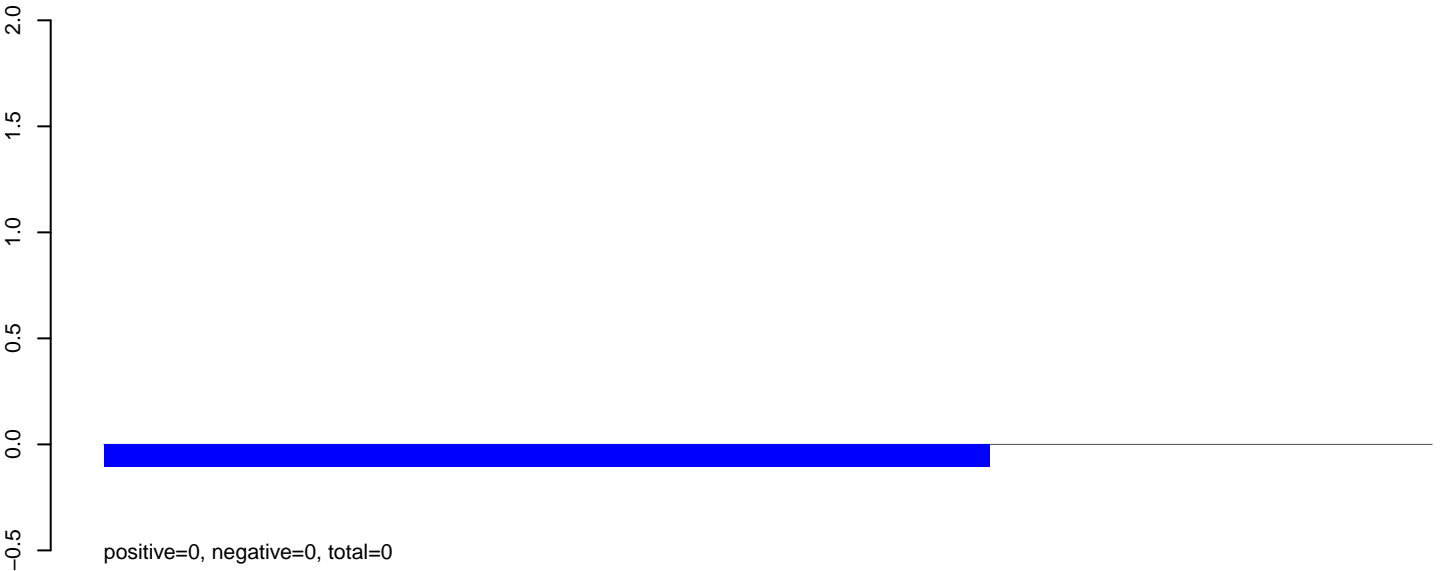
AnGam_Sua5b_cells.rep



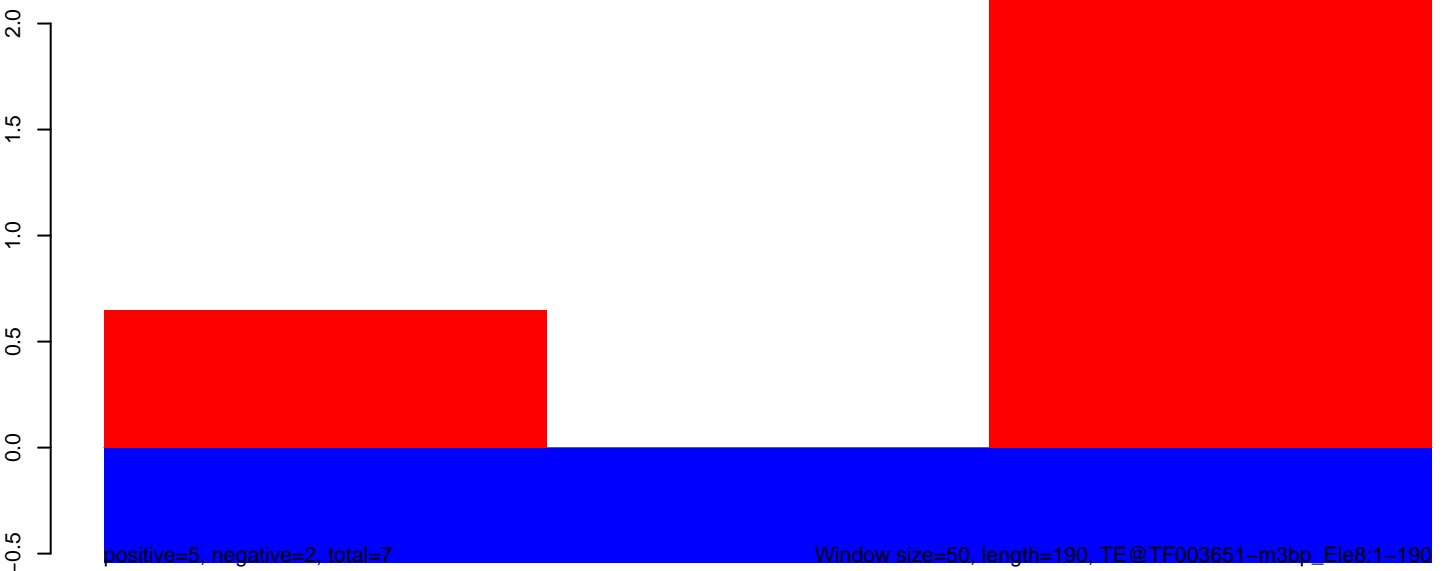
AnGam_Sua5b_cells.18_23.rep



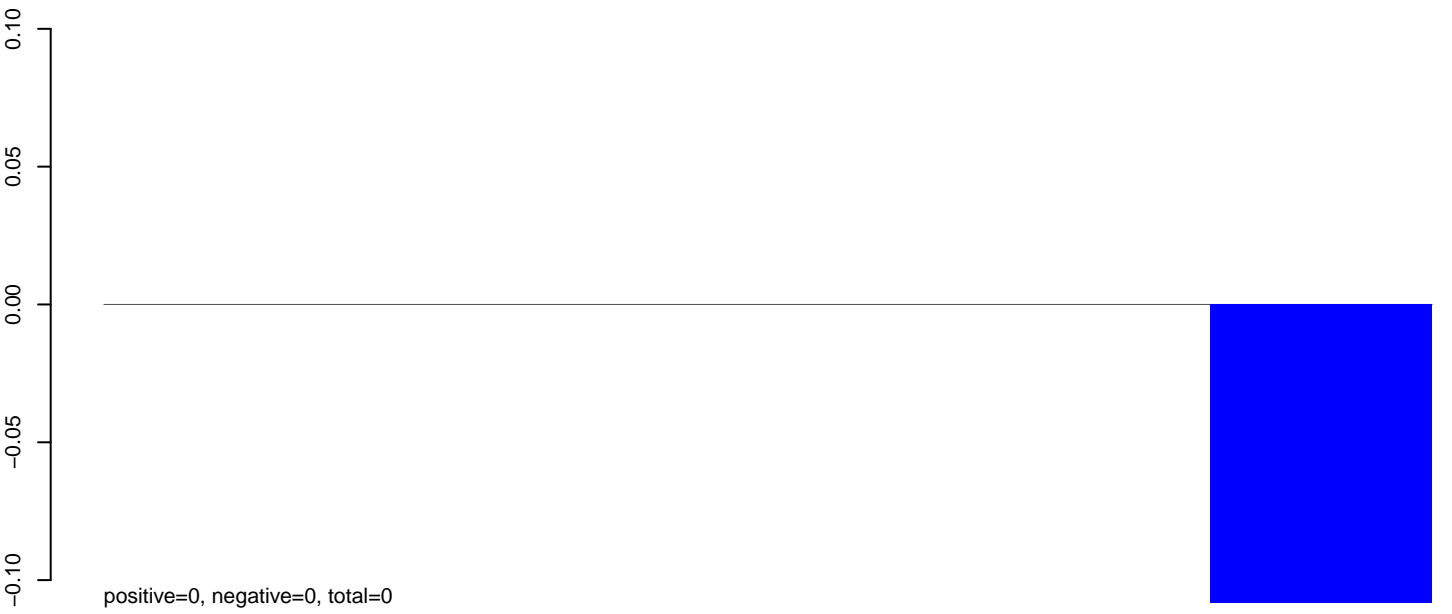
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



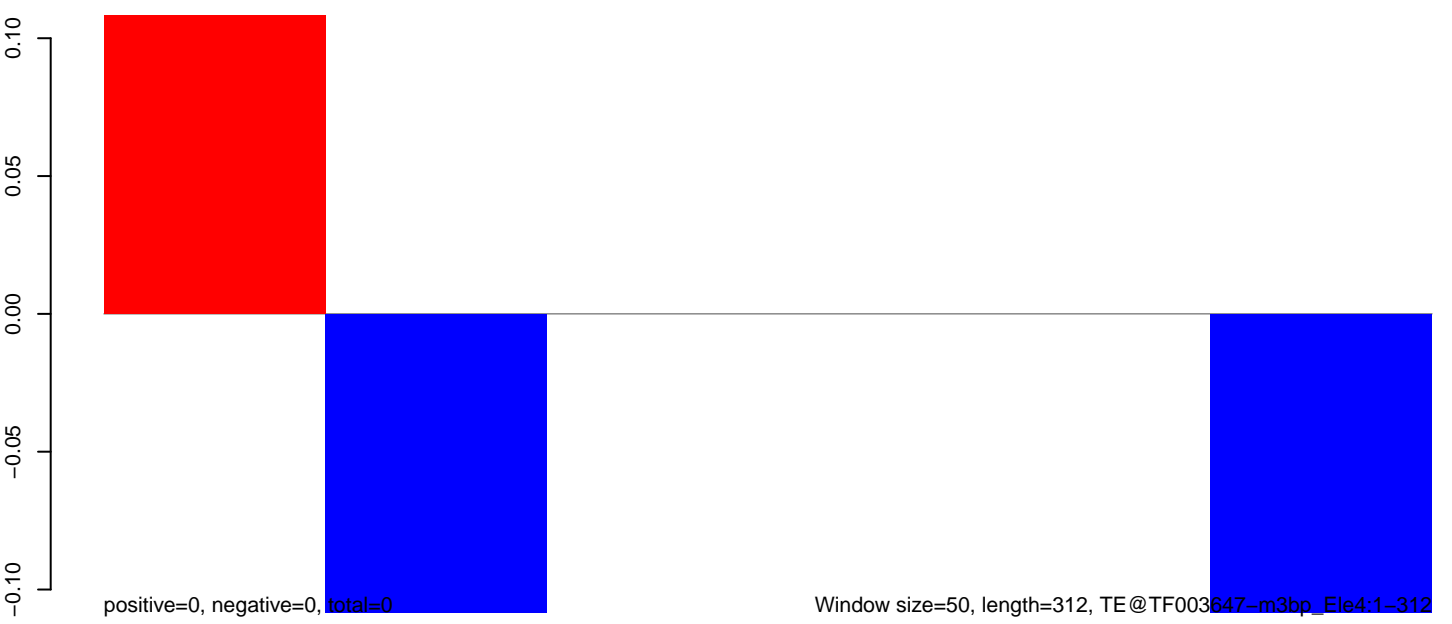
AnGam_Sua5b_cells.18_23.rep



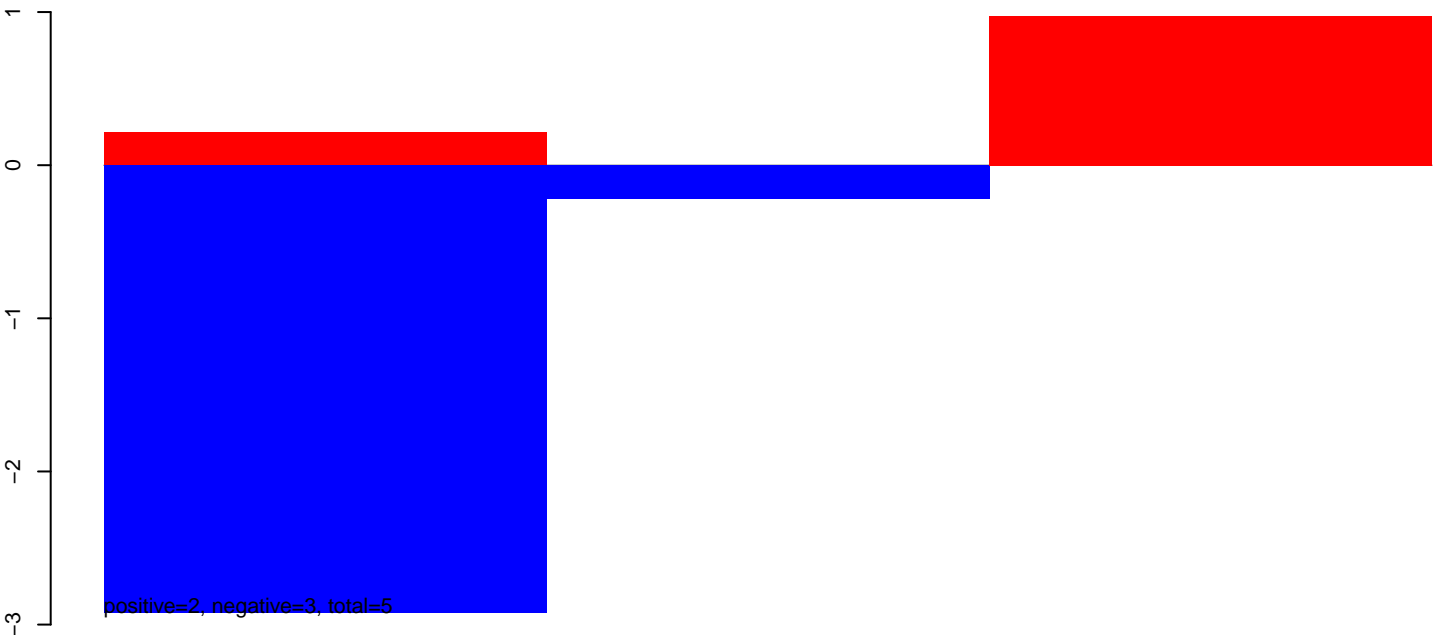
AnGam_Sua5b_cells.24_35.rep



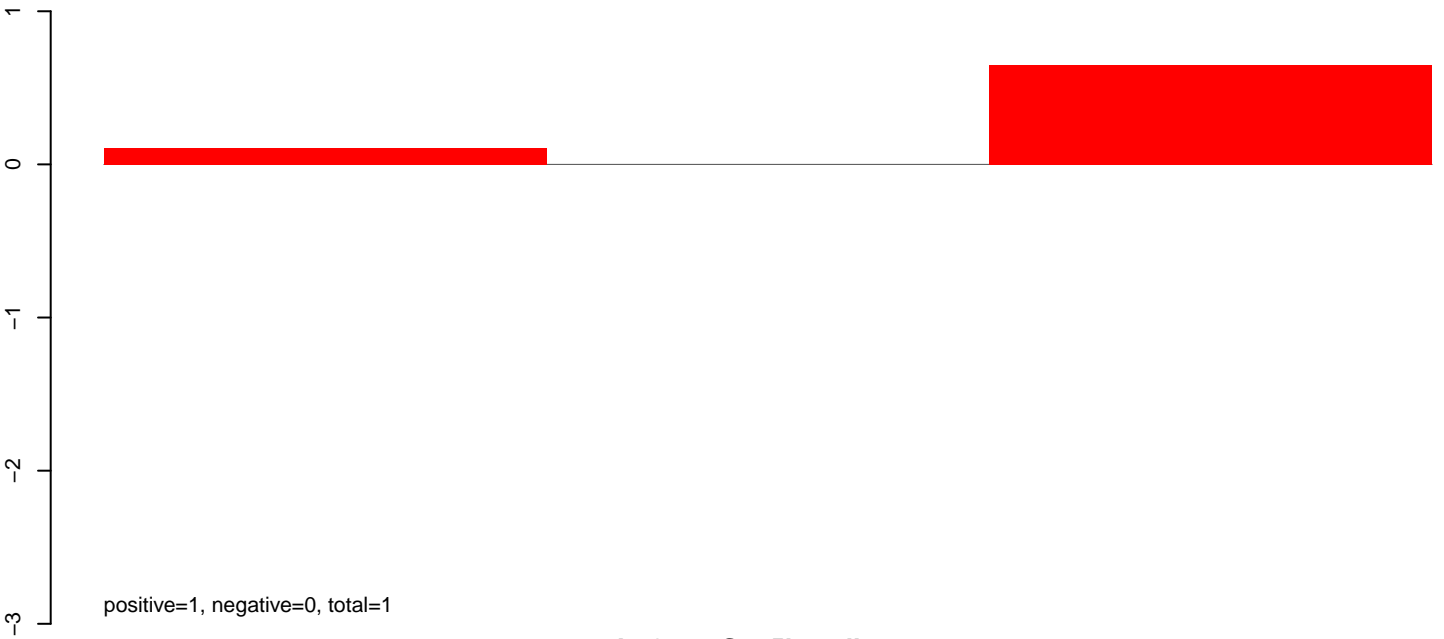
AnGam_Sua5b_cells.rep



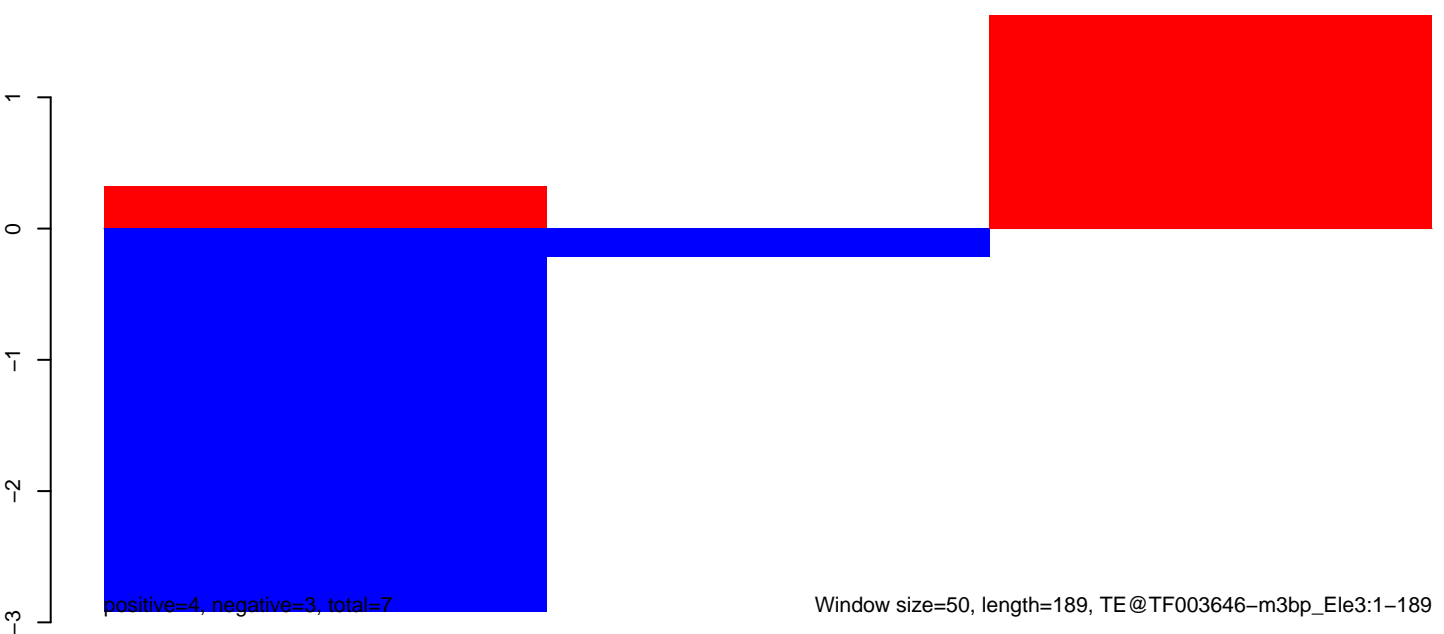
AnGam_Sua5b_cells.18_23.rep



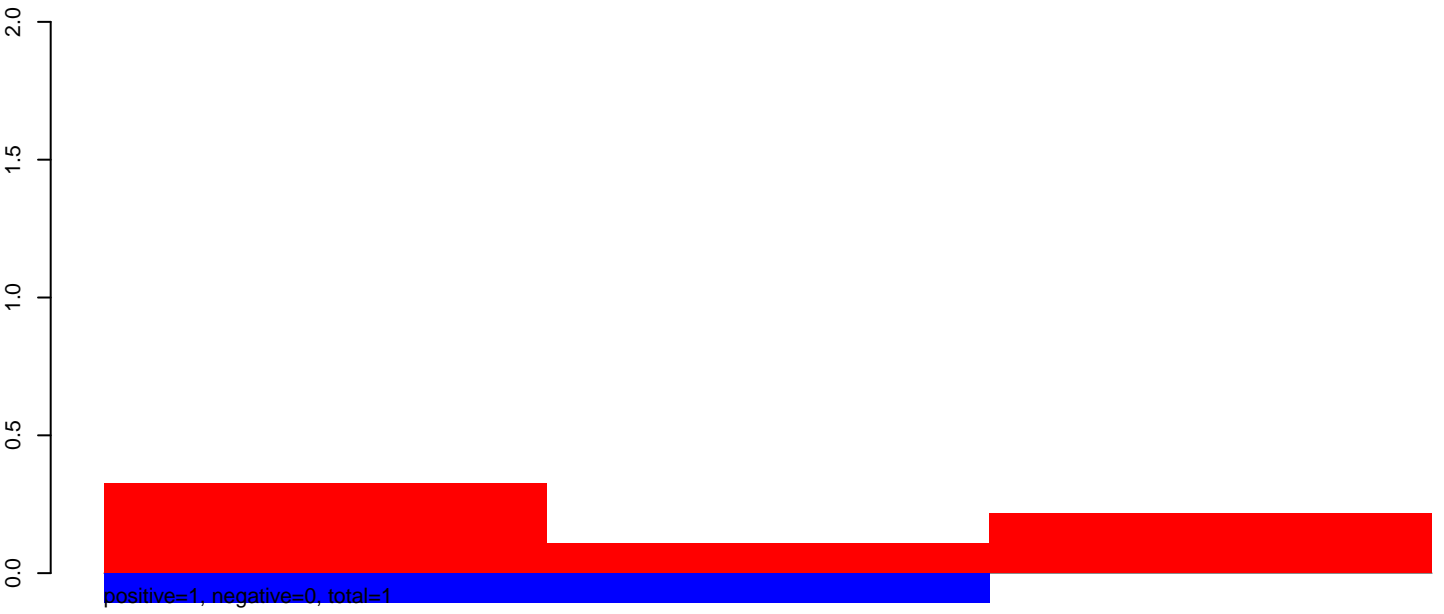
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



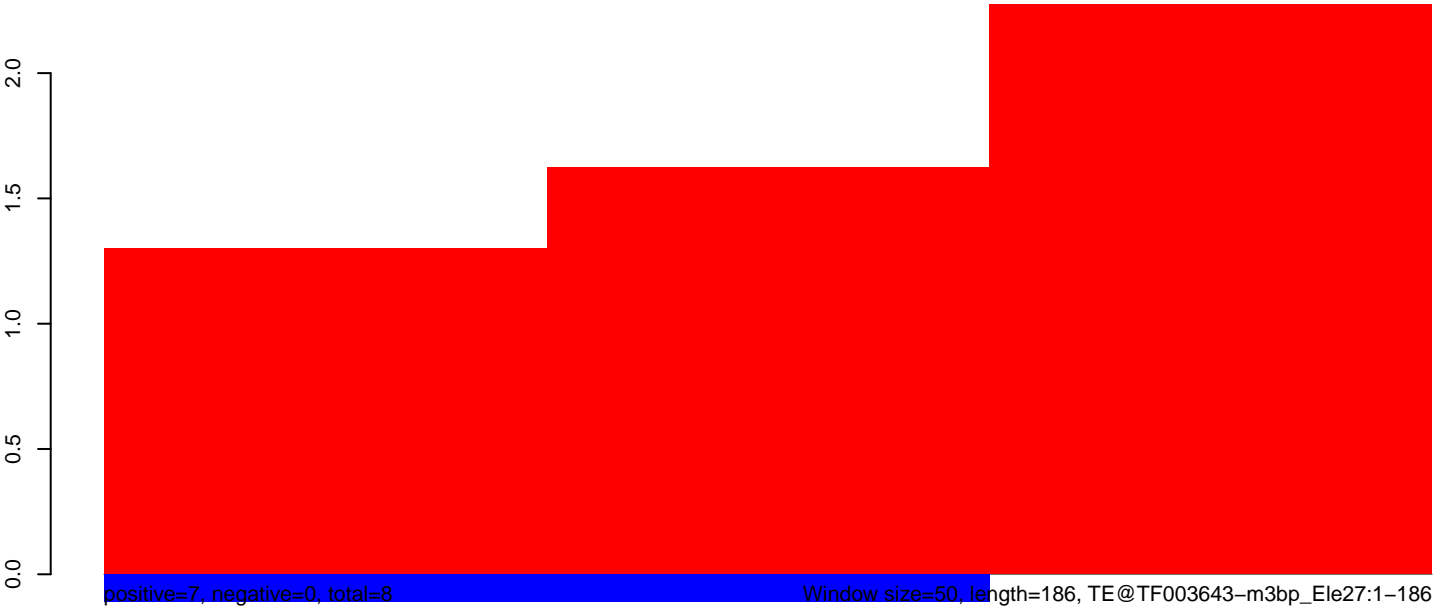
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



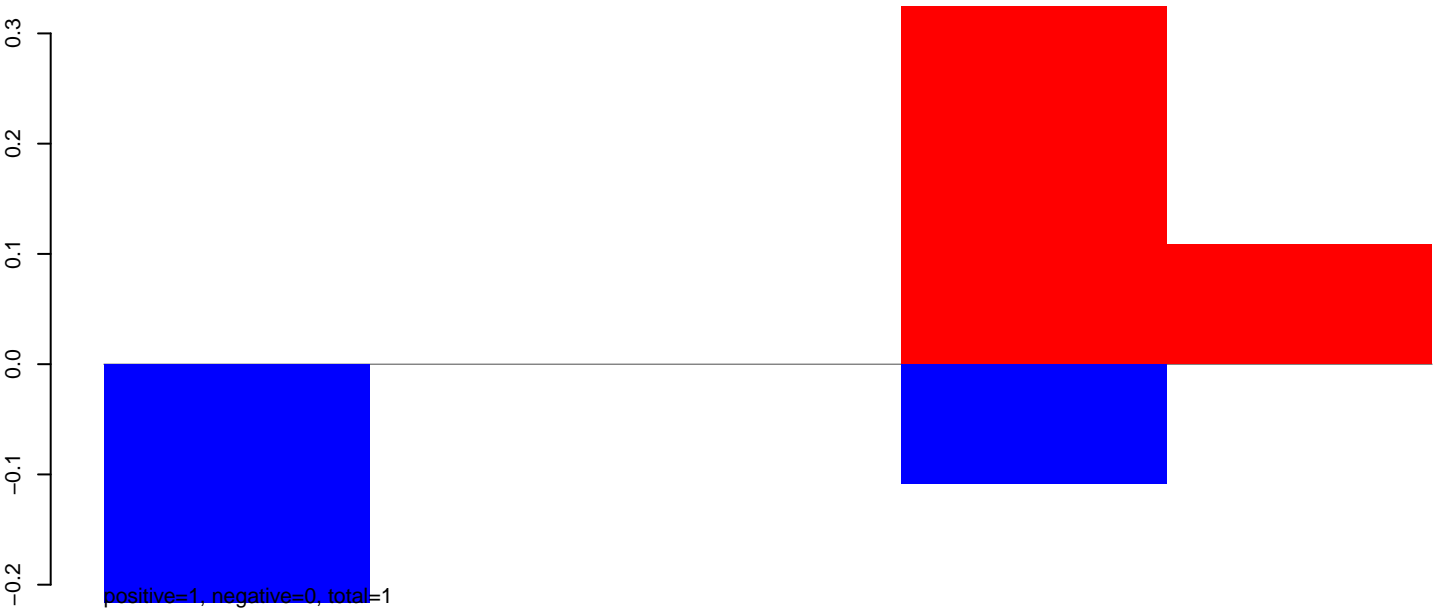
AnGam_Sua5b_cells.rep



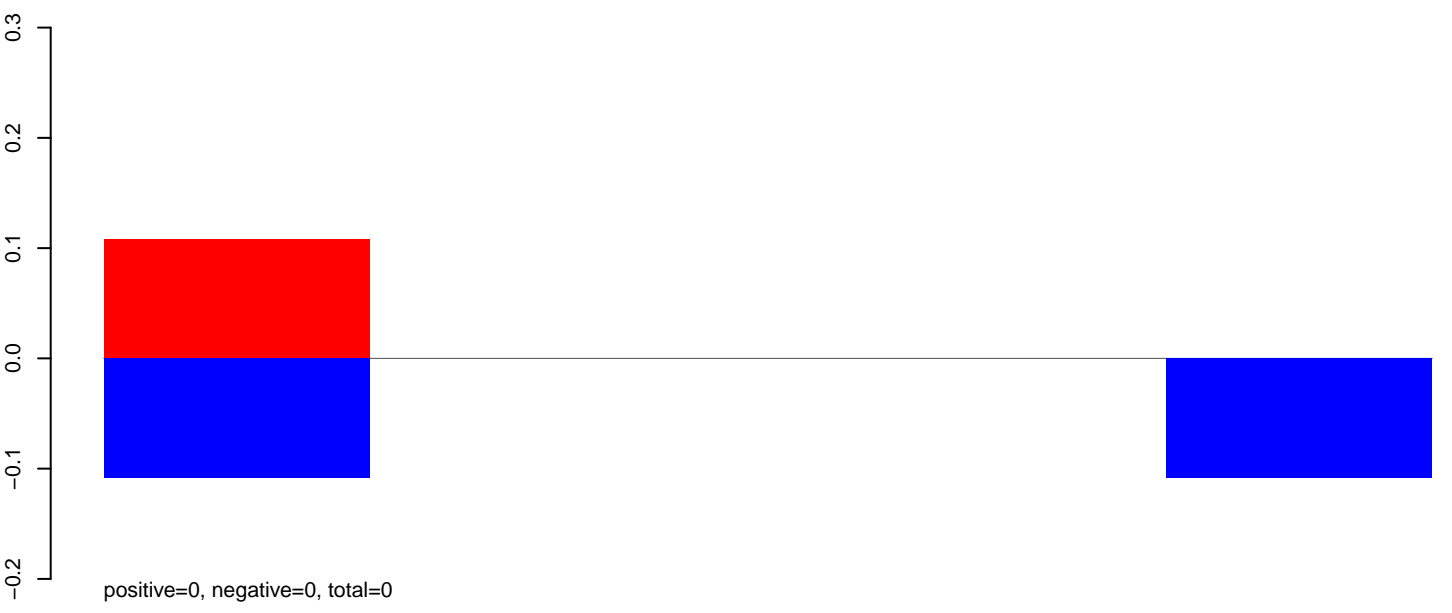
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50 100 150 200

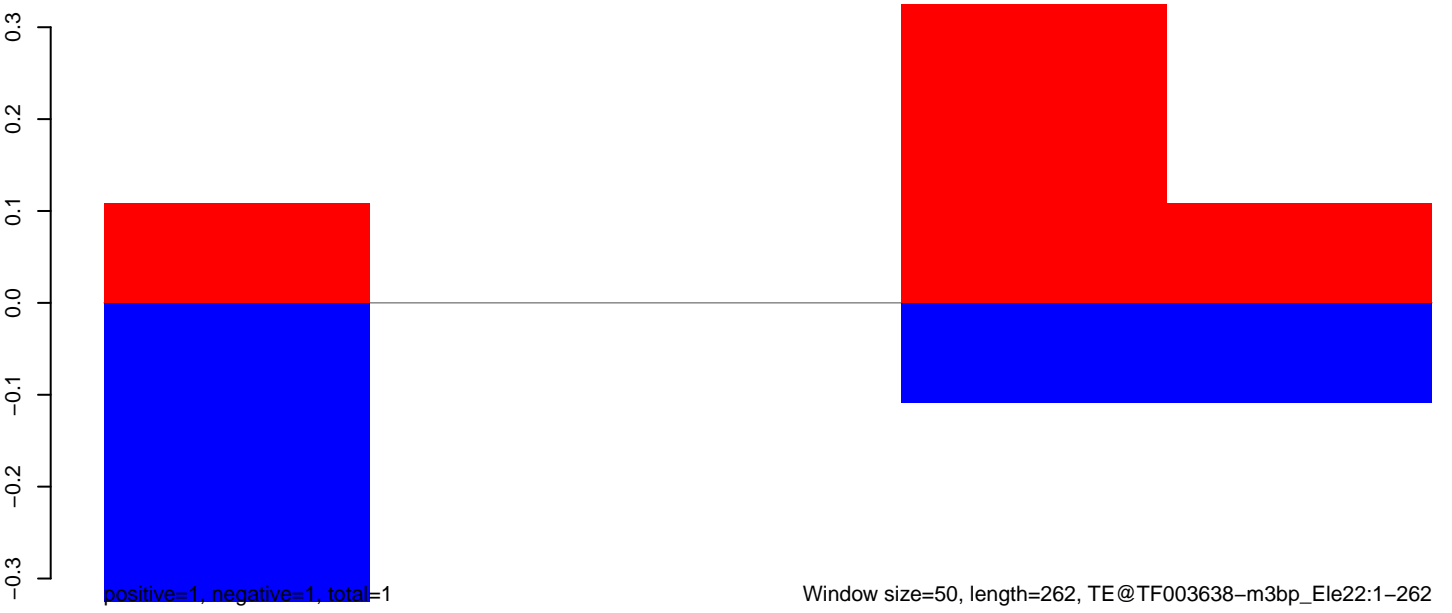
AnGam_Sua5b_cells.18_23.rep



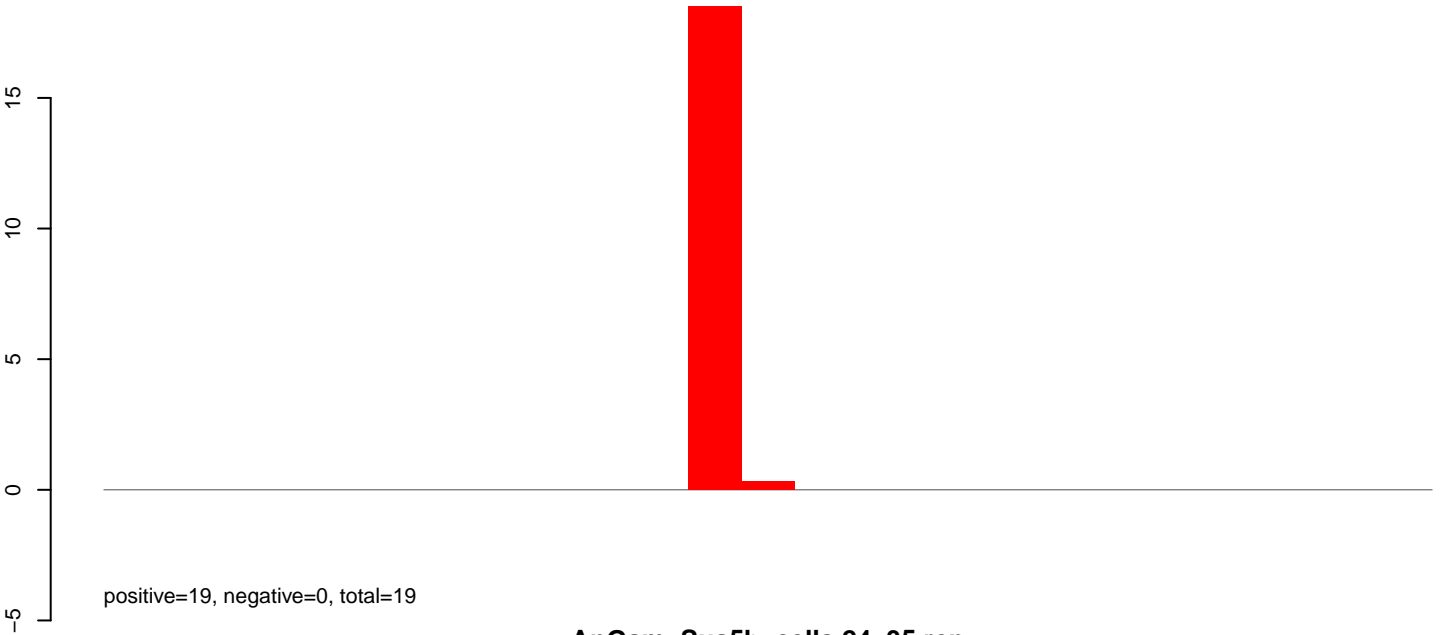
AnGam_Sua5b_cells.24_35.rep



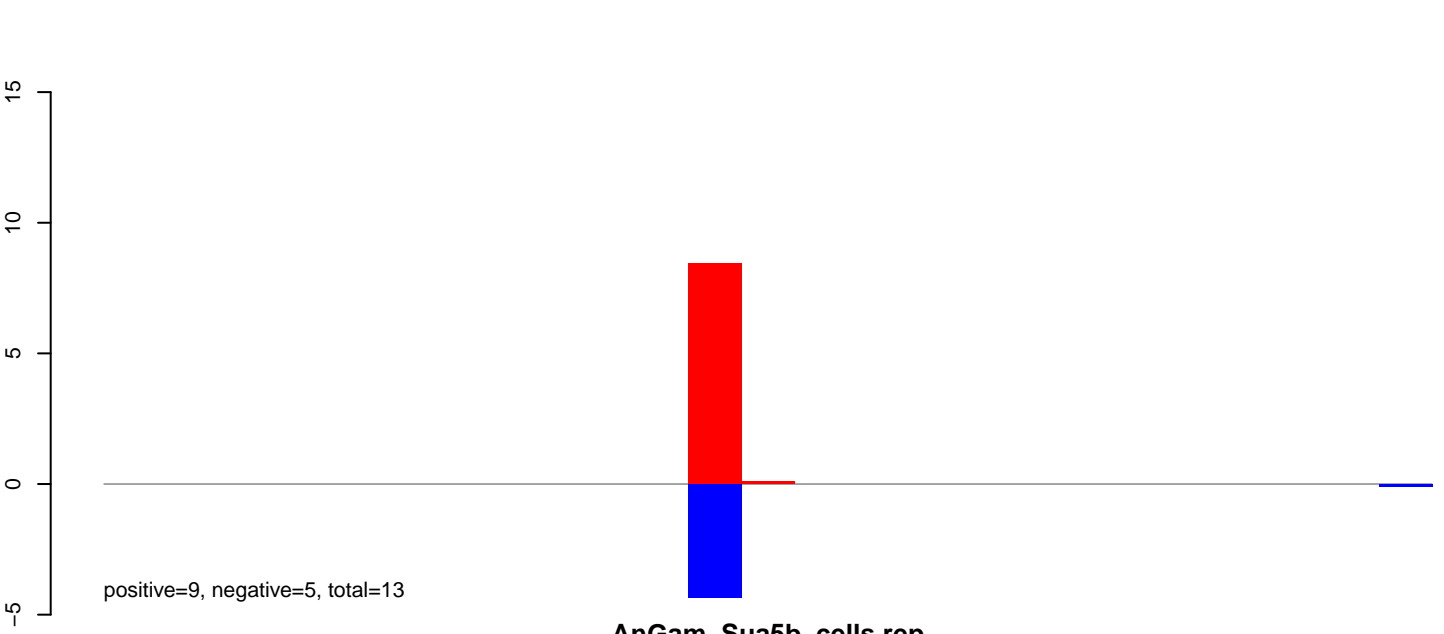
AnGam_Sua5b_cells.rep



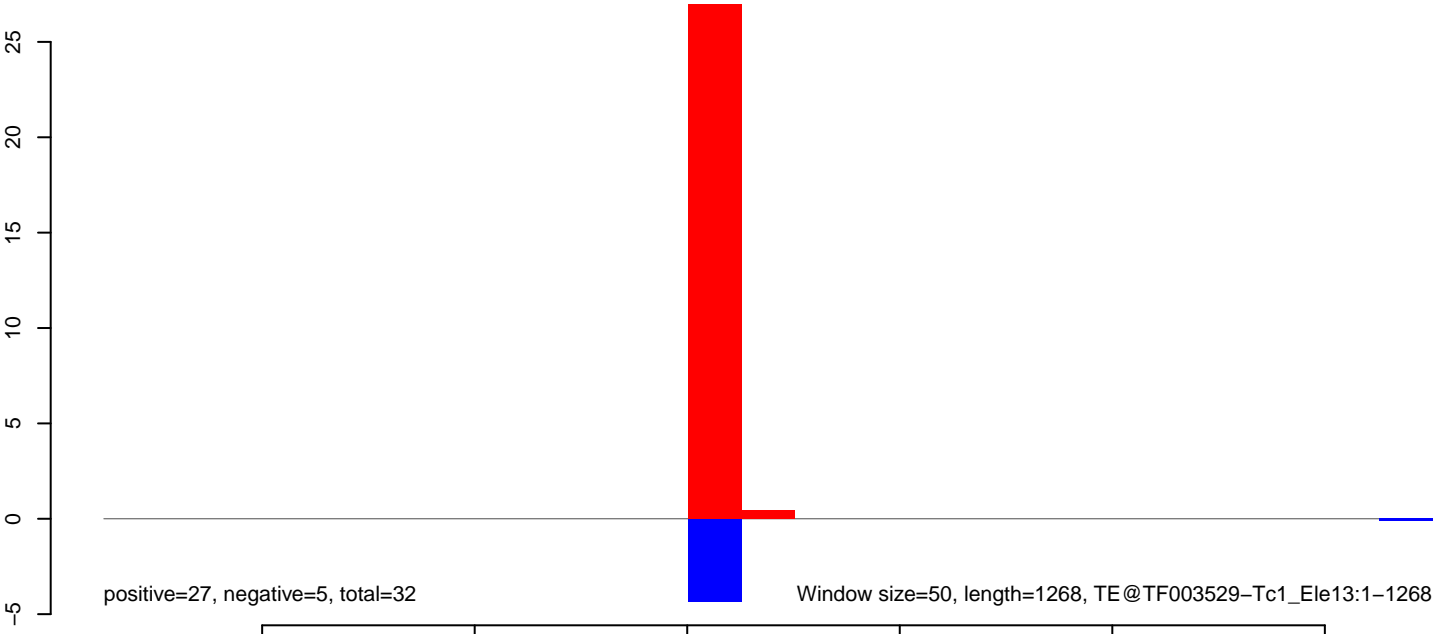
AnGam_Sua5b_cells.18_23.rep



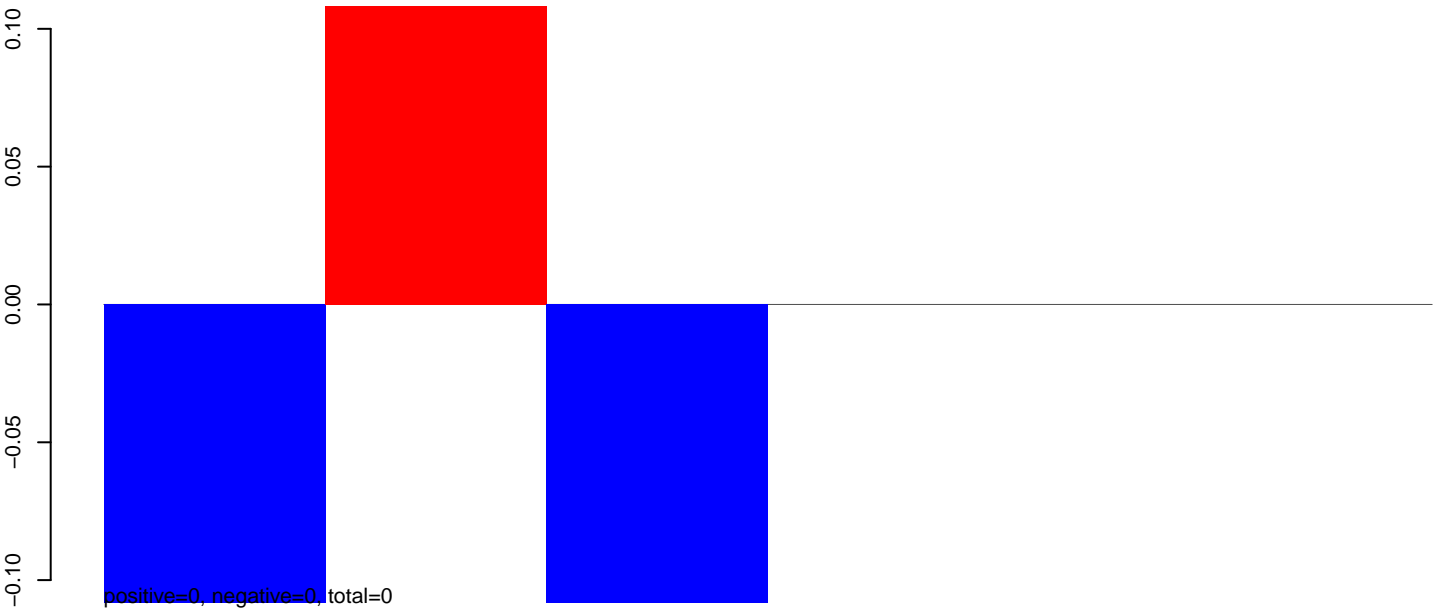
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



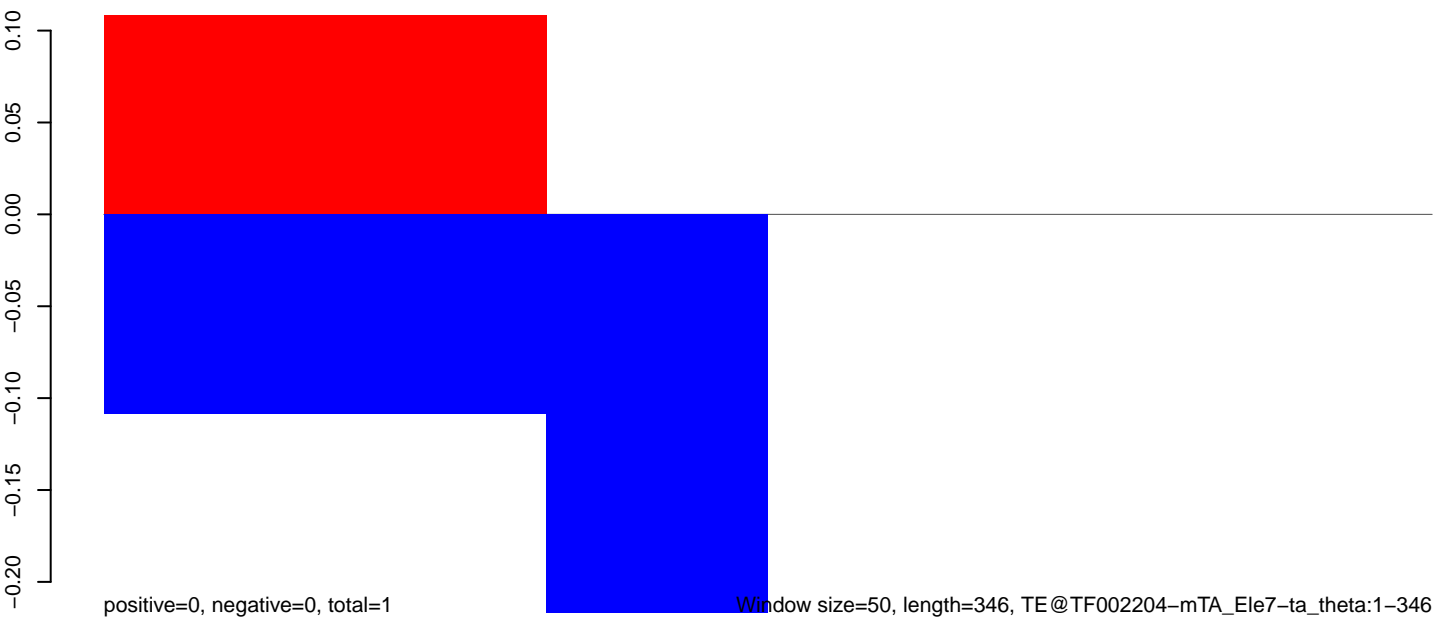
AnGam_Sua5b_cells.18_23.rep



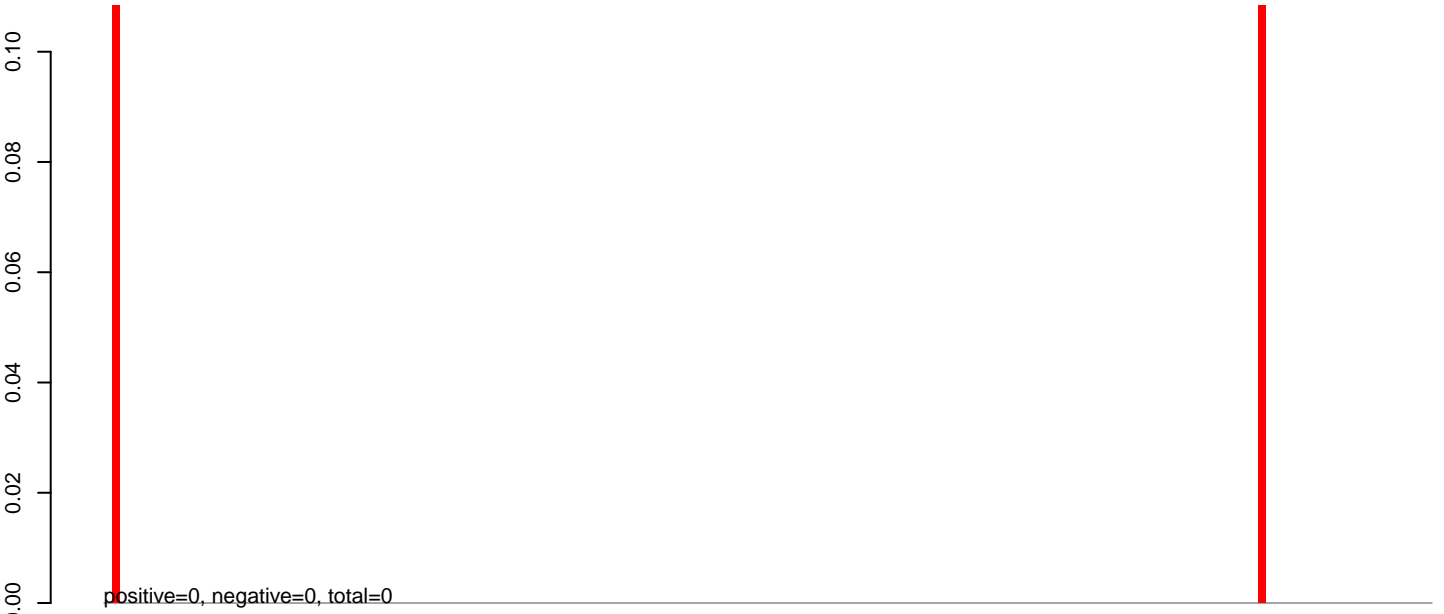
AnGam_Sua5b_cells.24_35.rep



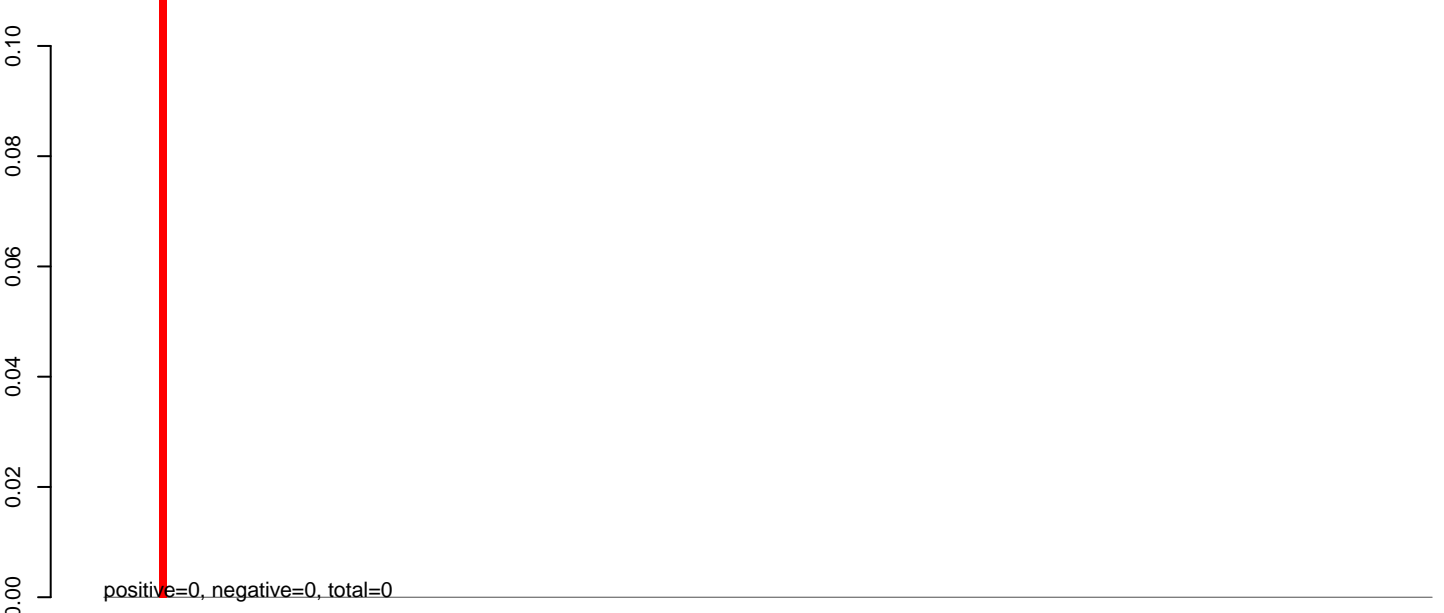
AnGam_Sua5b_cells.rep



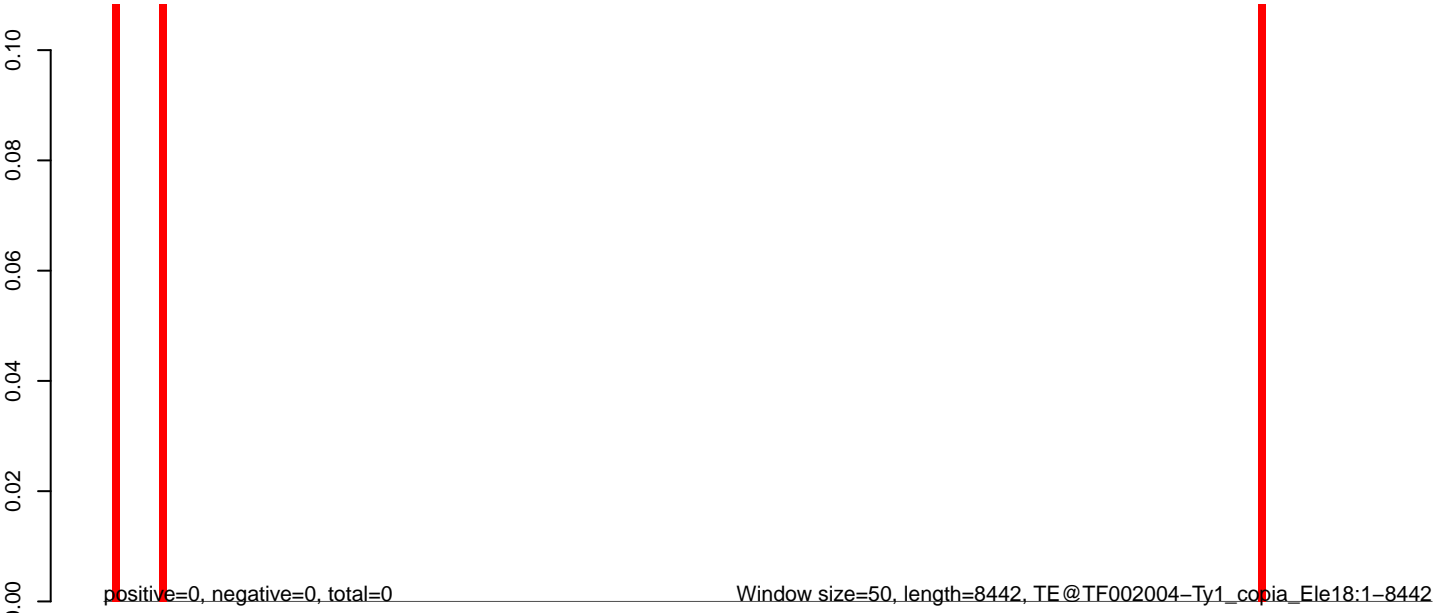
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

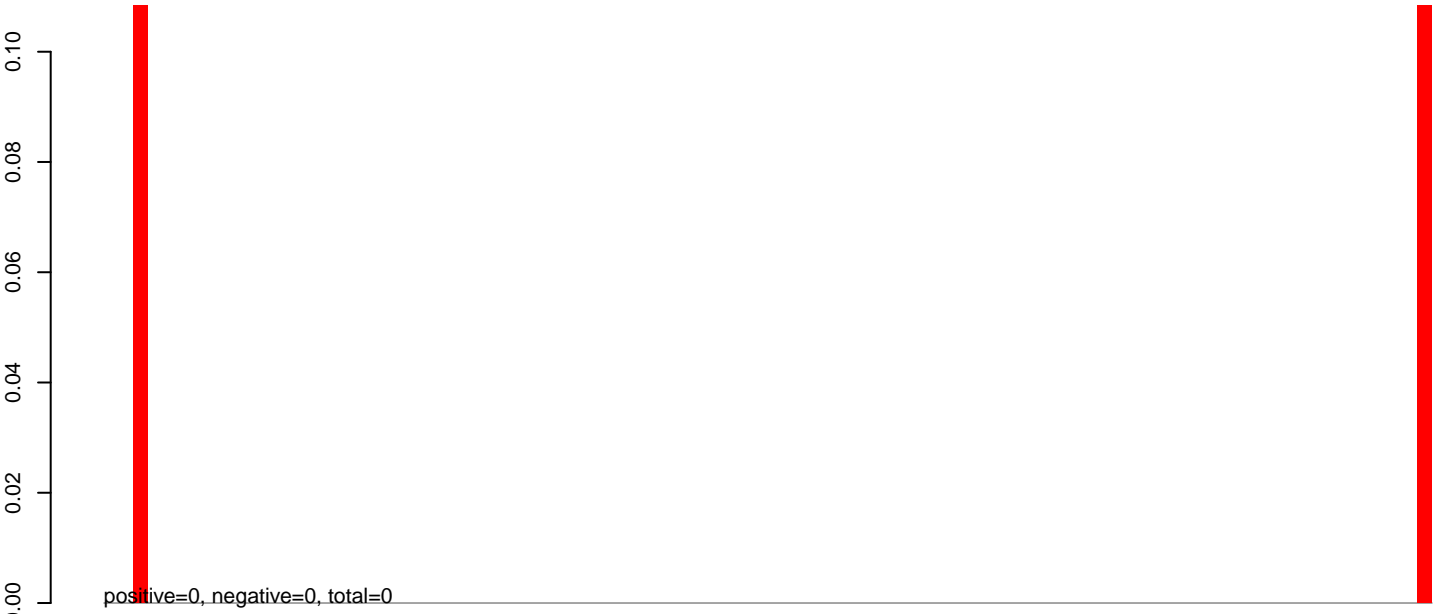


AnGam_Sua5b_cells.rep

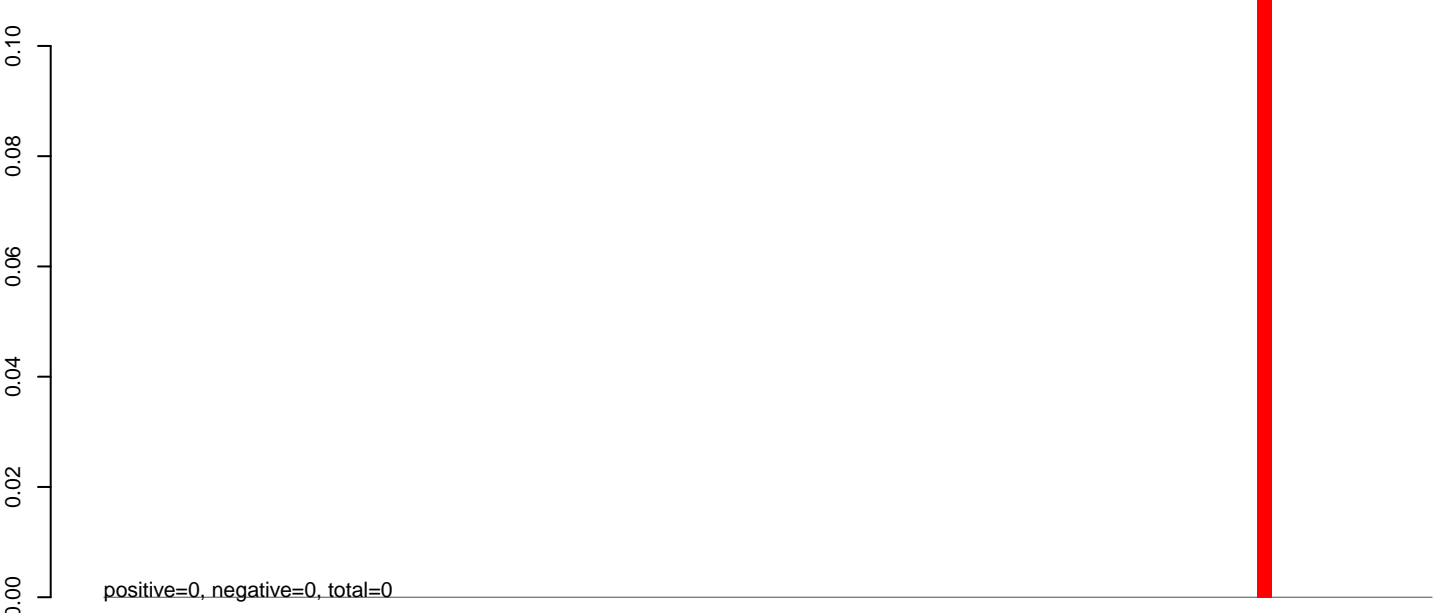


0 2000 4000 6000 8000

AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

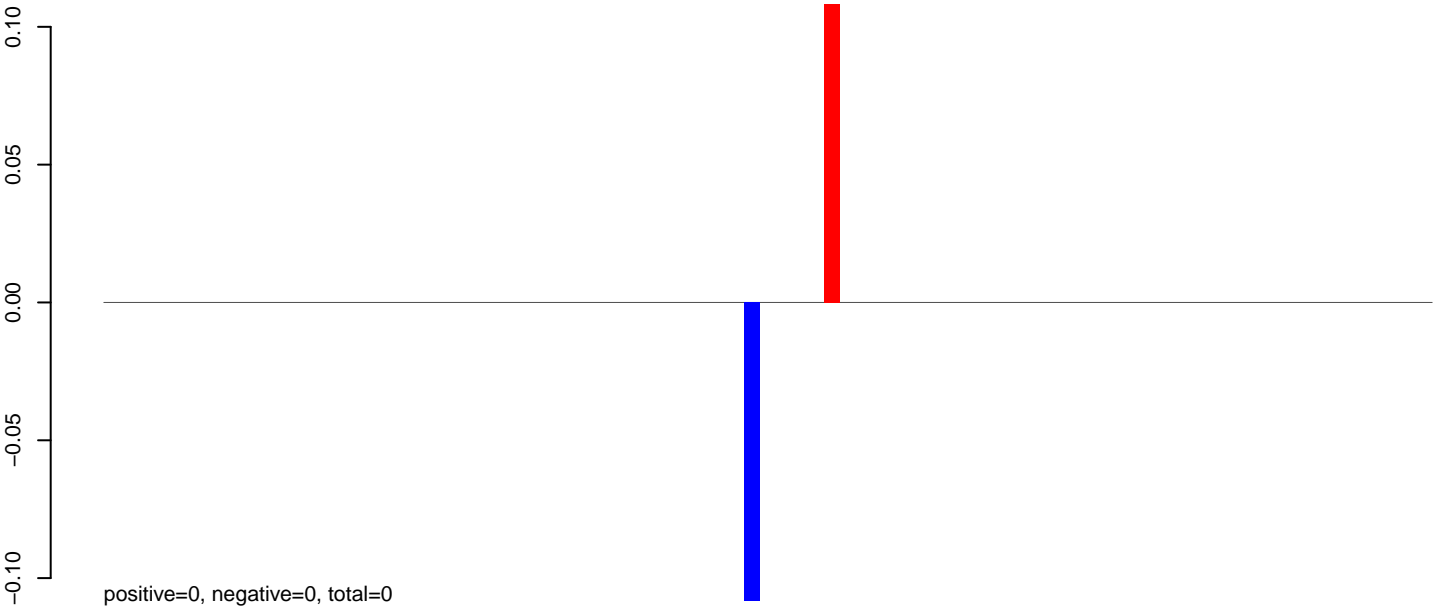


AnGam_Sua5b_cells.rep

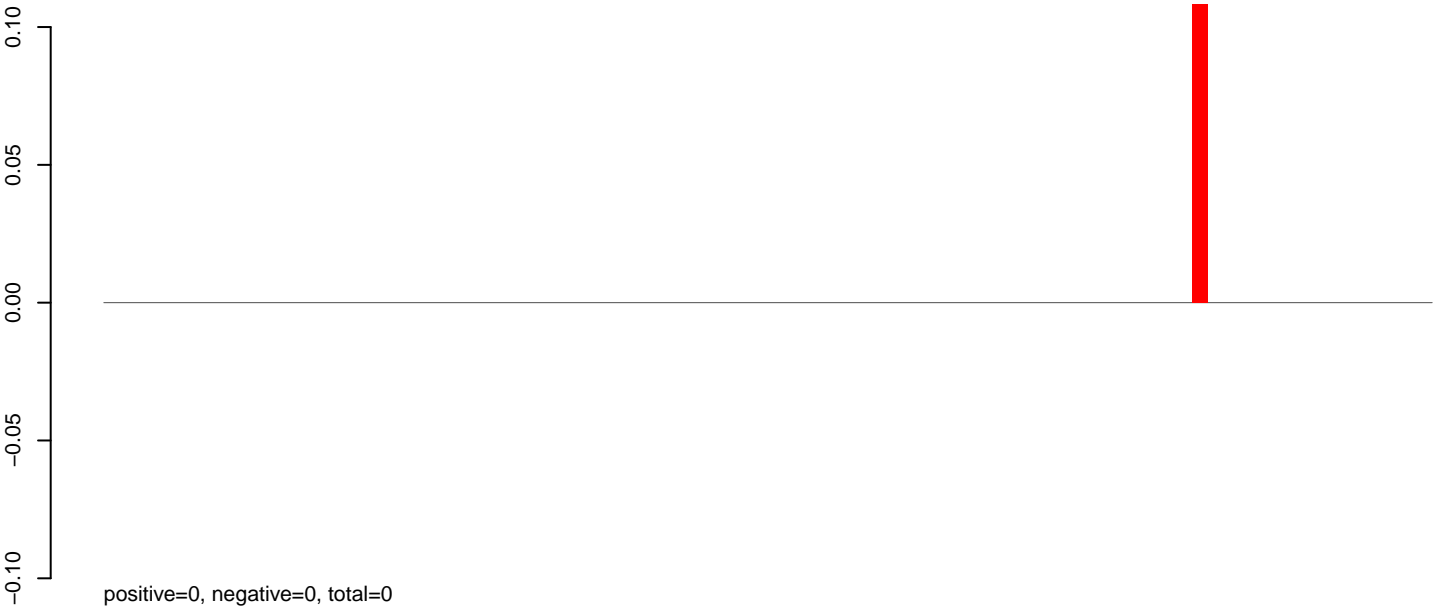


Window size=50, length=4582, TE@TF002001-Ty1_copia_Ele15:1-4582

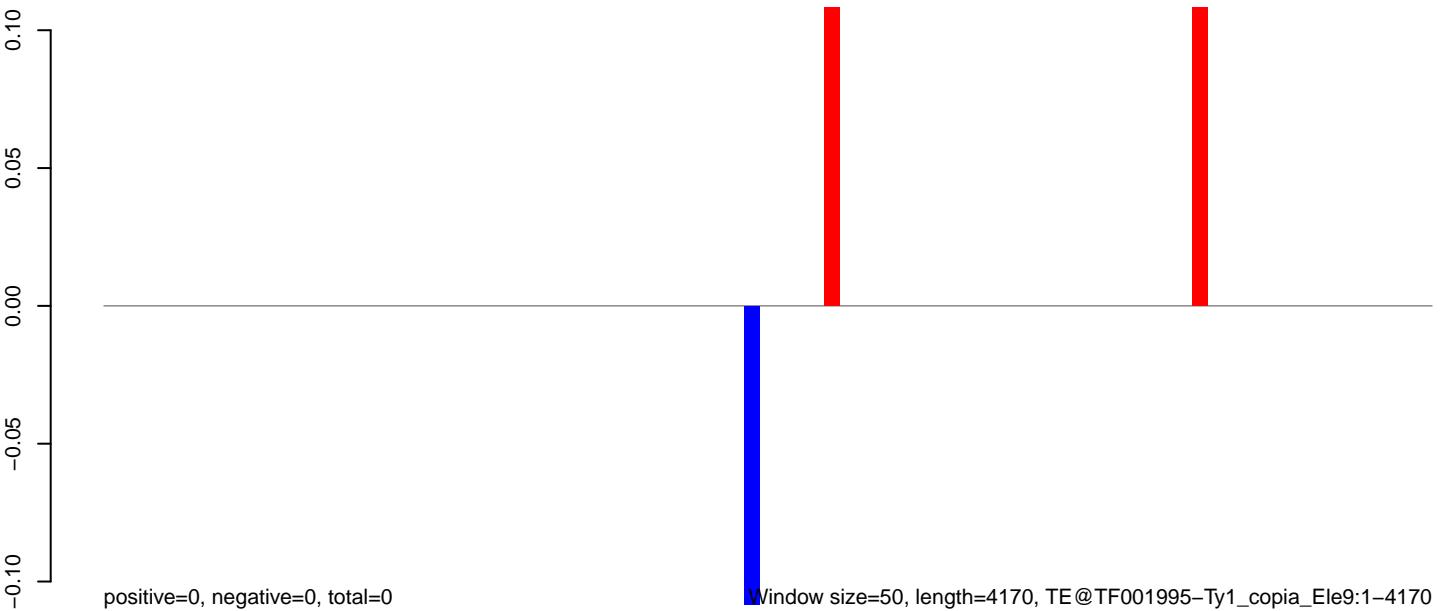
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



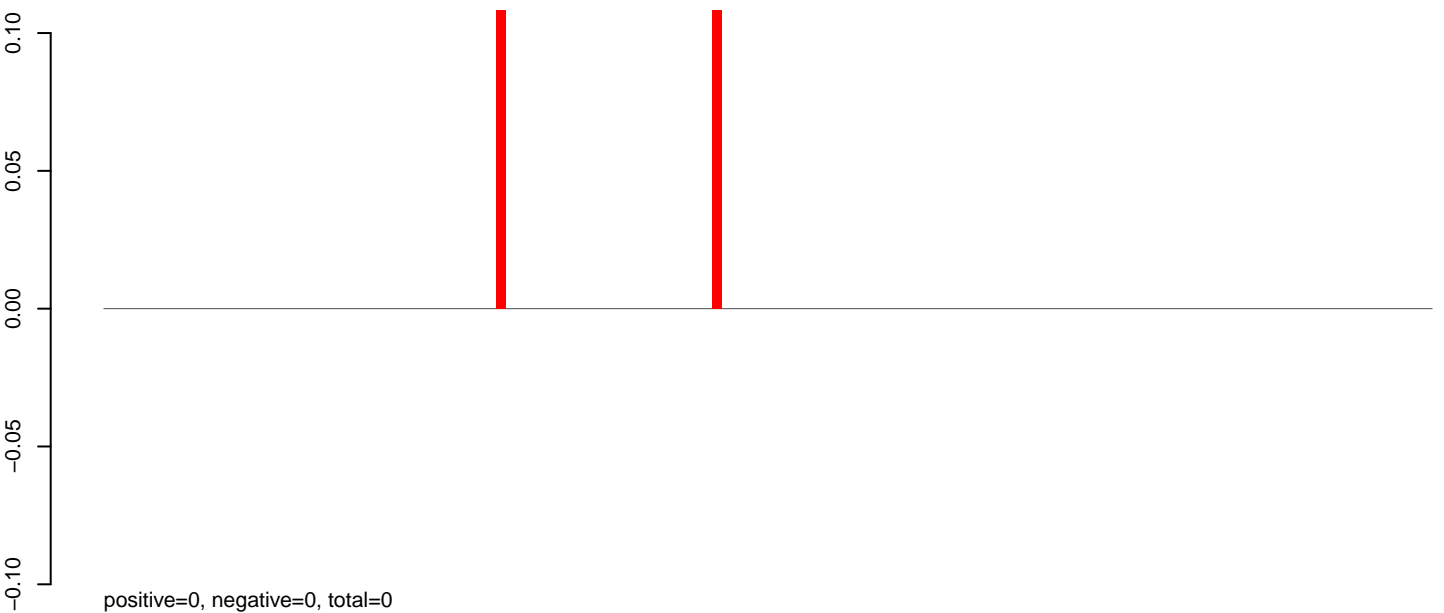
Window size=50, length=4170, TE@TF001995-Ty1_copia_Ele9:1-4170

0 1000 2000 3000 4000

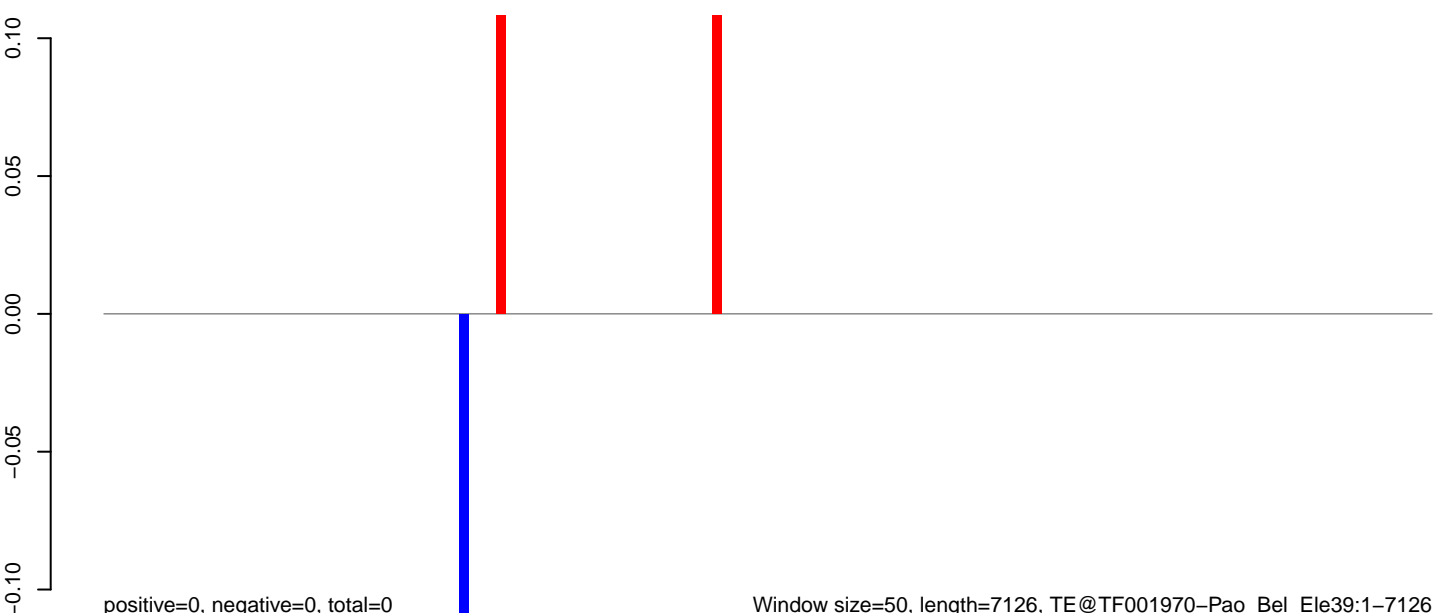
AnGam_Sua5b_cells.18_23.rep



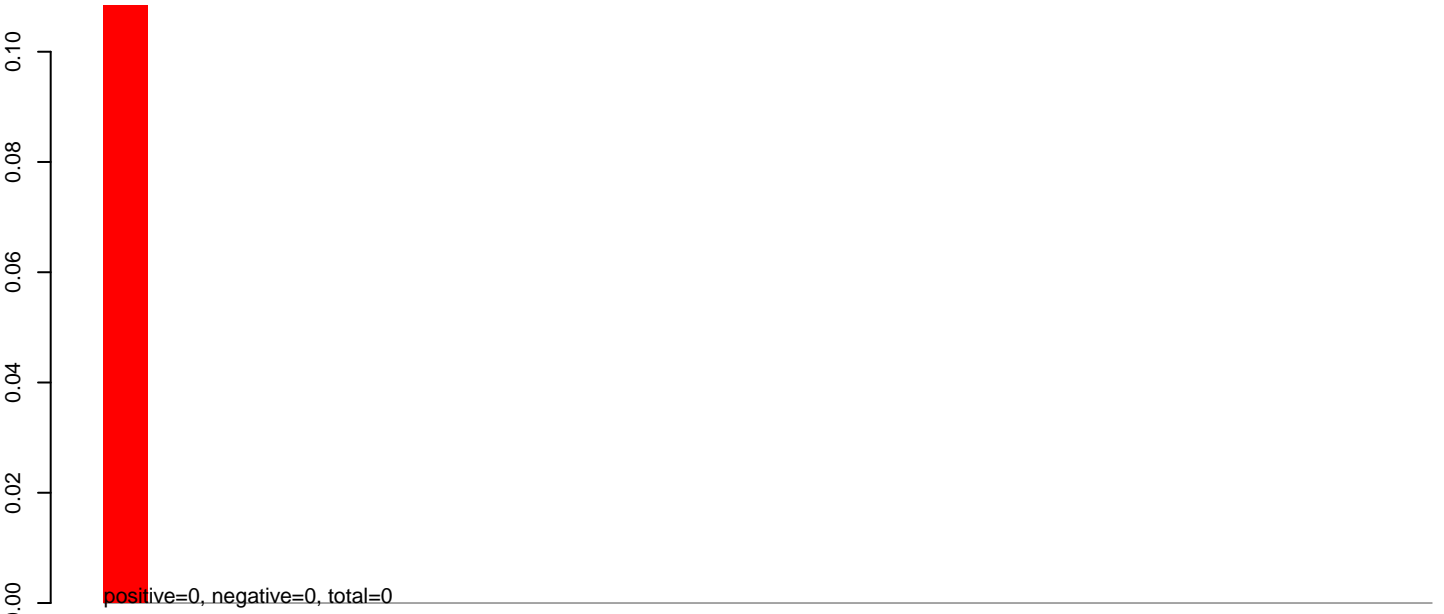
AnGam_Sua5b_cells.24_35.rep



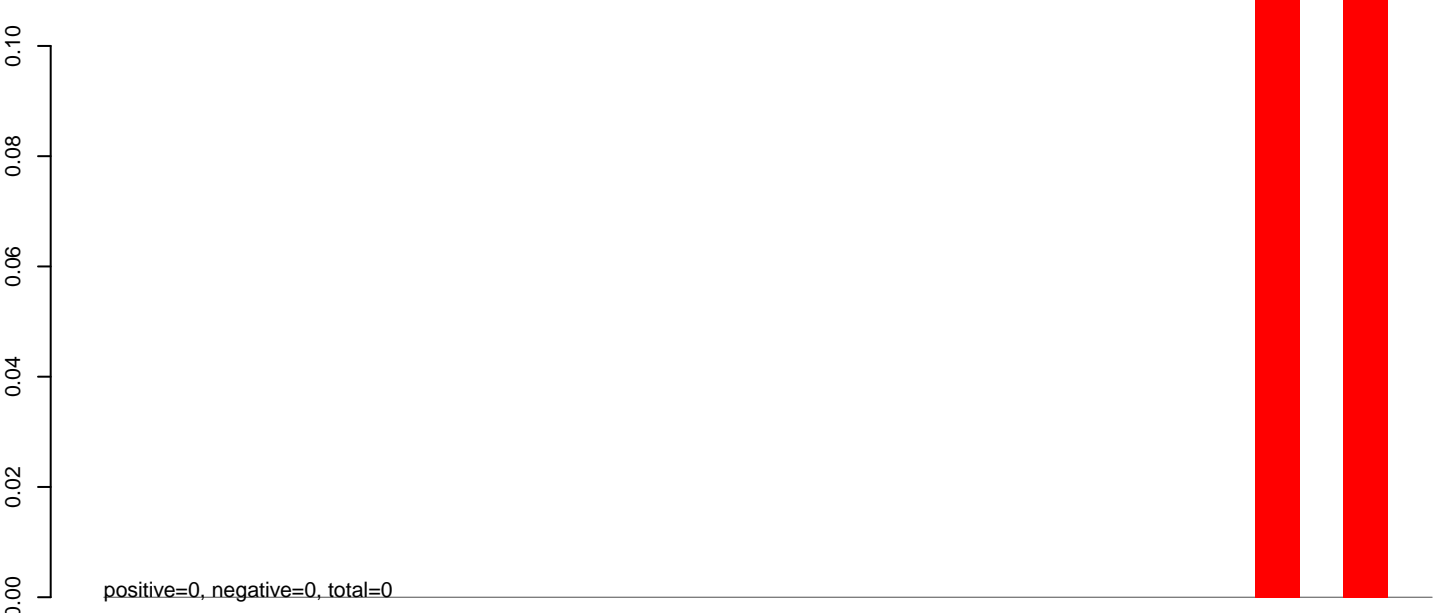
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



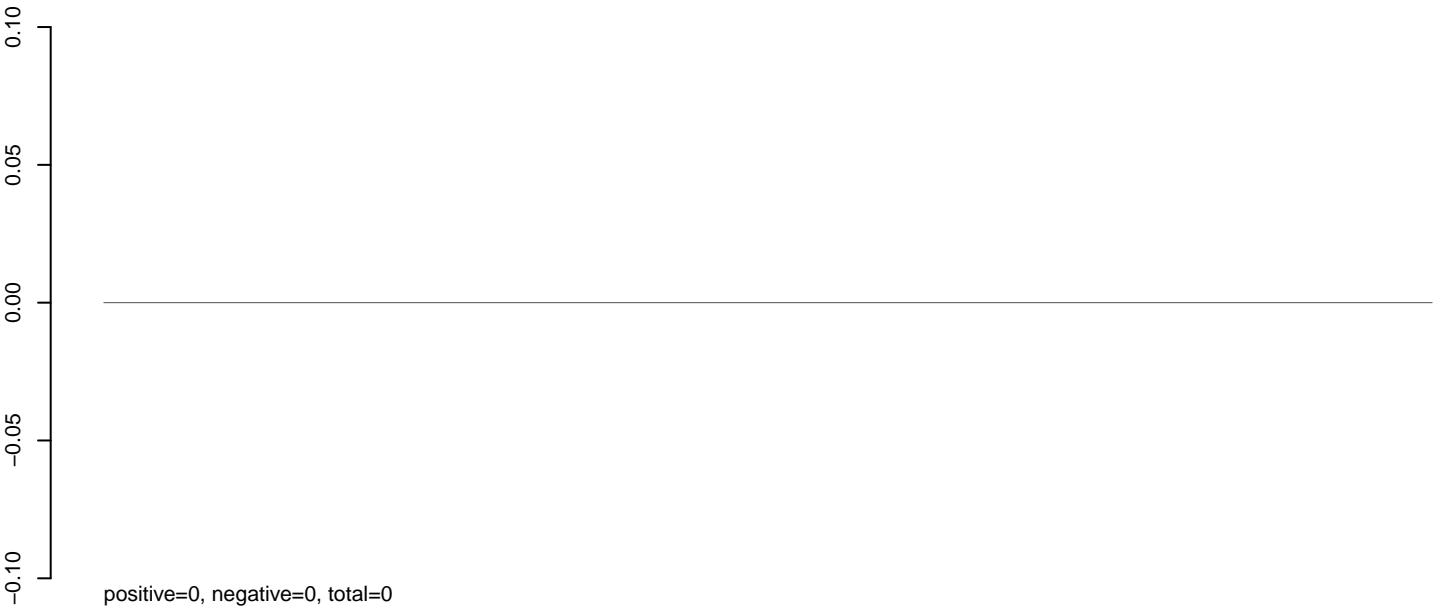
AnGam_Sua5b_cells.rep



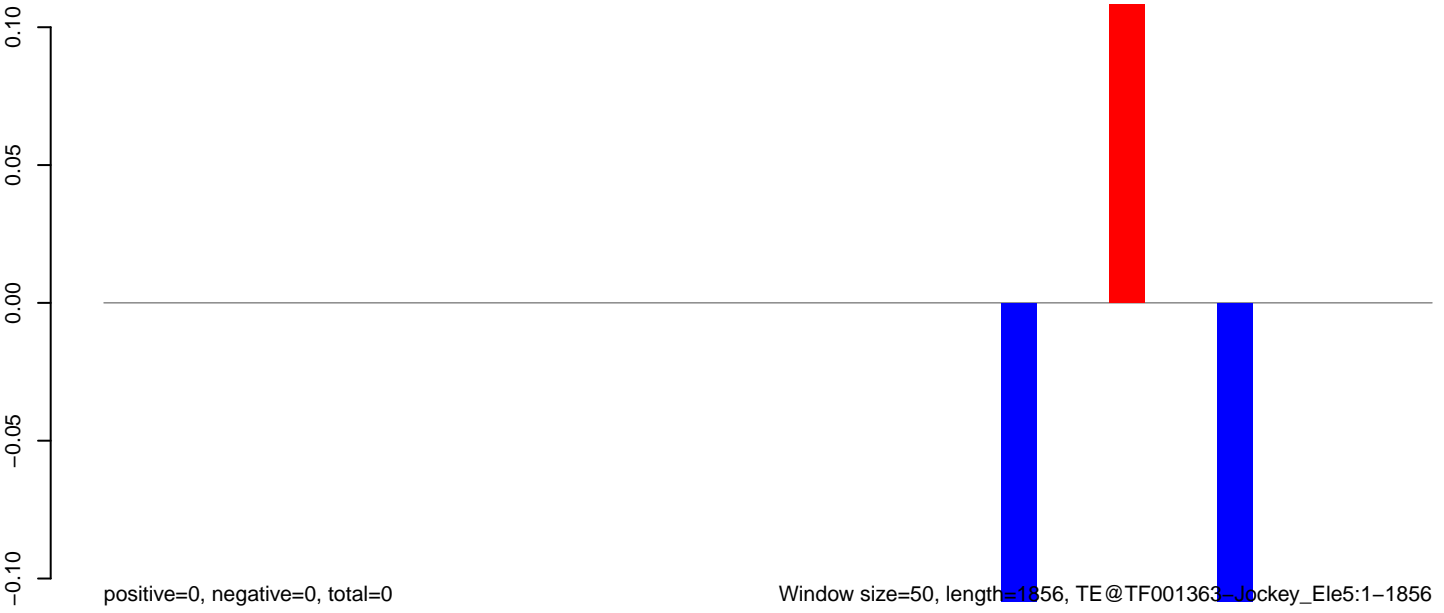
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



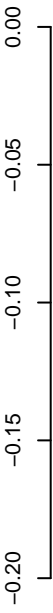
AnGam_Sua5b_cells.rep



Window size=50, length=1856, TE@TF001363-Jockey_Ele5:1-1856

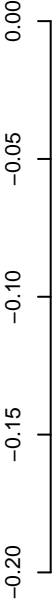
0 500 1000 1500

AnGam_Sua5b_cells.18_23.rep



positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep



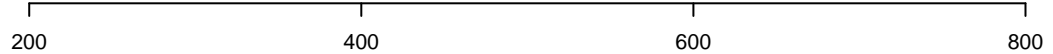
positive=0, negative=1, total=1

AnGam_Sua5b_cells.rep

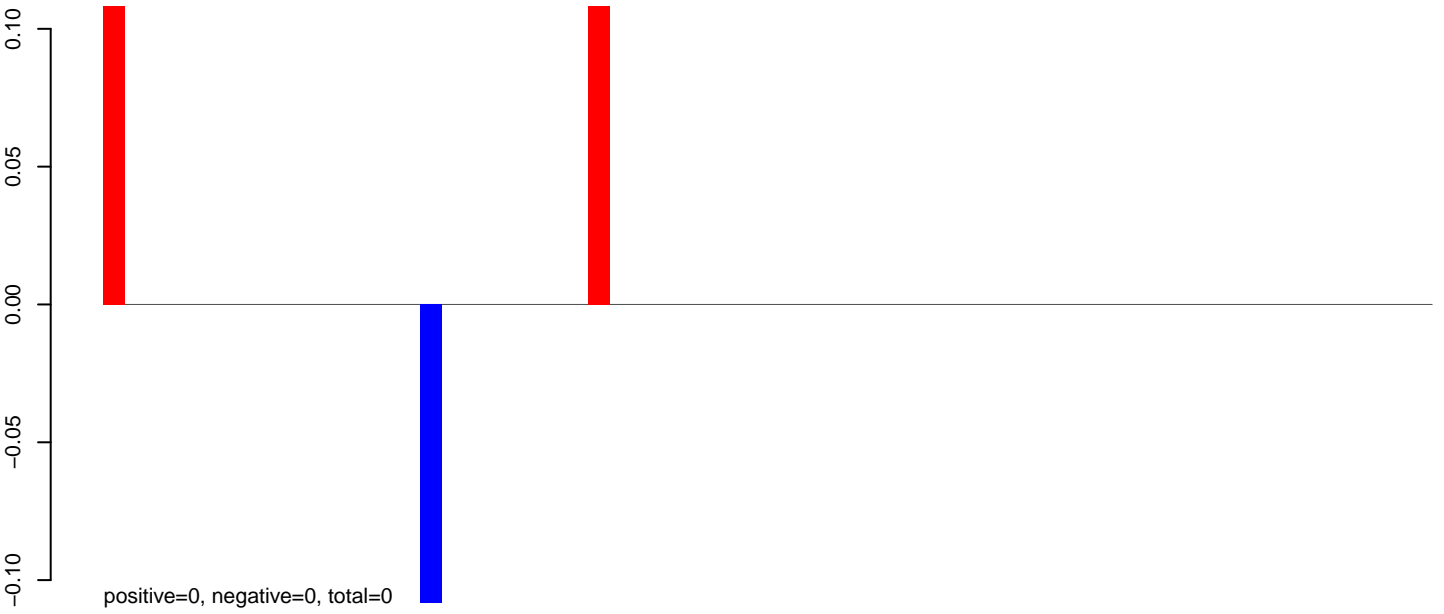


positive=0, negative=1, total=1

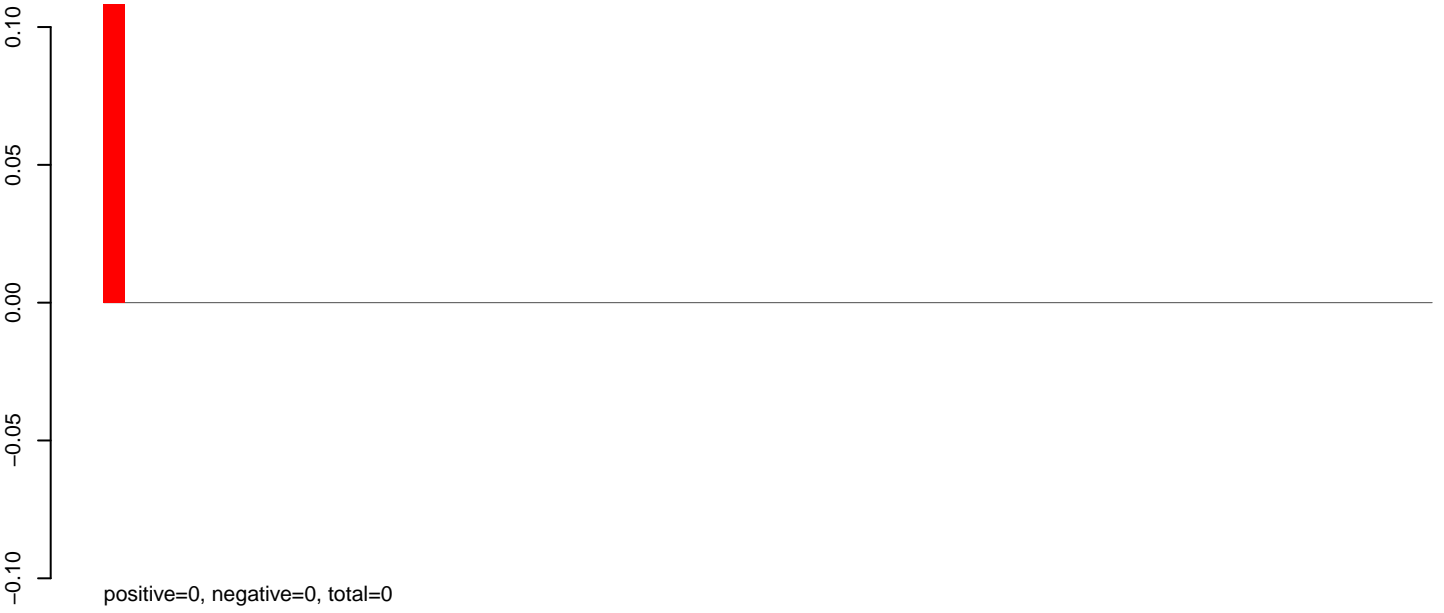
Window size=50, length=847, TE@TF001360-I_Ele7:1-847



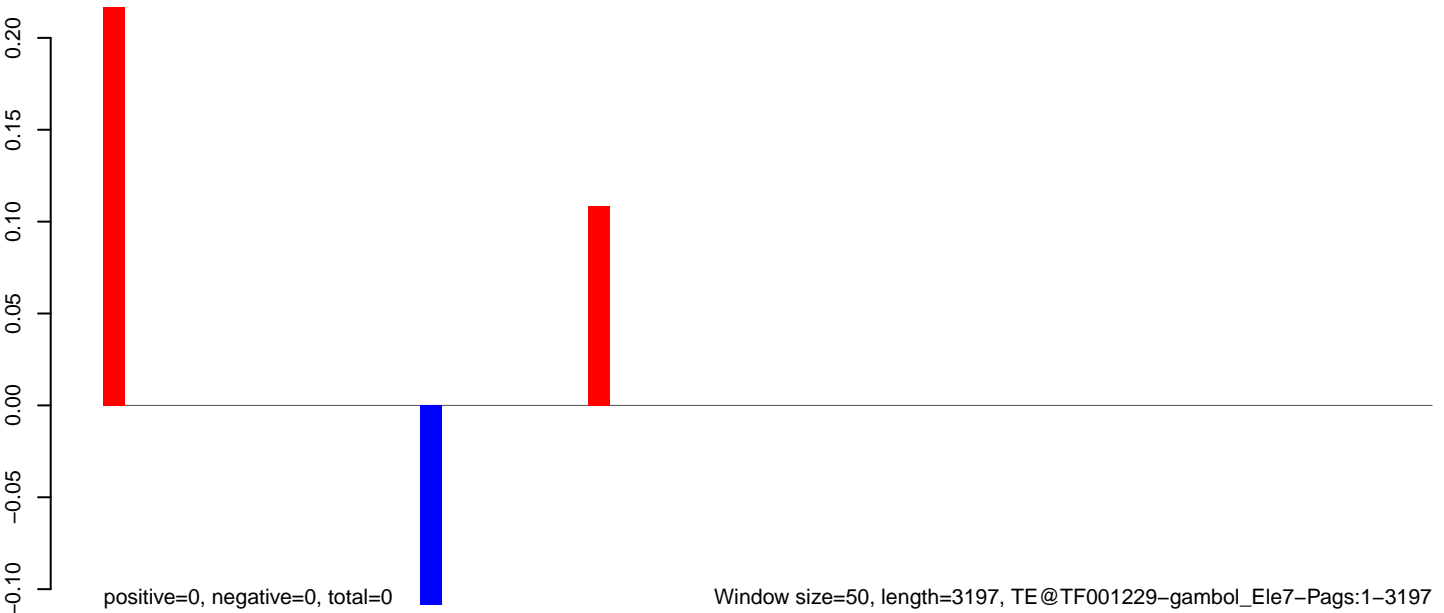
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



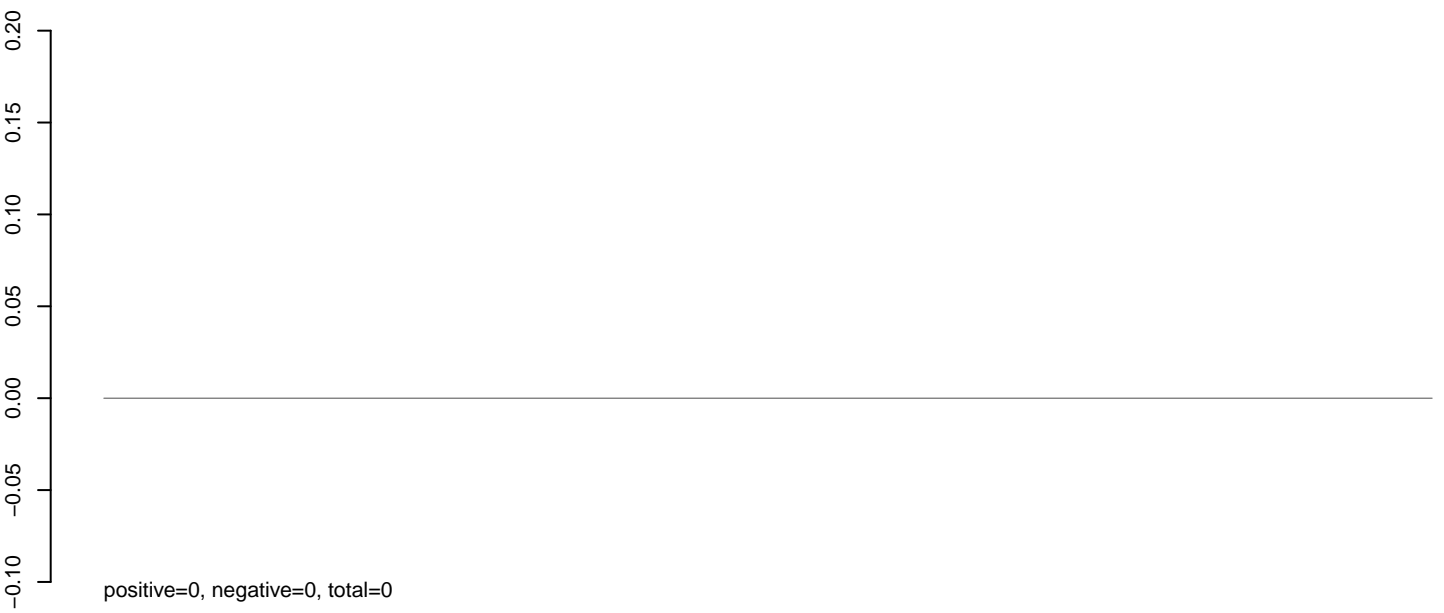
AnGam_Sua5b_cells.rep



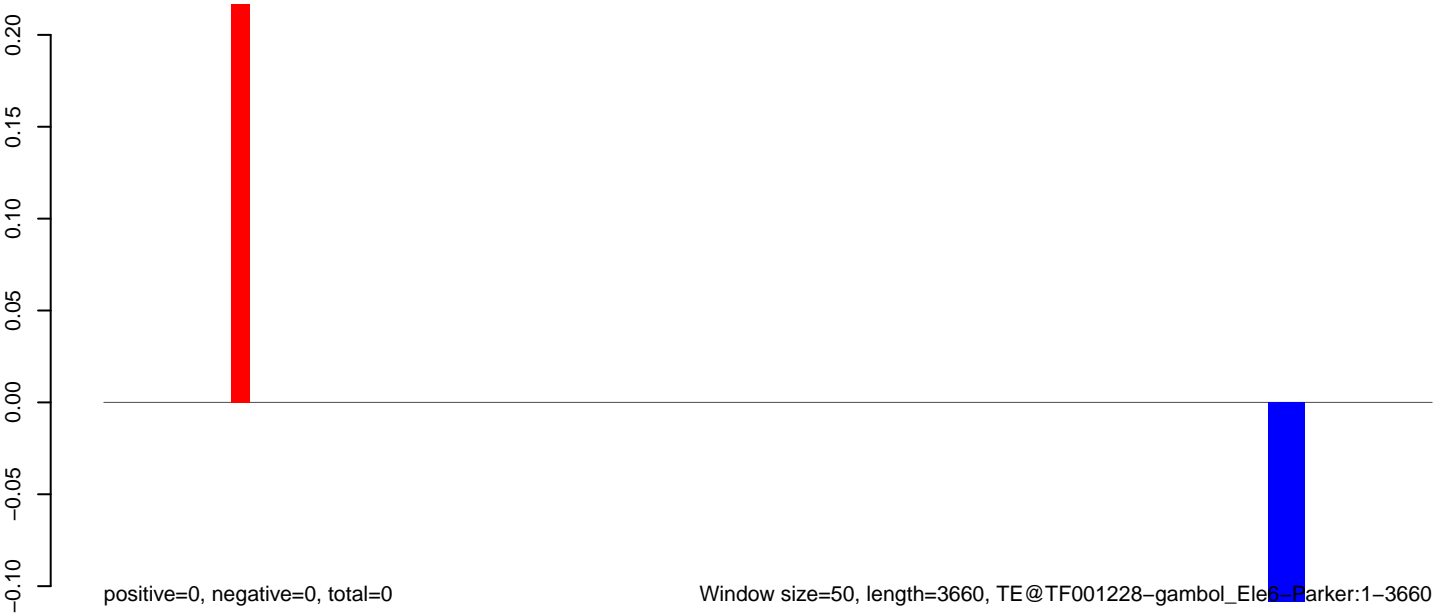
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



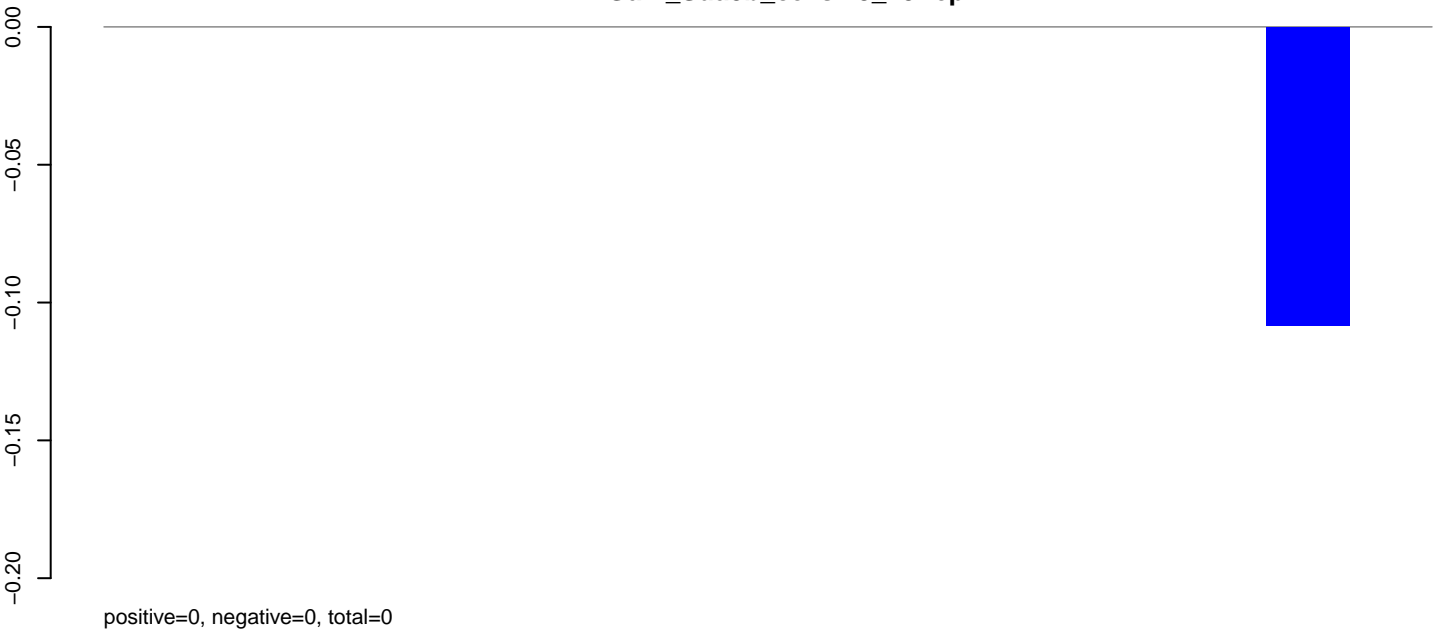
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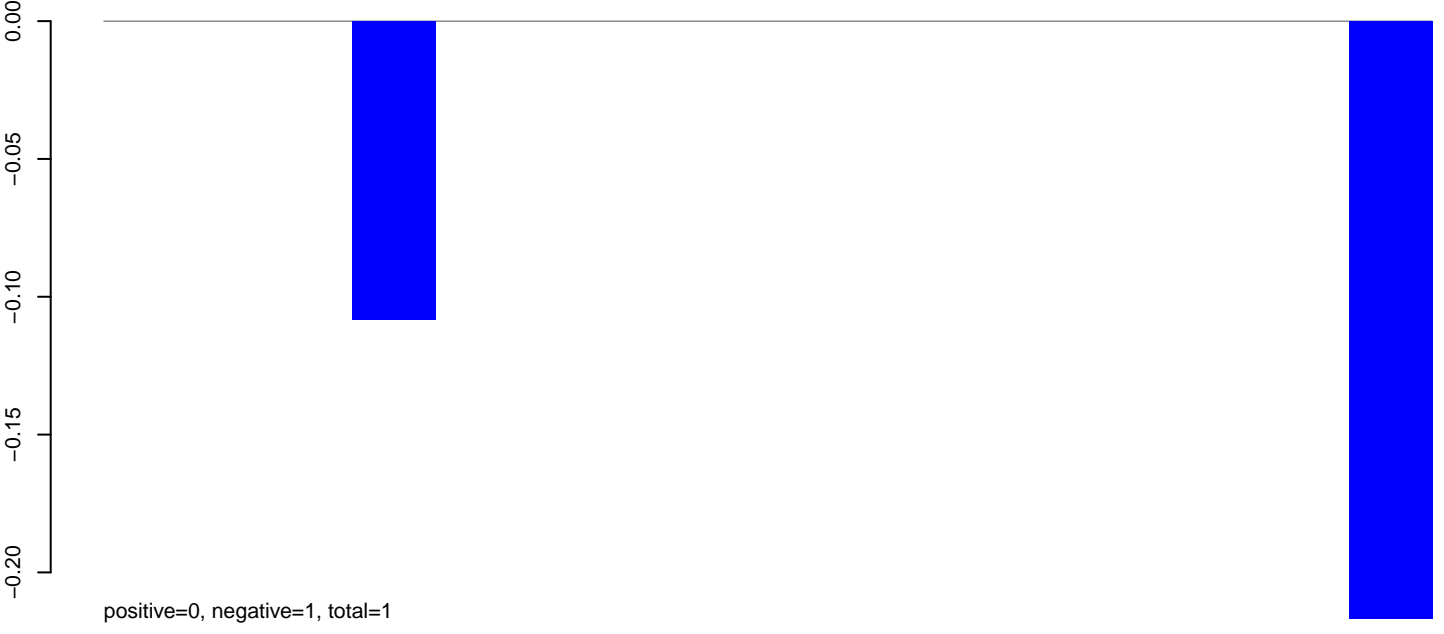
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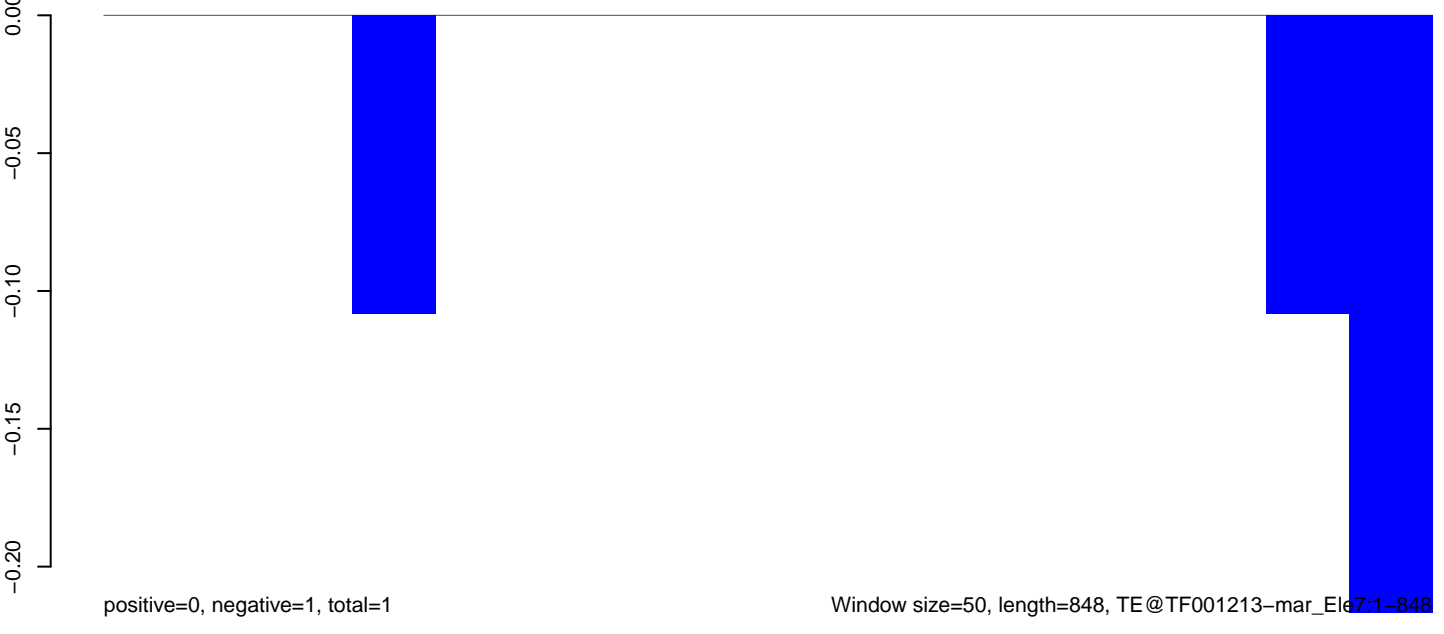
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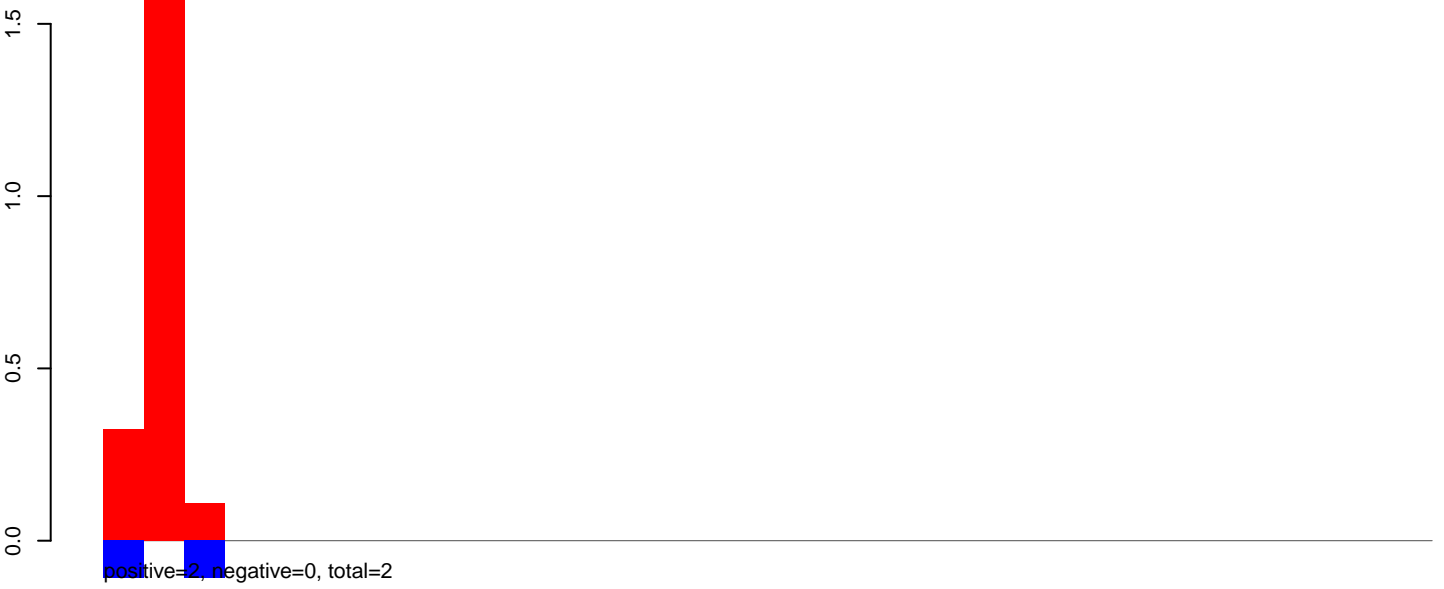
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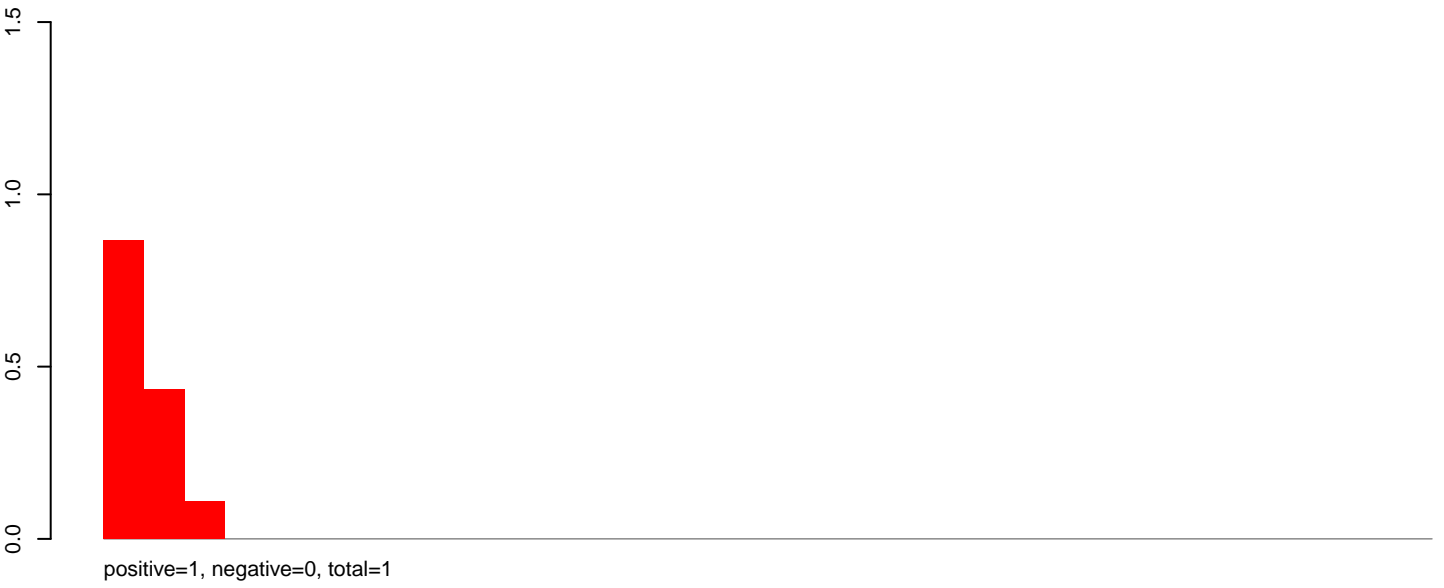
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AnGam_Sua5b_cells.18_23.rep



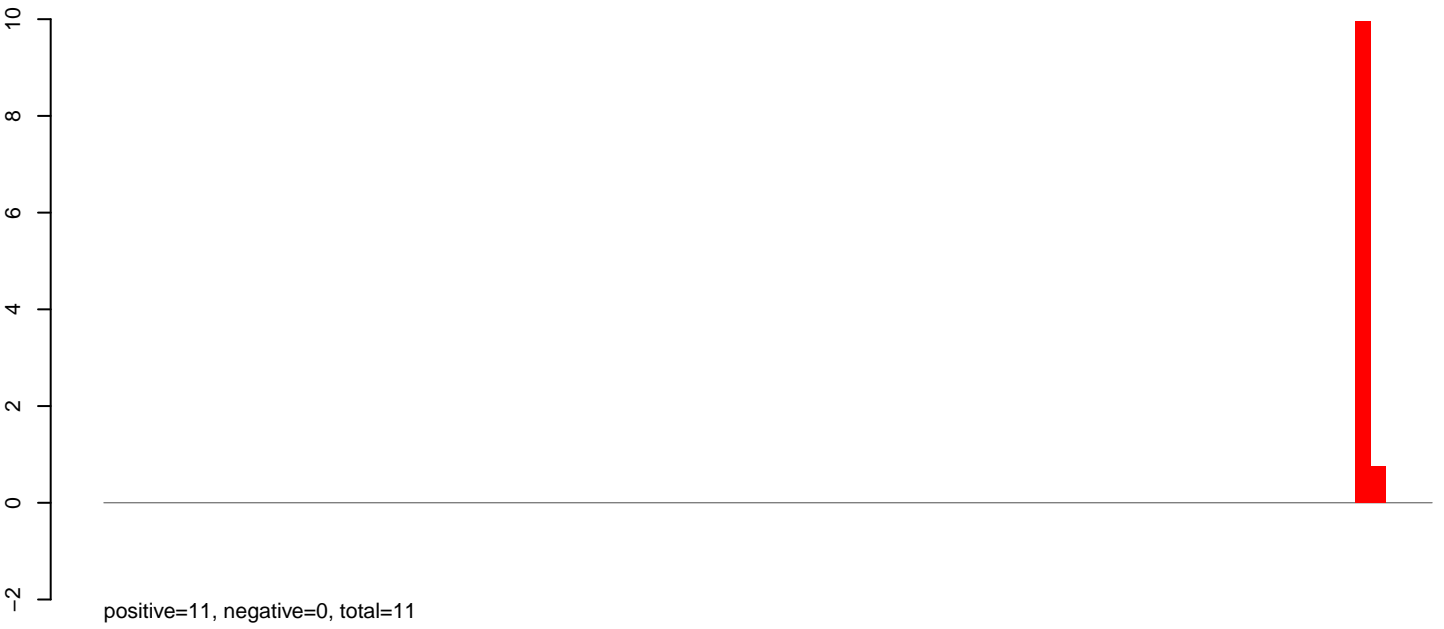
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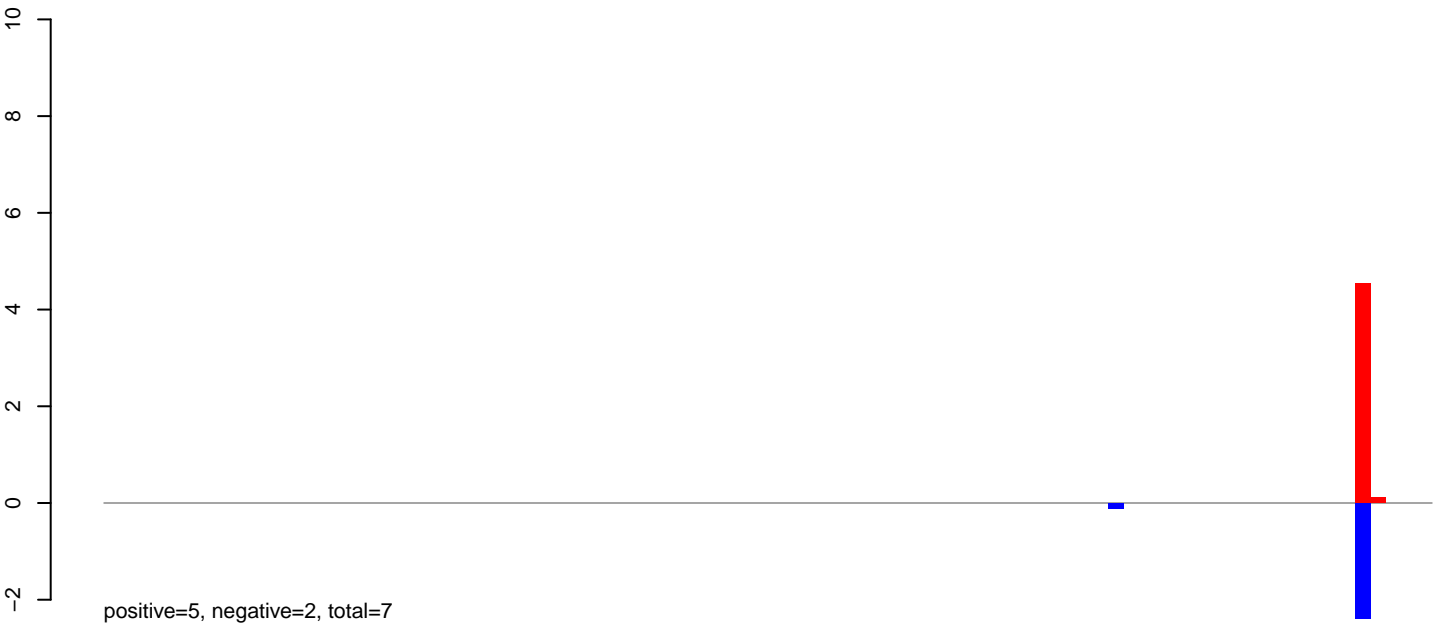
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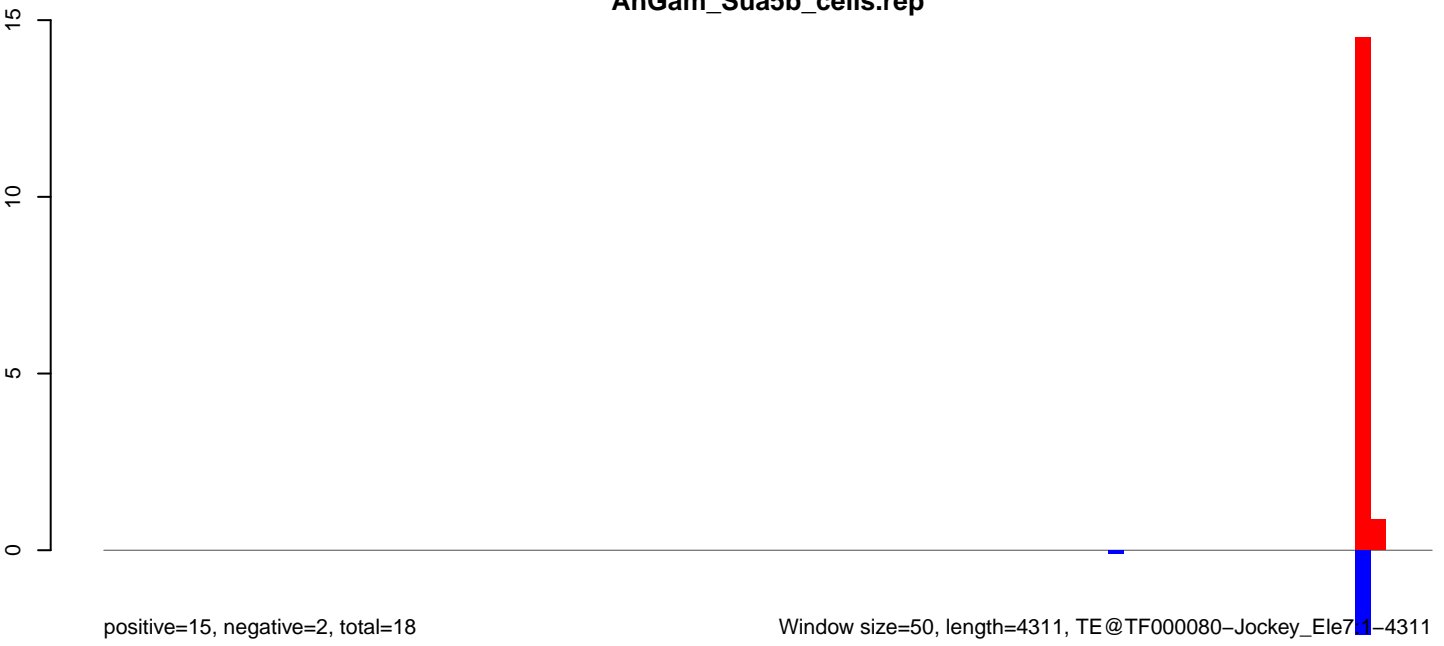
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AnGam_Sua5b_cells.24_35.rep



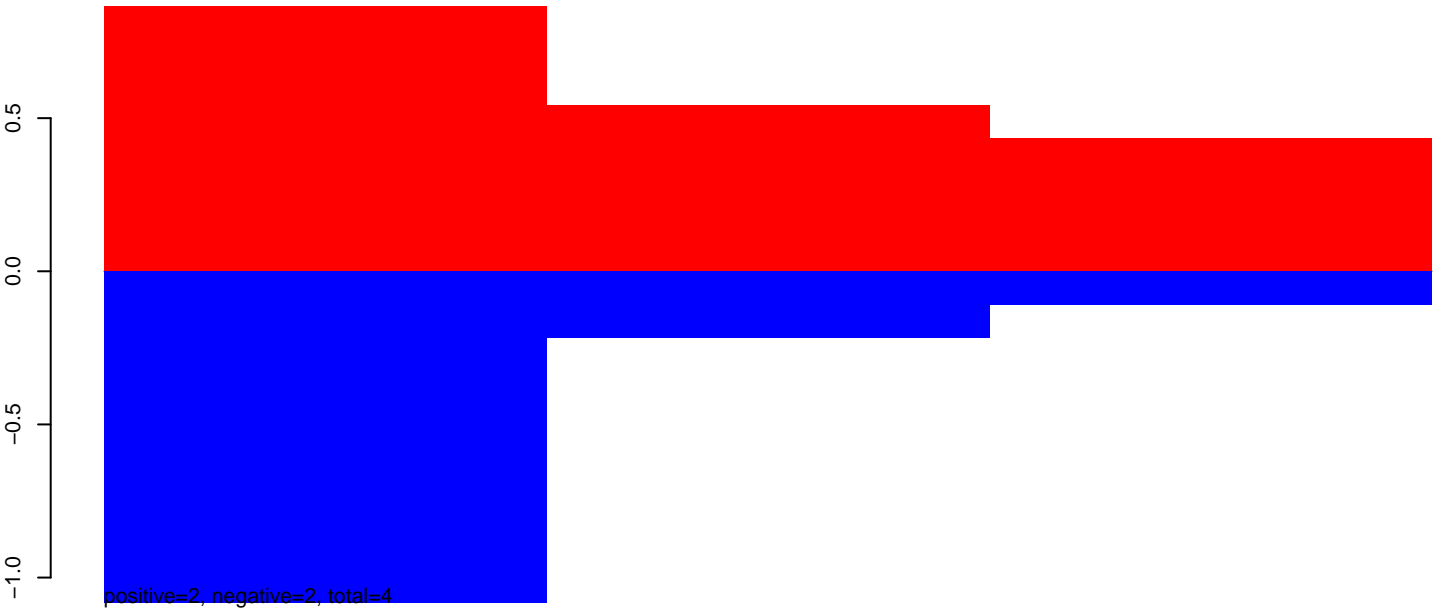
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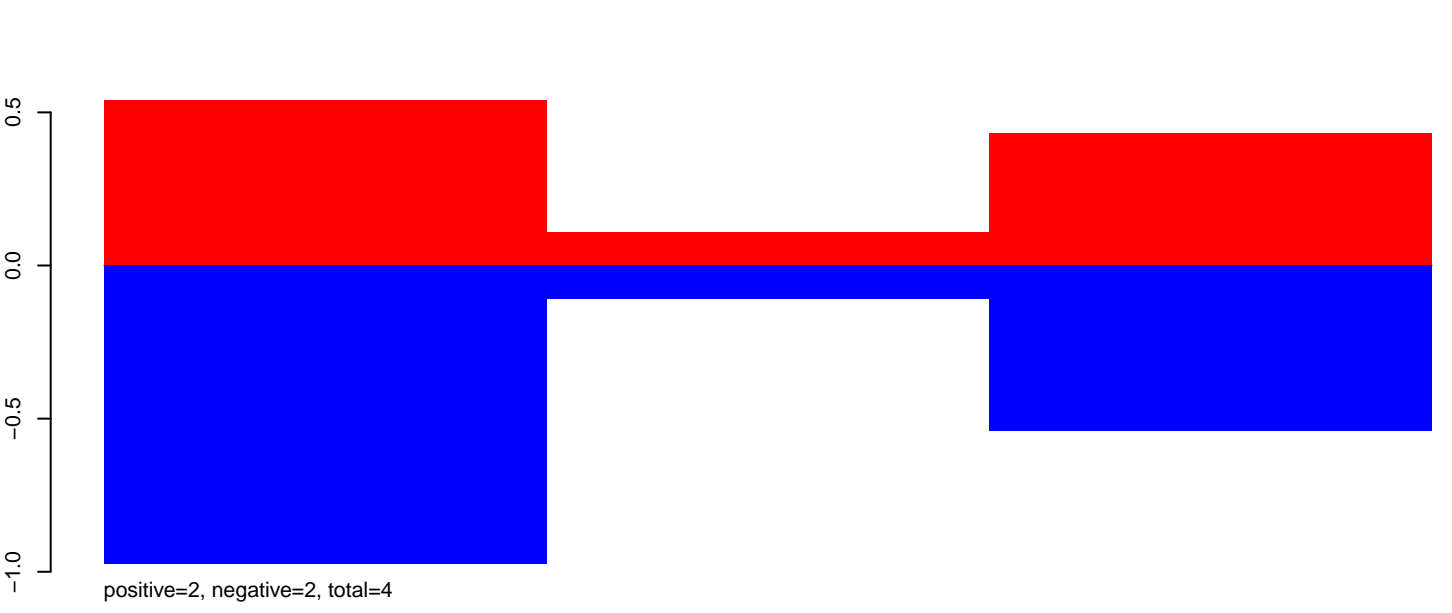
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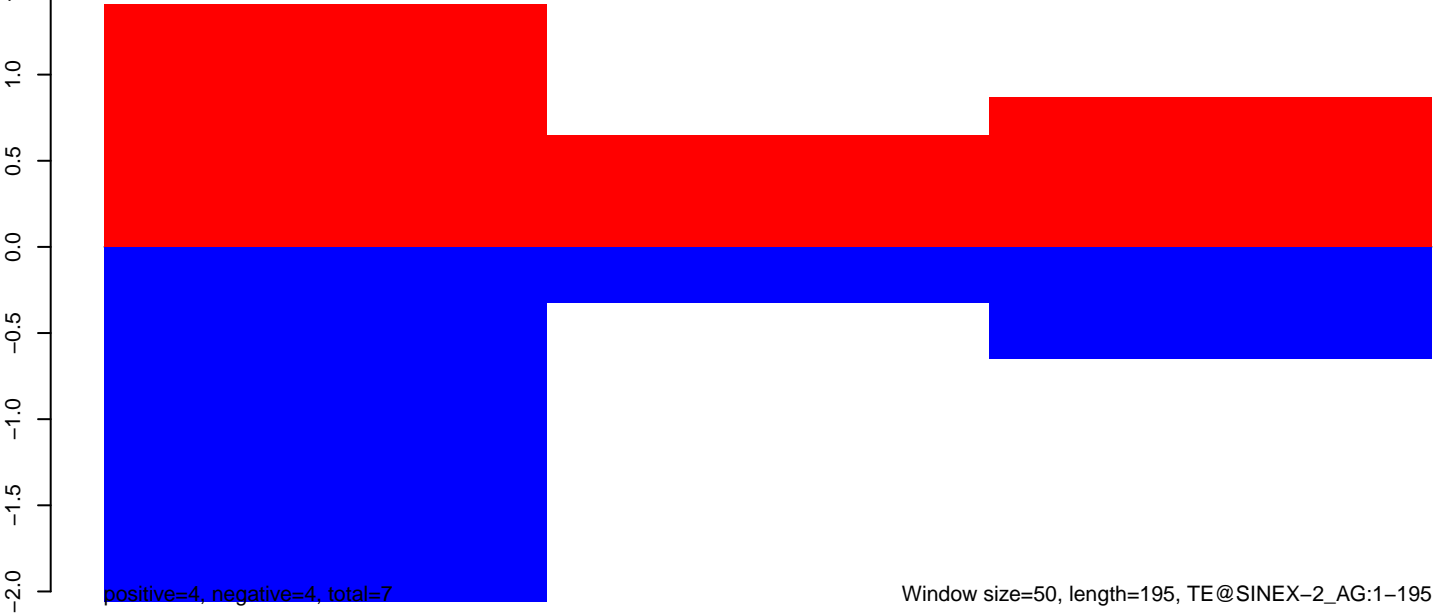
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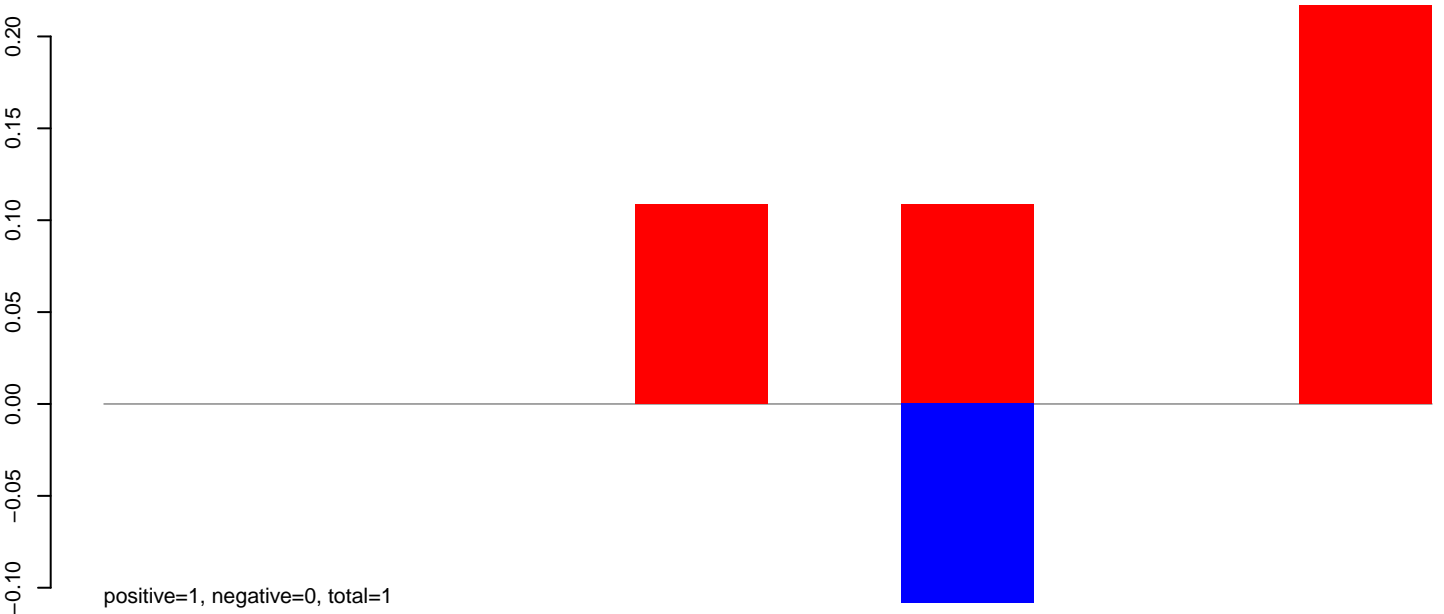
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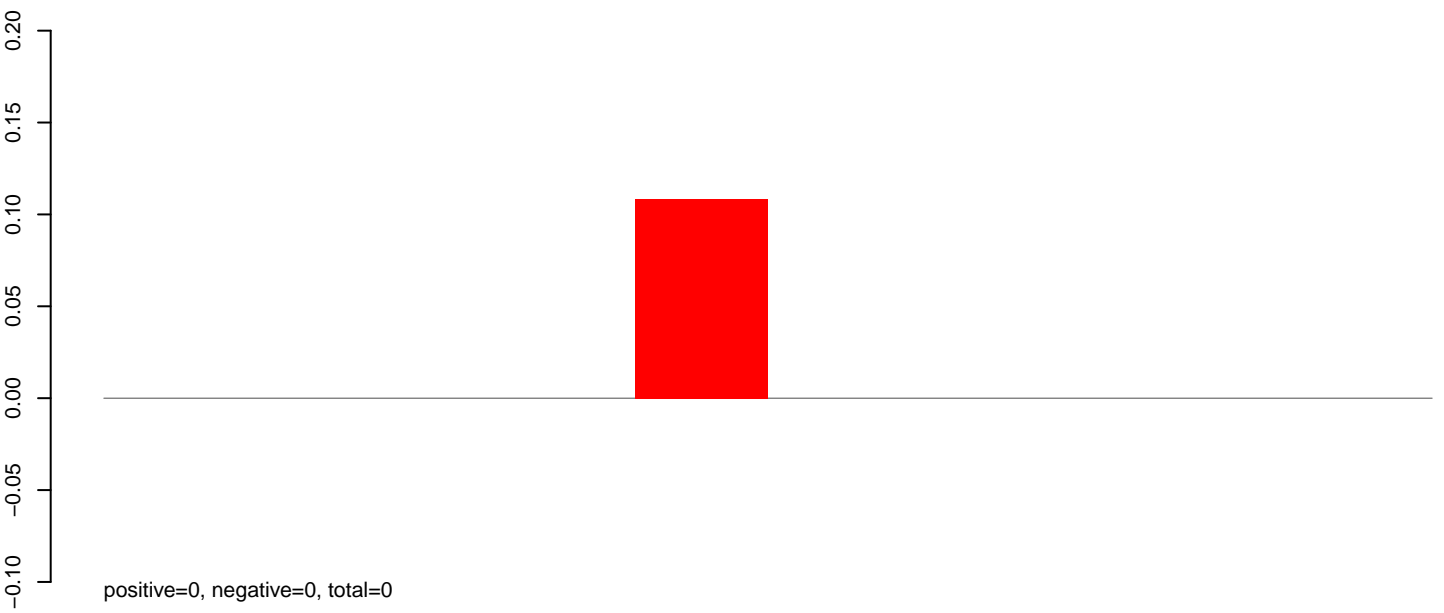
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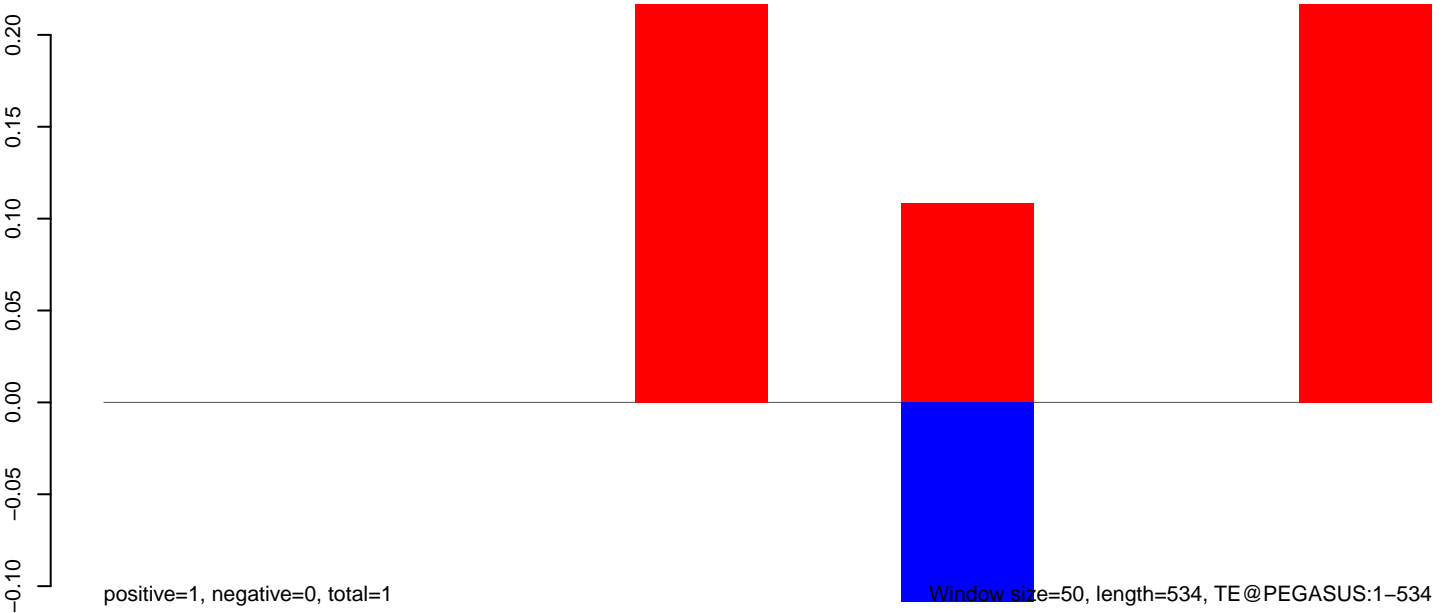
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AnGam_Sua5b_cells.24_35.rep



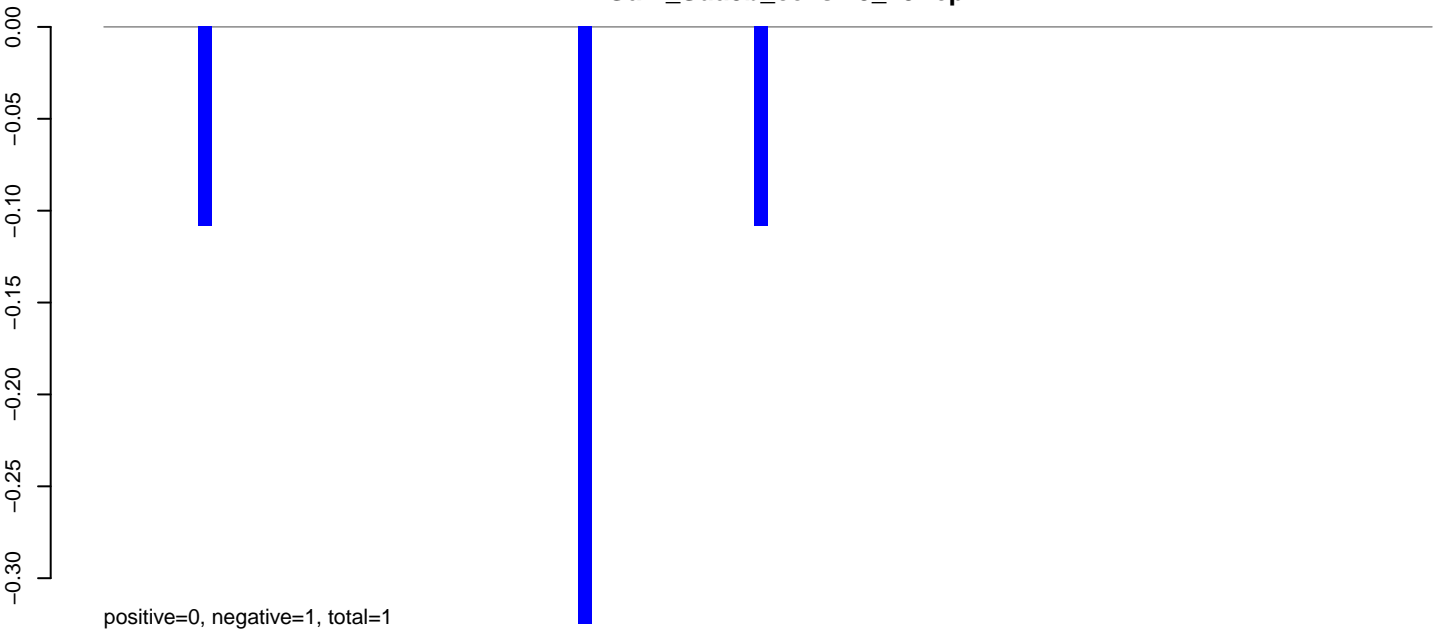
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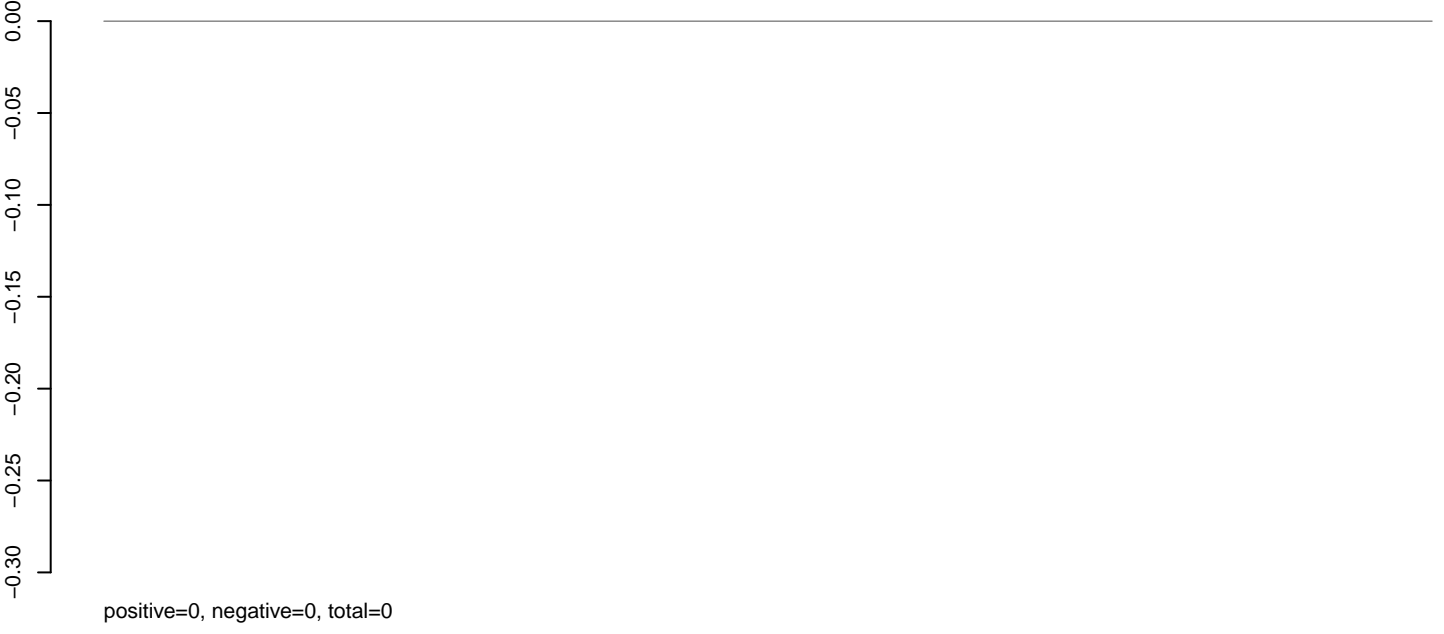
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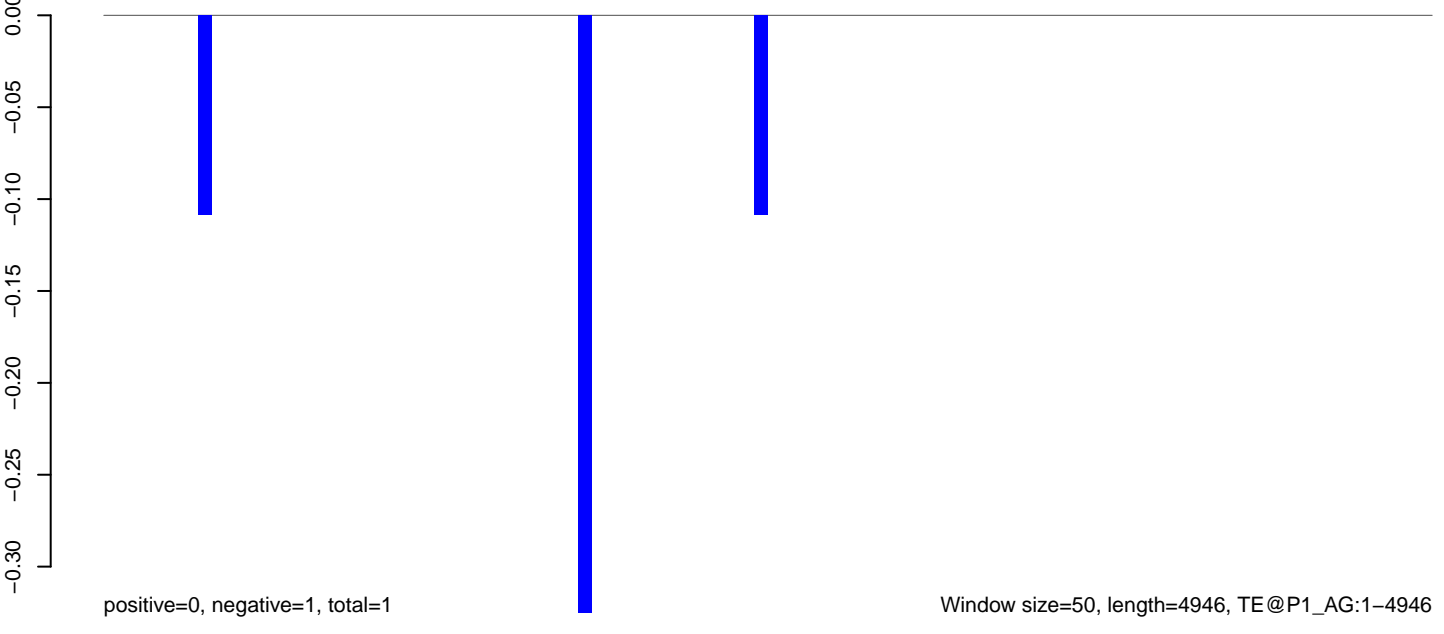
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AnGam_Sua5b_cells.24_35.rep



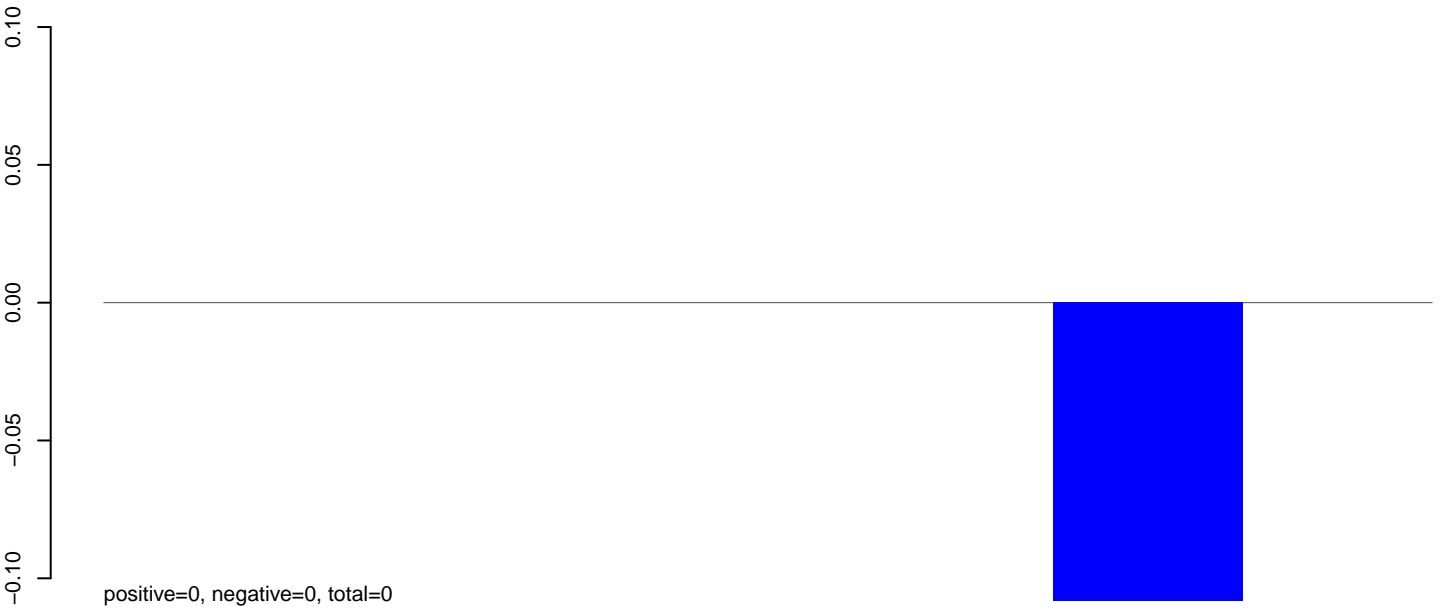
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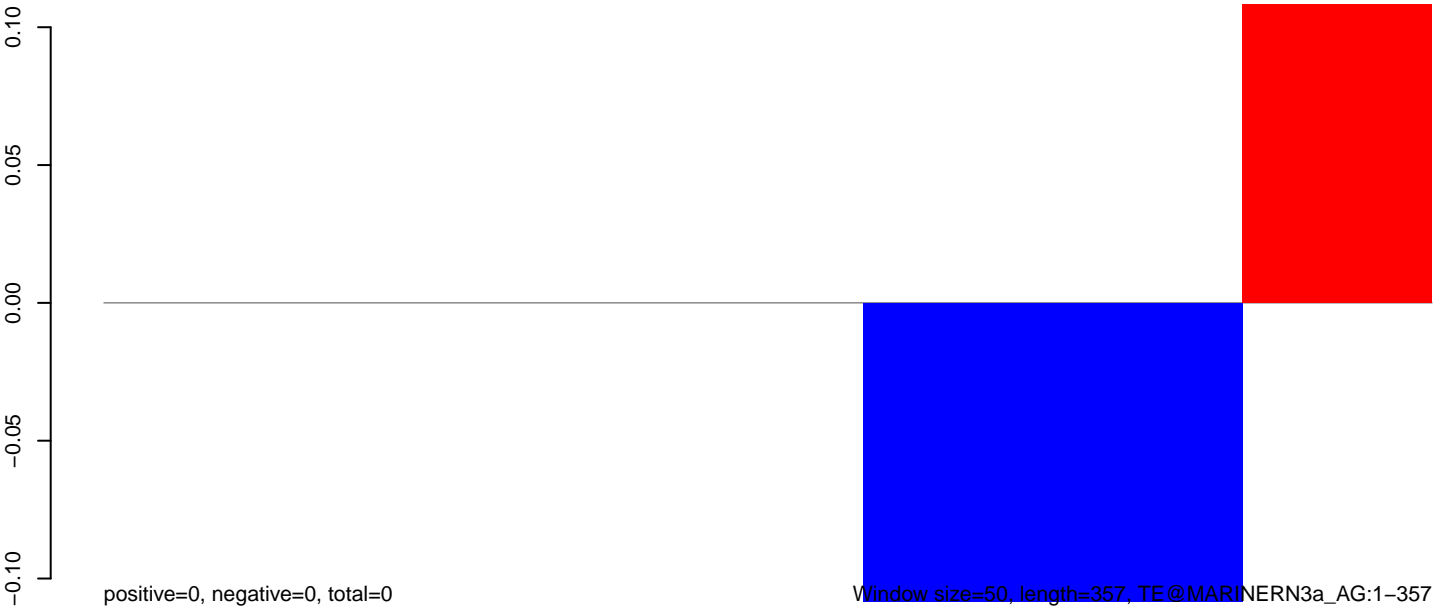
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AnGam_Sua5b_cells.24_35.rep



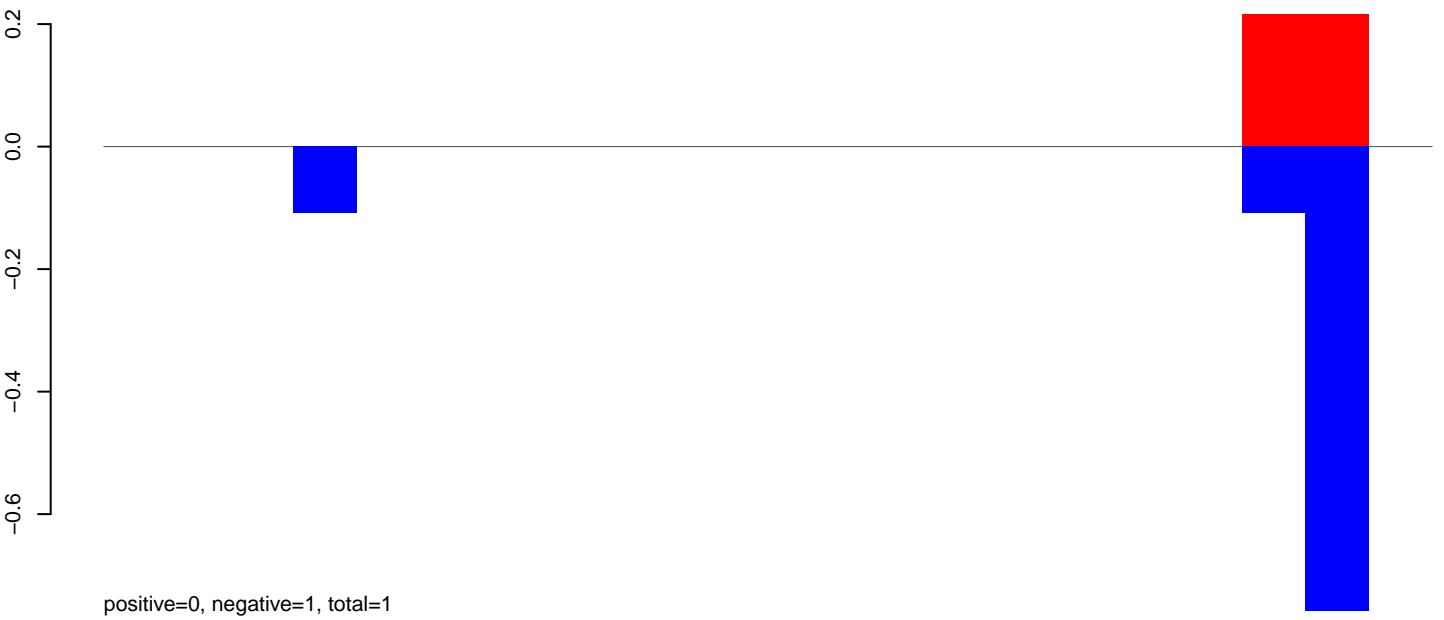
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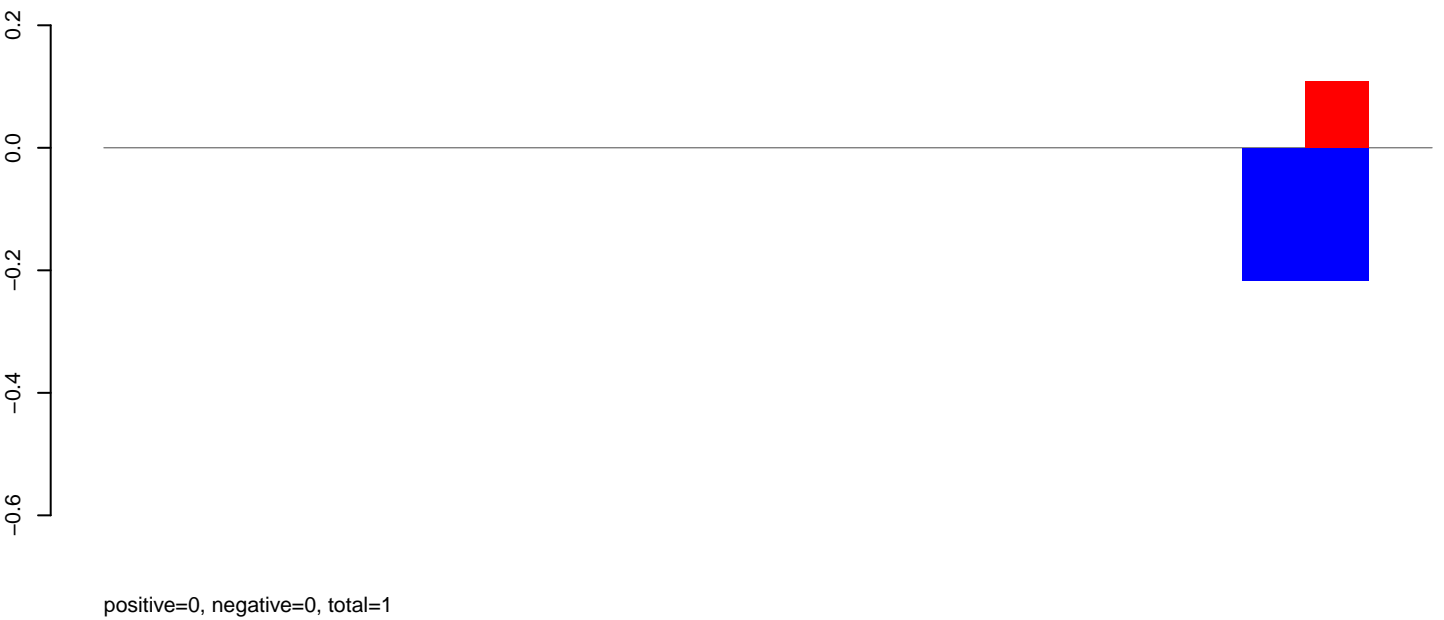
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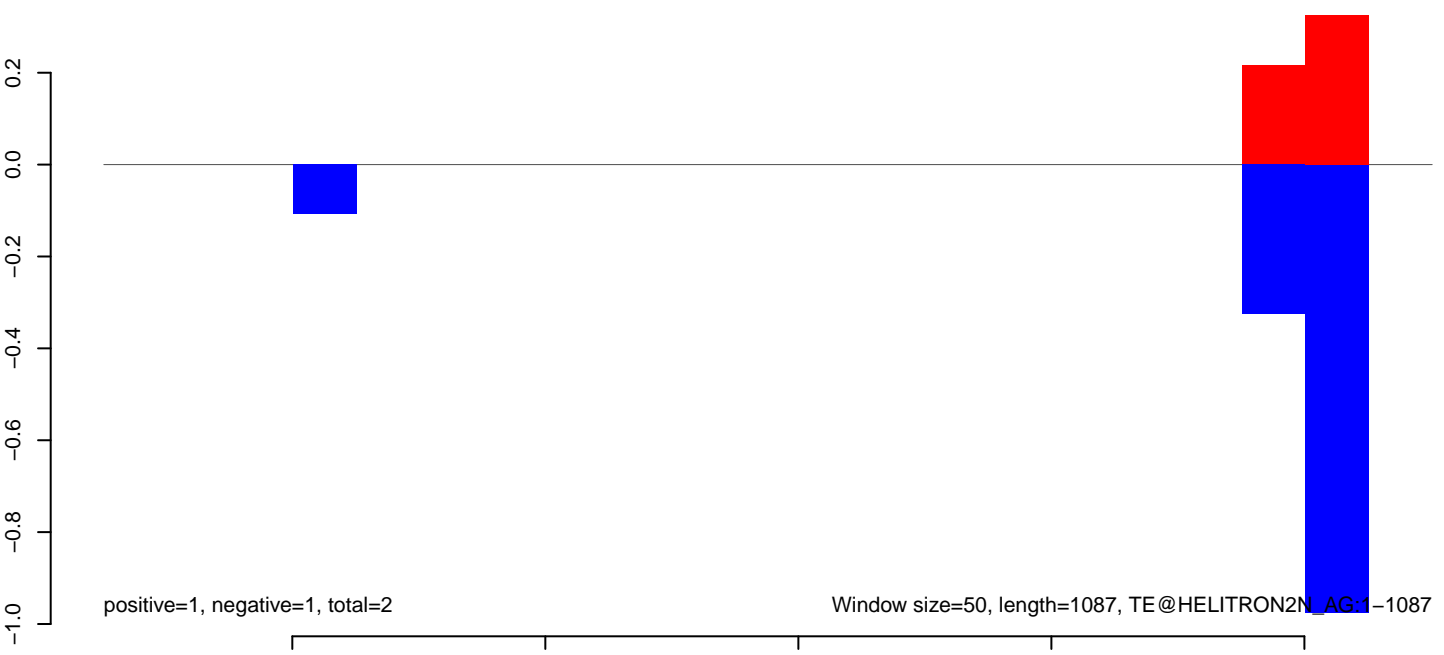
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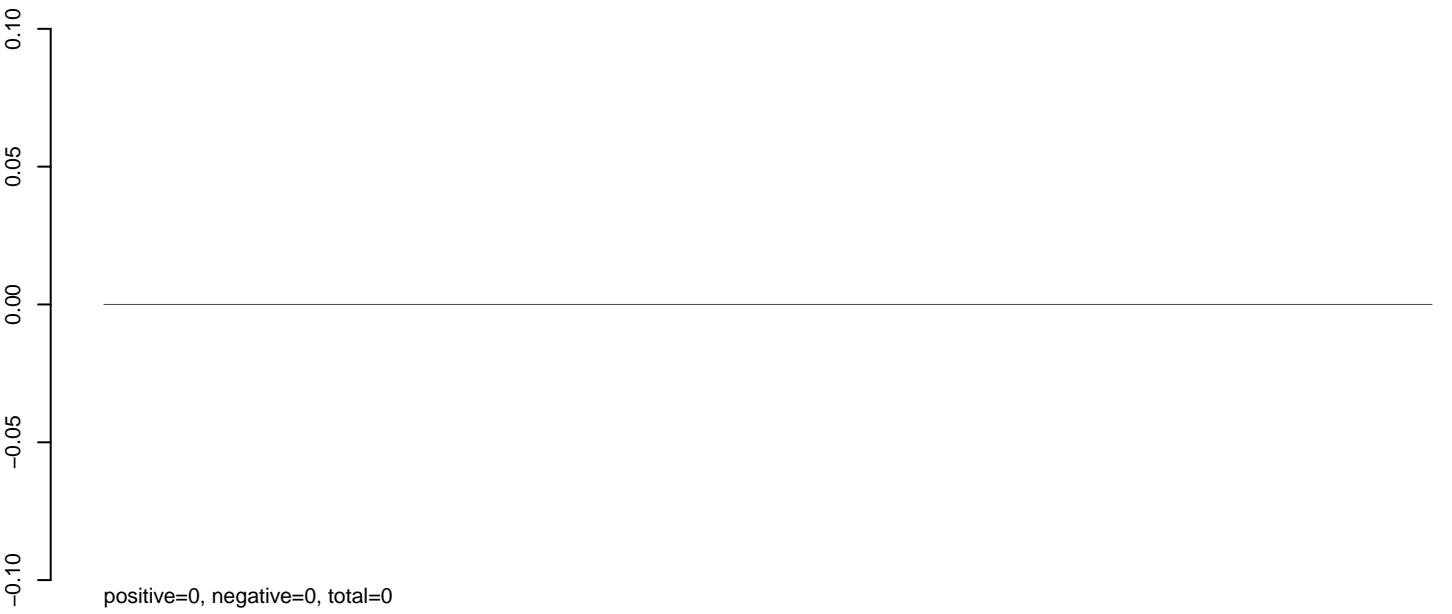
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AnGam_Sua5b_cells.rep



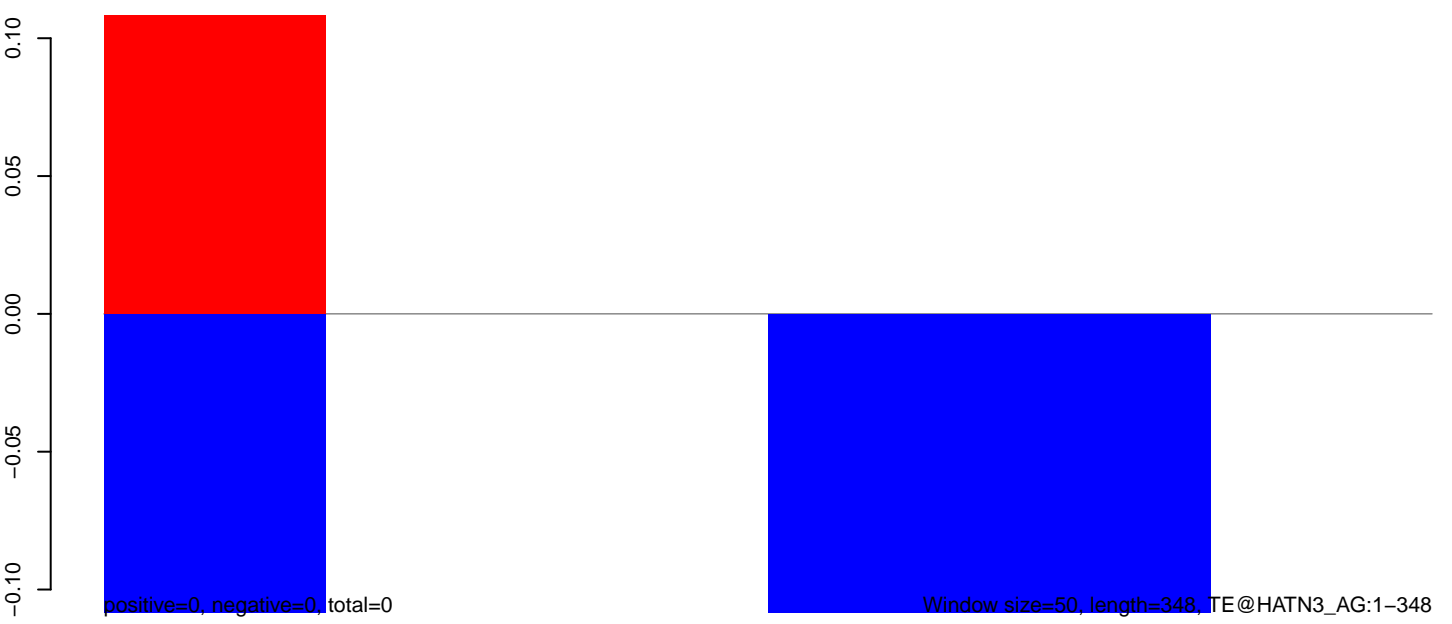
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AnGam_Sua5b_cells.24_35.rep

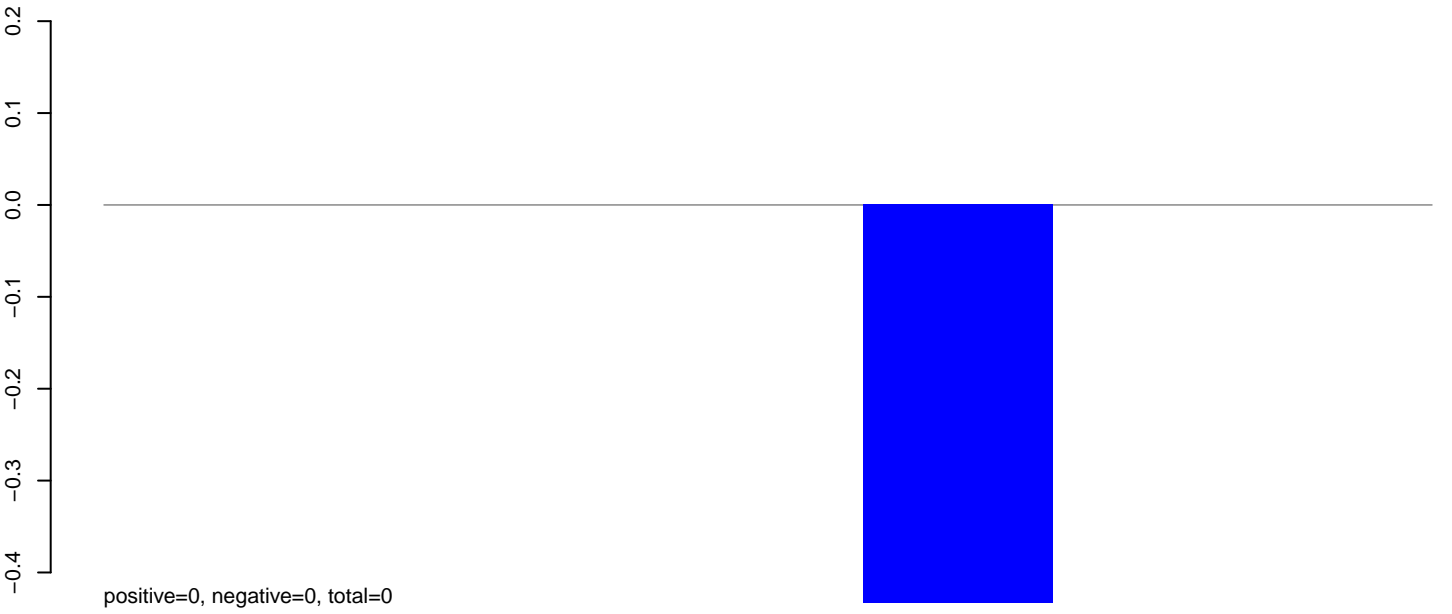


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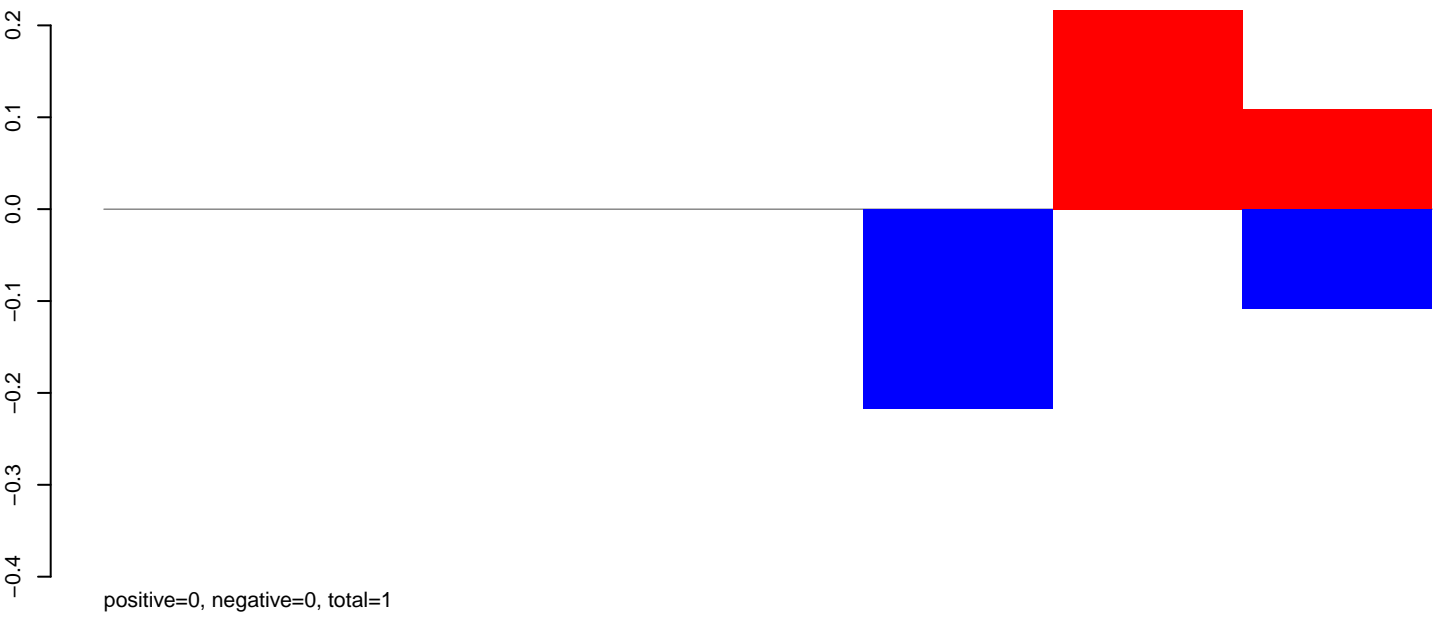


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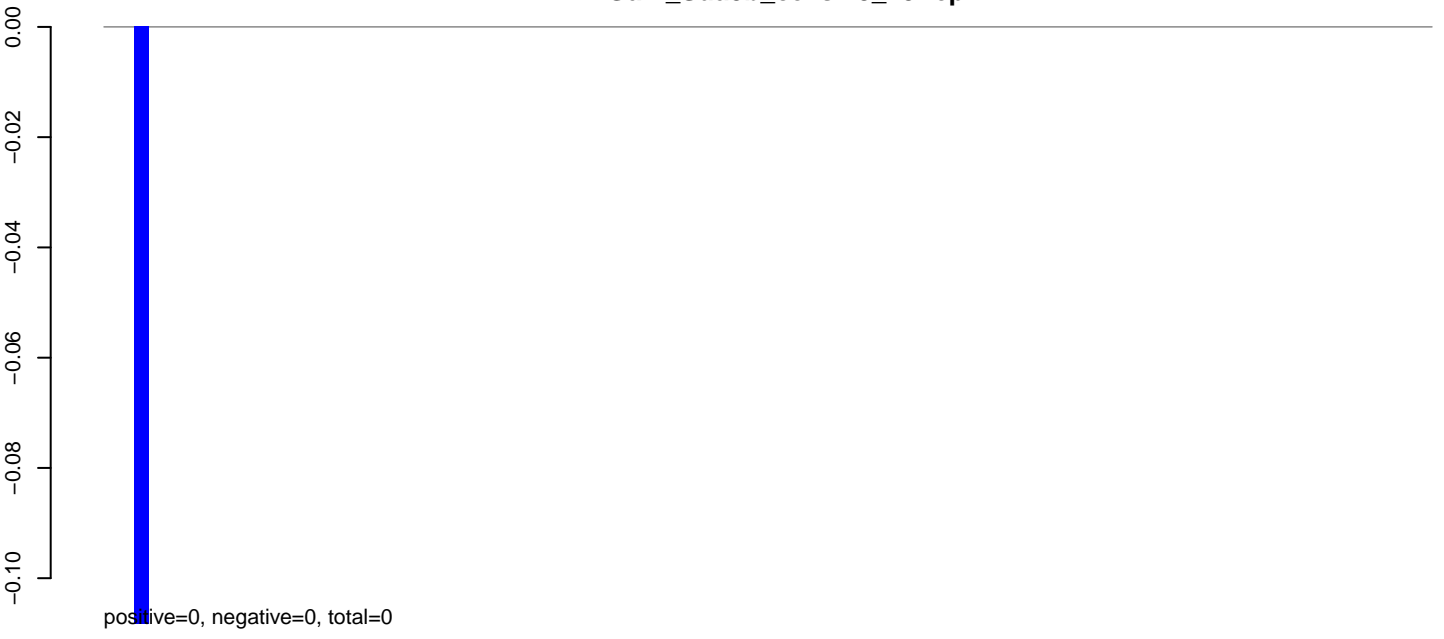
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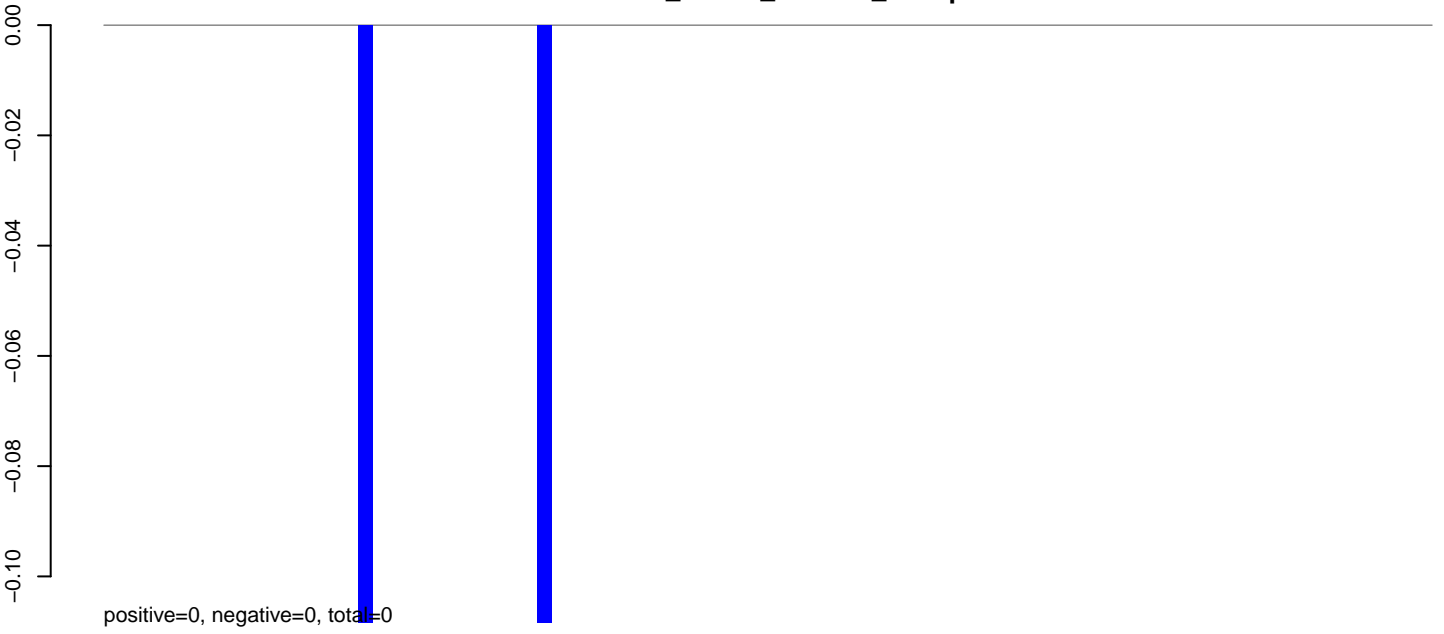
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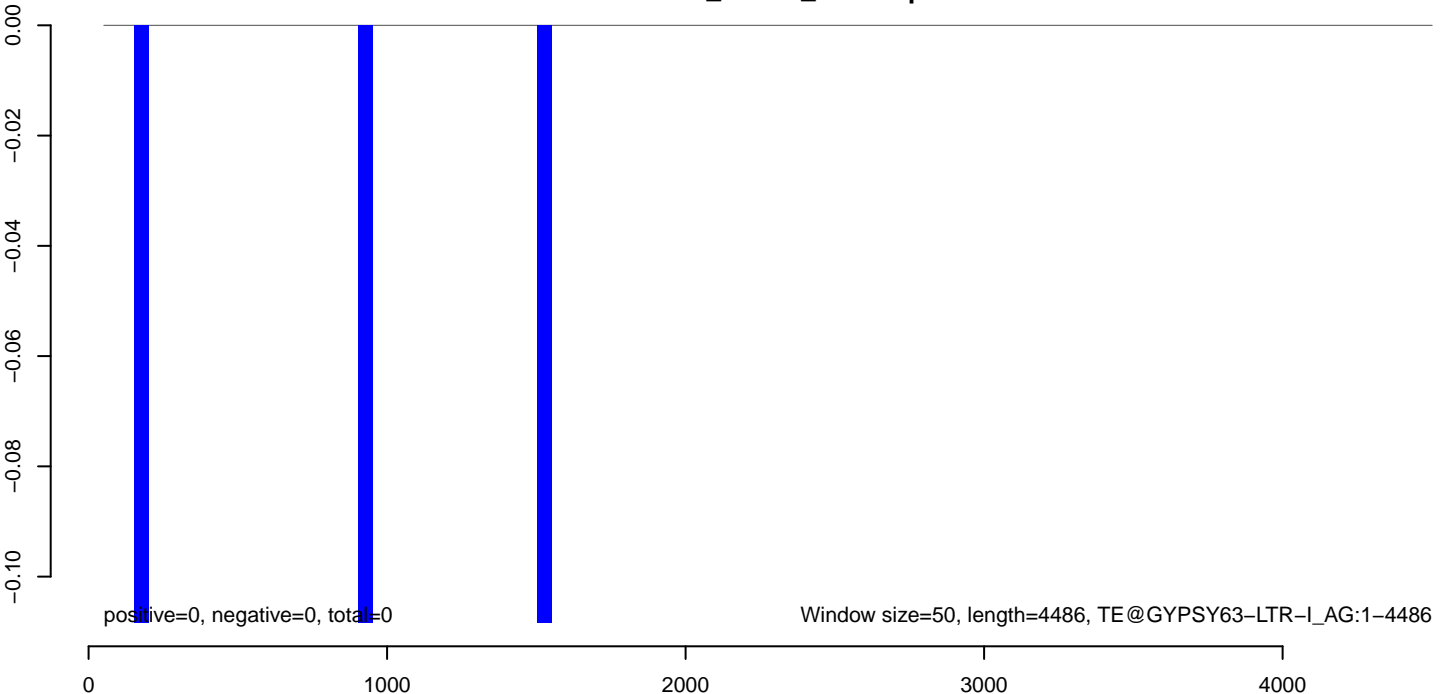
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AnGam_Sua5b_cells.24_35.rep



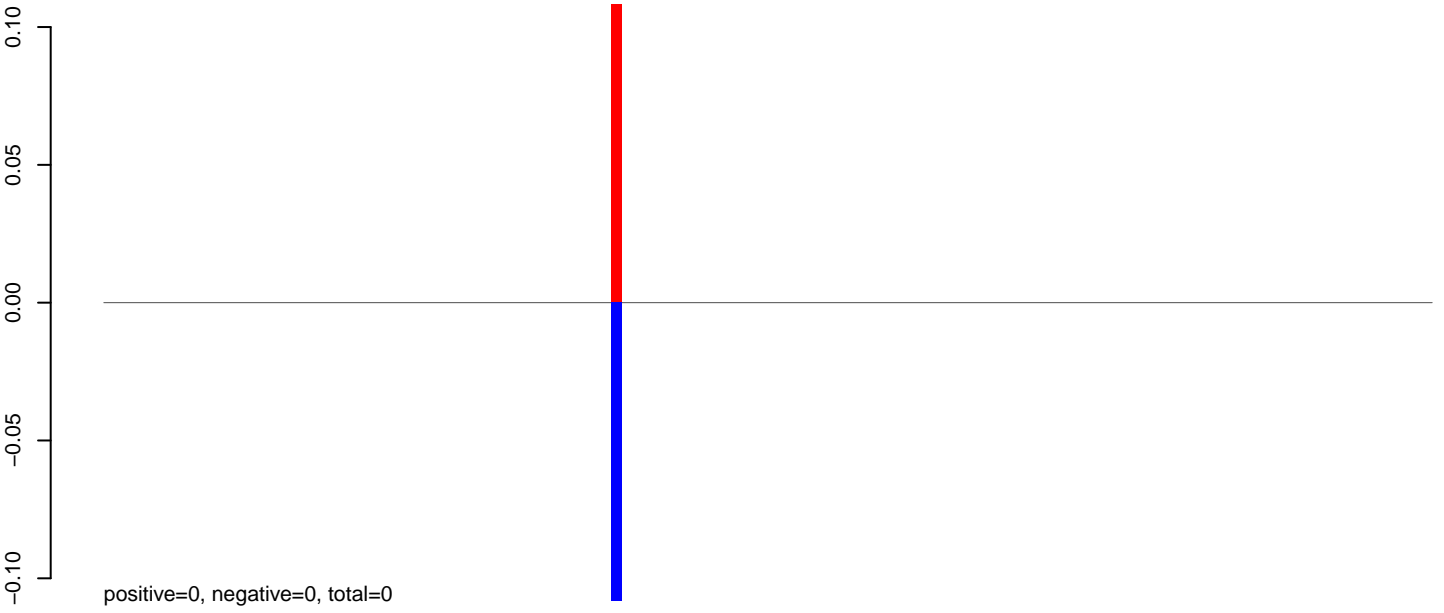
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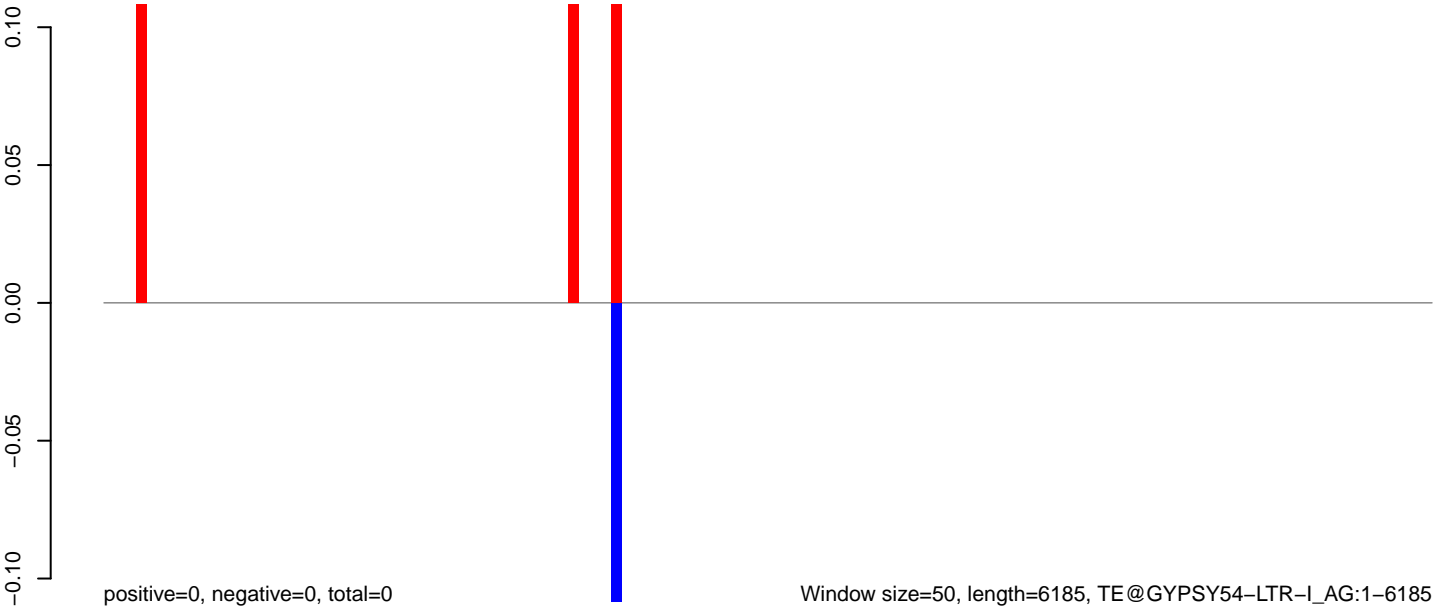
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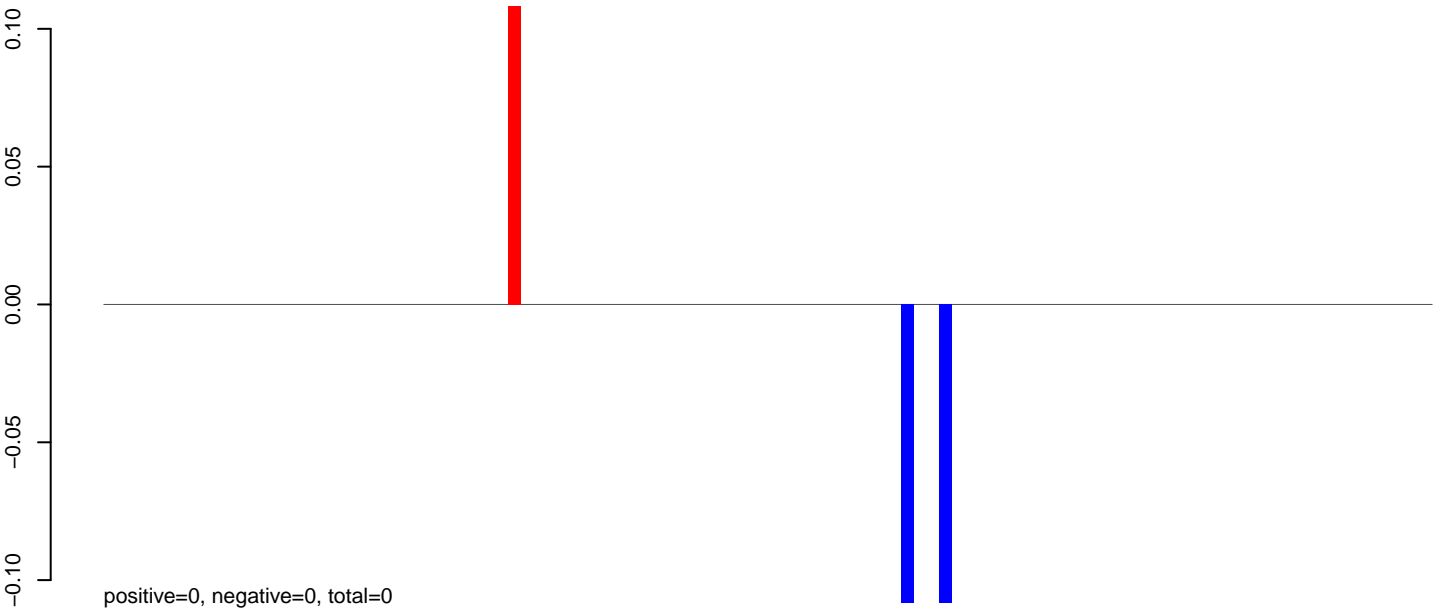
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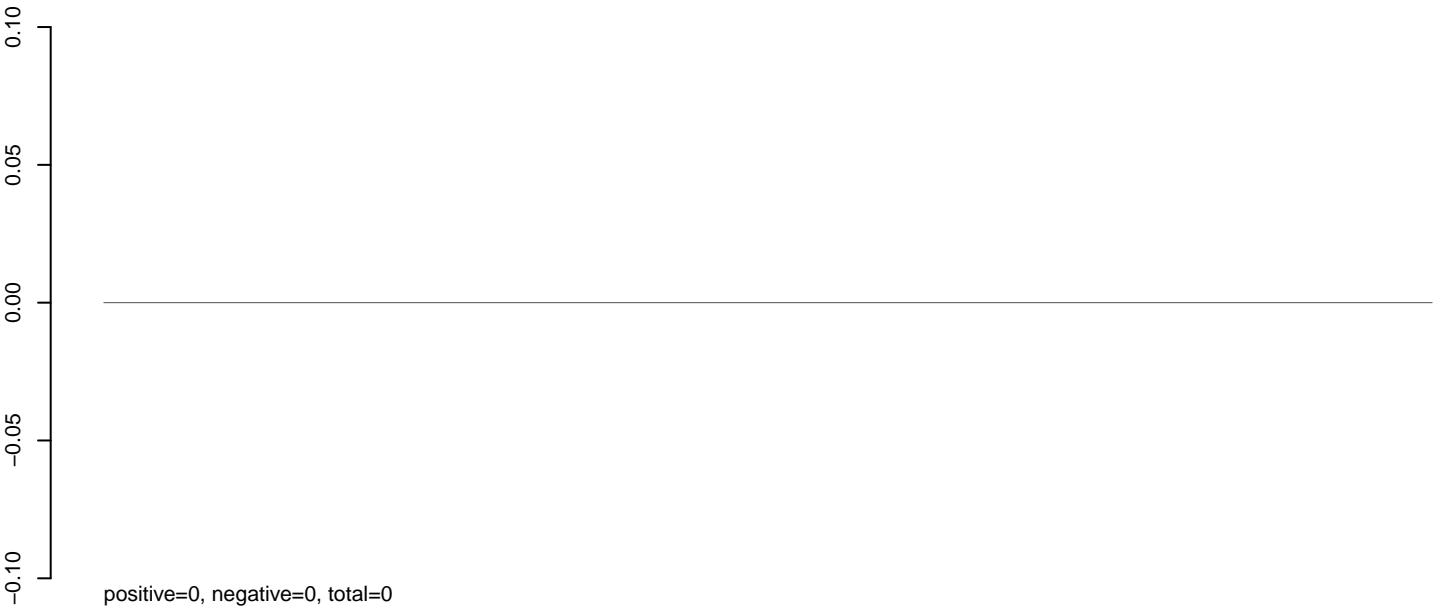
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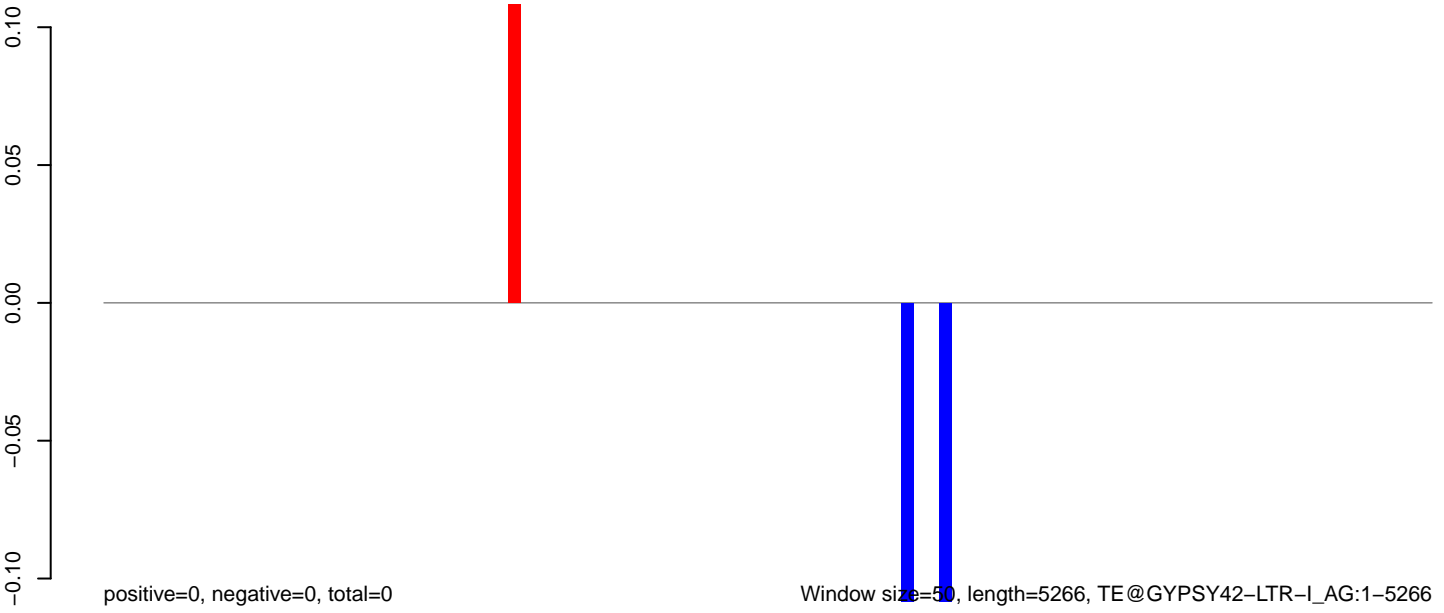
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AnGam_Sua5b_cells.24_35.rep

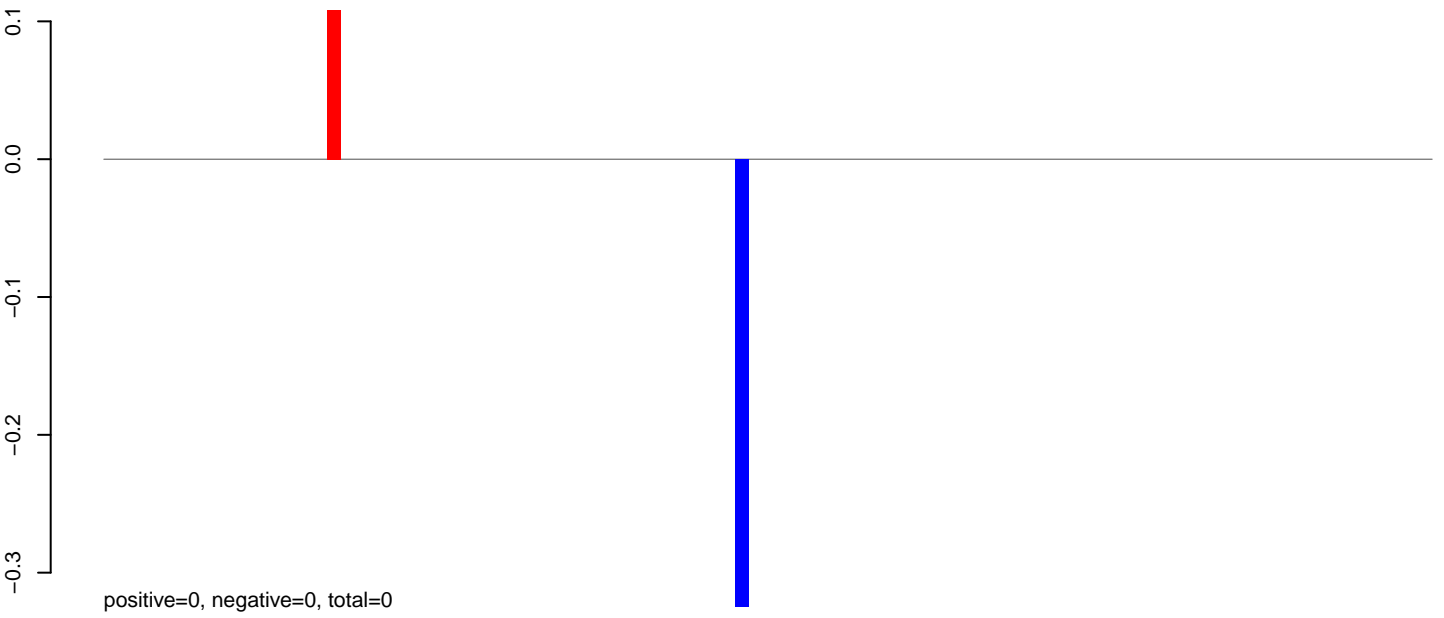


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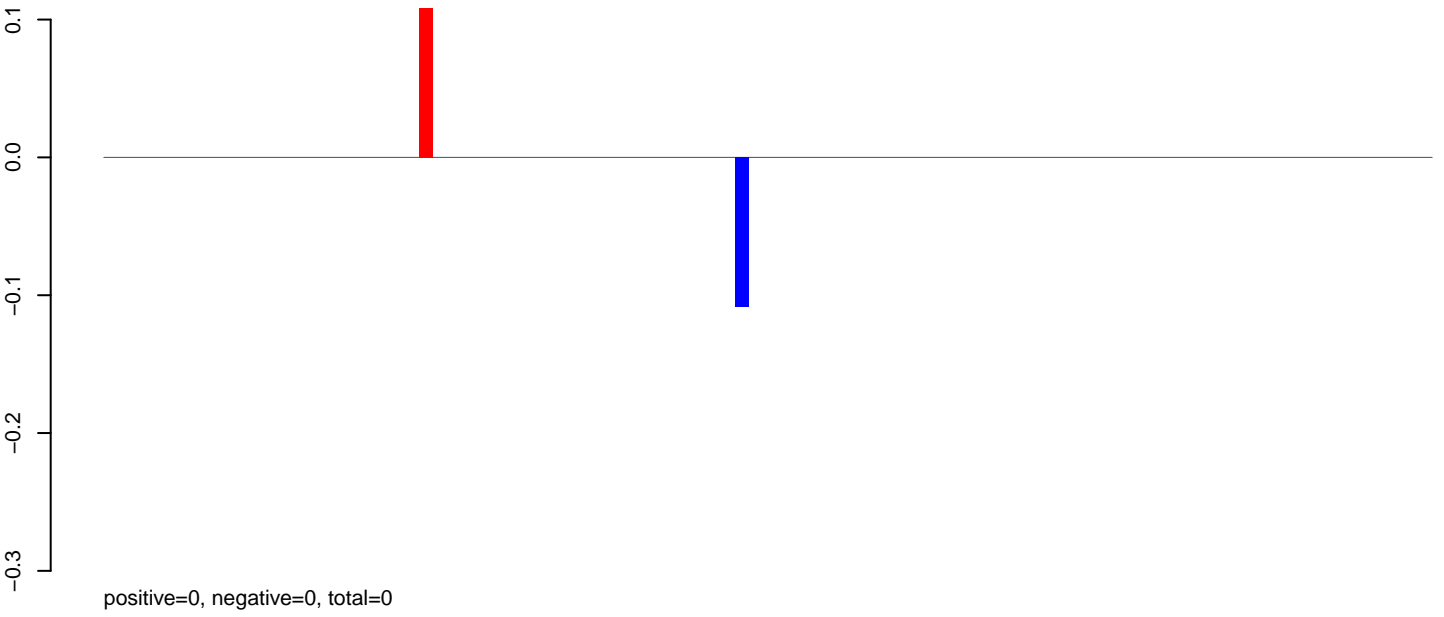


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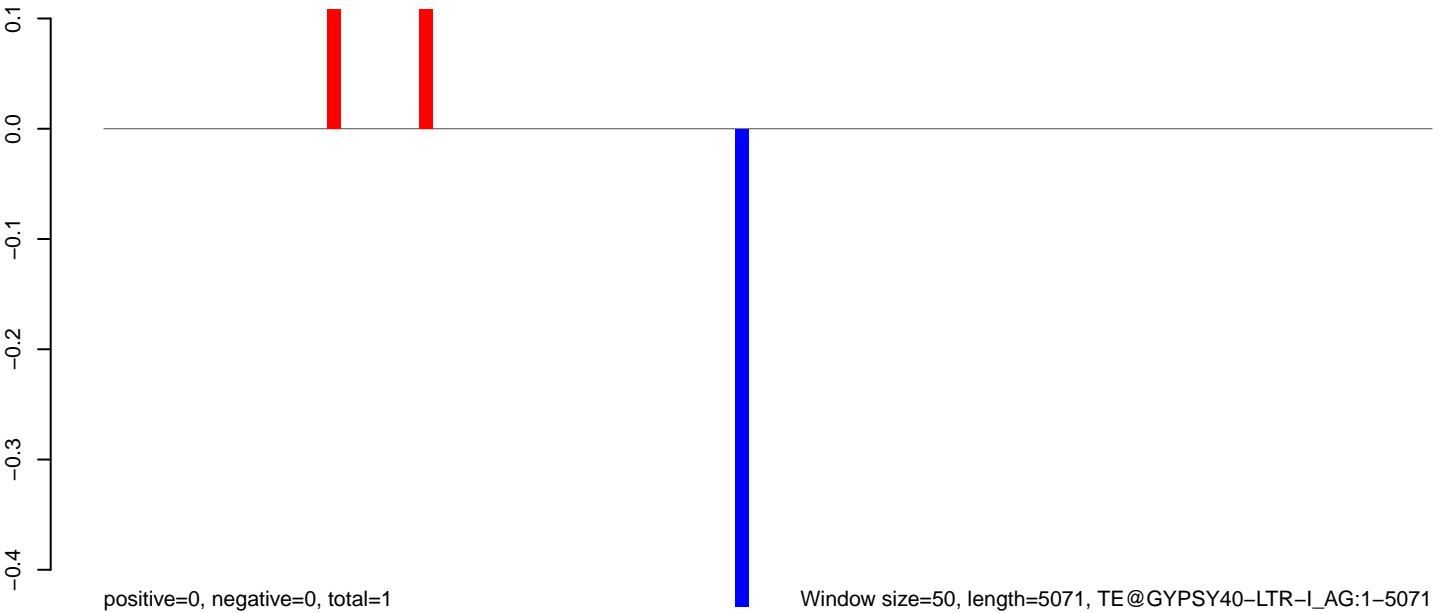
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AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

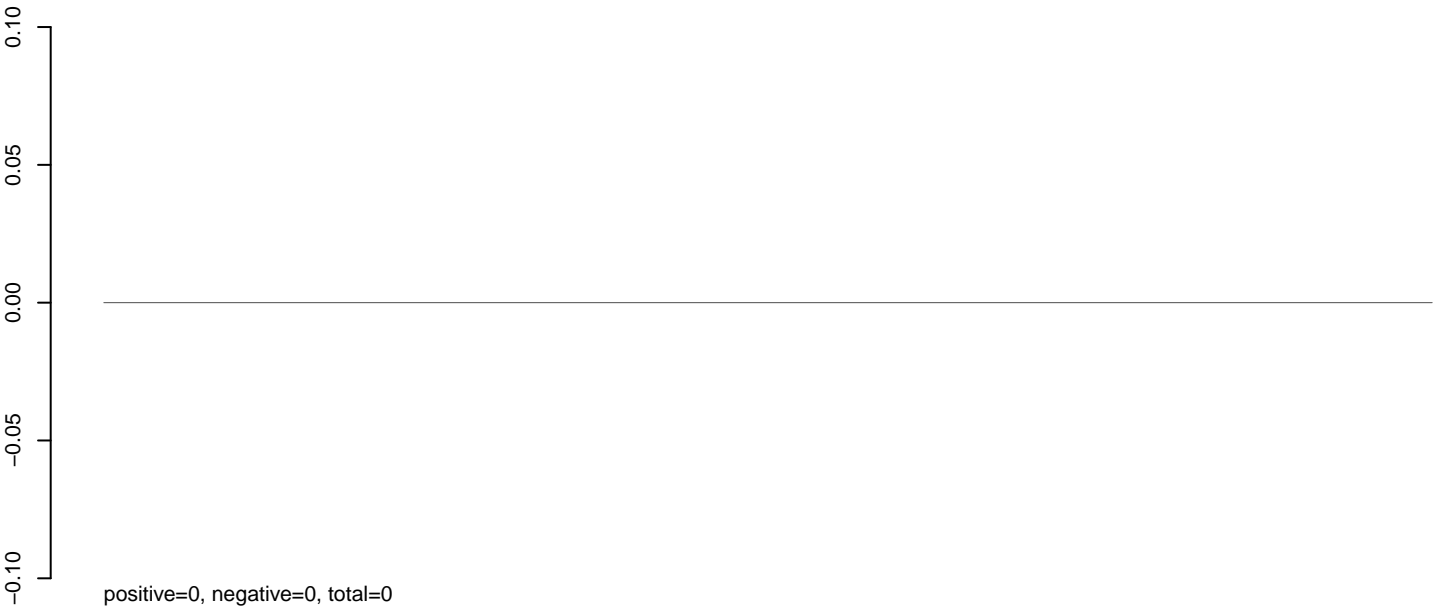


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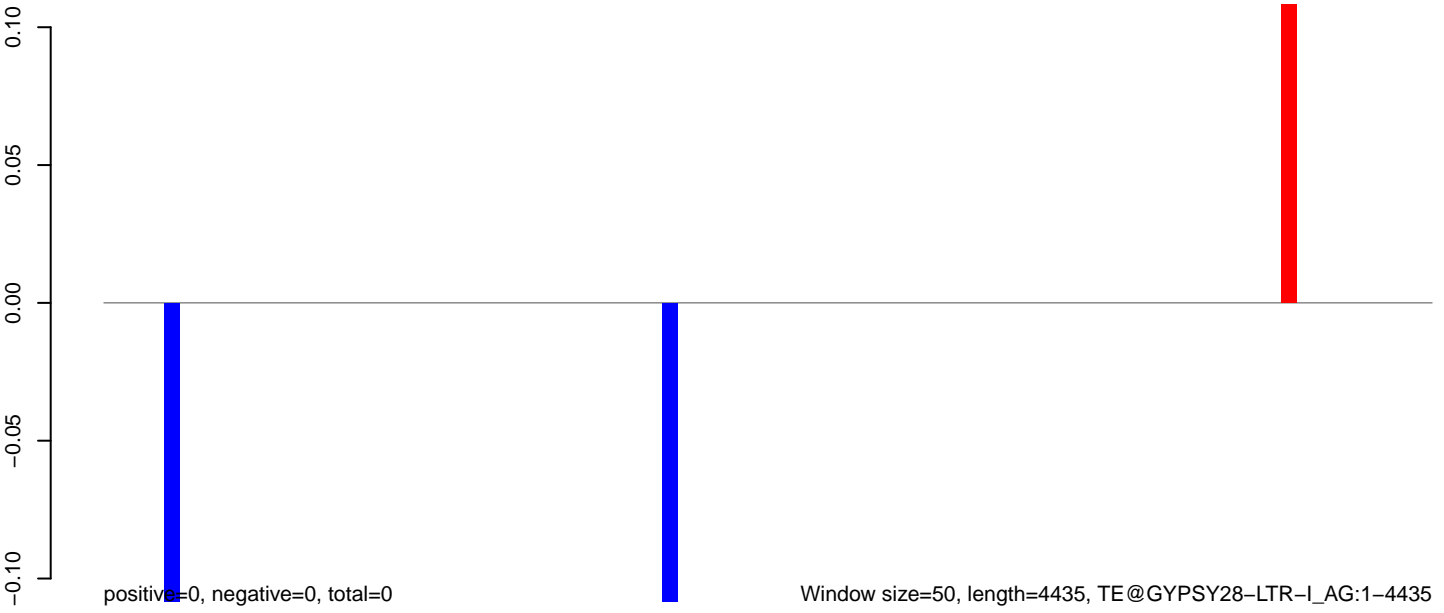
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AnGam_Sua5b_cells.24_35.rep



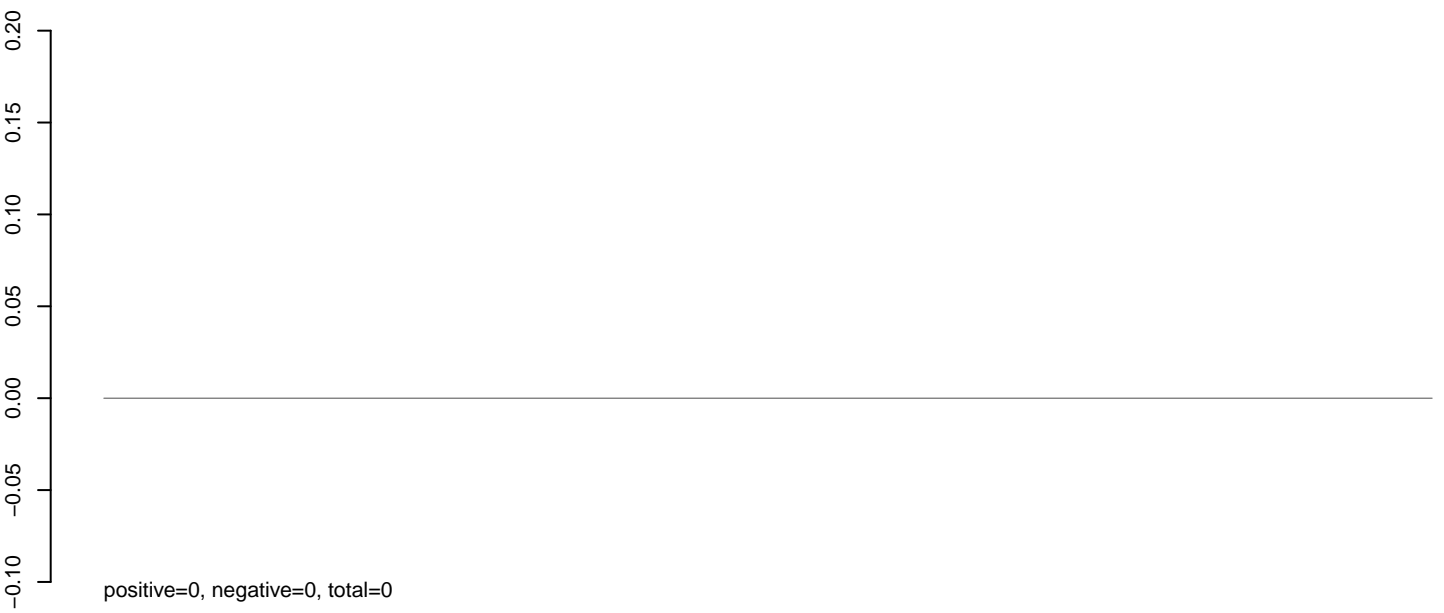
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AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



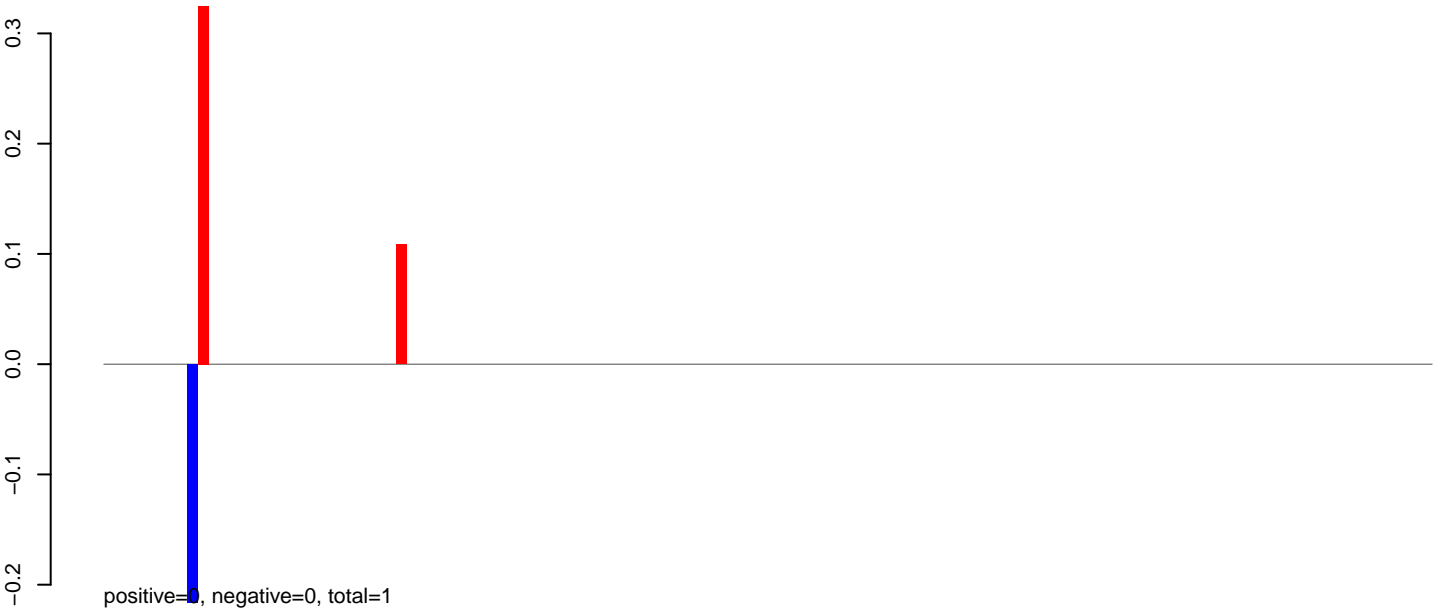
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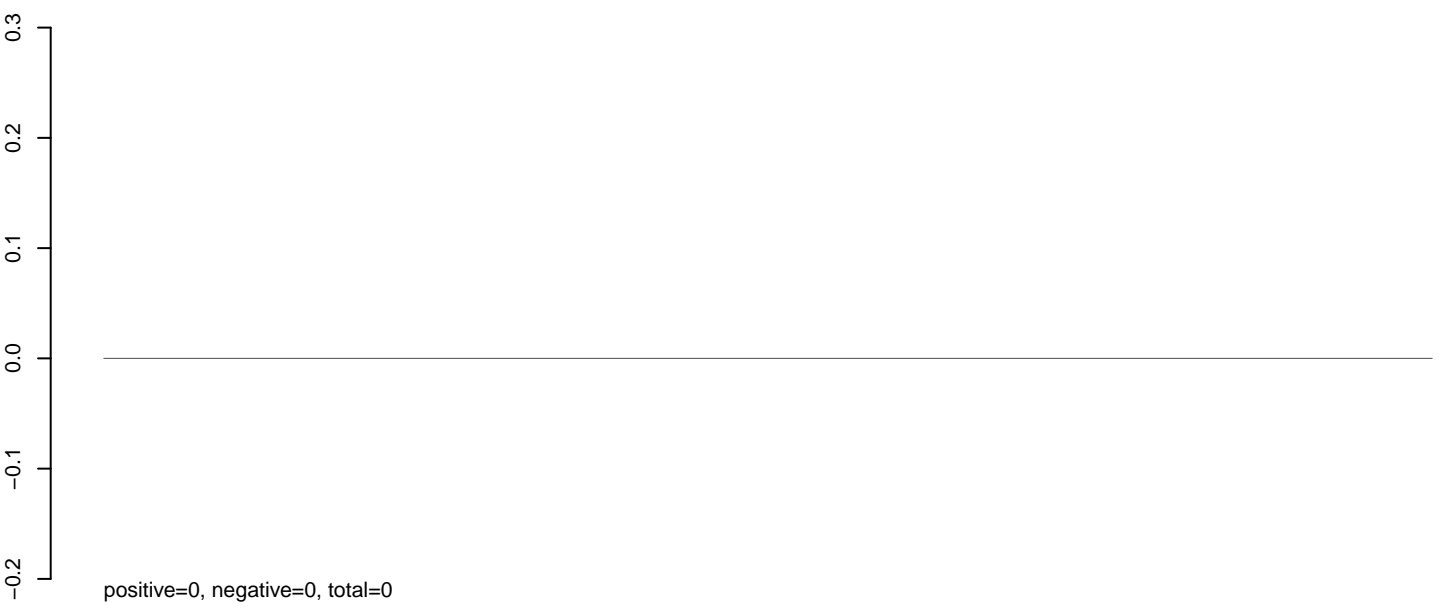
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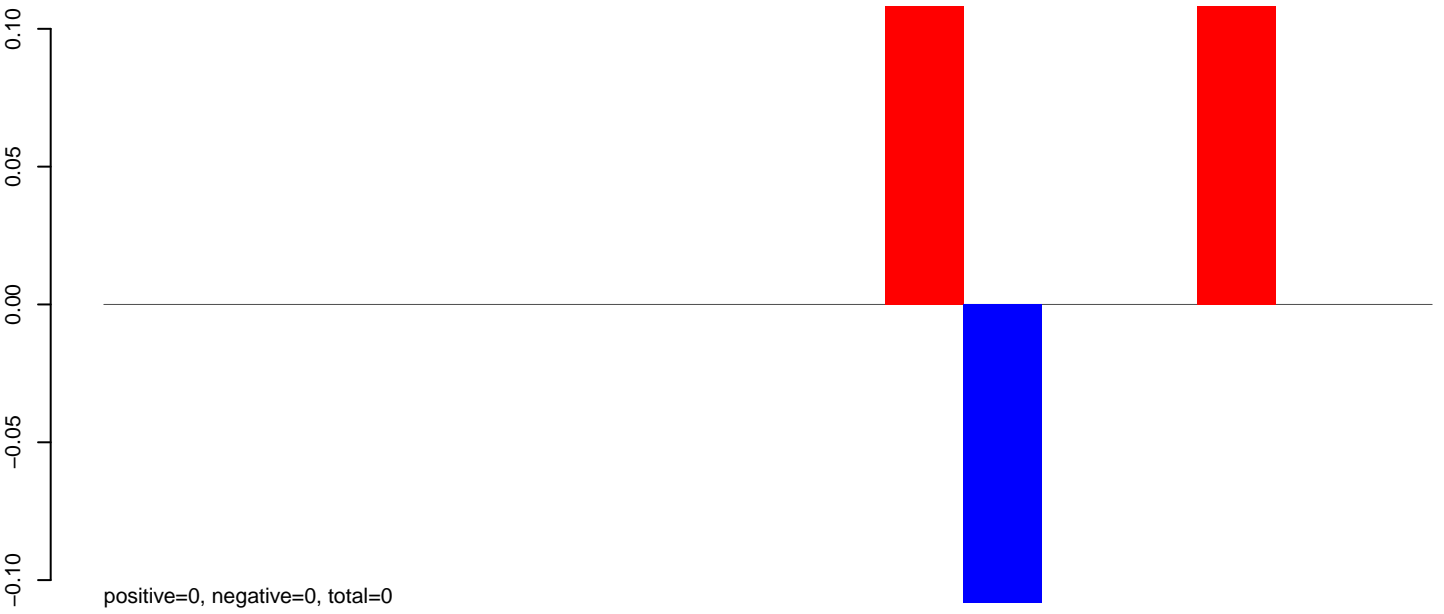
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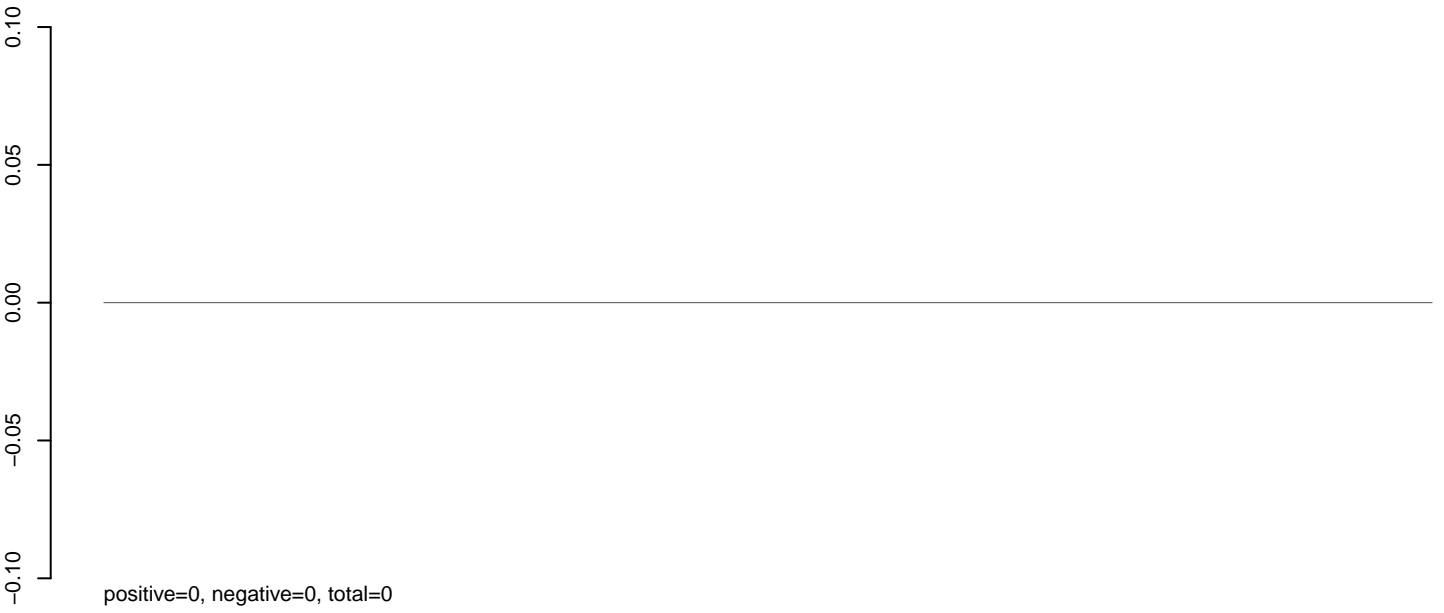
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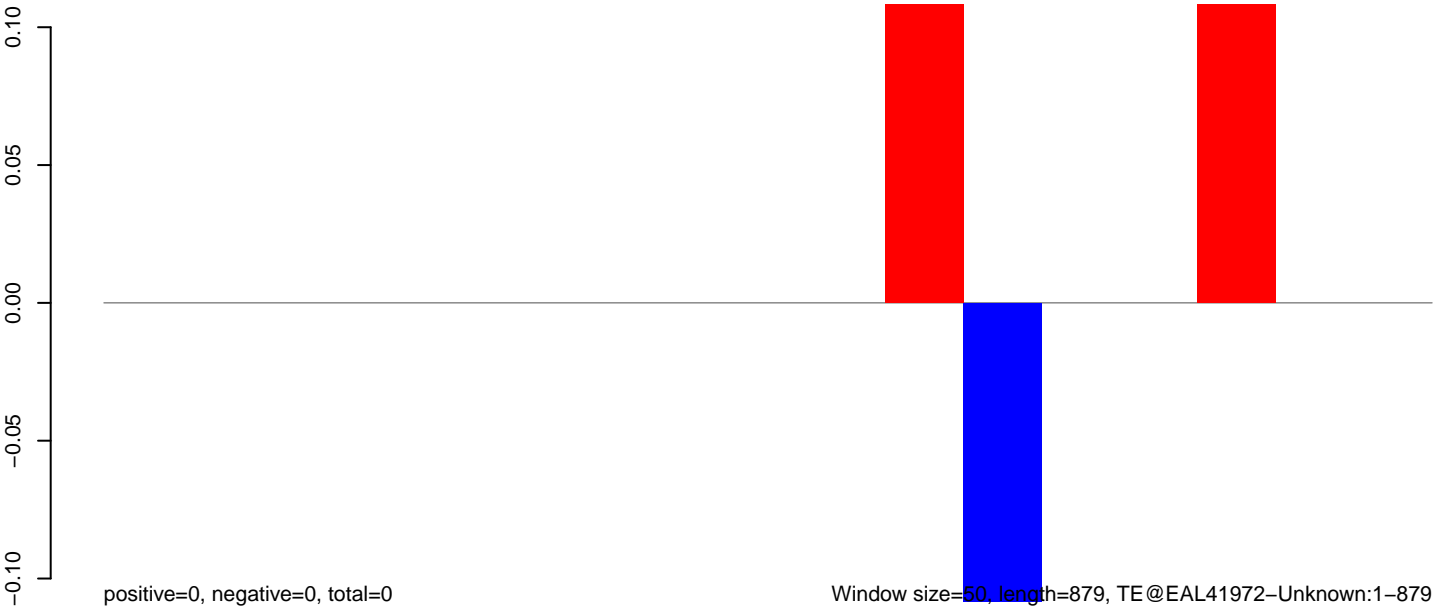
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AnGam_Sua5b_cells.24_35.rep

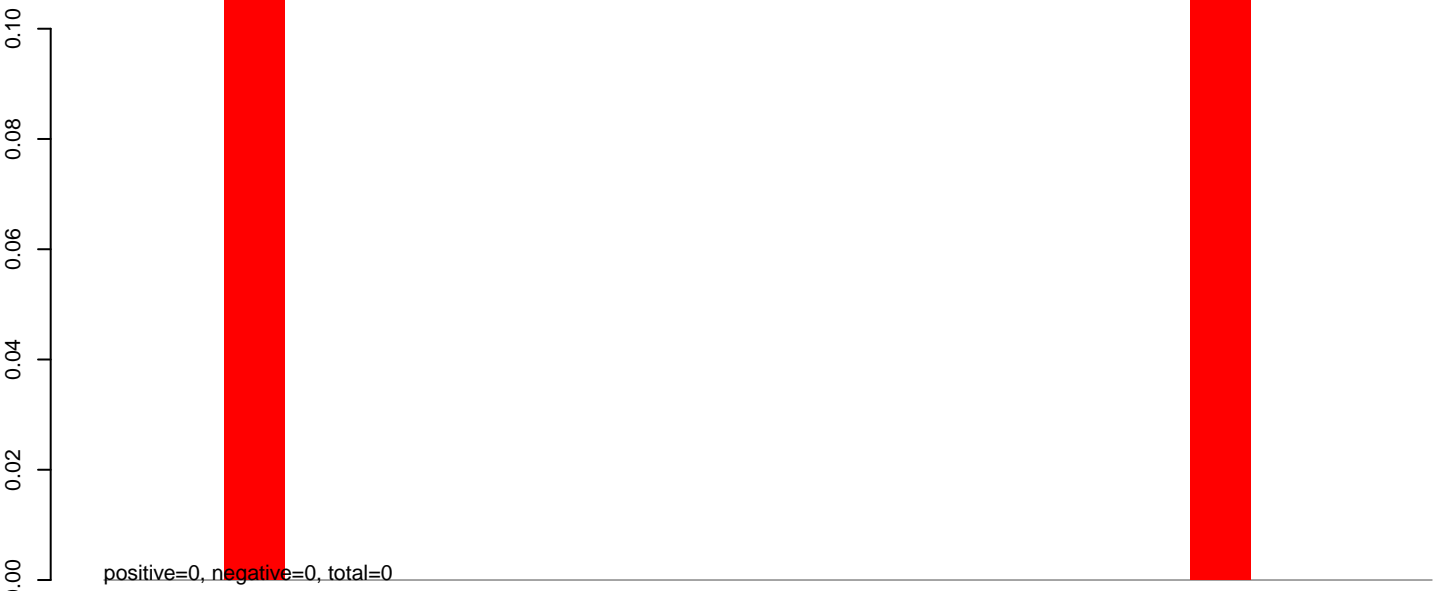


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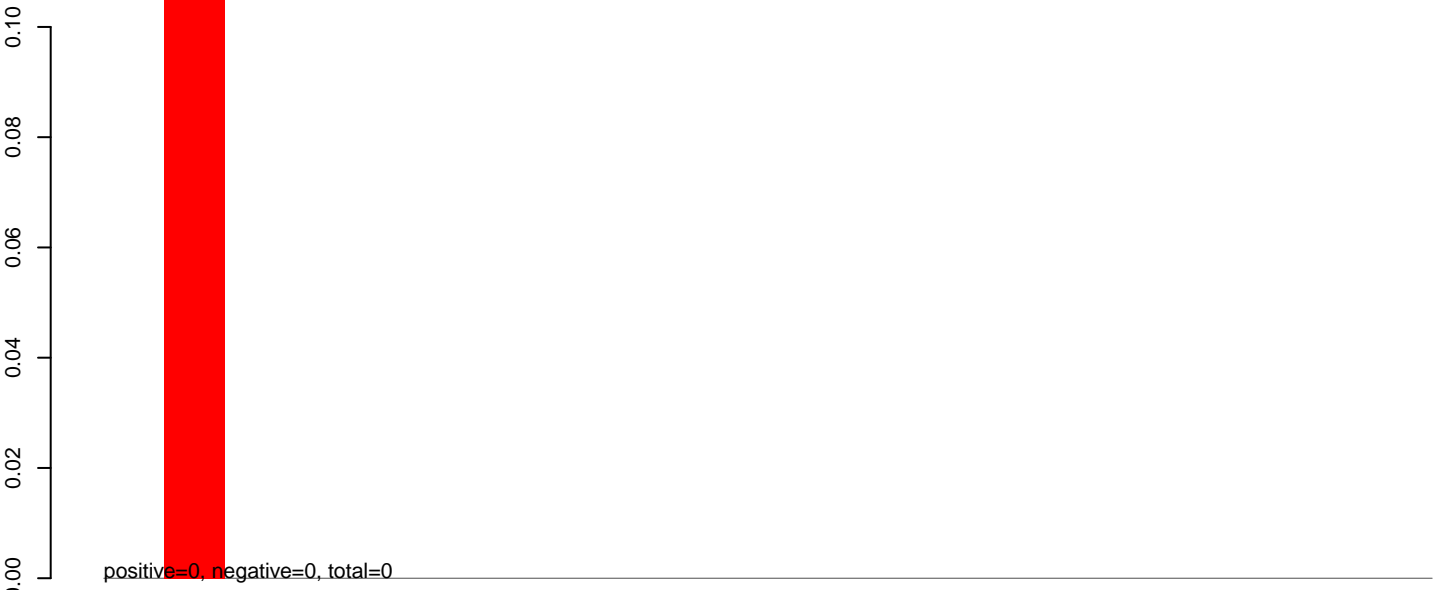


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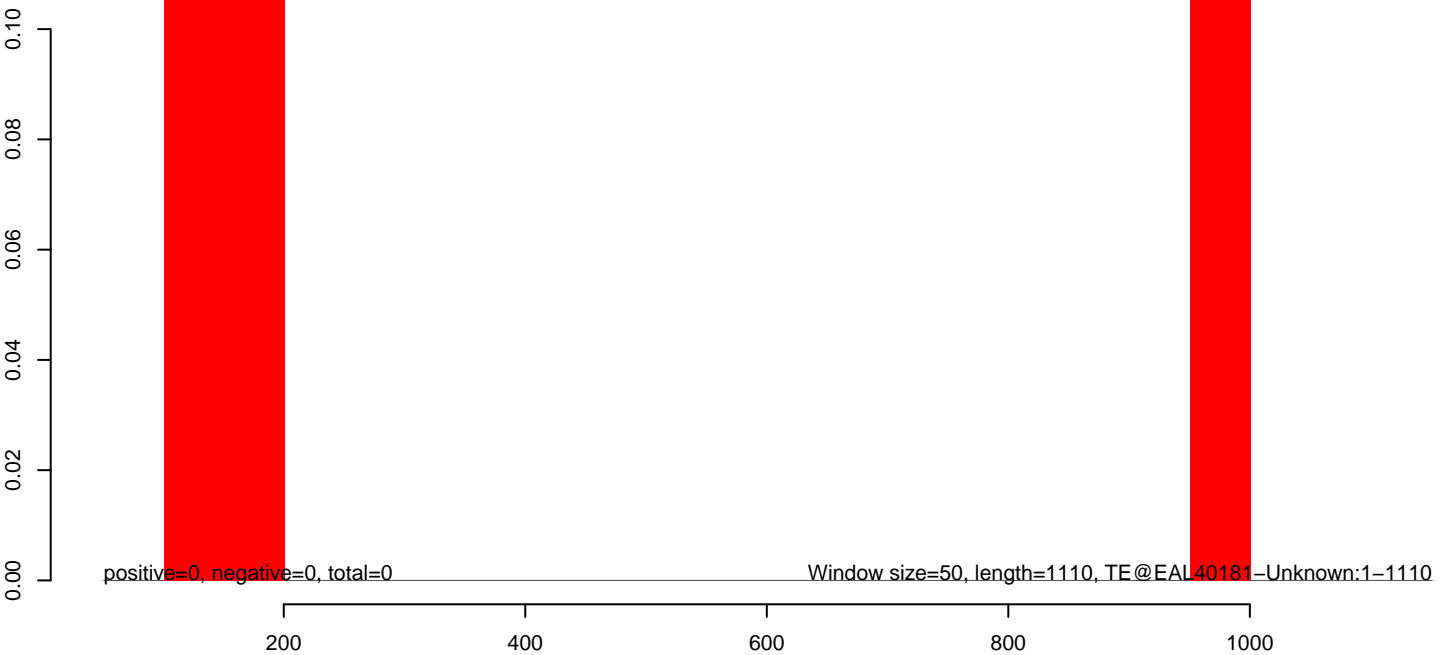
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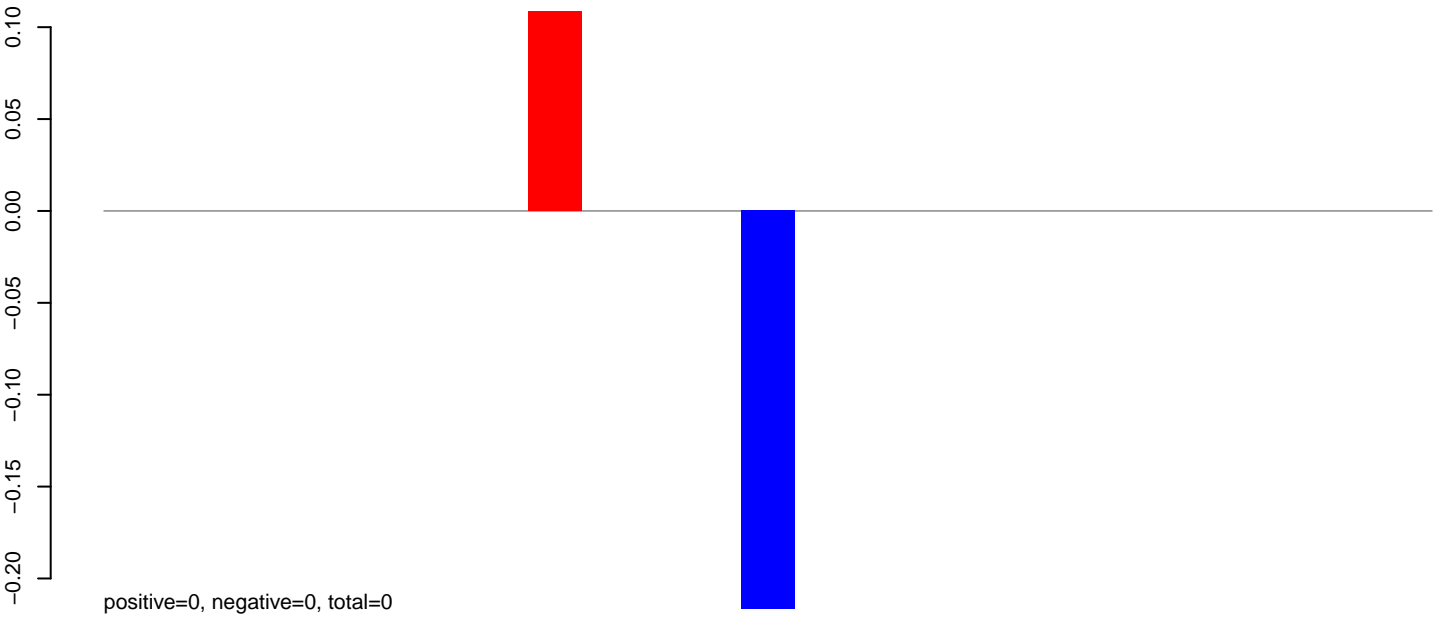
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AnGam_Sua5b_cells.rep



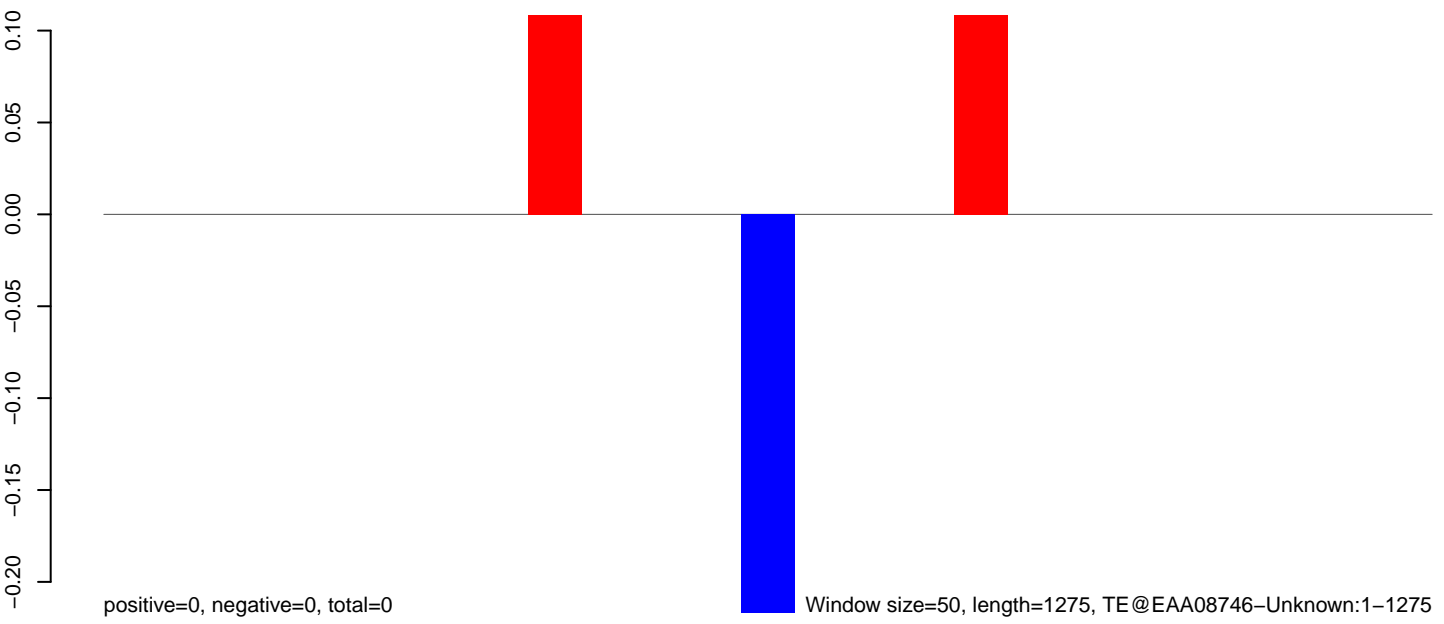
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AnGam_Sua5b_cells.24_35.rep



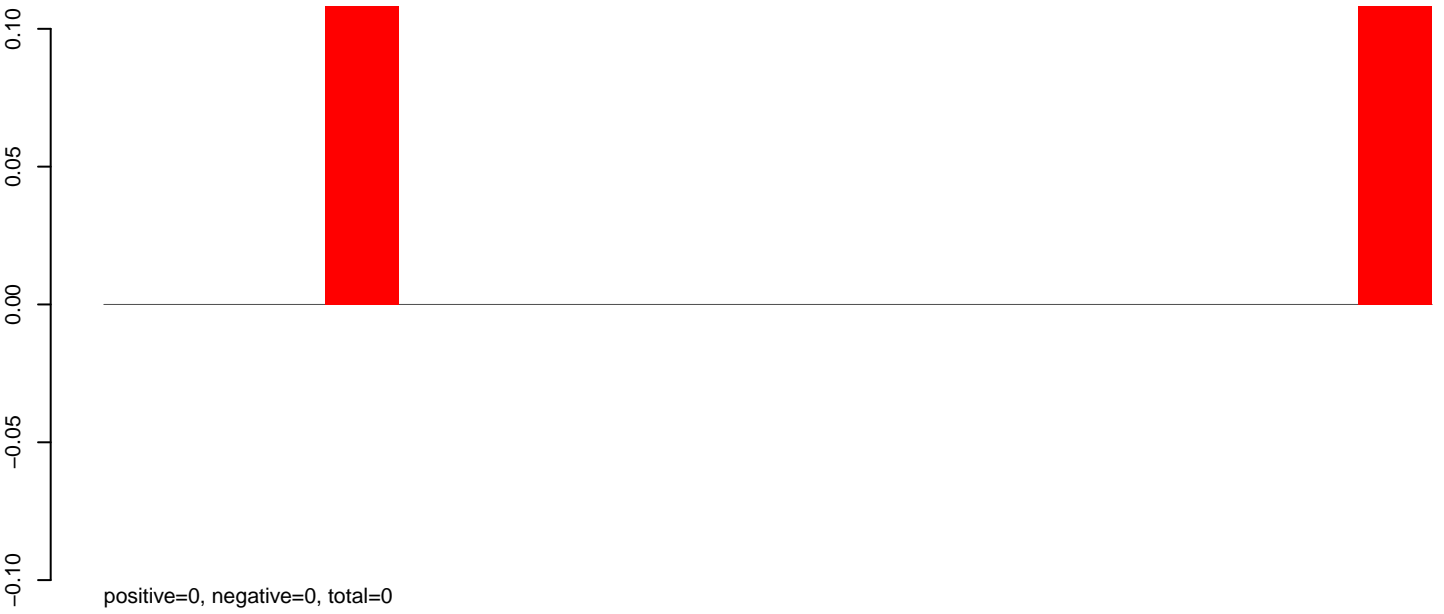
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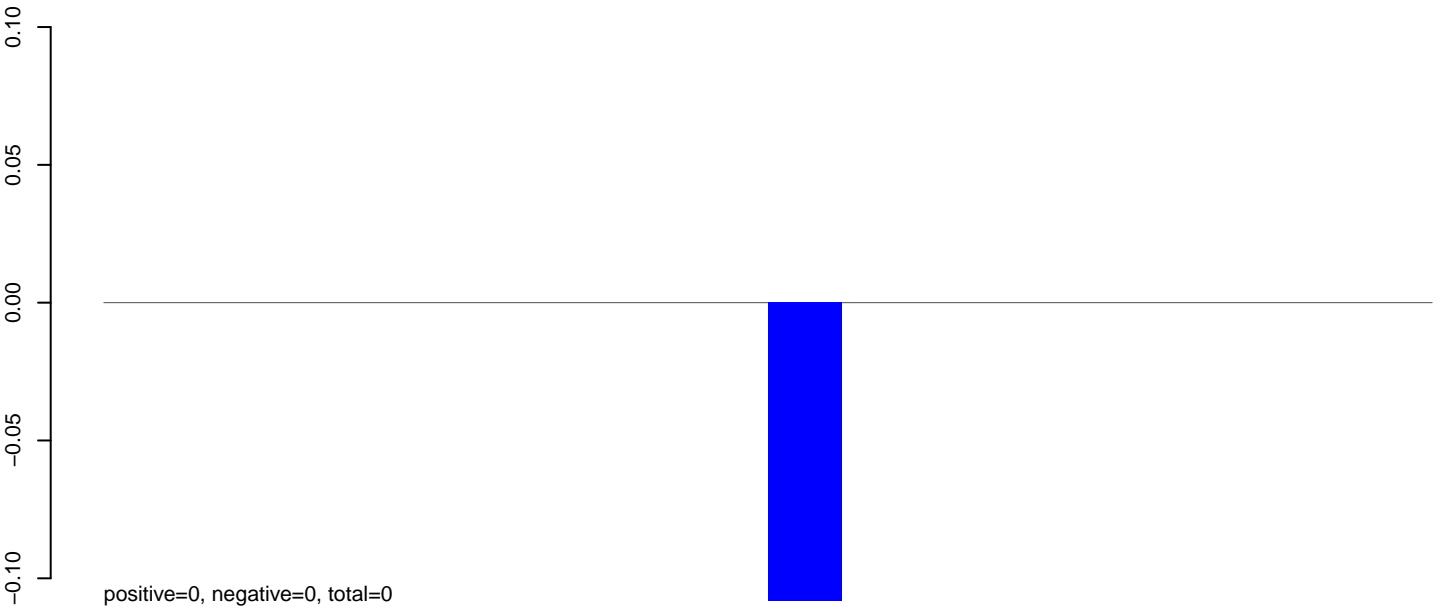
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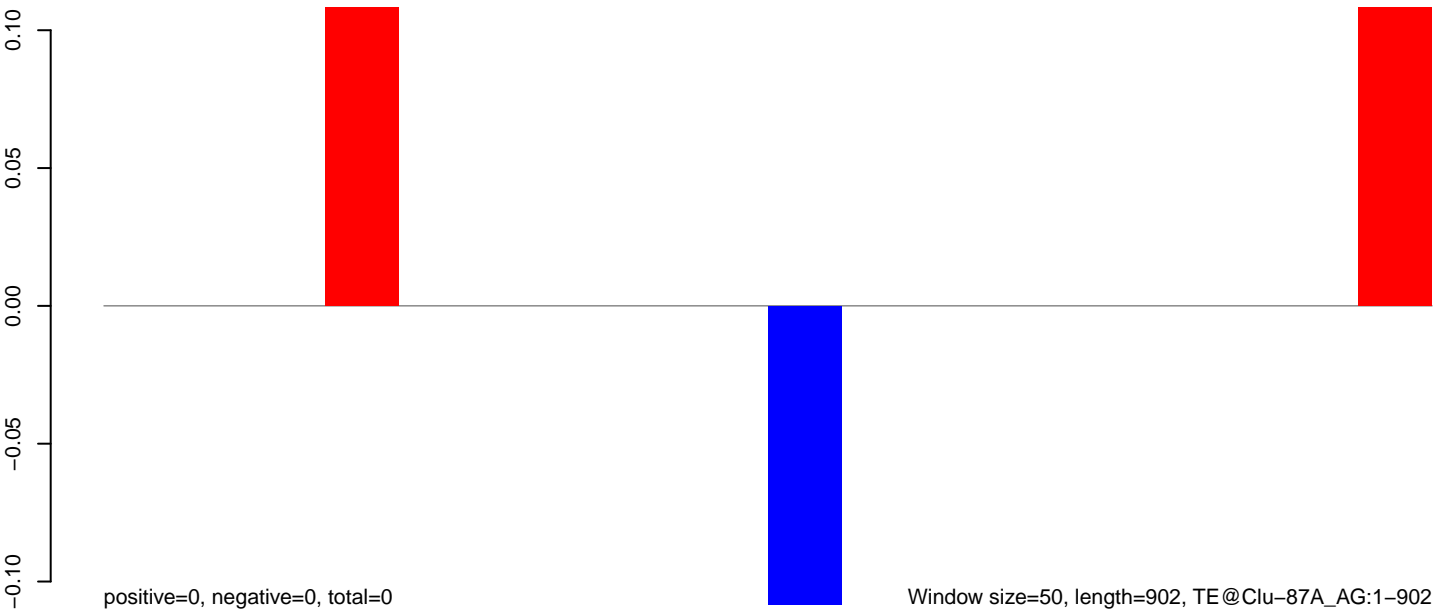
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AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



200

400

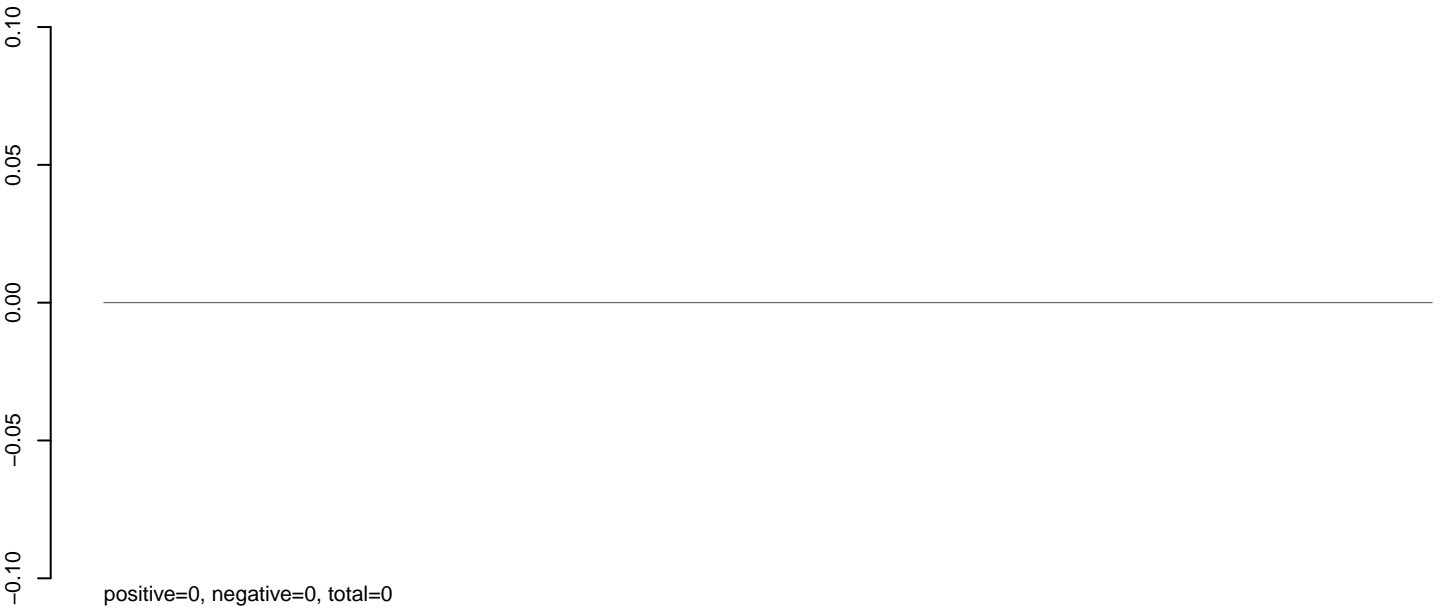
600

800

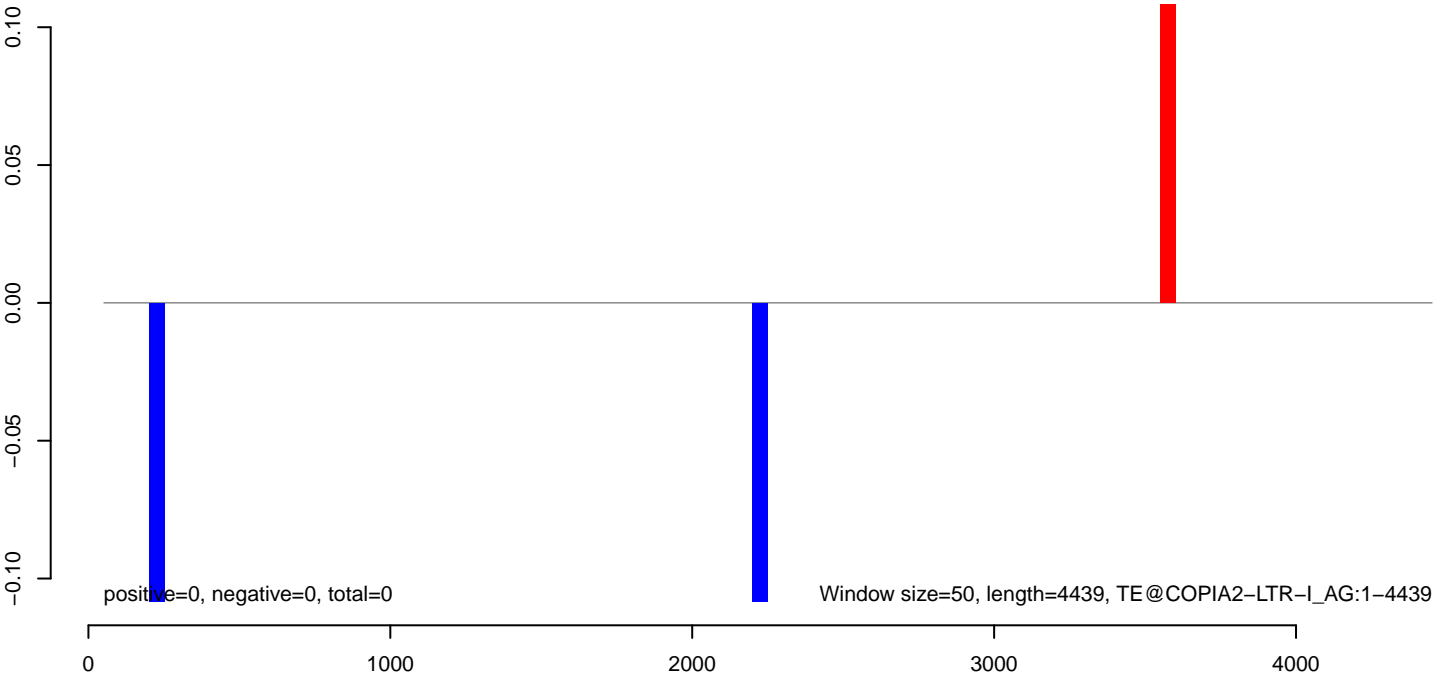
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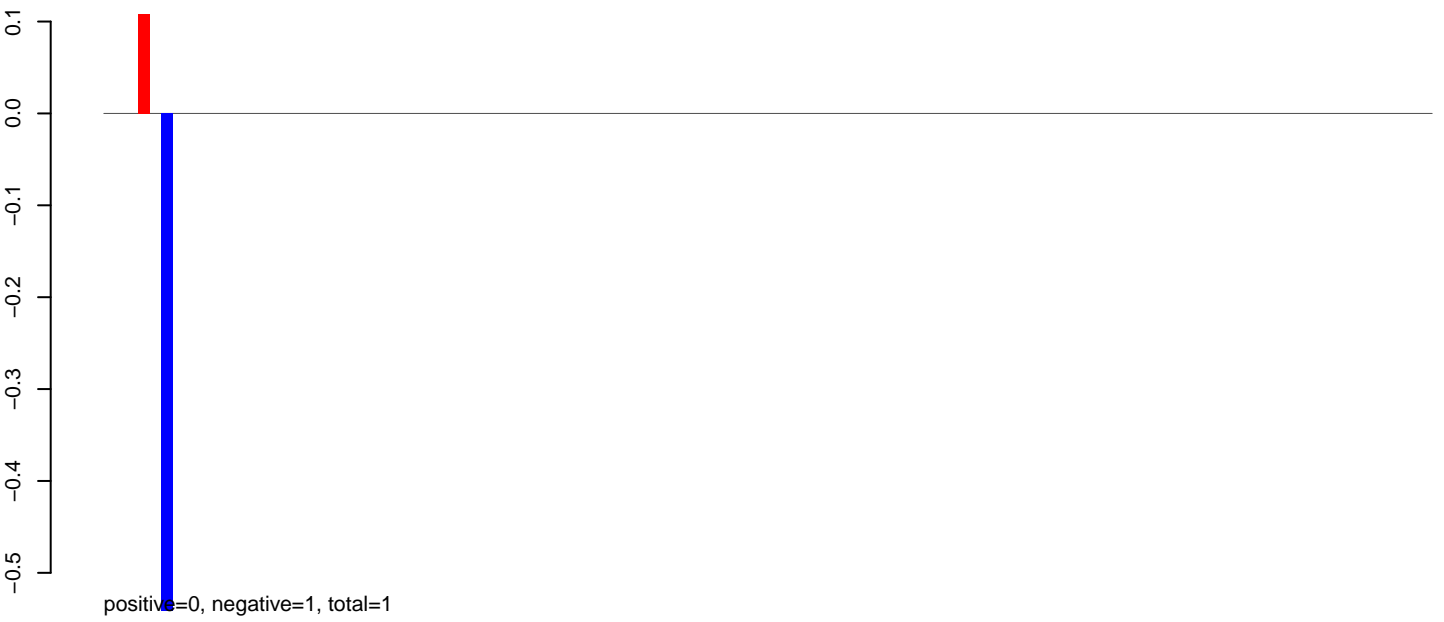
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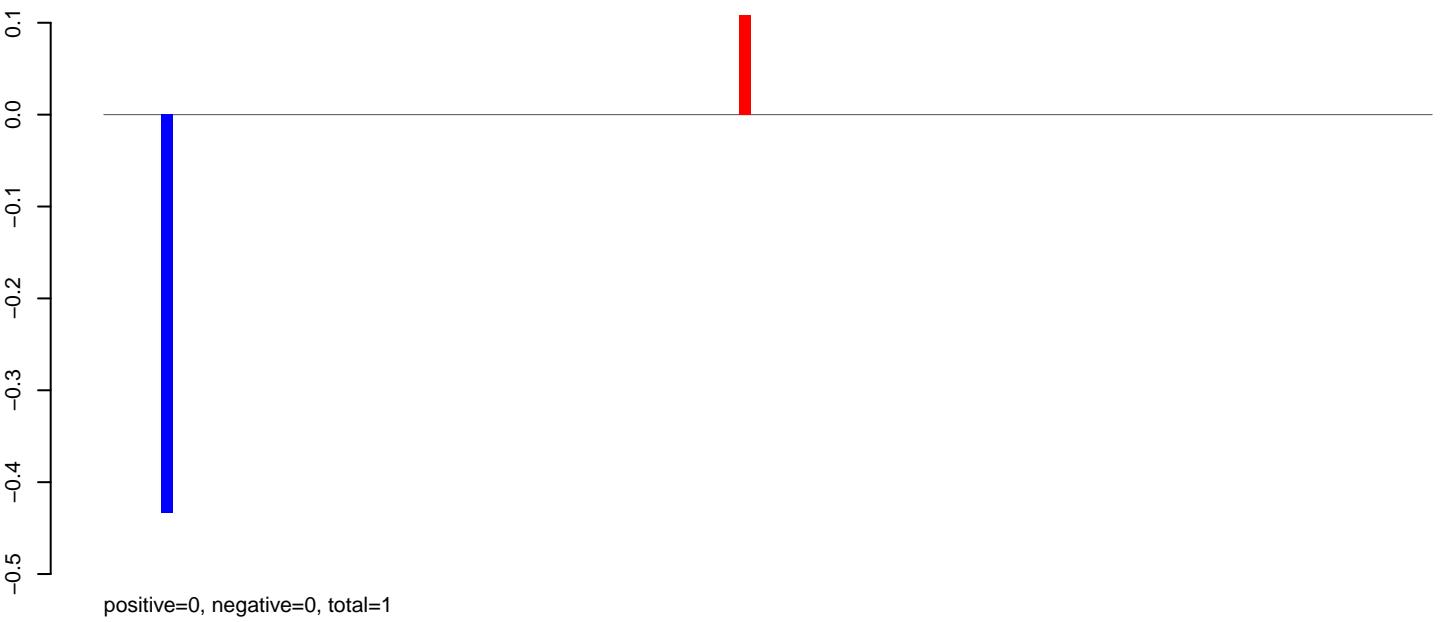
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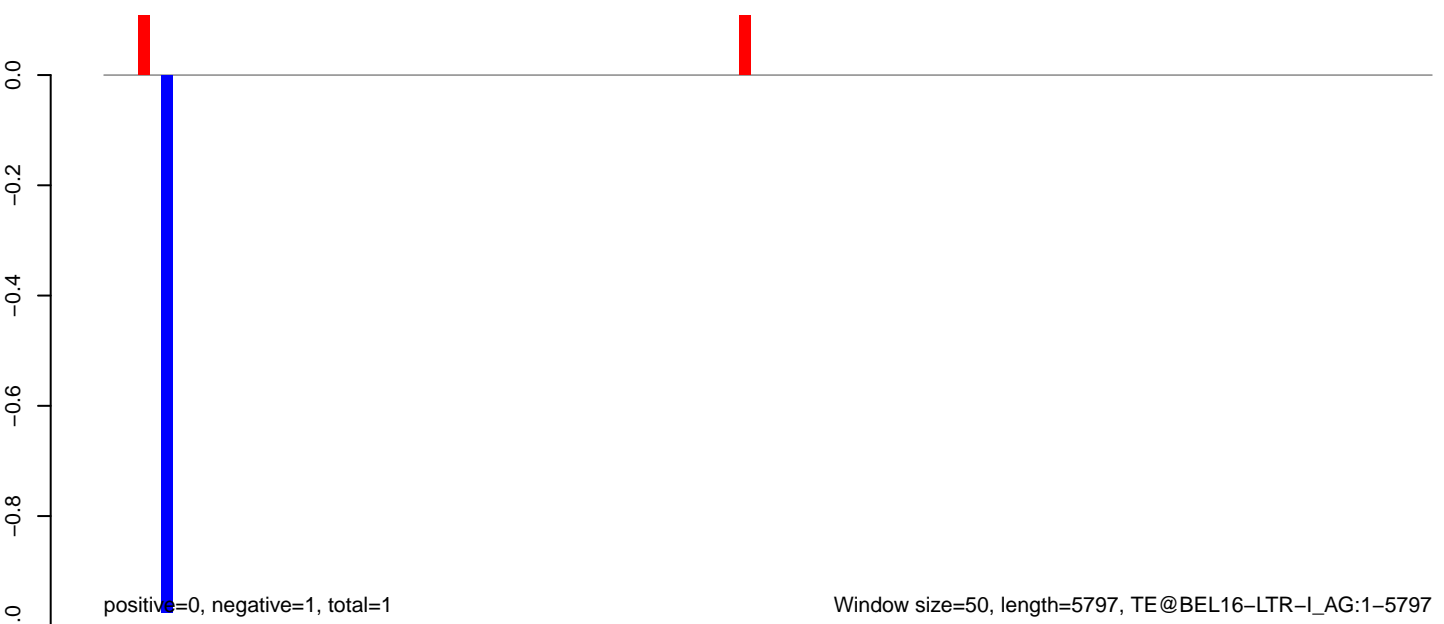
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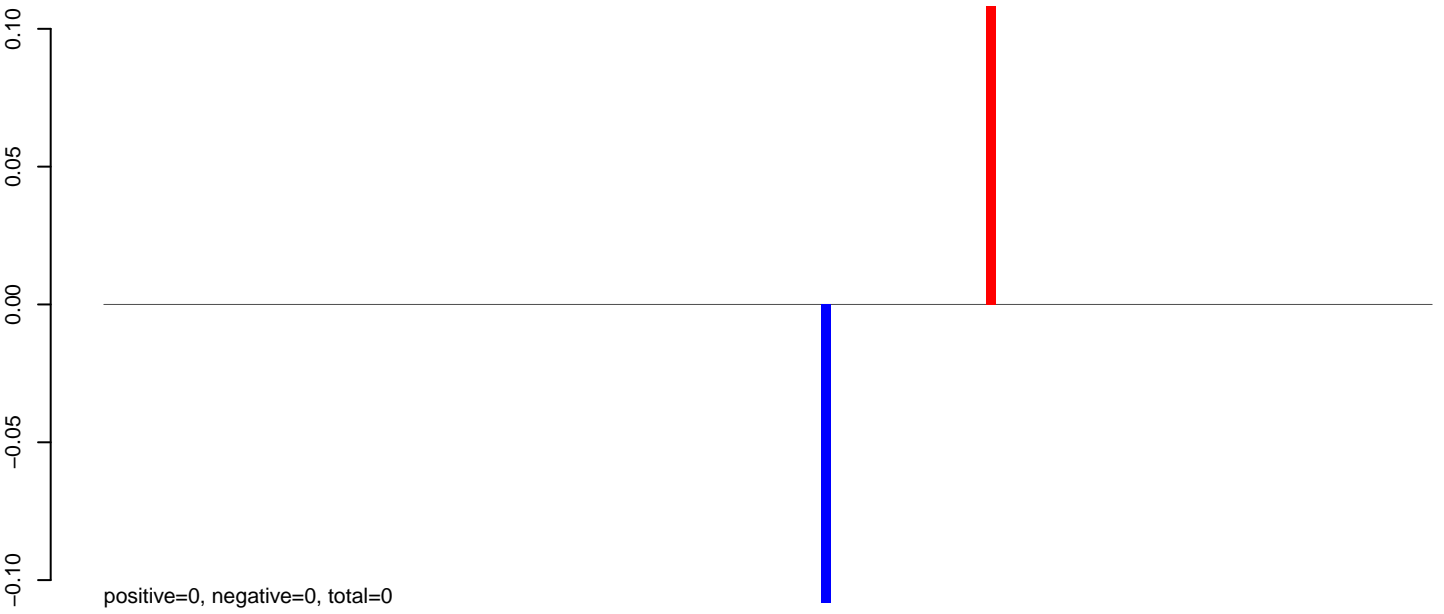
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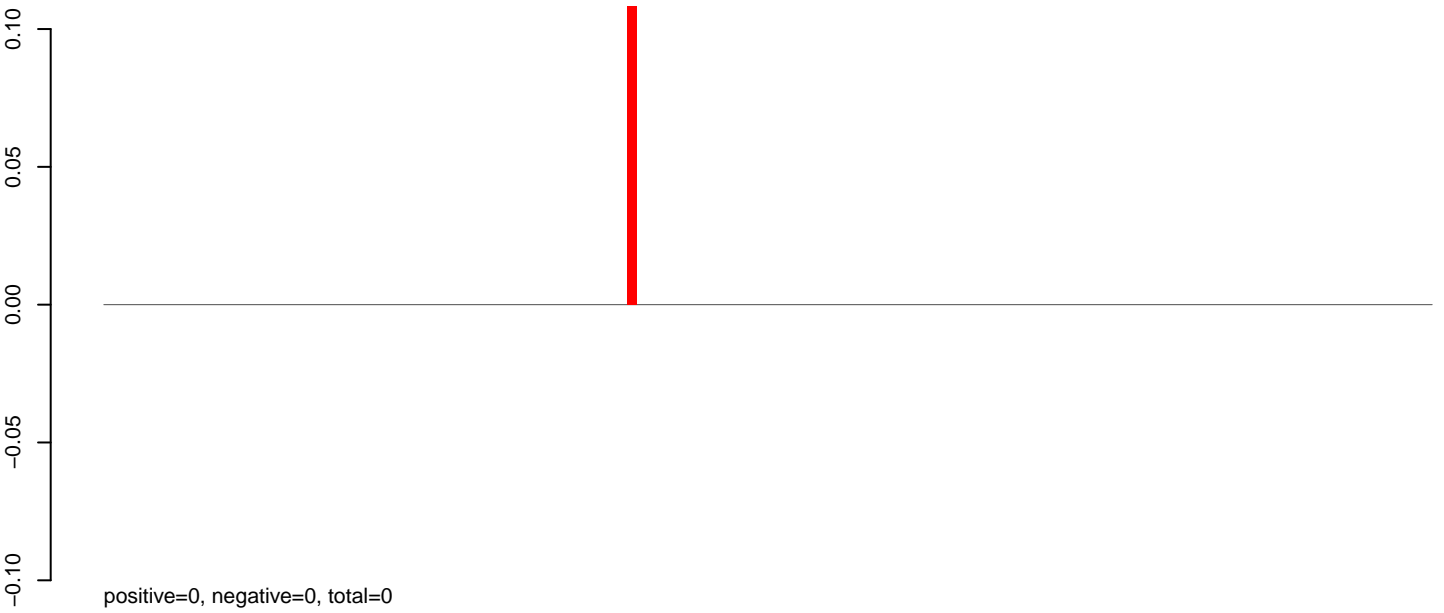
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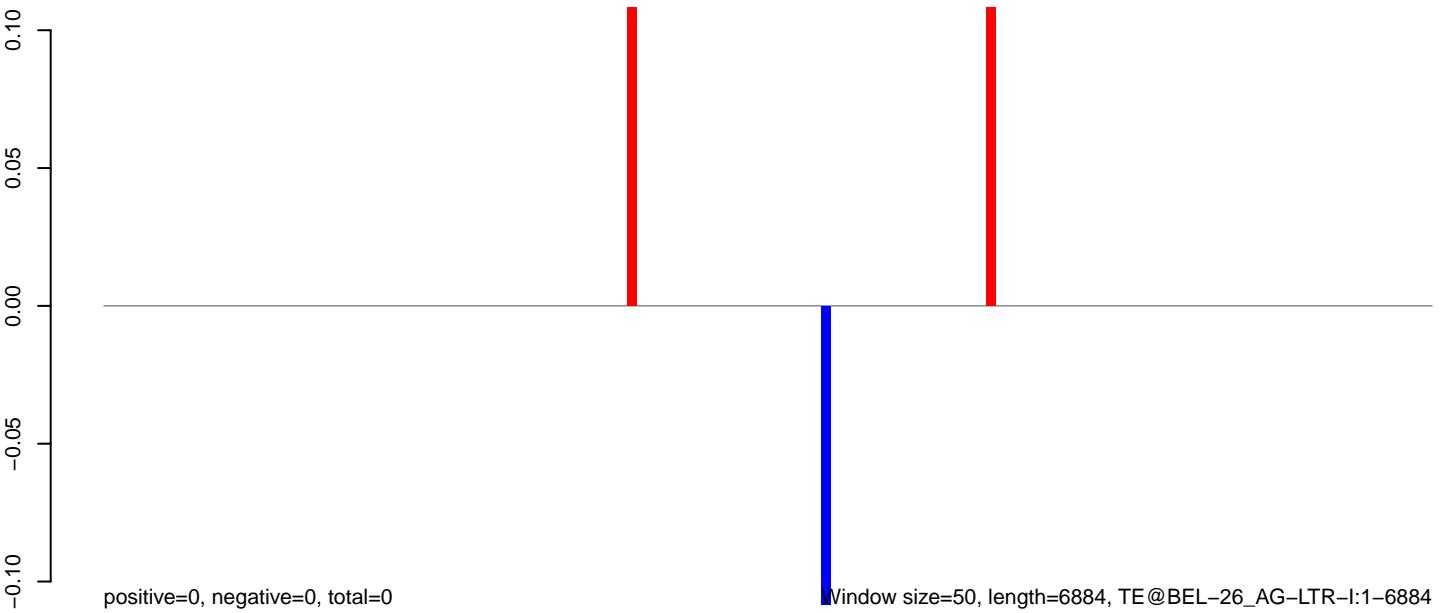
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AnGam_Sua5b_cells.24_35.rep

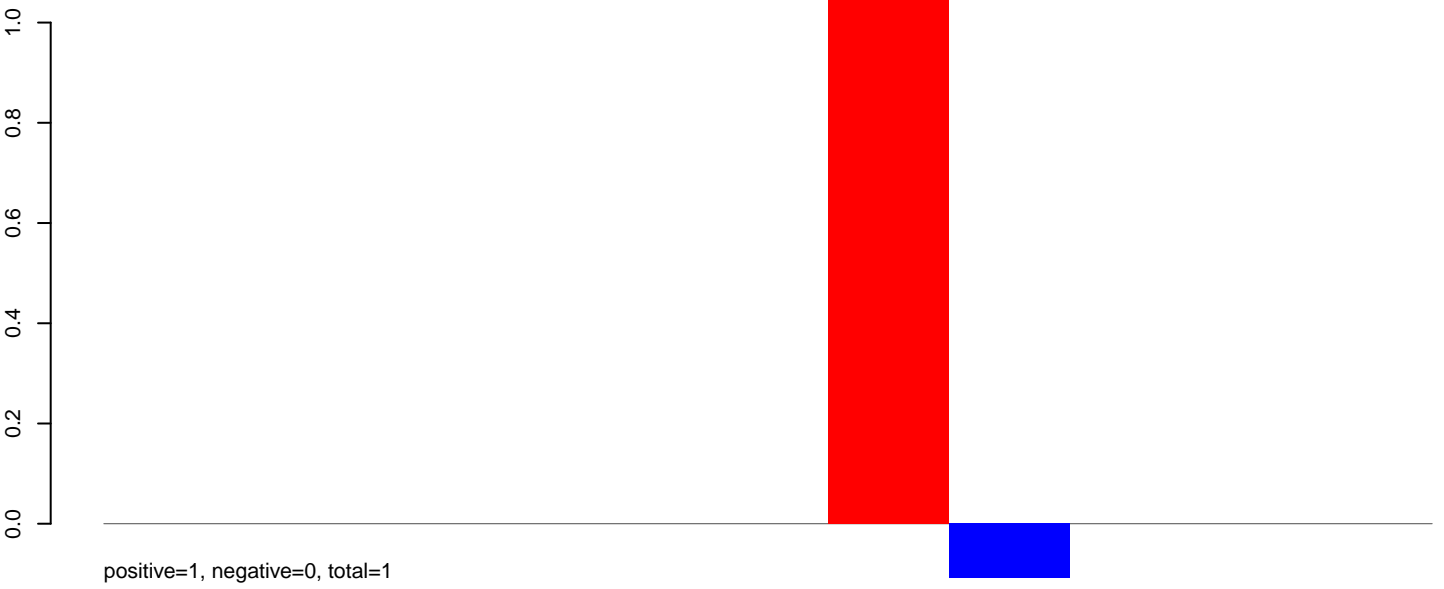


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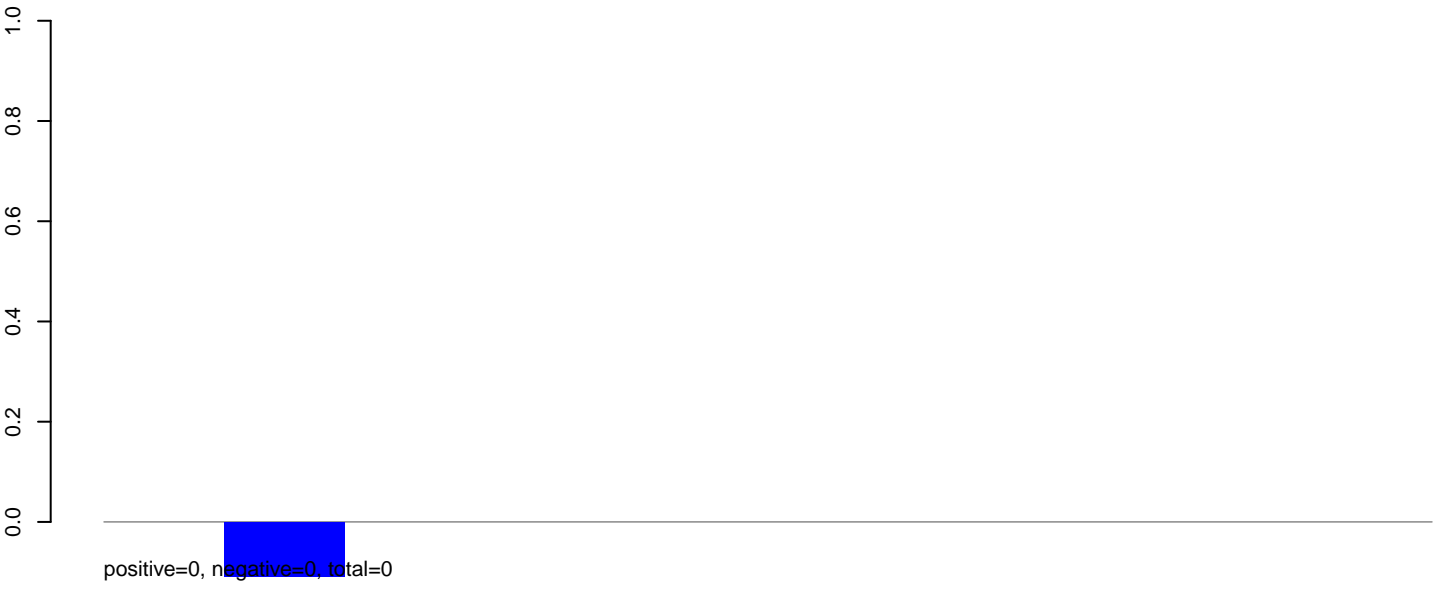


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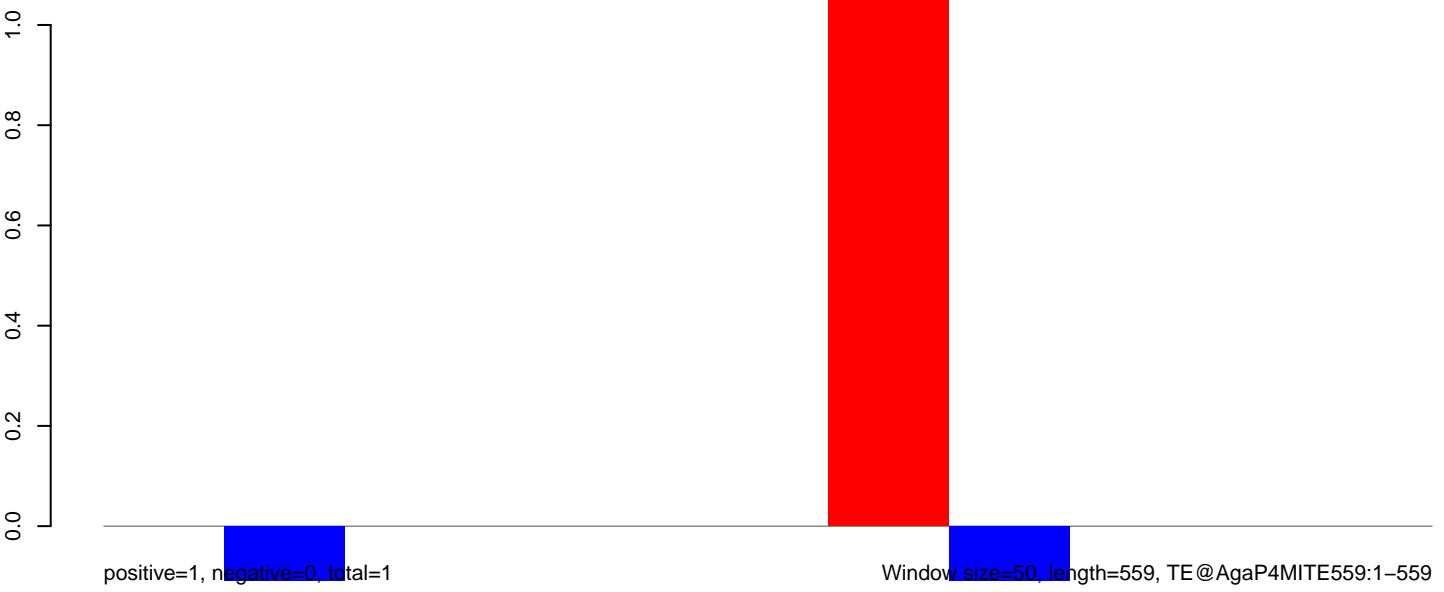
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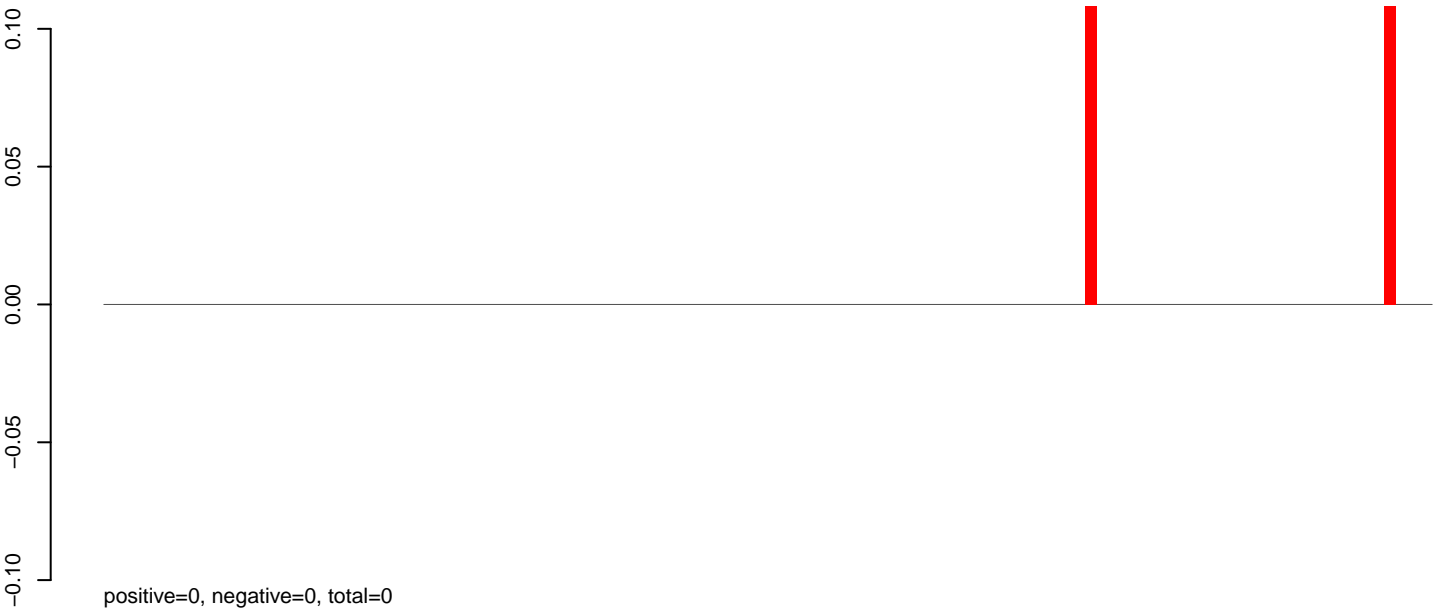


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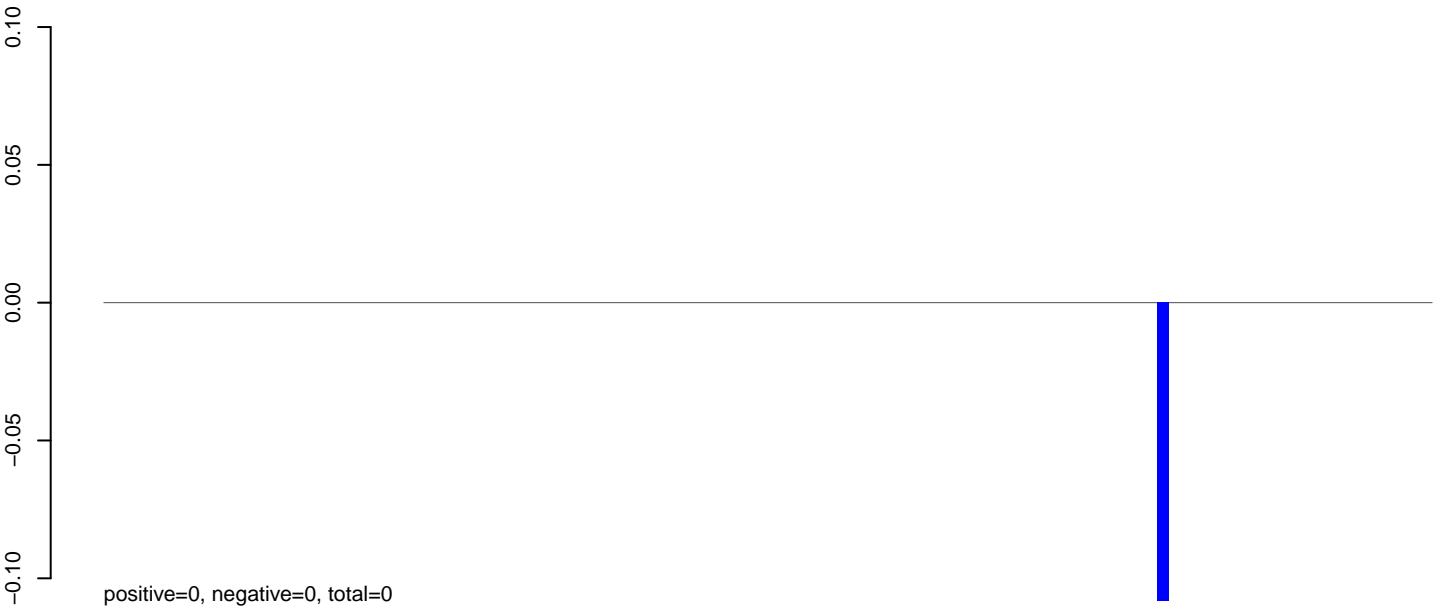


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AnGam_Sua5b_cells.18_23.rep



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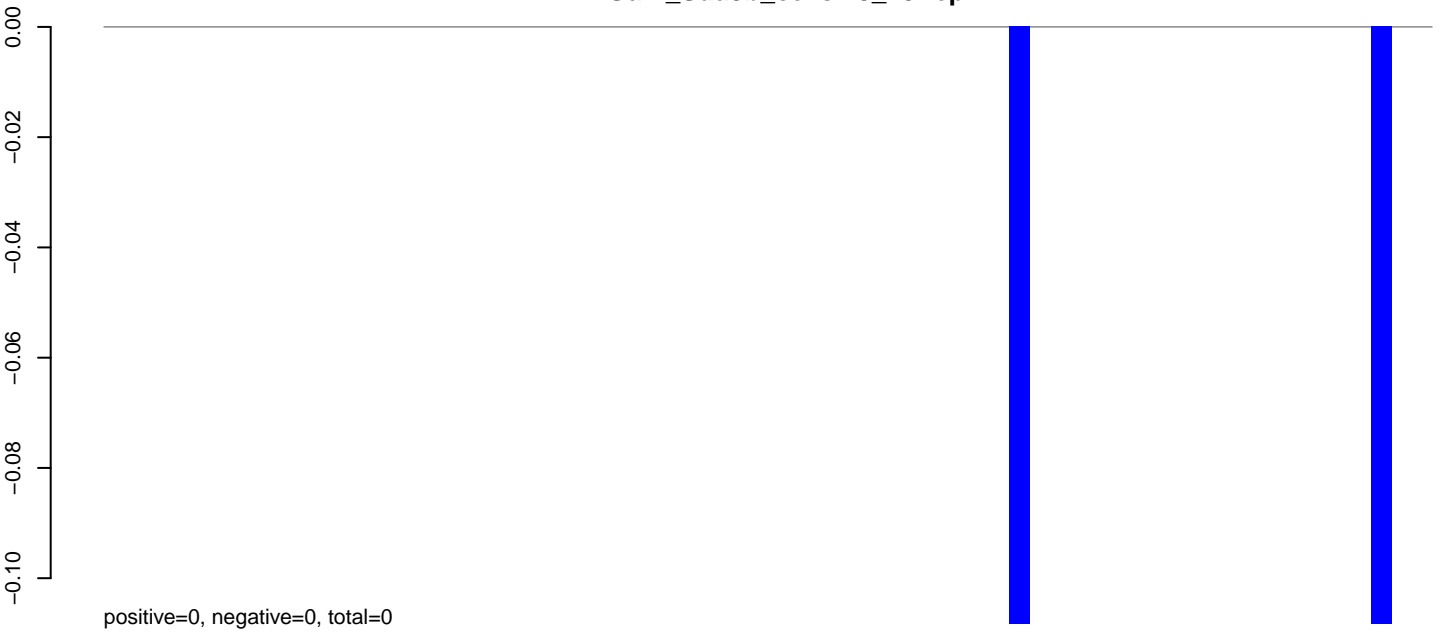


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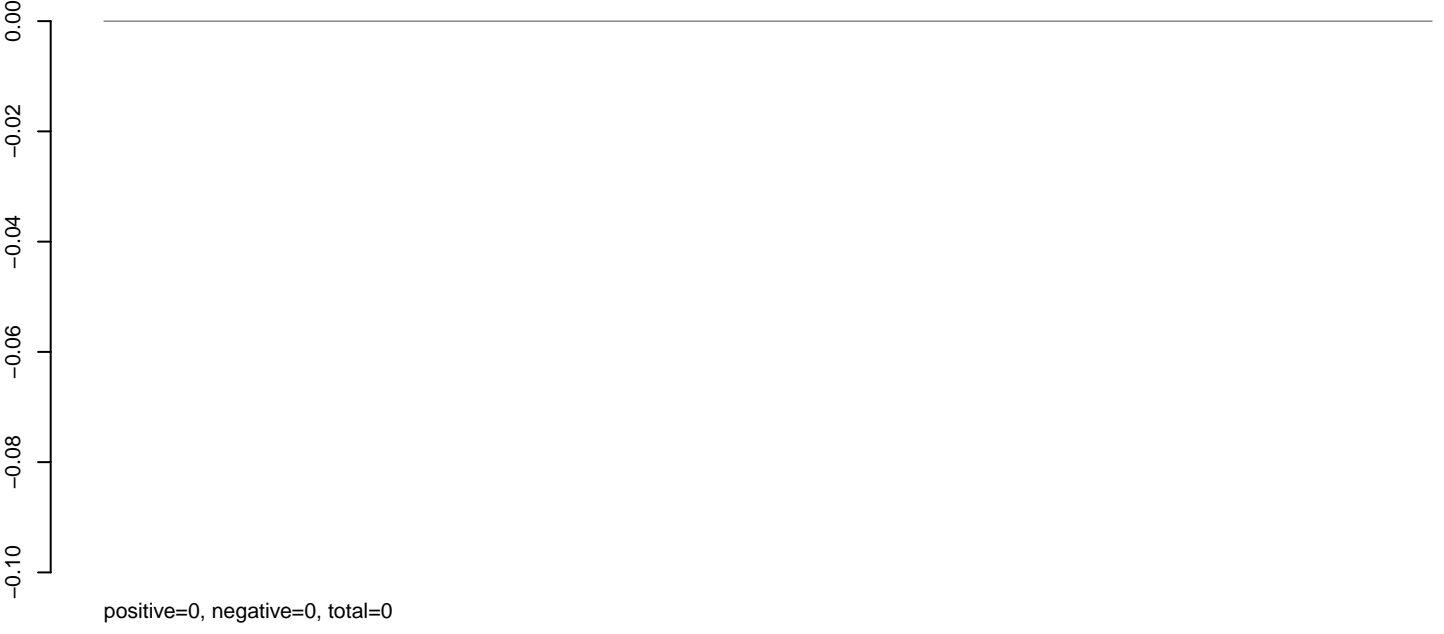


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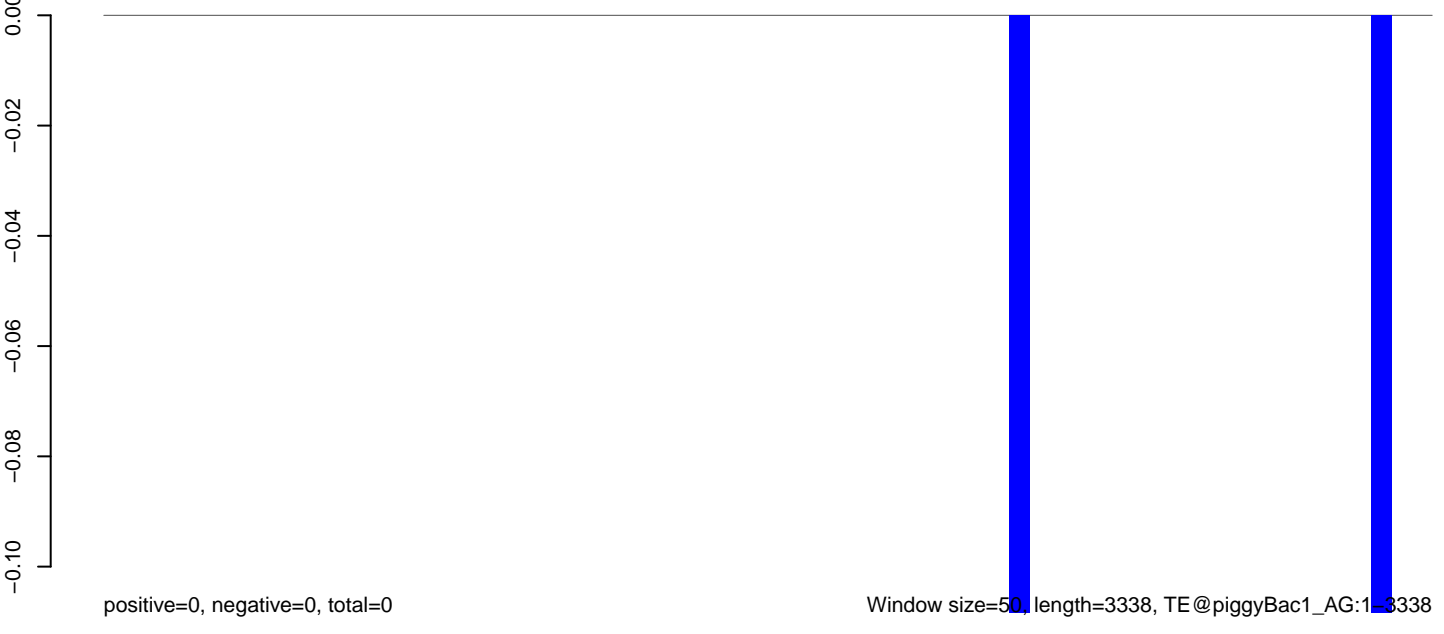
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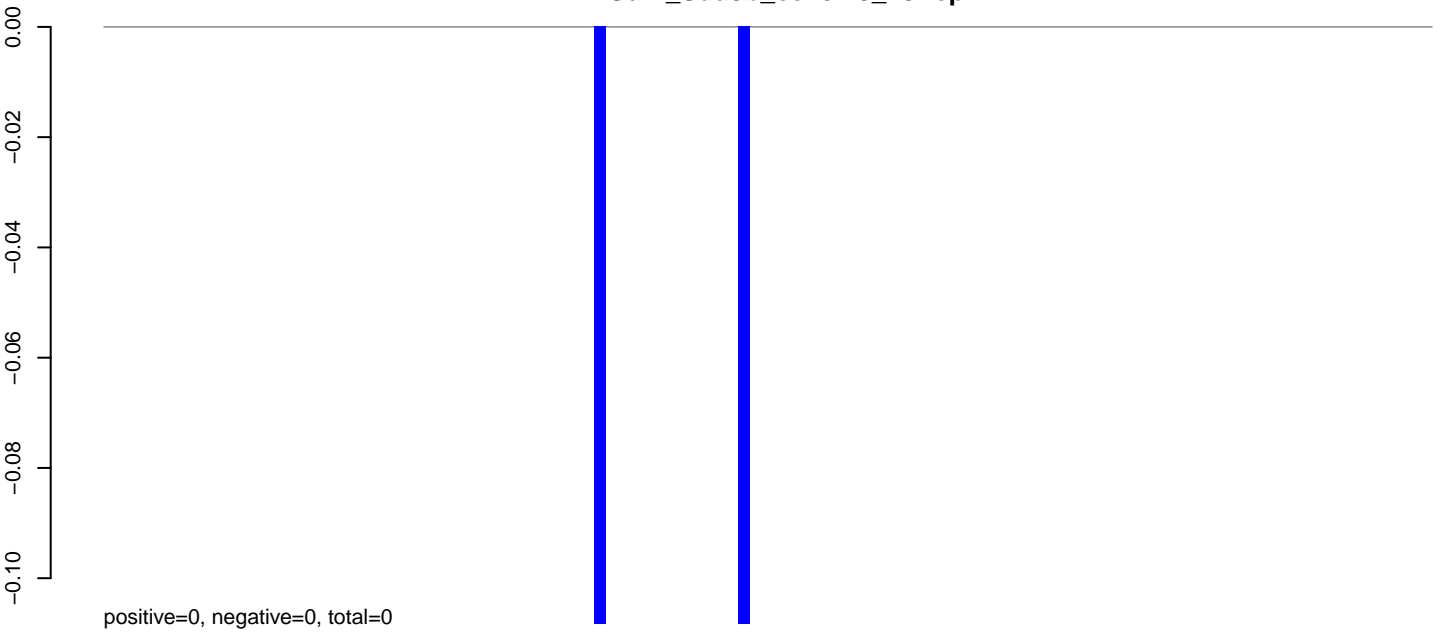
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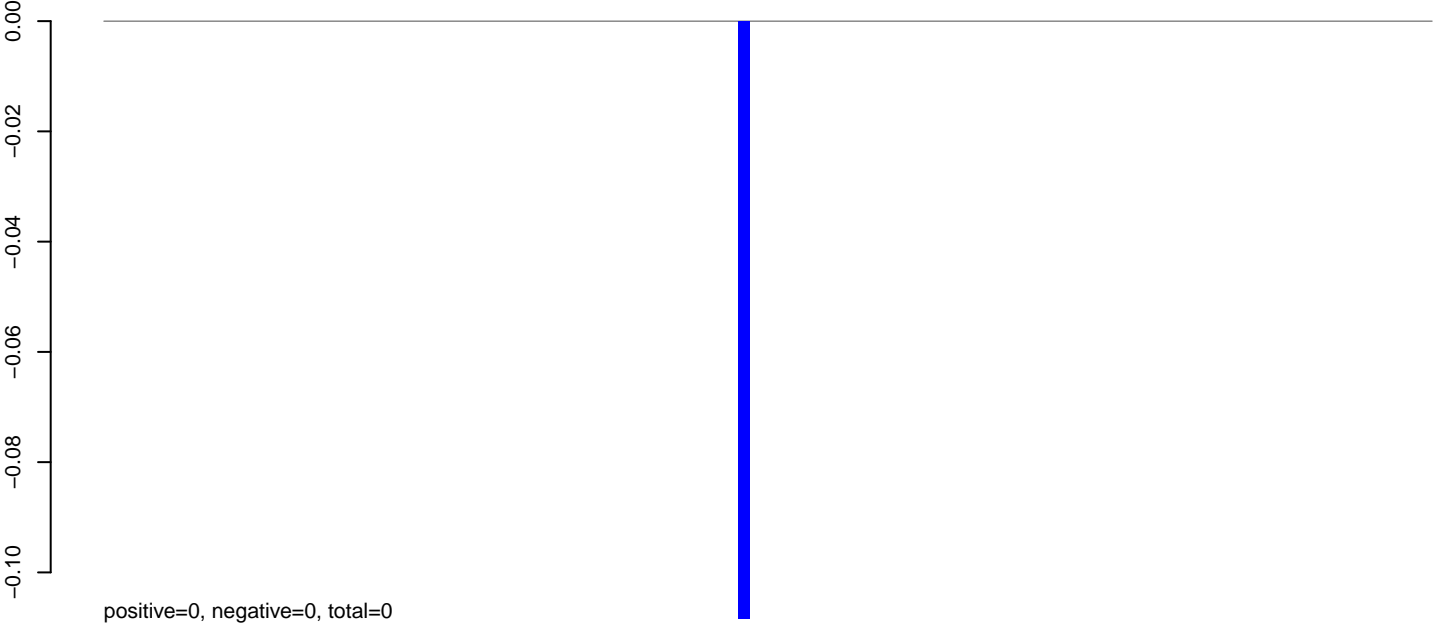
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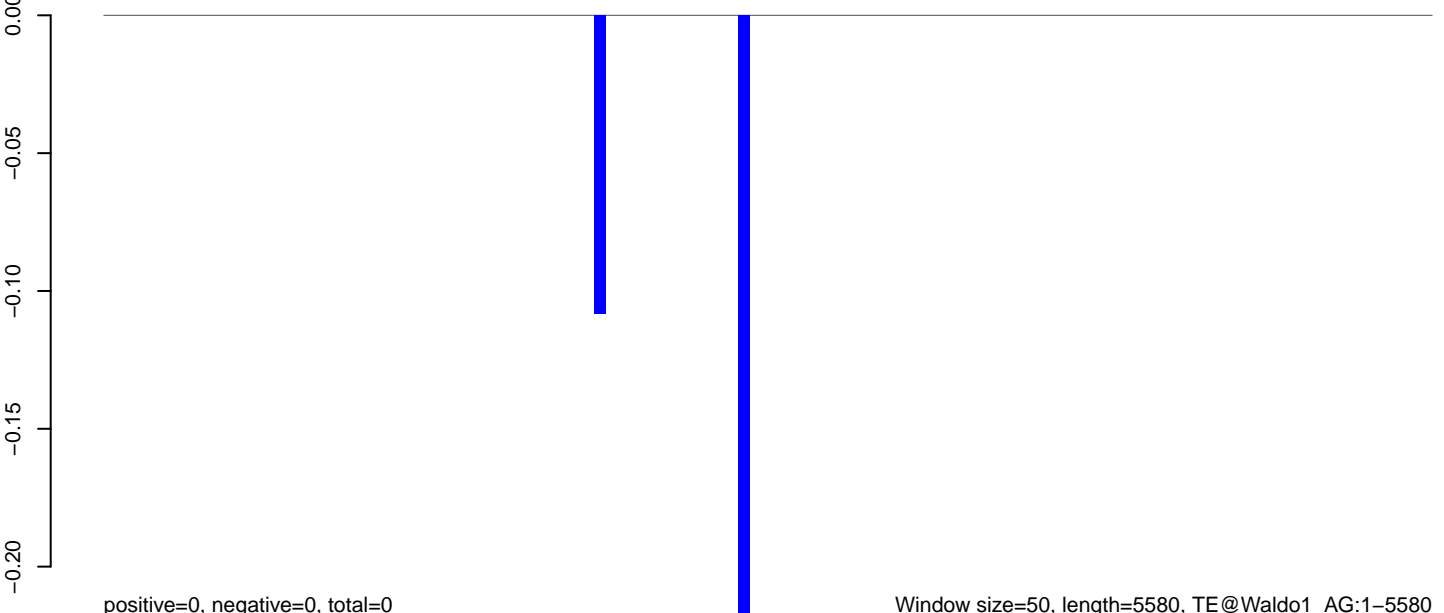
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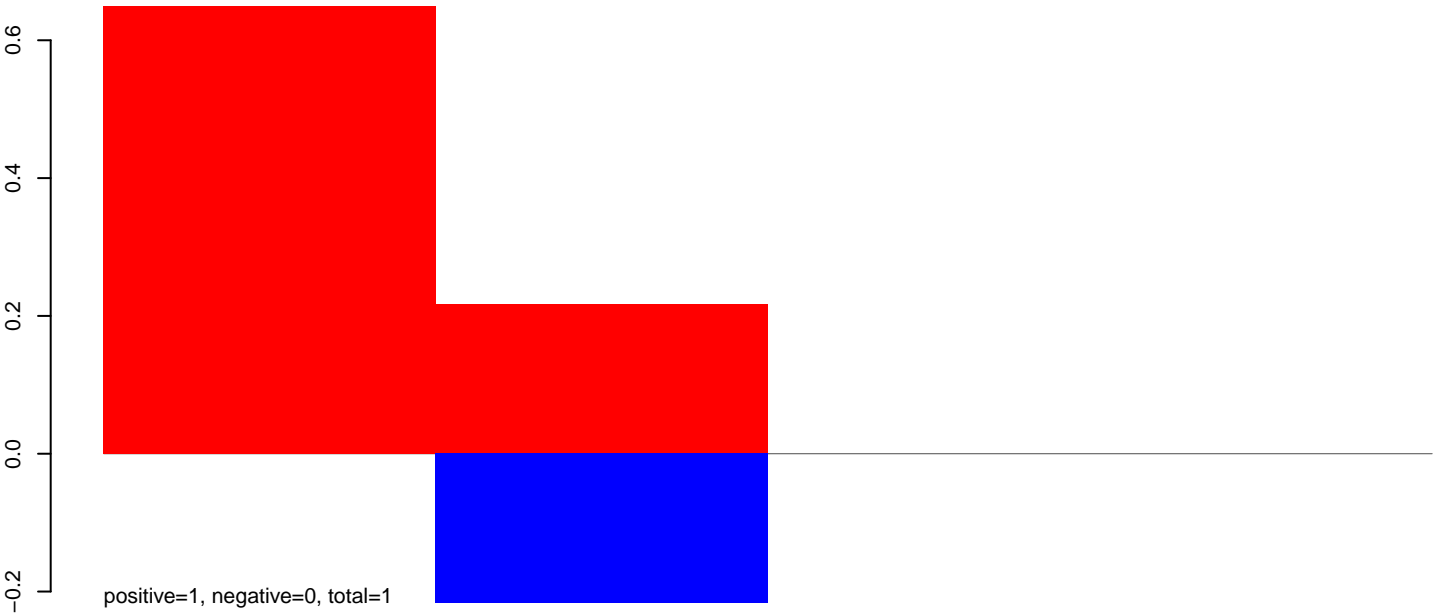
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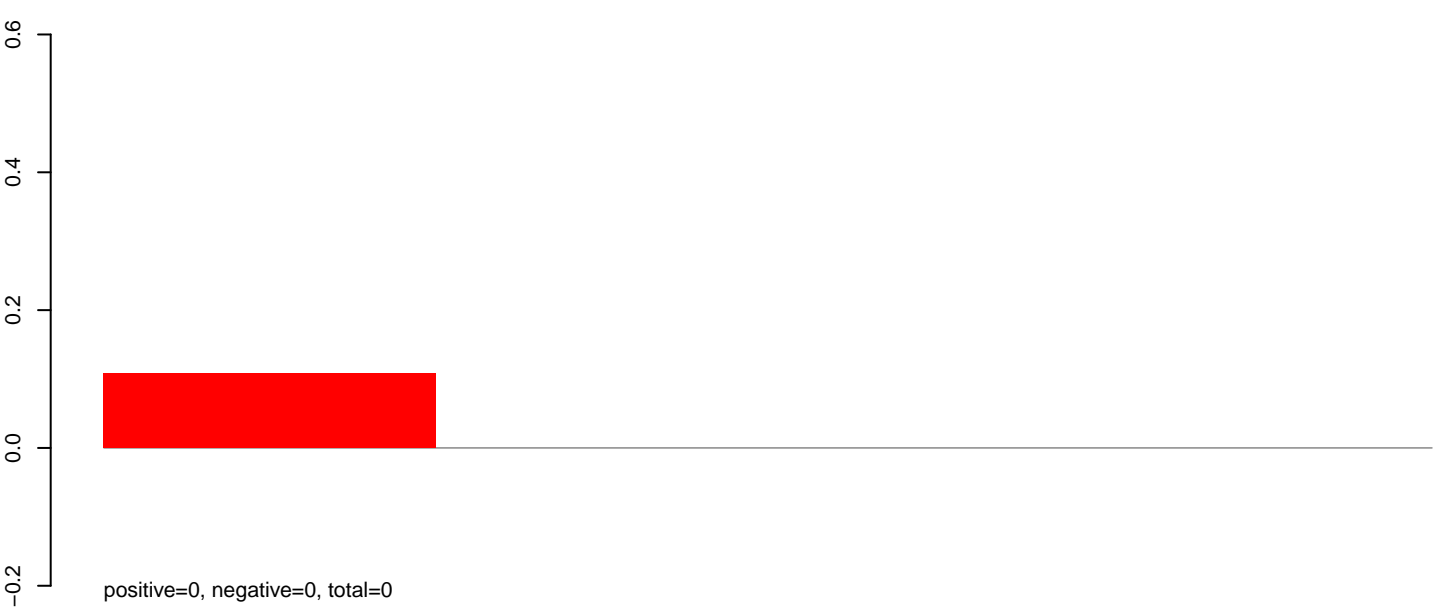
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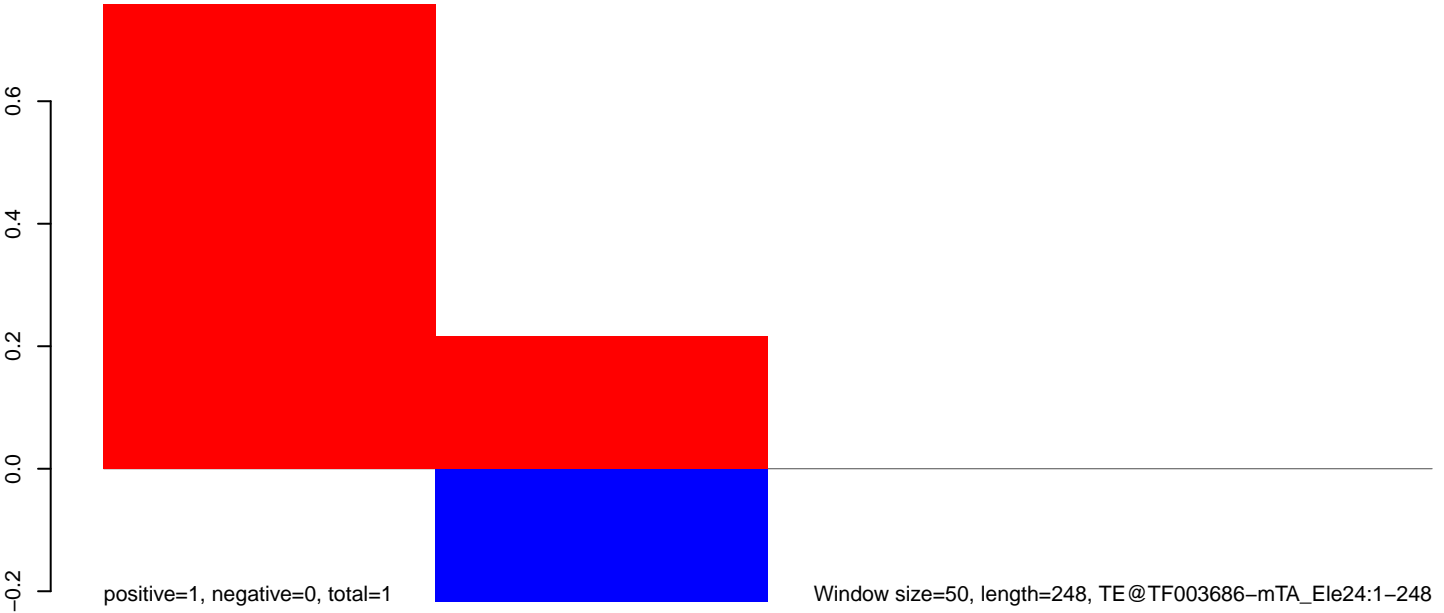
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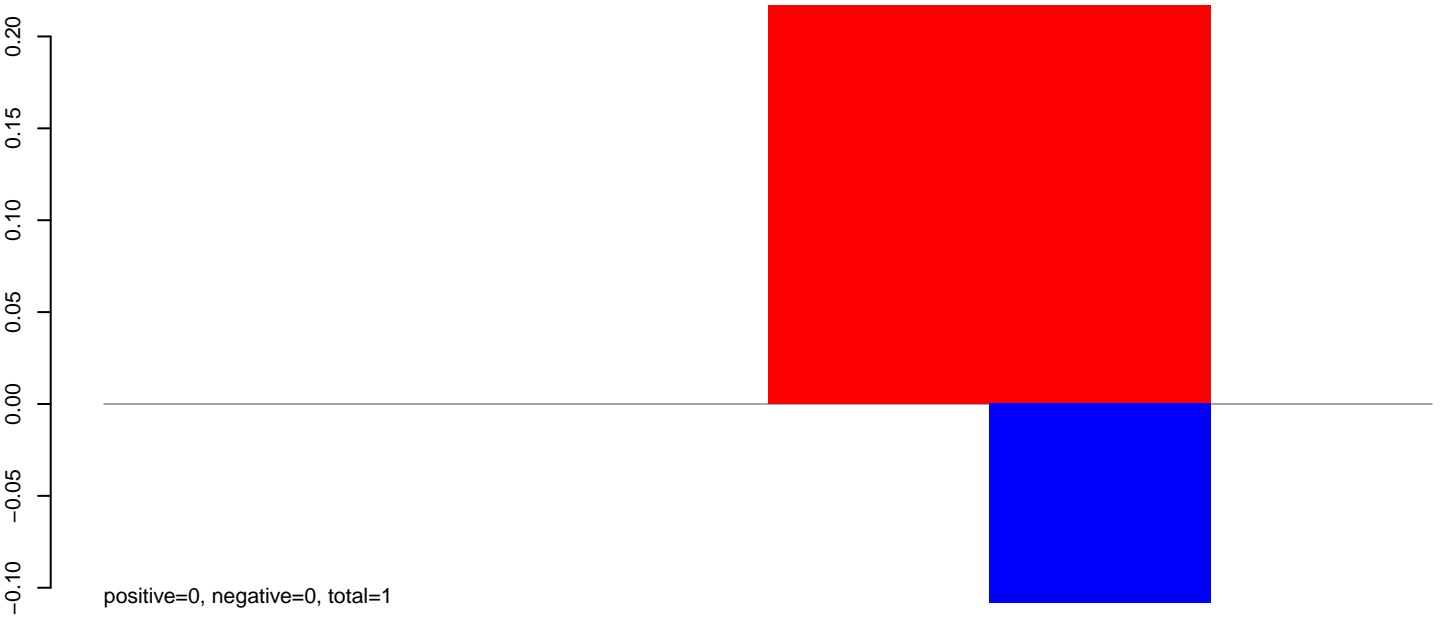
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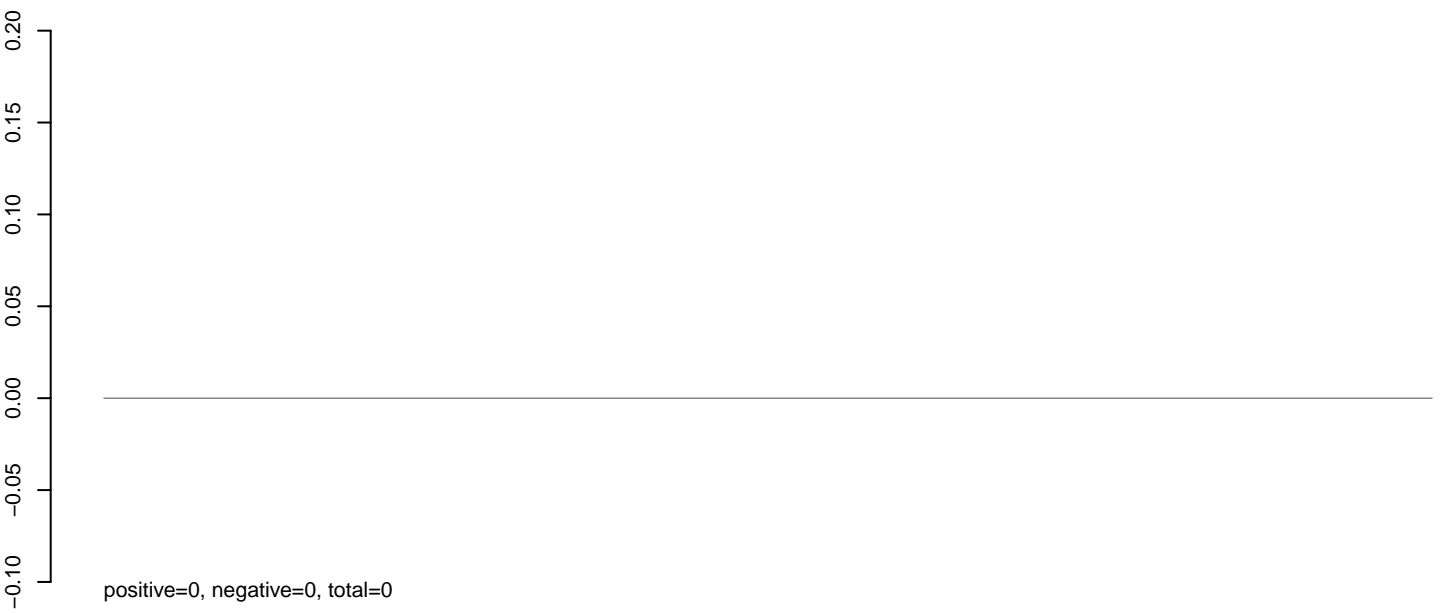
AnGam_Sua5b_cells.rep



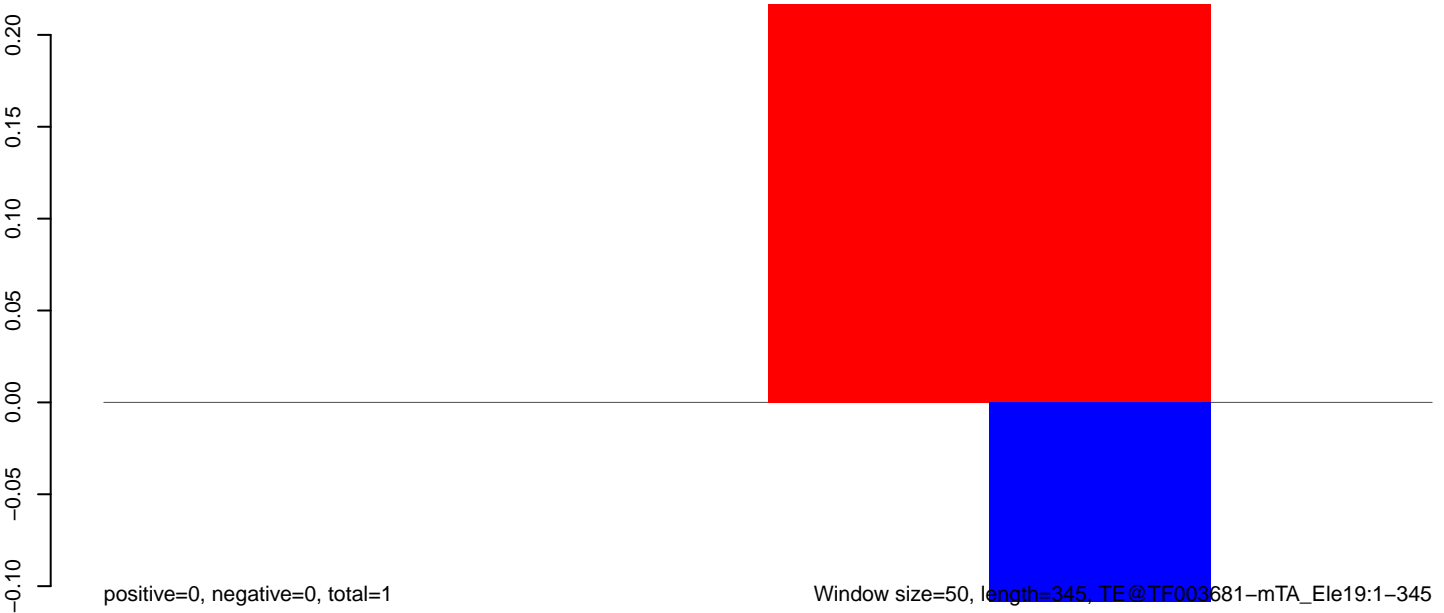
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

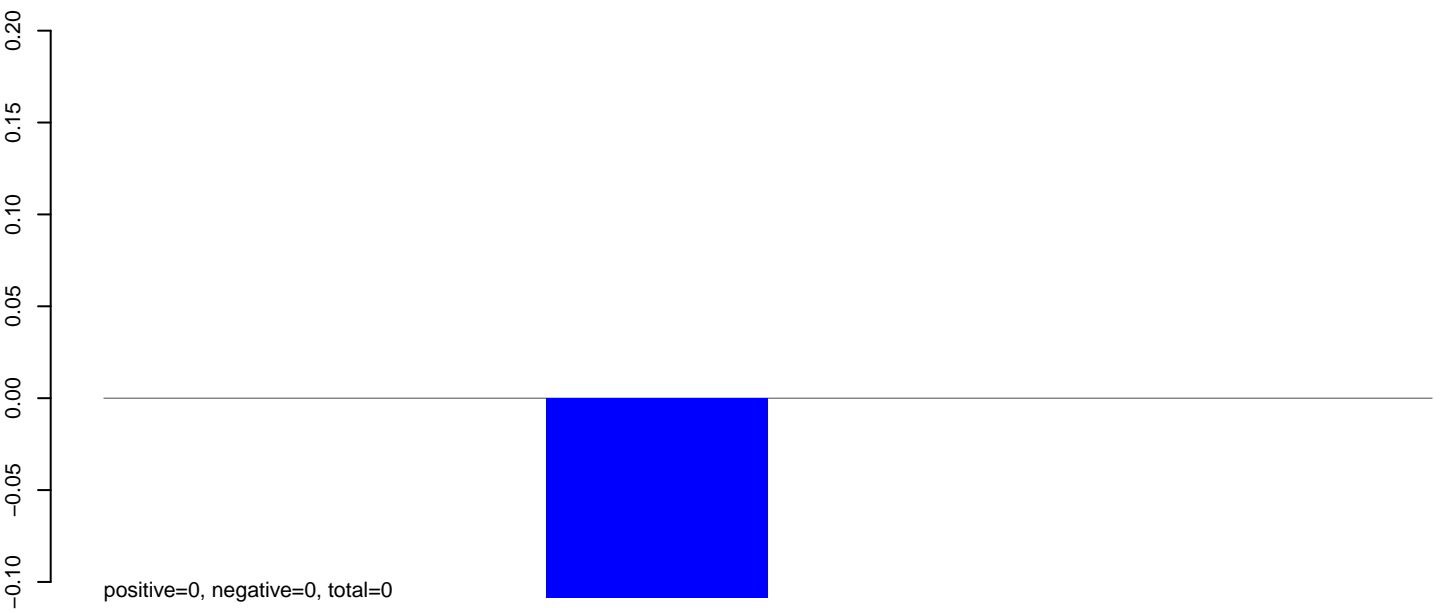


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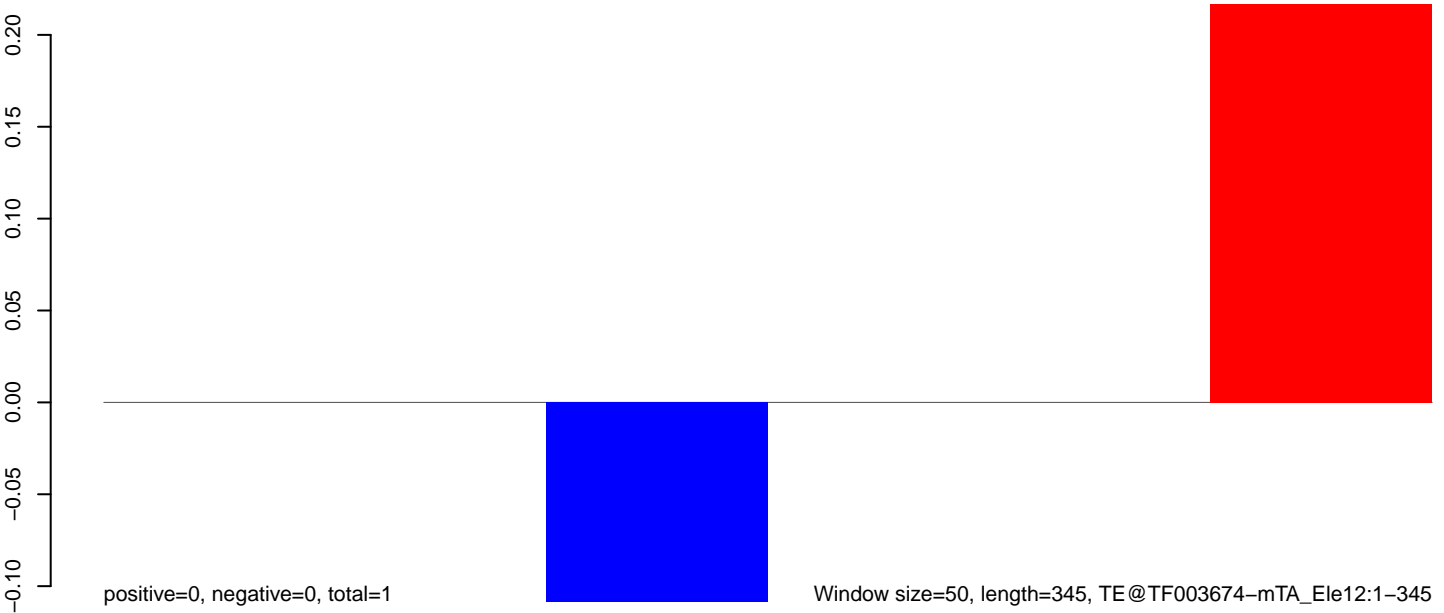
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AnGam_Sua5b_cells.24_35.rep

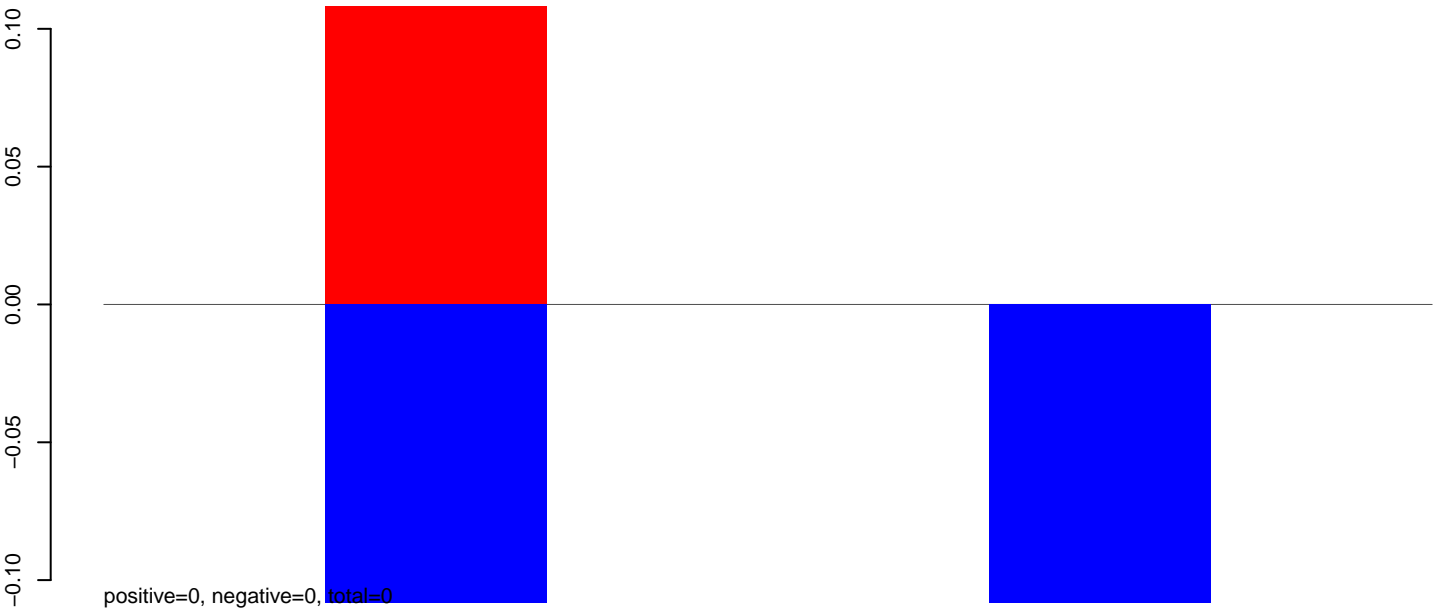


AnGam_Sua5b_cells.rep

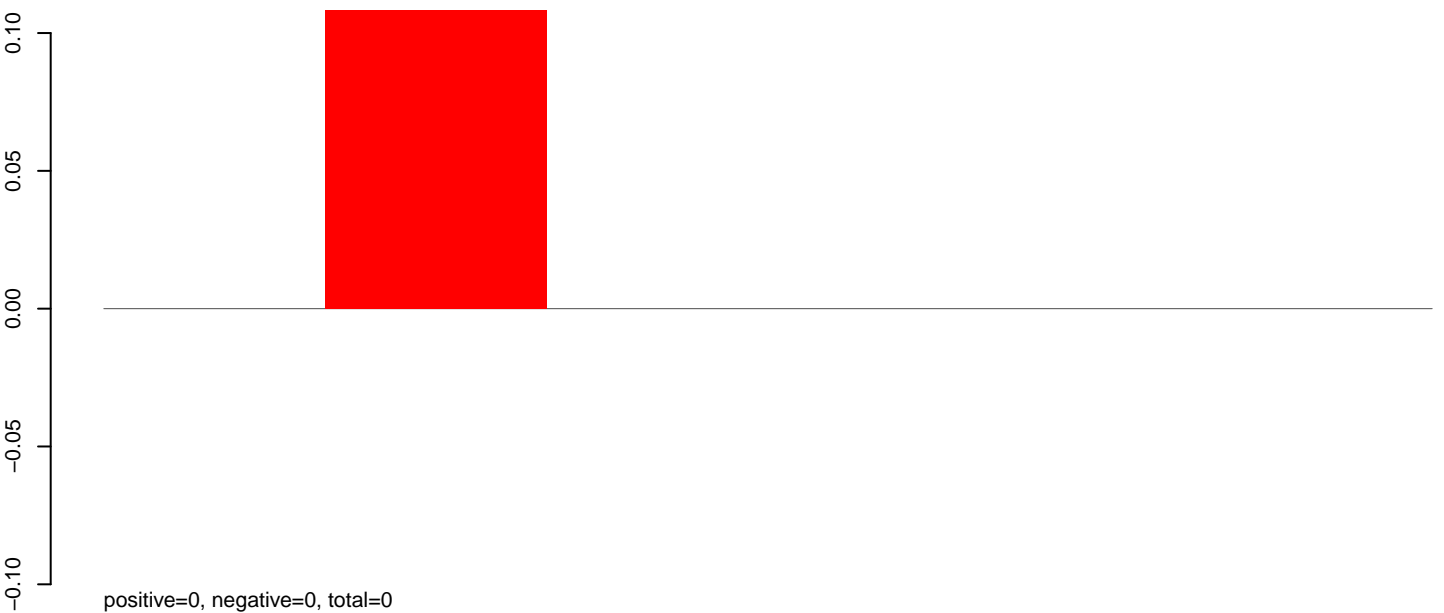


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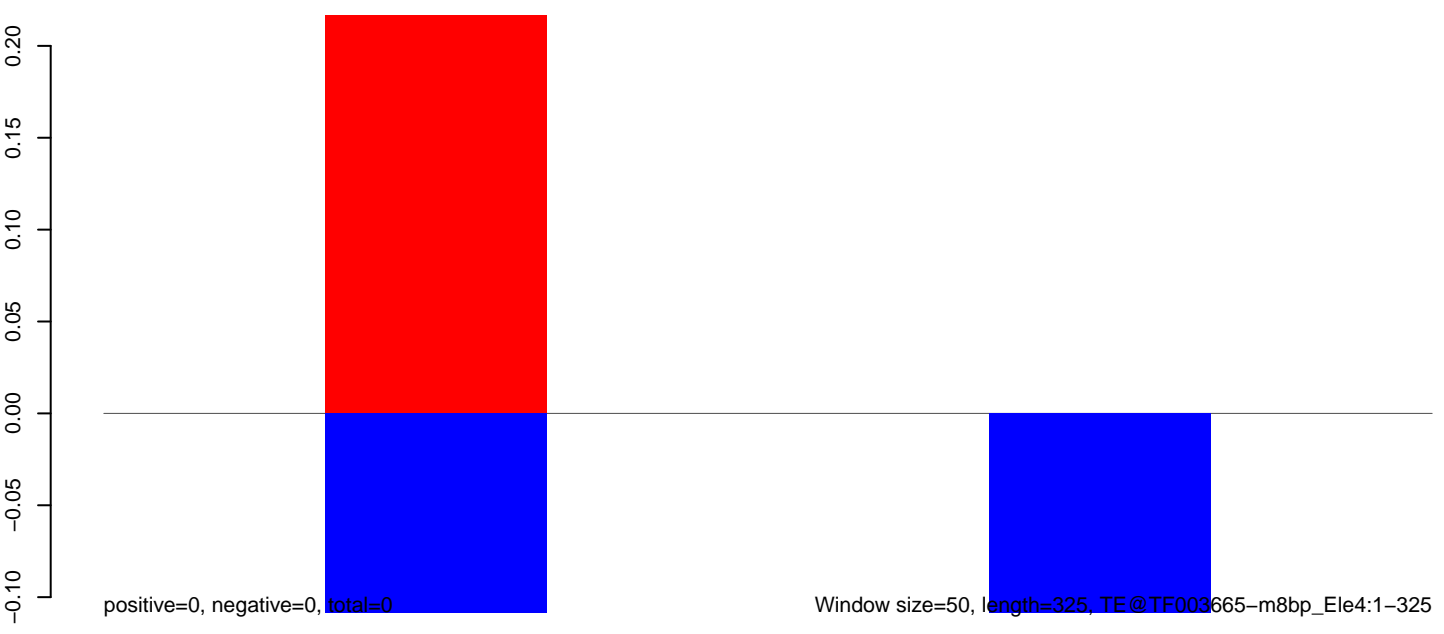
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

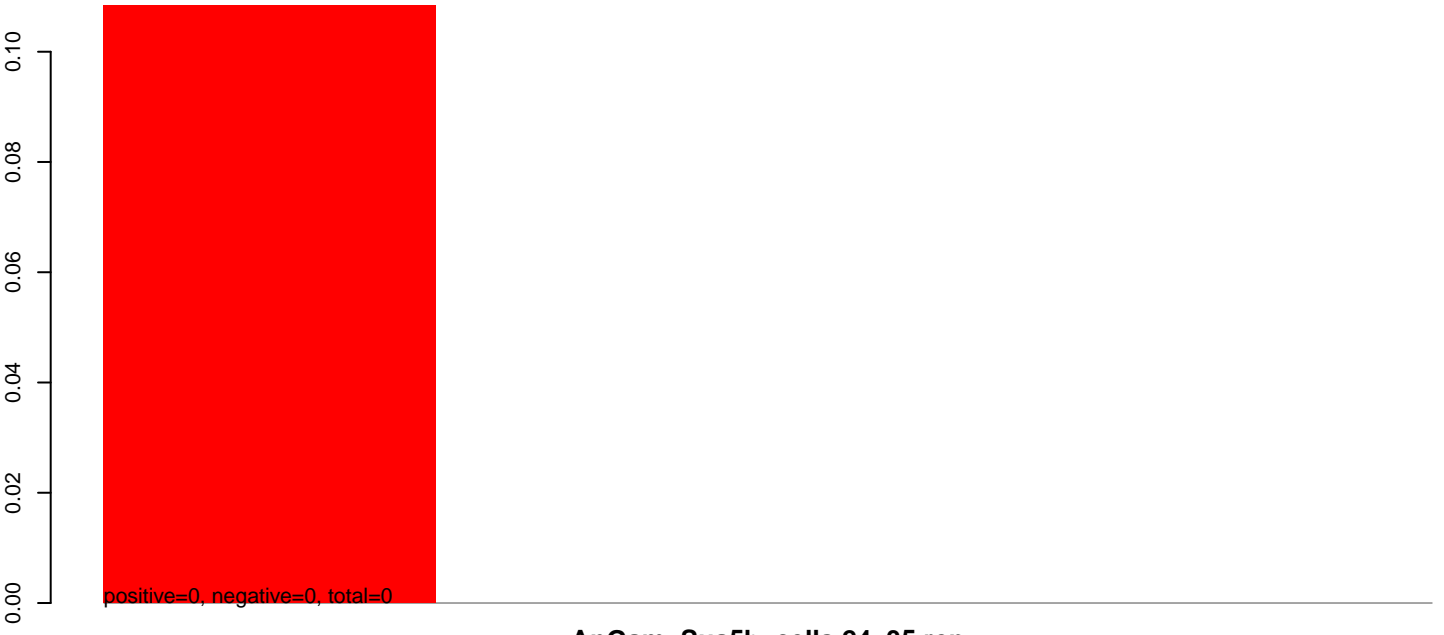


AnGam_Sua5b_cells.rep

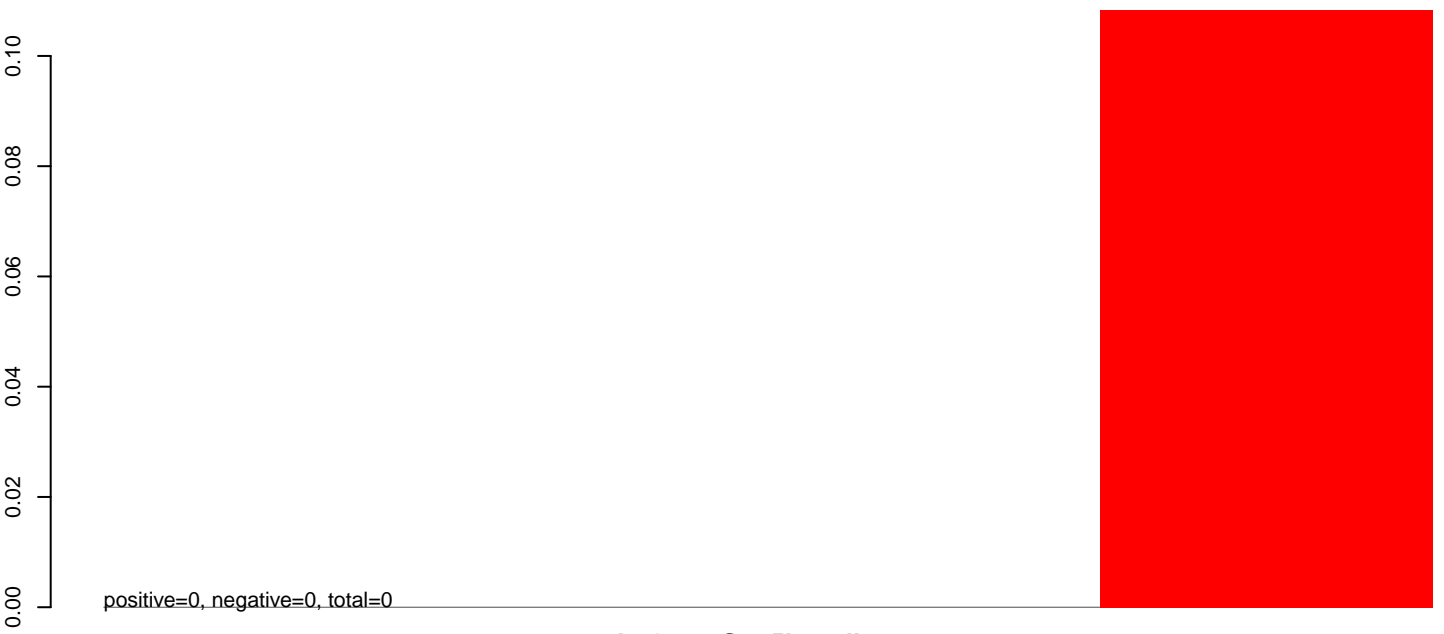


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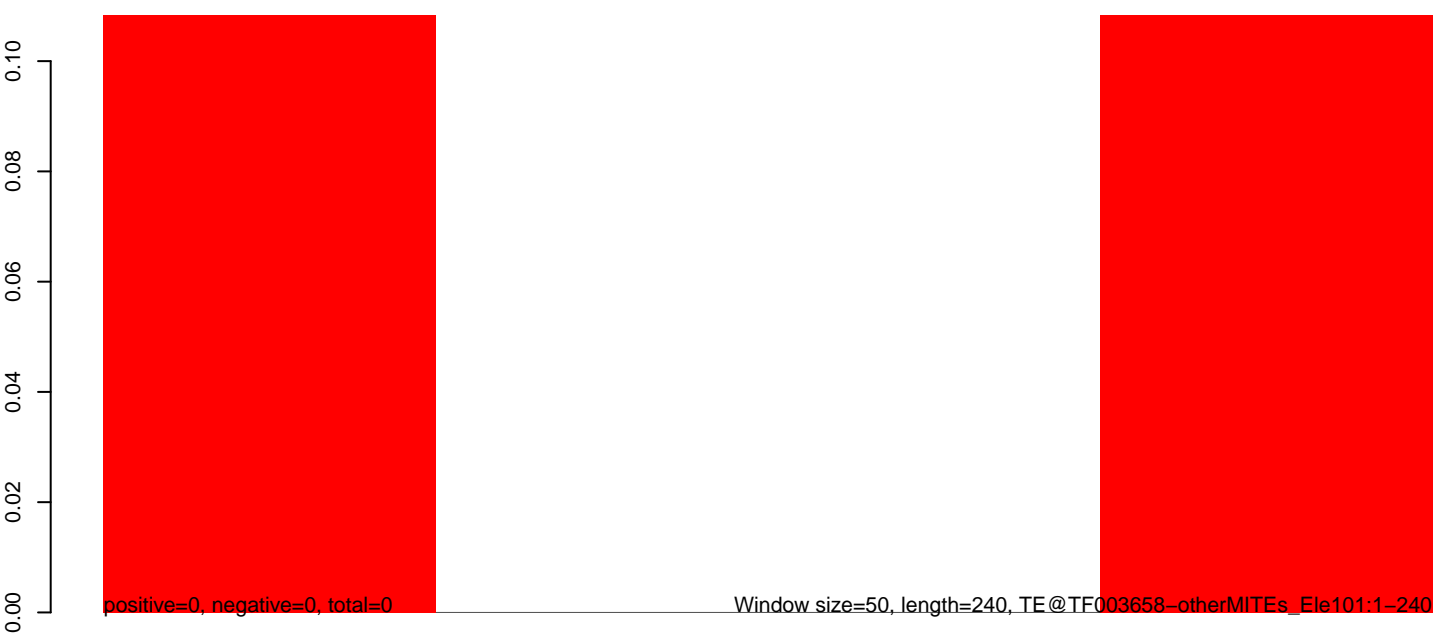
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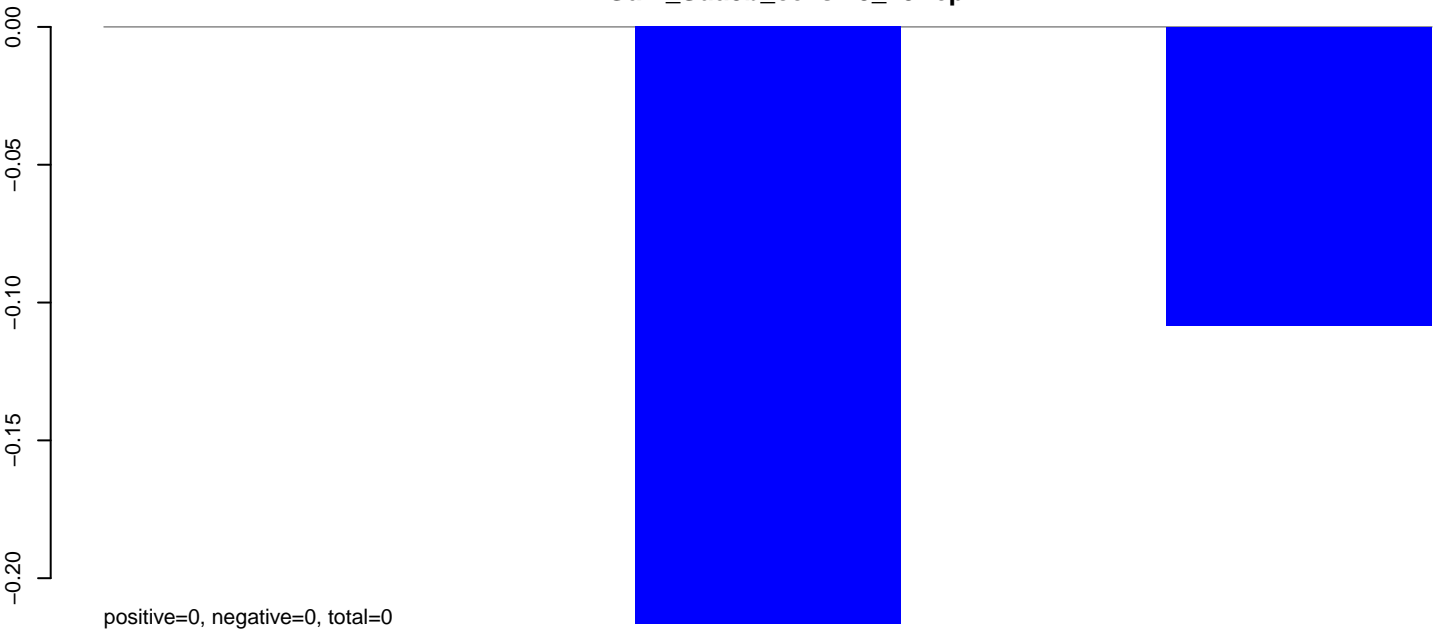
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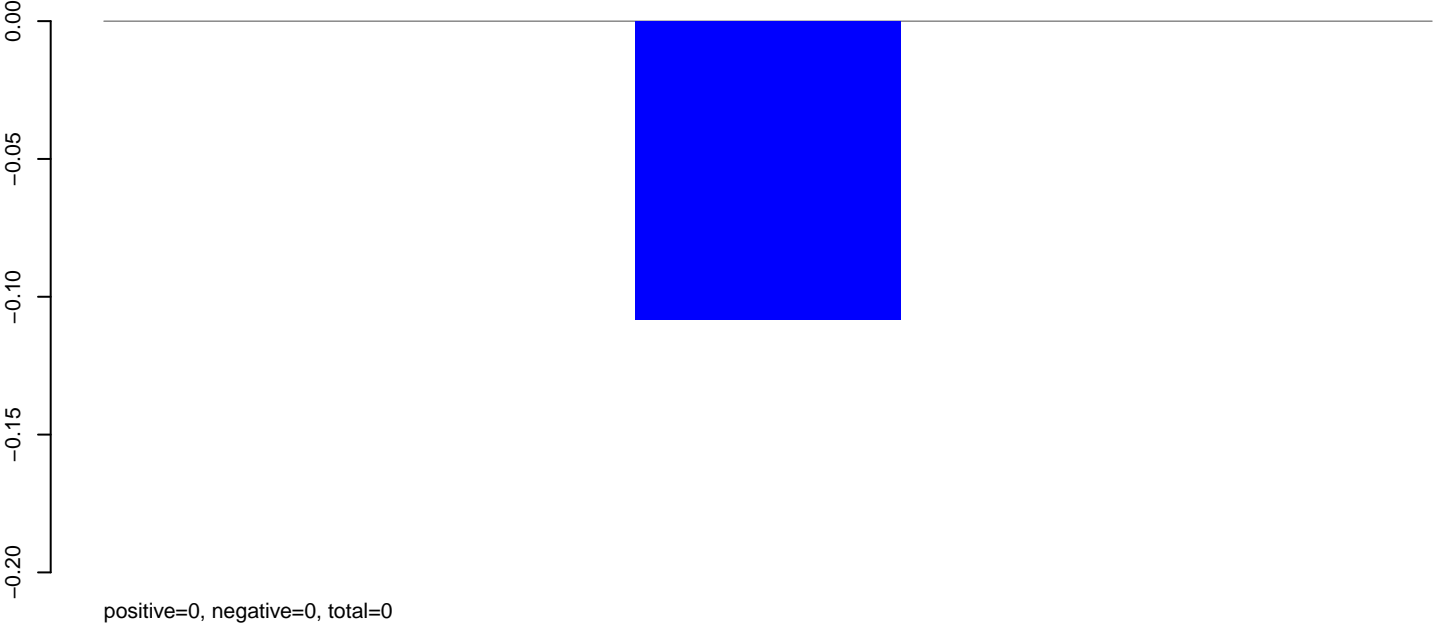
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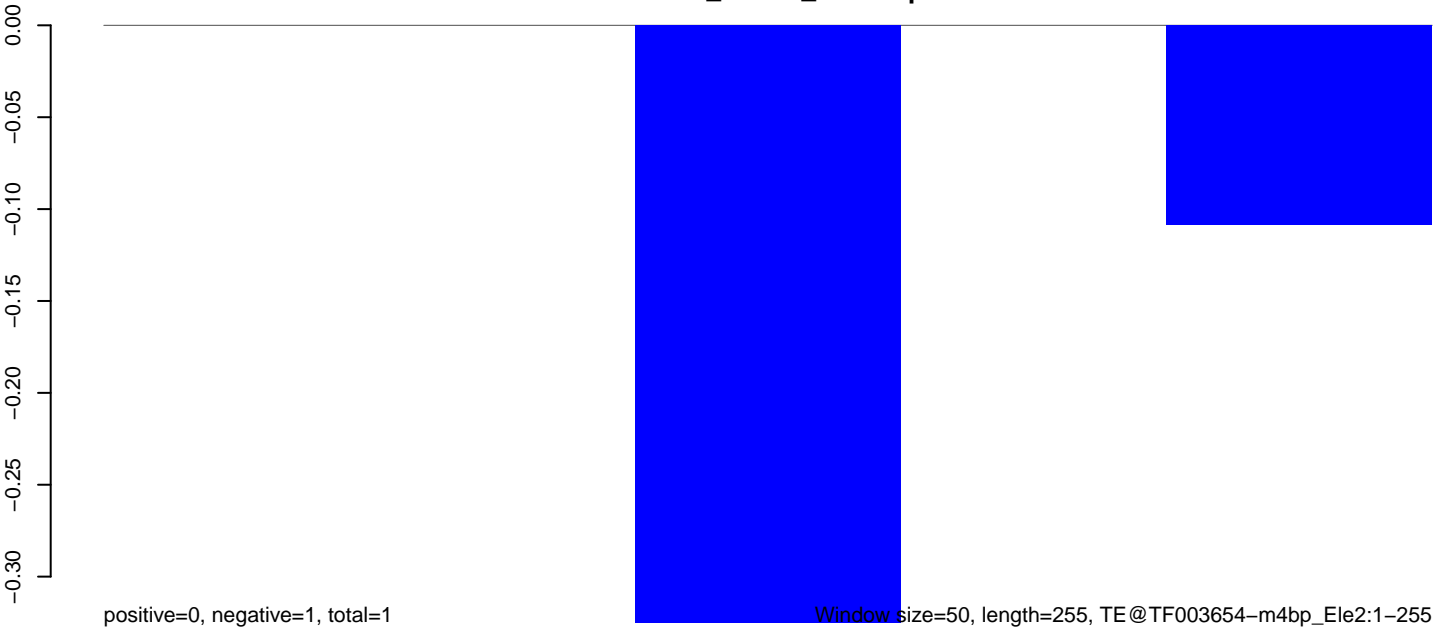
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AnGam_Sua5b_cells.24_35.rep



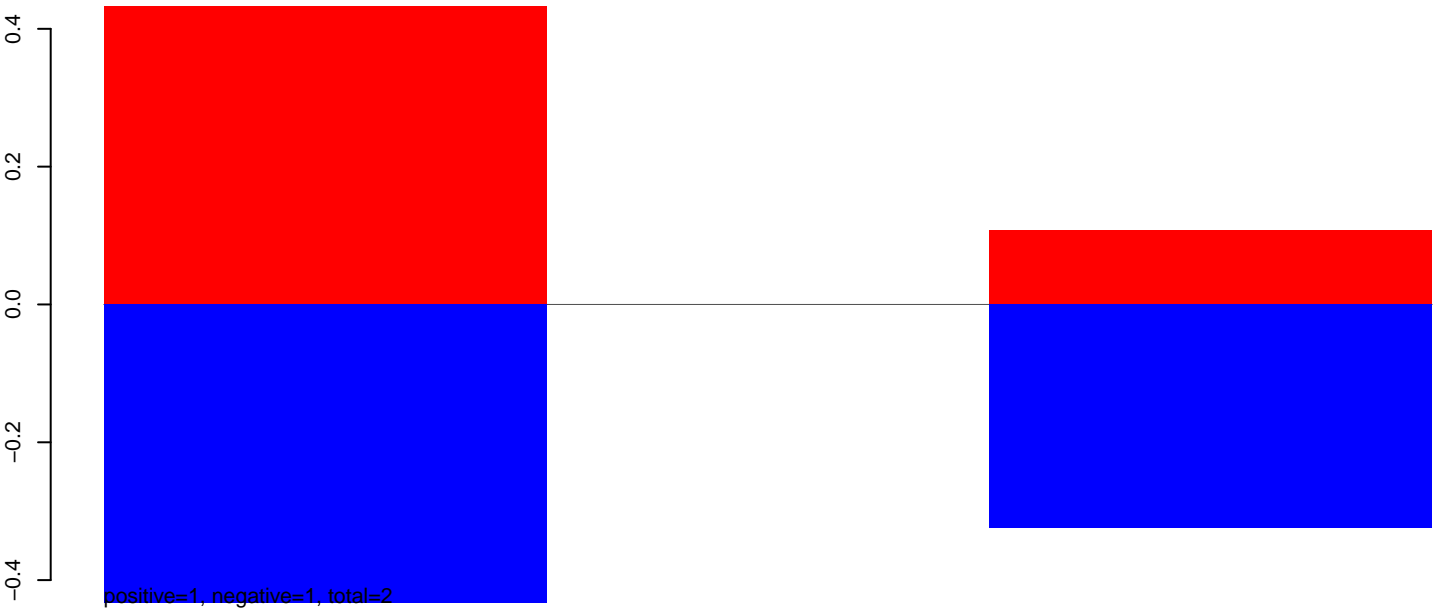
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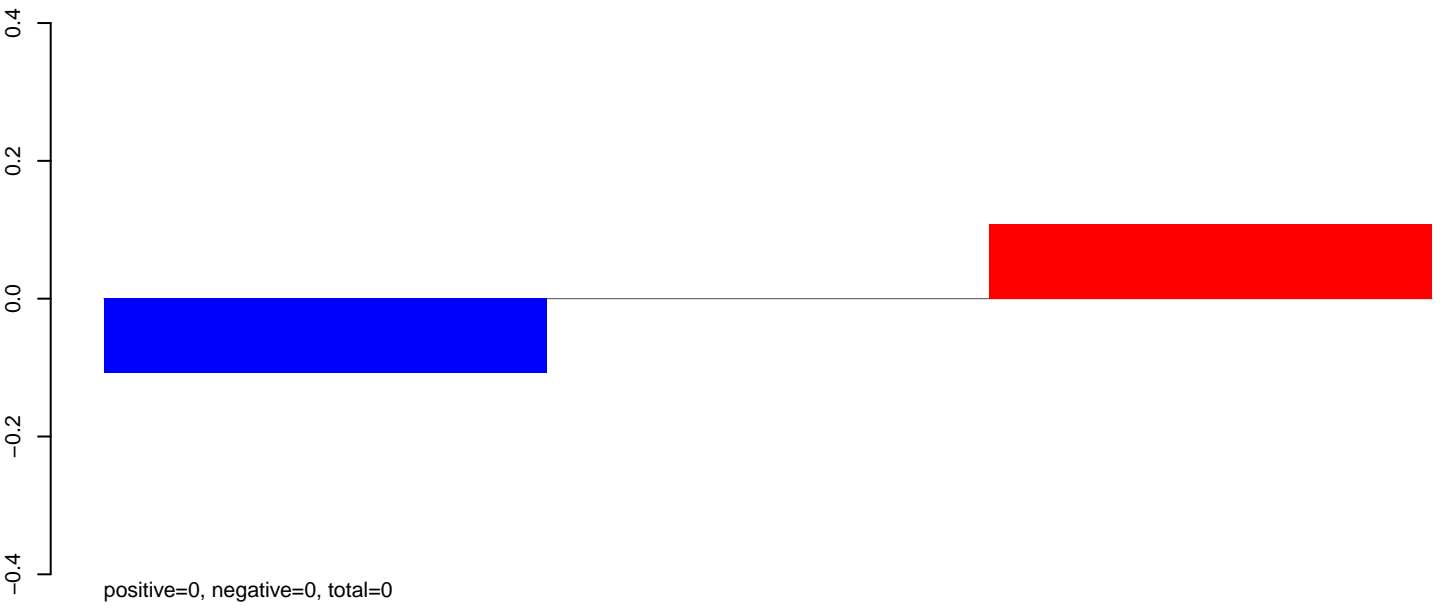
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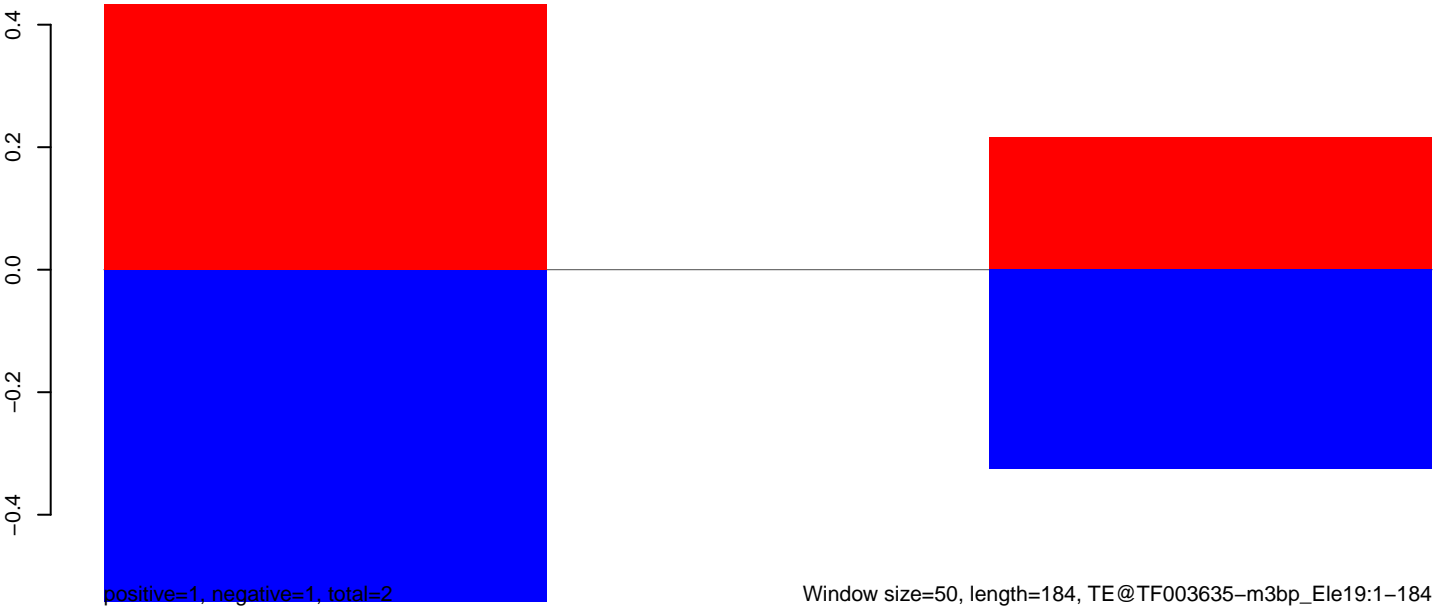
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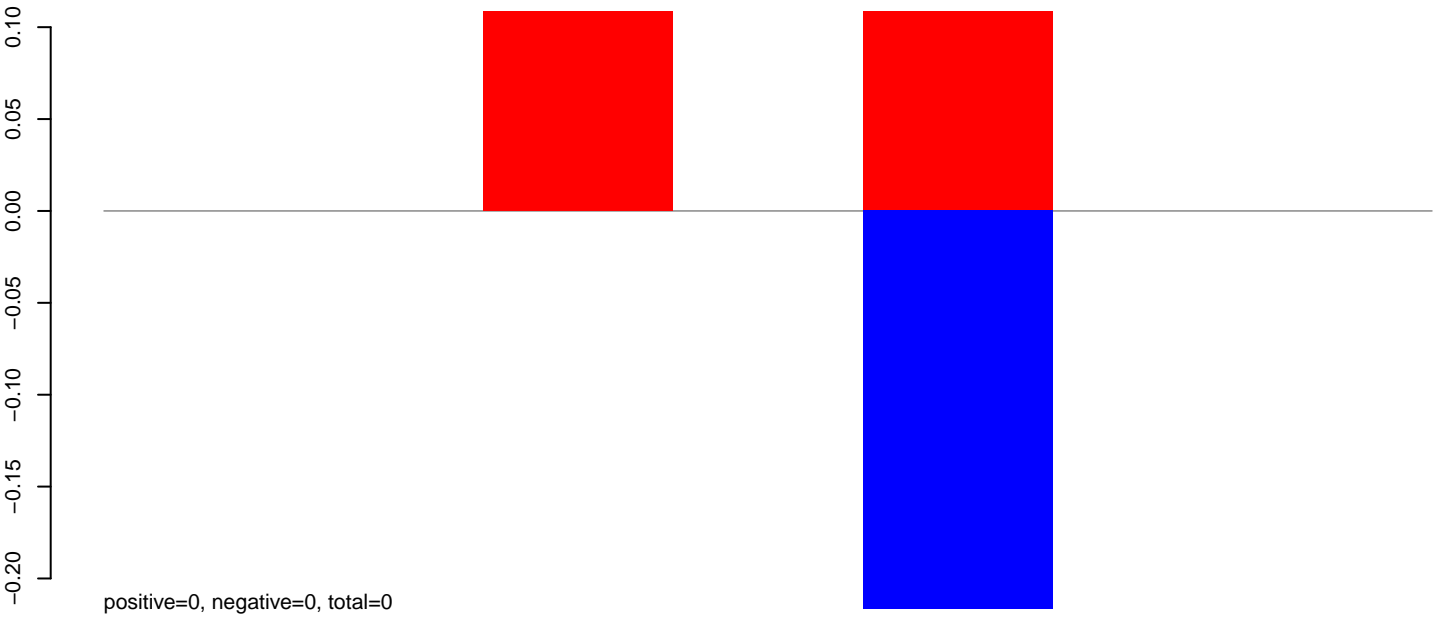
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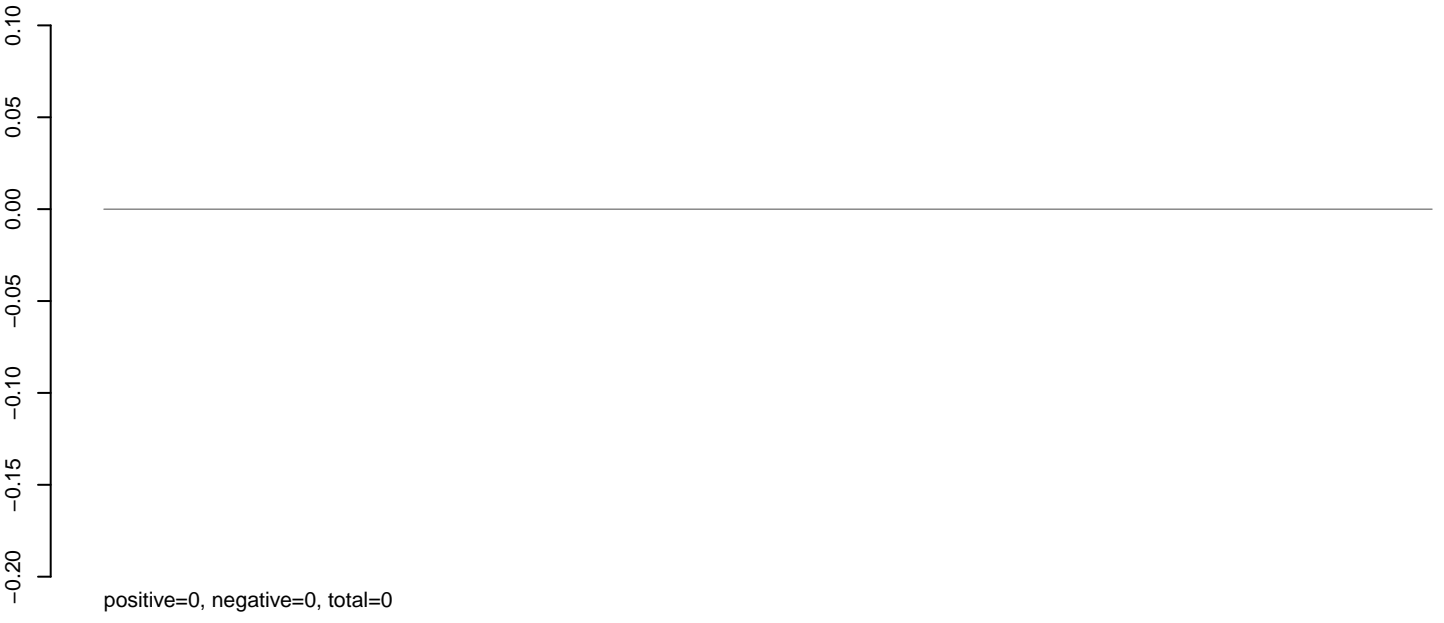
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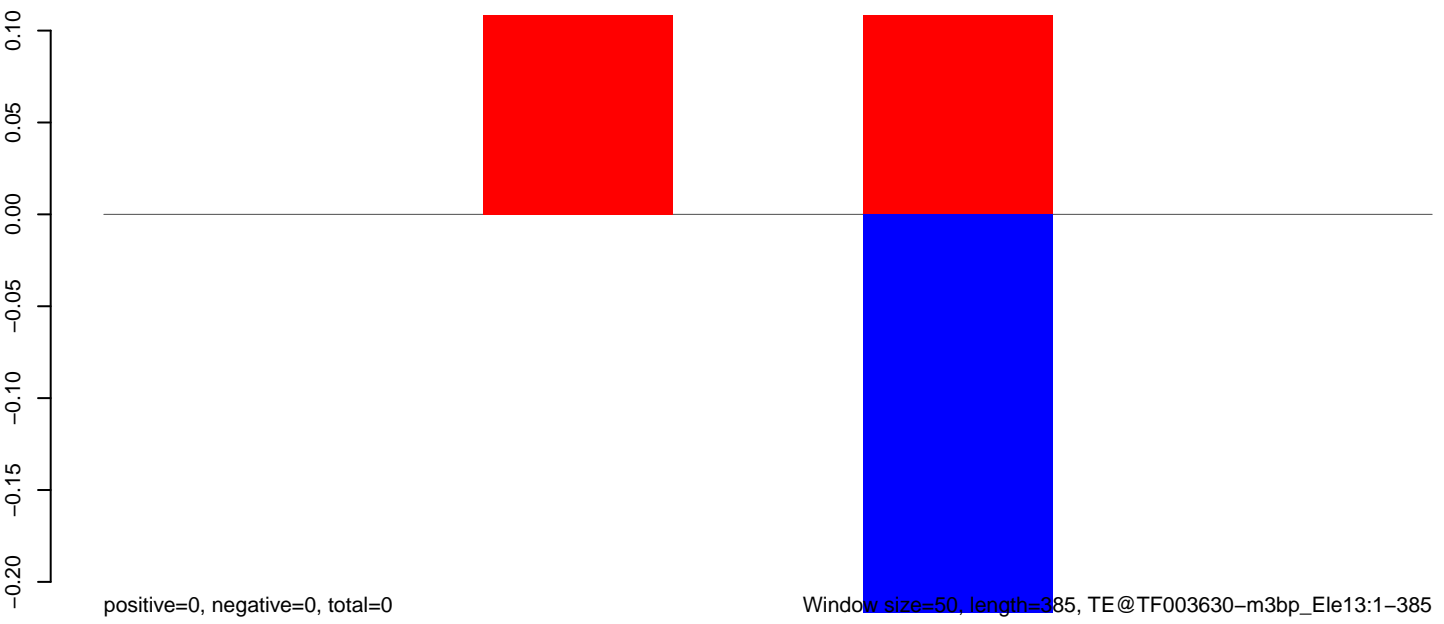
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AnGam_Sua5b_cells.24_35.rep



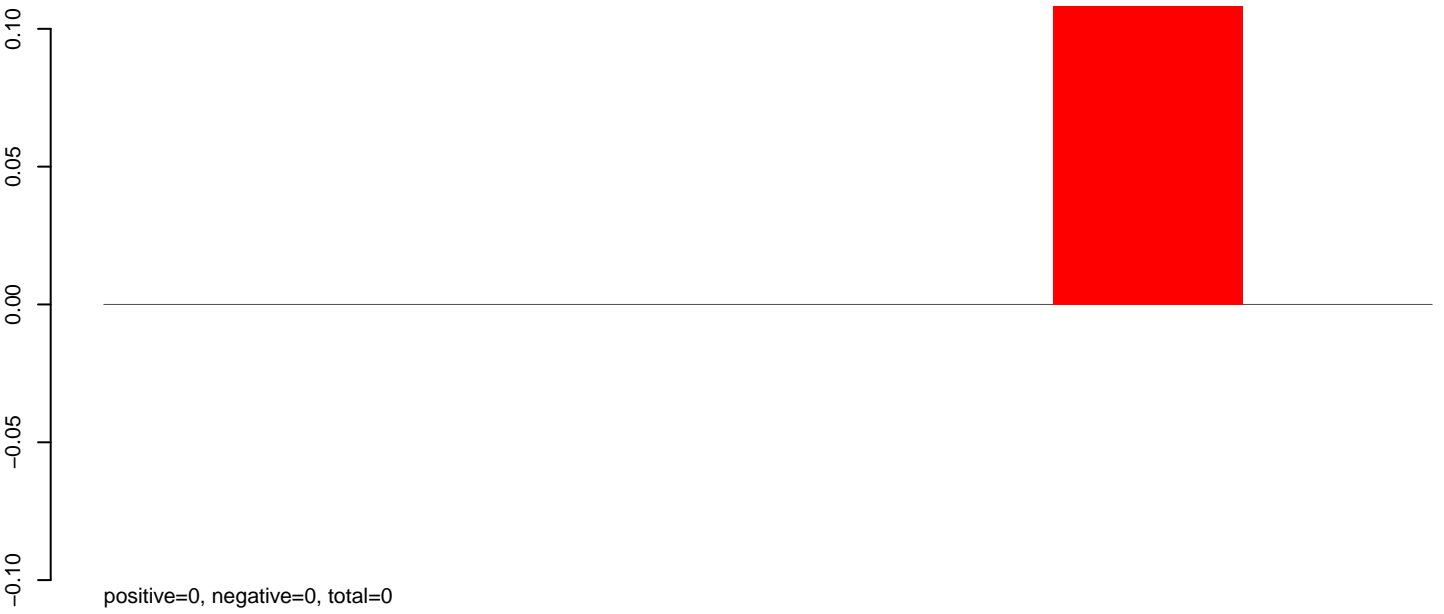
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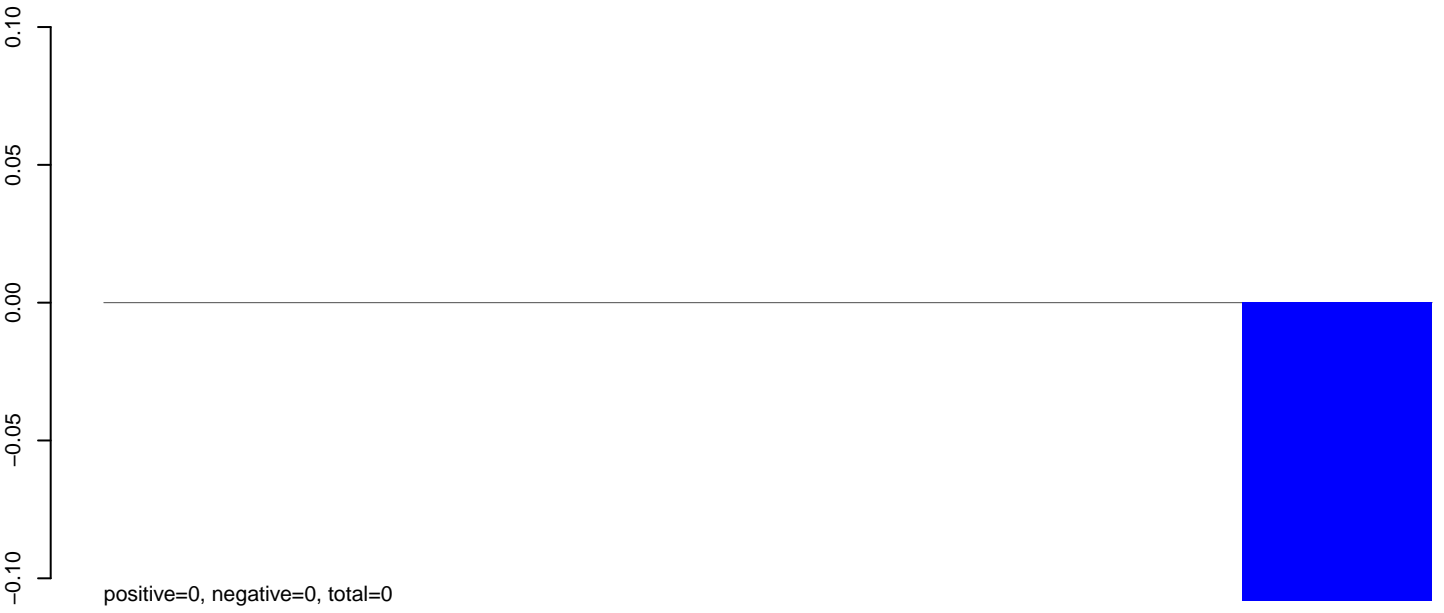
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AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep

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-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.rep

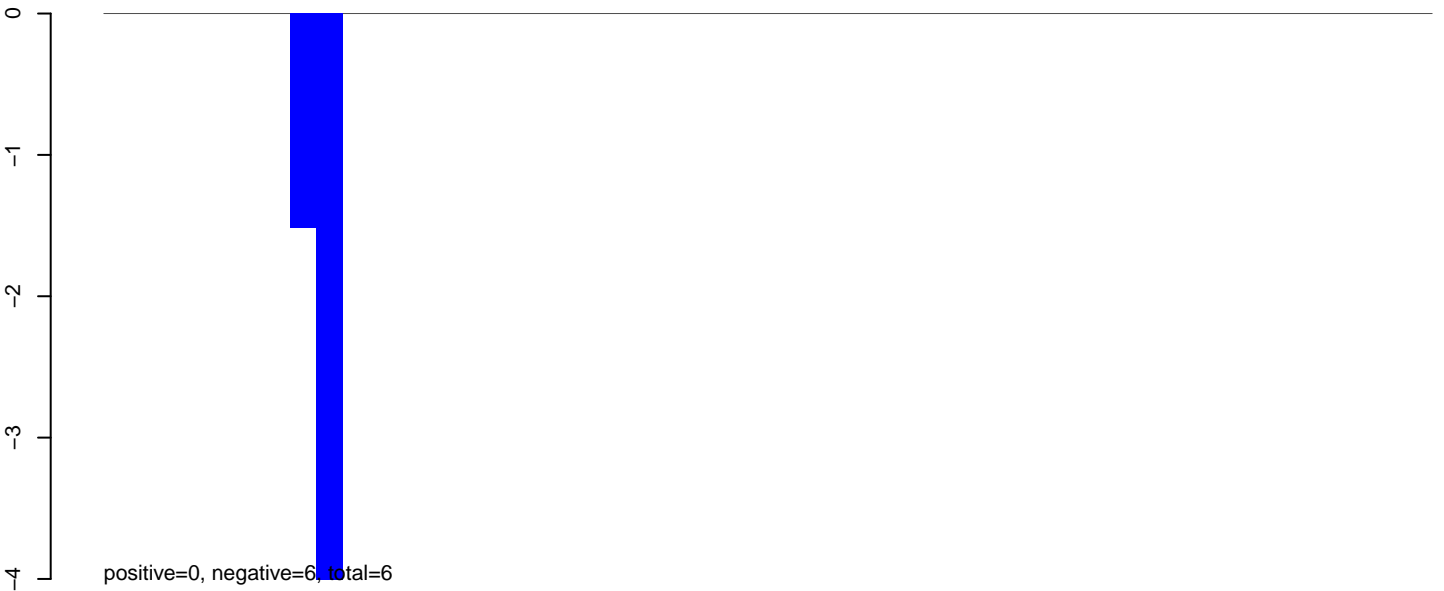
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positive=0, negative=0, total=0

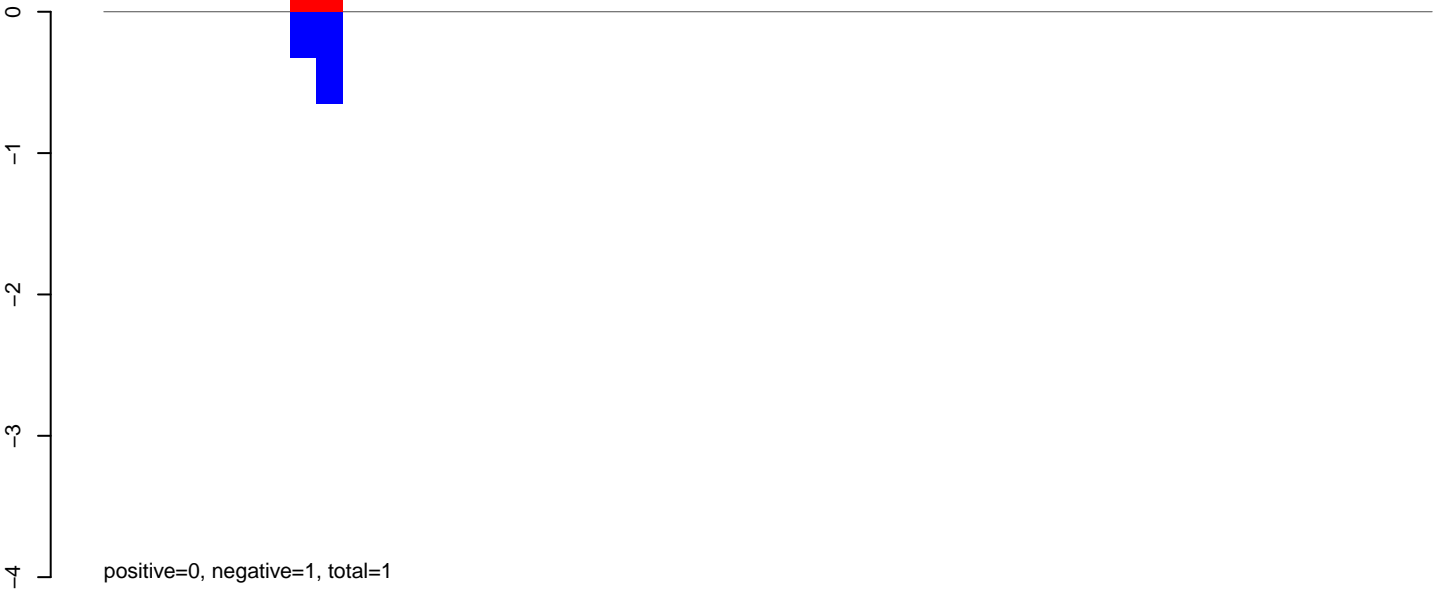
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AnGam_Sua5b_cells.18_23.rep



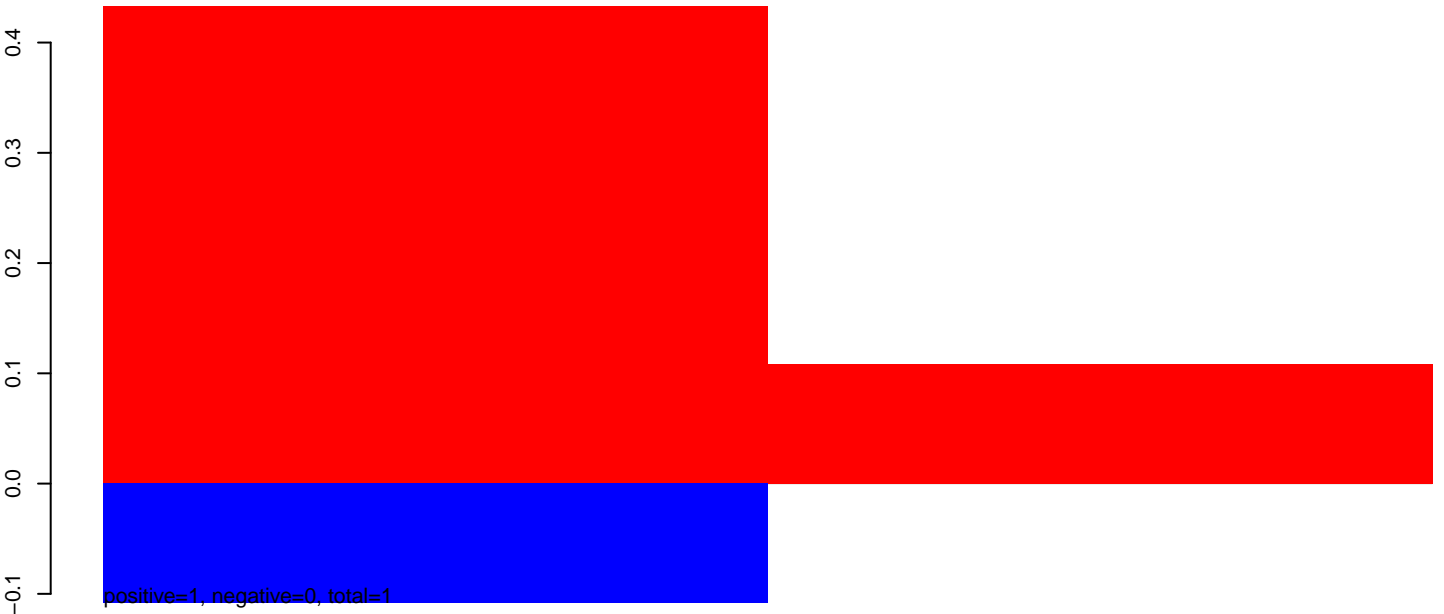
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



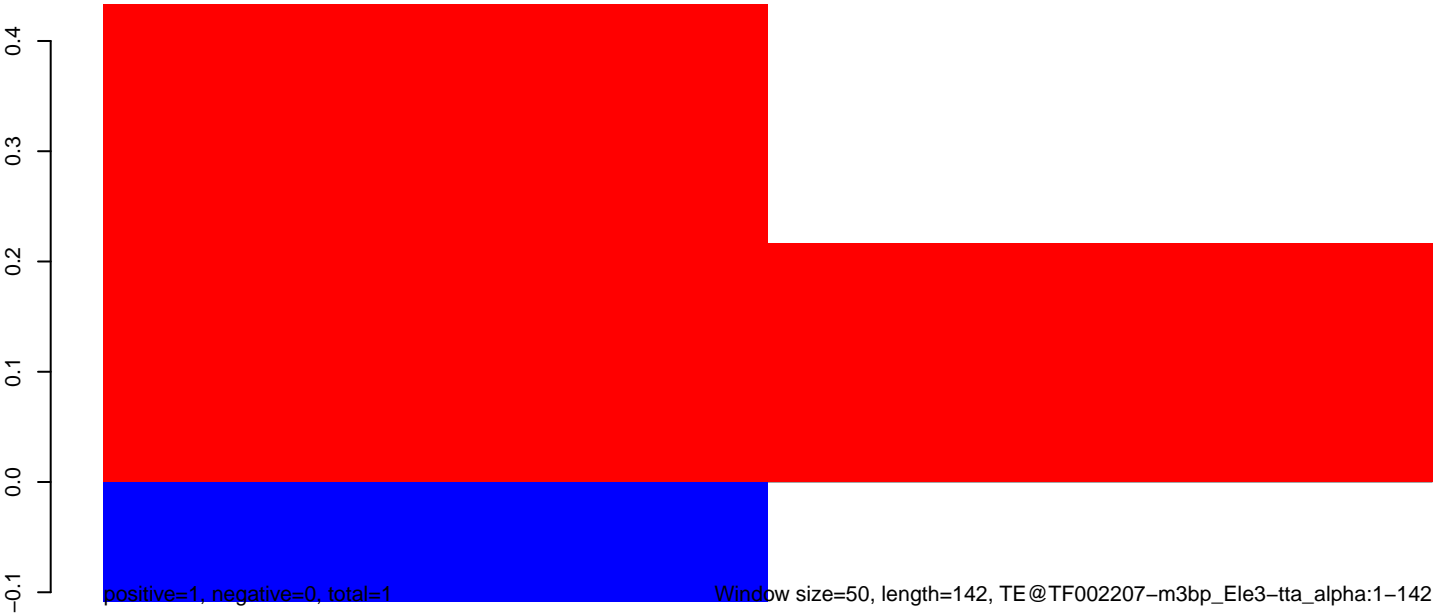
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AnGam_Sua5b_cells.24_35.rep

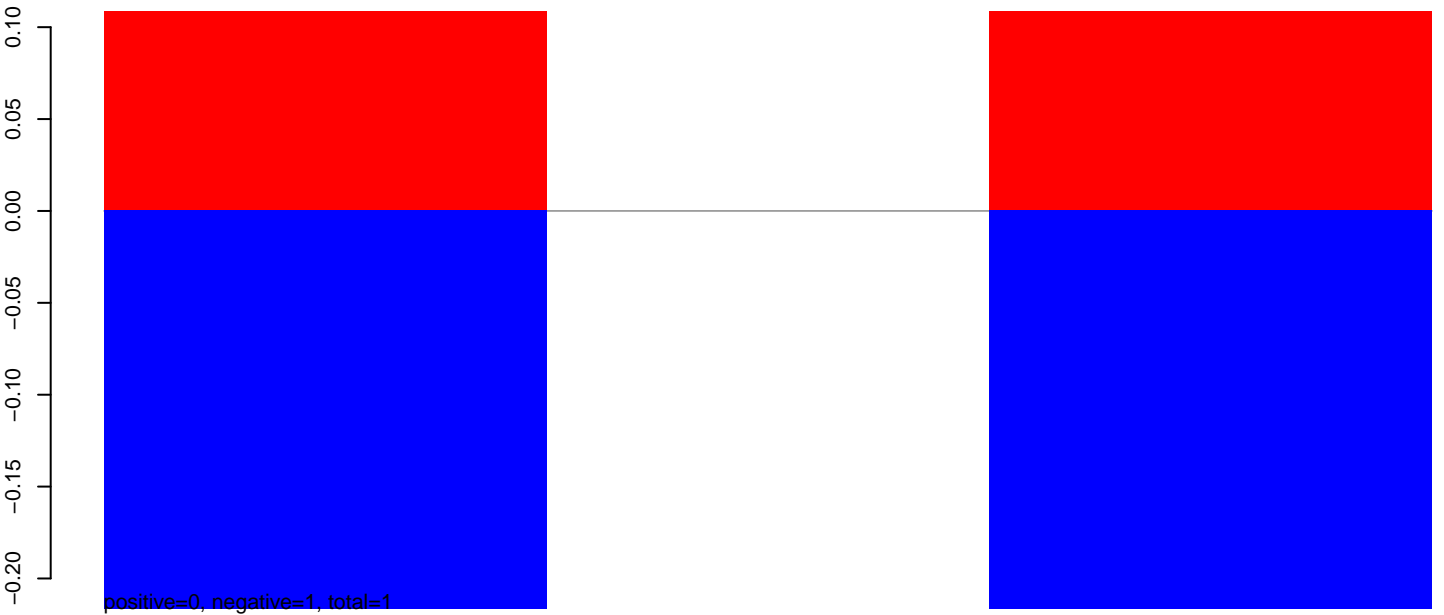


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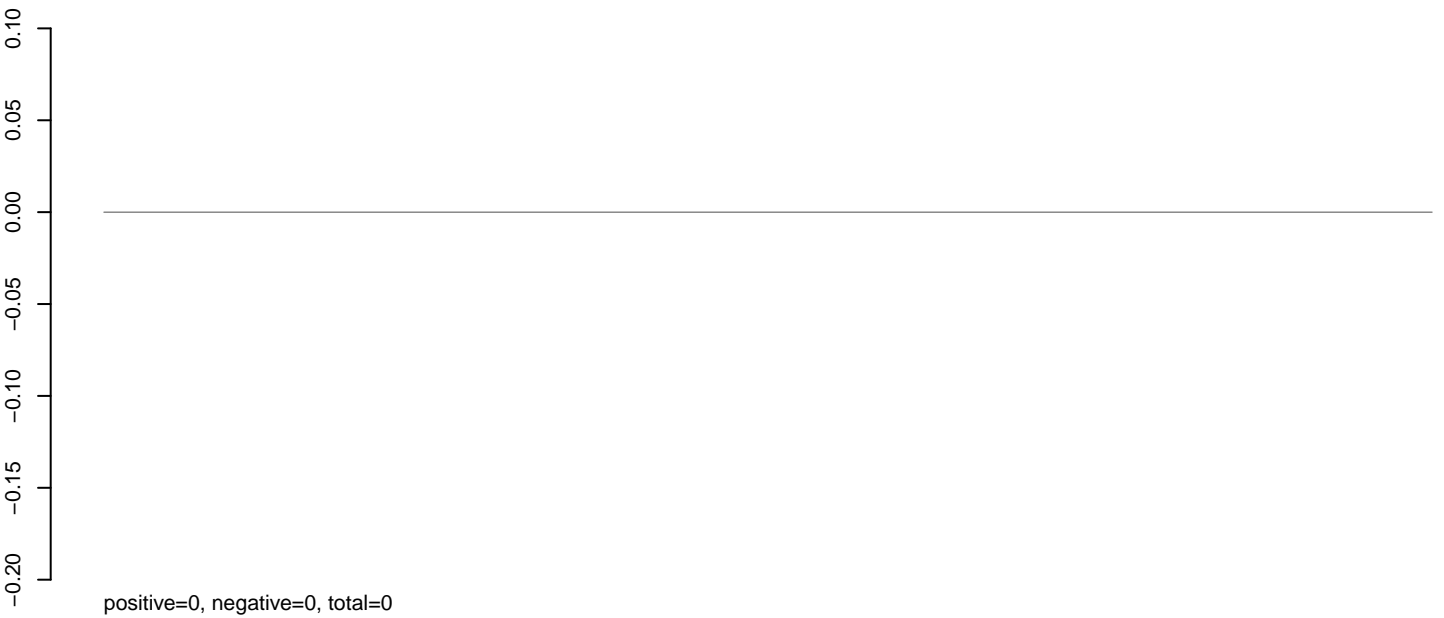


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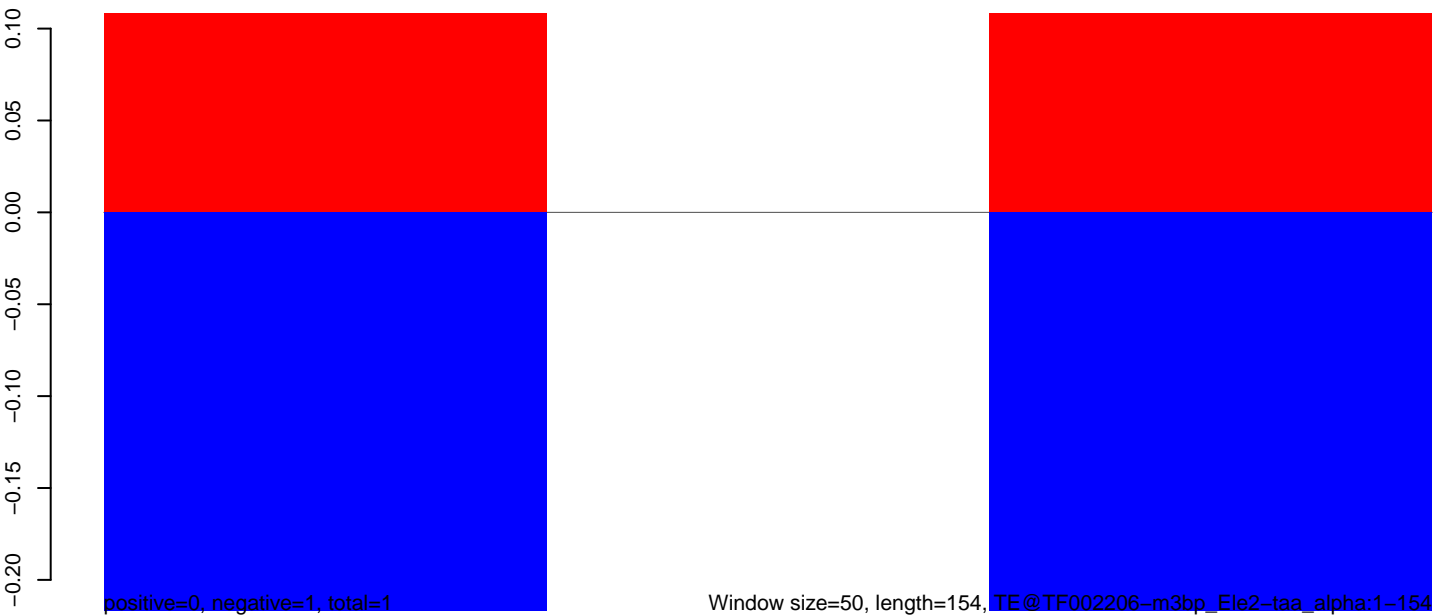
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

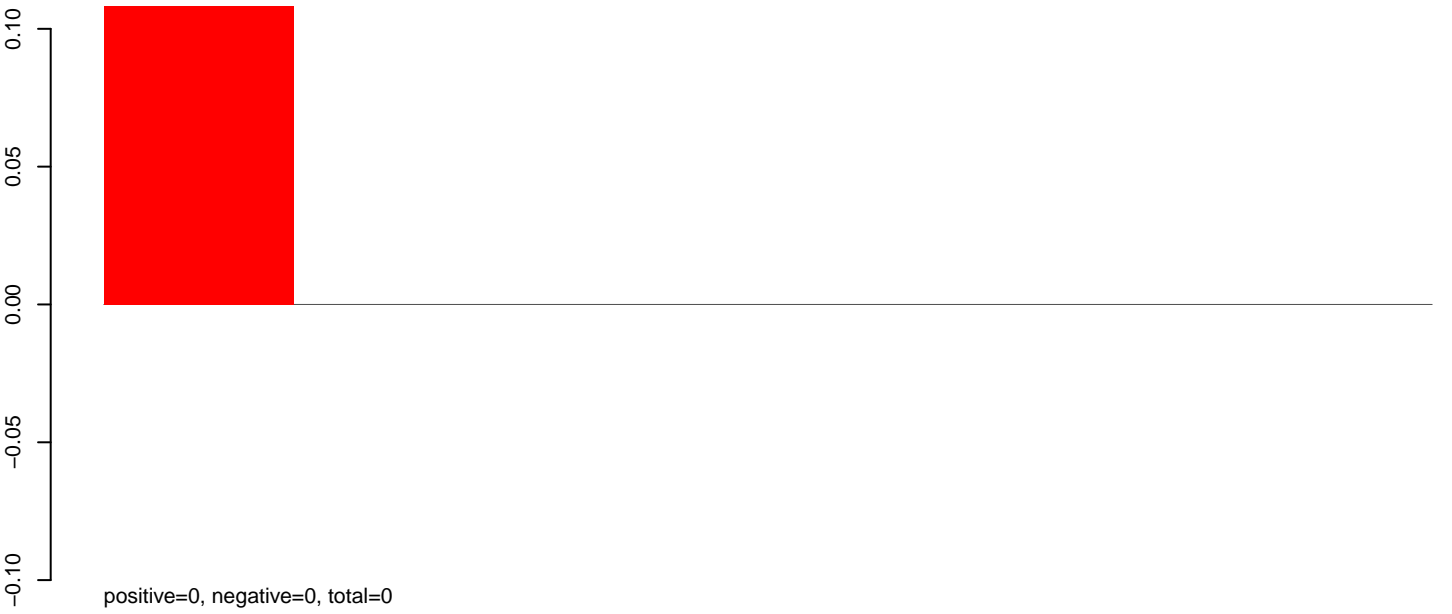


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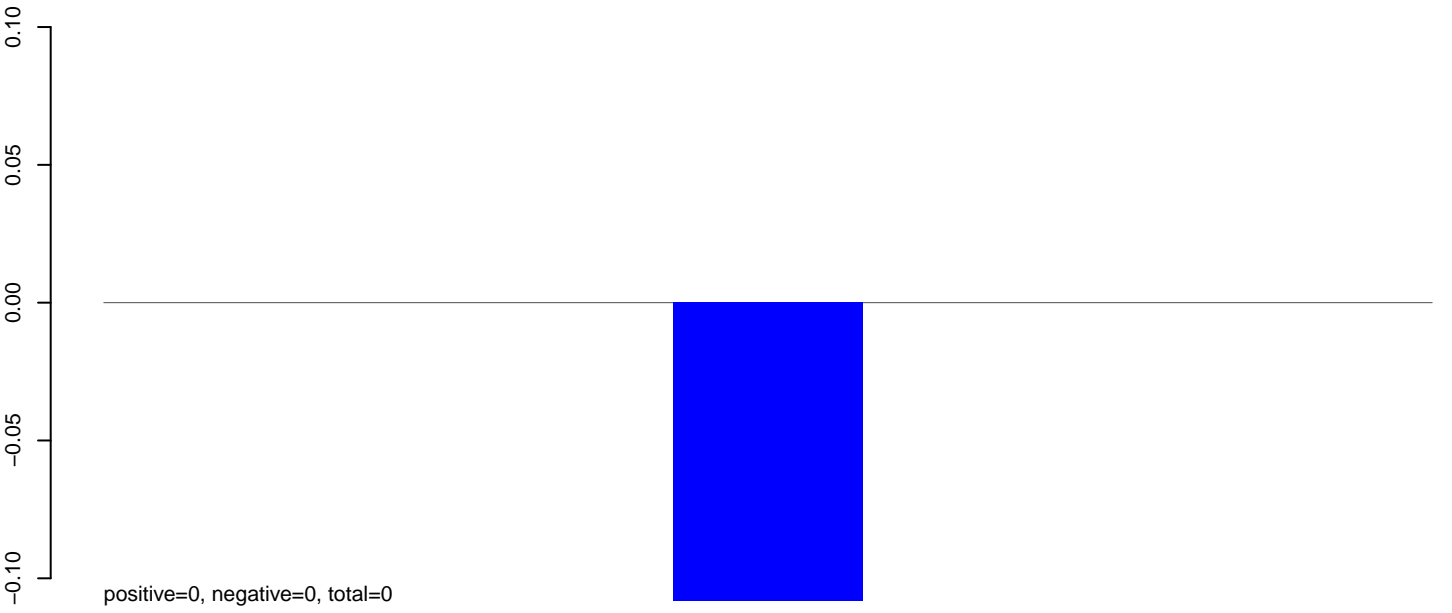


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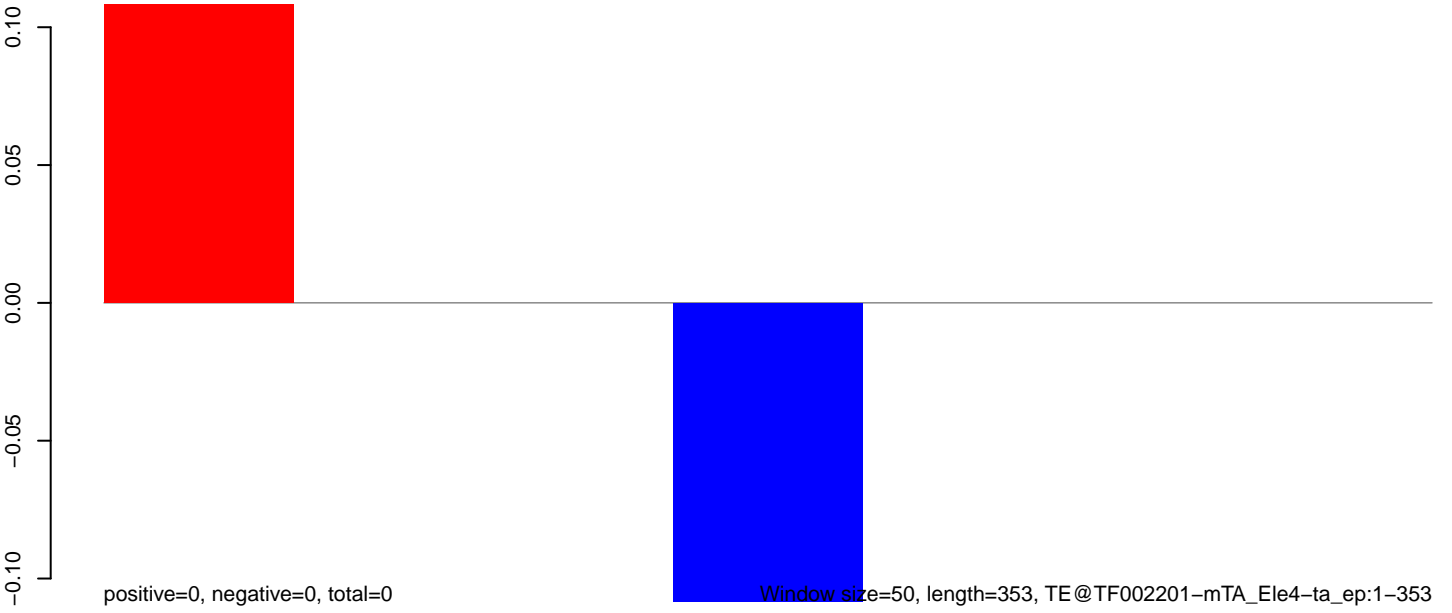
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AnGam_Sua5b_cells.24_35.rep

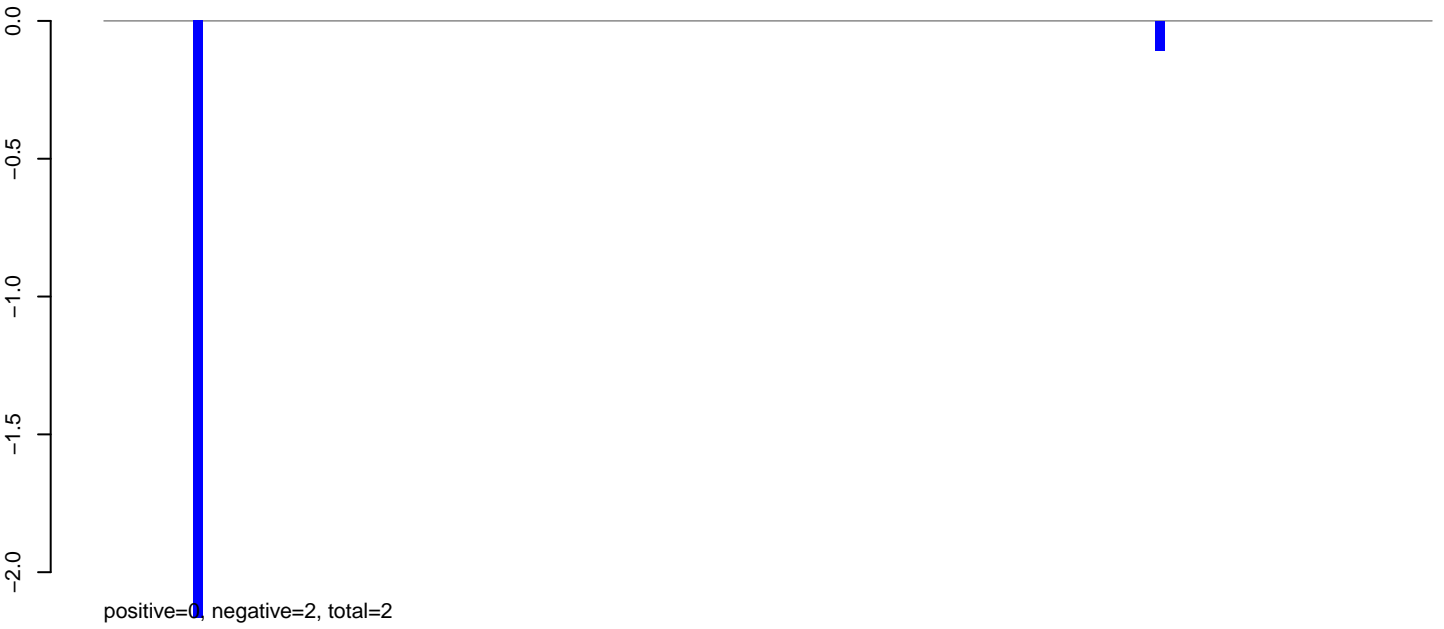


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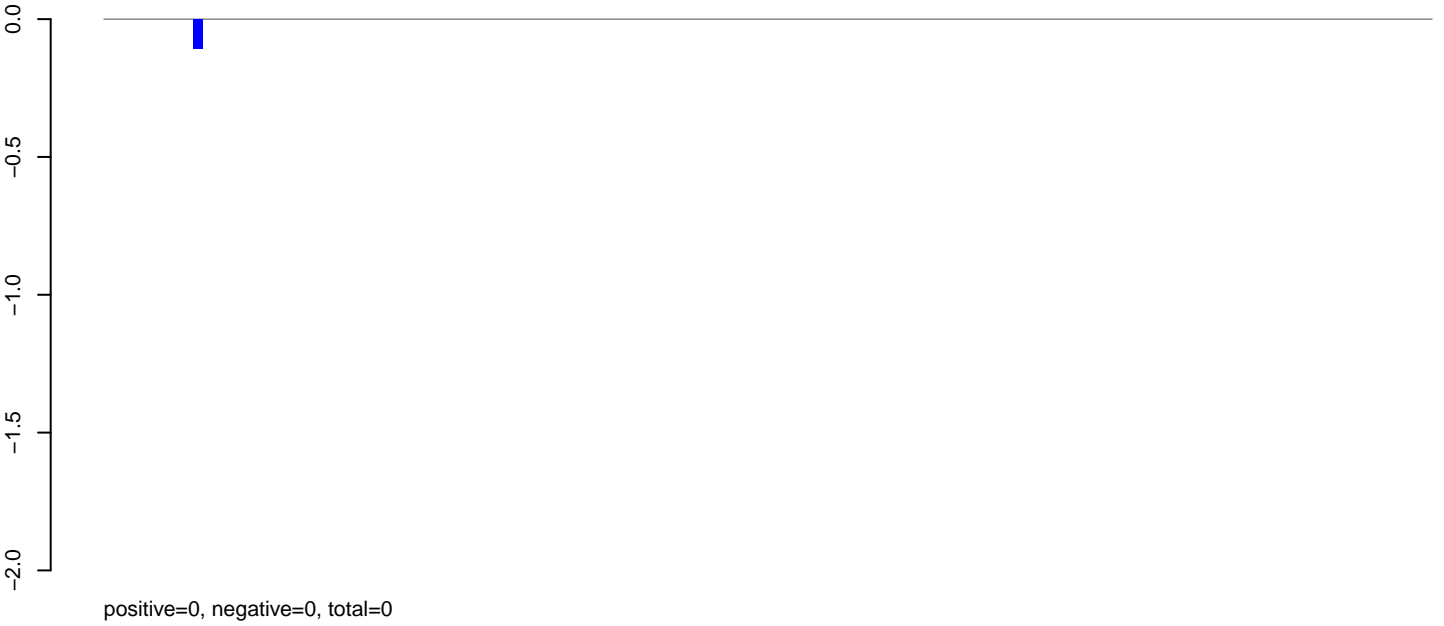


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AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



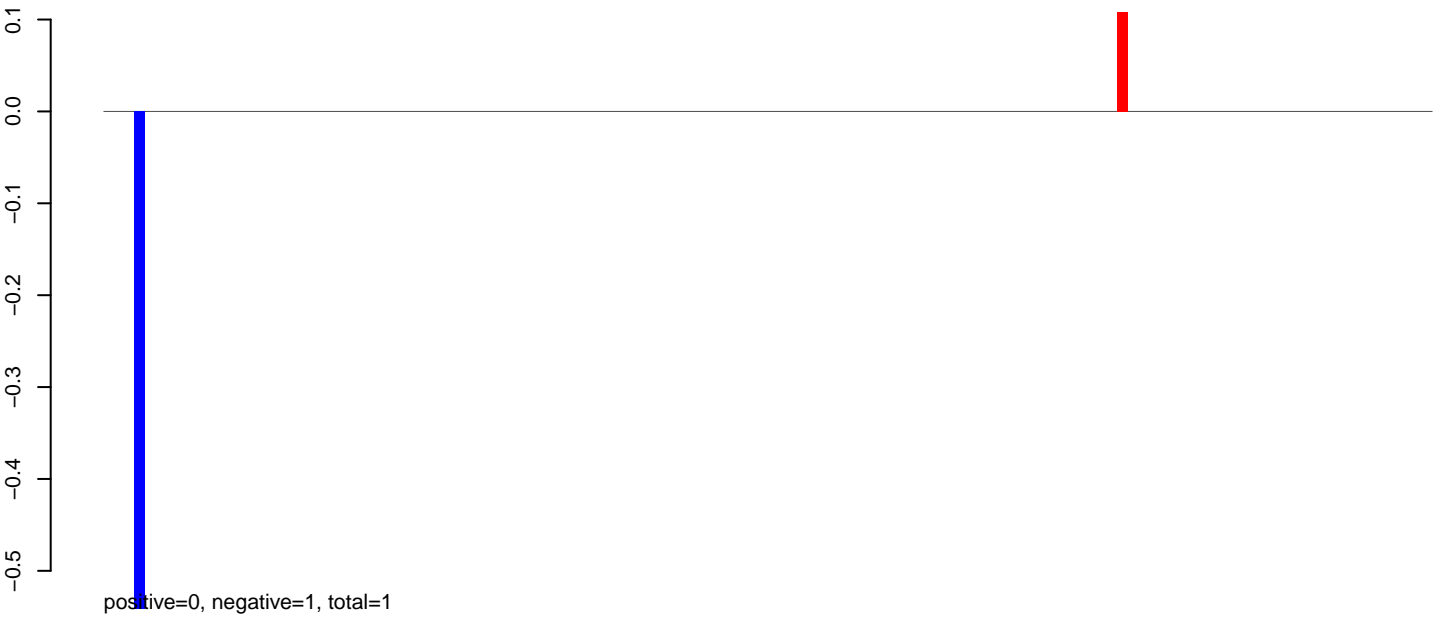
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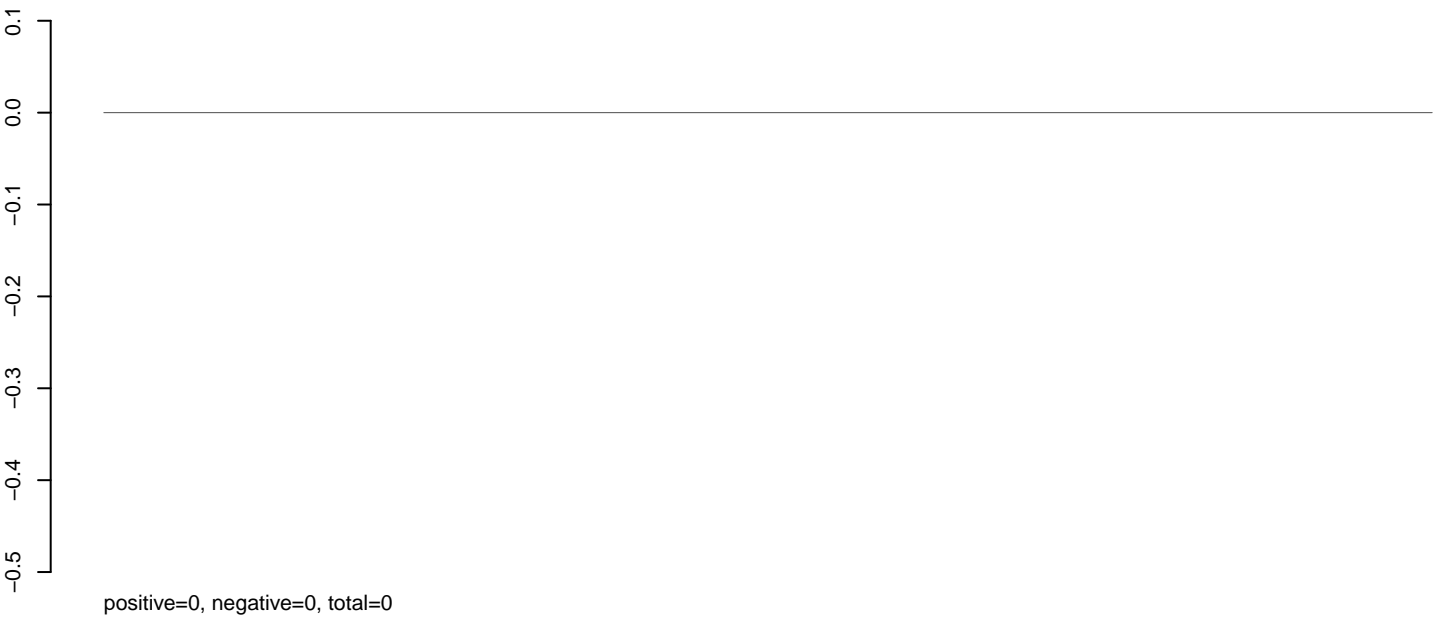
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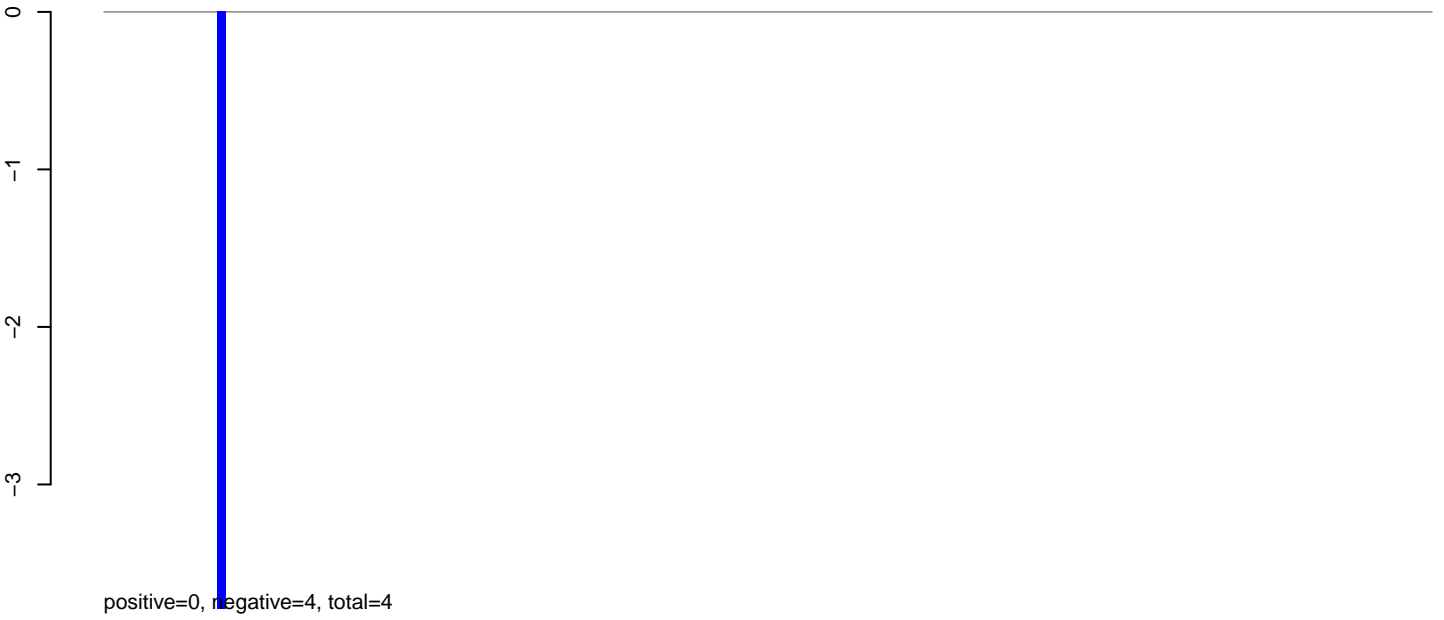
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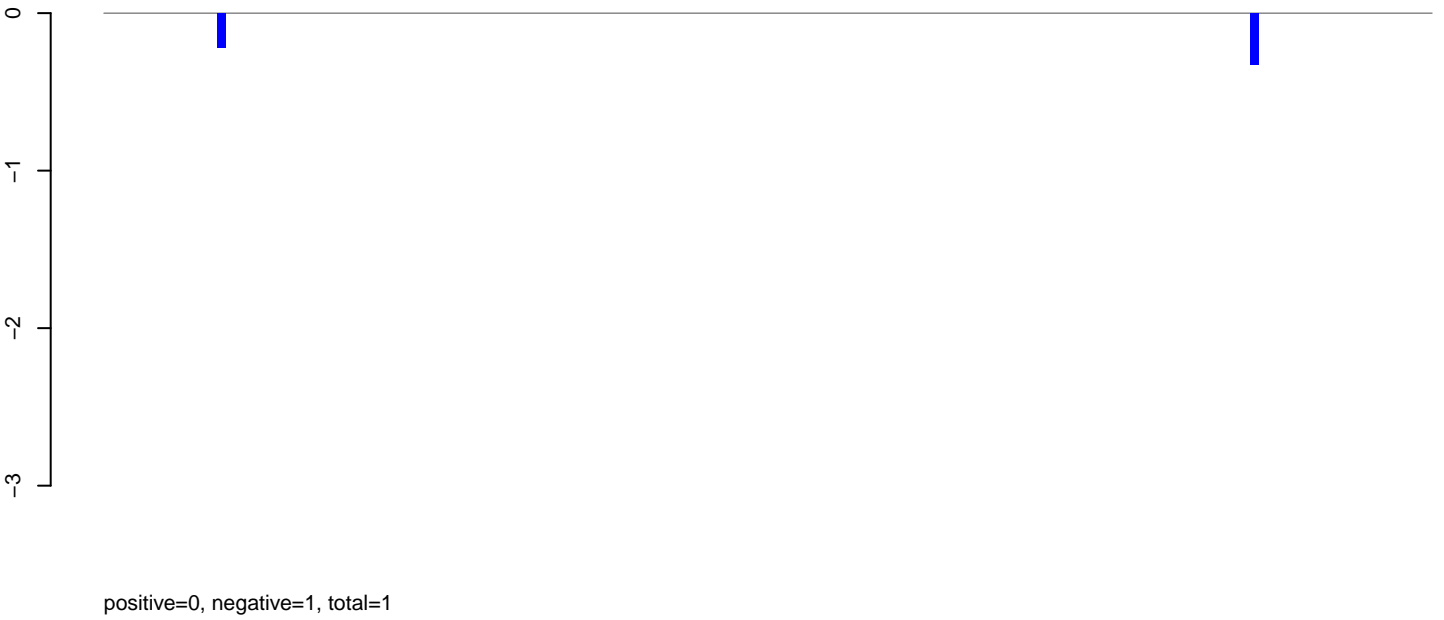
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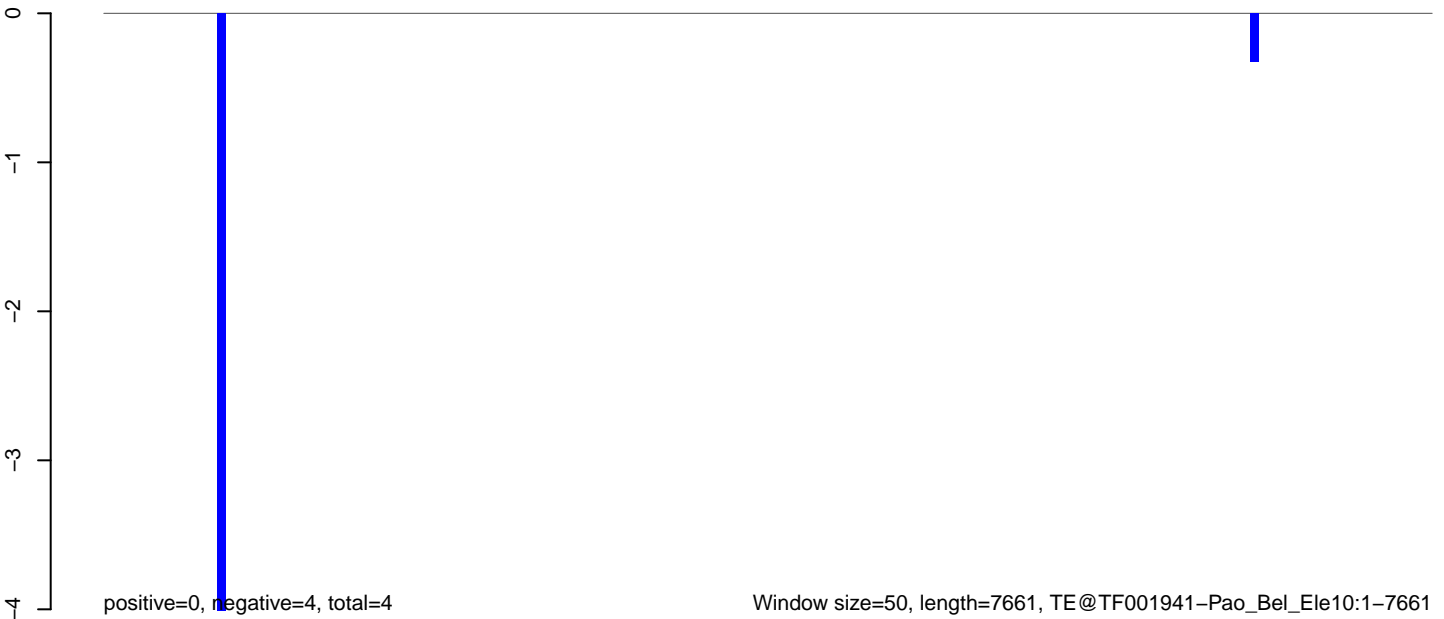
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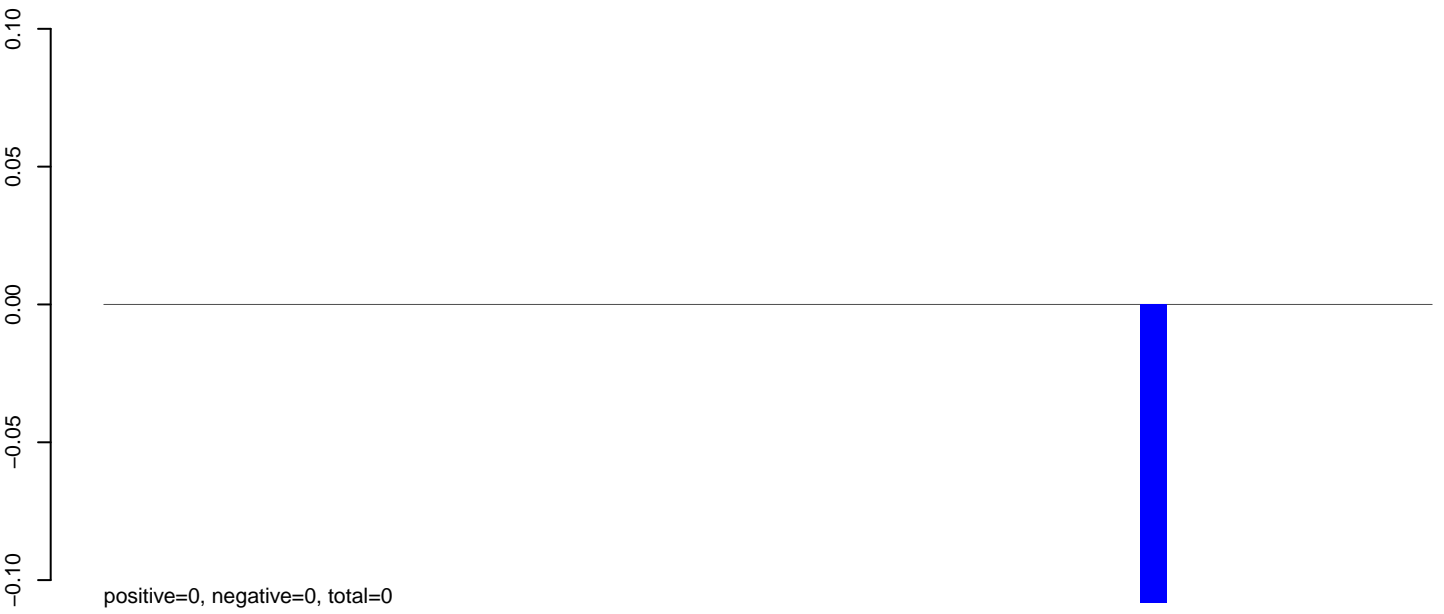
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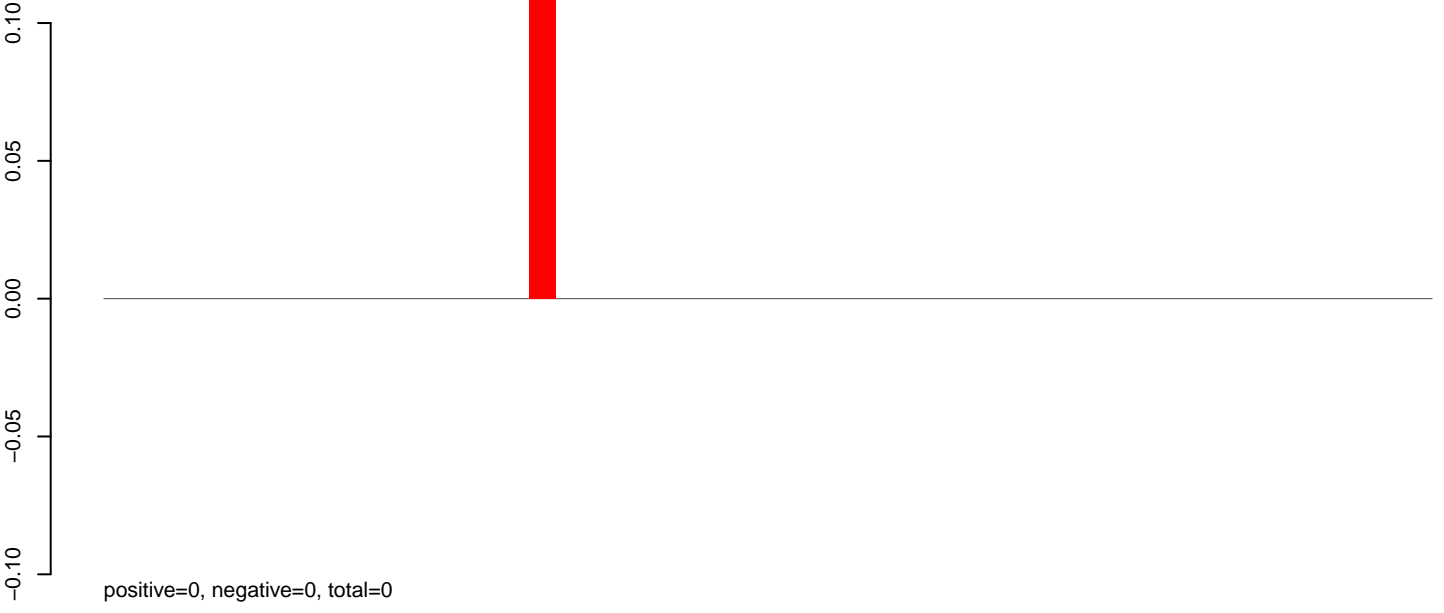
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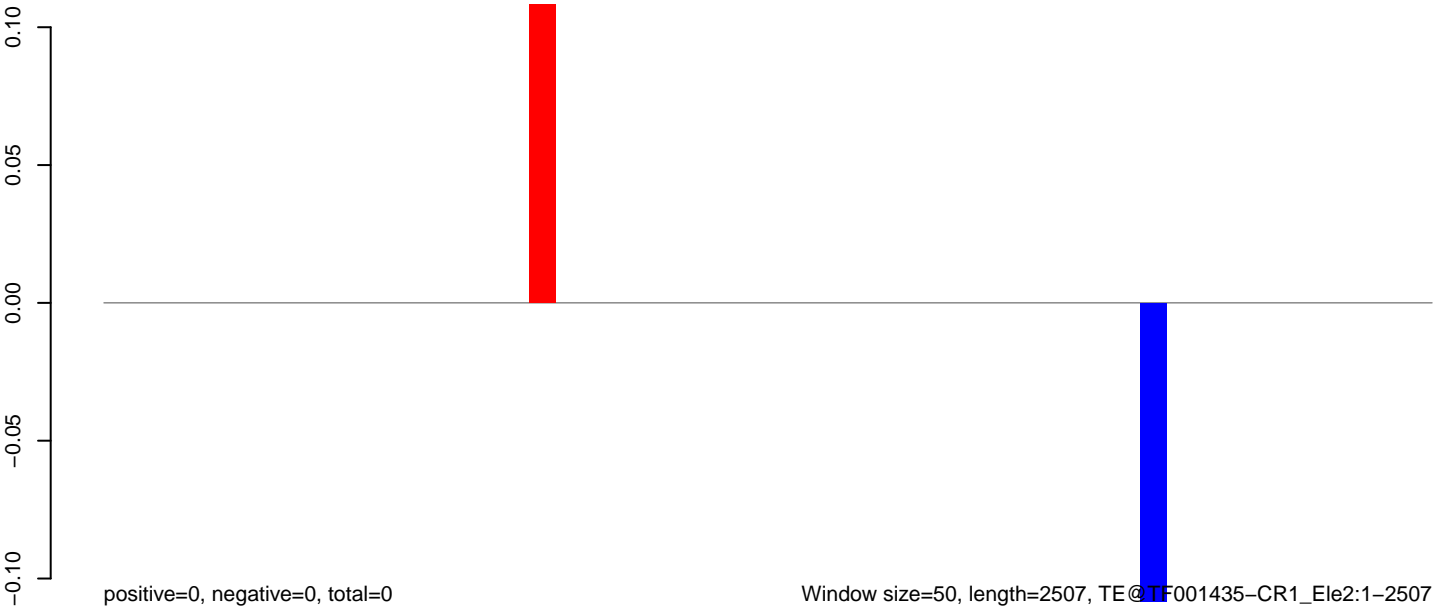
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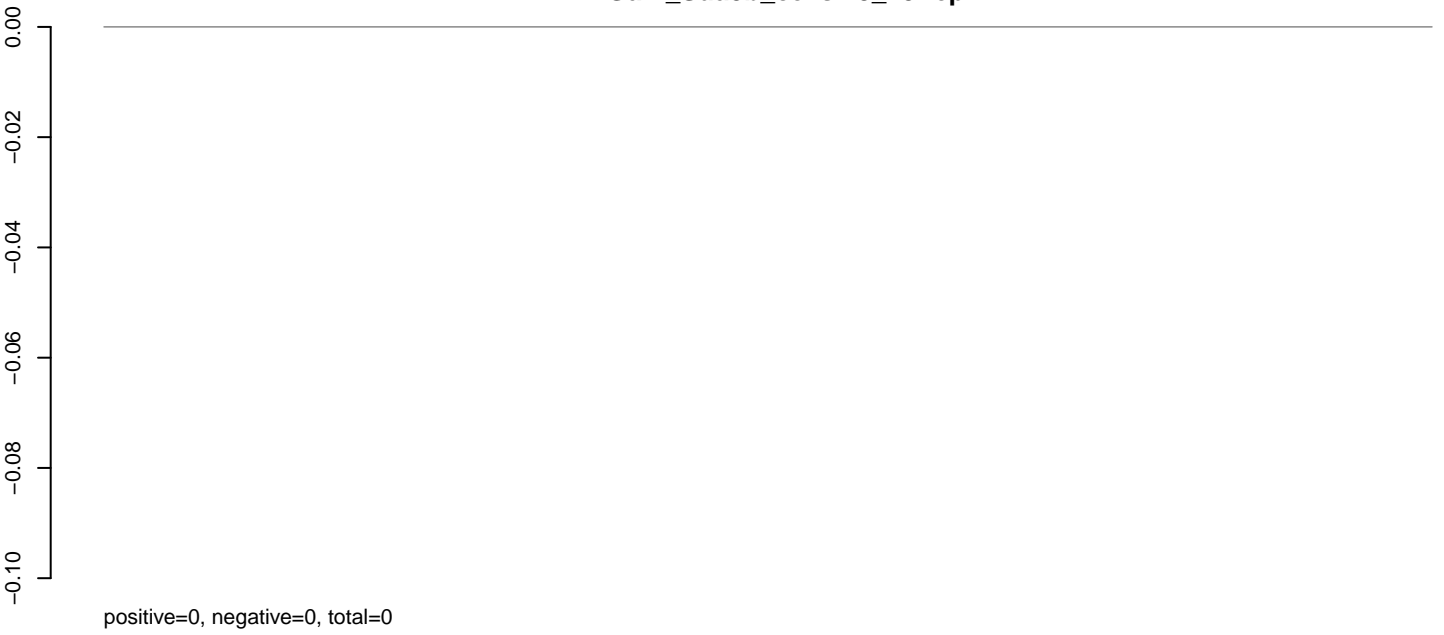
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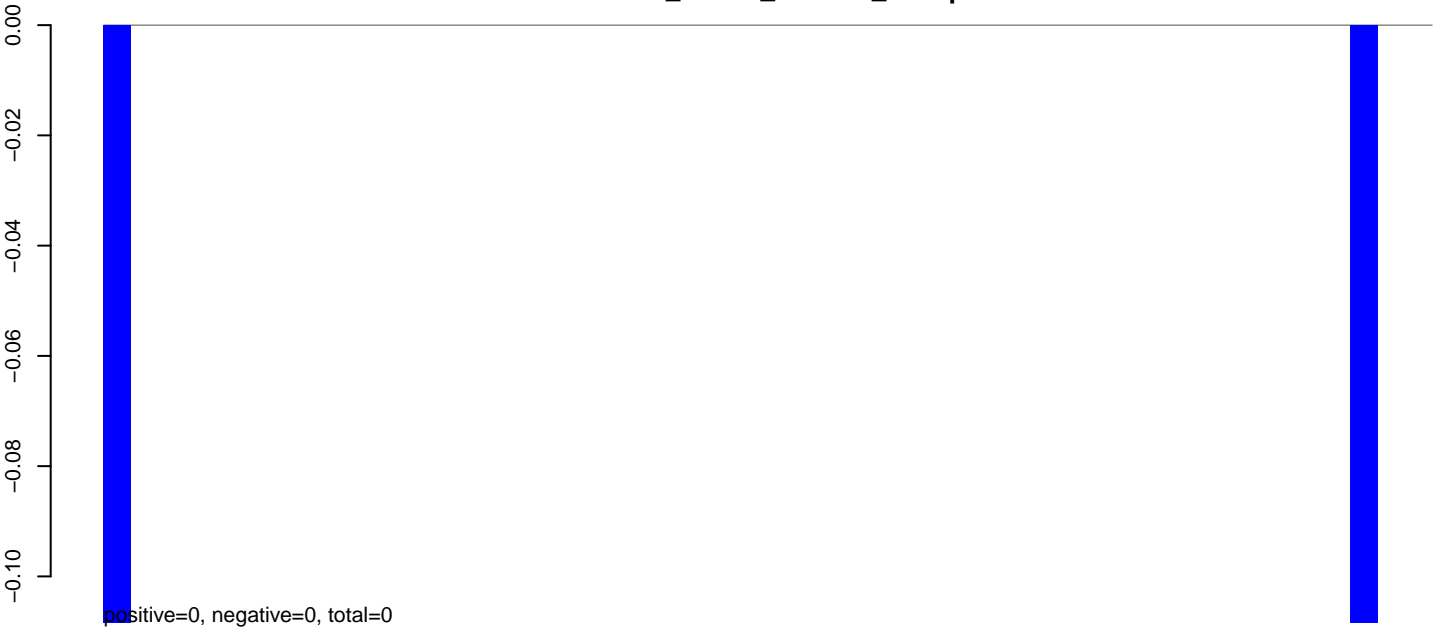
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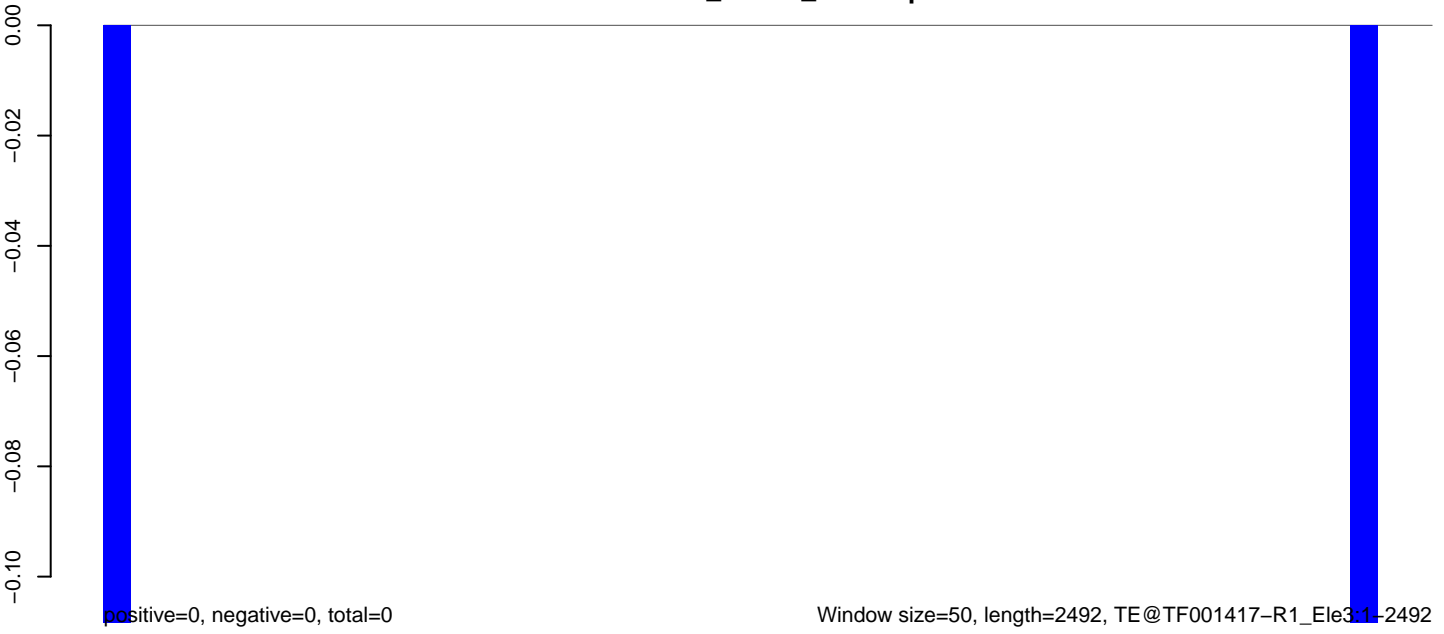
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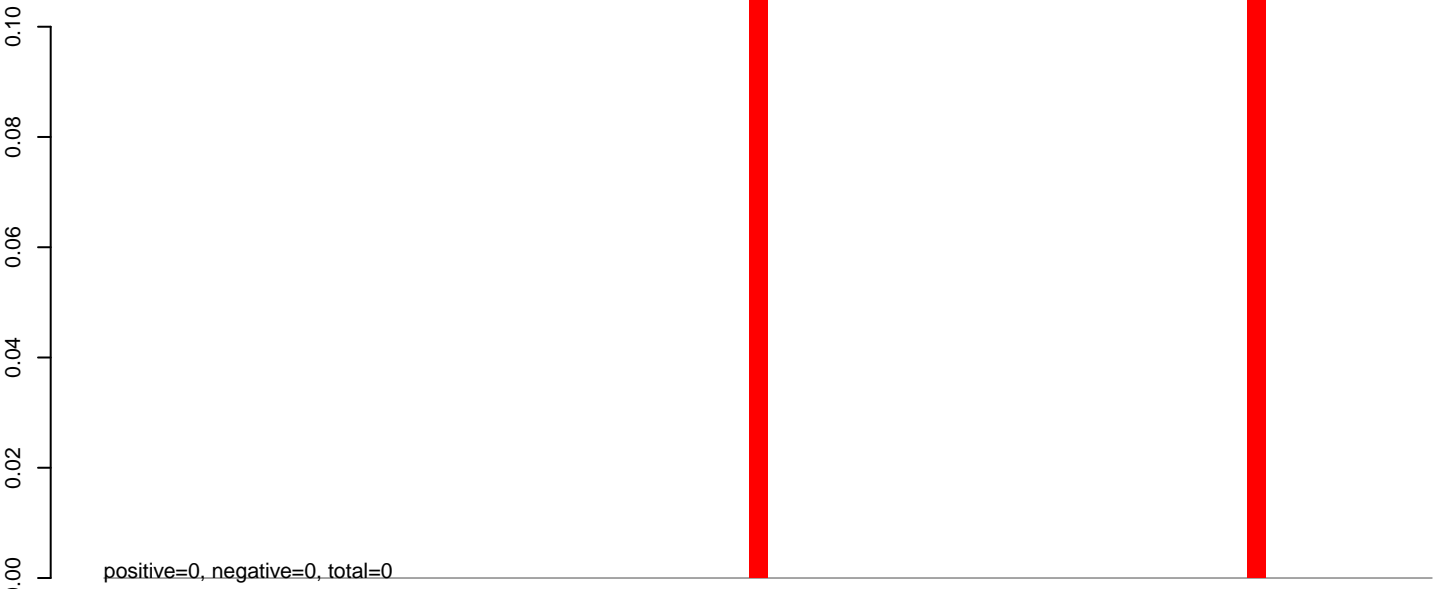
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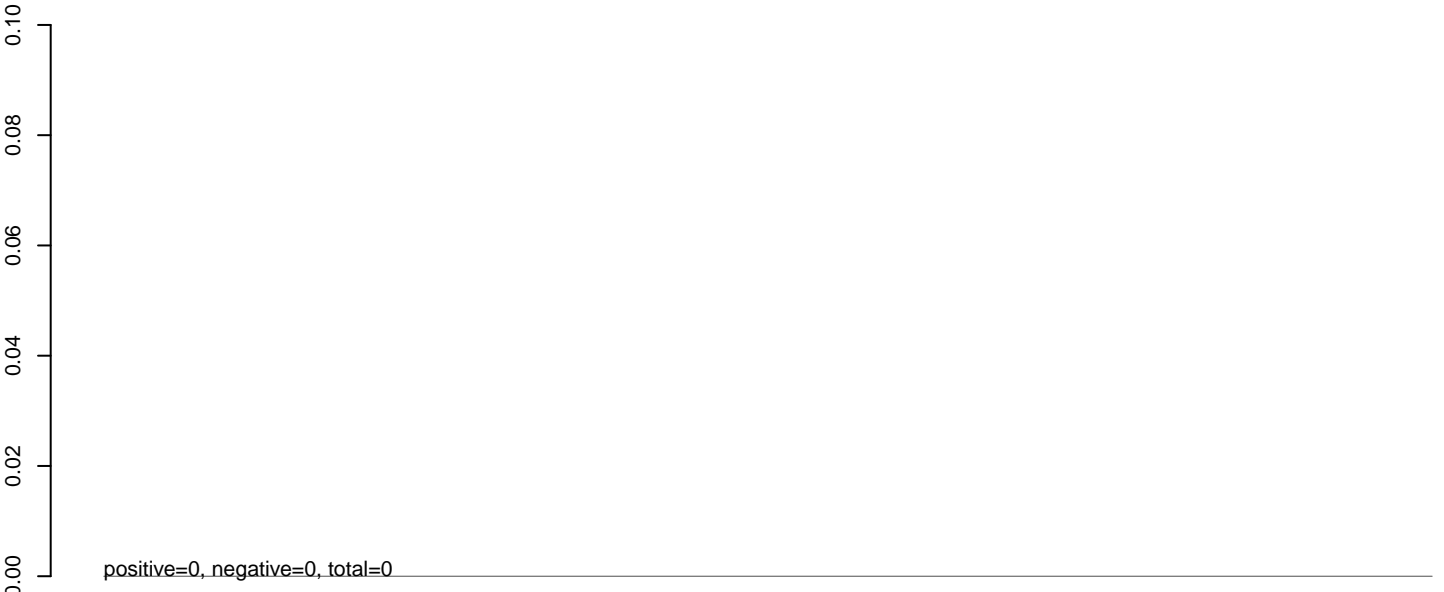
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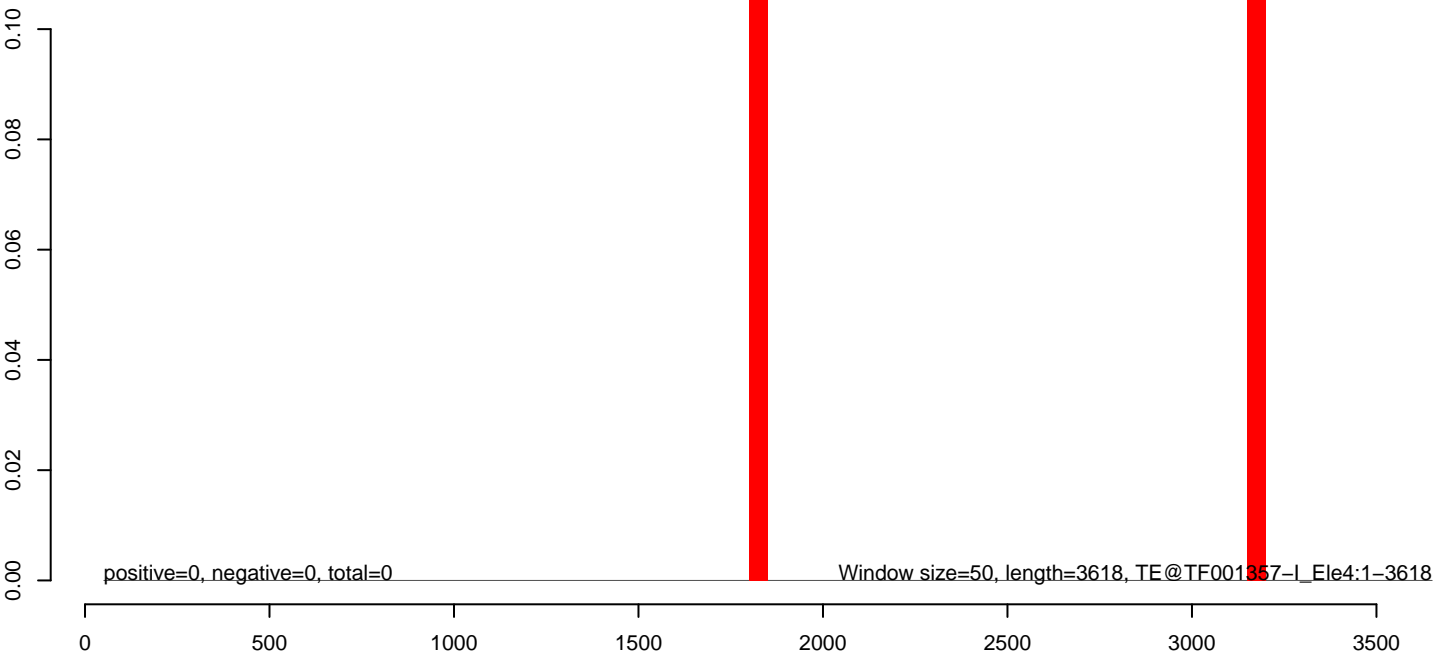
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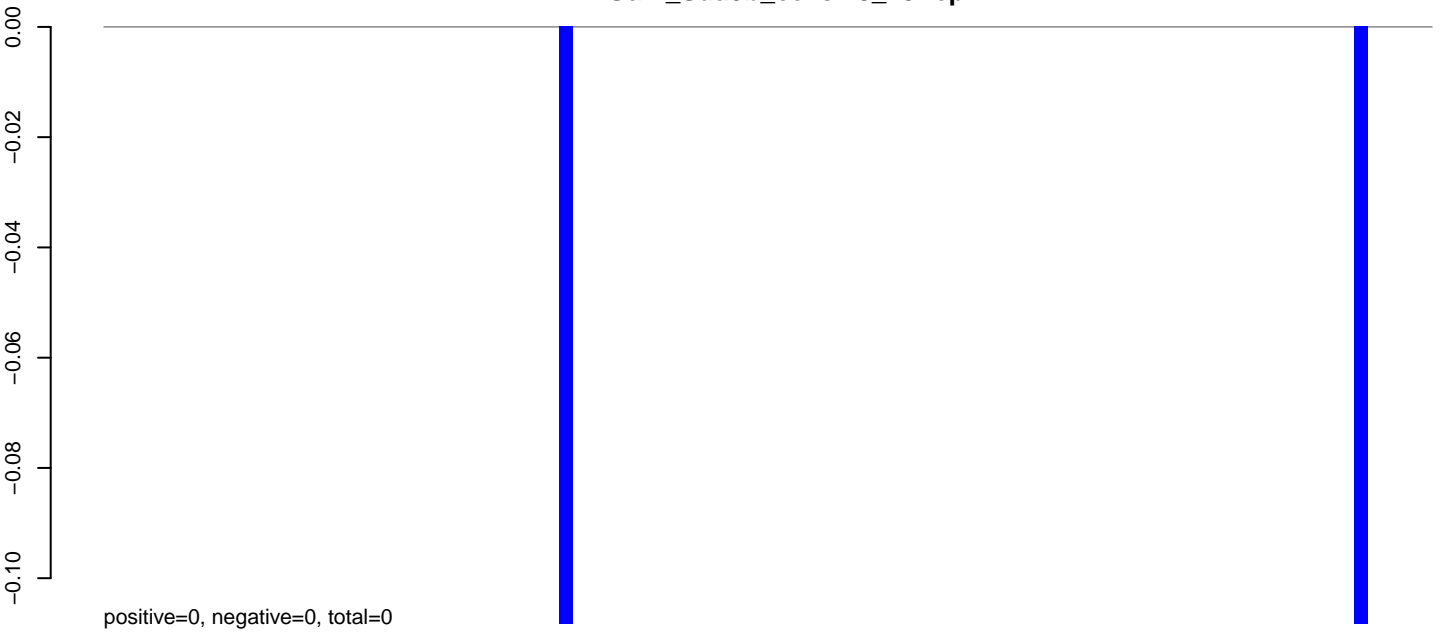
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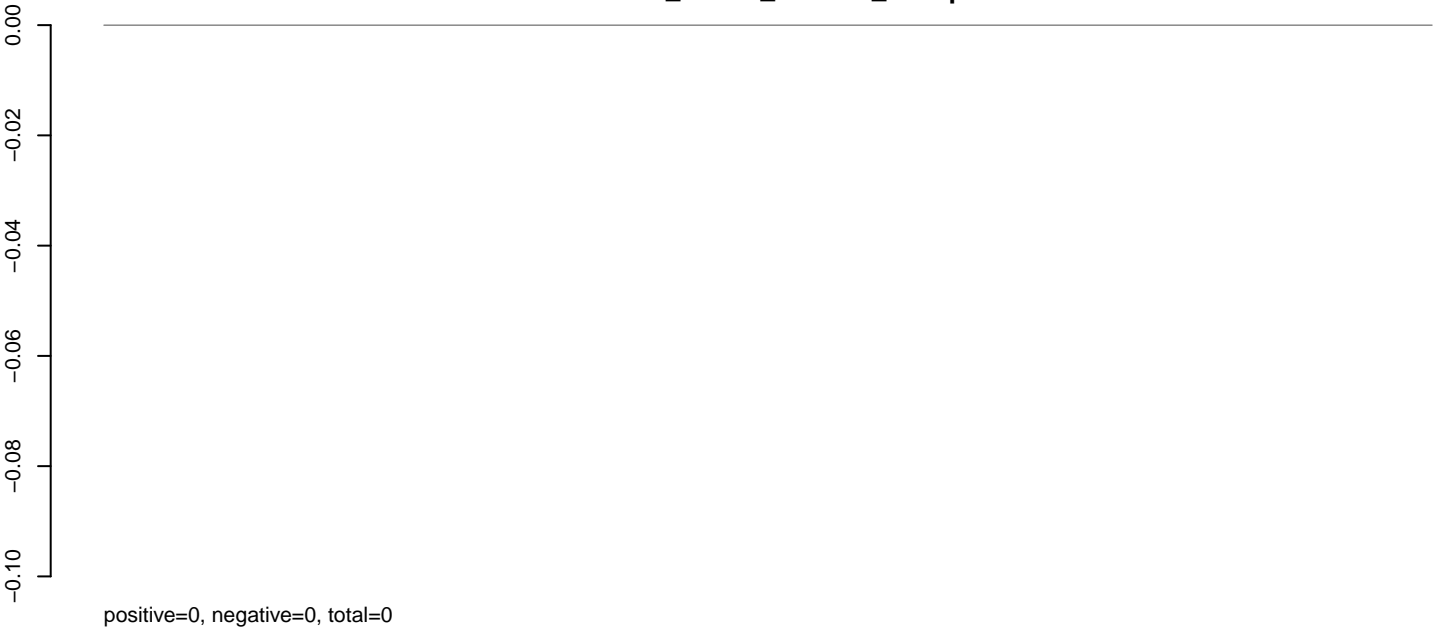
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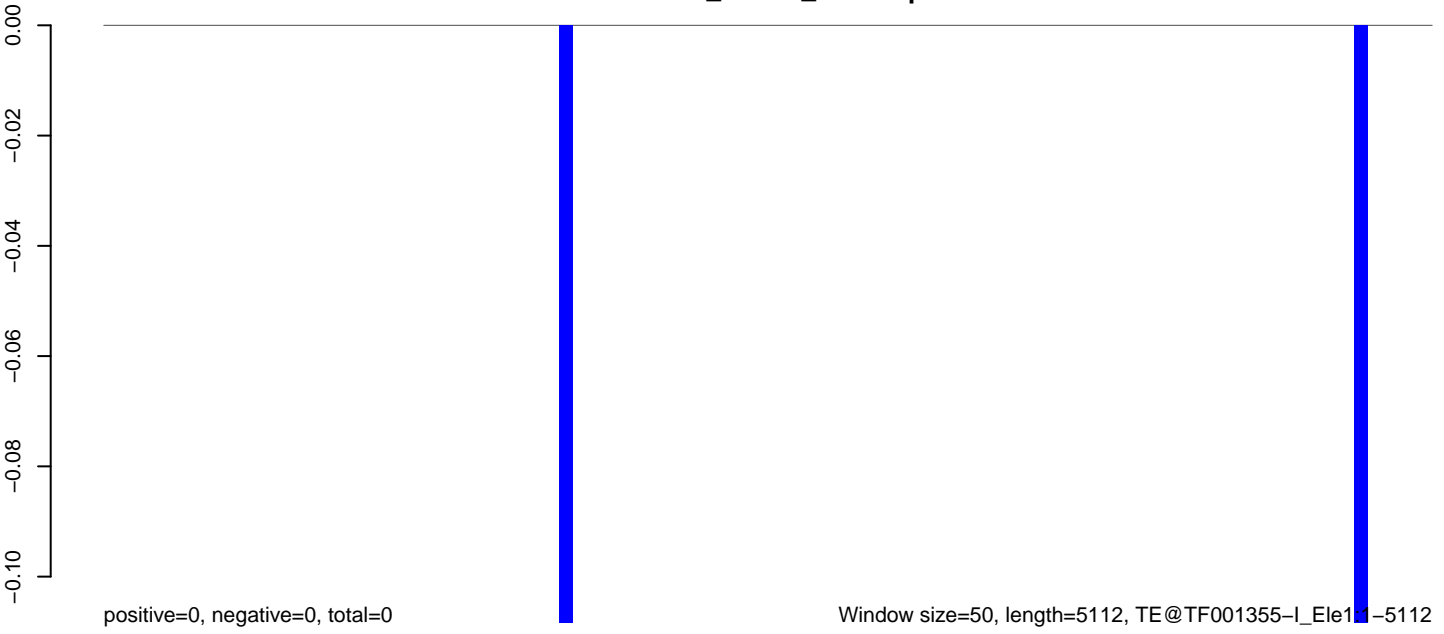
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AnGam_Sua5b_cells.24_35.rep

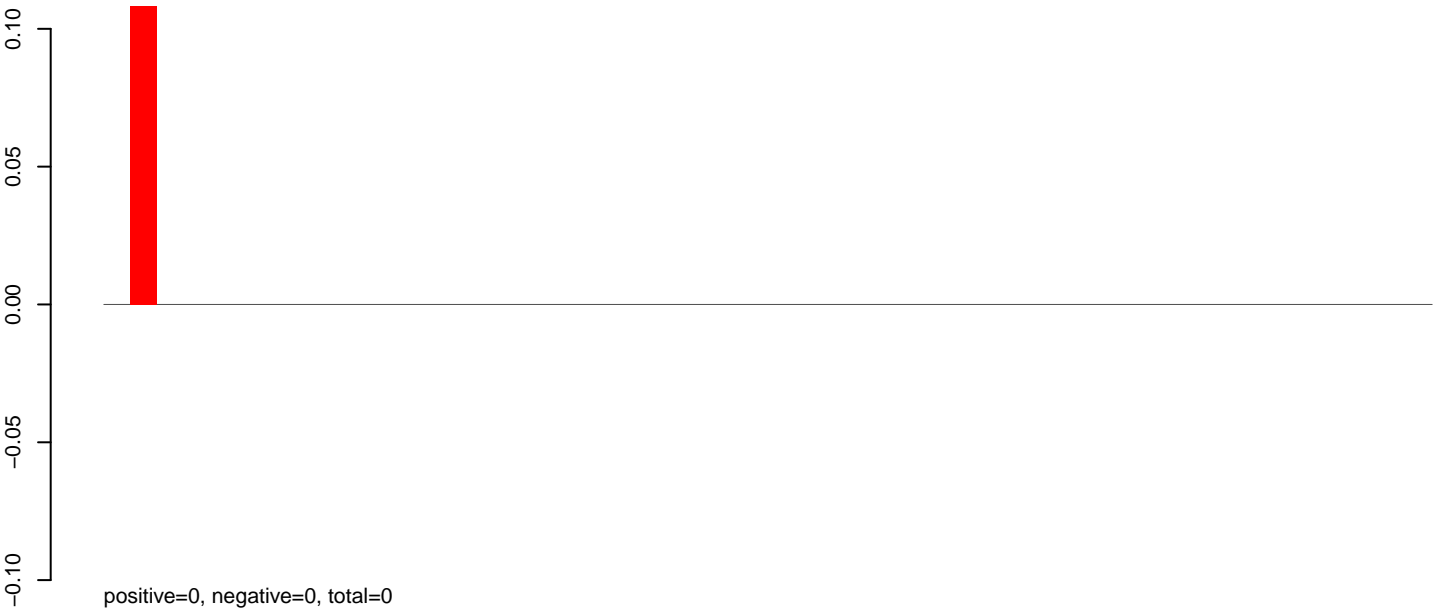


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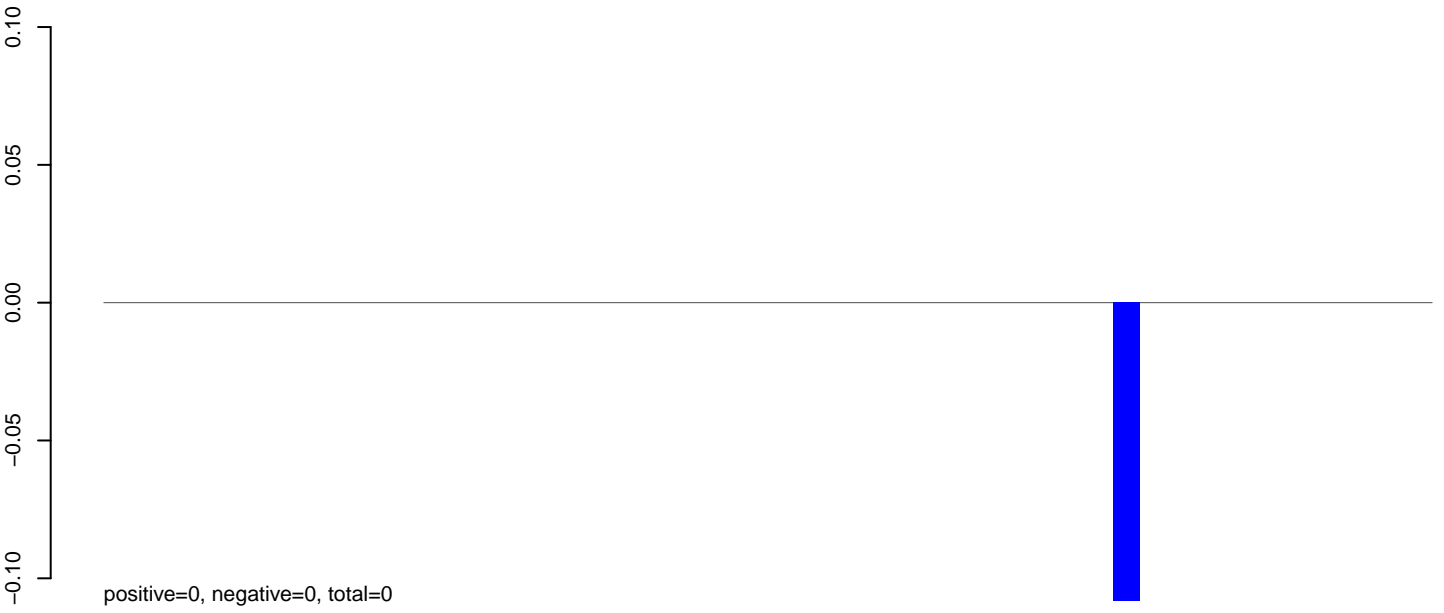


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AnGam_Sua5b_cells.18_23.rep



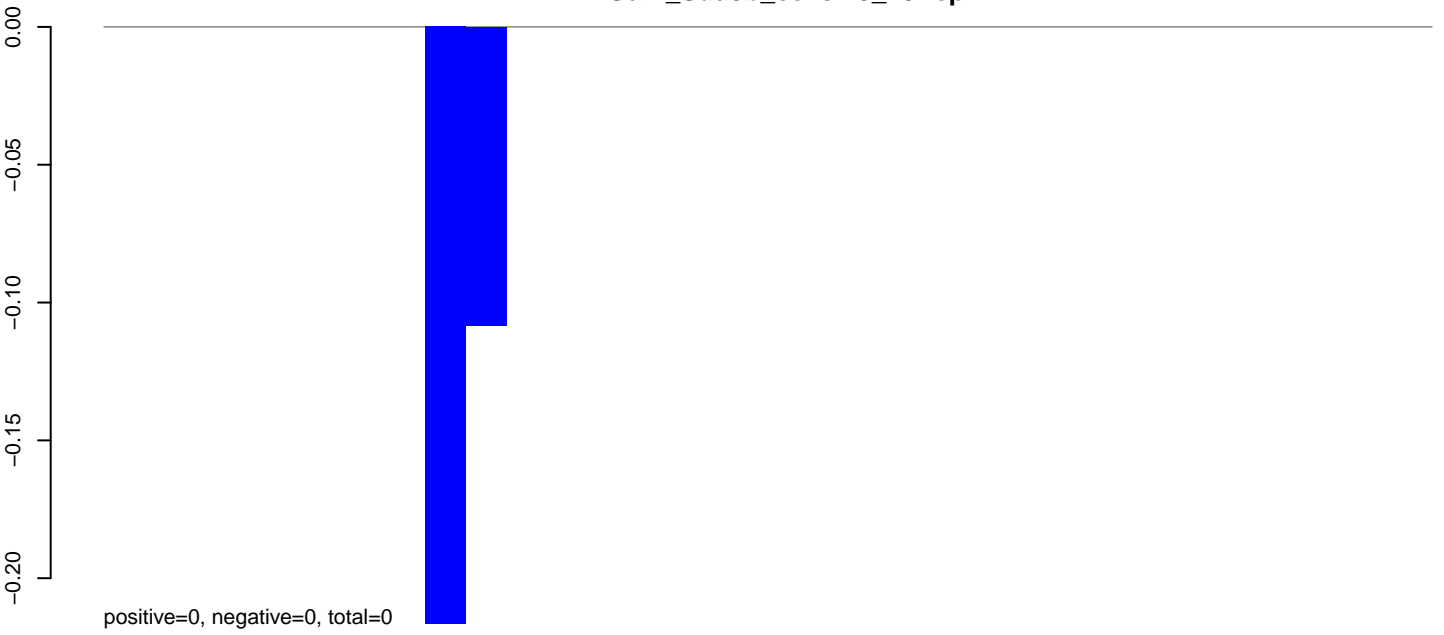
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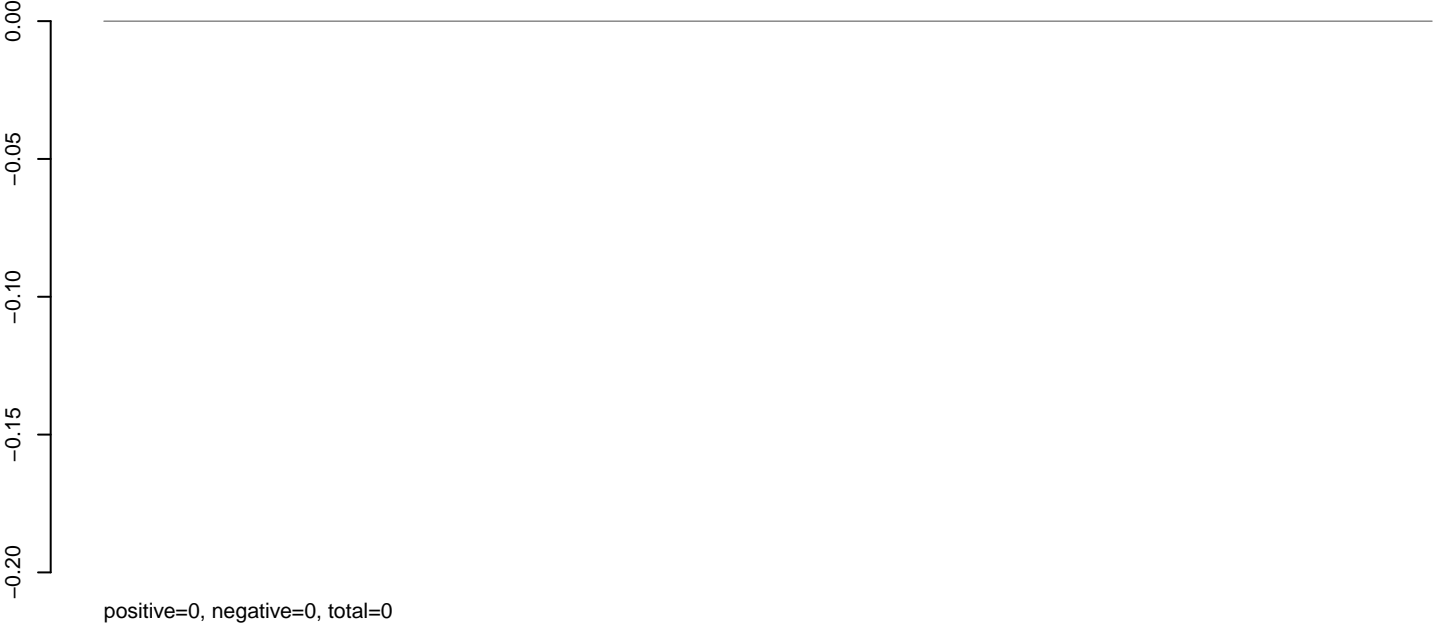
AnGam_Sua5b_cells.rep



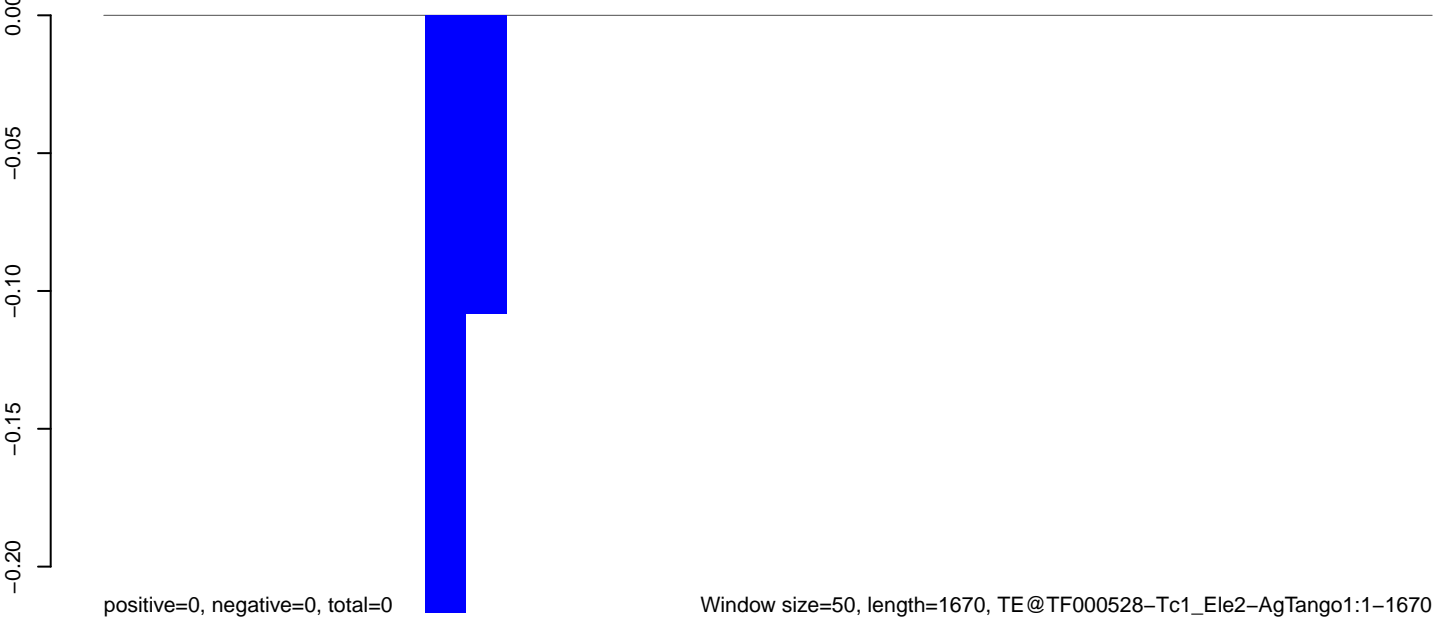
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep

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-0.04
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positive=0, negative=0, total=0

AnGam_Sua5b_cells.rep

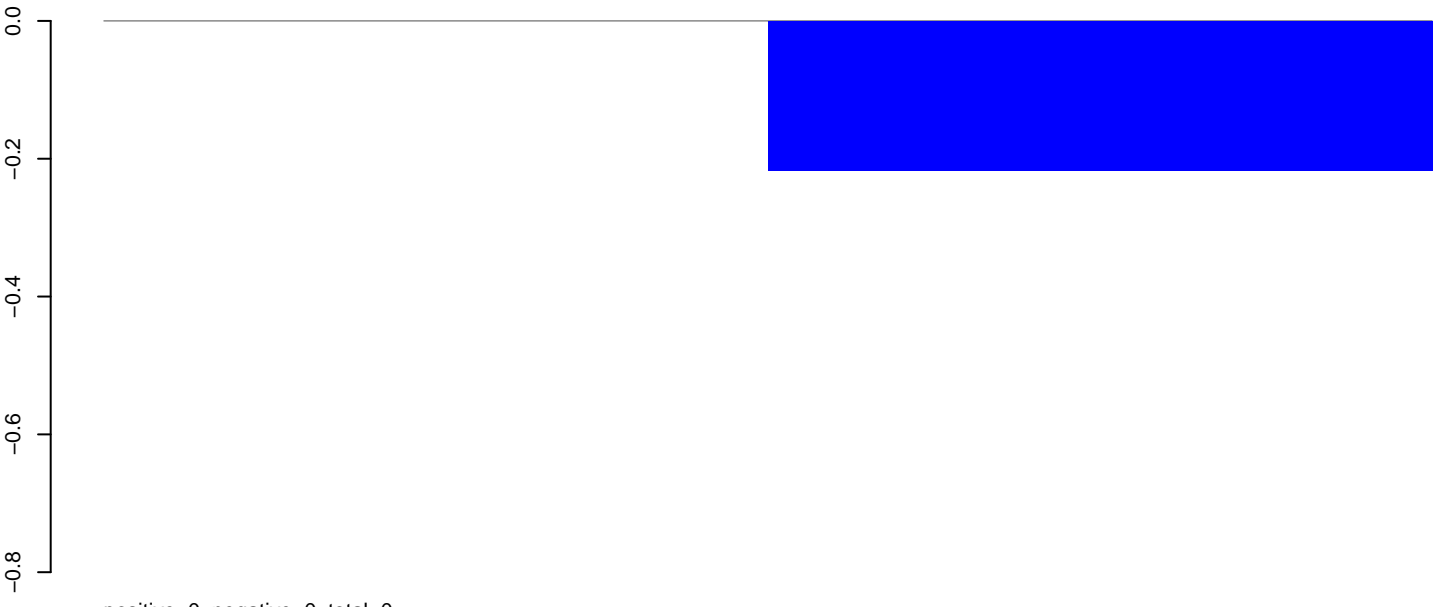
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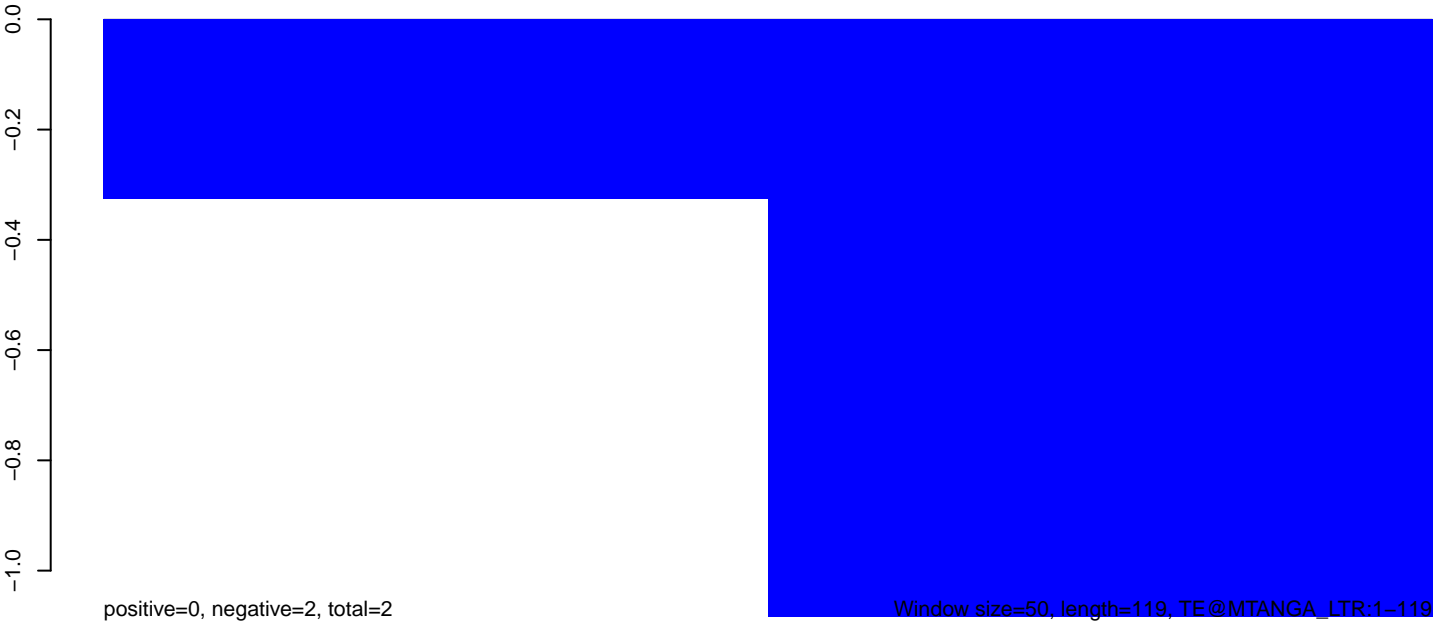
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AnGam_Sua5b_cells.24_35.rep



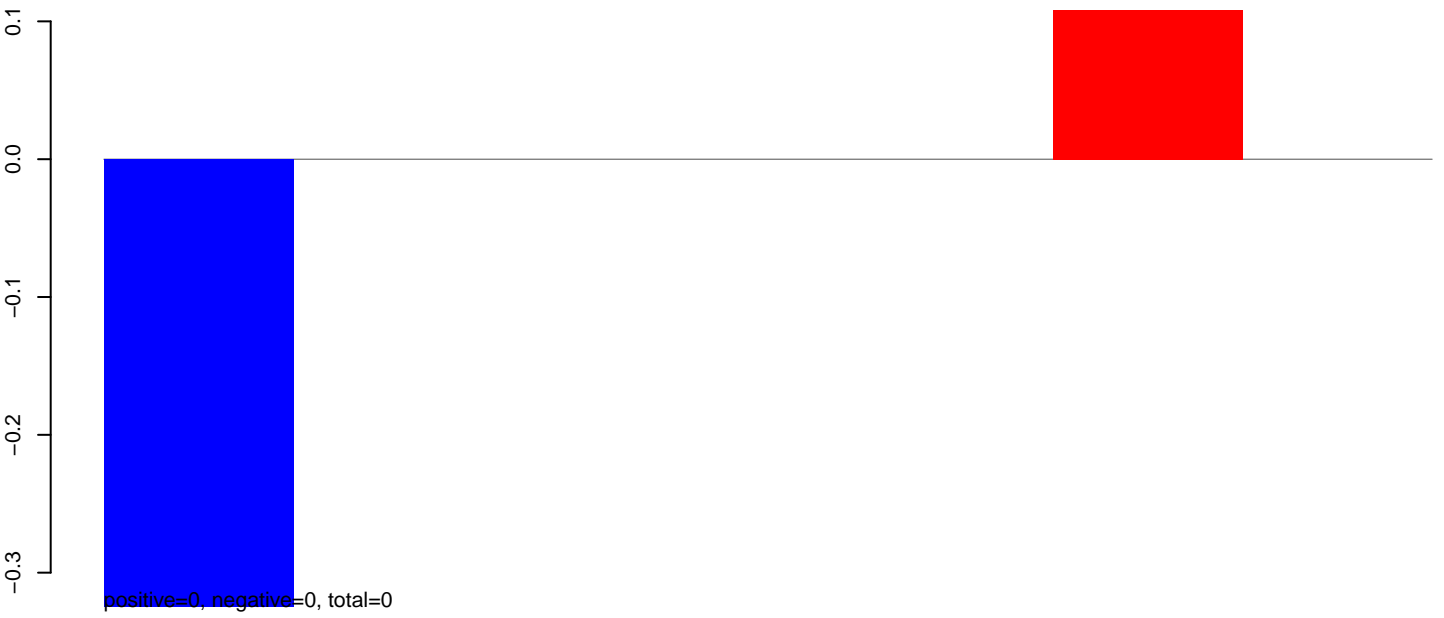
AnGam_Sua5b_cells.rep



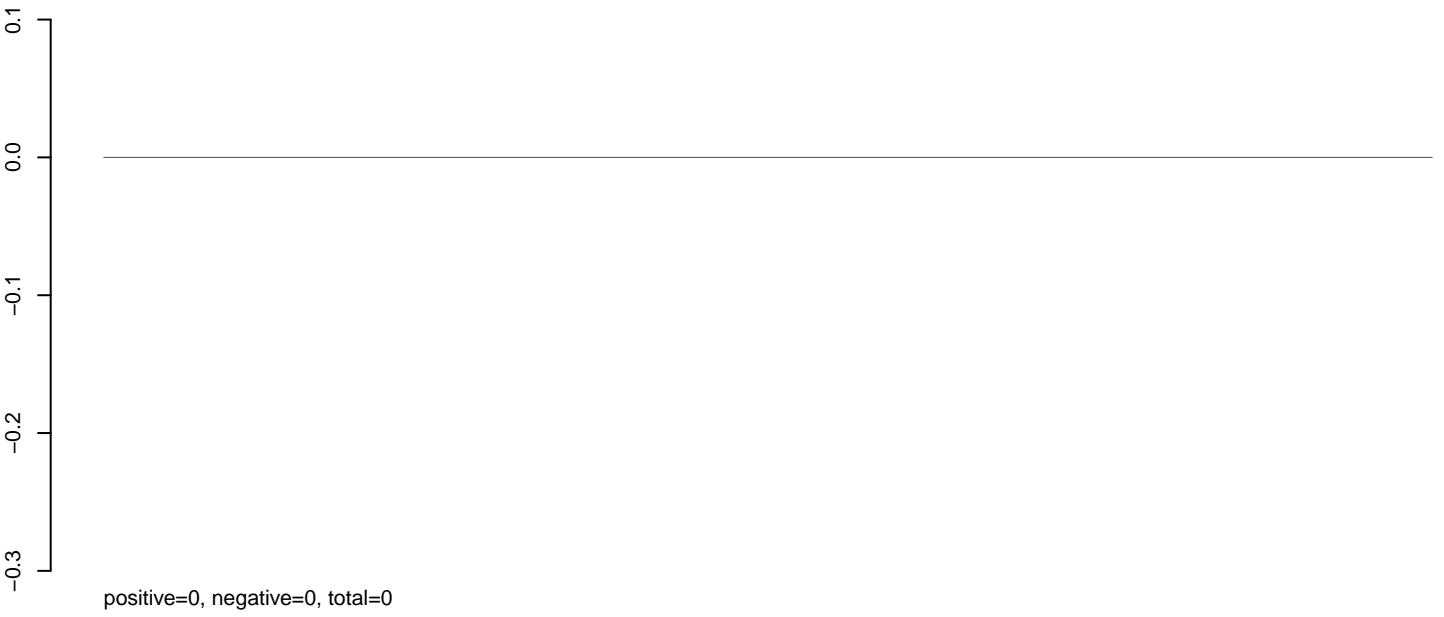
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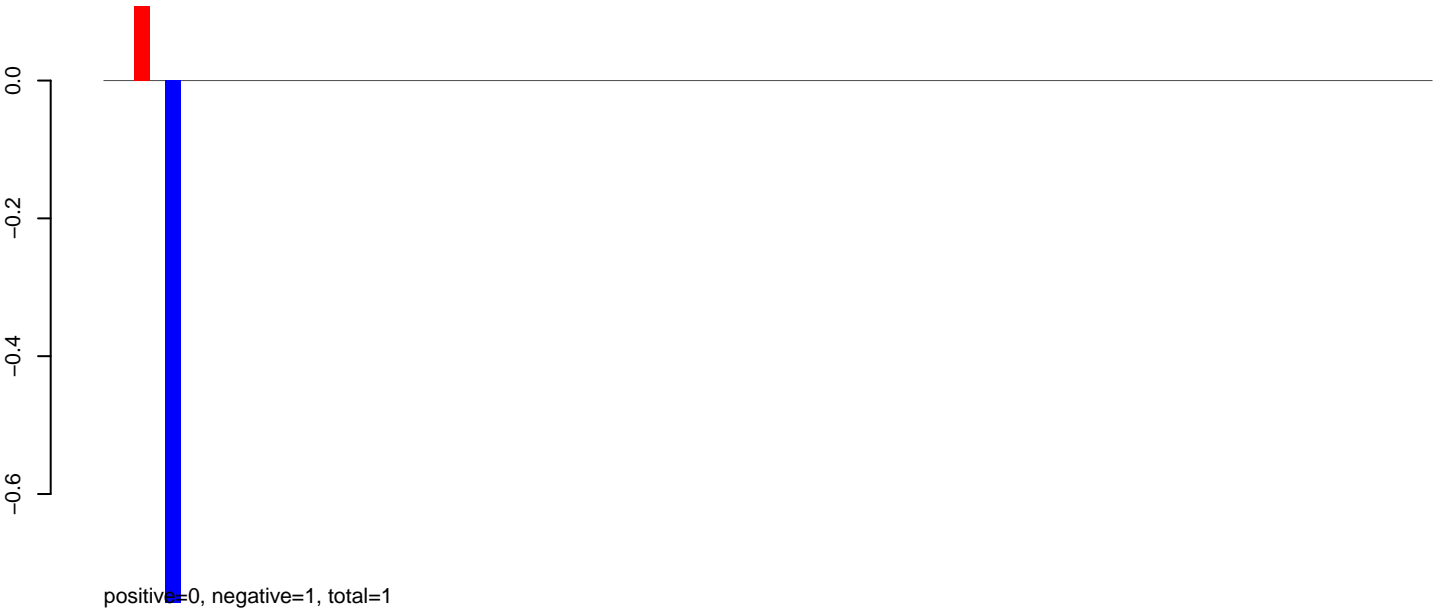
AnGam_Sua5b_cells.24_35.rep



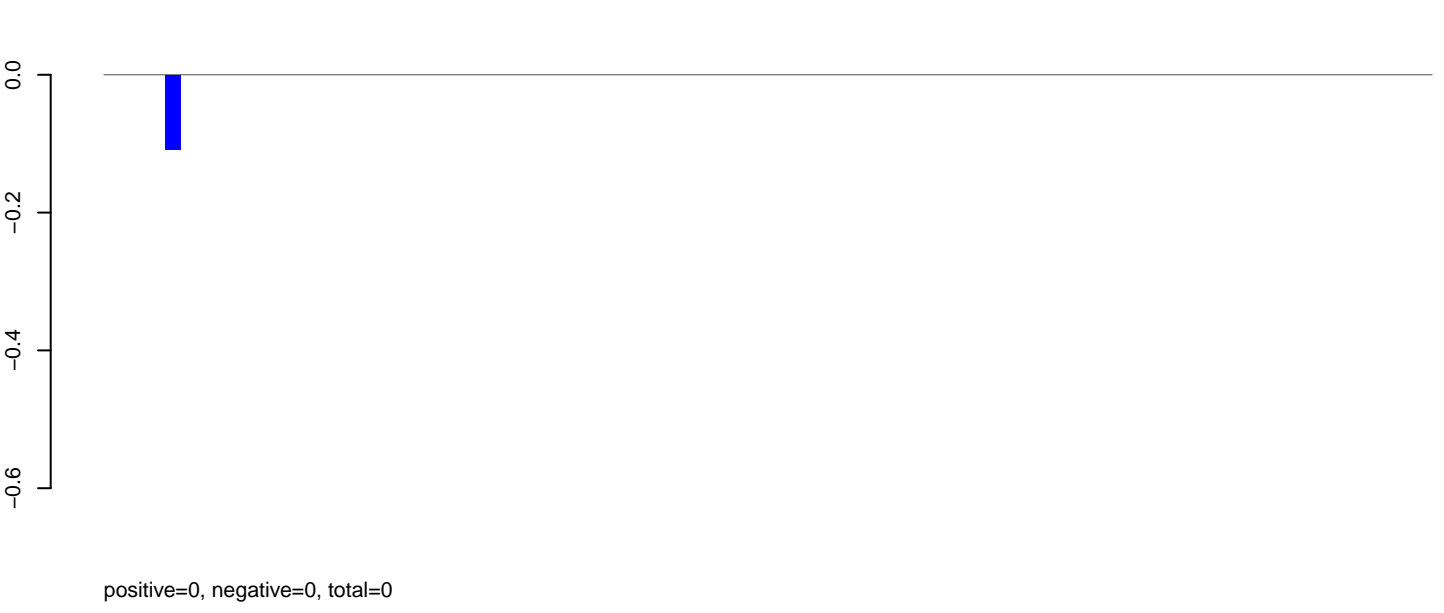
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AnGam_Sua5b_cells.18_23.rep



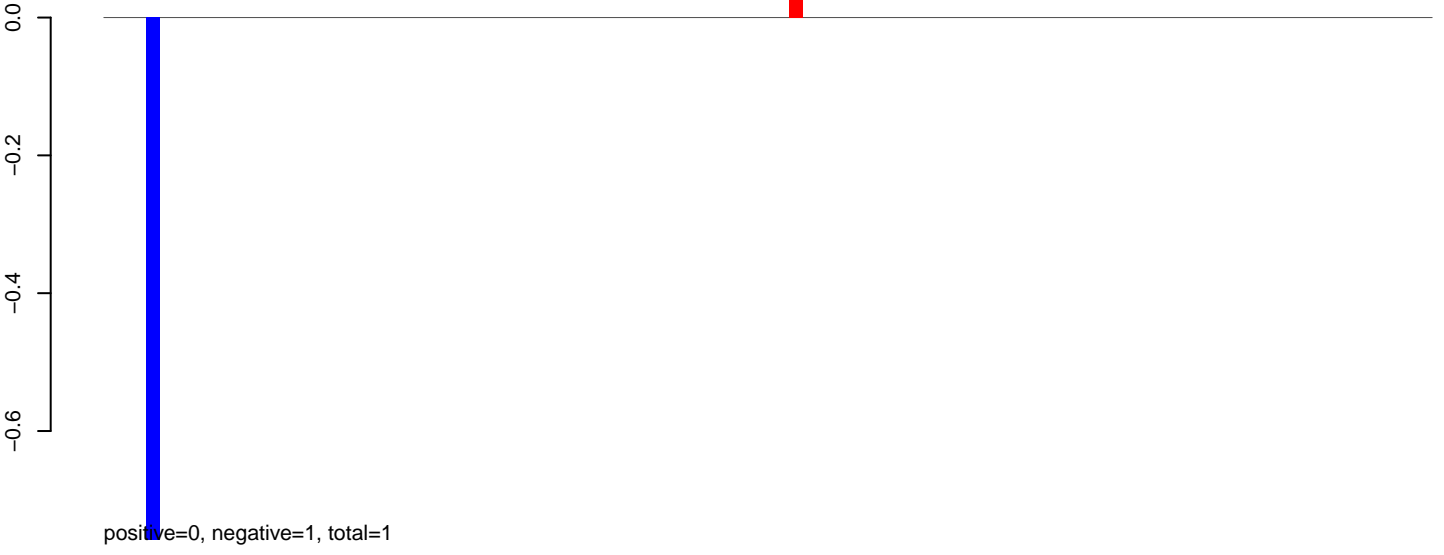
AnGam_Sua5b_cells.24_35.rep



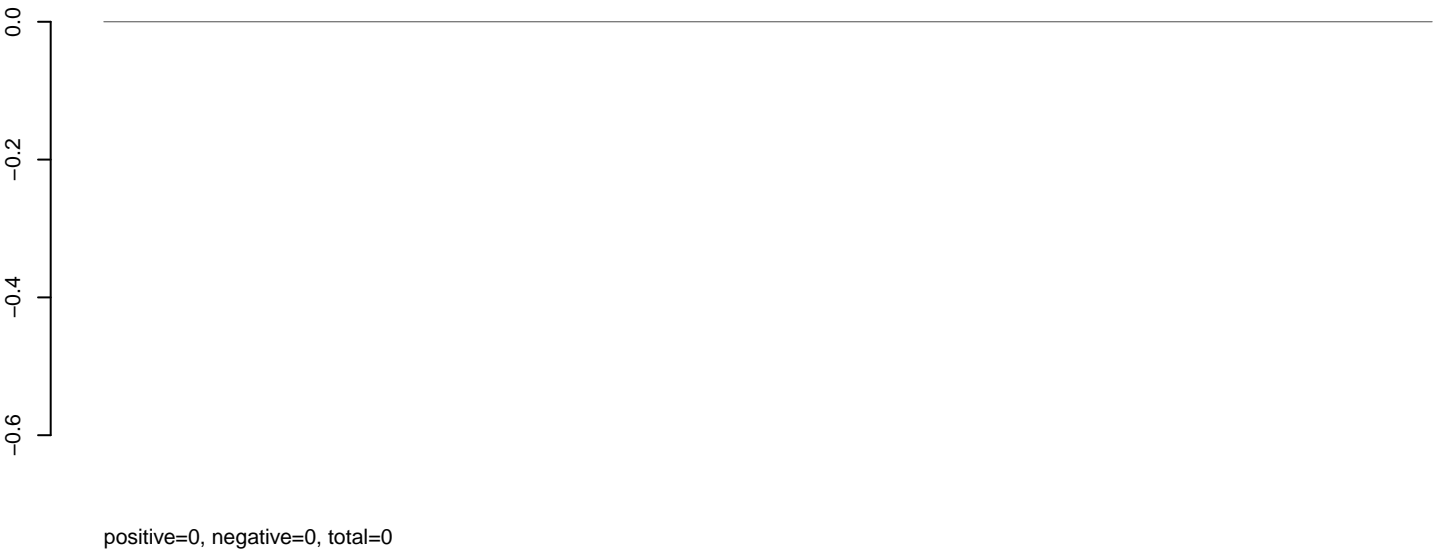
AnGam_Sua5b_cells.rep



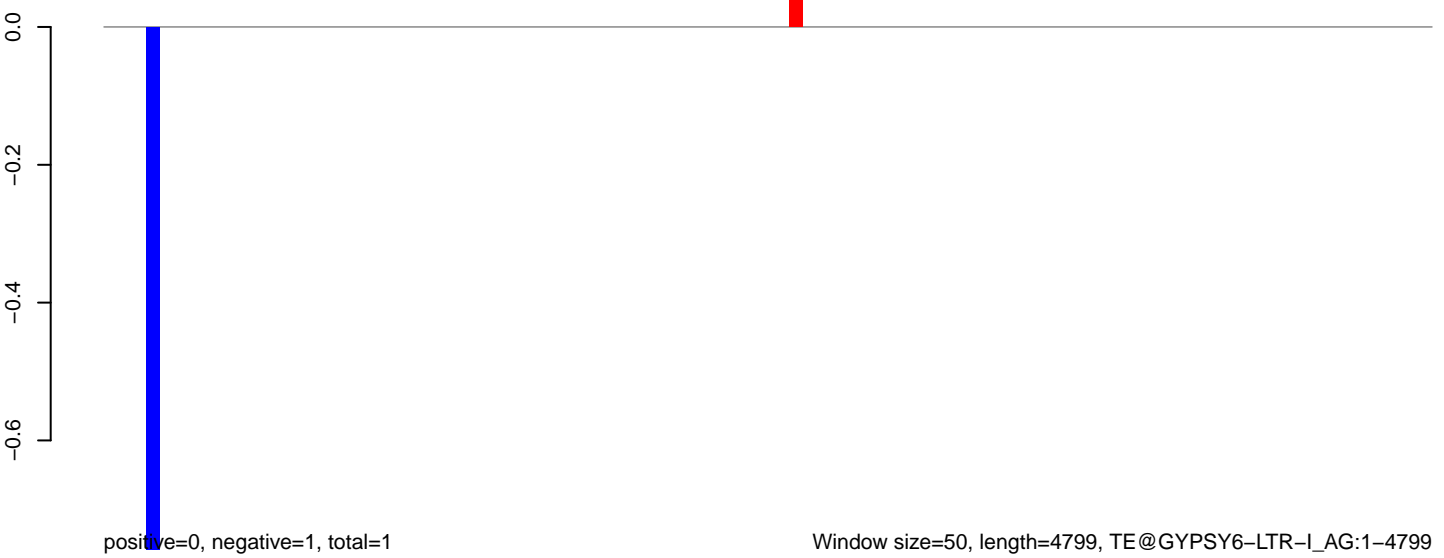
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AnGam_Sua5b_cells.24_35.rep



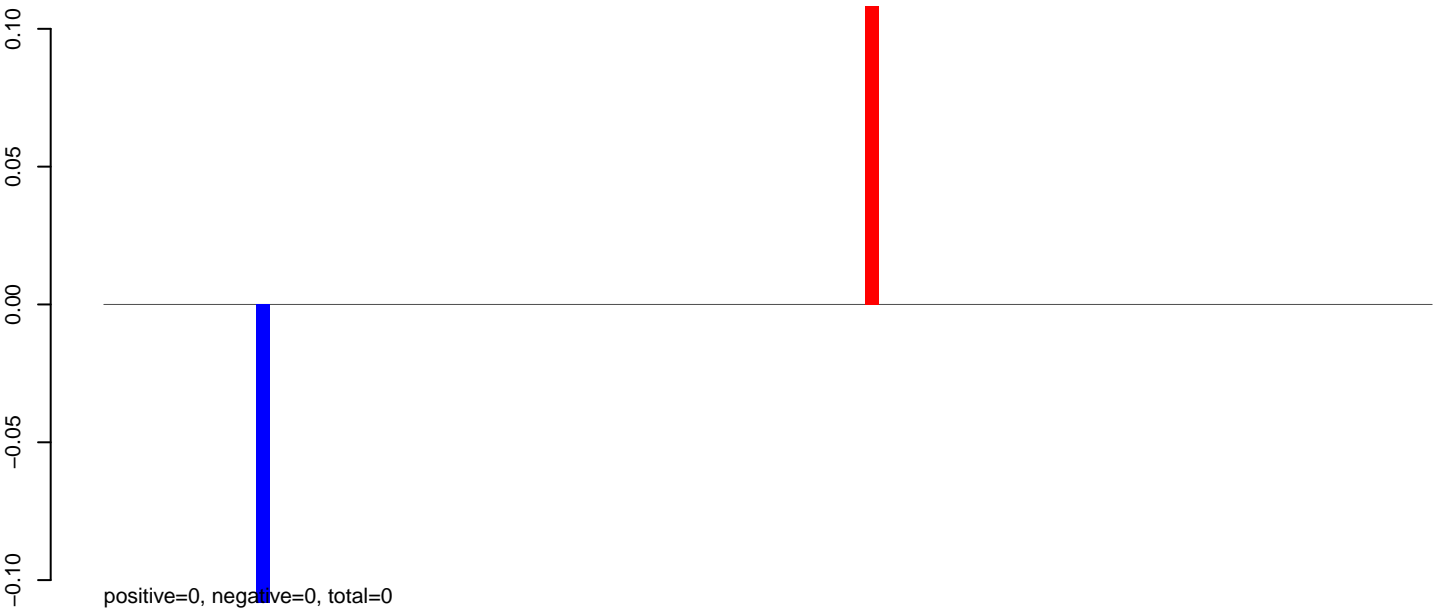
AnGam_Sua5b_cells.rep



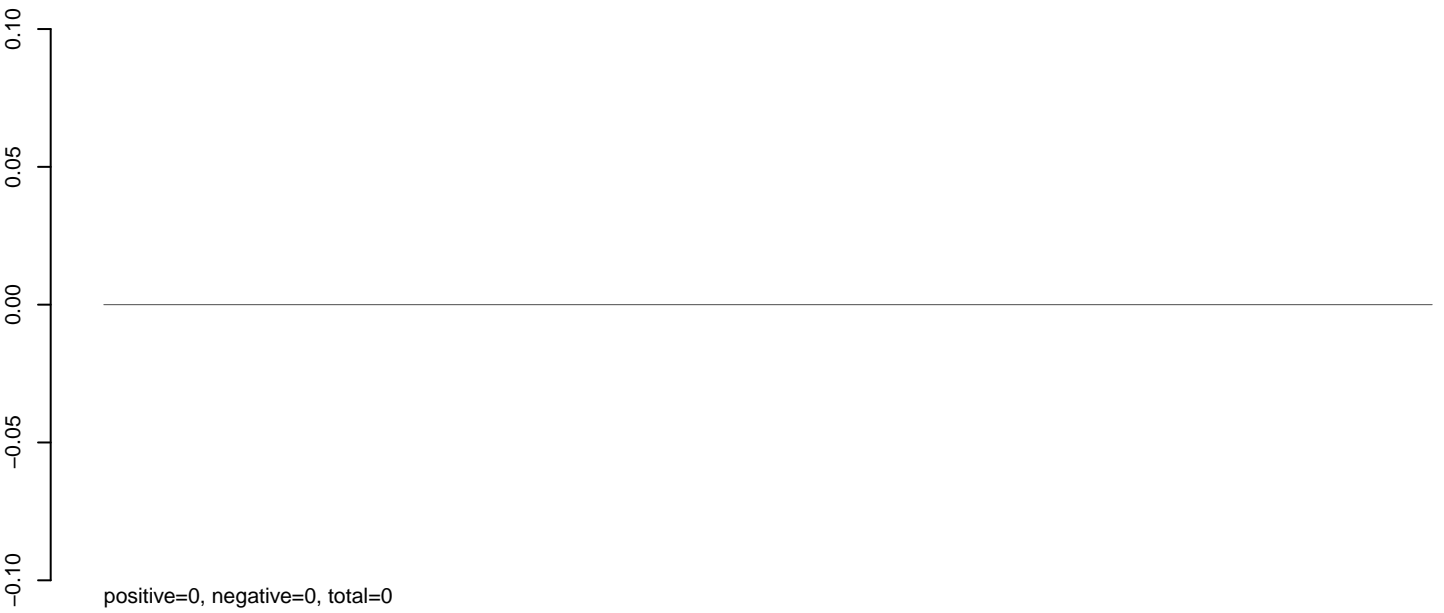
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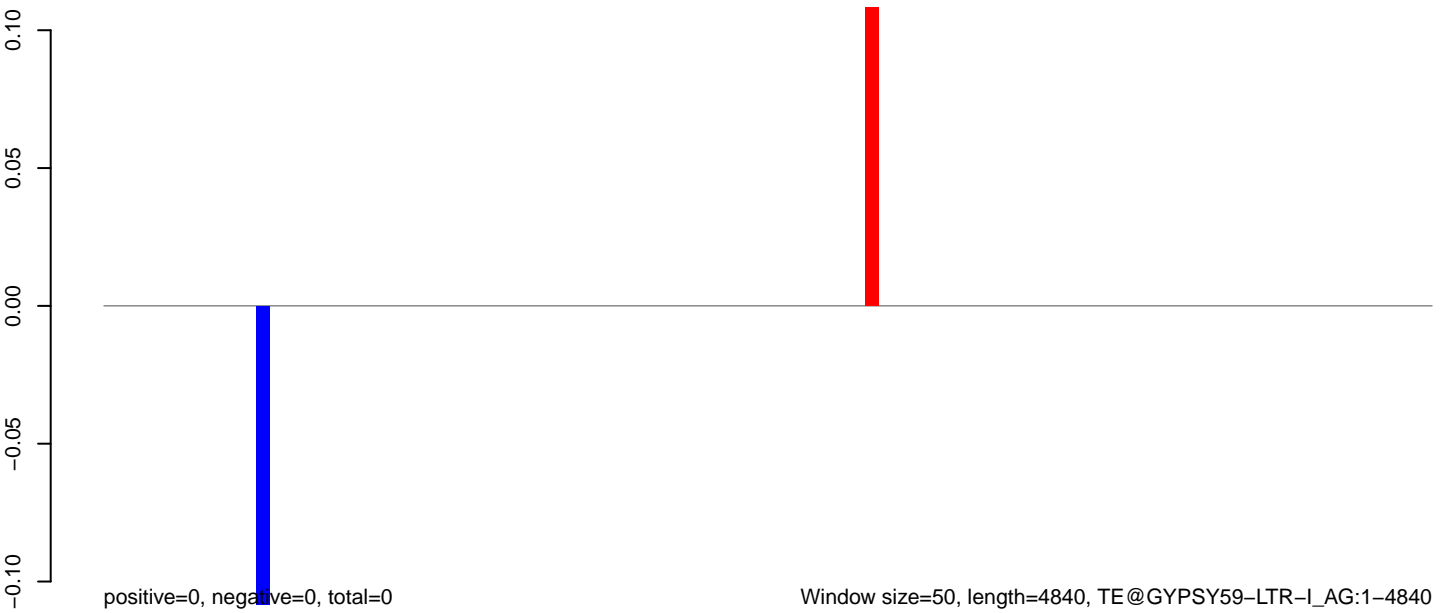
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AnGam_Sua5b_cells.24_35.rep

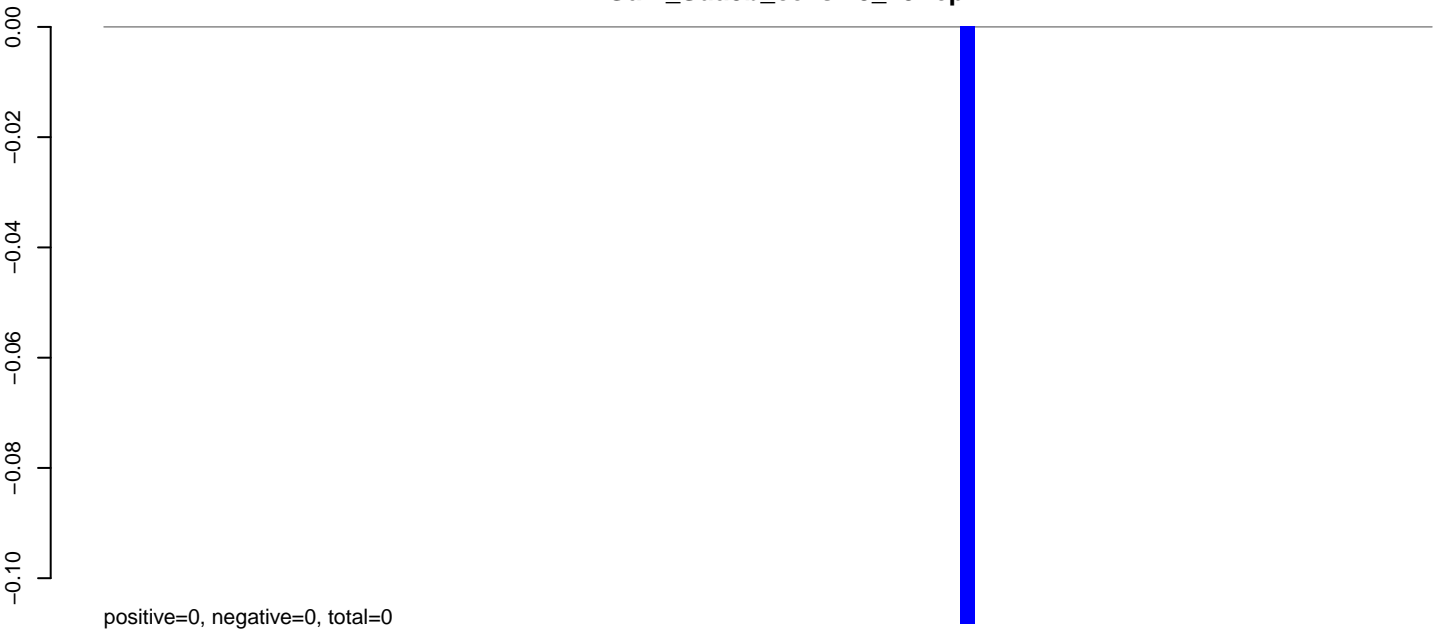


AnGam_Sua5b_cells.rep

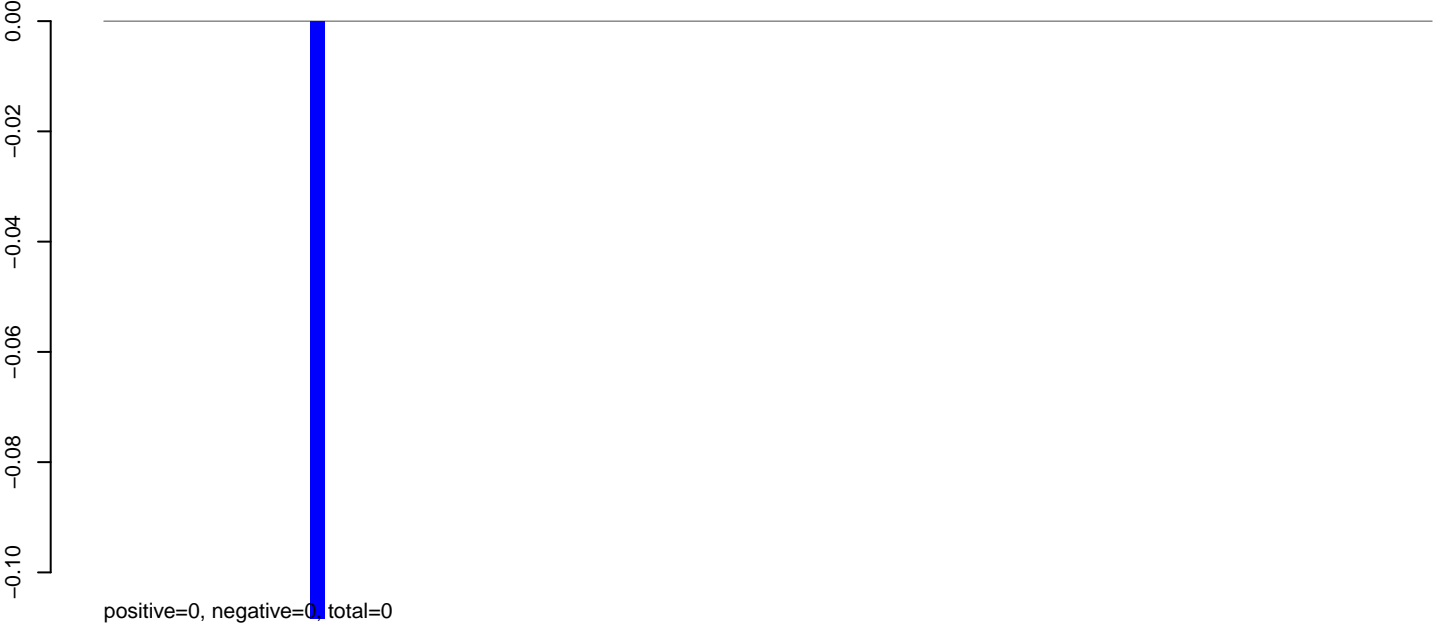


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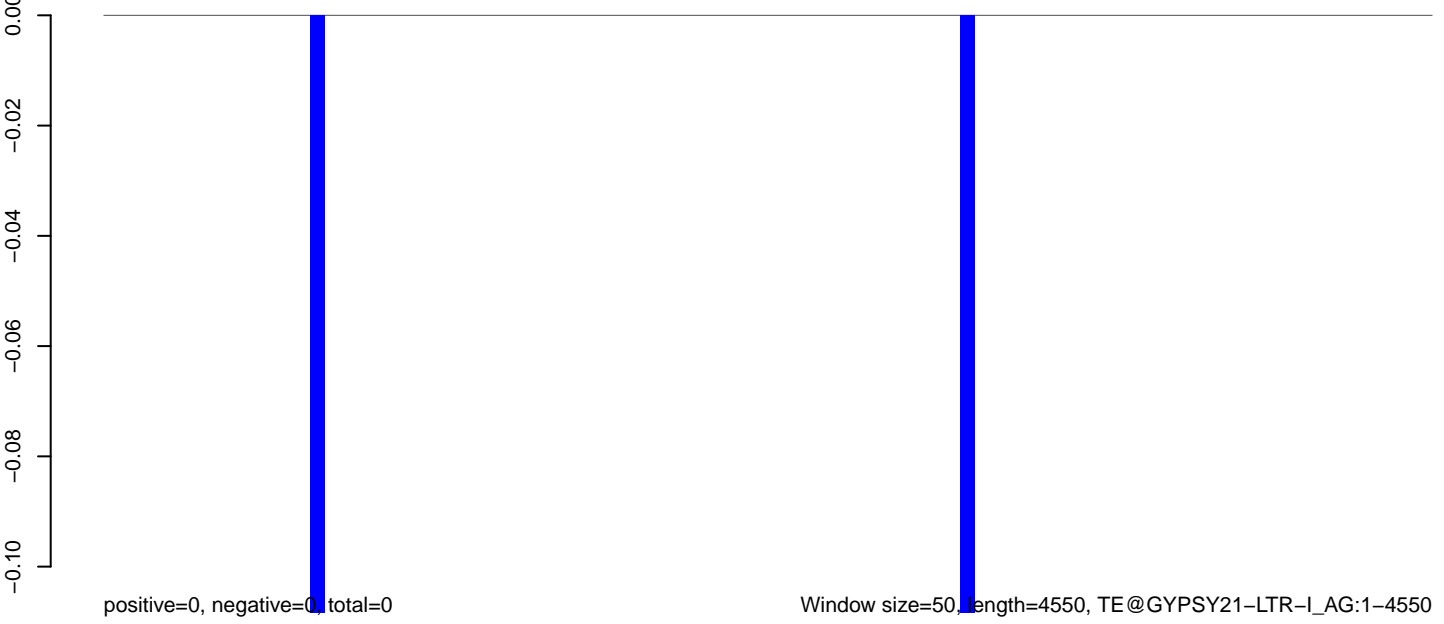
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AnGam_Sua5b_cells.24_35.rep

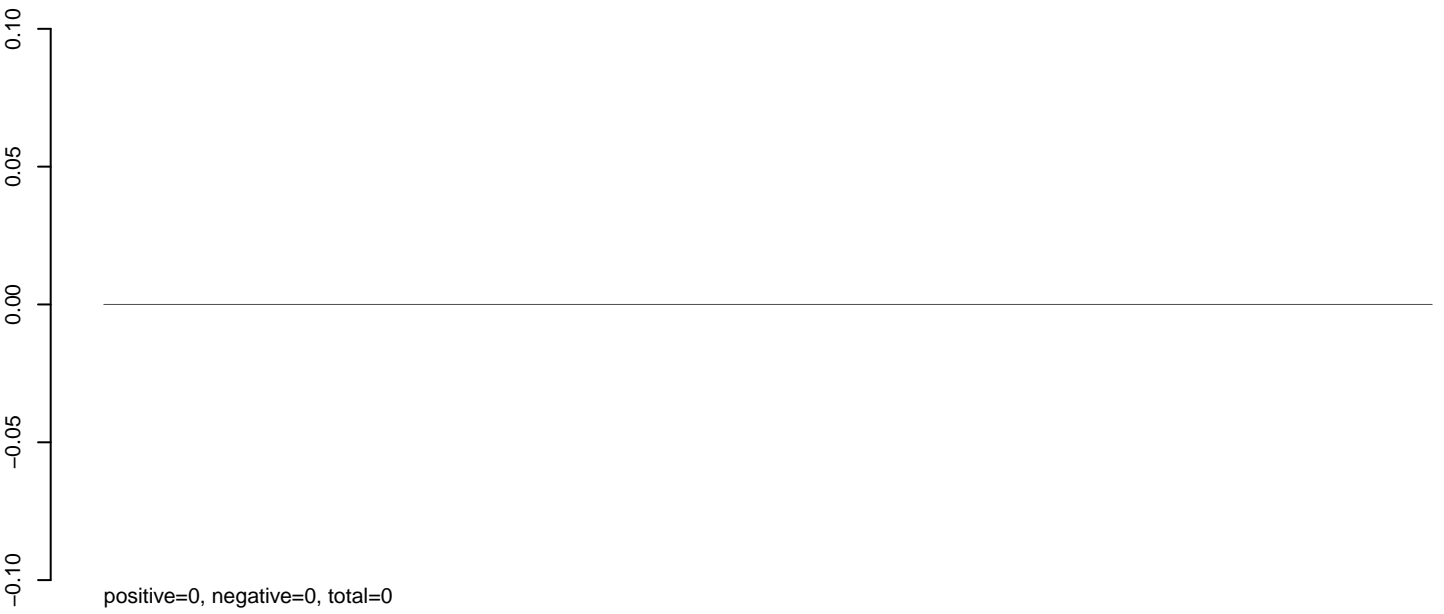


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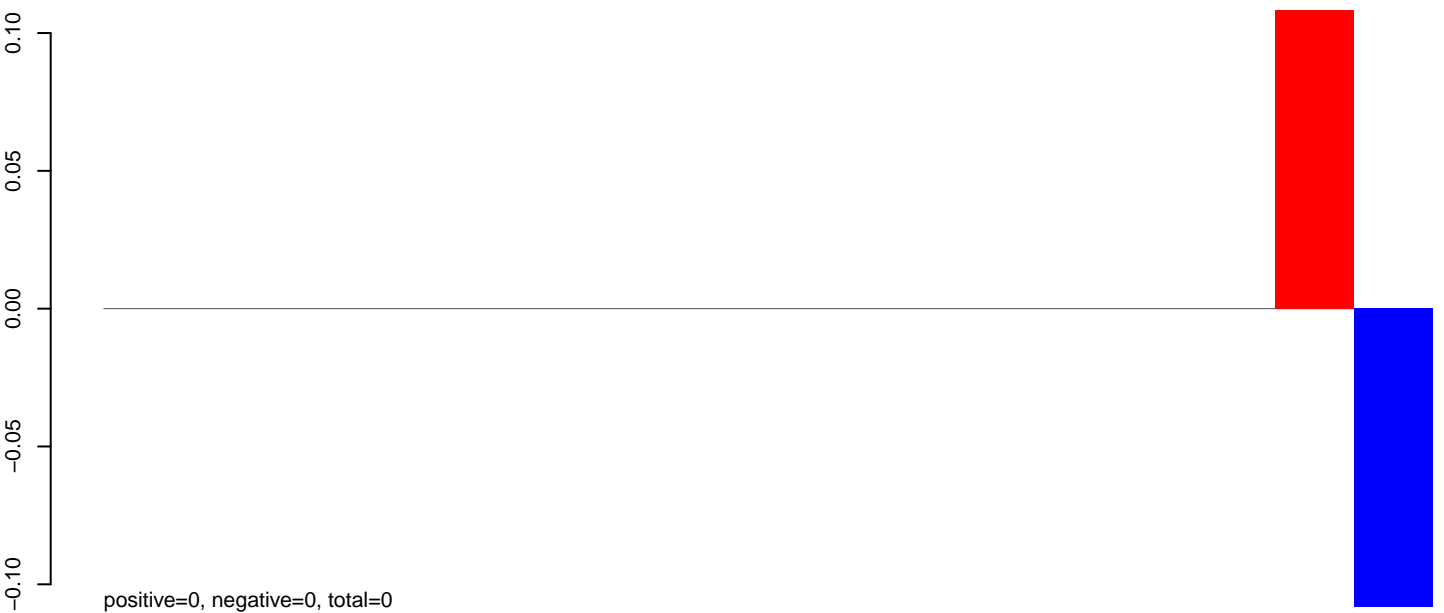


0 1000 2000 3000 4000

AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



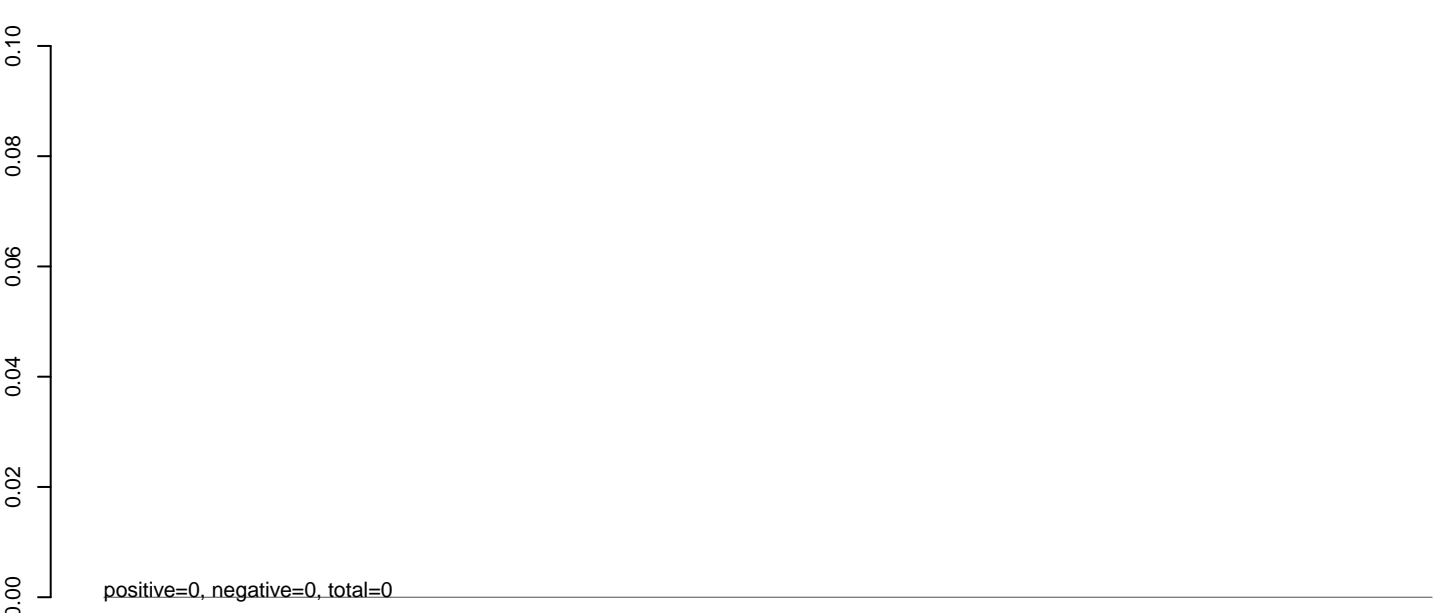
AnGam_Sua5b_cells.rep



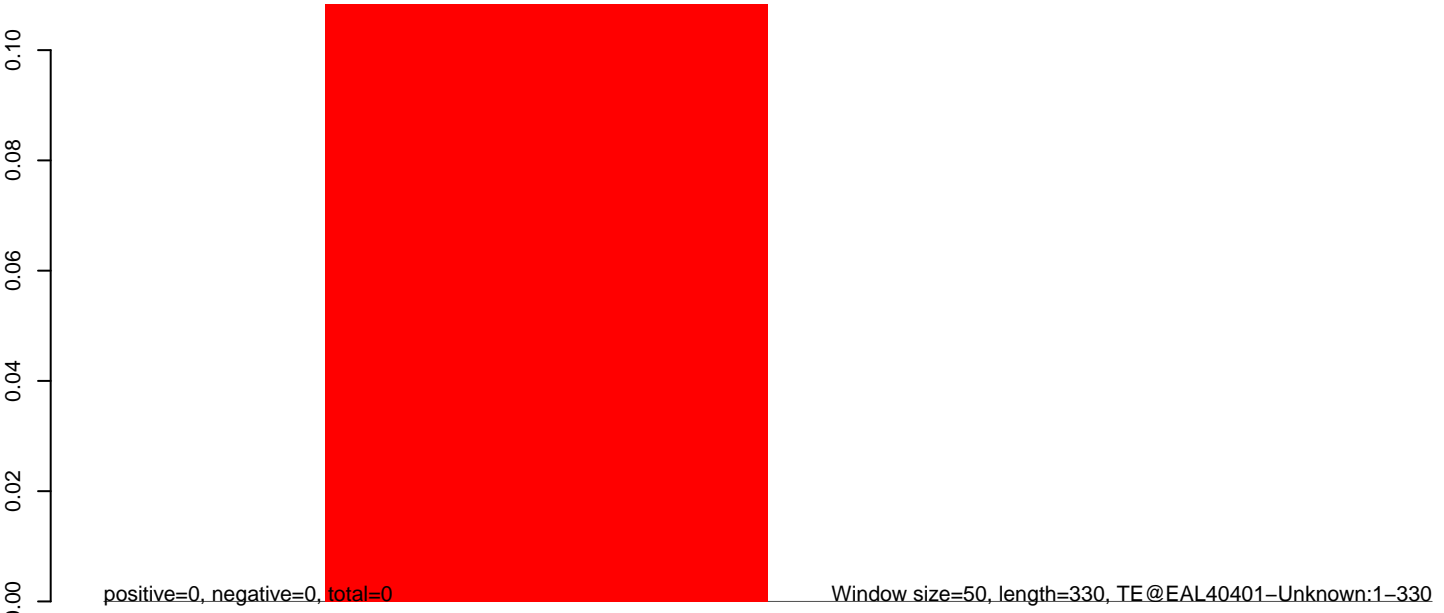
AnGam_Sua5b_cells.18_23.rep



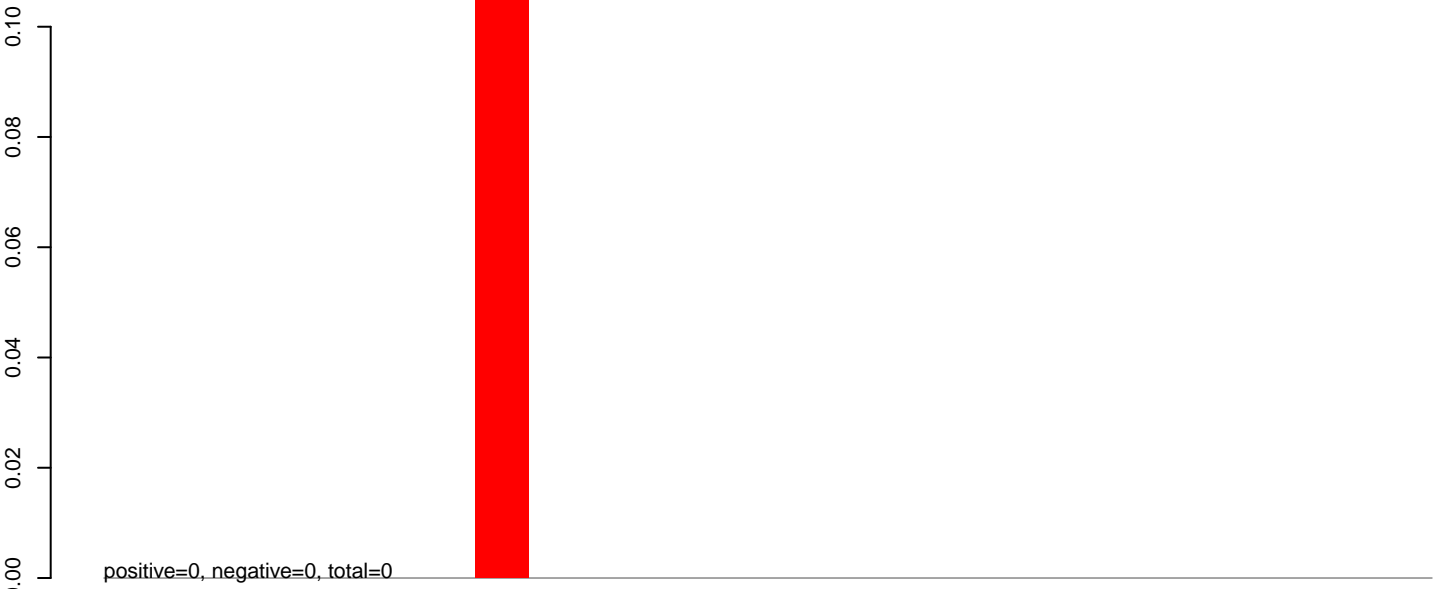
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



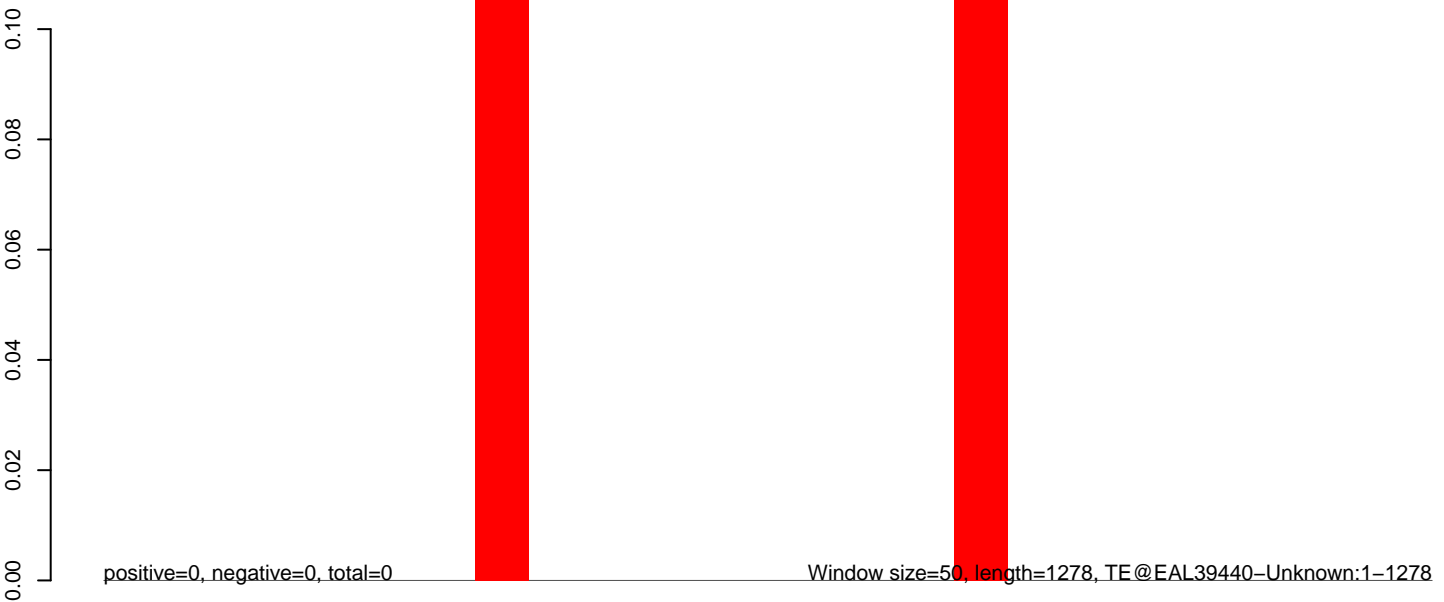
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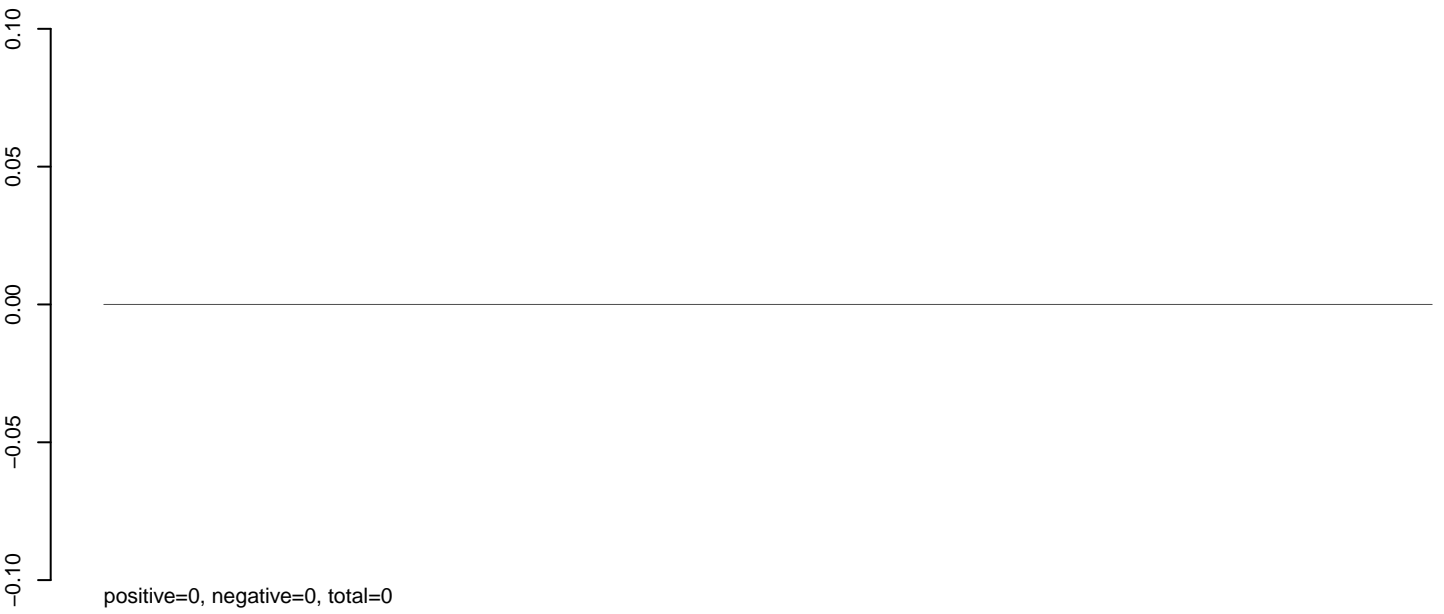
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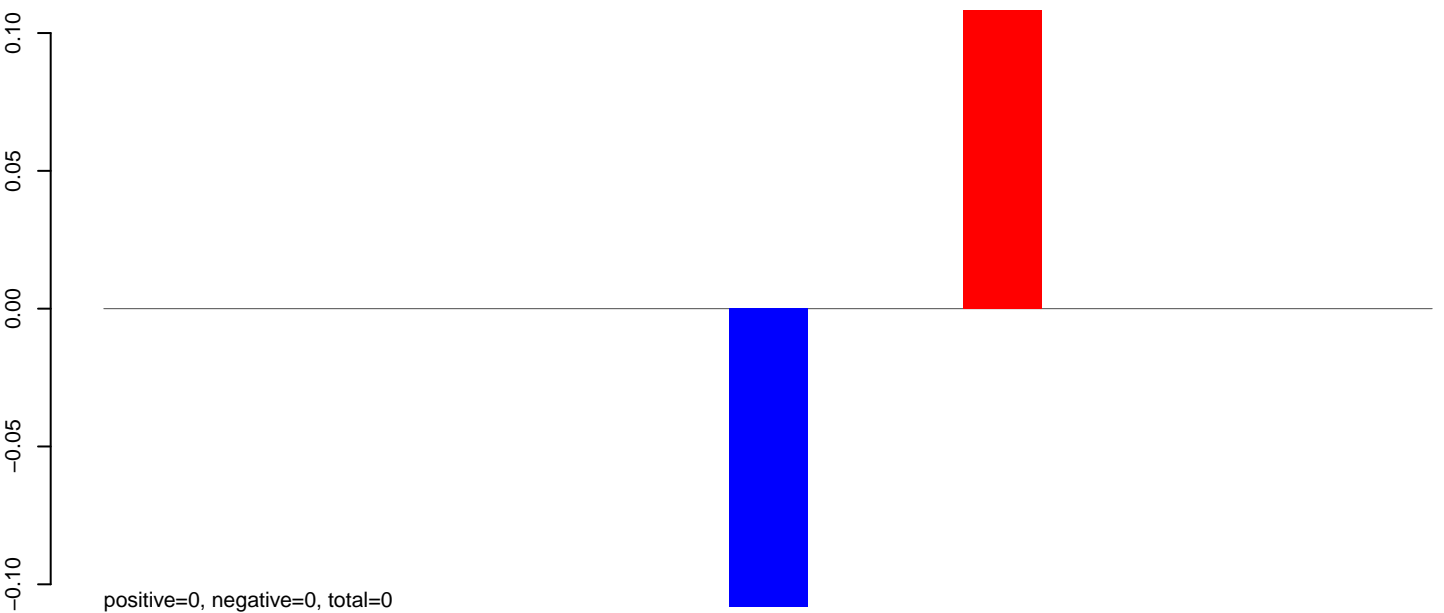
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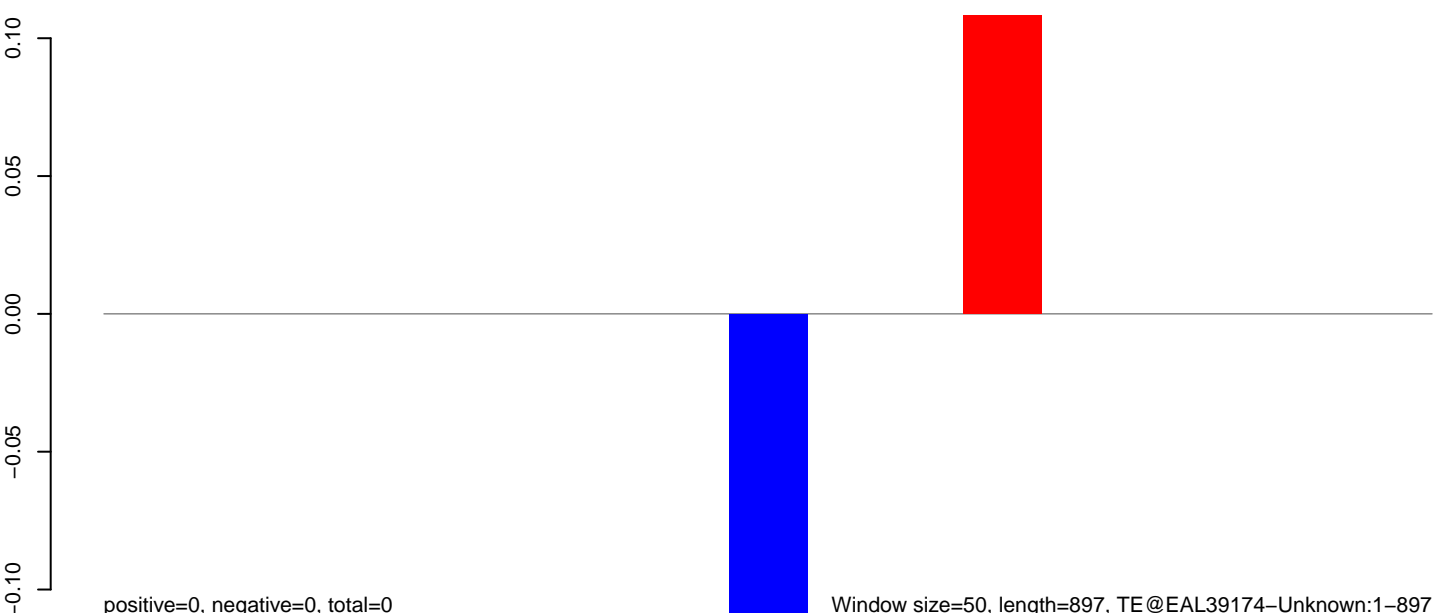
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AnGam_Sua5b_cells.24_35.rep



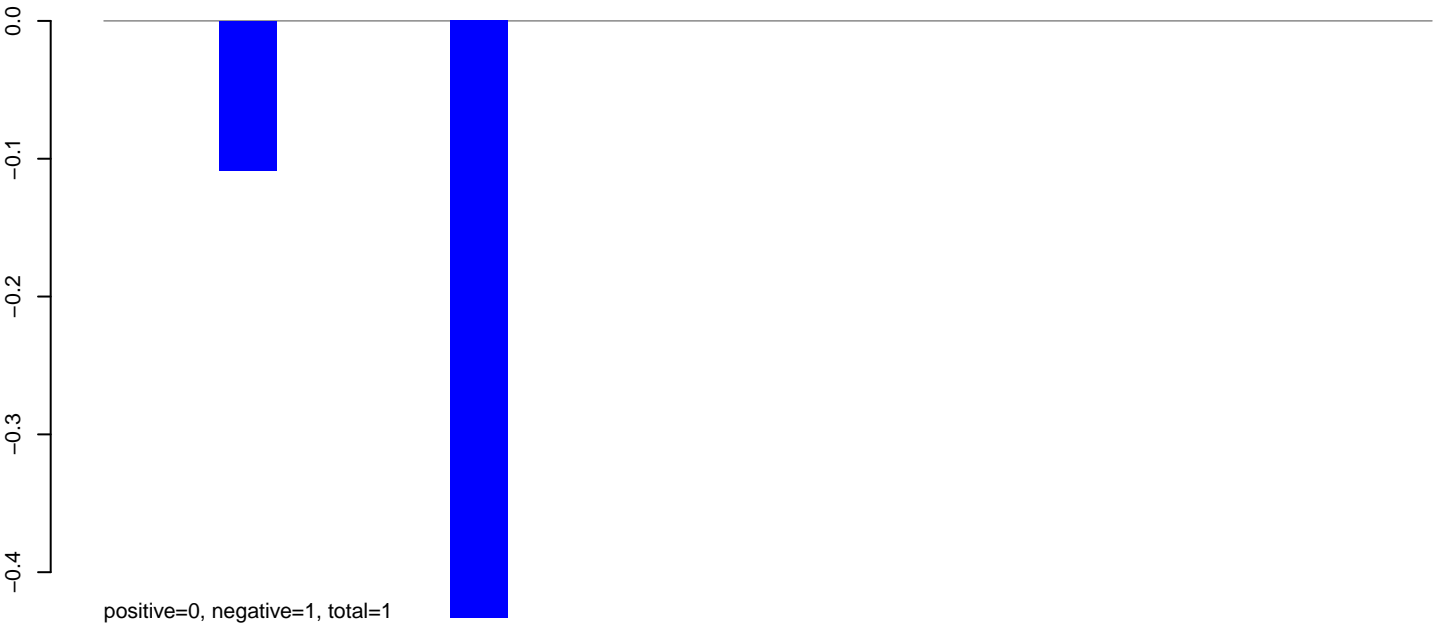
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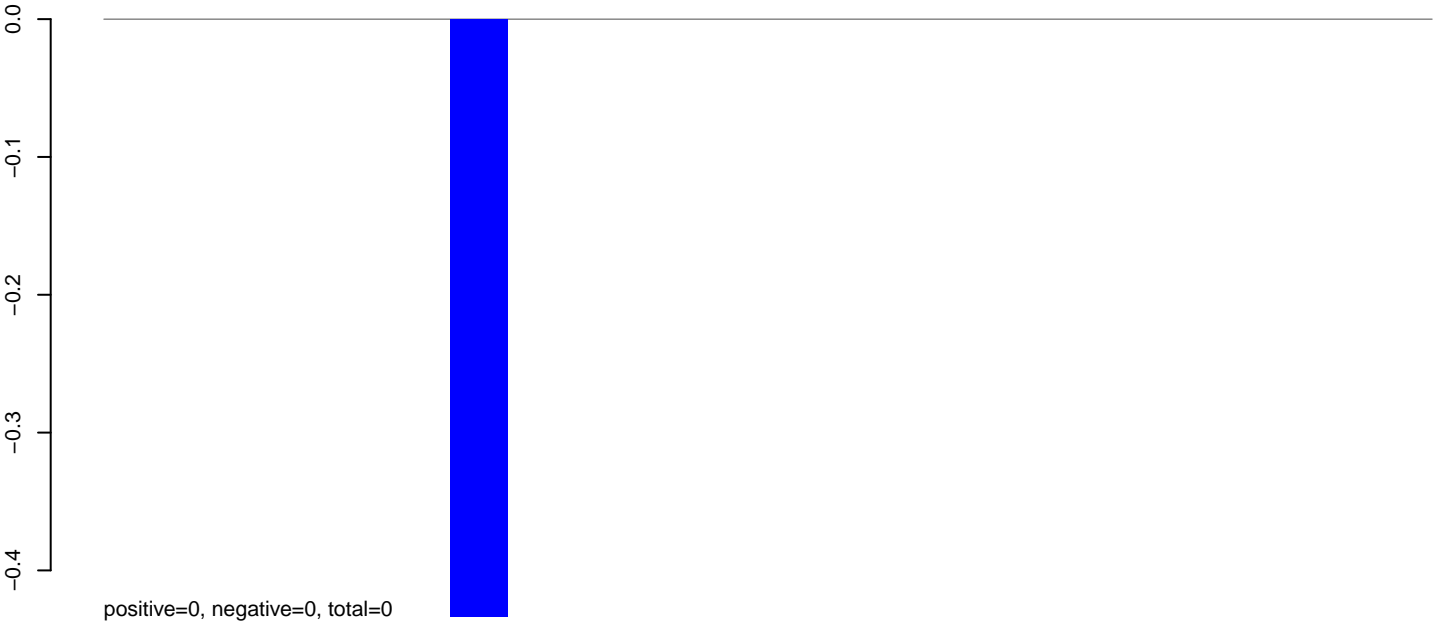
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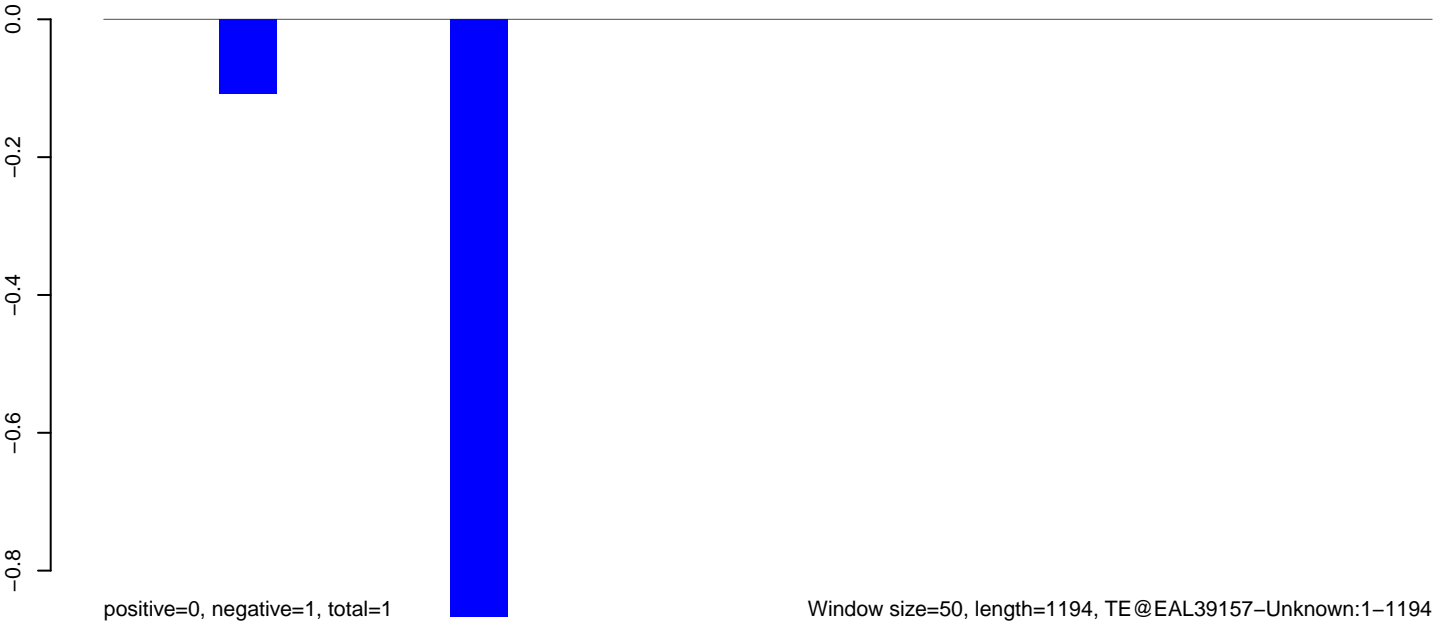
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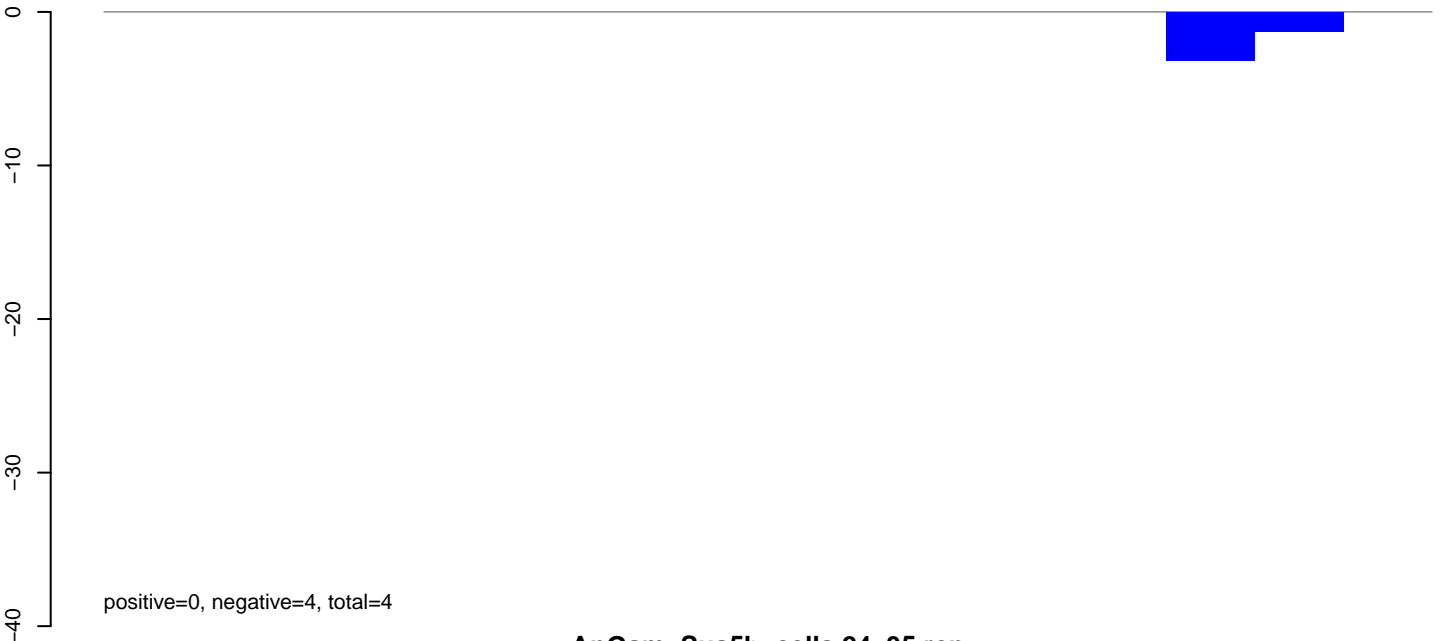
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AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



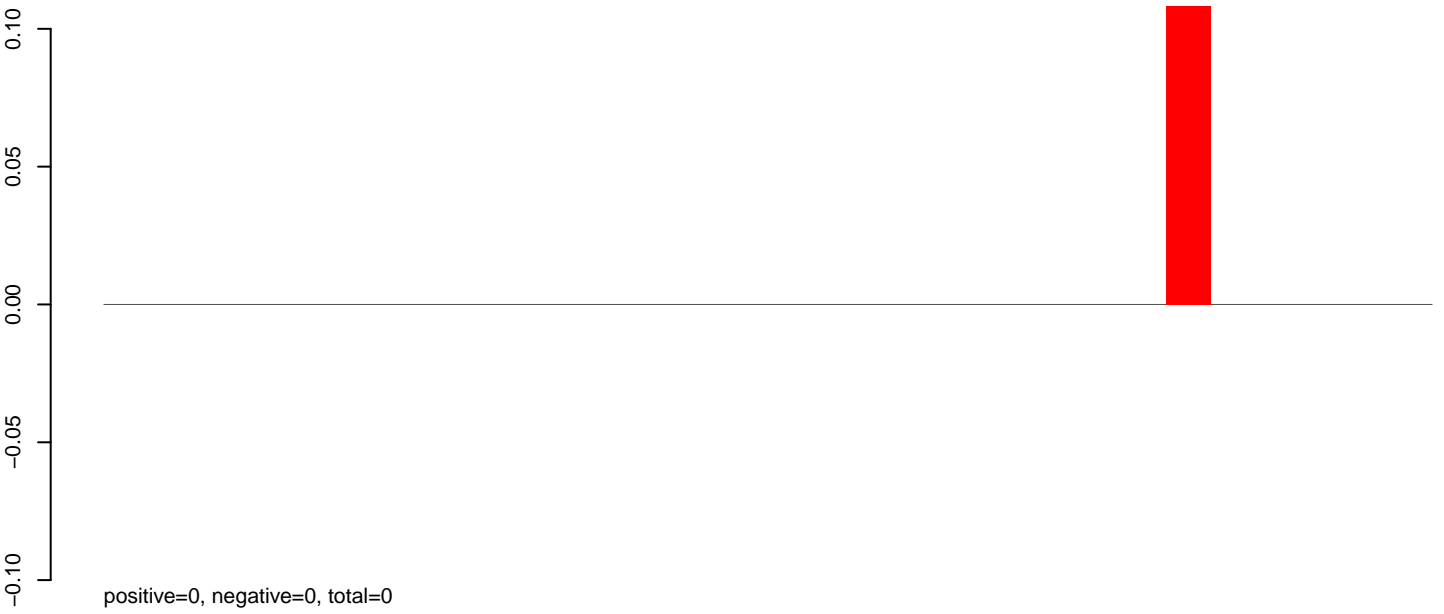
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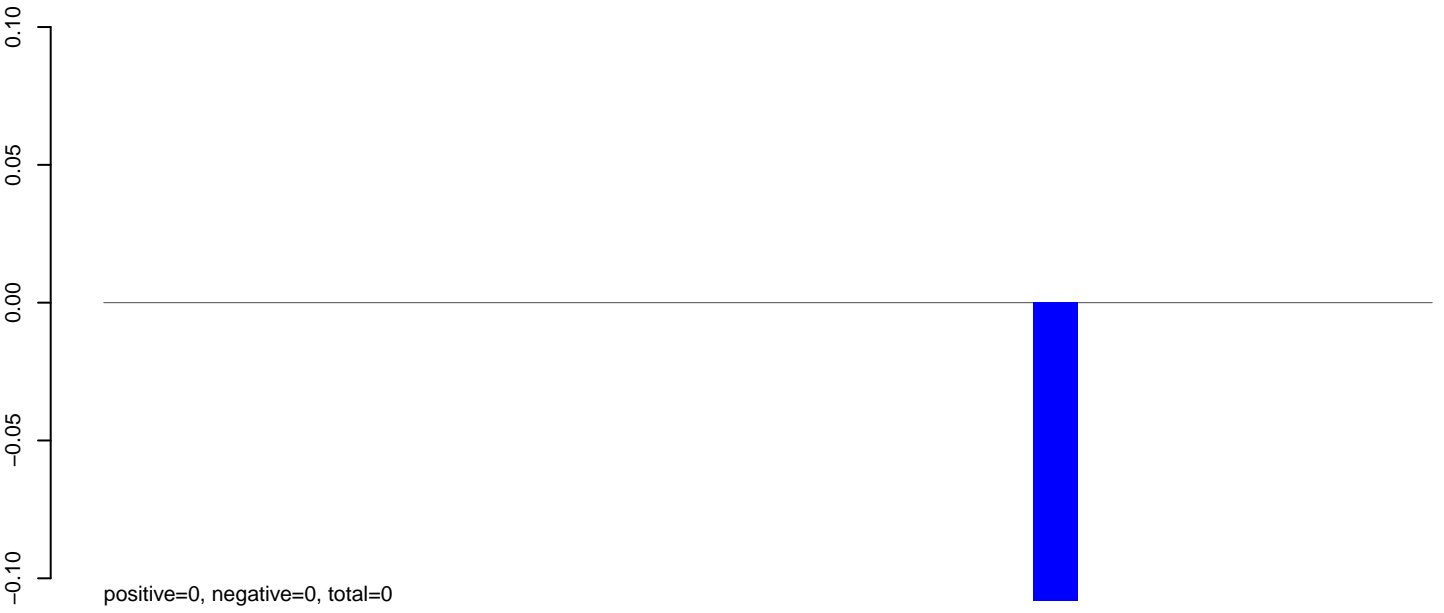
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AnGam_Sua5b_cells.18_23.rep



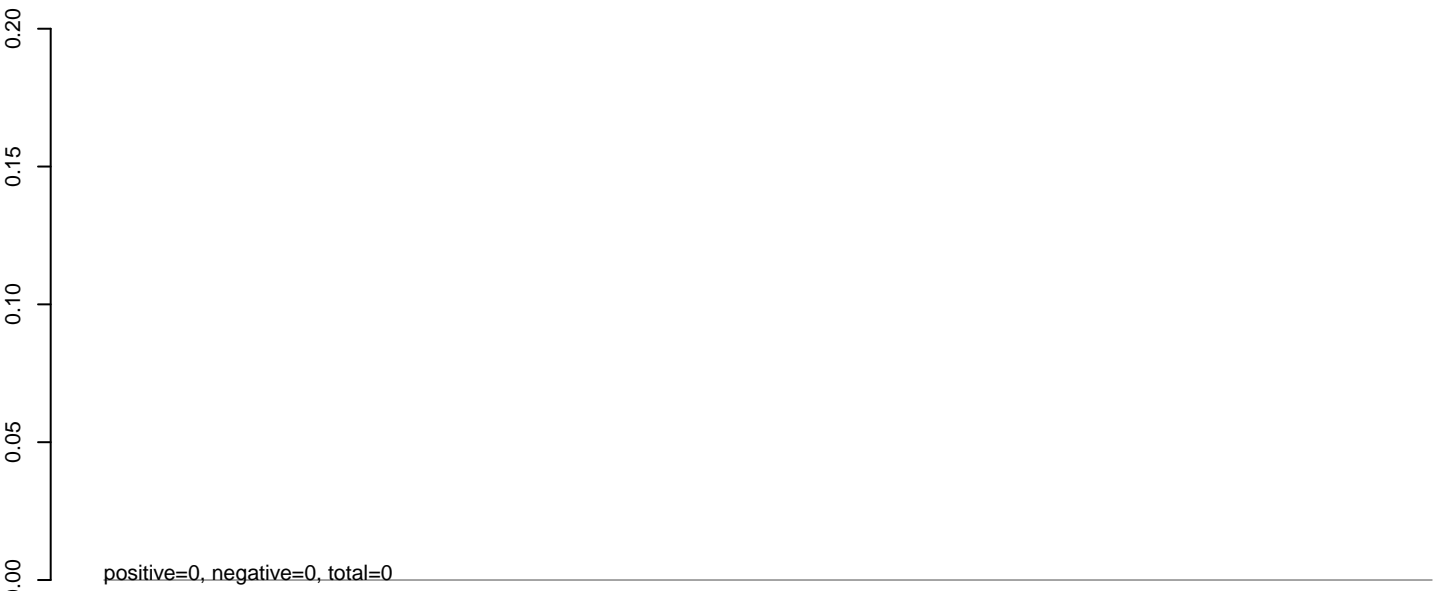
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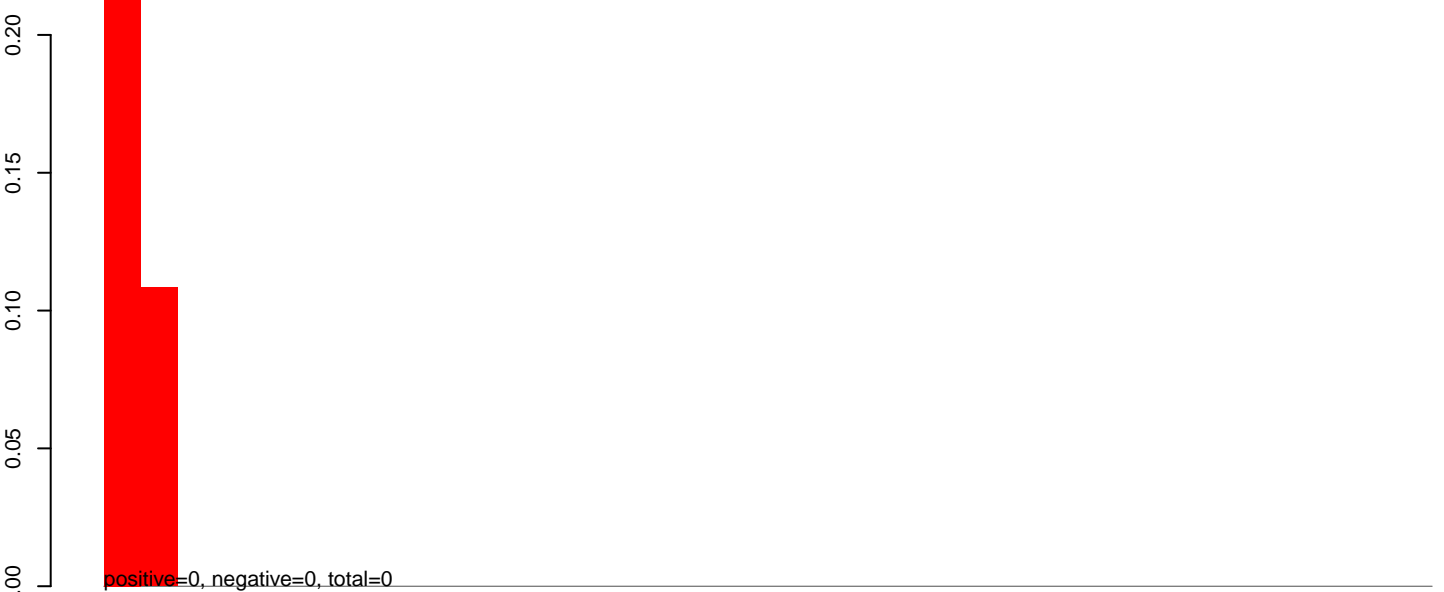
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AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



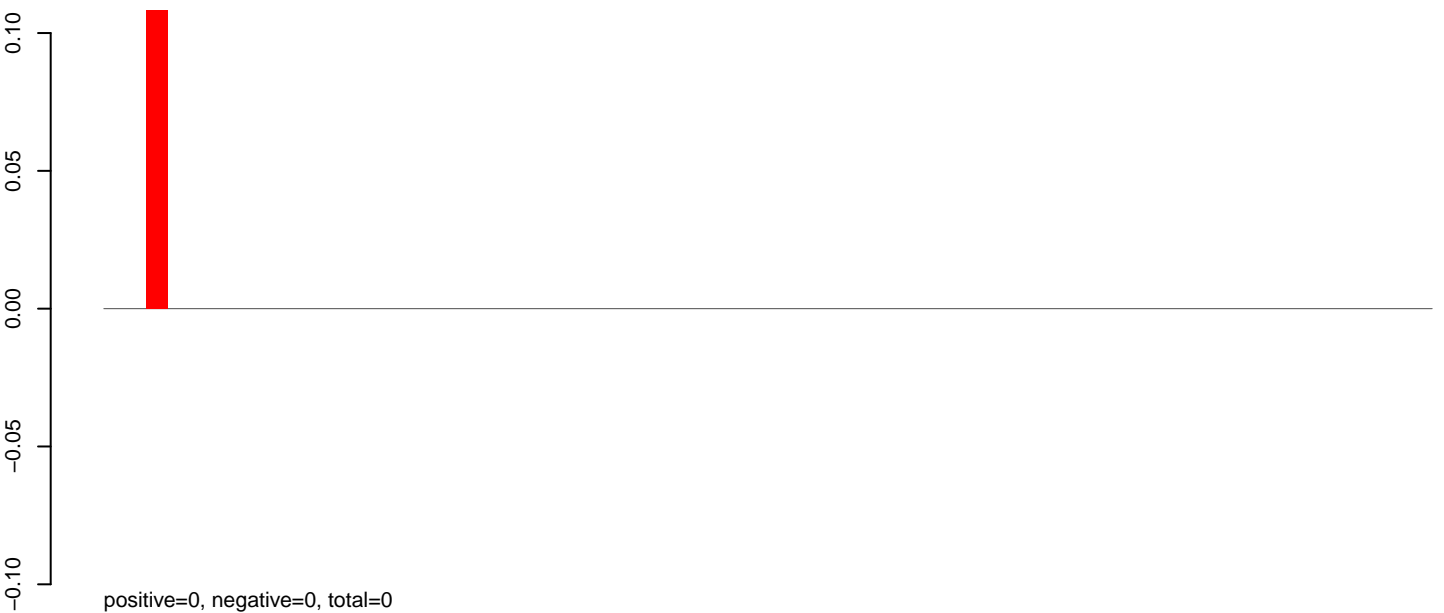
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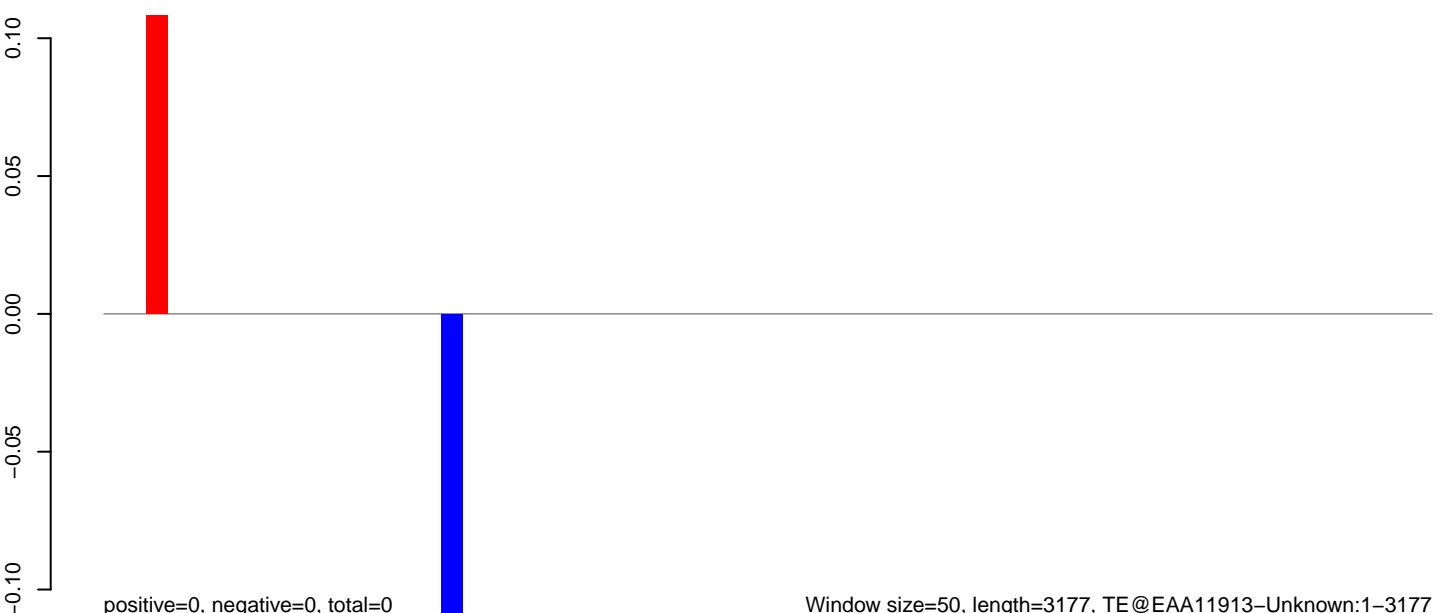
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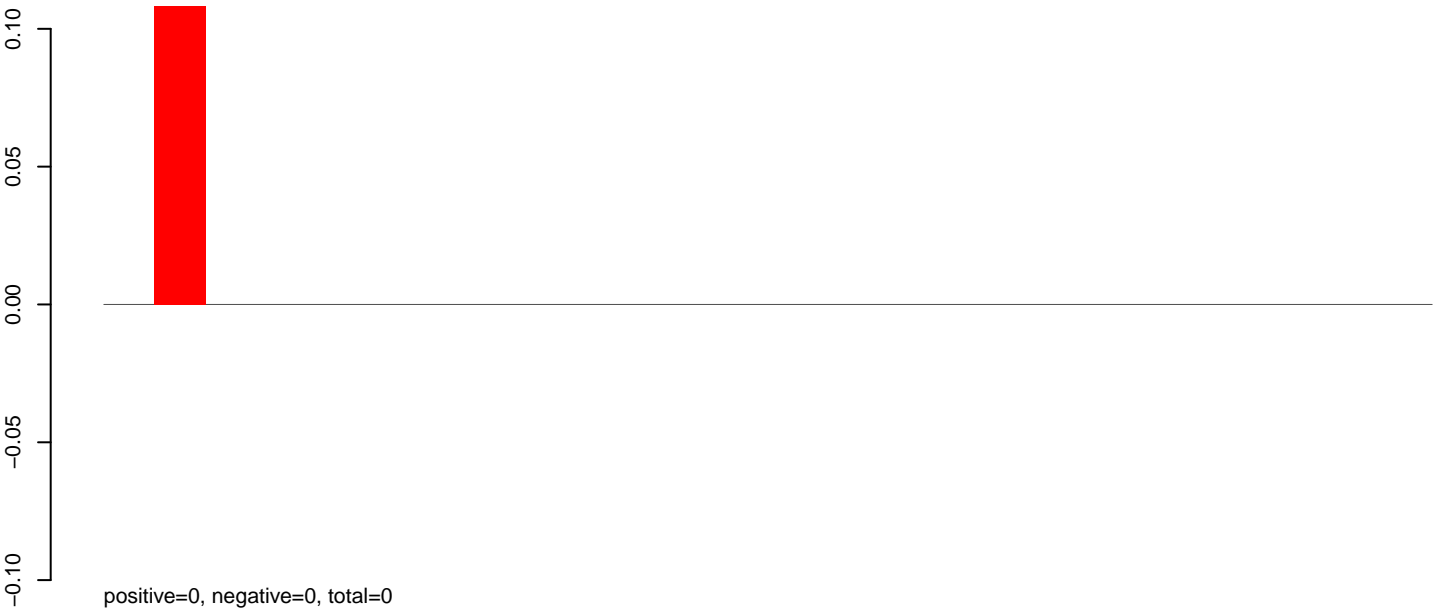
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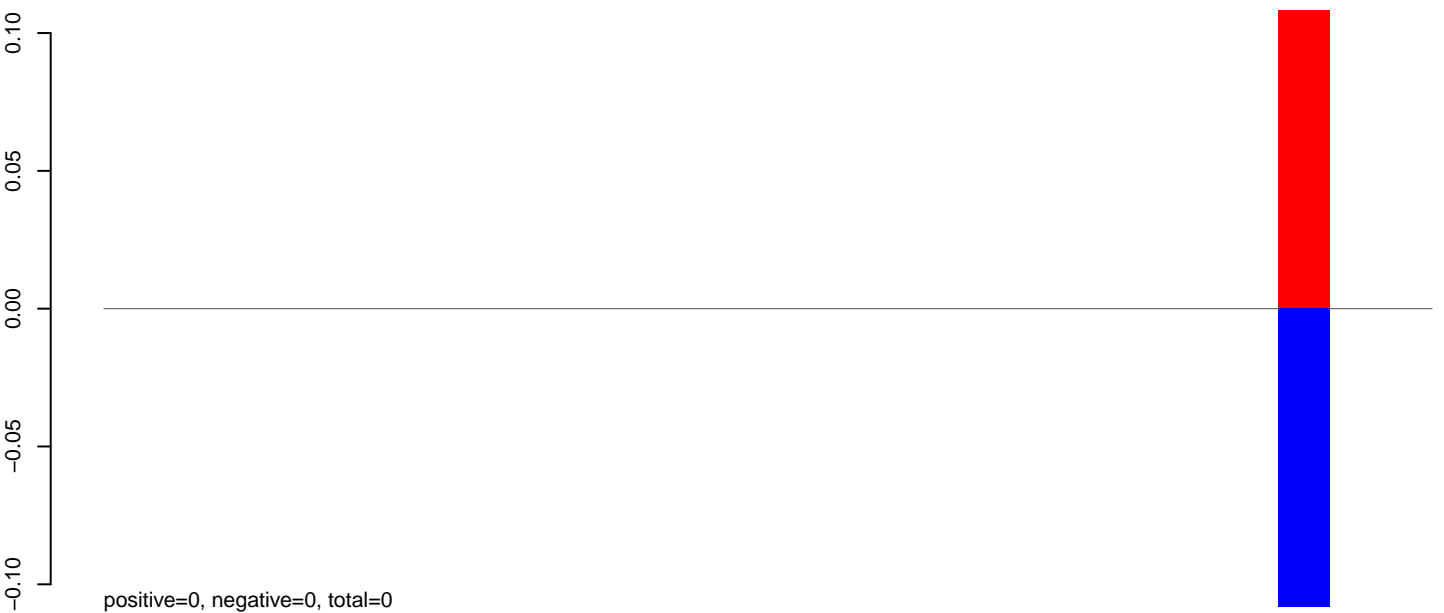
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AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



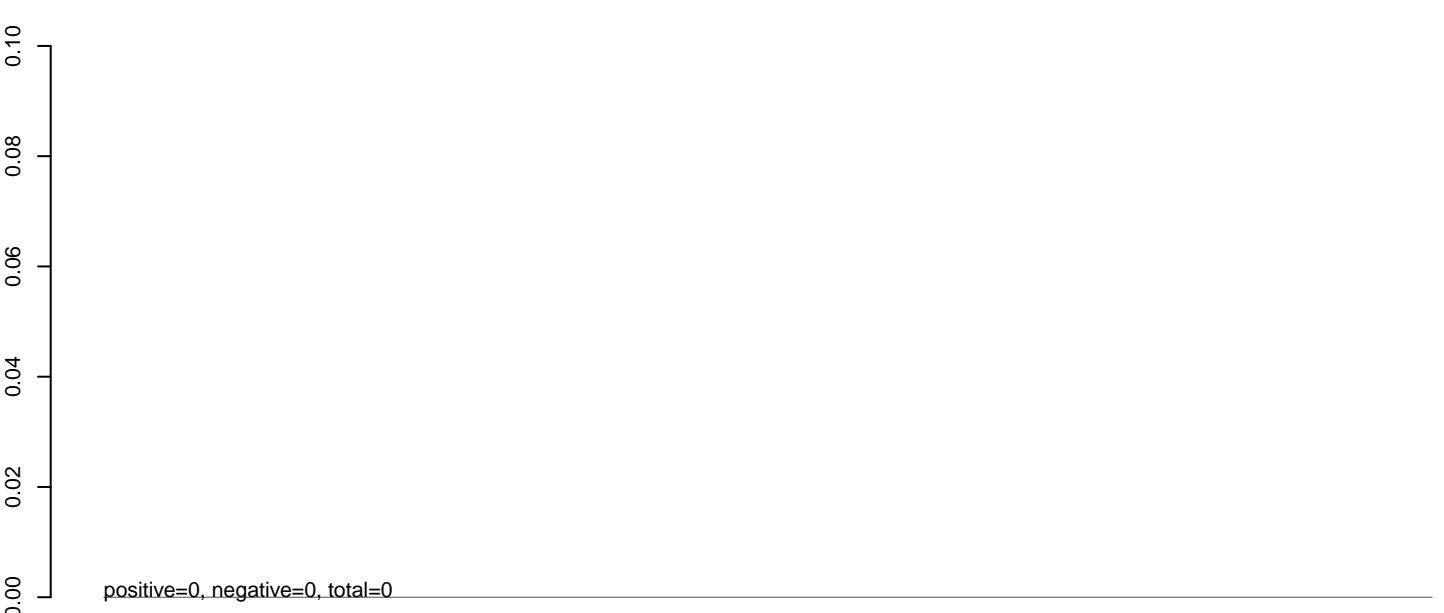
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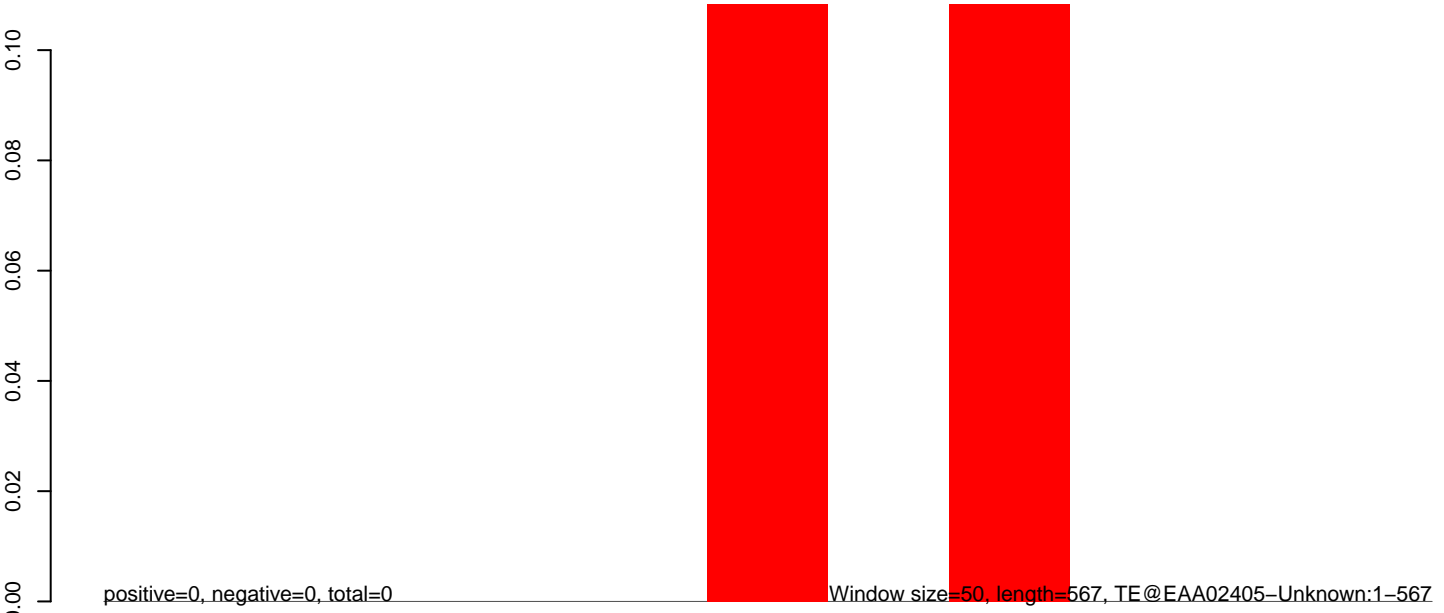
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AnGam_Sua5b_cells.24_35.rep

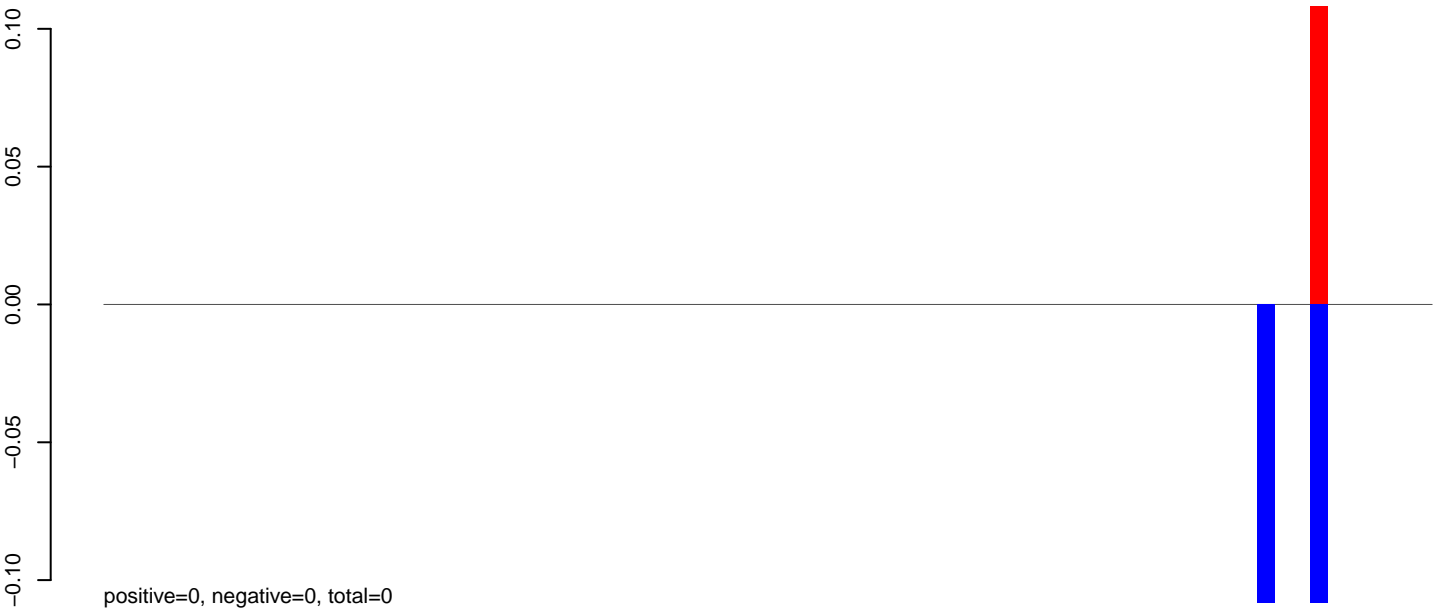


AnGam_Sua5b_cells.rep

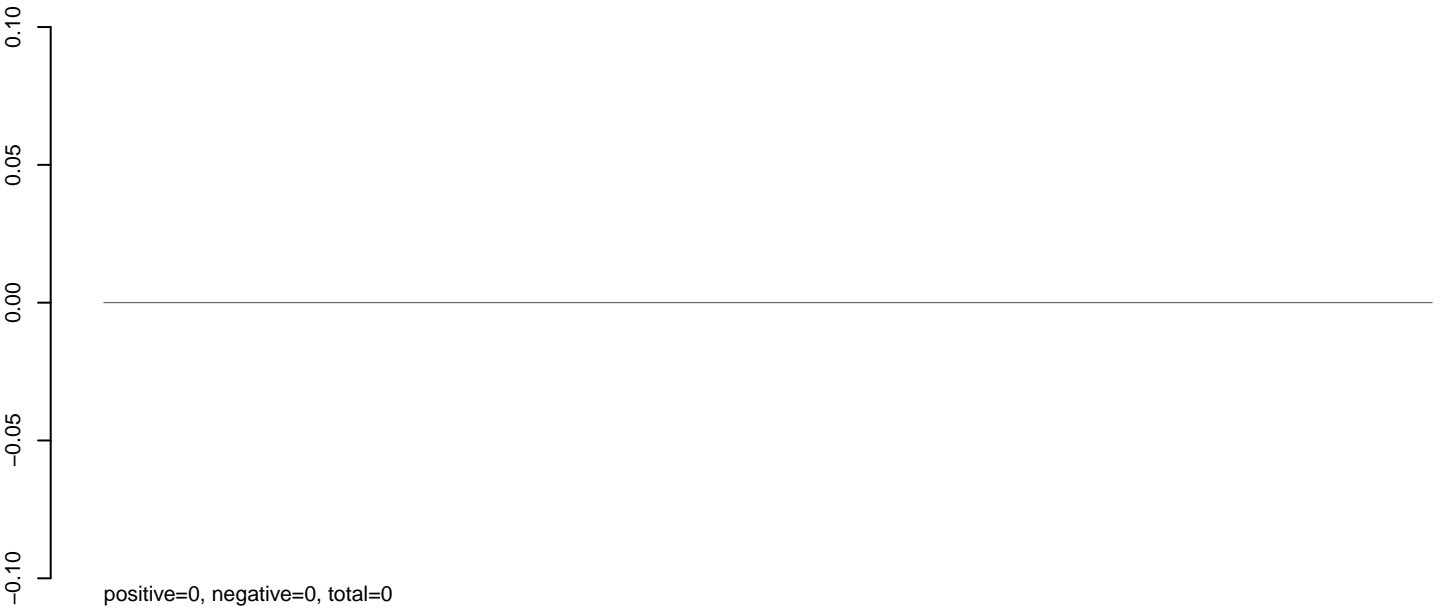


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AnGam_Sua5b_cells.18_23.rep



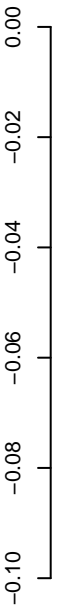
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

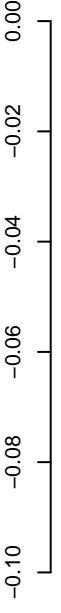


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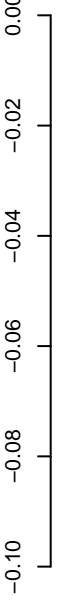
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AnGam_Sua5b_cells.24_35.rep



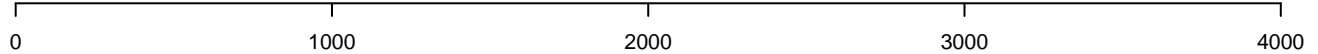
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AnGam_Sua5b_cells.rep

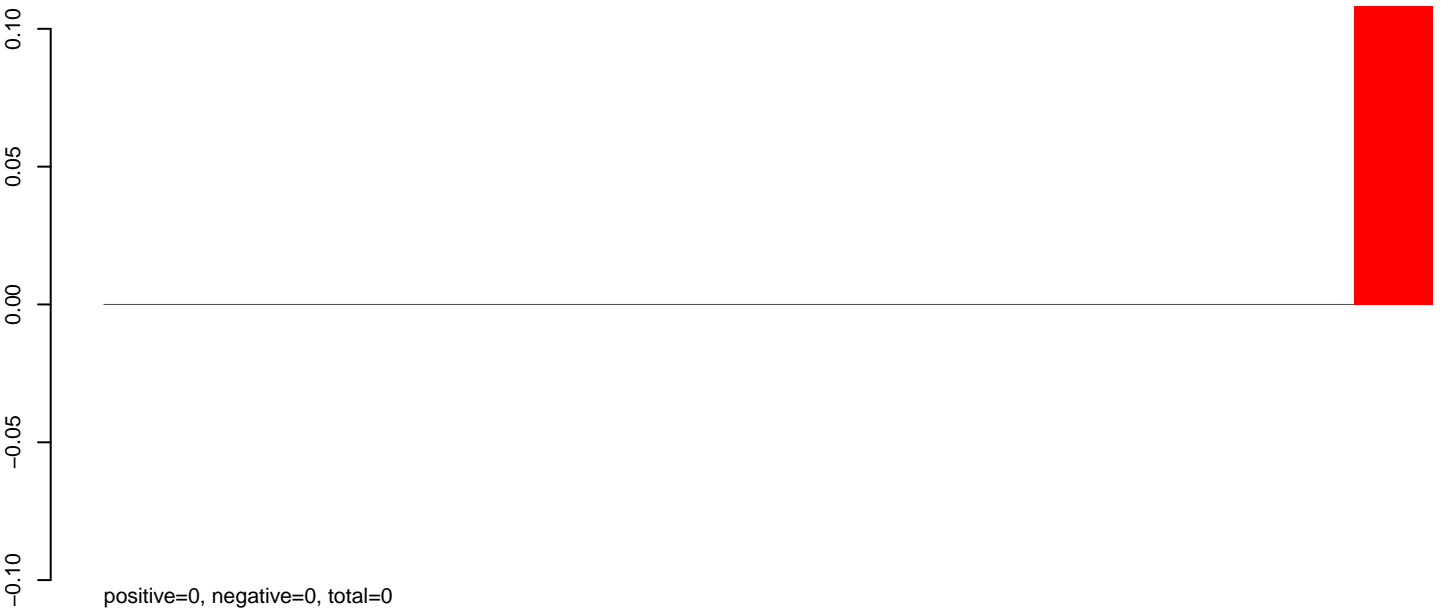


positive=0, negative=0, total=0

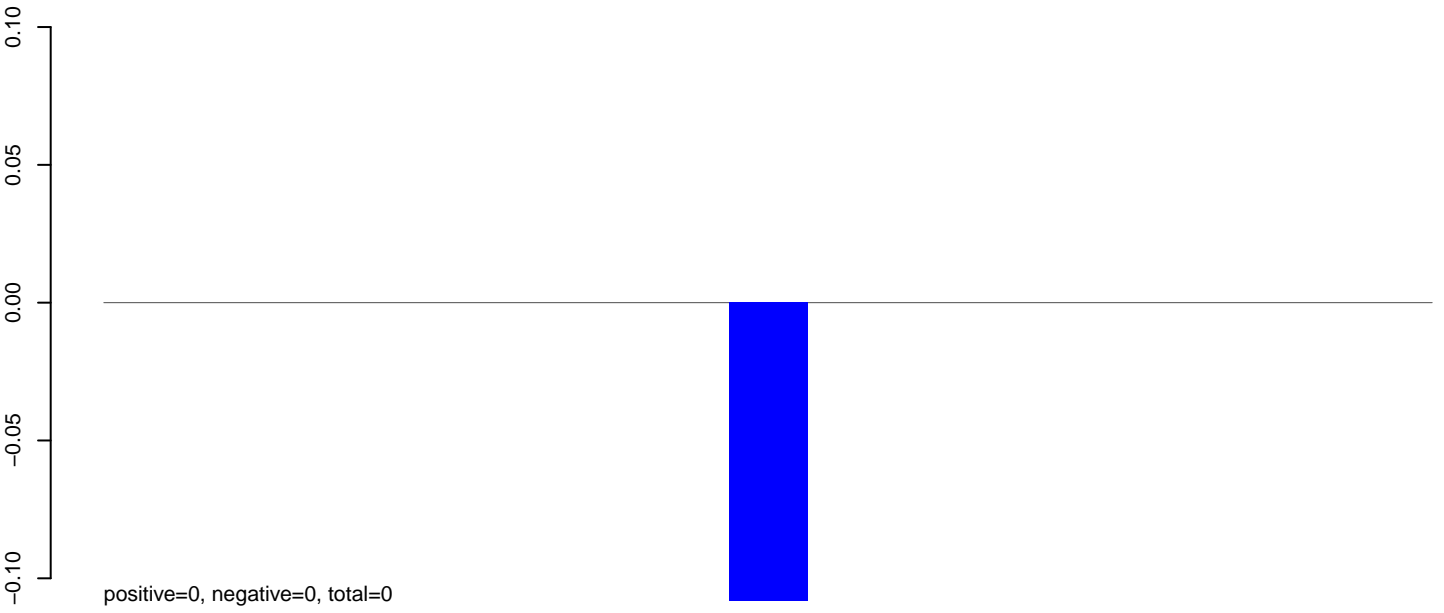
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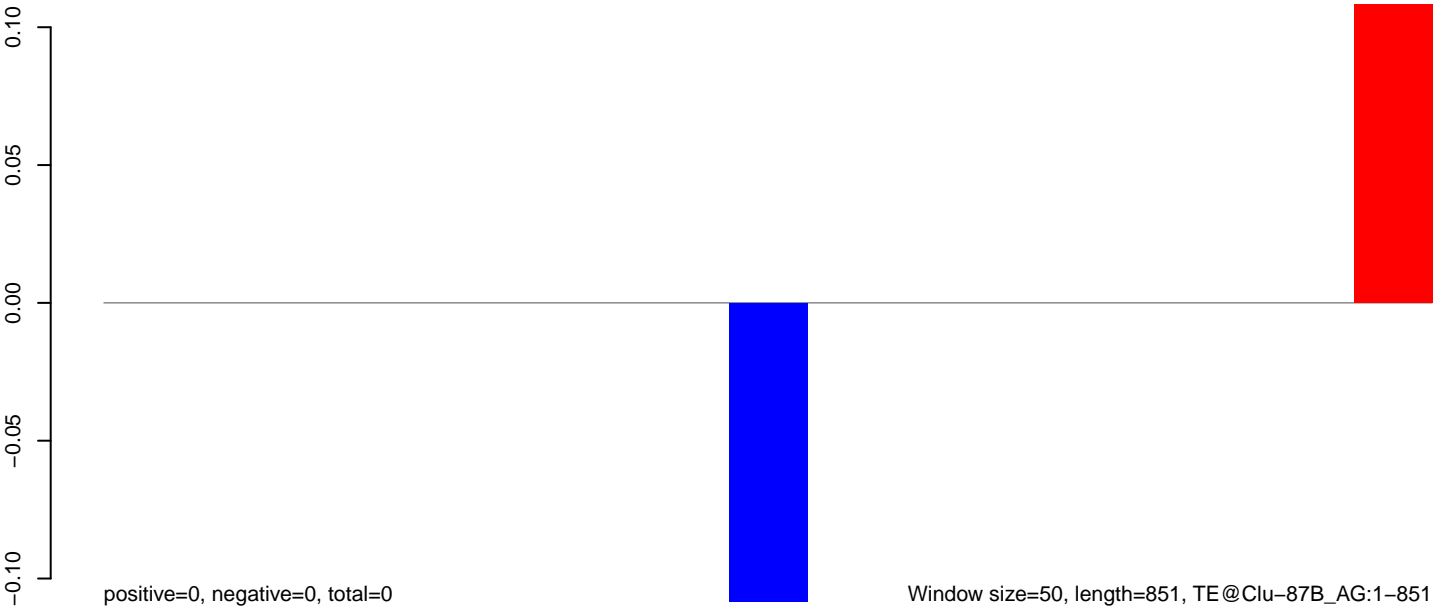
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

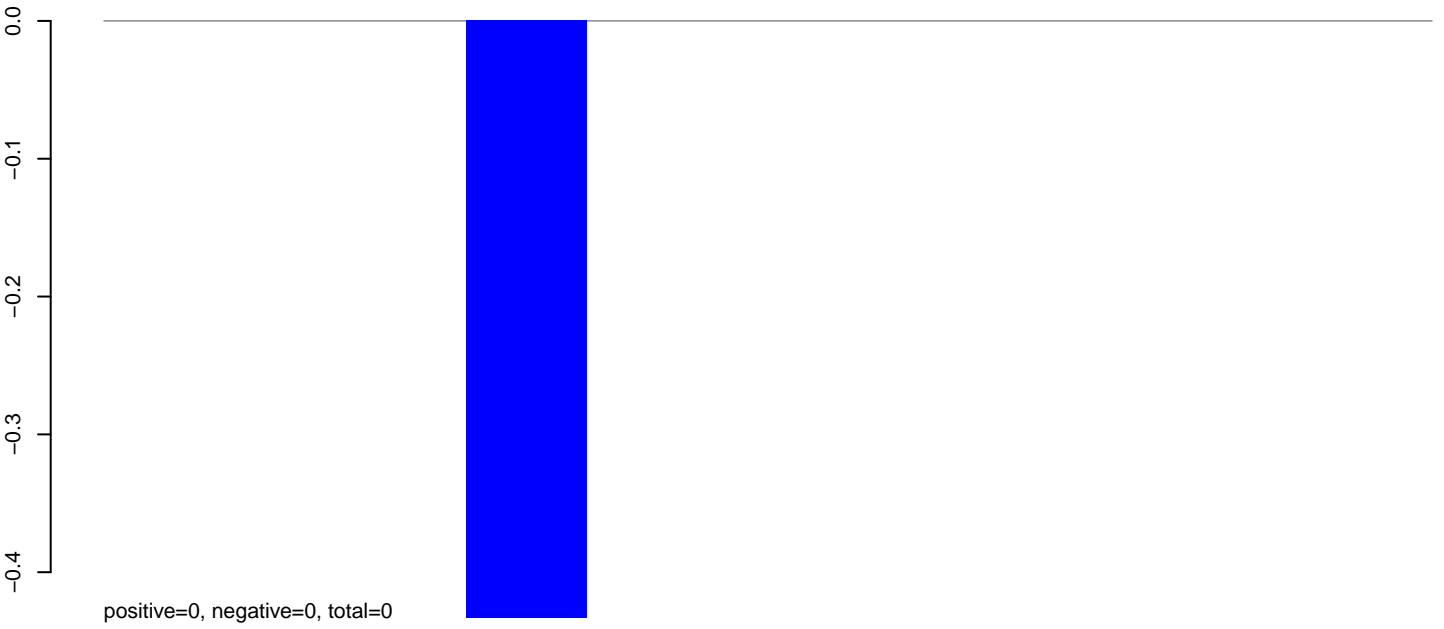


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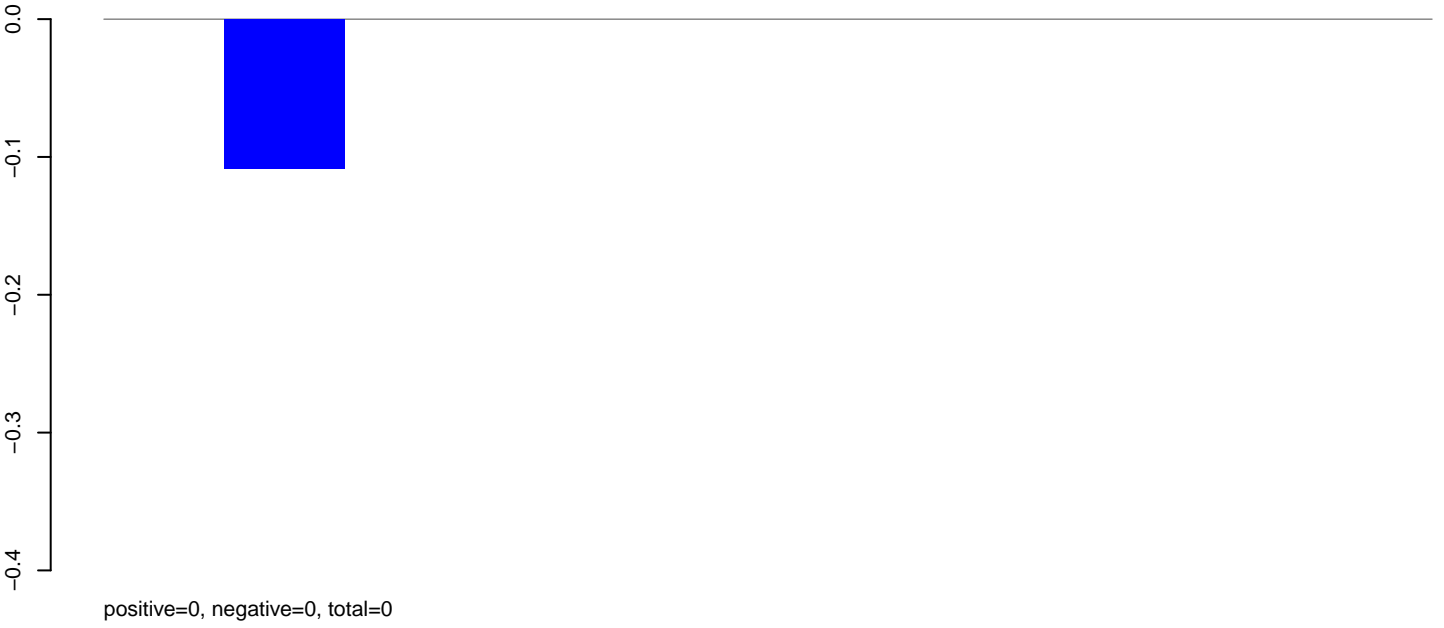


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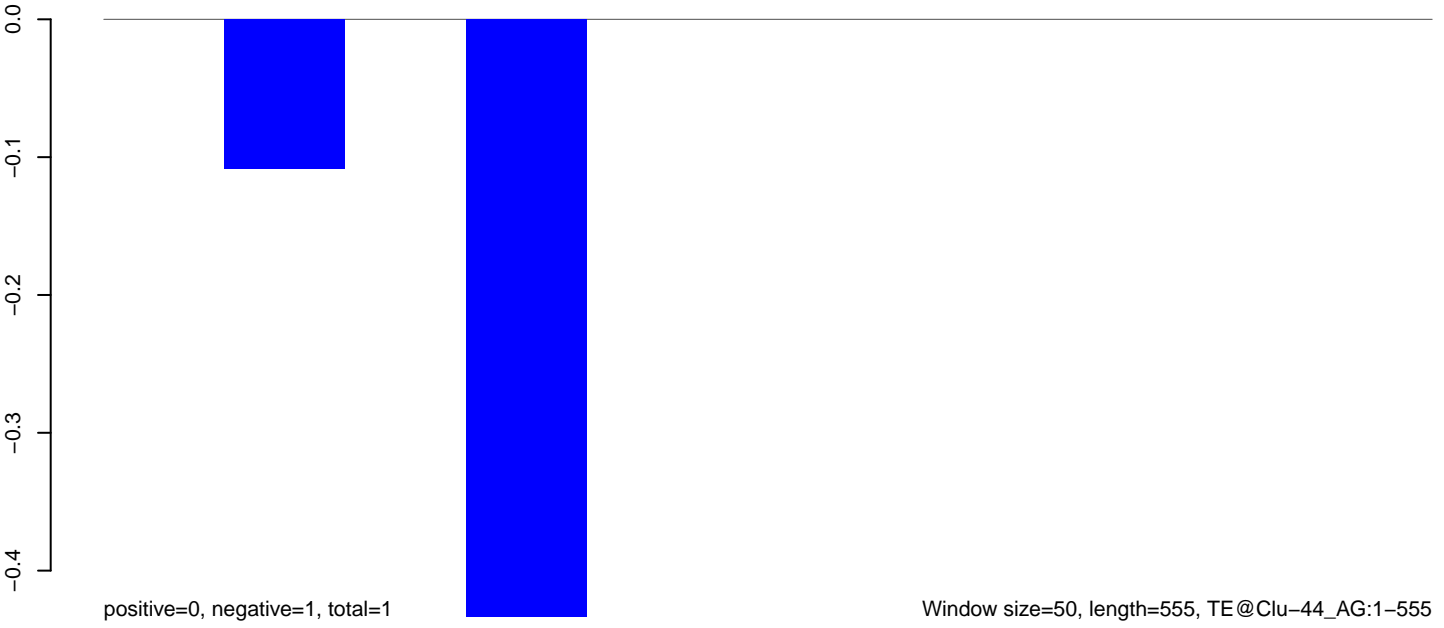
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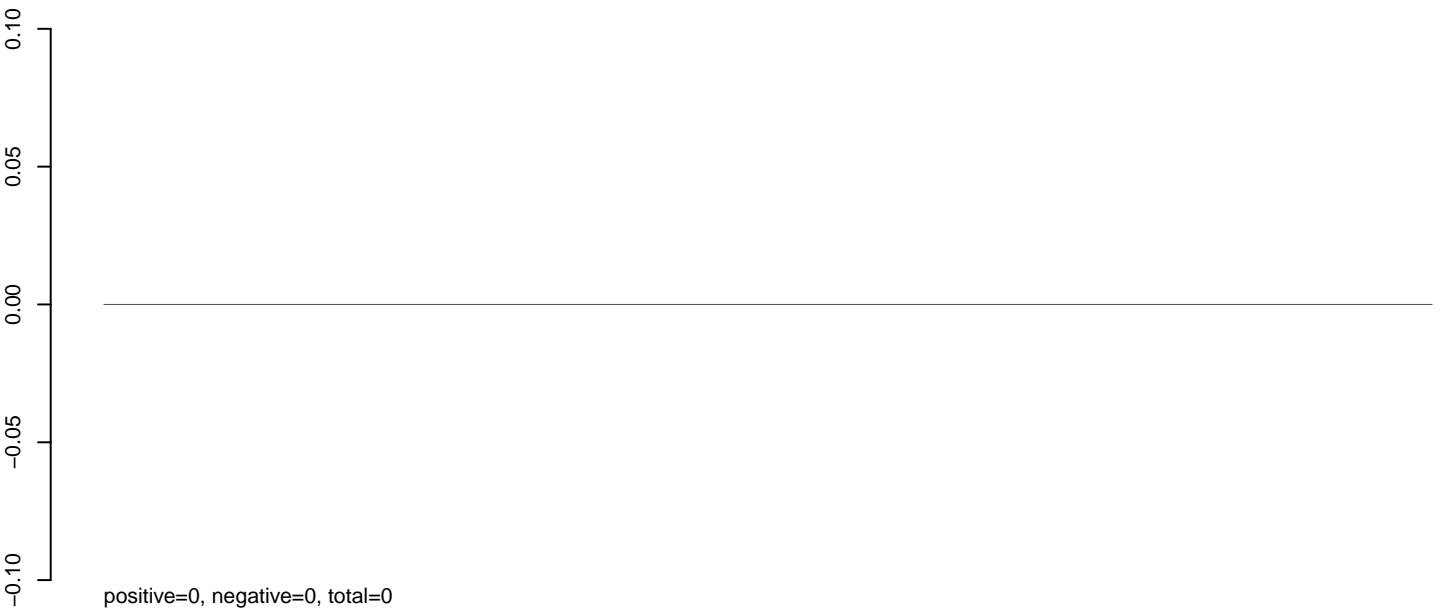
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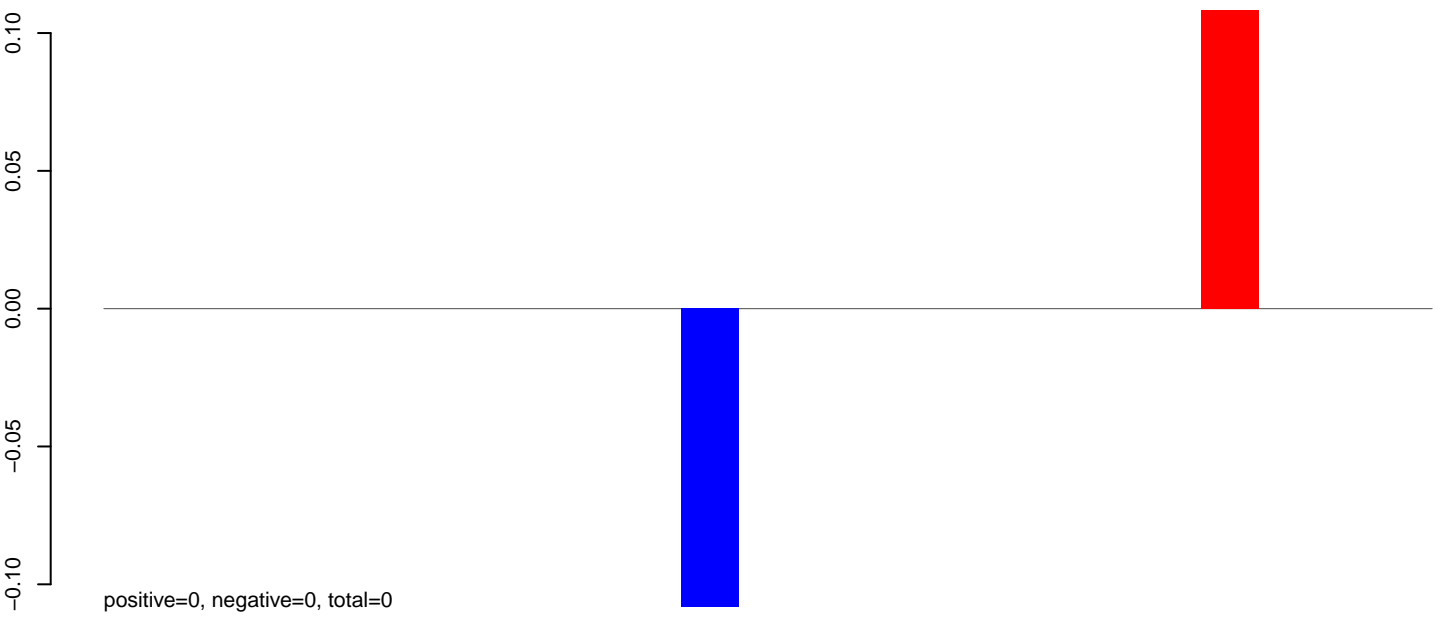
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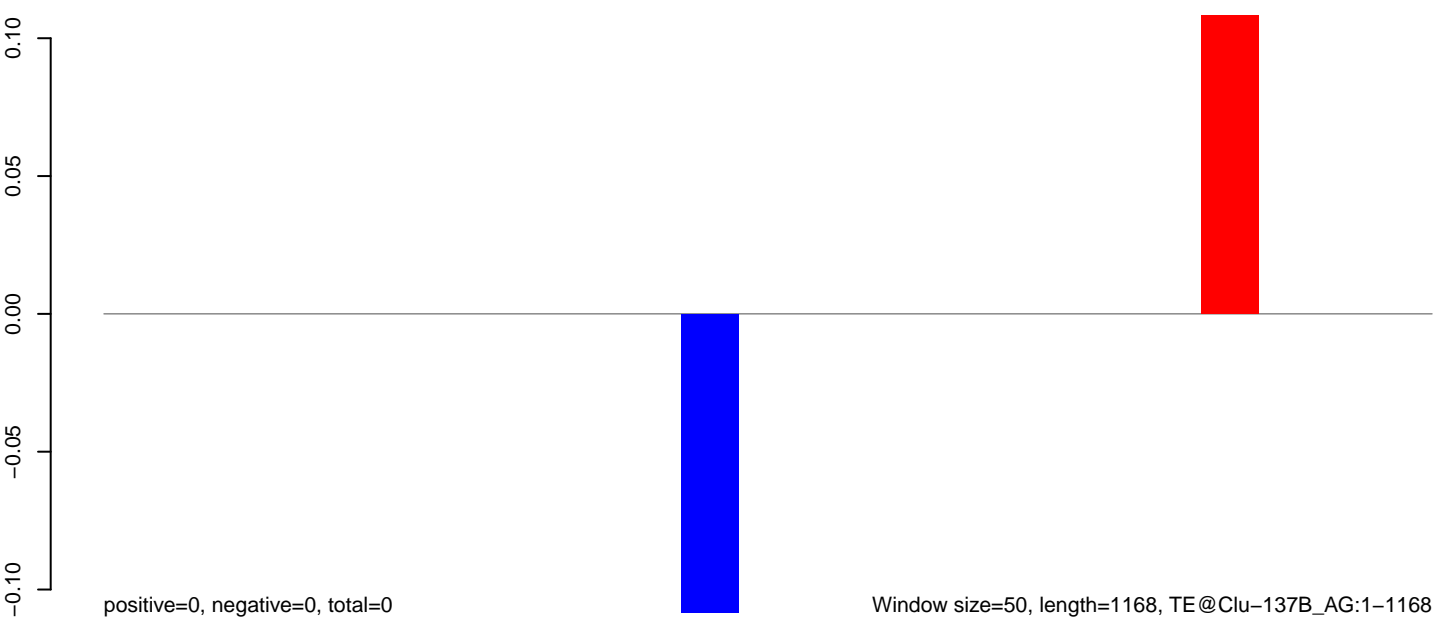
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AnGam_Sua5b_cells.24_35.rep



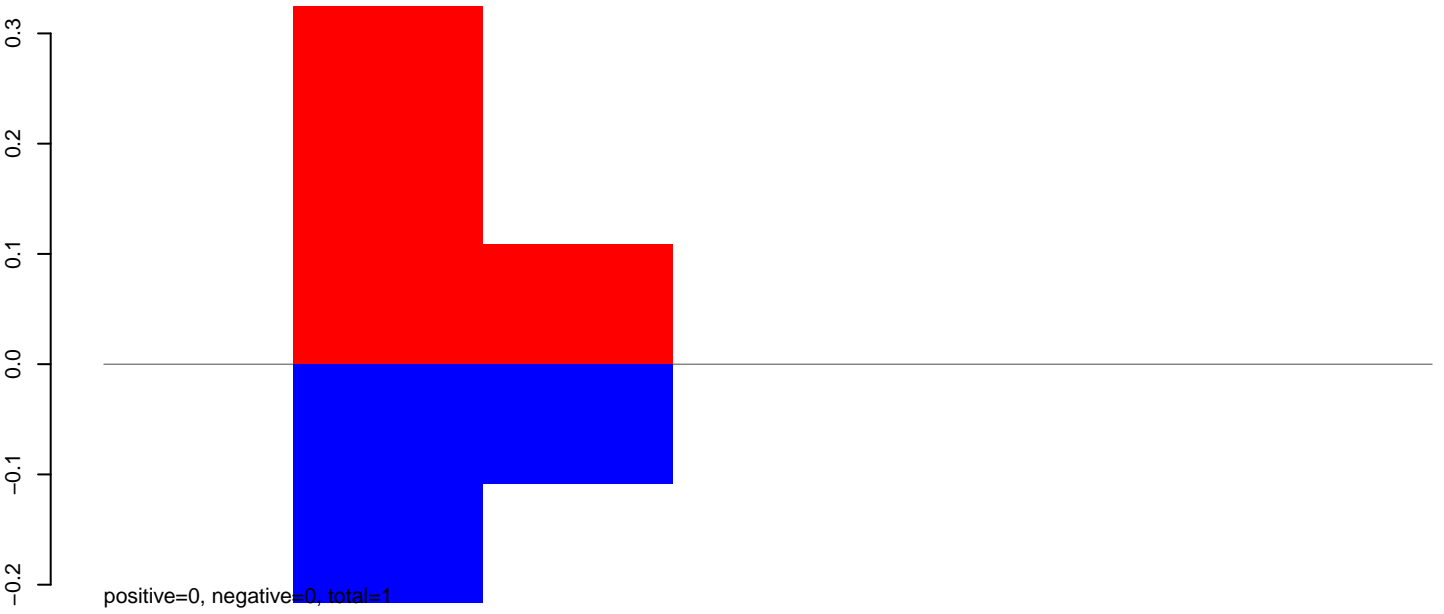
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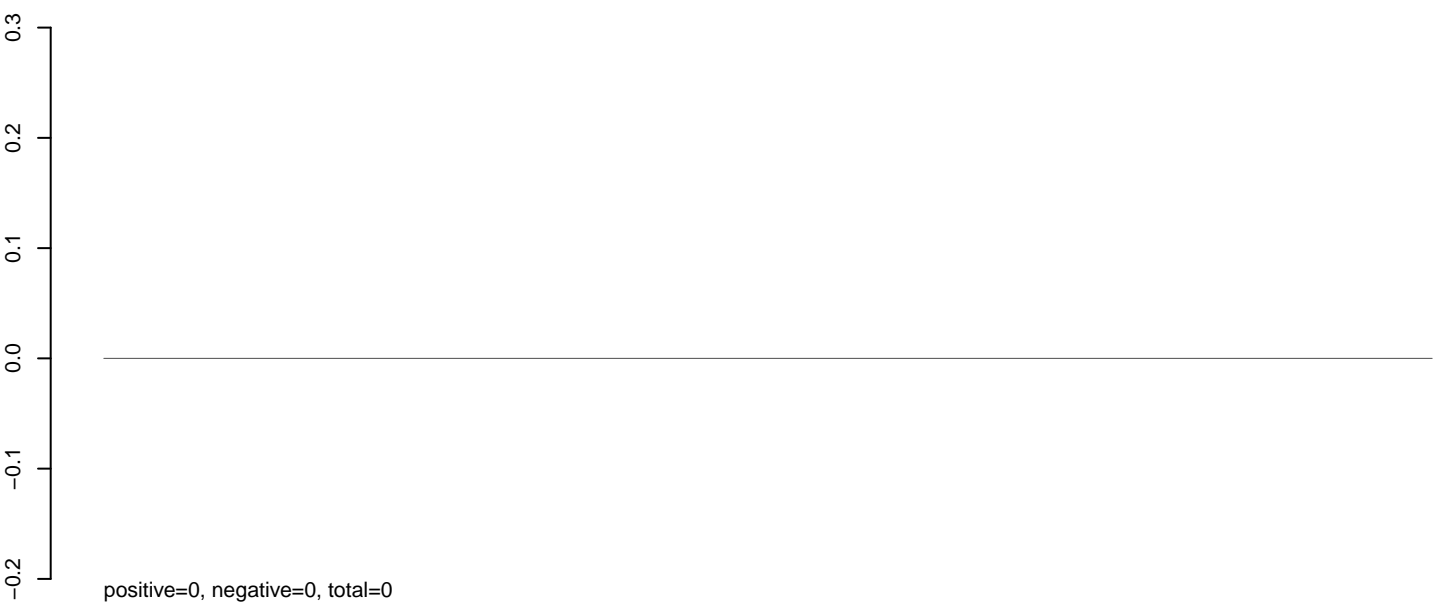
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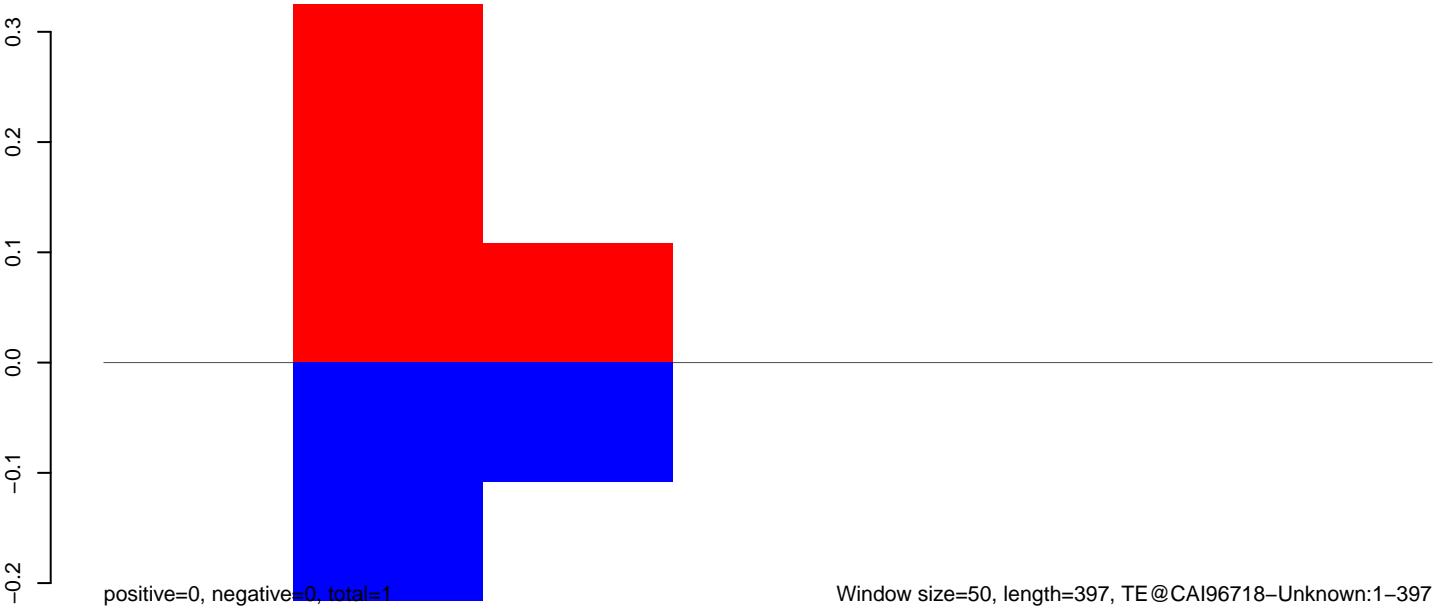
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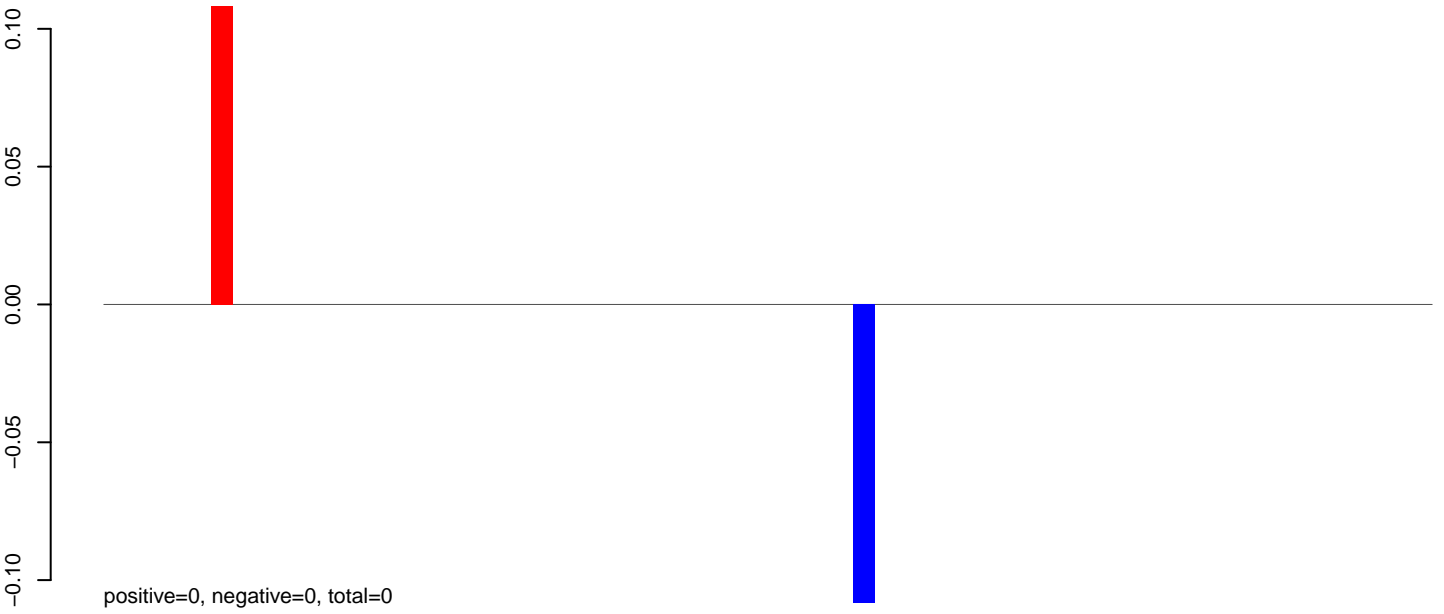
AnGam_Sua5b_cells.24_35.rep



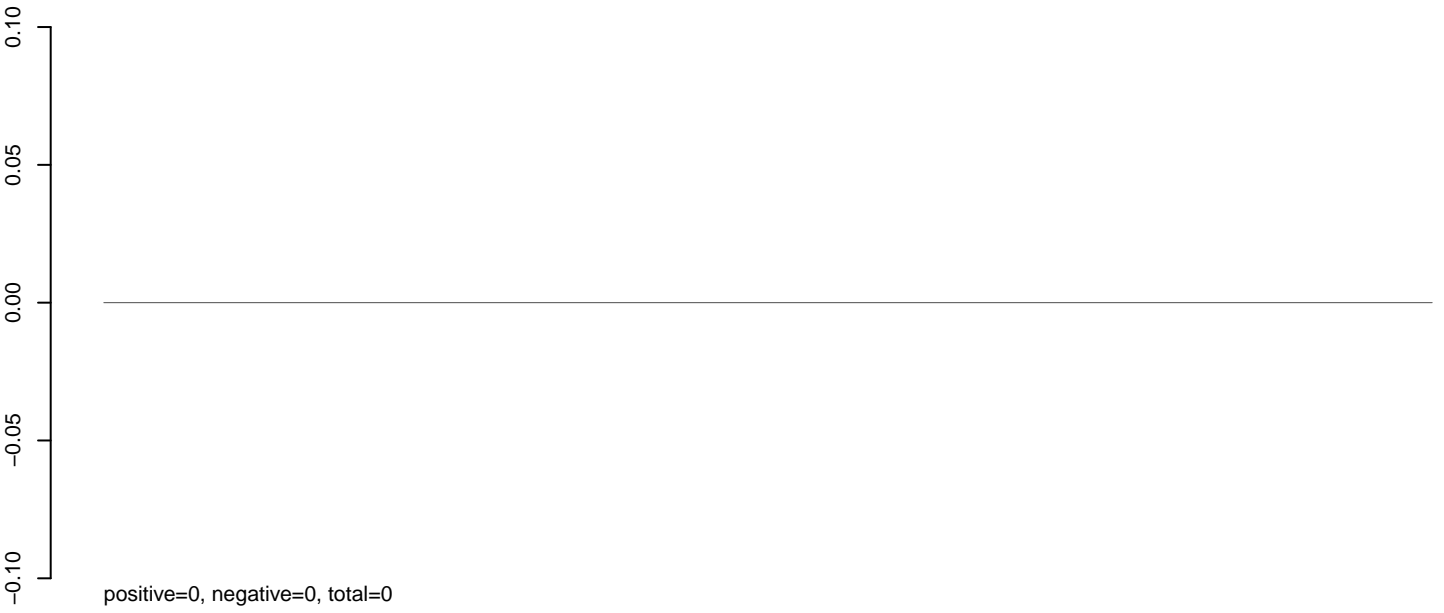
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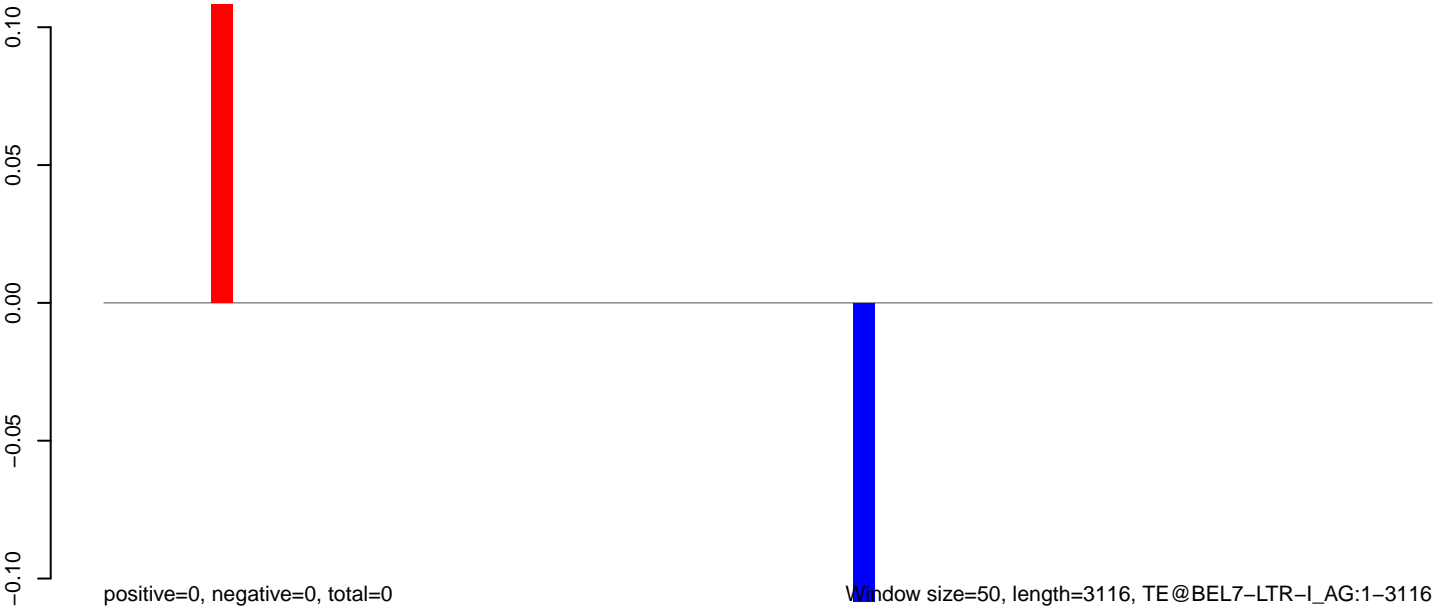
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AnGam_Sua5b_cells.24_35.rep

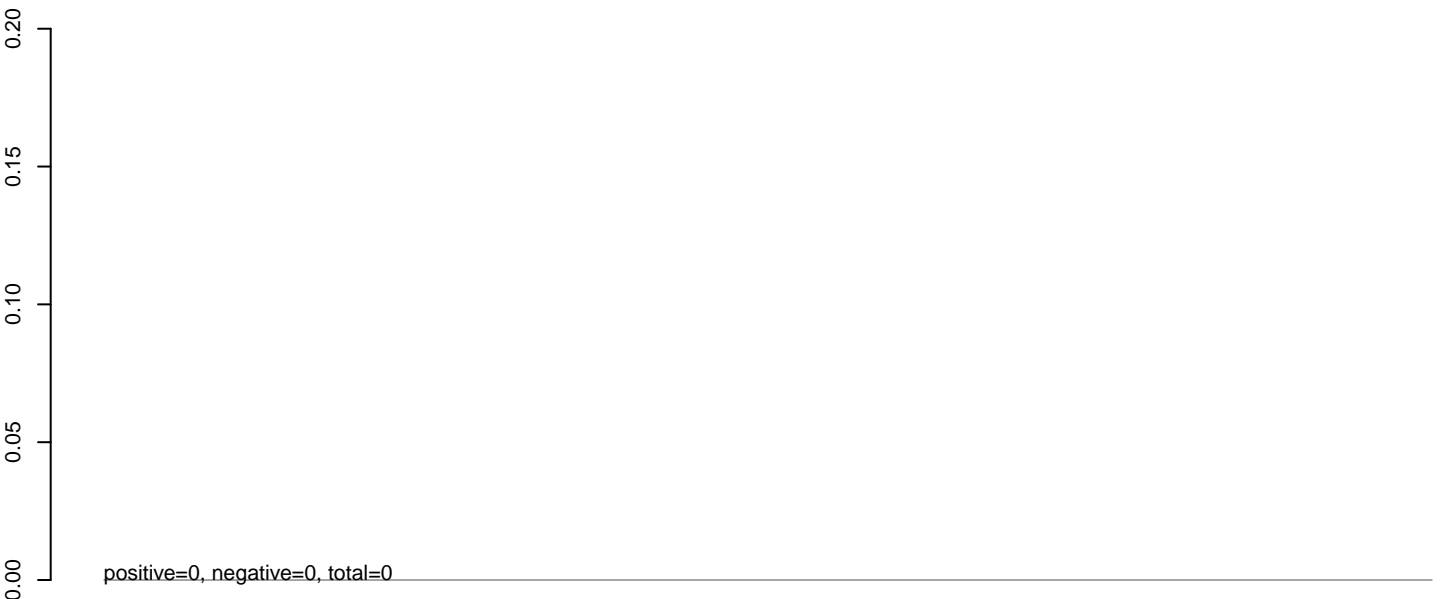


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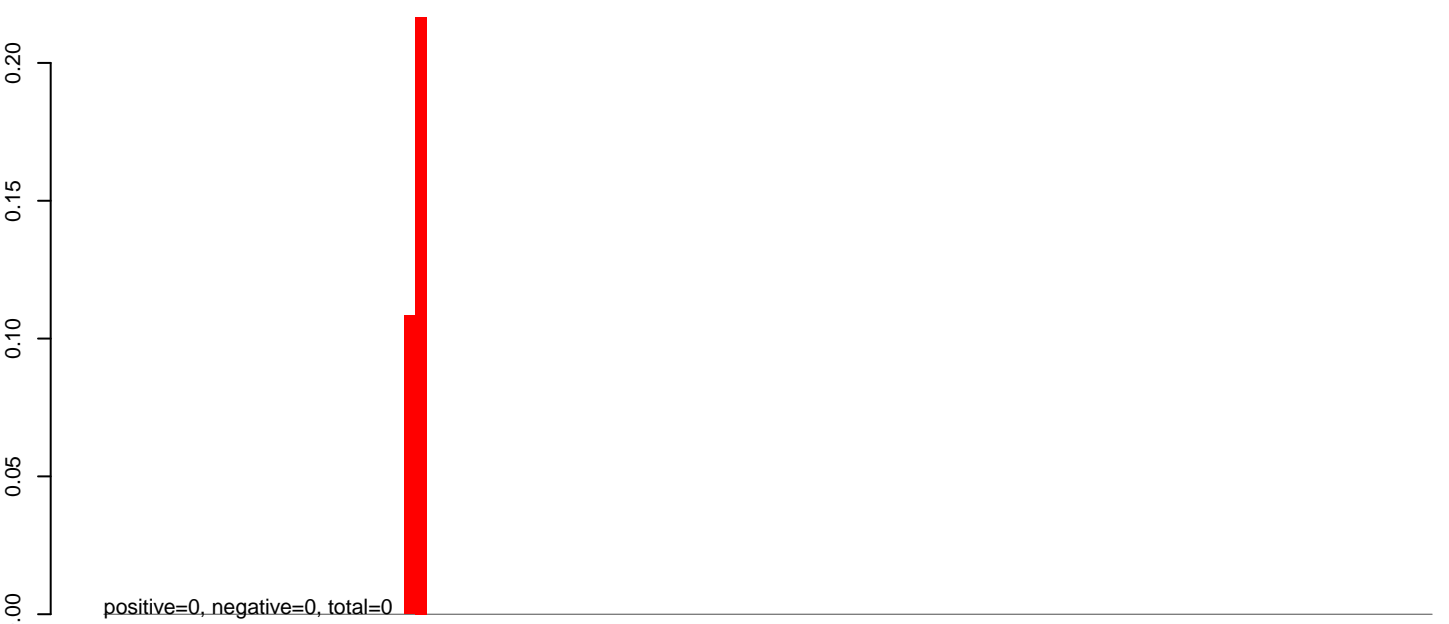


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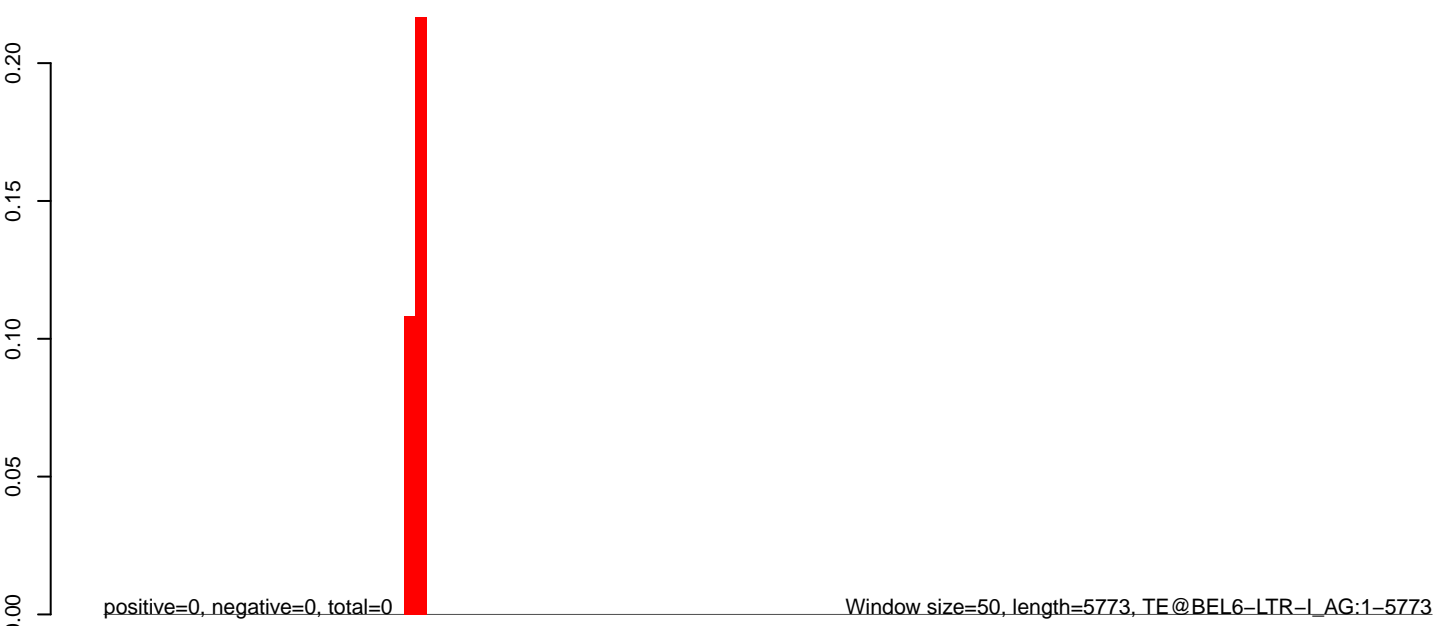
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AnGam_Sua5b_cells.24_35.rep

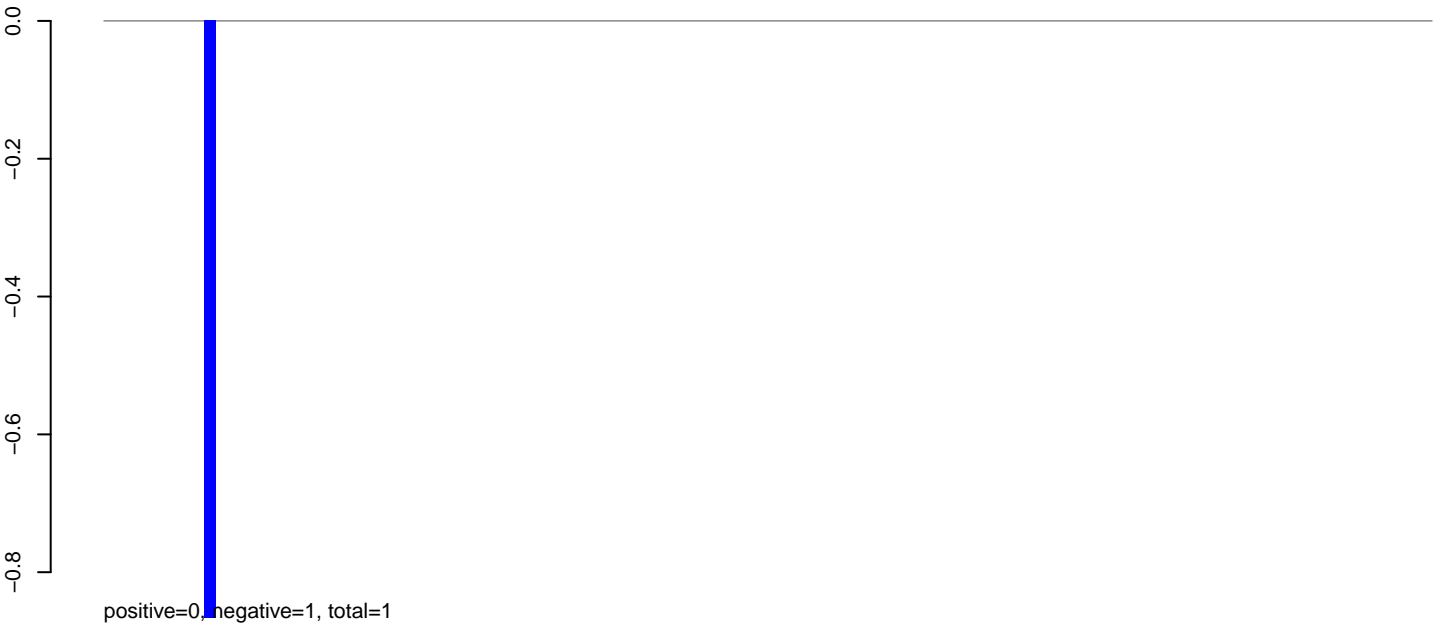


AnGam_Sua5b_cells.rep

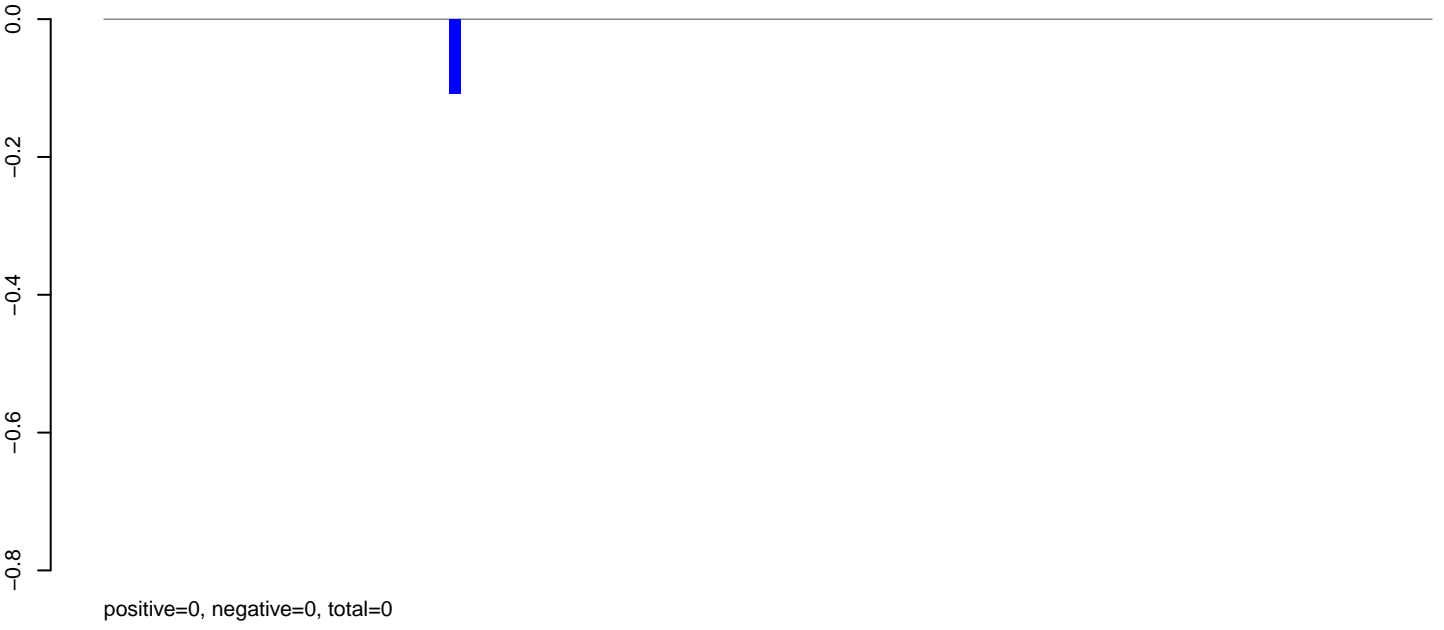


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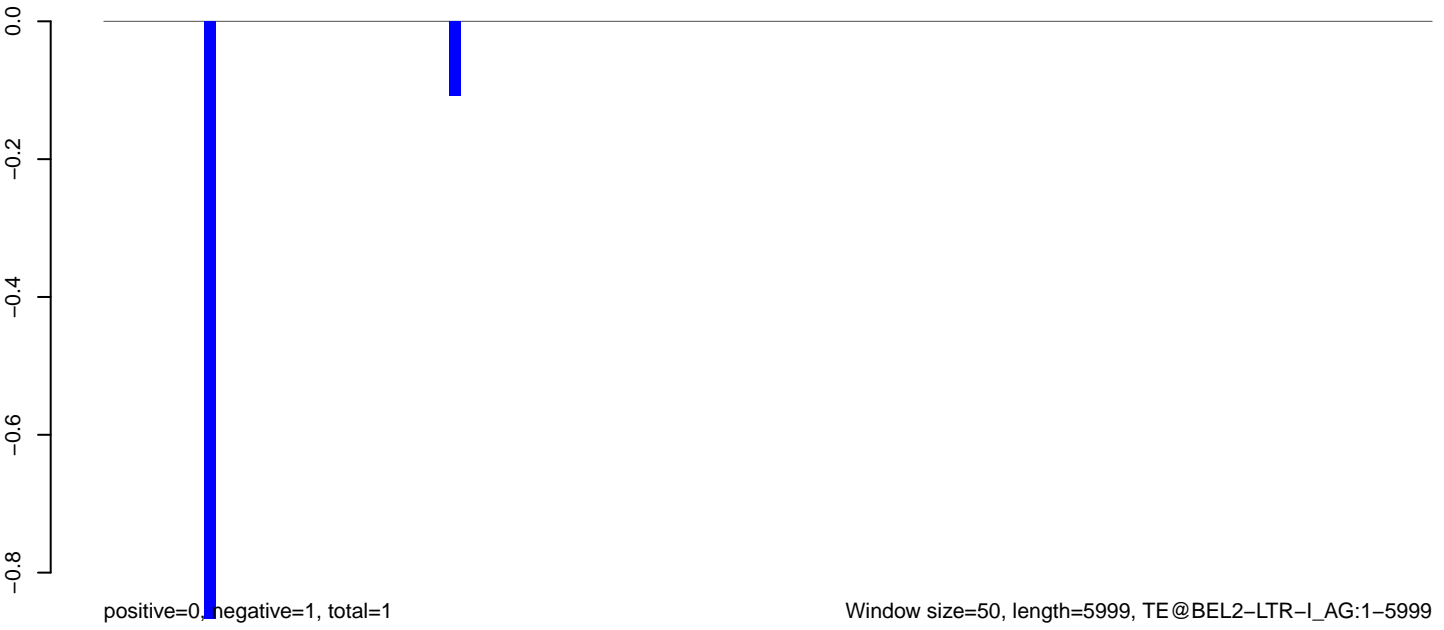
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AnGam_Sua5b_cells.24_35.rep

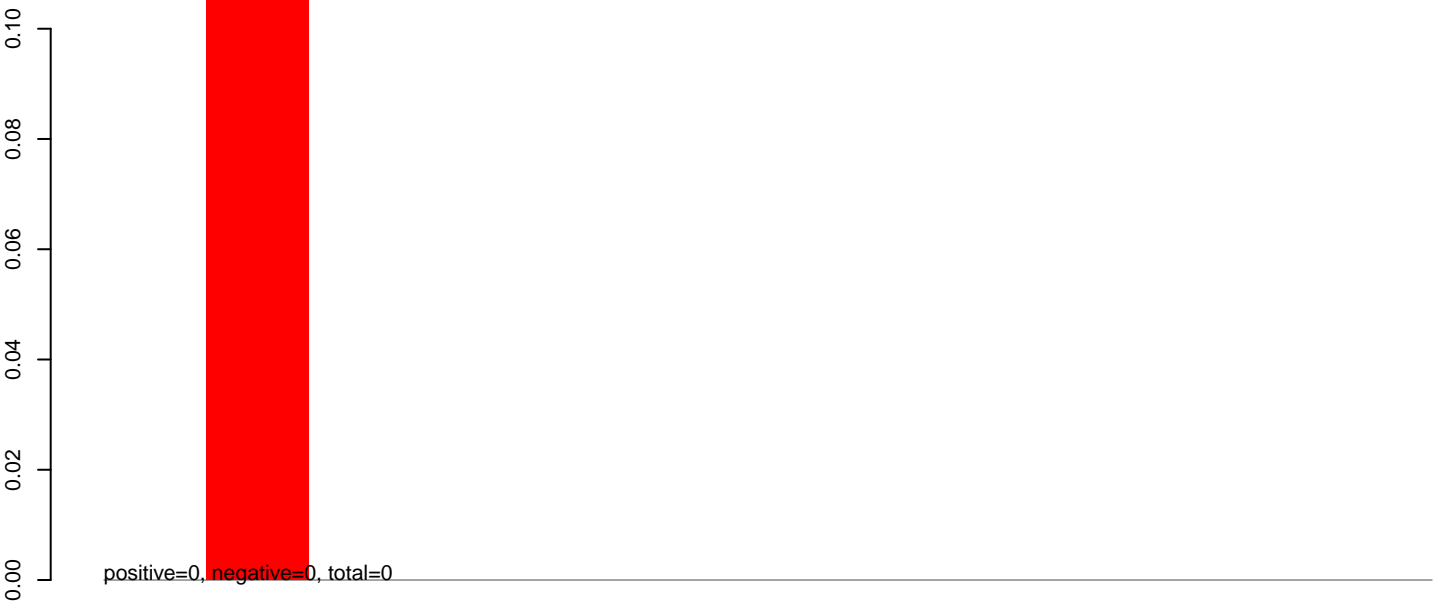


AnGam_Sua5b_cells.rep



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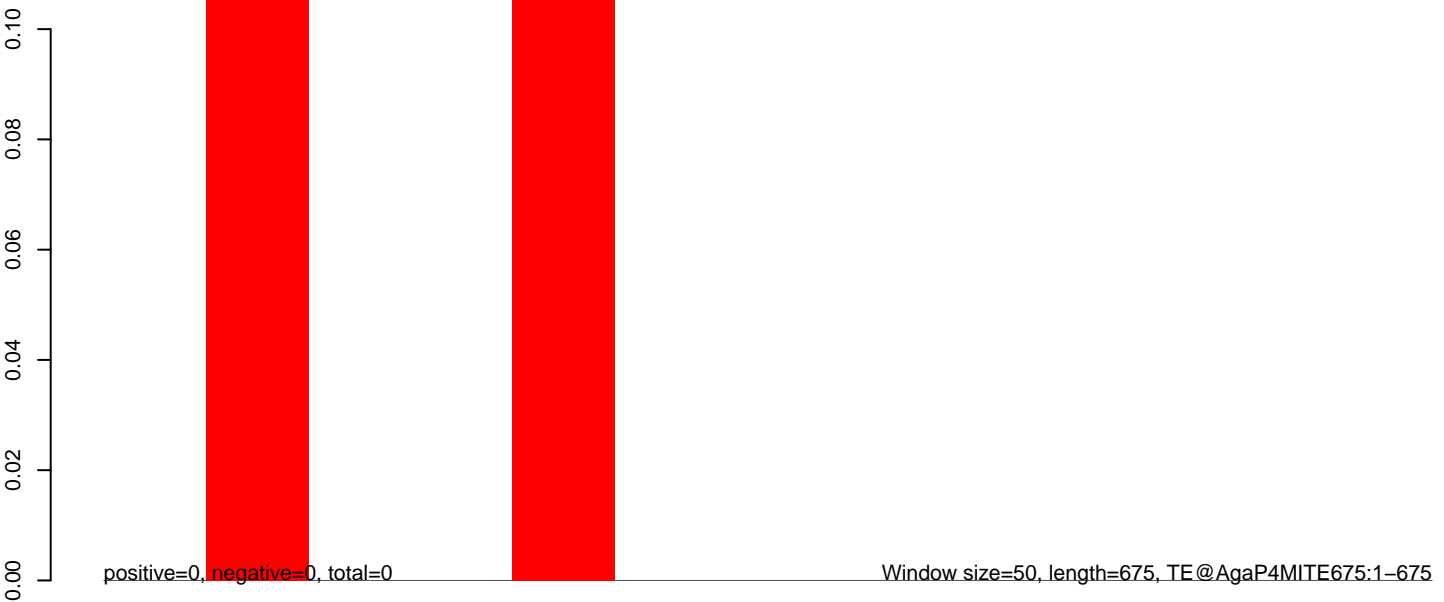
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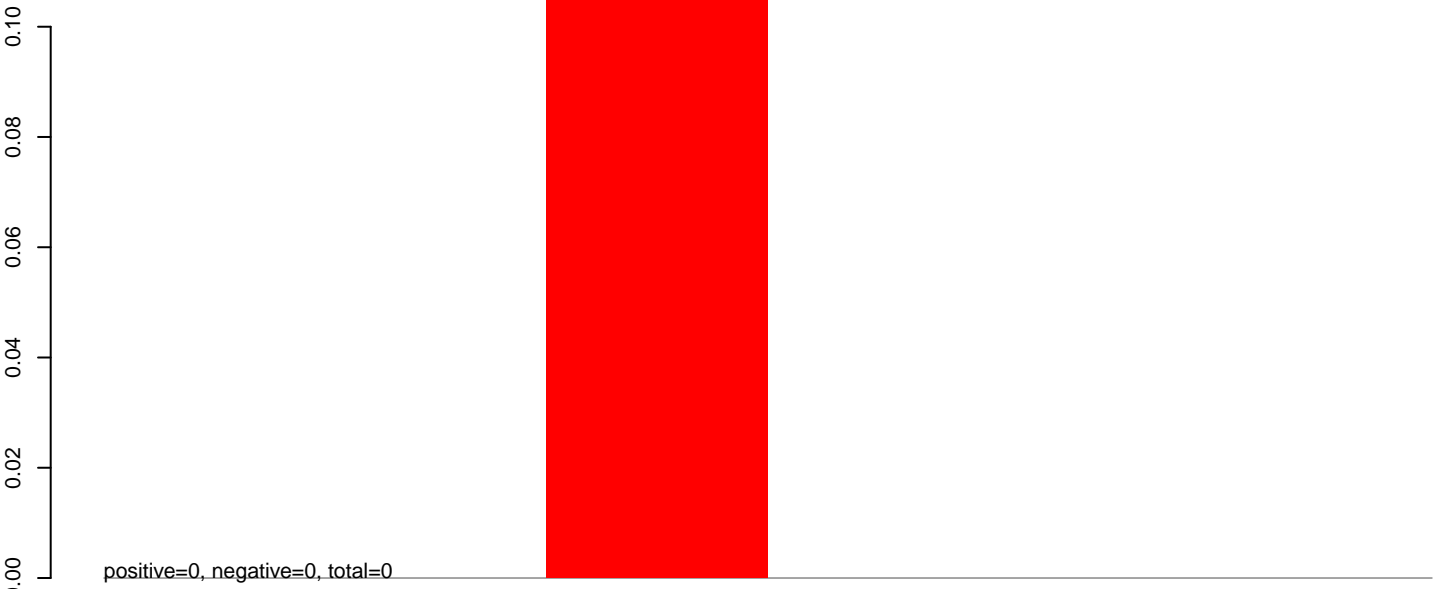
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



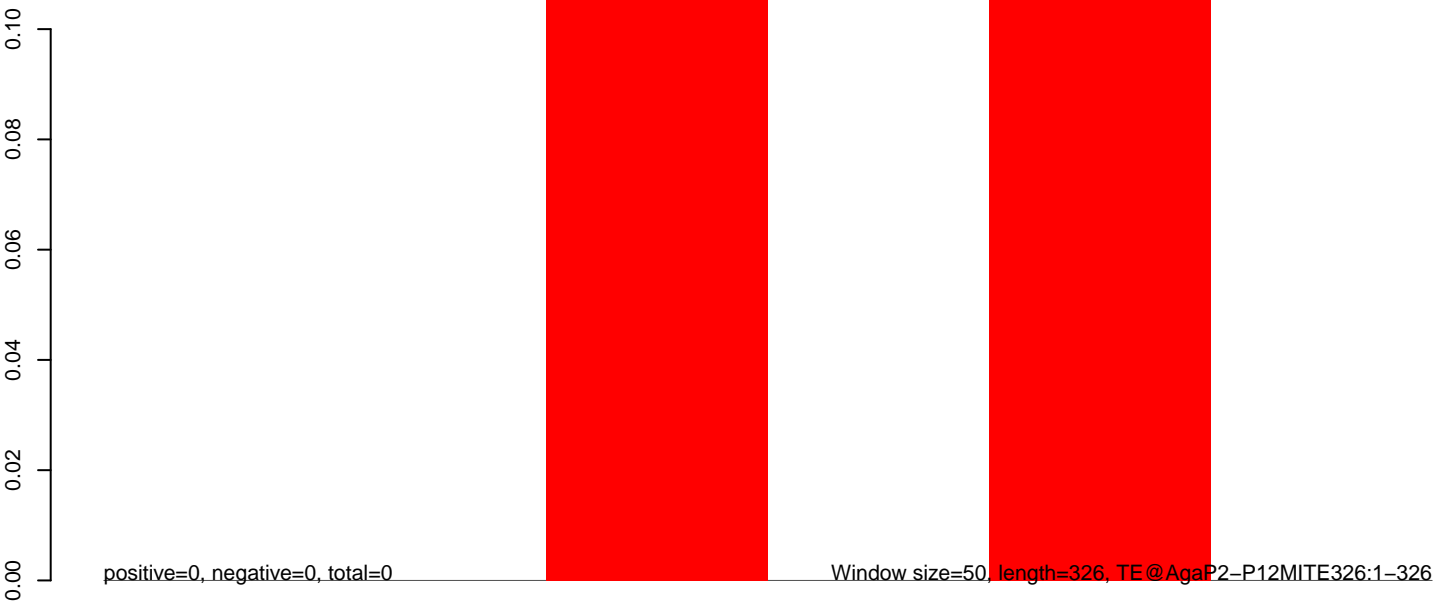
AnGam_Sua5b_cells.18_23.rep



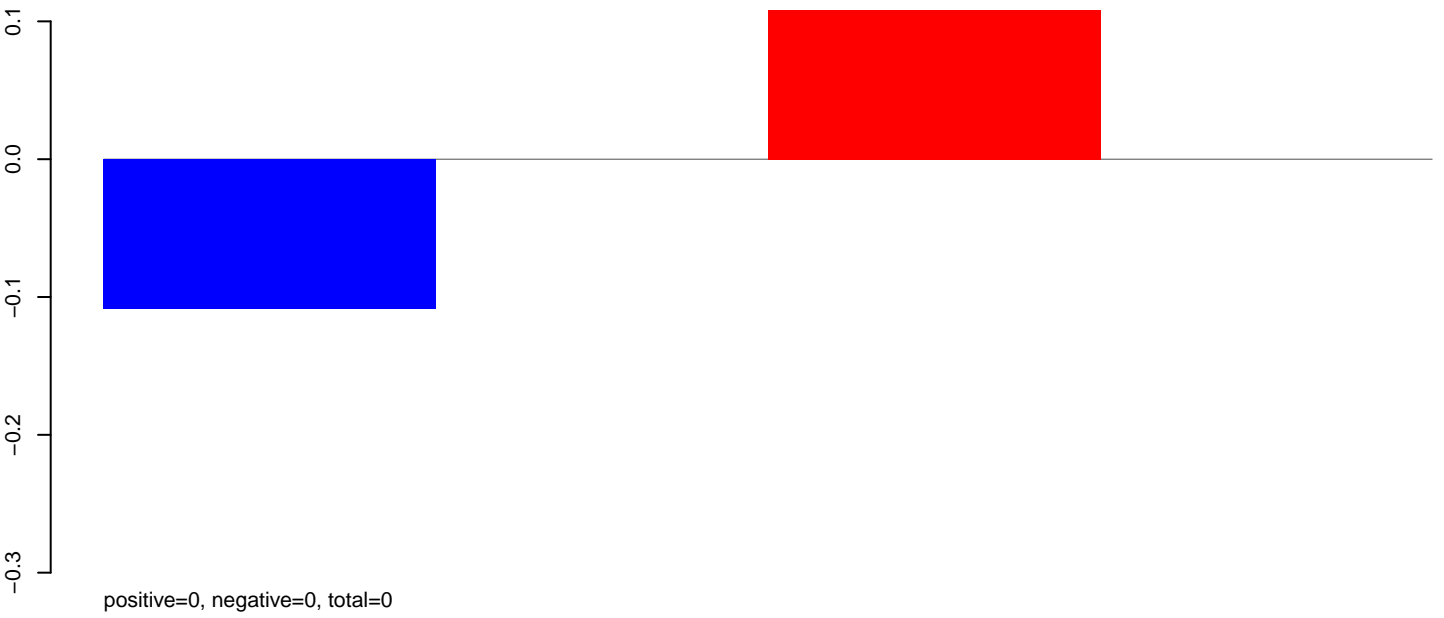
AnGam_Sua5b_cells.24_35.rep



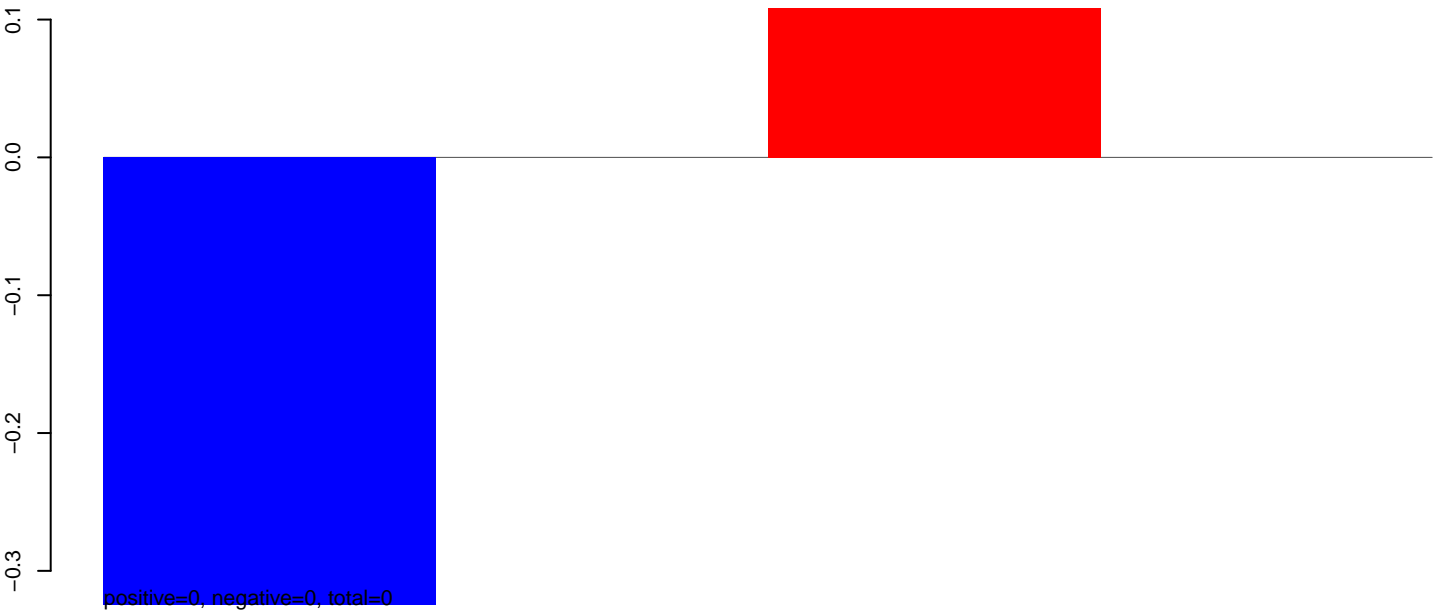
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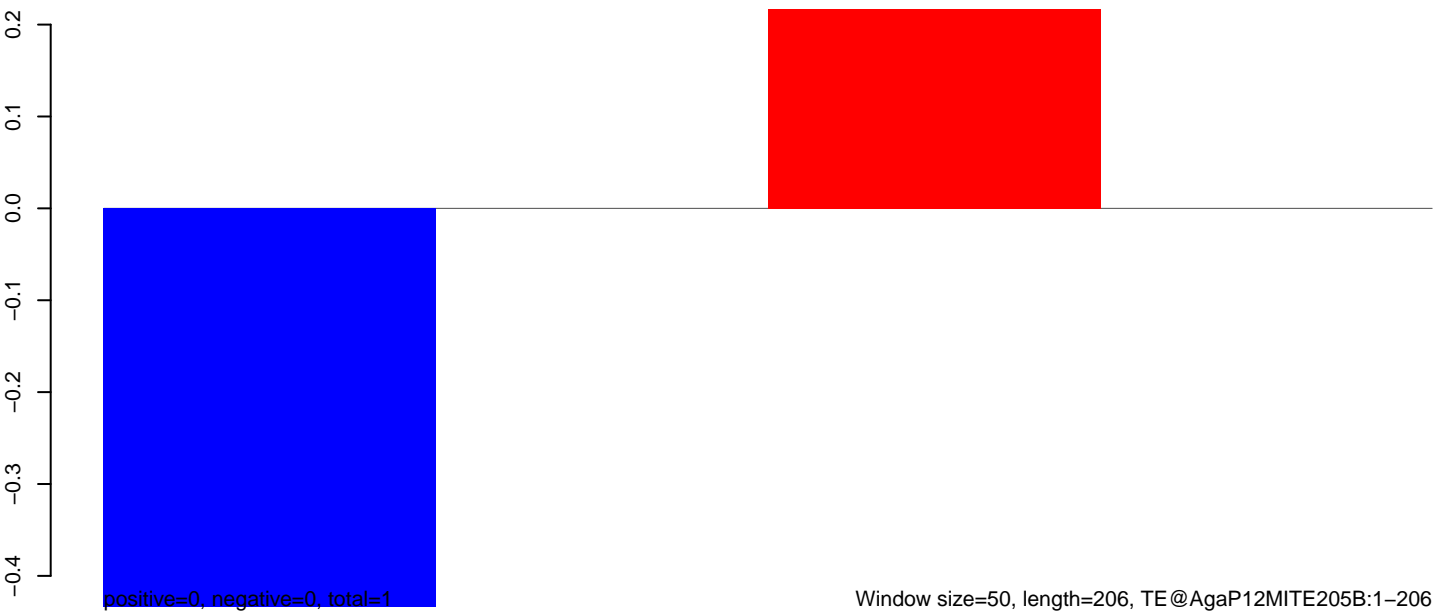
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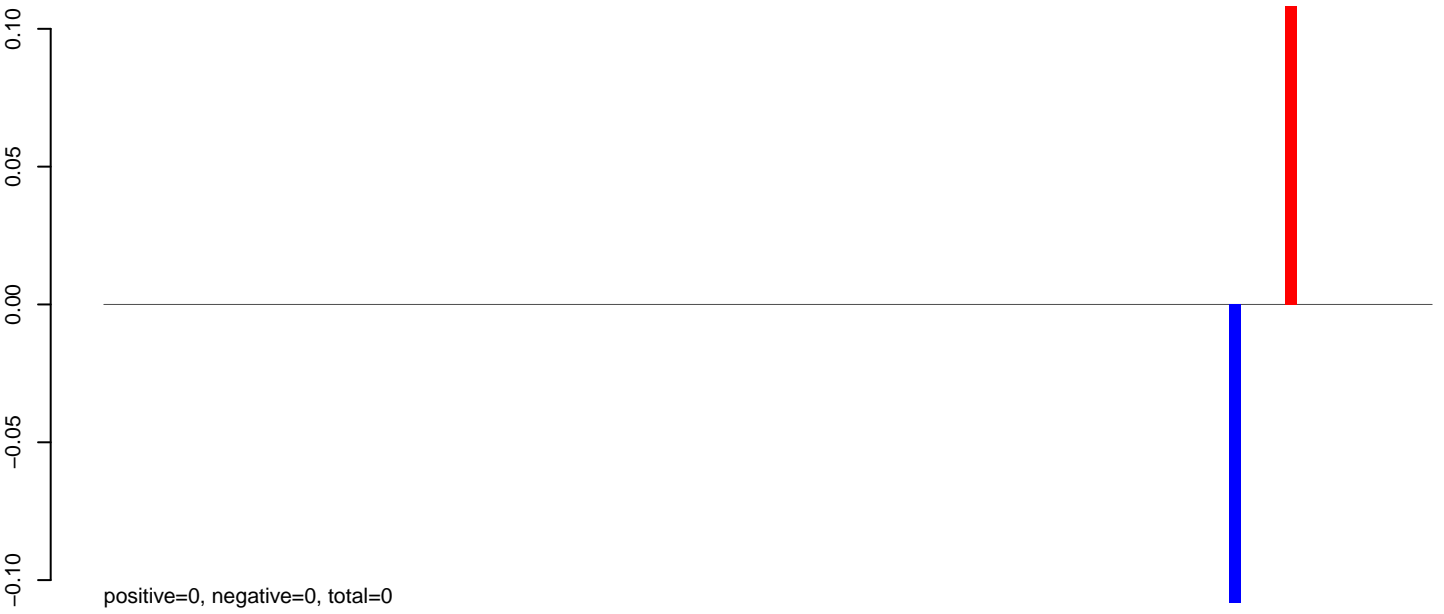
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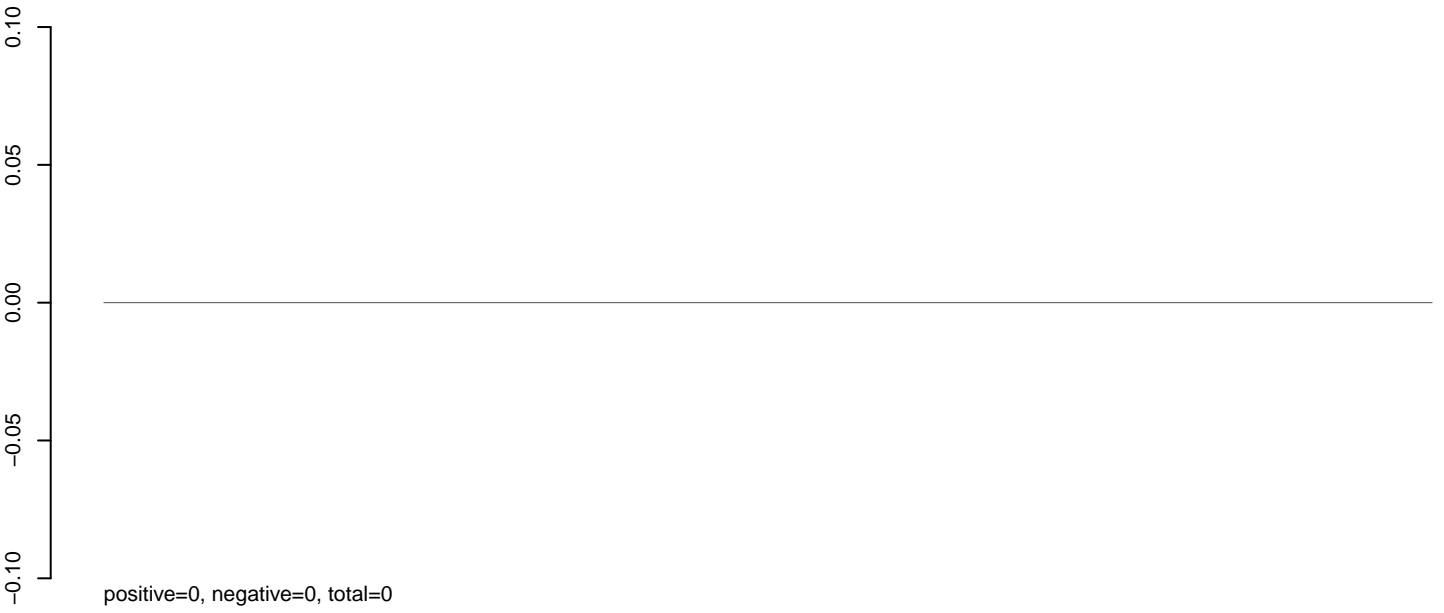
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AnGam_Sua5b_cells.18_23.rep



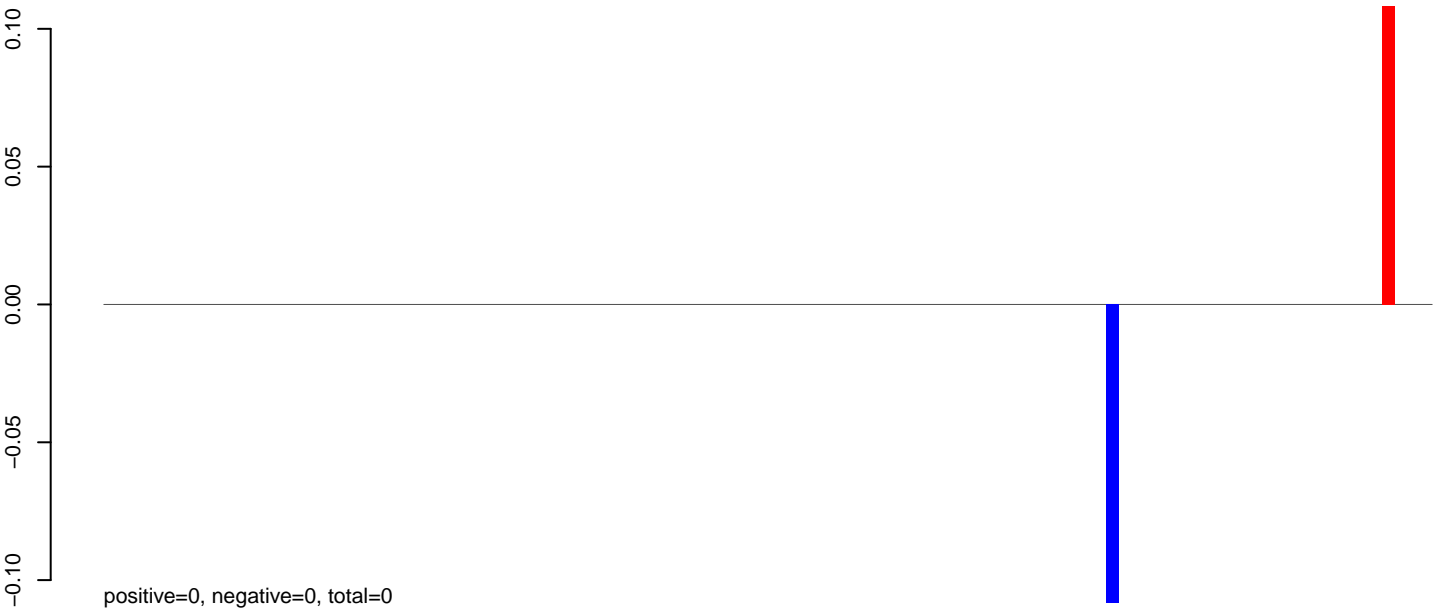
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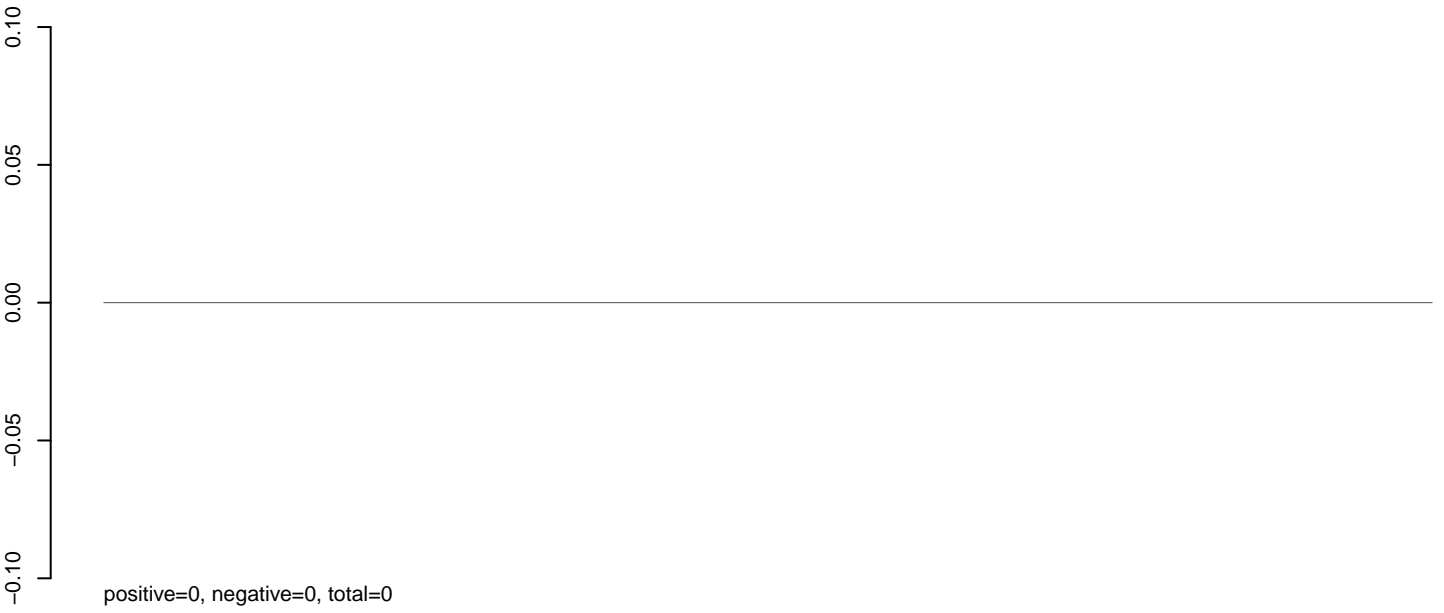
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AnGam_Sua5b_cells.18_23.rep



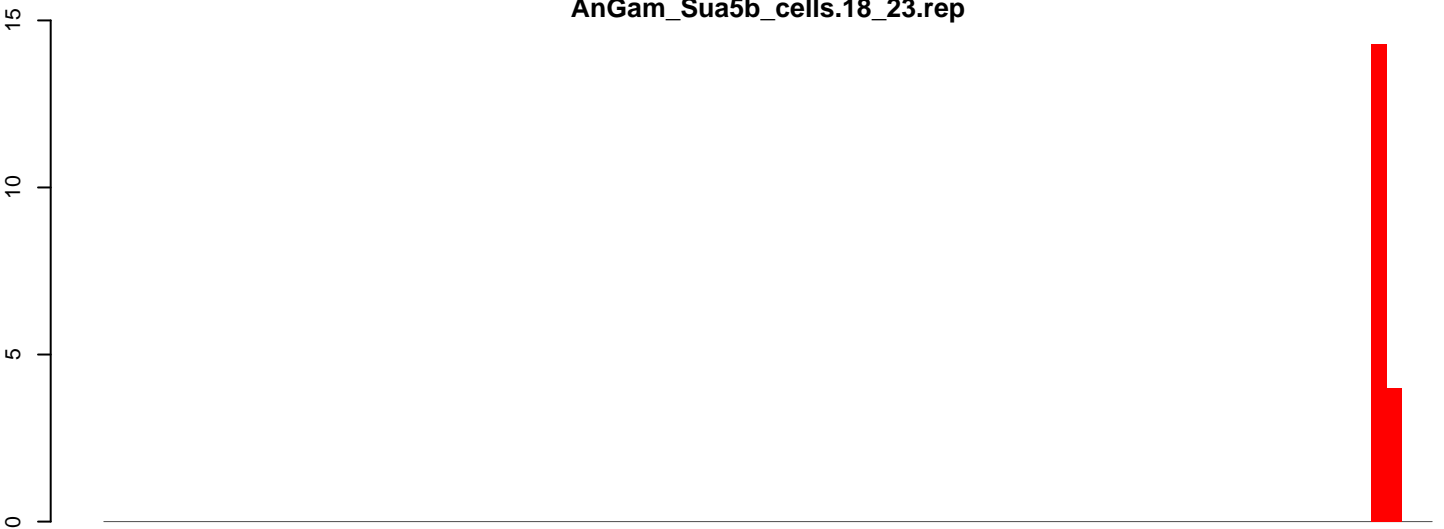
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AnGam_Sua5b_cells.rep

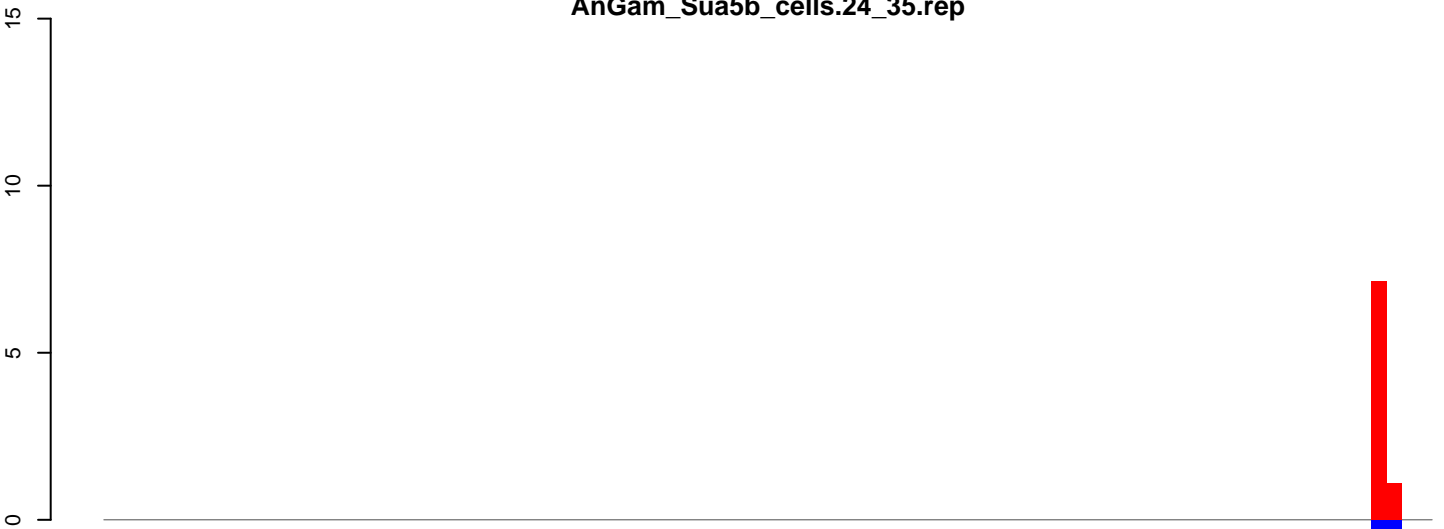


AnGam_Sua5b_cells.18_23.rep



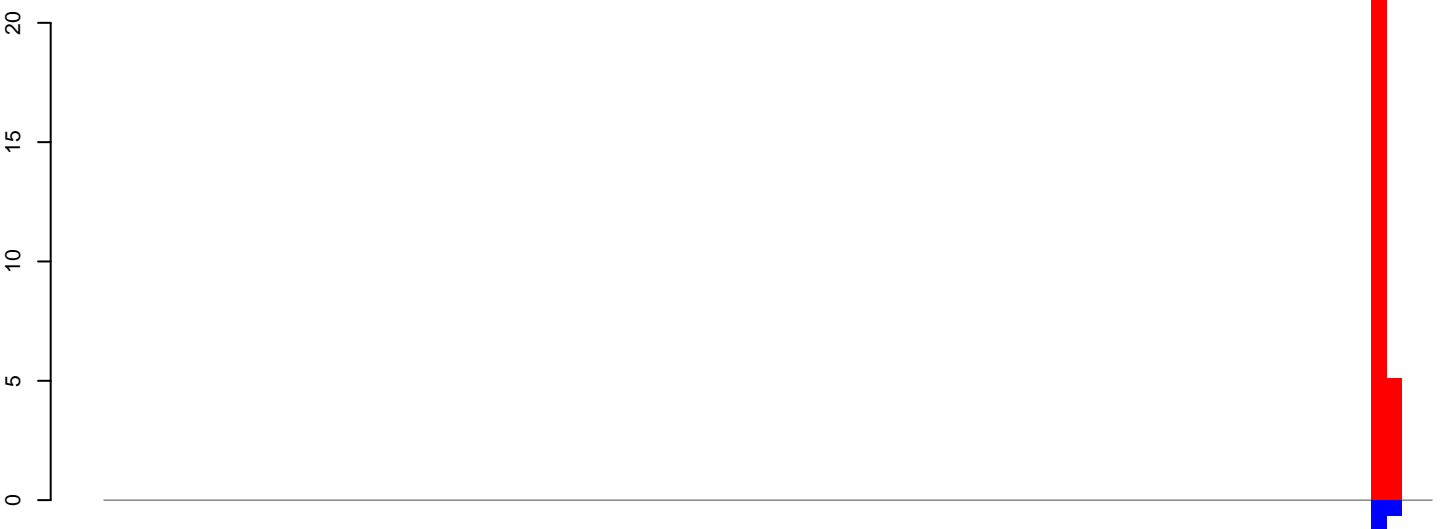
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AnGam_Sua5b_cells.24_35.rep



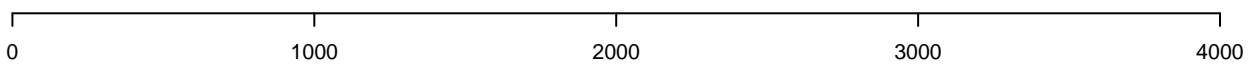
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AnGam_Sua5b_cells.rep

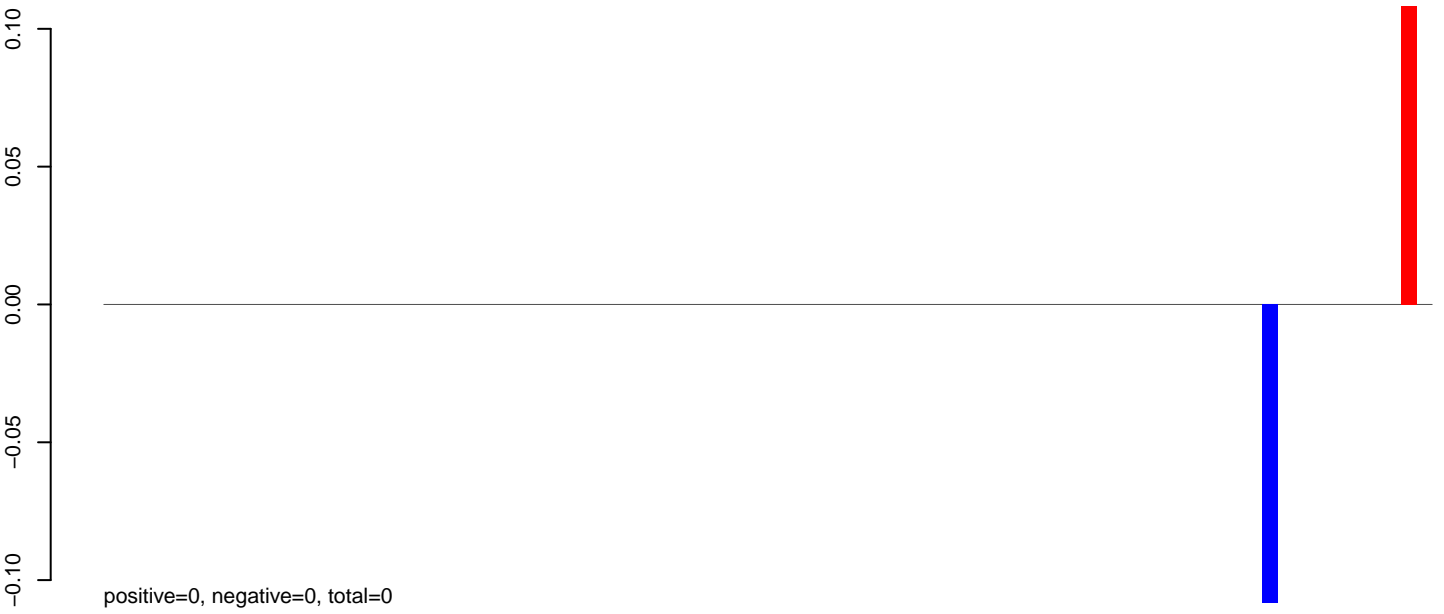


positive=27, negative=4, total=31

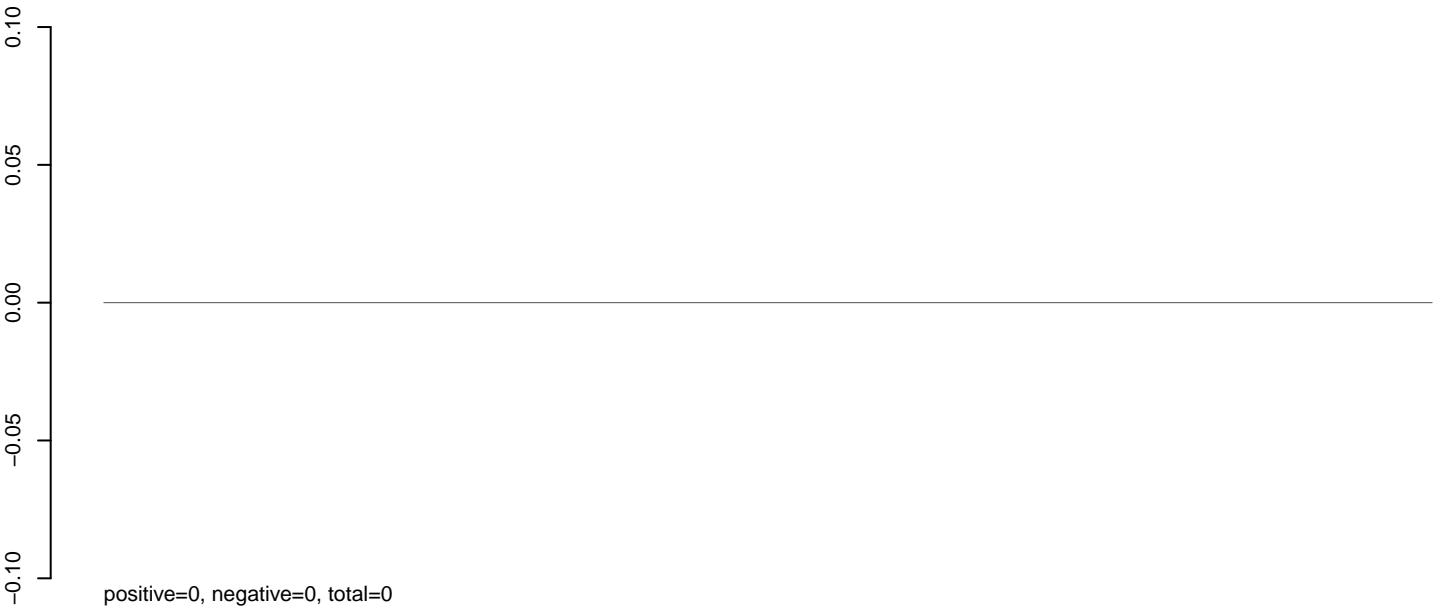
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AnGam_Sua5b_cells.18_23.rep



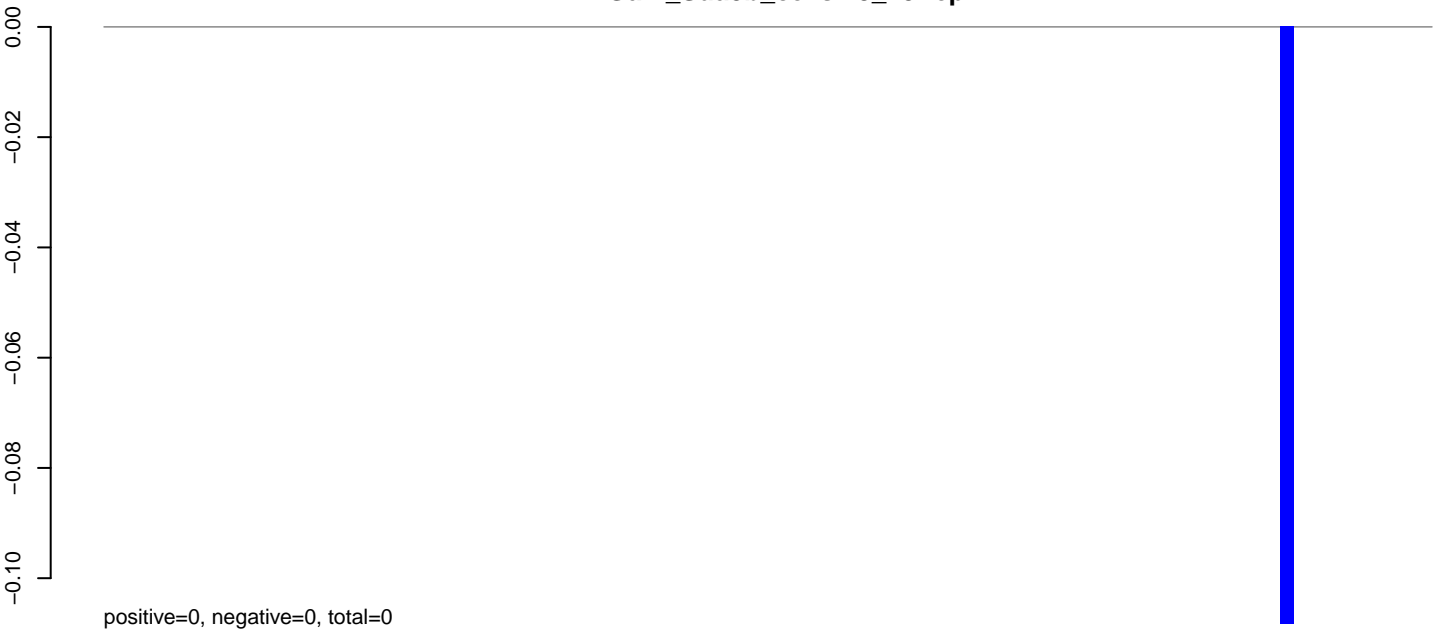
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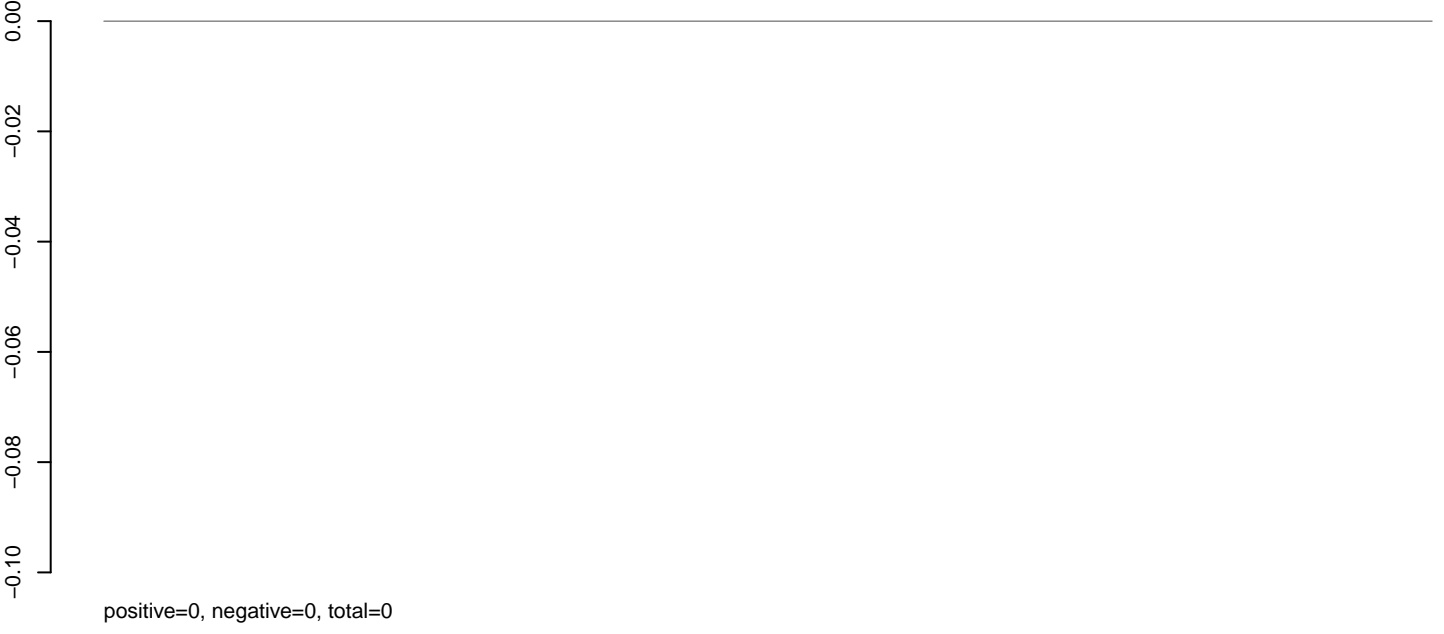
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AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



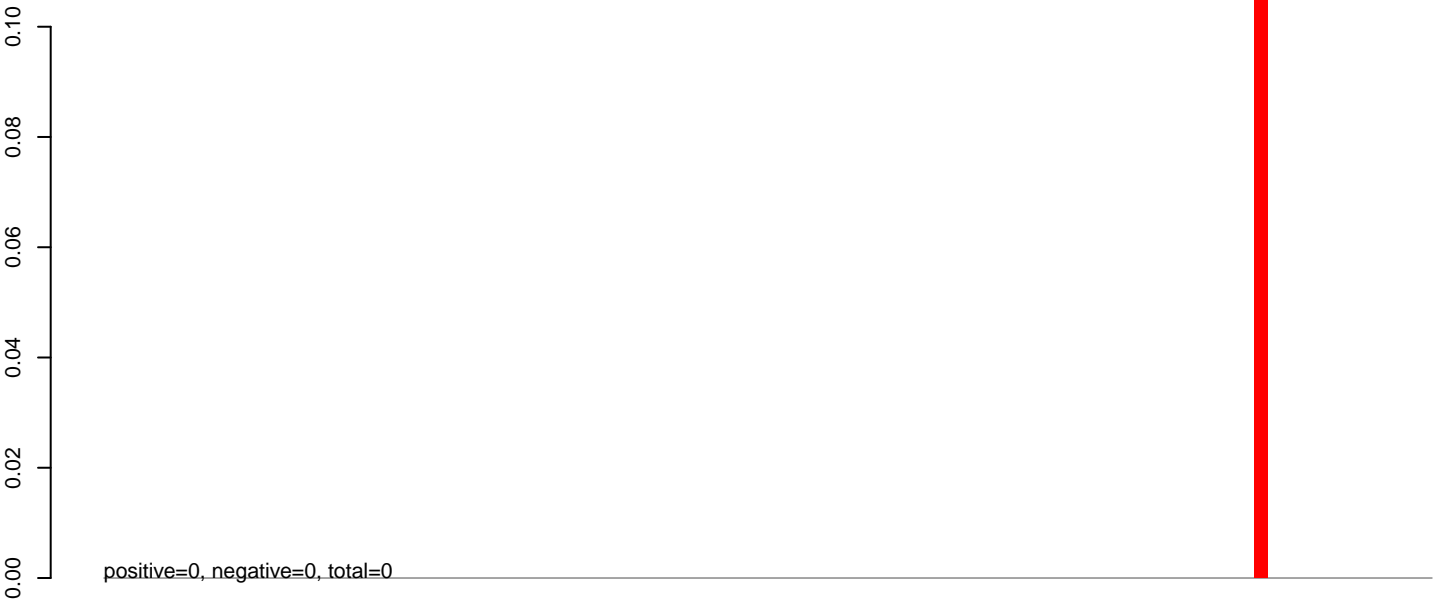
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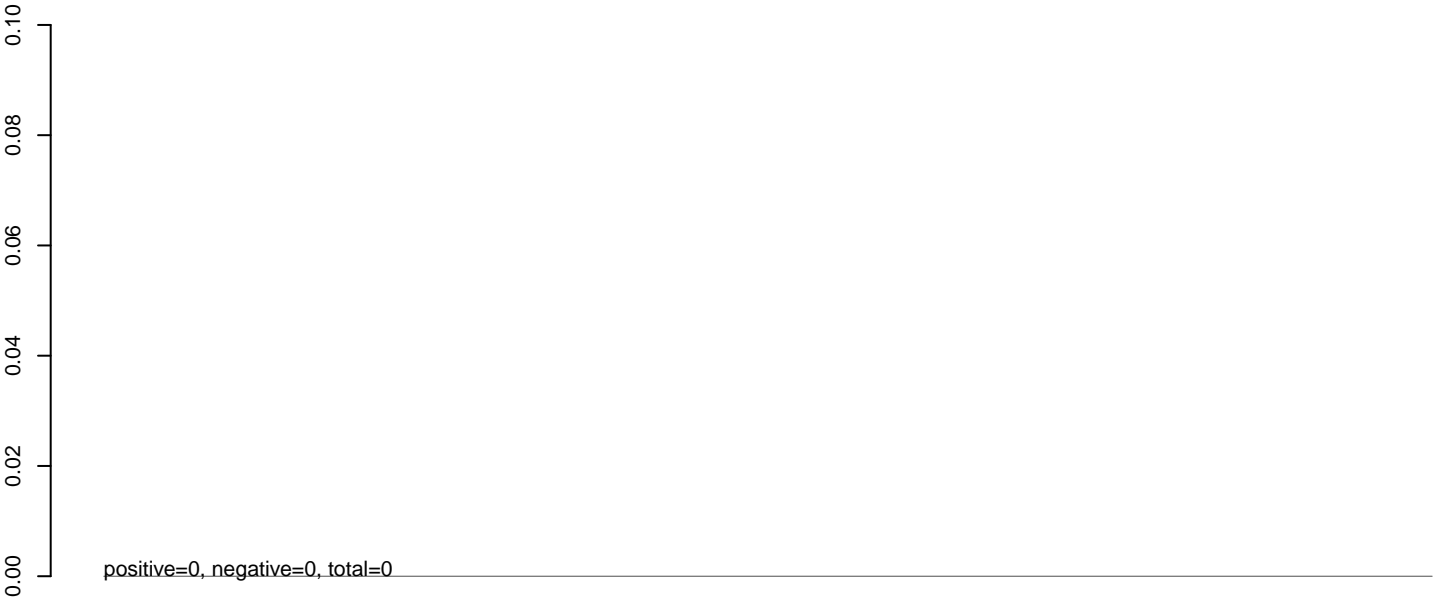
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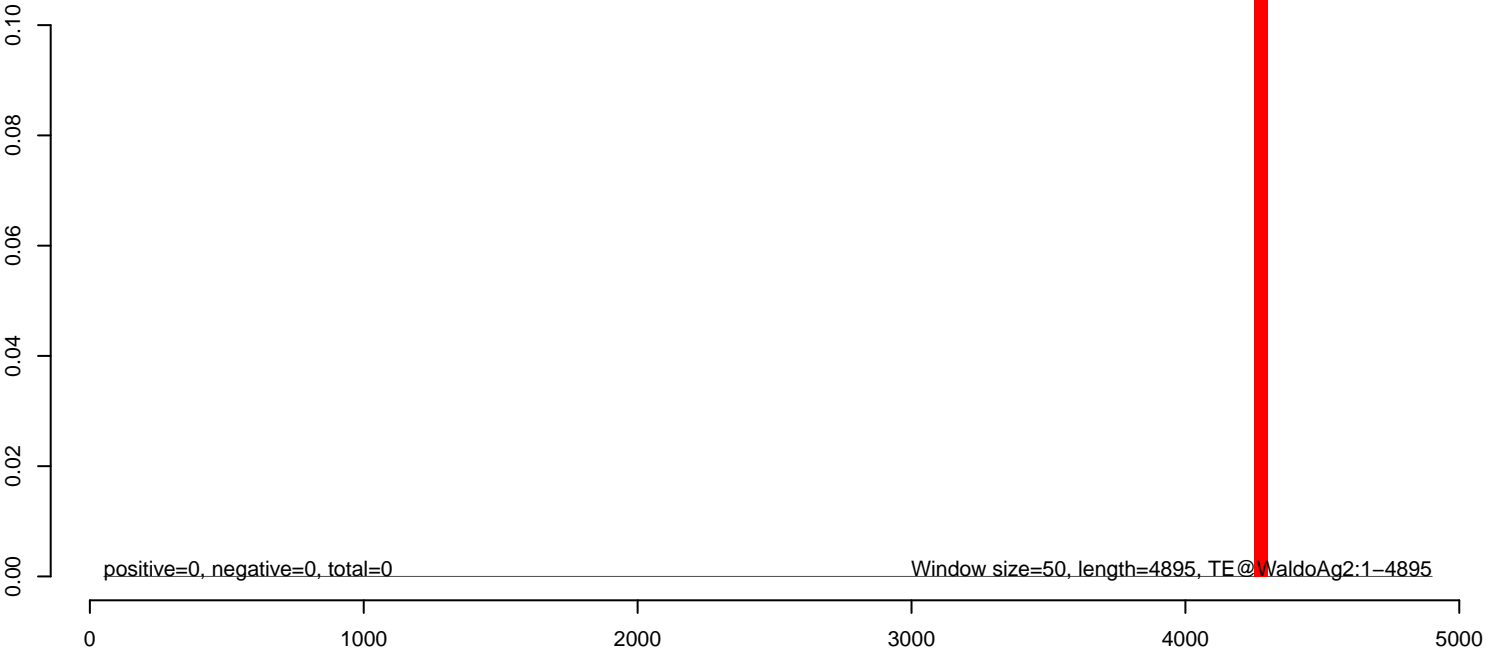
AnGam_Sua5b_cells.18_23.rep



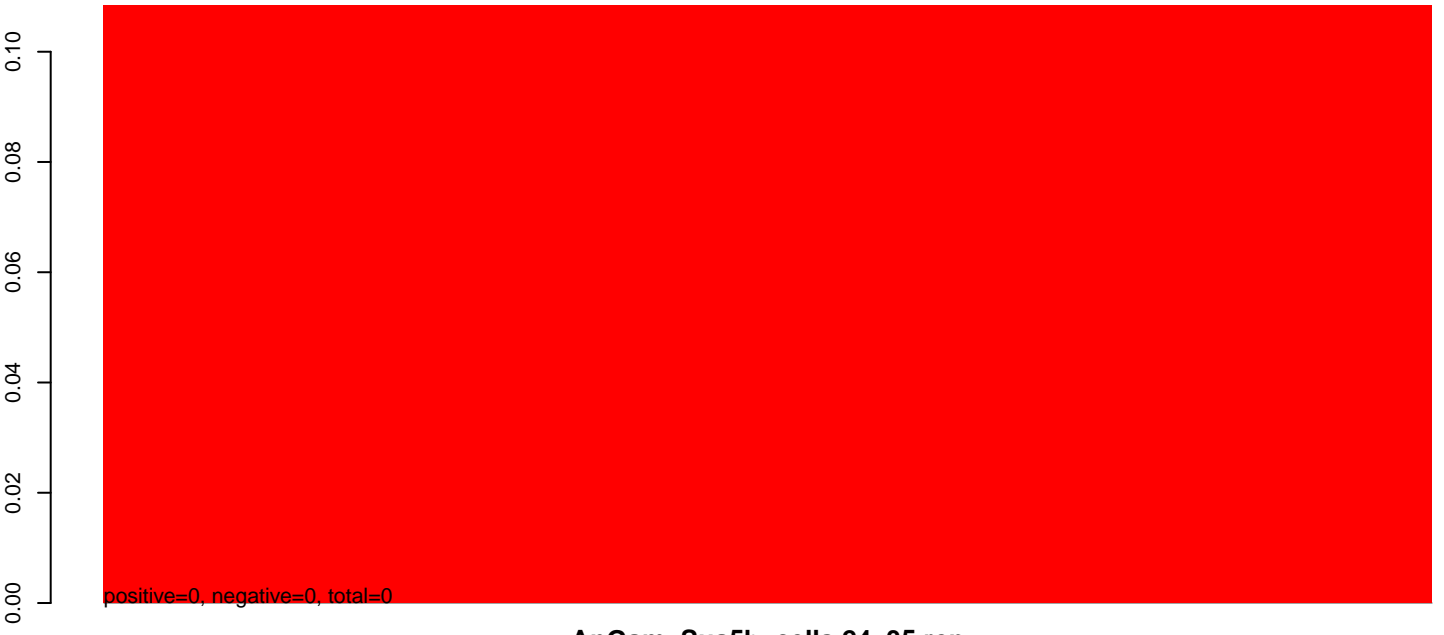
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep

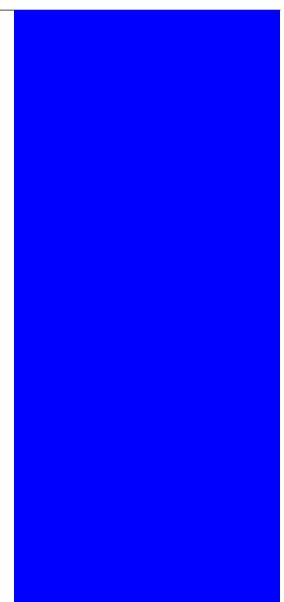
0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0



AnGam_Sua5b_cells.rep

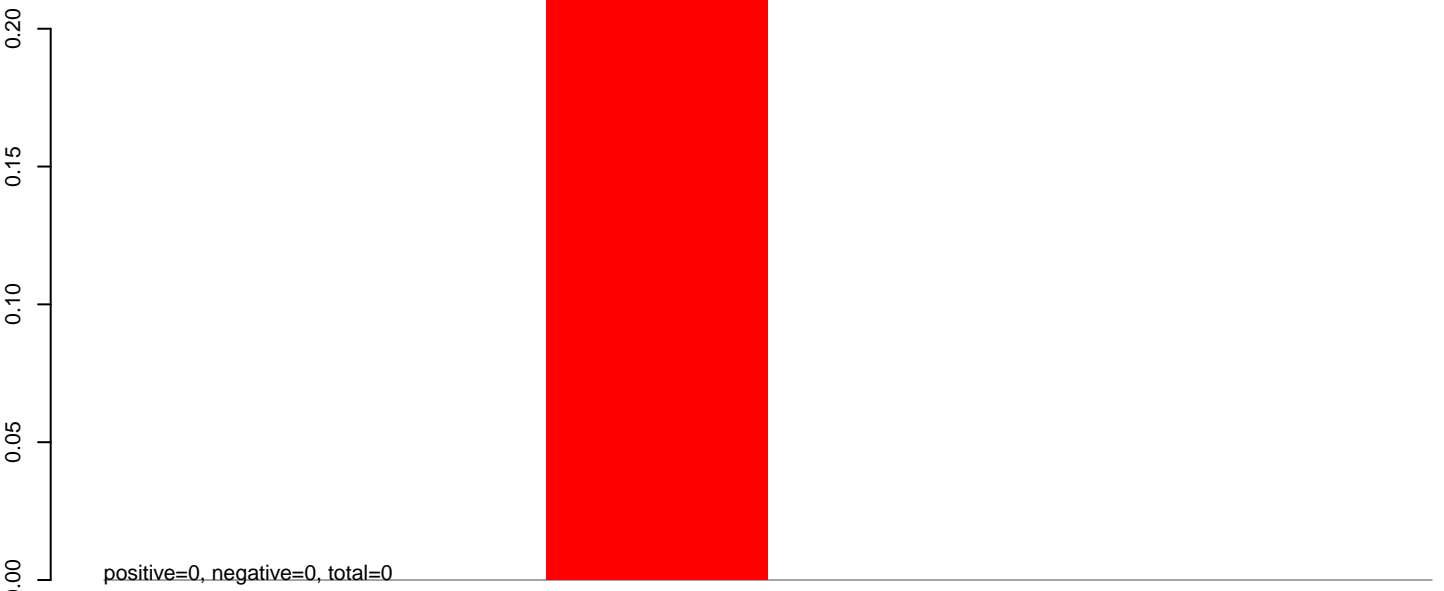
0.00
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-0.06
-0.08
-0.10

positive=0, negative=0, total=0

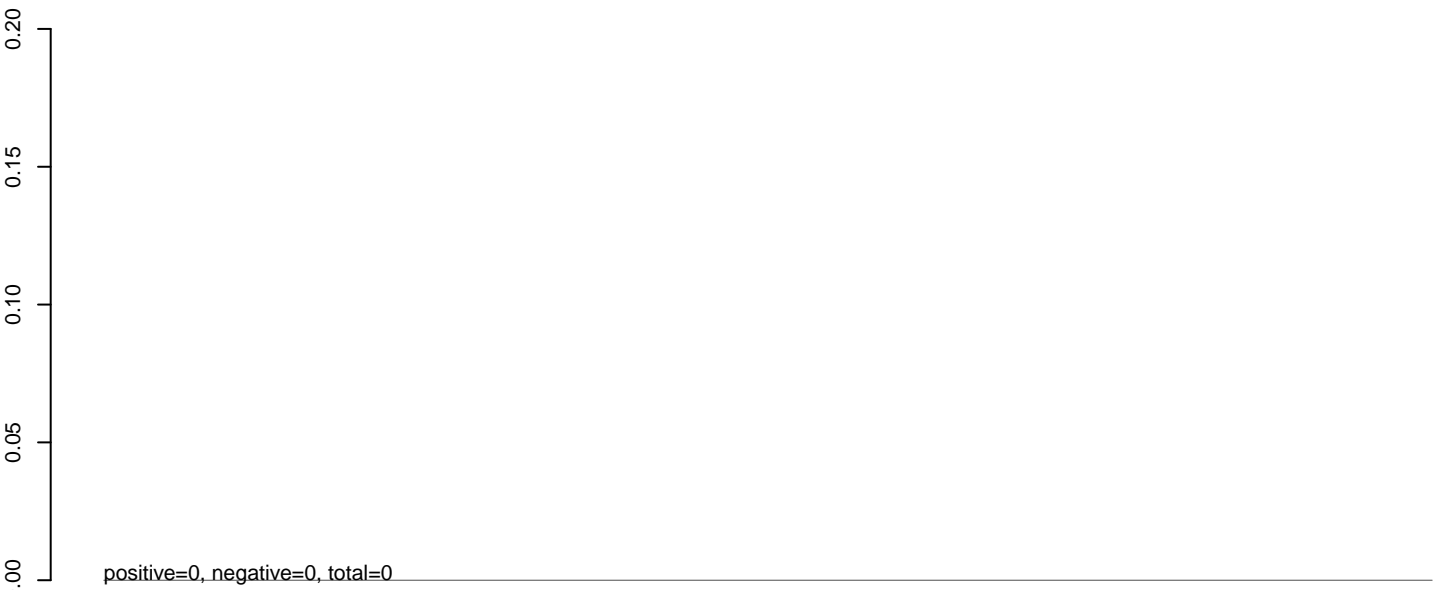
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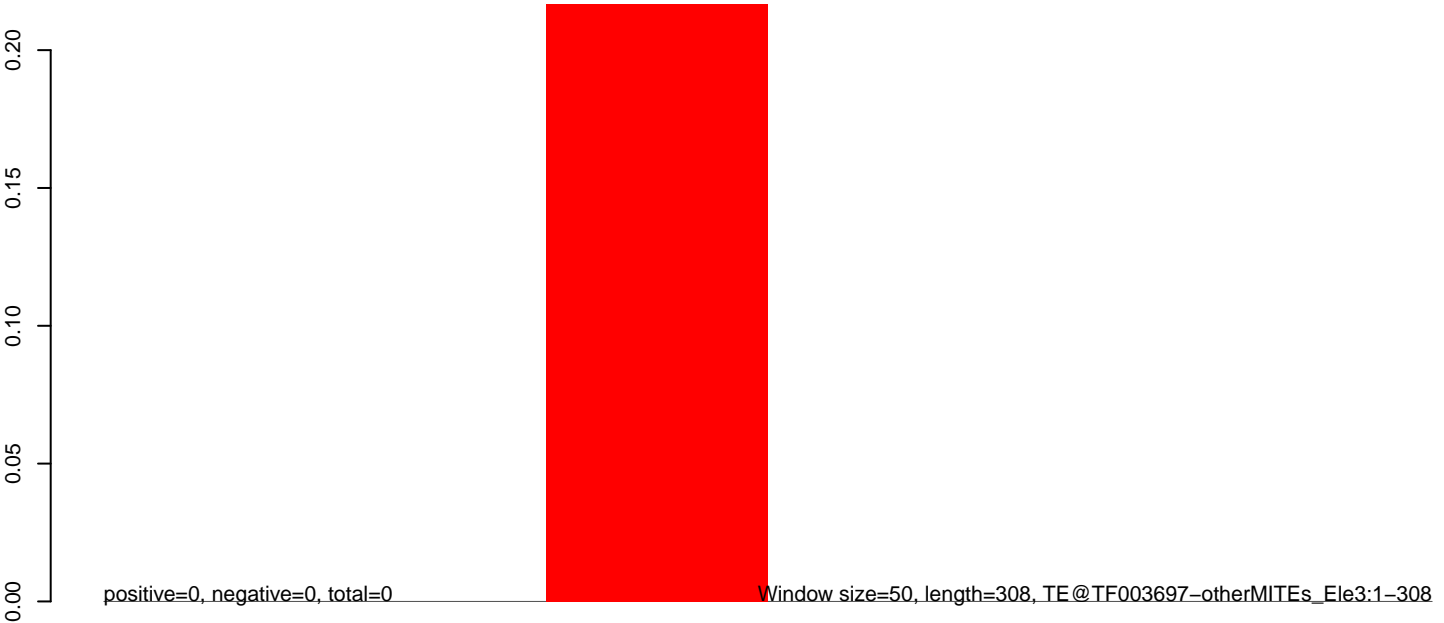
AnGam_Sua5b_cells.18_23.rep



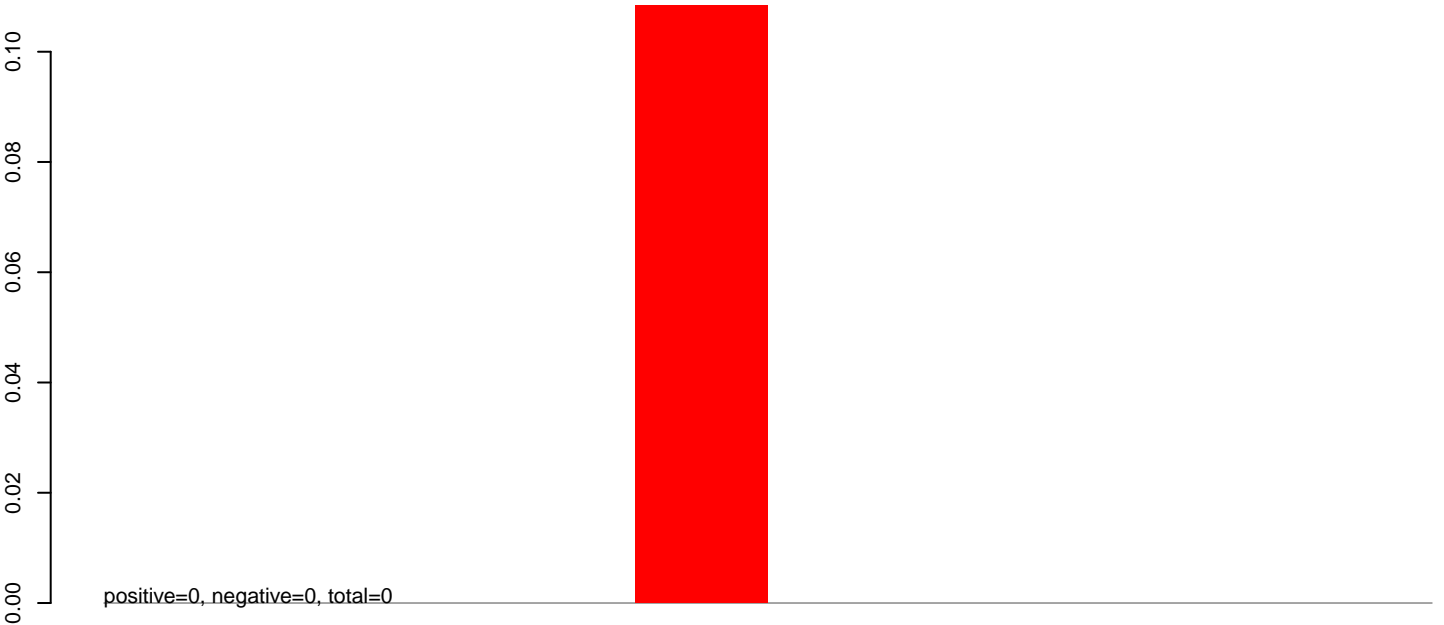
AnGam_Sua5b_cells.24_35.rep



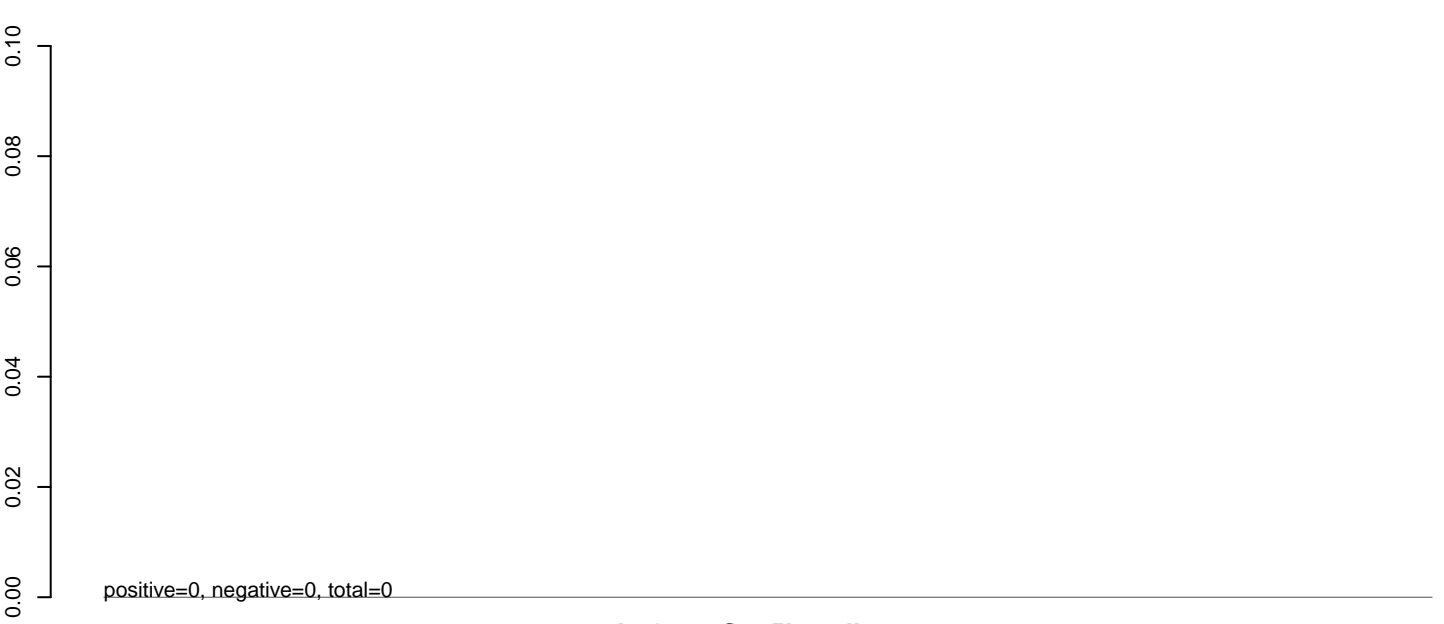
AnGam_Sua5b_cells.rep



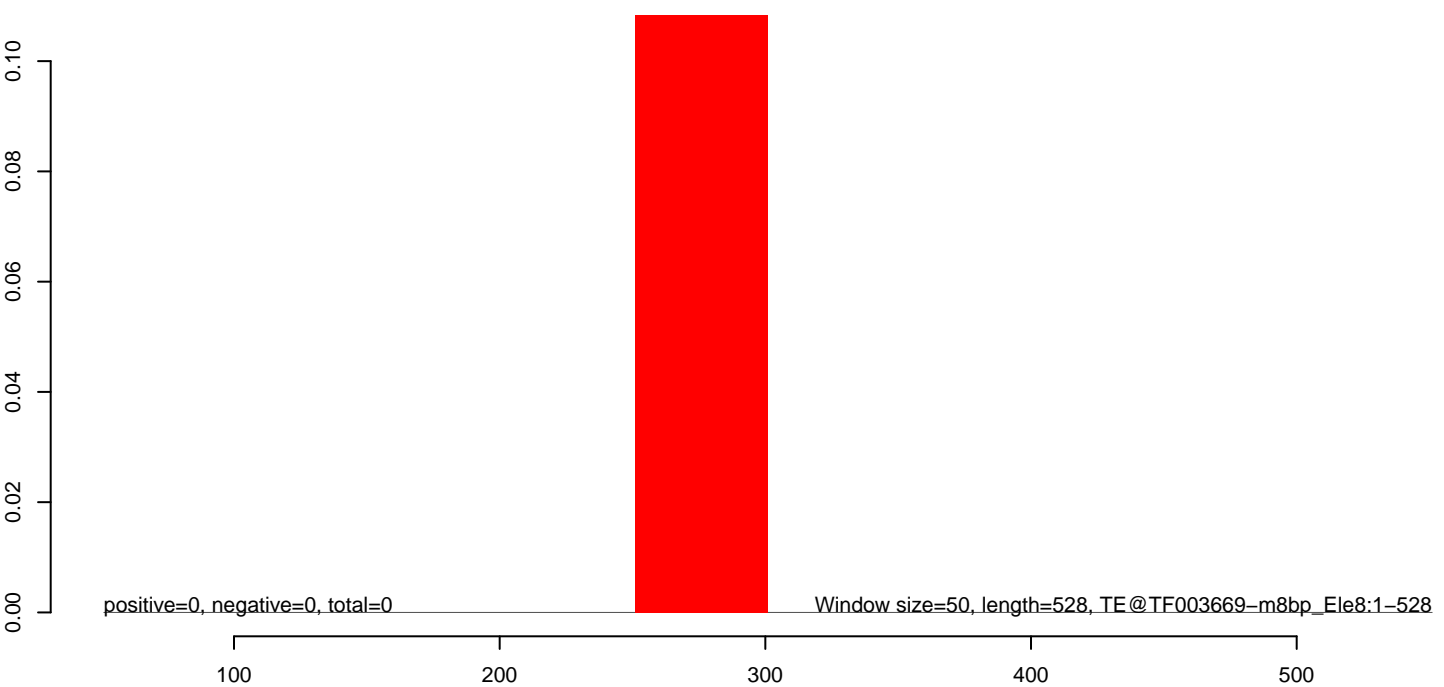
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.rep

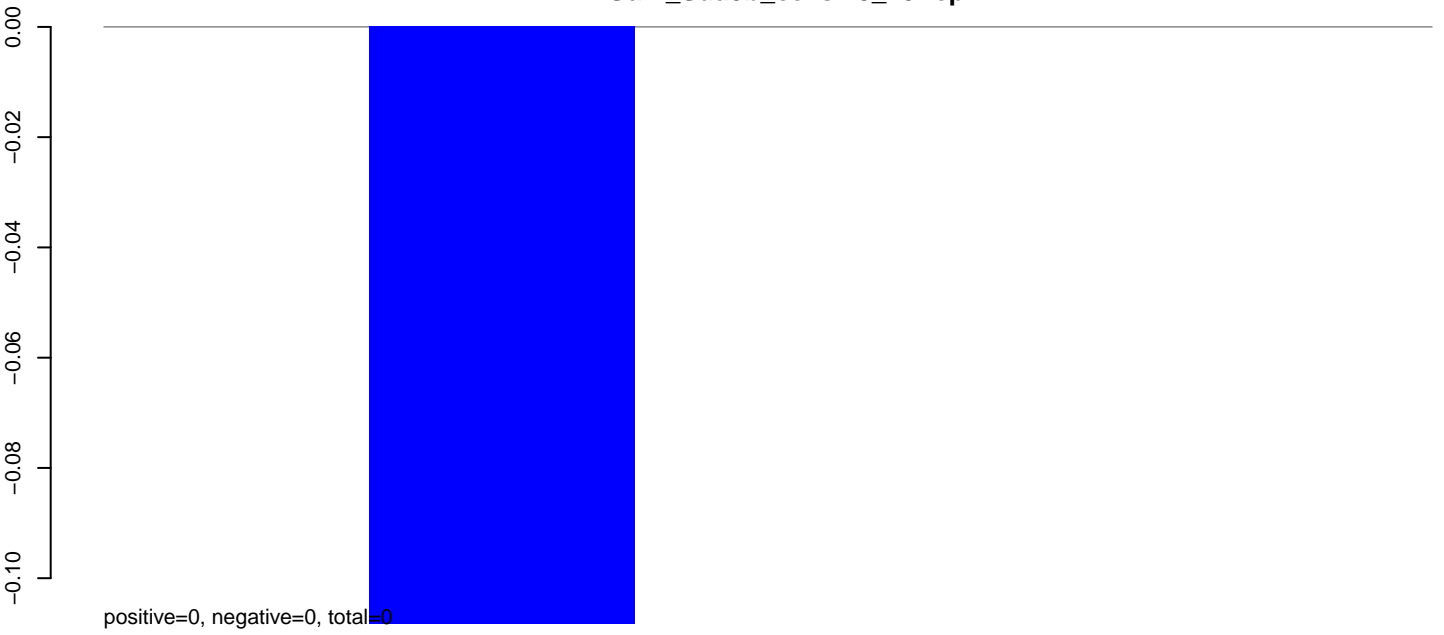
0.00
-0.02
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-0.06
-0.08
-0.10

positive=0, negative=0, total=0

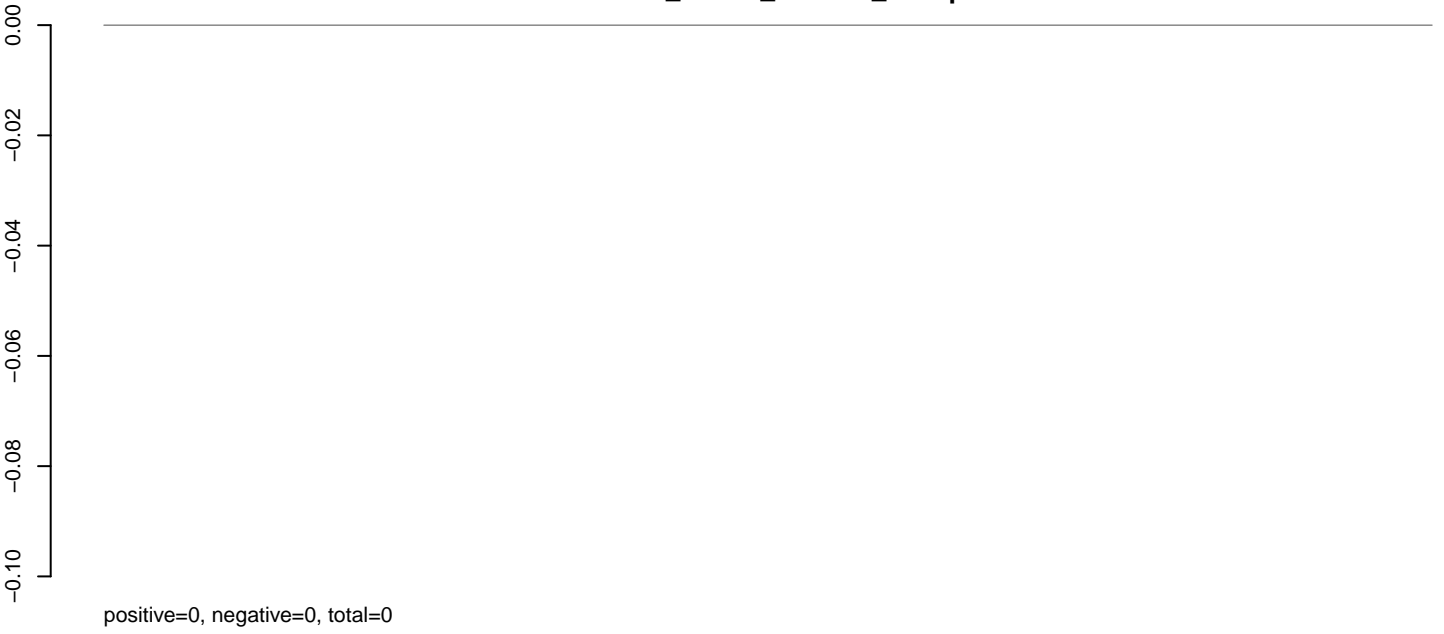
Window size=50, length=326, TE@TF003666-m8bp_Ele5:1-326

50 100 150 200 250 300 350

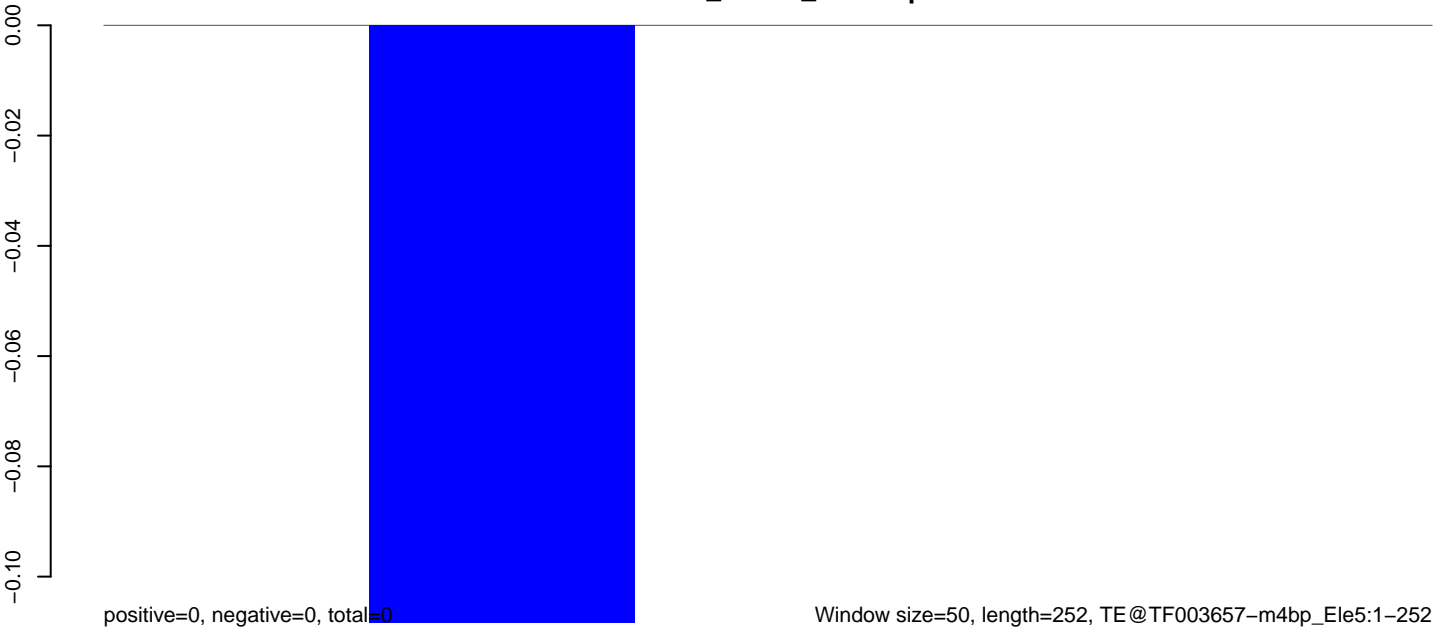
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



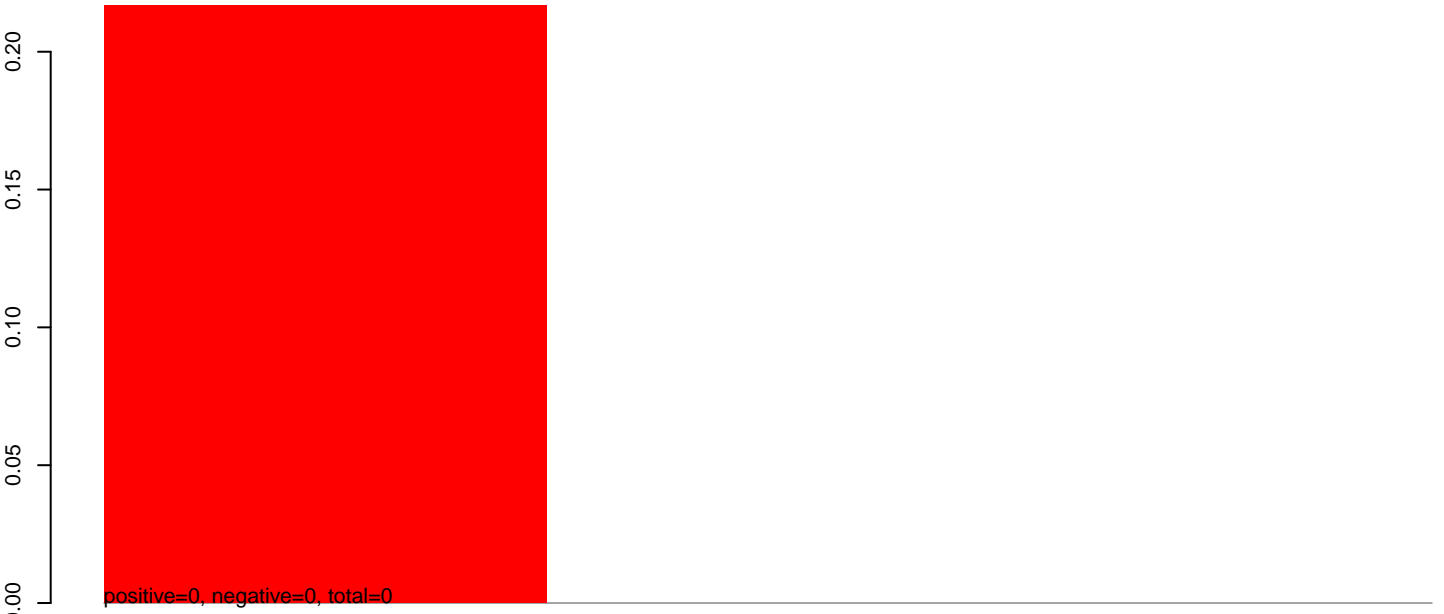
AnGam_Sua5b_cells.rep



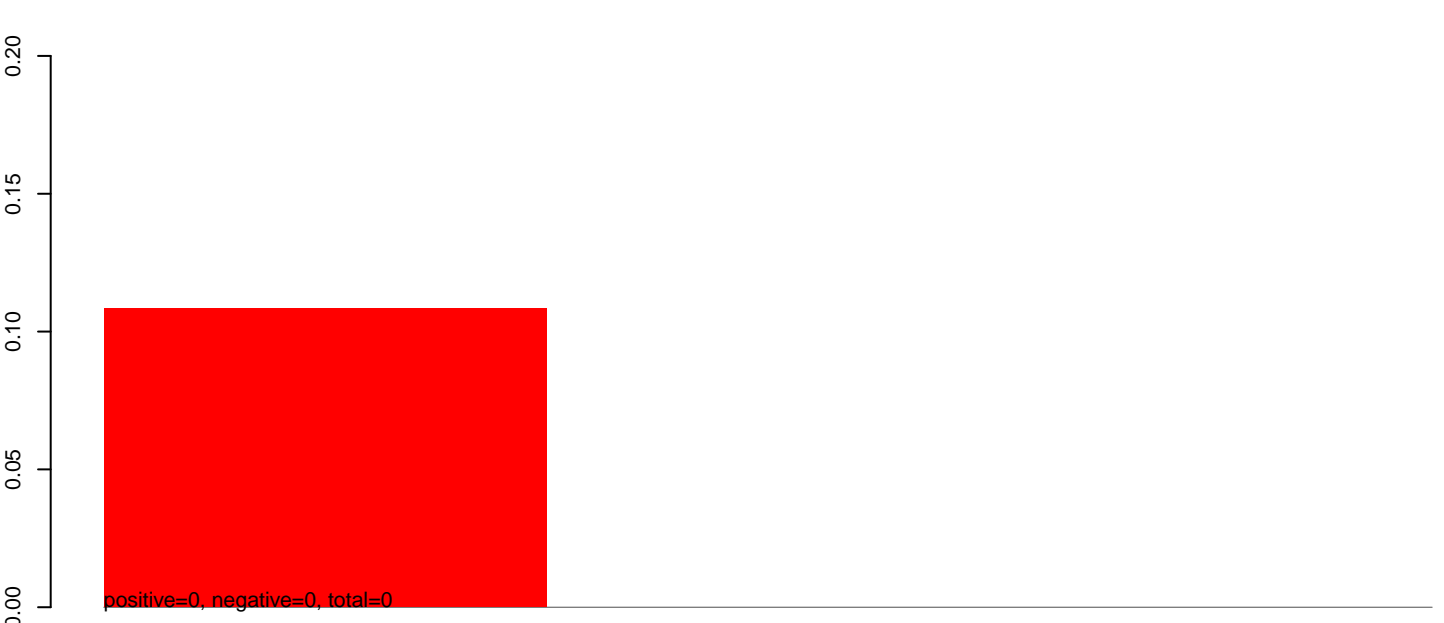
Window size=50, length=252, TE@TF003657-m4bp_Ele5:1-252

50 100 150 200 250 300

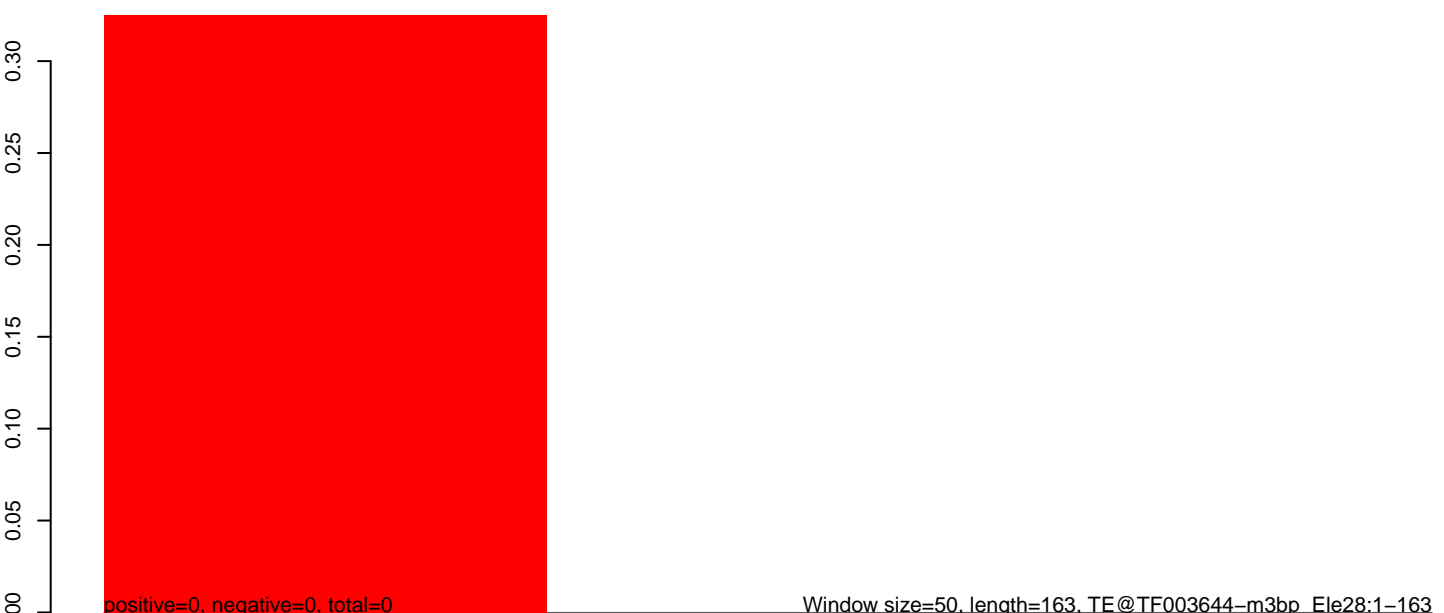
AnGam_Sua5b_cells.18_23.rep



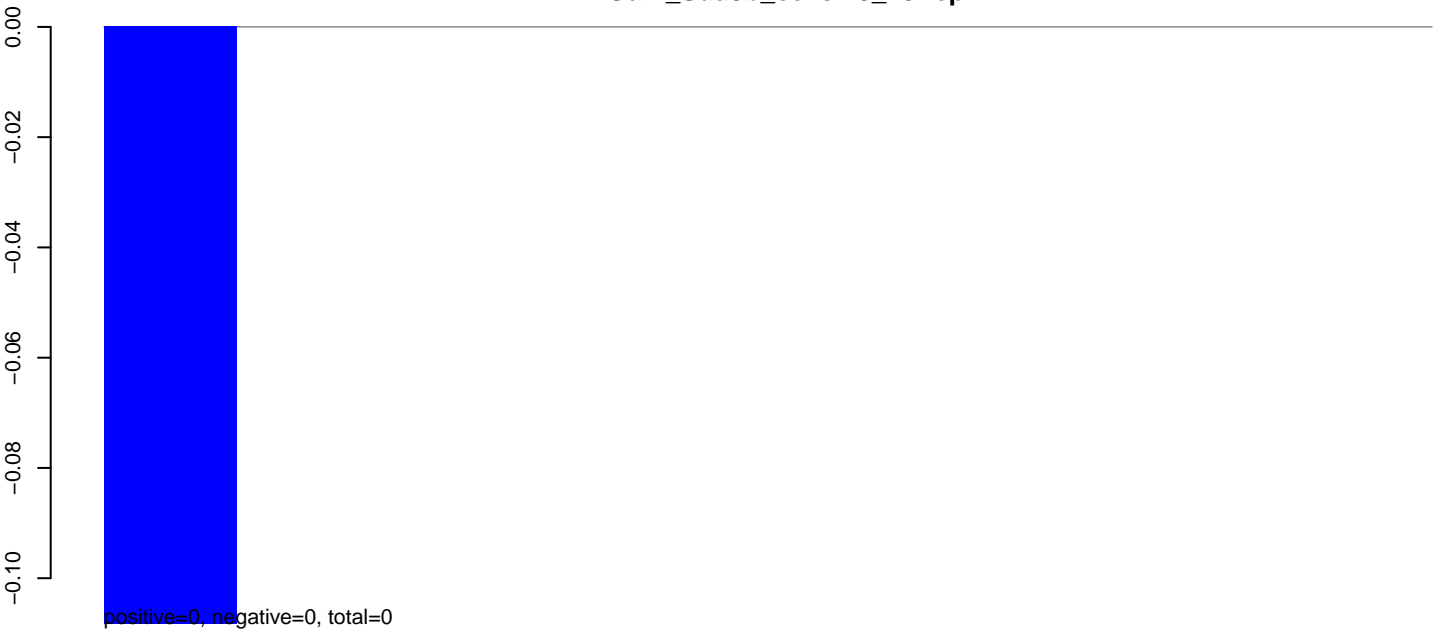
AnGam_Sua5b_cells.24_35.rep



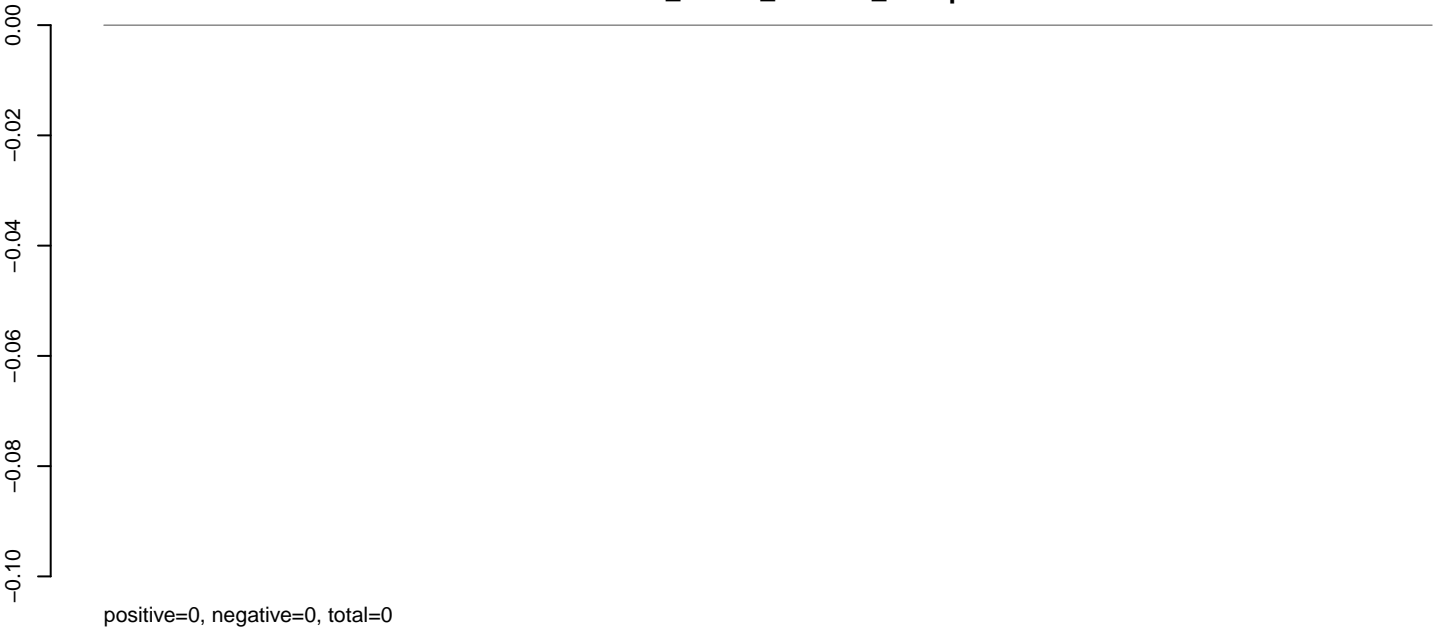
AnGam_Sua5b_cells.rep



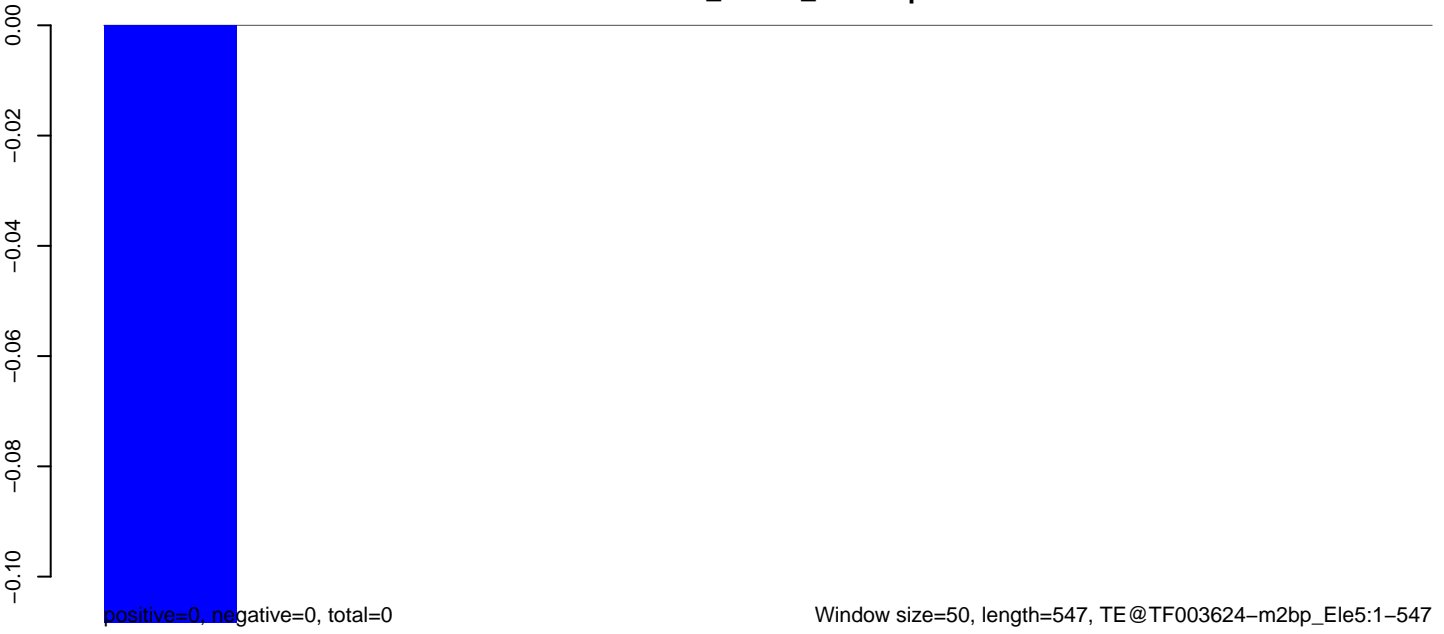
AnGam_Sua5b_cells.18_23.rep



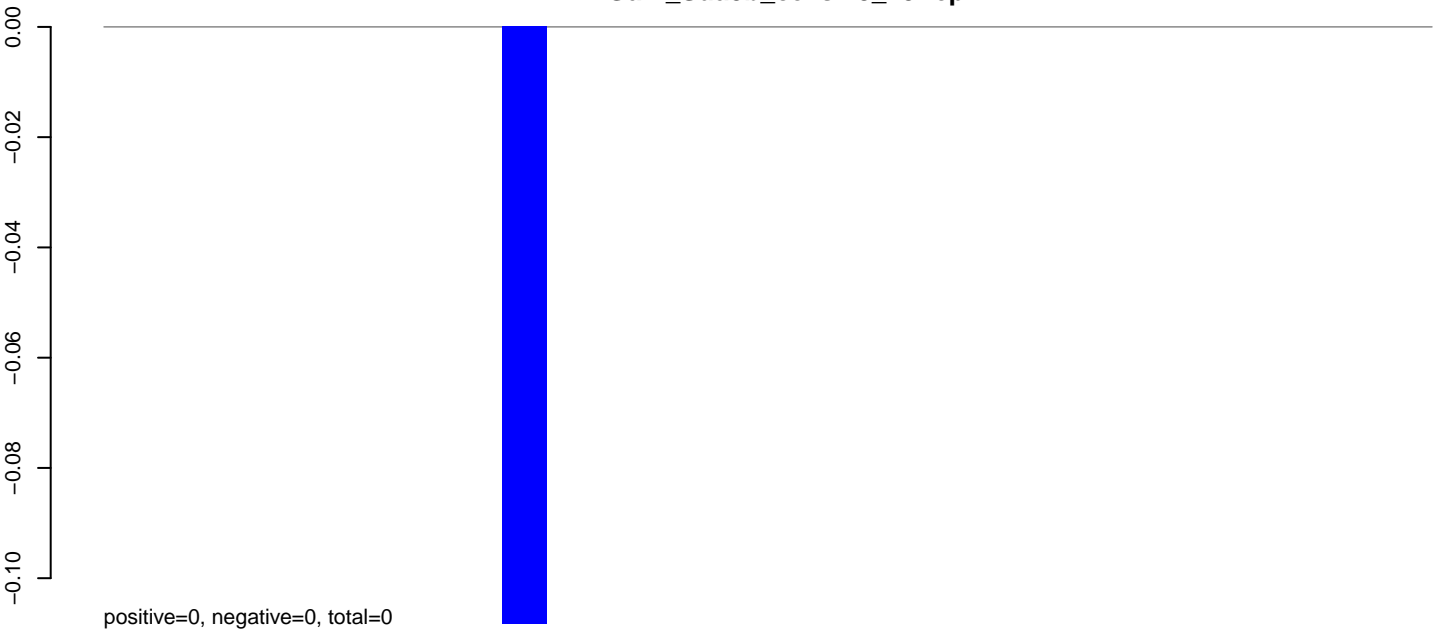
AnGam_Sua5b_cells.24_35.rep



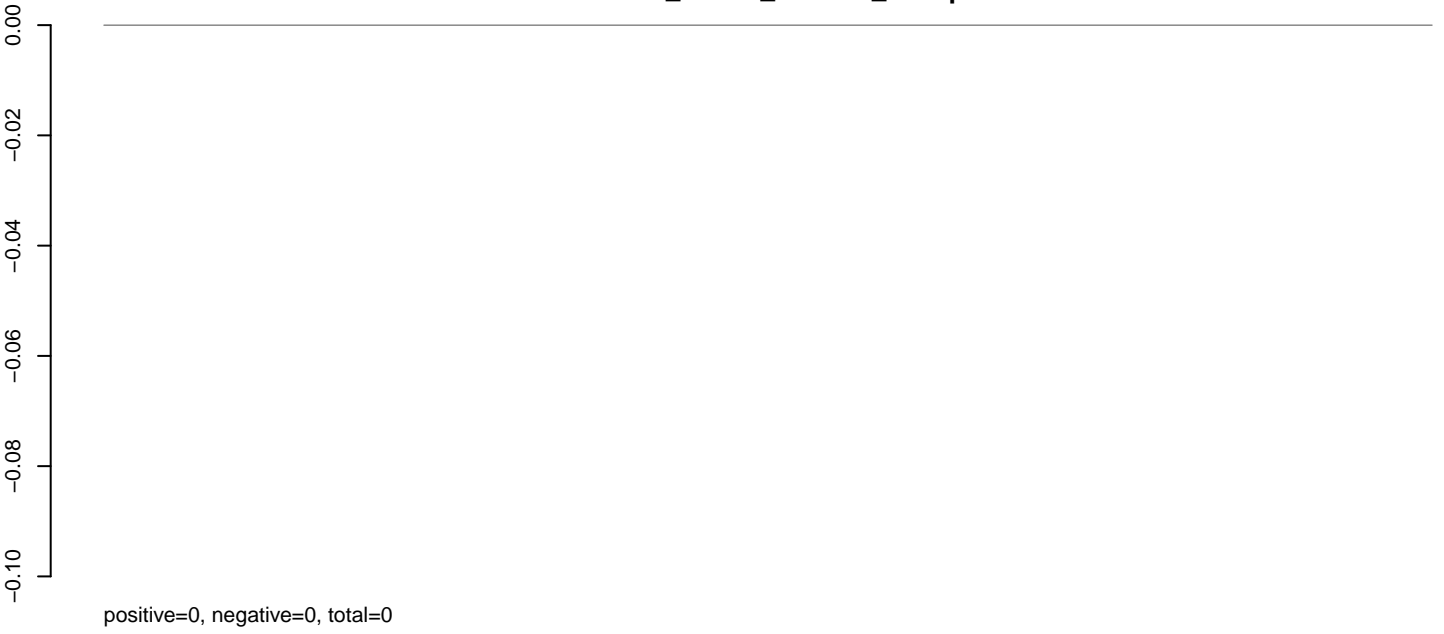
AnGam_Sua5b_cells.rep



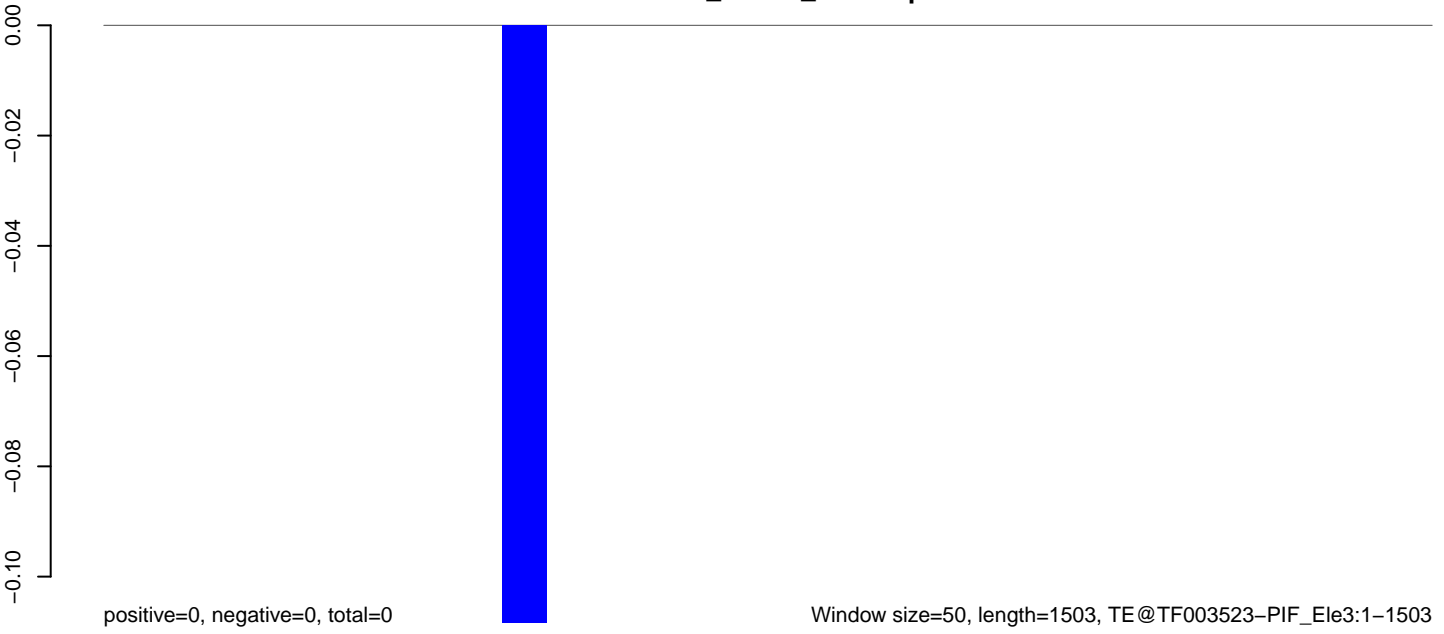
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

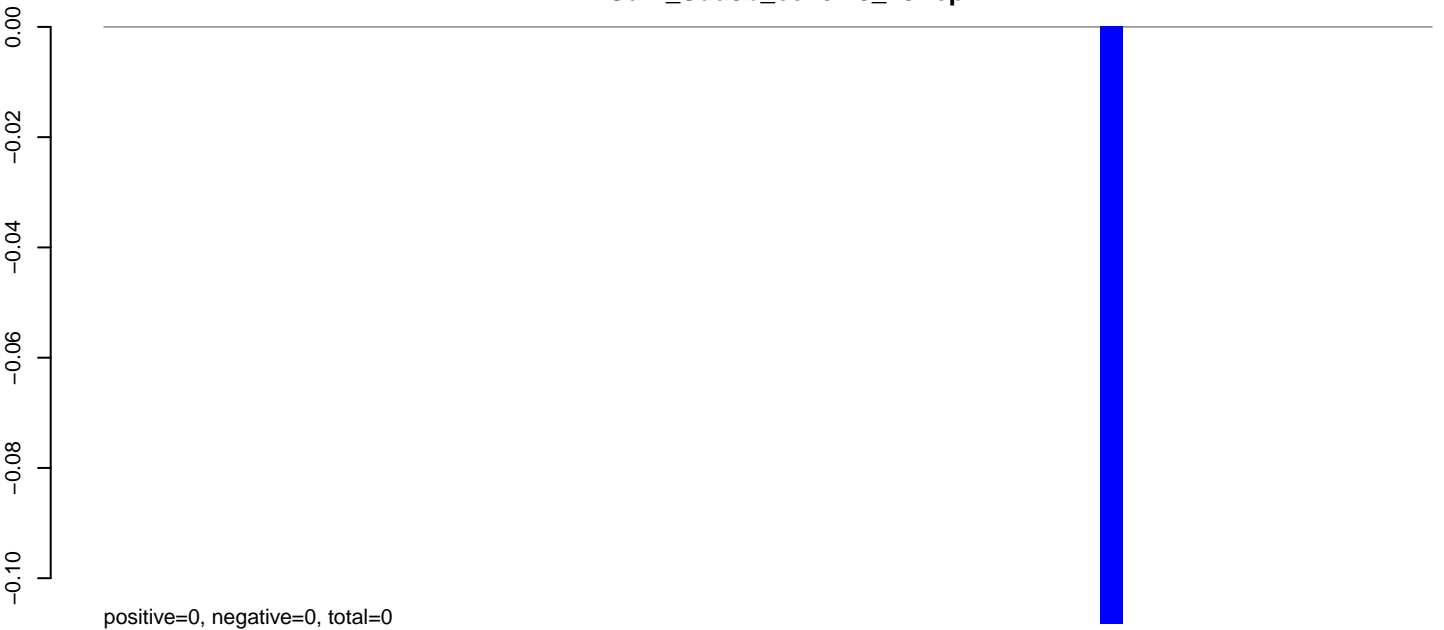


AnGam_Sua5b_cells.rep

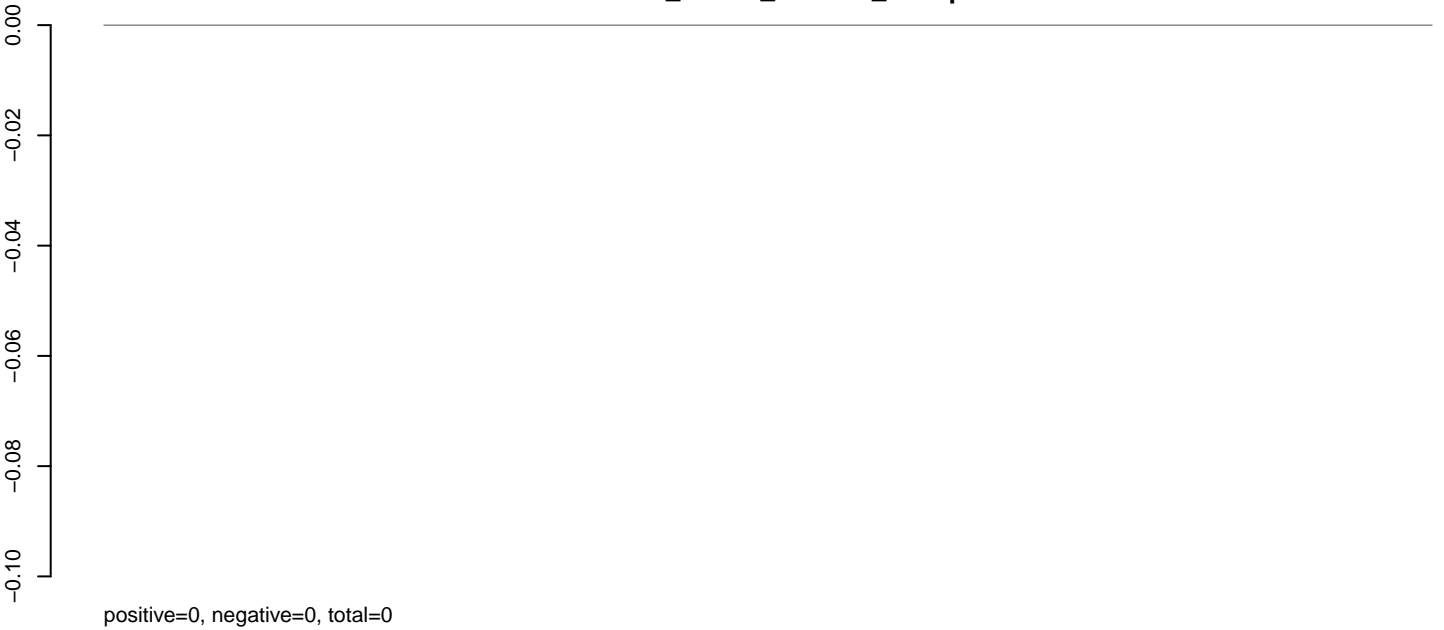


0 500 1000 1500

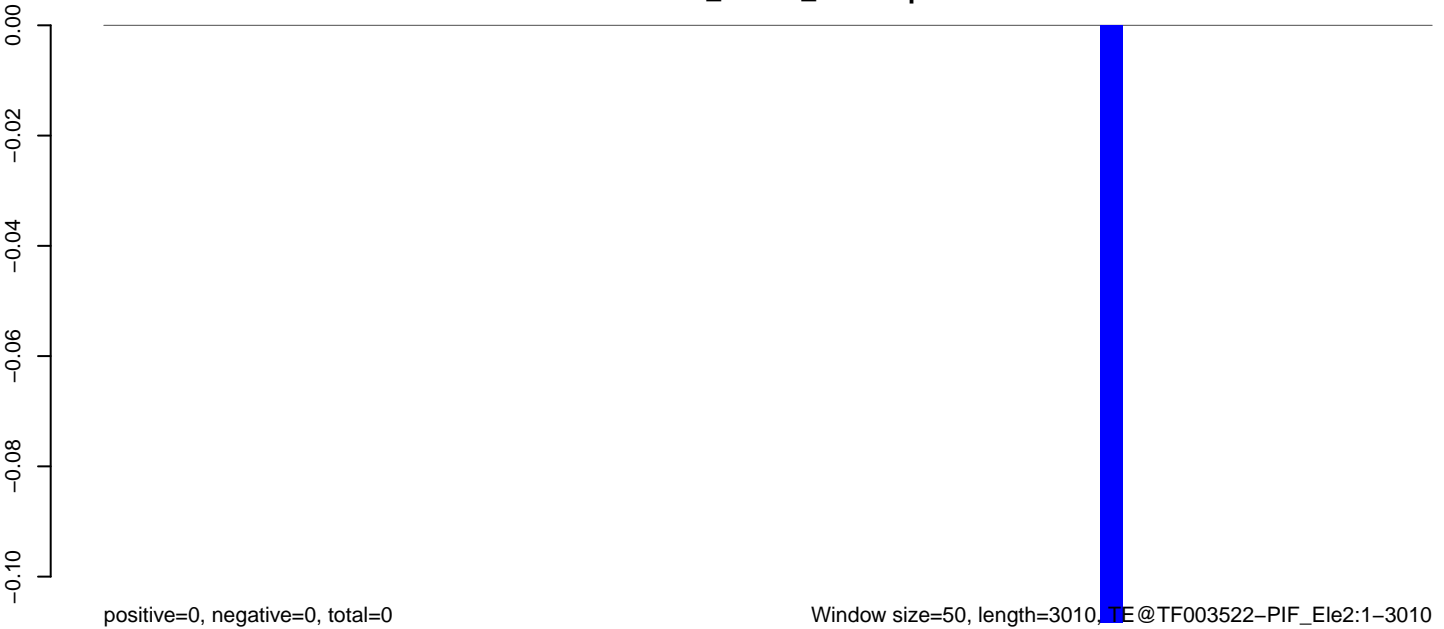
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

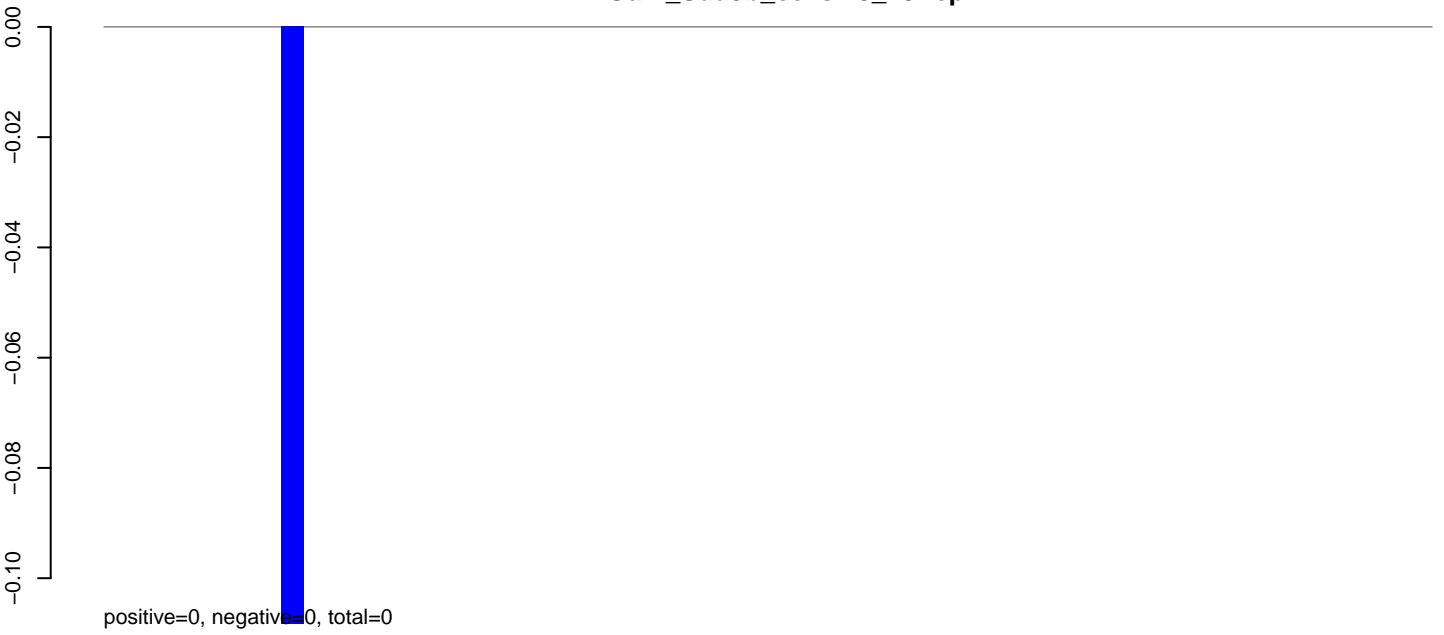


AnGam_Sua5b_cells.rep

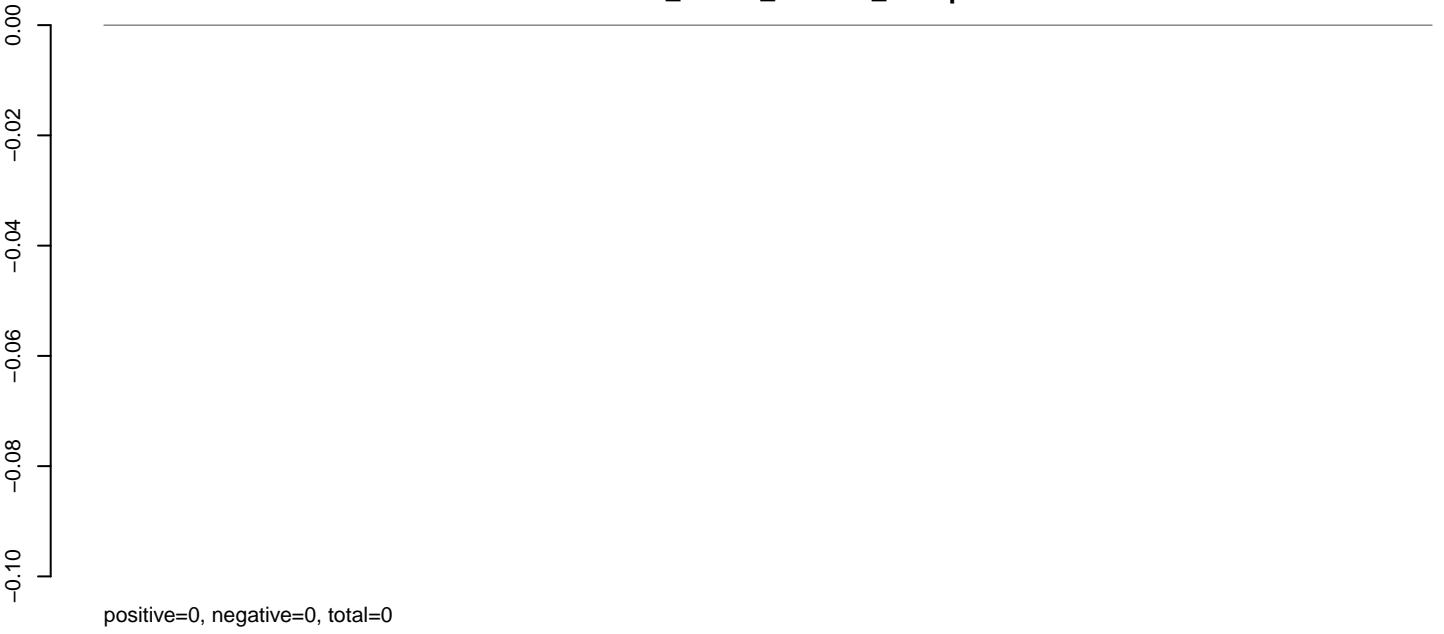


0 500 1000 1500 2000 2500 3000

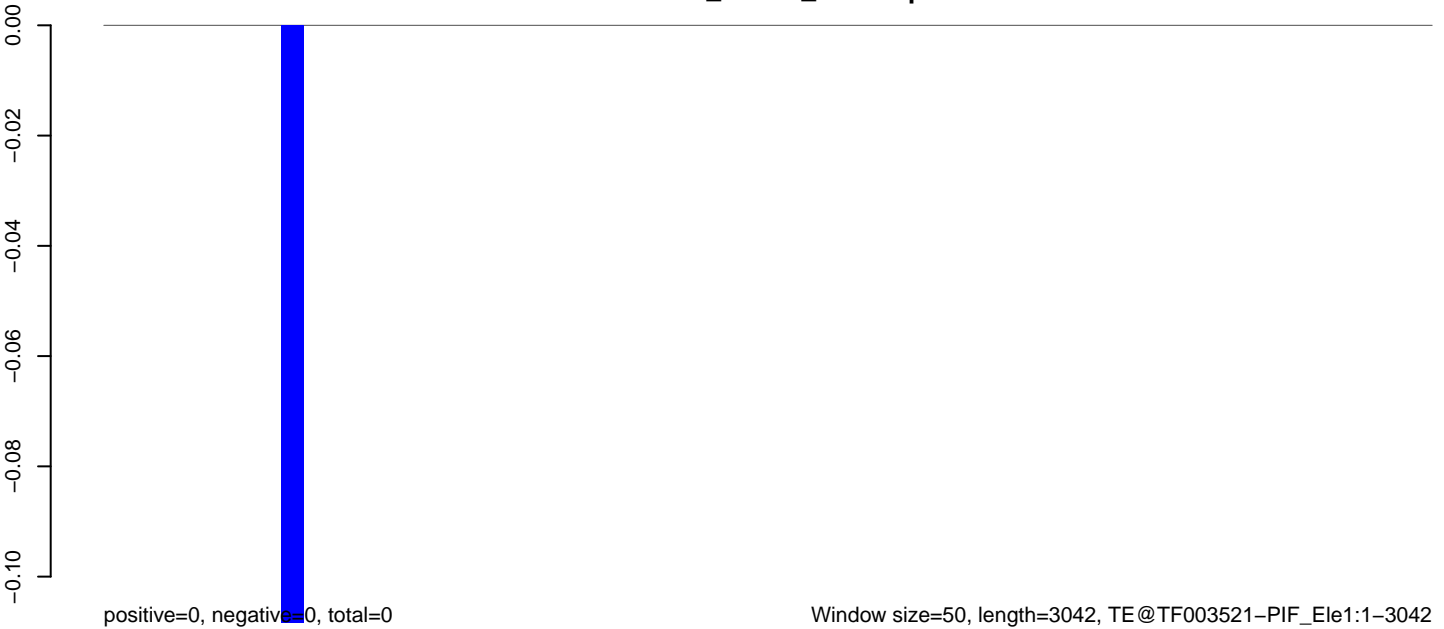
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



Window size=50, length=3042, TE@TF003521-PIF_Ele1:1-3042

0 500 1000 1500 2000 2500 3000

AnGam_Sua5b_cells.18_23.rep

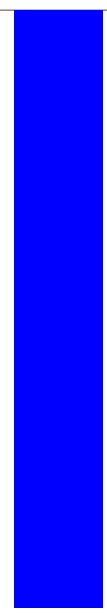
0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0



AnGam_Sua5b_cells.rep

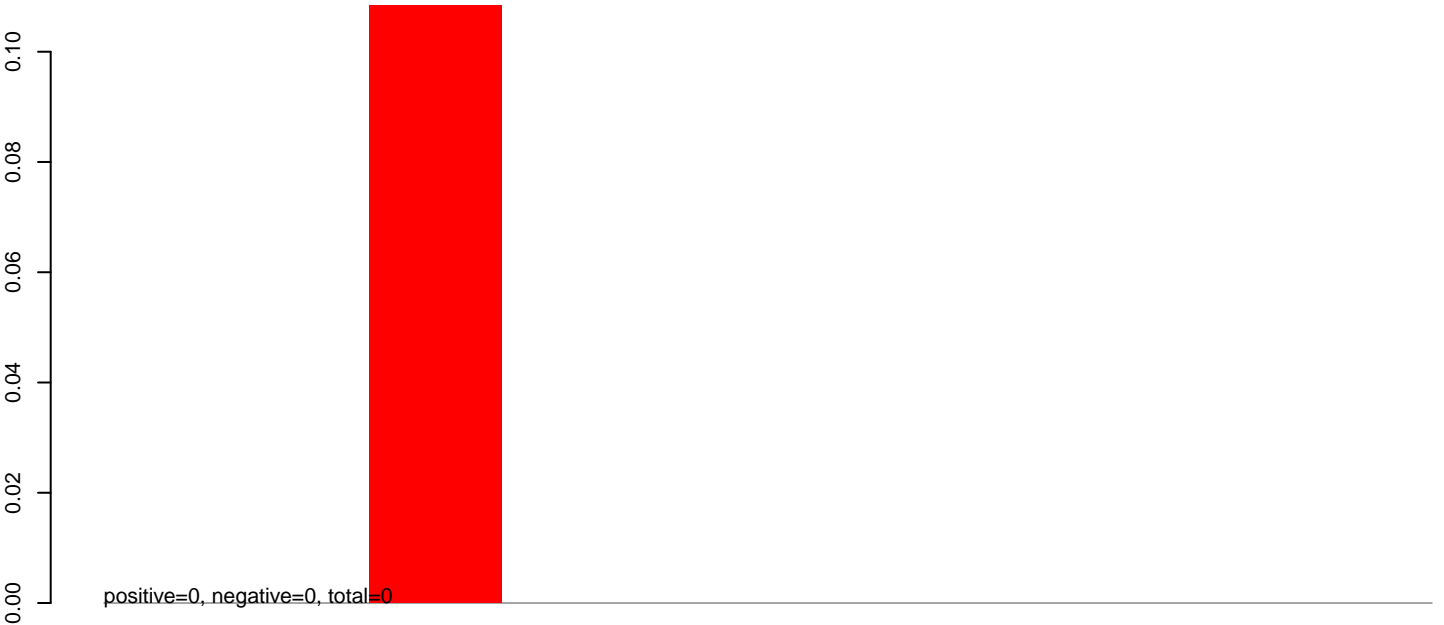
0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

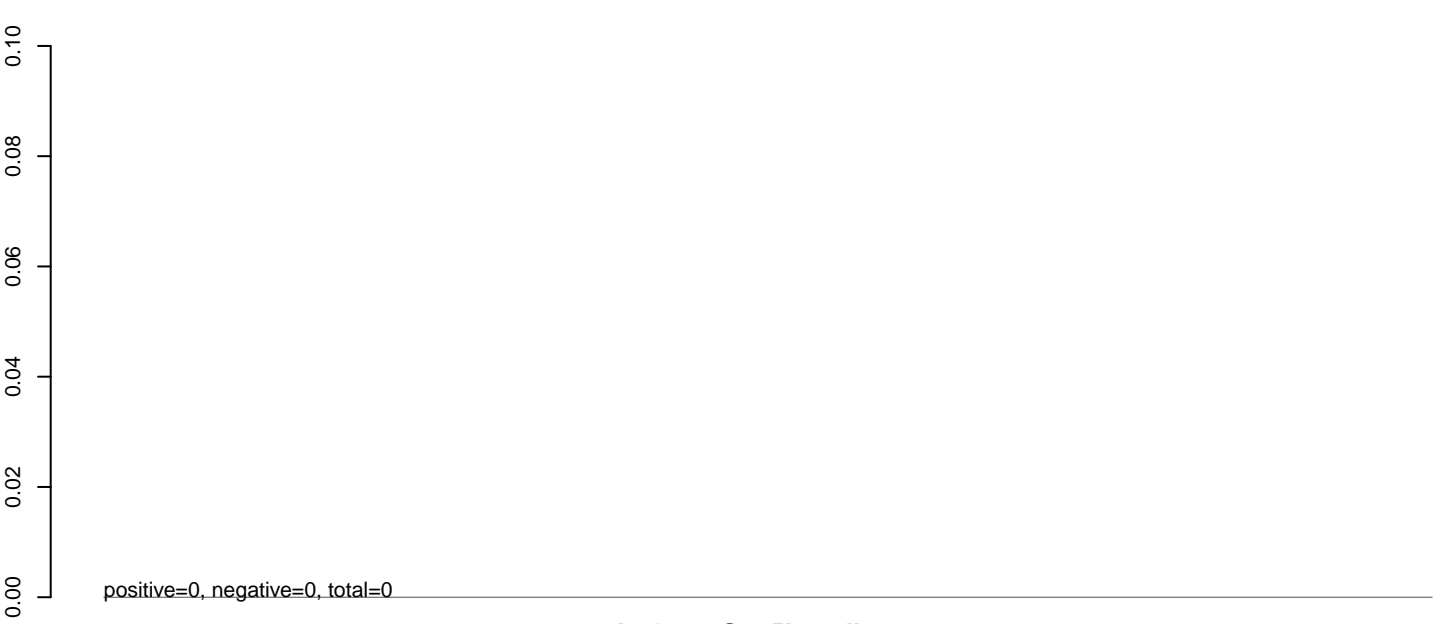
Window size=50, length=800, TE@TF003520-piggyBac_Ele3:1-800

200 400 600 800

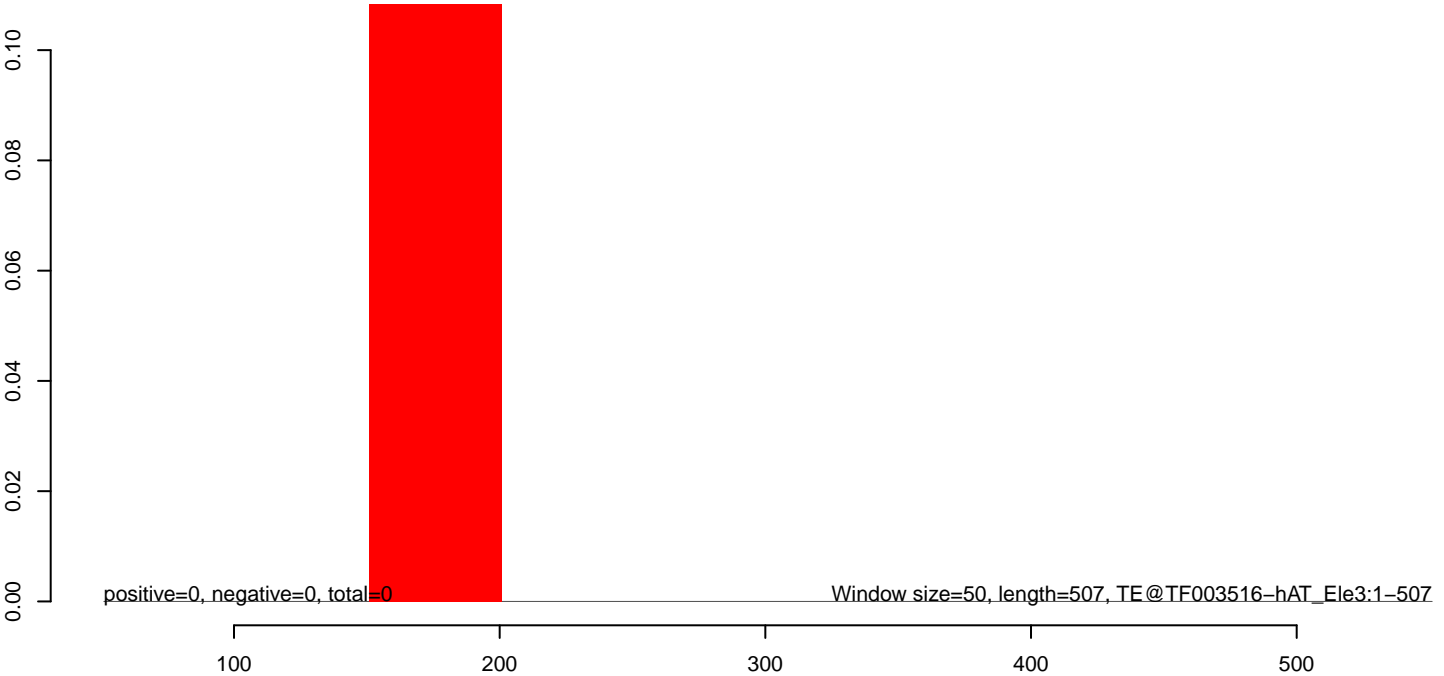
AnGam_Sua5b_cells.18_23.rep



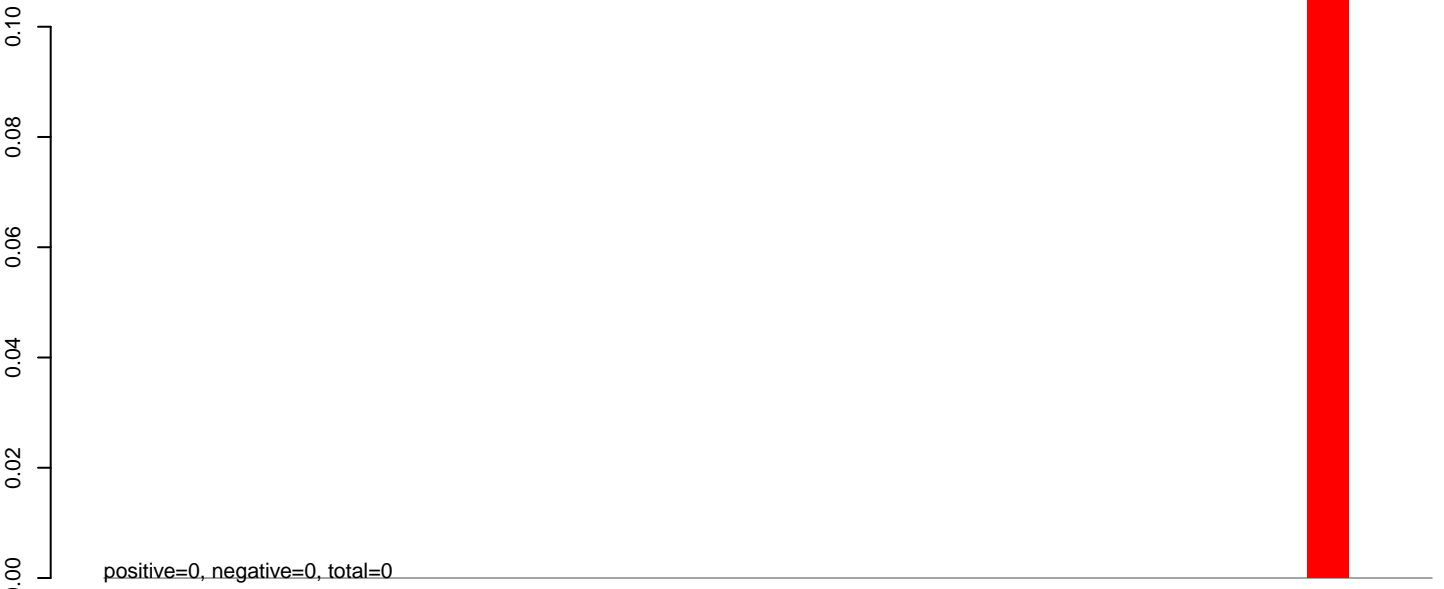
AnGam_Sua5b_cells.24_35.rep



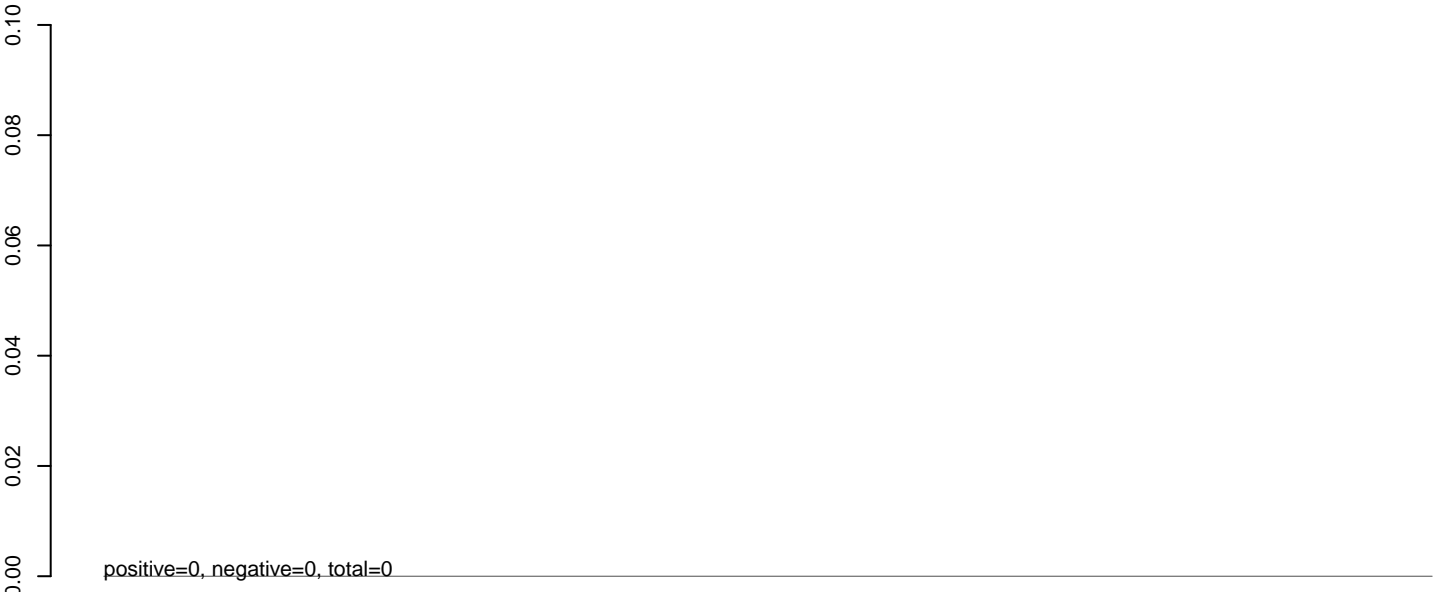
AnGam_Sua5b_cells.rep



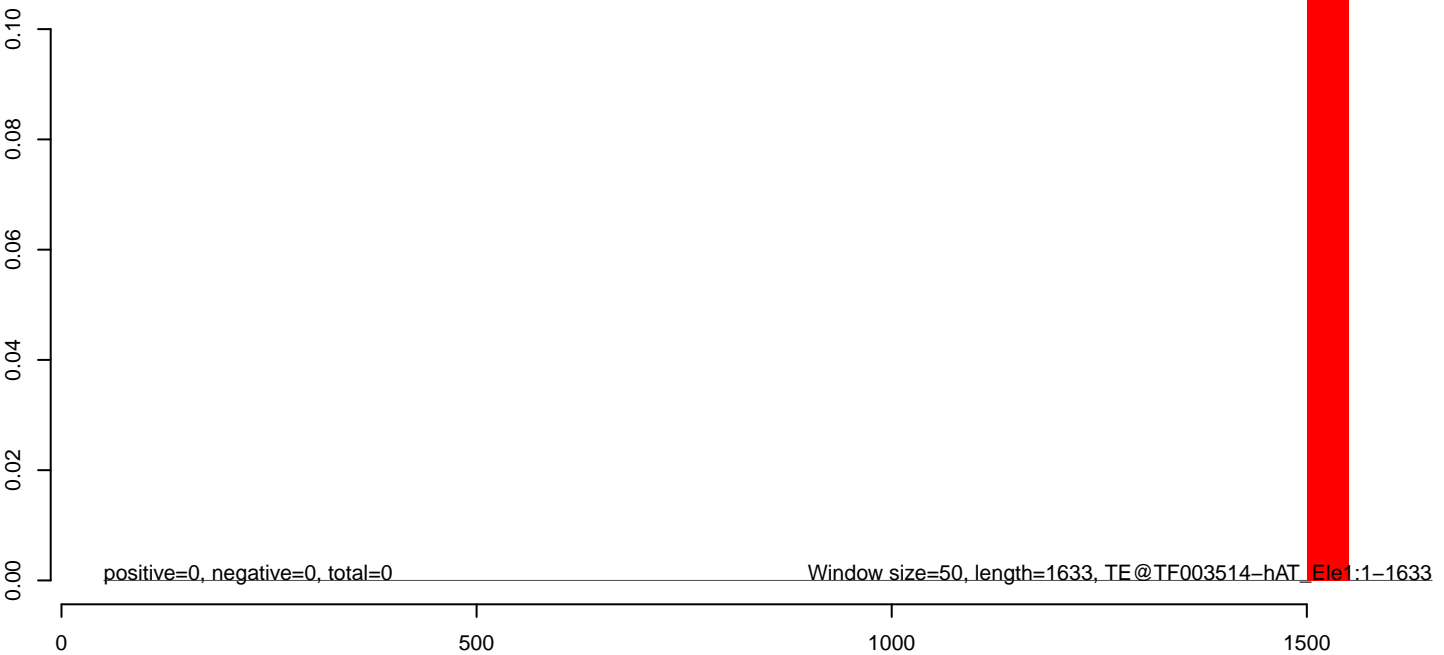
AnGam_Sua5b_cells.18_23.rep



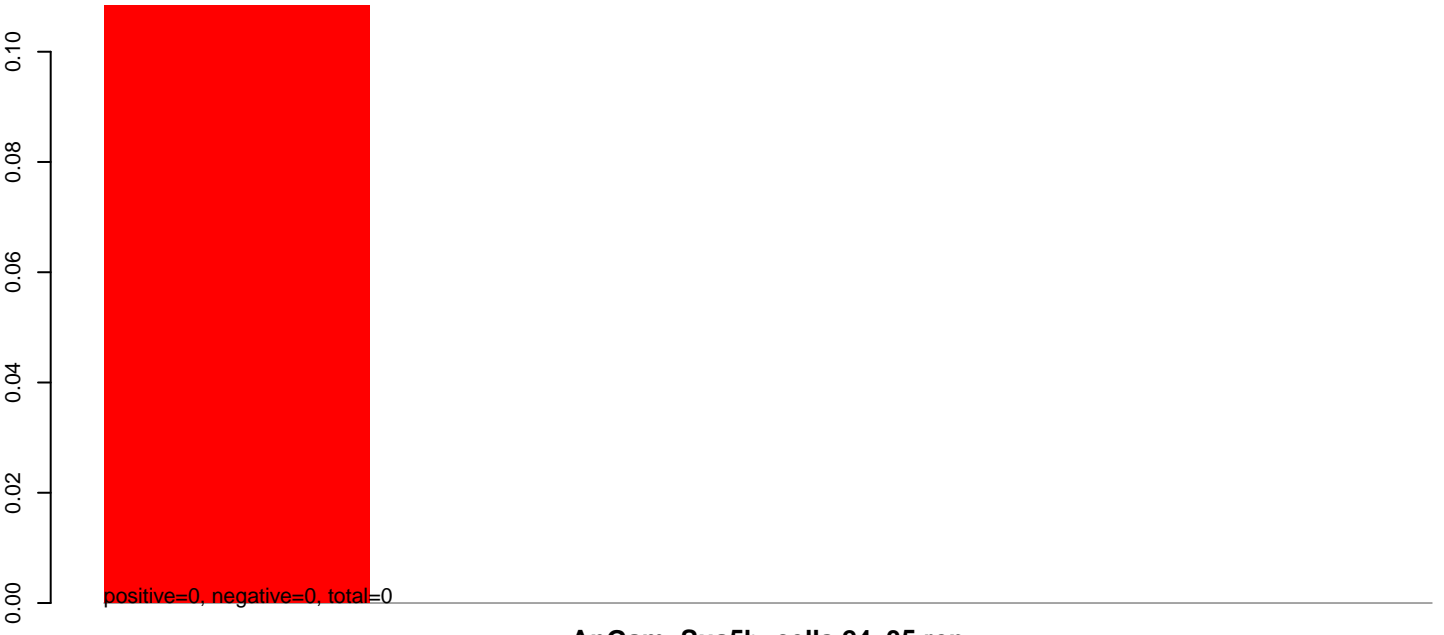
AnGam_Sua5b_cells.24_35.rep



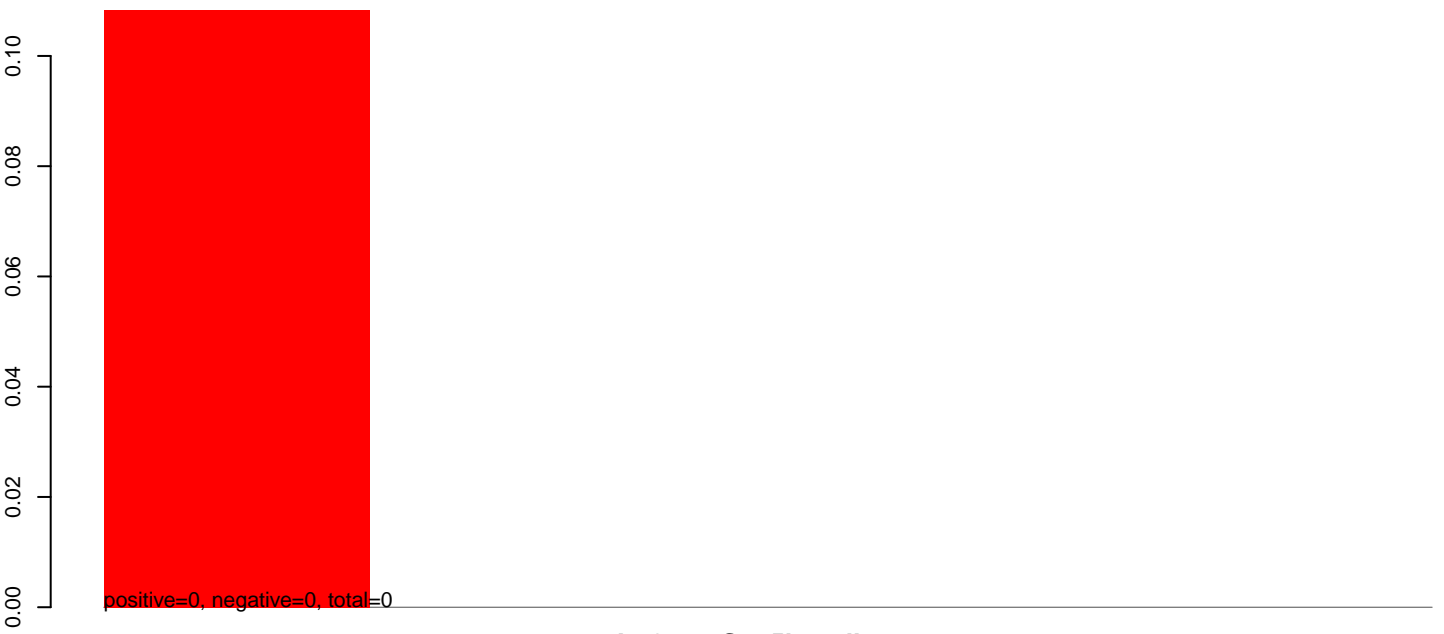
AnGam_Sua5b_cells.rep



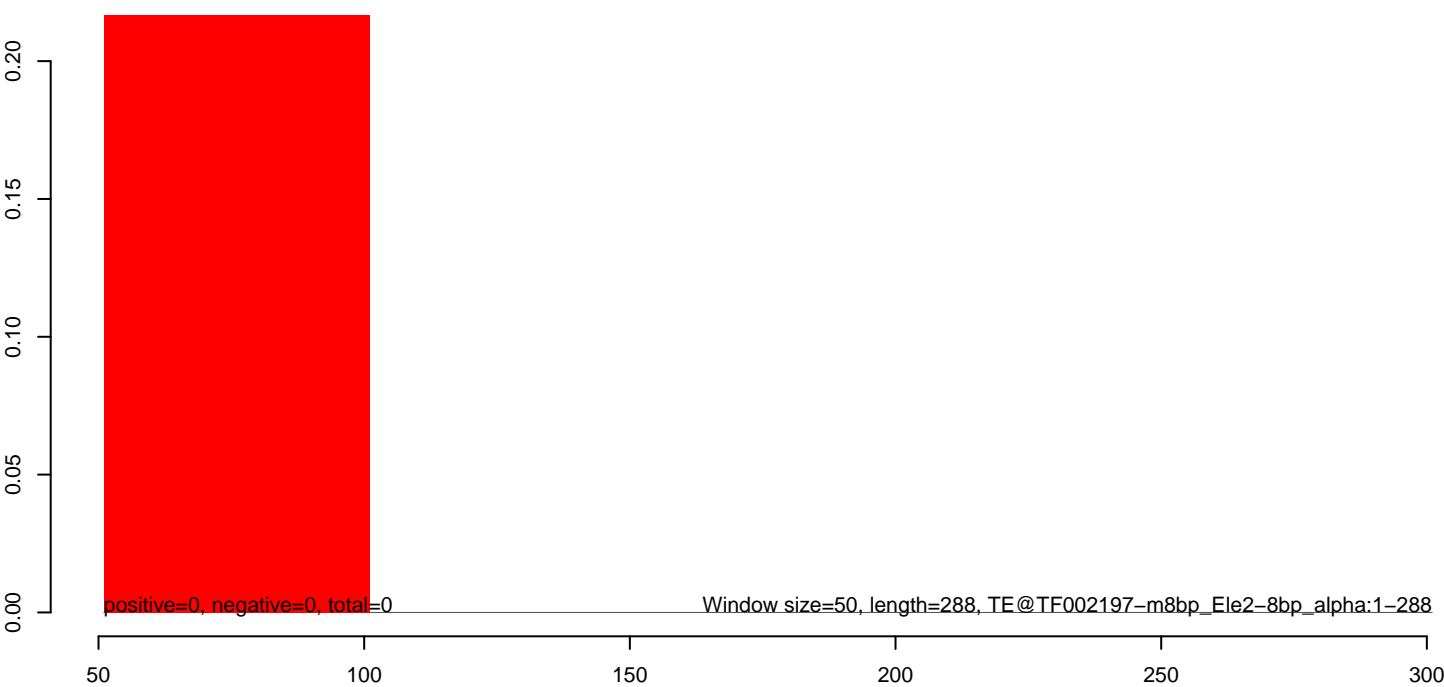
AnGam_Sua5b_cells.18_23.rep



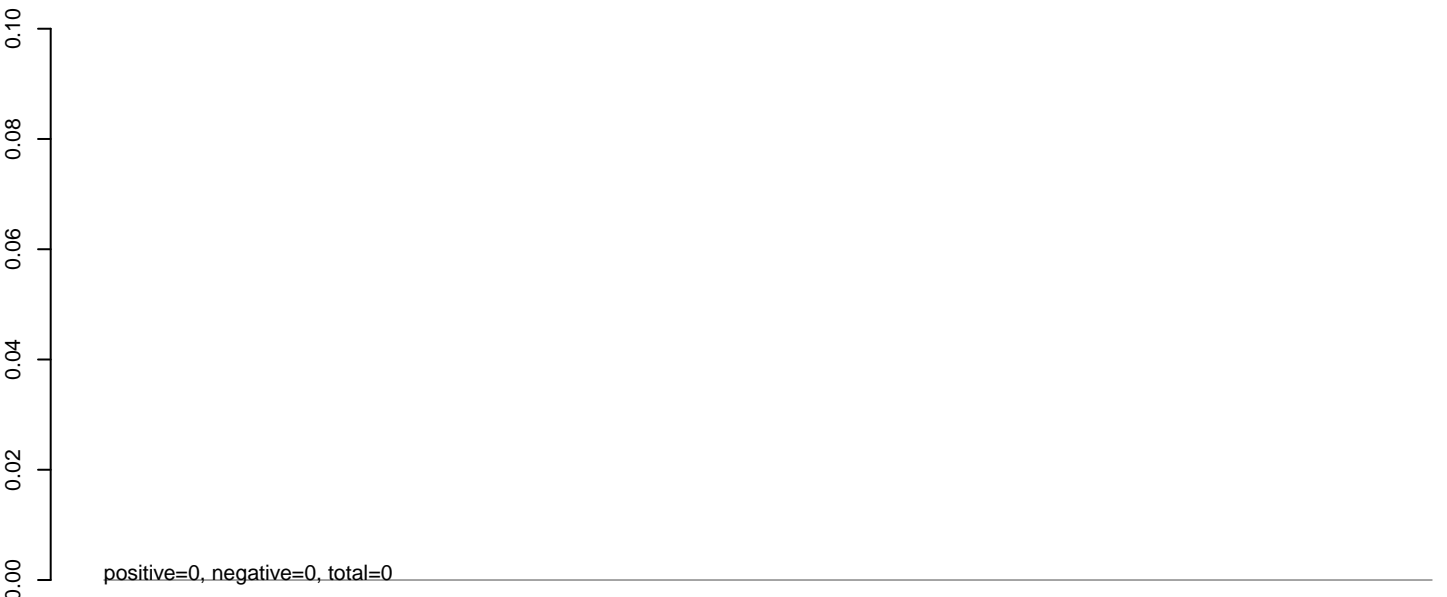
AnGam_Sua5b_cells.24_35.rep



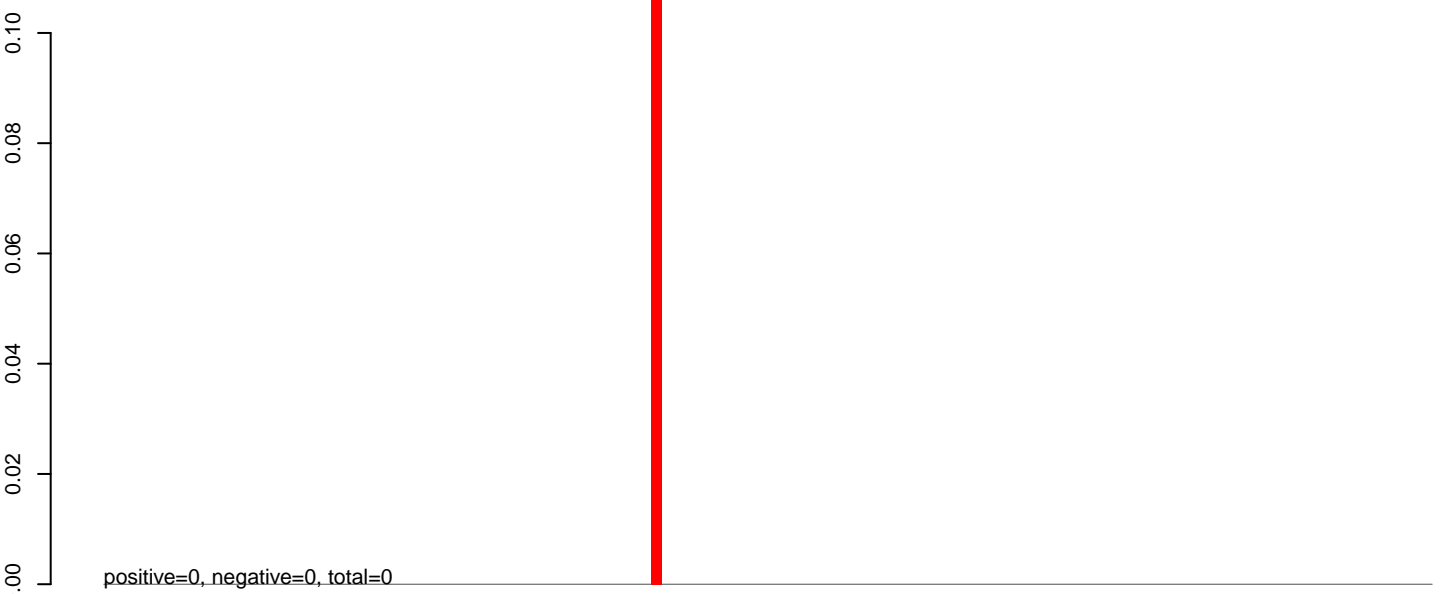
AnGam_Sua5b_cells.rep



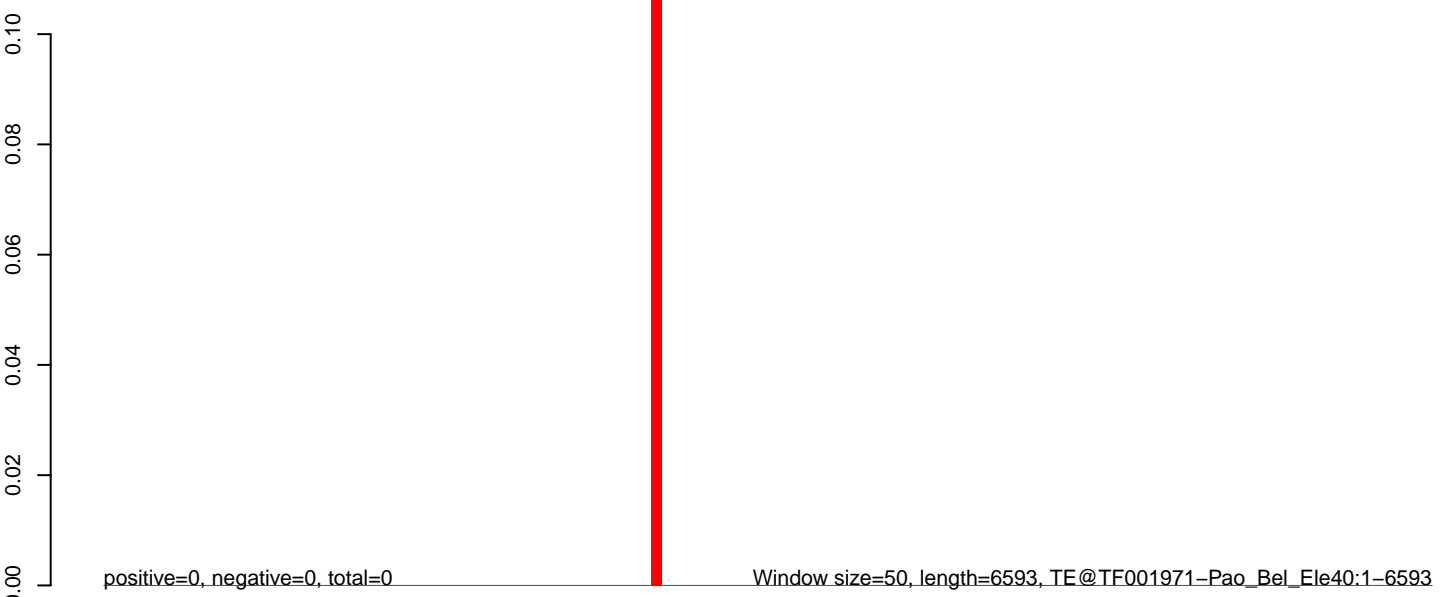
AnGam_Sua5b_cells.18_23.rep



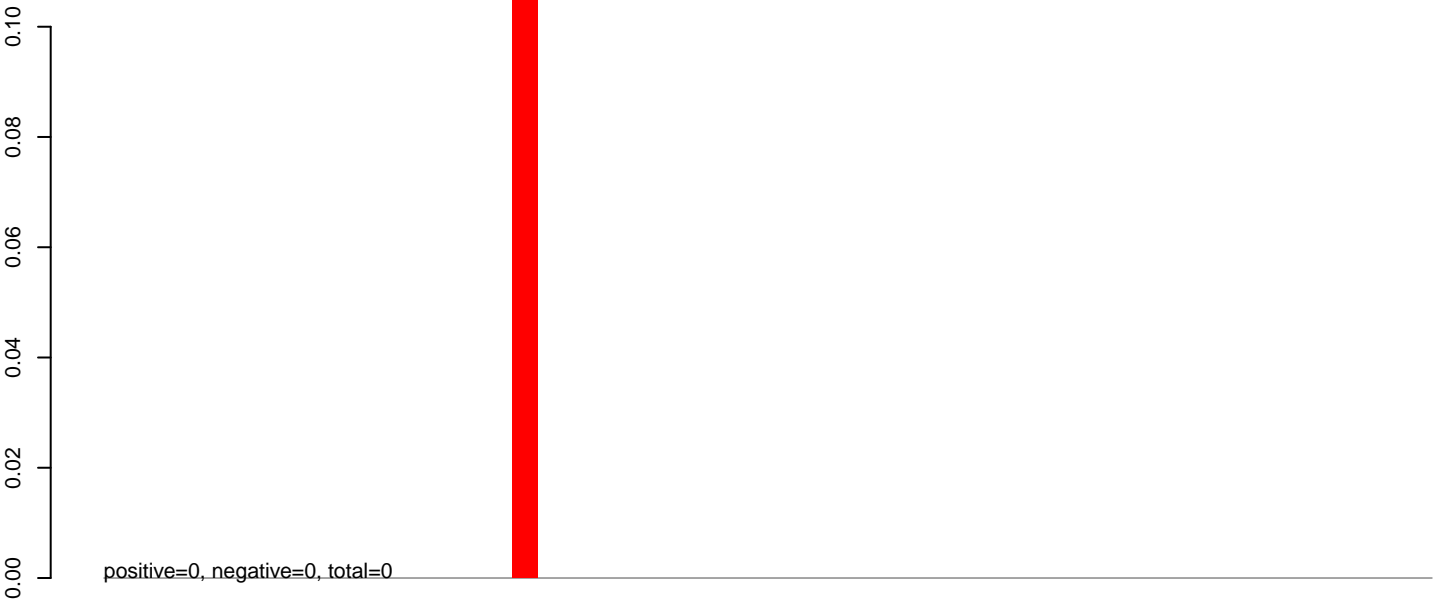
AnGam_Sua5b_cells.24_35.rep



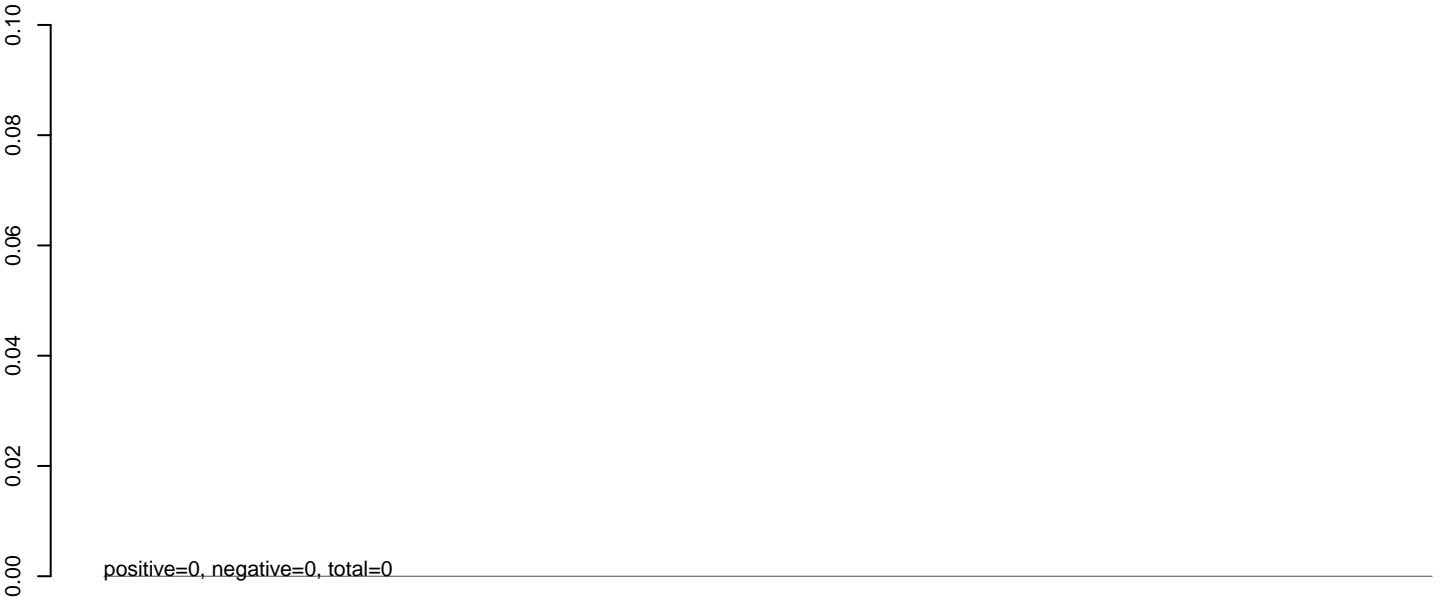
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

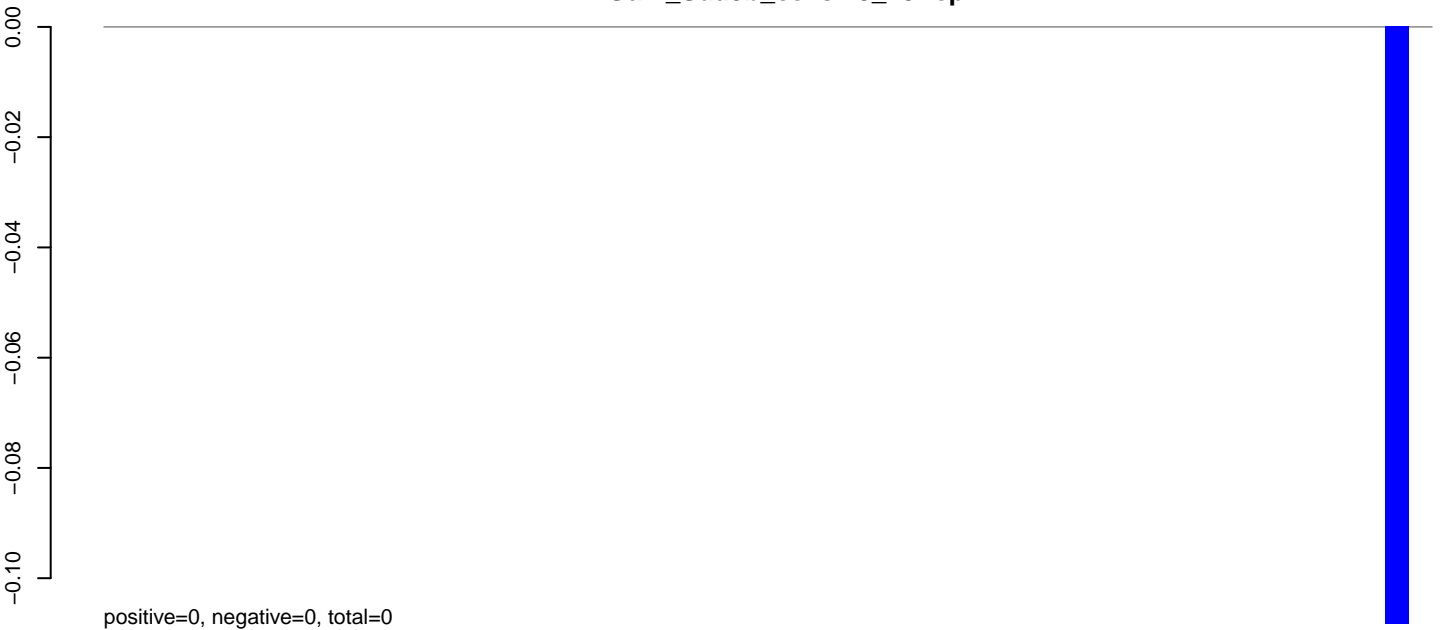


AnGam_Sua5b_cells.rep

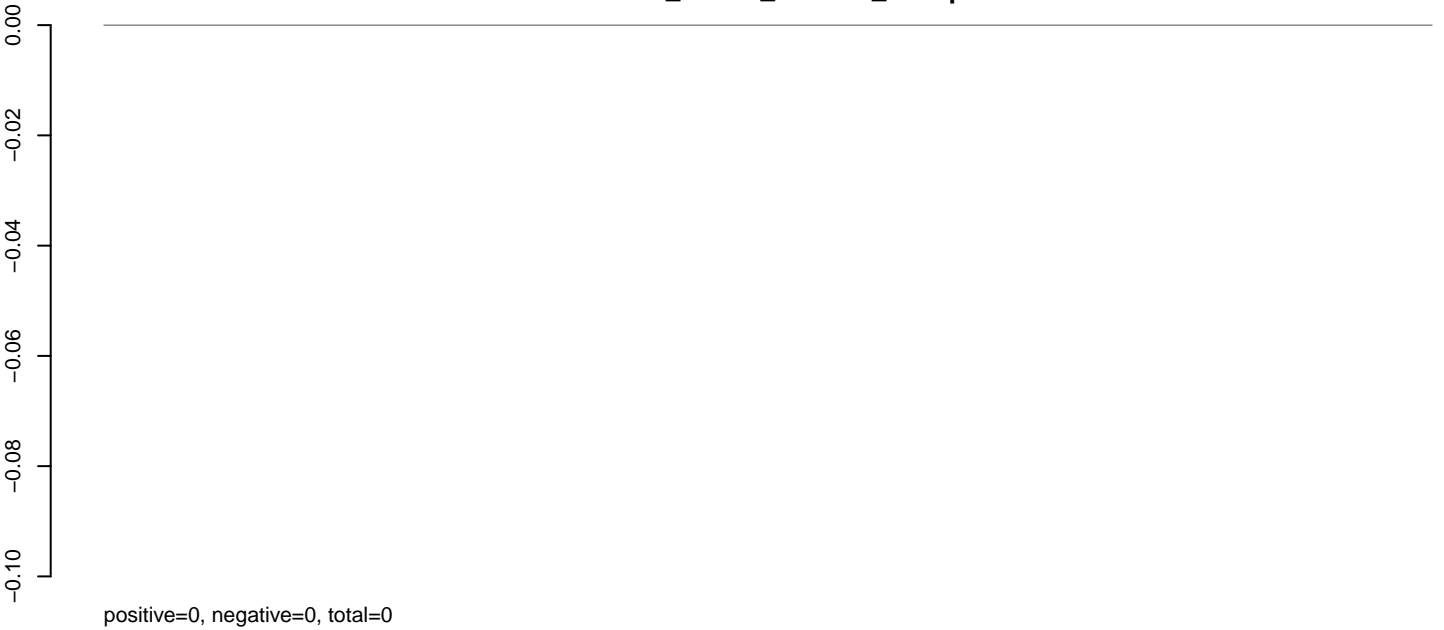


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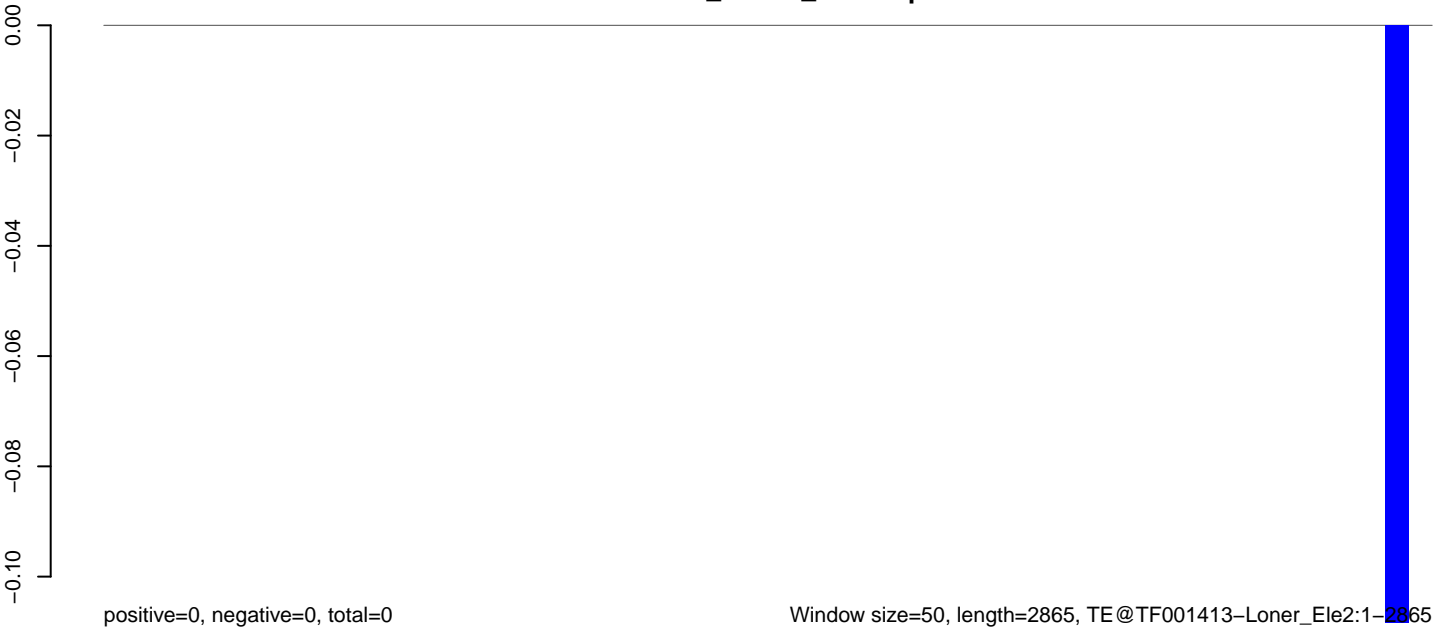
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



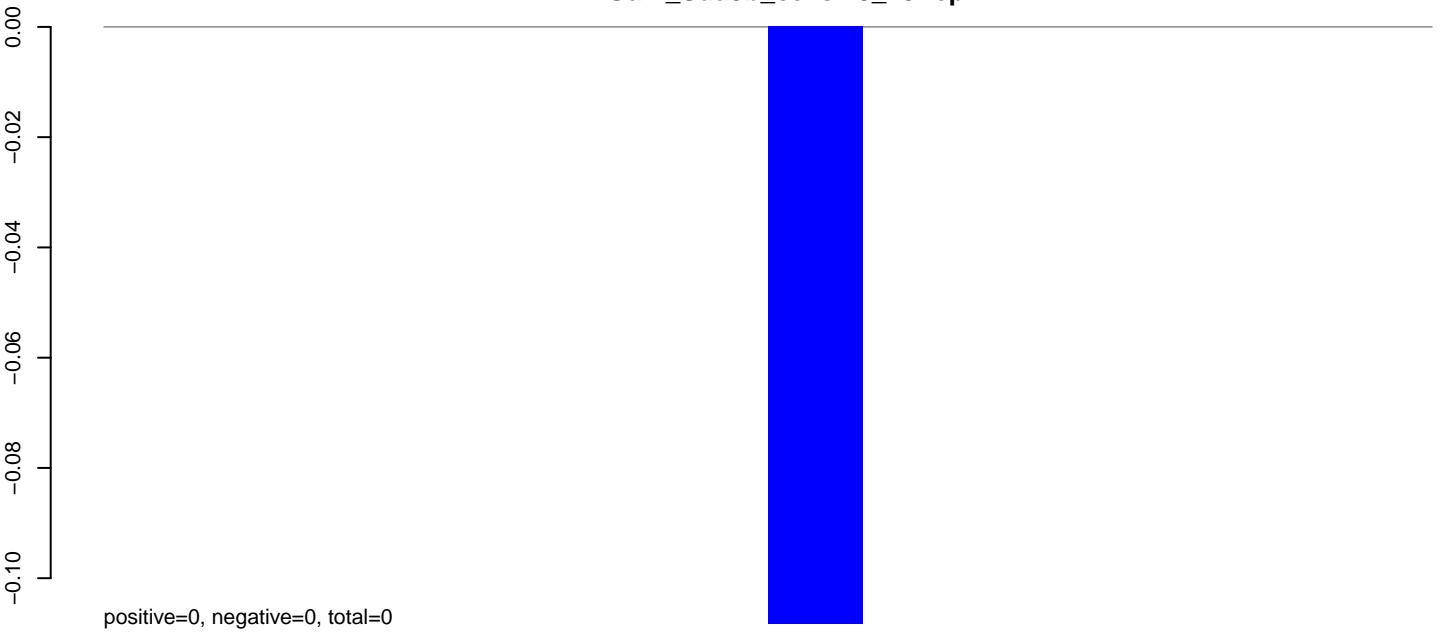
AnGam_Sua5b_cells.rep



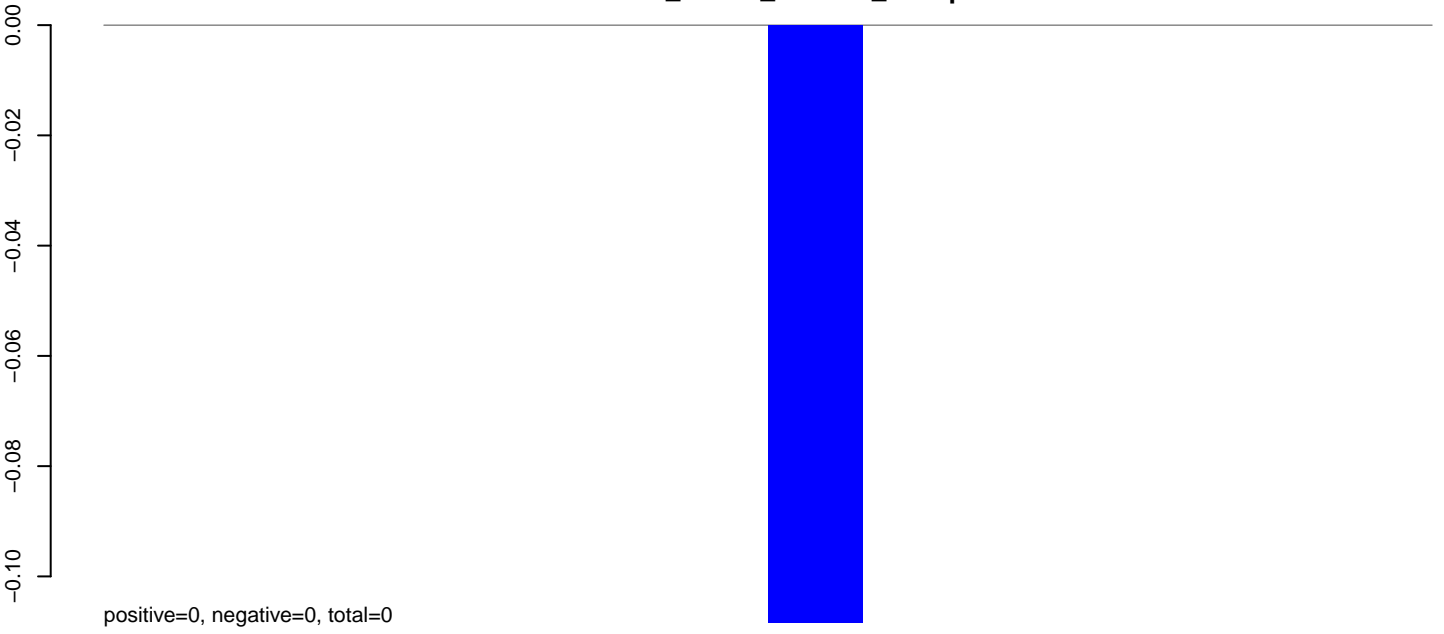
Window size=50, length=2865, TE@TF001413-Loner_Ele2:1-2865

0 500 1000 1500 2000 2500 3000

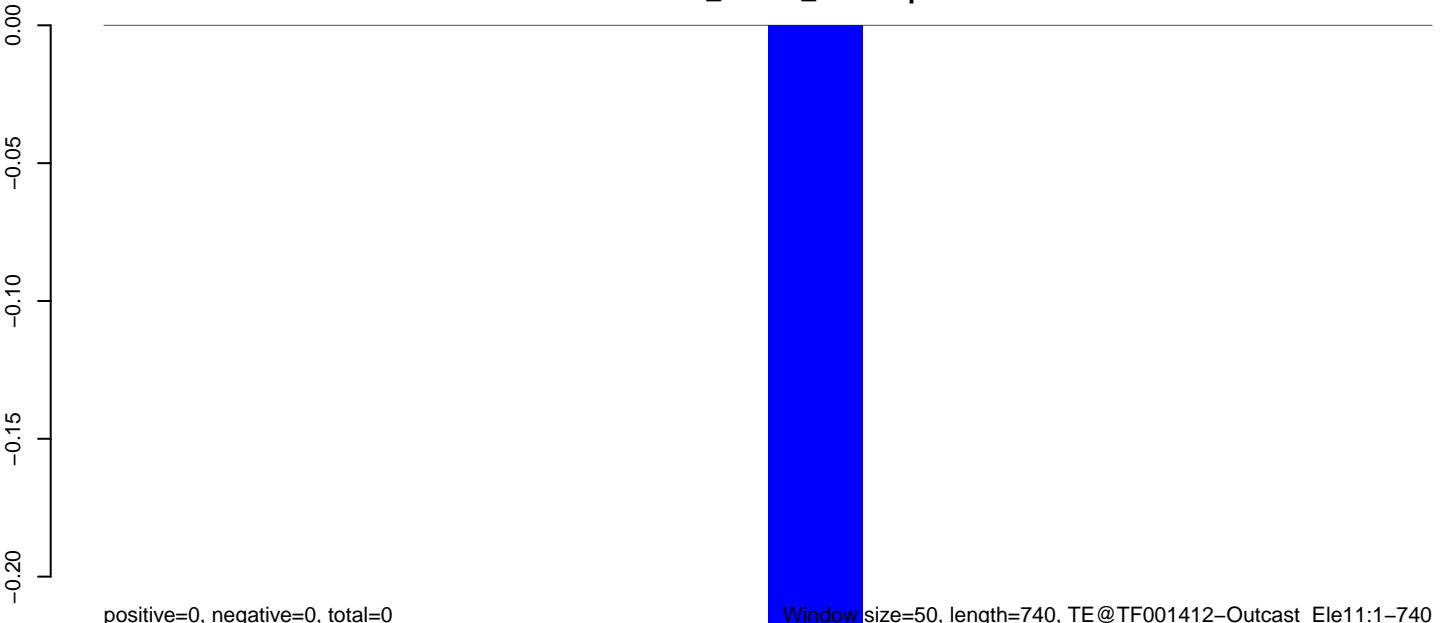
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



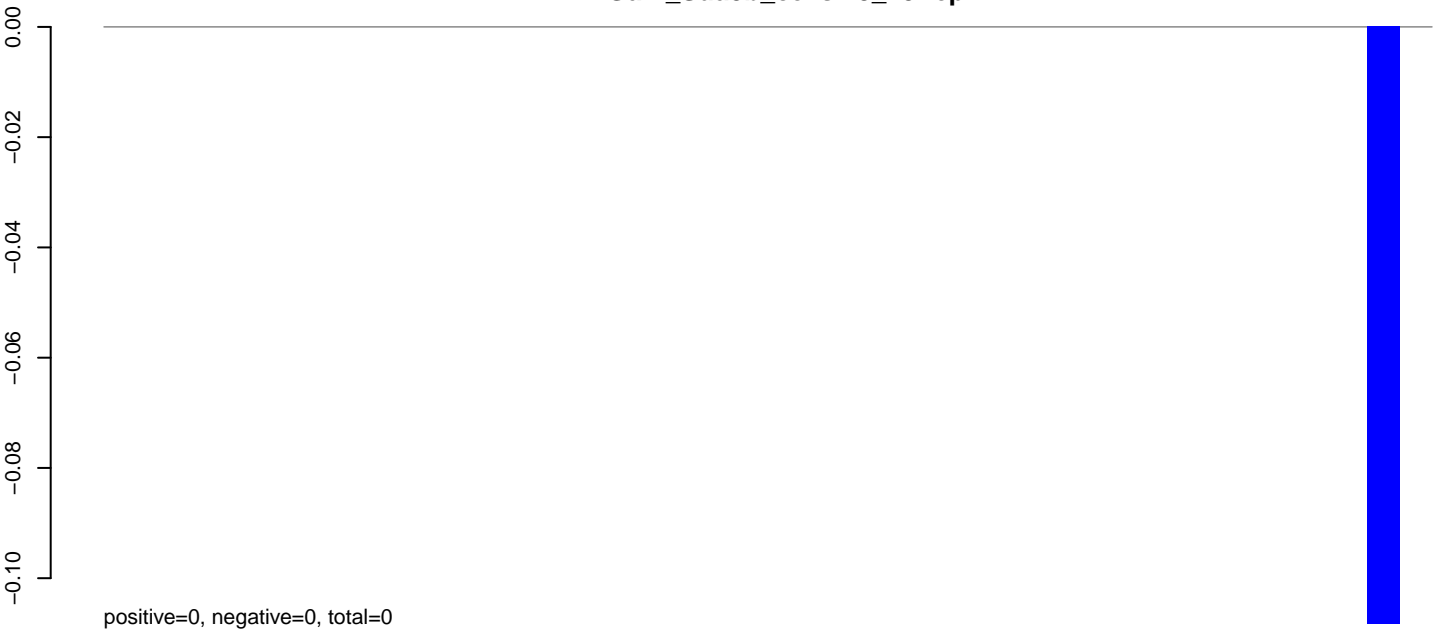
AnGam_Sua5b_cells.rep



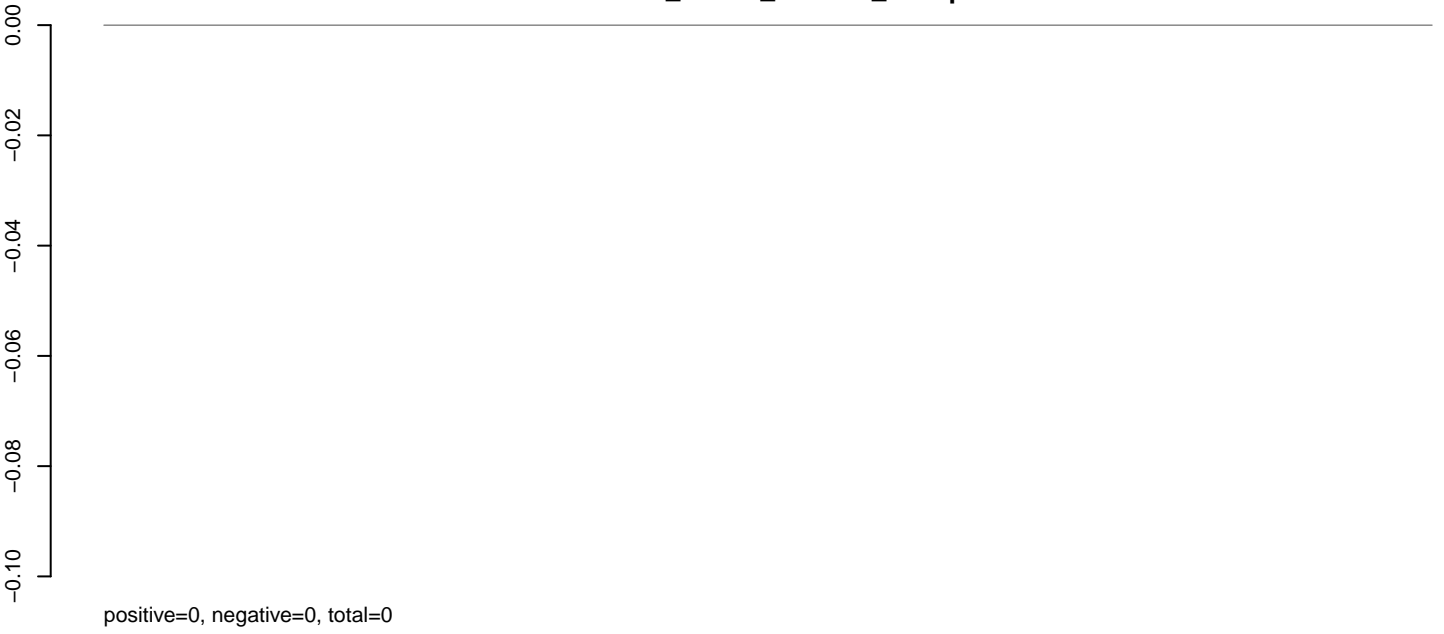
Window size=50, length=740, TE@TF001412-Outcast_Ele11:1-740

100 200 300 400 500 600 700

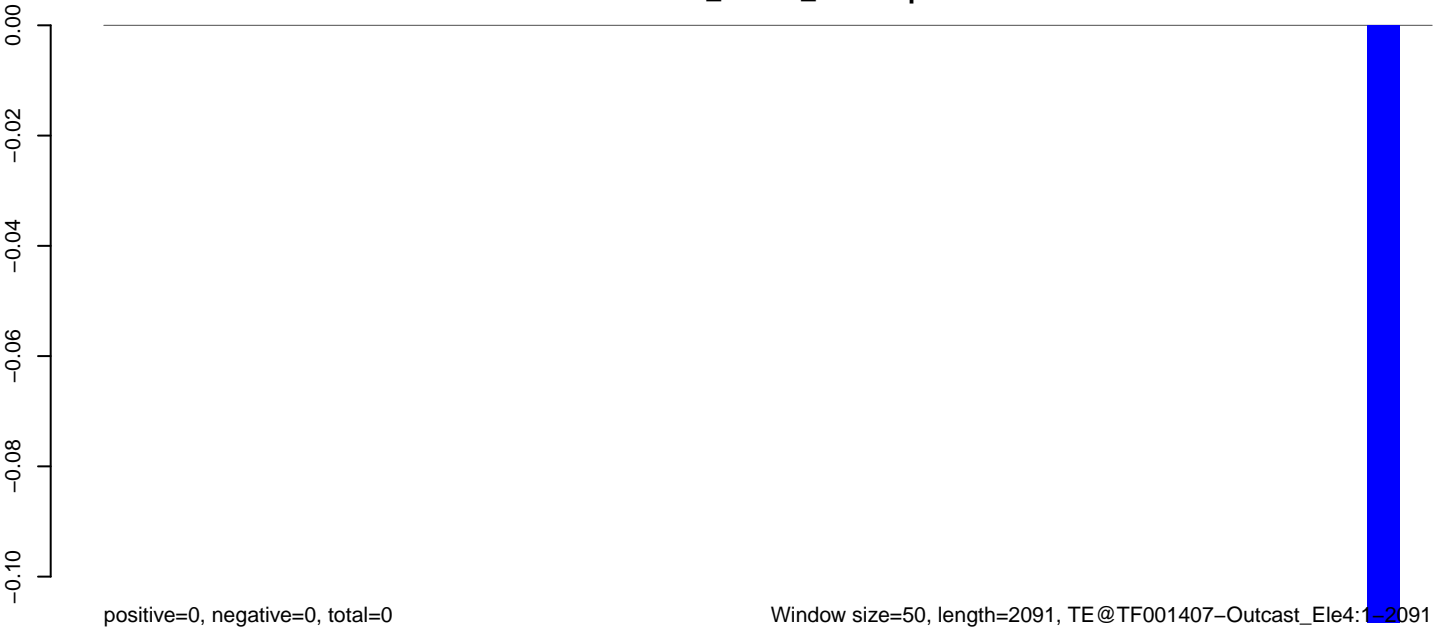
AnGam_Sua5b_cells.18_23.rep



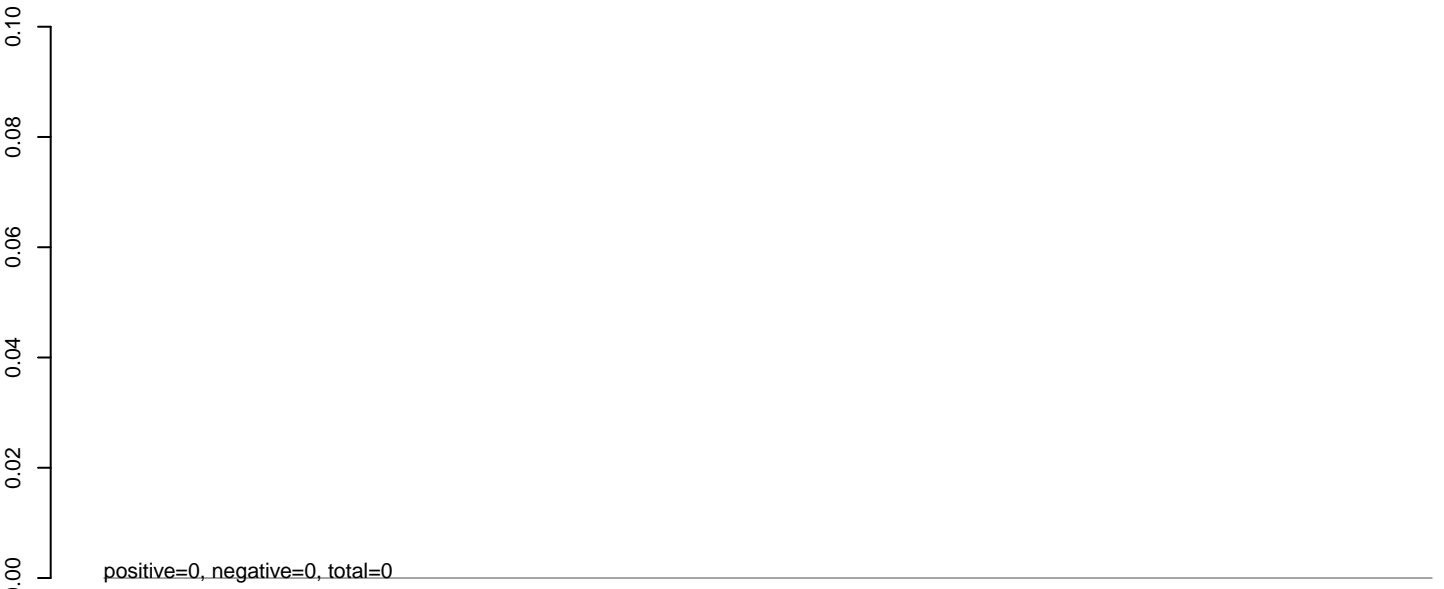
AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



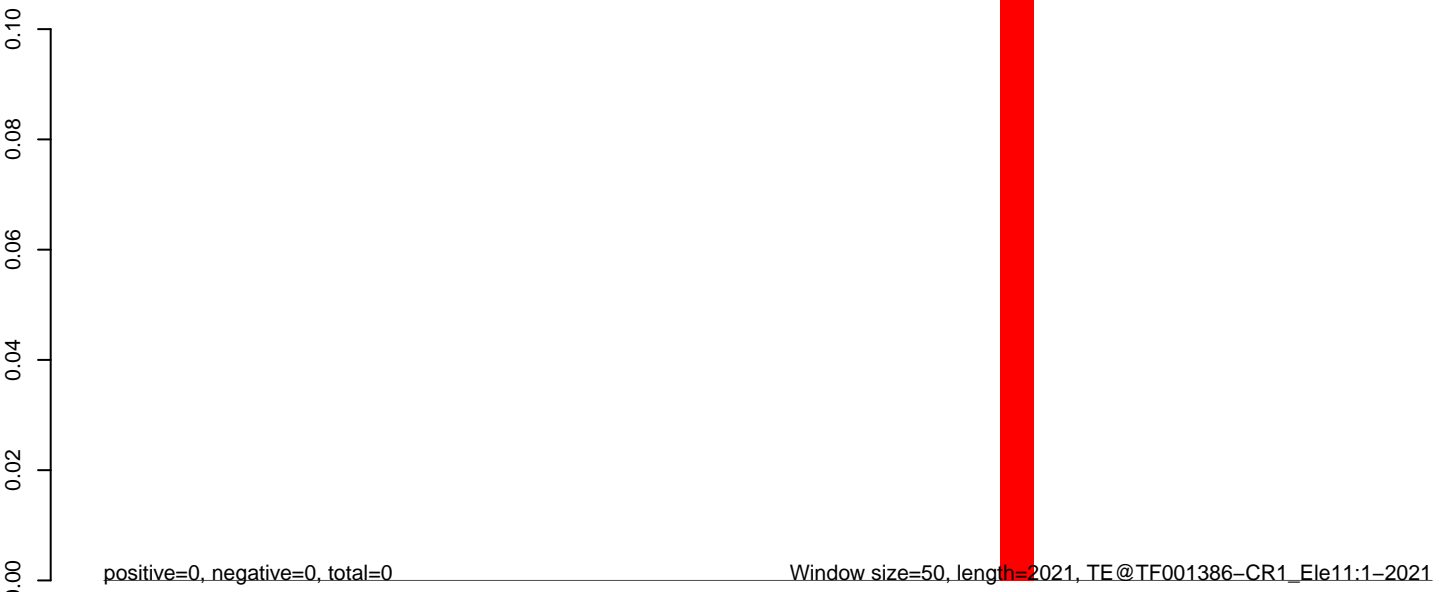
AnGam_Sua5b_cells.18_23.rep



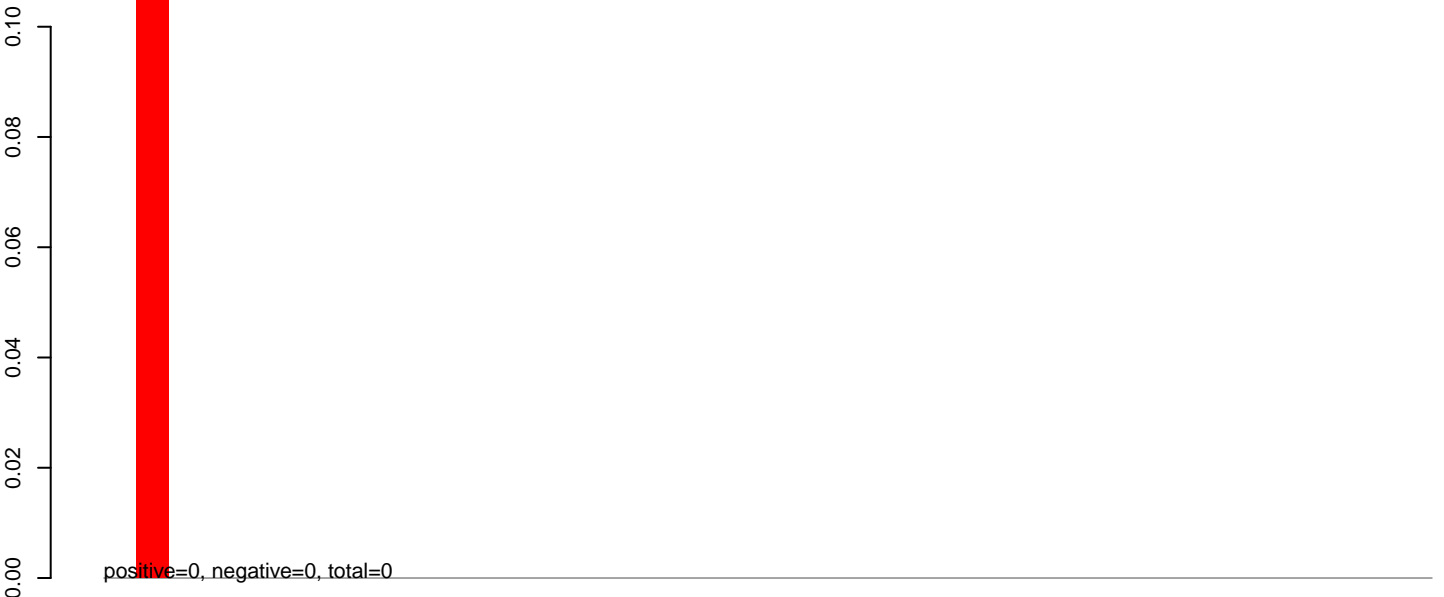
AnGam_Sua5b_cells.24_35.rep



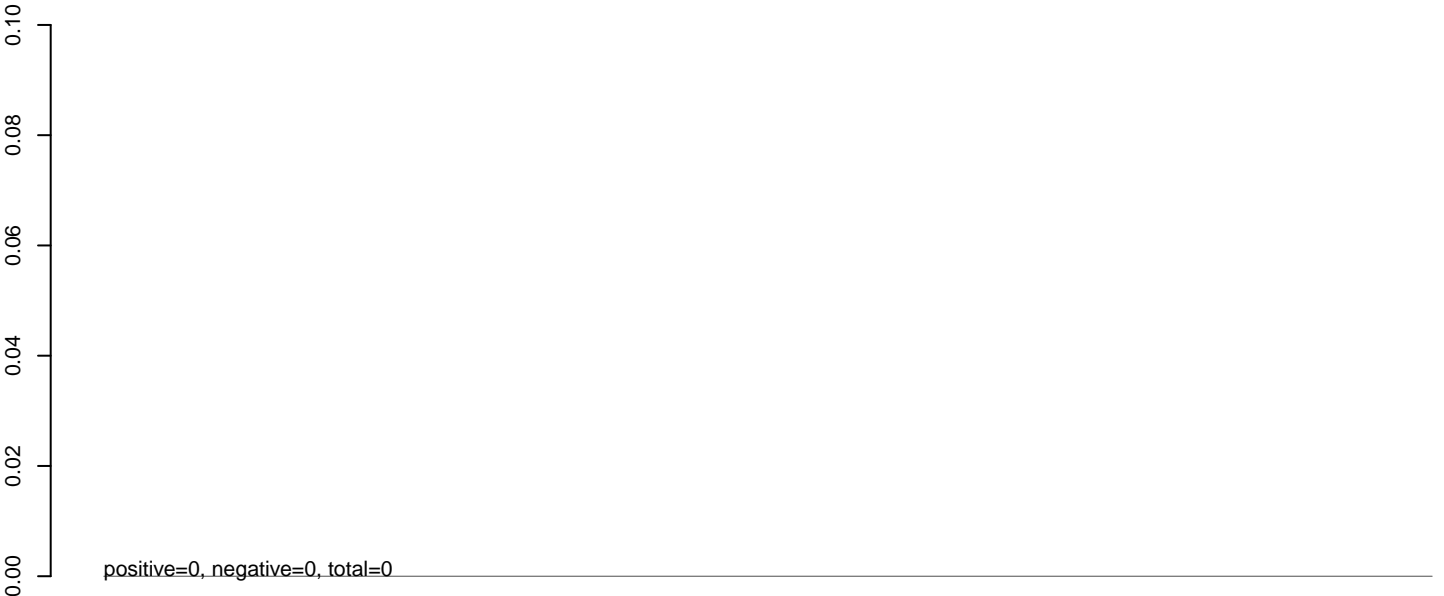
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



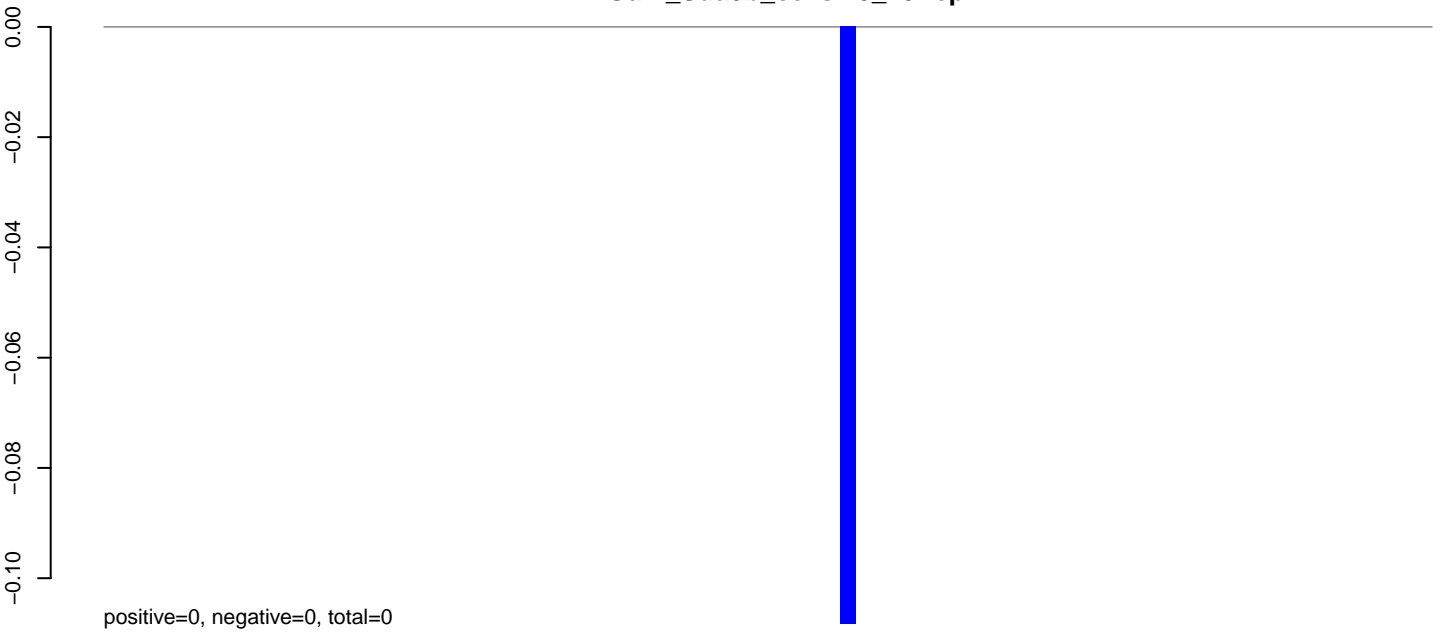
AnGam_Sua5b_cells.24_35.rep



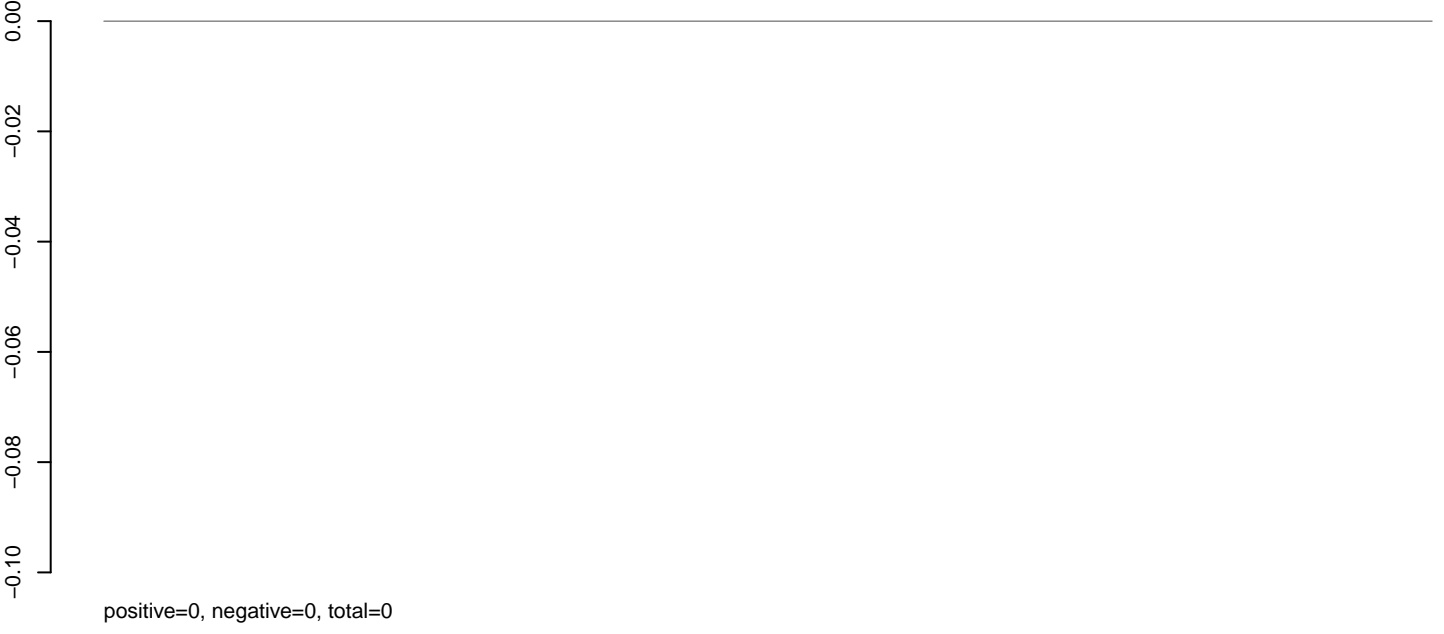
AnGam_Sua5b_cells.rep



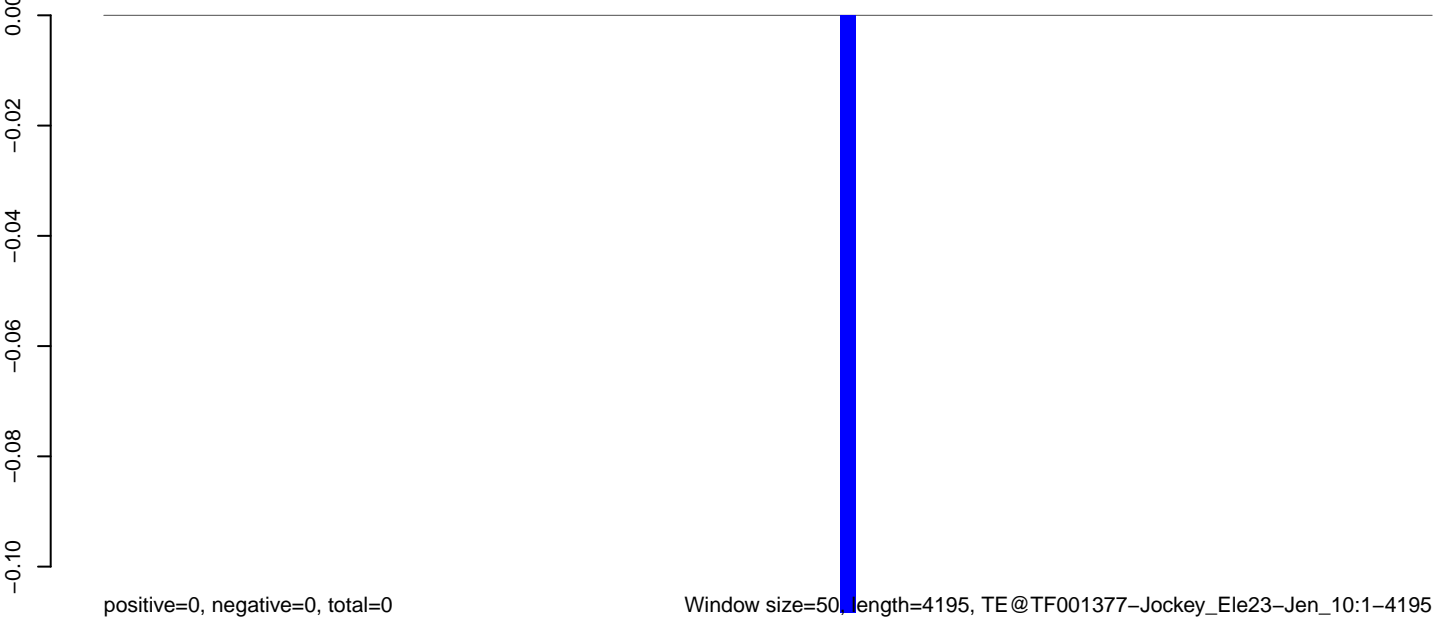
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



Window size=50, length=4195, TE@TF001377-Jockey_Ele23-Jen_10:1-4195

0 1000 2000 3000 4000

AnGam_Sua5b_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.rep

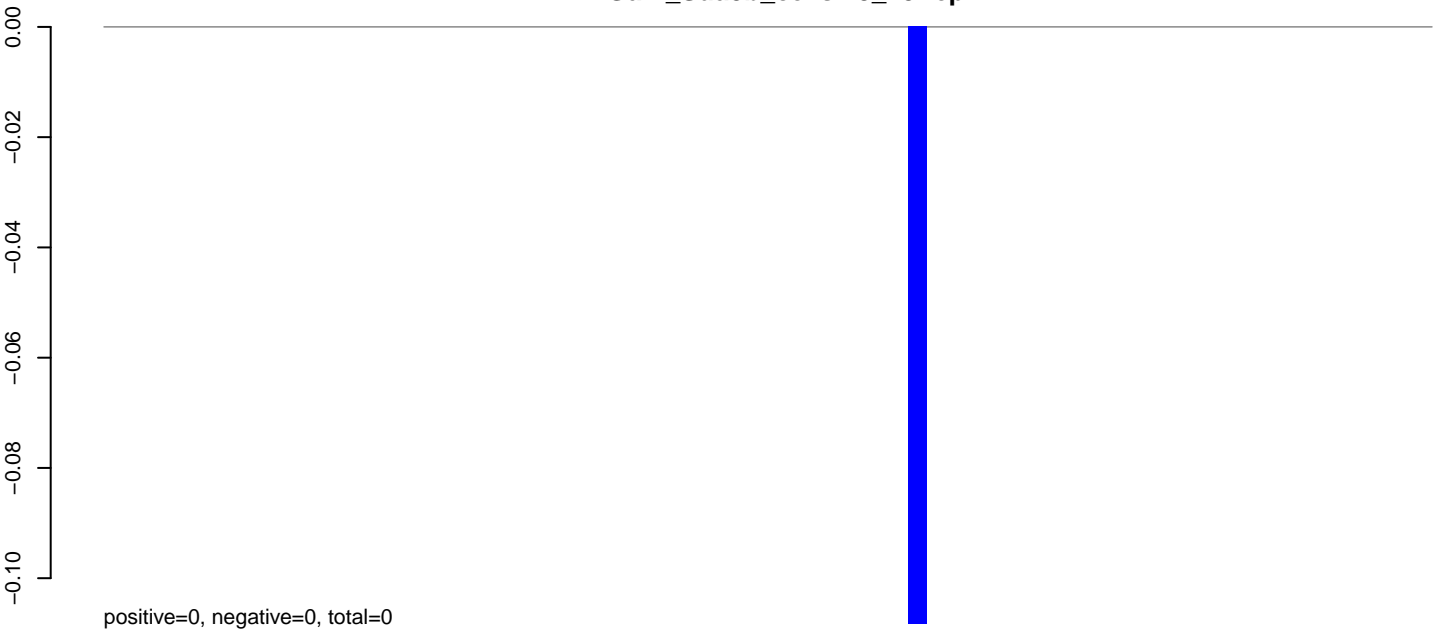
0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

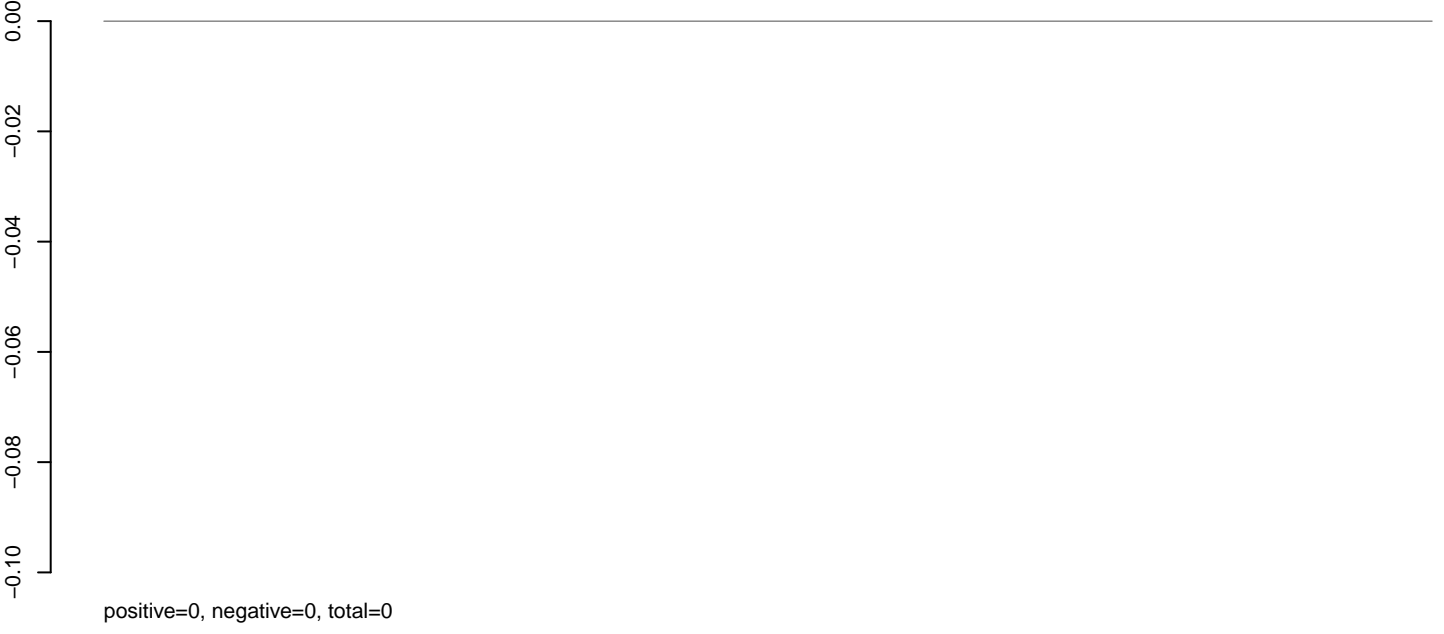
Window size=50, length=3735, TE@TF001372-Jockey_Ele18-Jen_5:1-3735

0 1000 2000 3000

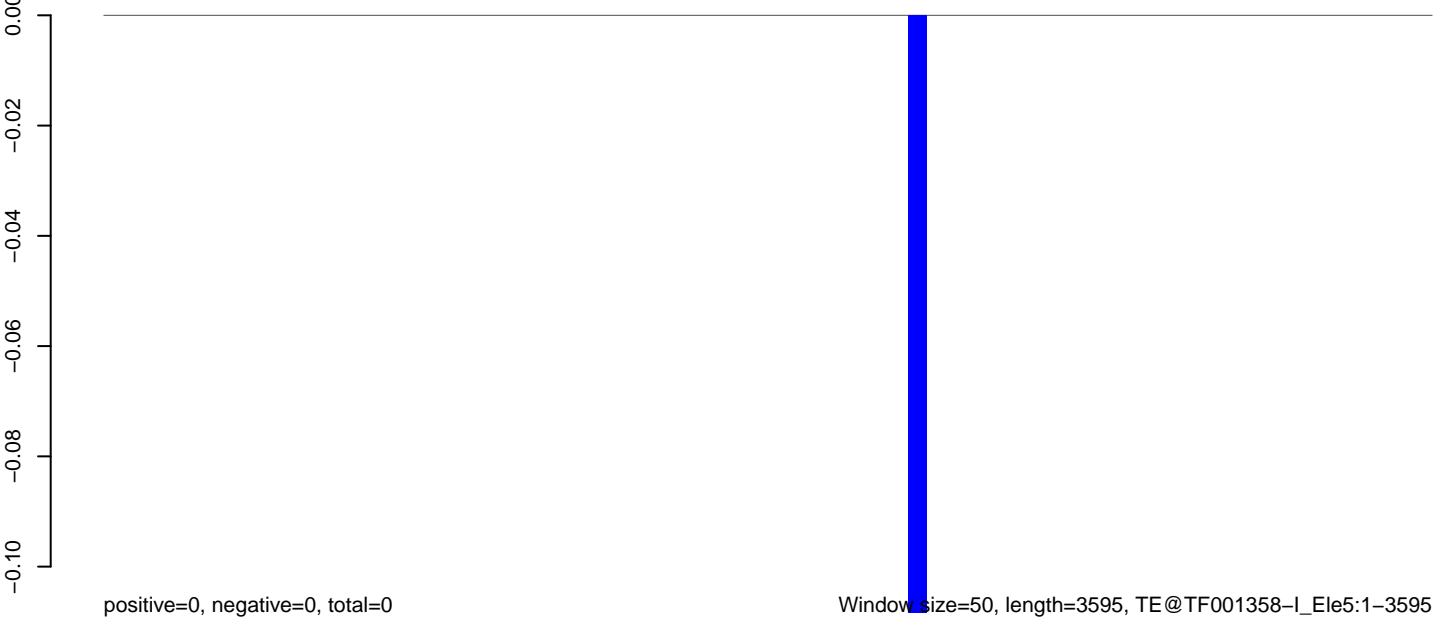
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

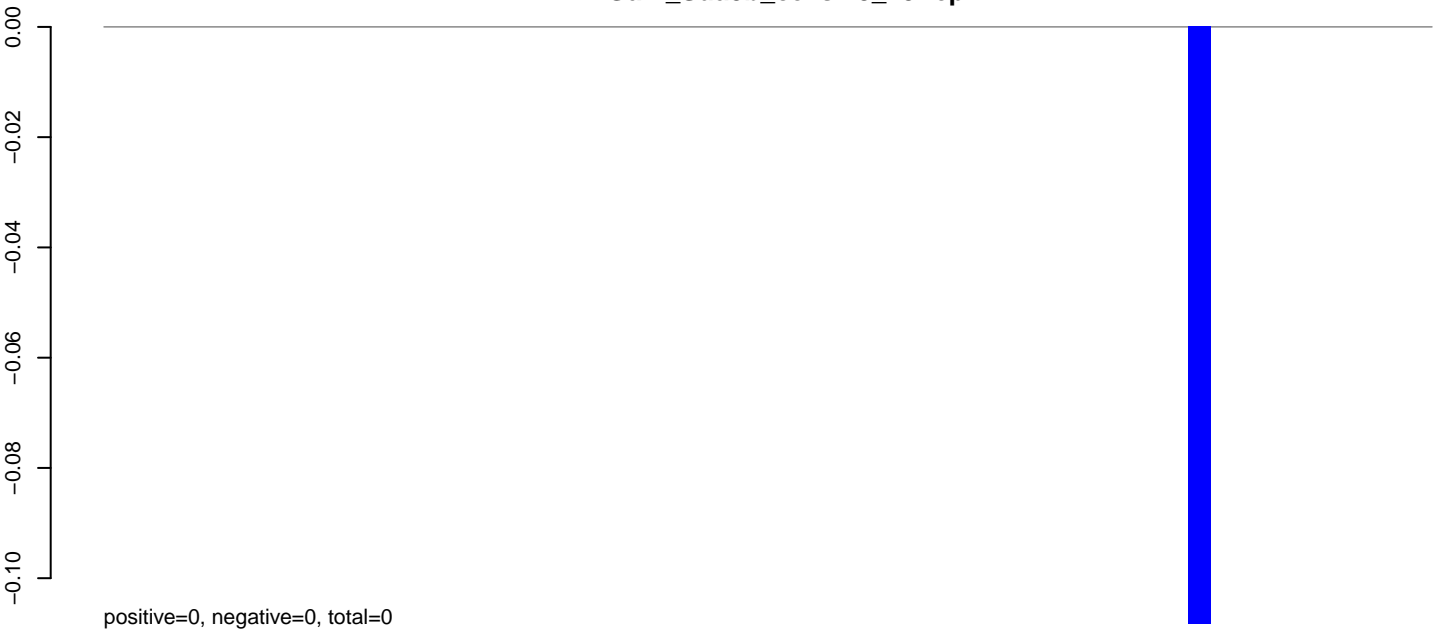


AnGam_Sua5b_cells.rep

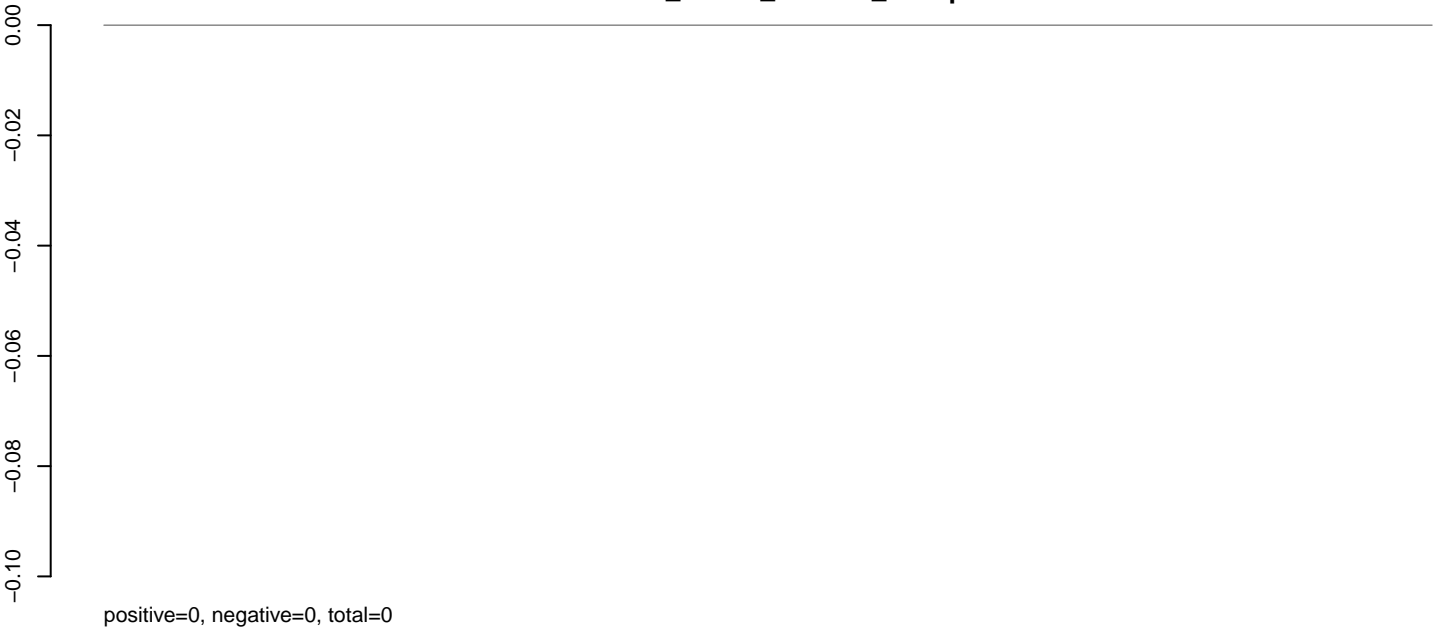


0 500 1000 1500 2000 2500 3000 3500

AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

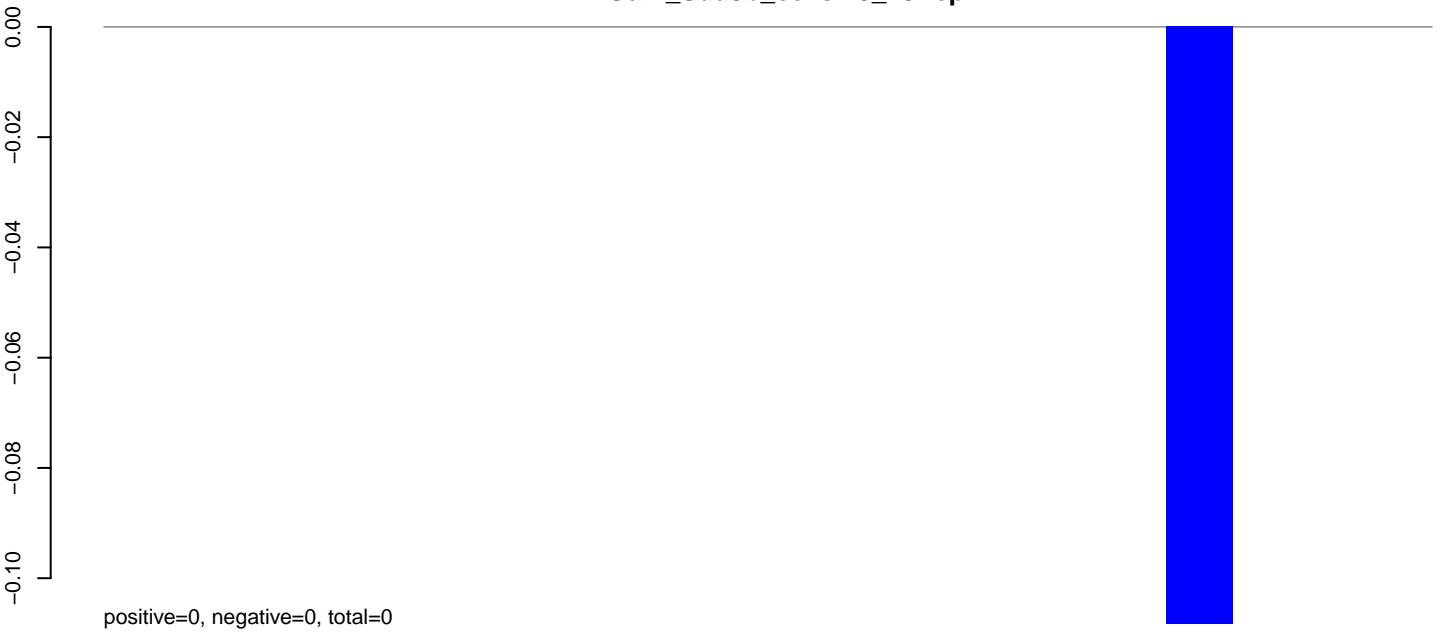


AnGam_Sua5b_cells.rep

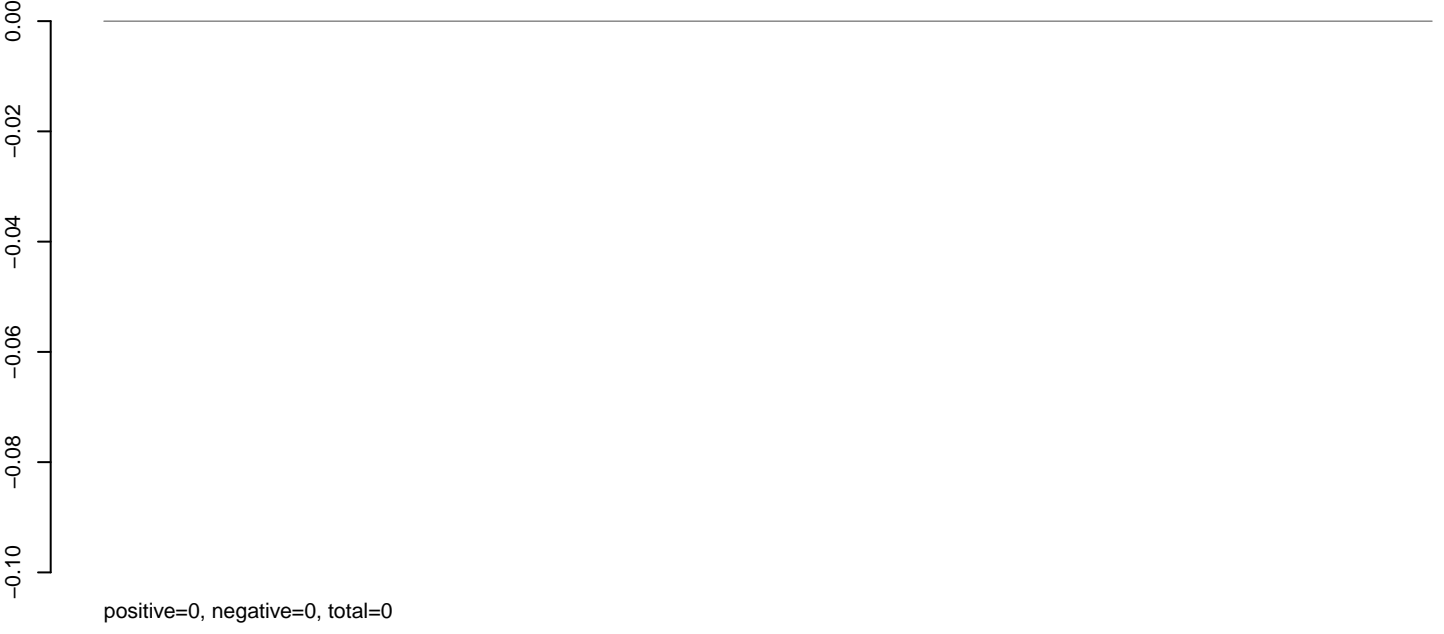


0 500 1000 1500 2000 2500 3000

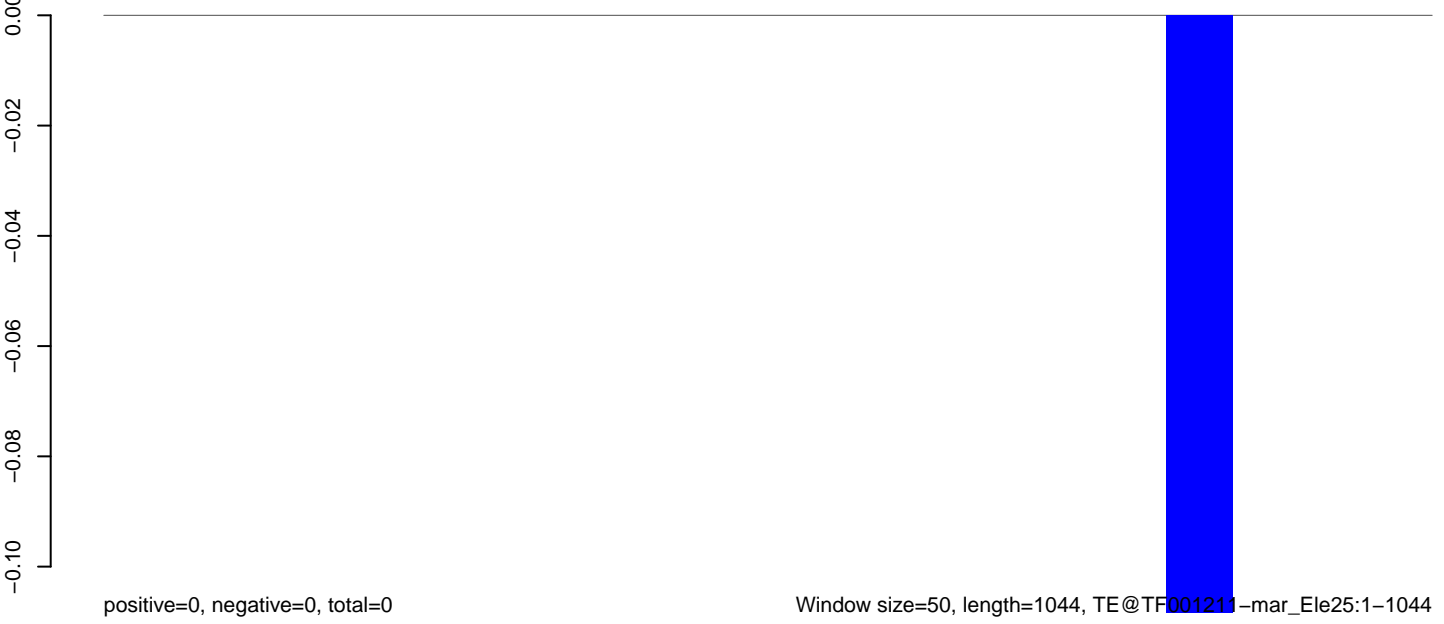
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



Window size=50, length=1044, TE@TF001211-mar_Ele25:1-1044

200 400 600 800 1000

AnGam_Sua5b_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0



AnGam_Sua5b_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0



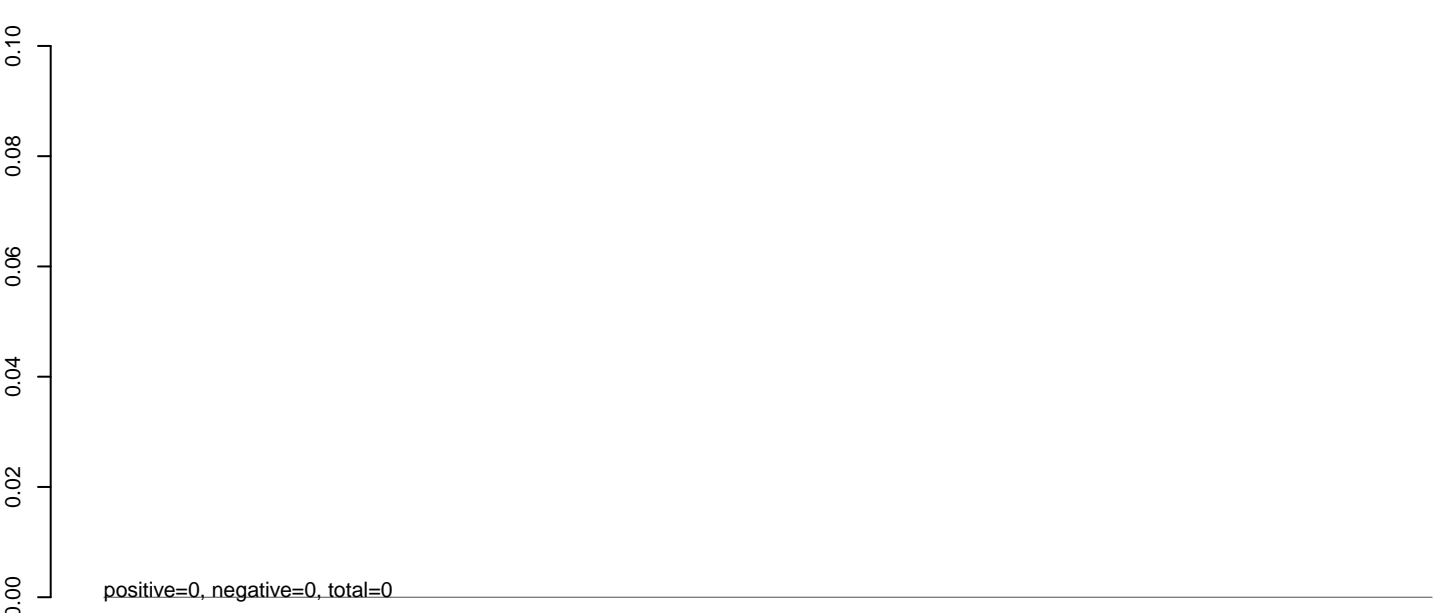
Window size=50, length=780, TE@TF001209-mar_Ele1:1-780

200 400 600 800

AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



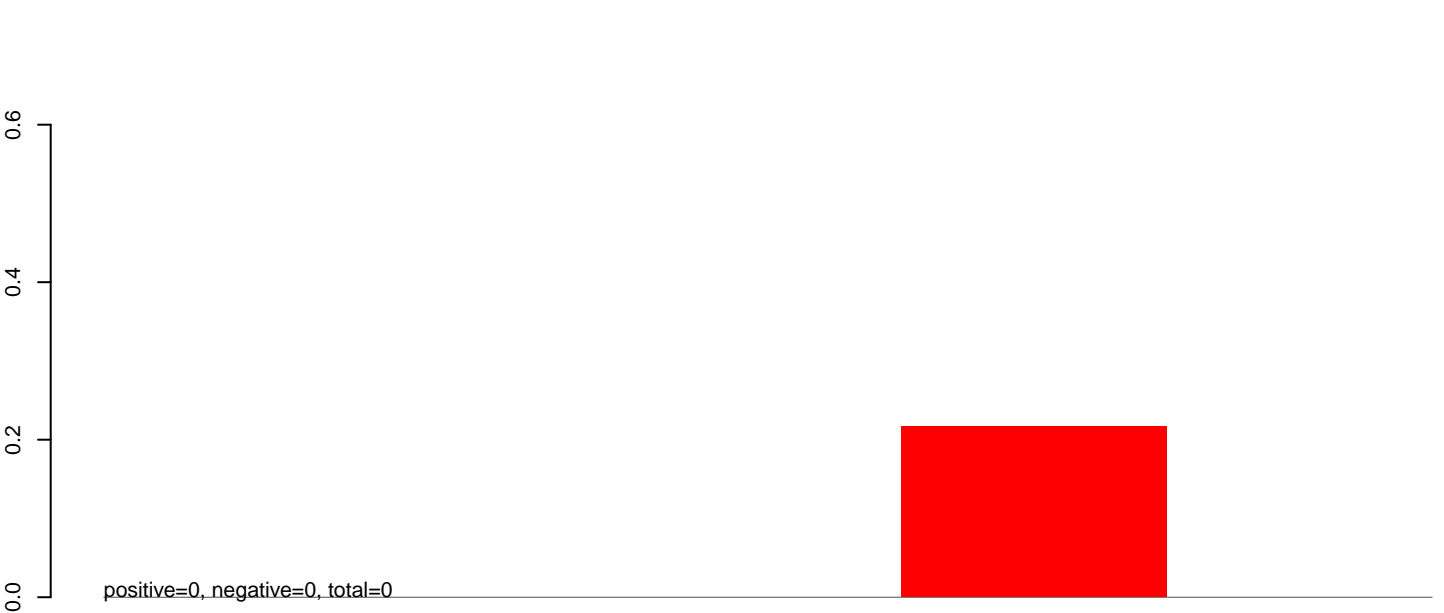
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



50 100 150 200 250 300

AnGam_Sua5b_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.rep

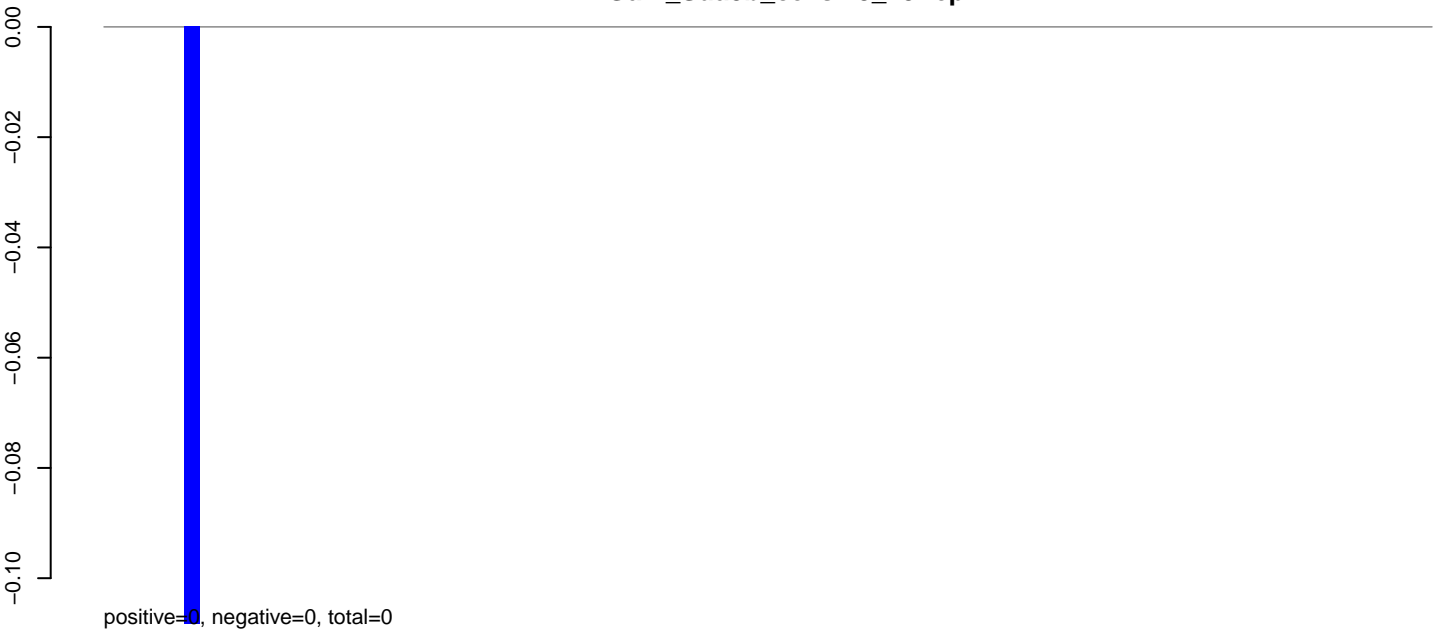
0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

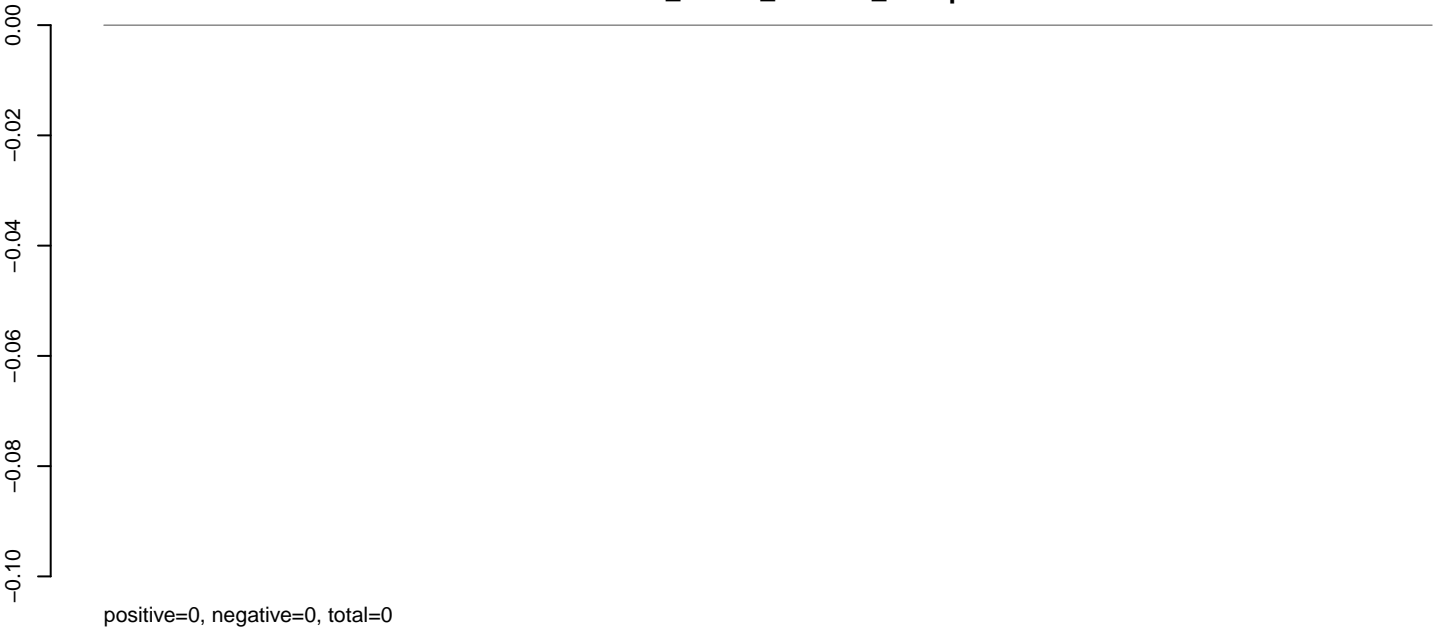
Window size=50, length=4394, TE@P3_AG:1-4394

0 1000 2000 3000 4000

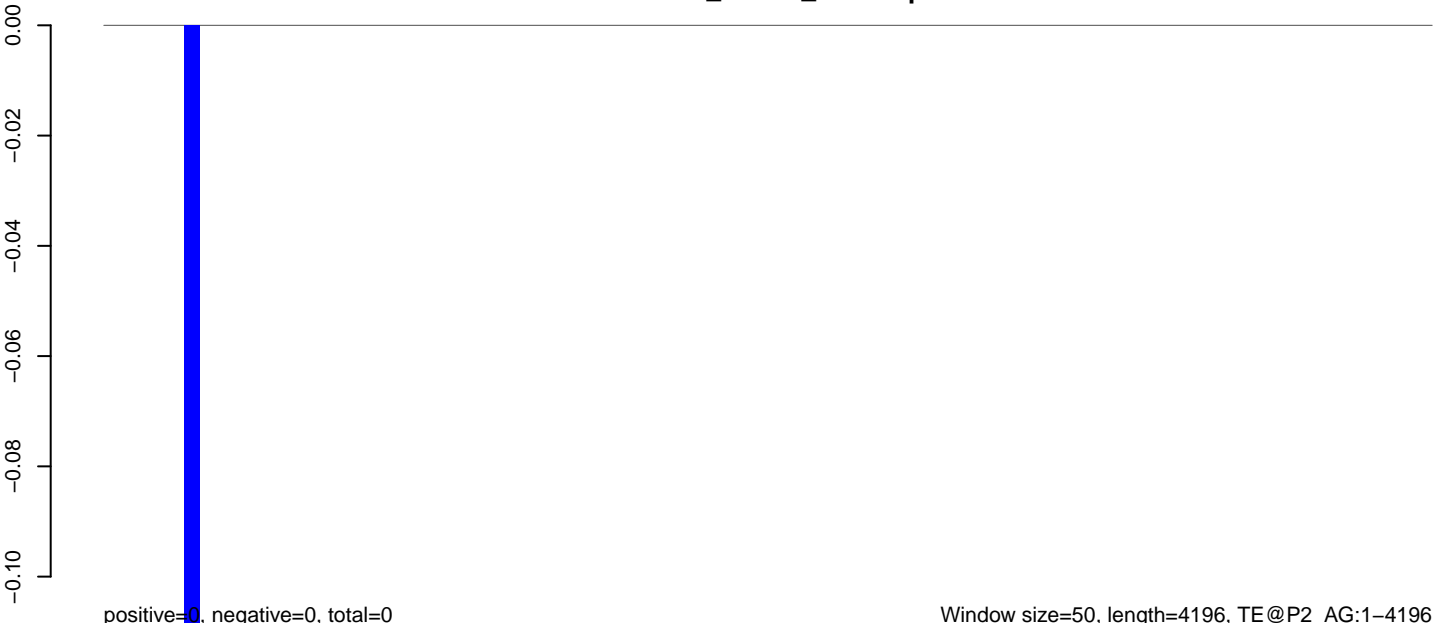
AnGam_Sua5b_cells.18_23.rep



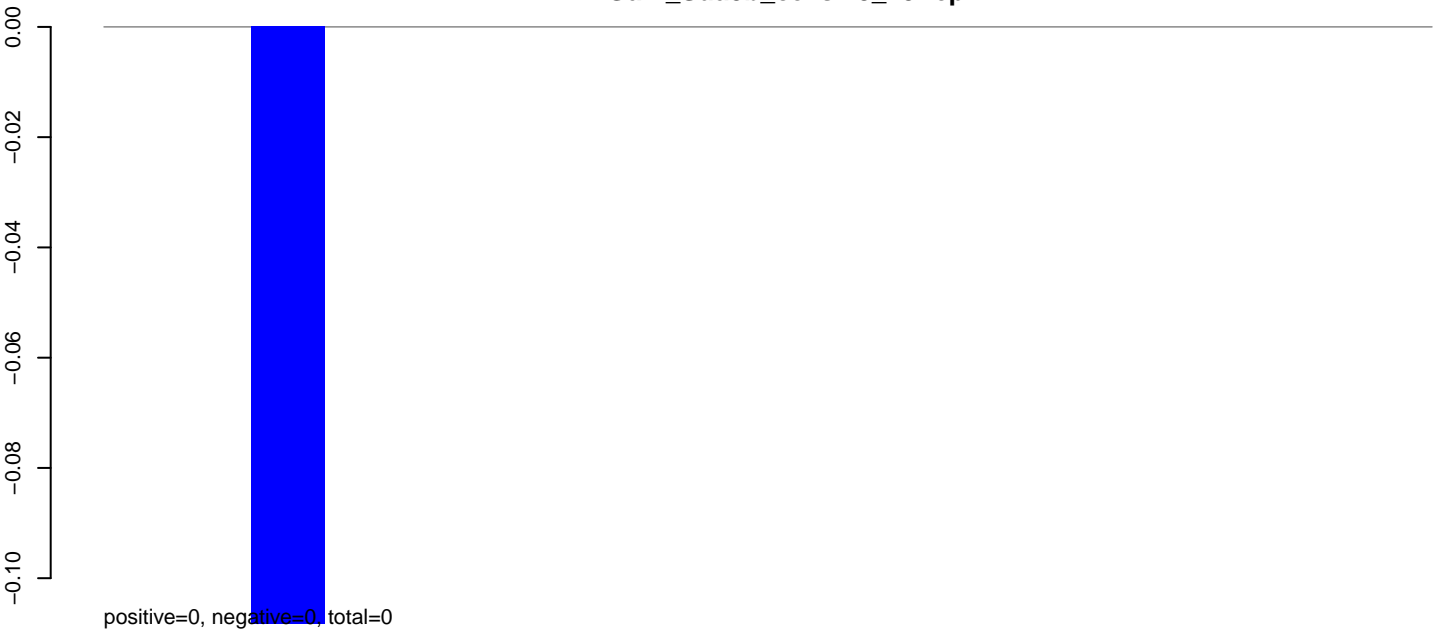
AnGam_Sua5b_cells.24_35.rep



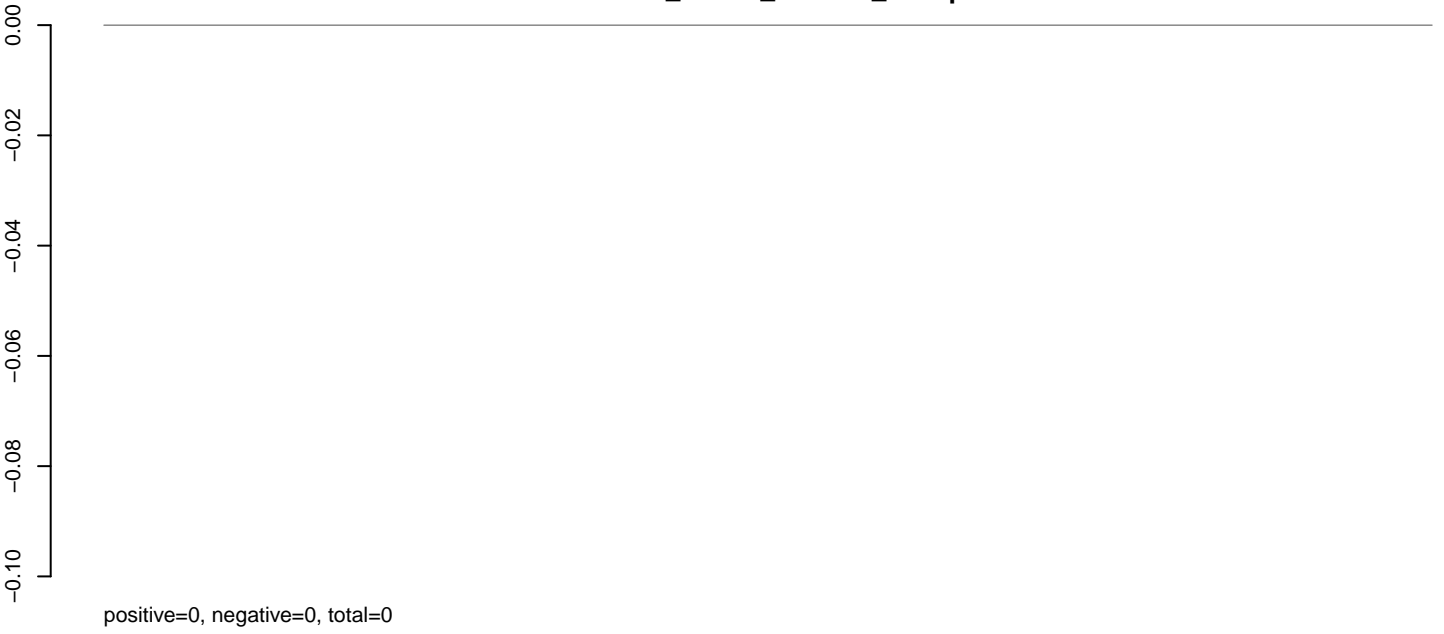
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



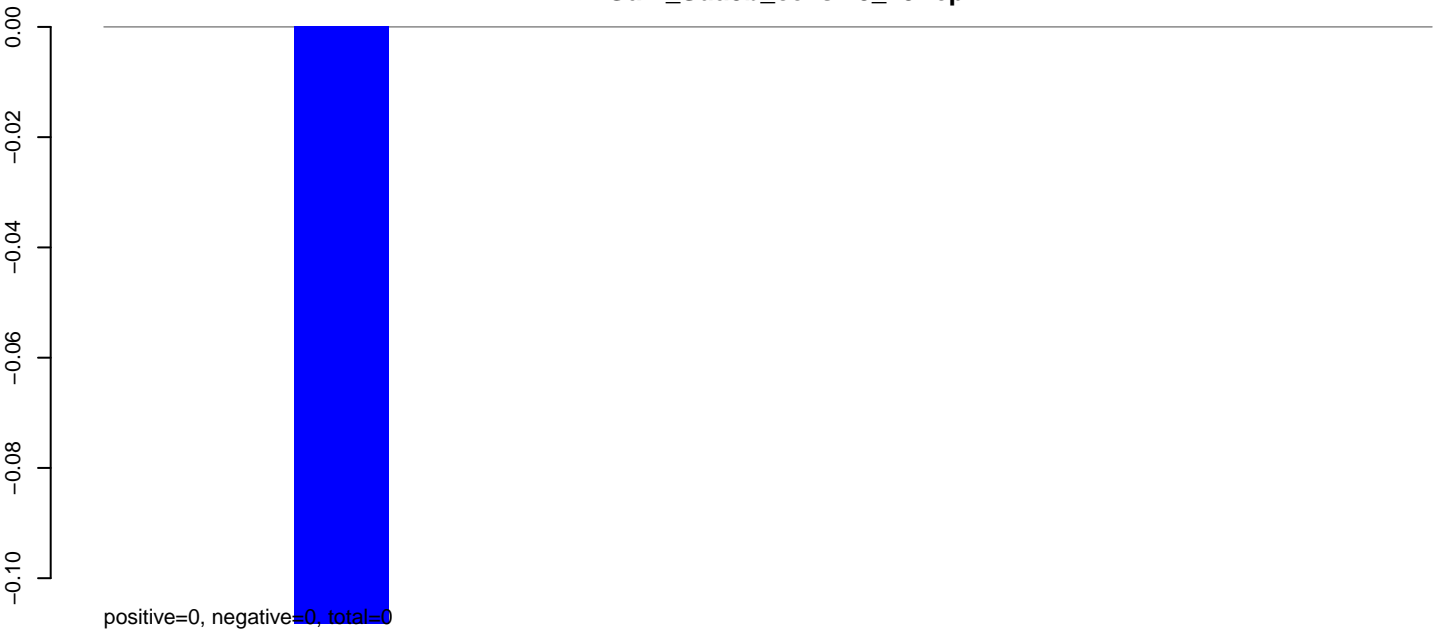
AnGam_Sua5b_cells.24_35.rep



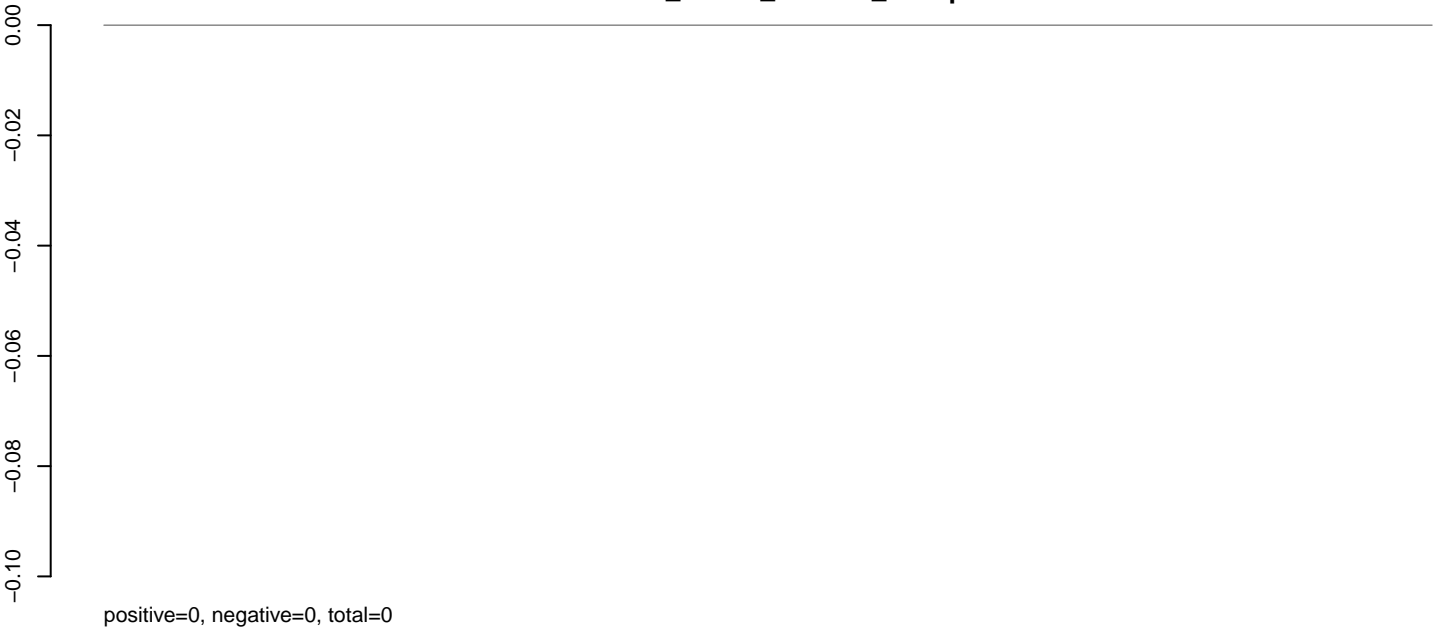
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



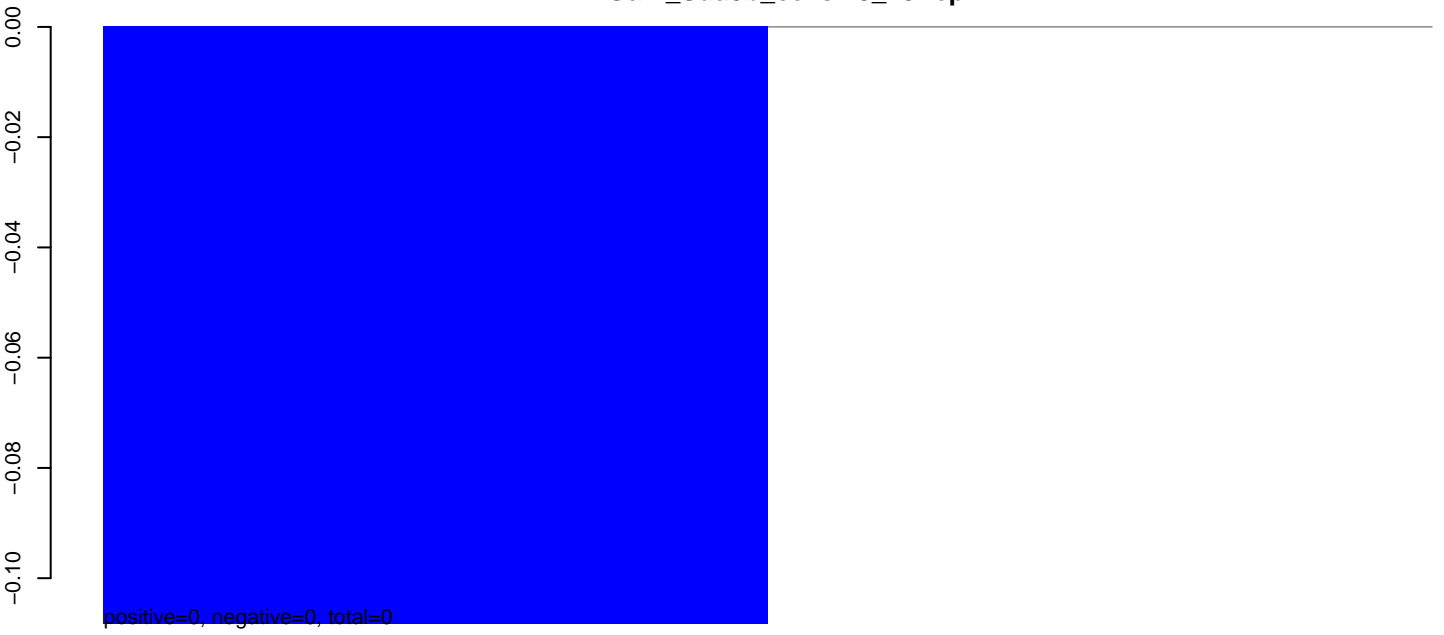
AnGam_Sua5b_cells.24_35.rep



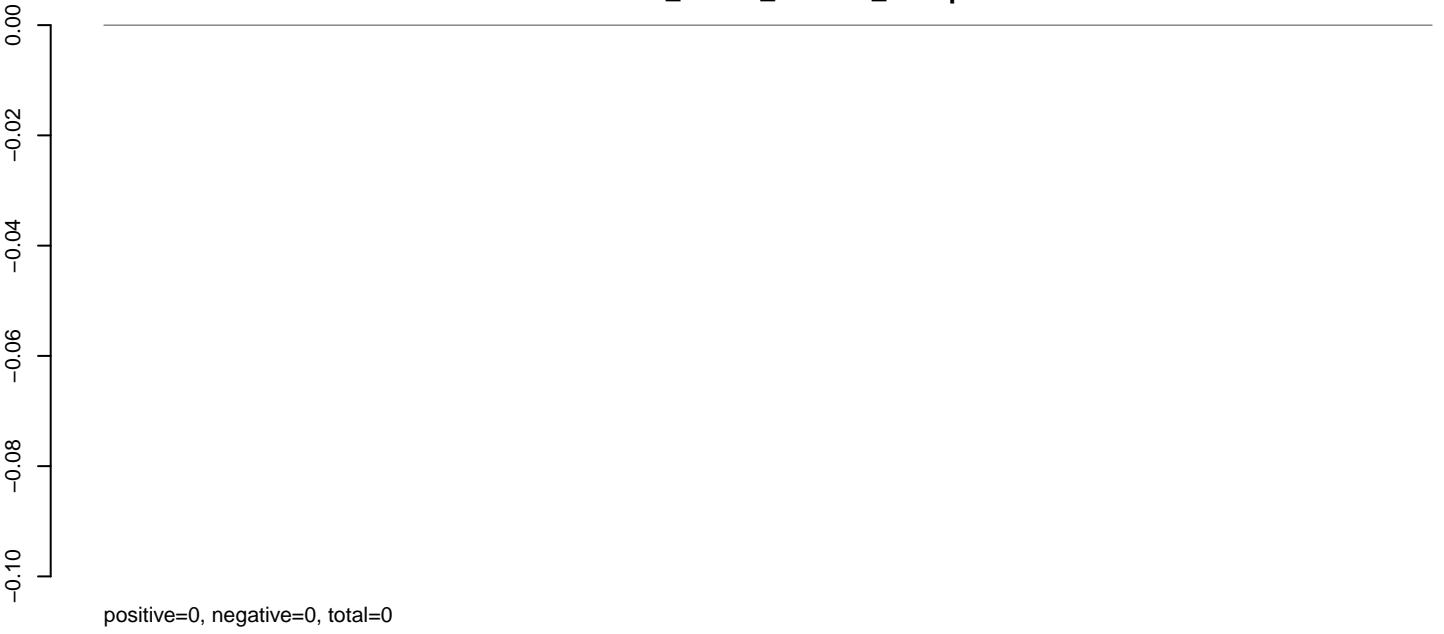
AnGam_Sua5b_cells.rep



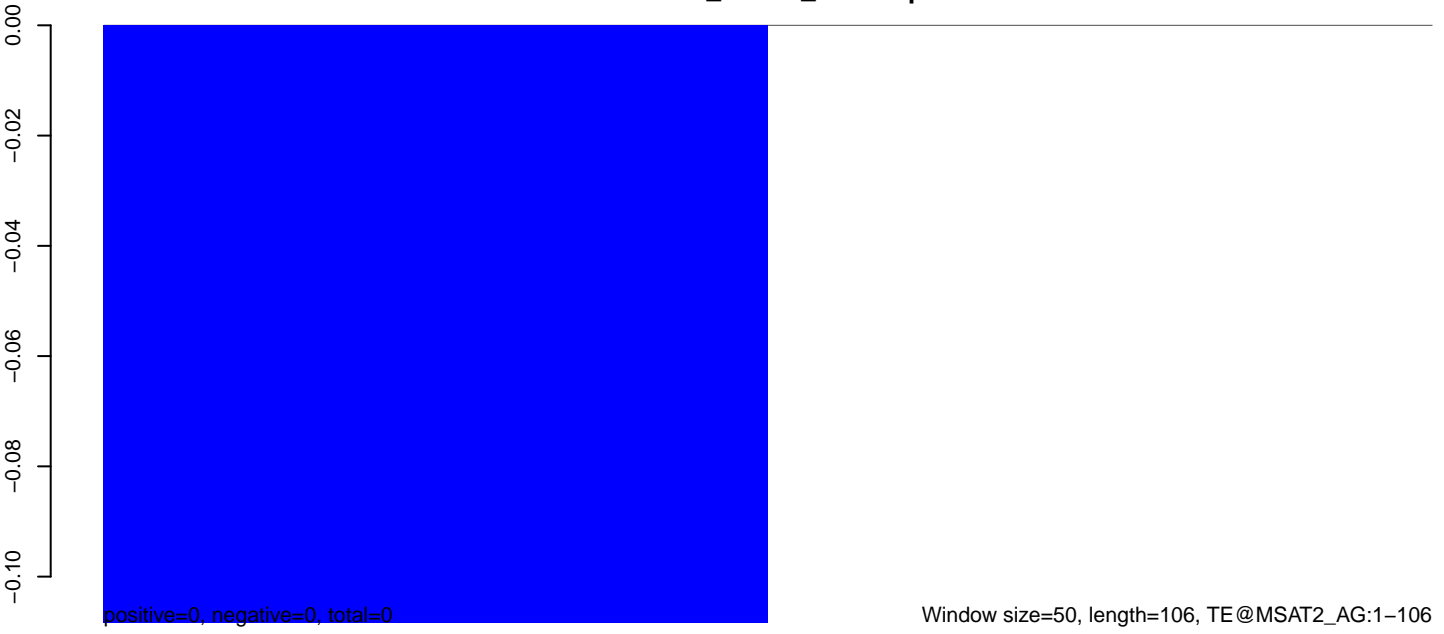
AnGam_Sua5b_cells.18_23.rep



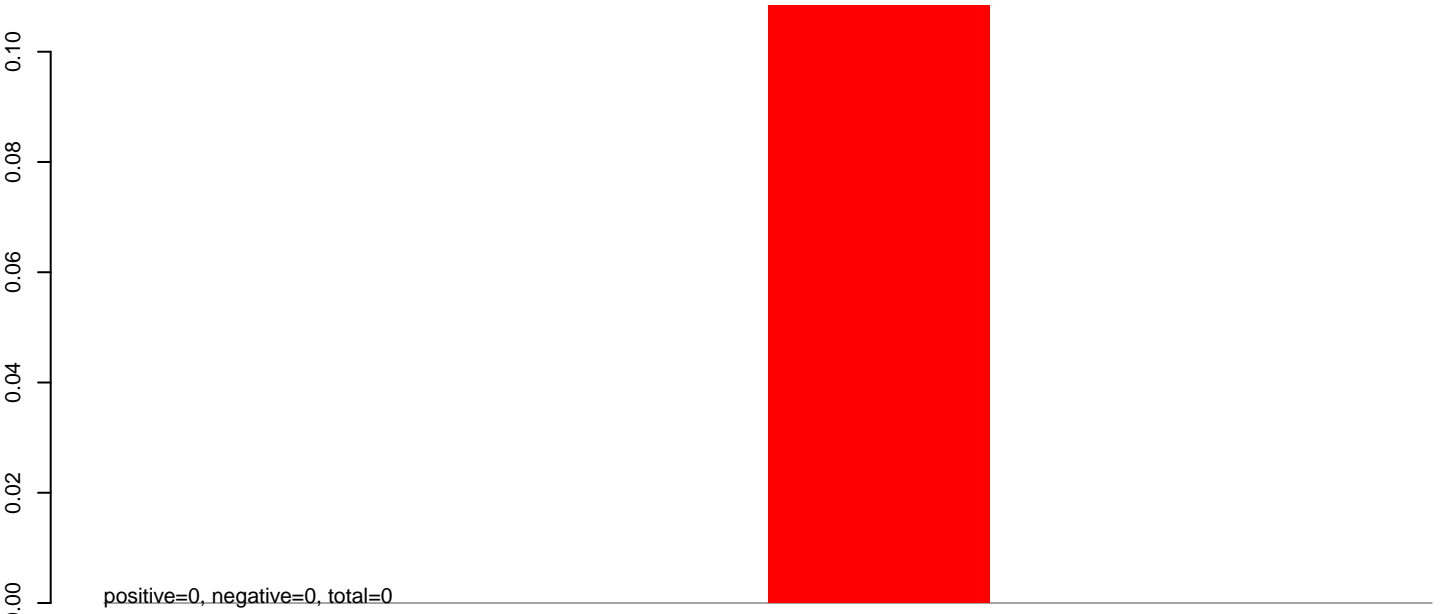
AnGam_Sua5b_cells.24_35.rep



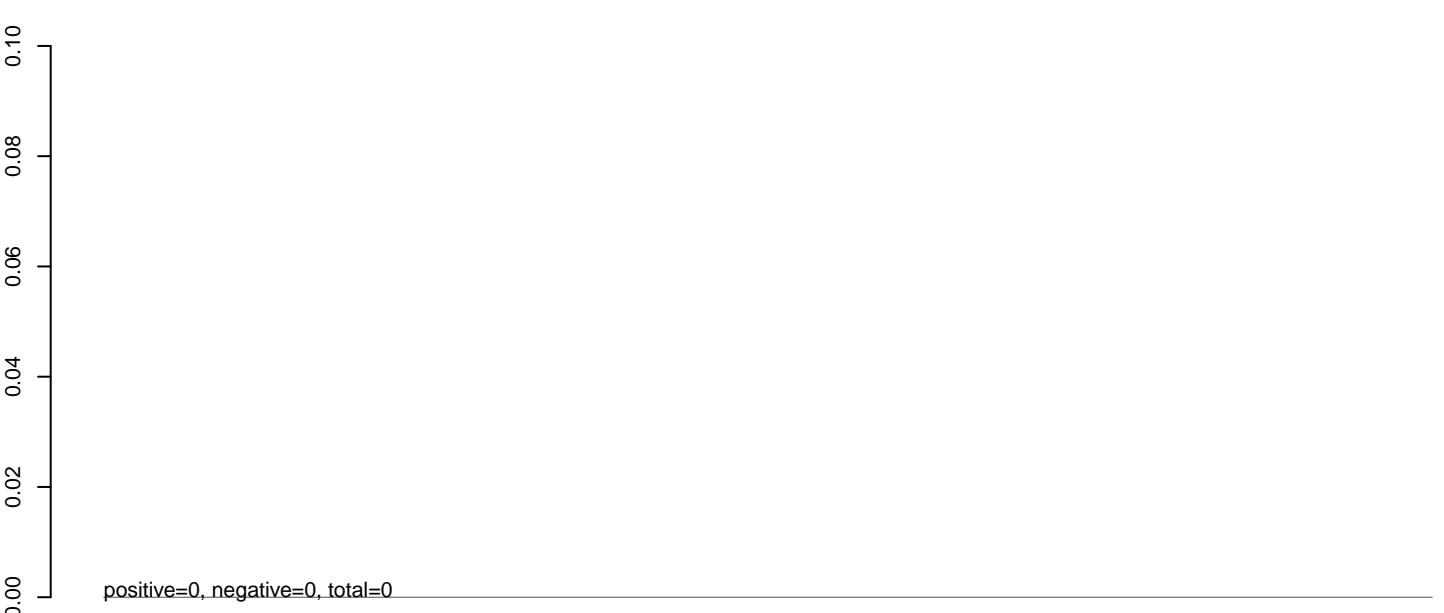
AnGam_Sua5b_cells.rep



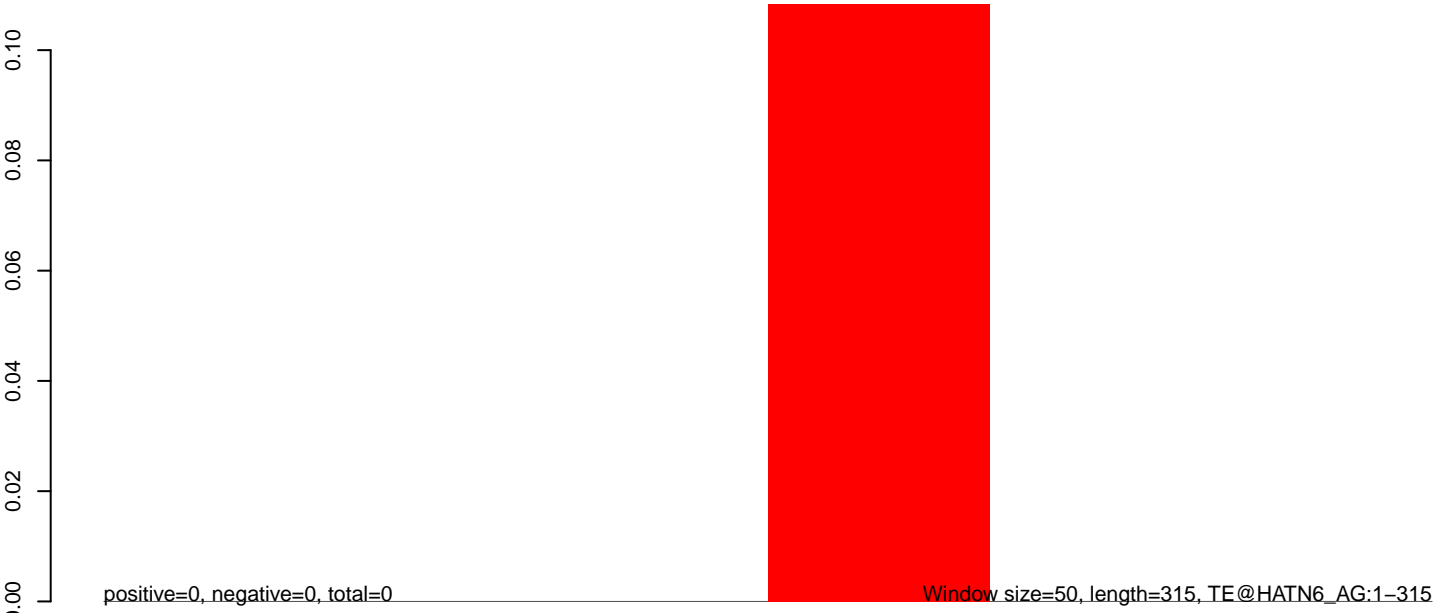
AnGam_Sua5b_cells.18_23.rep



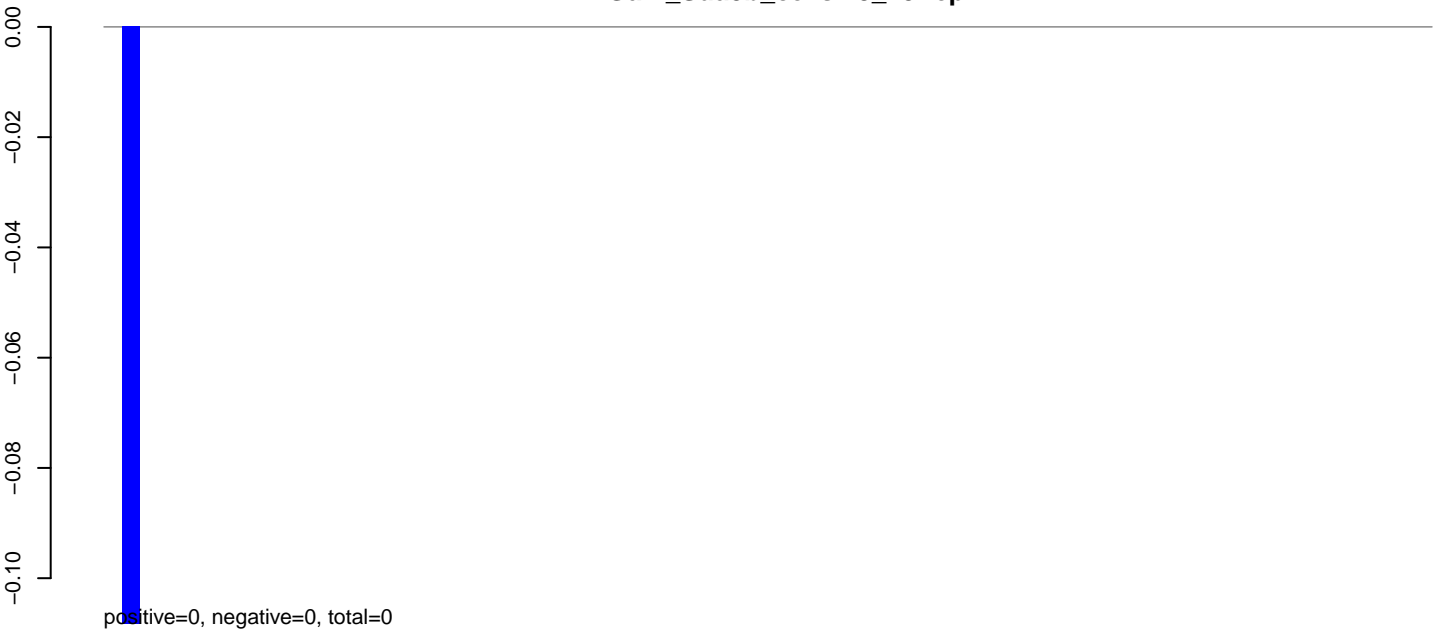
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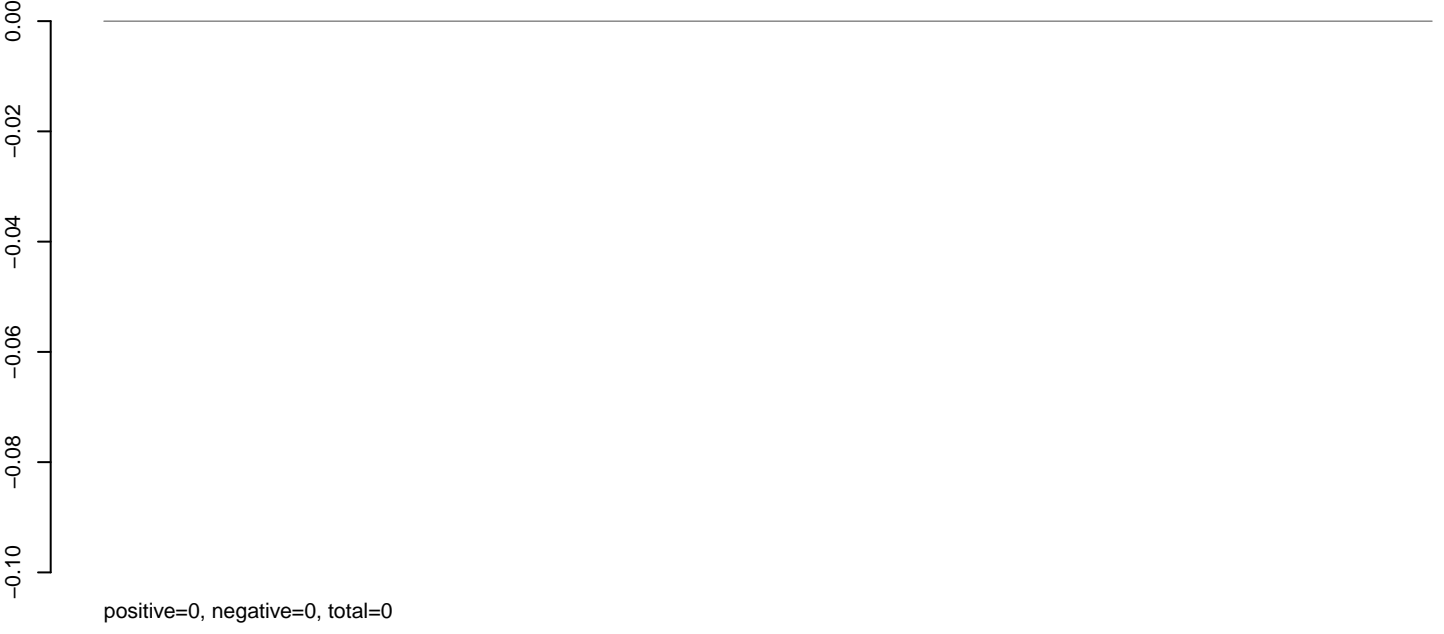
AnGam_Sua5b_cells.rep



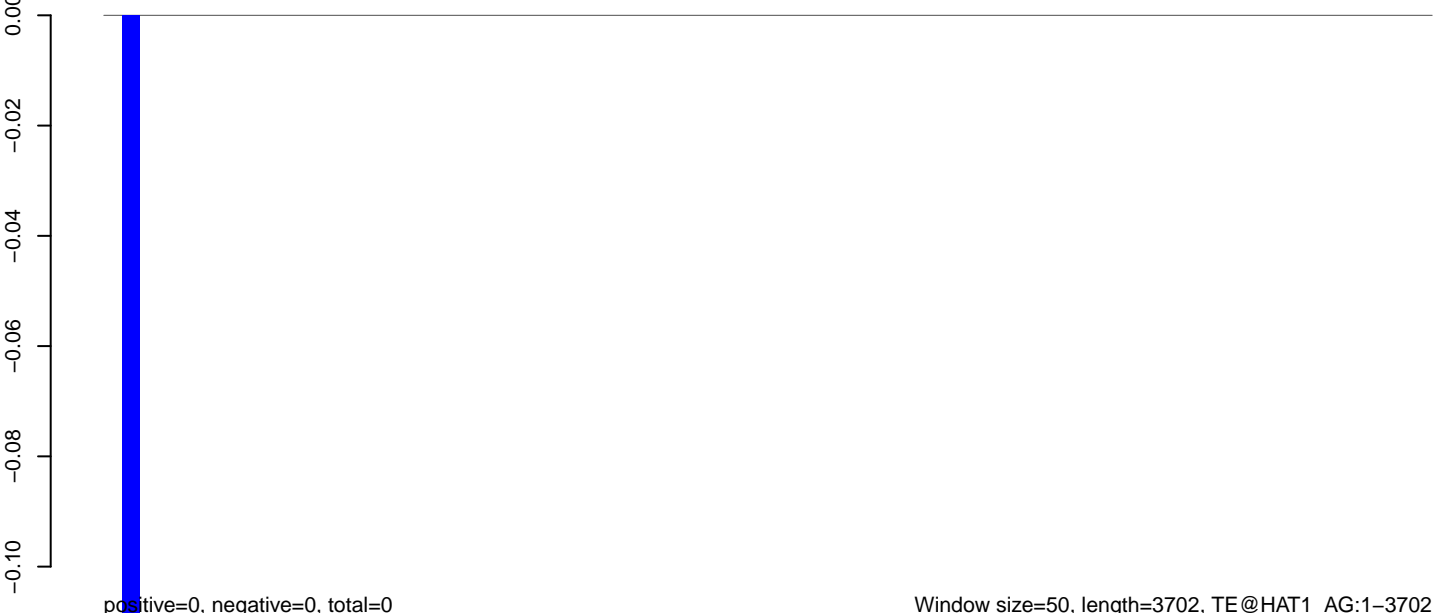
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



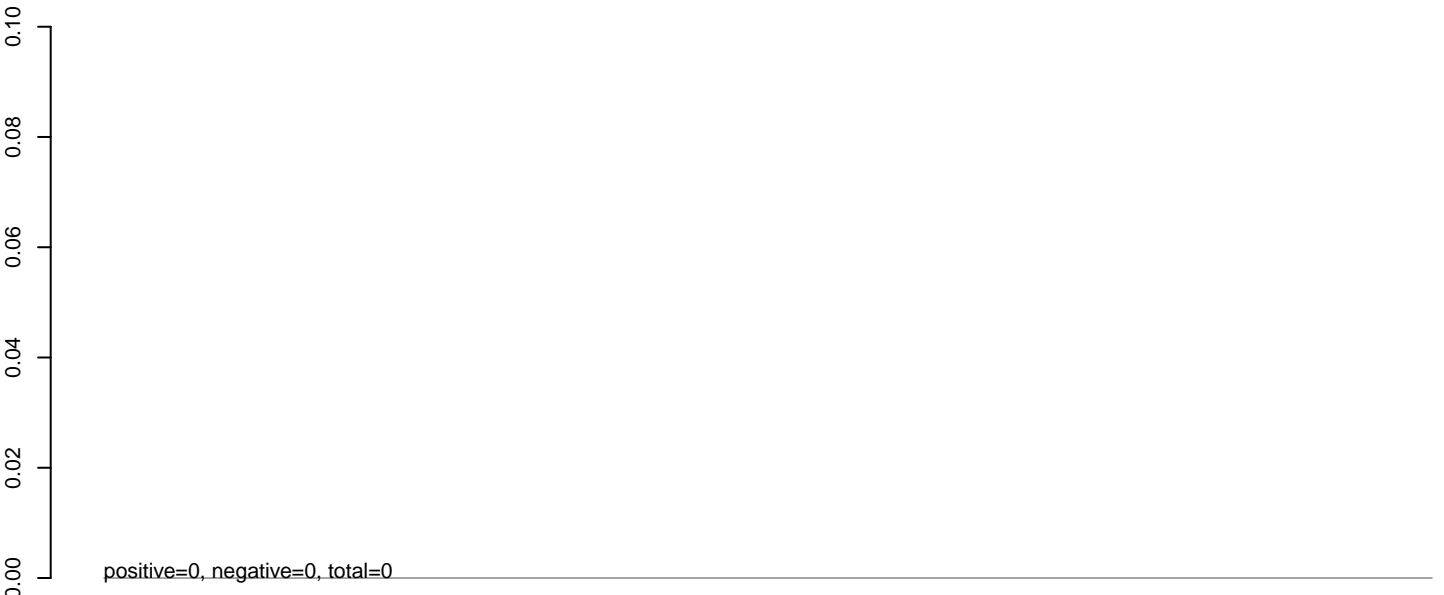
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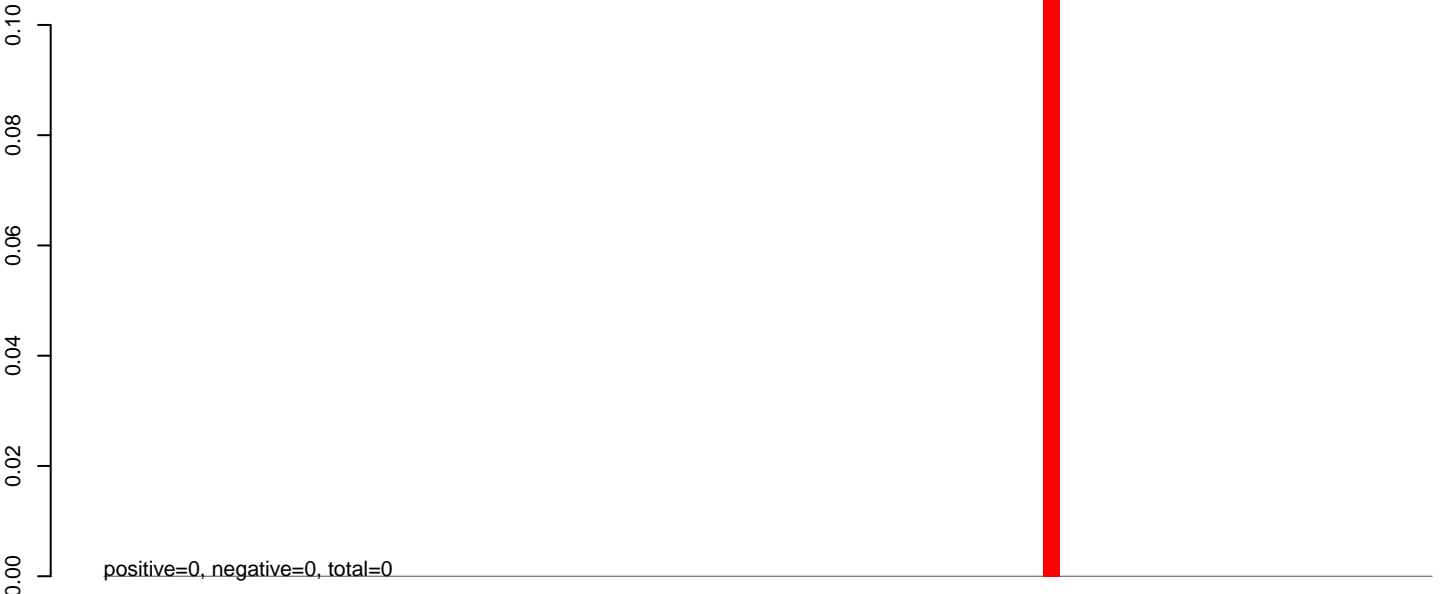
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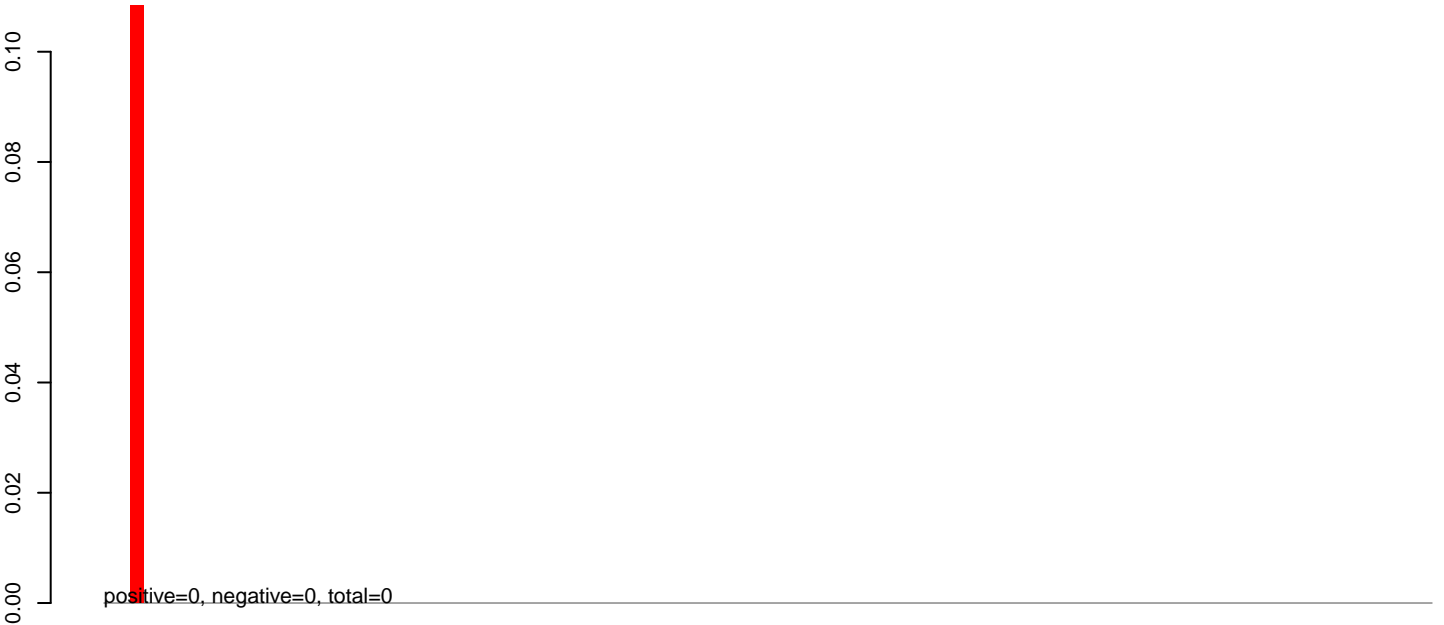
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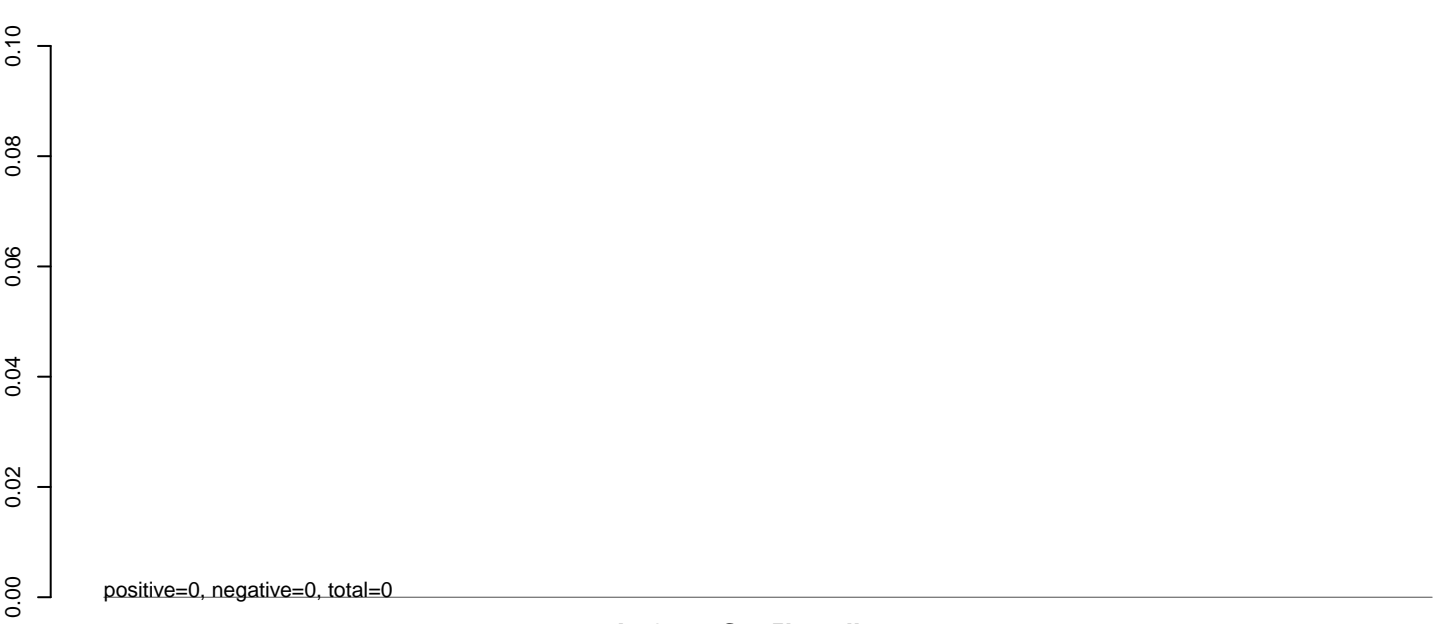
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AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

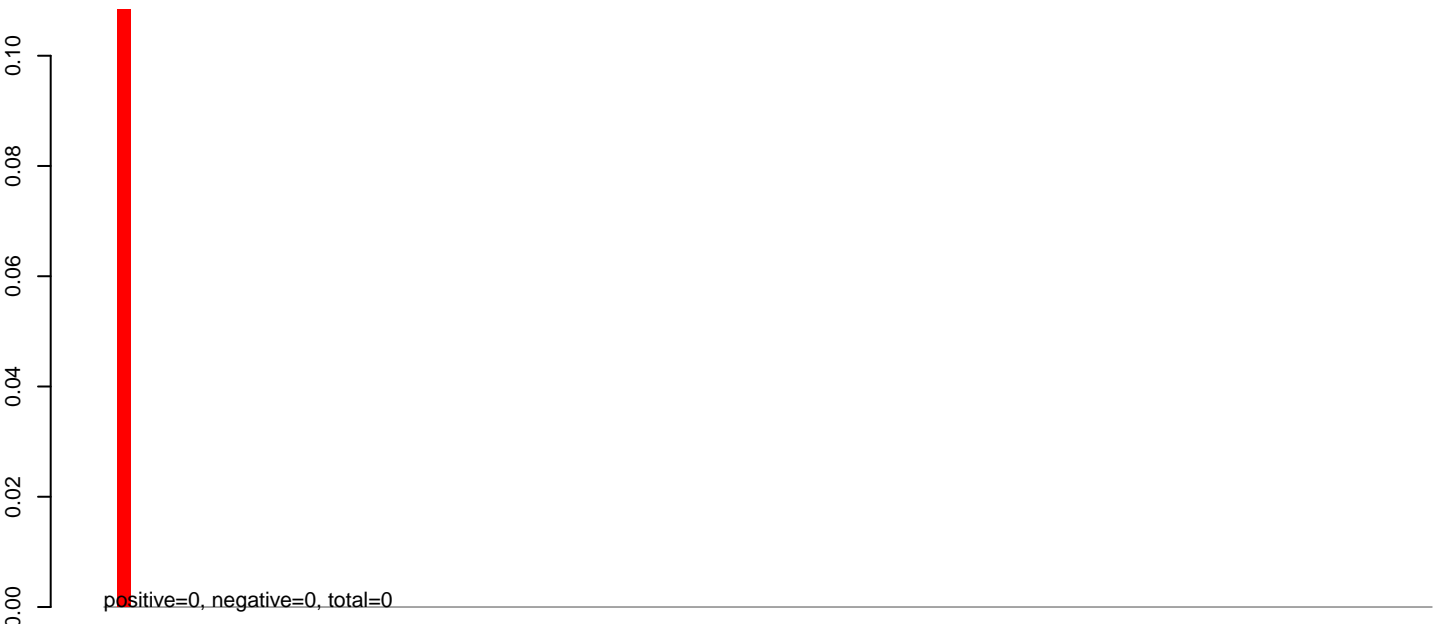


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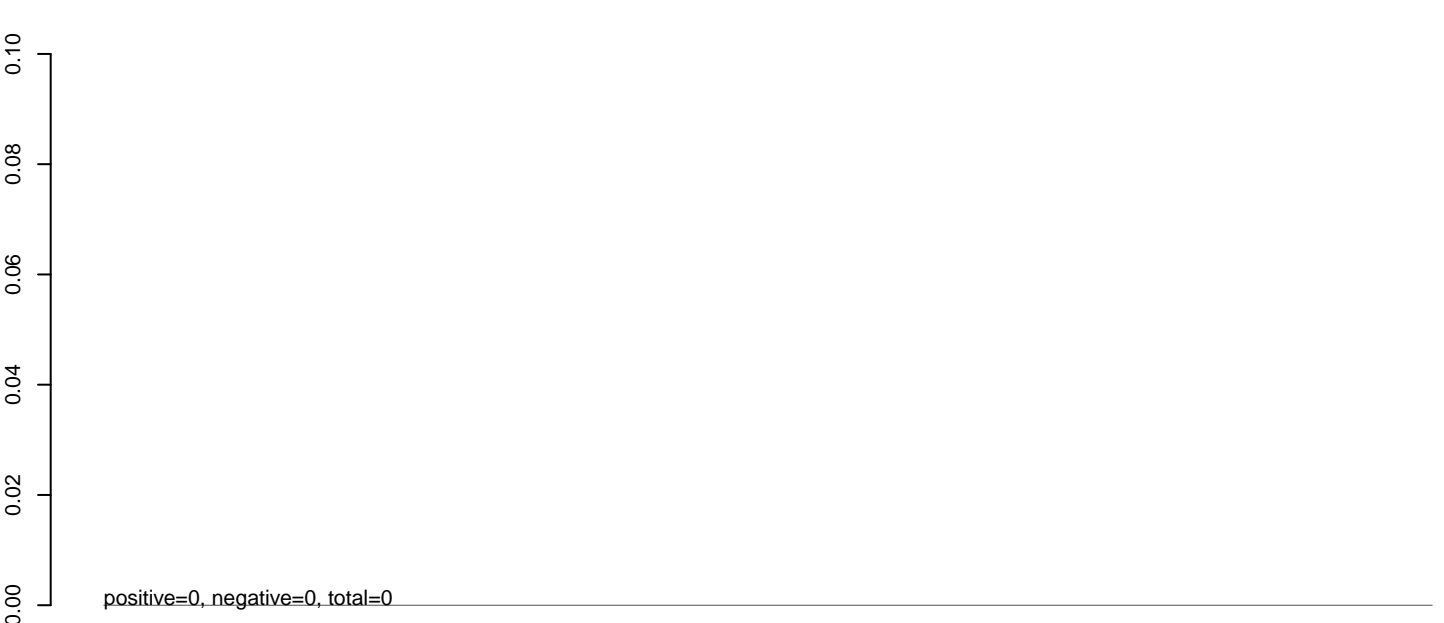


0 1000 2000 3000 4000 5000

AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

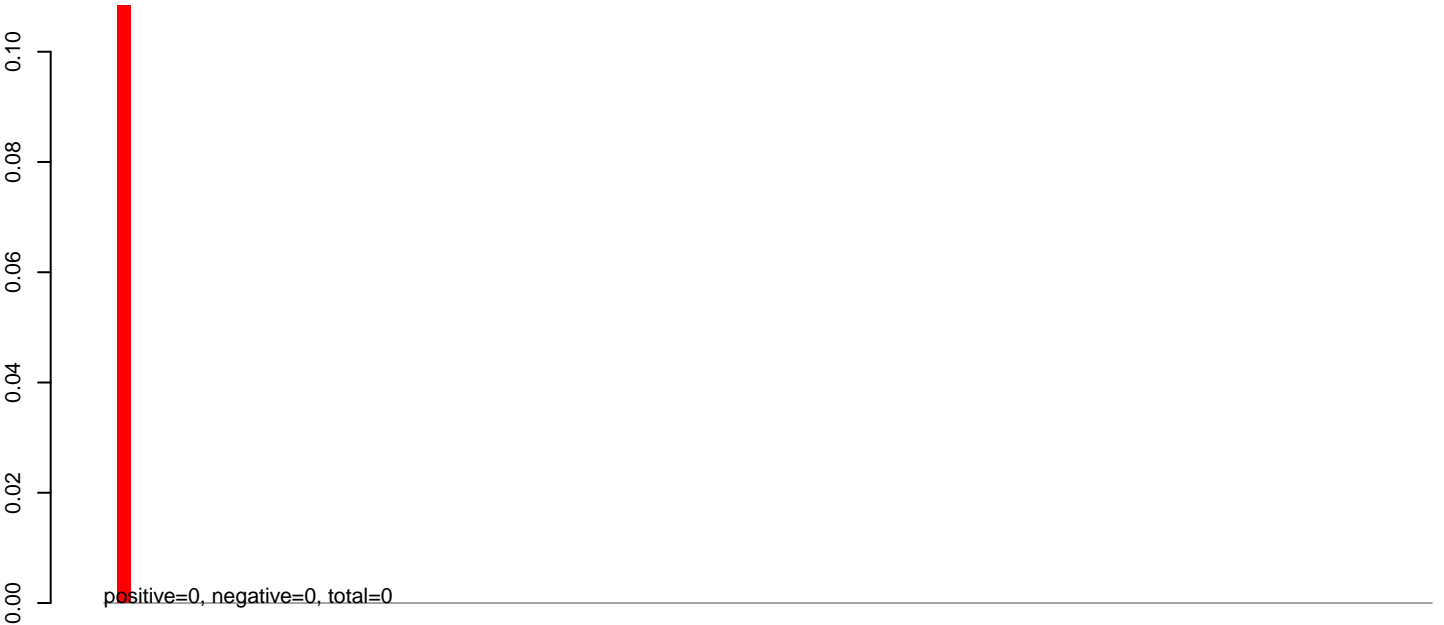


AnGam_Sua5b_cells.rep

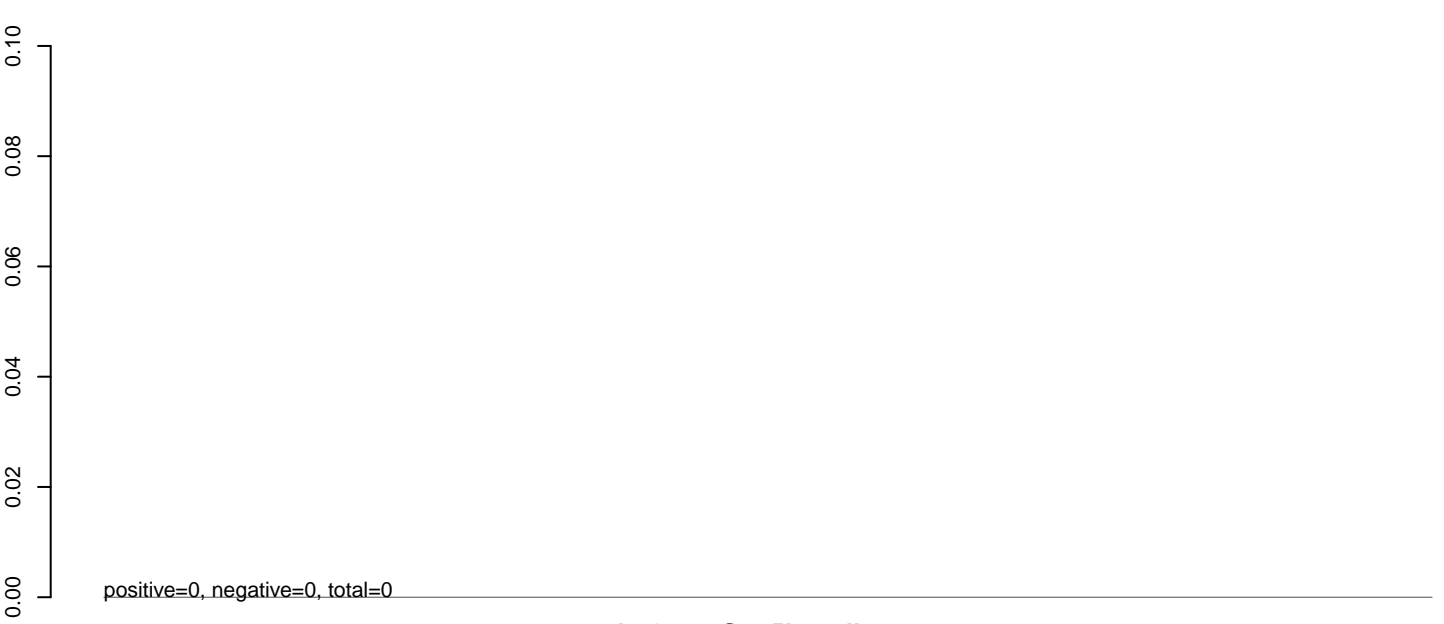


0 1000 2000 3000 4000 5000

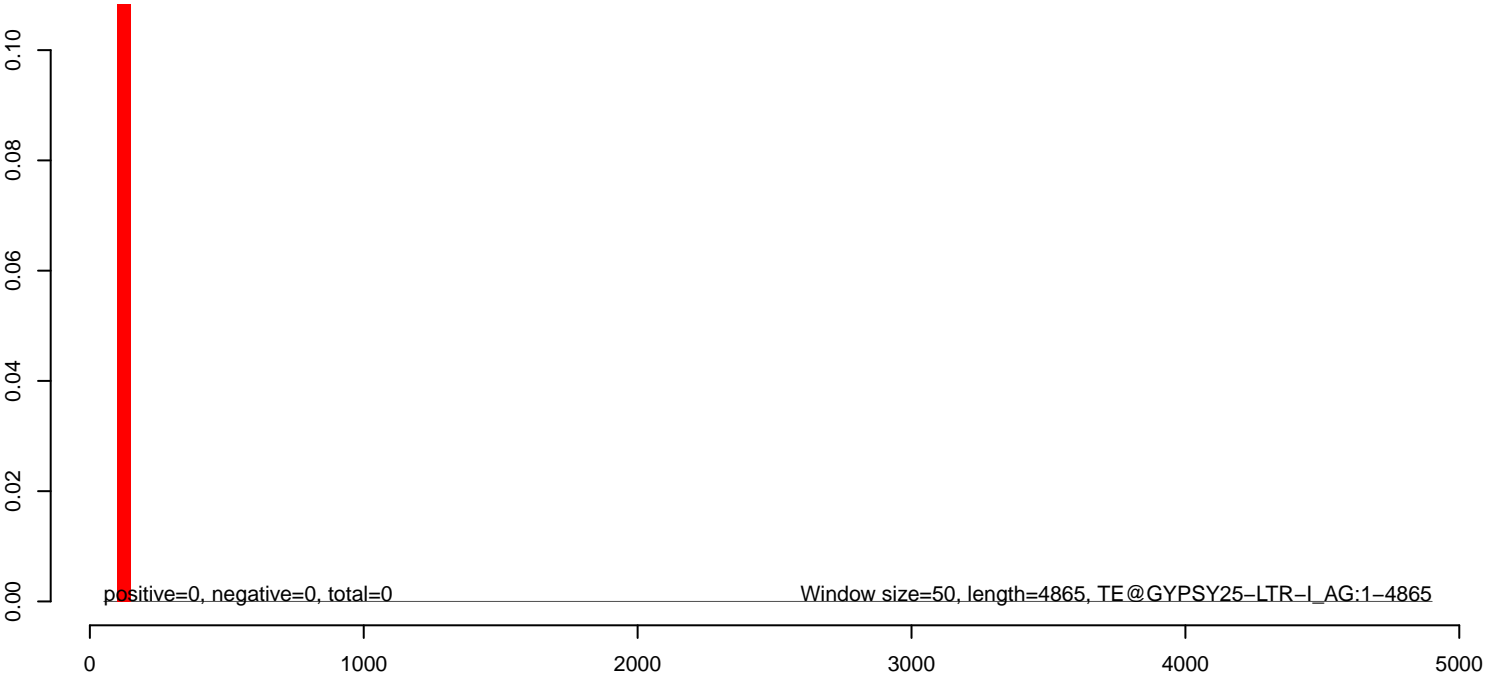
AnGam_Sua5b_cells.18_23.rep



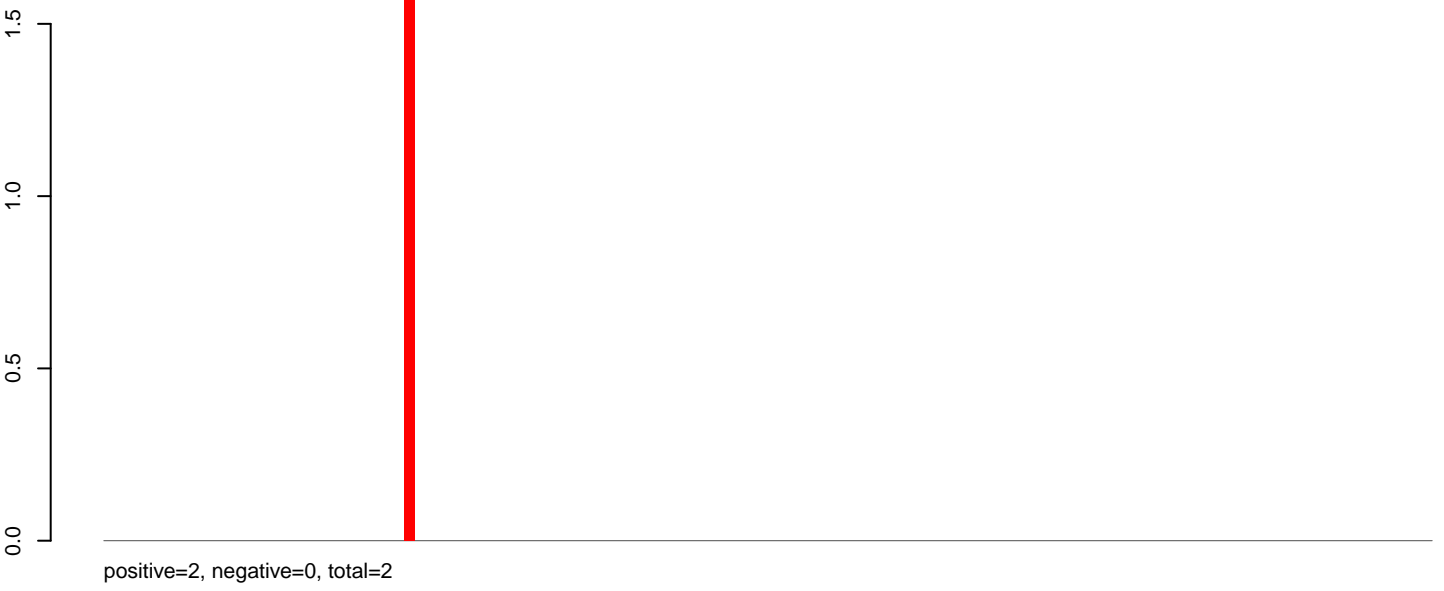
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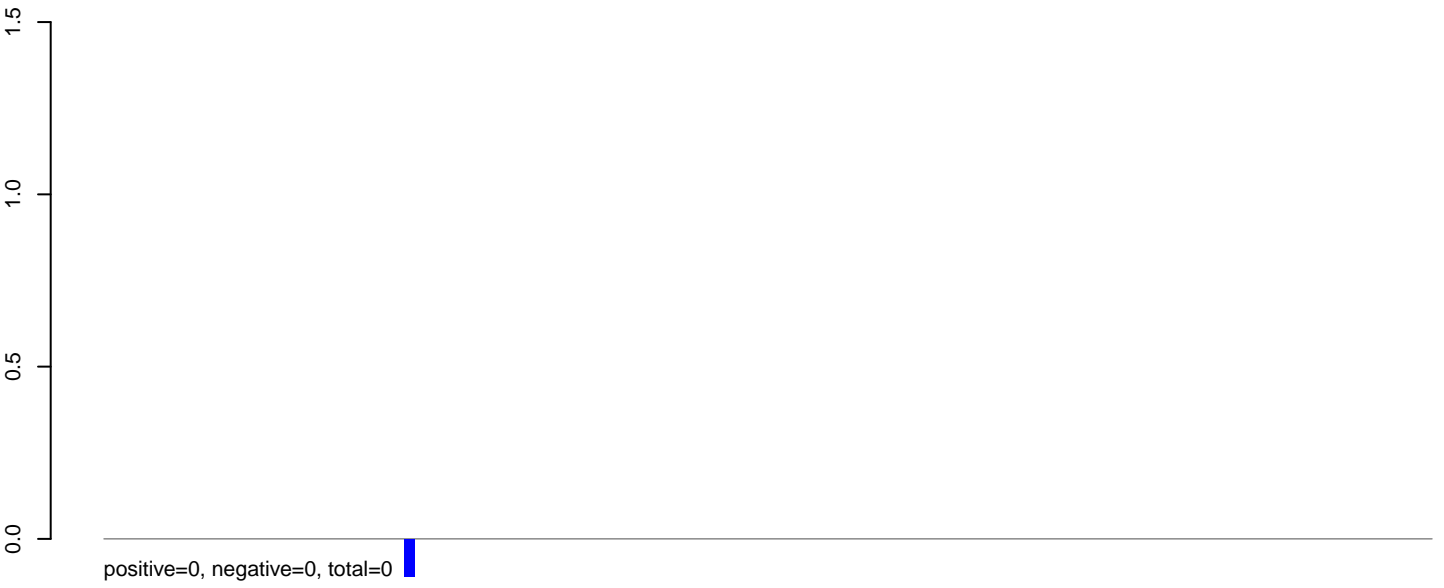
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AnGam_Sua5b_cells.18_23.rep



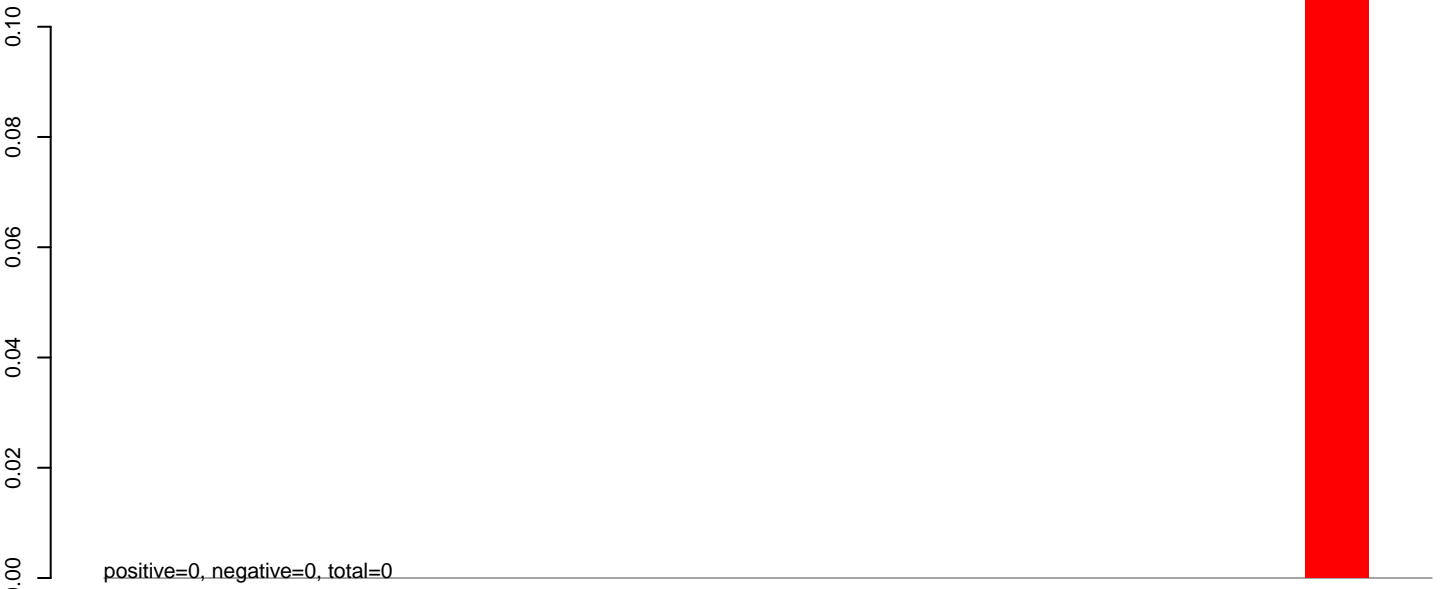
AnGam_Sua5b_cells.24_35.rep



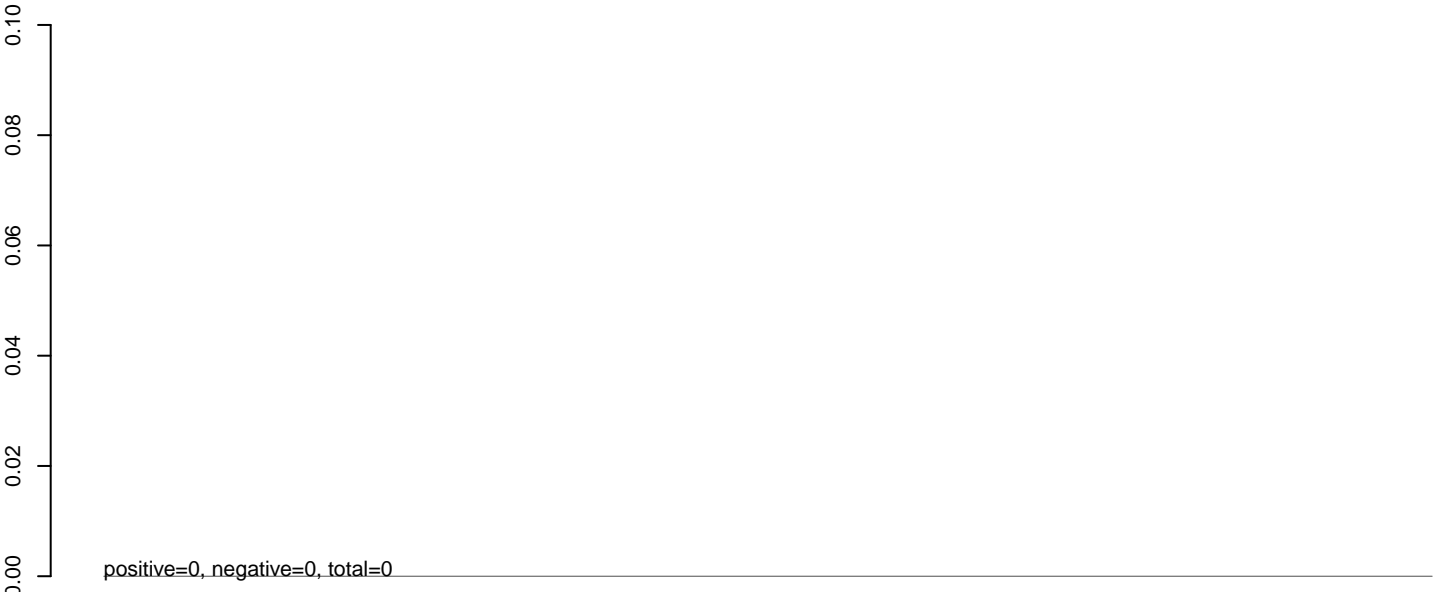
AnGam_Sua5b_cells.rep



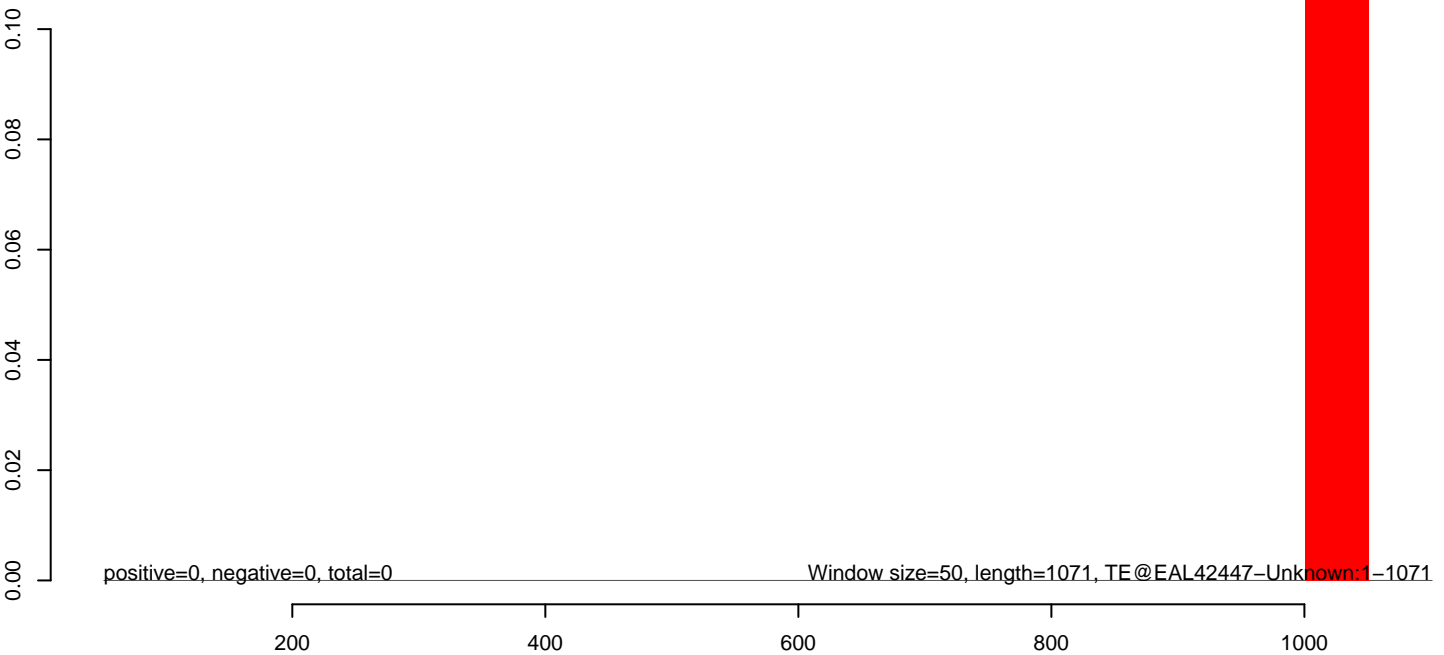
AnGam_Sua5b_cells.18_23.rep



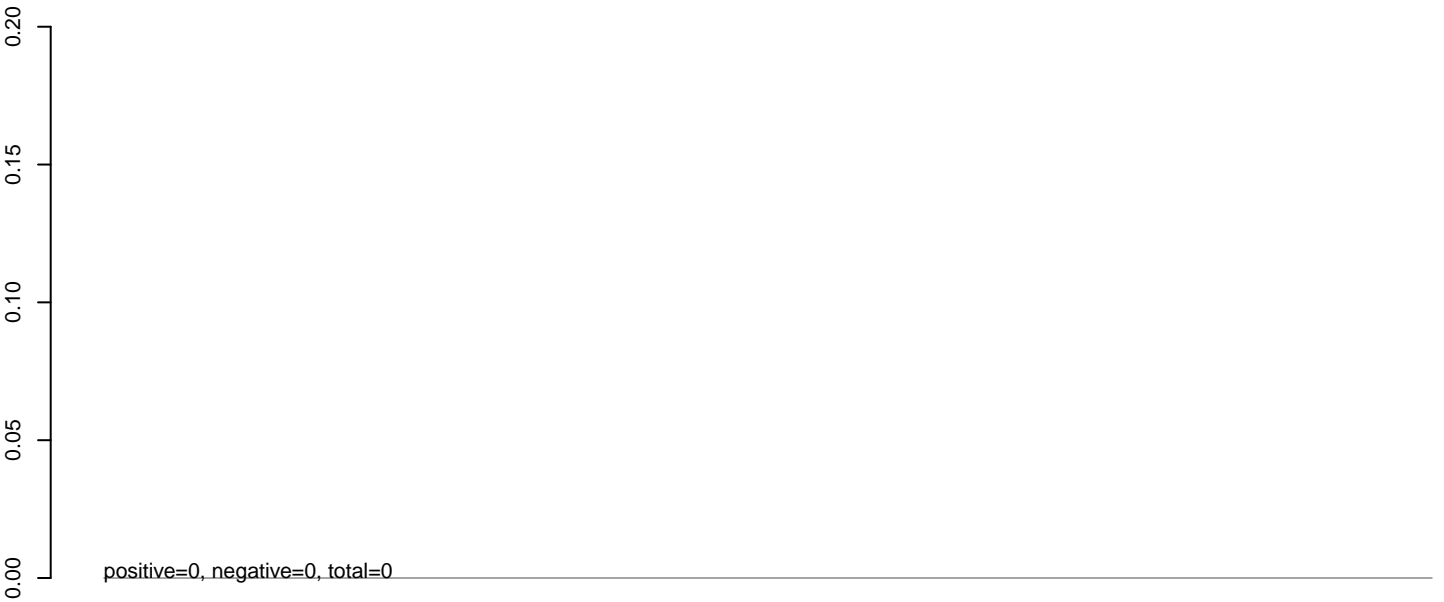
AnGam_Sua5b_cells.24_35.rep



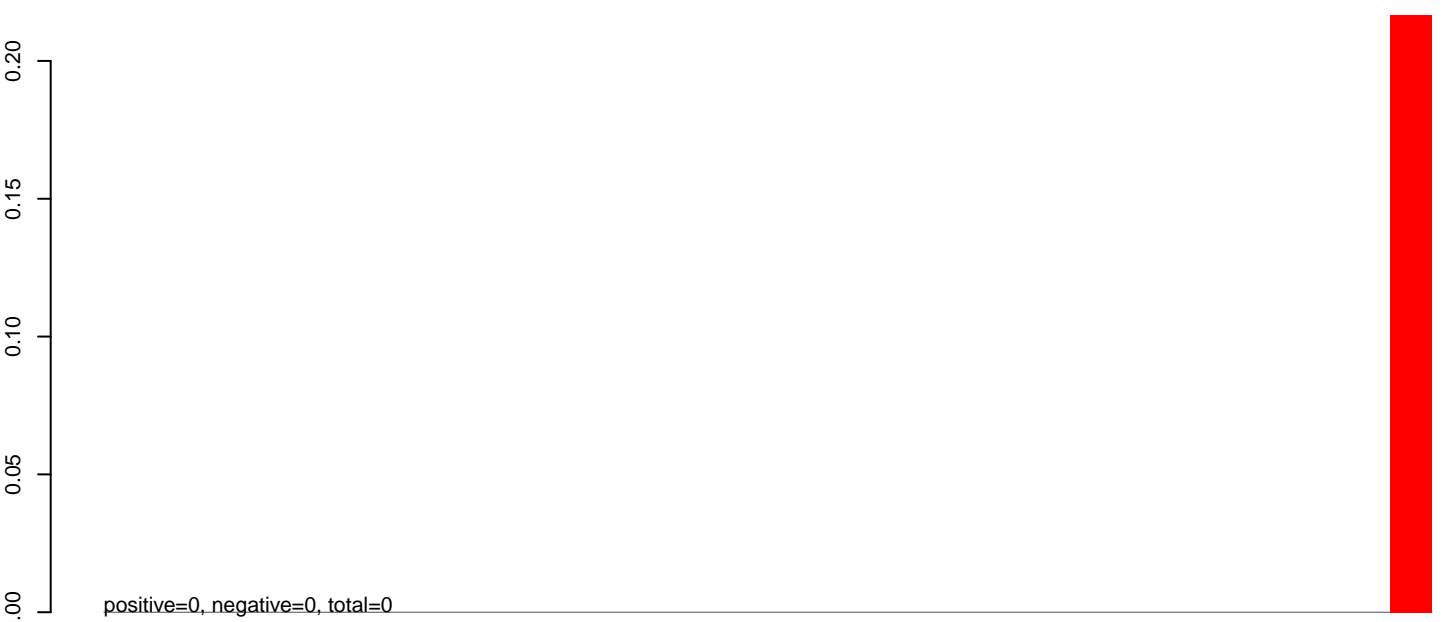
AnGam_Sua5b_cells.rep



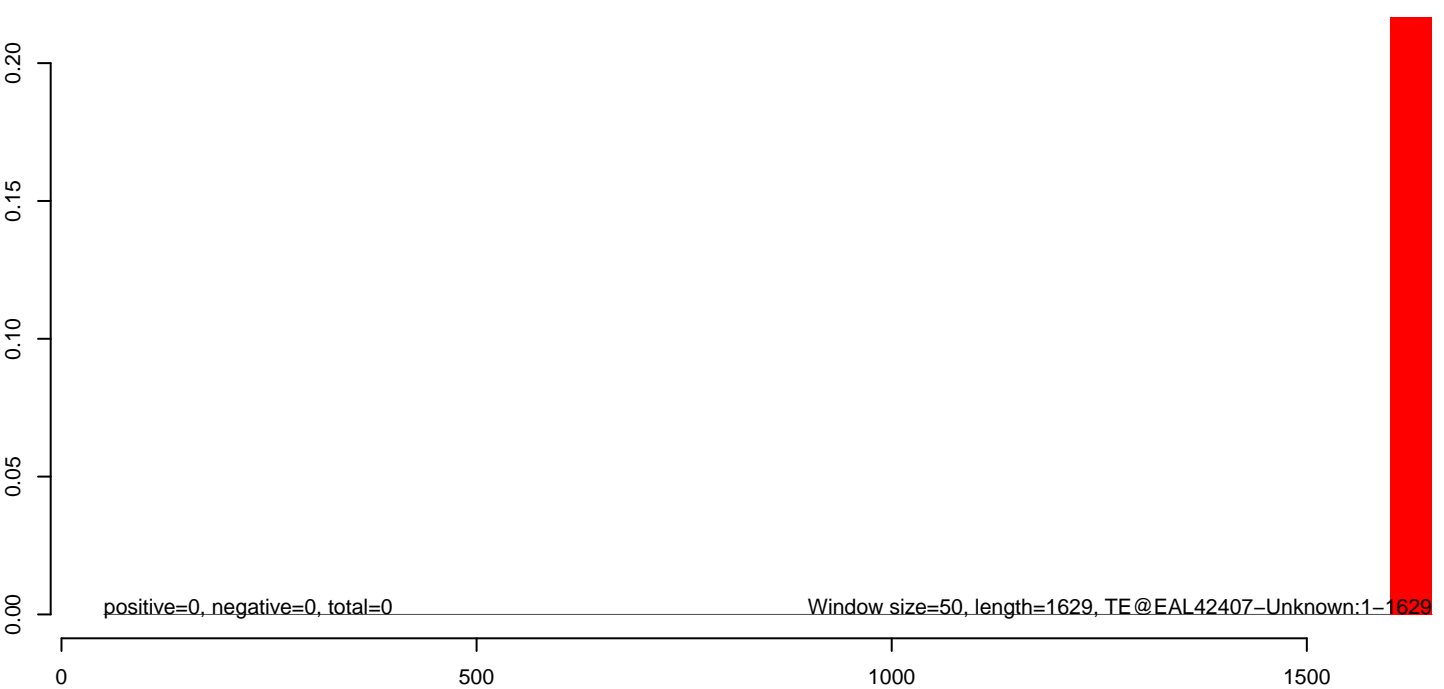
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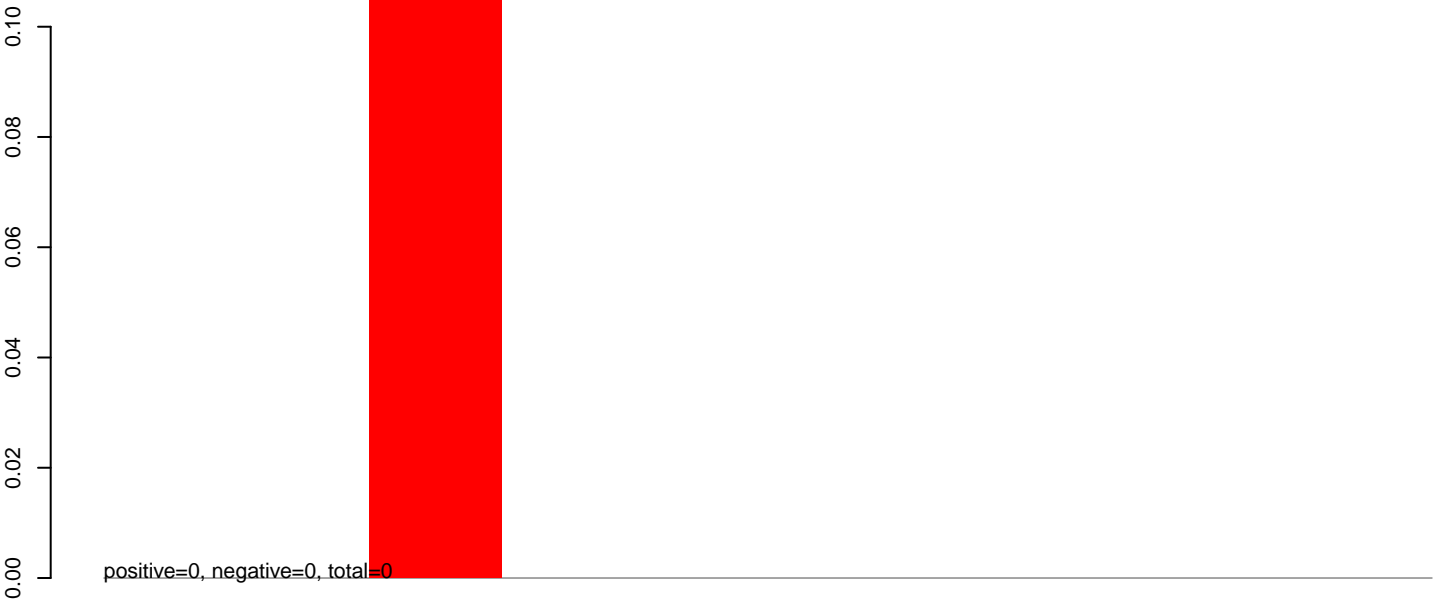
AnGam_Sua5b_cells.24_35.rep



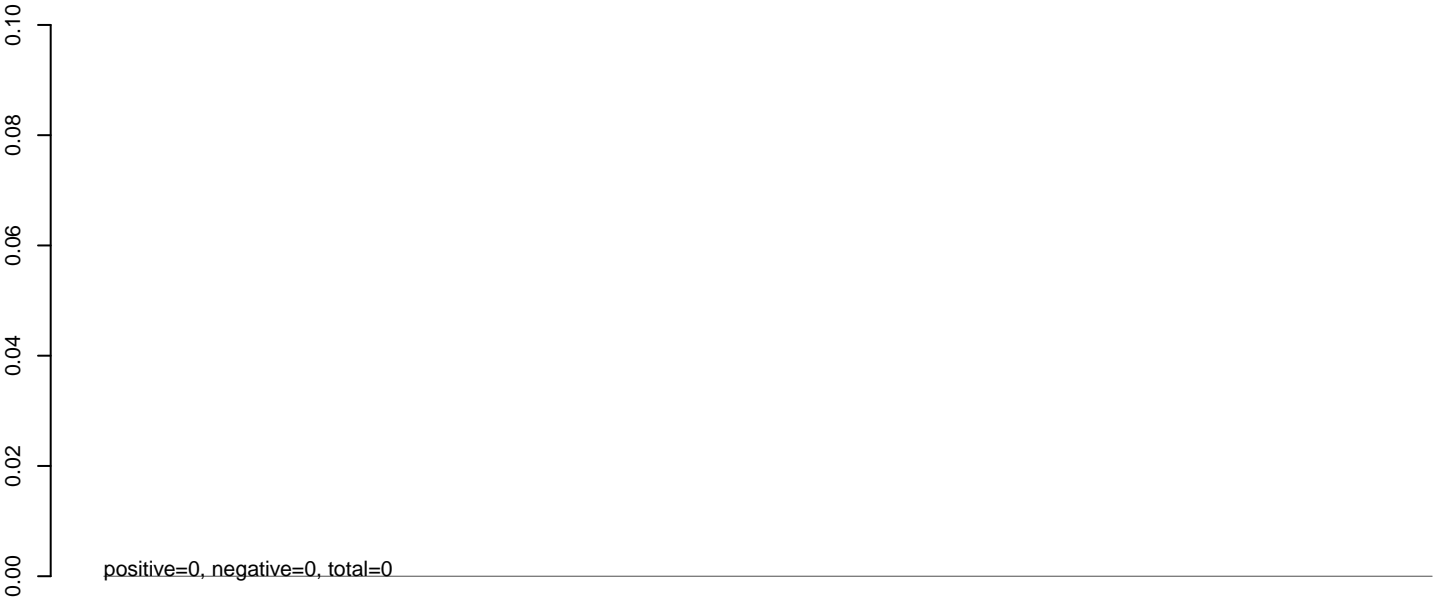
AnGam_Sua5b_cells.rep



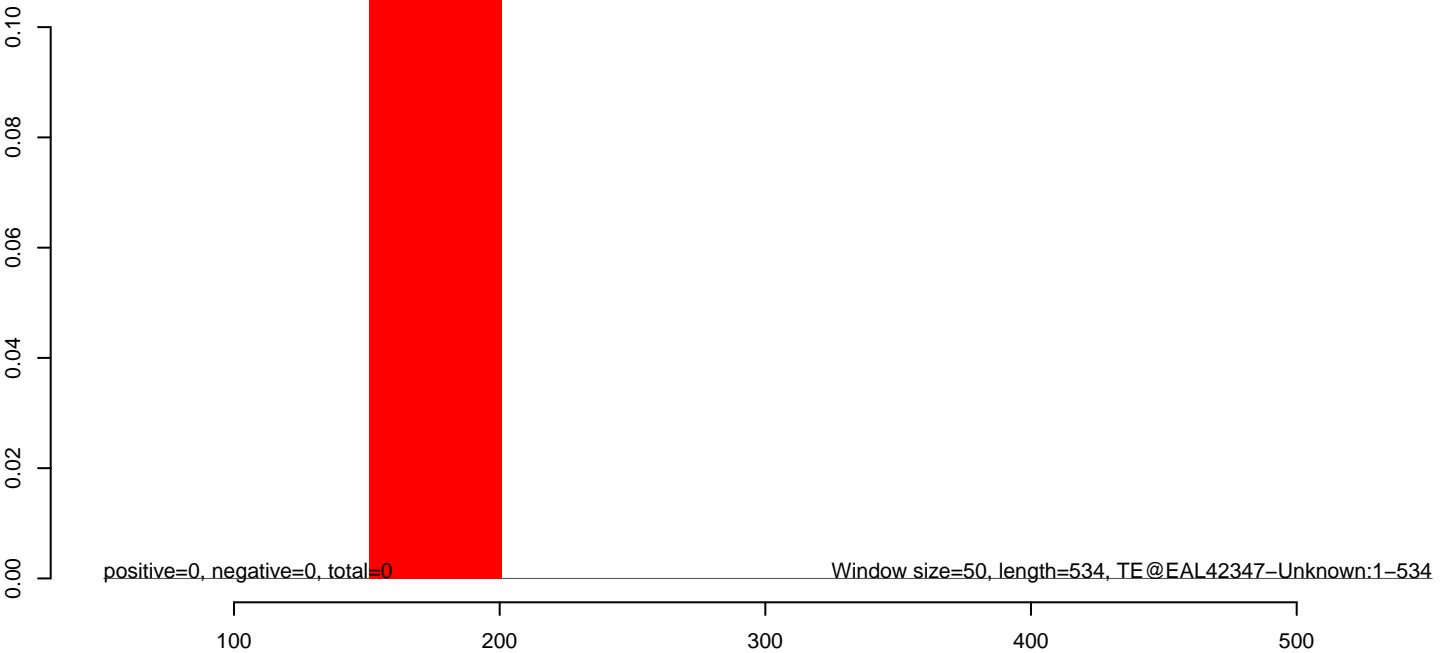
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.rep

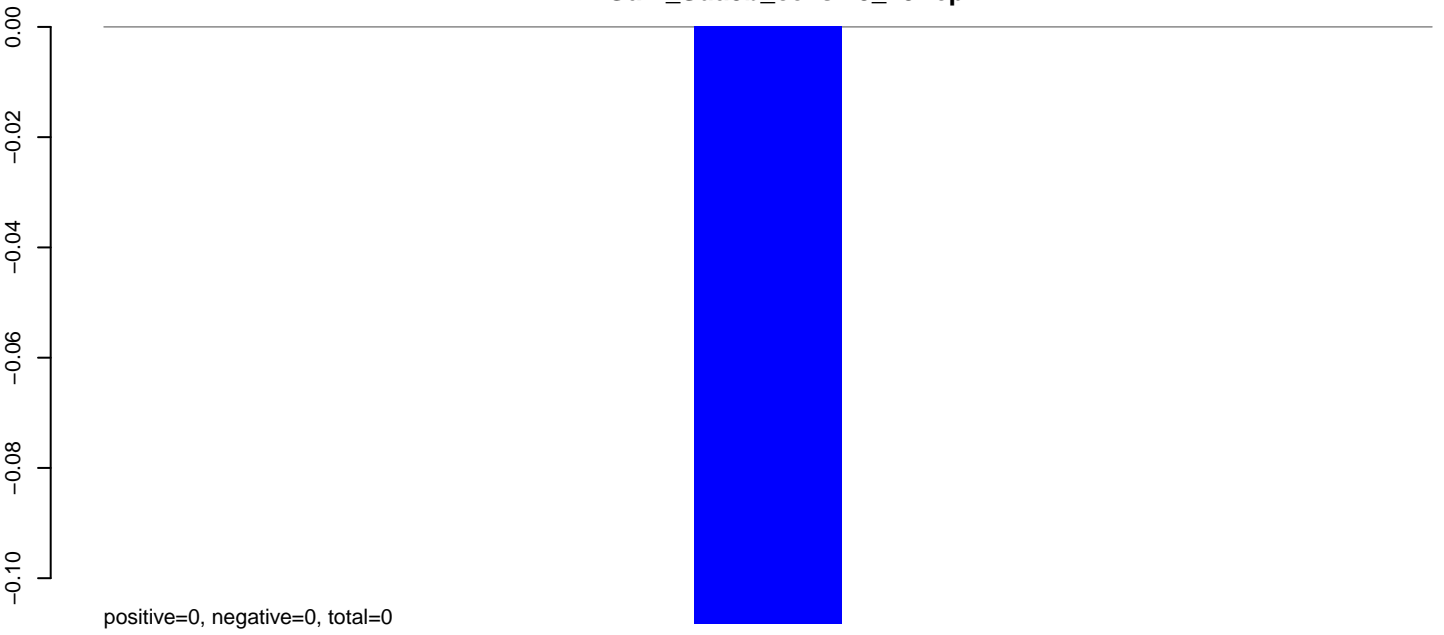
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positive=0, negative=0, total=0

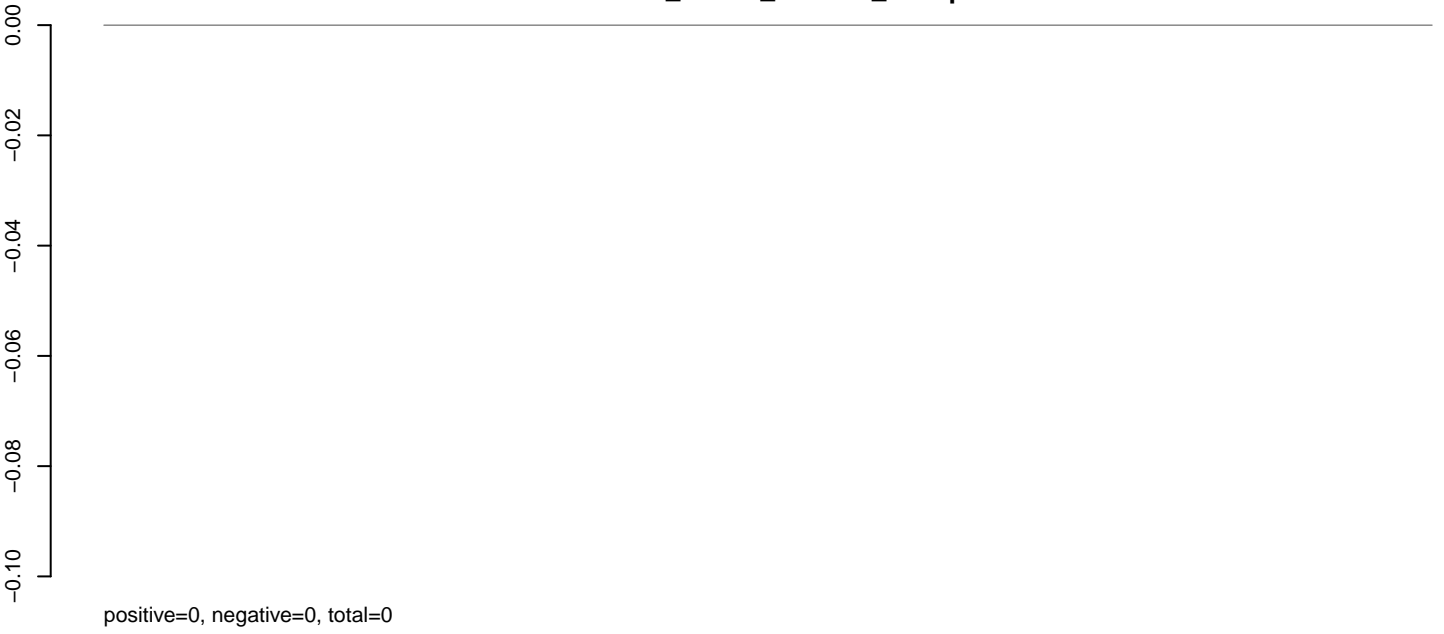
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200 400 600 800 1000 1200

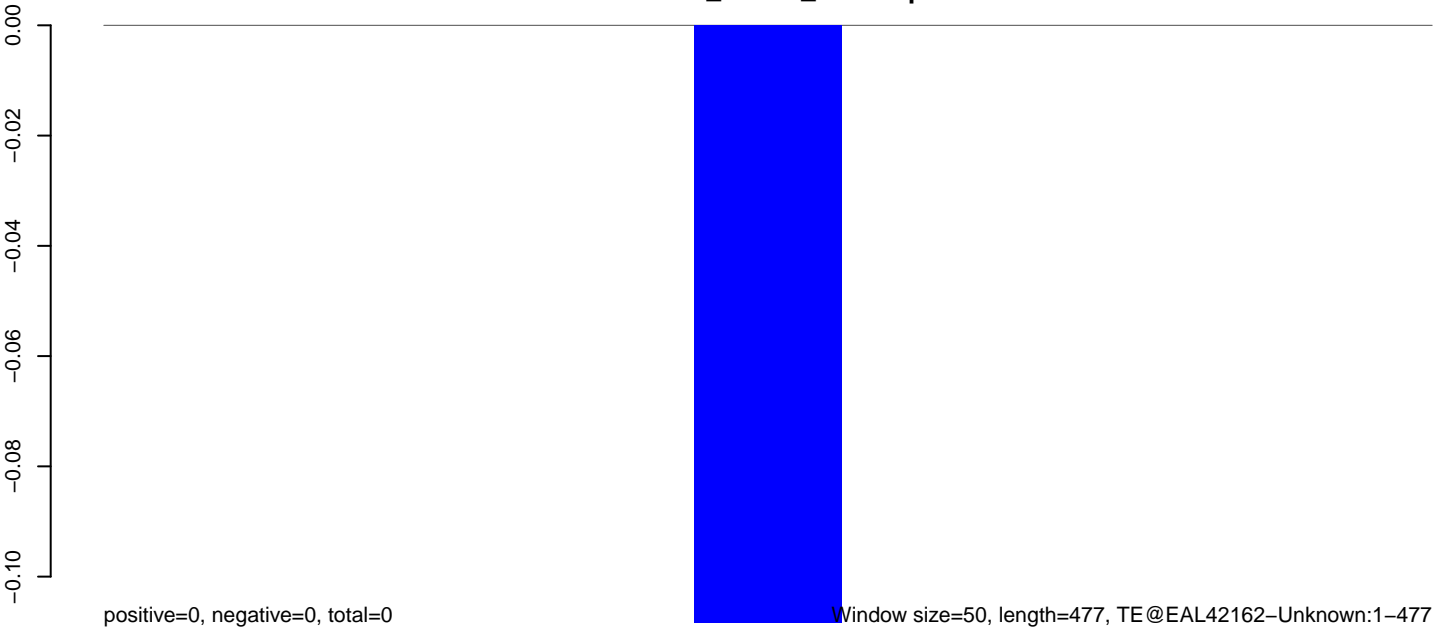
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

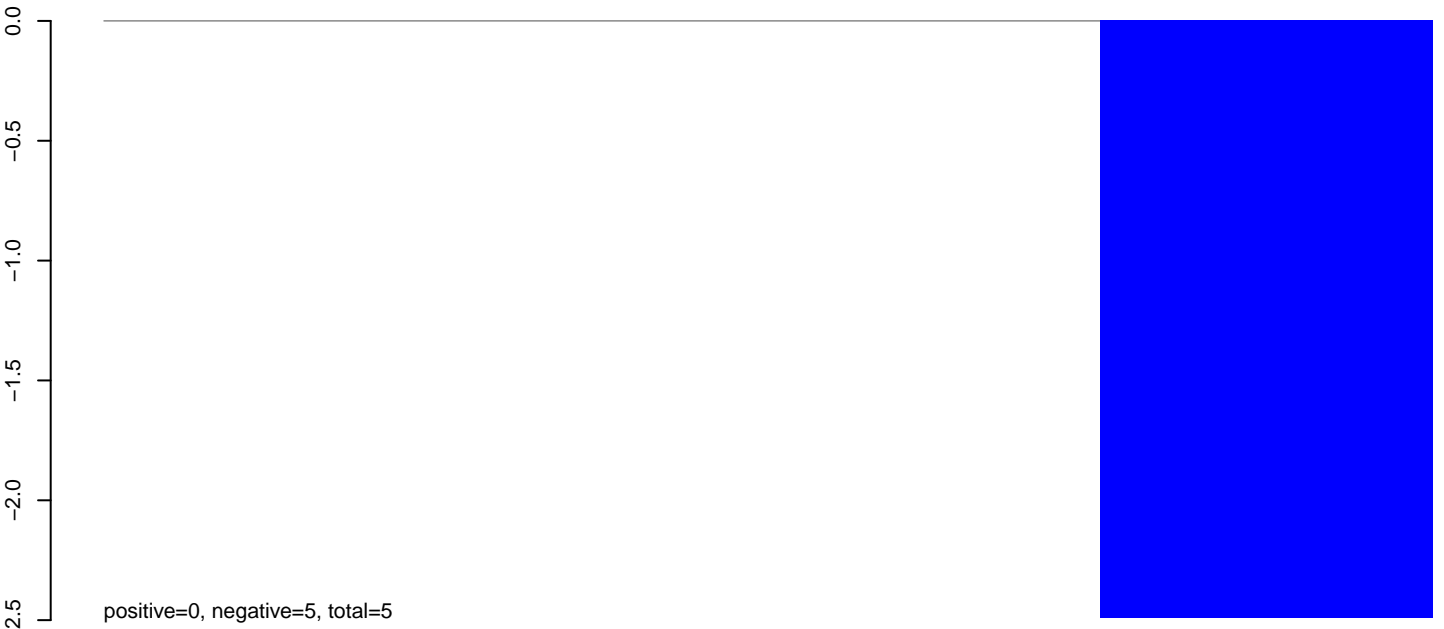


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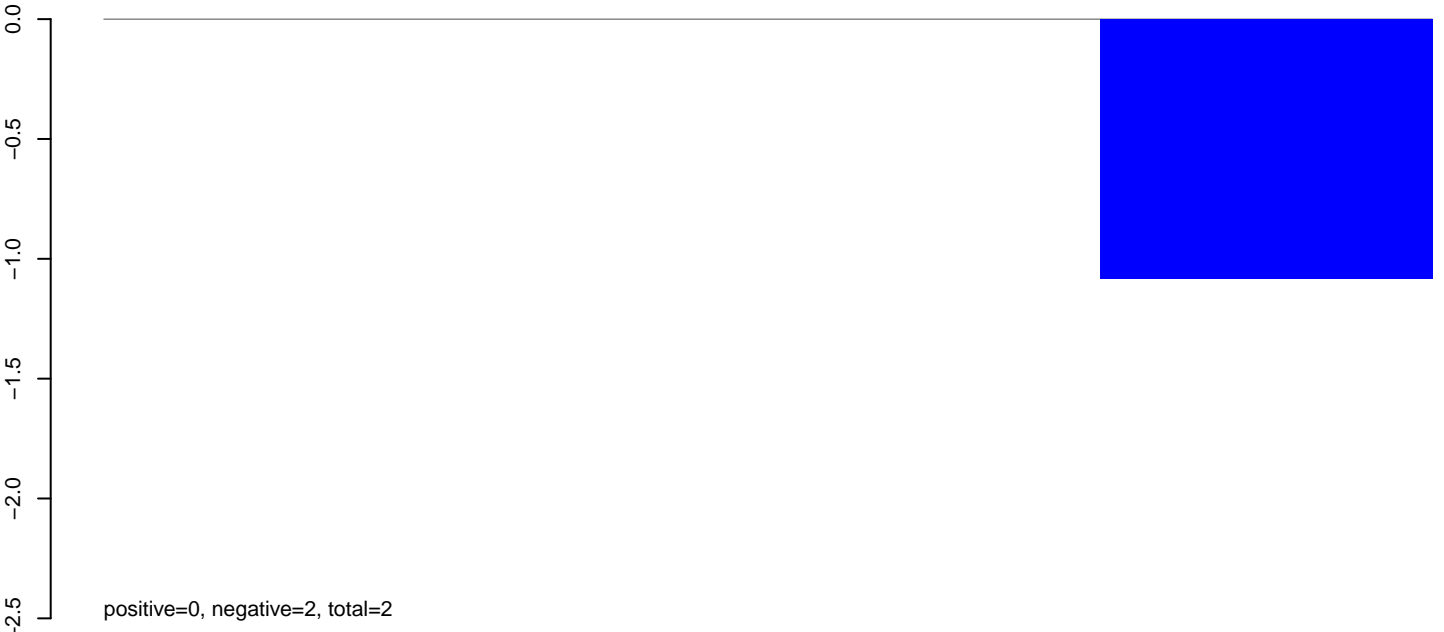


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AnGam_Sua5b_cells.18_23.rep



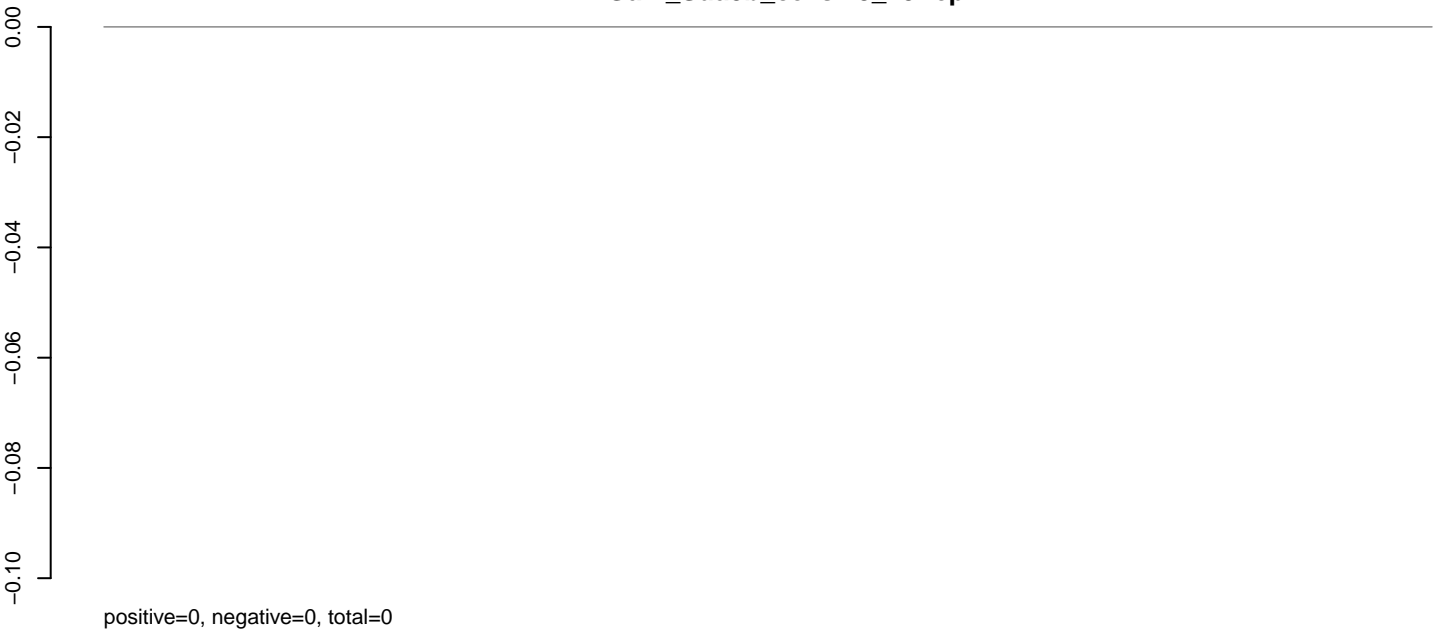
AnGam_Sua5b_cells.24_35.rep



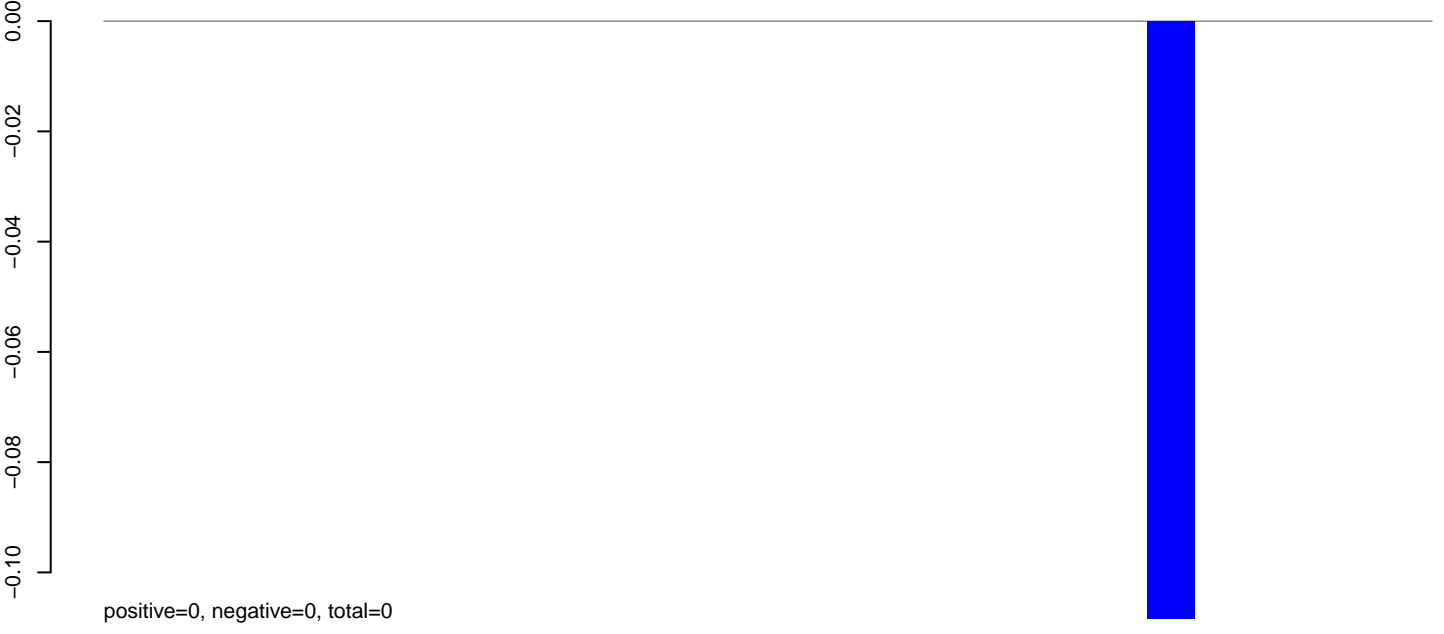
AnGam_Sua5b_cells.rep



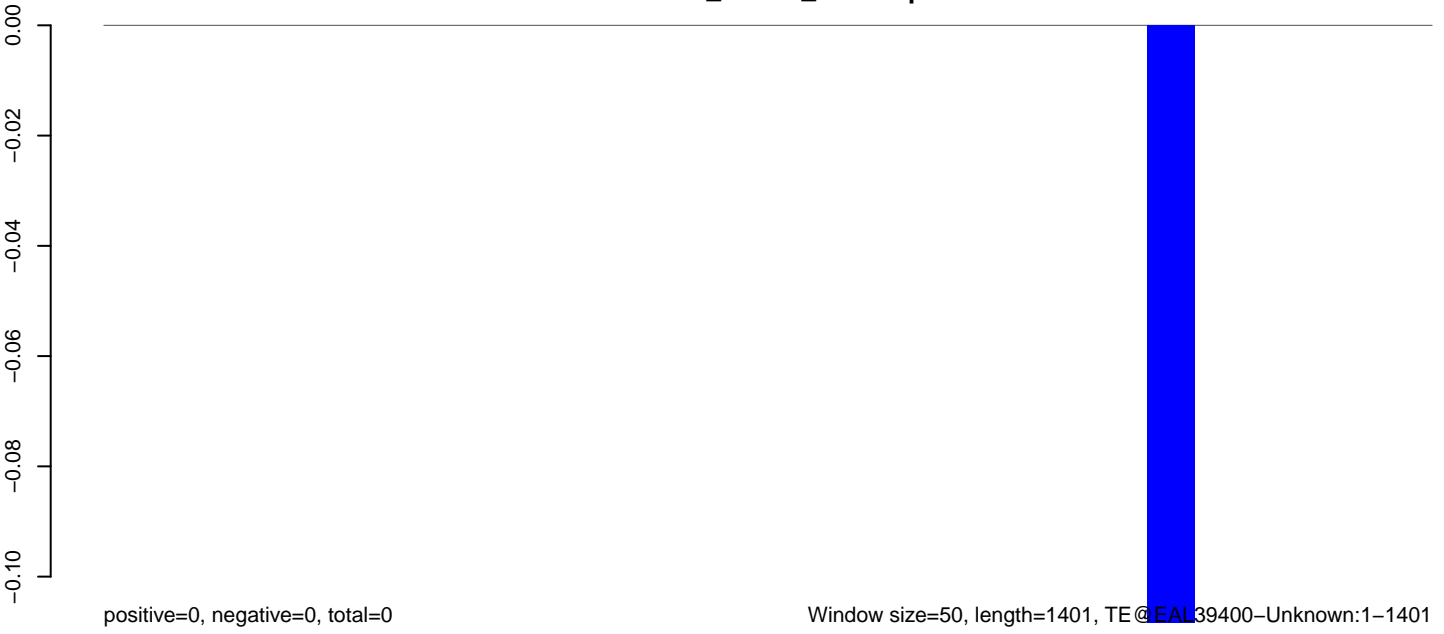
AnGam_Sua5b_cells.18_23.rep



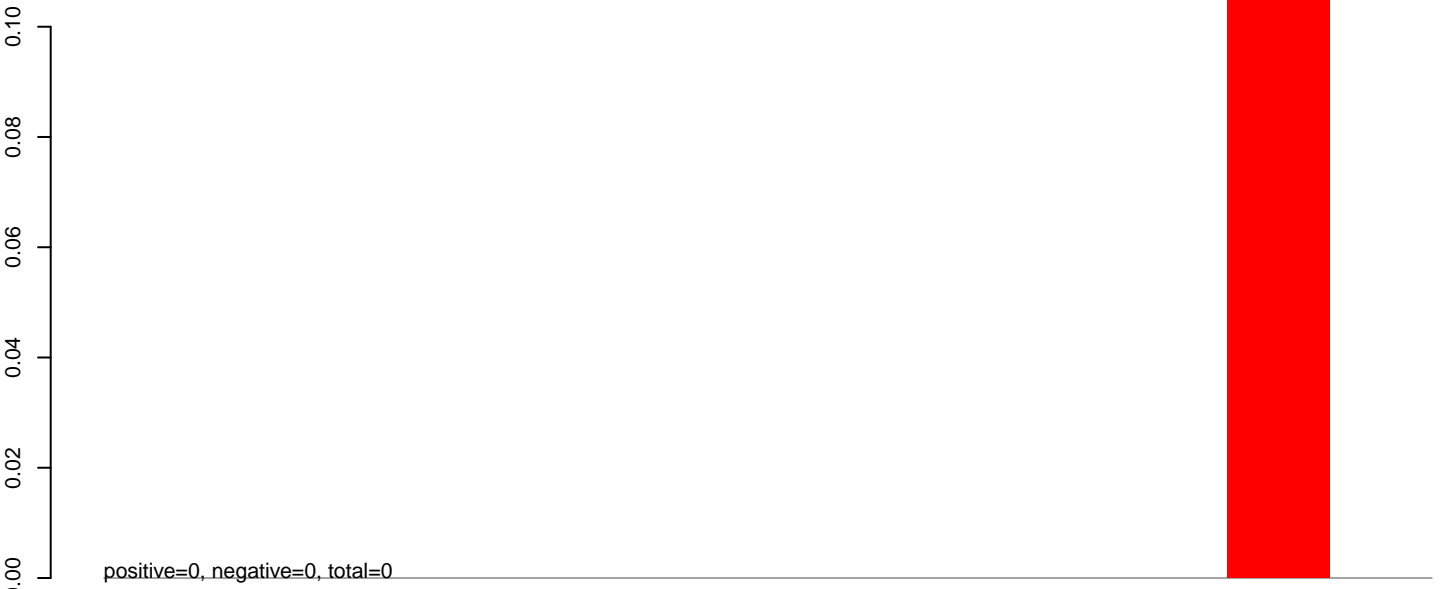
AnGam_Sua5b_cells.24_35.rep



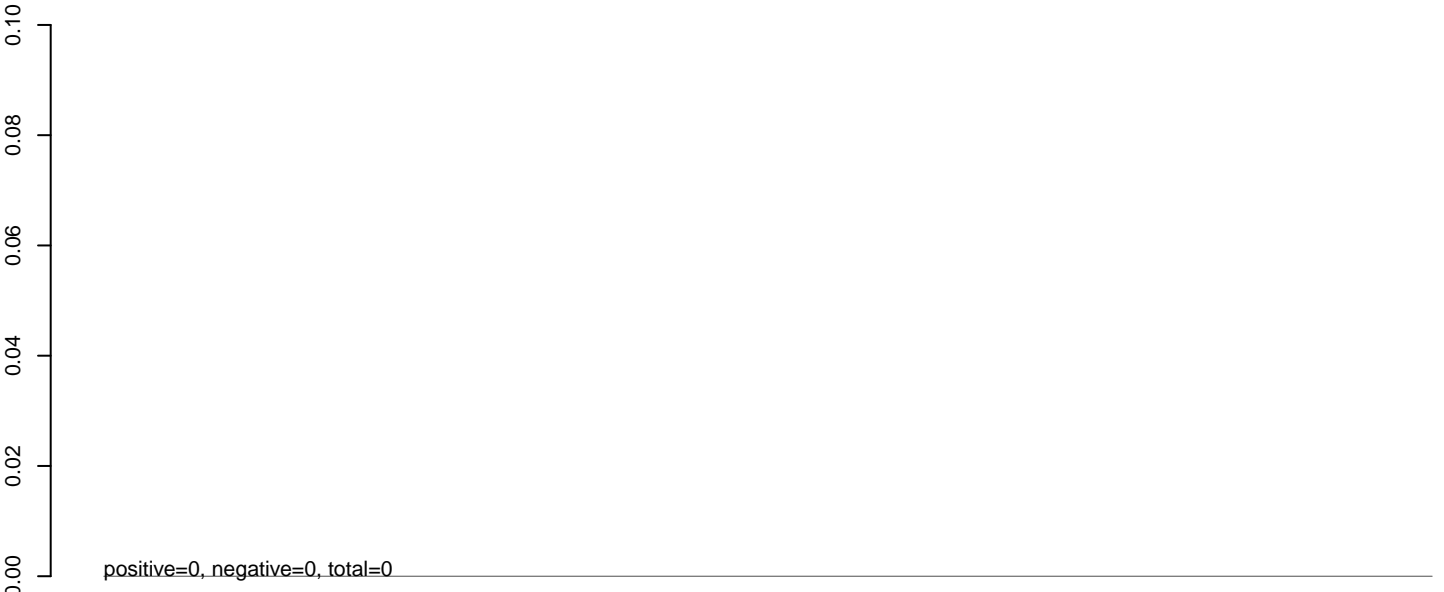
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

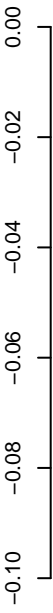


AnGam_Sua5b_cells.rep



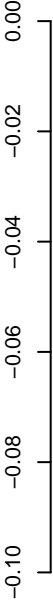
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AnGam_Sua5b_cells.18_23.rep



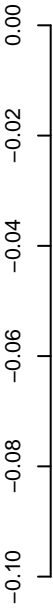
positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep



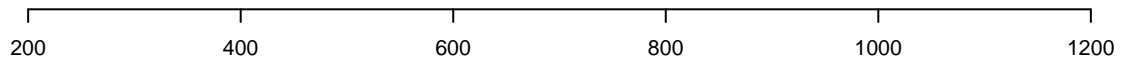
positive=0, negative=0, total=0

AnGam_Sua5b_cells.rep

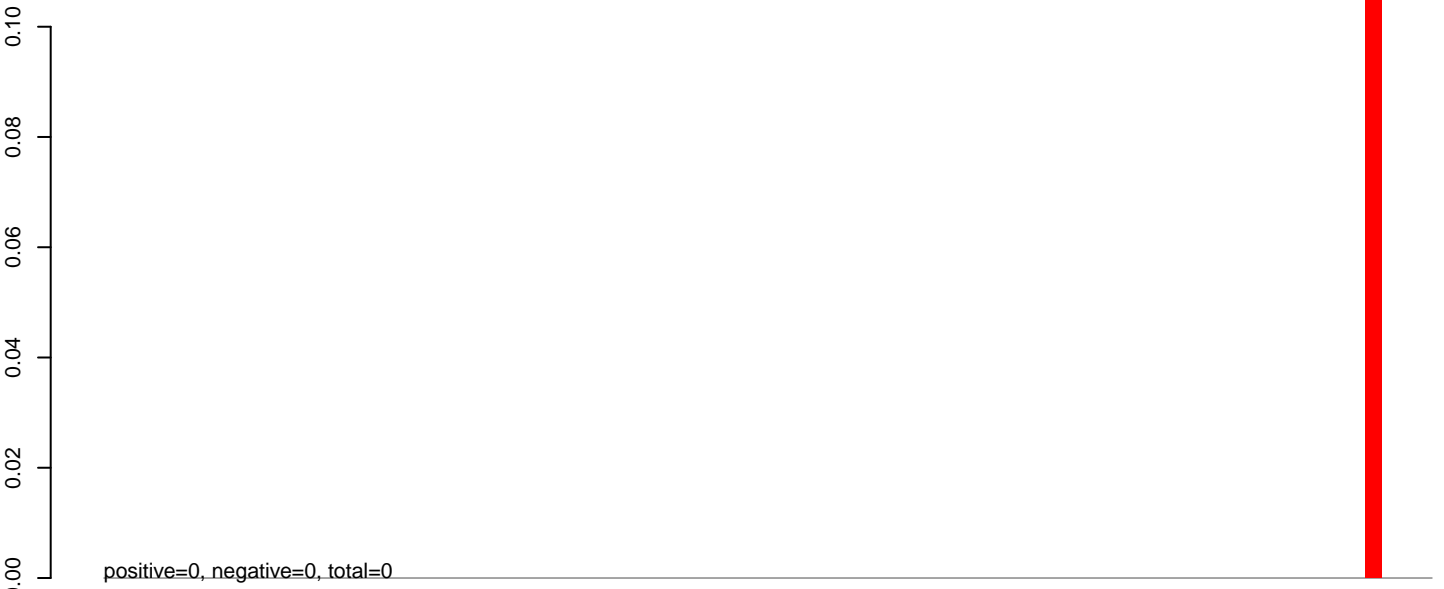


positive=0, negative=0, total=0

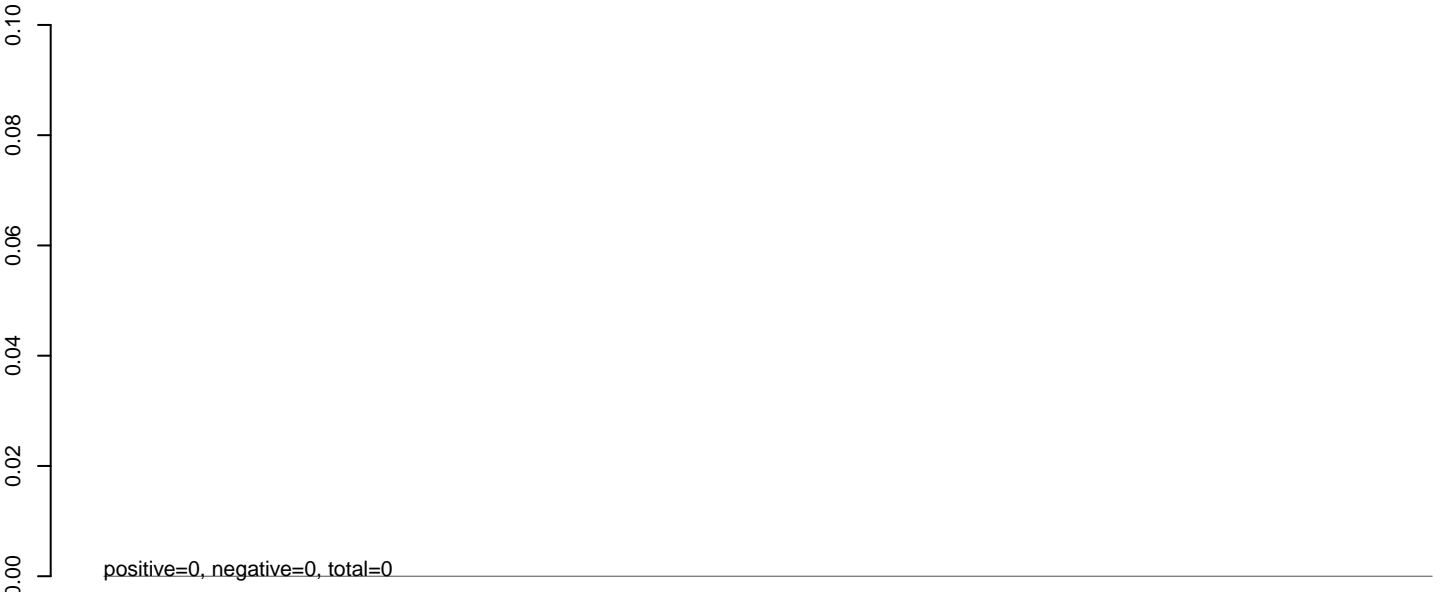
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AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

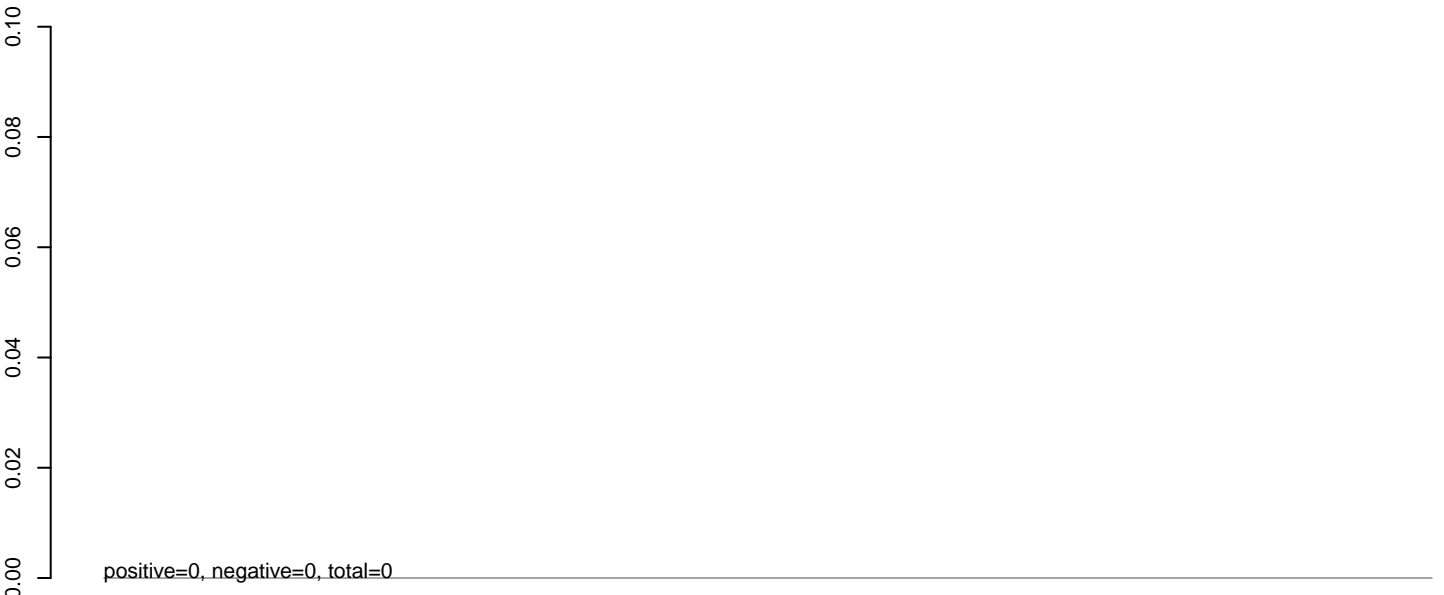


AnGam_Sua5b_cells.rep

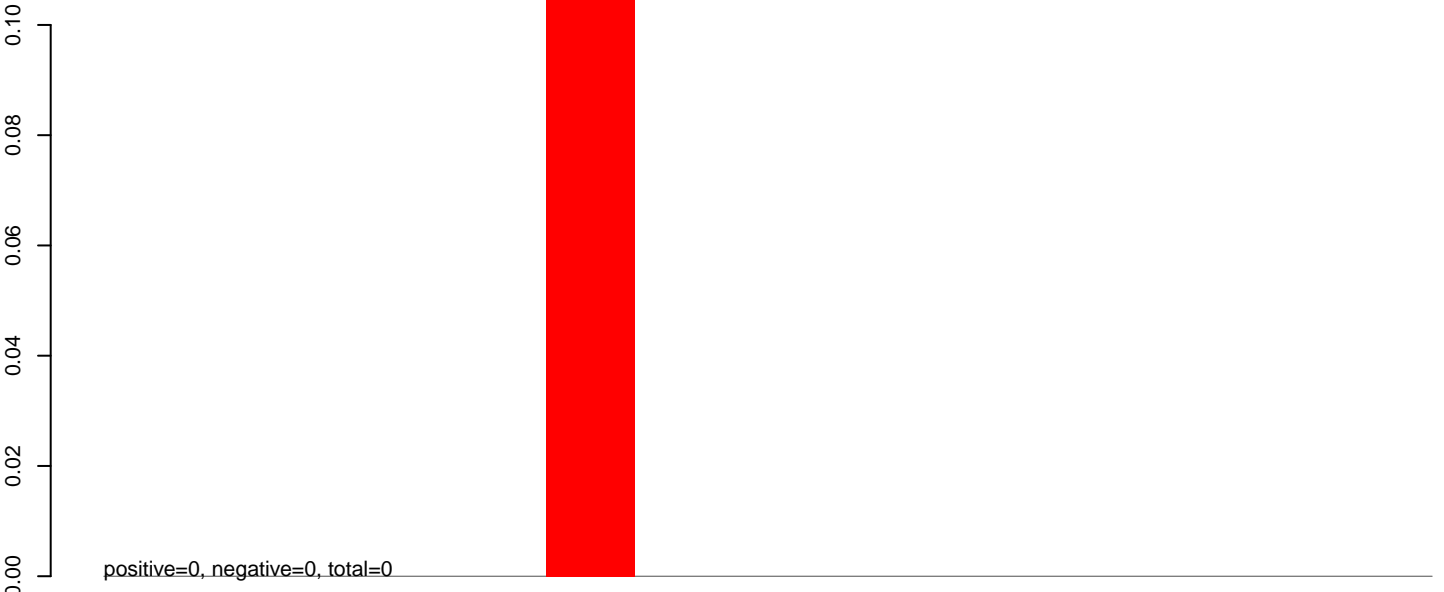


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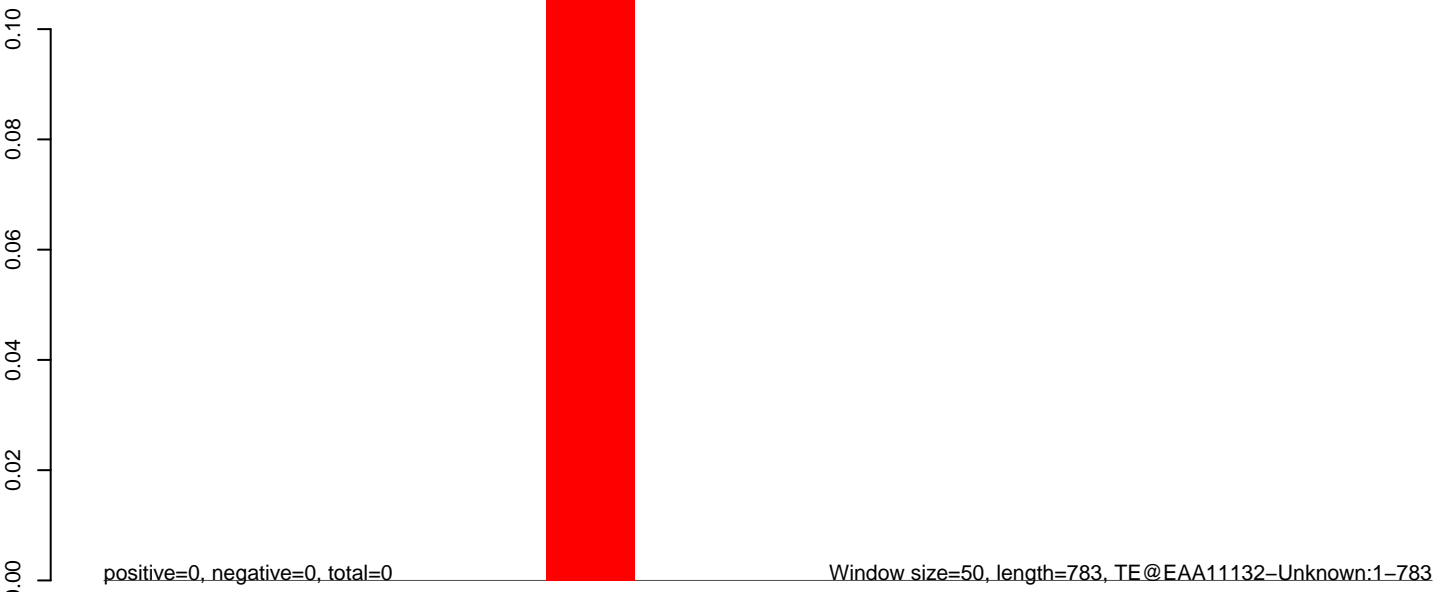
AnGam_Sua5b_cells.18_23.rep



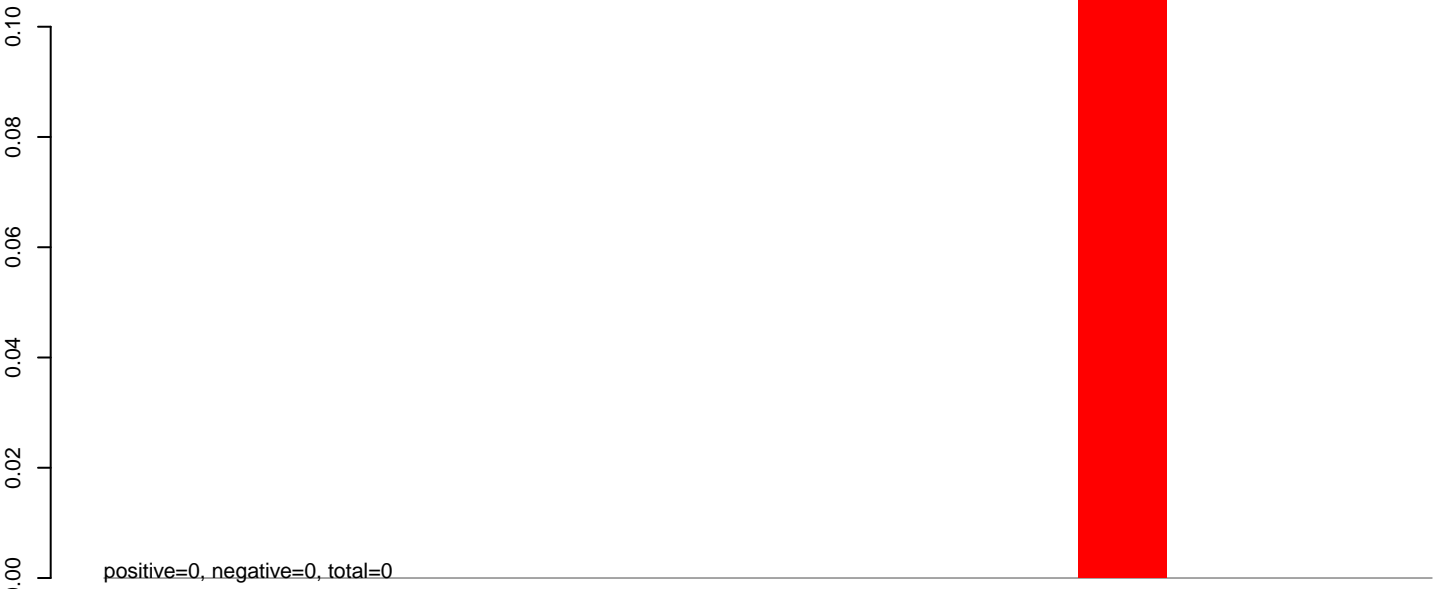
AnGam_Sua5b_cells.24_35.rep



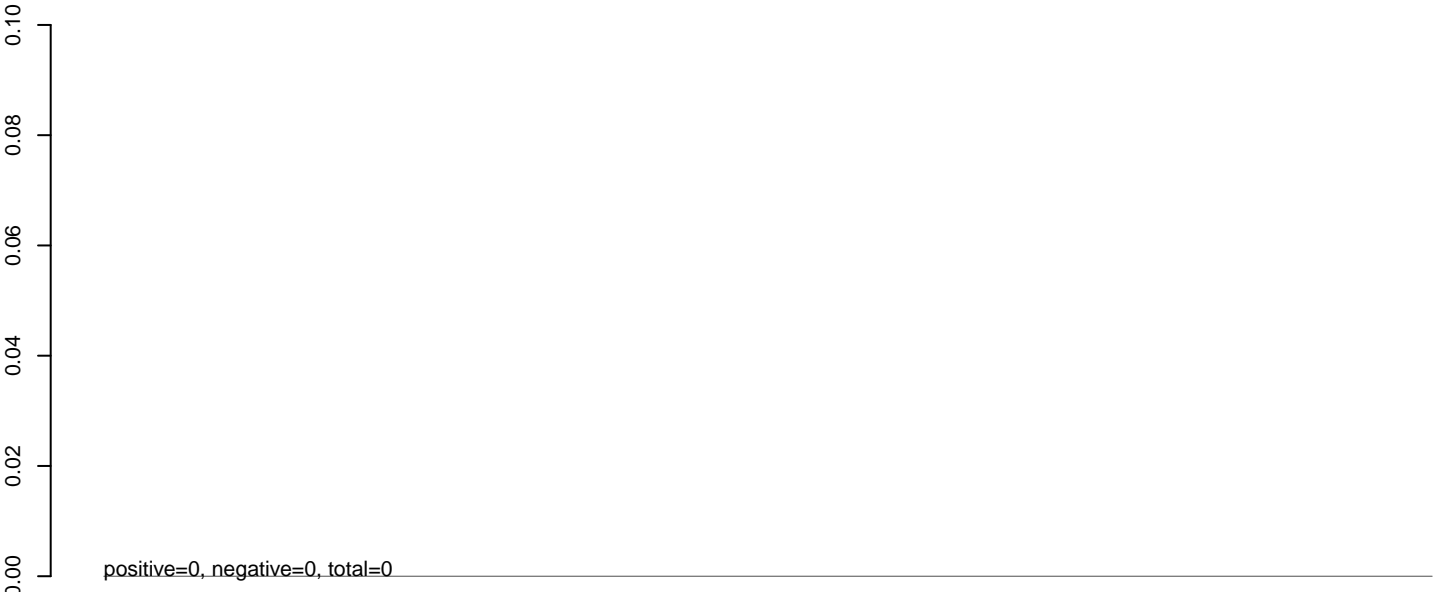
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.rep

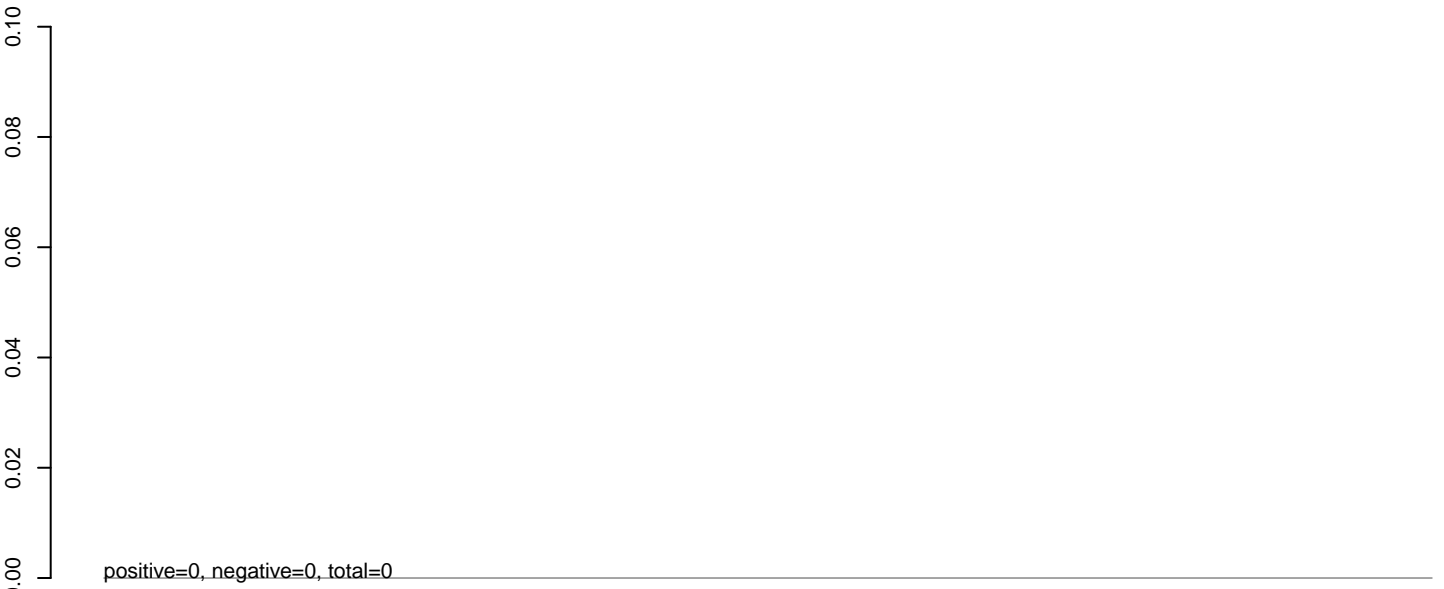
0.00
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-0.08
-0.10

positive=0, negative=0, total=0

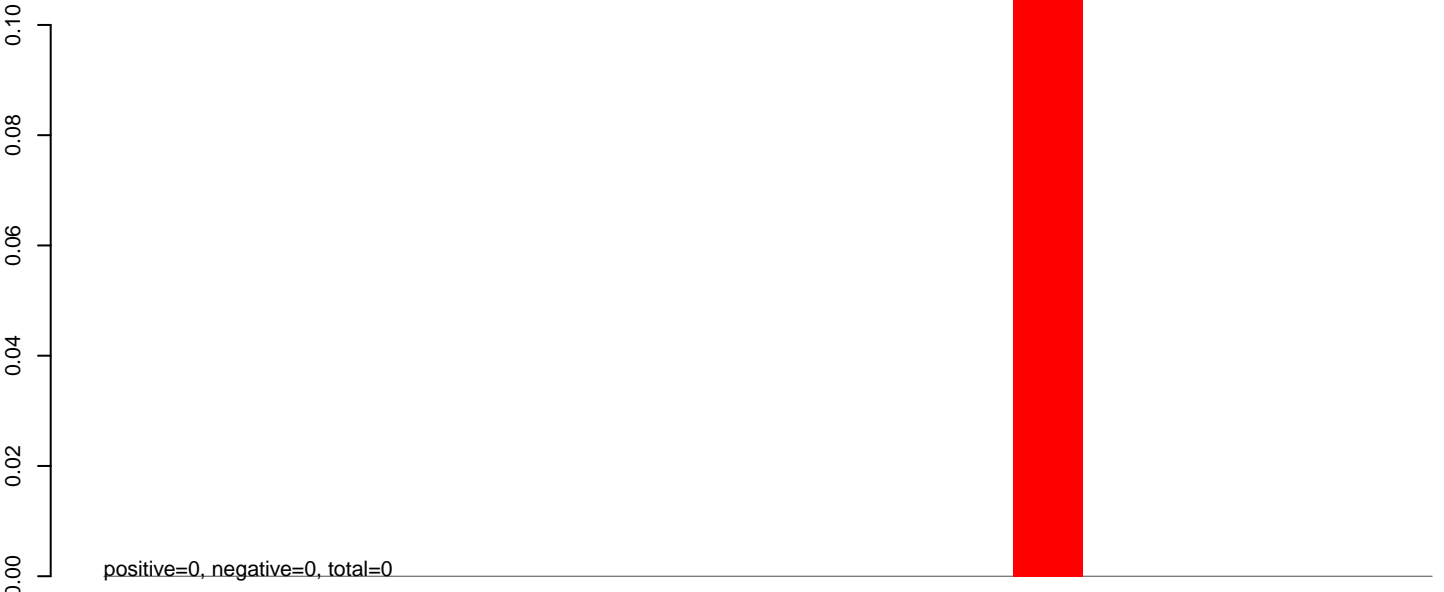
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AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep

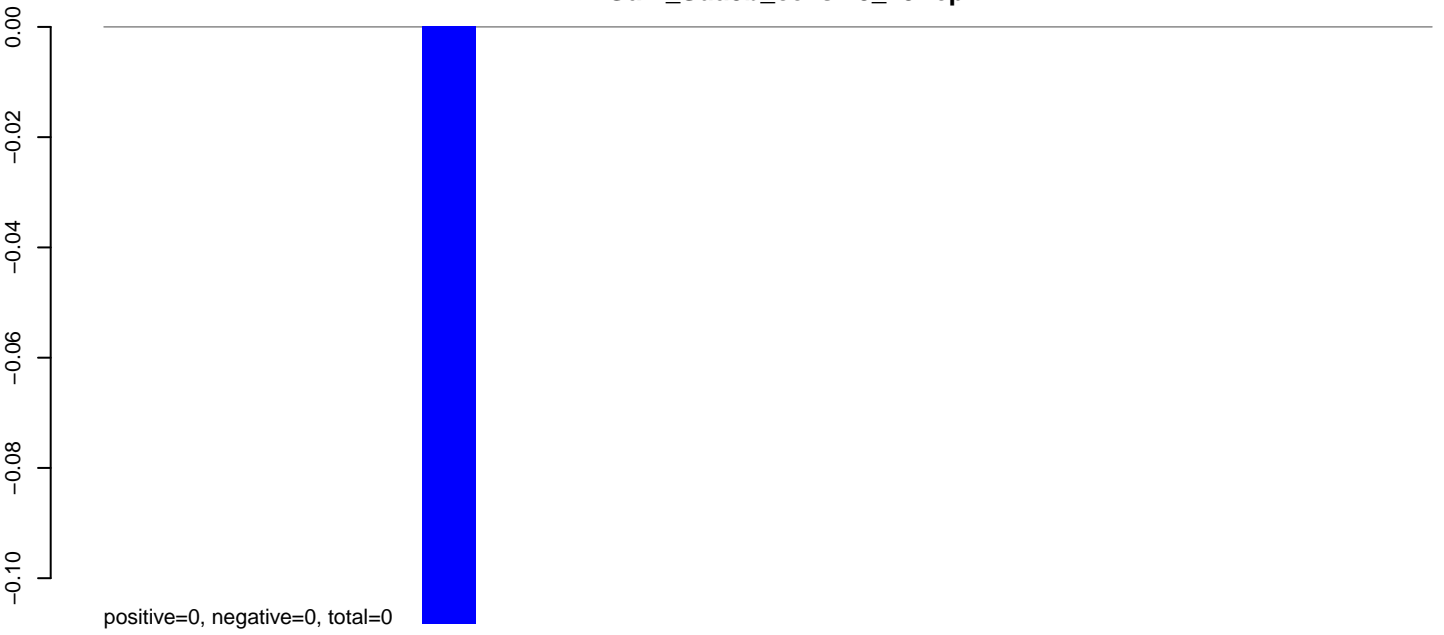


AnGam_Sua5b_cells.rep

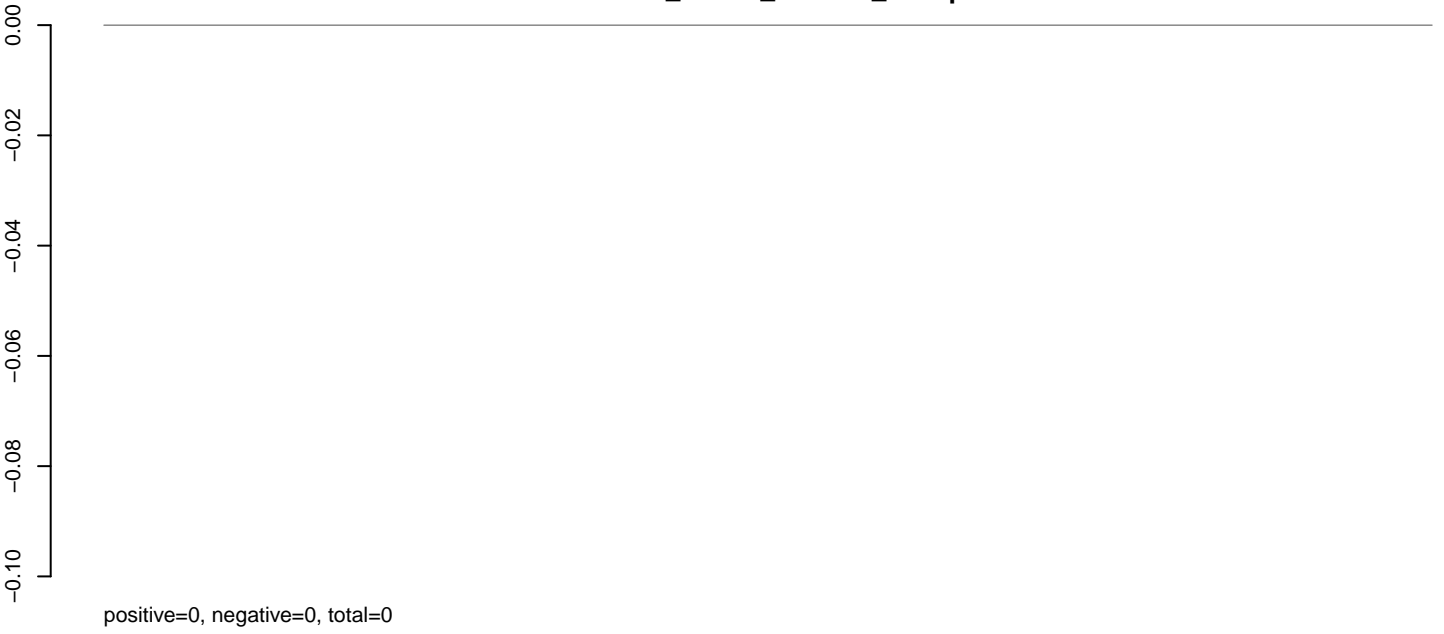


200 400 600 800 1000

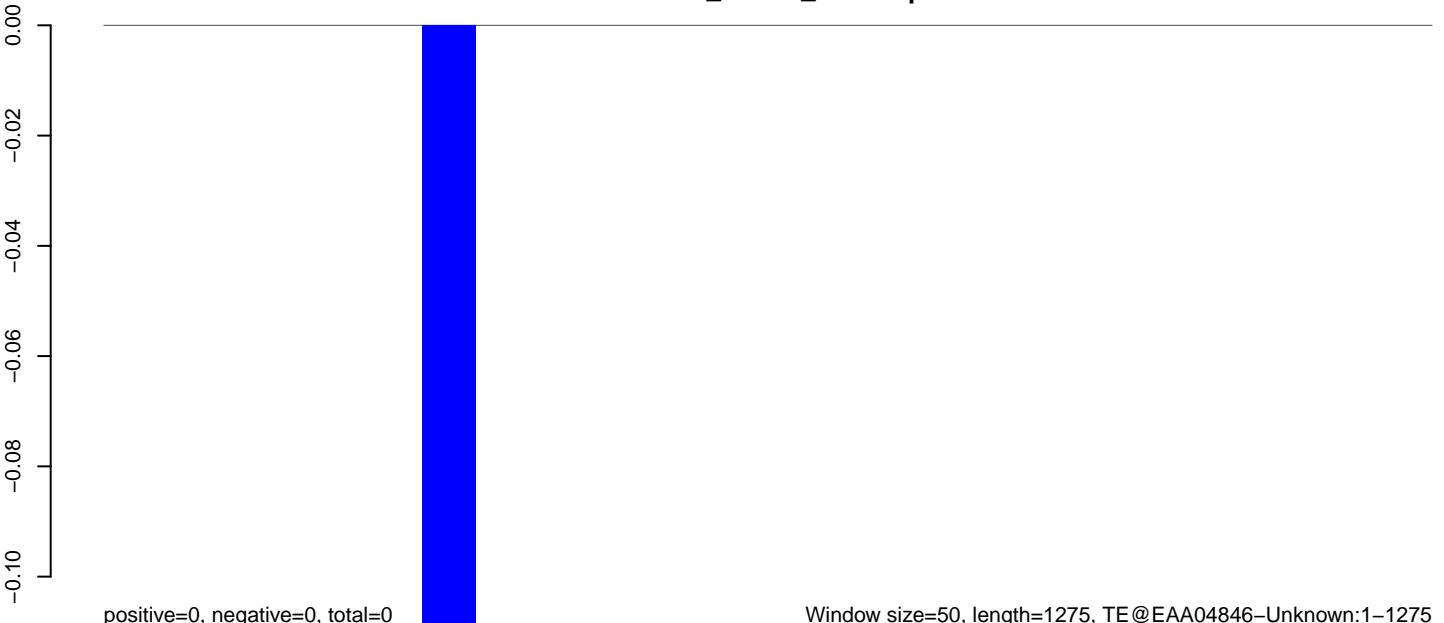
AnGam_Sua5b_cells.18_23.rep



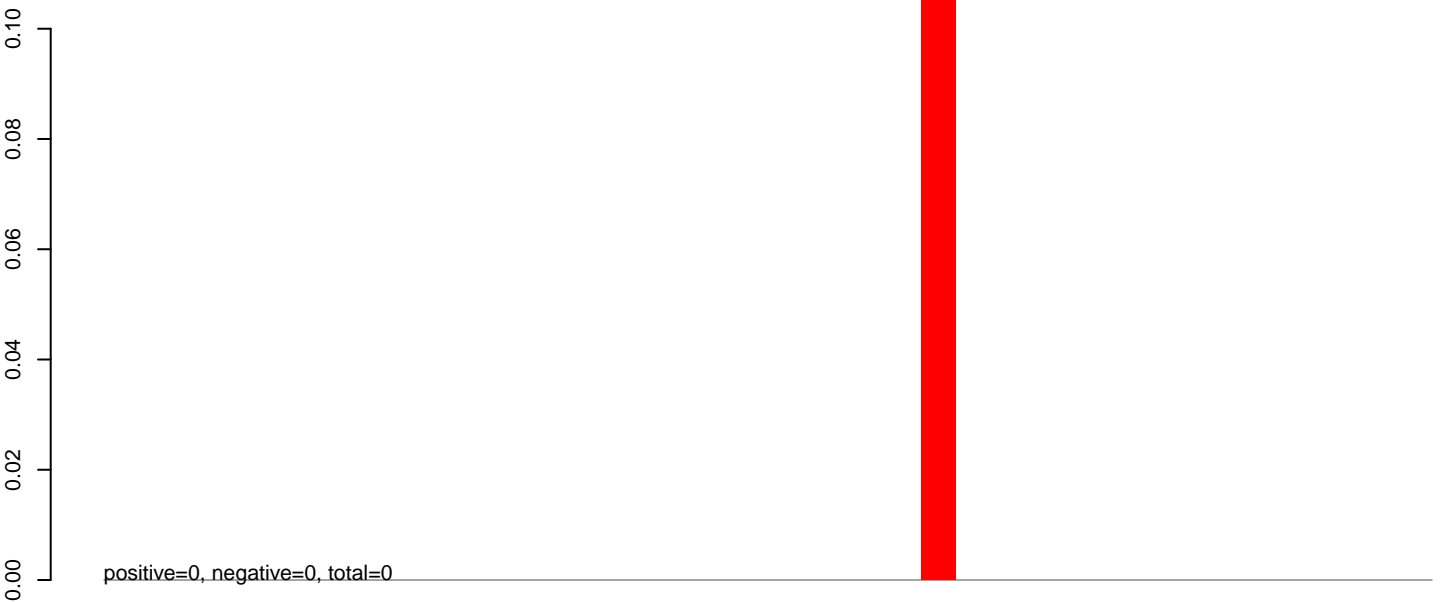
AnGam_Sua5b_cells.24_35.rep



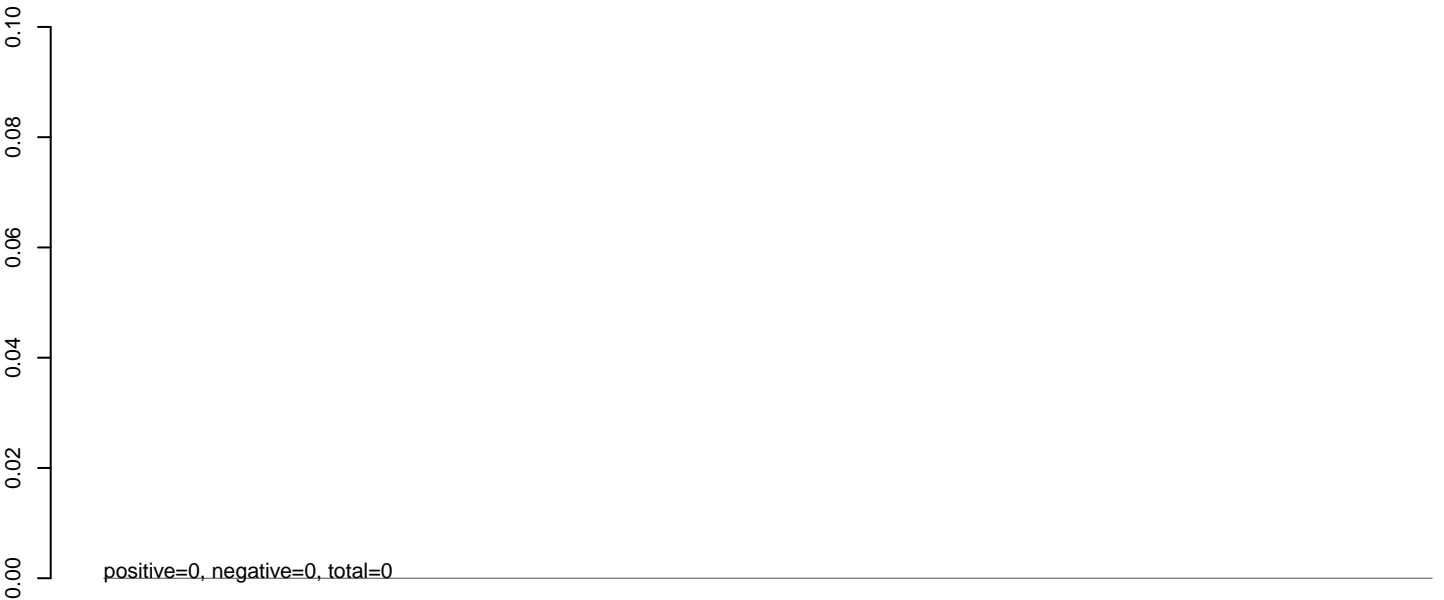
AnGam_Sua5b_cells.rep



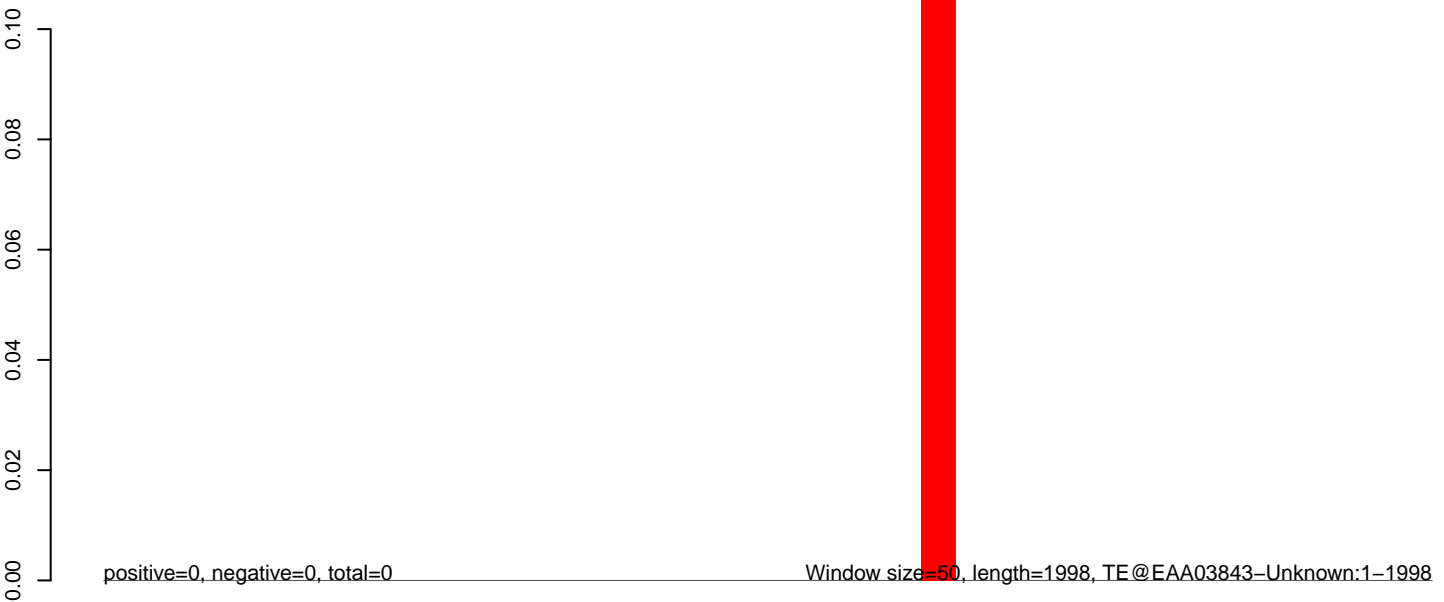
AnGam_Sua5b_cells.18_23.rep



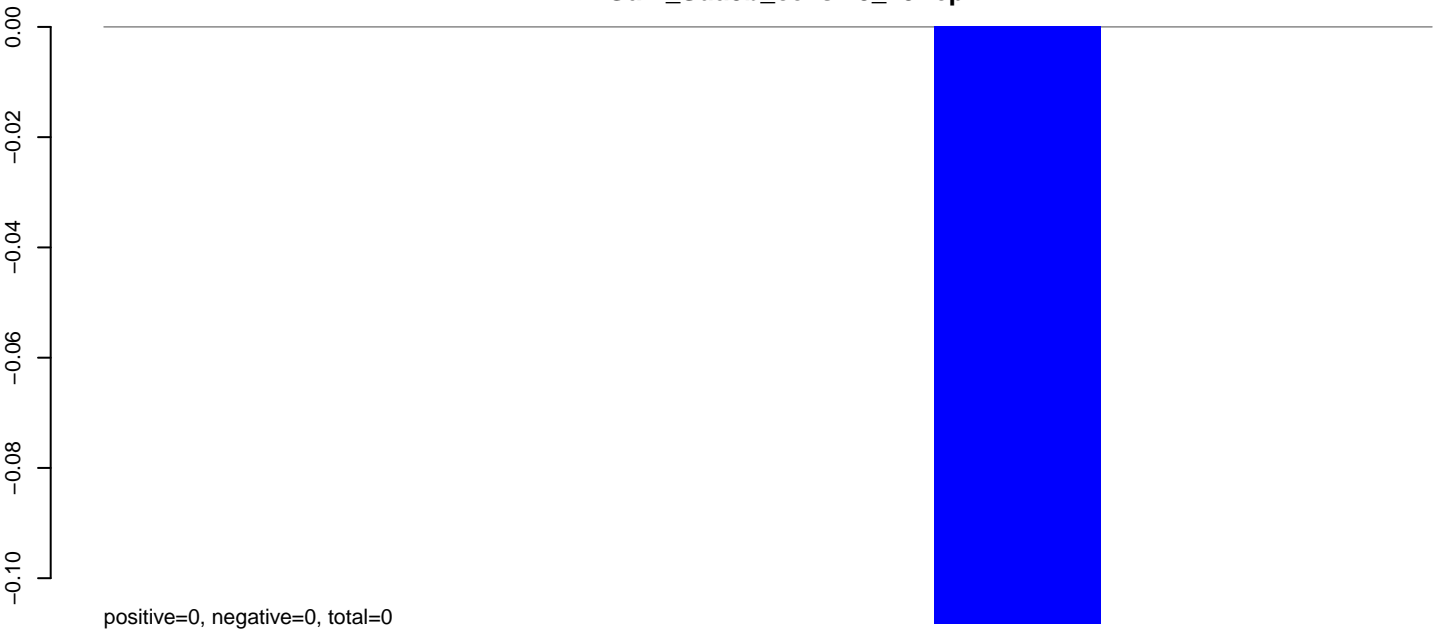
AnGam_Sua5b_cells.24_35.rep



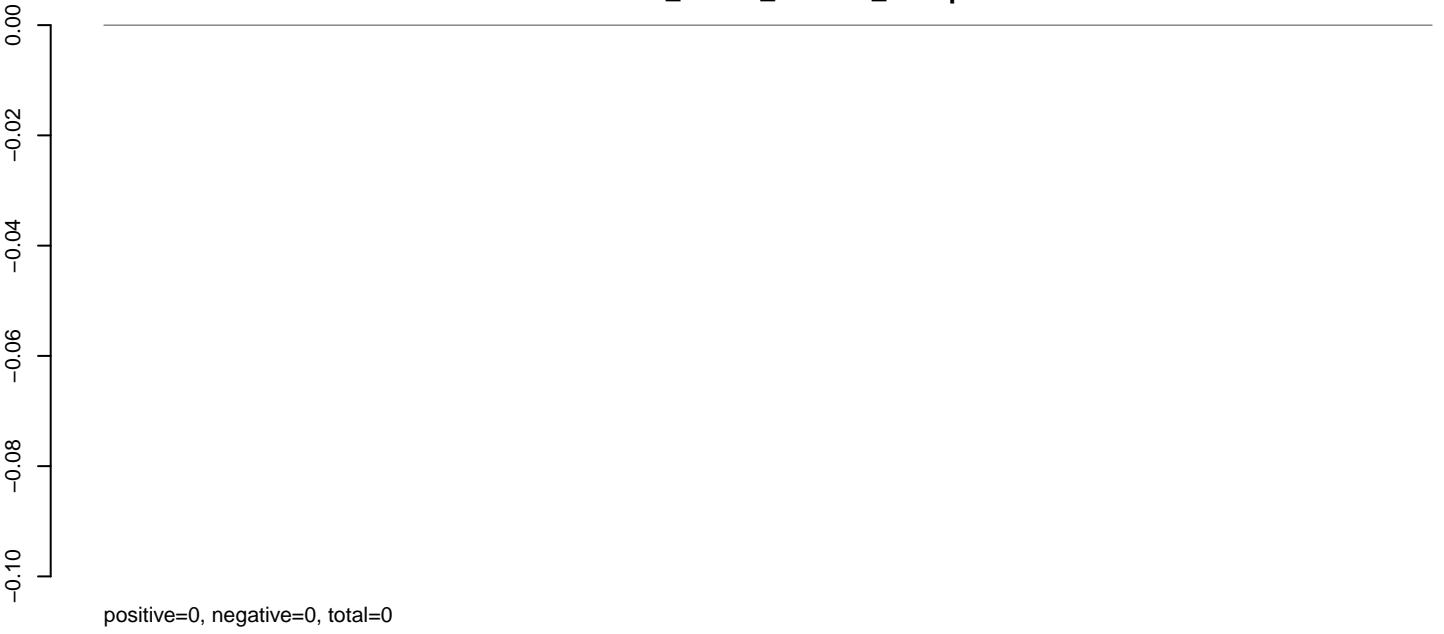
AnGam_Sua5b_cells.rep



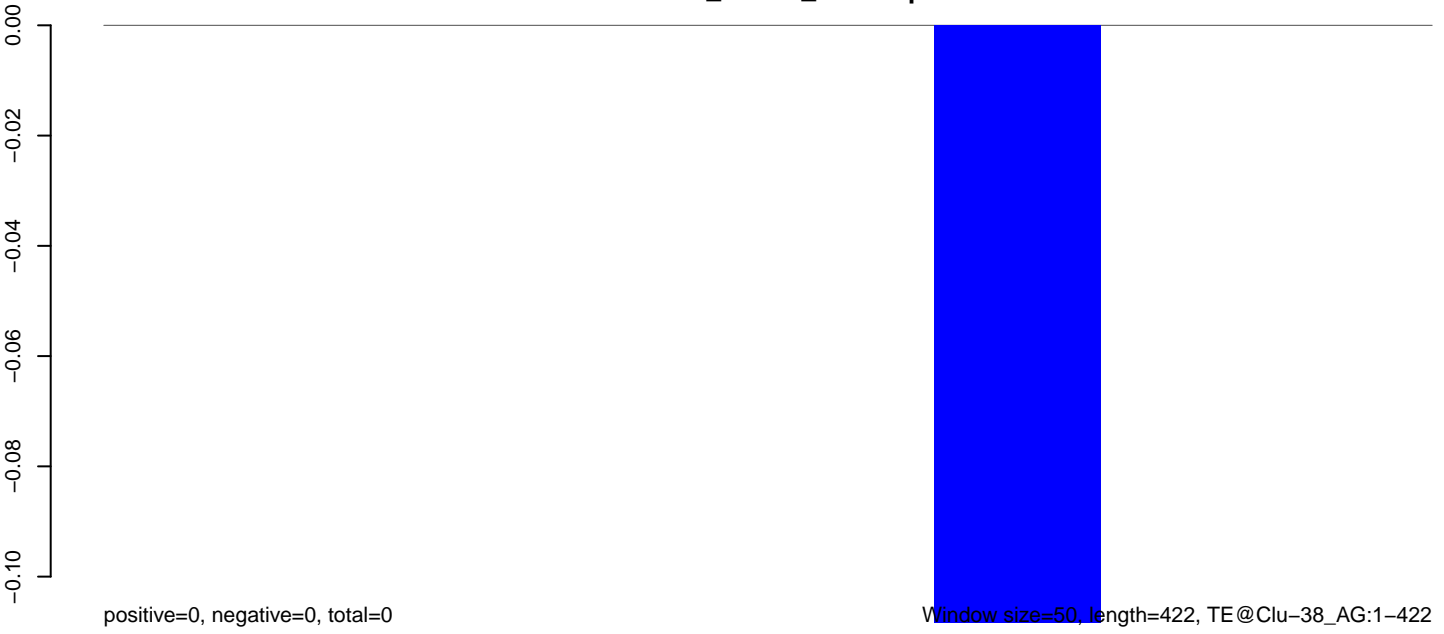
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



Window size=50, length=422, TE@Clu-38_AG:1-422

100 200 300 400

AnGam_Sua5b_cells.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AnGam_Sua5b_cells.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

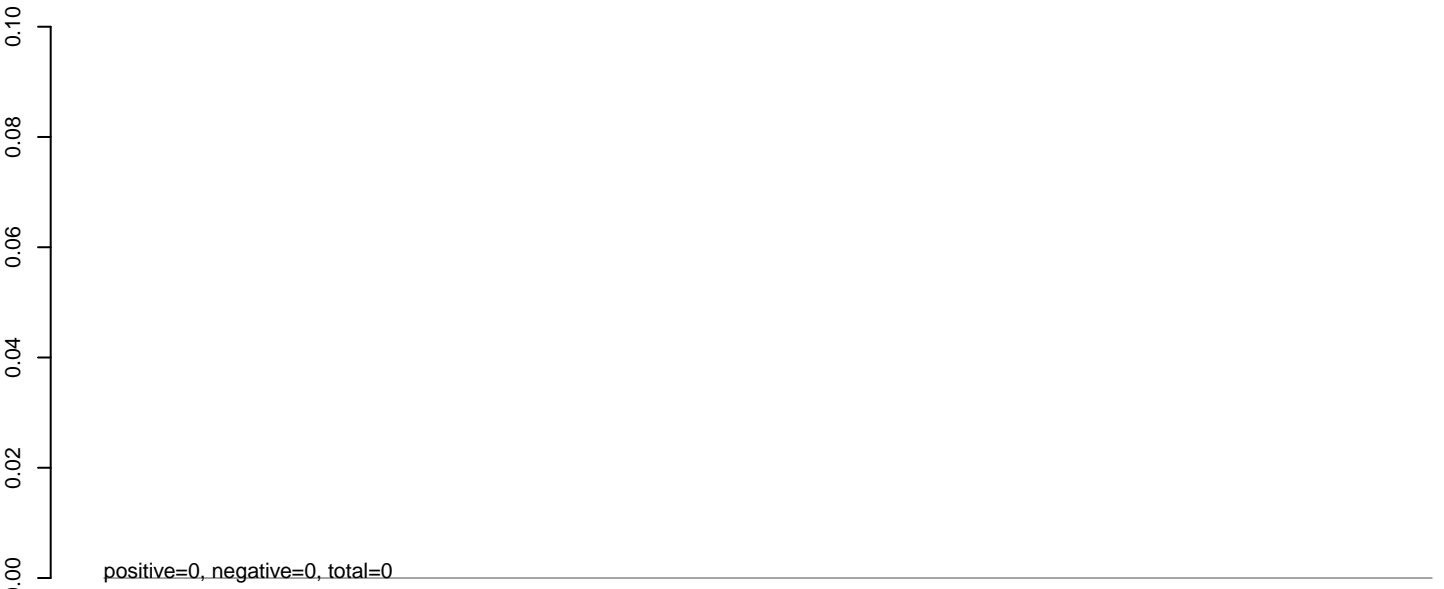
positive=0, negative=0, total=0

Window size=50, length=463, TE @ Agam_L10438_mariner-DNA-mariner:1-463

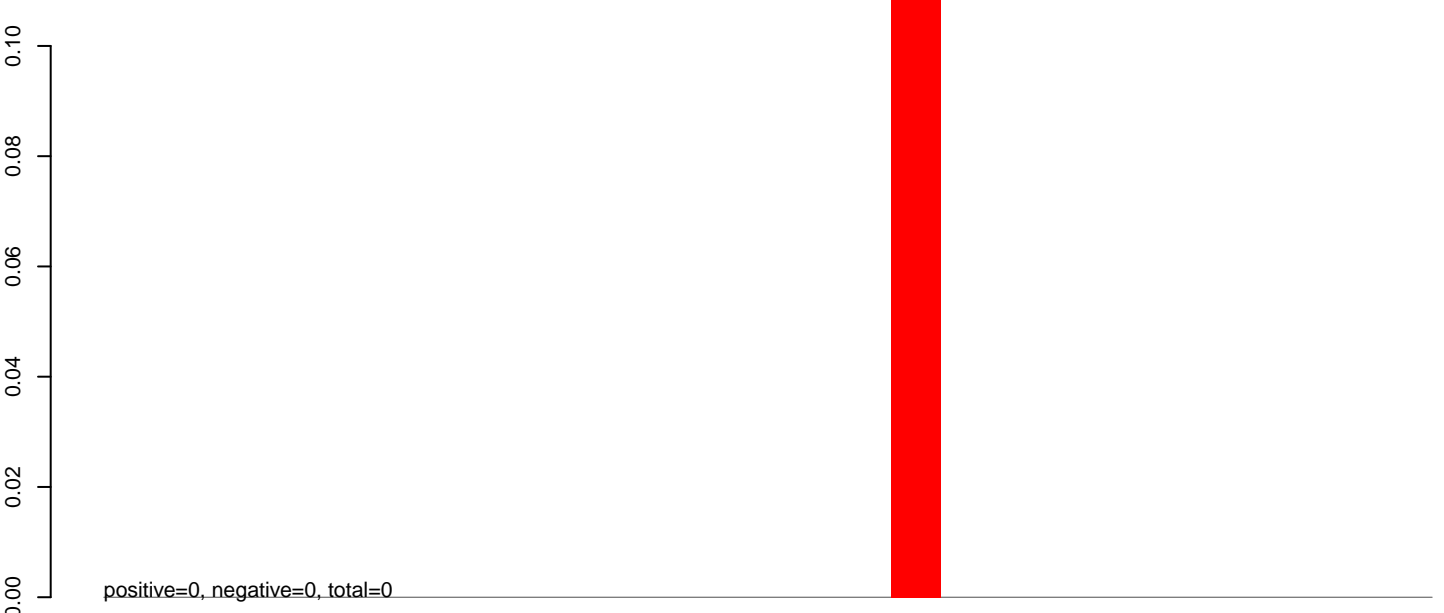
100 200 300 400 500



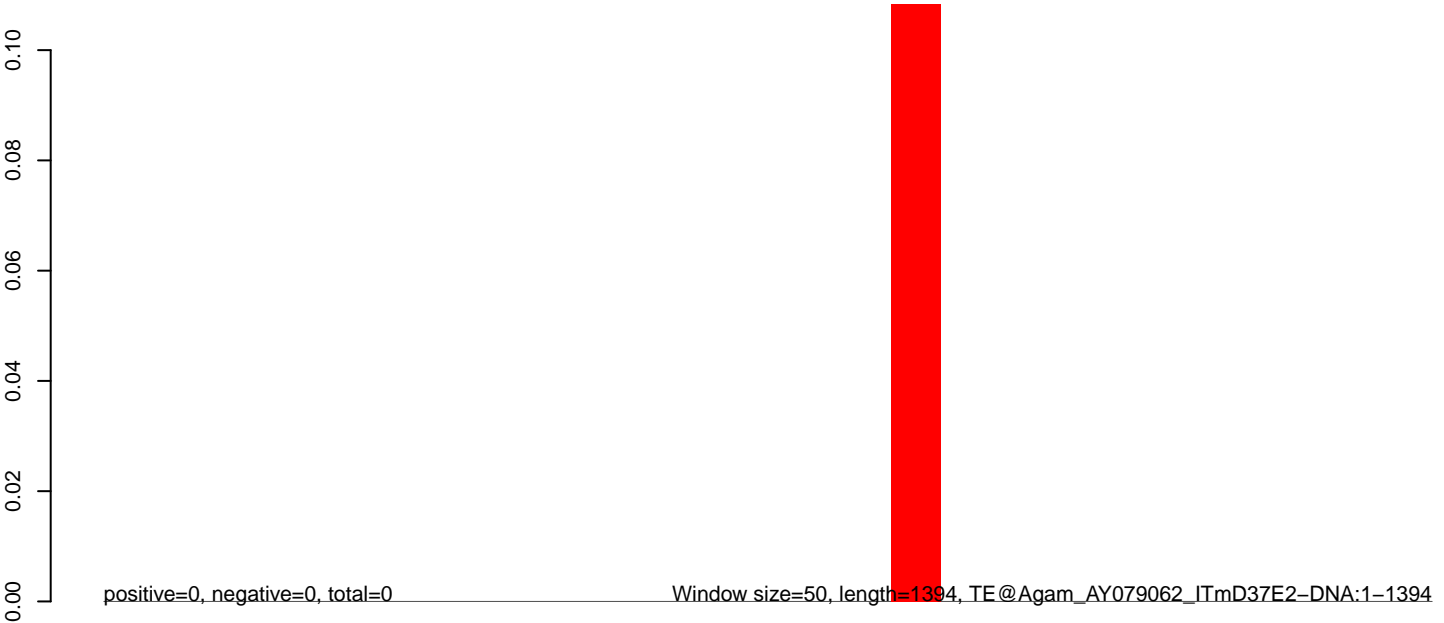
AnGam_Sua5b_cells.18_23.rep



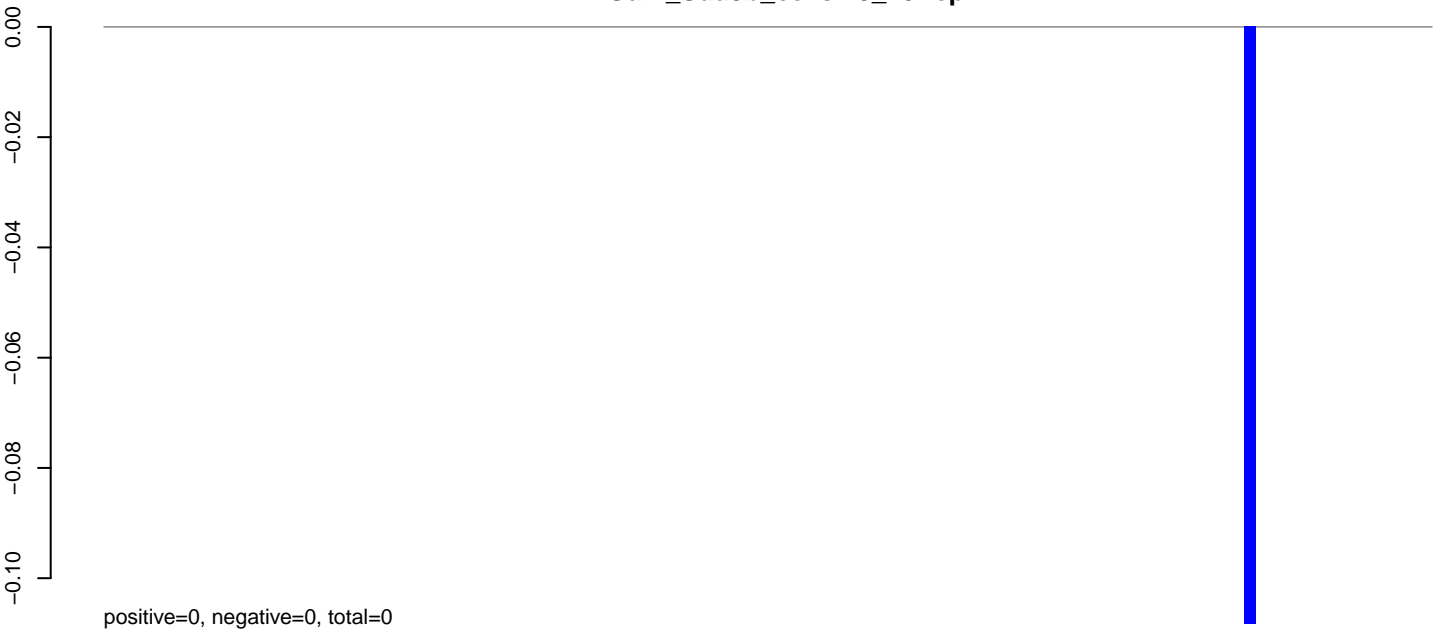
AnGam_Sua5b_cells.24_35.rep



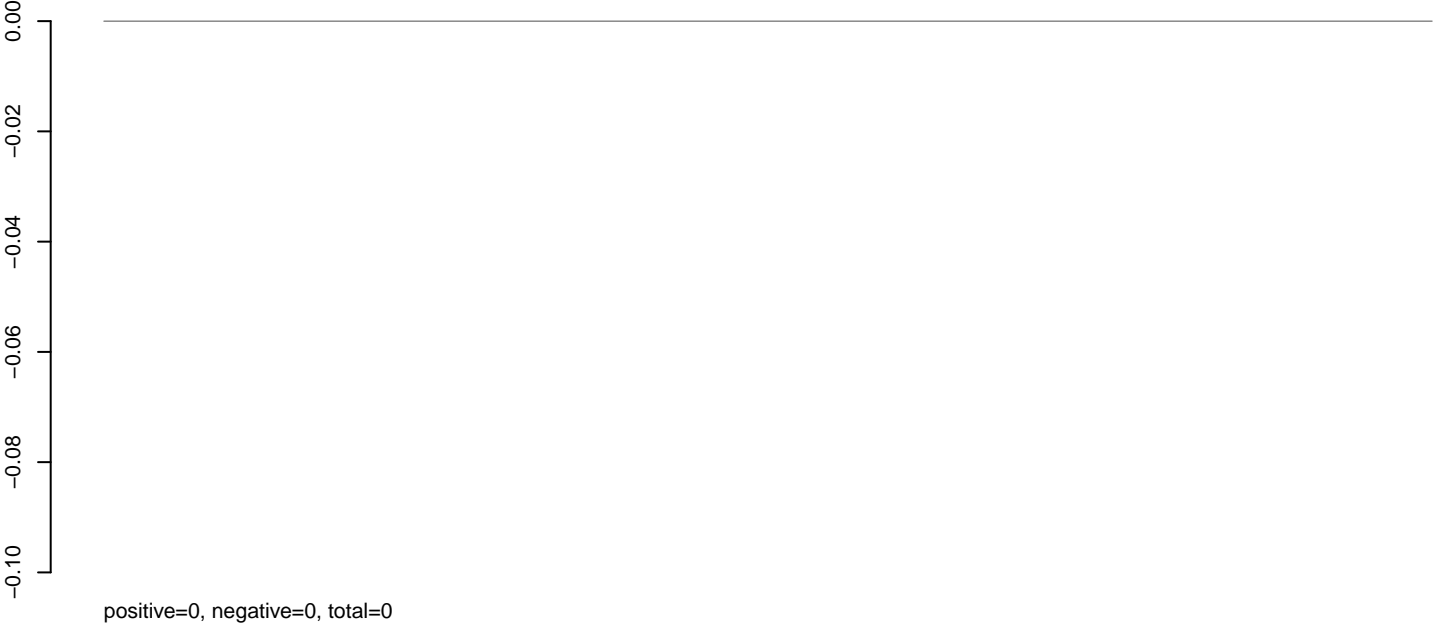
AnGam_Sua5b_cells.rep



AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



Window size=50, length=6047, TE@Agam_AB090824_KagaAg1-LINE-KagaAg1:1-6047

0 1000 2000 3000 4000 5000 6000

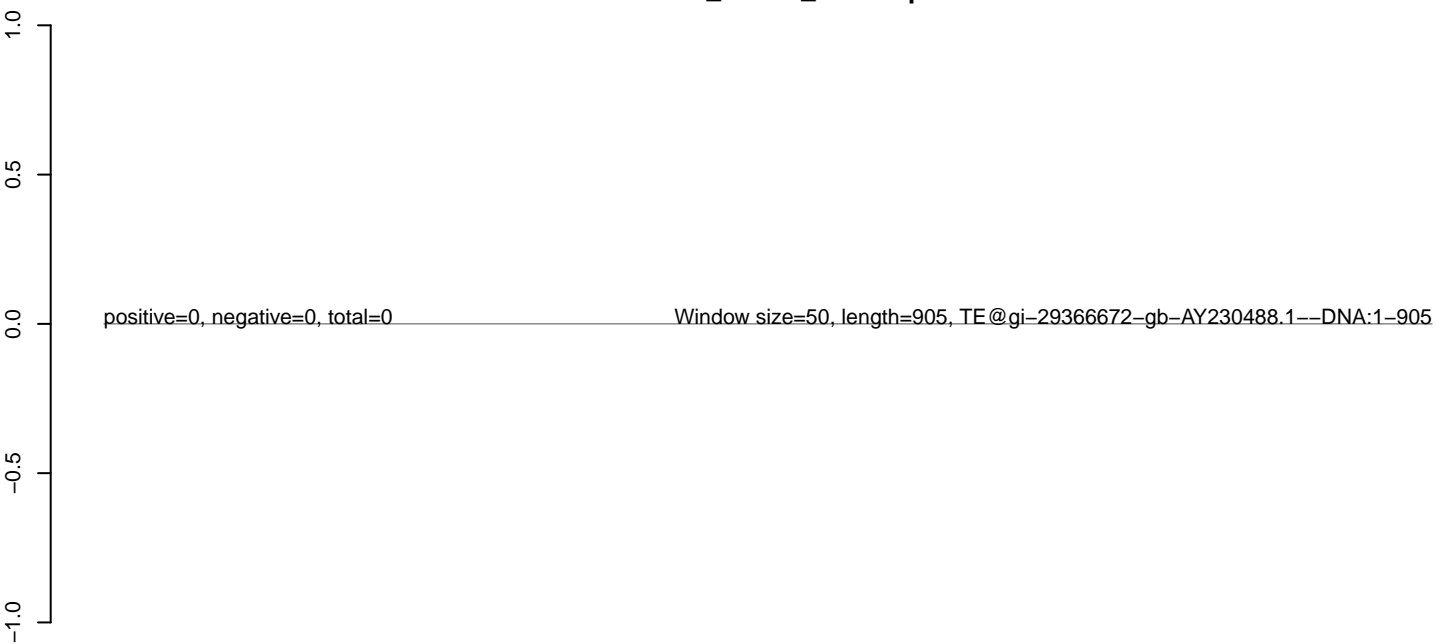
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



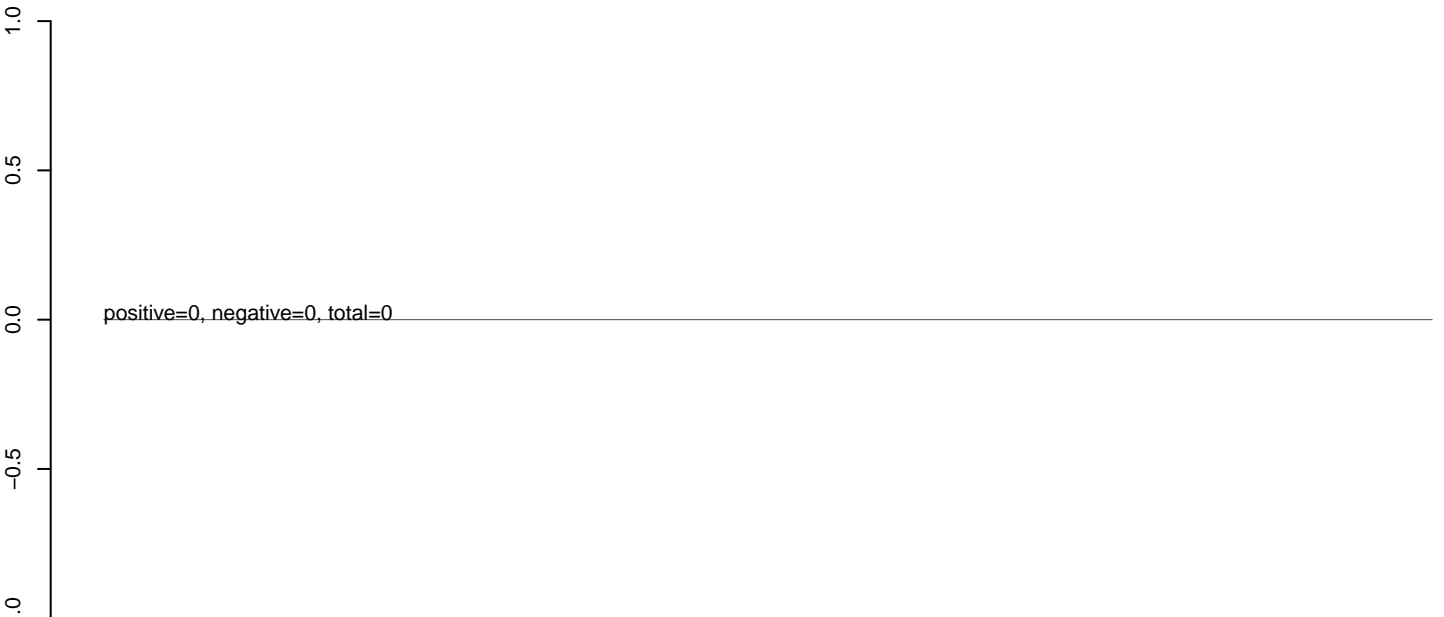
AnGam_Sua5b_cells.rep



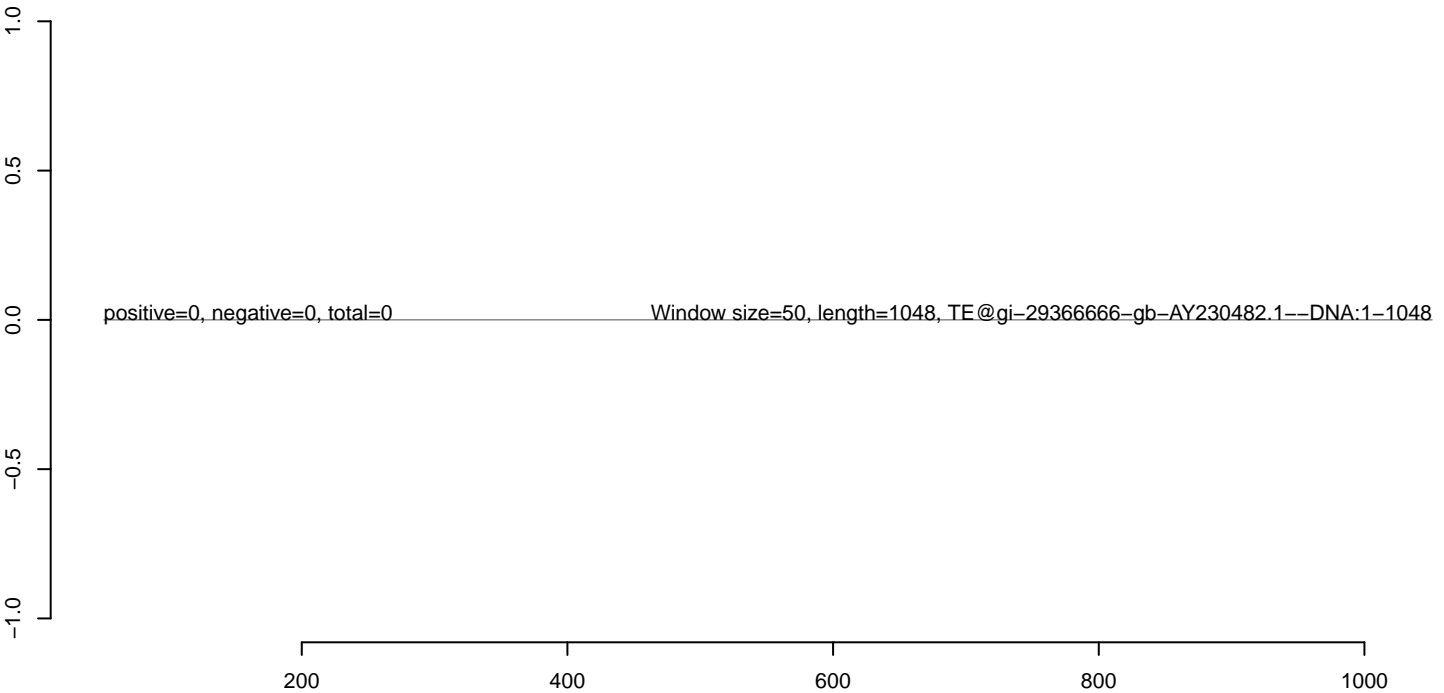
AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



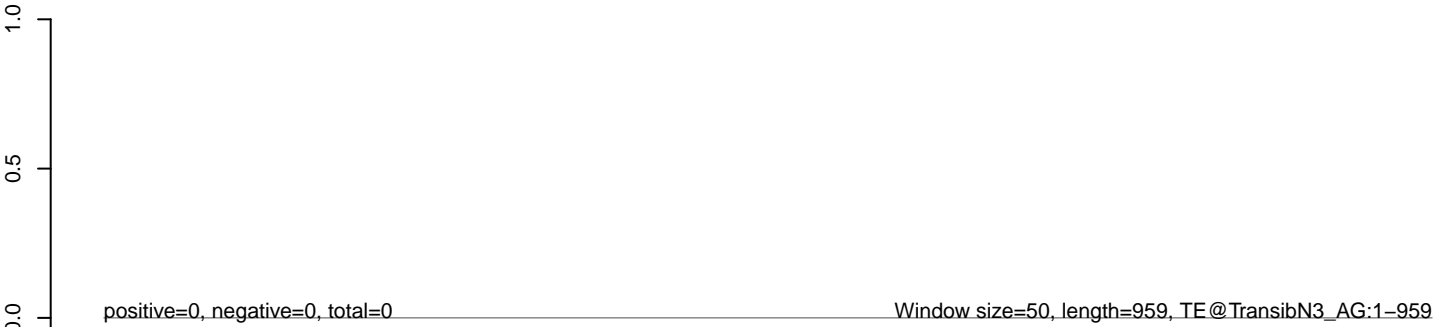
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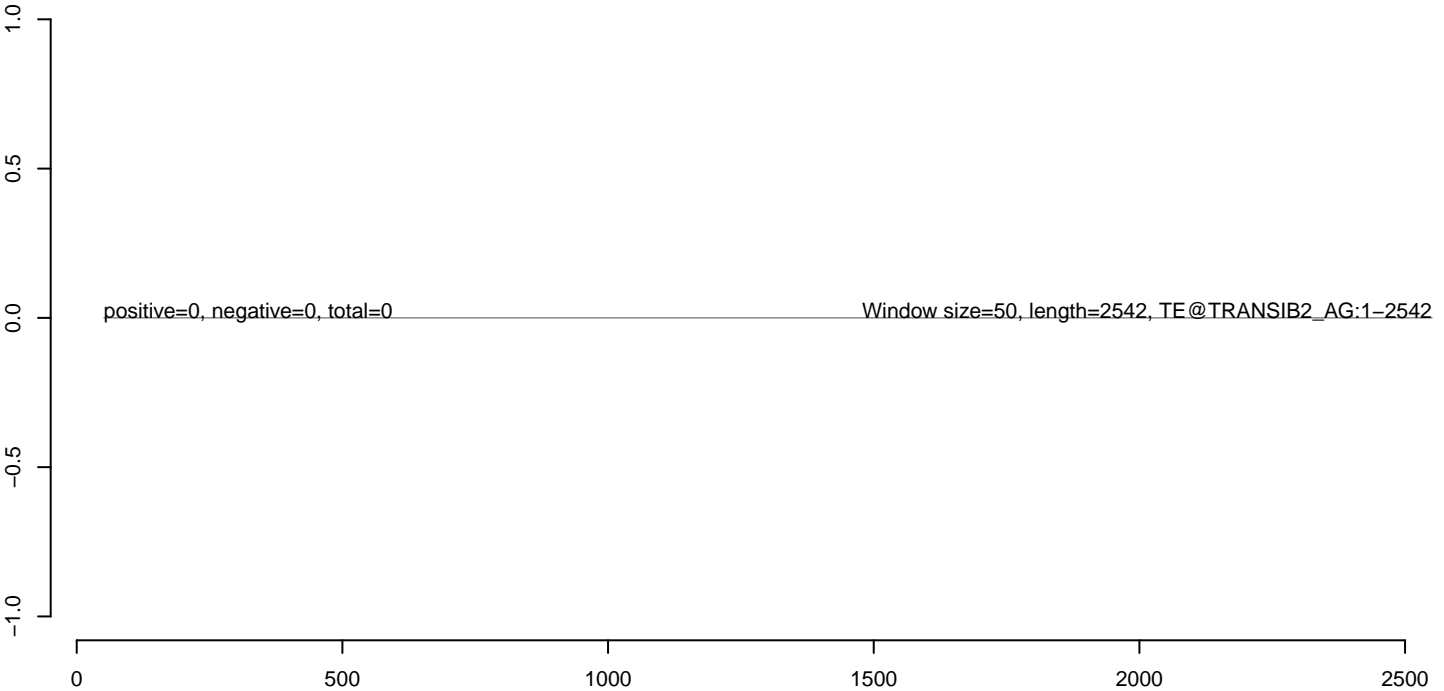
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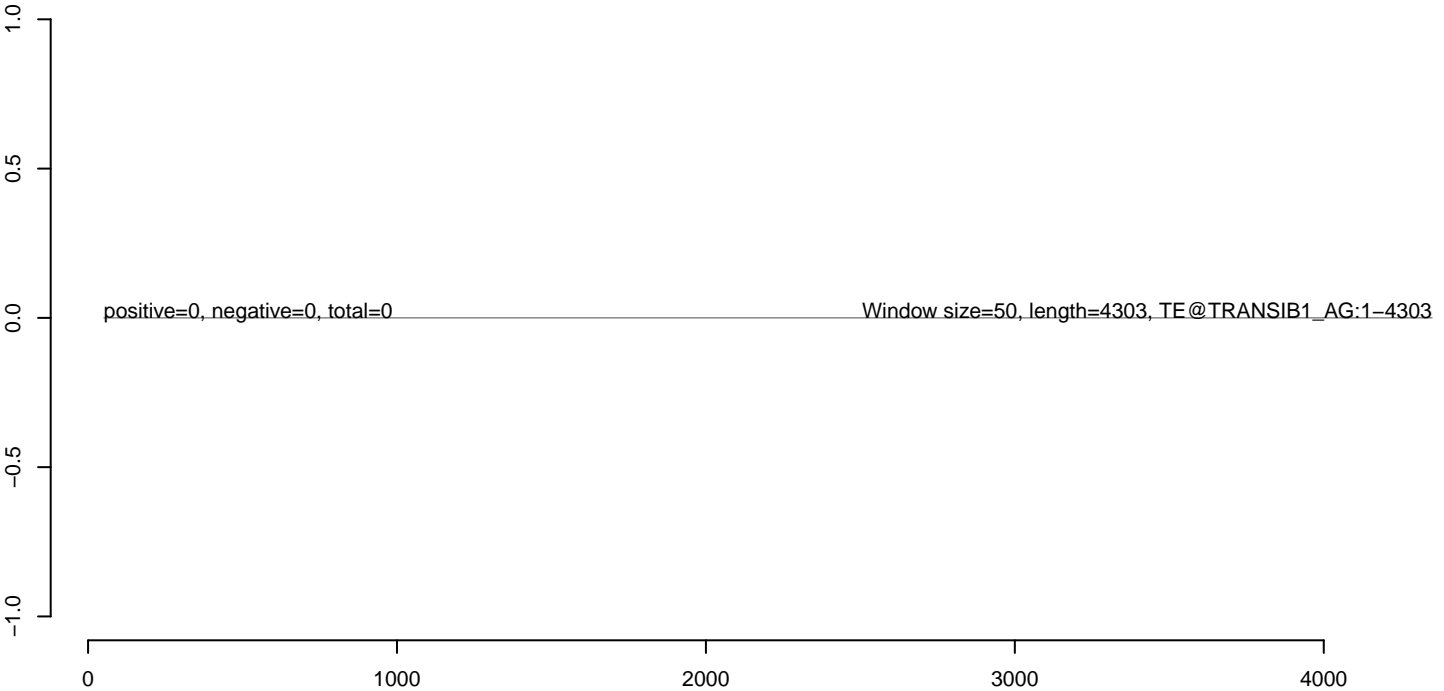
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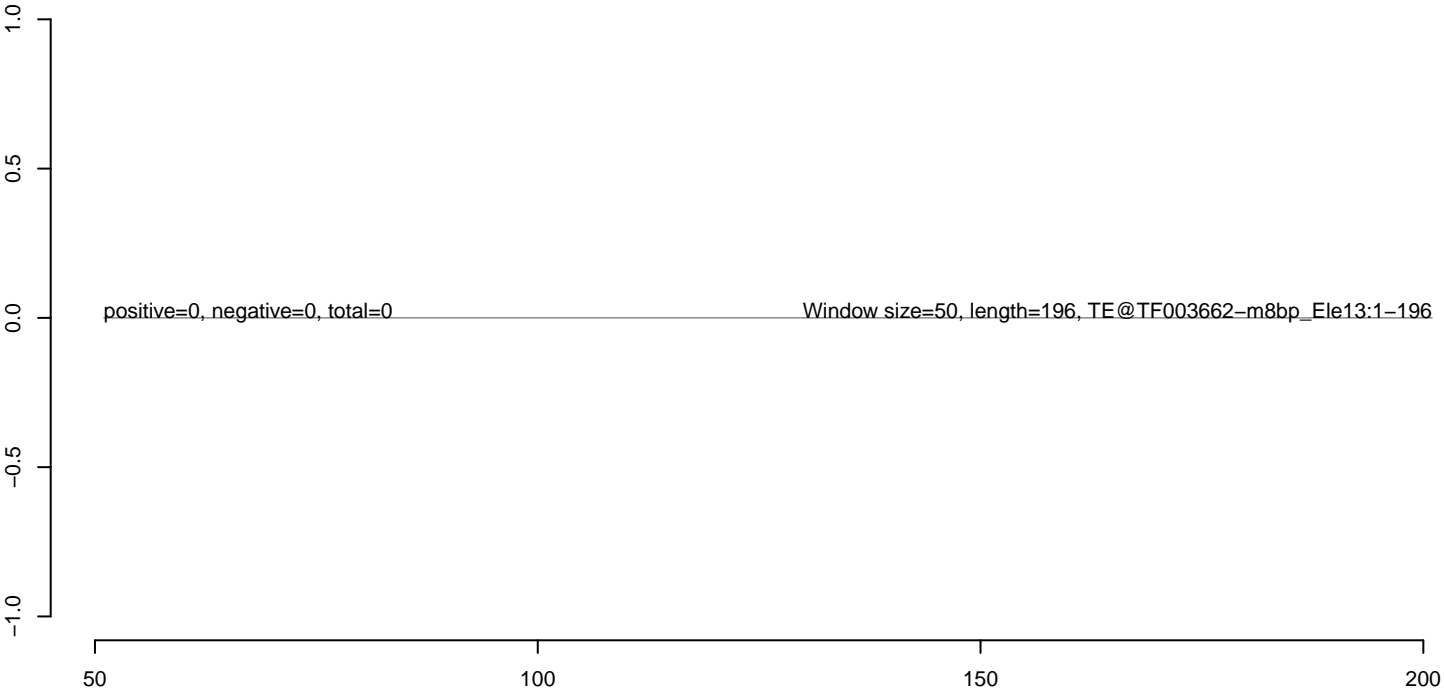
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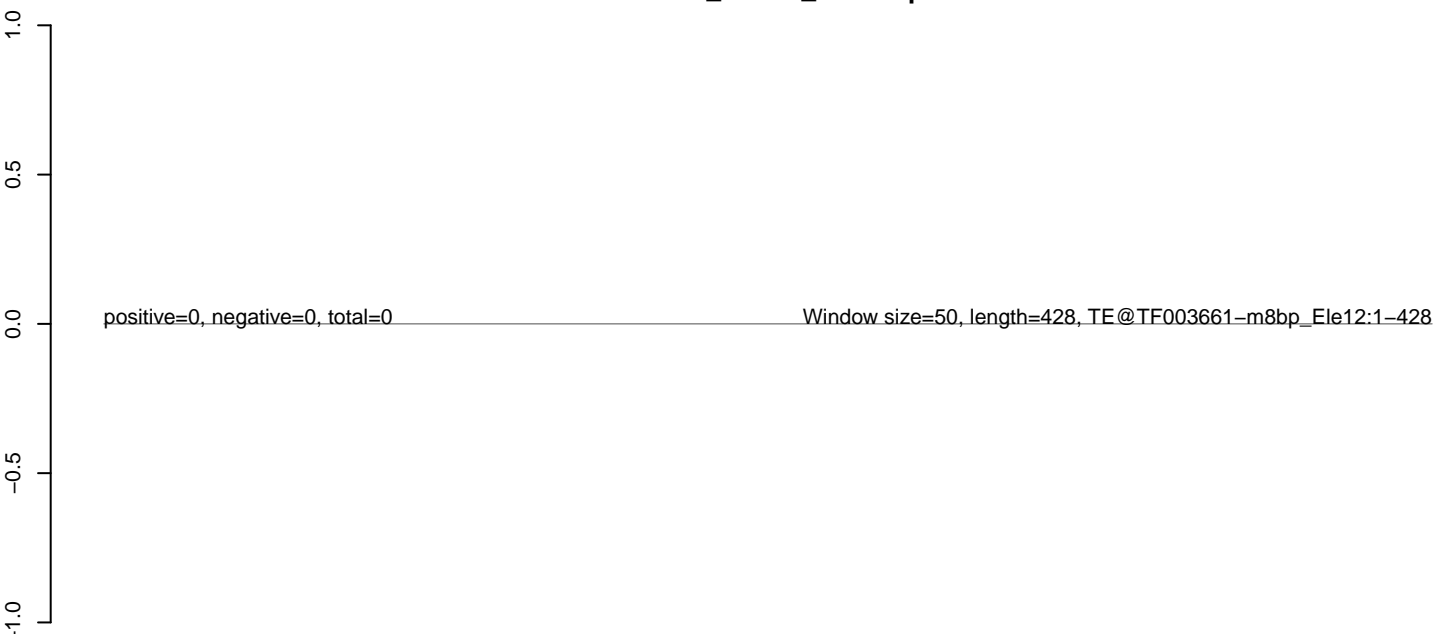
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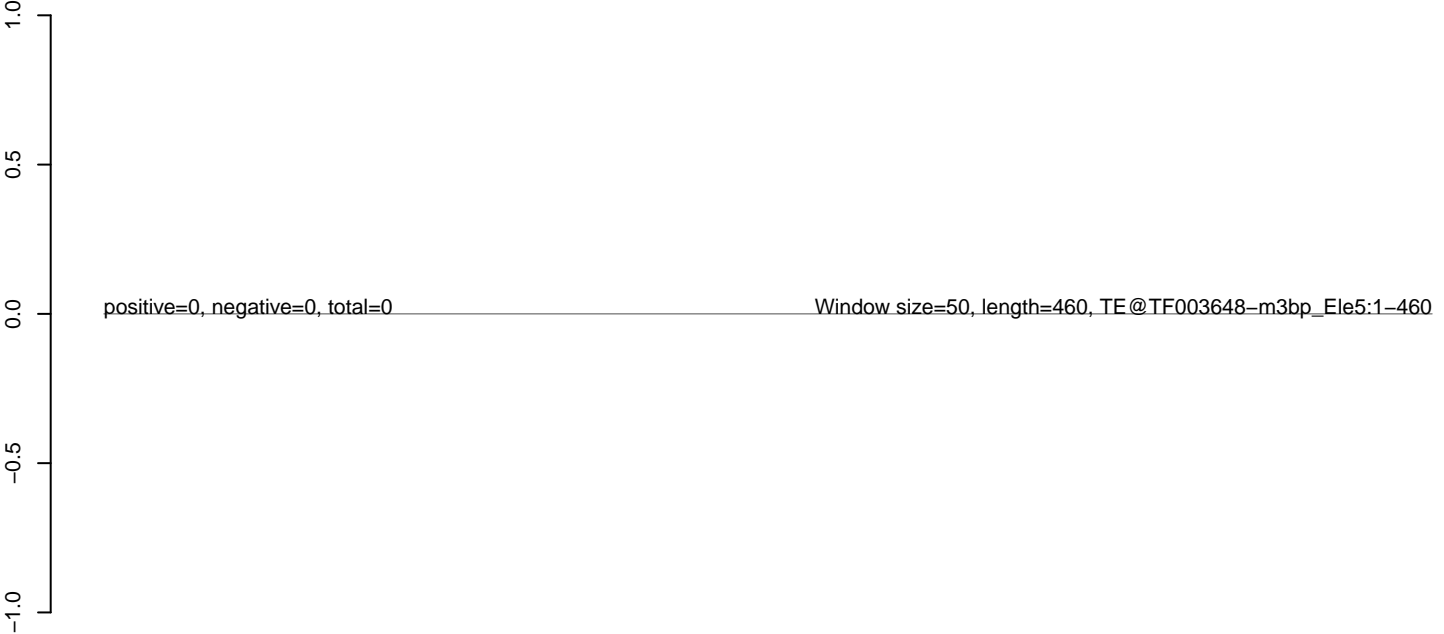
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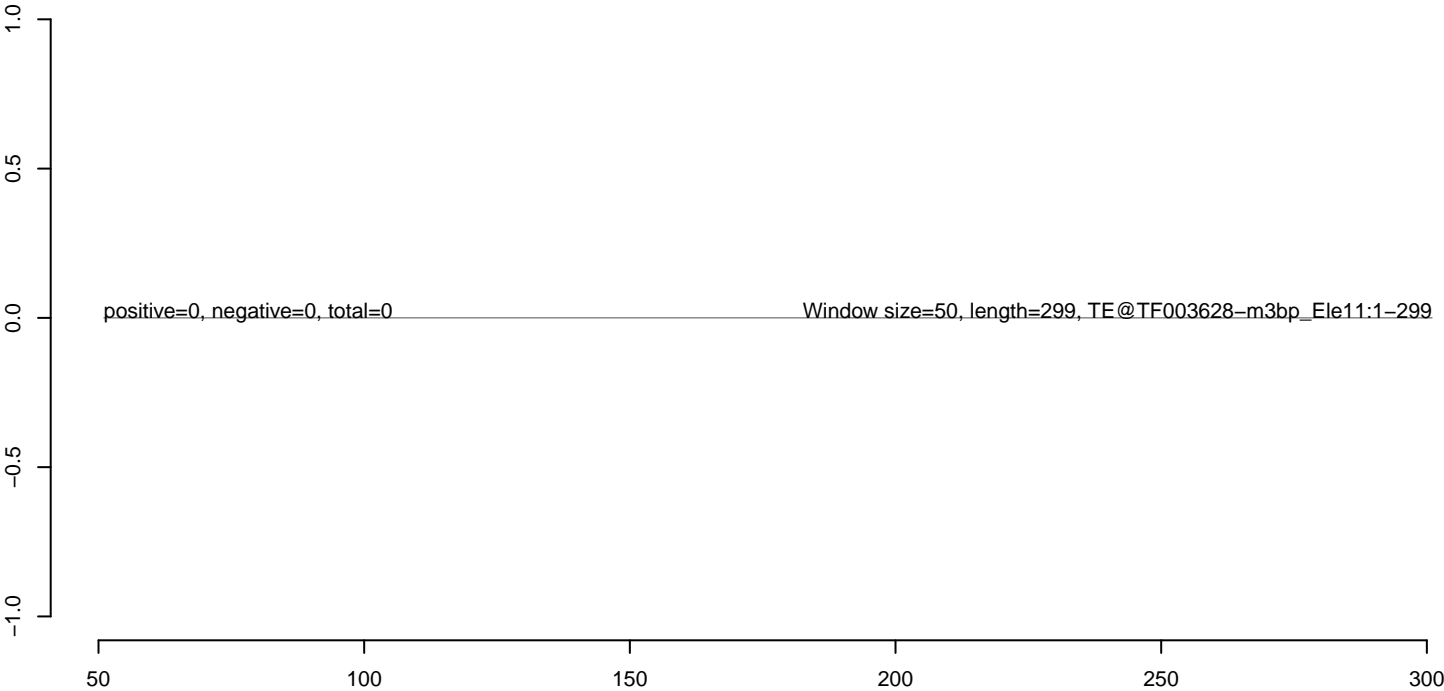
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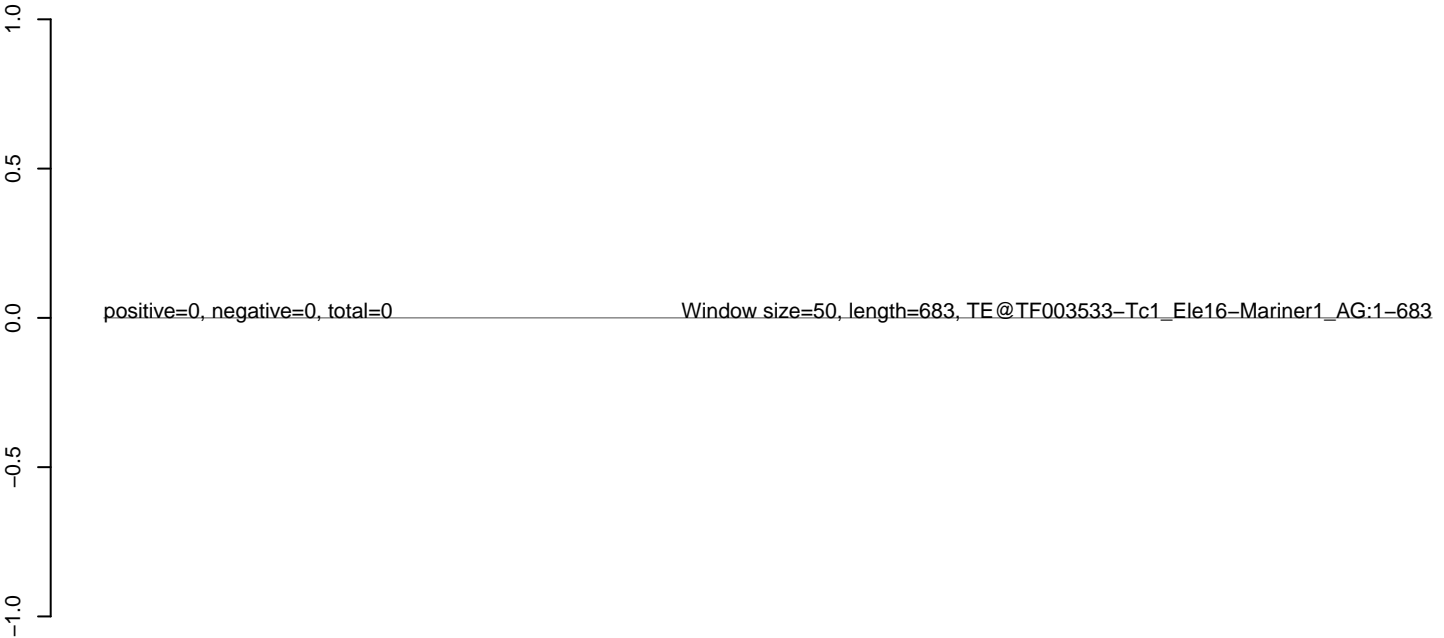
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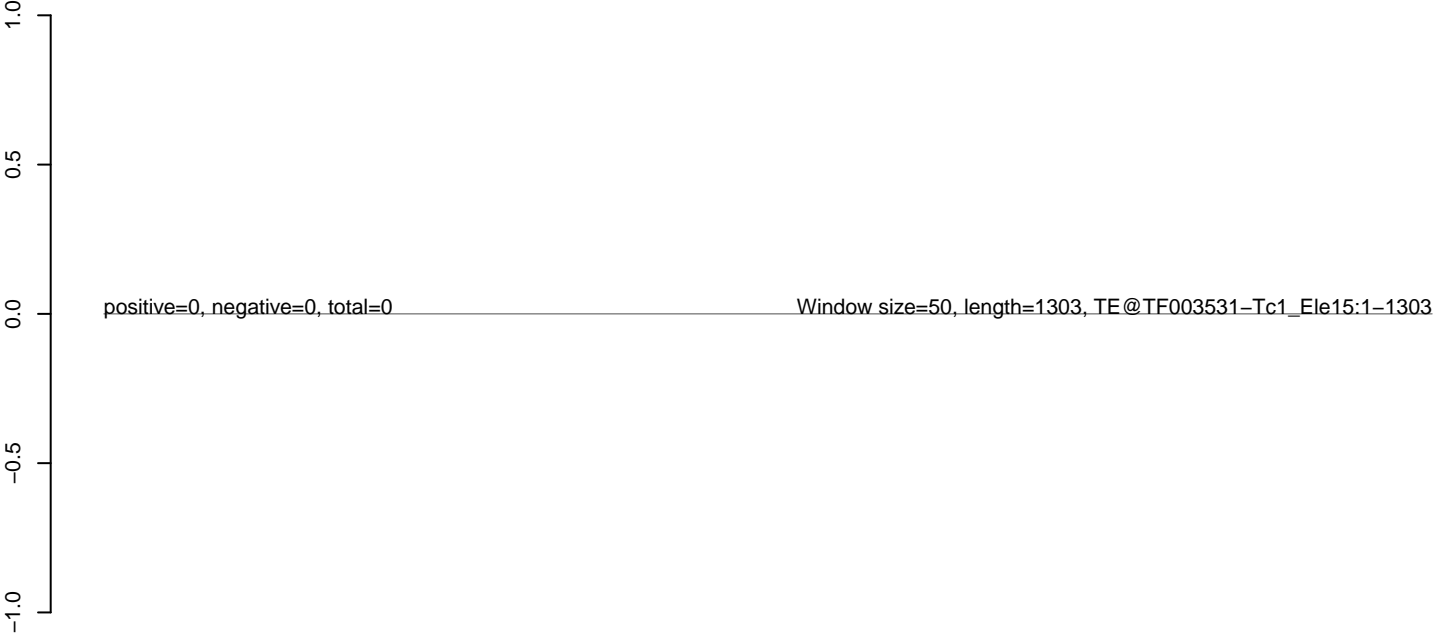
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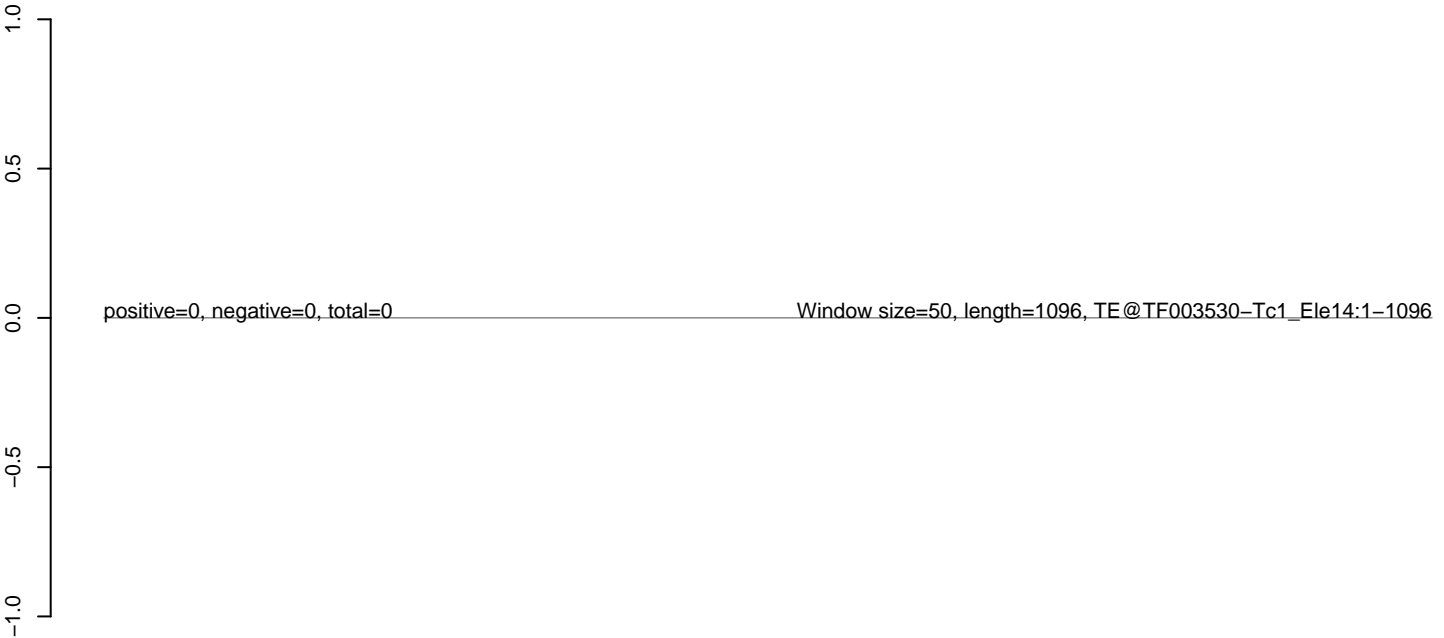
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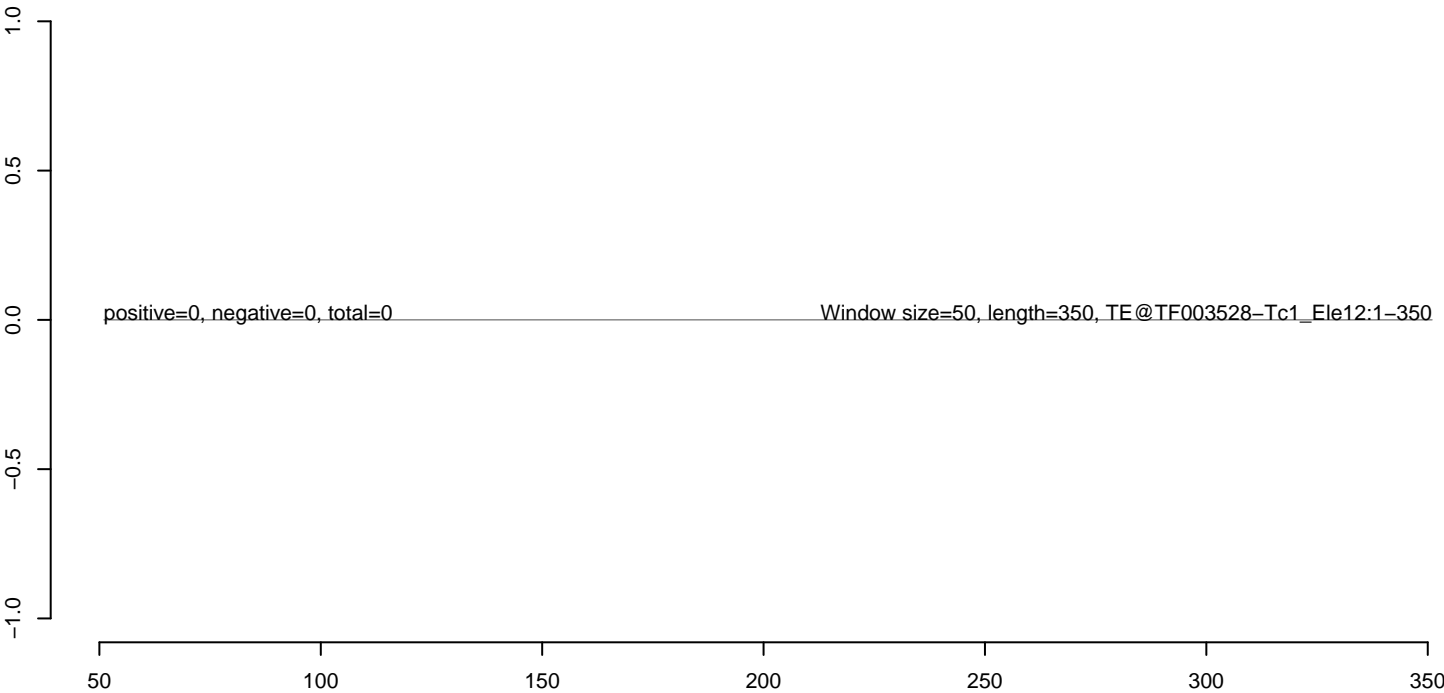
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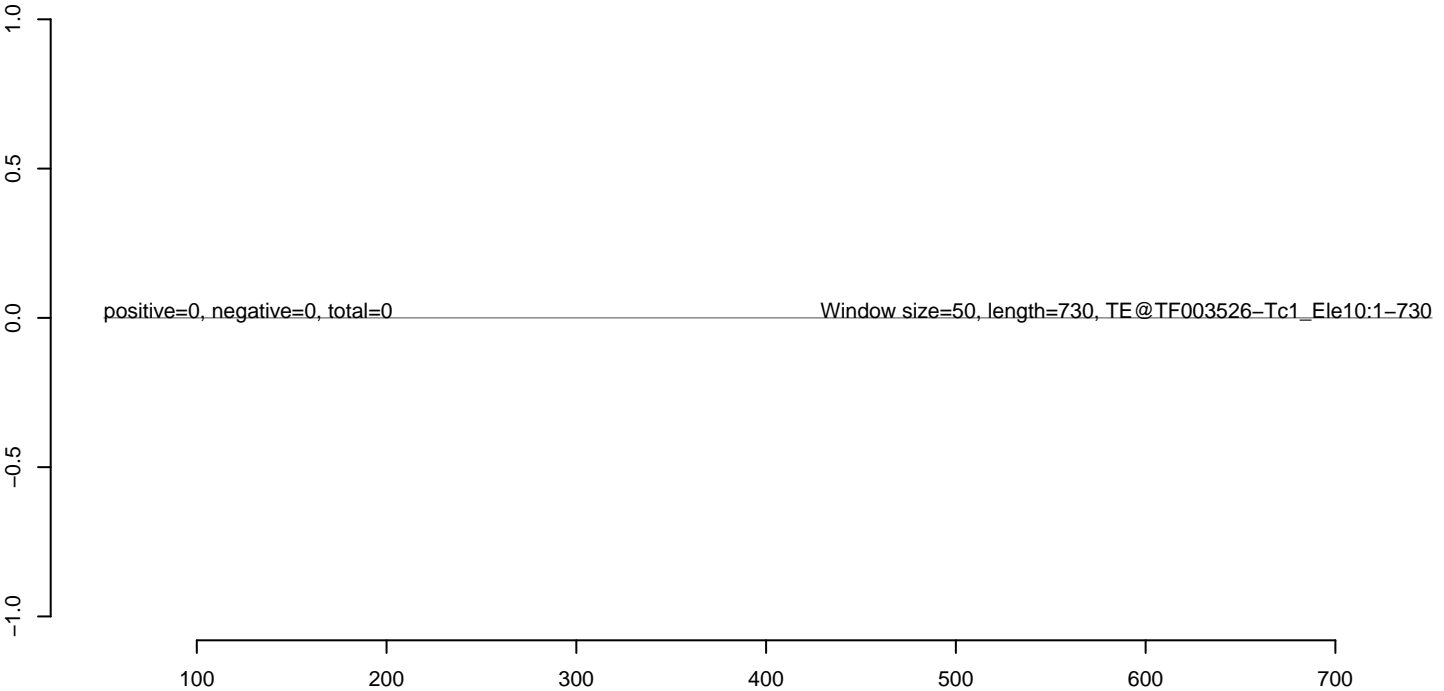
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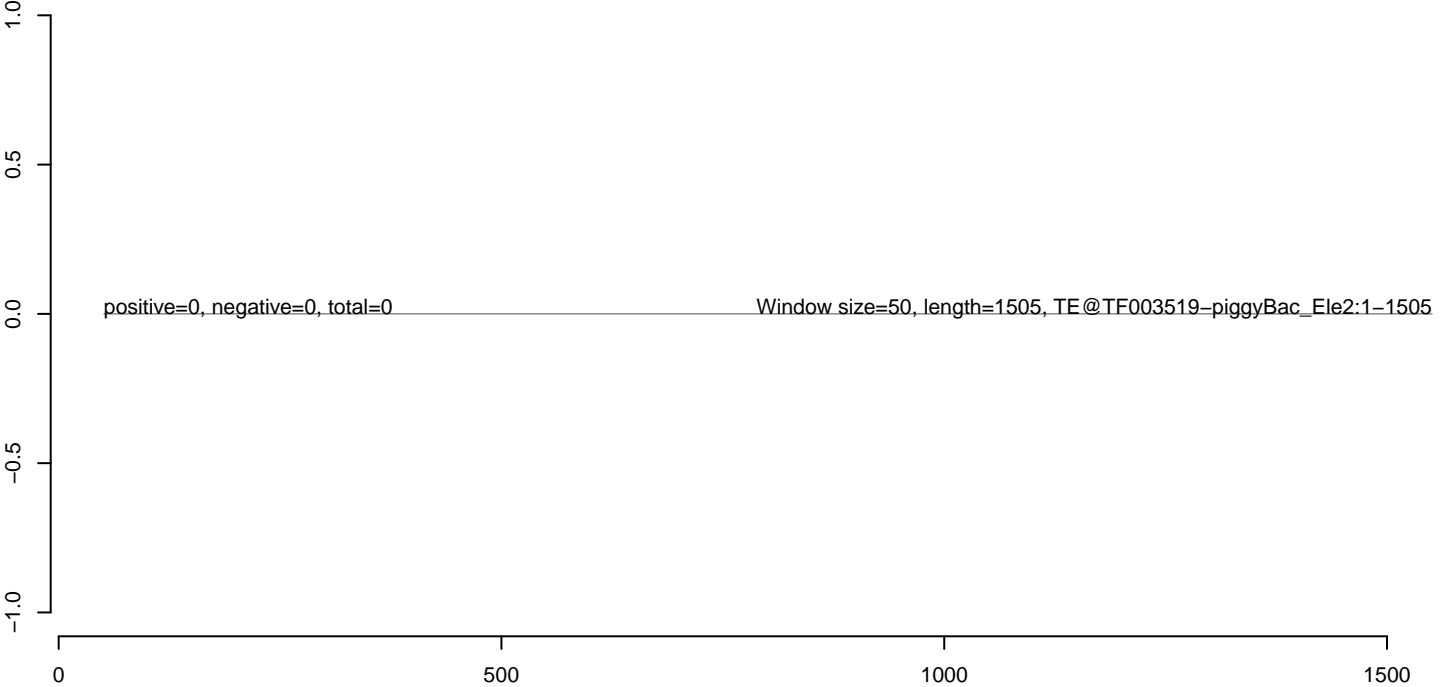
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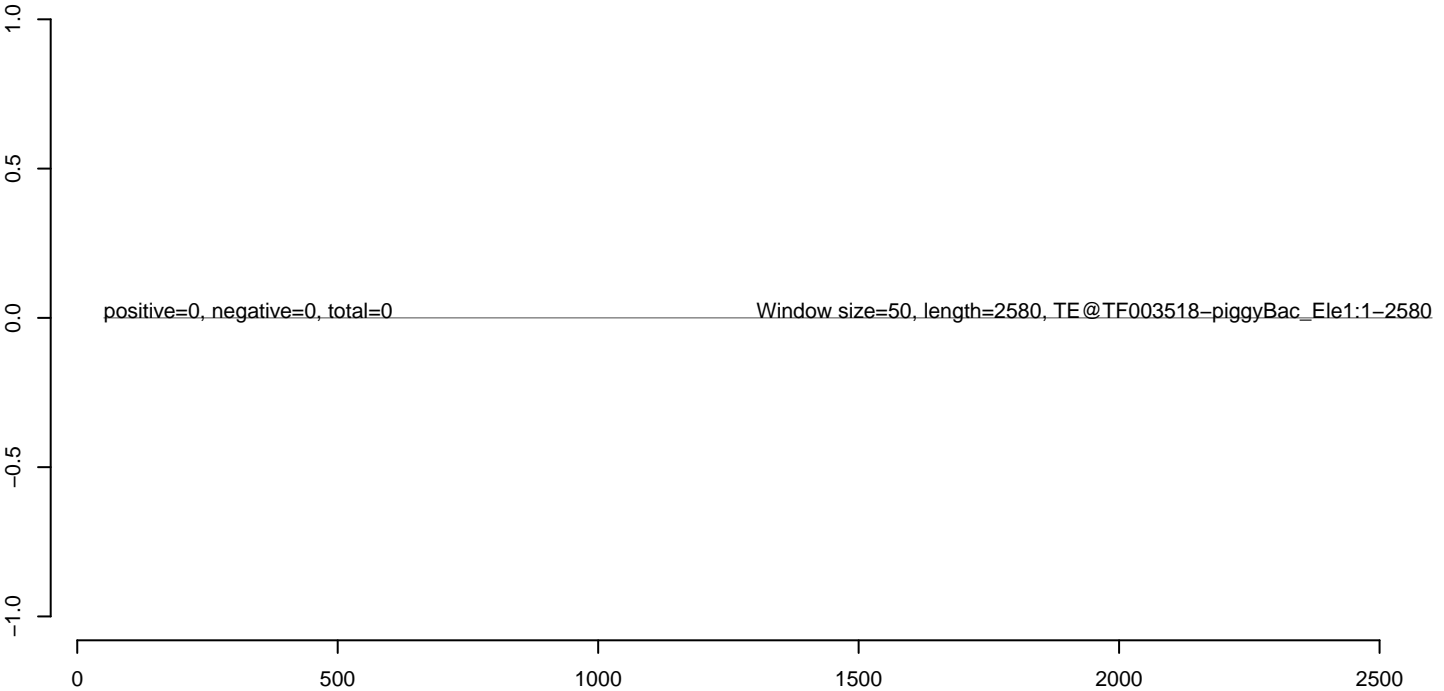
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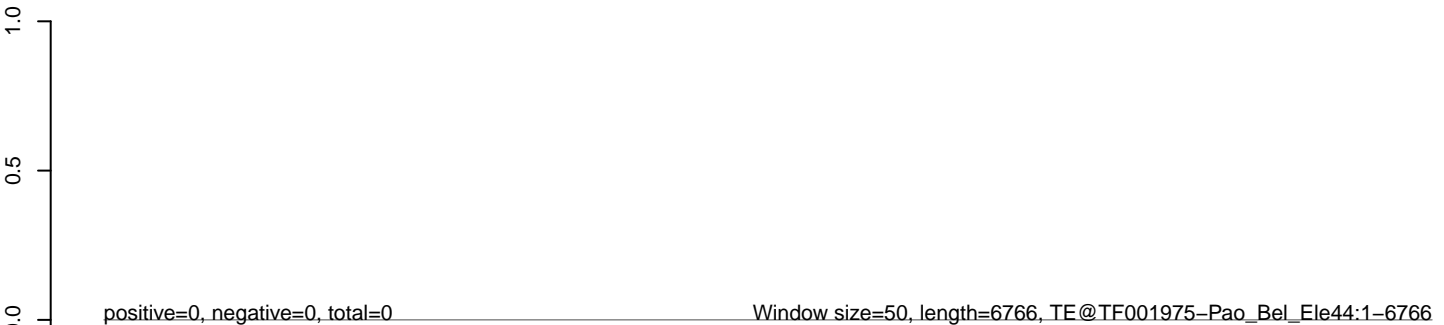
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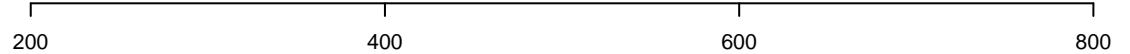
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AnGam_Sua5b_cells.rep



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AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



100 200 300 400 500 600 700

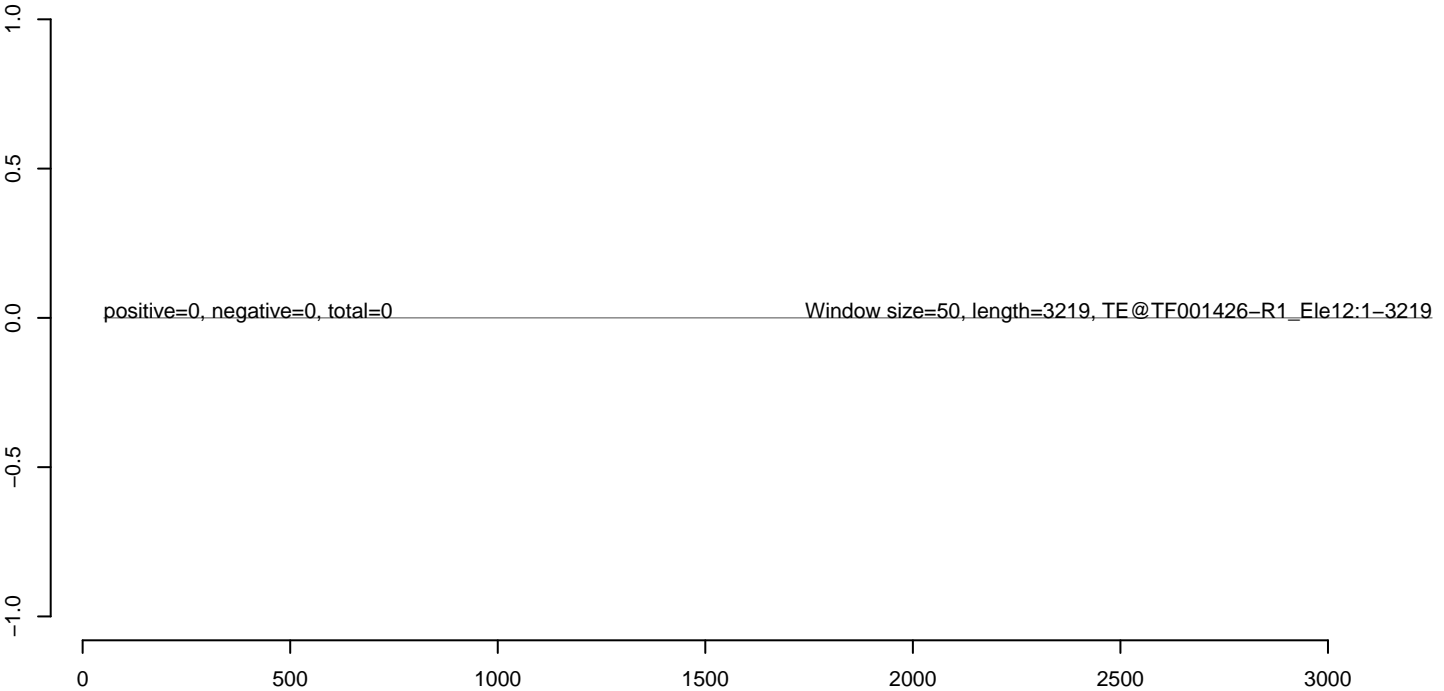
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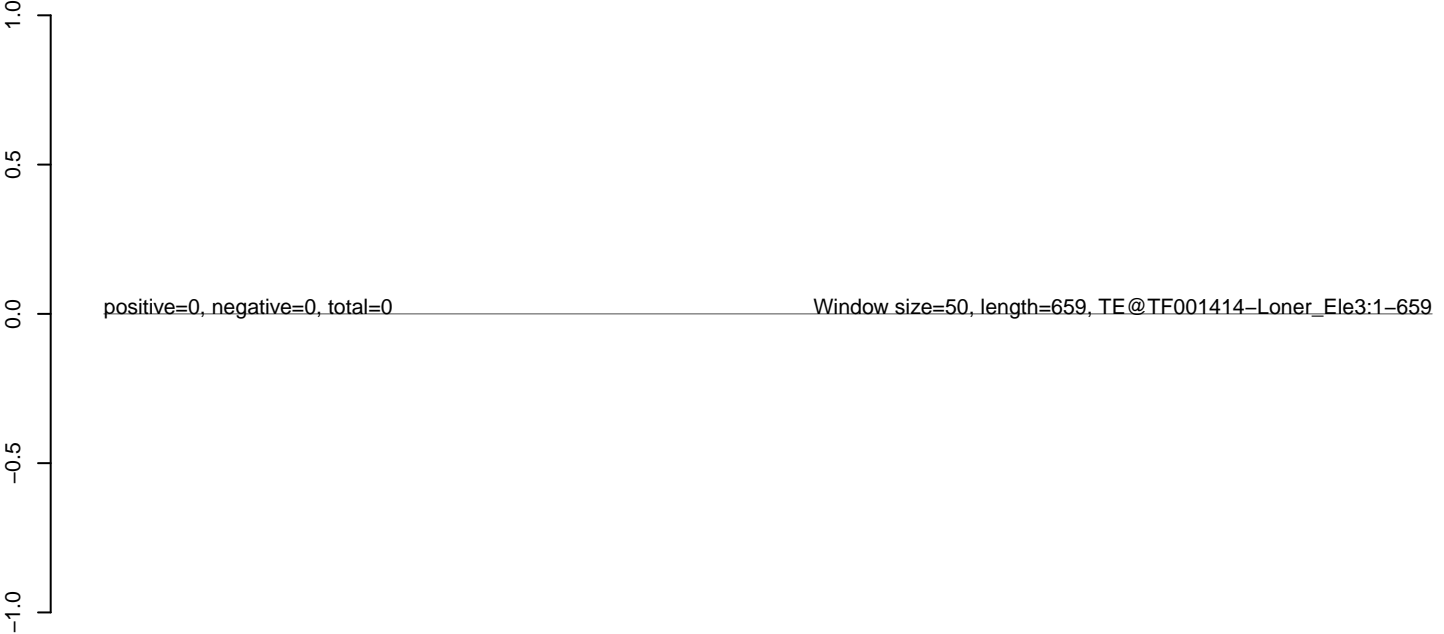
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AnGam_Sua5b_cells.rep



100 200 300 400 500 600 700

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AnGam_Sua5b_cells.rep



100 200 300 400 500 600 700

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AnGam_Sua5b_cells.rep



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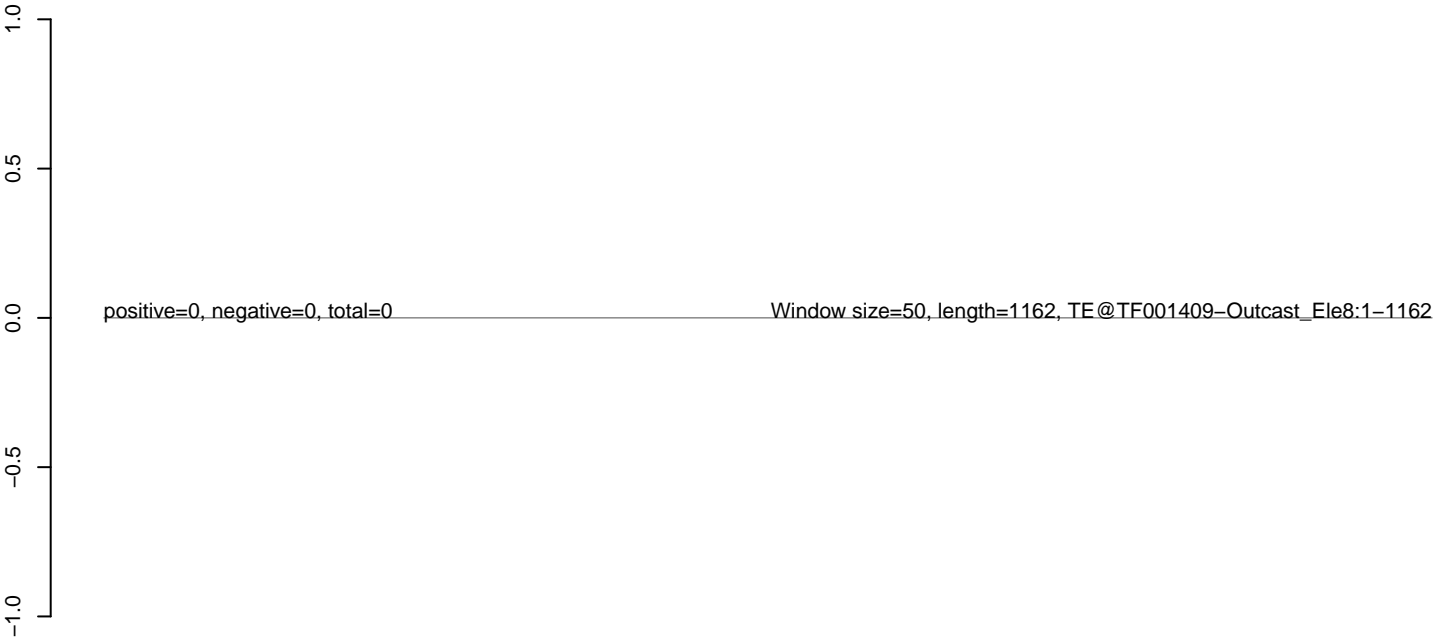
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AnGam_Sua5b_cells.rep



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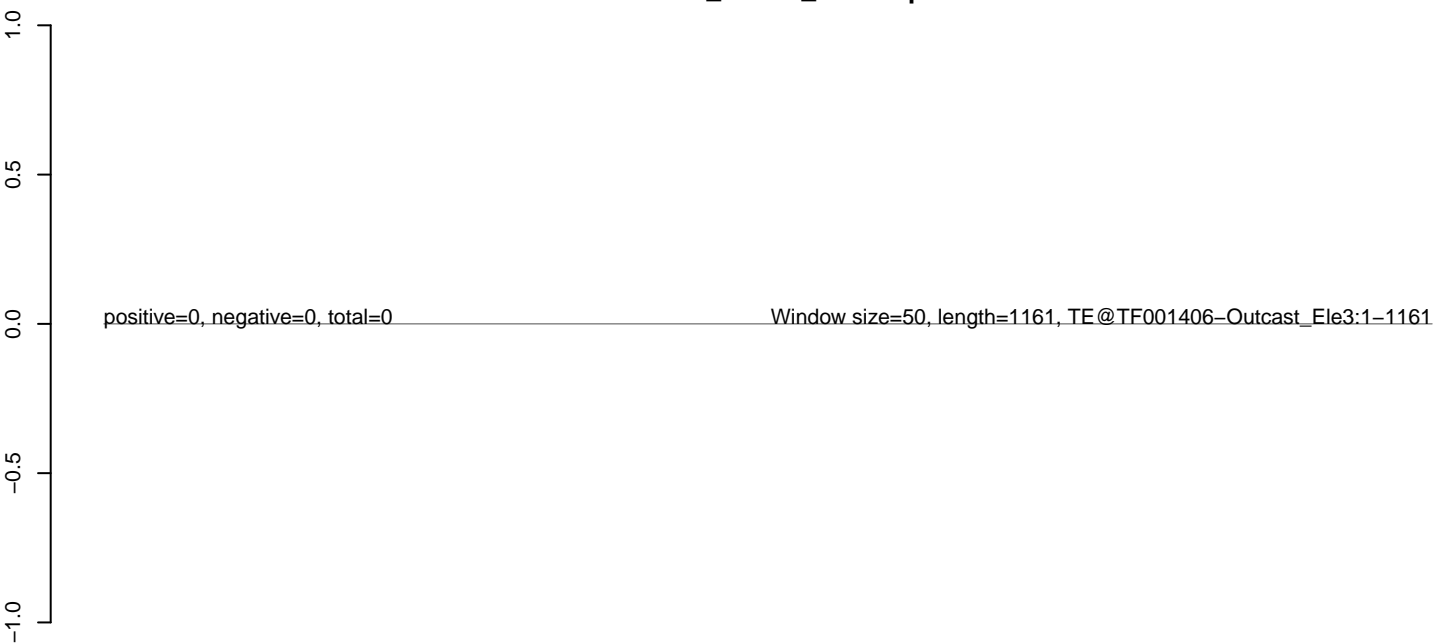
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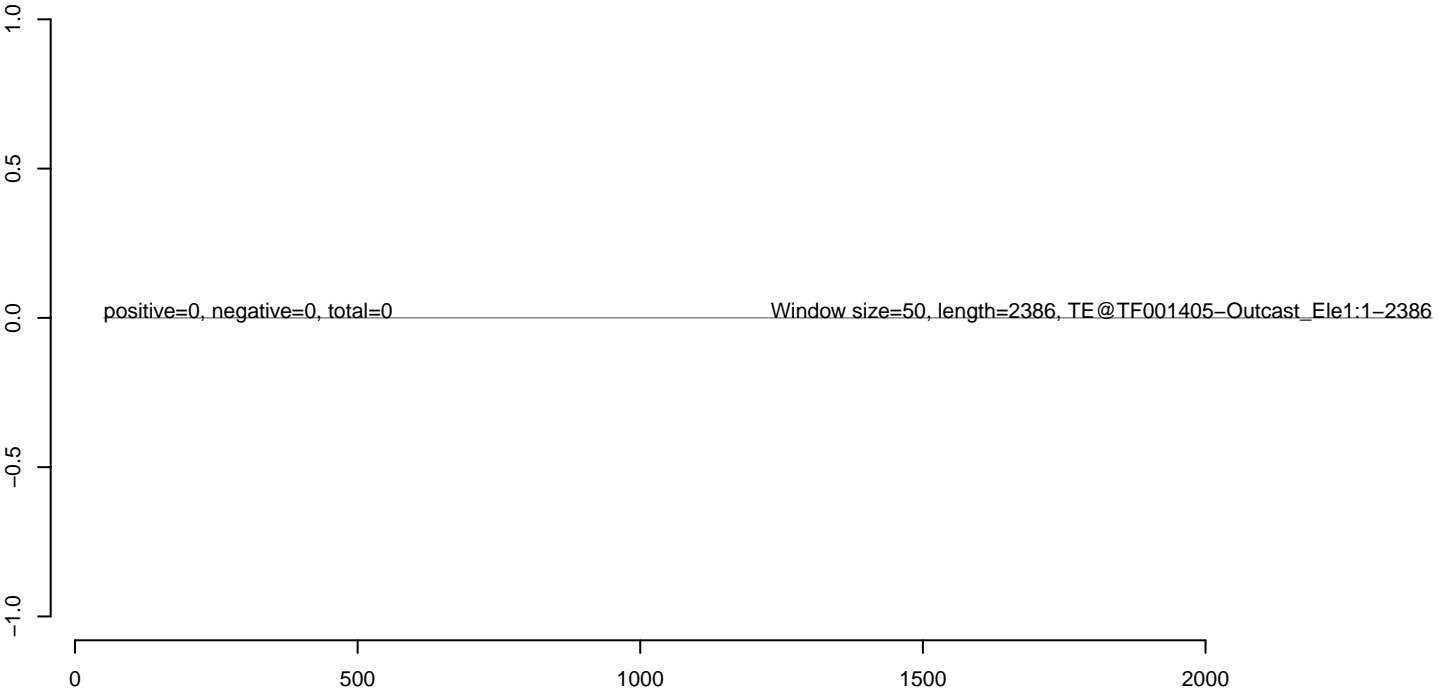
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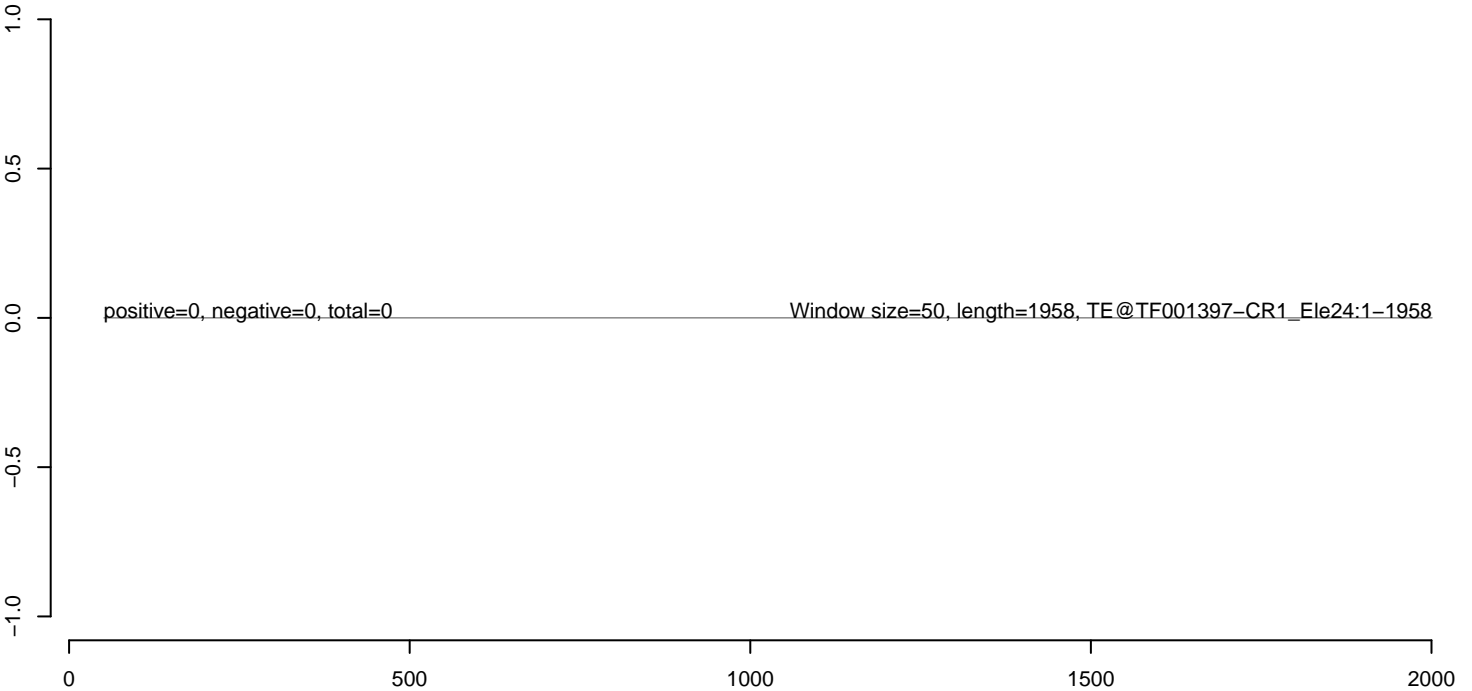
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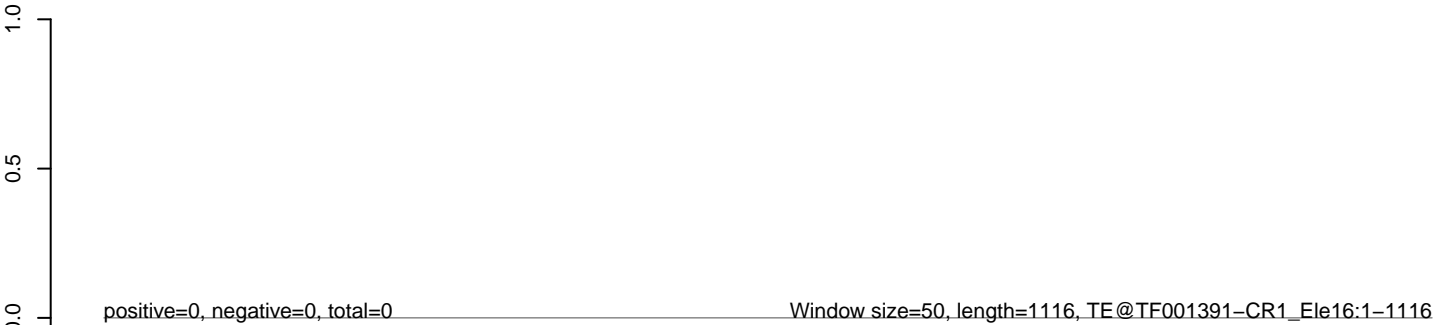
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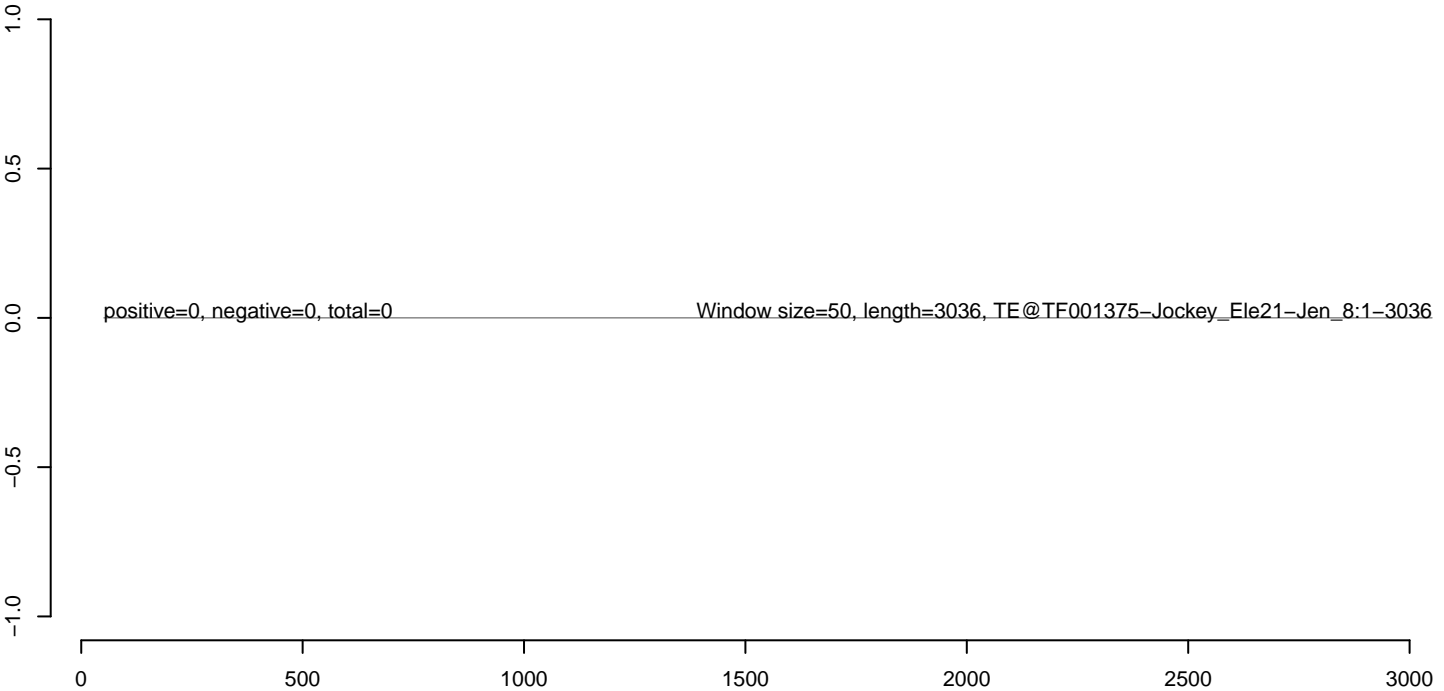
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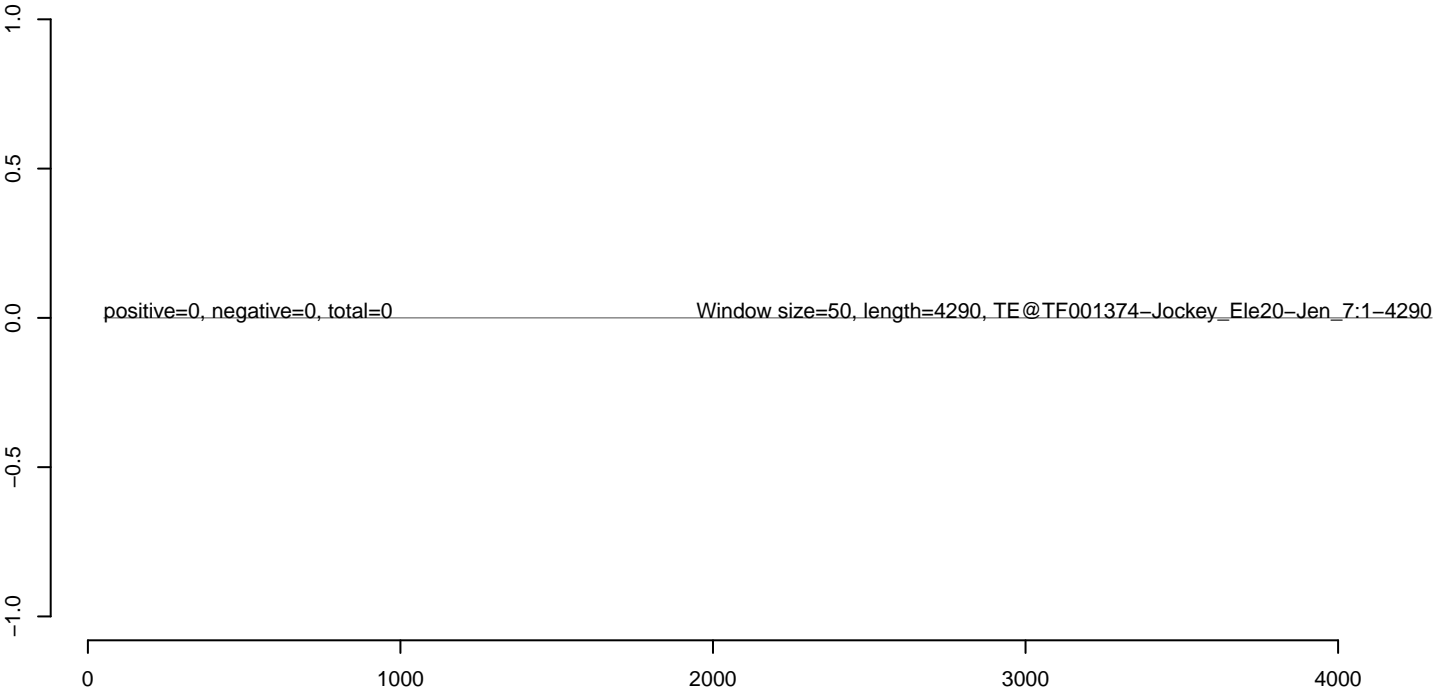
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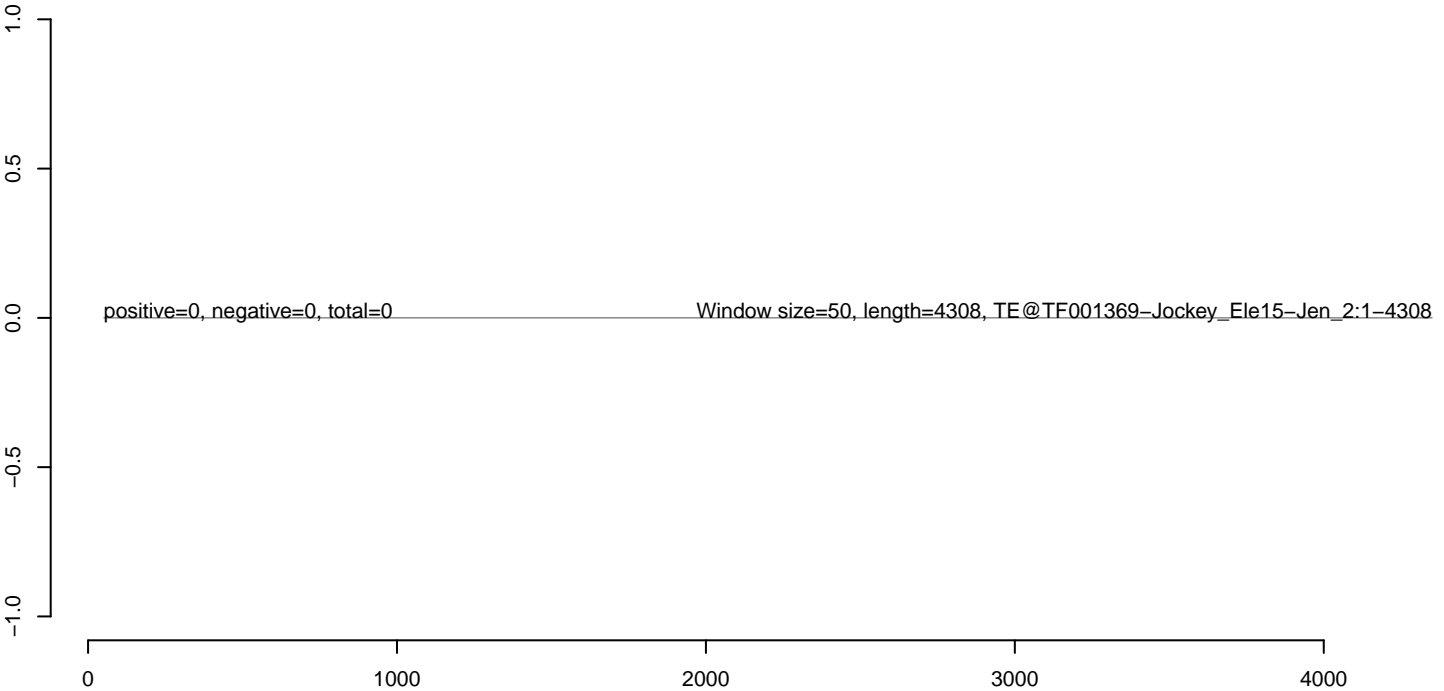
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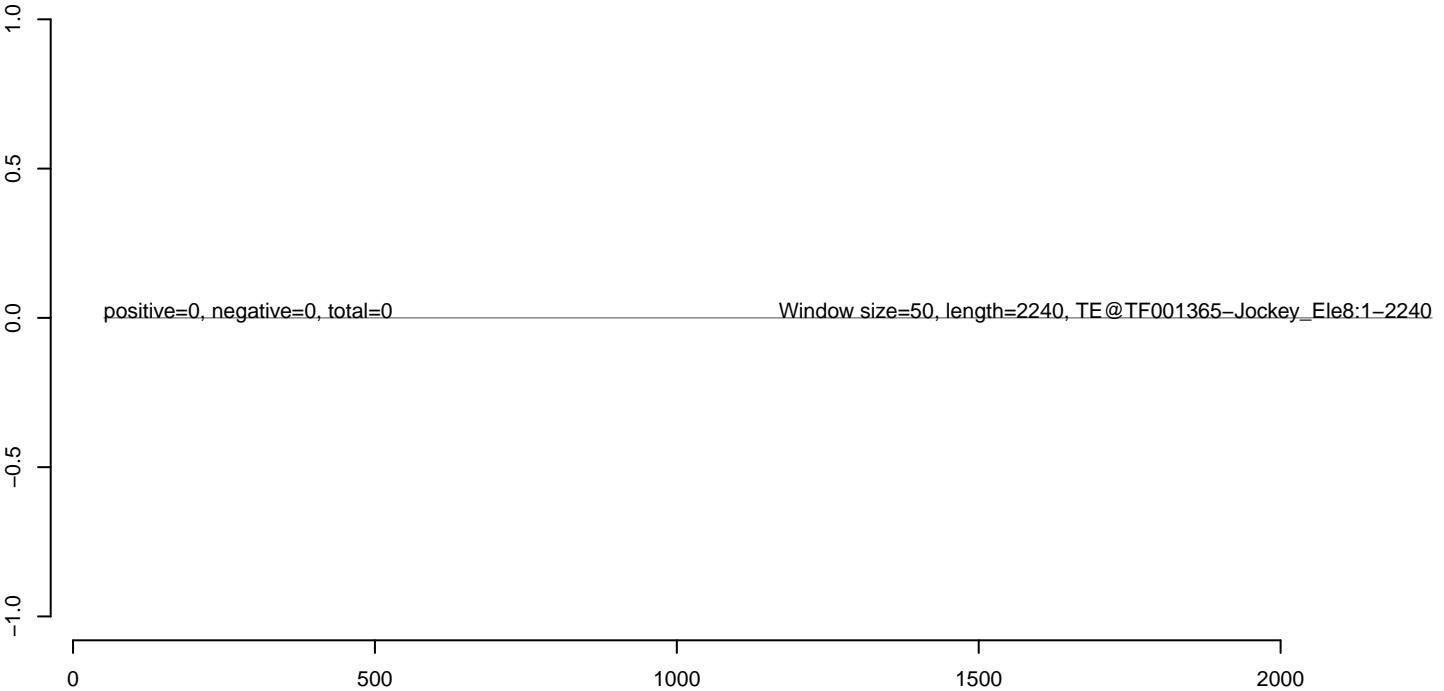
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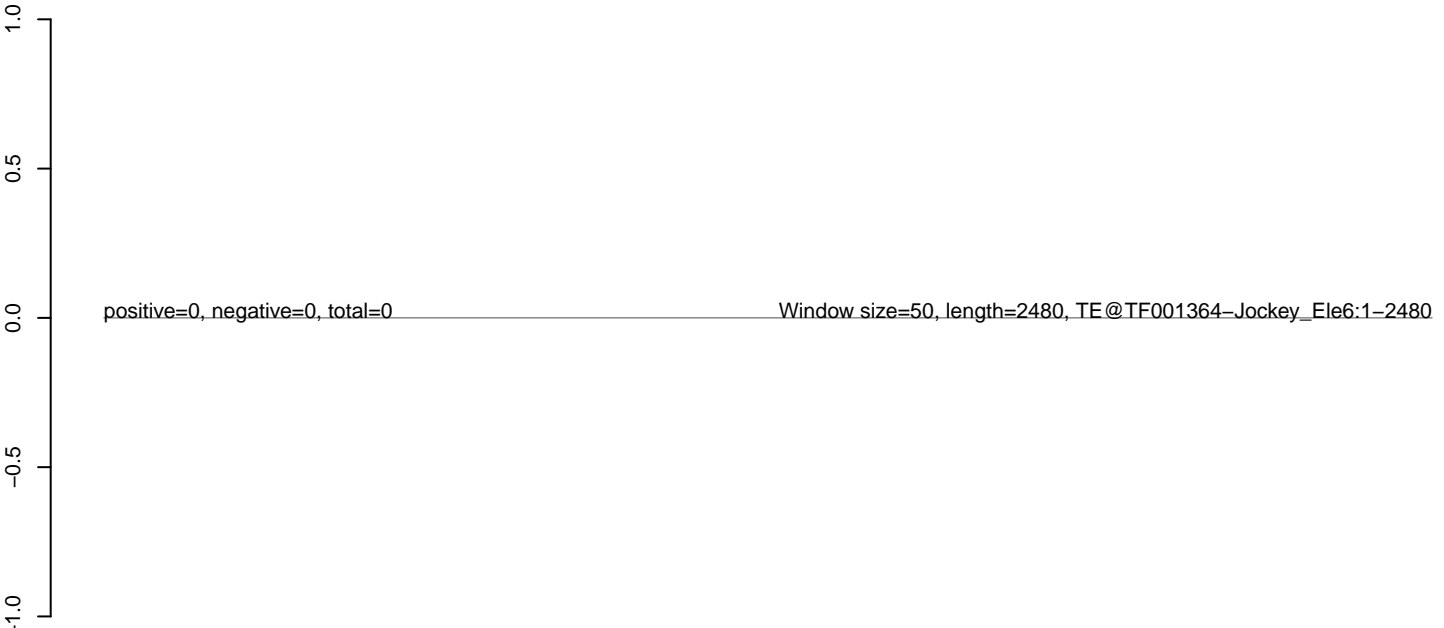
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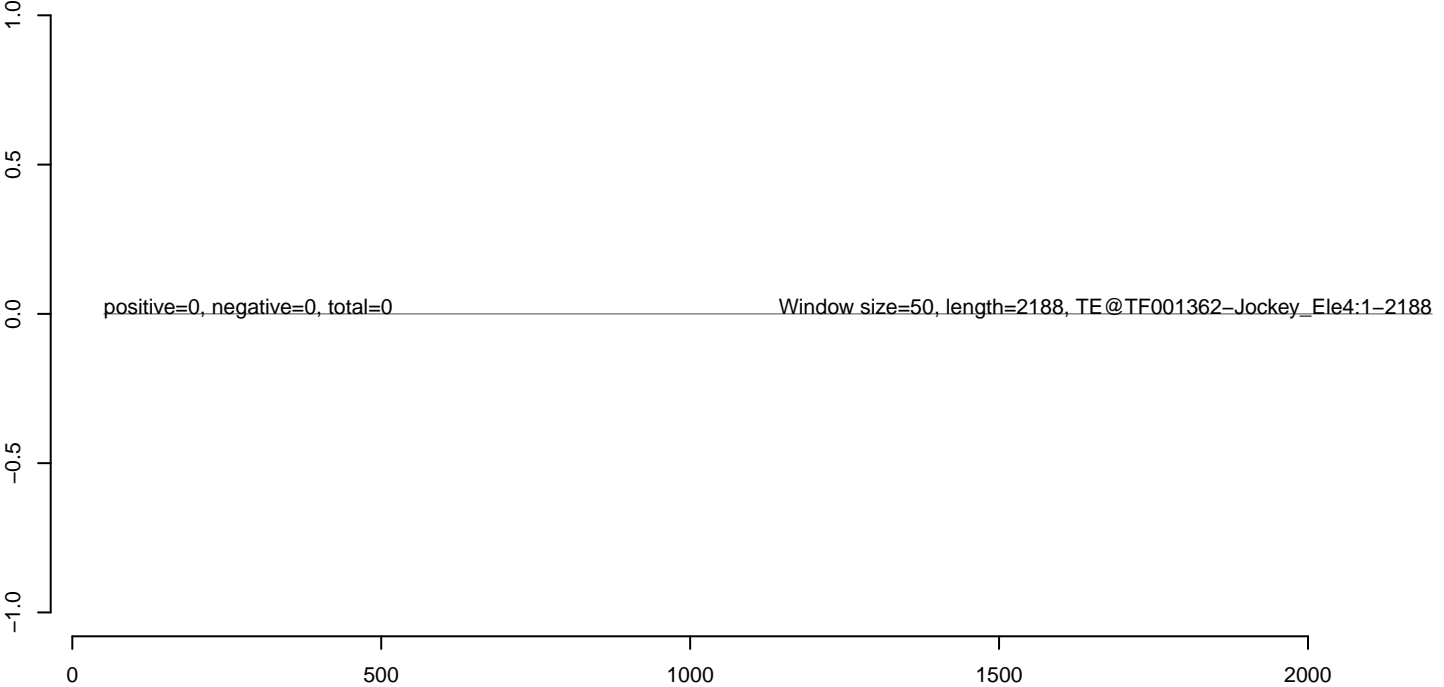
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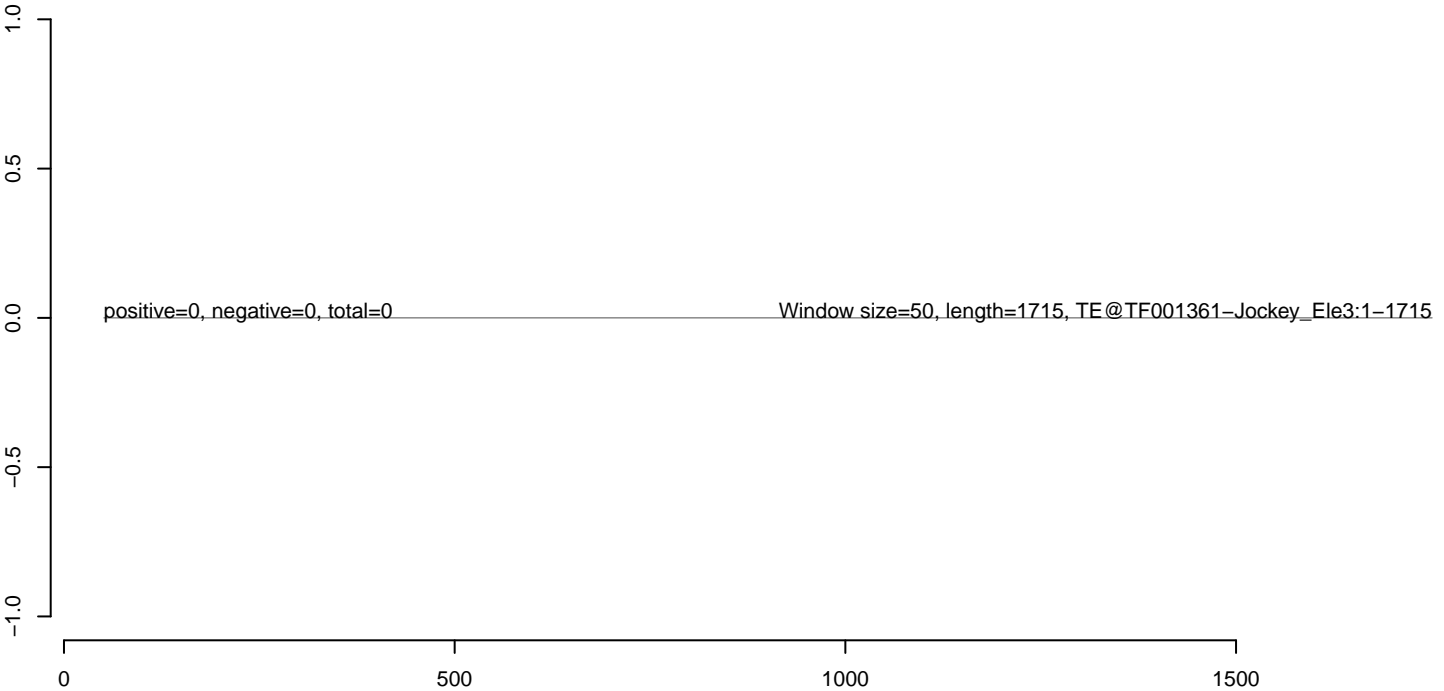
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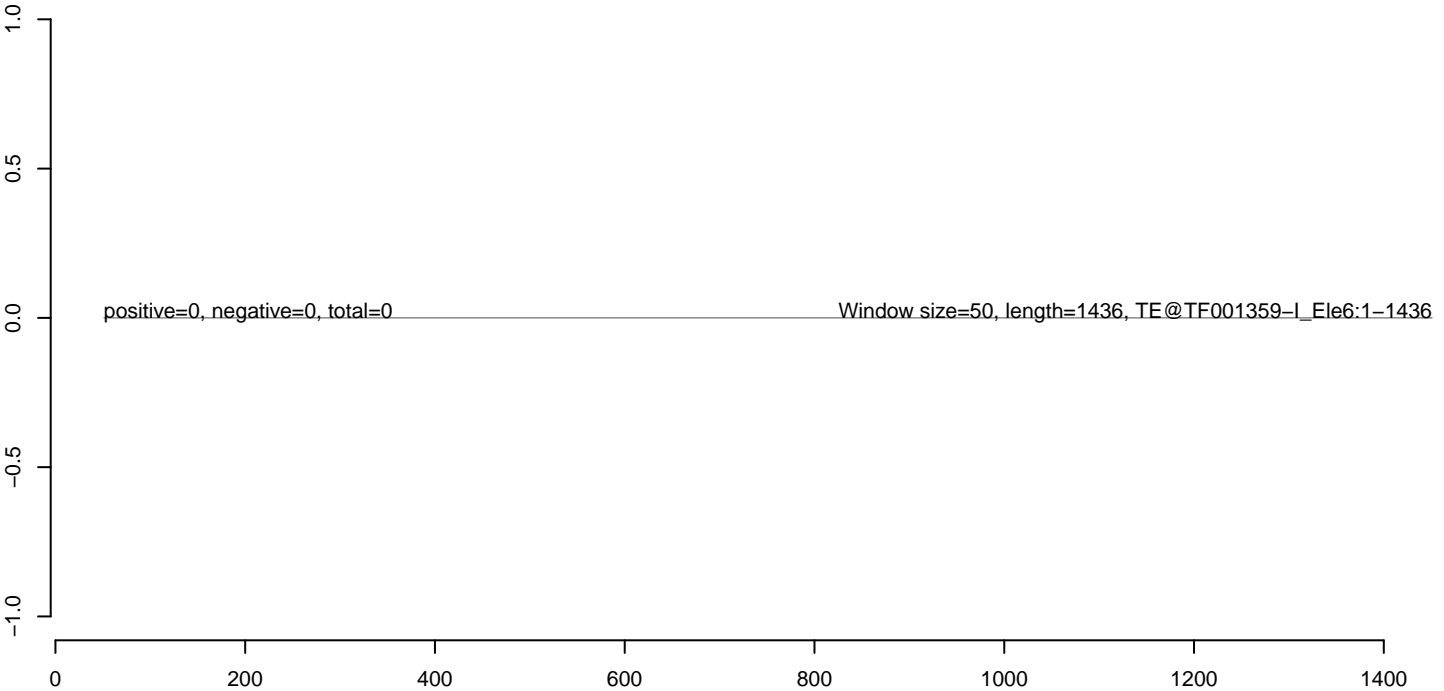
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AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



200 400 600 800

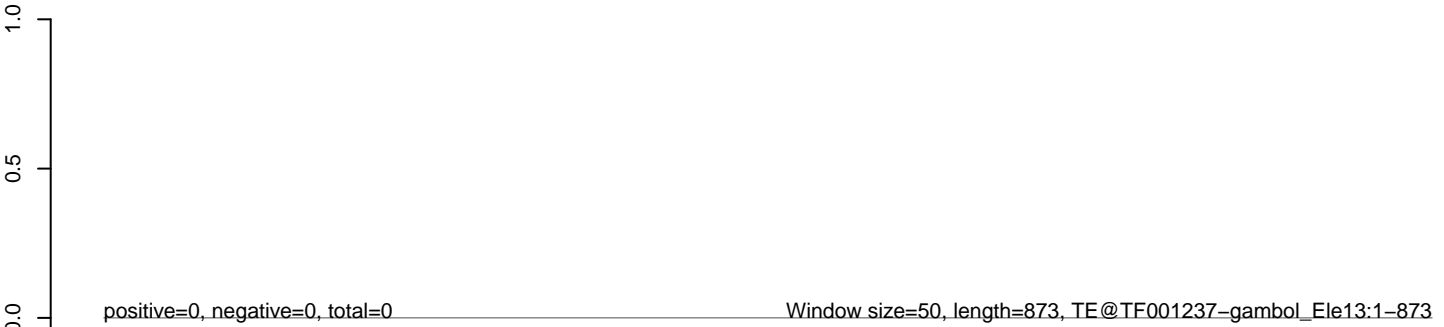
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AnGam_Sua5b_cells.rep



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AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



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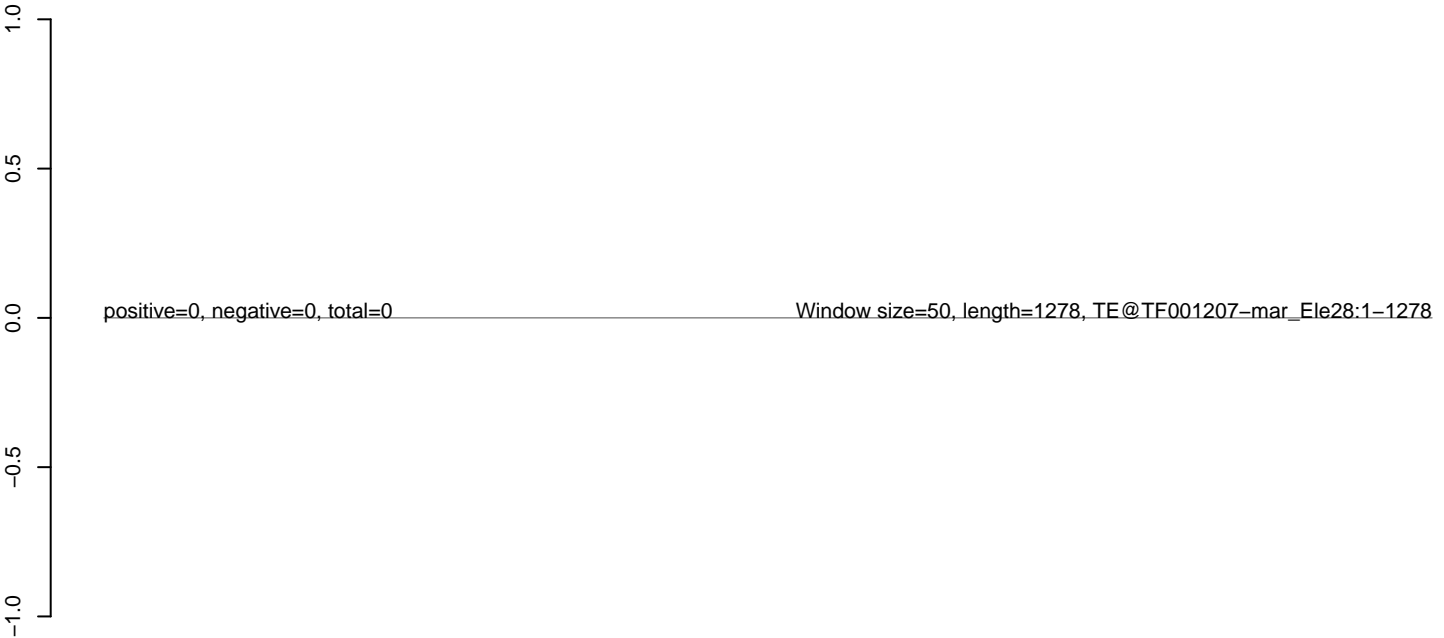
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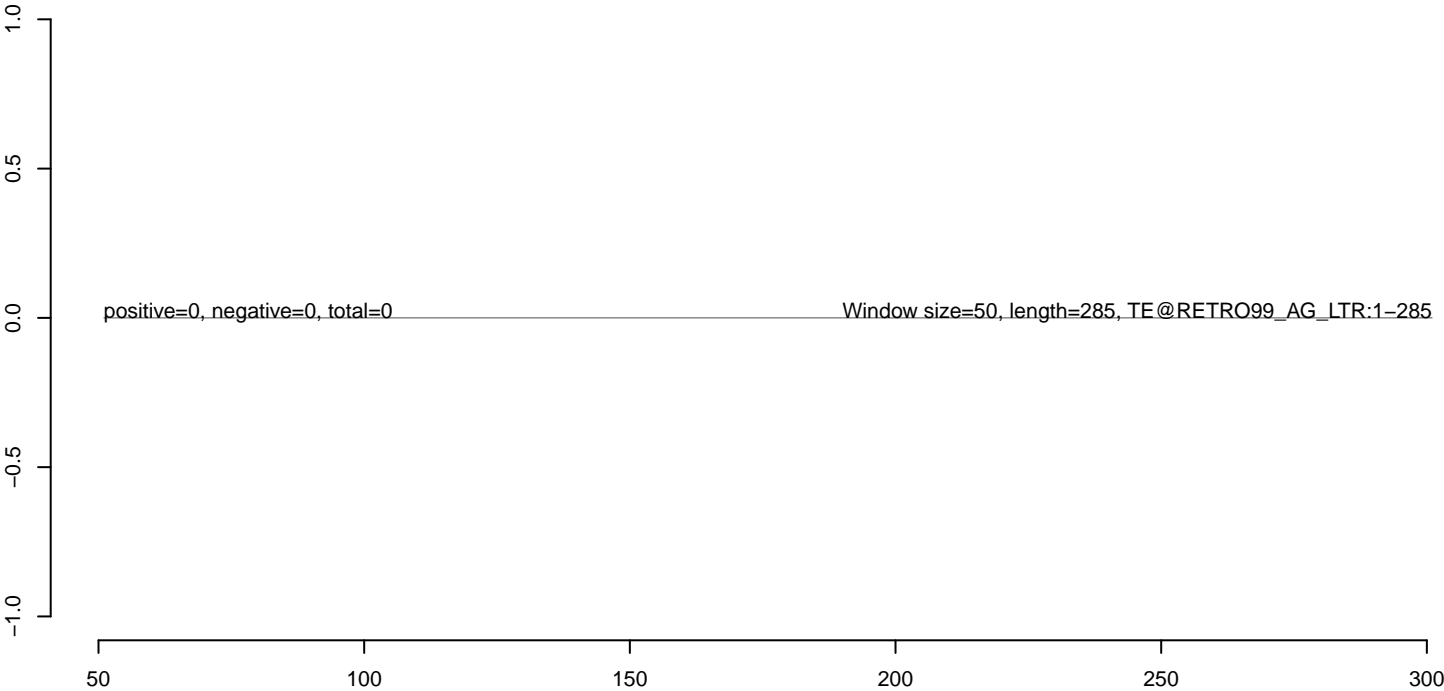
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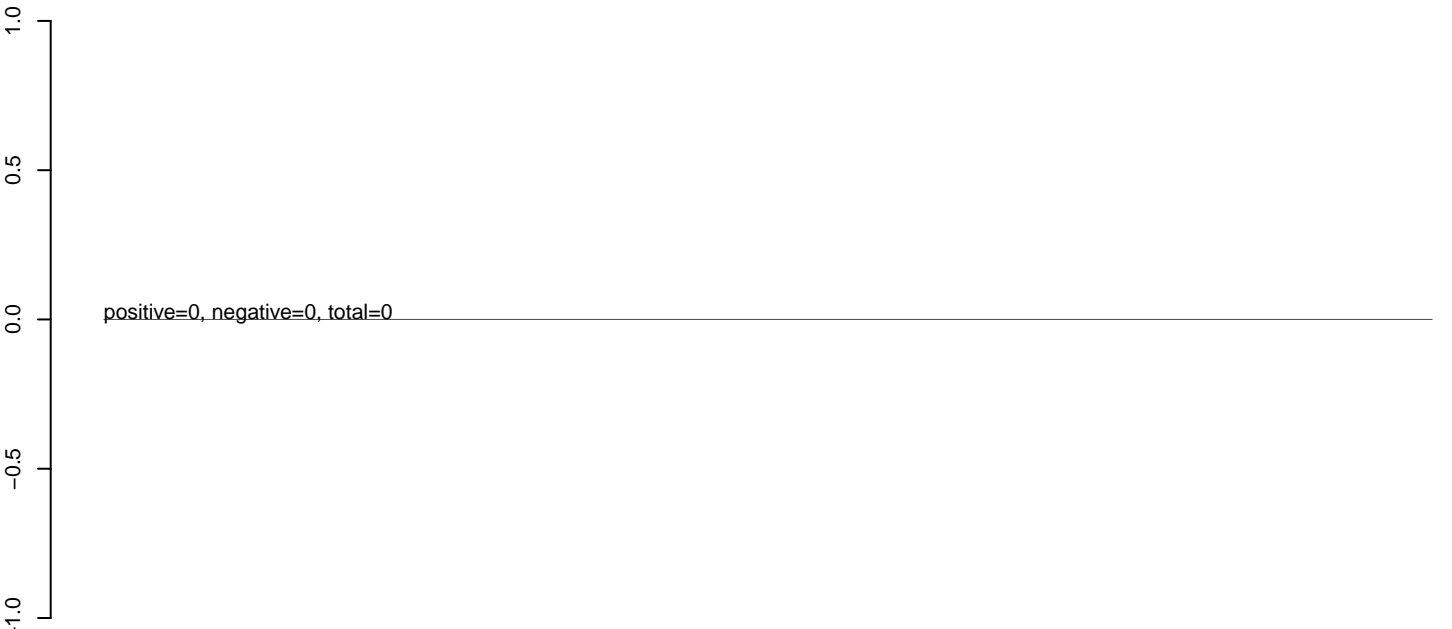
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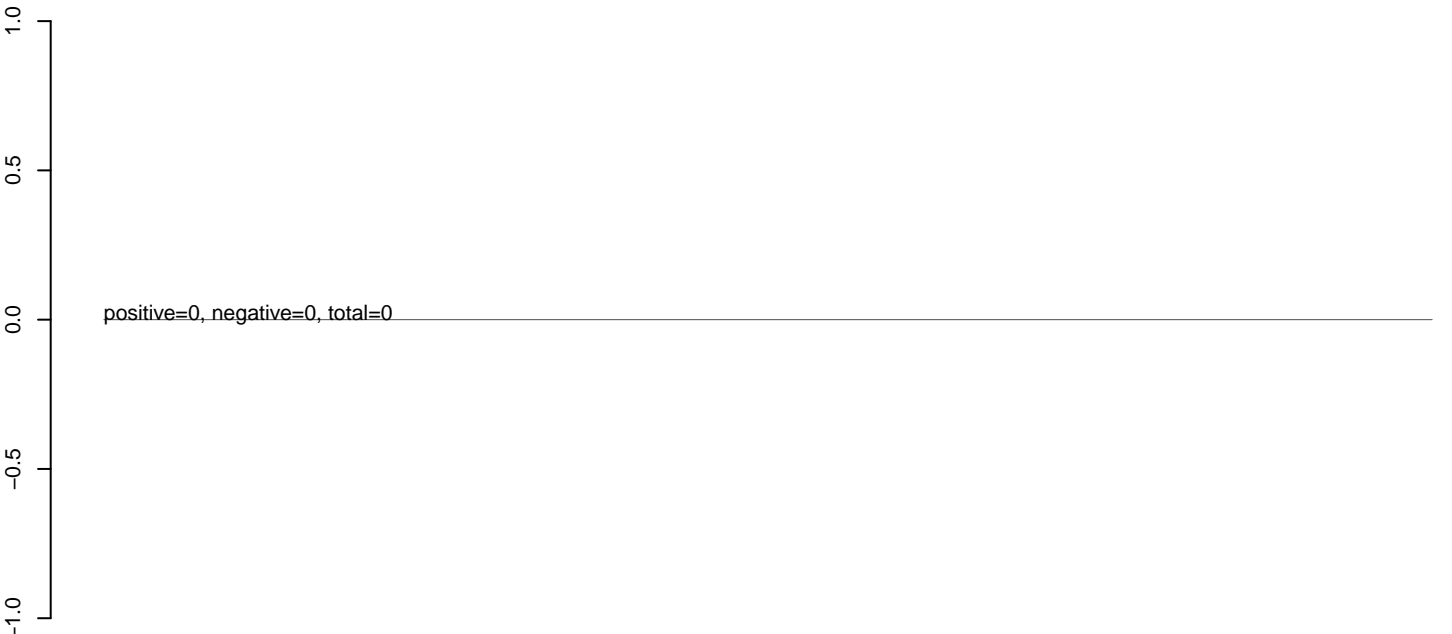
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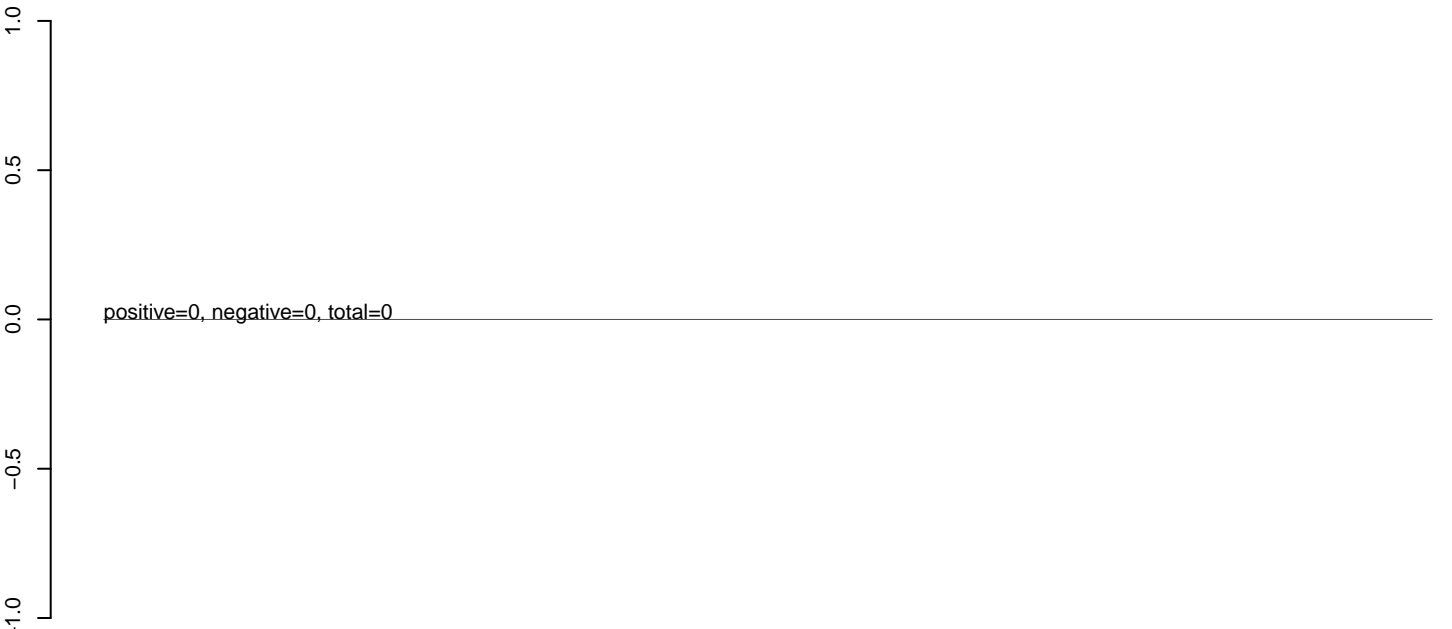
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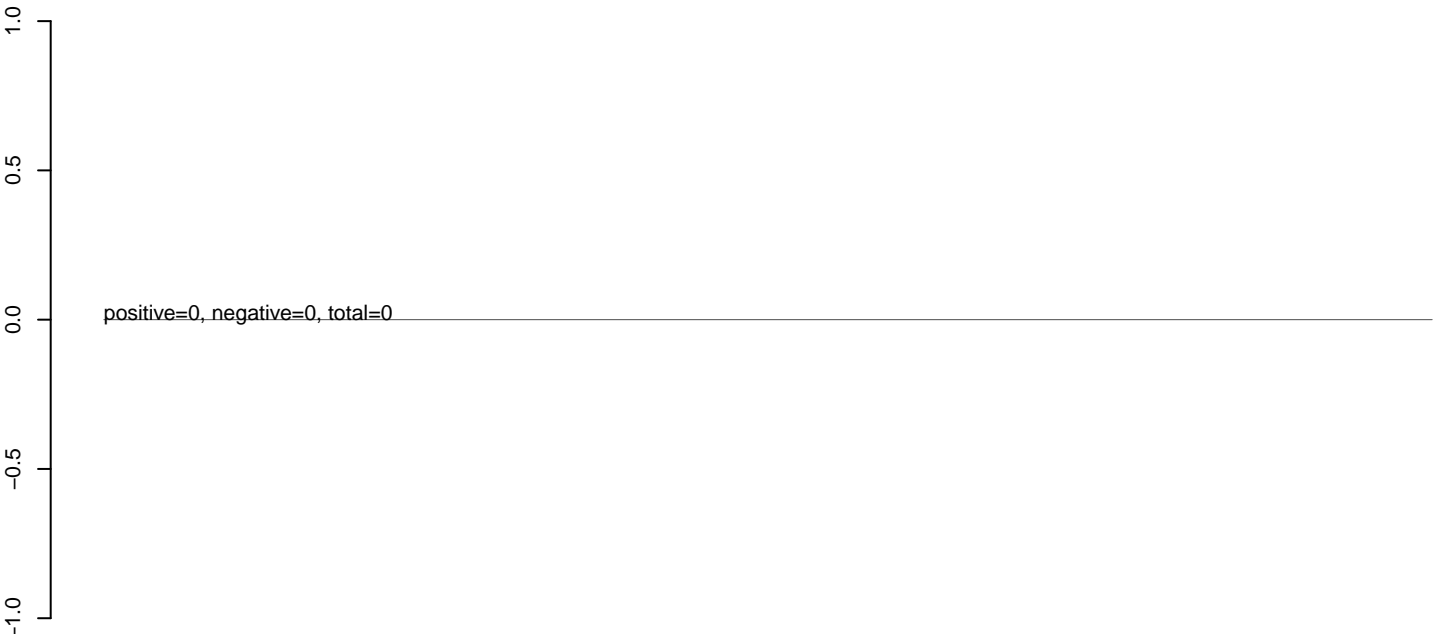
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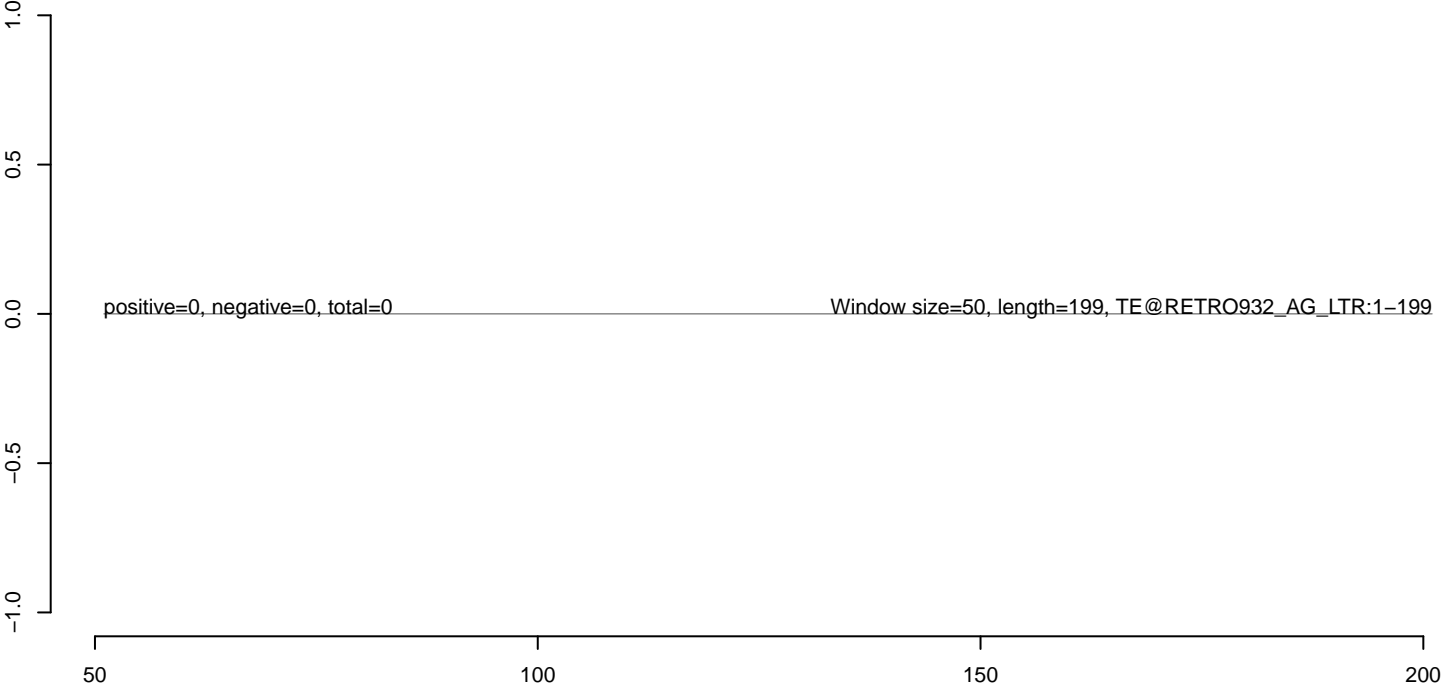
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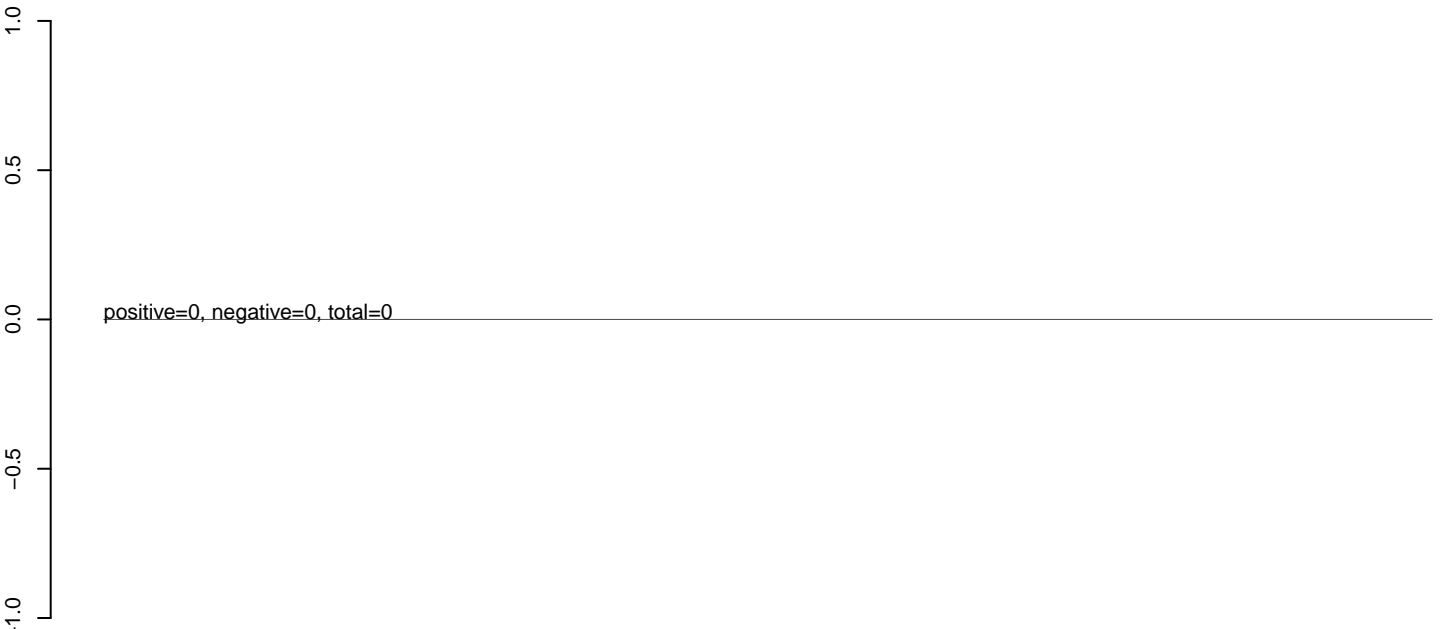
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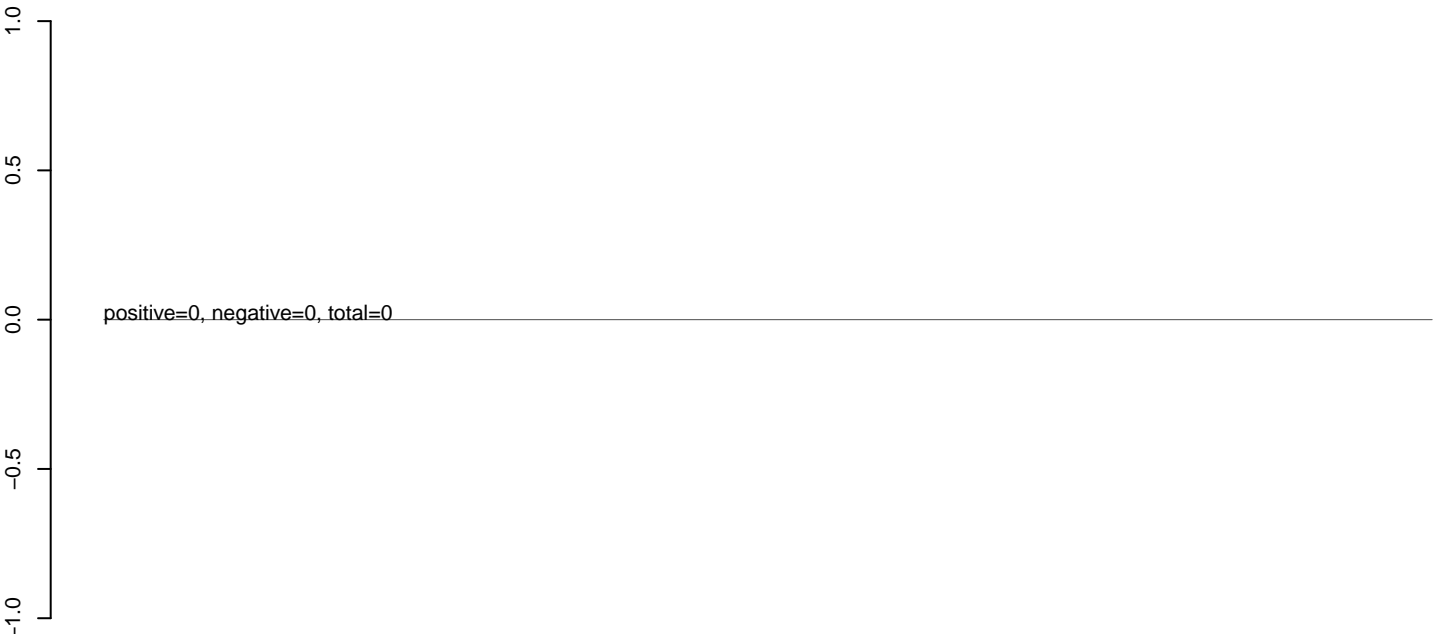
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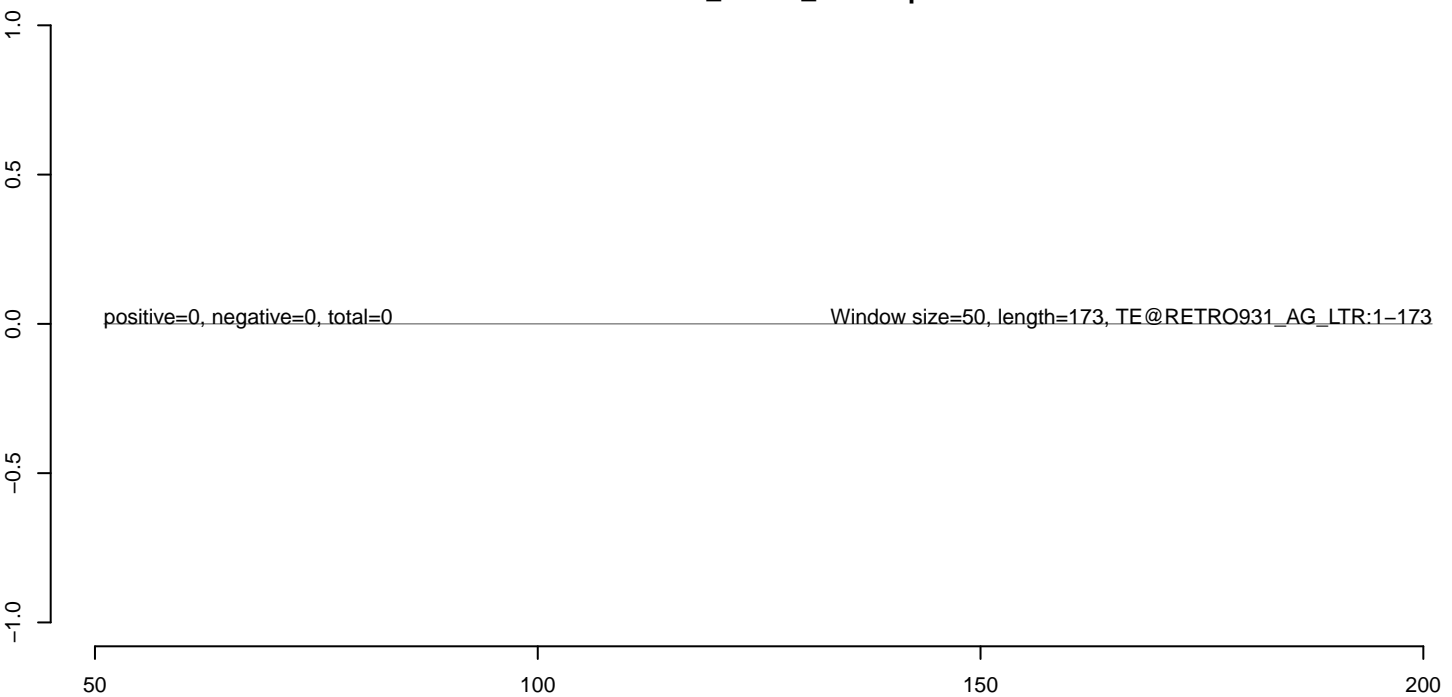
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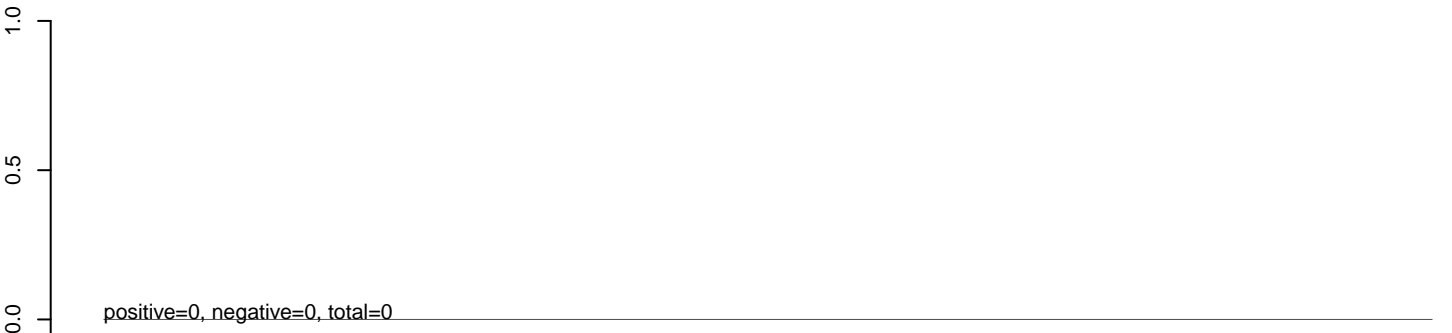
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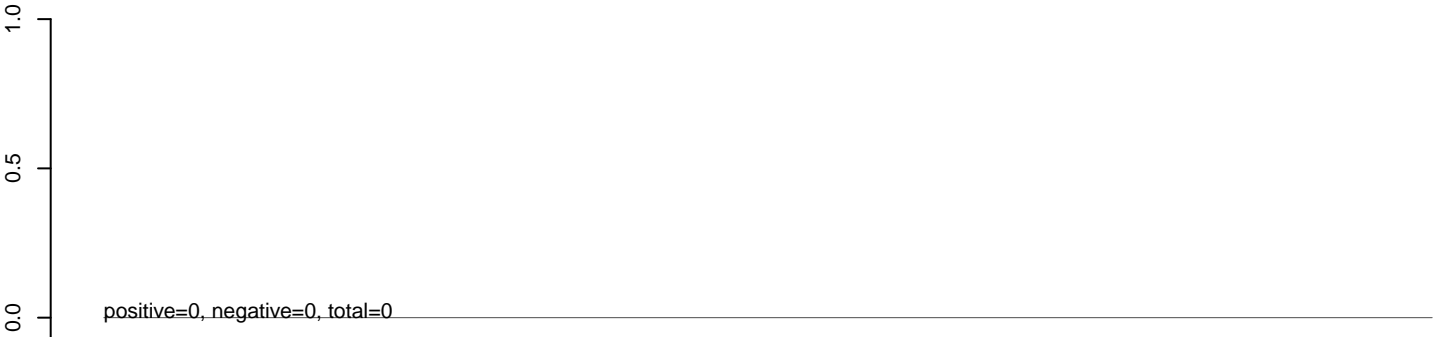
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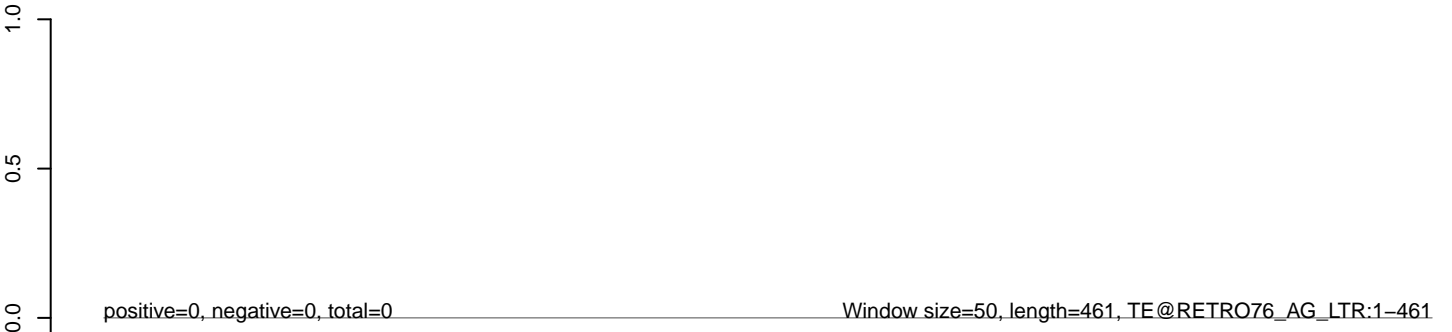
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100 200 300 400 500

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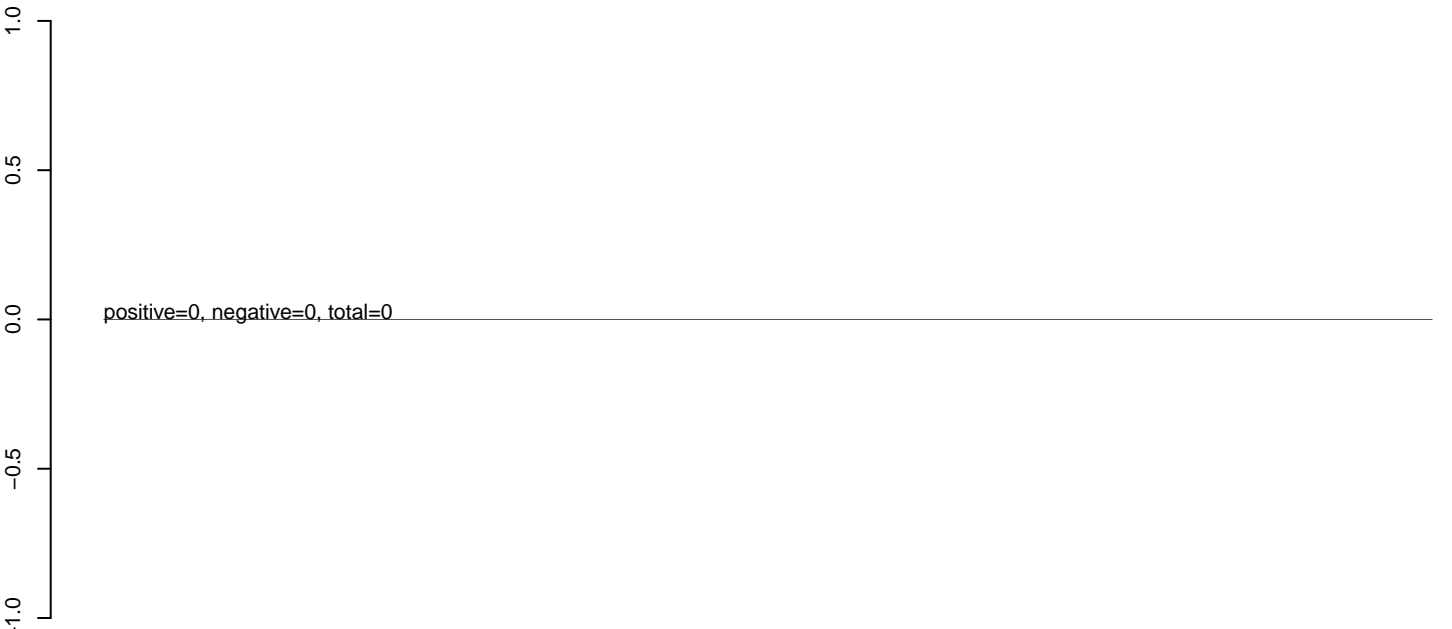
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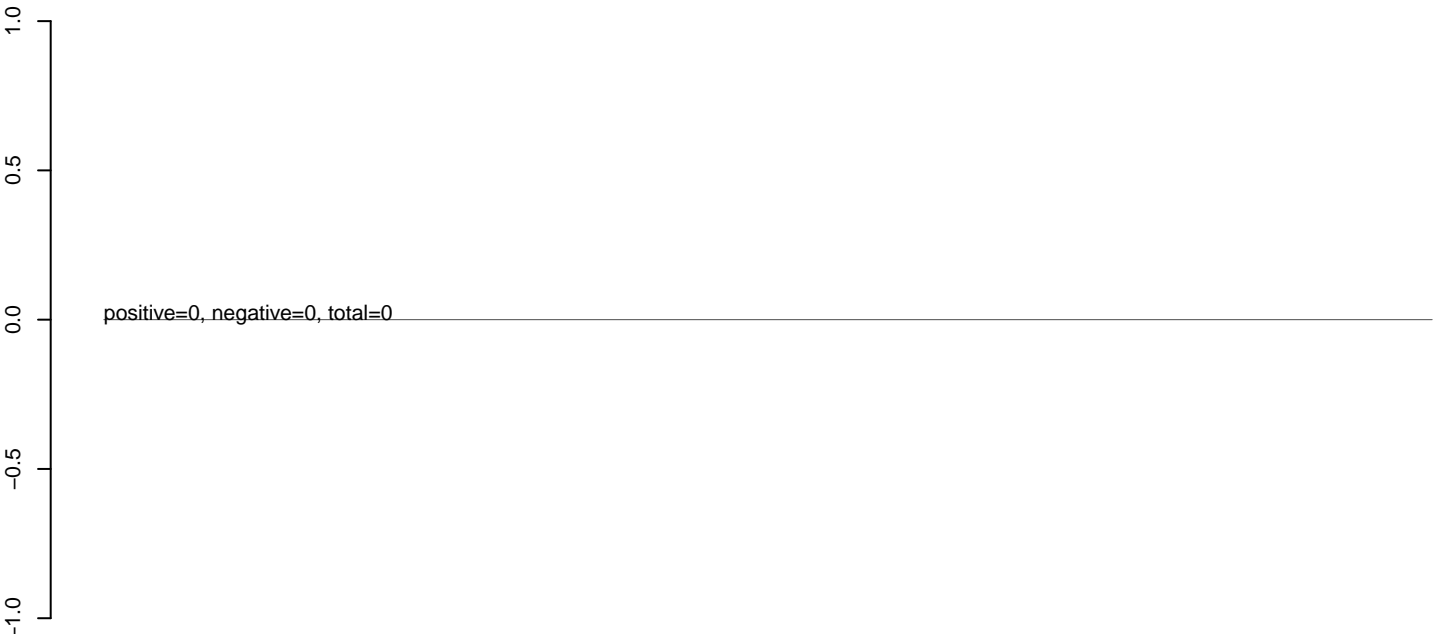
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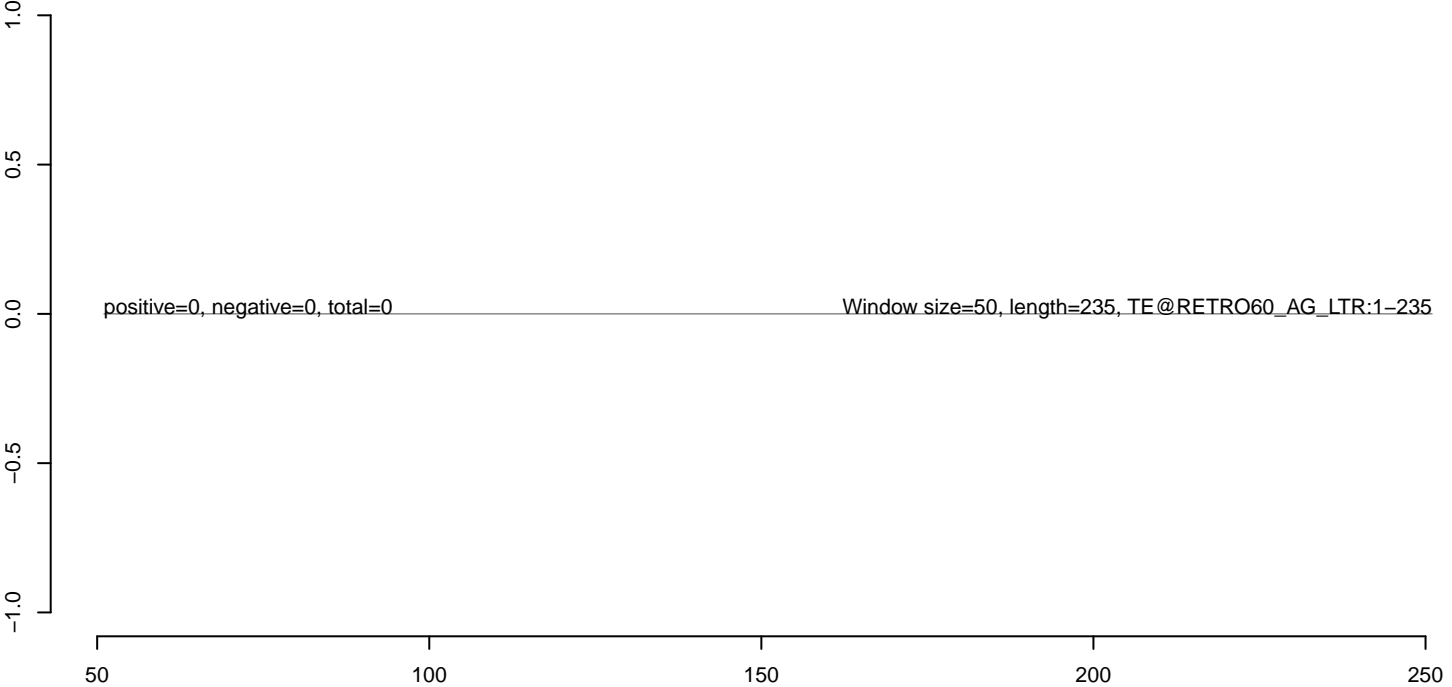
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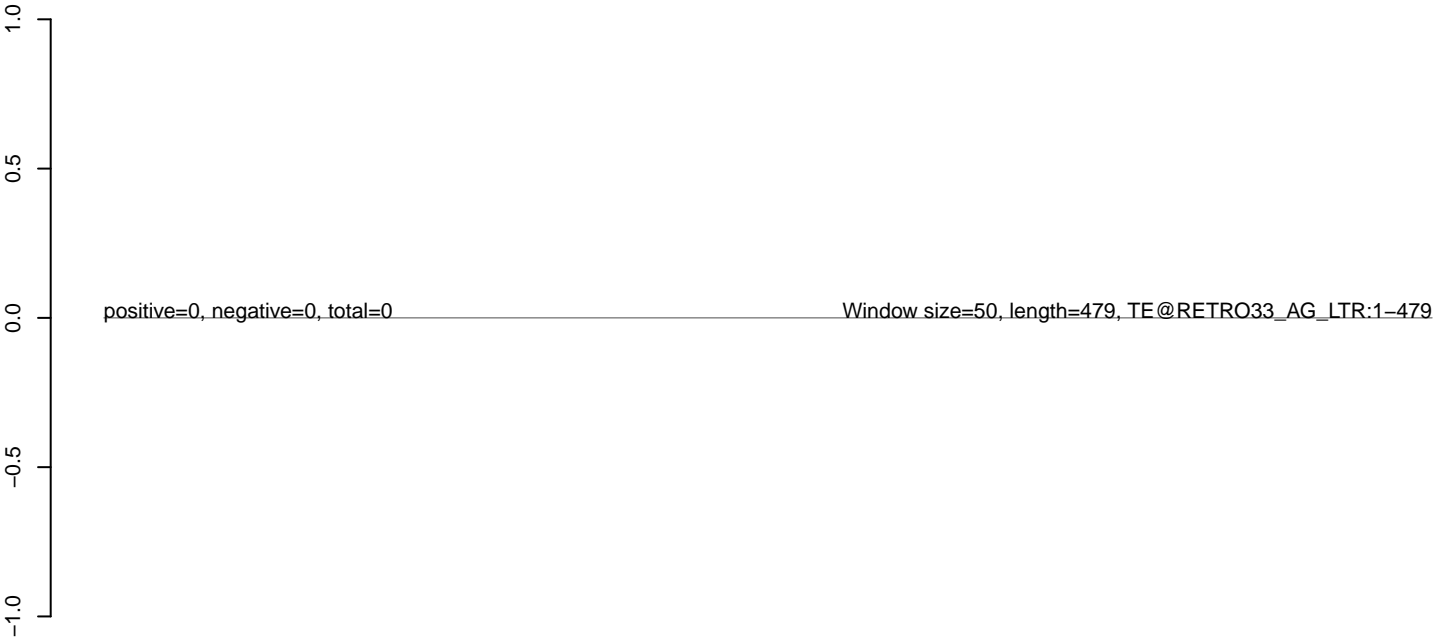
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AnGam_Sua5b_cells.rep



100 200 300 400 500

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AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



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AnGam_Sua5b_cells.rep



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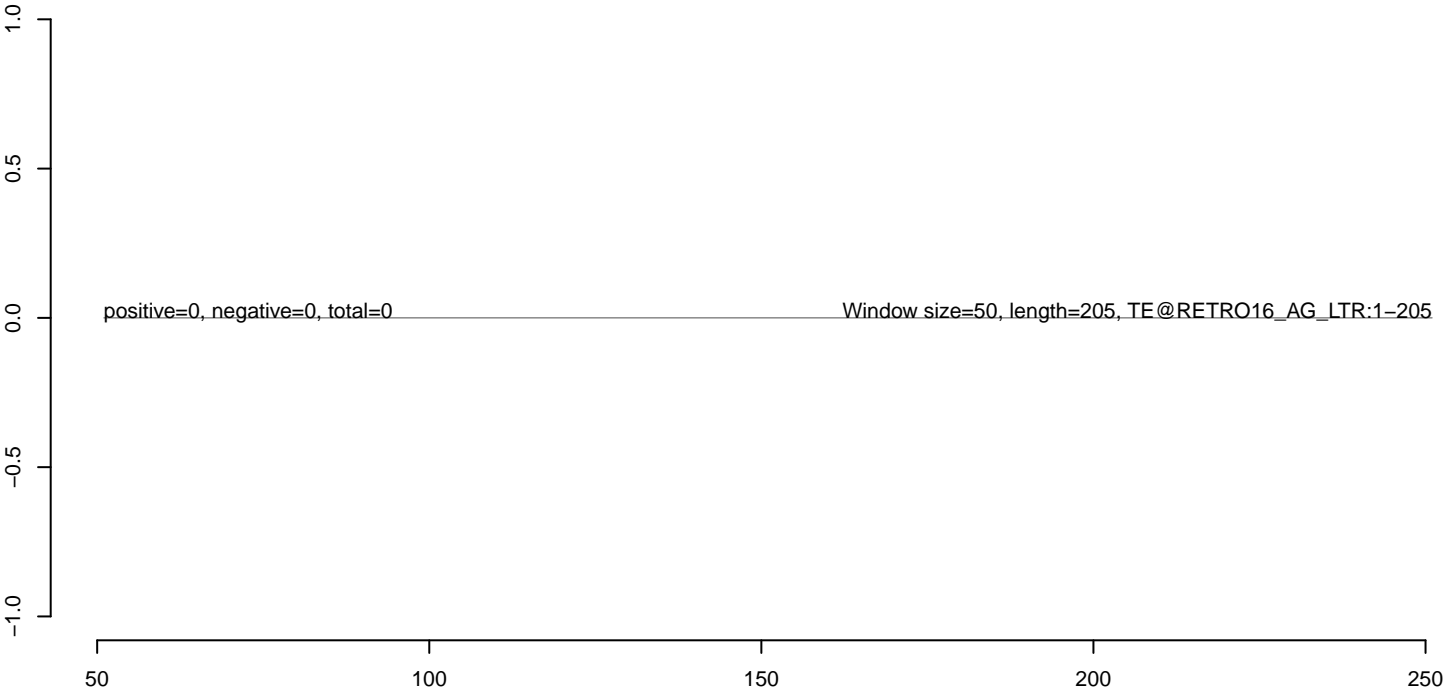
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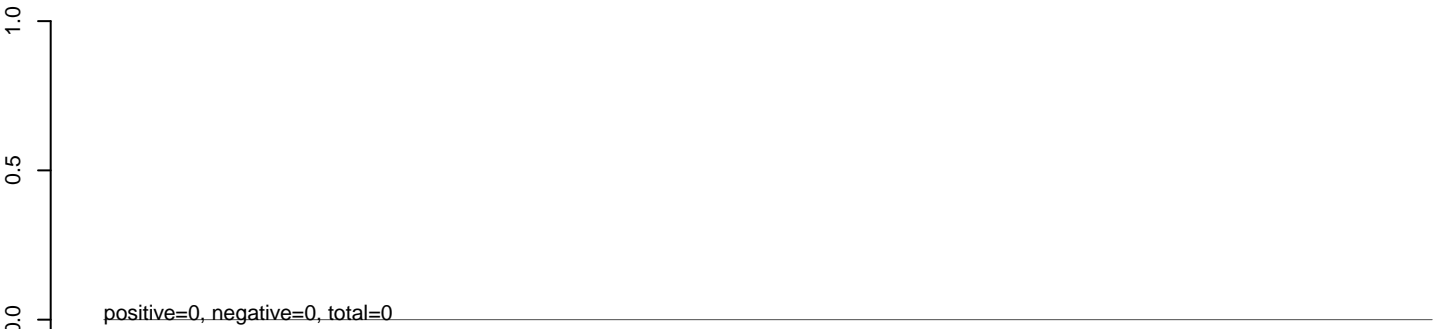
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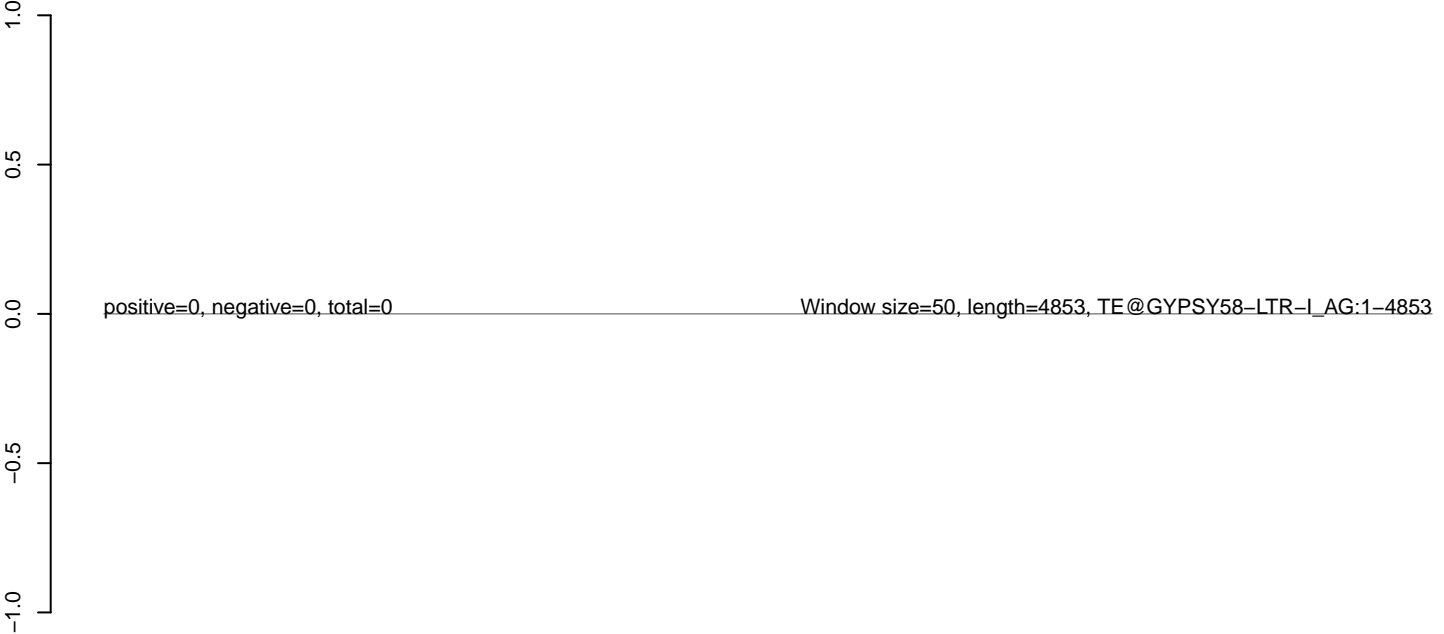
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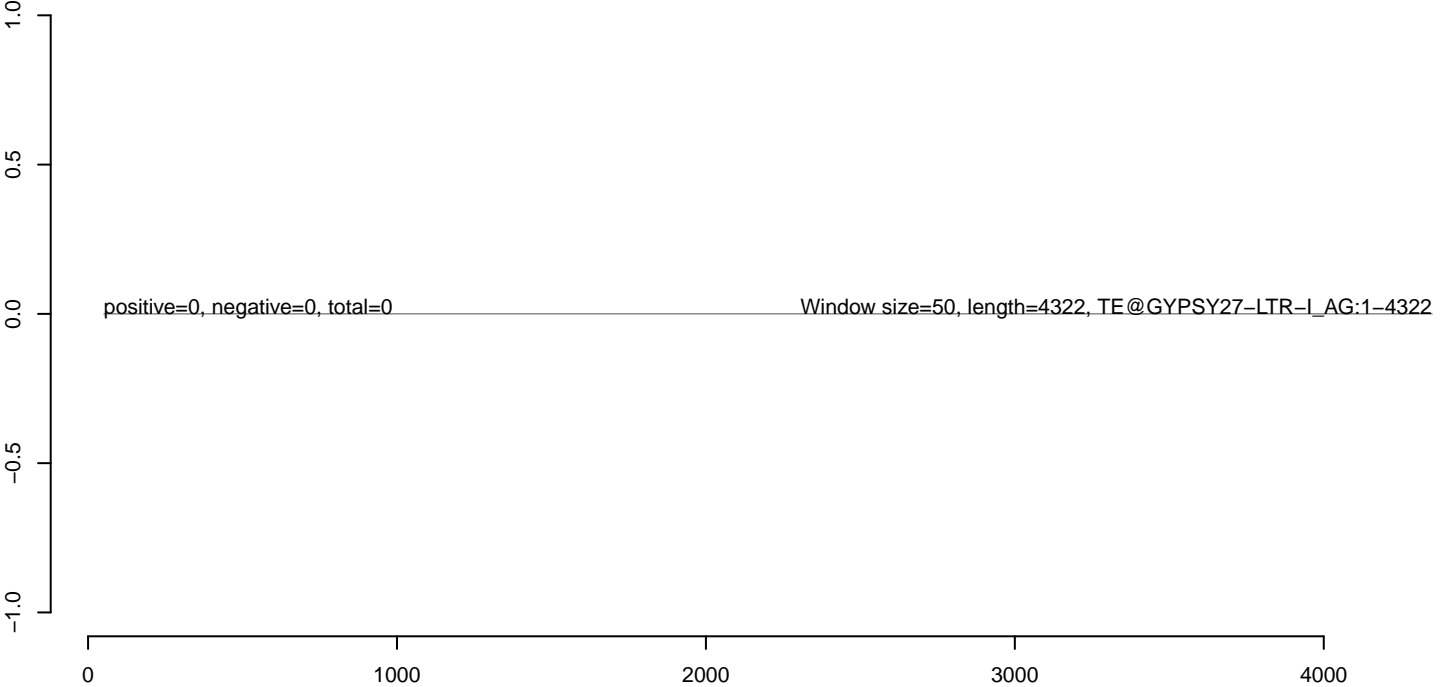
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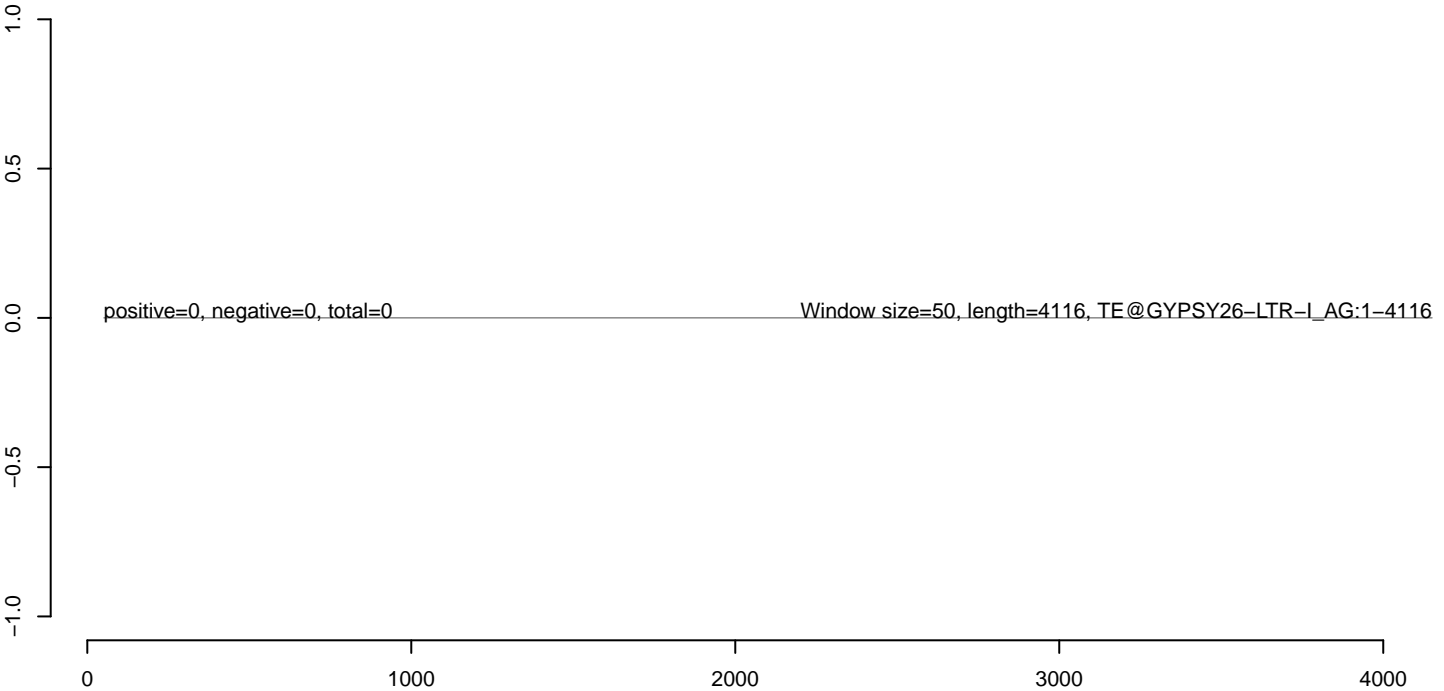
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AnGam_Sua5b_cells.rep



200 400 600 800 1000

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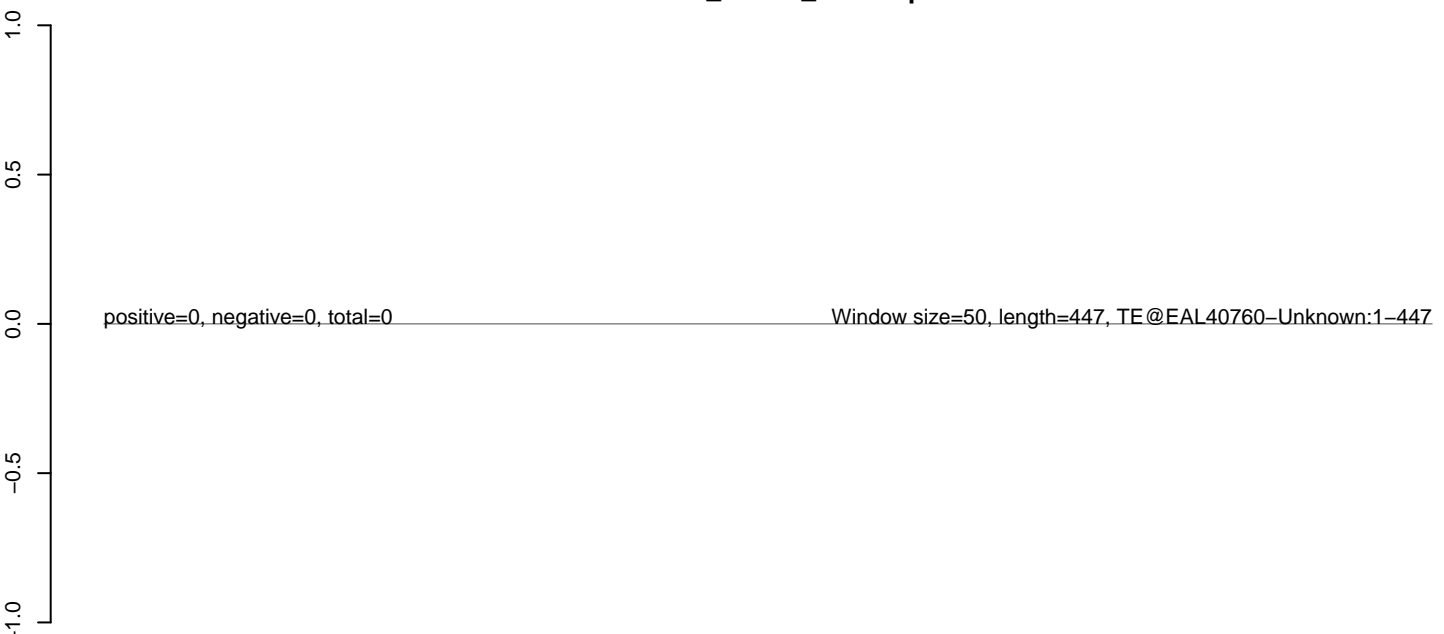
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100 200 300 400

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AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



200 400 600 800 1000

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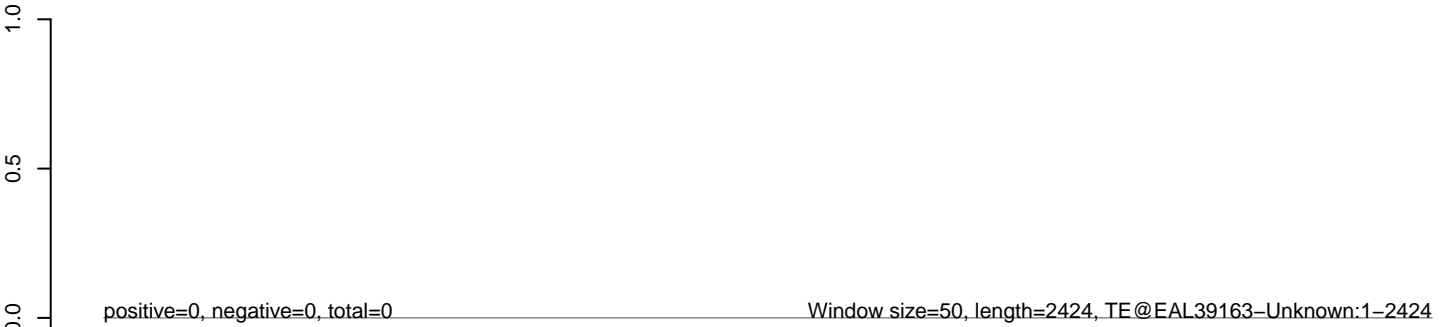
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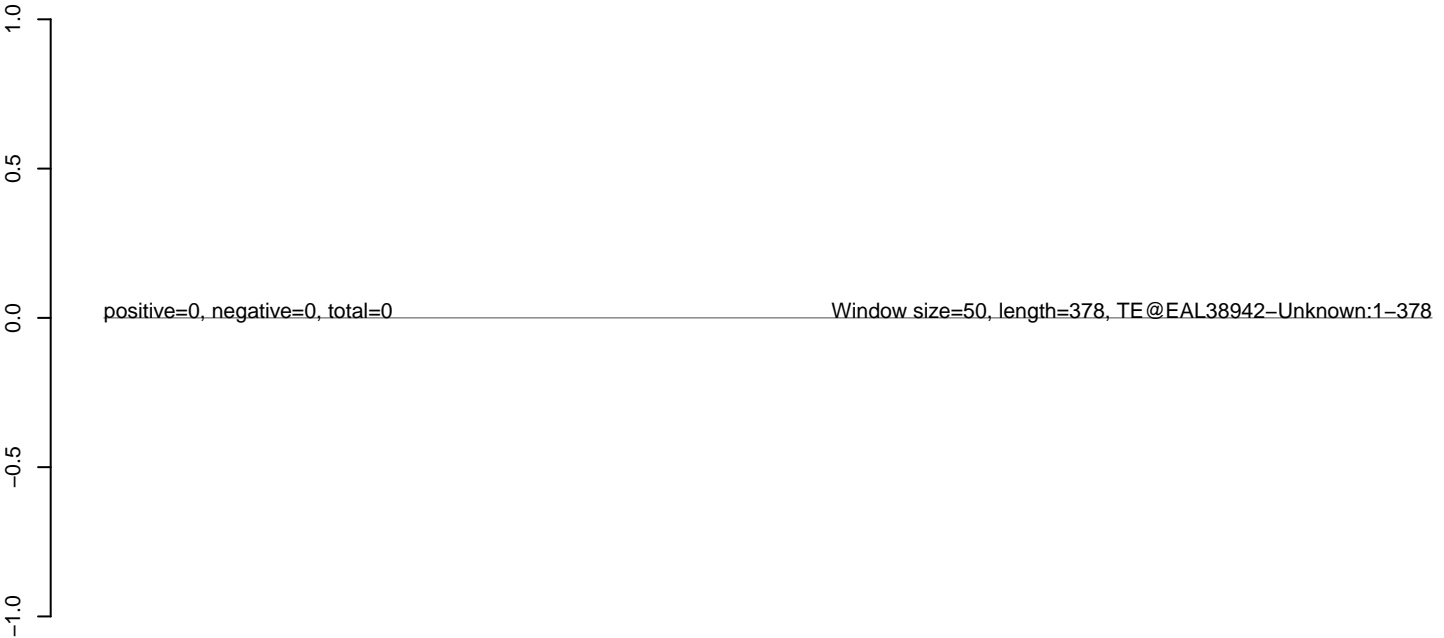
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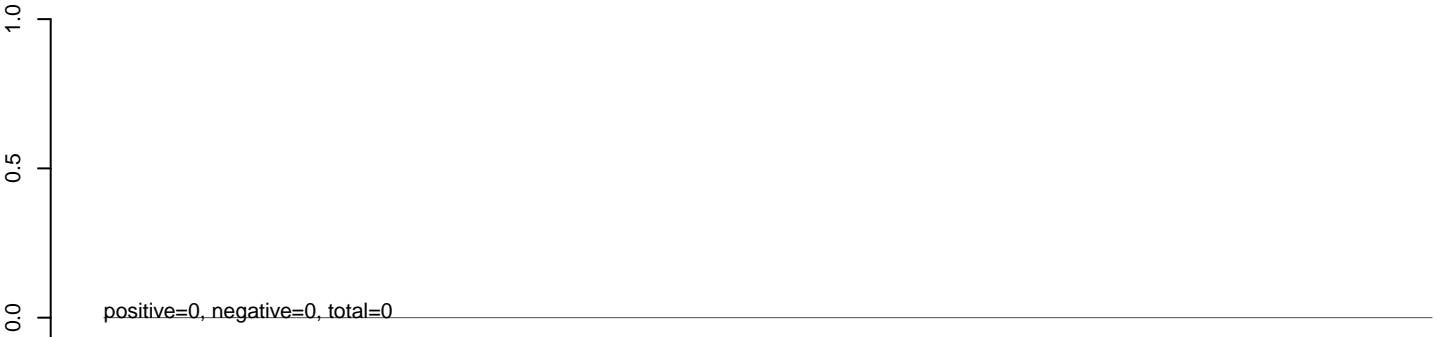
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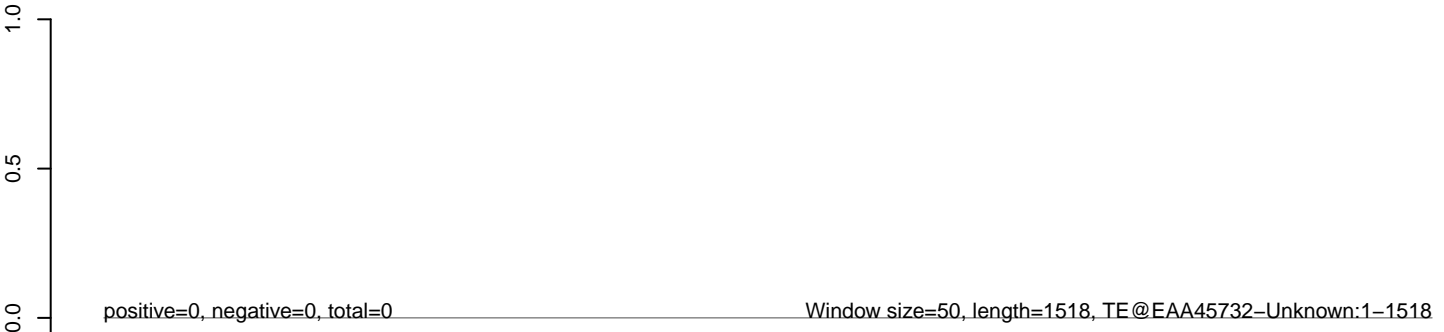
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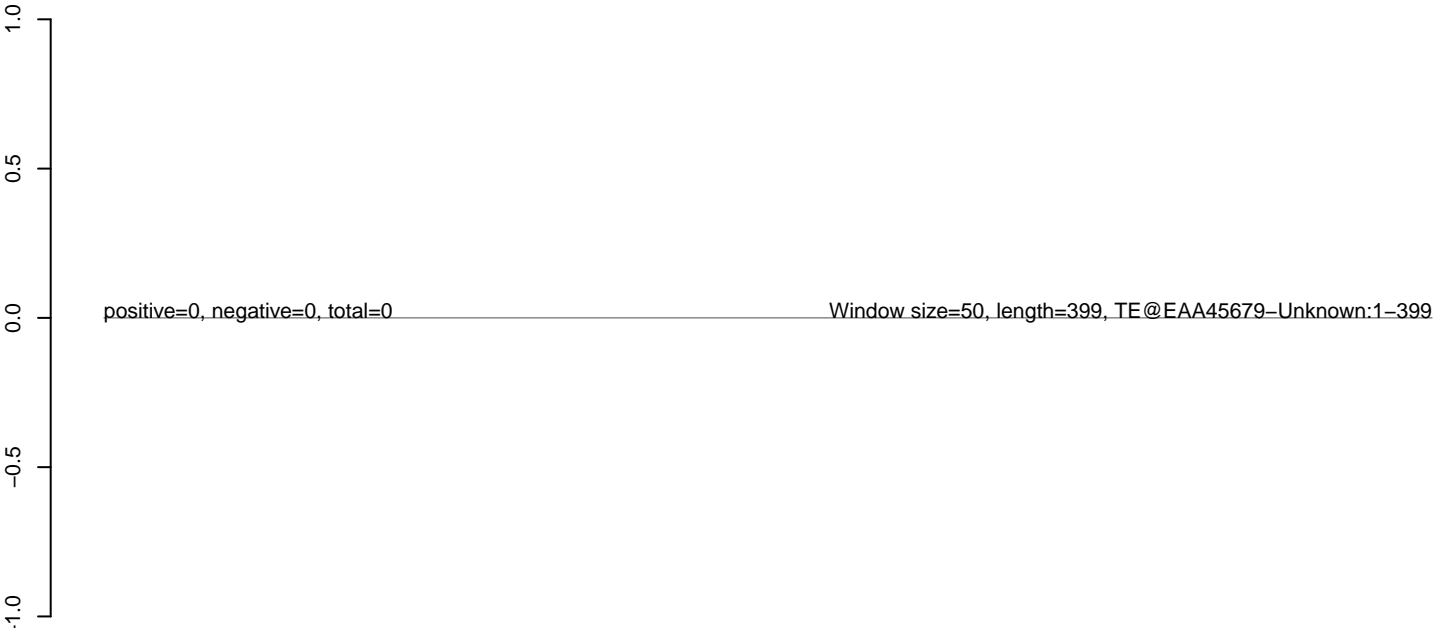
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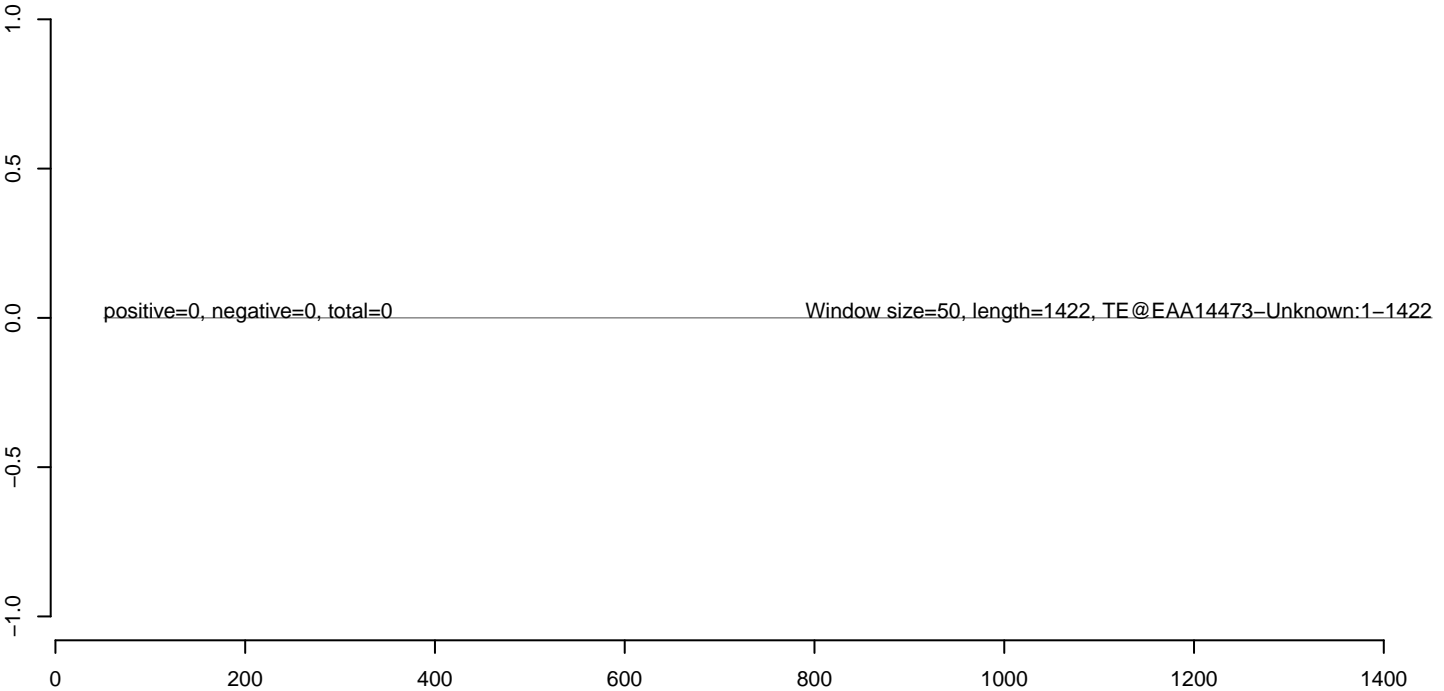
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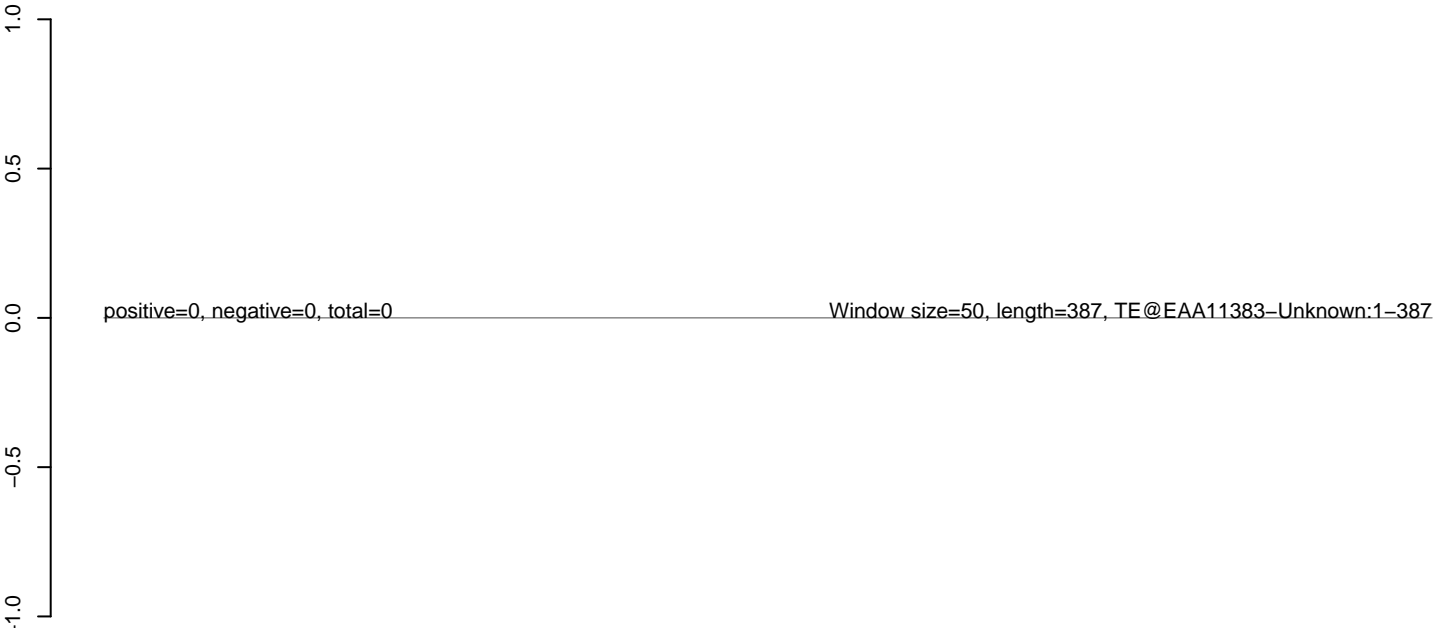
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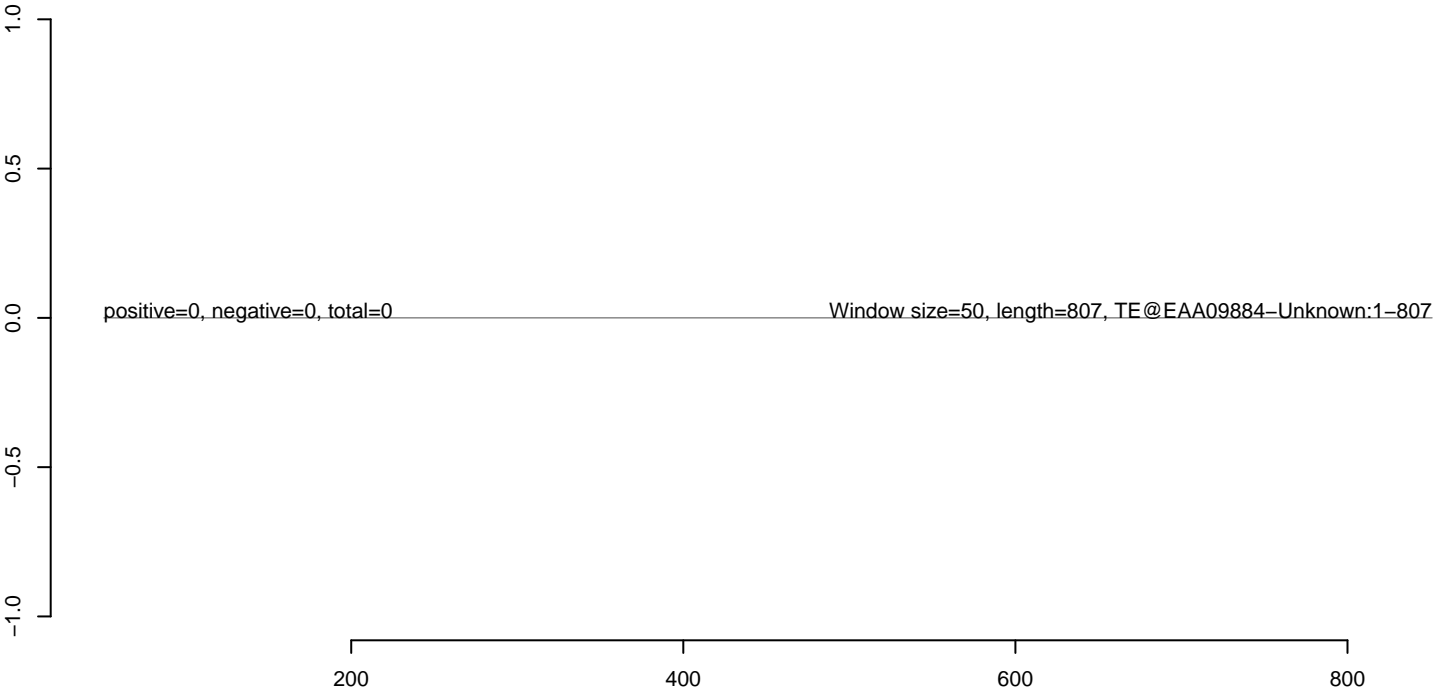
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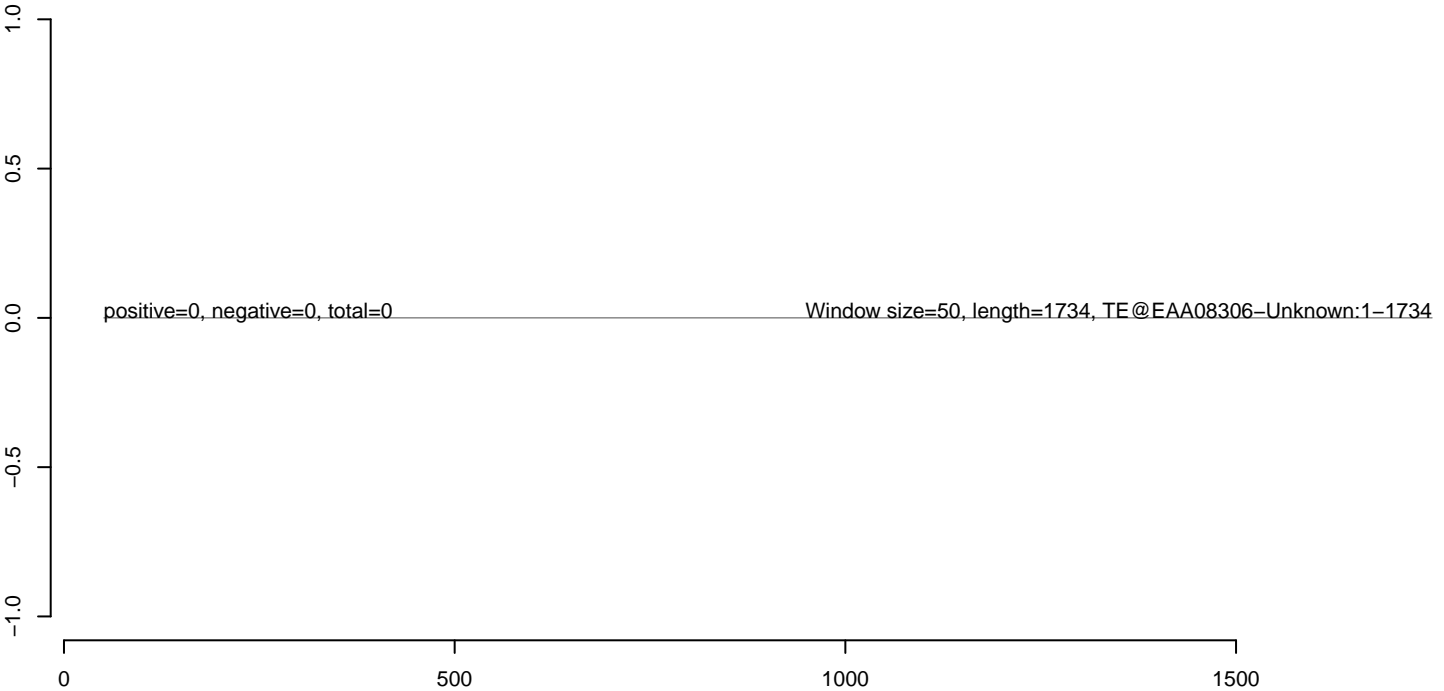
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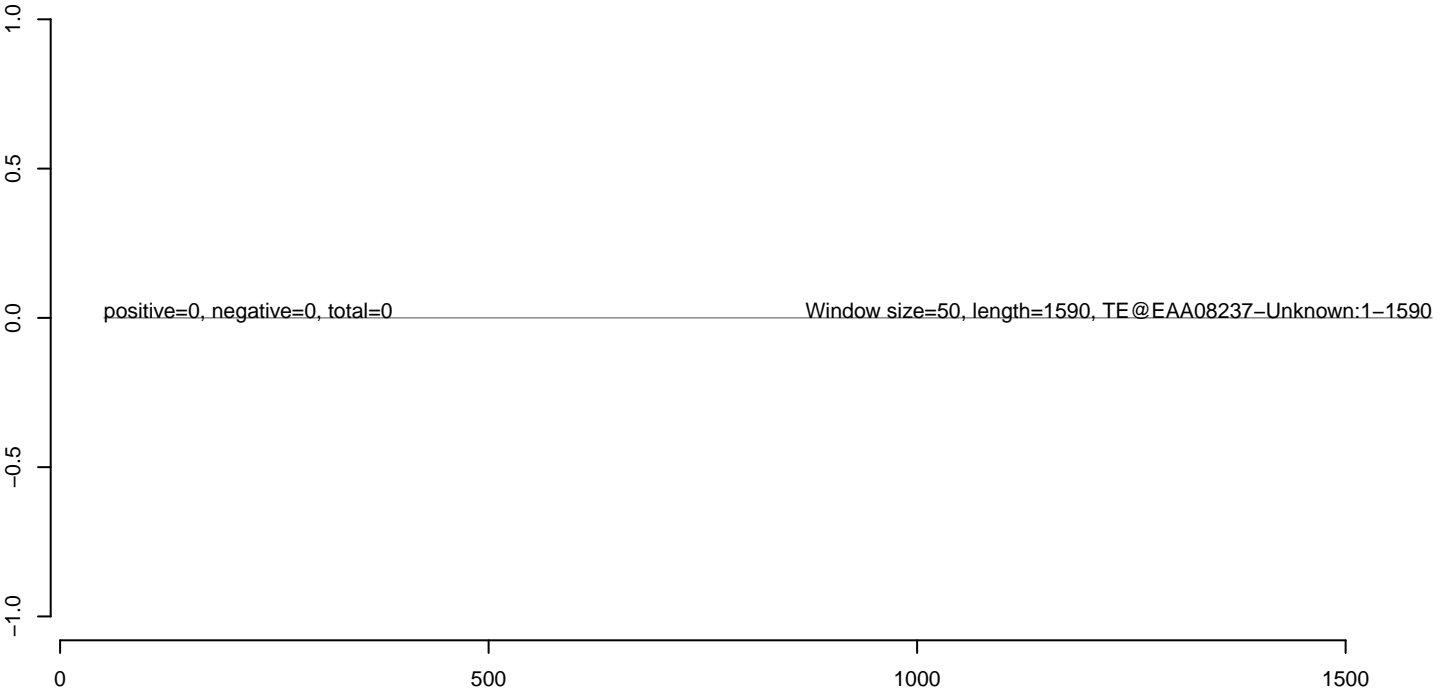
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AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



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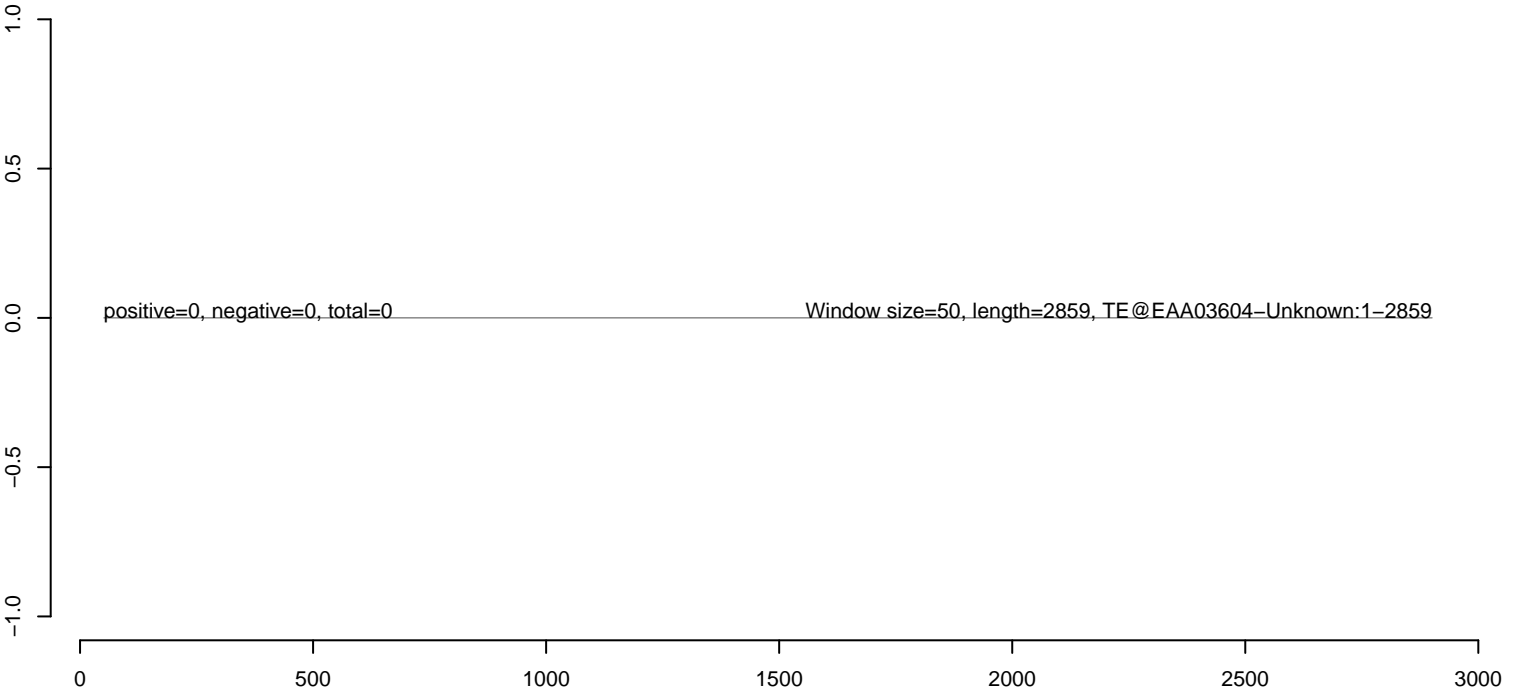
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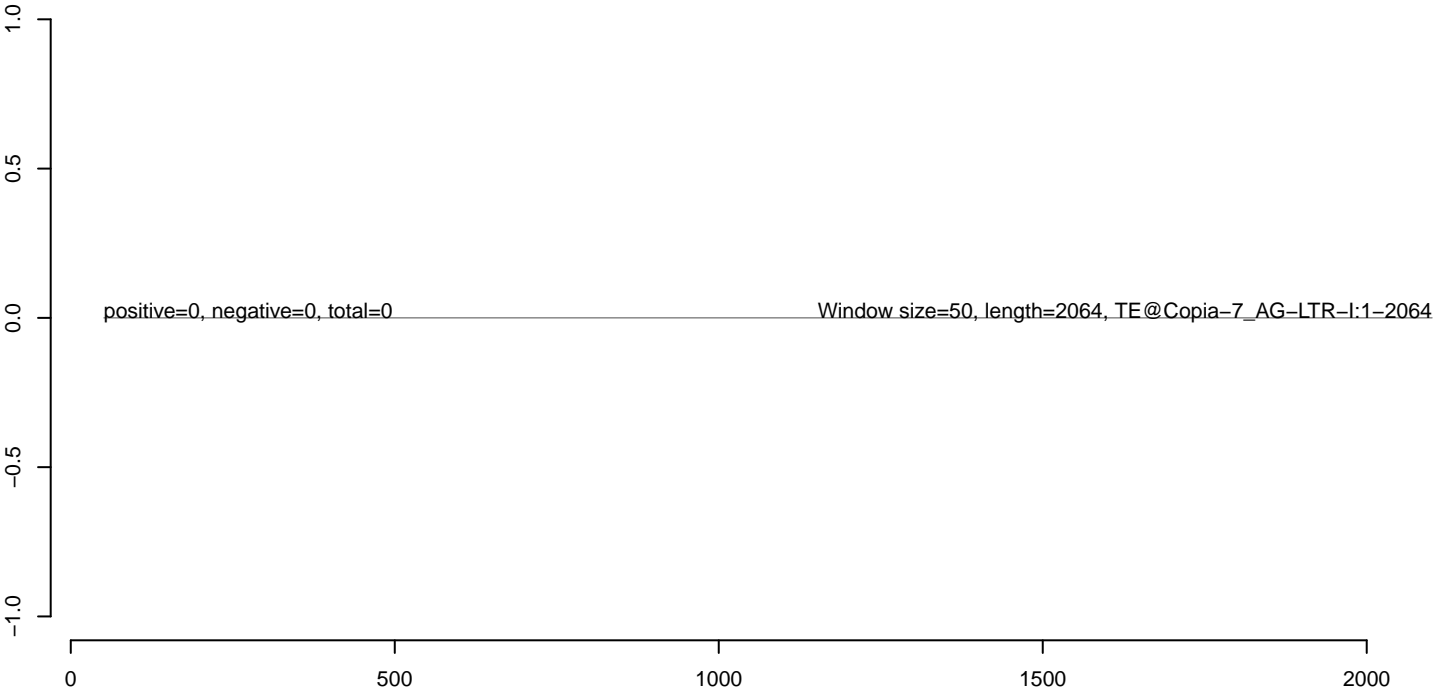
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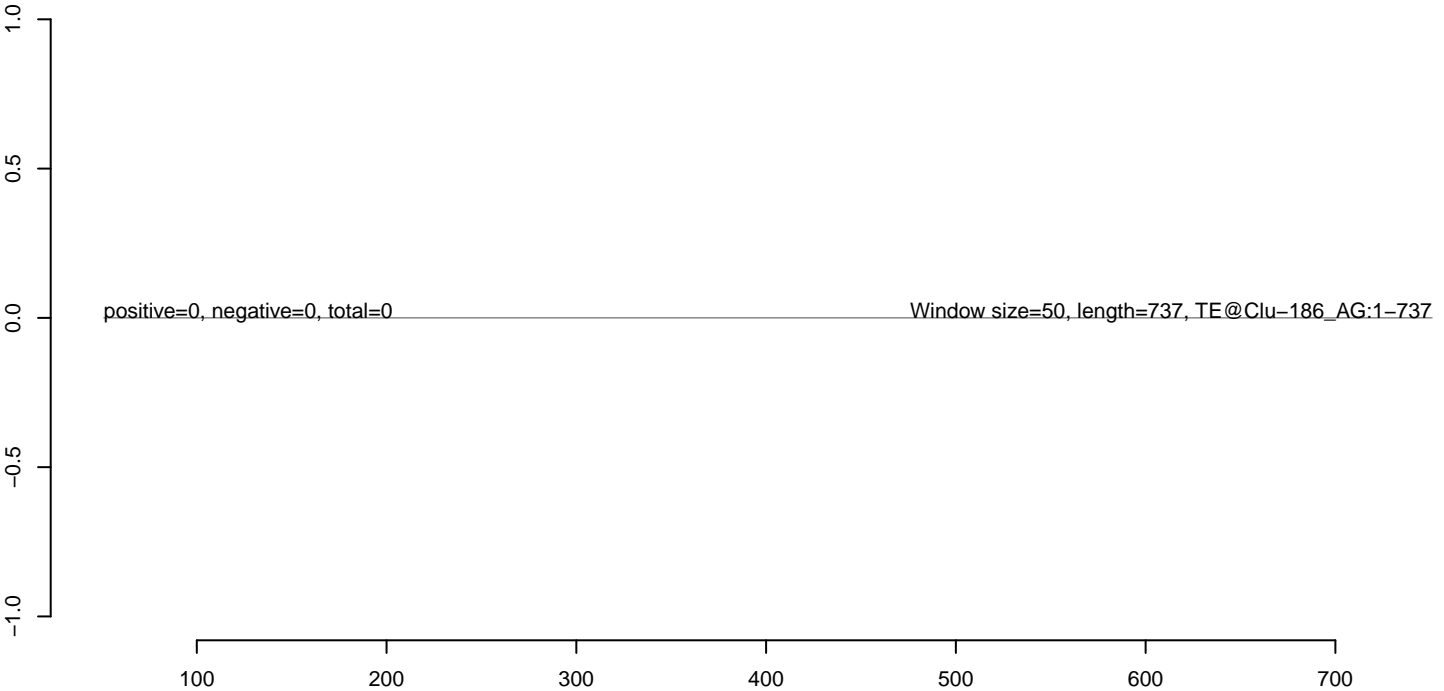
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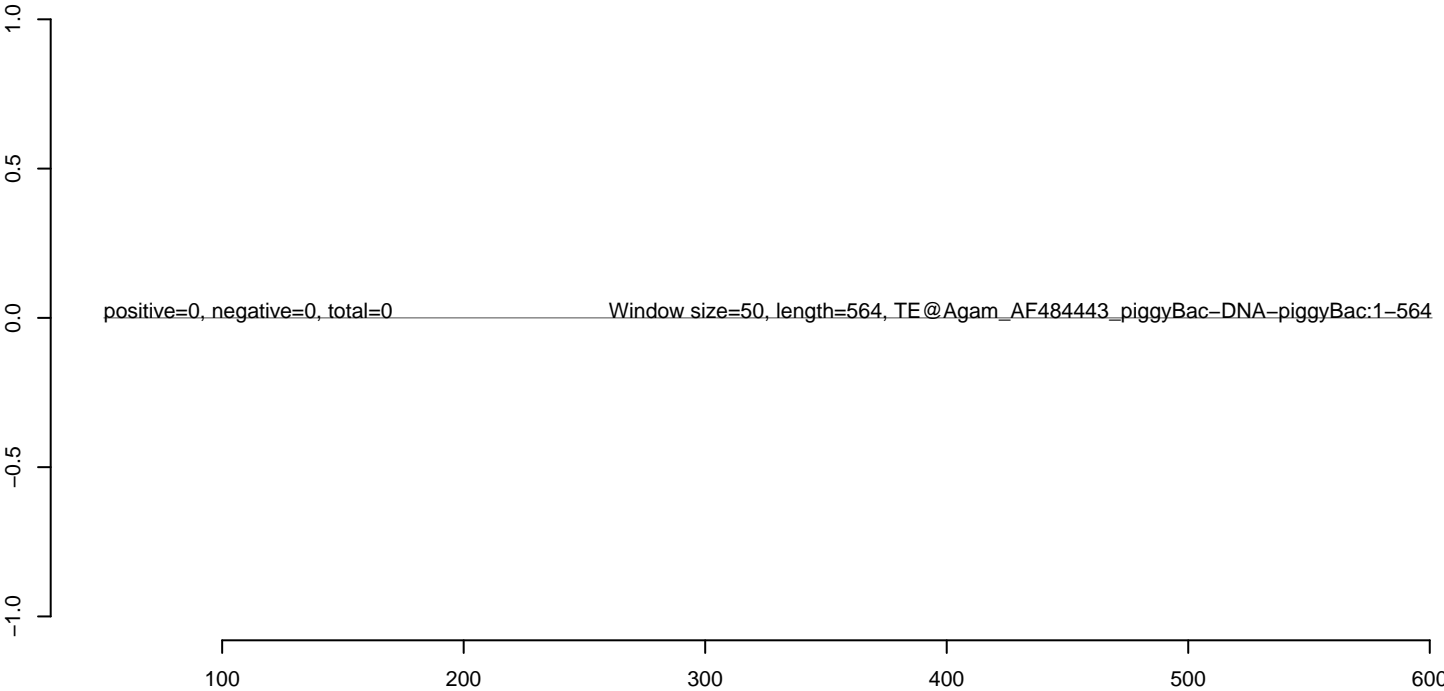
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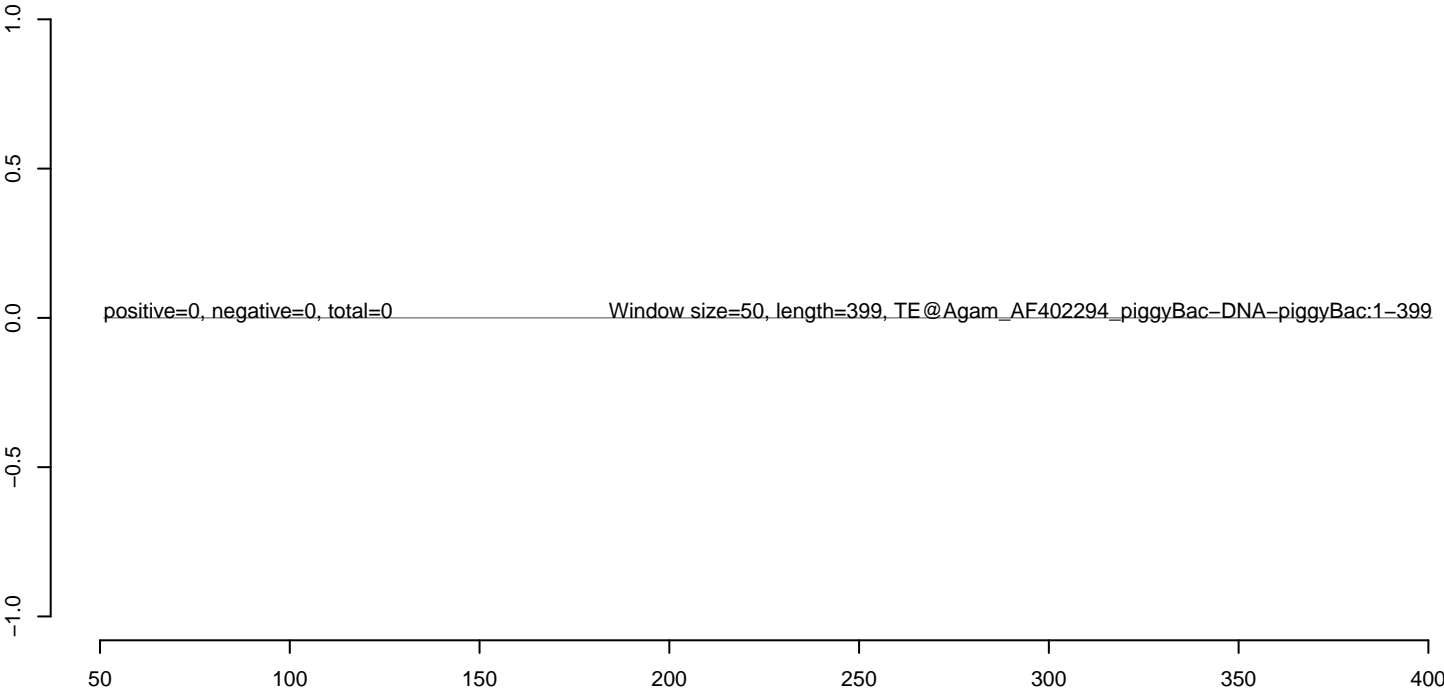
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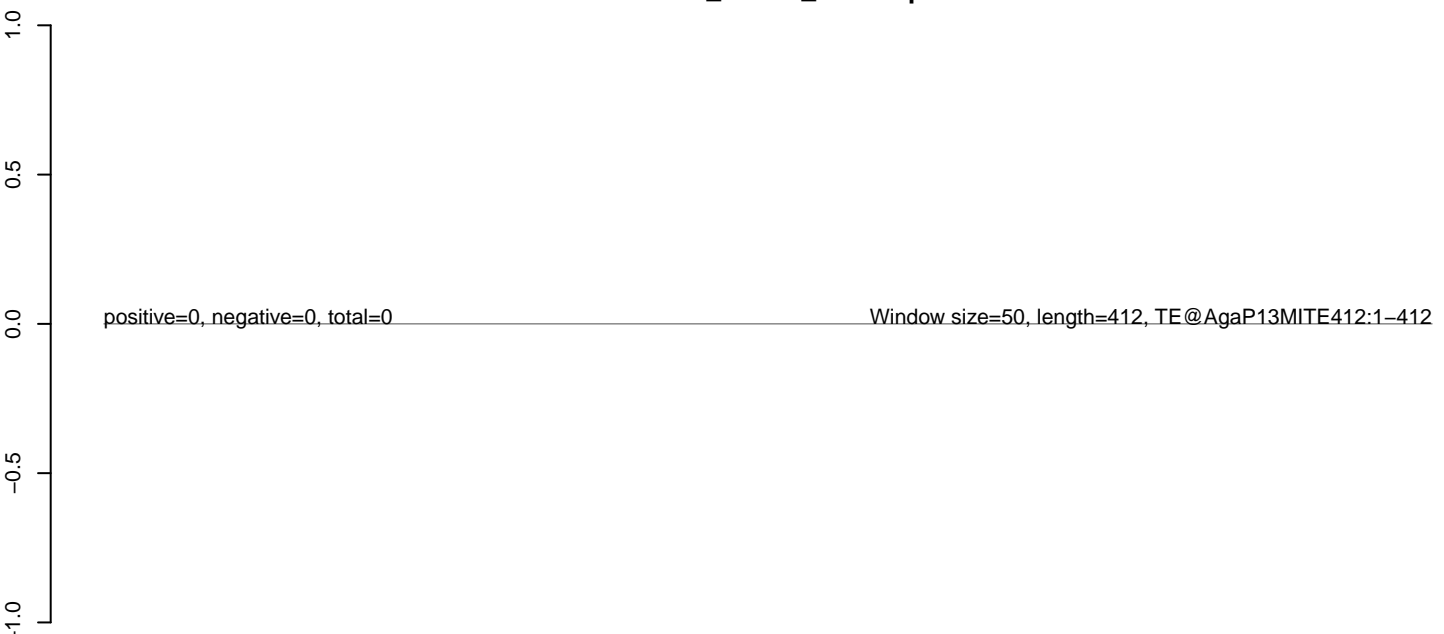
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AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep



0 1000 2000 3000 4000 5000

AnGam_Sua5b_cells.18_23.rep



AnGam_Sua5b_cells.24_35.rep



AnGam_Sua5b_cells.rep

