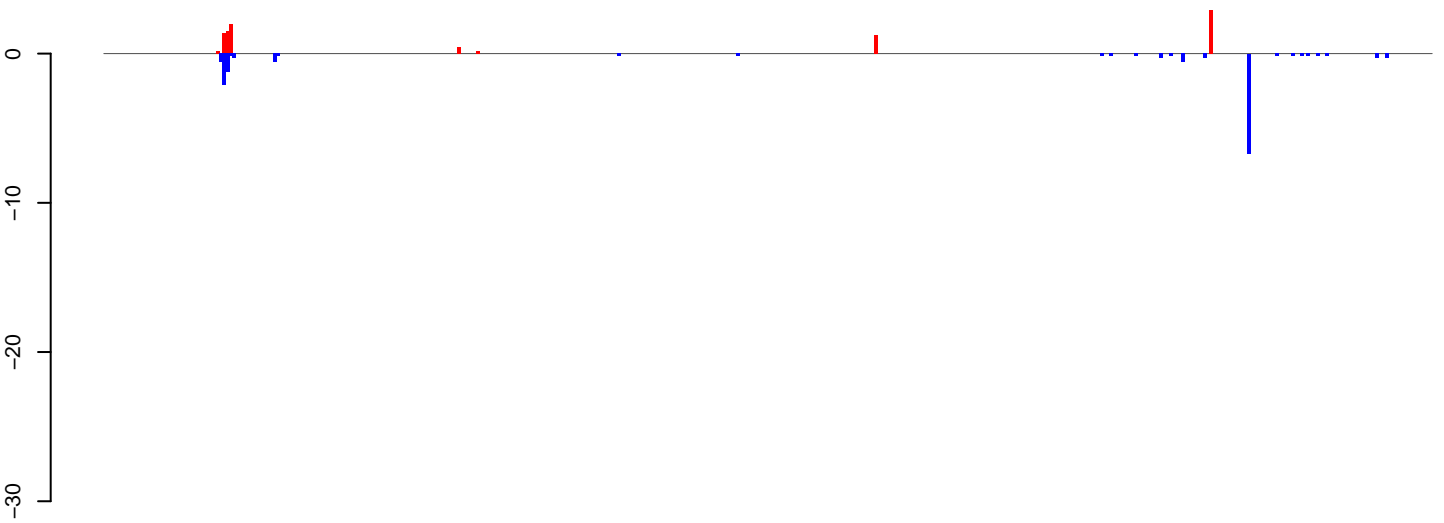
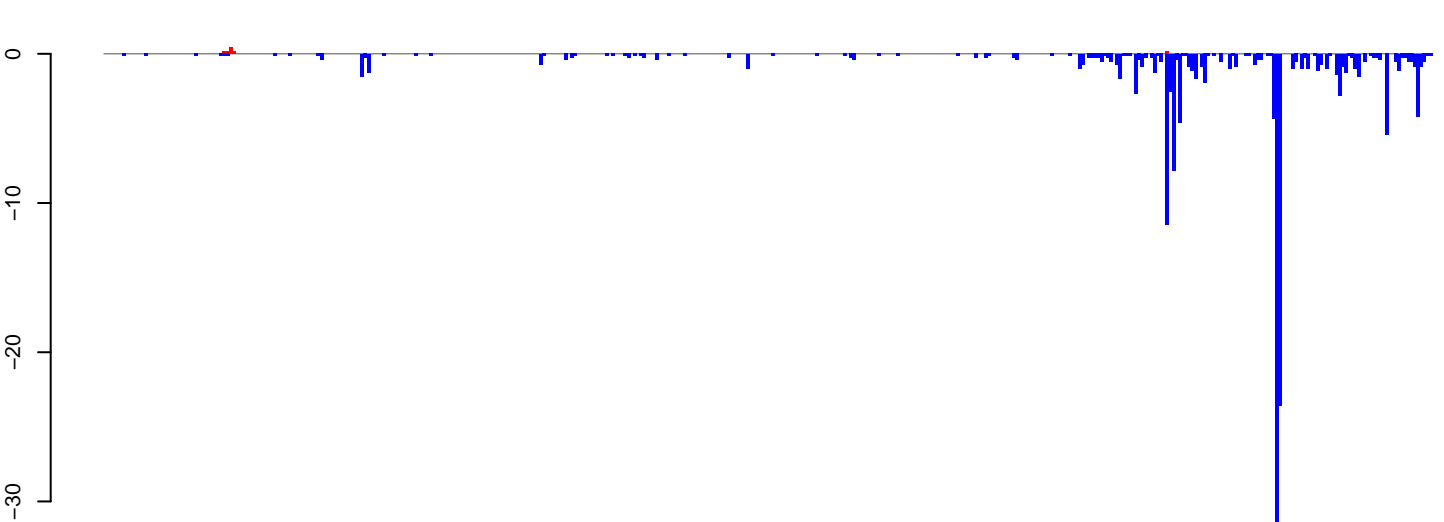


AeAlbo_C636_2dpi_mock3.18_23.rep



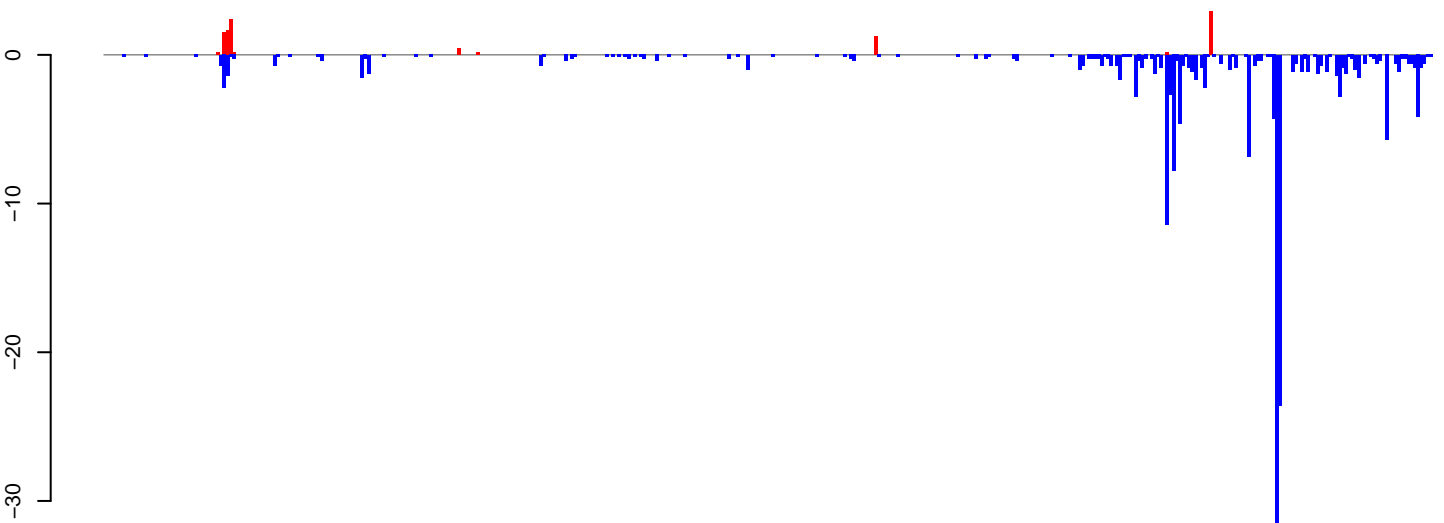
positive=10, negative=15, total=25

AeAlbo_C636_2dpi_mock3.24_35.rep



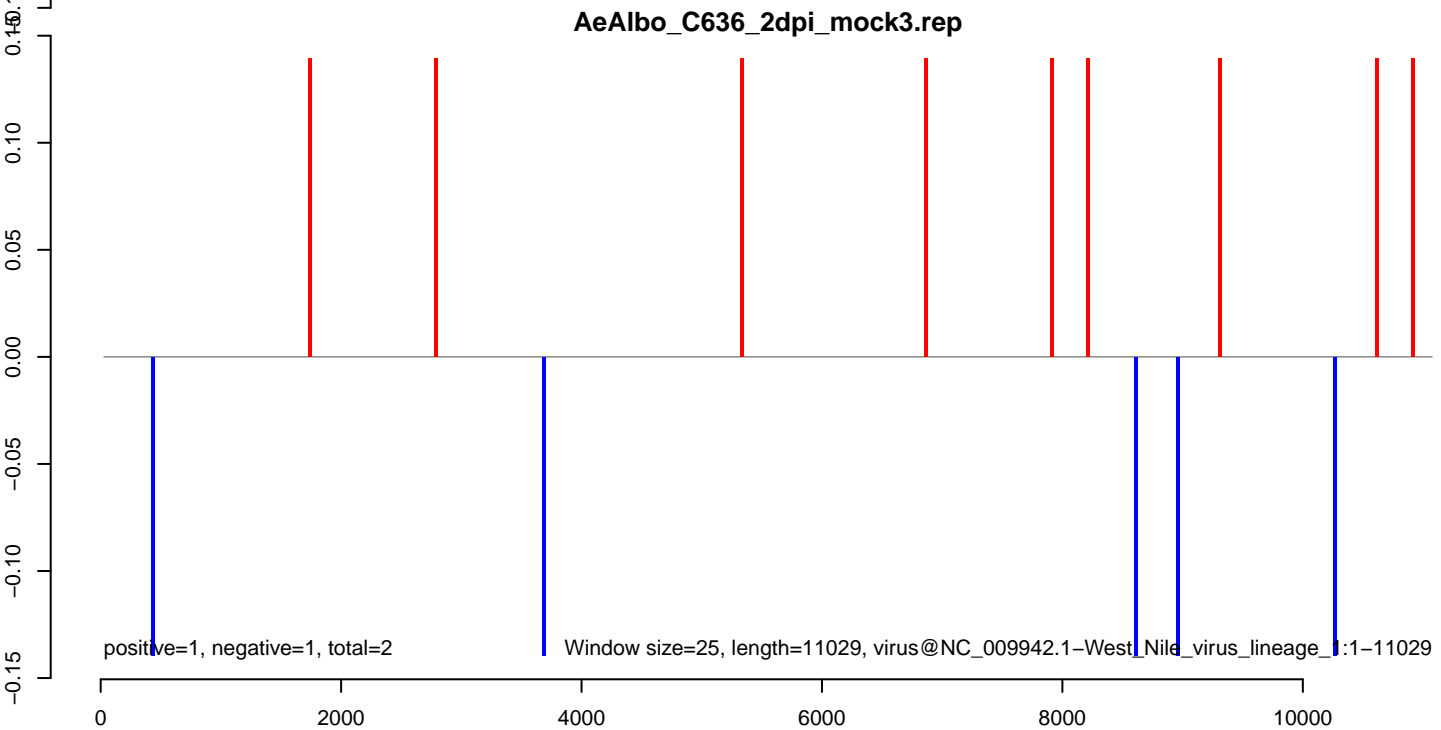
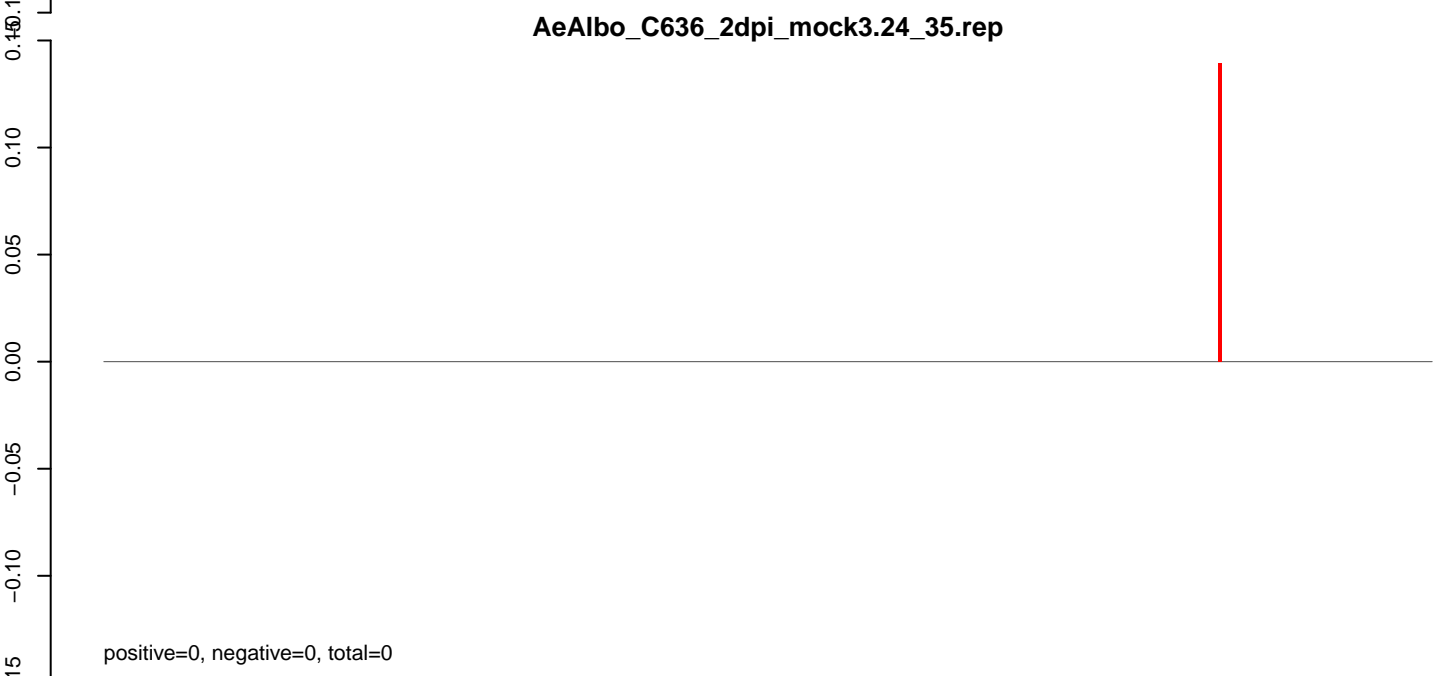
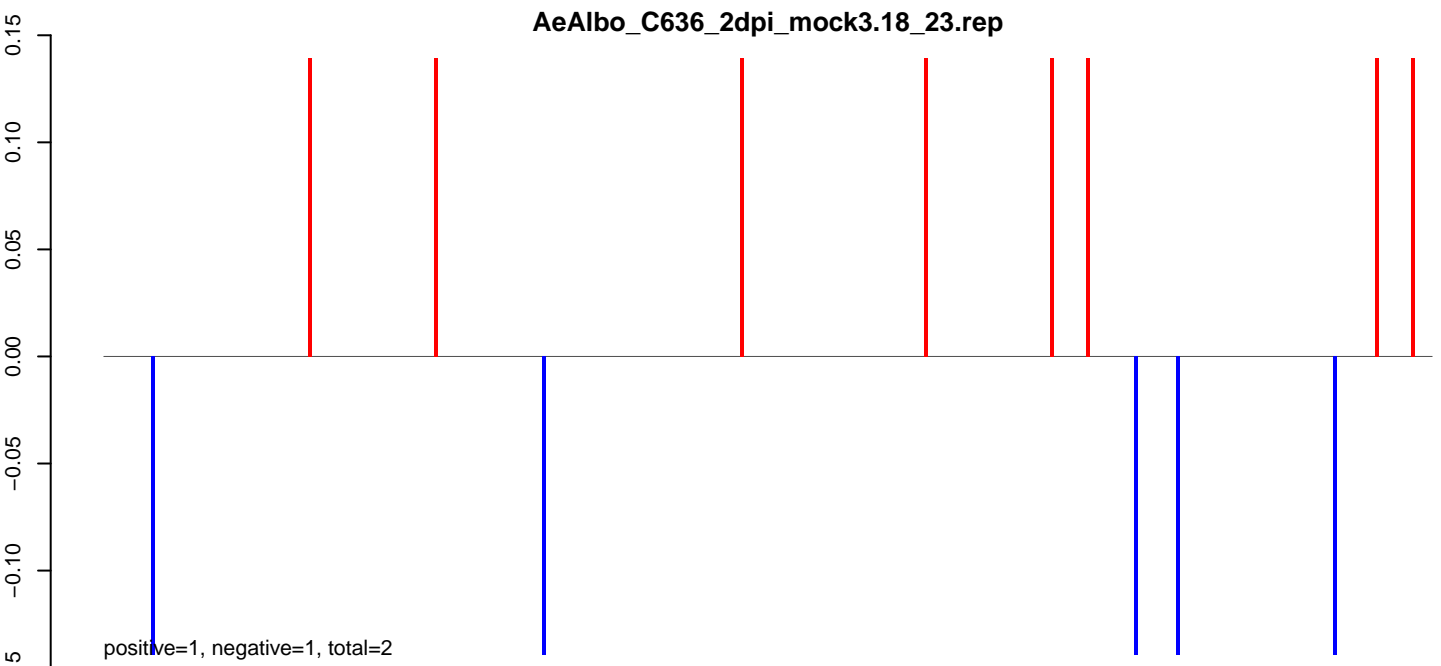
positive=1, negative=163, total=164

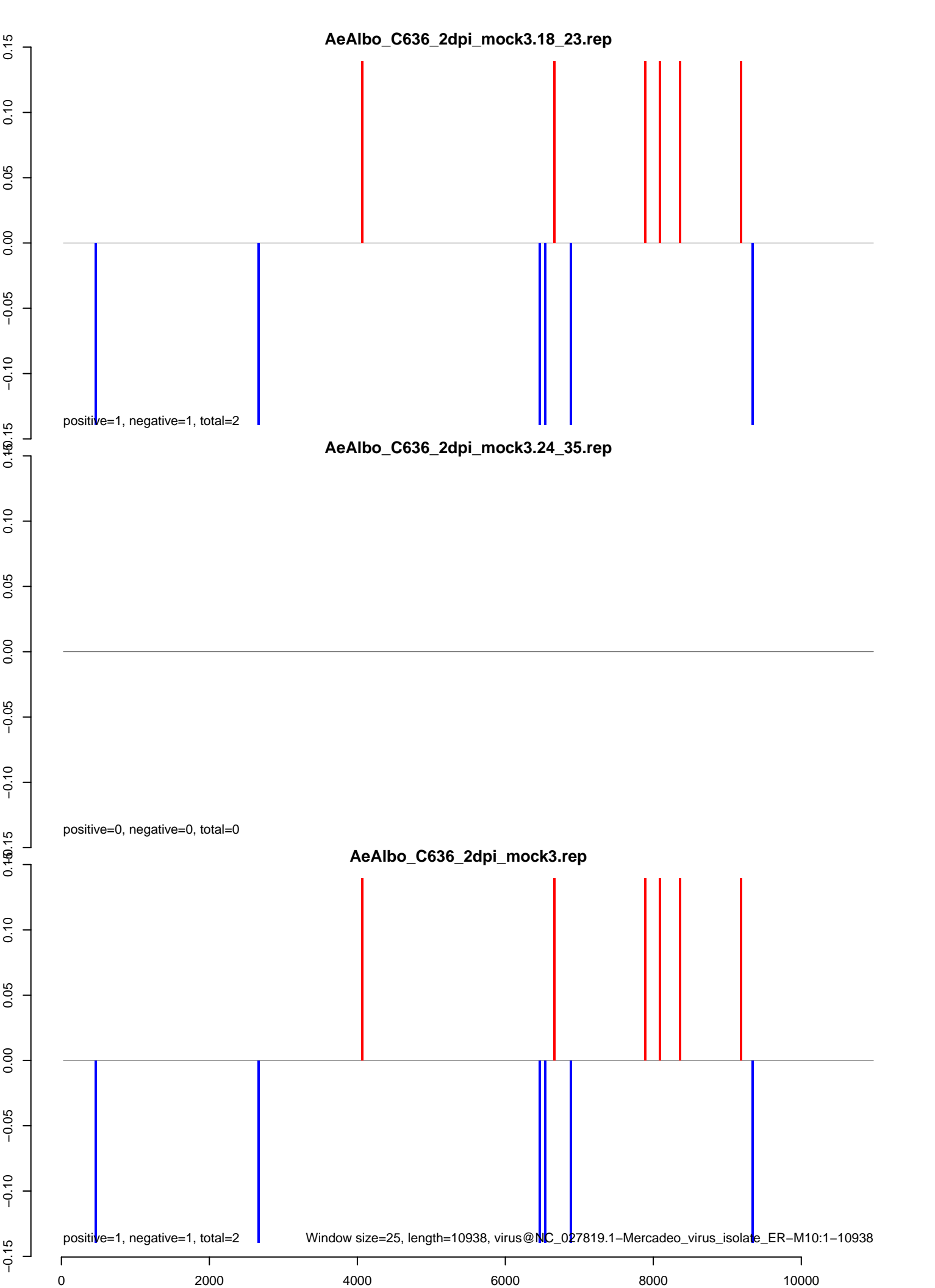
AeAlbo_C636_2dpi_mock3.rep



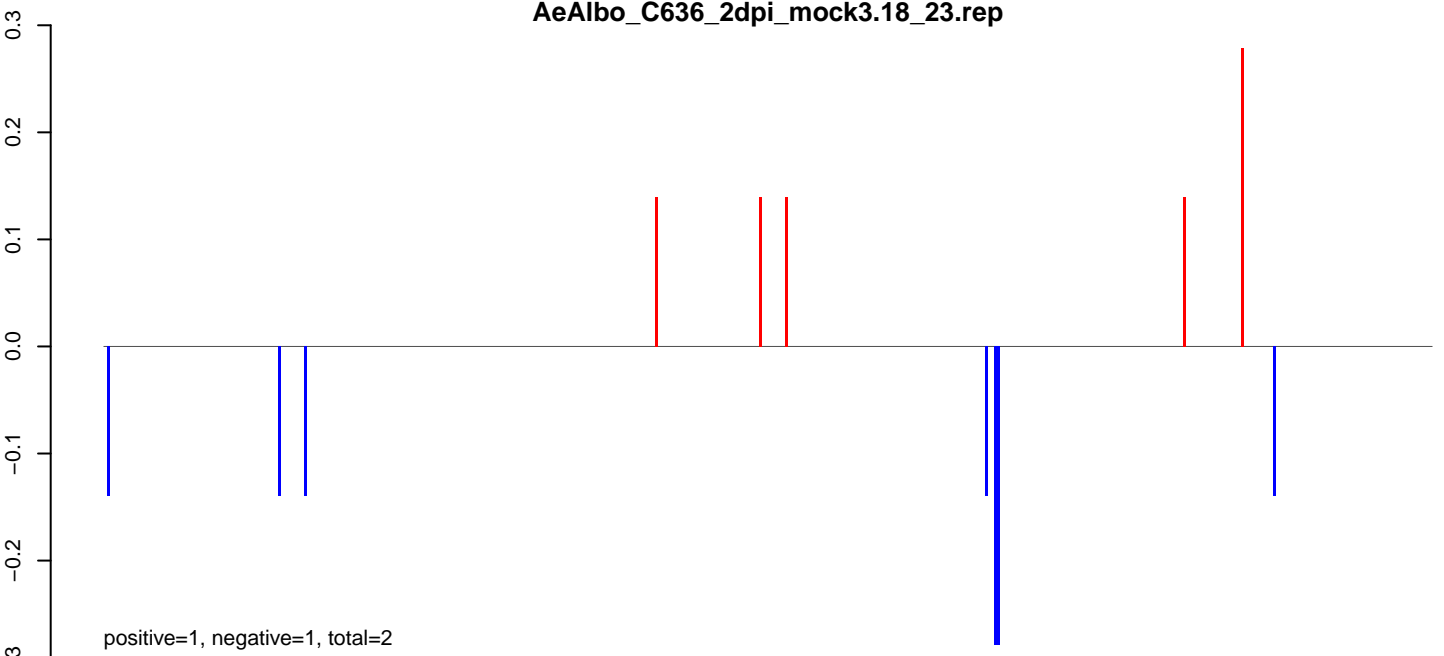
size=25, length=10587, positive=10, negative=15, total=25
size=164, length=10587, positive=1, negative=163, total=164
size=25, length=10587, positive=10, negative=15, total=25
size=164, length=10587, positive=1, negative=163, total=164

0 2000 4000 6000 8000 10000

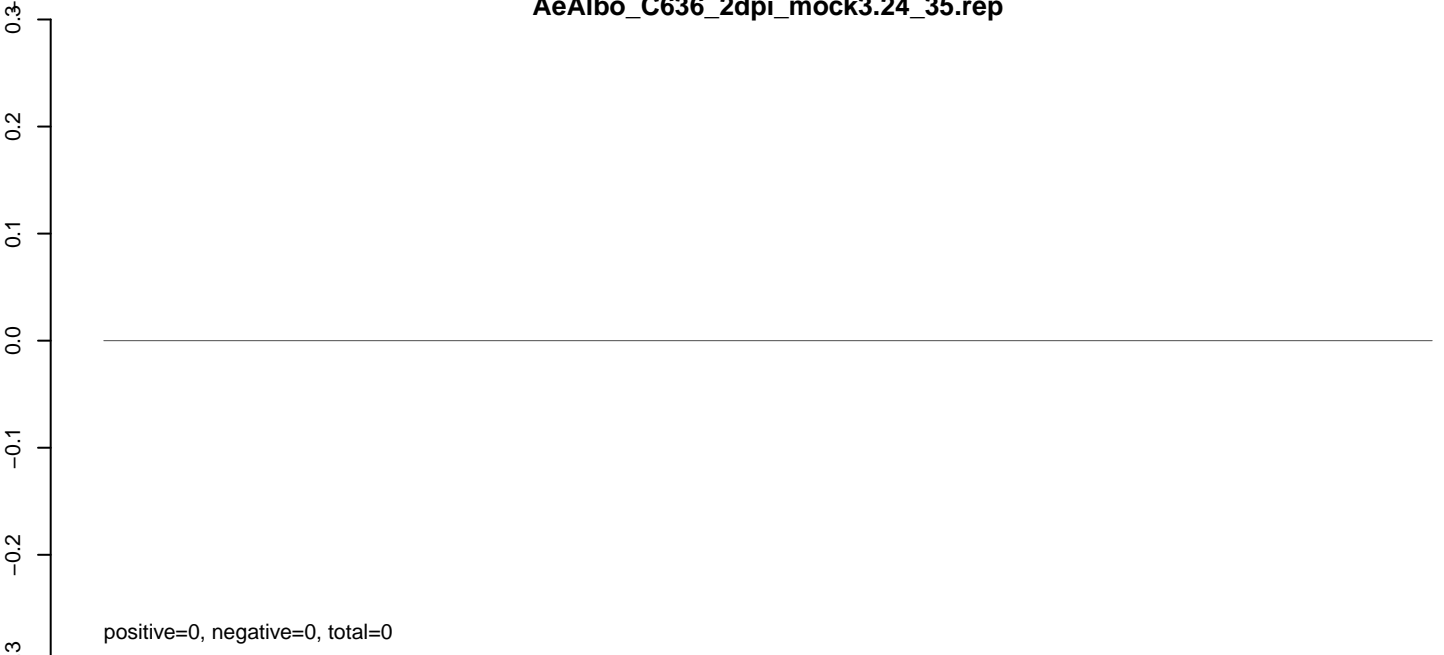




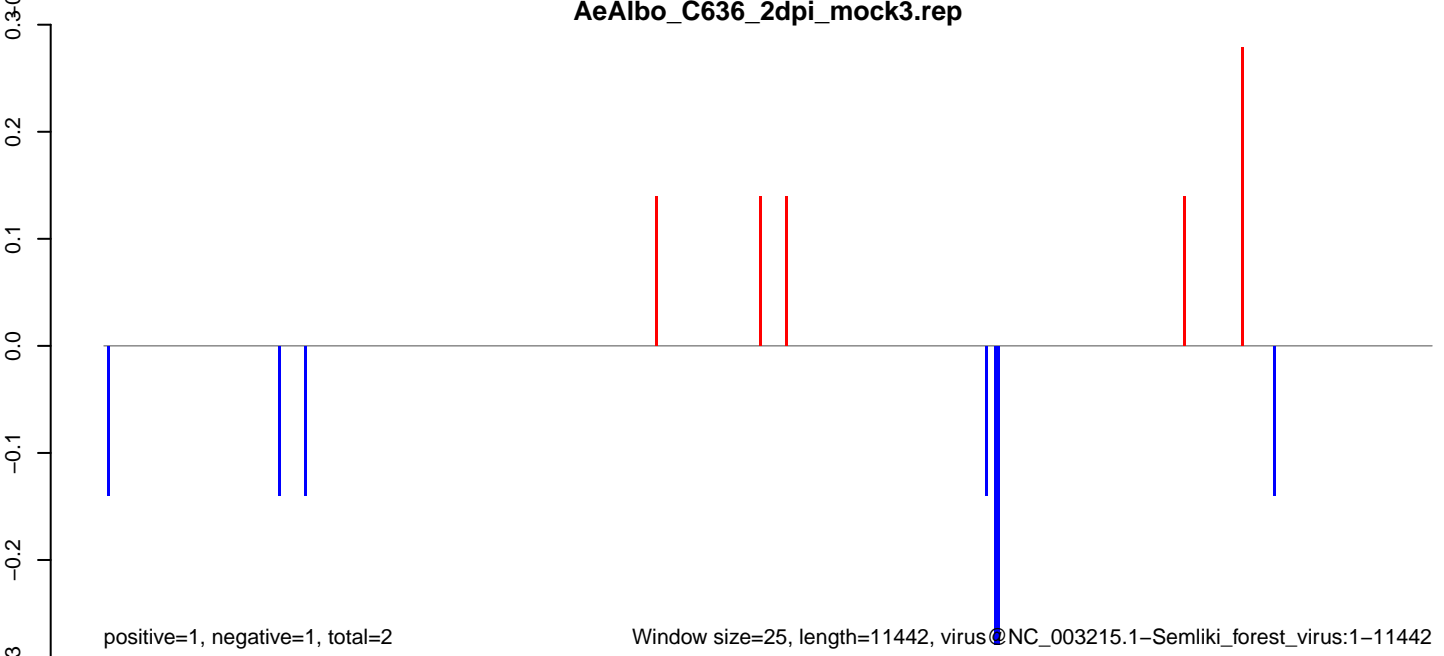
AeAlbo_C636_2dpi_mock3.18_23.rep



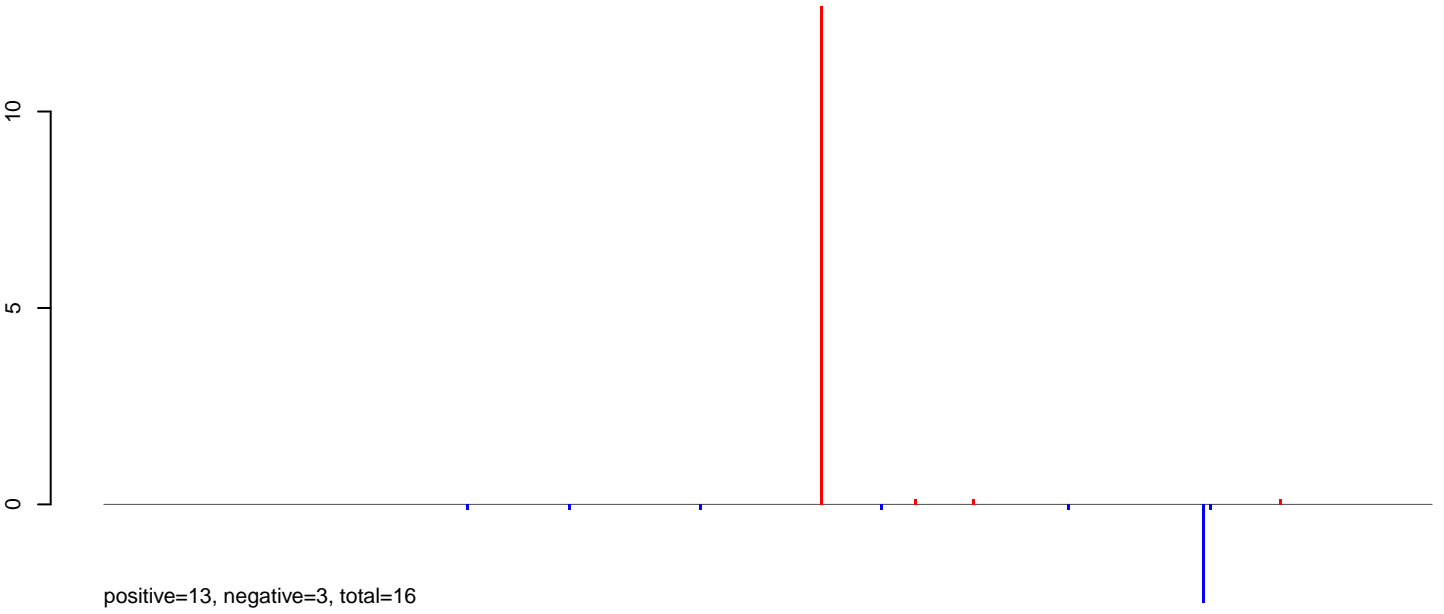
AeAlbo_C636_2dpi_mock3.24_35.rep



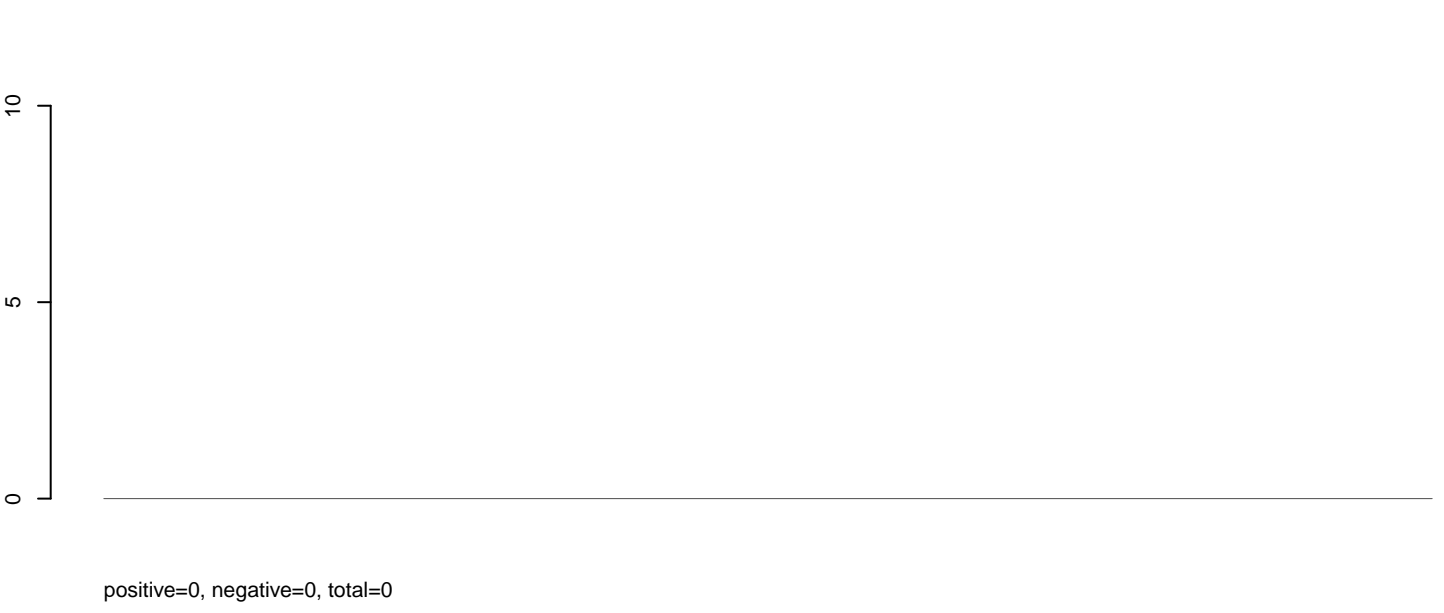
AeAlbo_C636_2dpi_mock3.rep



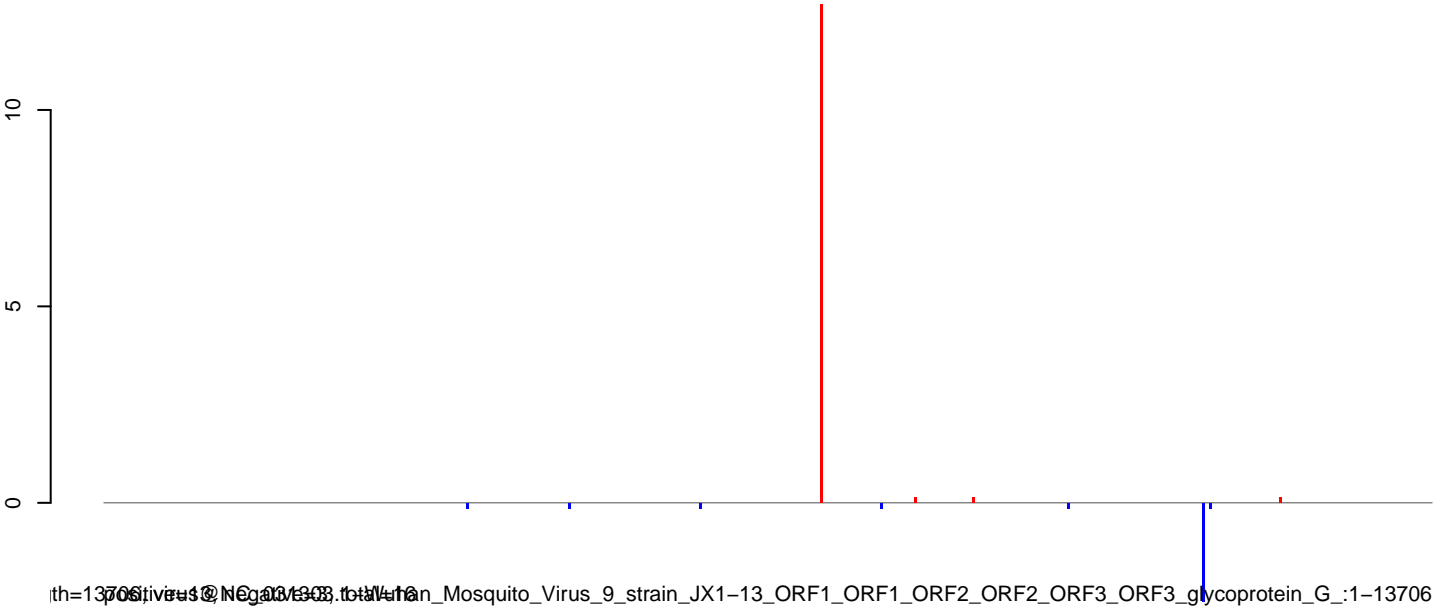
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

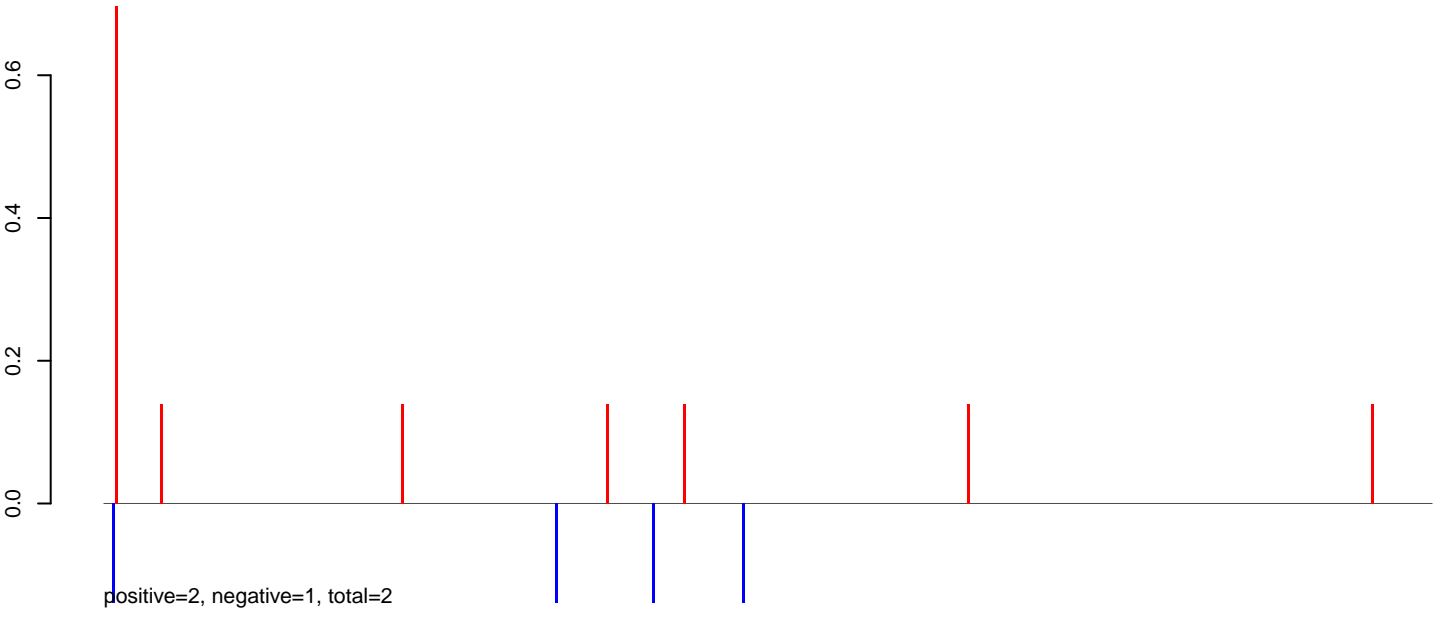


AeAlbo_C636_2dpi_mock3.rep



th=13706, positive=13, negative=3, total=16
Aedes albopictus Neglab0193, total=16
Aedes albopictus Neglab0193_Mosquito_Virus_9_strain_JX1-13_ORF1_ORF1_ORF2_ORF2_ORF3_ORF3_glycoprotein_G_:1-13706

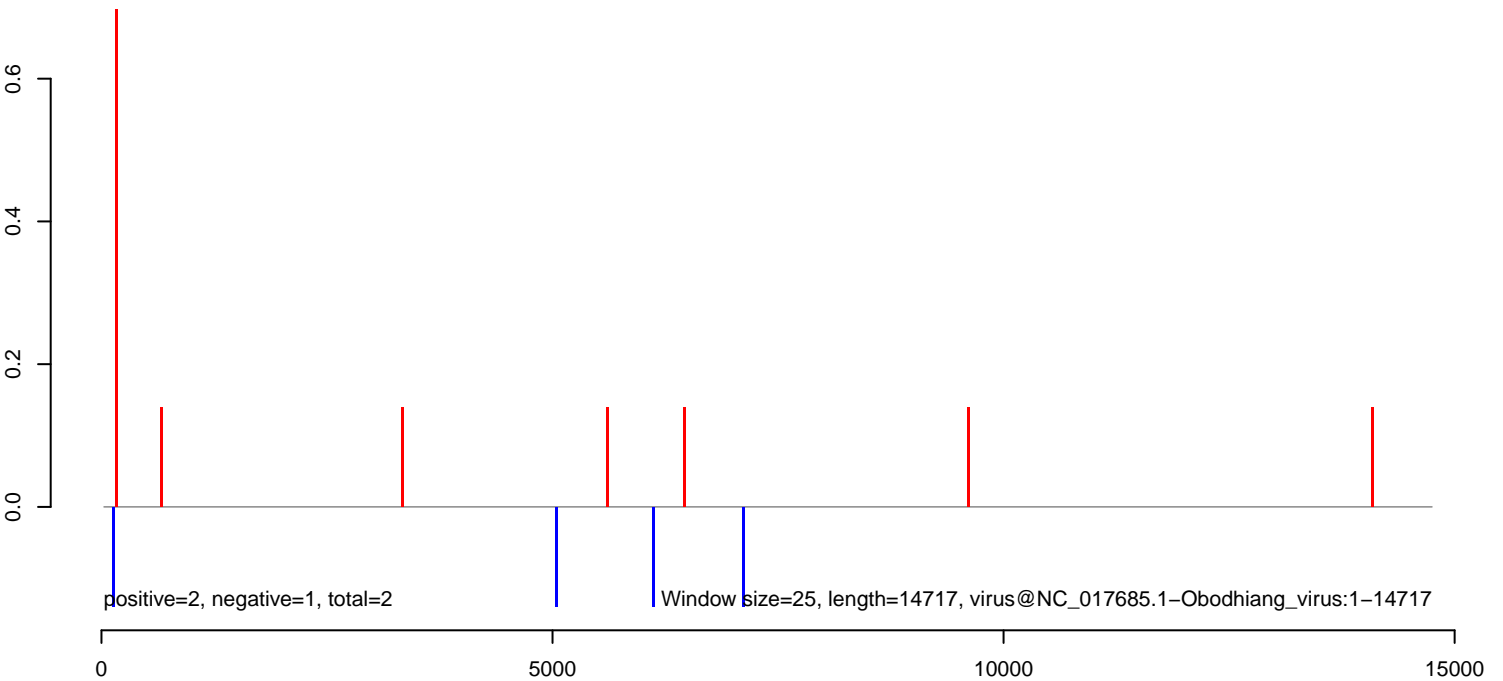
AeAlbo_C636_2dpi_mock3.18_23.rep



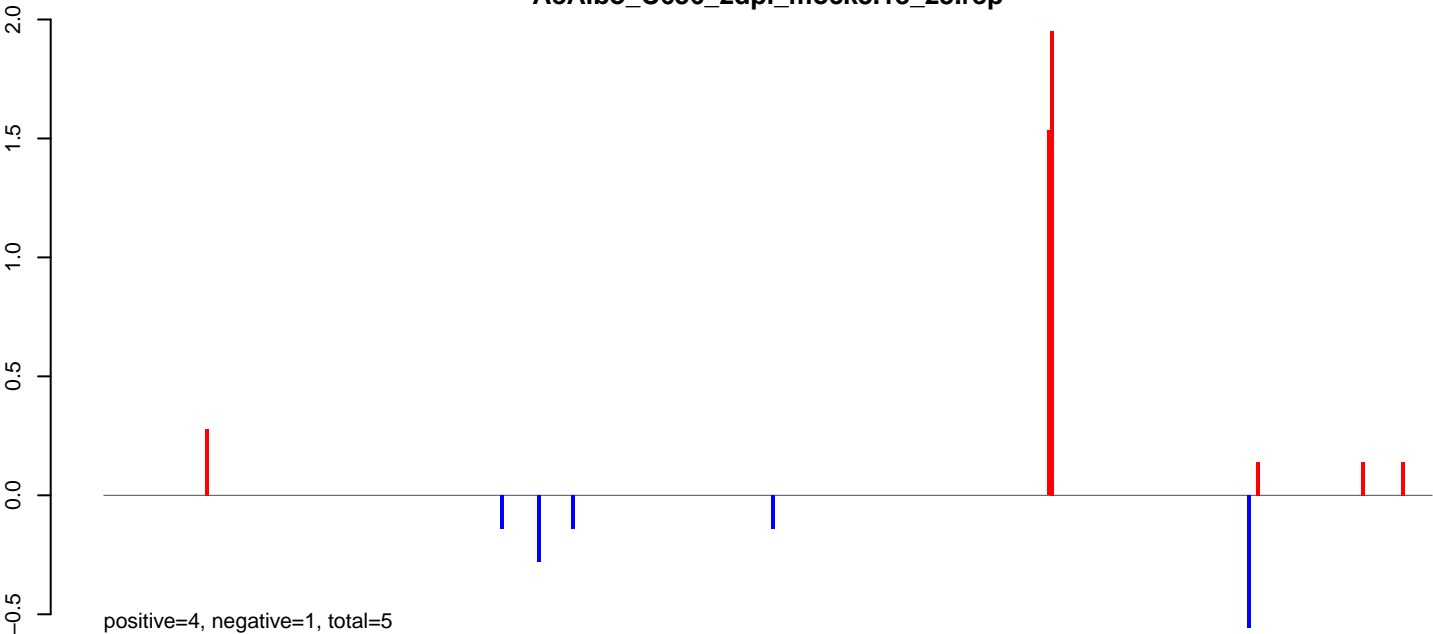
AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



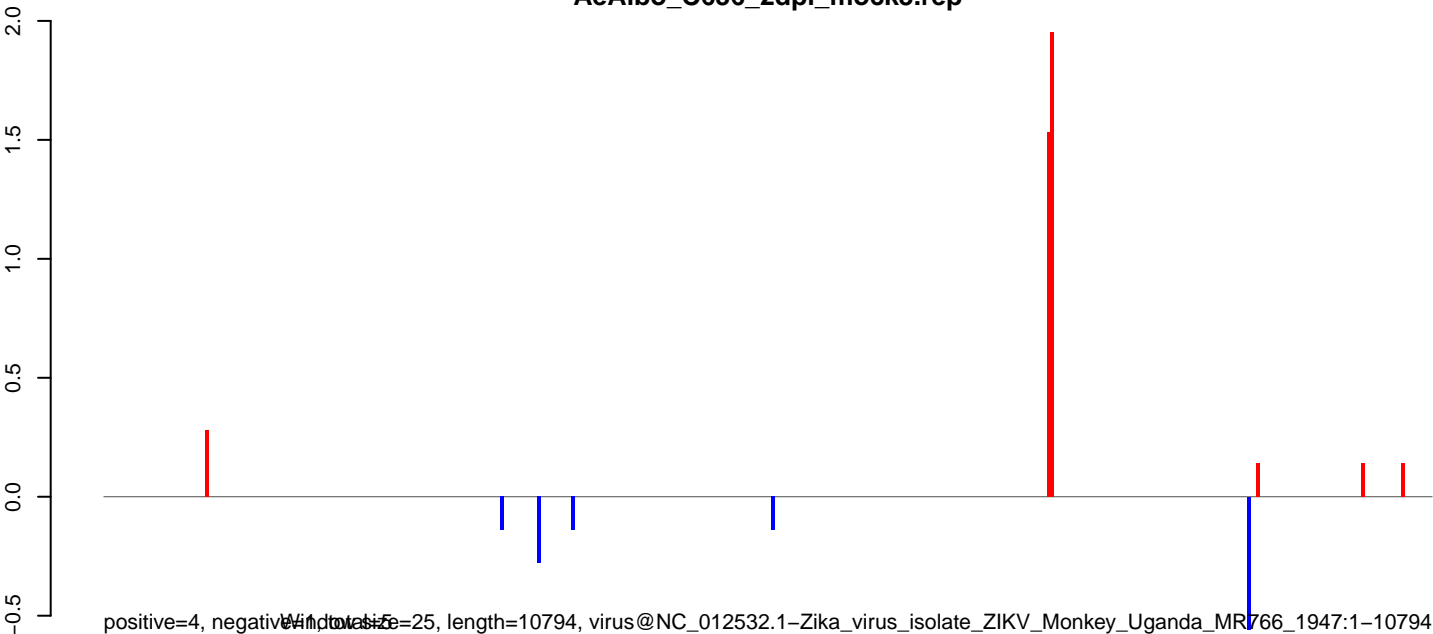
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



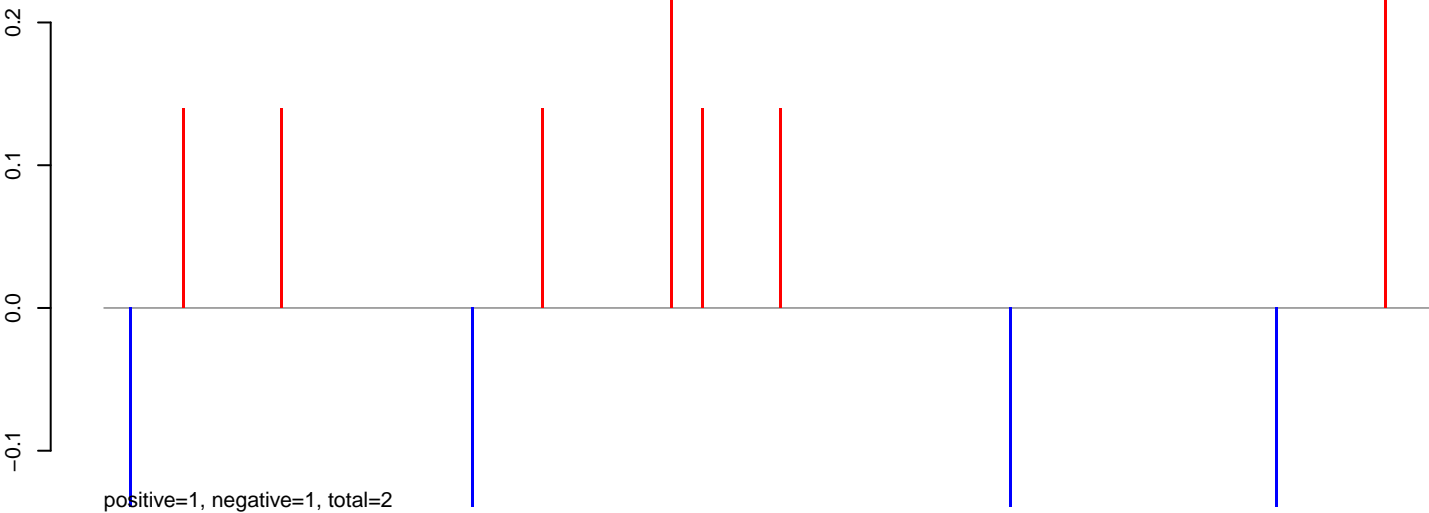
AeAlbo_C636_2dpi_mock3.rep



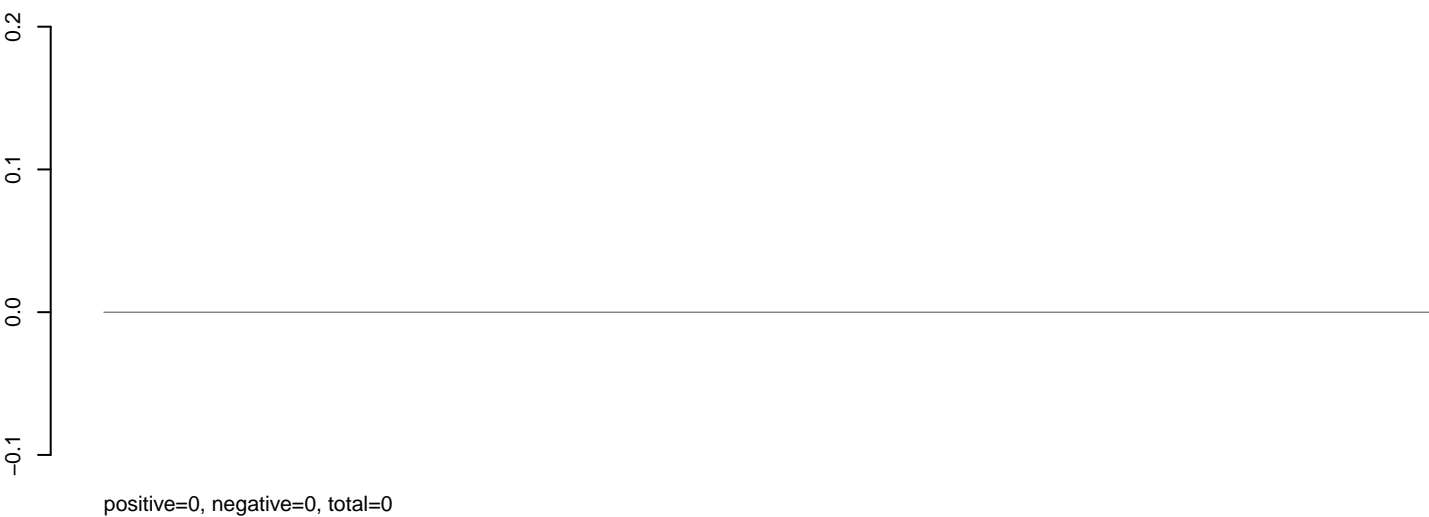
Window size=25, length=10794, virus@NC_012532.1-Zika_virus_isolate_ZIKV_Monkey_Uganda_MR766_1947:1-10794

0 2000 4000 6000 8000 10000

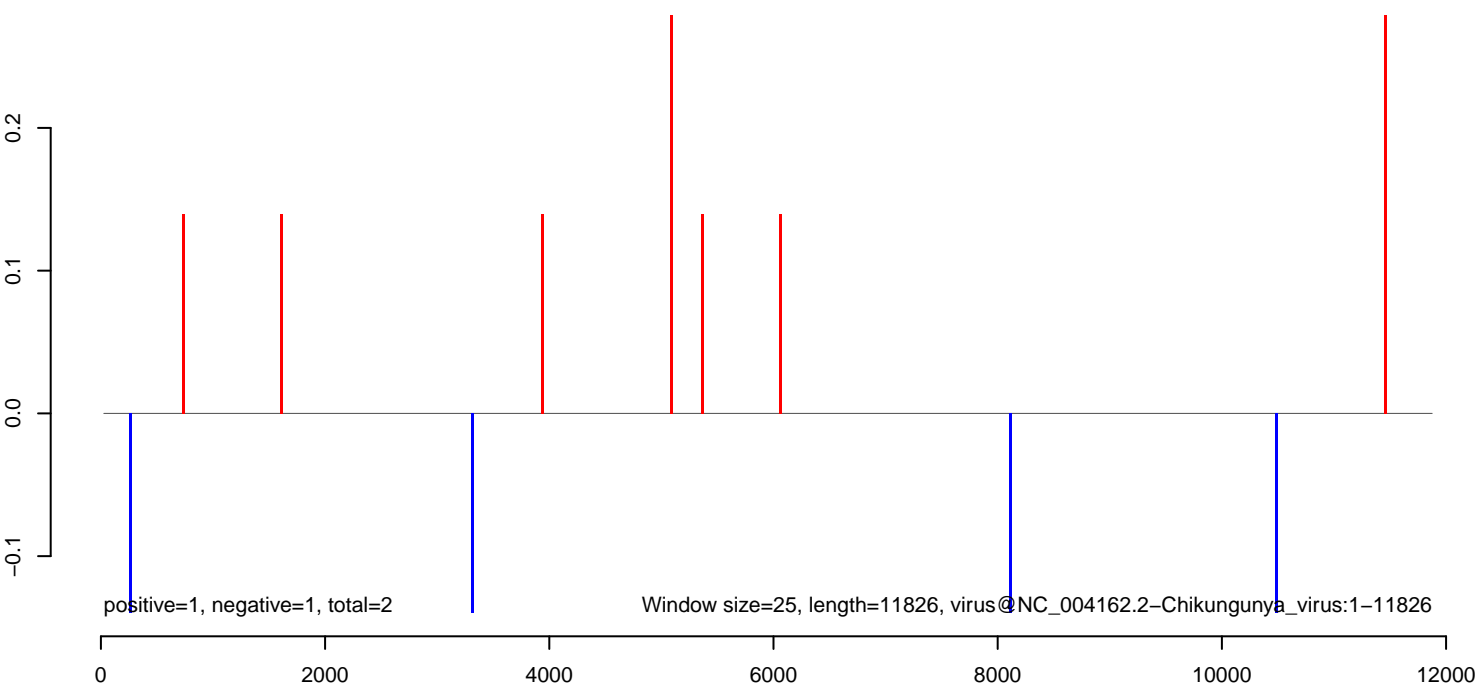
AeAlbo_C636_2dpi_mock3.18_23.rep



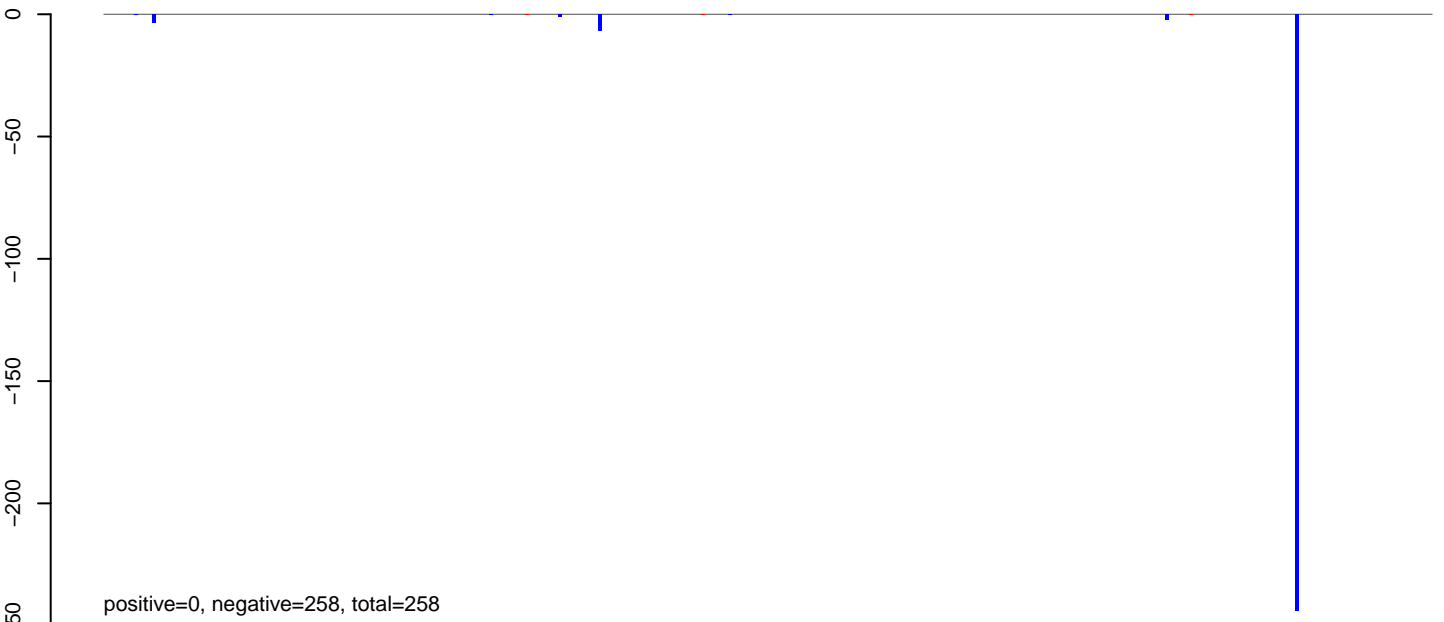
AeAlbo_C636_2dpi_mock3.24_35.rep



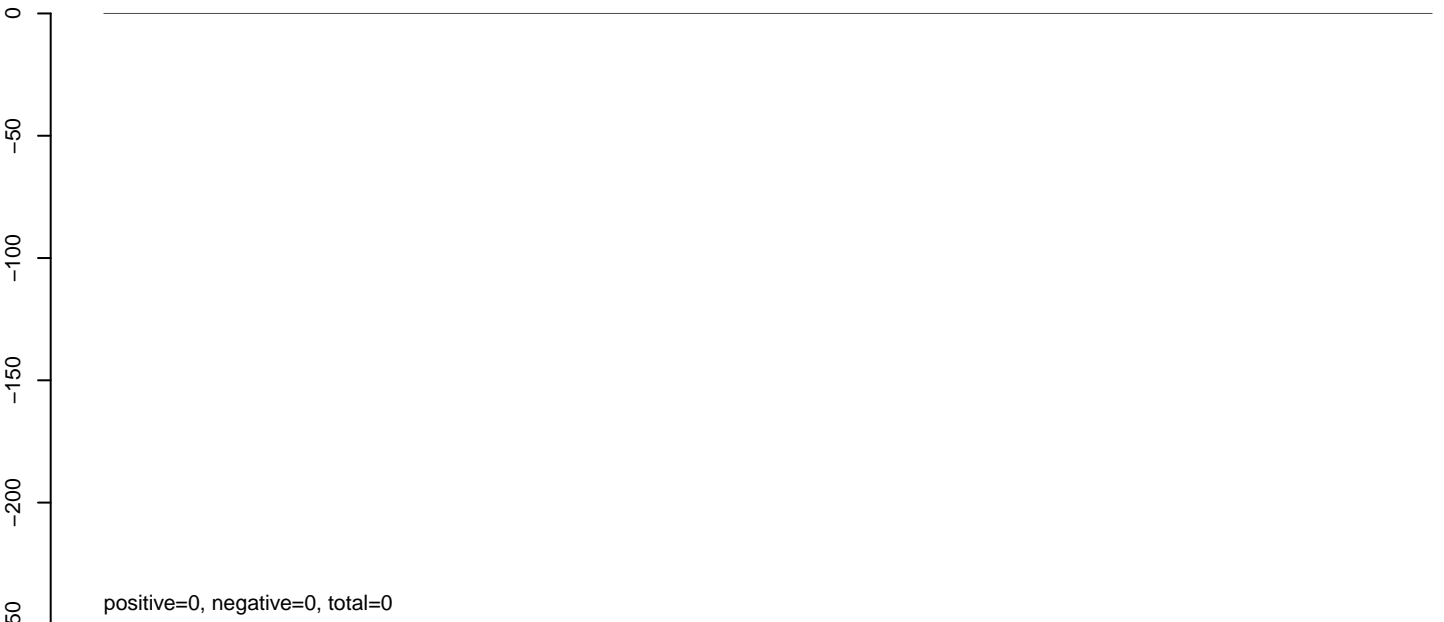
AeAlbo_C636_2dpi_mock3.rep



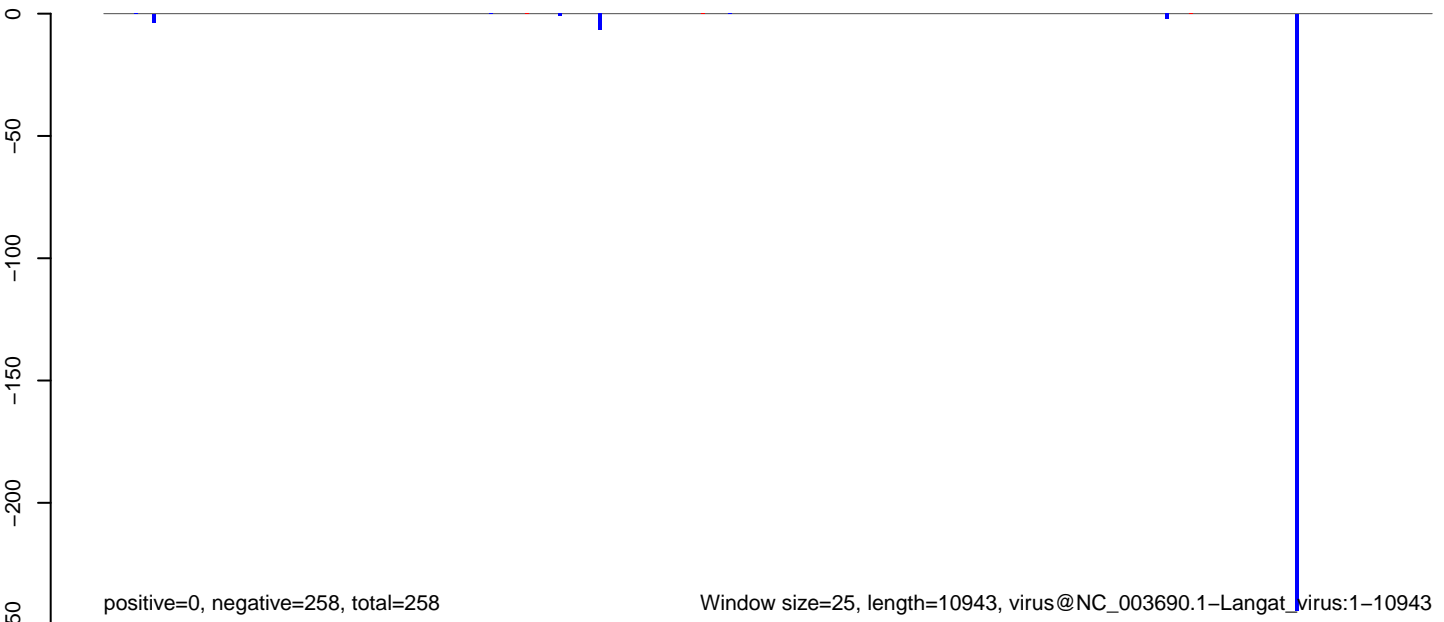
AeAlbo_C636_2dpi_mock3.18_23.rep



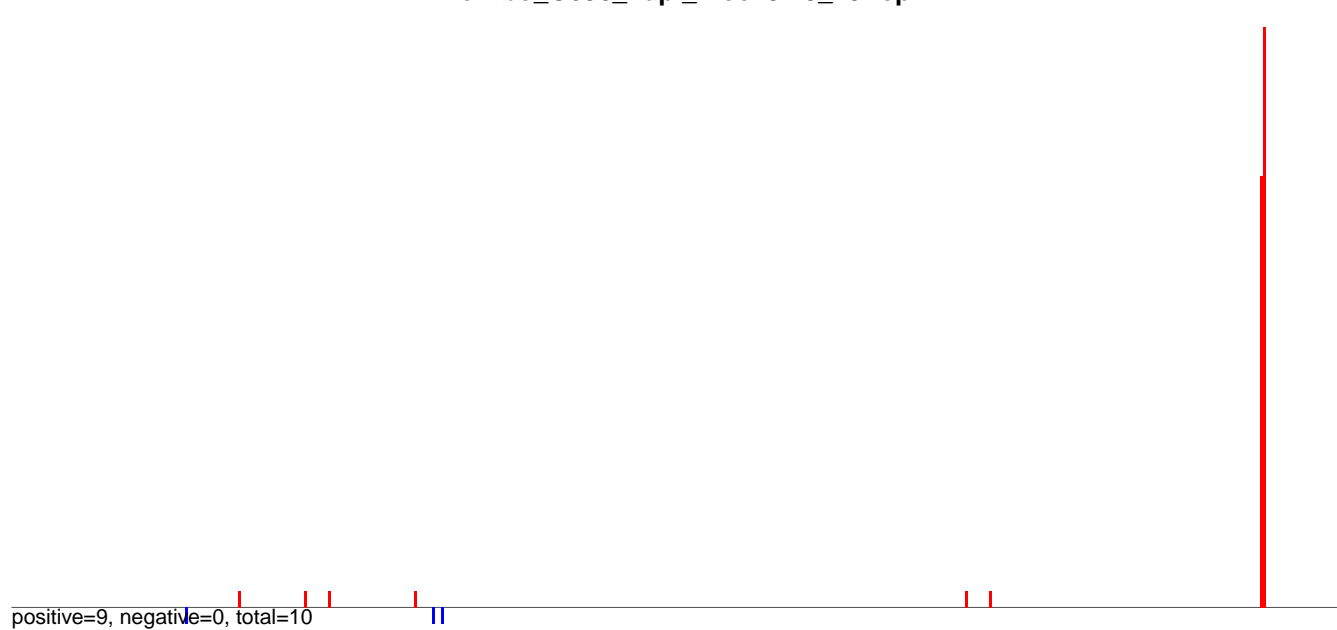
AeAlbo_C636_2dpi_mock3.24_35.rep



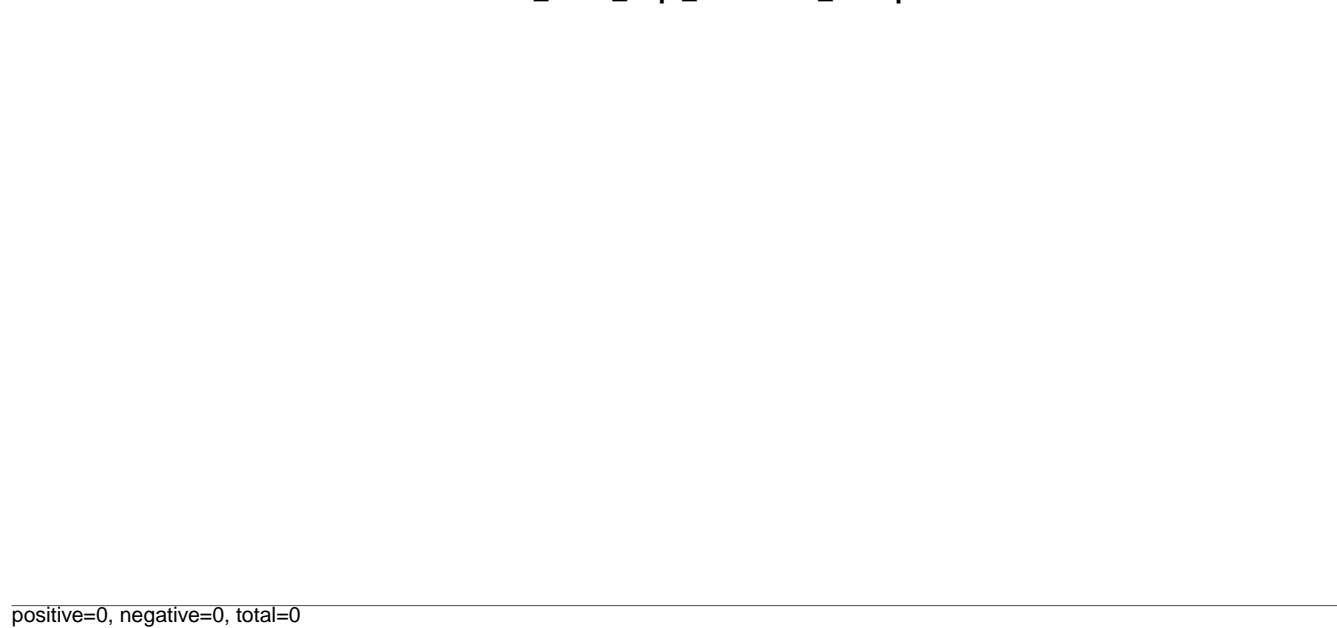
AeAlbo_C636_2dpi_mock3.rep



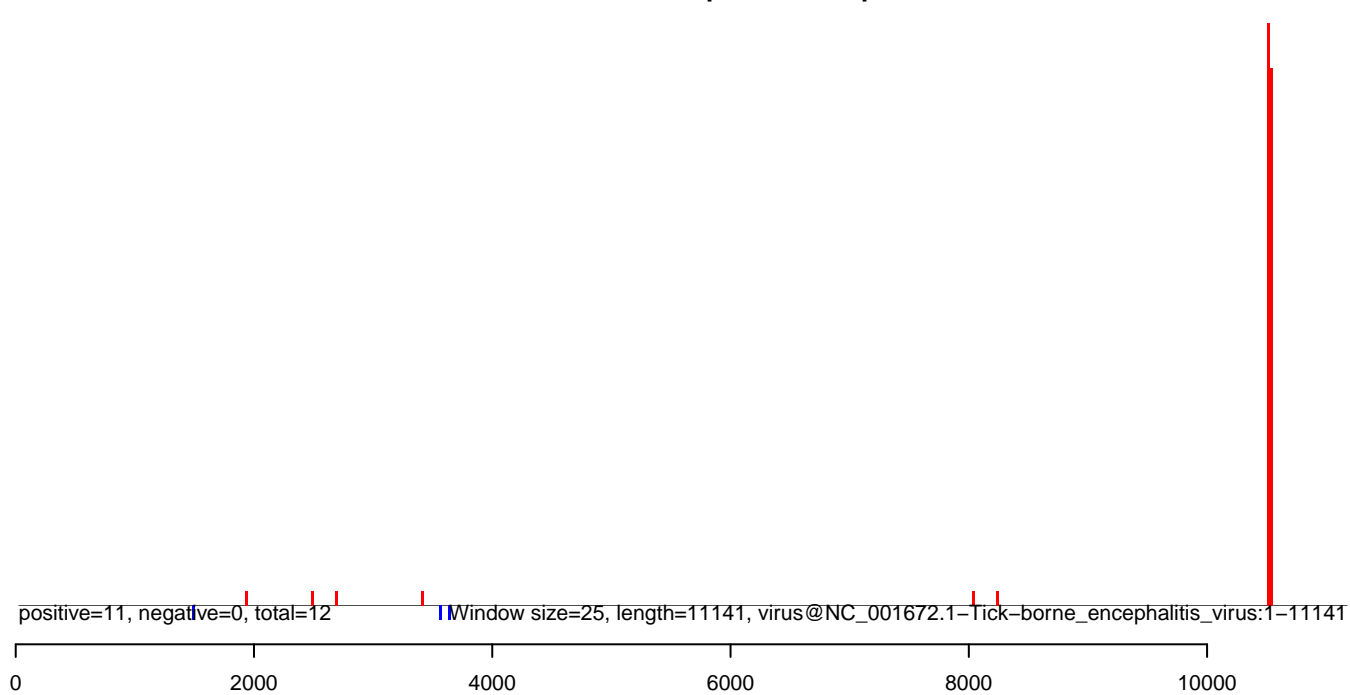
AeAlbo_C636_2dpi_mock3.18_23.rep



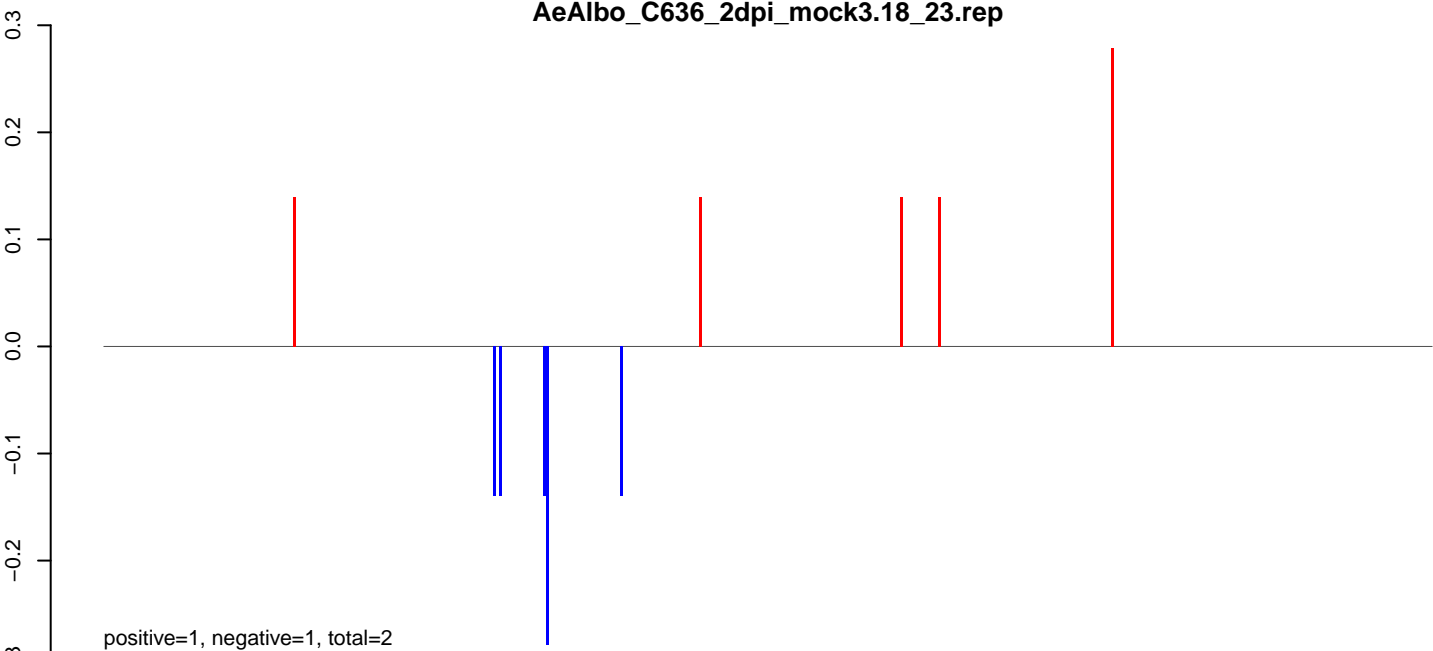
AeAlbo_C636_2dpi_mock3.24_35.rep



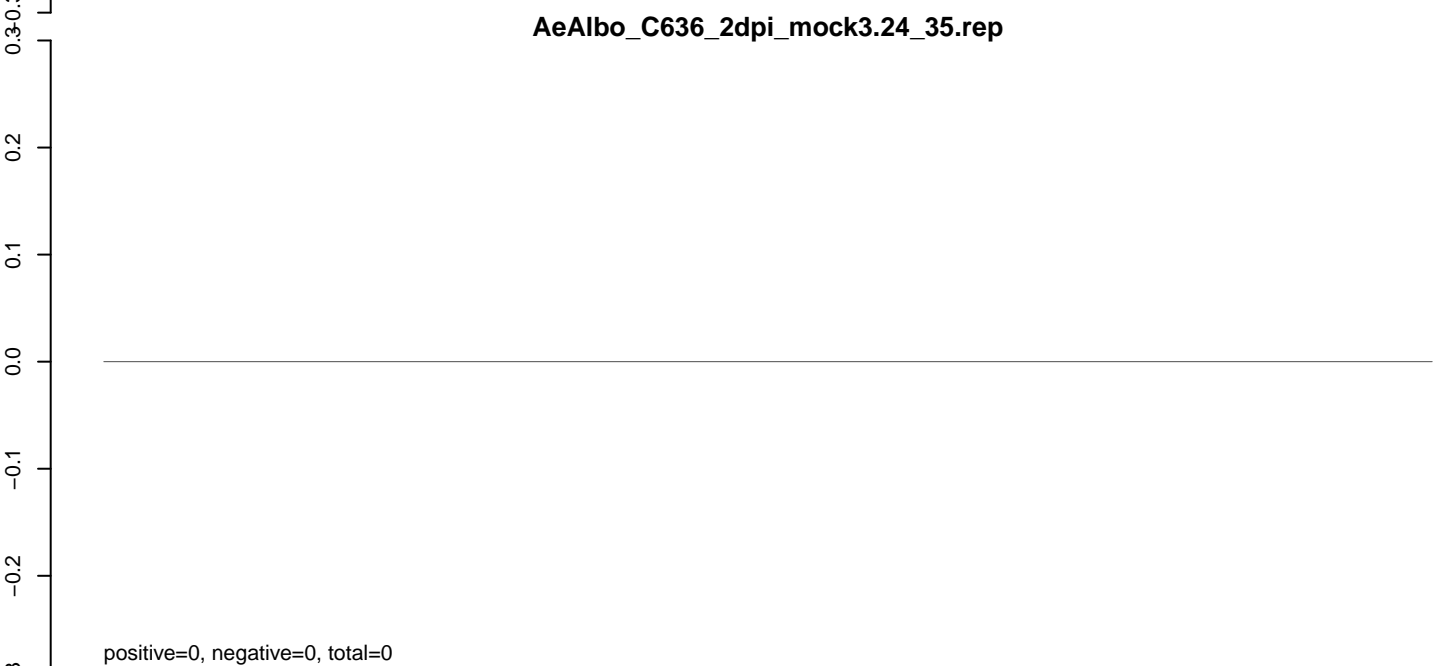
AeAlbo_C636_2dpi_mock3.rep



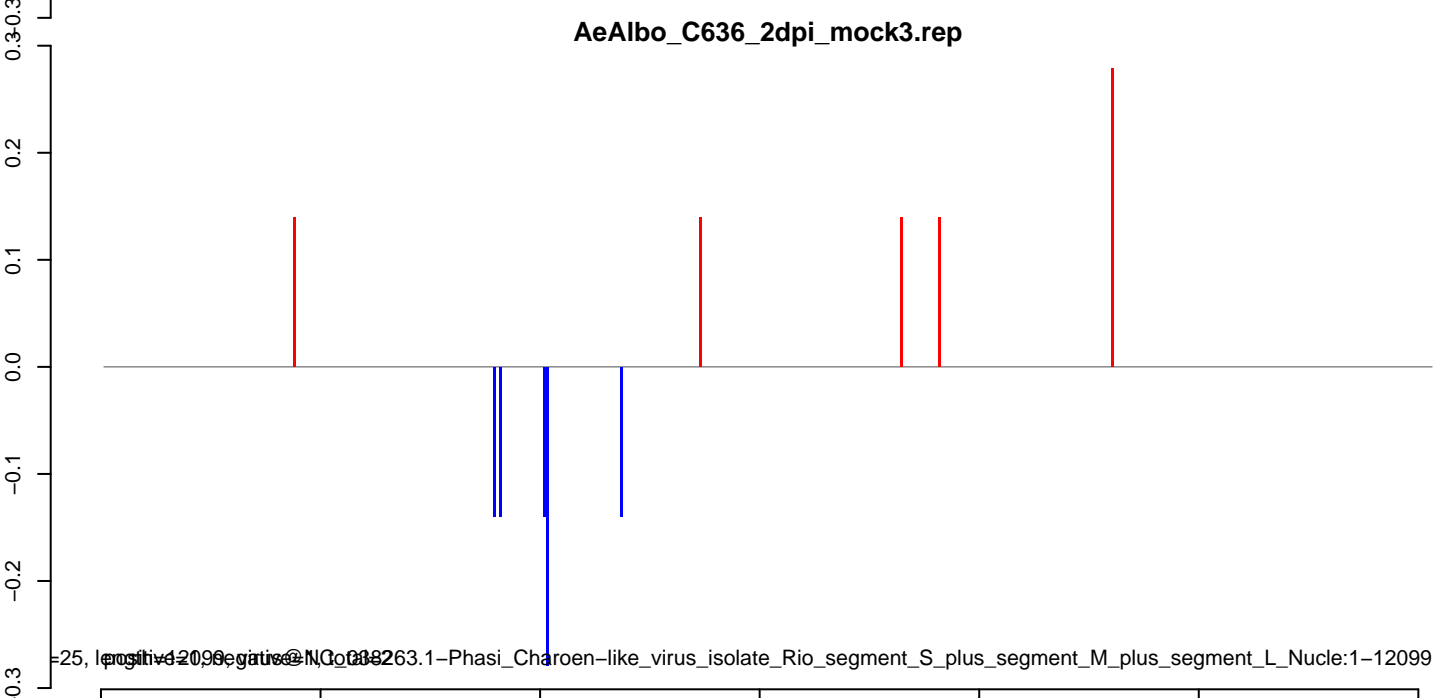
AeAlbo_C636_2dpi_mock3.18_23.rep



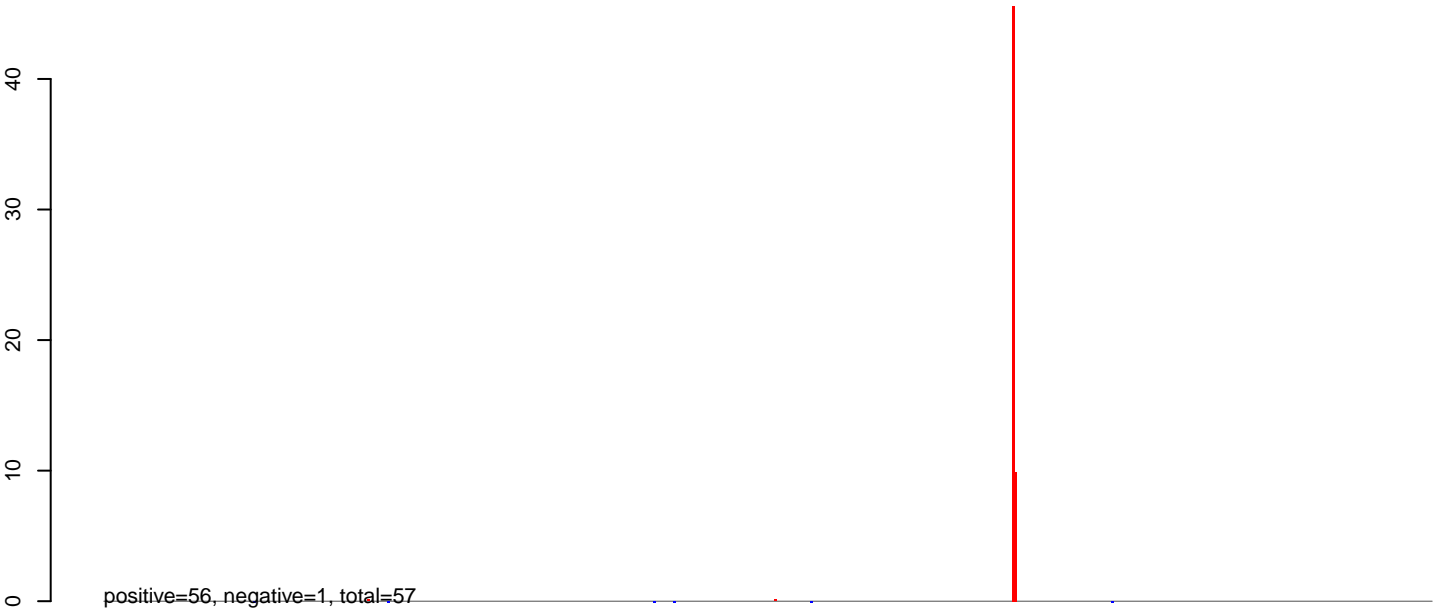
AeAlbo_C636_2dpi_mock3.24_35.rep



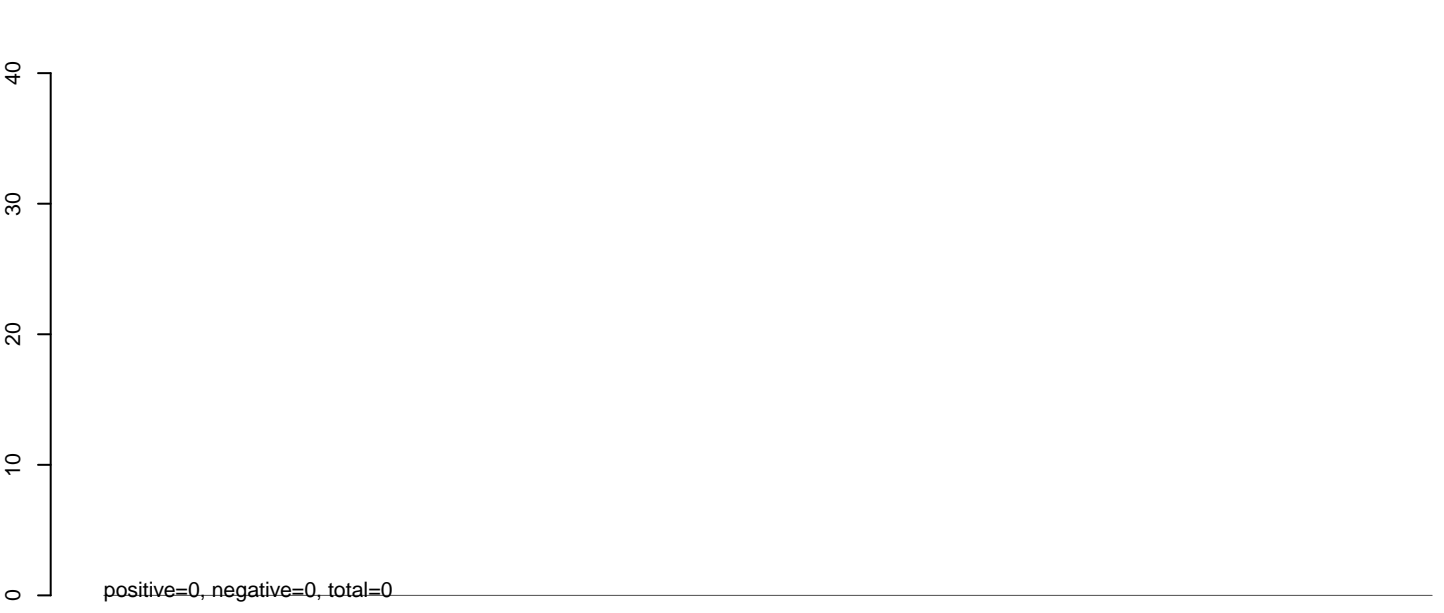
AeAlbo_C636_2dpi_mock3.rep



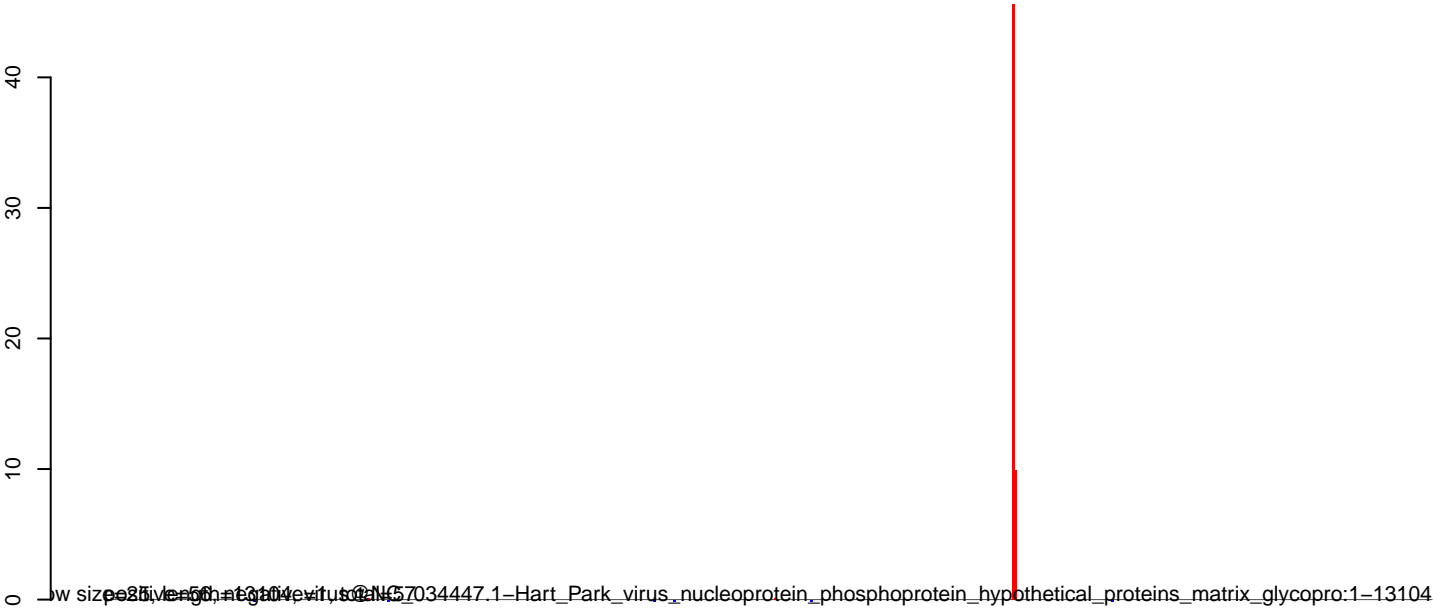
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



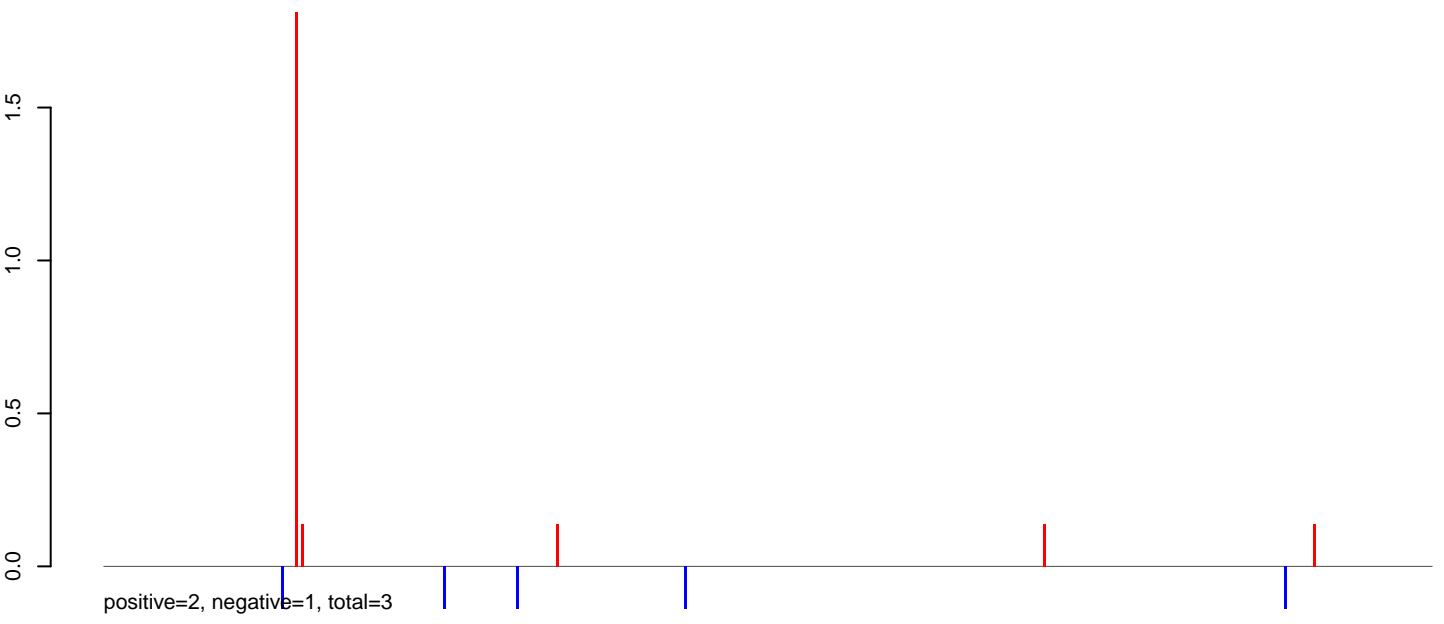
AeAlbo_C636_2dpi_mock3.rep



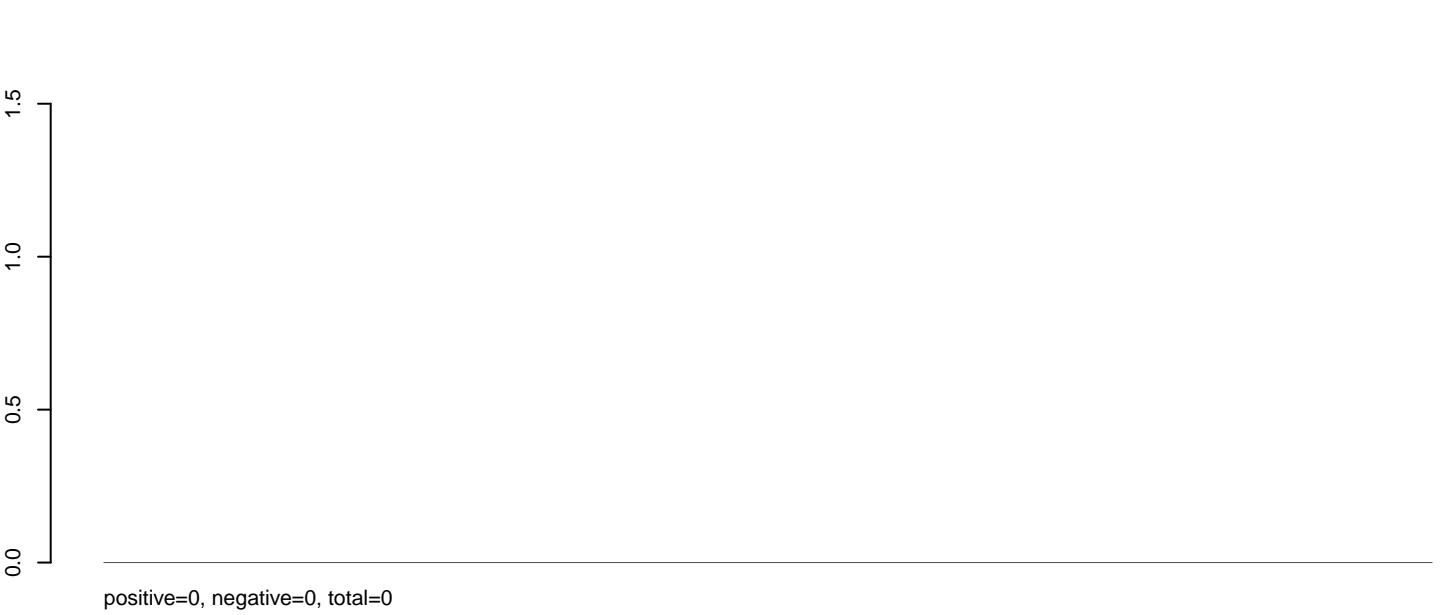
low size positive=56, negative=1, total=57
57034447.1-Hart_Park_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopro:1-13104

0 2000 4000 6000 8000 10000 12000

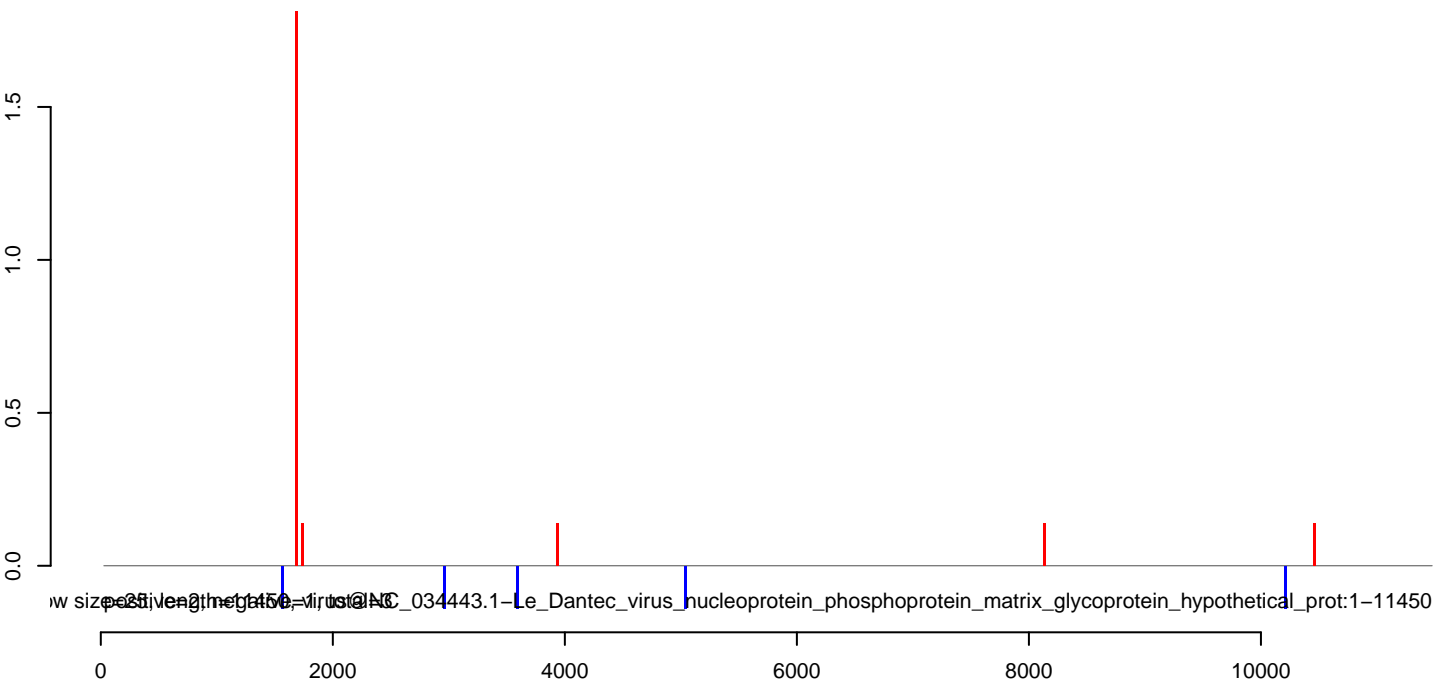
AeAlbo_C636_2dpi_mock3.18_23.rep



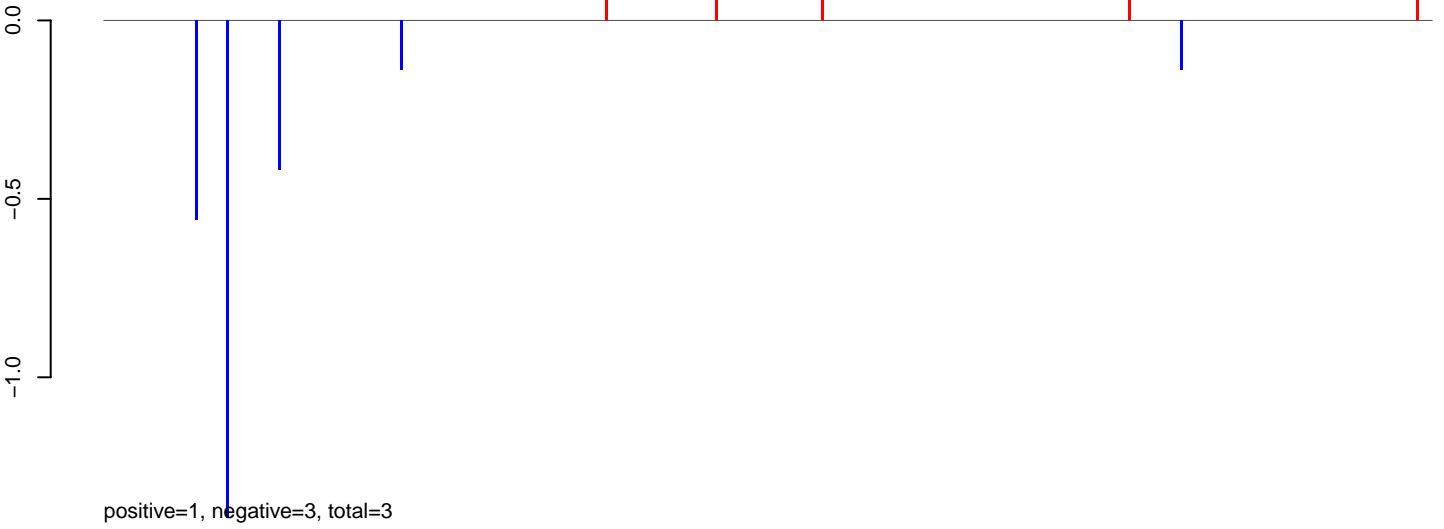
AeAlbo_C636_2dpi_mock3.24_35.rep



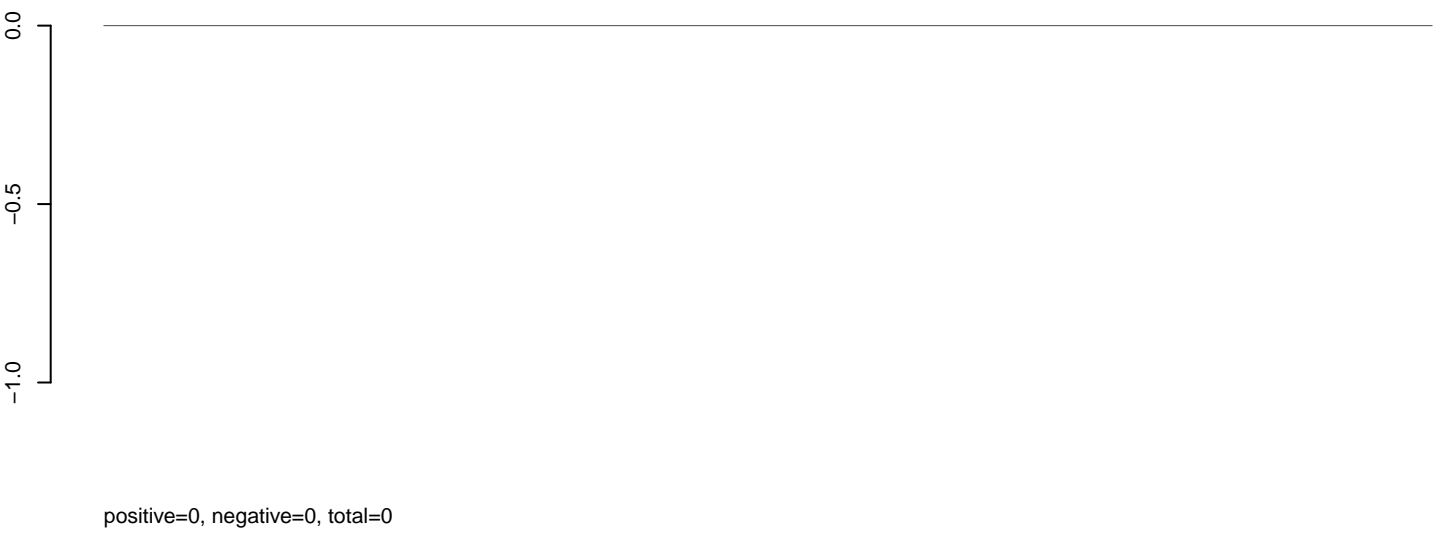
AeAlbo_C636_2dpi_mock3.rep



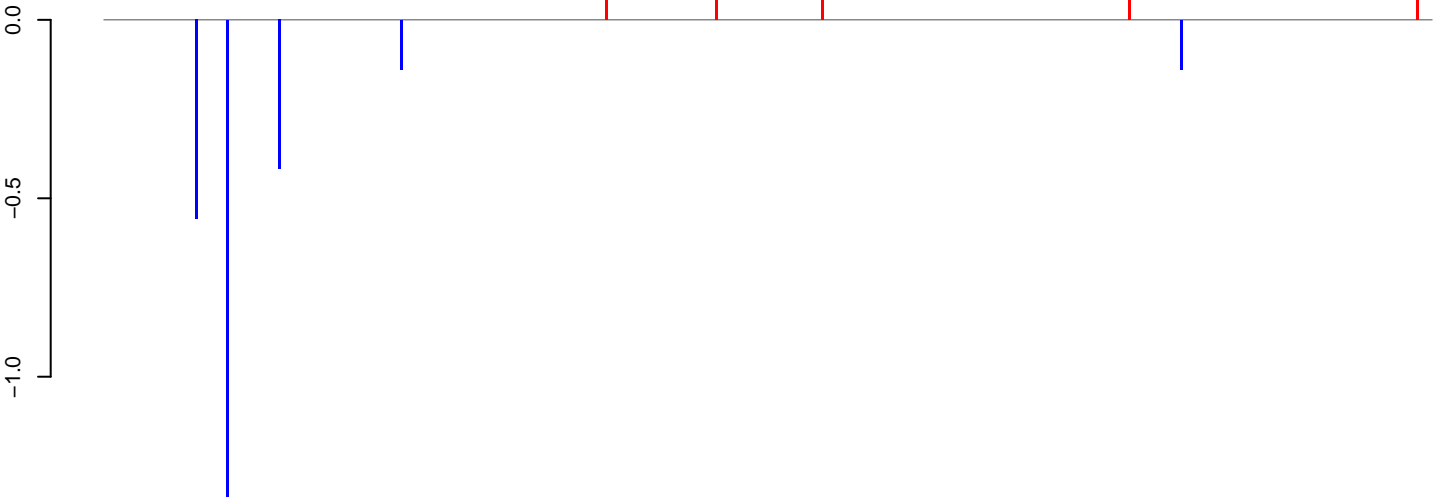
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



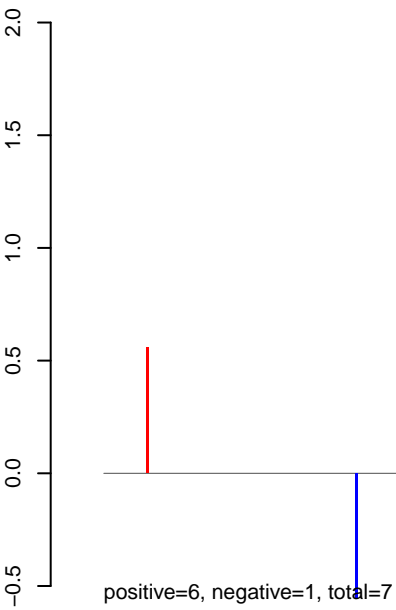
AeAlbo_C636_2dpi_mock3.rep



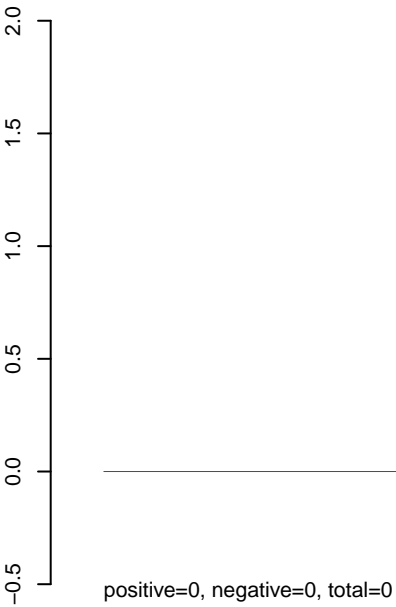
length=12774, version=1, source=NC_033241.1, Xincheng_Mosquito_Virus_strain_XC1-6_ORF1_ORF1_ORF2_ORF2_glycoprotein_G_and_RNA-de:1-12774

0 2000 4000 6000 8000 10000 12000

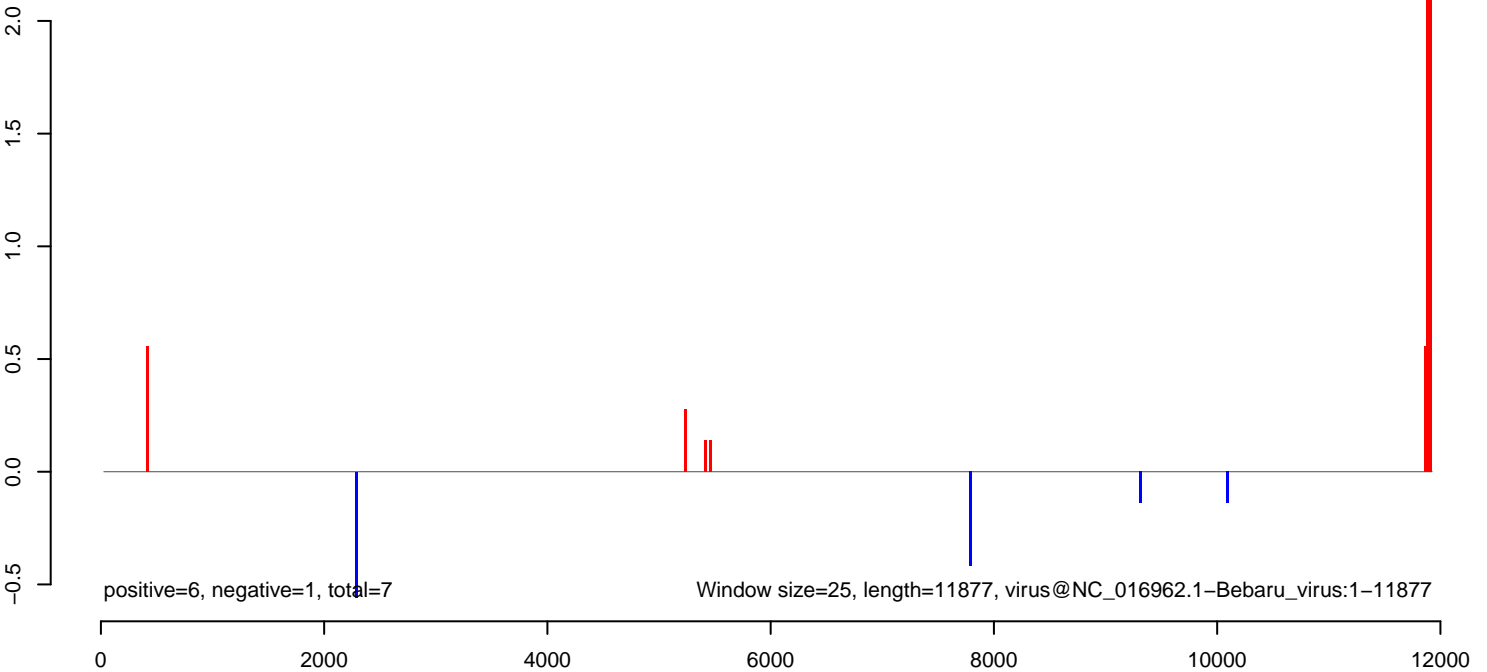
AeAlbo_C636_2dpi_mock3.18_23.rep



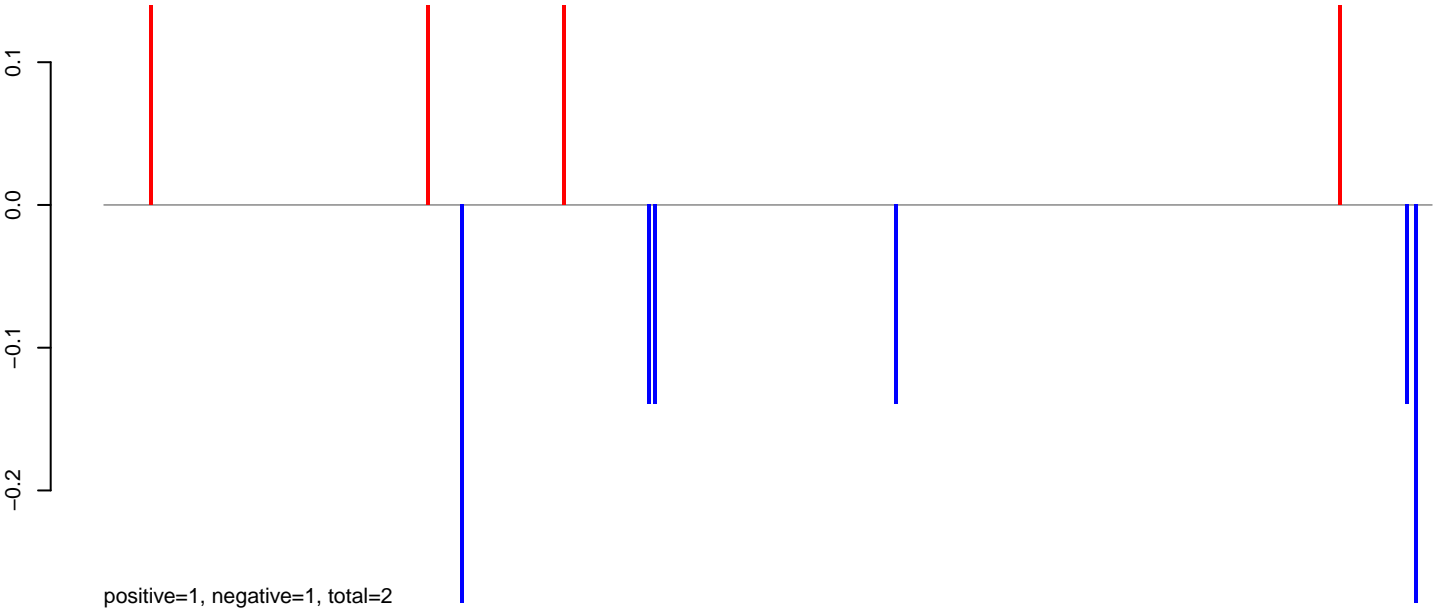
AeAlbo_C636_2dpi_mock3.24_35.rep



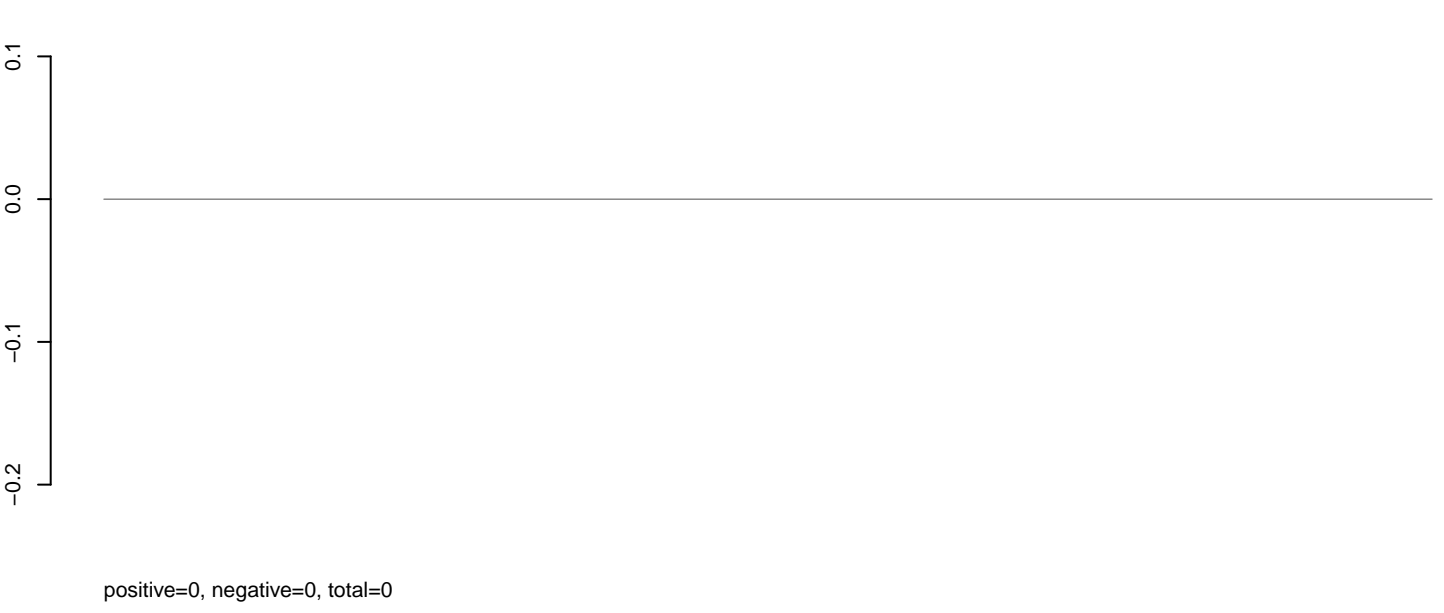
AeAlbo_C636_2dpi_mock3.rep



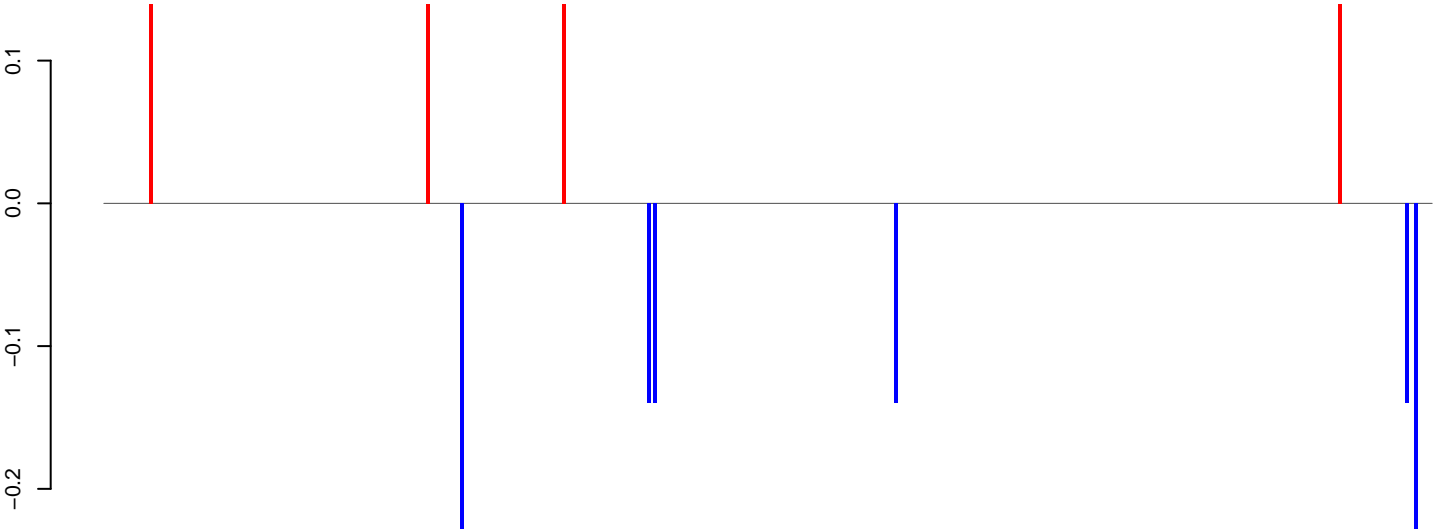
AeAlbo_C636_2dpi_mock3.18_23.rep



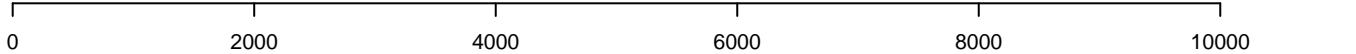
AeAlbo_C636_2dpi_mock3.24_35.rep



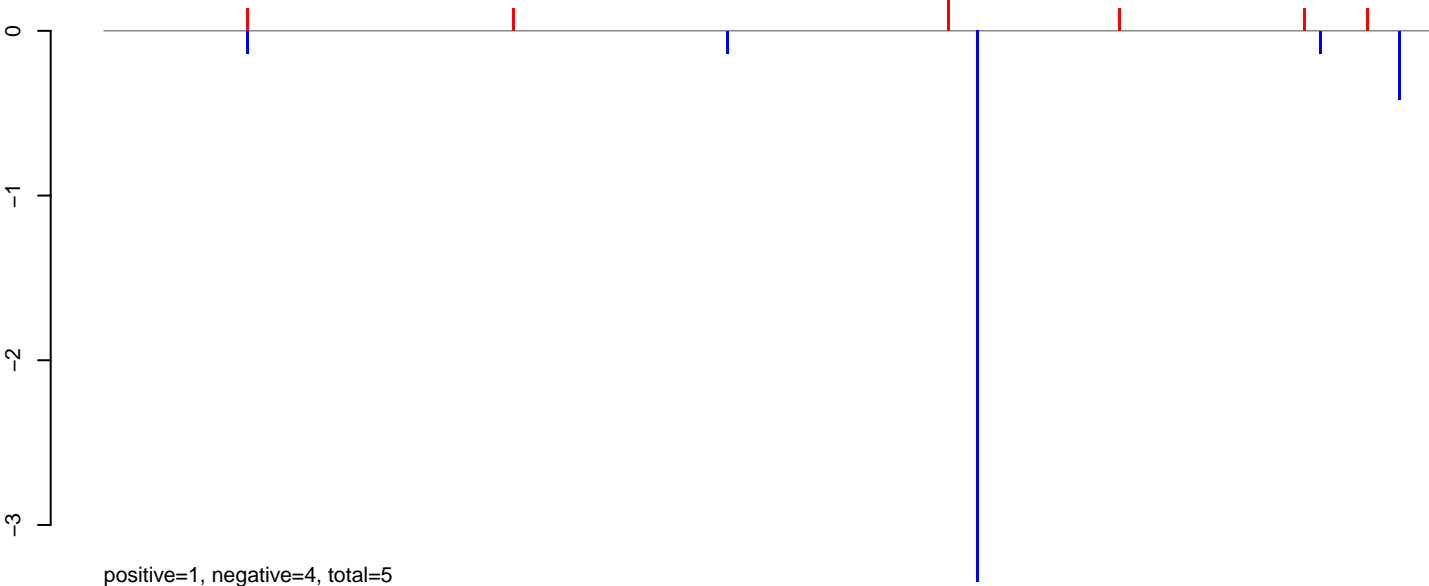
AeAlbo_C636_2dpi_mock3.rep



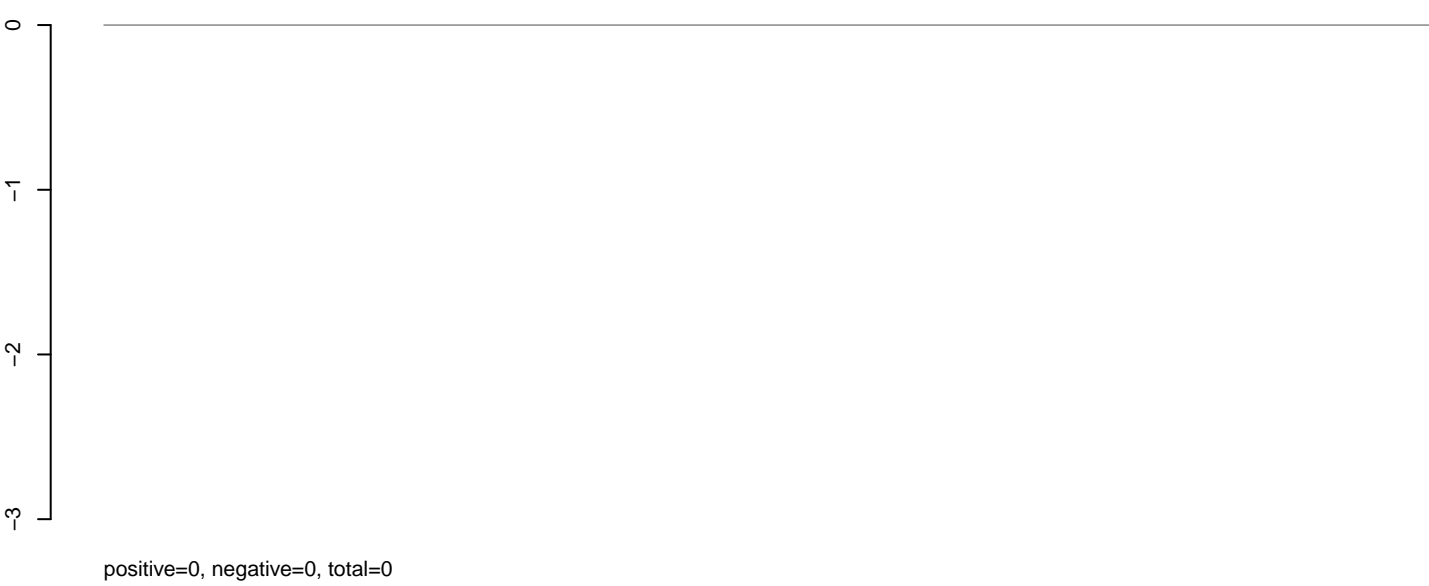
Windows size=25, length=20990, virus@NC_015843.2-Tembusu_virus_flavivirus_polyprotein_flavivirus_polyprotein_gene_gene:1-10990



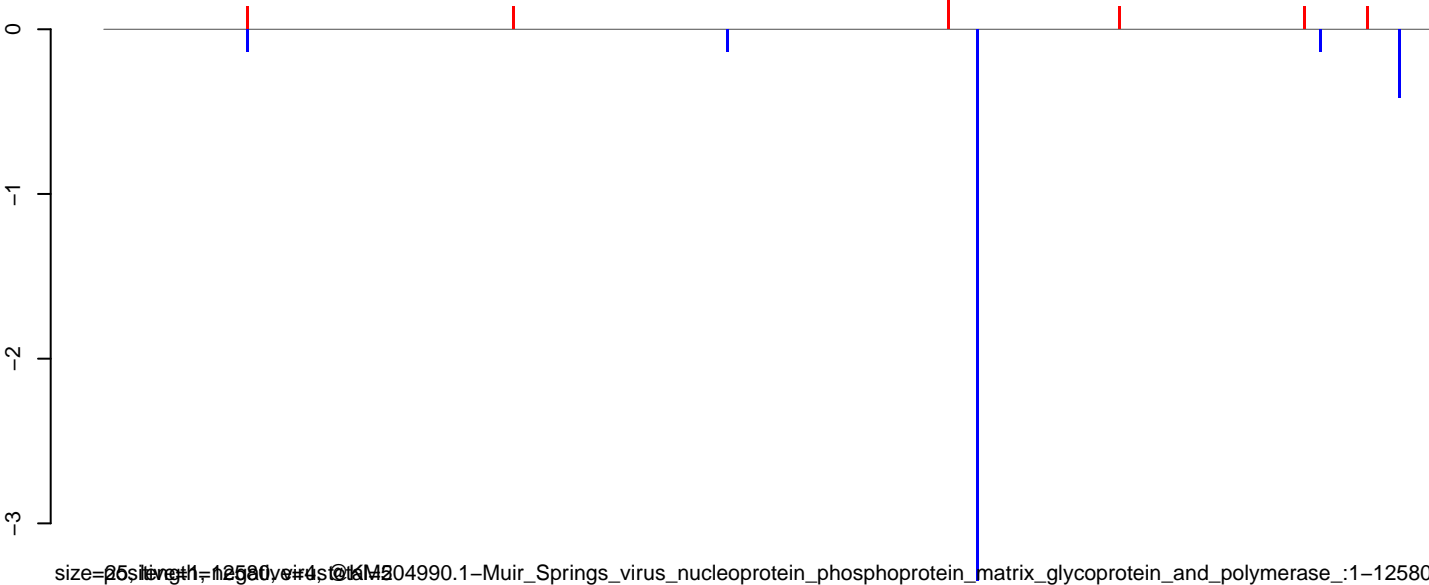
AeAlbo_C636_2dpi_mock3.18_23.rep



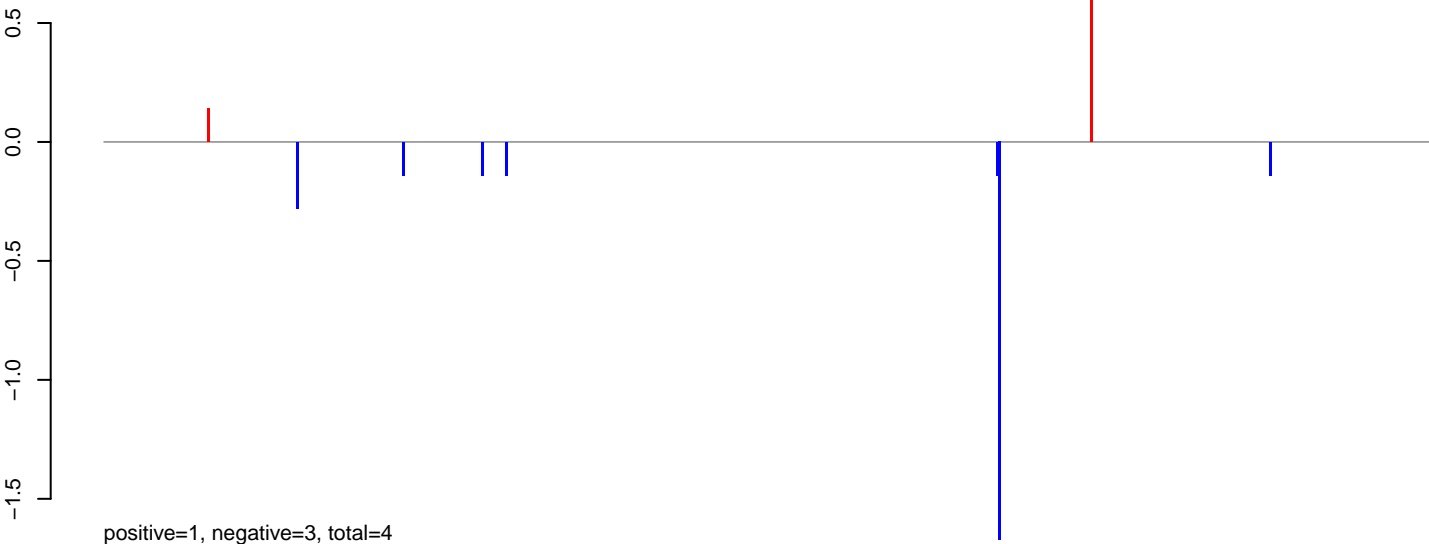
AeAlbo_C636_2dpi_mock3.24_35.rep



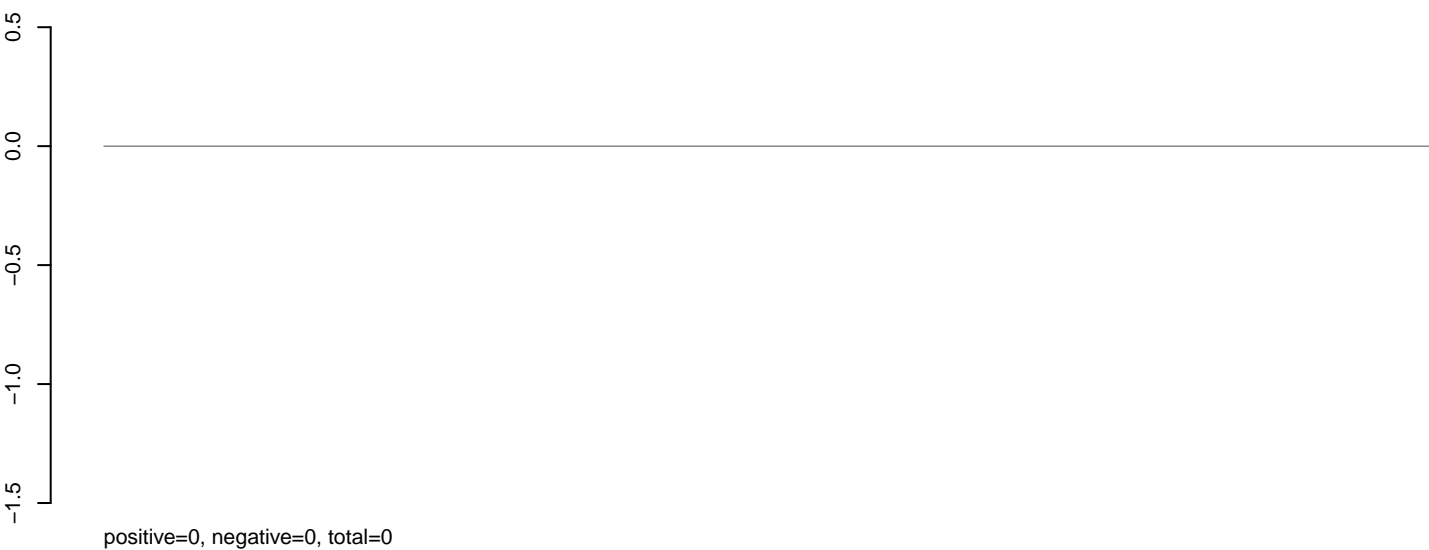
AeAlbo_C636_2dpi_mock3.rep



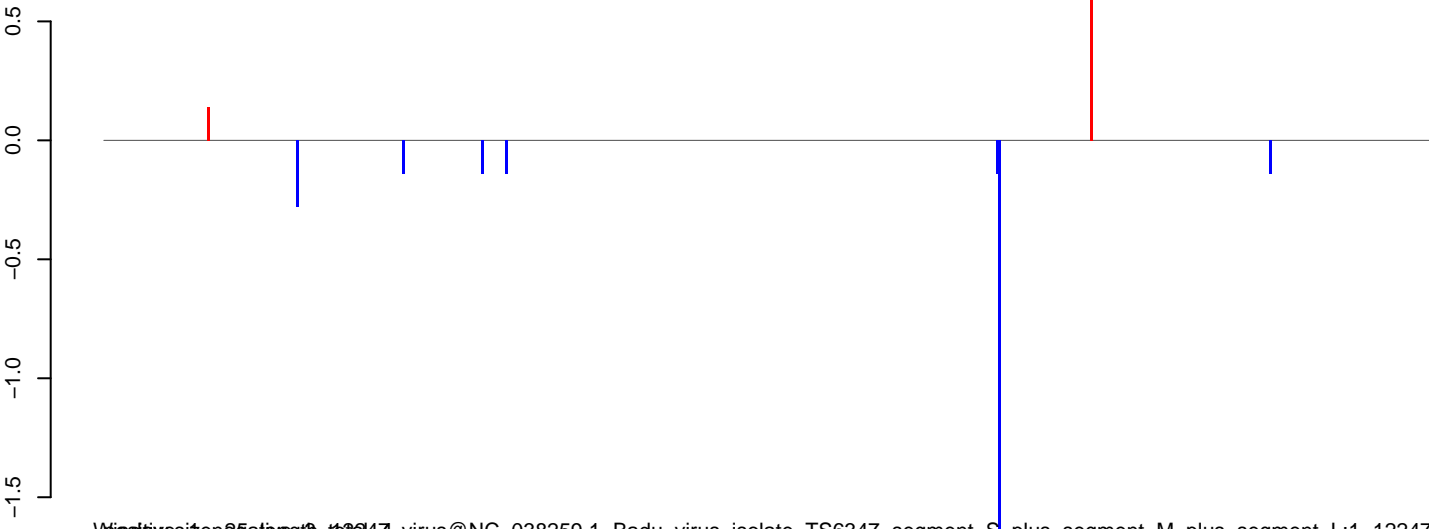
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



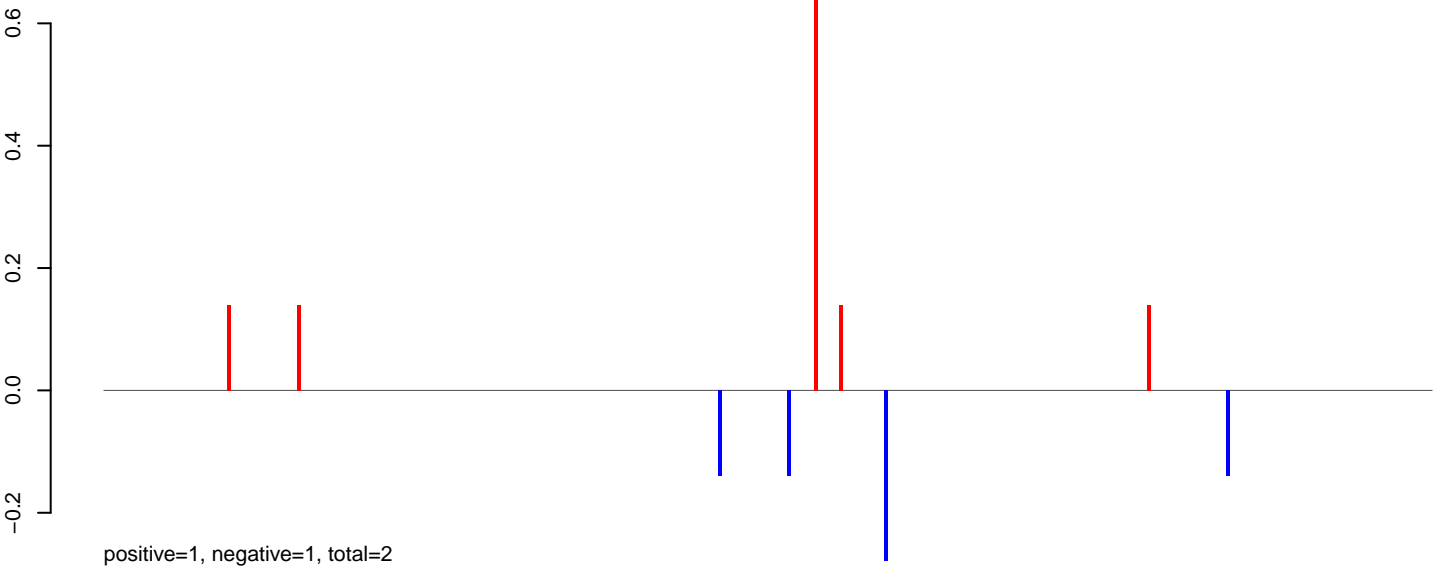
AeAlbo_C636_2dpi_mock3.rep



positives=2, negatives=4, total=6, virus@NC_038259.1-Badu_virus_isolate_TS6347_segment_S_plus_segment_M_plus_segment_L:1-12247

0 2000 4000 6000 8000 10000 12000

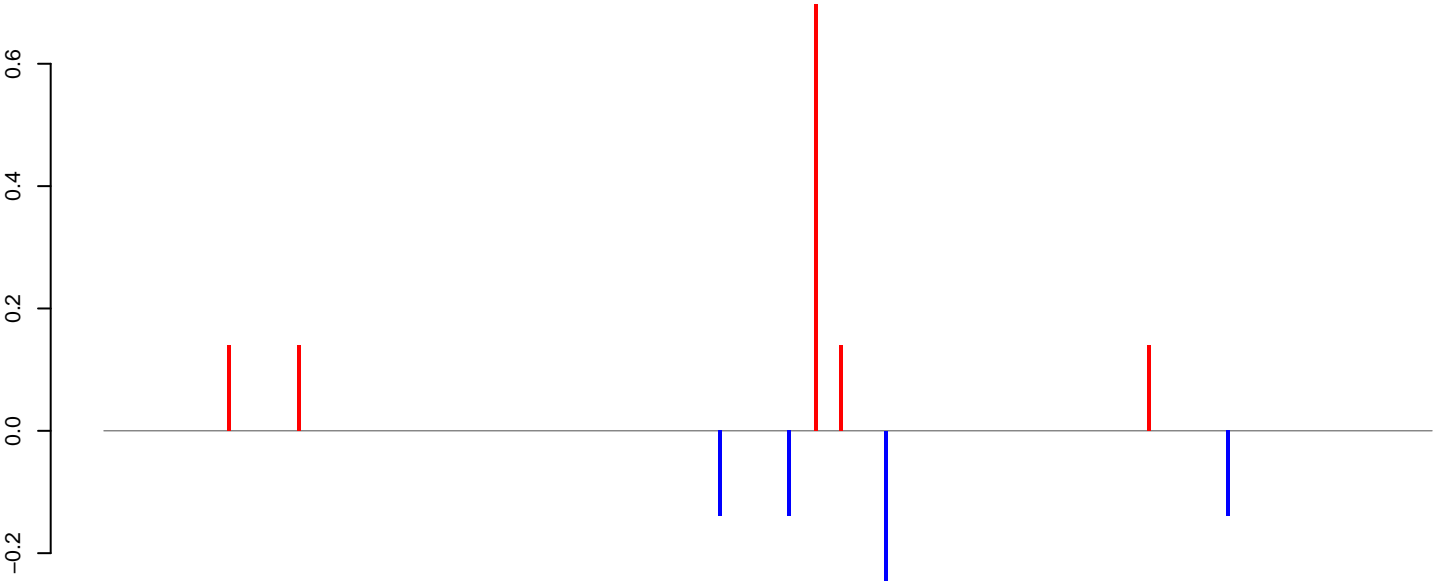
AeAlbo_C636_2dpi_mock3.18_23.rep



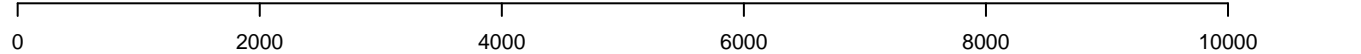
AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



size=25, strength=10000, virus=0, file=C:\2034539.1-Nkolbisson_virus_nucleoprotein_phosphoprotein_matrix_glycoprotein_and_polymerase_g:1-10968



AeAlbo_C636_2dpi_mock3.18_23.rep

0.1
0.0
-0.1
-0.2

positive=0, negative=1, total=2

AeAlbo_C636_2dpi_mock3.24_35.rep

0.1
0.0
-0.1
-0.2

positive=0, negative=0, total=0

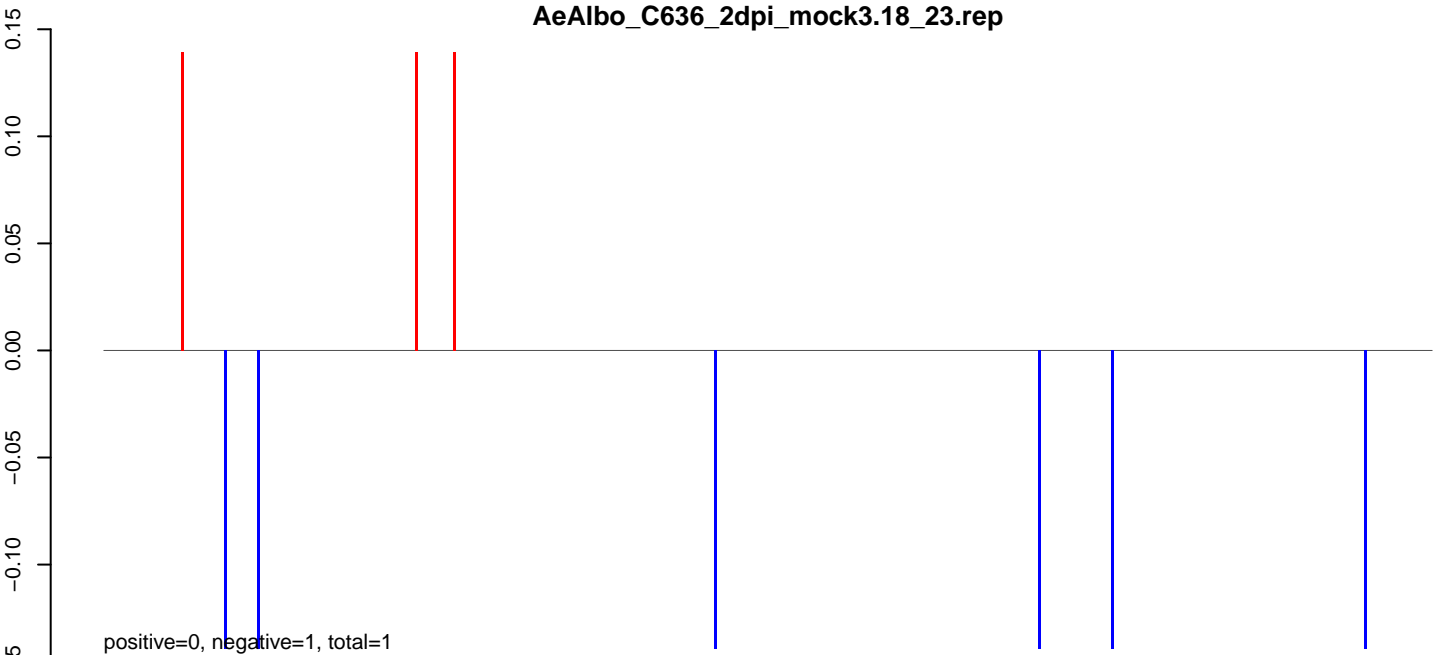
AeAlbo_C636_2dpi_mock3.rep

0.1
0.0
-0.1
-0.2

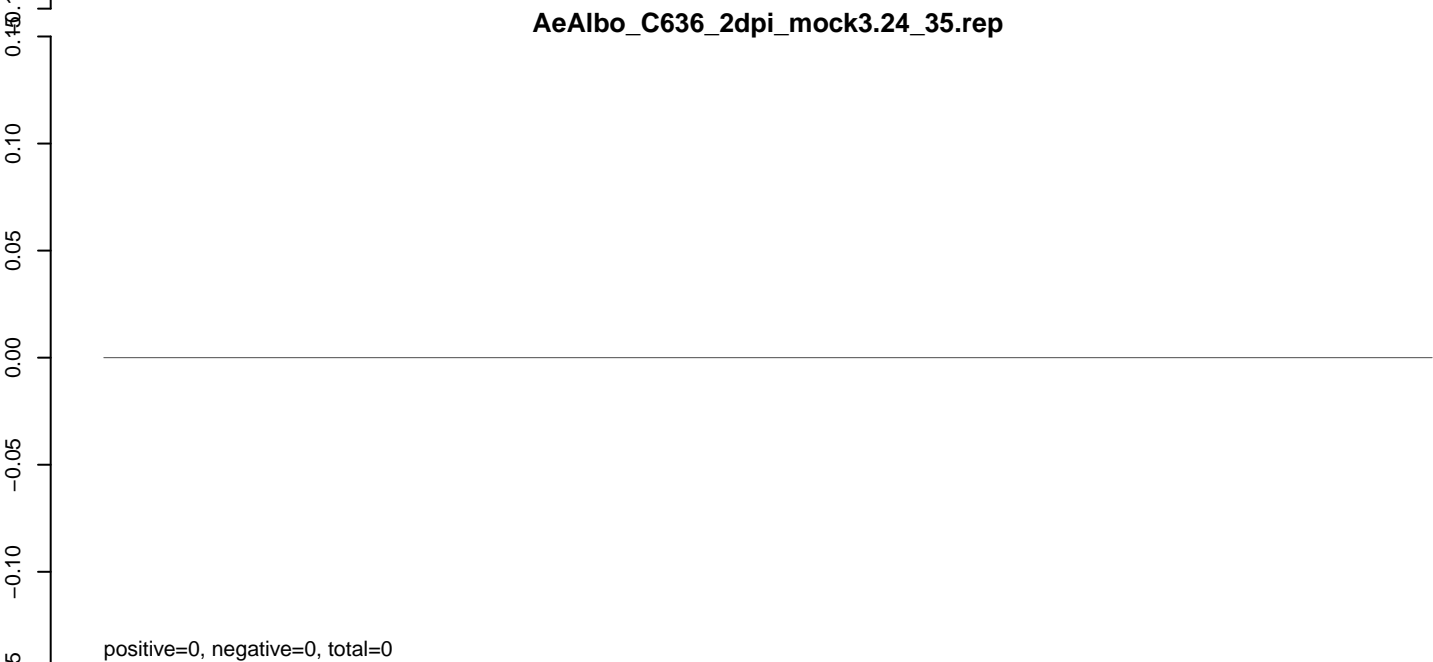
size=25, length=13593, use @NC_02034534.1-Rochambeau_virus_nucleoprotein_phosphoprotein_matrix_hypthetical_proteins_glycopr:1-13593

0 2000 4000 6000 8000 10000 12000 14000

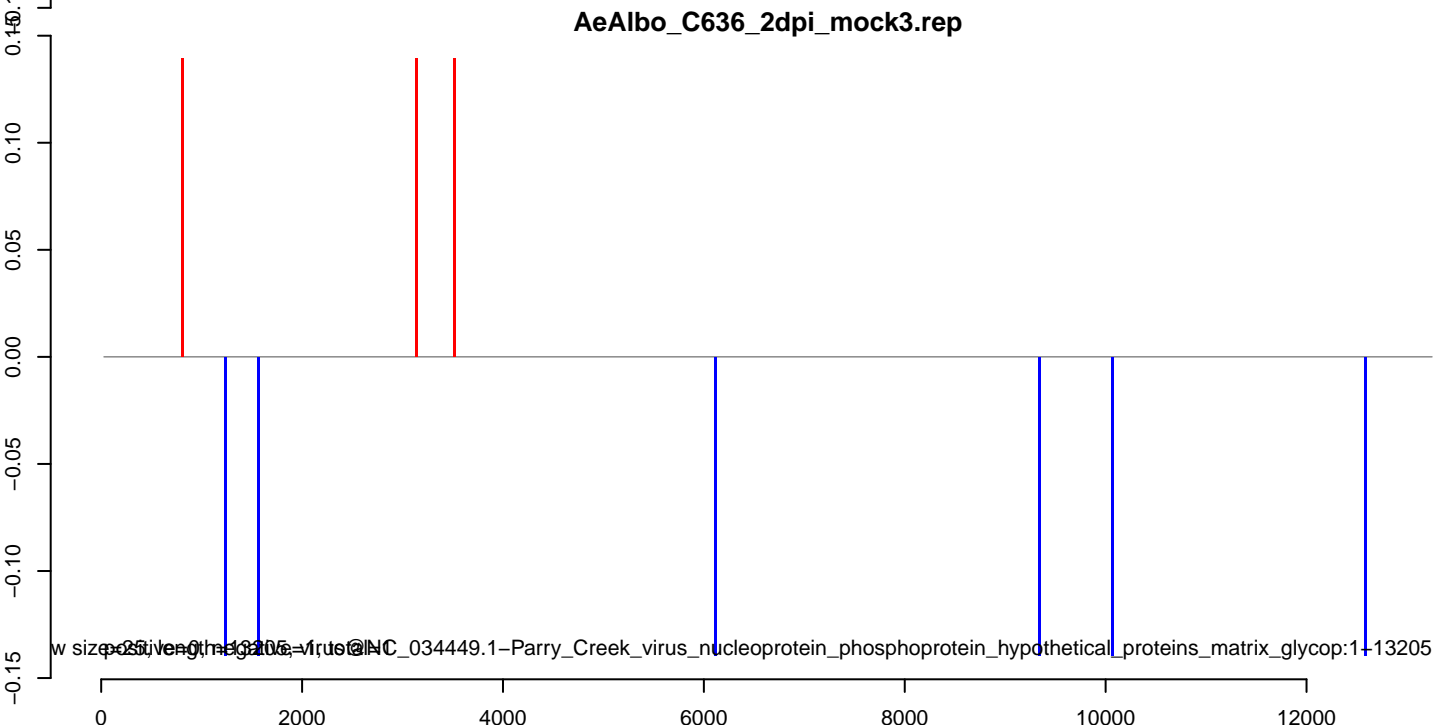
AeAlbo_C636_2dpi_mock3.18_23.rep



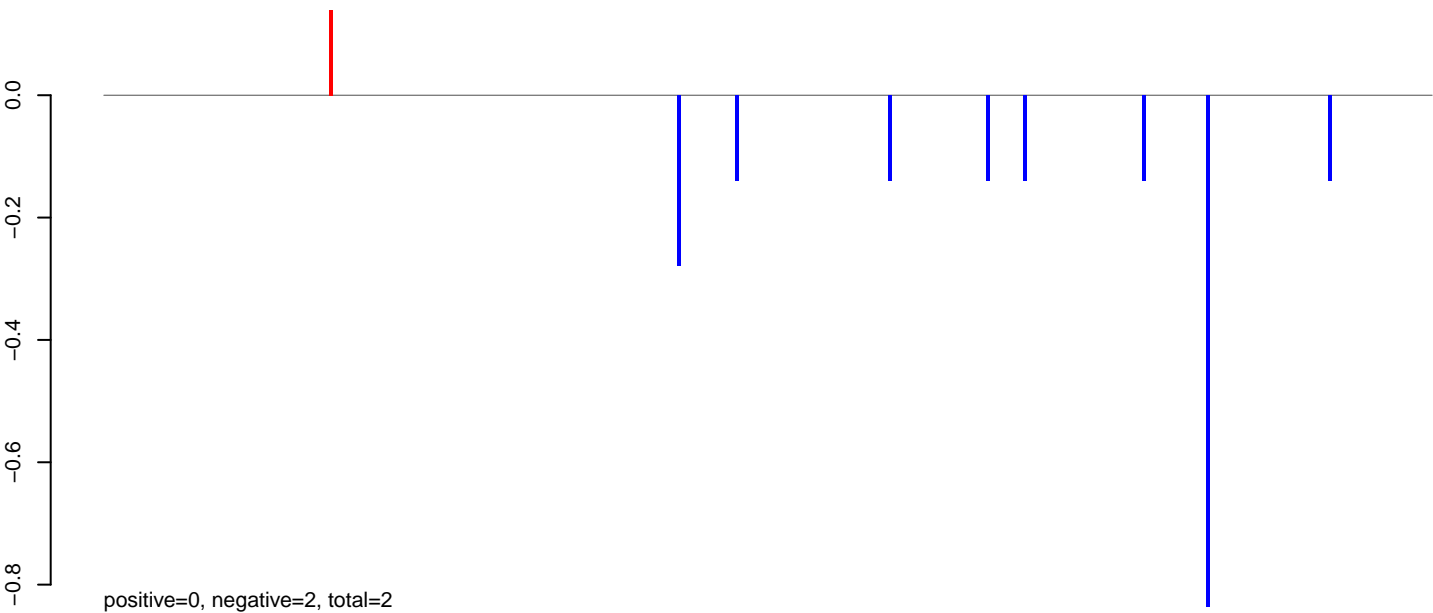
AeAlbo_C636_2dpi_mock3.24_35.rep



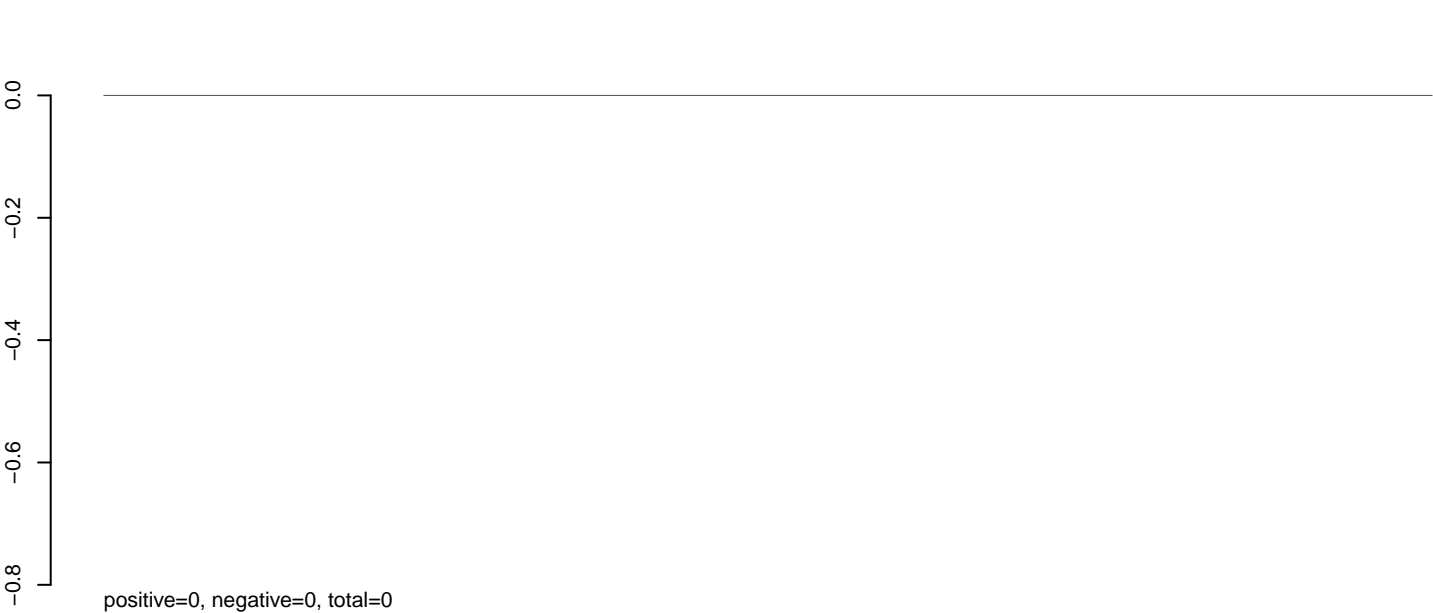
AeAlbo_C636_2dpi_mock3.rep



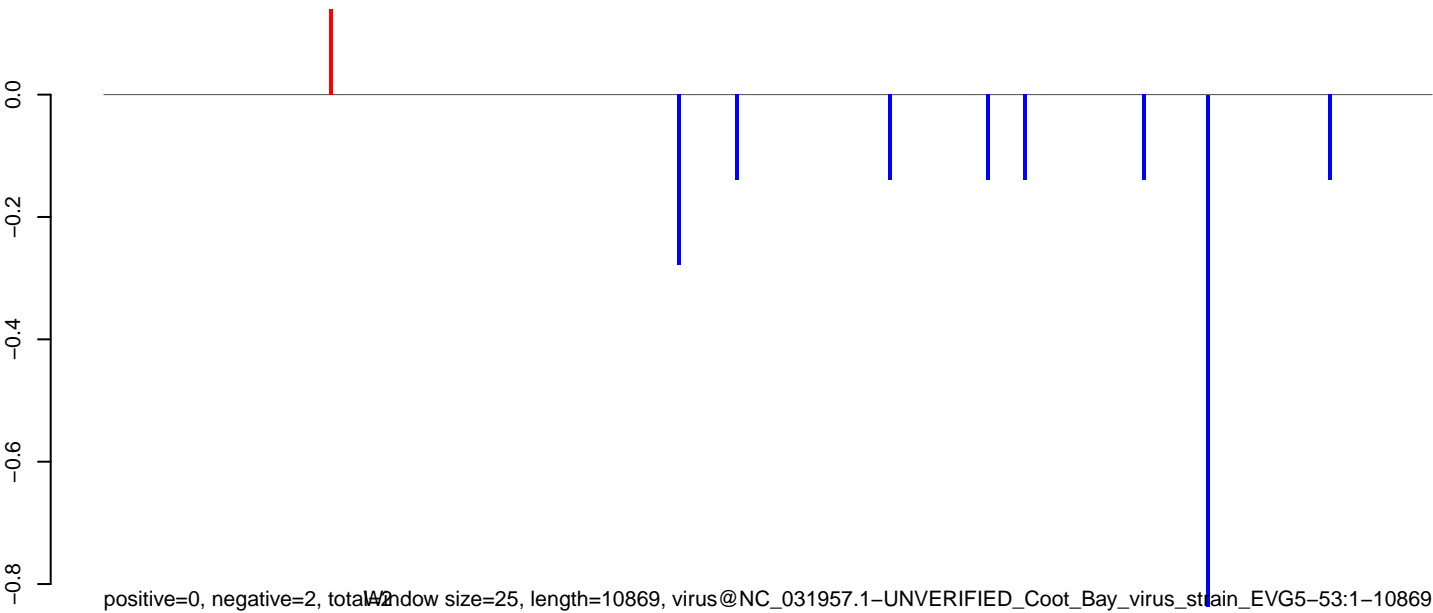
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



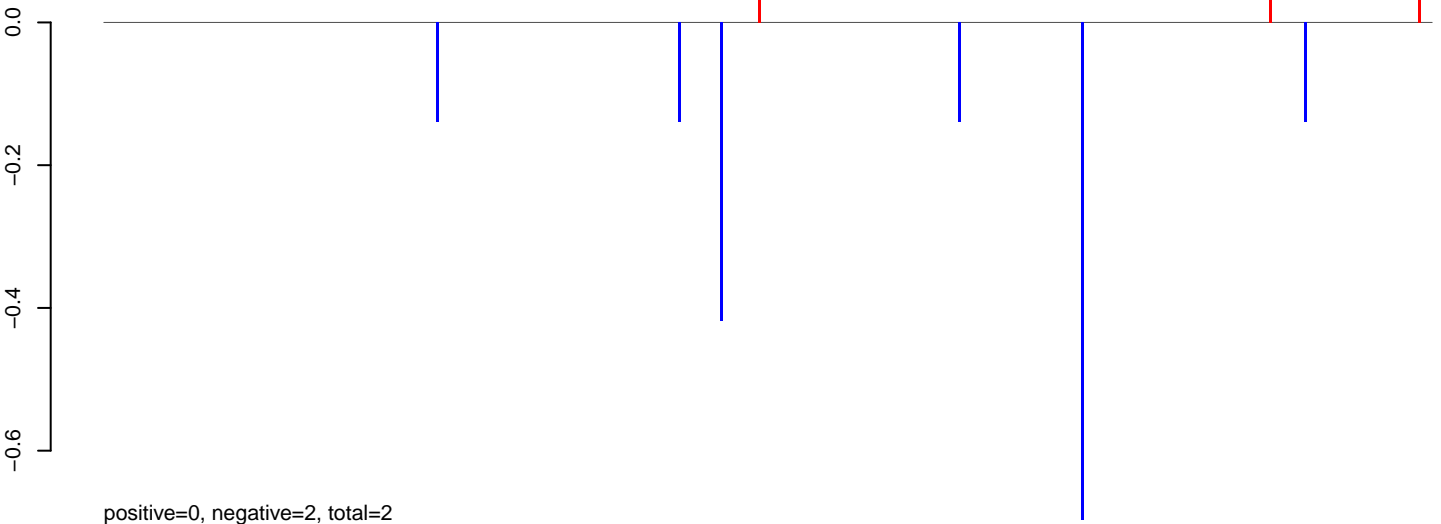
AeAlbo_C636_2dpi_mock3.rep



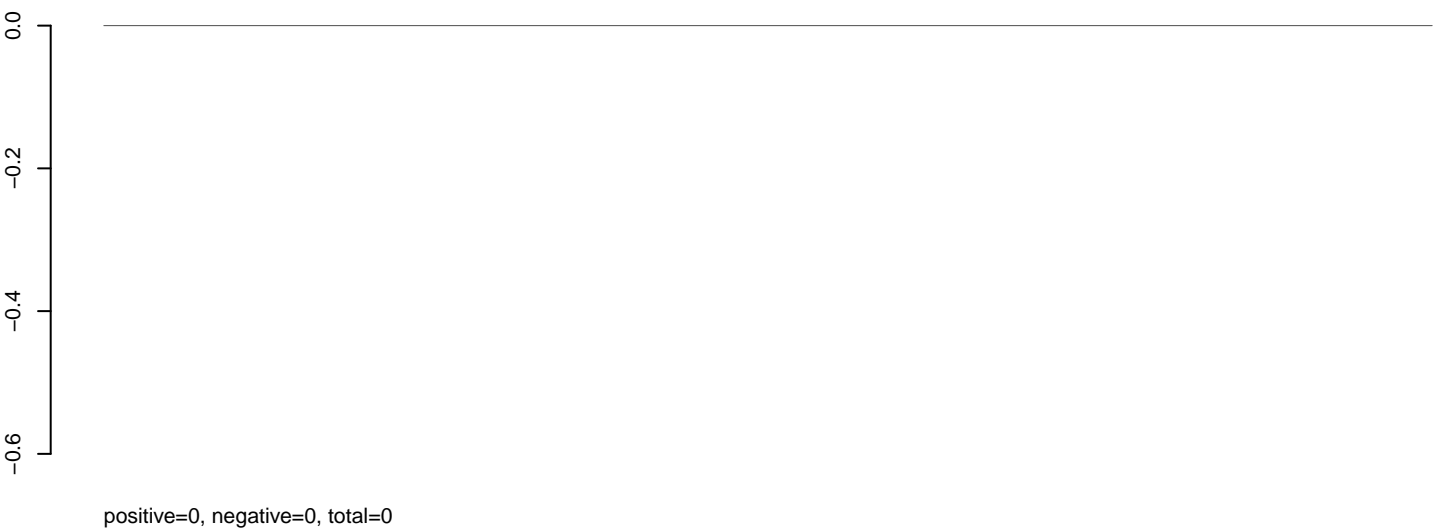
Window size=25, length=10869, virus@NC_031957.1-UNVERIFIED_Coot_Bay_virus_strain_EVG5-53:1-10869

0 2000 4000 6000 8000 10000

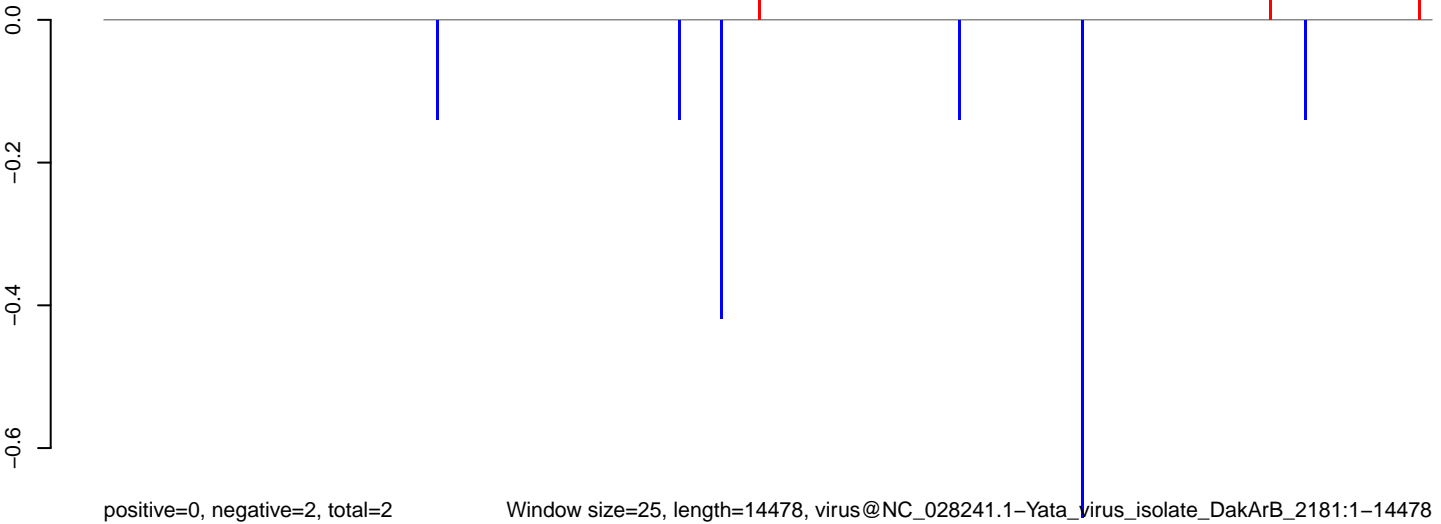
AeAlbo_C636_2dpi_mock3.18_23.rep



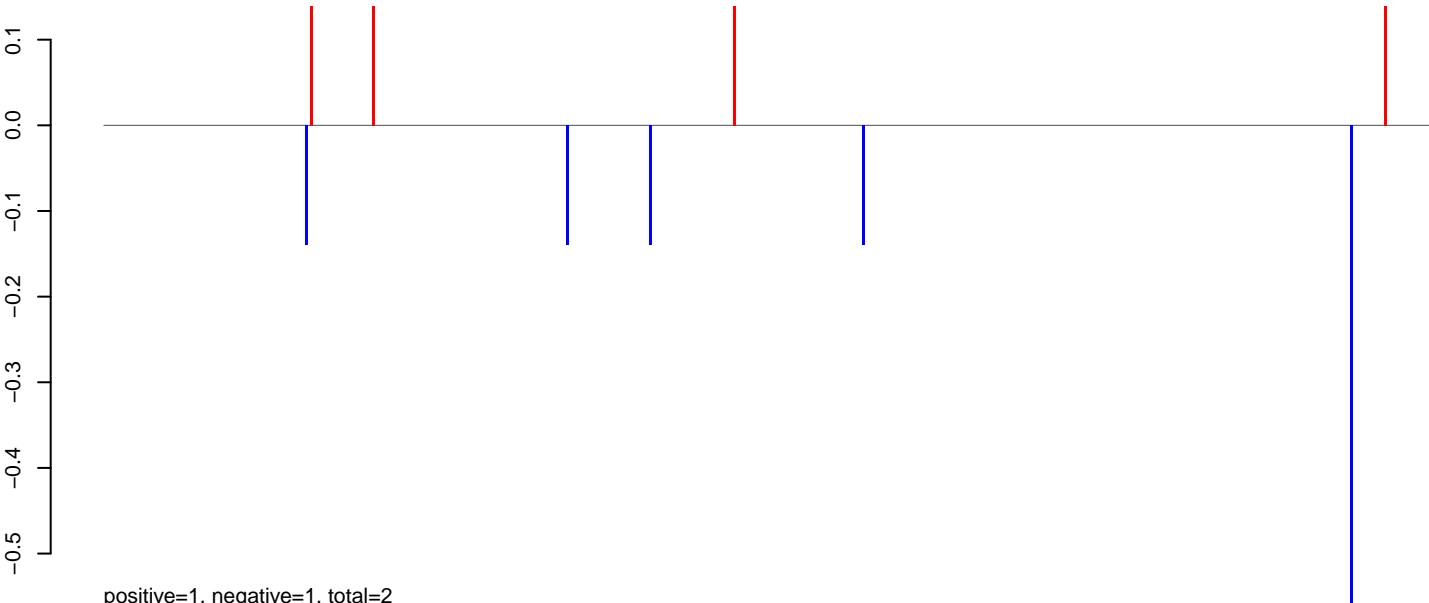
AeAlbo_C636_2dpi_mock3.24_35.rep



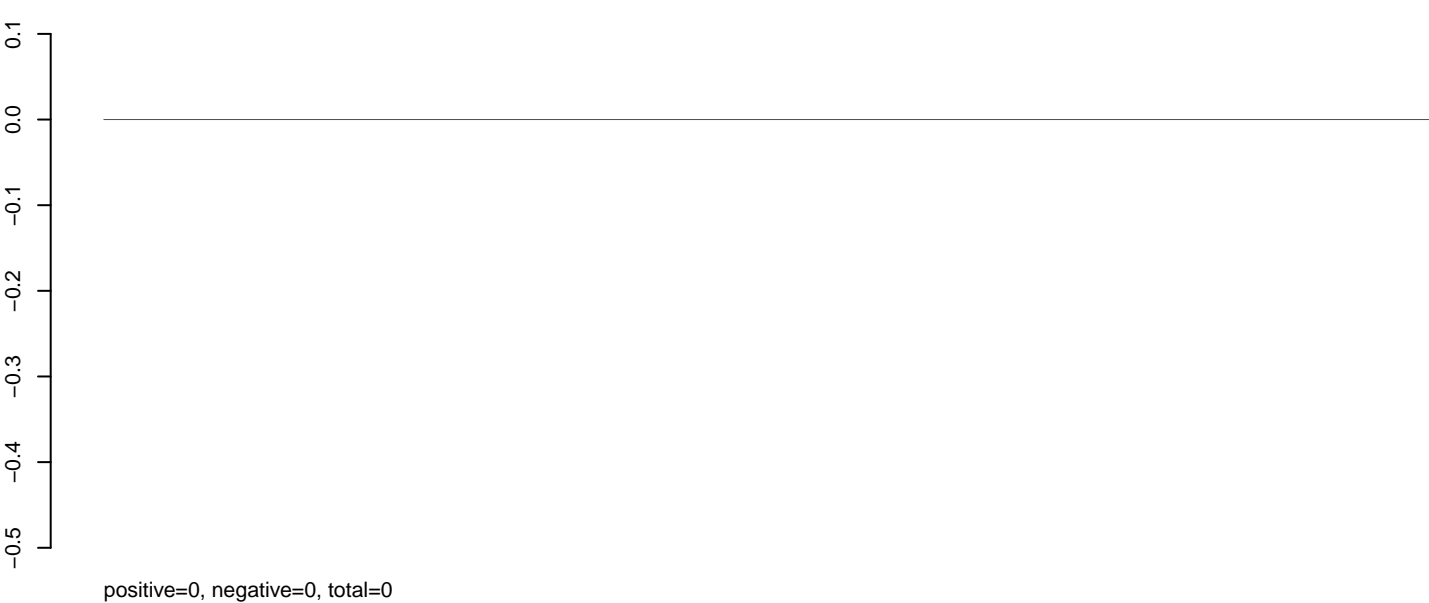
AeAlbo_C636_2dpi_mock3.rep



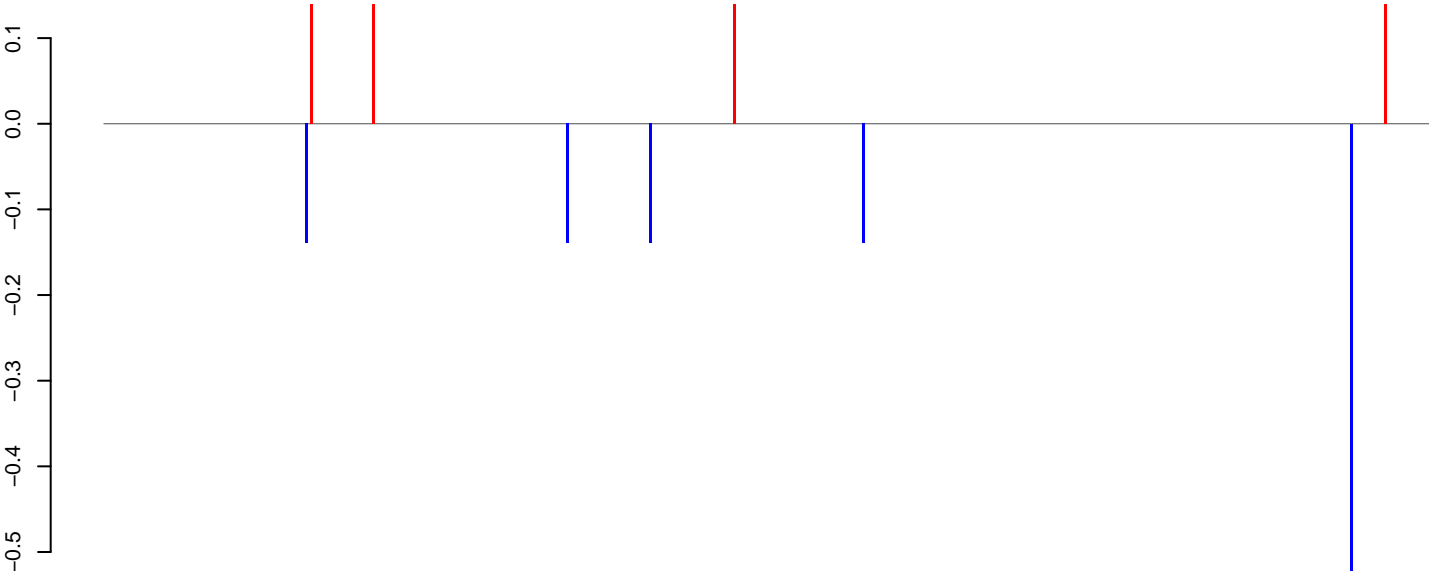
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

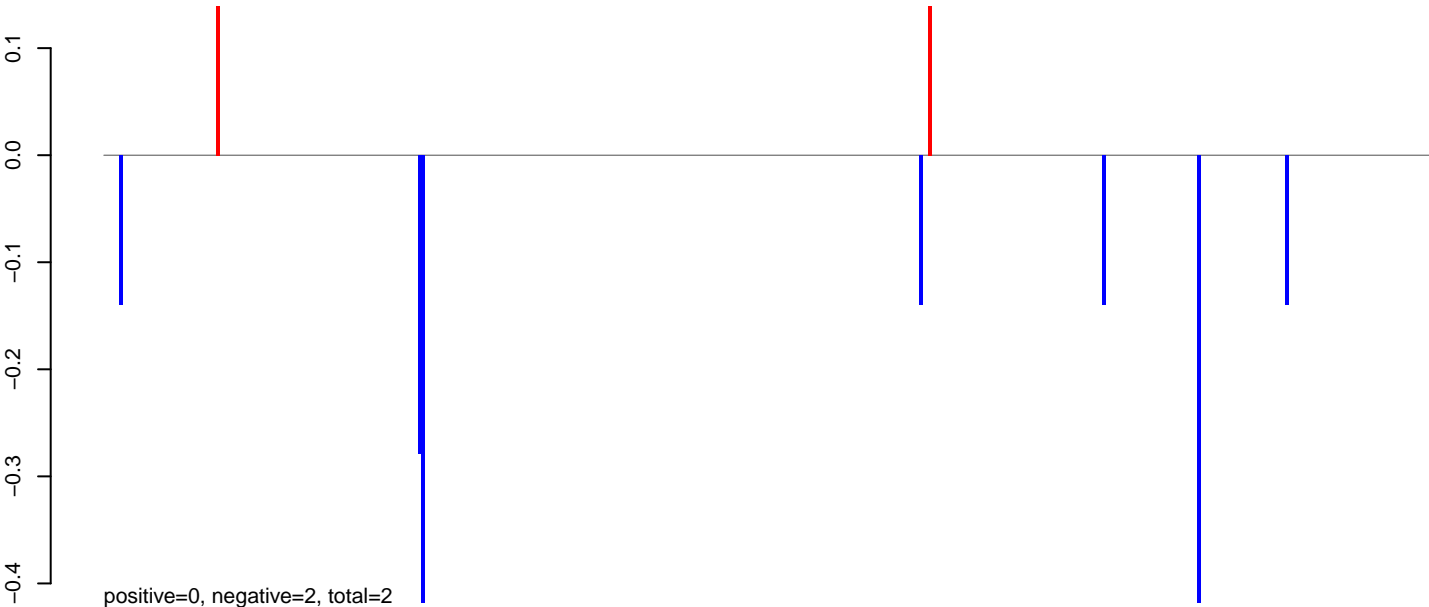


AeAlbo_C636_2dpi_mock3.rep

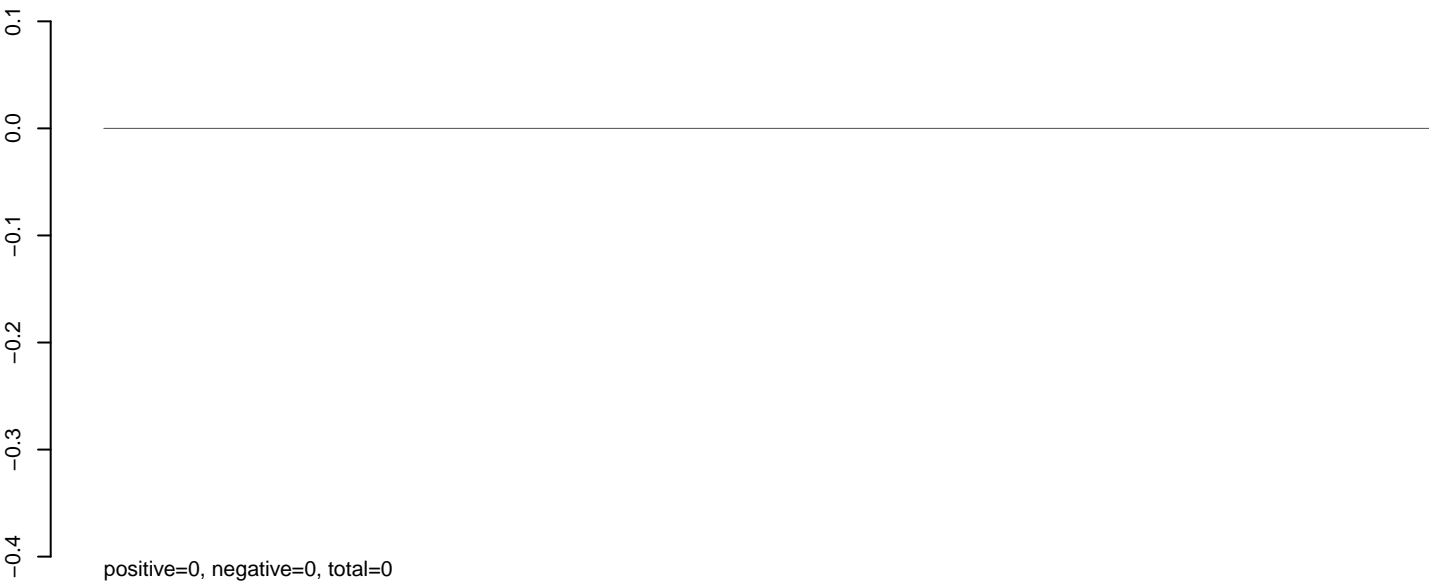


Window size=25, length=15444, virus@NC_025400.1-MalakaI_virus_isolate_SudAr_1169-64:1-15444

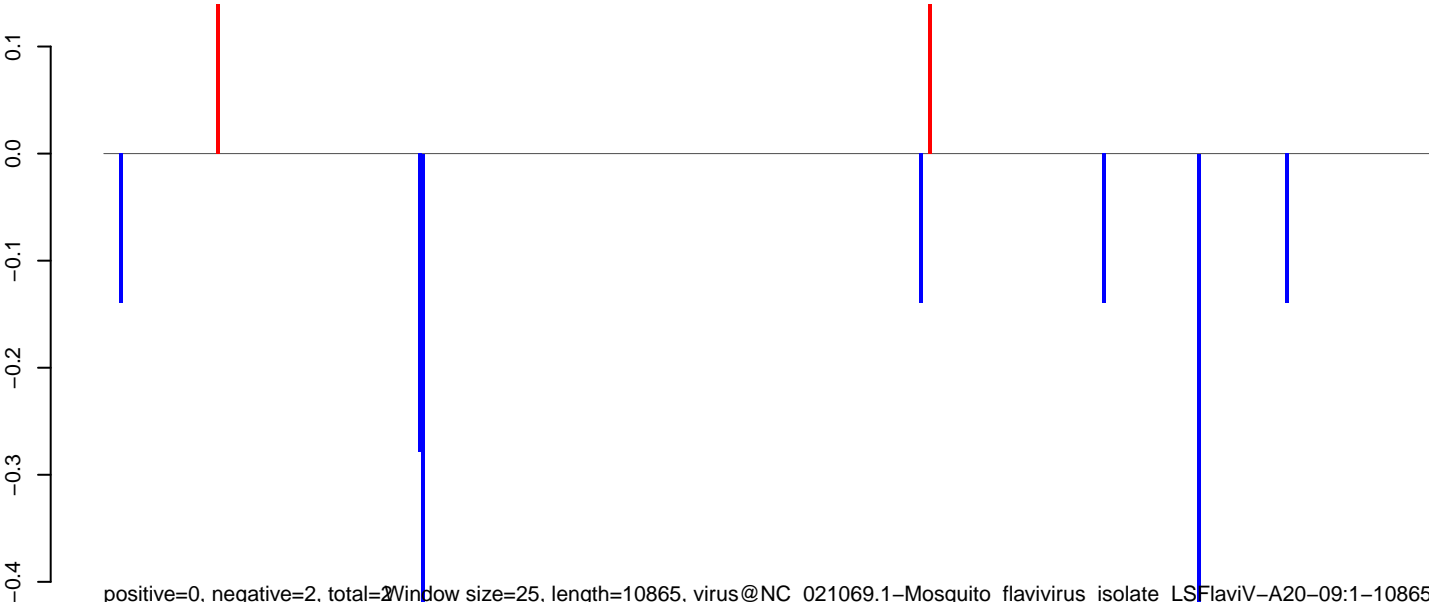
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

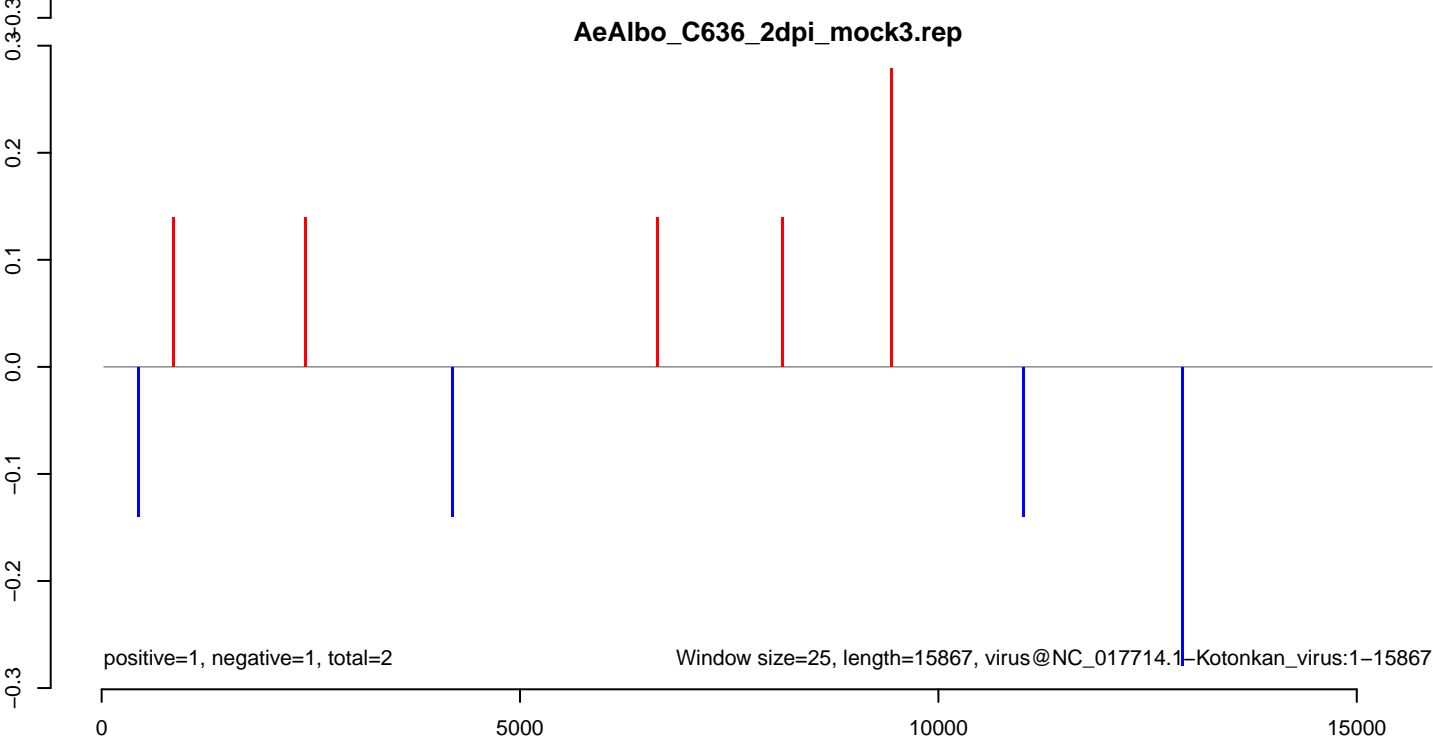
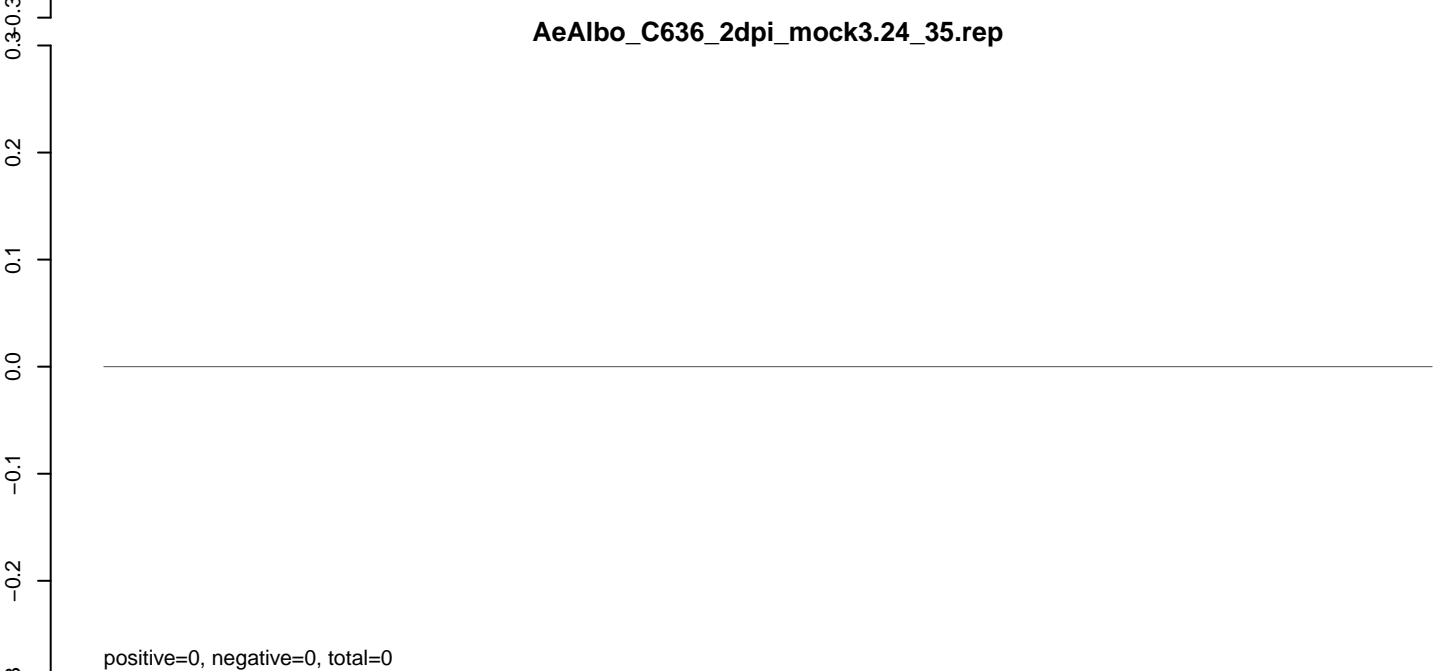
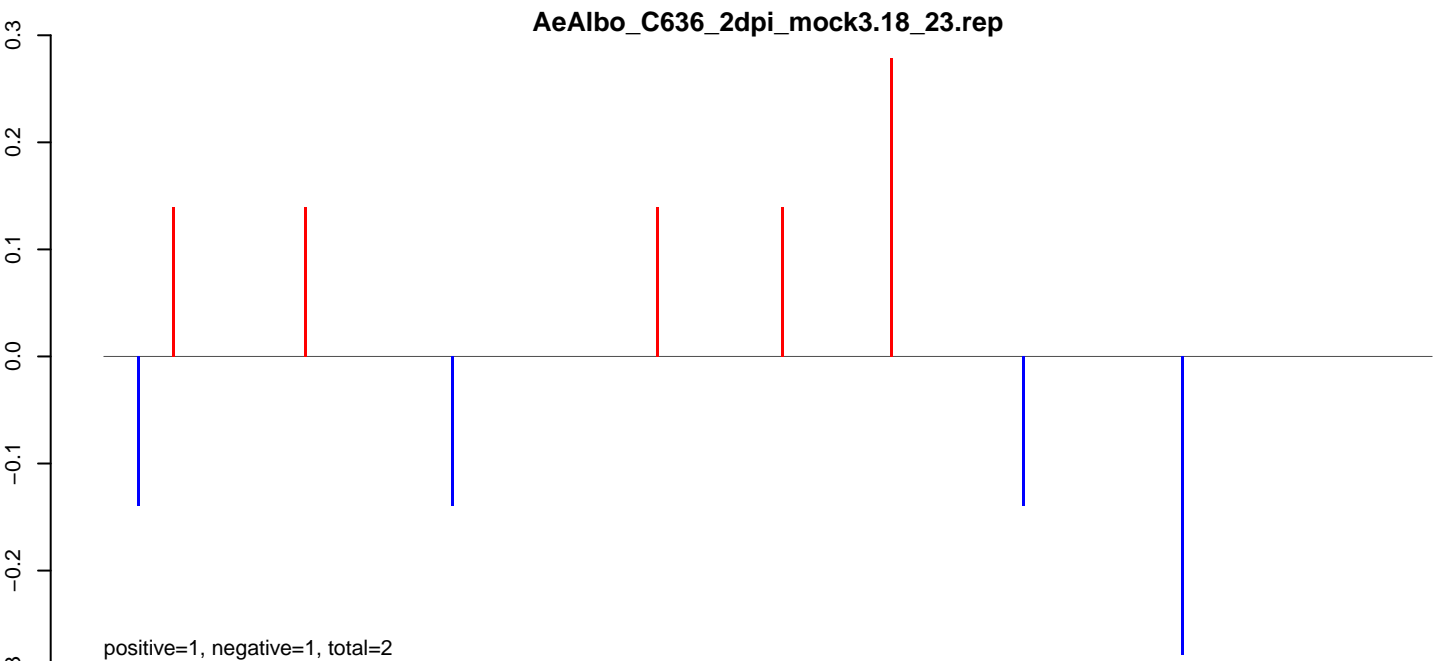


AeAlbo_C636_2dpi_mock3.rep

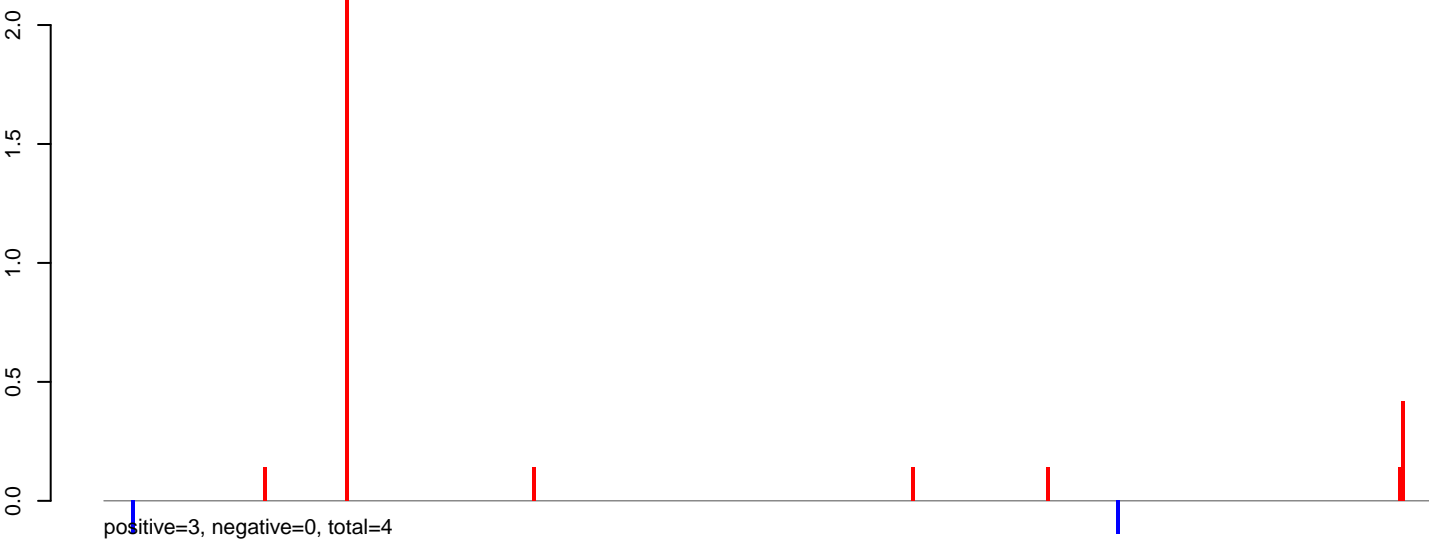


Window size=25, length=10865, virus@NC_021069.1-Mosquito_flavivirus_isolate_LSFlaviV-A20-09:1-10865

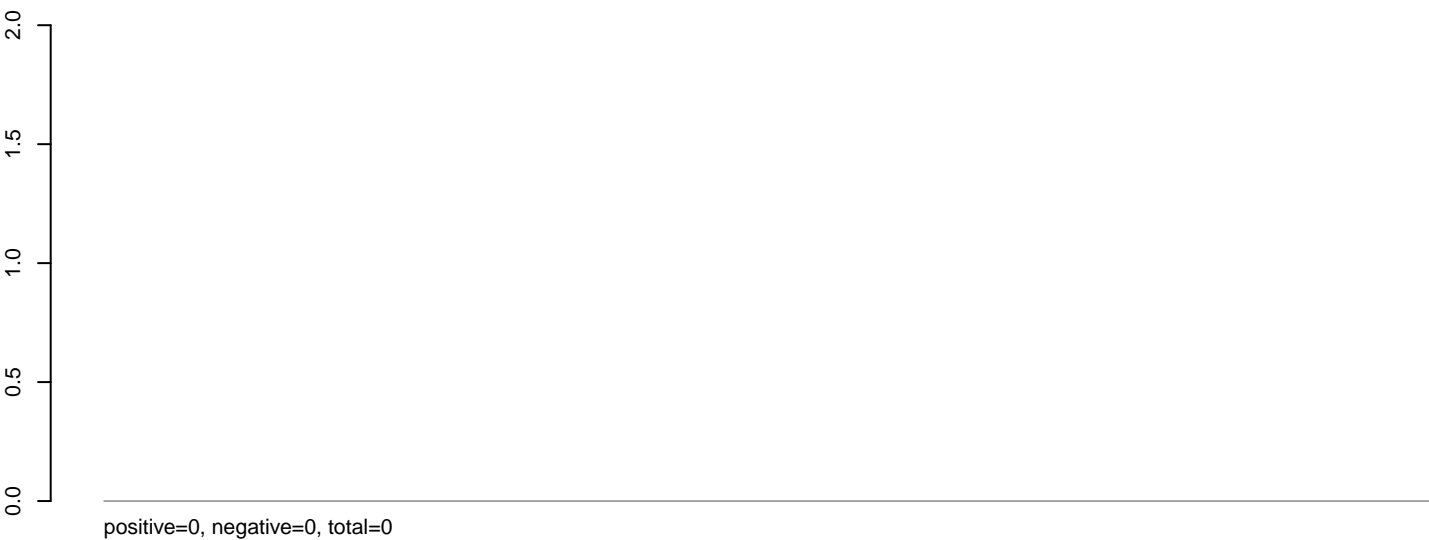
0 2000 4000 6000 8000 10000



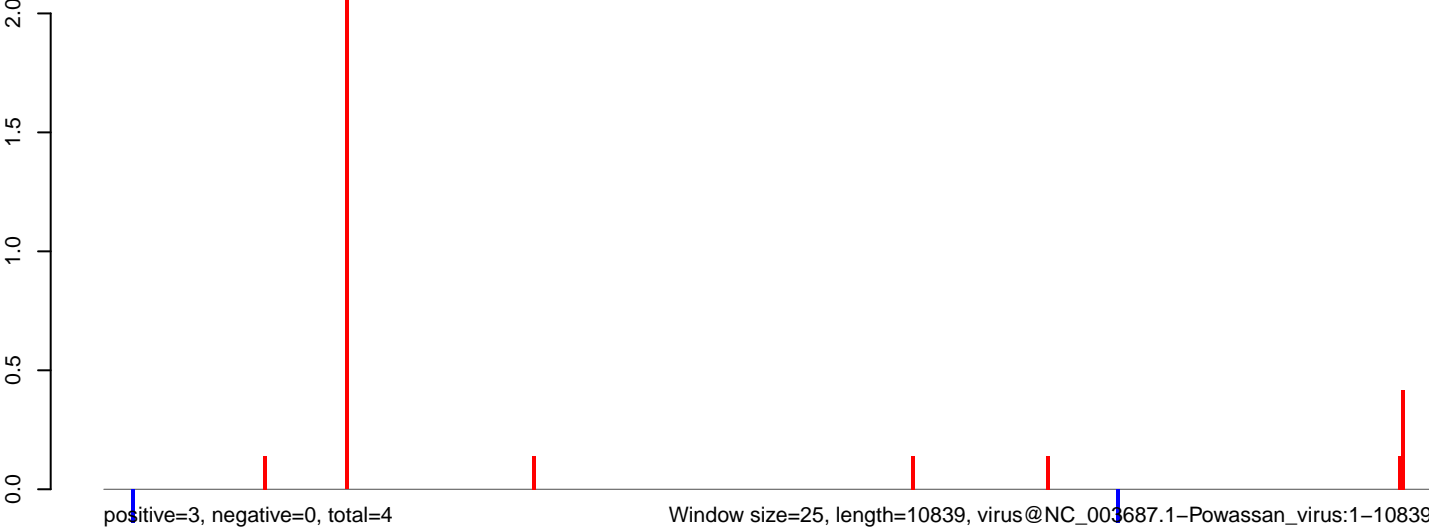
AeAlbo_C636_2dpi_mock3.18_23.rep



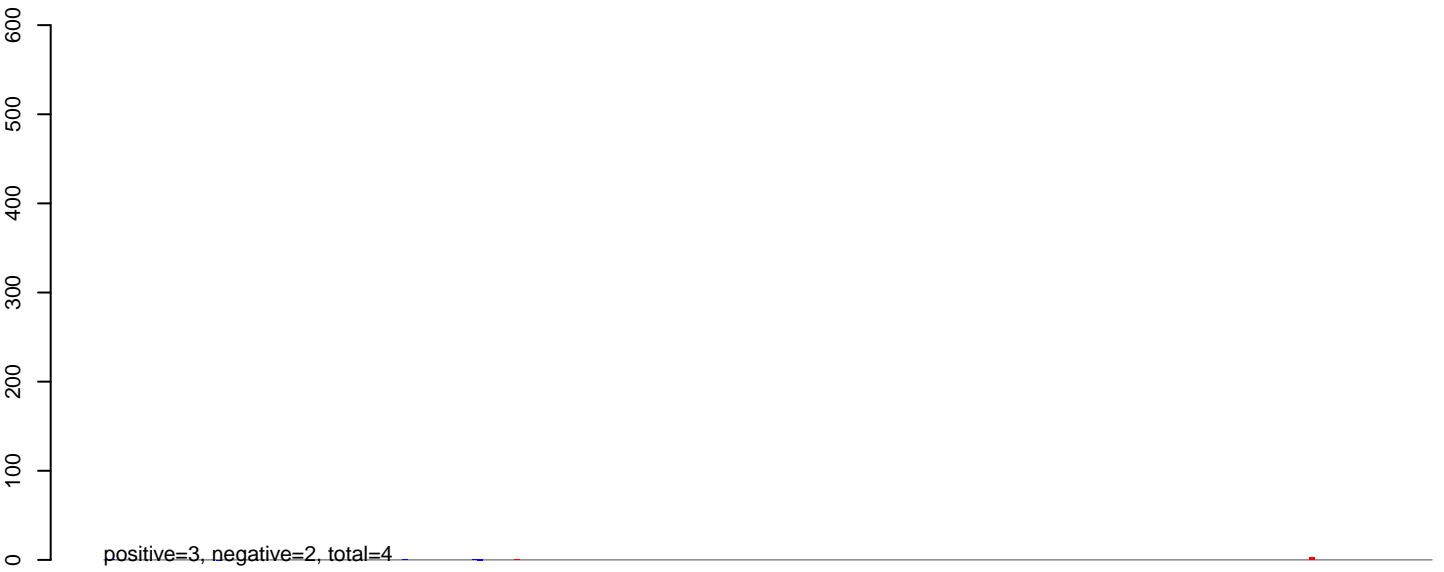
AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



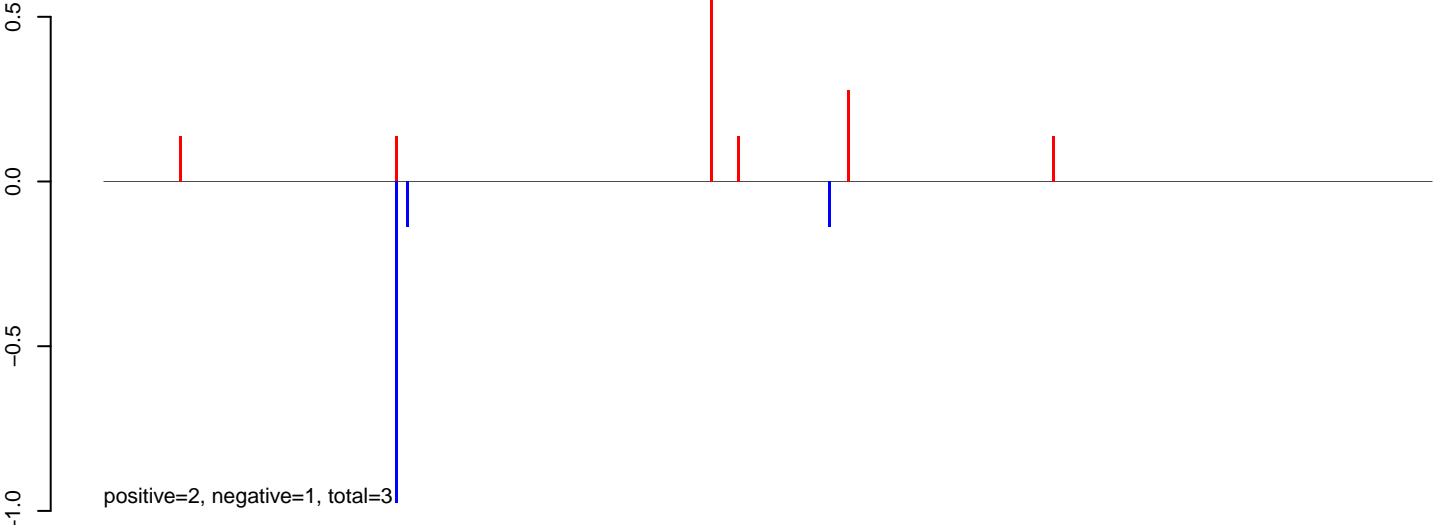
AeAlbo_C636_2dpi_mock3.rep



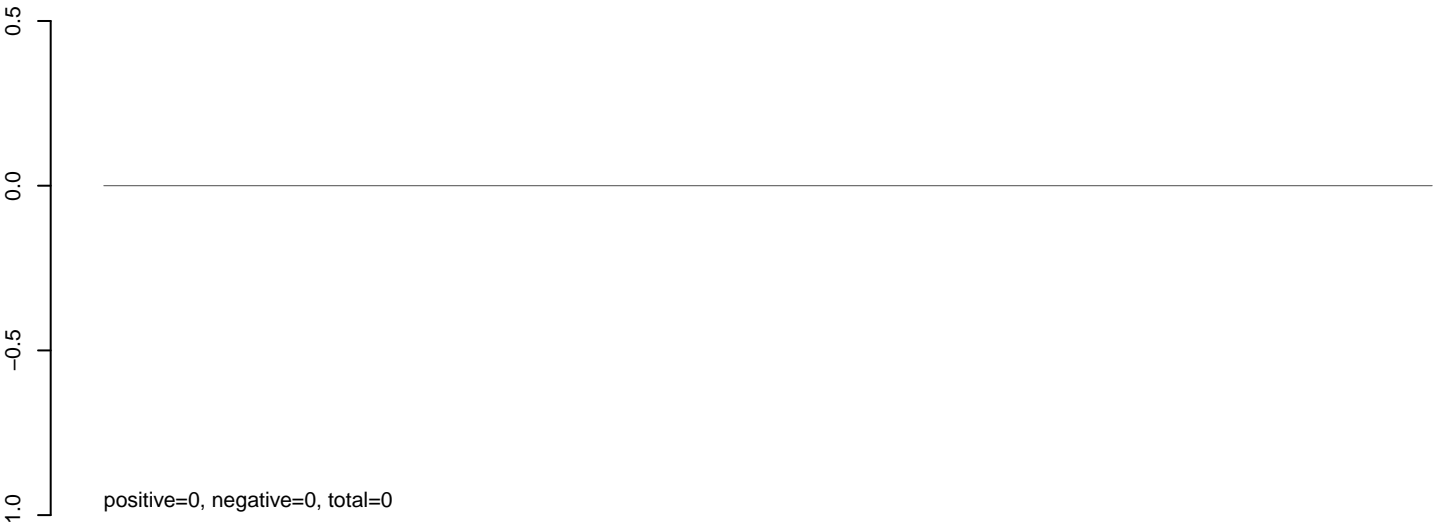
positive=671, negative=3, total=674, length=6203, virus@NC_035674.1-Australian_Anopheles_totivirus_isolate_AATV_150840:1-6203

0 1000 2000 3000 4000 5000 6000

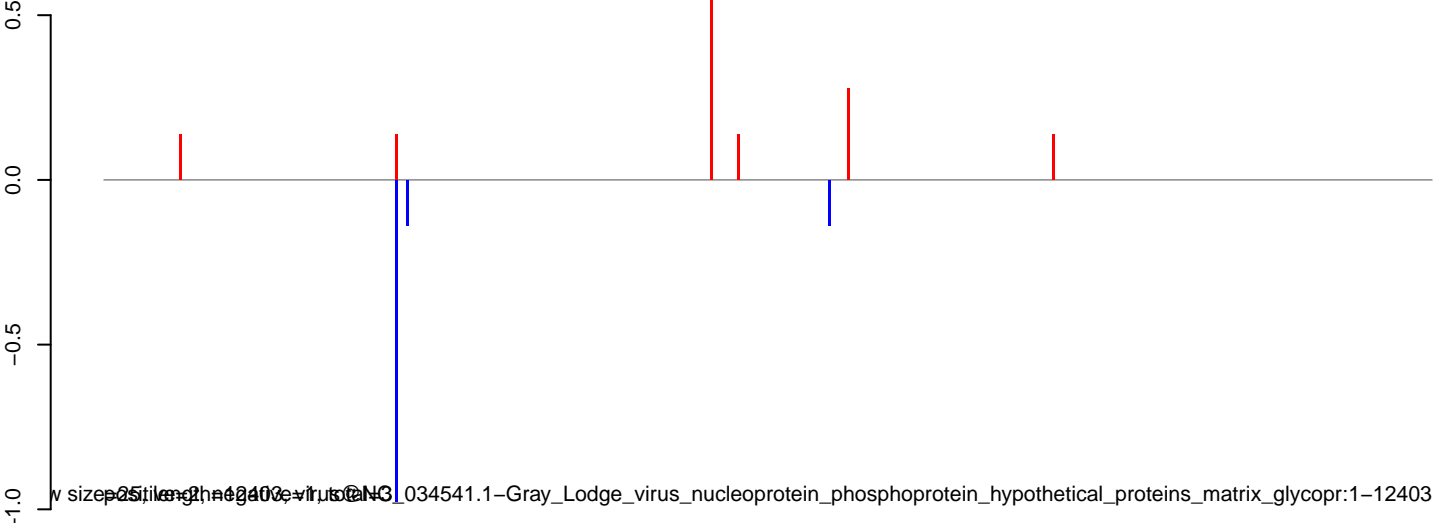
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

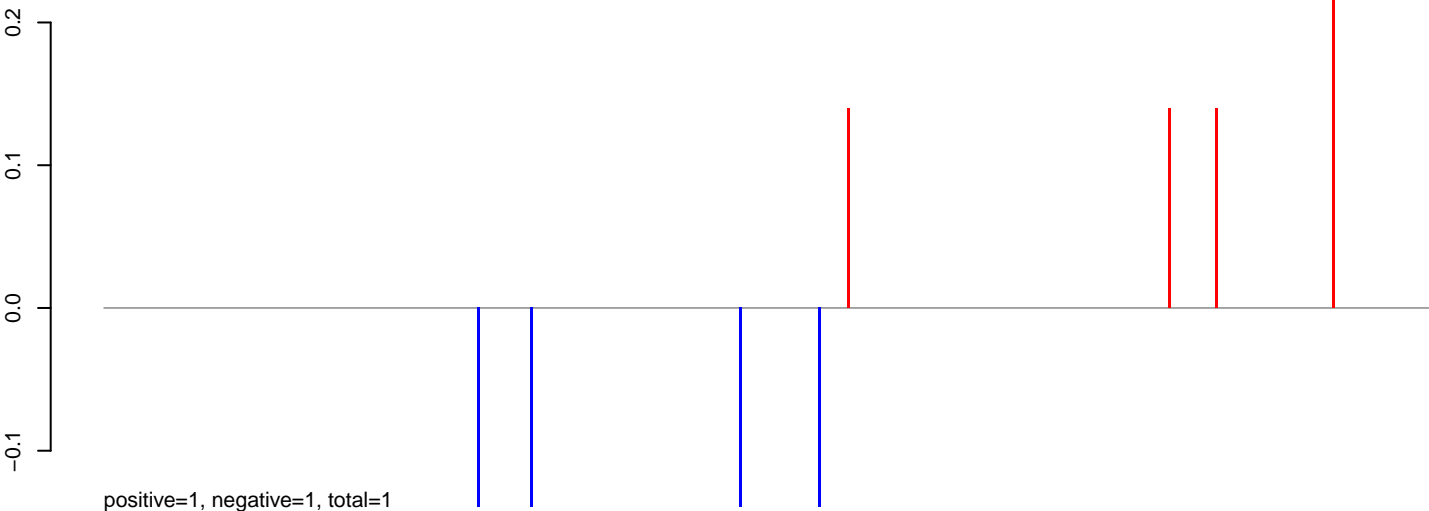


AeAlbo_C636_2dpi_mock3.rep

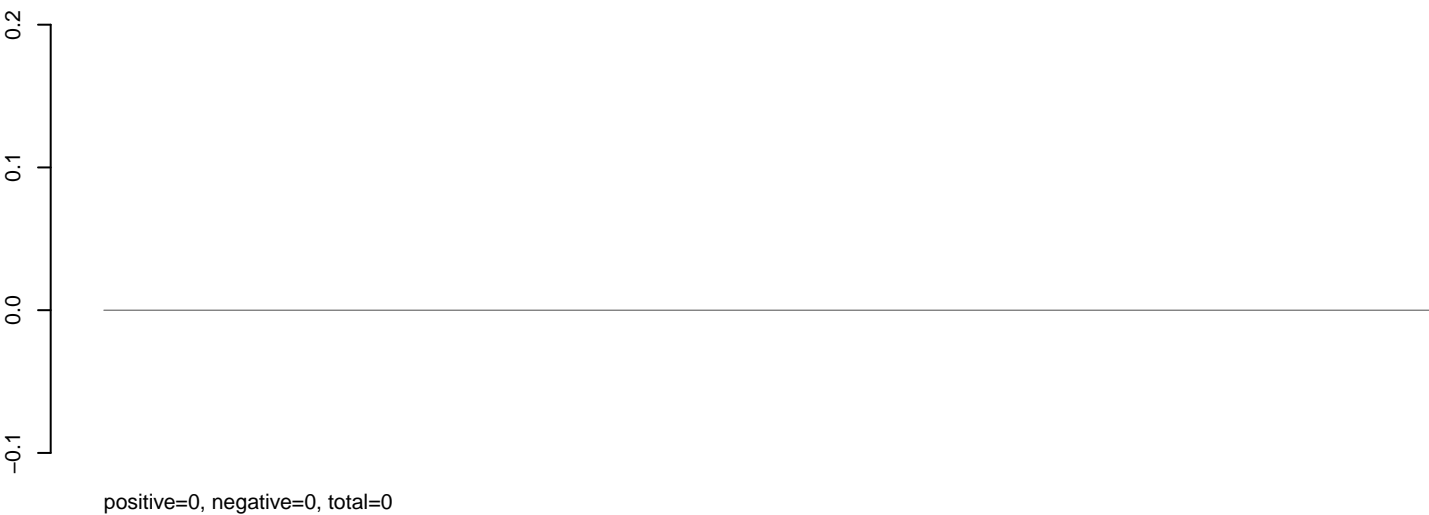


view sizes 25, line graph 400, virus @ NC_034541.1-Gray_Lodge_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopr:1-12403

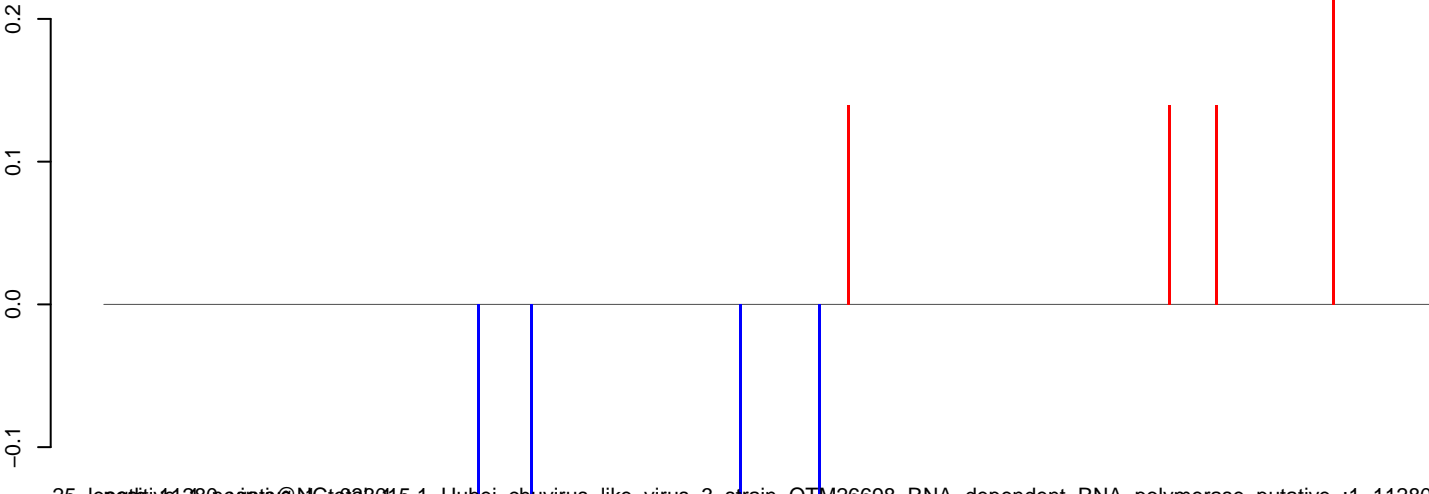
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



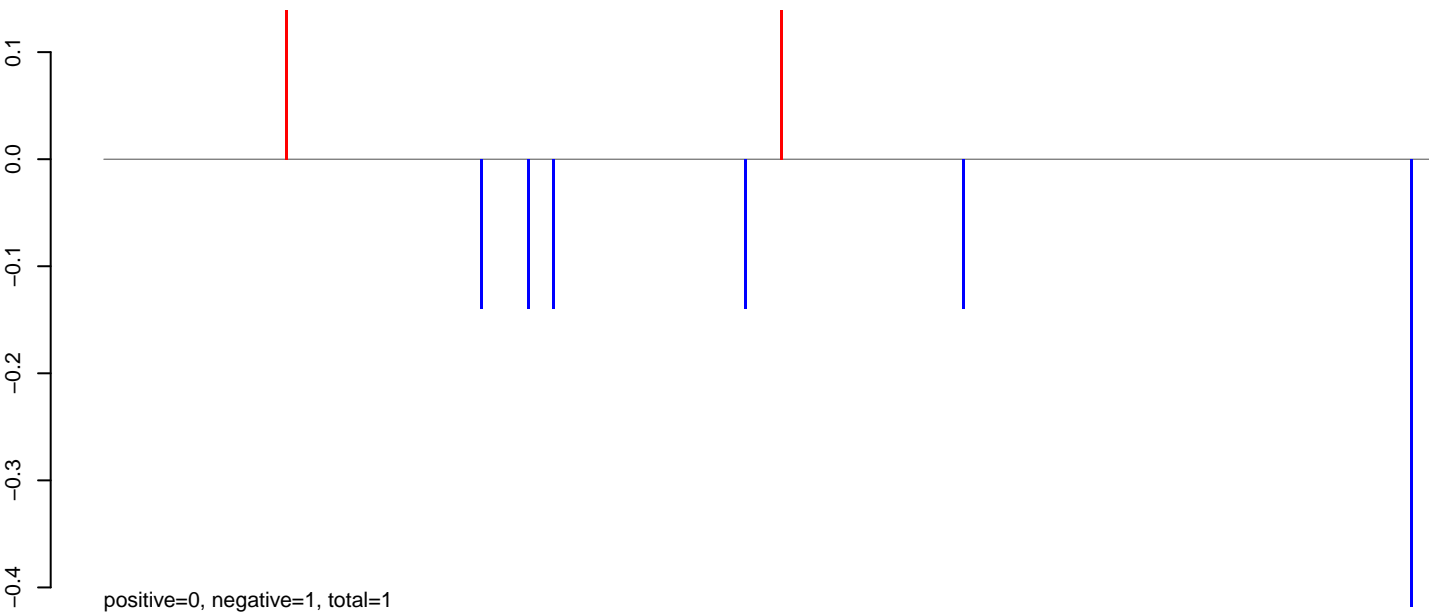
AeAlbo_C636_2dpi_mock3.rep



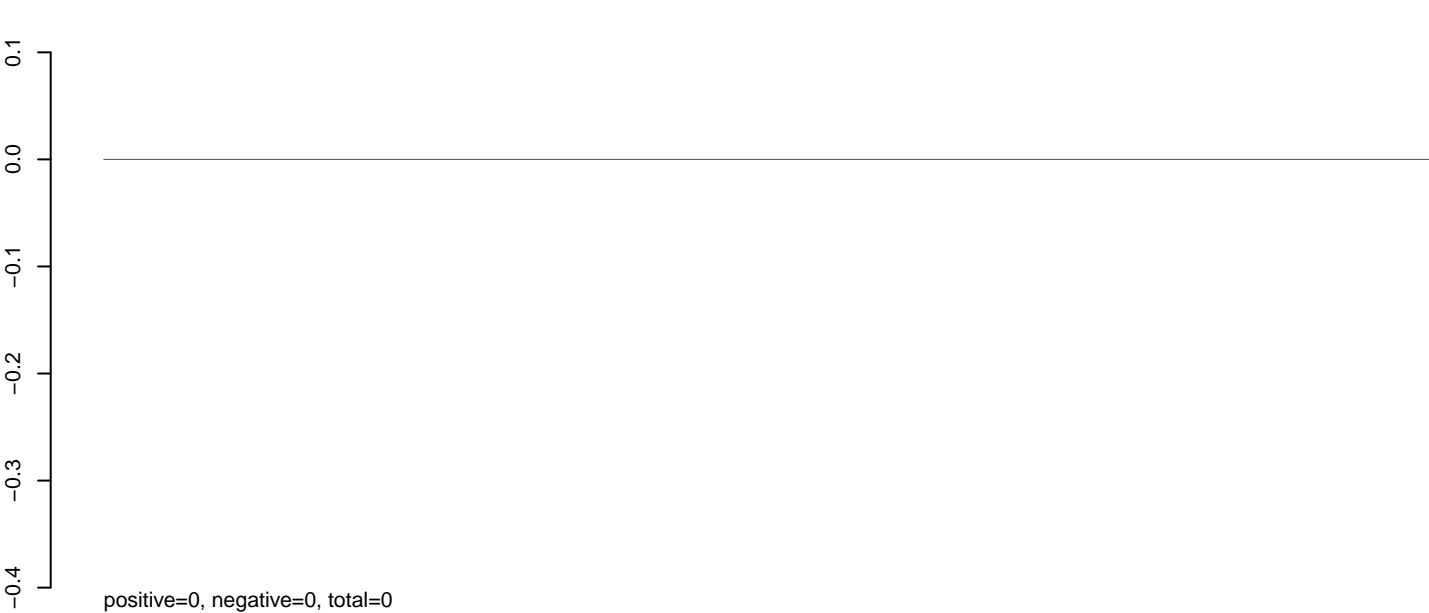
25, length=280, negative=0, NC_023801.1-Hubei_chuvirus-like_virus_3_strain_QTM26698_RNA-dependent_RNA_polymerase_putative_:1-11280

0 2000 4000 6000 8000 10000

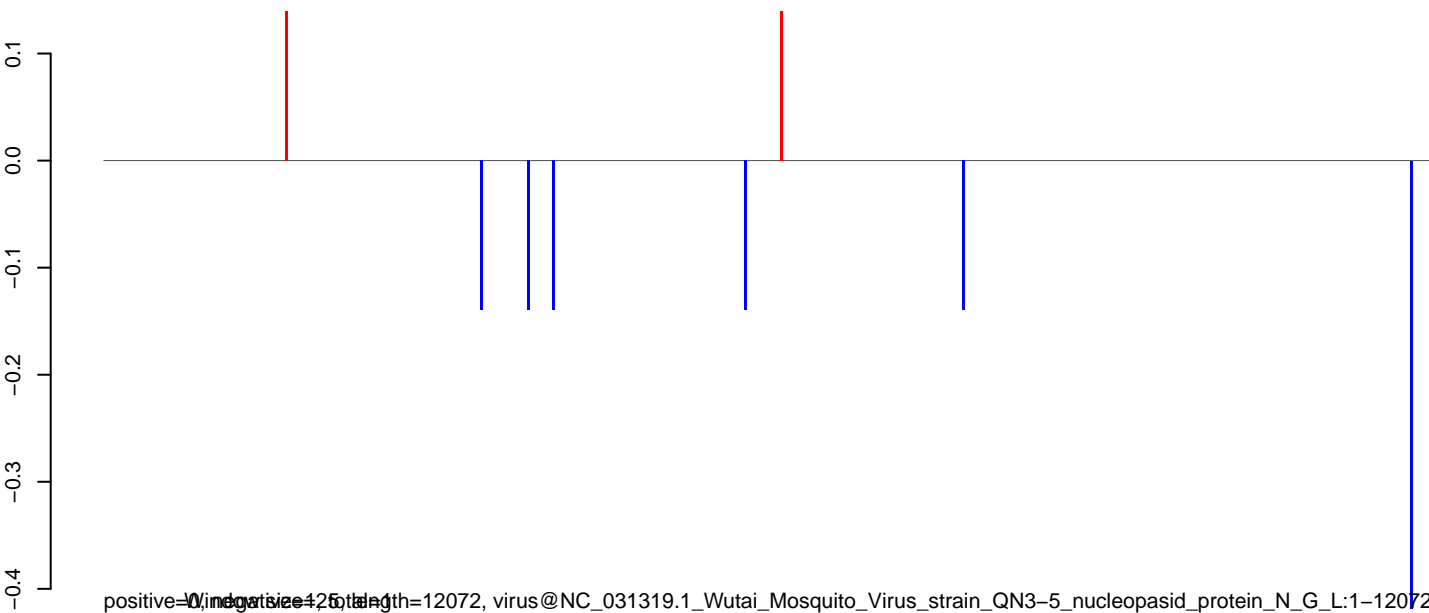
AeAlbo_C636_2dpi_mock3.18_23.rep



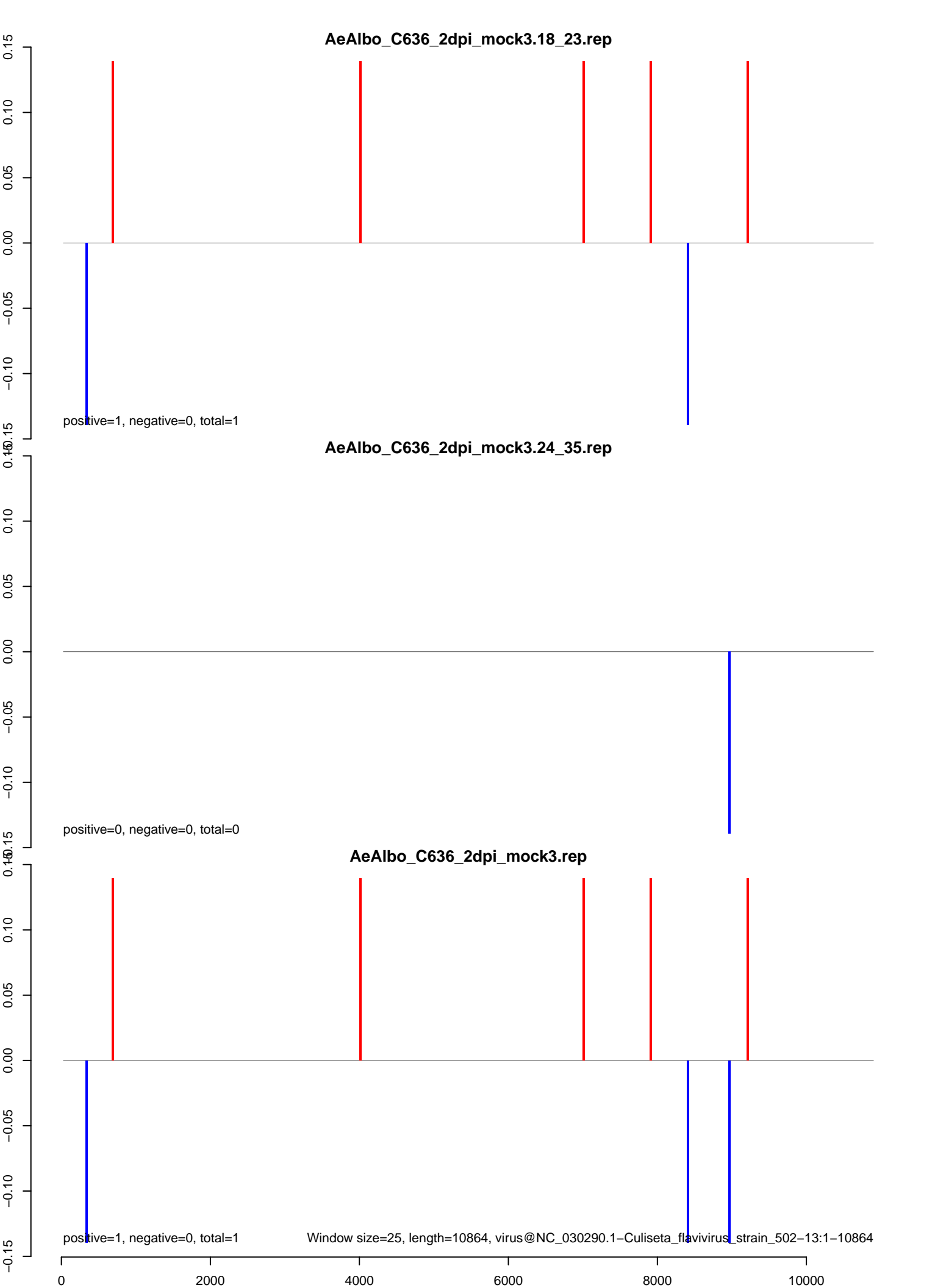
AeAlbo_C636_2dpi_mock3.24_35.rep



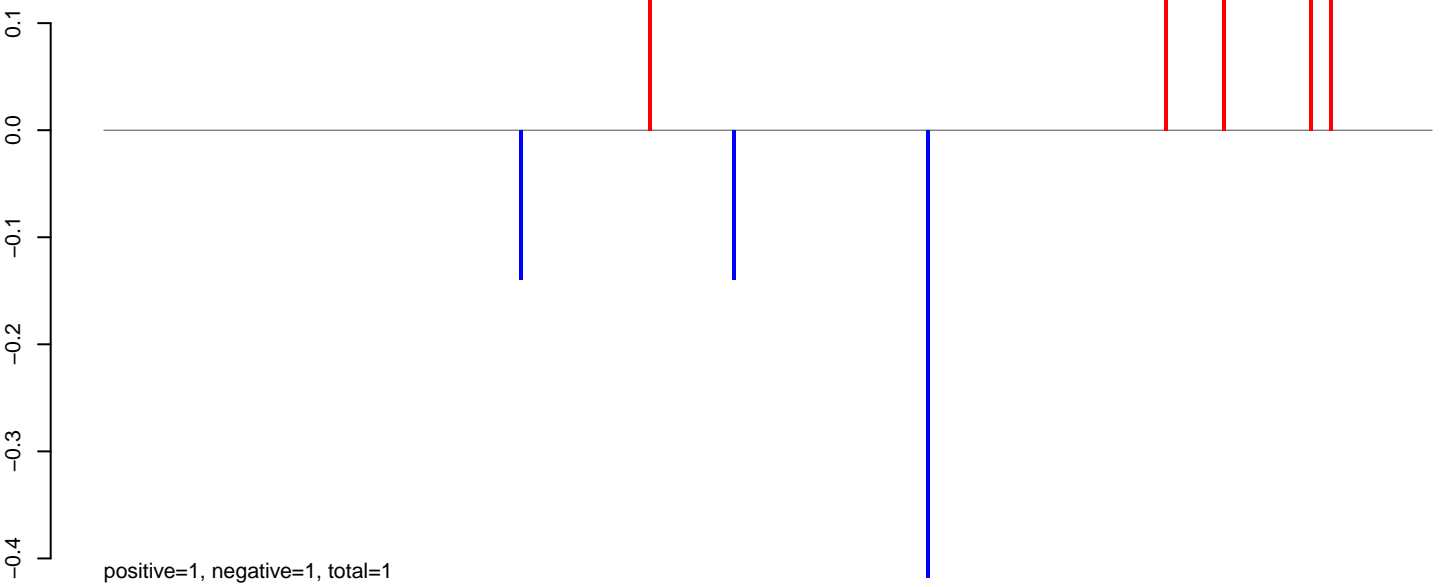
AeAlbo_C636_2dpi_mock3.rep



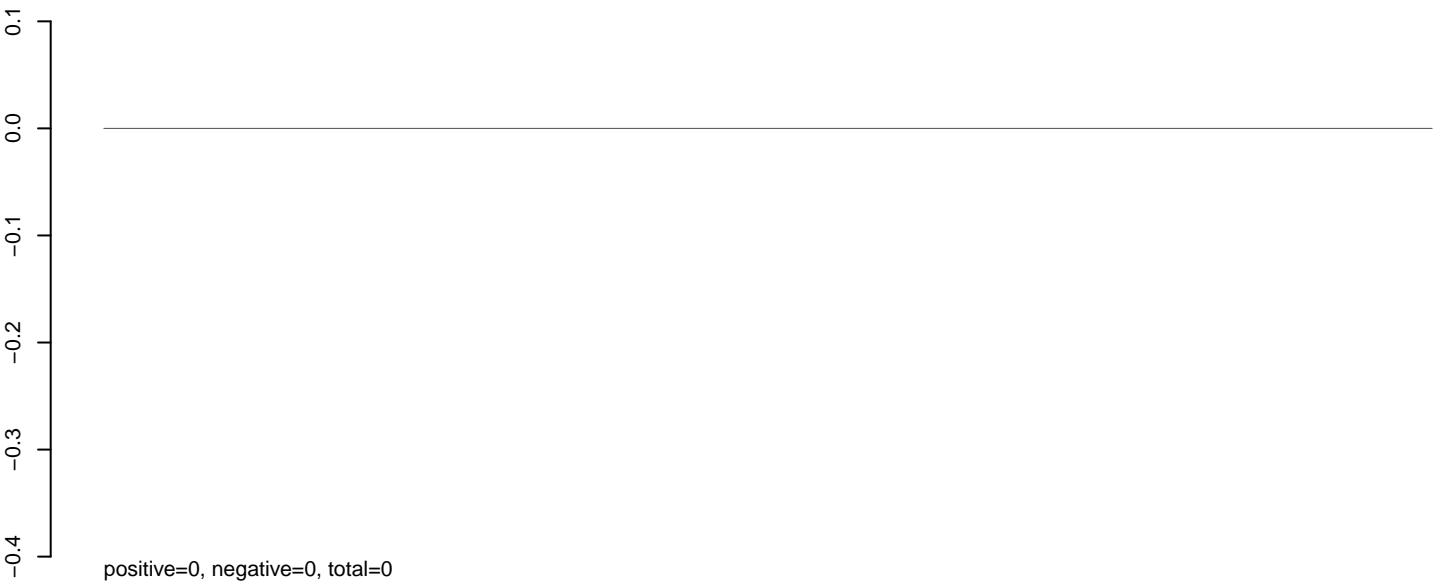
positive=0, negative=1, total=1
positive=0, negative=0, total=0
positive=0, negative=1, total=1
virus@NC_031319.1_Wutai_Mosquito_Virus_strain_QN3-5_nucleopasid_protein_N_G_L:1-12072



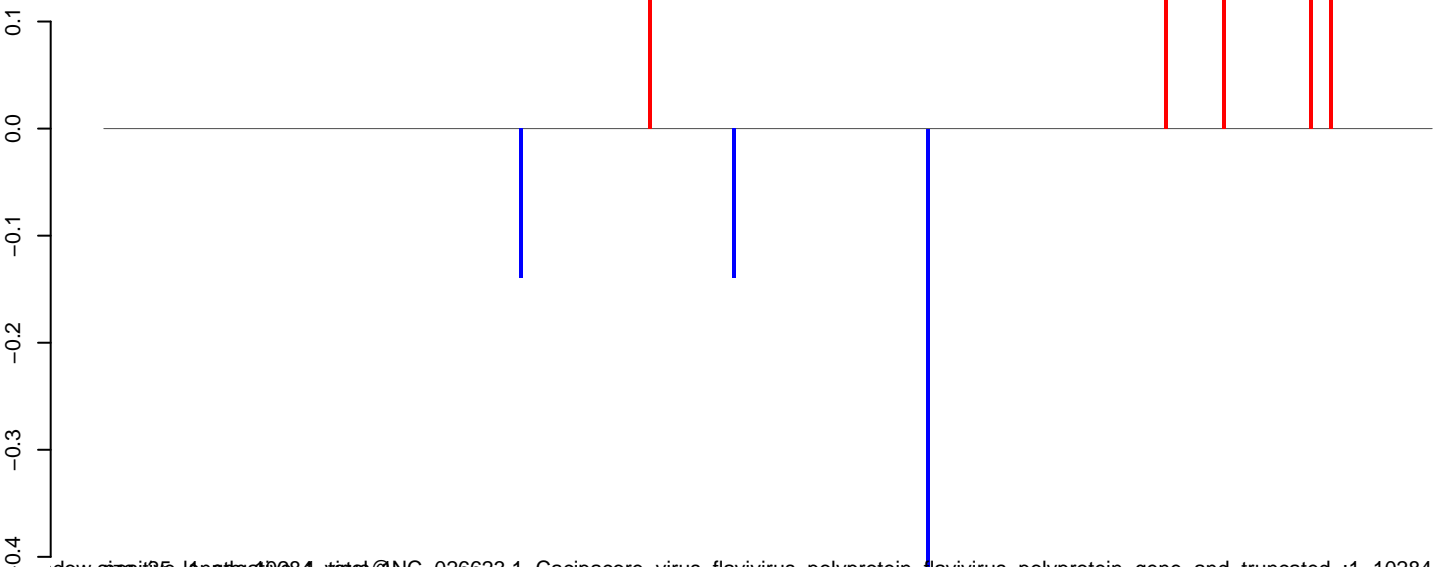
AeAlbo_C636_2dpi_mock3.18_23.rep



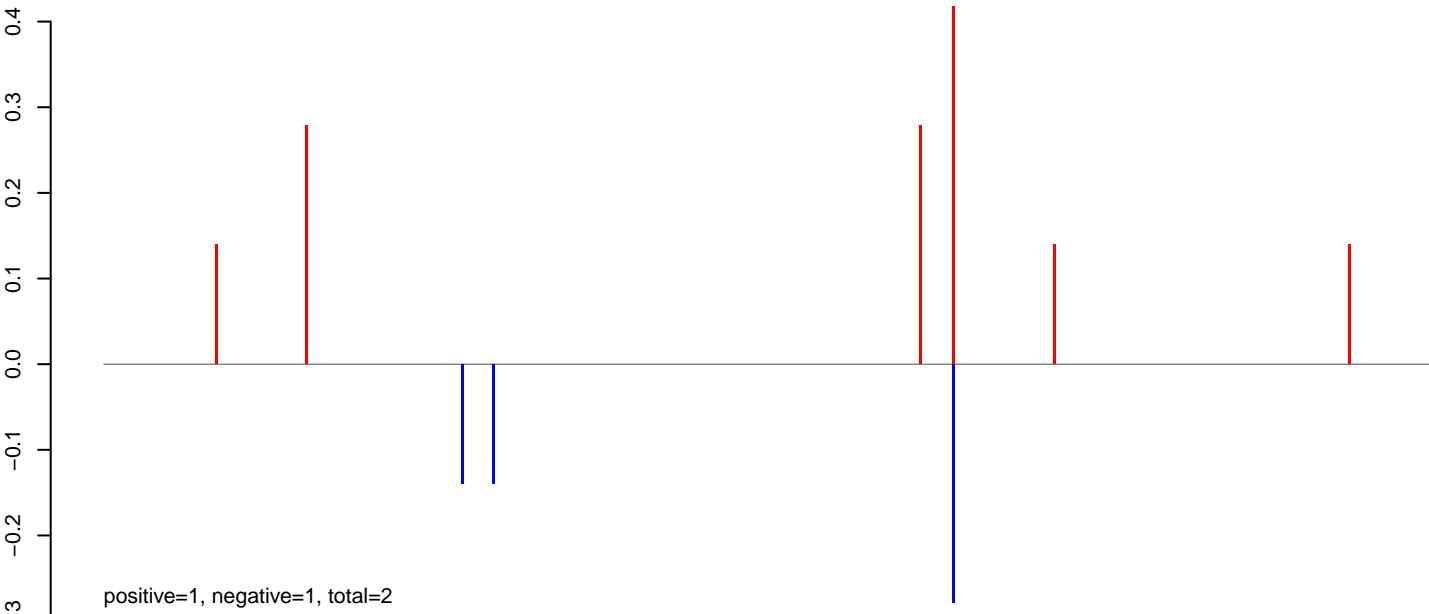
AeAlbo_C636_2dpi_mock3.24_35.rep



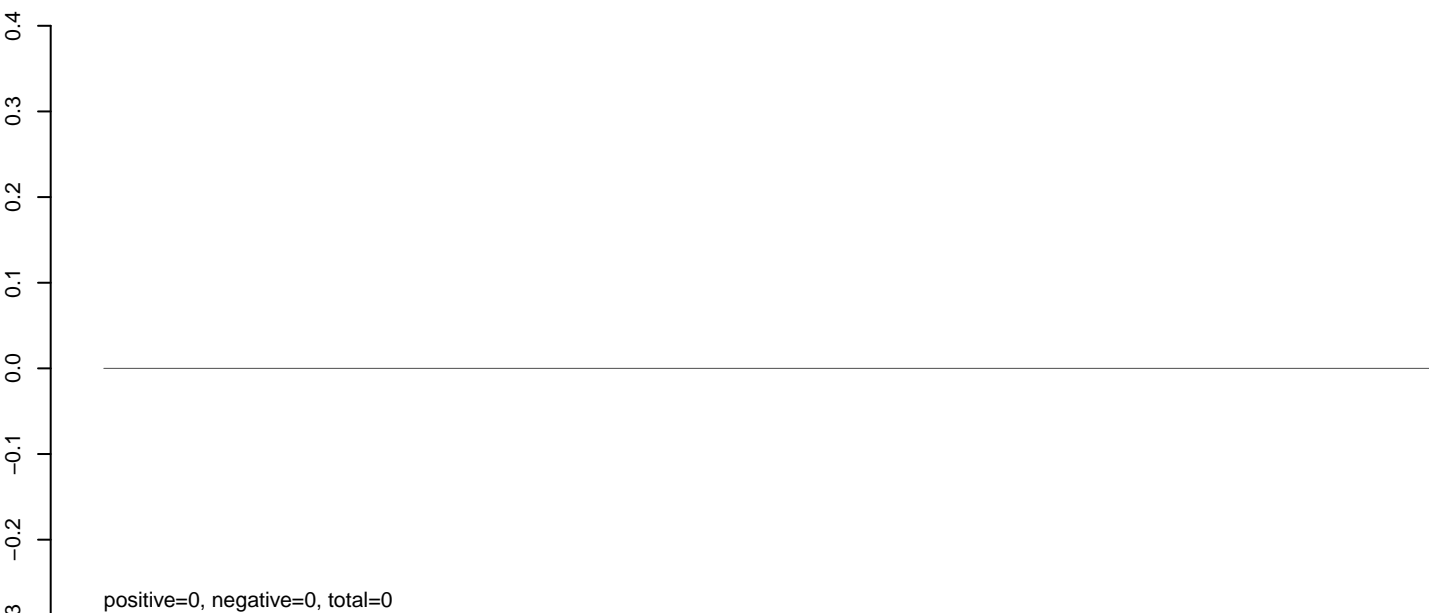
AeAlbo_C636_2dpi_mock3.rep



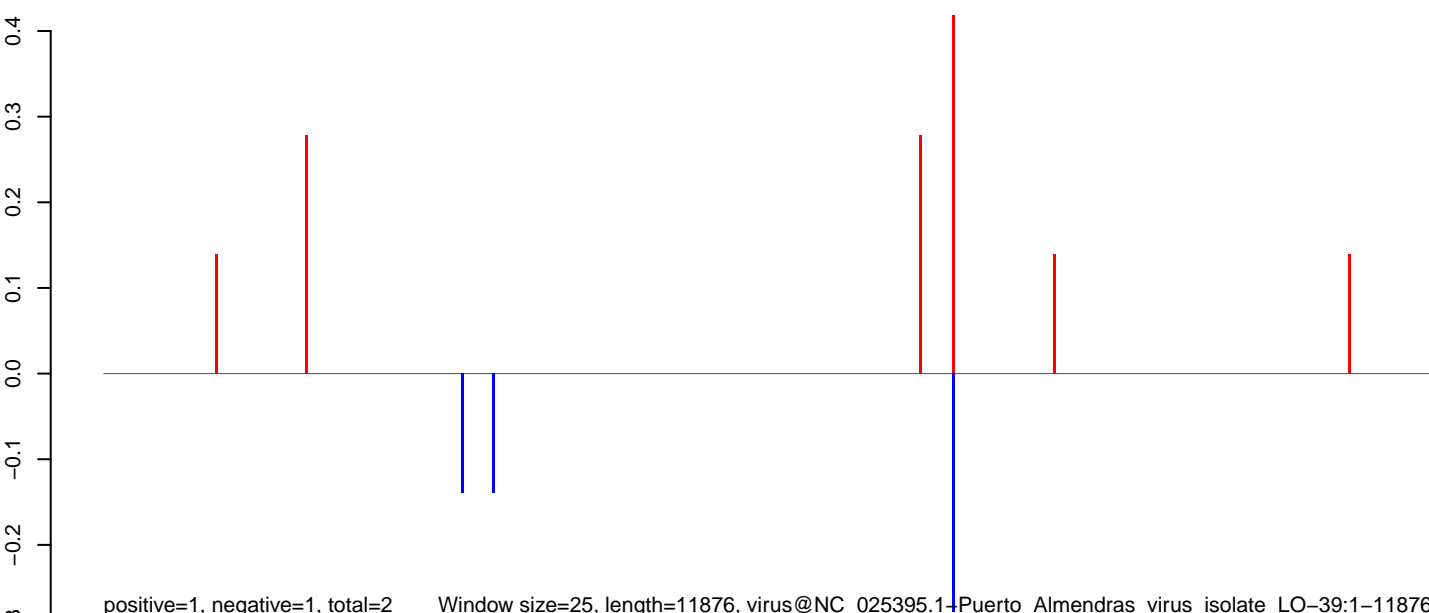
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

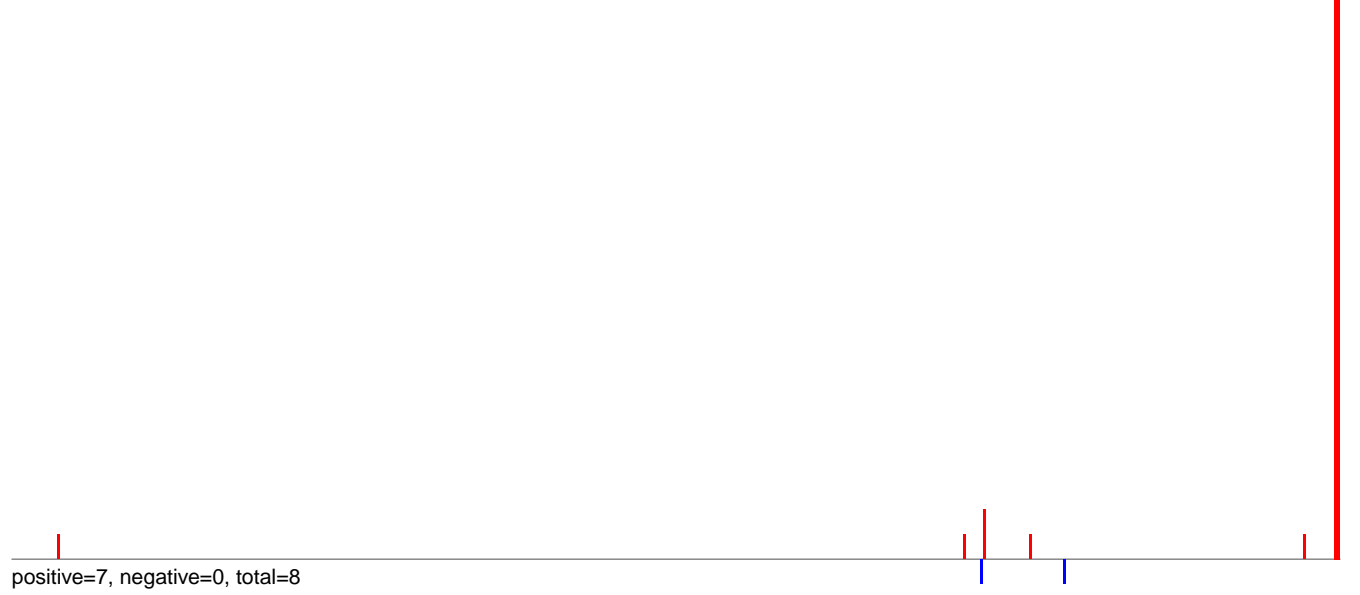


AeAlbo_C636_2dpi_mock3.rep

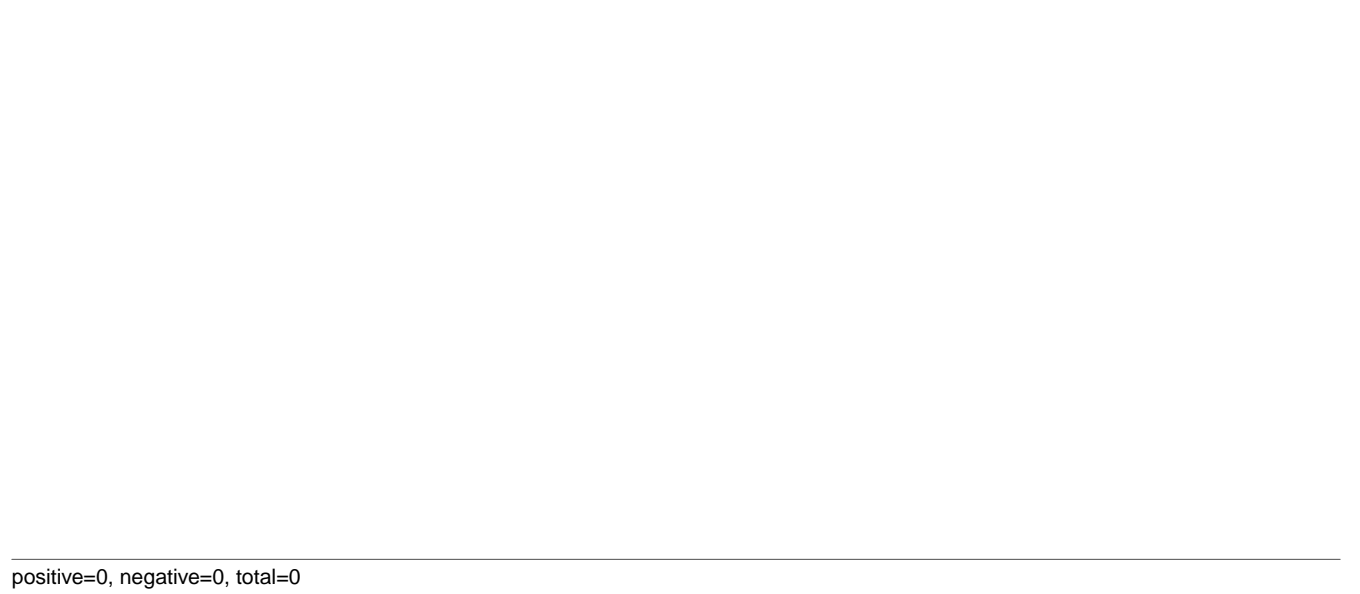


Window size=25, length=11876, virus@NC_025395.1-Puerto_Almendras_virus_isolate_LO-39:1-11876

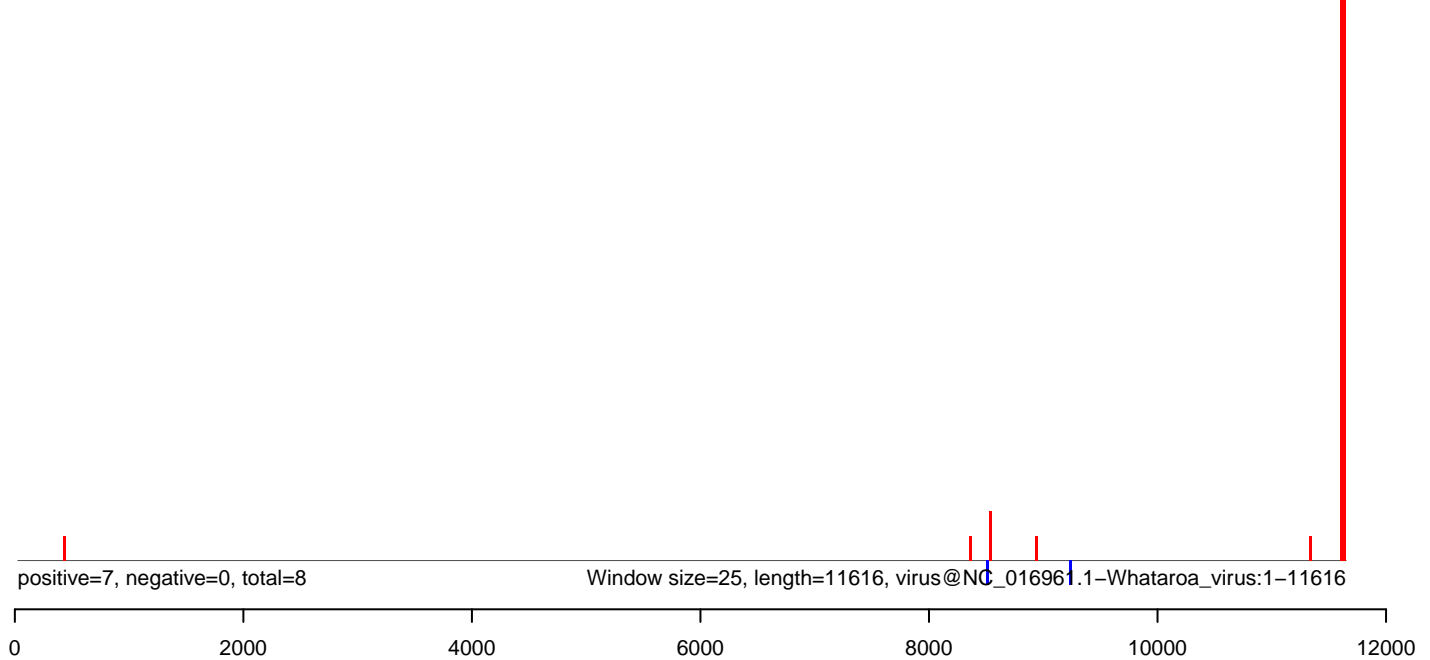
AeAlbo_C636_2dpi_mock3.18_23.rep

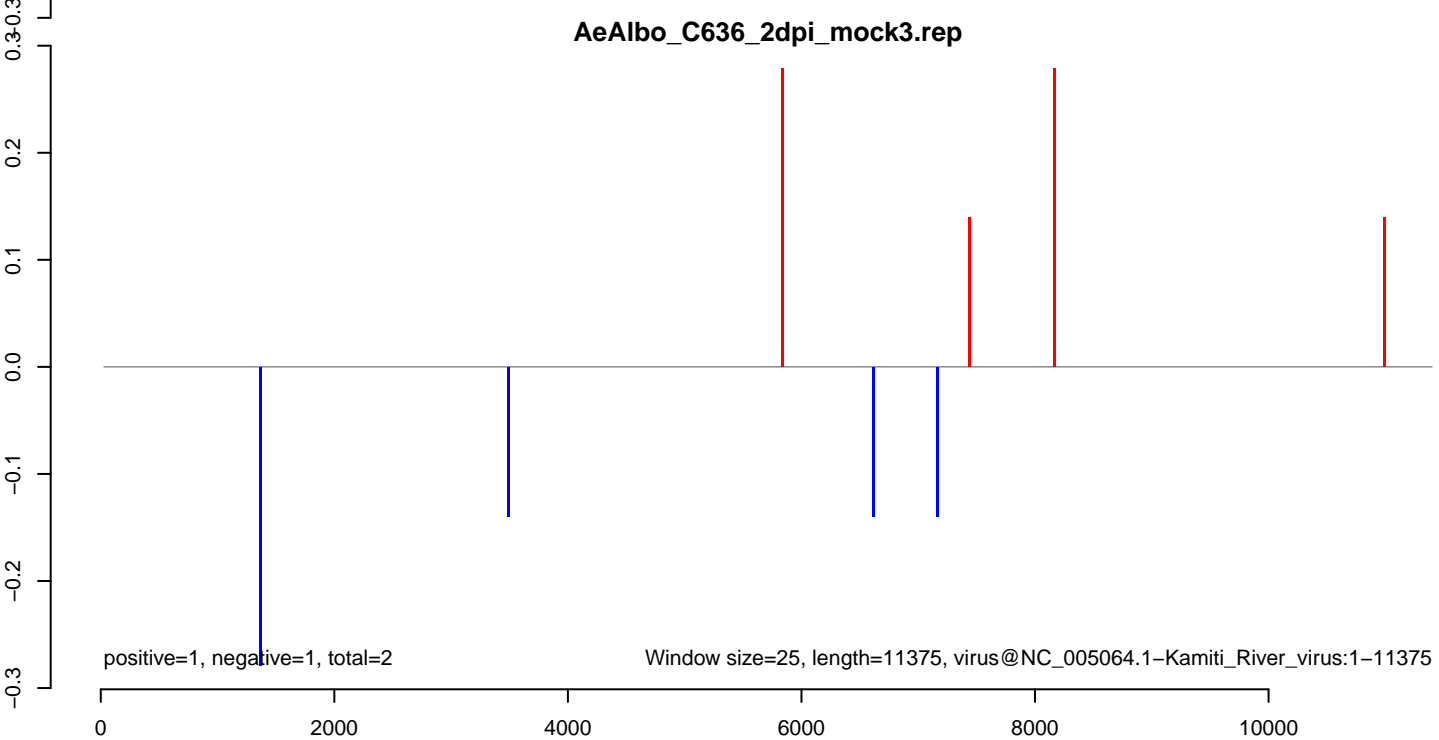
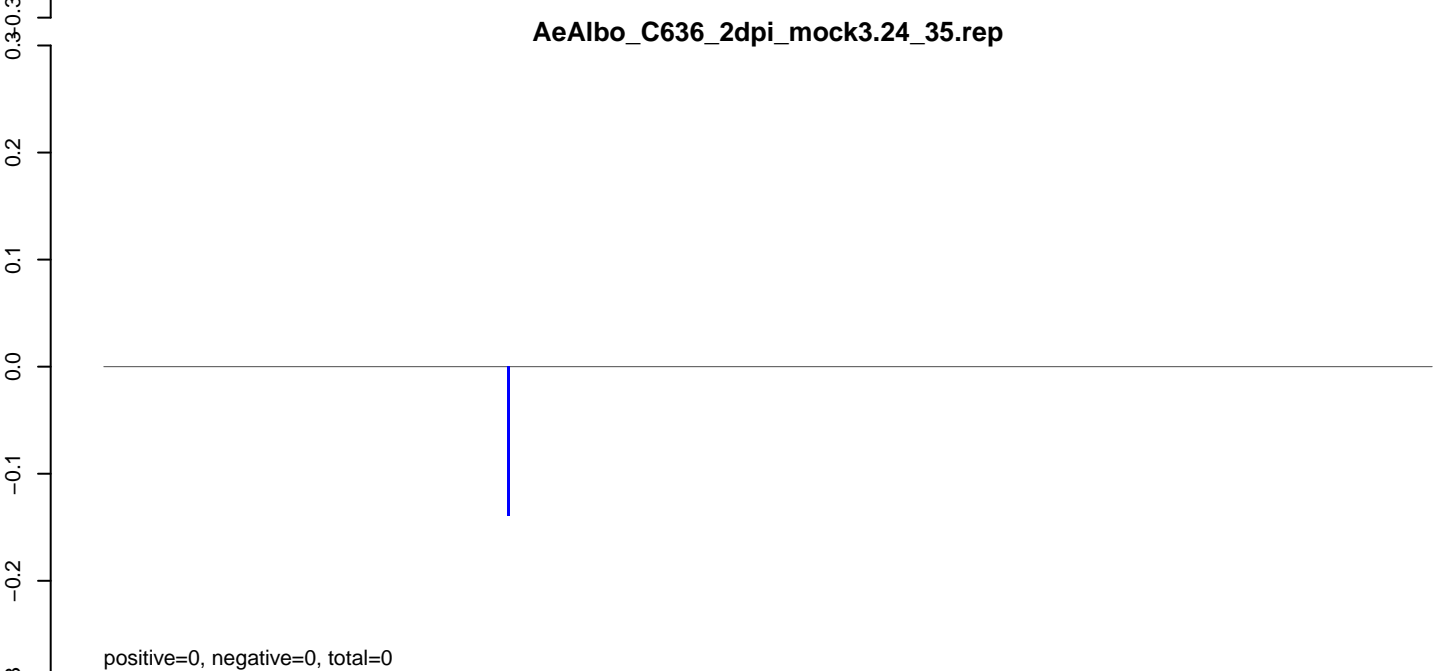
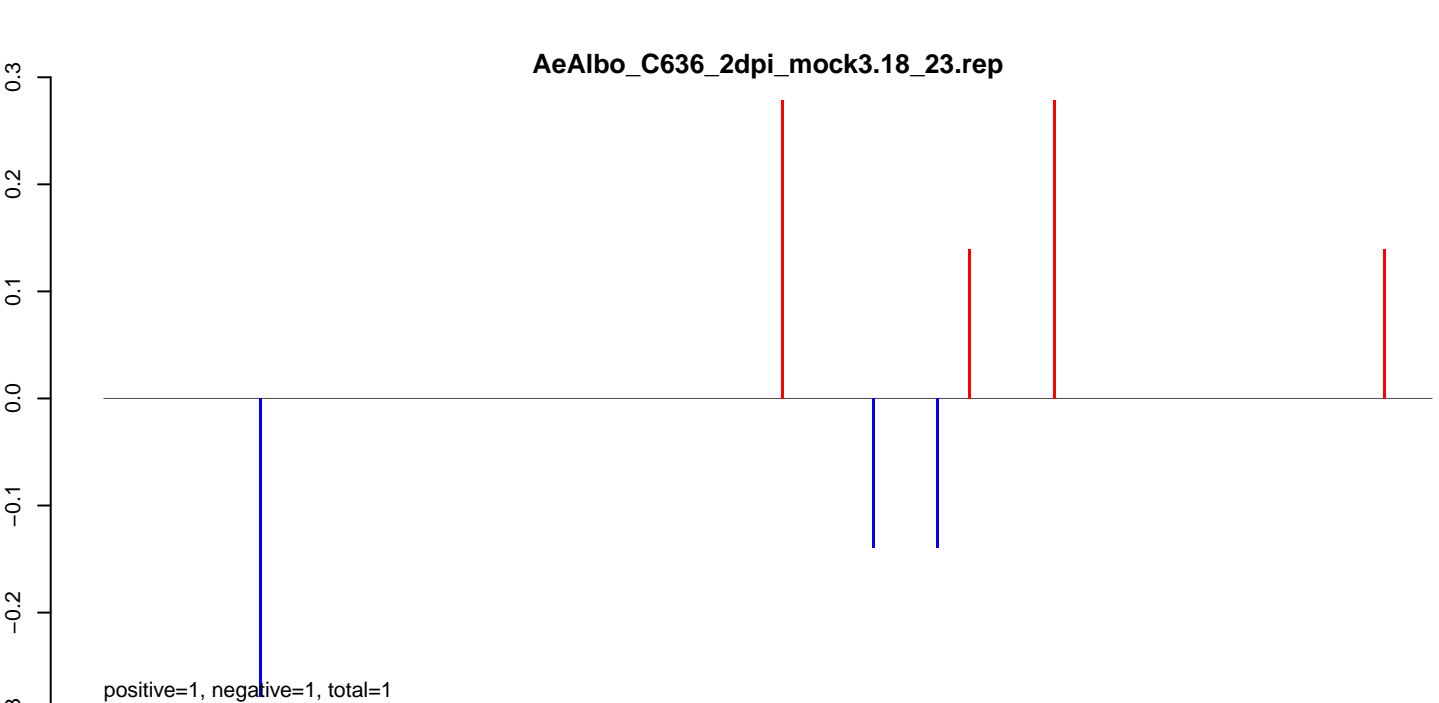


AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep





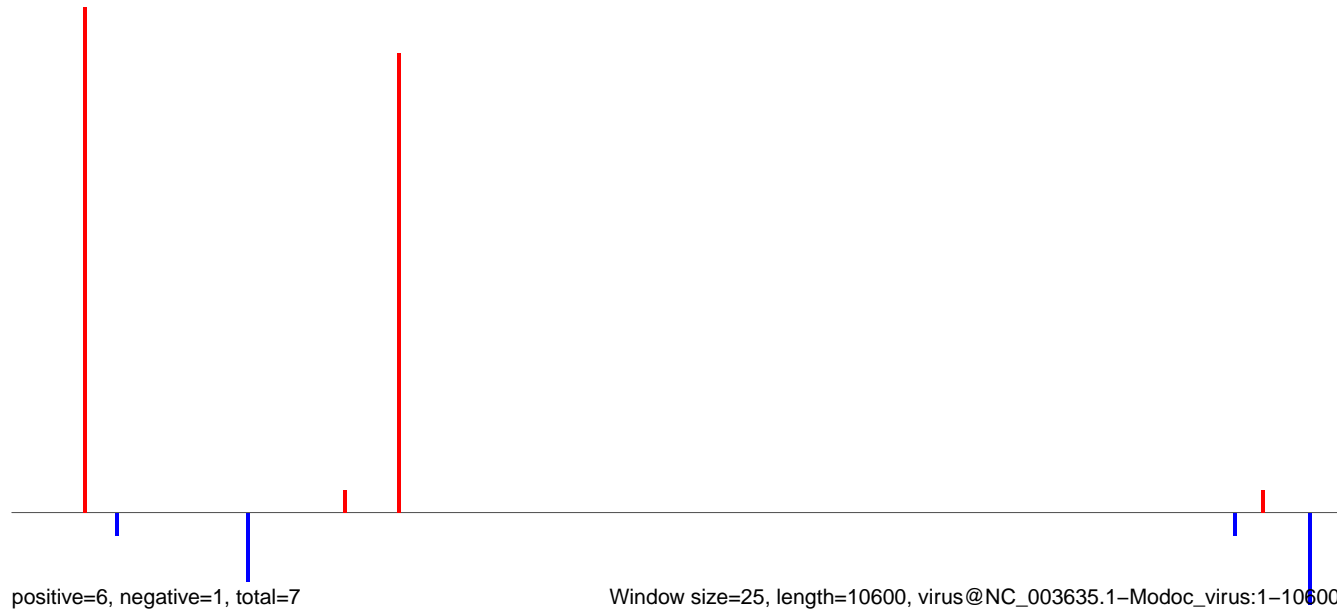
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



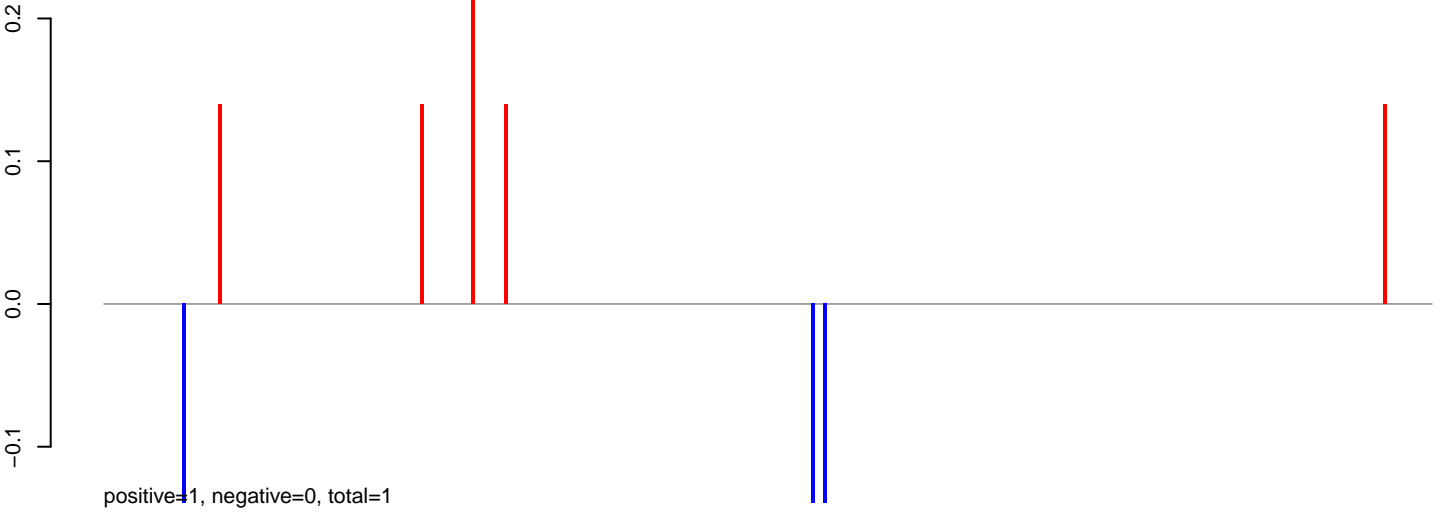
AeAlbo_C636_2dpi_mock3.rep



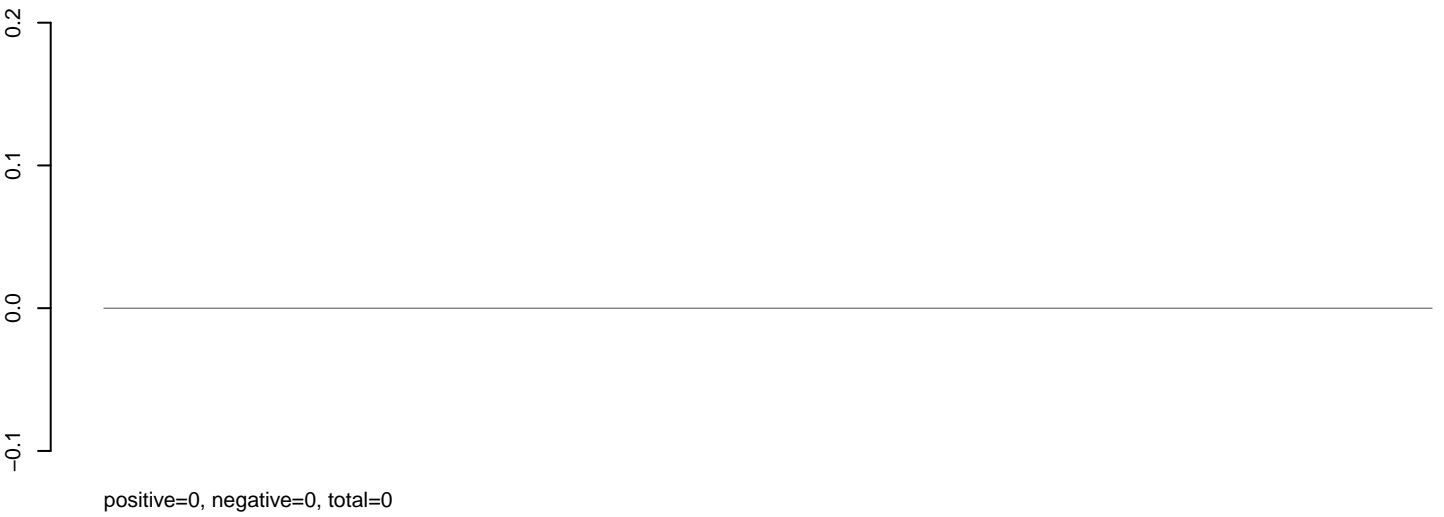
Window size=25, length=10600, virus@NC_003635.1-Modoc_virus:1-10600

0 2000 4000 6000 8000 10000

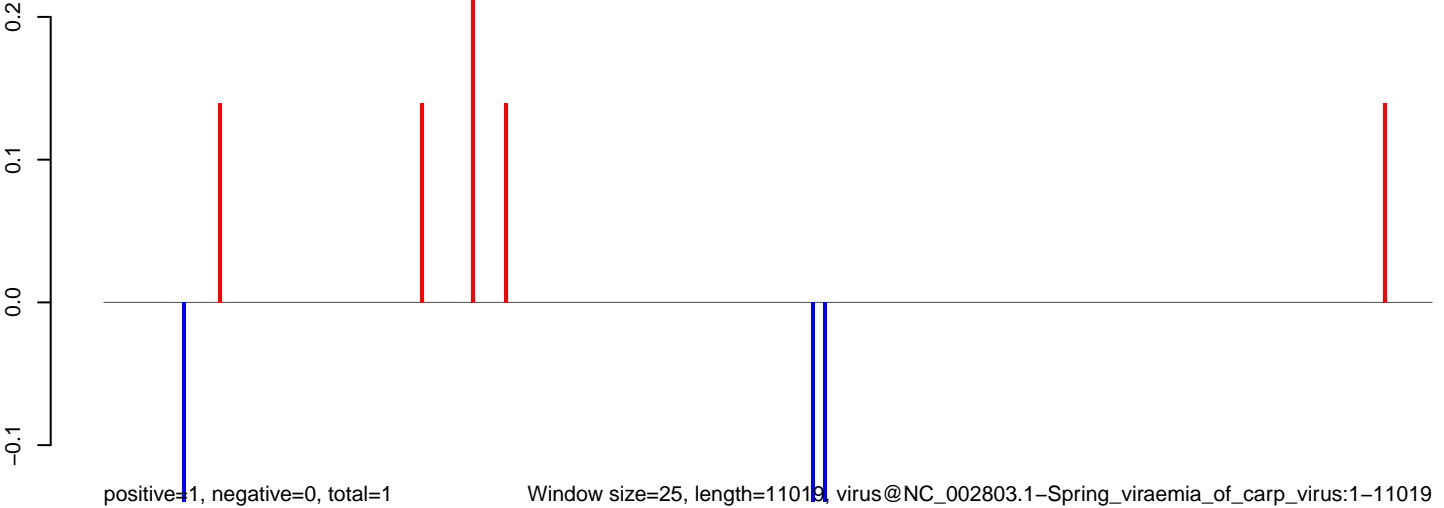
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



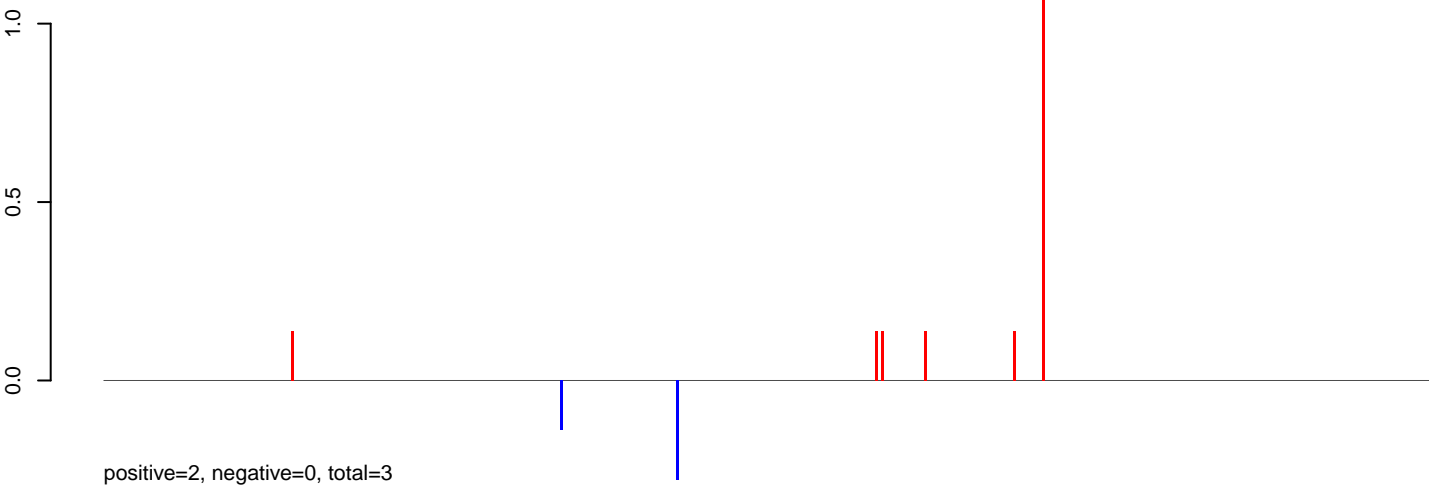
AeAlbo_C636_2dpi_mock3.rep



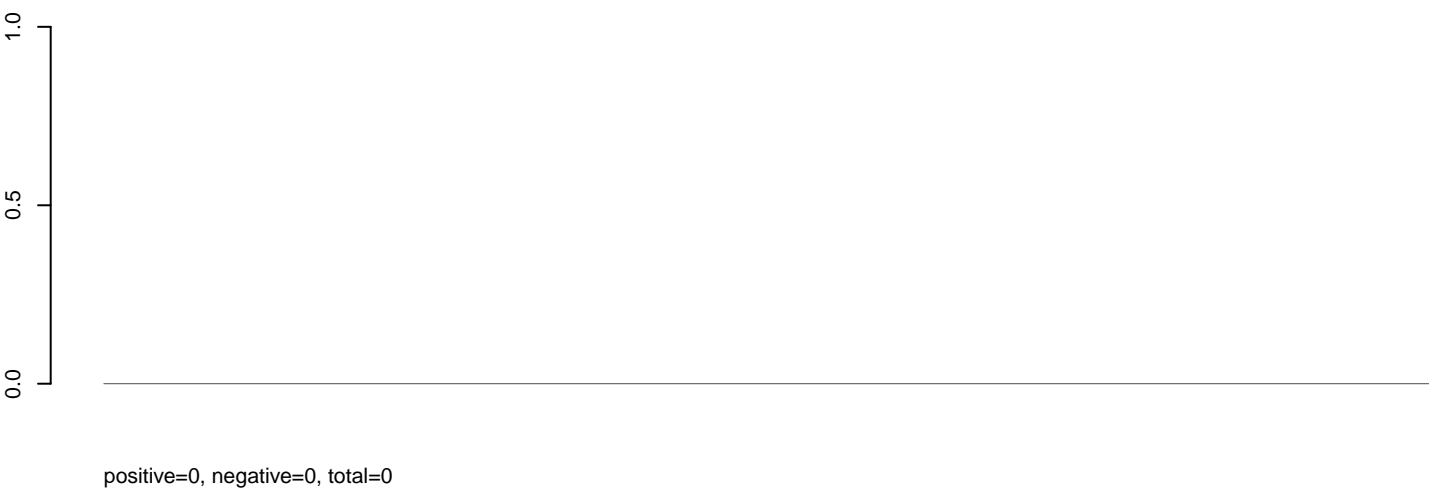
Window size=25, length=11019, virus@NC_002803.1-Spring_viraemia_of_carp_virus:1-11019

0 2000 4000 6000 8000 10000

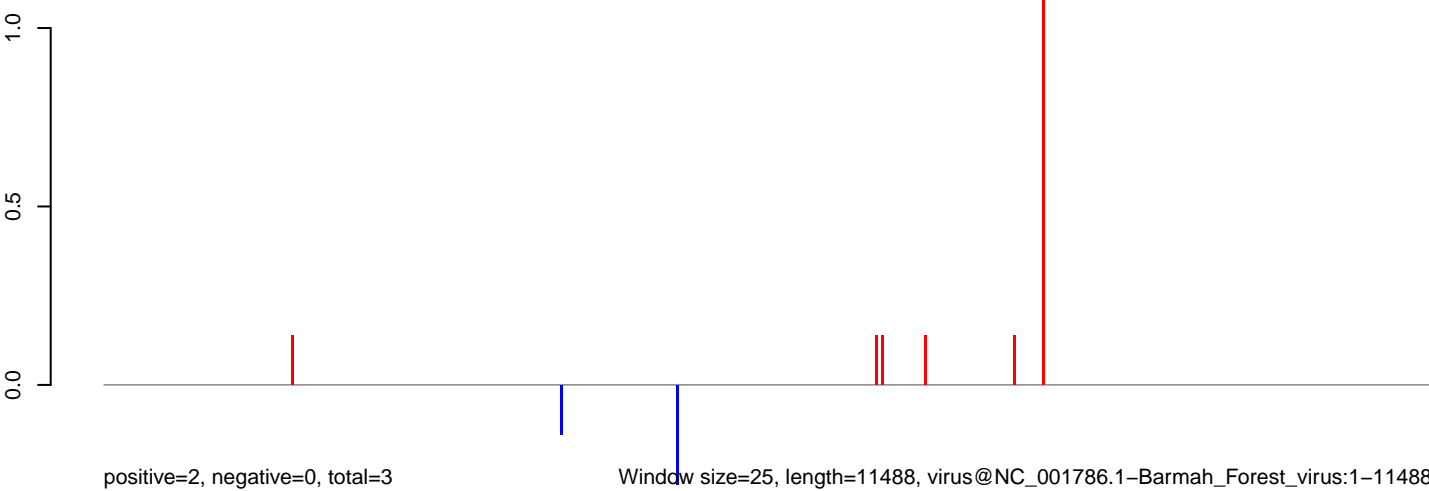
AeAlbo_C636_2dpi_mock3.18_23.rep



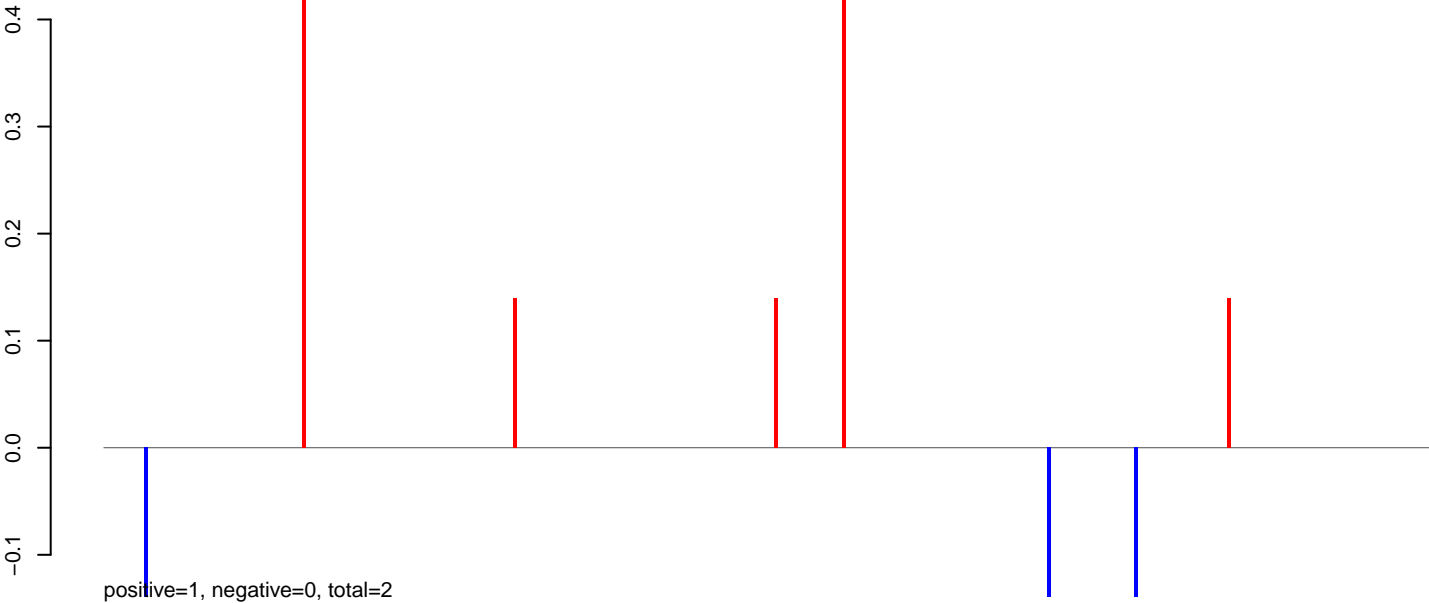
AeAlbo_C636_2dpi_mock3.24_35.rep



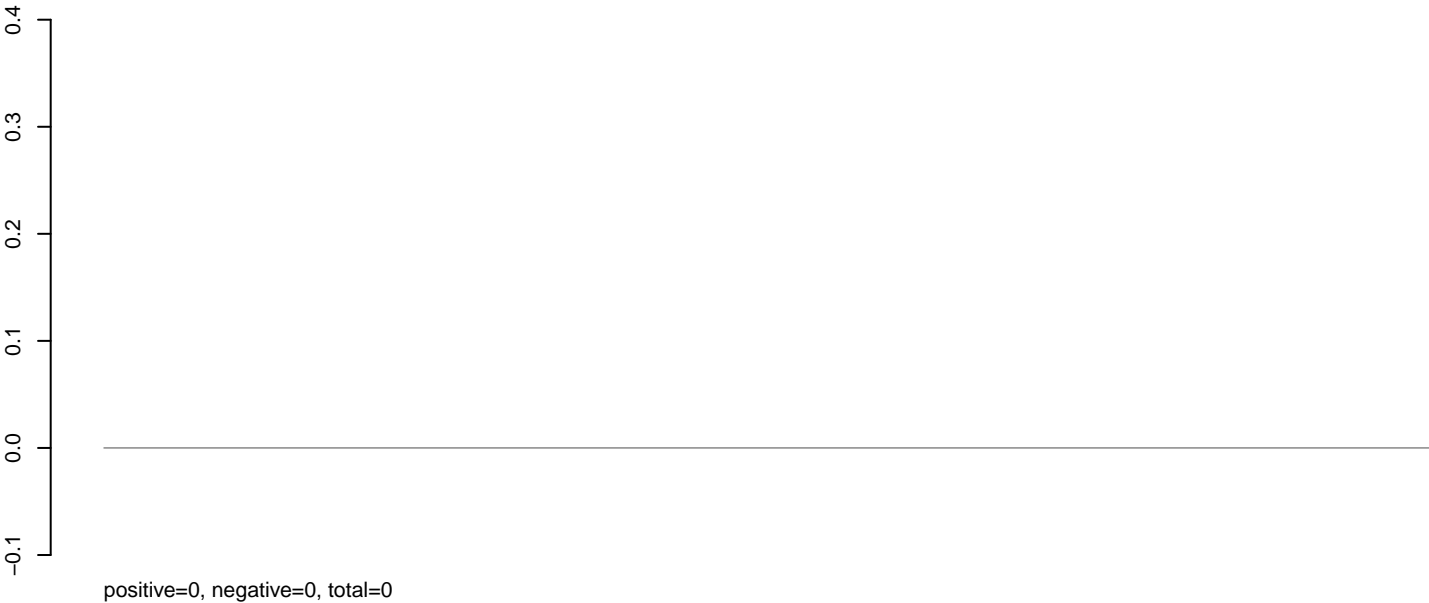
AeAlbo_C636_2dpi_mock3.rep



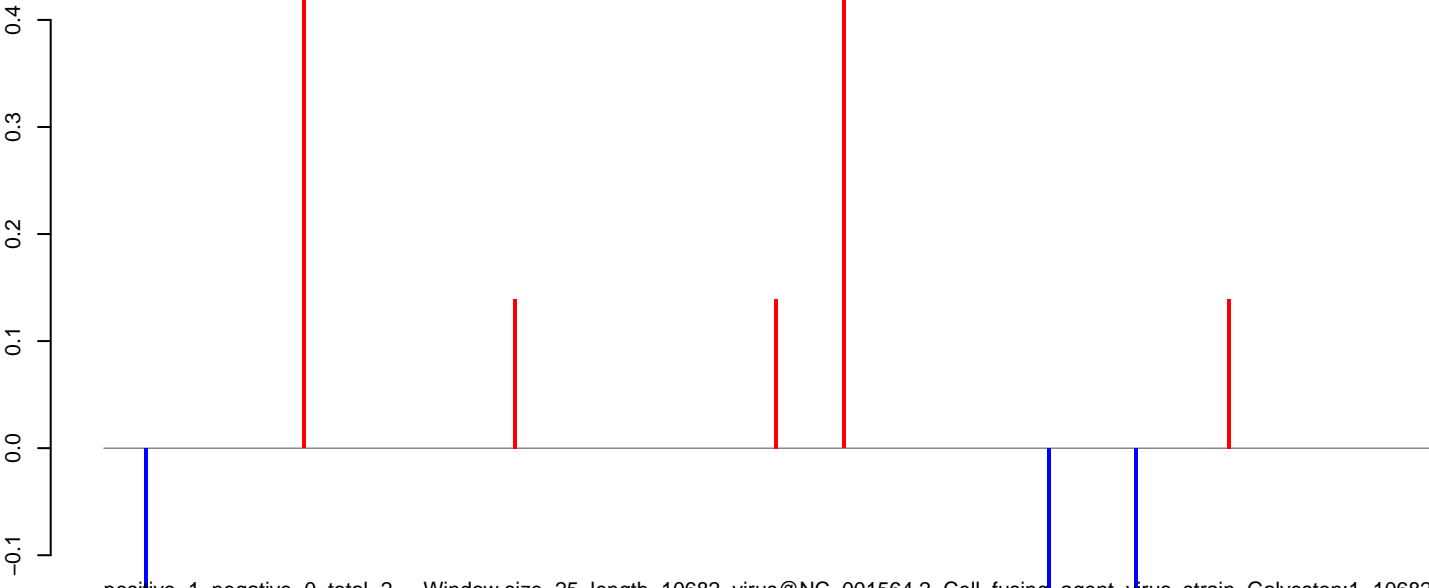
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



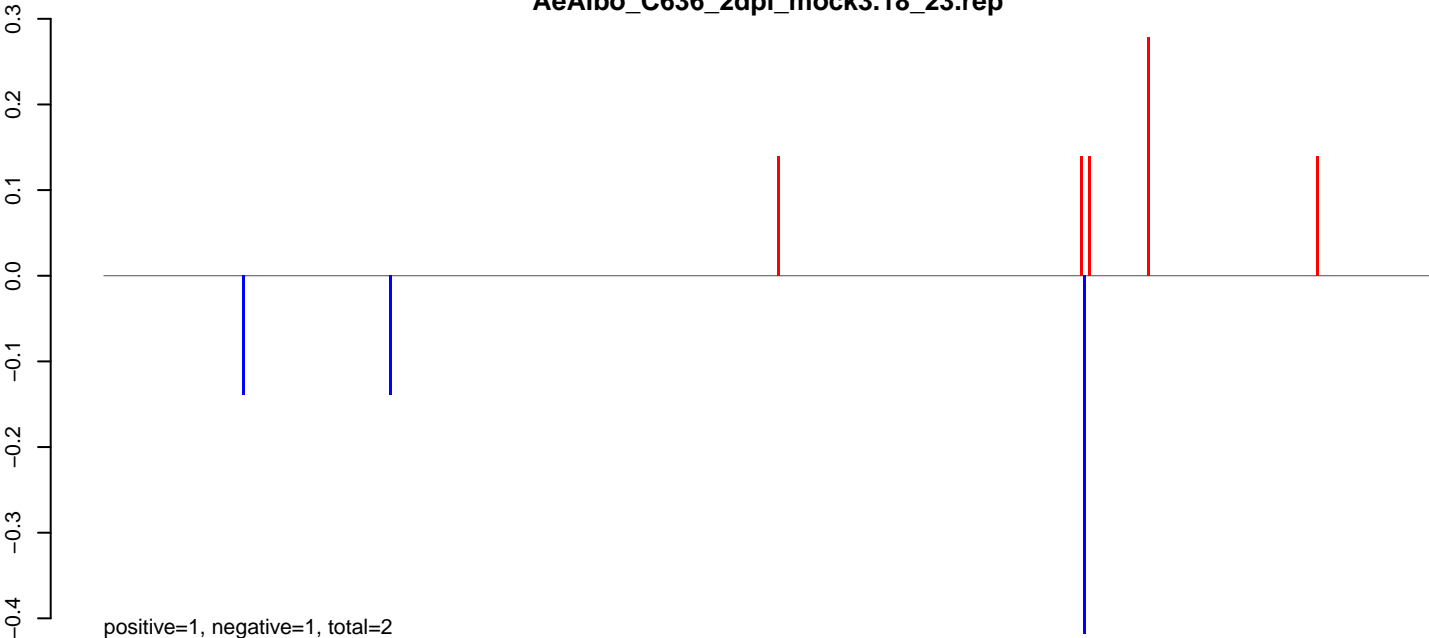
AeAlbo_C636_2dpi_mock3.rep



Window size=25, length=10682, virus@NC_001564.2-Cell_fusing_agent_virus_strain_Galveston:1-10682

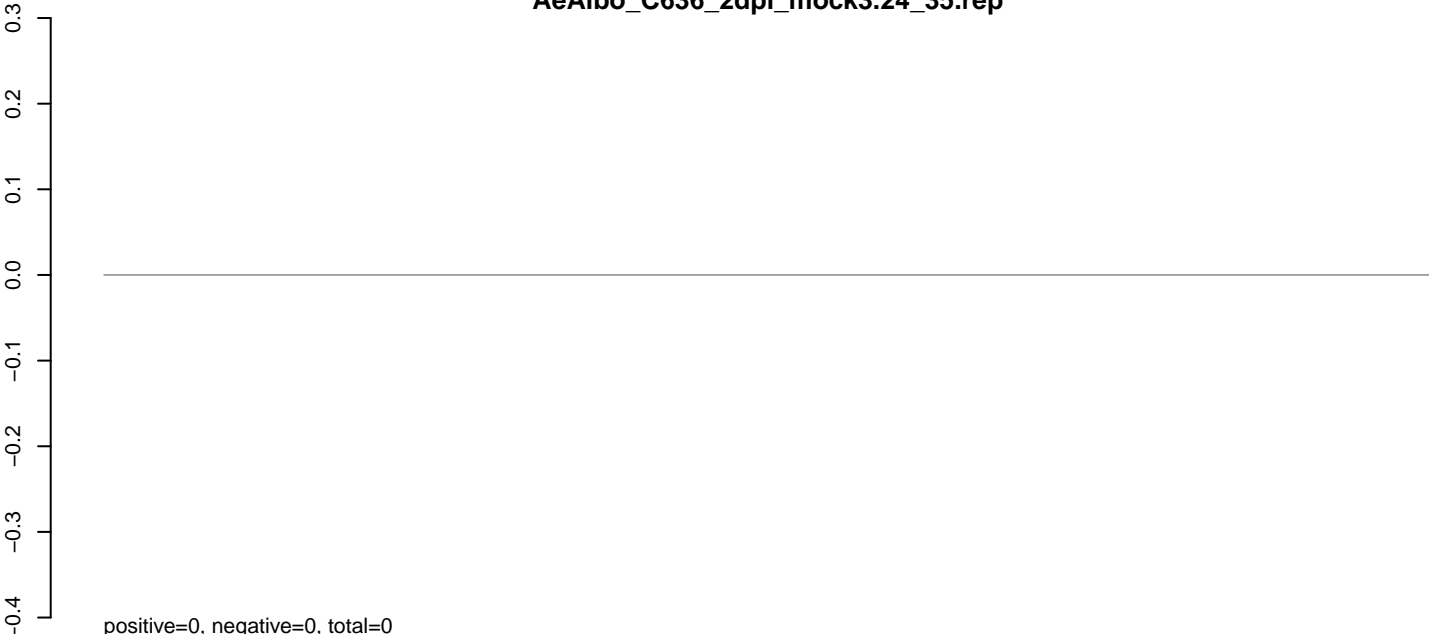
0 2000 4000 6000 8000 10000

AeAlbo_C636_2dpi_mock3.18_23.rep



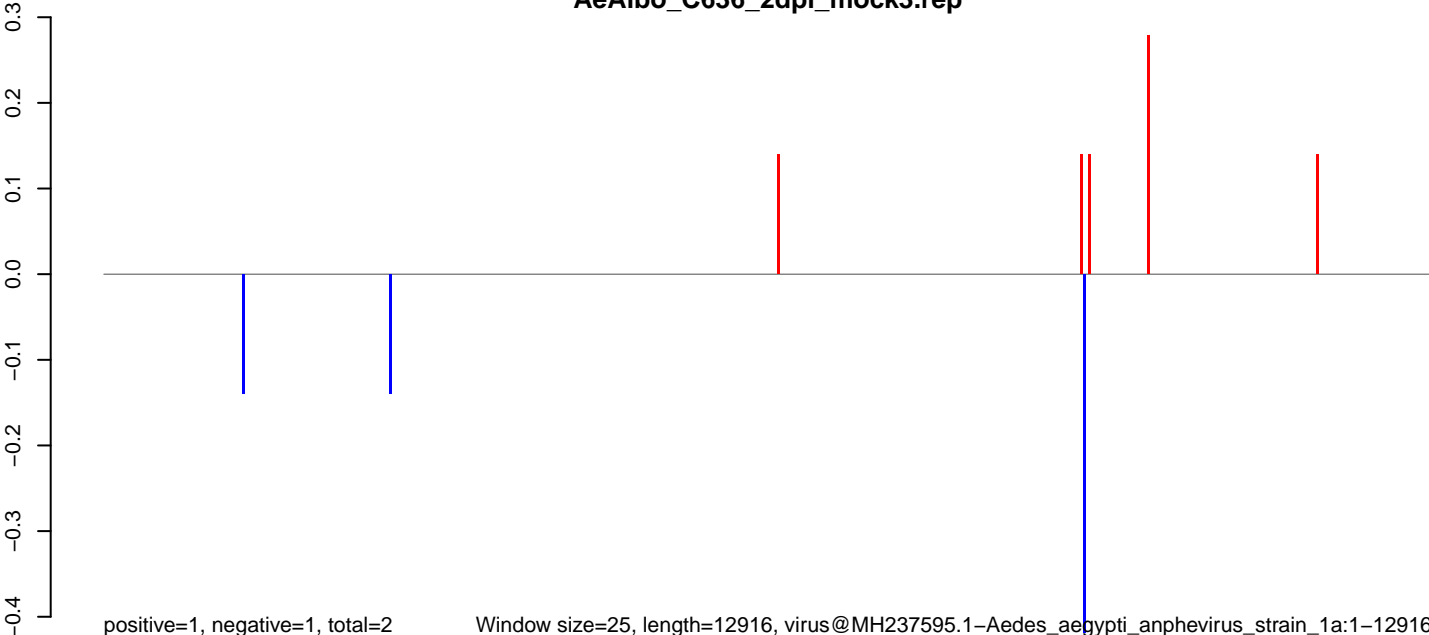
positive=1, negative=1, total=2

AeAlbo_C636_2dpi_mock3.24_35.rep



positive=0, negative=0, total=0

AeAlbo_C636_2dpi_mock3.rep

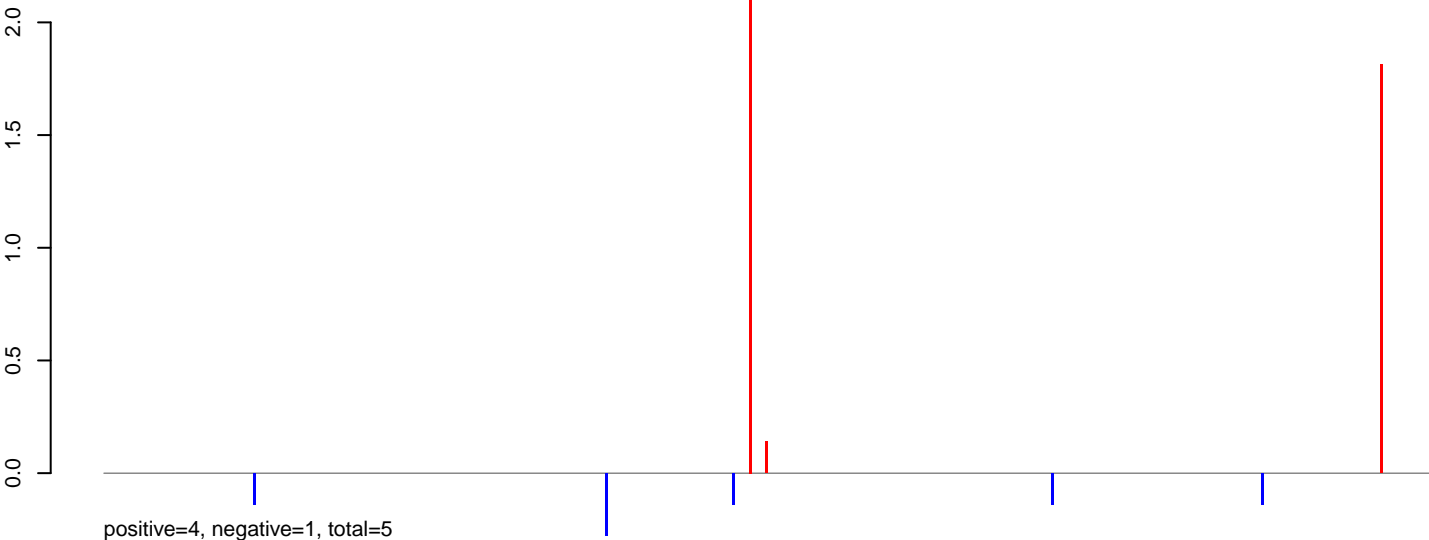


positive=1, negative=1, total=2

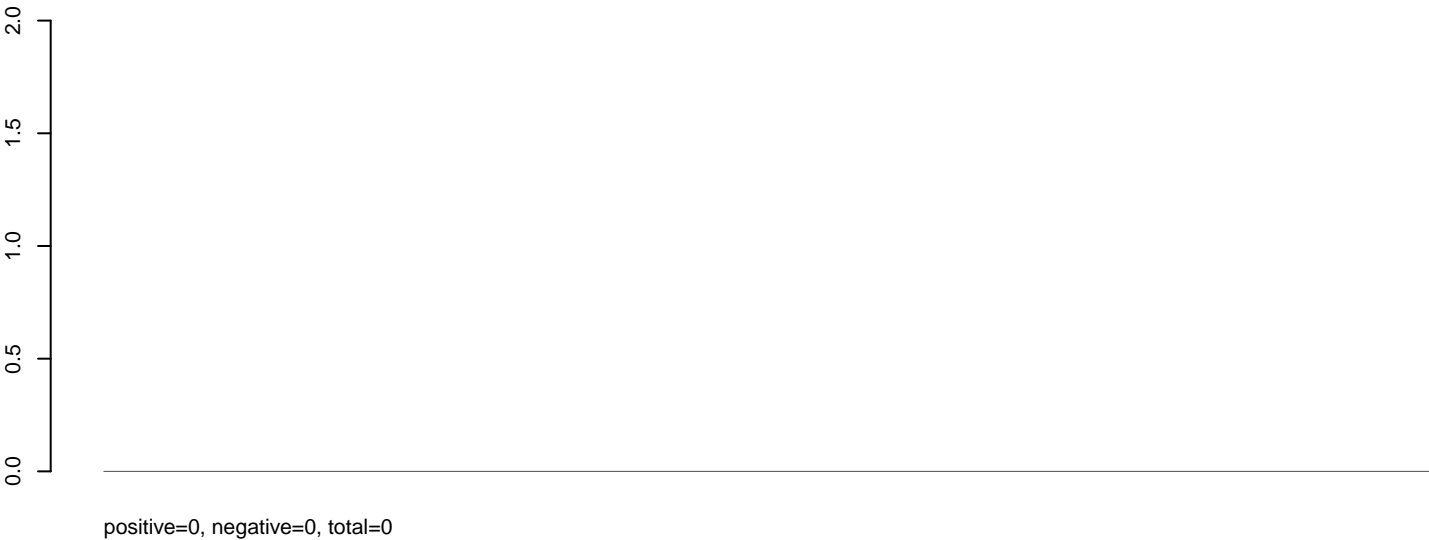
Window size=25, length=12916, virus@MH237595.1-Aedes_aegypti_anphevirus_strain_1a:1-12916

0 2000 4000 6000 8000 10000 12000

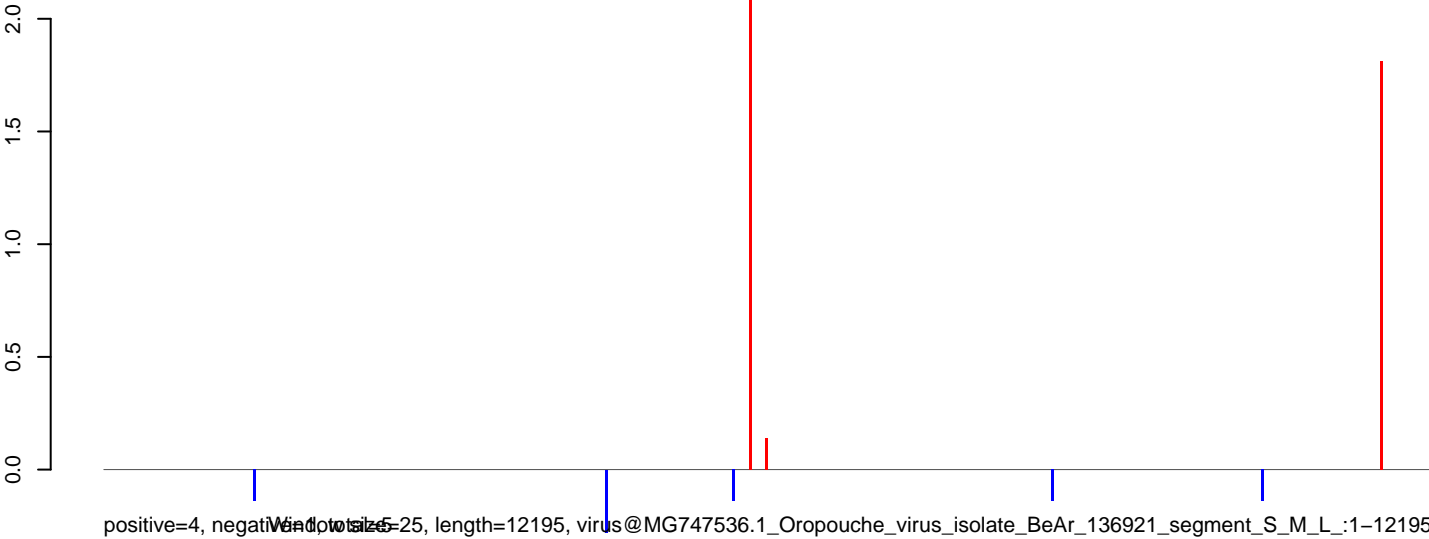
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

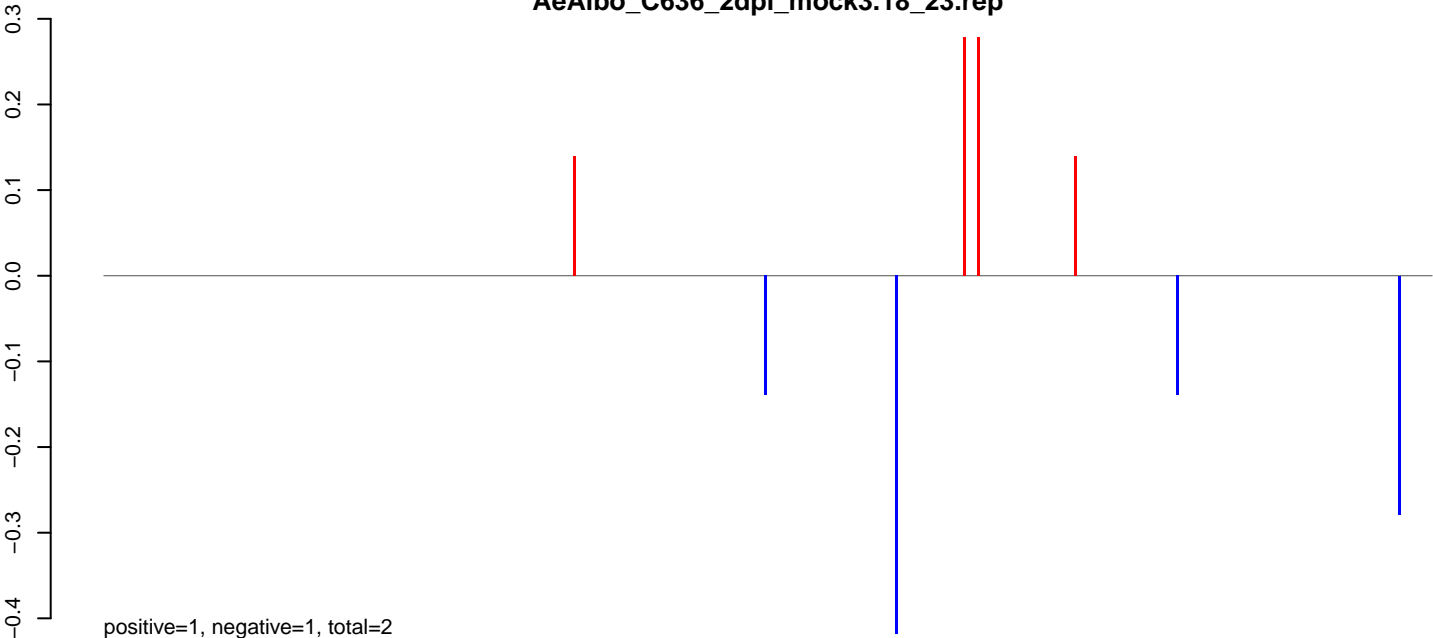


AeAlbo_C636_2dpi_mock3.rep

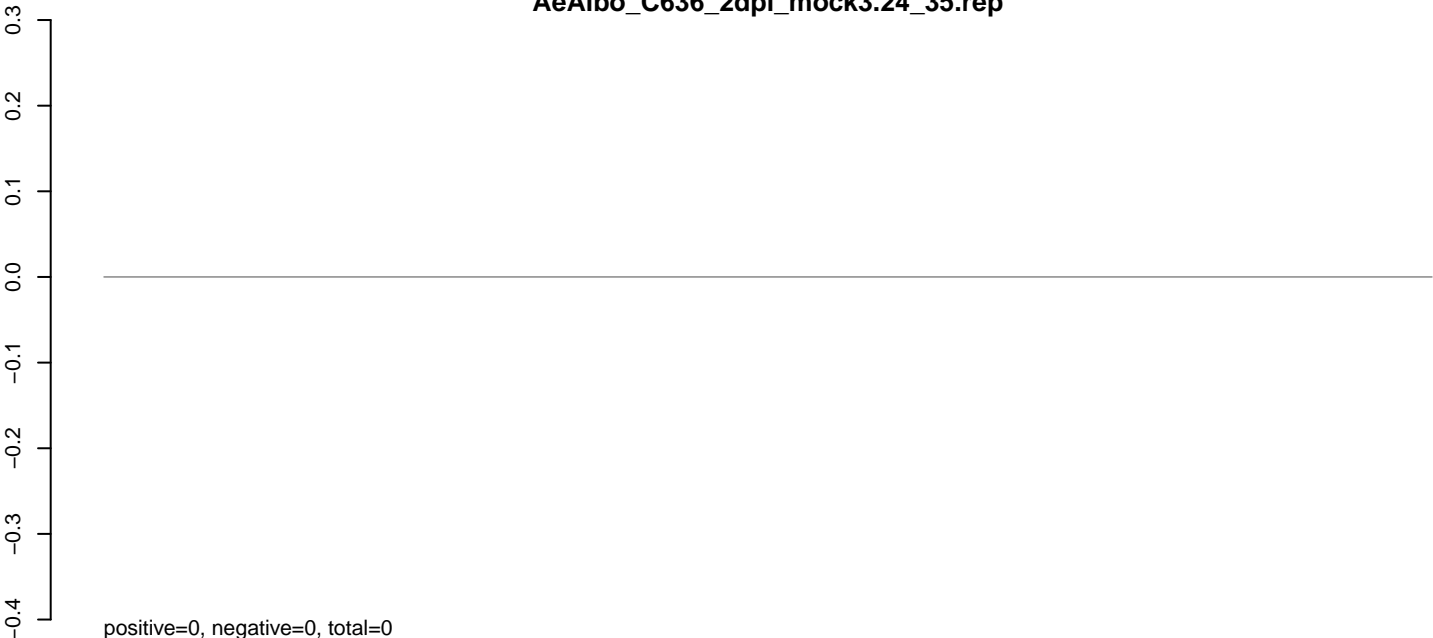


Window size=25, length=12195, virus@MG747536.1_Oropouche_virus_isolate_BeAr_136921_segment_S_M_L:1-12195

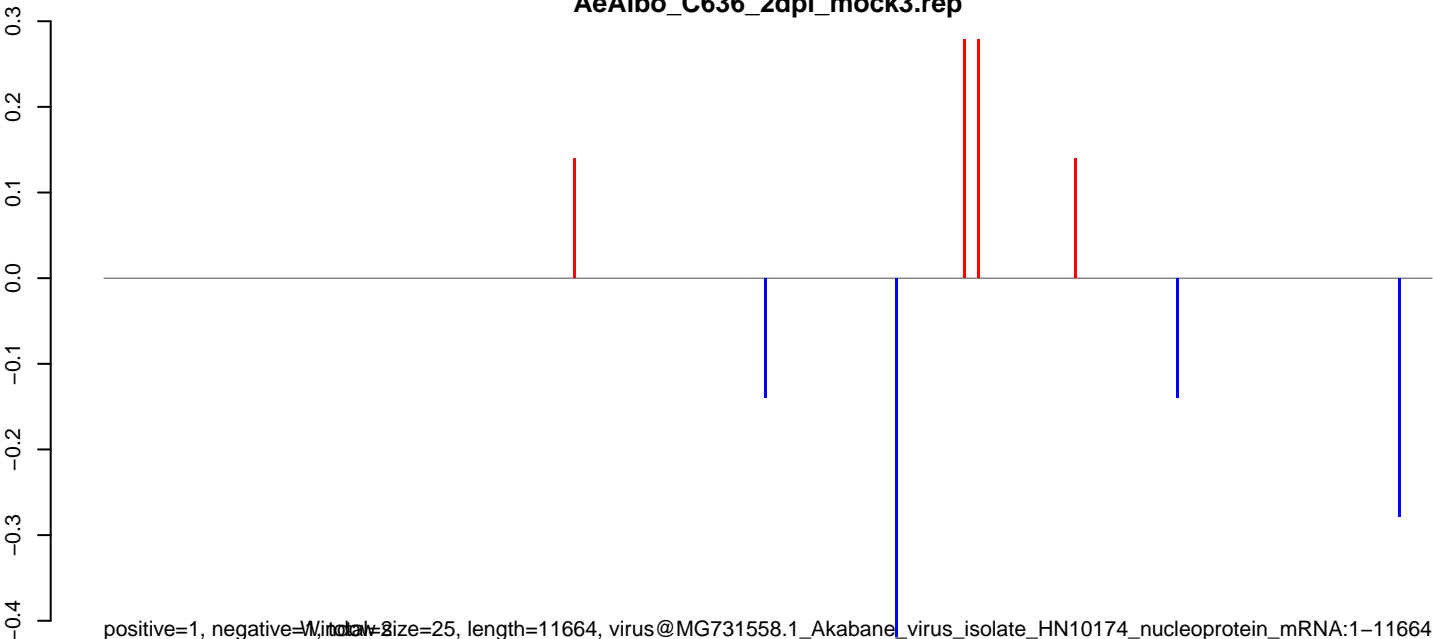
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



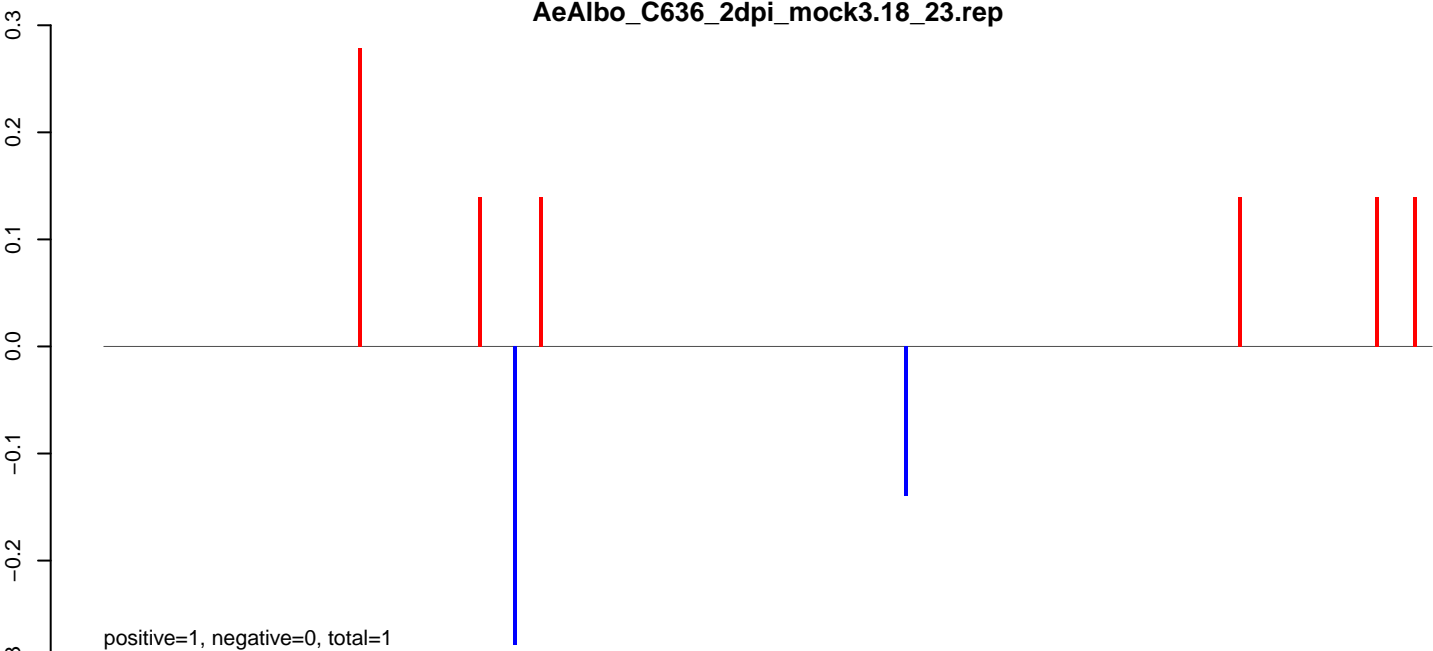
AeAlbo_C636_2dpi_mock3.rep



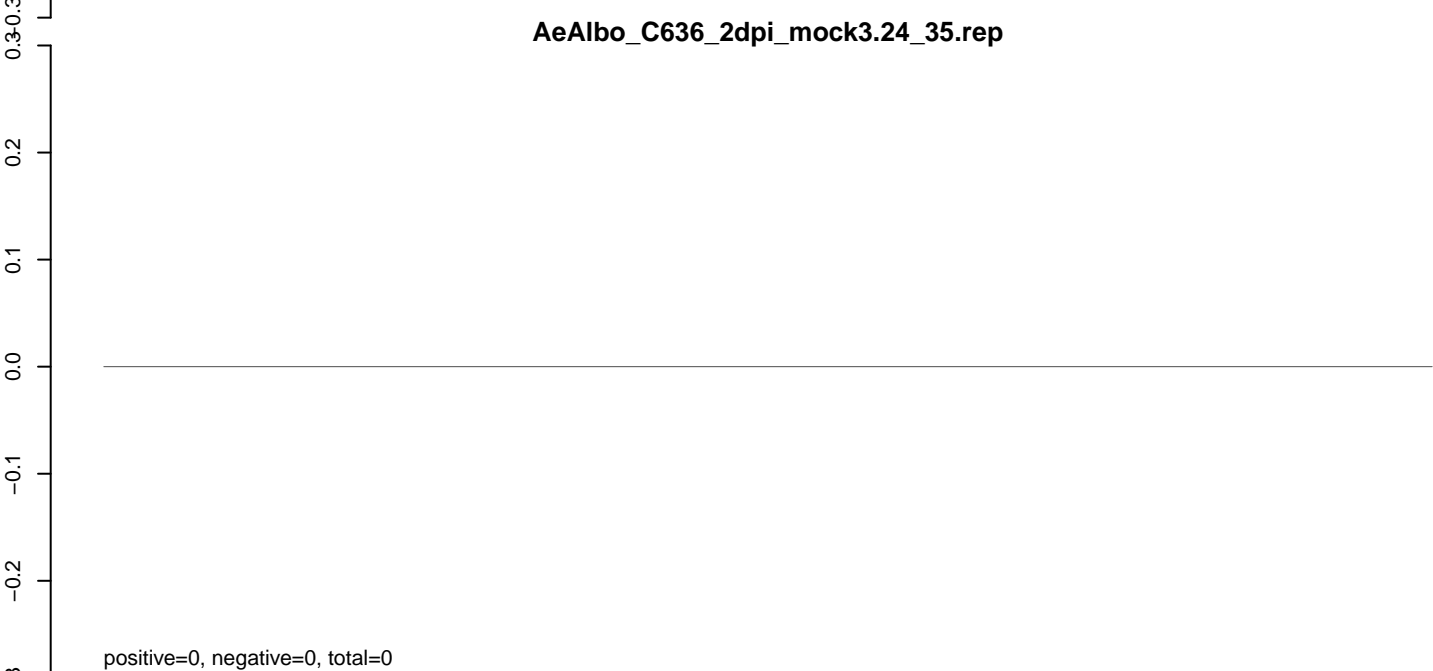
Window size=25, length=11664, virus@MG731558.1_Akabane_virus_isolate_HN10174_nucleoprotein_mRNA:1-11664

0 2000 4000 6000 8000 10000 12000

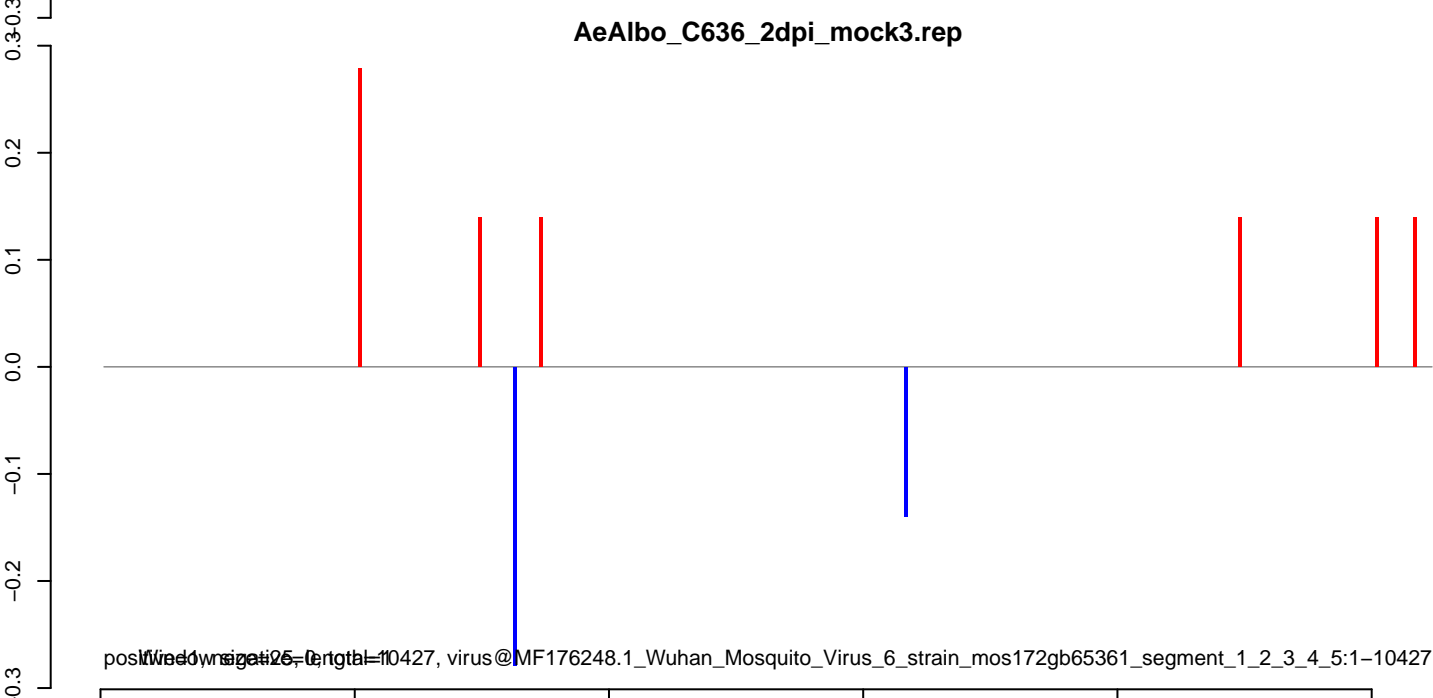
AeAlbo_C636_2dpi_mock3.18_23.rep



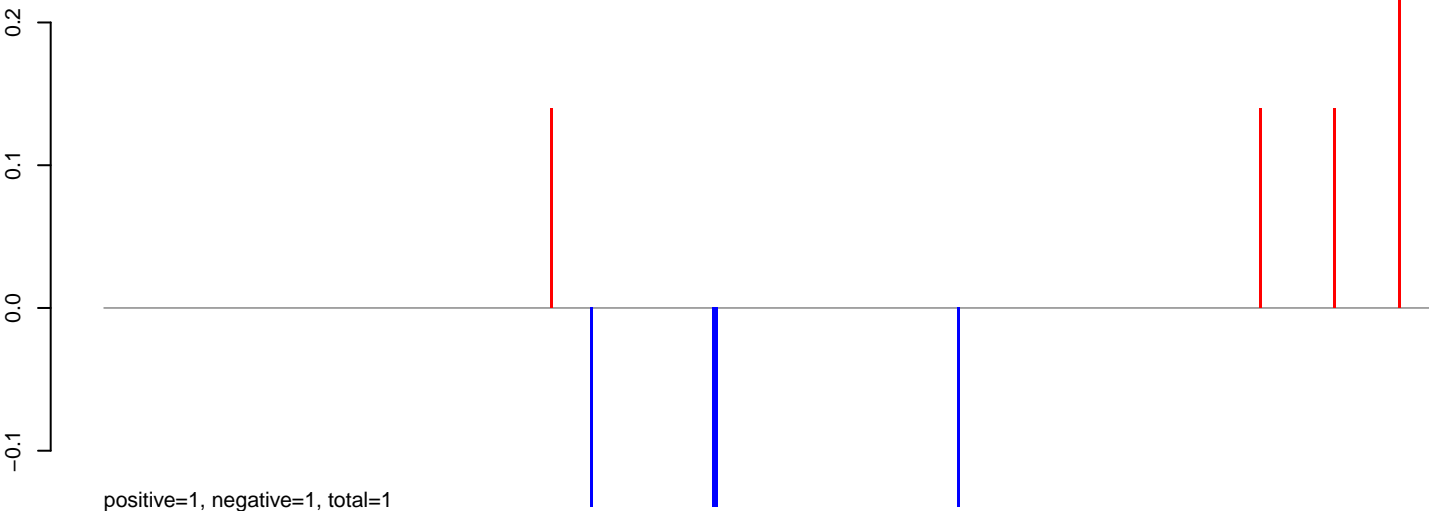
AeAlbo_C636_2dpi_mock3.24_35.rep



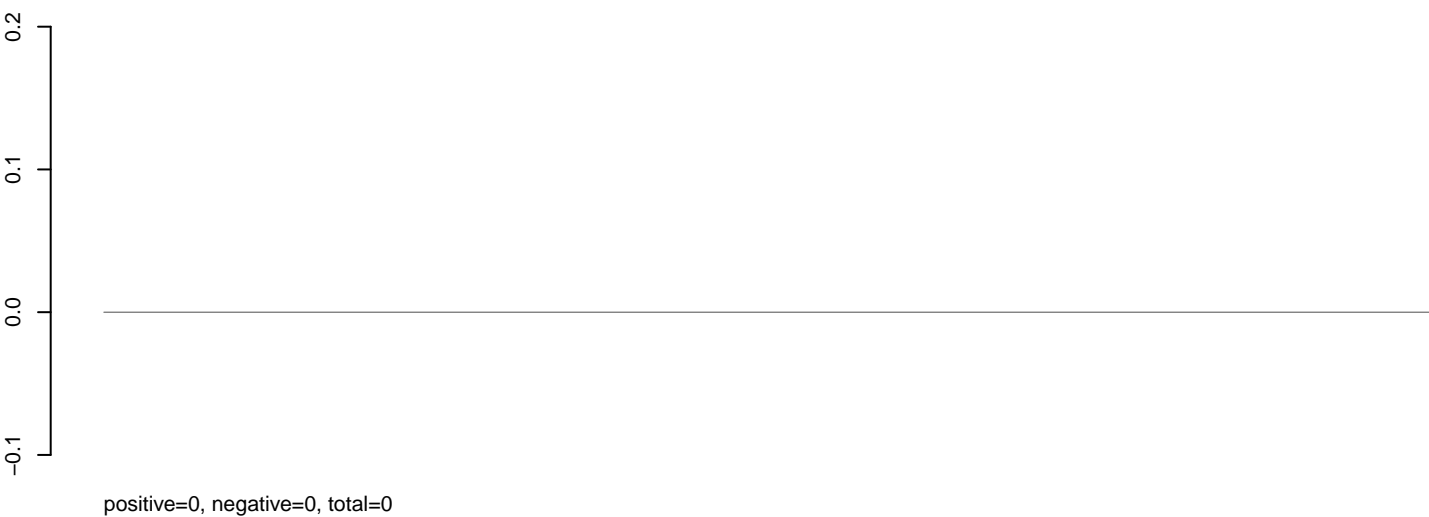
AeAlbo_C636_2dpi_mock3.rep



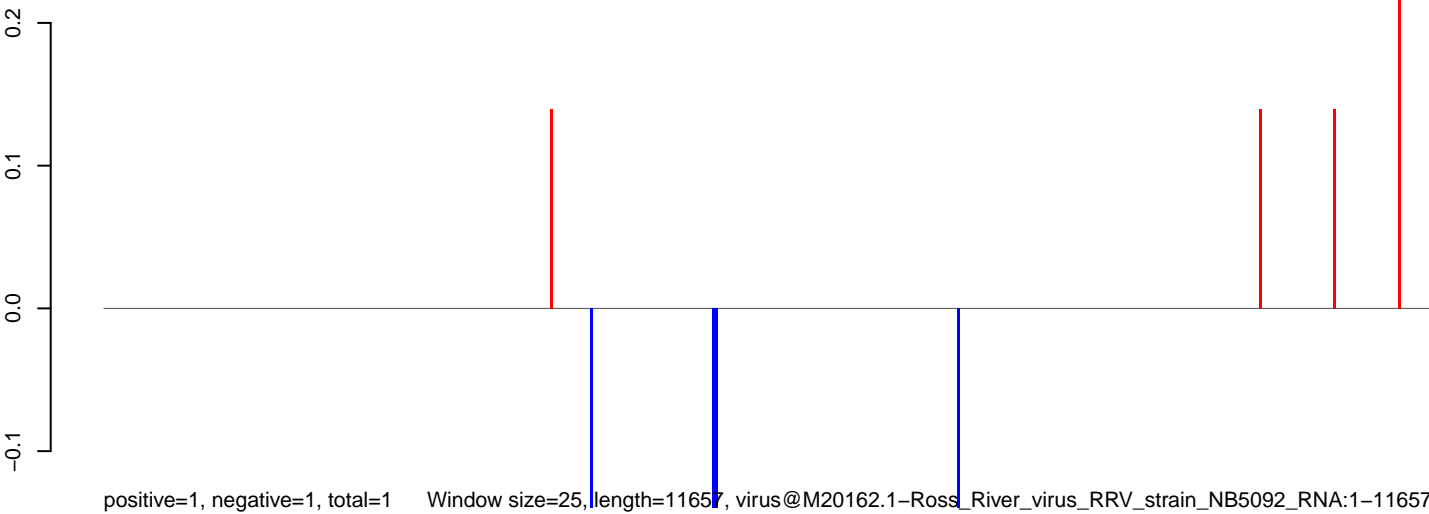
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



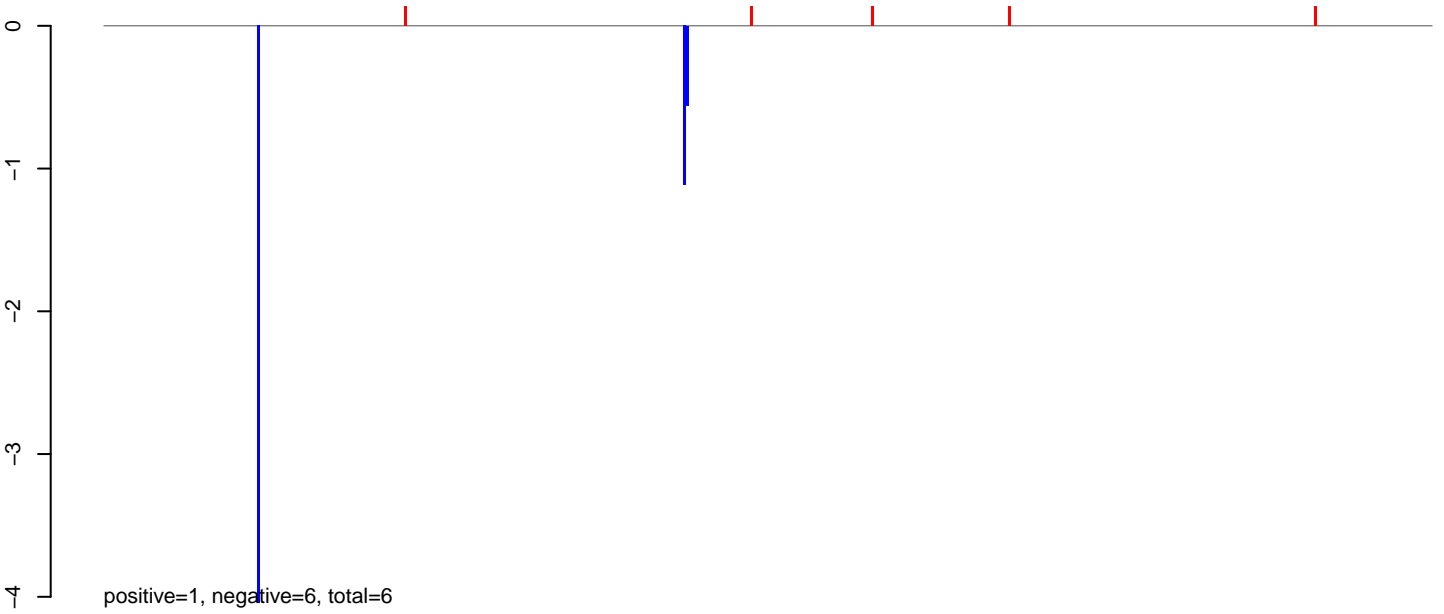
AeAlbo_C636_2dpi_mock3.rep



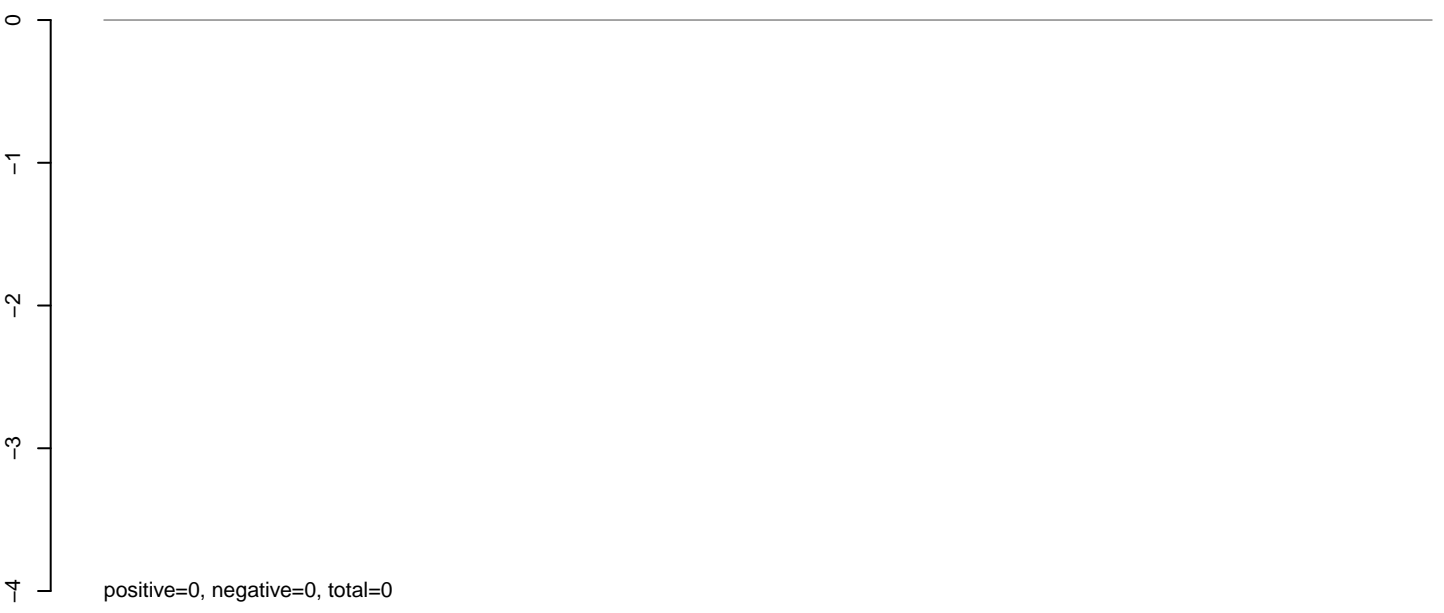
Window size=25, length=11657, virus@M20162.1-Ross_River_virus_RRV_strain_NB5092_RNA:1-11657

0 2000 4000 6000 8000 10000 12000

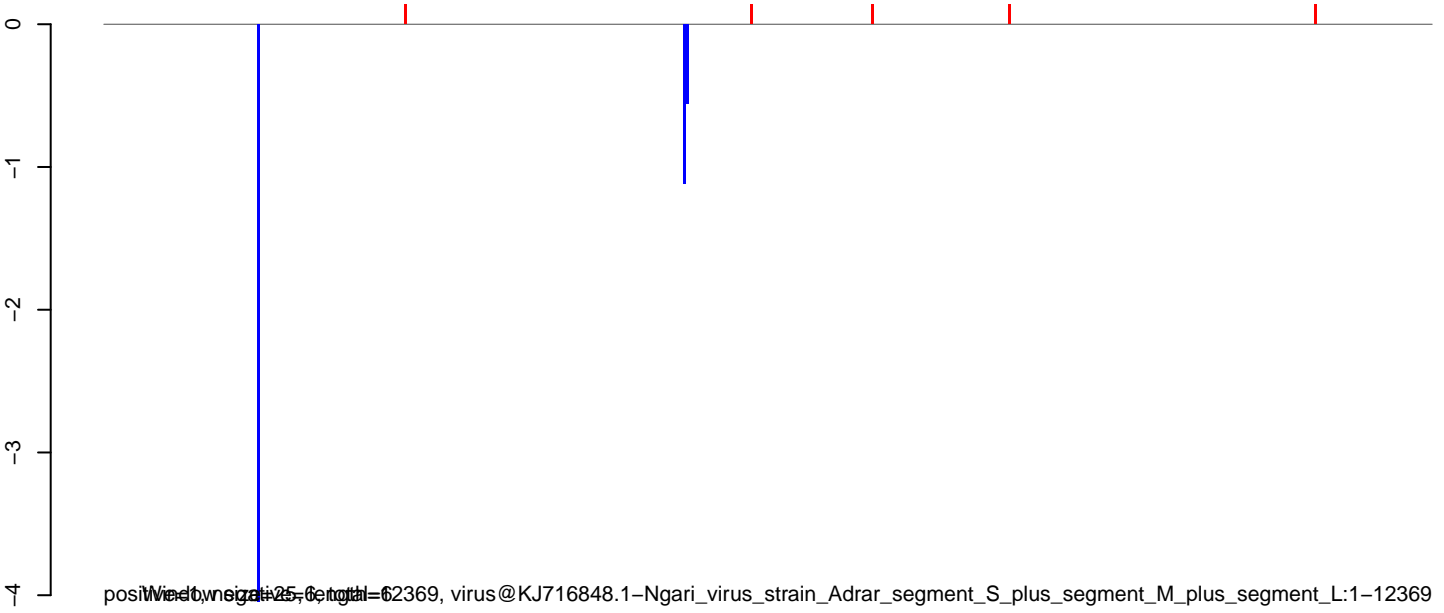
AeAlbo_C636_2dpi_mock3.18_23.rep



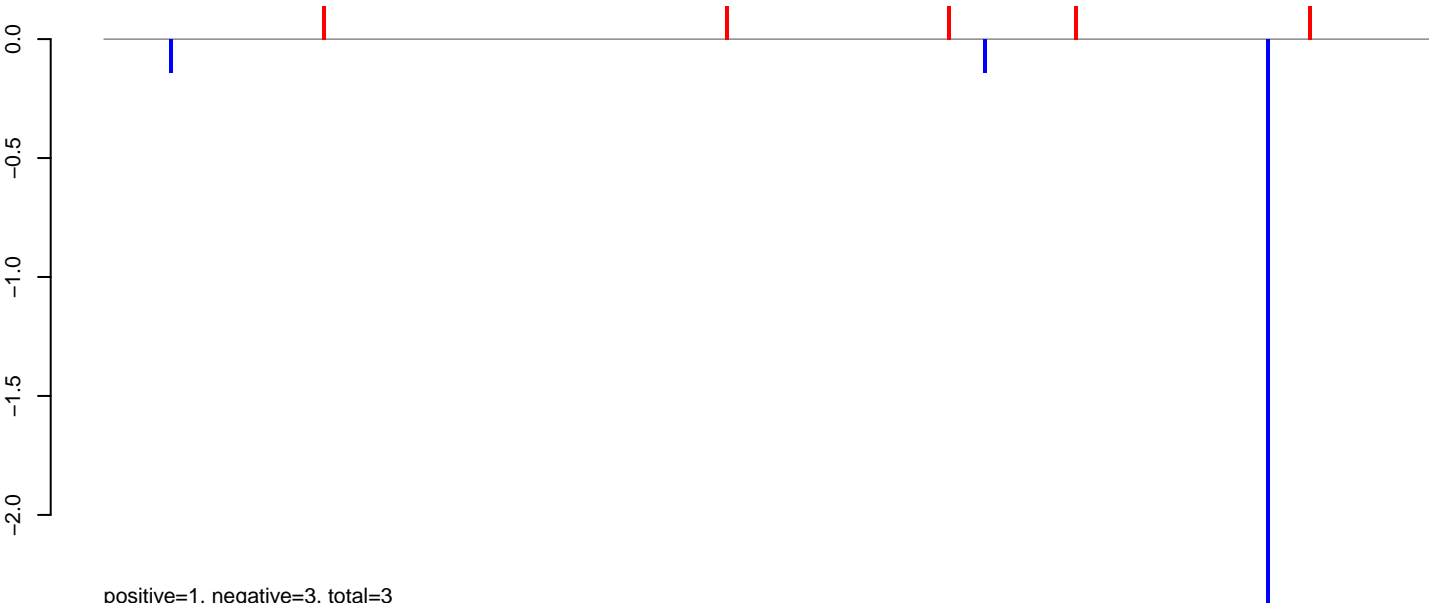
AeAlbo_C636_2dpi_mock3.24_35.rep



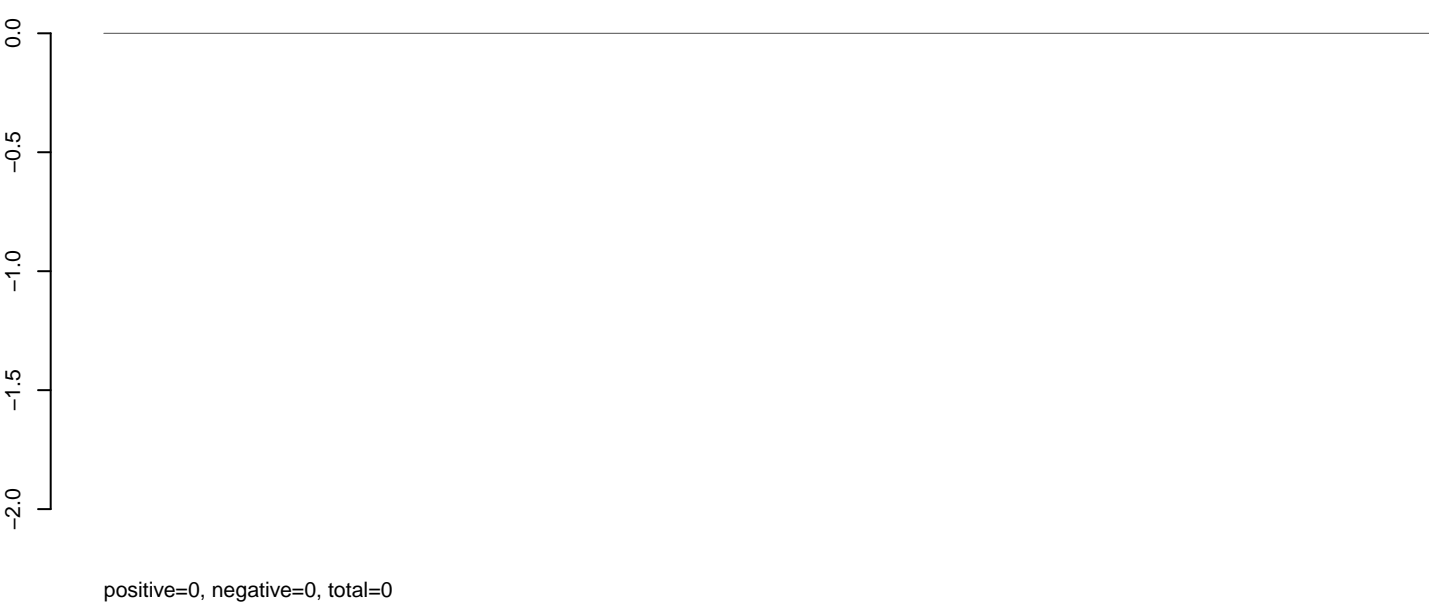
AeAlbo_C636_2dpi_mock3.rep



AeAlbo_C636_2dpi_mock3.18_23.rep



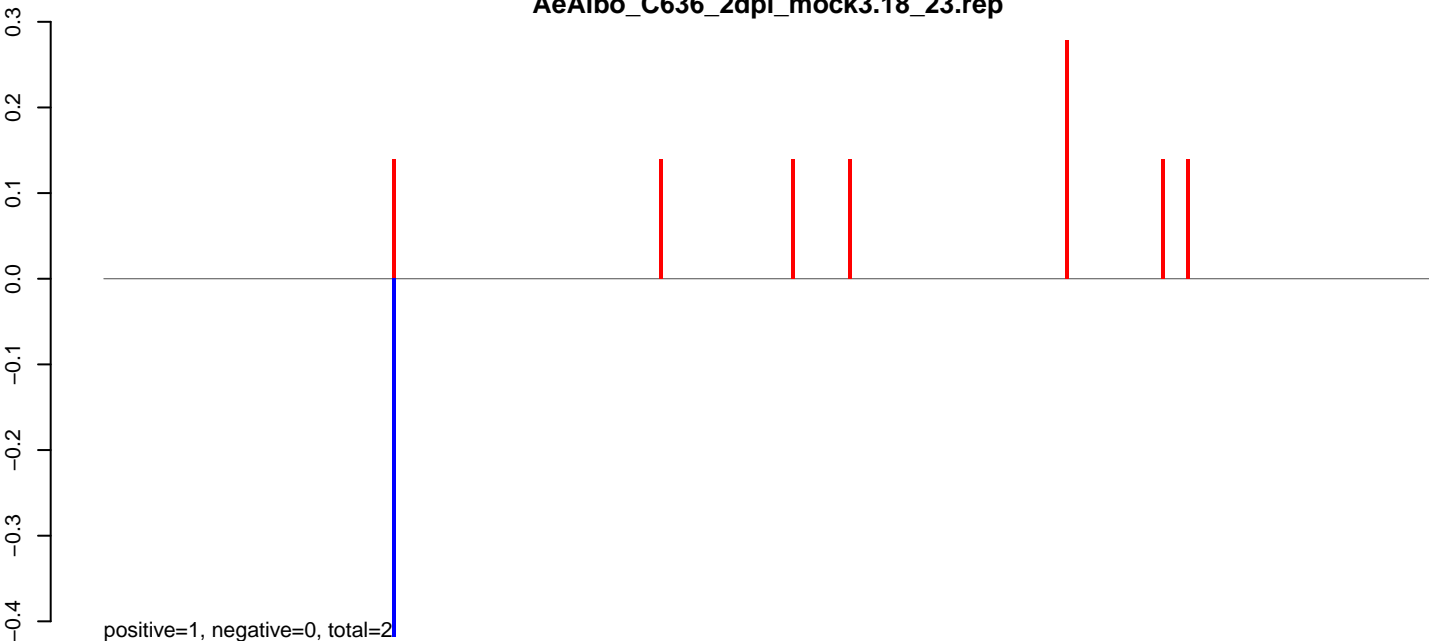
AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep

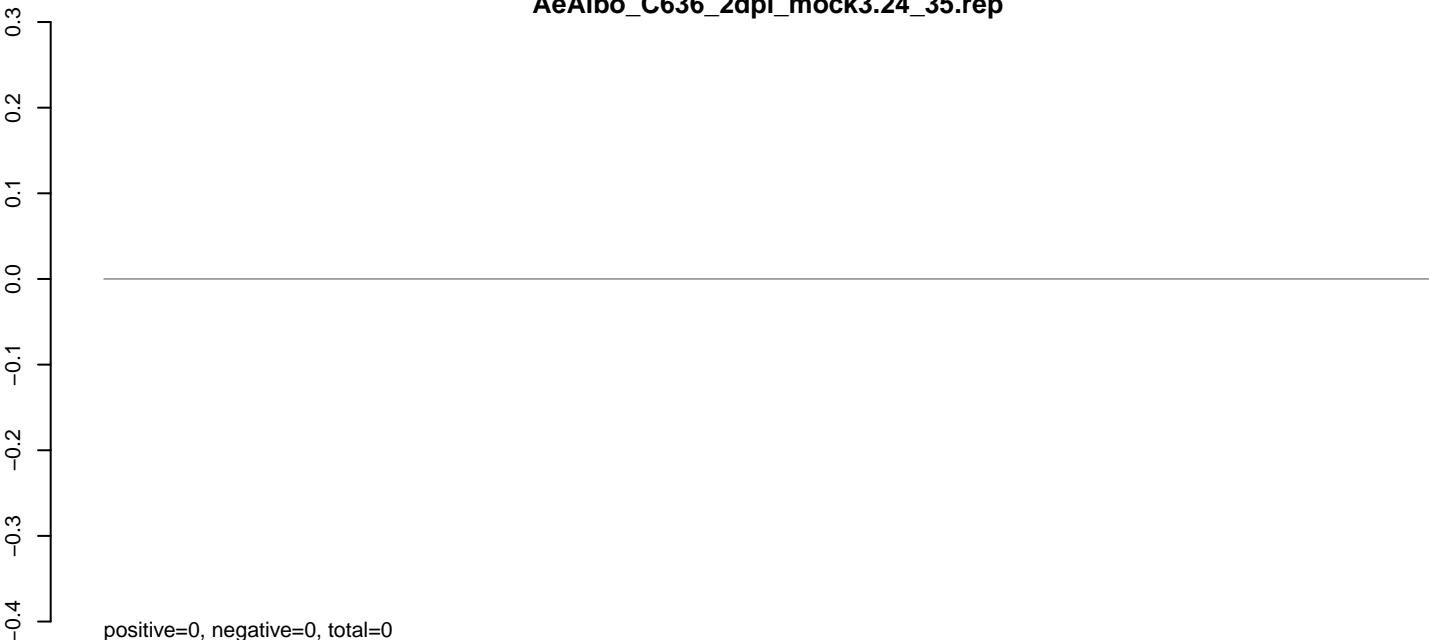


AeAlbo_C636_2dpi_mock3.18_23.rep



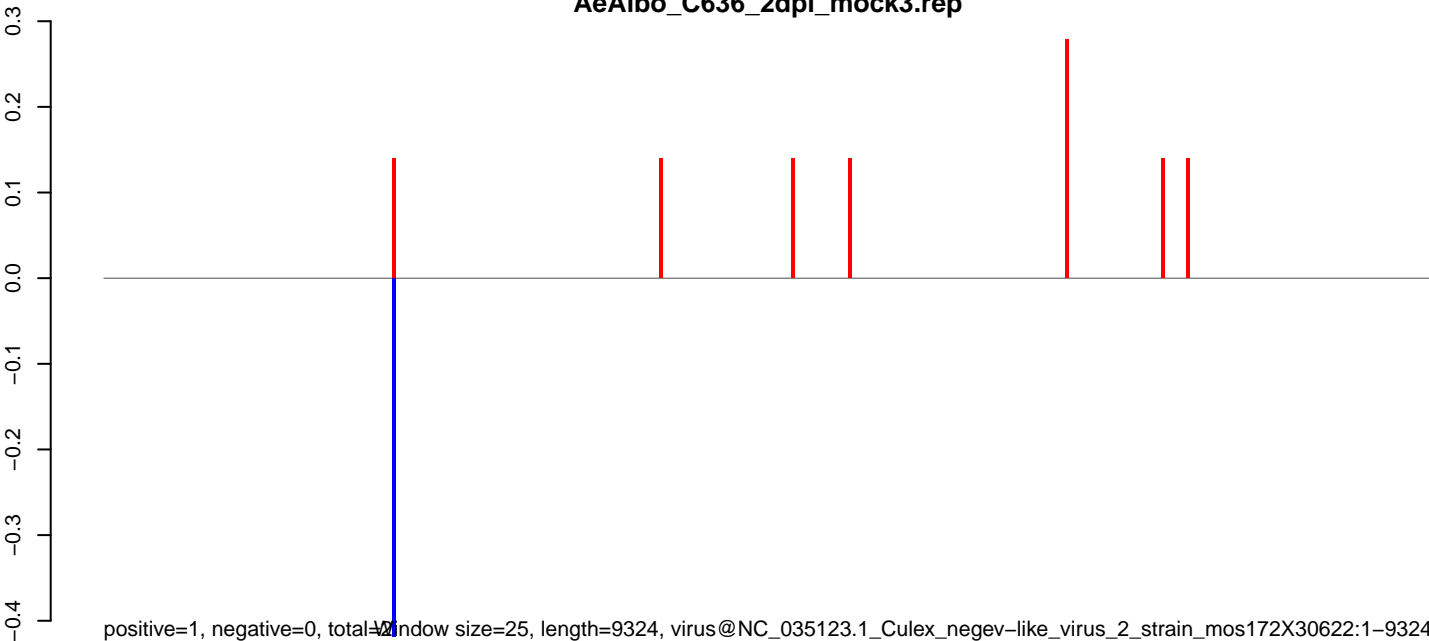
positive=1, negative=0, total=2

AeAlbo_C636_2dpi_mock3.24_35.rep



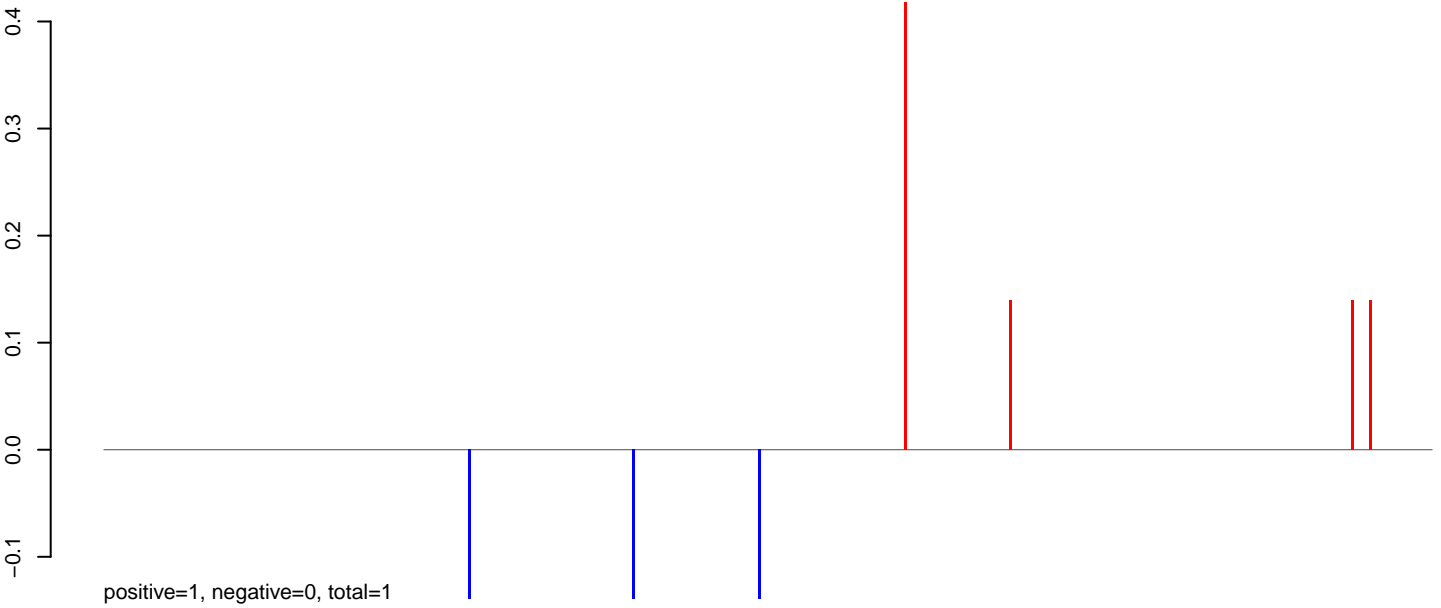
positive=0, negative=0, total=0

AeAlbo_C636_2dpi_mock3.rep



positive=1, negative=0, total=1, window size=25, length=9324, virus@NC_035123.1_Culex_negev-like_virus_2_strain_mos172X30622:1-9324

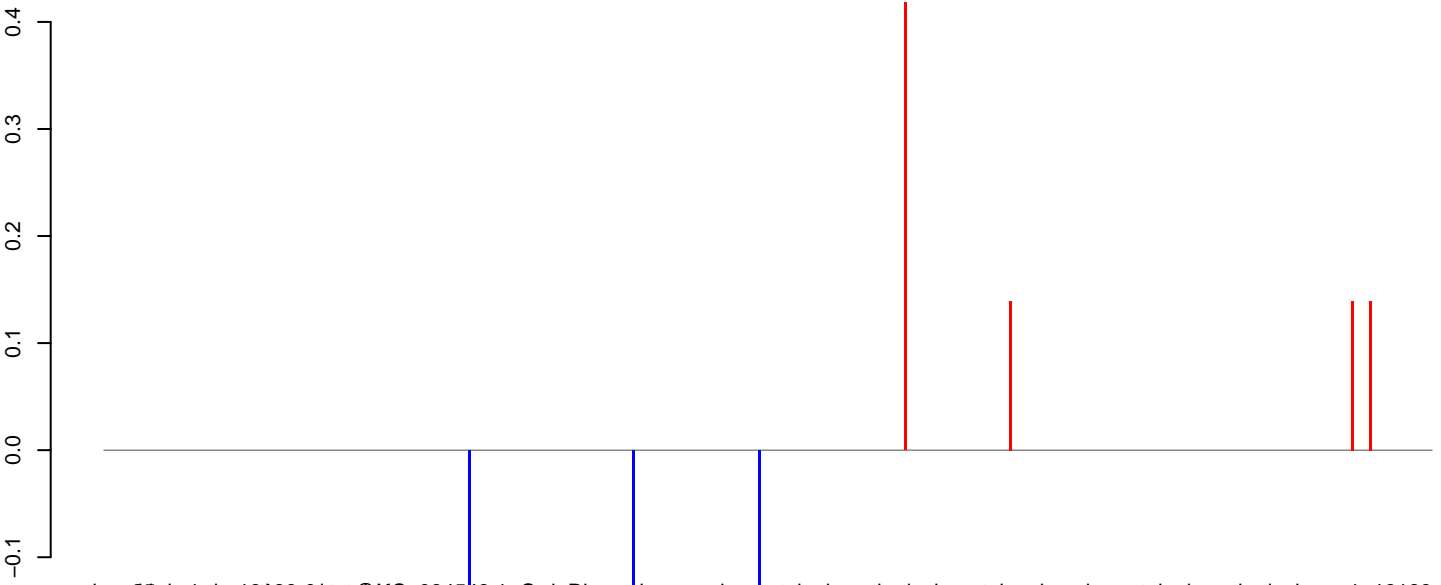
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



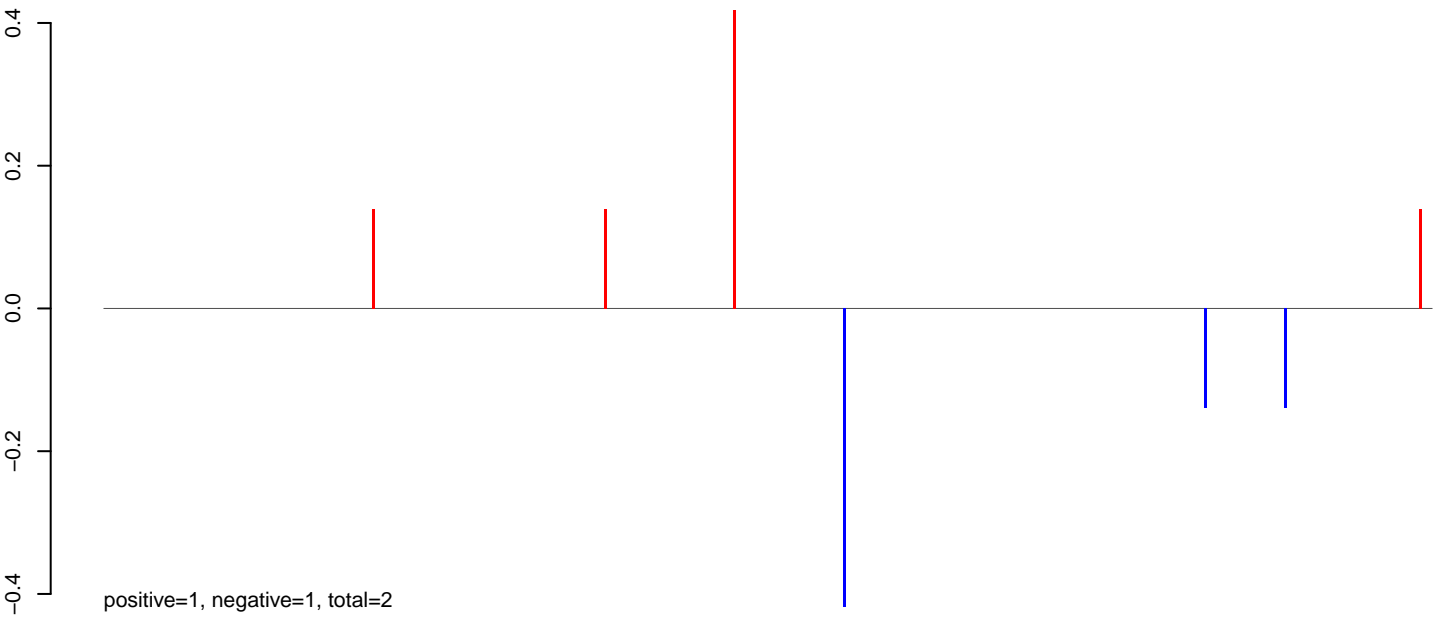
AeAlbo_C636_2dpi_mock3.rep



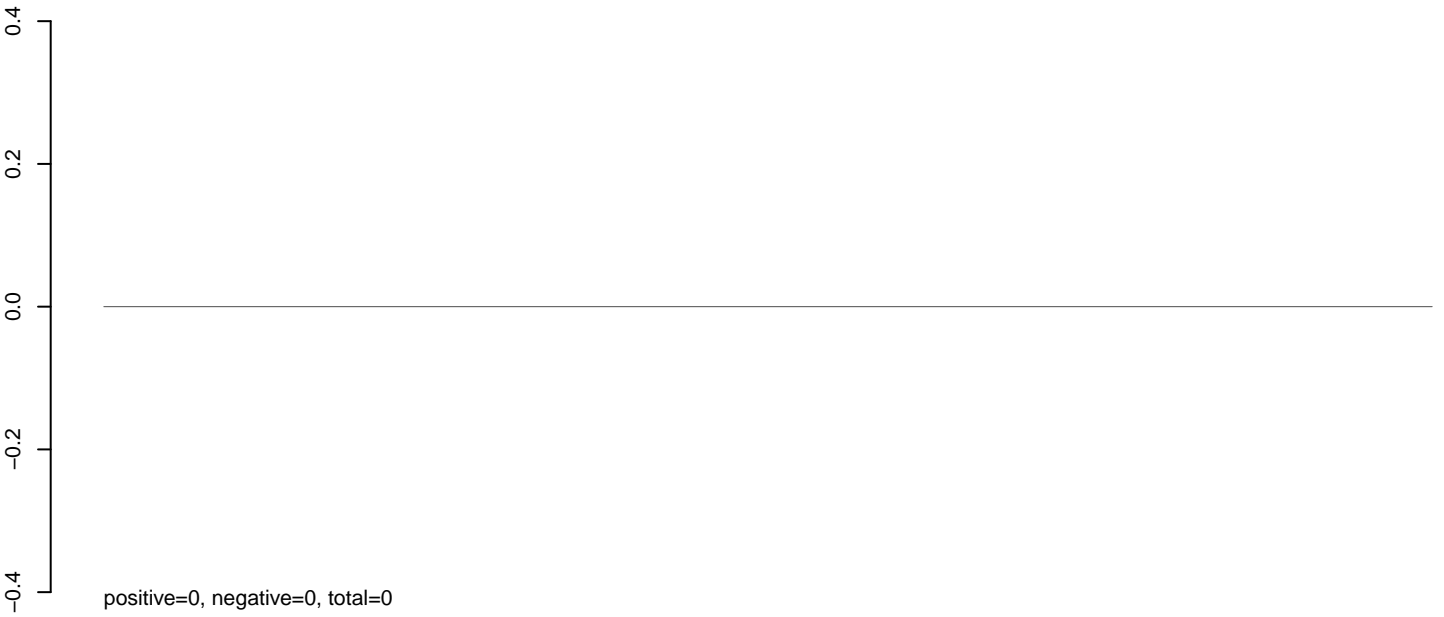
NC_034543.1-Ord_River_virus_nucleoprotein_hypothetical_protein_hypothetical_pro:1-13189

0 2000 4000 6000 8000 10000 12000

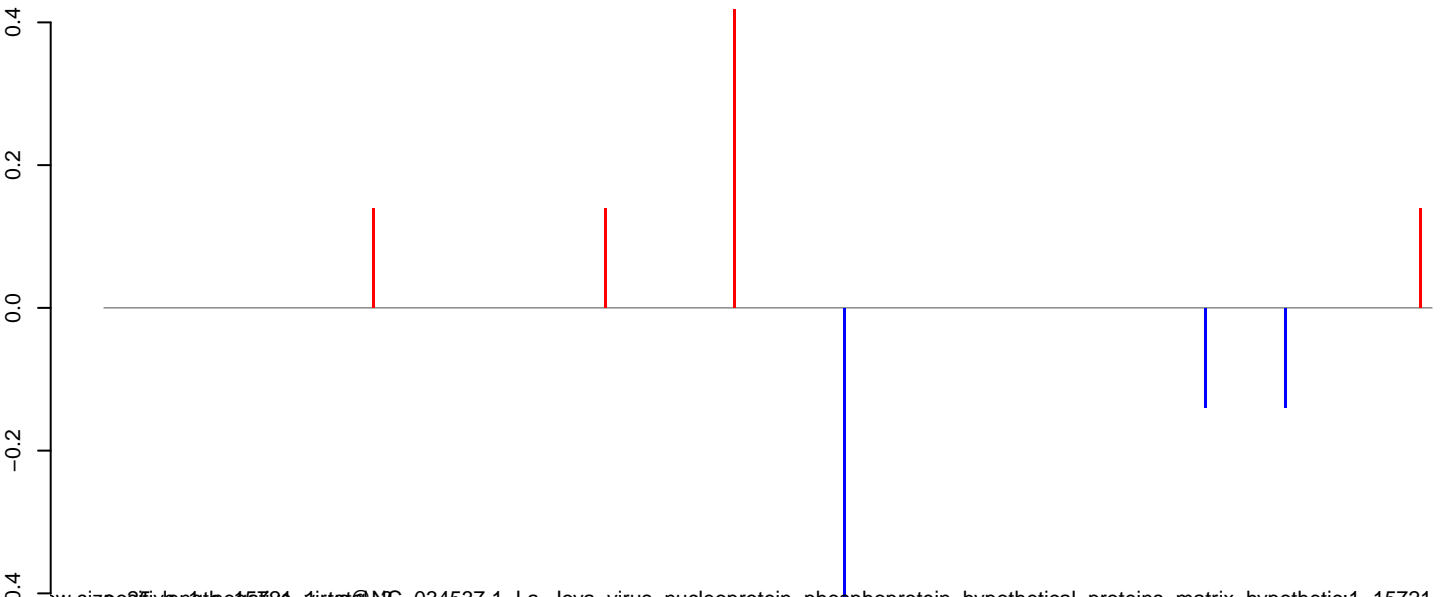
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



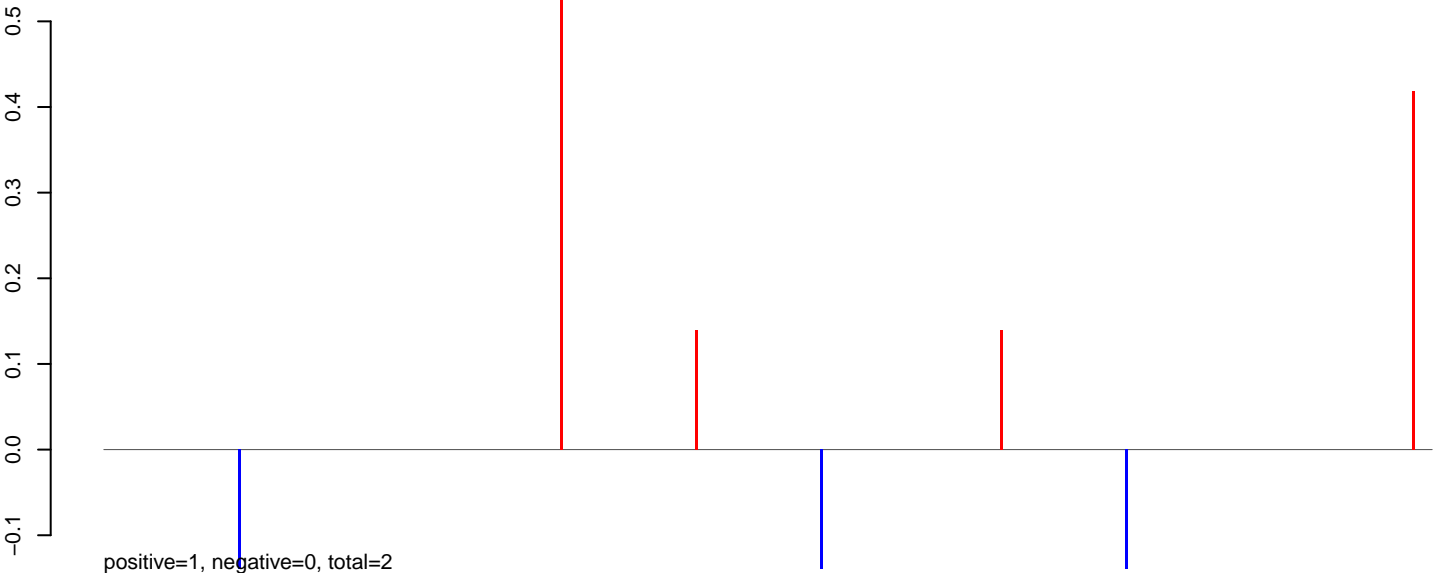
AeAlbo_C636_2dpi_mock3.rep



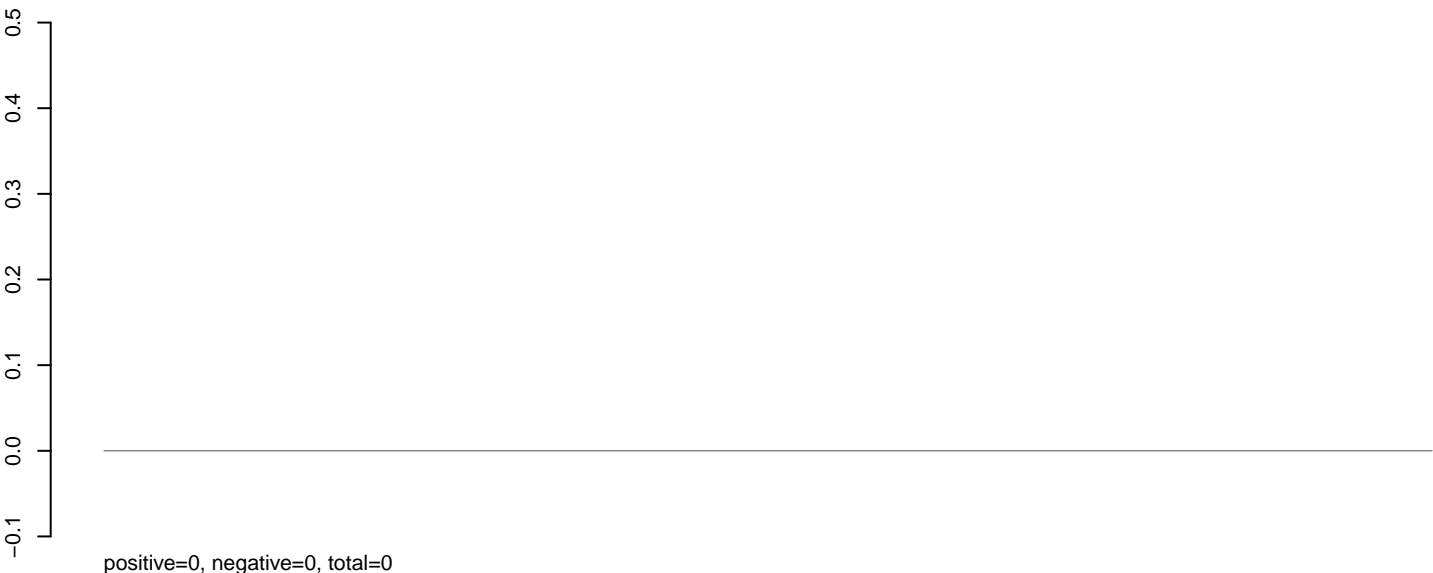
row size=25, length=5724, virus=AC_034537.1-La_Joya_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_hypothetic:1-15721

0 5000 10000 15000

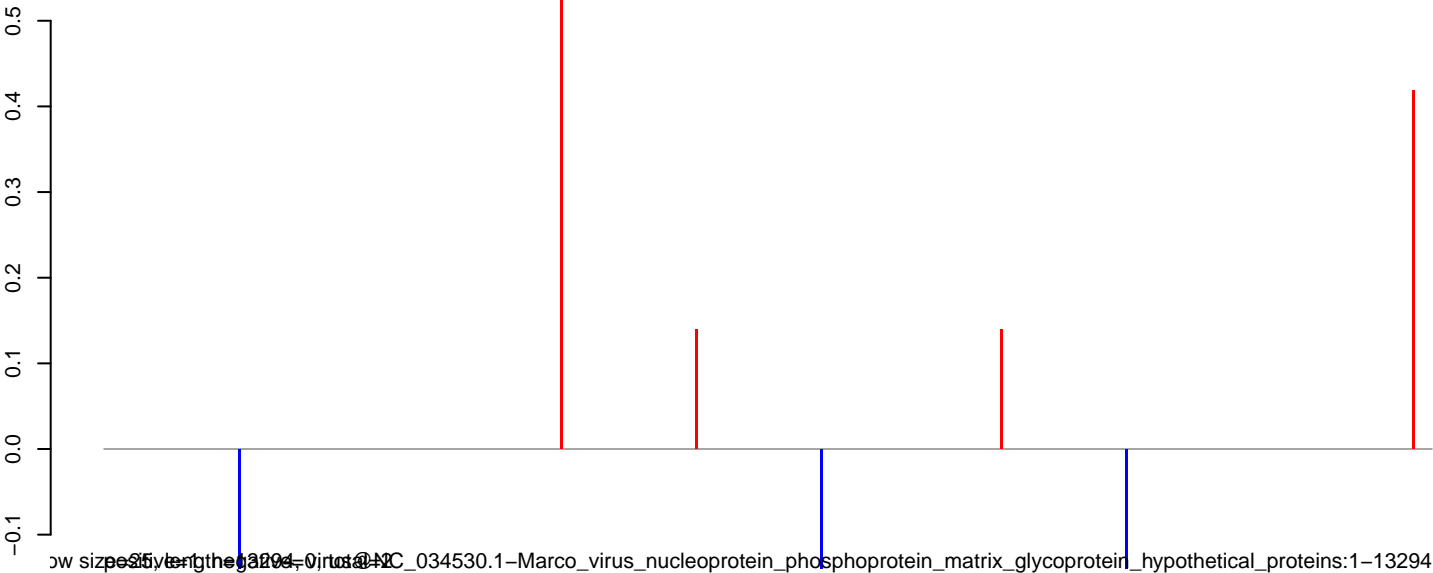
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



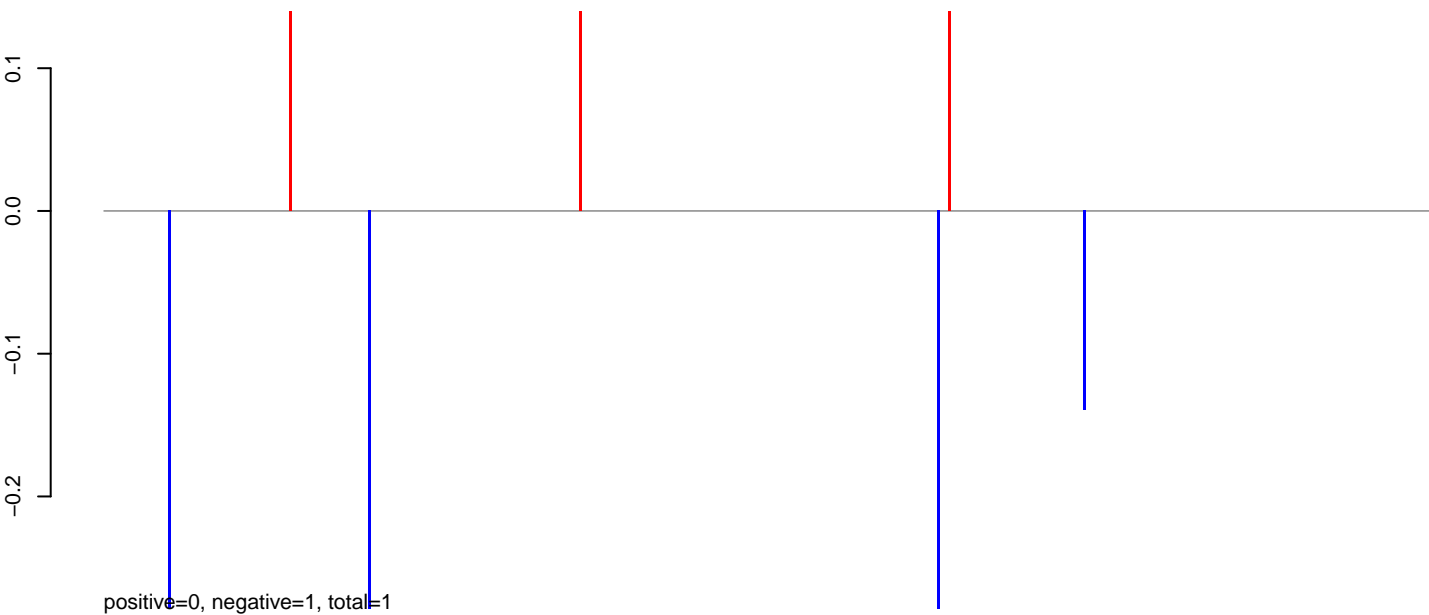
AeAlbo_C636_2dpi_mock3.rep



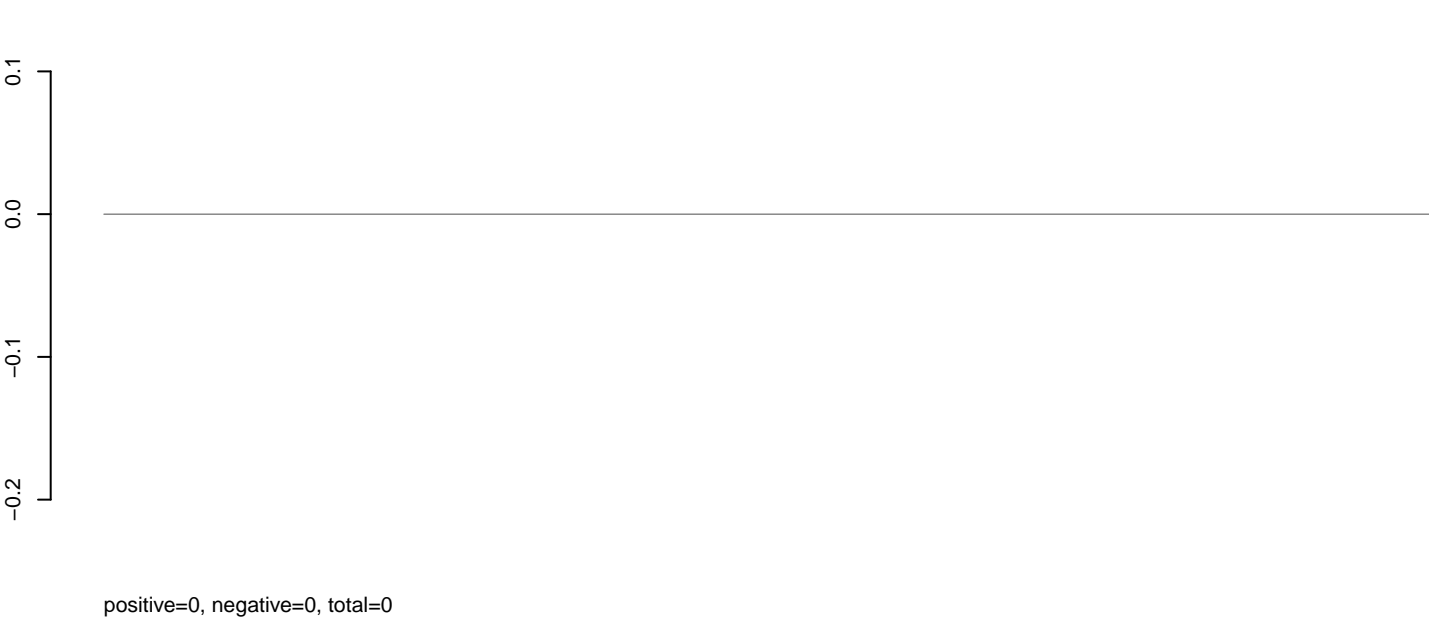
show sizes of the genes in the NC_034530.1-Marco_virus_nucleoprotein_phosphoprotein_matrix_glycoprotein_hypothetical_proteins:1-13294

0 2000 4000 6000 8000 10000 12000

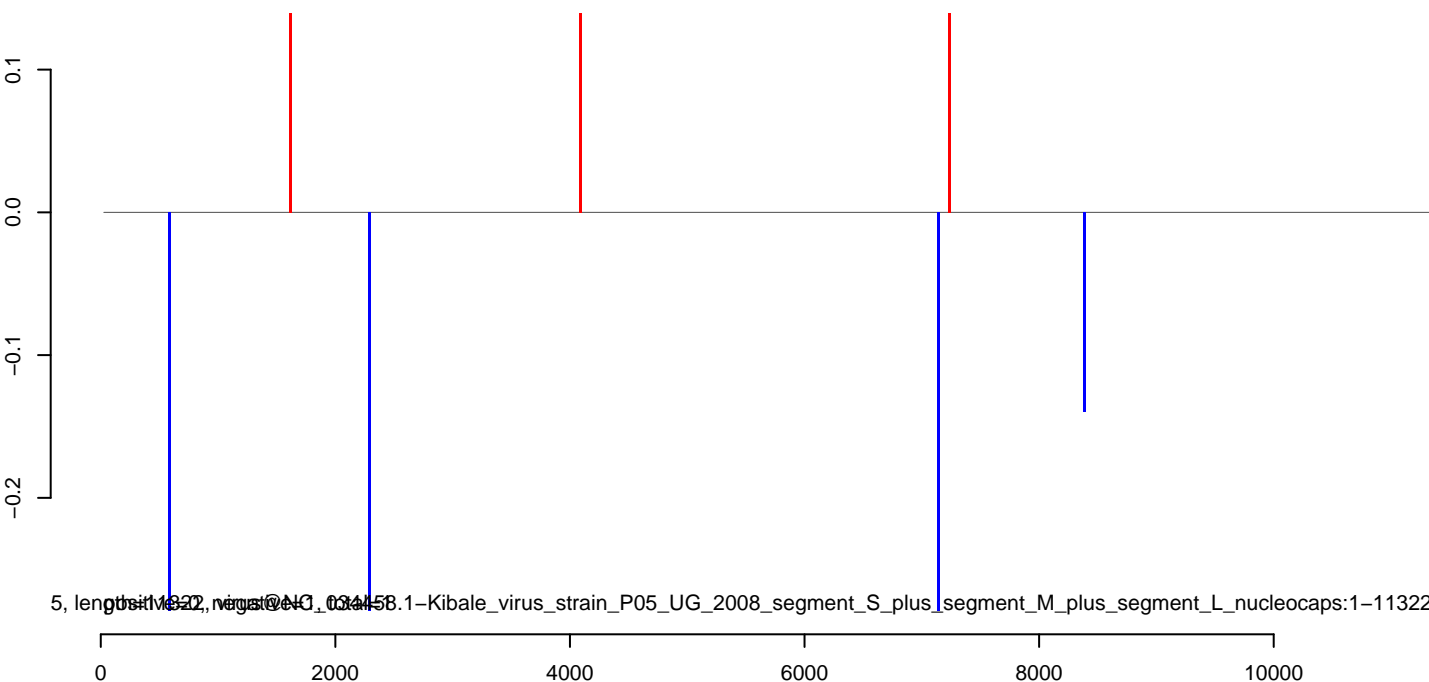
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



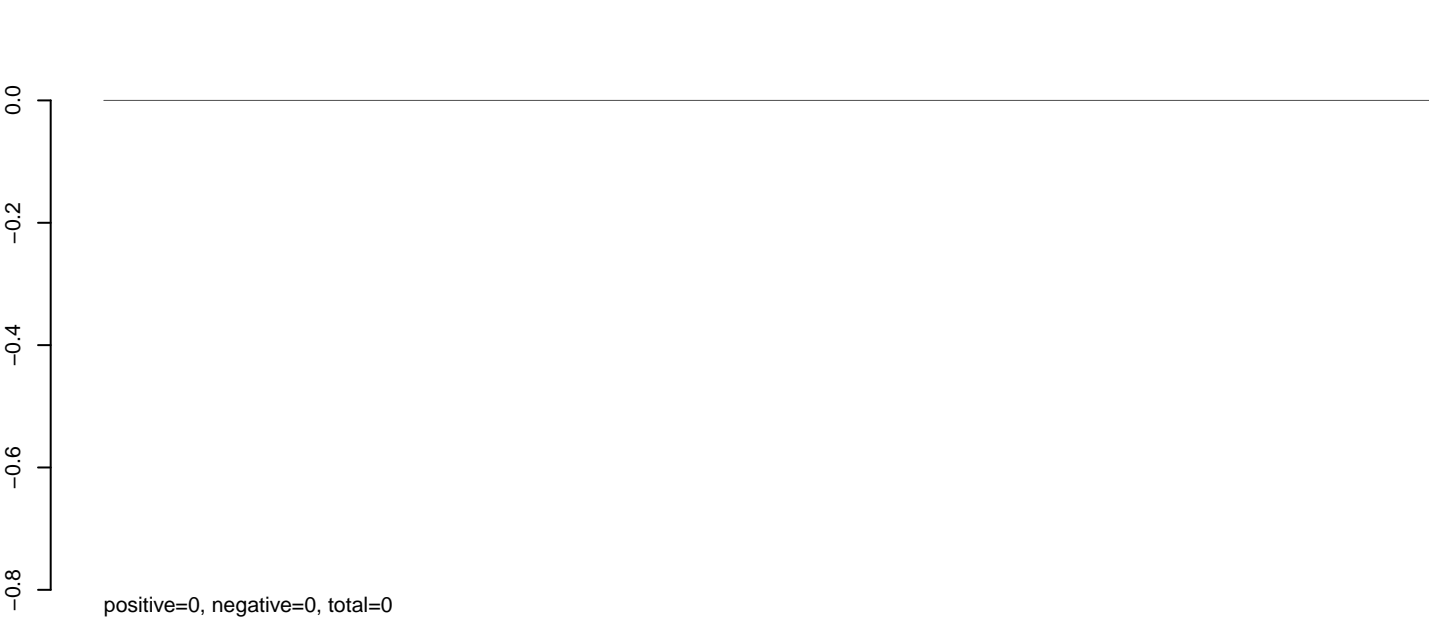
AeAlbo_C636_2dpi_mock3.rep



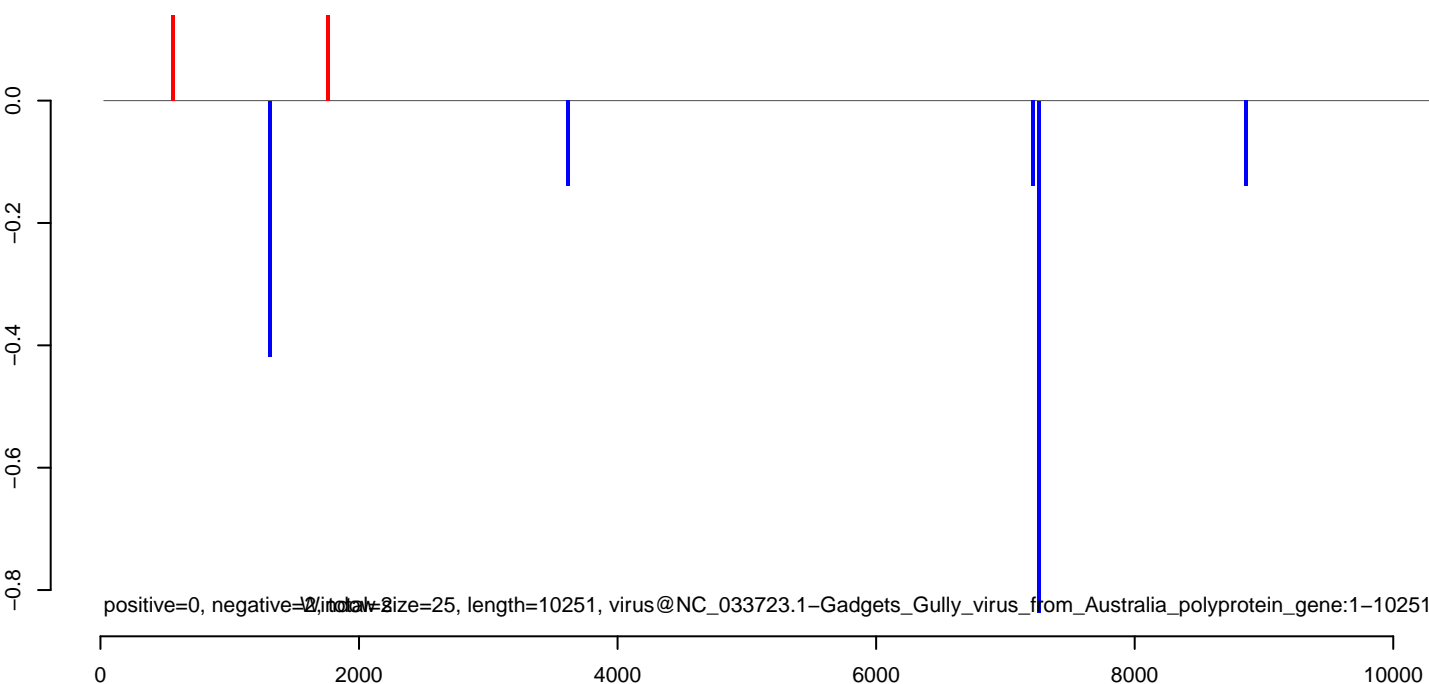
AeAlbo_C636_2dpi_mock3.18_23.rep



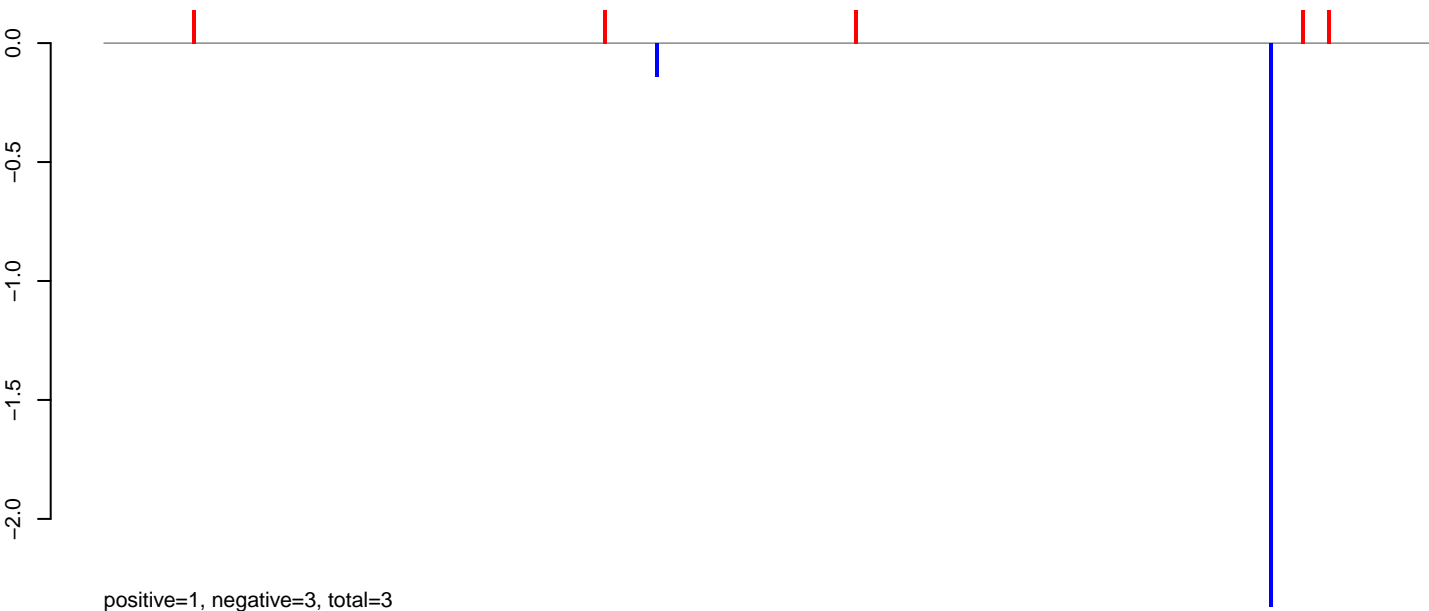
AeAlbo_C636_2dpi_mock3.24_35.rep



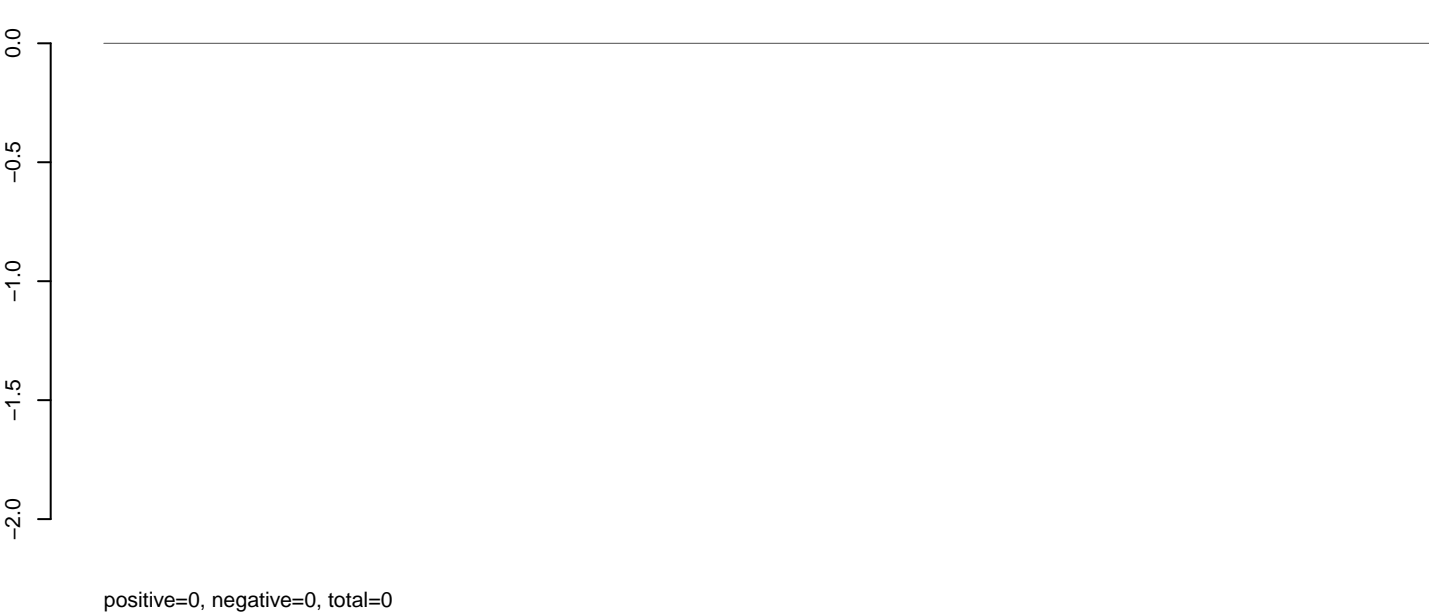
AeAlbo_C636_2dpi_mock3.rep



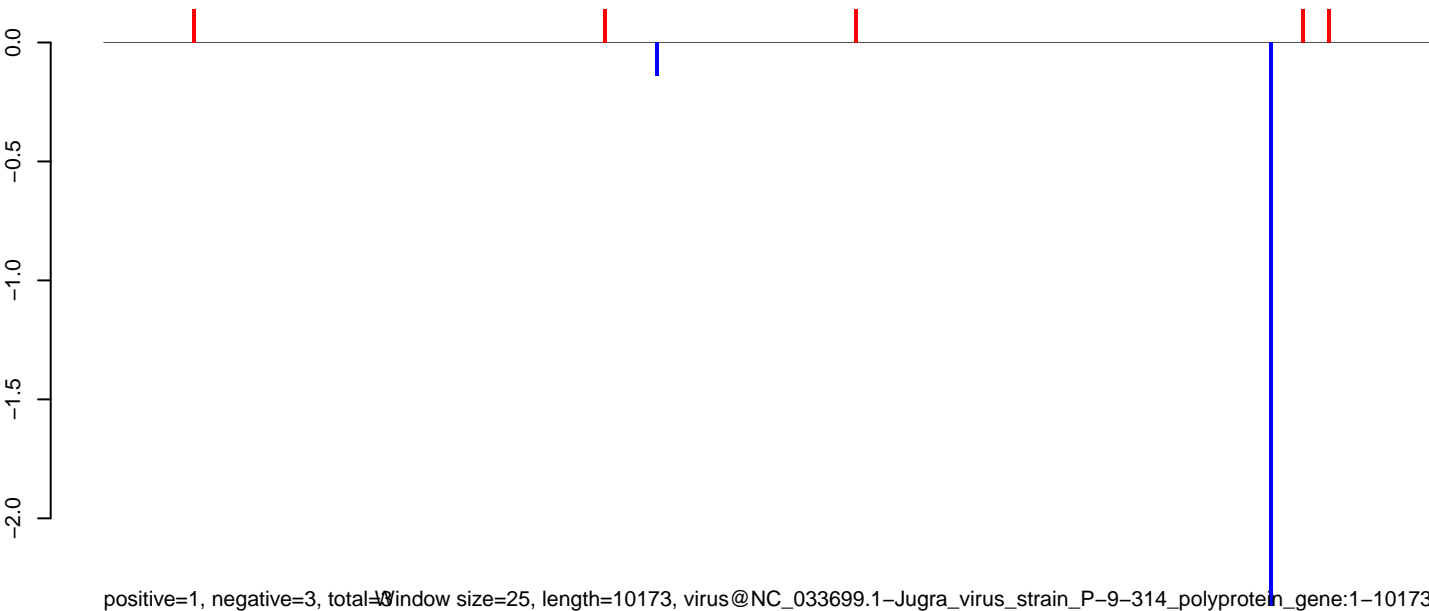
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



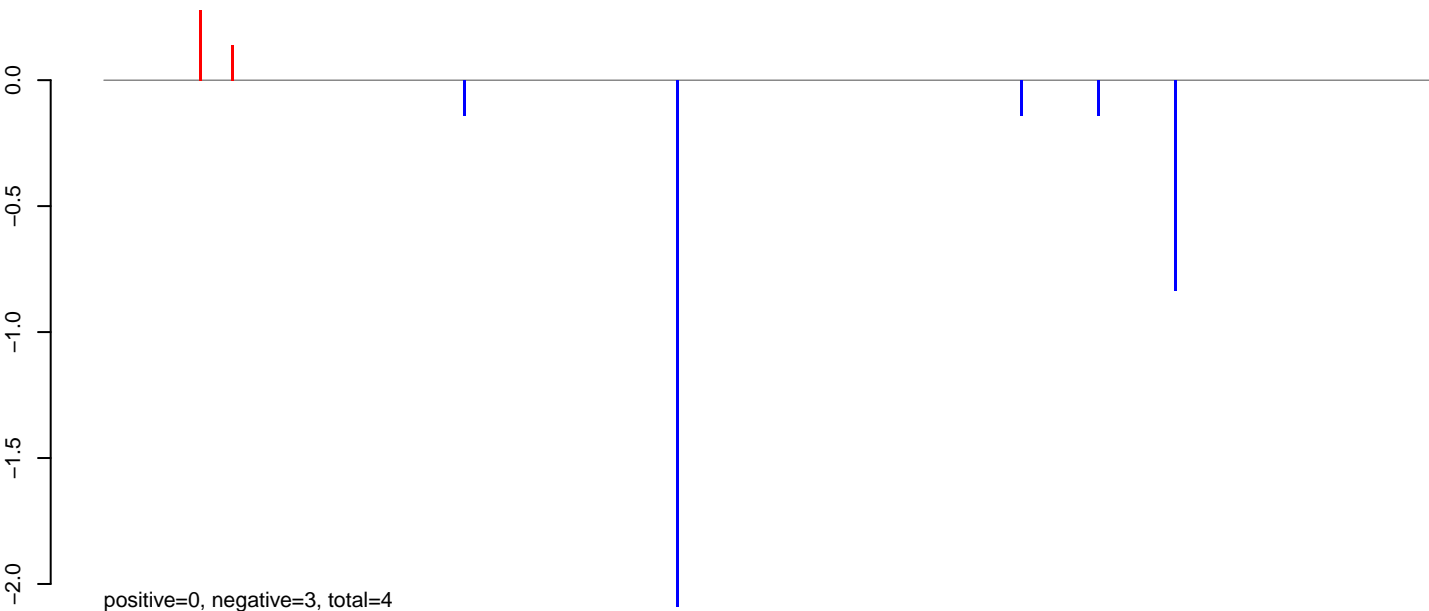
AeAlbo_C636_2dpi_mock3.rep



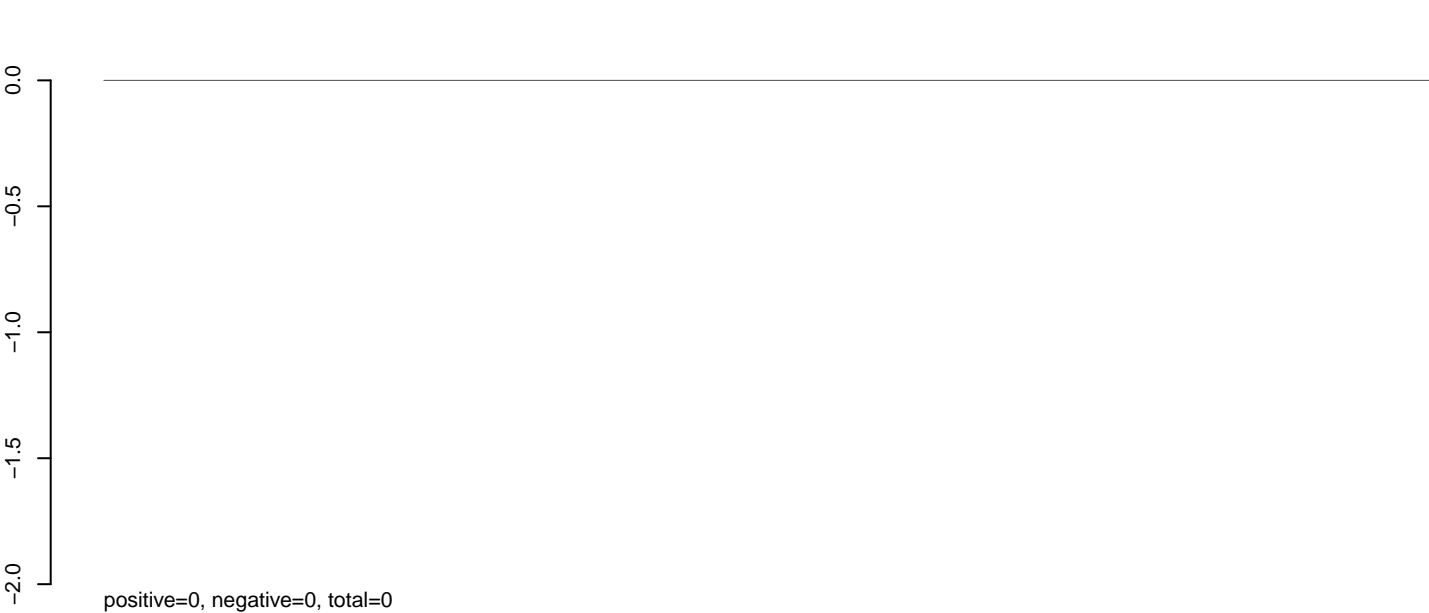
Window size=25, length=10173, virus@NC_033699.1-Jugra_virus_strain_P-9-314_polyprotein_gene:1-10173

0 2000 4000 6000 8000 10000

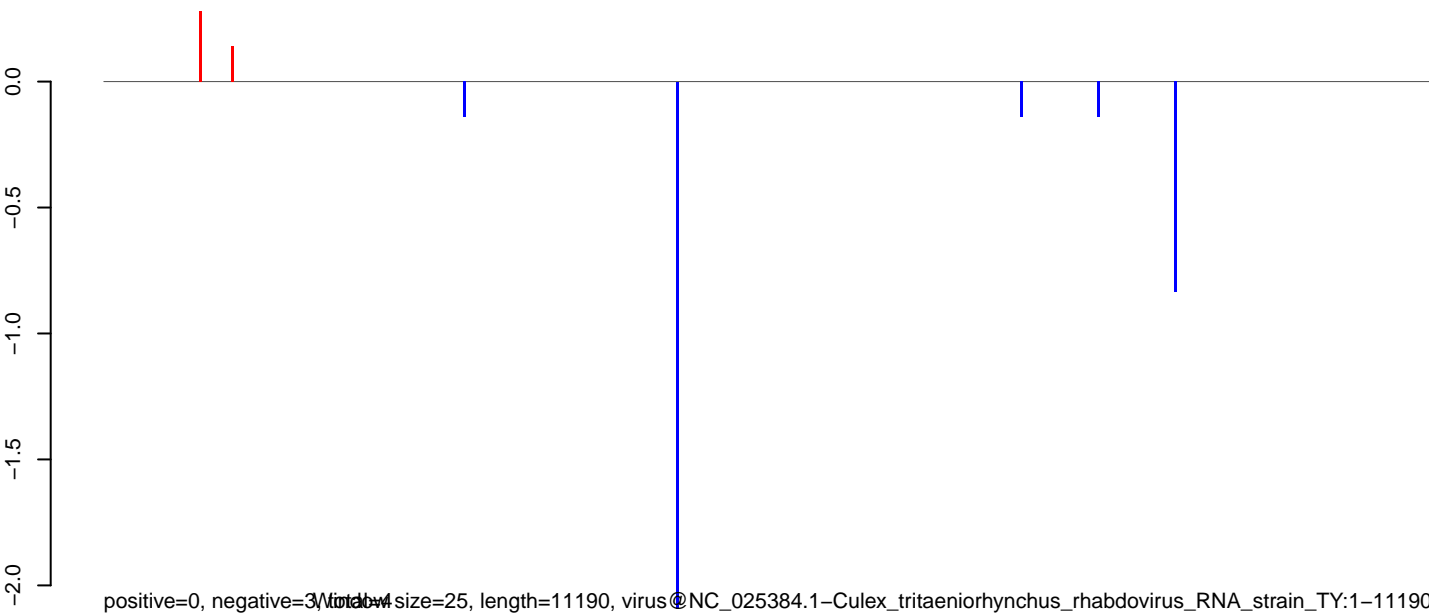
AeAlbo_C636_2dpi_mock3.18_23.rep



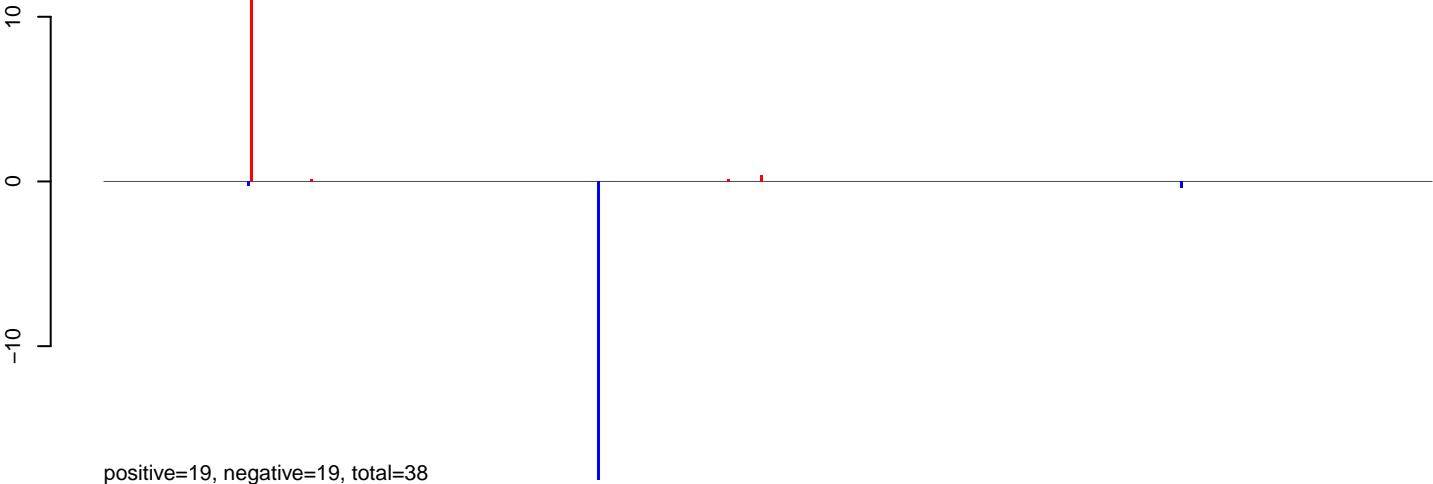
AeAlbo_C636_2dpi_mock3.24_35.rep



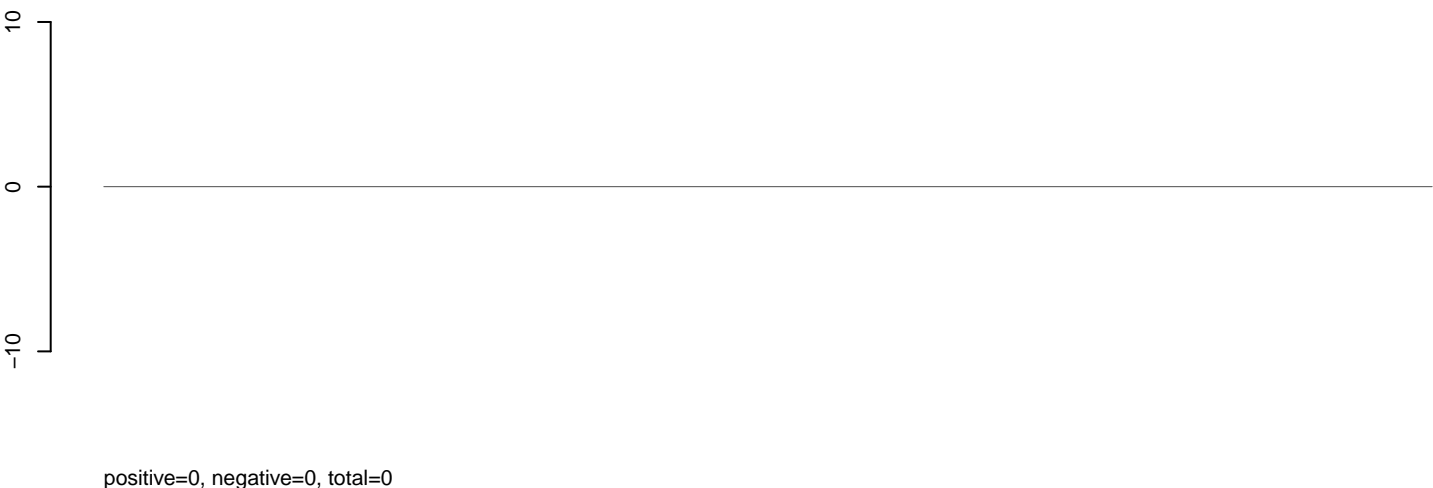
AeAlbo_C636_2dpi_mock3.rep



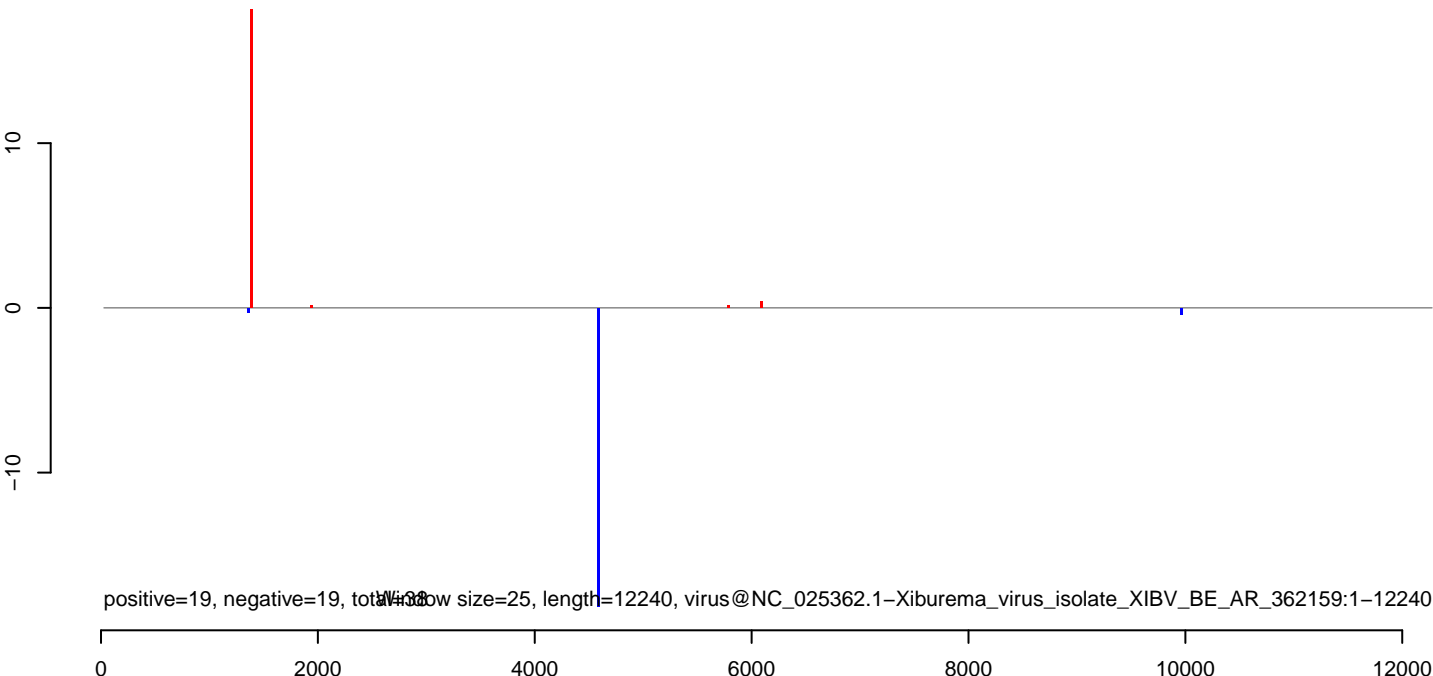
AeAlbo_C636_2dpi_mock3.18_23.rep



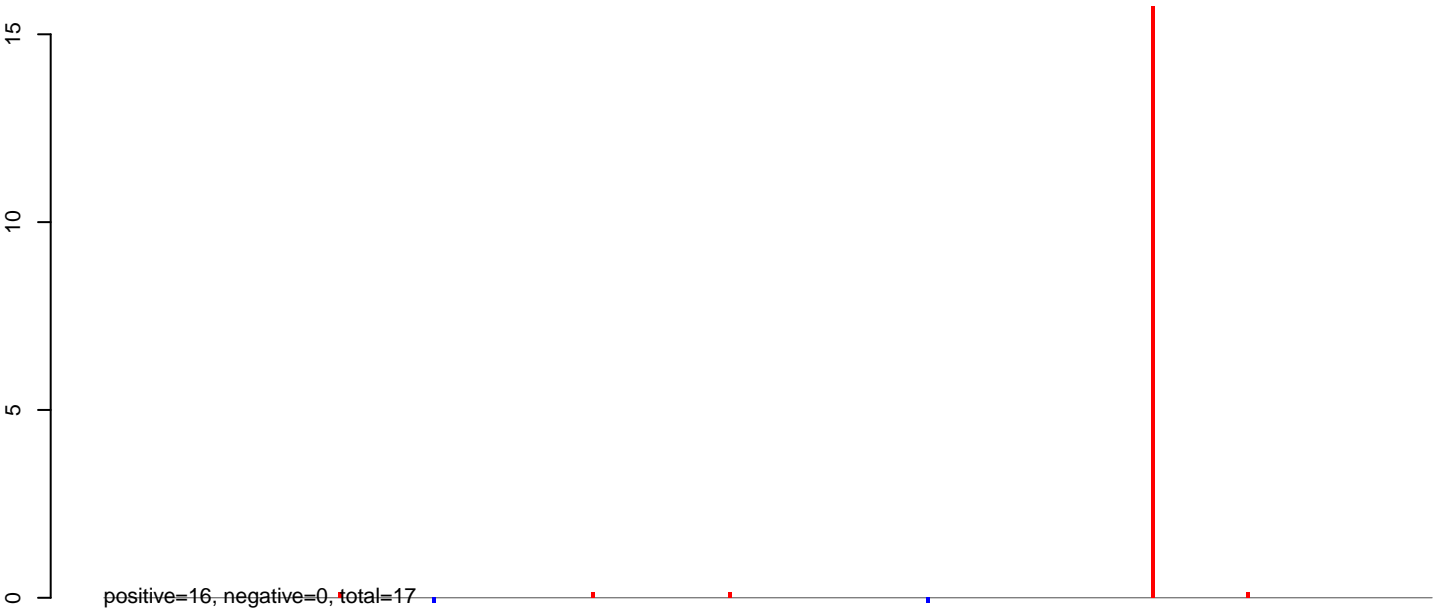
AeAlbo_C636_2dpi_mock3.24_35.rep



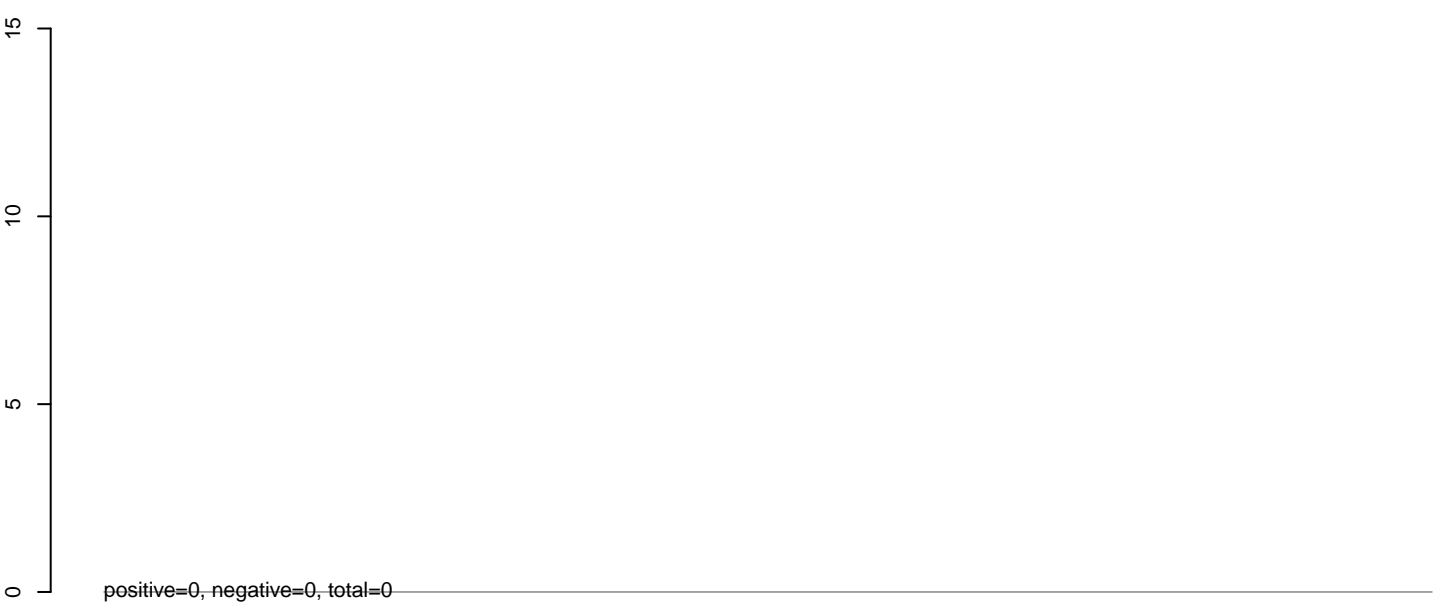
AeAlbo_C636_2dpi_mock3.rep



AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



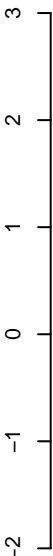
AeAlbo_C636_2dpi_mock3.rep



Window-size=25, length=10878, virus@NC_024299.2-Nienokoue_virus_isolate_B51_C1_2004:1-10878

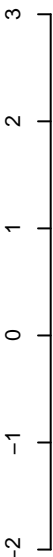
0 2000 4000 6000 8000 10000

AeAlbo_C636_2dpi_mock3.18_23.rep



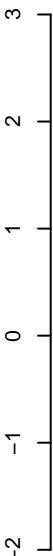
positive=7, negative=3, total=10

AeAlbo_C636_2dpi_mock3.24_35.rep



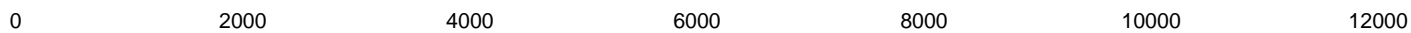
positive=0, negative=0, total=0

AeAlbo_C636_2dpi_mock3.rep

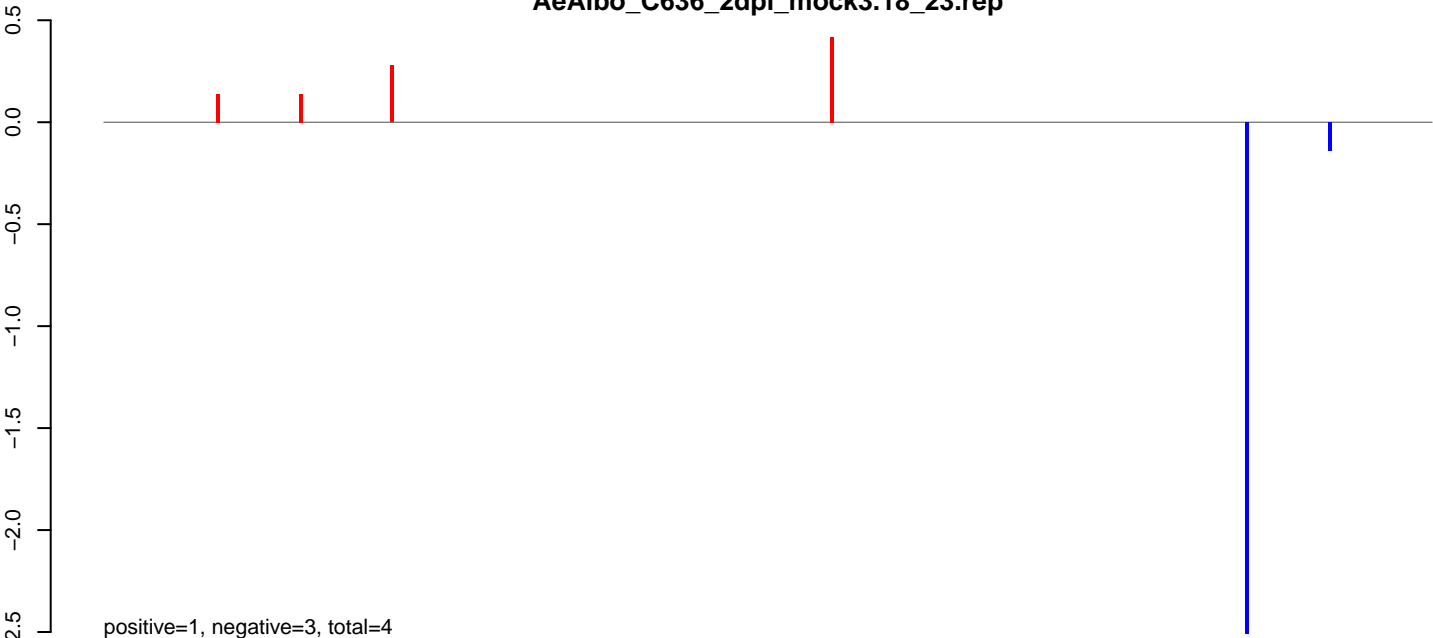


positive=7, negative=3, total=10

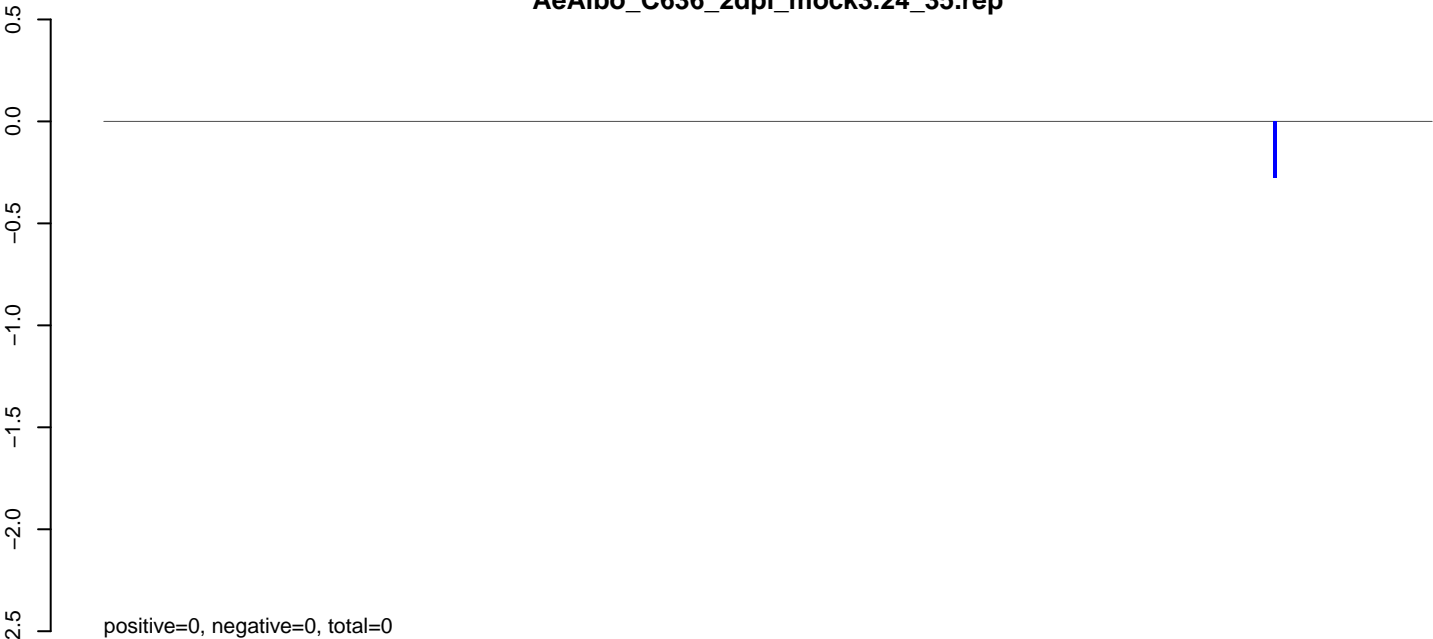
Window size=25, length=11688, virus@NC_016959.1-Ndumu_virus:1-11688



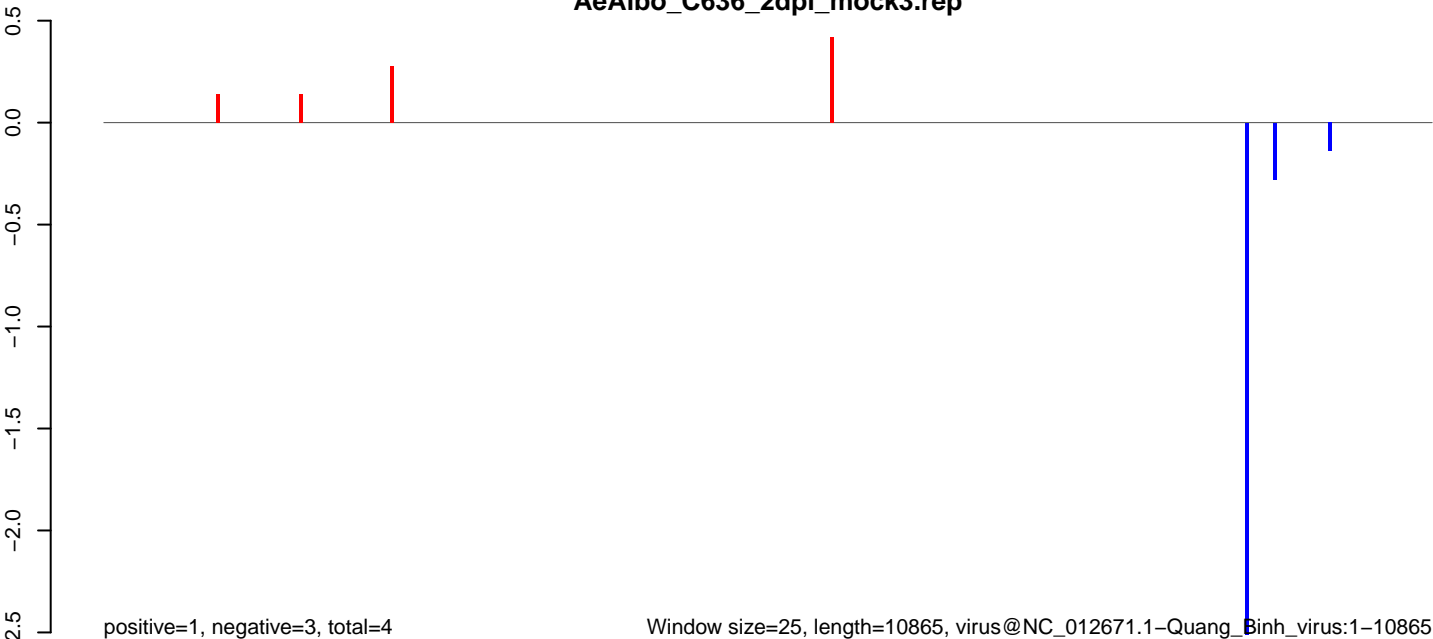
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



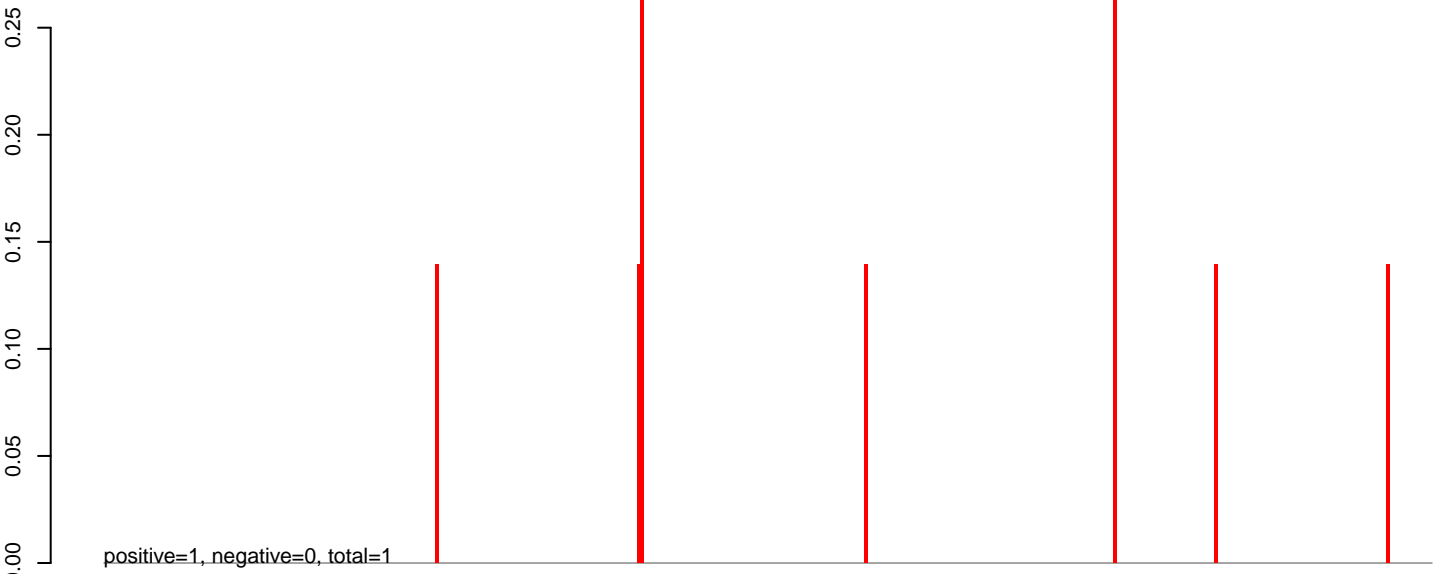
AeAlbo_C636_2dpi_mock3.rep



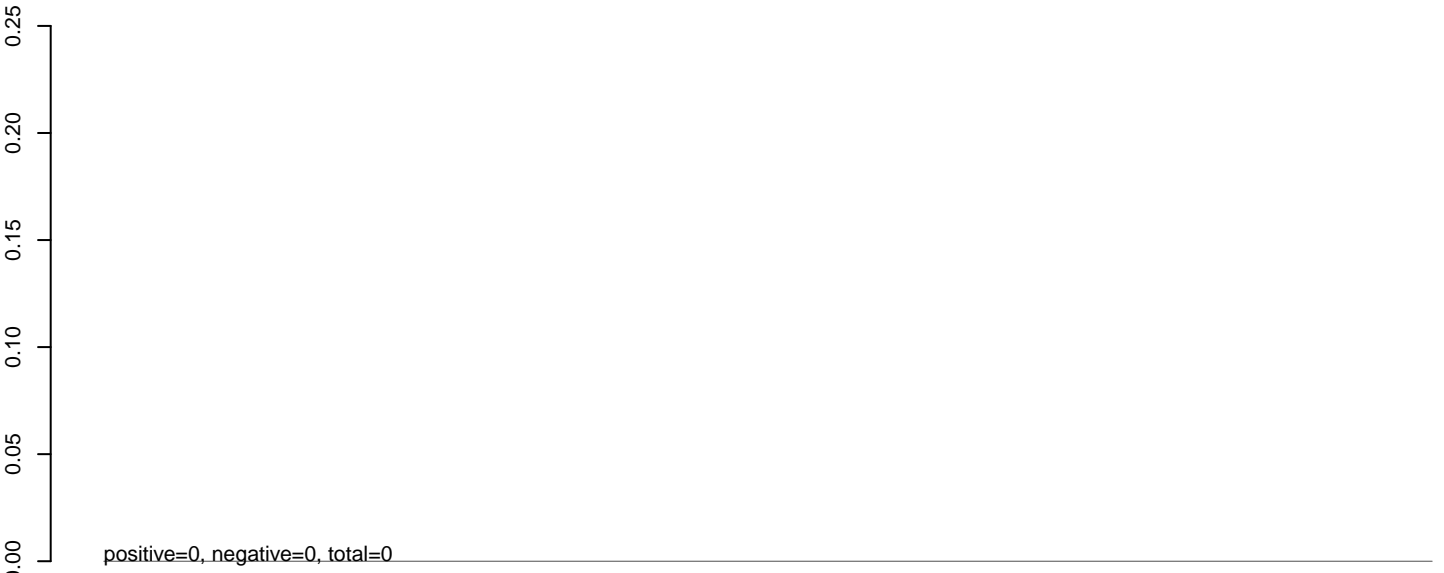
Window size=25, length=10865, virus@NC_012671.1-Quang_Binh_virus:1-10865

0 2000 4000 6000 8000 10000

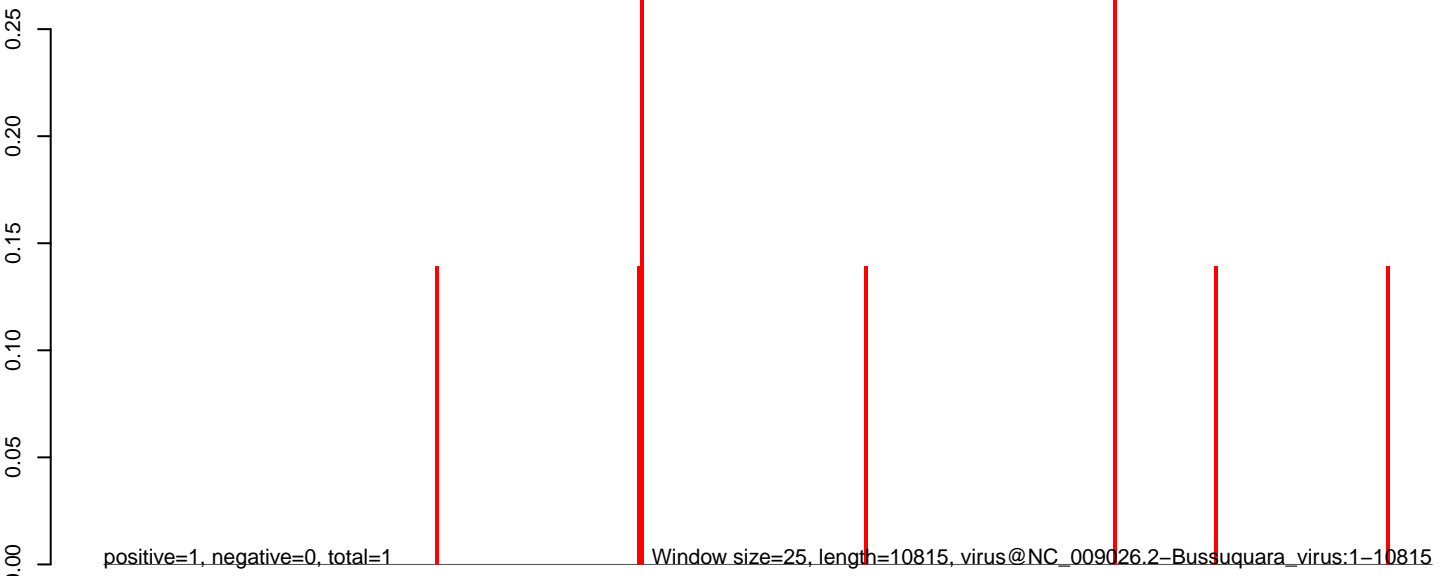
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

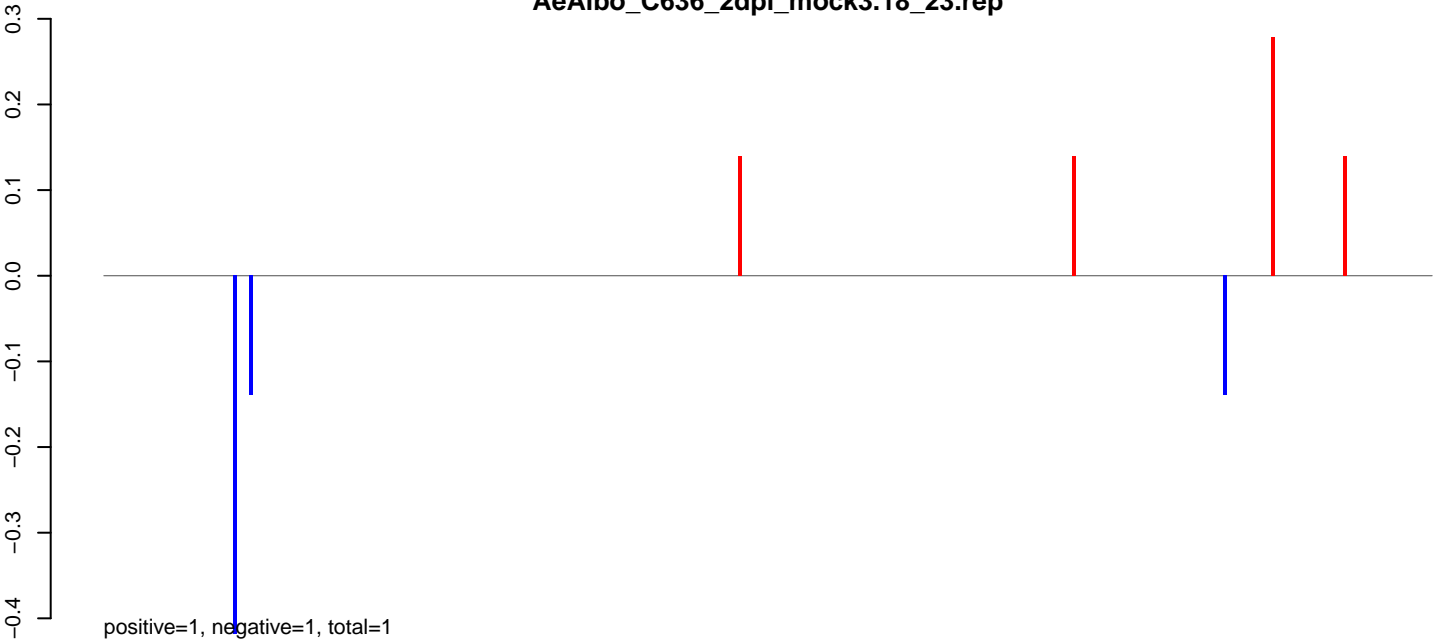


AeAlbo_C636_2dpi_mock3.rep

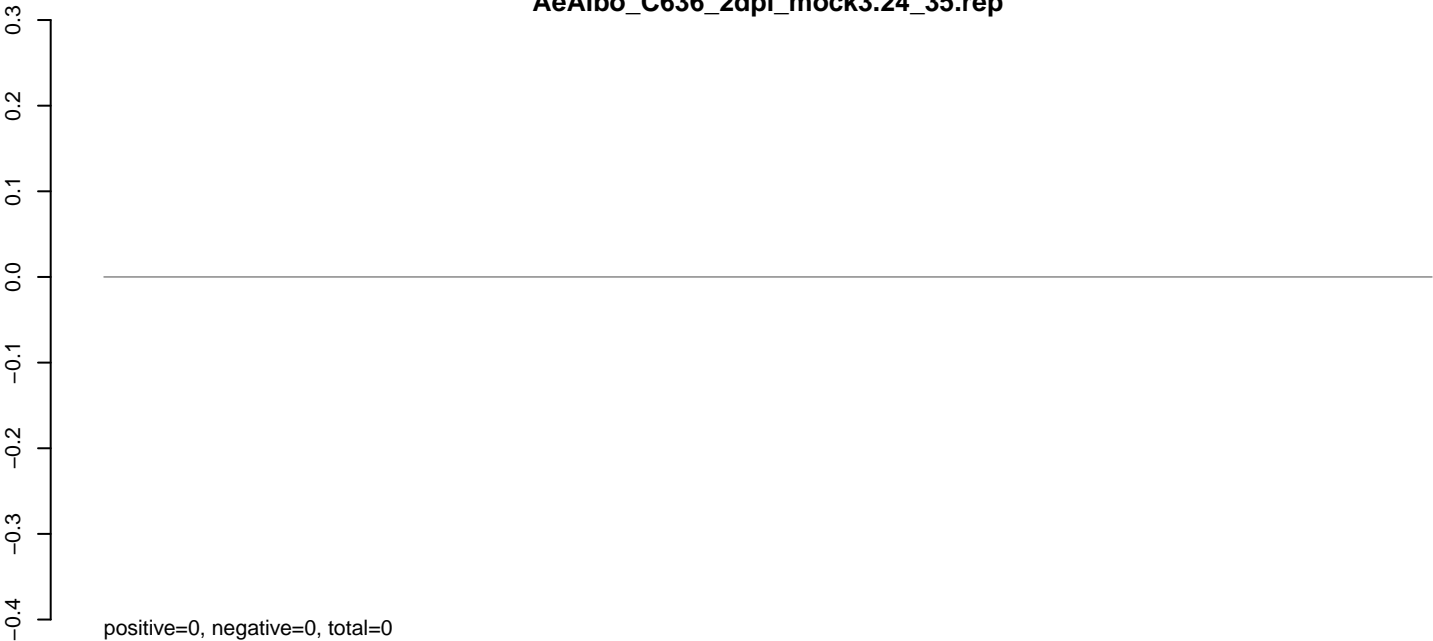


0 2000 4000 6000 8000 10000

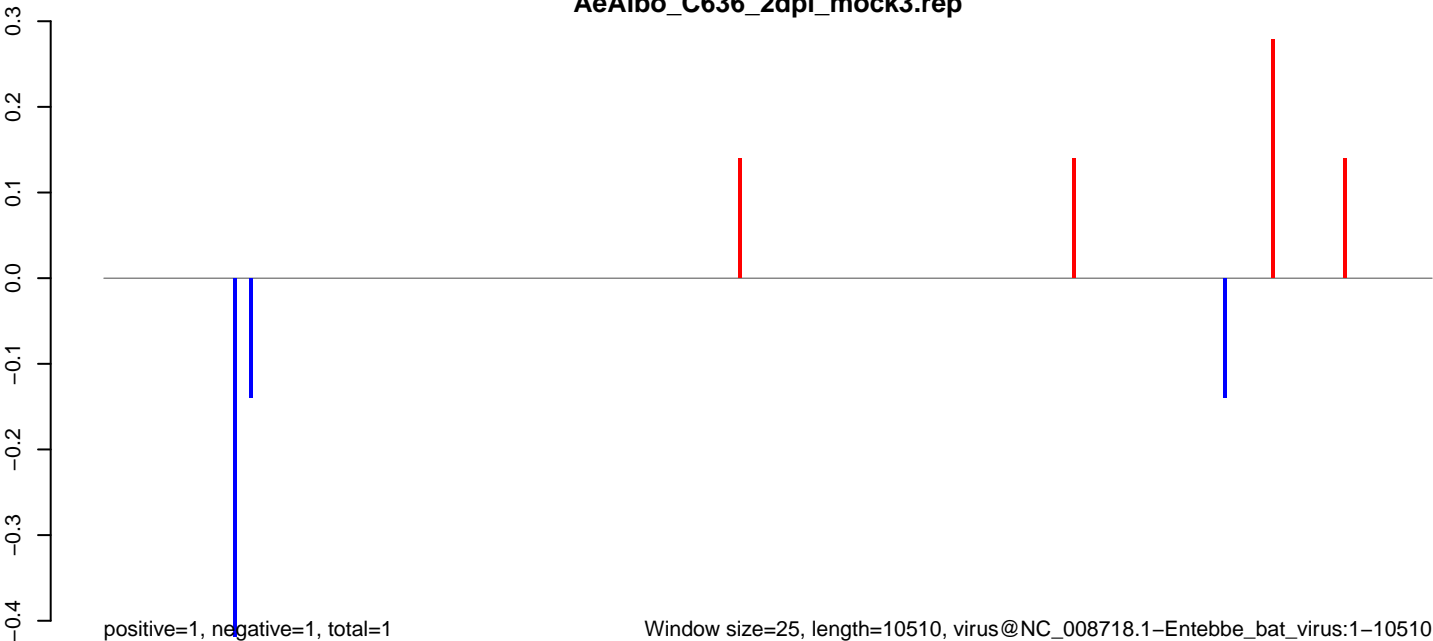
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



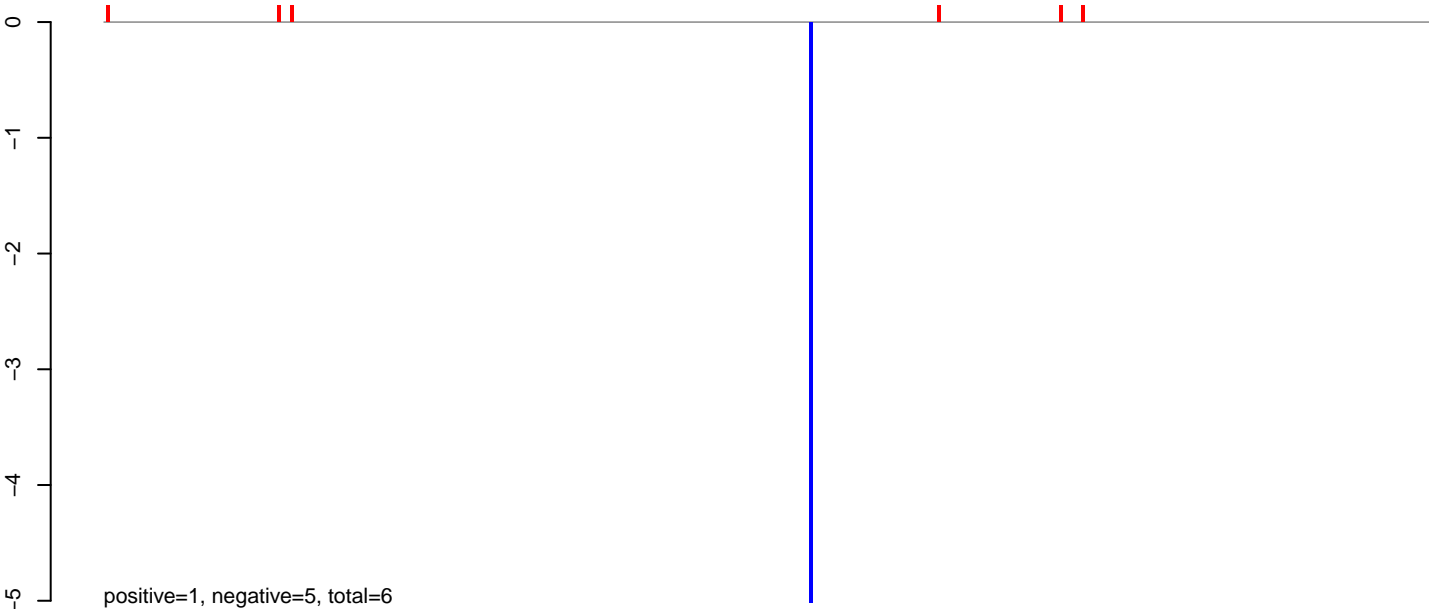
AeAlbo_C636_2dpi_mock3.rep



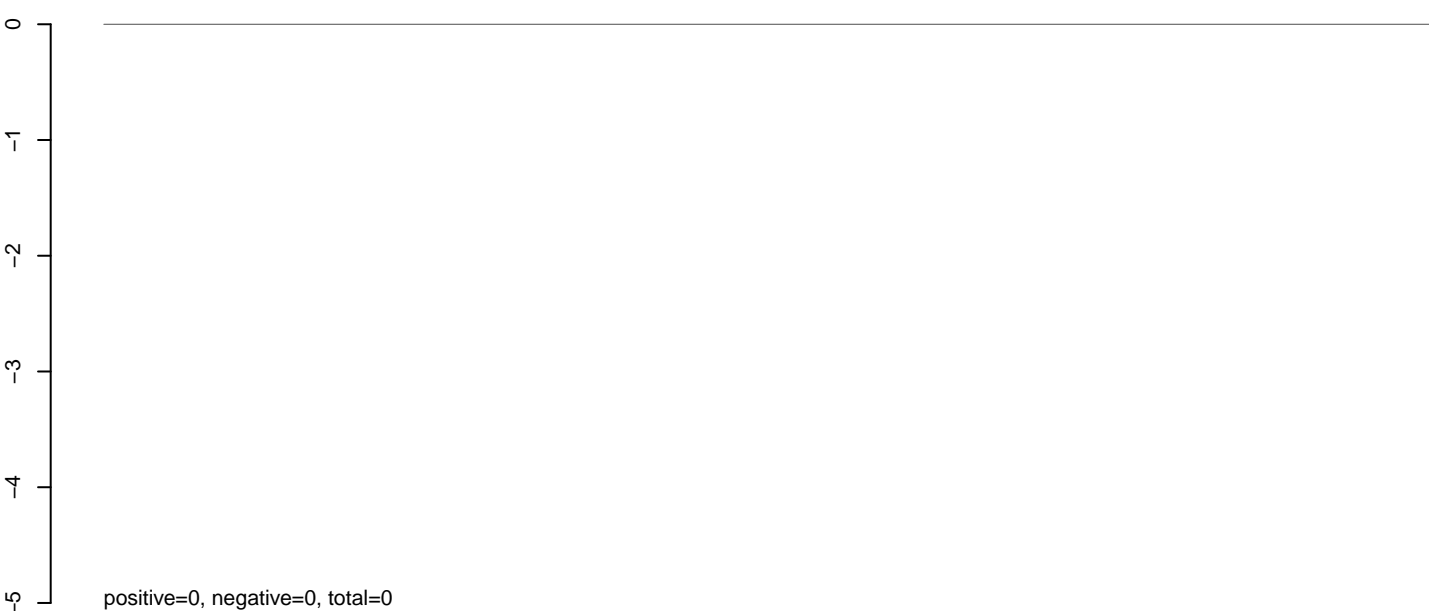
Window size=25, length=10510, virus@NC_008718.1-Entebbe_bat_virus:1-10510

0 2000 4000 6000 8000 10000

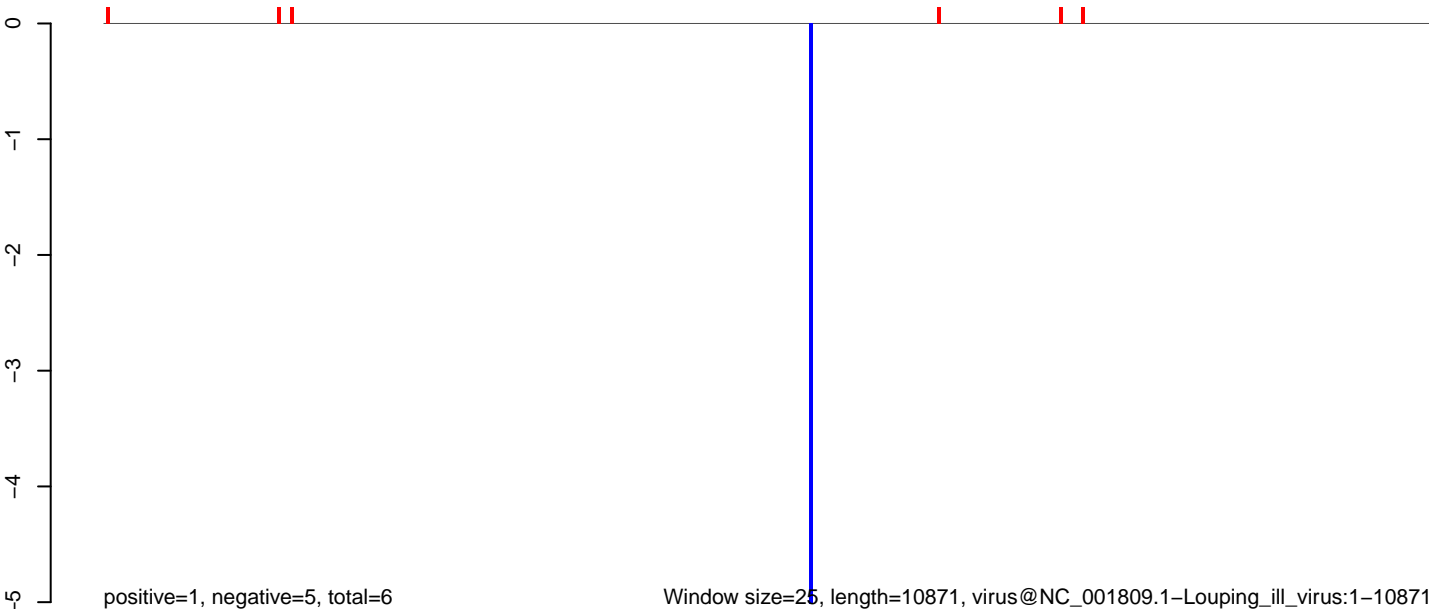
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



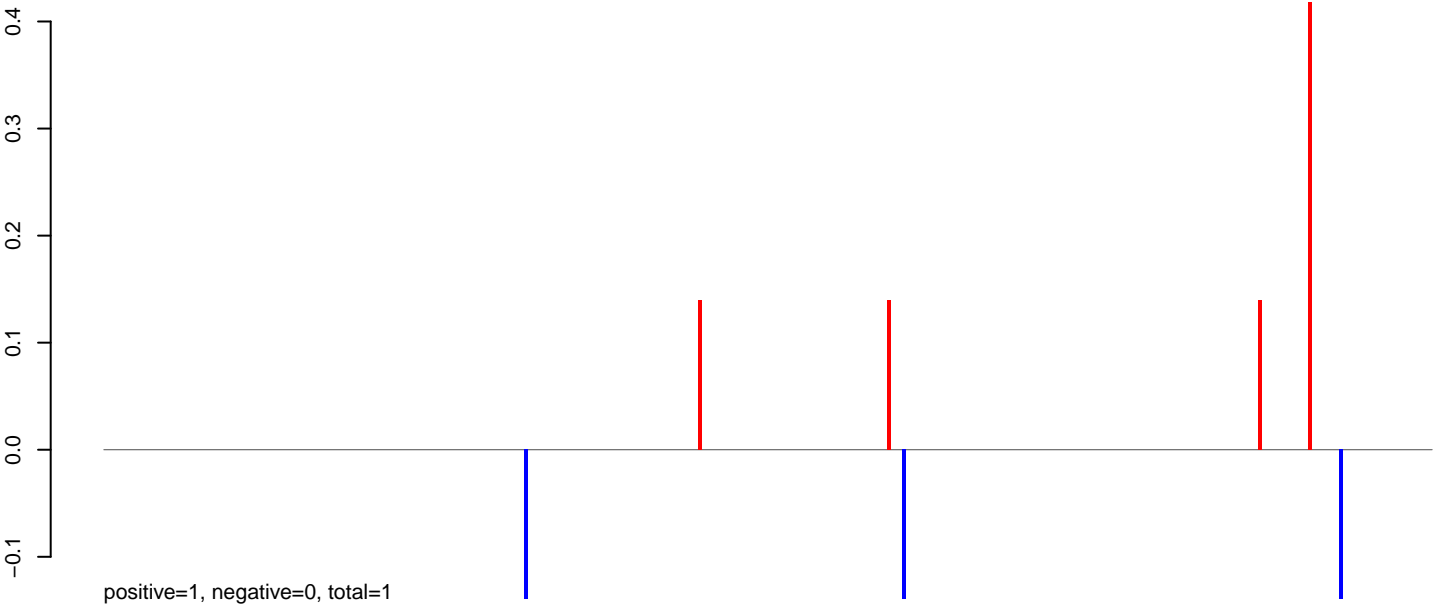
AeAlbo_C636_2dpi_mock3.rep



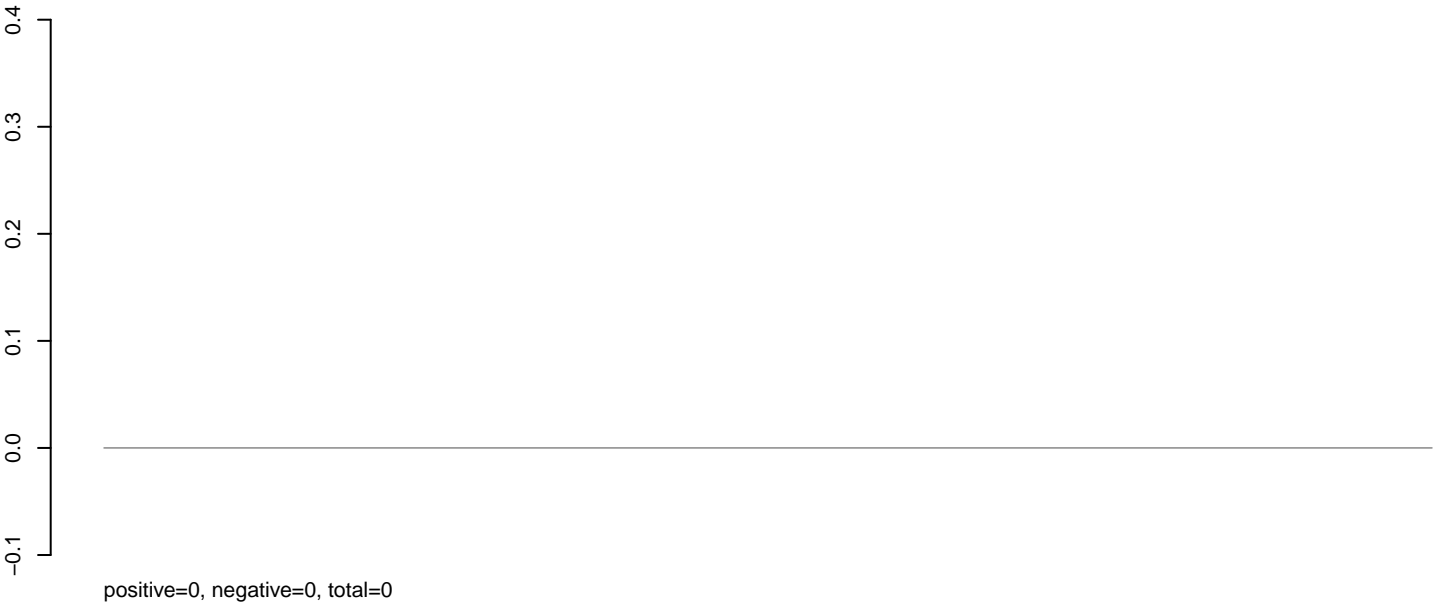
Window size=25, length=10871, virus@NC_001809.1-Louping_ill_virus:1-10871

0 2000 4000 6000 8000 10000

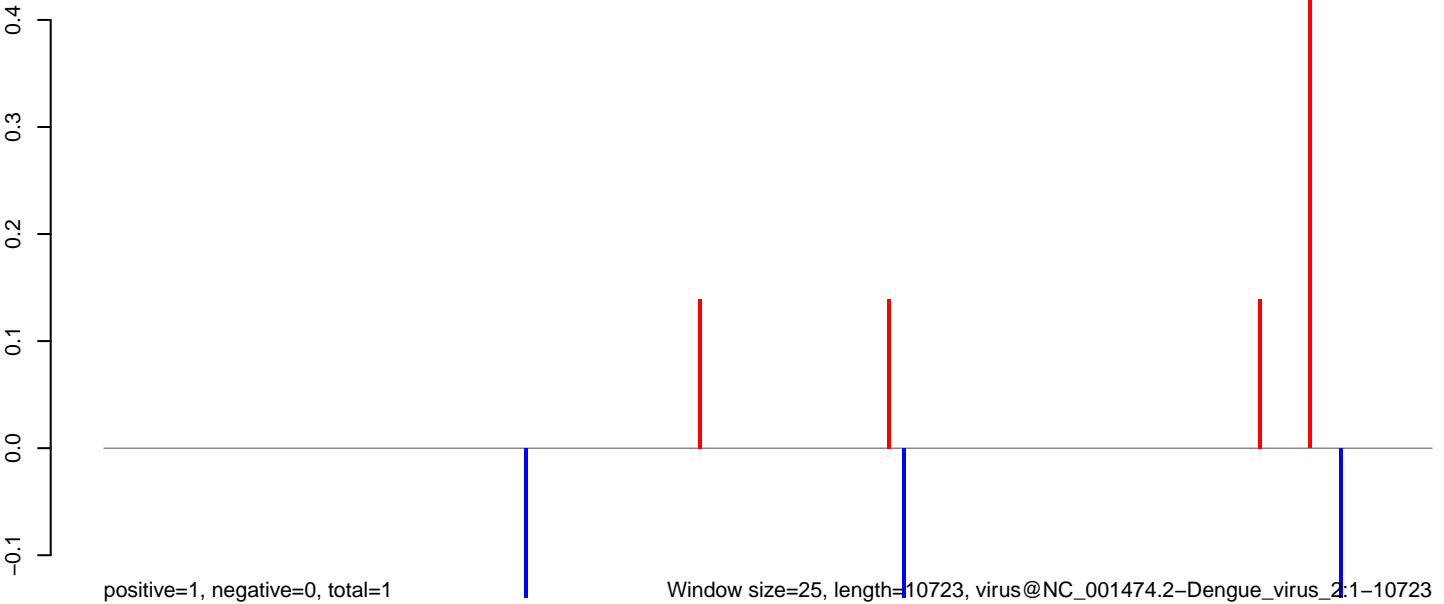
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

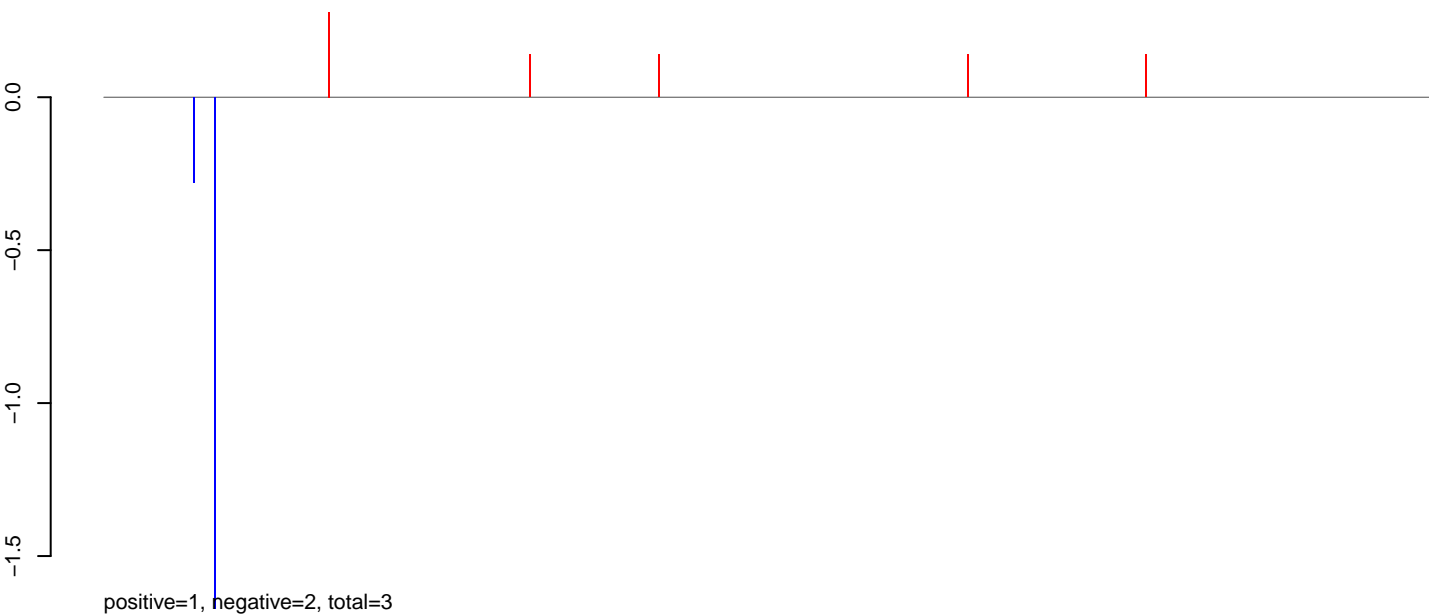


AeAlbo_C636_2dpi_mock3.rep

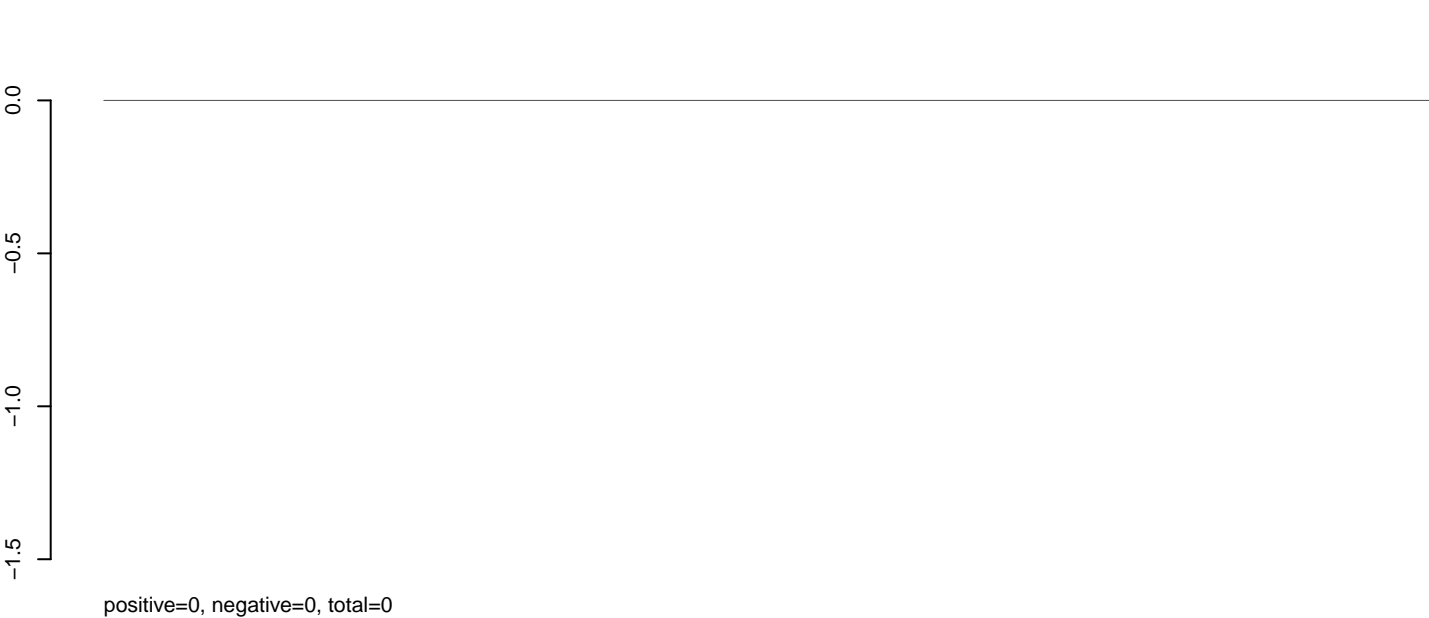


0 2000 4000 6000 8000 10000

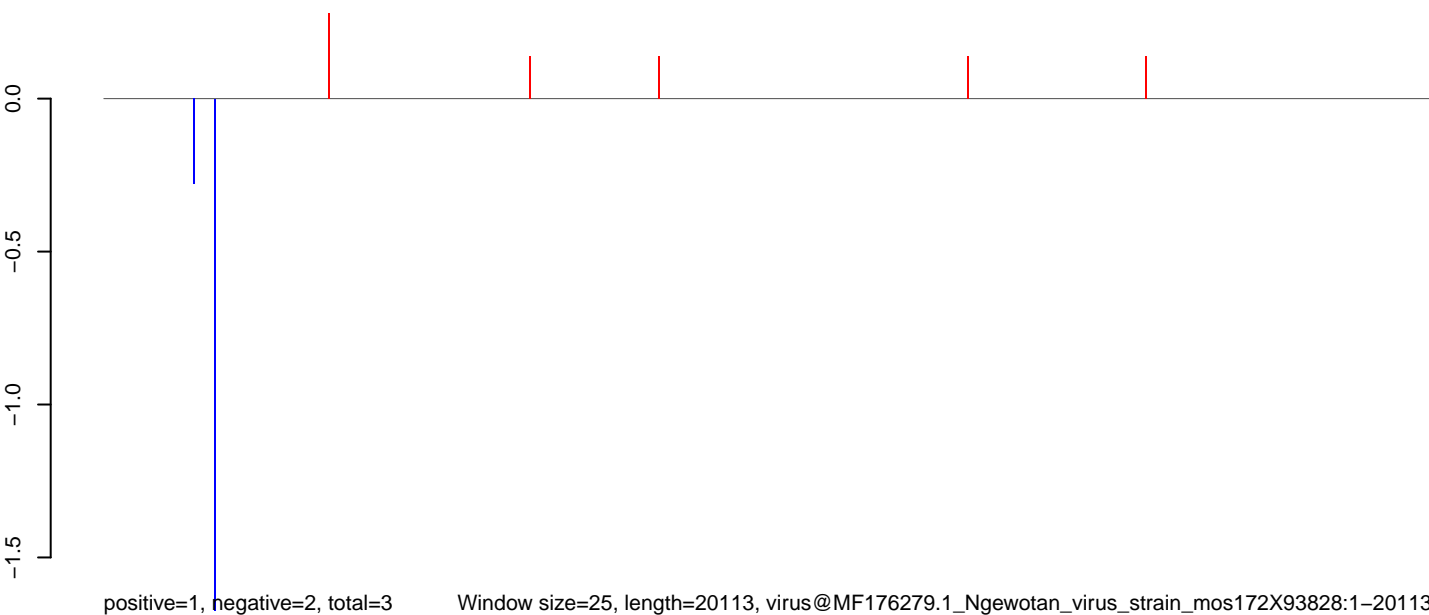
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



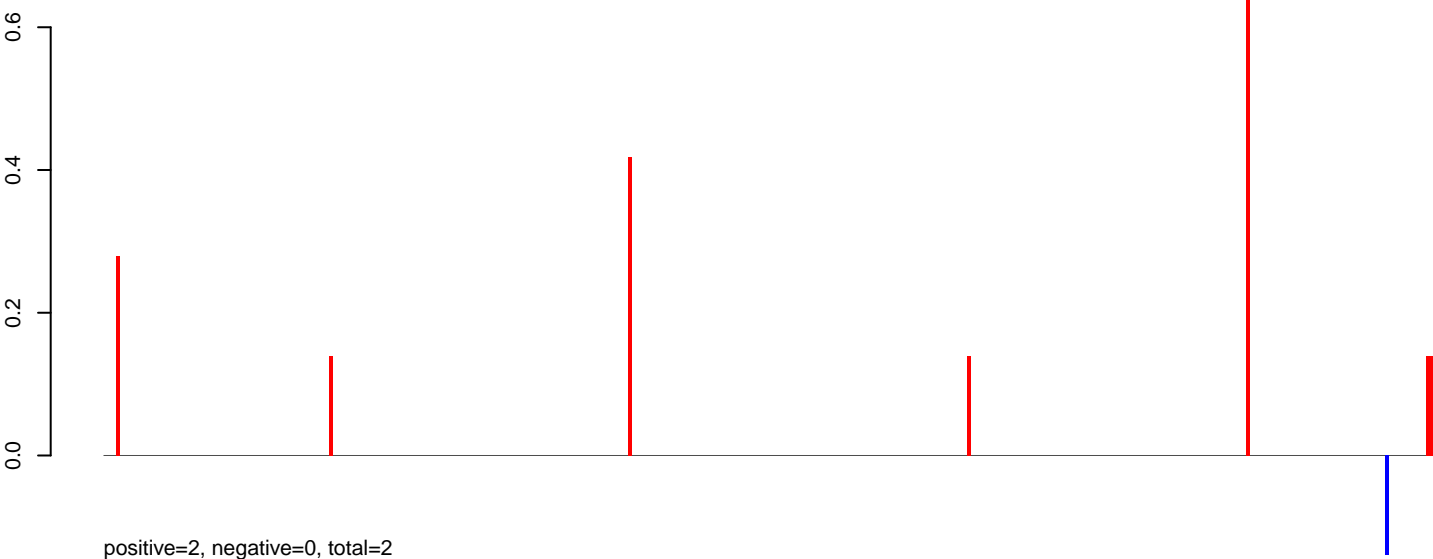
AeAlbo_C636_2dpi_mock3.rep



Window size=25, length=20113, virus@MF176279.1_Ngewotan_virus_strain_mos172X93828:1-20113

0 5000 10000 15000 20000

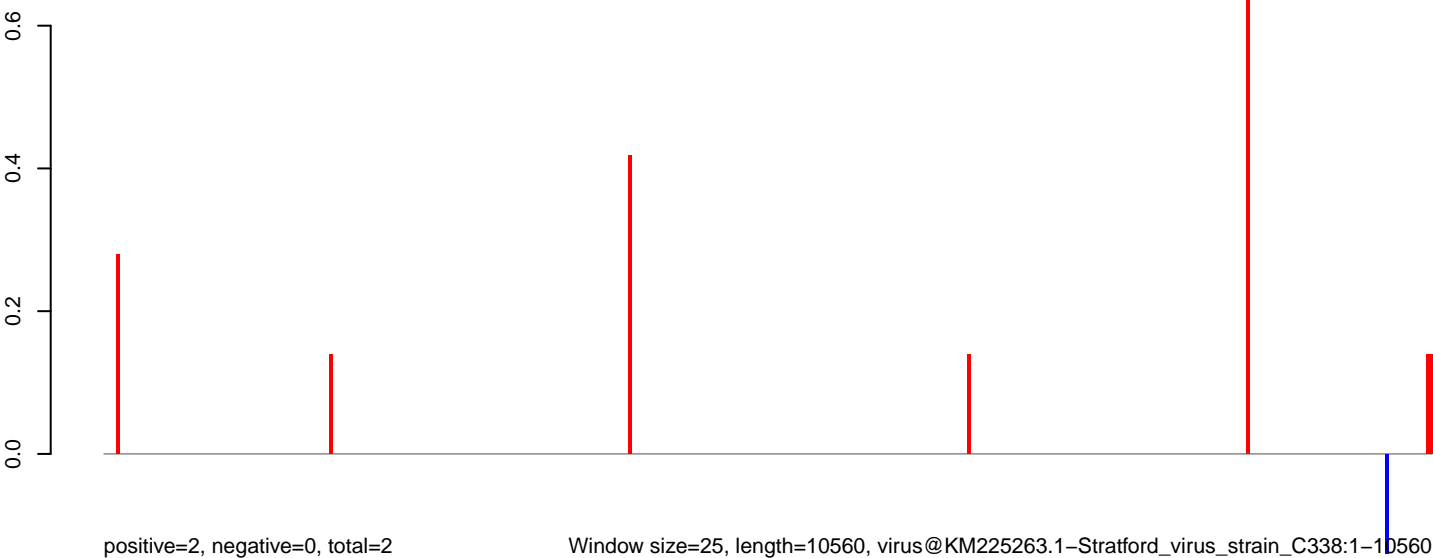
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep

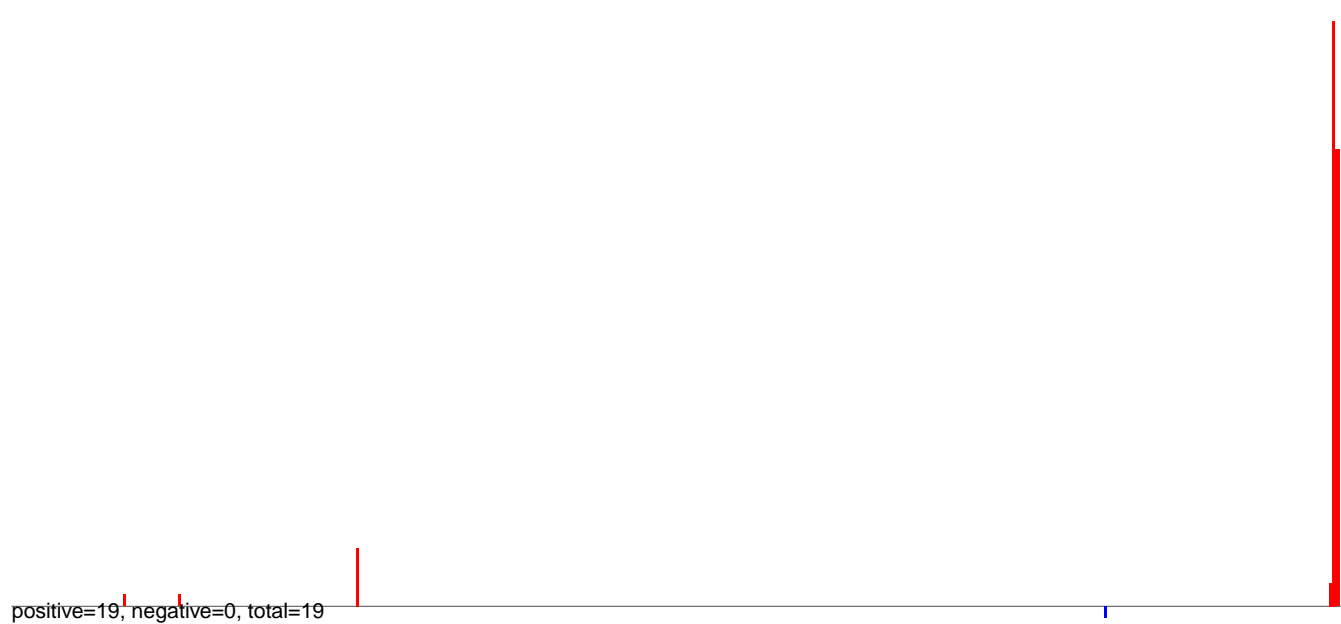


positive=2, negative=0, total=2

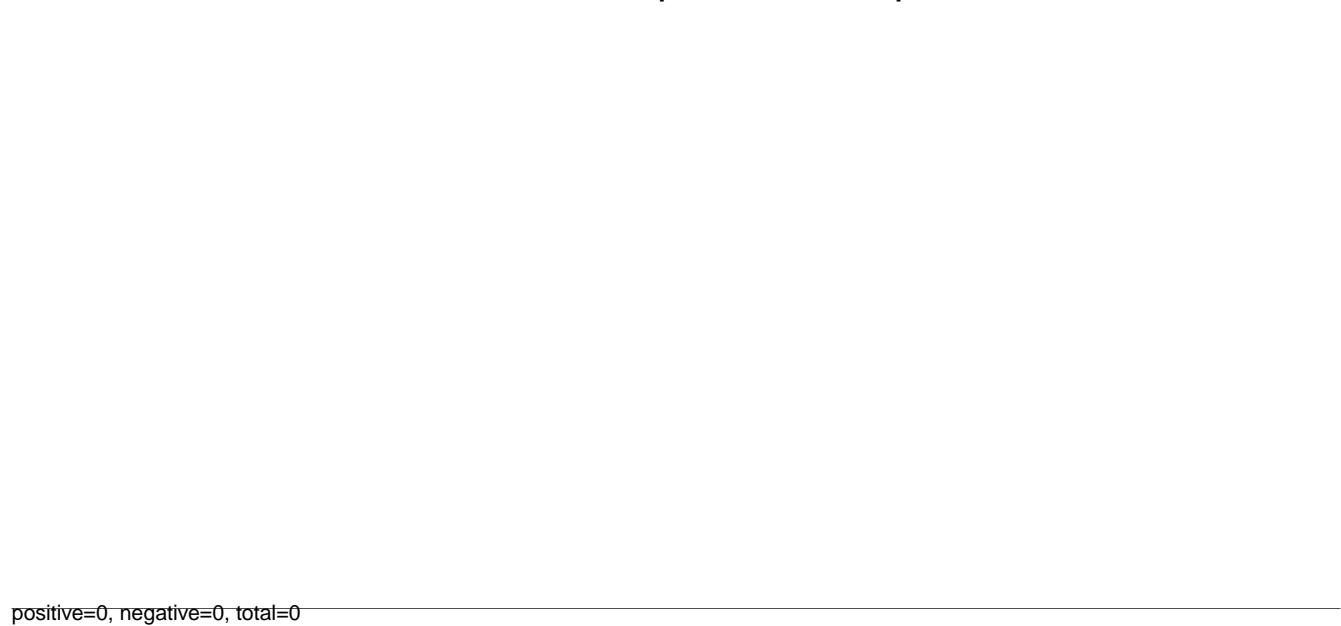
Window size=25, length=10560, virus@KM225263.1-Stratford_virus_strain_C338:1-10560

0 2000 4000 6000 8000 10000

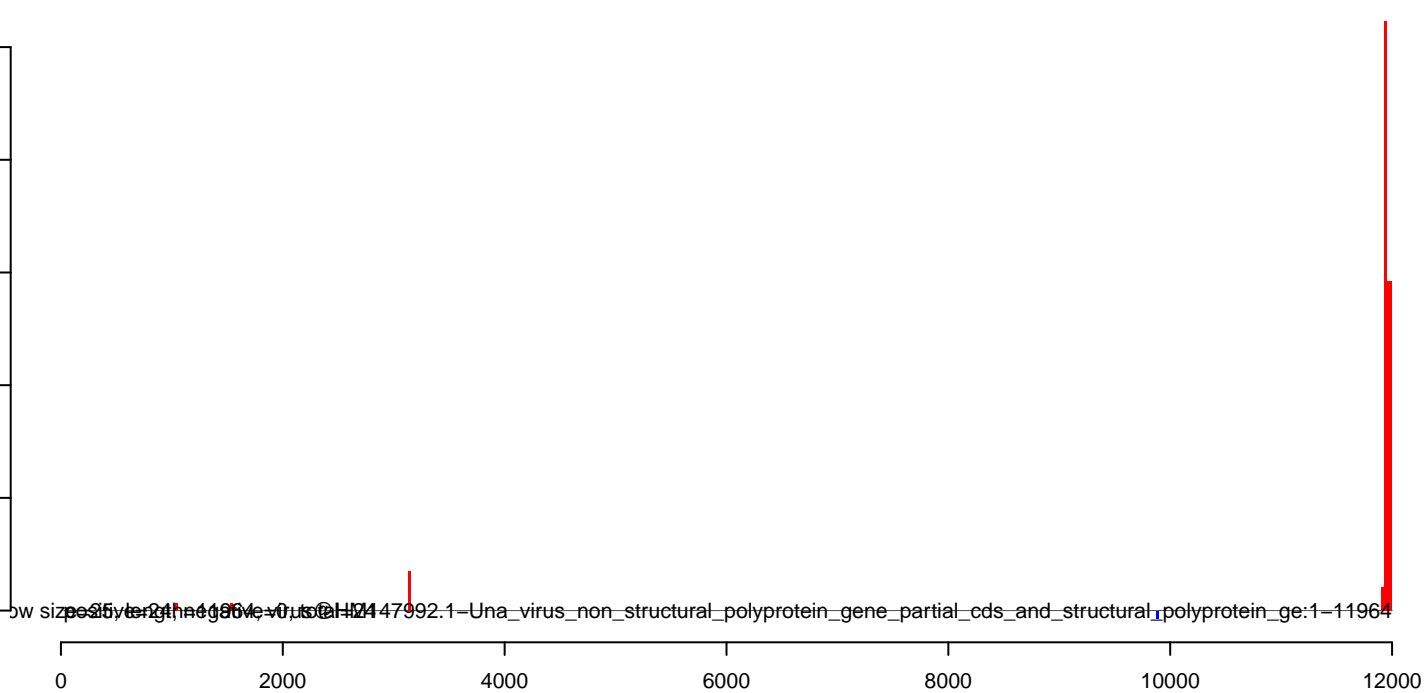
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



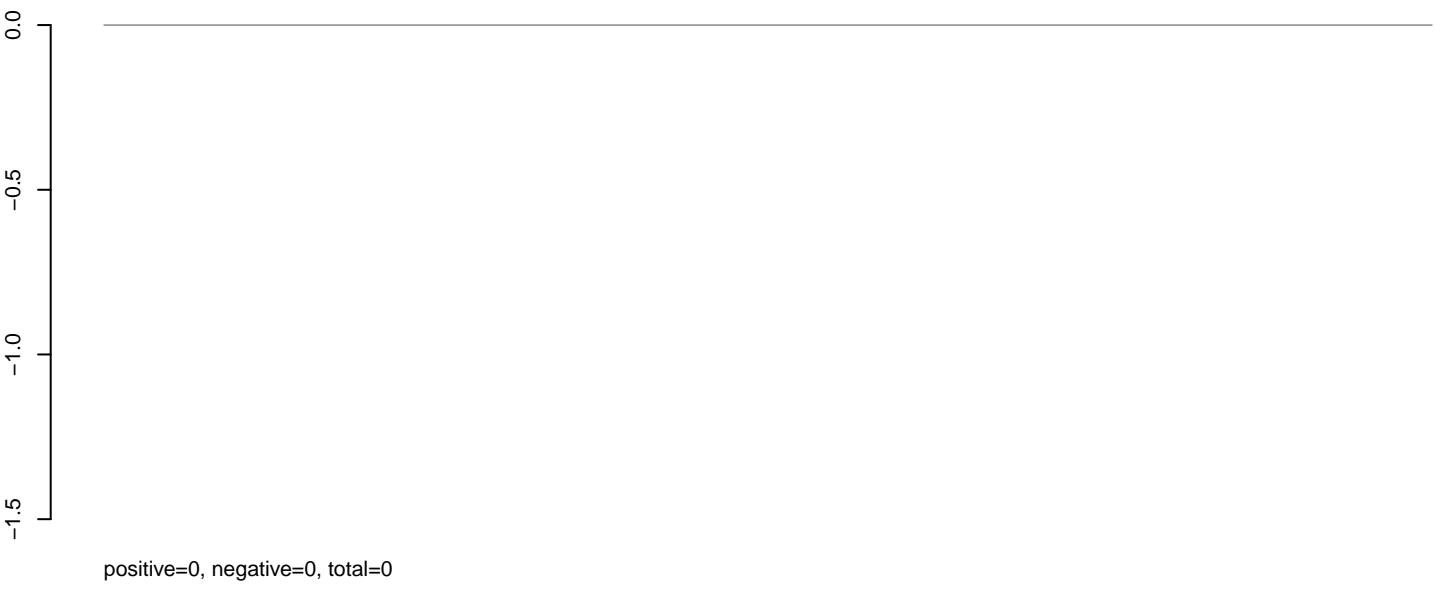
AeAlbo_C636_2dpi_mock3.rep



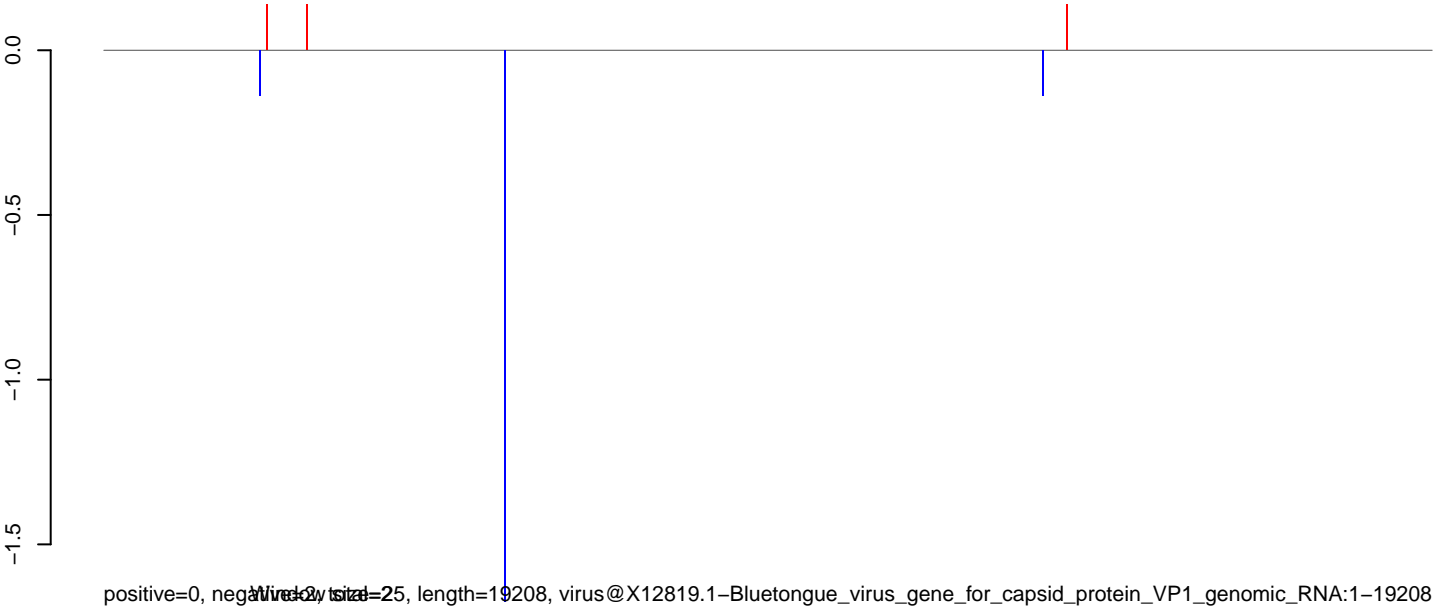
AeAlbo_C636_2dpi_mock3.18_23.rep



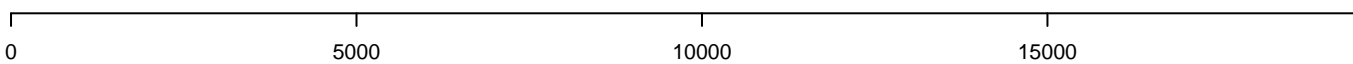
AeAlbo_C636_2dpi_mock3.24_35.rep



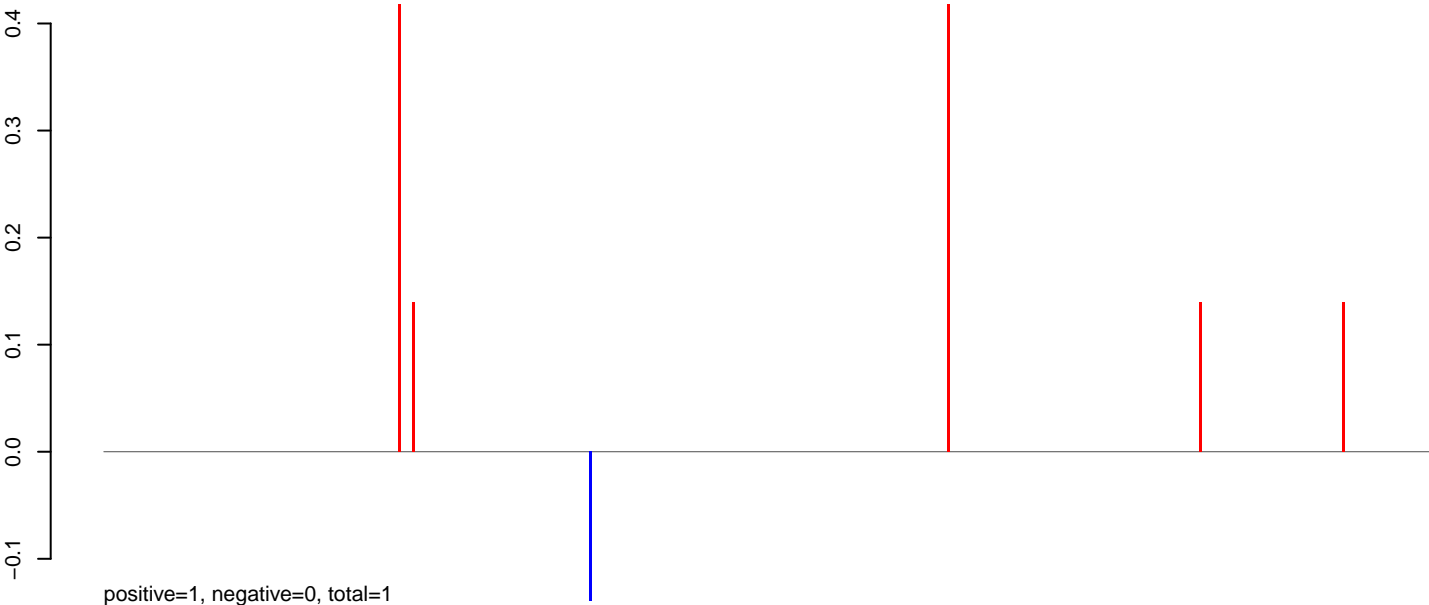
AeAlbo_C636_2dpi_mock3.rep



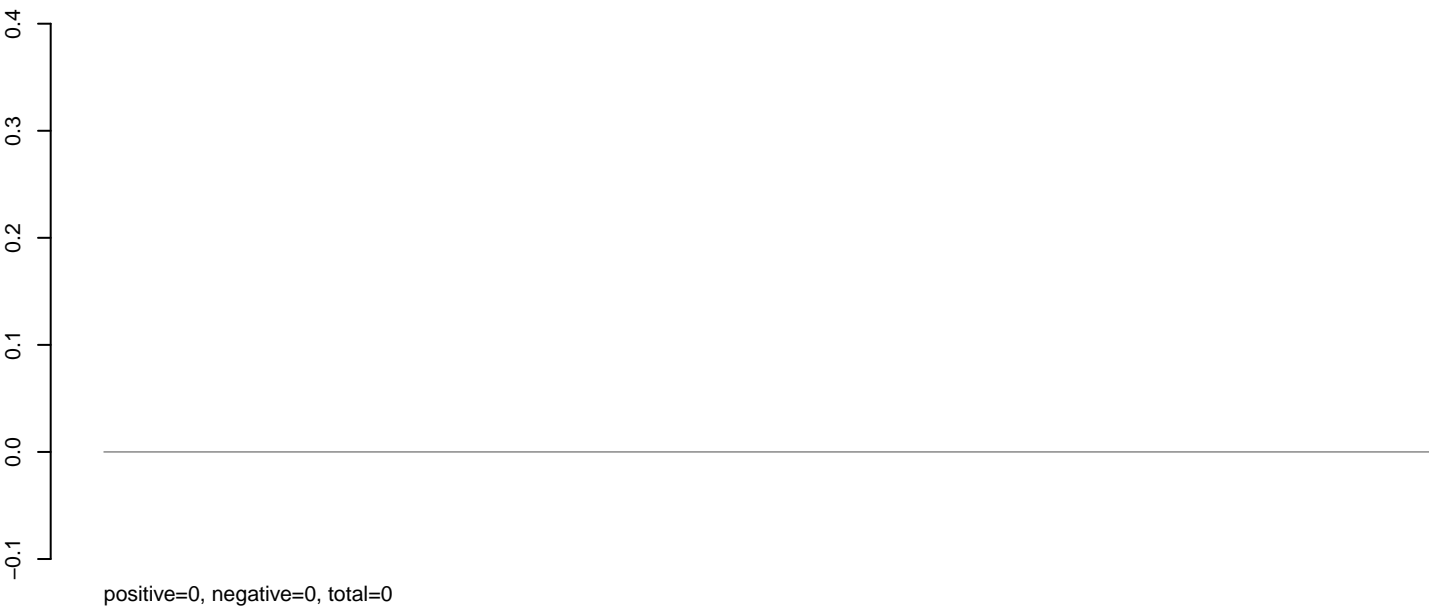
virus@X12819.1-Bluetongue_virus_gene_for_capsid_protein_VP1_genomic_RNA:1-19208



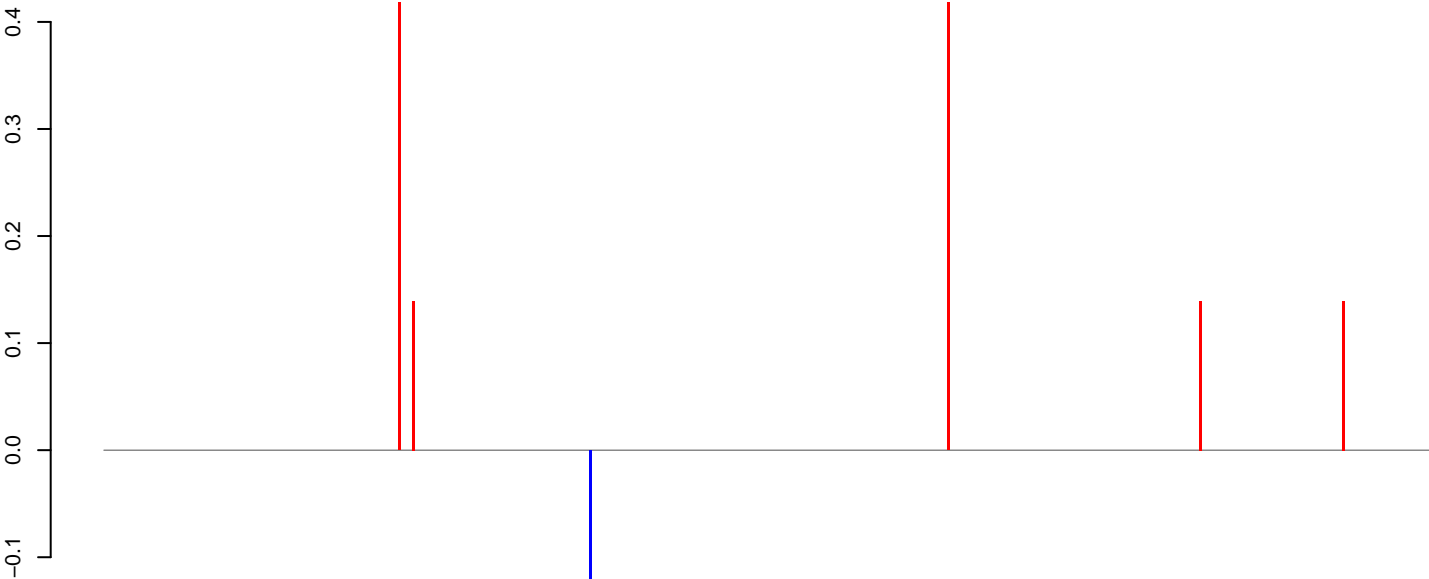
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



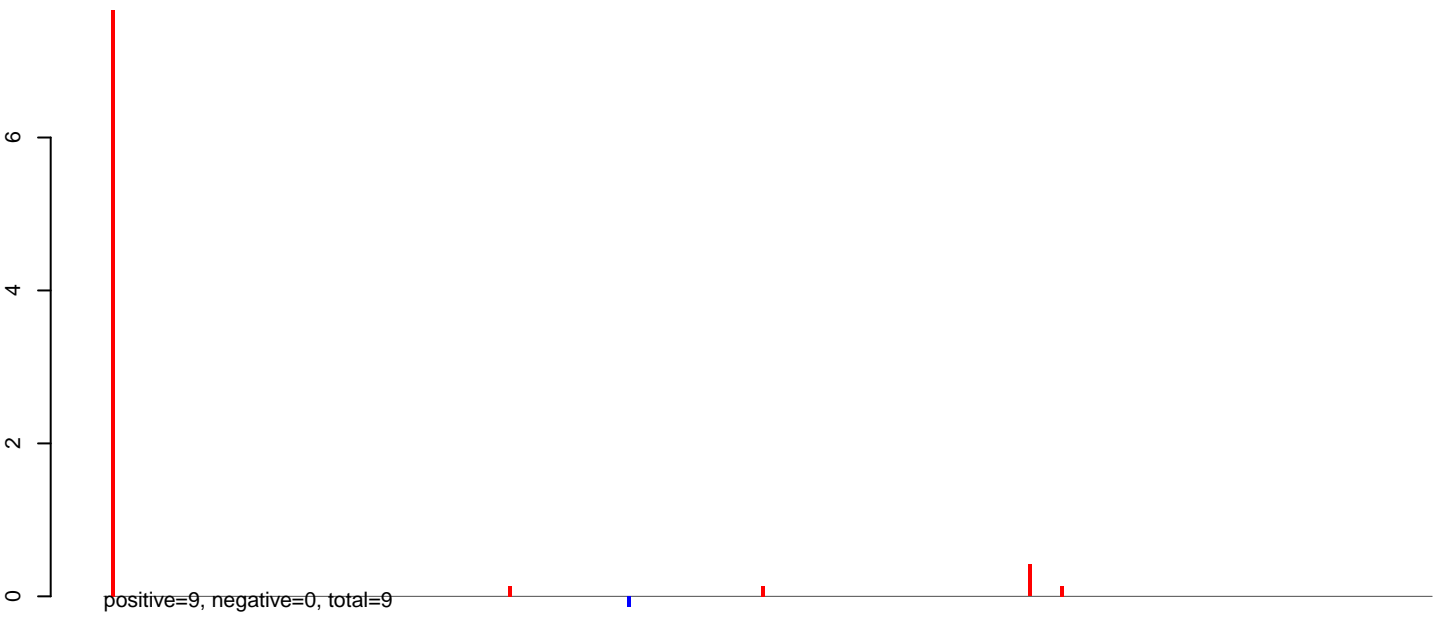
AeAlbo_C636_2dpi_mock3.rep



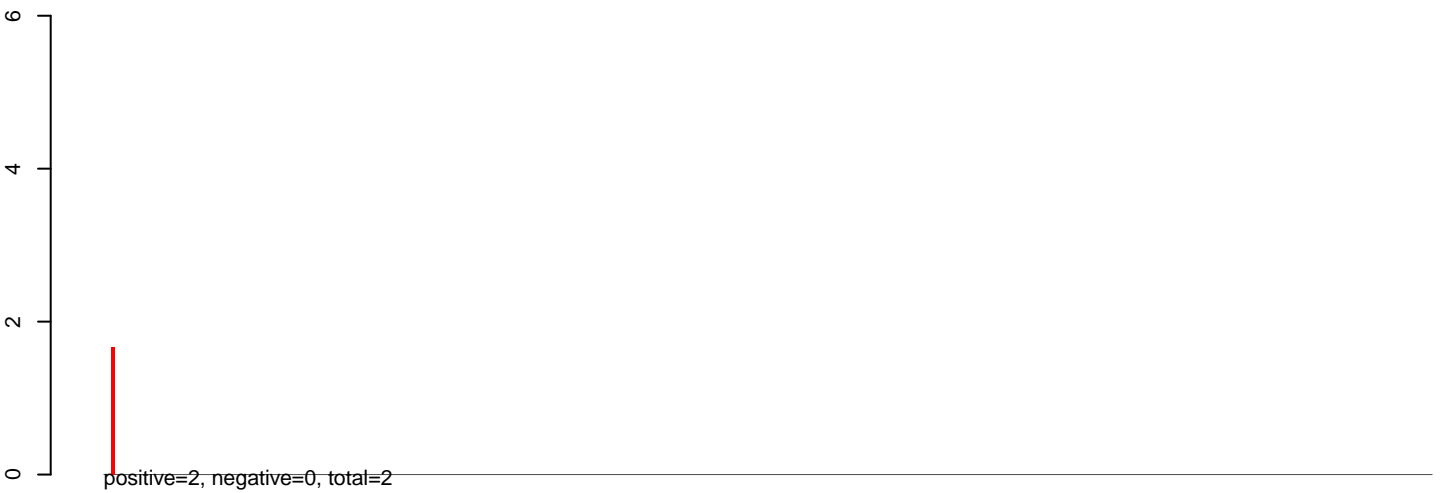
row size=25, length=14098, total=0 NC_038707.1-Jonchet_virus_strain_B81-CI-2004_segment_S_plus_segment_M_plus_segment_L:1-14098

0 2000 4000 6000 8000 10000 12000 14000

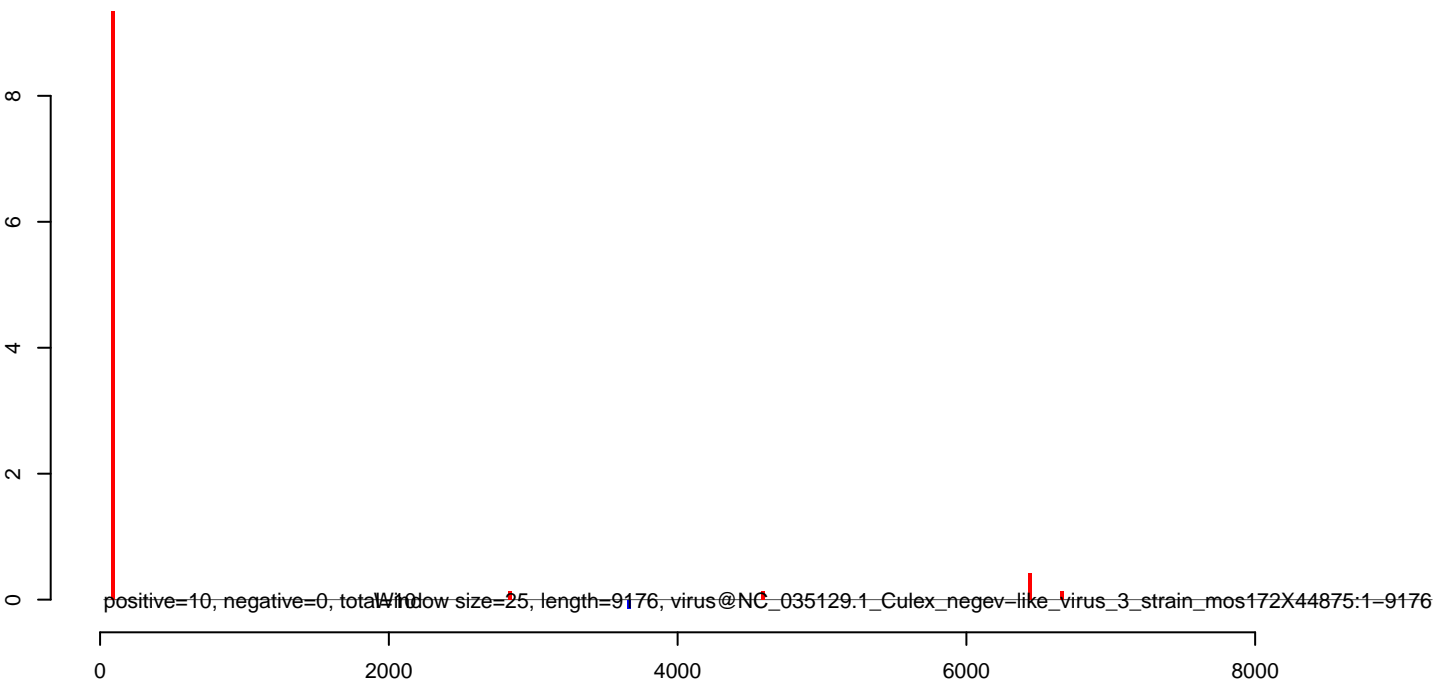
AeAlbo_C636_2dpi_mock3.18_23.rep



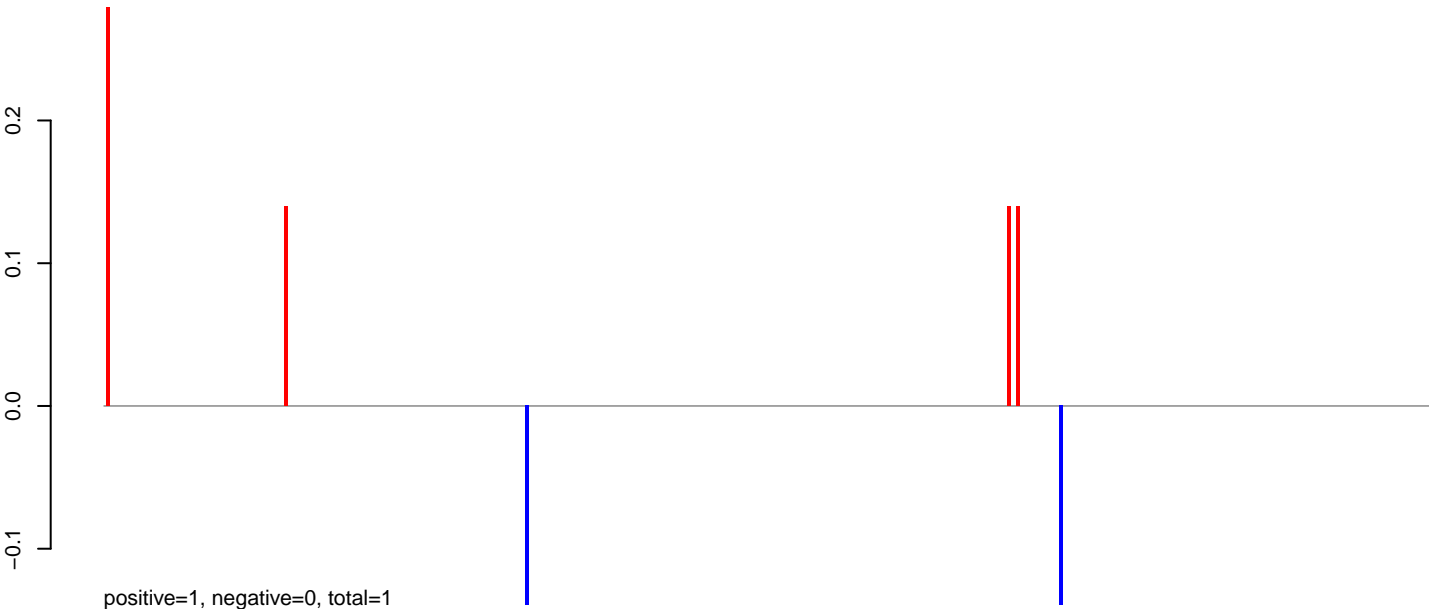
AeAlbo_C636_2dpi_mock3.24_35.rep



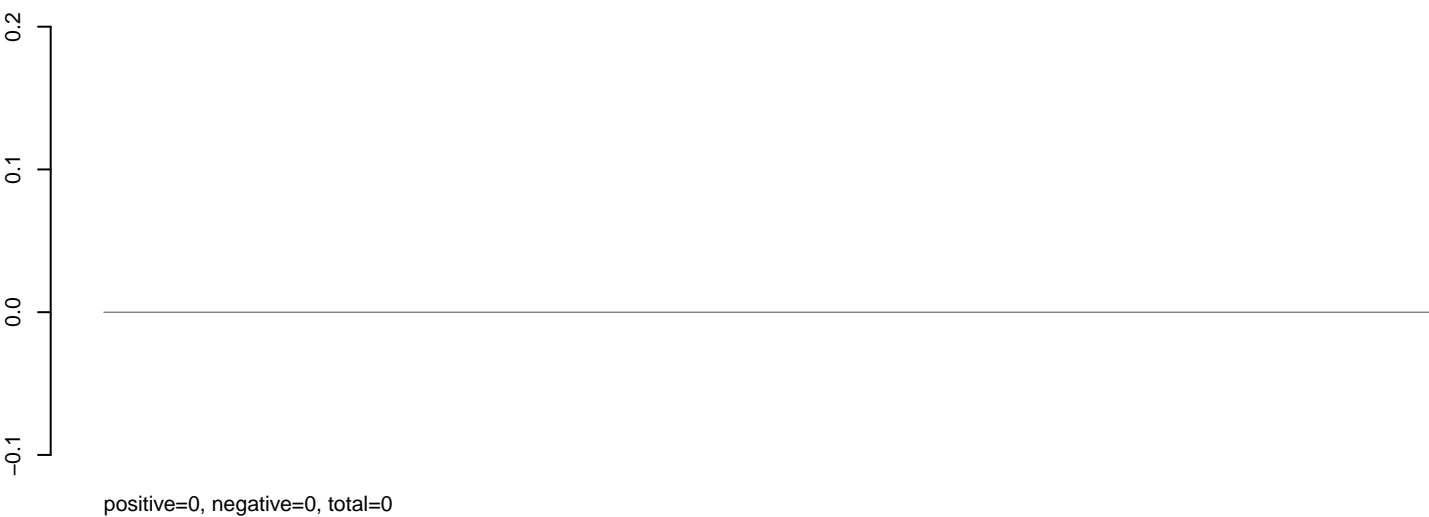
AeAlbo_C636_2dpi_mock3.rep



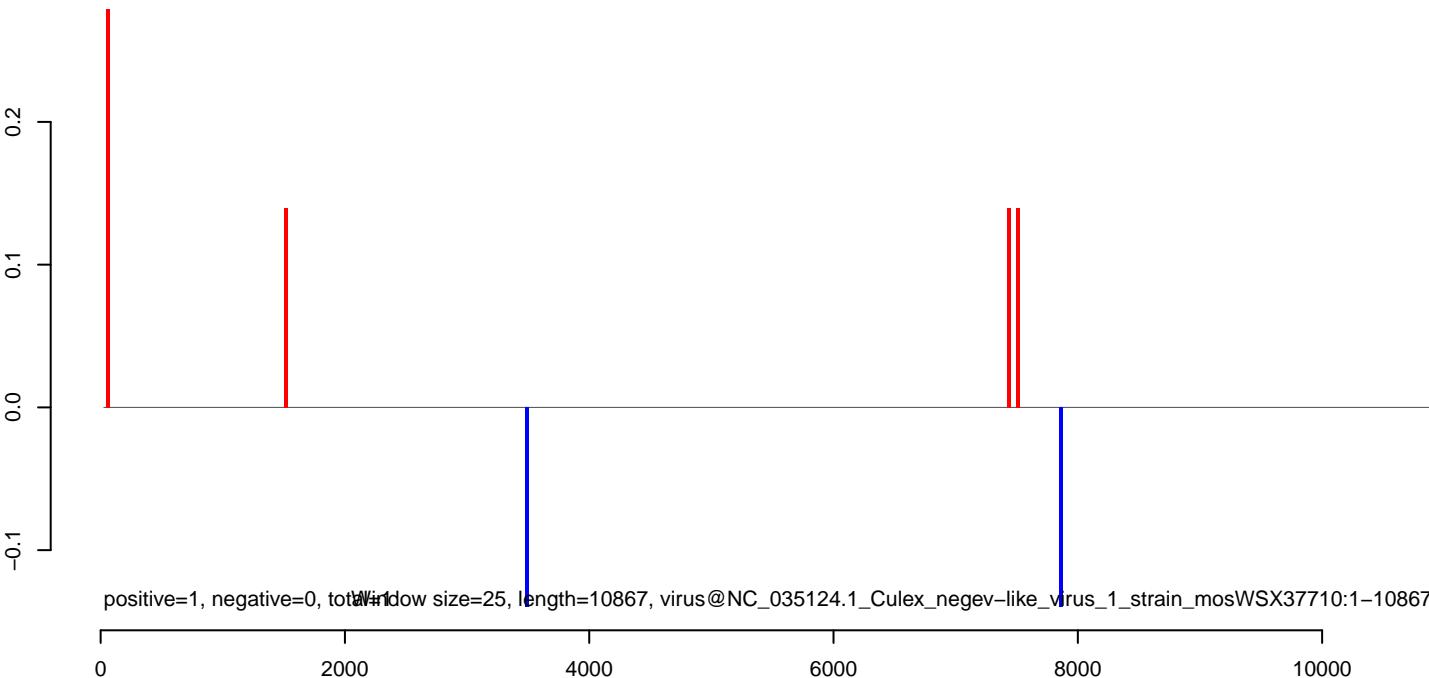
AeAlbo_C636_2dpi_mock3.18_23.rep



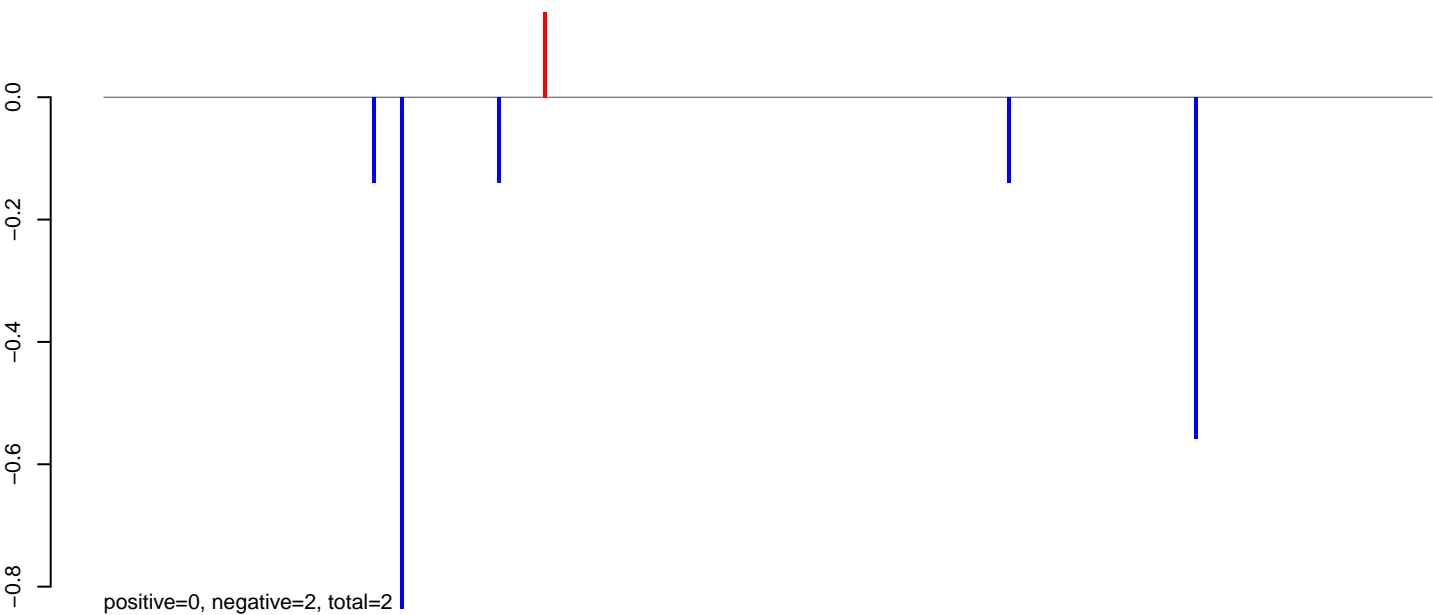
AeAlbo_C636_2dpi_mock3.24_35.rep



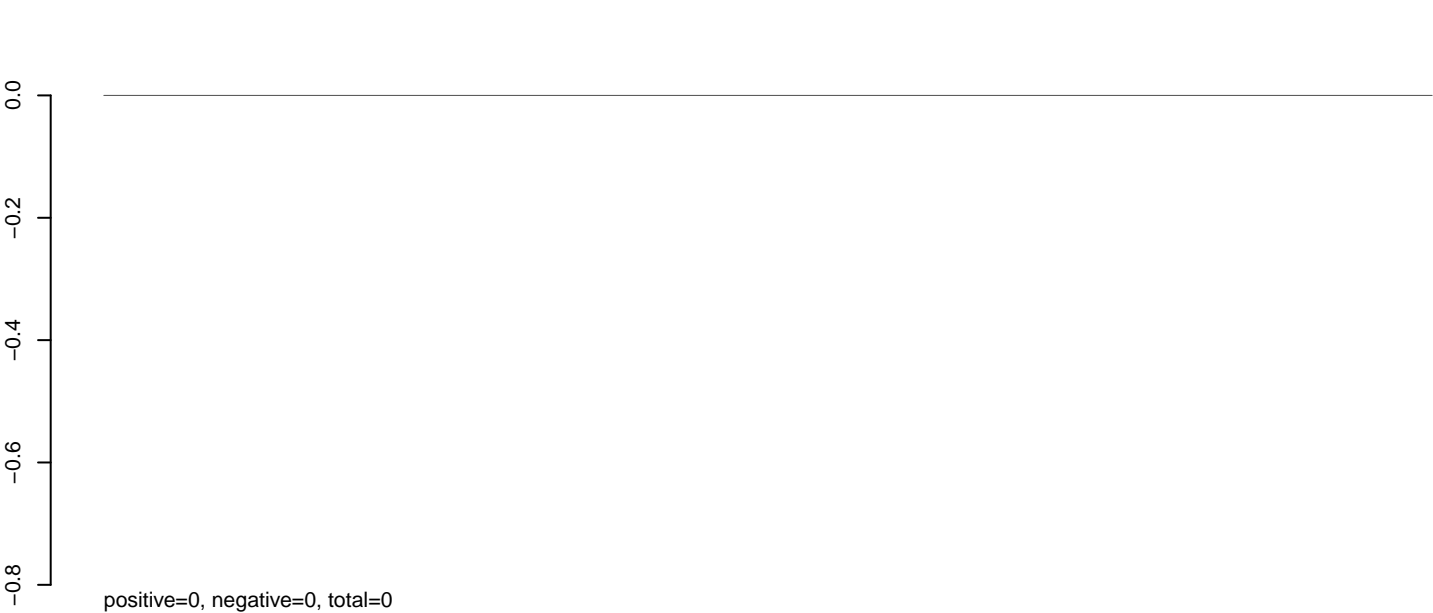
AeAlbo_C636_2dpi_mock3.rep



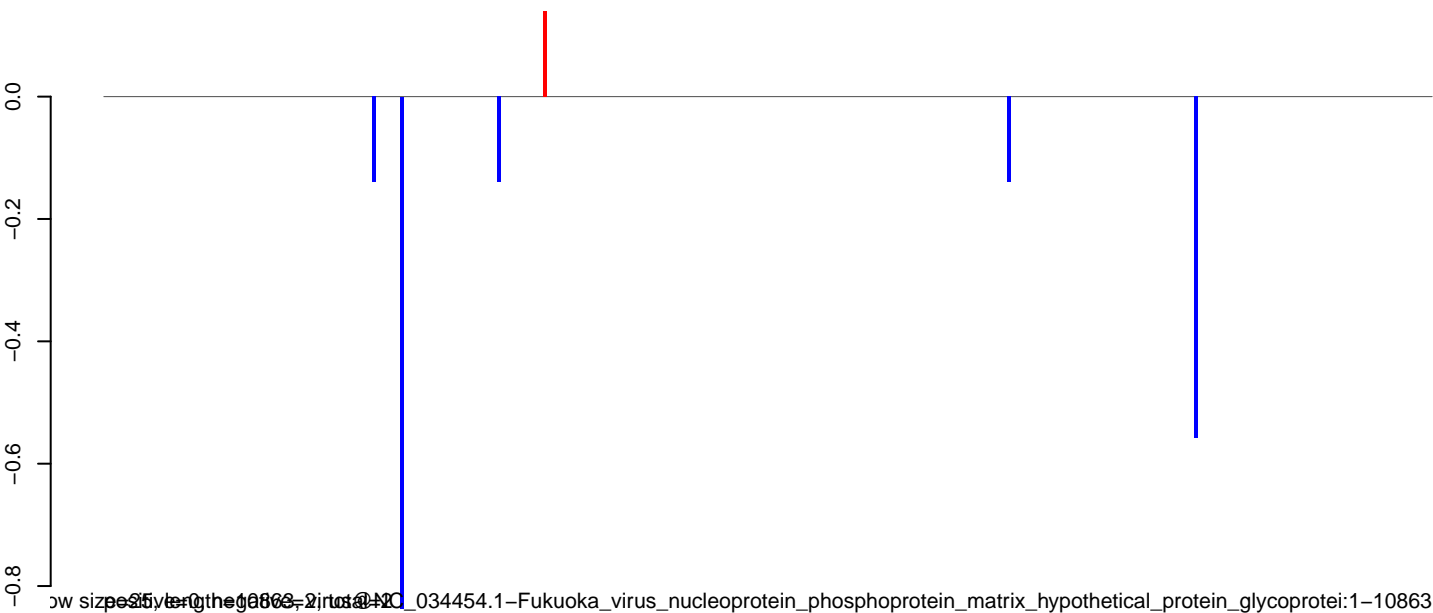
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



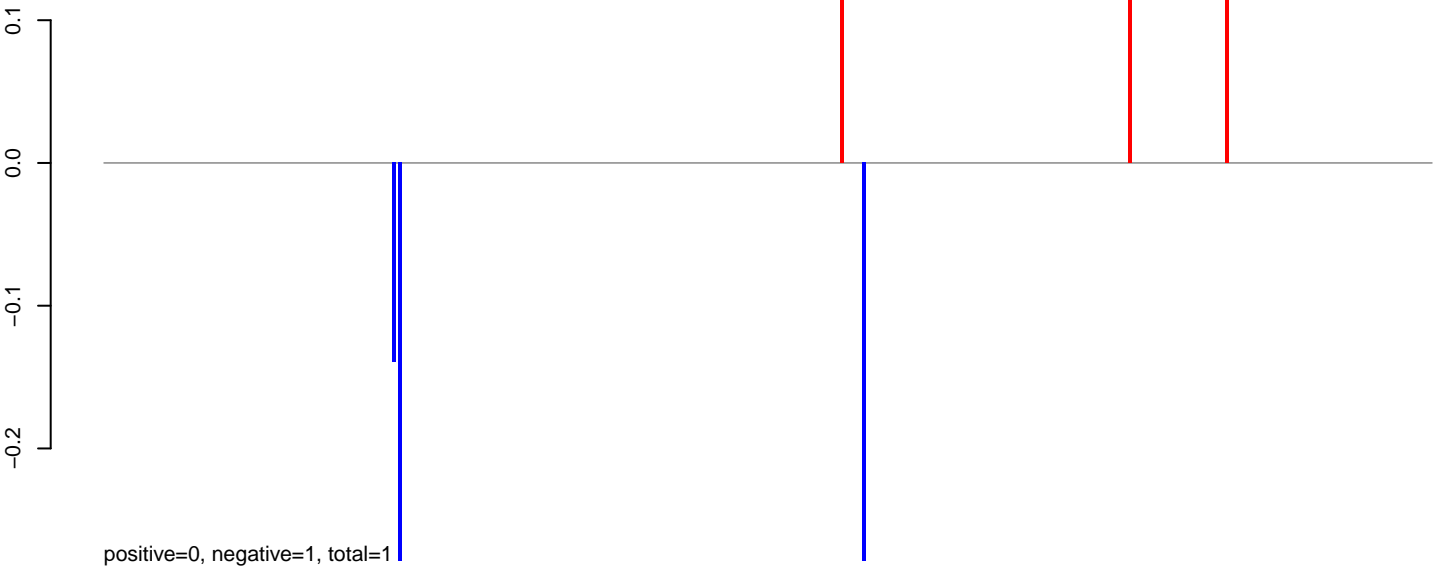
AeAlbo_C636_2dpi_mock3.rep



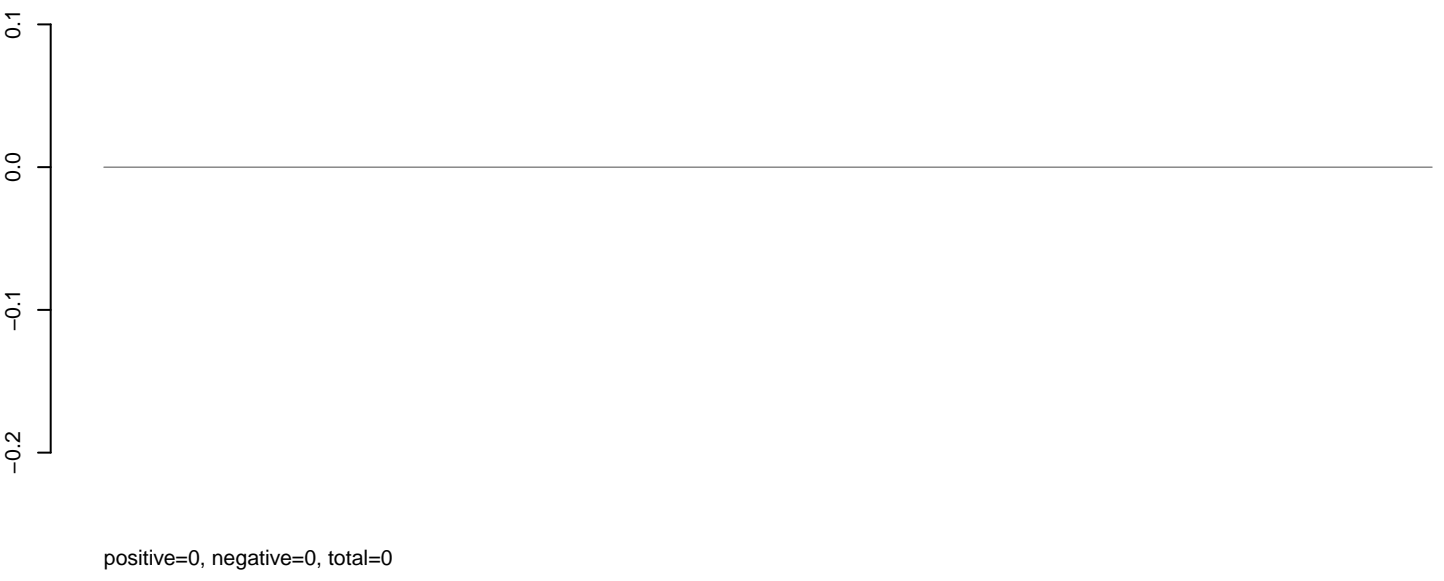
bow size style font the 1863-2 into 4 NC_034454.1-Fukuoka_virus_nucleoprotein_phosphoprotein_matrix_hypothetical_protein_glycoprotei:1-10863

0 2000 4000 6000 8000 10000

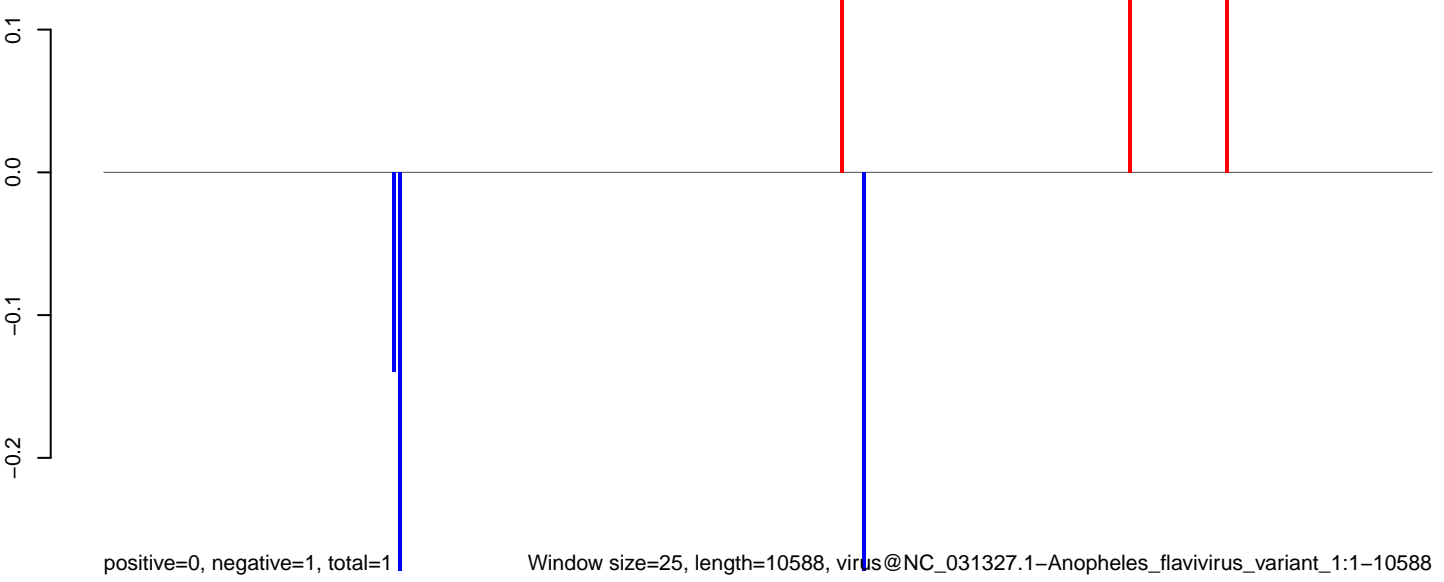
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



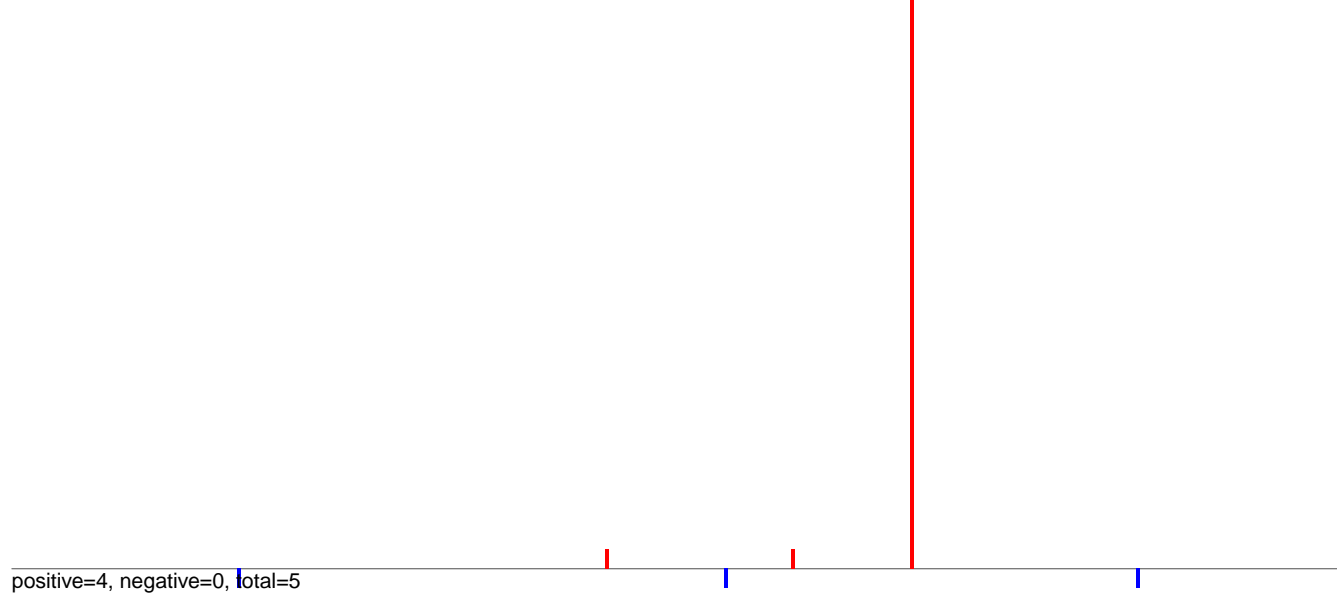
AeAlbo_C636_2dpi_mock3.rep



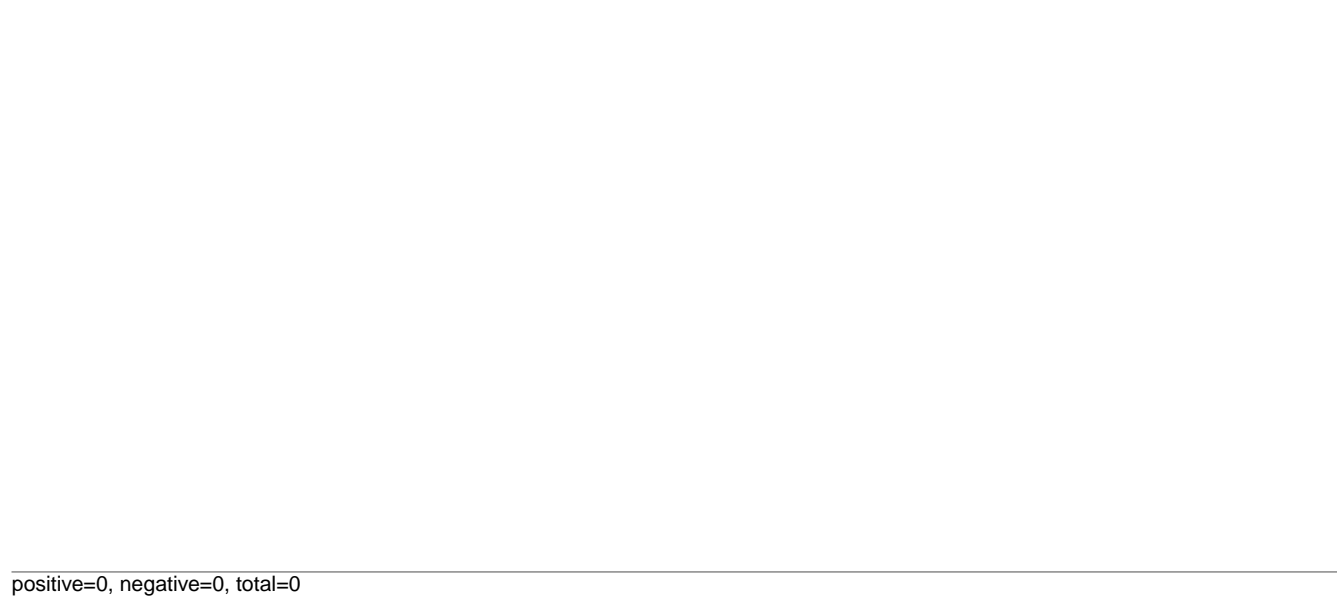
Window size=25, length=10588, virus@NC_031327.1-Anopheles_flavivirus_variant_1:1-10588

0 2000 4000 6000 8000 10000

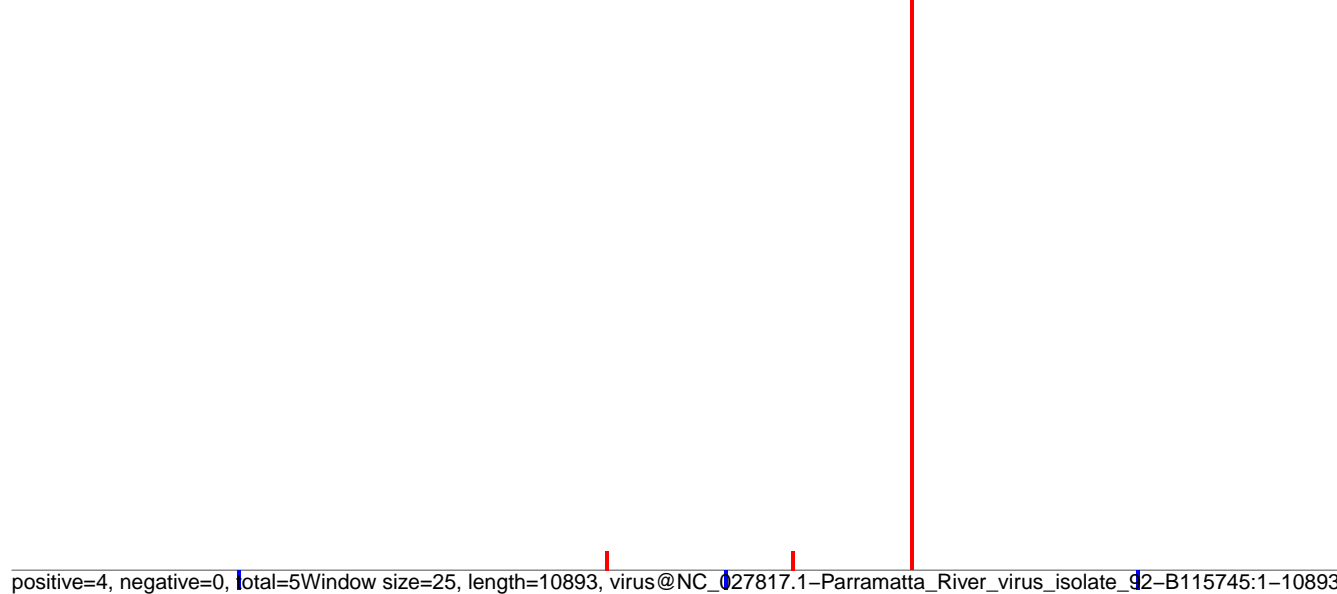
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



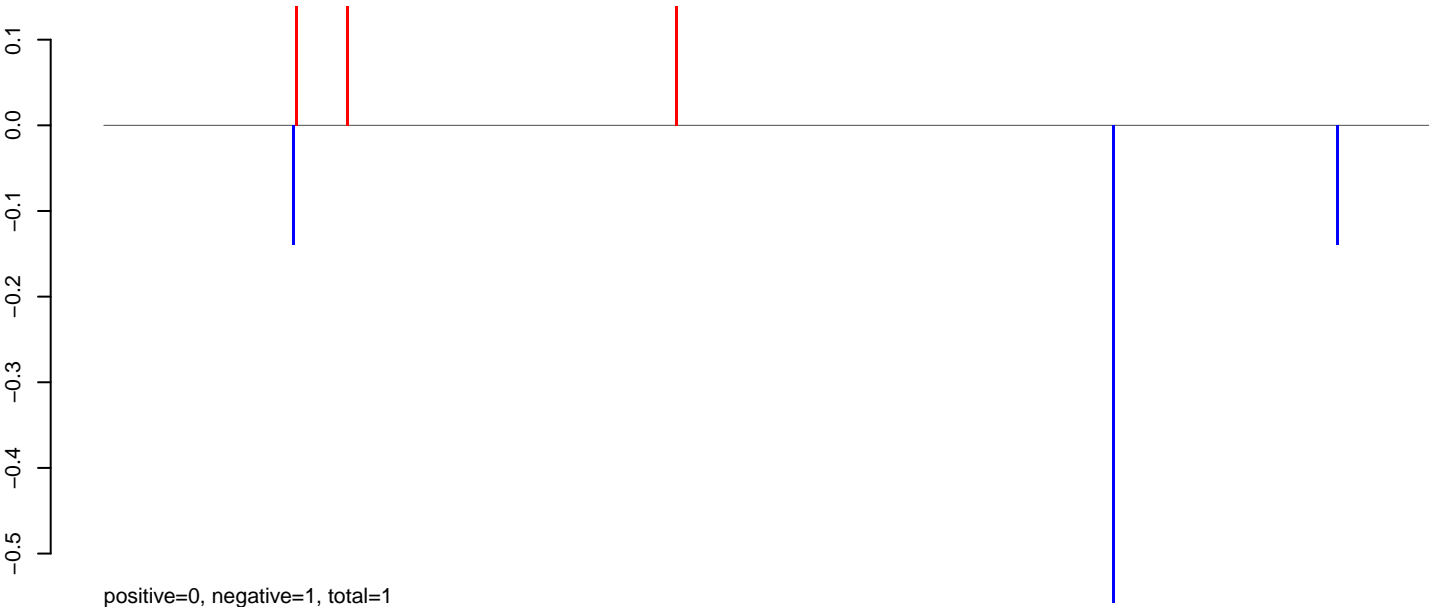
AeAlbo_C636_2dpi_mock3.rep



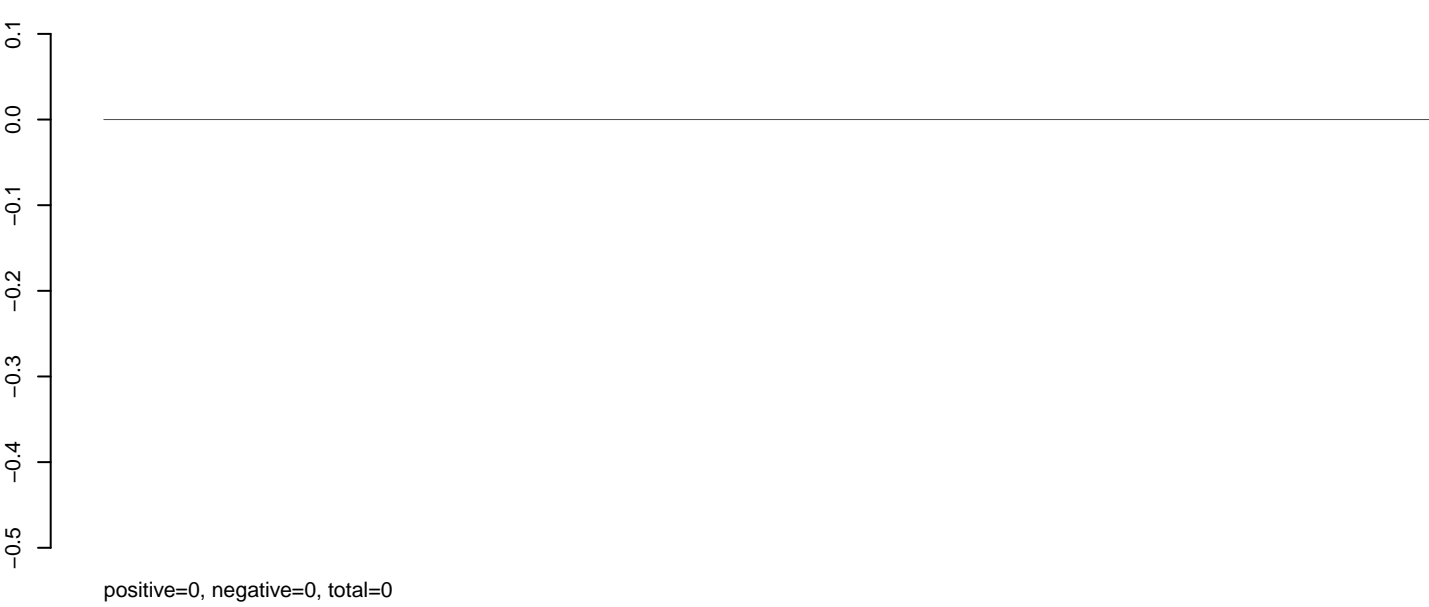
Window size=25, length=10893, virus@NC_027817.1-Parramatta_River_virus_isolate_92-B115745:1-10893

0 2000 4000 6000 8000 10000

AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



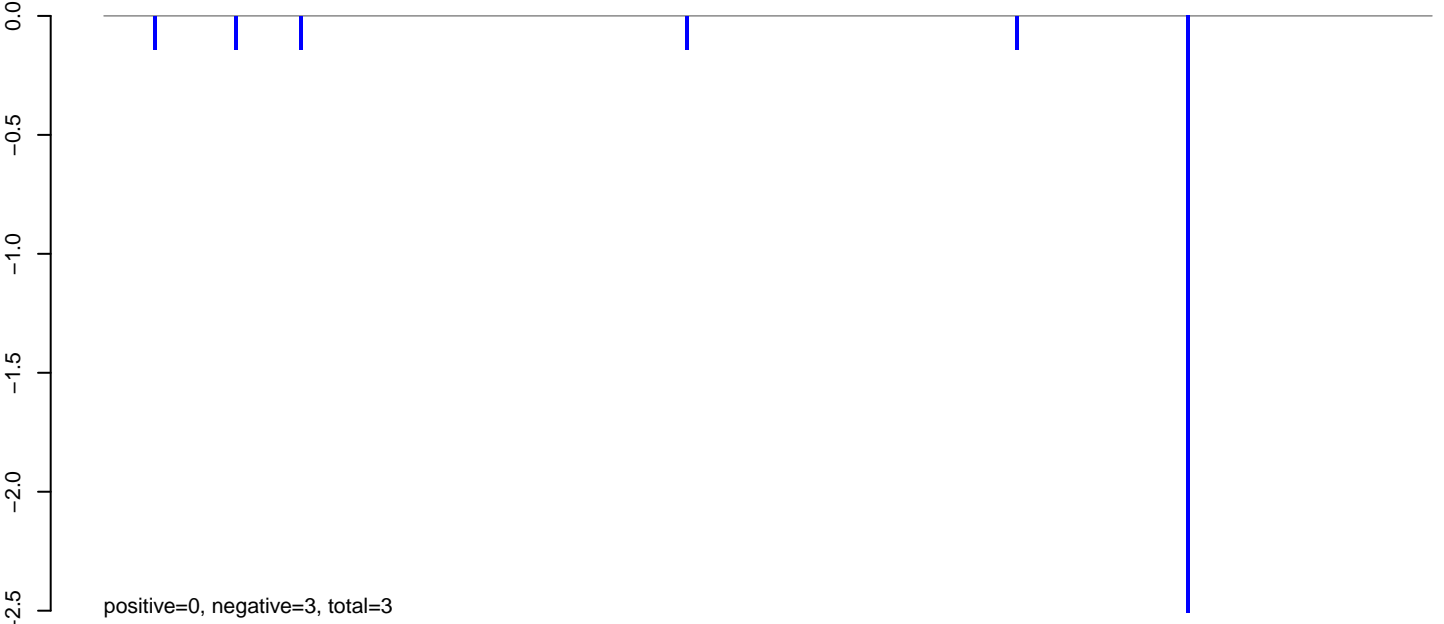
AeAlbo_C636_2dpi_mock3.rep



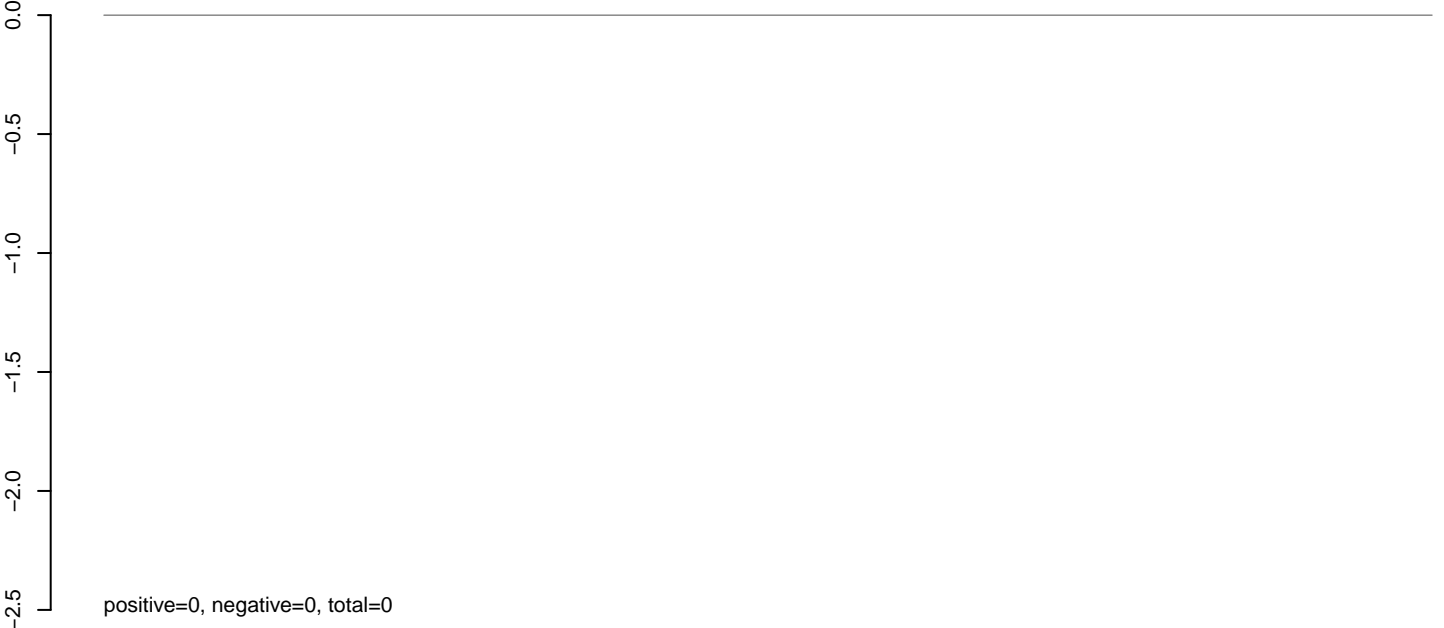
Window size=25, length=11097, virus@NC_025356.1-Pike_fry_rhabdovirus_isolate_F4:1-11097

0 2000 4000 6000 8000 10000

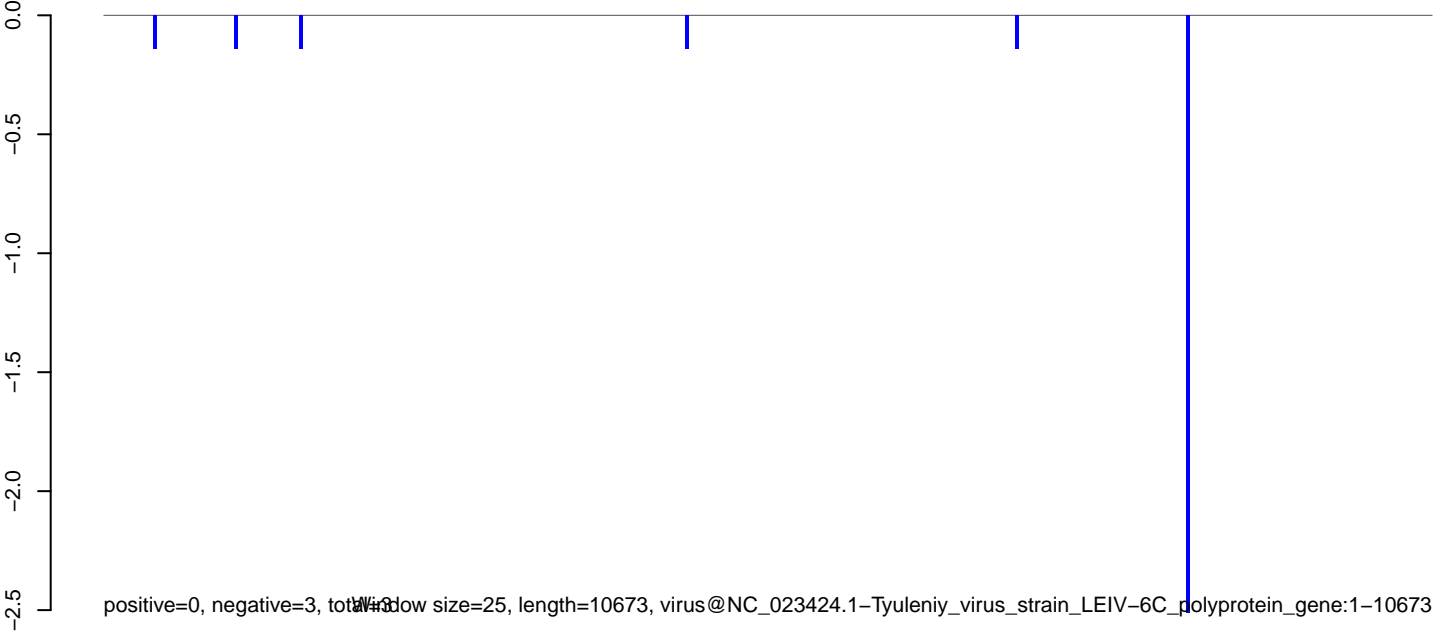
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



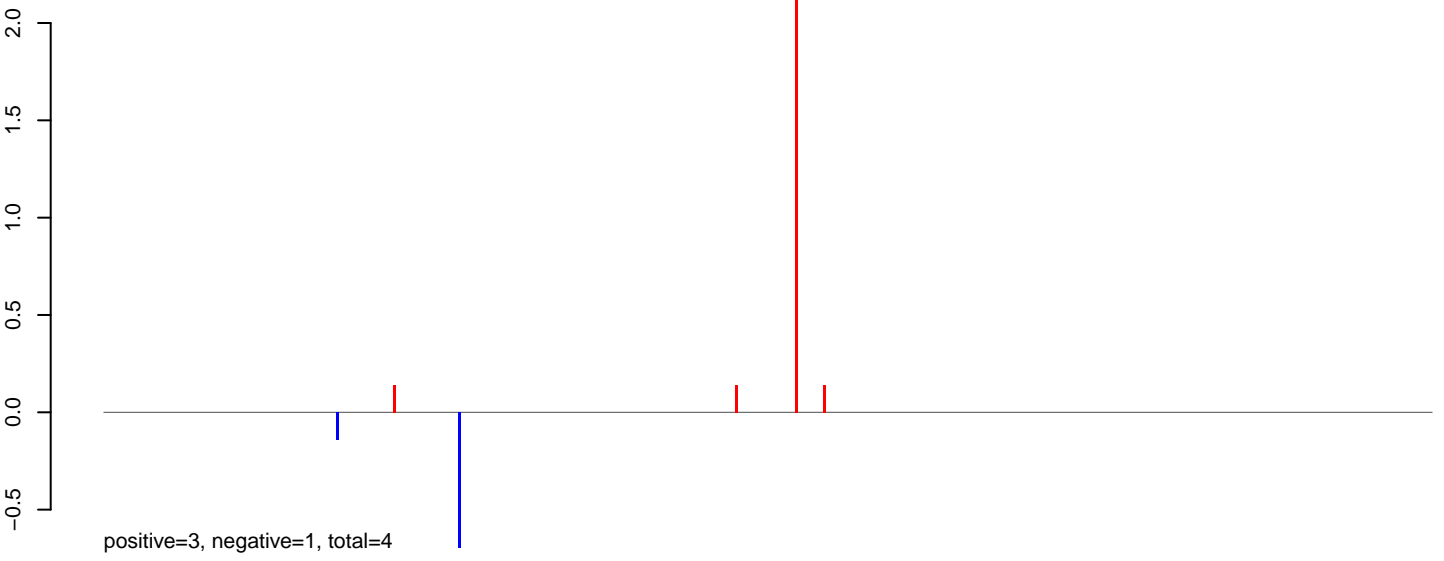
AeAlbo_C636_2dpi_mock3.rep



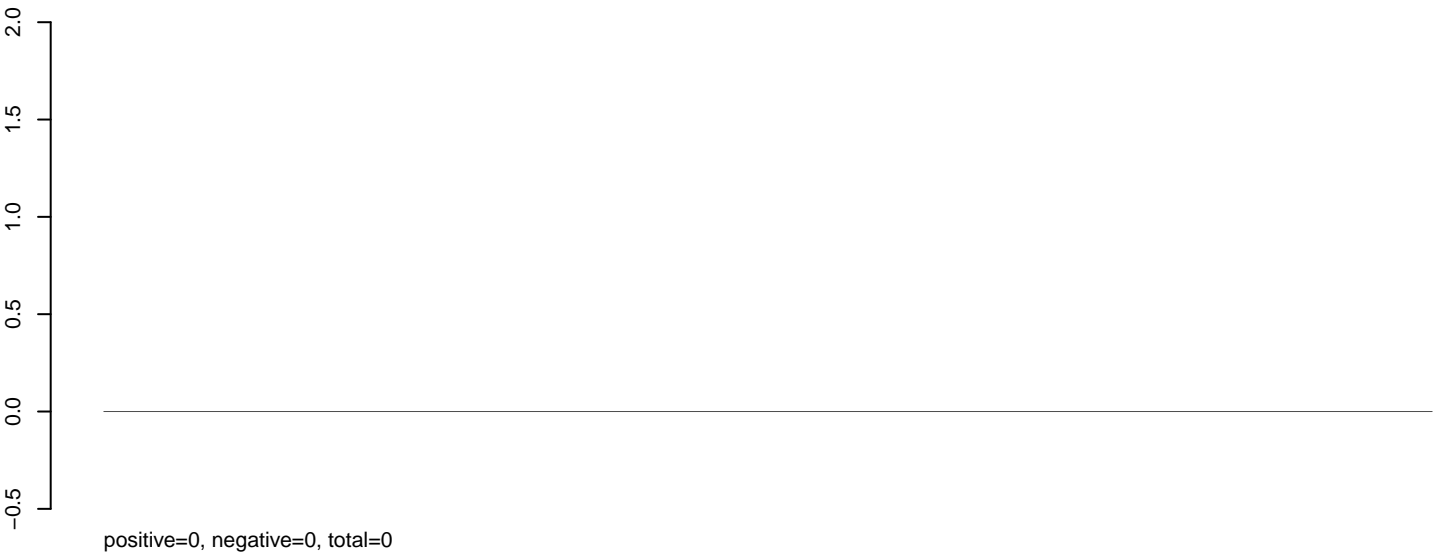
Window size=25, length=10673, virus@NC_023424.1-Tyuleniy_virus_strain_LEIV-6C_polyprotein_gene:1-10673

0 2000 4000 6000 8000 10000

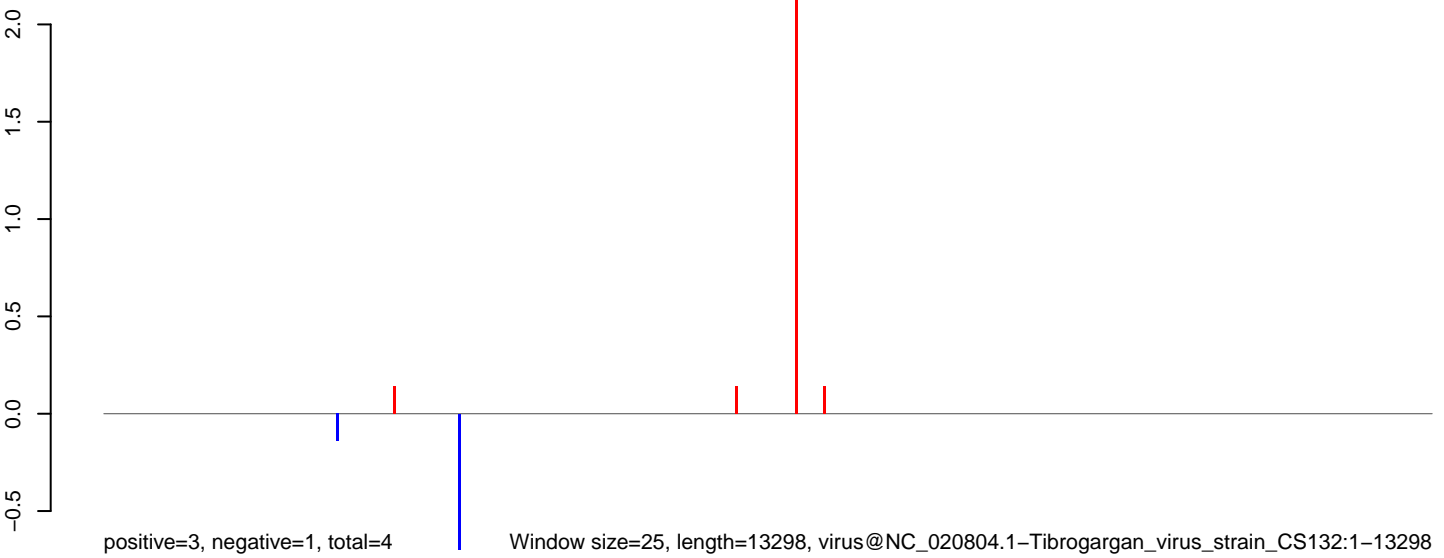
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



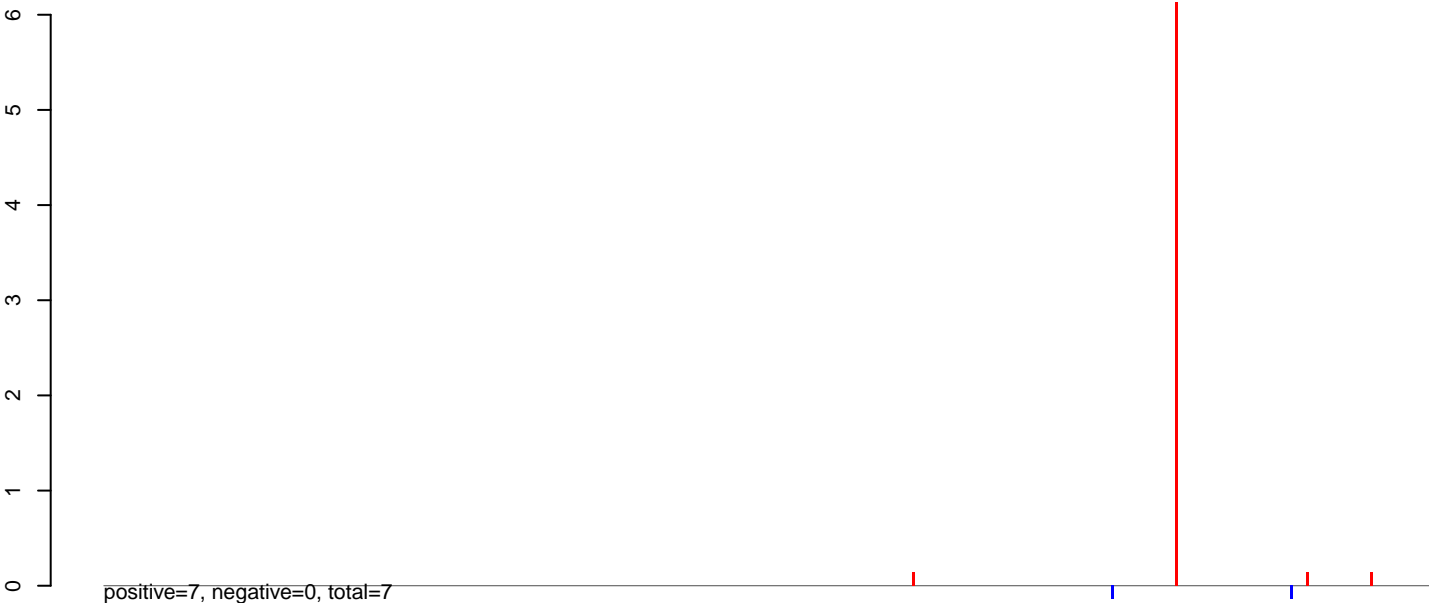
AeAlbo_C636_2dpi_mock3.rep



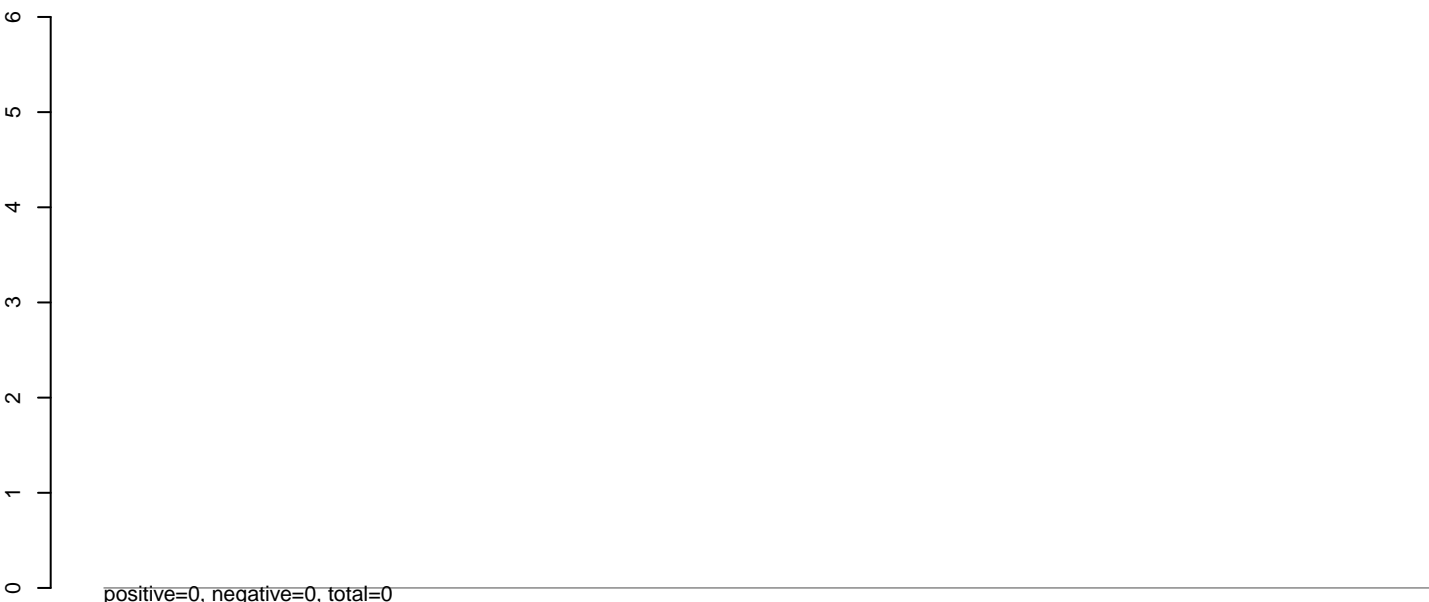
Window size=25, length=13298, virus@NC_020804.1-Tibrogargan_virus_strain_CS132:1-13298

0 2000 4000 6000 8000 10000 12000

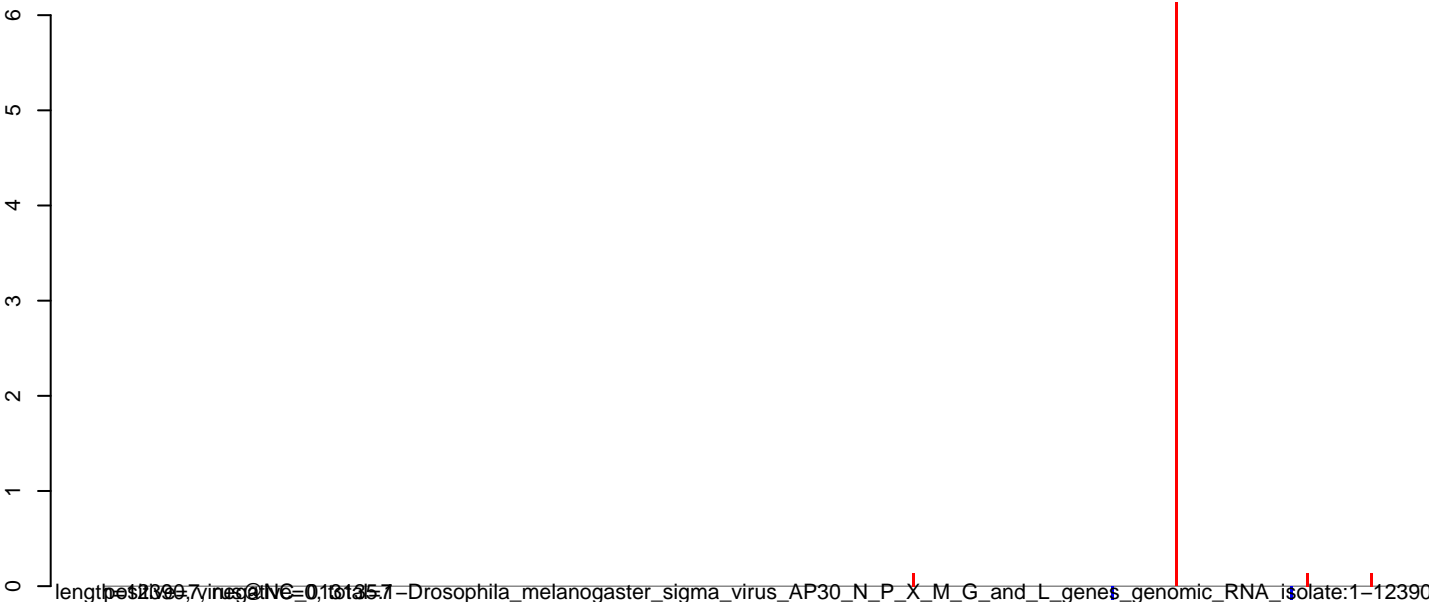
AeAlbo_C636_2dpi_mock3.18_23.rep



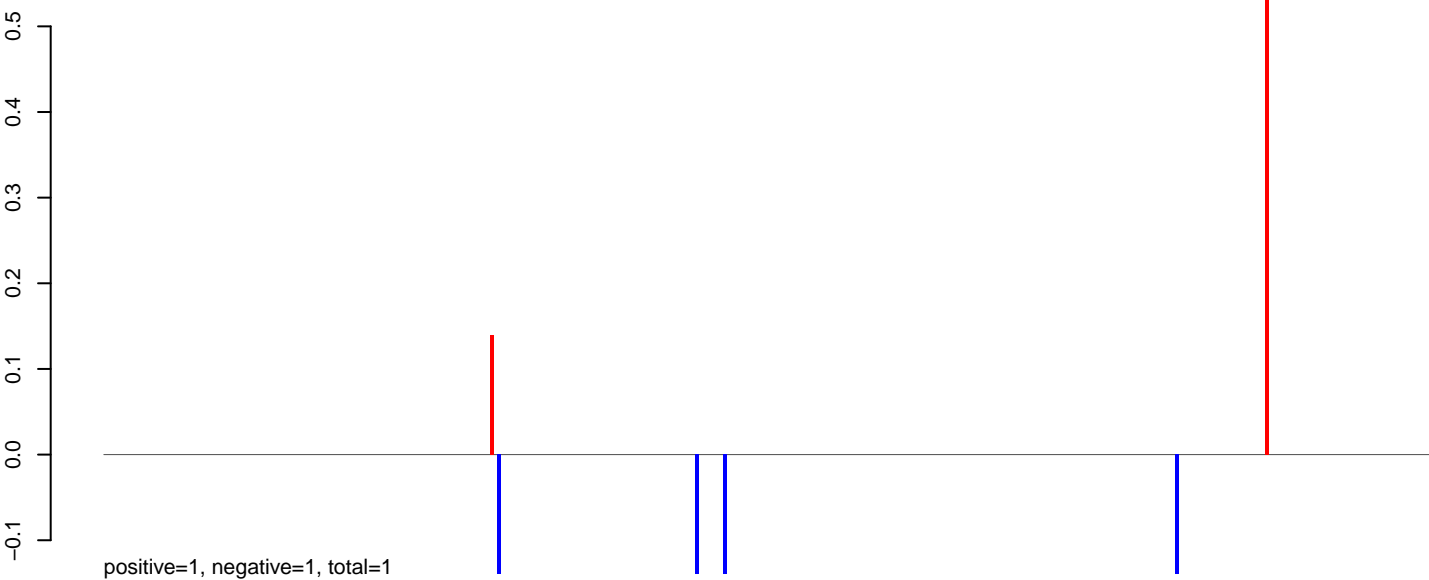
AeAlbo_C636_2dpi_mock3.24_35.rep



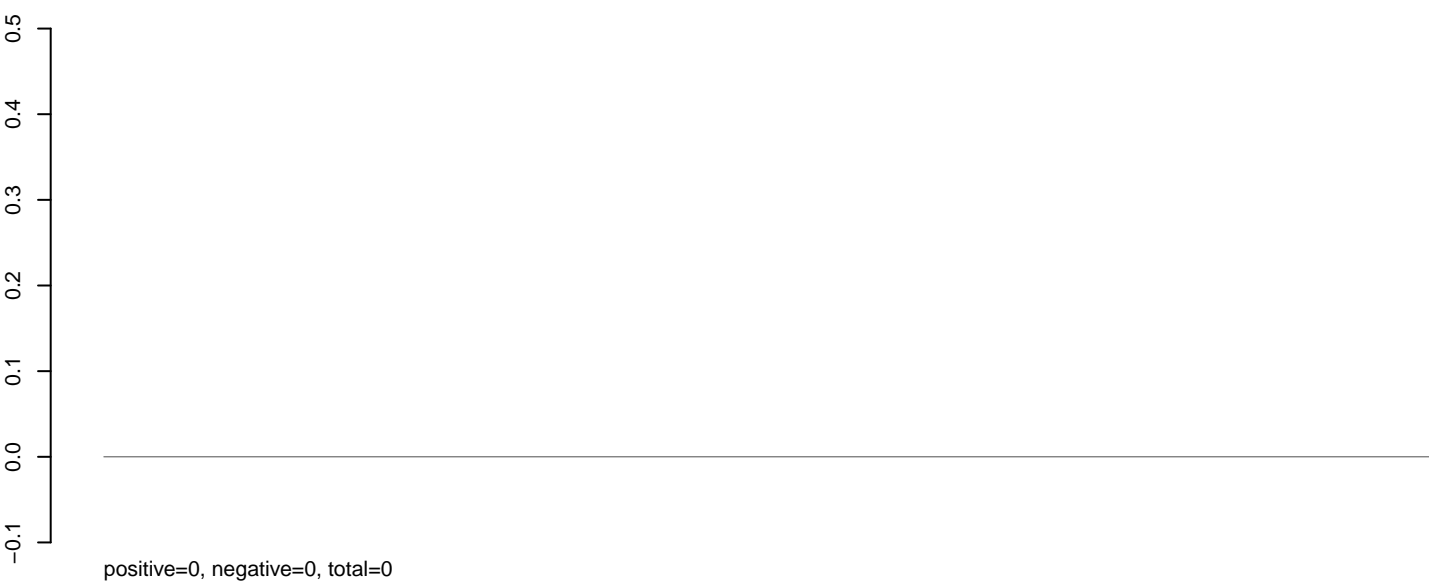
AeAlbo_C636_2dpi_mock3.rep



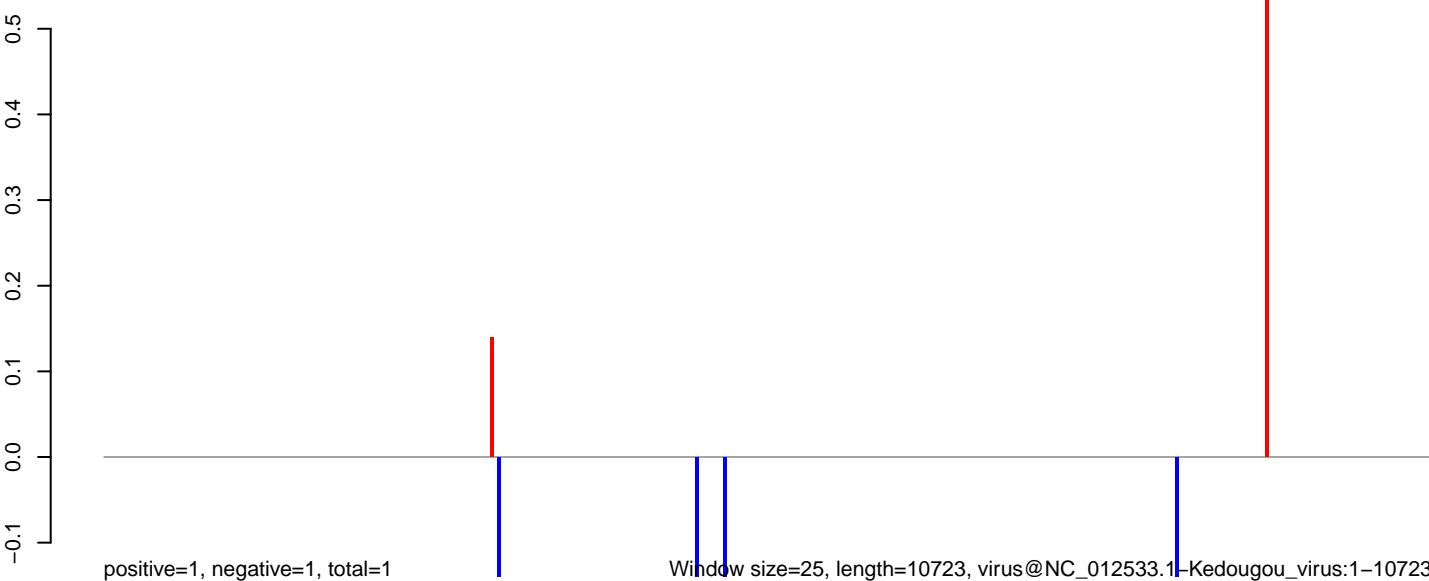
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep

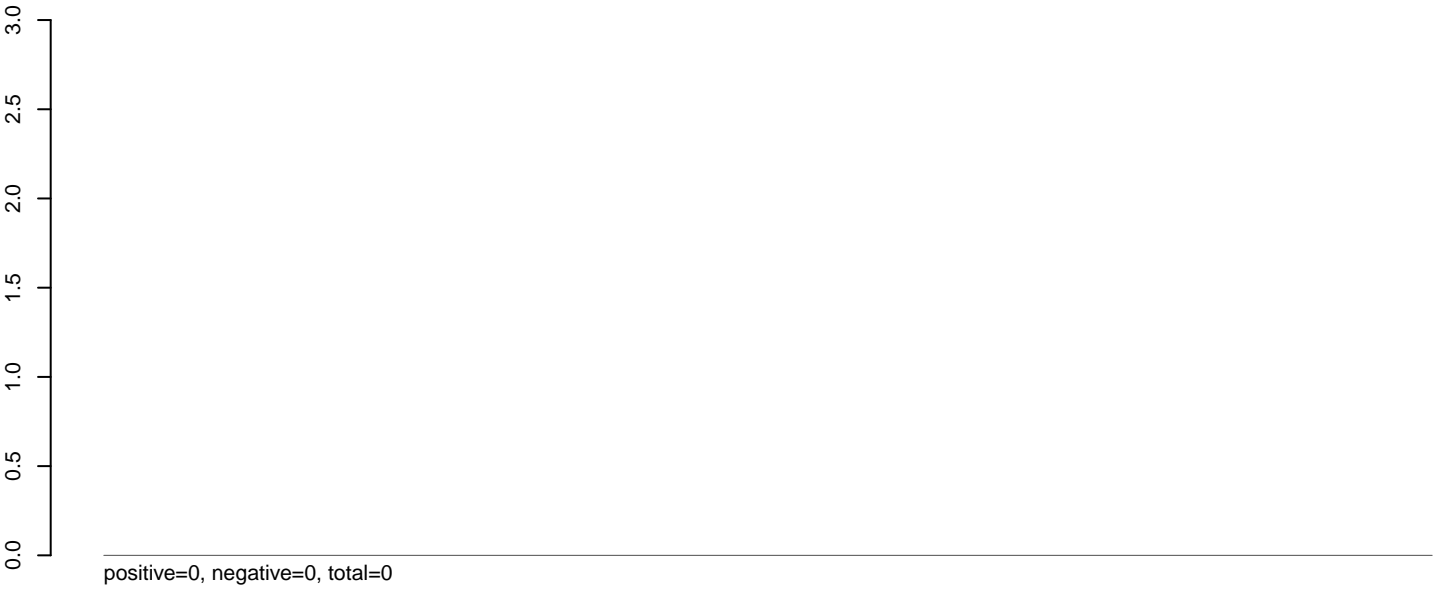


0 2000 4000 6000 8000 10000

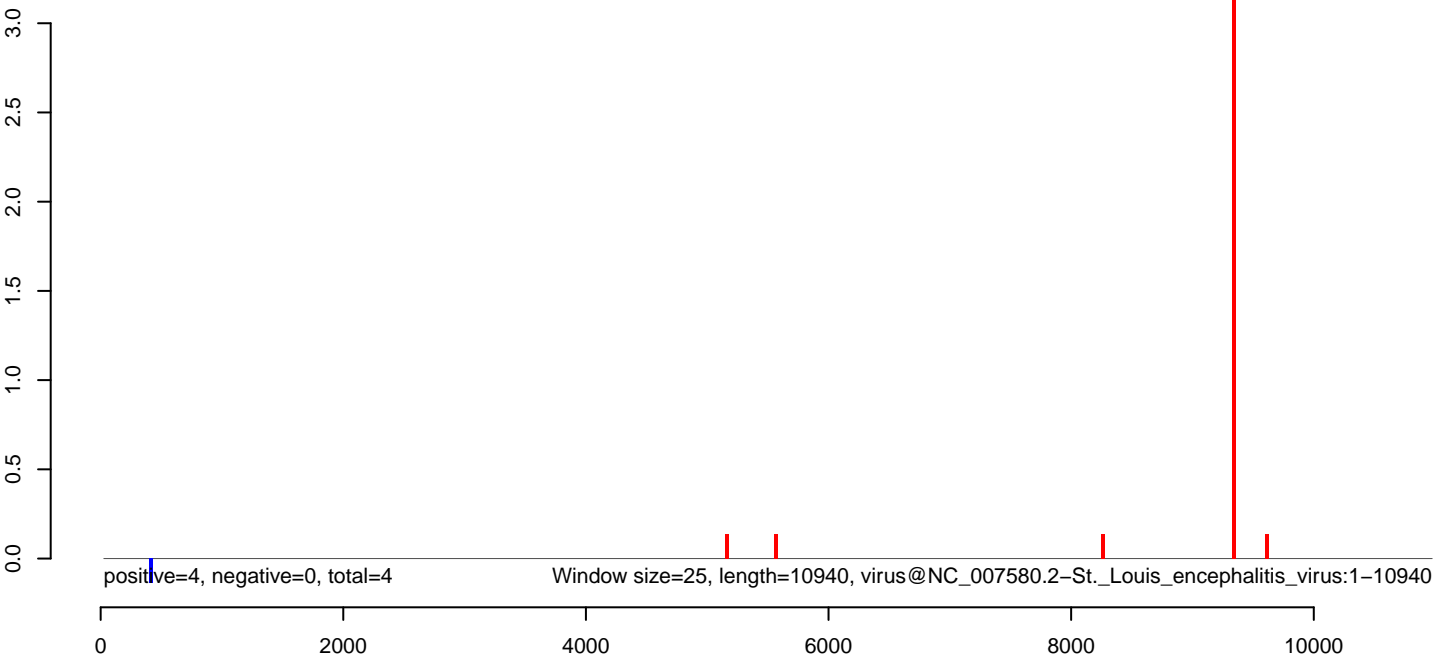
AeAlbo_C636_2dpi_mock3.18_23.rep



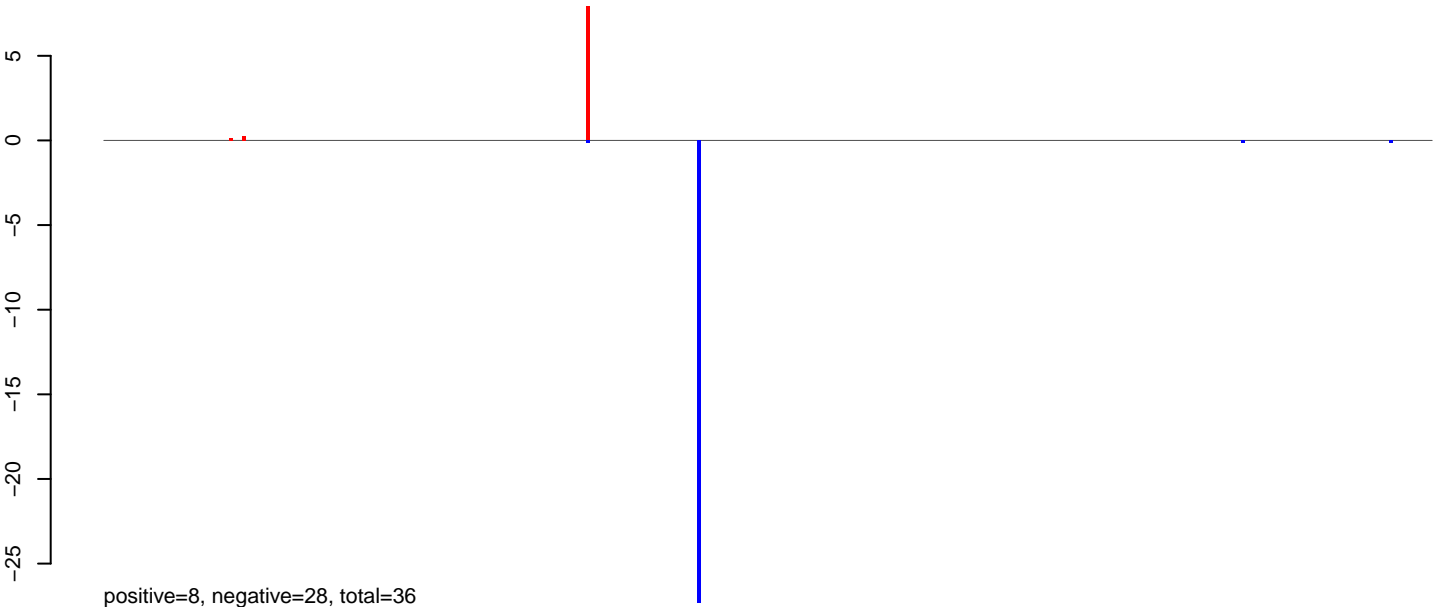
AeAlbo_C636_2dpi_mock3.24_35.rep



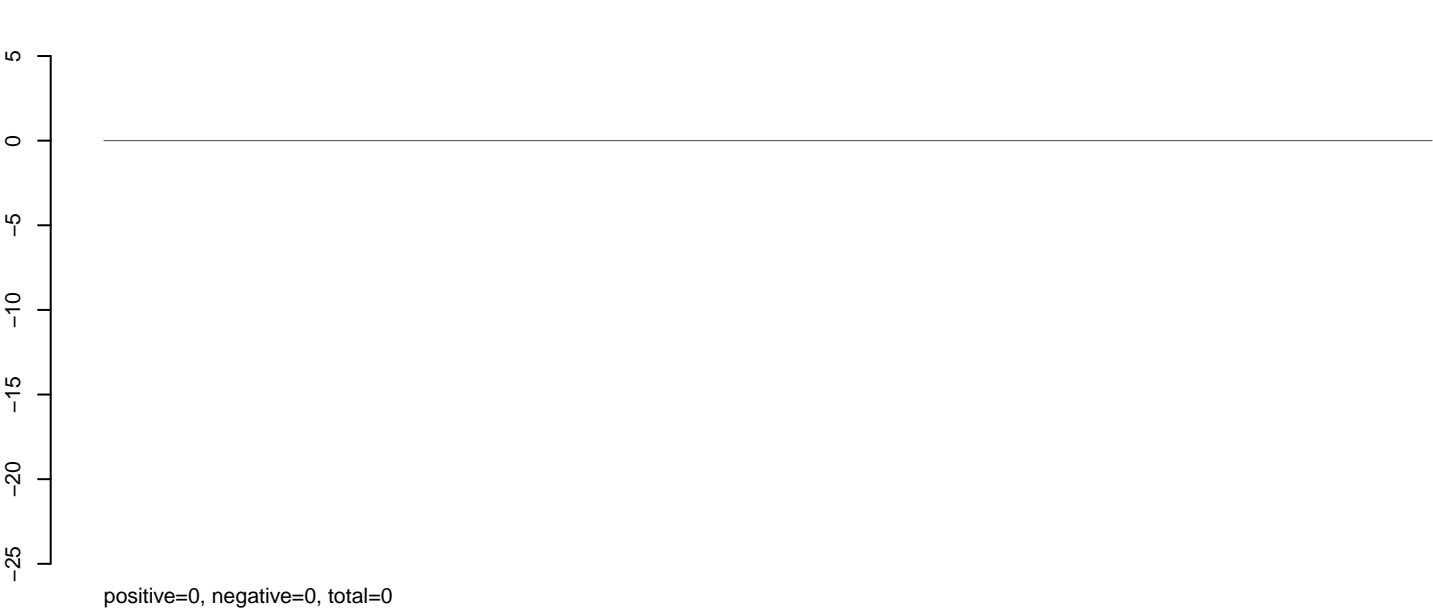
AeAlbo_C636_2dpi_mock3.rep



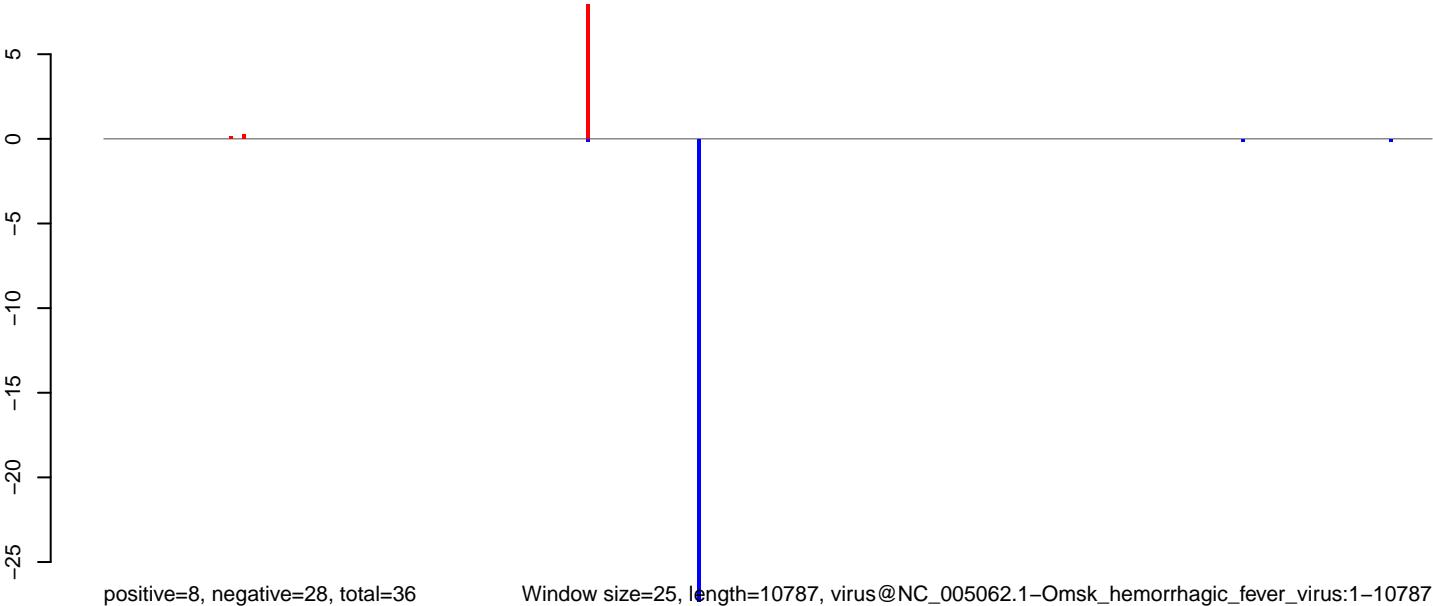
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



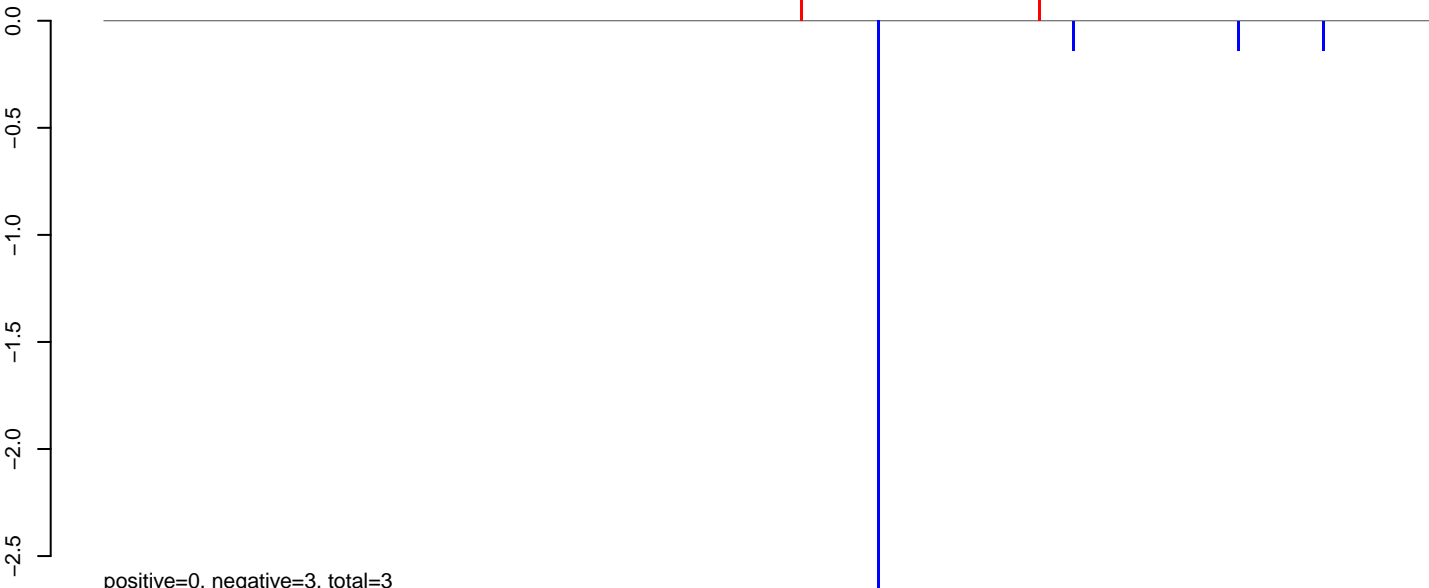
AeAlbo_C636_2dpi_mock3.rep



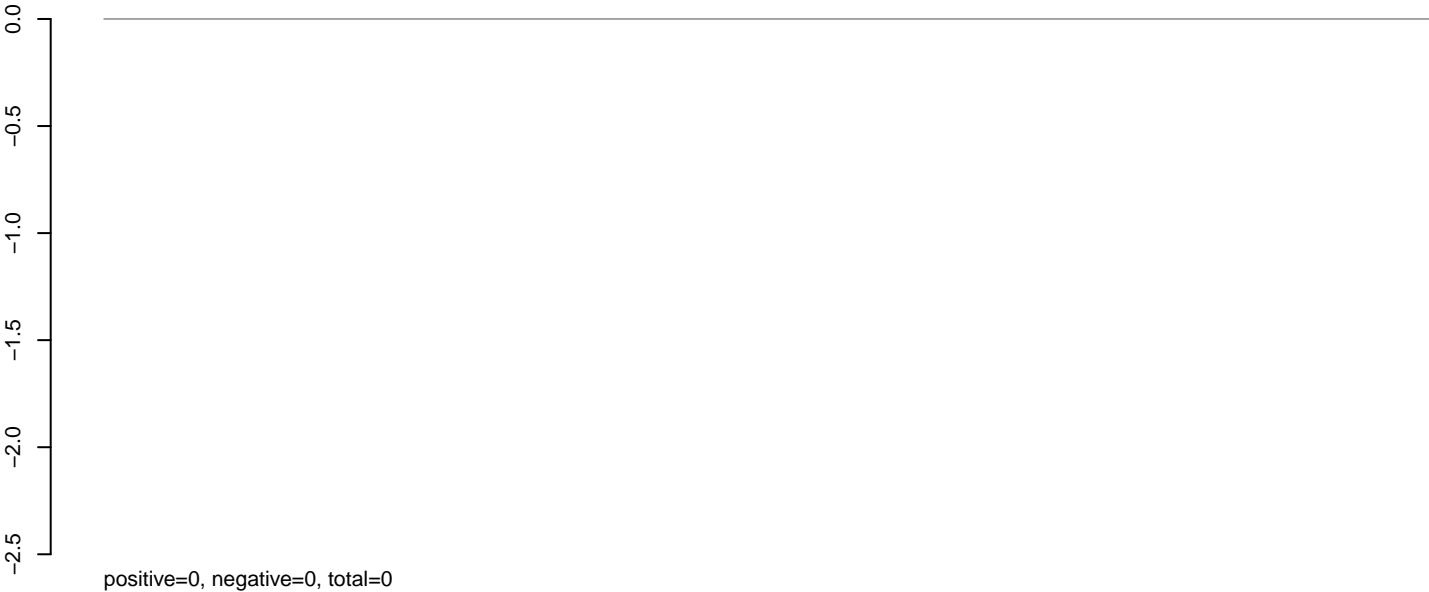
Window size=25, length=10787, virus@NC_005062.1-Omsk_hemorrhagic_fever_virus:1-10787

0 2000 4000 6000 8000 10000

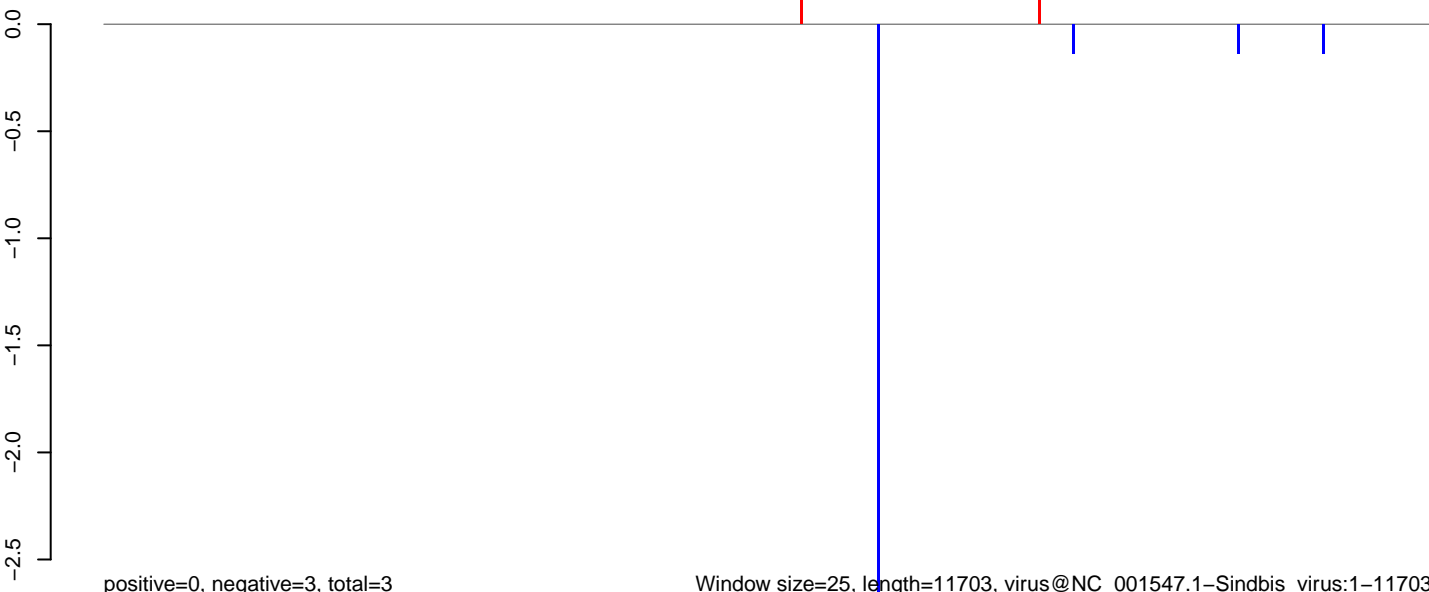
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

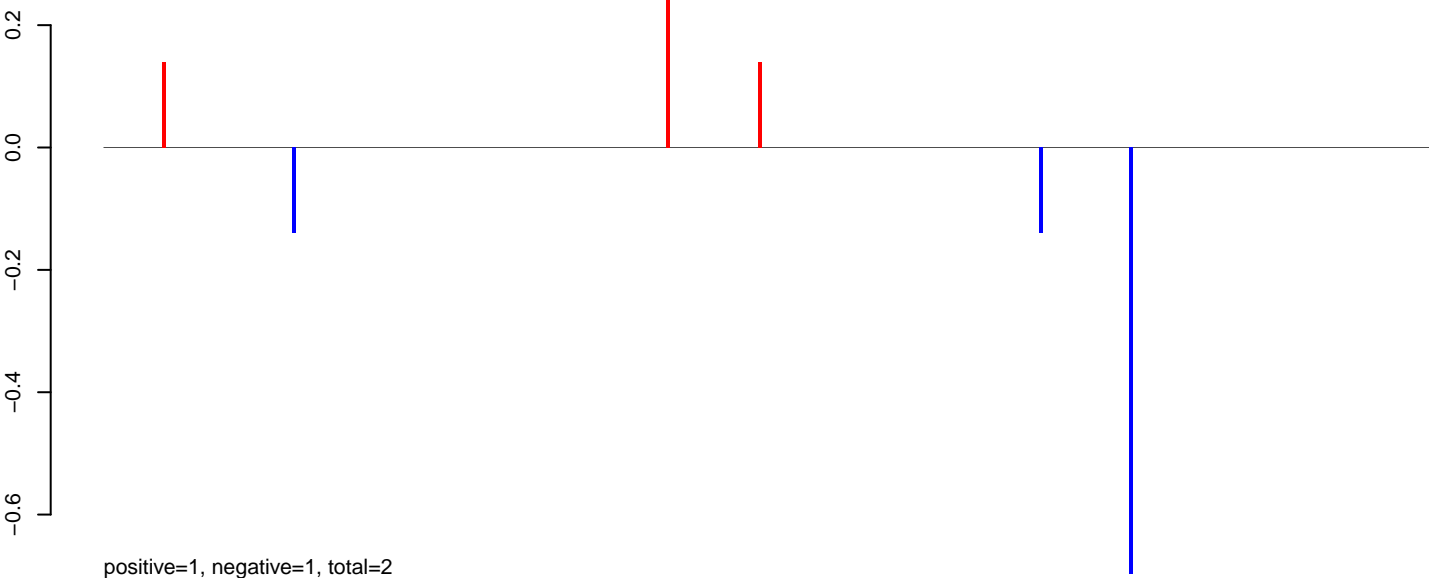


AeAlbo_C636_2dpi_mock3.rep

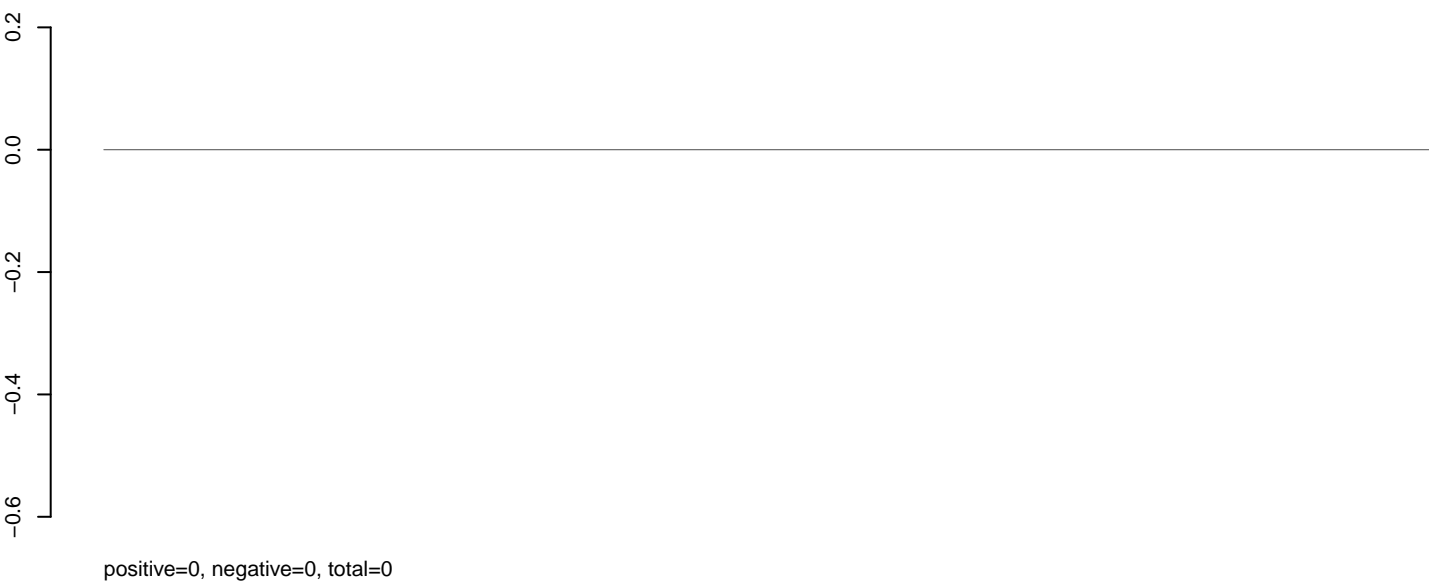


0 2000 4000 6000 8000 10000 12000

AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



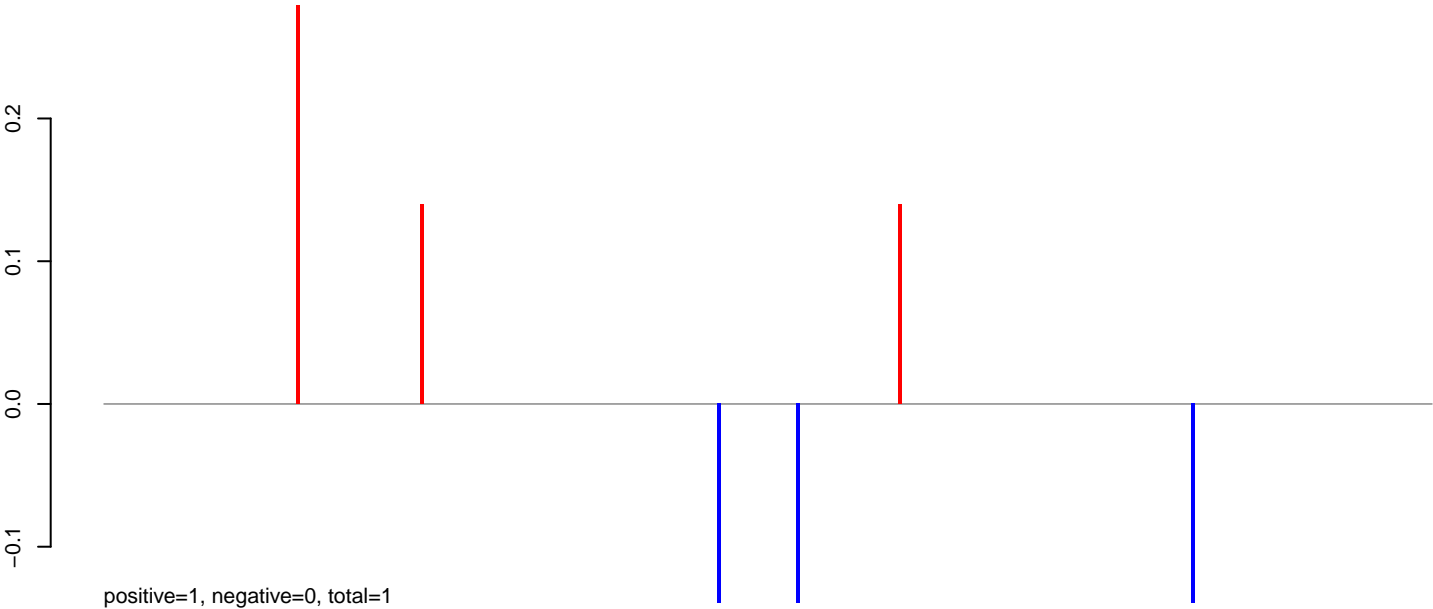
AeAlbo_C636_2dpi_mock3.rep



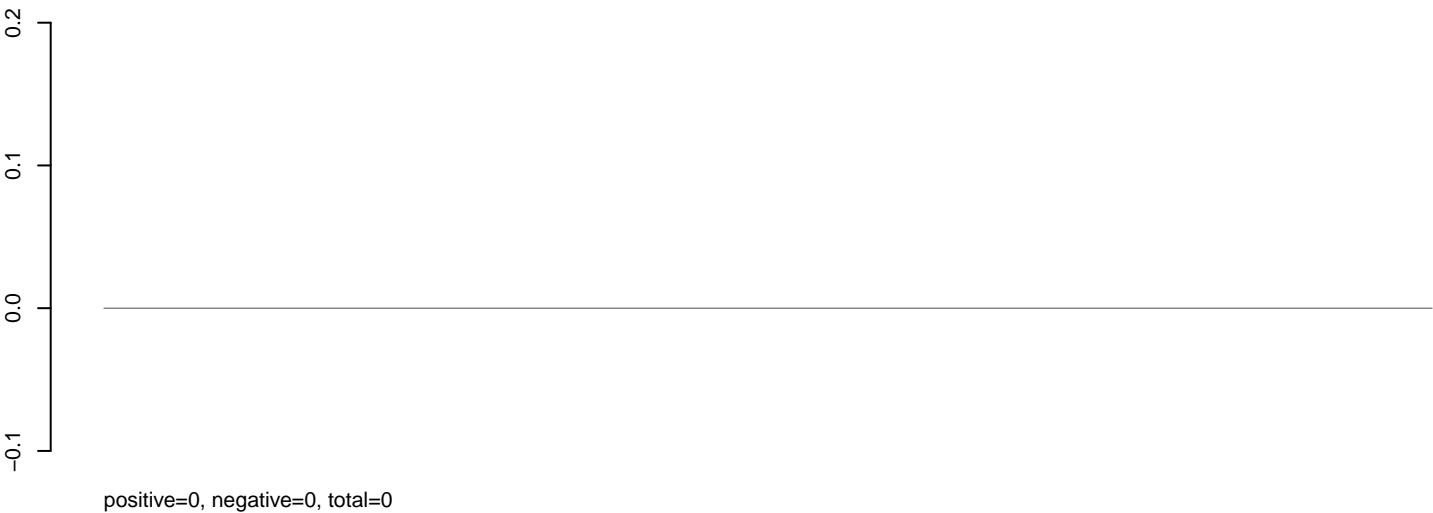
Window size=25, length=10735, virus@NC_001477.1-Dengue_virus_1:1-10735

0 2000 4000 6000 8000 10000

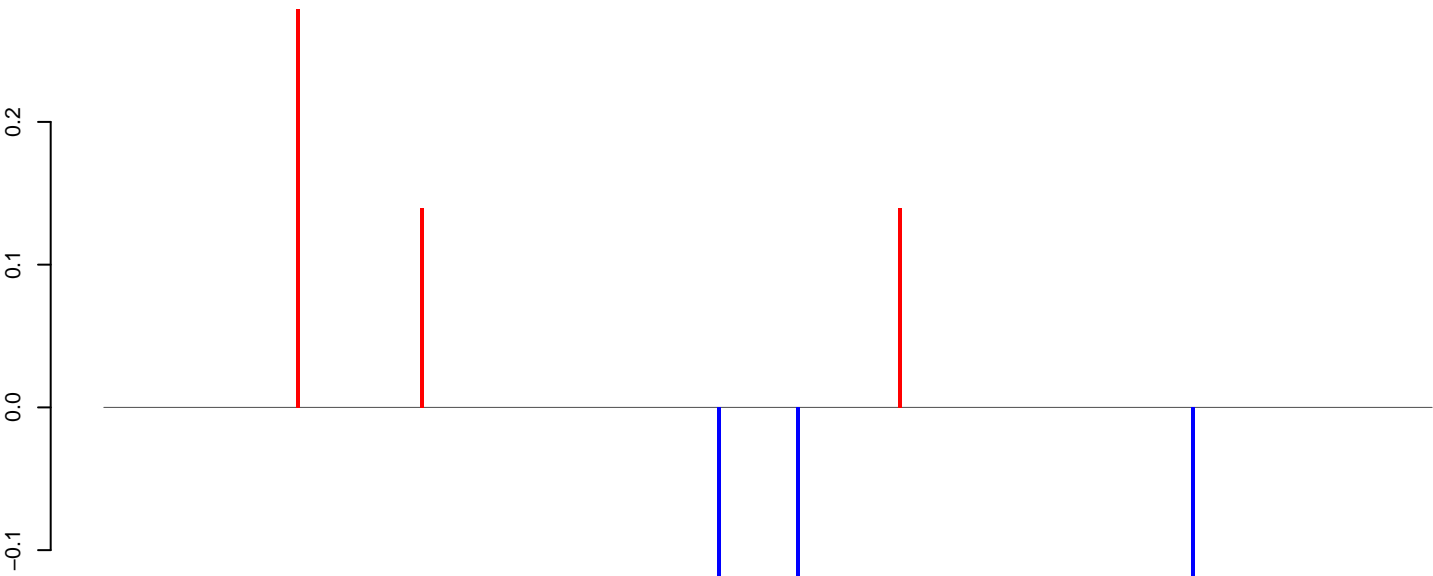
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



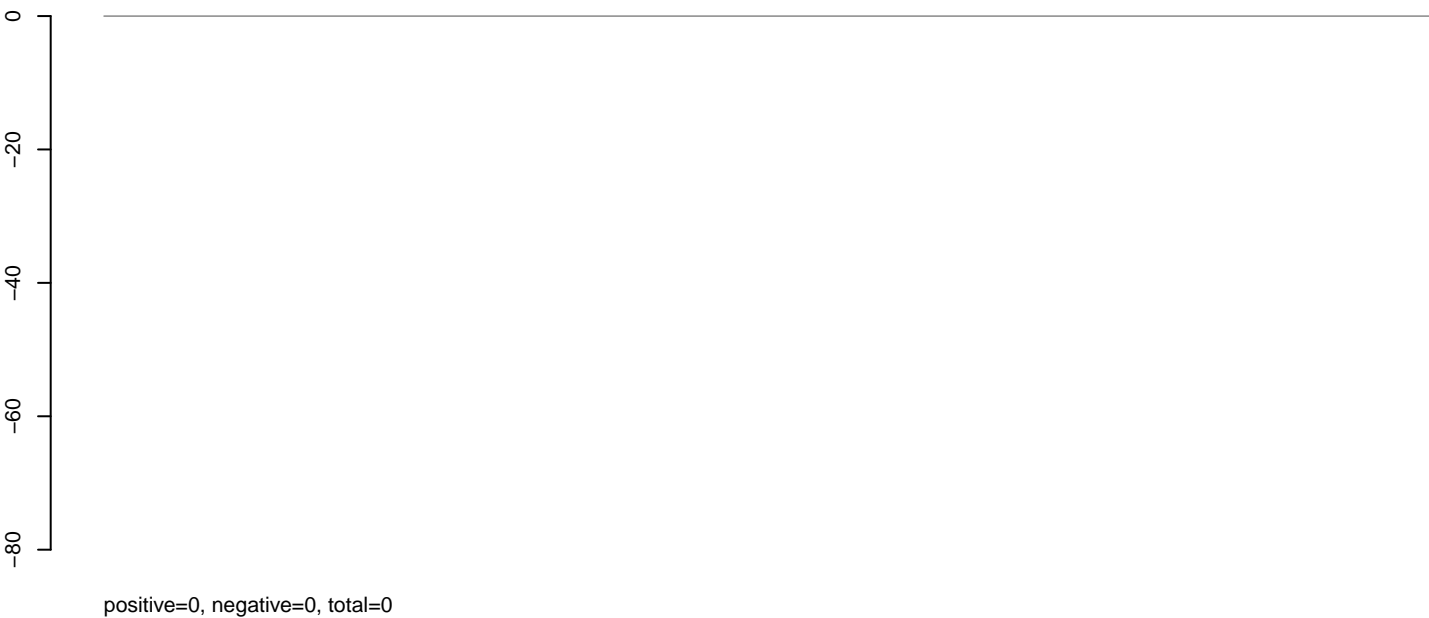
idow size=125, length=8819, virus=MF176325.1_Aedes_alboannulatus_Orthomyxo-like_virus_strain_mosWSAb70818_segment_1_2_3_4:1-8819

0 2000 4000 6000 8000

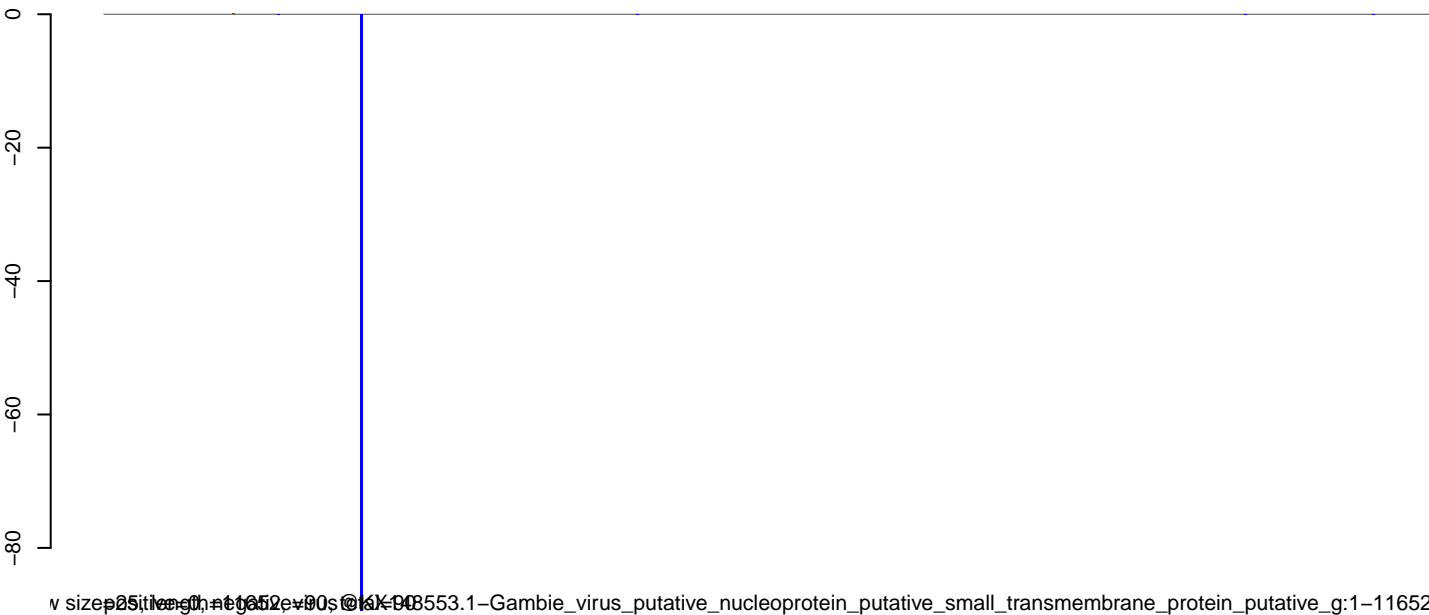
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



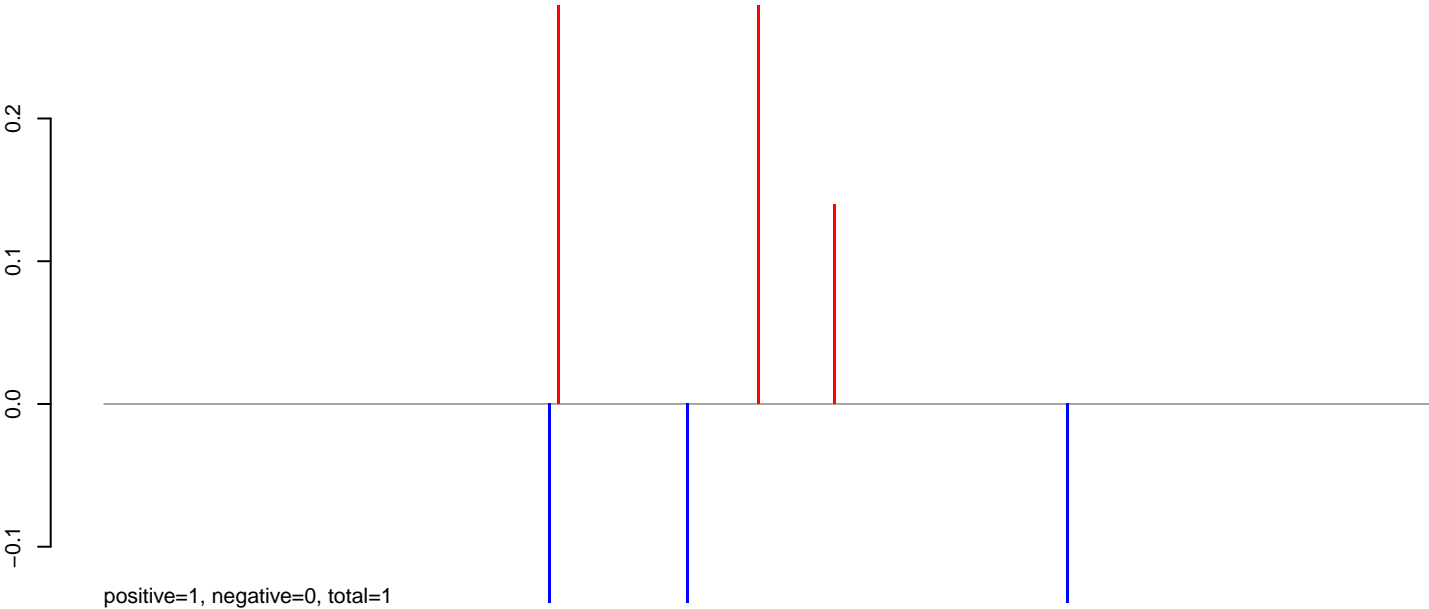
AeAlbo_C636_2dpi_mock3.rep



v size=25, length=11652, total=90, @1X=90
1-Gambie_virus_putative_nucleoprotein_putative_small_transmembrane_protein_putative_g:1-11652

0 2000 4000 6000 8000 10000 12000

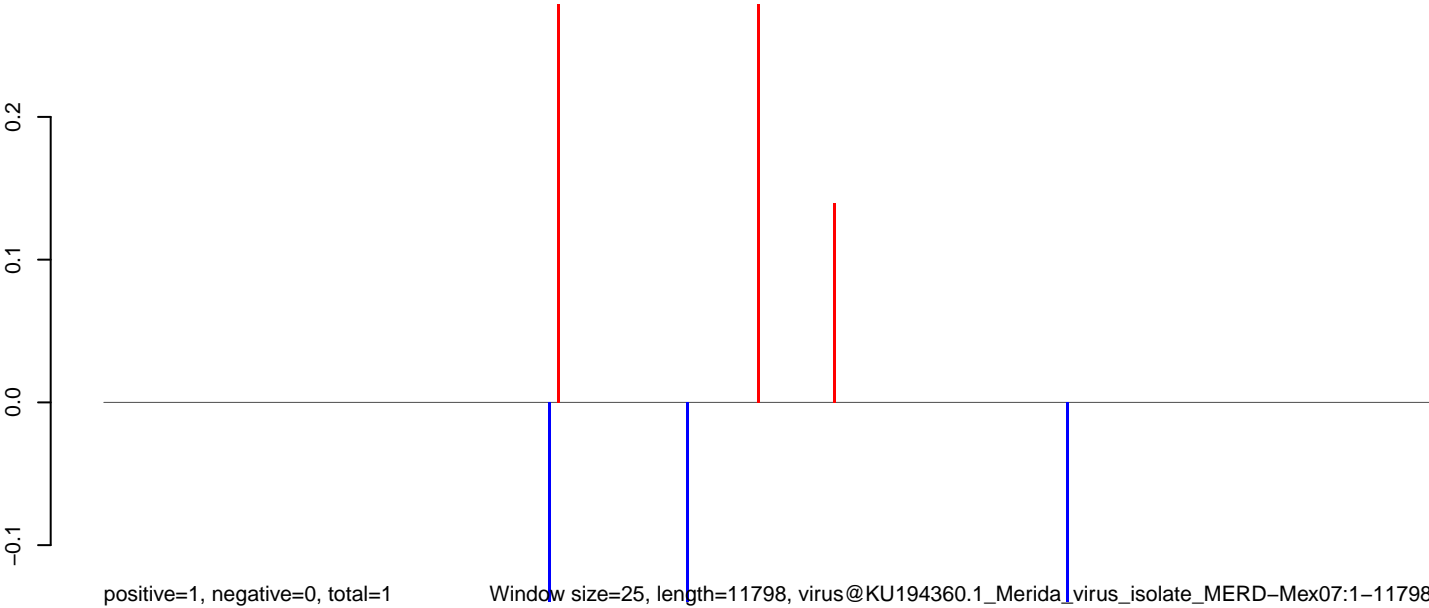
AeAlbo_C636_2dpi_mock3.18_23.rep



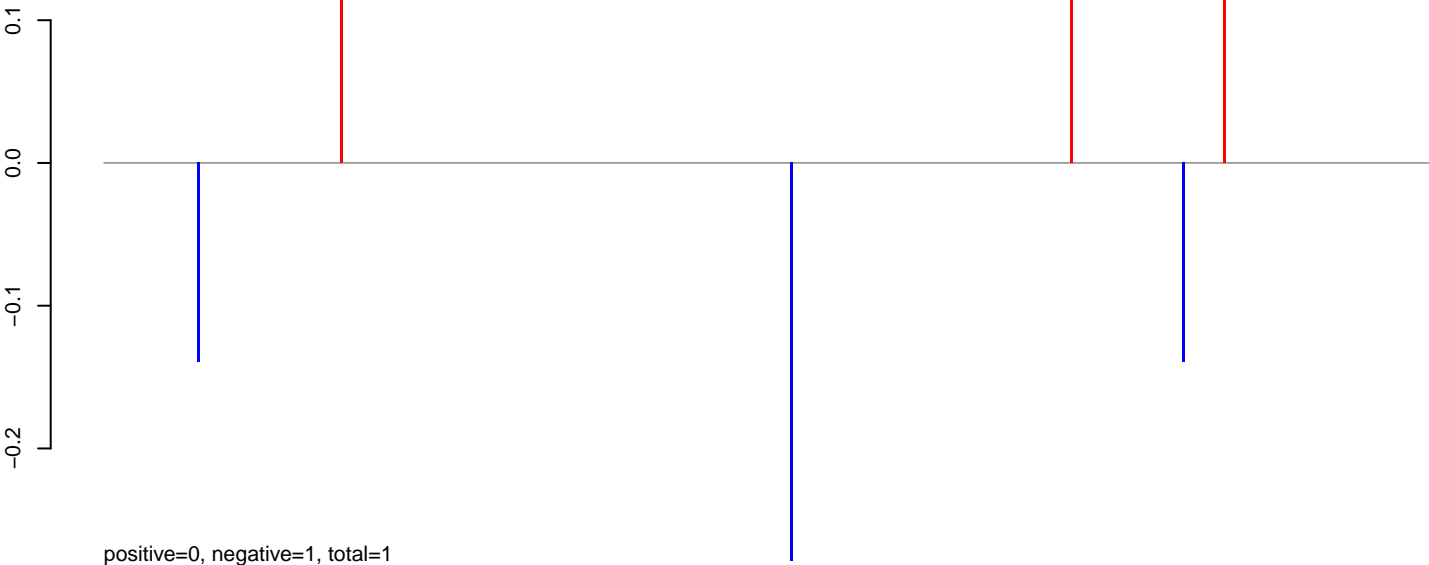
AeAlbo_C636_2dpi_mock3.24_35.rep



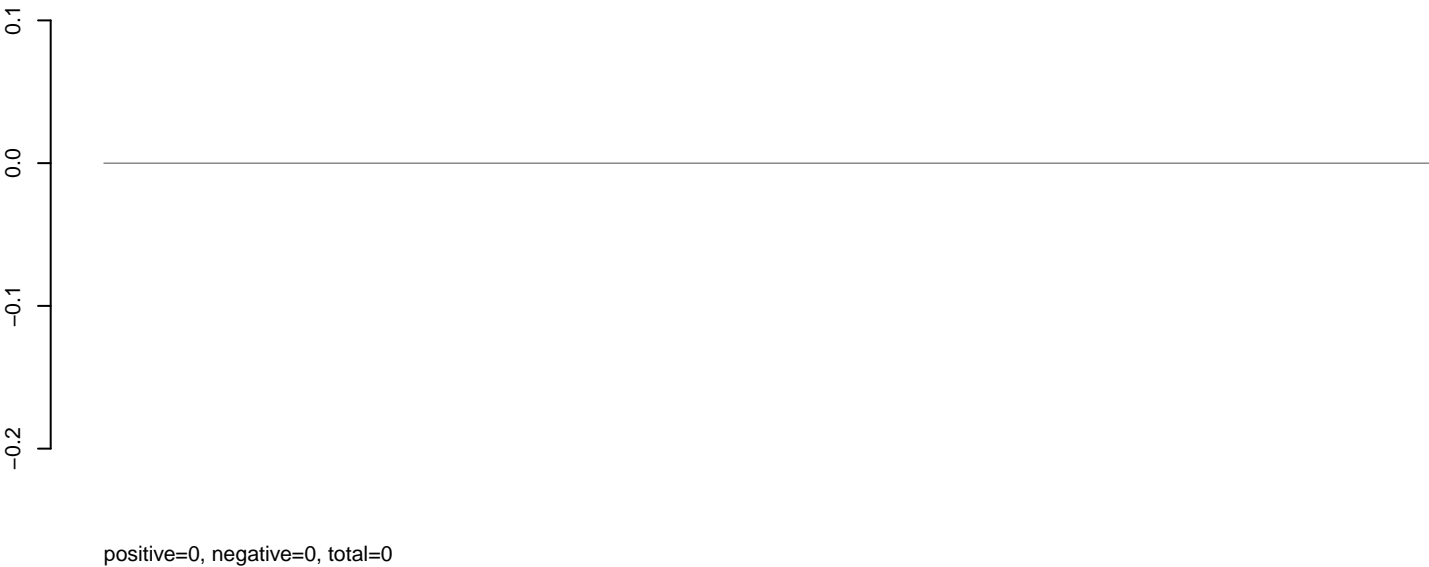
AeAlbo_C636_2dpi_mock3.rep



AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



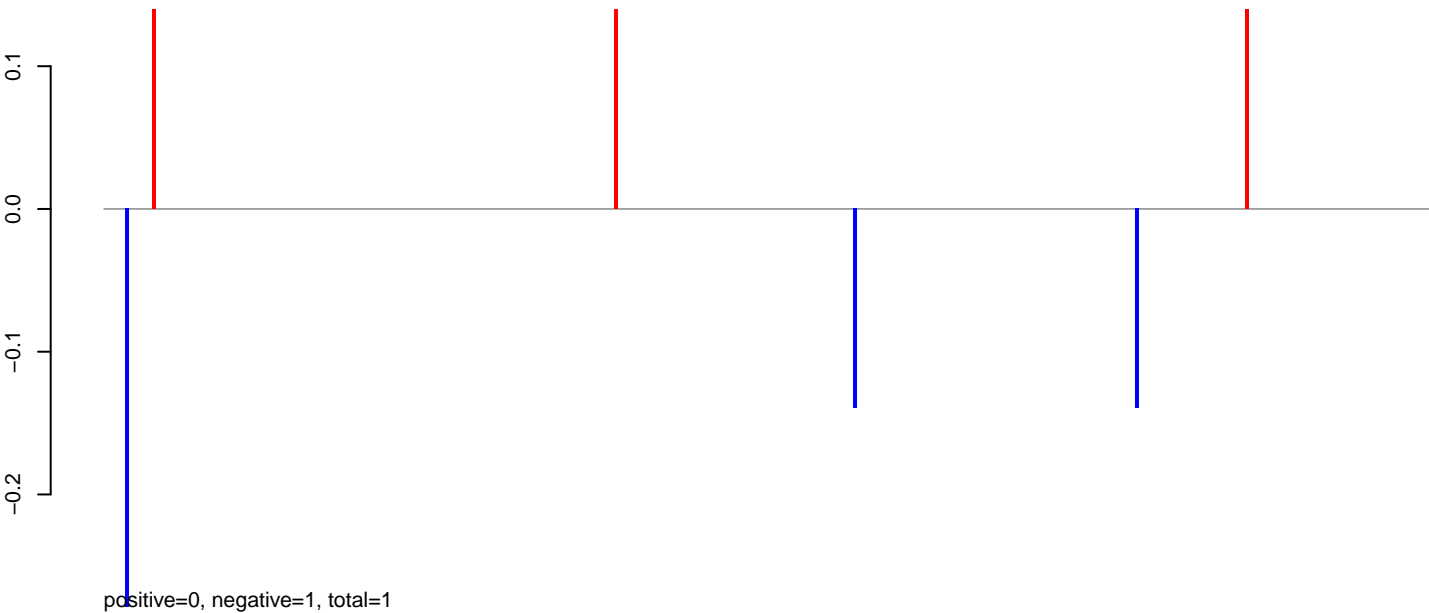
AeAlbo_C636_2dpi_mock3.rep



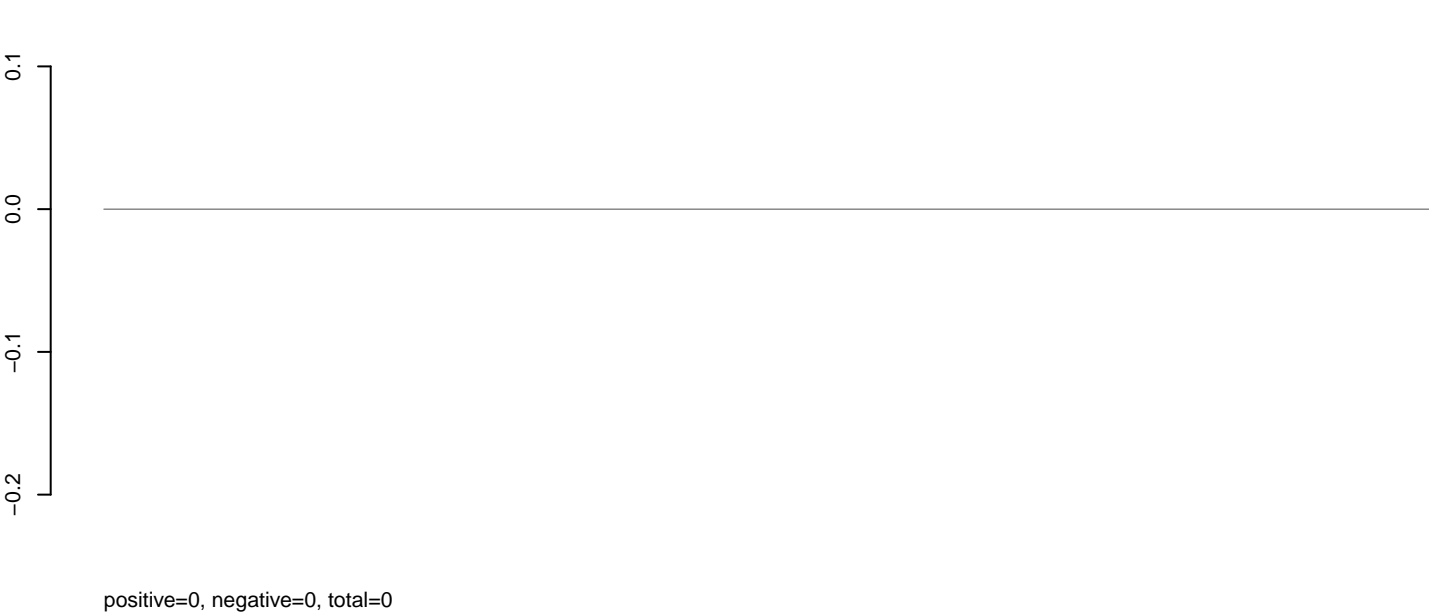
low size def of the 2004, virus @ 1X853181.1-Schmallenberg_virus_isolate_Germany_segment_S_plus_segment_M_plus_segment_L:1-12094

0 2000 4000 6000 8000 10000 12000

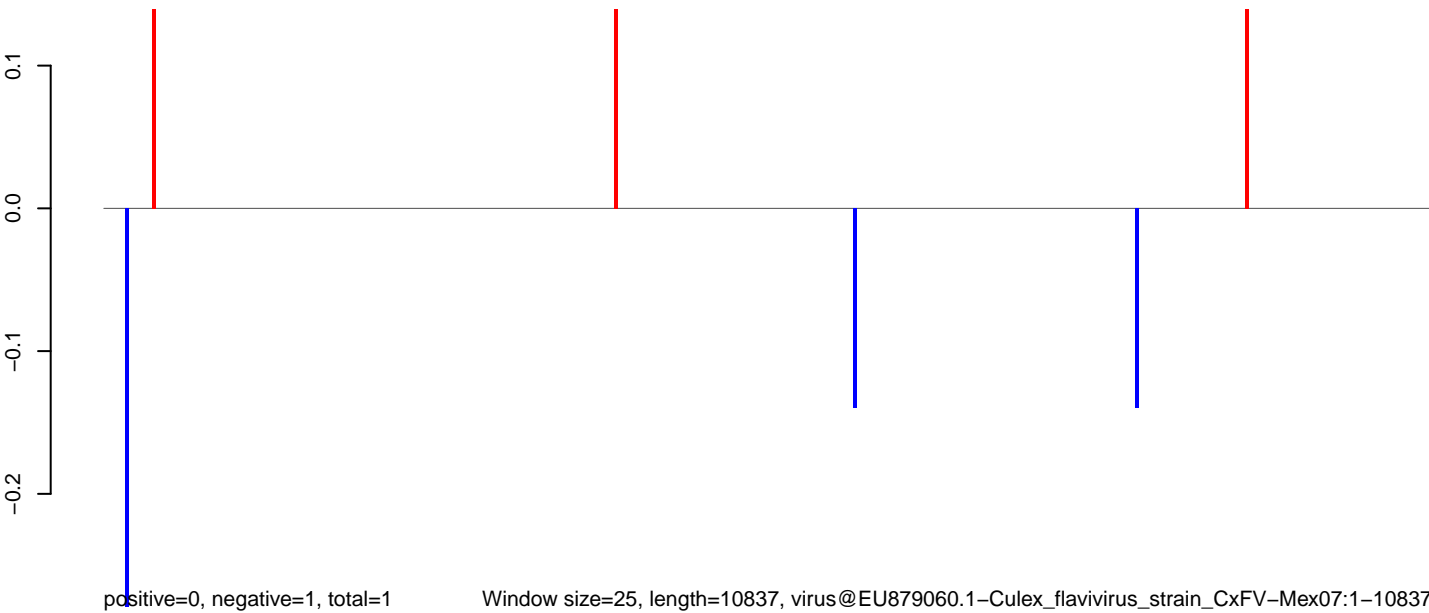
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



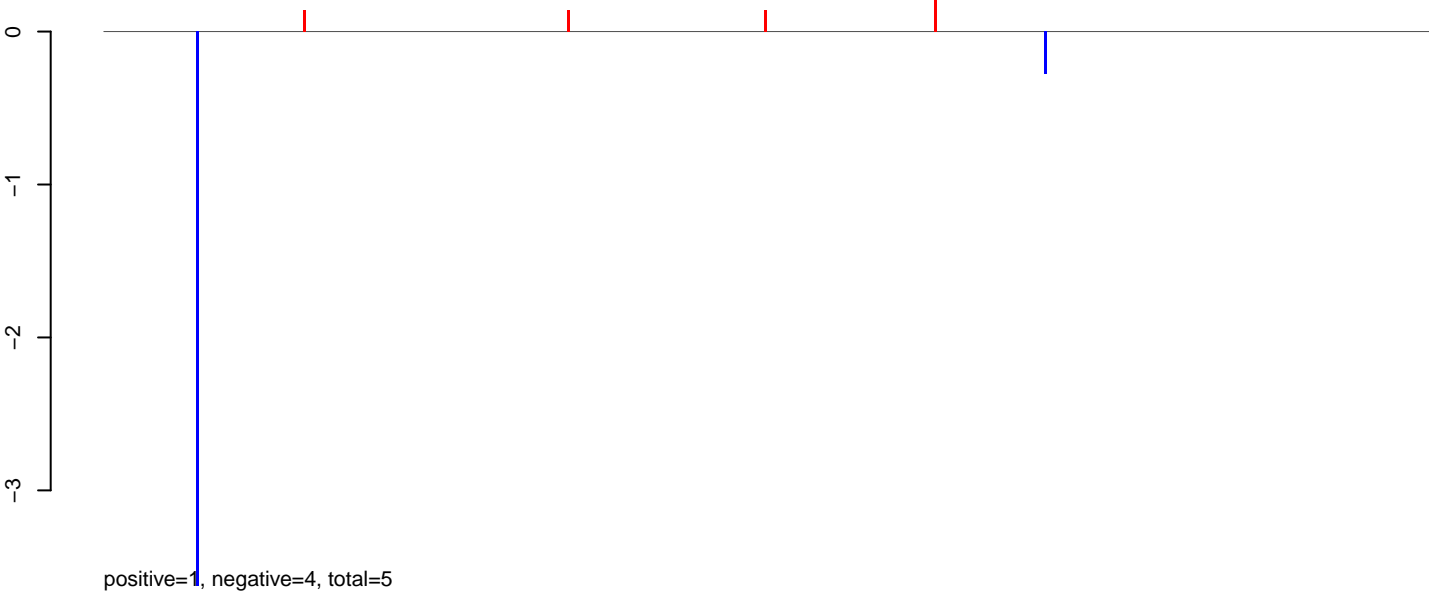
AeAlbo_C636_2dpi_mock3.rep



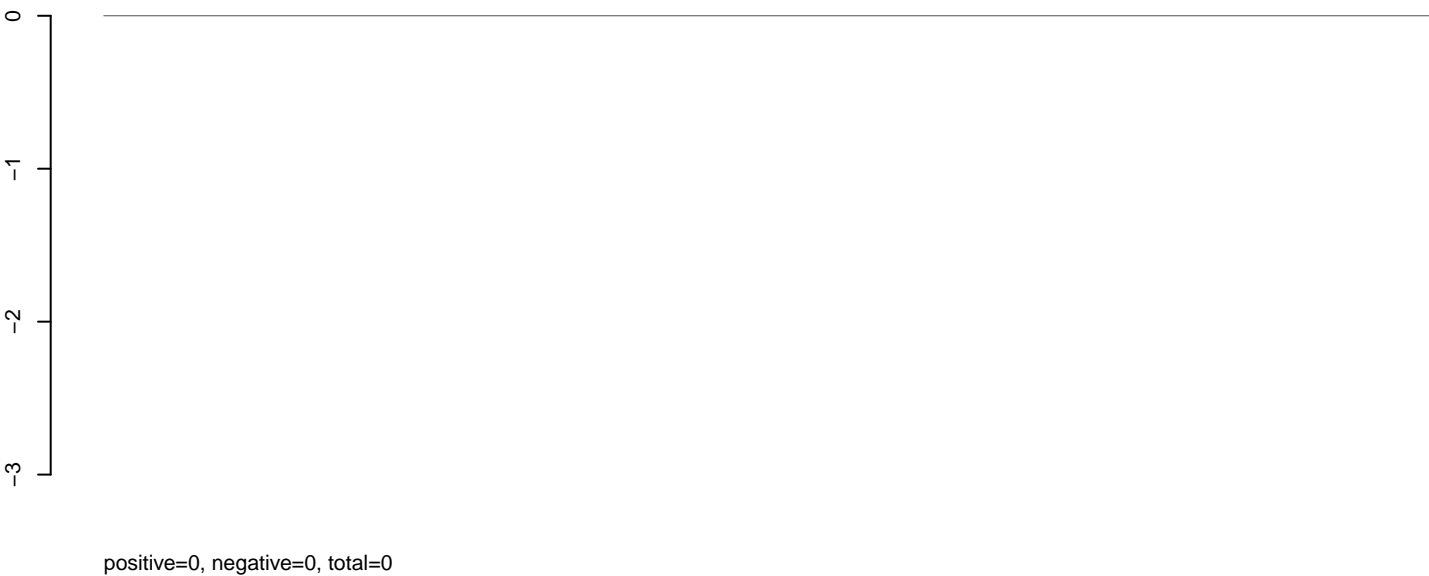
Window size=25, length=10837, virus@EU879060.1-Culex_flavivirus_strain_CxFV-Mex07:1-10837

0 2000 4000 6000 8000 10000

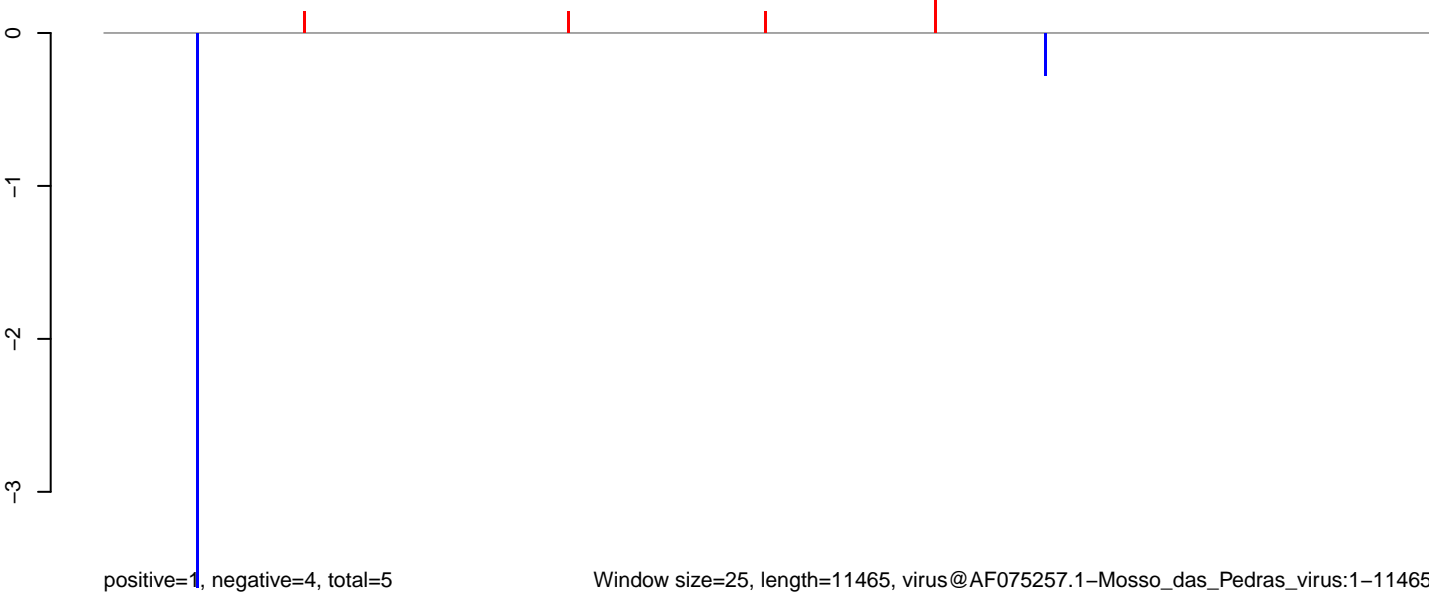
AeAlbo_C636_2dpi_mock3.18_23.rep



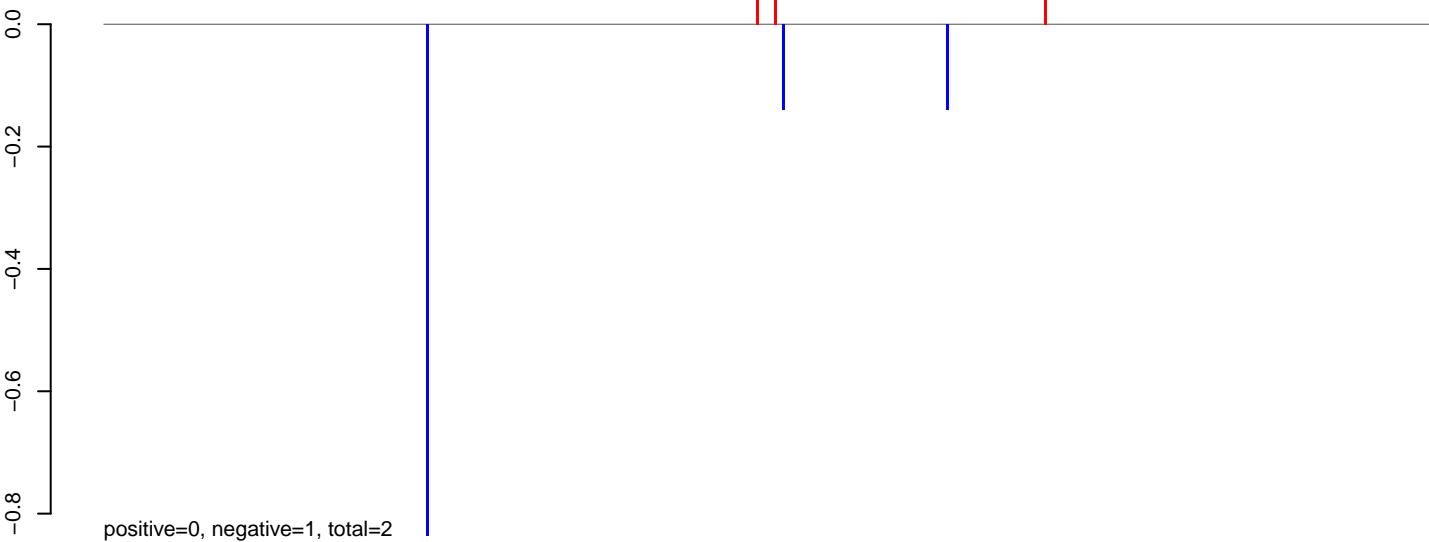
AeAlbo_C636_2dpi_mock3.24_35.rep



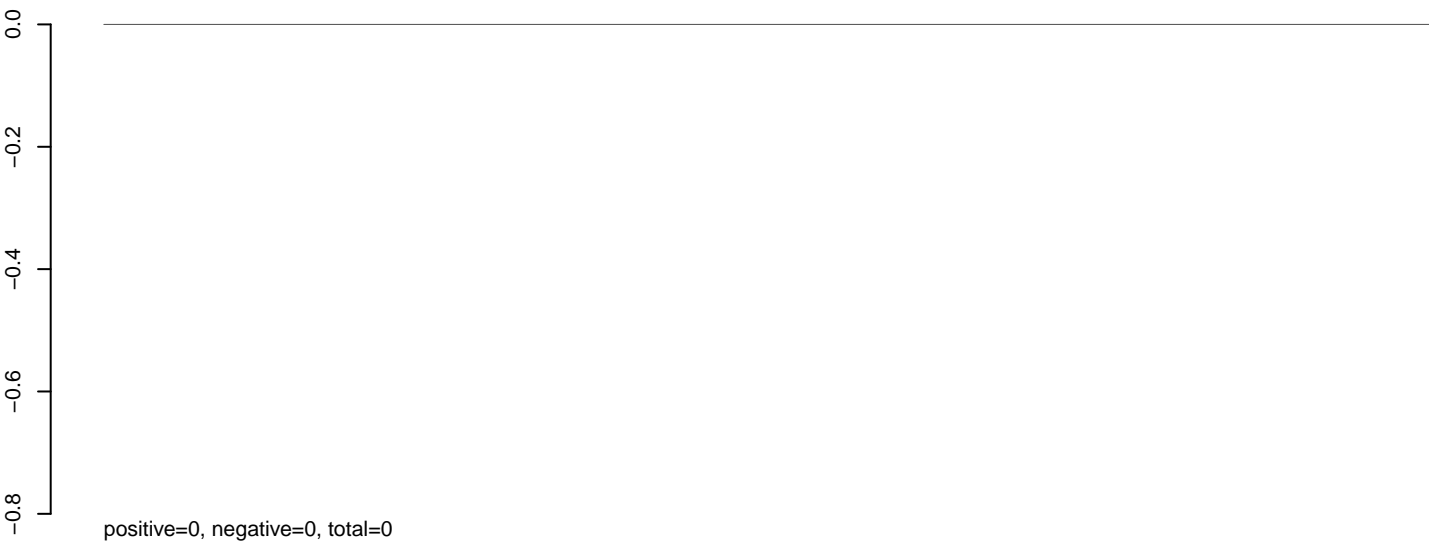
AeAlbo_C636_2dpi_mock3.rep



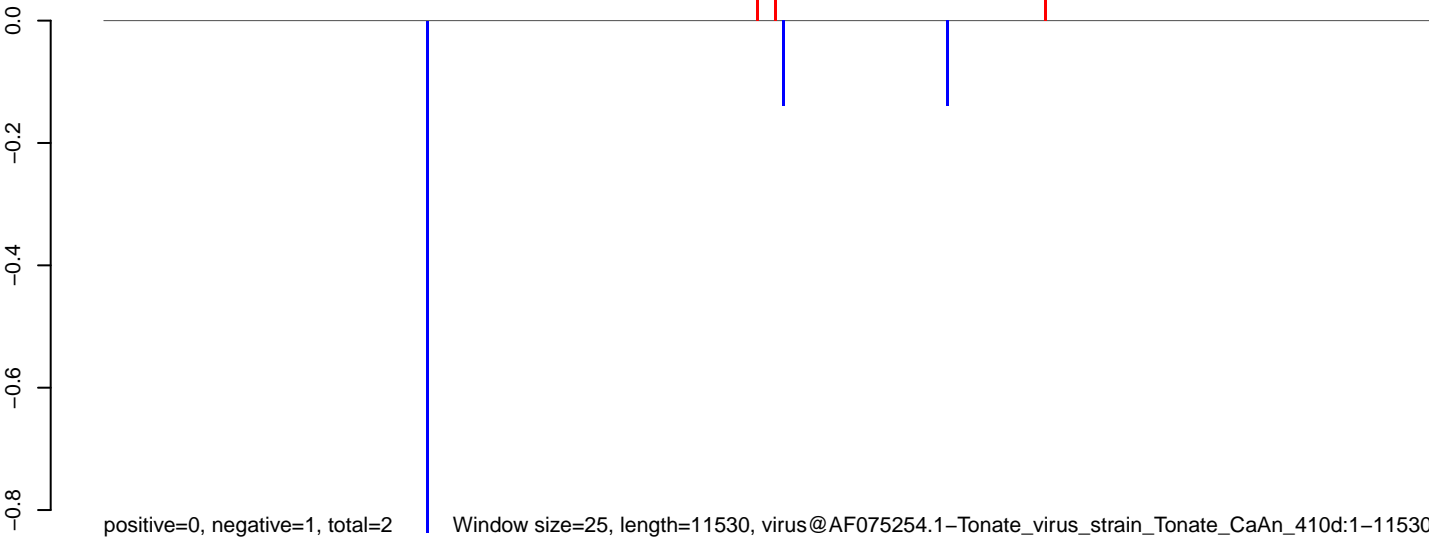
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

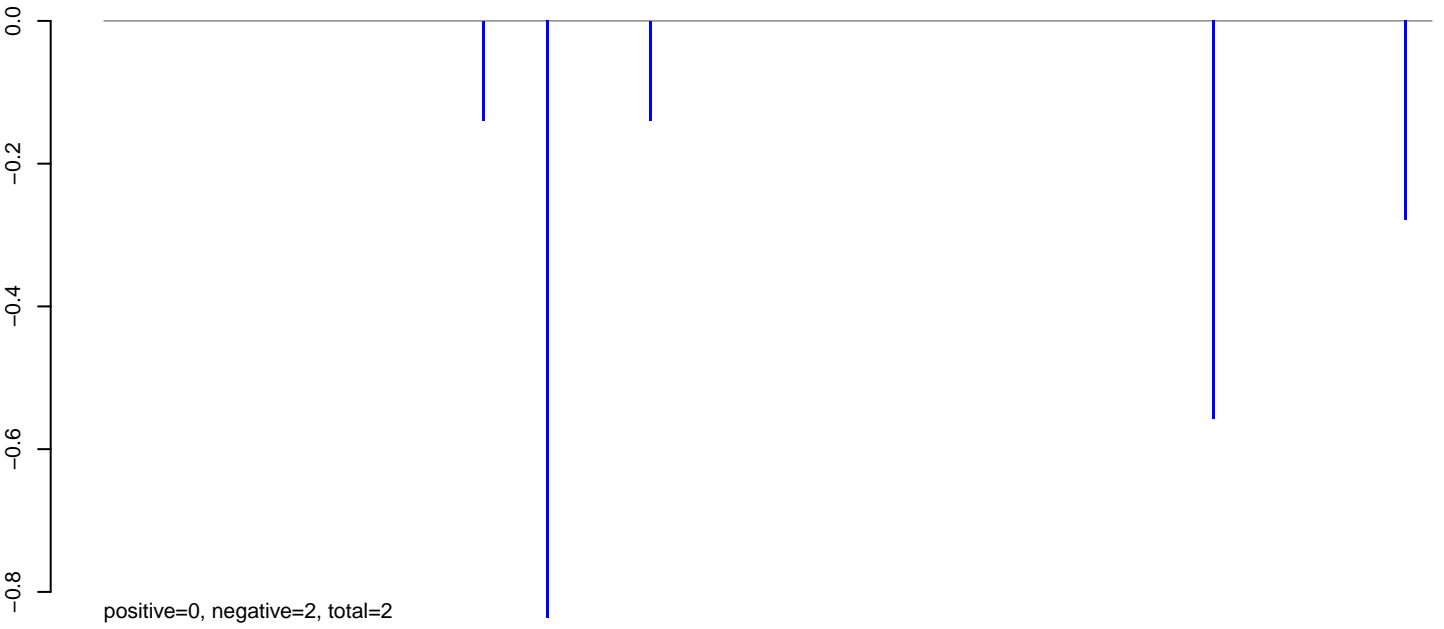


AeAlbo_C636_2dpi_mock3.rep

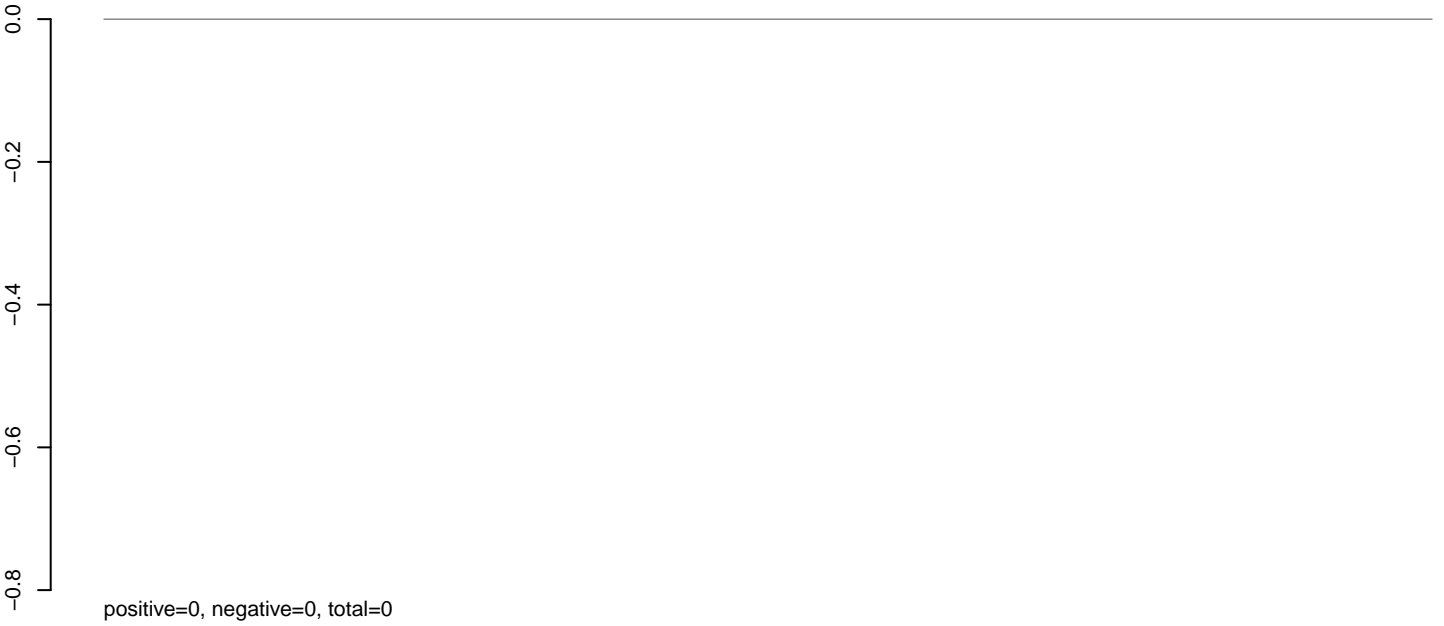


0 2000 4000 6000 8000 10000 12000

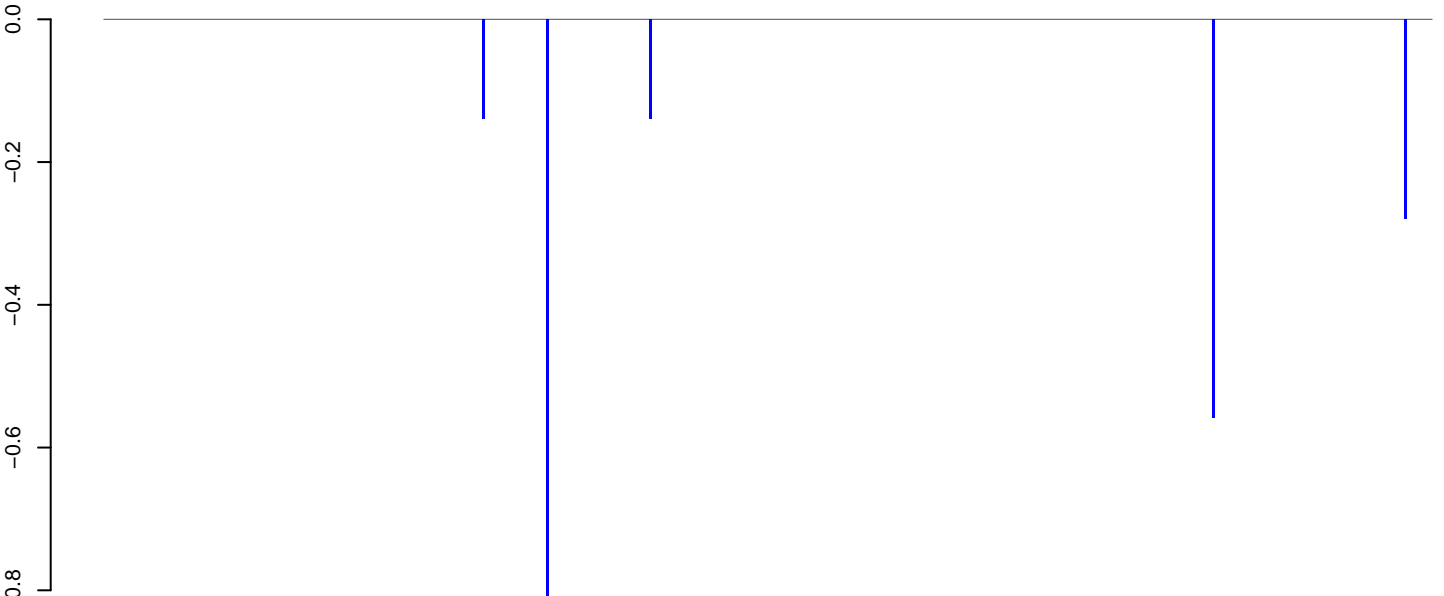
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



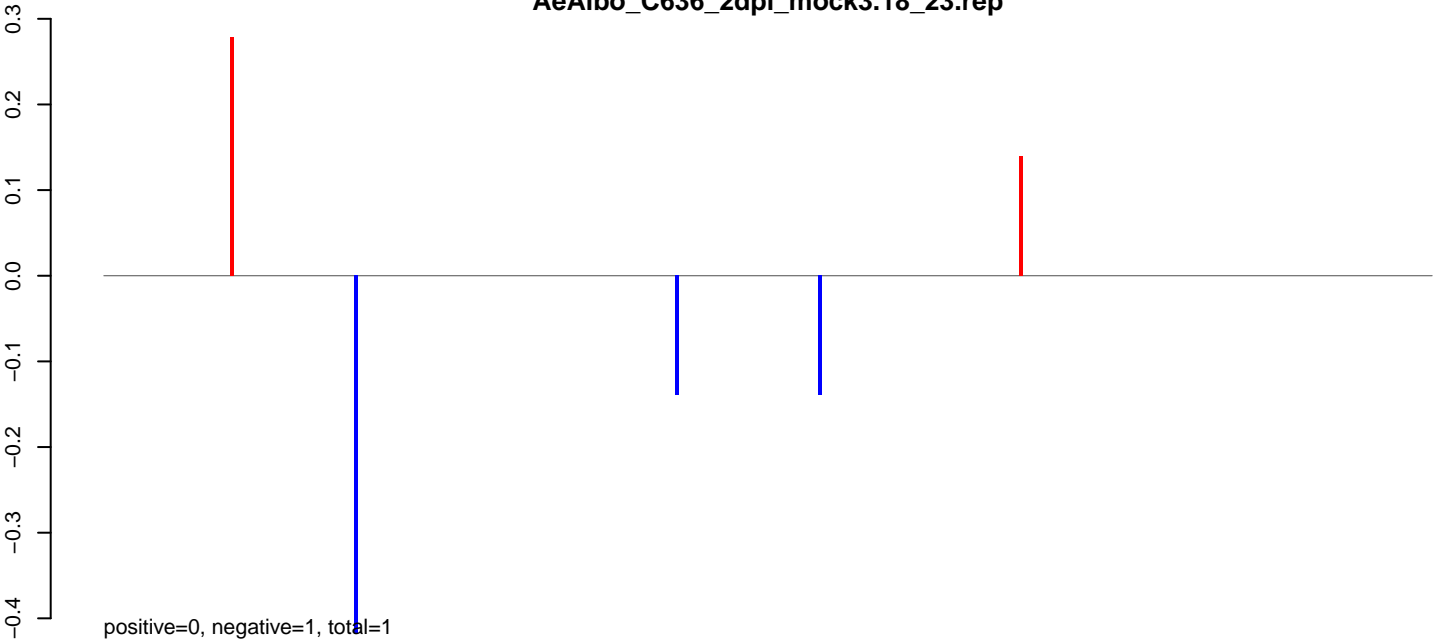
AeAlbo_C636_2dpi_mock3.rep



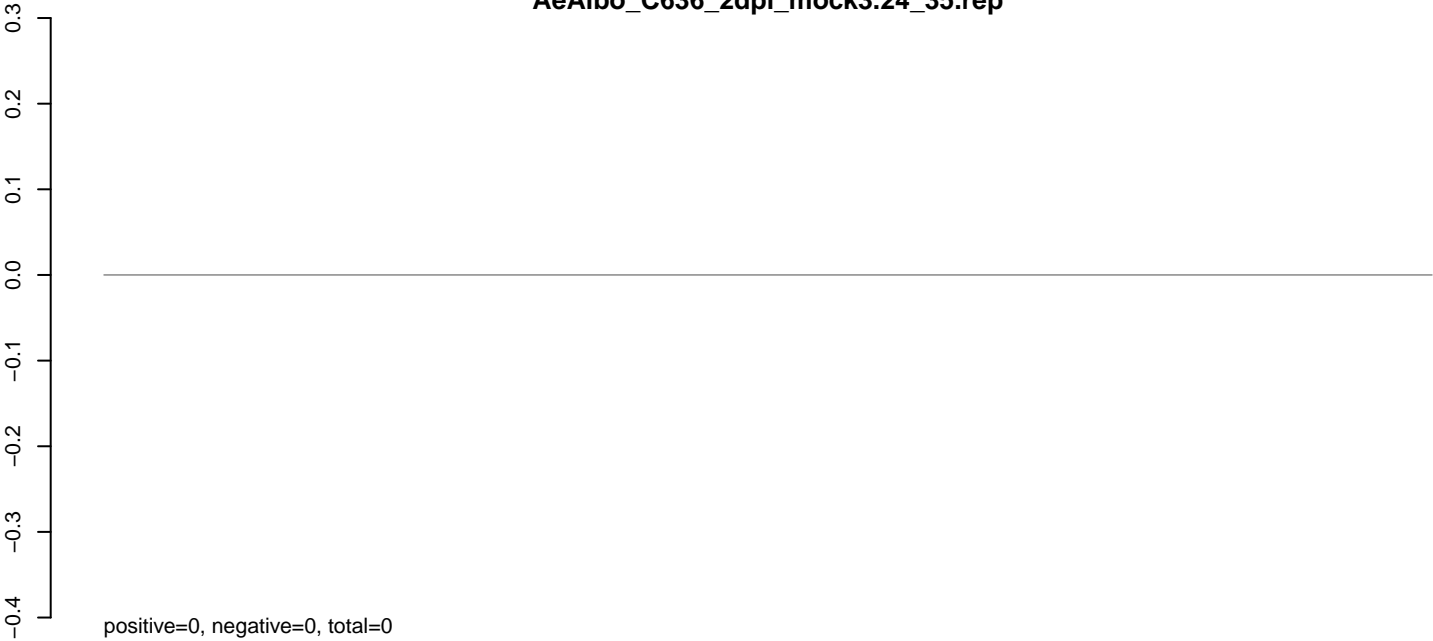
w size=51, length=12957, use@NC_034448.1-Mosqueiro_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopro:1-12957

0 2000 4000 6000 8000 10000 12000

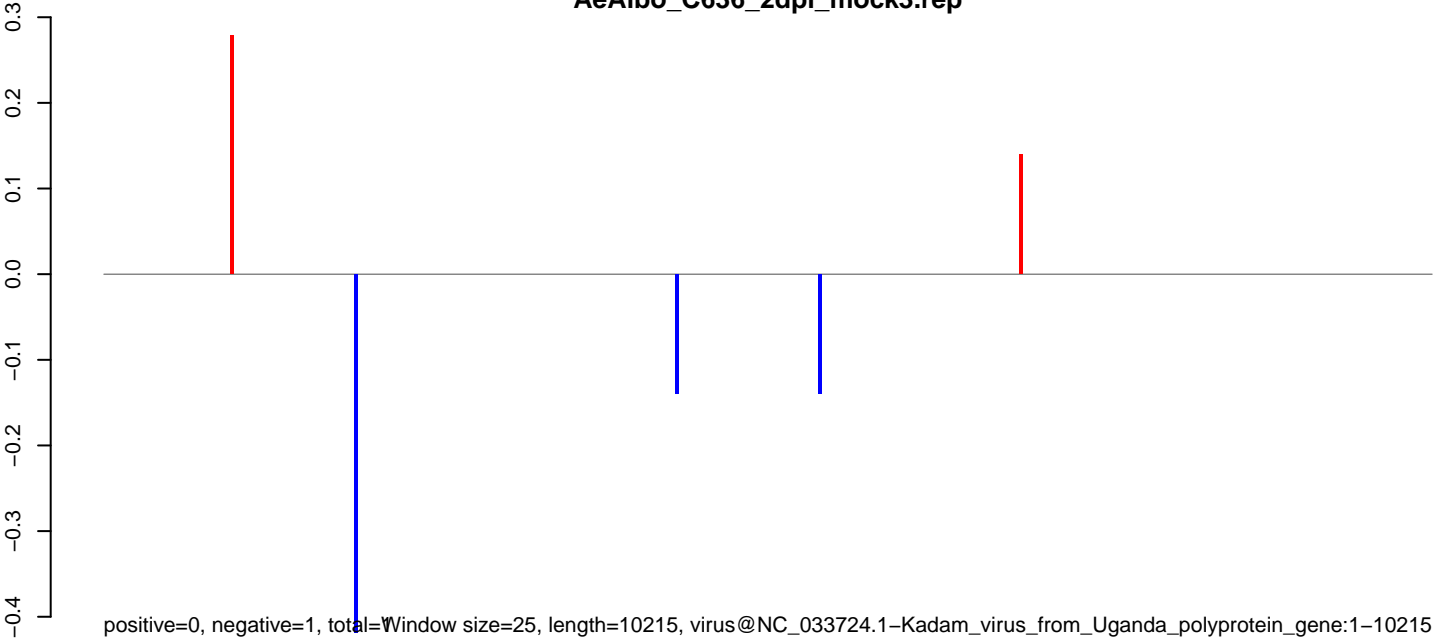
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



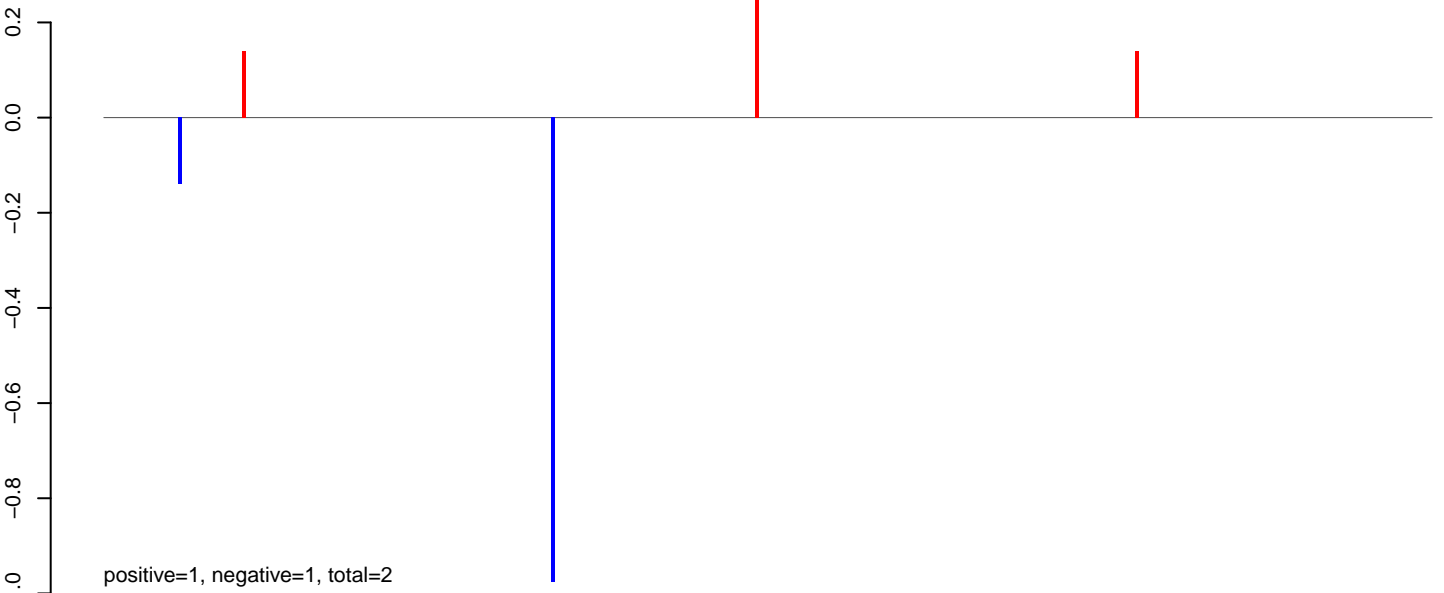
AeAlbo_C636_2dpi_mock3.rep



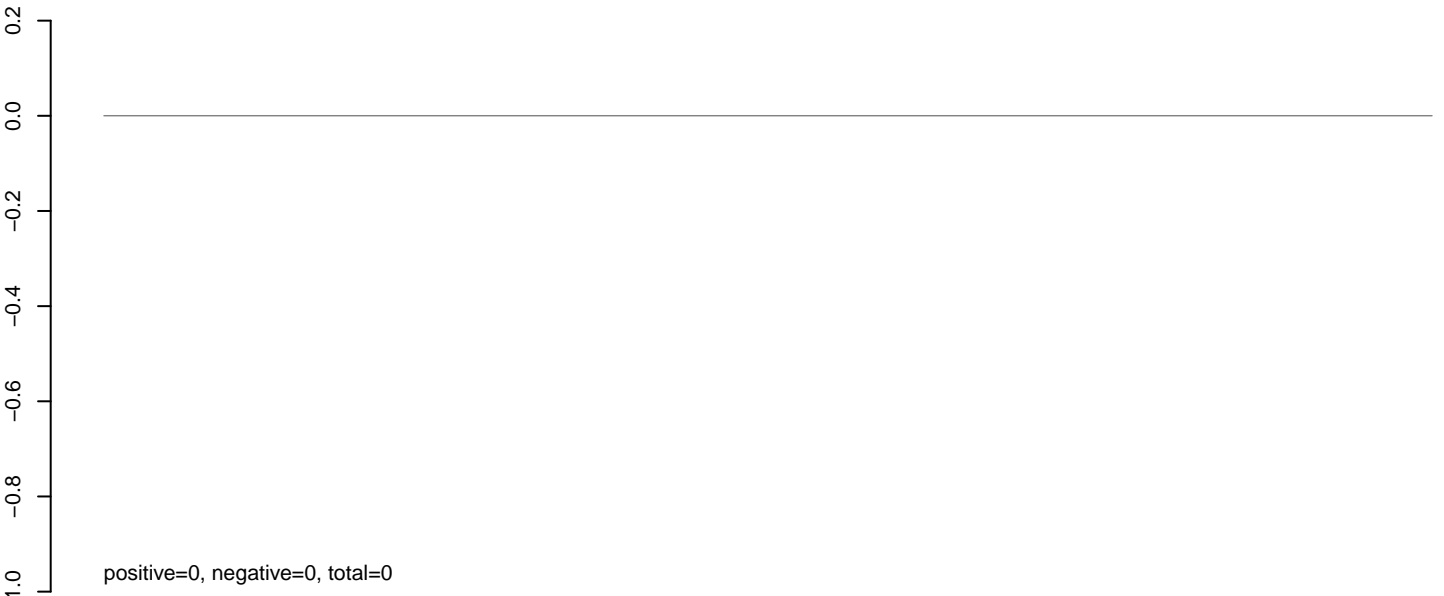
Window size=25, length=10215, virus@NC_033724.1-Kadam_virus_from_Uganda_polyprotein_gene:1-10215

0 2000 4000 6000 8000 10000

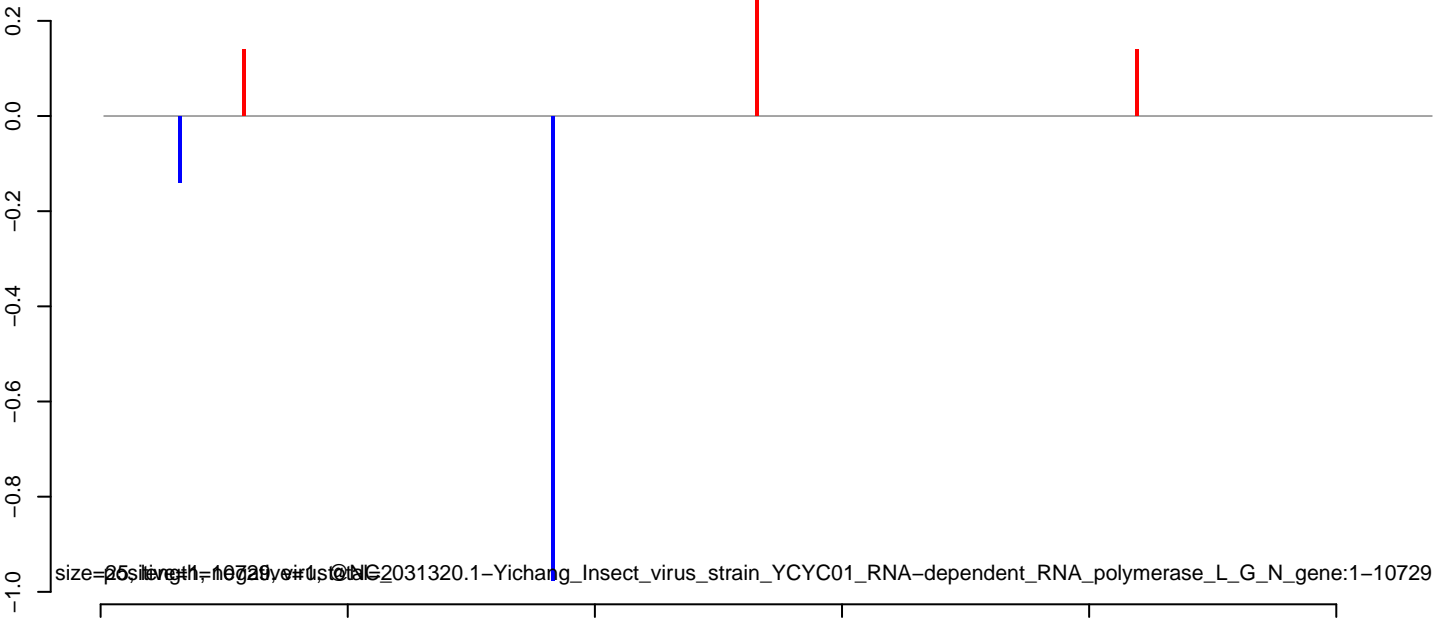
AeAlbo_C636_2dpi_mock3.18_23.rep

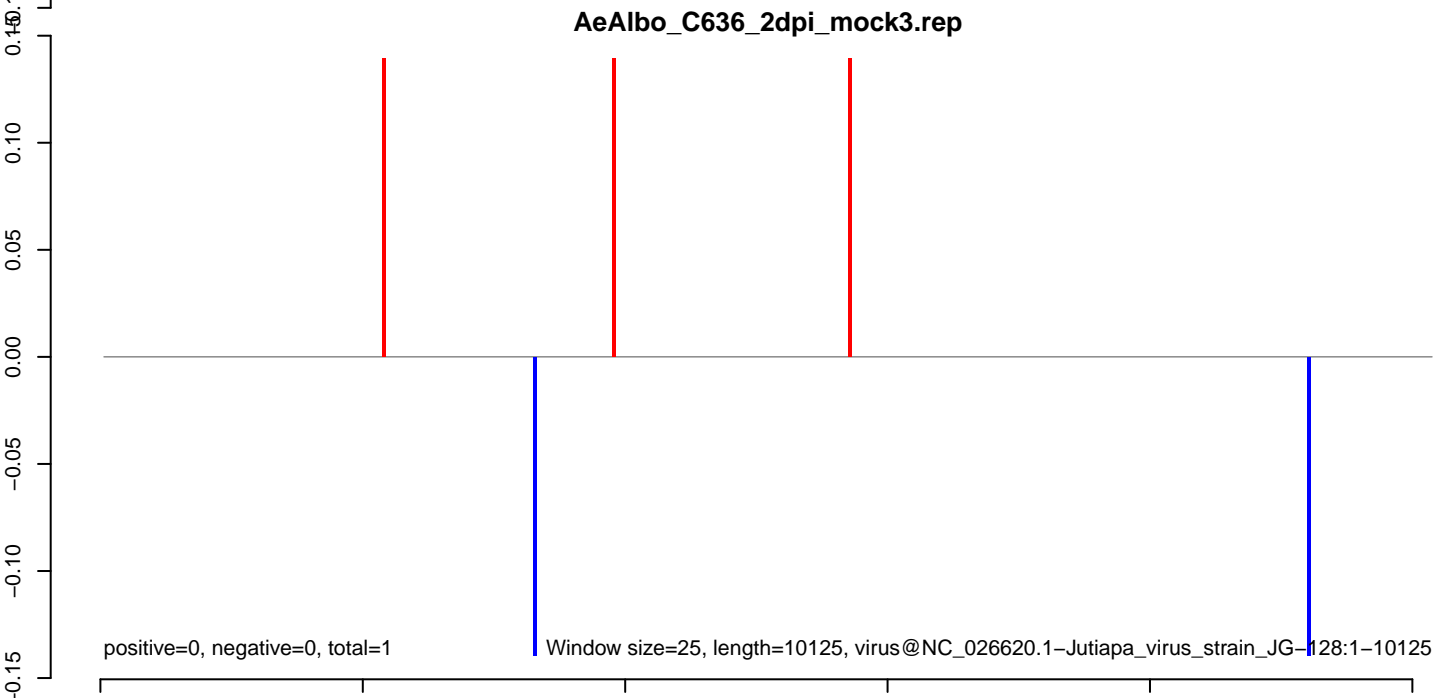
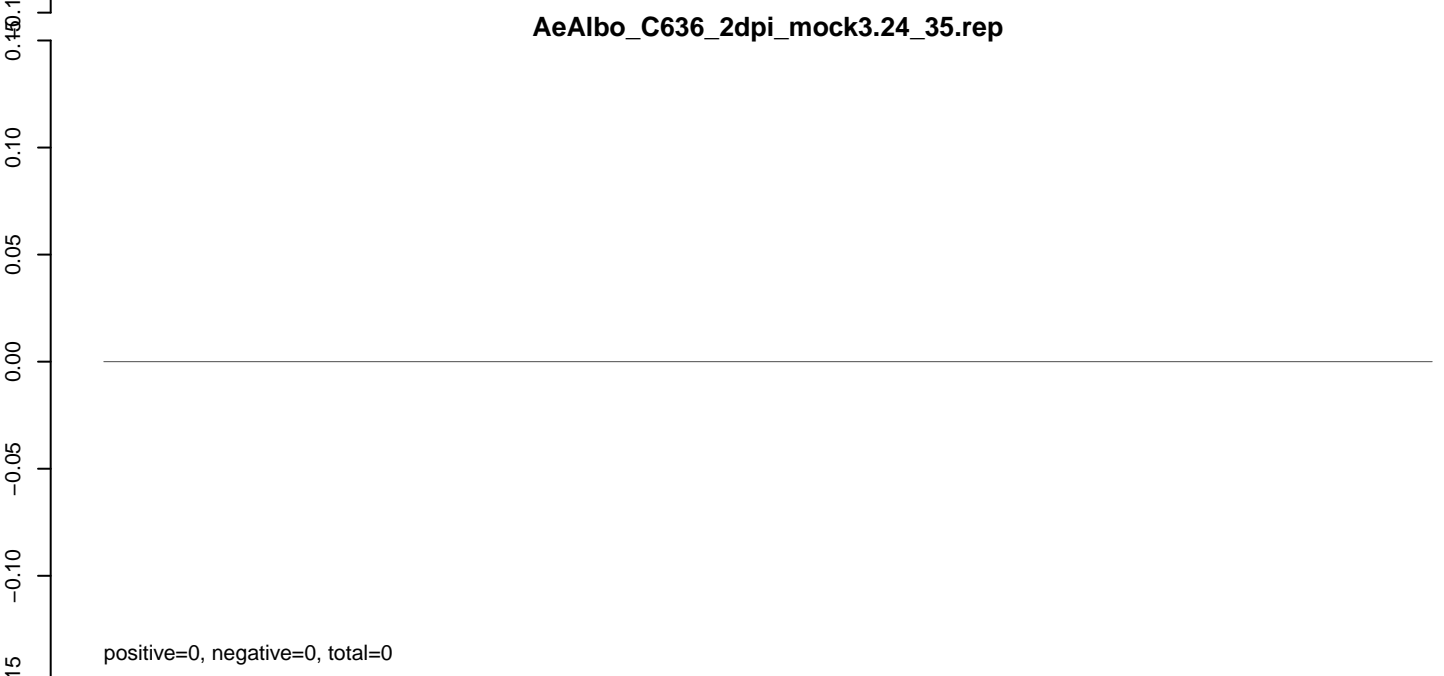
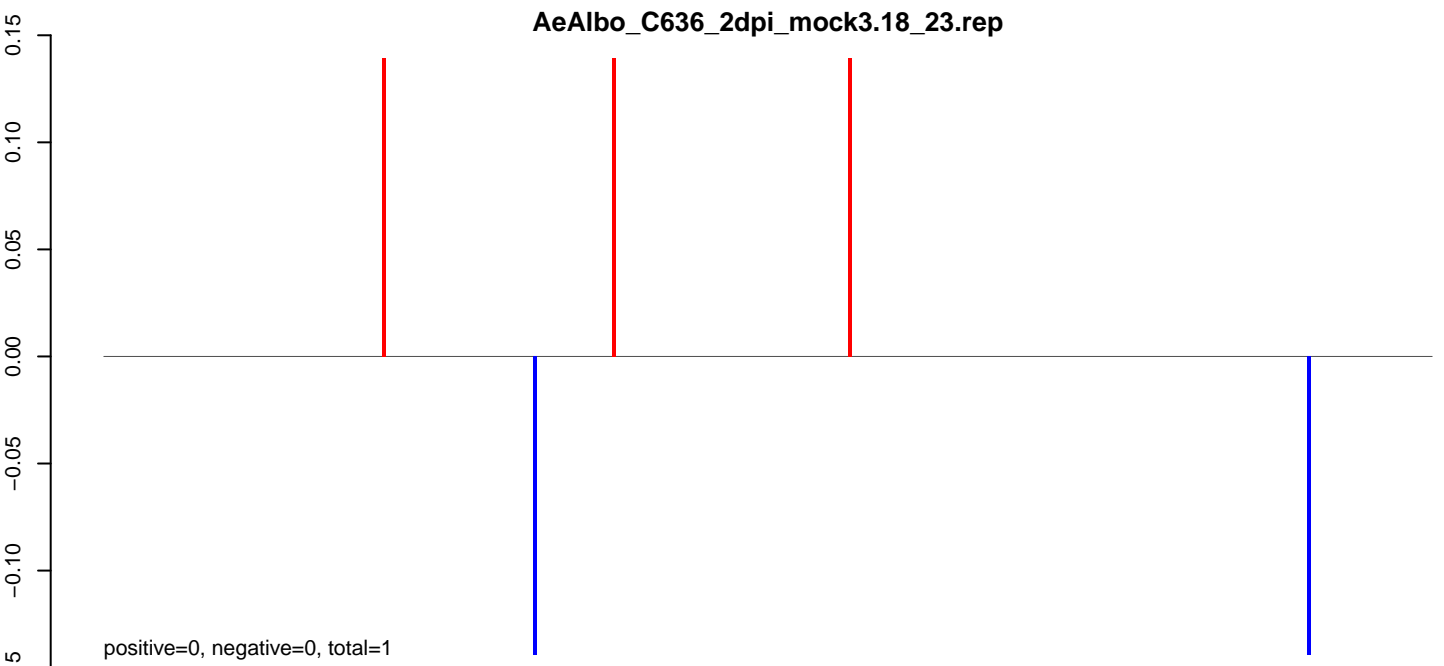


AeAlbo_C636_2dpi_mock3.24_35.rep

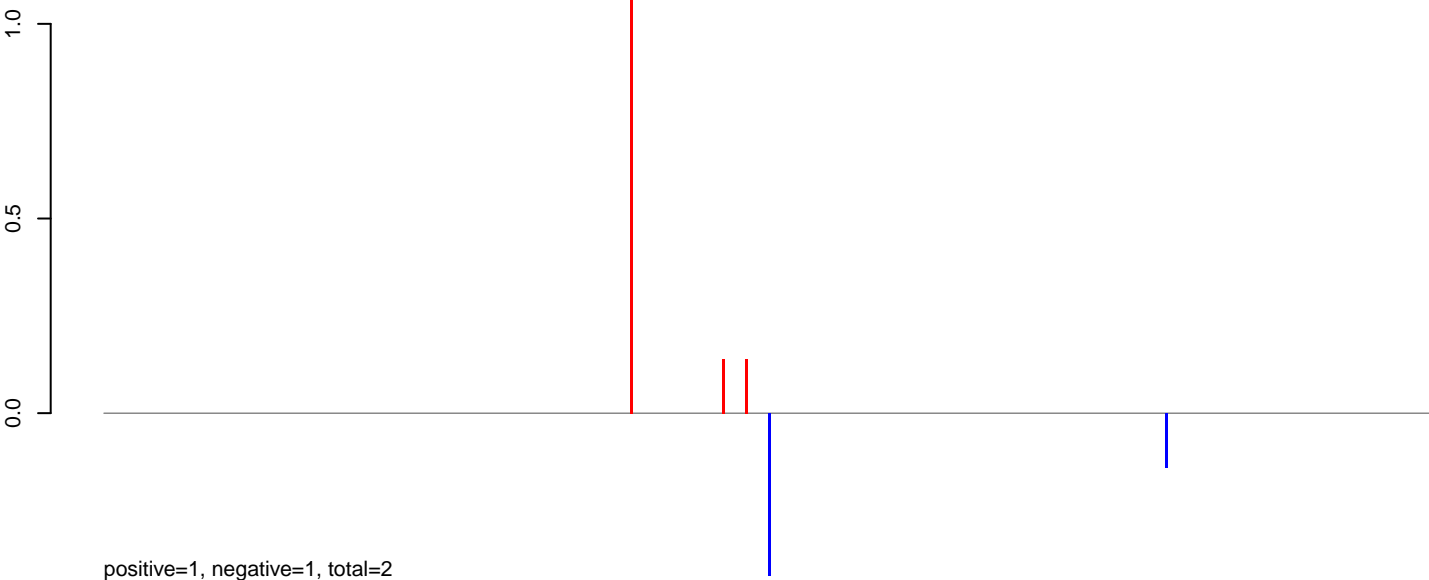


AeAlbo_C636_2dpi_mock3.rep





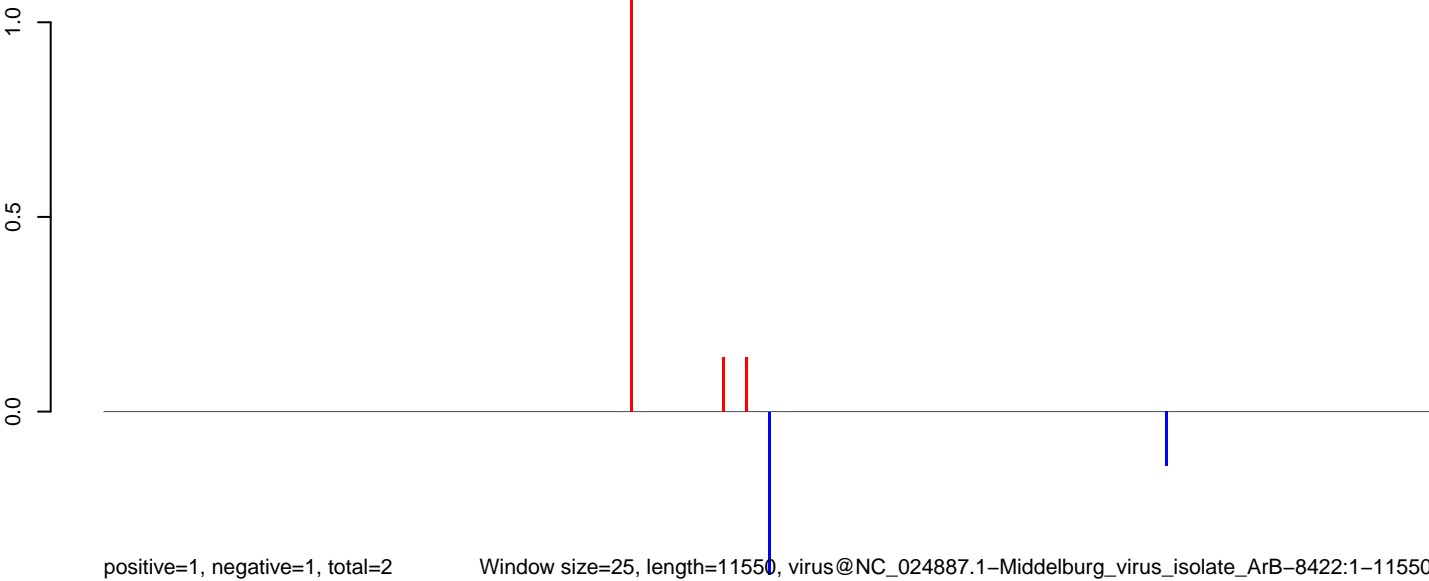
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

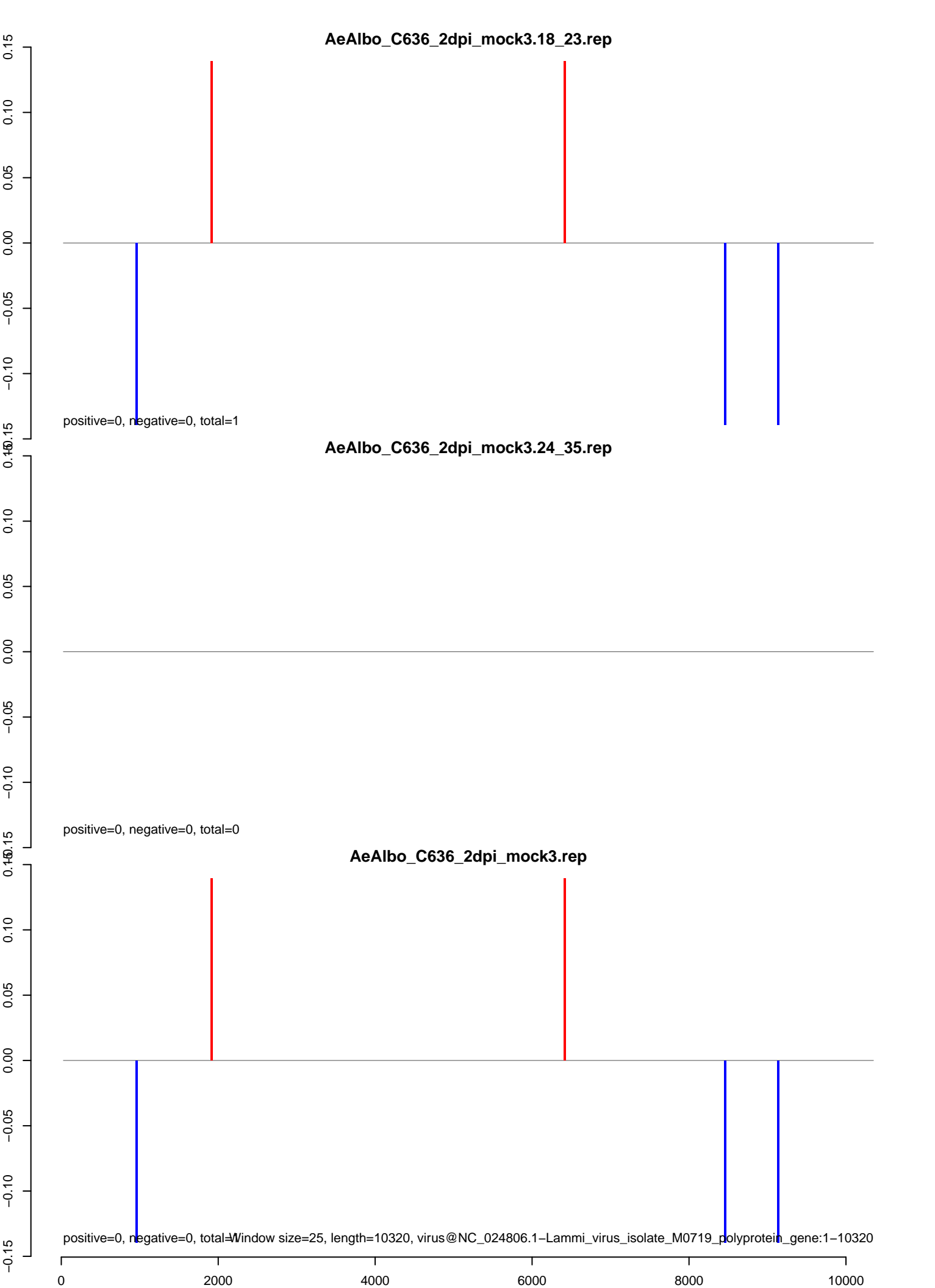


AeAlbo_C636_2dpi_mock3.rep

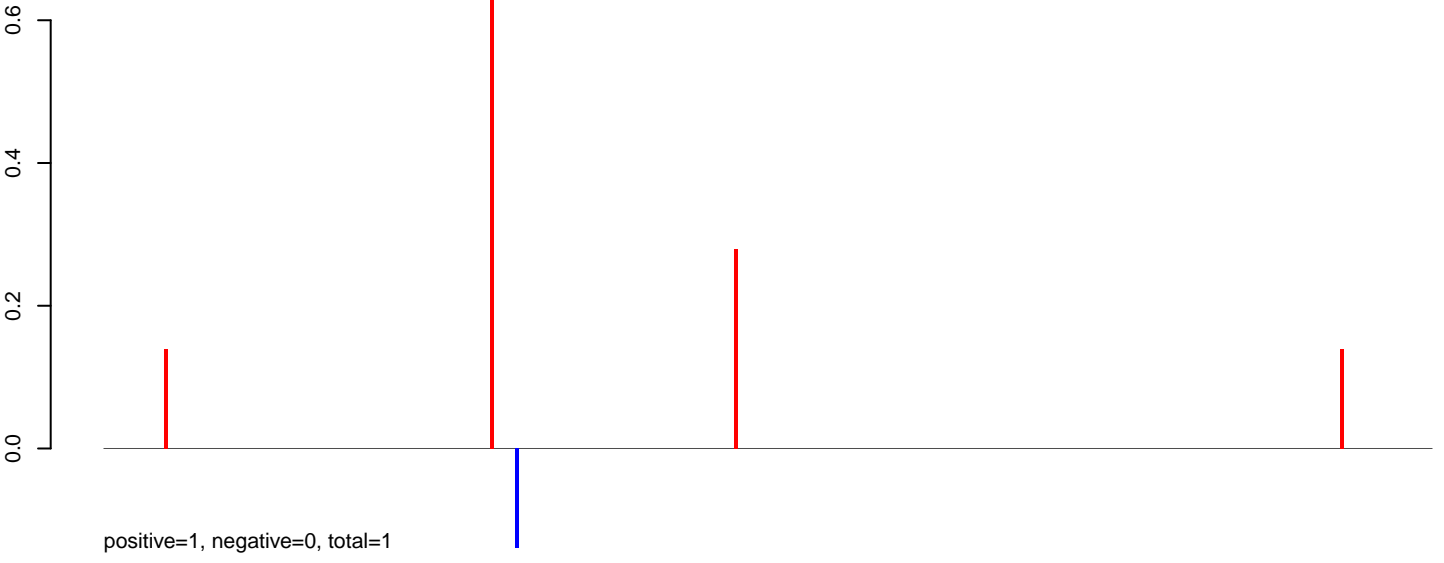


Window size=25, length=11550, virus@NC_024887.1-Middelburg_virus_isolate_ArB-8422:1-11550

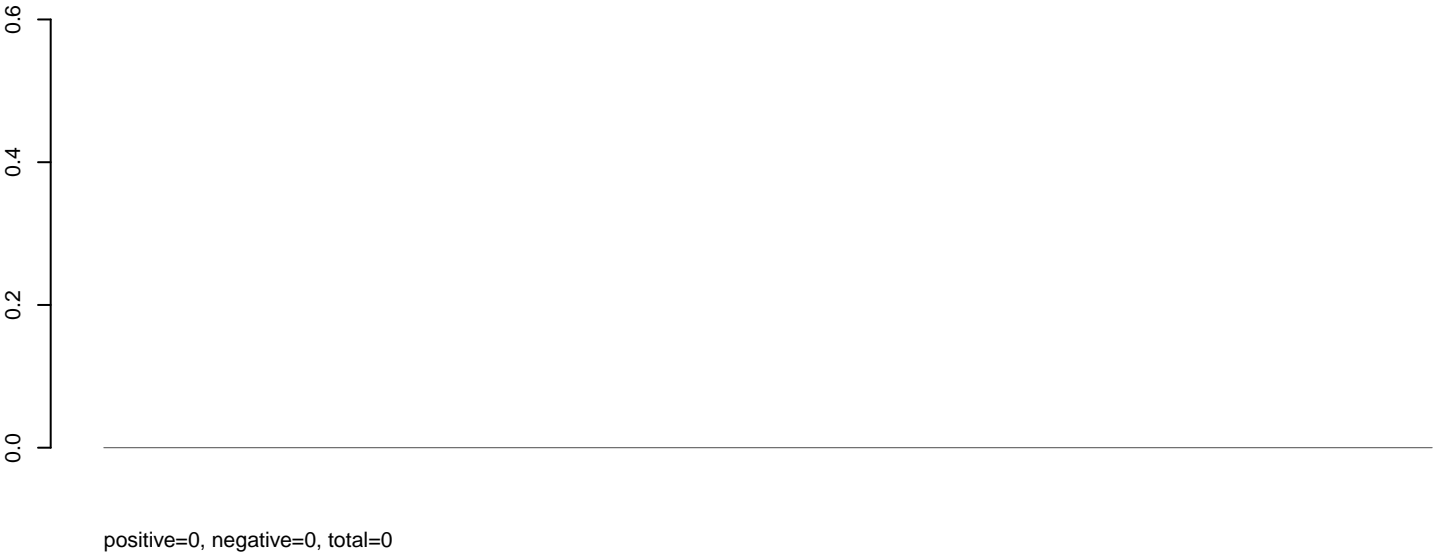
0 2000 4000 6000 8000 10000 12000



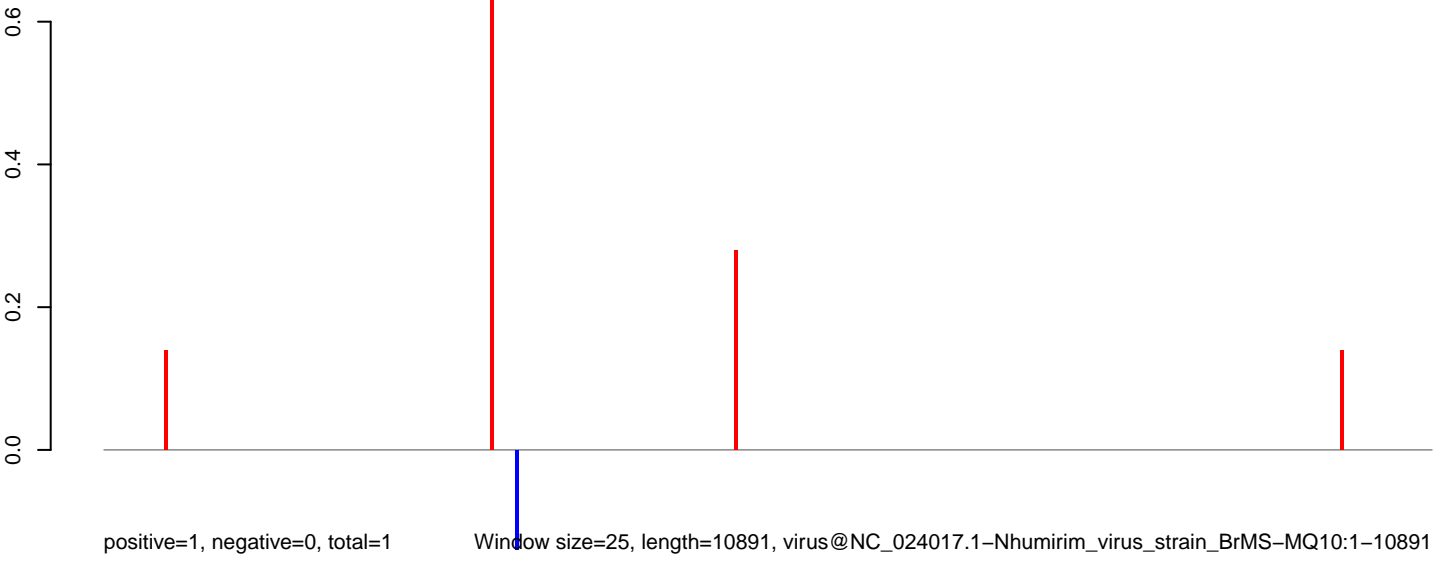
AeAlbo_C636_2dpi_mock3.18_23.rep

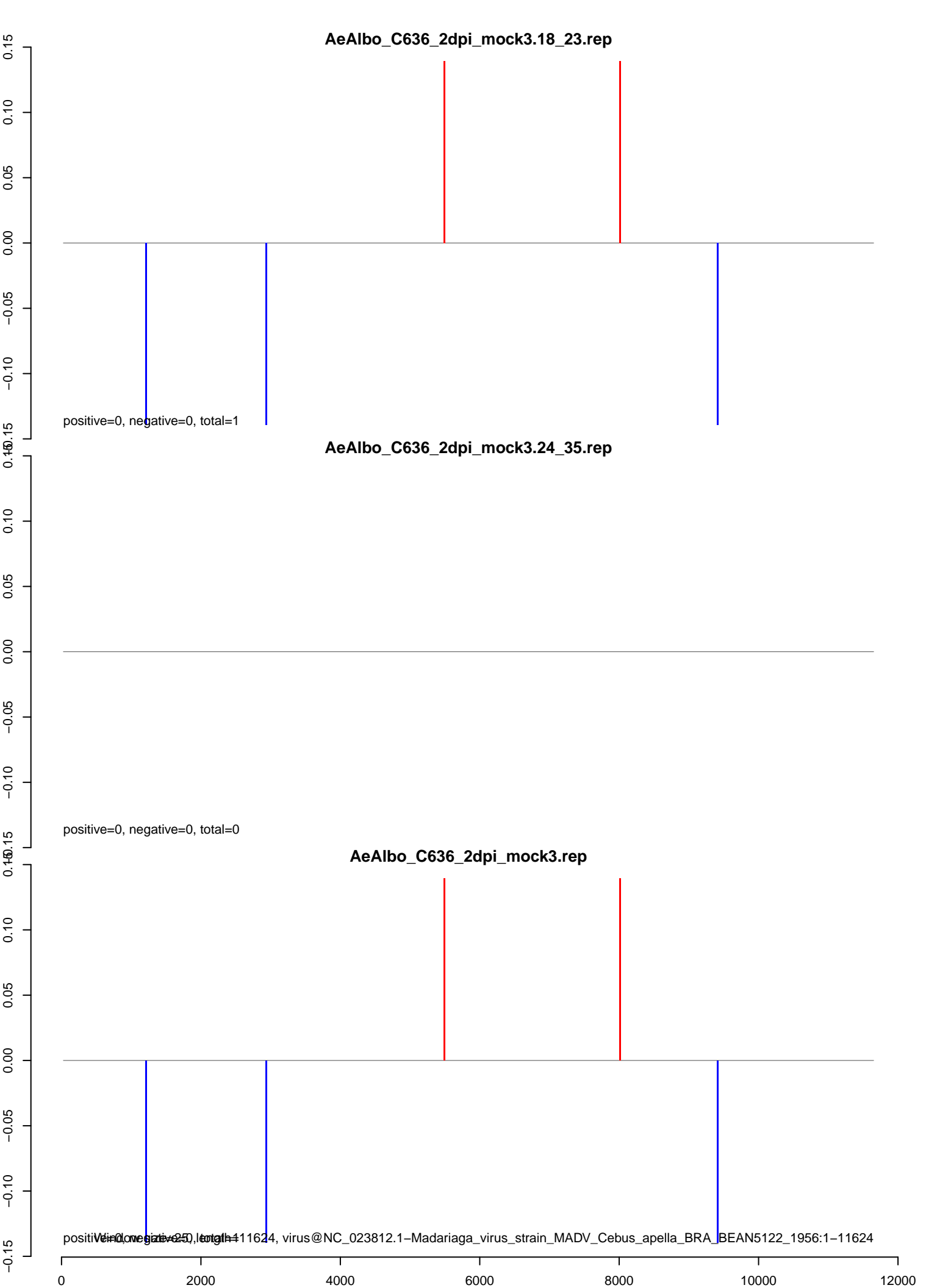


AeAlbo_C636_2dpi_mock3.24_35.rep

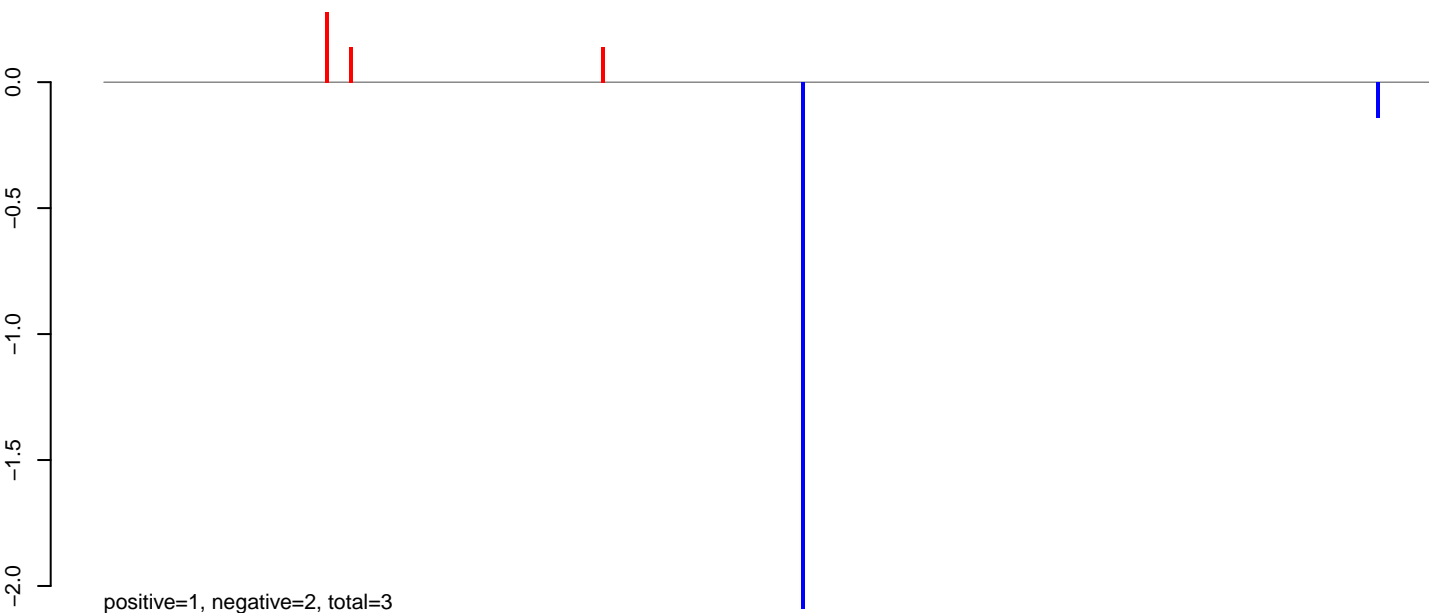


AeAlbo_C636_2dpi_mock3.rep

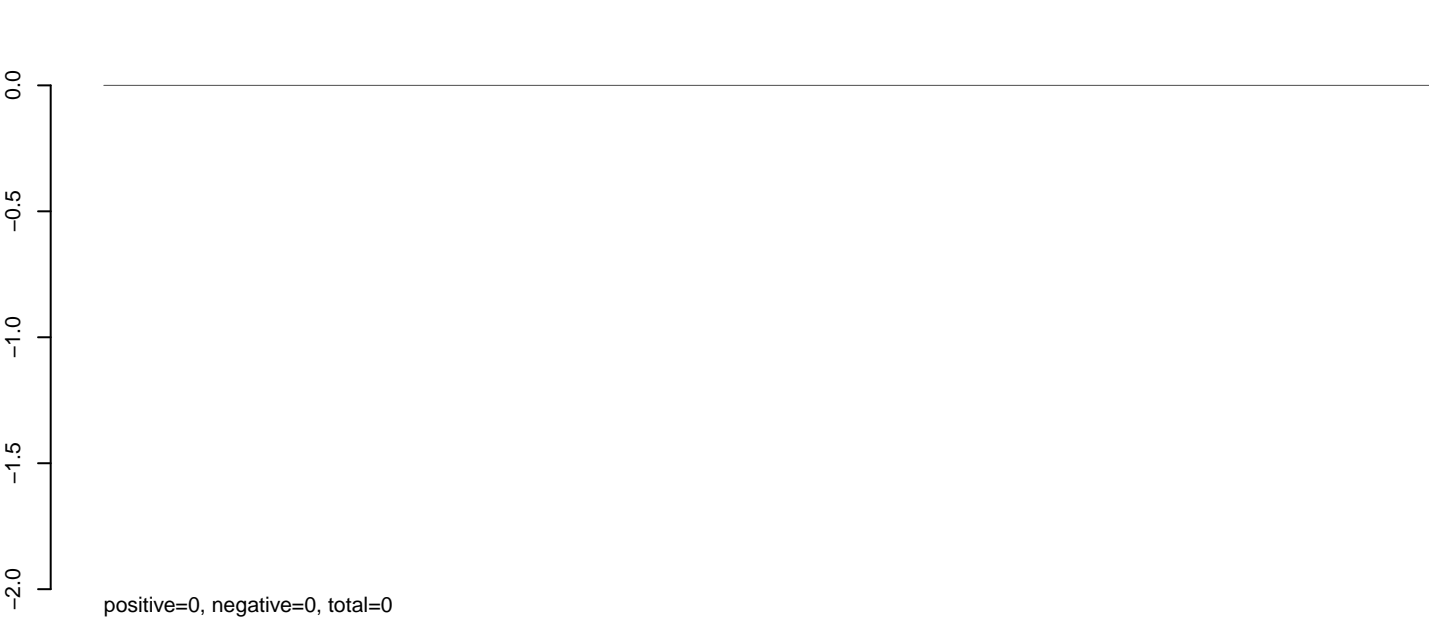




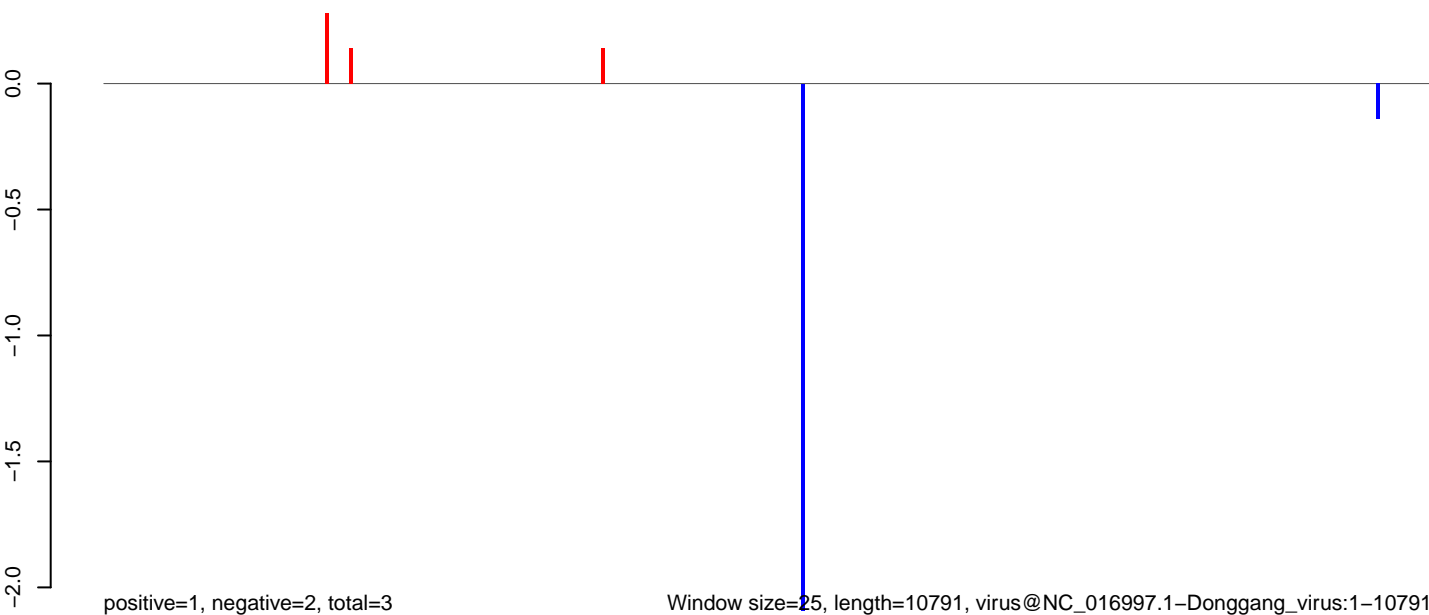
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

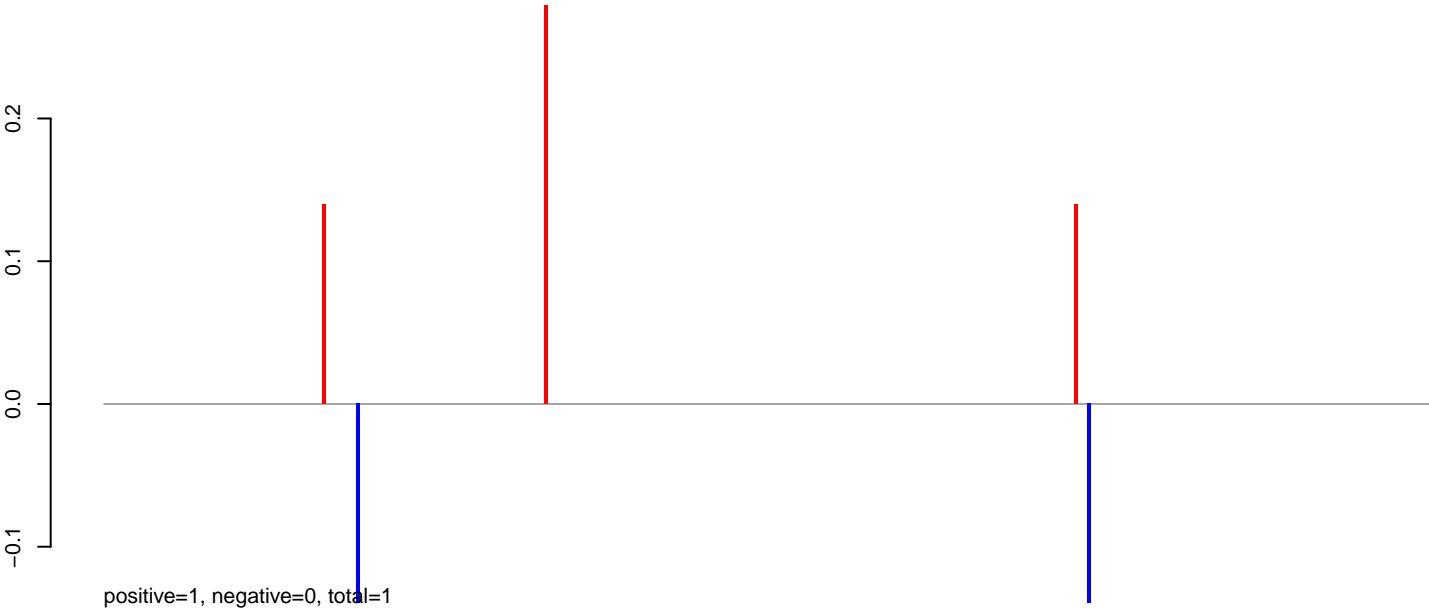


AeAlbo_C636_2dpi_mock3.rep

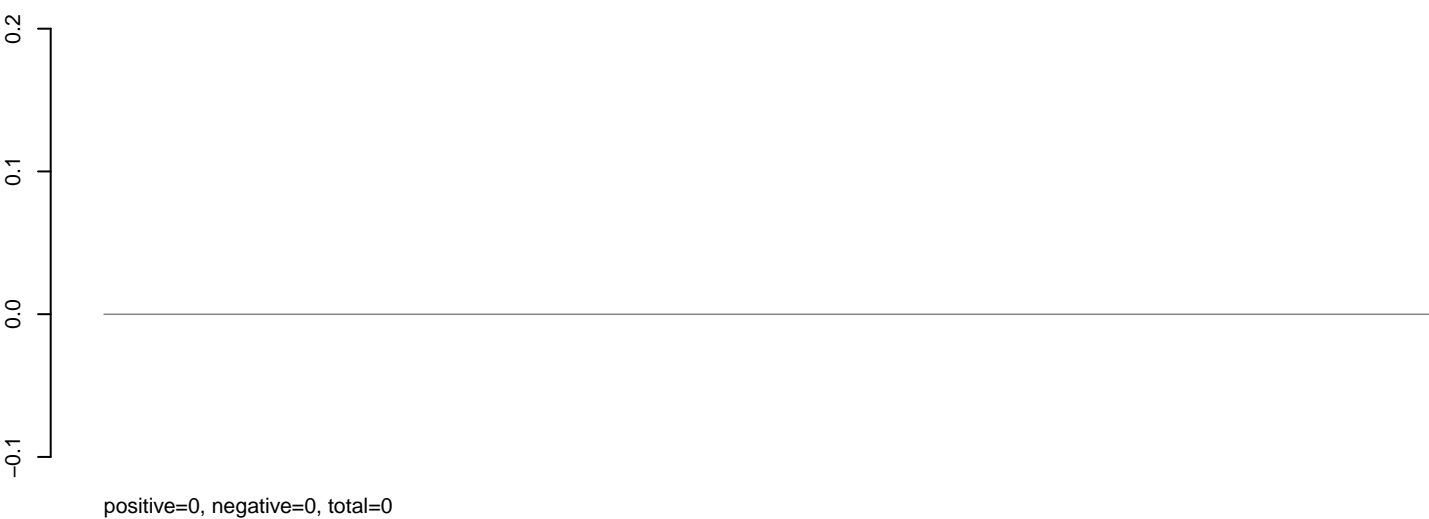


0 2000 4000 6000 8000 10000

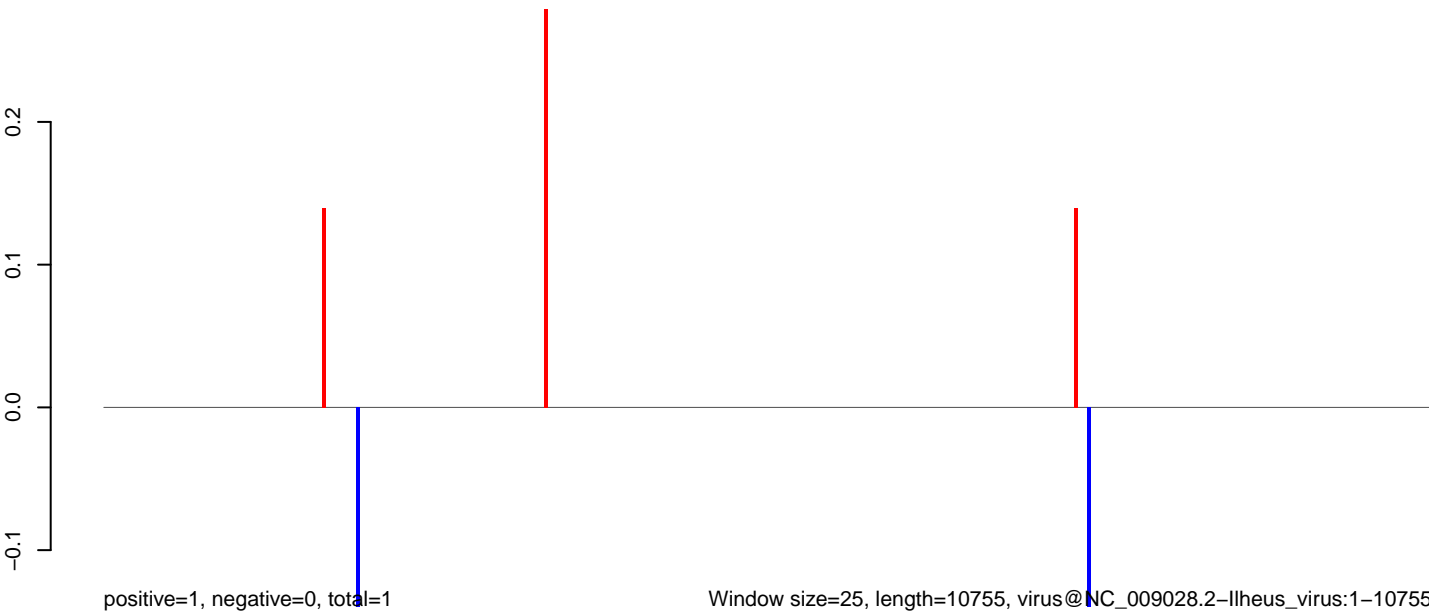
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



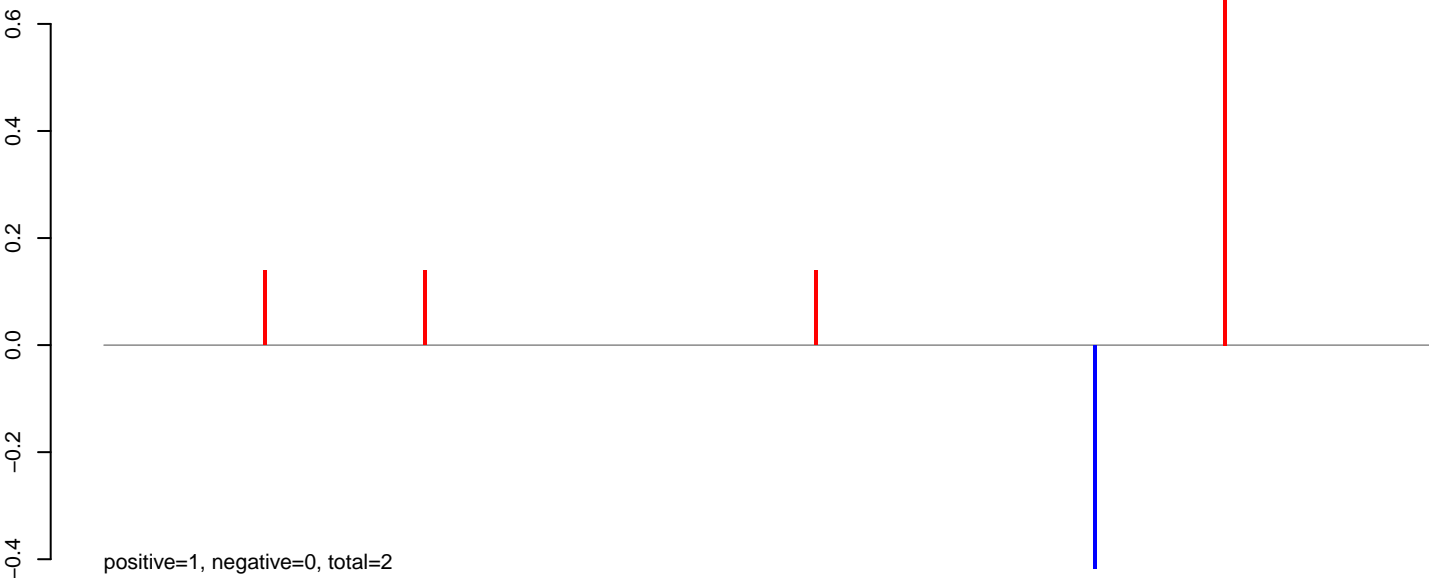
AeAlbo_C636_2dpi_mock3.rep



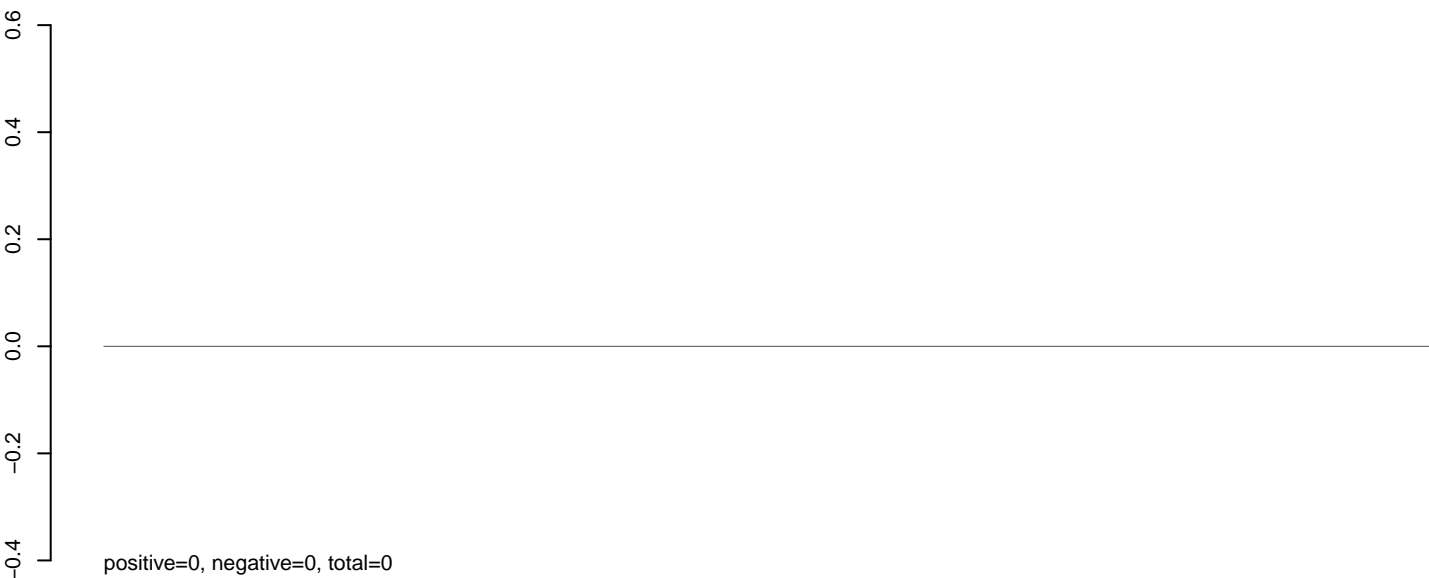
Window size=25, length=10755, virus@NC_009028.2-Ilheus_virus:1-10755

0 2000 4000 6000 8000 10000

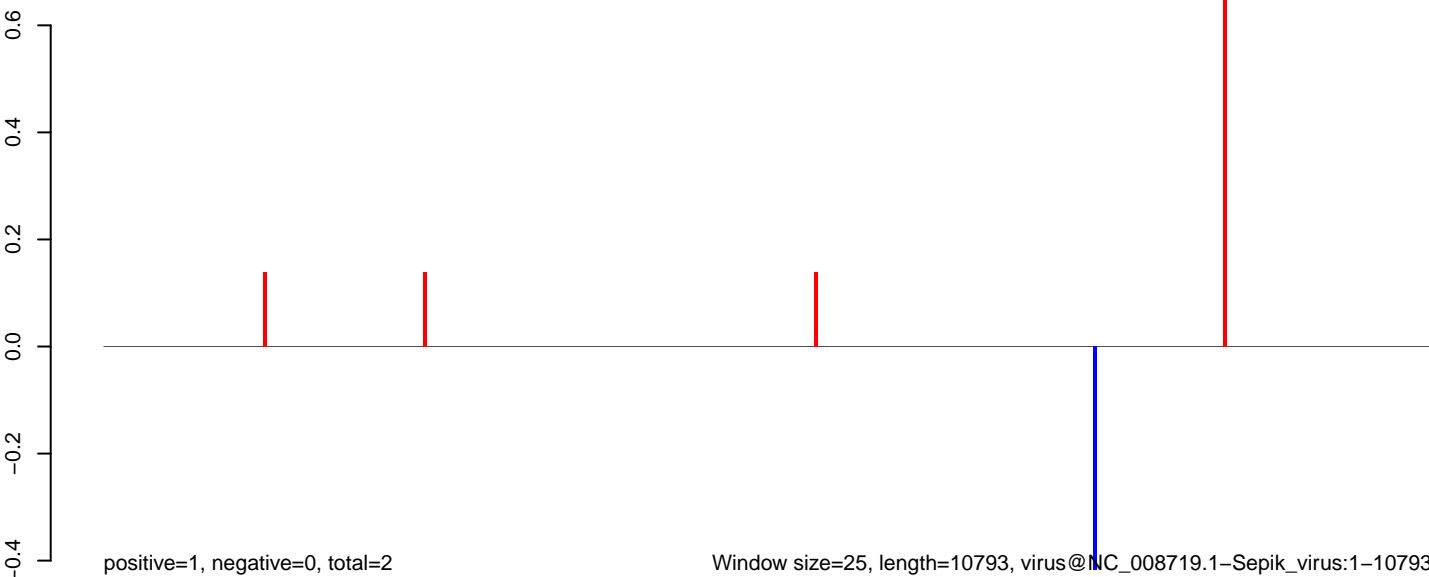
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

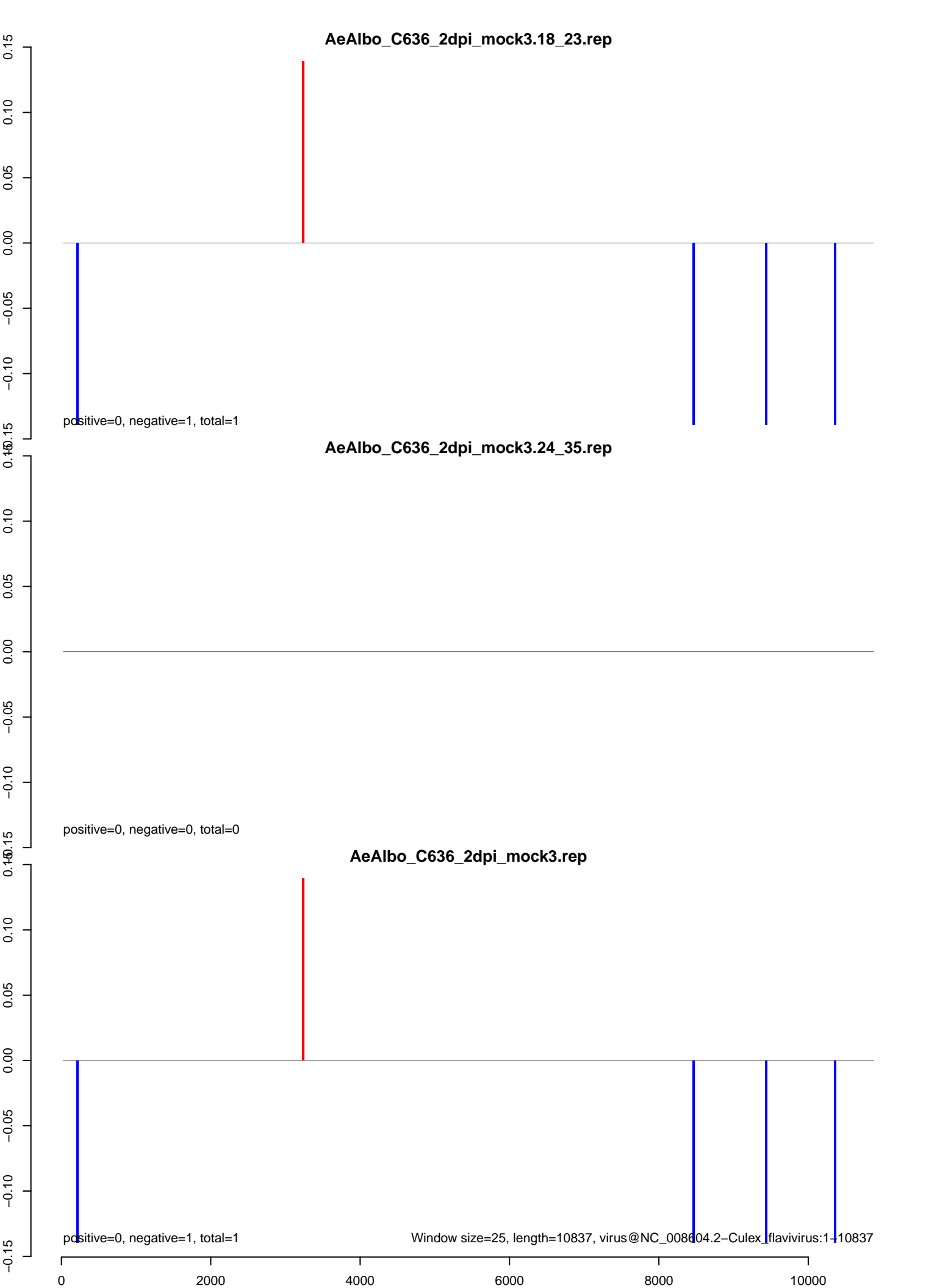


AeAlbo_C636_2dpi_mock3.rep

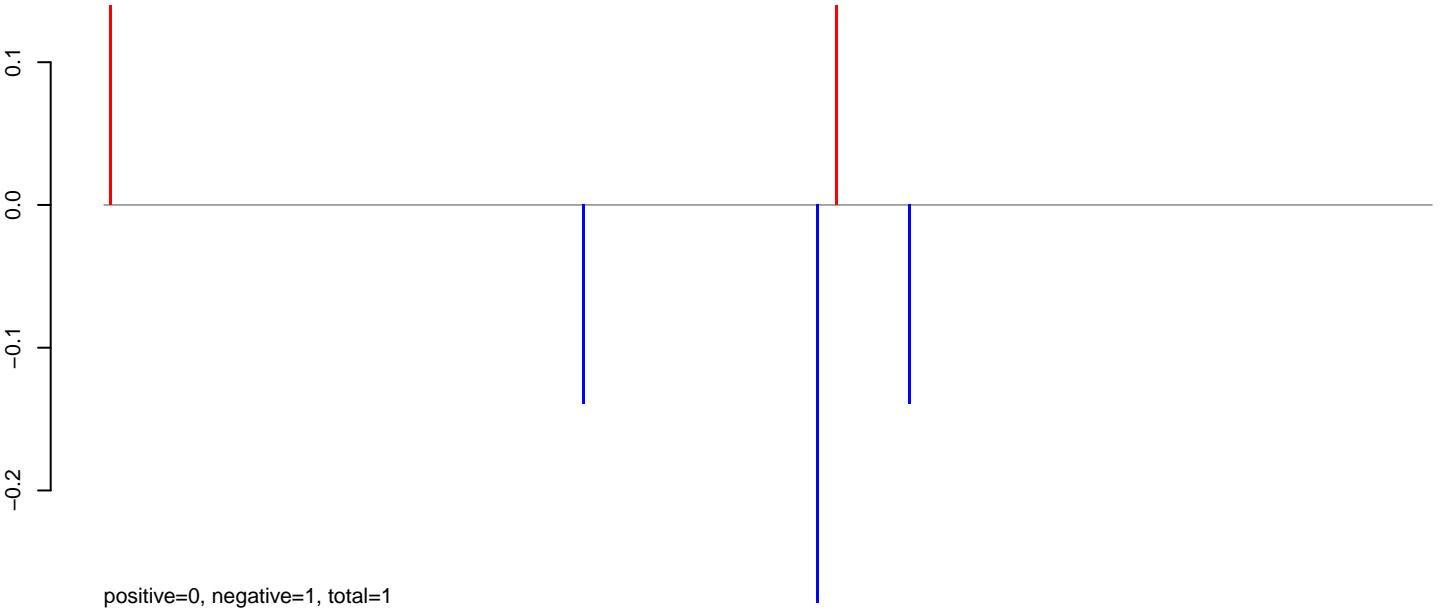


Window size=25, length=10793, virus@NC_008719.1-Sepik_virus:1-10793

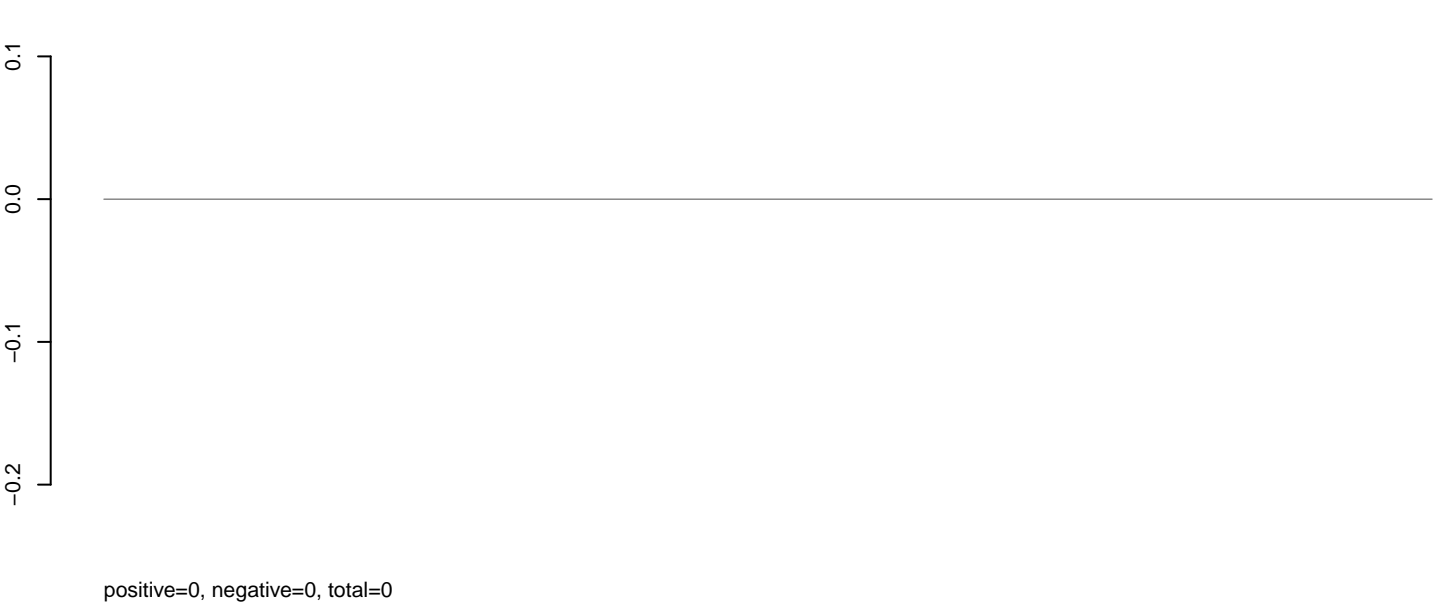
0 2000 4000 6000 8000 10000



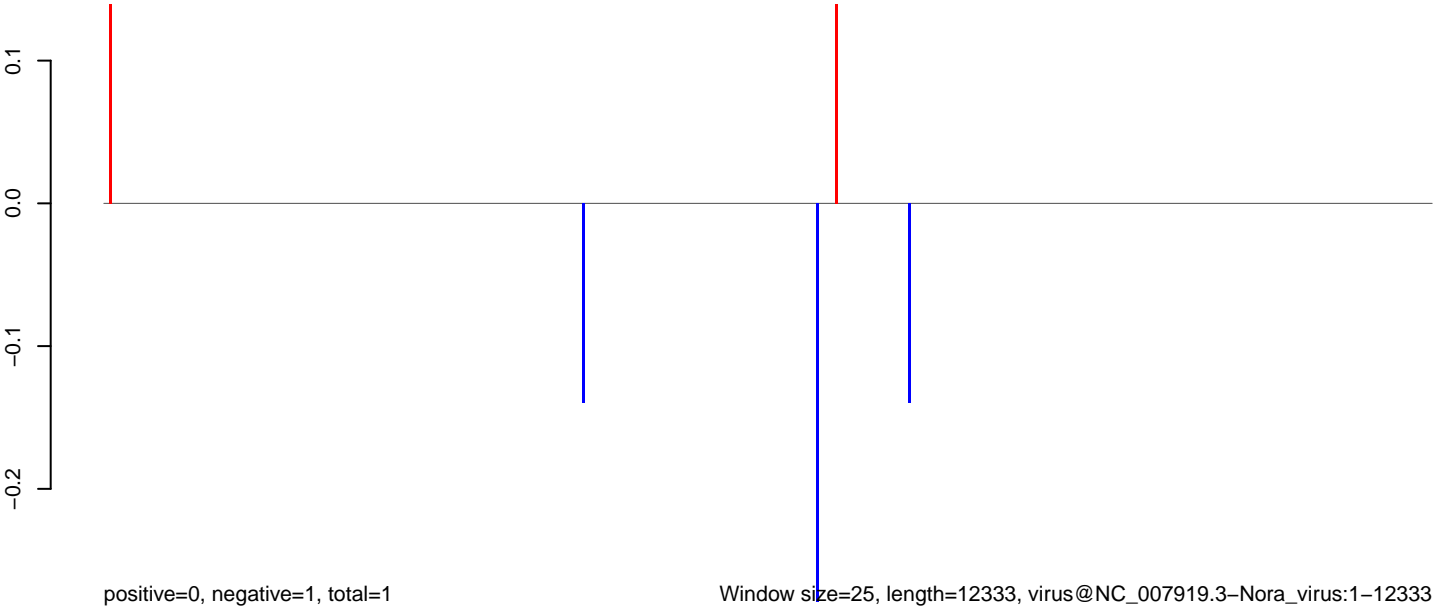
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



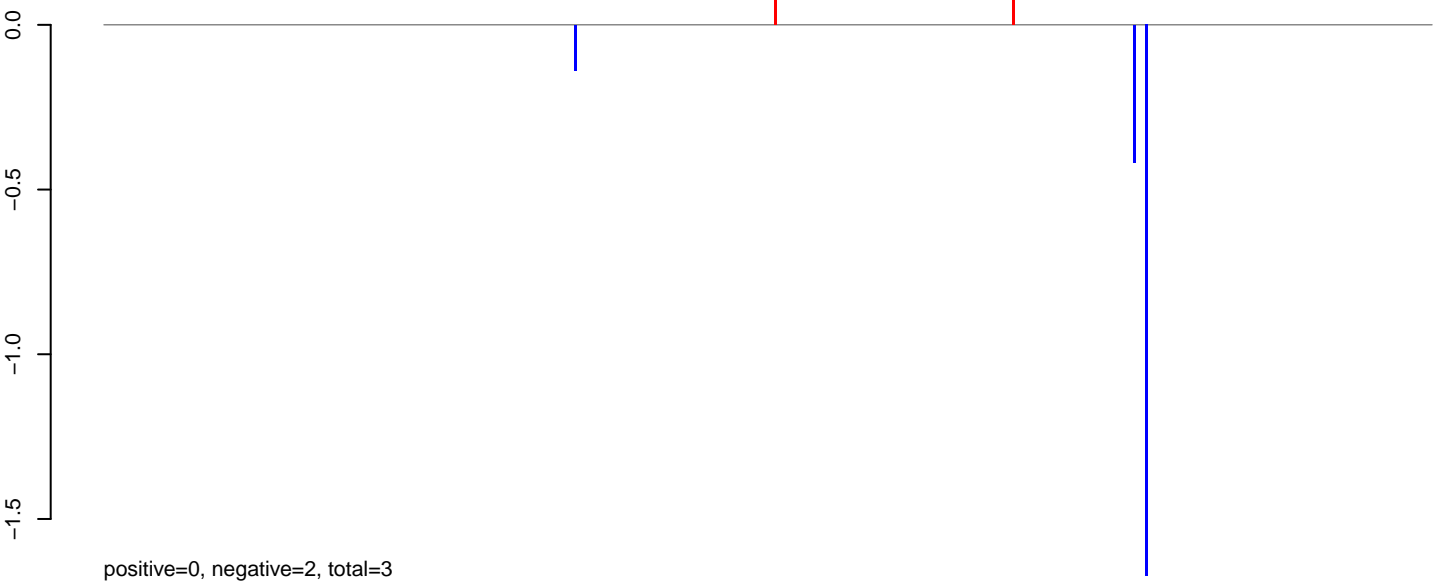
AeAlbo_C636_2dpi_mock3.rep



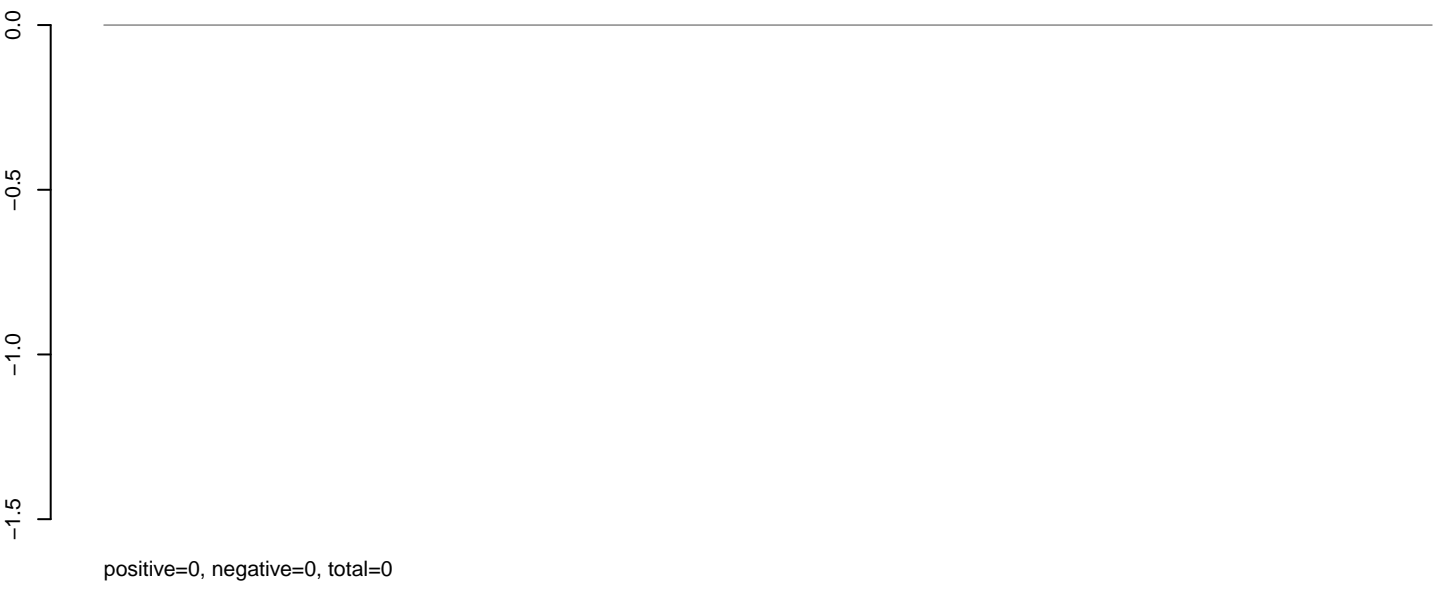
Window size=25, length=12333, virus@NC_007919.3-Nora_virus:1-12333

0 2000 4000 6000 8000 10000 12000

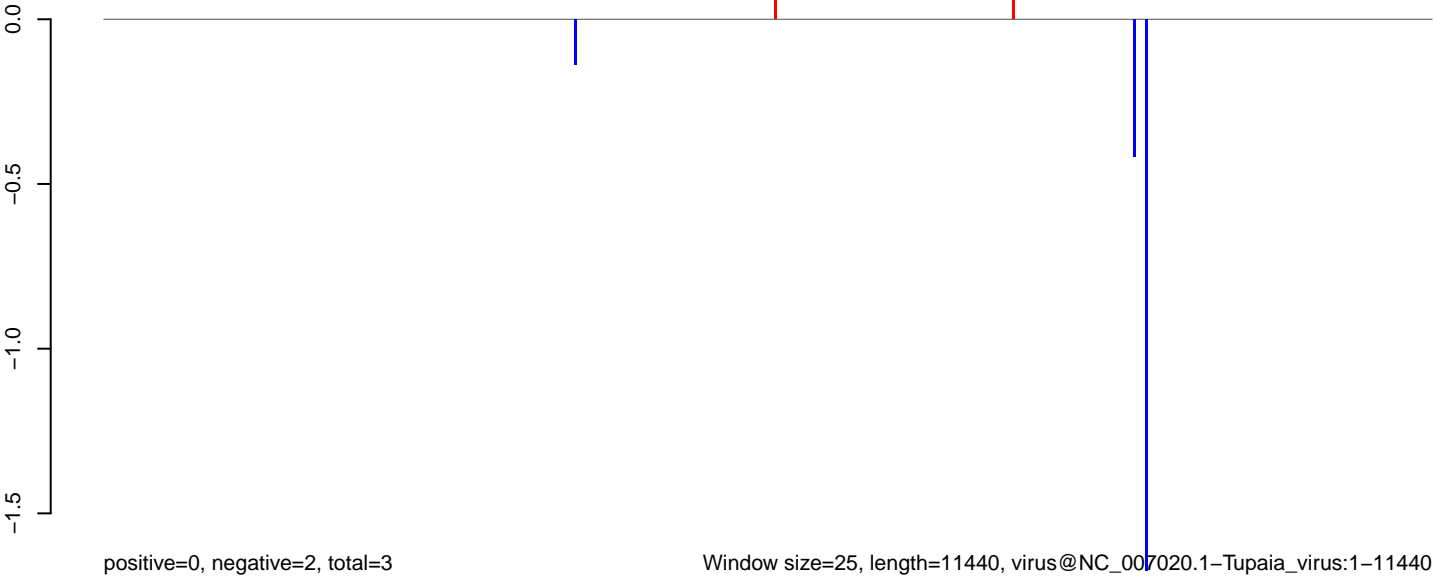
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

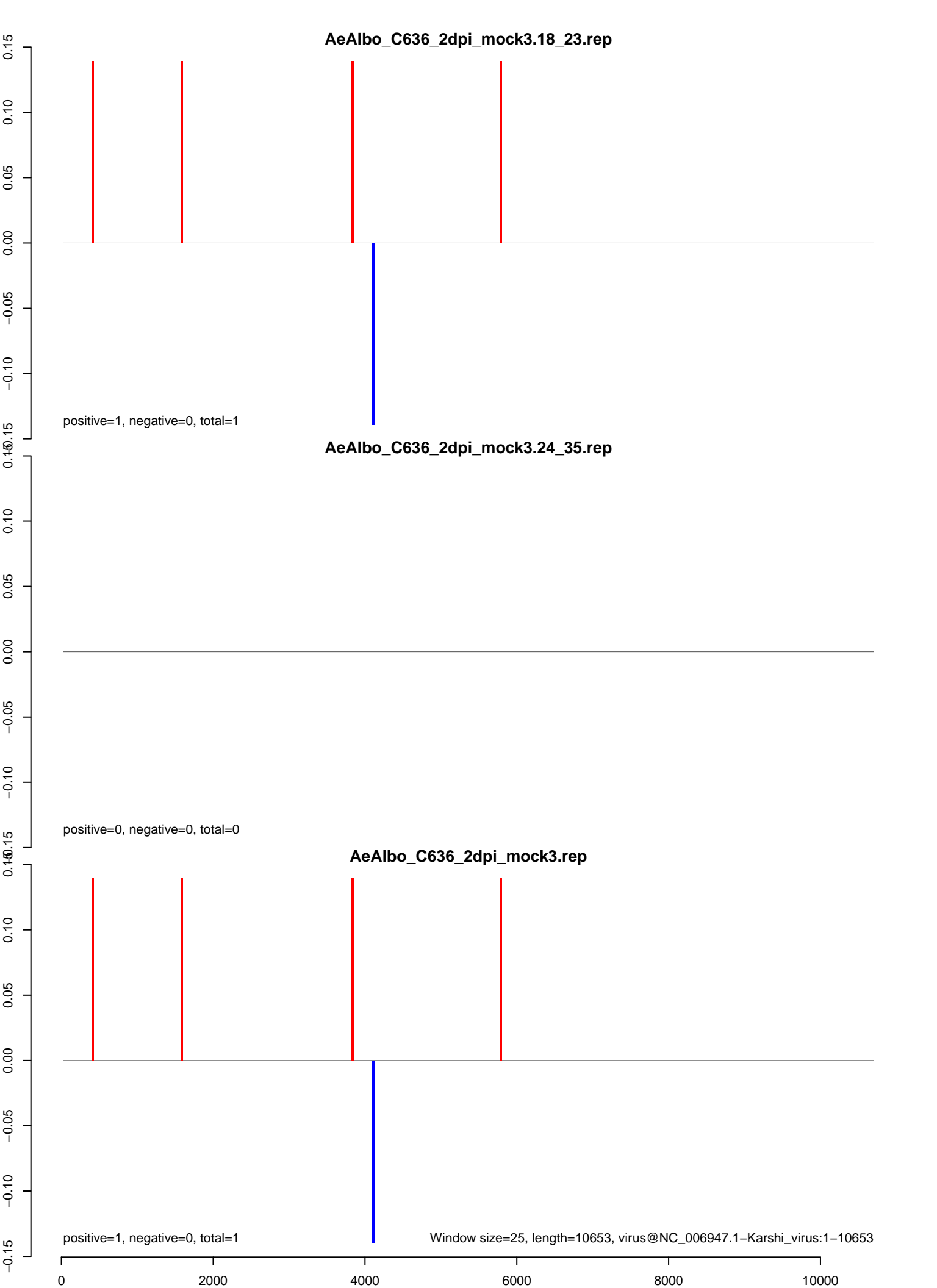


AeAlbo_C636_2dpi_mock3.rep

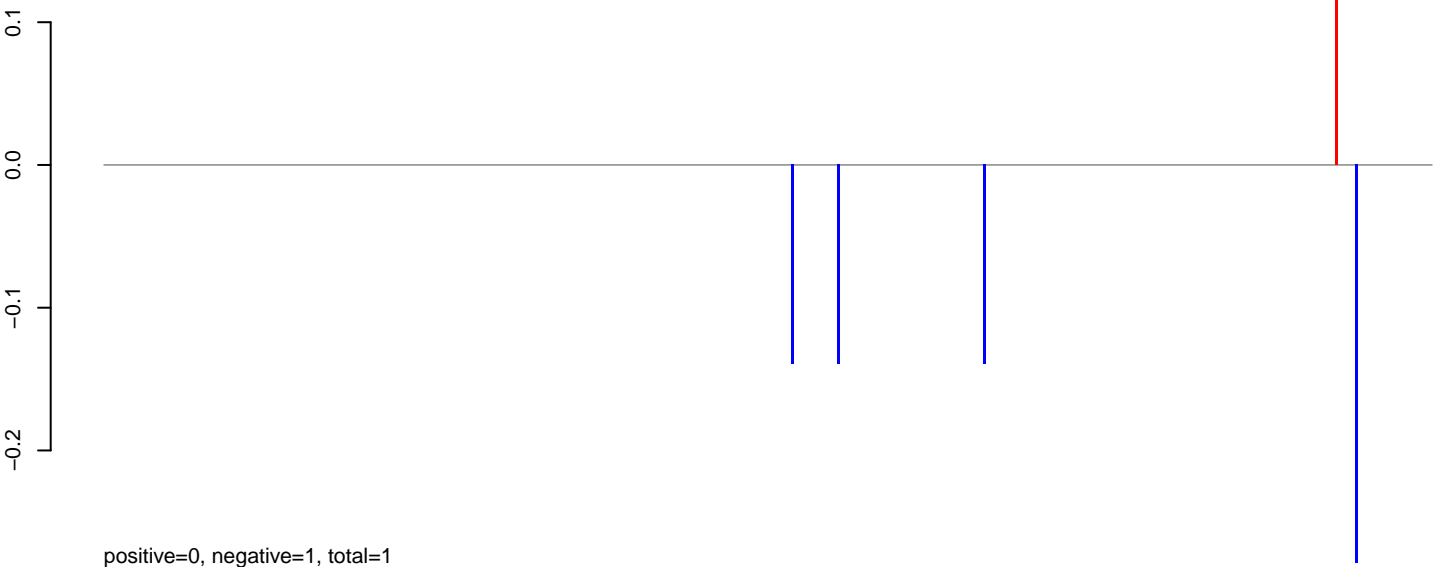


Window size=25, length=11440, virus@NC_007020.1-Tupaia_virus:1-11440

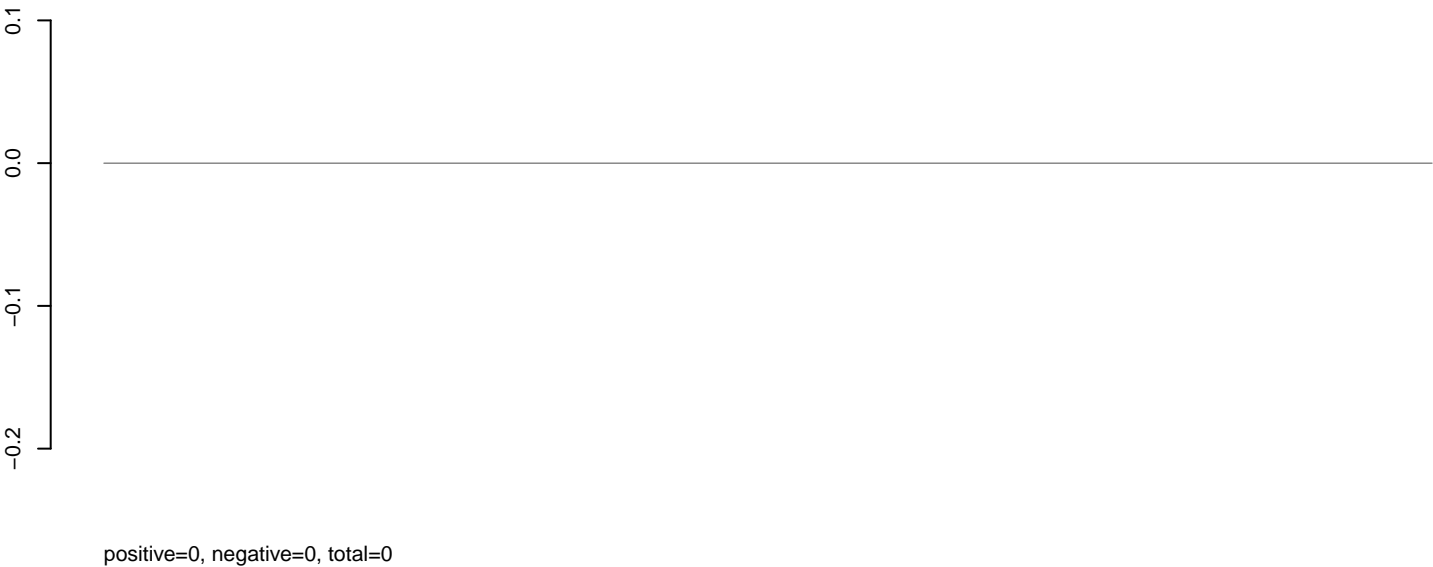
0 2000 4000 6000 8000 10000



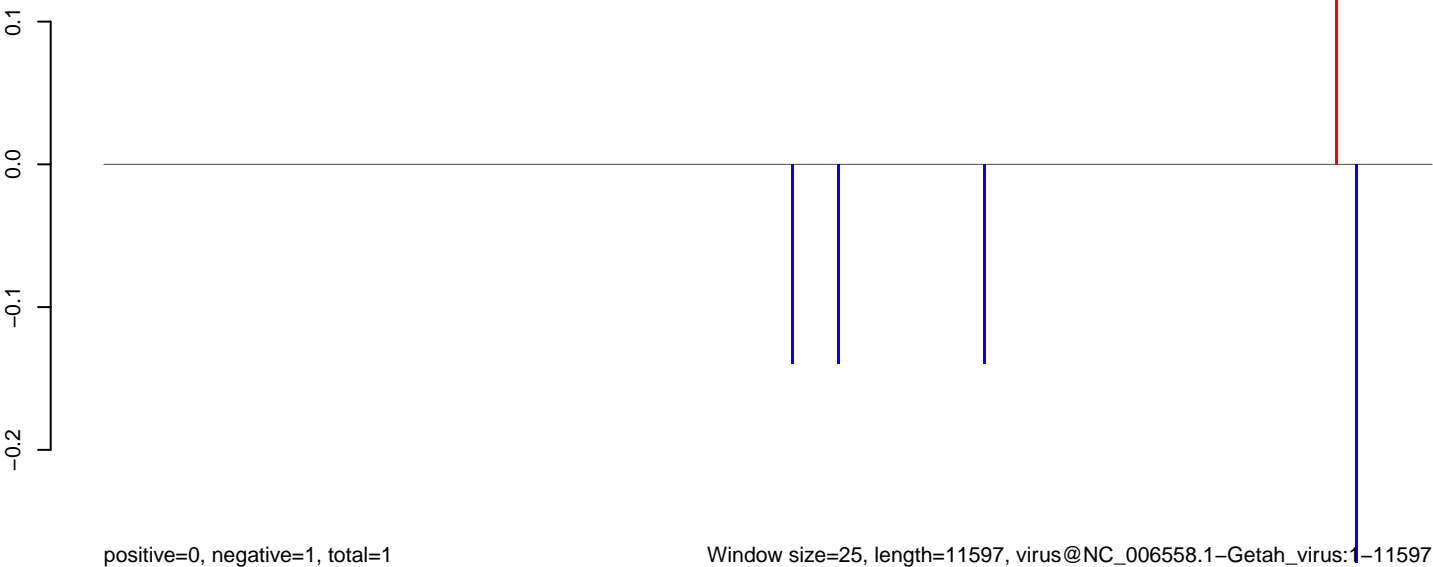
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

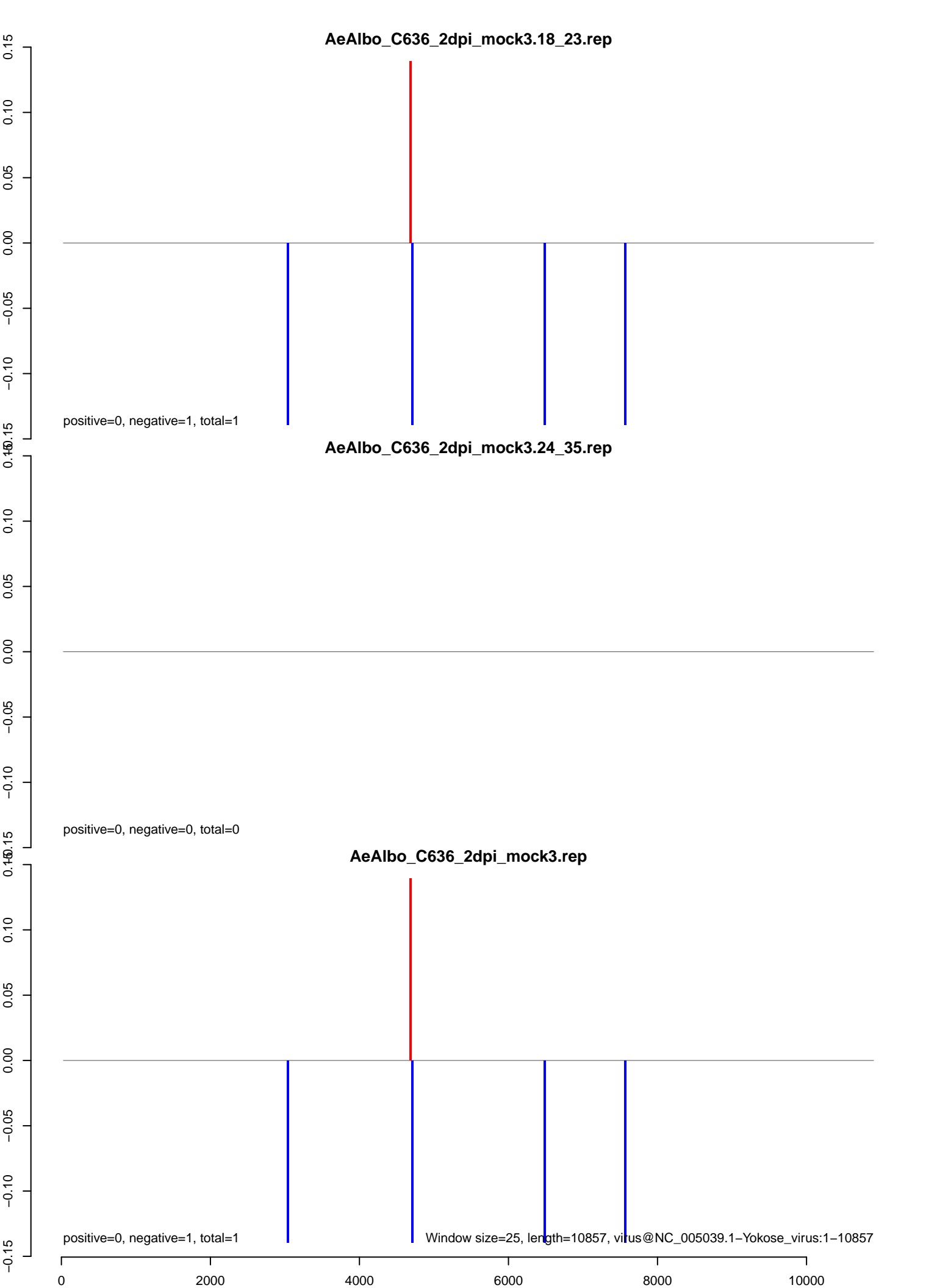


AeAlbo_C636_2dpi_mock3.rep

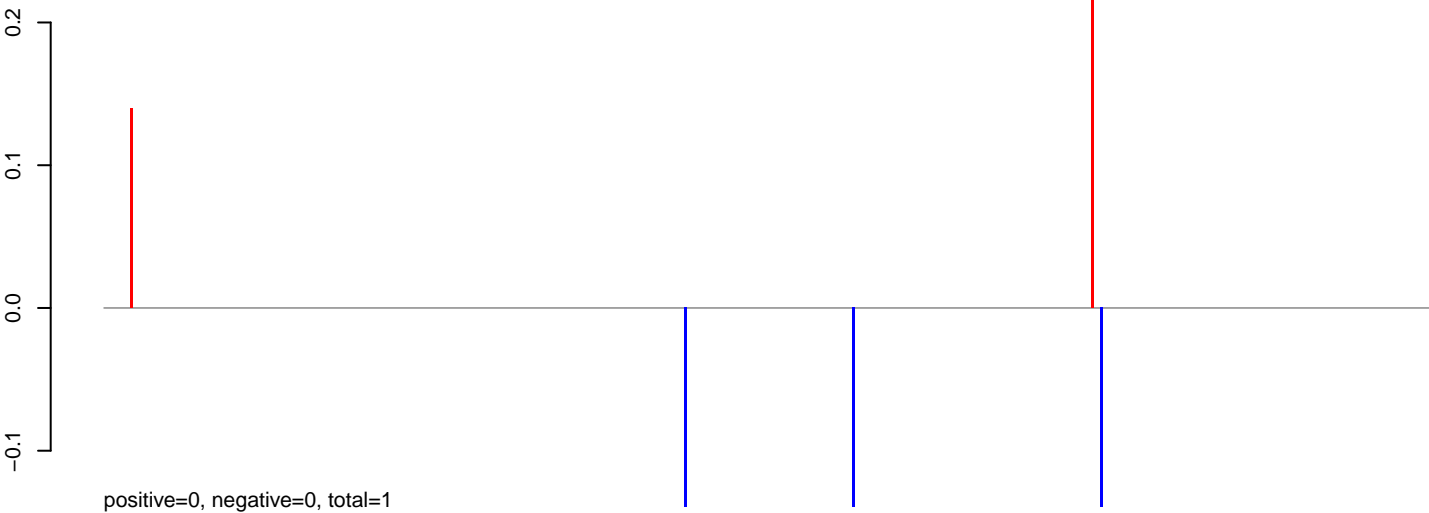


Window size=25, length=11597, virus@NC_006558.1-Getah_virus:1-11597

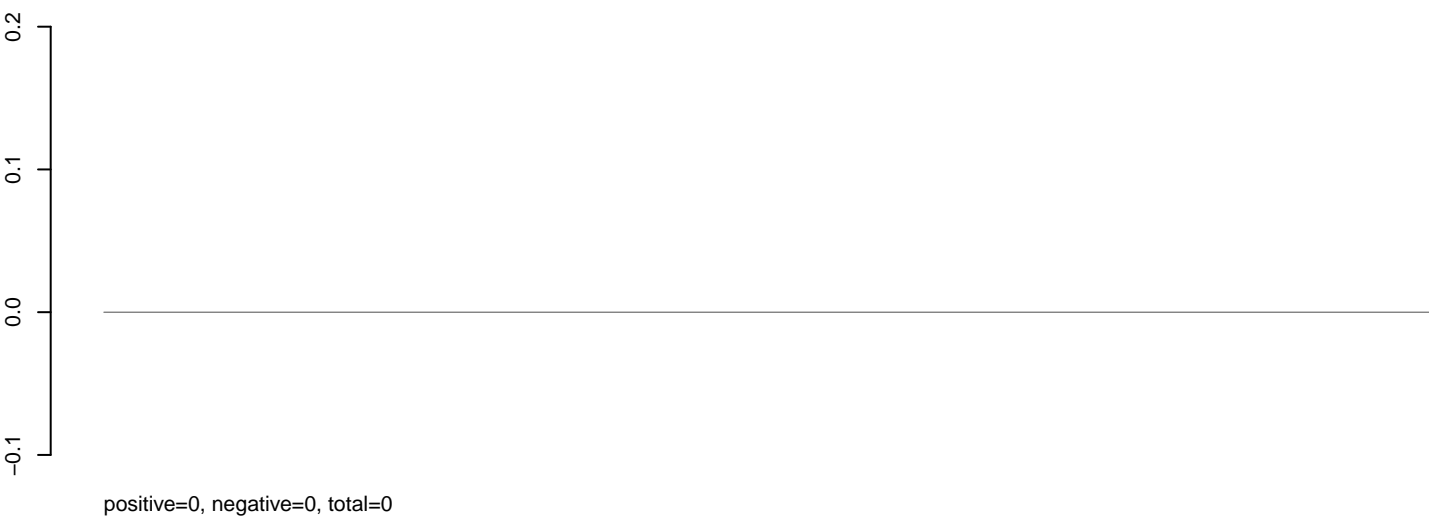
0 2000 4000 6000 8000 10000 12000



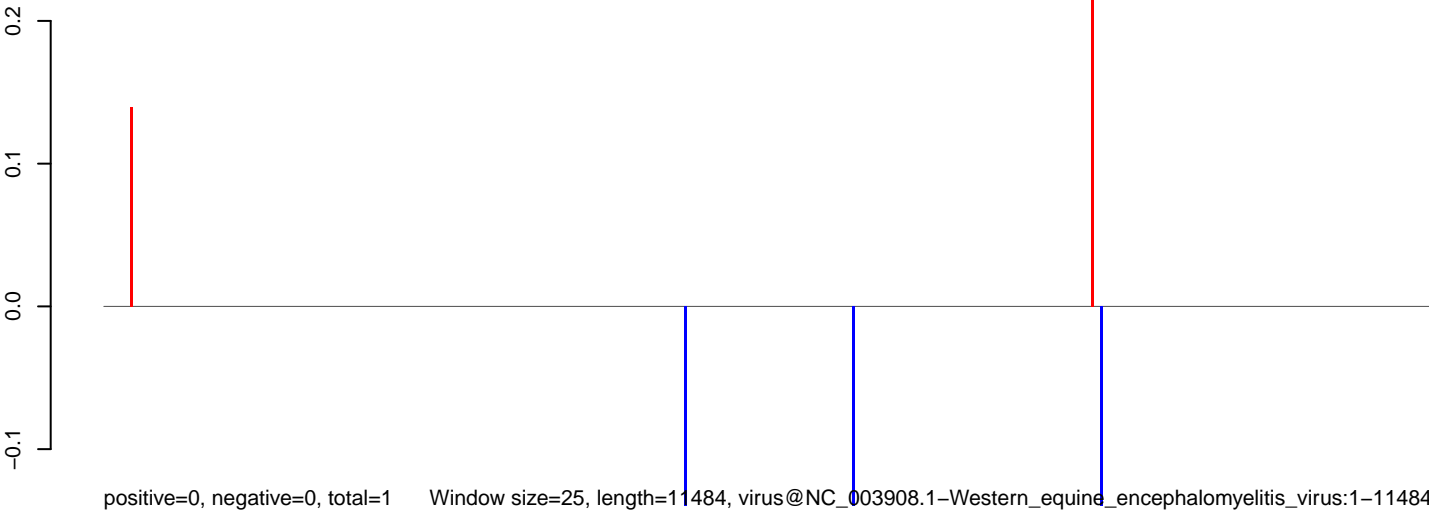
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



Window size=25, length=11484, virus@NC_003908.1-Western_equine_encephalomyelitis_virus:1-11484

AeAlbo_C636_2dpi_mock3.18_23.rep

positive=5, negative=0, total=5

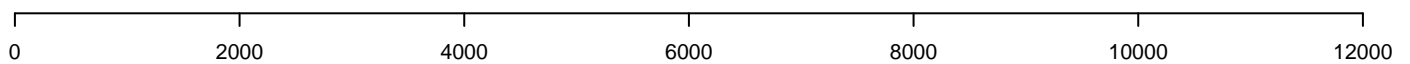
AeAlbo_C636_2dpi_mock3.24_35.rep

positive=0, negative=0, total=0

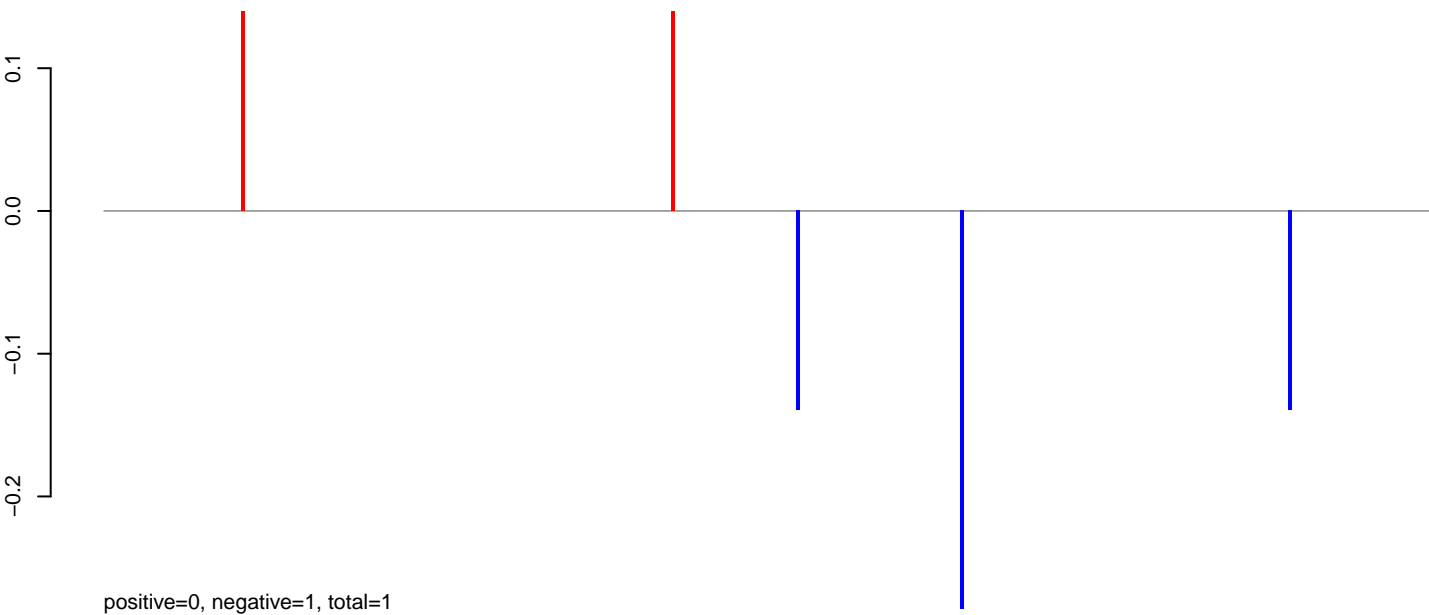
AeAlbo_C636_2dpi_mock3.rep

positive=5, negative=0, total=5

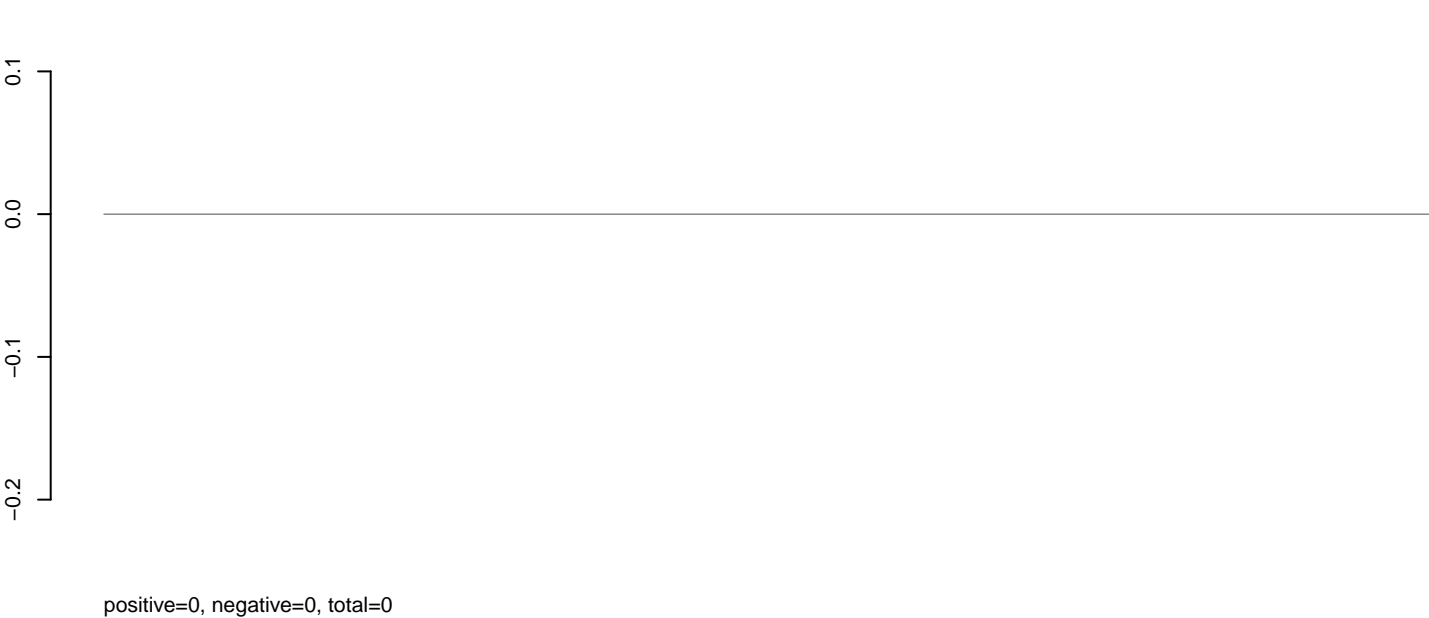
Window size=25, length=11824, virus@NC_003900.1-Aura_virus:1-11824



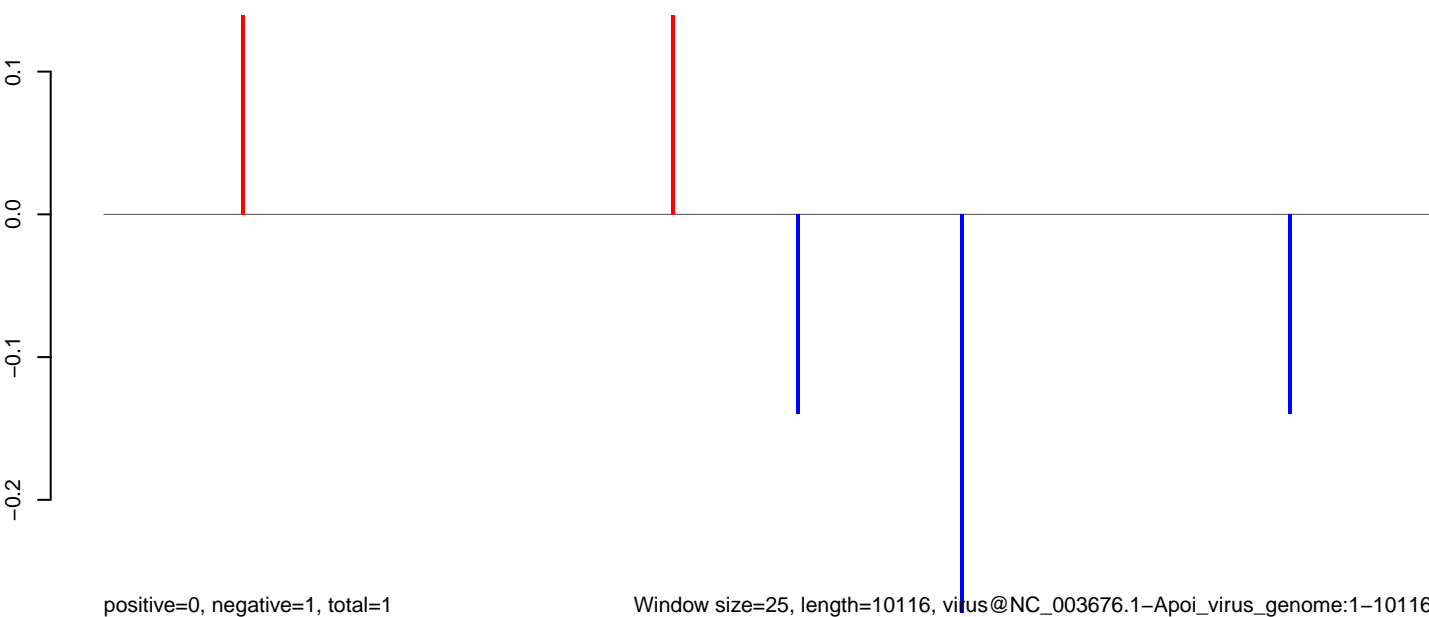
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



Window size=25, length=10116, virus@NC_003676.1-Apoi_virus_genome:1-10116

0 2000 4000 6000 8000 10000

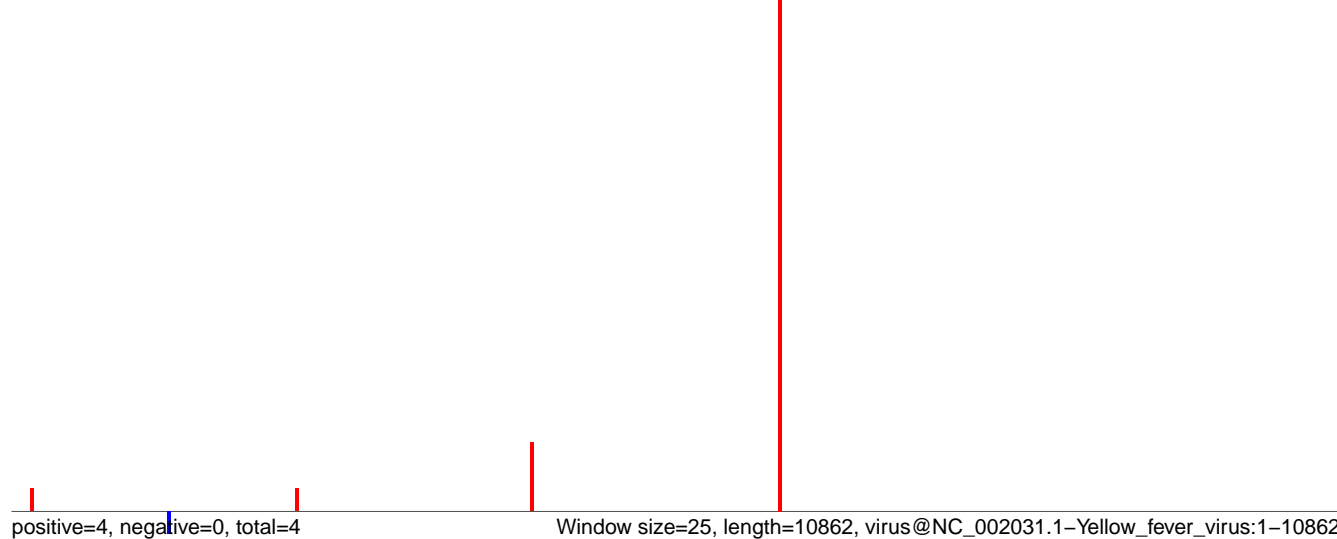
AeAlbo_C636_2dpi_mock3.18_23.rep



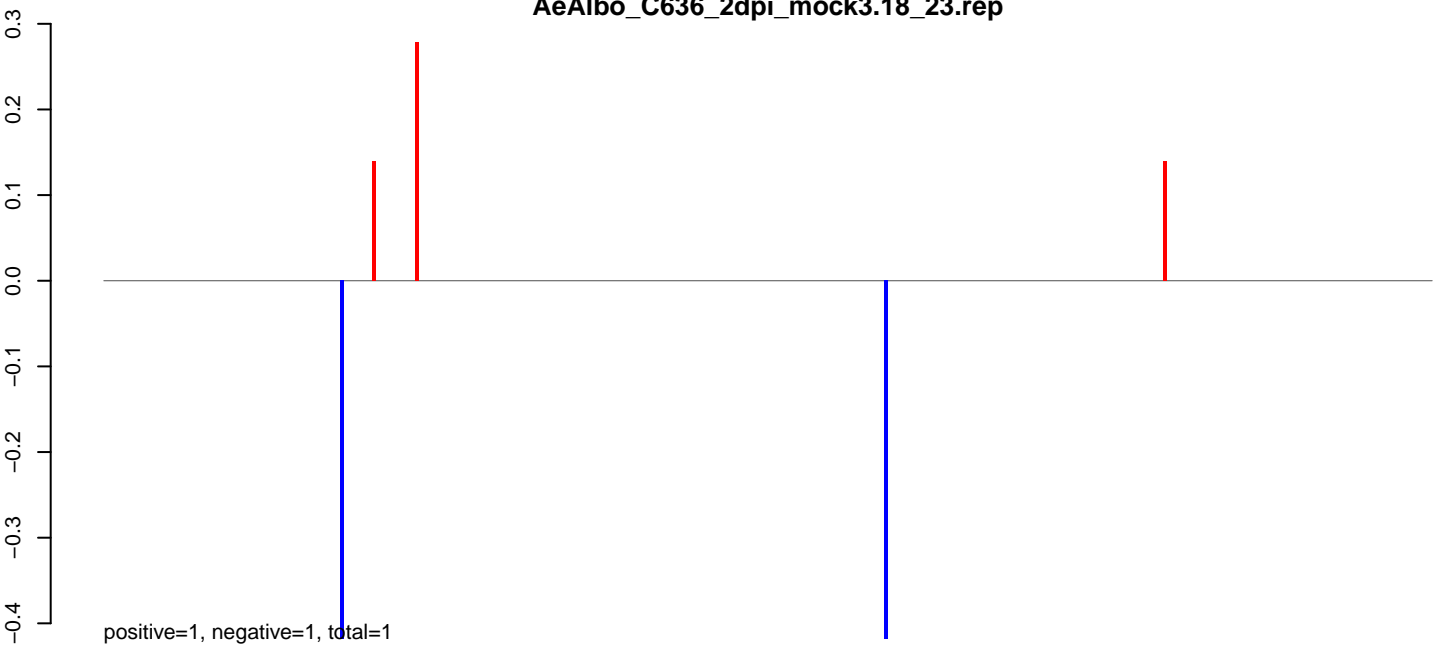
AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep

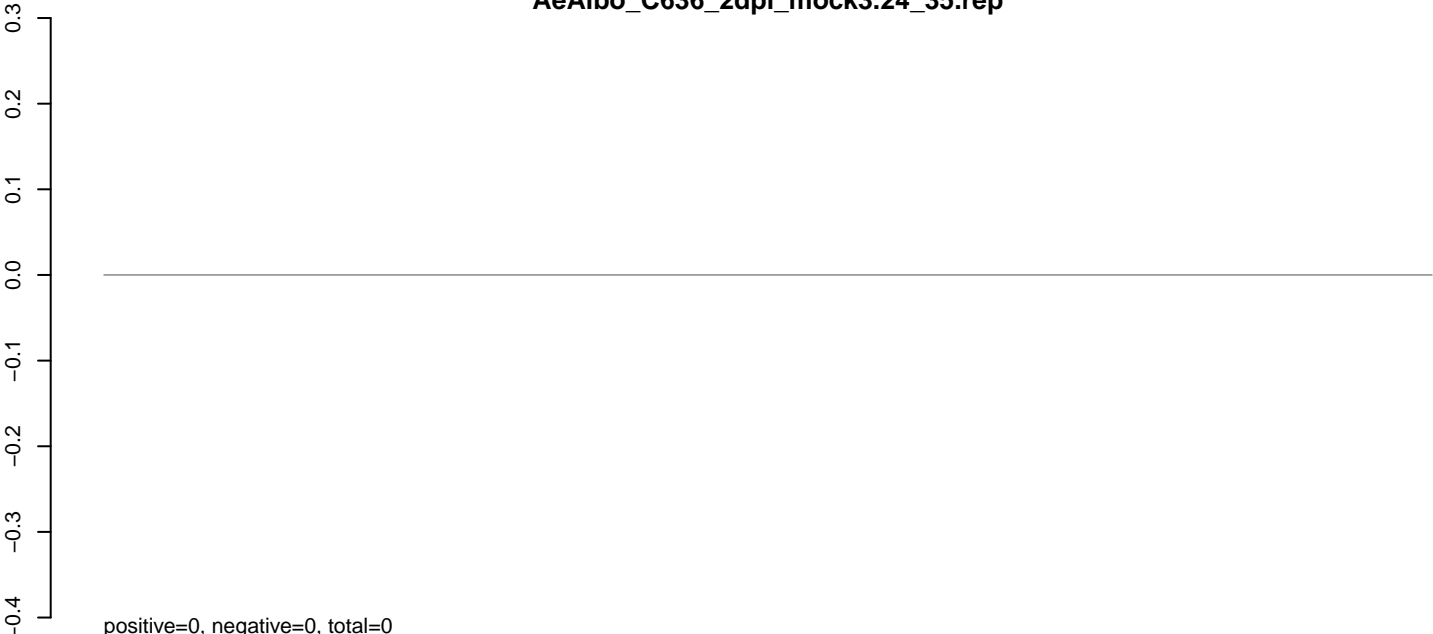


AeAlbo_C636_2dpi_mock3.18_23.rep



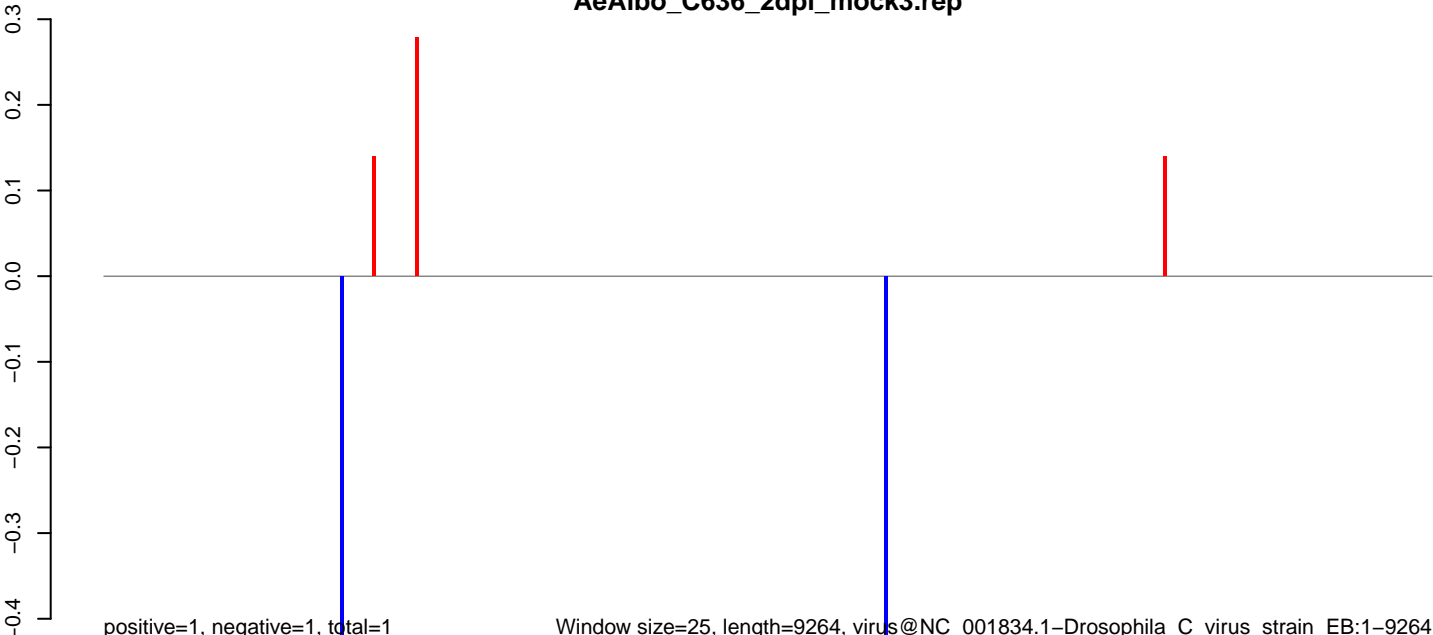
positive=1, negative=1, total=1

AeAlbo_C636_2dpi_mock3.24_35.rep



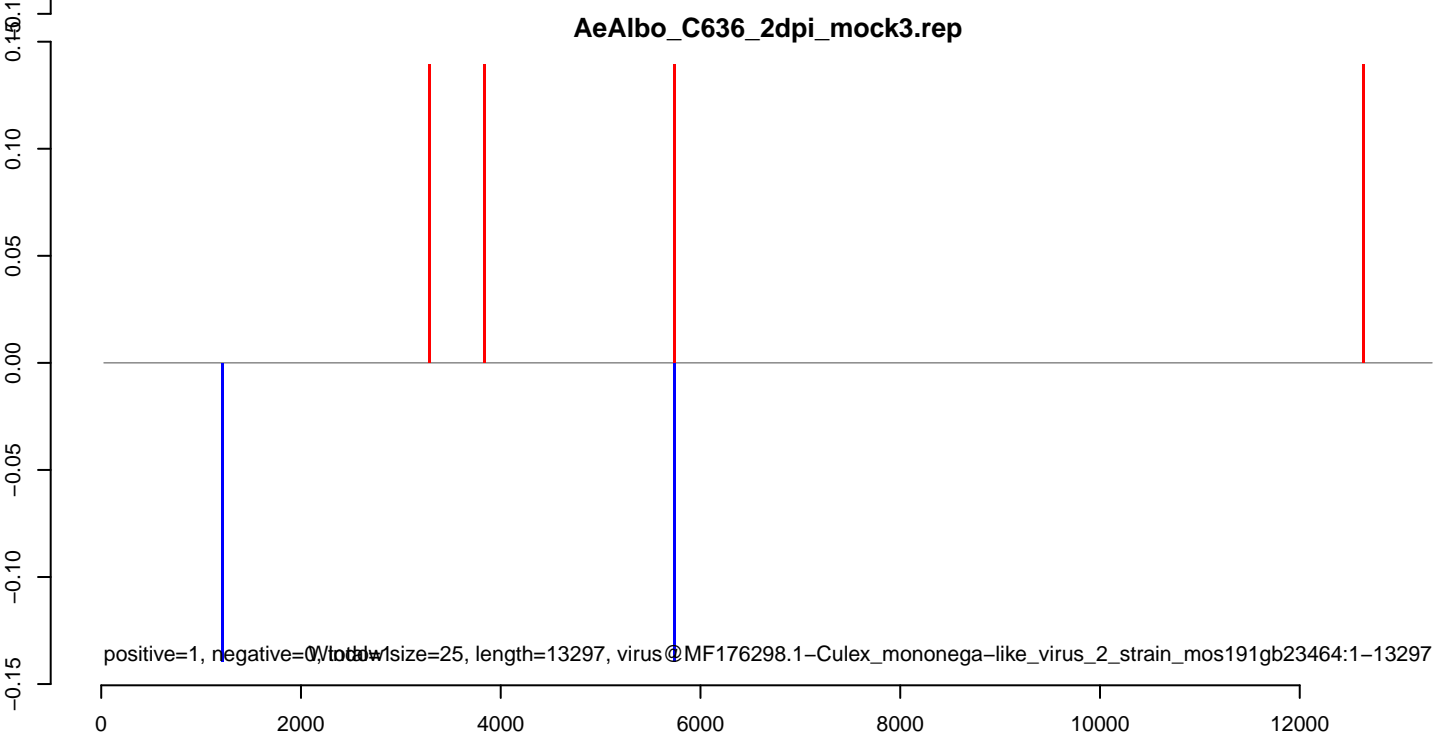
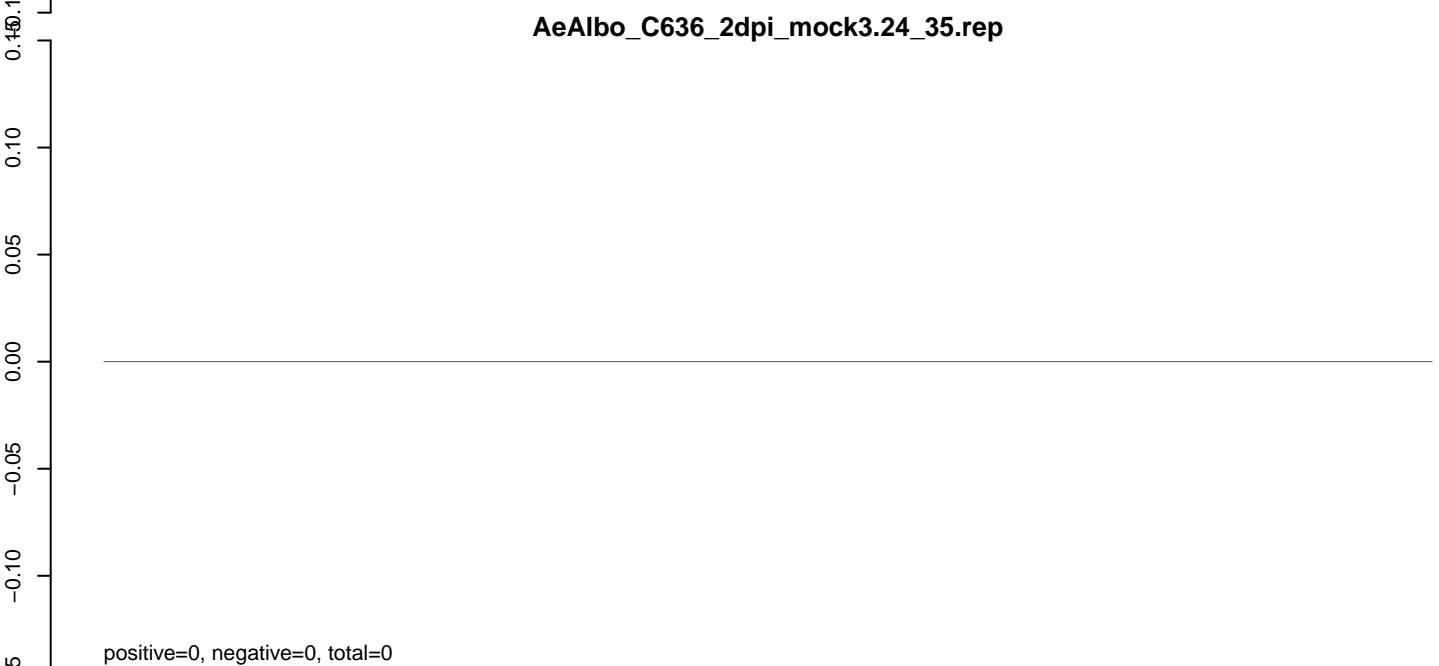
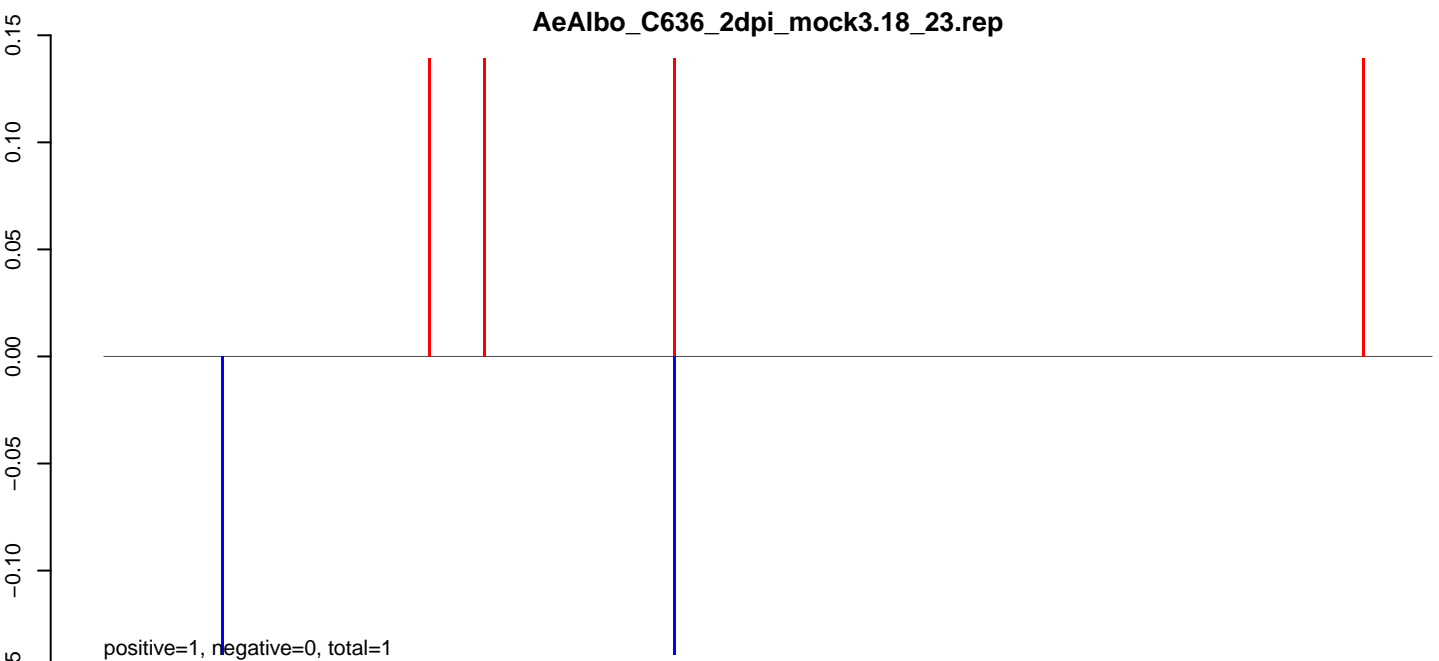
positive=0, negative=0, total=0

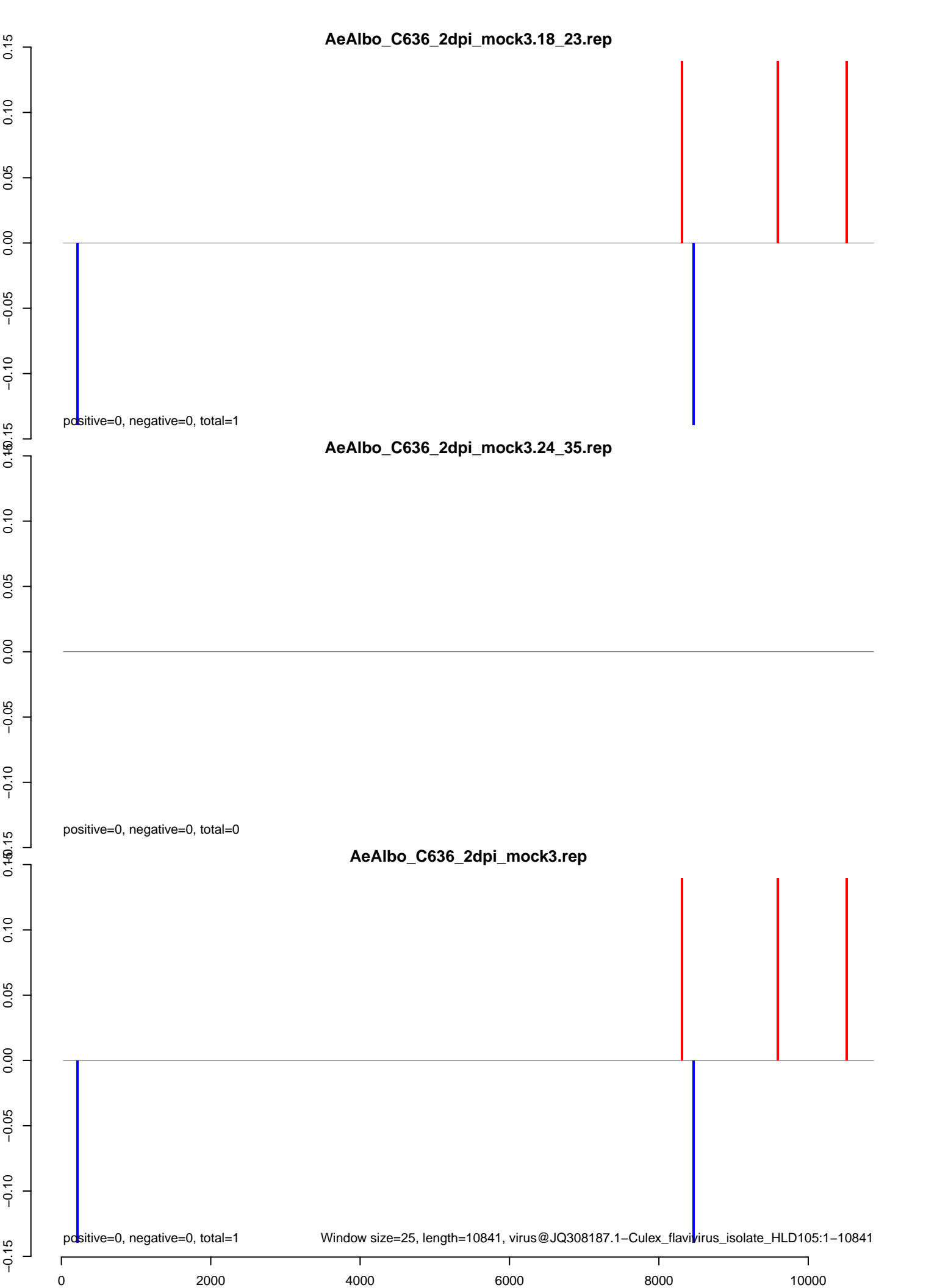
AeAlbo_C636_2dpi_mock3.rep

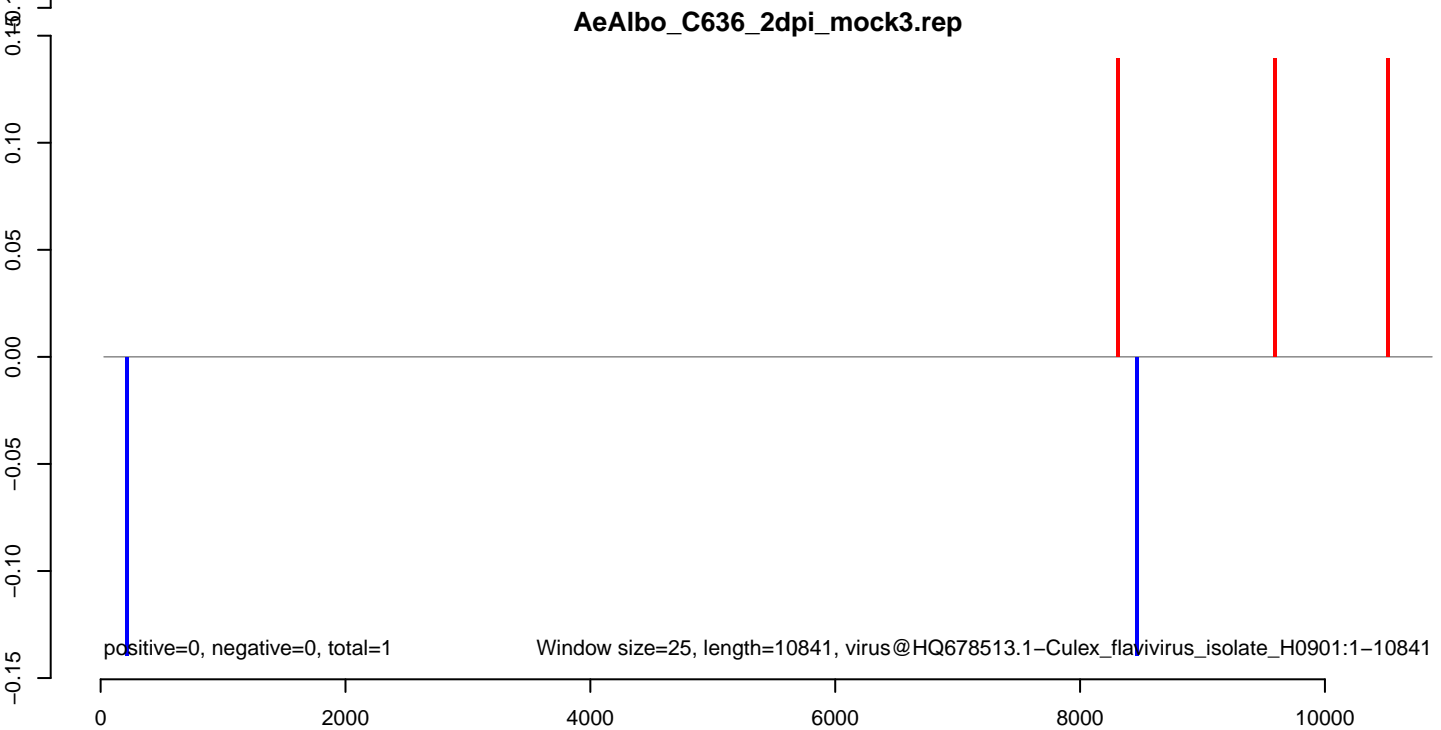
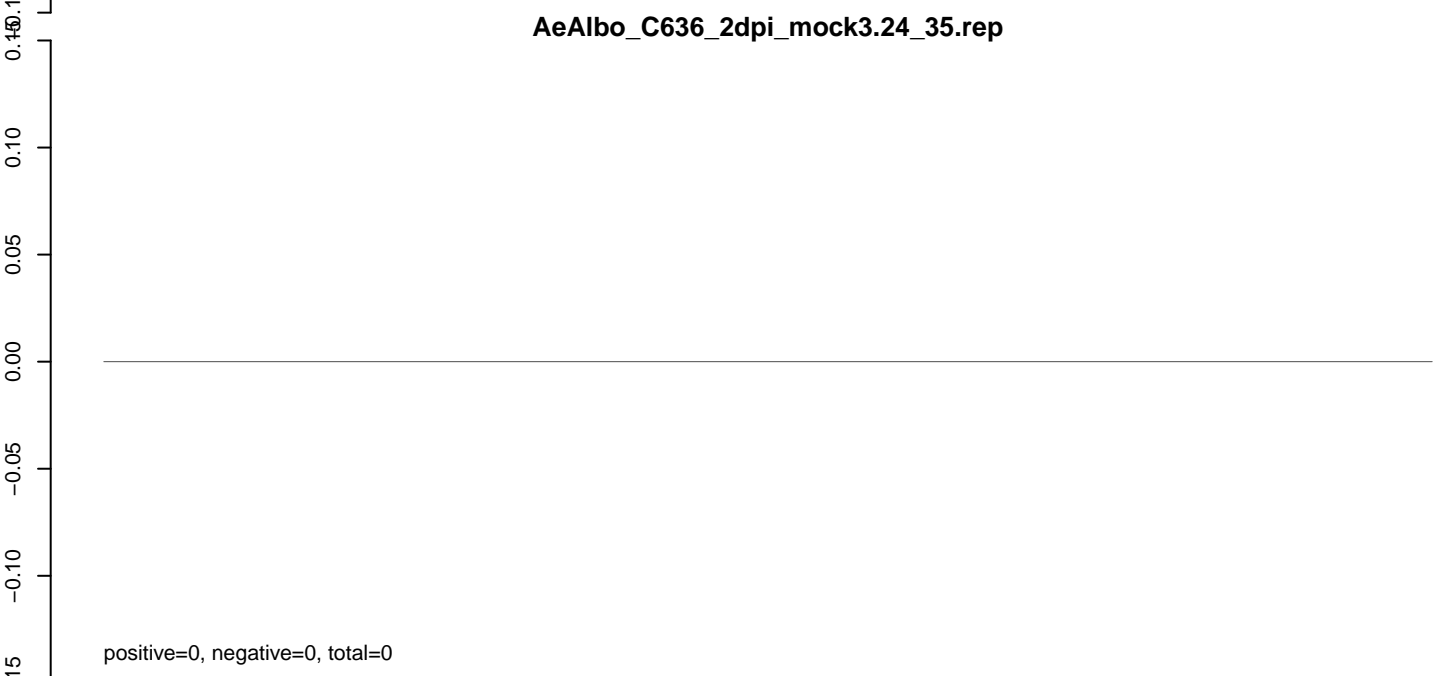
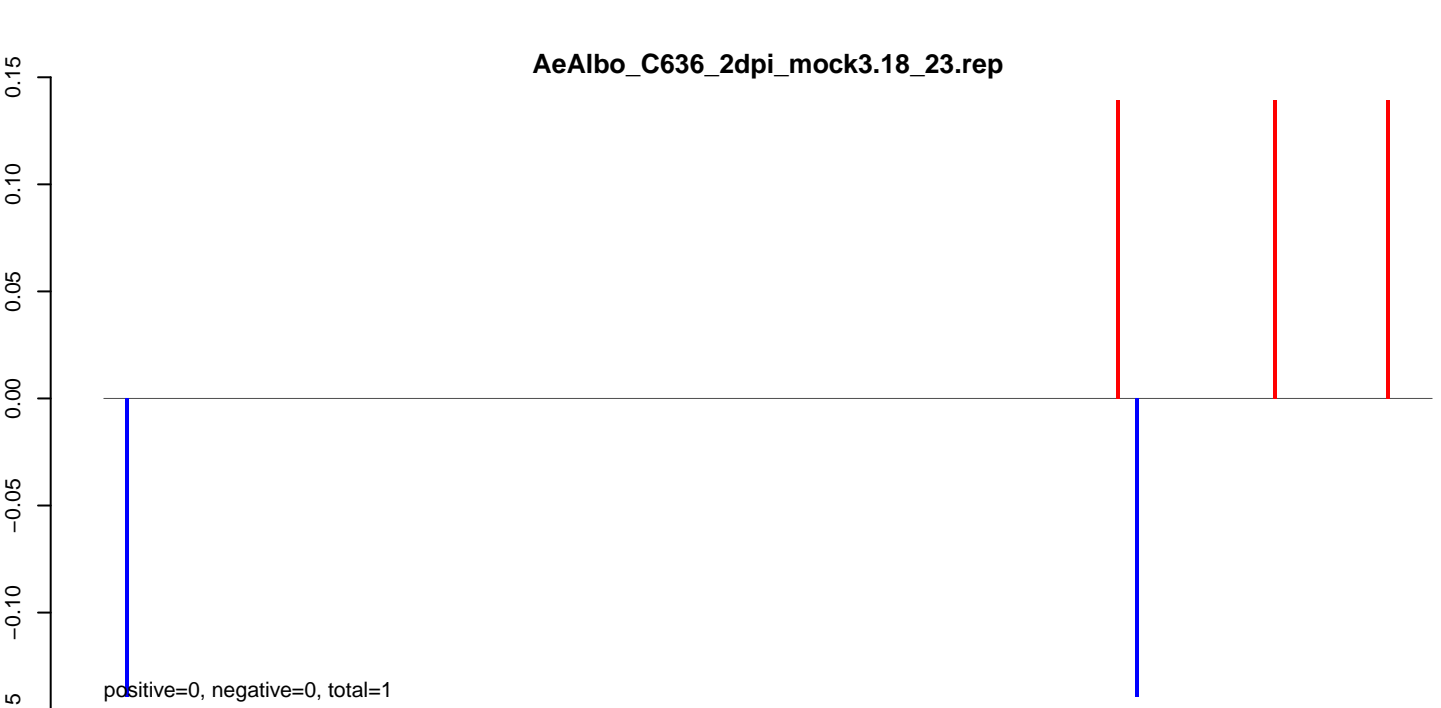


positive=1, negative=1, total=1

Window size=25, length=9264, virus@NC_001834.1-Drosophila_C_virus_strain_EB:1-9264



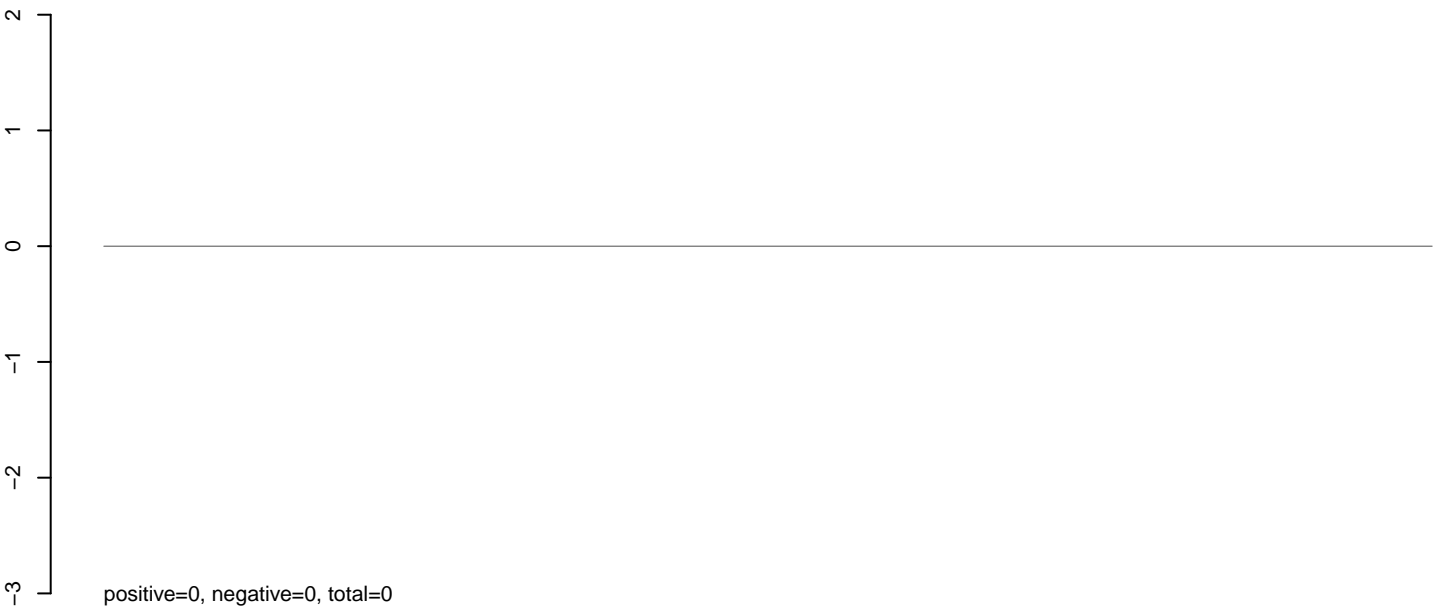




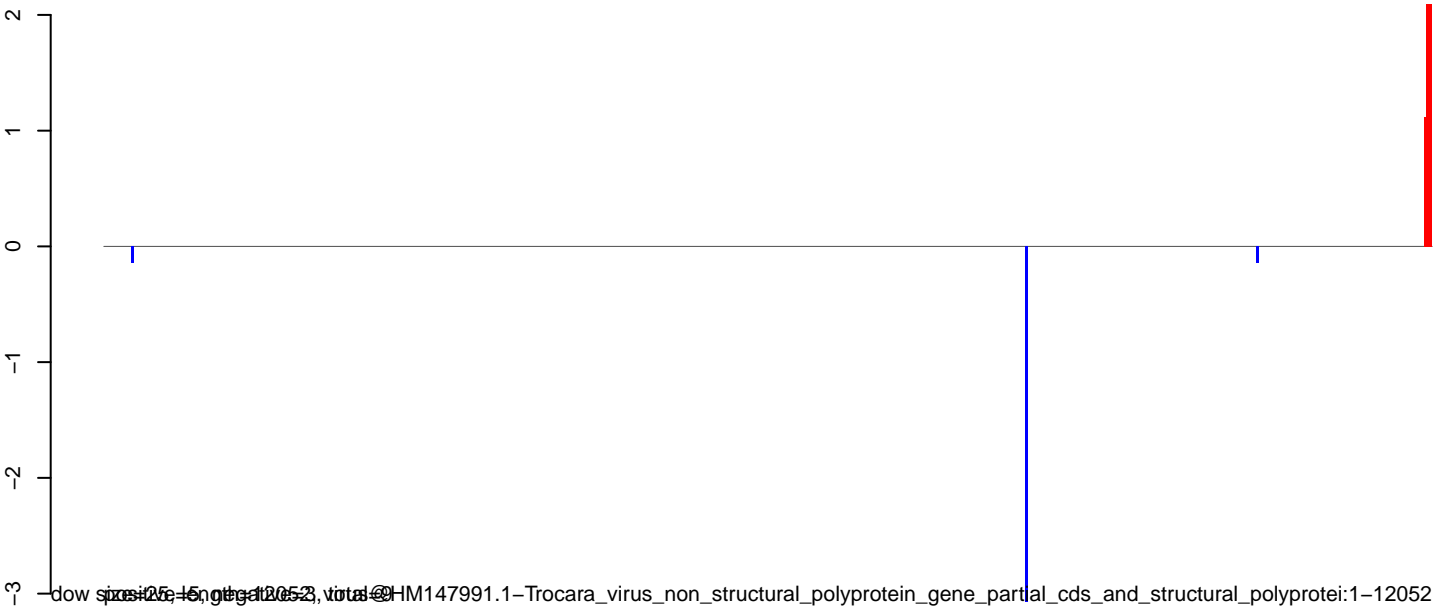
AeAlbo_C636_2dpi_mock3.18_23.rep

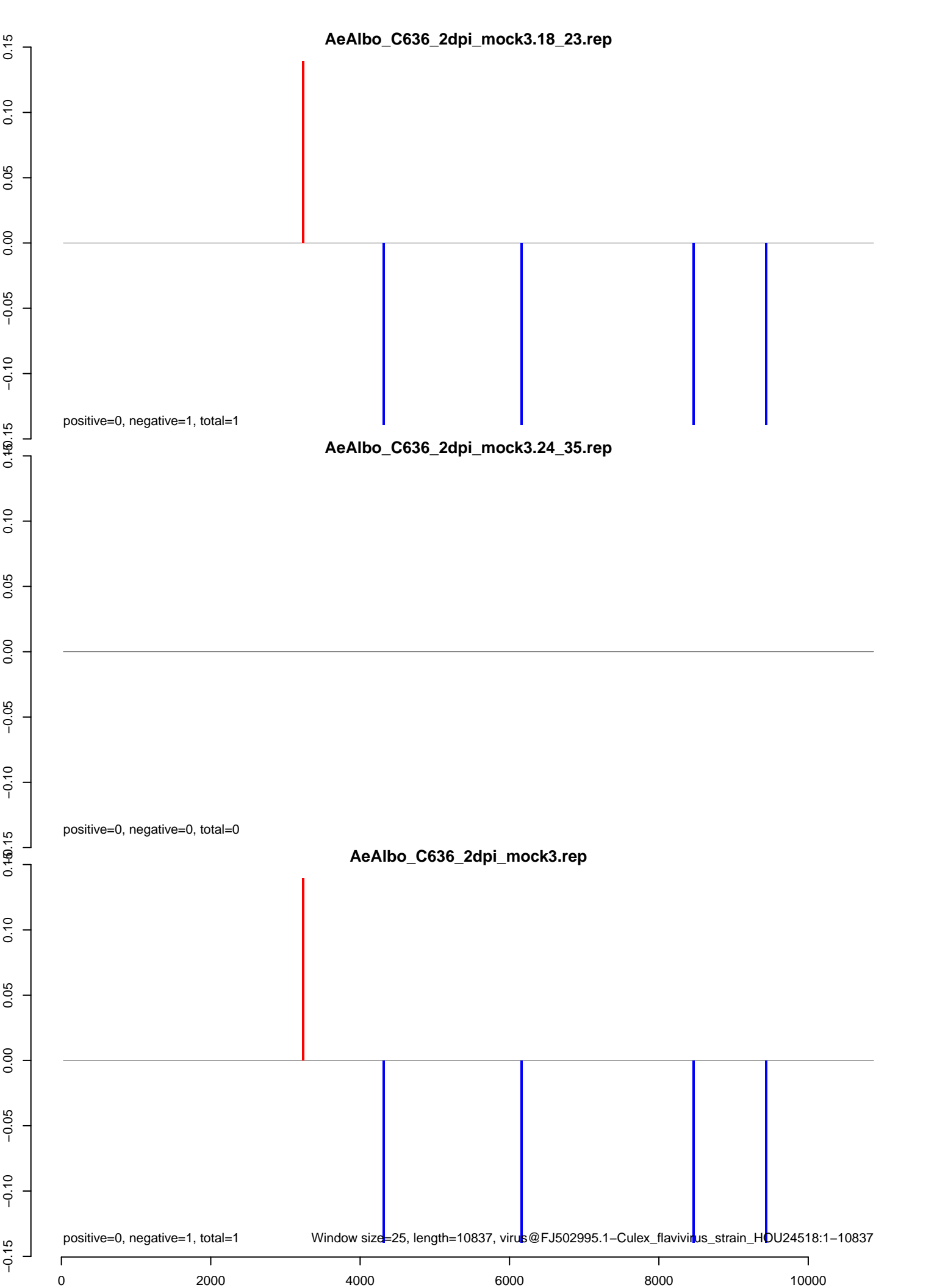


AeAlbo_C636_2dpi_mock3.24_35.rep

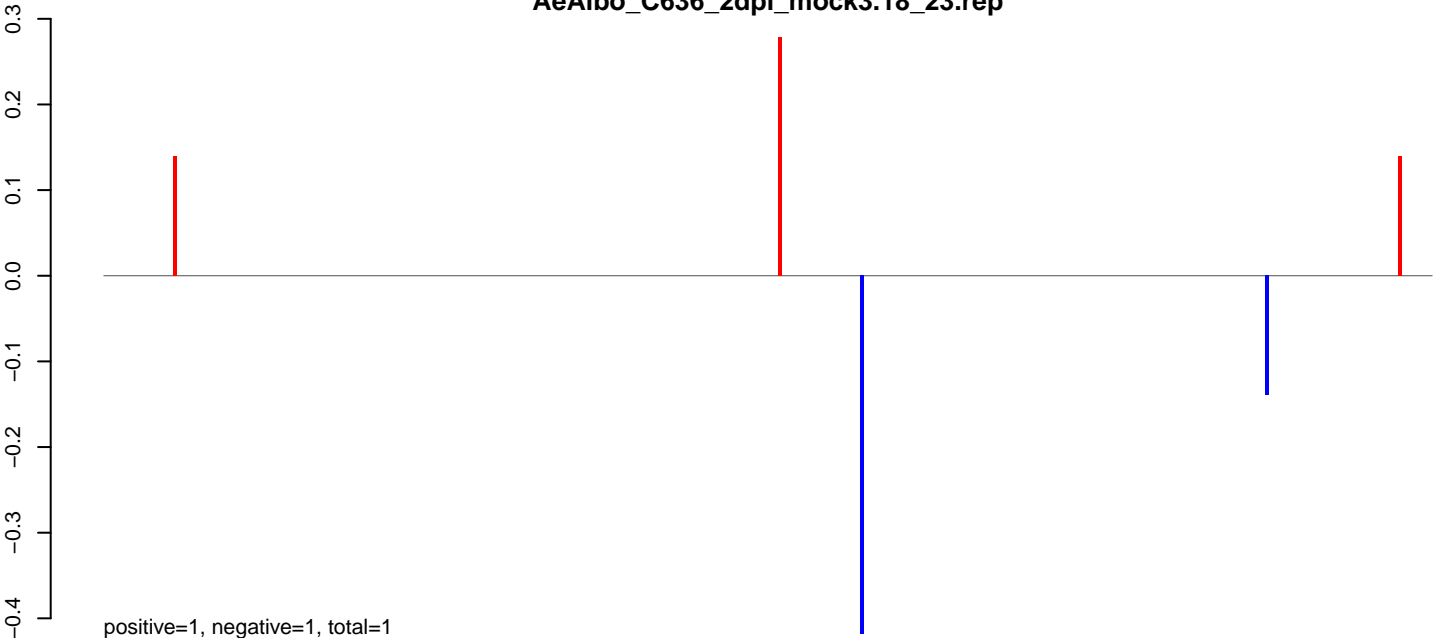


AeAlbo_C636_2dpi_mock3.rep

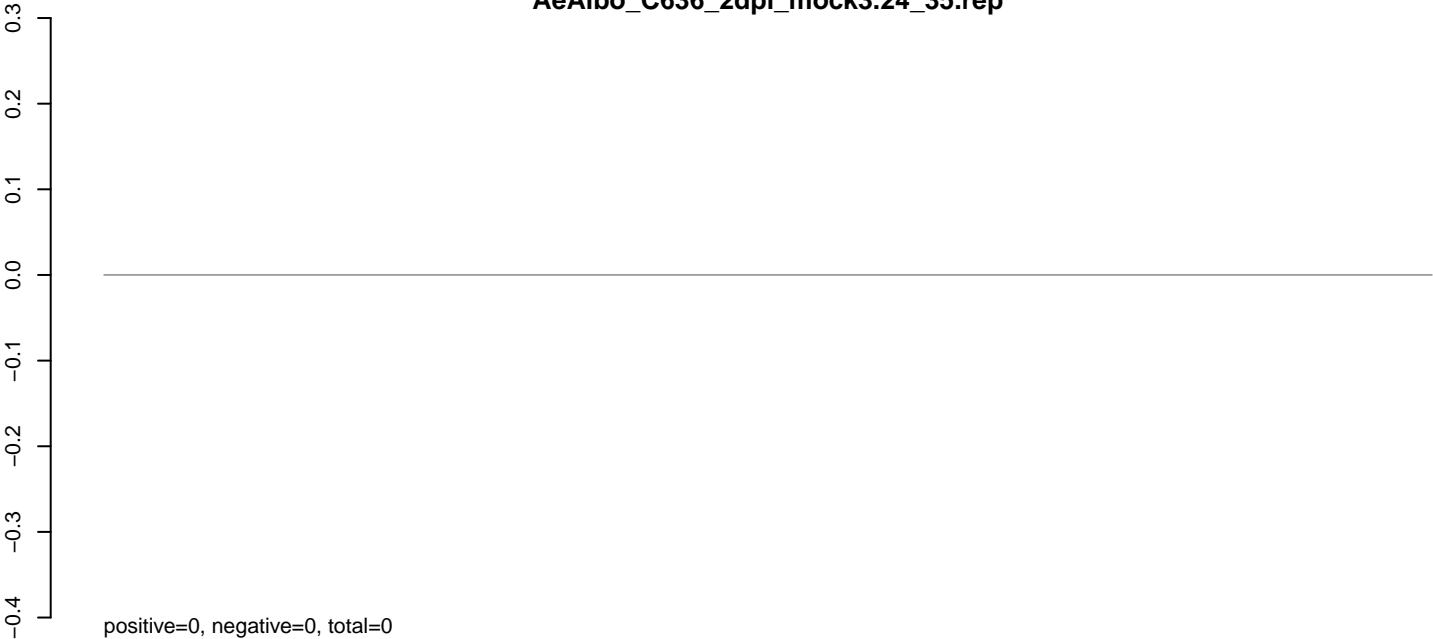




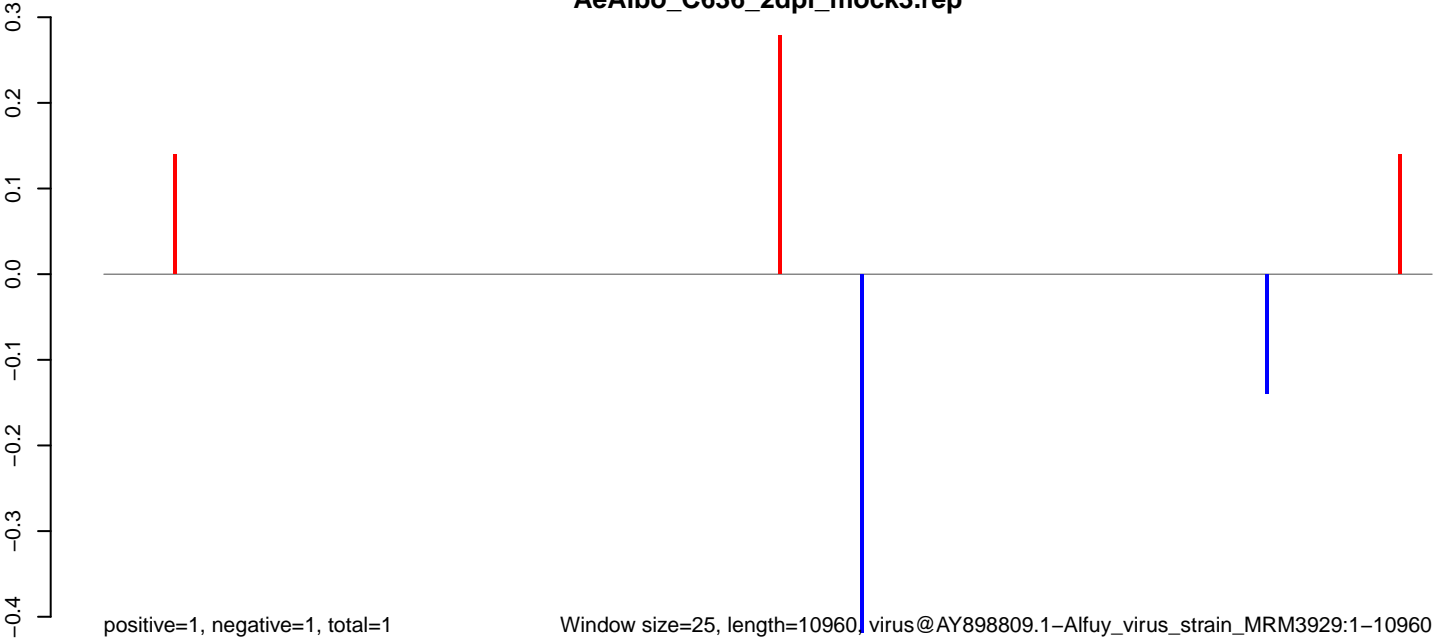
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



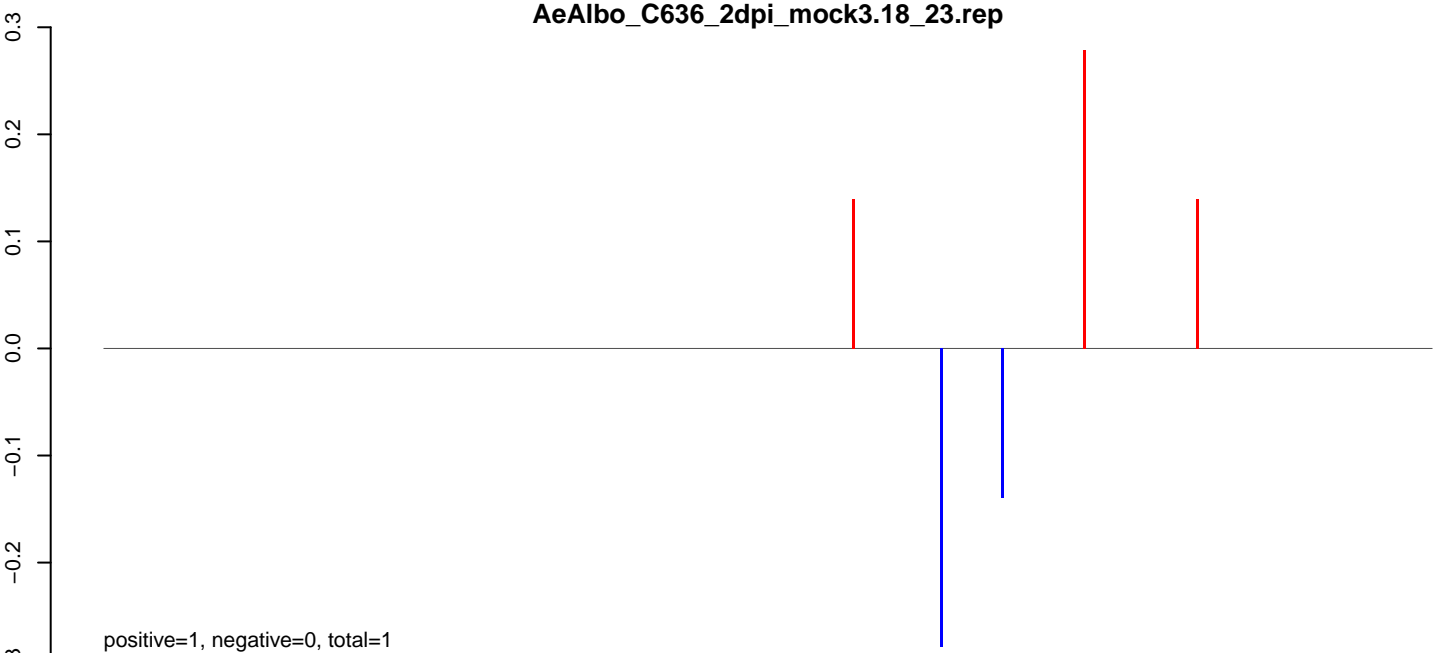
AeAlbo_C636_2dpi_mock3.rep



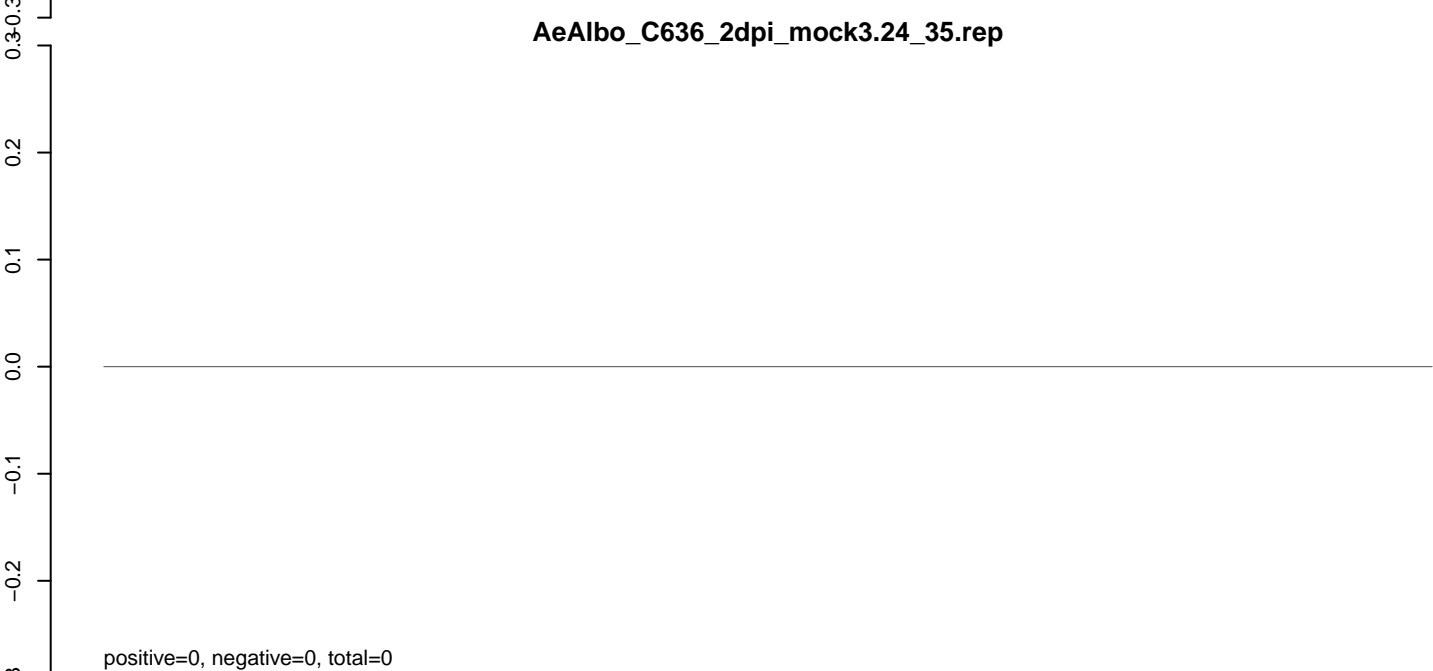
Window size=25, length=10960, virus@AY898809.1-Alfuy_virus_strain_MRM3929:1-10960

0 2000 4000 6000 8000 10000

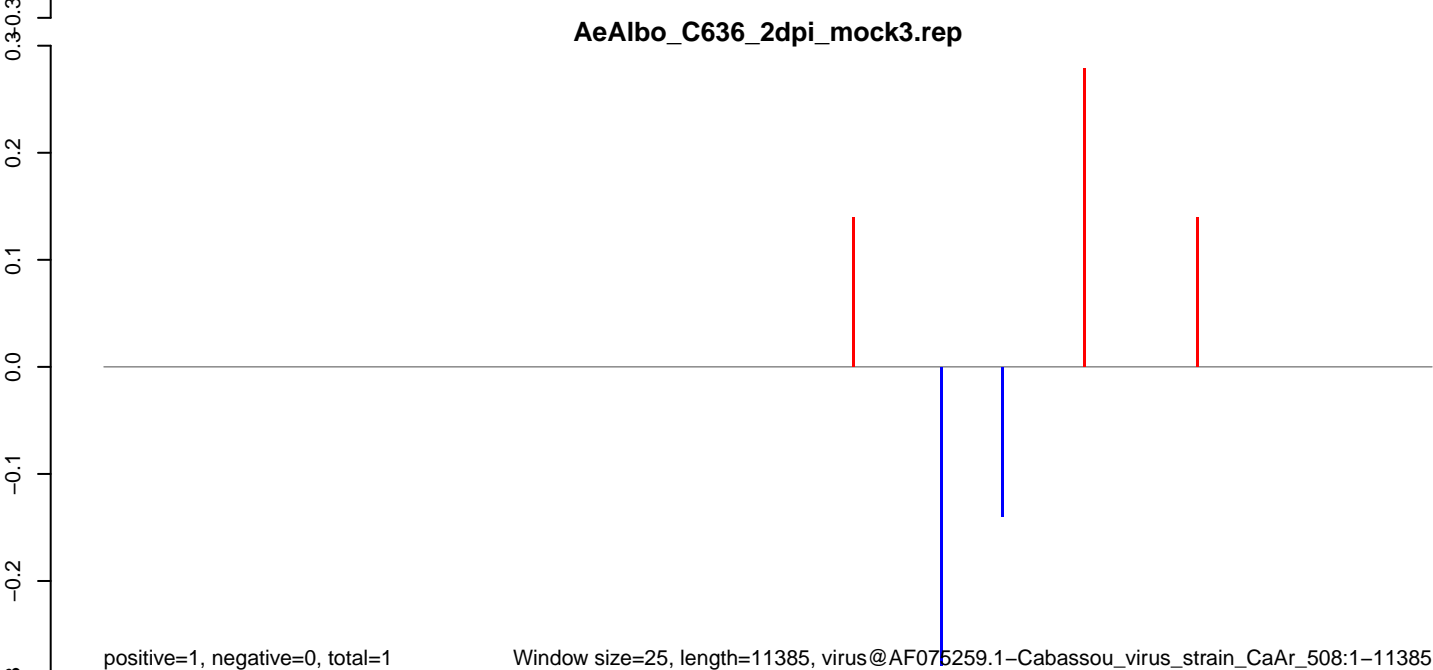
AeAlbo_C636_2dpi_mock3.18_23.rep



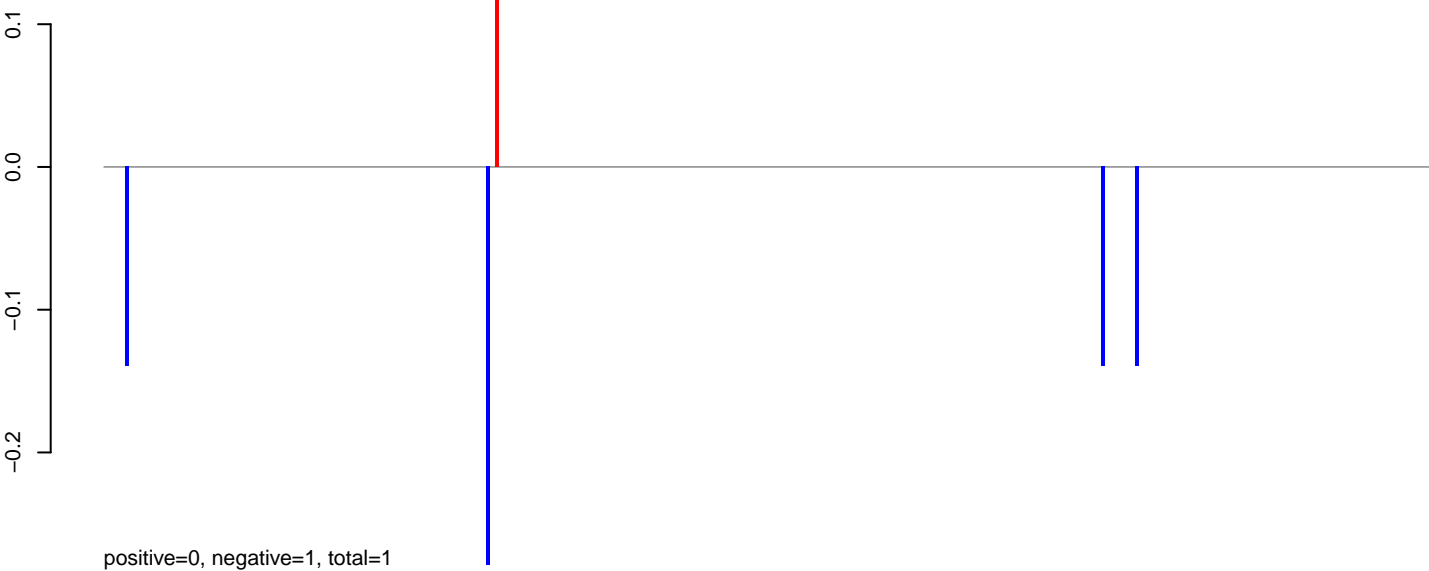
AeAlbo_C636_2dpi_mock3.24_35.rep



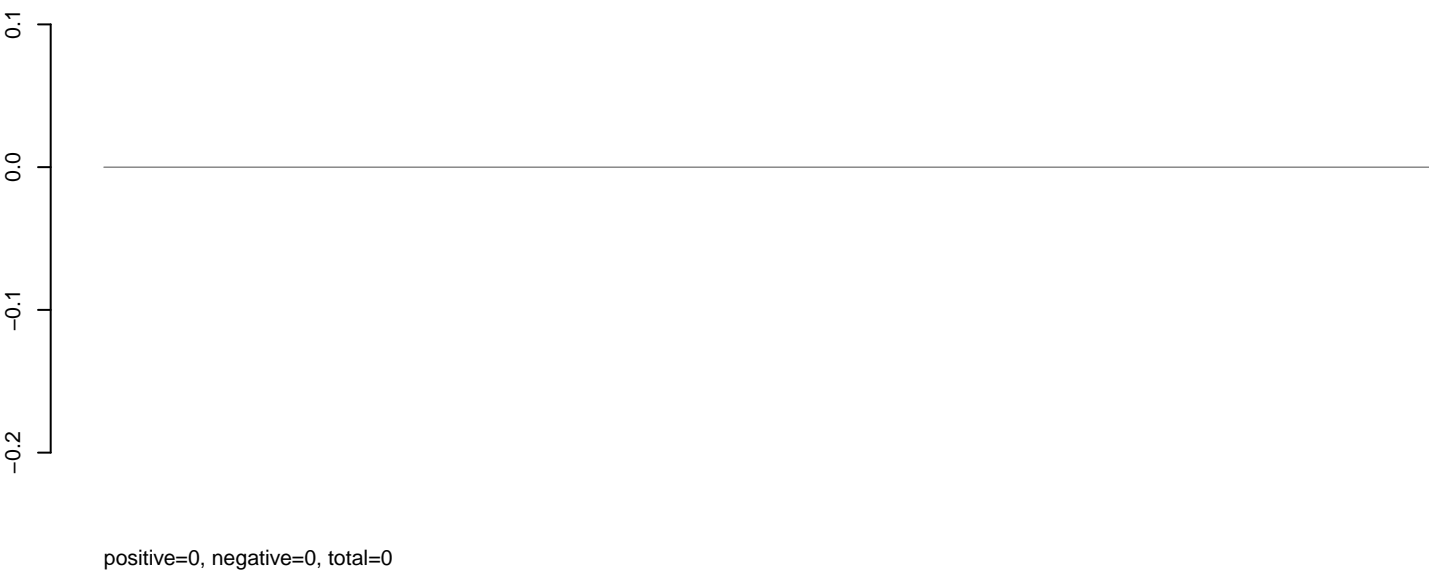
AeAlbo_C636_2dpi_mock3.rep



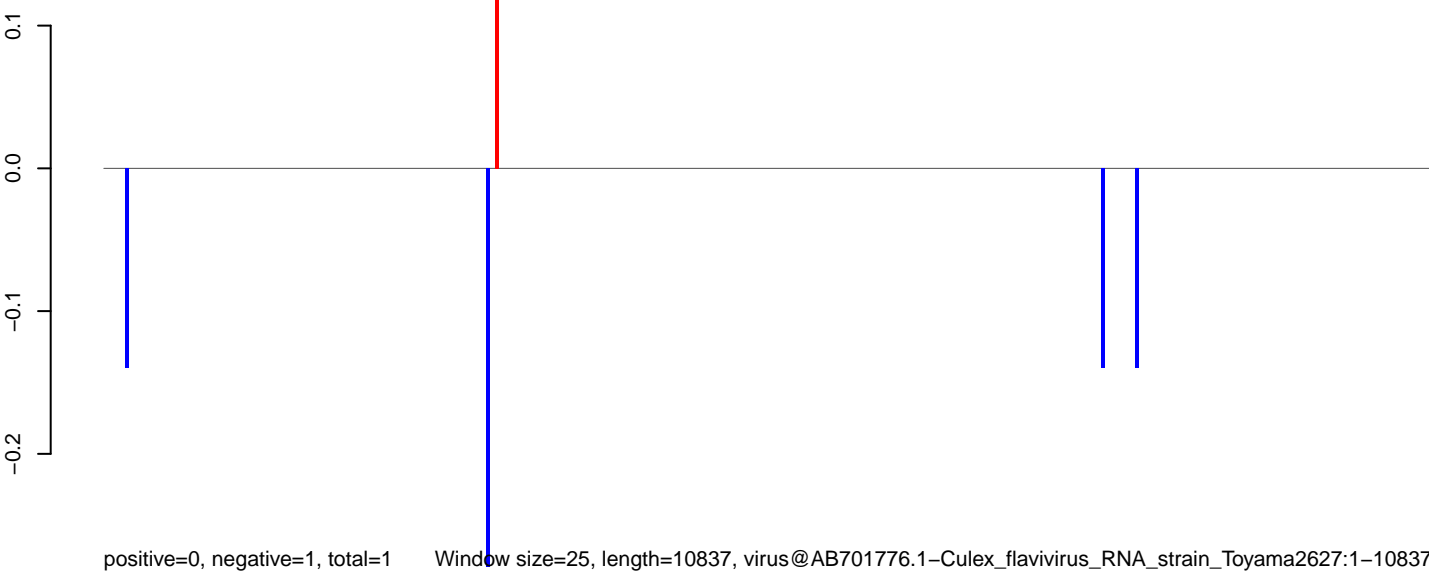
AeAlbo_C636_2dpi_mock3.18_23.rep



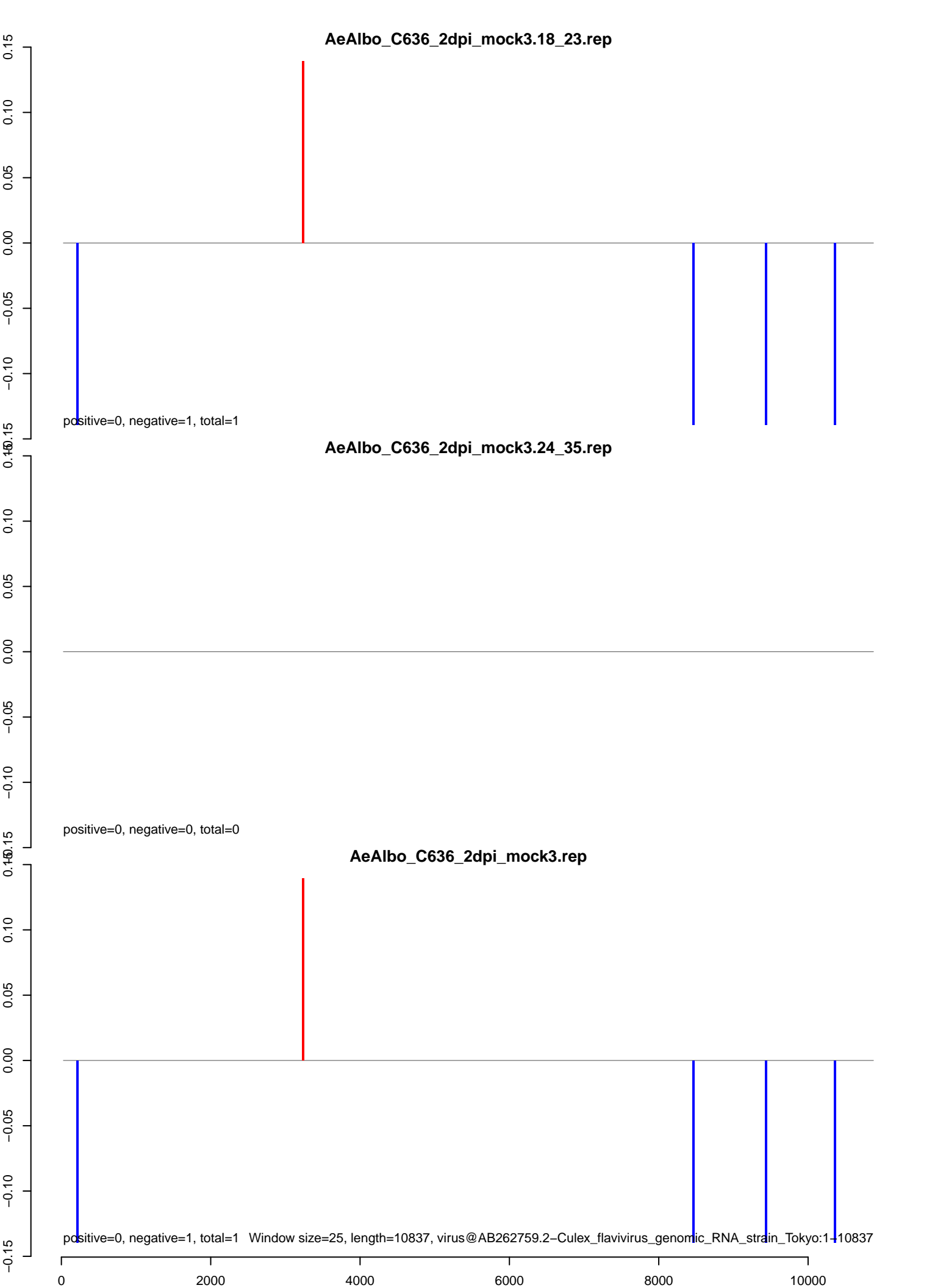
AeAlbo_C636_2dpi_mock3.24_35.rep



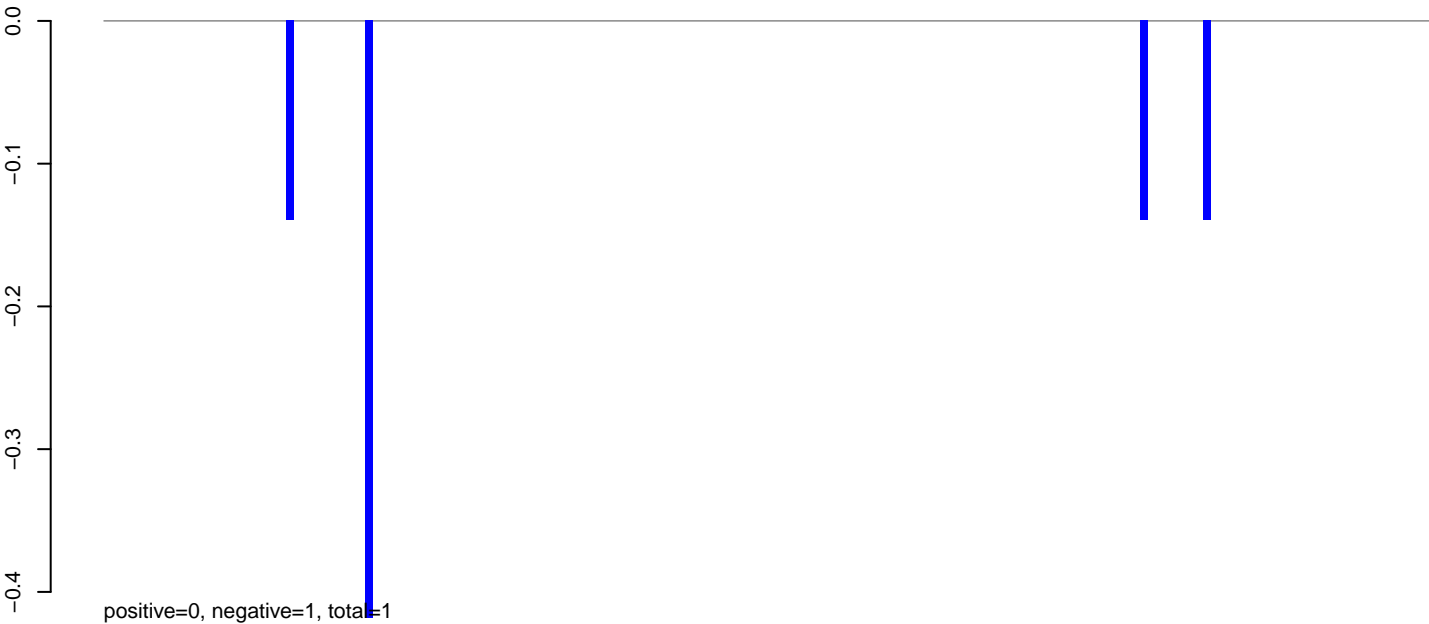
AeAlbo_C636_2dpi_mock3.rep



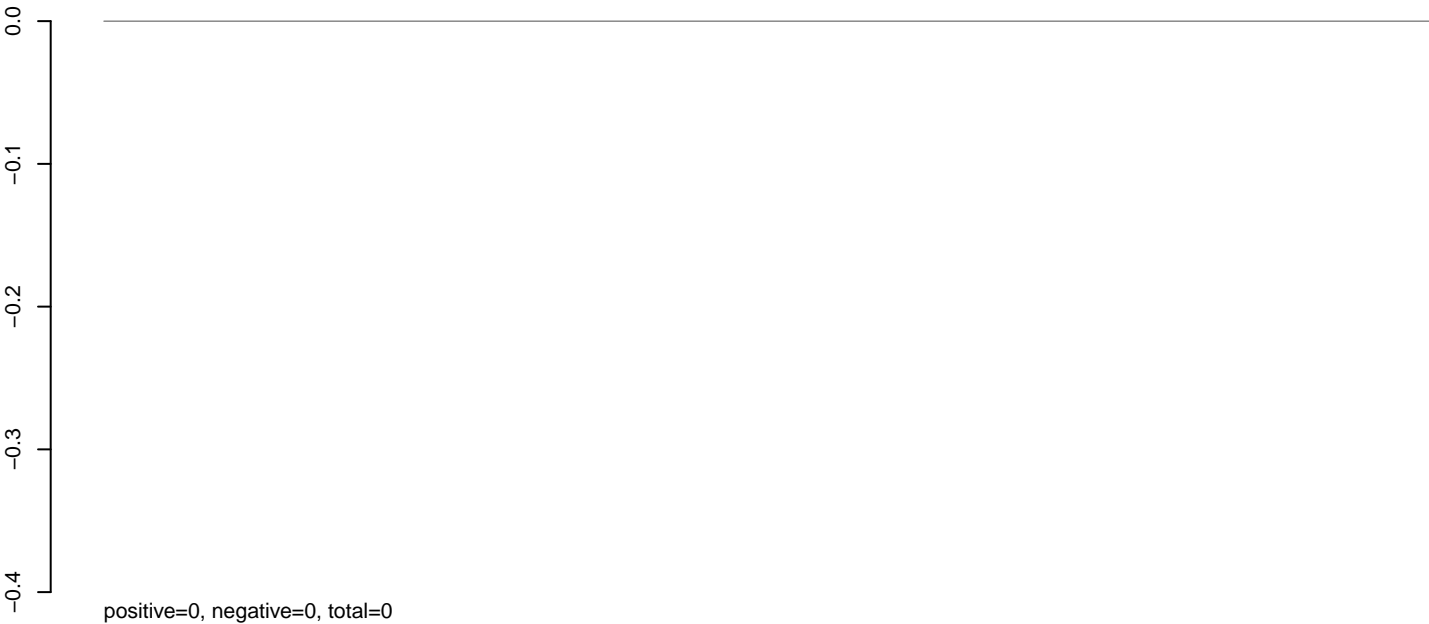
0 2000 4000 6000 8000 10000



AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

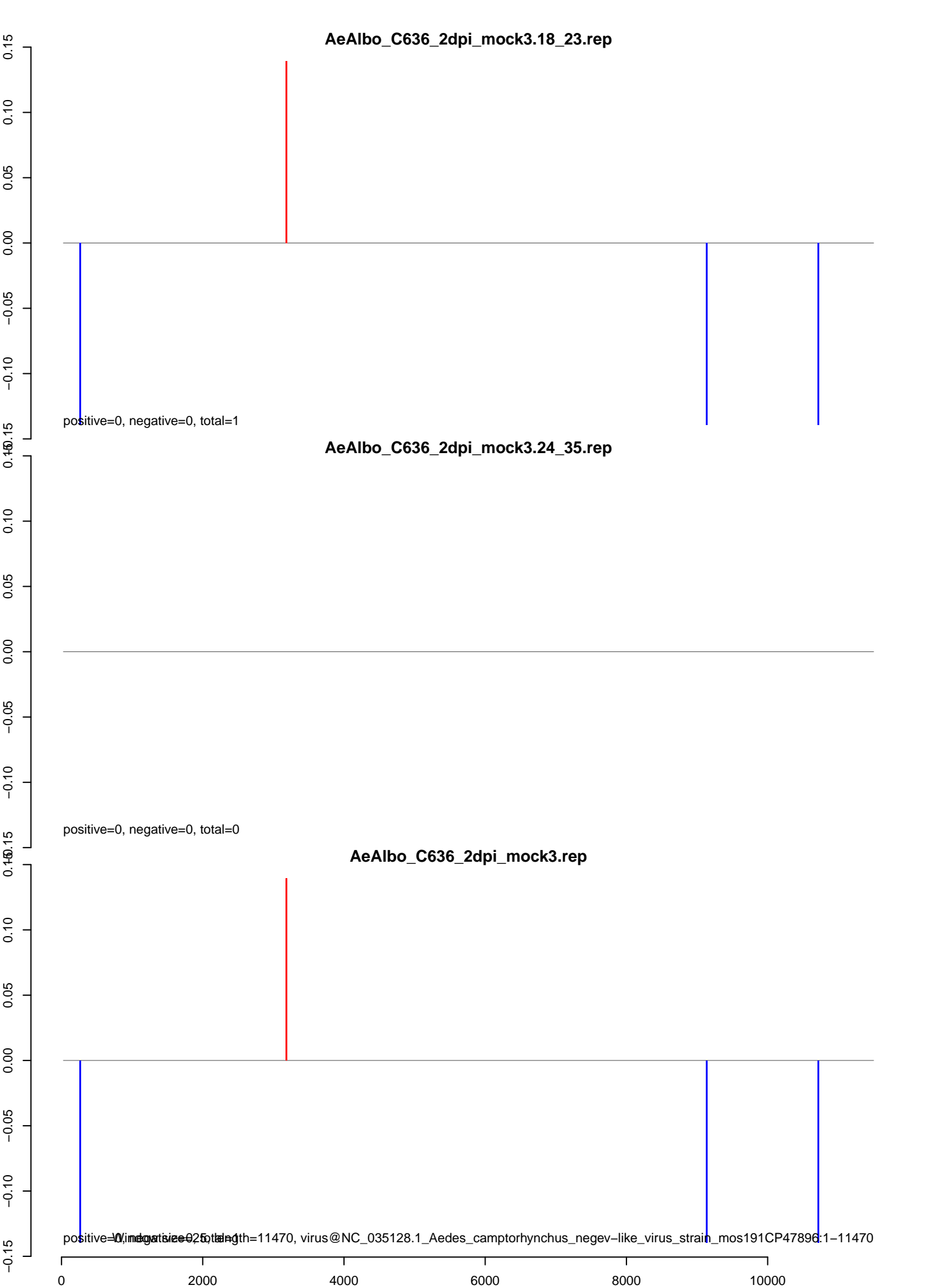


AeAlbo_C636_2dpi_mock3.rep

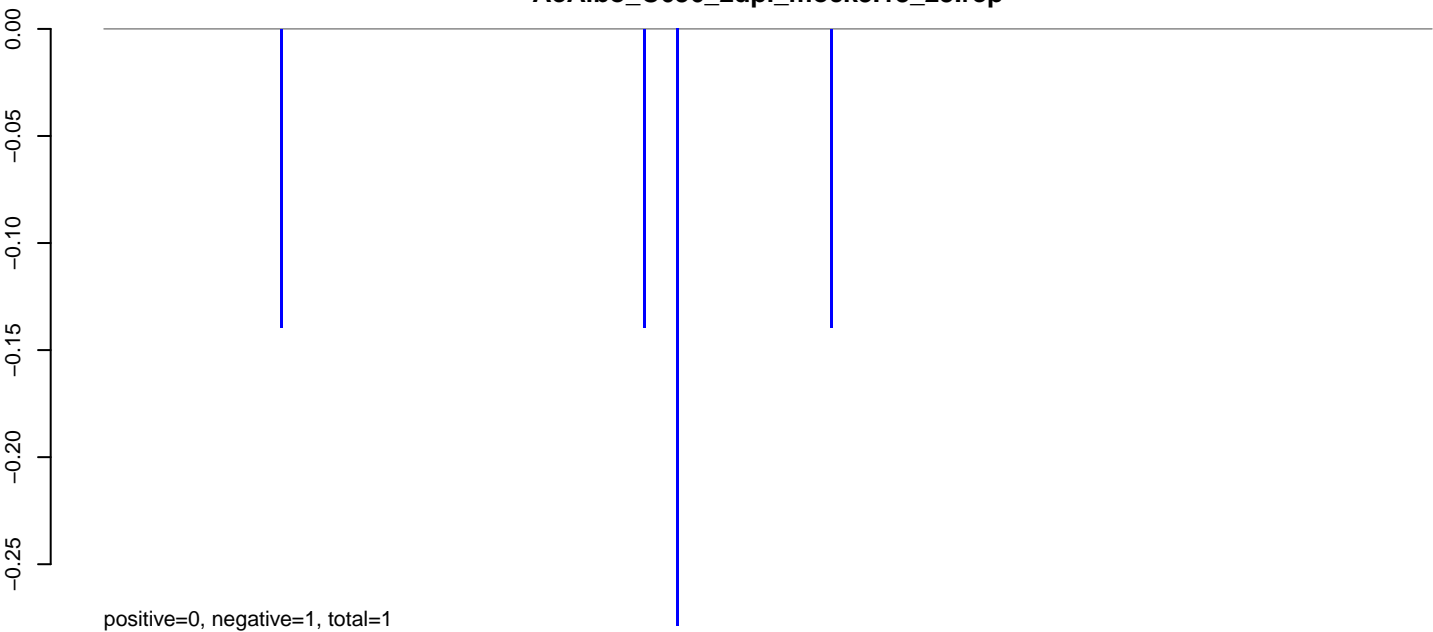


Window size=25, length=4176, virus@X74945.1-Aedes_albopictus_densovirus_2_DNA:1-4176

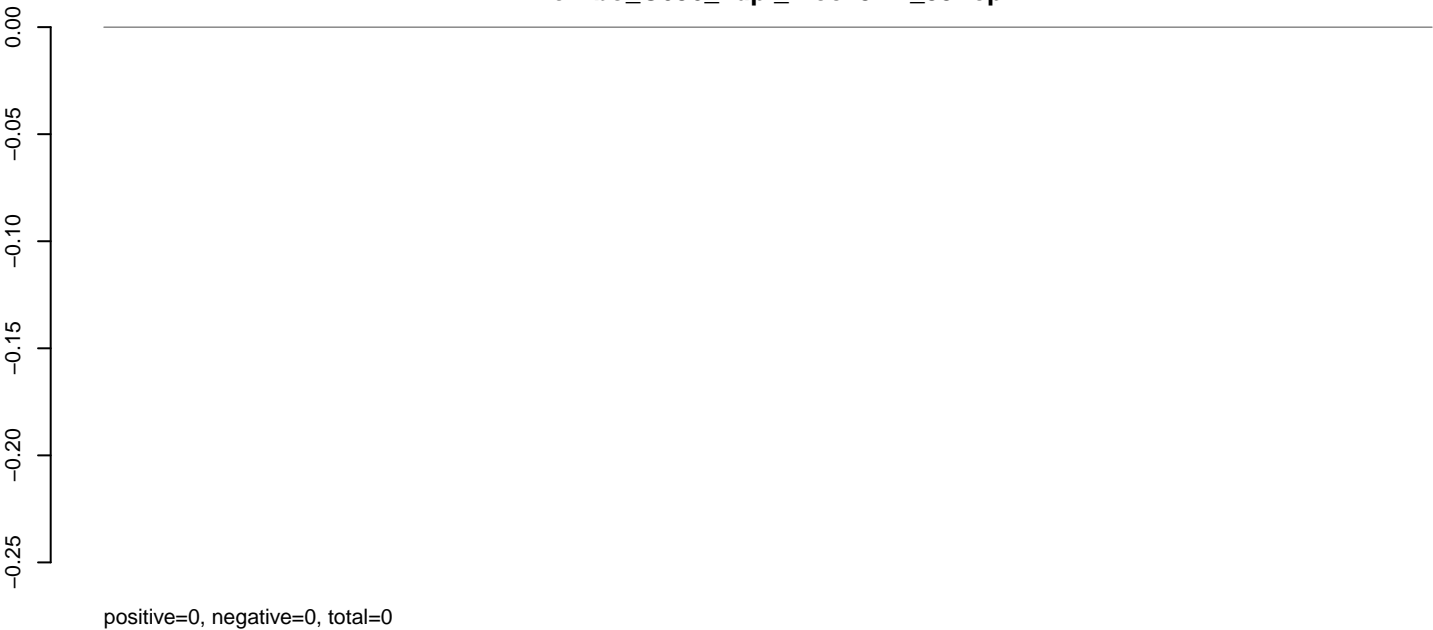
0 1000 2000 3000 4000



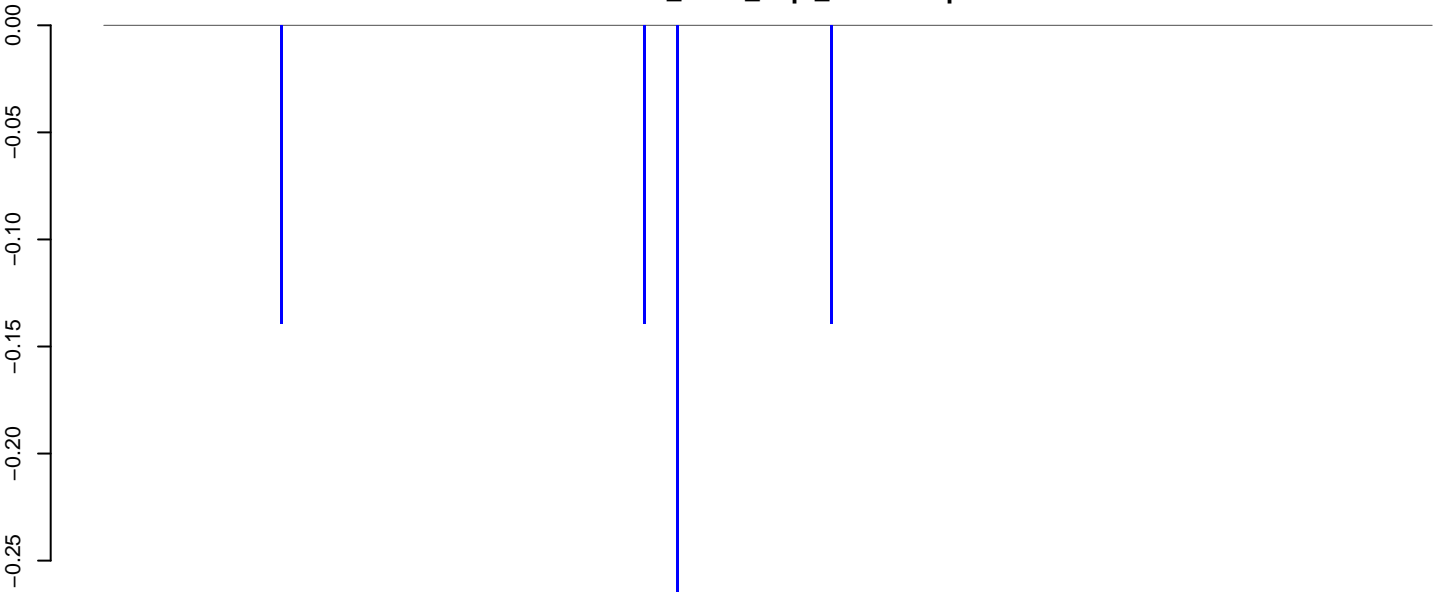
AeAlbo_C636_2dpi_mock3.18_23.rep



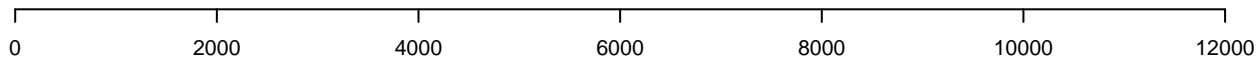
AeAlbo_C636_2dpi_mock3.24_35.rep



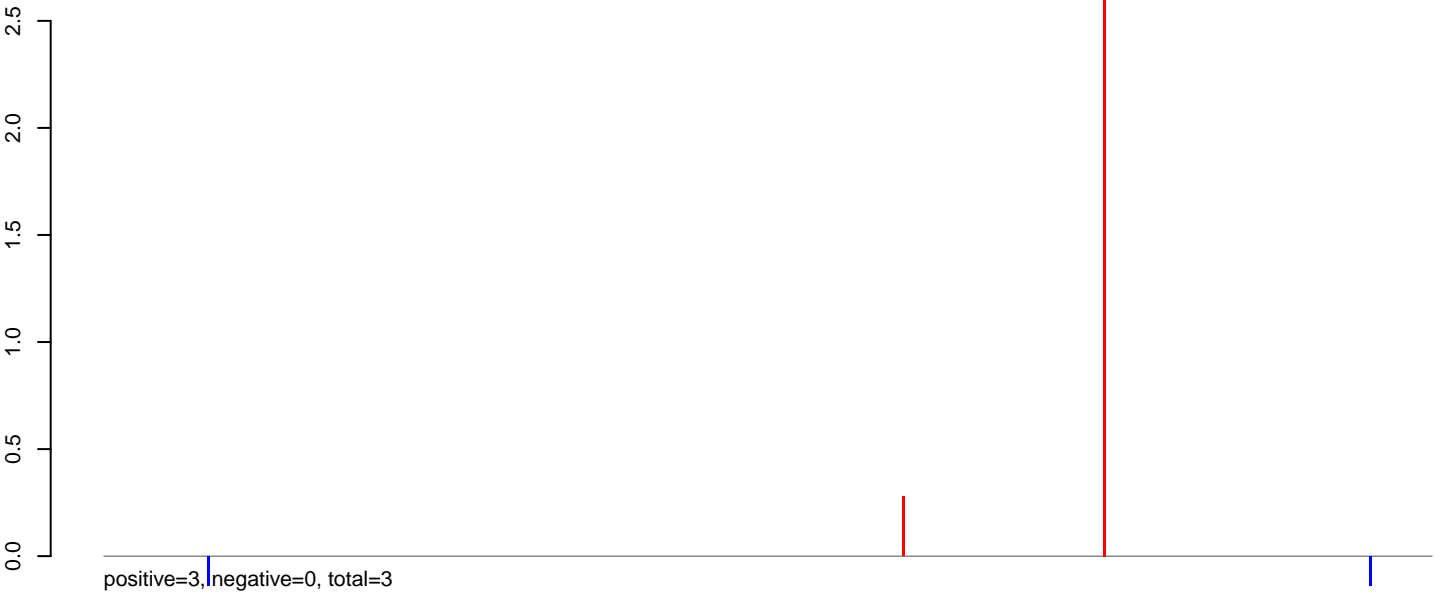
AeAlbo_C636_2dpi_mock3.rep



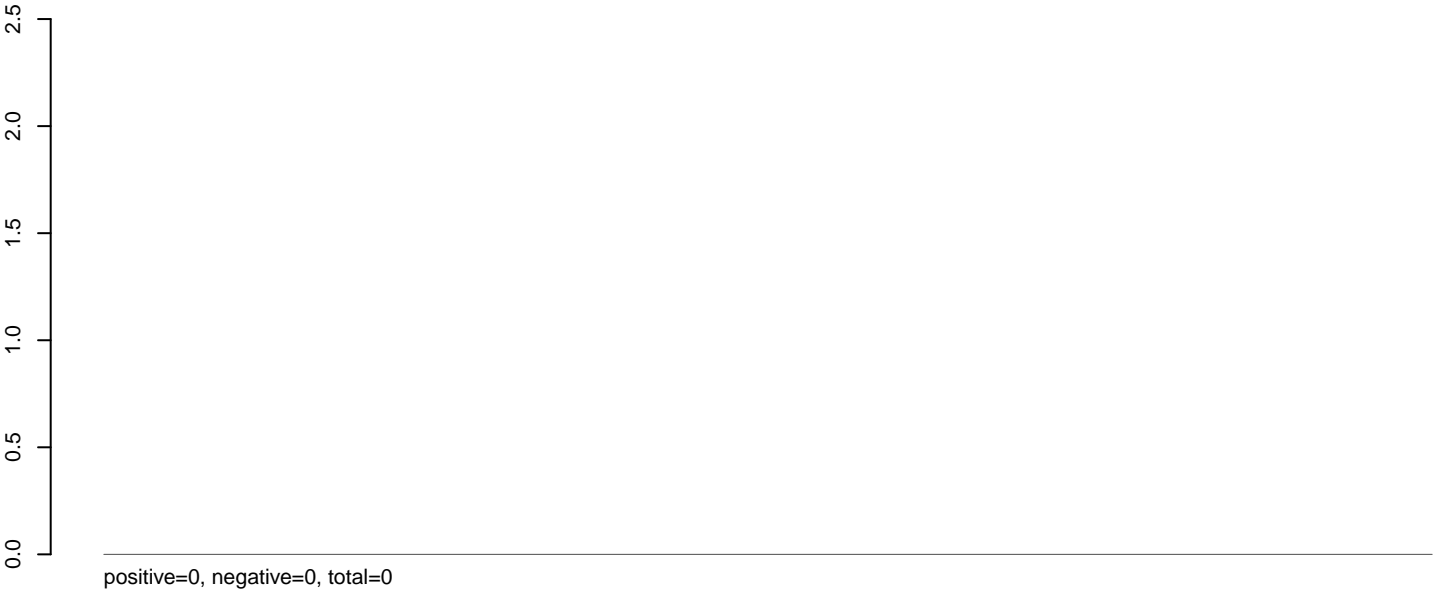
row size=25, col=1, the gaps are 1155, 1105 and 1105. NC_034538.1-Joinjakaka_virus_nucleoprotein_phosphoprotein_hypothetical_protein_matrix_glycopro:1-13155



AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

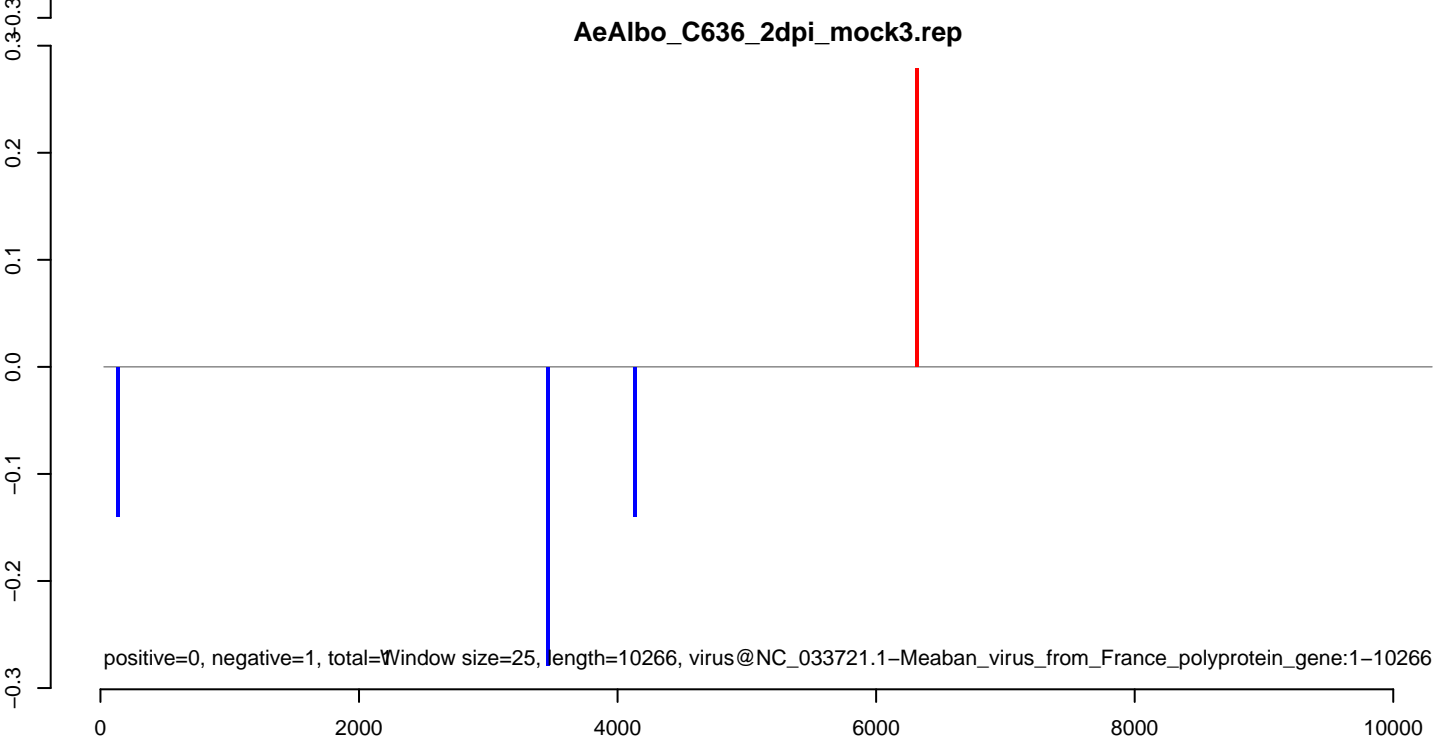
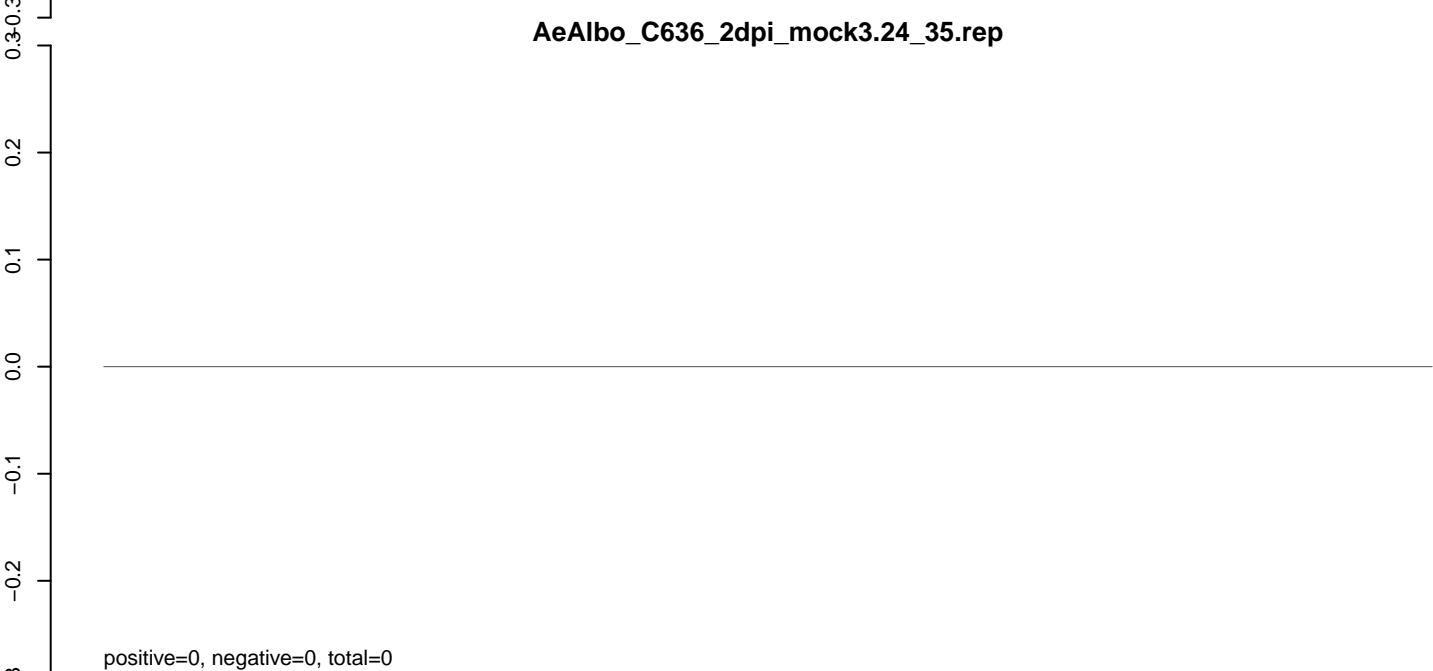
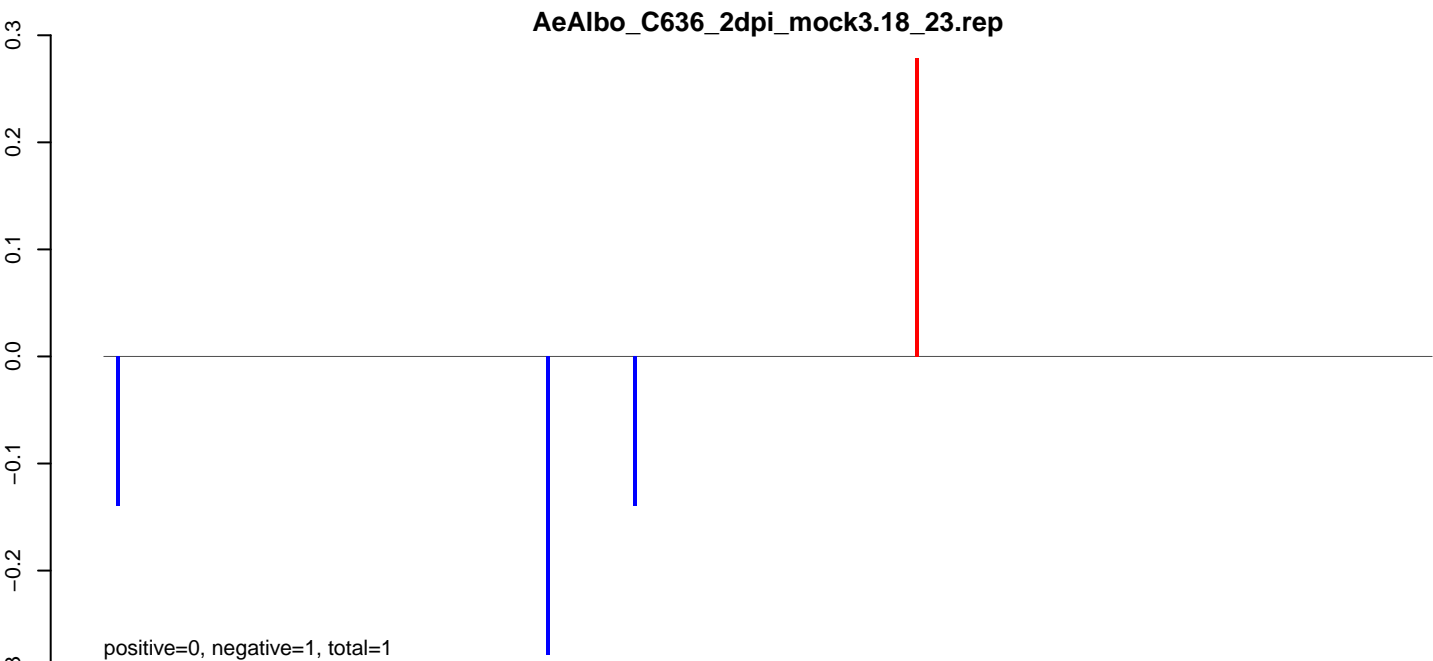


AeAlbo_C636_2dpi_mock3.rep

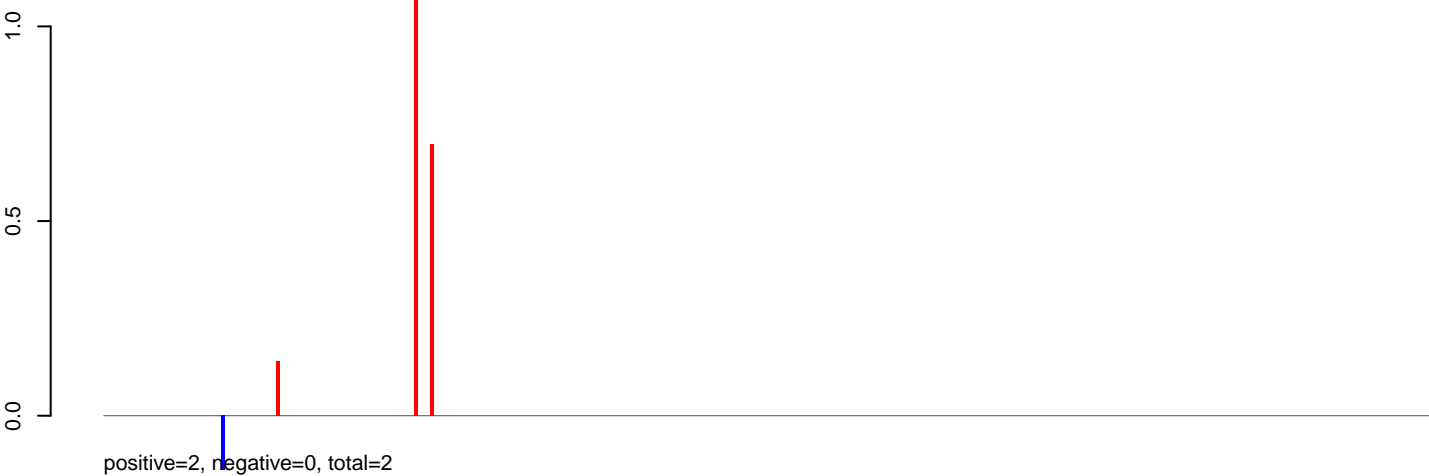


w size=25, v=0, t=0, u=0, use=MS_034450.1-Kamese_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycoprotei:1-13209

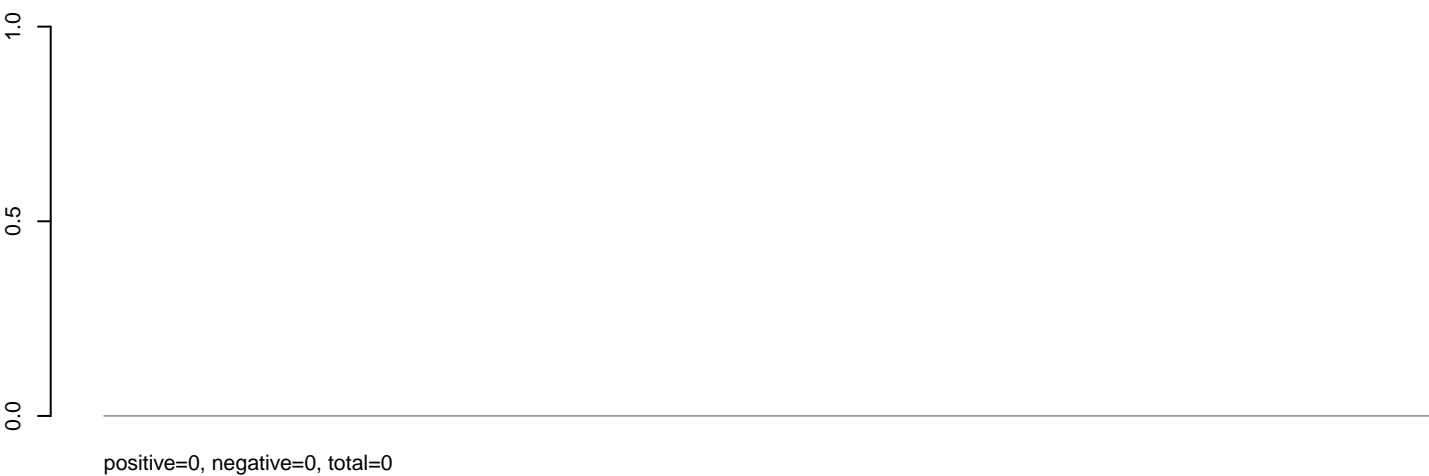
0 2000 4000 6000 8000 10000 12000



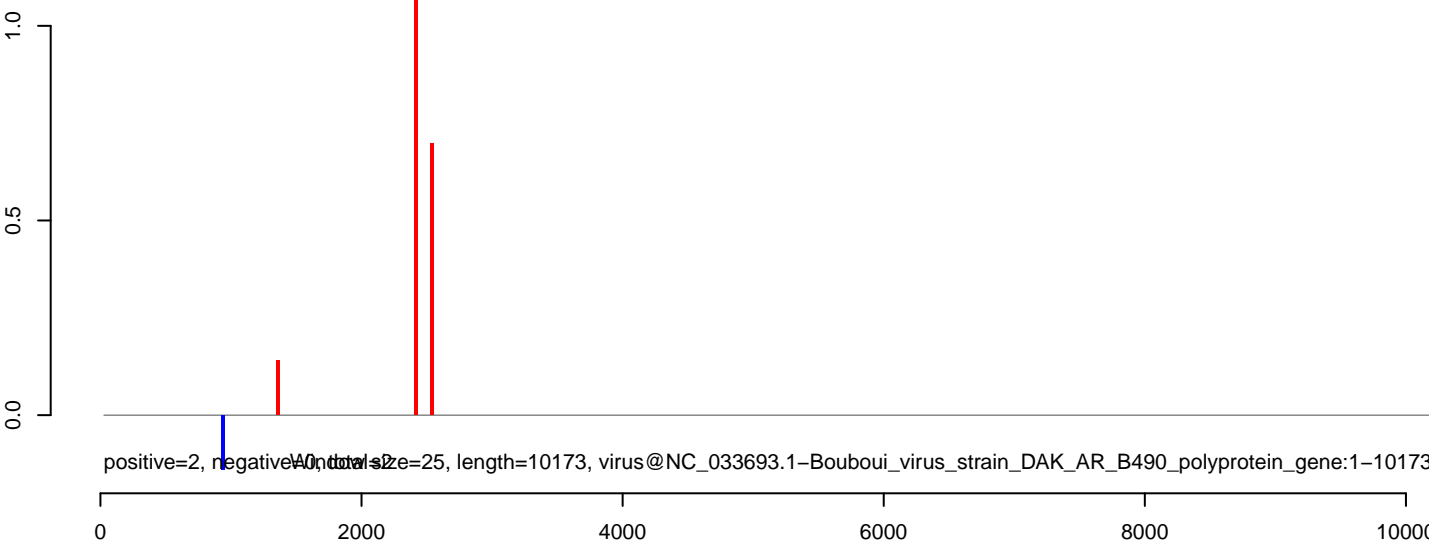
AeAlbo_C636_2dpi_mock3.18_23.rep



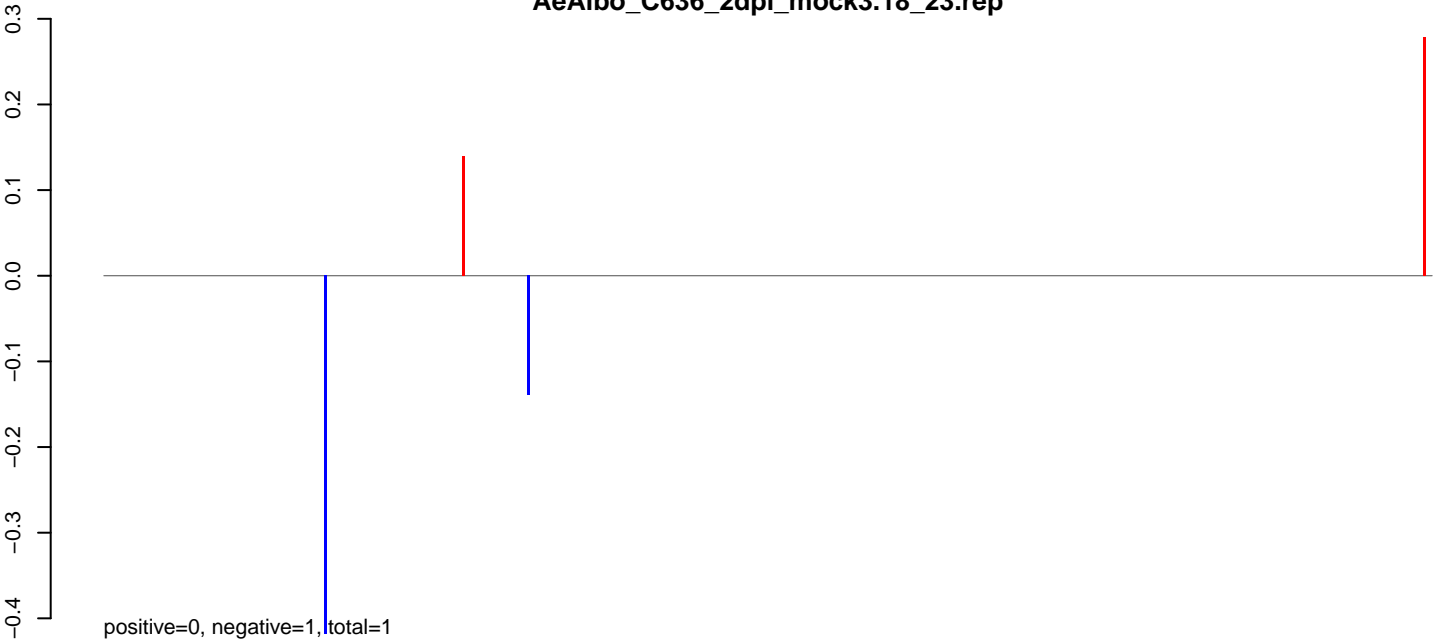
AeAlbo_C636_2dpi_mock3.24_35.rep



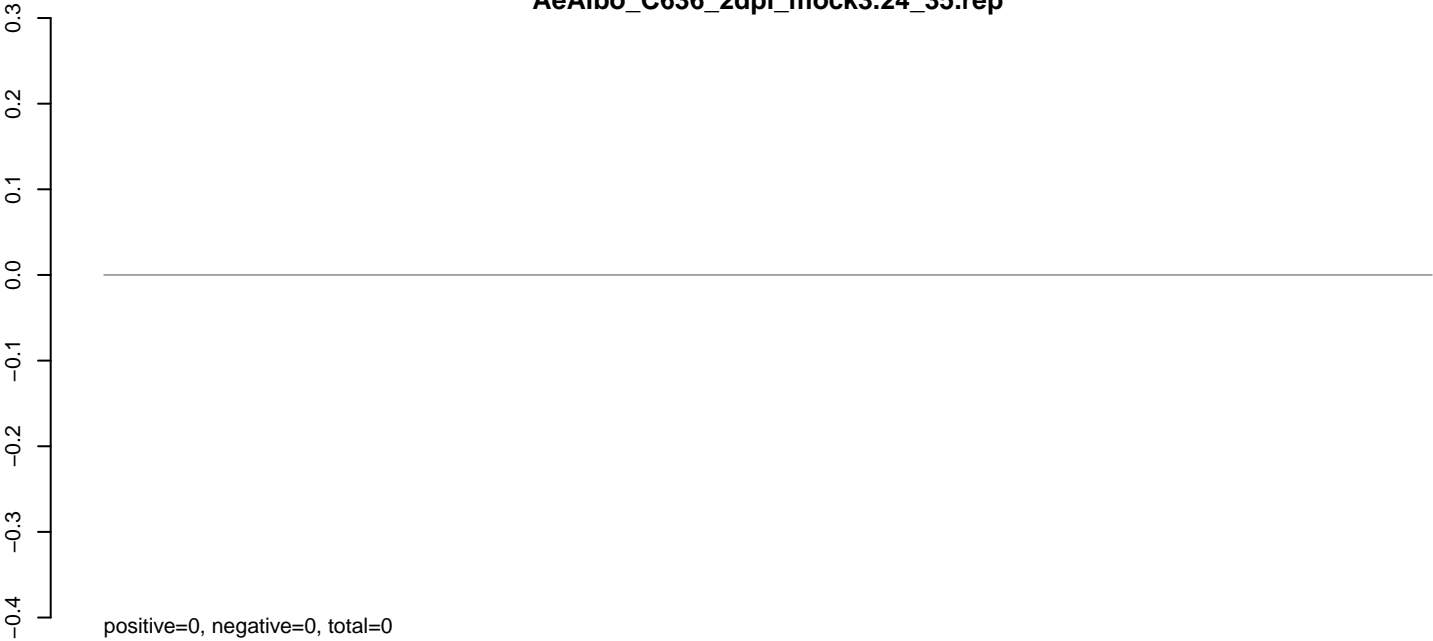
AeAlbo_C636_2dpi_mock3.rep



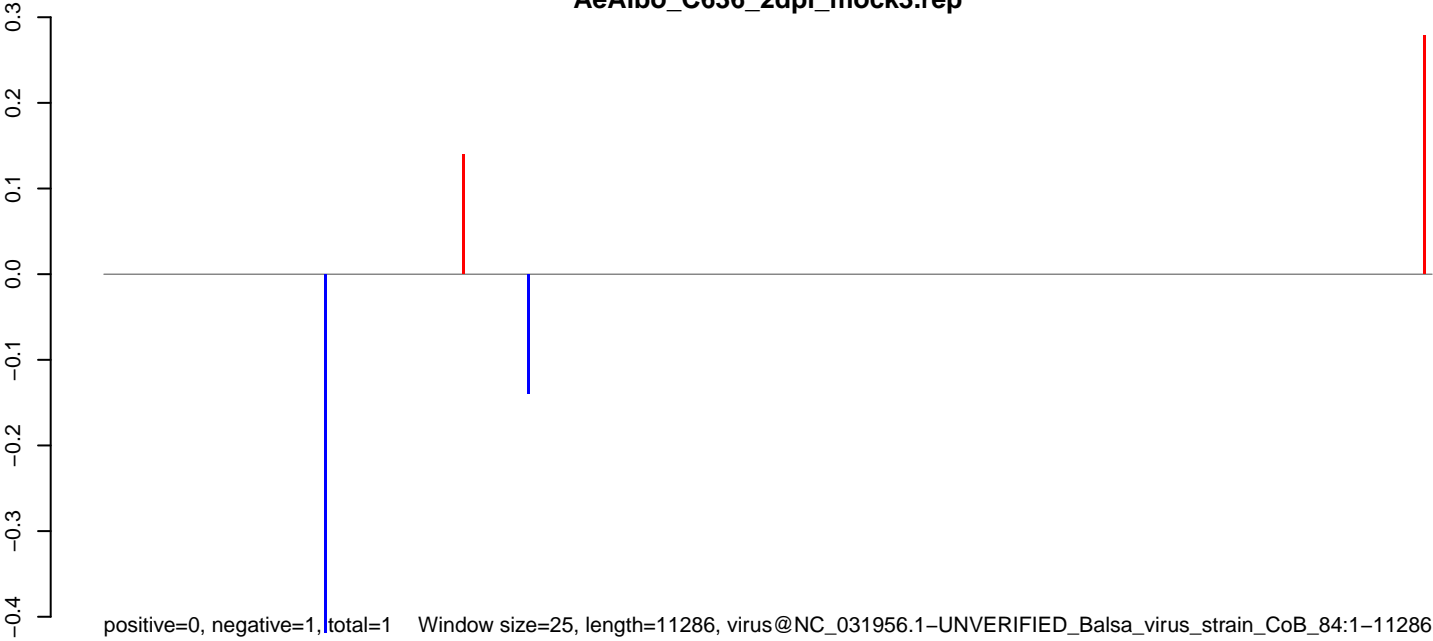
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



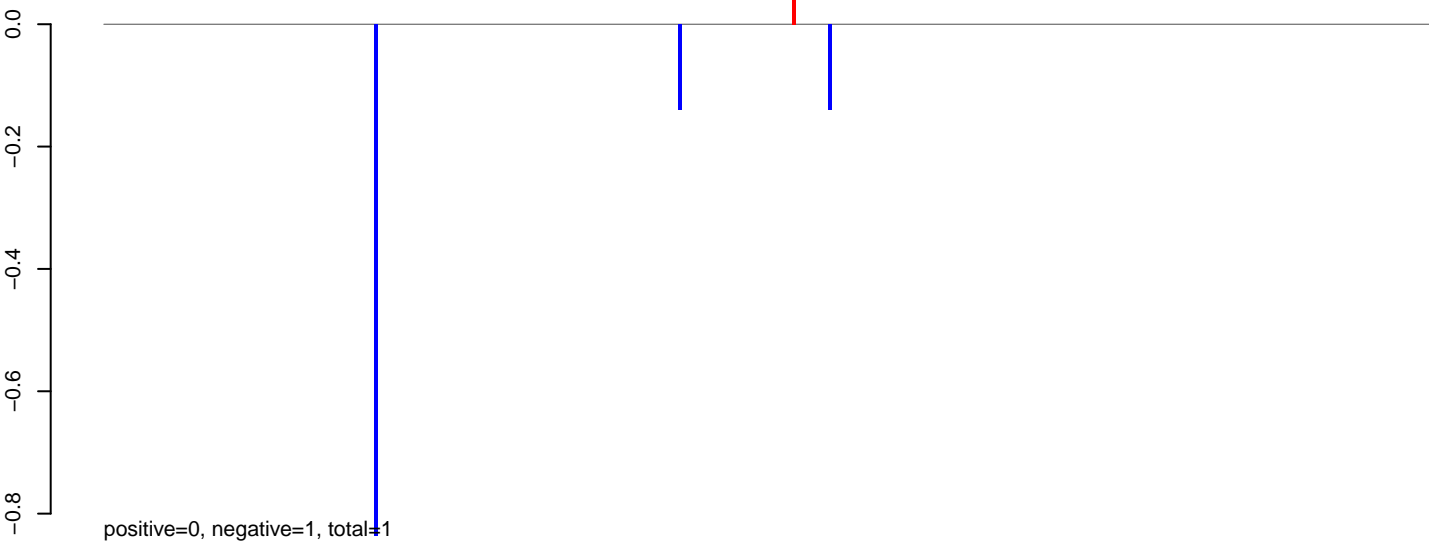
AeAlbo_C636_2dpi_mock3.rep



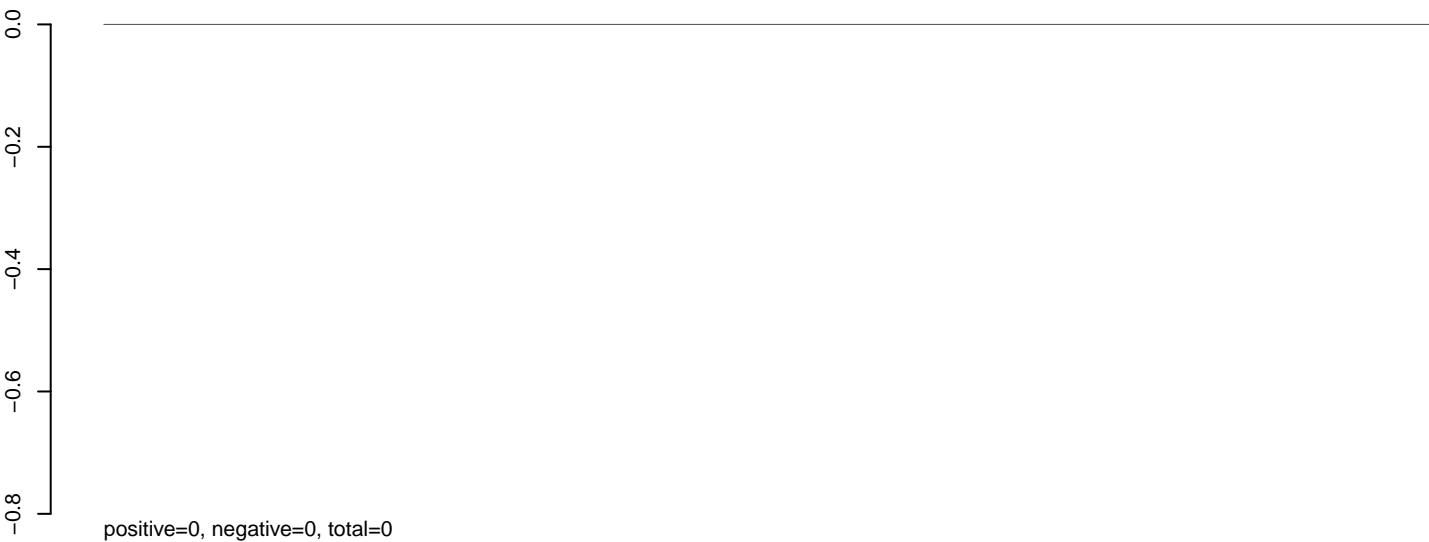
Window size=25, length=11286, virus@NC_031956.1-UNVERIFIED_Balsa_virus_strain_CoB_84:1-11286

0 2000 4000 6000 8000 10000

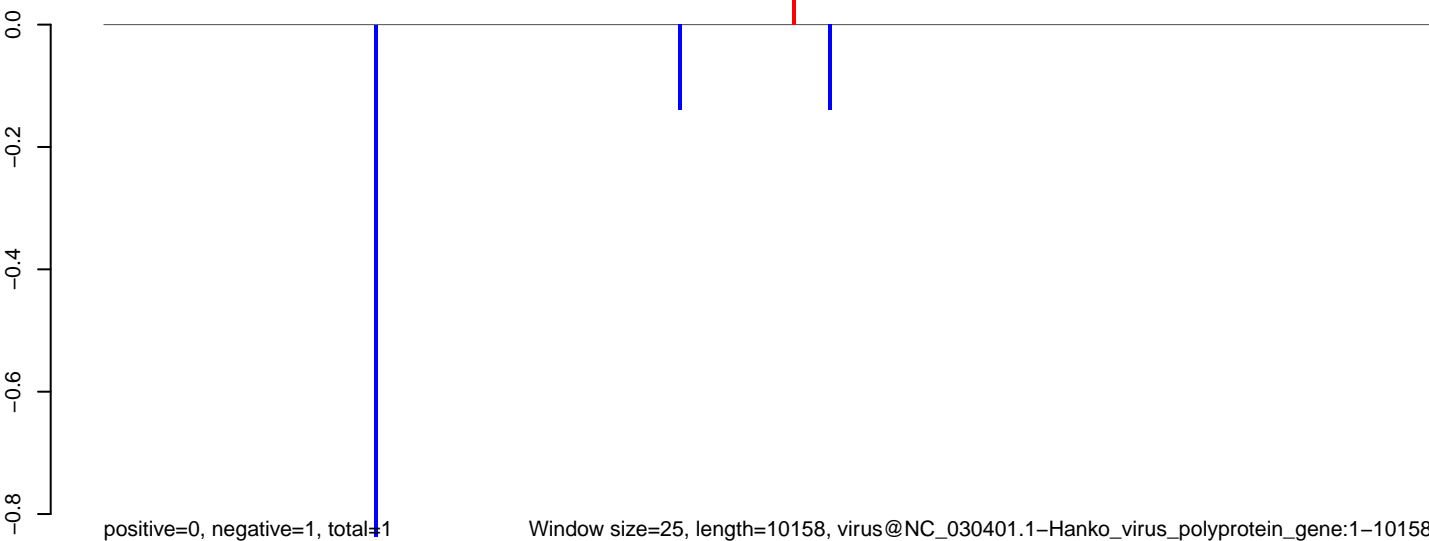
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

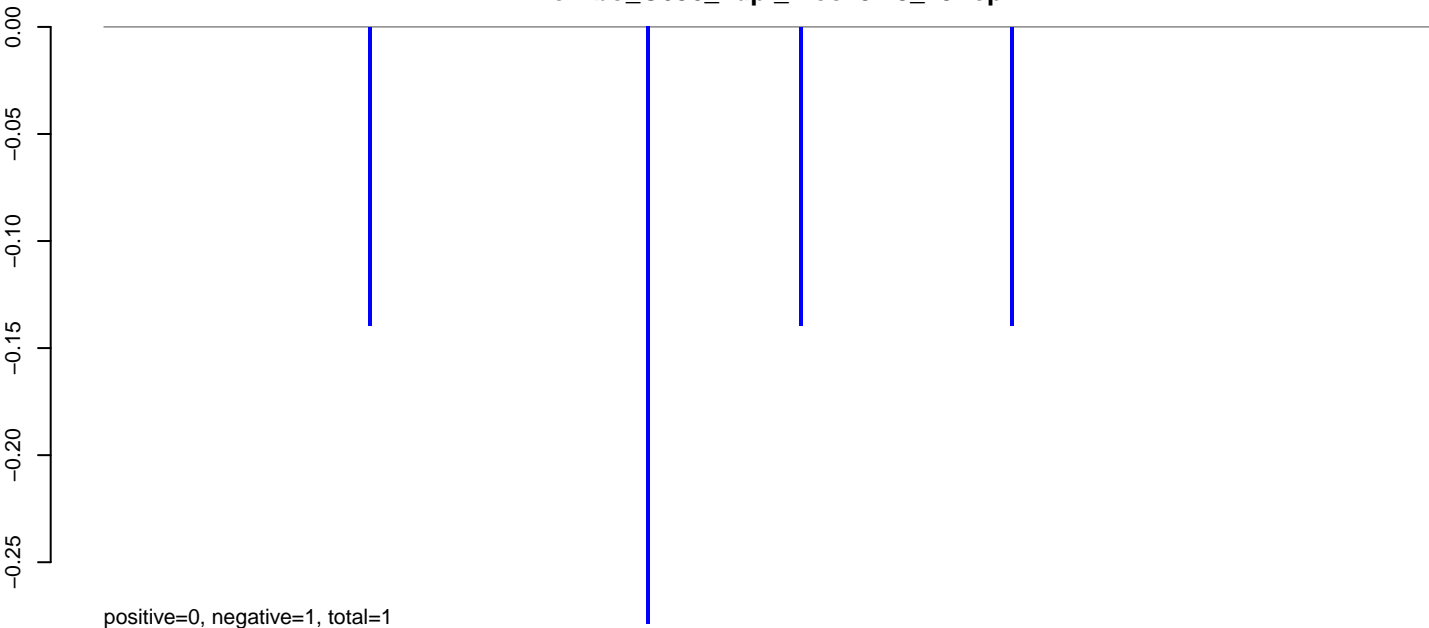


AeAlbo_C636_2dpi_mock3.rep

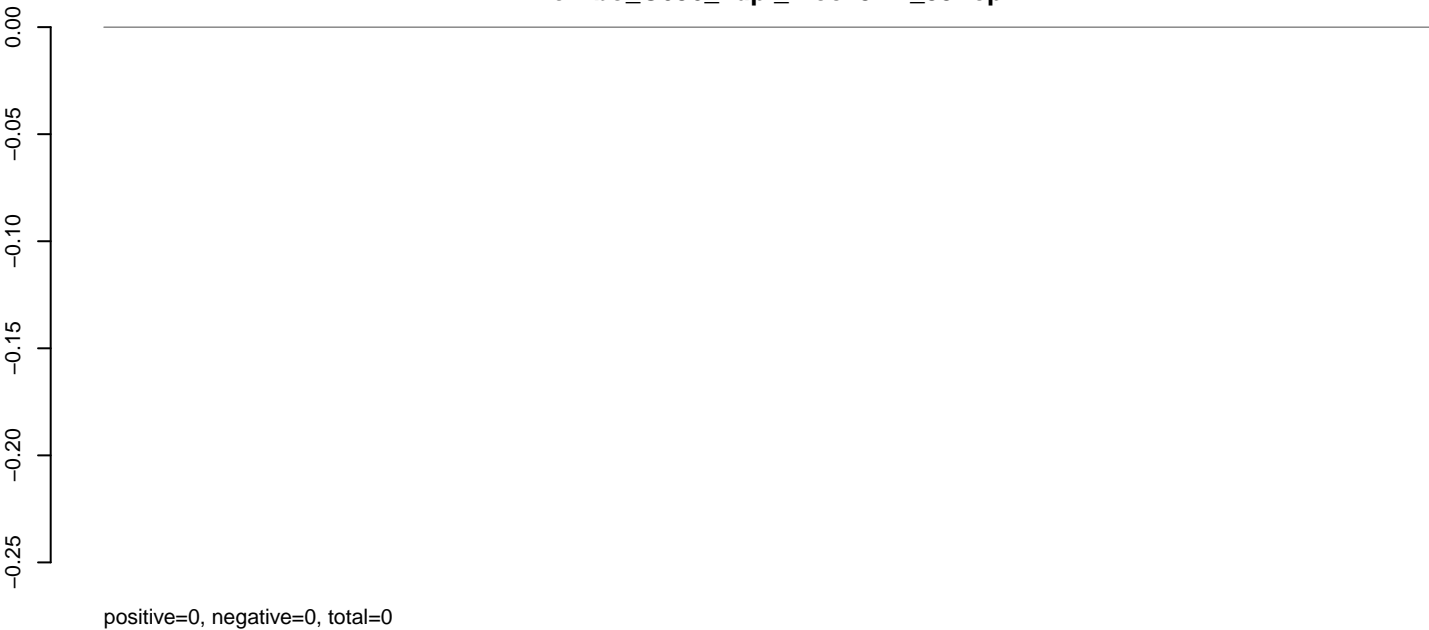


Window size=25, length=10158, virus@NC_030401.1-Hanko_virus_polyprotein_gene:1-10158

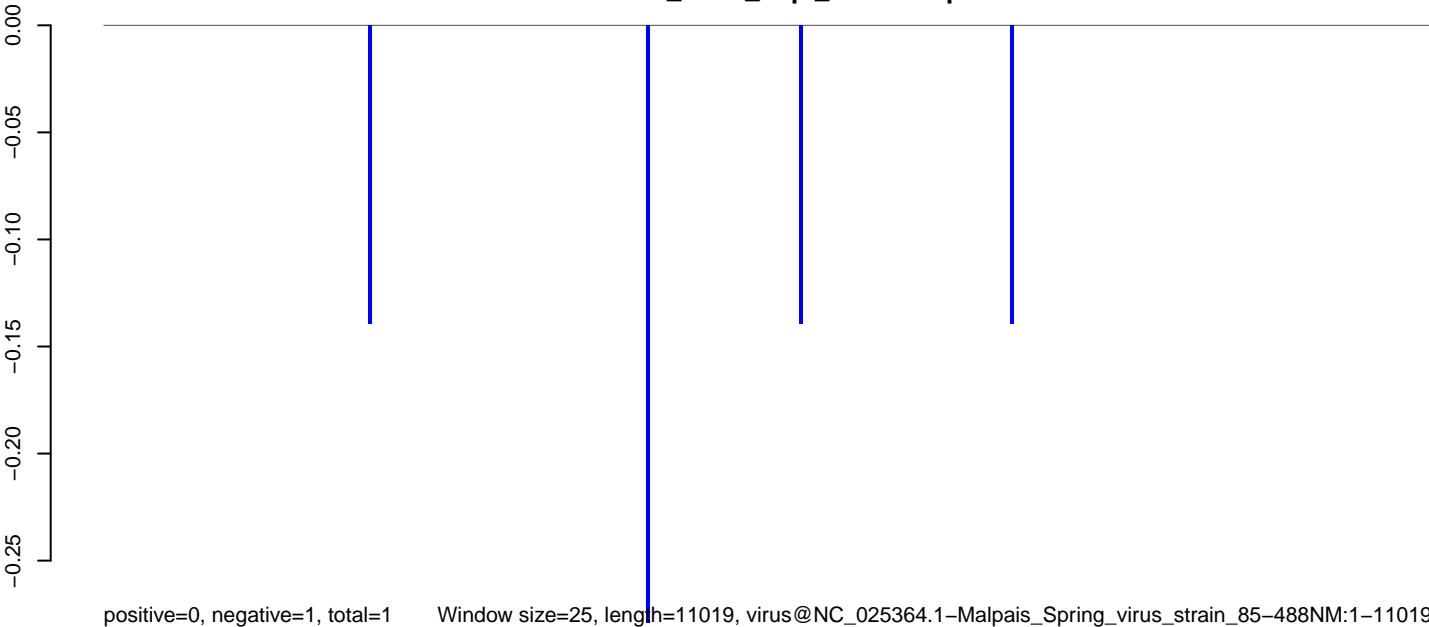
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep

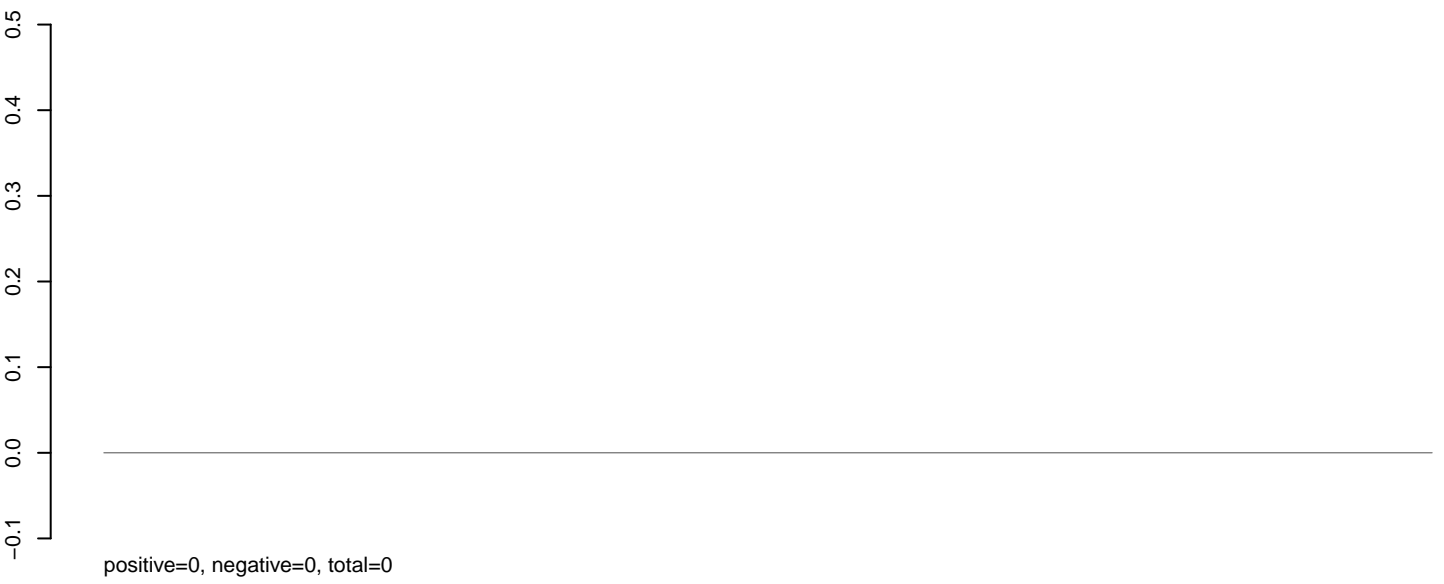


Window size=25, length=11019, virus@NC_025364.1-Malpais_Spring_virus_strain_85-488NM:1-11019

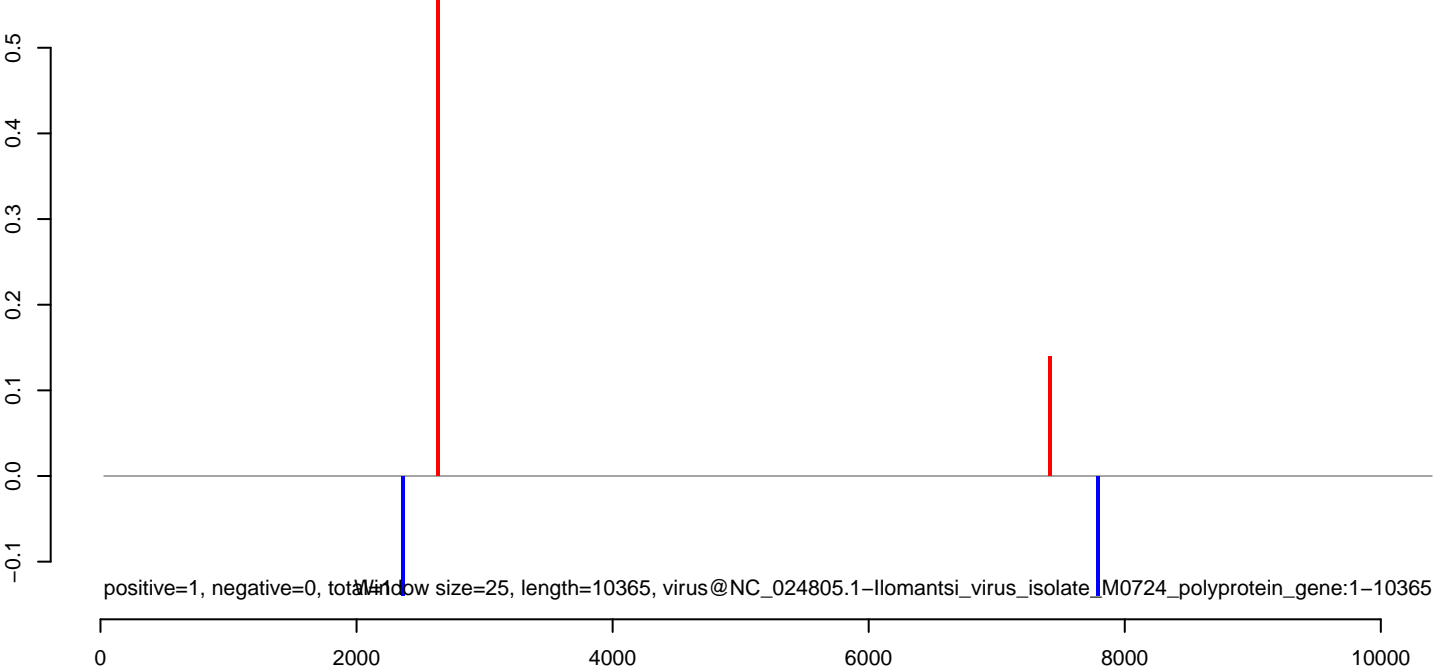
AeAlbo_C636_2dpi_mock3.18_23.rep



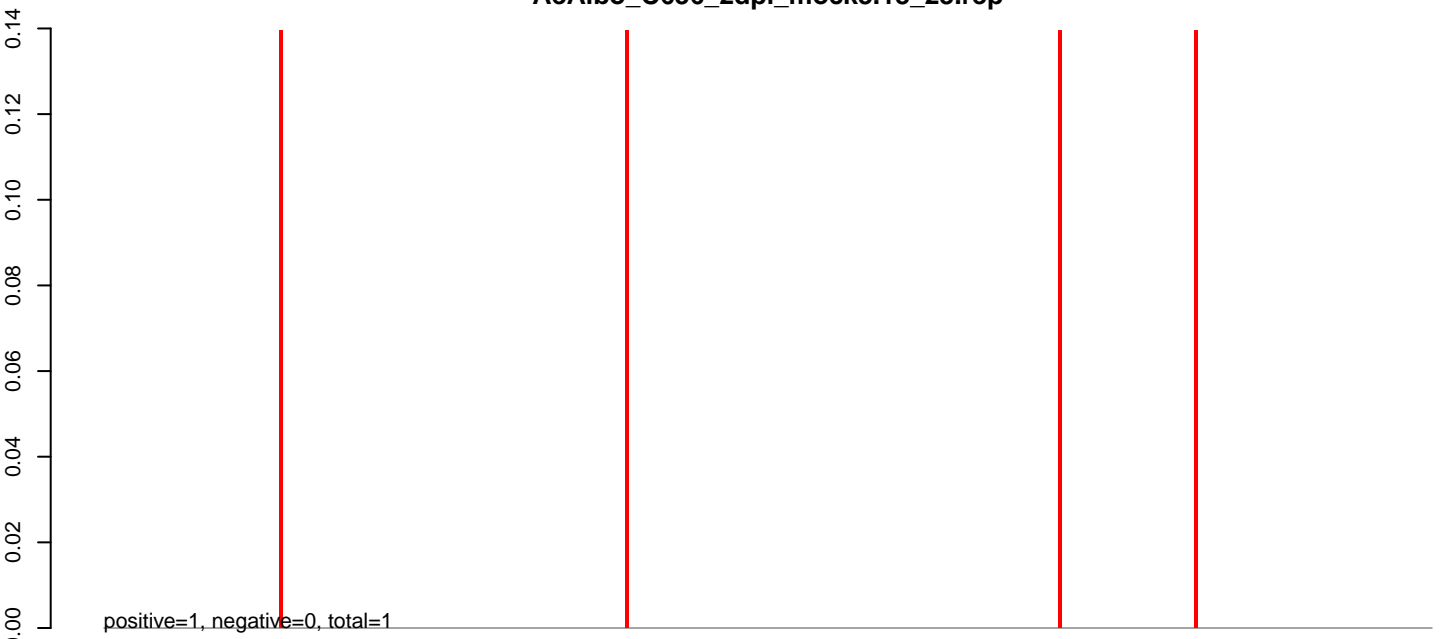
AeAlbo_C636_2dpi_mock3.24_35.rep



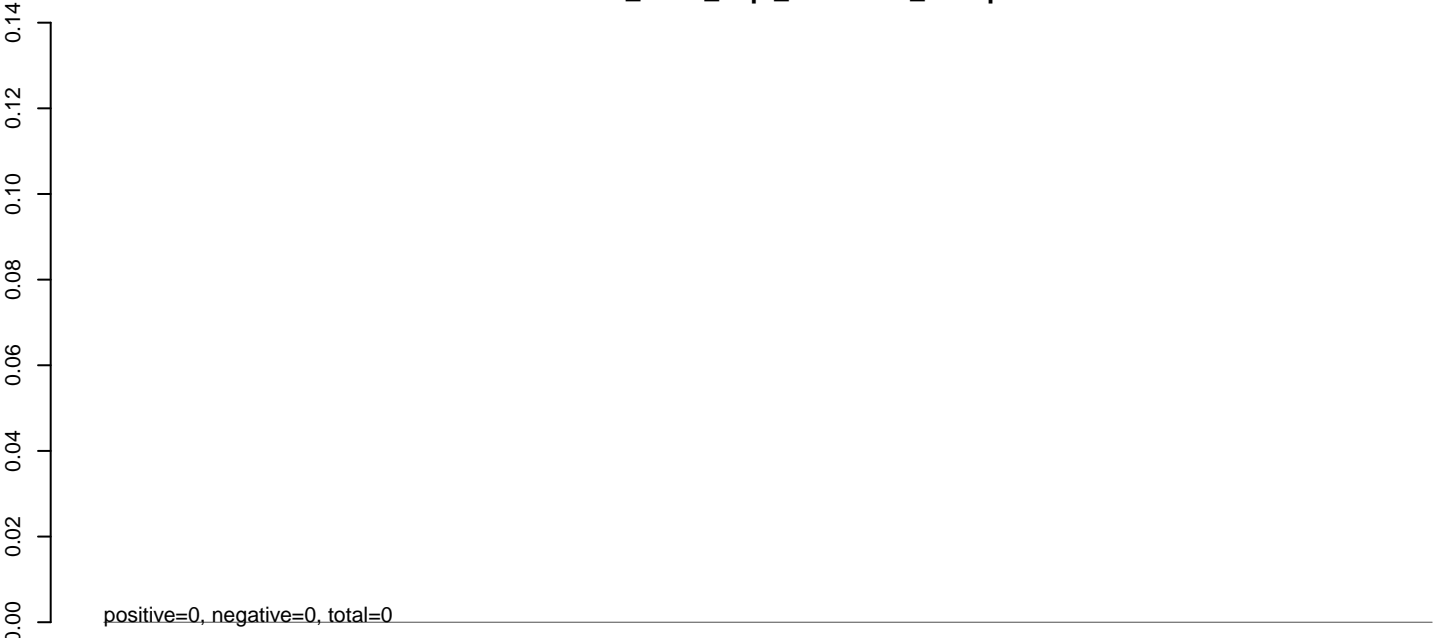
AeAlbo_C636_2dpi_mock3.rep



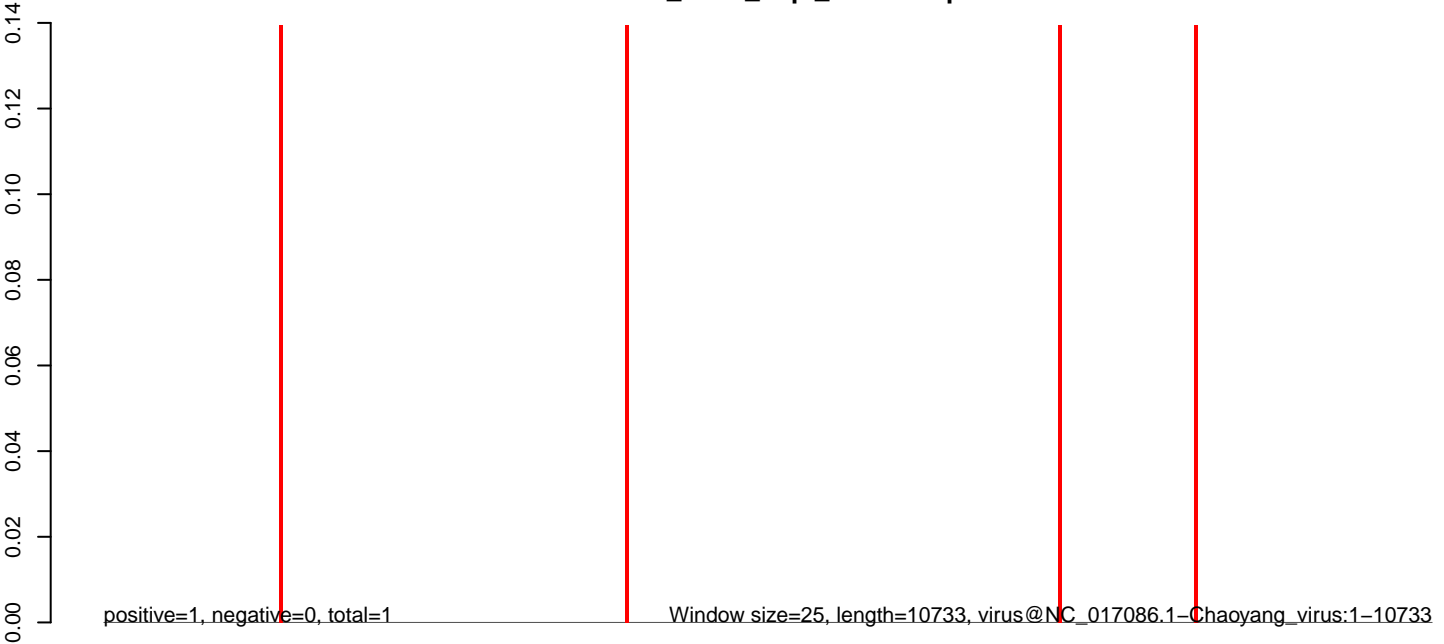
AeAlbo_C636_2dpi_mock3.18_23.rep



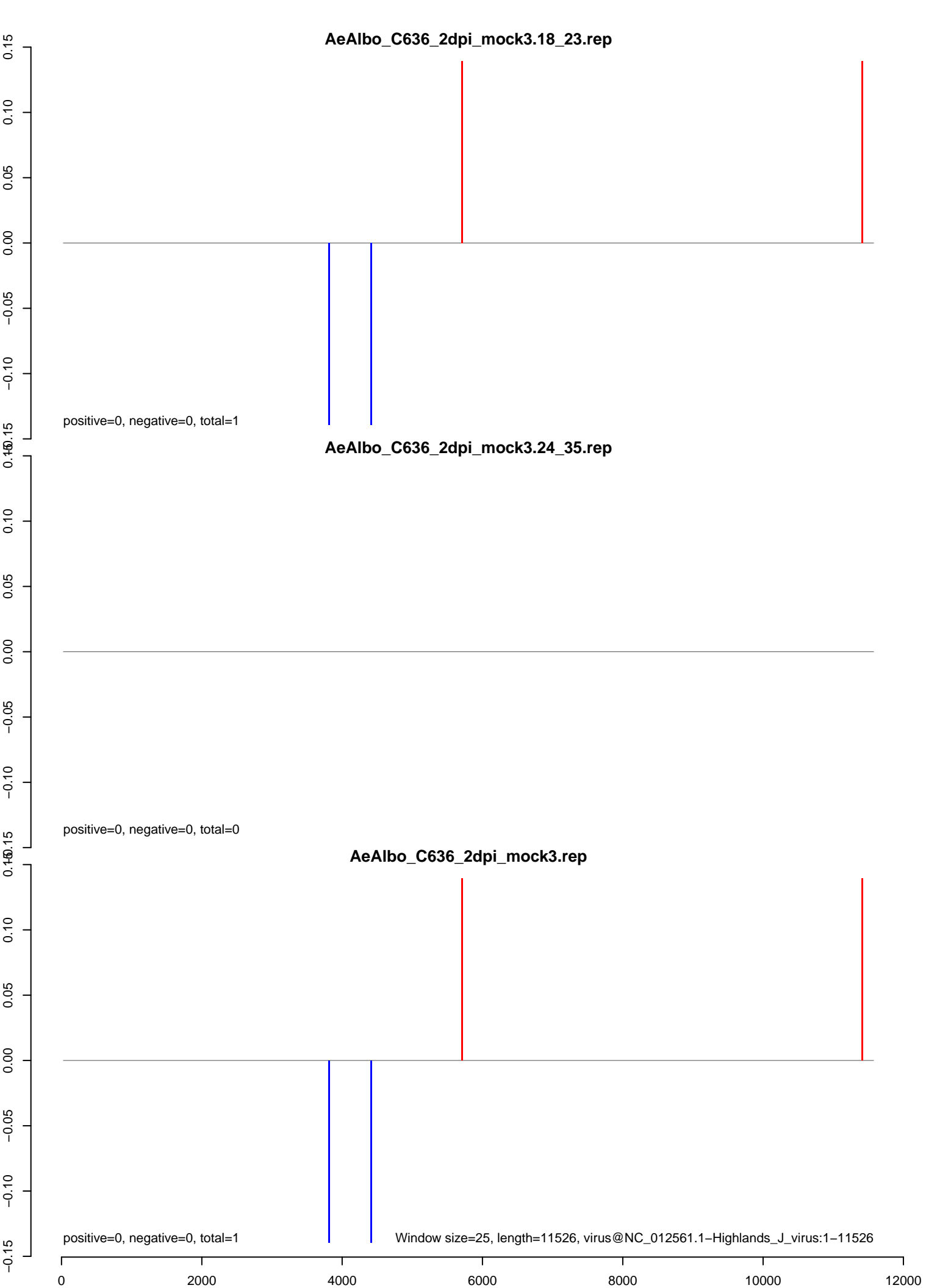
AeAlbo_C636_2dpi_mock3.24_35.rep



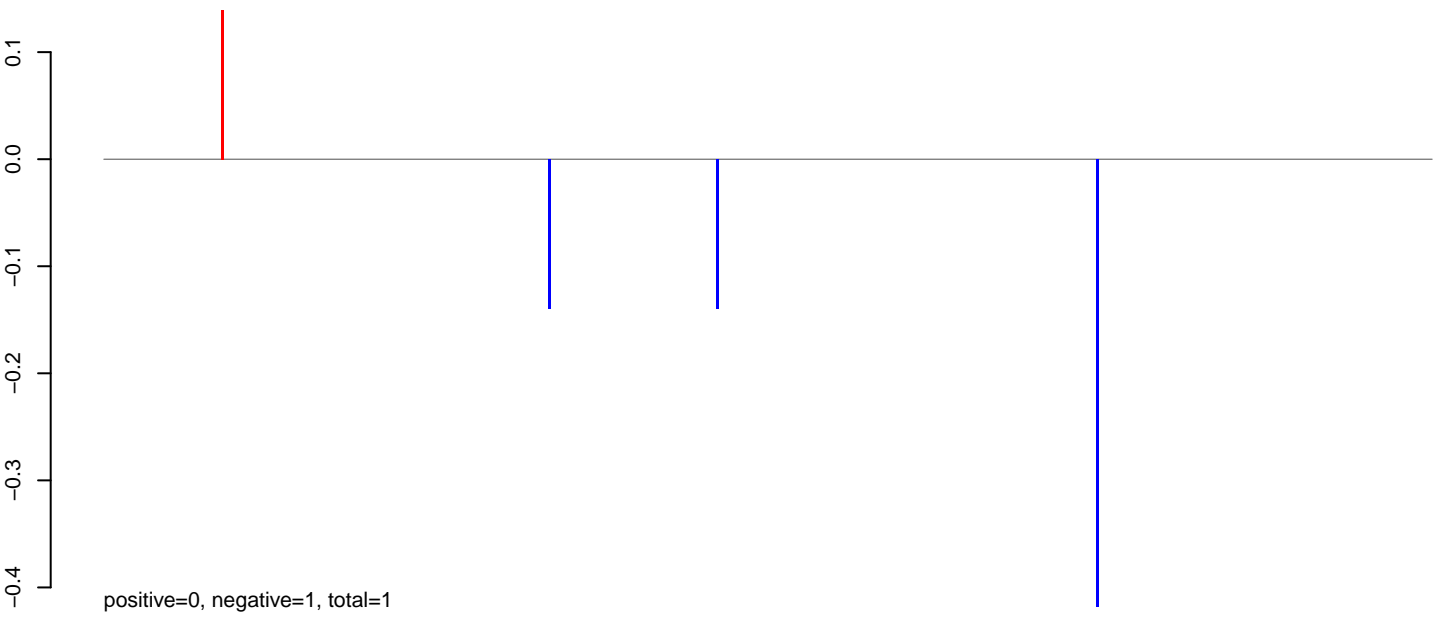
AeAlbo_C636_2dpi_mock3.rep



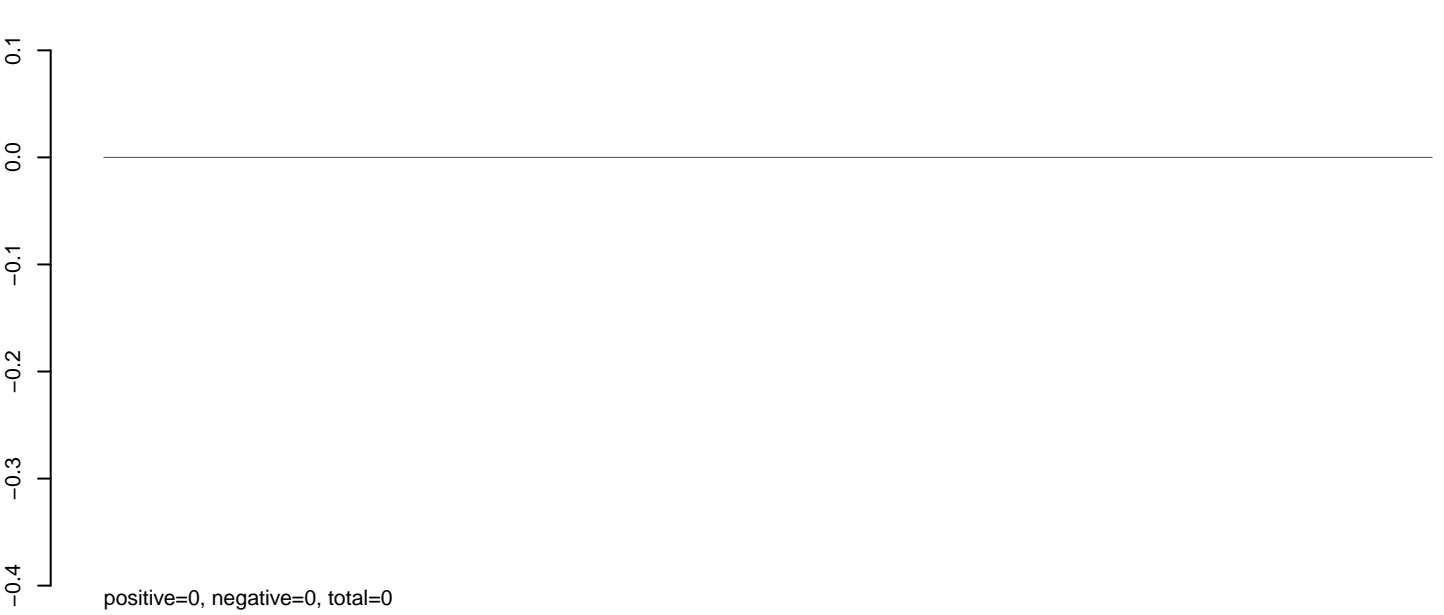
0 2000 4000 6000 8000 10000



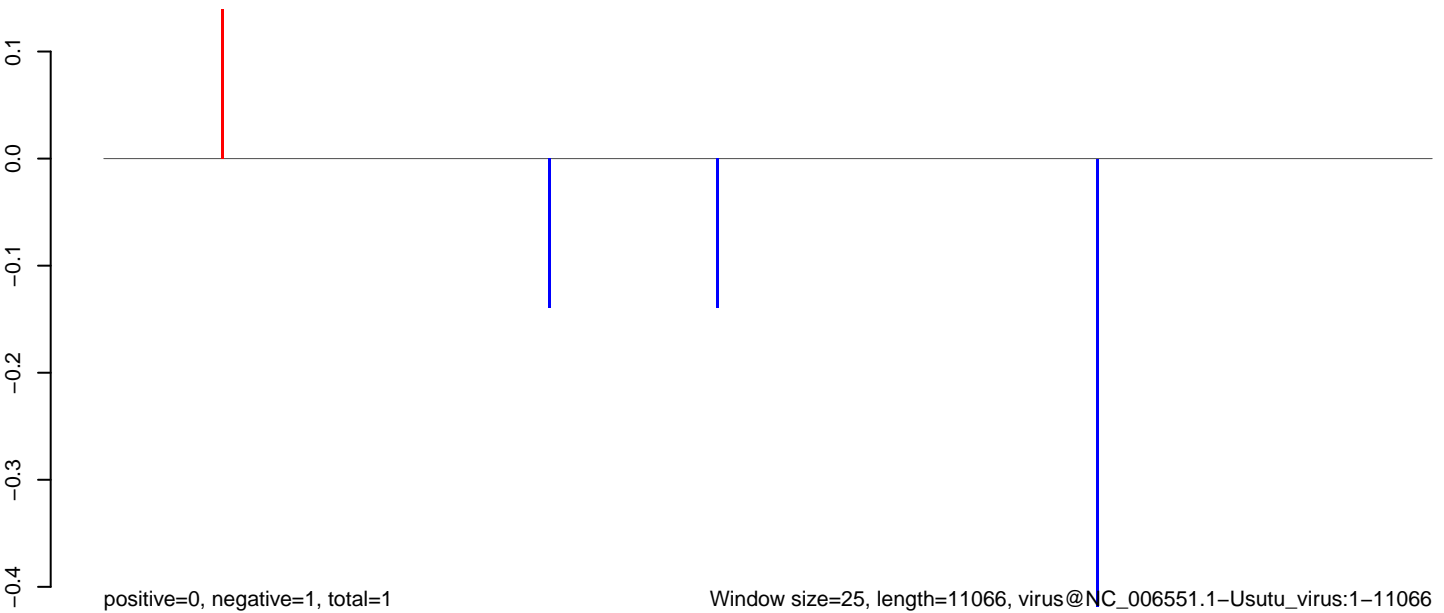
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

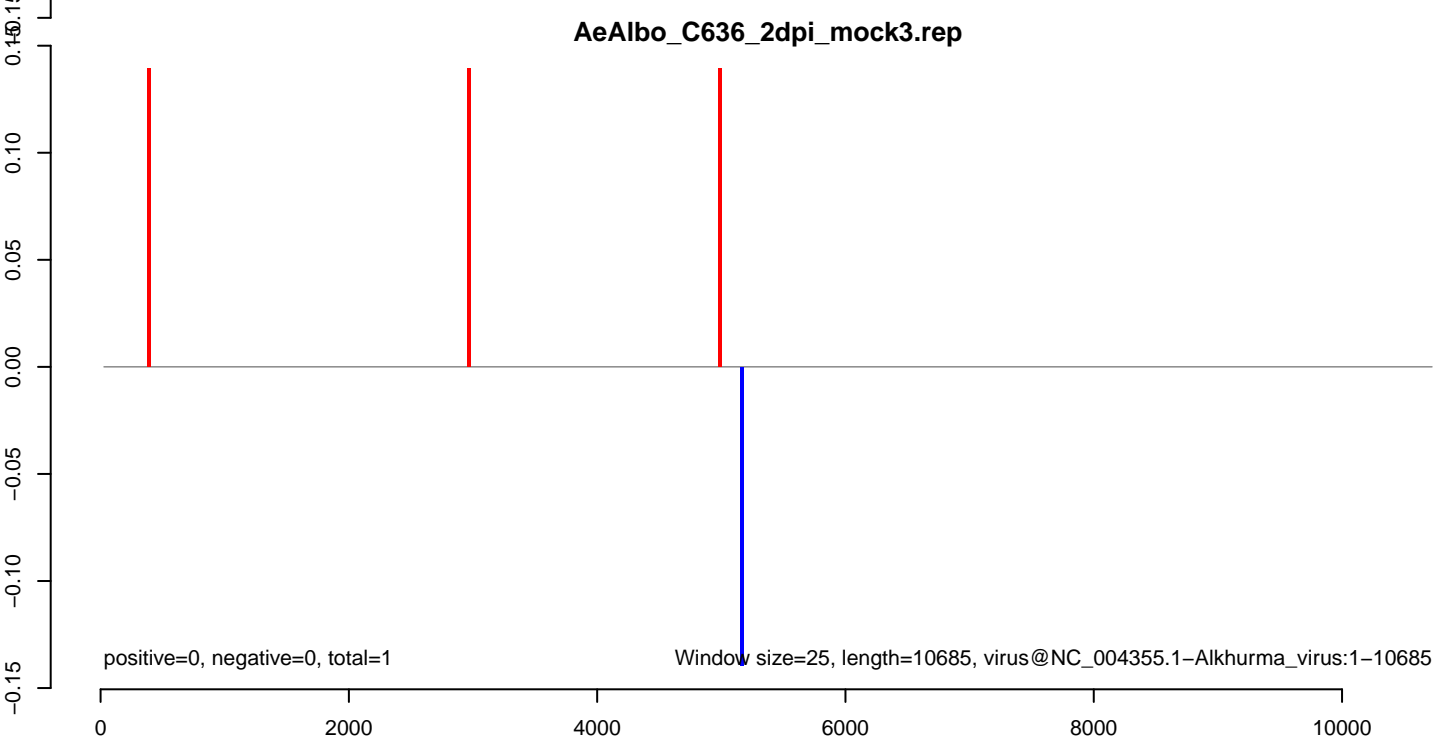
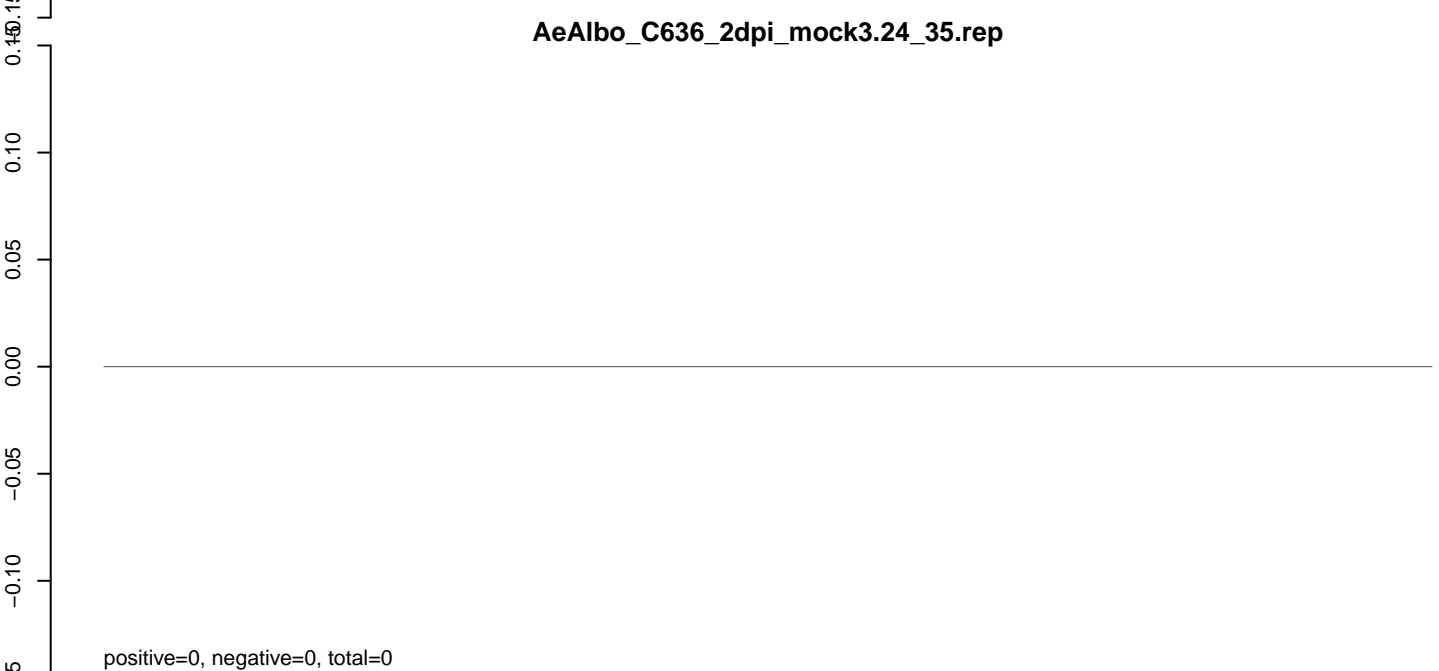
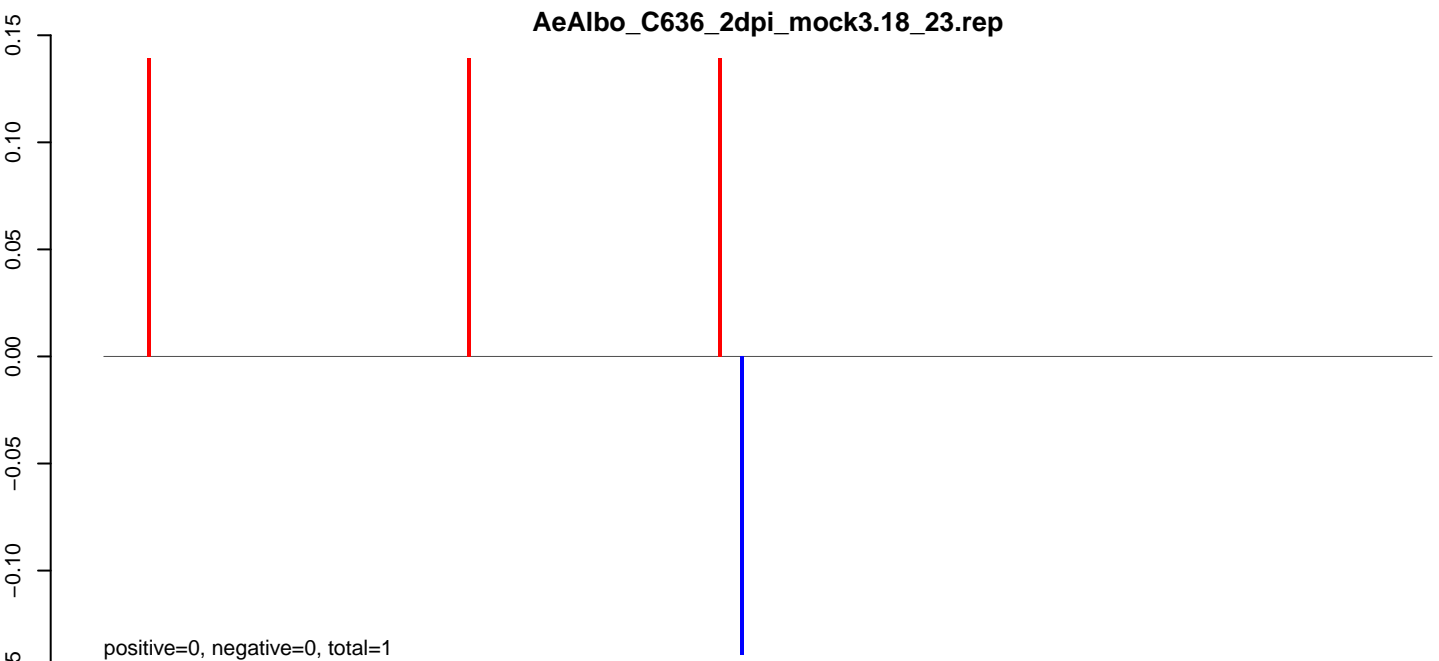


AeAlbo_C636_2dpi_mock3.rep

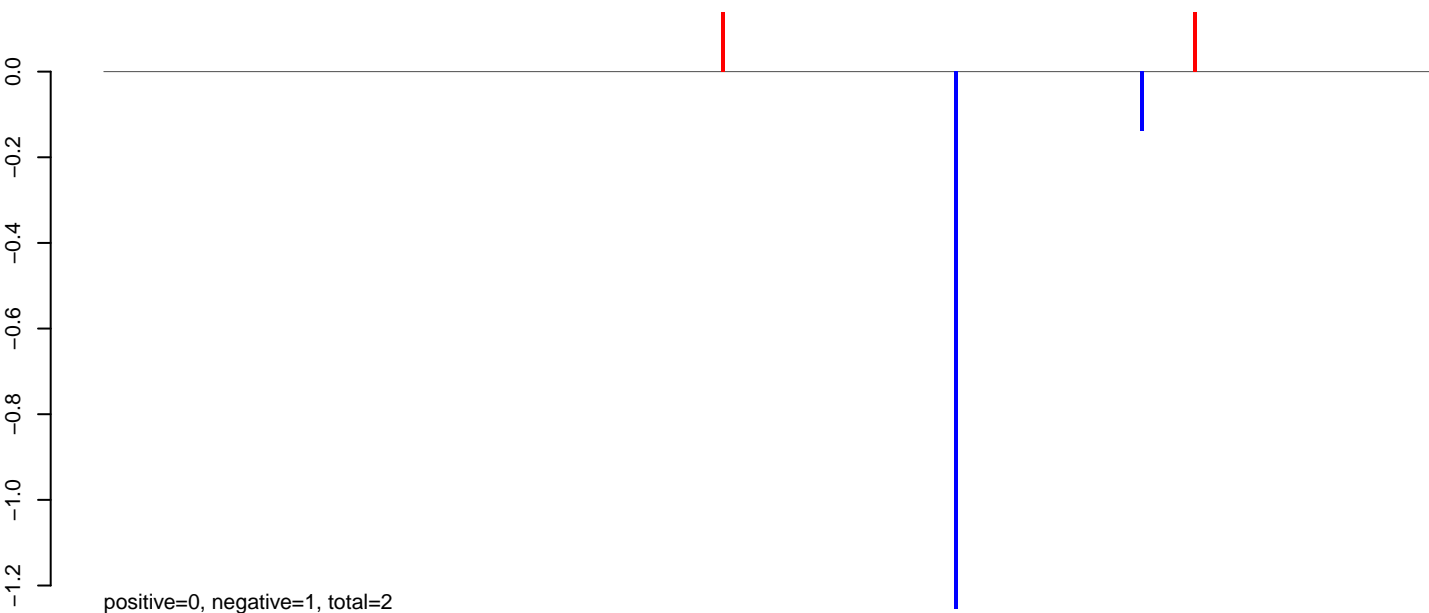


Window size=25, length=11066, virus@NC_006551.1-Usutu_virus:1-11066

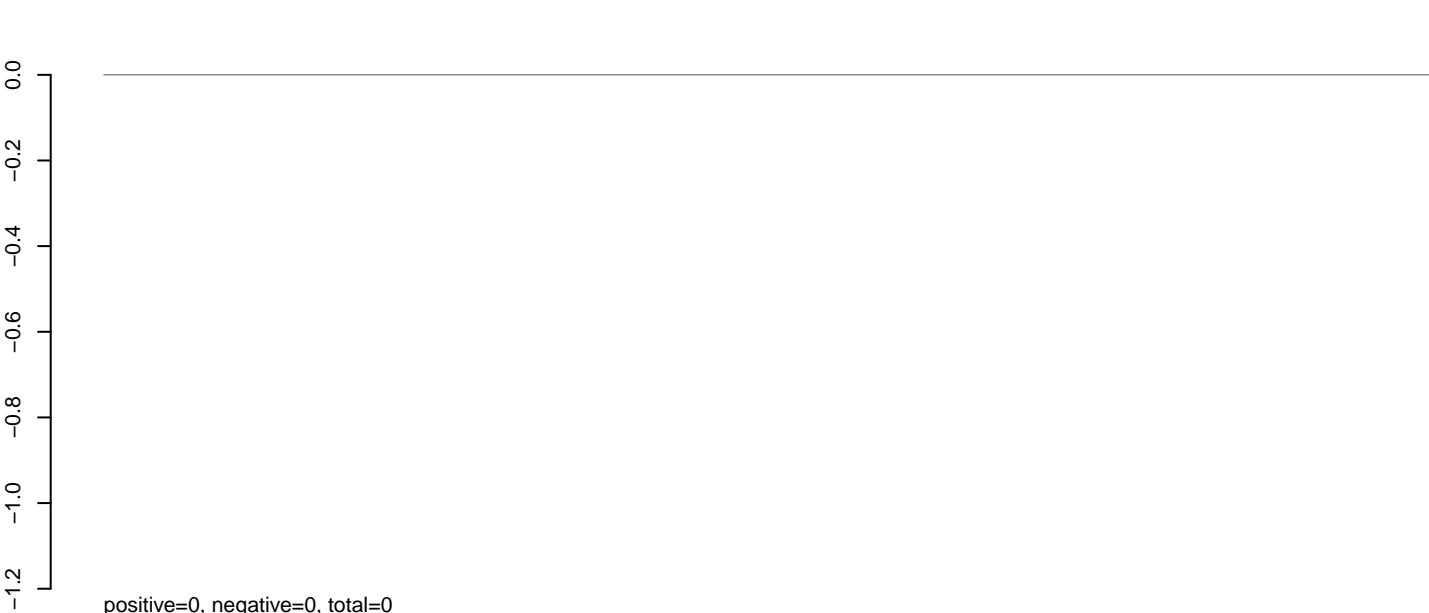
0 2000 4000 6000 8000 10000



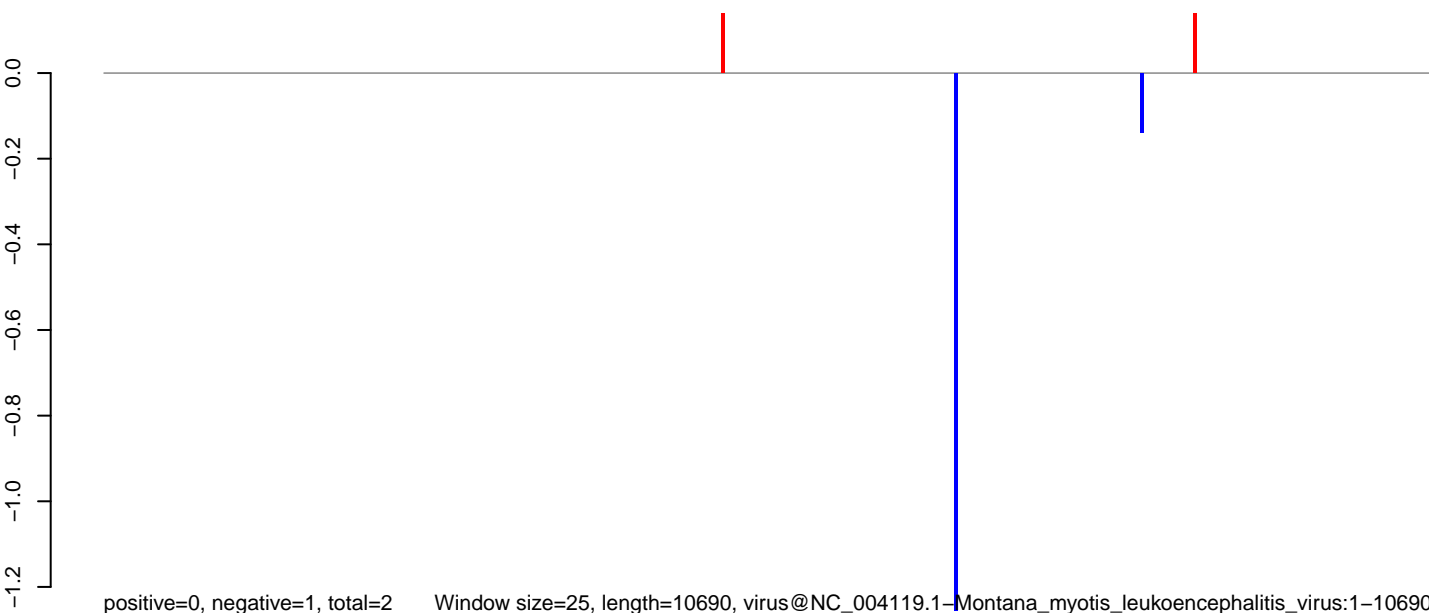
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

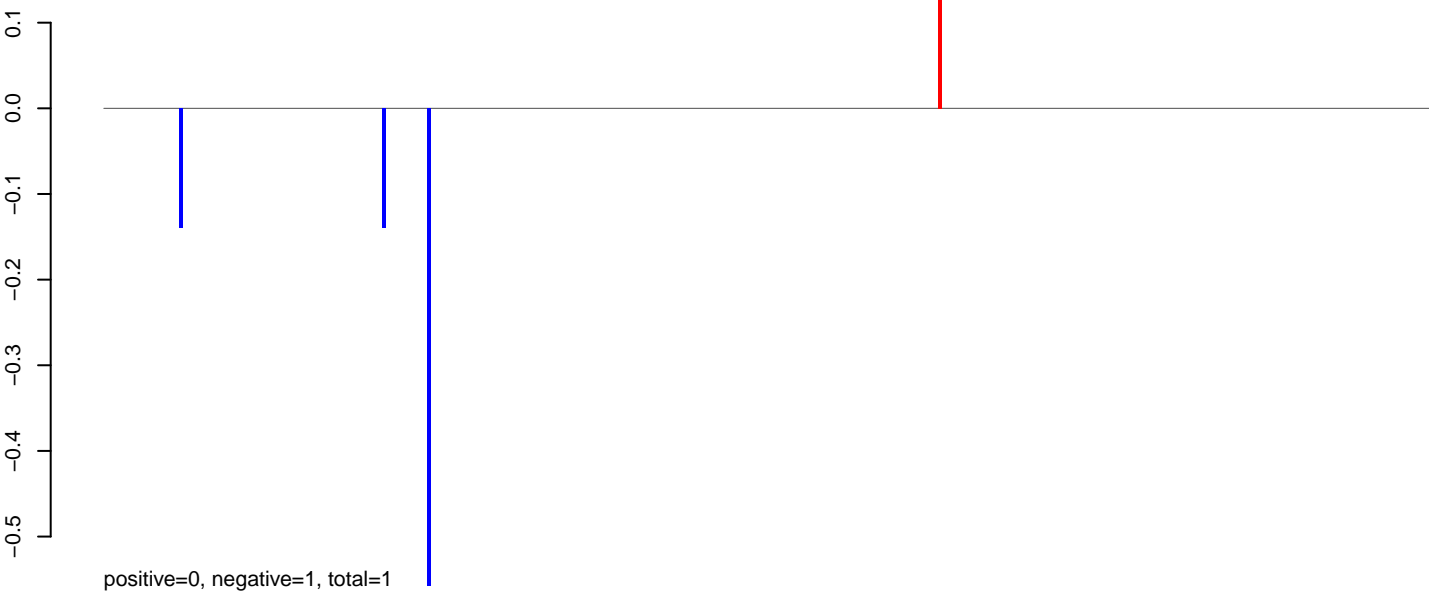


AeAlbo_C636_2dpi_mock3.rep

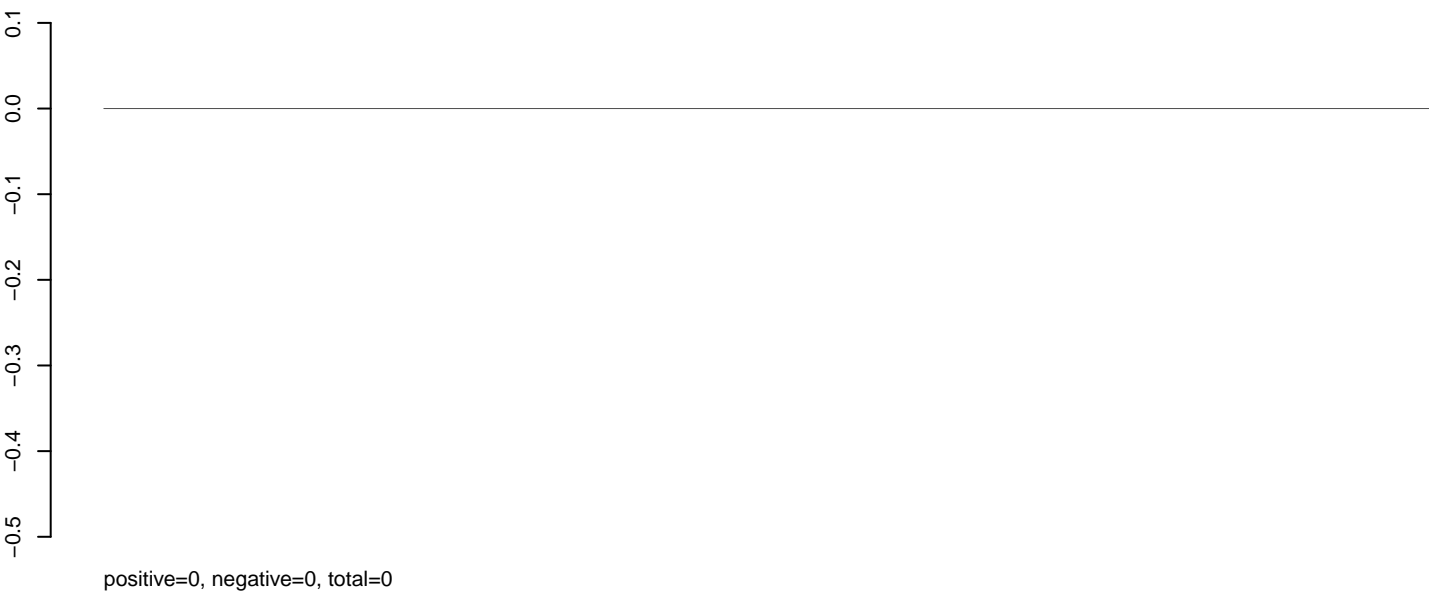


0 2000 4000 6000 8000 10000

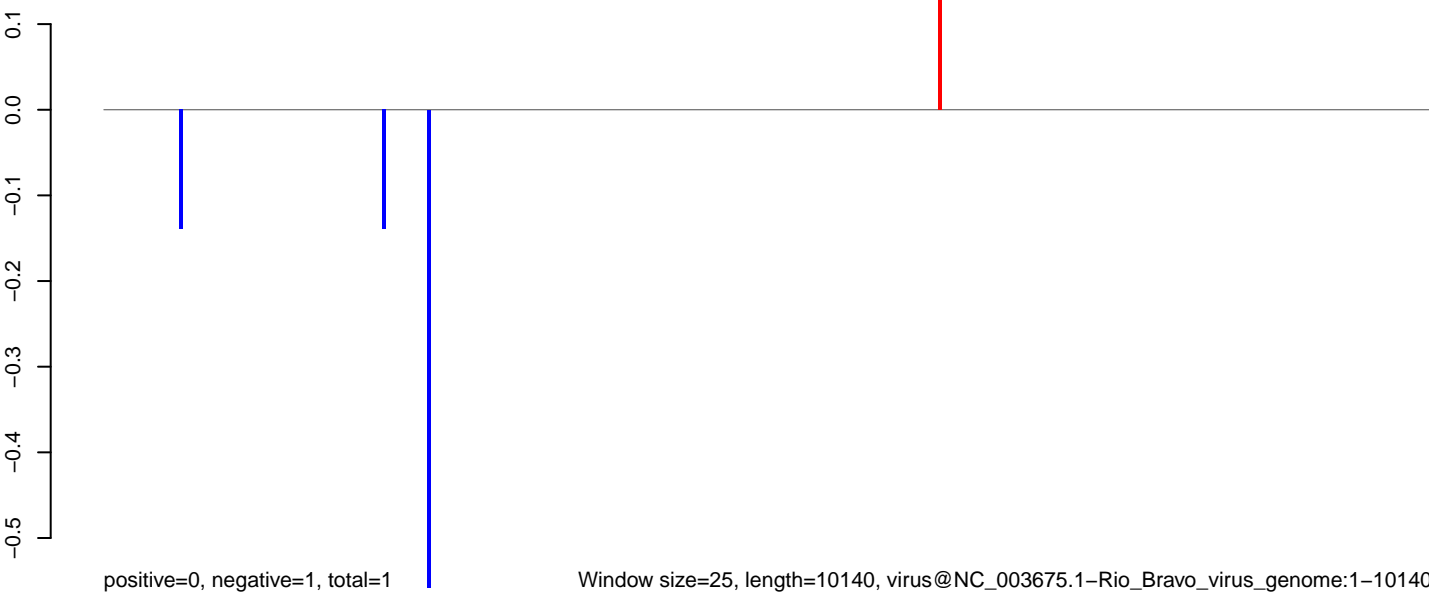
AeAlbo_C636_2dpi_mock3.18_23.rep

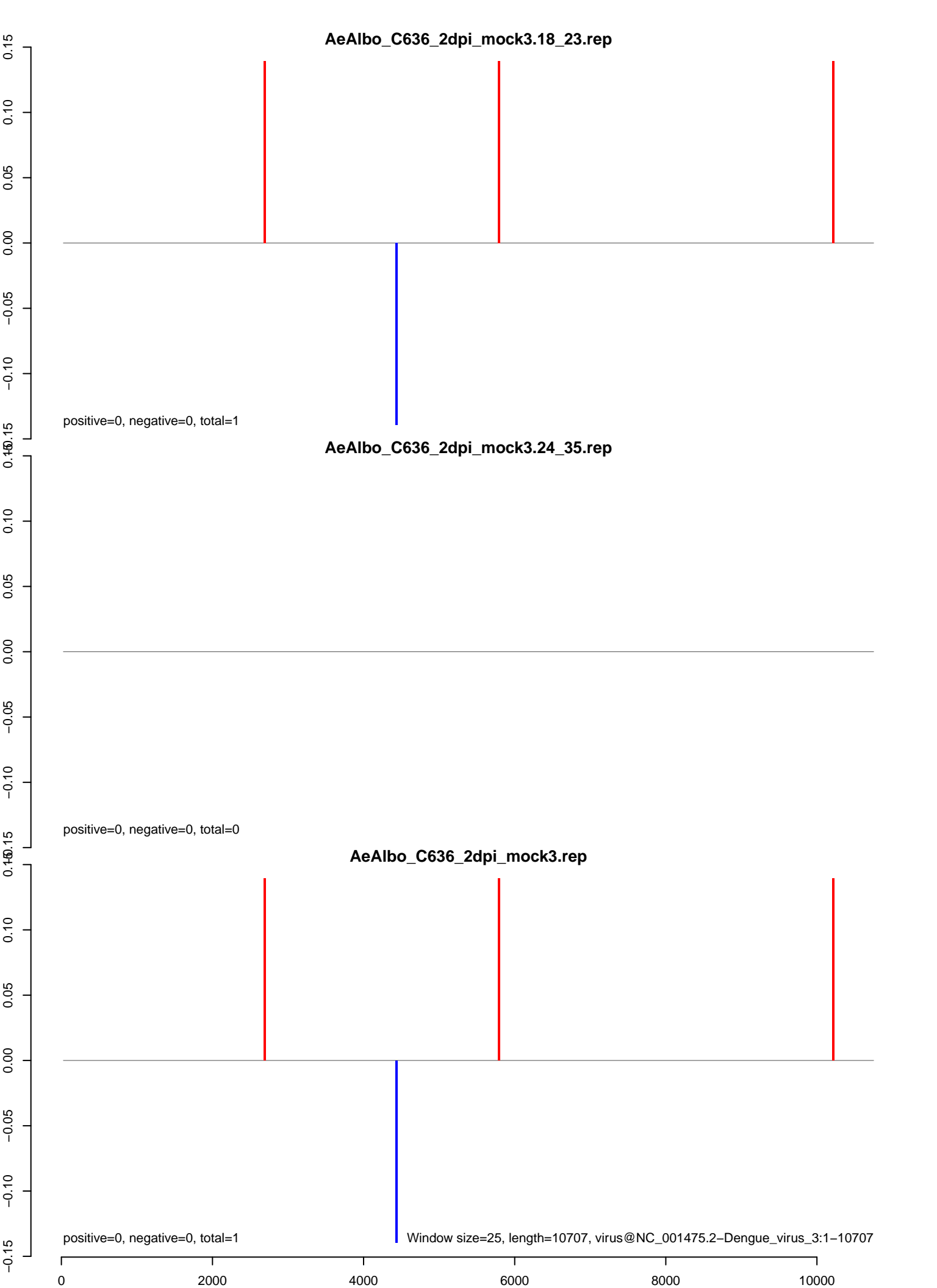


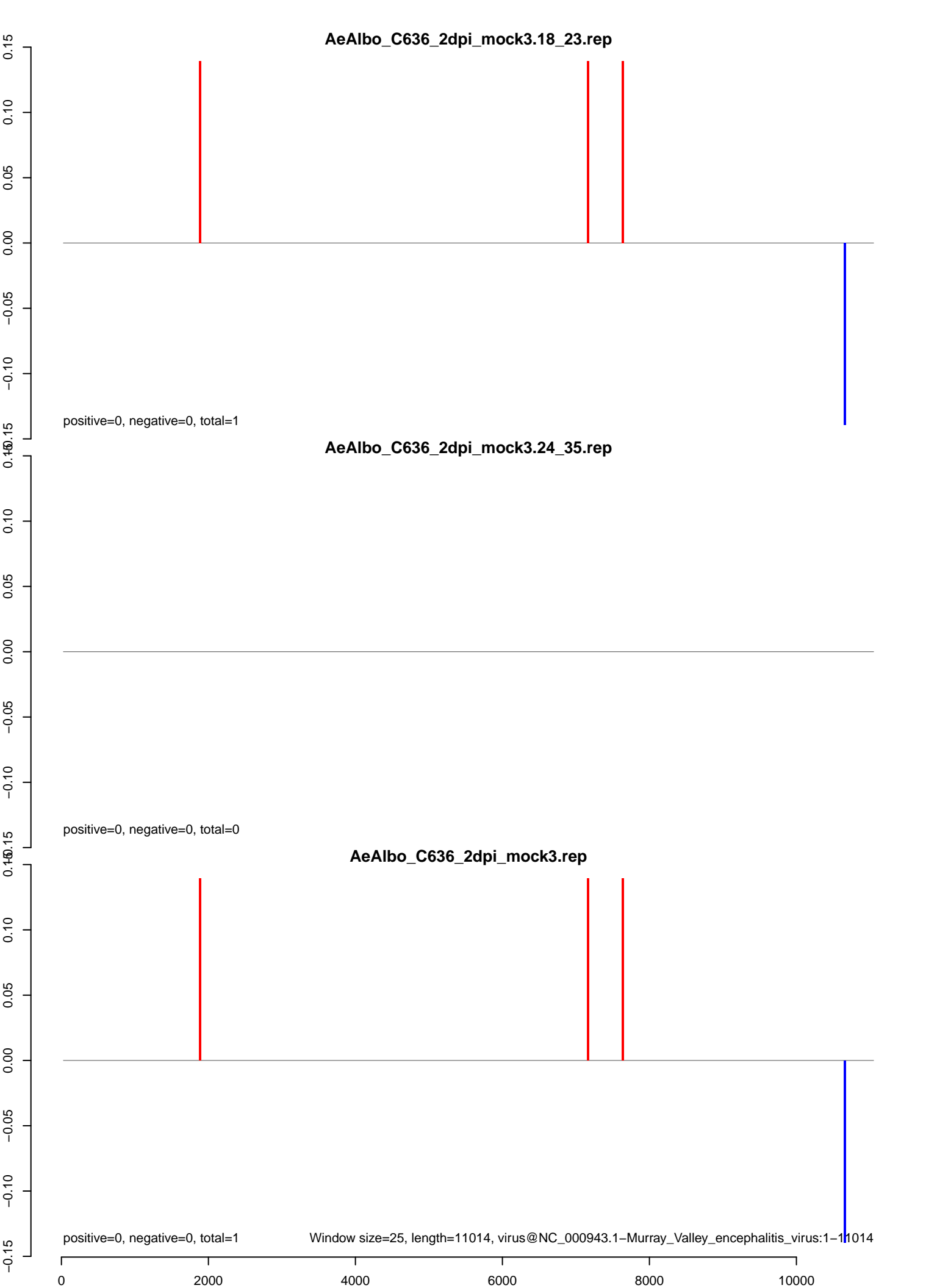
AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



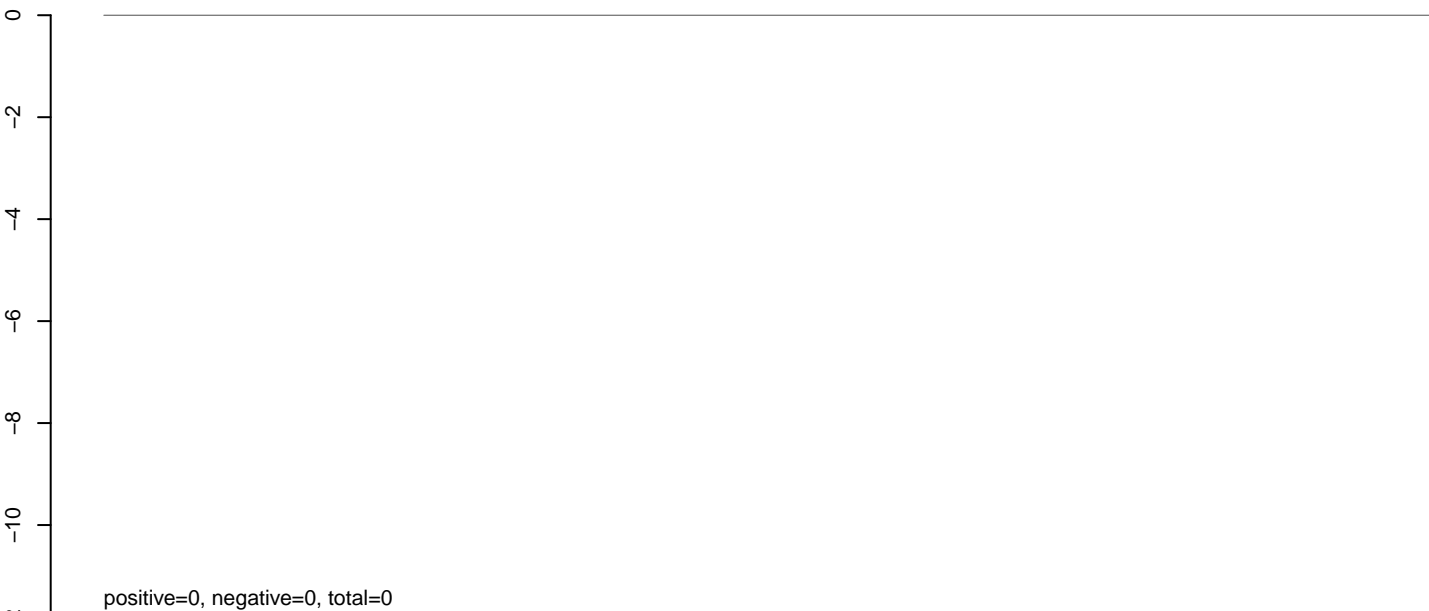




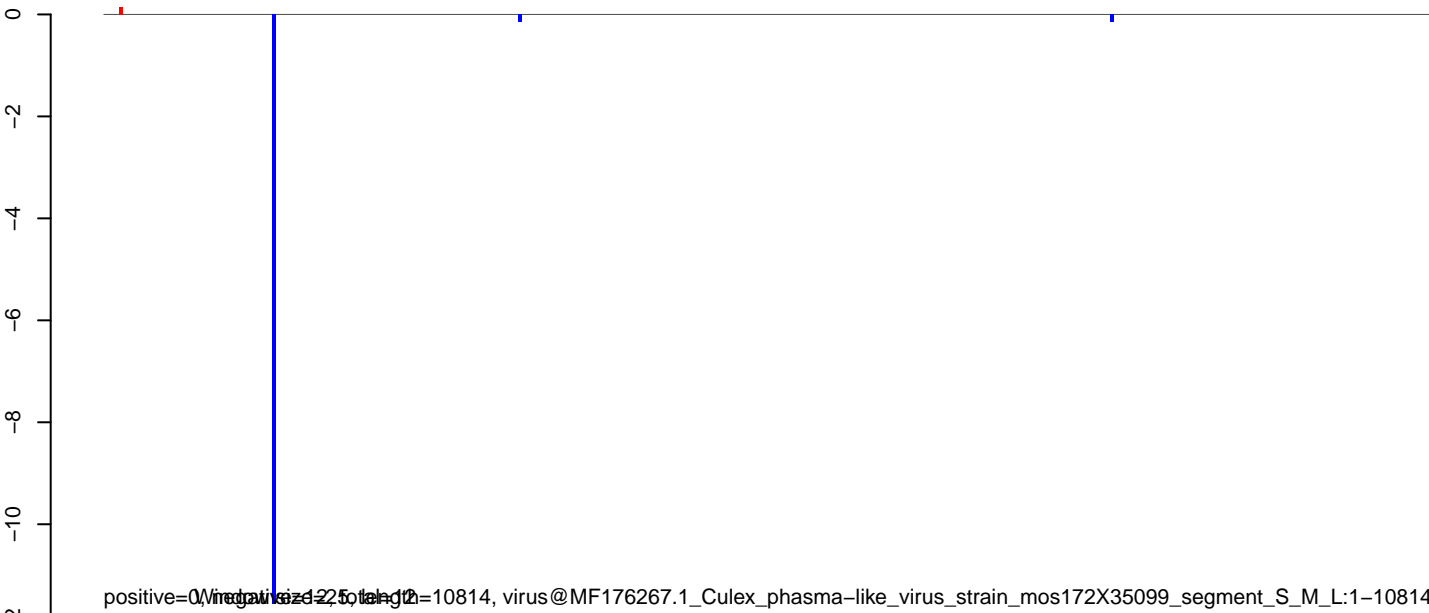
AeAlbo_C636_2dpi_mock3.18_23.rep

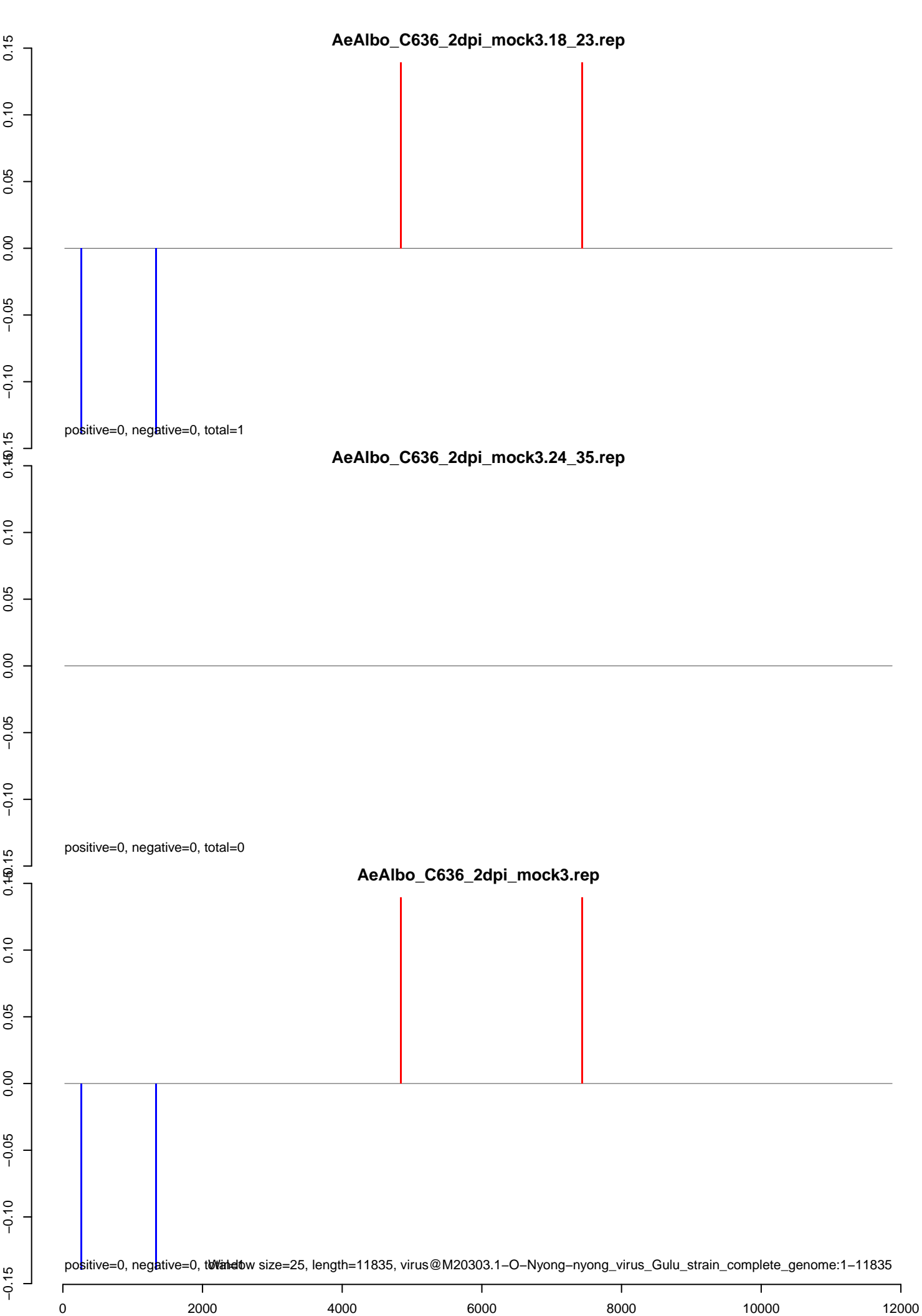


AeAlbo_C636_2dpi_mock3.24_35.rep

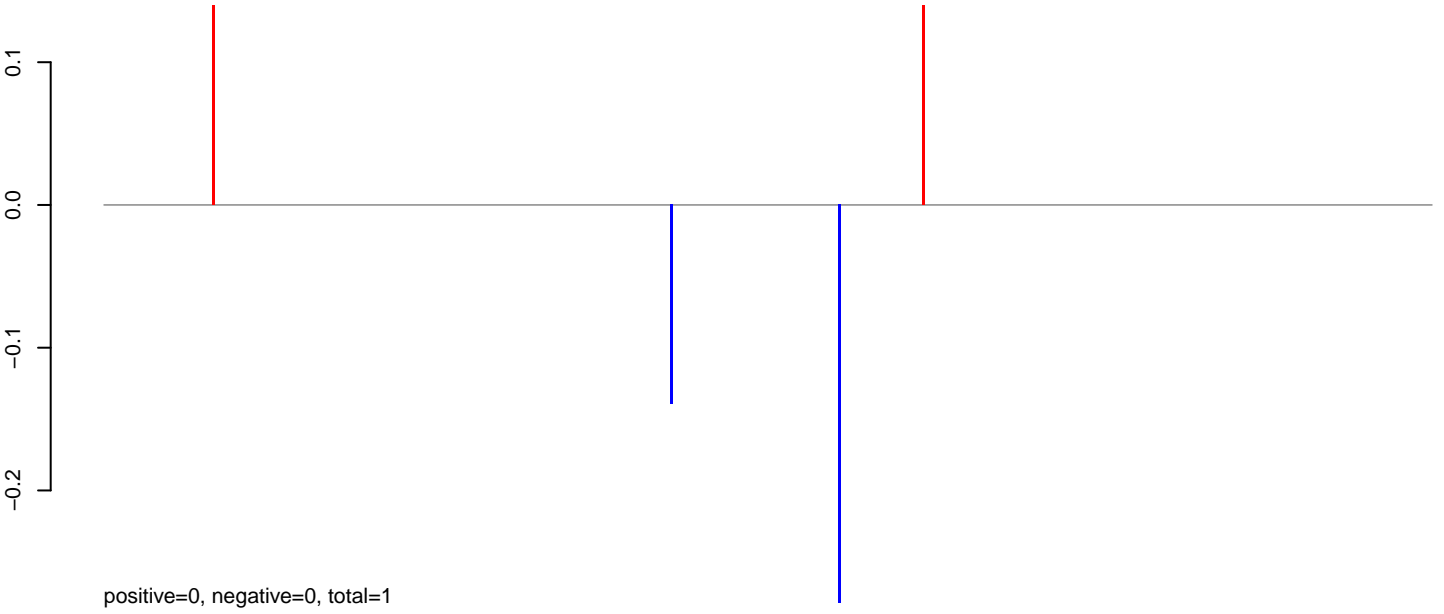


AeAlbo_C636_2dpi_mock3.rep

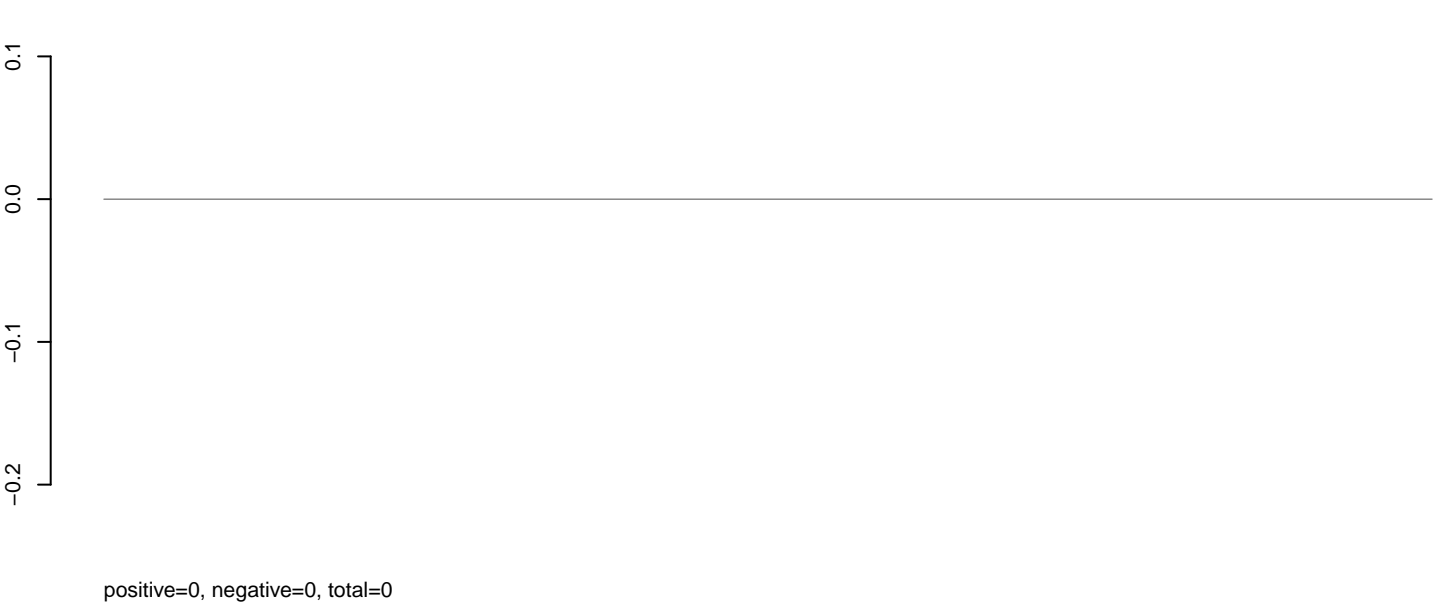




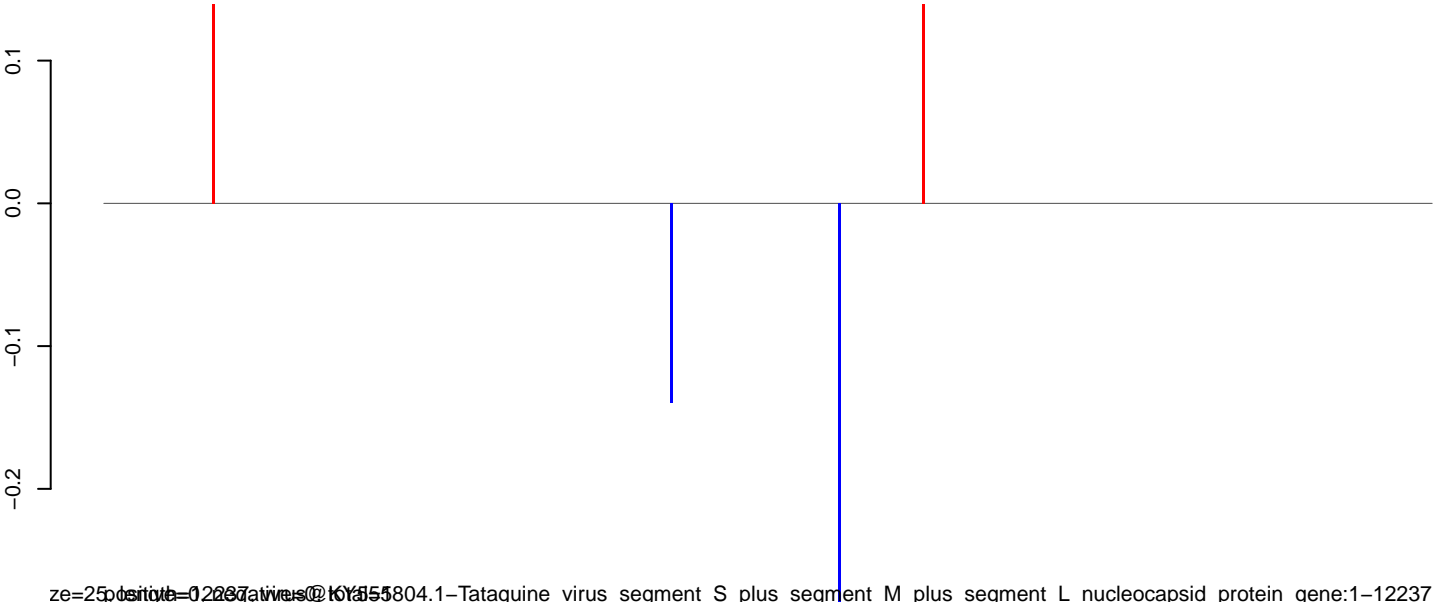
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



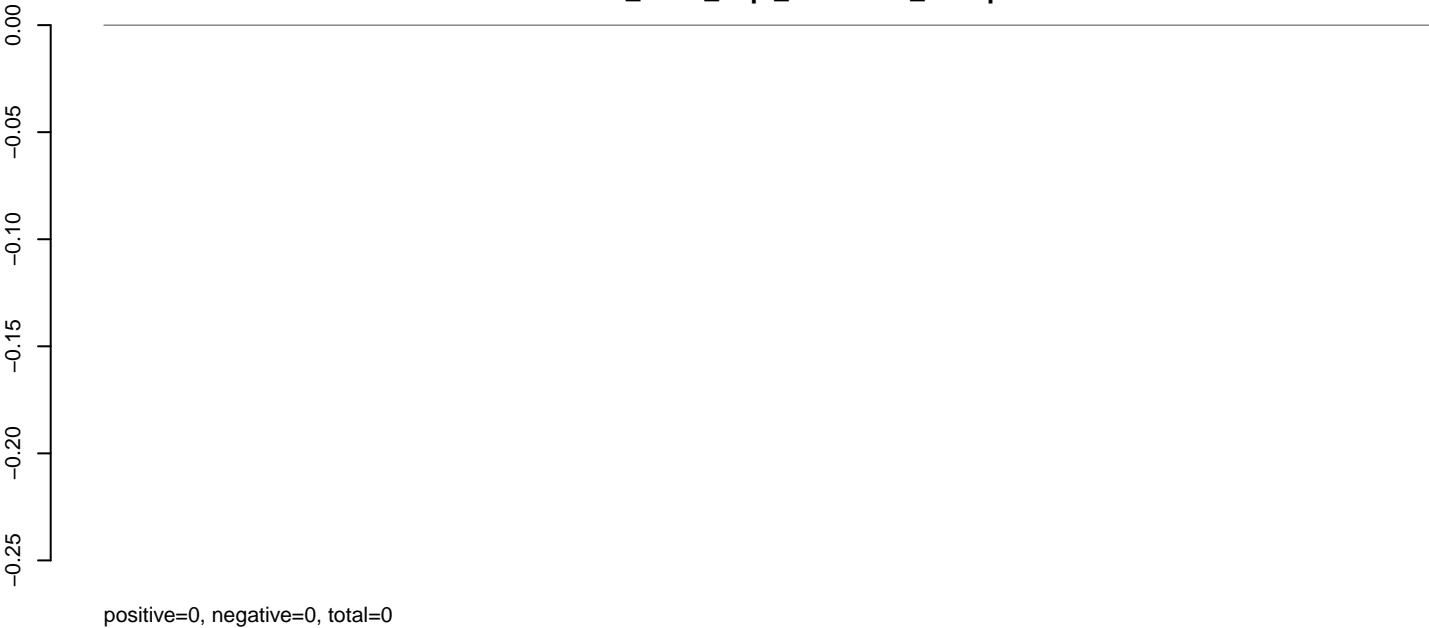
ze=25012237 positives@16155804.1-Tataguine_virus_segment_S_plus_segment_M_plus_segment_L_nucleocapsid_protein_gene:1-12237

0 2000 4000 6000 8000 10000 12000

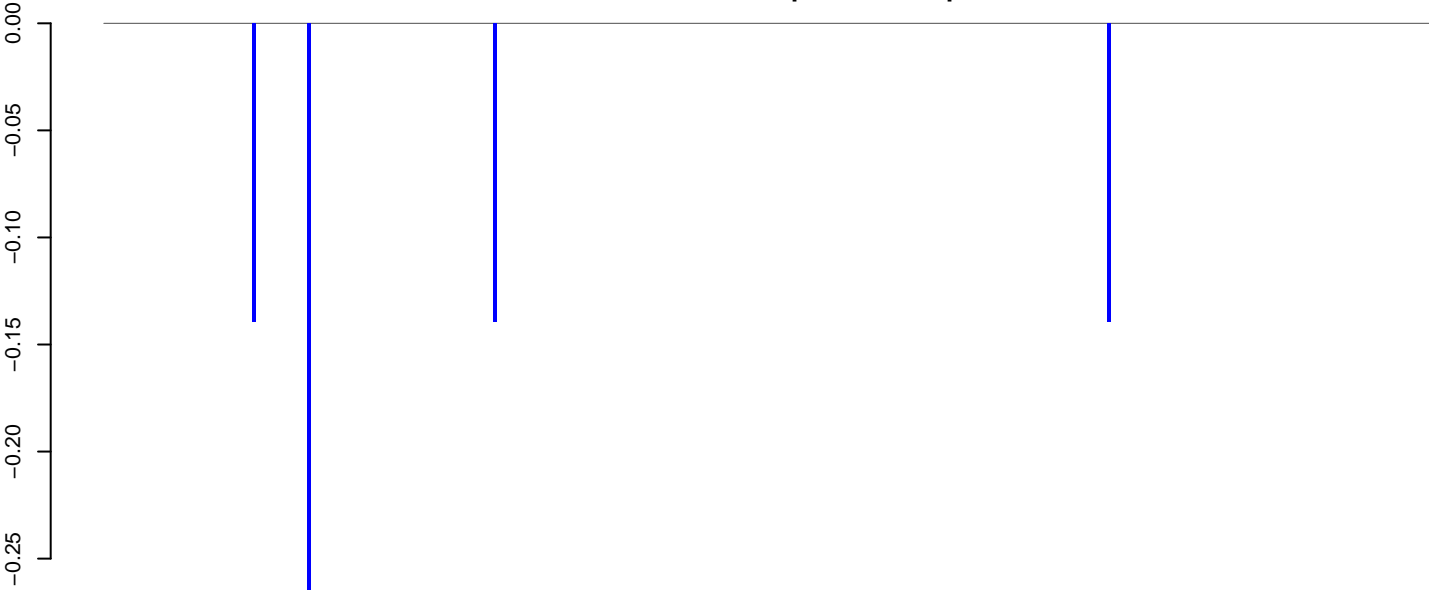
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



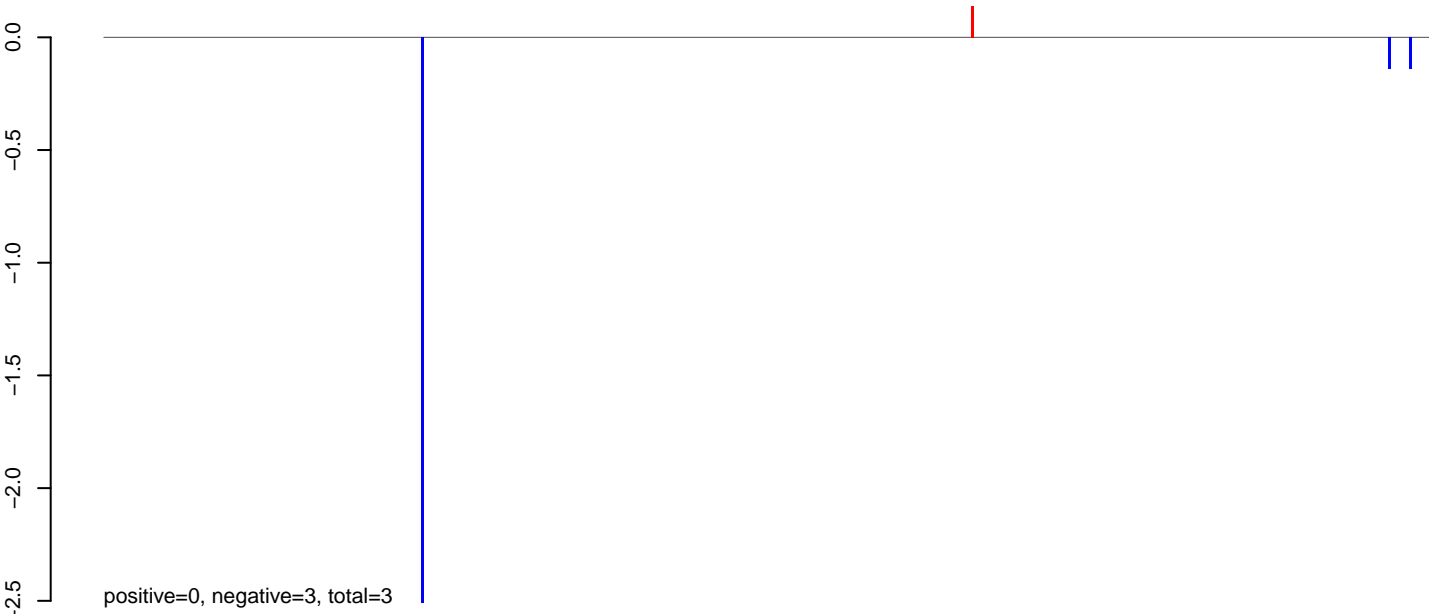
AeAlbo_C636_2dpi_mock3.rep



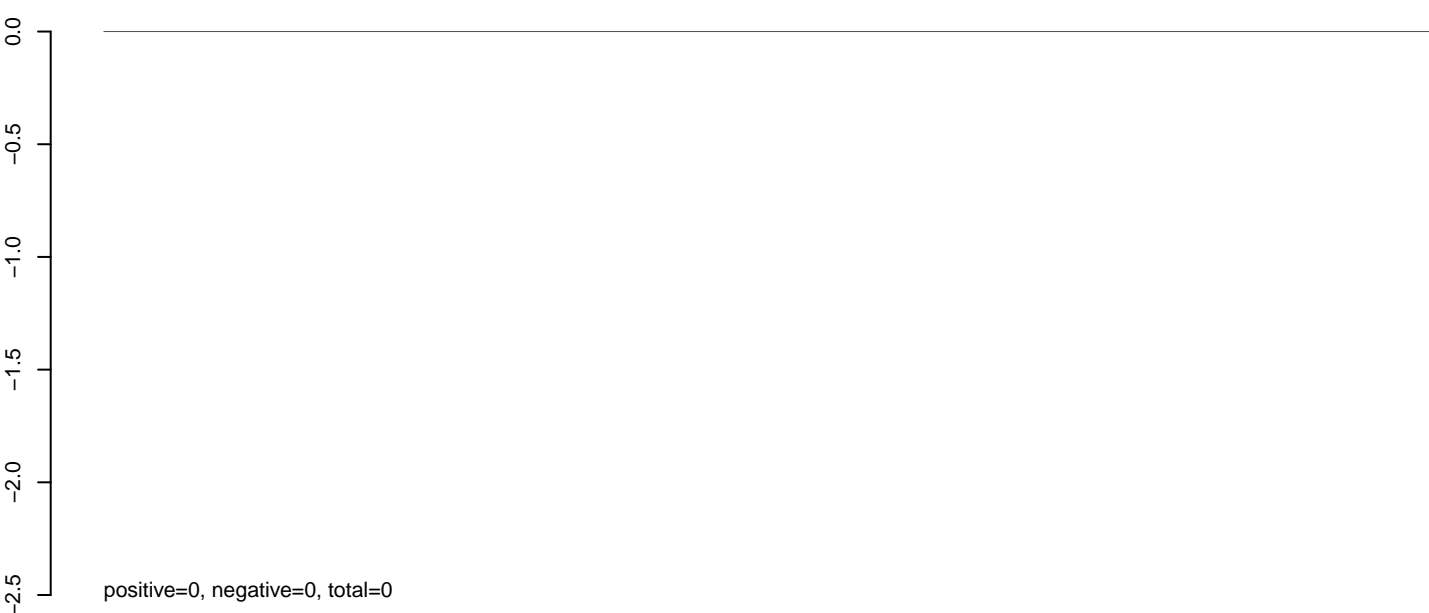
Window size=25, length=10749, virus@KX669682.1-Calbertado_virus_isolate_Cal_mos_2_polyprotein_gene:1-10749

0 2000 4000 6000 8000 10000

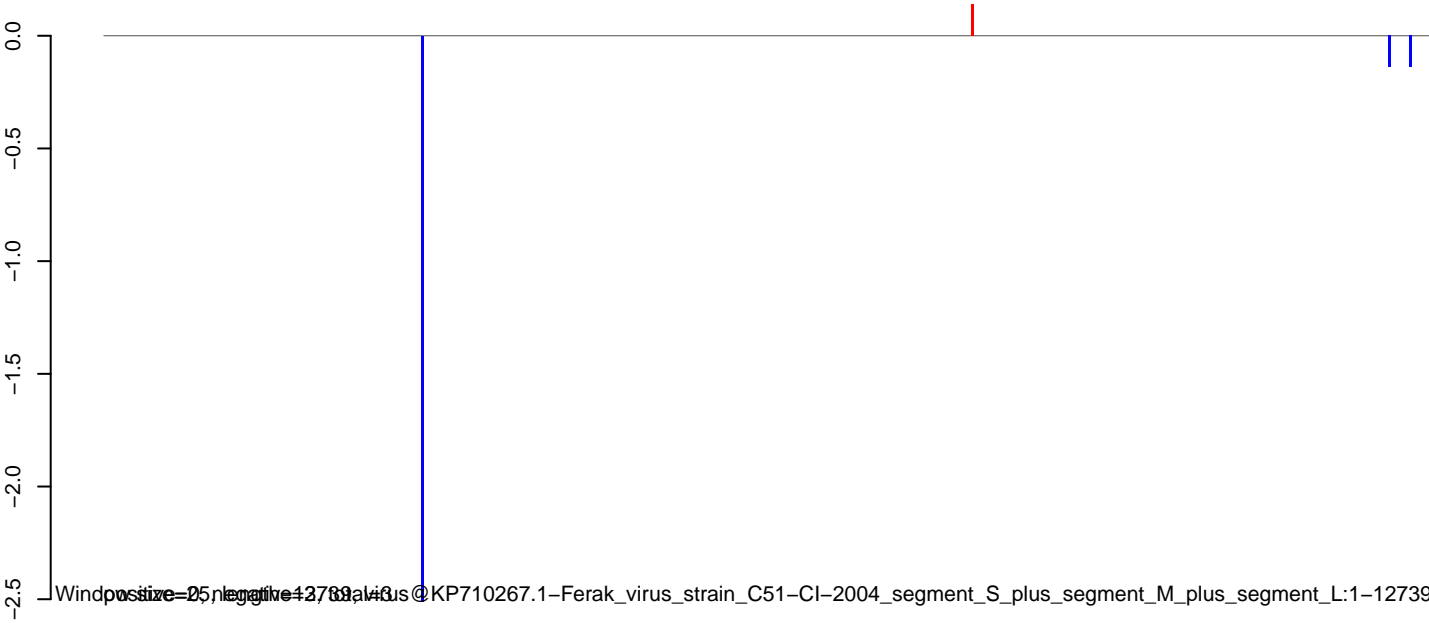
AeAlbo_C636_2dpi_mock3.18_23.rep

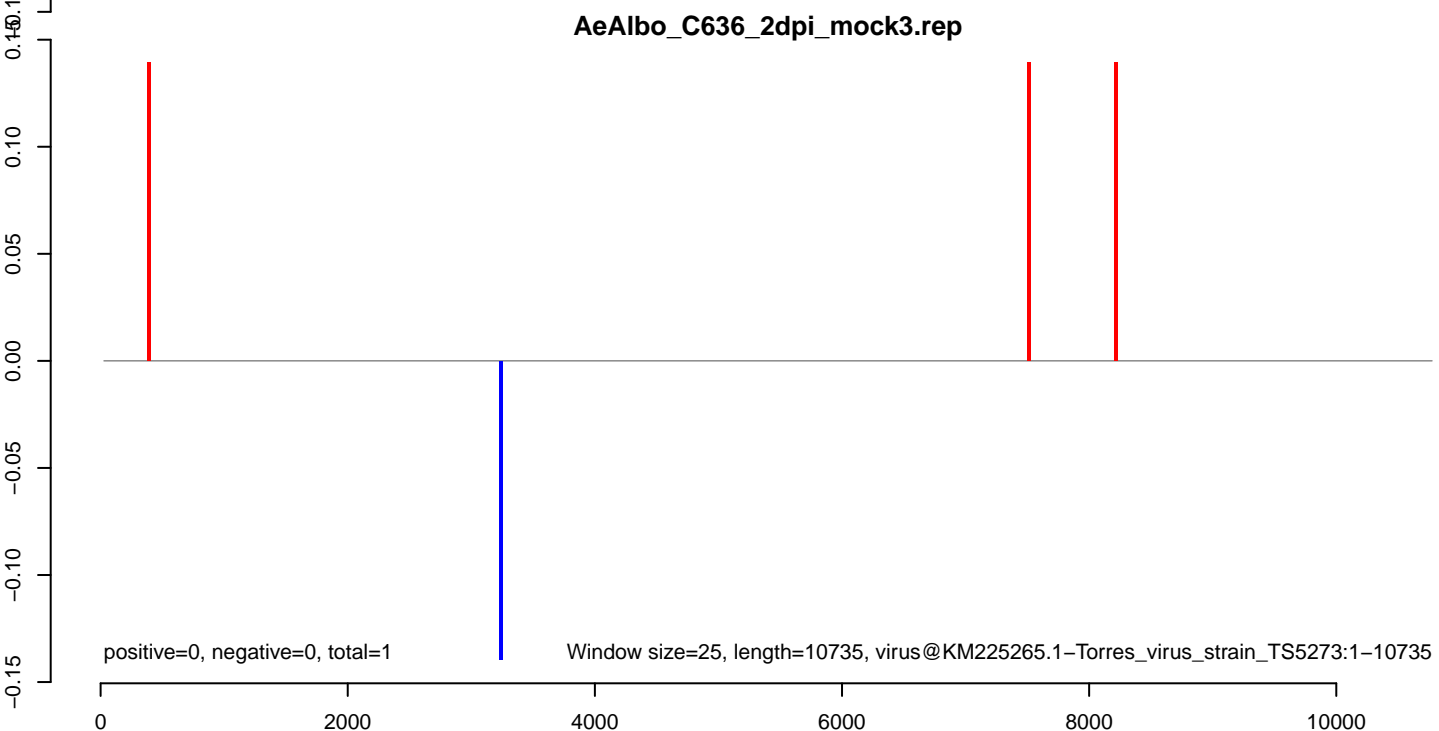
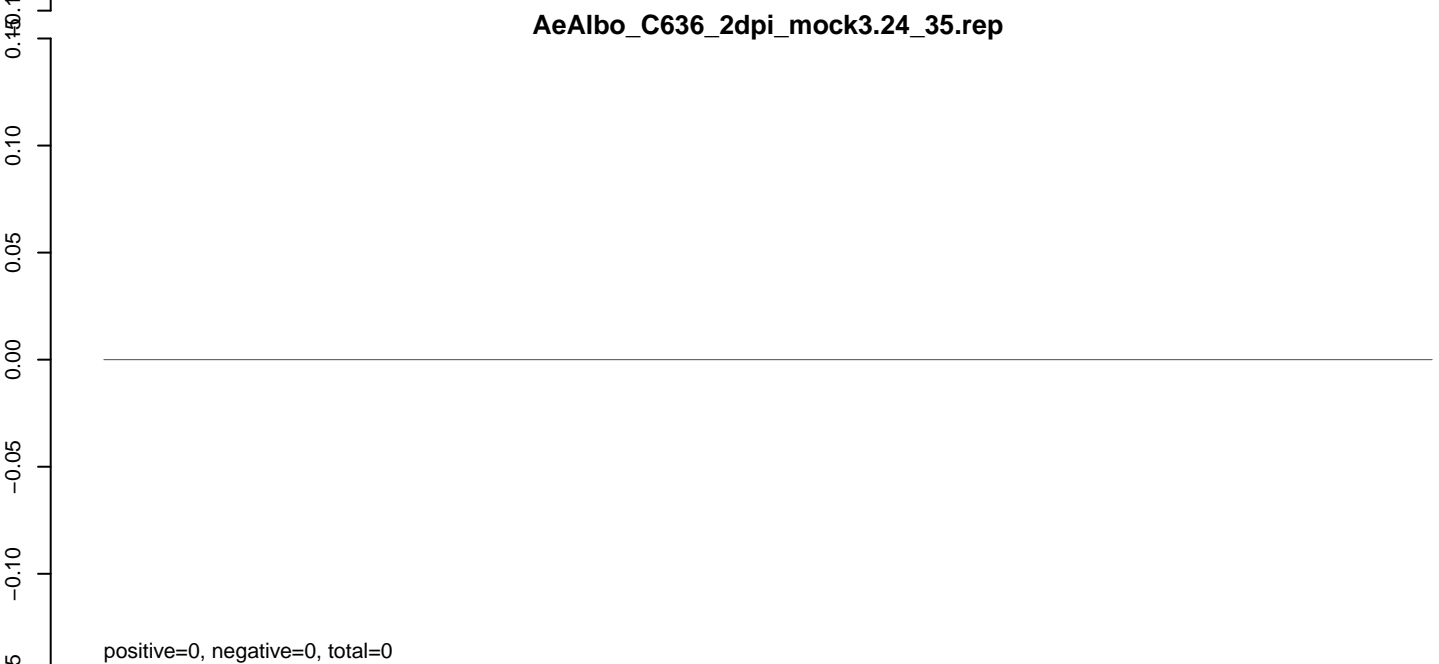
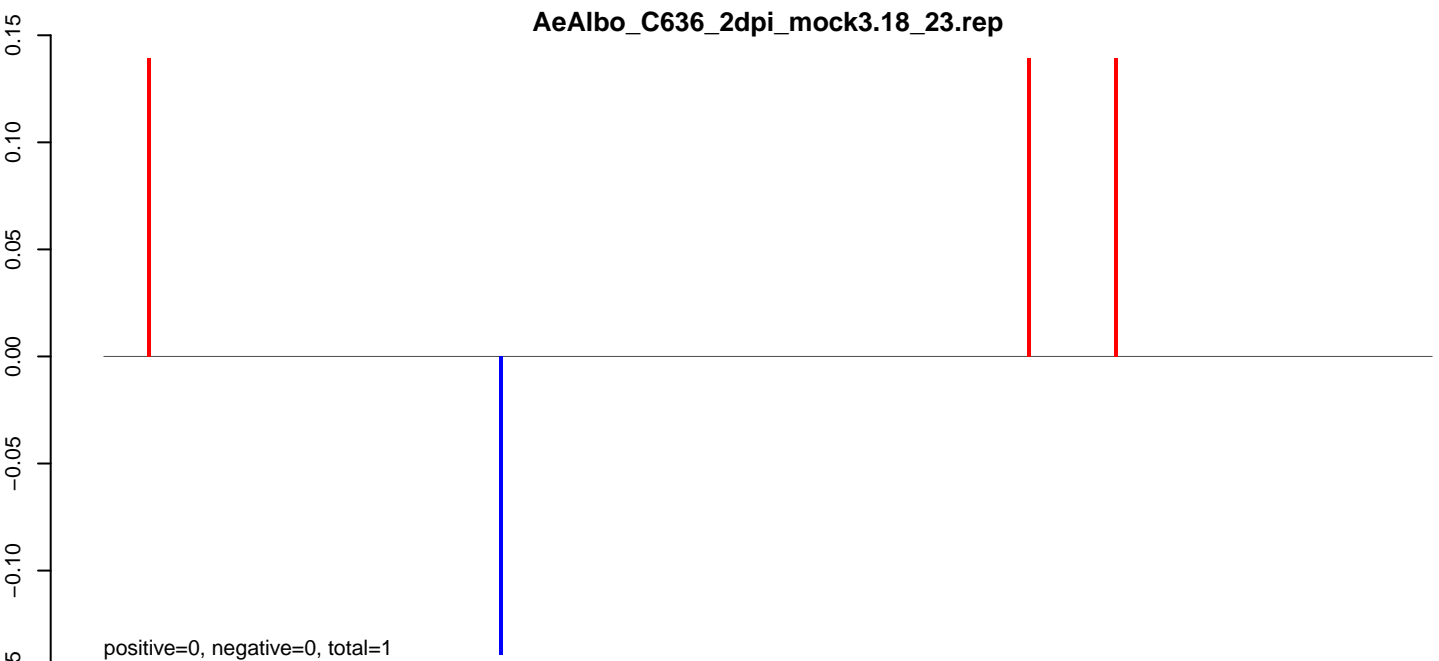


AeAlbo_C636_2dpi_mock3.24_35.rep

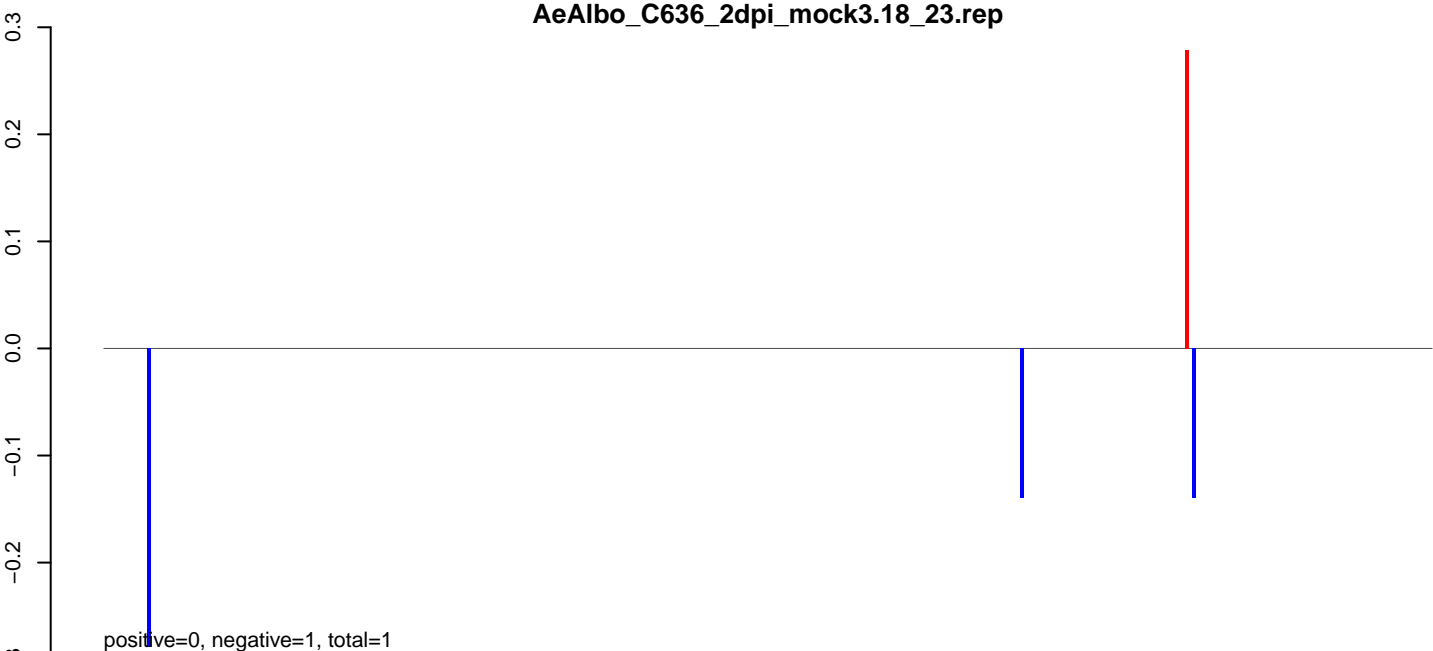


AeAlbo_C636_2dpi_mock3.rep



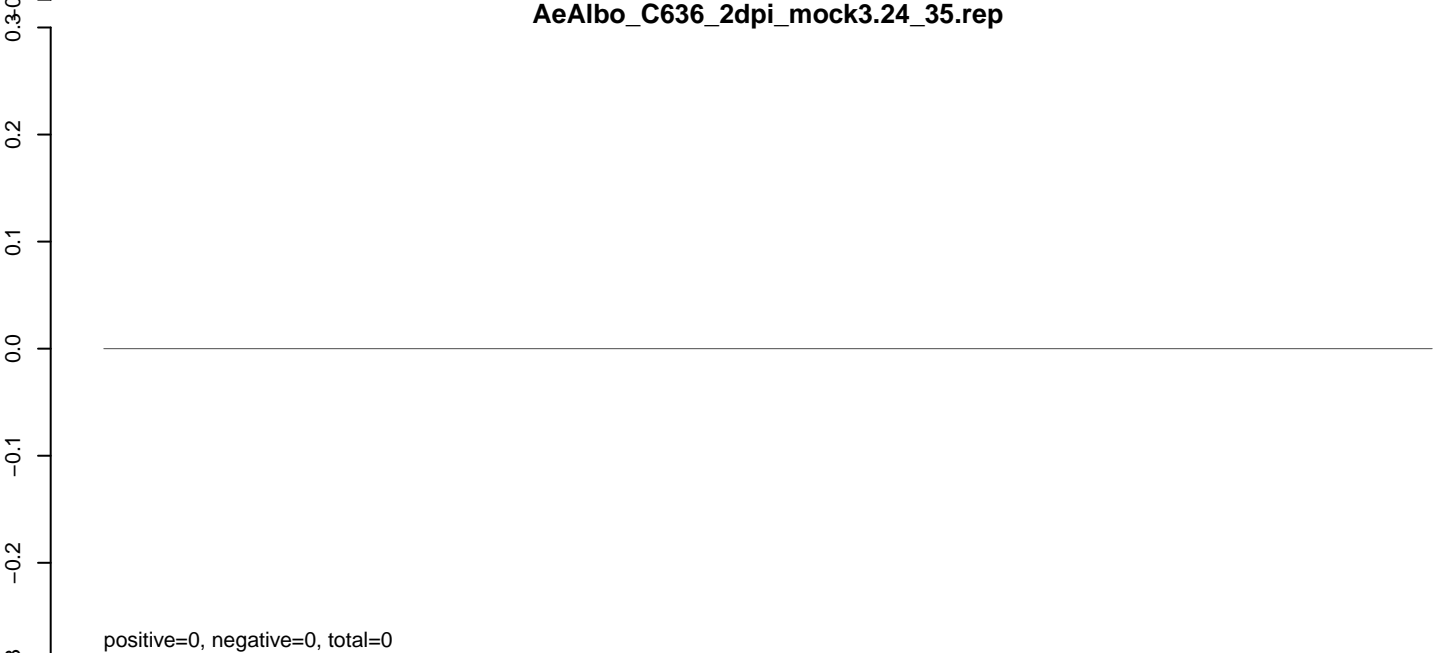


AeAlbo_C636_2dpi_mock3.18_23.rep



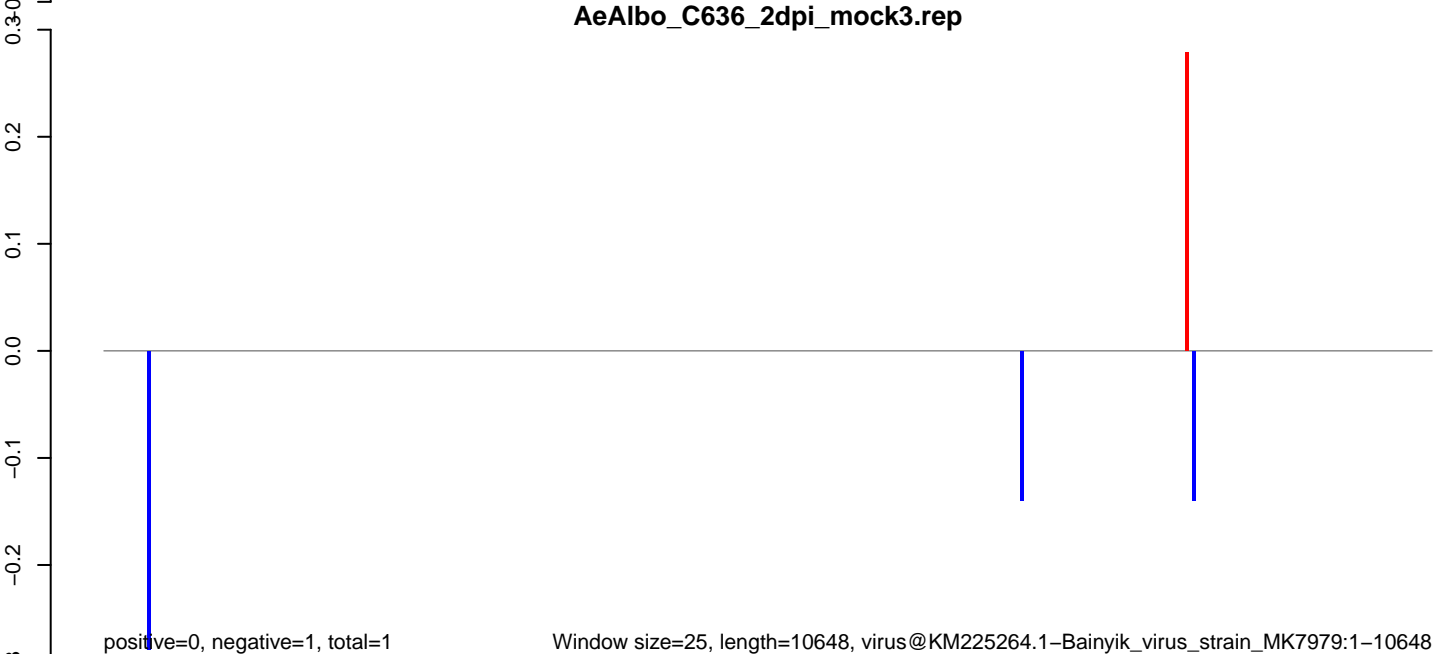
positive=0, negative=1, total=1

AeAlbo_C636_2dpi_mock3.24_35.rep



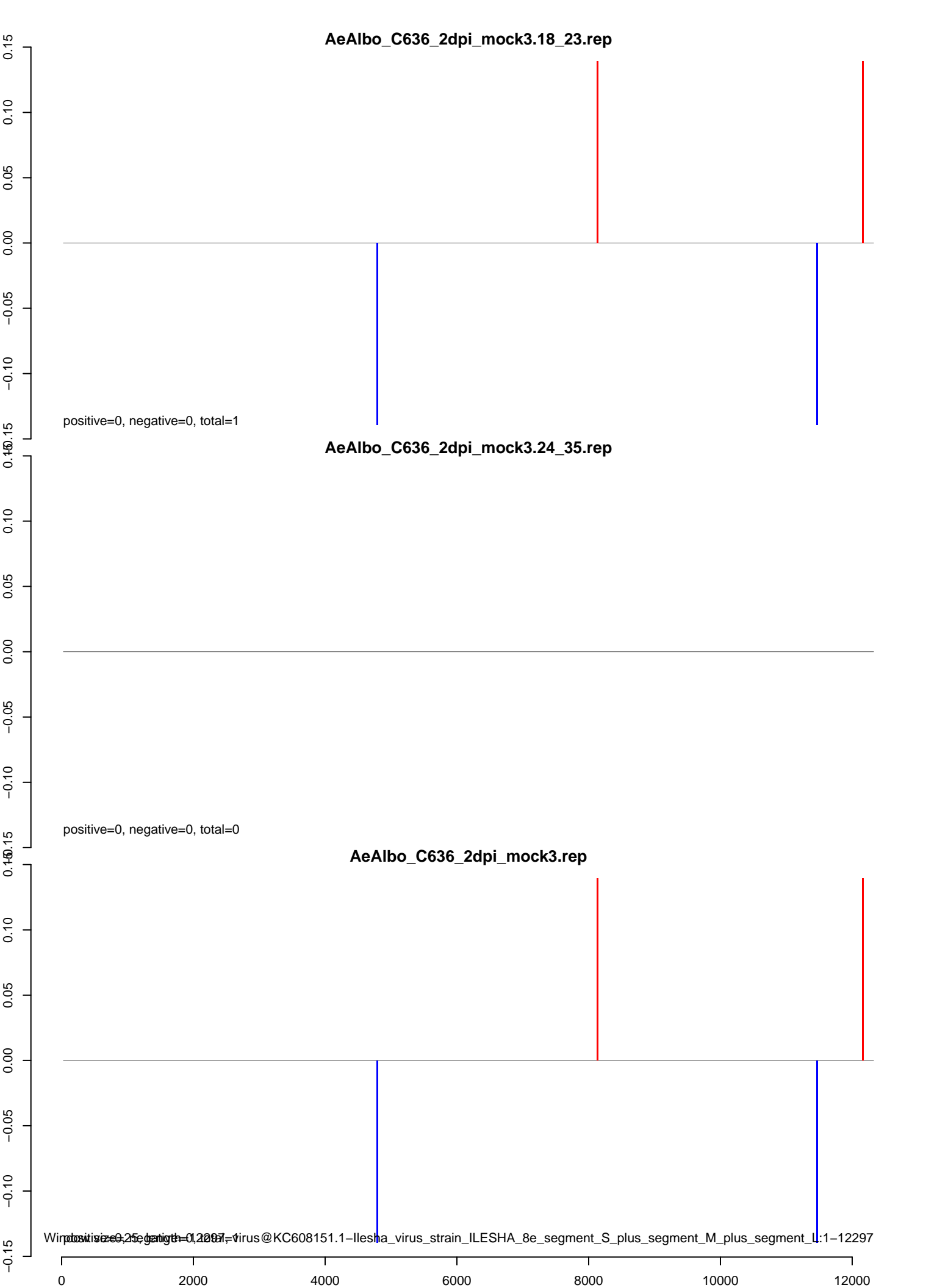
positive=0, negative=0, total=0

AeAlbo_C636_2dpi_mock3.rep

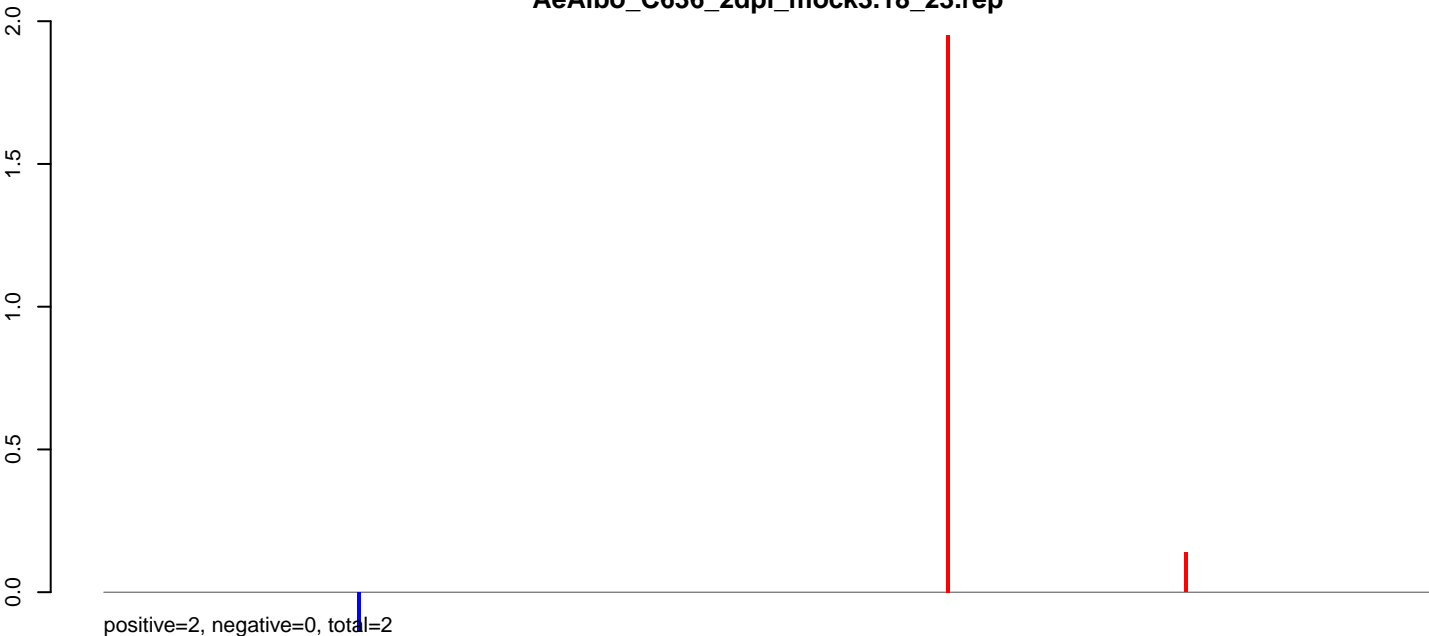


positive=0, negative=1, total=1

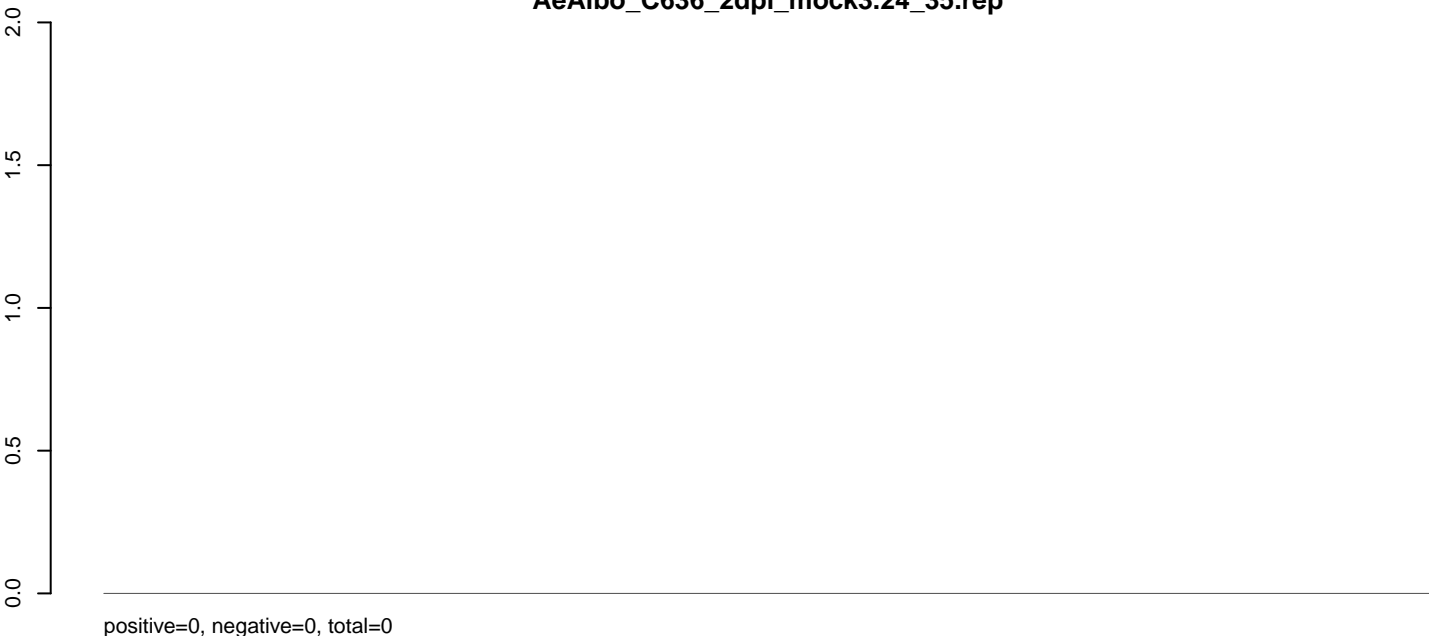
Window size=25, length=10648, virus@KM225264.1-Bainyik_virus_strain_MK7979:1-10648



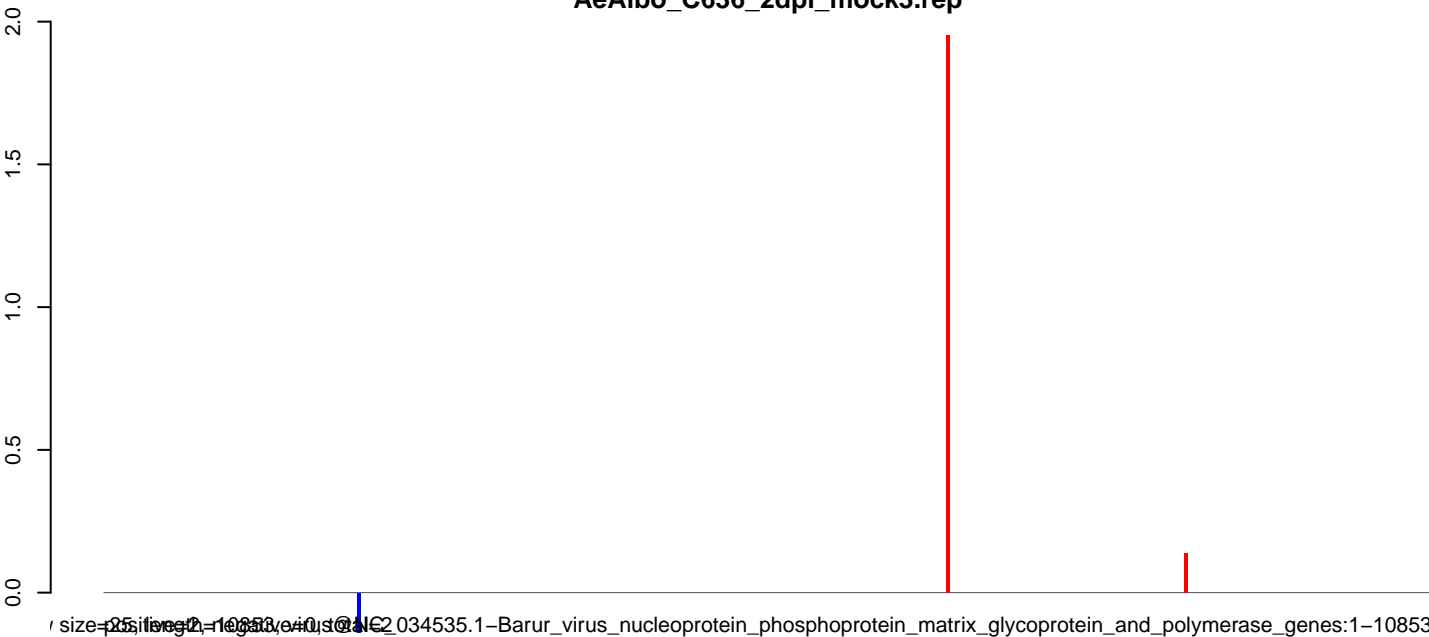
AeAlbo_C636_2dpi_mock3.18_23.rep



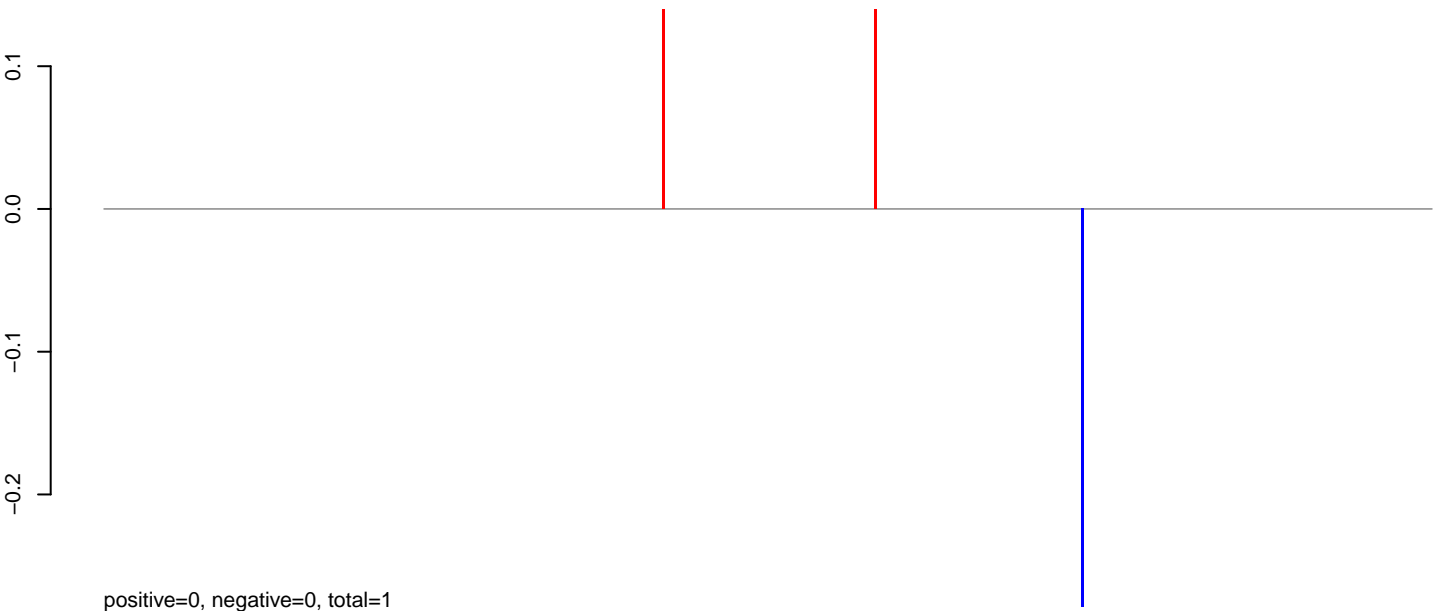
AeAlbo_C636_2dpi_mock3.24_35.rep



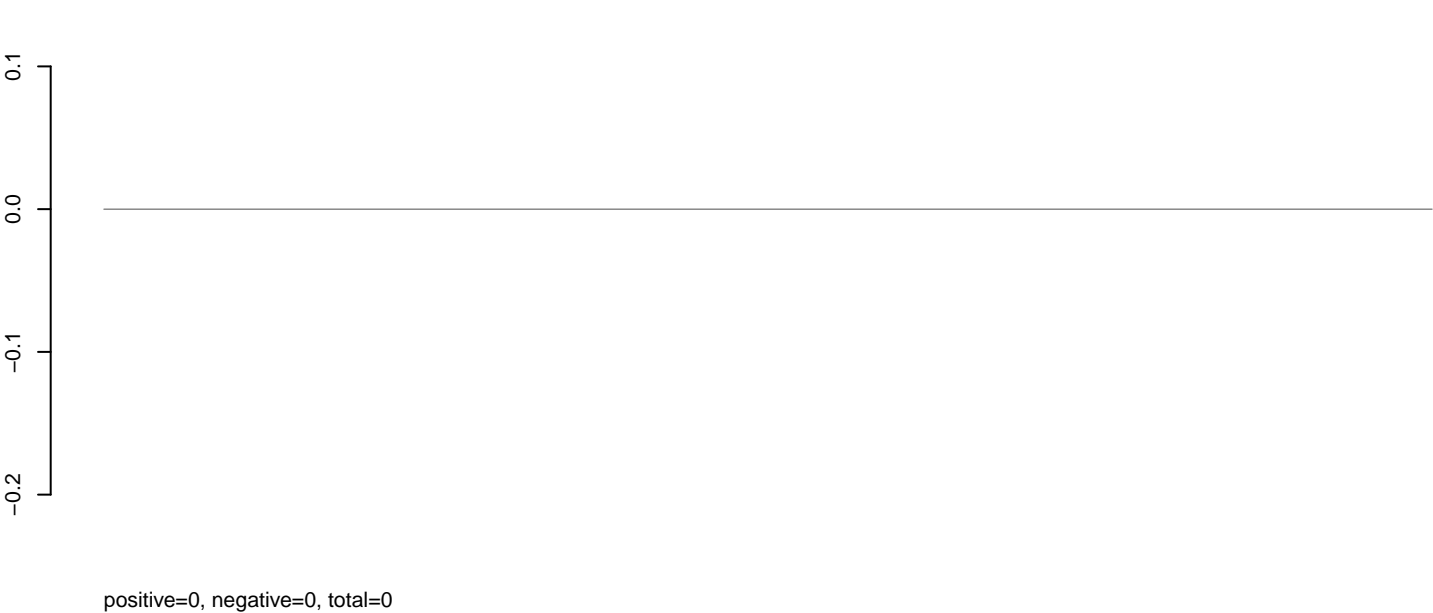
AeAlbo_C636_2dpi_mock3.rep



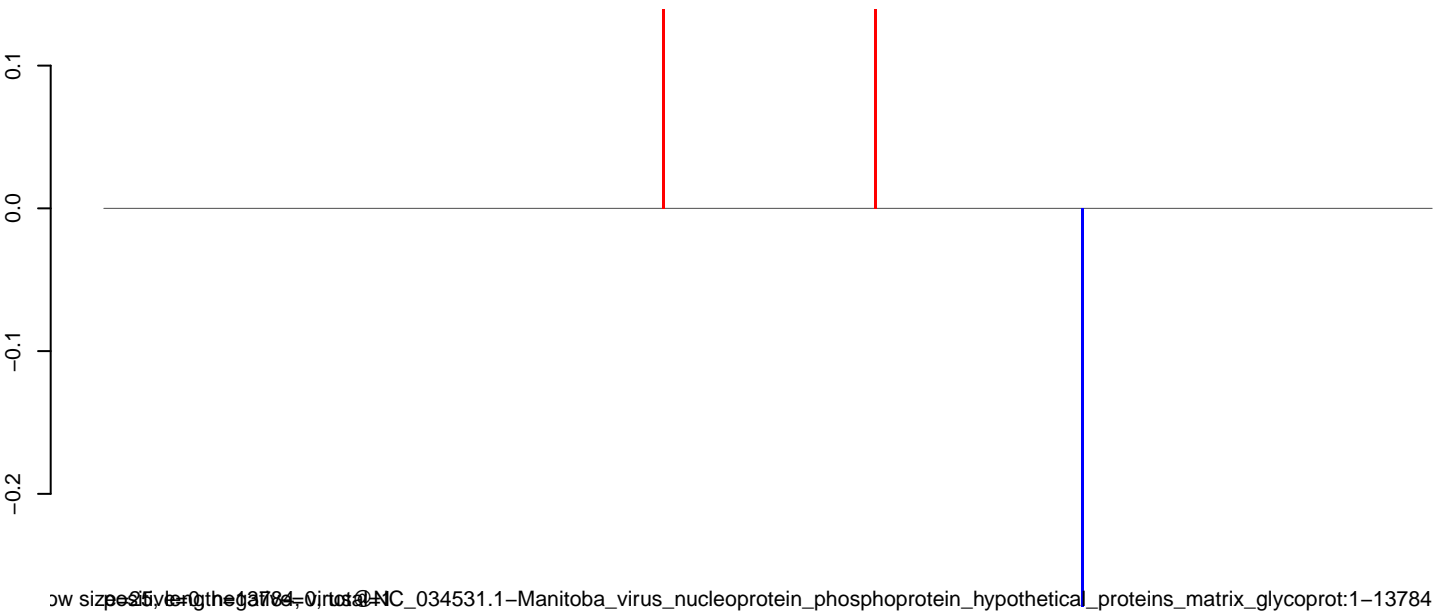
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



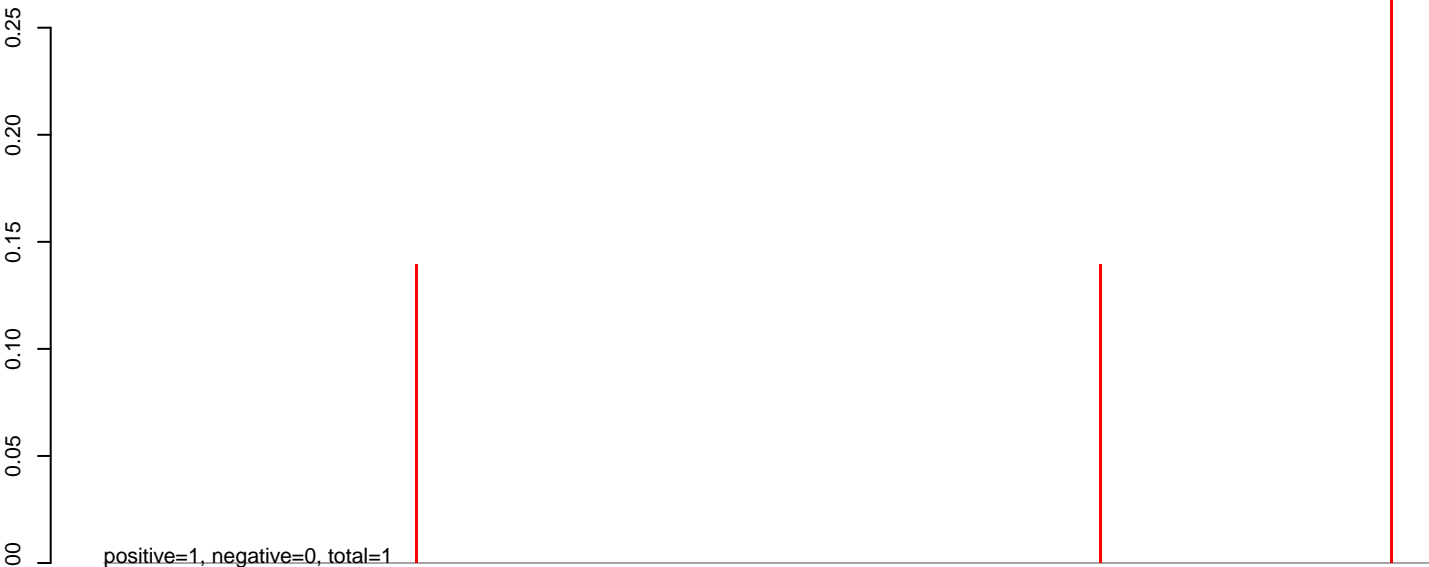
AeAlbo_C636_2dpi_mock3.rep



NC_034531.1-Manitoba_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycoprot:1-13784

0 2000 4000 6000 8000 10000 12000 14000

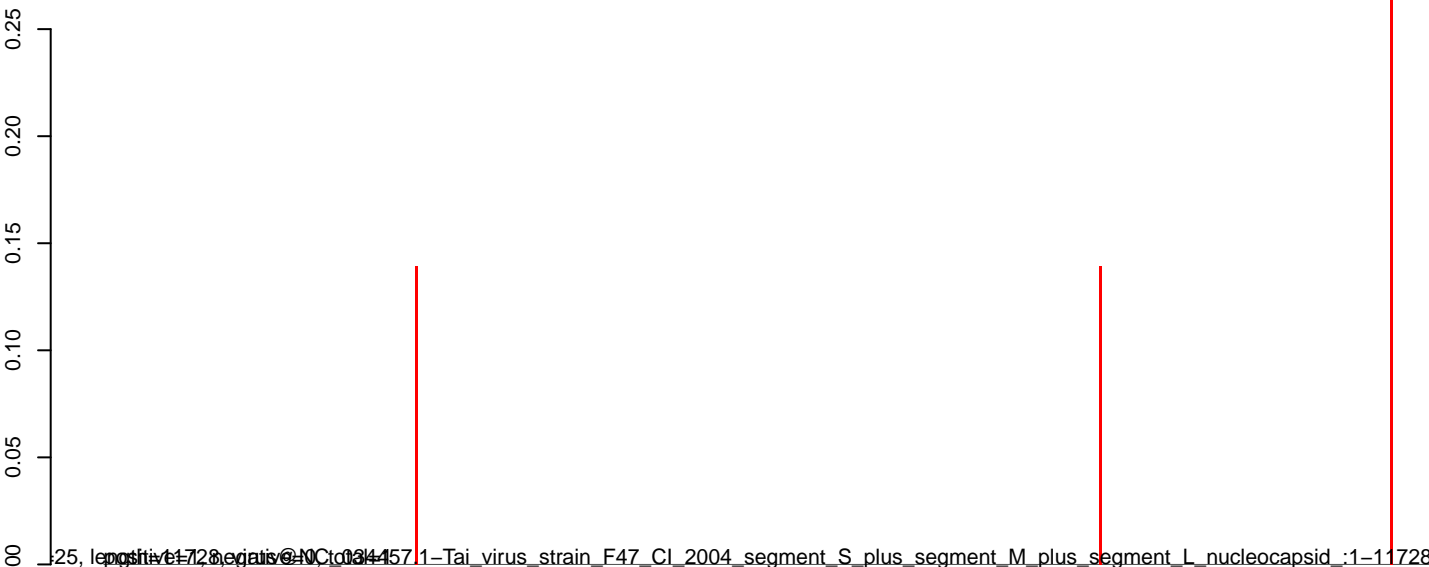
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



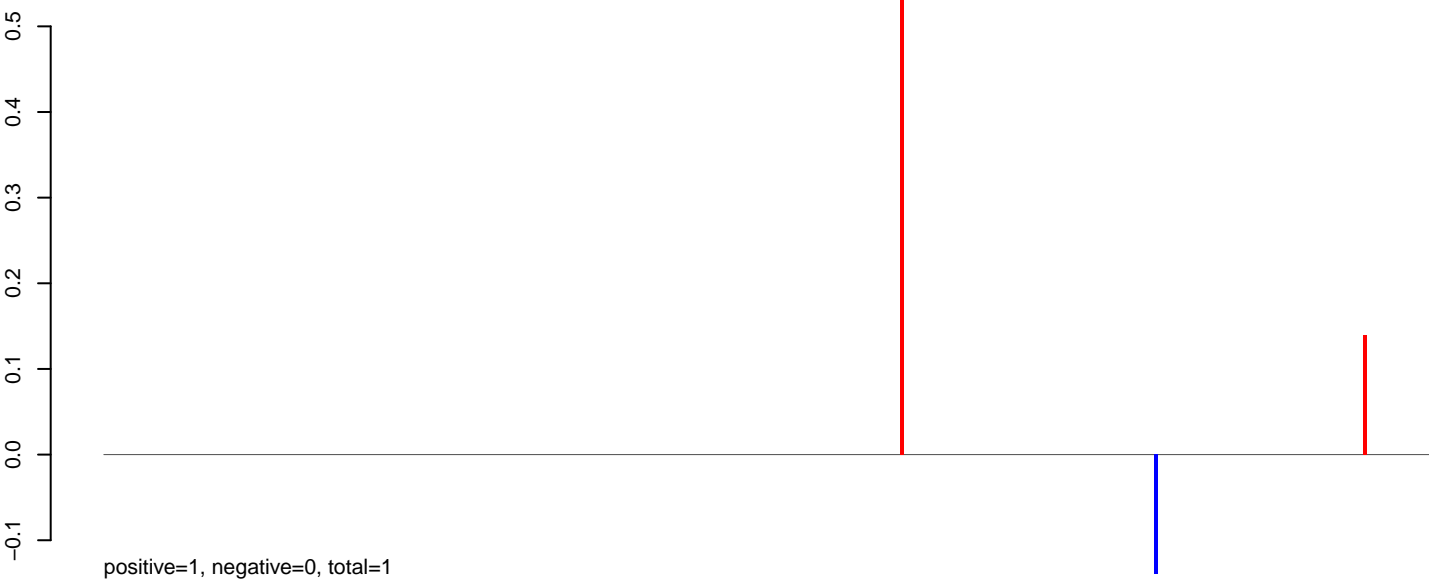
AeAlbo_C636_2dpi_mock3.rep



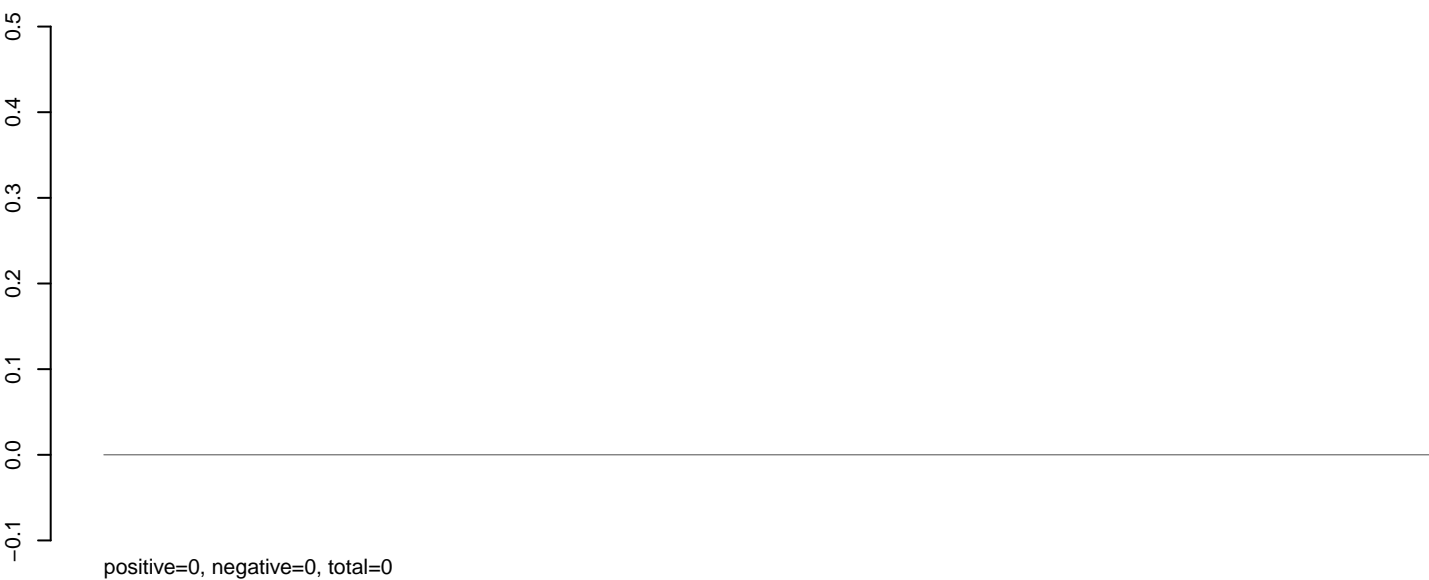
25, length=11728, negative=0, total=1, 1-Tai_virus_strain_F47_CI_2004_segment_S_plus_segment_M_plus_segment_L_nucleocapsid :1-11728

0 2000 4000 6000 8000 10000 12000

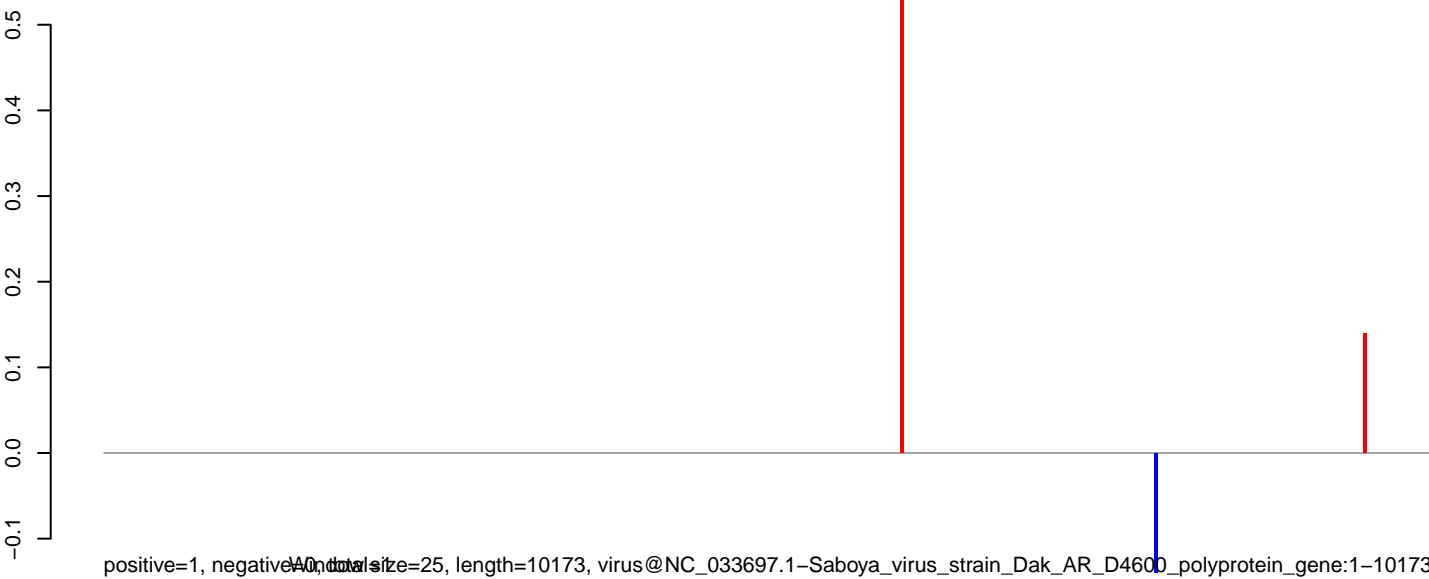
AeAlbo_C636_2dpi_mock3.18_23.rep



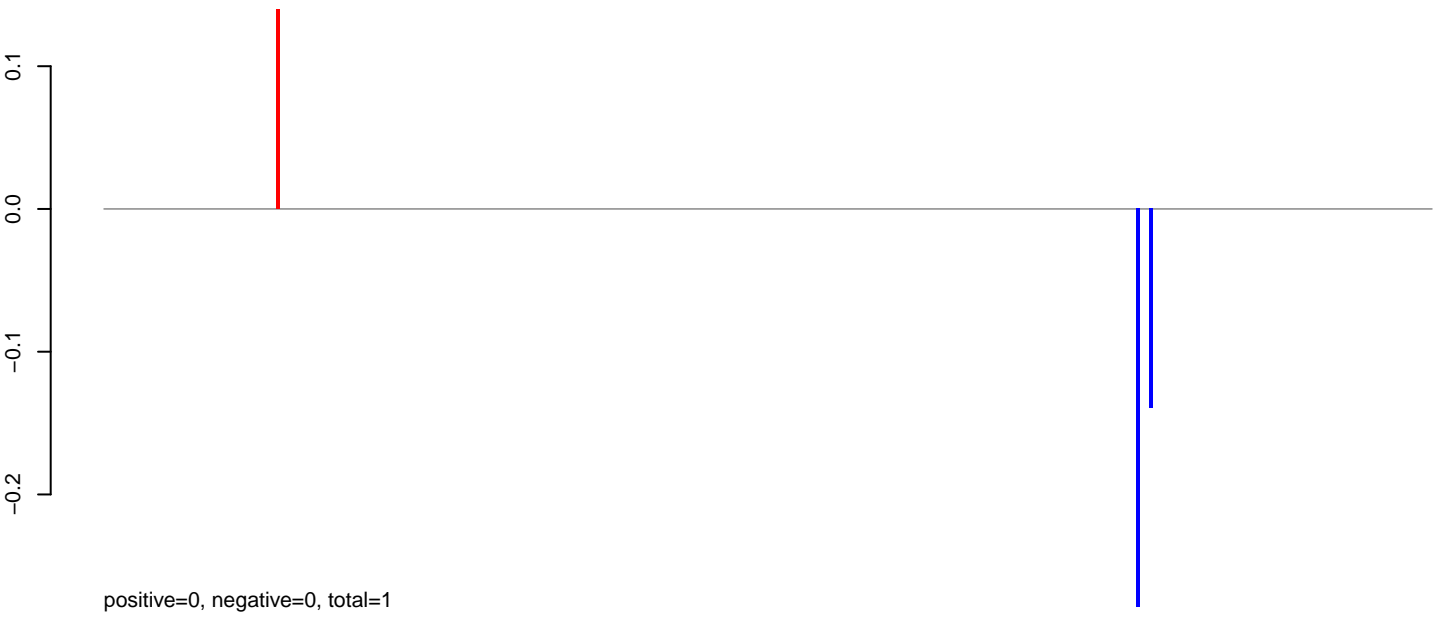
AeAlbo_C636_2dpi_mock3.24_35.rep



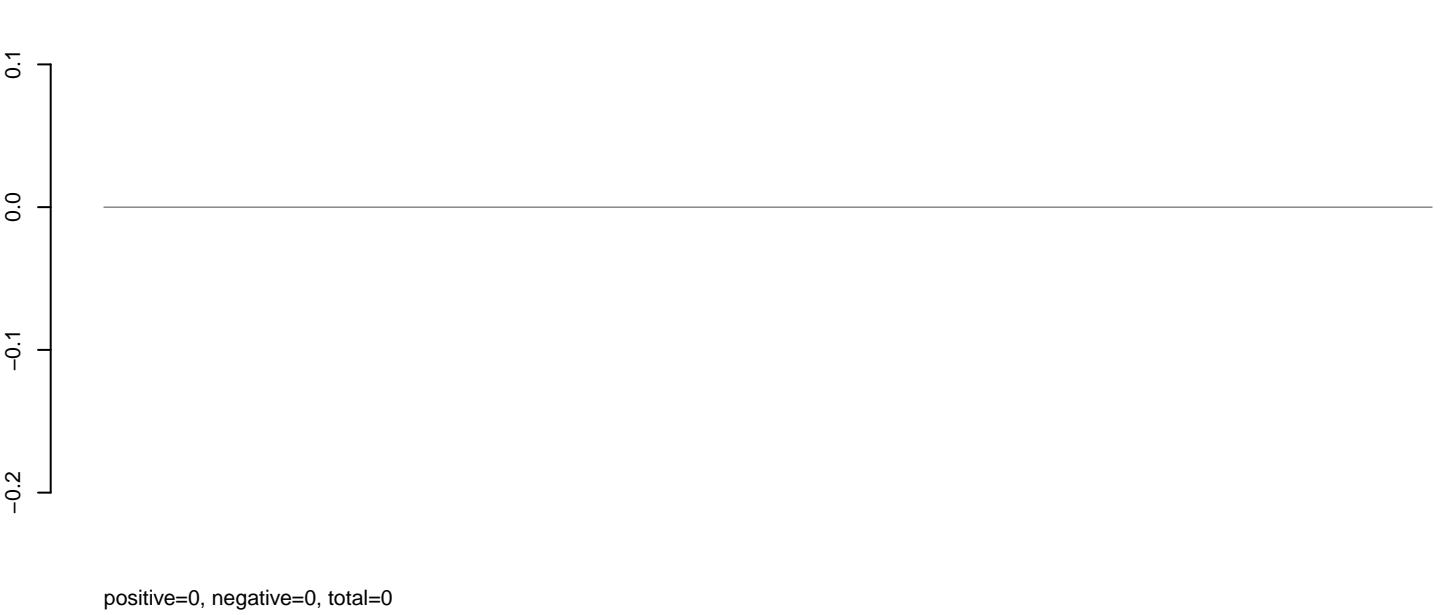
AeAlbo_C636_2dpi_mock3.rep



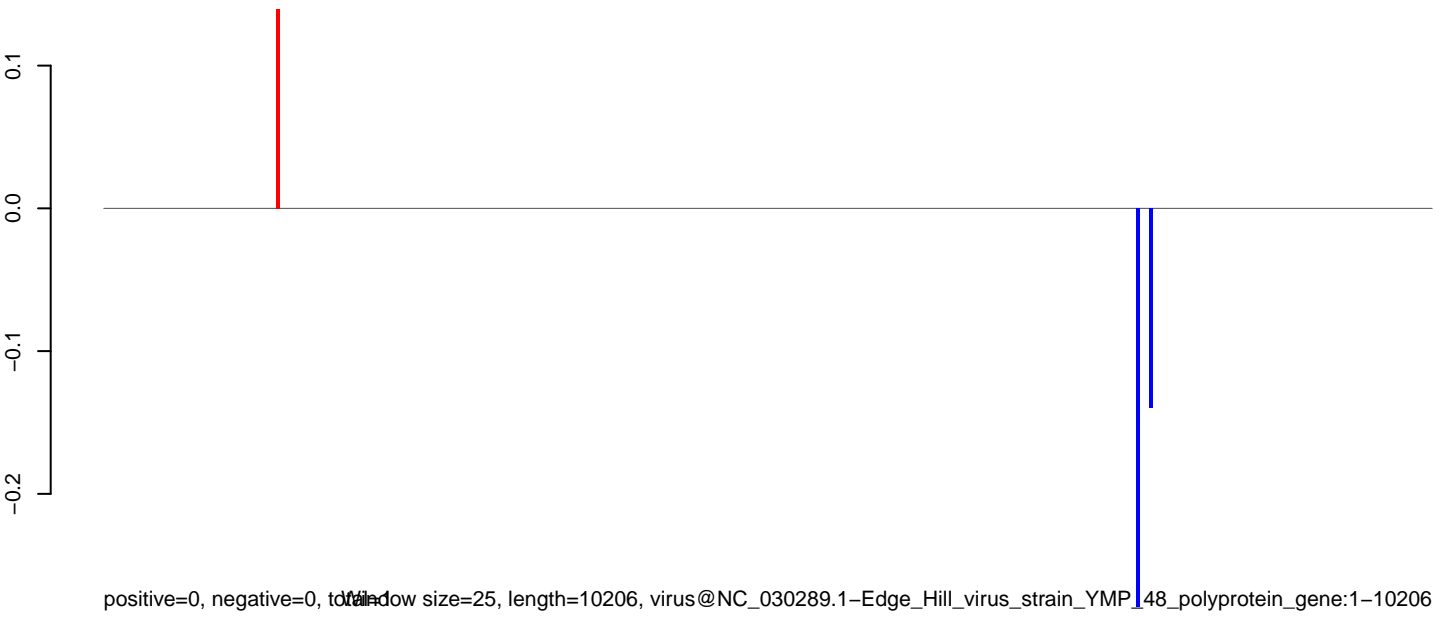
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

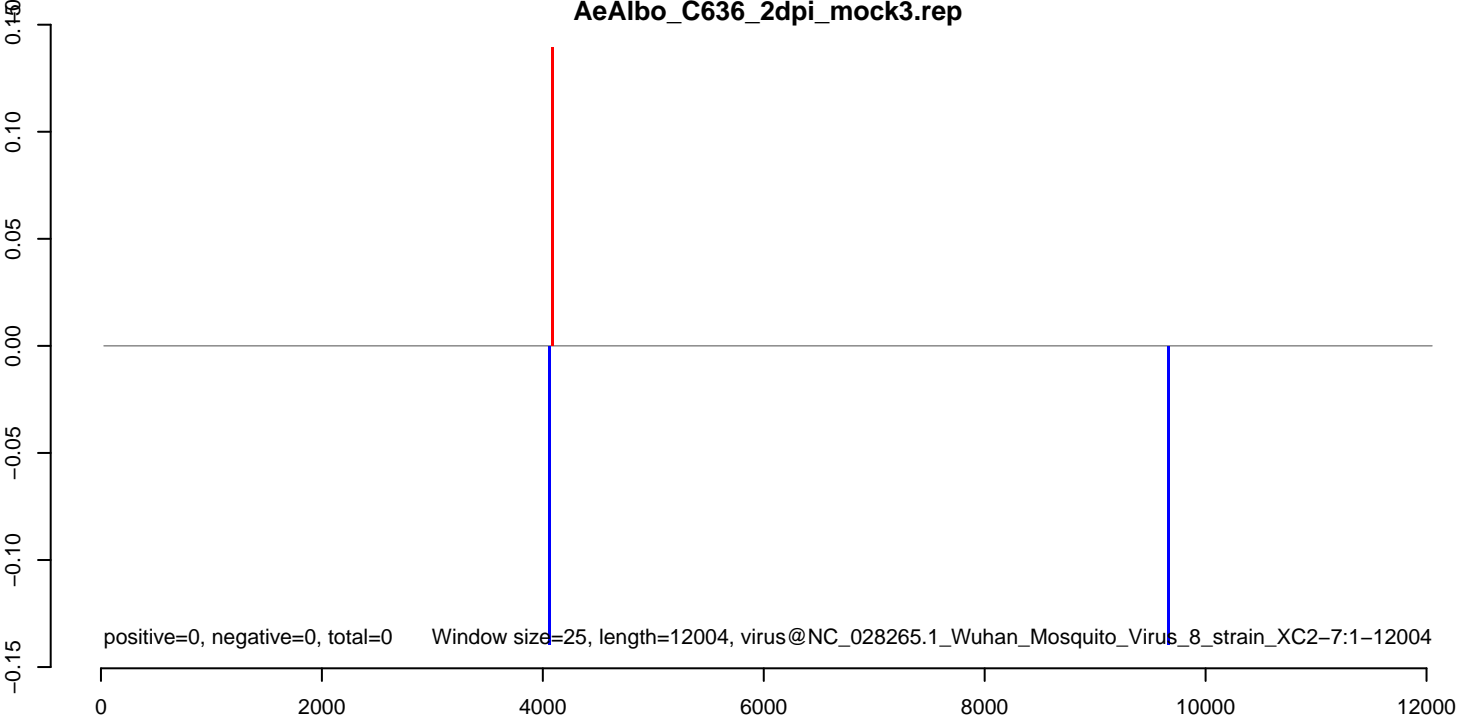
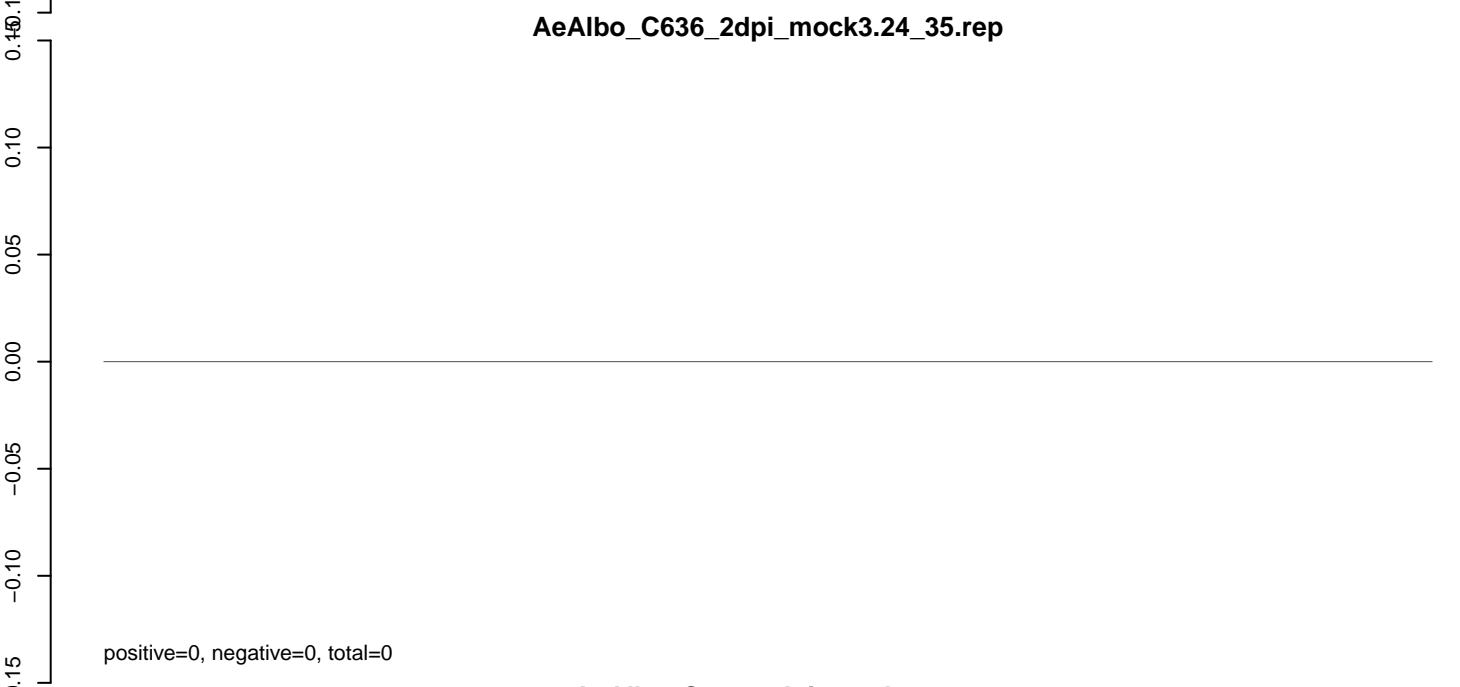
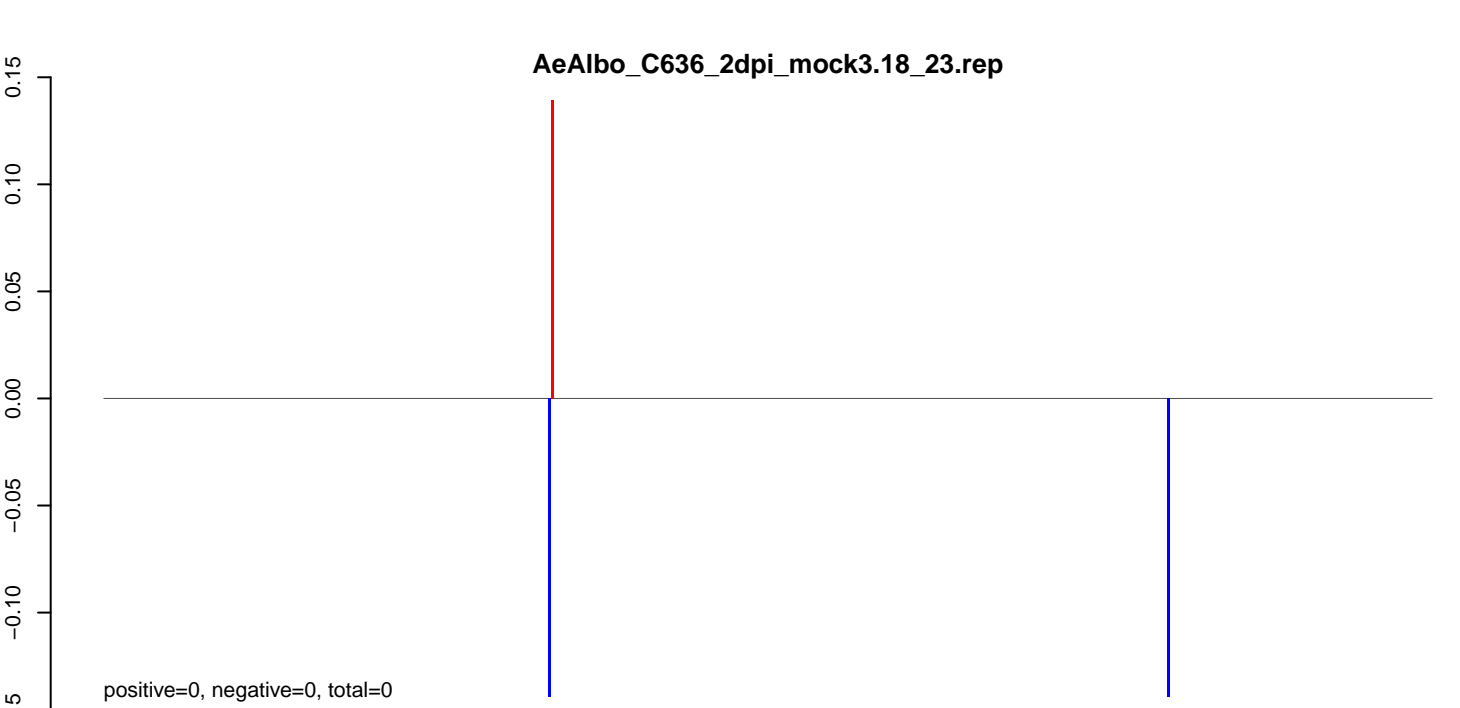


AeAlbo_C636_2dpi_mock3.rep

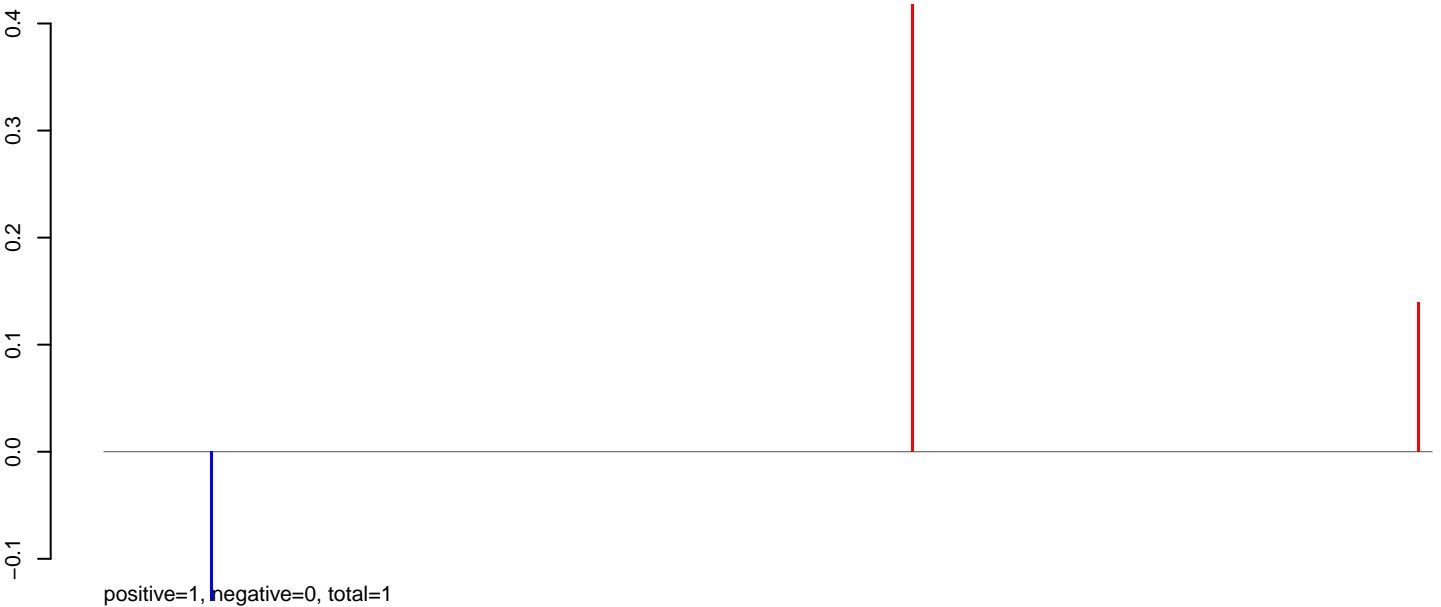


positive=0, negative=0, total=1 window size=25, length=10206, virus@NC_030289.1-Edge_Hill_virus_strain_YMP_148_polyprotein_gene:1-10206

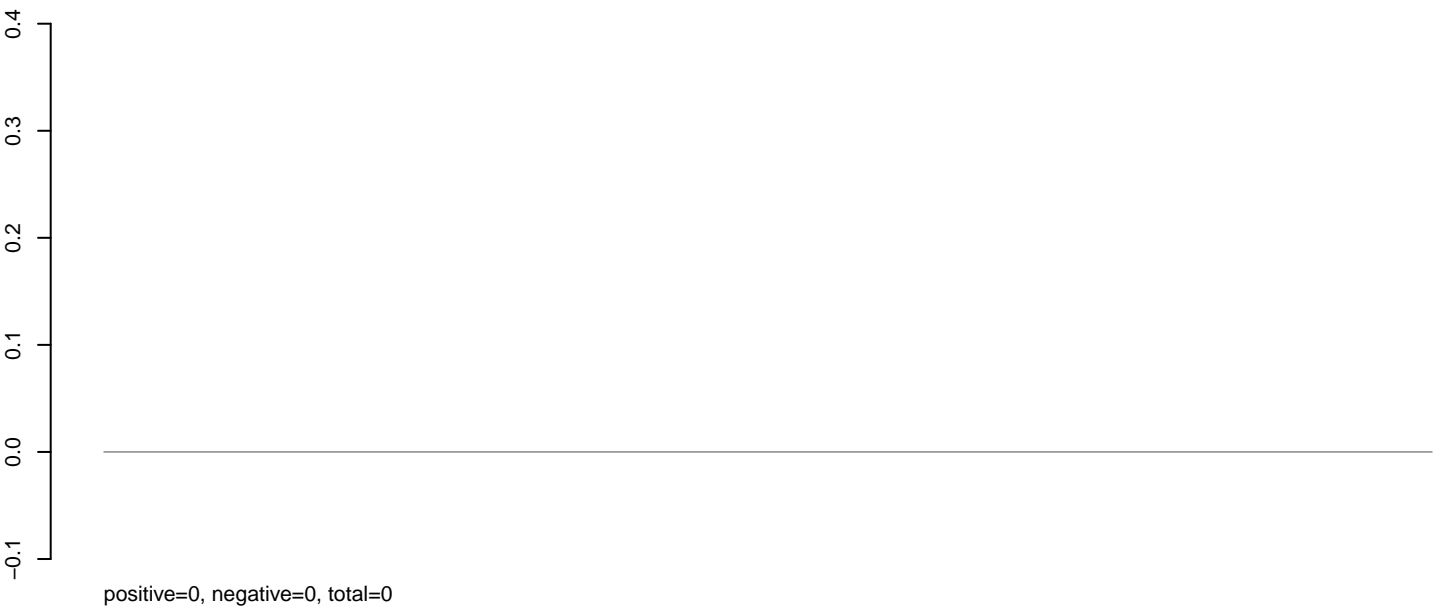
0 2000 4000 6000 8000 10000



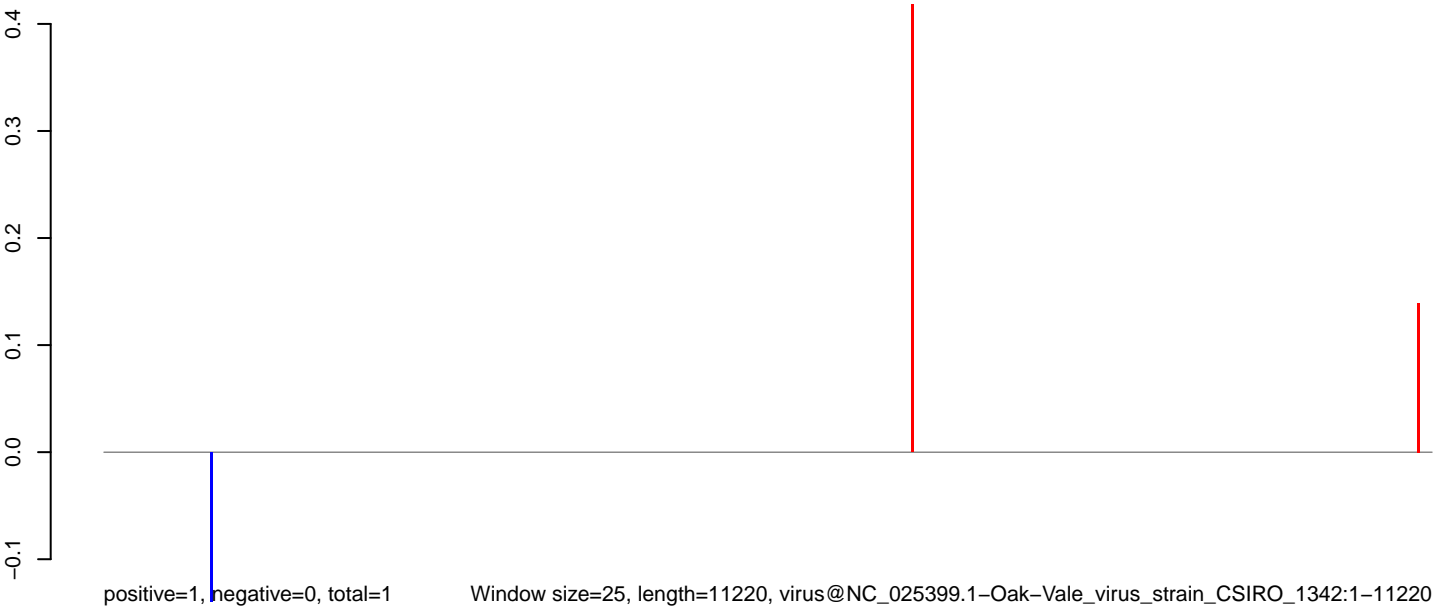
AeAlbo_C636_2dpi_mock3.18_23.rep



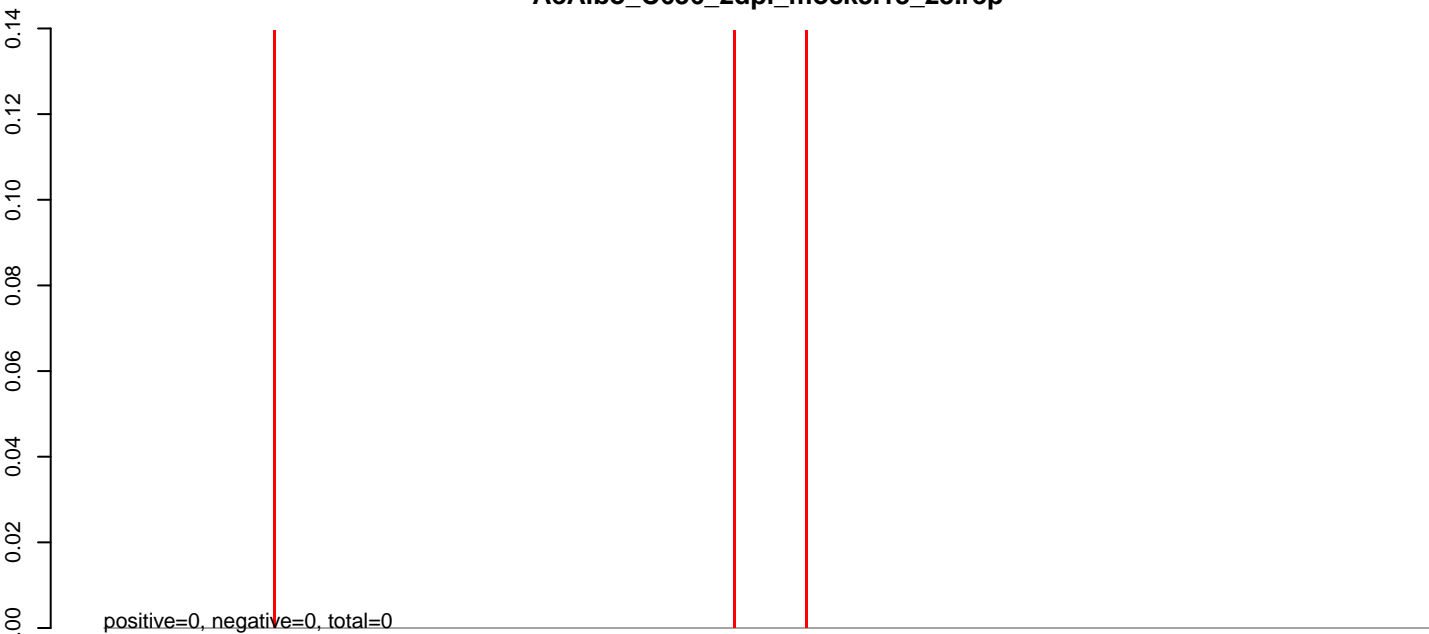
AeAlbo_C636_2dpi_mock3.24_35.rep



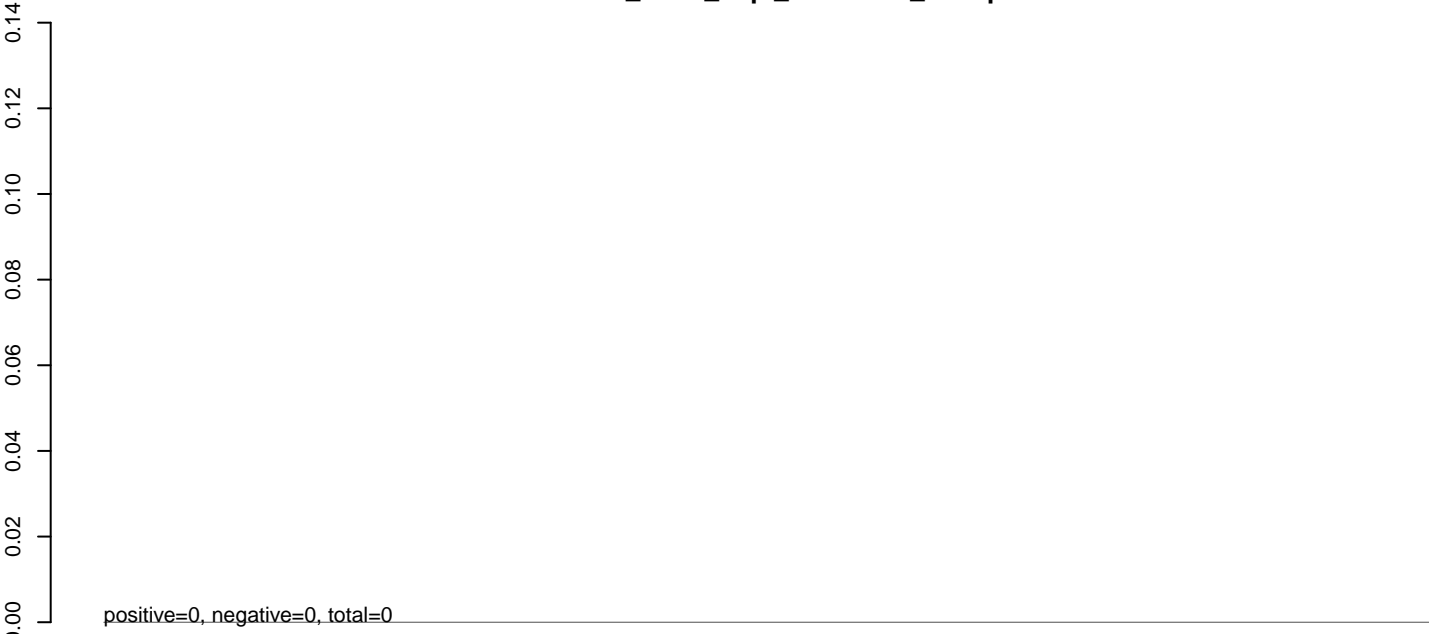
AeAlbo_C636_2dpi_mock3.rep



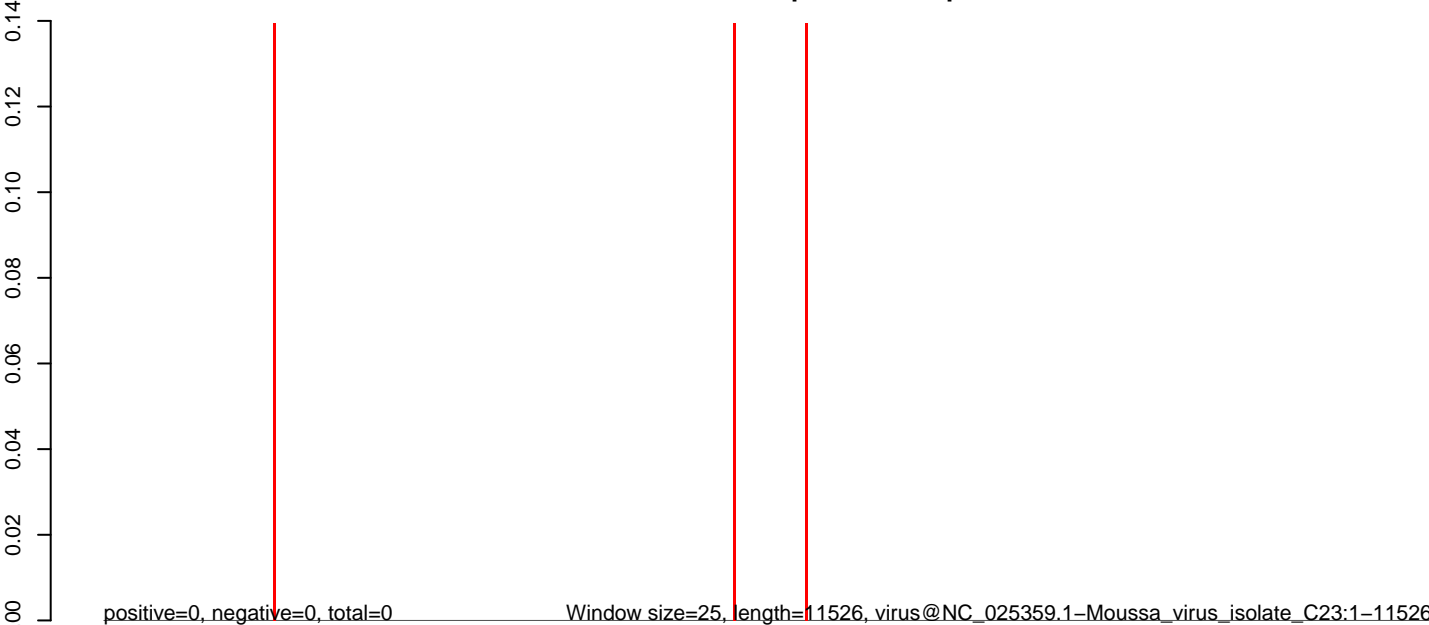
AeAlbo_C636_2dpi_mock3.18_23.rep



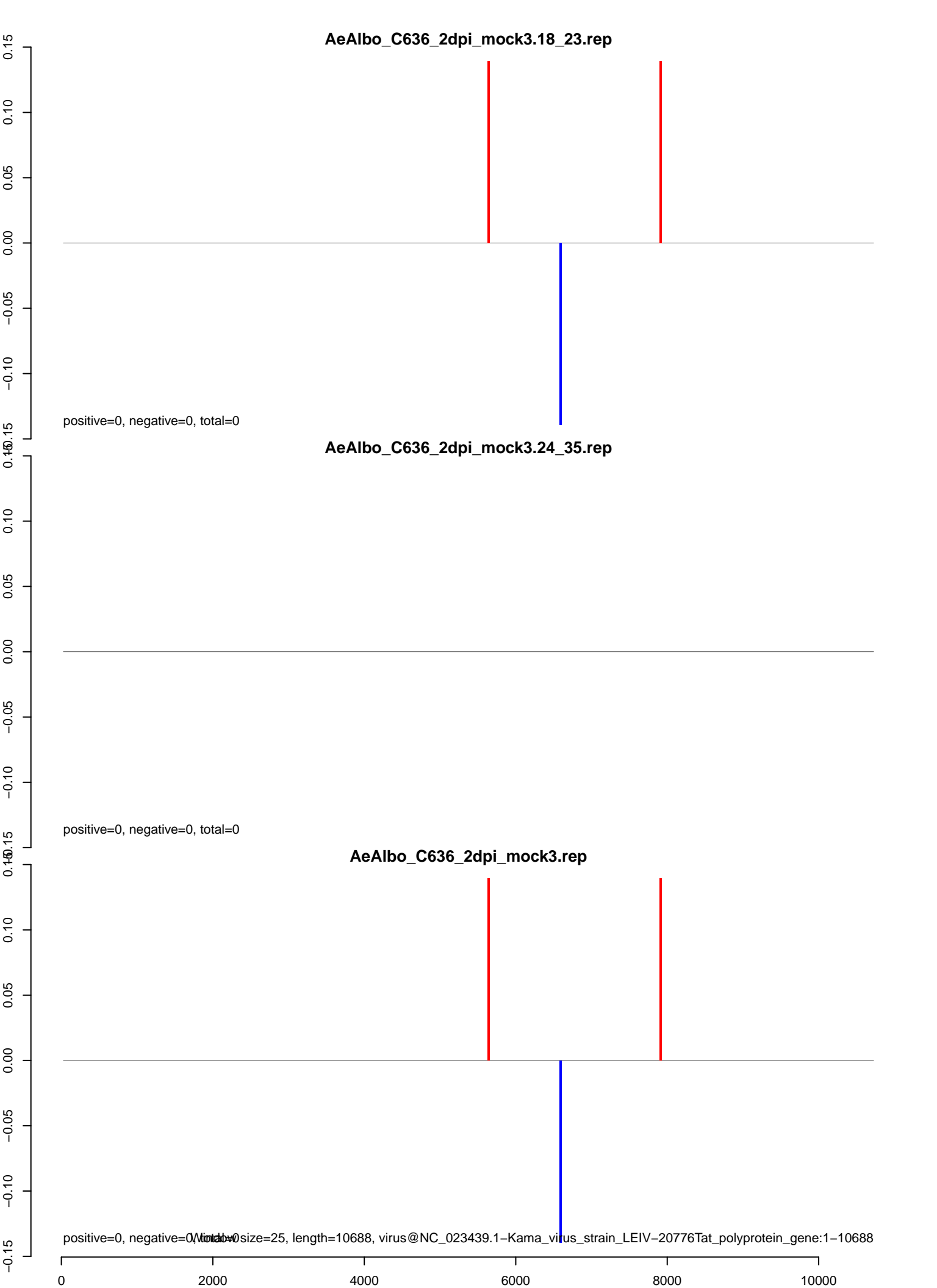
AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



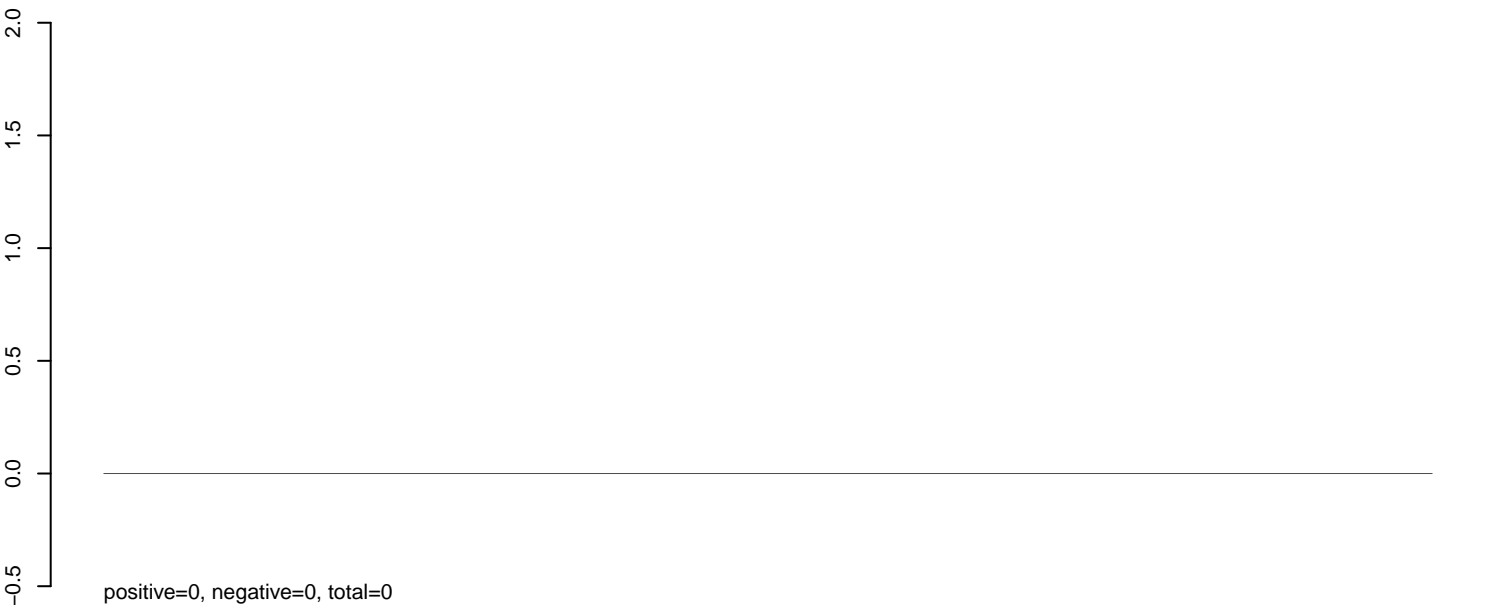
0 2000 4000 6000 8000 10000 12000



AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep

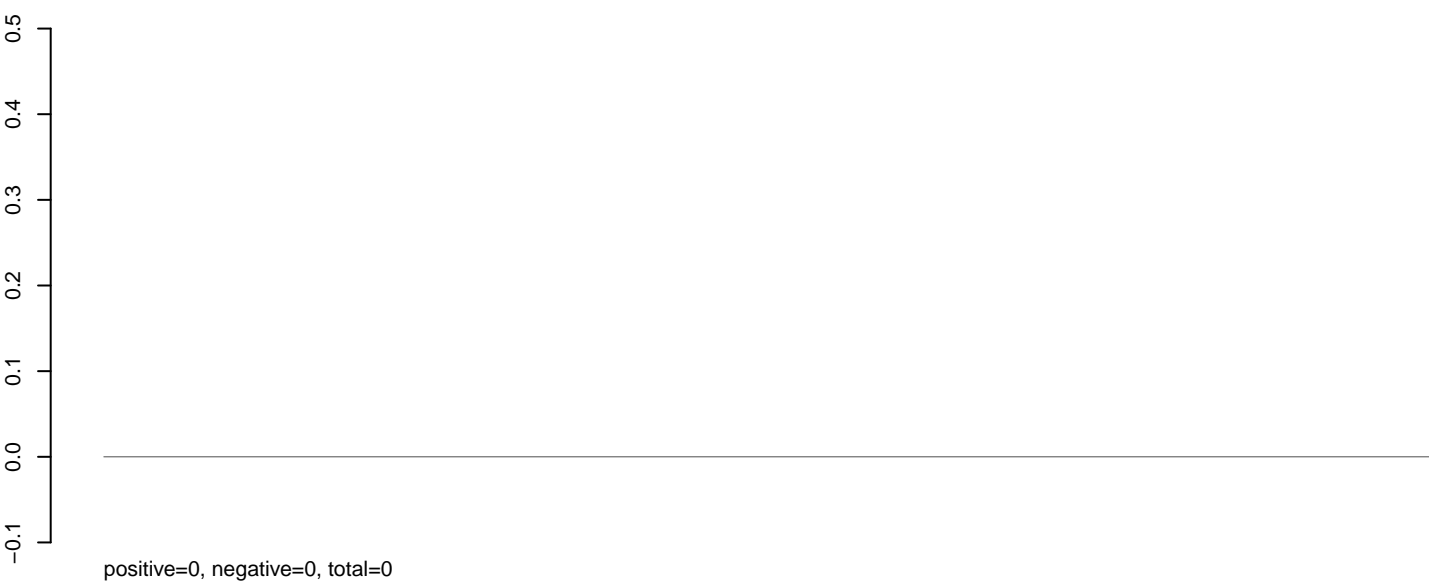


0 2000 4000 6000 8000 10000 12000

AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



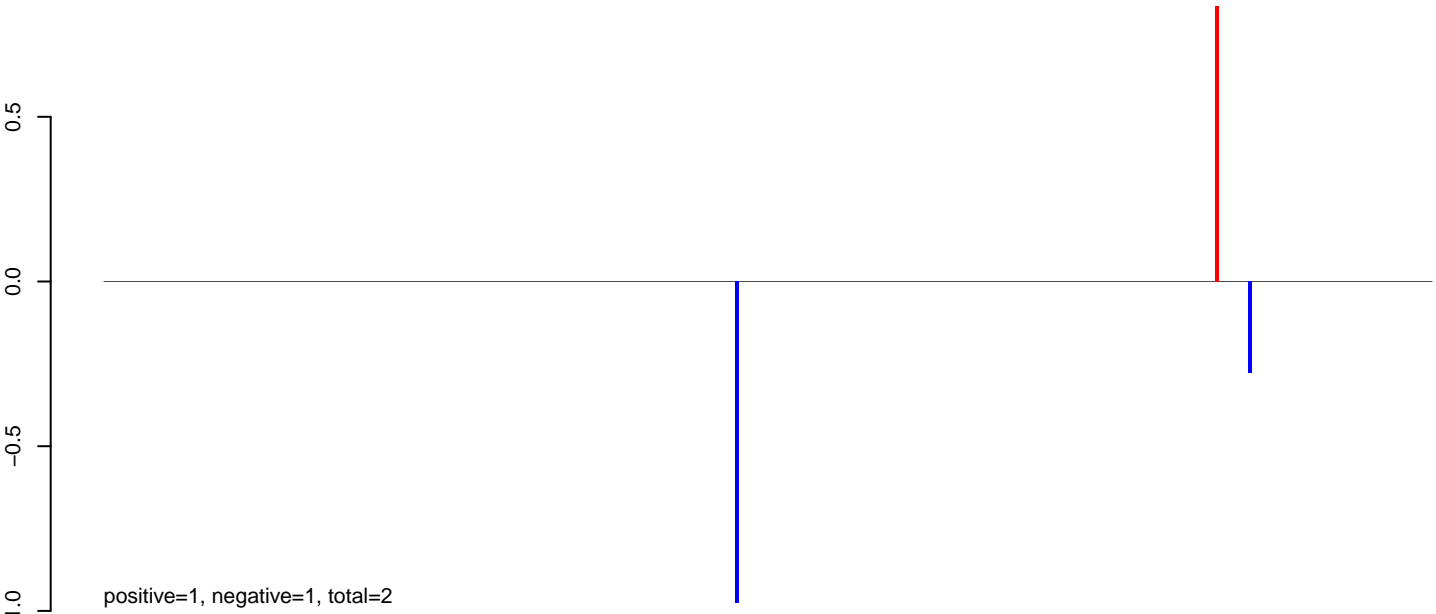
AeAlbo_C636_2dpi_mock3.rep



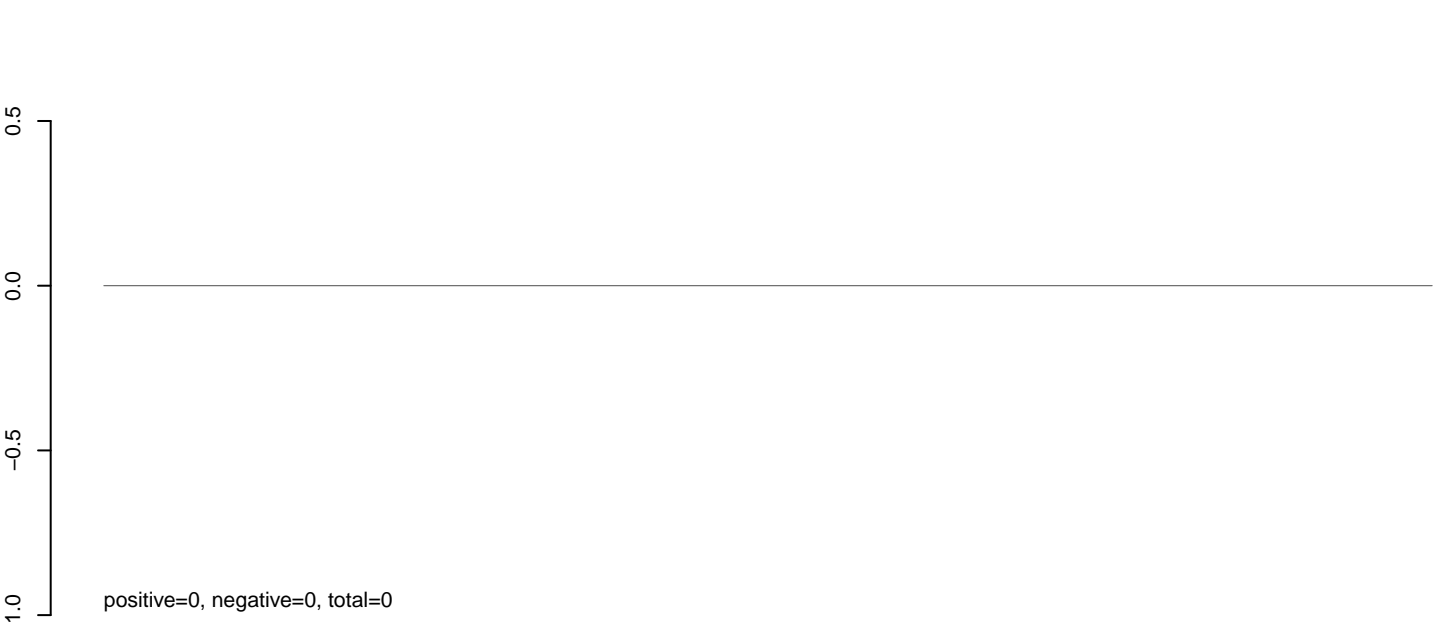
Window size=25, length=10814, virus@NC_012735.1-Wesselsbron_virus:1-10814

0 2000 4000 6000 8000 10000

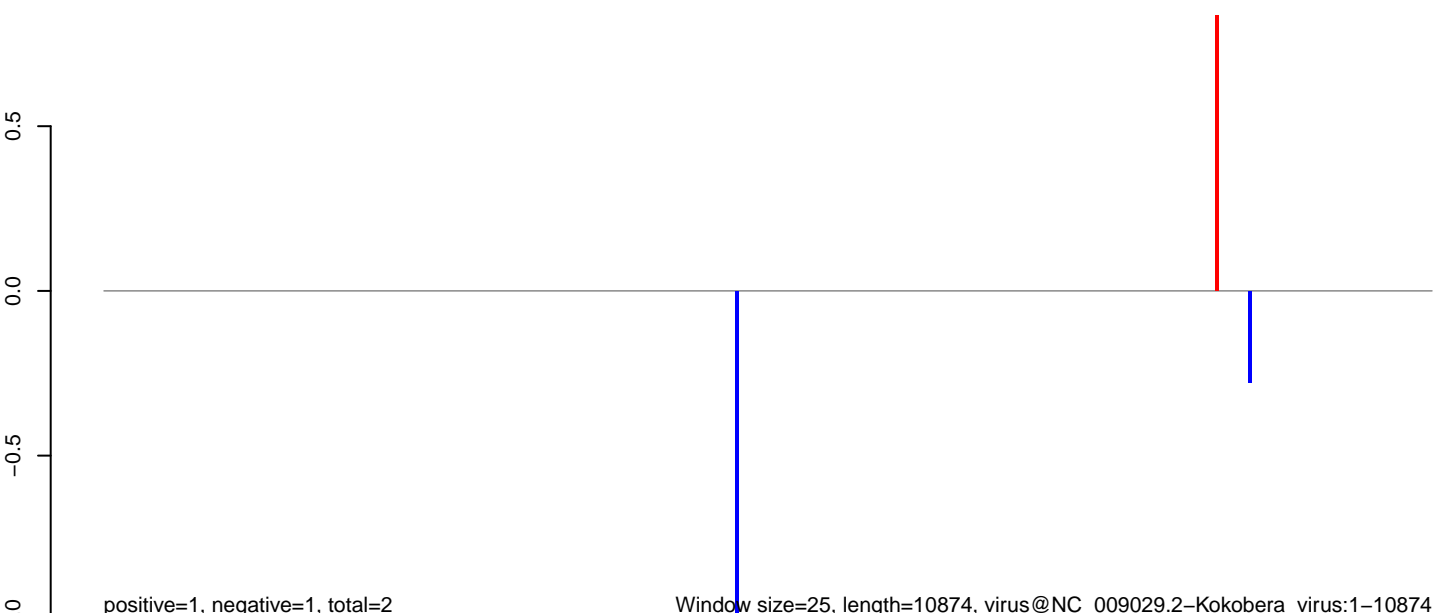
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



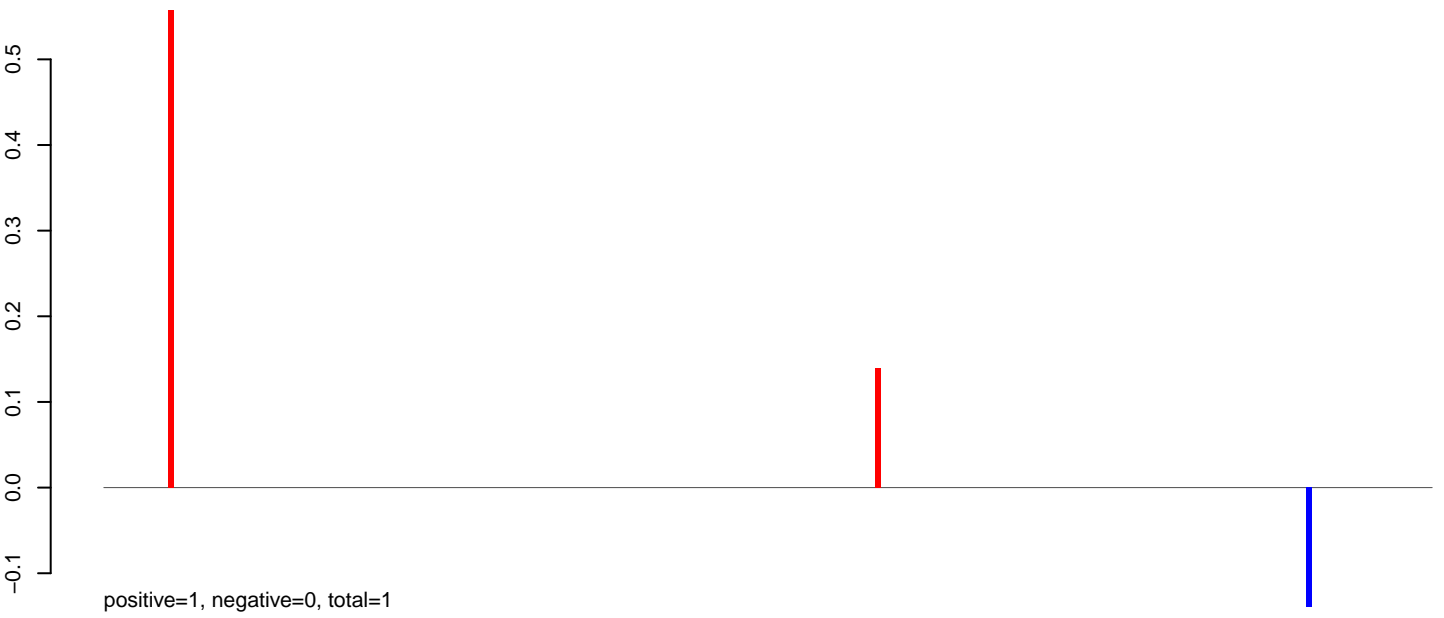
AeAlbo_C636_2dpi_mock3.rep



Window size=25, length=10874, virus@NC_009029.2-Kokobera_virus:1-10874

0 2000 4000 6000 8000 10000

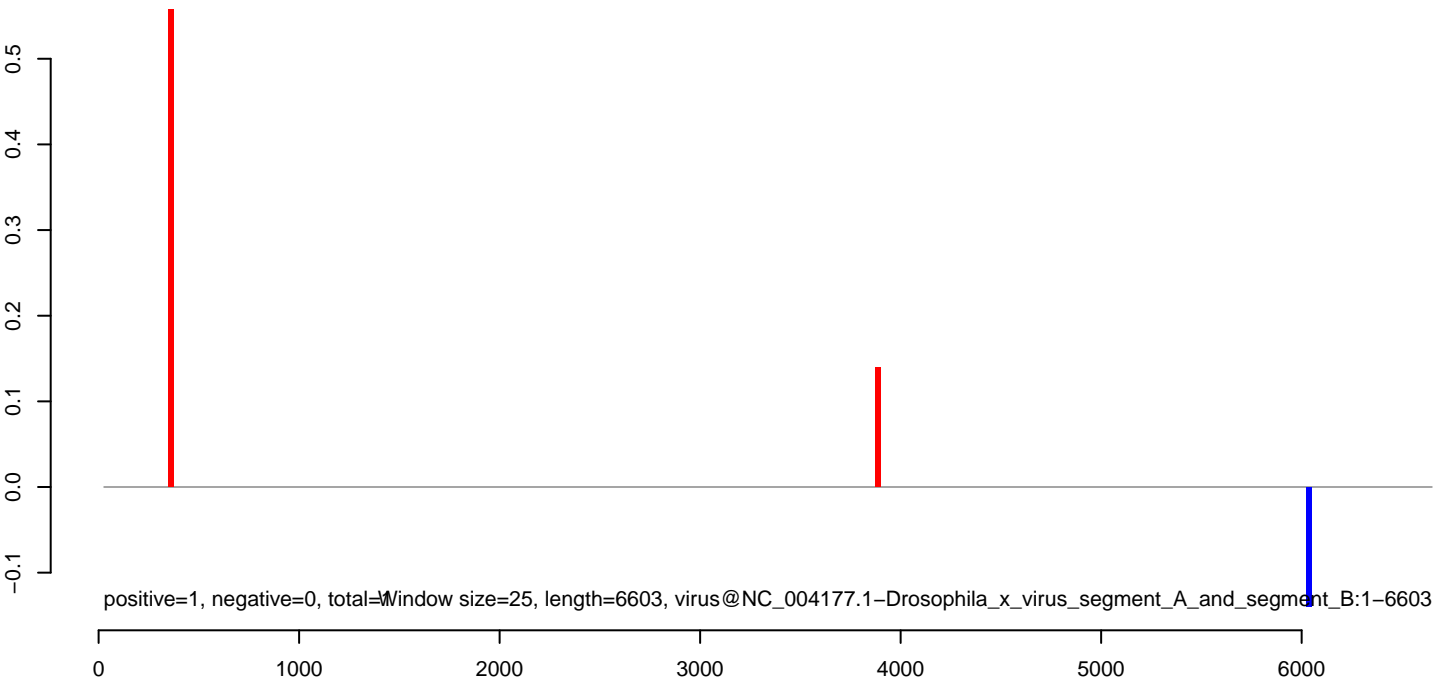
AeAlbo_C636_2dpi_mock3.18_23.rep



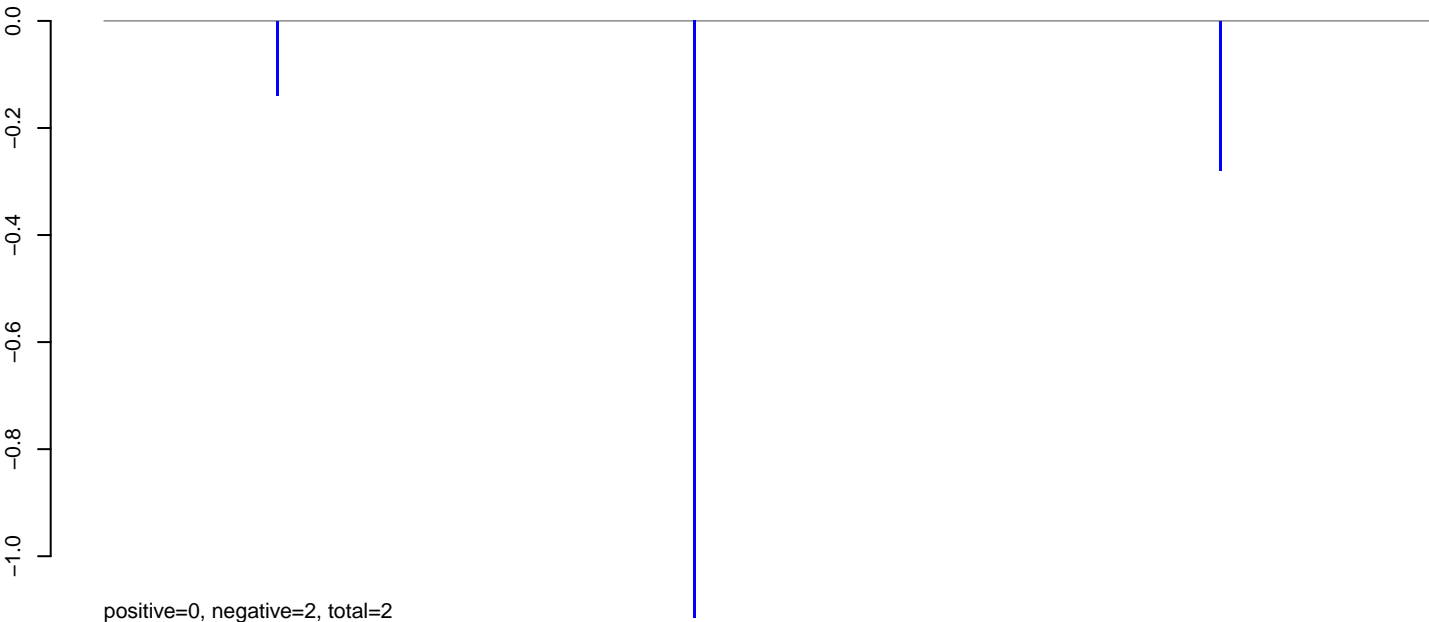
AeAlbo_C636_2dpi_mock3.24_35.rep



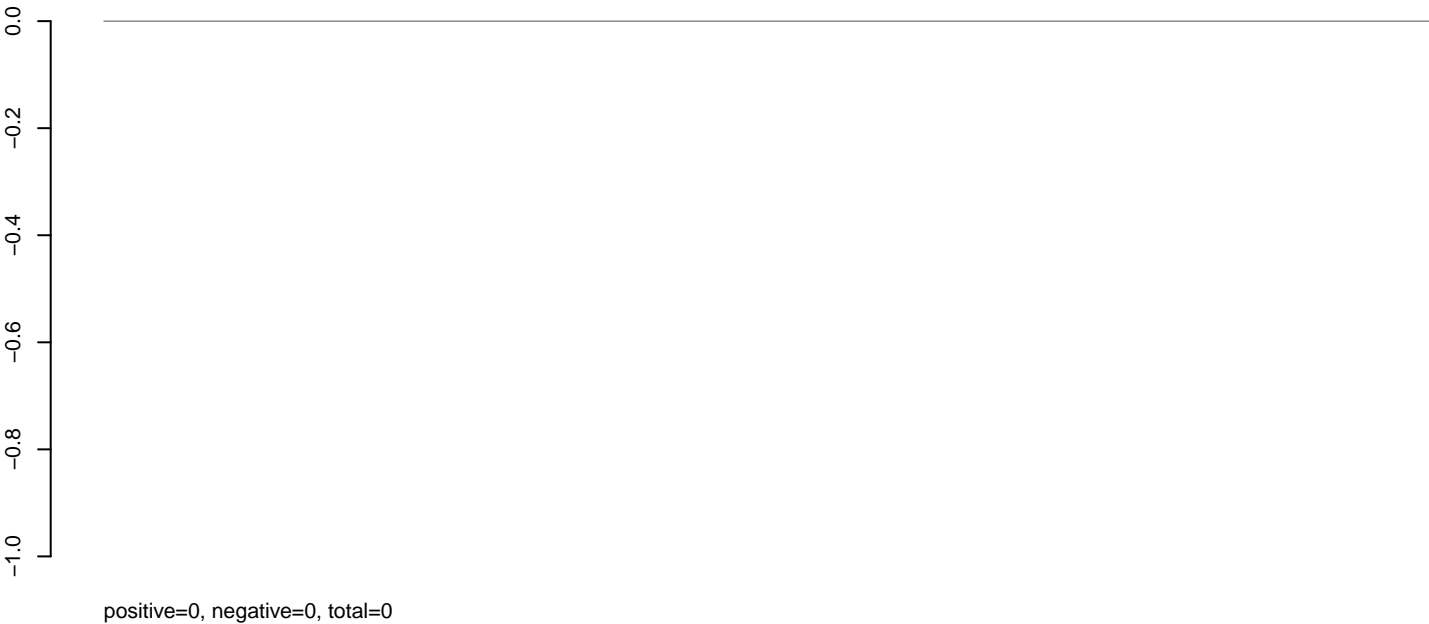
AeAlbo_C636_2dpi_mock3.rep



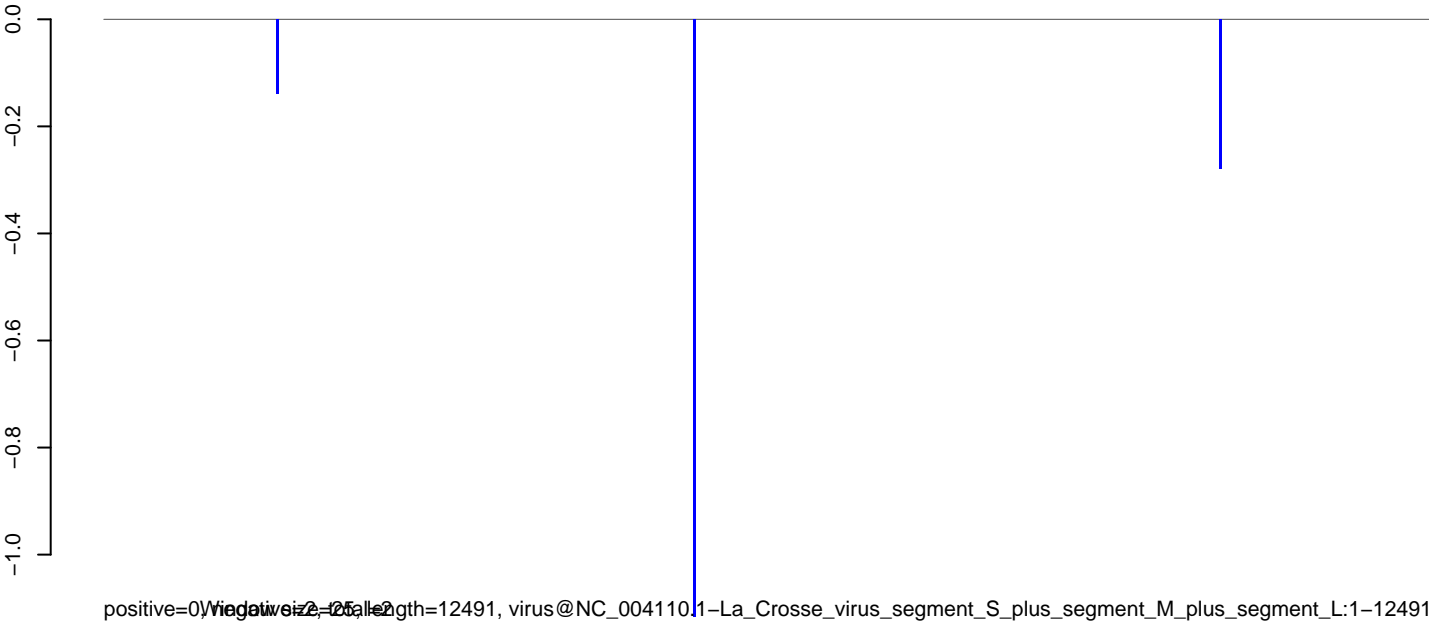
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



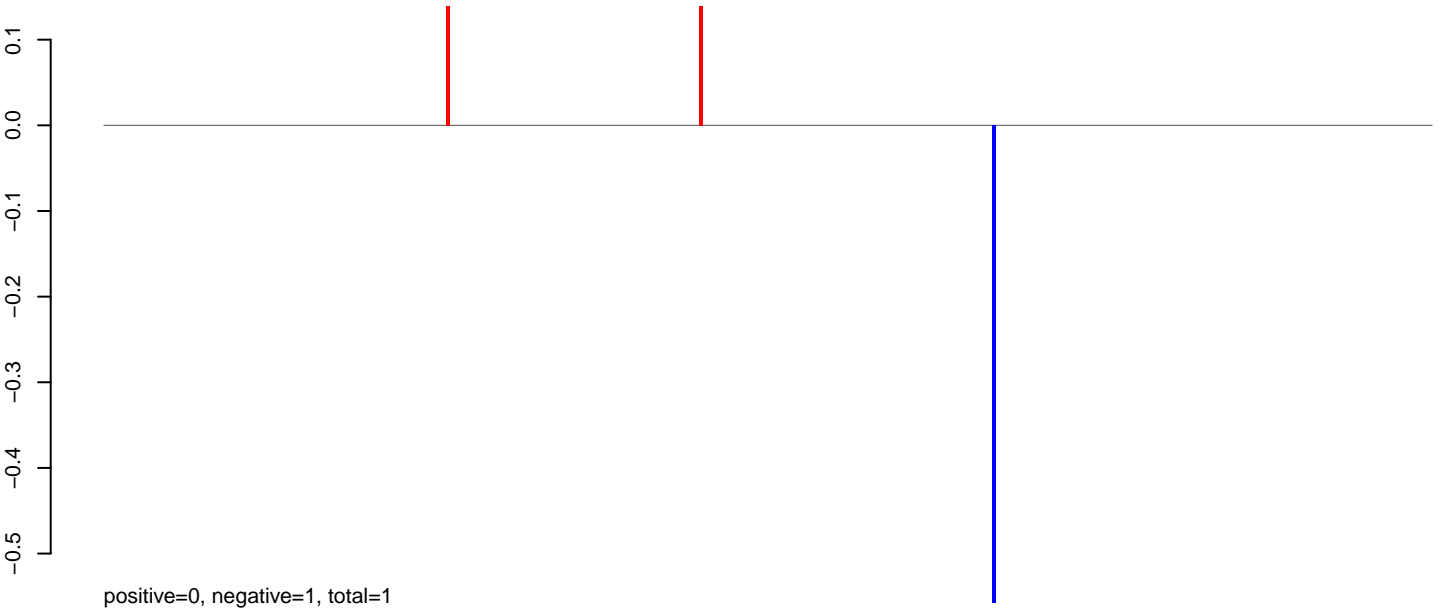
AeAlbo_C636_2dpi_mock3.rep



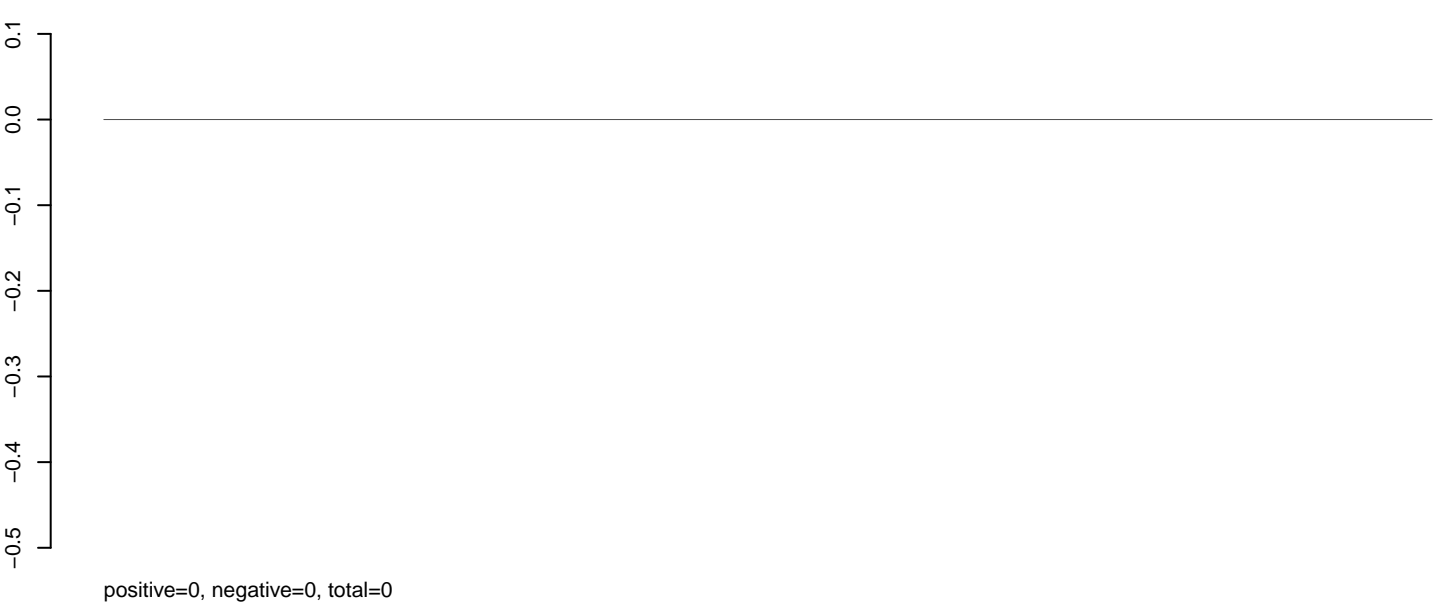
negative=2, length=12491, virus@NC_004110.1-La_Crosse_virus_segment_S_plus_segment_M_plus_segment_L:1-12491

0 2000 4000 6000 8000 10000 12000

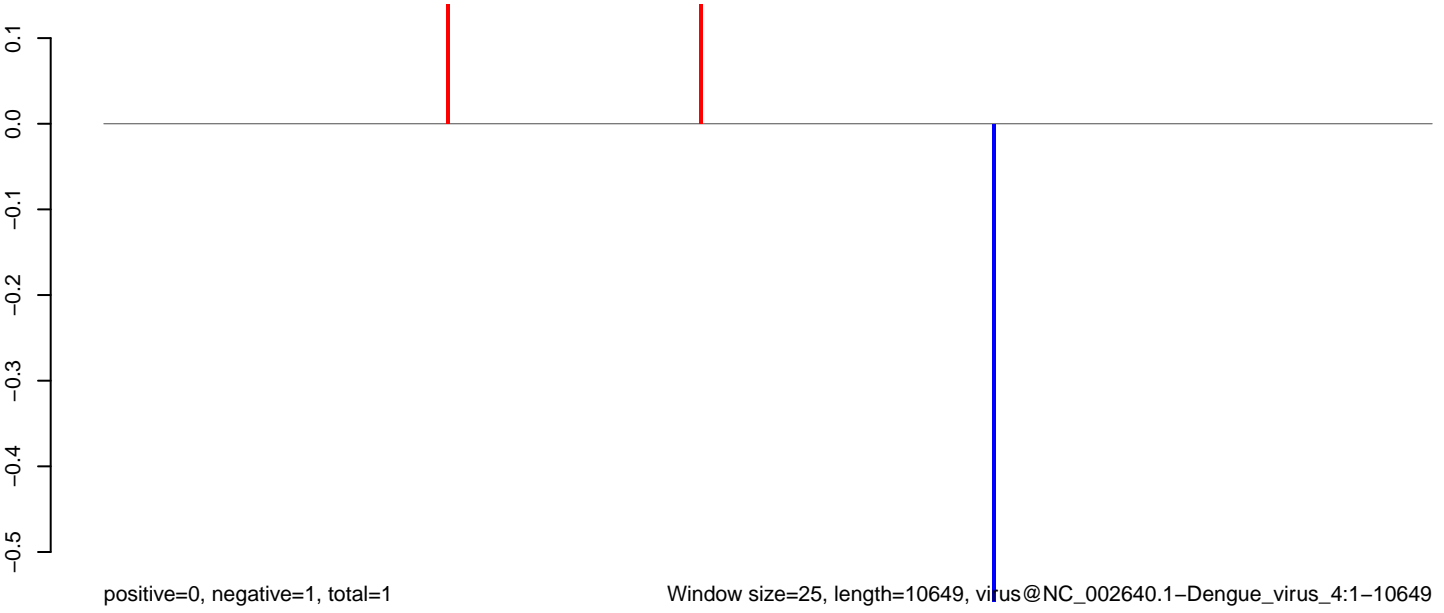
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



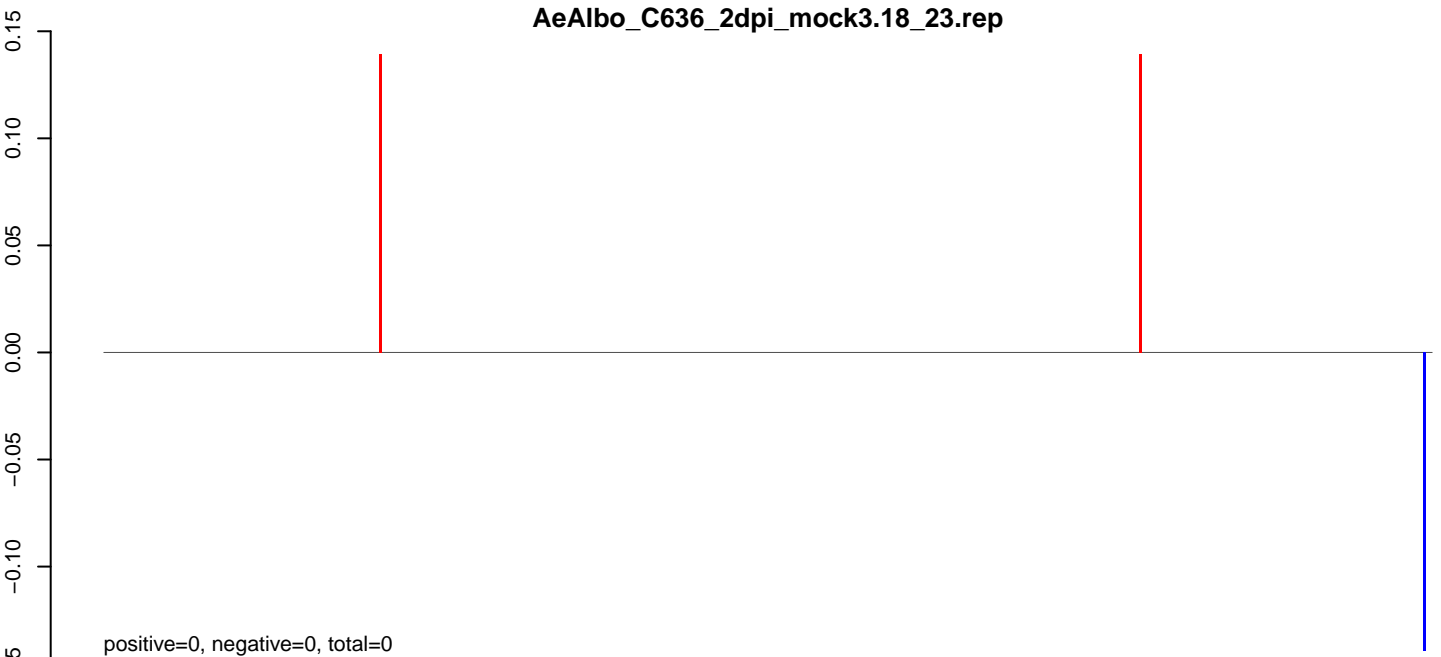
AeAlbo_C636_2dpi_mock3.rep



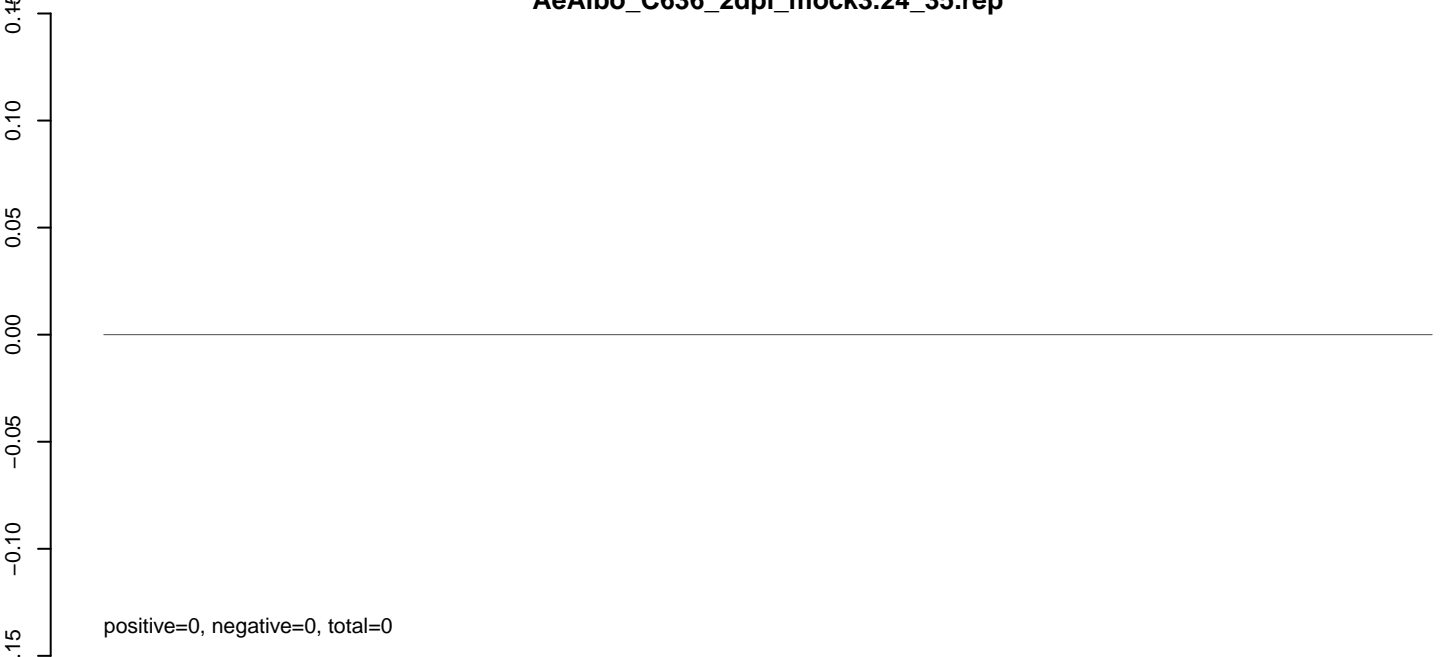
Window size=25, length=10649, virus@NC_002640.1-Dengue_virus_4:1-10649

0 2000 4000 6000 8000 10000

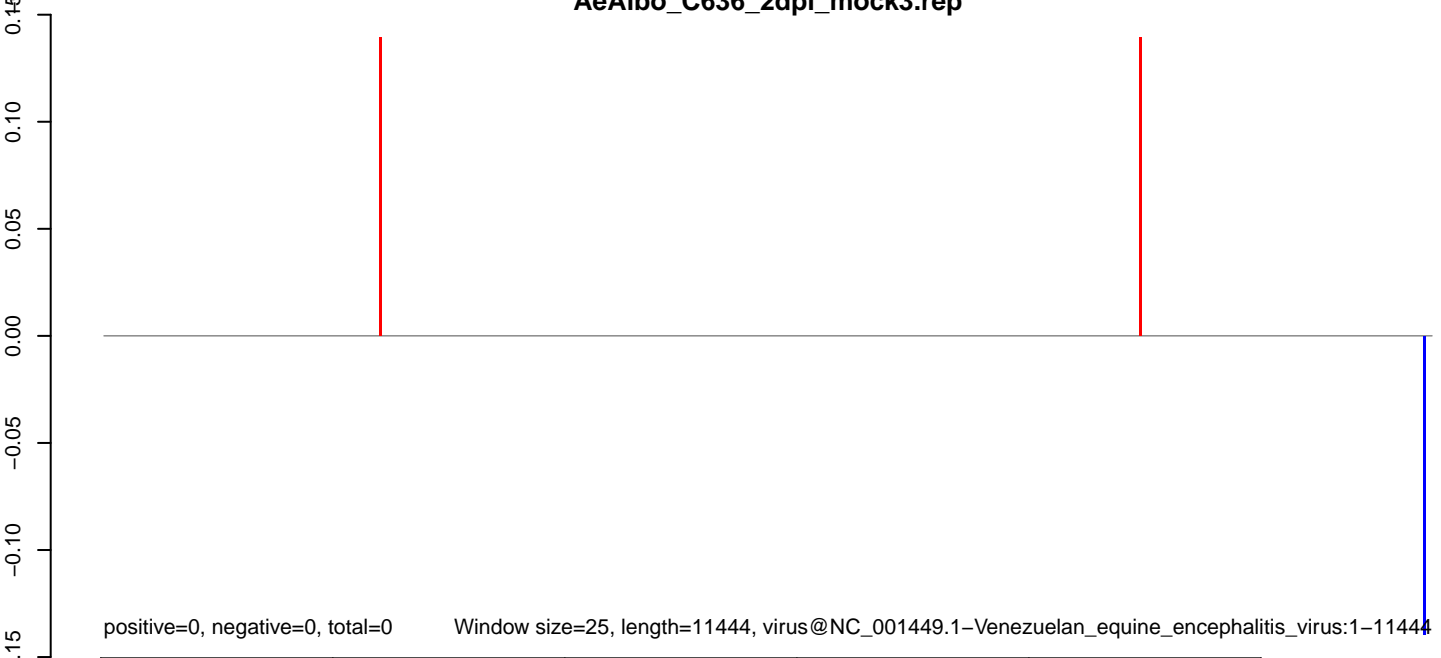
AeAlbo_C636_2dpi_mock3.18_23.rep



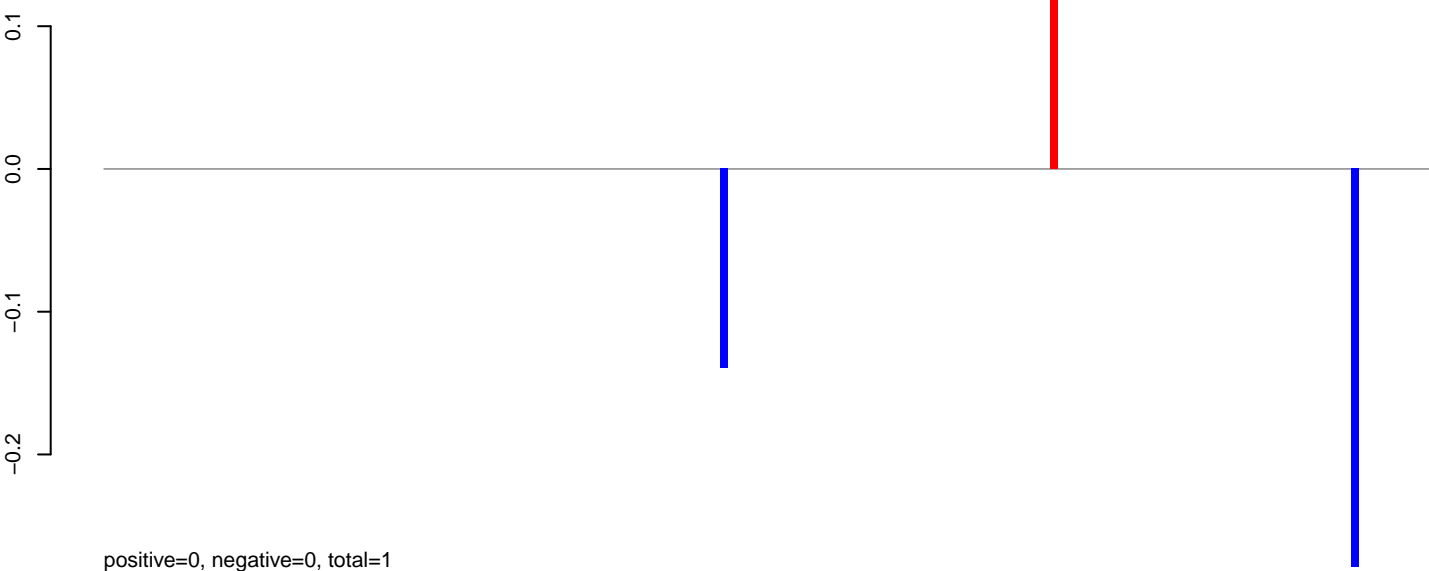
AeAlbo_C636_2dpi_mock3.24_35.rep



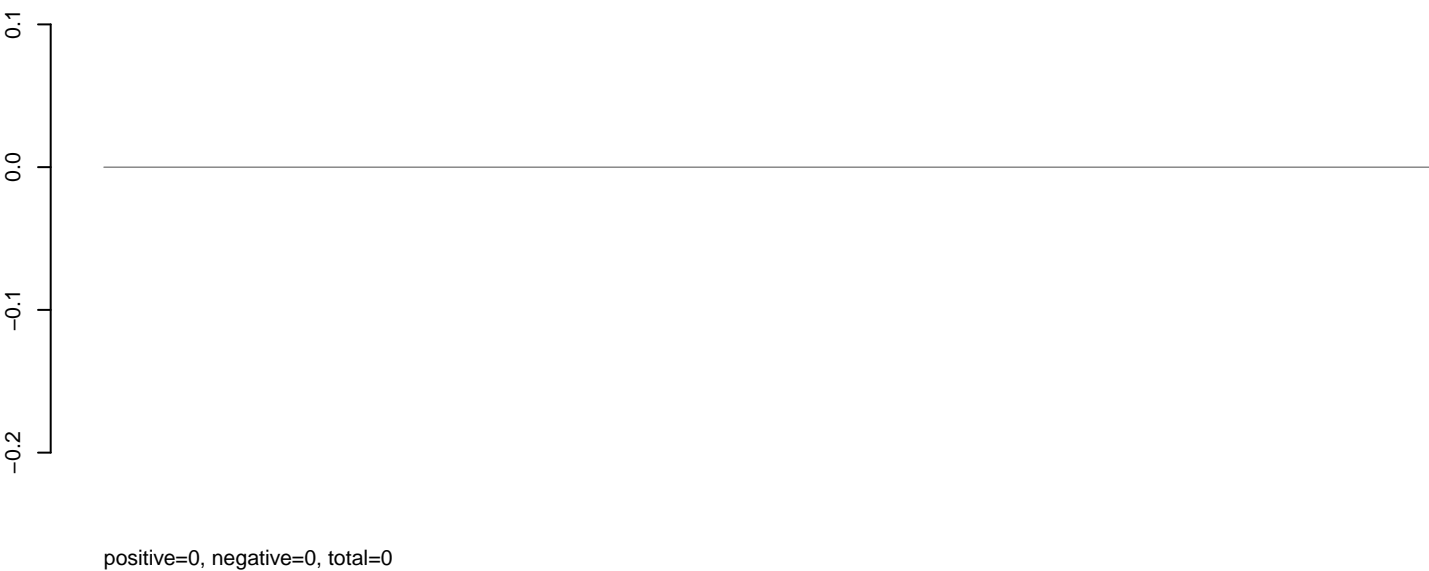
AeAlbo_C636_2dpi_mock3.rep



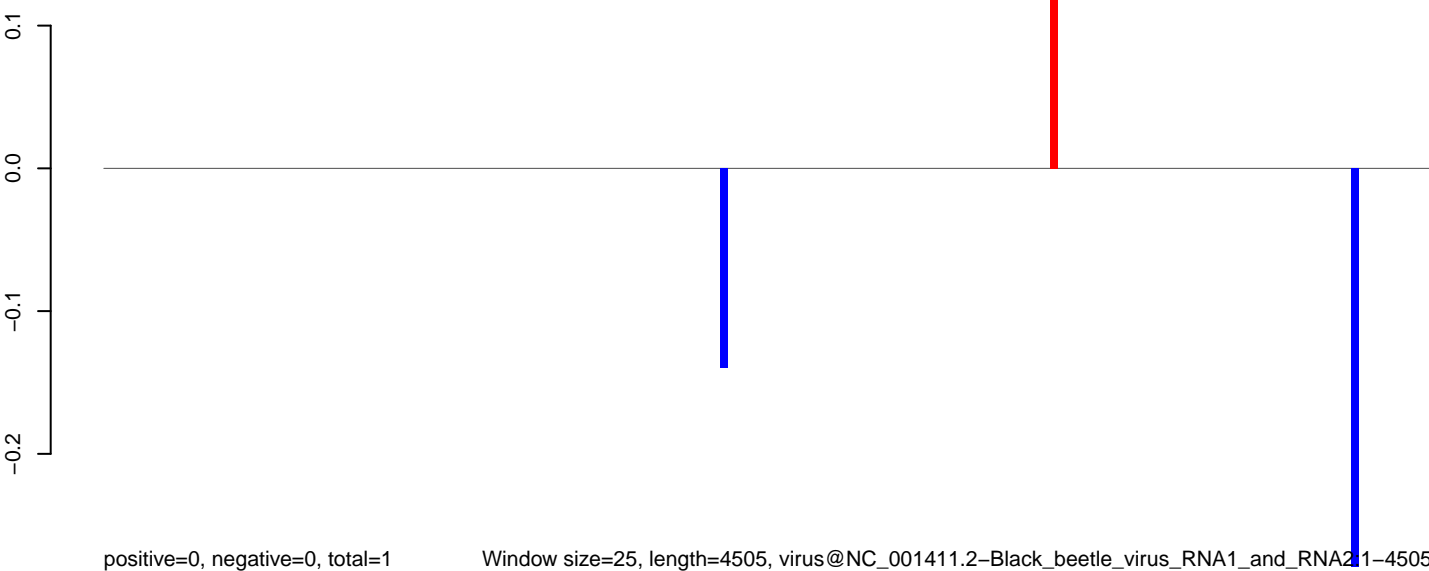
AeAlbo_C636_2dpi_mock3.18_23.rep



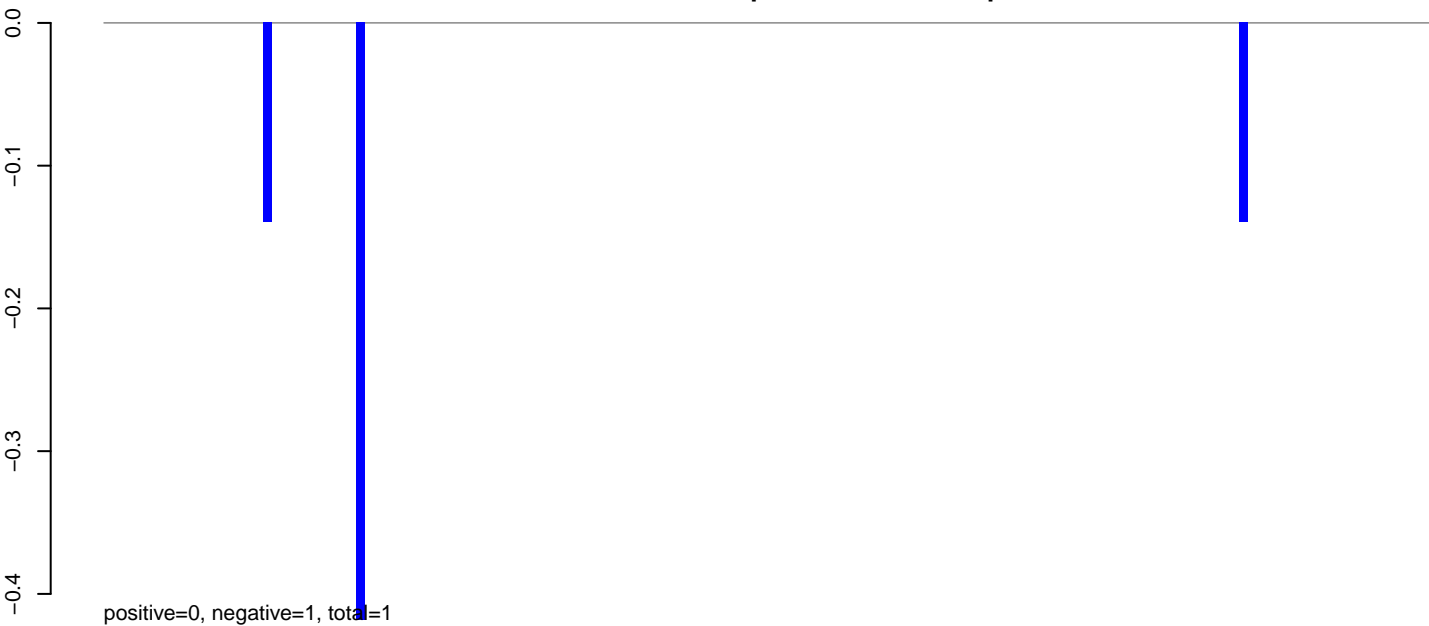
AeAlbo_C636_2dpi_mock3.24_35.rep



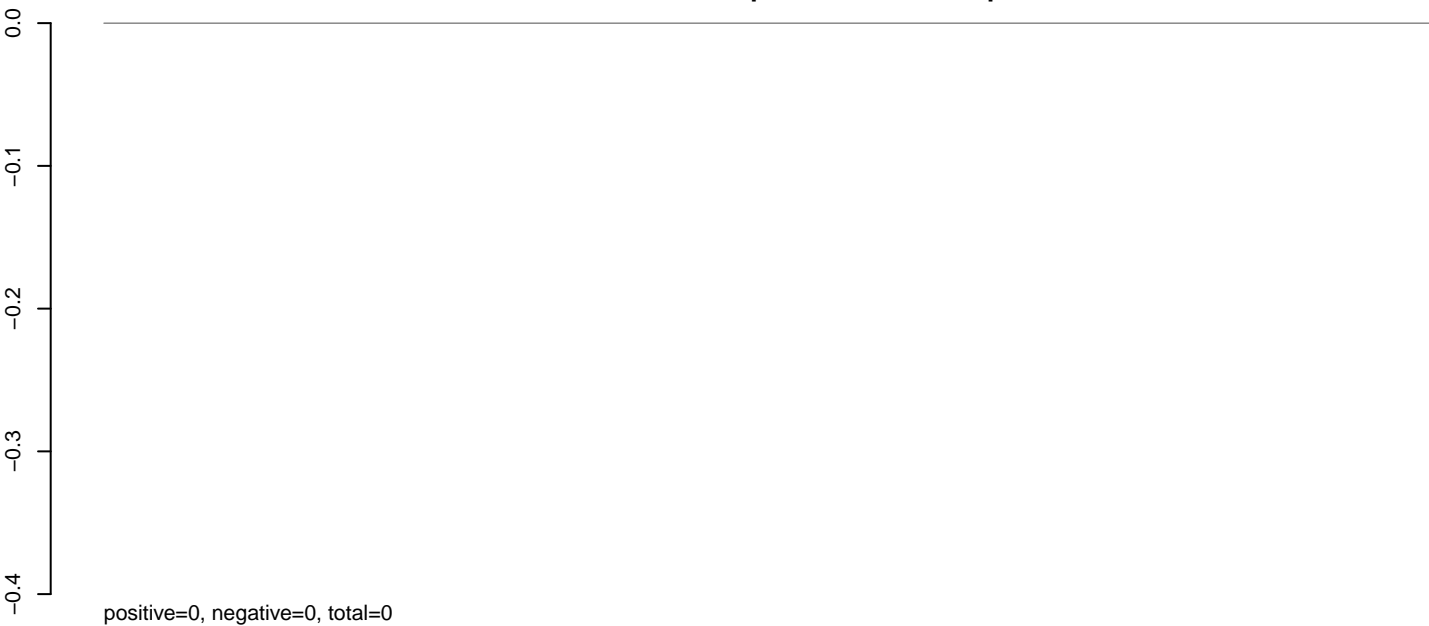
AeAlbo_C636_2dpi_mock3.rep



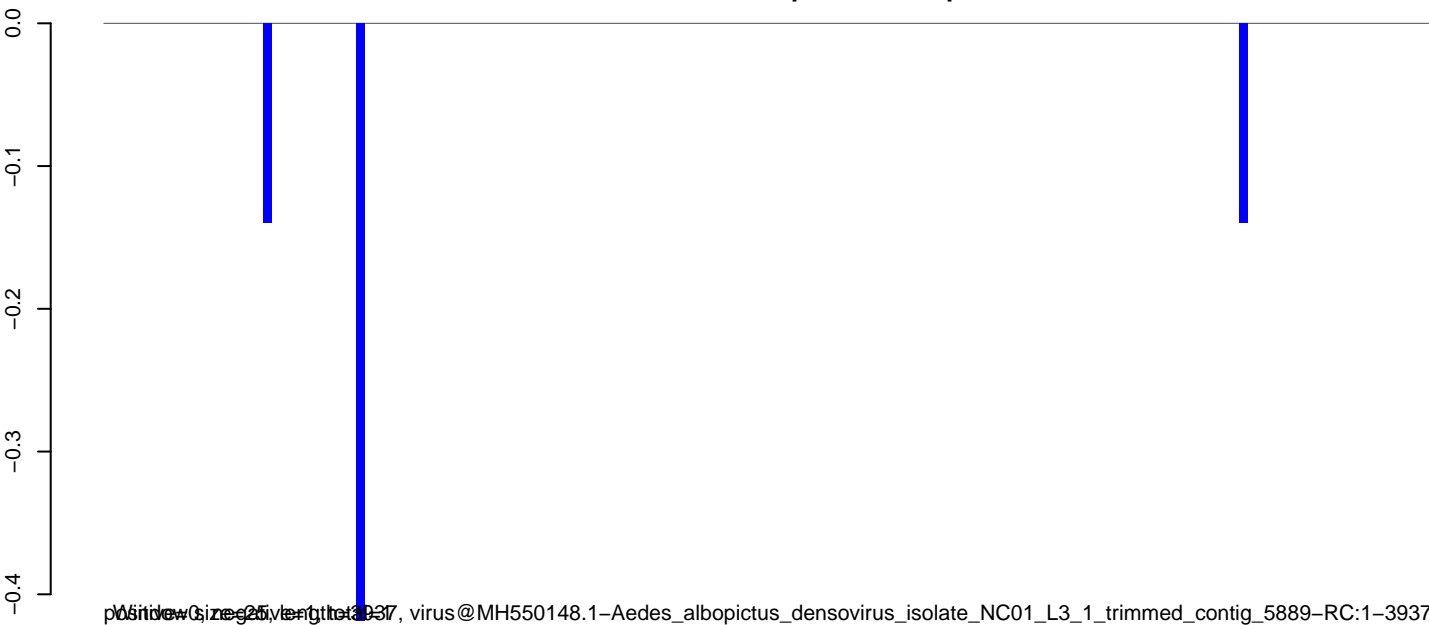
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

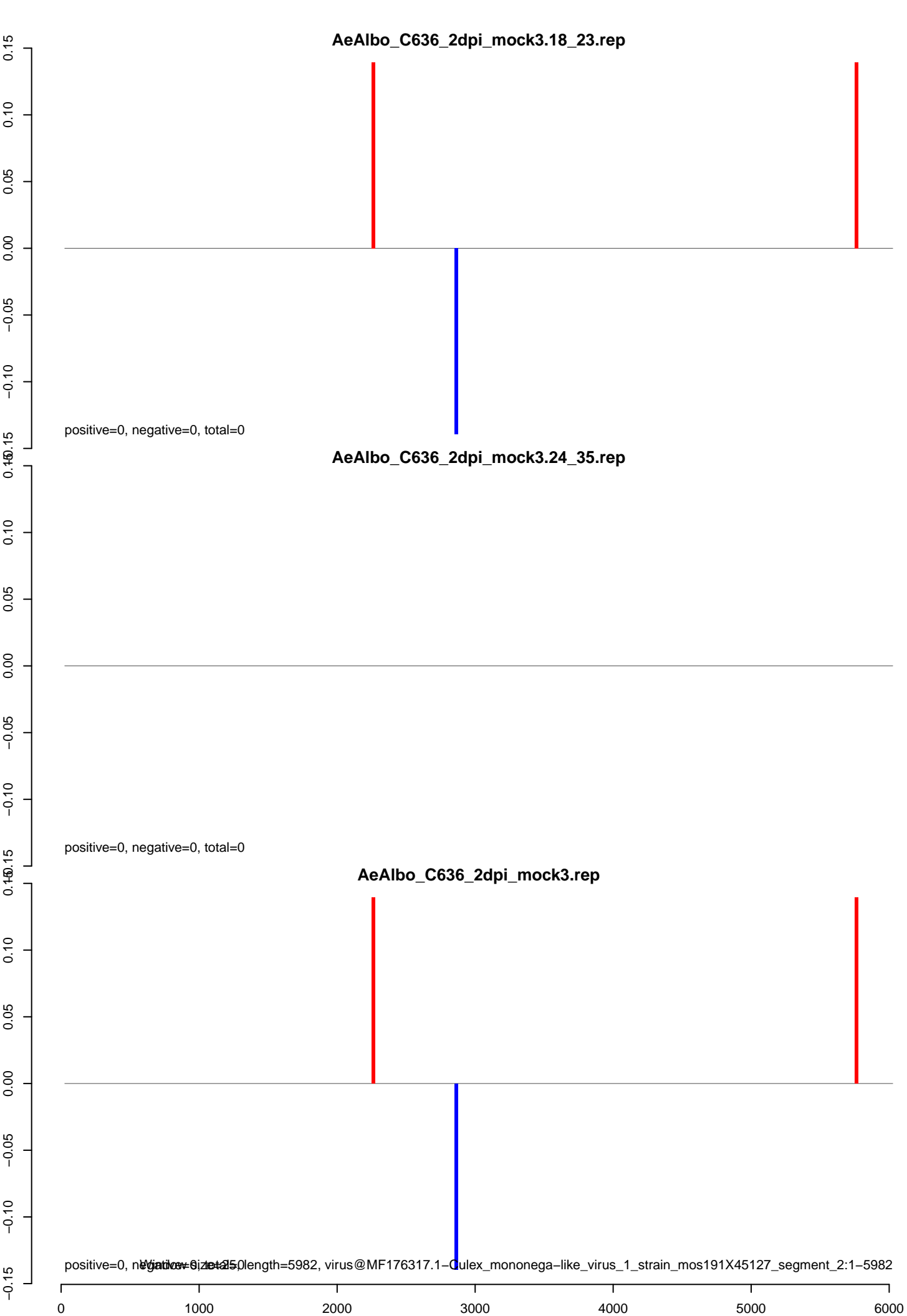


AeAlbo_C636_2dpi_mock3.rep

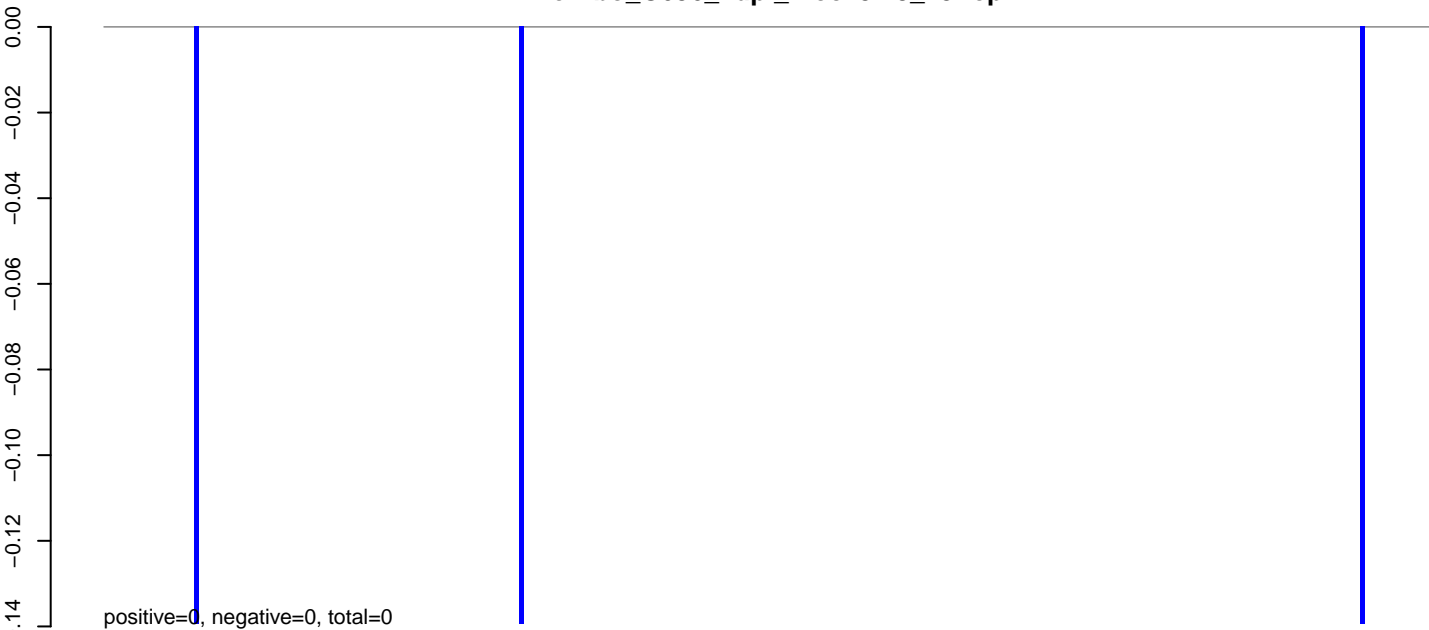


Window size: 25, length: 3937, virus@MH550148.1-Aedes_albopictus_densovirus_isolate_NC01_L3_1_trimmed_contig_5889-RC:1-3937

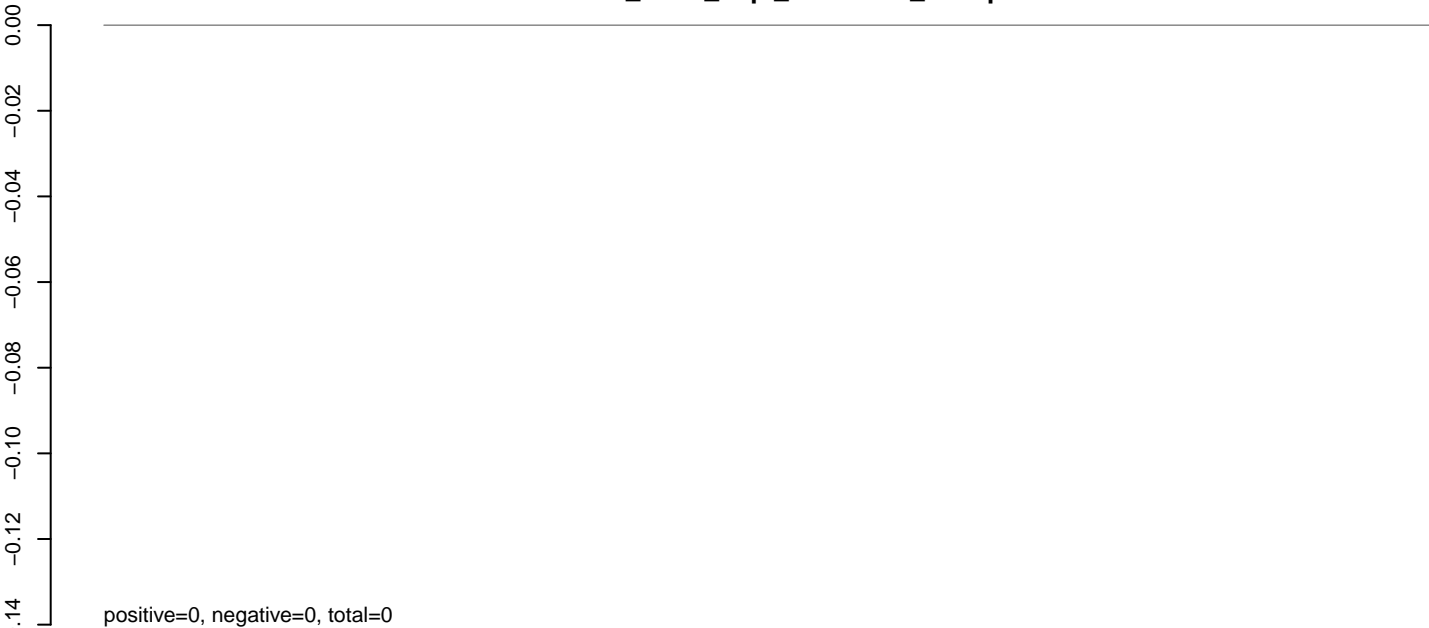
0 1000 2000 3000 4000



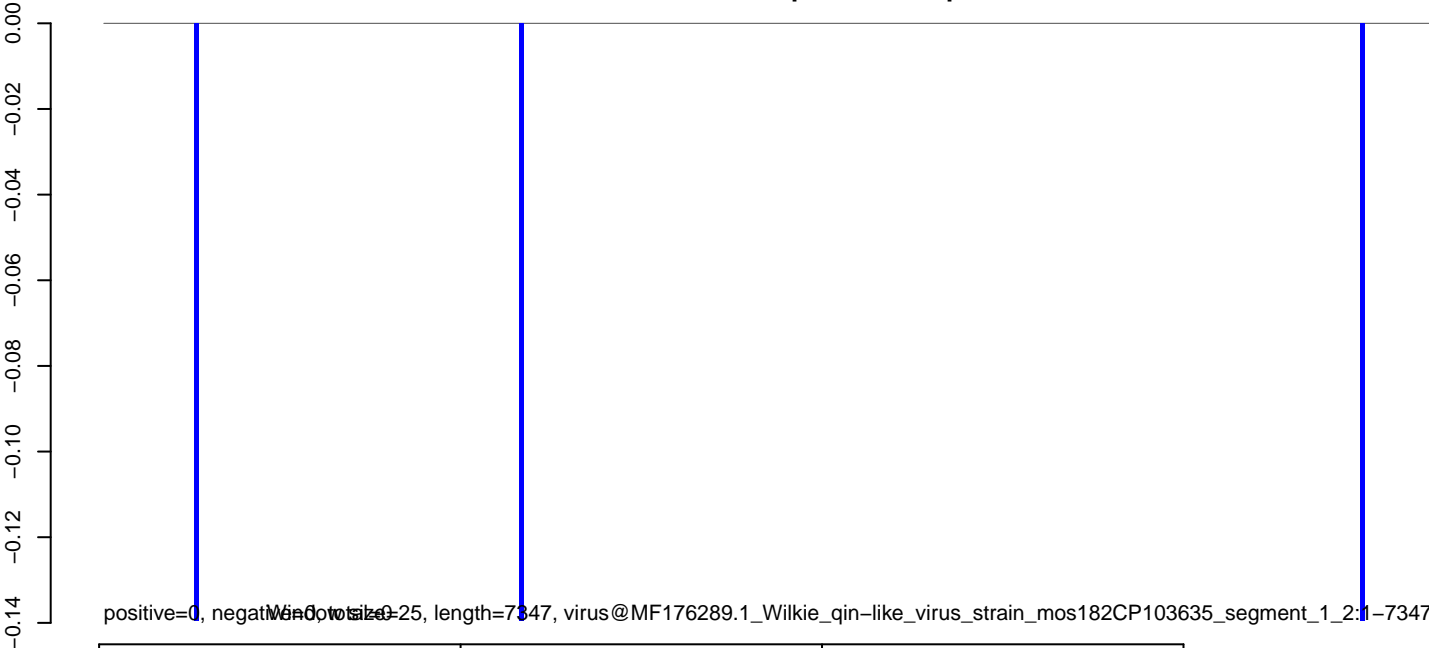
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



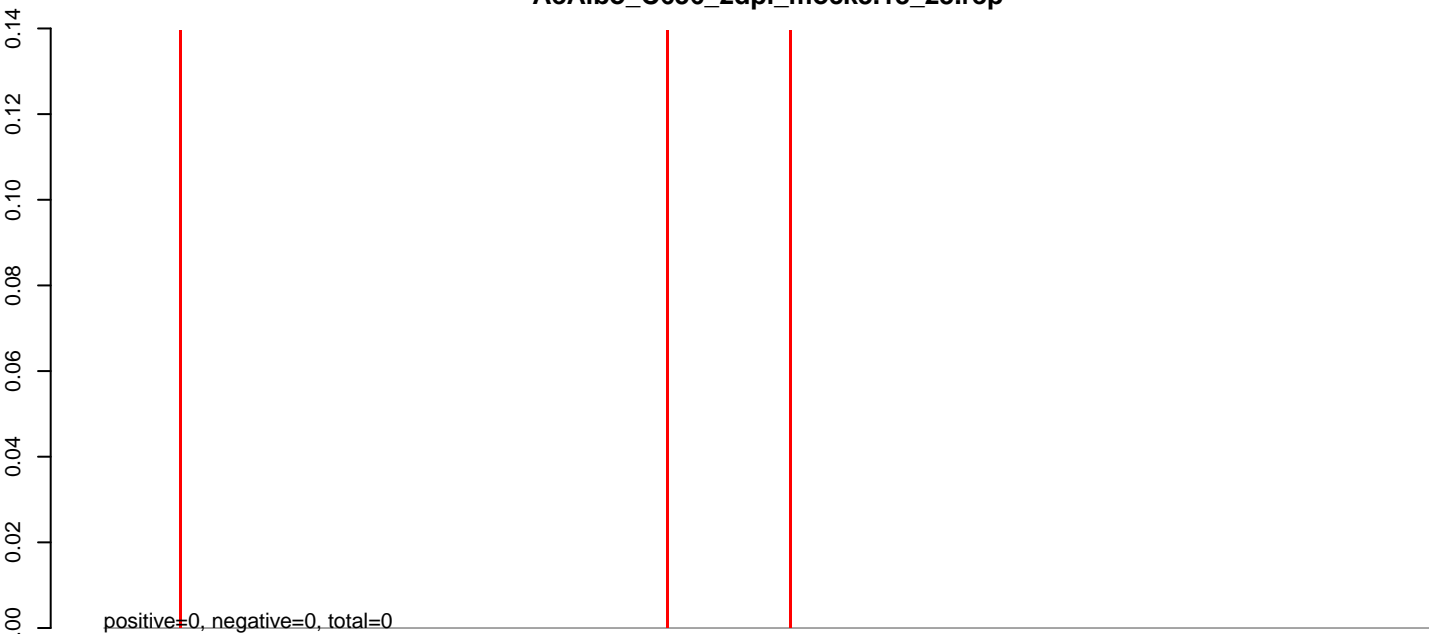
AeAlbo_C636_2dpi_mock3.rep



0 2000 4000 6000

Window size=25, length=7347, virus@MF176289.1_Wilkie_qin-like_virus_strain_mos182CP103635_segment_1_2:1-7347

AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

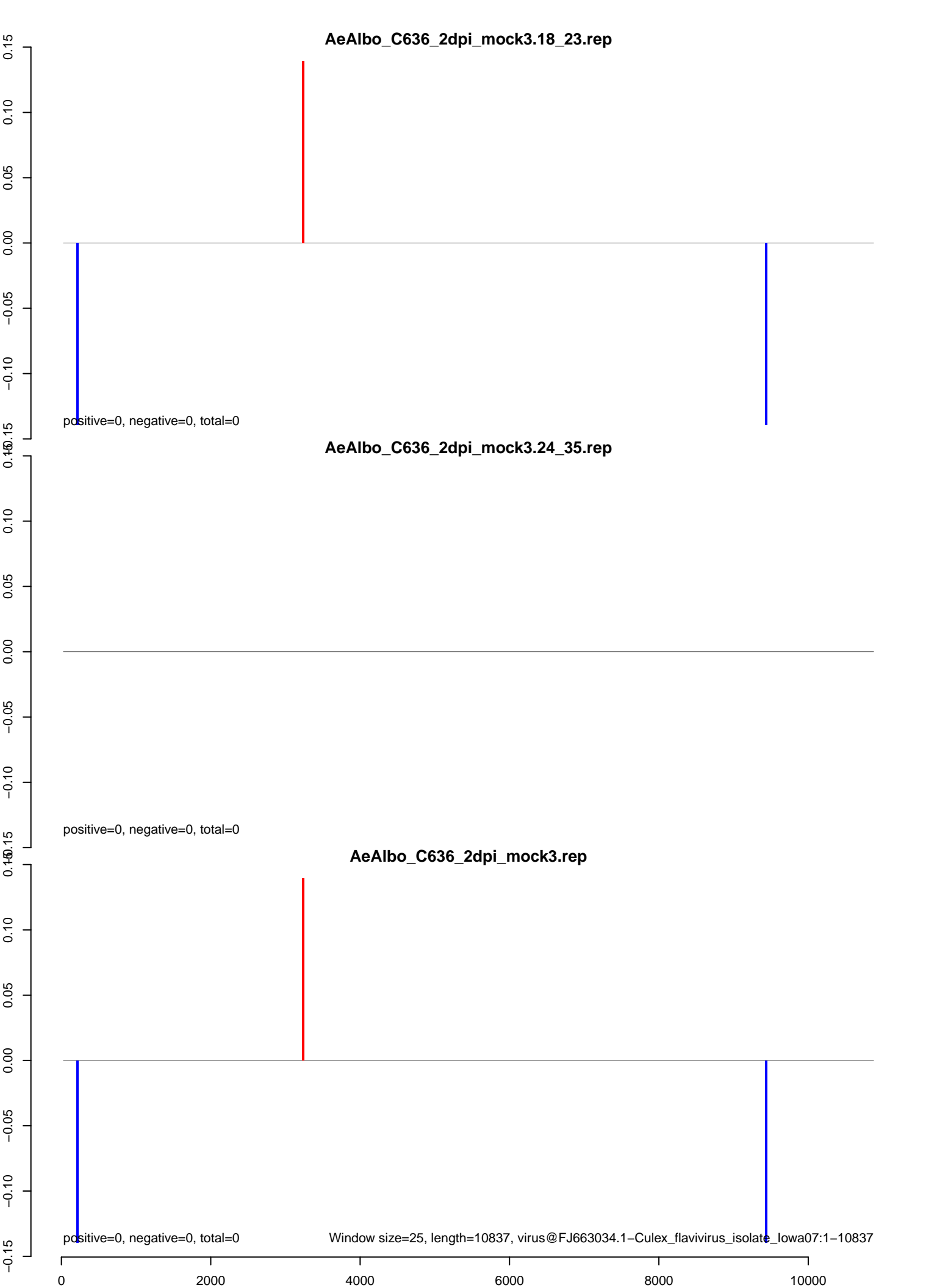


AeAlbo_C636_2dpi_mock3.rep

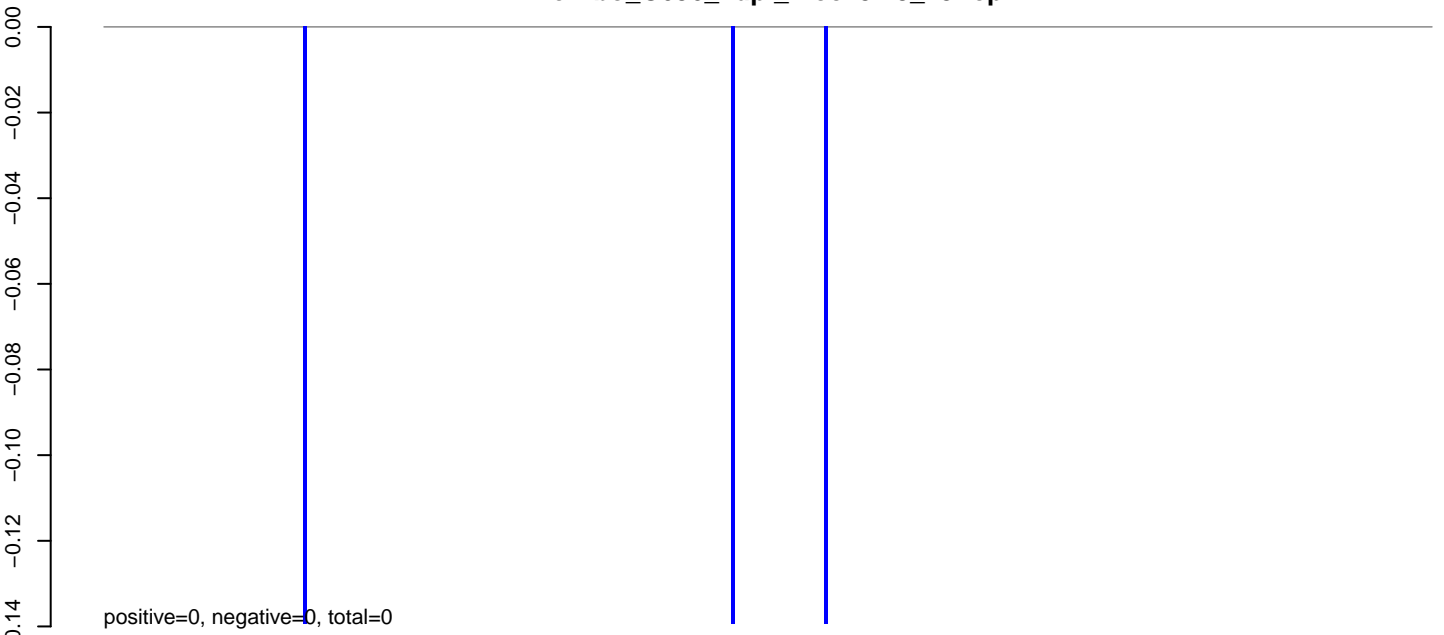


0 2000 4000 6000 8000 10000

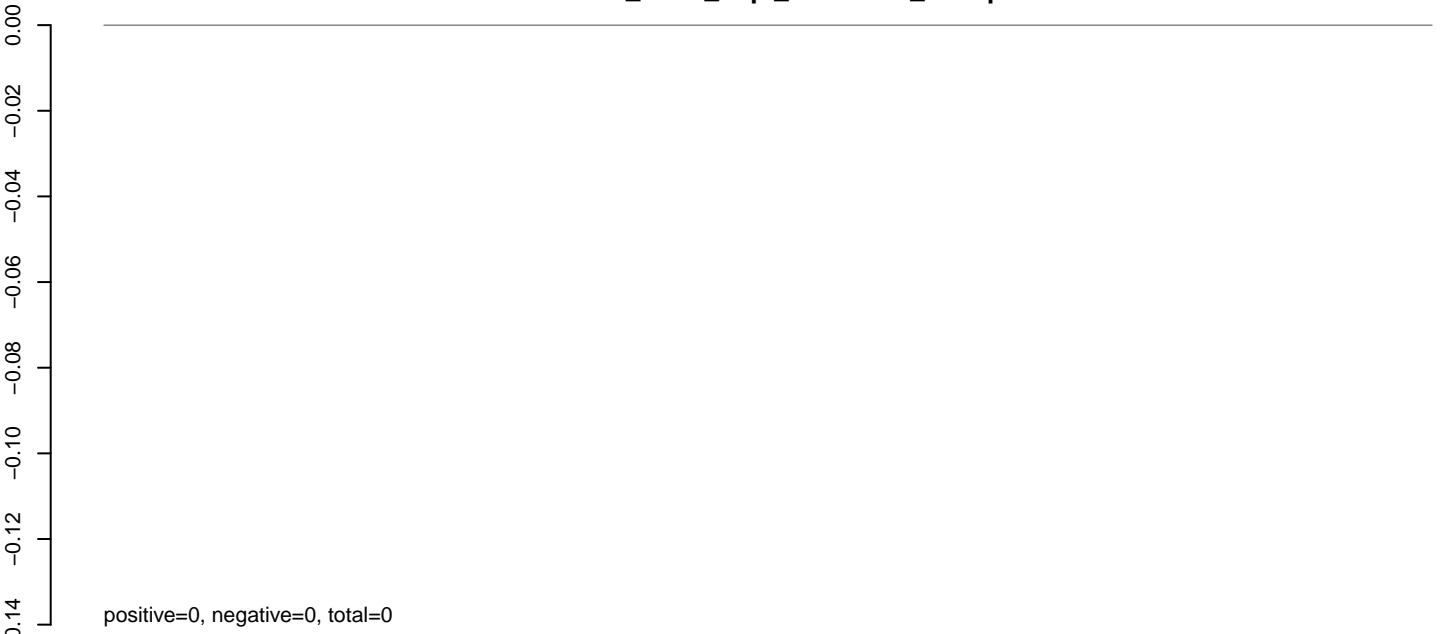
Window size=25, length=11079, virus@KC181923.1-Aedes flavivirus_strain_AEFV-SPFLD-MO-2011-MP6:1-11079



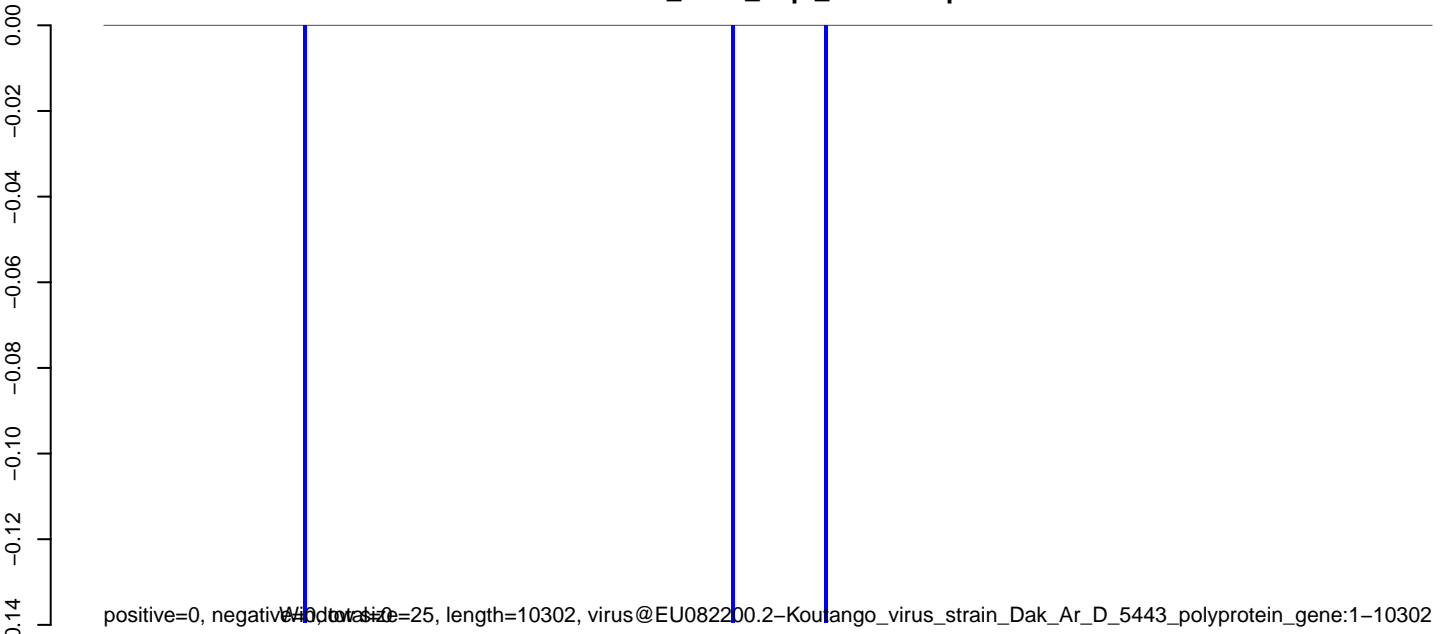
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

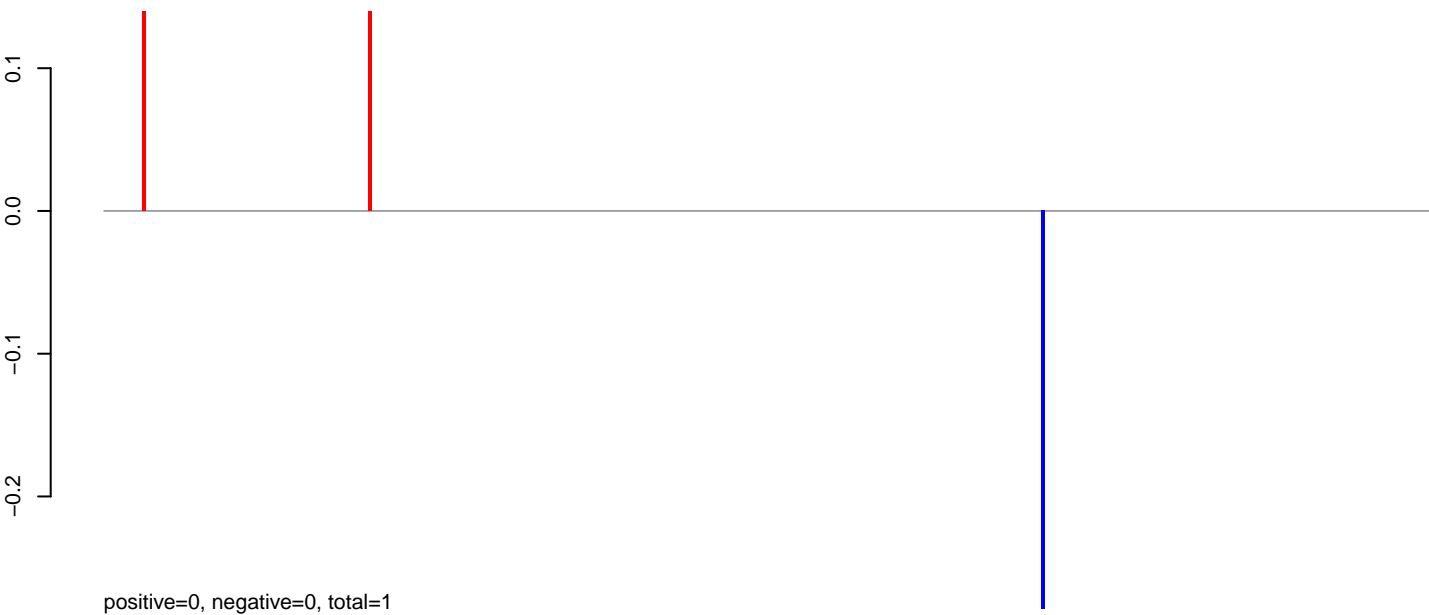


AeAlbo_C636_2dpi_mock3.rep

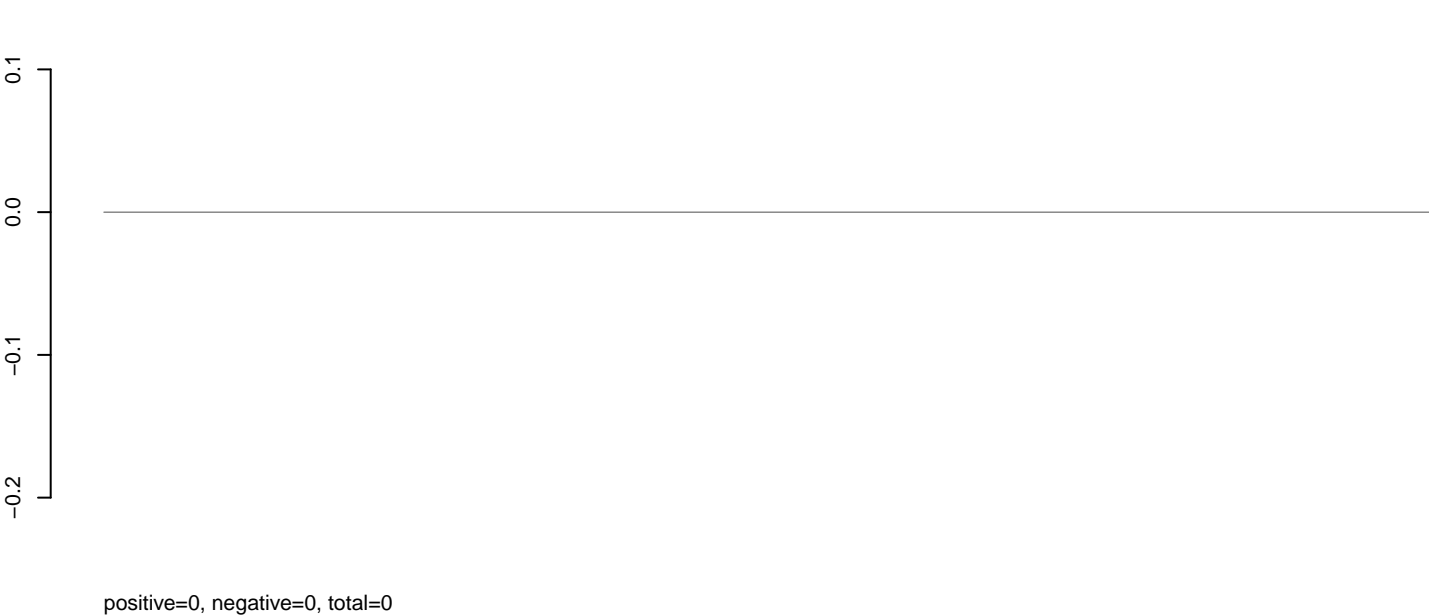


0 2000 4000 6000 8000 10000

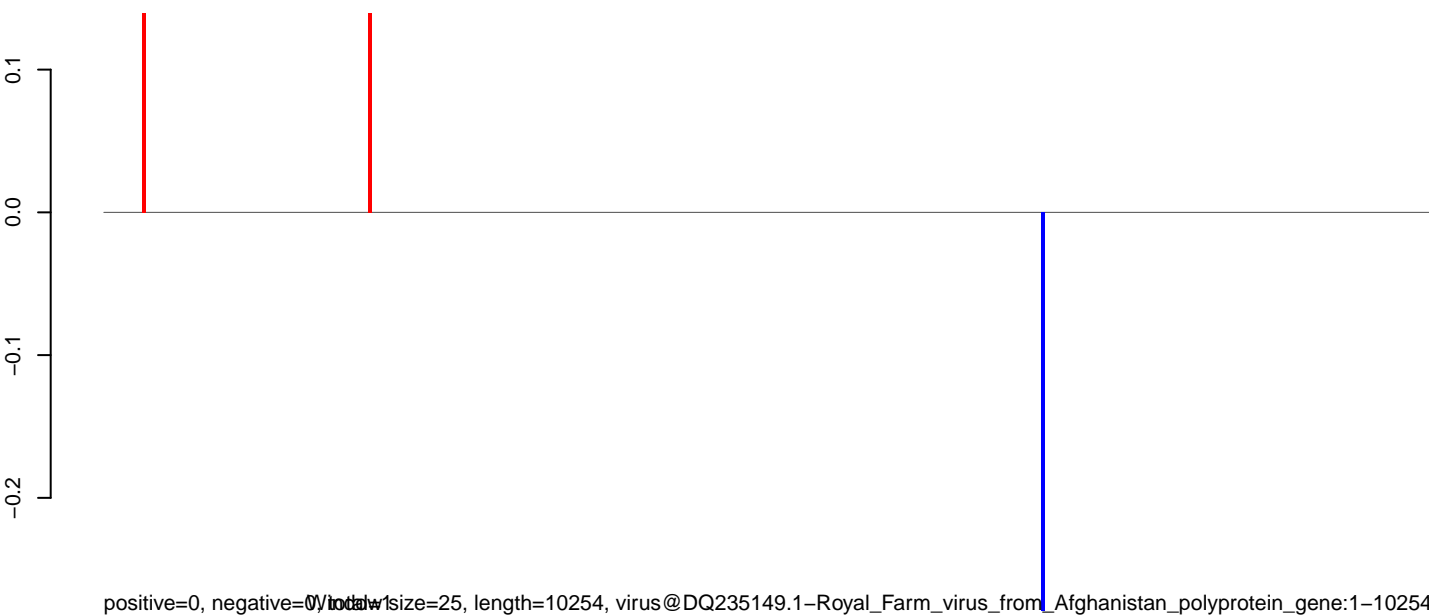
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

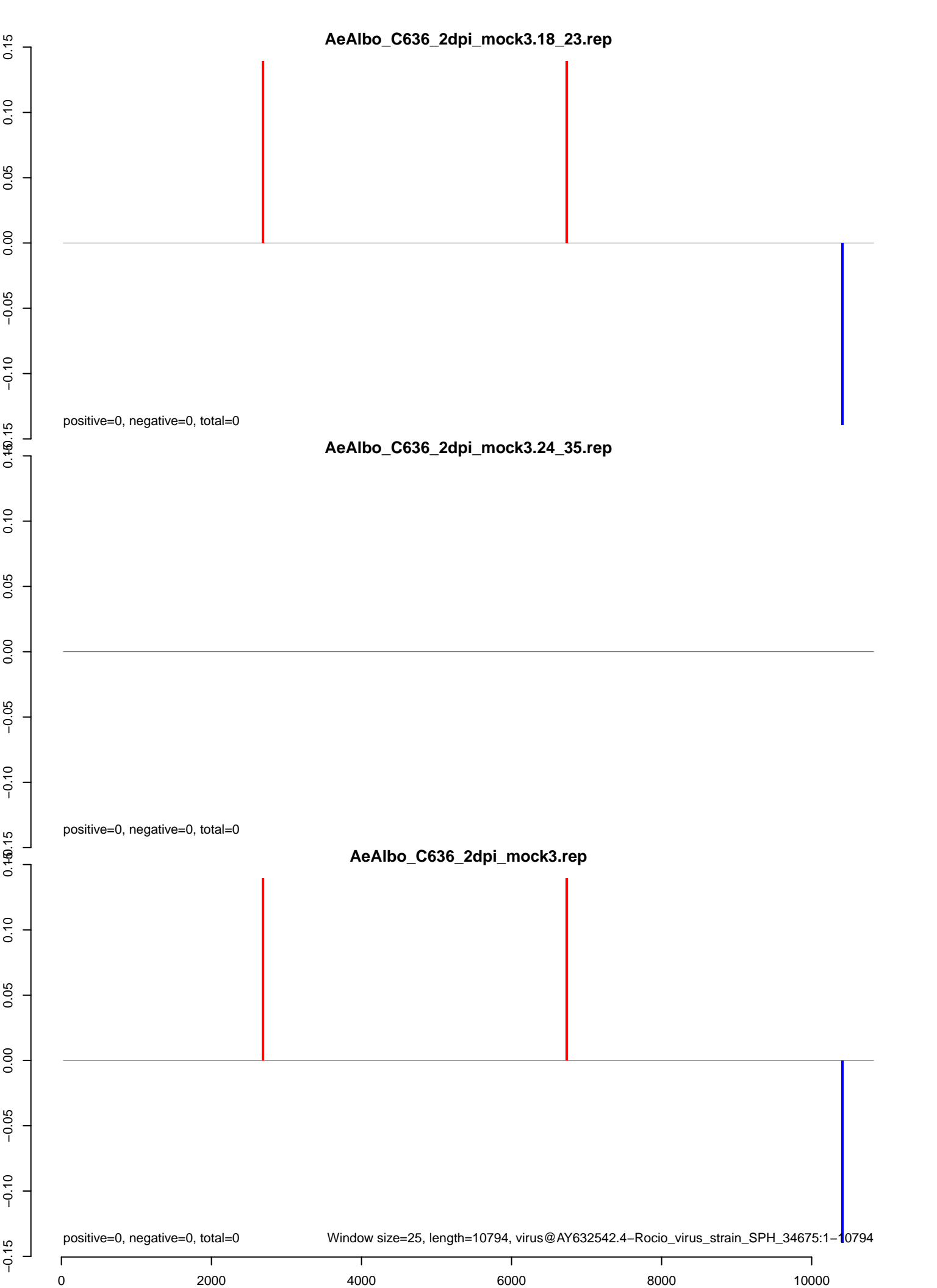


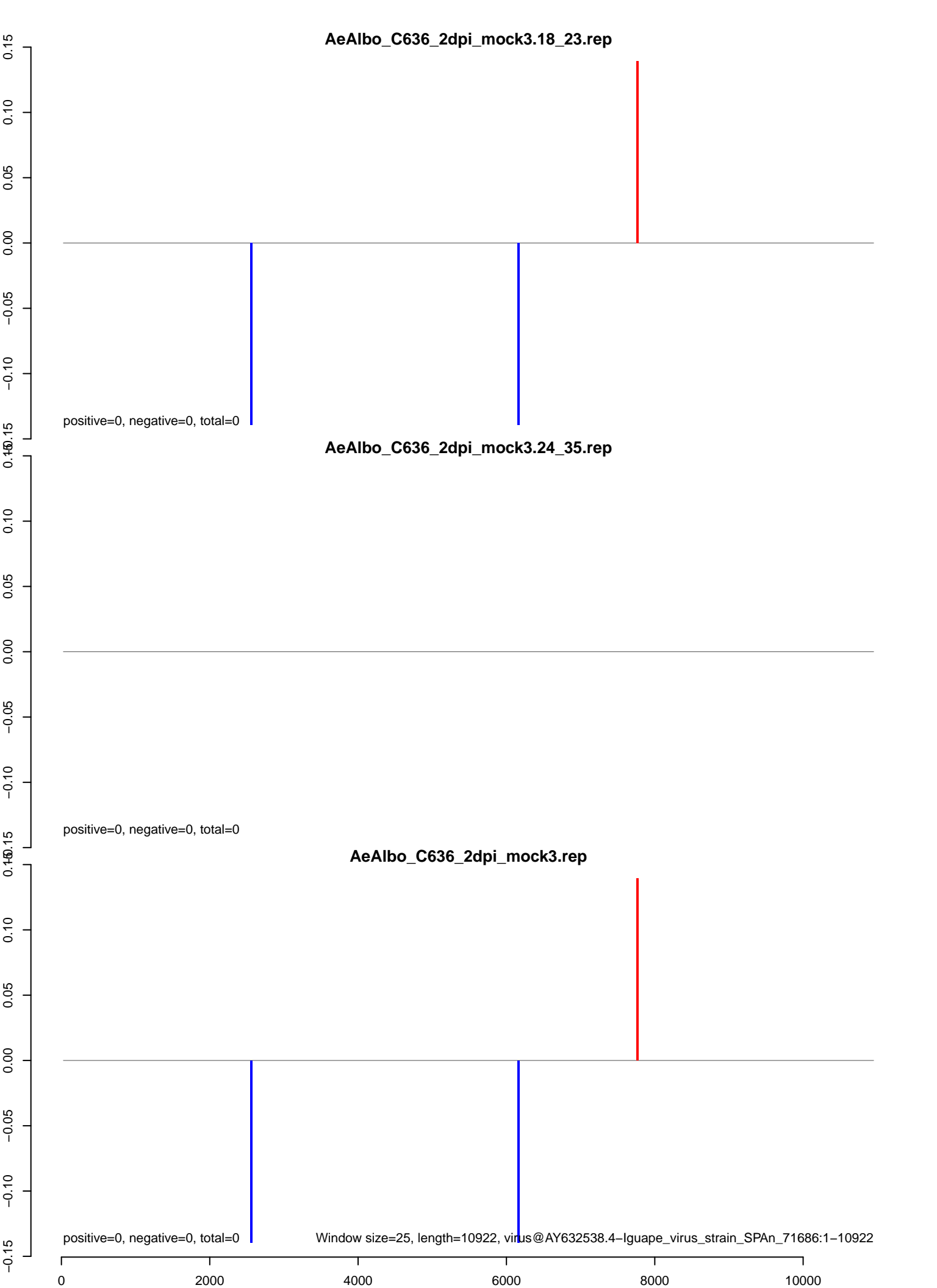
AeAlbo_C636_2dpi_mock3.rep



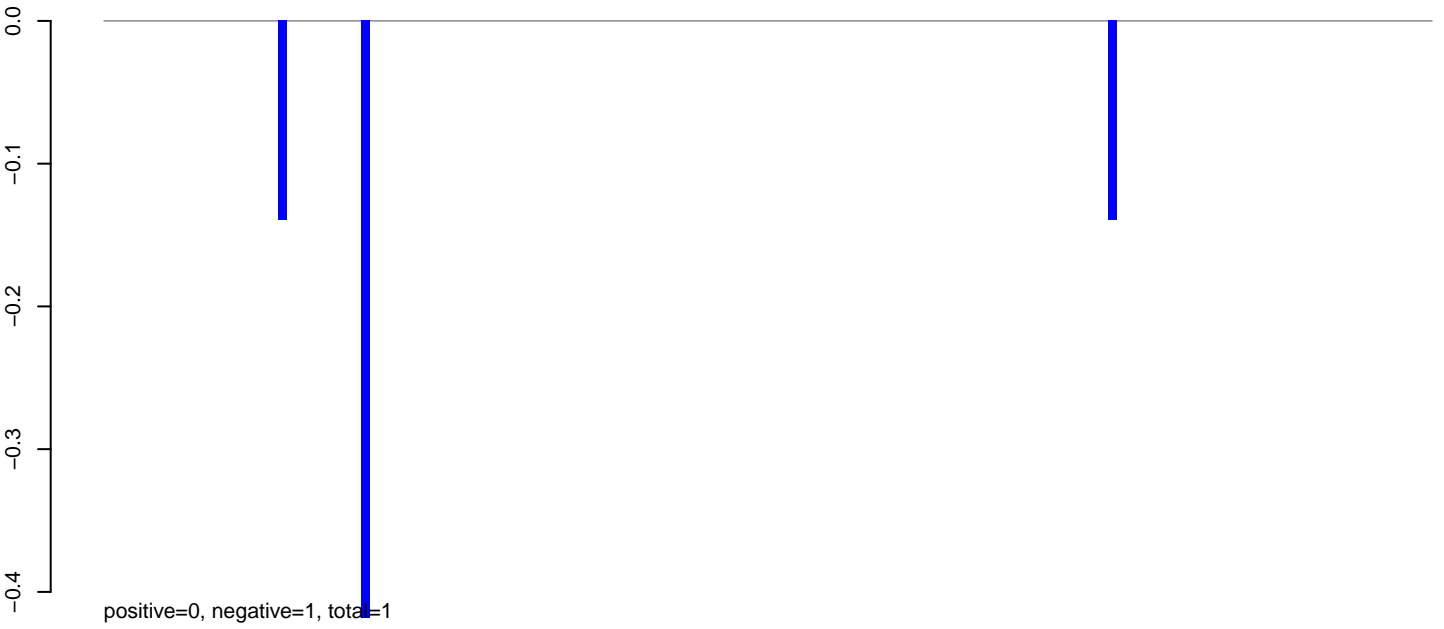
Window size=25, length=10254, virus@DQ235149.1-Royal_Farm_virus_from_Afghanistan_polyprotein_gene:1-10254

0 2000 4000 6000 8000 10000

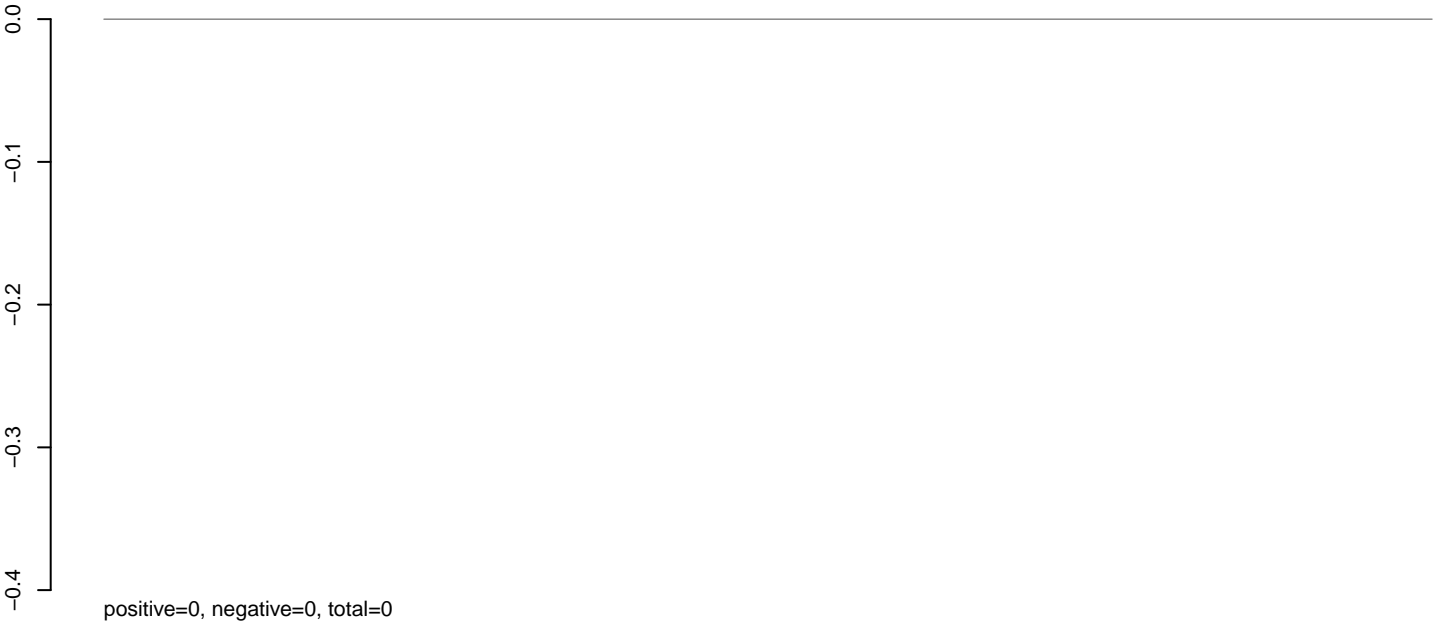




AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



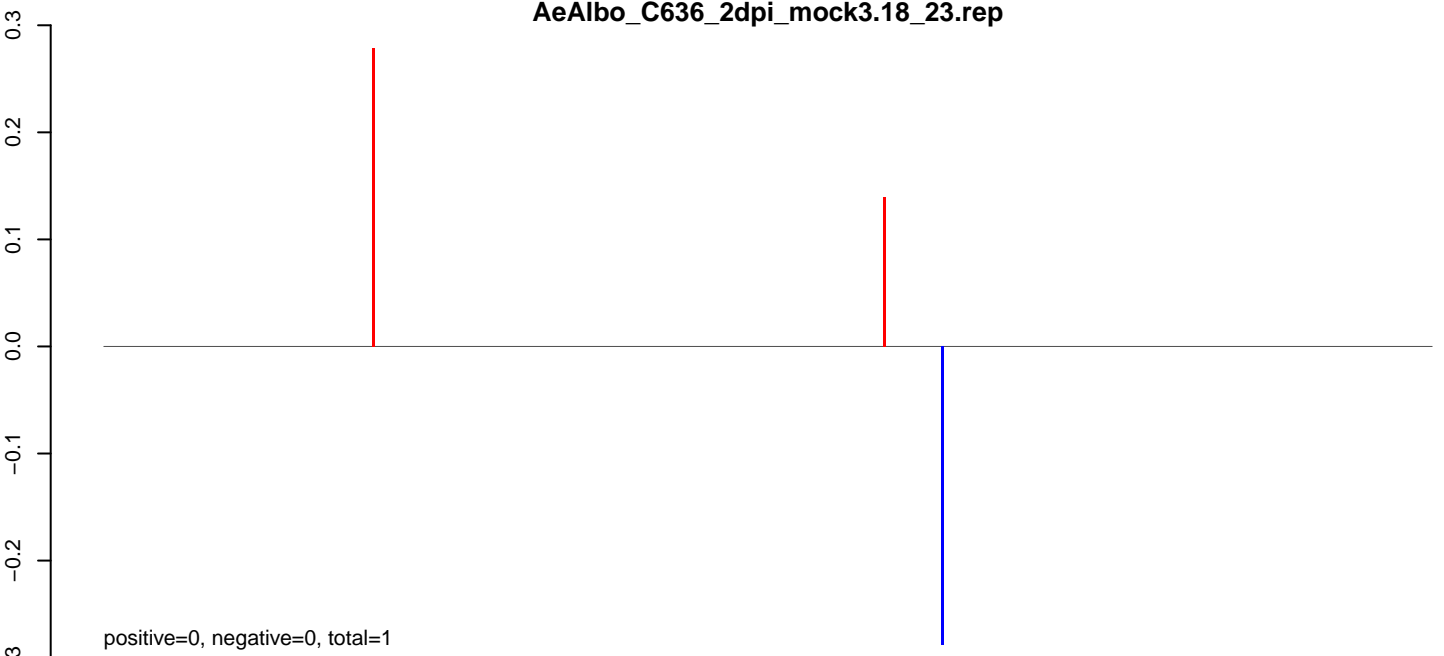
AeAlbo_C636_2dpi_mock3.rep



w size=251, length=9986, virus=Ca1605055.1-Haemagogus_equinus_densovirus_nonstructural_protein_1_nonstructural_protein_2_and_c:1-3986

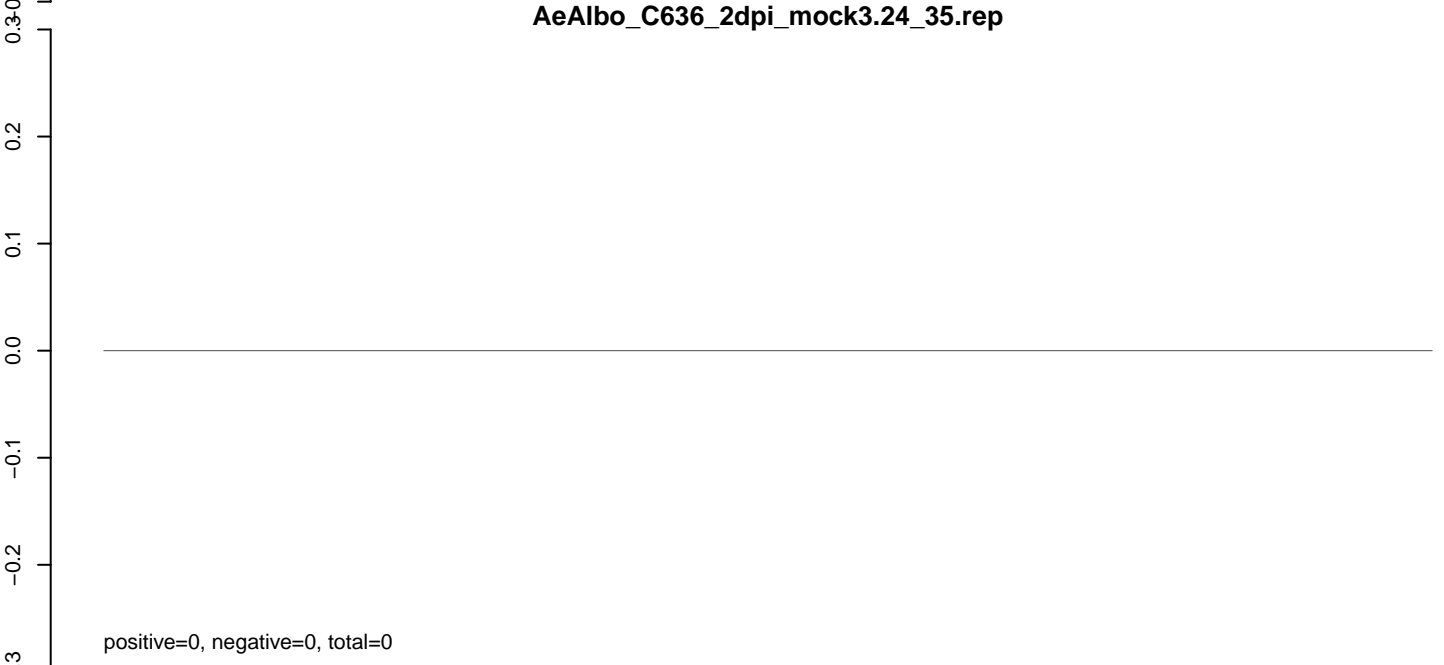
0 1000 2000 3000 4000

AeAlbo_C636_2dpi_mock3.18_23.rep



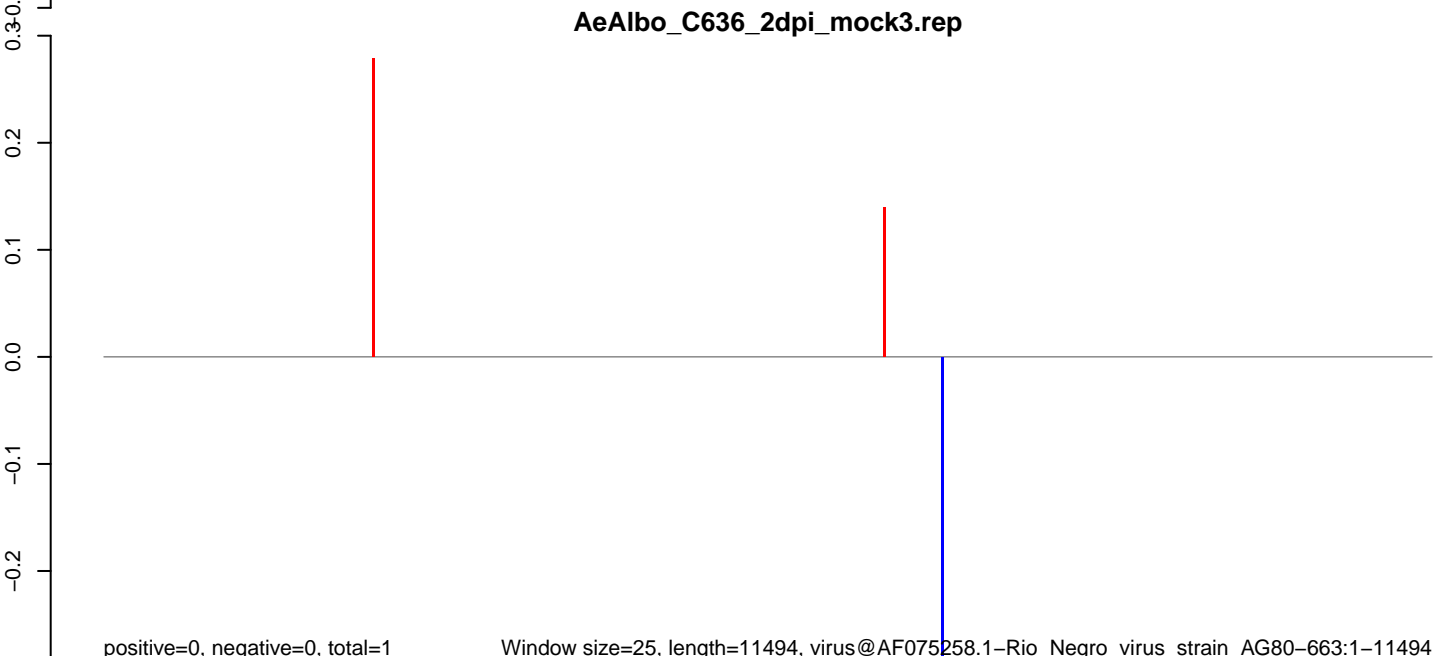
positive=0, negative=0, total=1

AeAlbo_C636_2dpi_mock3.24_35.rep



positive=0, negative=0, total=0

AeAlbo_C636_2dpi_mock3.rep

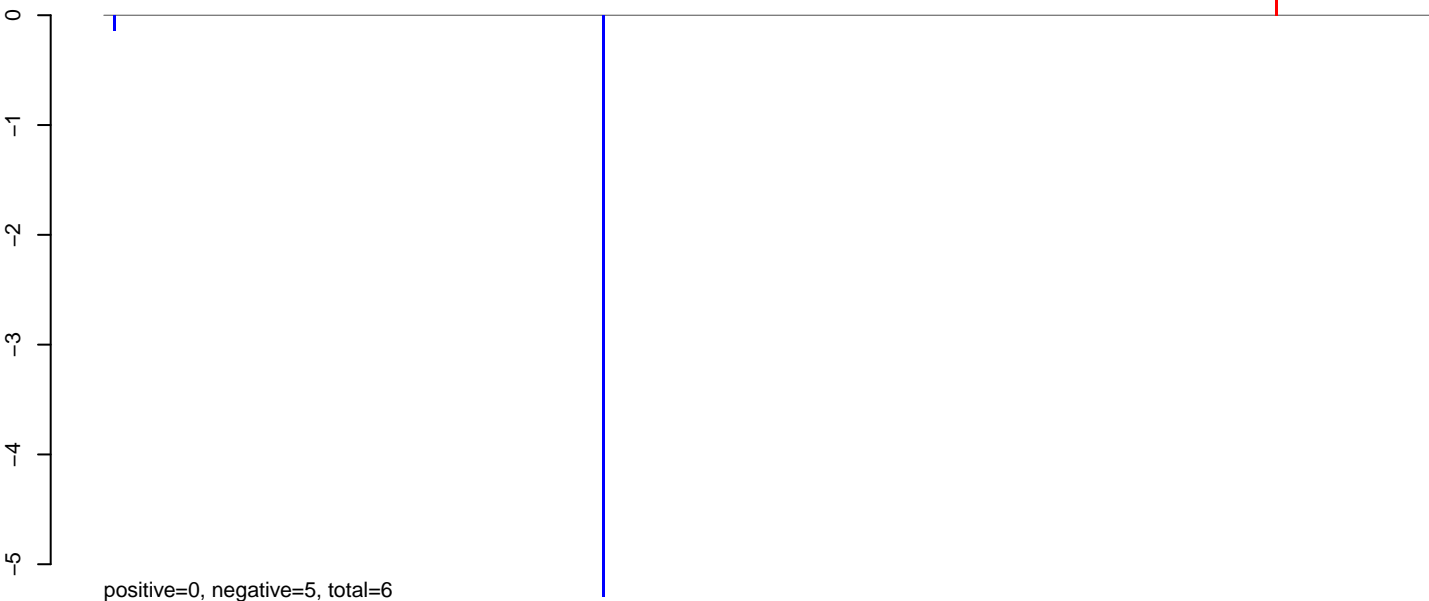


positive=0, negative=0, total=1

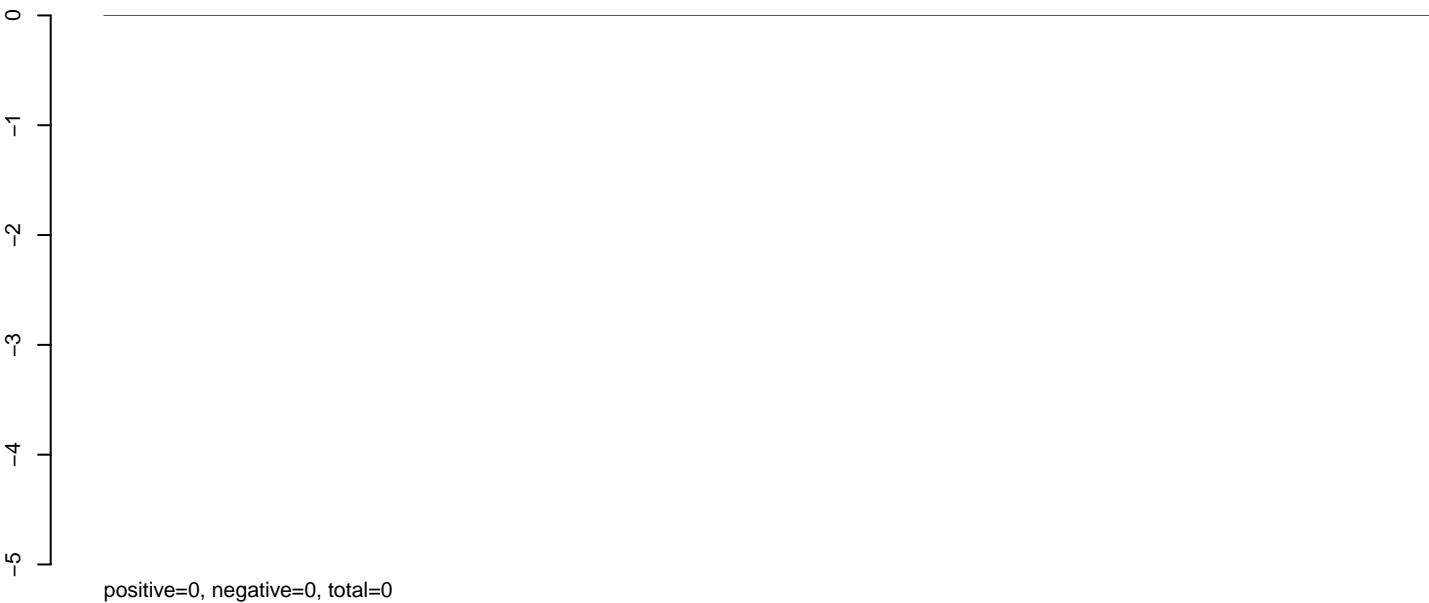
Window size=25, length=11494, virus@AF075258.1-Rio_Negro_virus_strain_AG80-663:1-11494

0 2000 4000 6000 8000 10000

AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



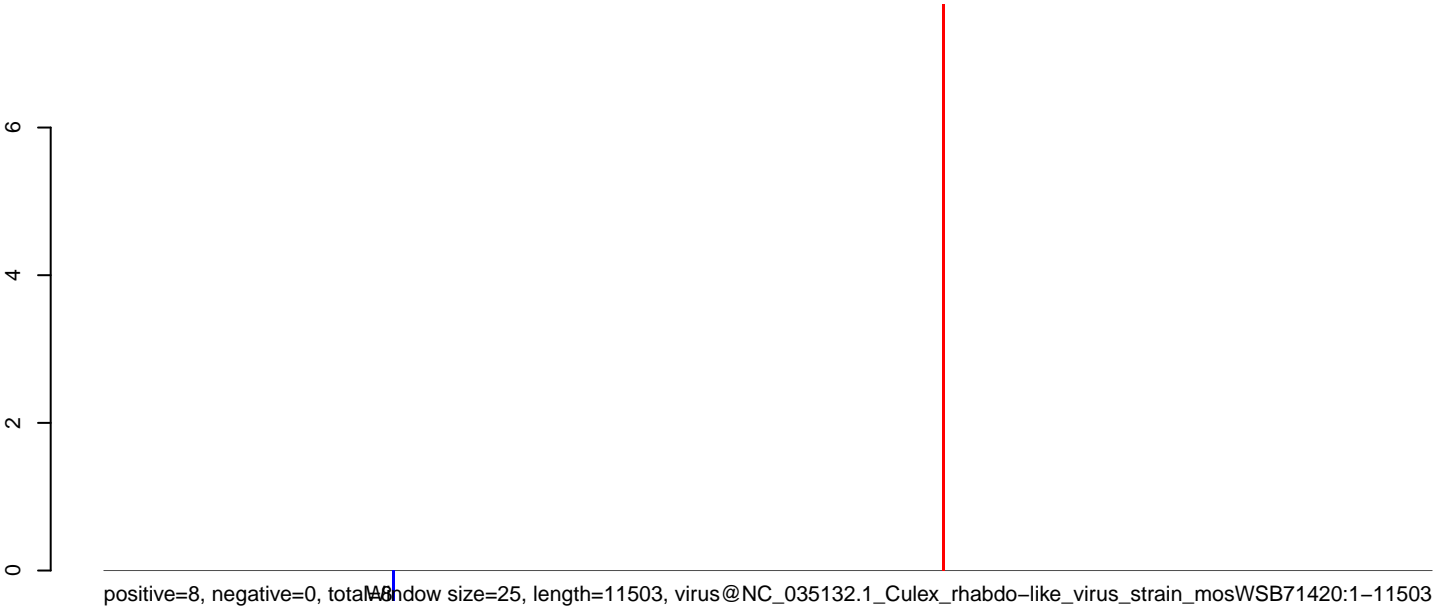
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

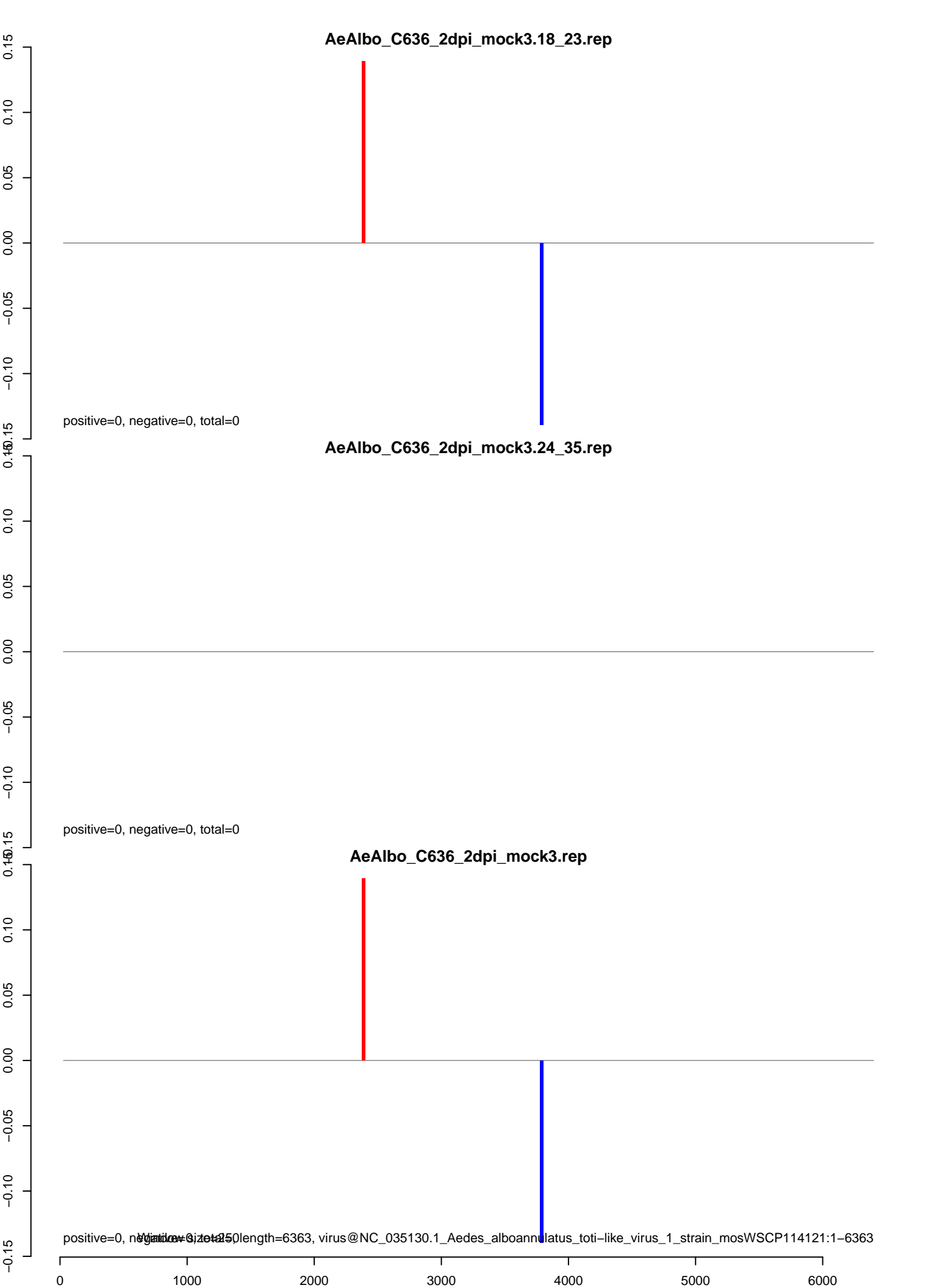


AeAlbo_C636_2dpi_mock3.rep



Window size=25, length=11503, virus@NC_035132.1_Culex_rhabdo-like_virus_strain_mosWSB71420:1-11503

0 2000 4000 6000 8000 10000 12000



AeAlbo_C636_2dpi_mock3.18_23.rep

positive=0, negative=0, total=0

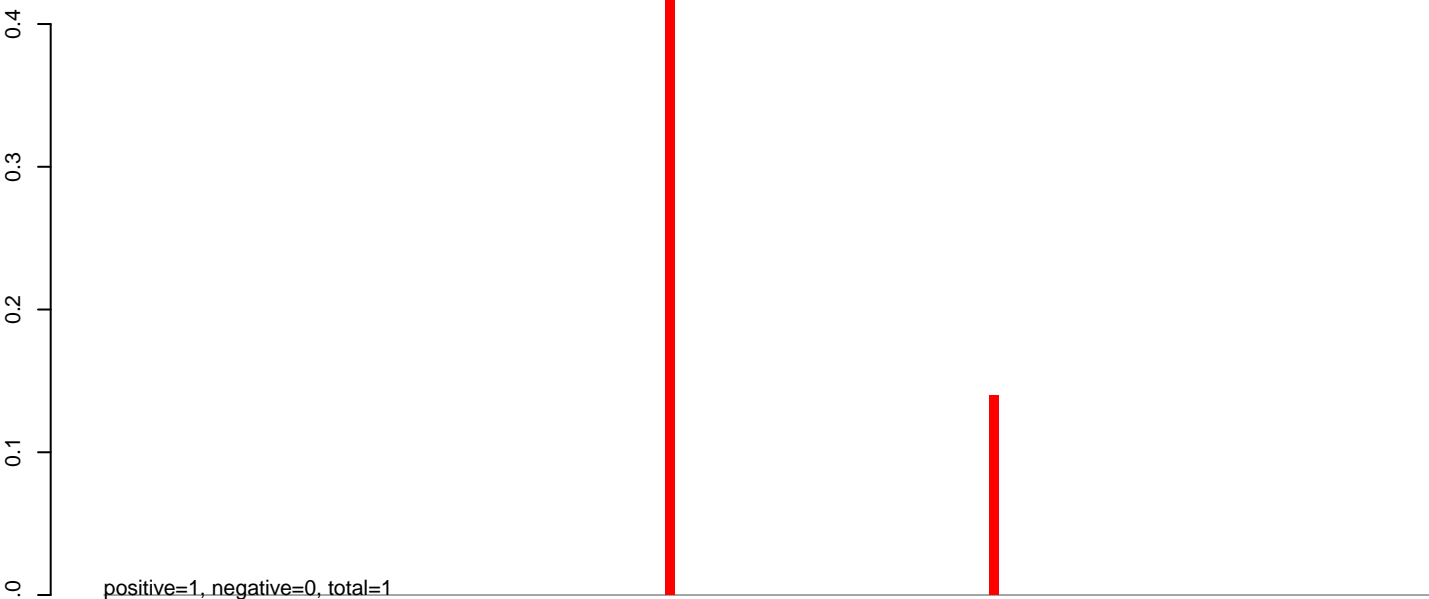
AeAlbo_C636_2dpi_mock3.24_35.rep

positive=0, negative=0, total=0

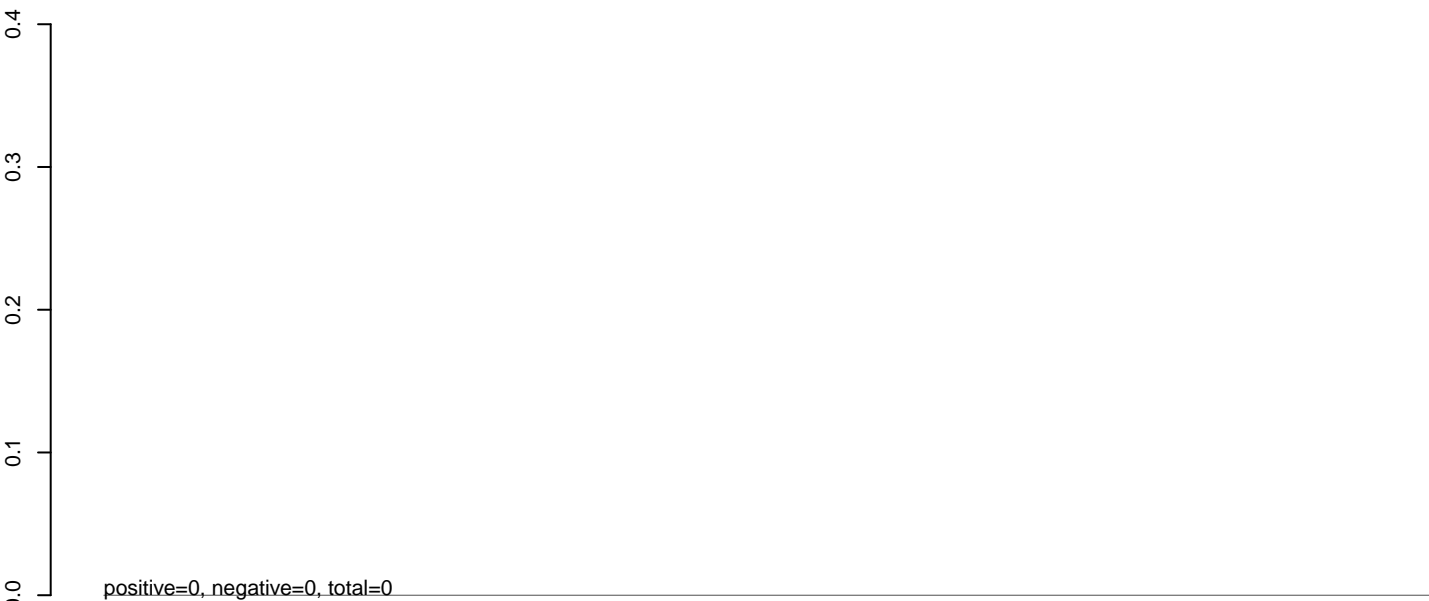
AeAlbo_C636_2dpi_mock3.rep

positive=0, negative=0, total=0
Window Size=25 length=6363, virus@NC_035130.1_Aedes_alboannulatus_toti-like_virus_1_strain_mosWSCP114121:1-6363

AeAlbo_C636_2dpi_mock3.18_23.rep



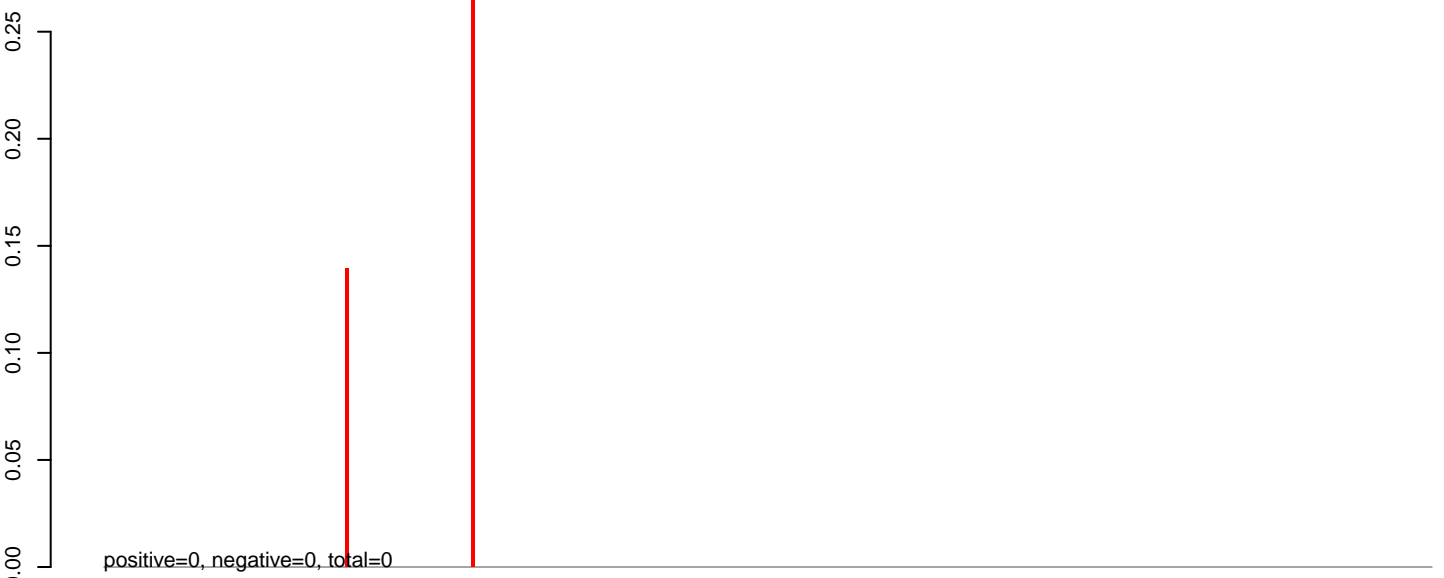
AeAlbo_C636_2dpi_mock3.24_35.rep



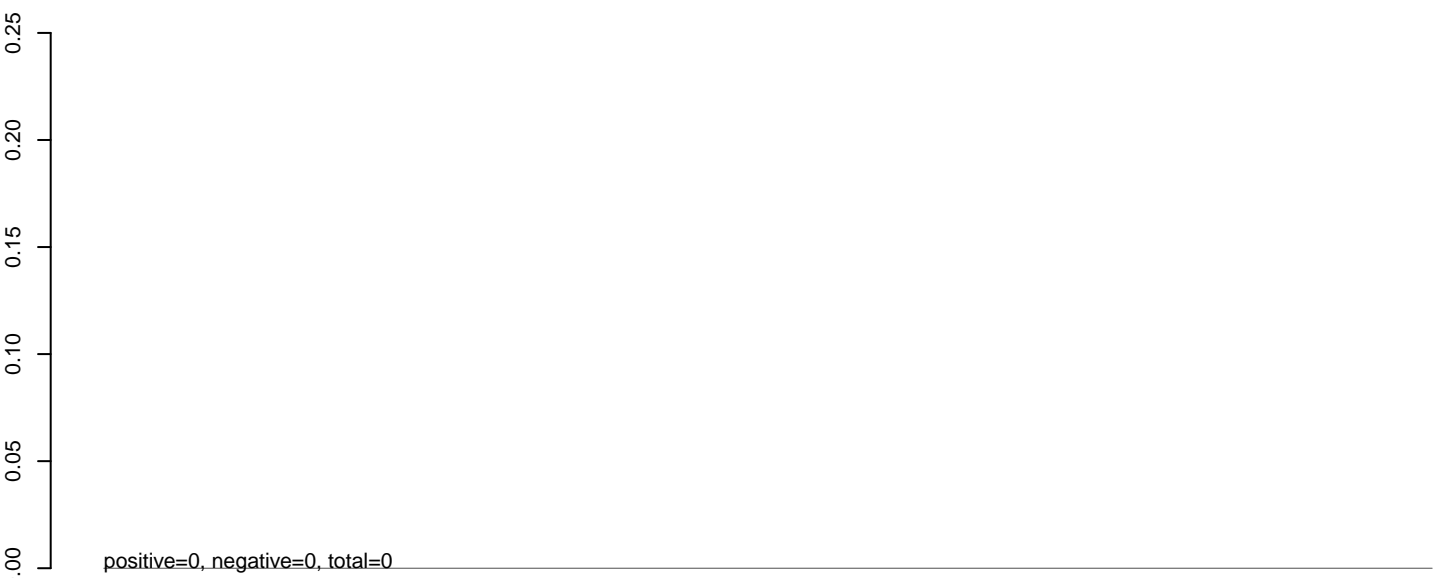
AeAlbo_C636_2dpi_mock3.rep



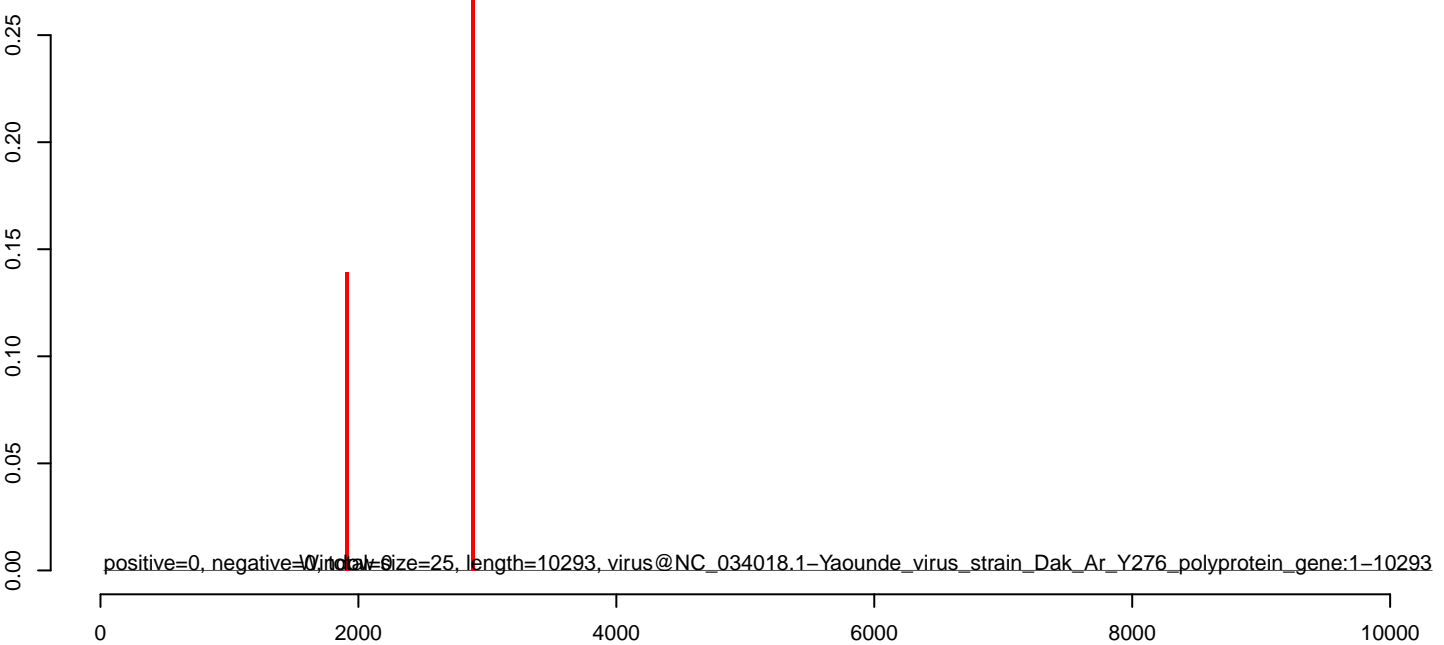
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



AeAlbo_C636_2dpi_mock3.18_23.rep

positive=6, negative=0, total=6

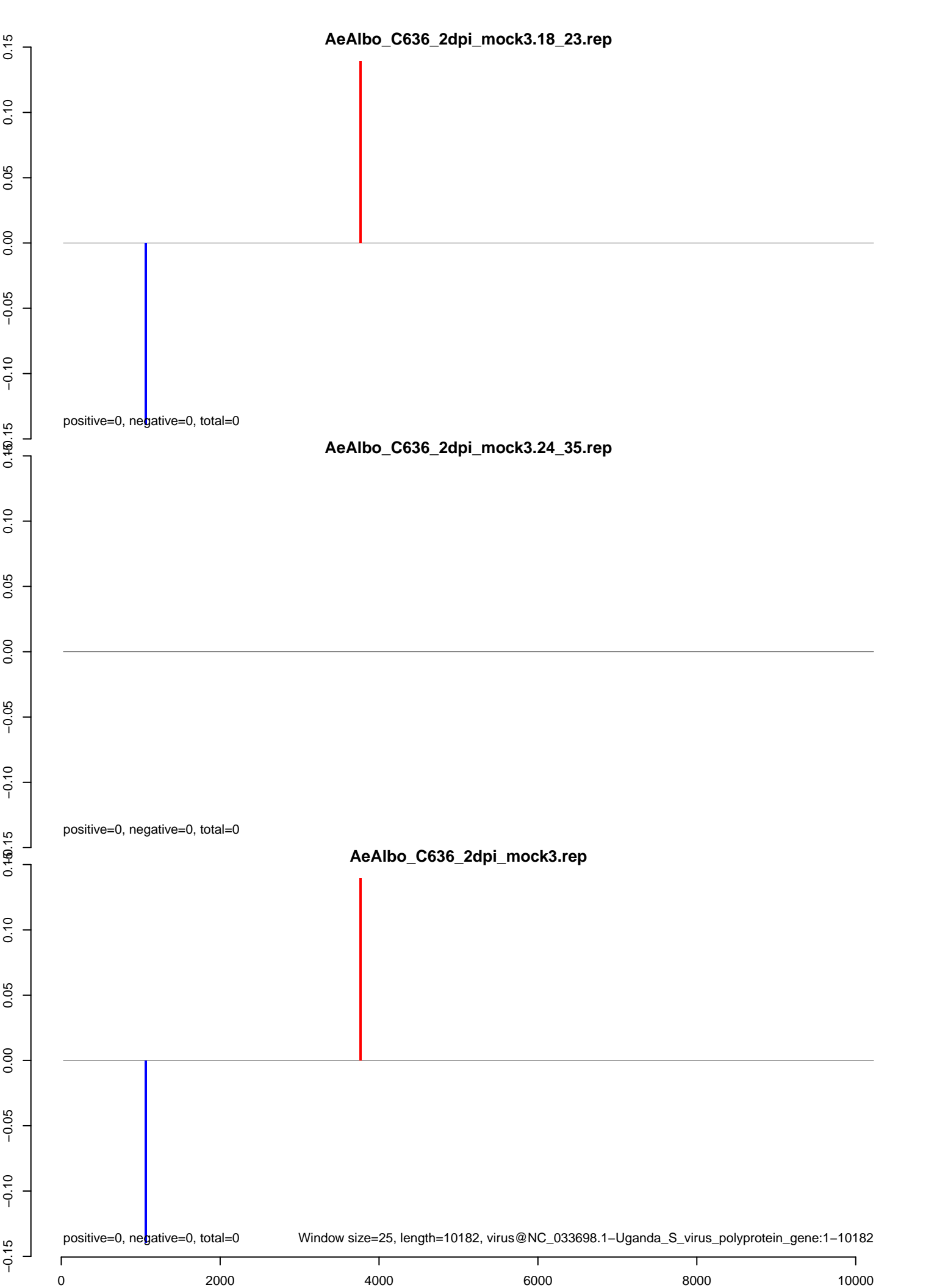
AeAlbo_C636_2dpi_mock3.24_35.rep

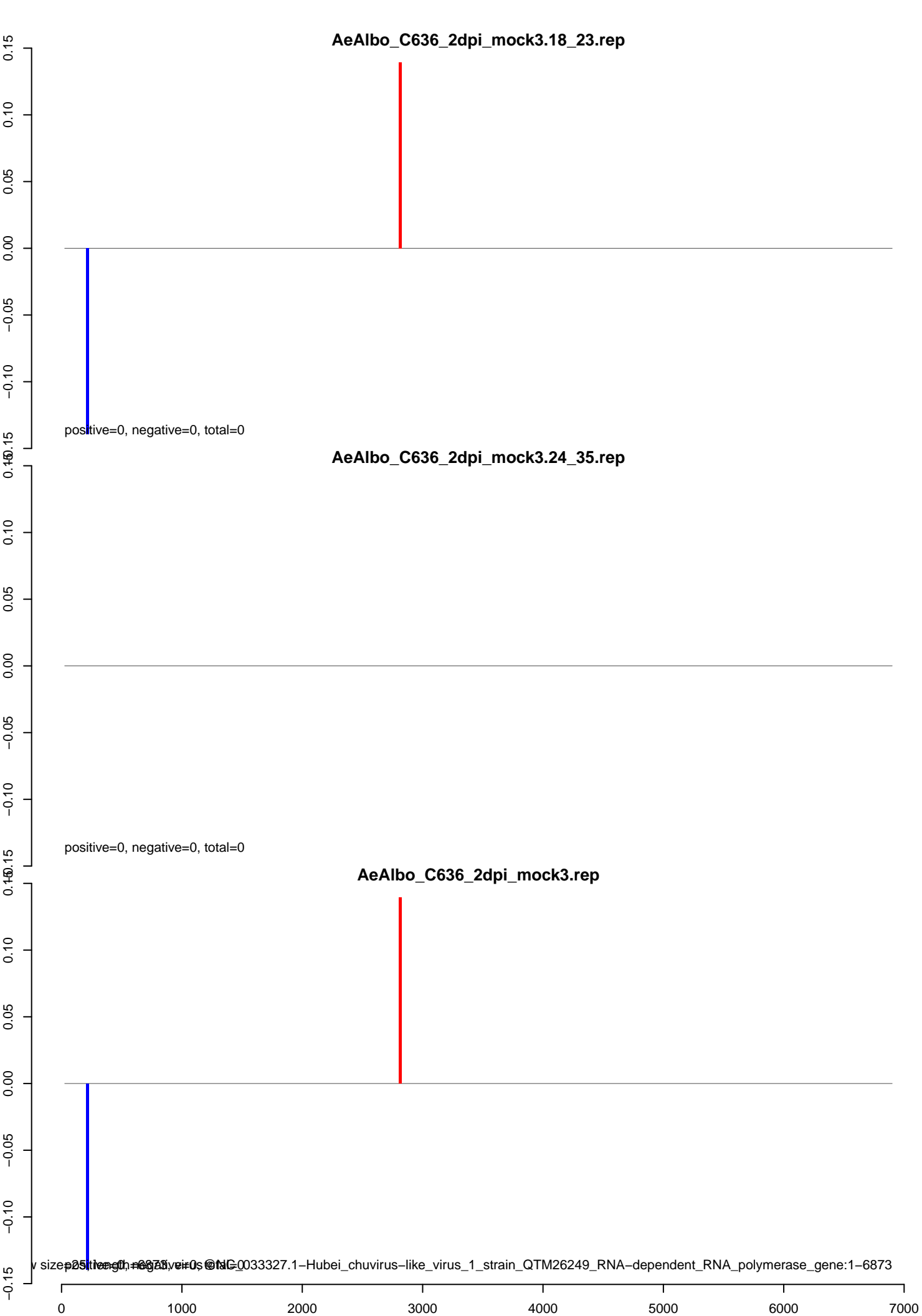
positive=0, negative=0, total=0

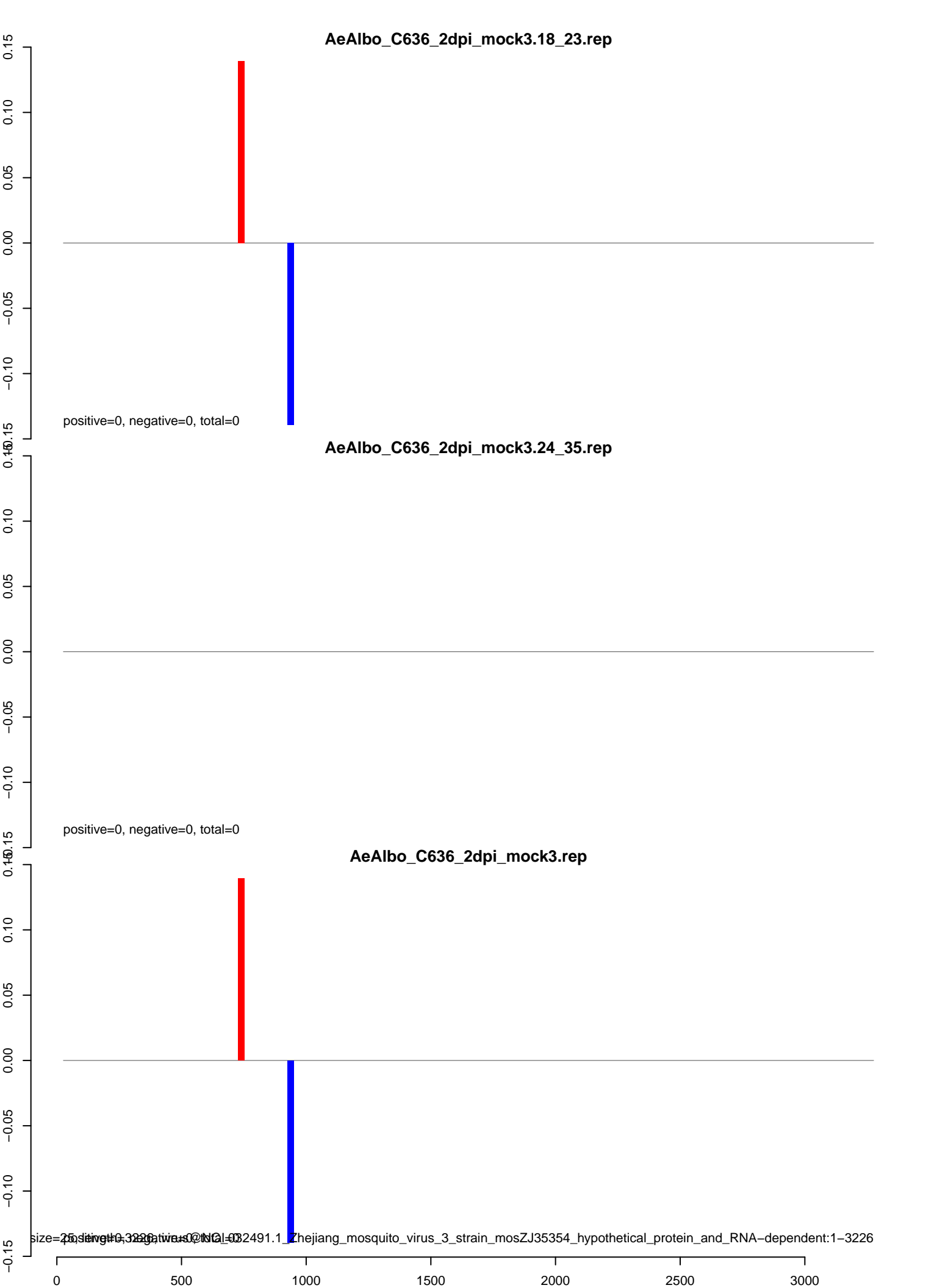
AeAlbo_C636_2dpi_mock3.rep

positive=6, negative=0, total=6, virus@NC_033716.1_Zhejiang_mosquito_virus_1_strain_mosZJ35497_hypothetical_protein_gene:1-9558

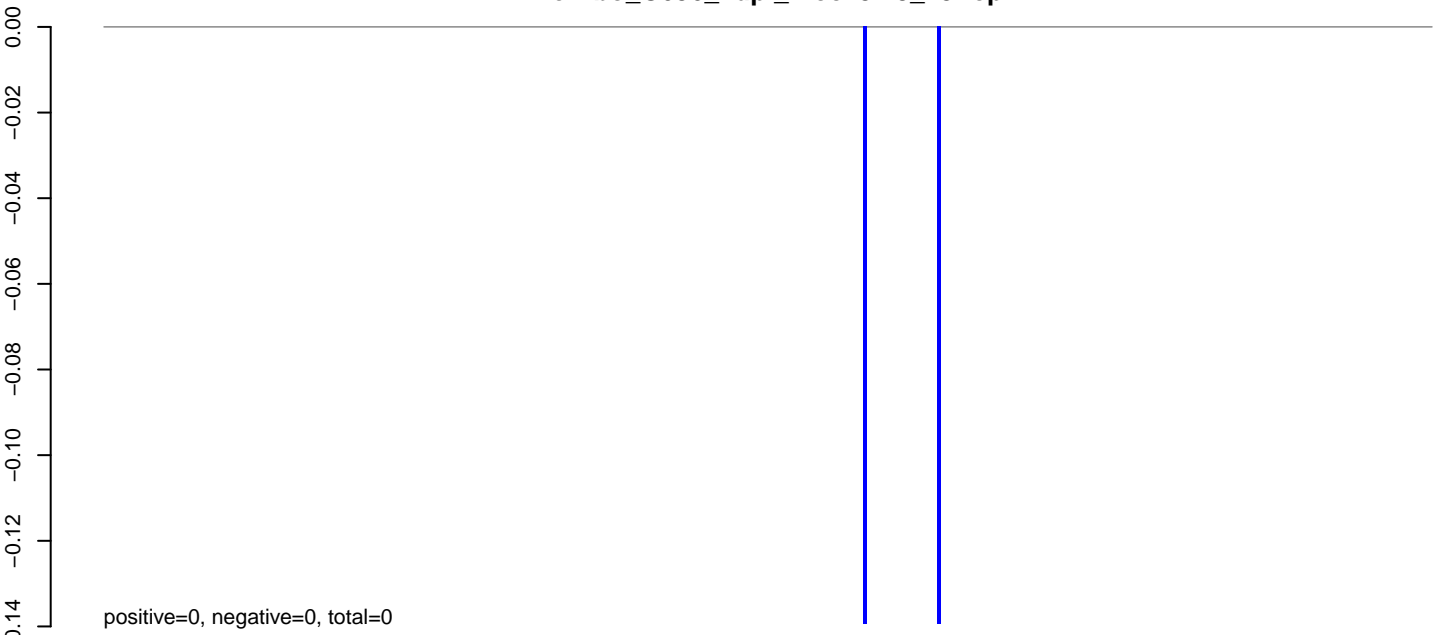
0 2000 4000 6000 8000



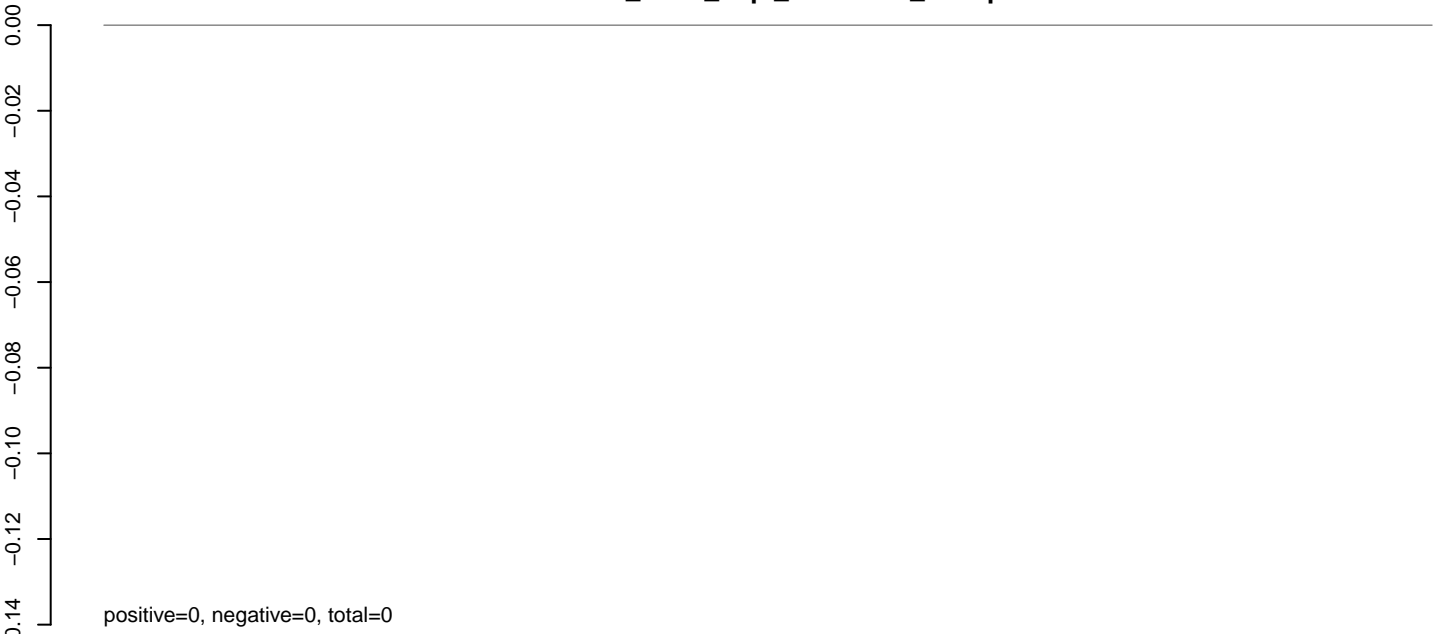




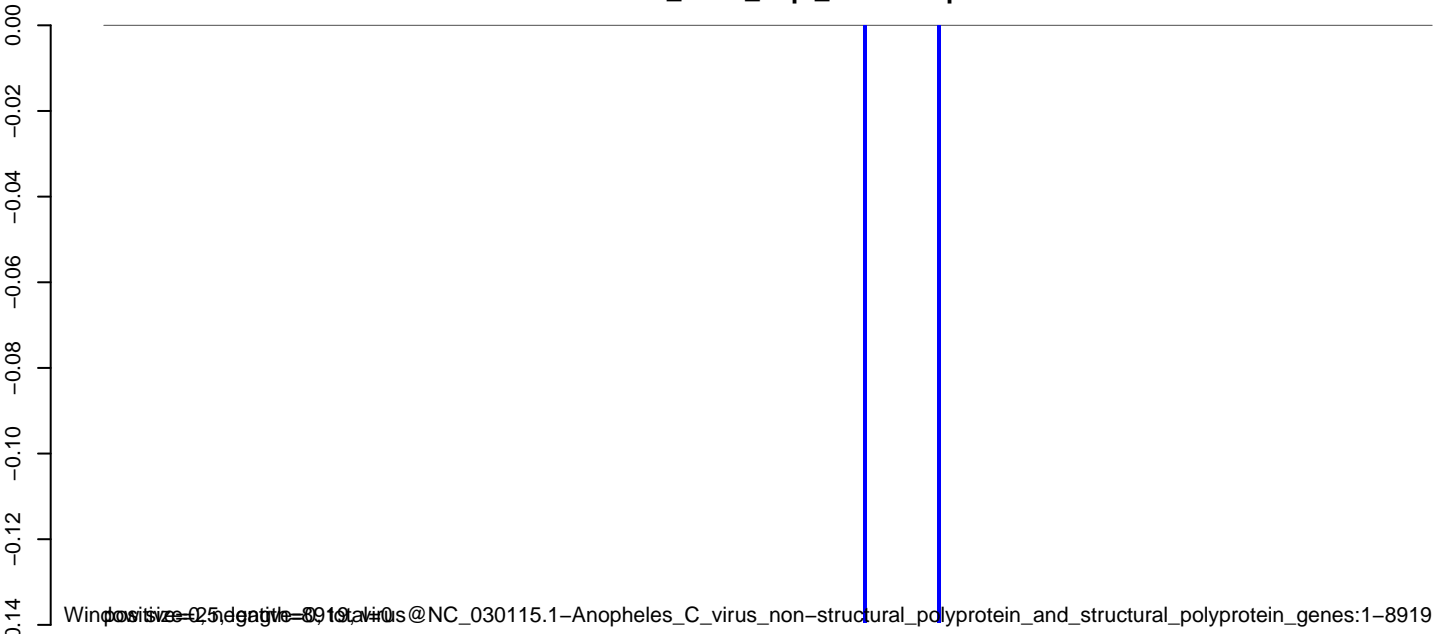
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

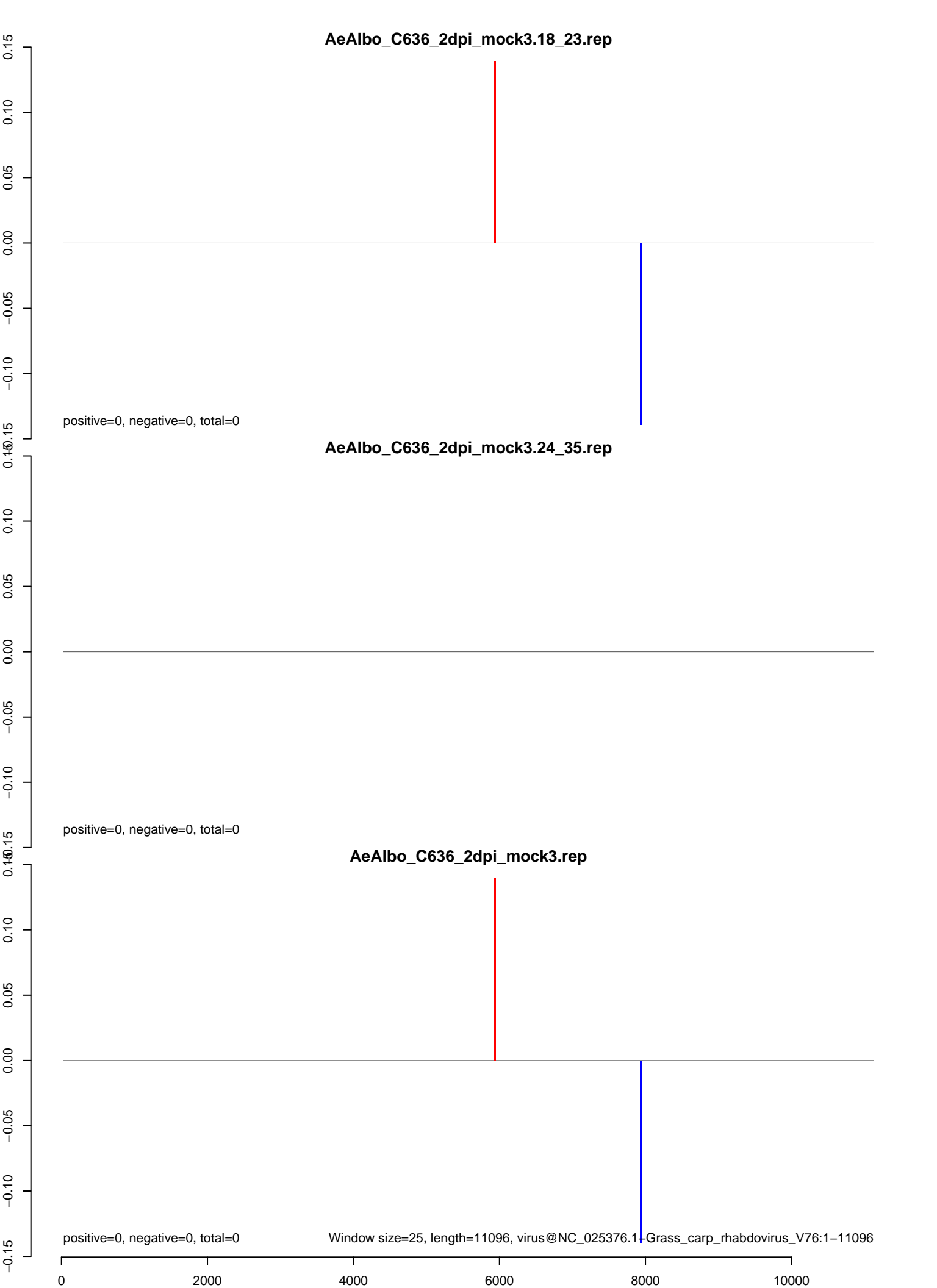


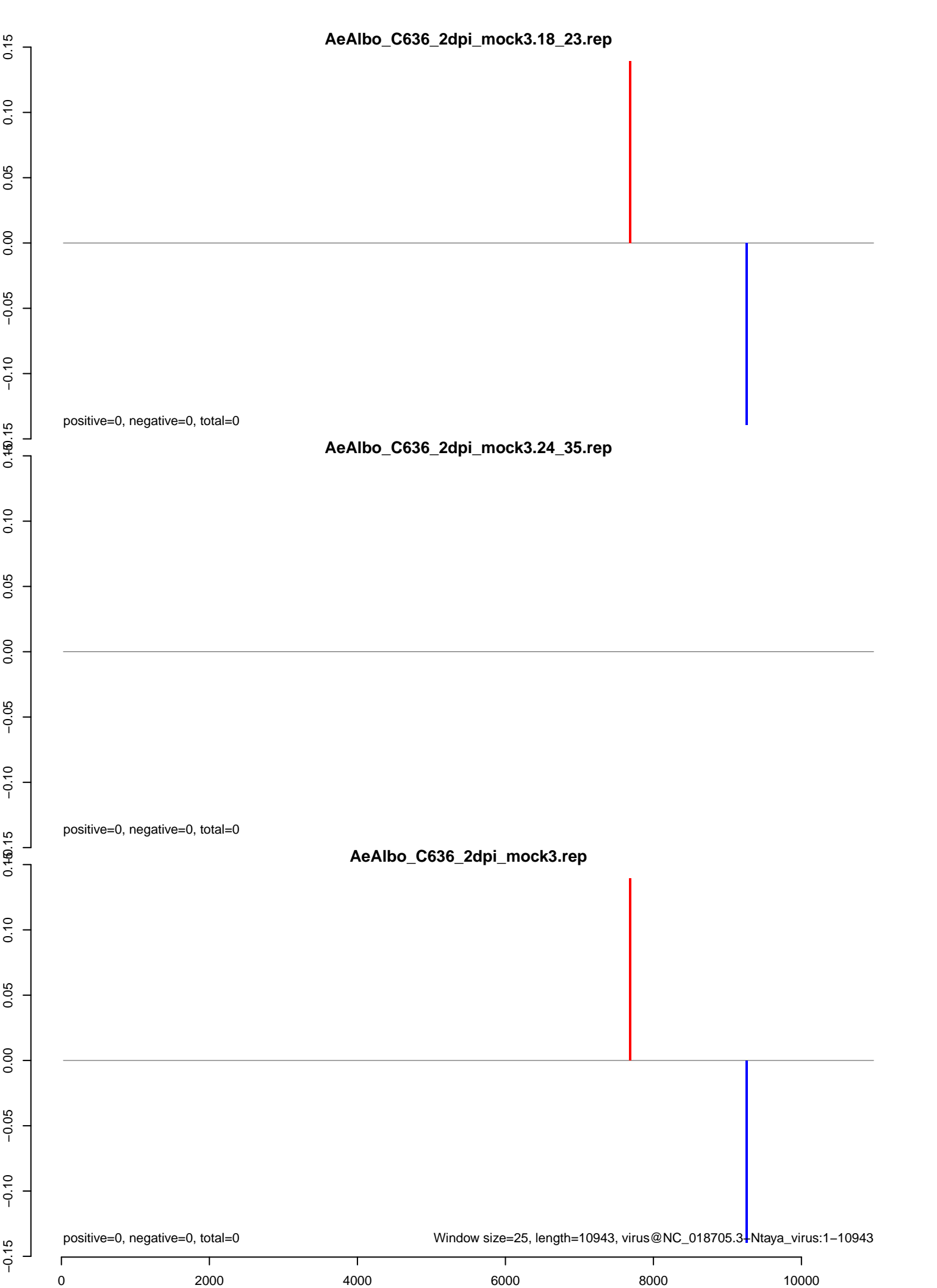
AeAlbo_C636_2dpi_mock3.rep



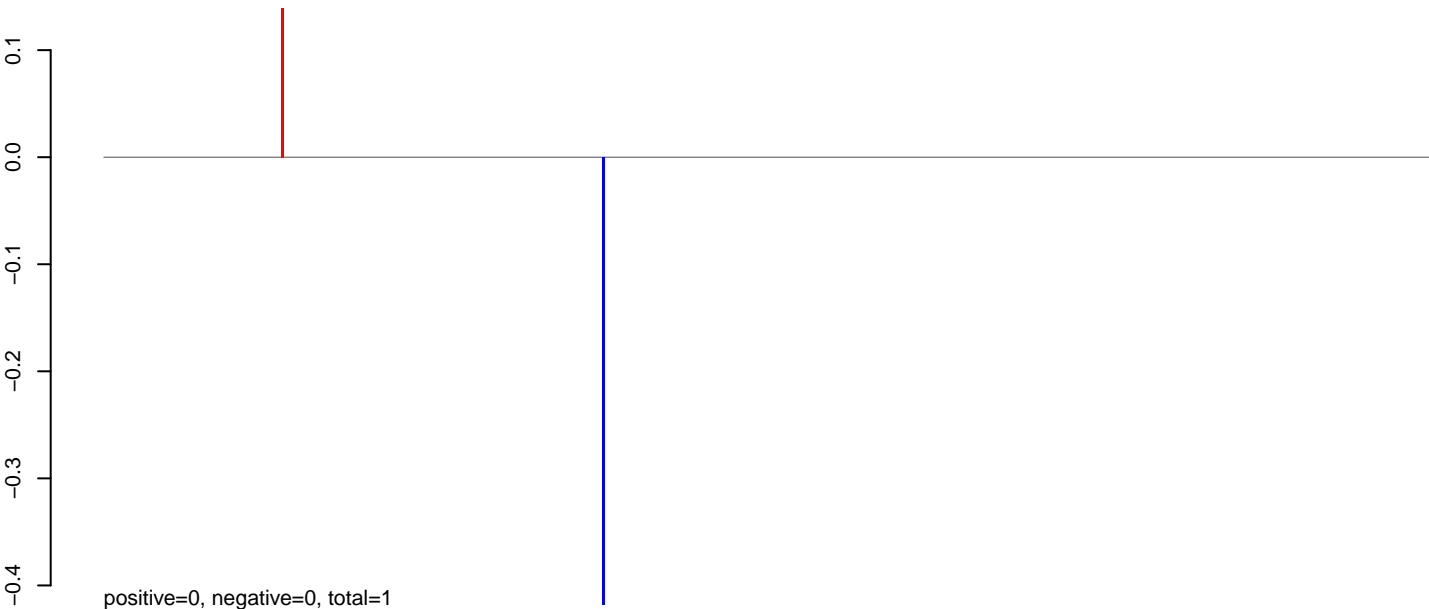
Window size=25, negative=8919, total=0 @ NC_030115.1-Anopheles_C_virus_non-structural_polyprotein_and_structural_polyprotein_genes:1-8919

0 2000 4000 6000 8000

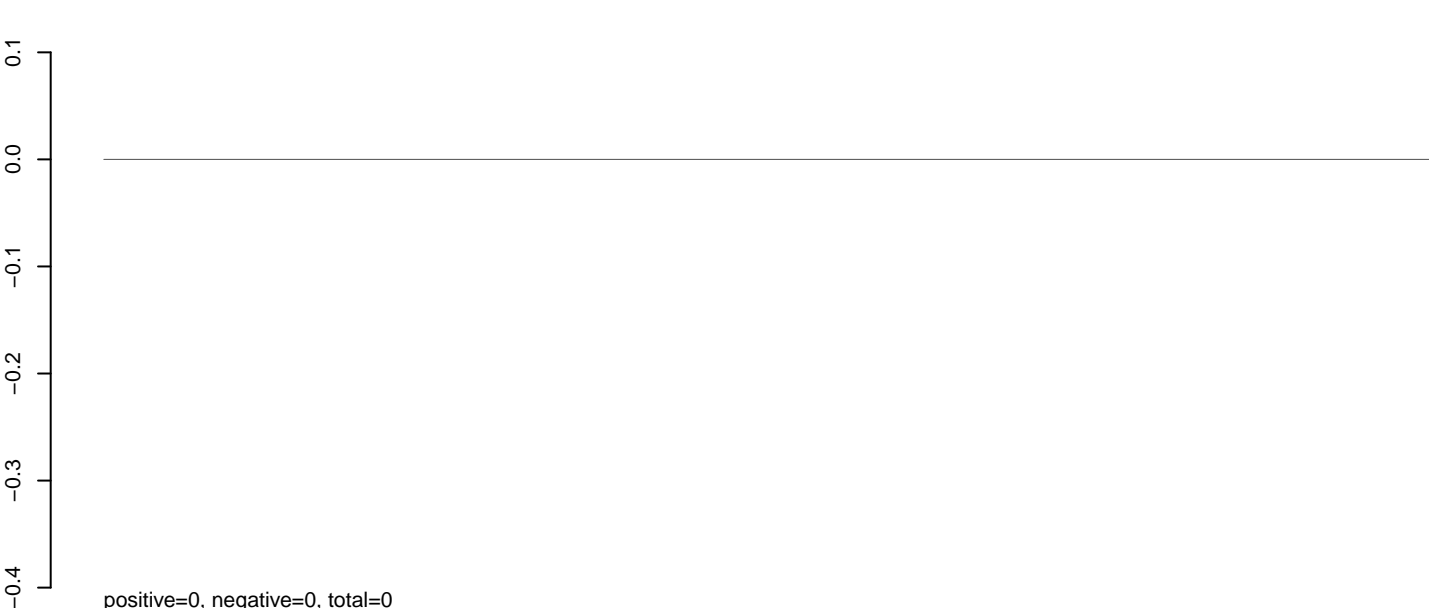




AeAlbo_C636_2dpi_mock3.18_23.rep



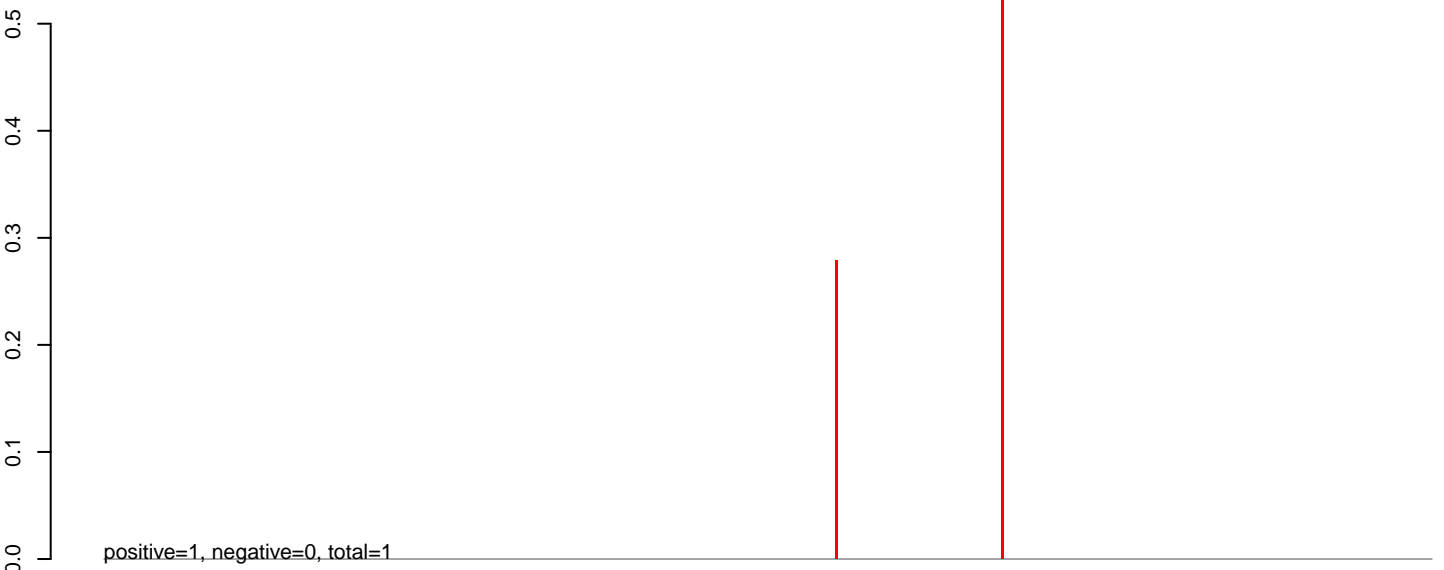
AeAlbo_C636_2dpi_mock3.24_35.rep



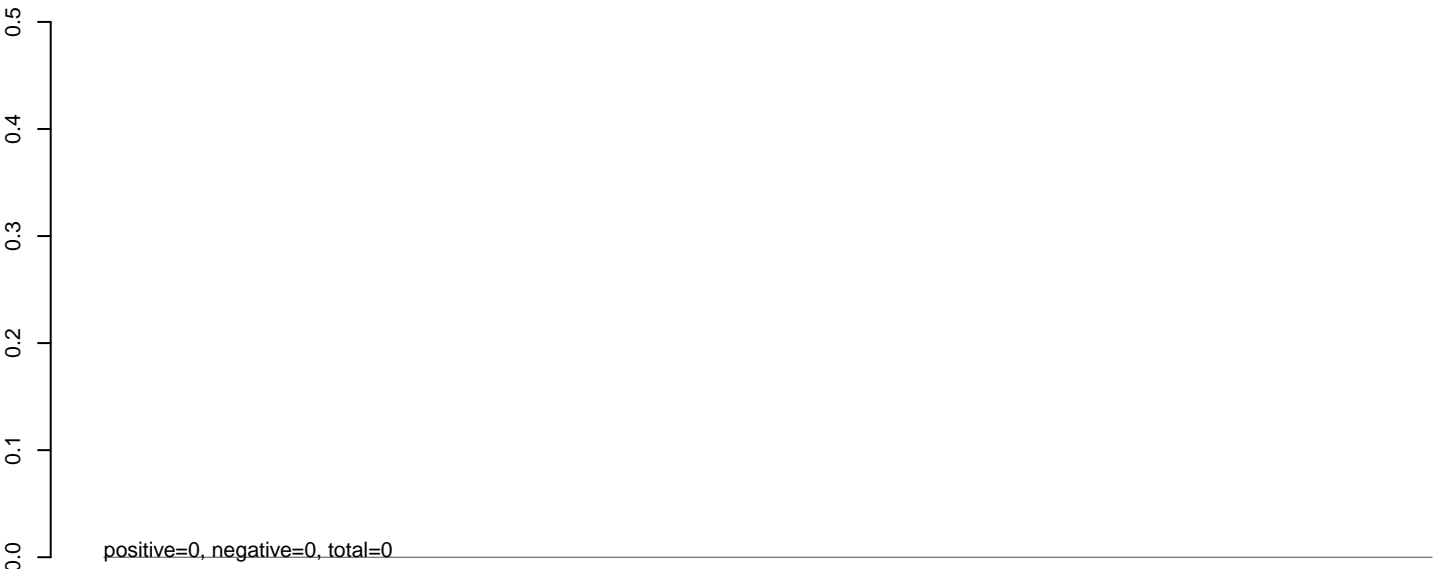
AeAlbo_C636_2dpi_mock3.rep



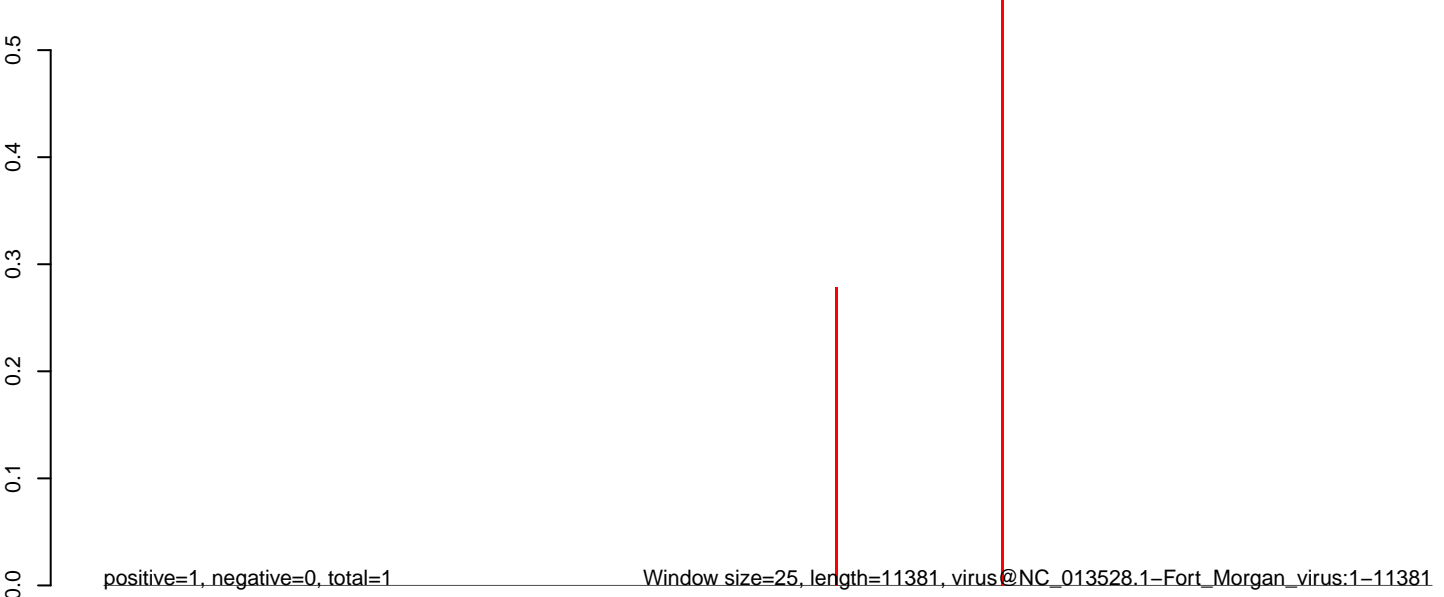
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

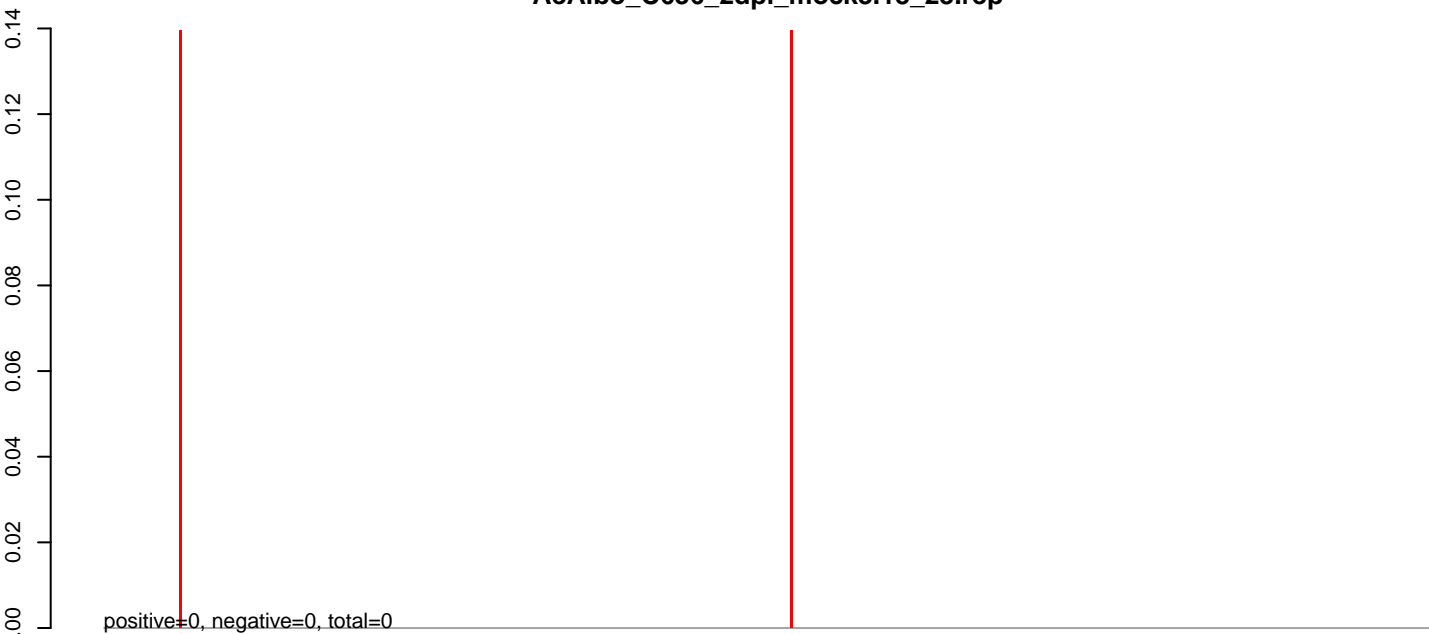


AeAlbo_C636_2dpi_mock3.rep



0 2000 4000 6000 8000 10000

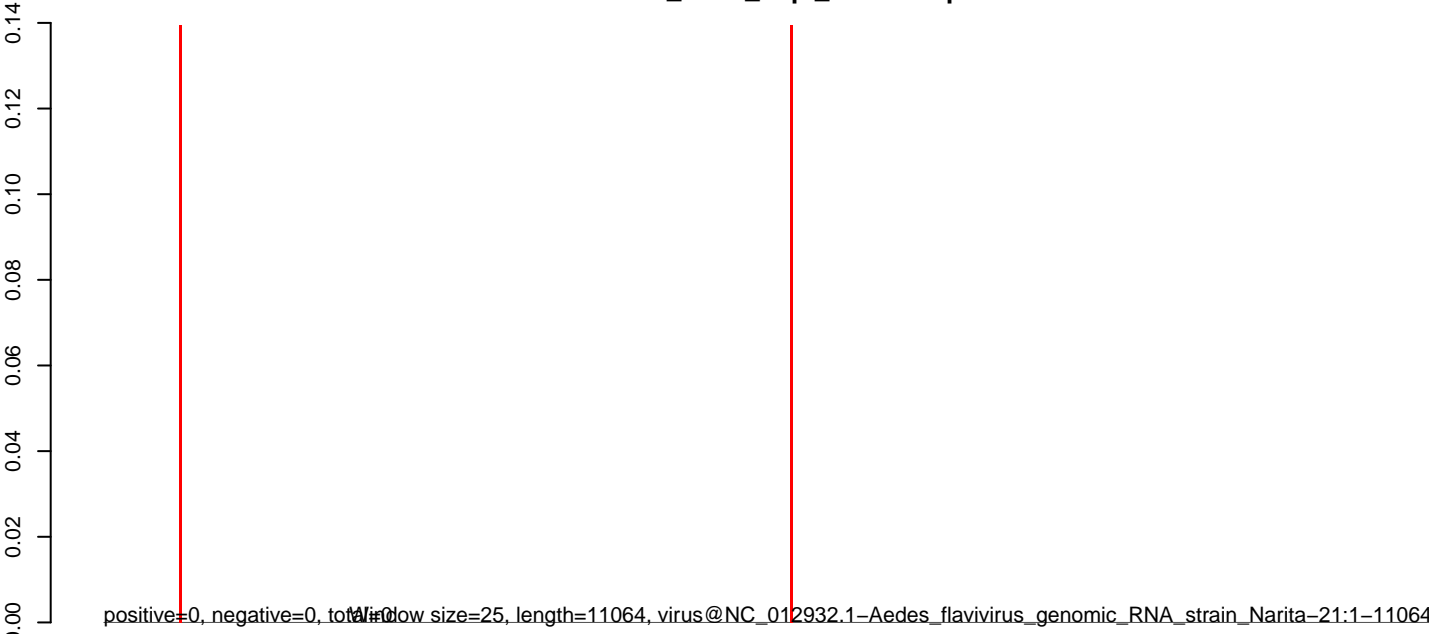
AeAlbo_C636_2dpi_mock3.18_23.rep



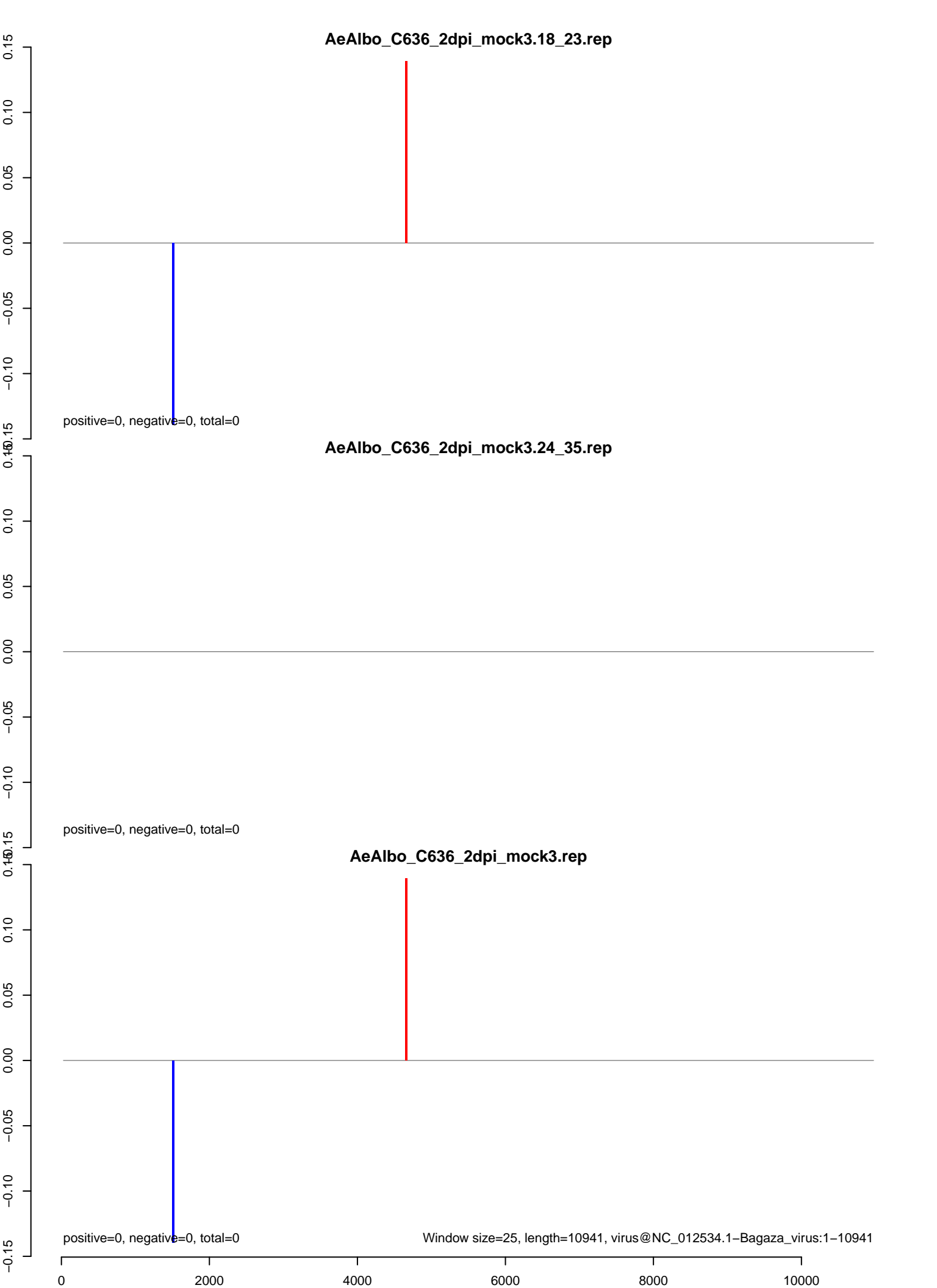
AeAlbo_C636_2dpi_mock3.24_35.rep



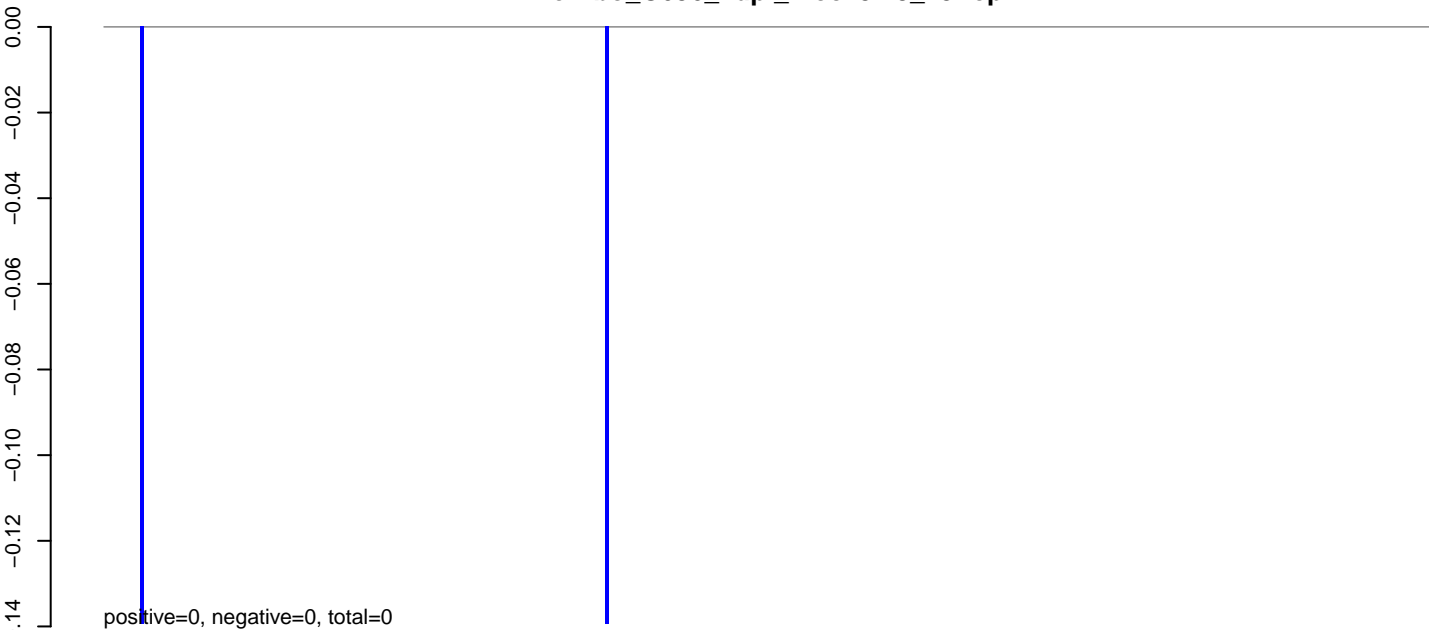
AeAlbo_C636_2dpi_mock3.rep



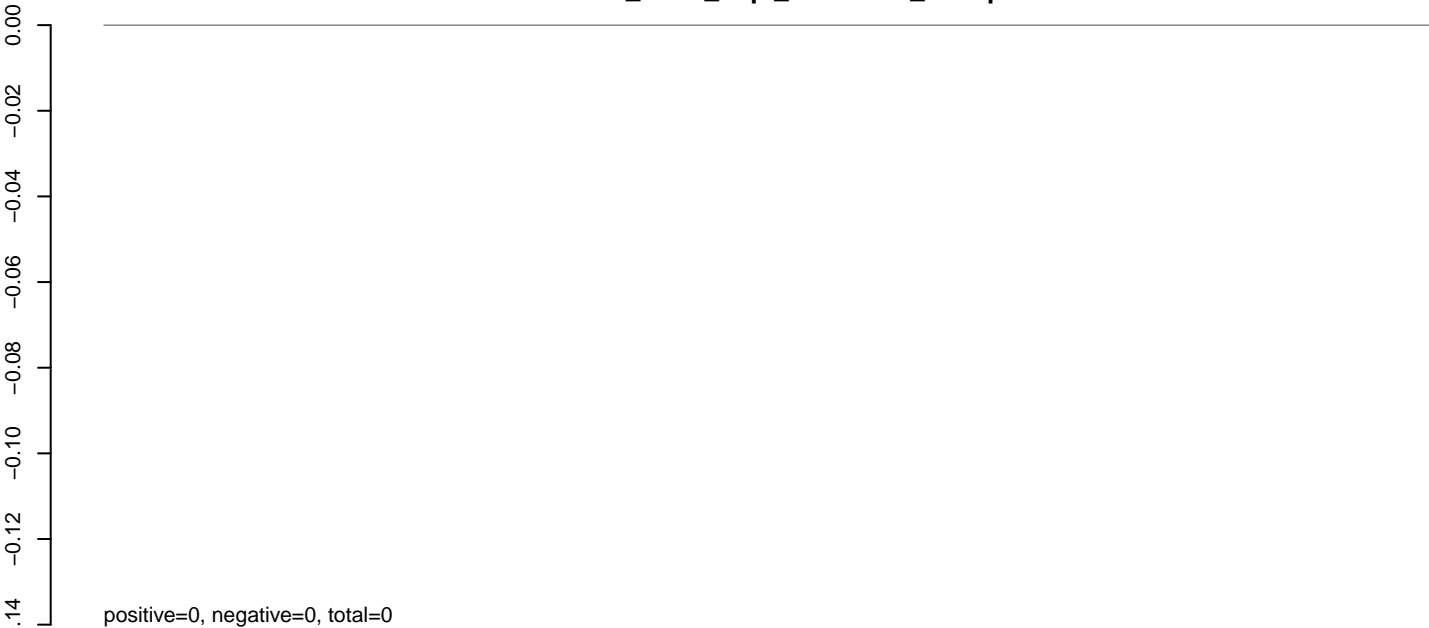
0 2000 4000 6000 8000 10000



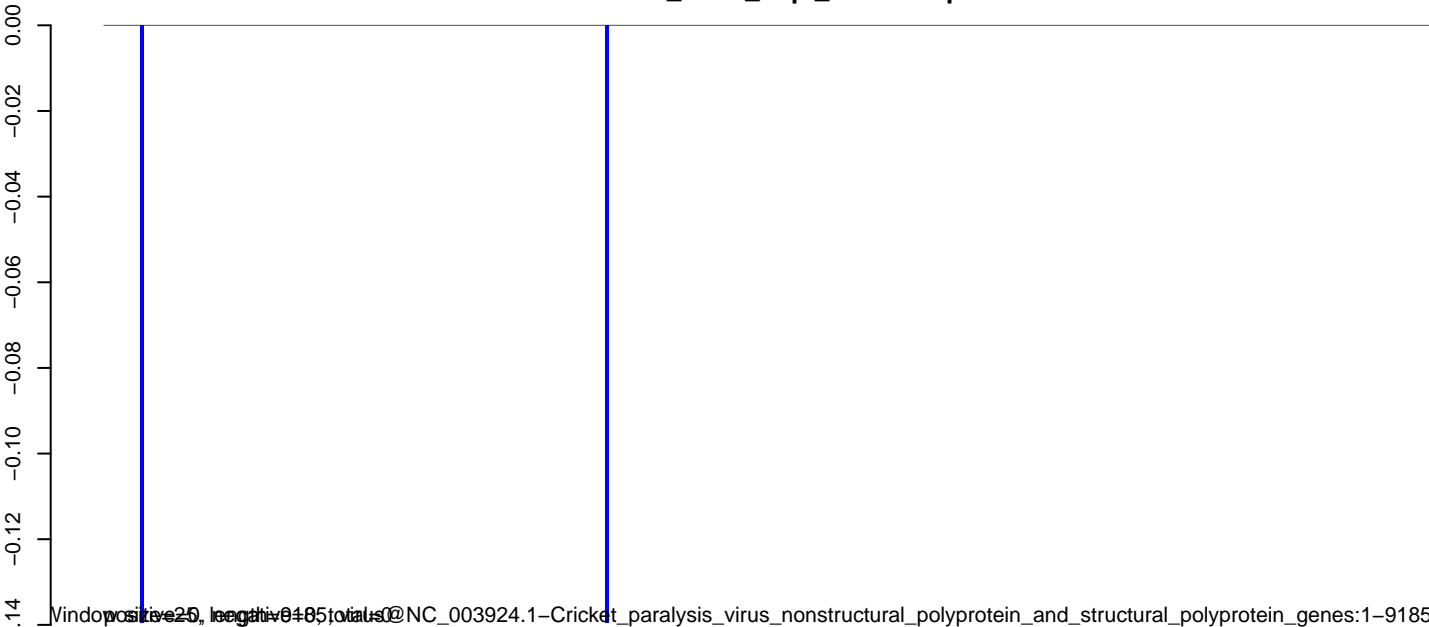
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



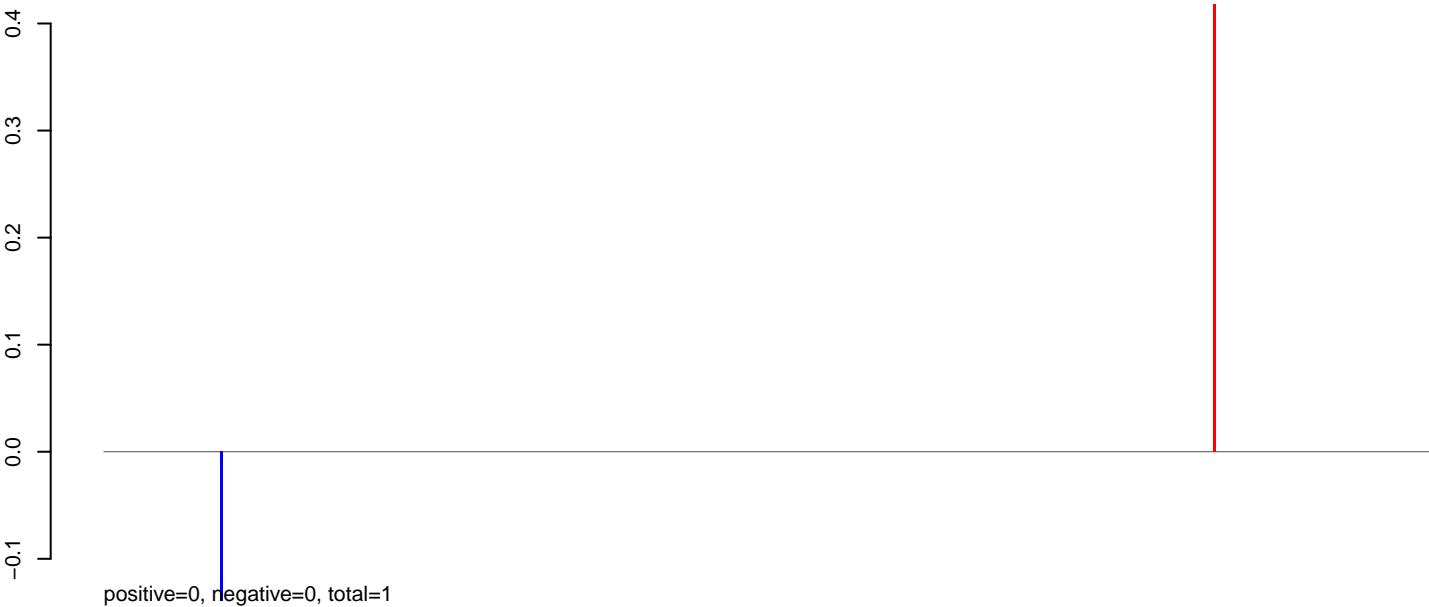
AeAlbo_C636_2dpi_mock3.rep



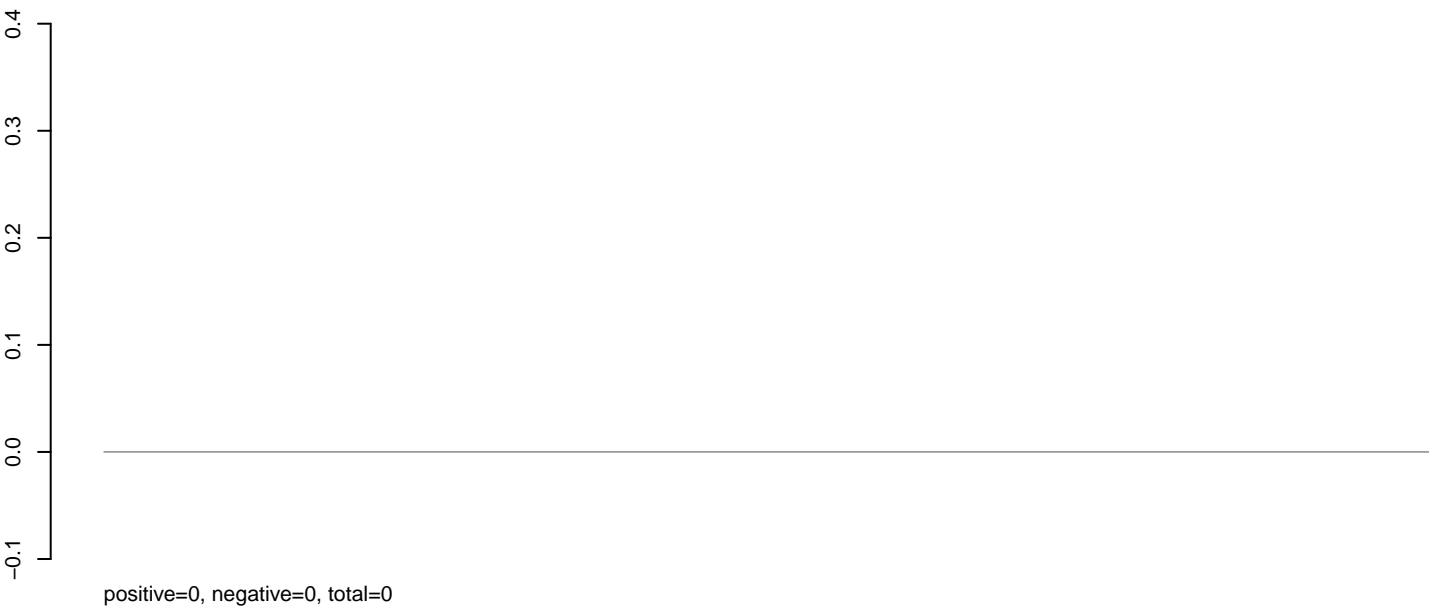
Vindor positive=25, negative=185, total=210 NC_003924.1-Cricket_paralysis_virus_nonstructural_polyprotein_and_structural_polyprotein_genes:1-9185

0 2000 4000 6000 8000

AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

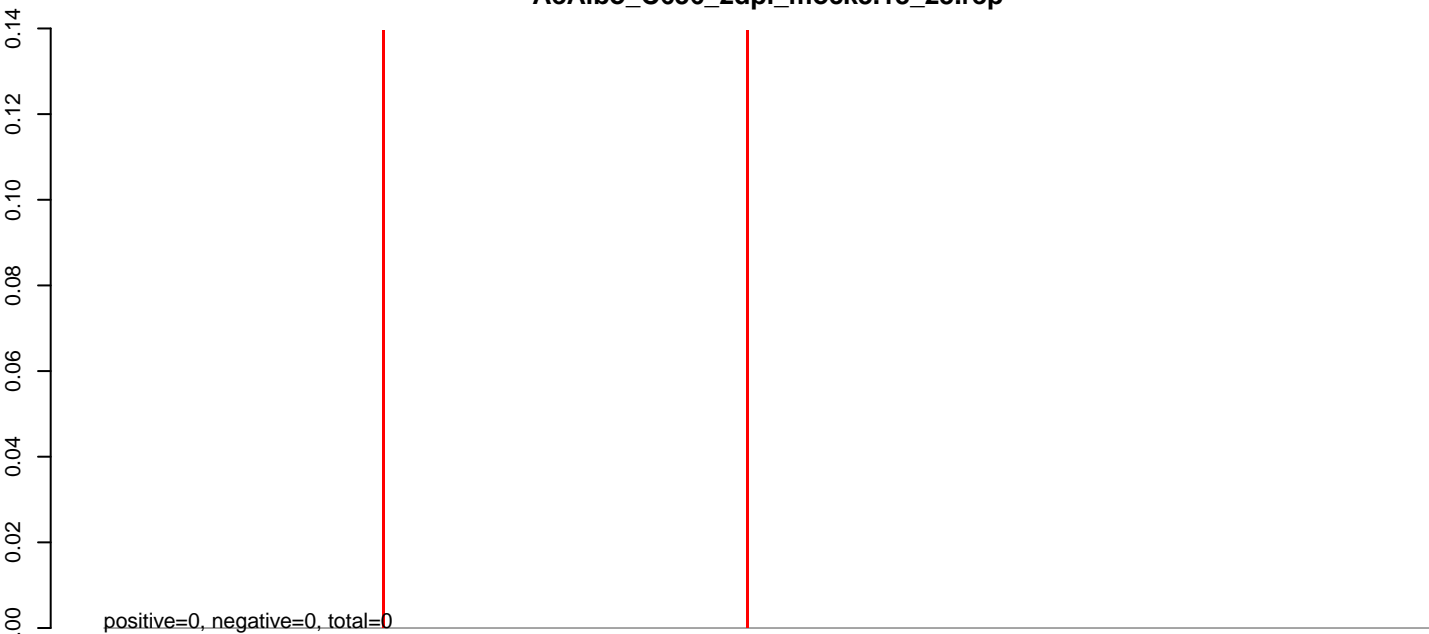


AeAlbo_C636_2dpi_mock3.rep

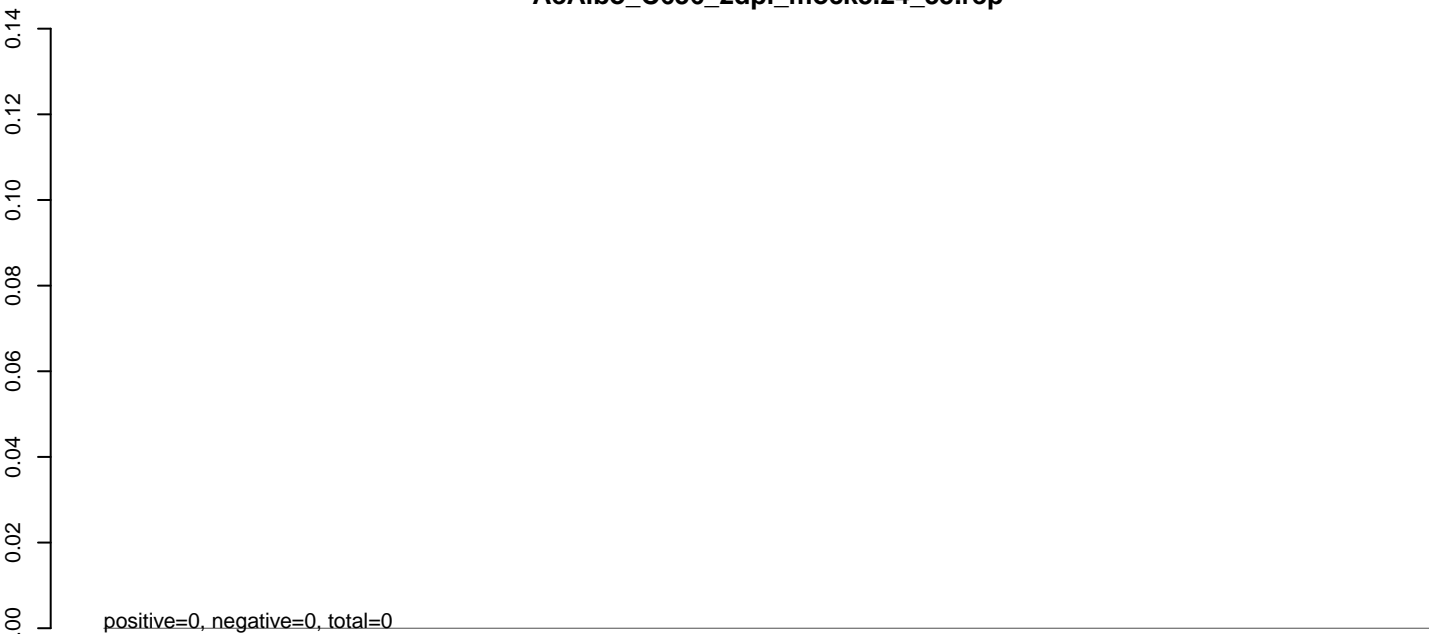


0 2000 4000 6000 8000 10000 12000

AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

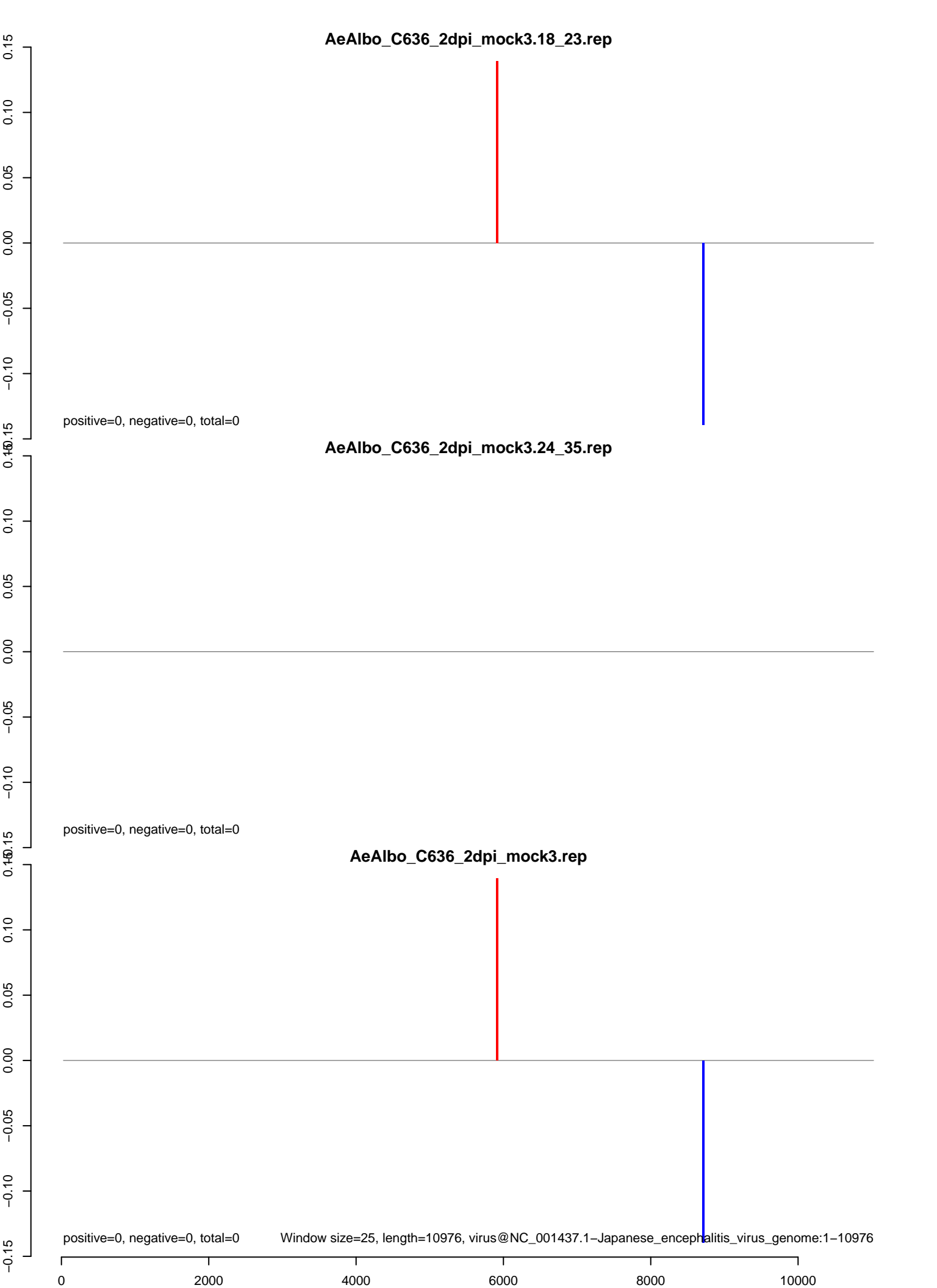


AeAlbo_C636_2dpi_mock3.rep

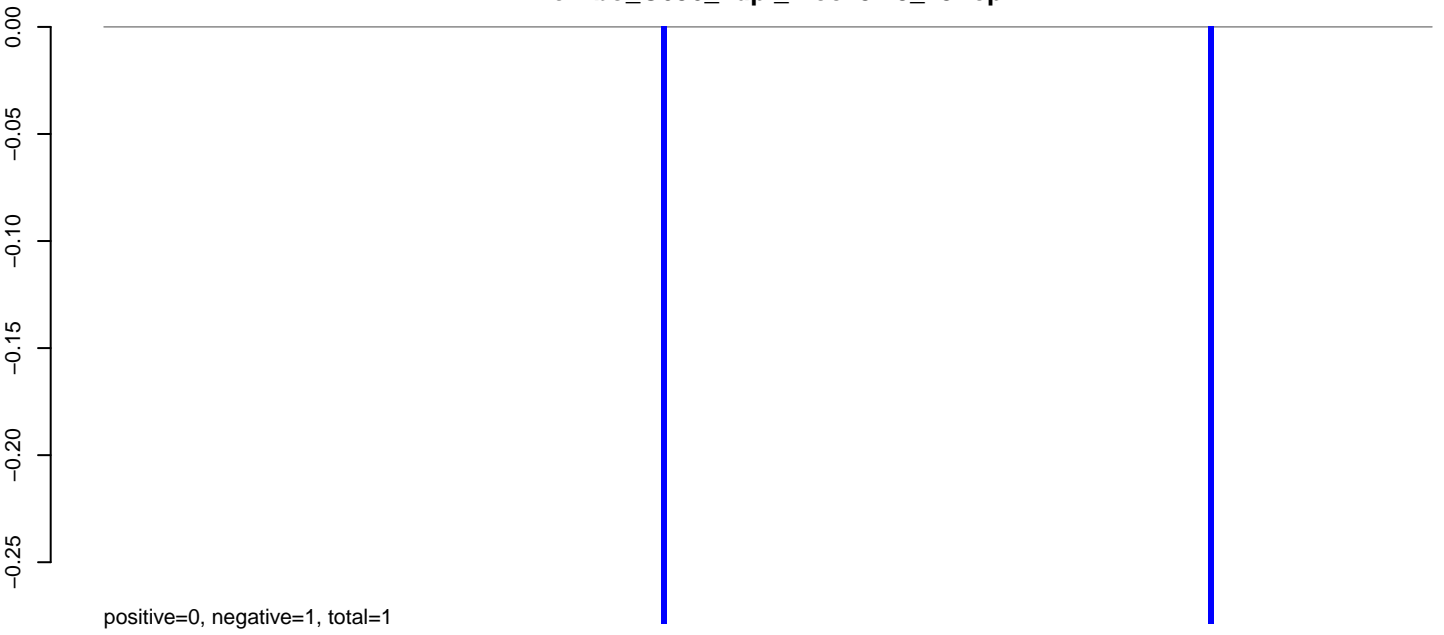


Window size=25, threshold=12294, virus@NC_001927.1-Bunyamwera_virus_segment_S_plus_segment_M_plus_segment_L:1-12294

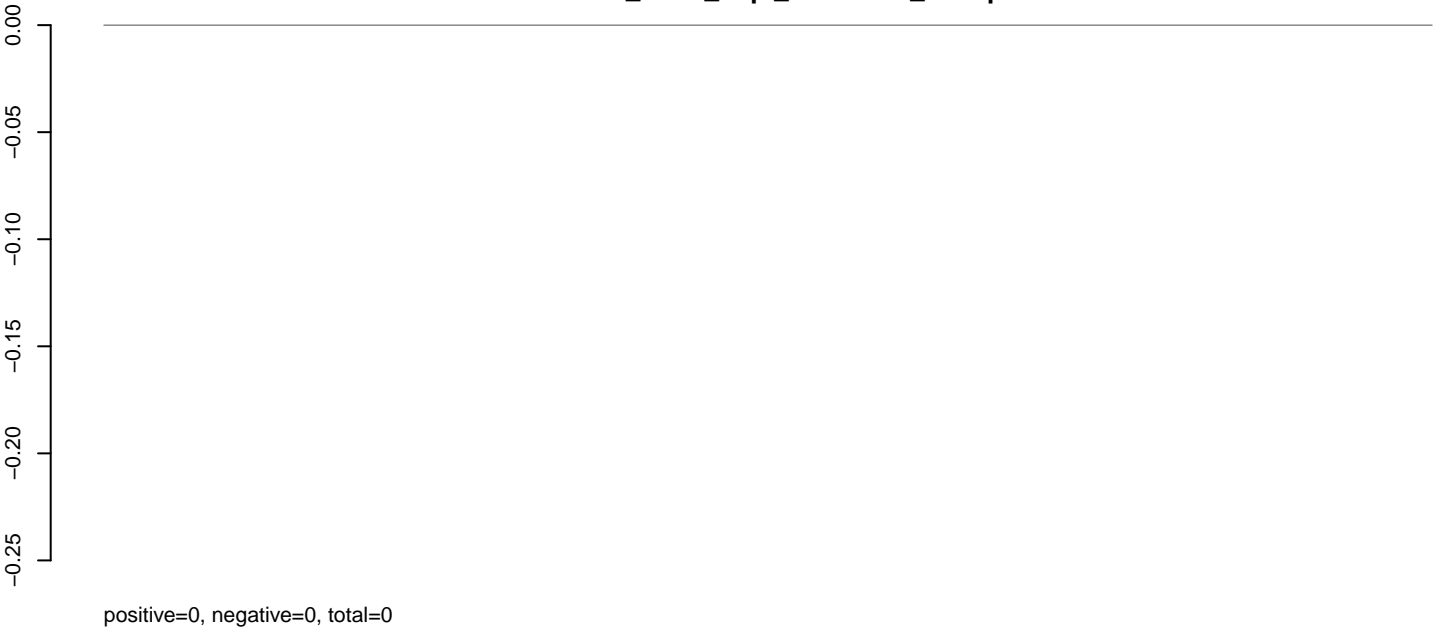
0 2000 4000 6000 8000 10000 12000



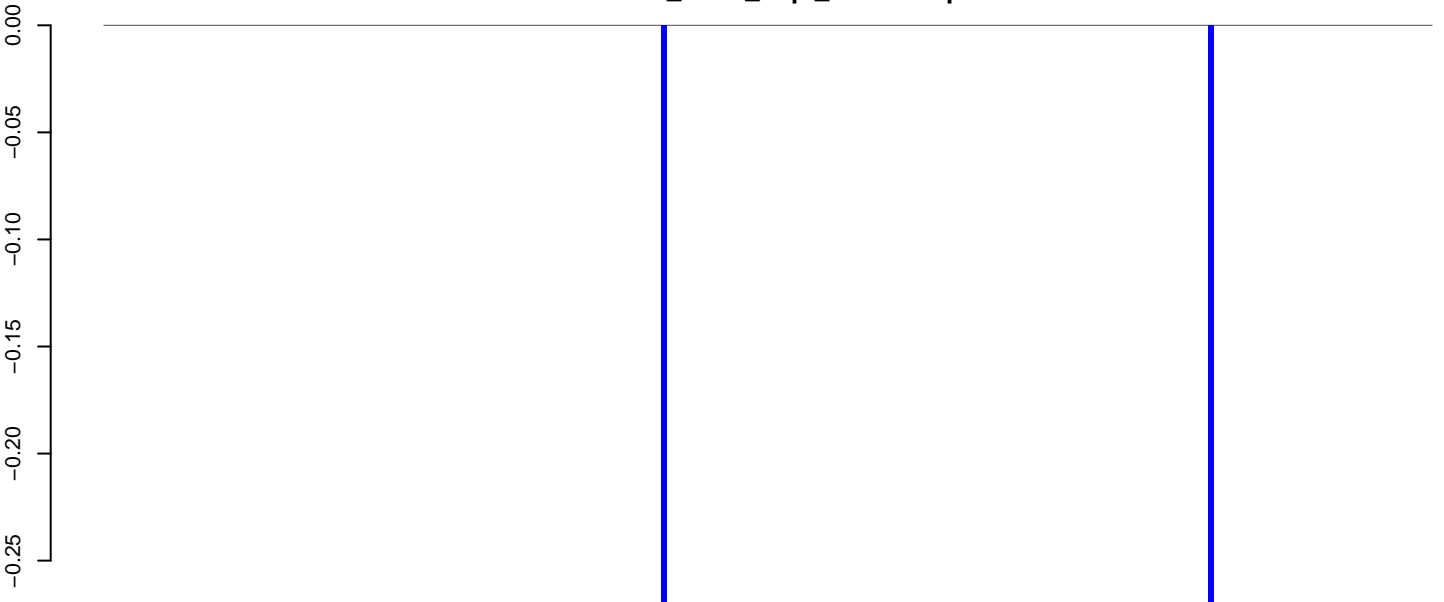
AeAlbo_C636_2dpi_mock3.18_23.rep



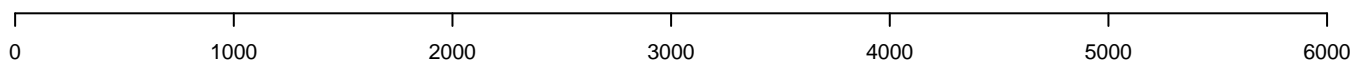
AeAlbo_C636_2dpi_mock3.24_35.rep



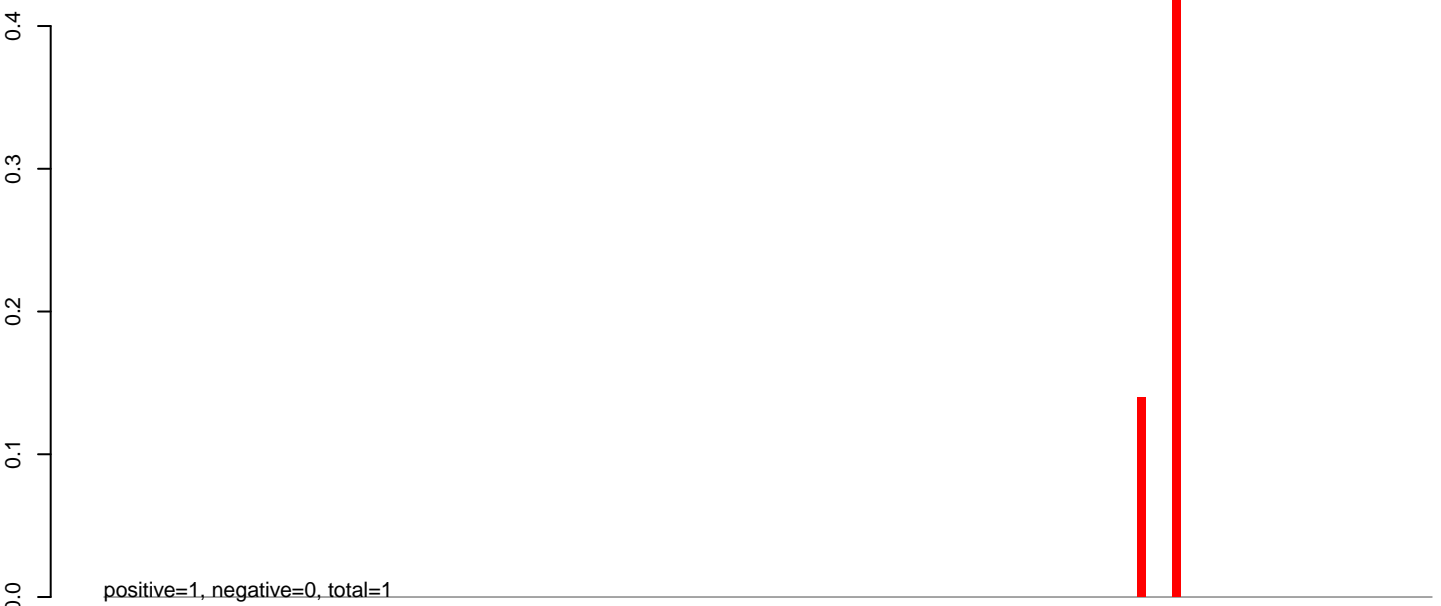
AeAlbo_C636_2dpi_mock3.rep



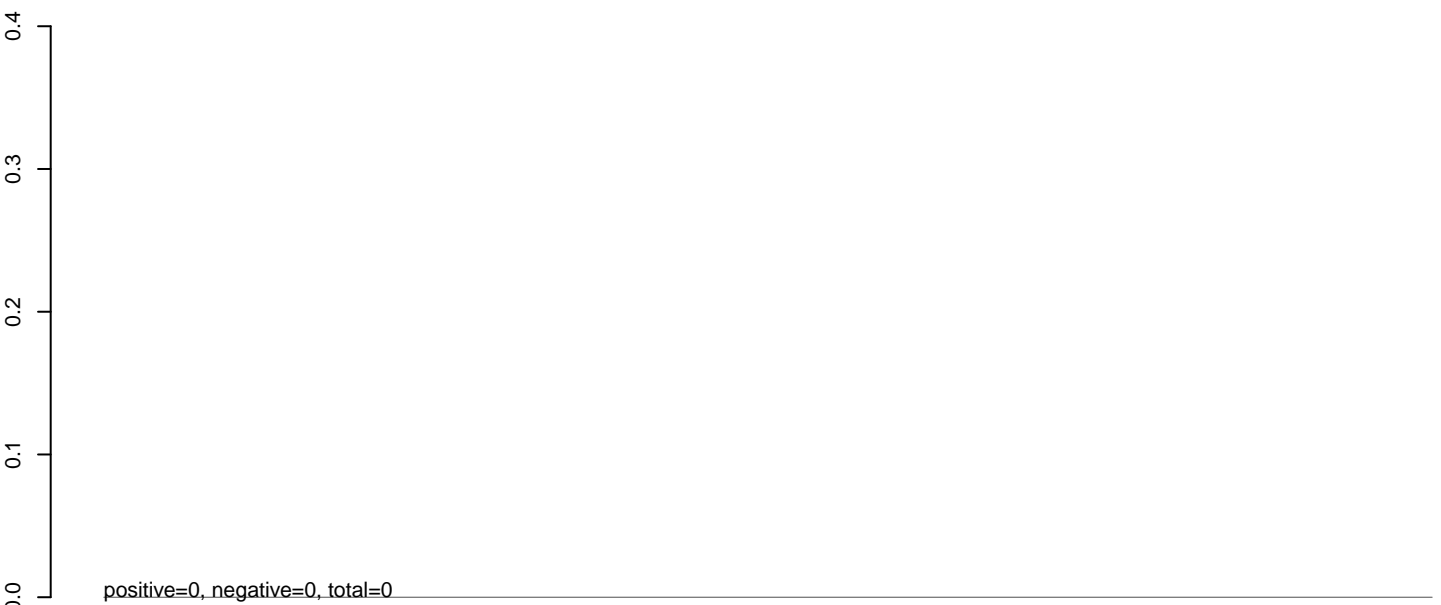
Window size=25, length=6065, virus@MH188054.1-Culex_Y_virus_strain_CYV_Kern_partial_genome:1-6065



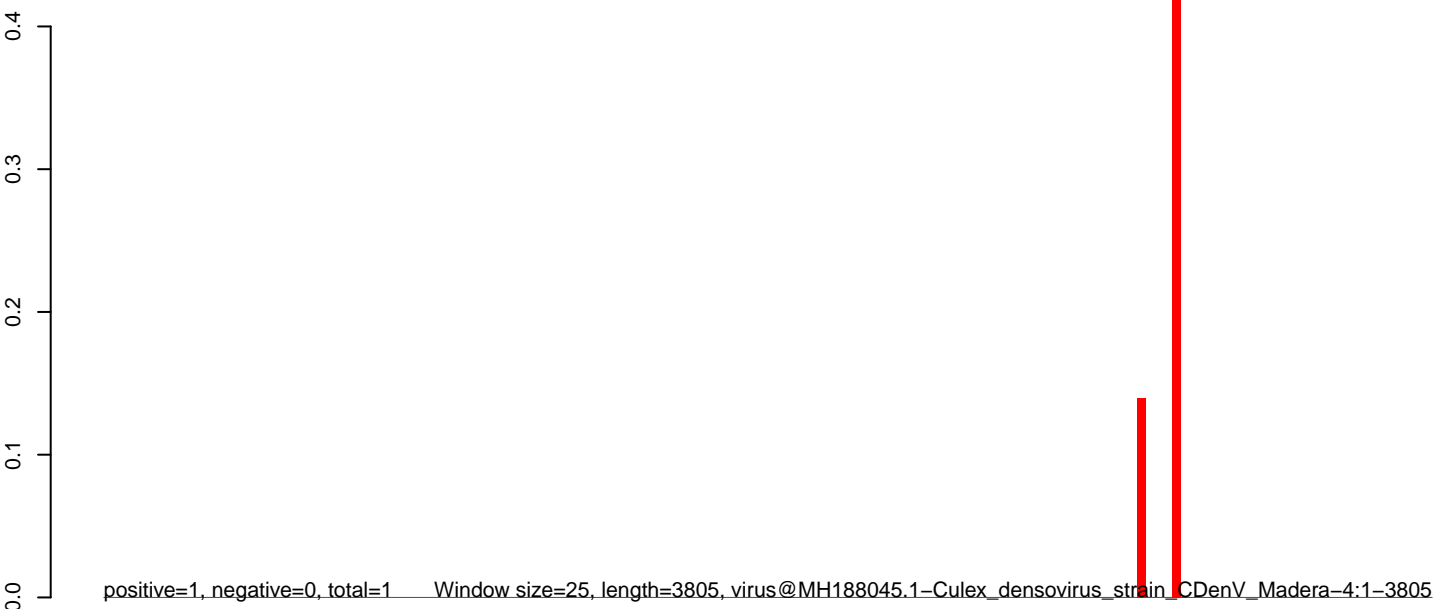
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



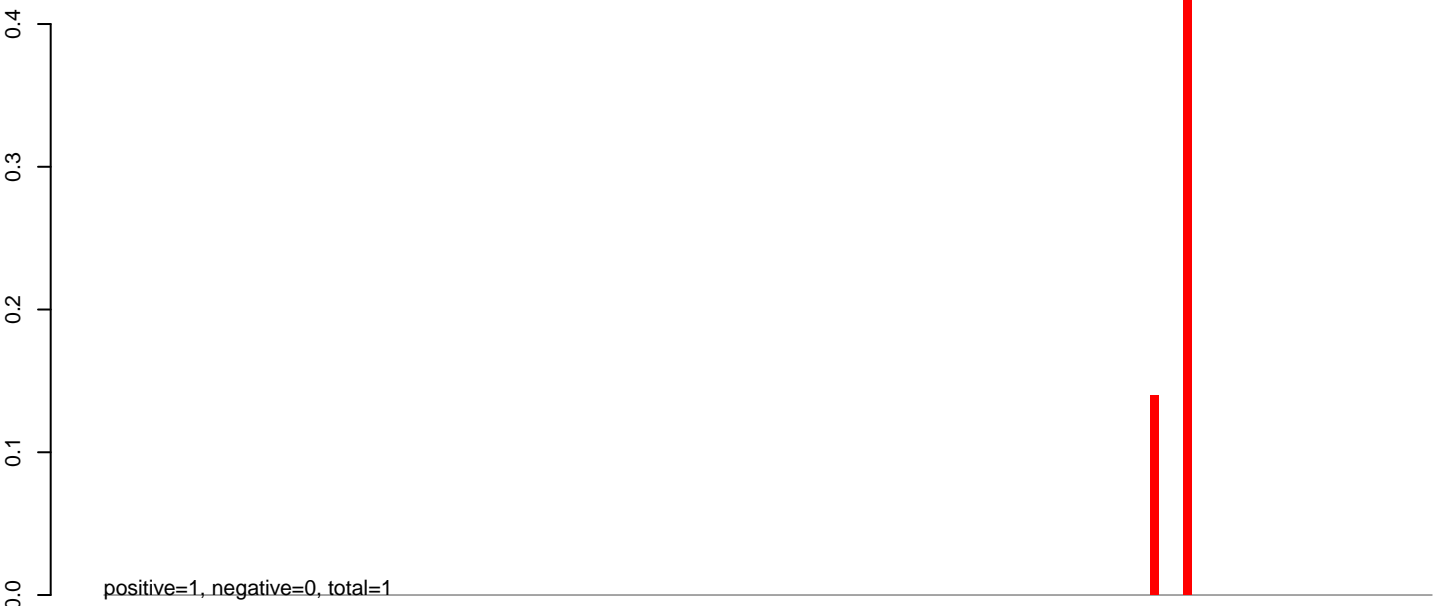
AeAlbo_C636_2dpi_mock3.rep



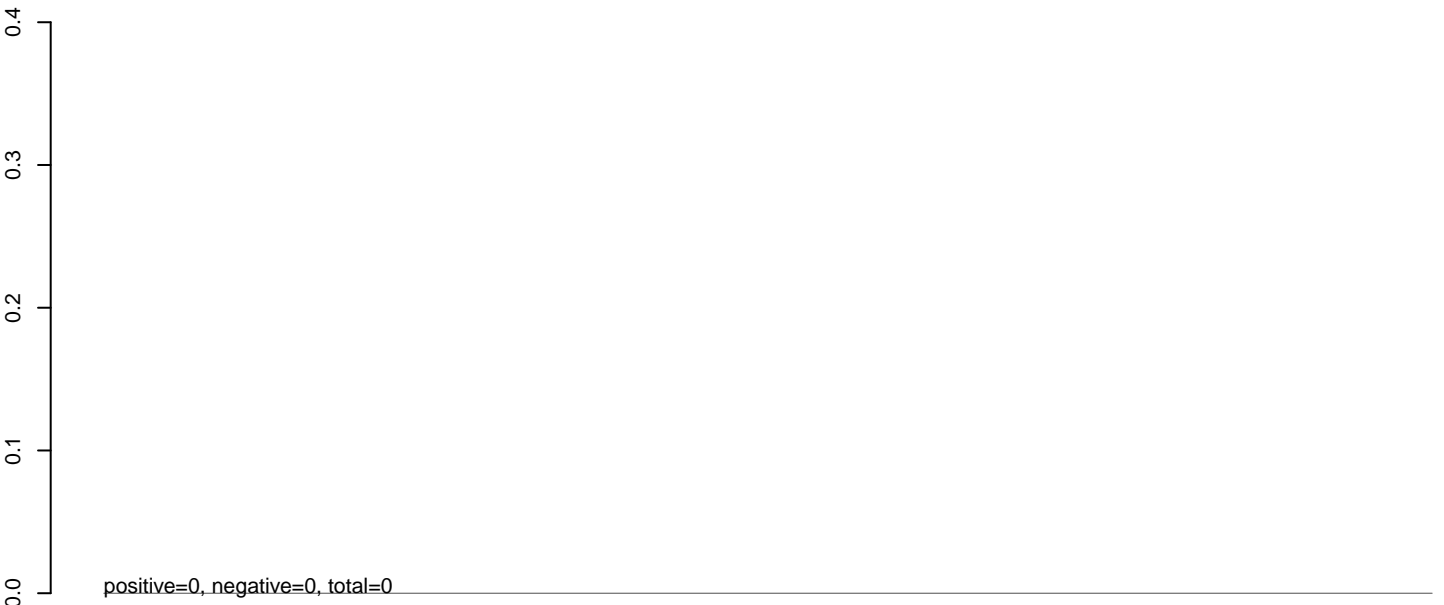
Window size=25, length=3805, virus@MH188045.1-Culex_densovirus_strain_CDENV_Madera-4:1-3805

0 1000 2000 3000 4000

AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



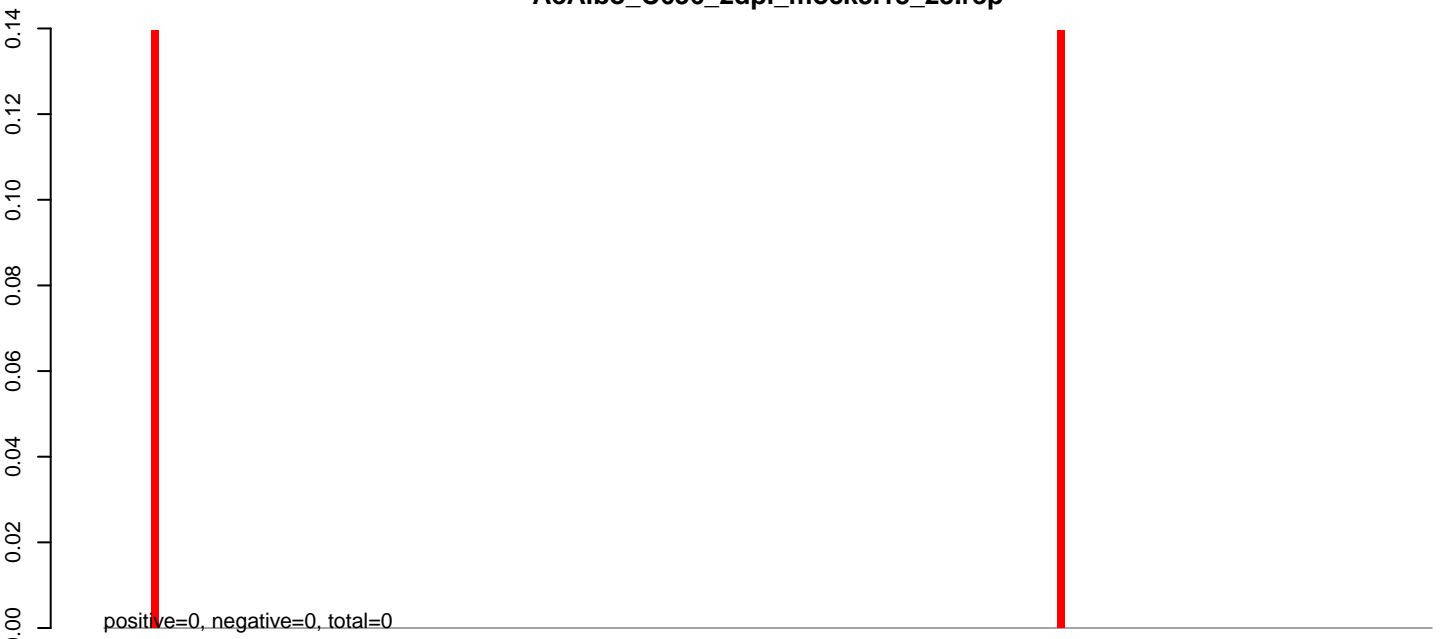
AeAlbo_C636_2dpi_mock3.rep



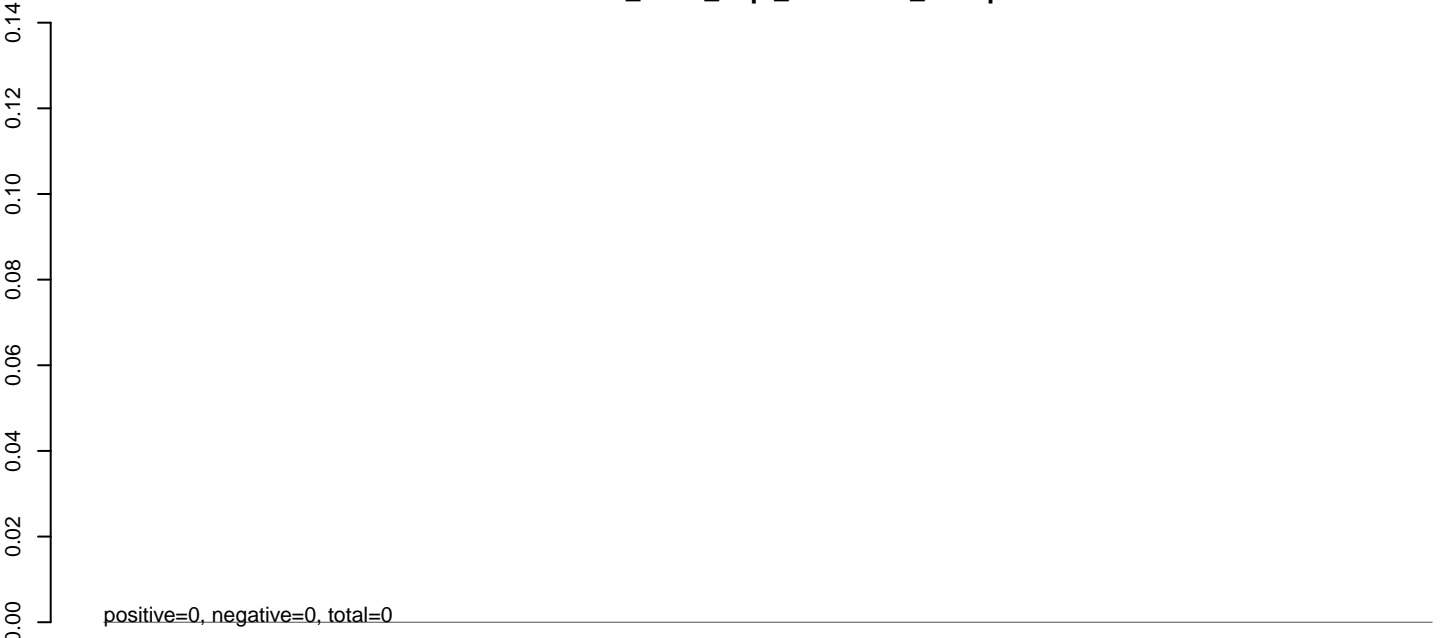
Window size=25, length=3991, virus@MH188044.1-Culex_densovirus_strain_CDenV_Madera-3:1-3991

0 1000 2000 3000 4000

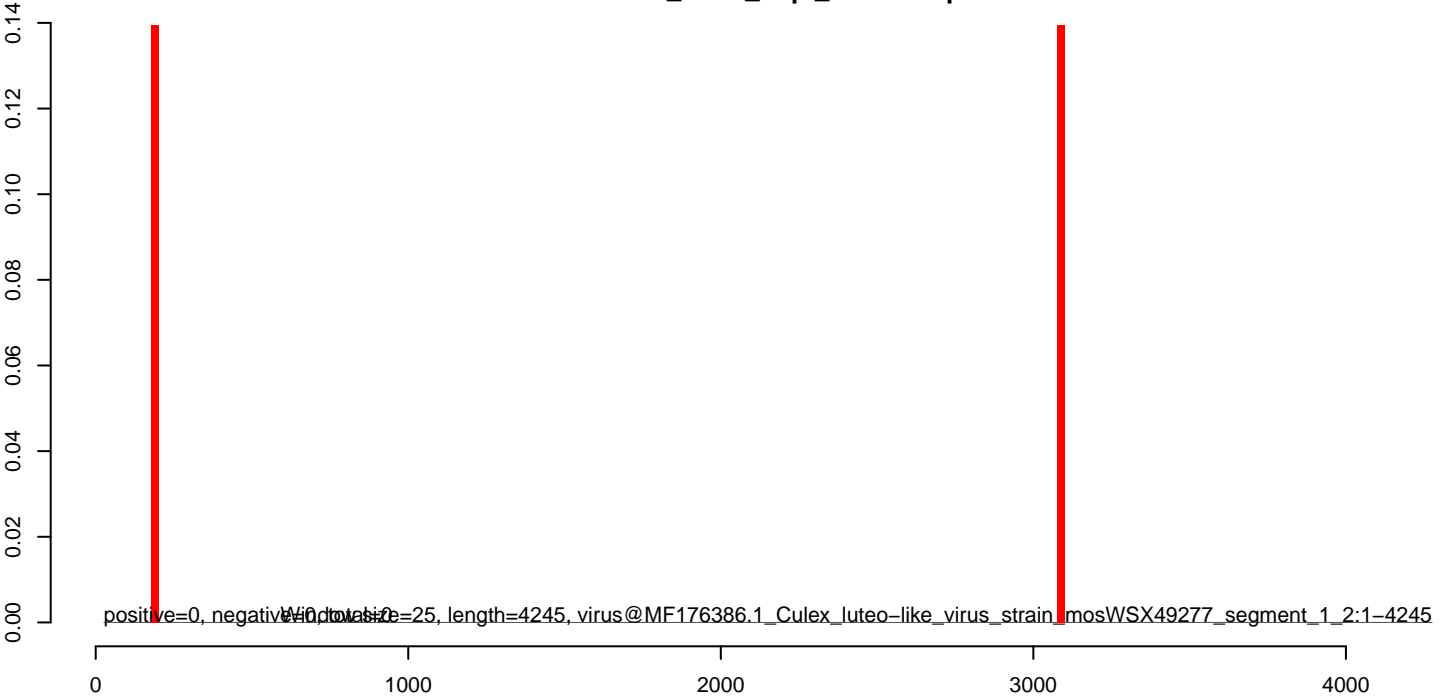
AeAlbo_C636_2dpi_mock3.18_23.rep



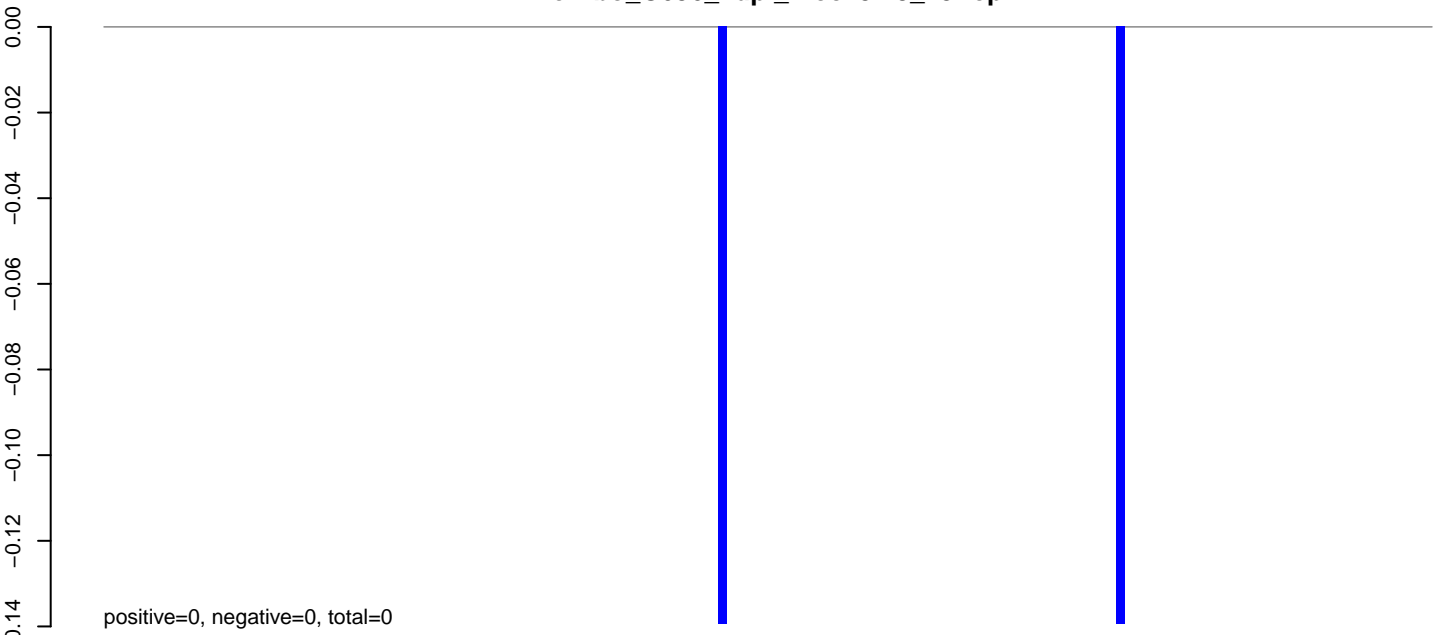
AeAlbo_C636_2dpi_mock3.24_35.rep



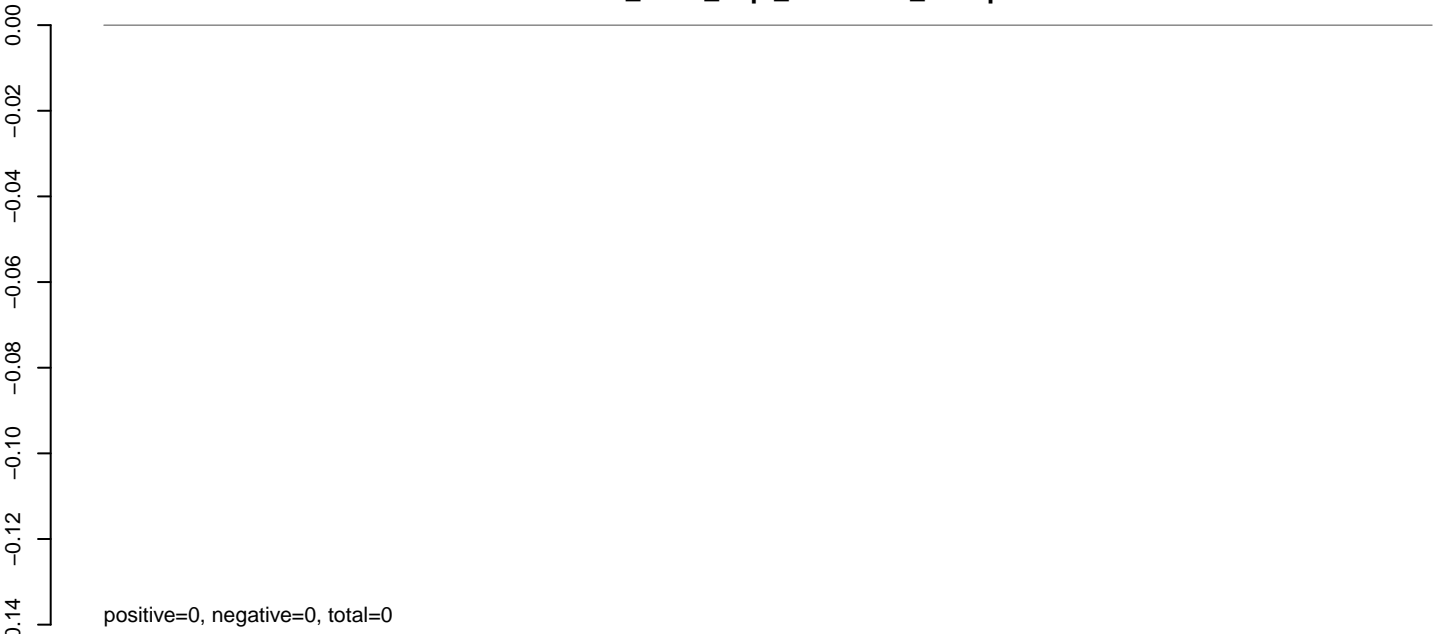
AeAlbo_C636_2dpi_mock3.rep



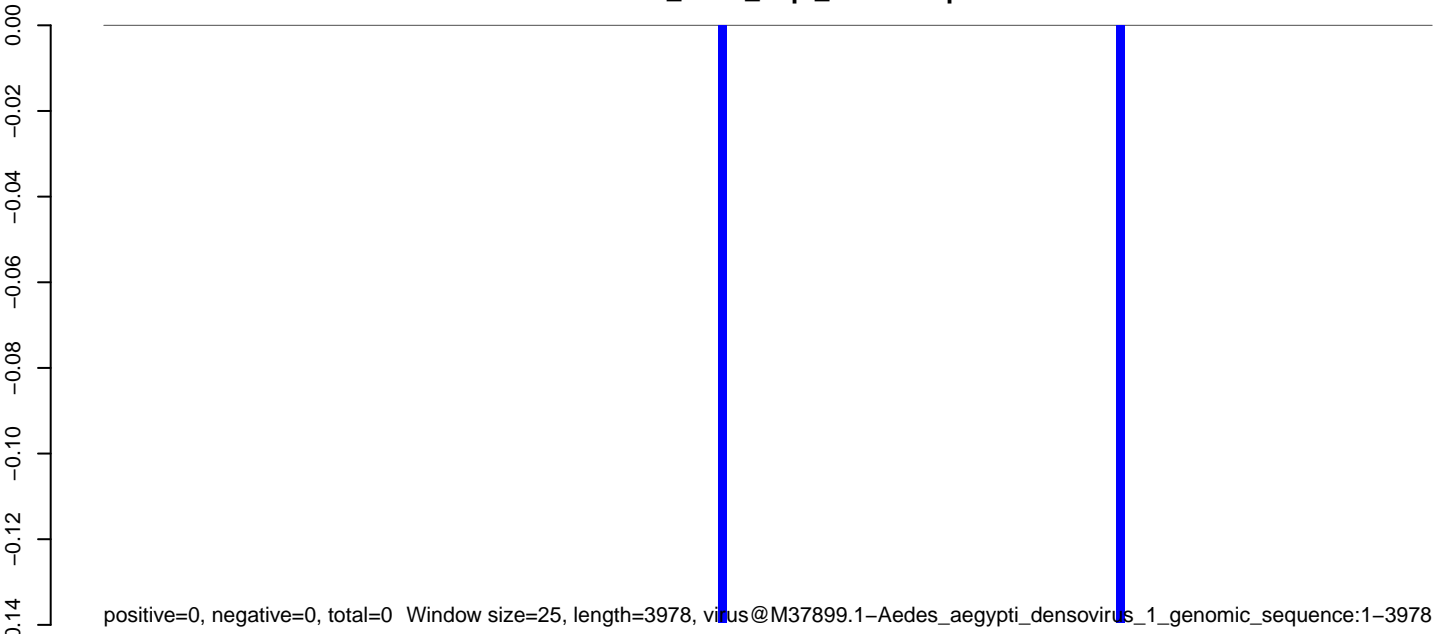
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

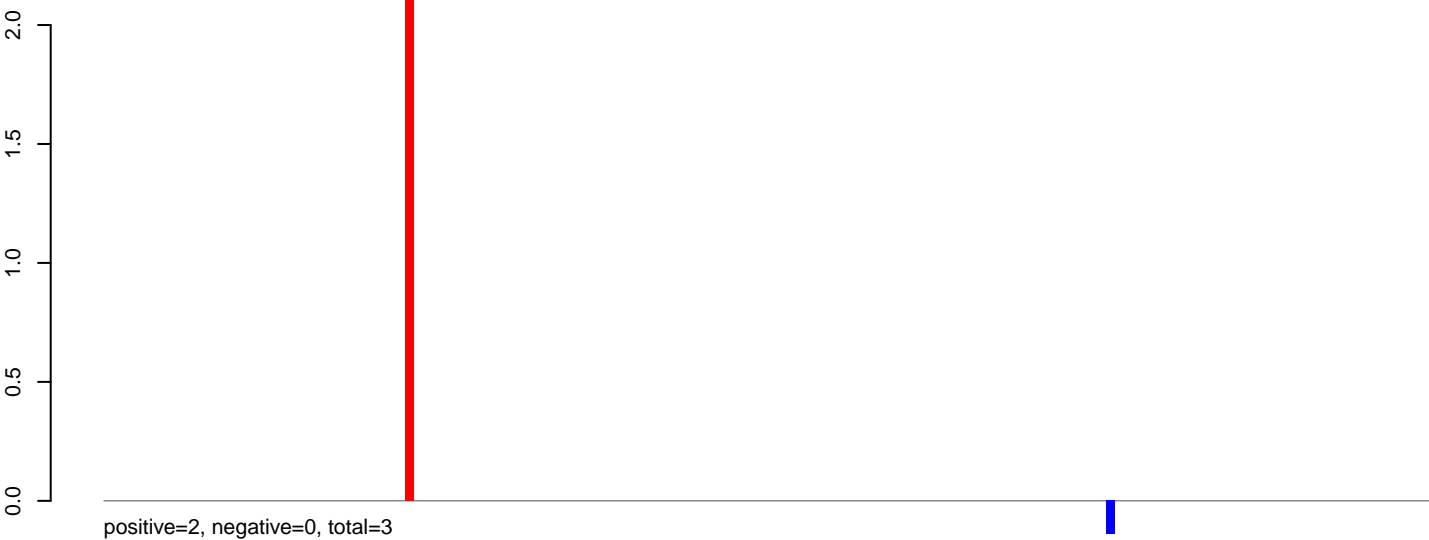


AeAlbo_C636_2dpi_mock3.rep

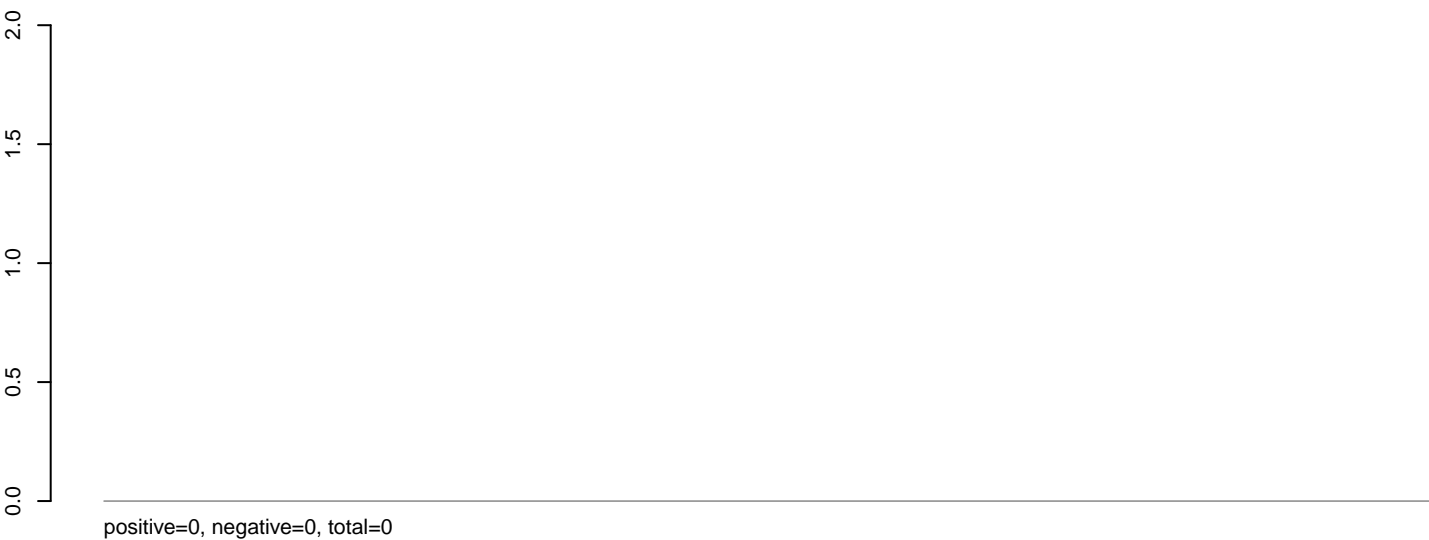


0 1000 2000 3000 4000

AeAlbo_C636_2dpi_mock3.18_23.rep

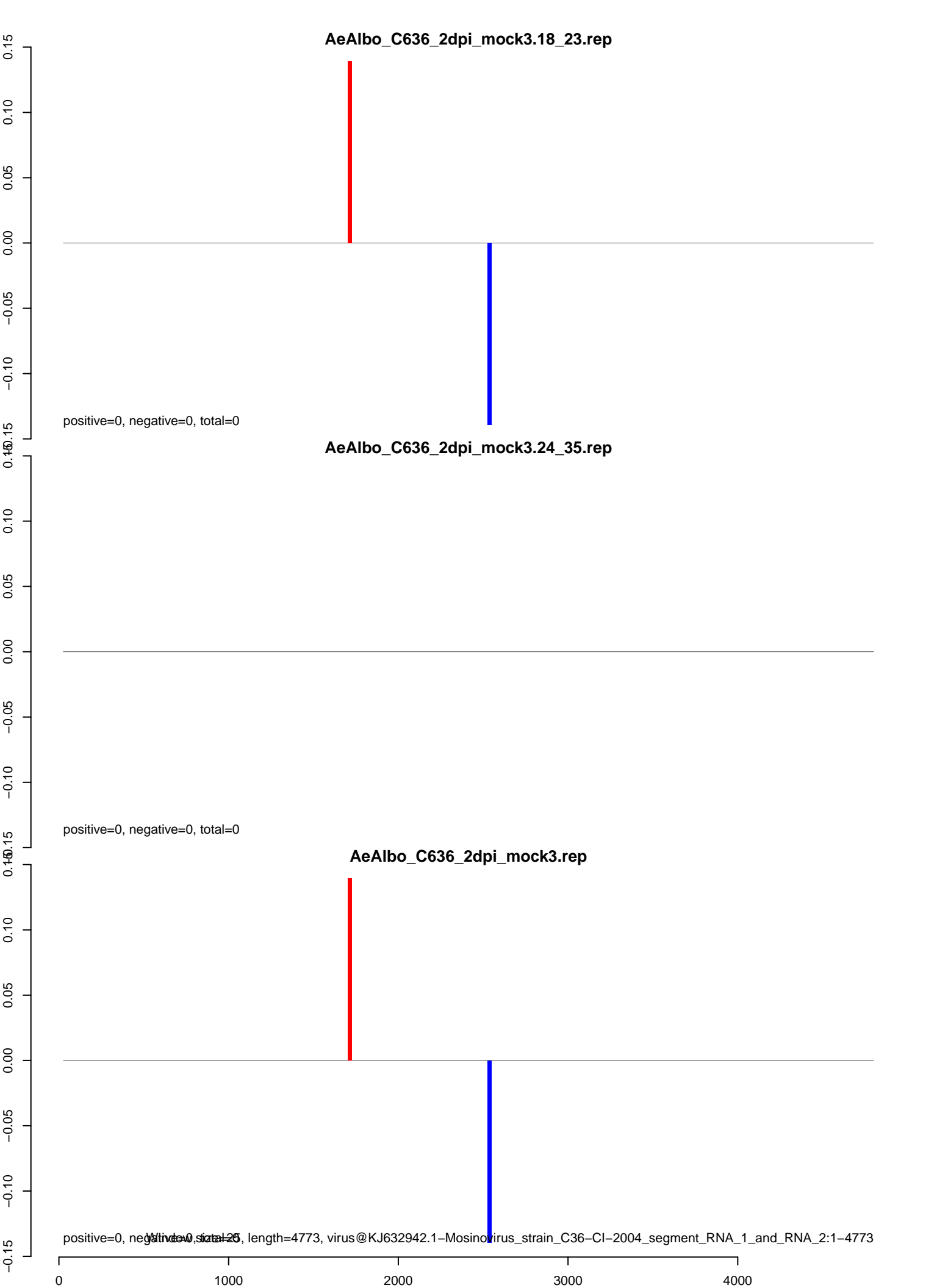


AeAlbo_C636_2dpi_mock3.24_35.rep

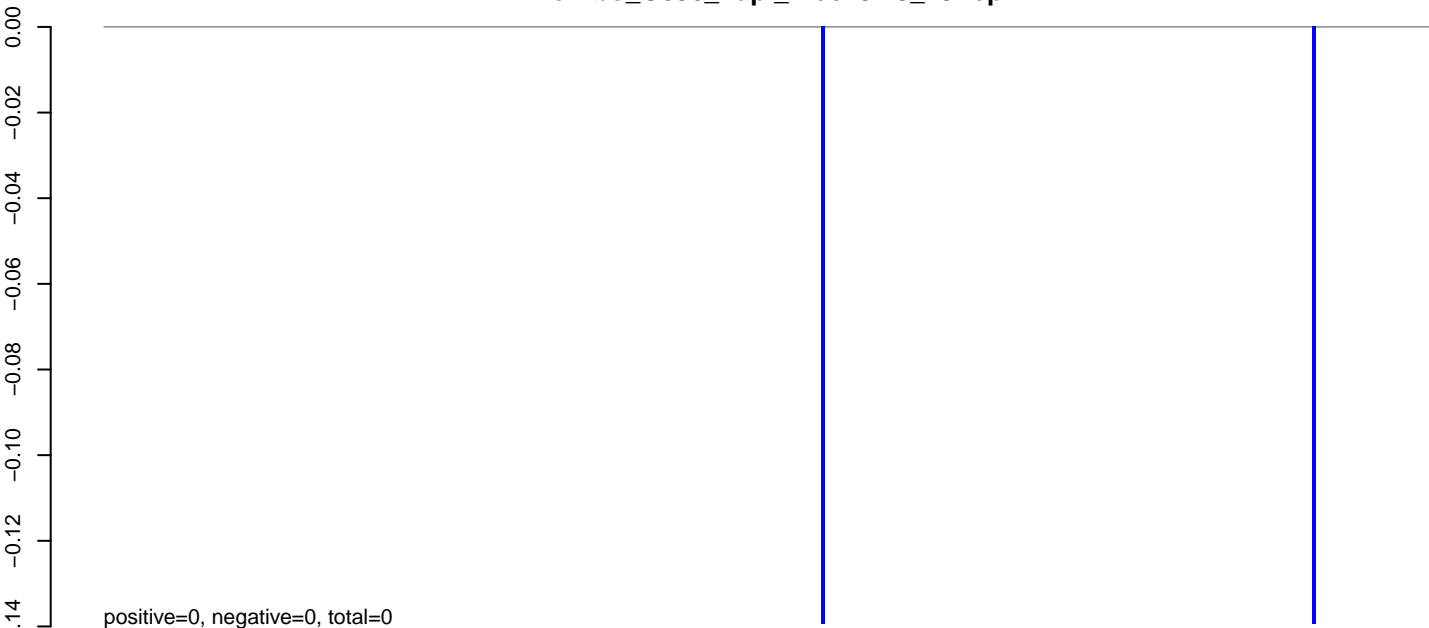


AeAlbo_C636_2dpi_mock3.rep

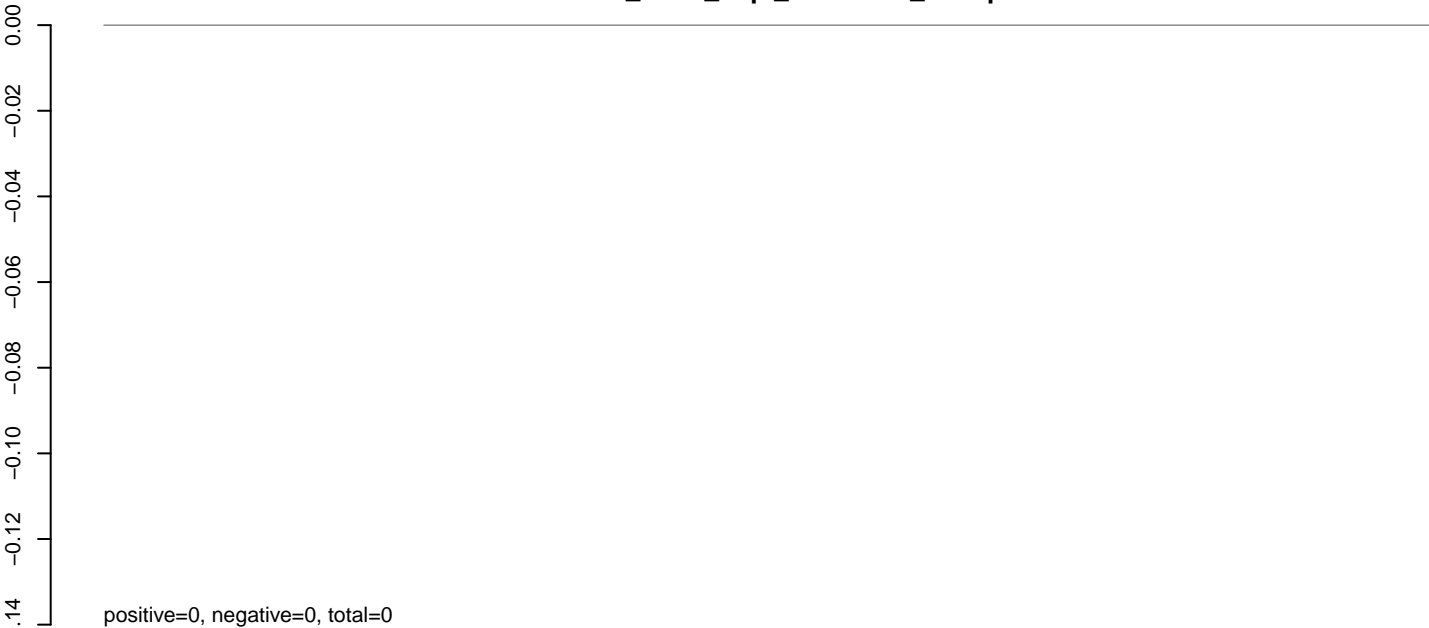




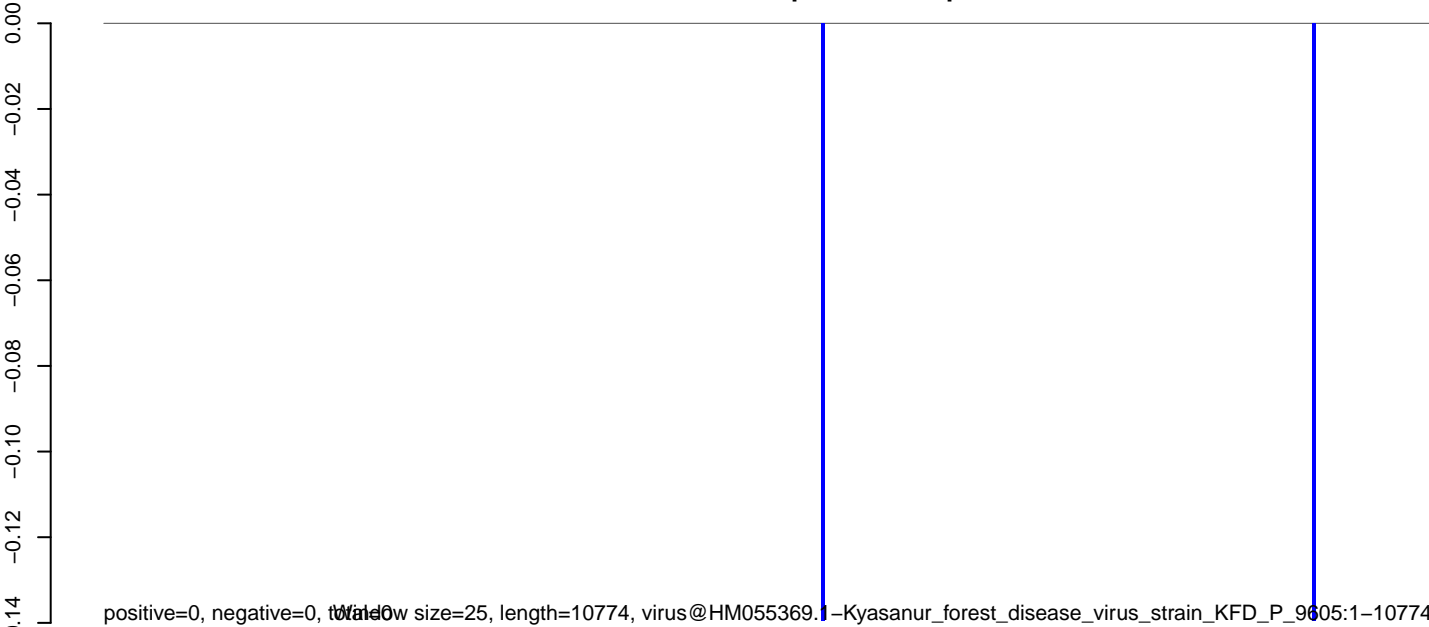
AeAlbo_C636_2dpi_mock3.18_23.rep



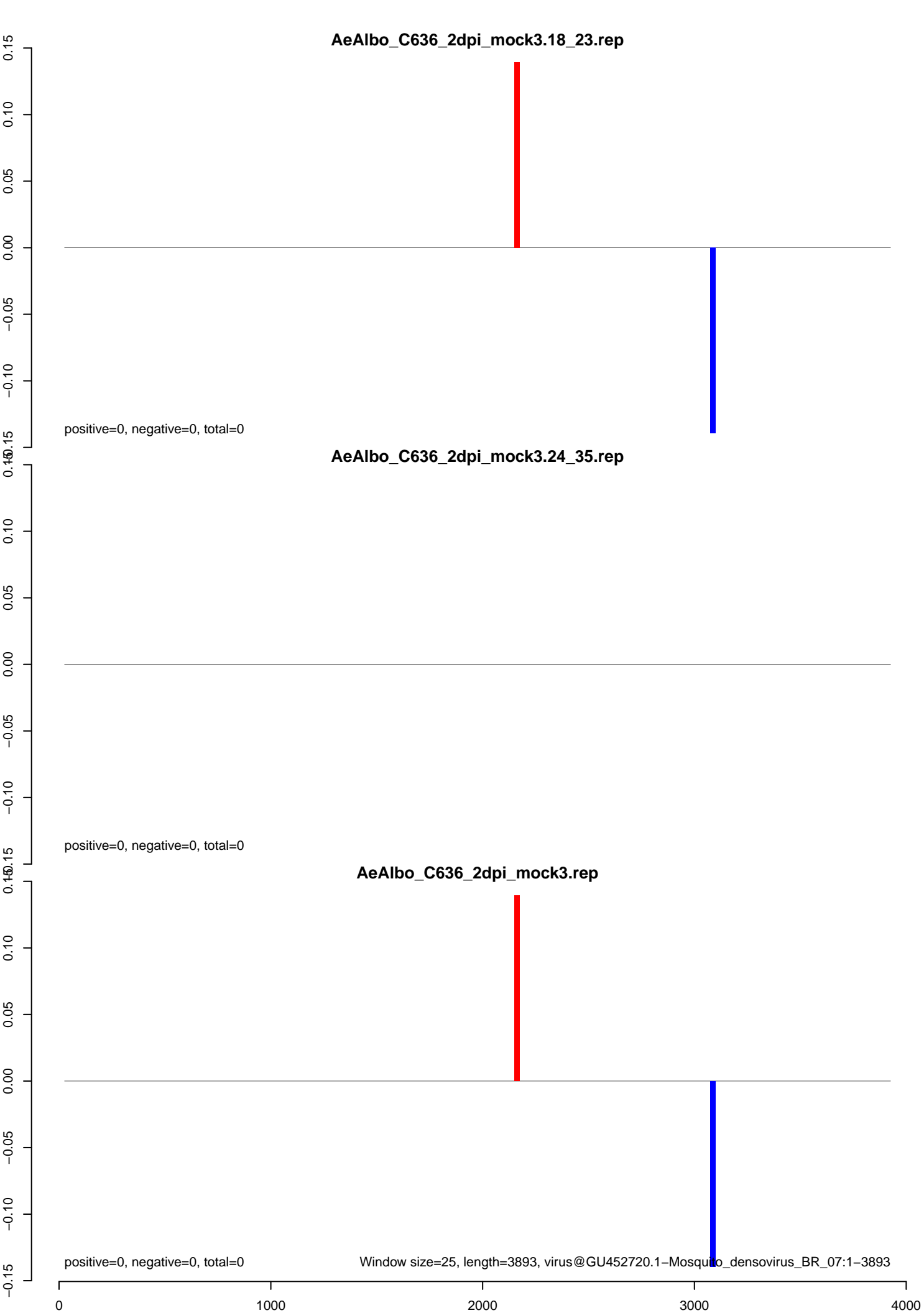
AeAlbo_C636_2dpi_mock3.24_35.rep



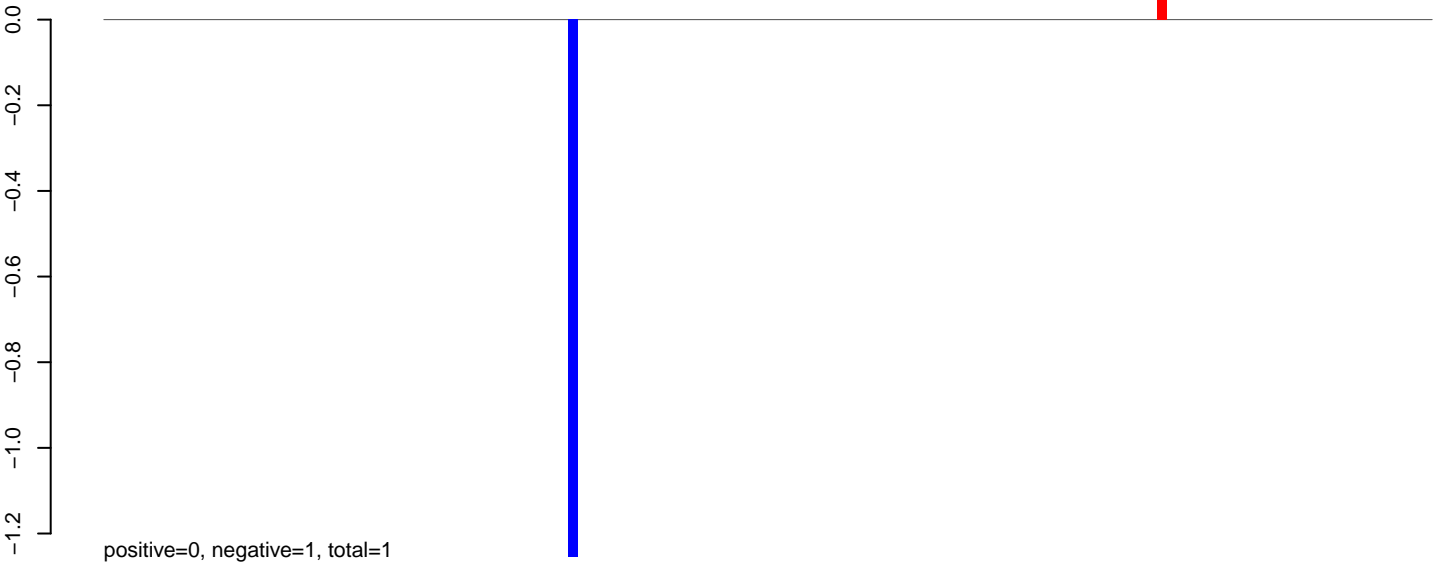
AeAlbo_C636_2dpi_mock3.rep



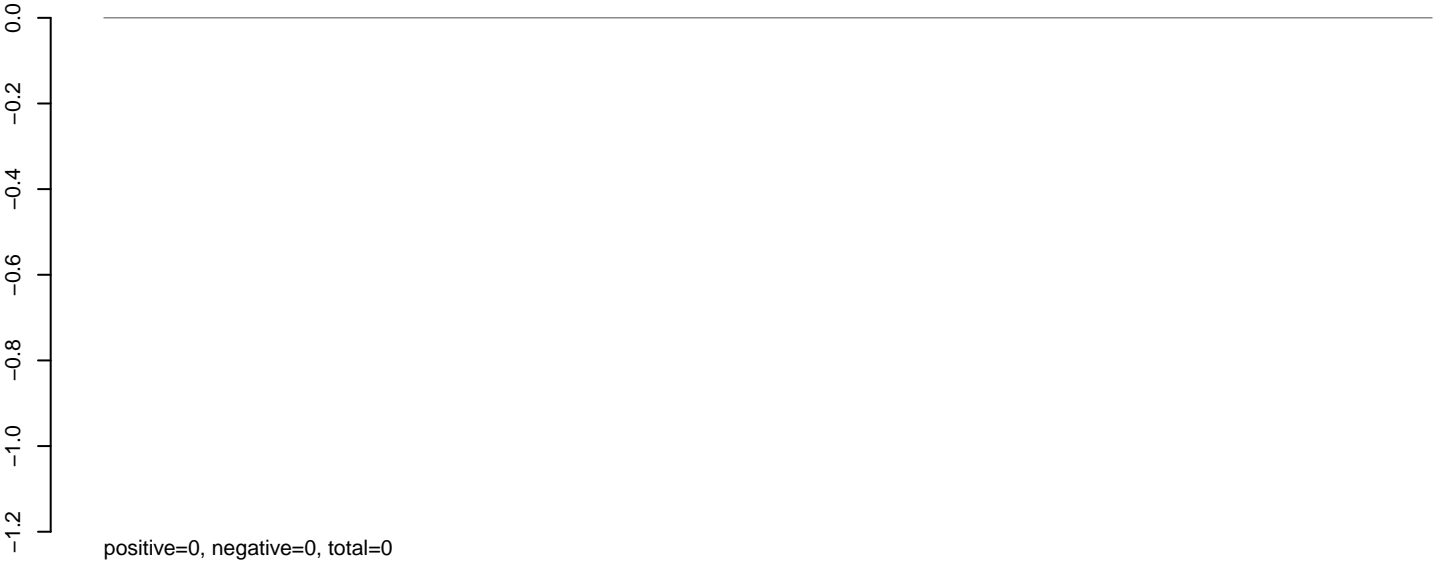
0 2000 4000 6000 8000 10000



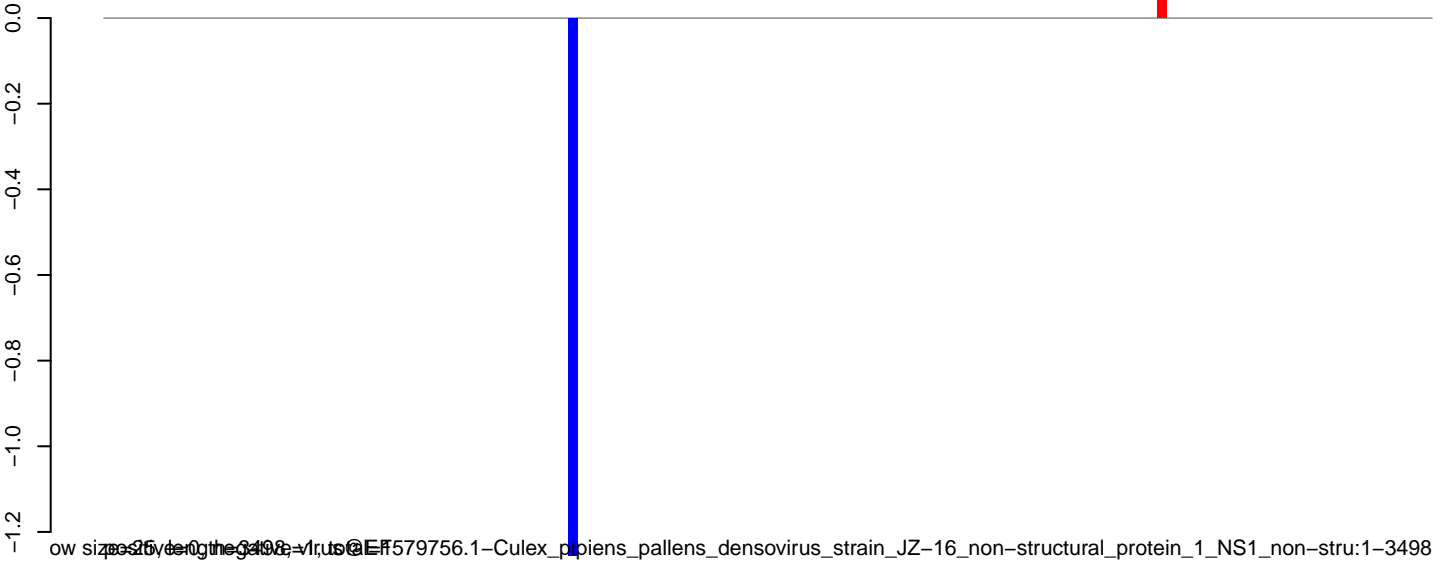
AeAlbo_C636_2dpi_mock3.18_23.rep



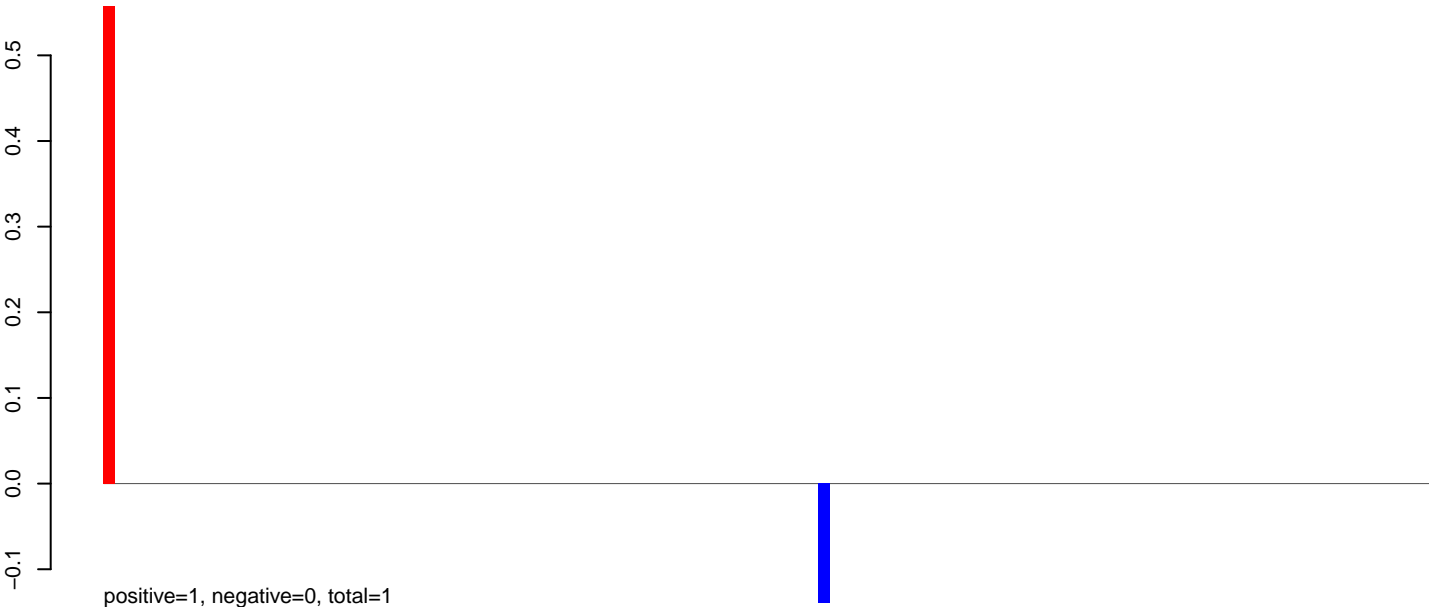
AeAlbo_C636_2dpi_mock3.24_35.rep



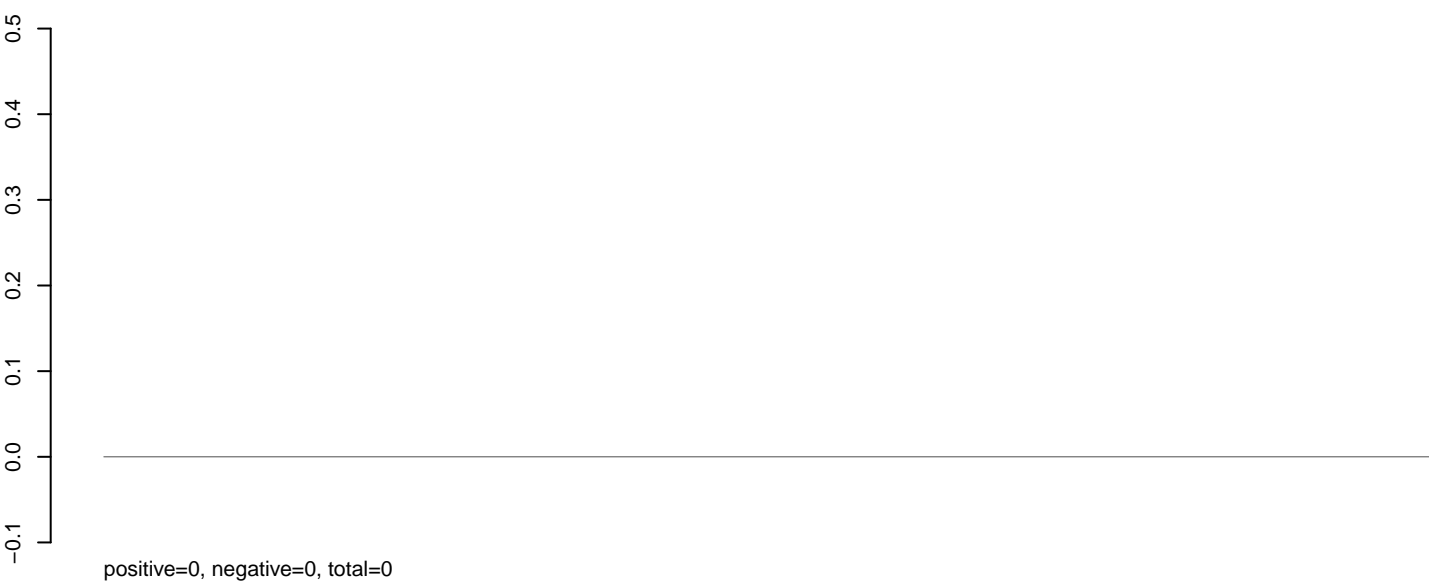
AeAlbo_C636_2dpi_mock3.rep



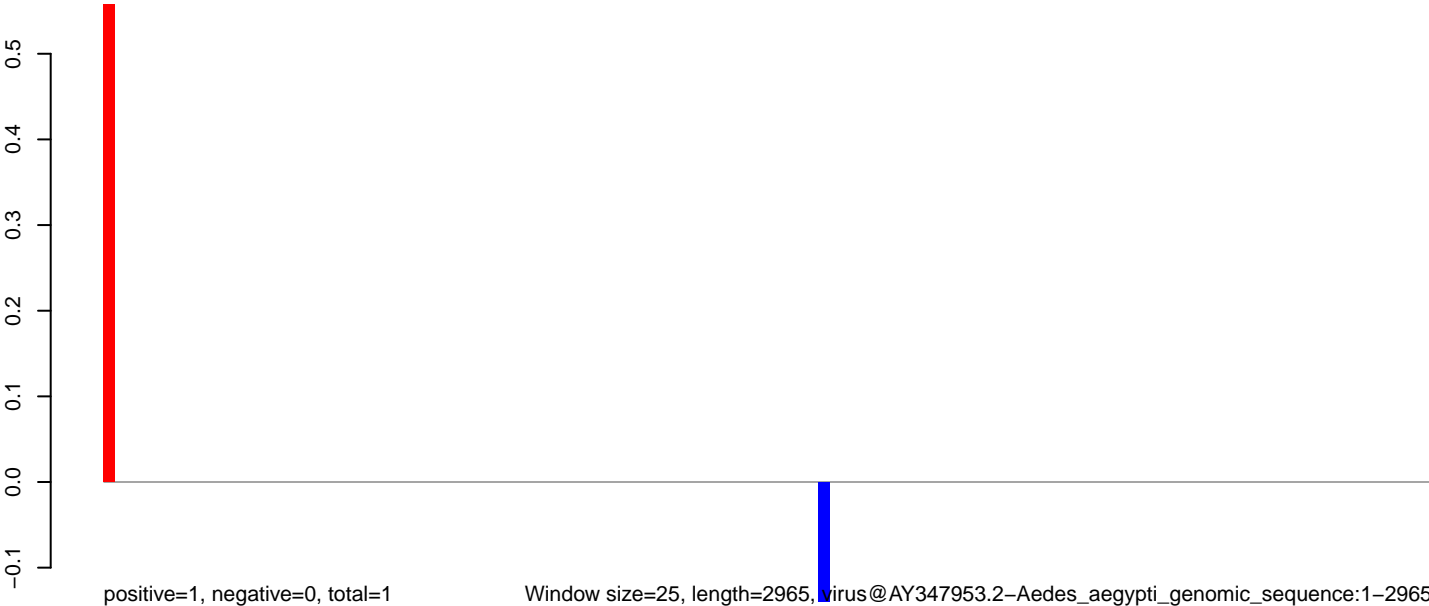
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

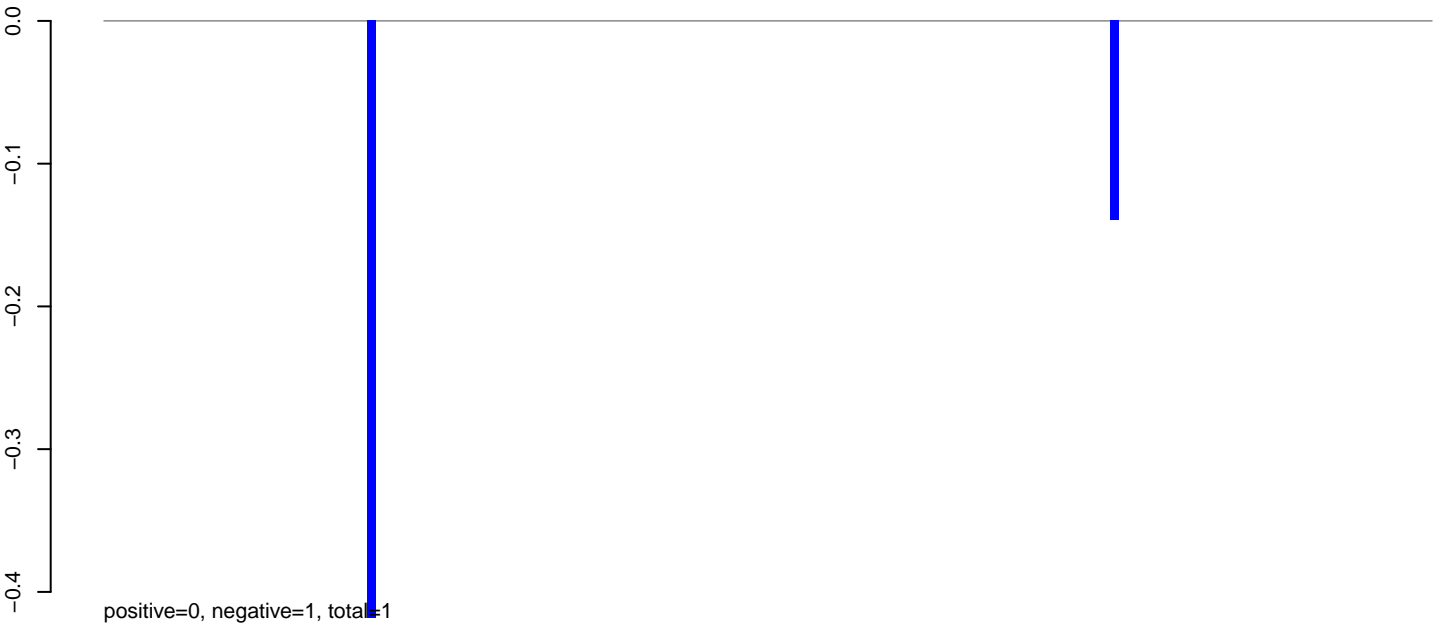


AeAlbo_C636_2dpi_mock3.rep

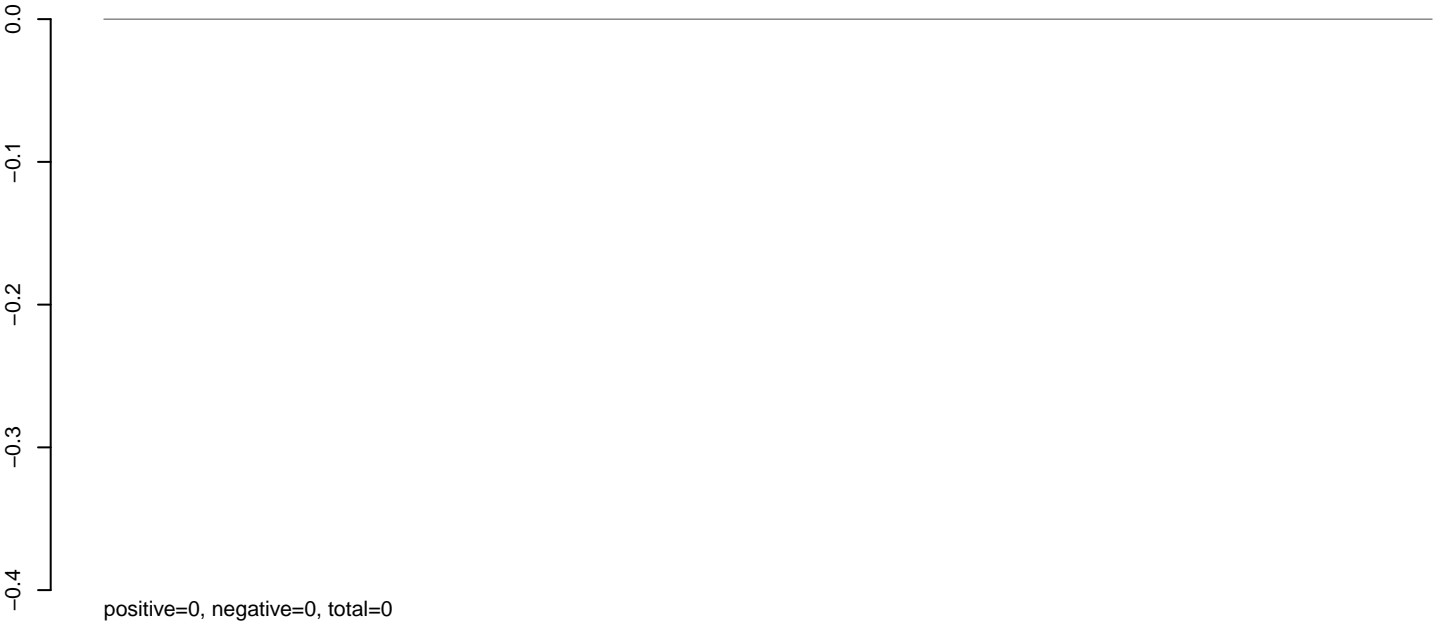


0 500 1000 1500 2000 2500 3000

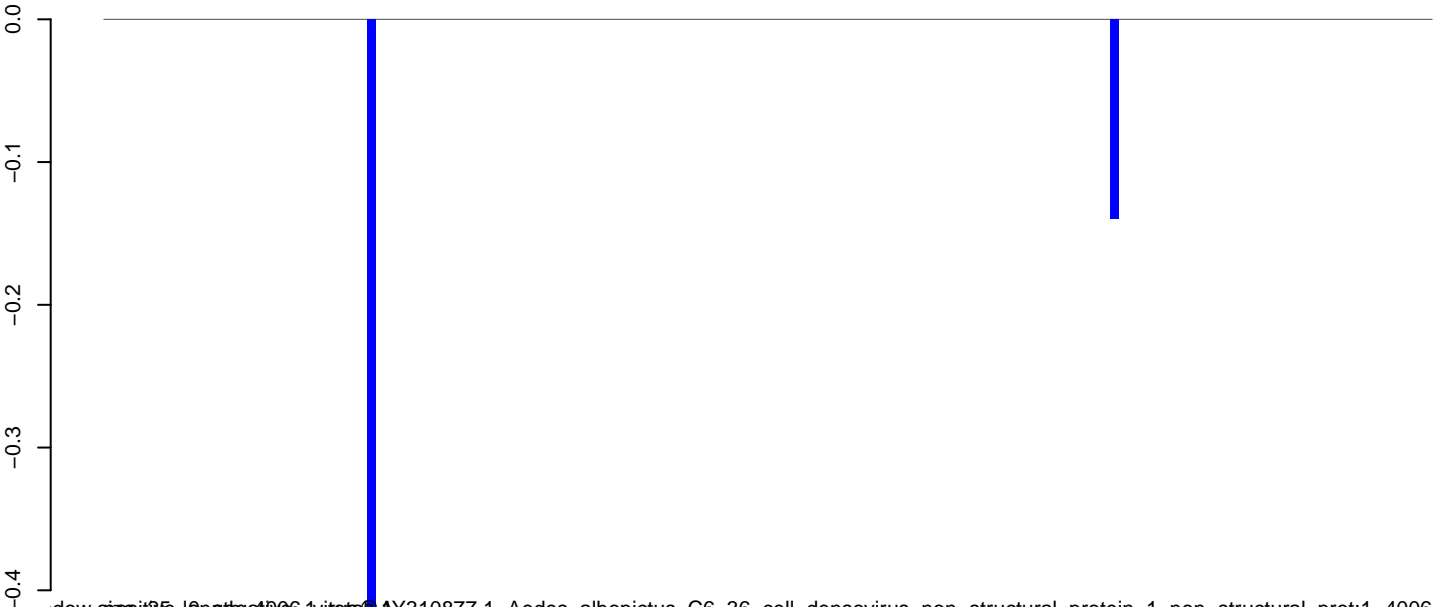
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



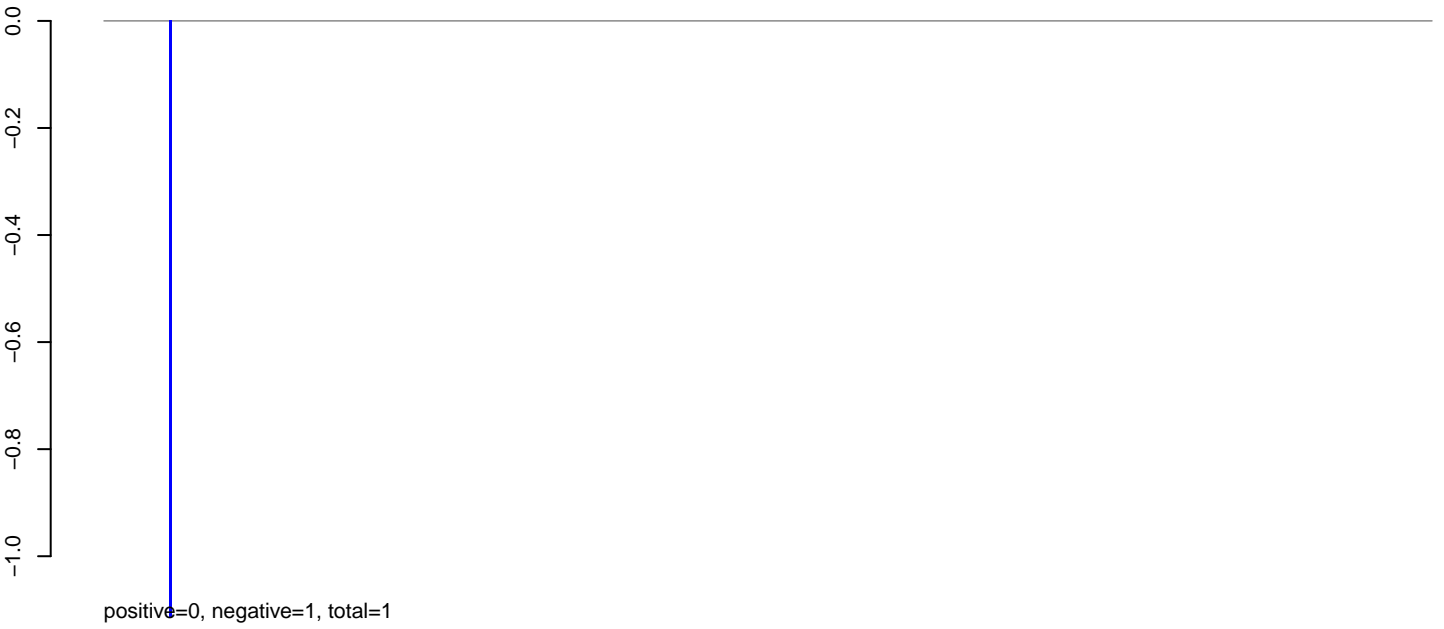
AeAlbo_C636_2dpi_mock3.rep



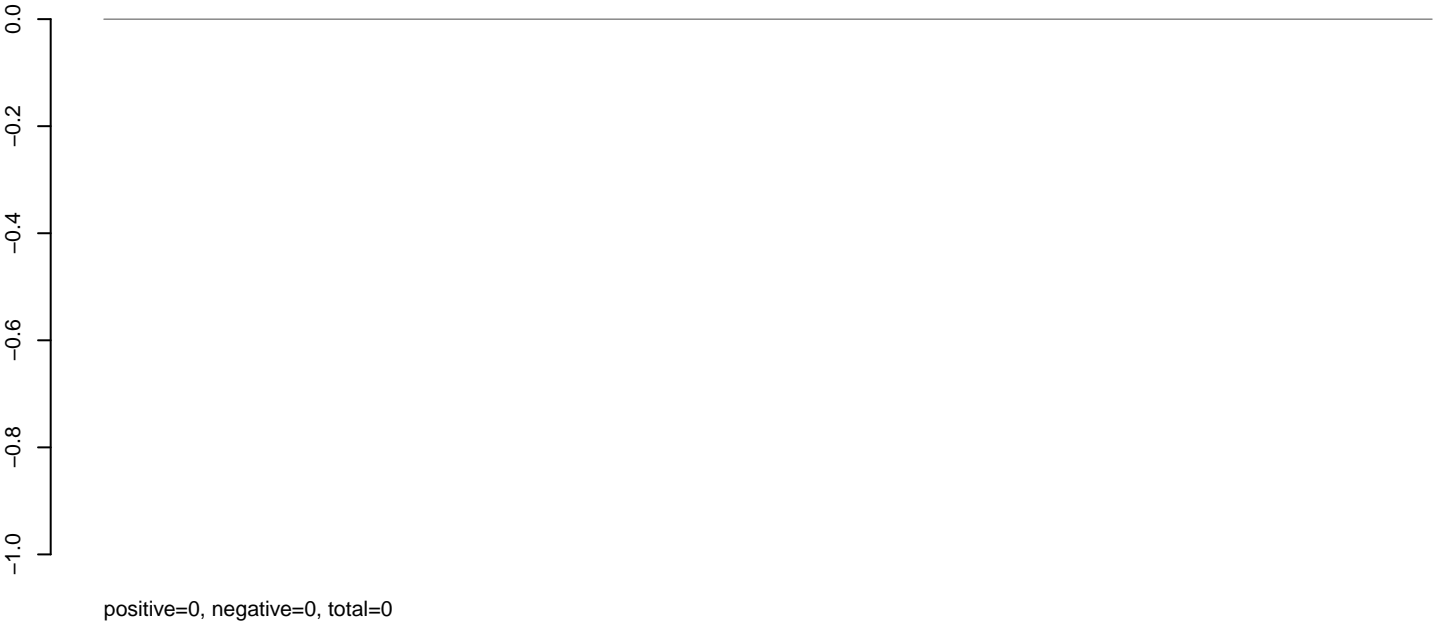
downloaded from [https://www.ncbi.nlm.nih.gov/trace.ncbi.nlm.nih.gov/projects/genome/assembly/AY310877.1-Aedes_albopictus_C6_36_cell_densovirus_non-structural_protein_1_non-structural_prot:1-4006](https://www.ncbi.nlm.nih.gov/trace.ncbi.nlm.nih.gov/projects/genome/assembly/AY310877.1/Aedes_albopictus_C6_36_cell_densovirus_non-structural_protein_1_non-structural_prot:1-4006)

0 1000 2000 3000 4000

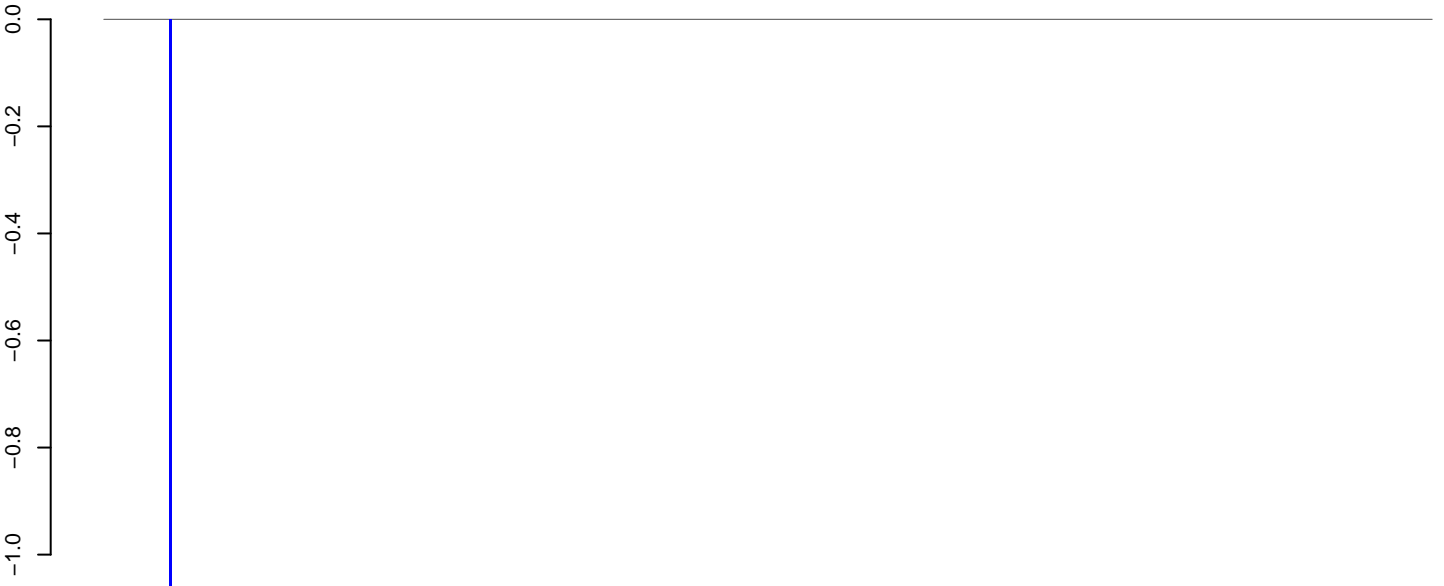
AeAlbo_C636_2dpi_mock3.18_23.rep



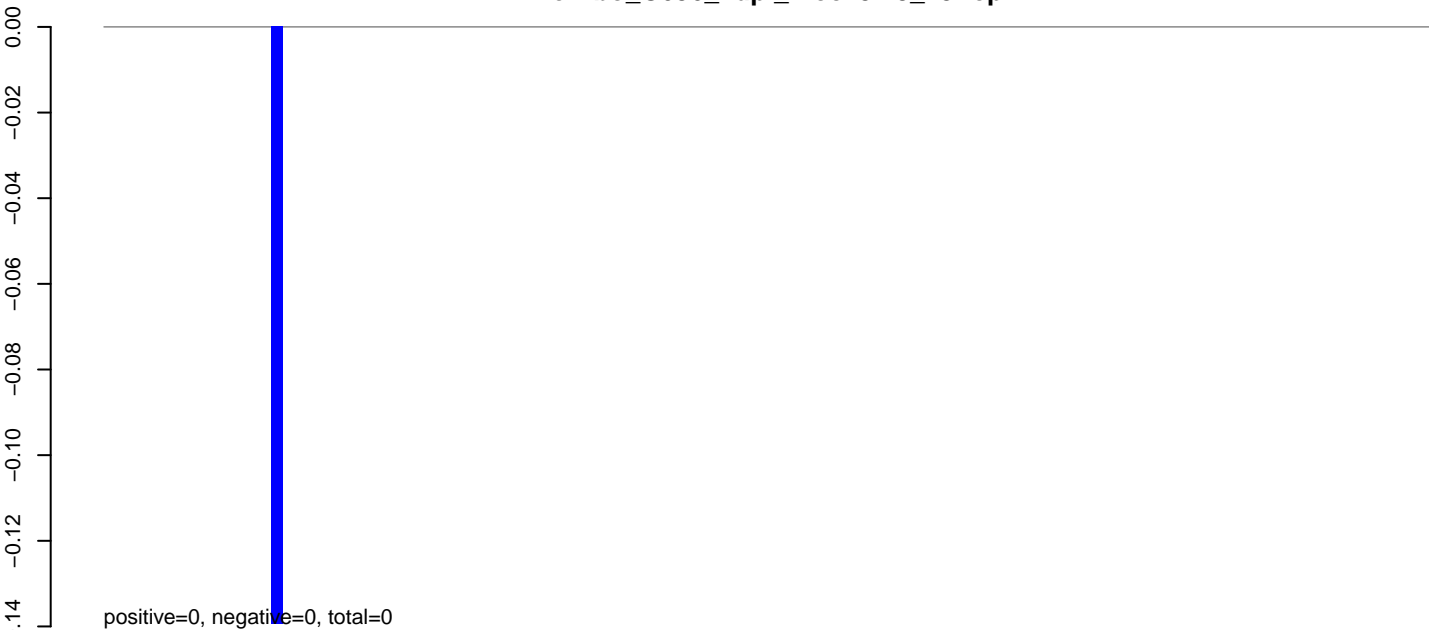
AeAlbo_C636_2dpi_mock3.24_35.rep



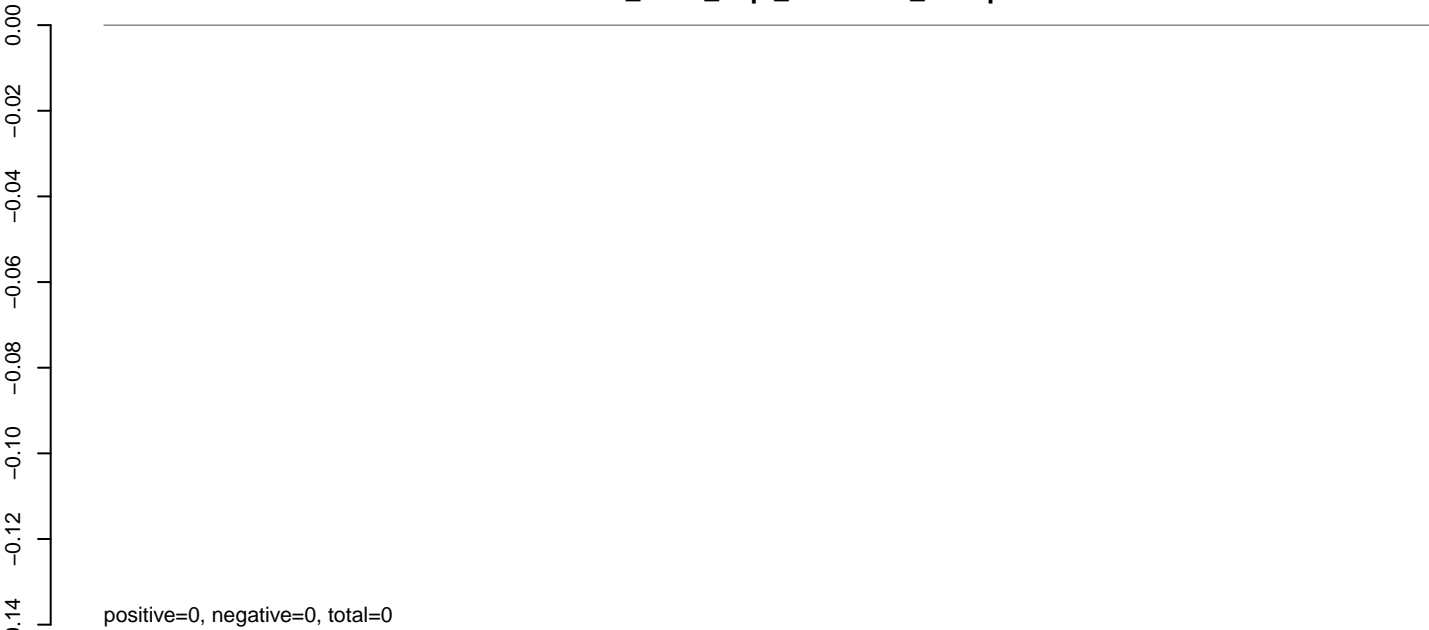
AeAlbo_C636_2dpi_mock3.rep



AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



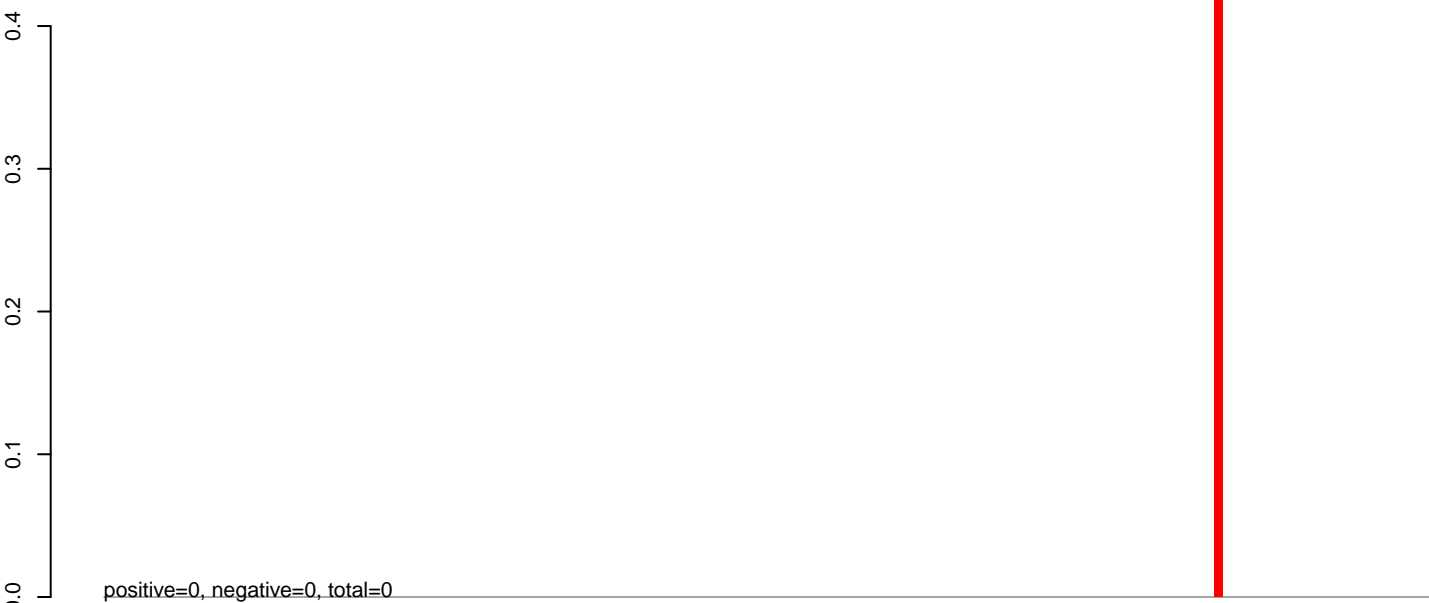
AeAlbo_C636_2dpi_mock3.rep



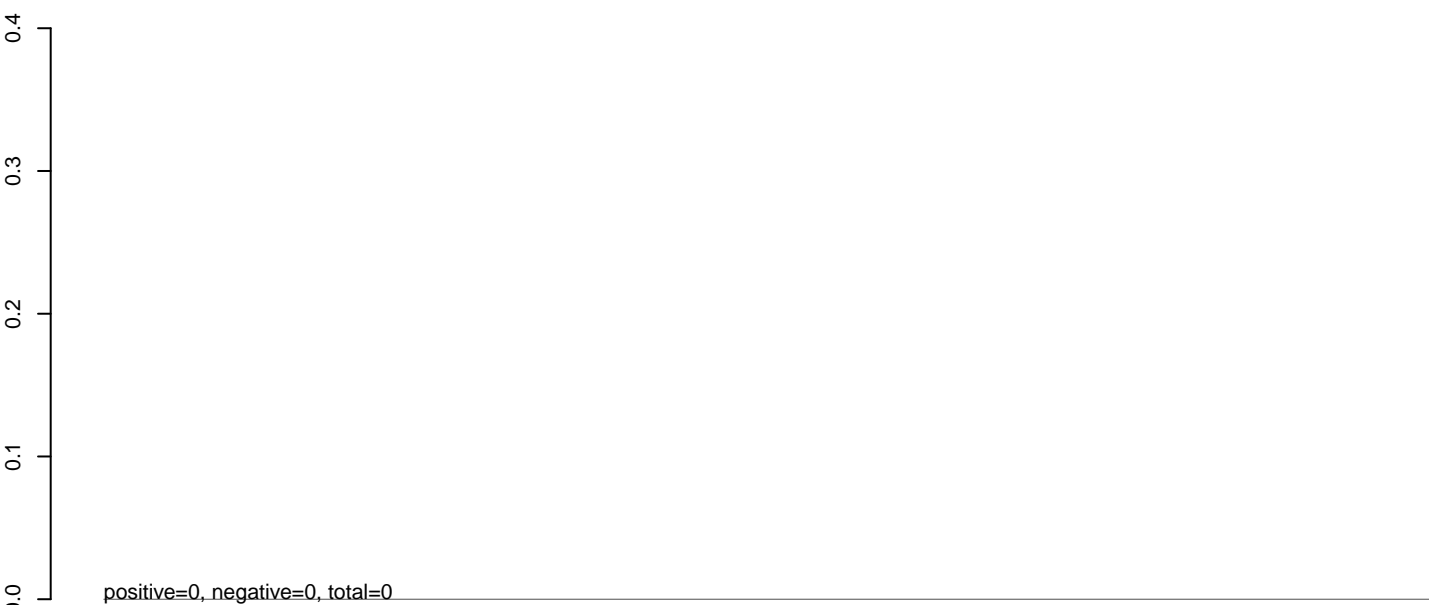
0 500 1000 1500 2000 2500 3000

positive=0, negative=0, total=0 window size=25, length=2968, virus@NC_035126.1_Wilkie_narna-like_virus_1_strain_mosWSCP70929:1-2968

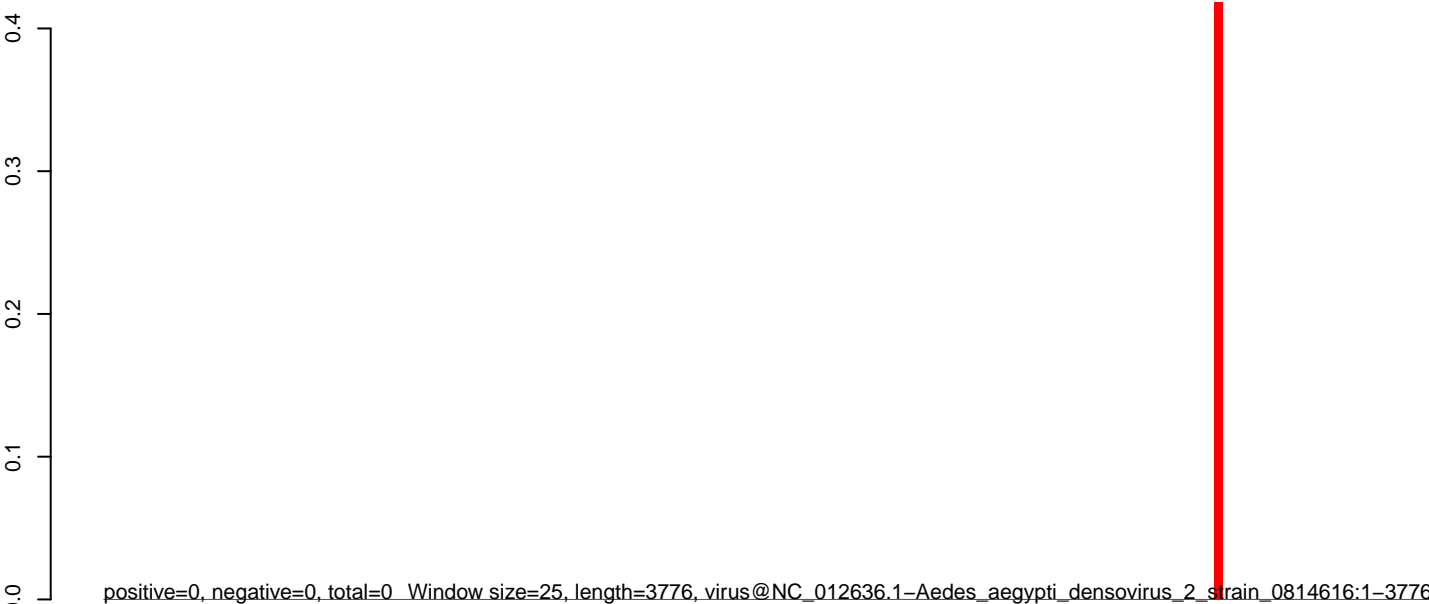
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

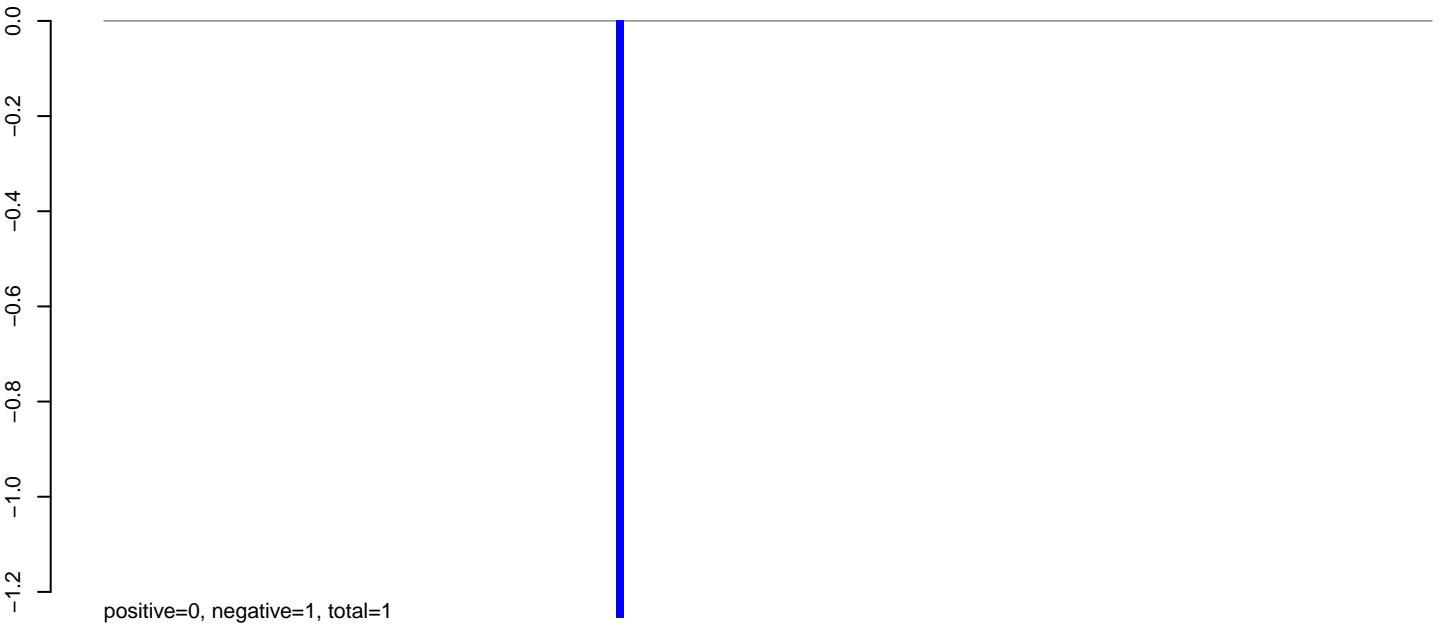


AeAlbo_C636_2dpi_mock3.rep

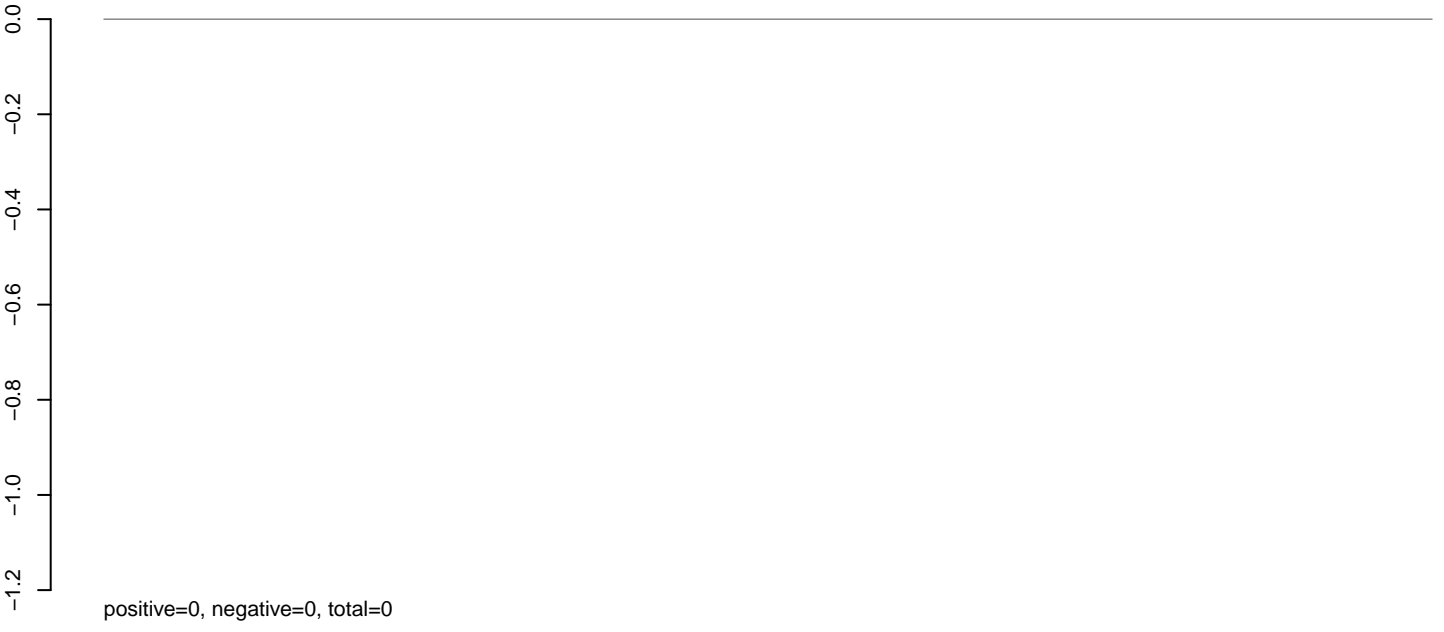


0 1000 2000 3000

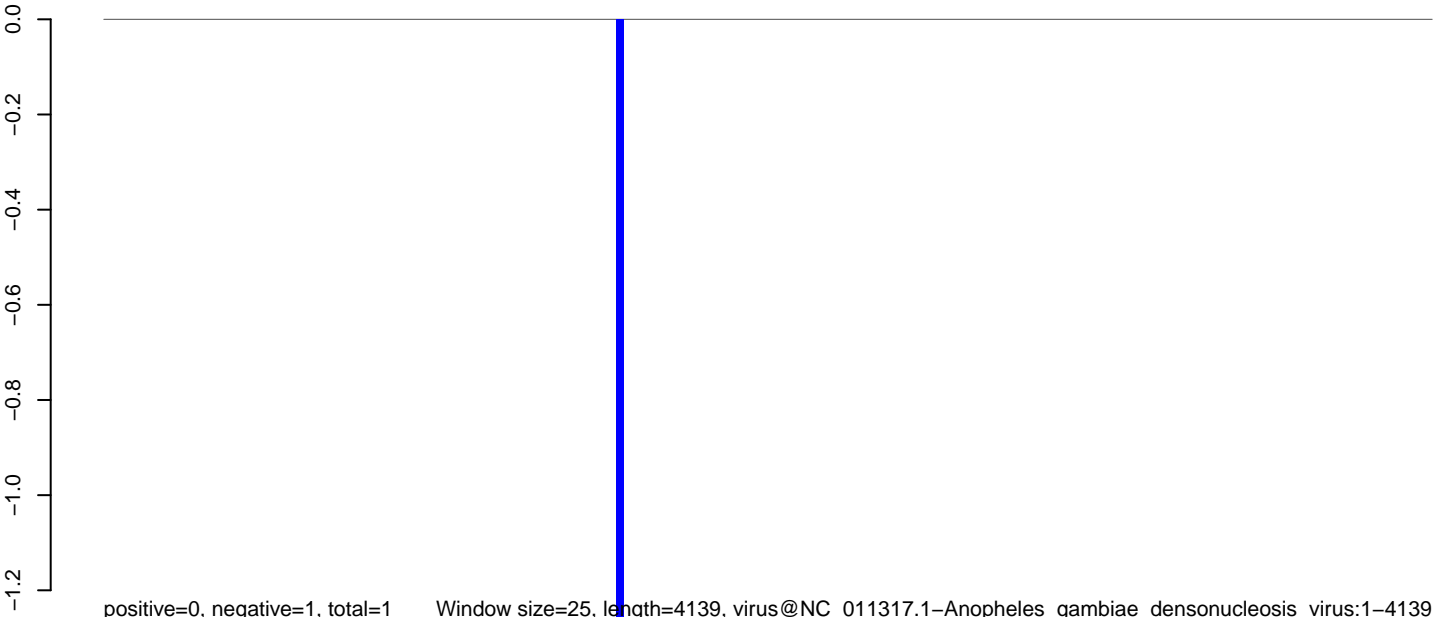
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



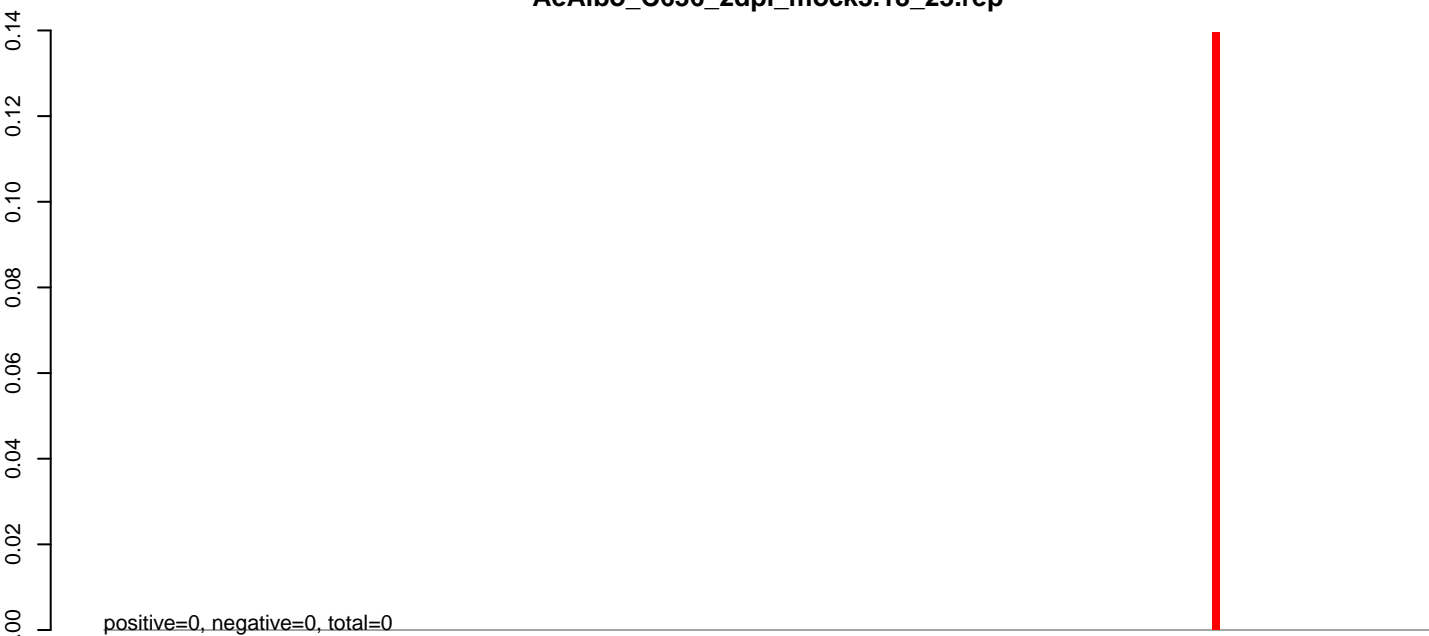
AeAlbo_C636_2dpi_mock3.rep



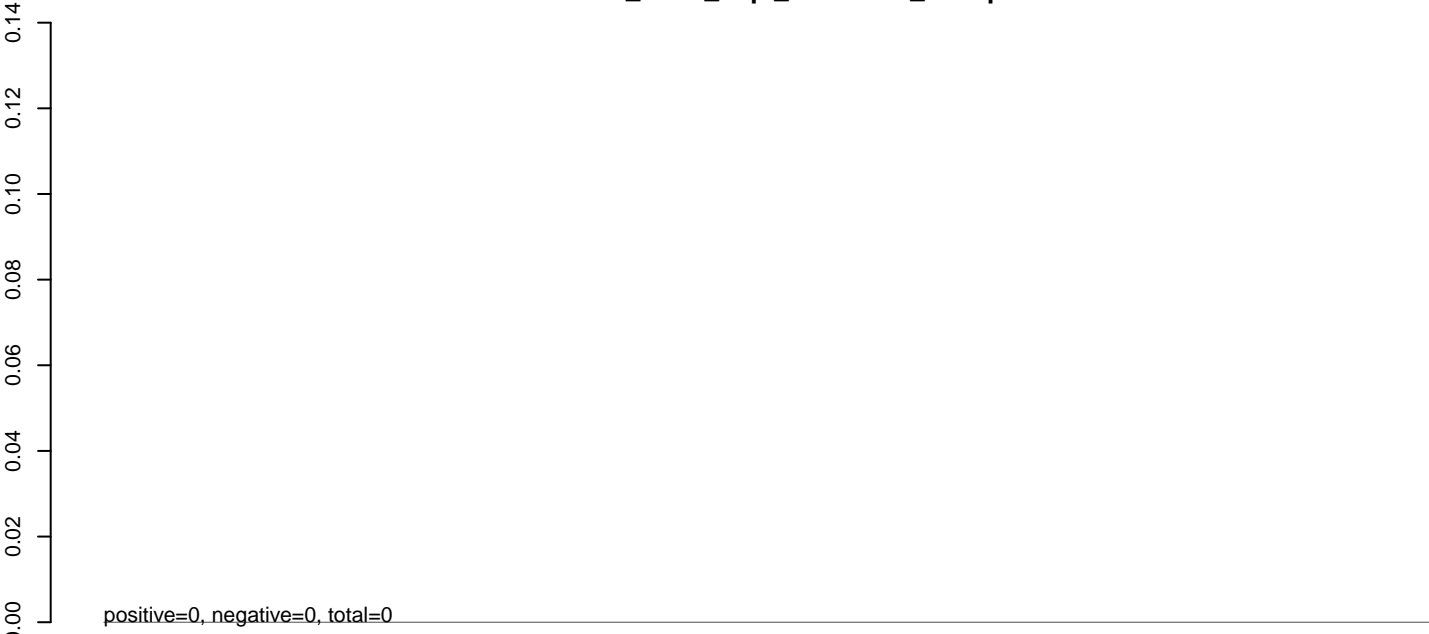
Window size=25, length=4139, virus@NC_011317.1-Anopheles_gambiae_densonucleosis_virus:1-4139

0 1000 2000 3000 4000

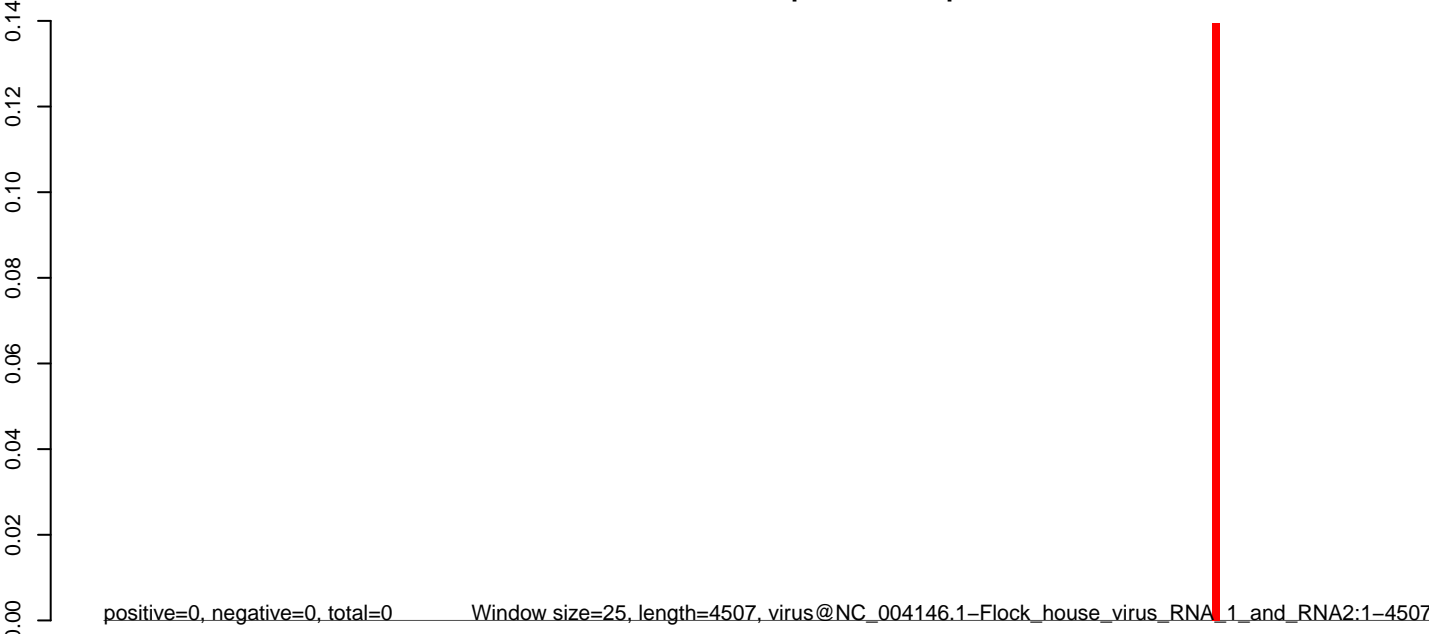
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

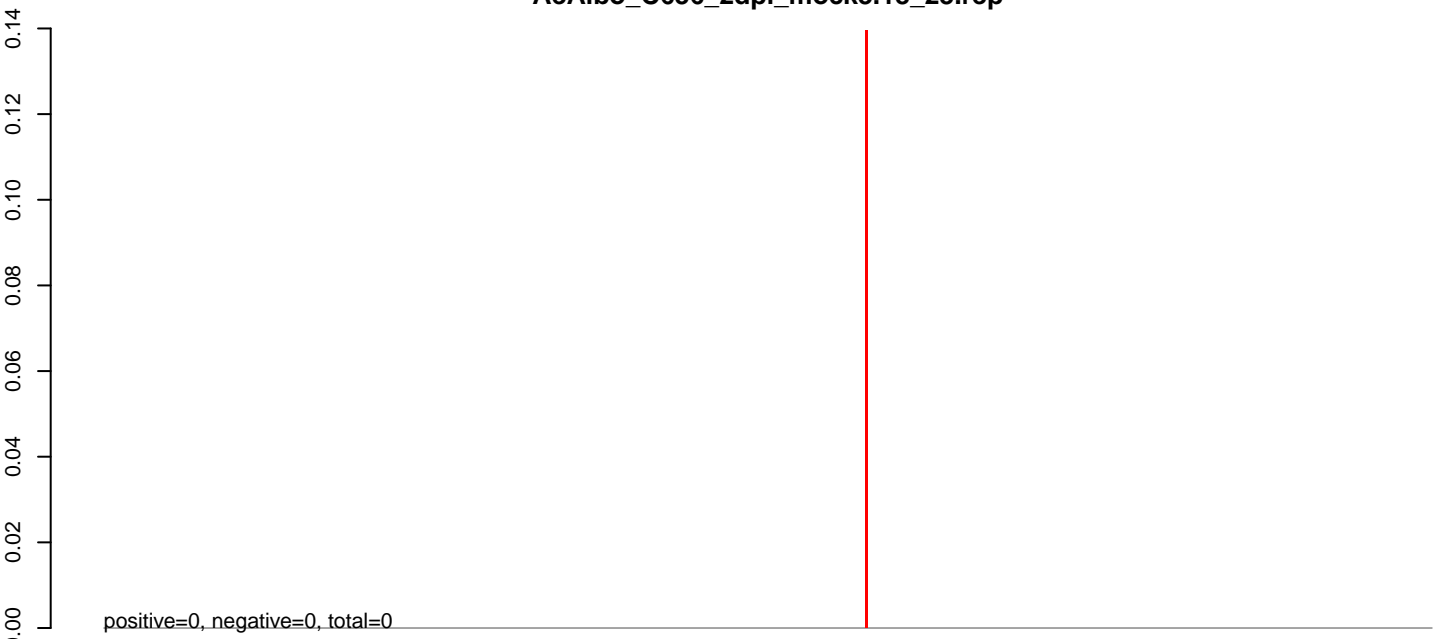


AeAlbo_C636_2dpi_mock3.rep

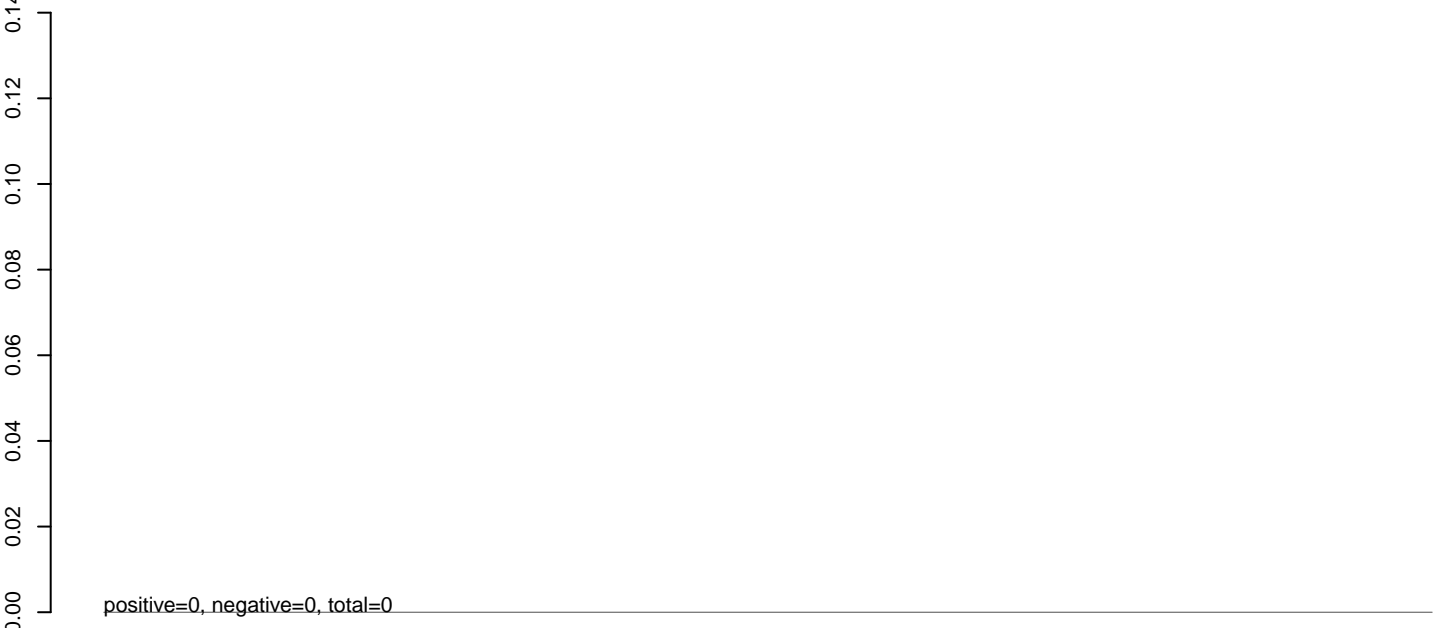


0 1000 2000 3000 4000

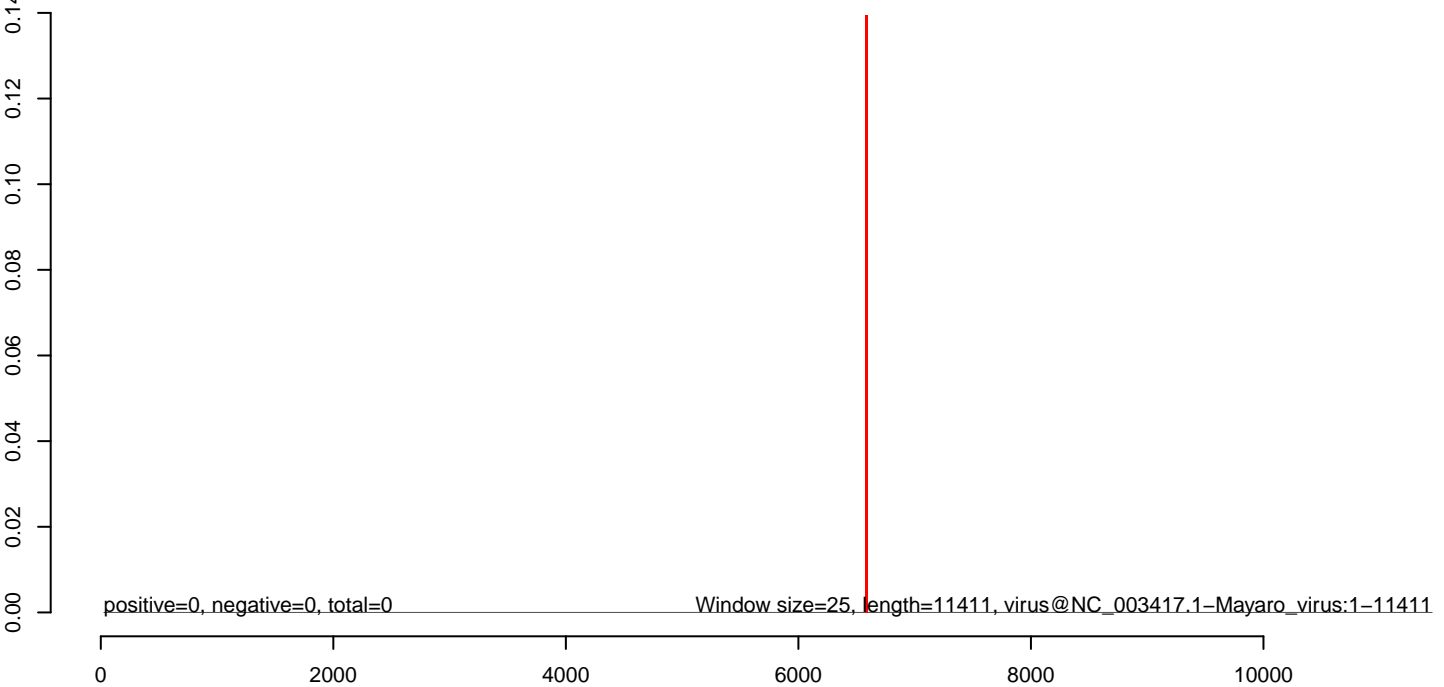
AeAlbo_C636_2dpi_mock3.18_23.rep



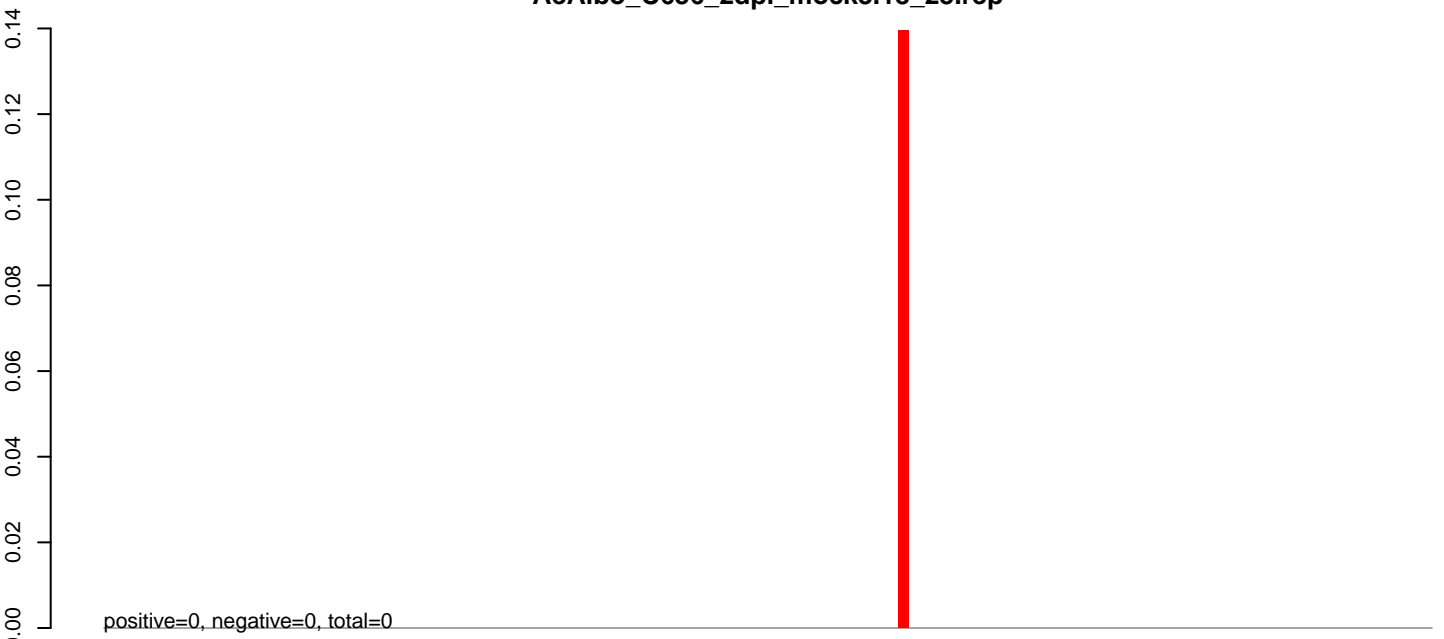
AeAlbo_C636_2dpi_mock3.24_35.rep



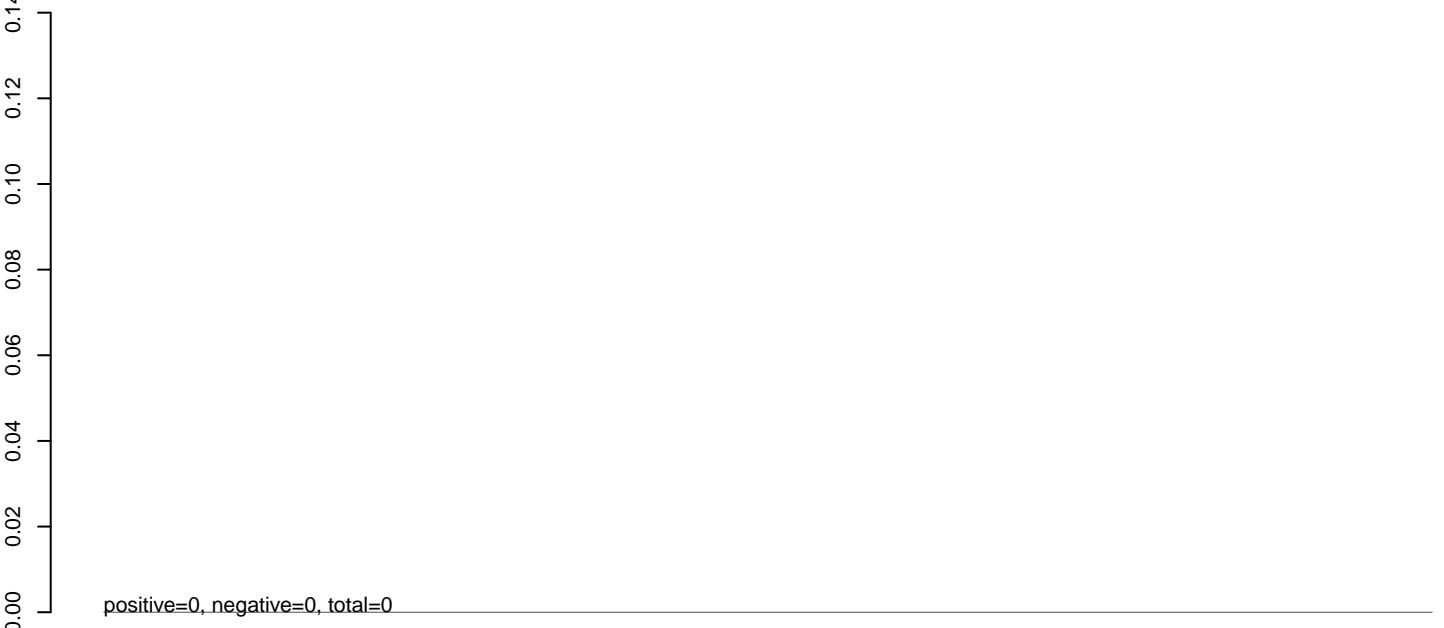
AeAlbo_C636_2dpi_mock3.rep



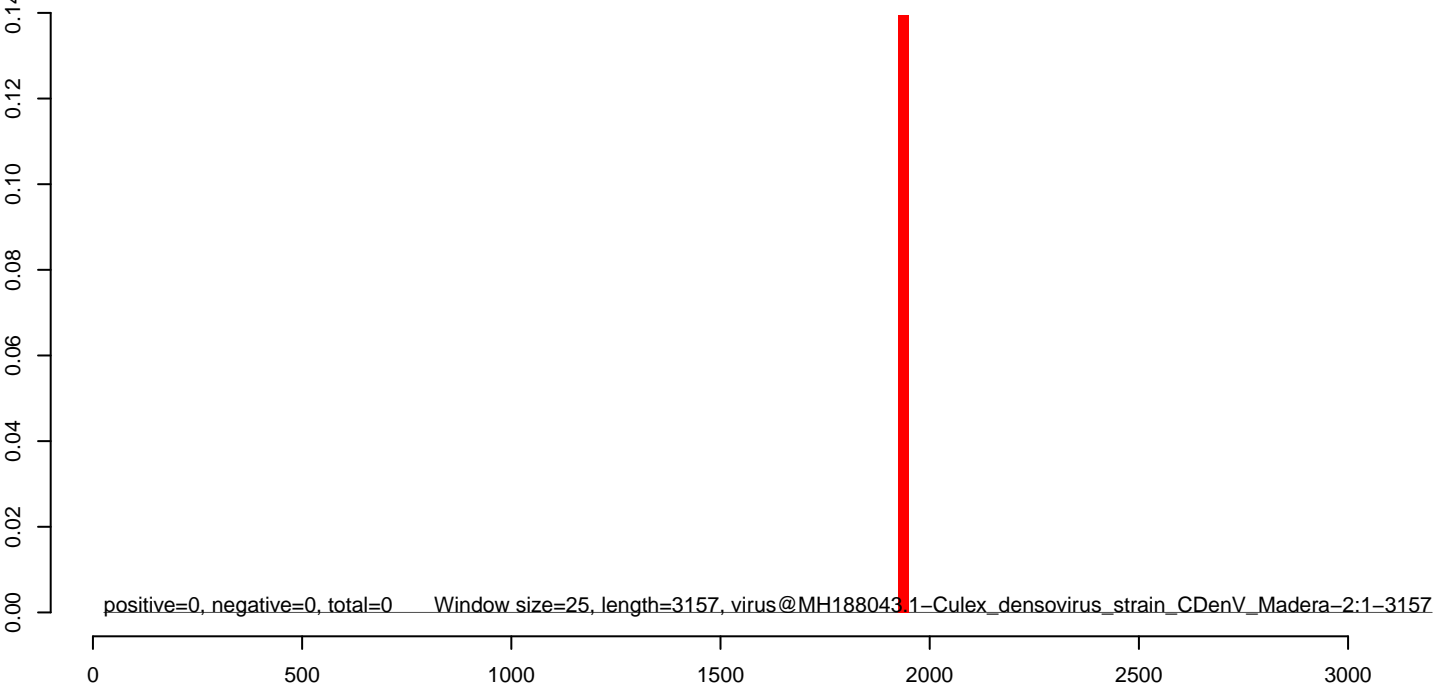
AeAlbo_C636_2dpi_mock3.18_23.rep



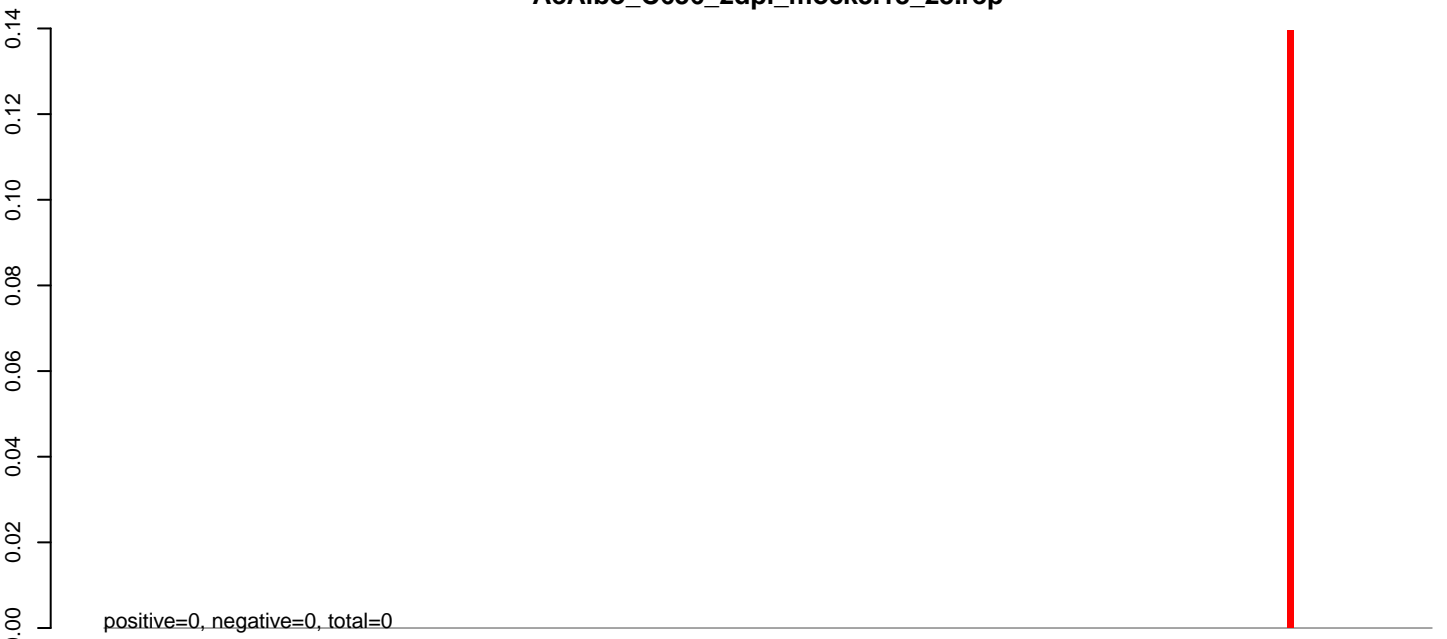
AeAlbo_C636_2dpi_mock3.24_35.rep



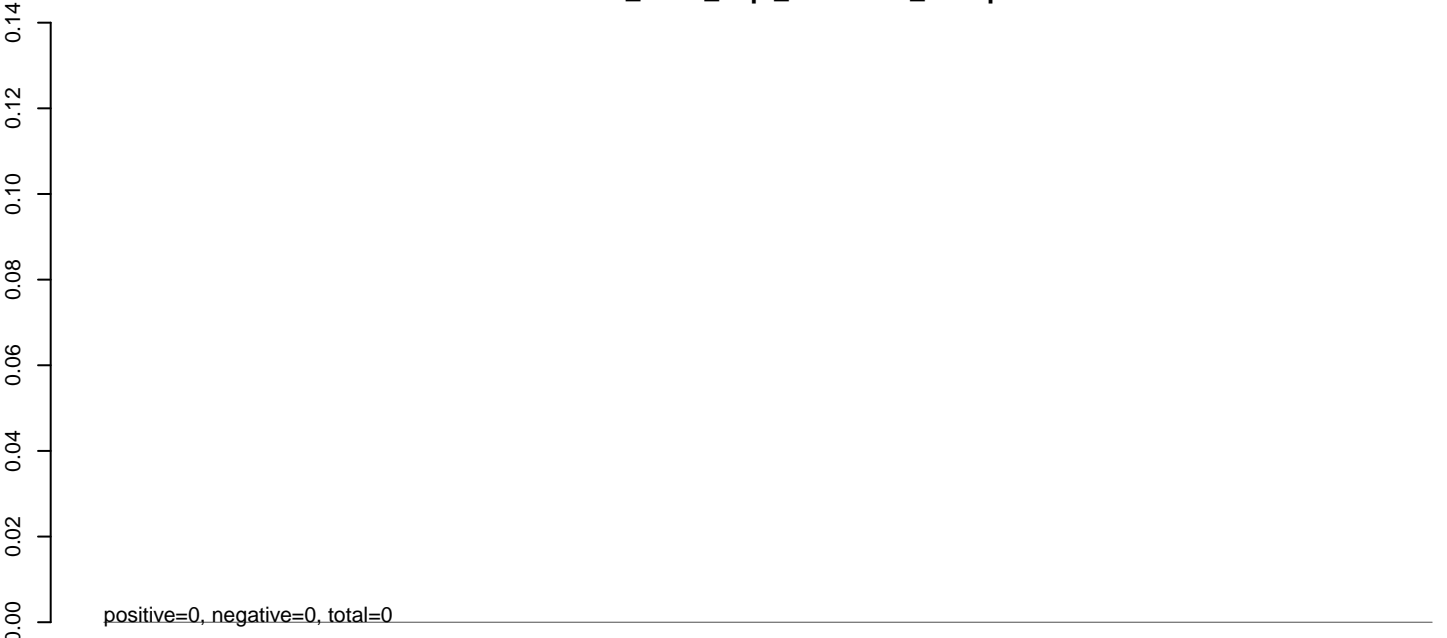
AeAlbo_C636_2dpi_mock3.rep



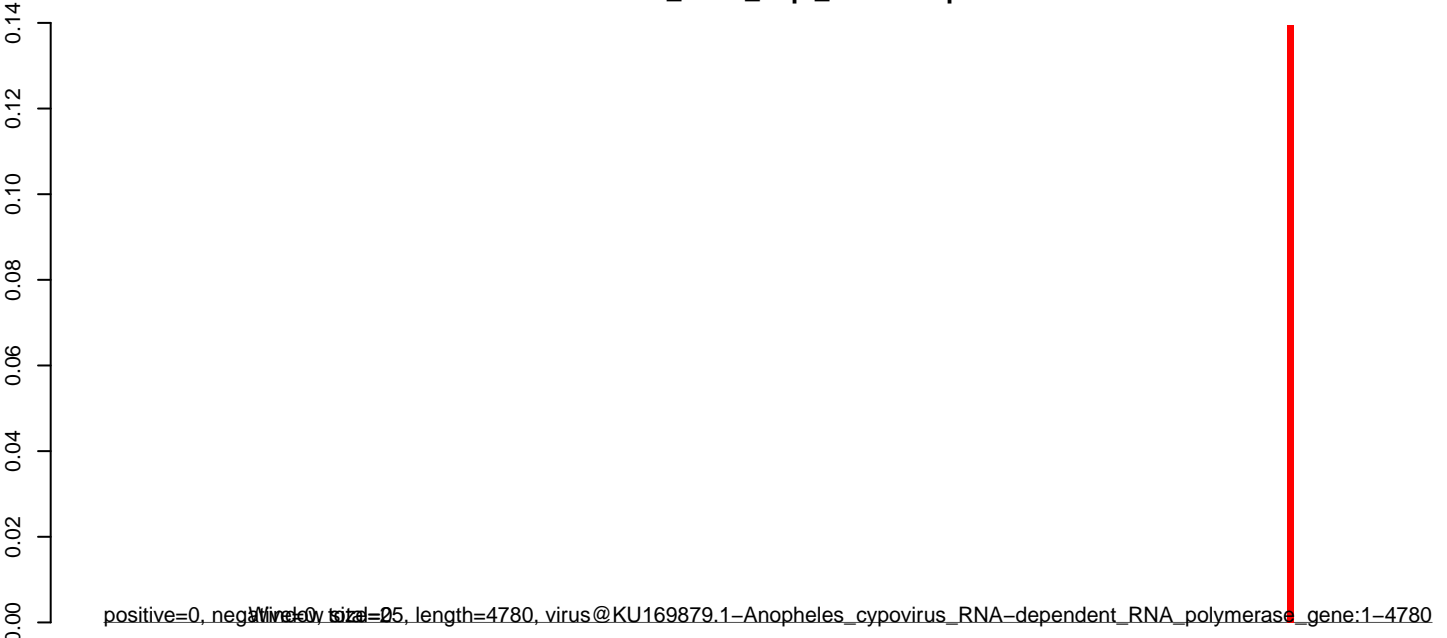
AeAlbo_C636_2dpi_mock3.18_23.rep



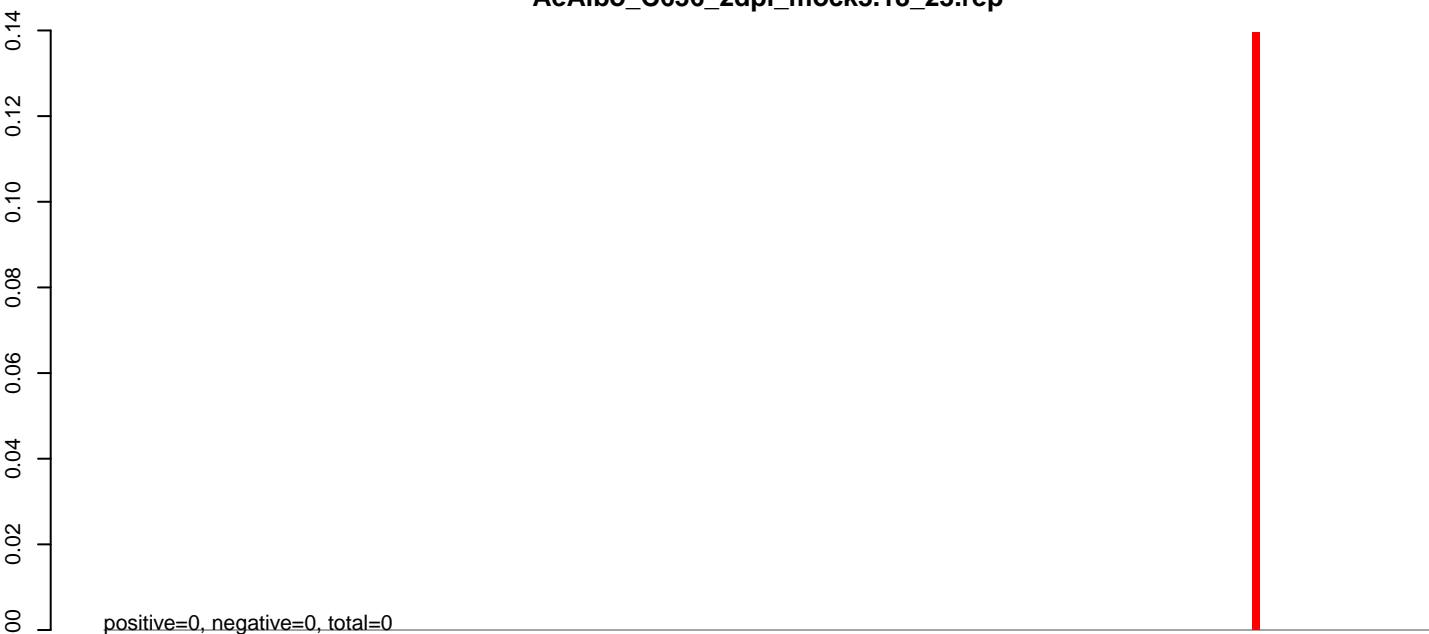
AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



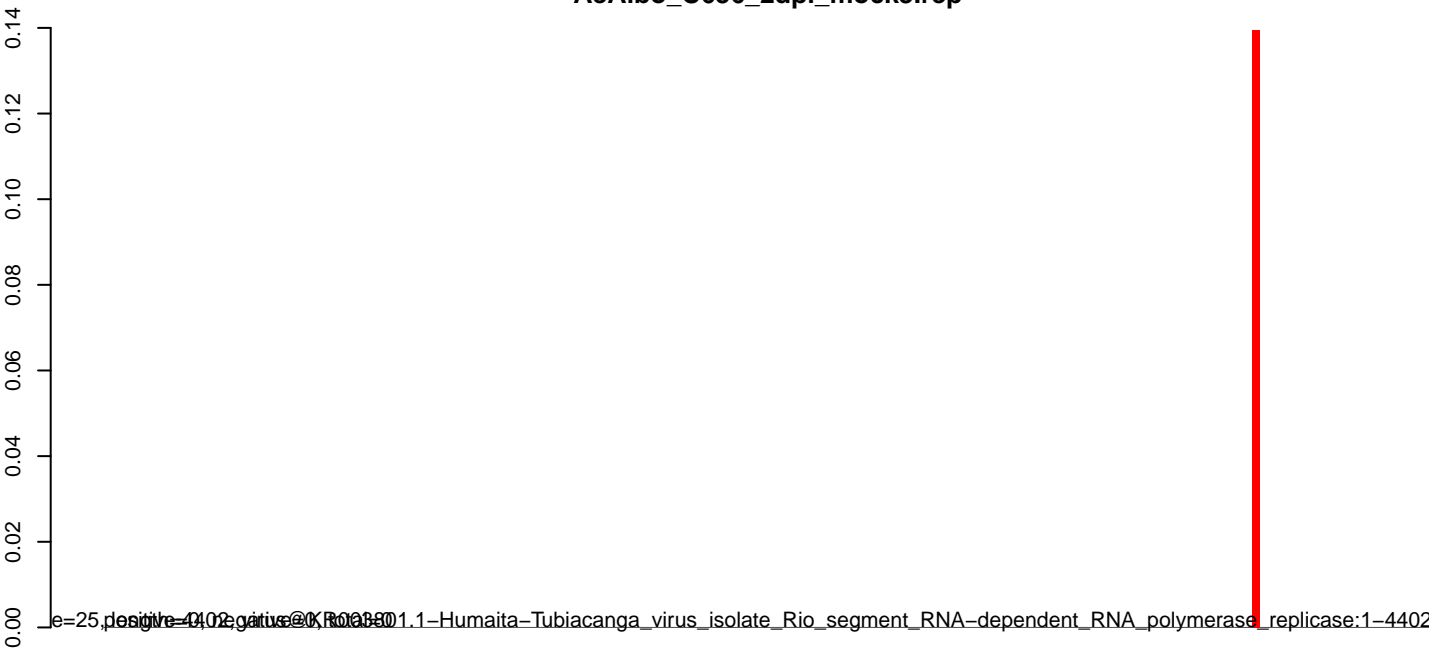
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep

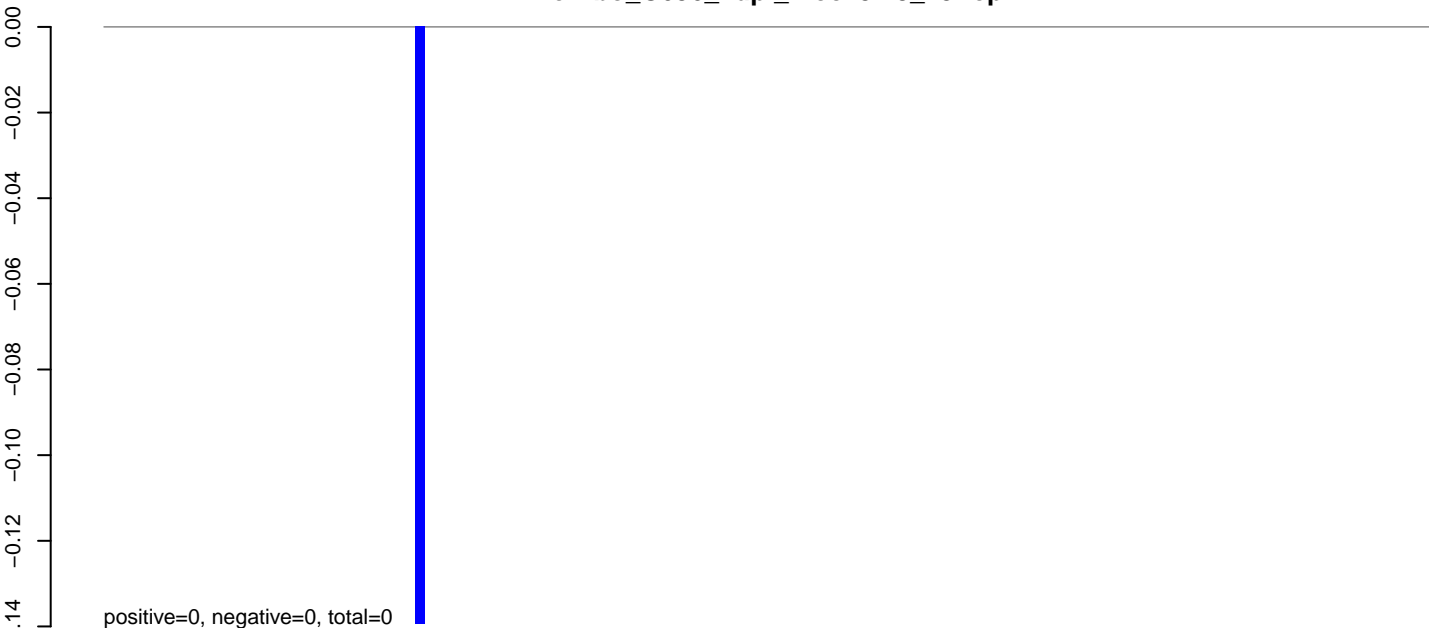


AeAlbo_C636_2dpi_mock3.rep

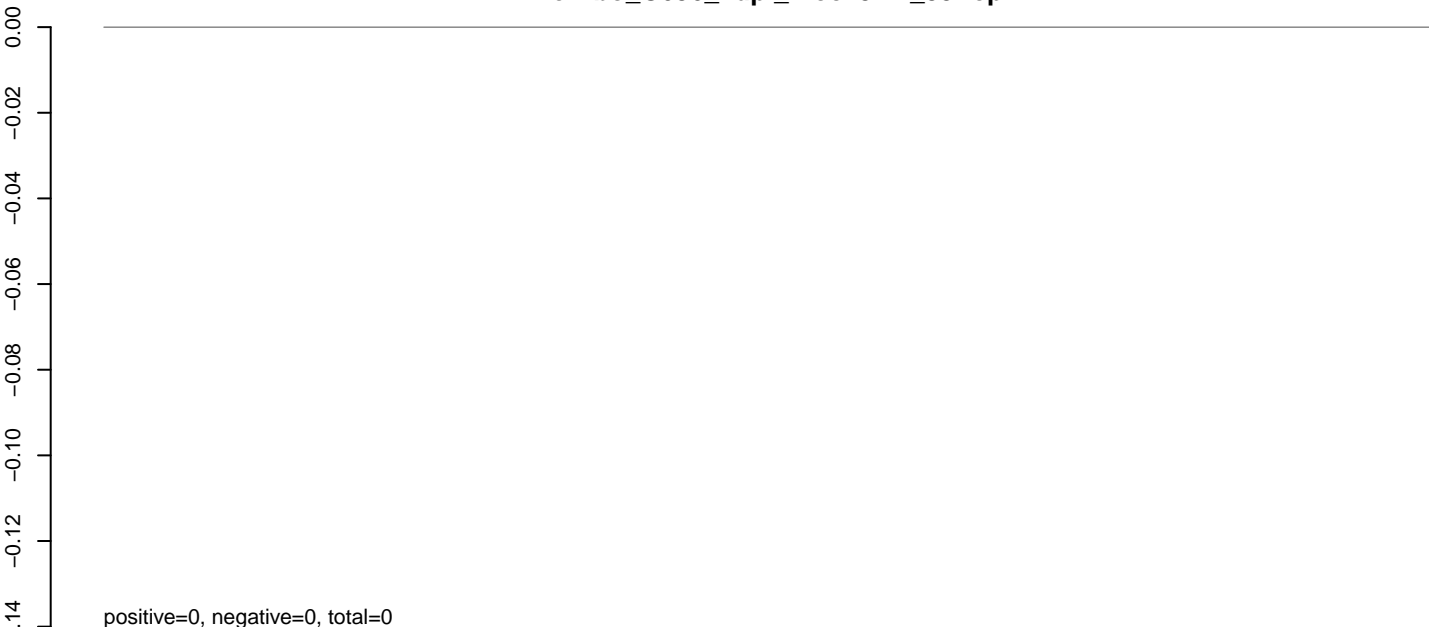


1.1-Humaita-Tubianga_virus_isolate_Rio_segment_RNA-dependent_RNA_polymerase_replicase:1-4402

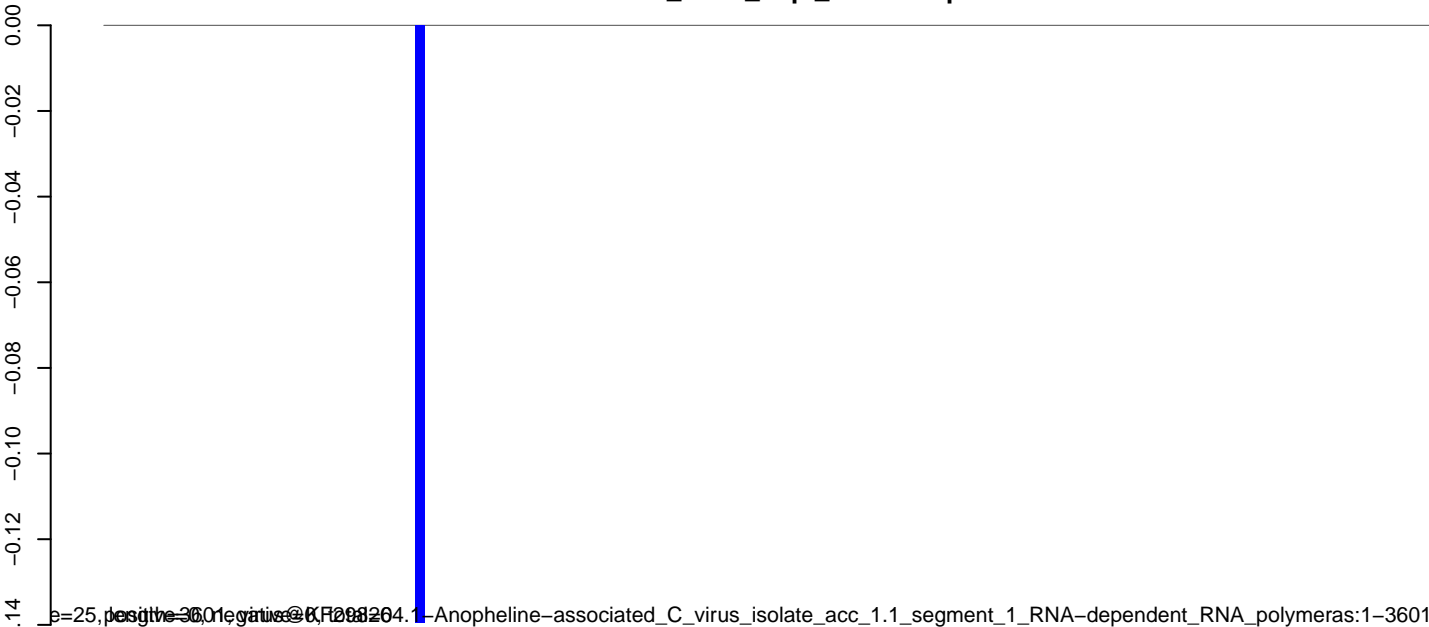
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



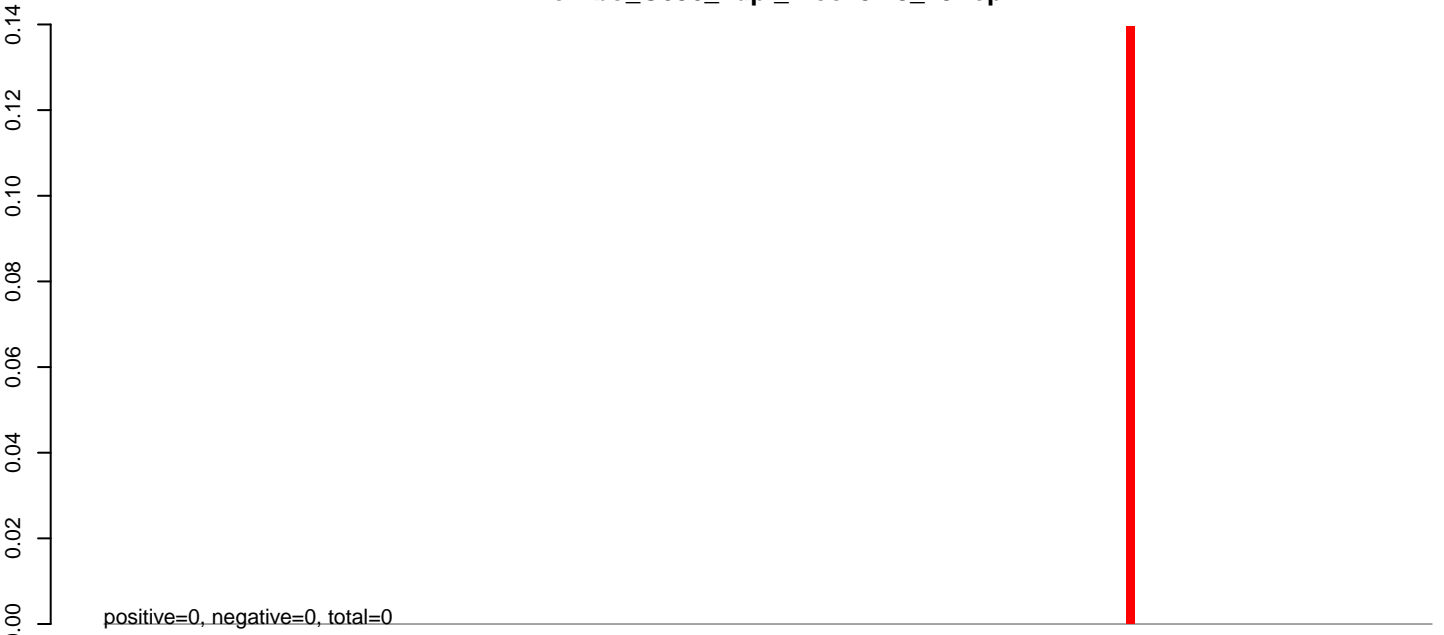
AeAlbo_C636_2dpi_mock3.rep



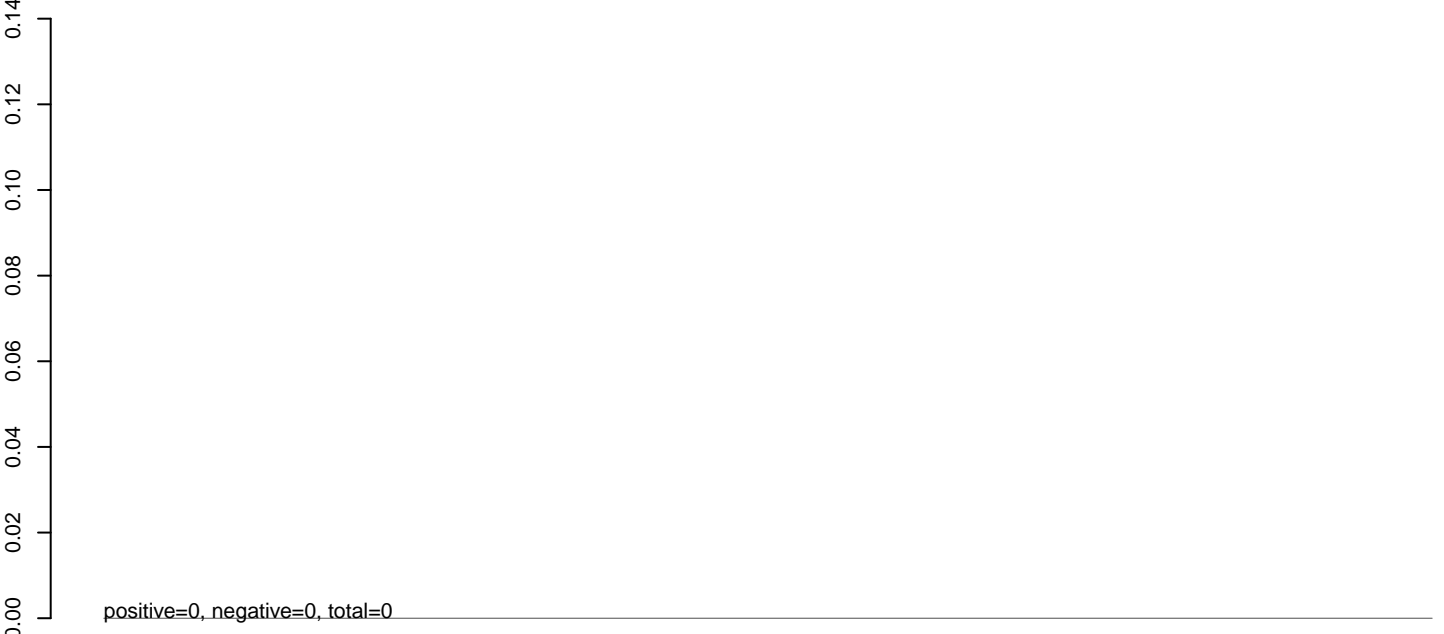
25, positive=0, negative=0, total=0.1264.1-Anopheline-associated_C_virus_isolate_acc.1.1_segment_1_RNA-dependent_RNA_polymerase:1-3601

0 1000 2000 3000

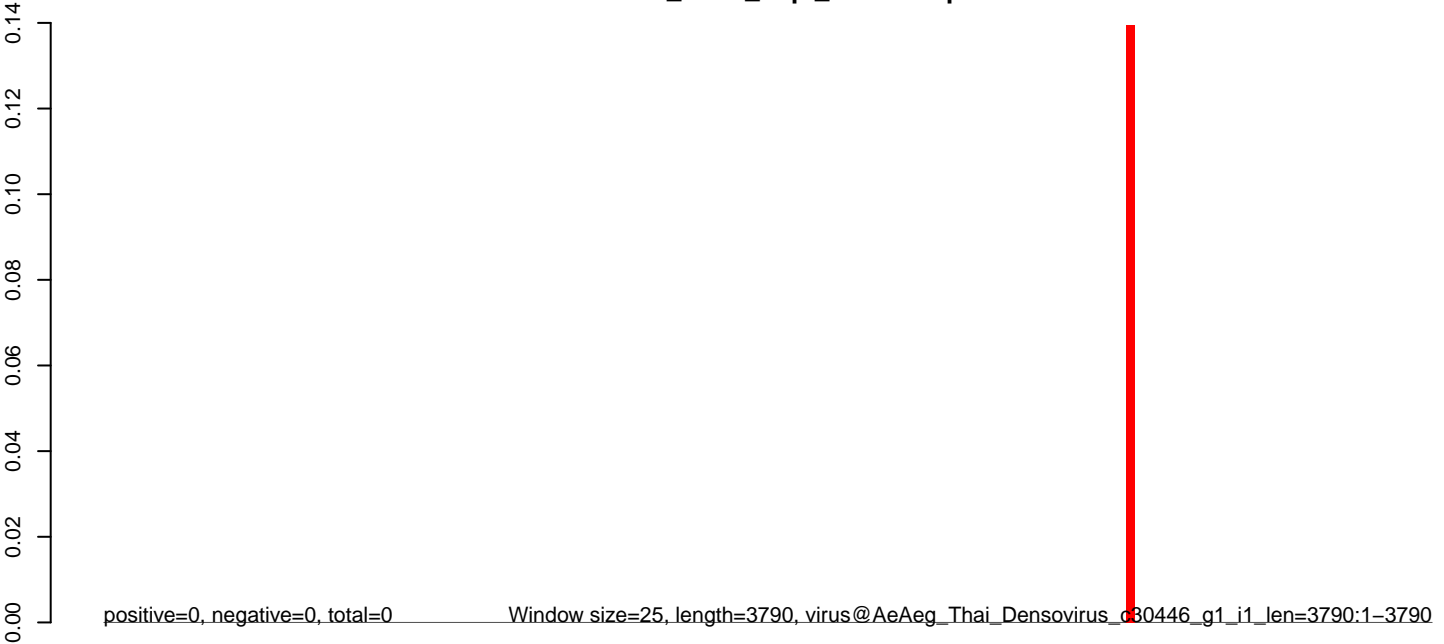
AeAlbo_C636_2dpi_mock3.18_23.rep



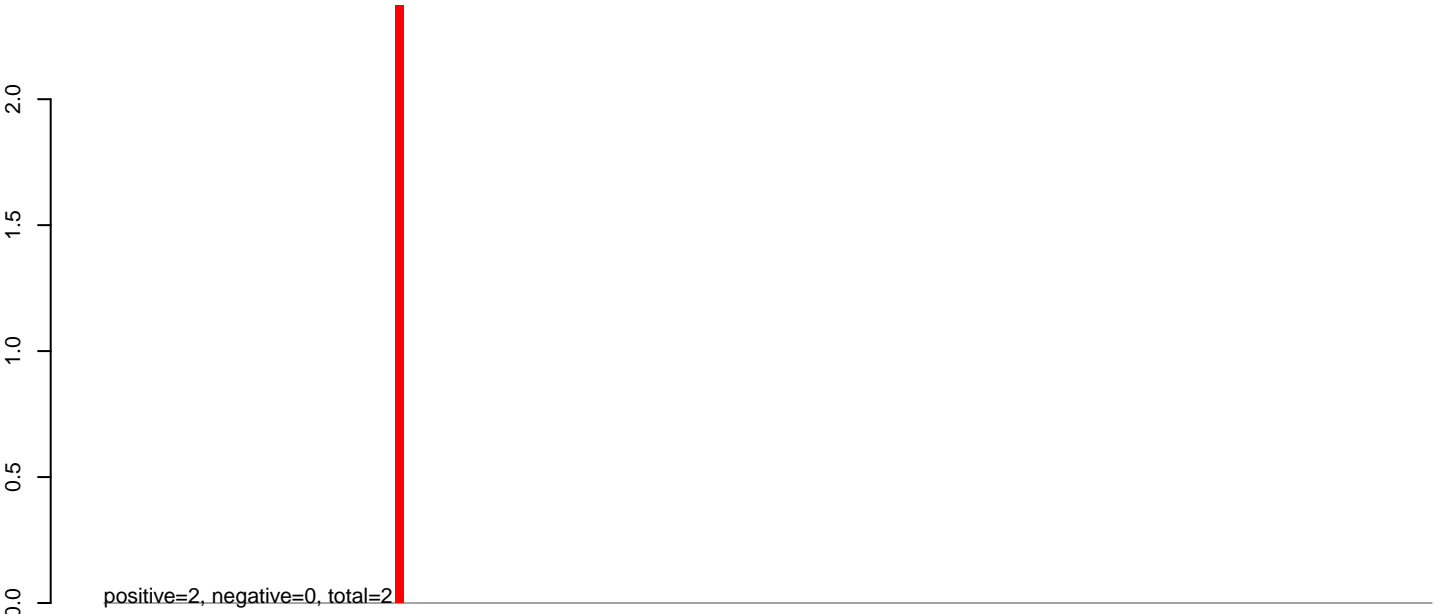
AeAlbo_C636_2dpi_mock3.24_35.rep



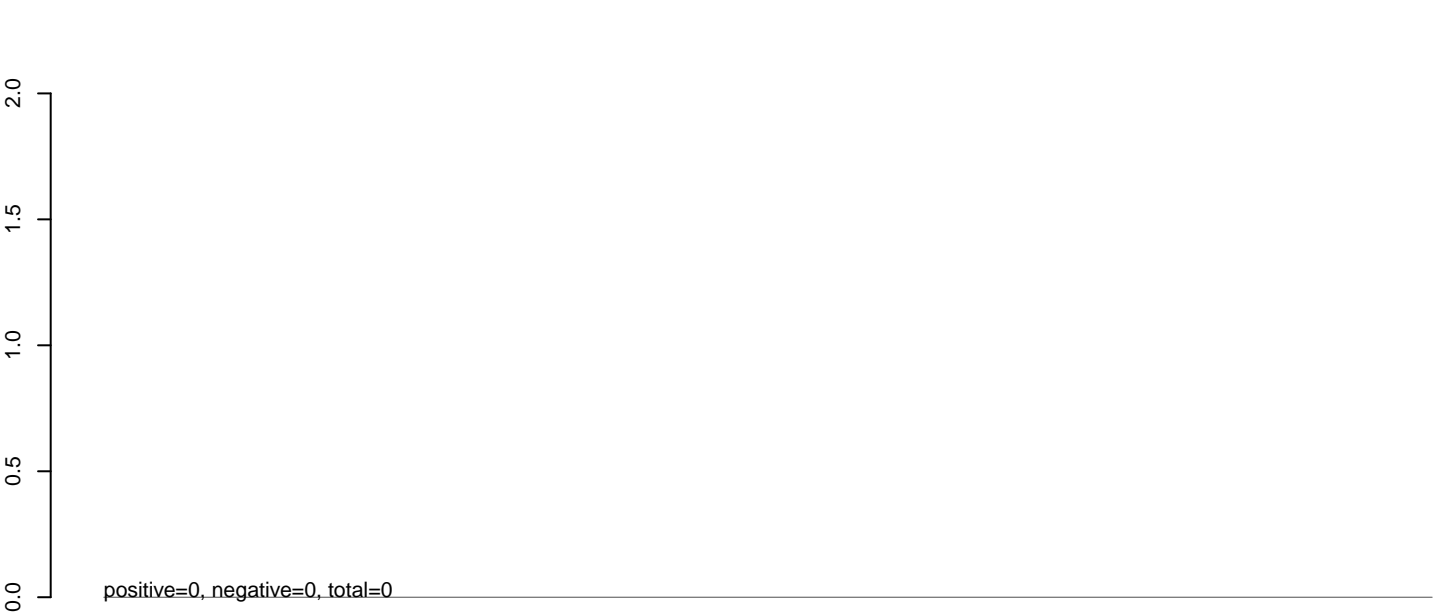
AeAlbo_C636_2dpi_mock3.rep



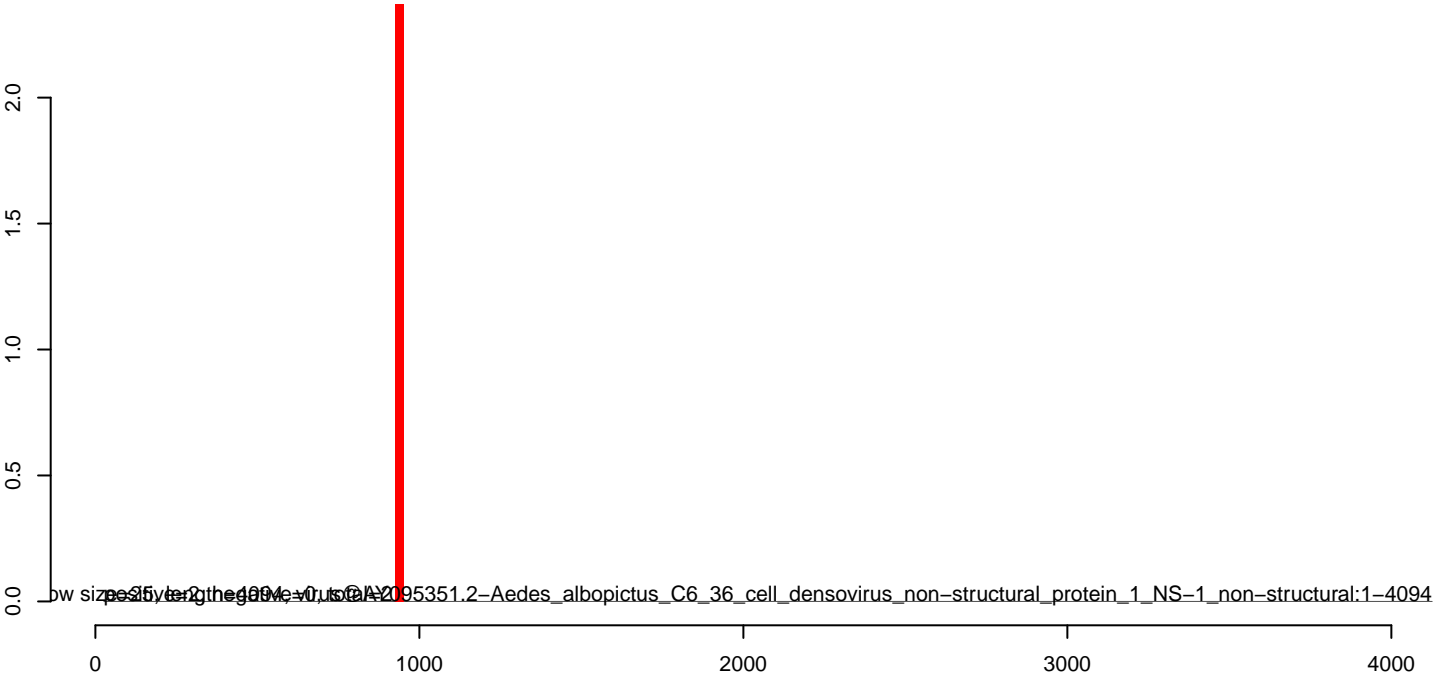
AeAlbo_C636_2dpi_mock3.18_23.rep



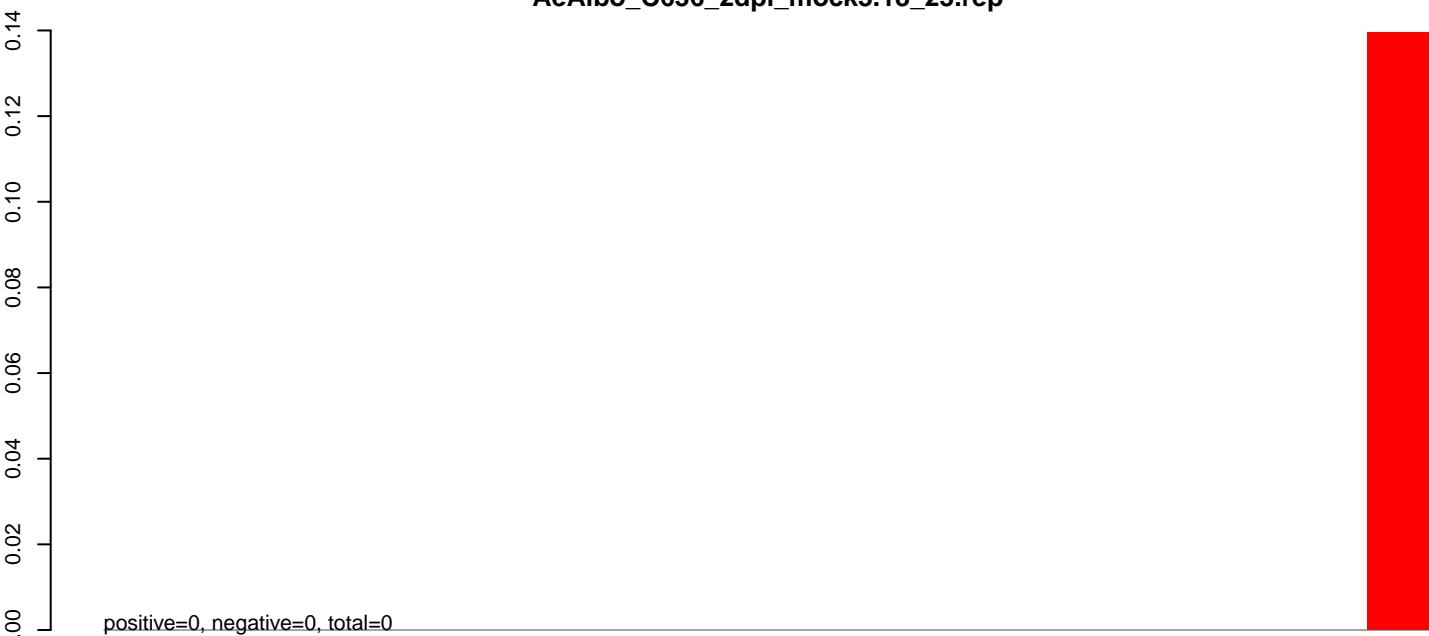
AeAlbo_C636_2dpi_mock3.24_35.rep



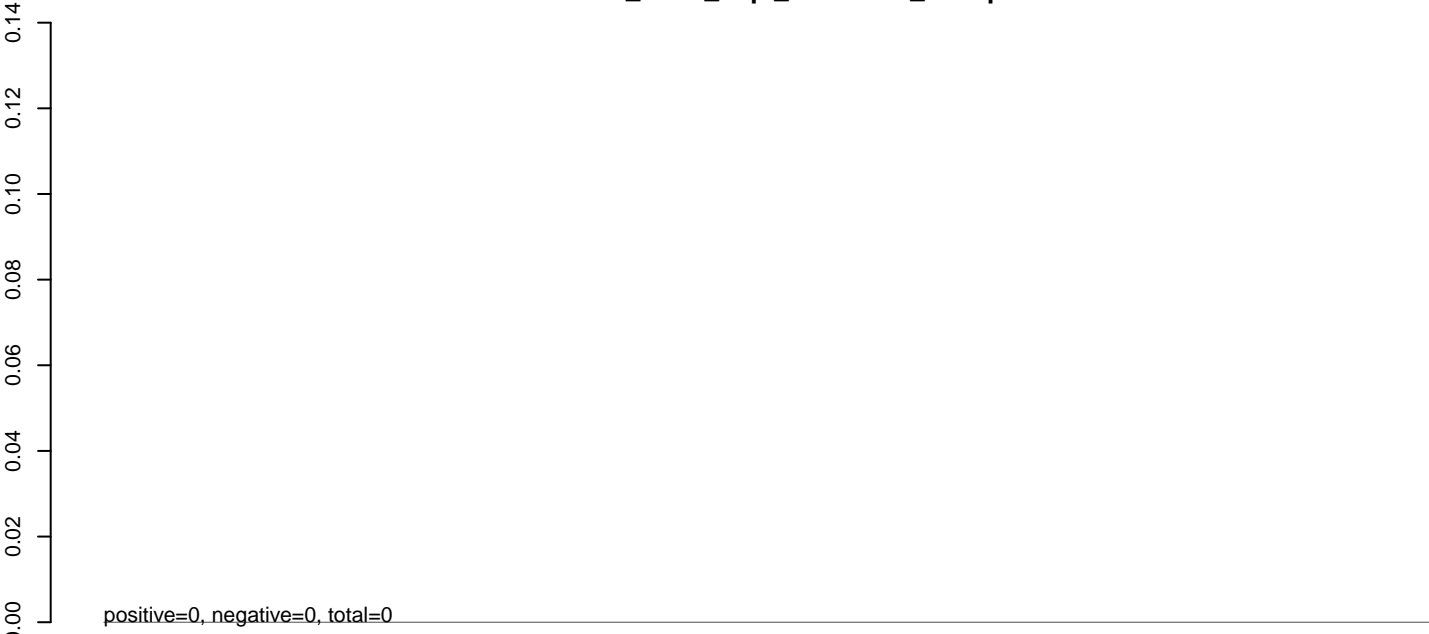
AeAlbo_C636_2dpi_mock3.rep



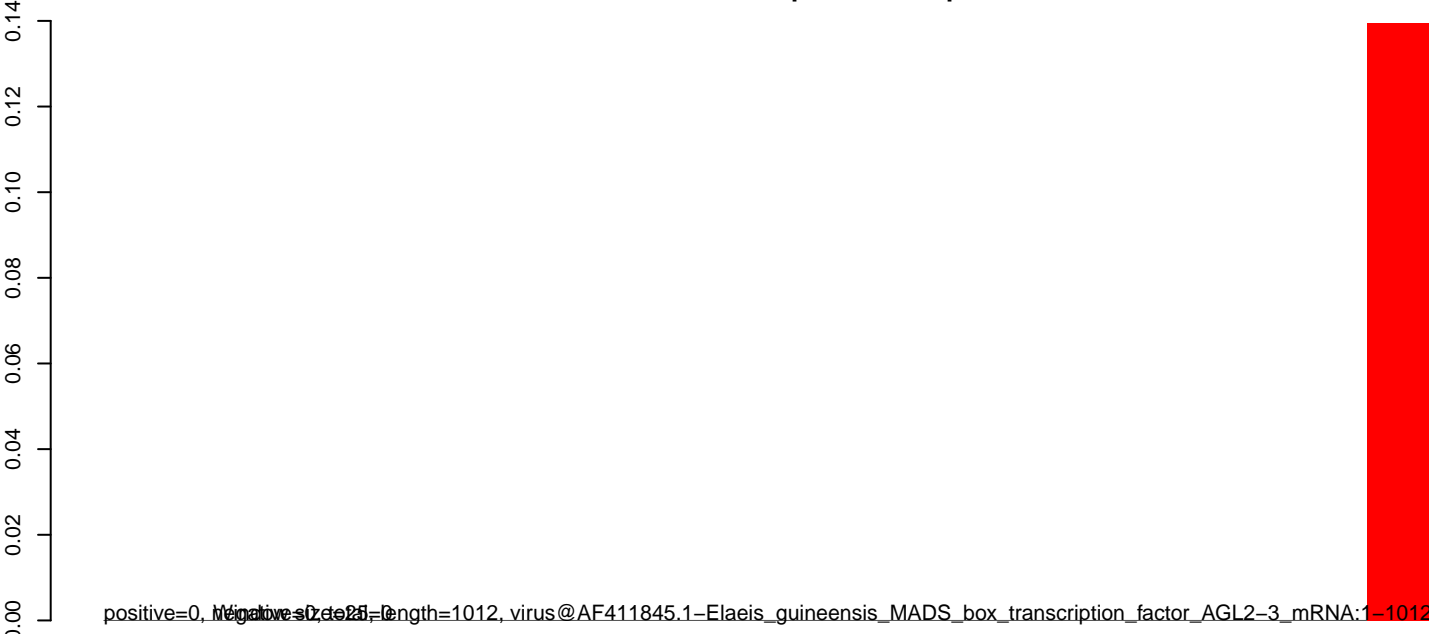
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



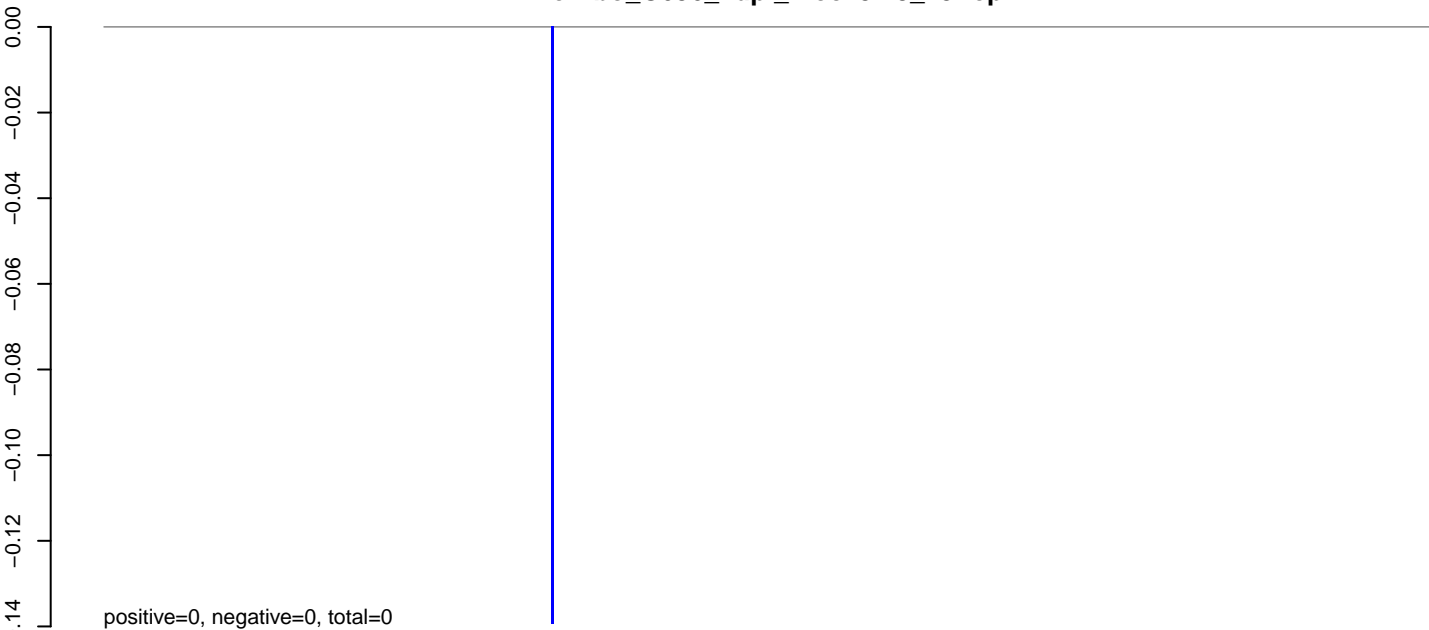
AeAlbo_C636_2dpi_mock3.rep



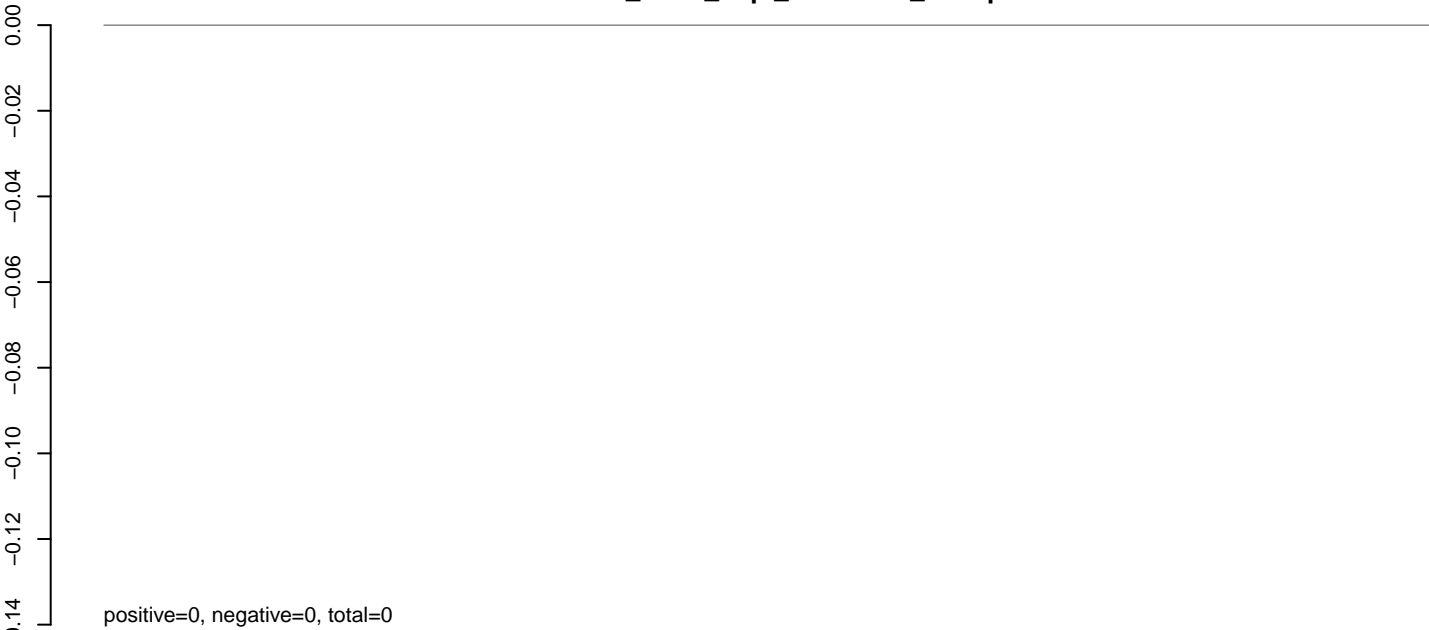
0 200 400 600 800 1000

positive=0, negative=0, total=0, length=1012, virus@AF411845.1-Elaeis_guineensis_MADS_box_transcription_factor_AGL2-3_mRNA:1-1012

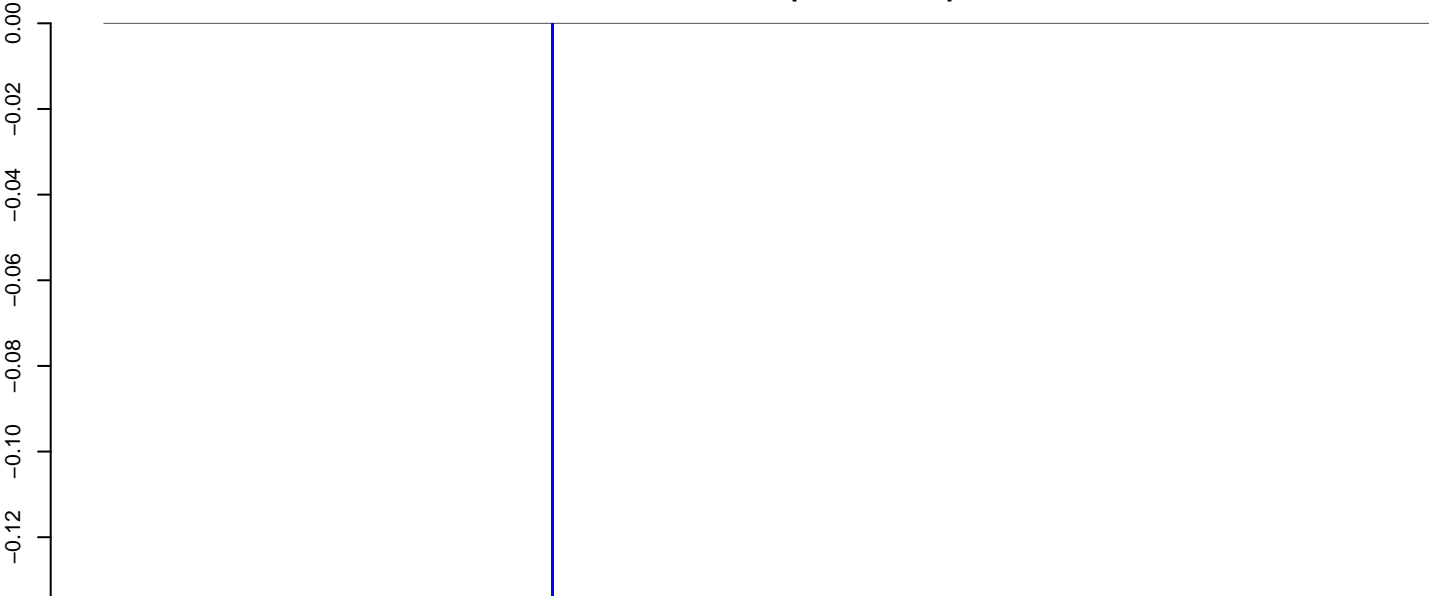
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



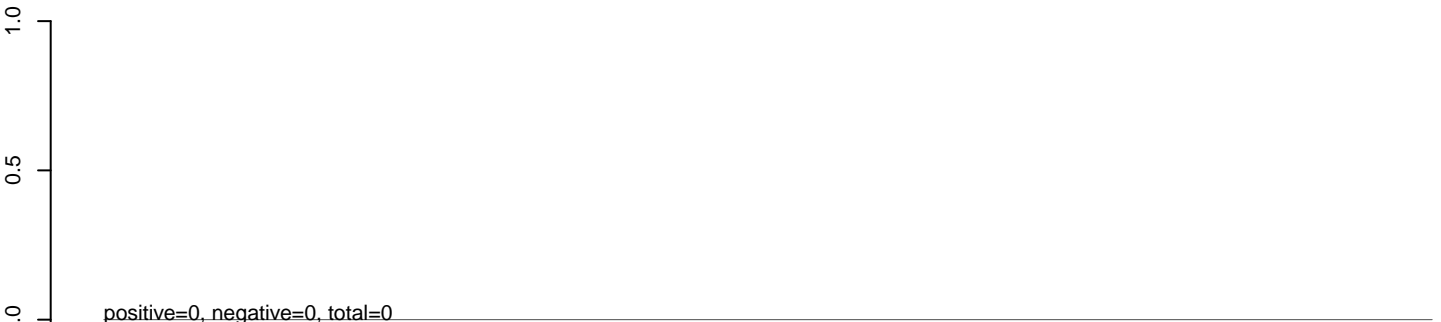
Window size=25, length=11344, virus@AF075256.1-Pixuna_virus_strain_Pixuna_BeAr_35645:1-11344

0 2000 4000 6000 8000 10000

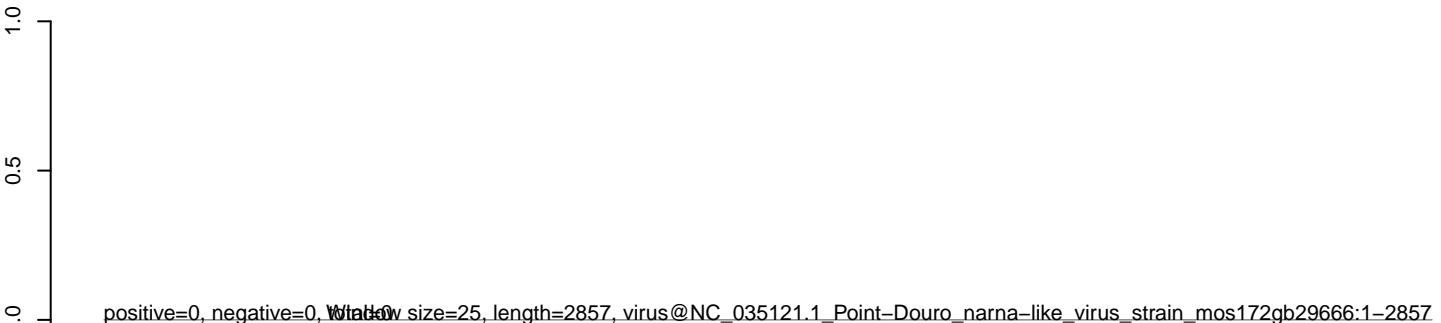
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



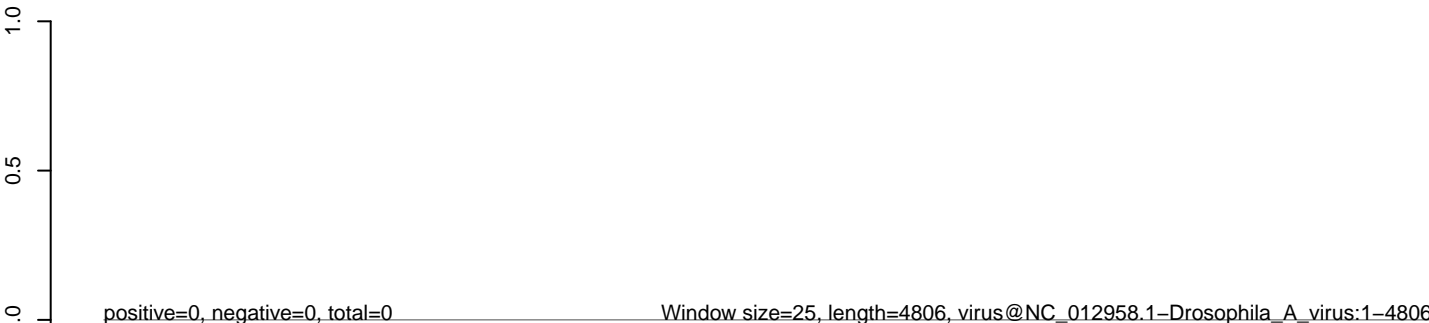
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



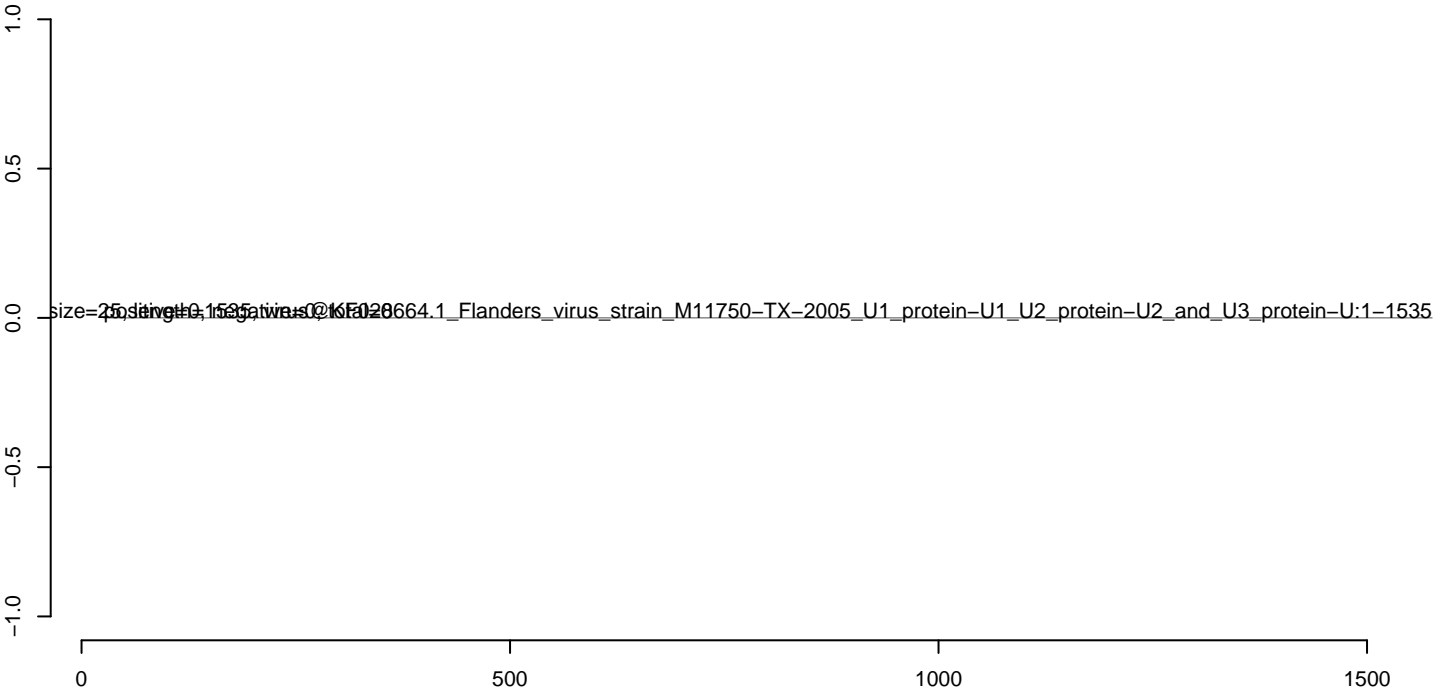
AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep



AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



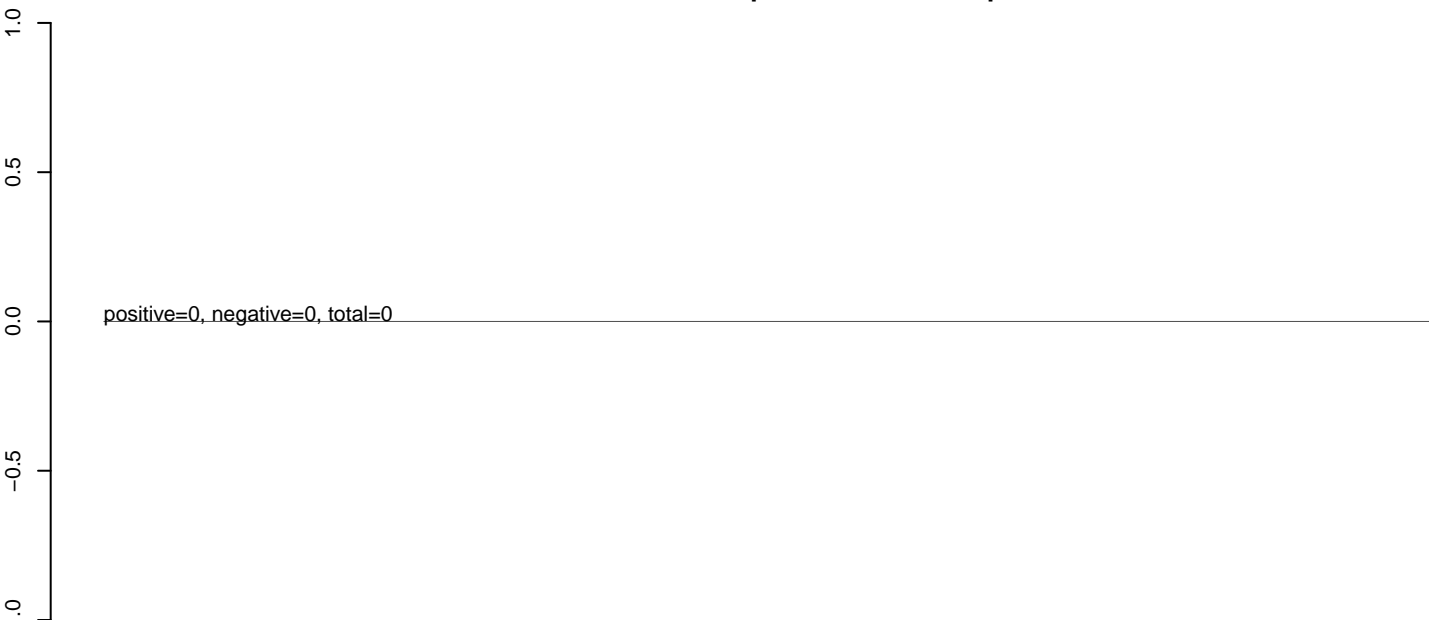
AeAlbo_C636_2dpi_mock3.rep



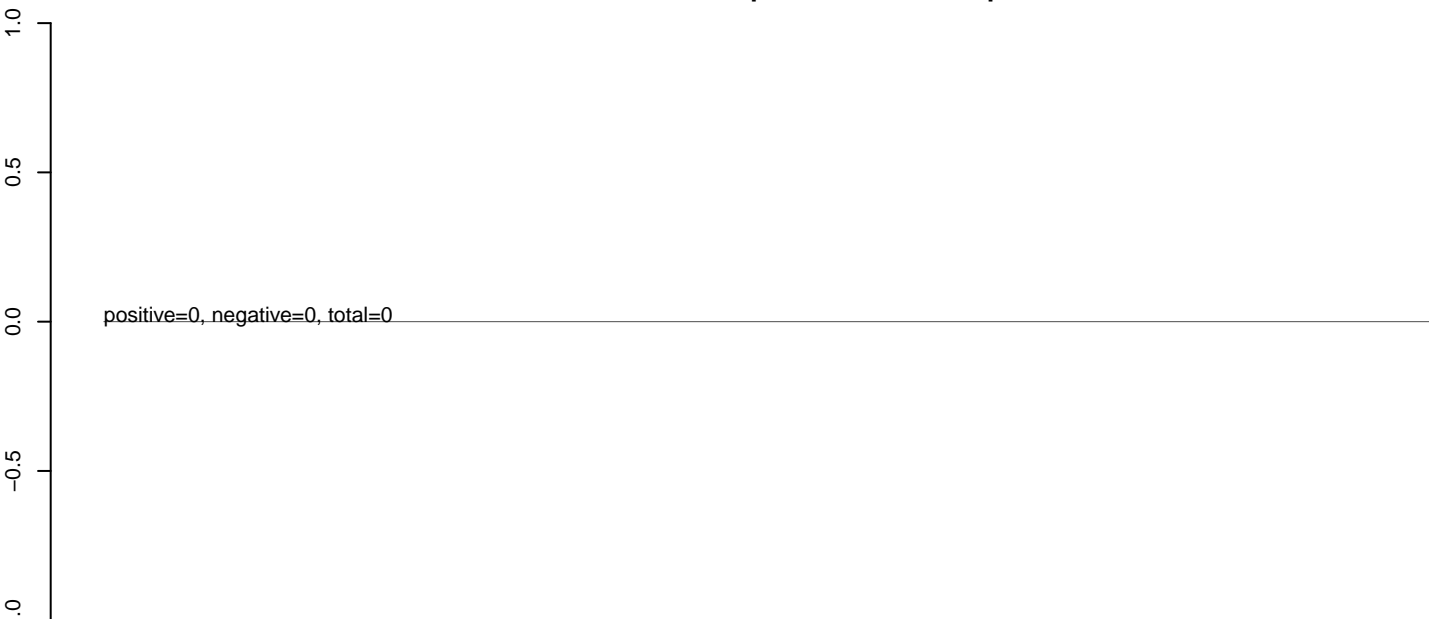
z=25 density=4.523 genome=0.02147965.1-Drosophila_melanogaster_American_nodavirus_ANV_SW-2009a_segment_RNA1_and_RNA2:1-4523

0 1000 2000 3000 4000

AeAlbo_C636_2dpi_mock3.18_23.rep



AeAlbo_C636_2dpi_mock3.24_35.rep



AeAlbo_C636_2dpi_mock3.rep

