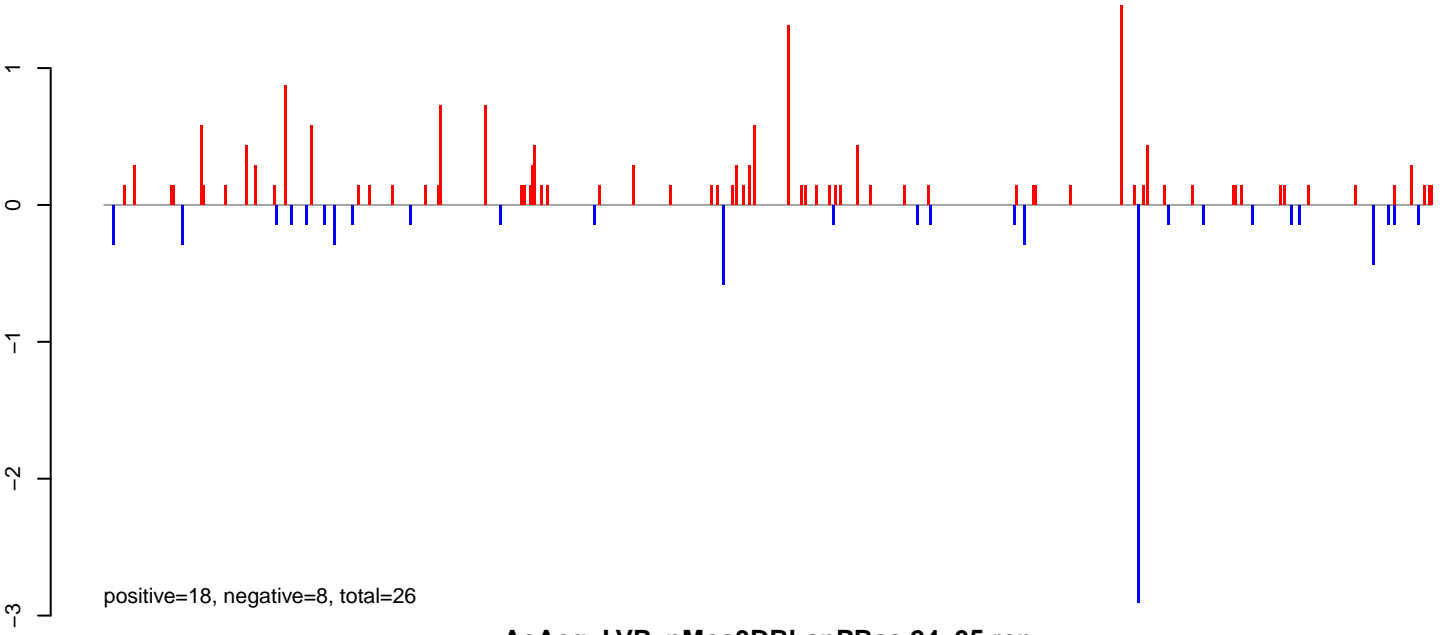
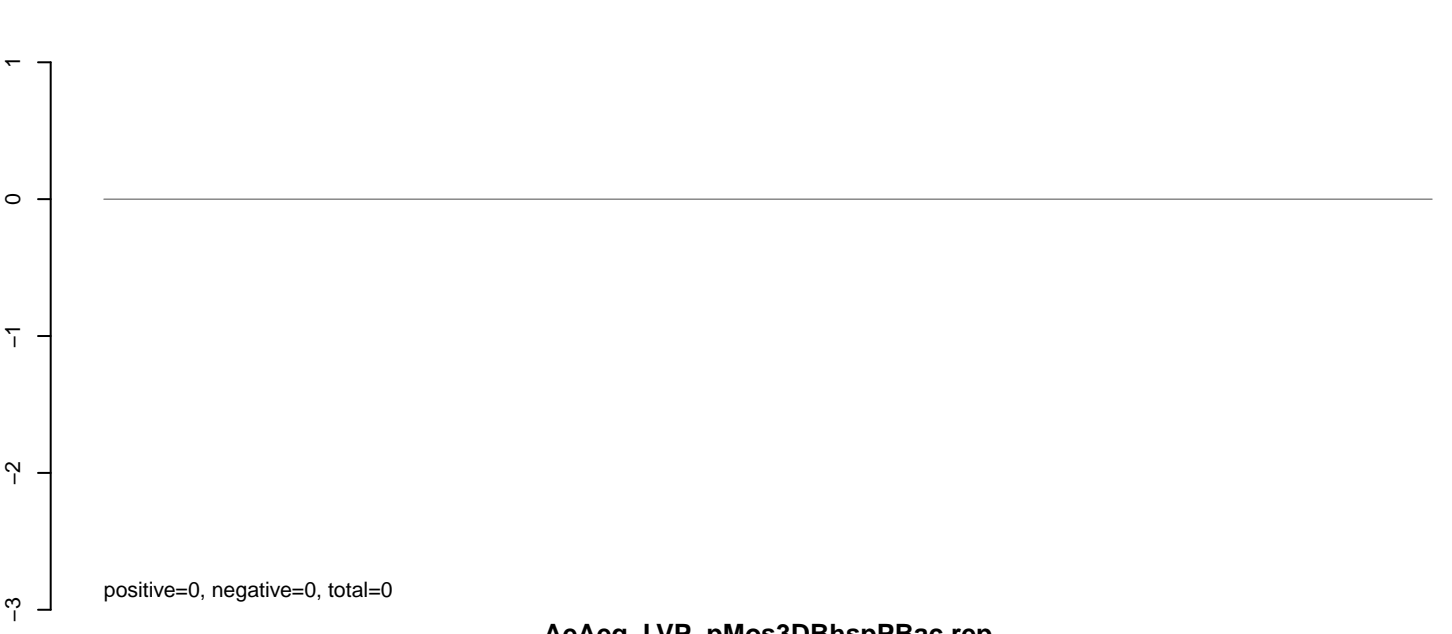


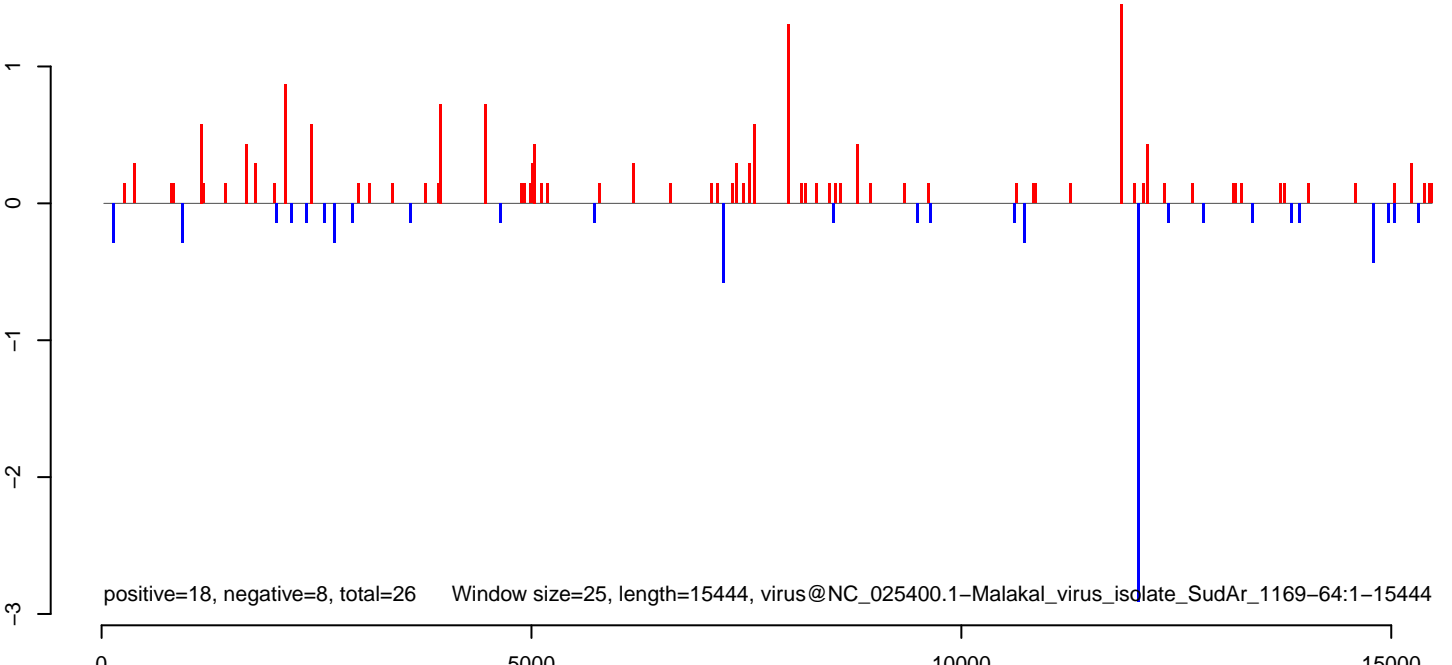
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



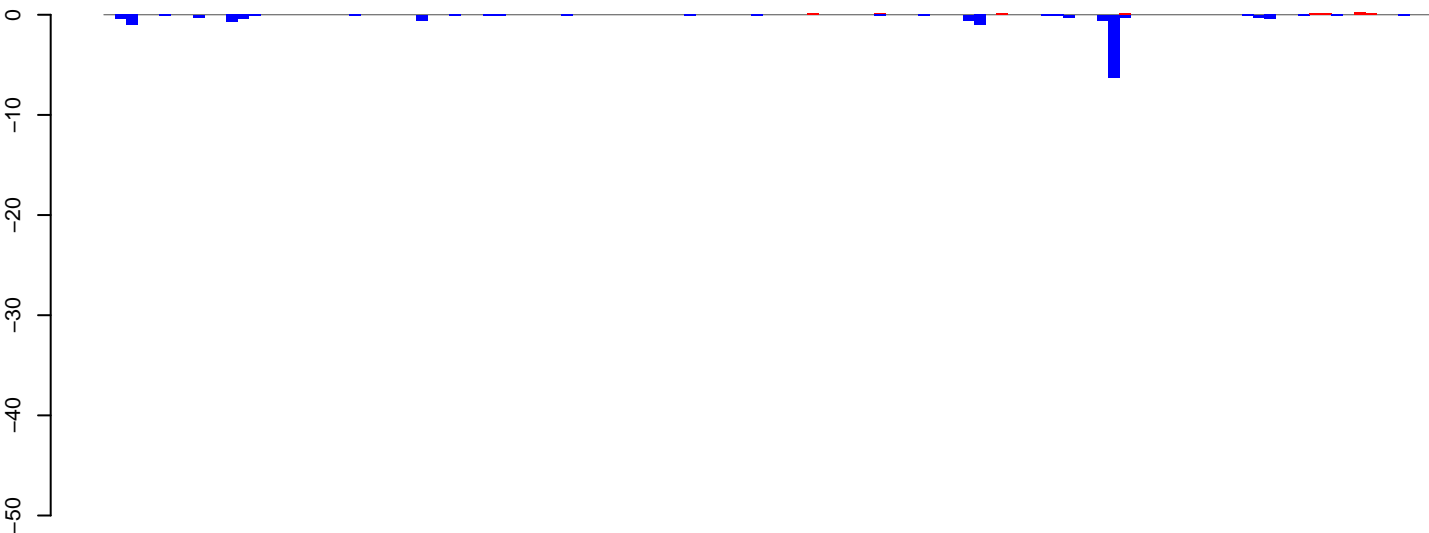
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep

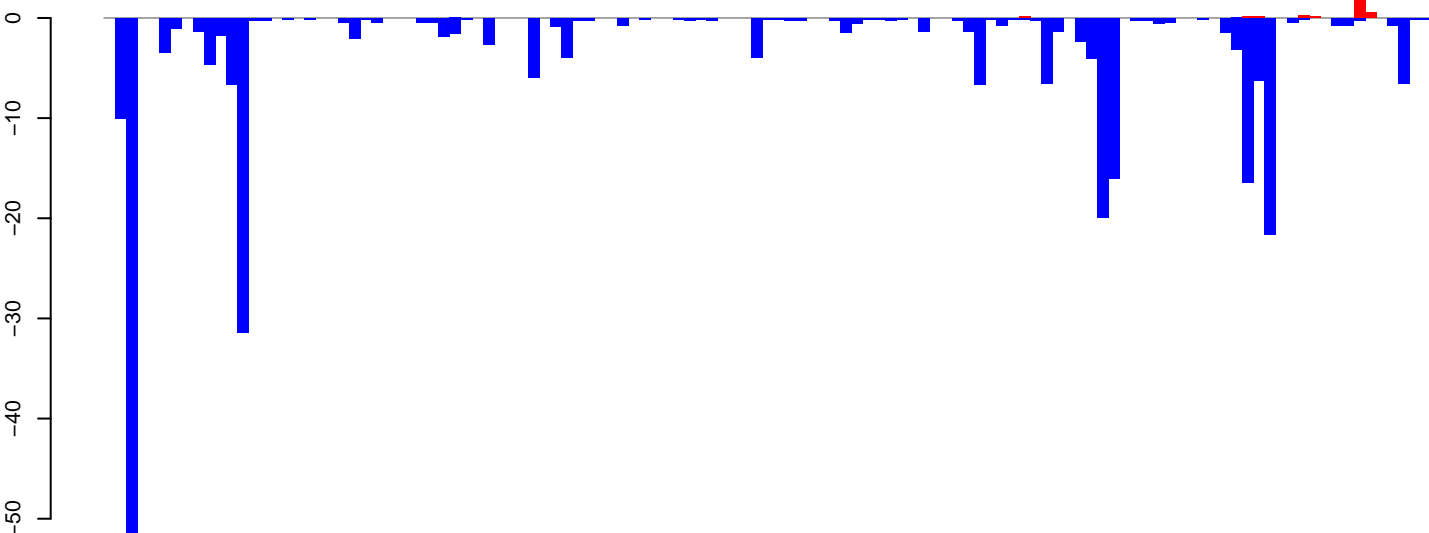


AeAeg_LVP_pMos3DBhspPBac.18_23.rep



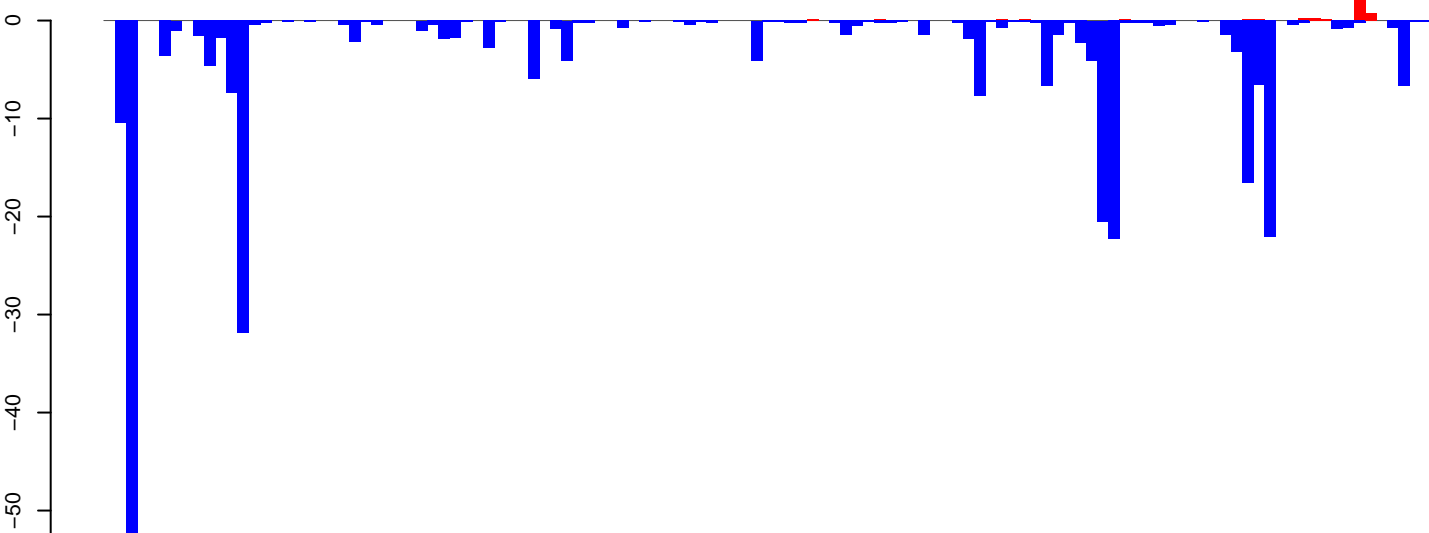
positive=1, negative=16, total=17

AeAeg_LVP_pMos3DBhspPBac.24_35.rep



positive=4, negative=272, total=276

AeAeg_LVP_pMos3DBhspPBac.rep

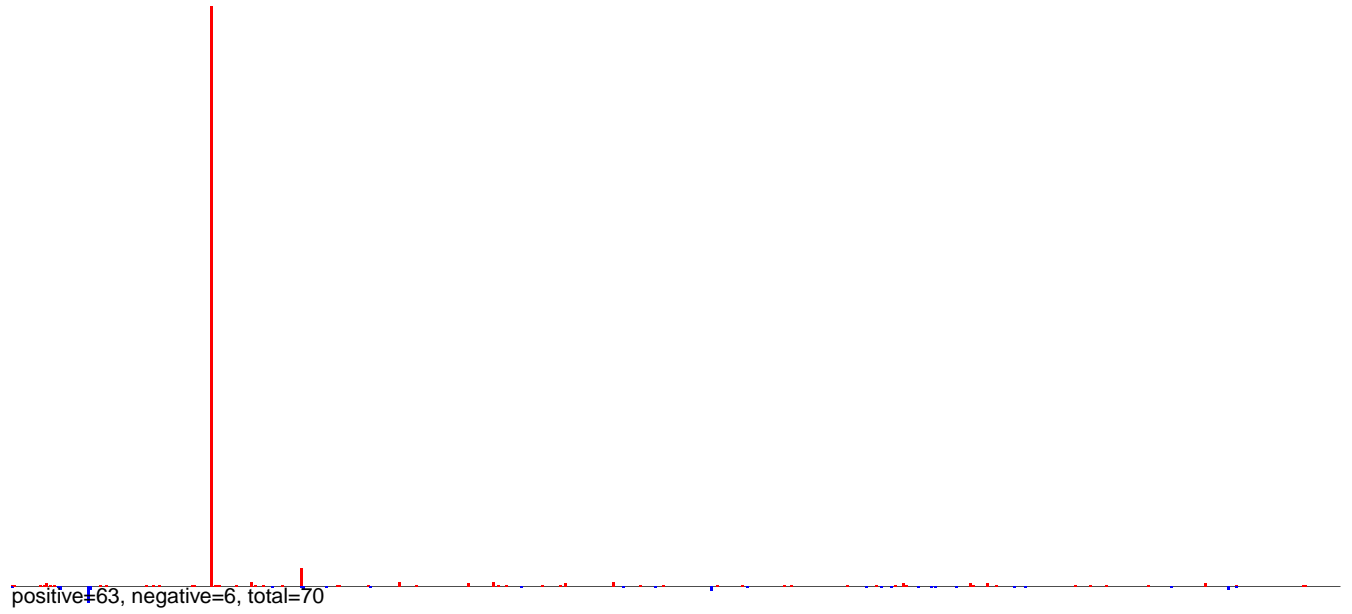


positive=5, negative=287, total=292

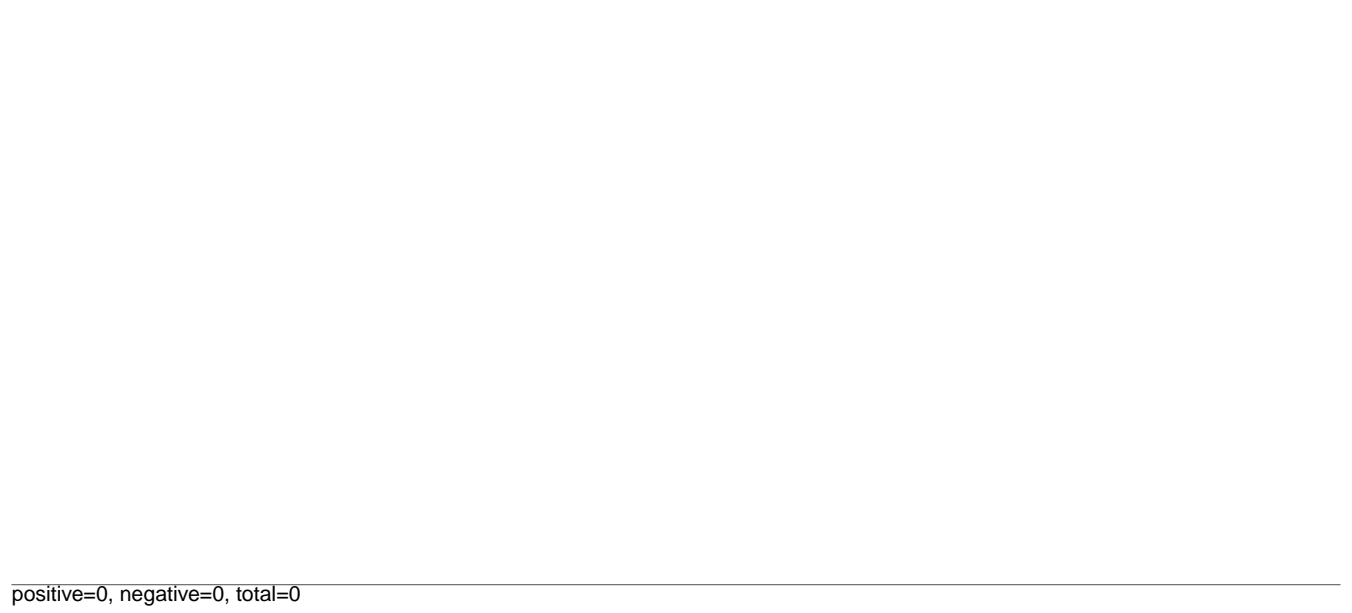
Window size=25, length=2965, virus@AY347953.2-Aedes_aegypti_genomic_sequence:1-2965

0 500 1000 1500 2000 2500 3000

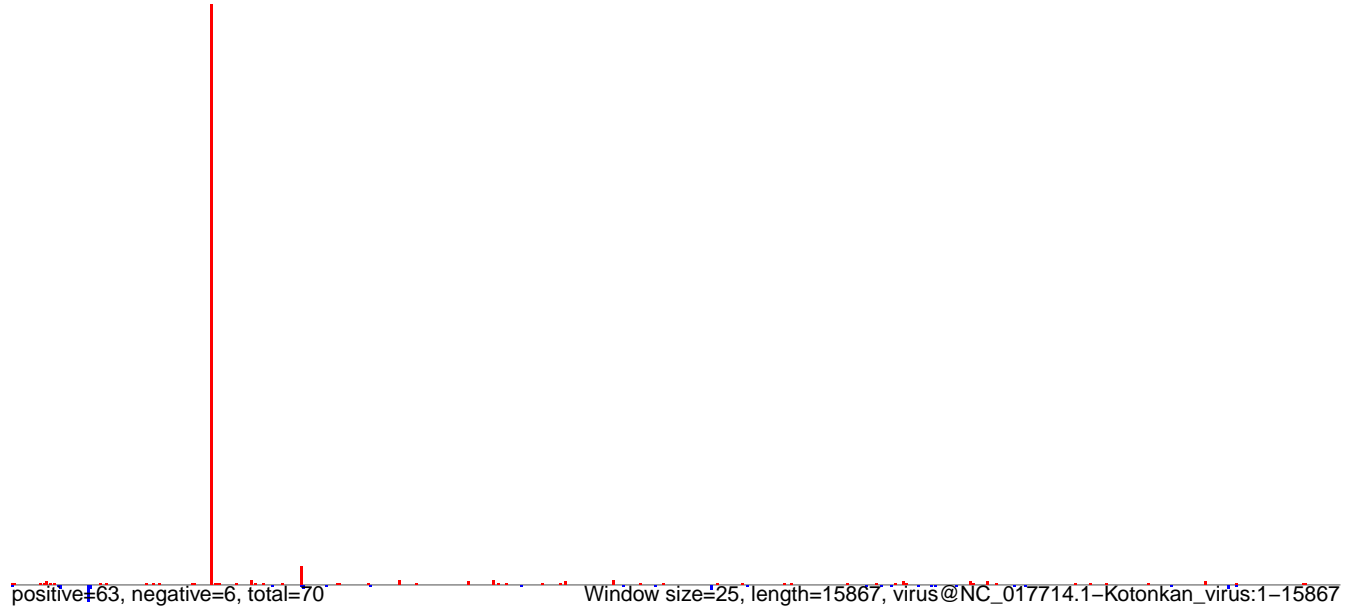
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

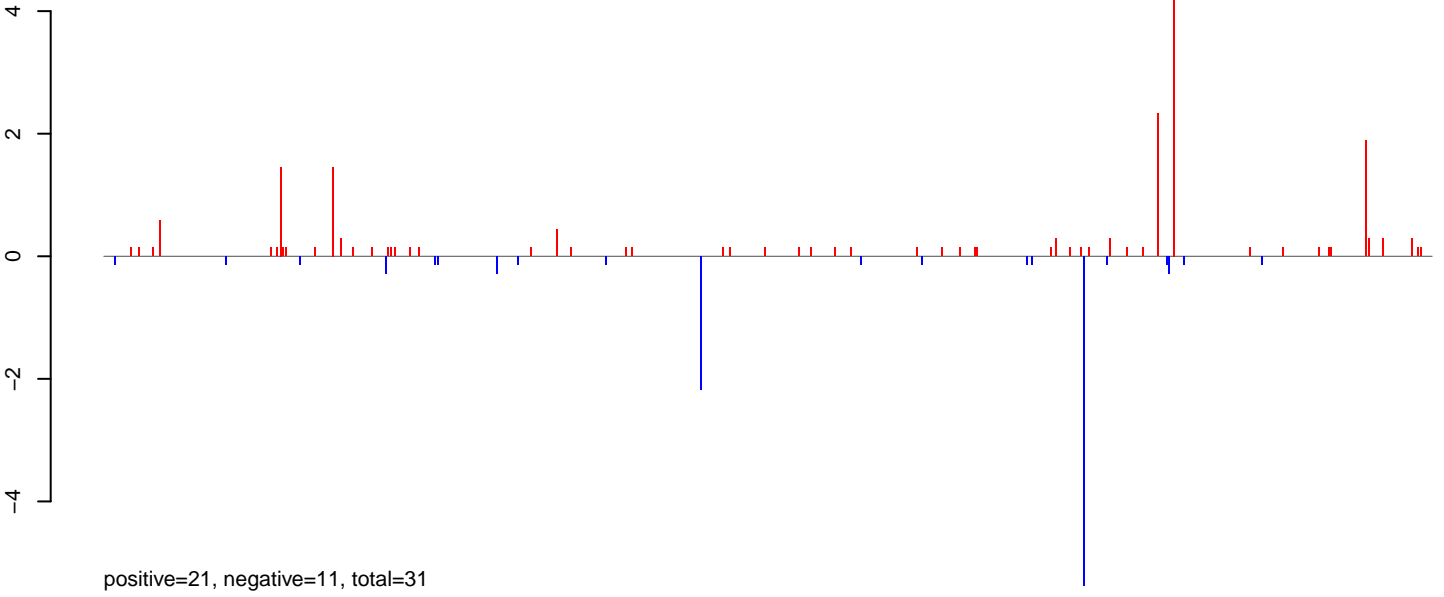


AeAeg_LVP_pMos3DBhspPBac.rep

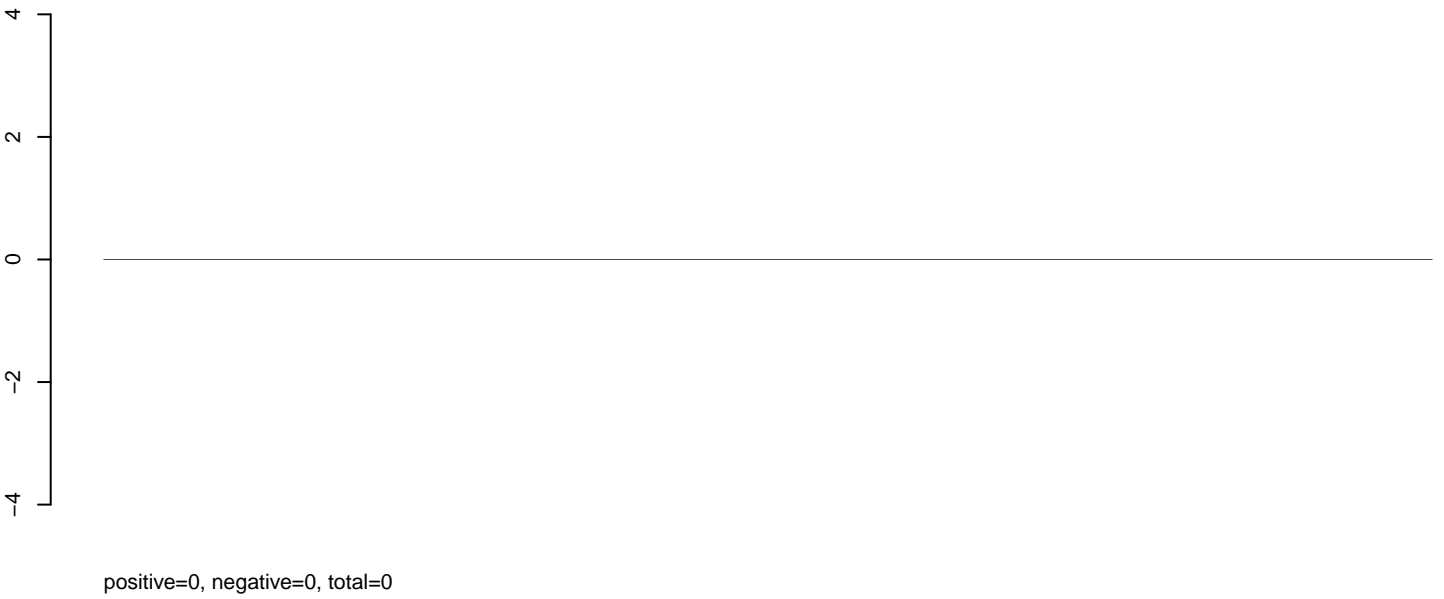


0 5000 10000 15000

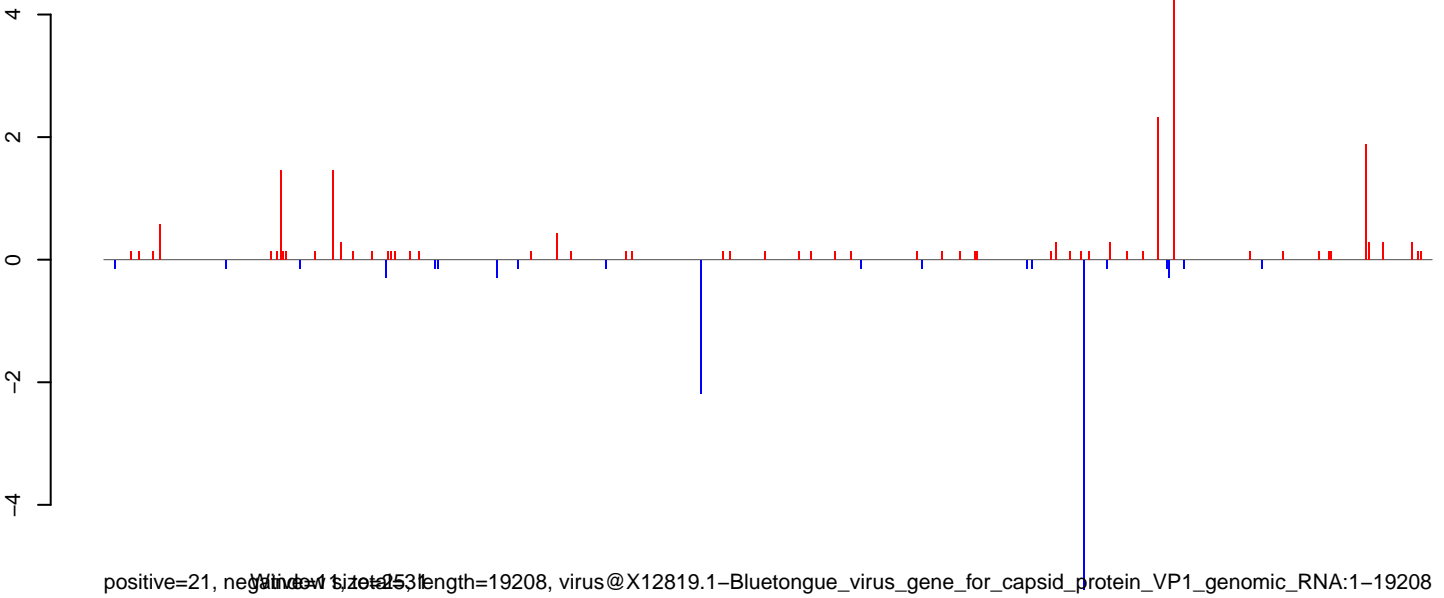
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



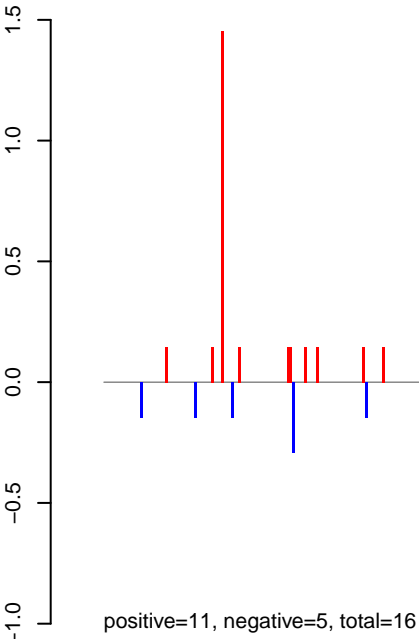
AeAeg_LVP_pMos3DBhspPBac.rep



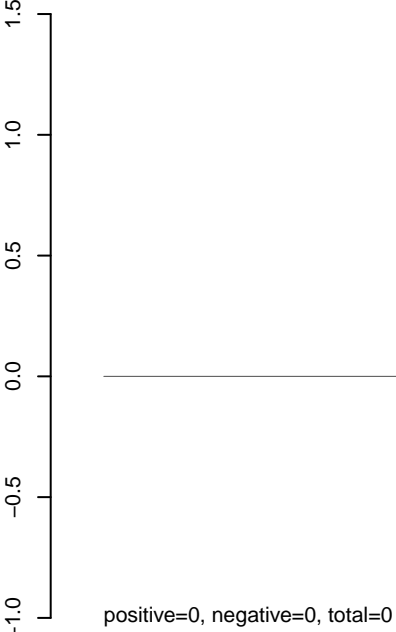
Window size=253 length=19208, virus@X12819.1-Bluetongue_virus_gene_for_capsid_protein_VP1_genomic_RNA:1-19208

0 5000 10000 15000 20000

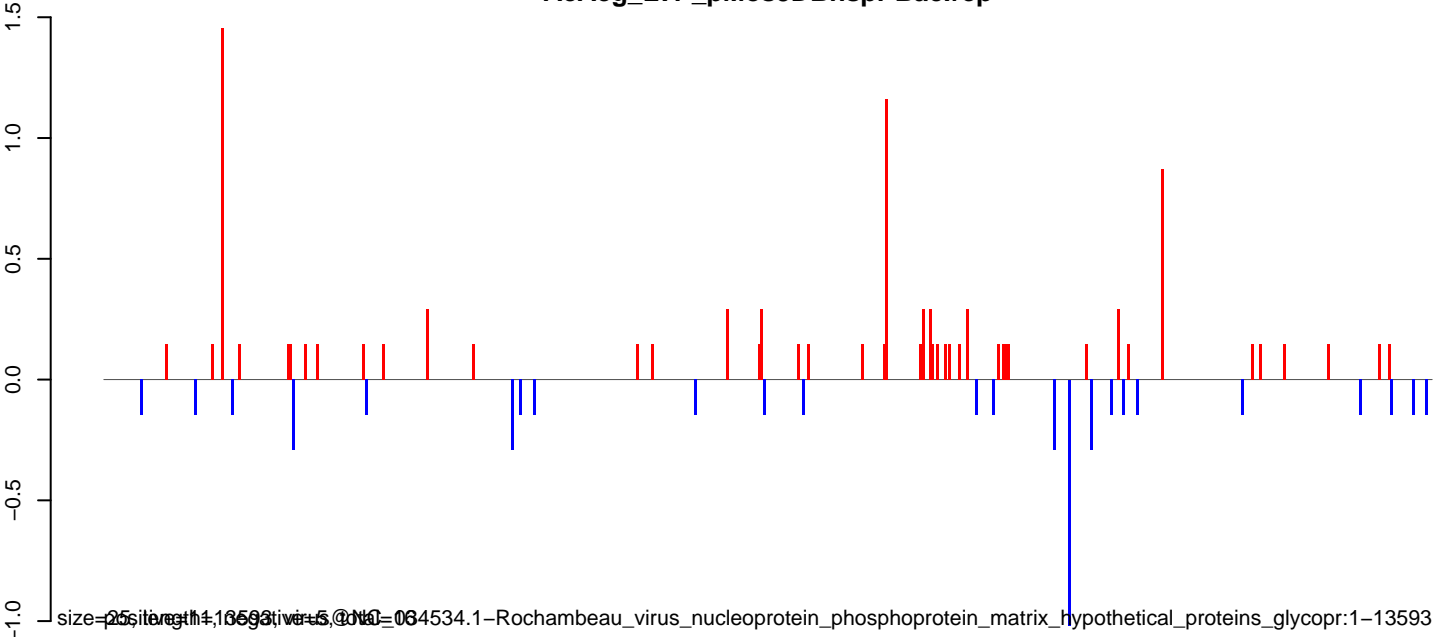
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



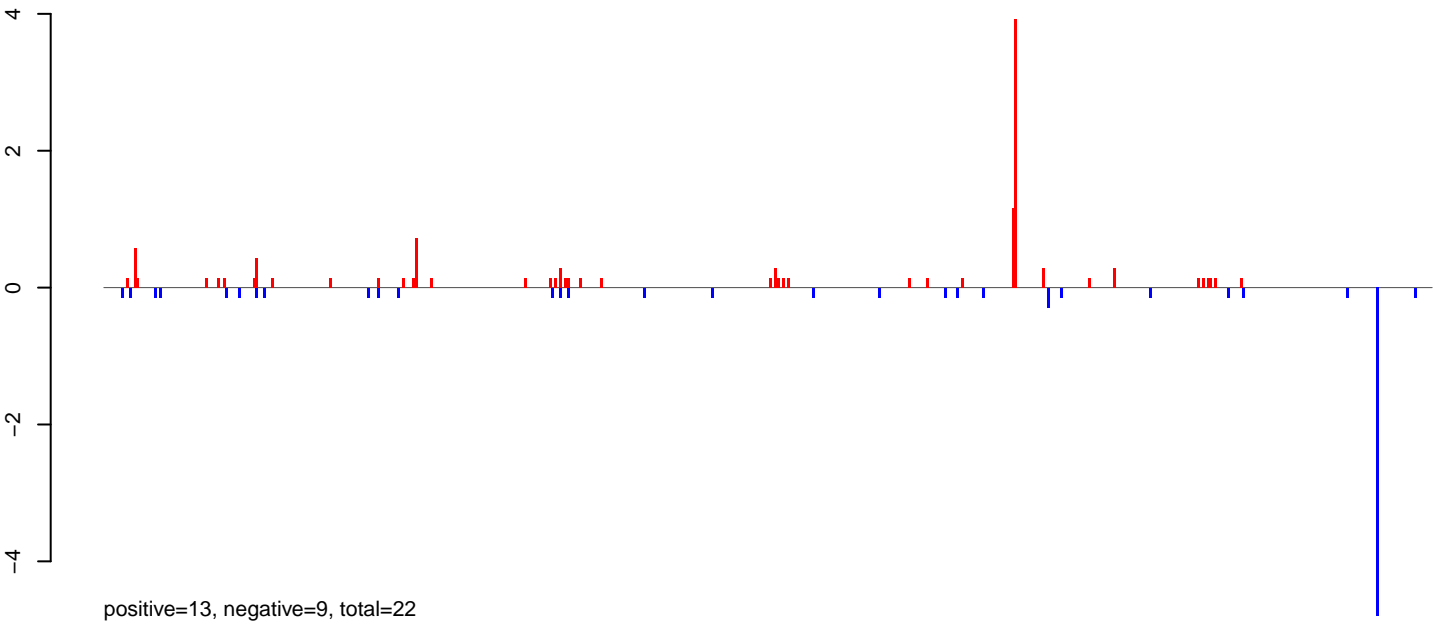
AeAeg_LVP_pMos3DBhspPBac.rep



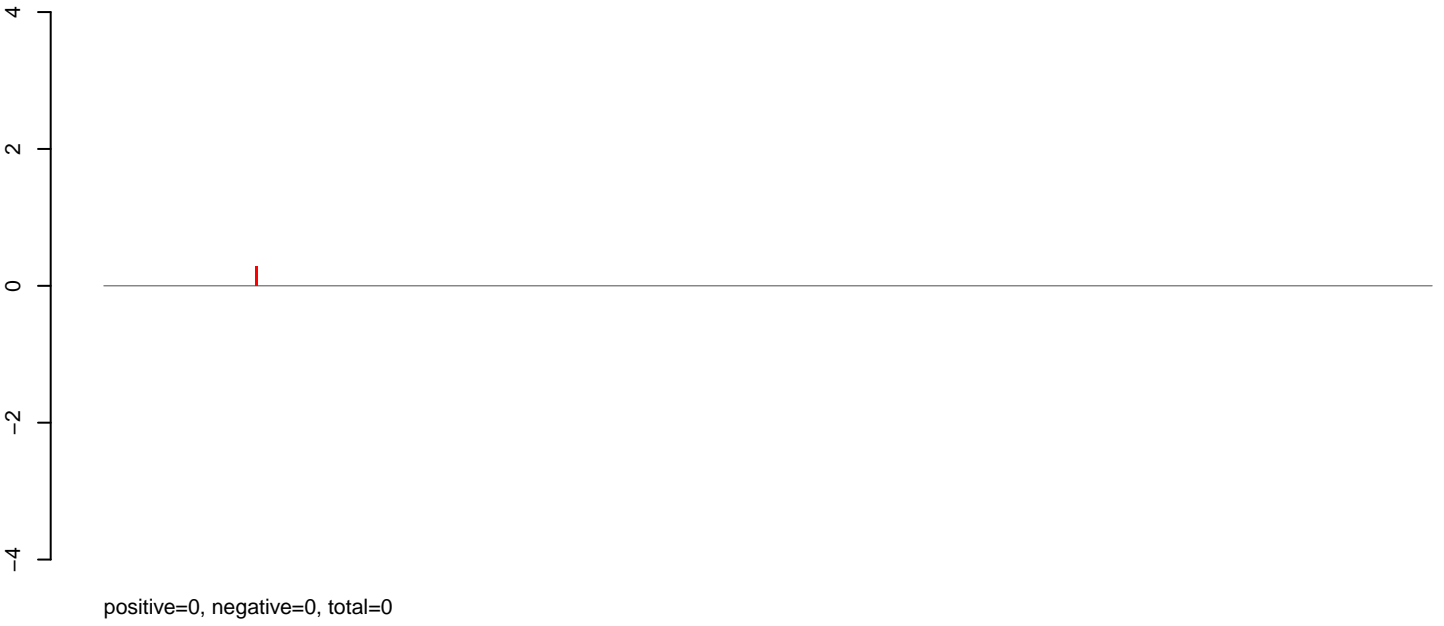
064534.1-Rochambeau_virus_nucleoprotein_phosphoprotein_matrix_hypothetical_proteins_glycopr:1-13593

0 2000 4000 6000 8000 10000 12000 14000

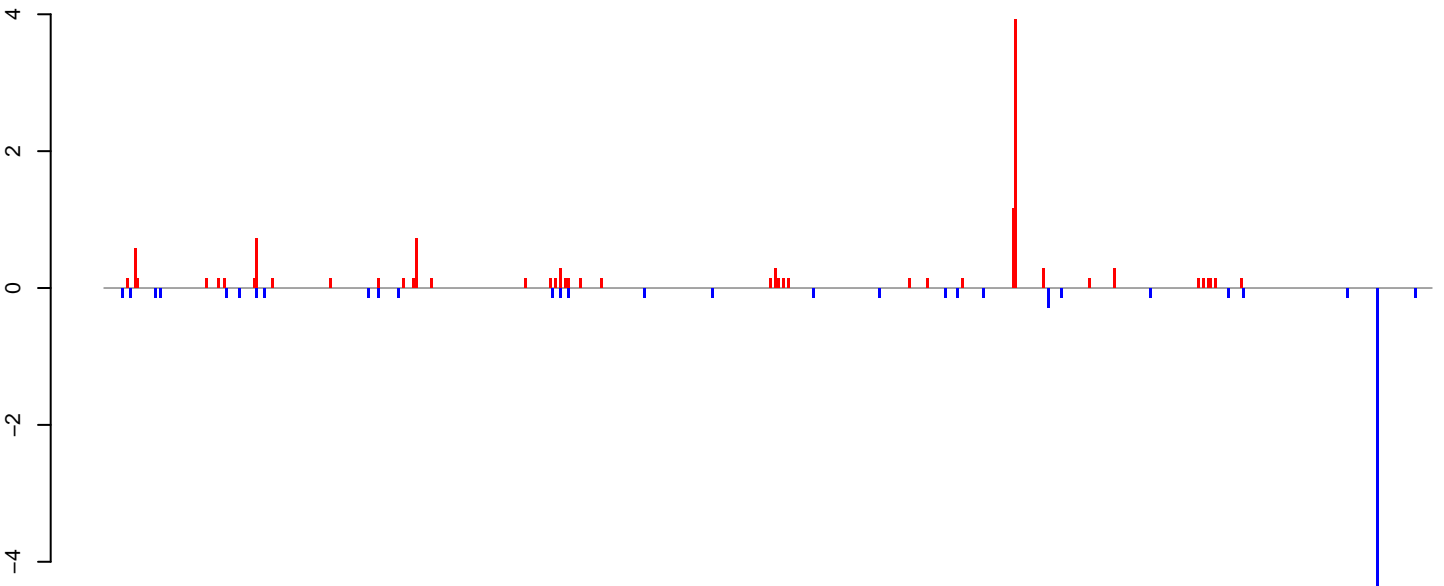
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



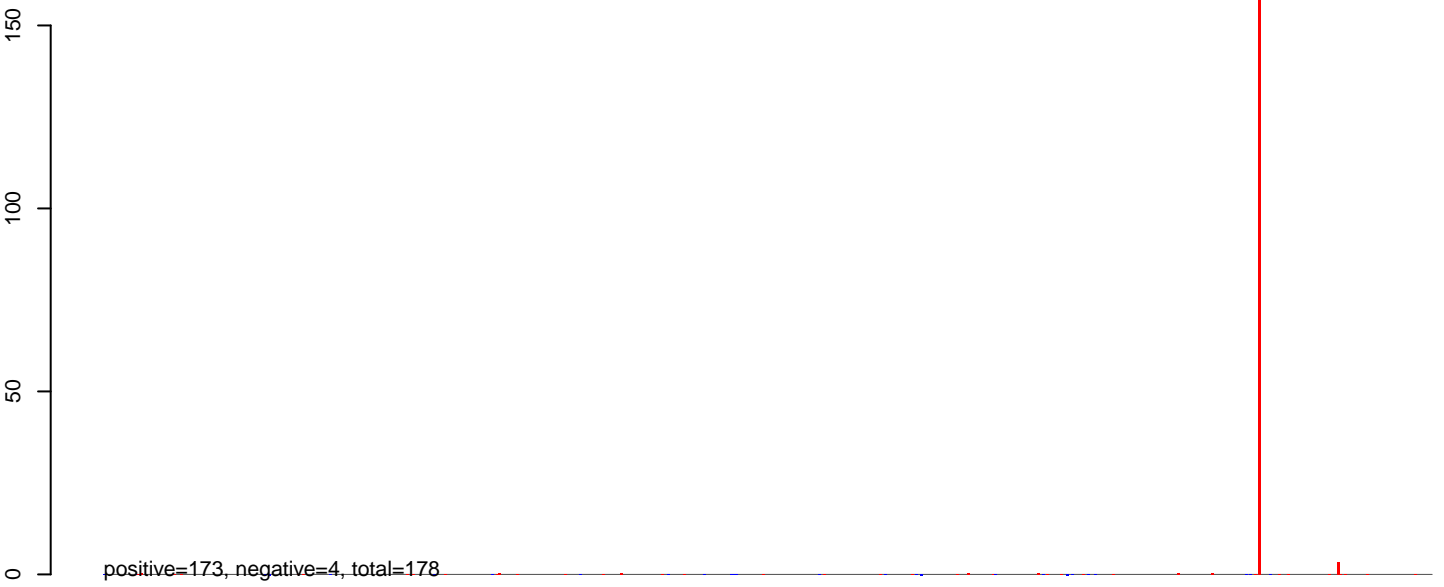
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



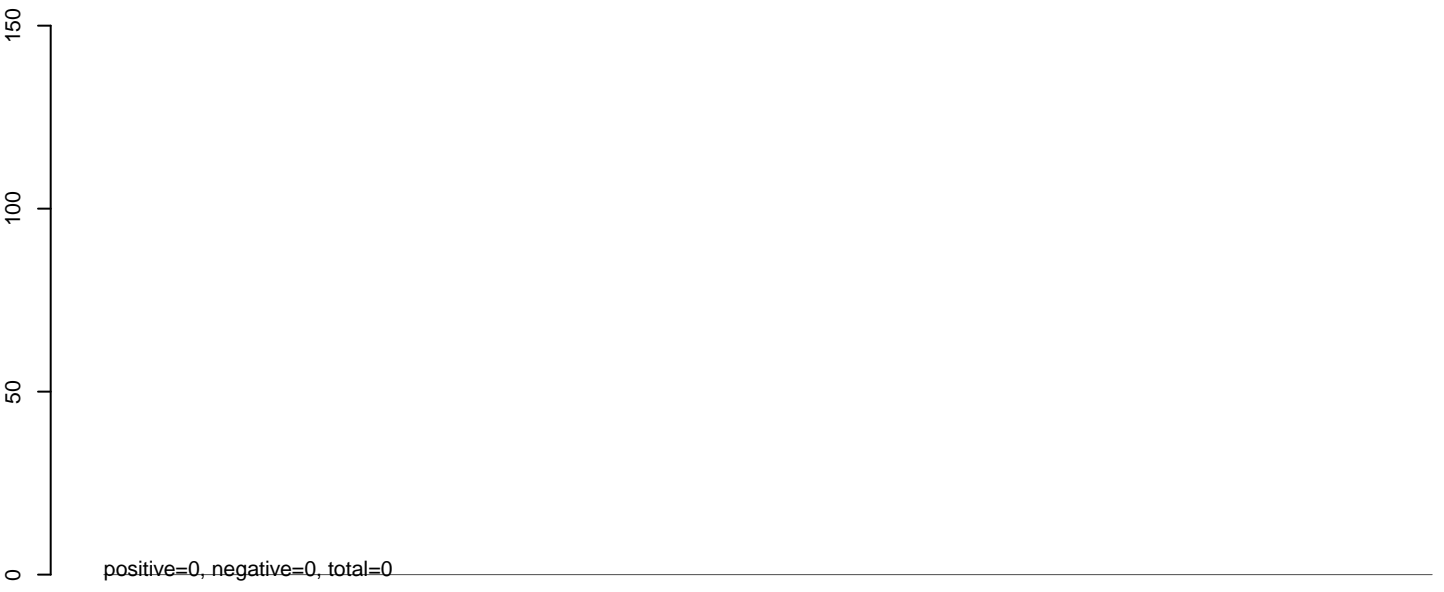
AeAeg_LVP_pMos3DBhspPBac.rep



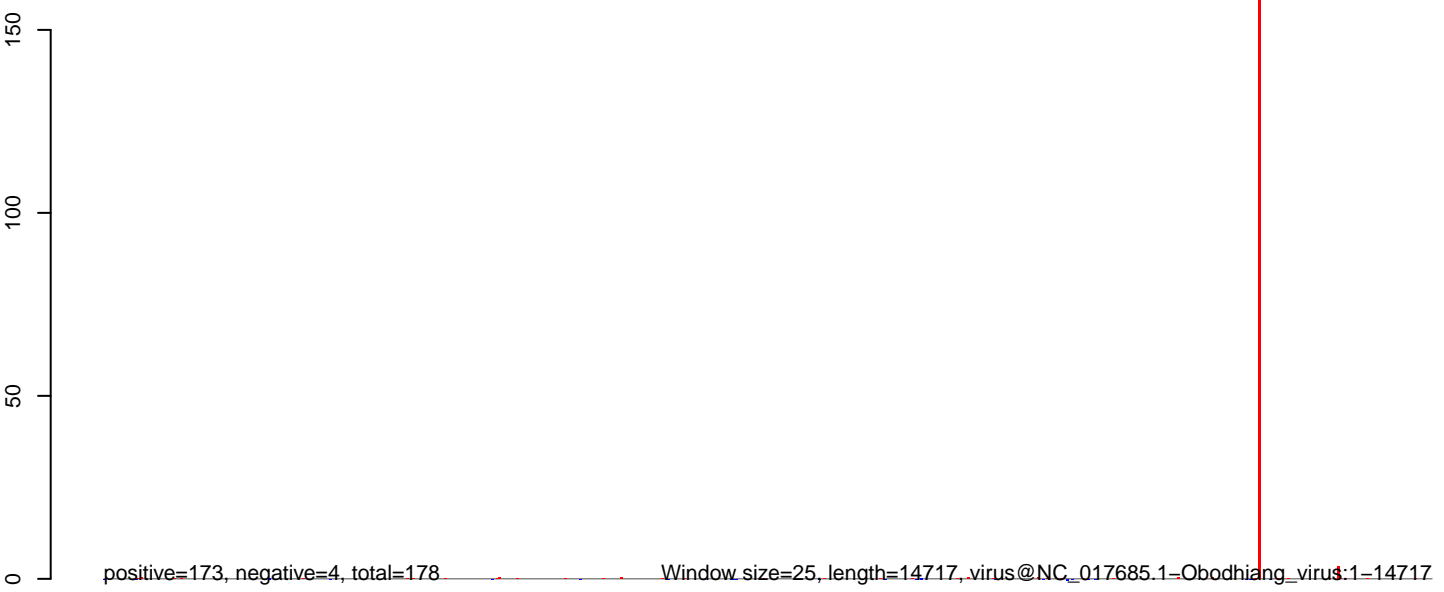
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



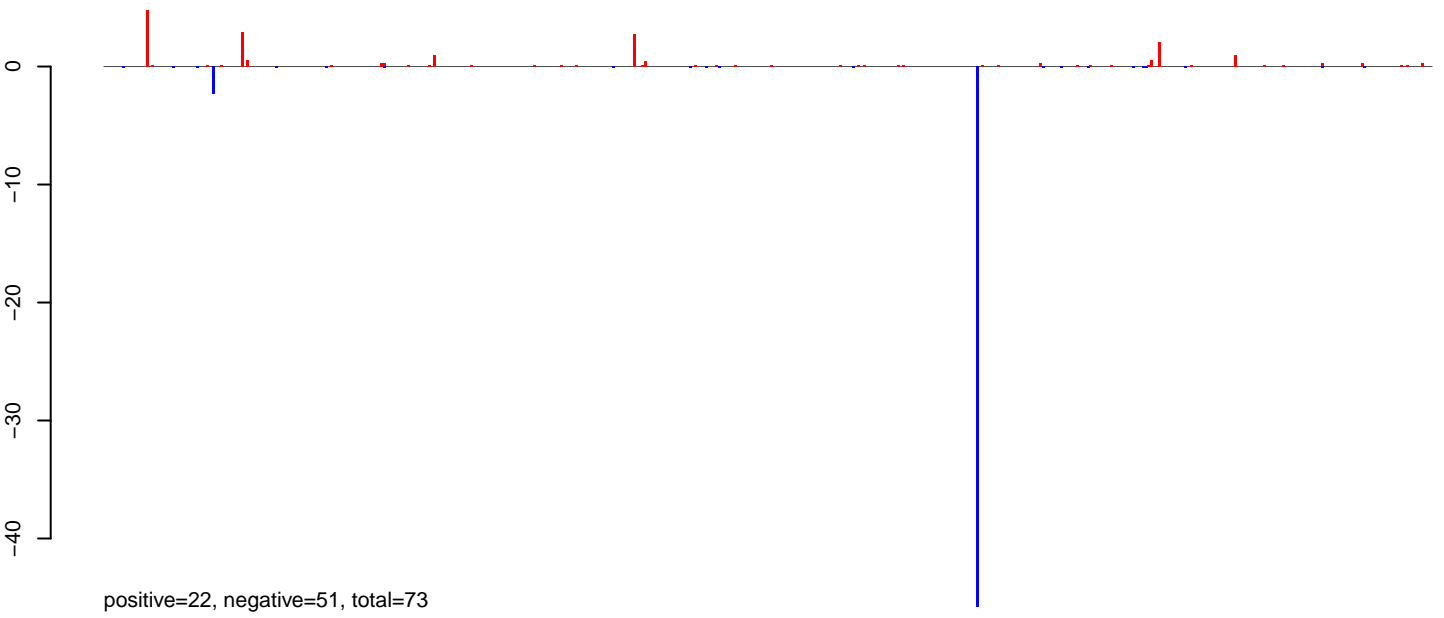
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



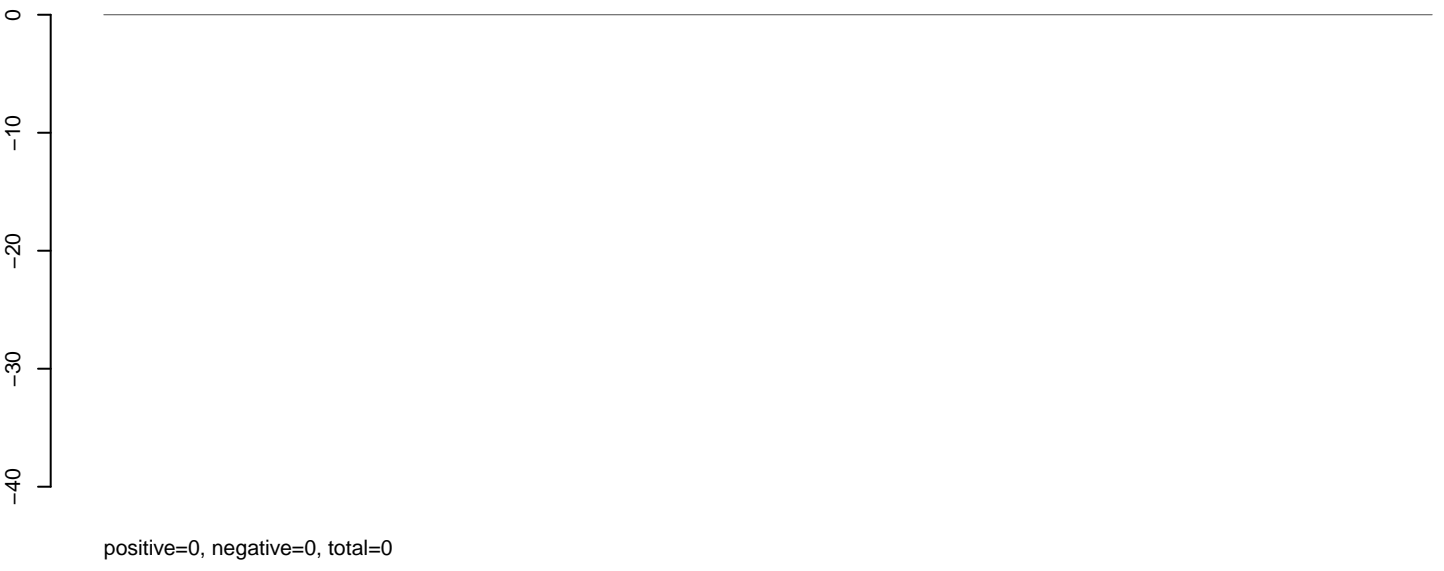
AeAeg_LVP_pMos3DBhspPBac.rep



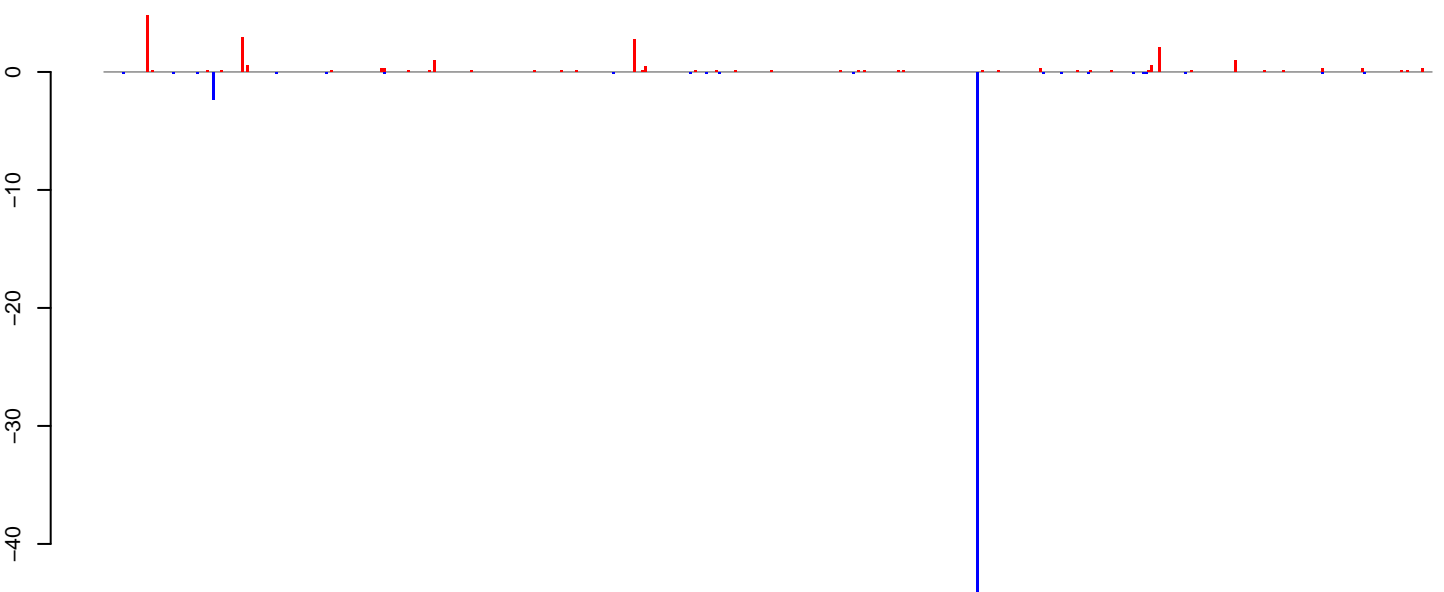
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



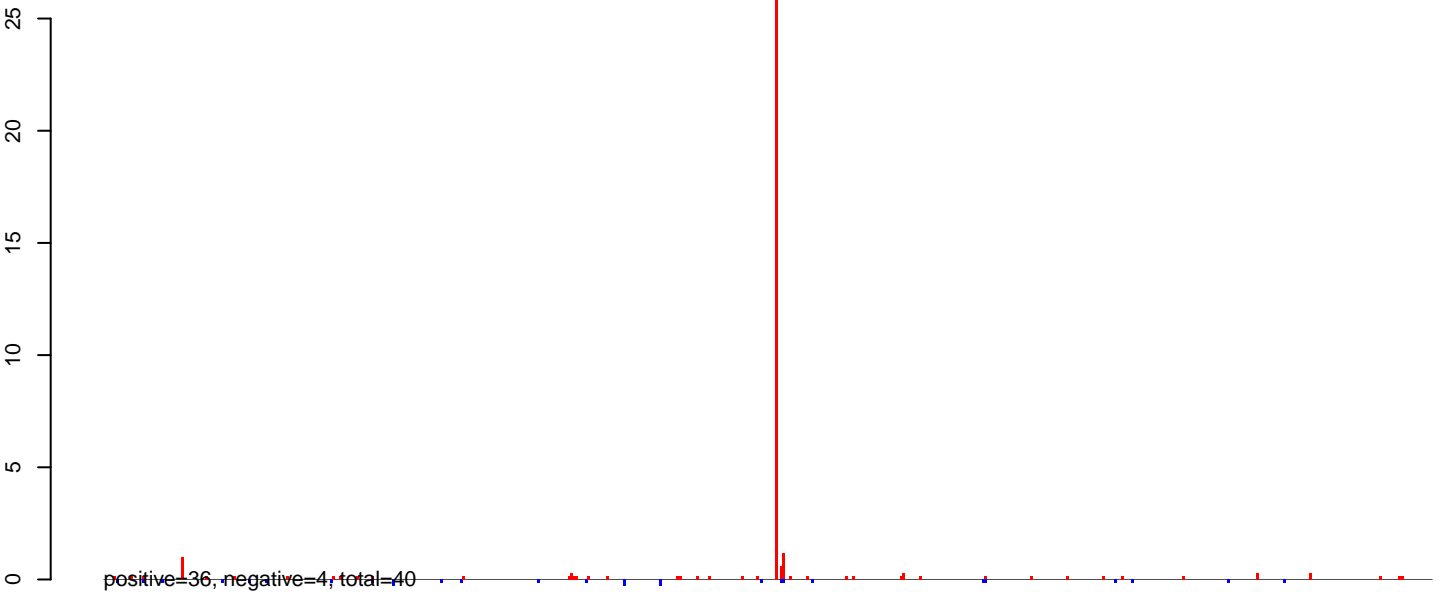
AeAeg_LVP_pMos3DBhspPBac.rep



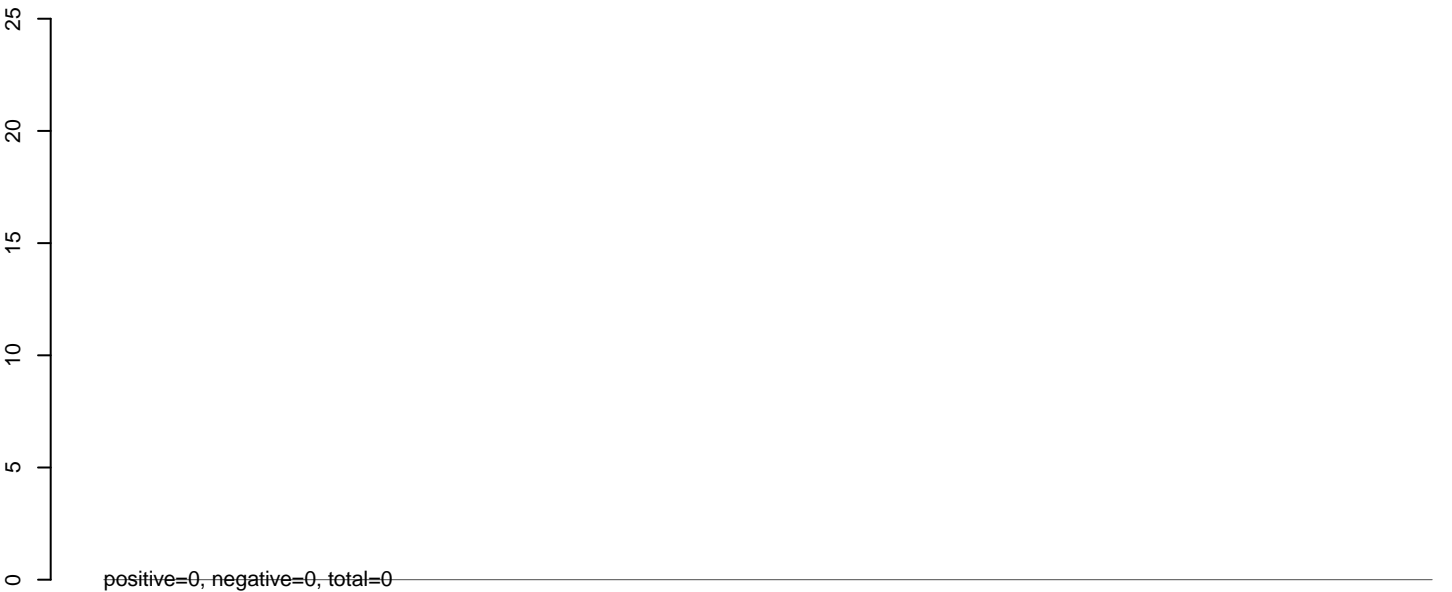
size=25, length=12580, positive=22, negative=51, total=73

0 2000 4000 6000 8000 10000 12000

AeAeg_LVP_pMos3DBhspPBac.18_23.rep



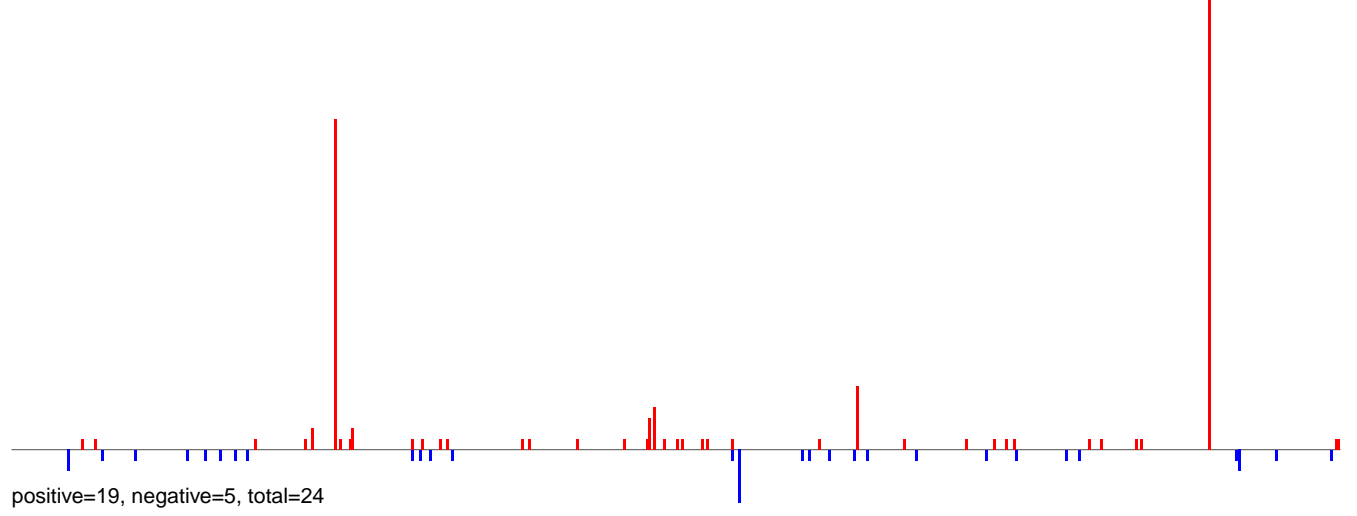
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



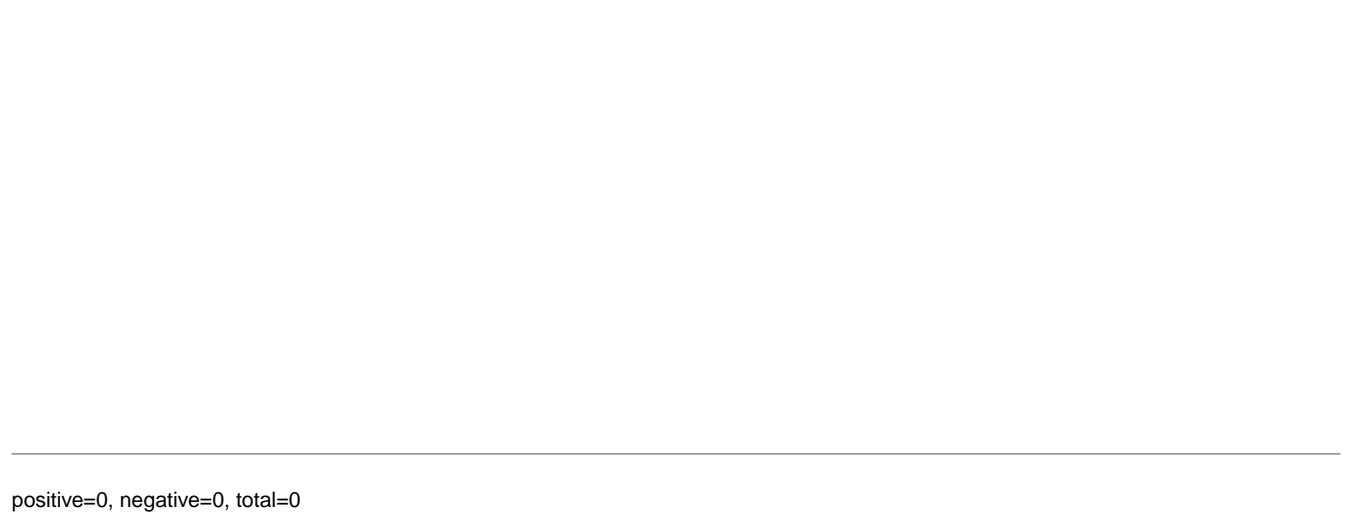
AeAeg_LVP_pMos3DBhspPBac.rep



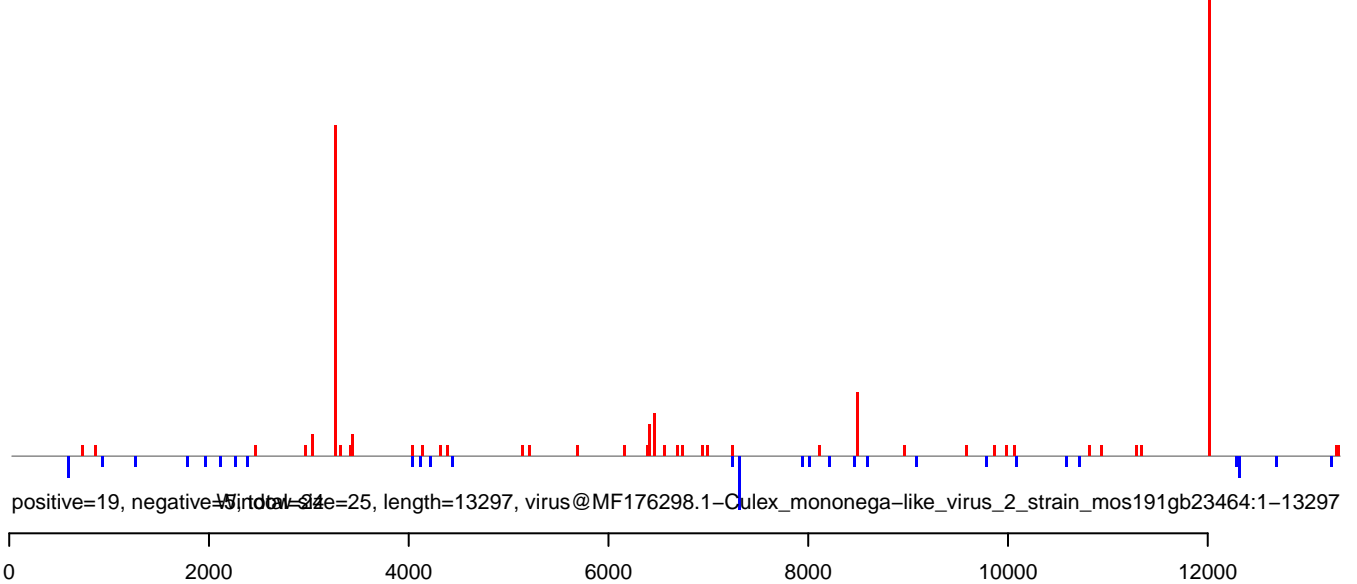
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



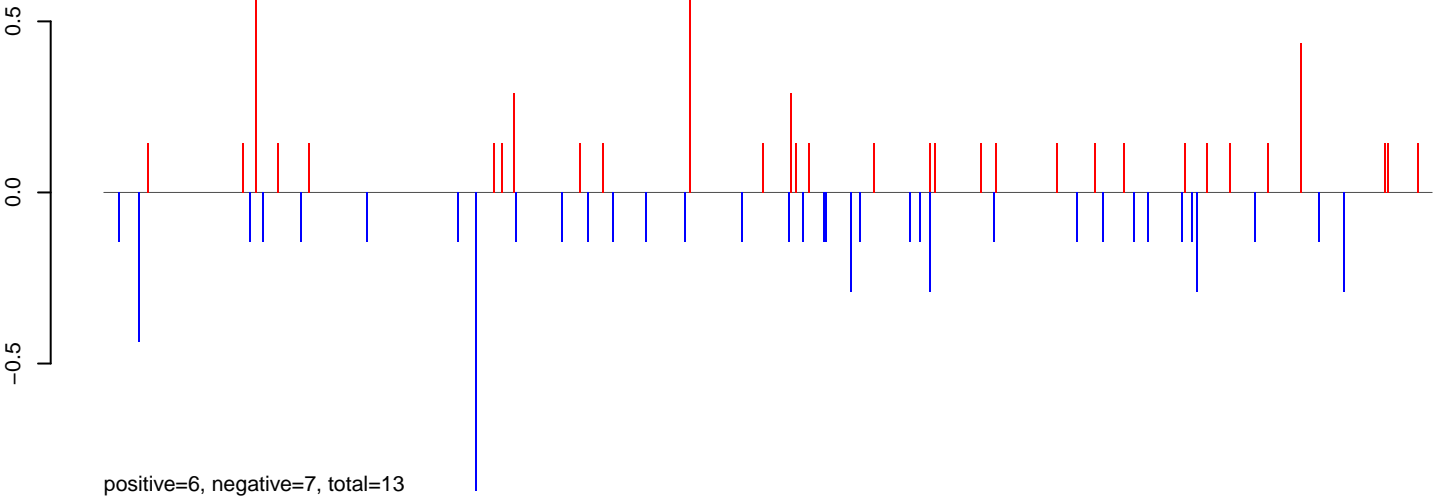
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



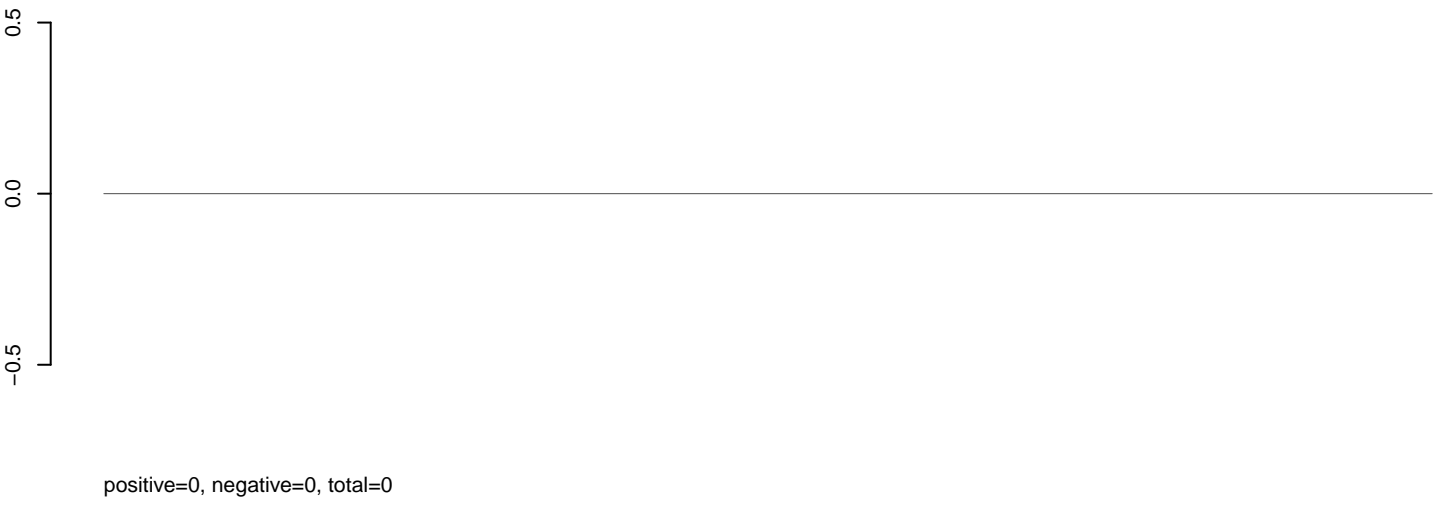
AeAeg_LVP_pMos3DBhspPBac.rep



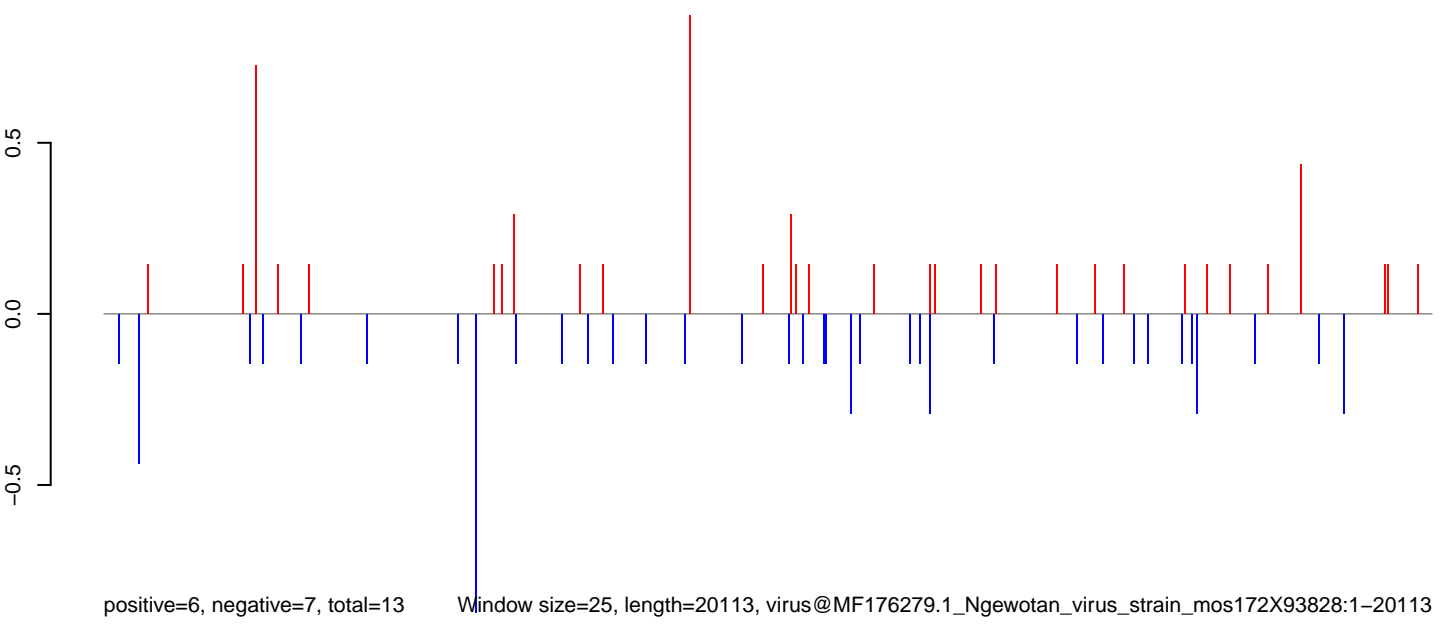
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

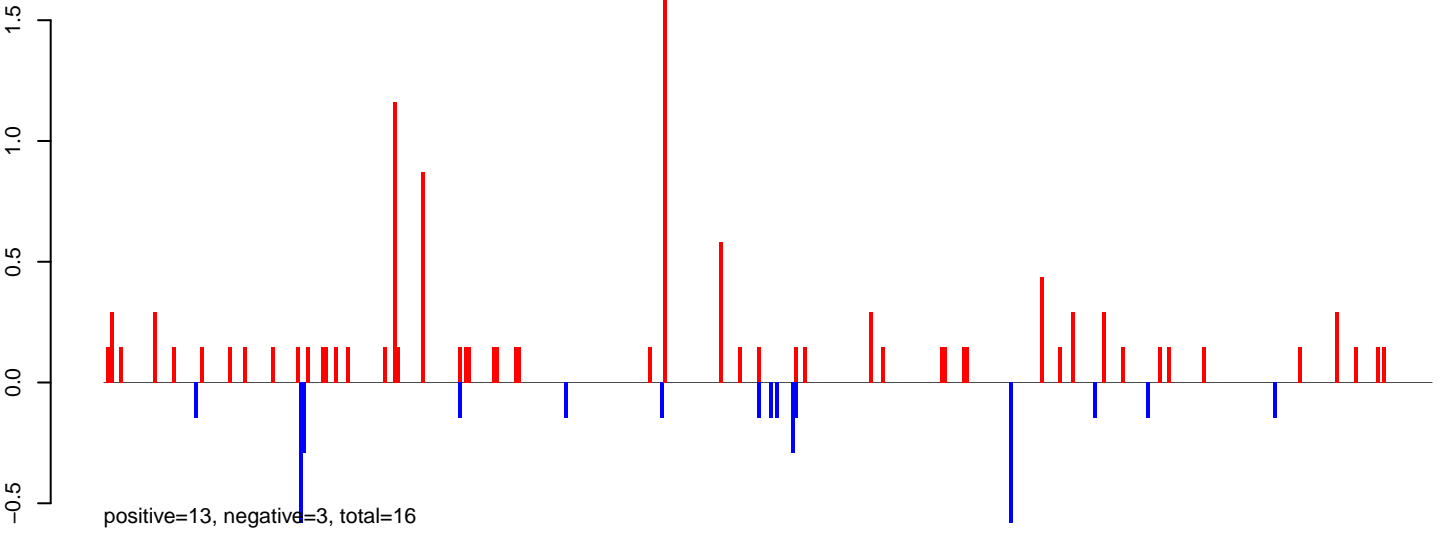


AeAeg_LVP_pMos3DBhspPBac.rep

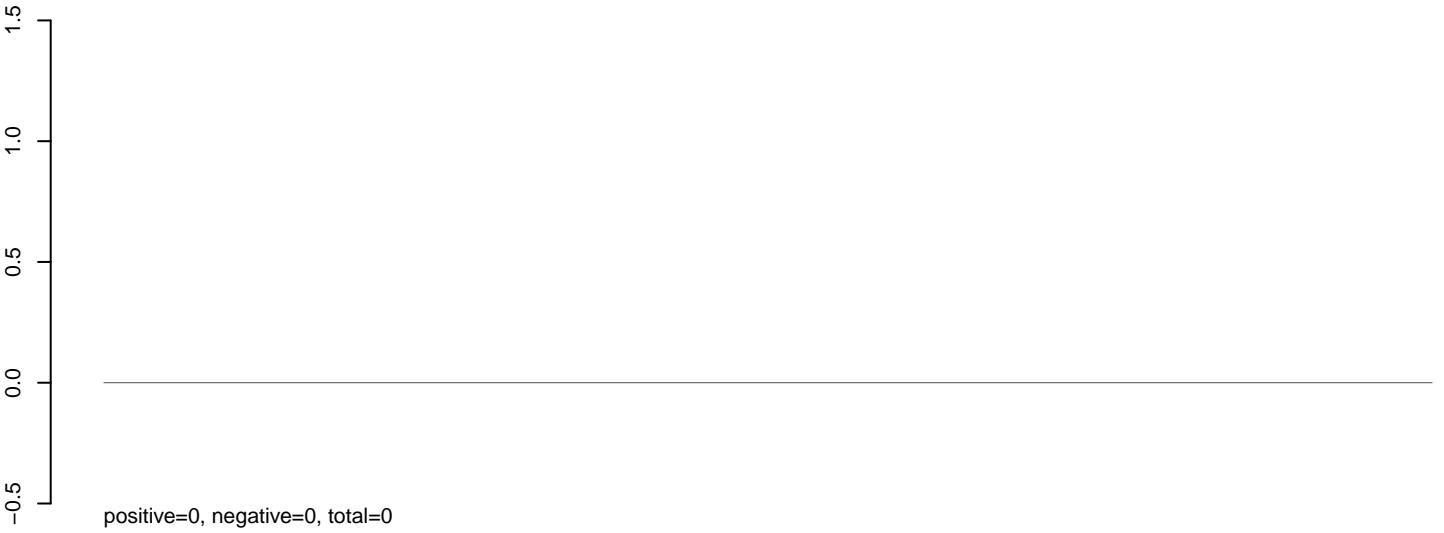


0 5000 10000 15000 20000

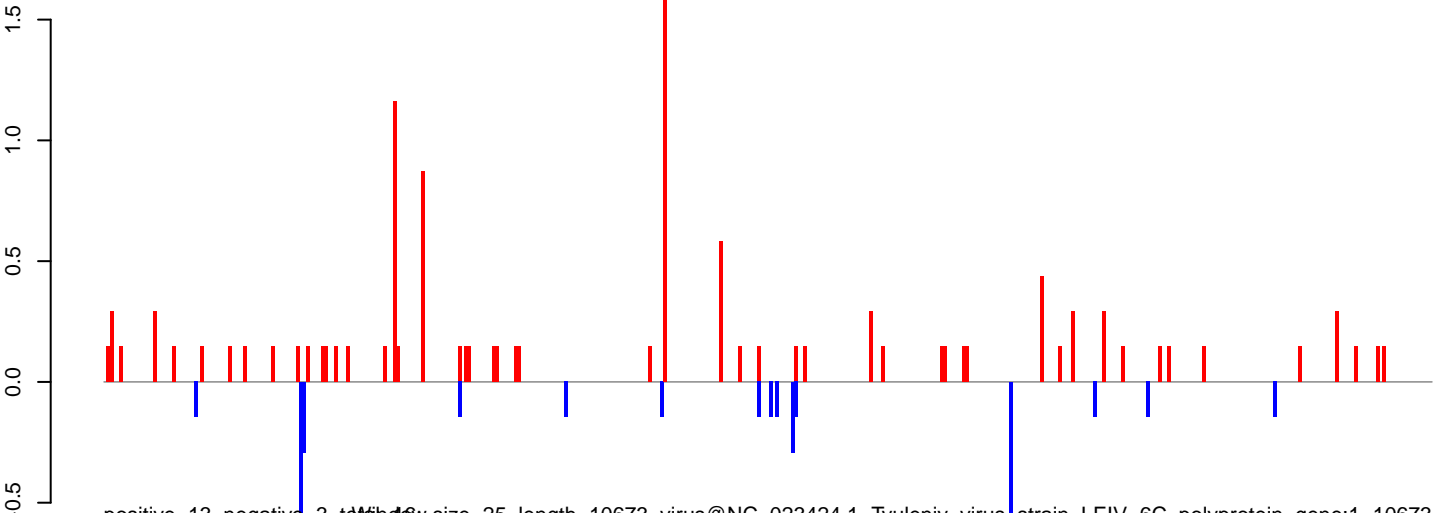
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



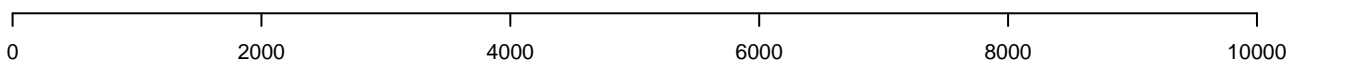
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



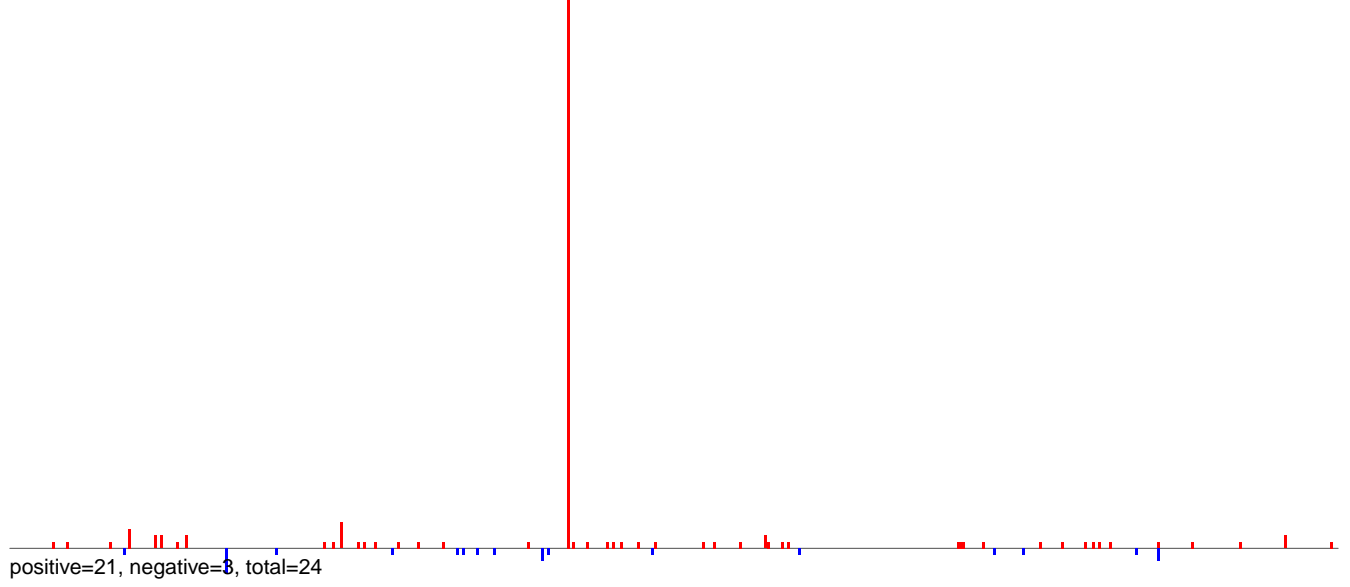
AeAeg_LVP_pMos3DBhspPBac.rep



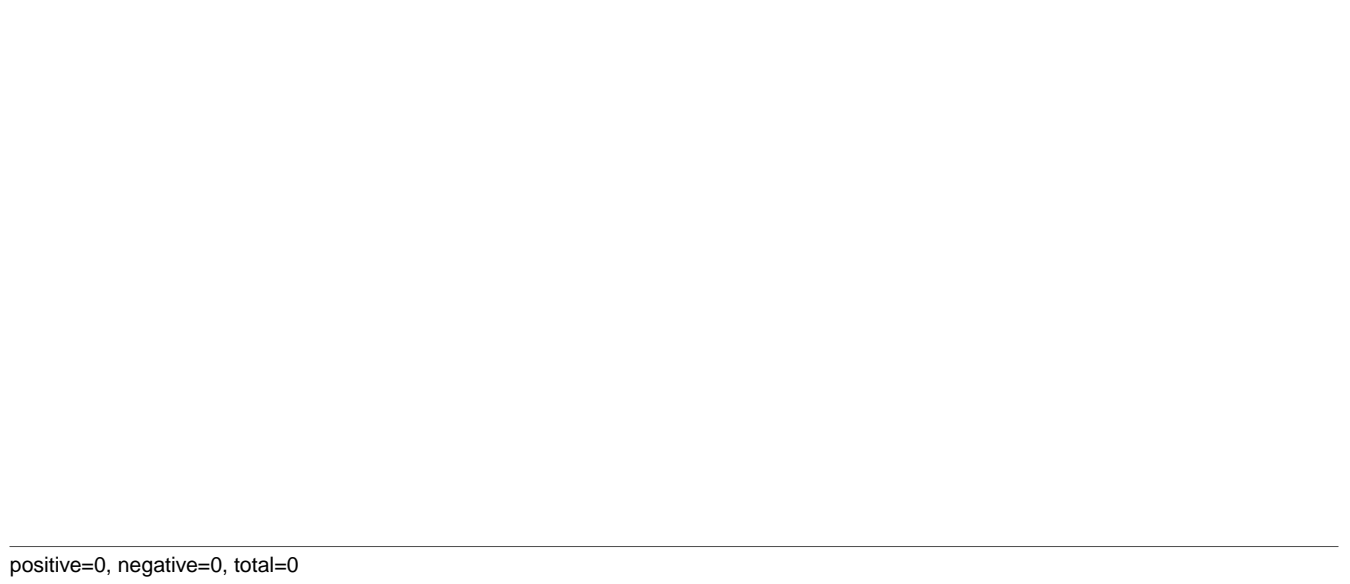
Window size=25, length=10673, virus@NC_023424.1-Tyuleniy_virus_strain_LEIV-6C_polyprotein_gene:1-10673



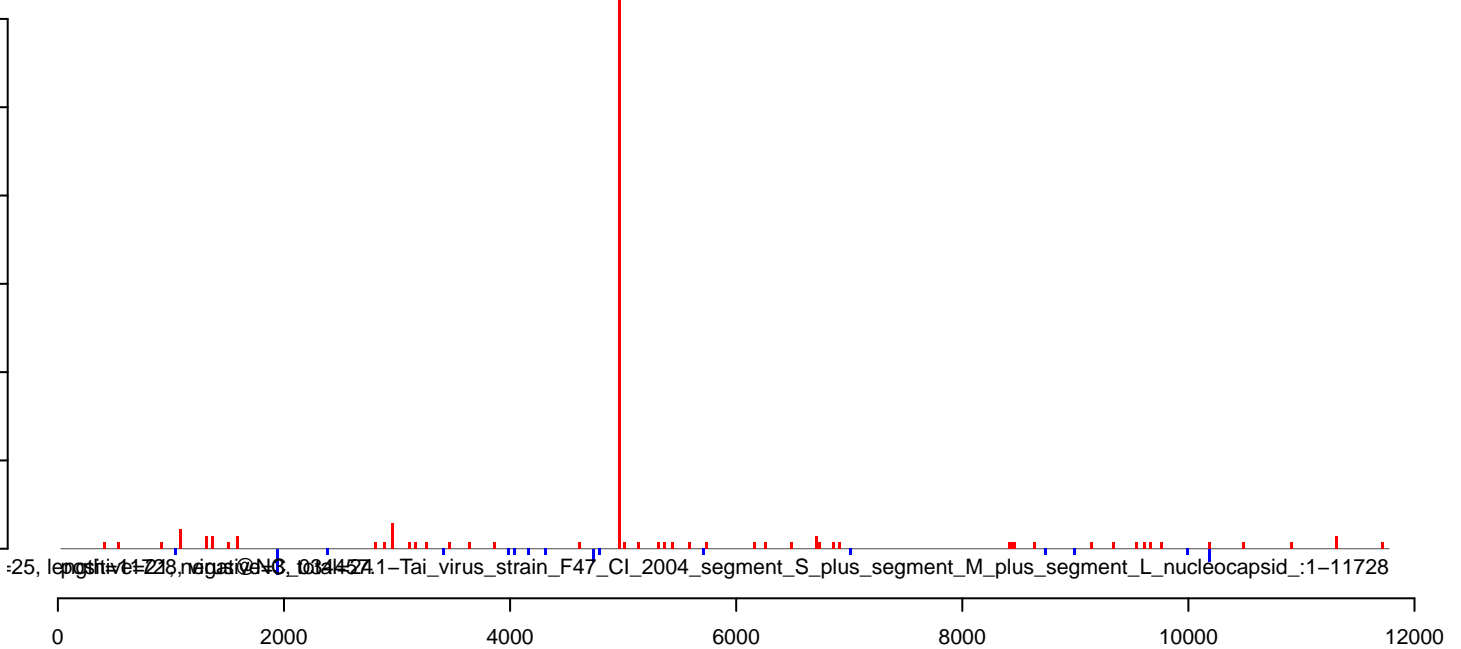
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



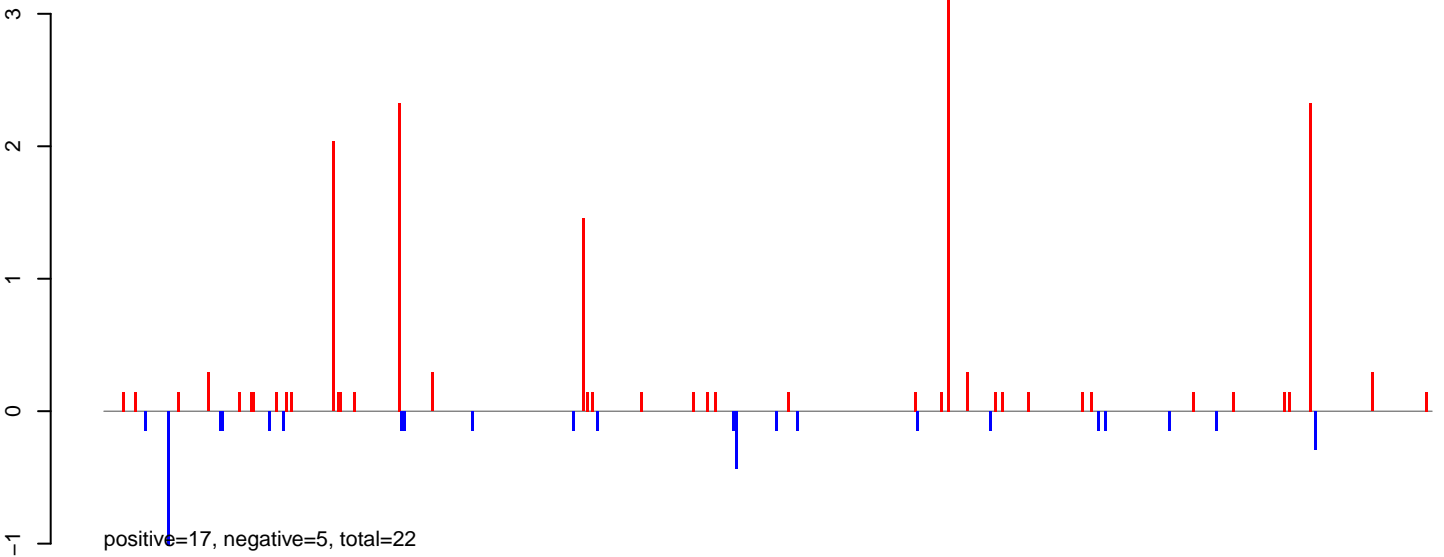
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



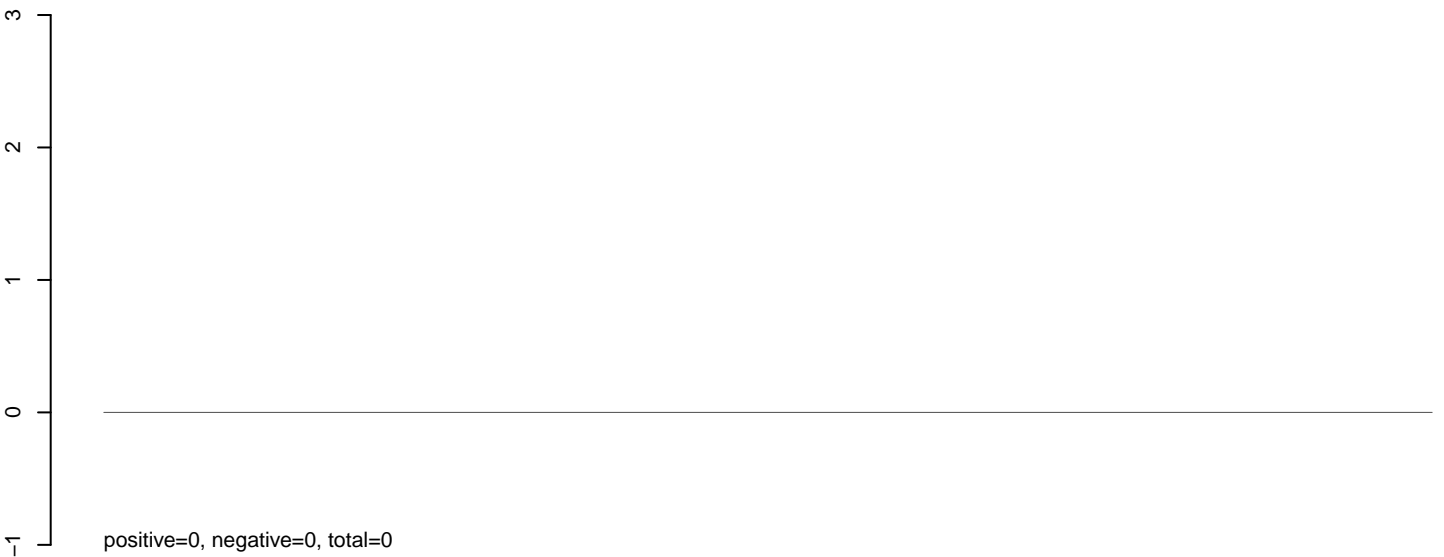
AeAeg_LVP_pMos3DBhspPBac.rep



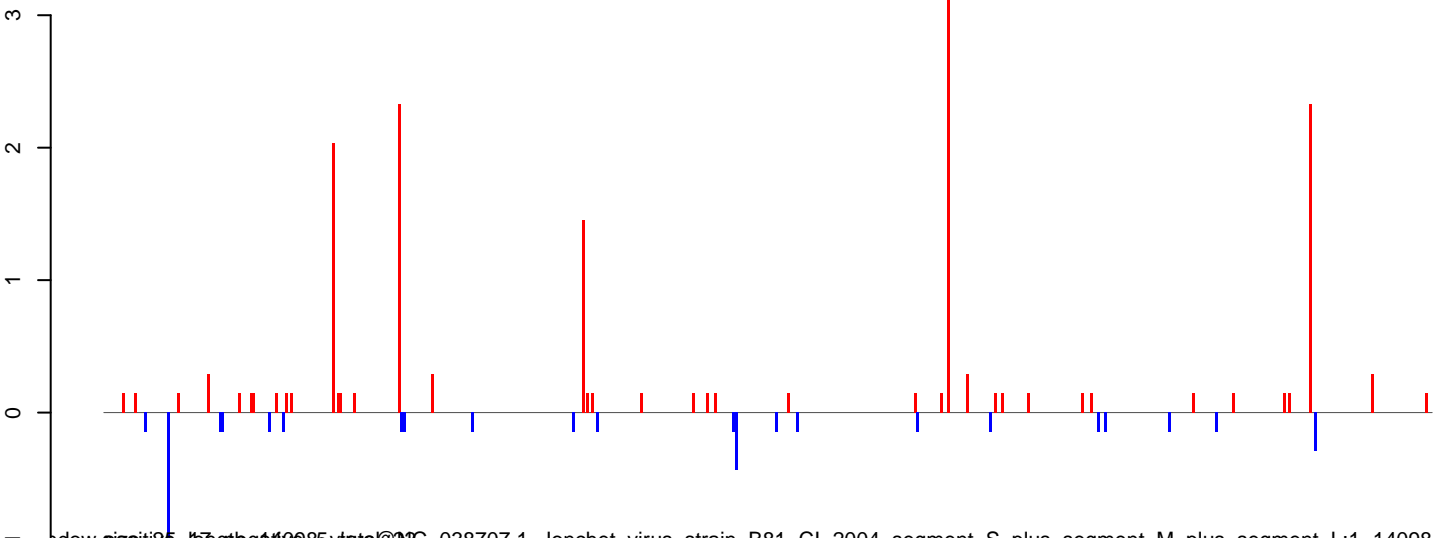
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



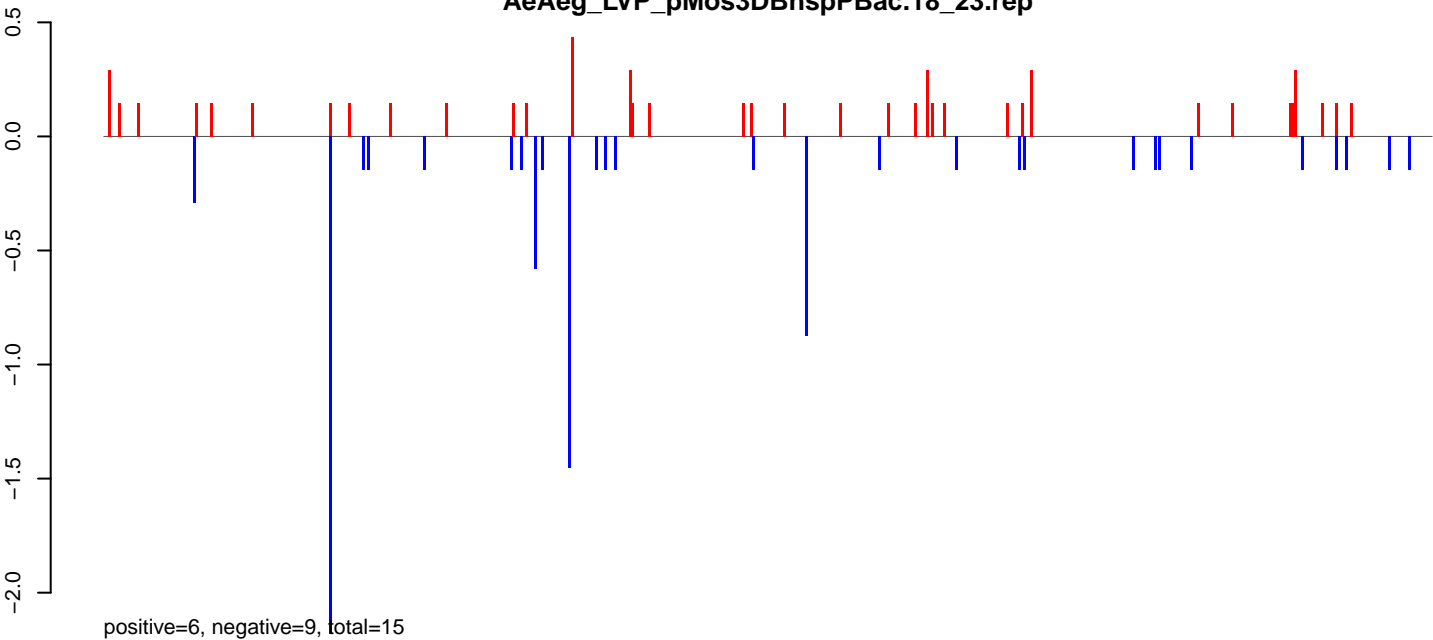
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



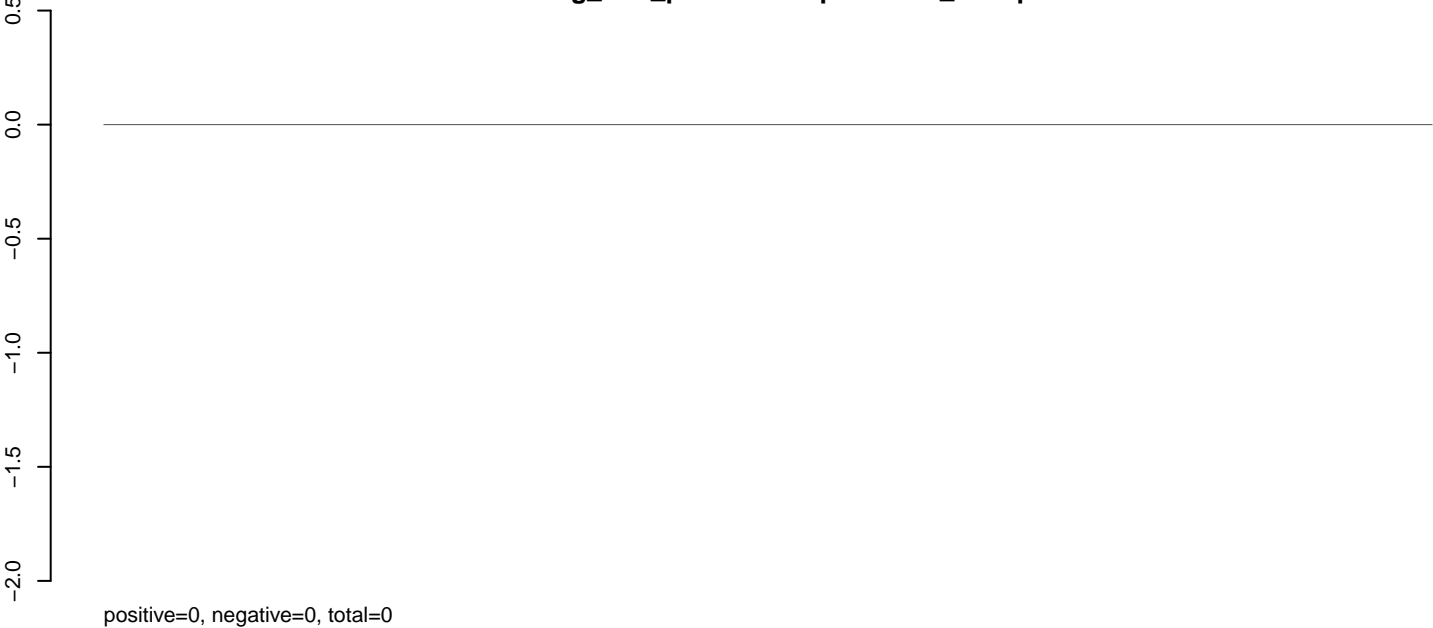
AeAeg_LVP_pMos3DBhspPBac.rep



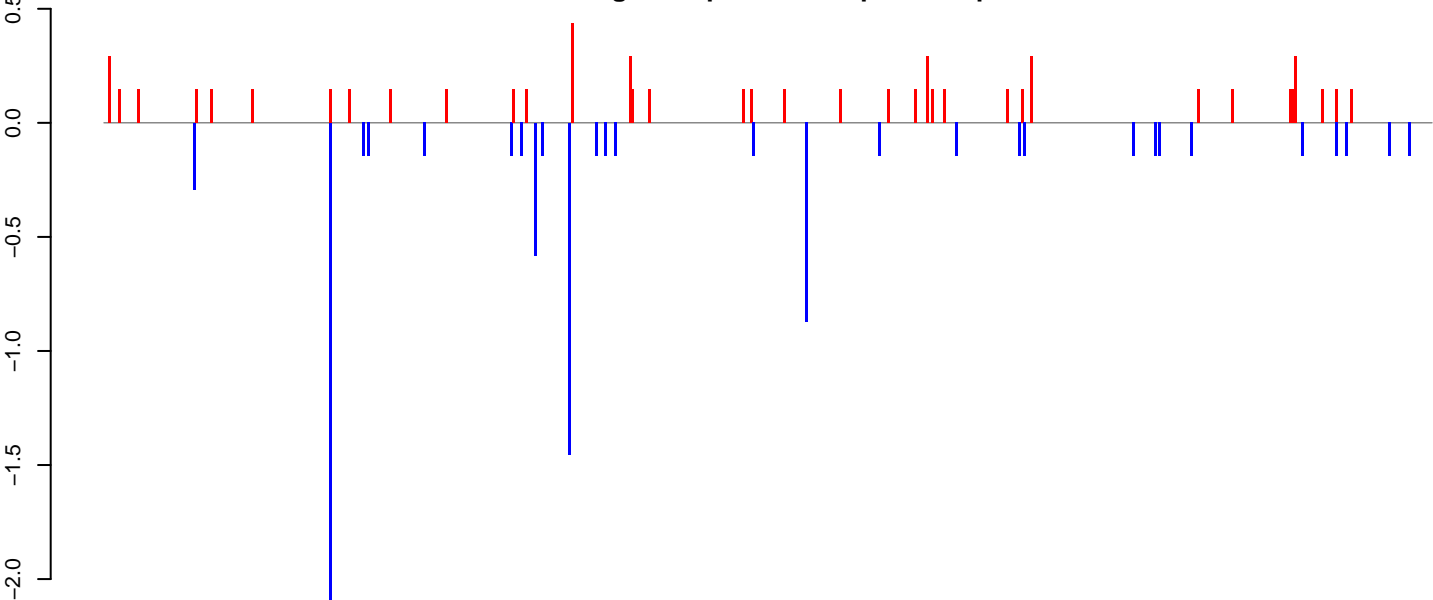
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



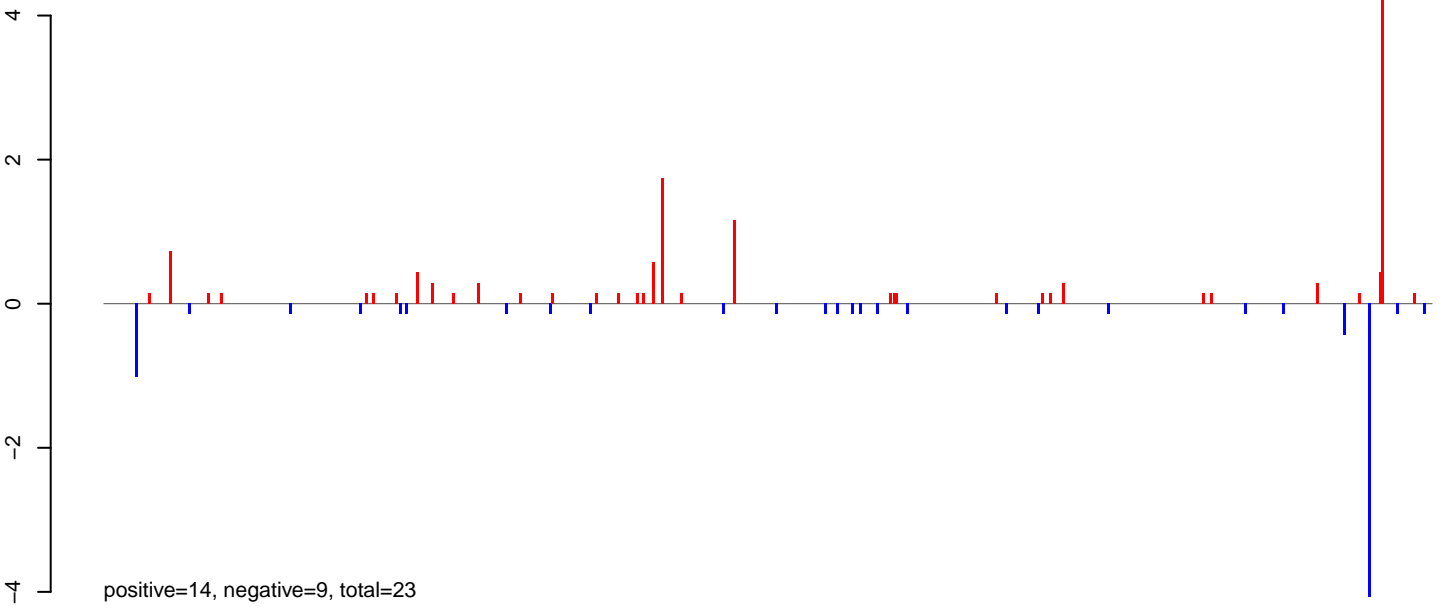
AeAeg_LVP_pMos3DBhspPBac.rep



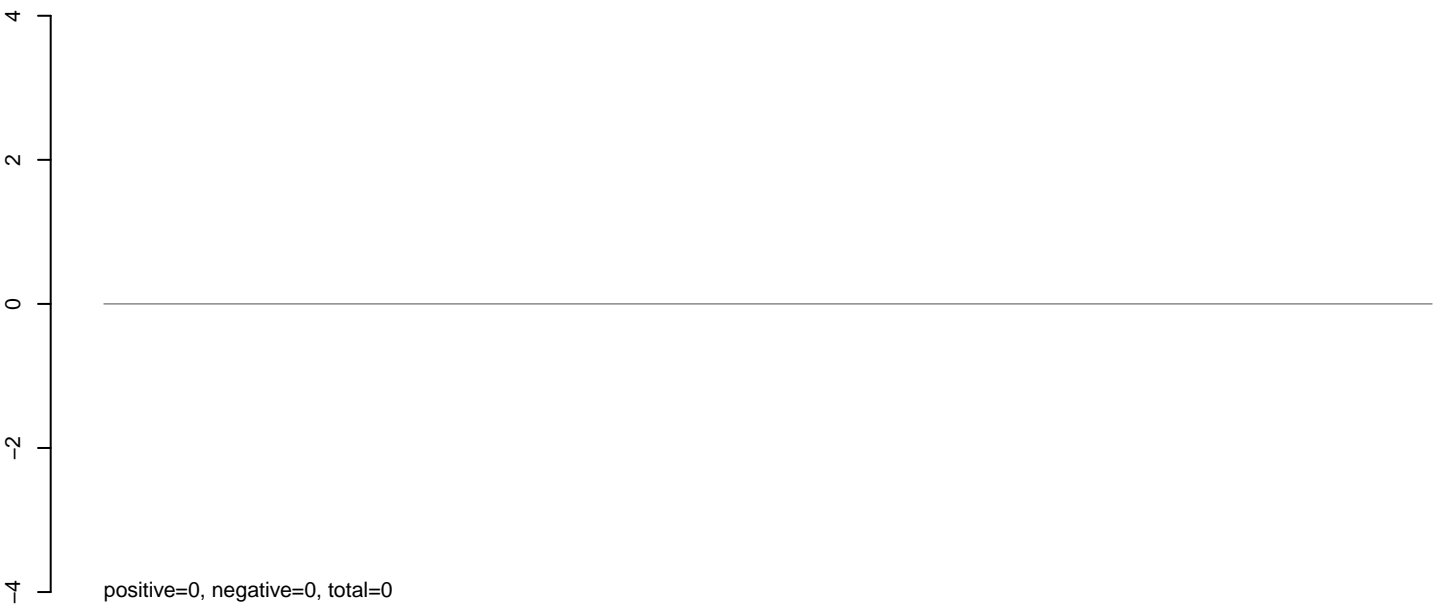
th=13706, virus=9, gain=0, total=15, An_Mosquito_Virus_9_strain_JX1-13_ORF1_ORF1_ORF2_ORF2_ORF3_ORF3_glycoprotein_G_:1-13706

0 2000 4000 6000 8000 10000 12000 14000

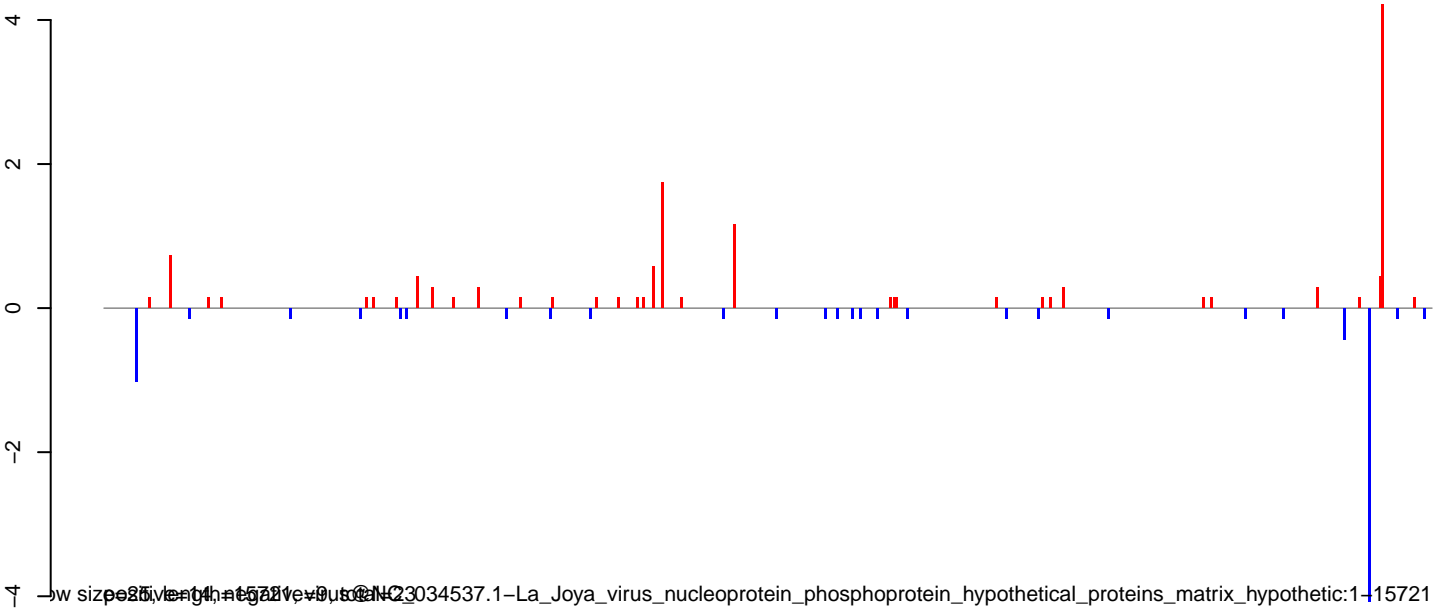
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

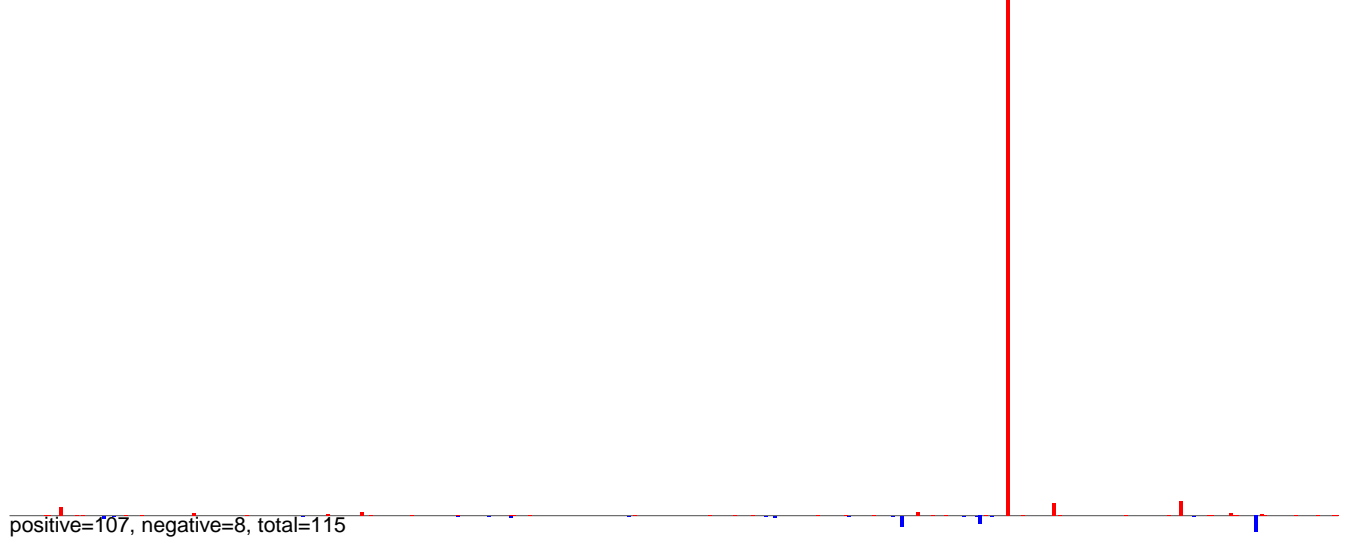


AeAeg_LVP_pMos3DBhspPBac.rep

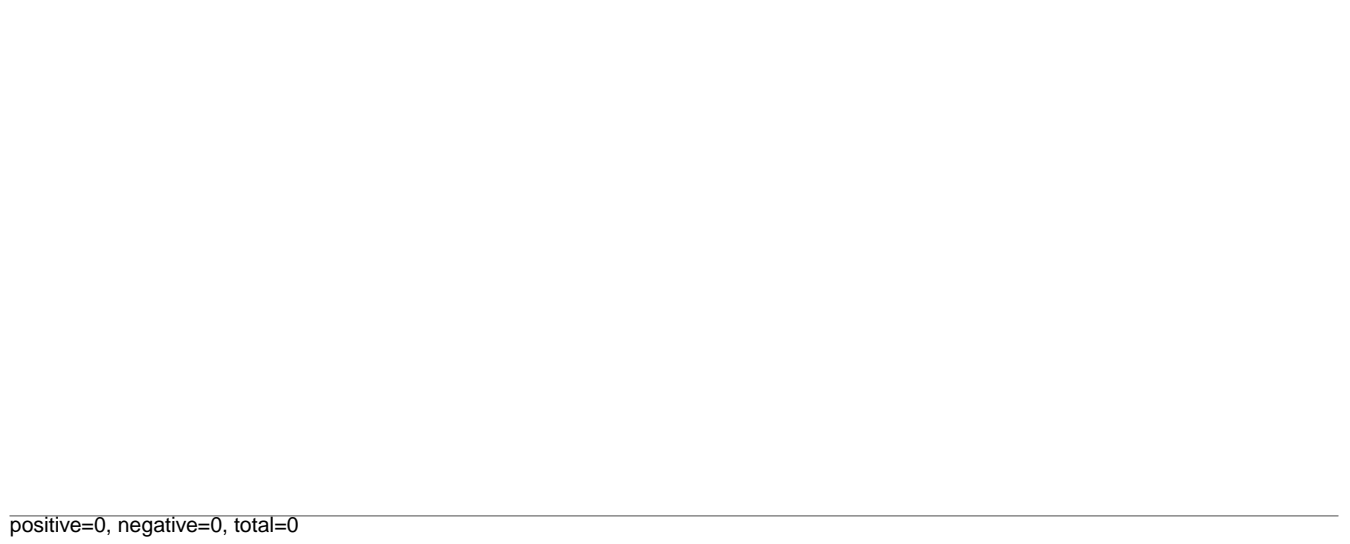


row size=25, length=15721, positive=14, negative=9, total=23
@NC_030345.1-La_Joya_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_hypothetic:1-15721

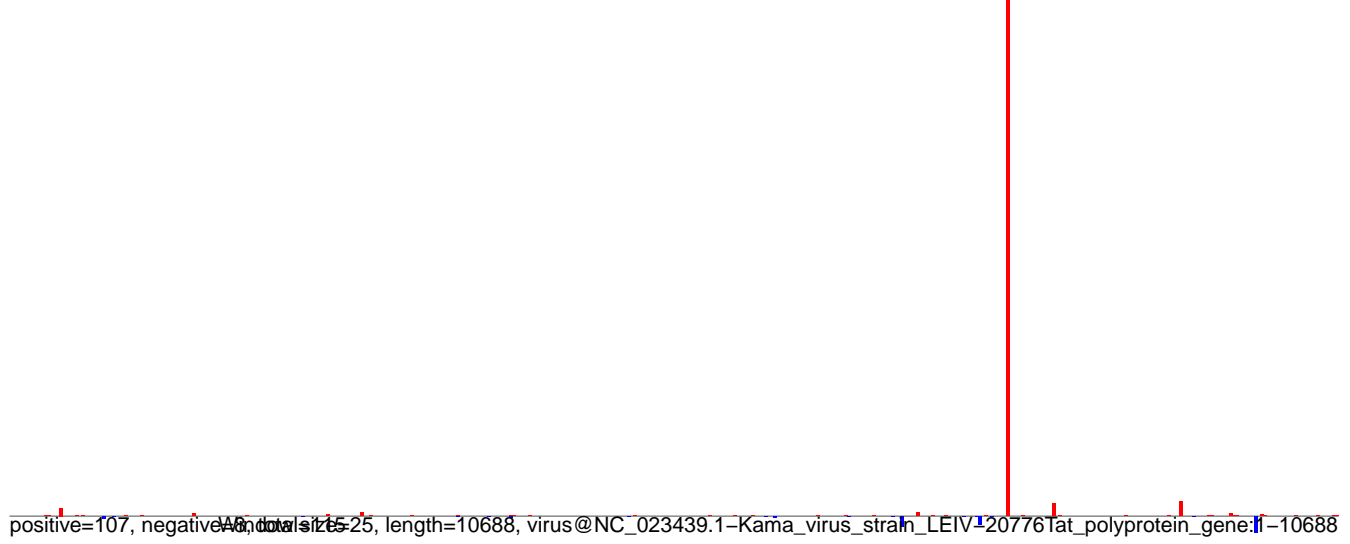
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

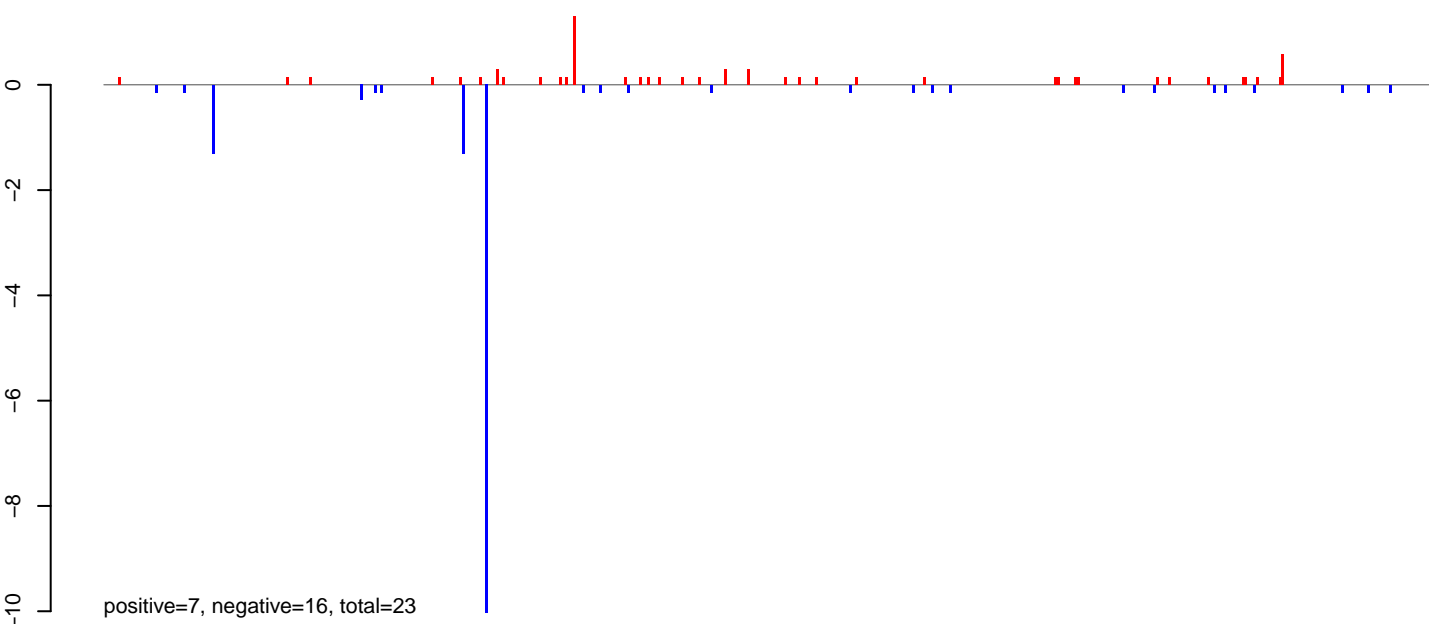


AeAeg_LVP_pMos3DBhspPBac.rep

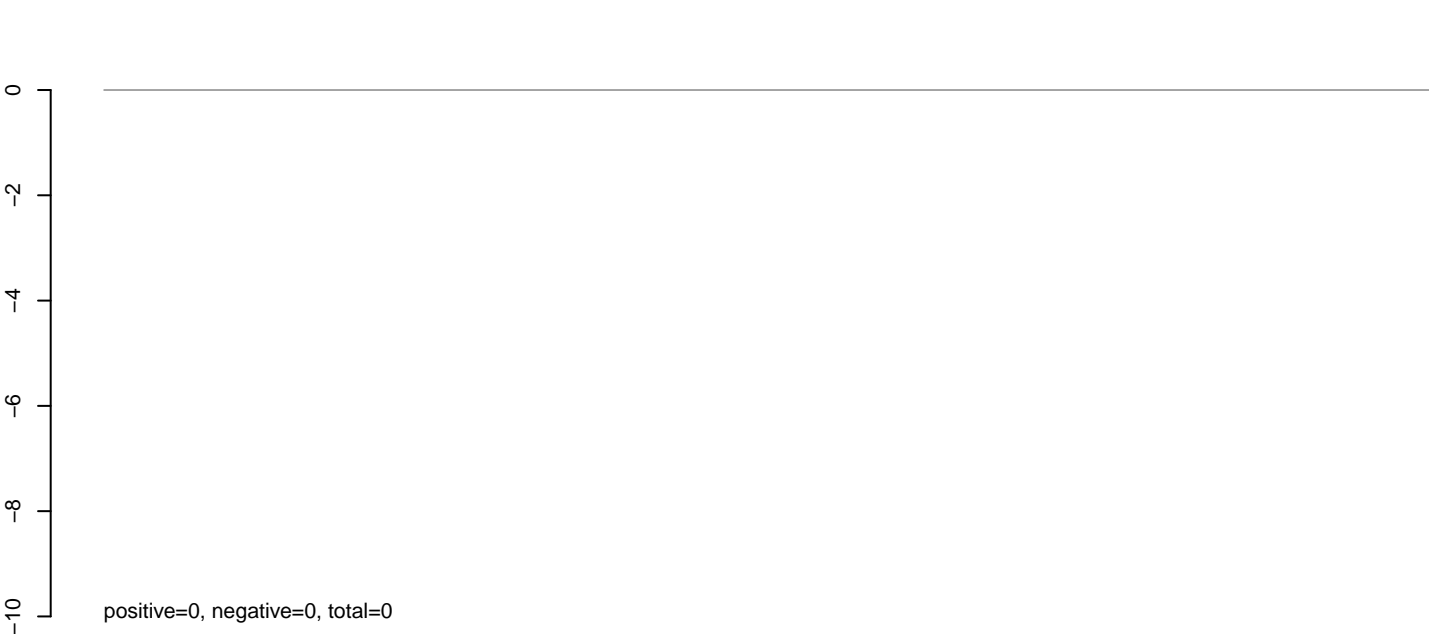


0 2000 4000 6000 8000 10000
Annotation: 1-10688, virus@NC_023439.1-Kama_virus_strain_LEIV-207761at_polyprotein_gene.1-10688

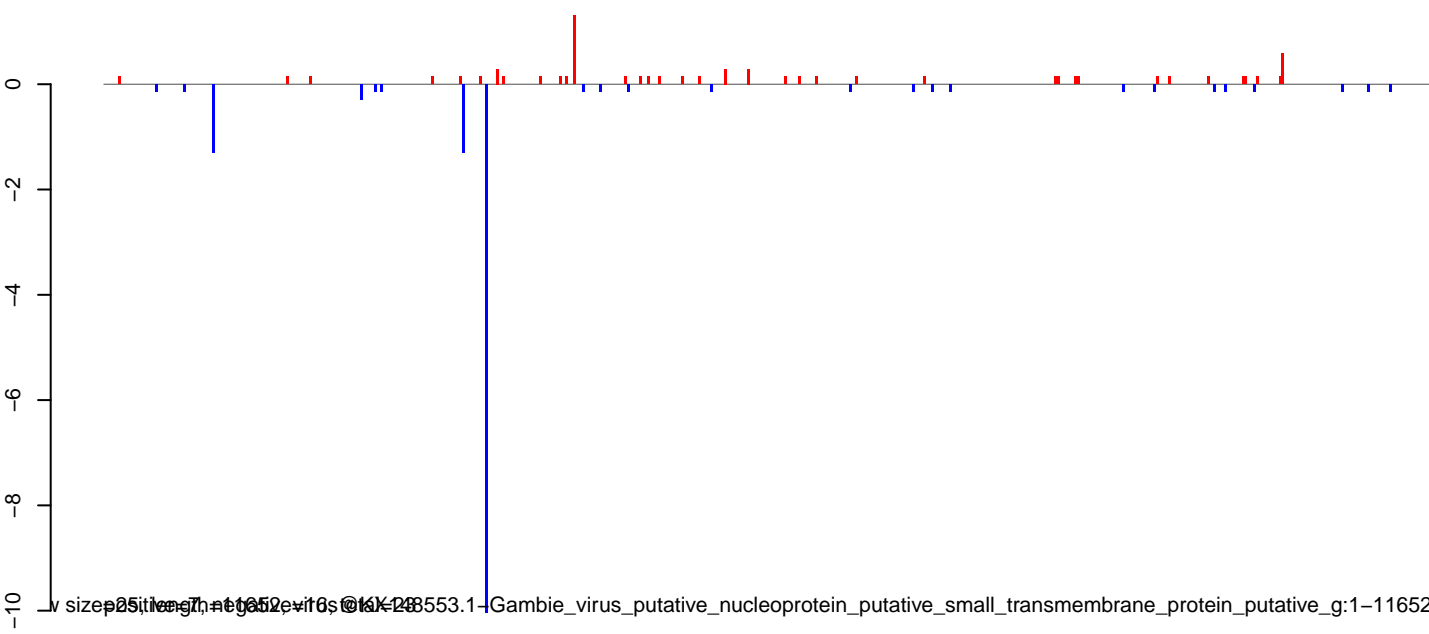
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



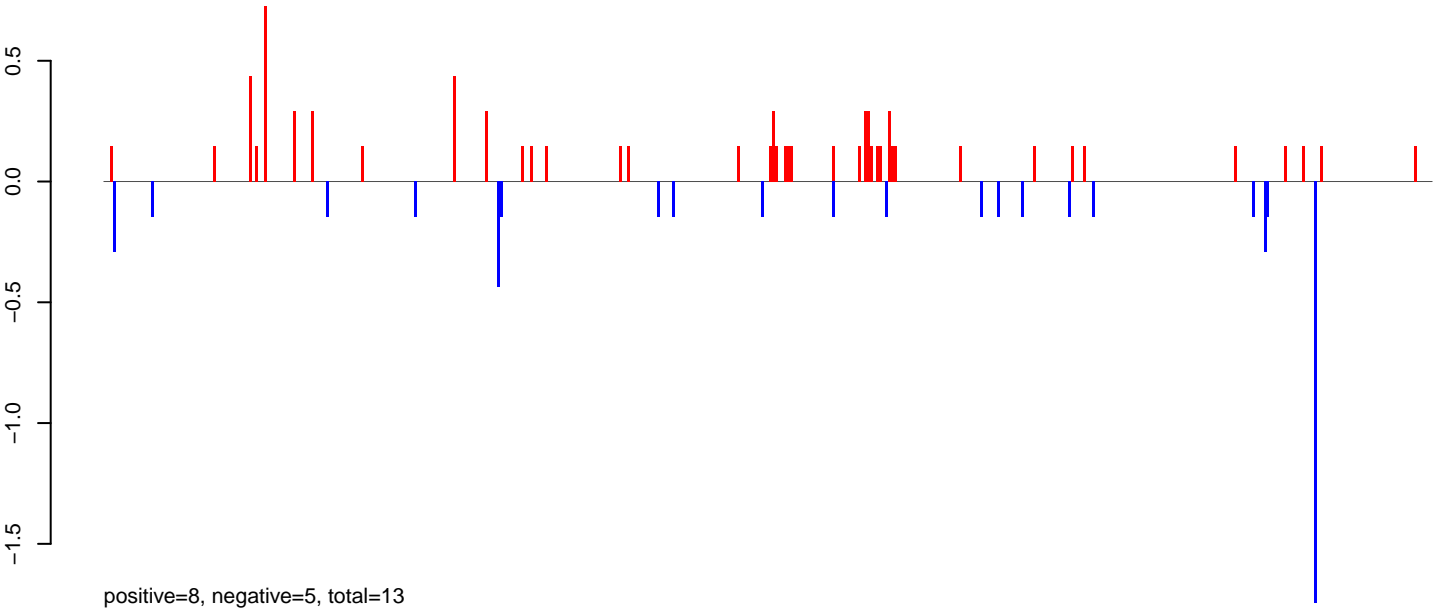
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



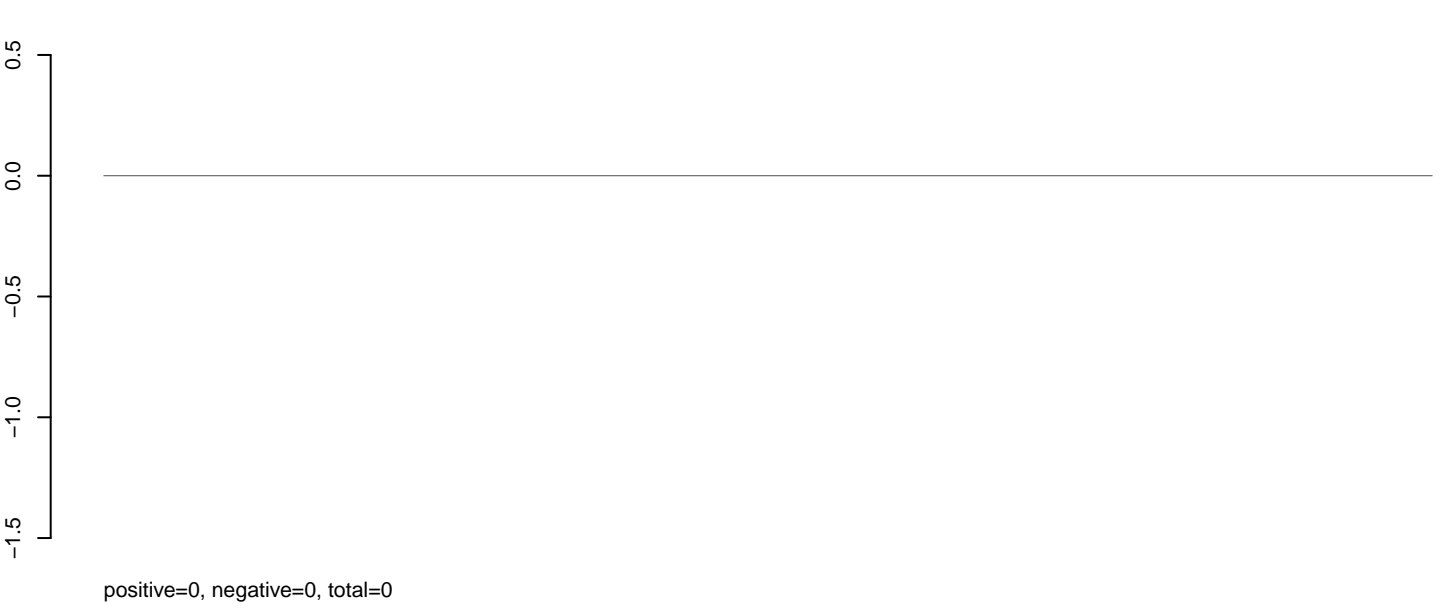
AeAeg_LVP_pMos3DBhspPBac.rep



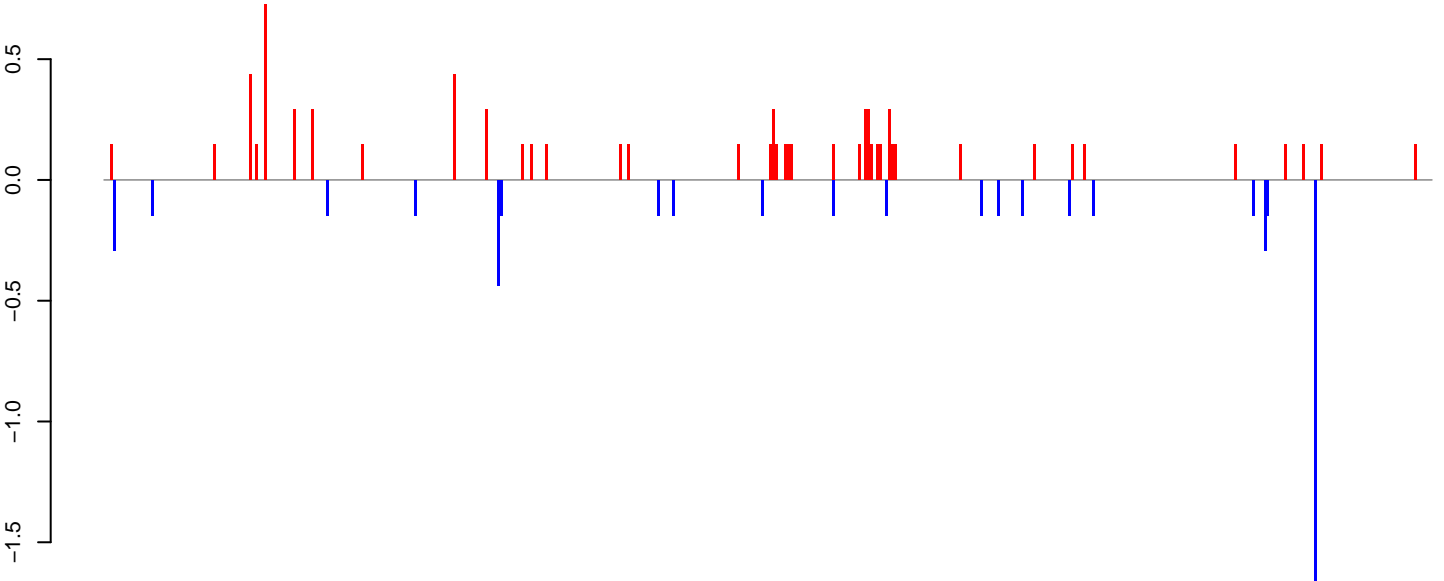
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



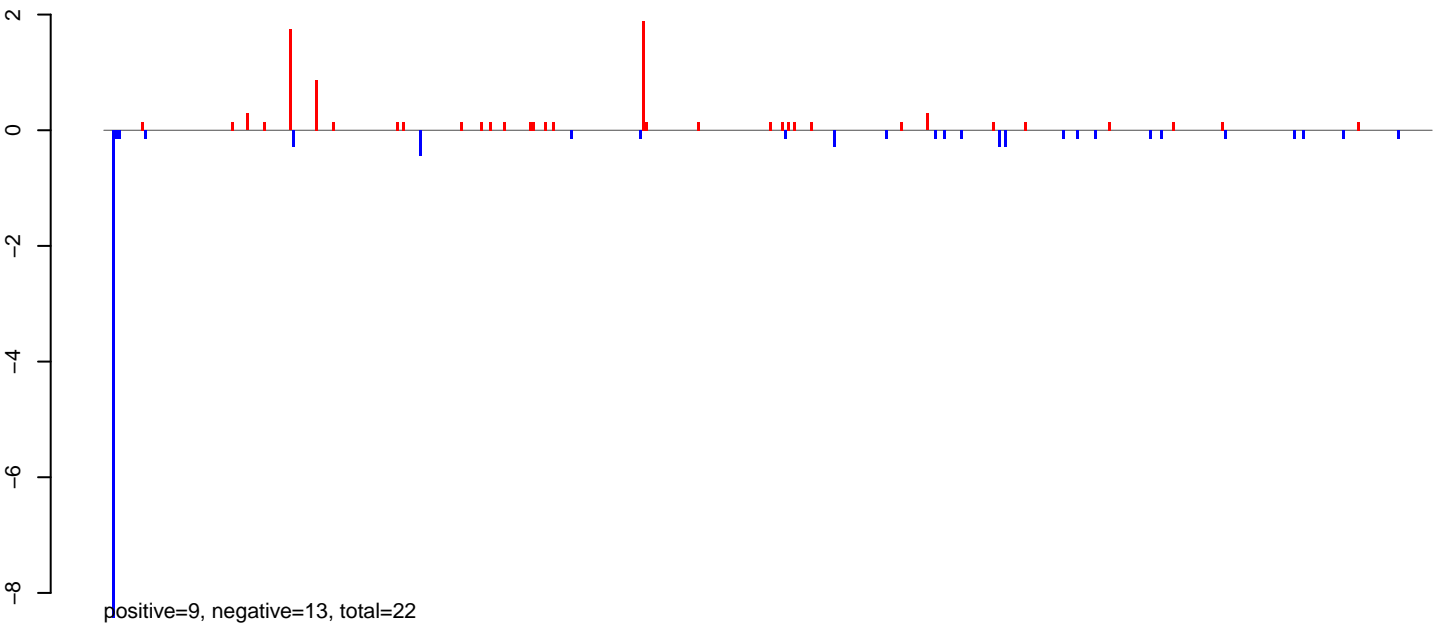
AeAeg_LVP_pMos3DBhspPBac.rep



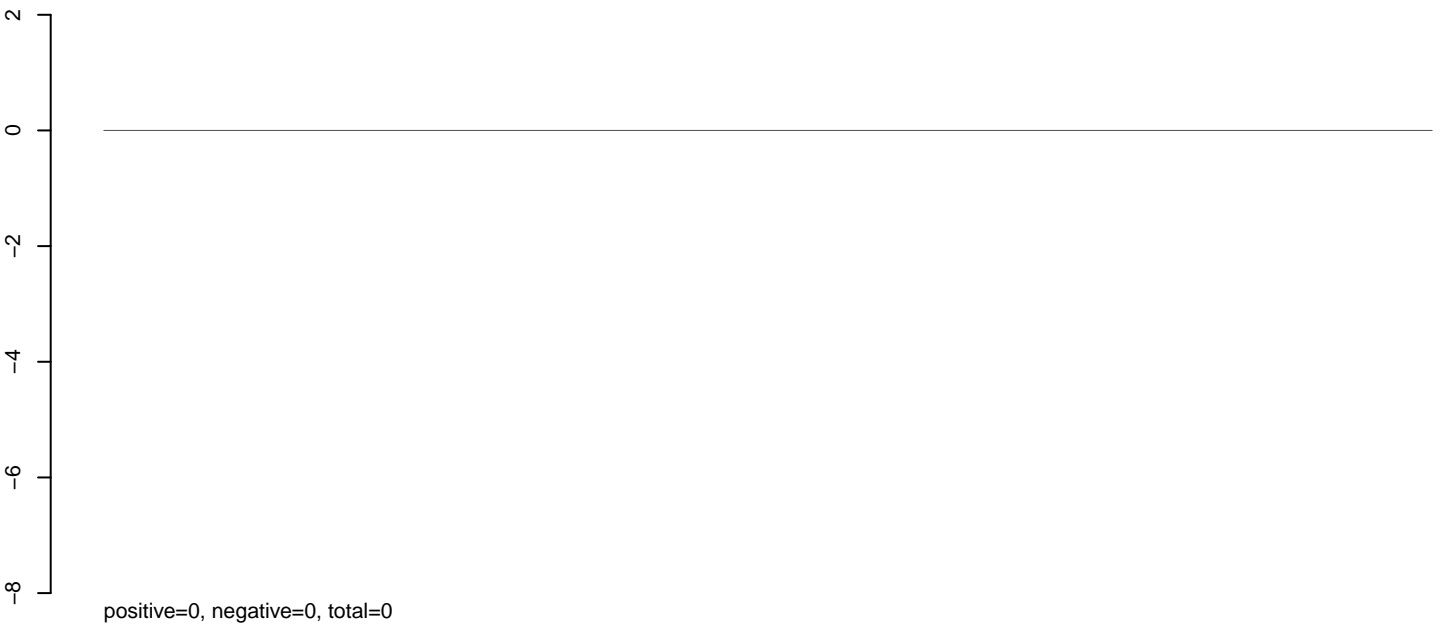
Window size=25, length=11220, virus@NC_025399.1-Oak-Vale_virus_strain_CSIRO_1342:1-11220

0 2000 4000 6000 8000 10000

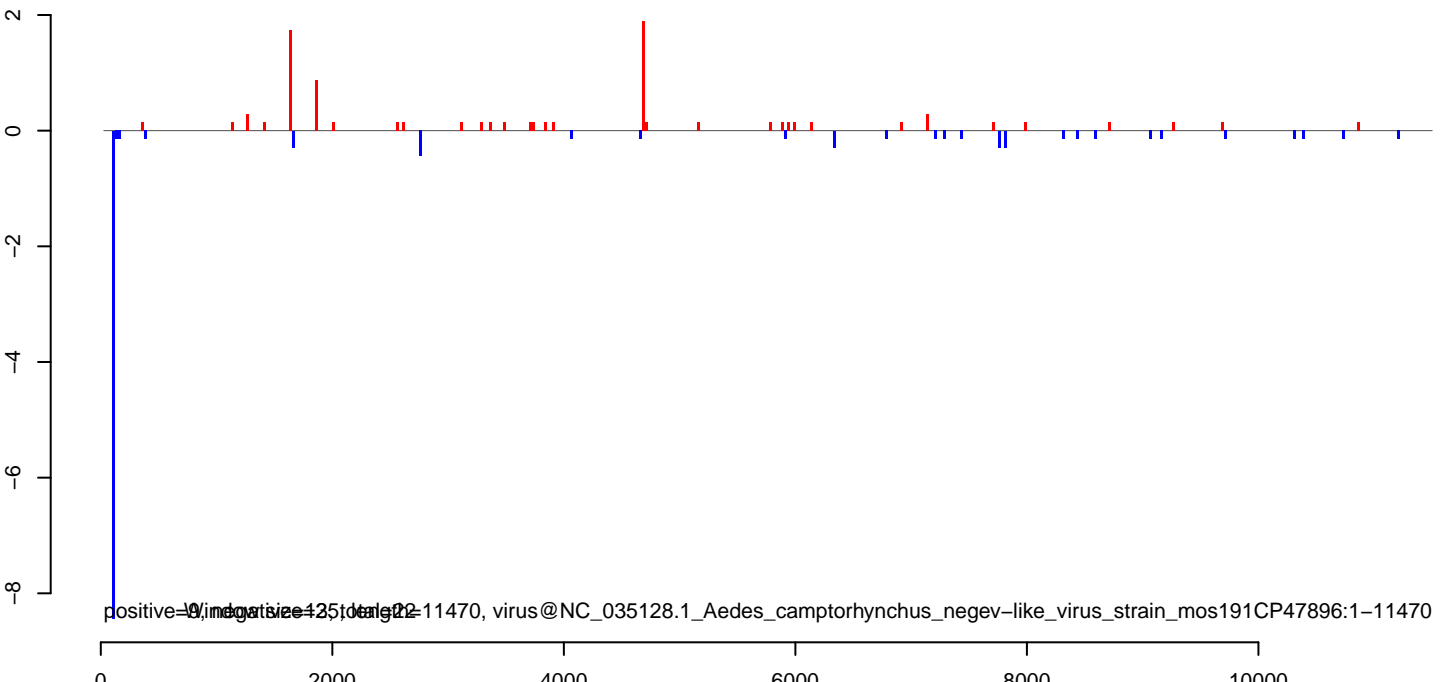
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

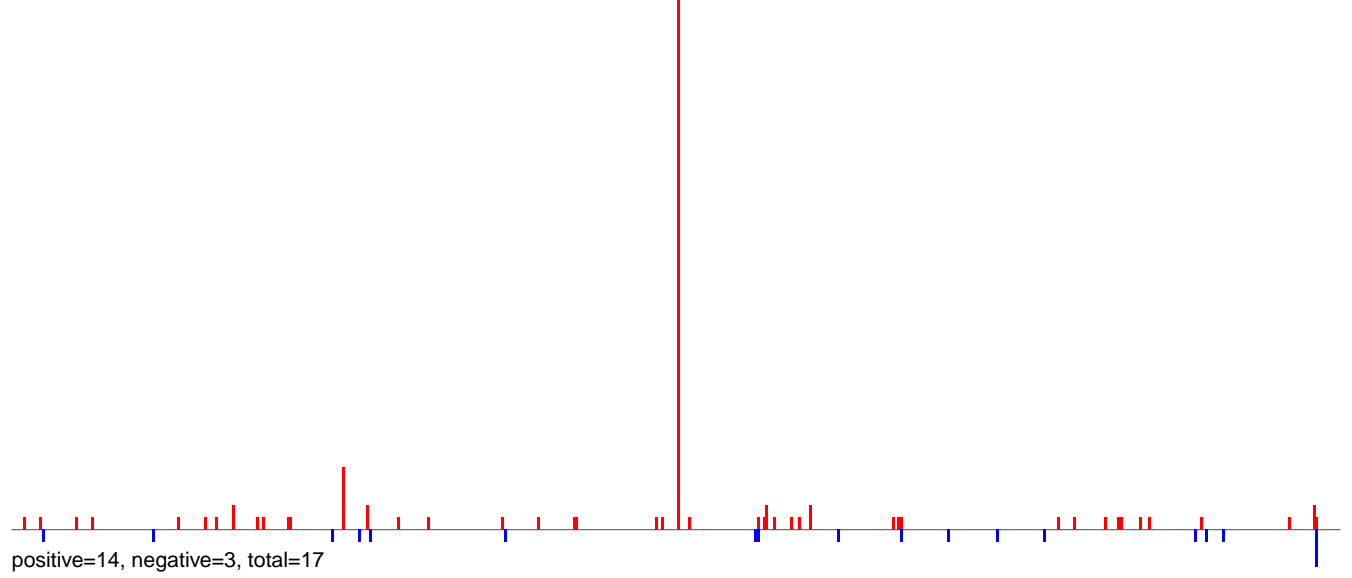


AeAeg_LVP_pMos3DBhspPBac.rep

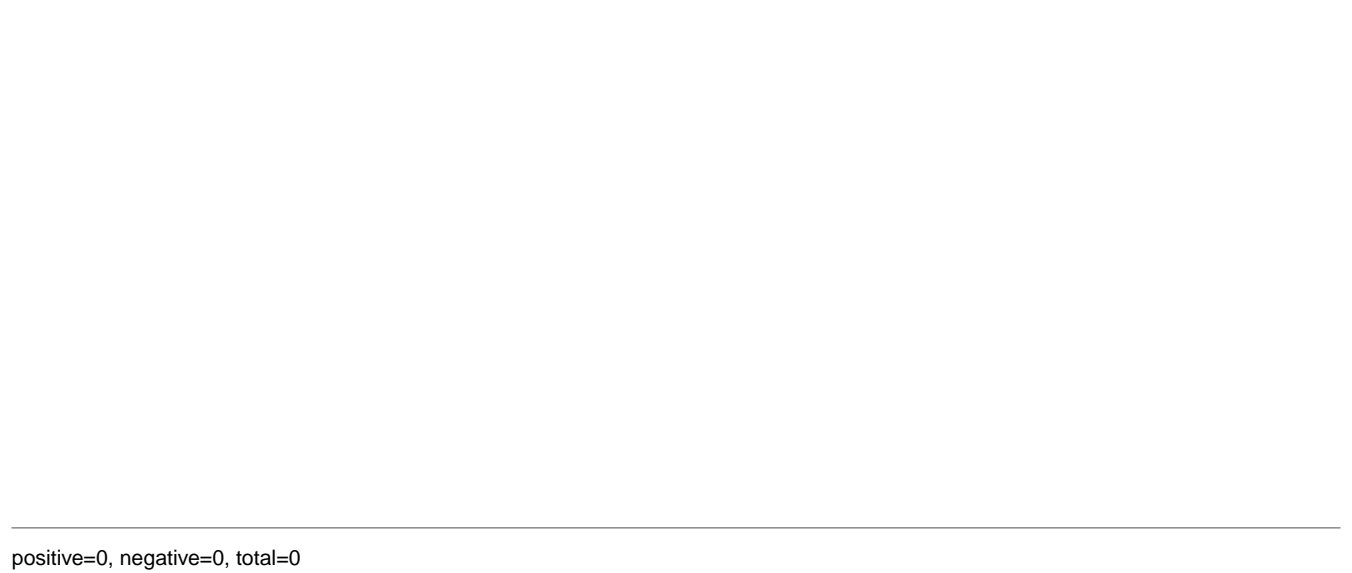


positive=9, negative=13, total=22, virus@NC_035128.1_Aedes_camptorhynchus_negev-like_virus_strain_mos191CP47896:1-11470

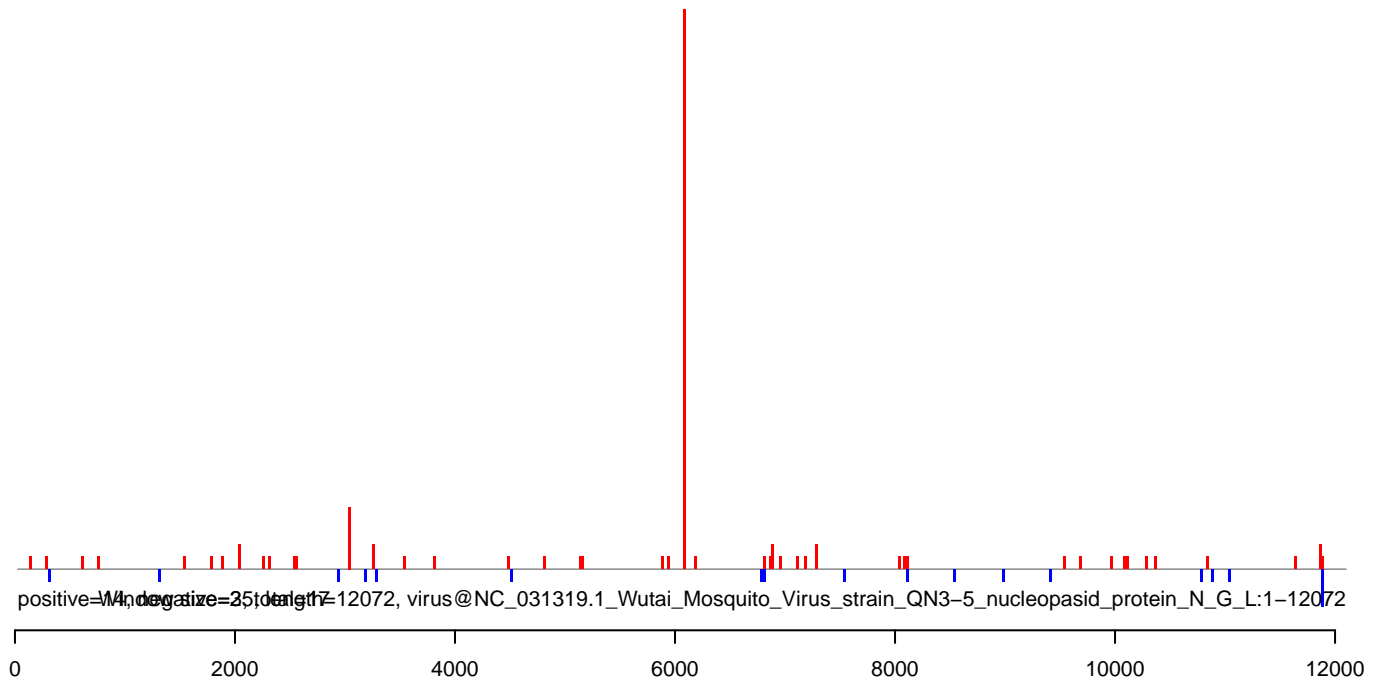
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



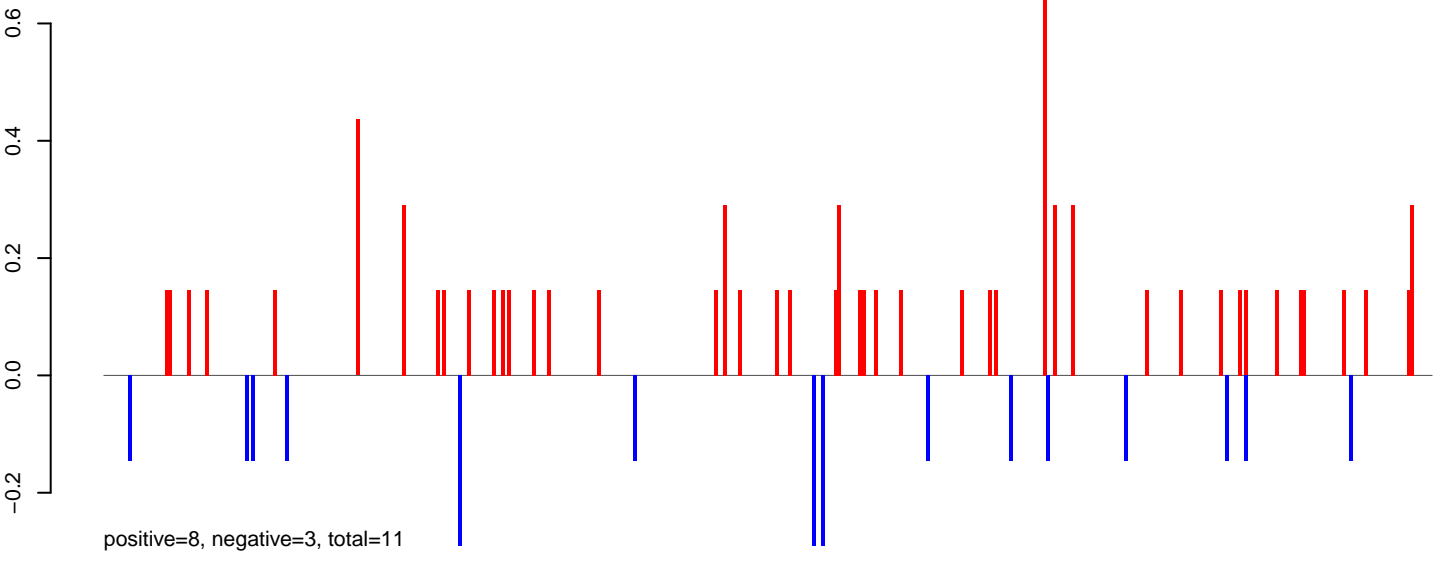
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



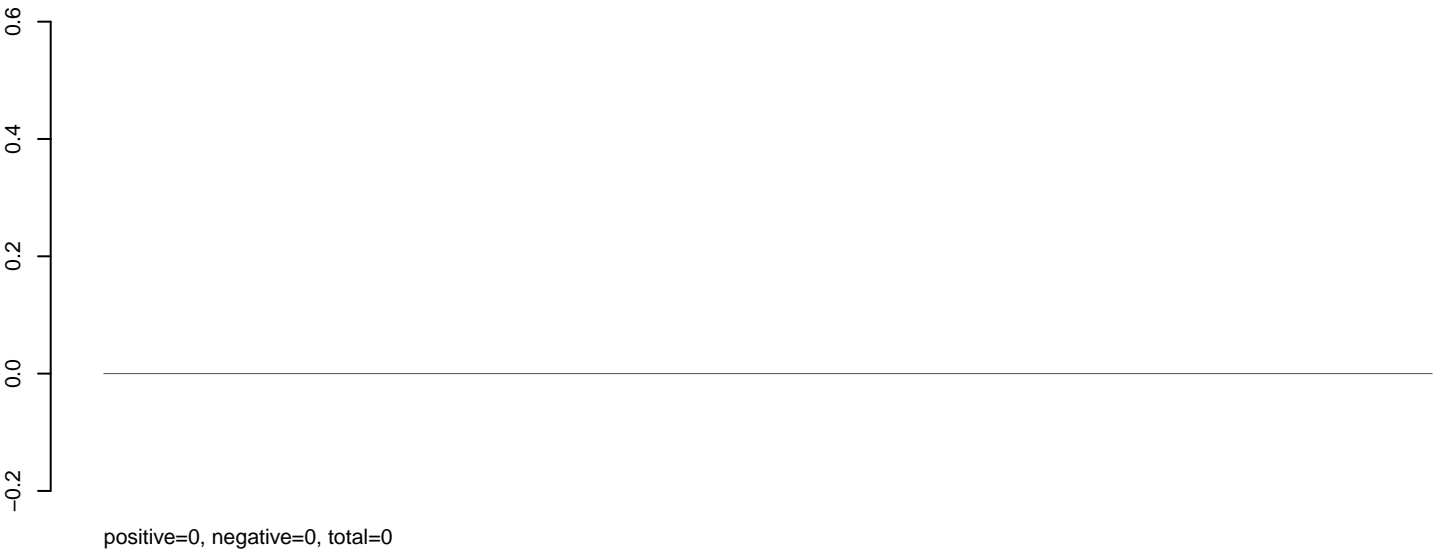
AeAeg_LVP_pMos3DBhspPBac.rep



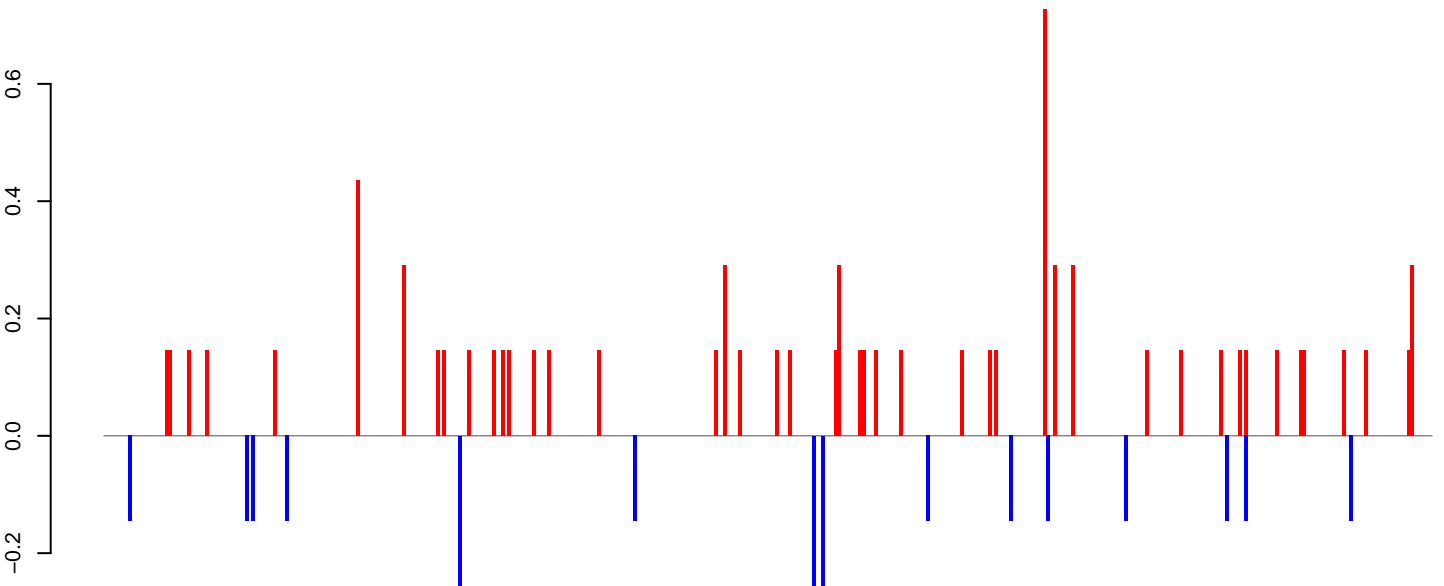
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

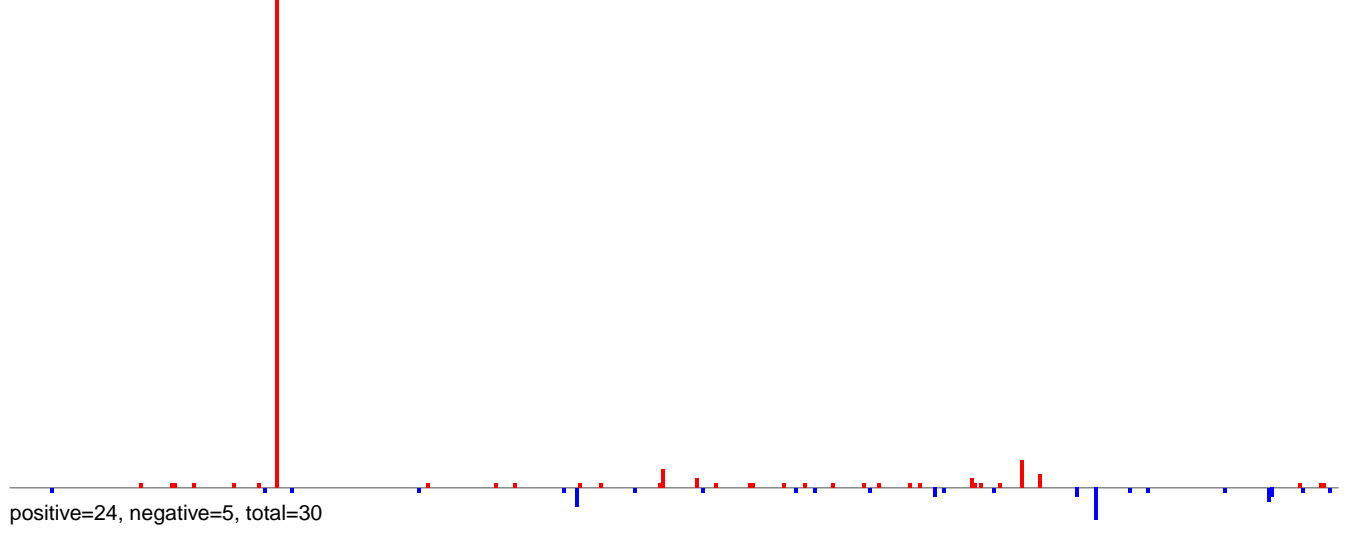


AeAeg_LVP_pMos3DBhspPBac.rep

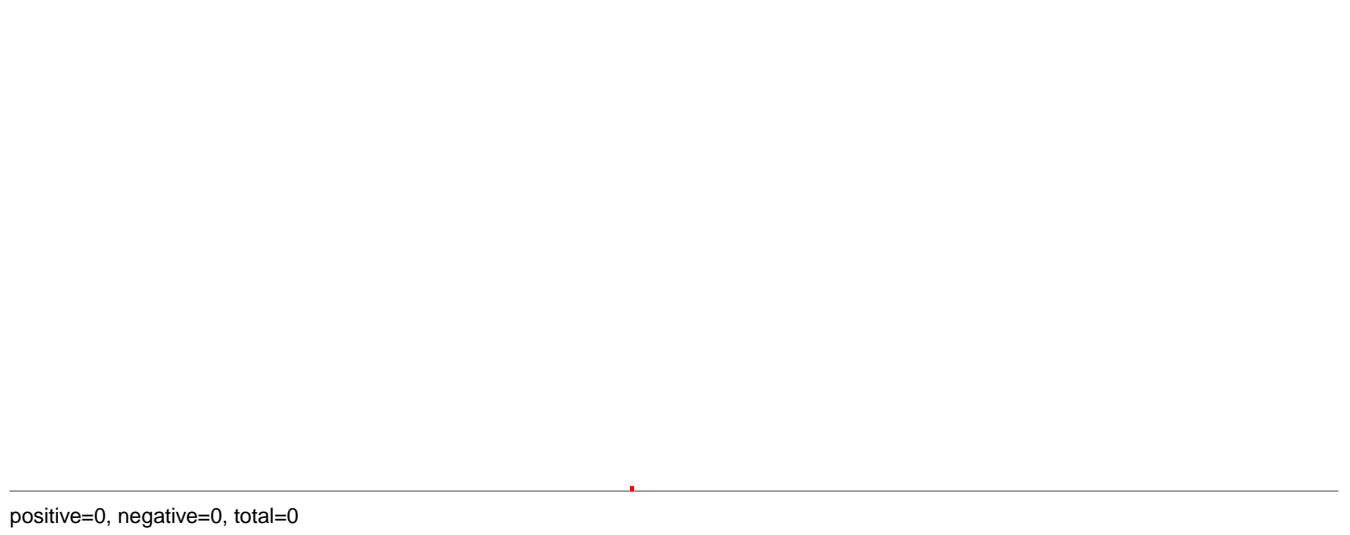


0 2000 4000 6000 8000 10000

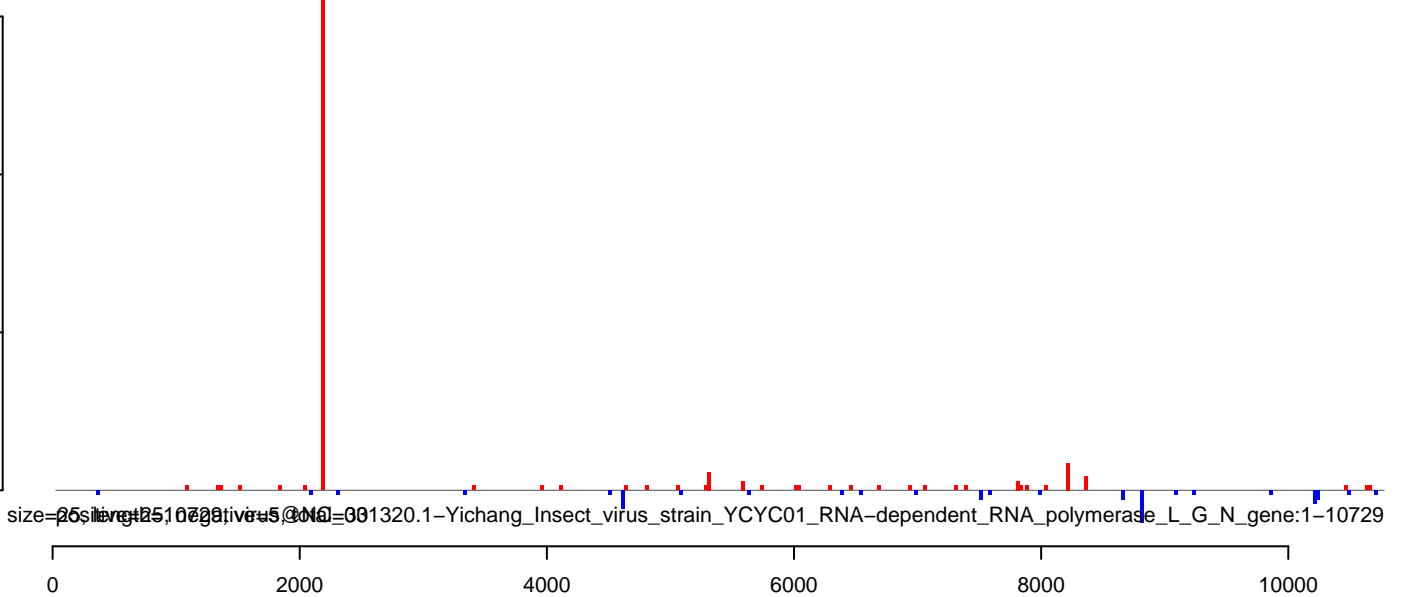
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



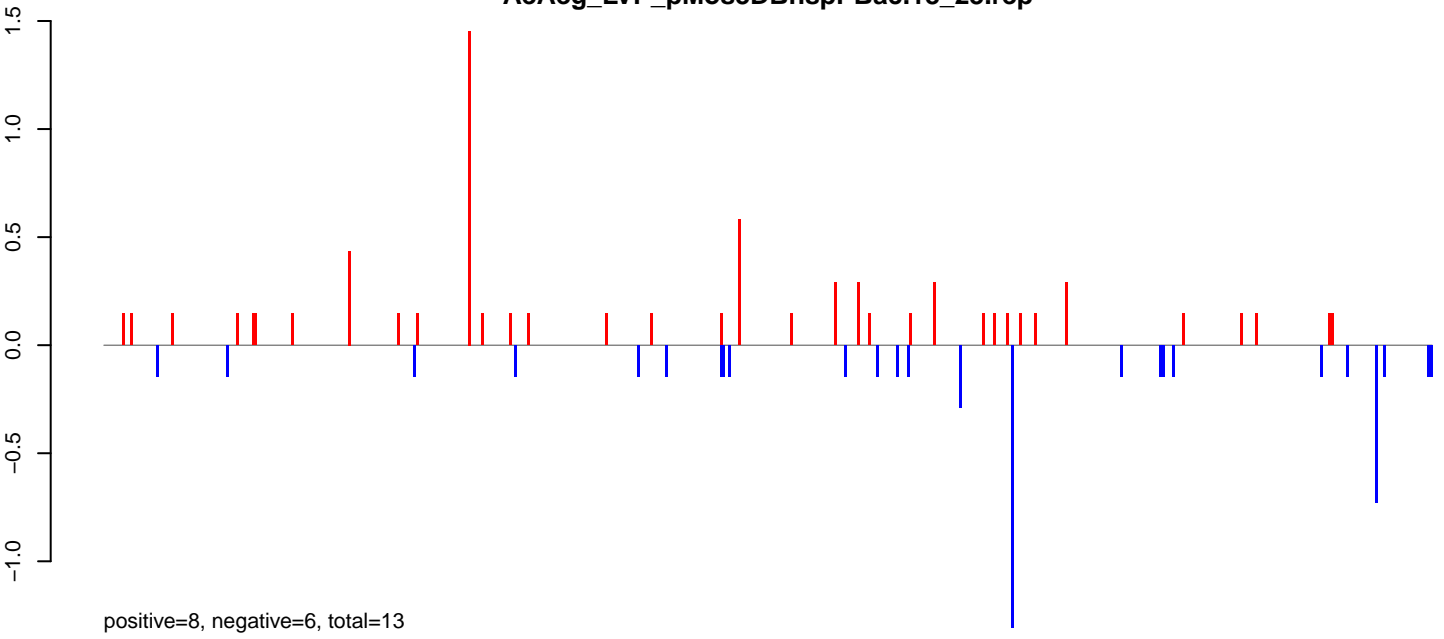
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



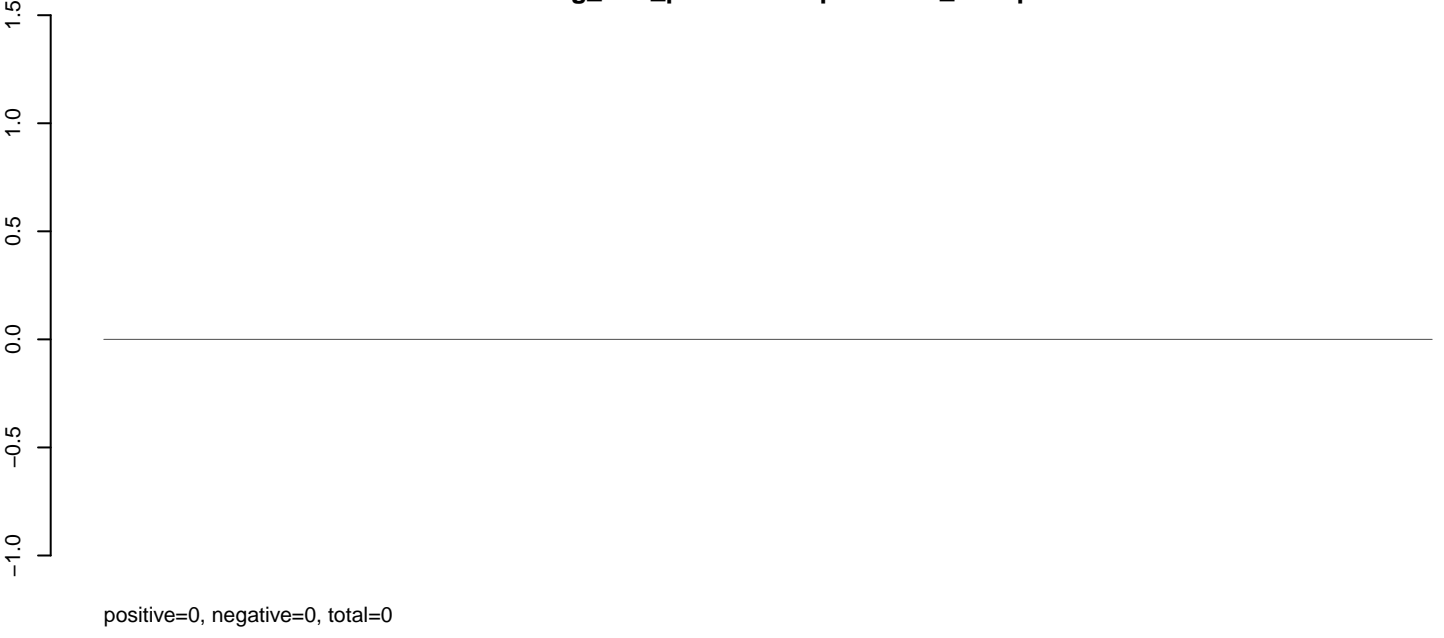
AeAeg_LVP_pMos3DBhspPBac.rep



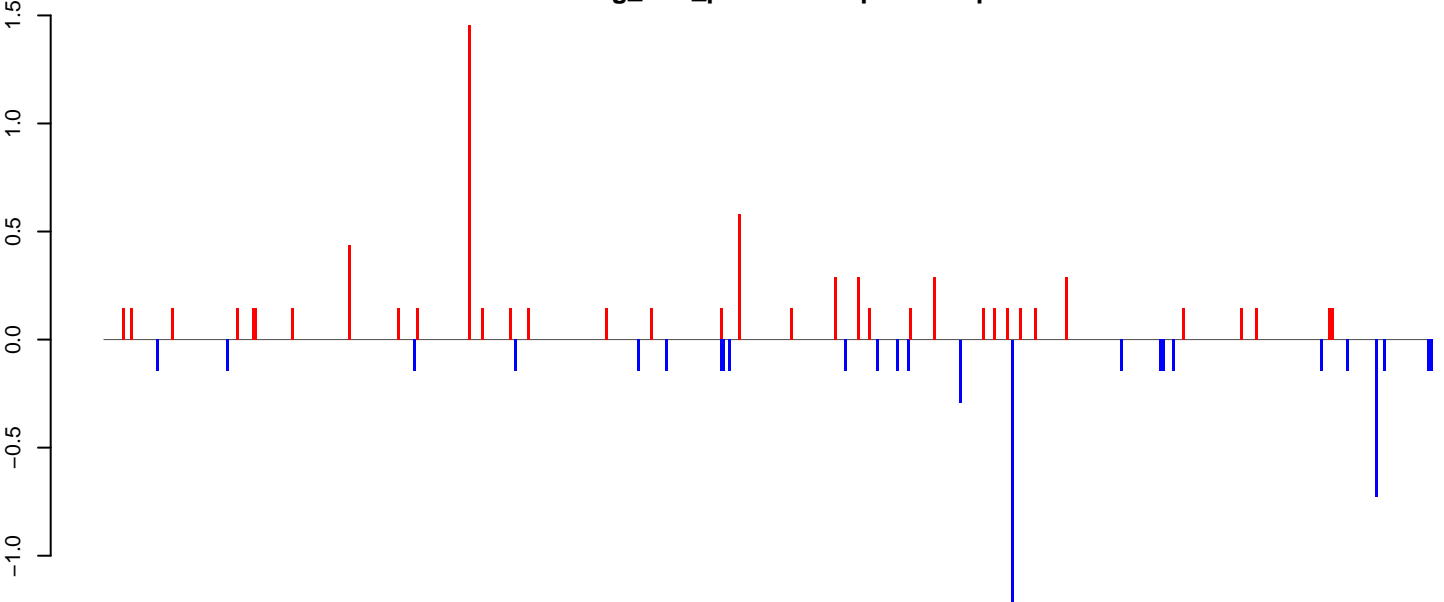
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



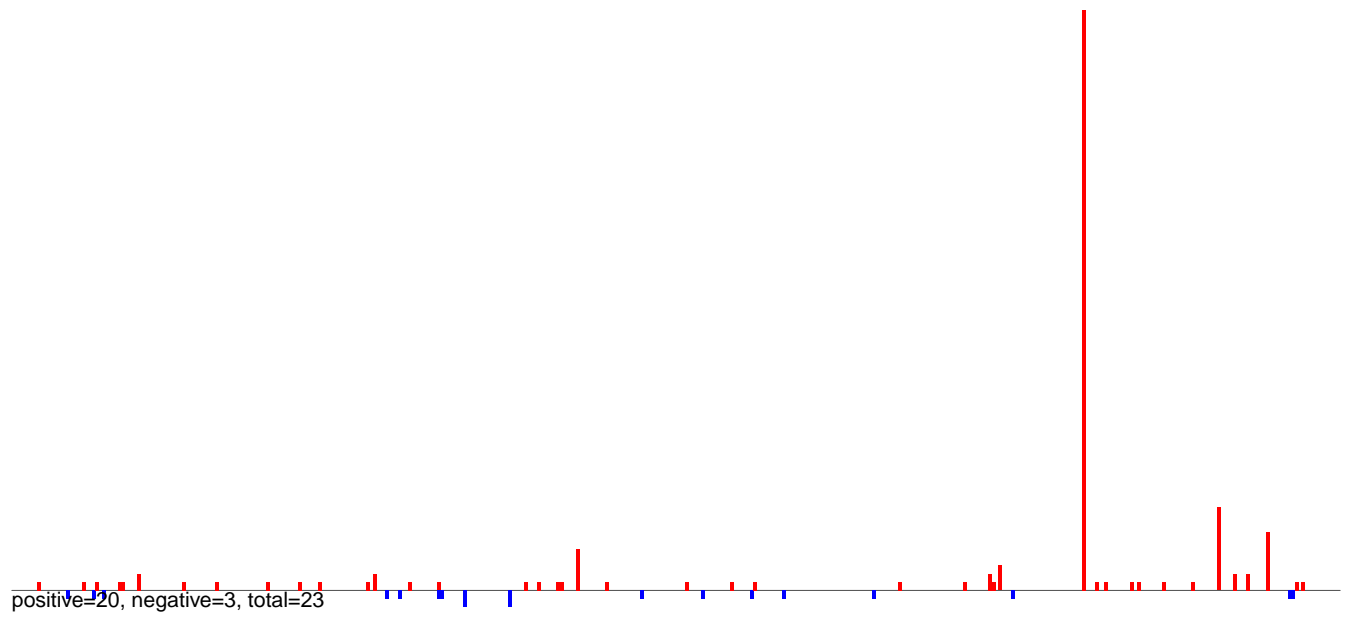
AeAeg_LVP_pMos3DBhspPBac.rep



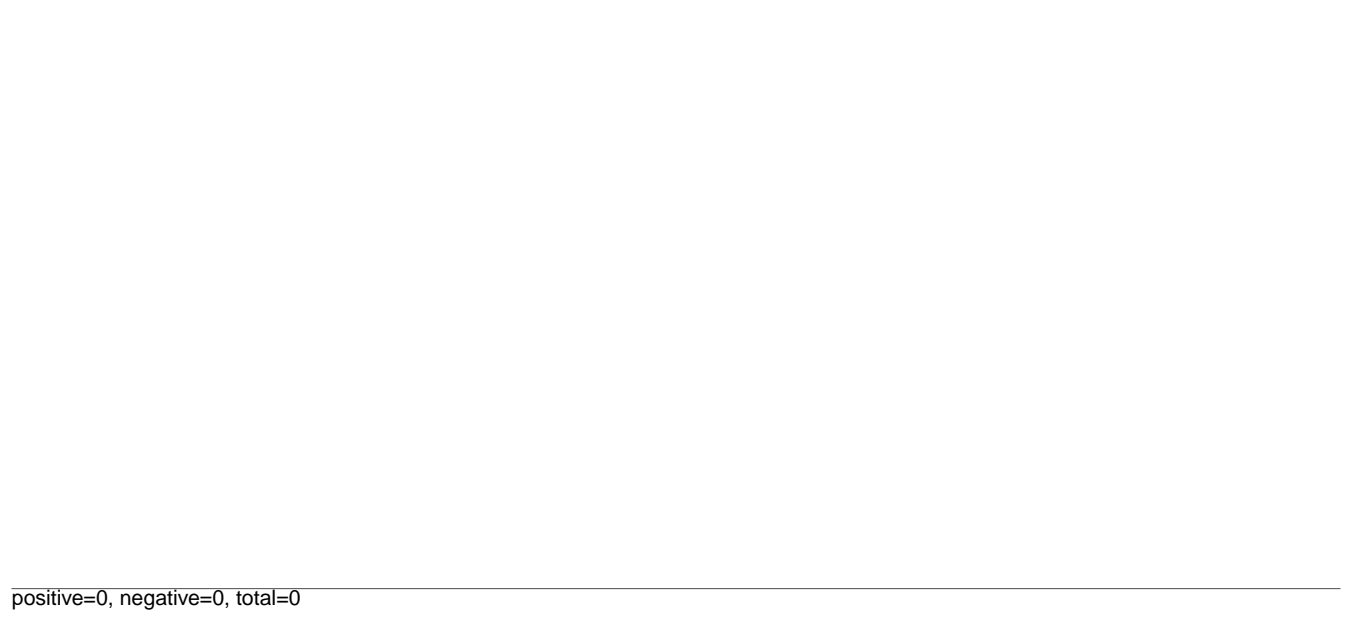
Cheng_Mosquito_Virus_strain_XC1-6_ORF1_ORF1_ORF2_ORF2_glycoprotein_G_and_RNA-de:1-12774

0 2000 4000 6000 8000 10000 12000

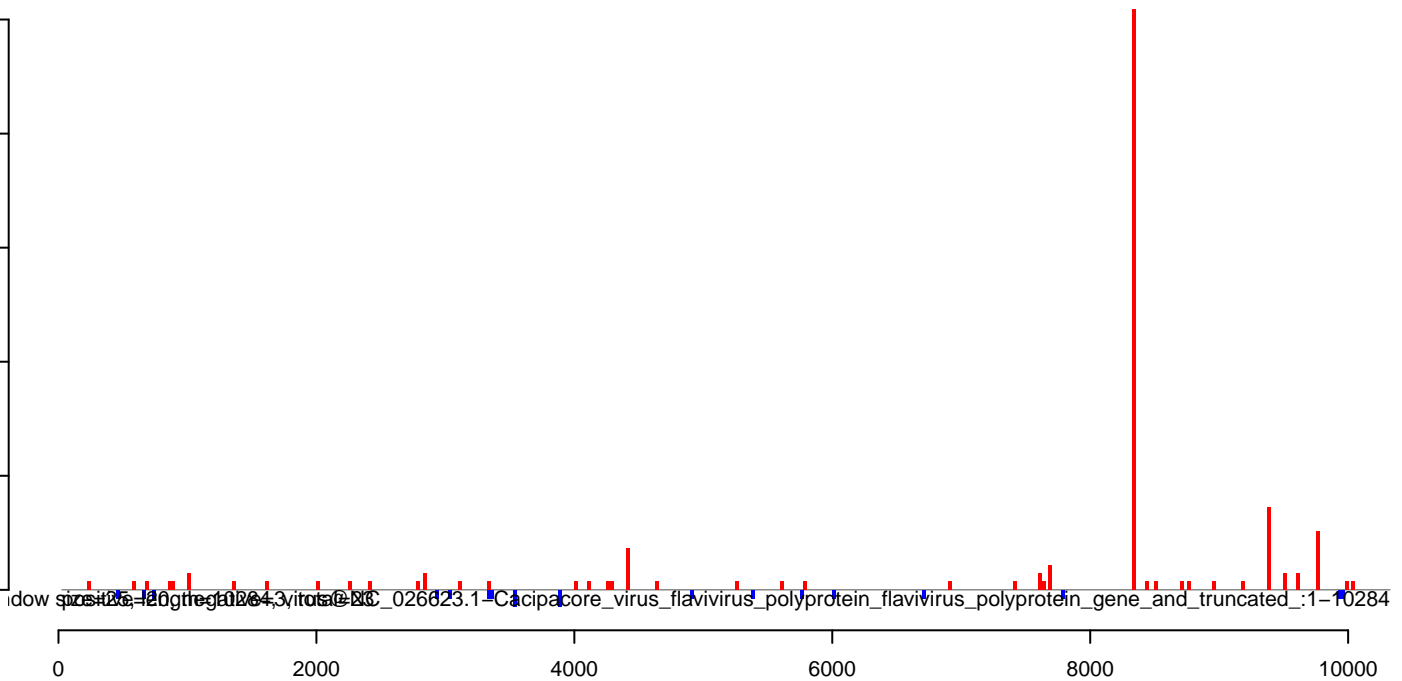
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



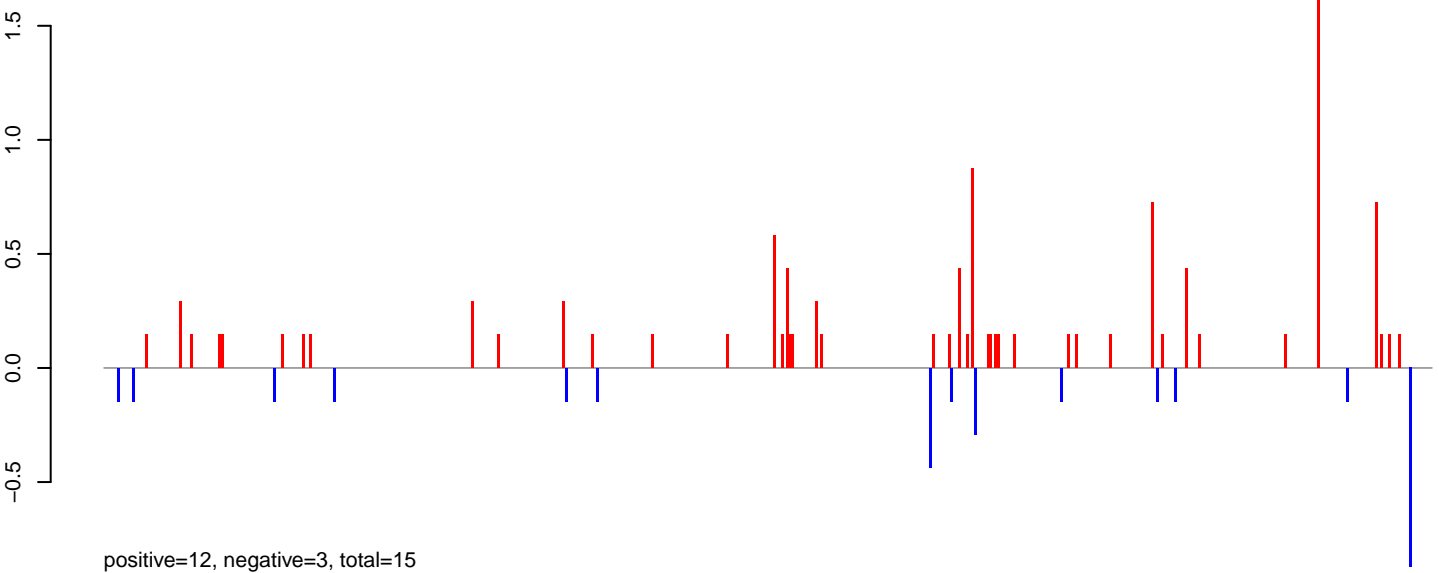
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



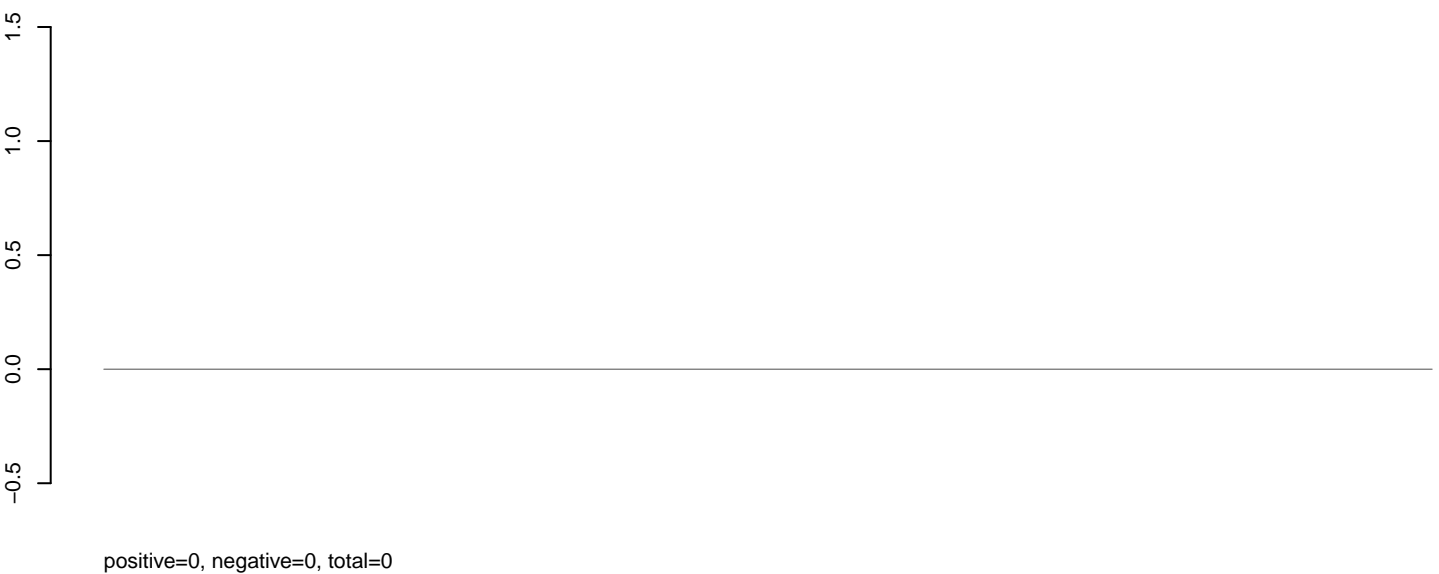
AeAeg_LVP_pMos3DBhspPBac.rep



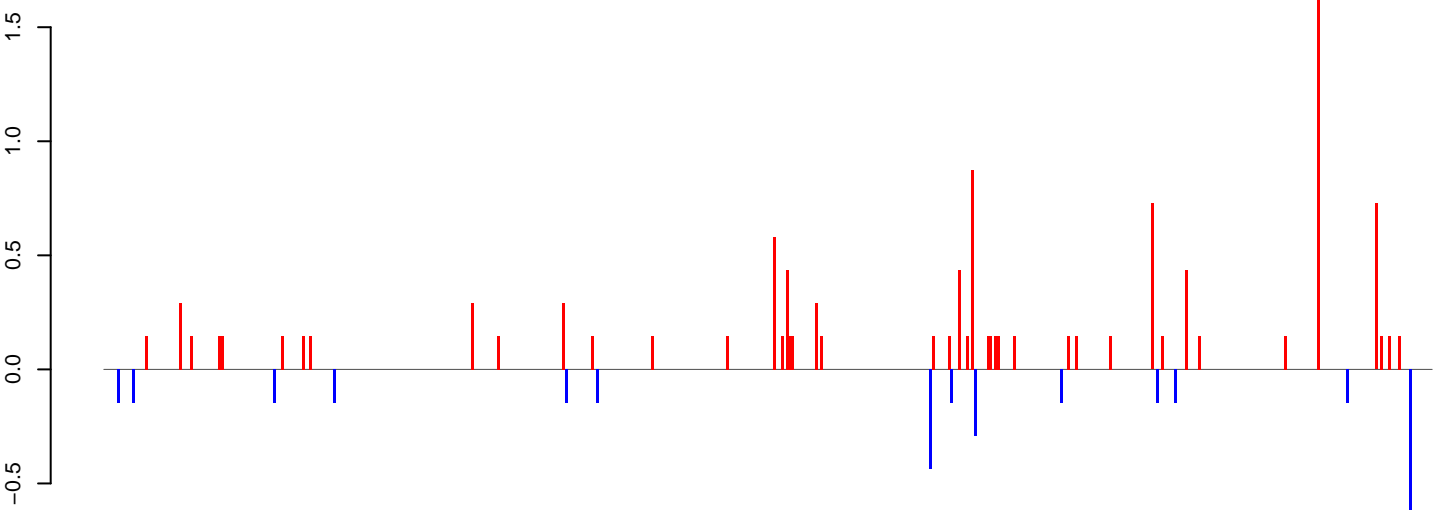
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



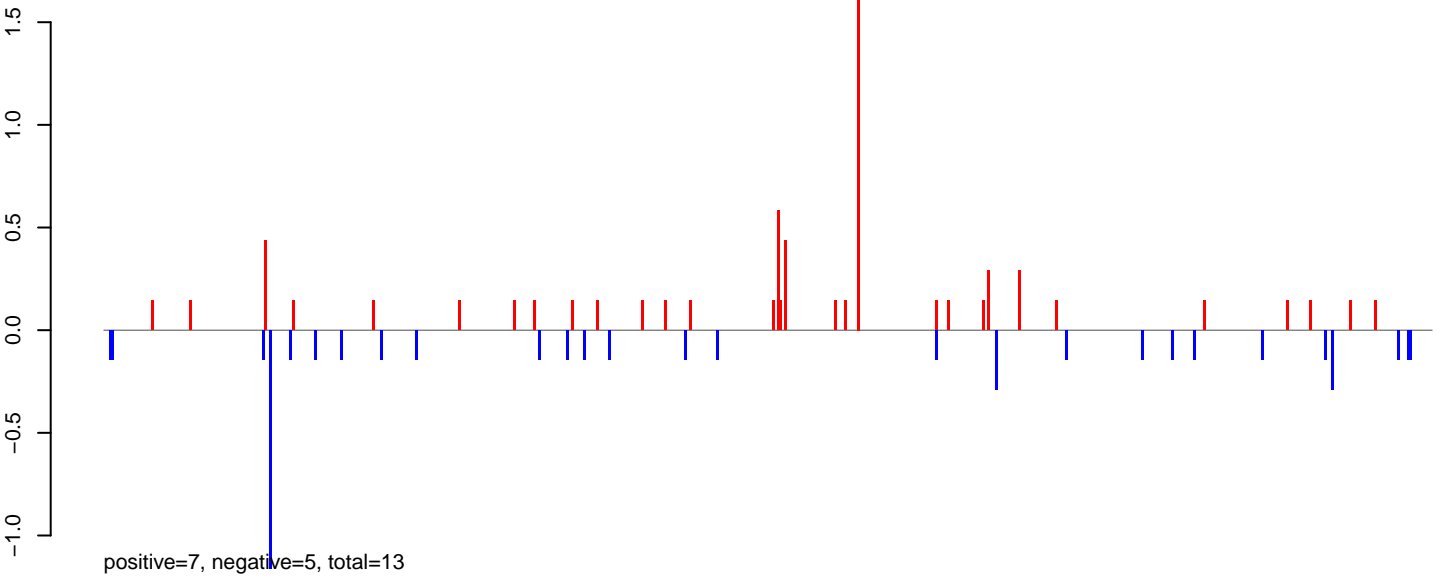
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



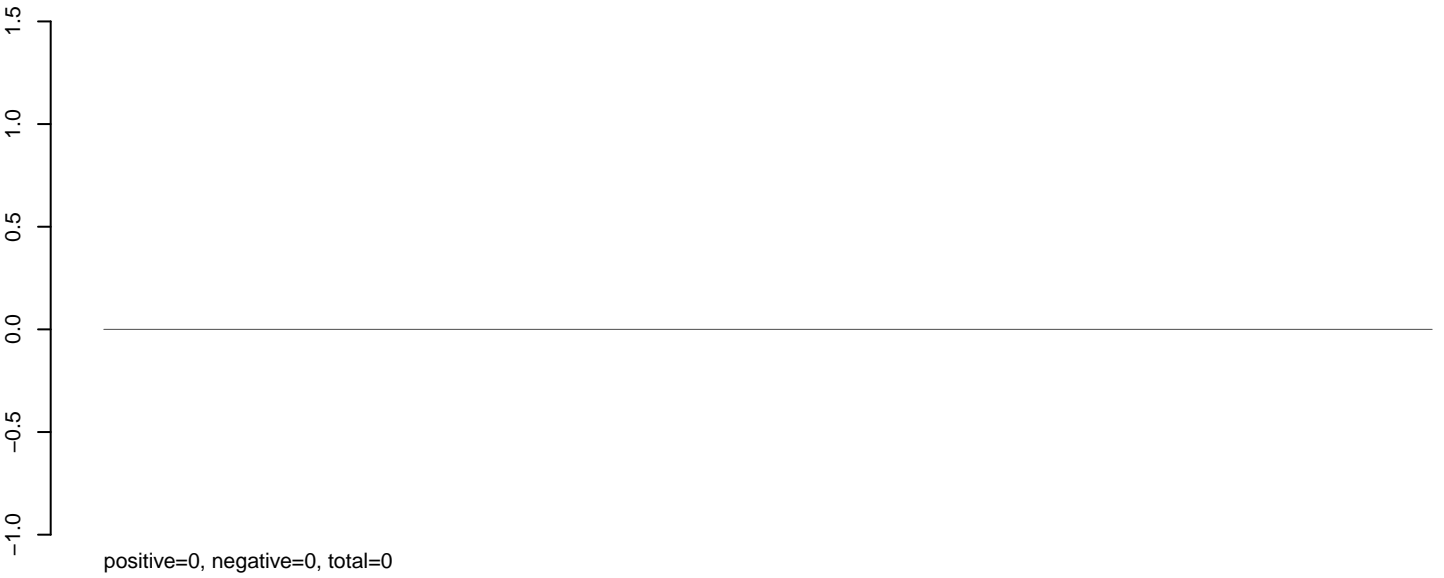
AeAeg_LVP_pMos3DBhspPBac.rep



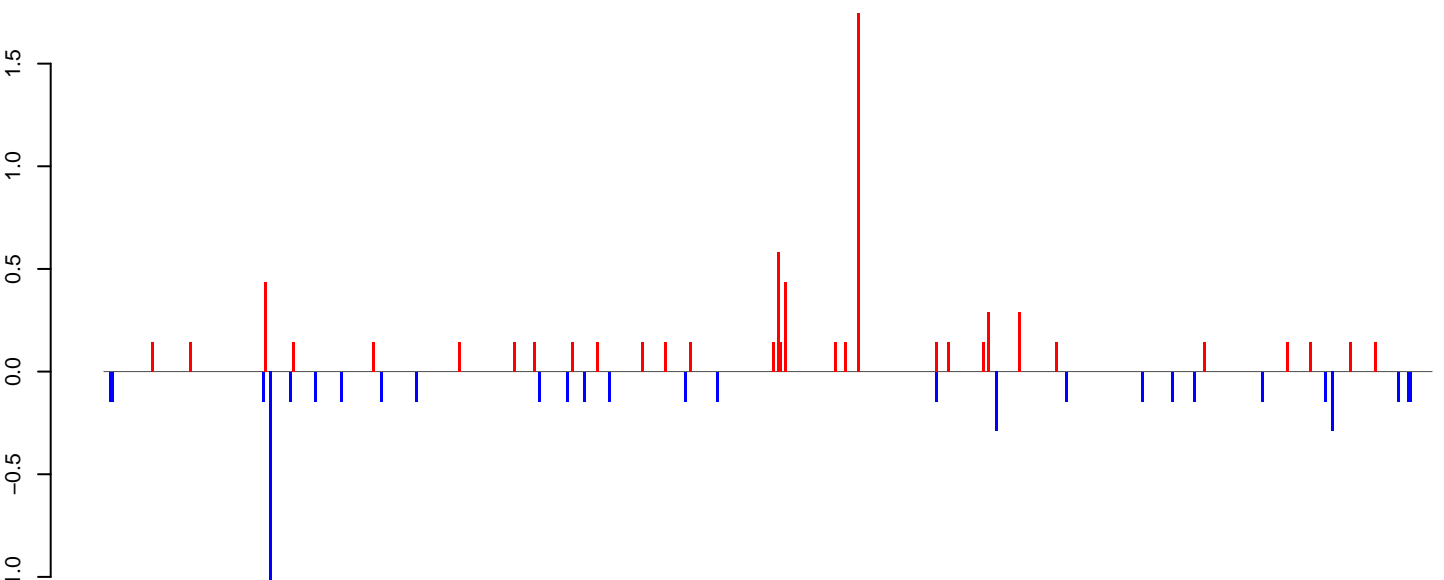
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



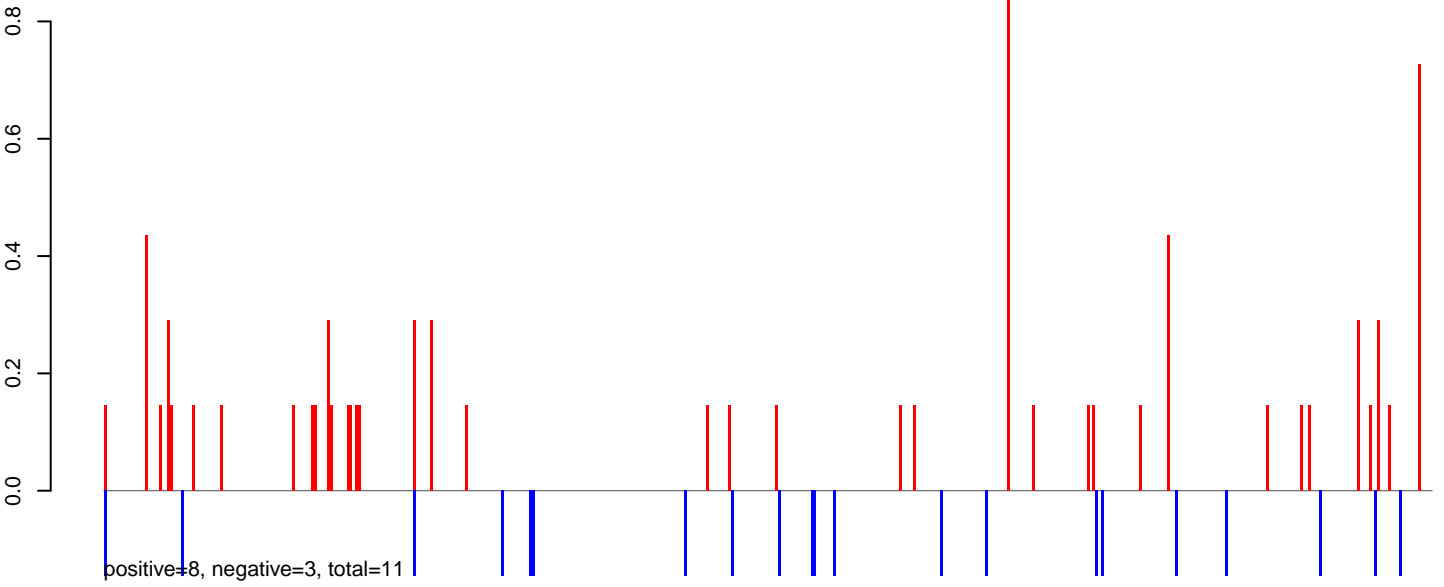
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



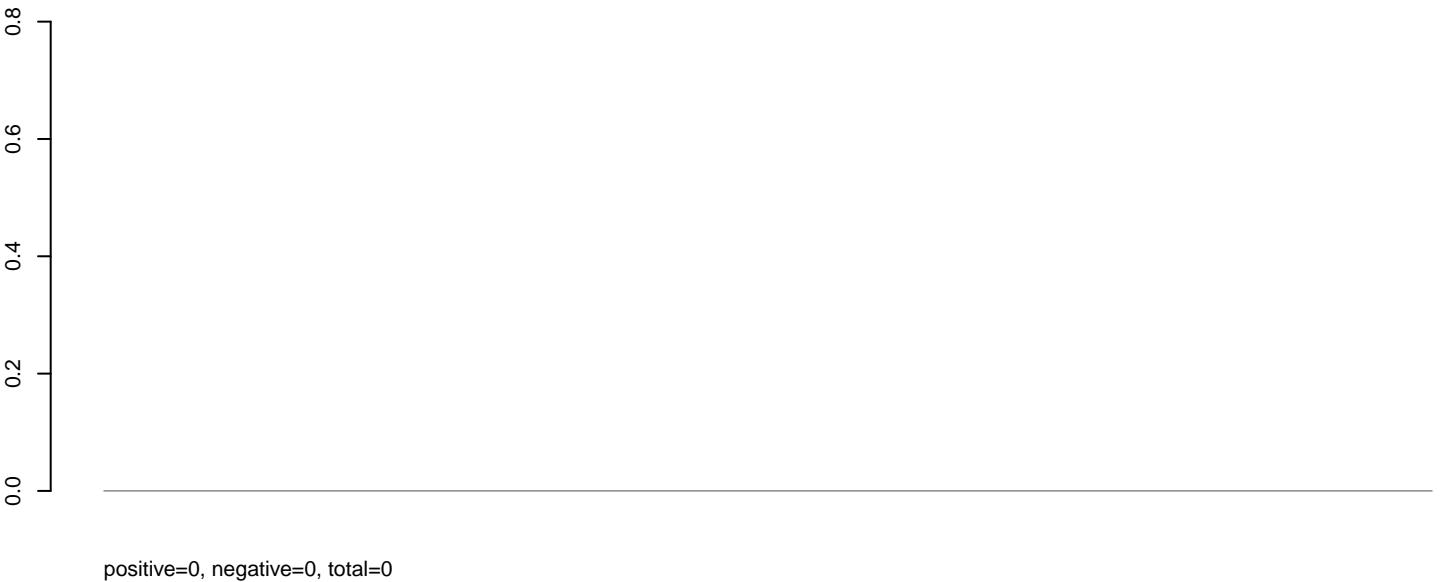
AeAeg_LVP_pMos3DBhspPBac.rep



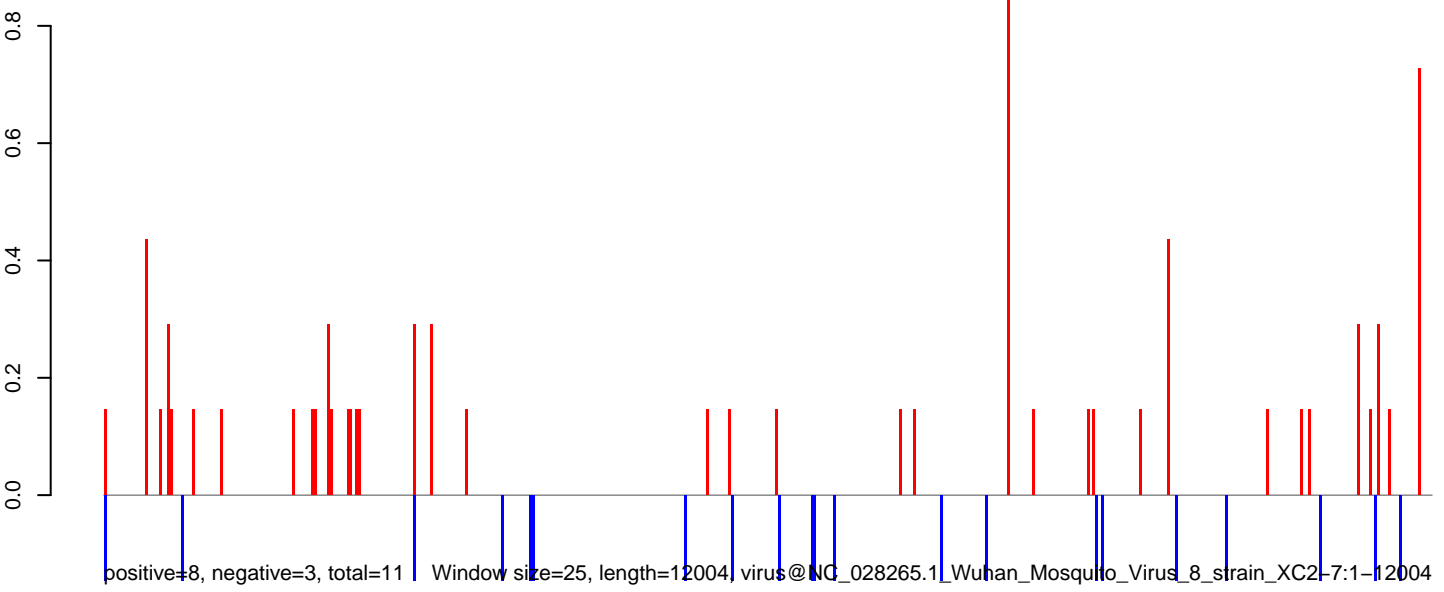
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

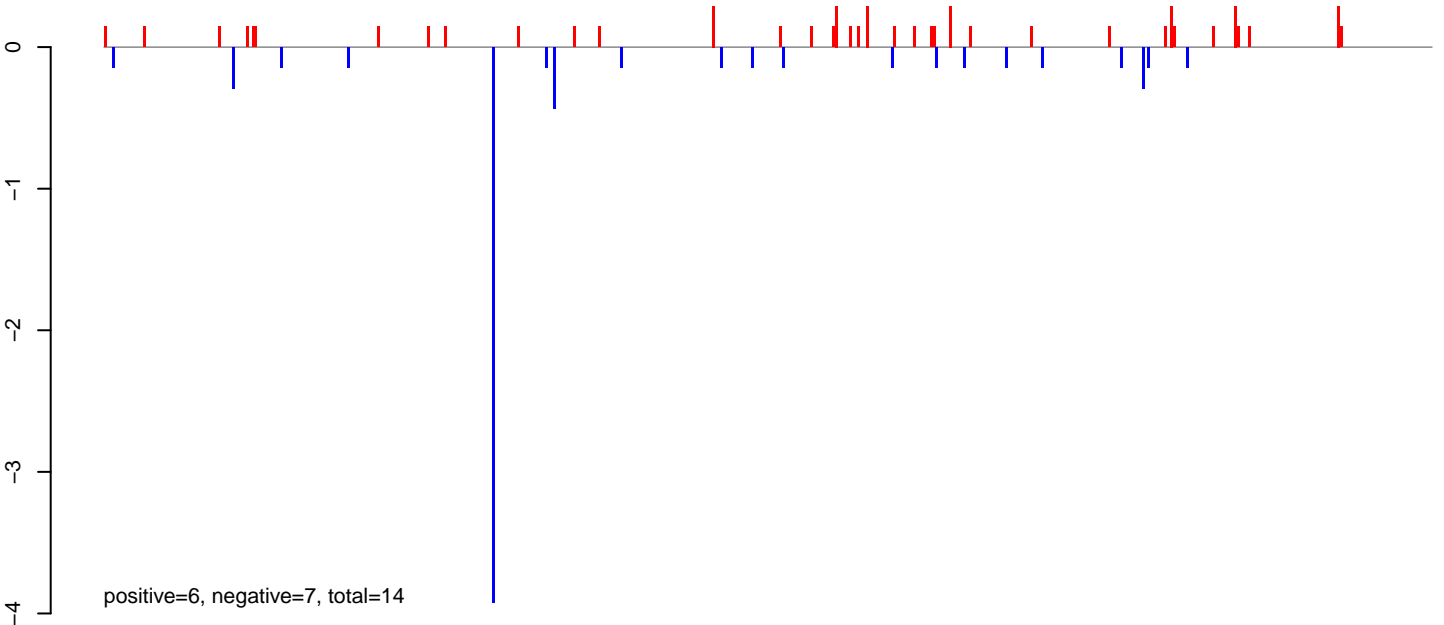


AeAeg_LVP_pMos3DBhspPBac.rep

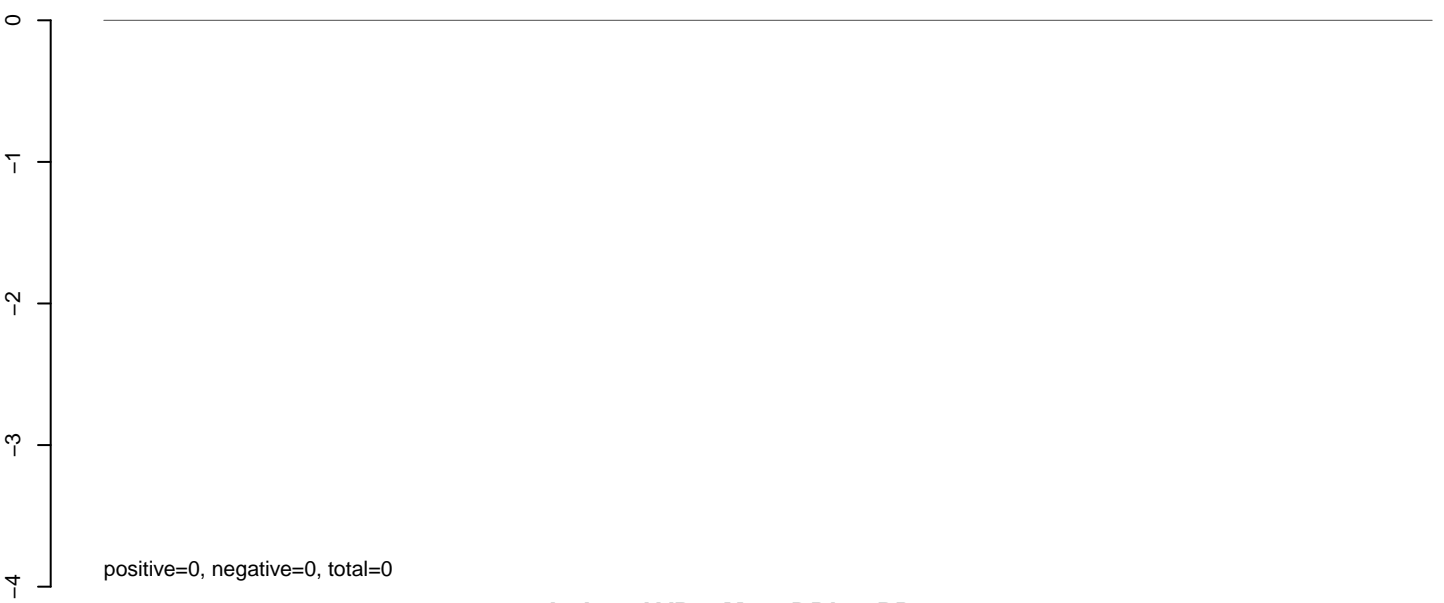


Window size=25, length=12004, virus@NC_028265.1_Wuhan_Mosquito_Virus_8_strain_XC2-7:1-12004

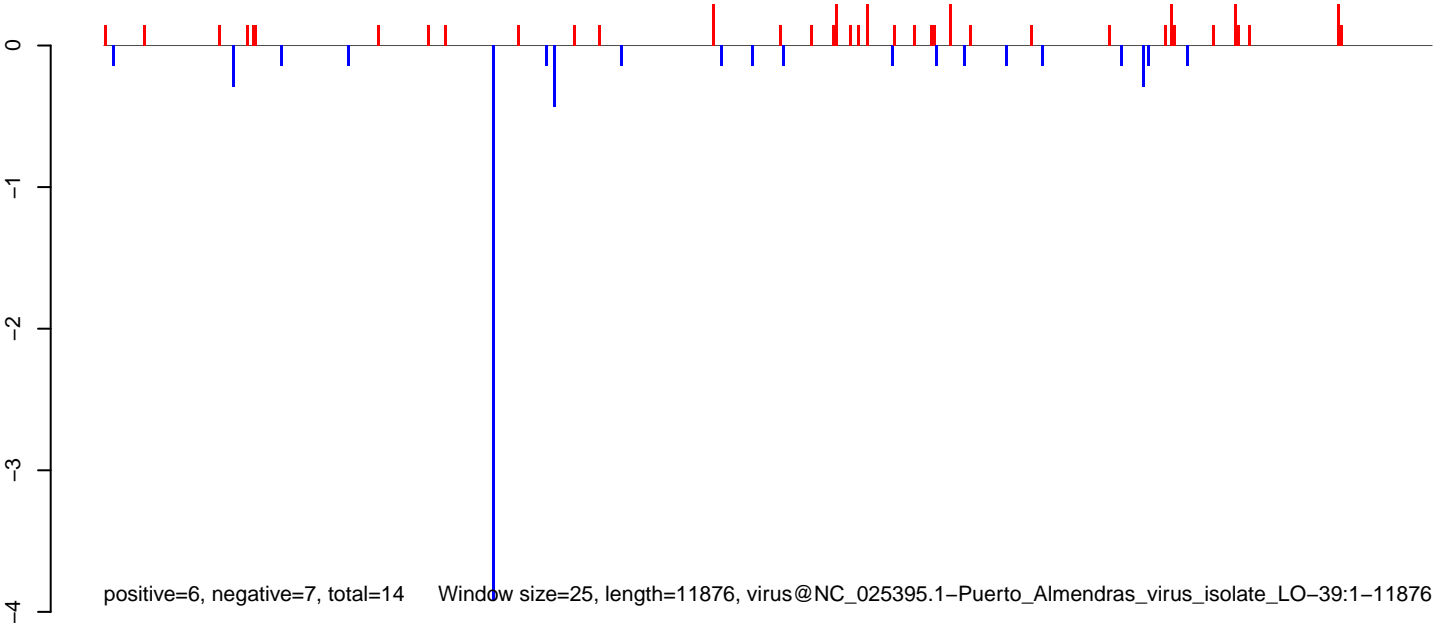
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



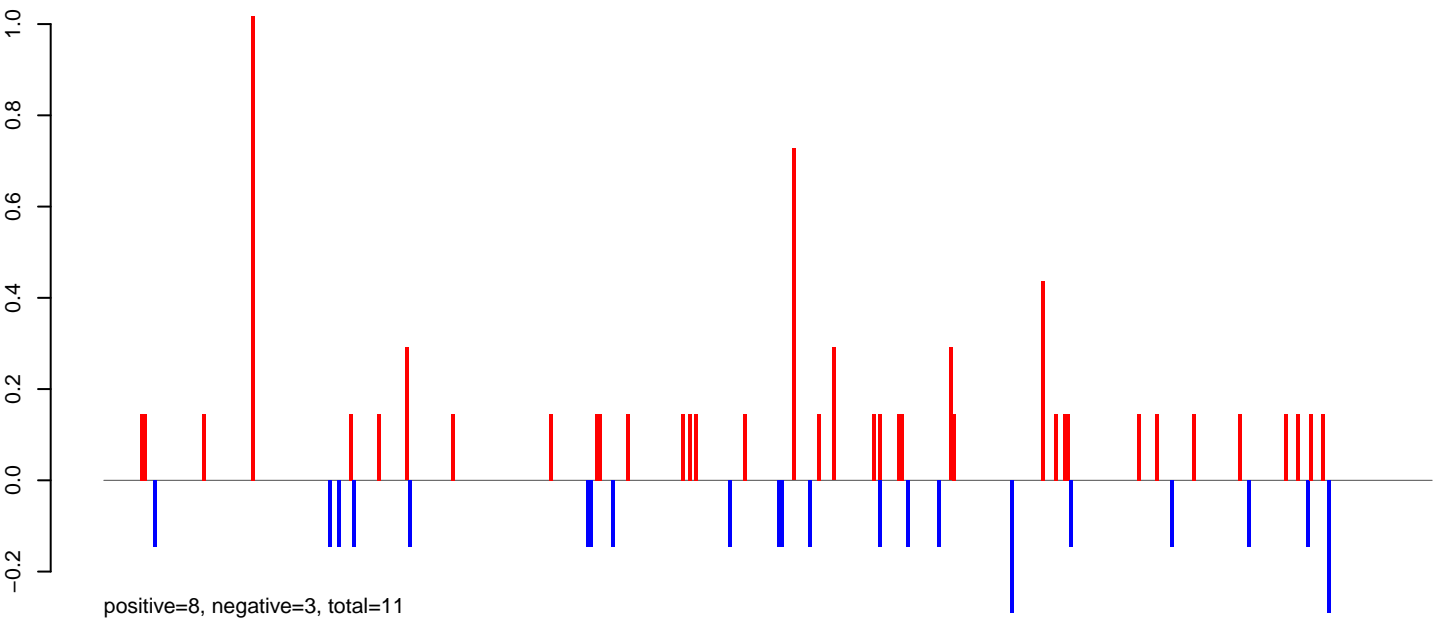
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



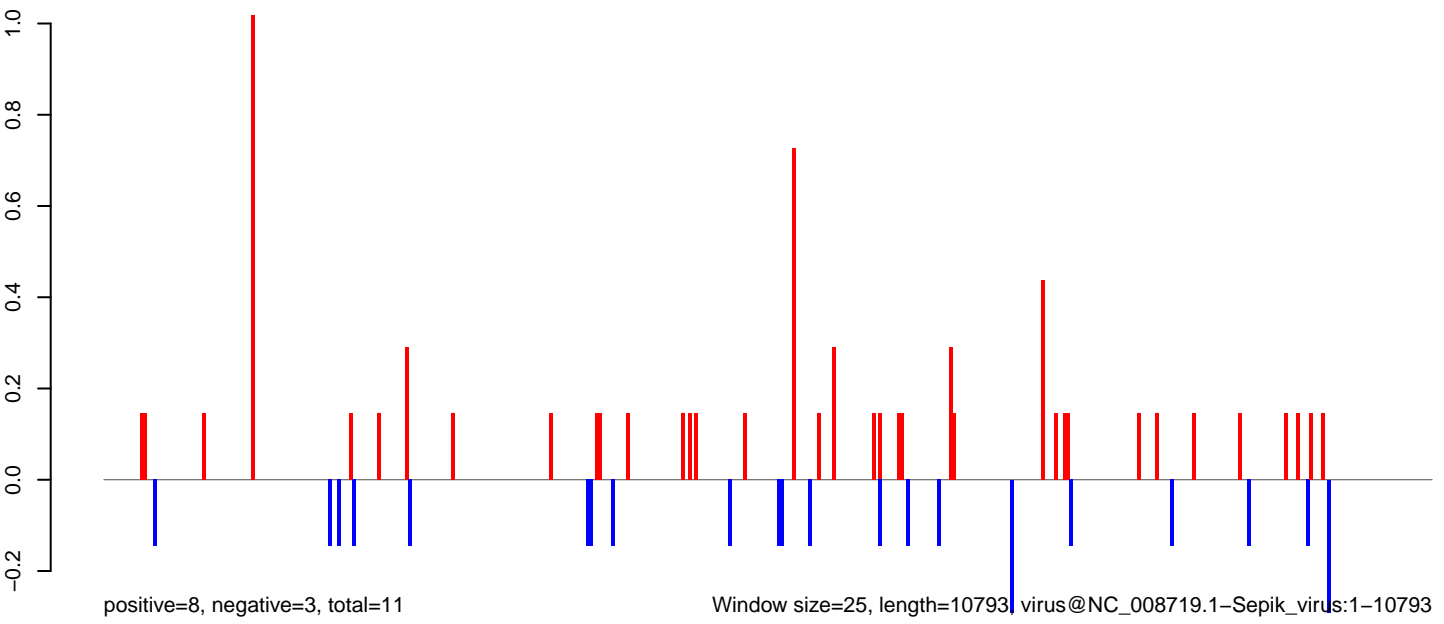
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



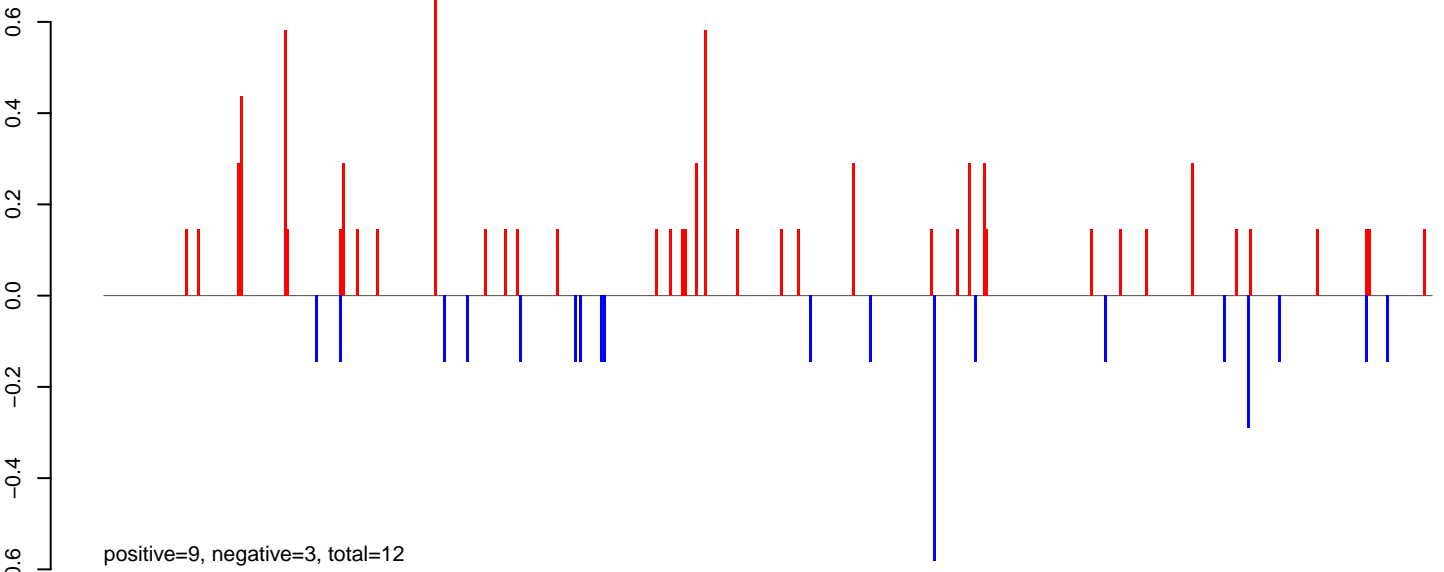
AeAeg_LVP_pMos3DBhspPBac.rep



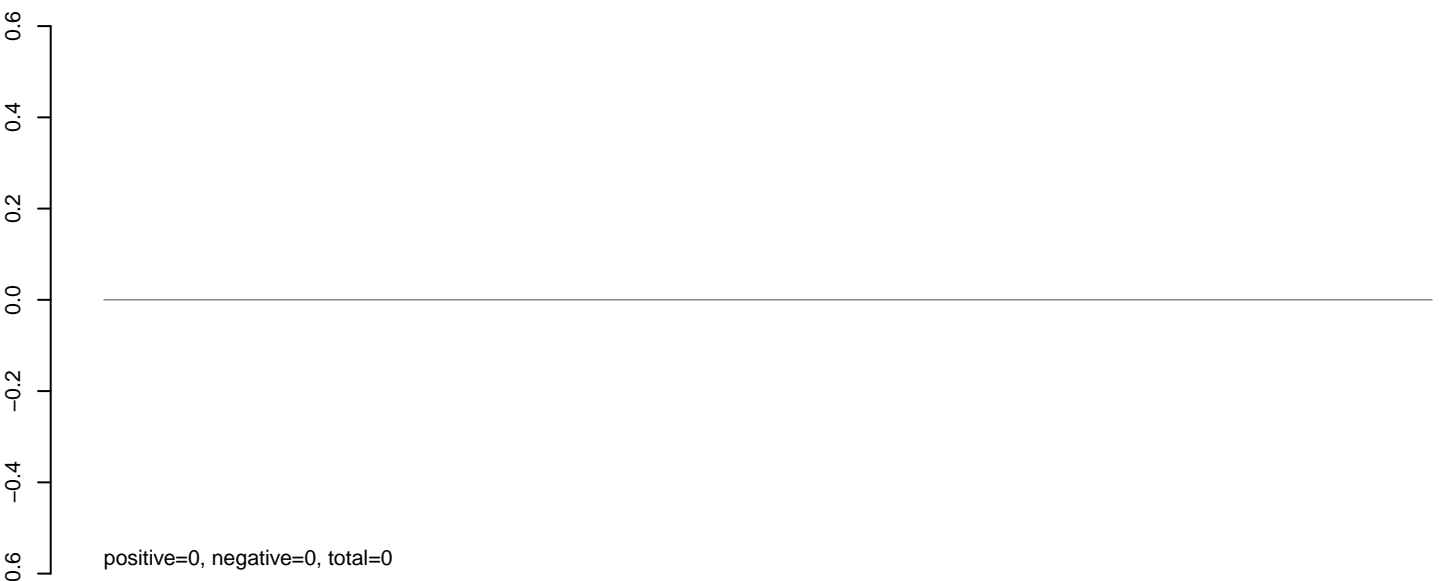
Window size=25, length=10793, virus@NC_008719.1-Sepik_virus:1-10793

0 2000 4000 6000 8000 10000

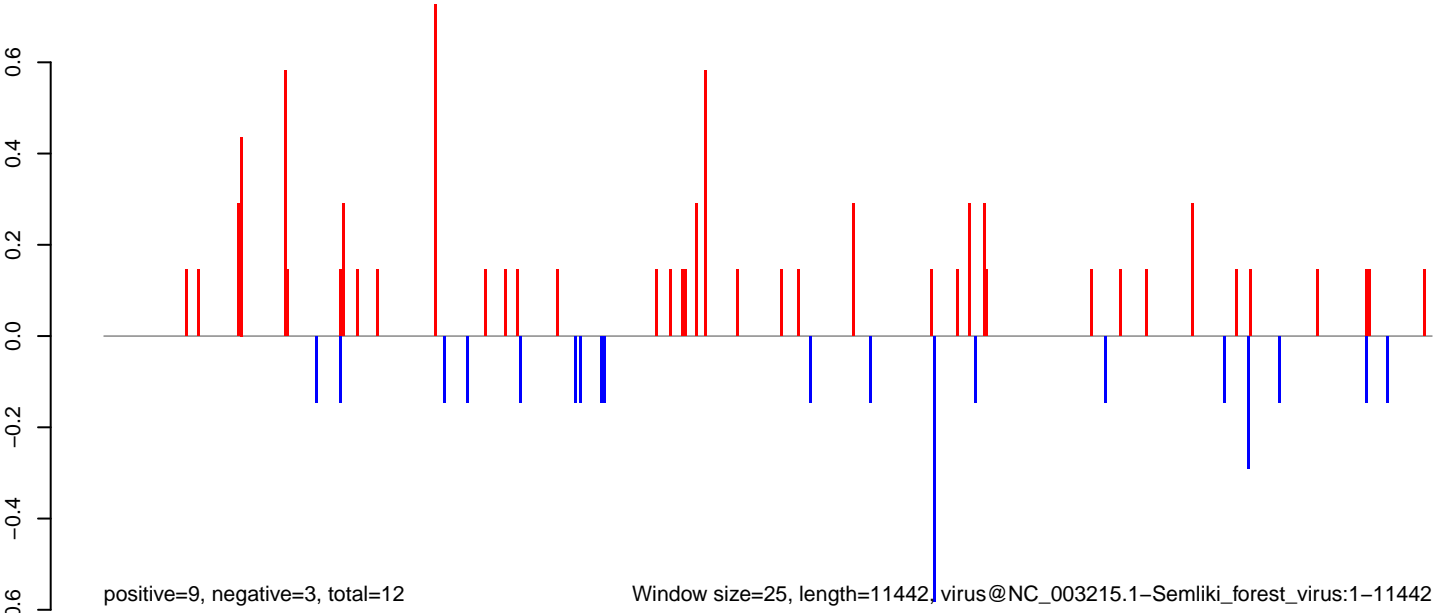
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

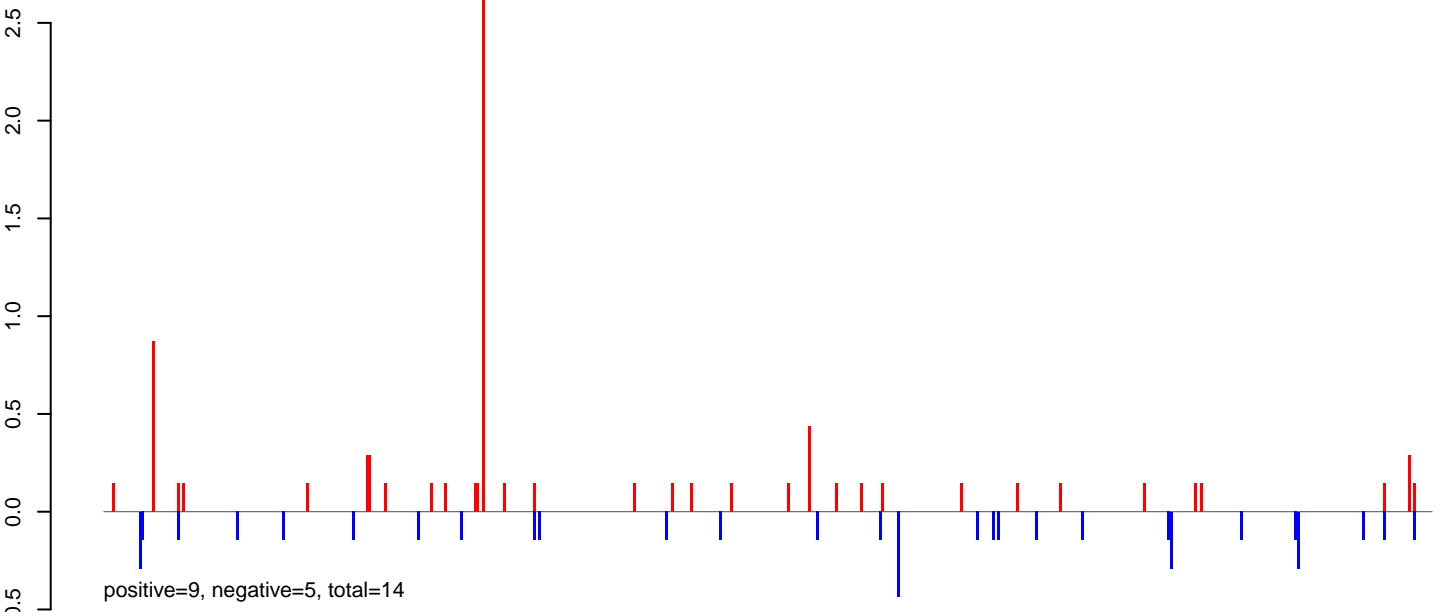


AeAeg_LVP_pMos3DBhspPBac.rep

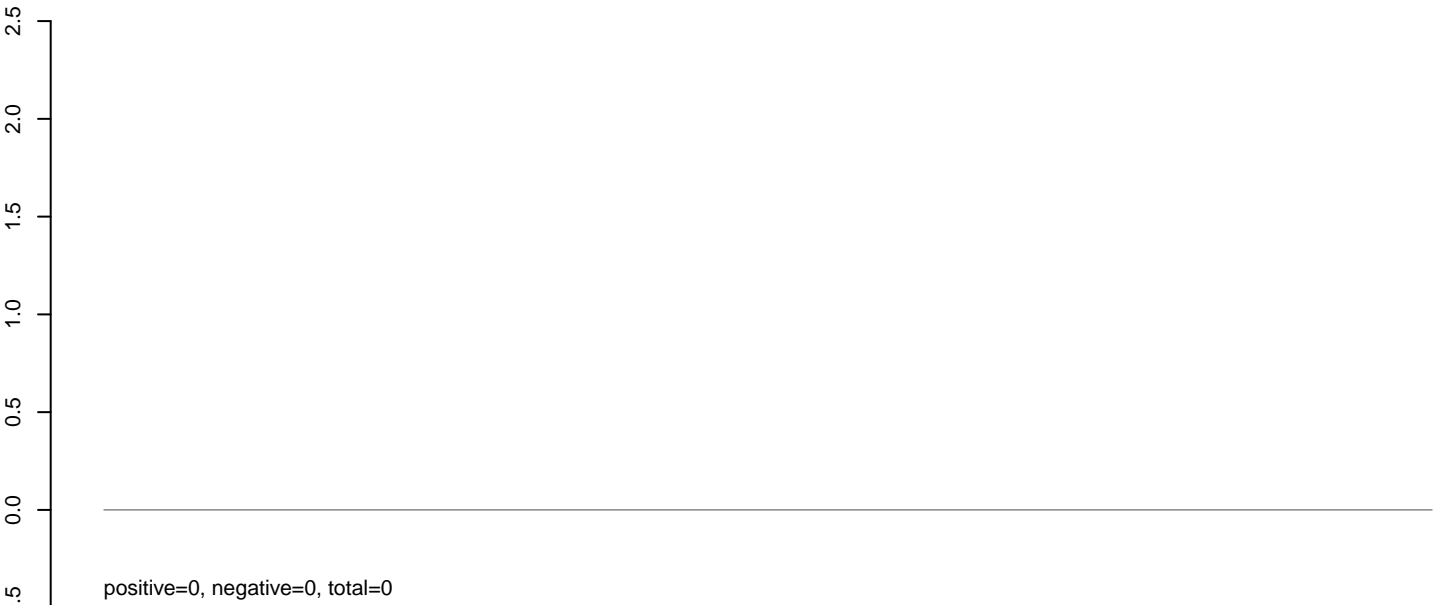


0 2000 4000 6000 8000 10000

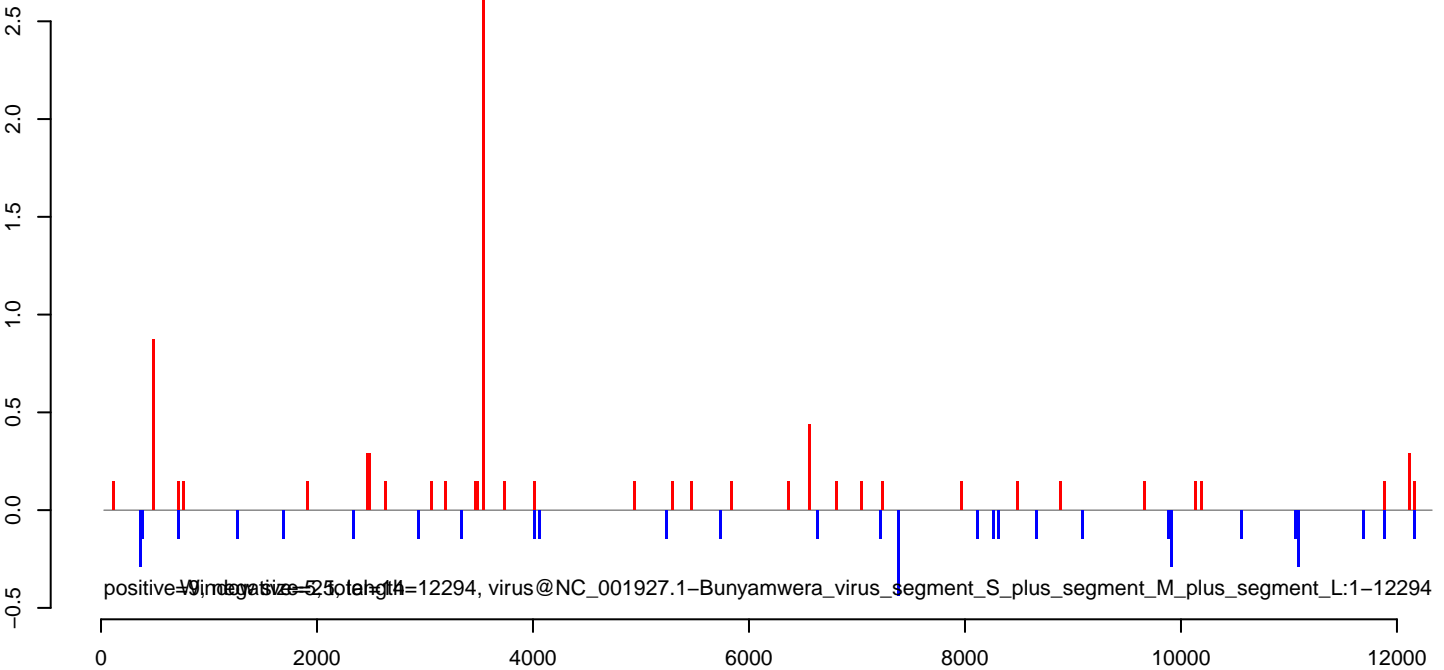
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



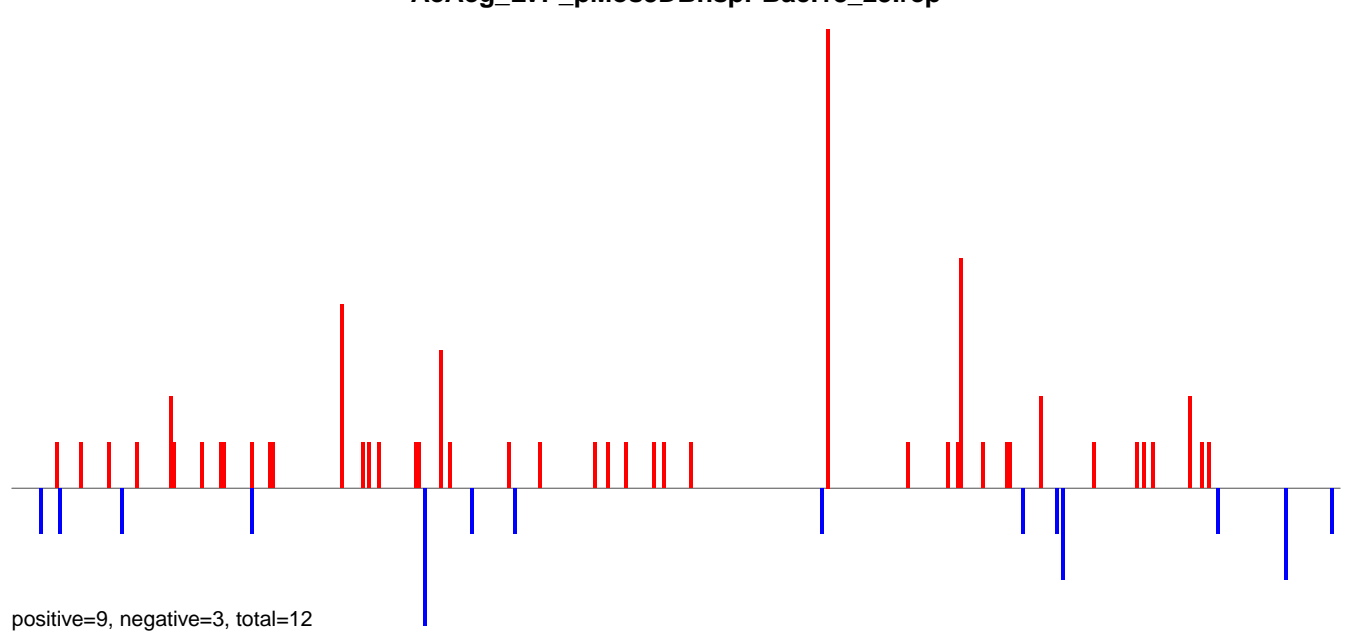
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



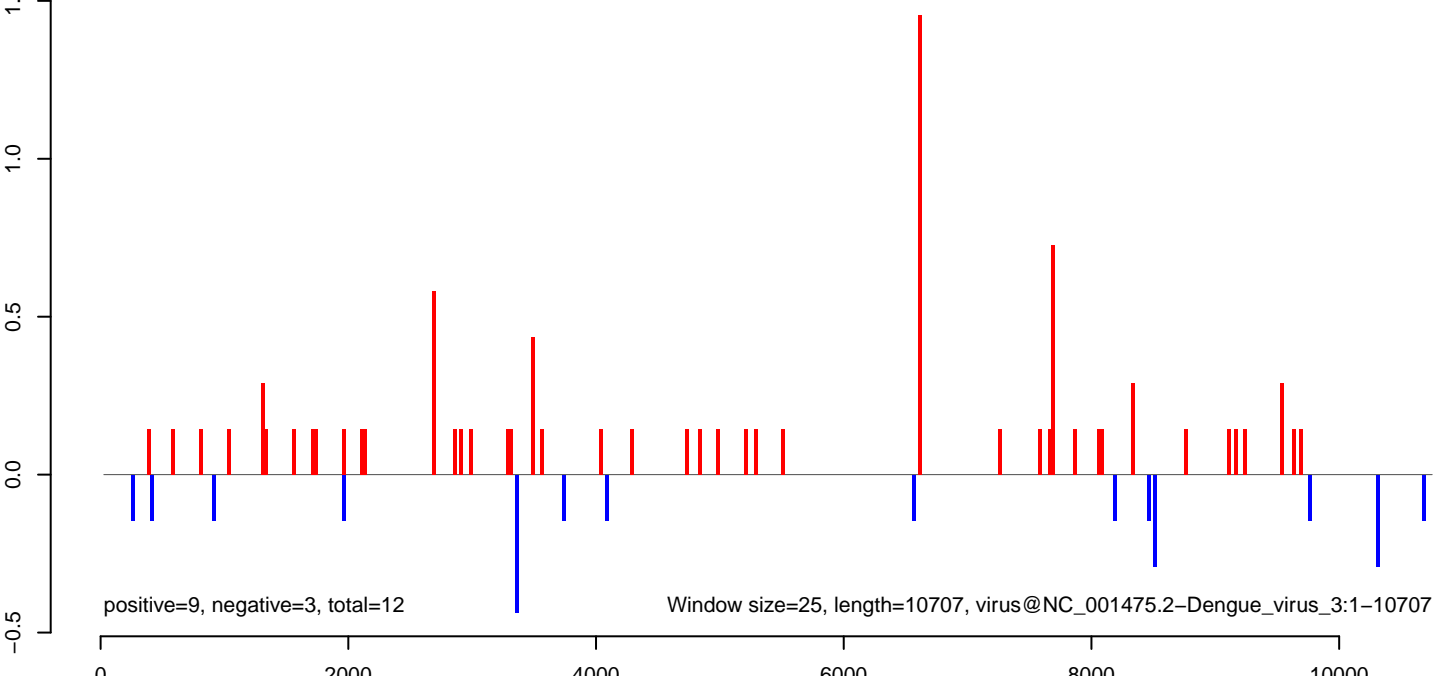
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



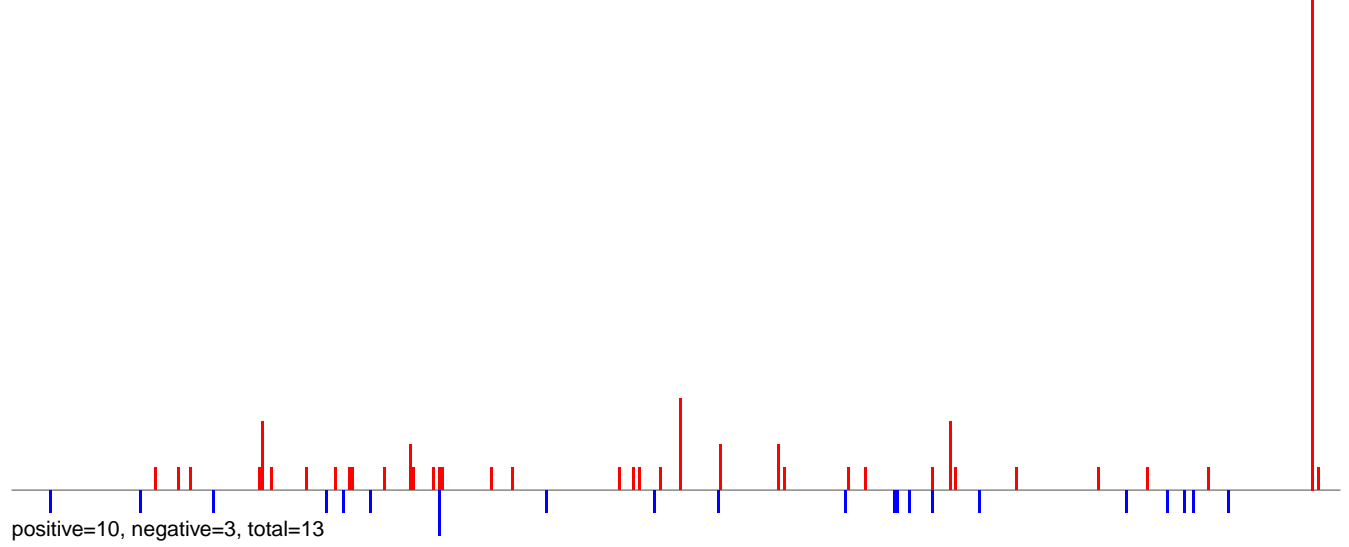
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



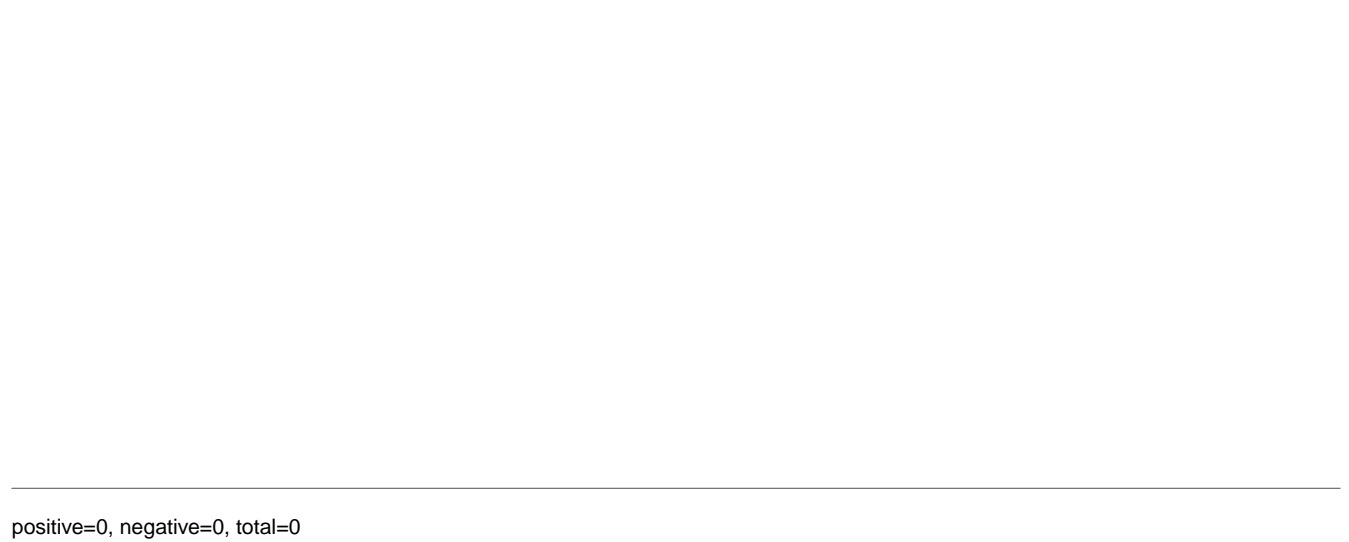
AeAeg_LVP_pMos3DBhspPBac.rep



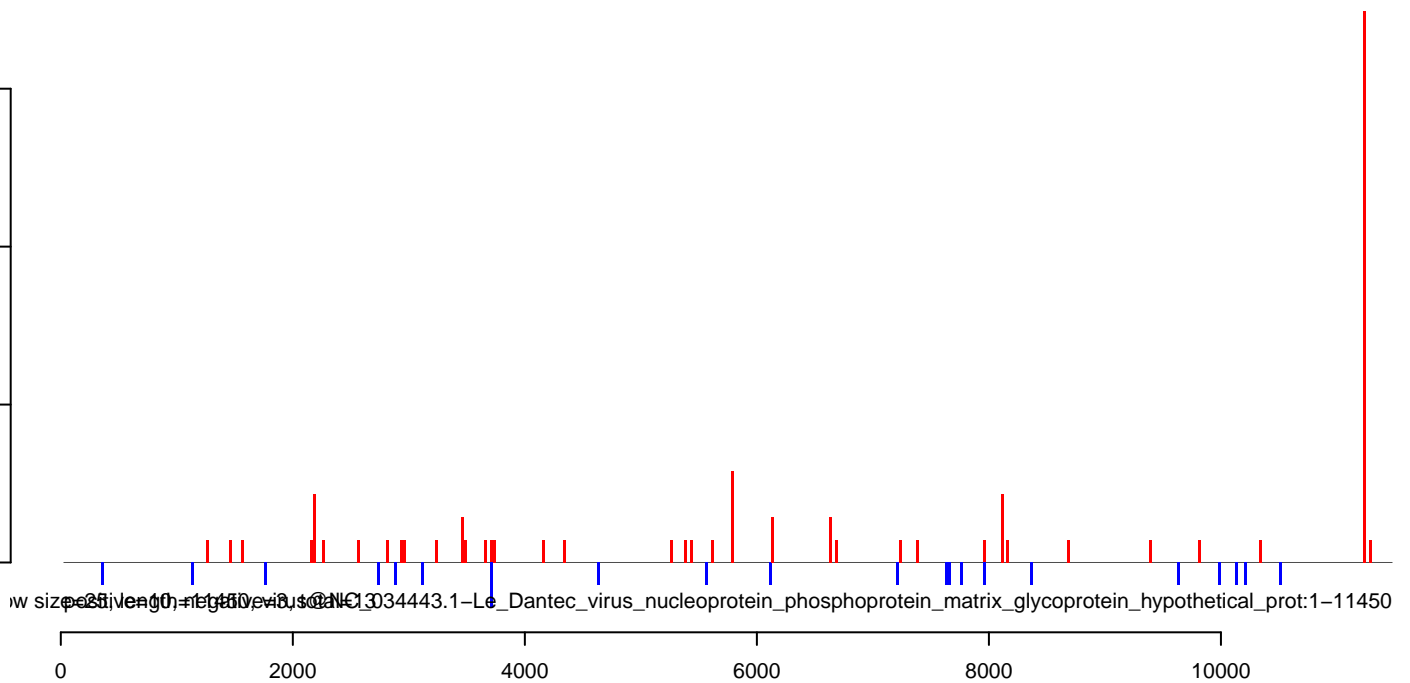
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



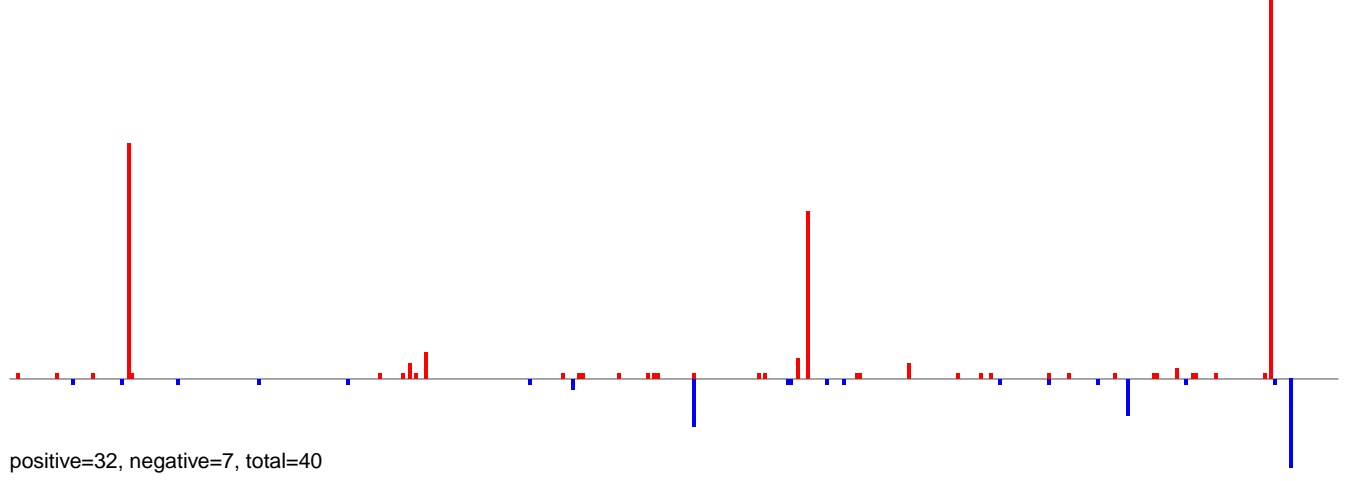
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



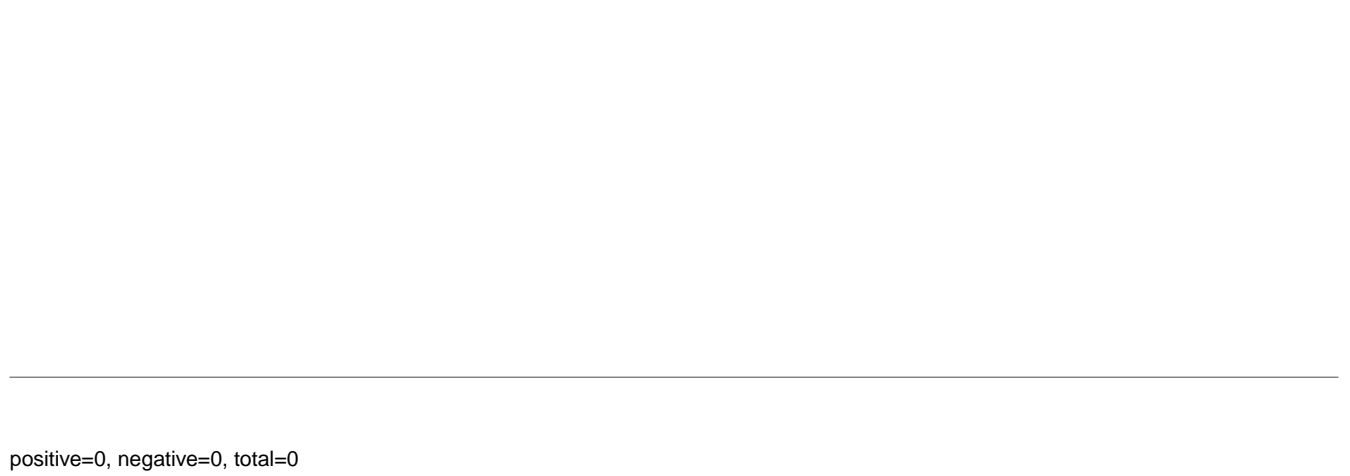
AeAeg_LVP_pMos3DBhspPBac.rep



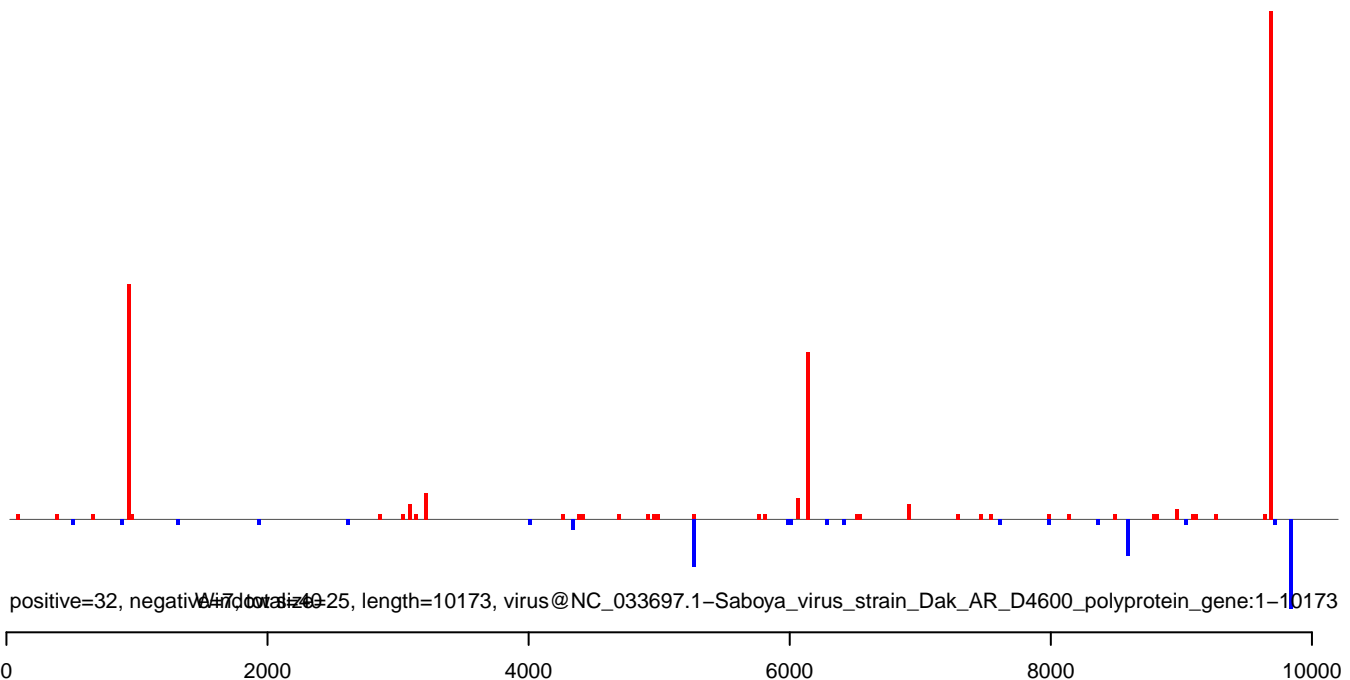
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



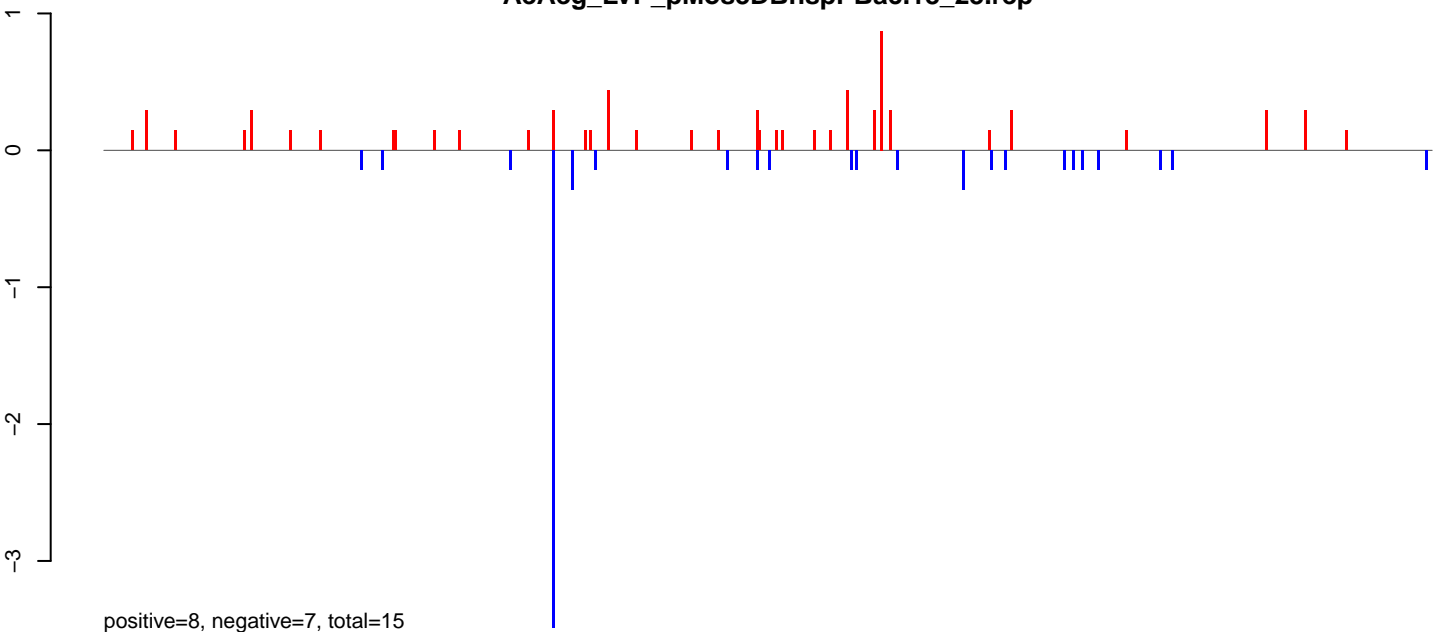
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



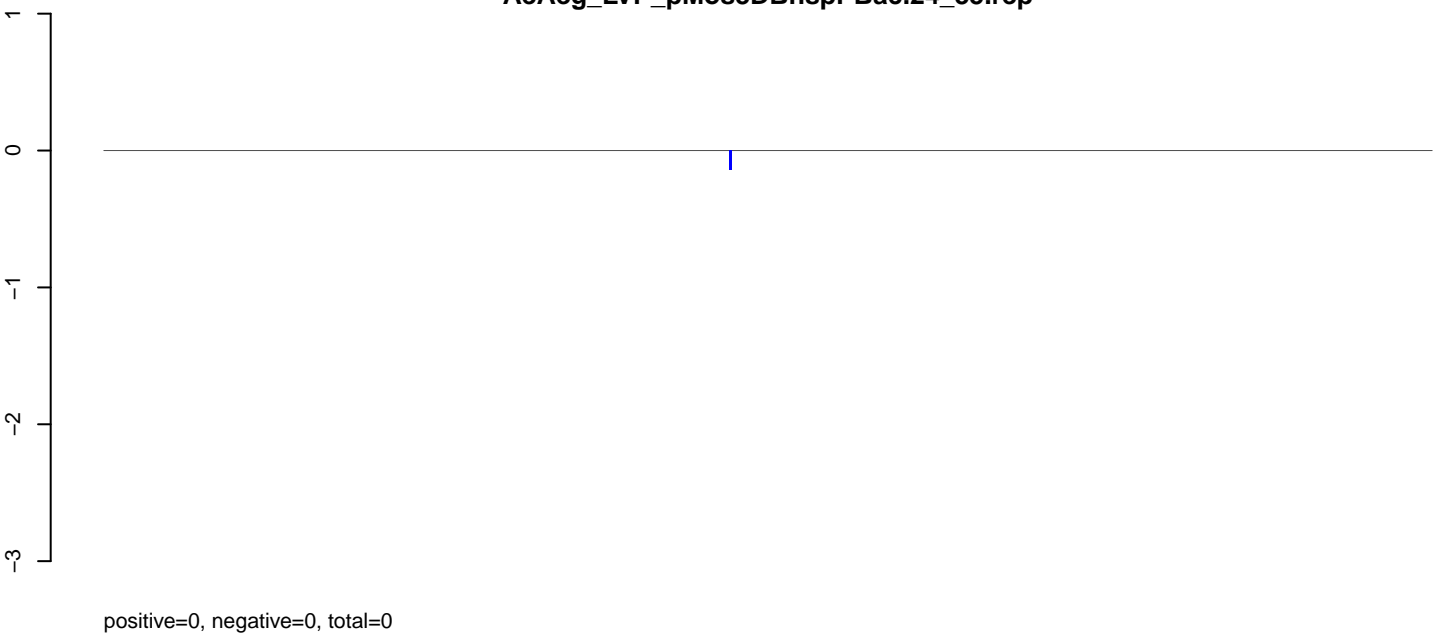
AeAeg_LVP_pMos3DBhspPBac.rep



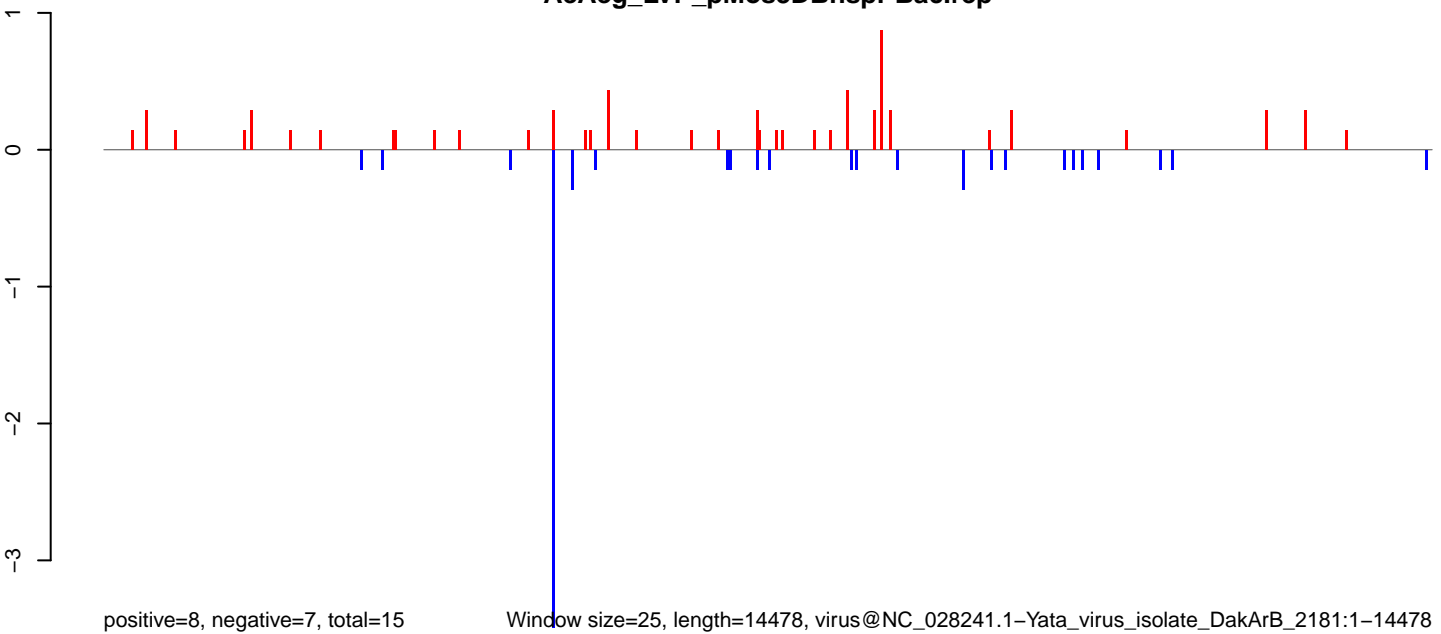
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



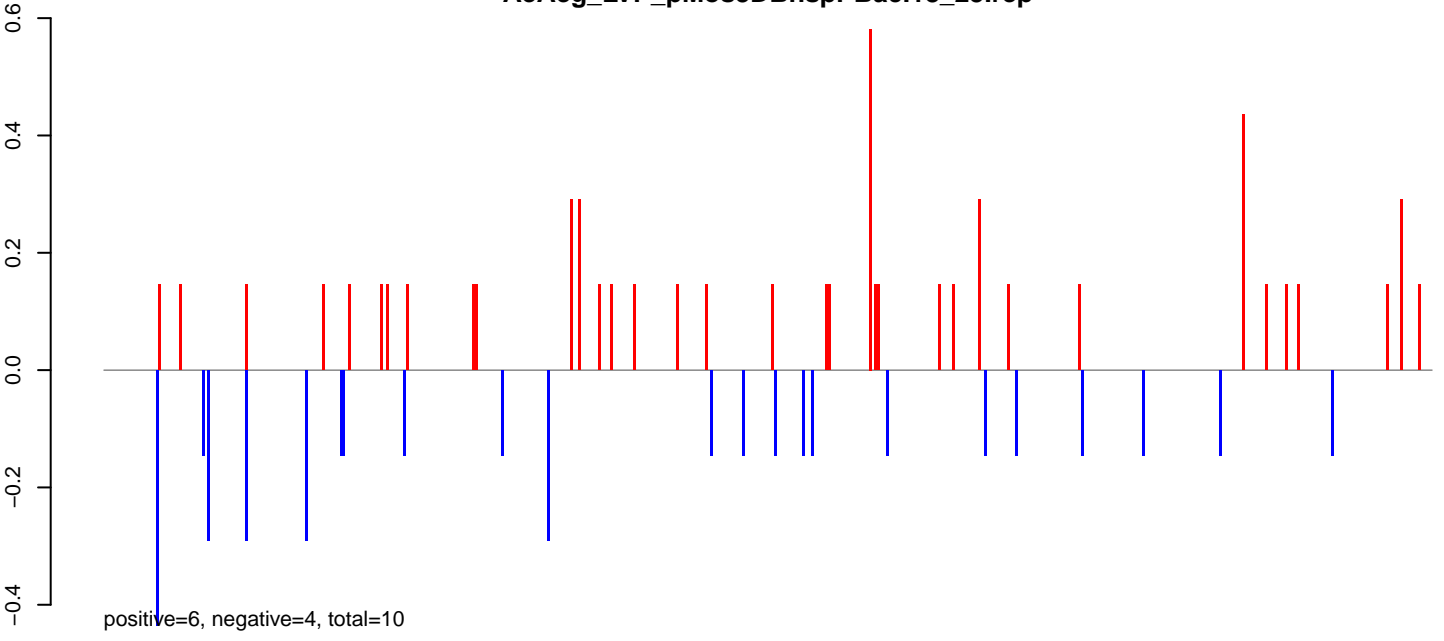
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



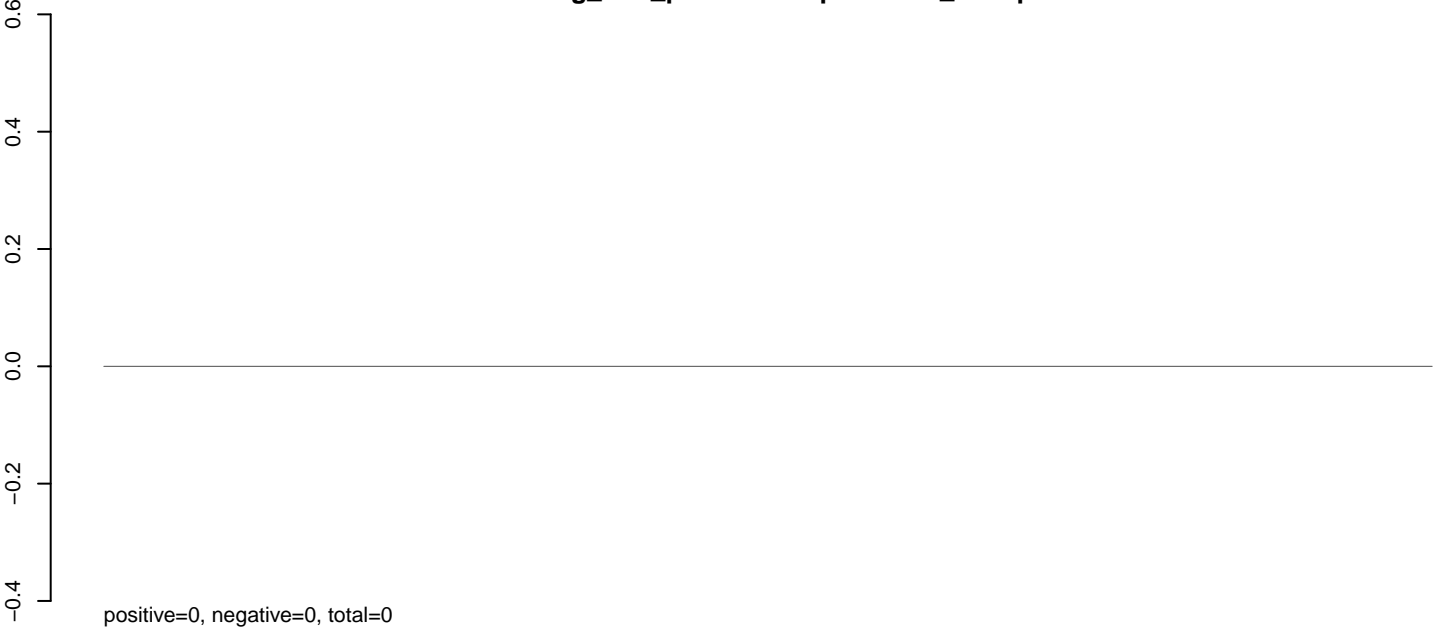
AeAeg_LVP_pMos3DBhspPBac.rep



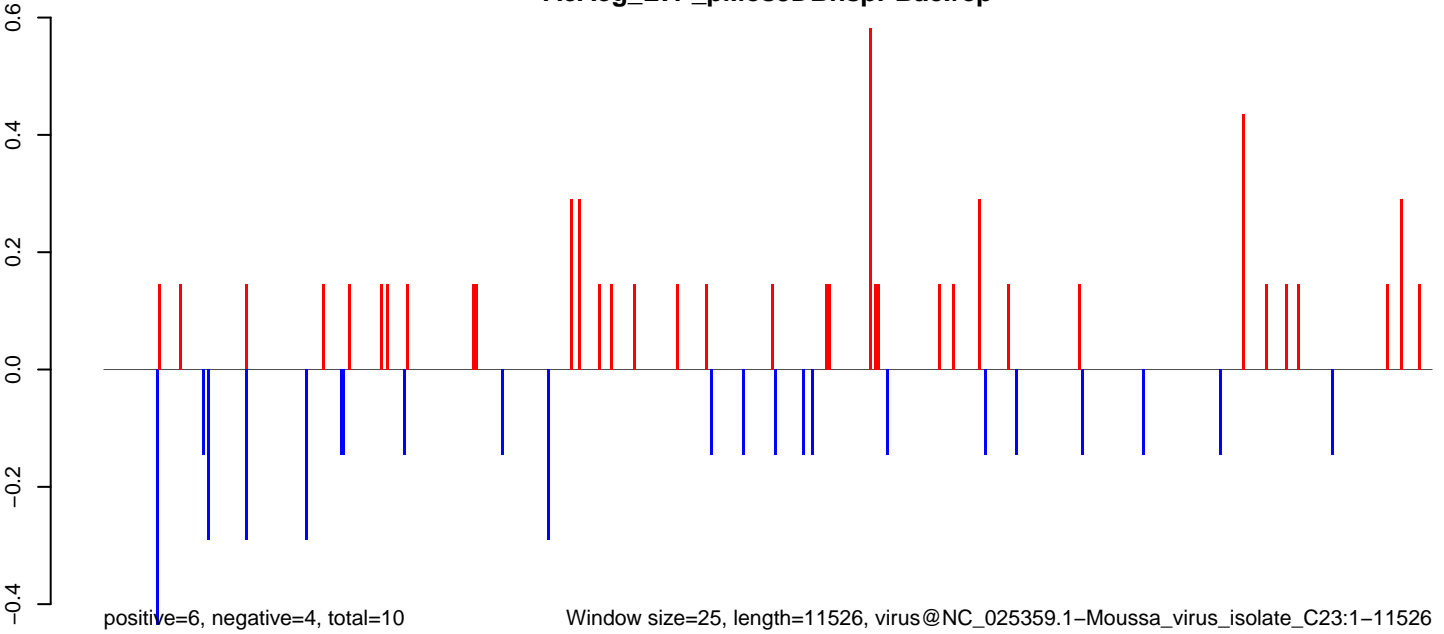
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

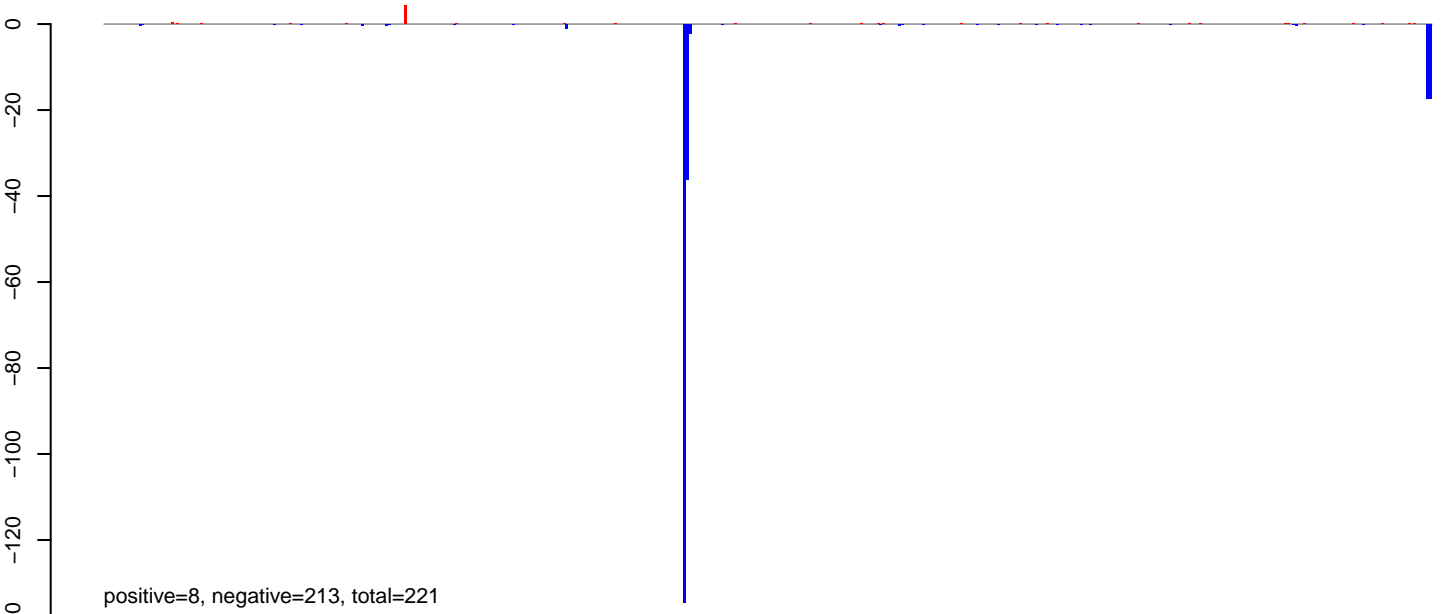


AeAeg_LVP_pMos3DBhspPBac.rep

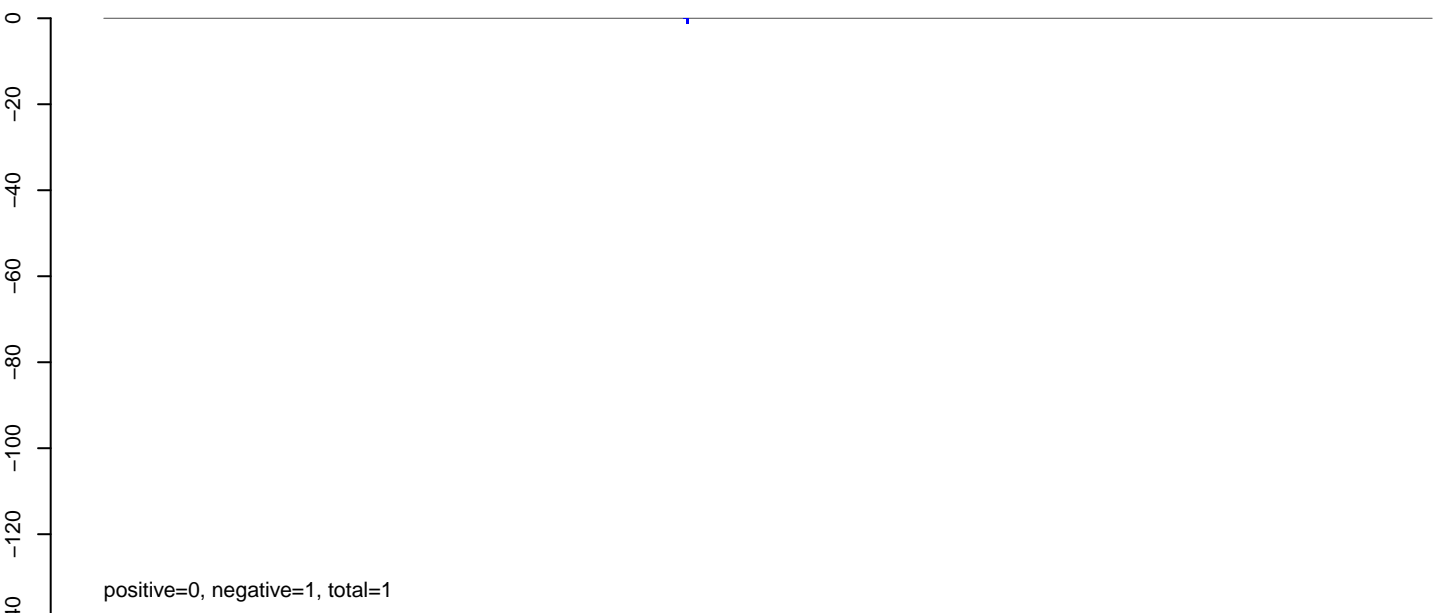


0 2000 4000 6000 8000 10000 12000

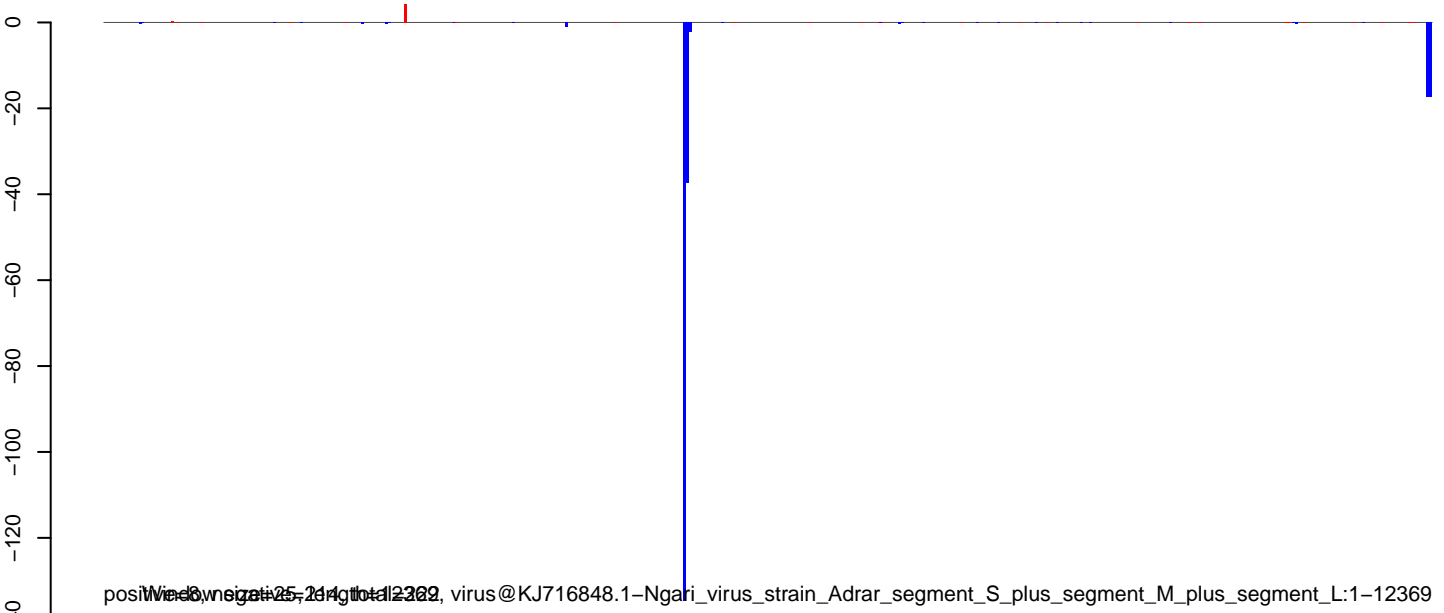
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



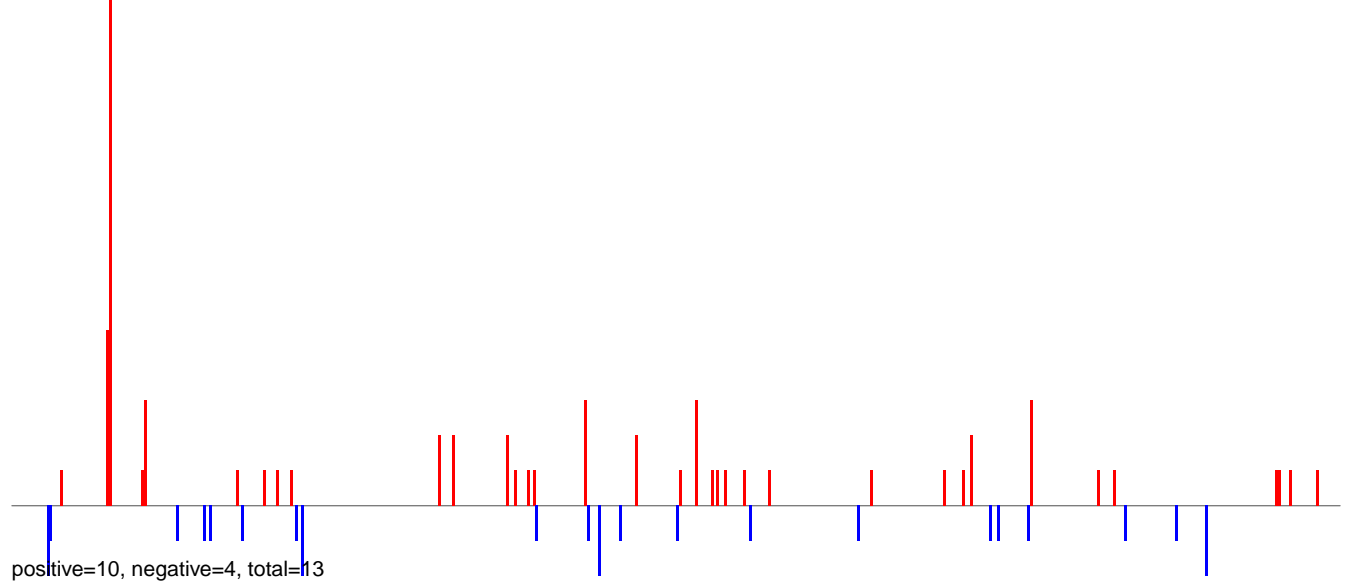
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



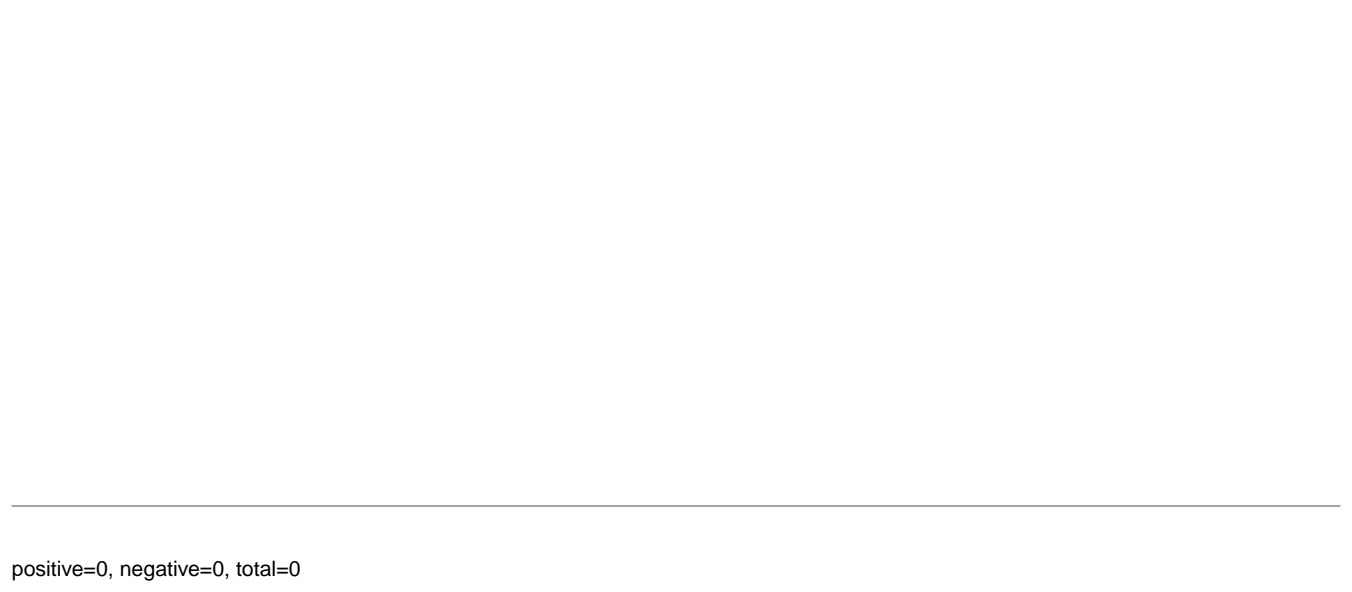
AeAeg_LVP_pMos3DBhspPBac.rep



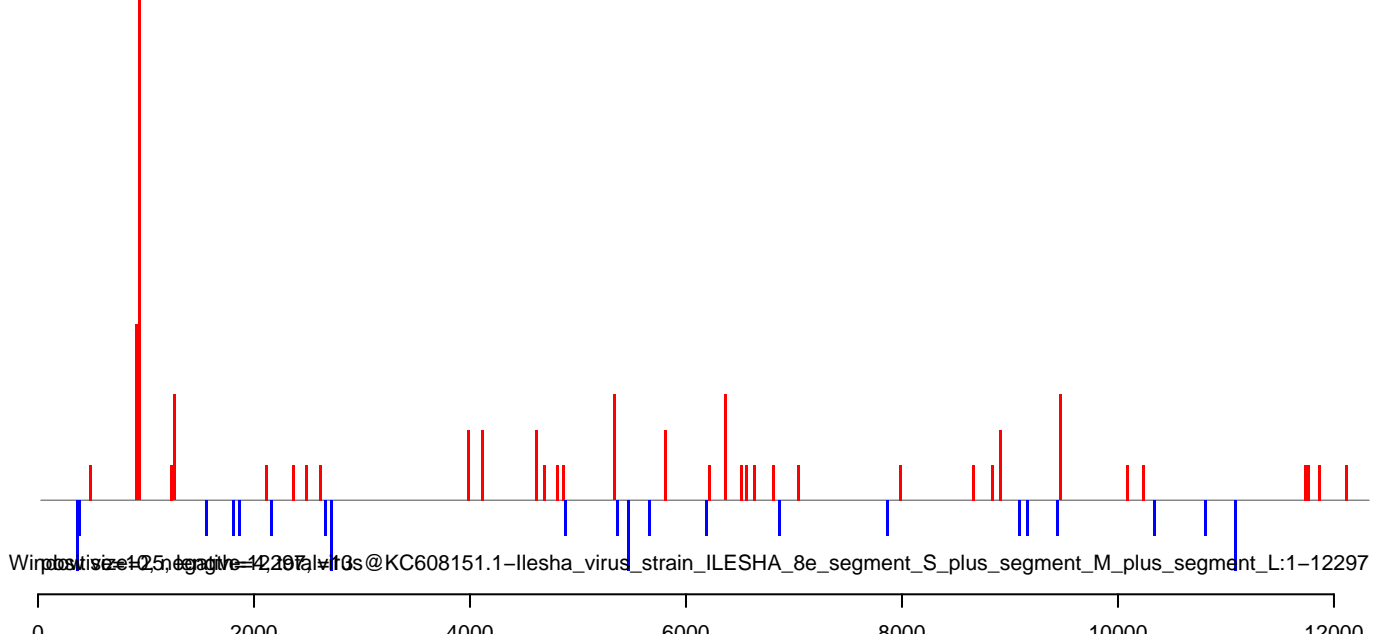
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



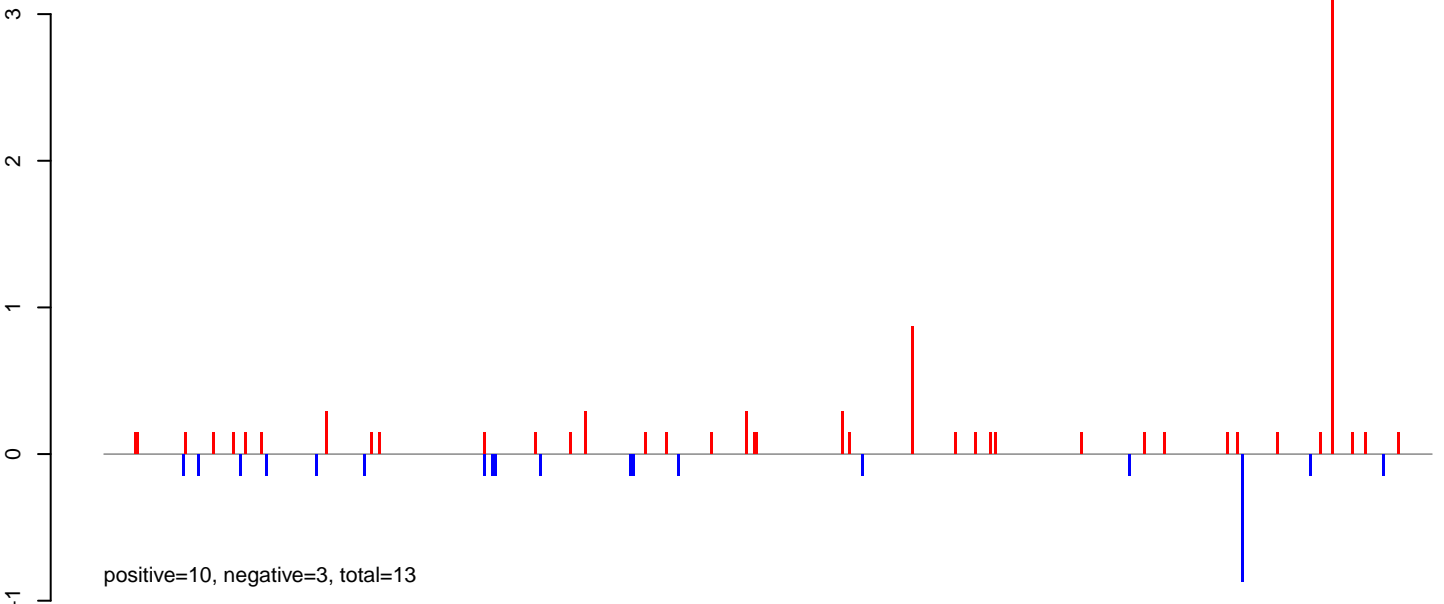
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



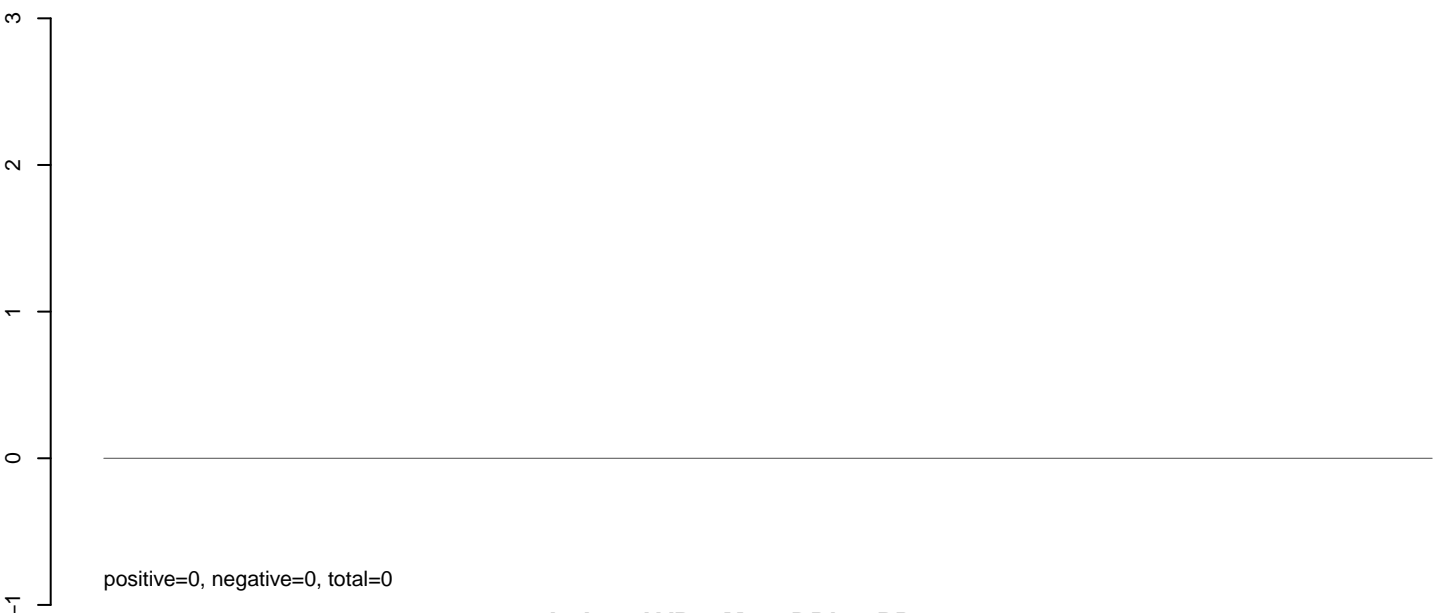
AeAeg_LVP_pMos3DBhspPBac.rep



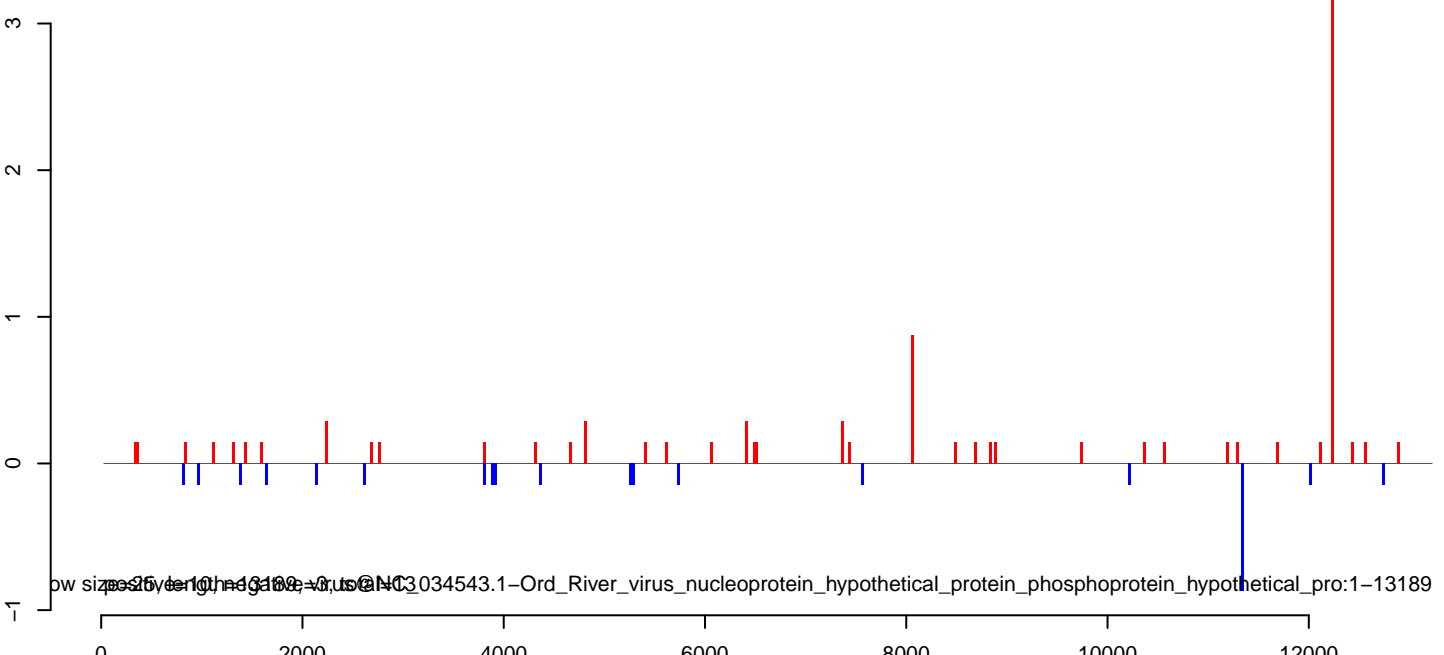
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



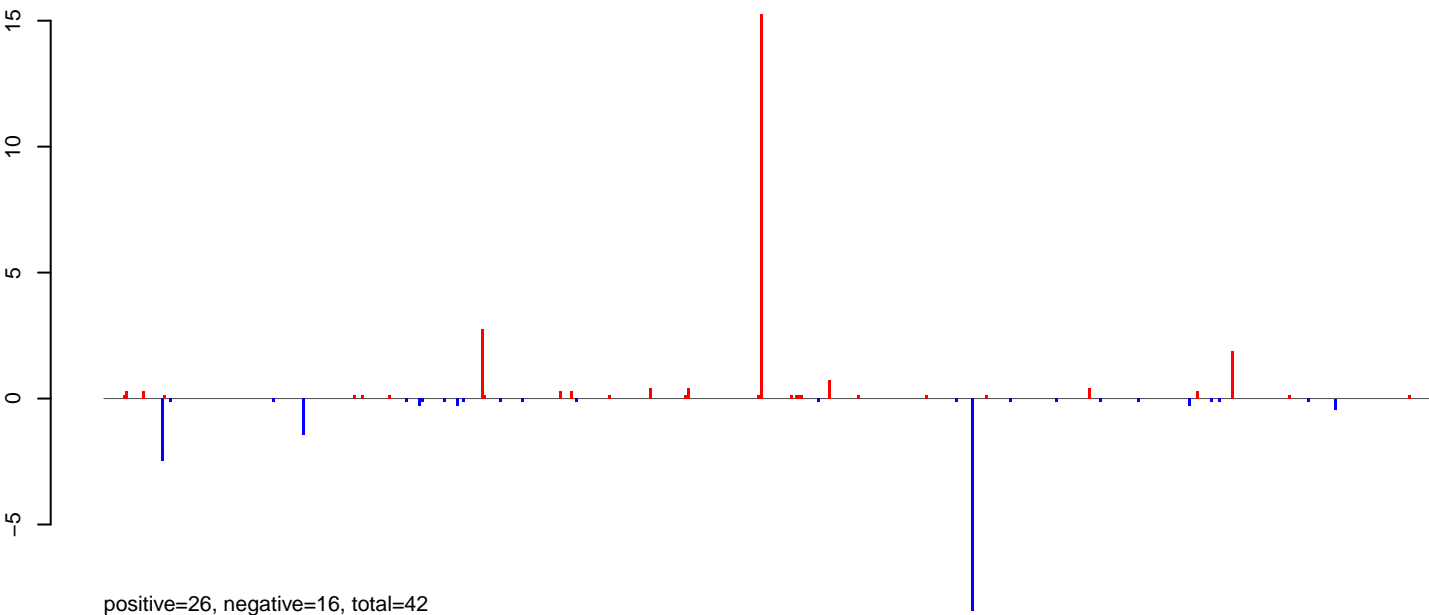
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



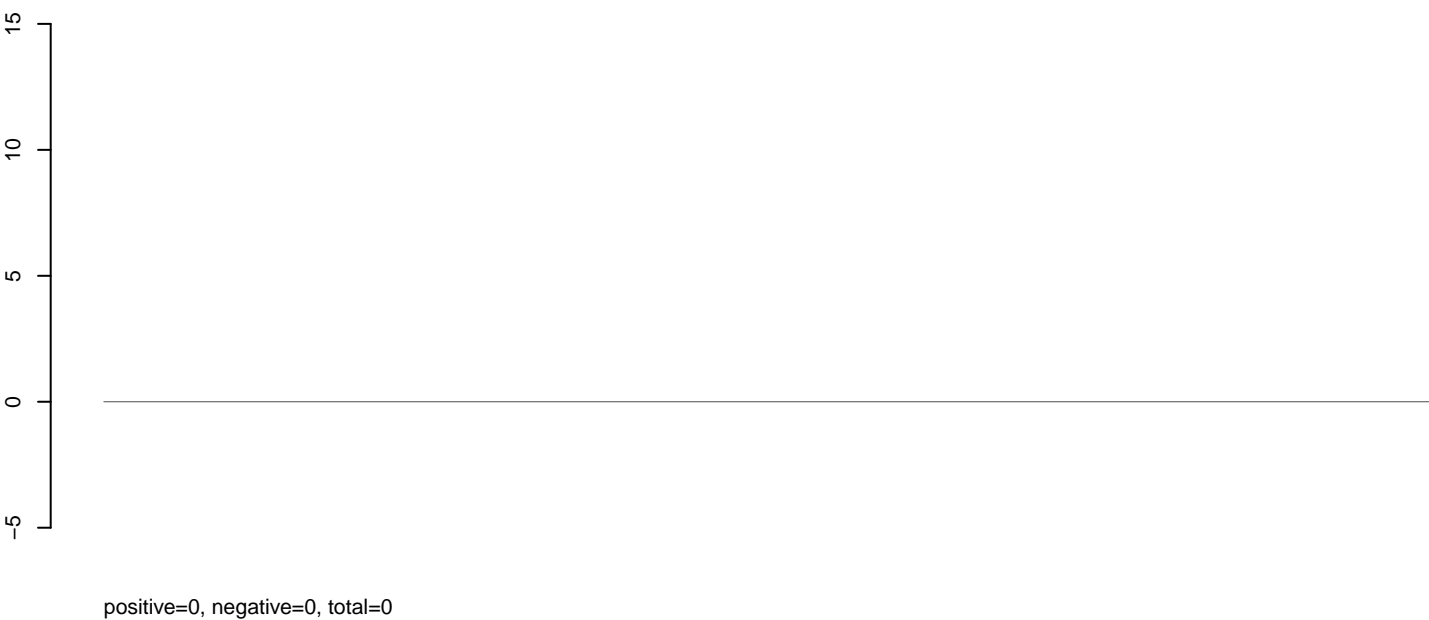
AeAeg_LVP_pMos3DBhspPBac.rep



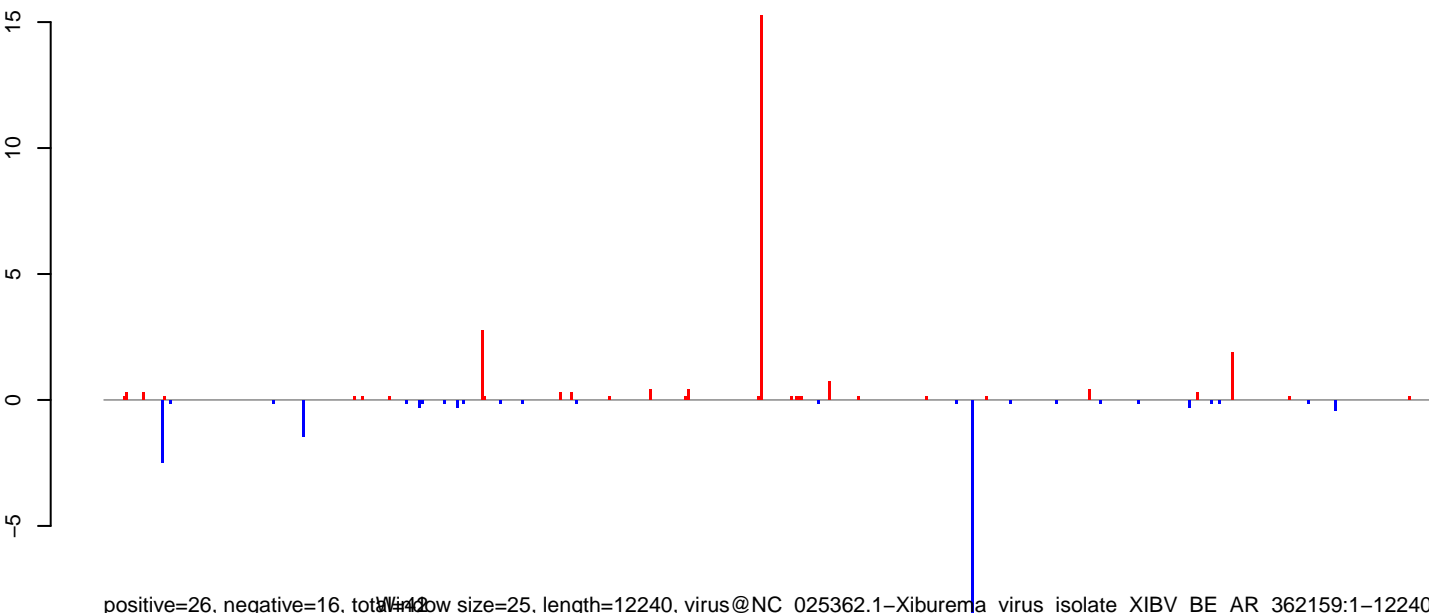
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



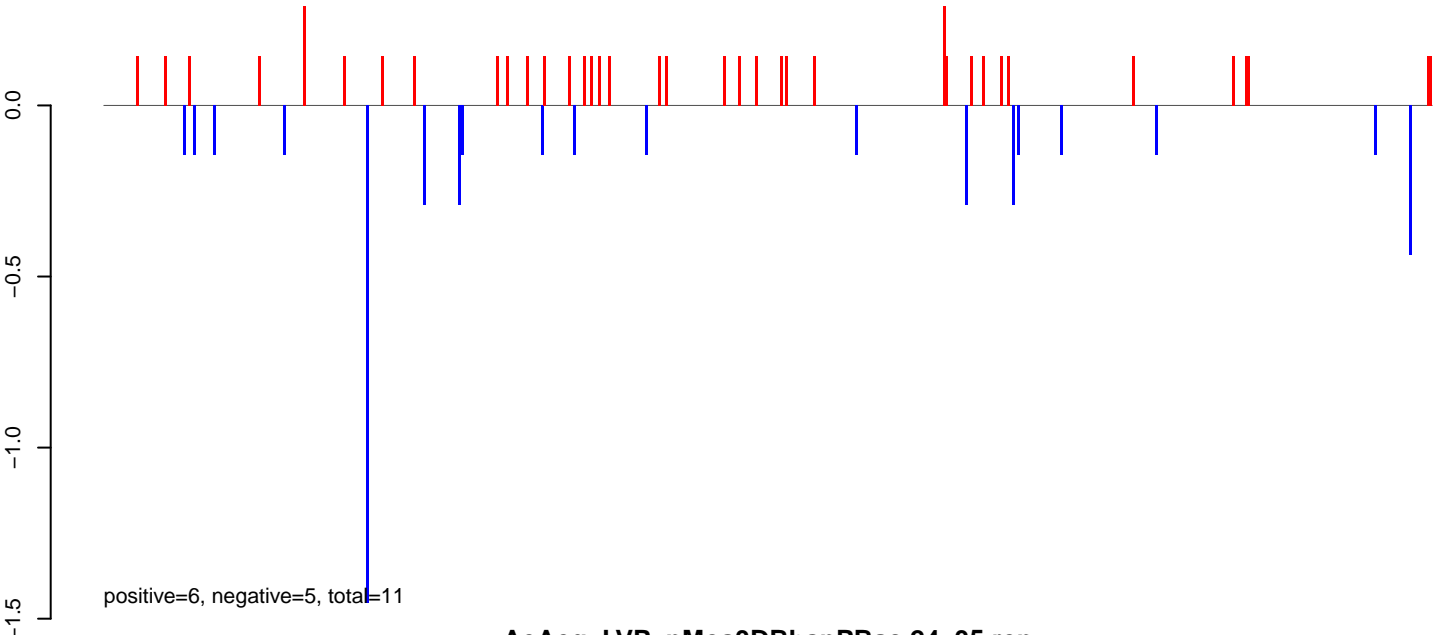
AeAeg_LVP_pMos3DBhspPBac.rep



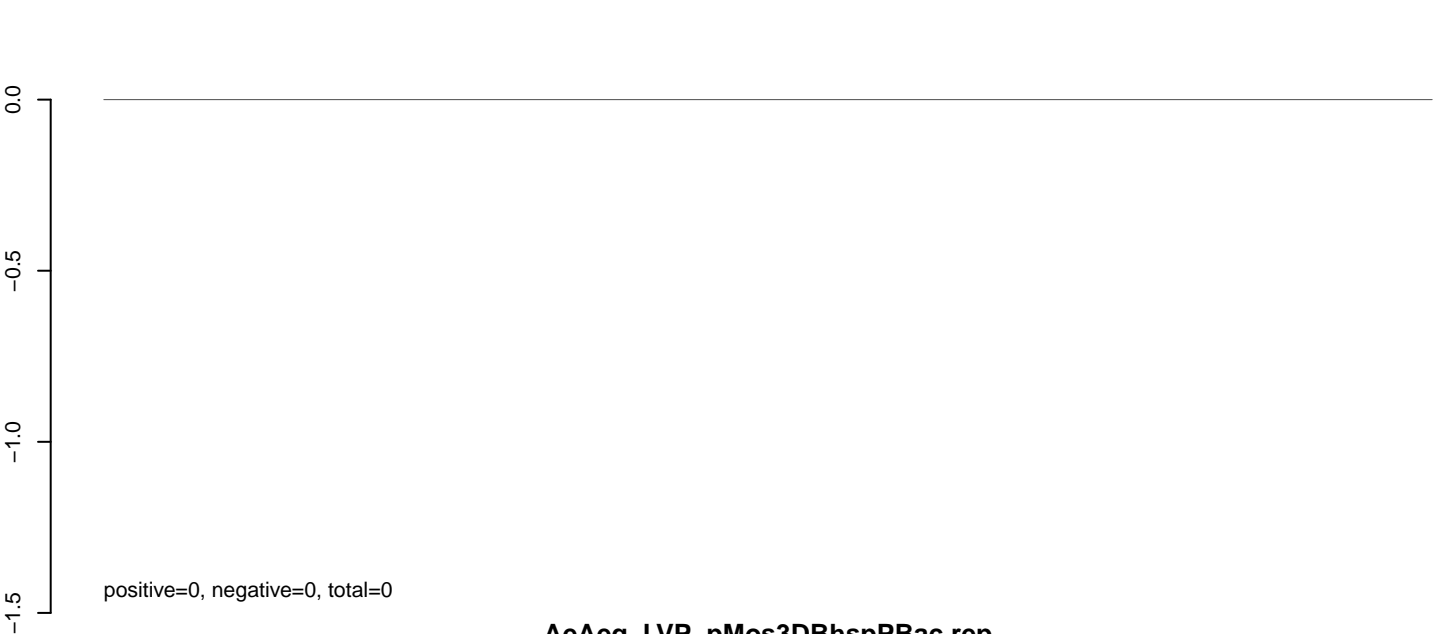
Window size=25, length=12240, virus@NC_025362.1-Xiburema_virus_isolate_XIBV_BE_AR_362159:1-12240

0 2000 4000 6000 8000 10000 12000

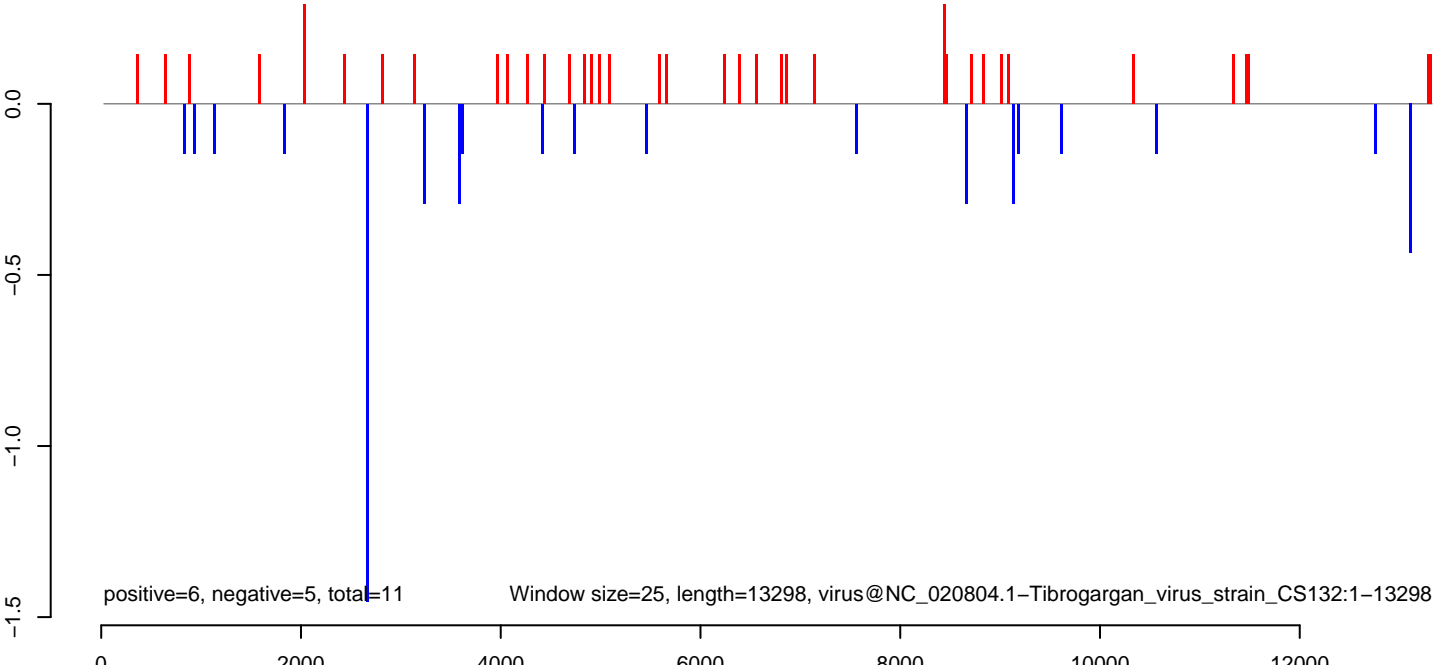
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



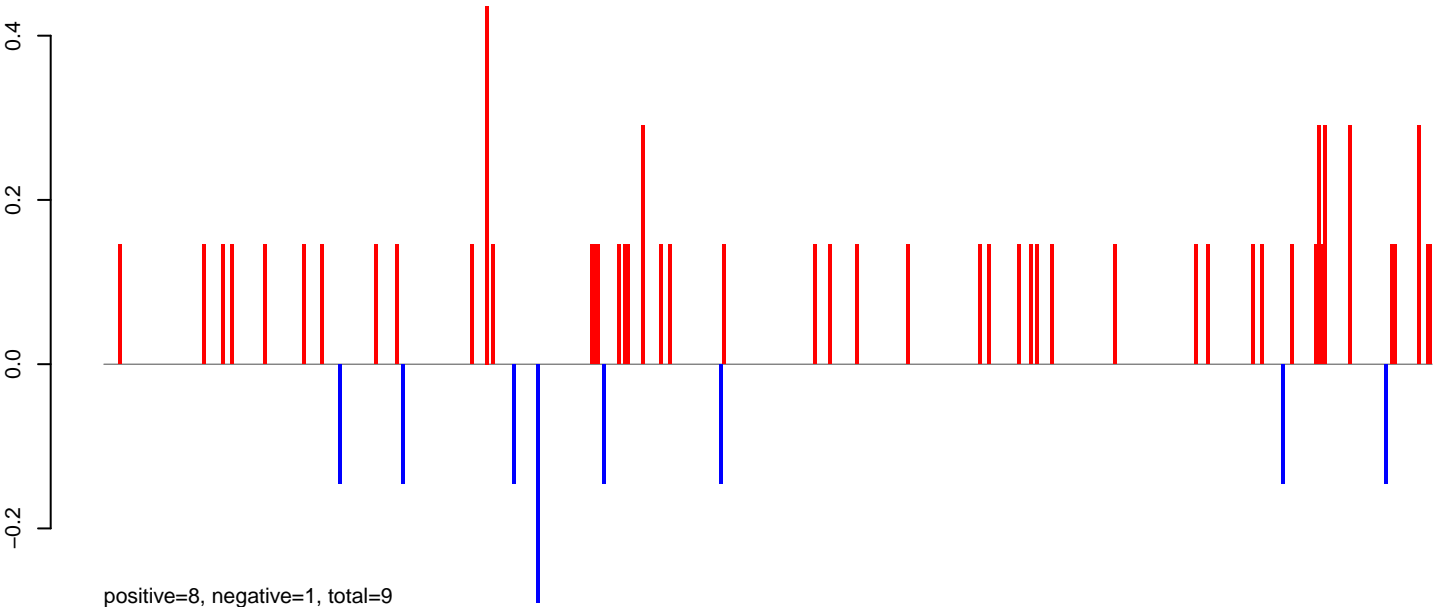
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



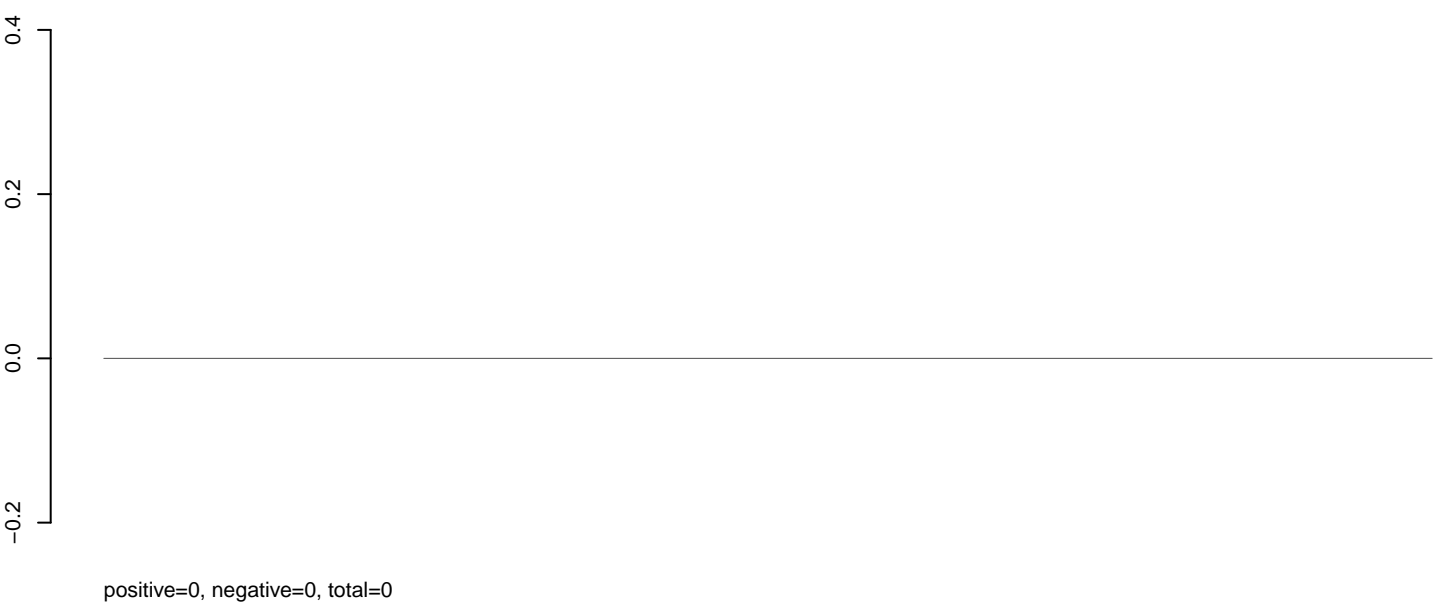
AeAeg_LVP_pMos3DBhspPBac.rep



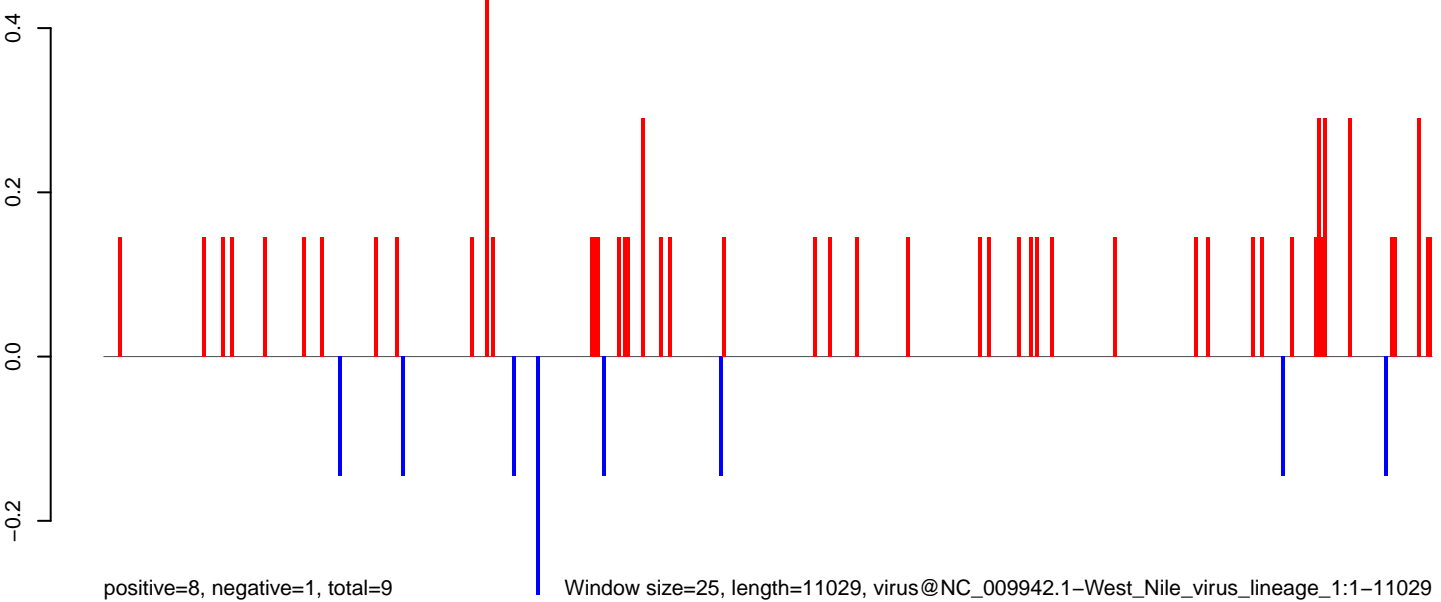
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

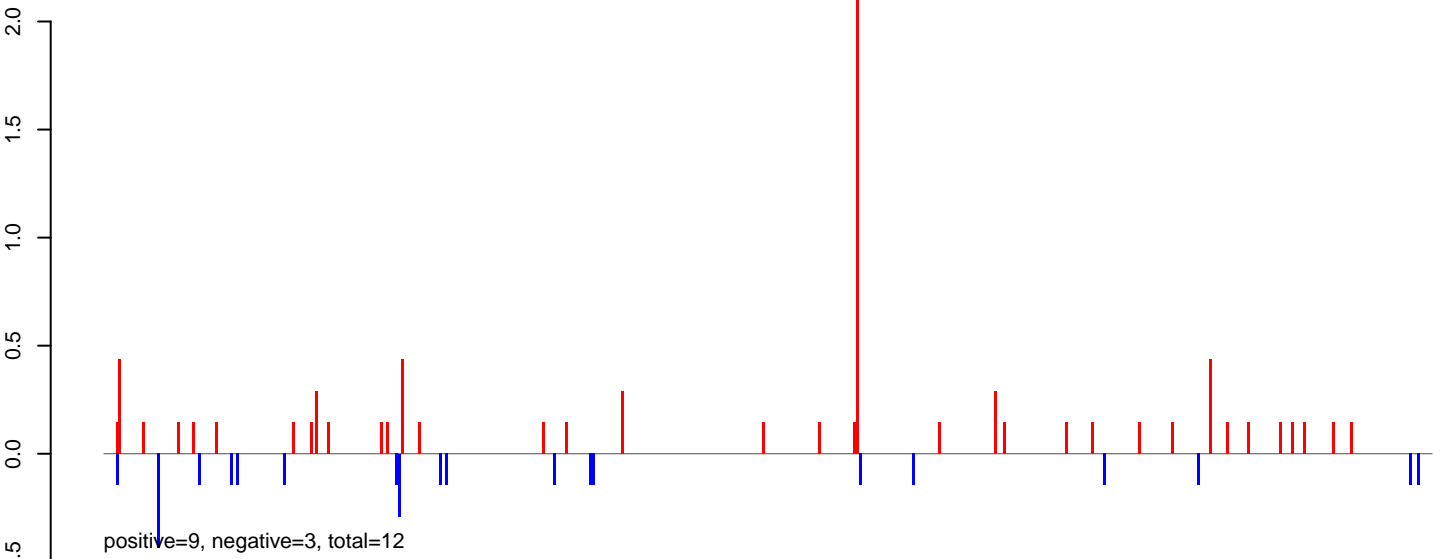


AeAeg_LVP_pMos3DBhspPBac.rep

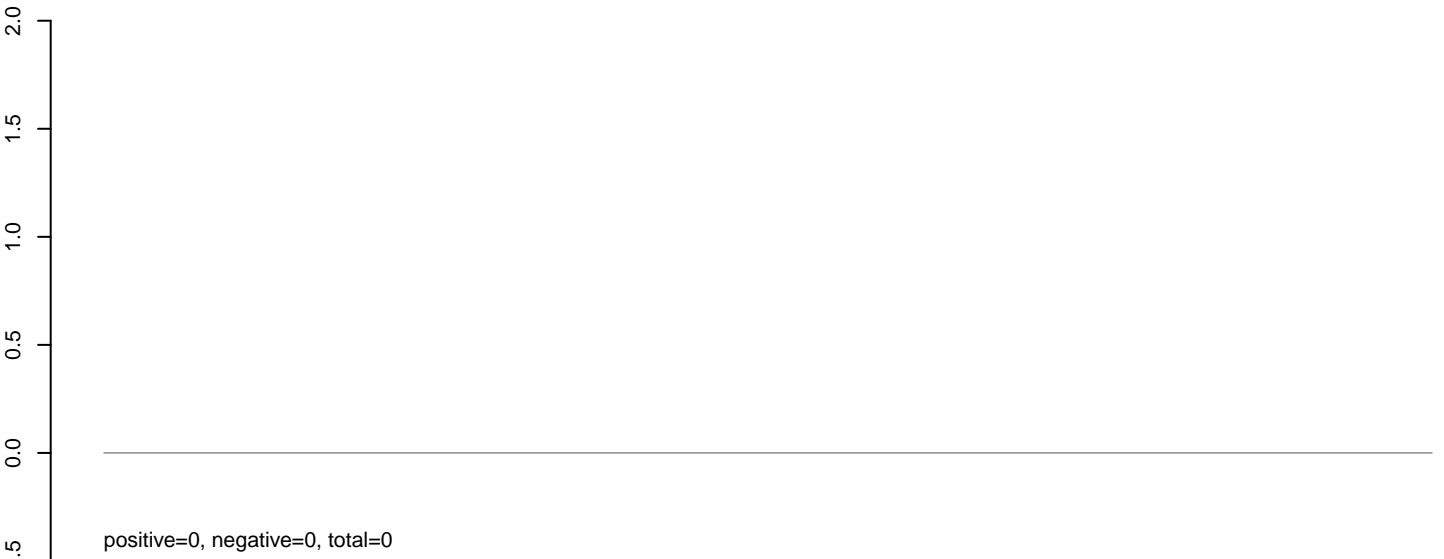


0 2000 4000 6000 8000 10000

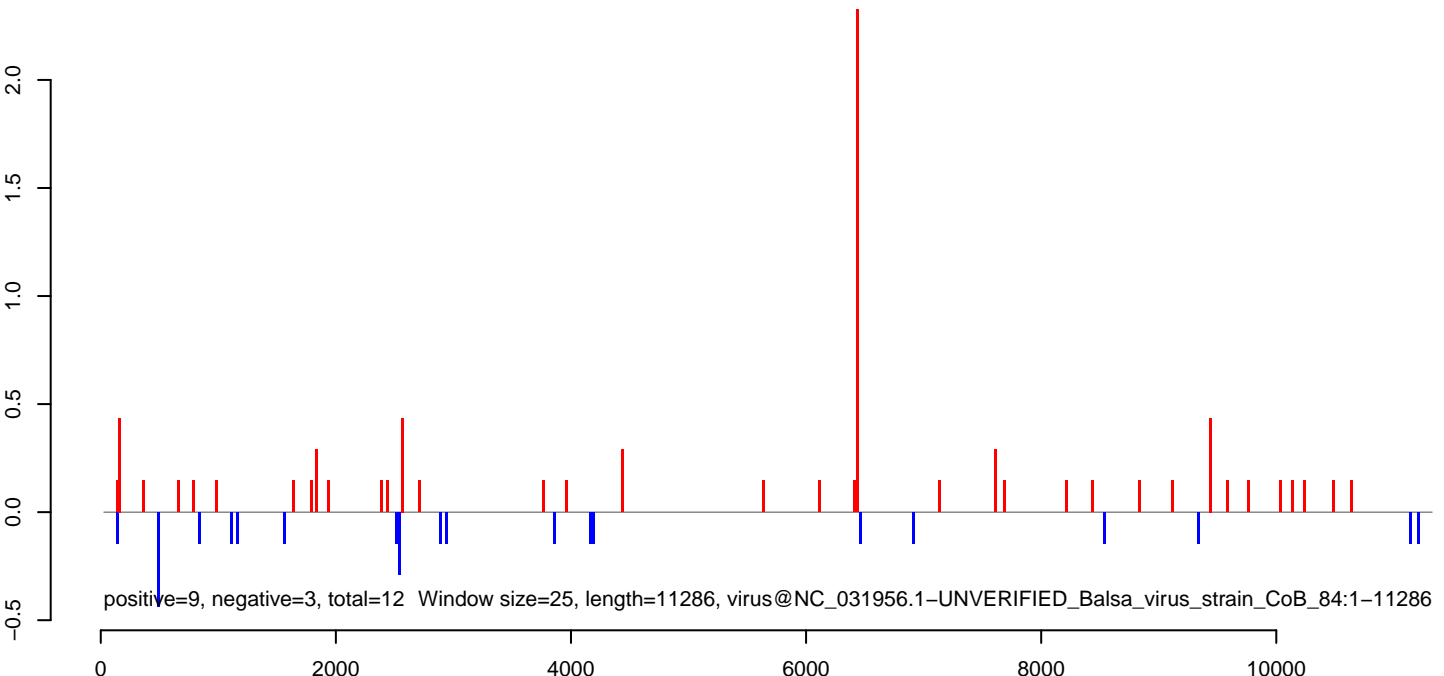
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



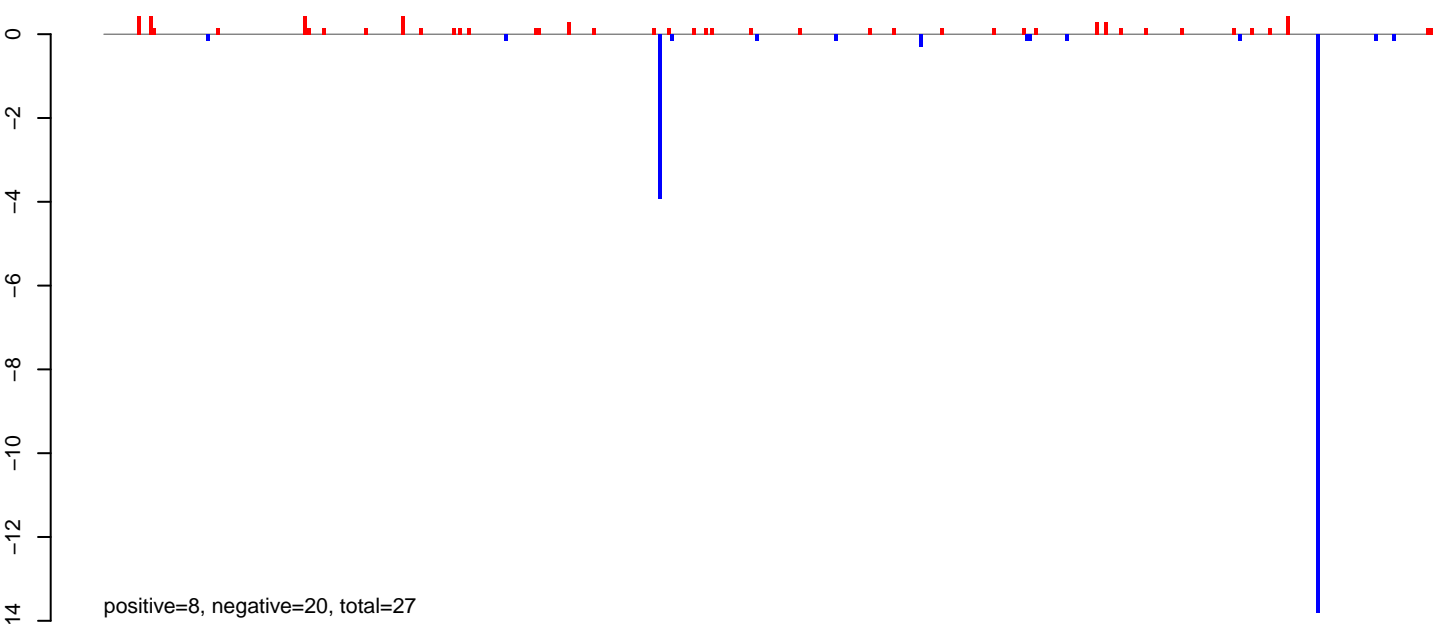
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



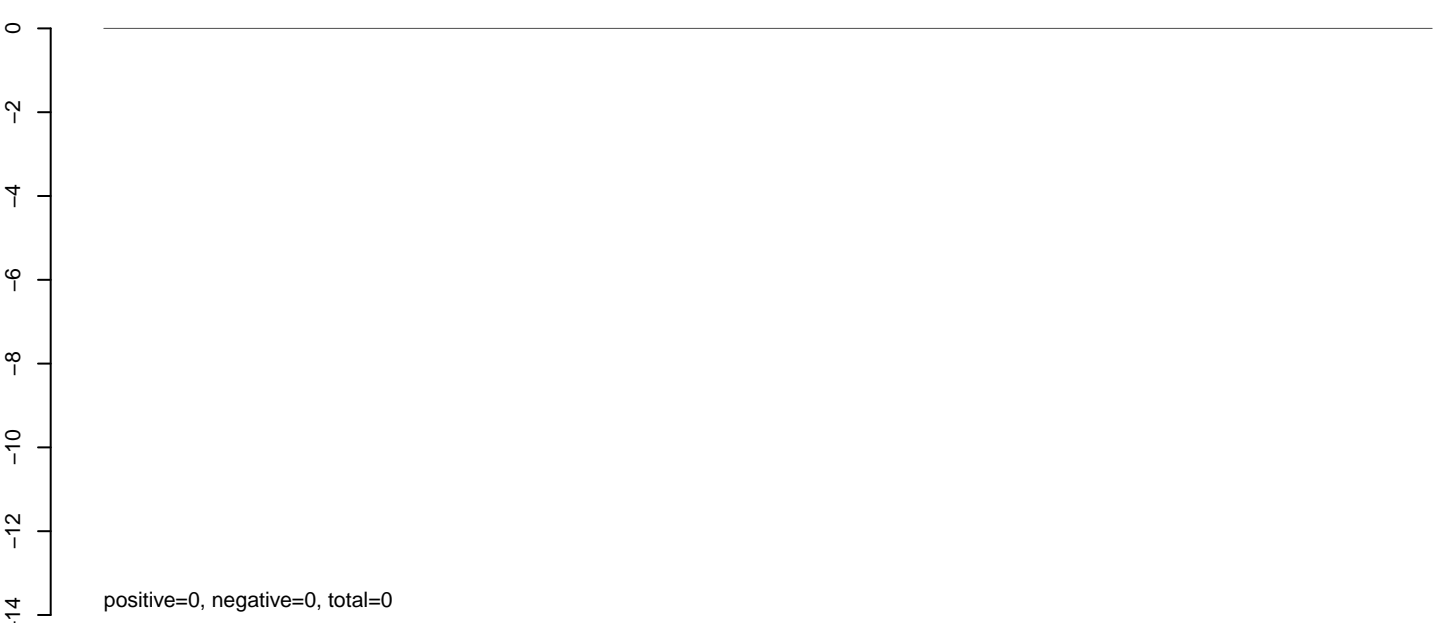
AeAeg_LVP_pMos3DBhspPBac.rep



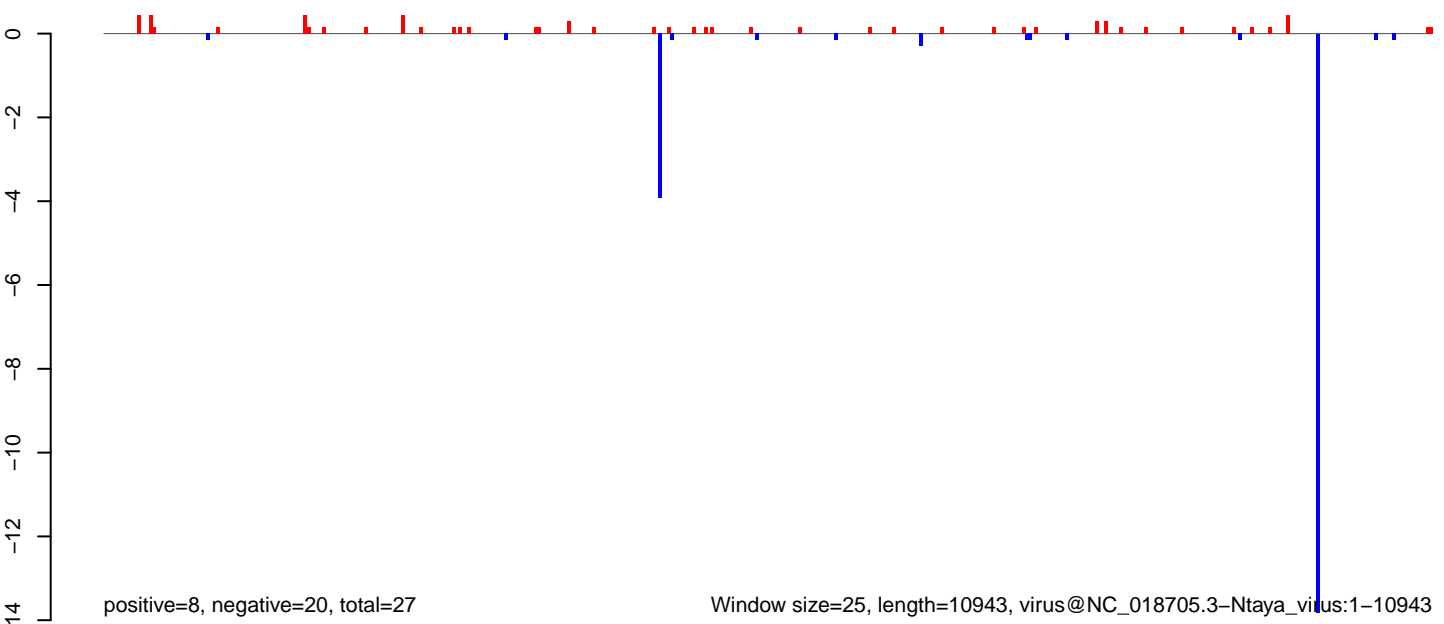
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



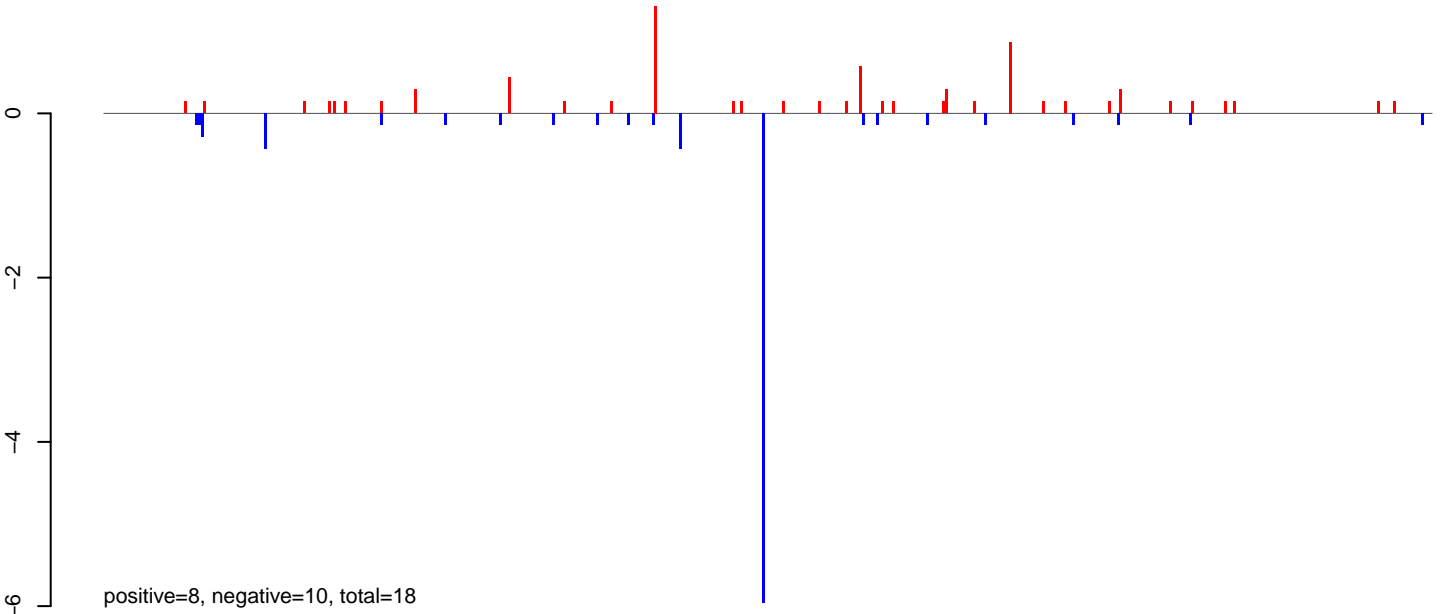
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



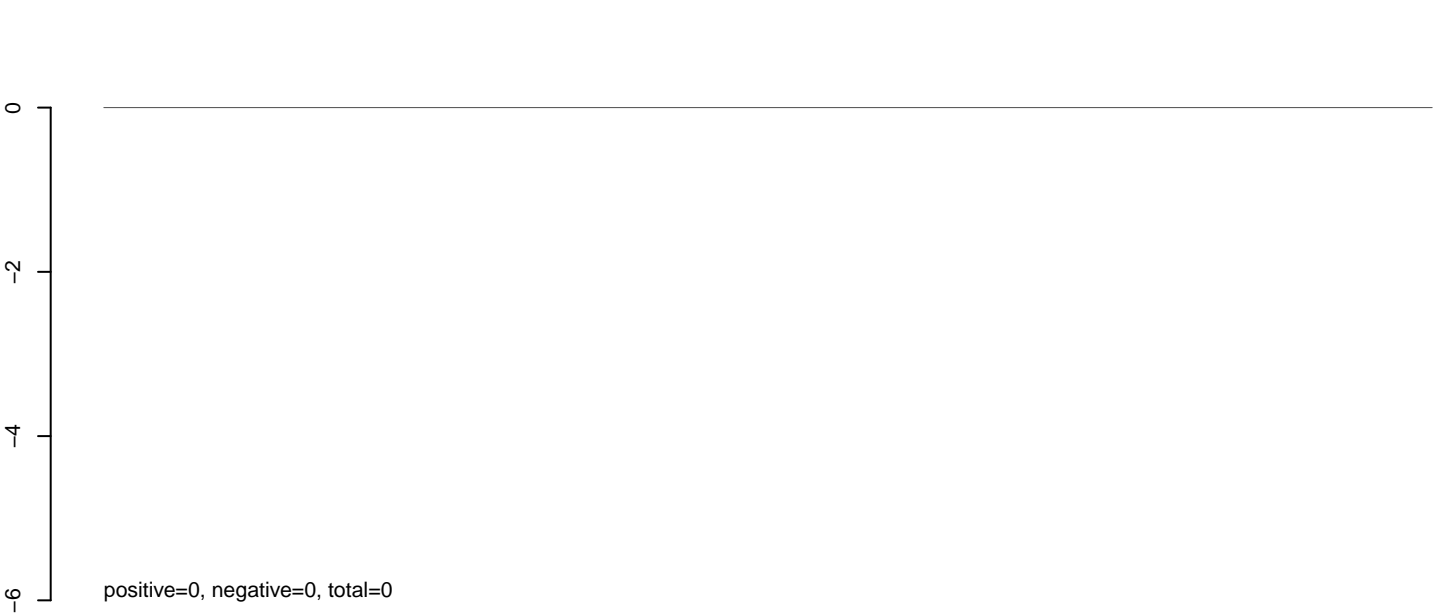
AeAeg_LVP_pMos3DBhspPBac.rep



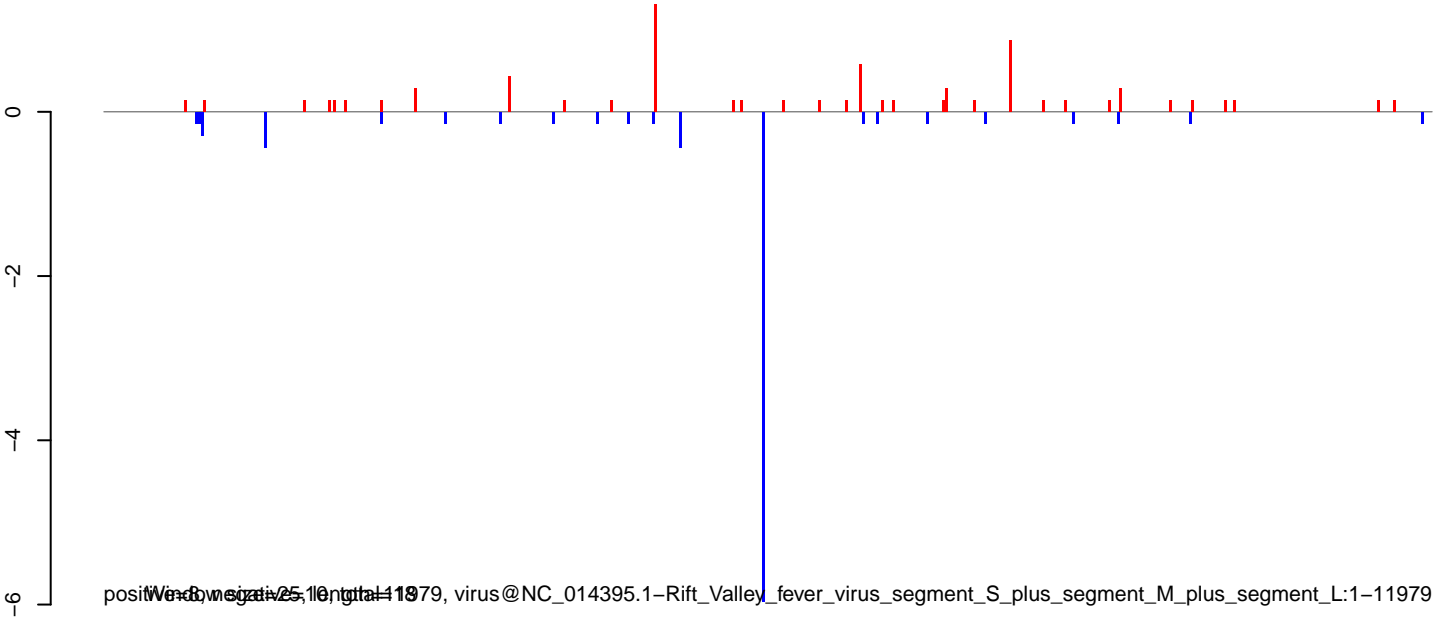
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



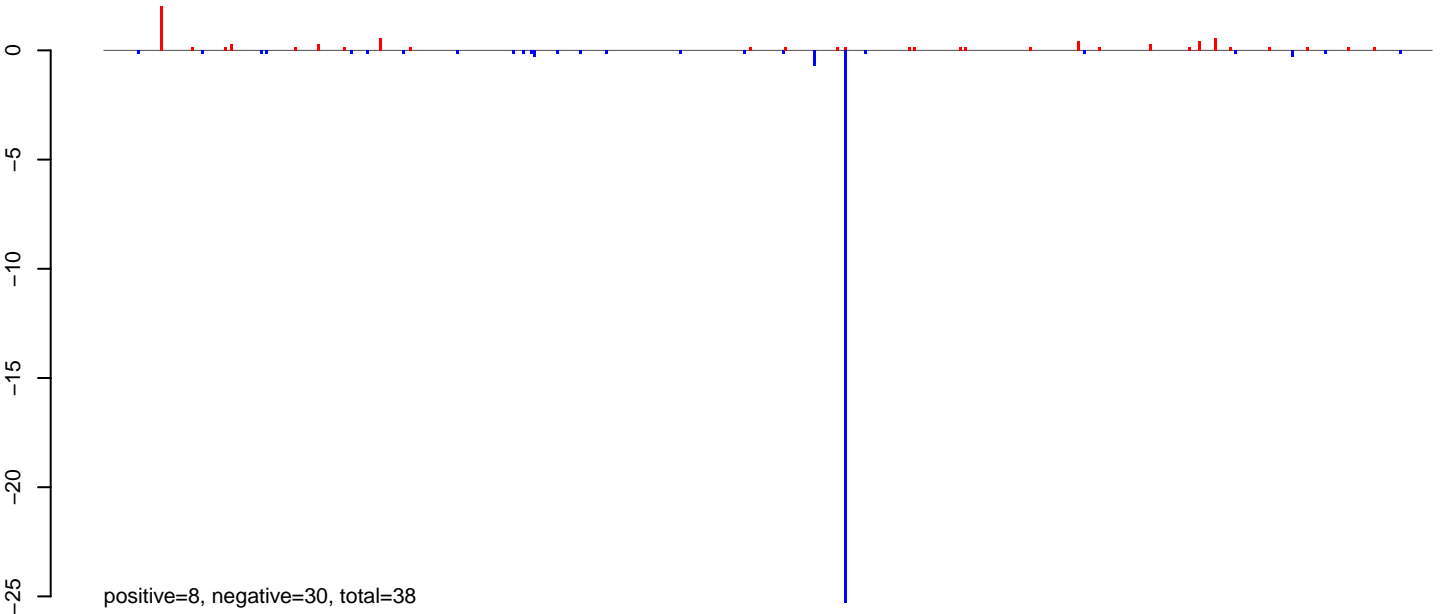
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



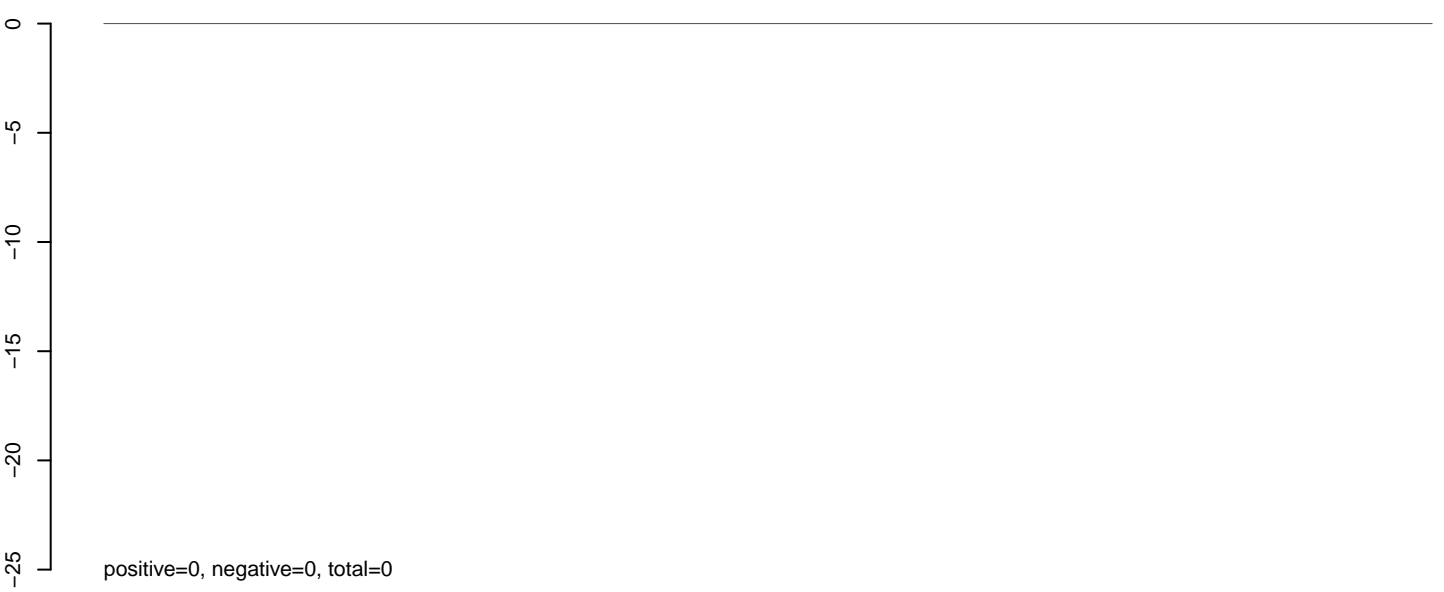
AeAeg_LVP_pMos3DBhspPBac.rep



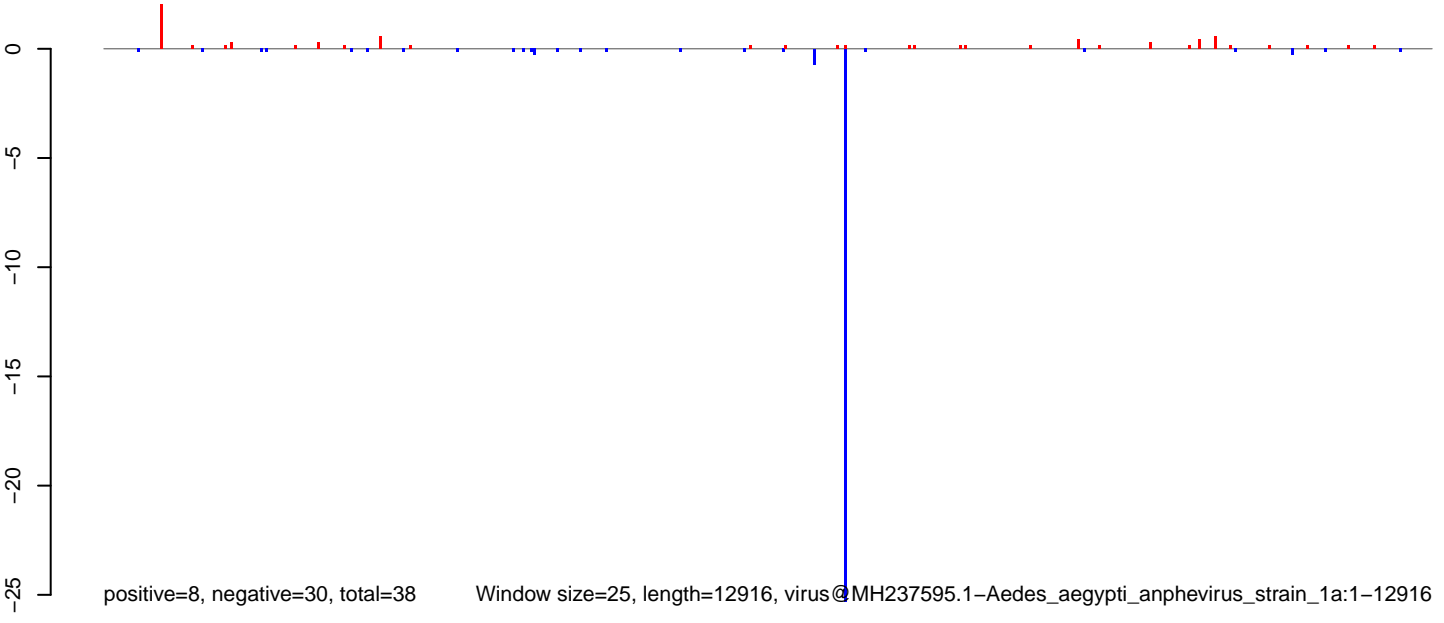
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



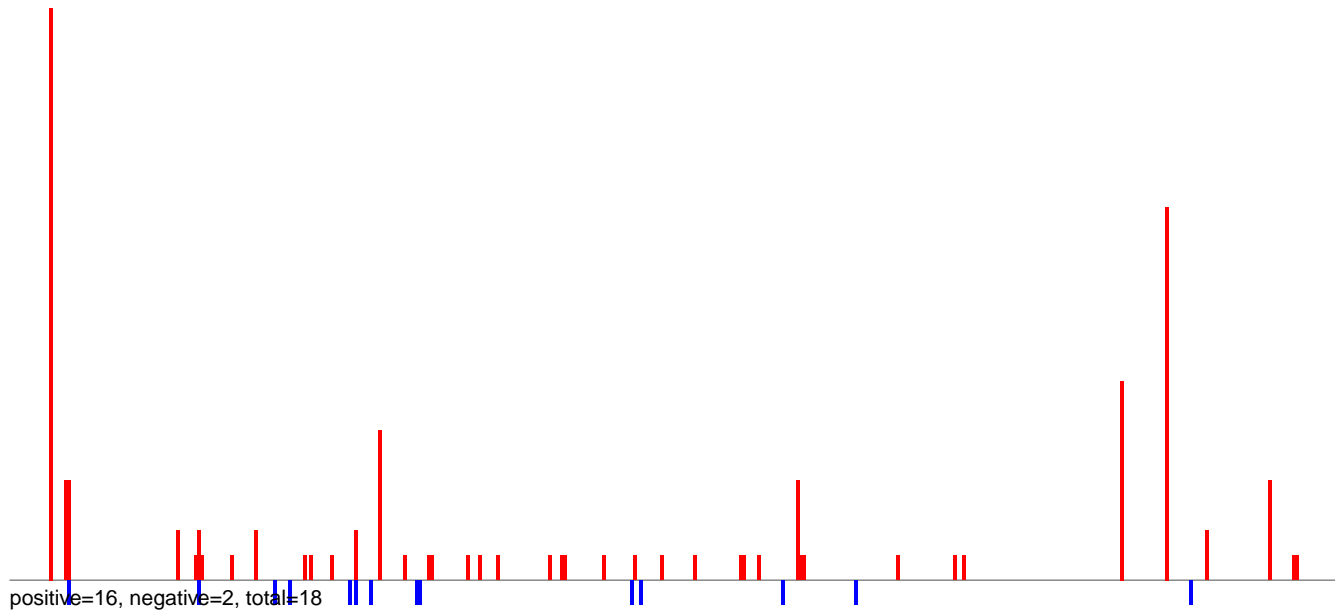
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



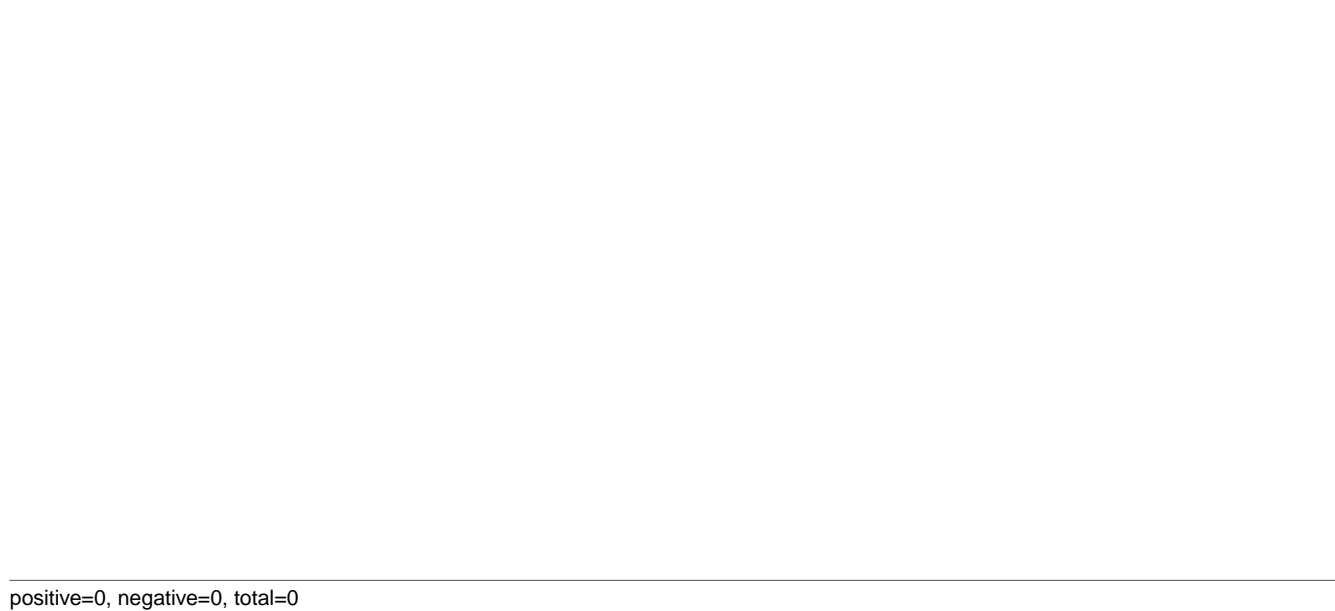
AeAeg_LVP_pMos3DBhspPBac.rep



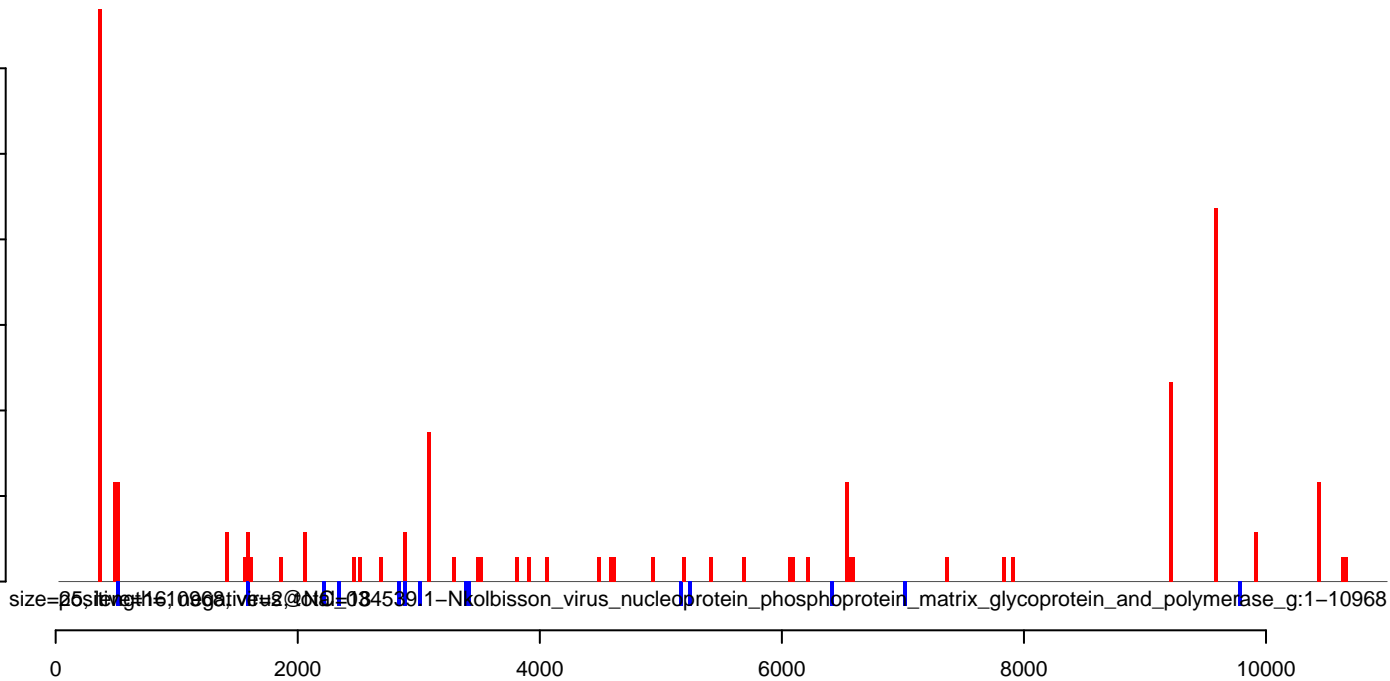
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



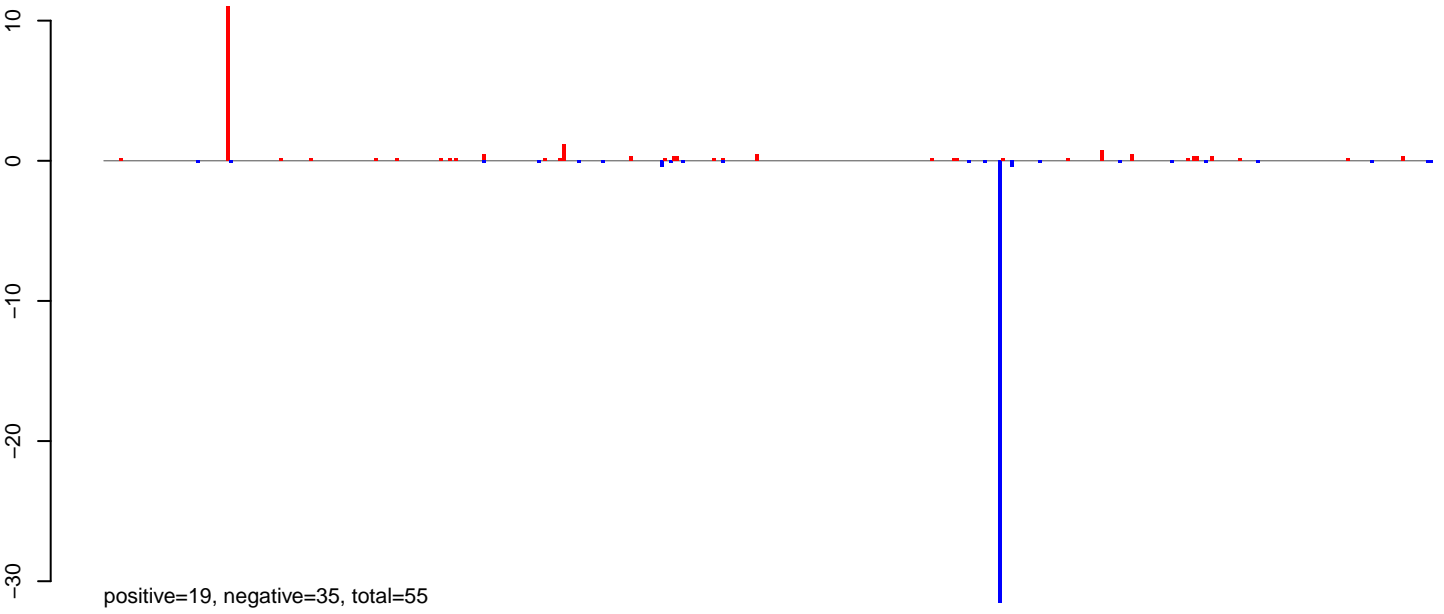
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



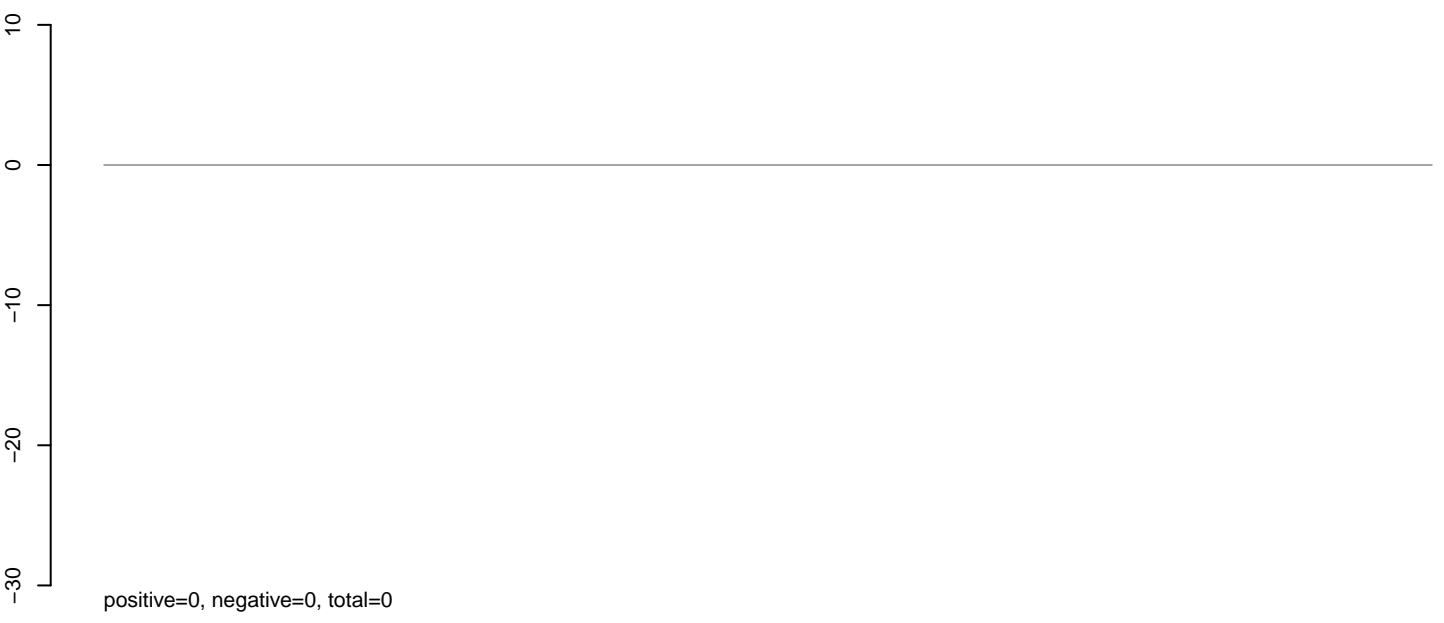
AeAeg_LVP_pMos3DBhspPBac.rep



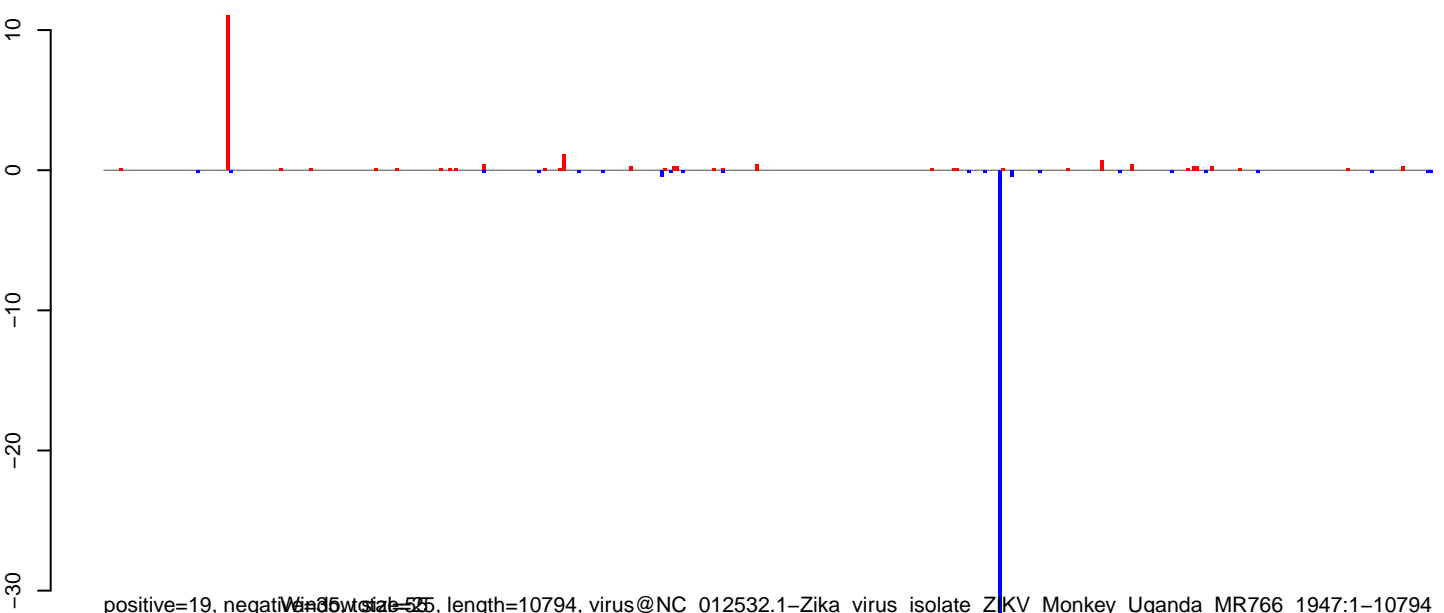
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



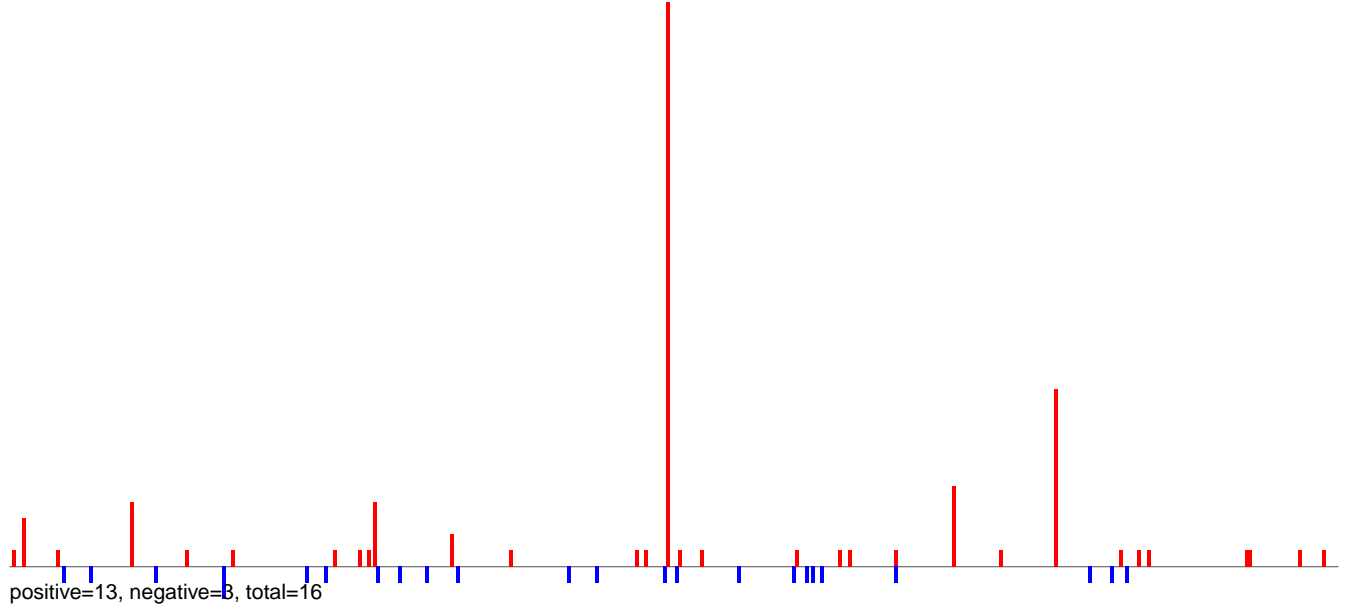
AeAeg_LVP_pMos3DBhspPBac.rep



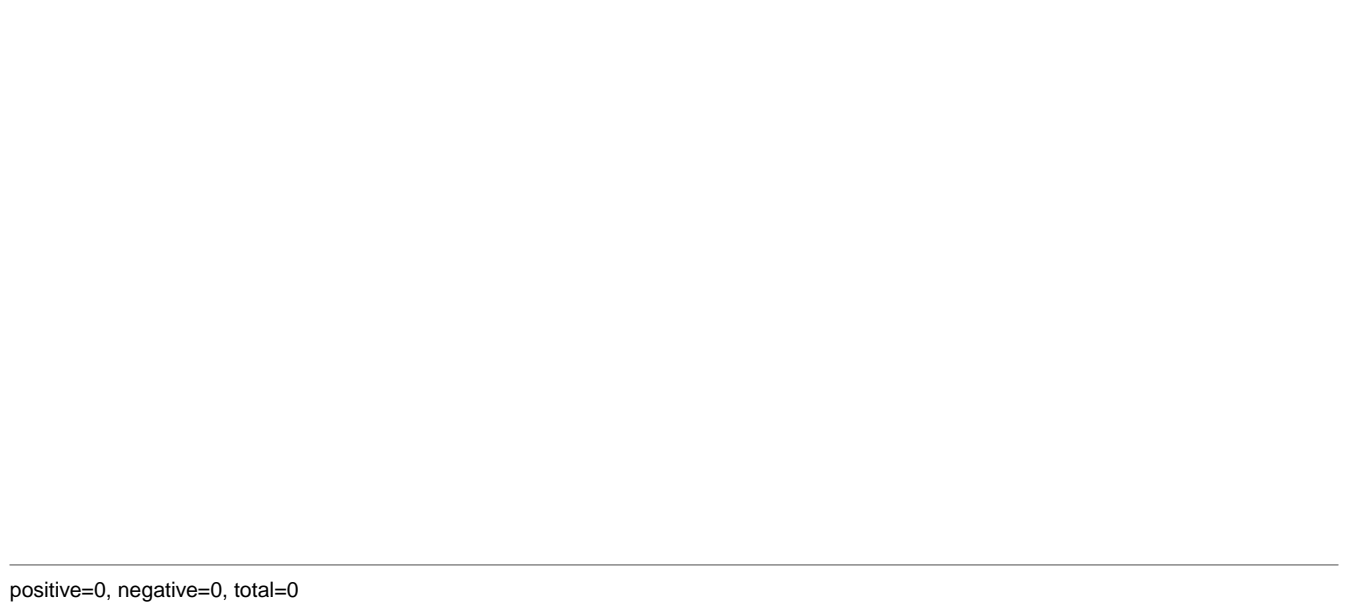
positive=19, negative=35, total=55, length=10794, virus@NC_012532.1-Zika_virus_isolate_ZIKV_Monkey_Uganda_MR766_1947:1-10794

0 2000 4000 6000 8000 10000

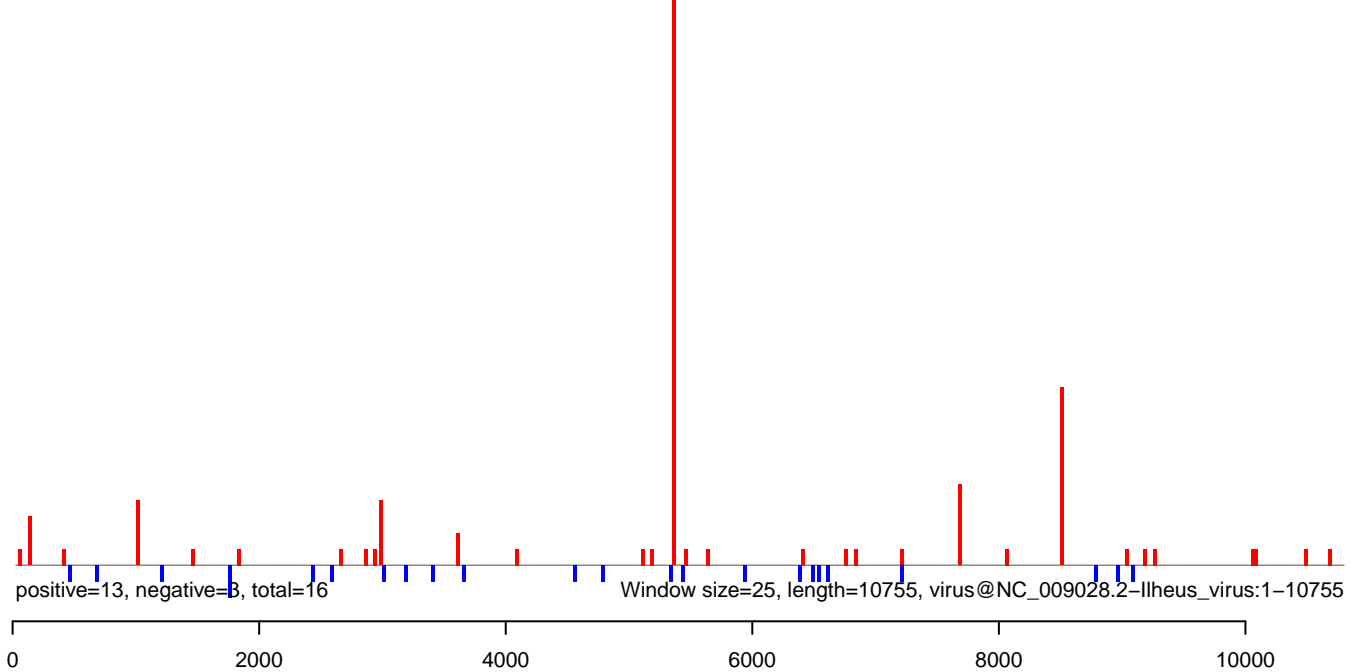
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



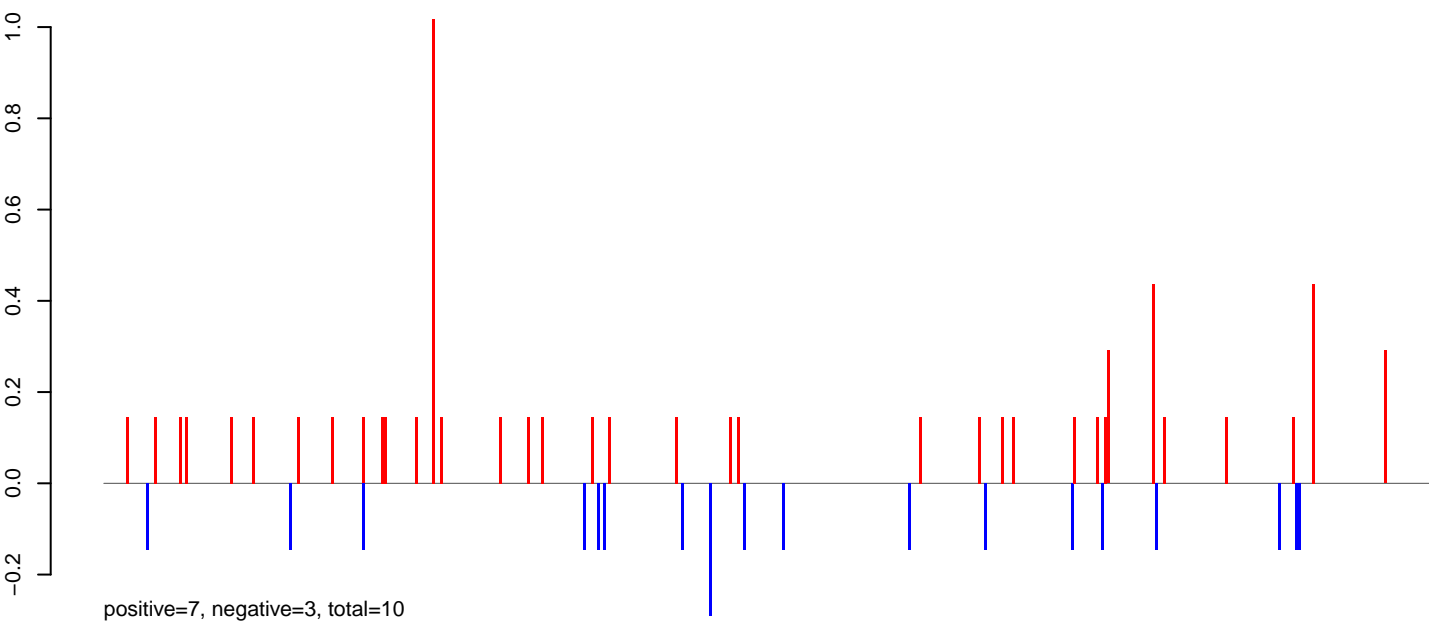
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



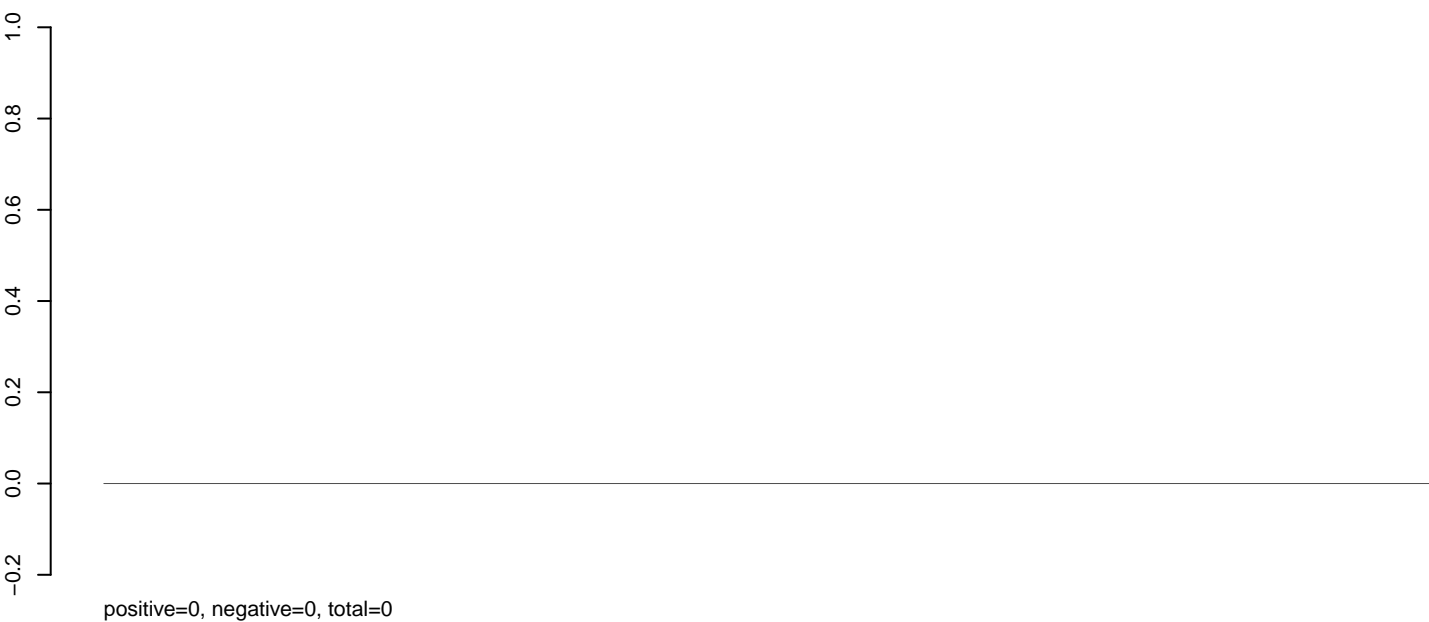
AeAeg_LVP_pMos3DBhspPBac.rep



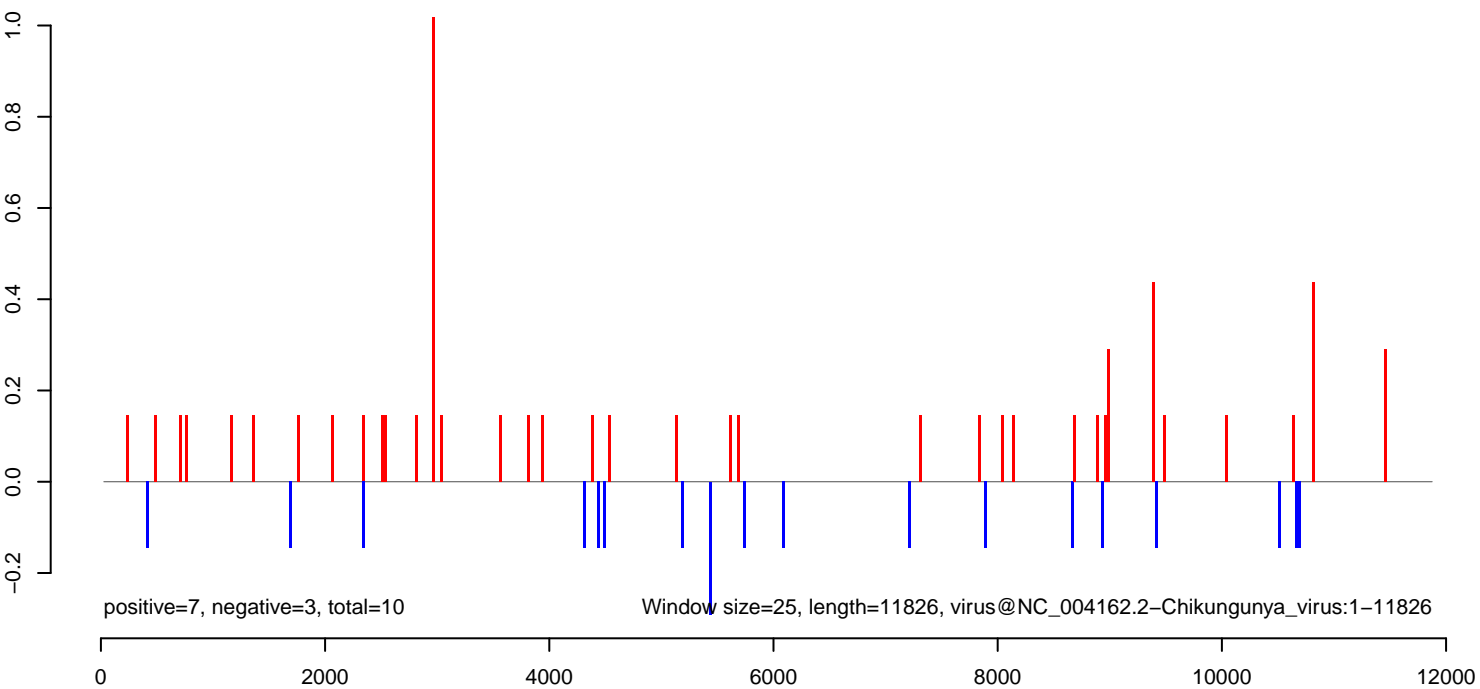
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



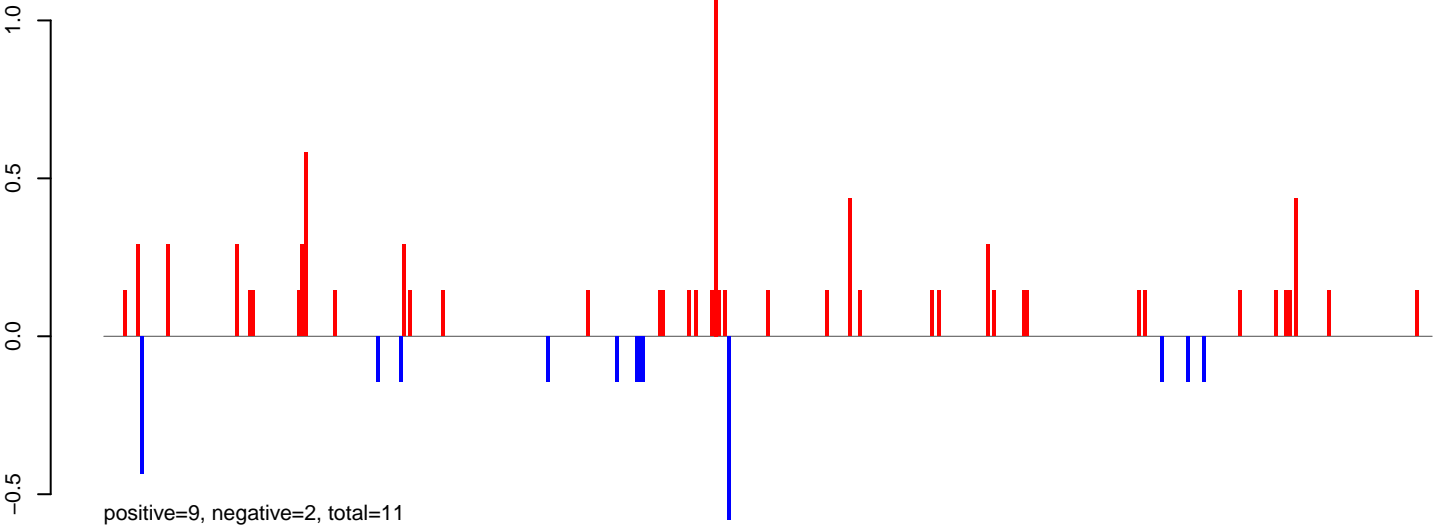
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



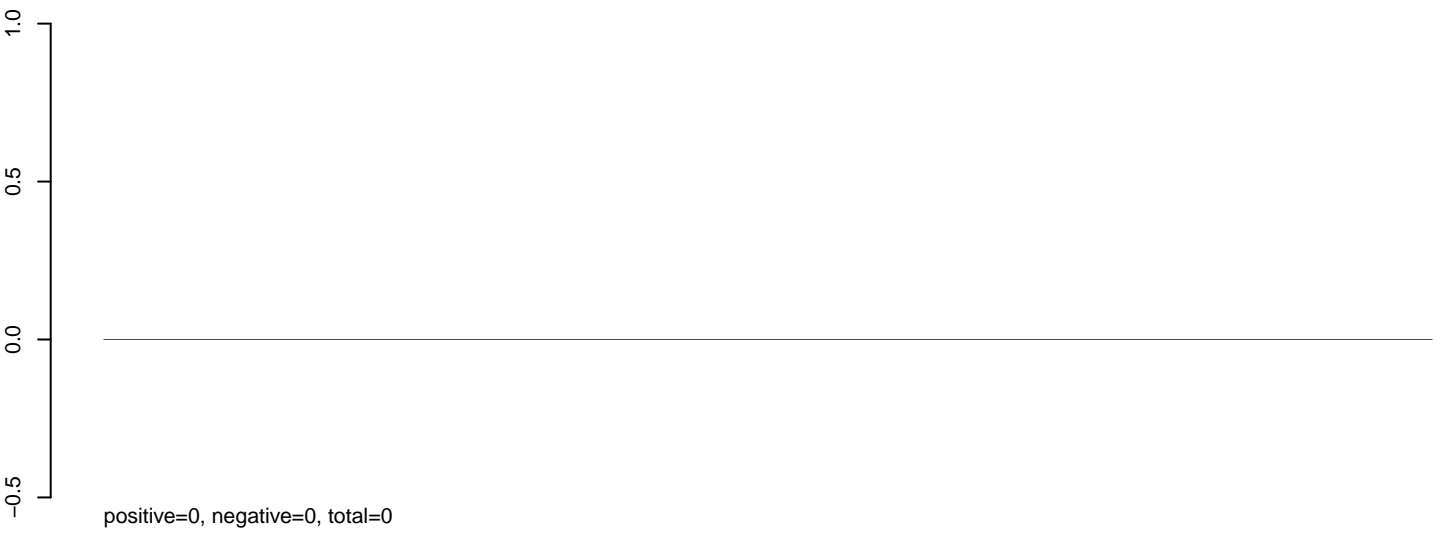
AeAeg_LVP_pMos3DBhspPBac.rep



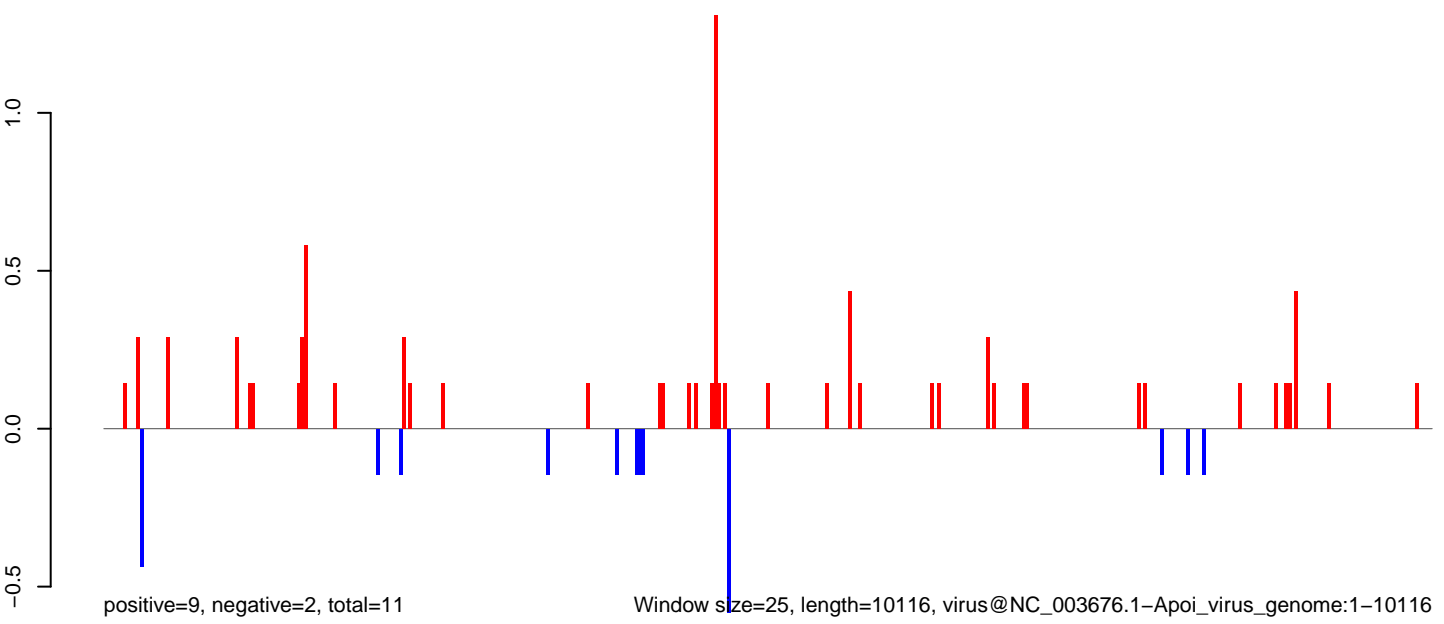
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



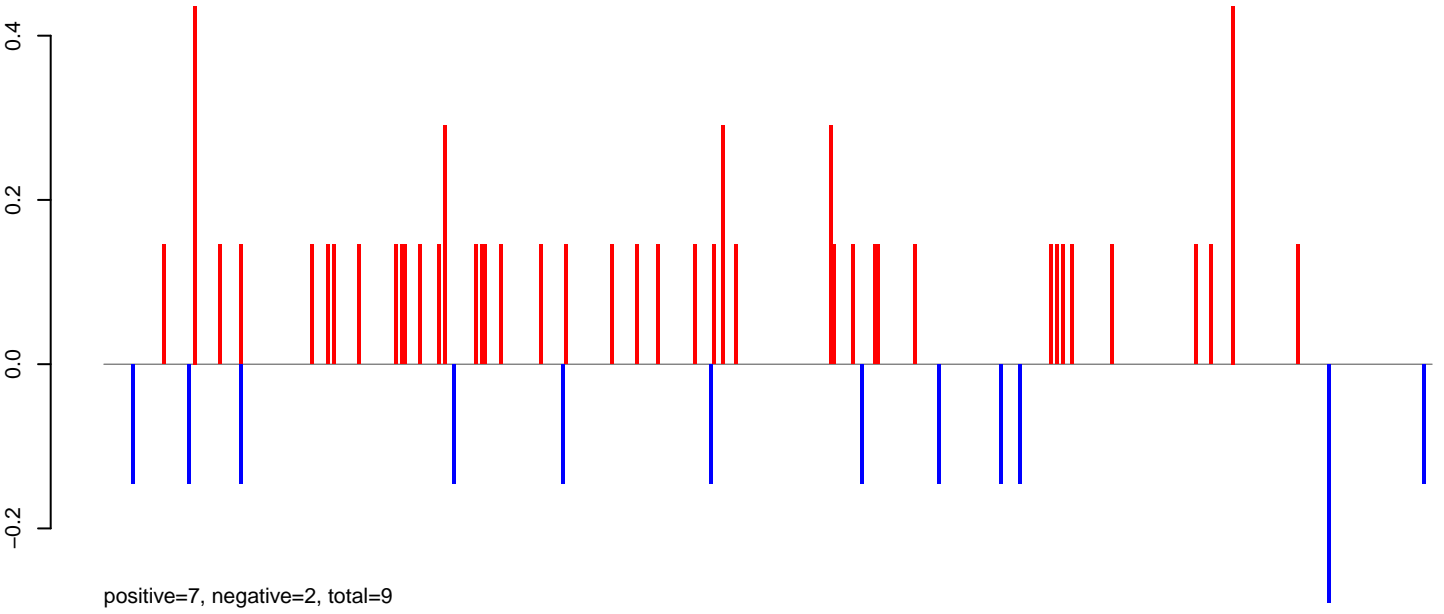
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



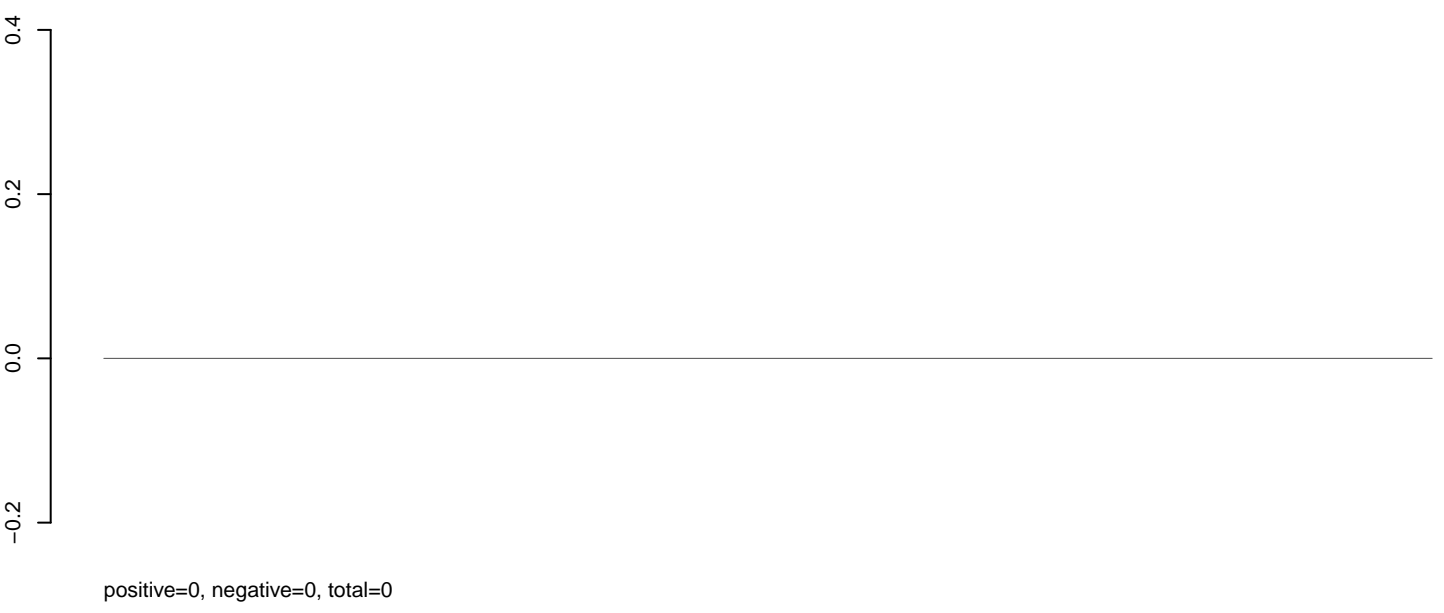
AeAeg_LVP_pMos3DBhspPBac.rep



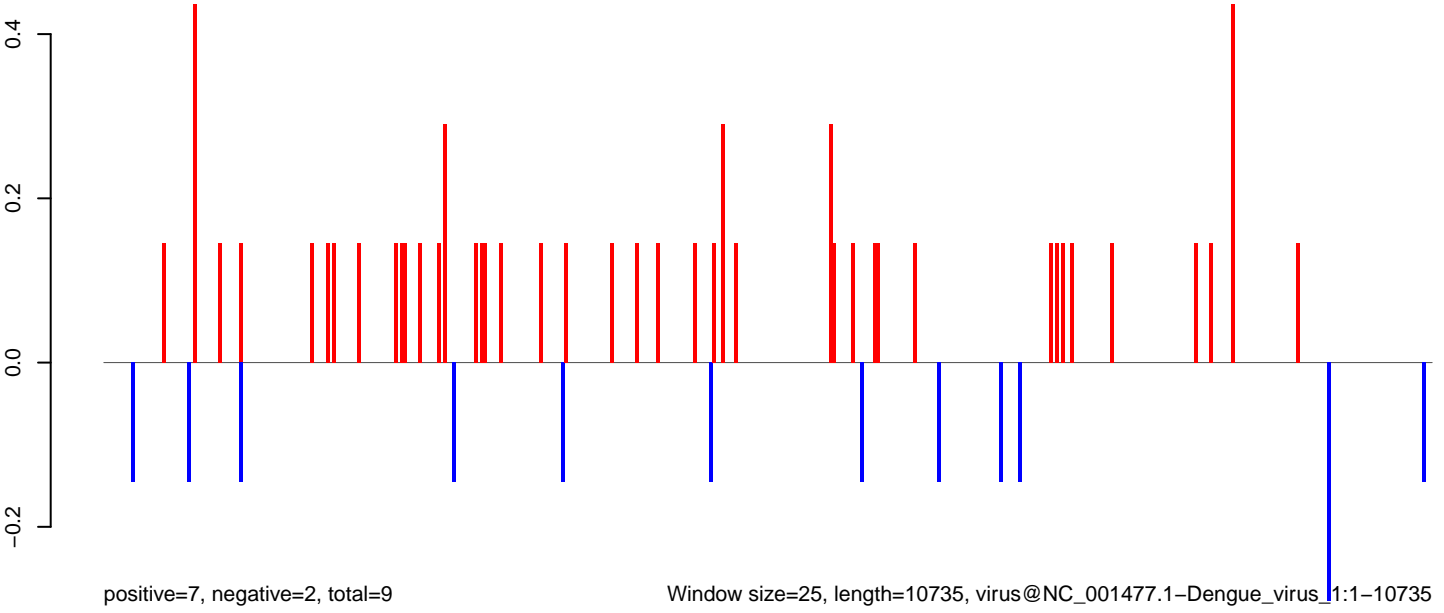
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



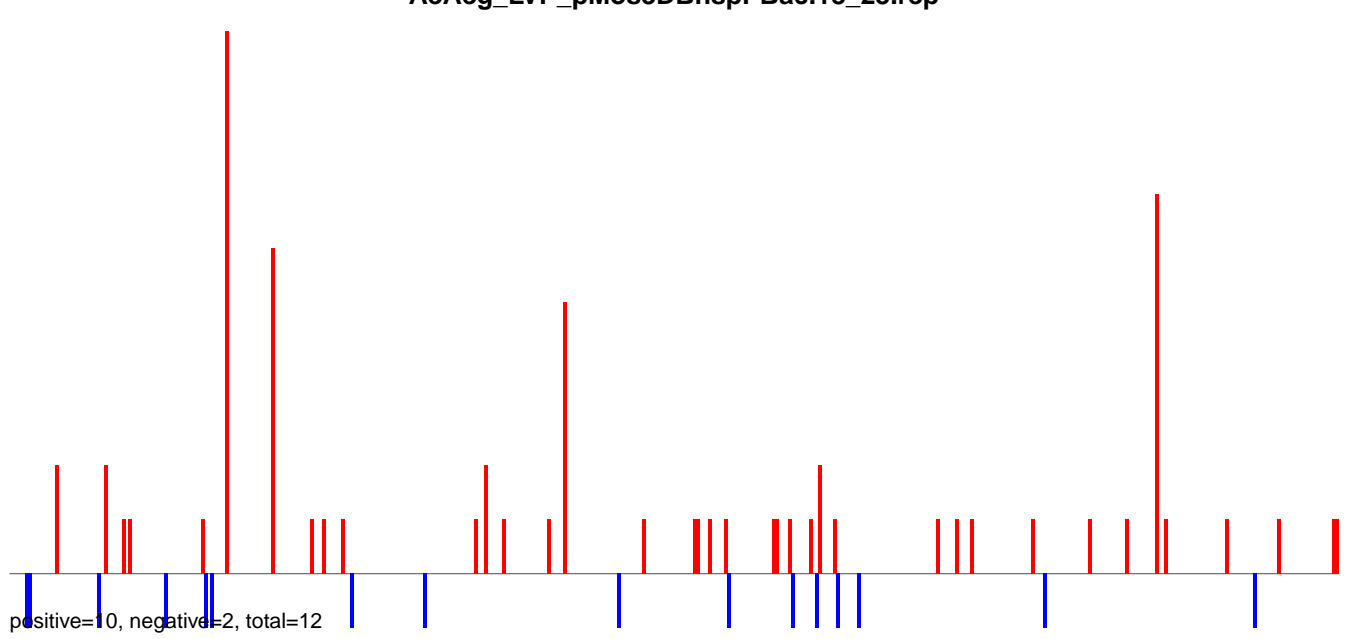
AeAeg_LVP_pMos3DBhspPBac.rep



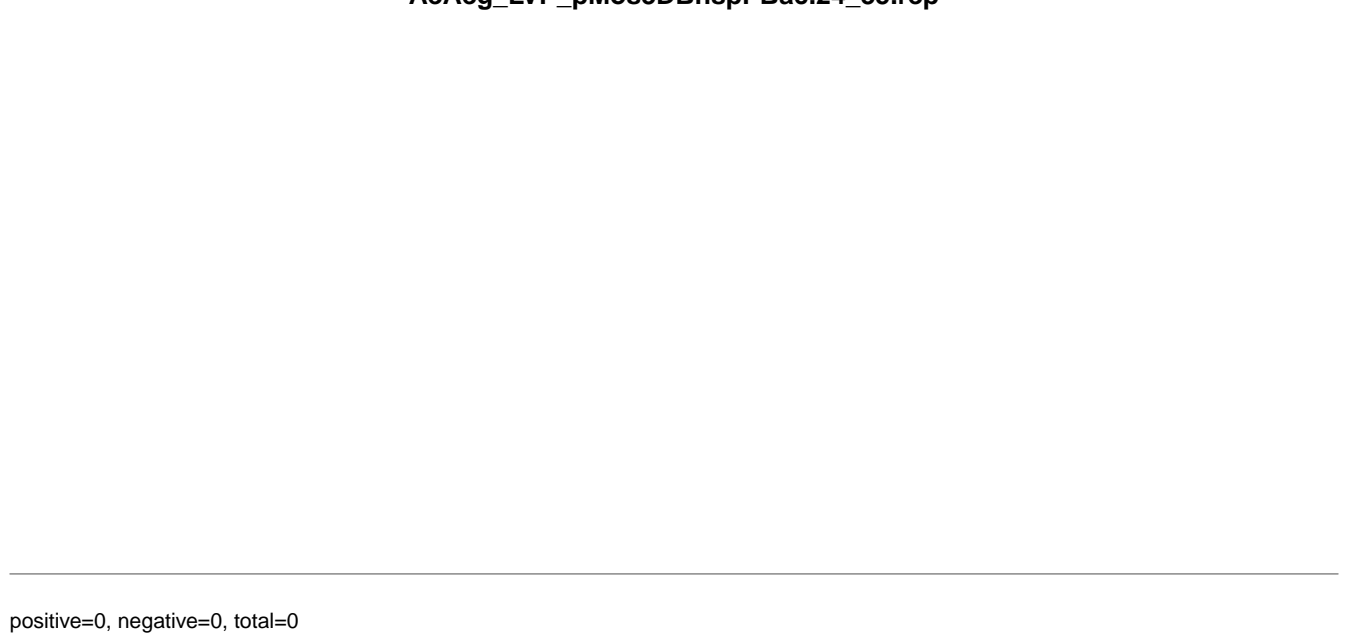
Window size=25, length=10735, virus@NC_001477.1-Dengue_virus_1:1-10735

0 2000 4000 6000 8000 10000

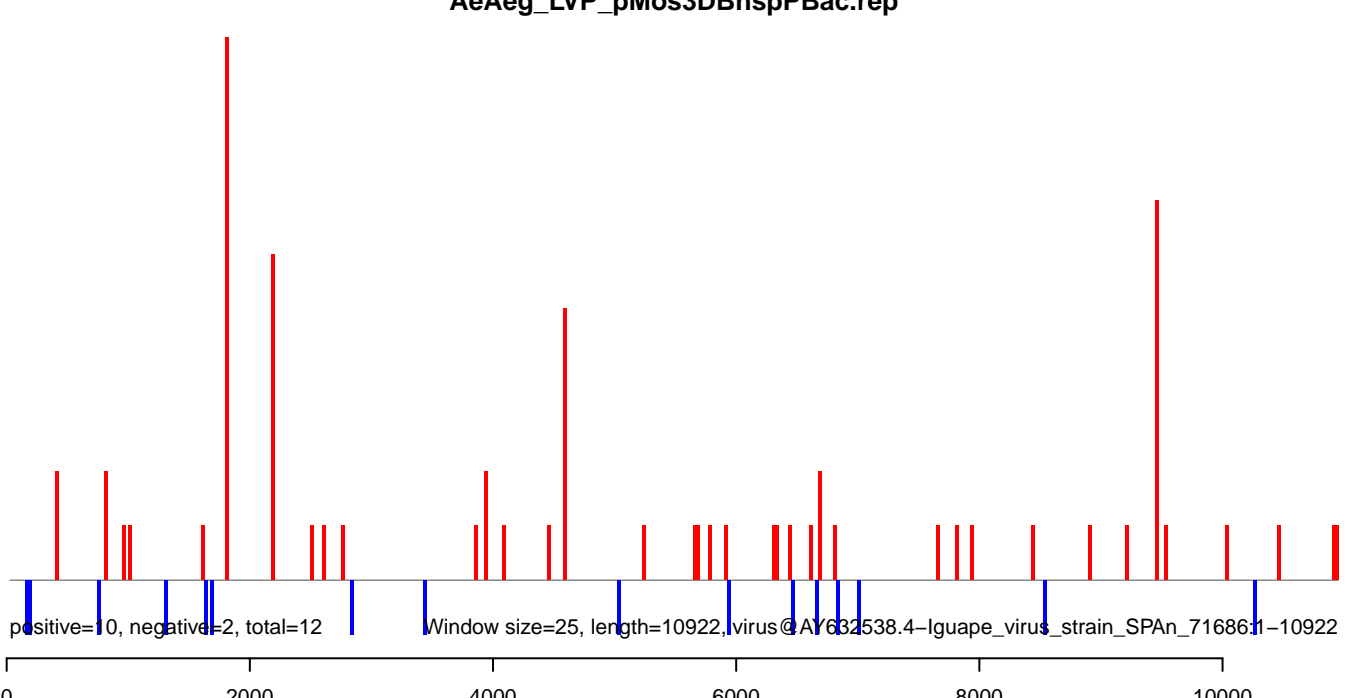
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



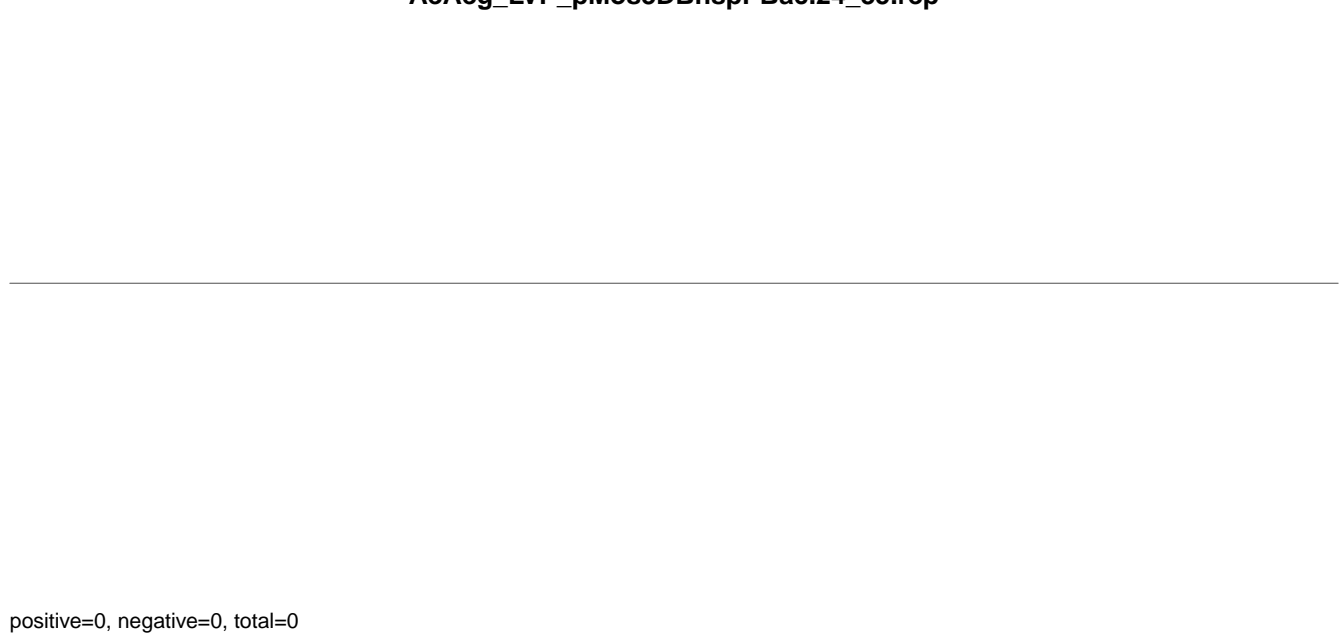
AeAeg_LVP_pMos3DBhspPBac.rep



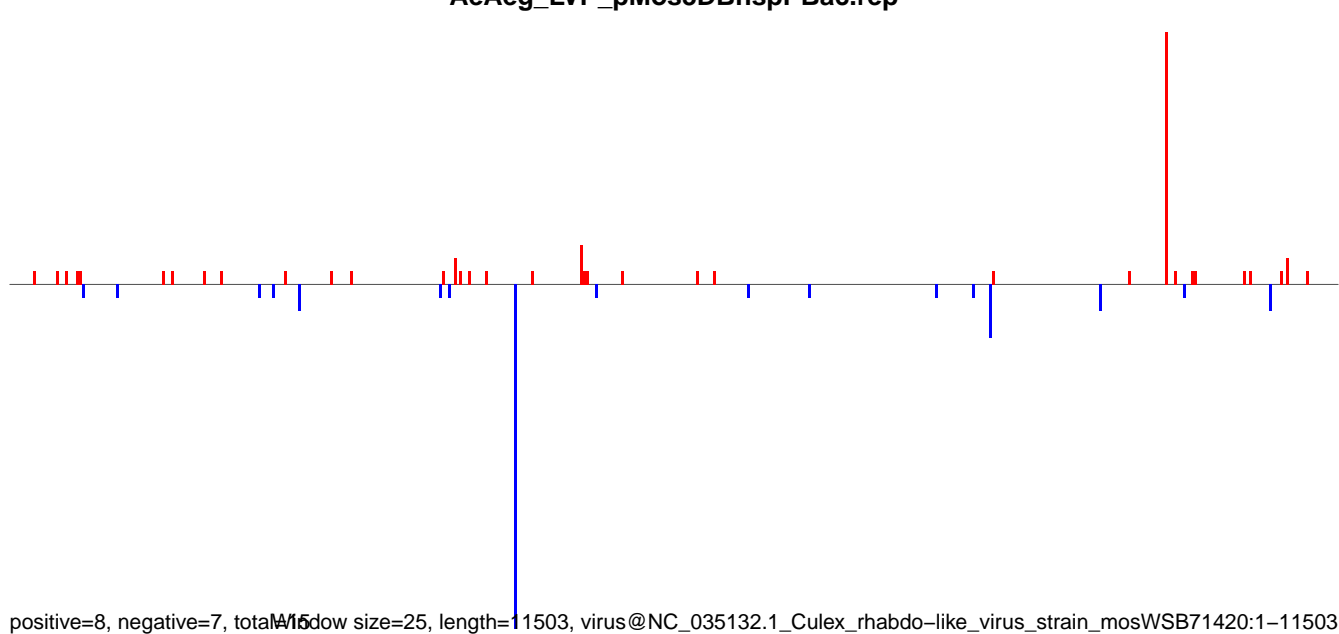
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

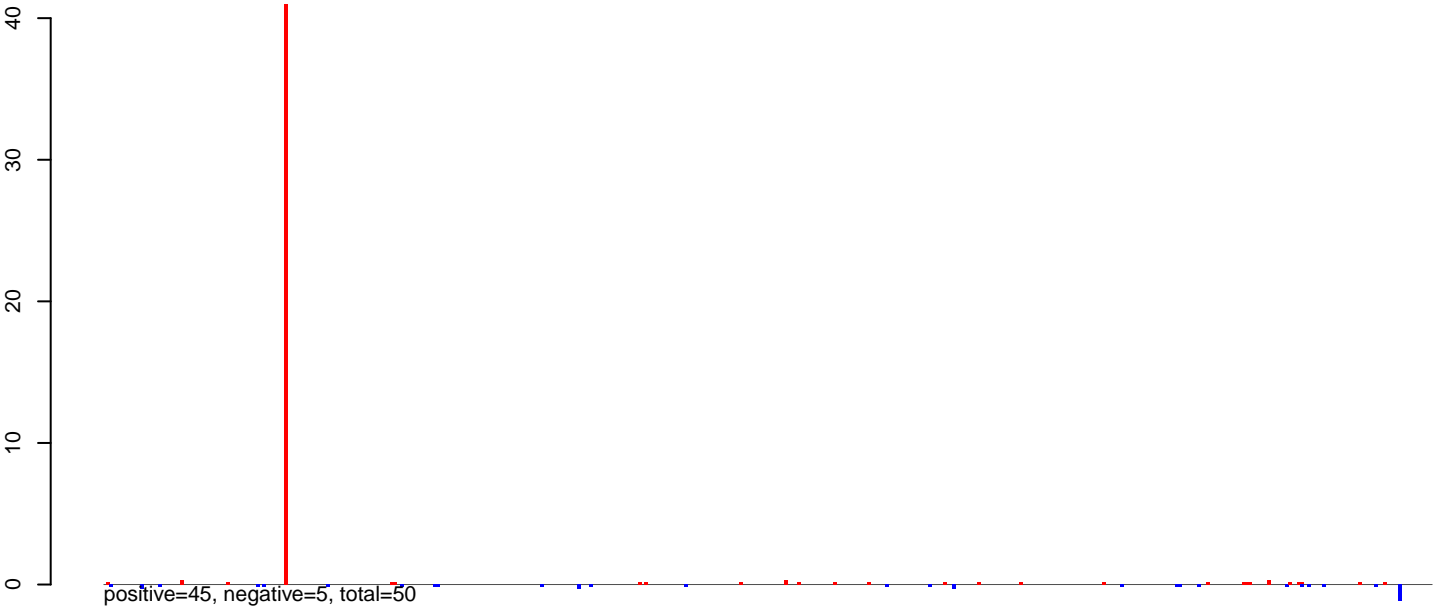


AeAeg_LVP_pMos3DBhspPBac.rep

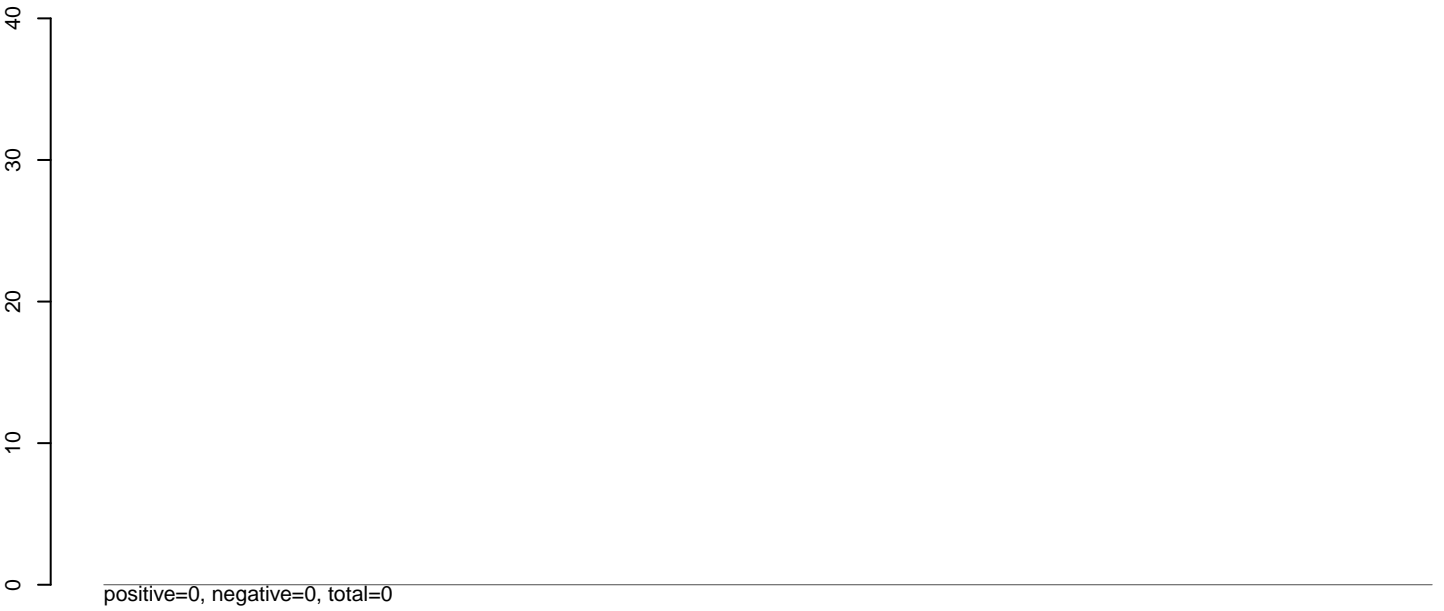


Window size=25, length=11503, virus@NC_035132.1_Culex_rhabdo-like_virus_strain_mosWSB71420:1-11503

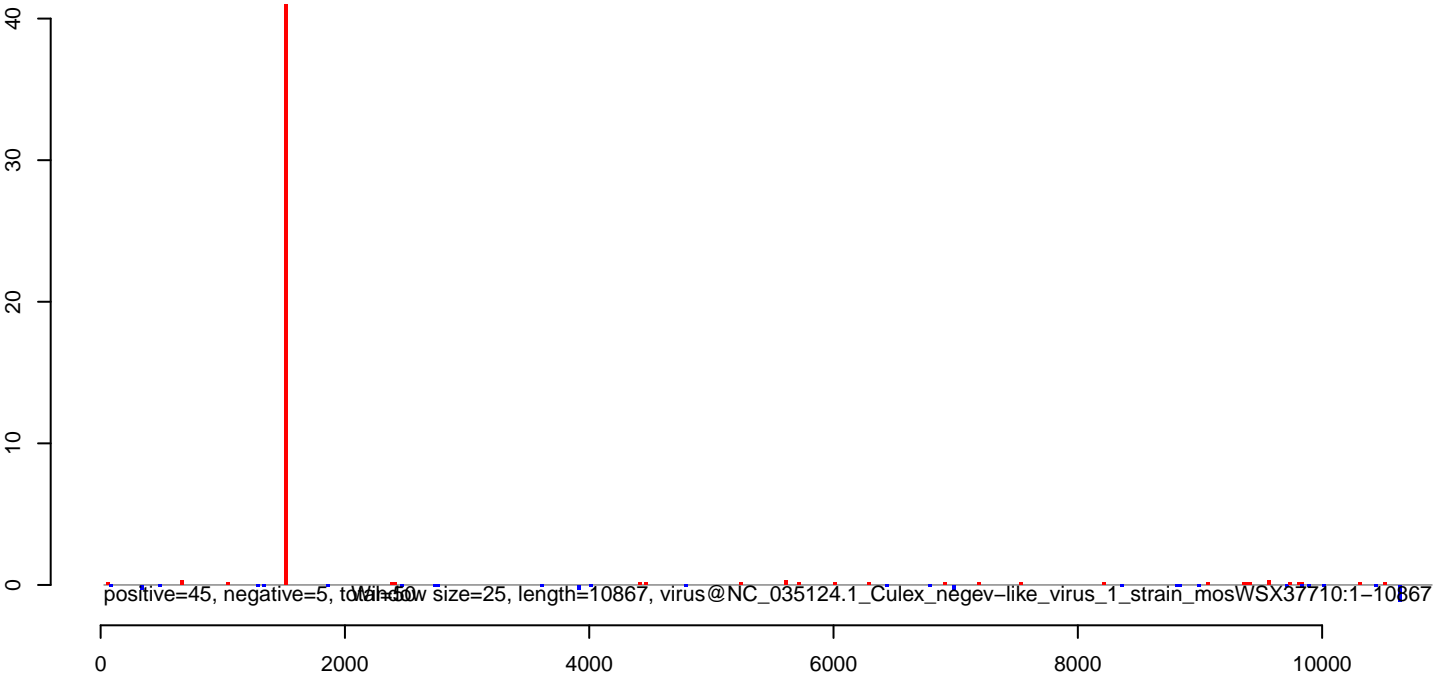
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



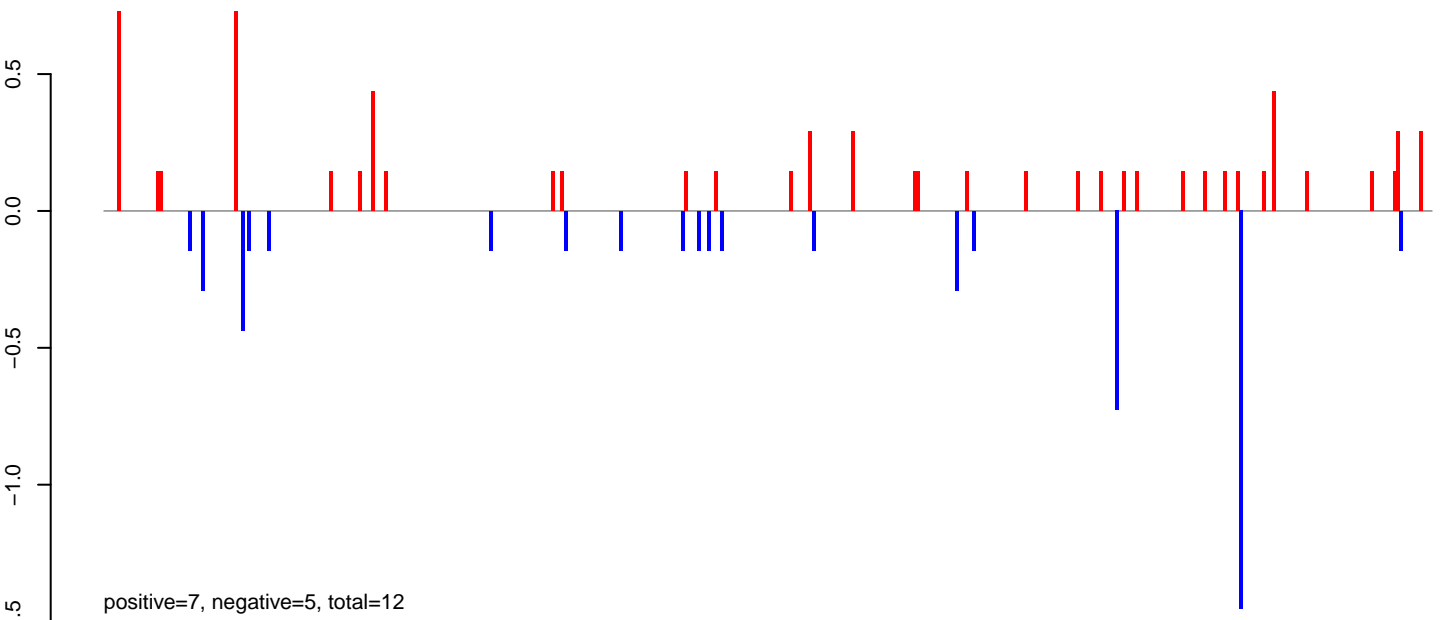
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



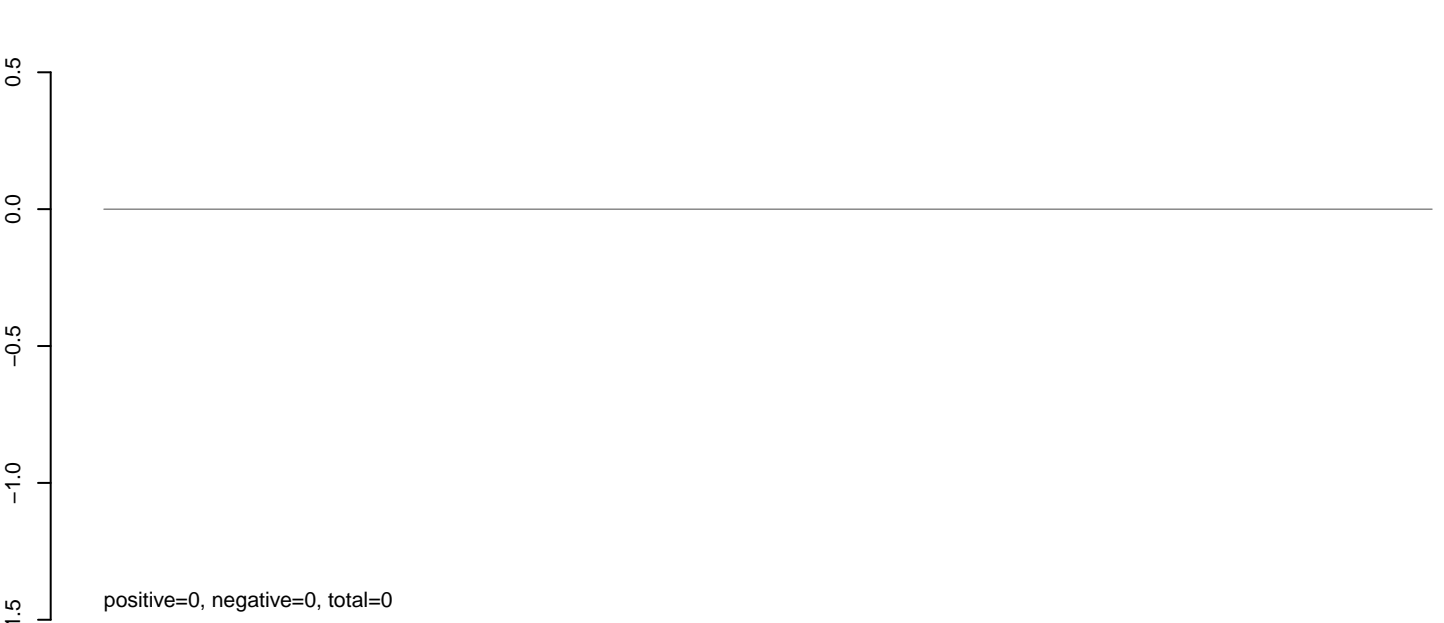
AeAeg_LVP_pMos3DBhspPBac.rep



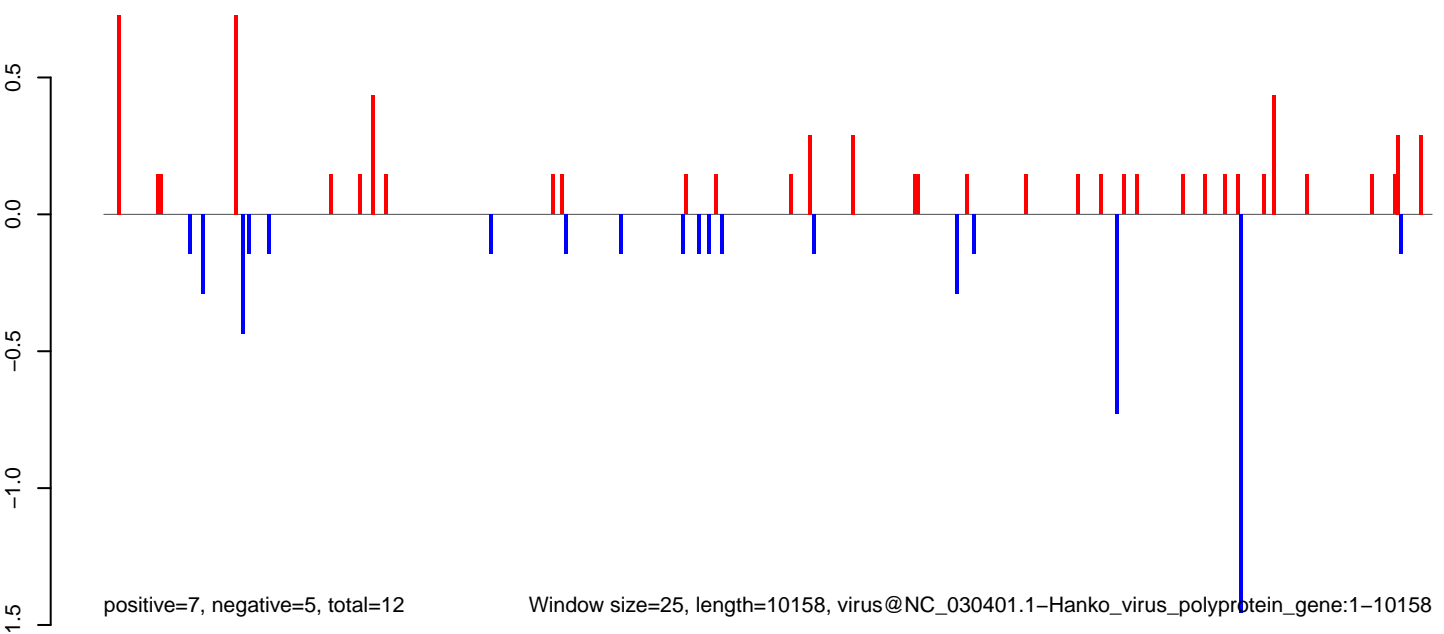
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



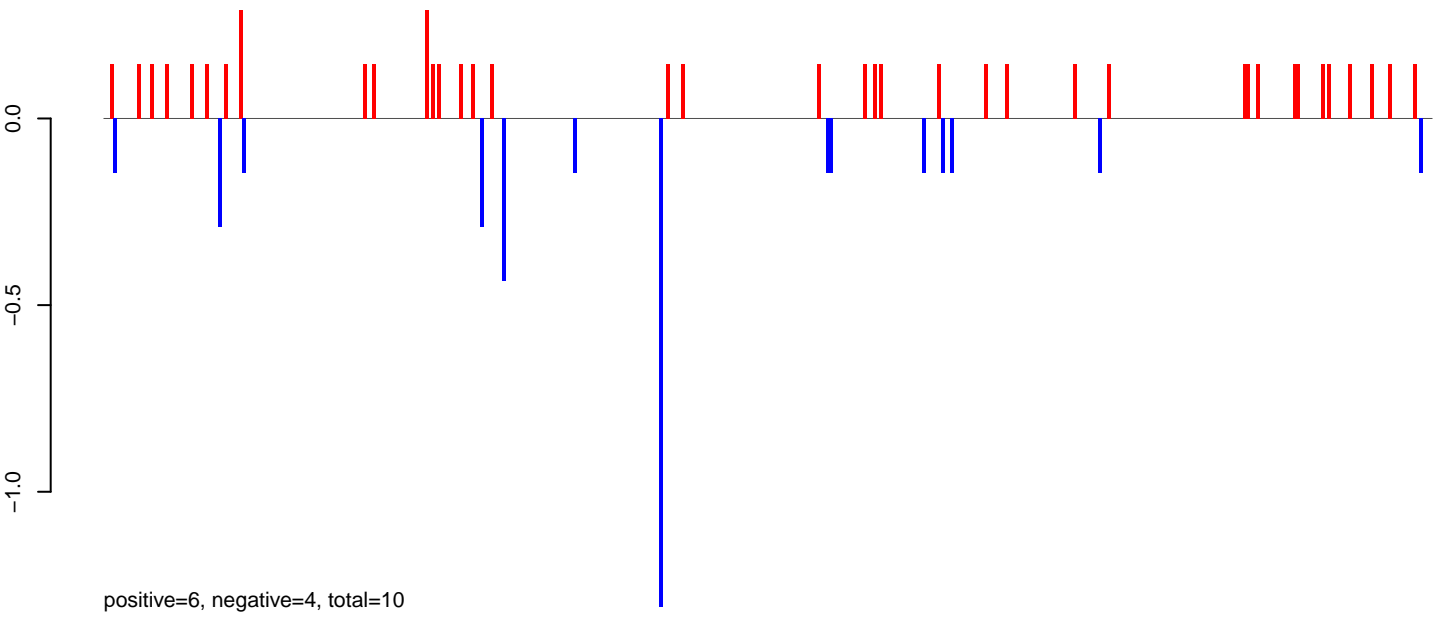
AeAeg_LVP_pMos3DBhspPBac.rep



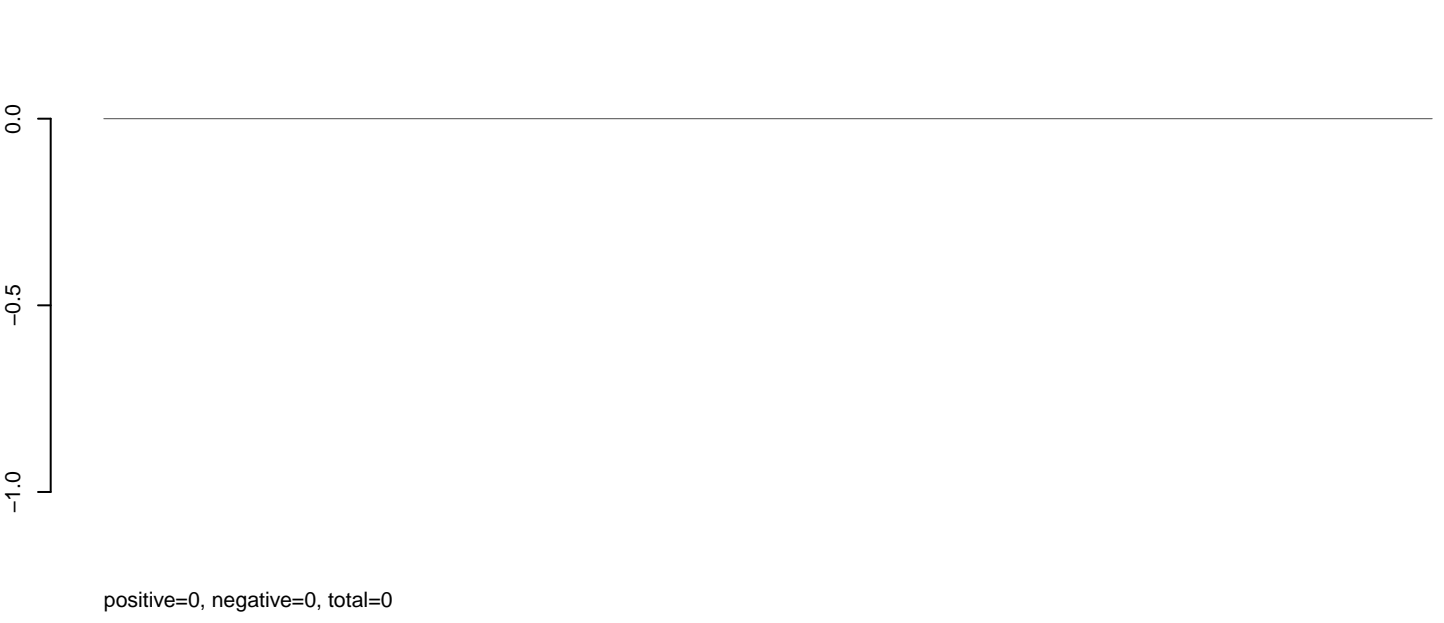
Window size=25, length=10158, virus@NC_030401.1-Hanko_virus_polyprotein_gene:1-10158

0 2000 4000 6000 8000 10000

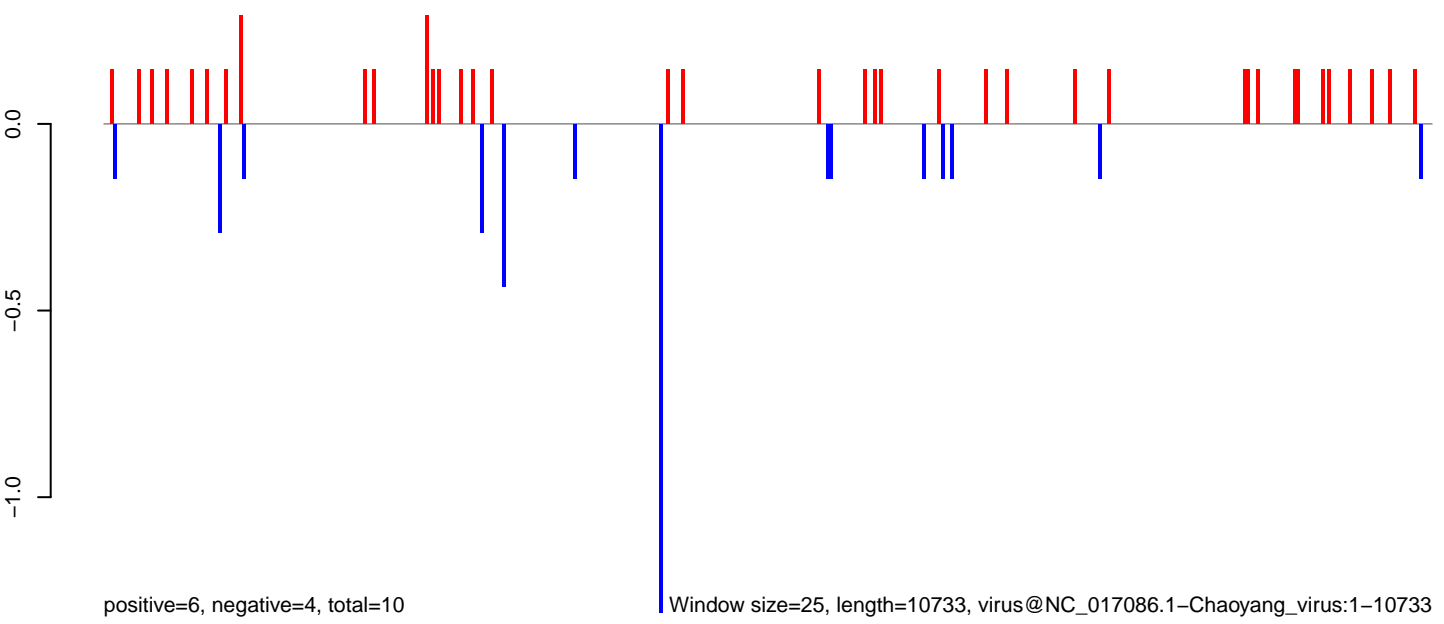
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

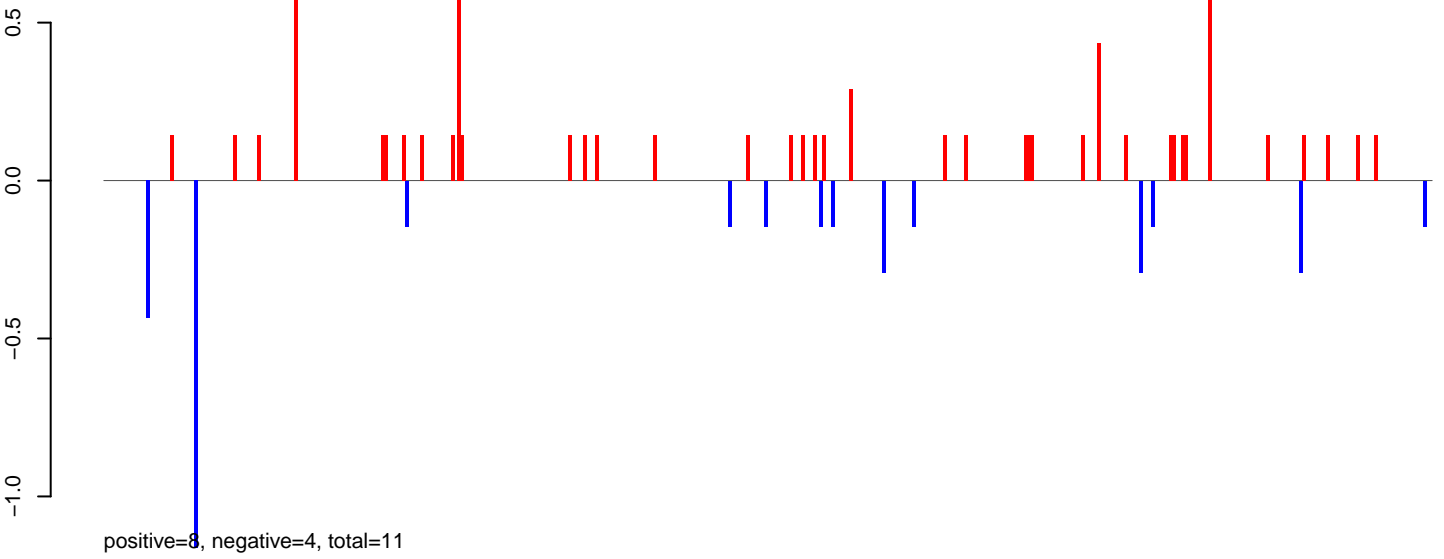


AeAeg_LVP_pMos3DBhspPBac.rep

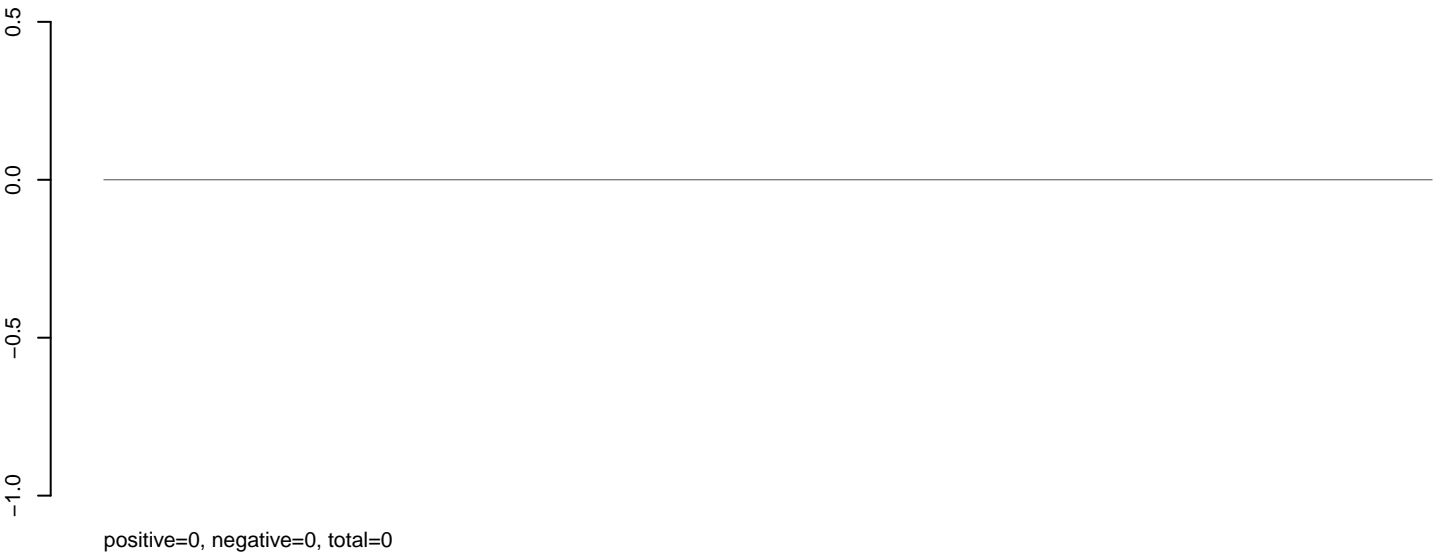


0 2000 4000 6000 8000 10000

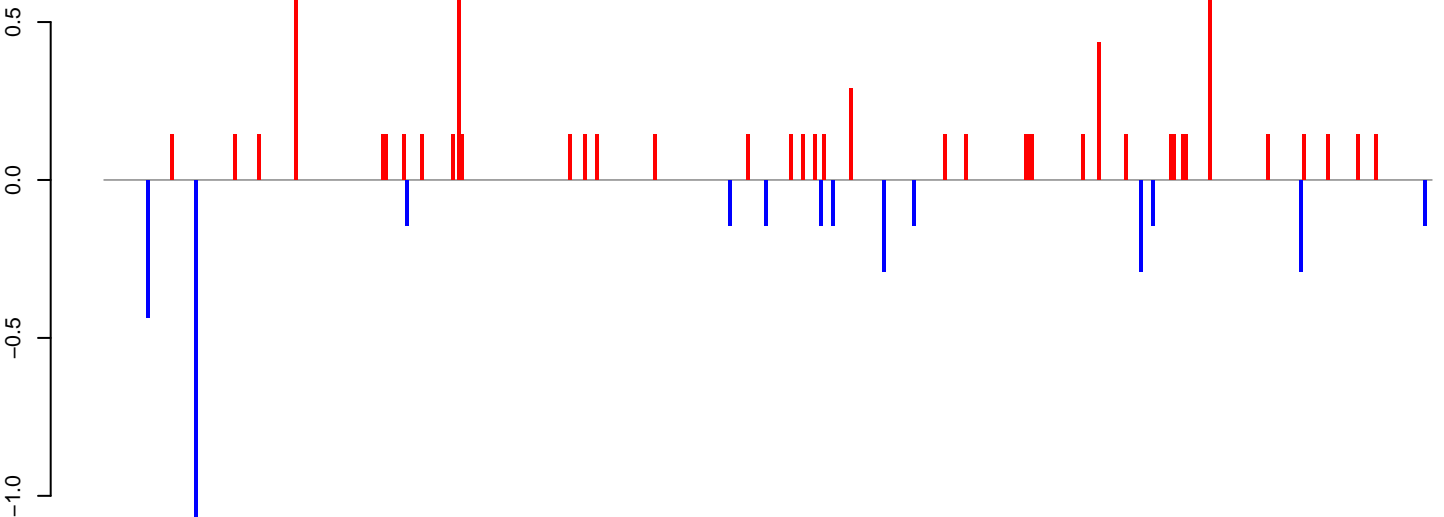
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



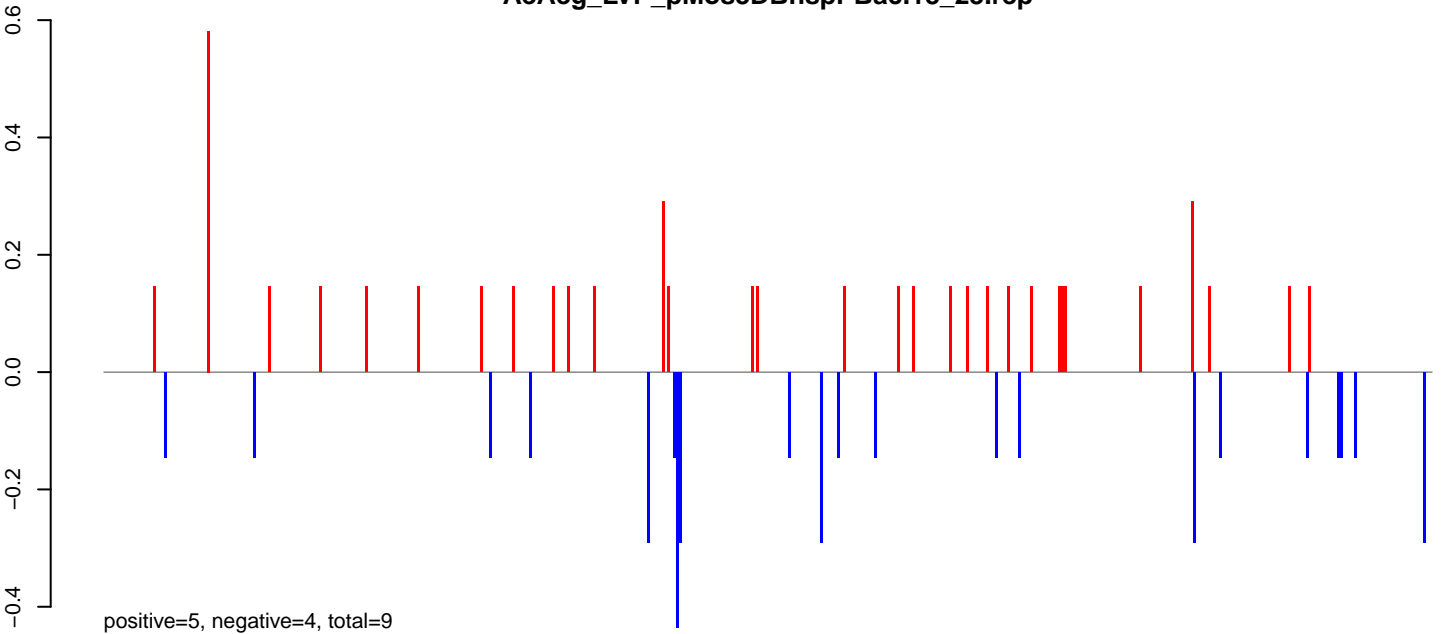
AeAeg_LVP_pMos3DBhspPBac.rep



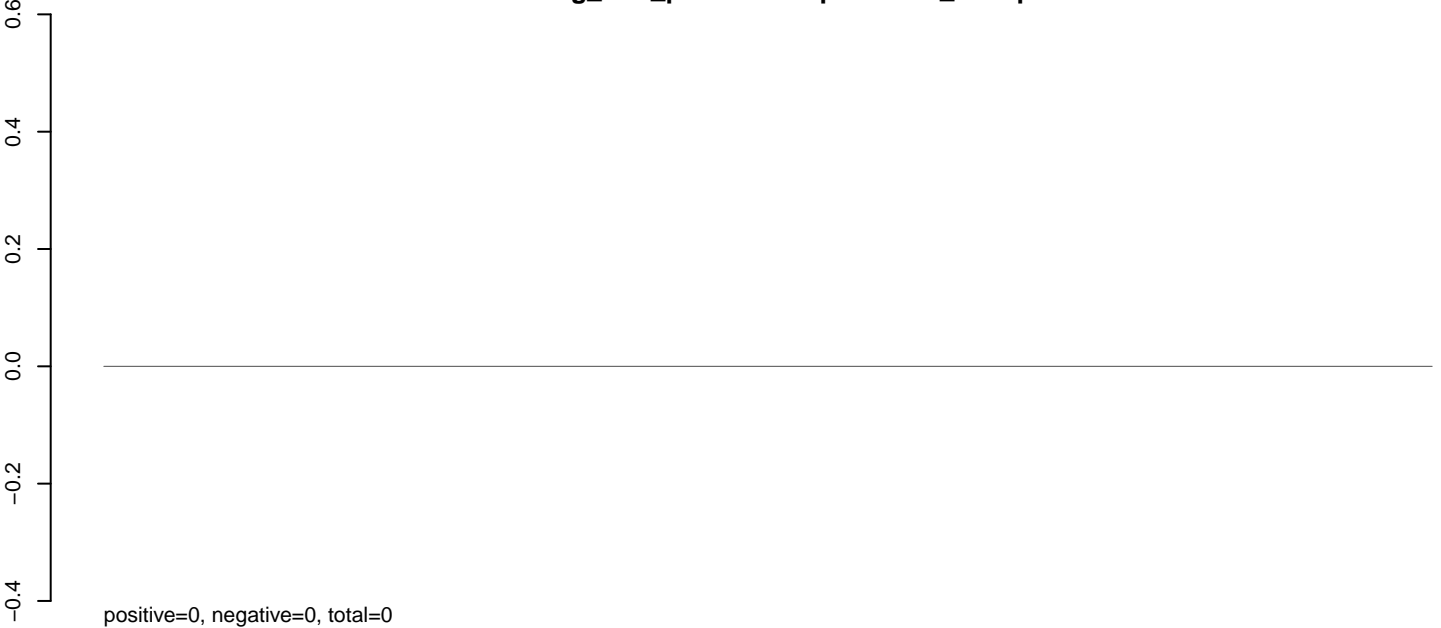
Widowsigat25-4, length=10990, virus@NC_015843.2-Tembusu_virus_flavivirus_polyprotein_flavivirus_polyprotein_gene_gene:1-10990

0 2000 4000 6000 8000 10000

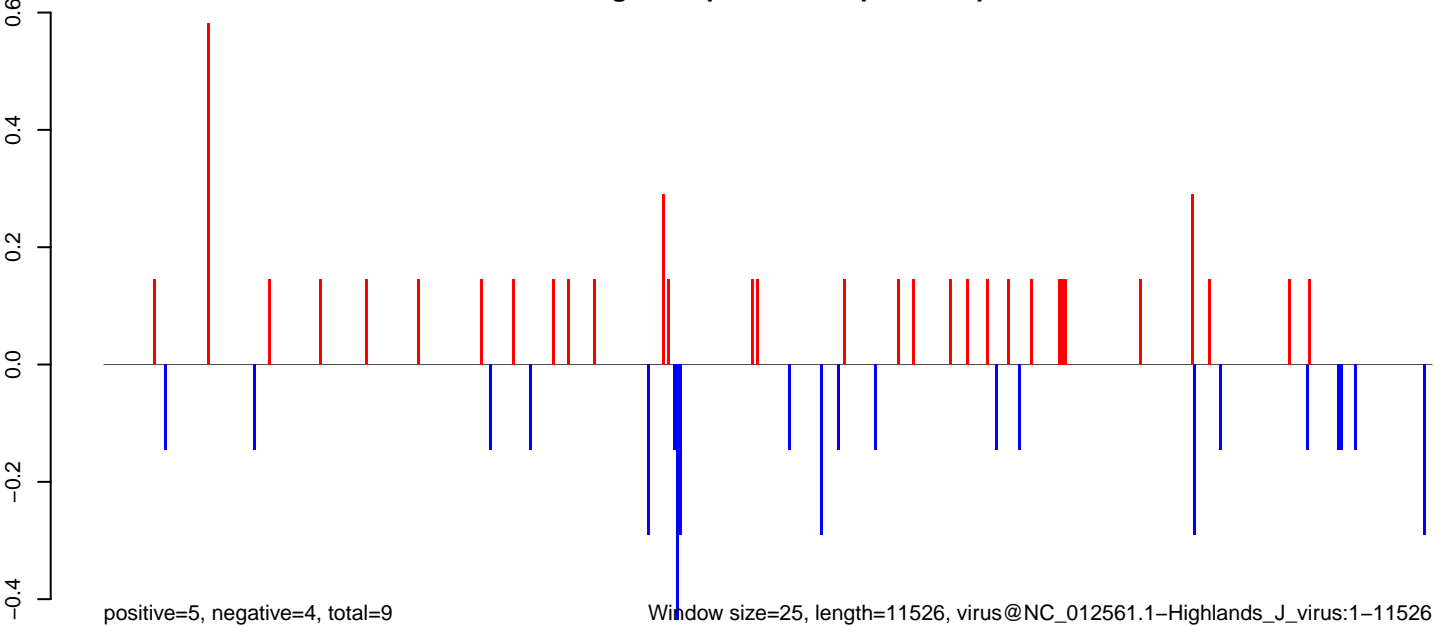
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

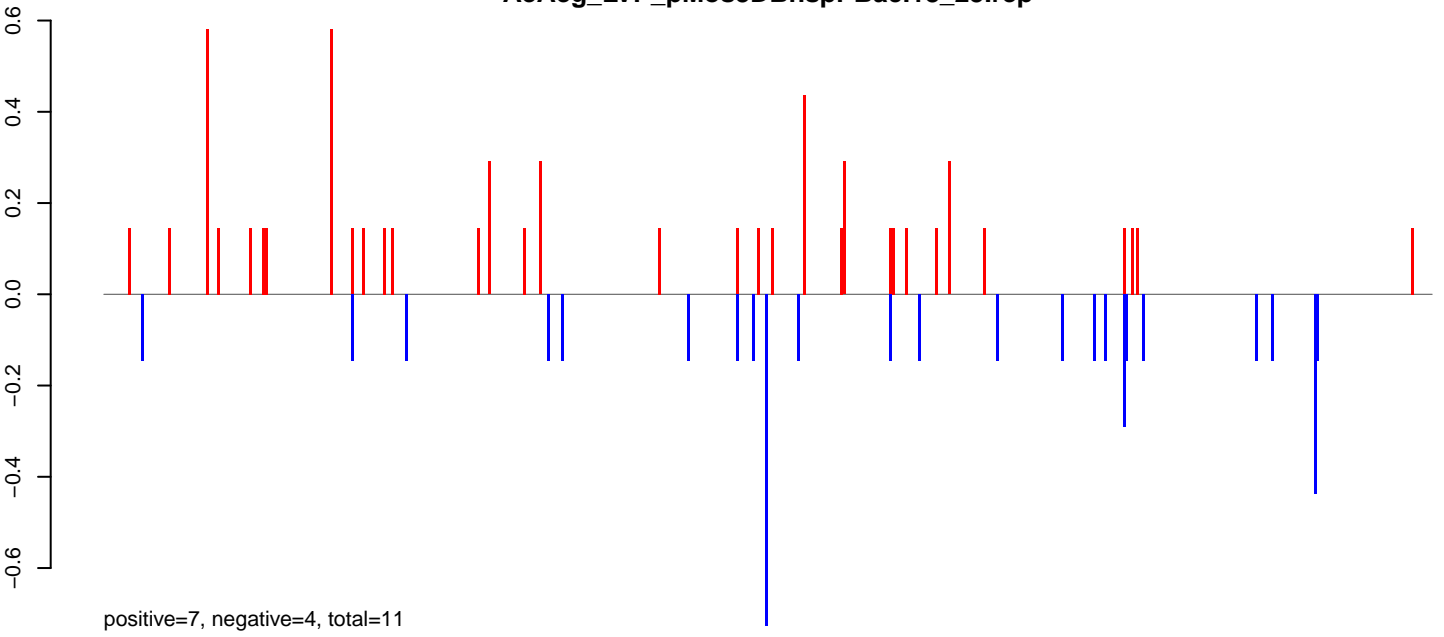


AeAeg_LVP_pMos3DBhspPBac.rep

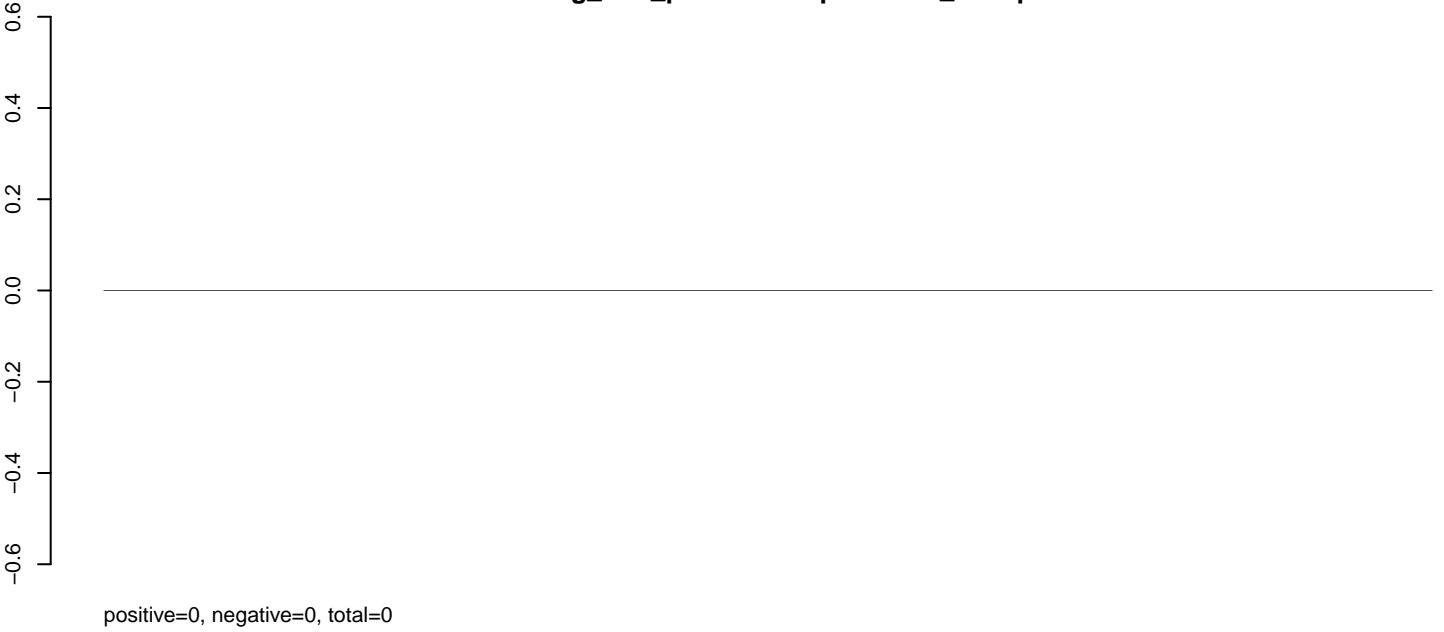


0 2000 4000 6000 8000 10000 12000

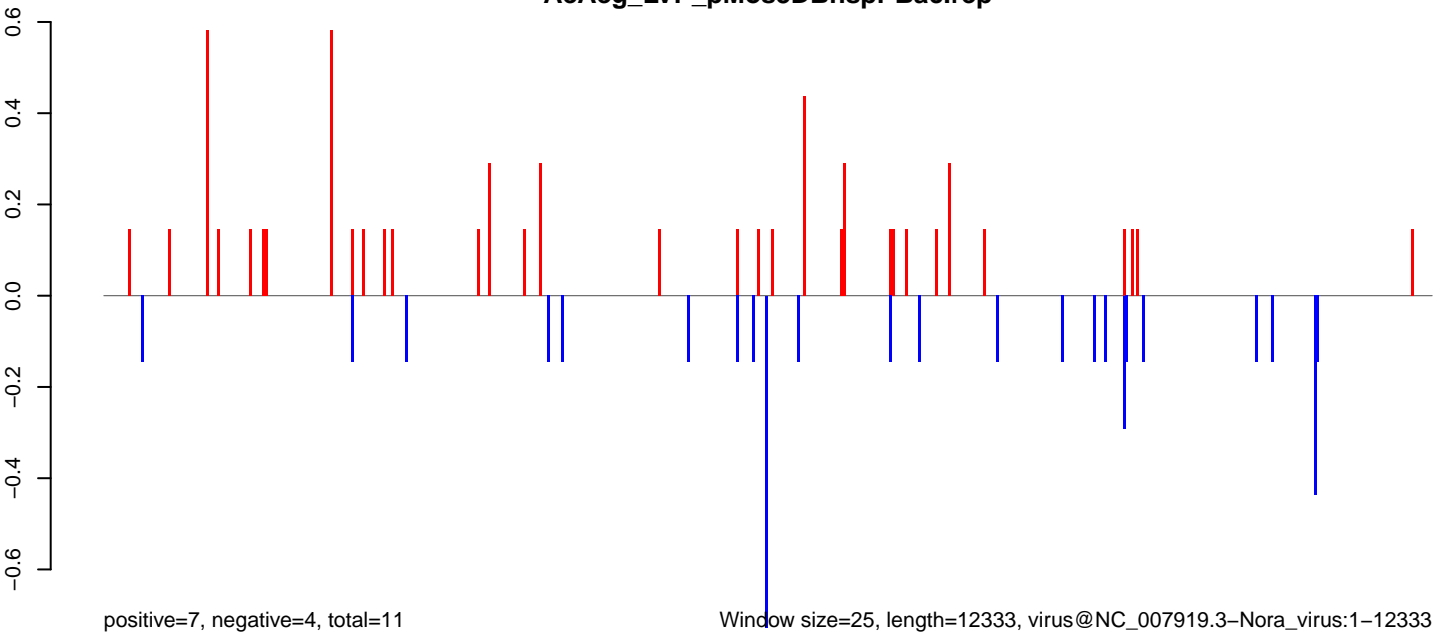
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

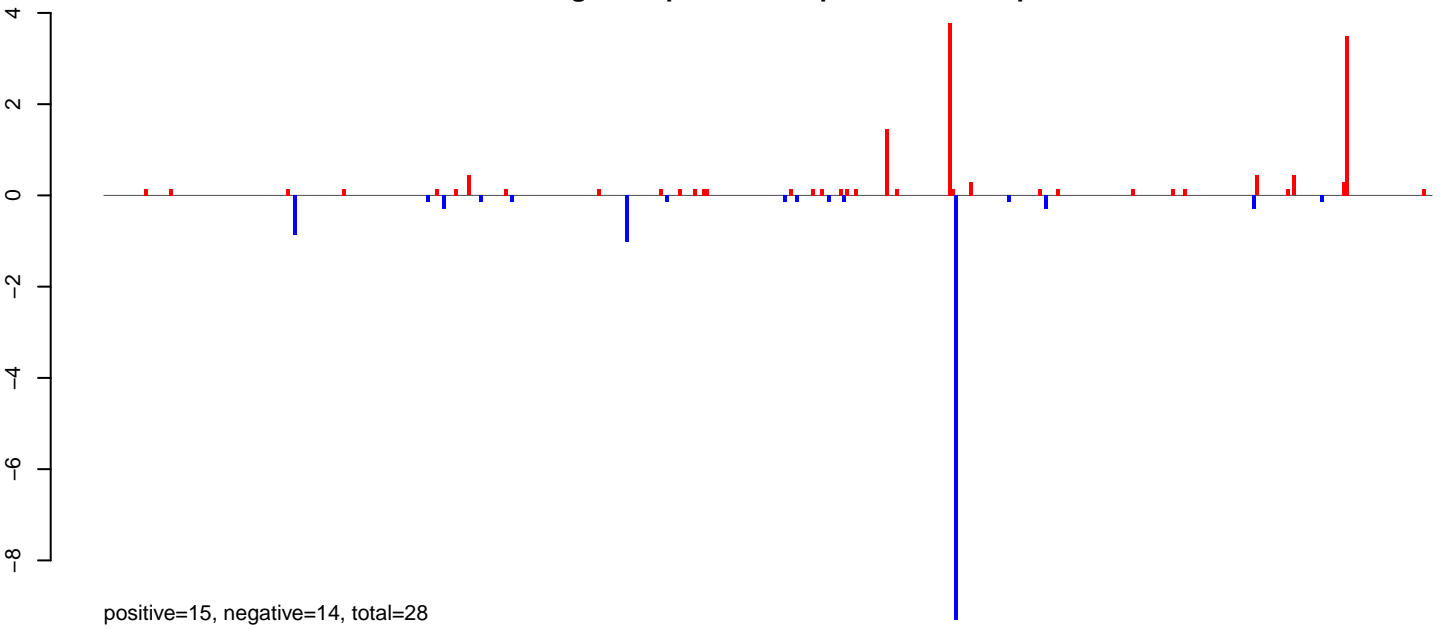


AeAeg_LVP_pMos3DBhspPBac.rep

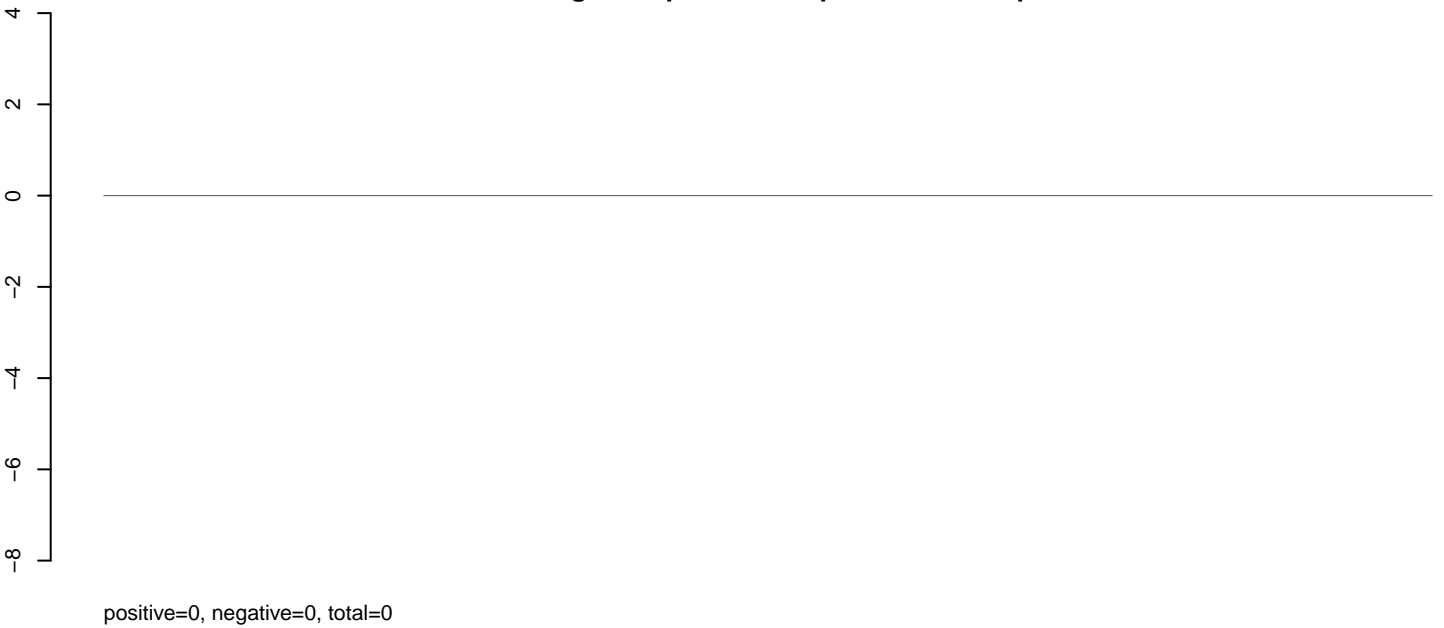


0 2000 4000 6000 8000 10000 12000

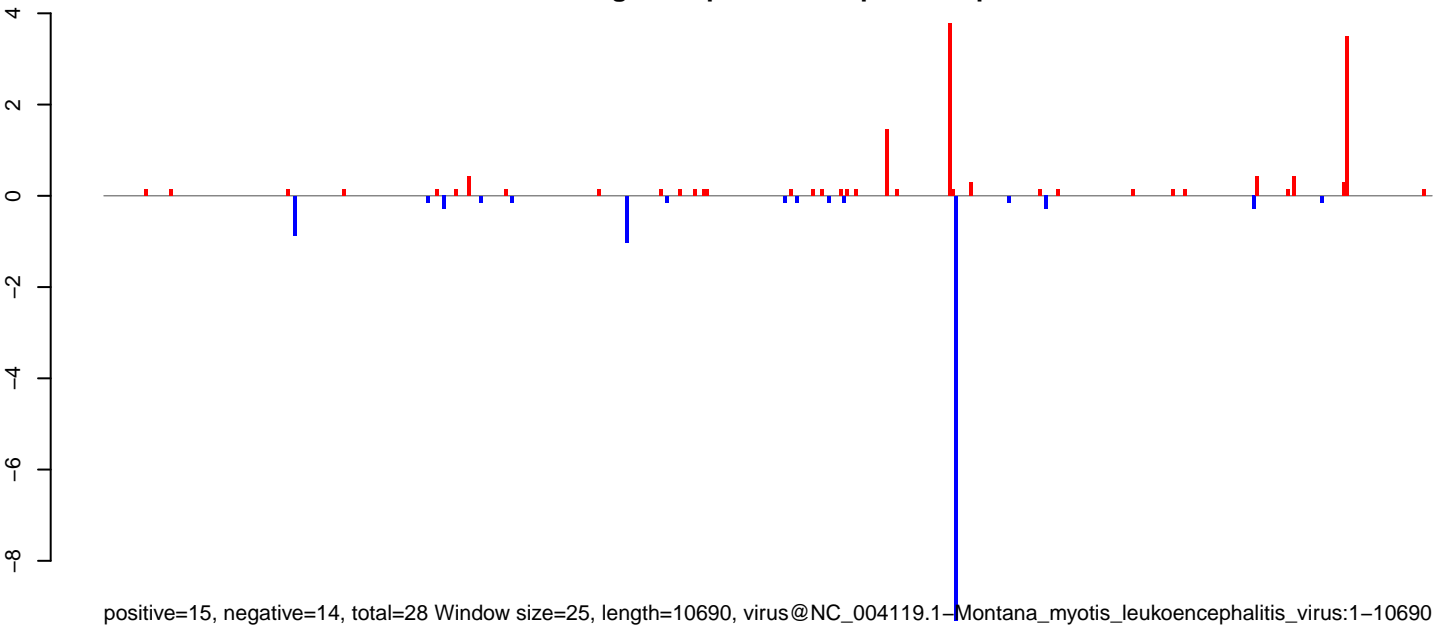
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

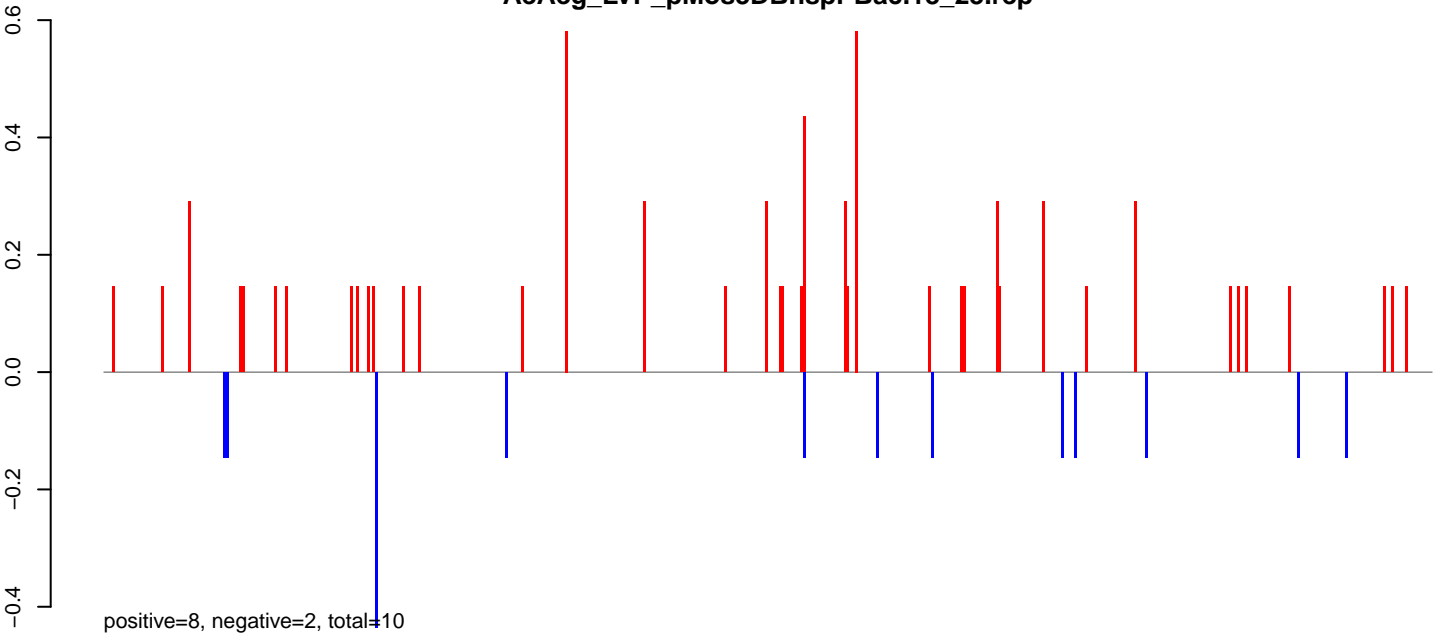


AeAeg_LVP_pMos3DBhspPBac.rep

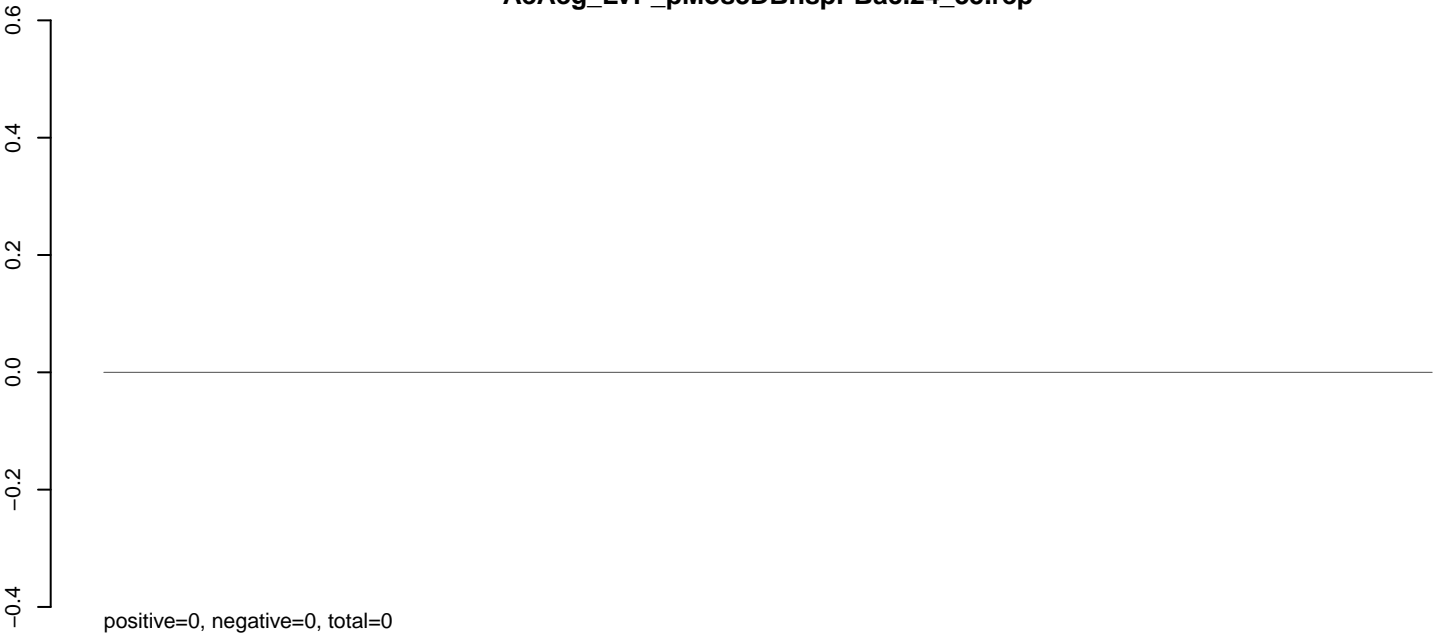


0 2000 4000 6000 8000 10000

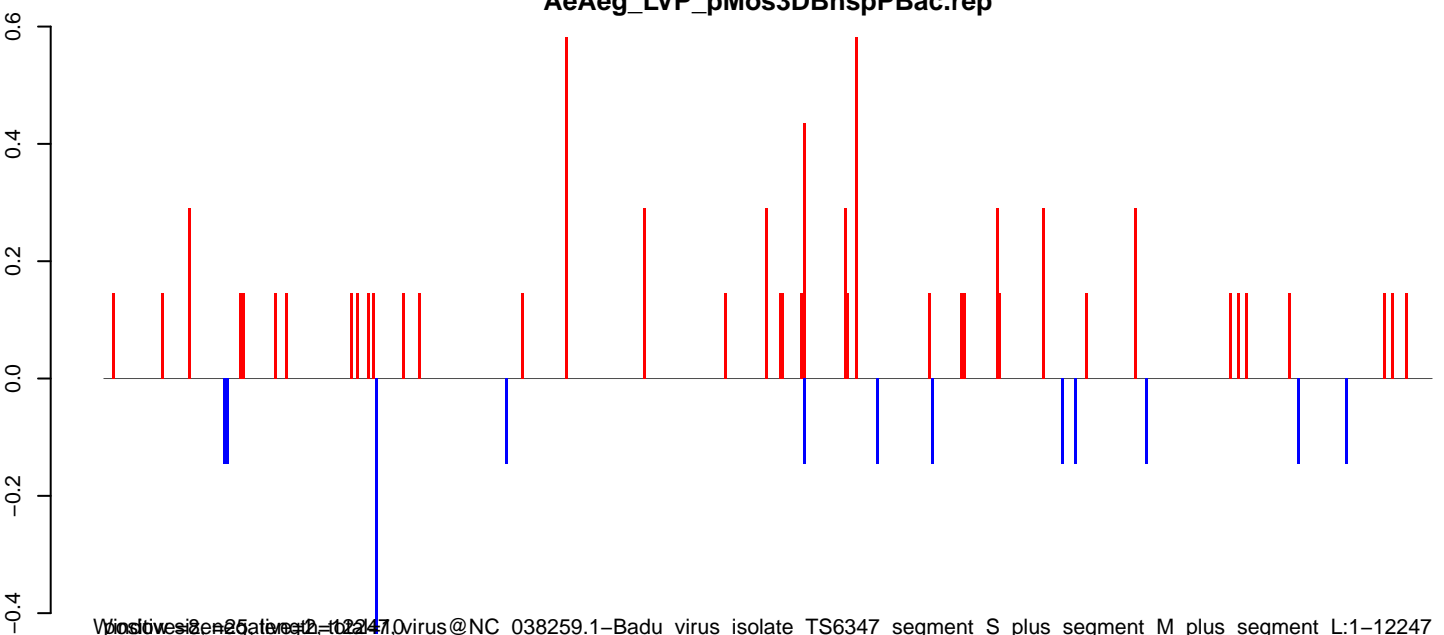
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



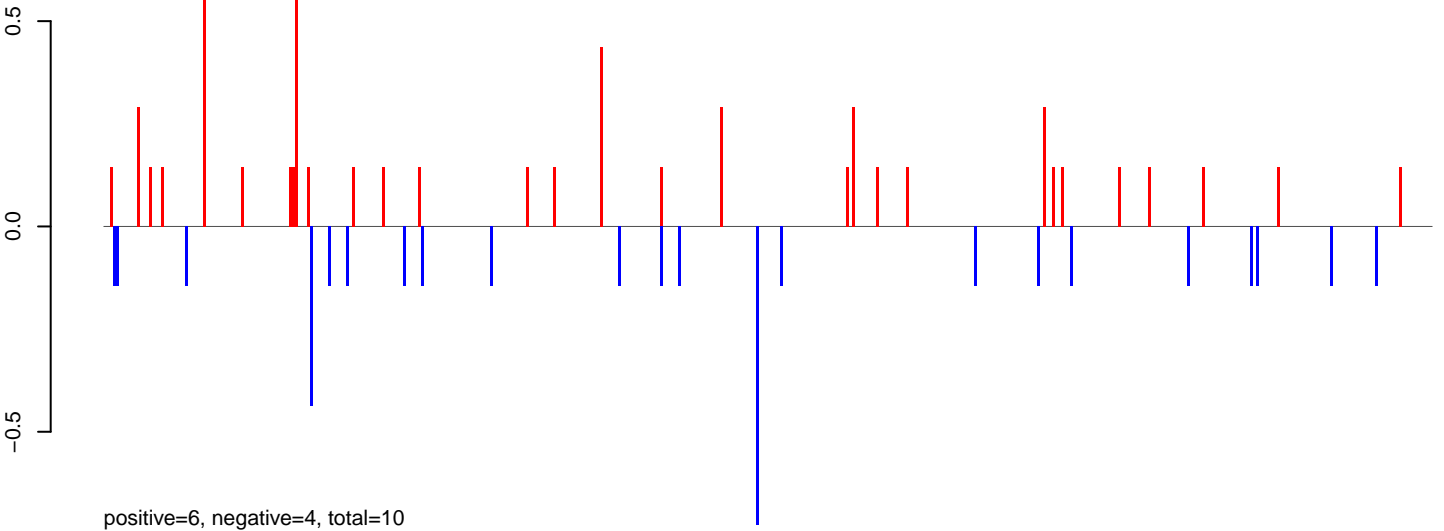
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



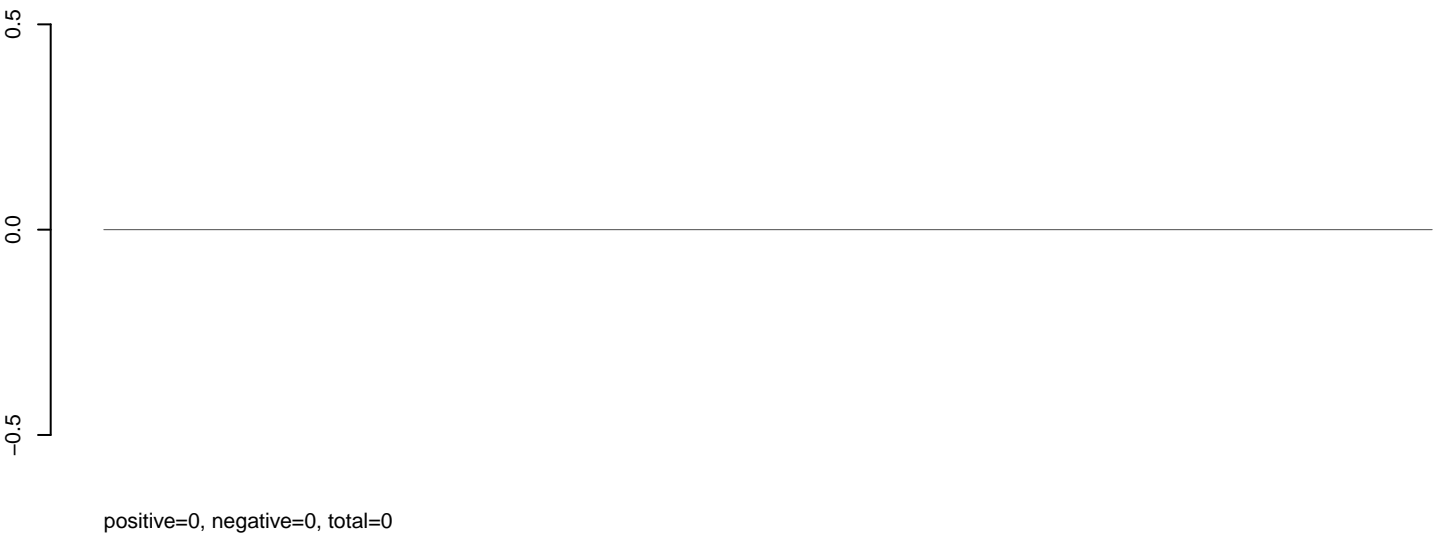
AeAeg_LVP_pMos3DBhspPBac.rep



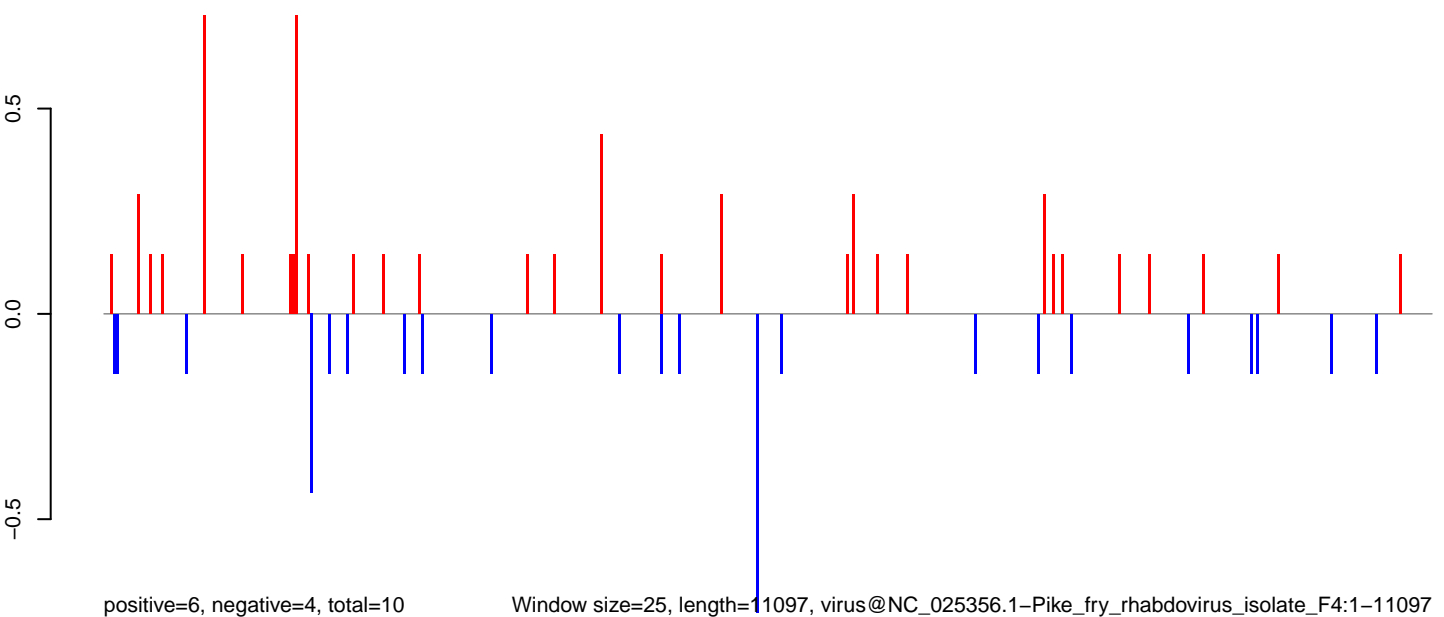
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

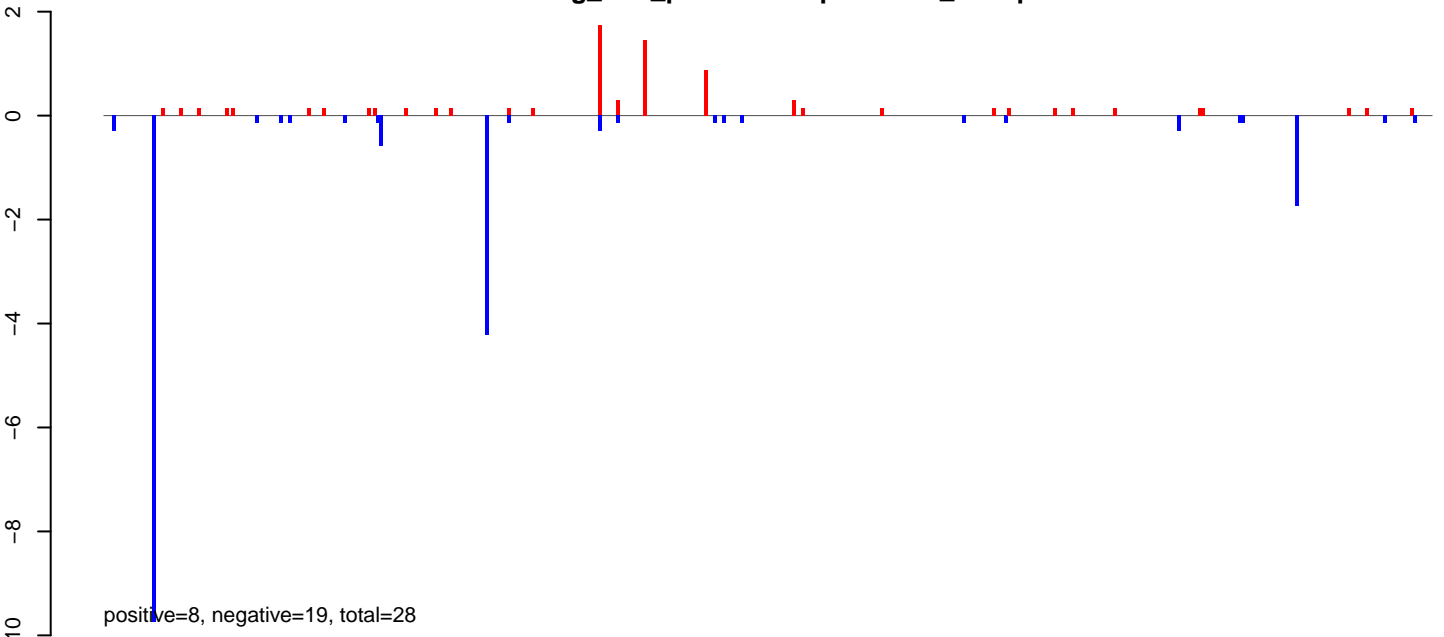


AeAeg_LVP_pMos3DBhspPBac.rep

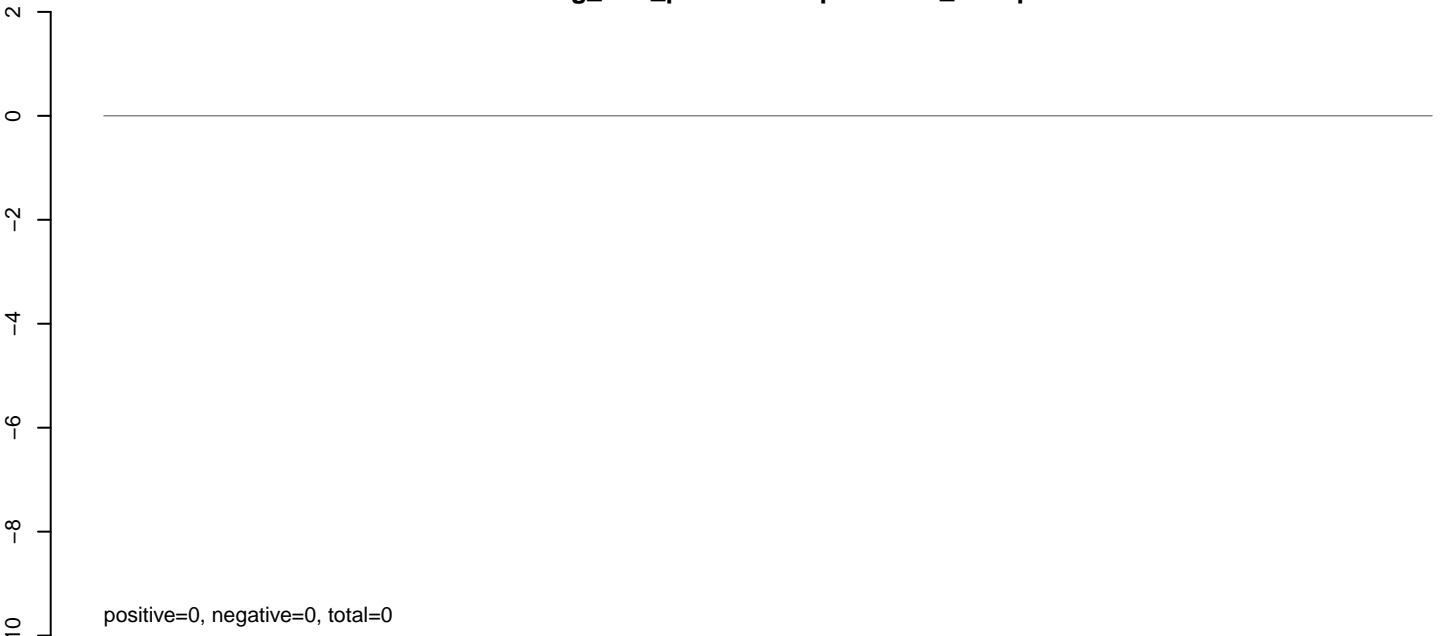


0 2000 4000 6000 8000 10000

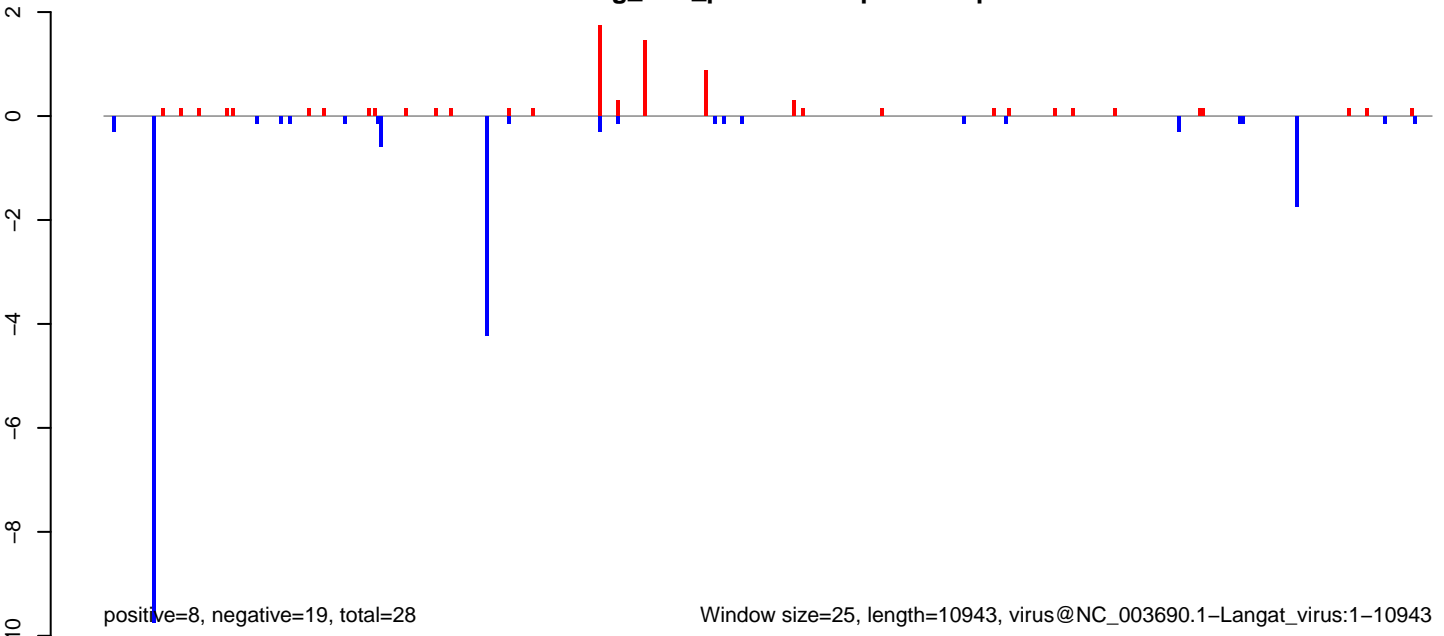
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



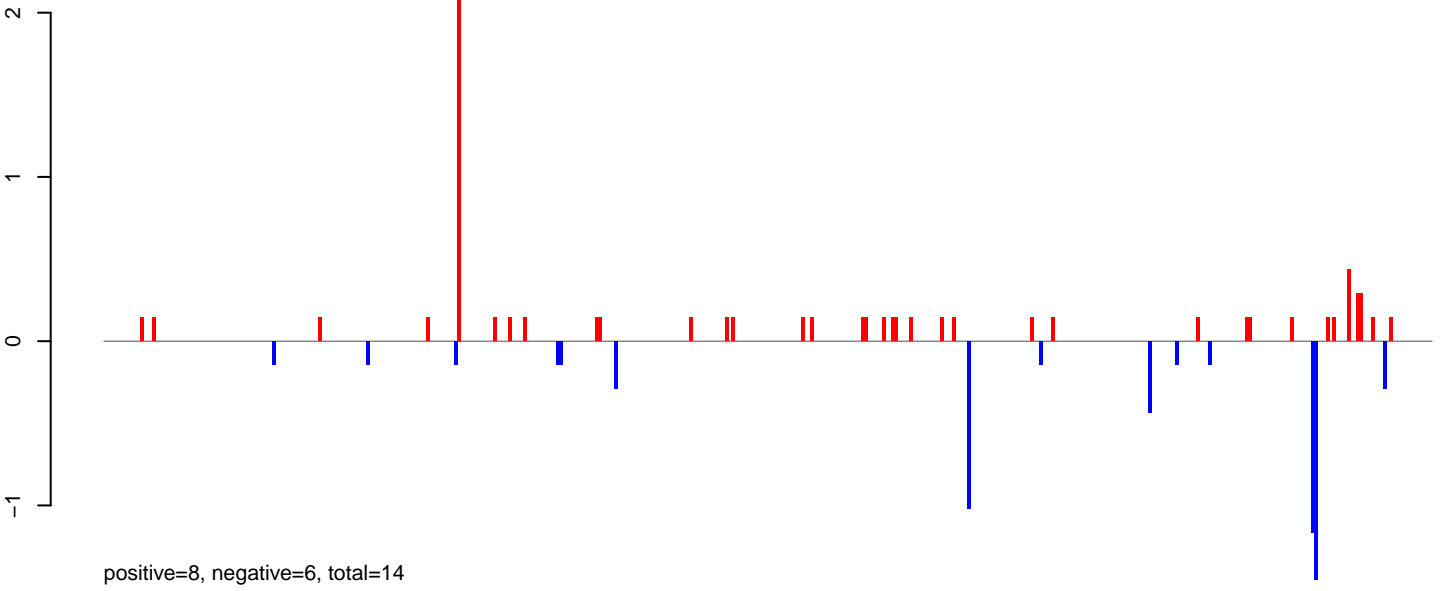
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



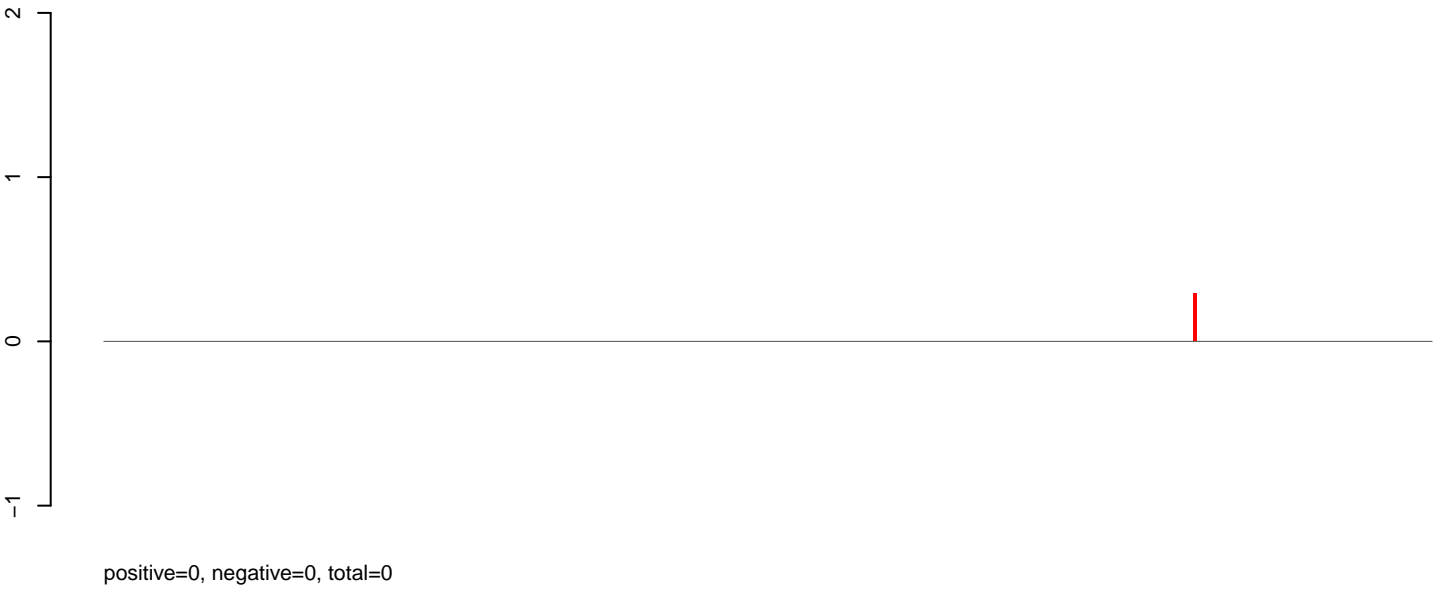
AeAeg_LVP_pMos3DBhspPBac.rep



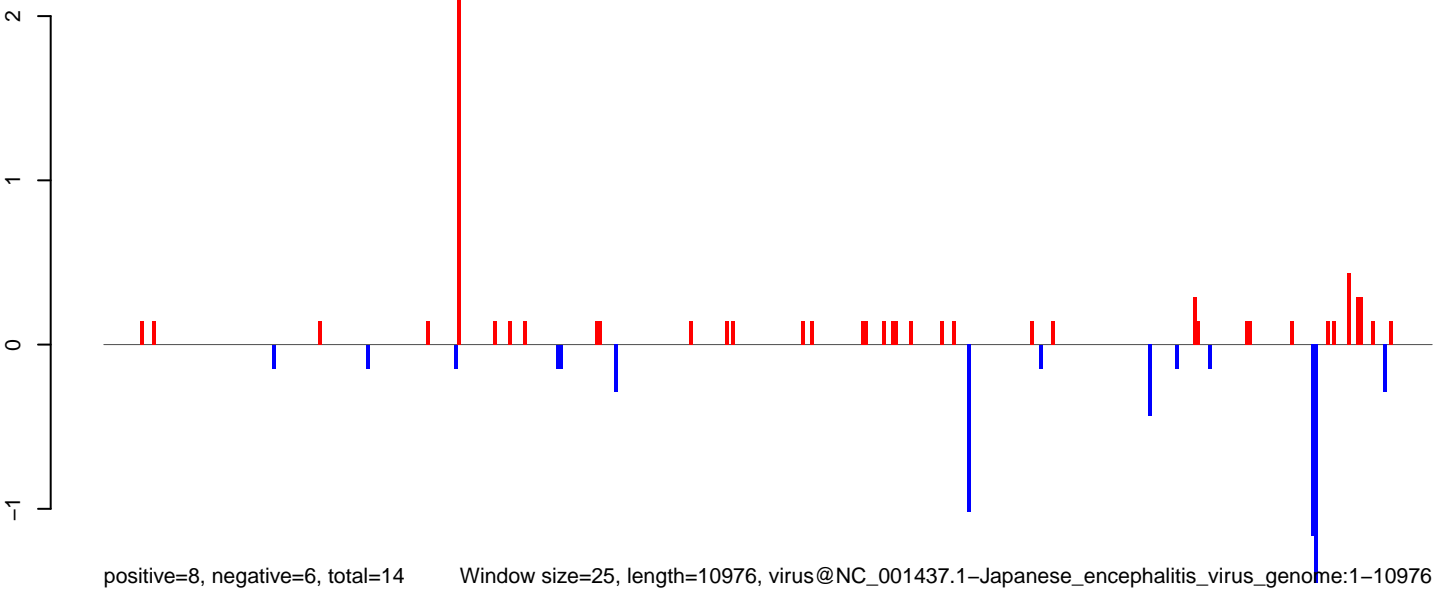
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



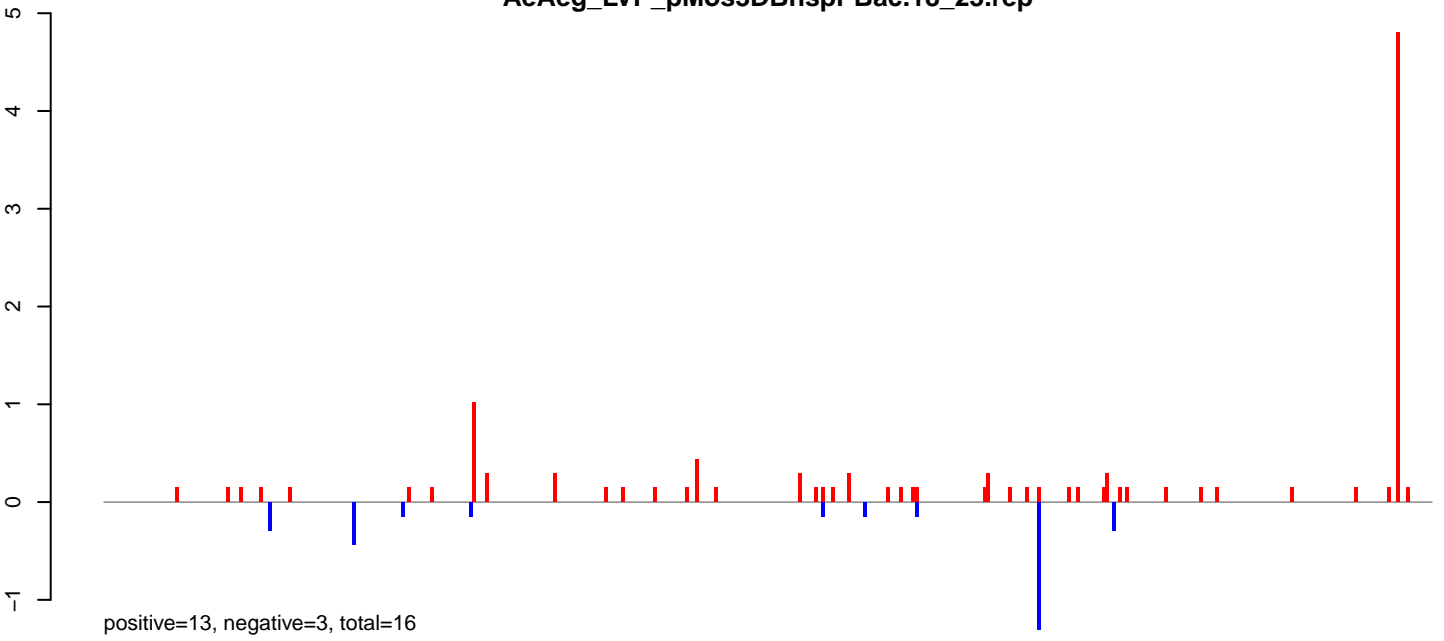
AeAeg_LVP_pMos3DBhspPBac.rep



Window size=25, length=10976, virus@NC_001437.1-Japanese_encephalitis_virus_genome:1-10976

0 2000 4000 6000 8000 10000

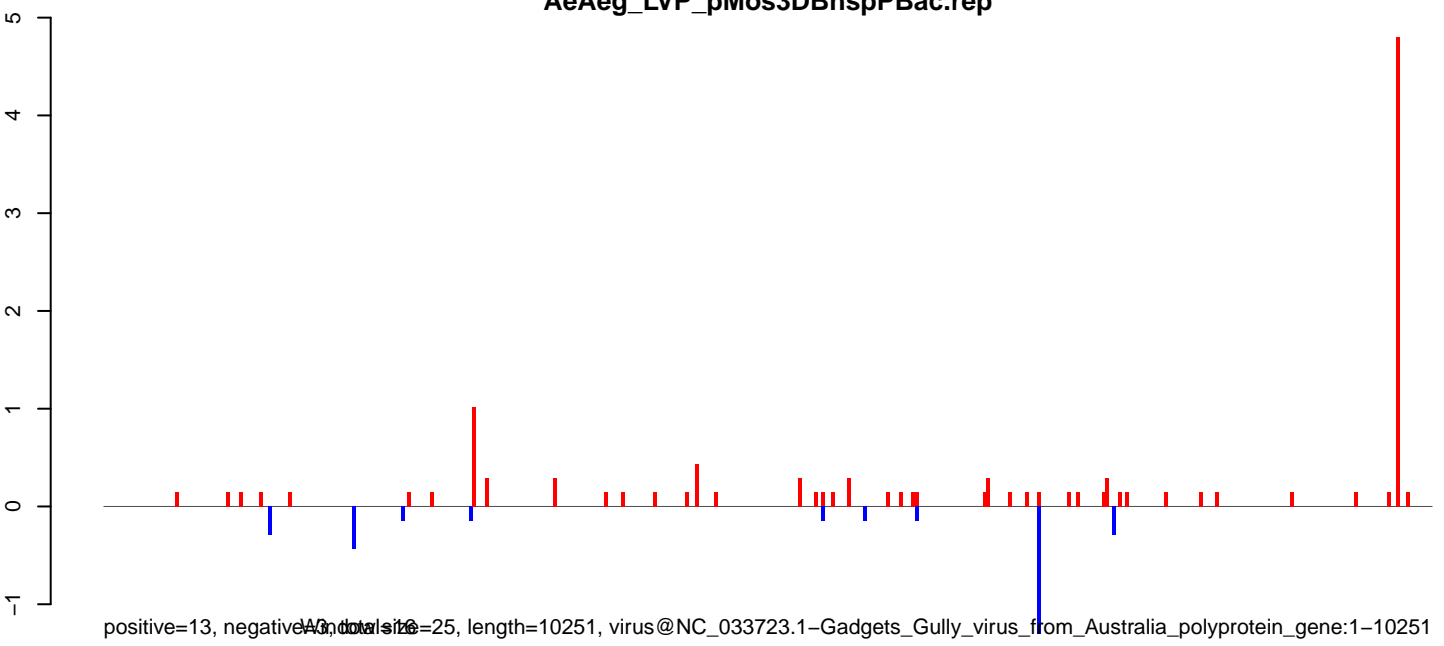
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

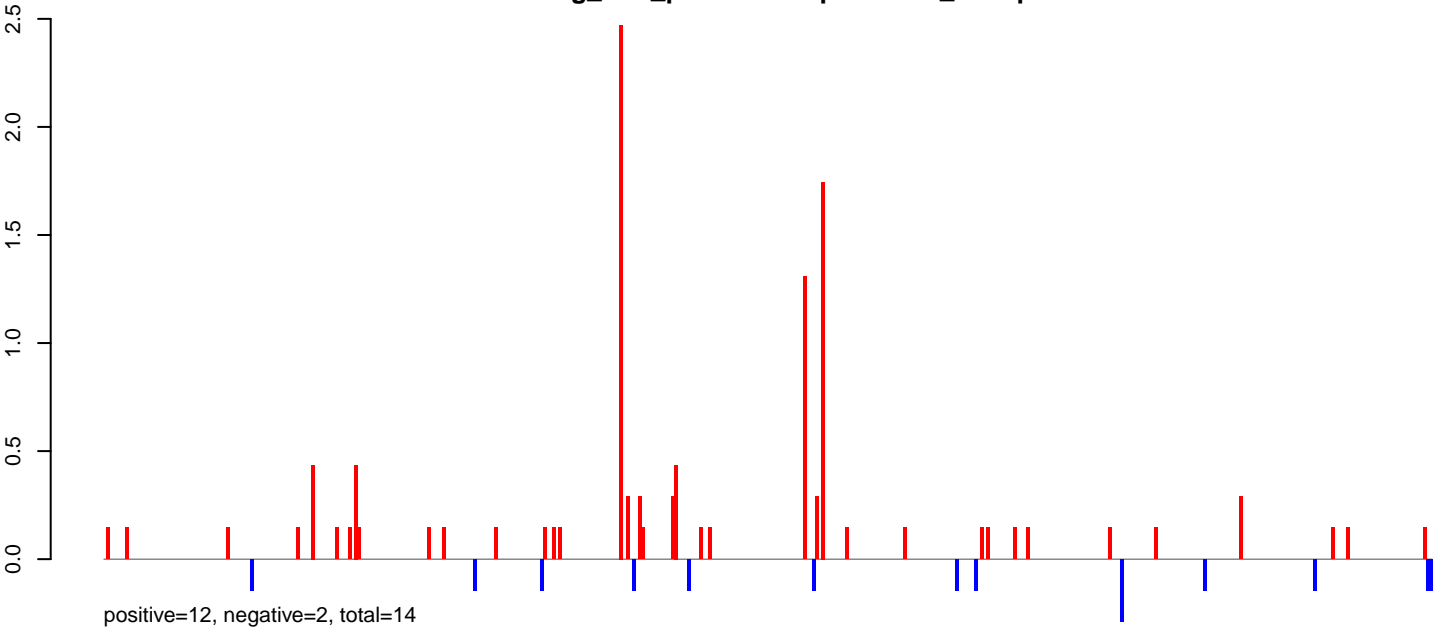


AeAeg_LVP_pMos3DBhspPBac.rep

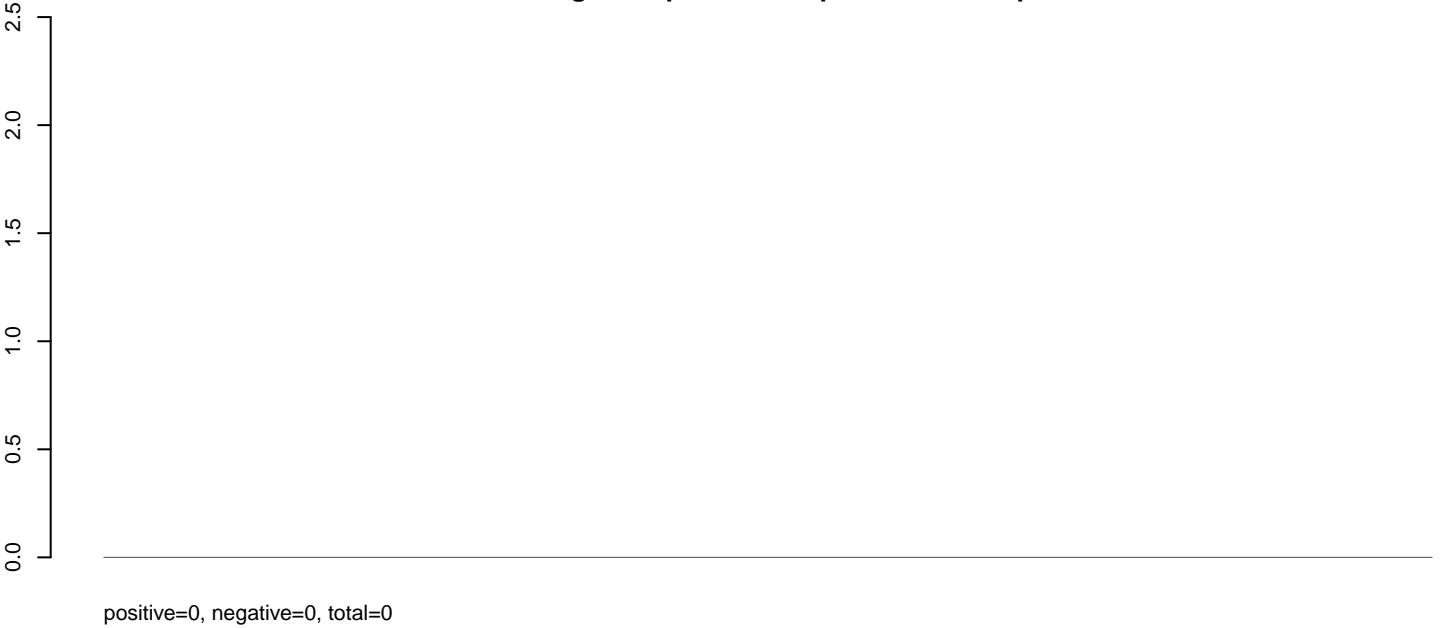


0 2000 4000 6000 8000 10000
positive=13, negative=25, total=38, length=10251, virus@NC_033723.1-Gadgets_Gully_virus_from_Australia_polyprotein_gene:1-10251

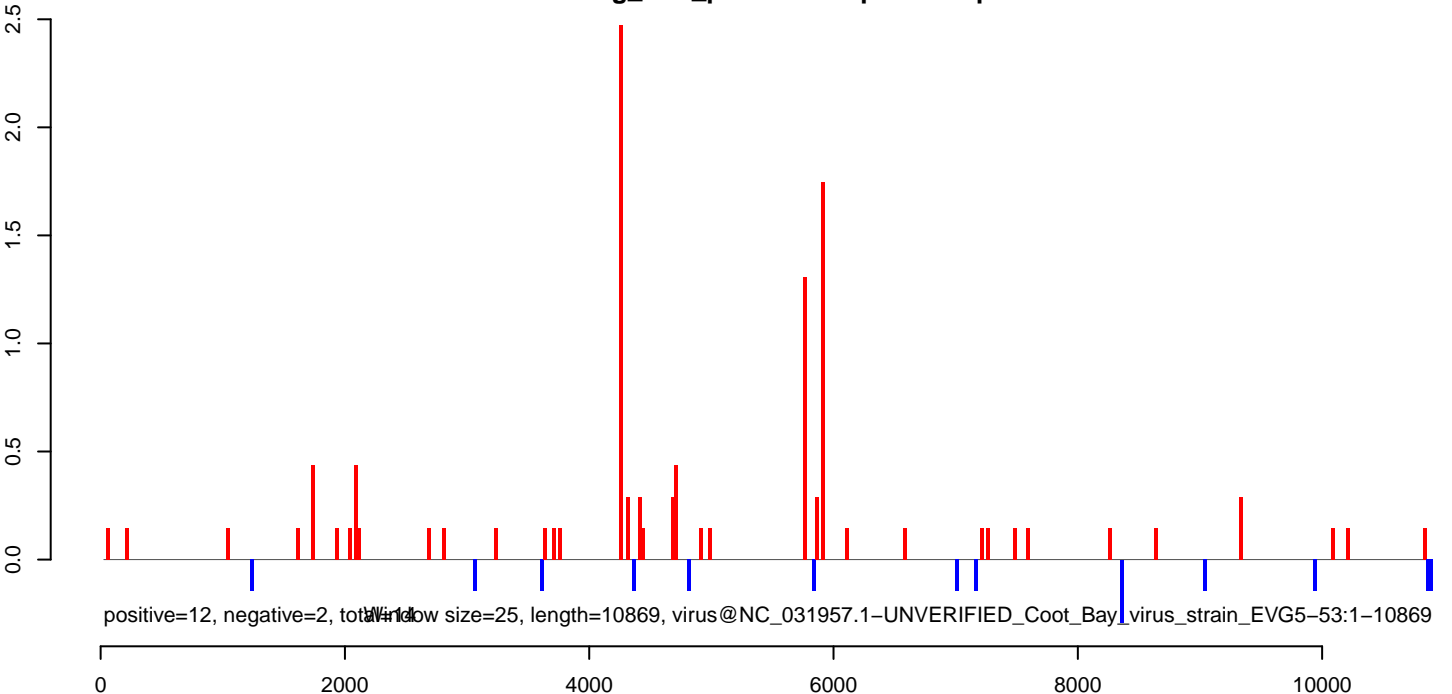
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



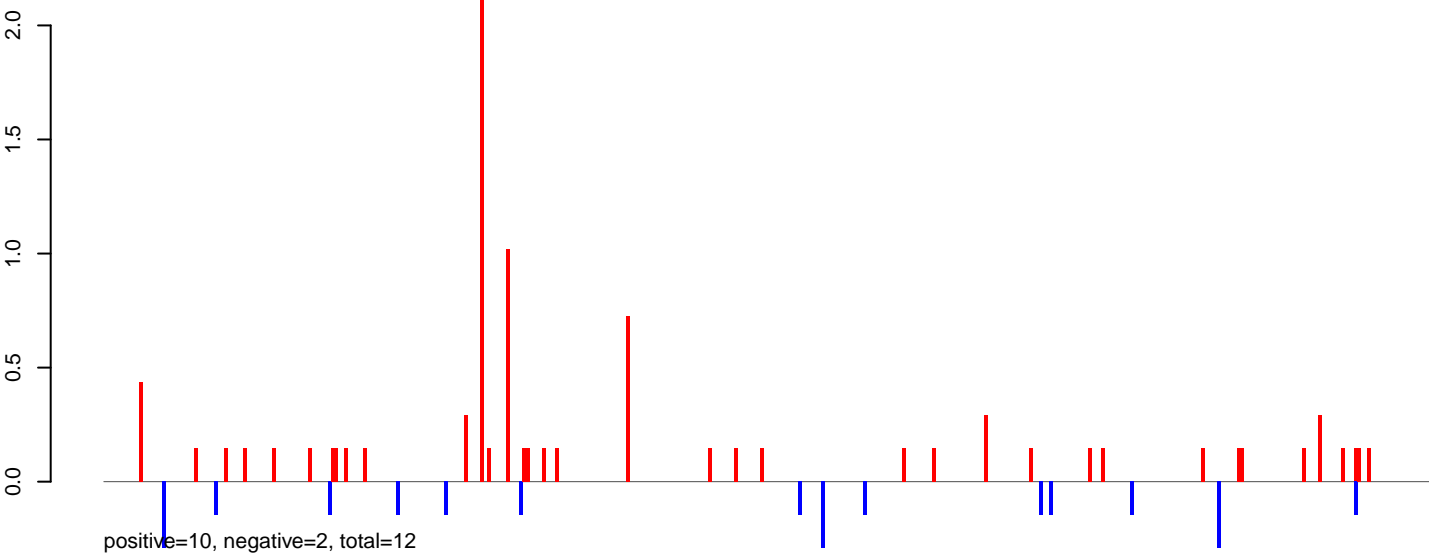
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



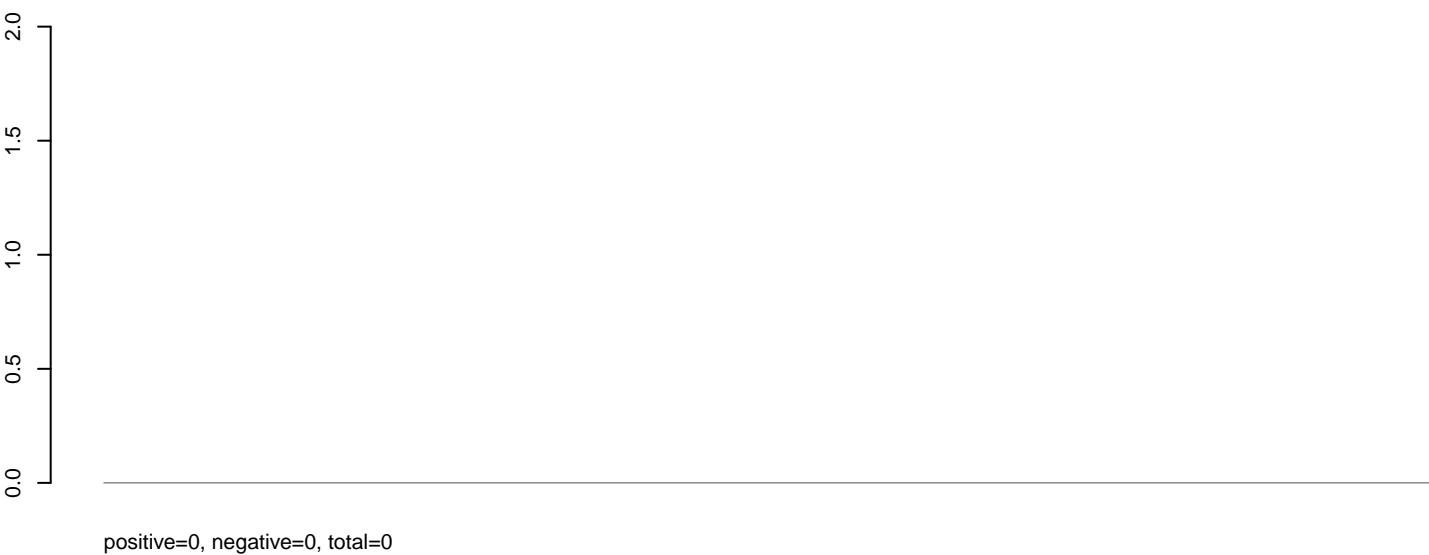
AeAeg_LVP_pMos3DBhspPBac.rep



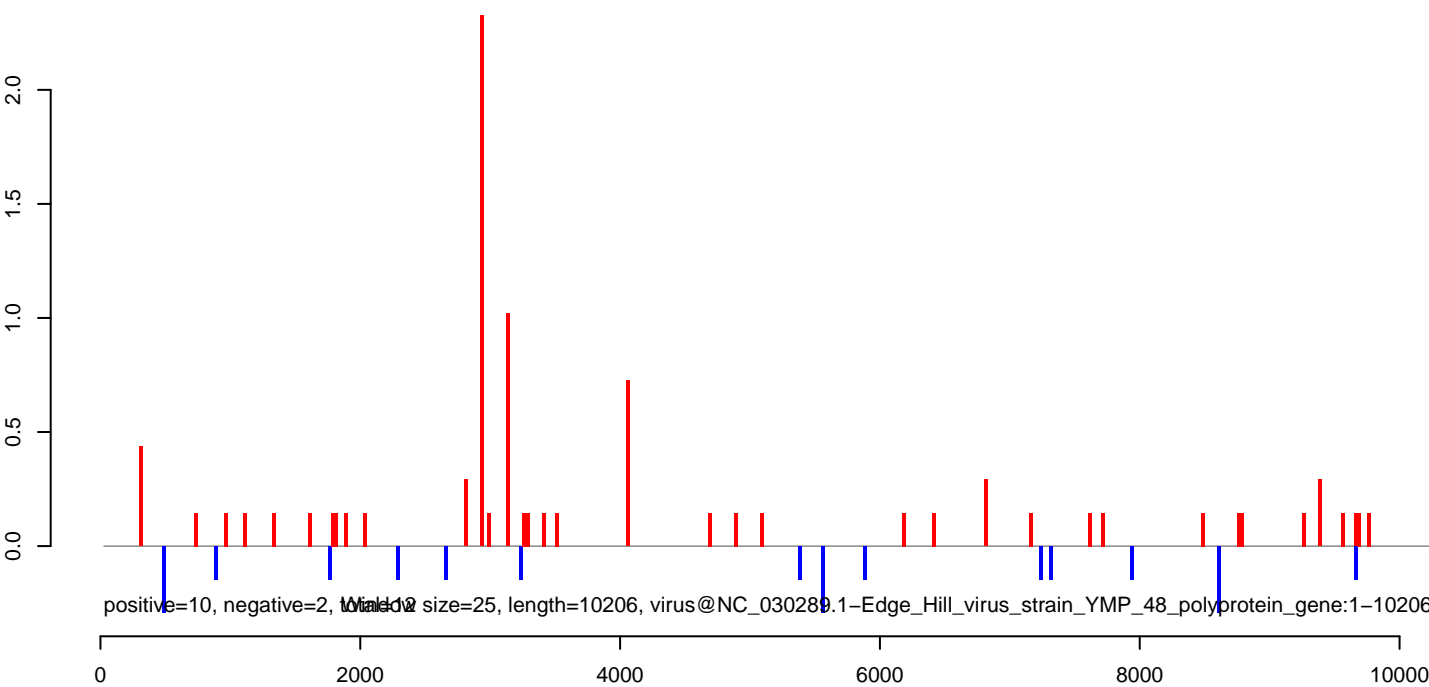
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



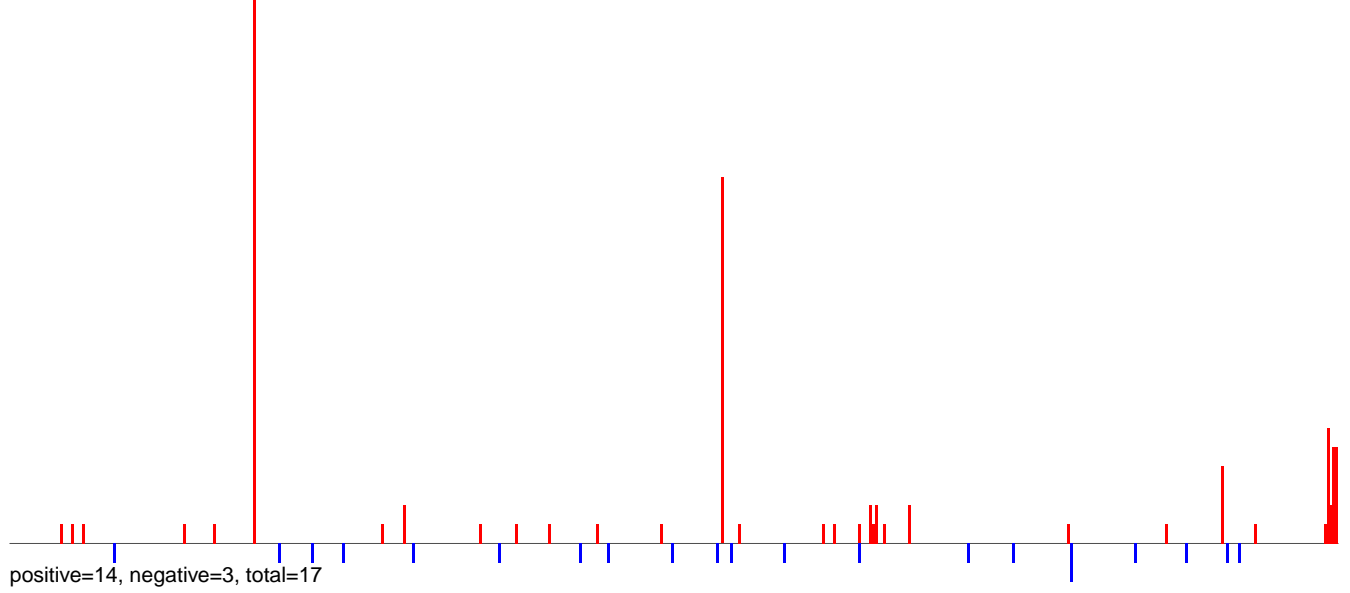
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



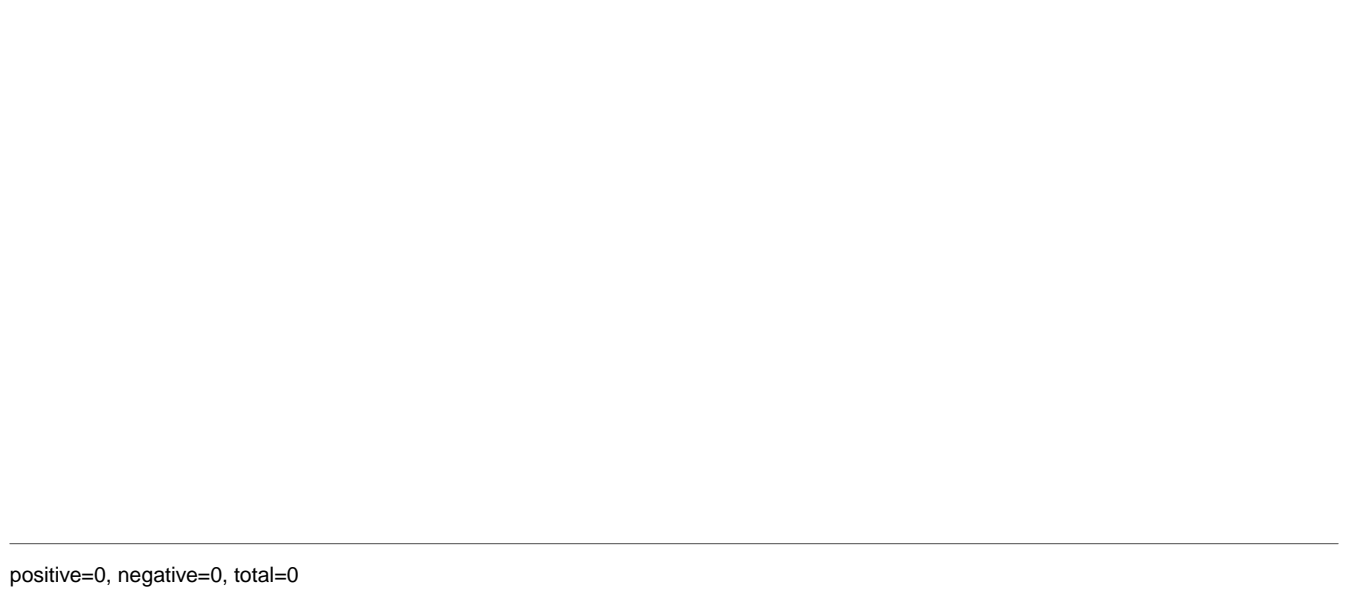
AeAeg_LVP_pMos3DBhspPBac.rep



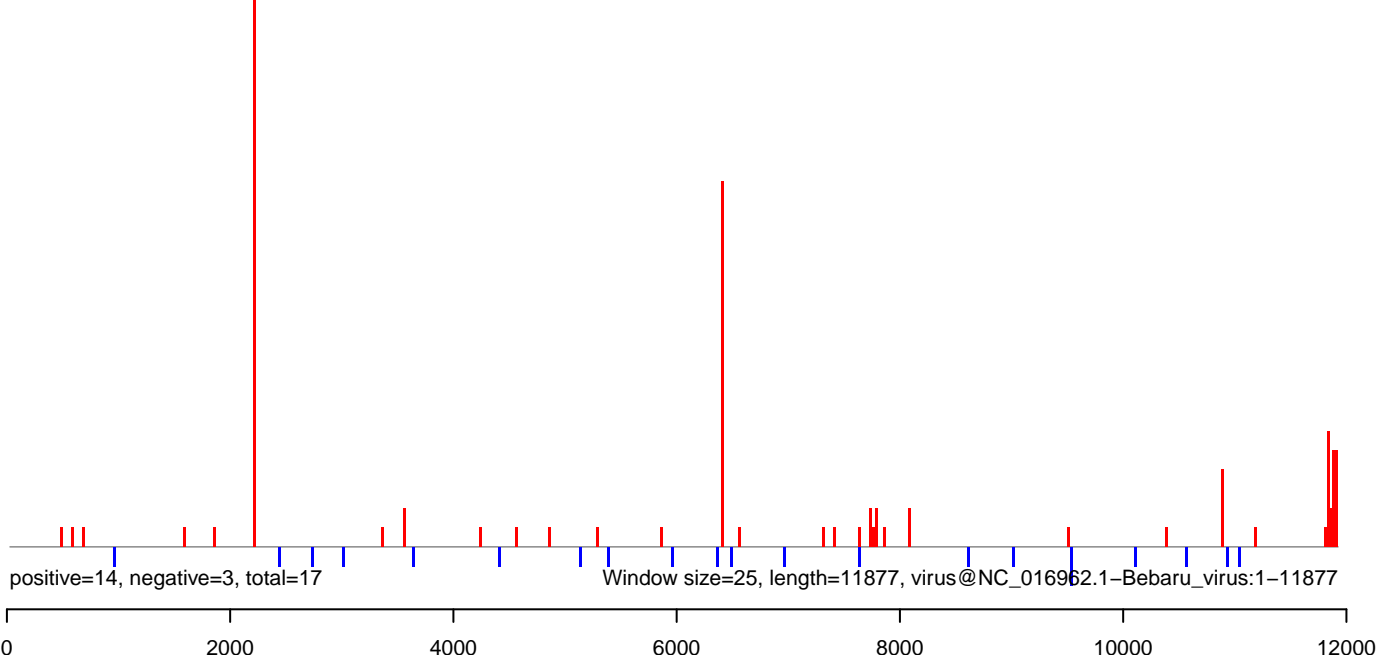
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



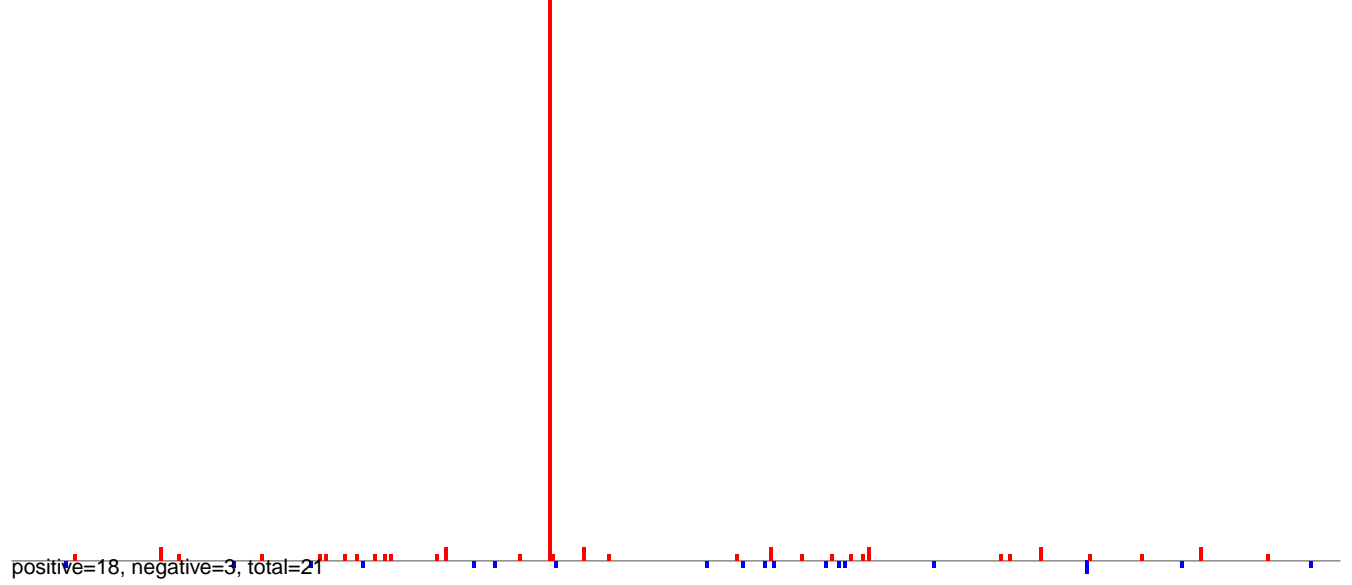
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



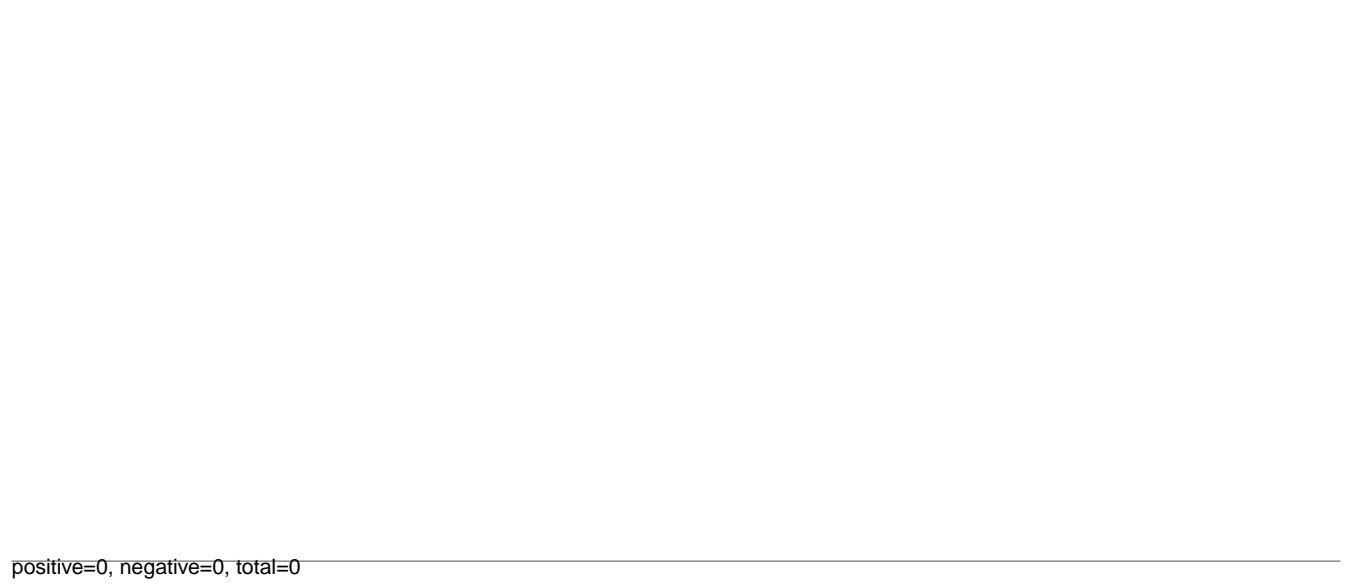
AeAeg_LVP_pMos3DBhspPBac.rep



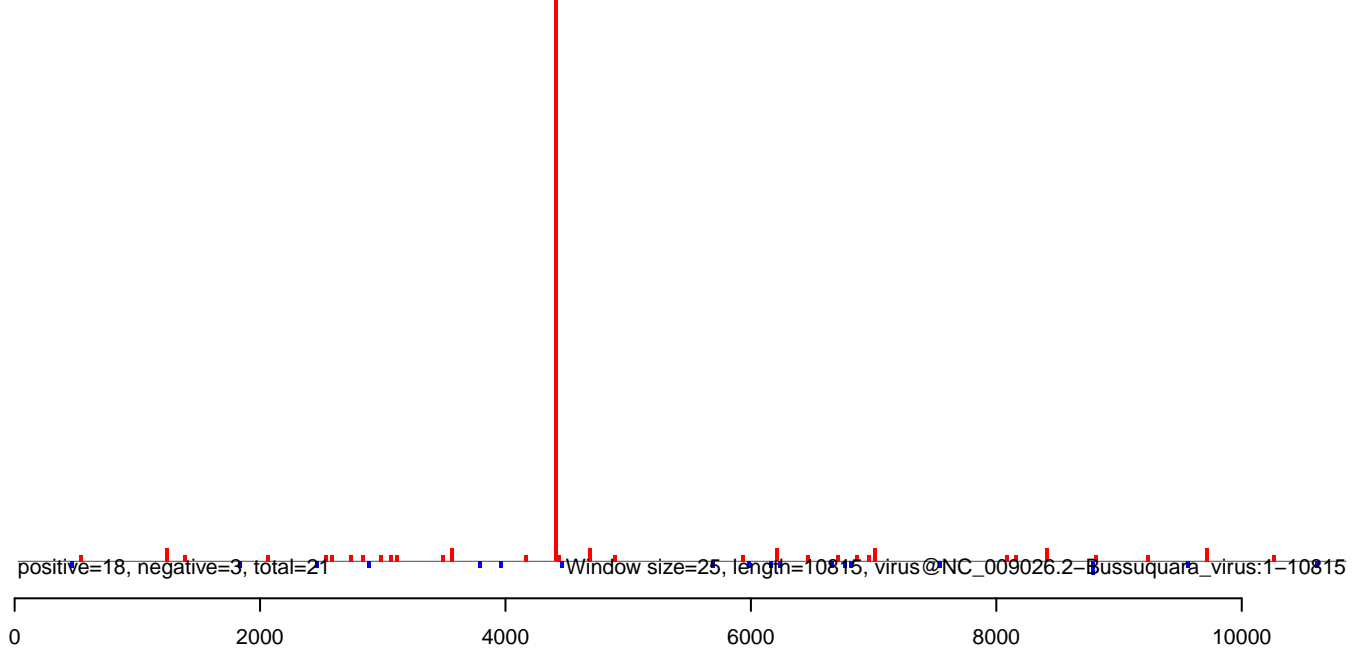
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



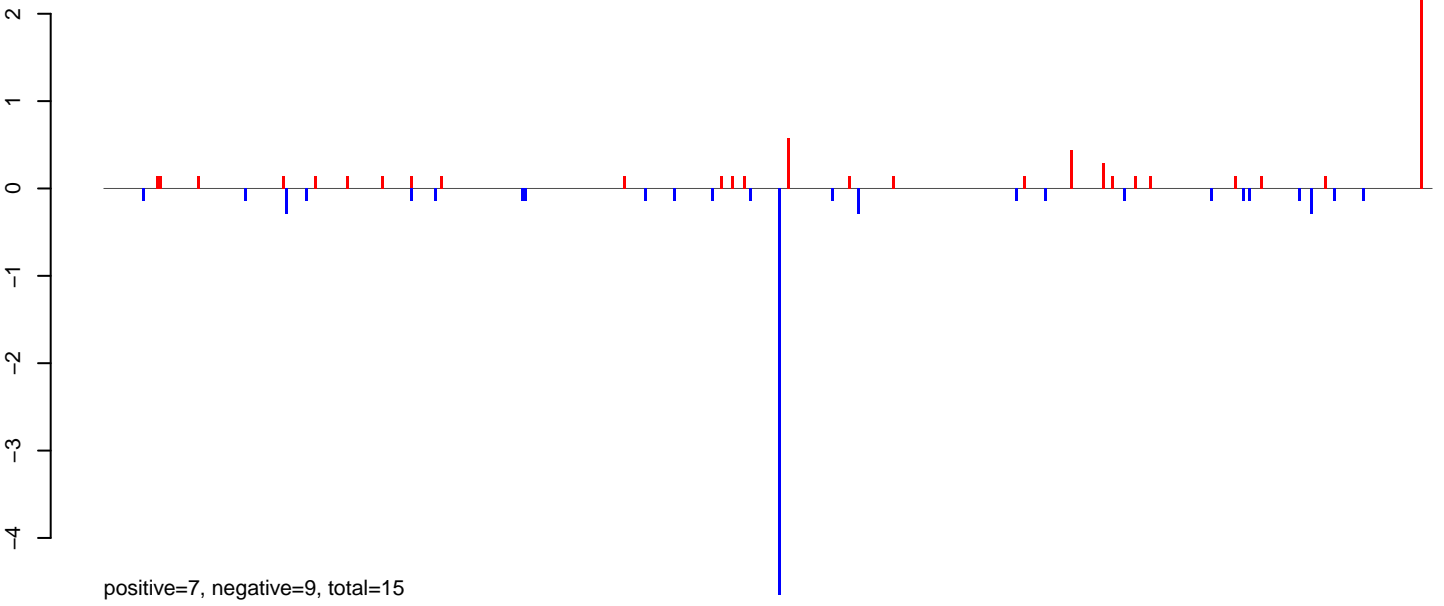
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



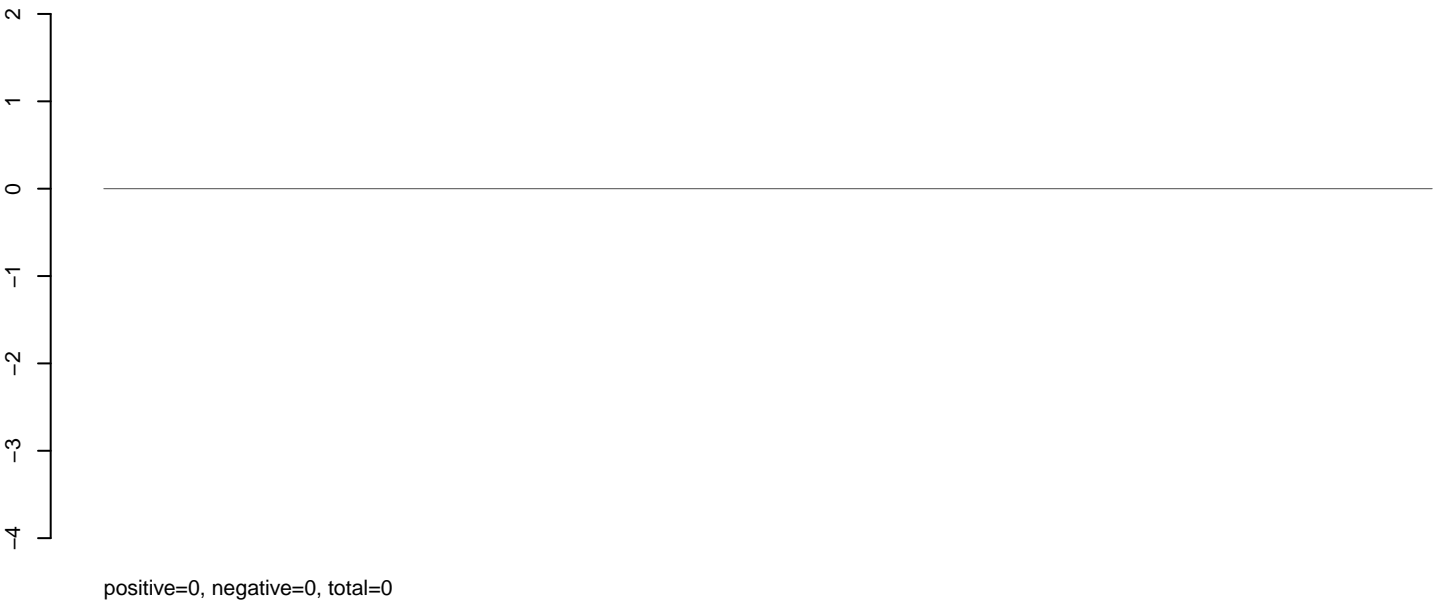
AeAeg_LVP_pMos3DBhspPBac.rep



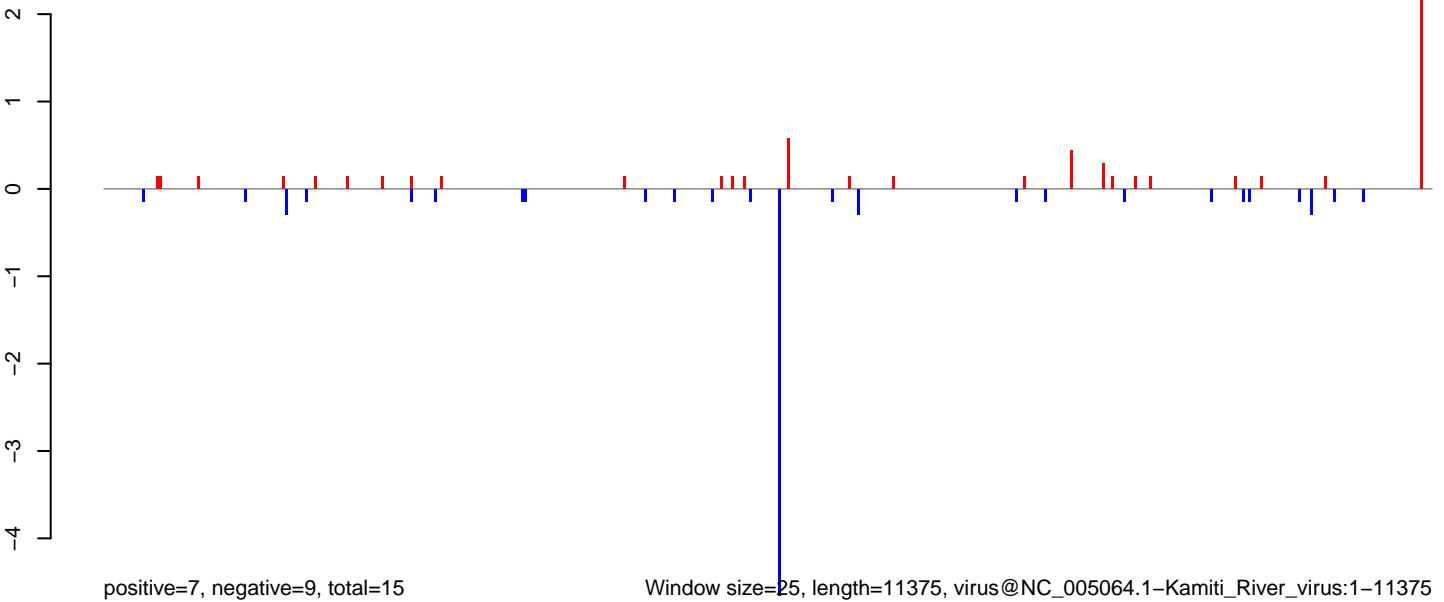
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

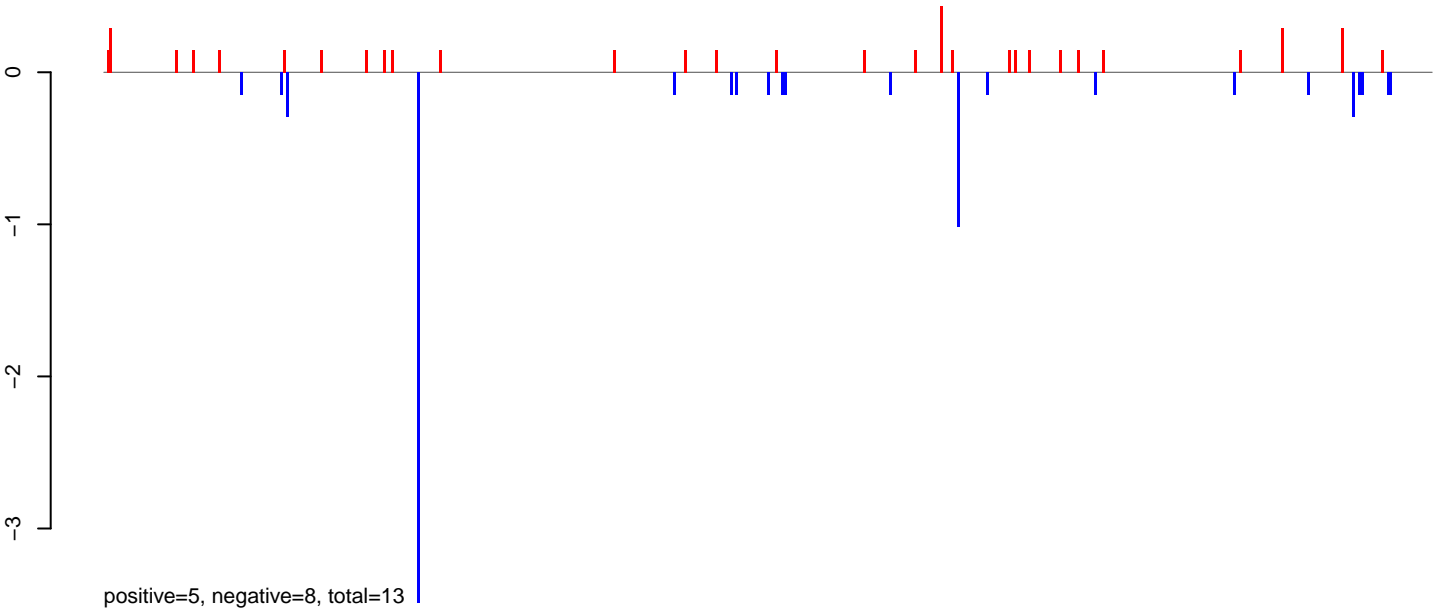


AeAeg_LVP_pMos3DBhspPBac.rep

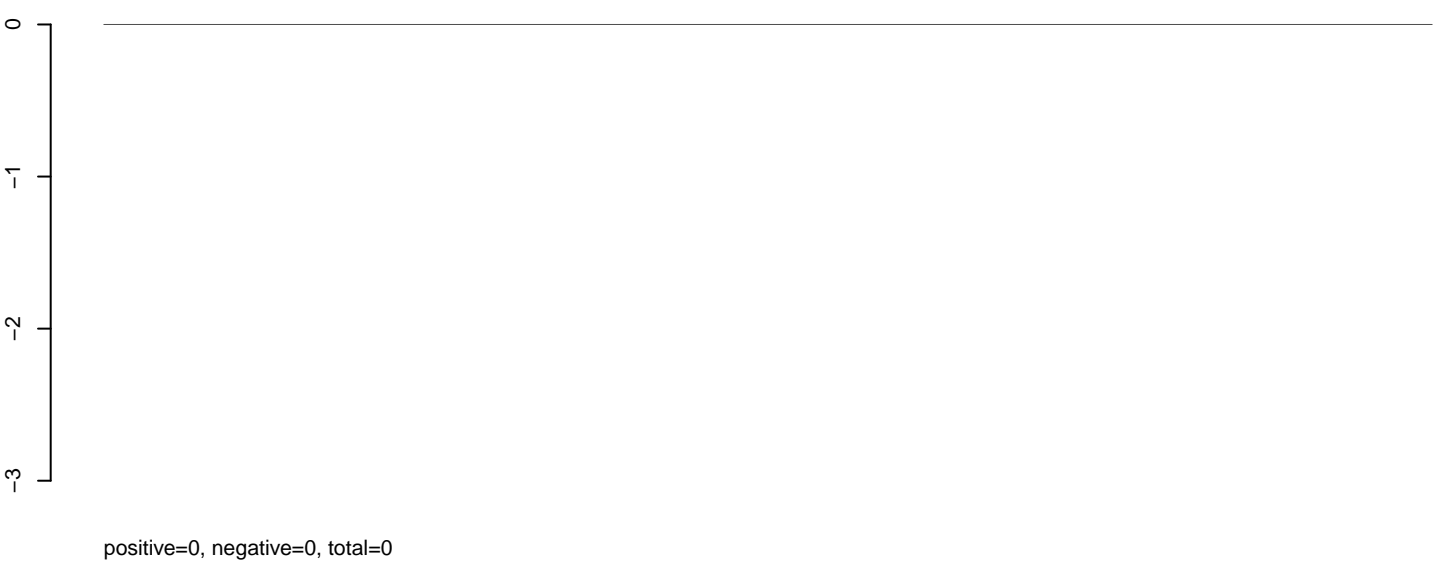


0 2000 4000 6000 8000 10000

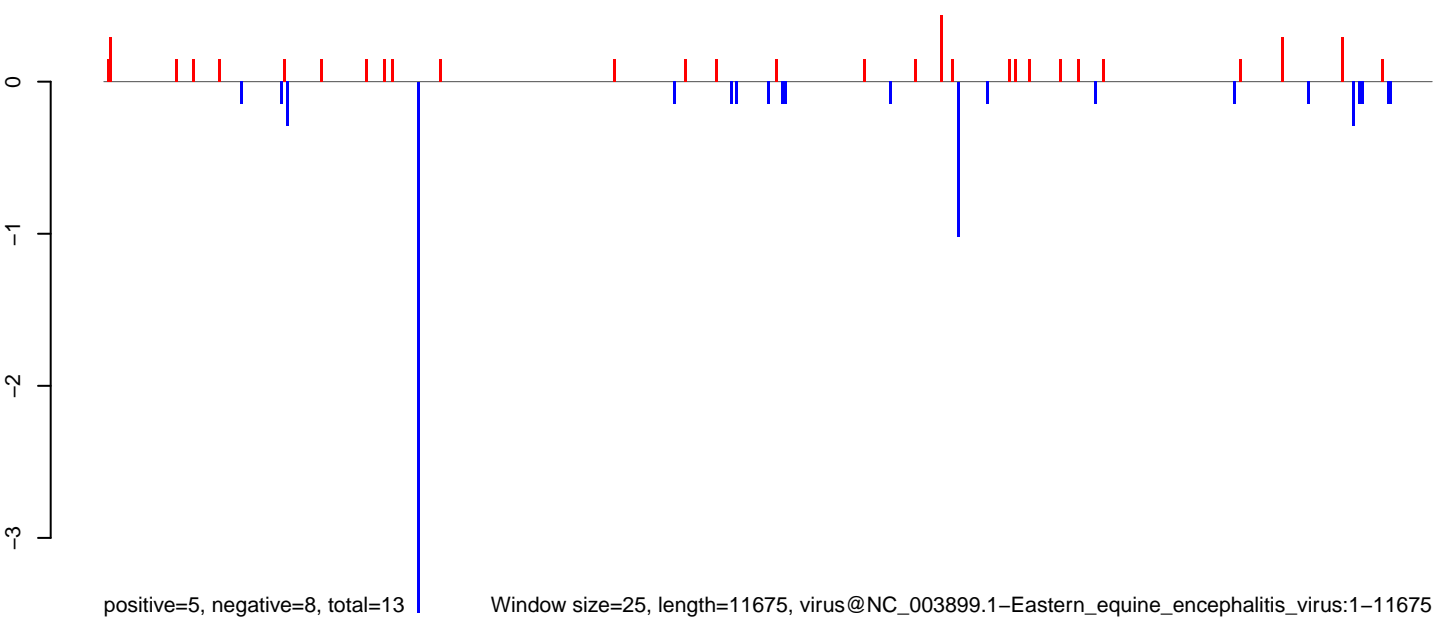
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



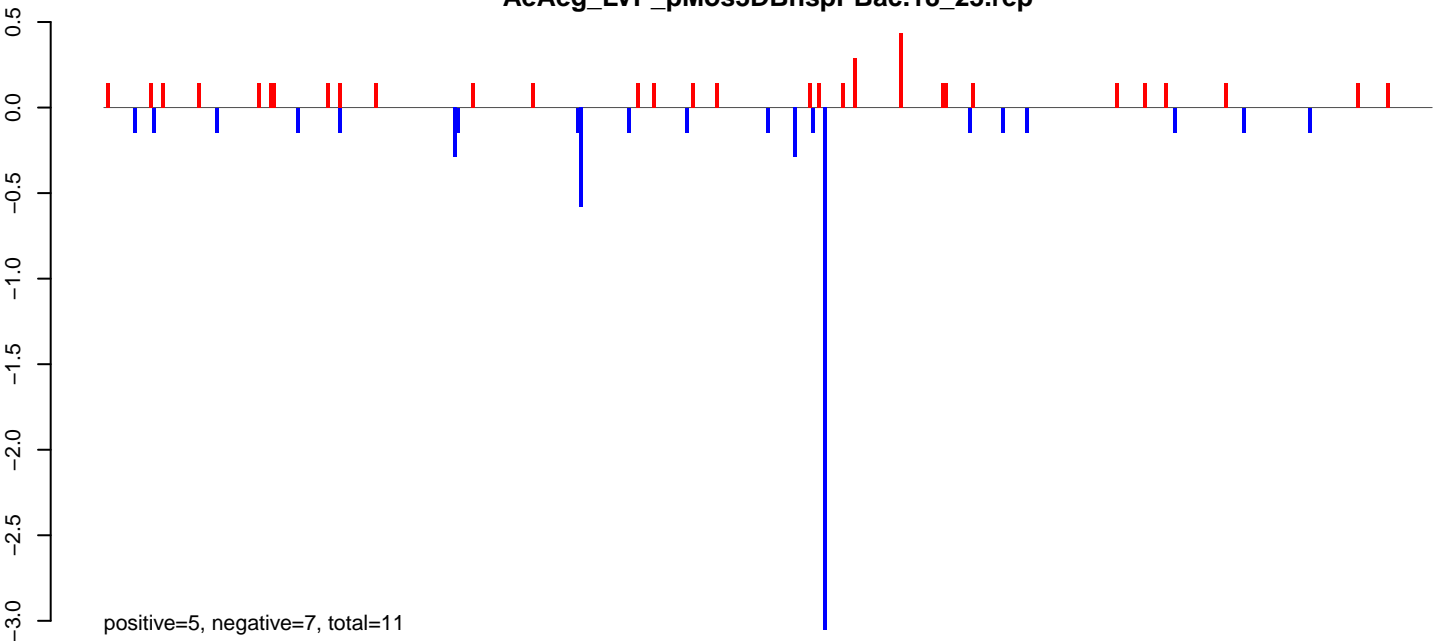
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



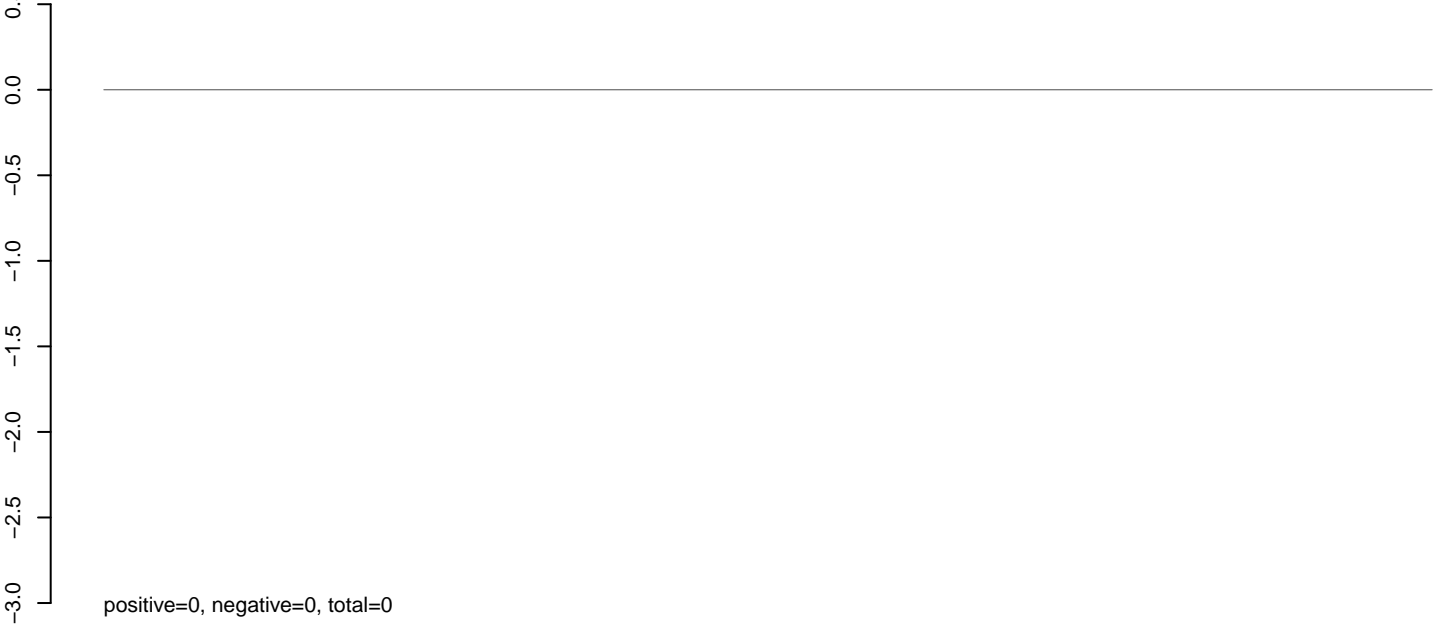
AeAeg_LVP_pMos3DBhspPBac.rep



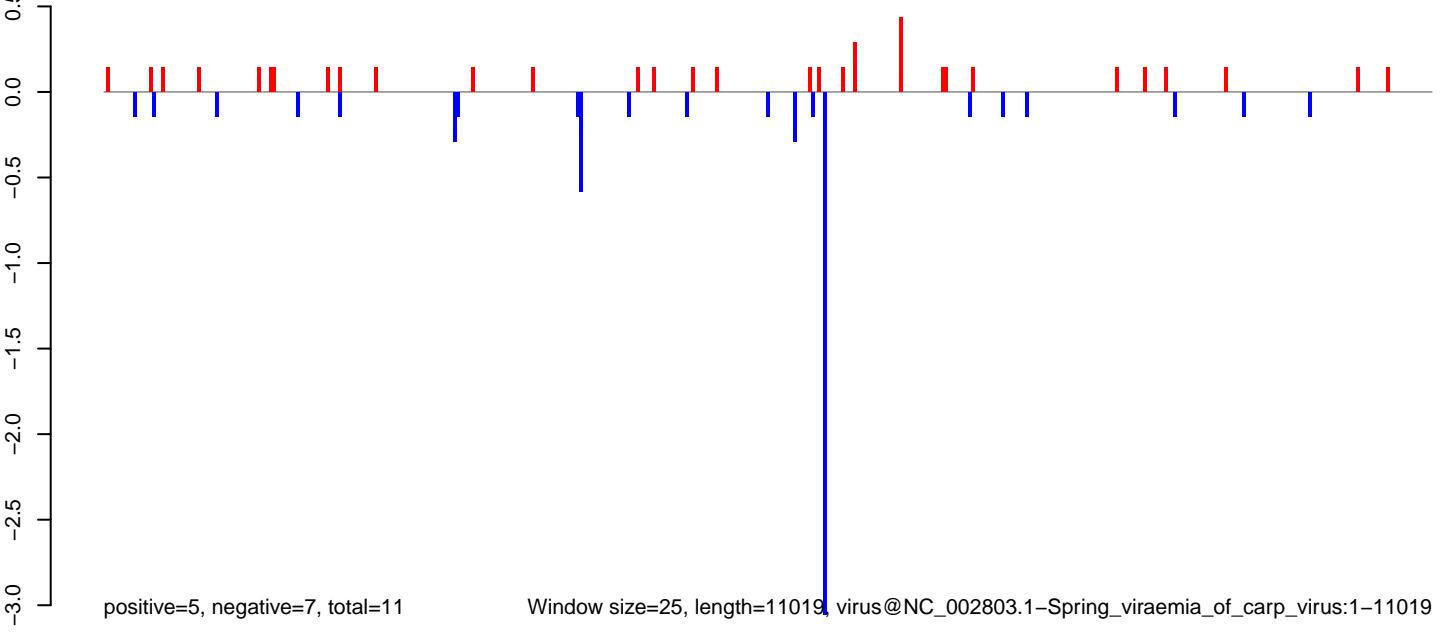
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



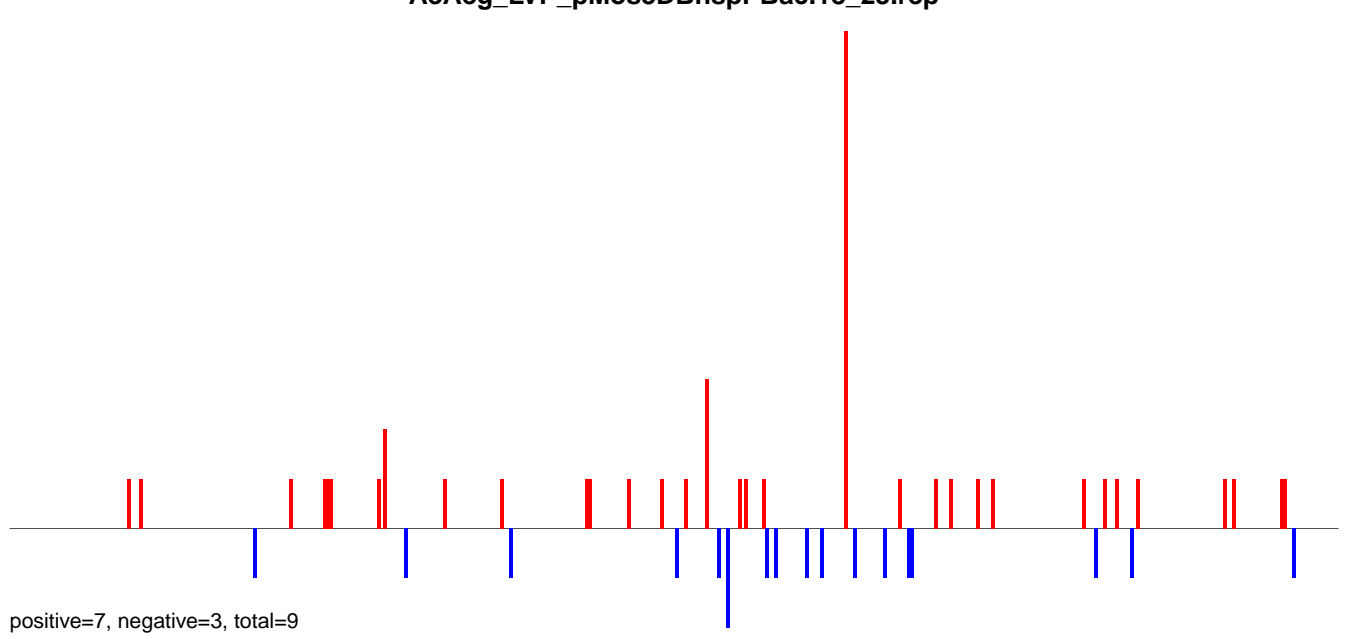
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



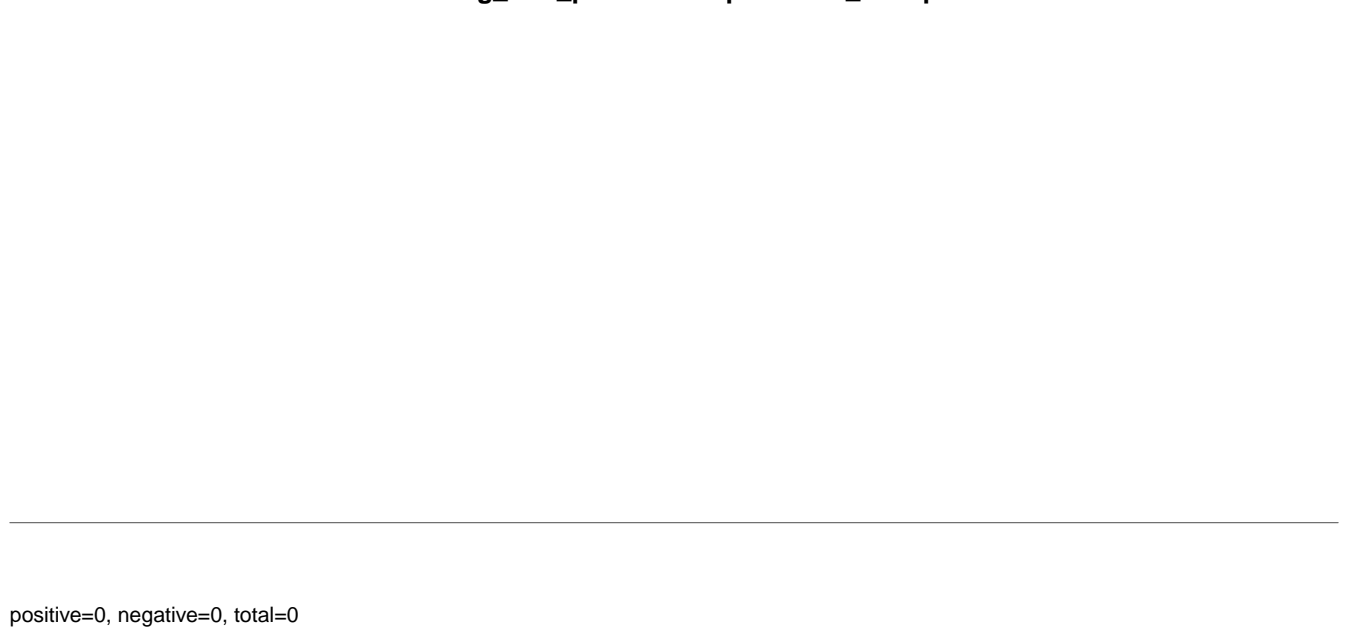
AeAeg_LVP_pMos3DBhspPBac.rep



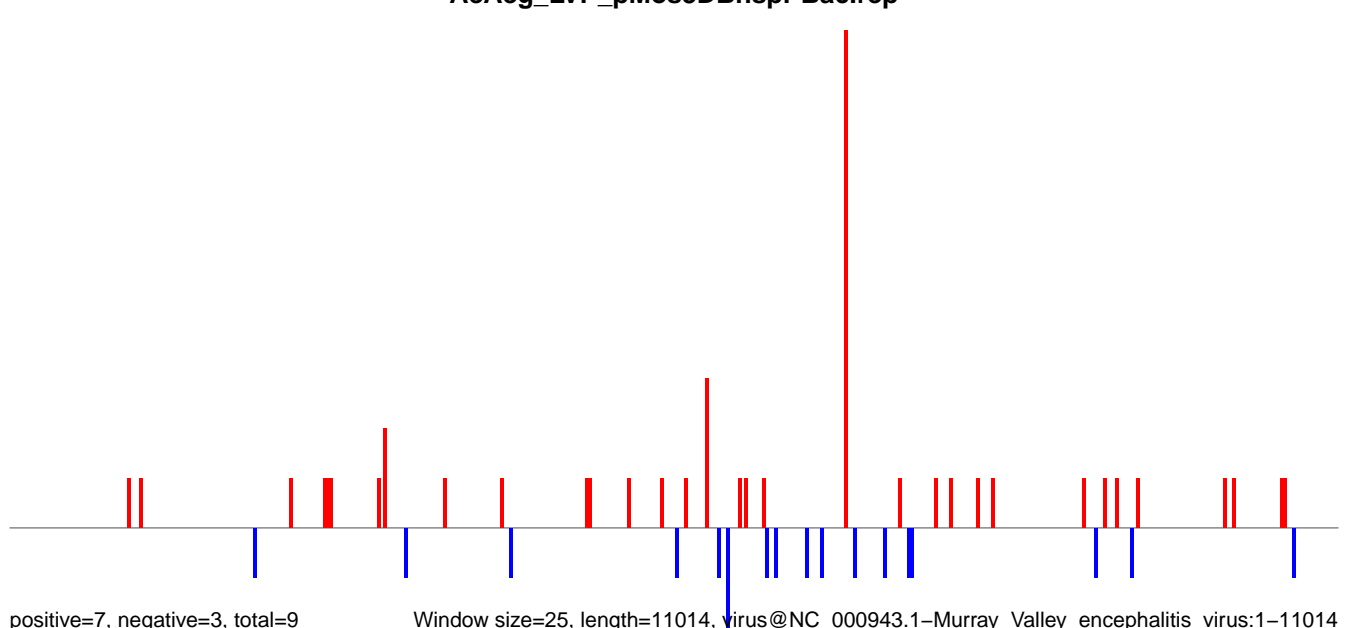
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



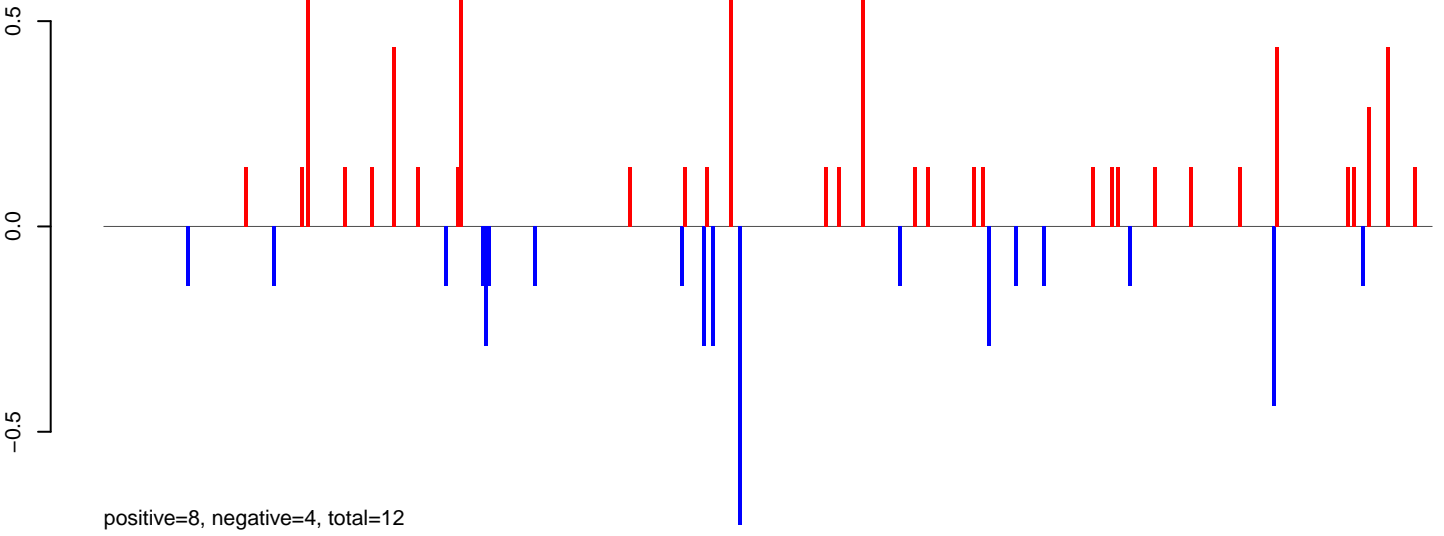
AeAeg_LVP_pMos3DBhspPBac.rep



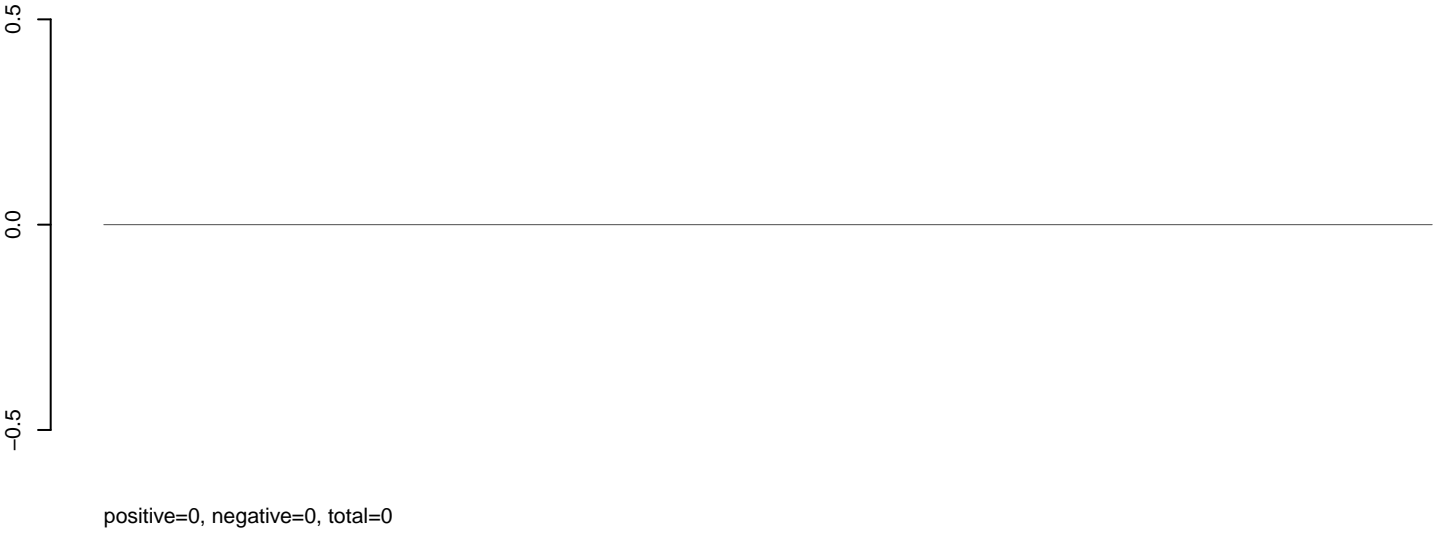
Window size=25, length=11014, virus@NC_000943.1-Murray_Valley_encephalitis_virus:1-11014

0 2000 4000 6000 8000 10000

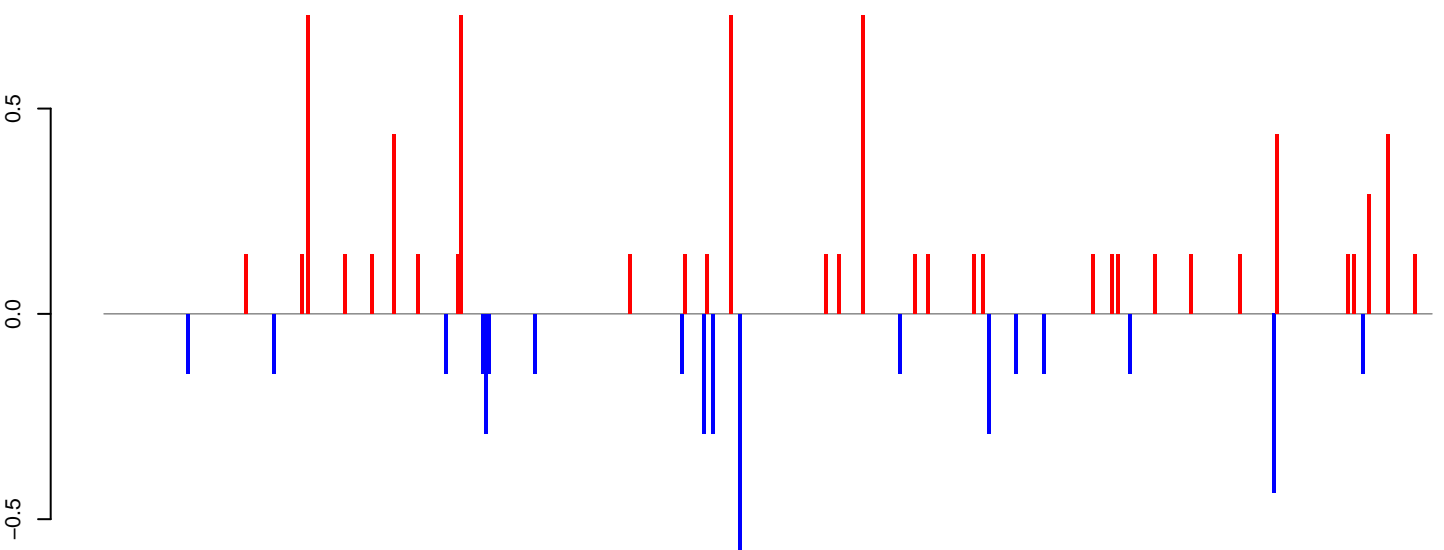
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



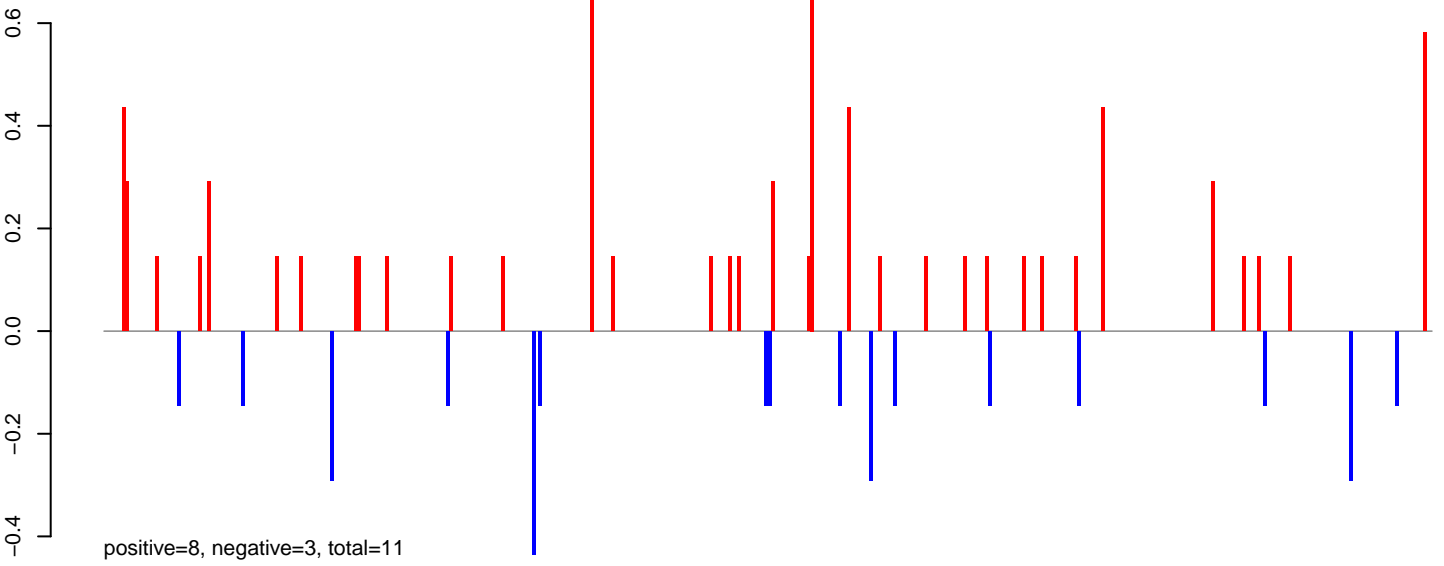
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



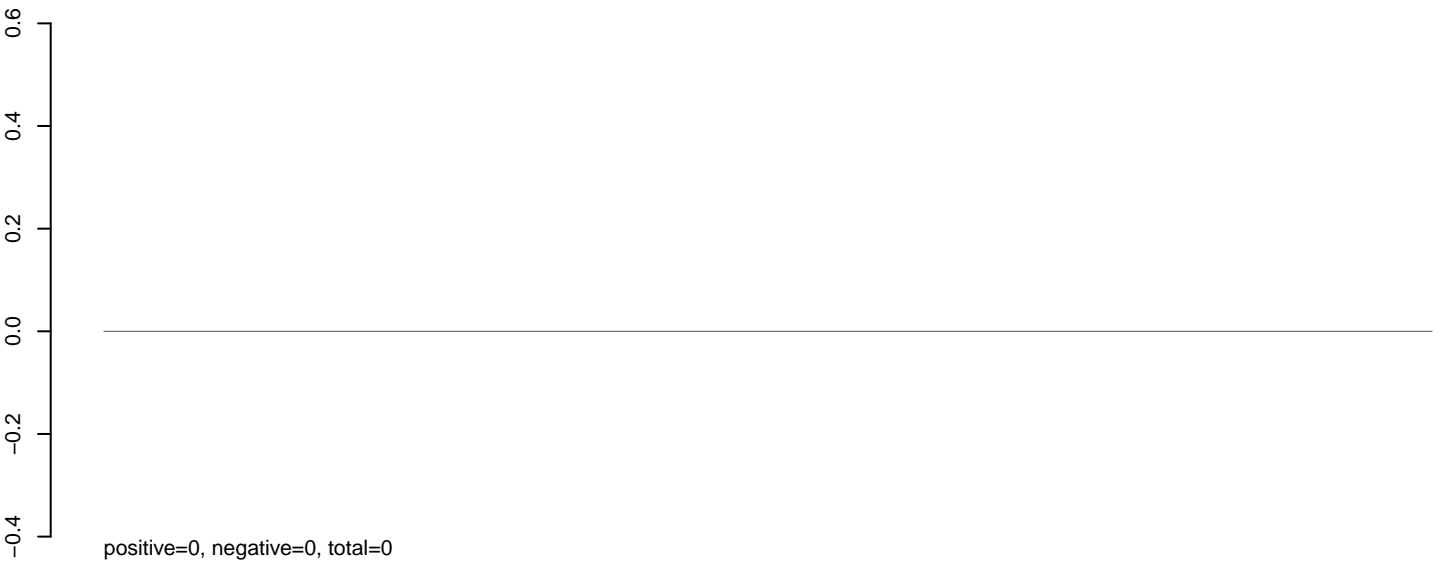
AeAeg_LVP_pMos3DBhspPBac.rep



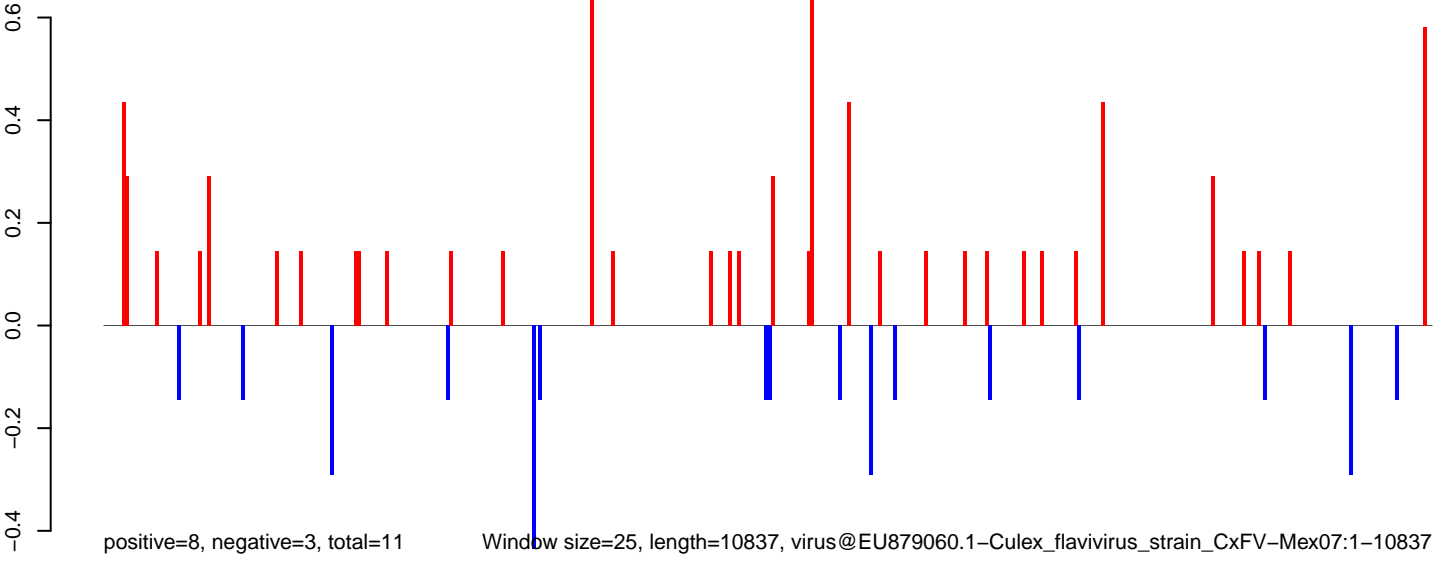
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



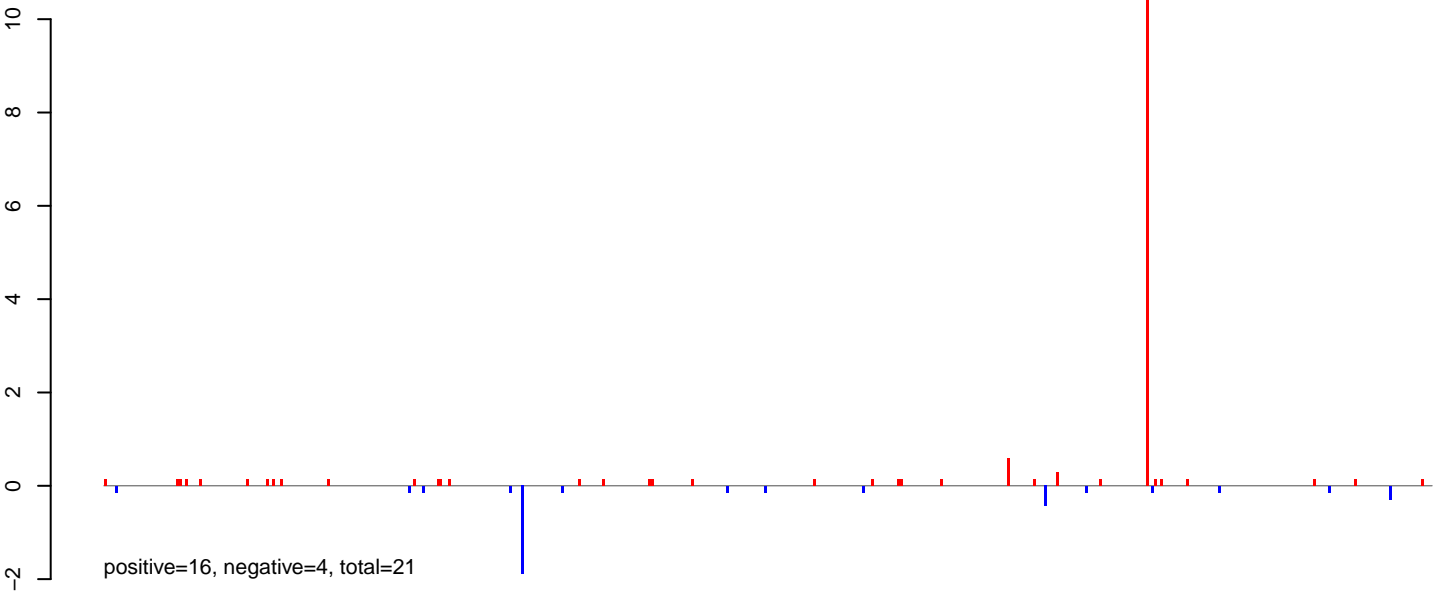
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



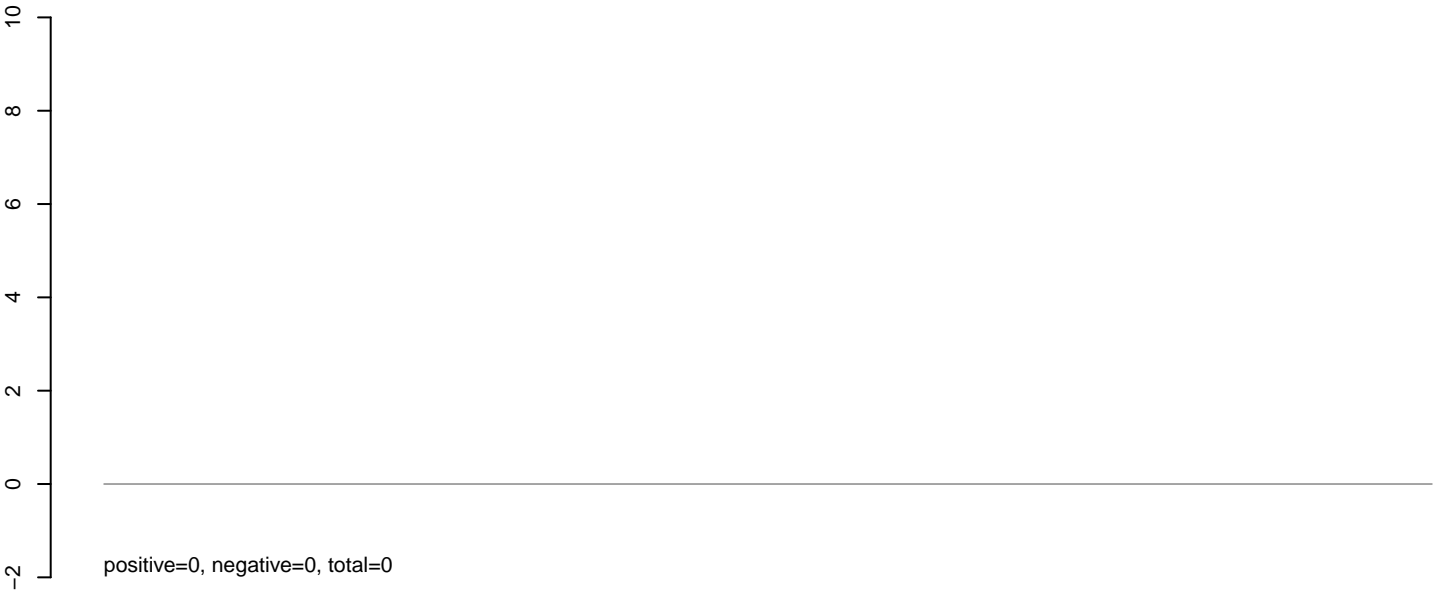
AeAeg_LVP_pMos3DBhspPBac.rep



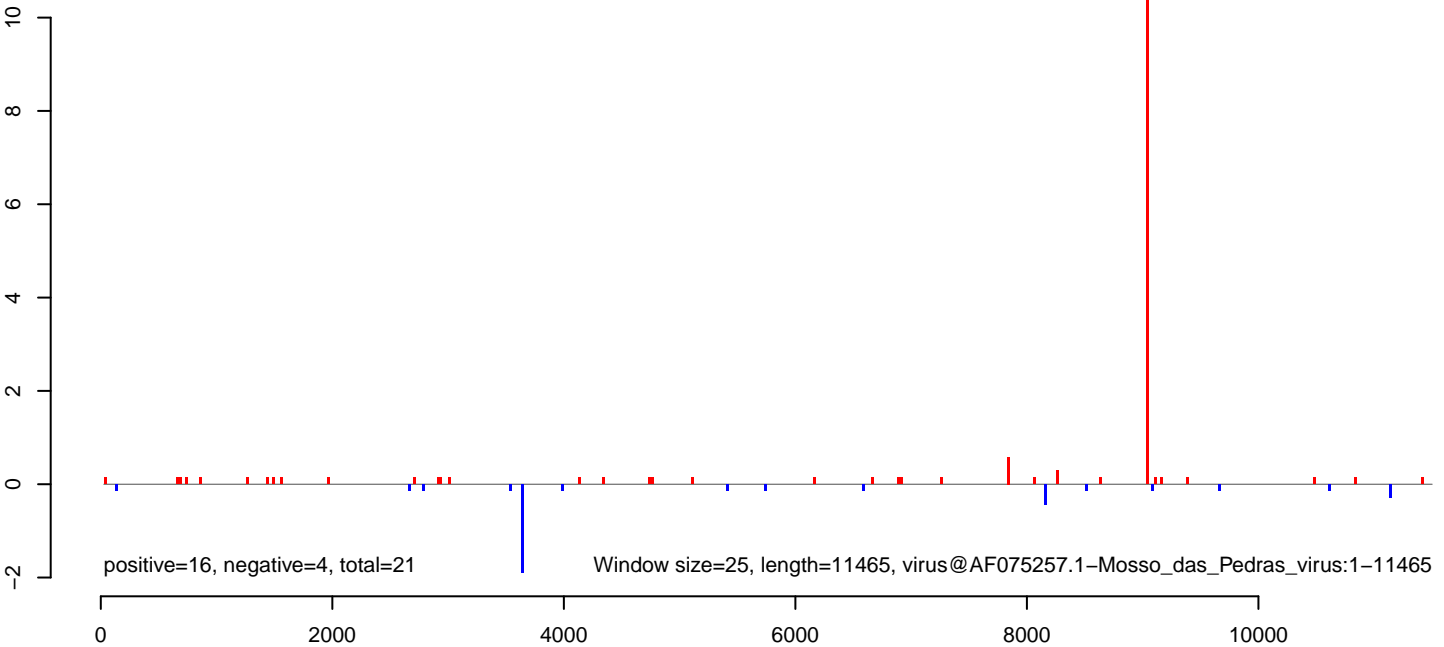
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



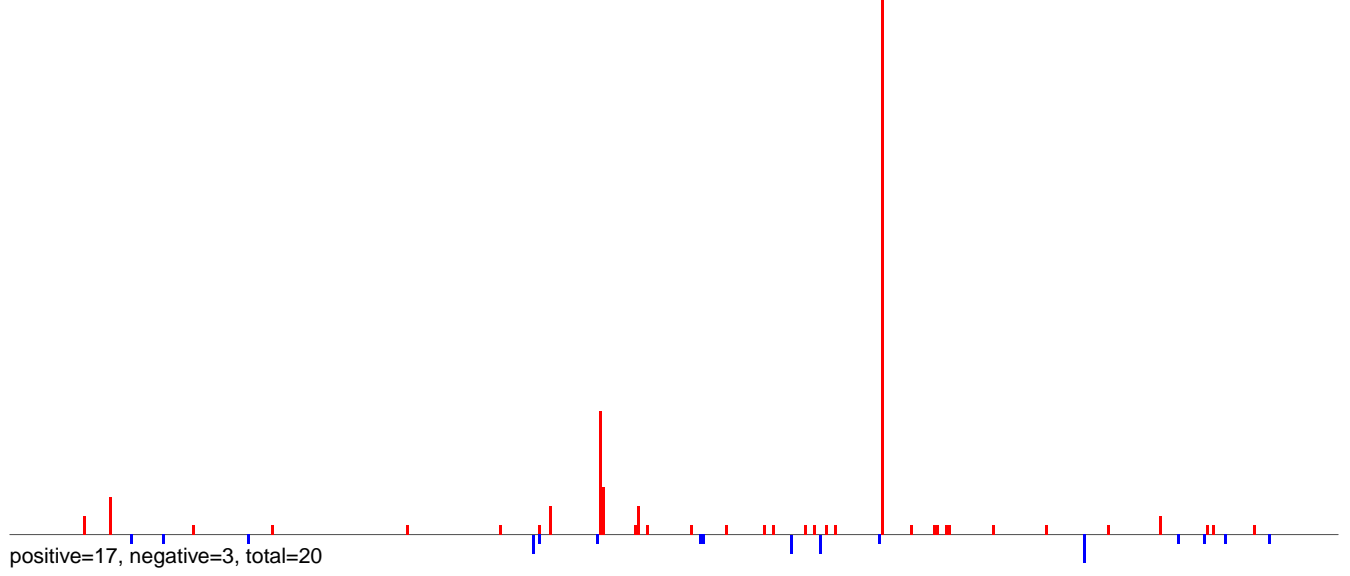
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



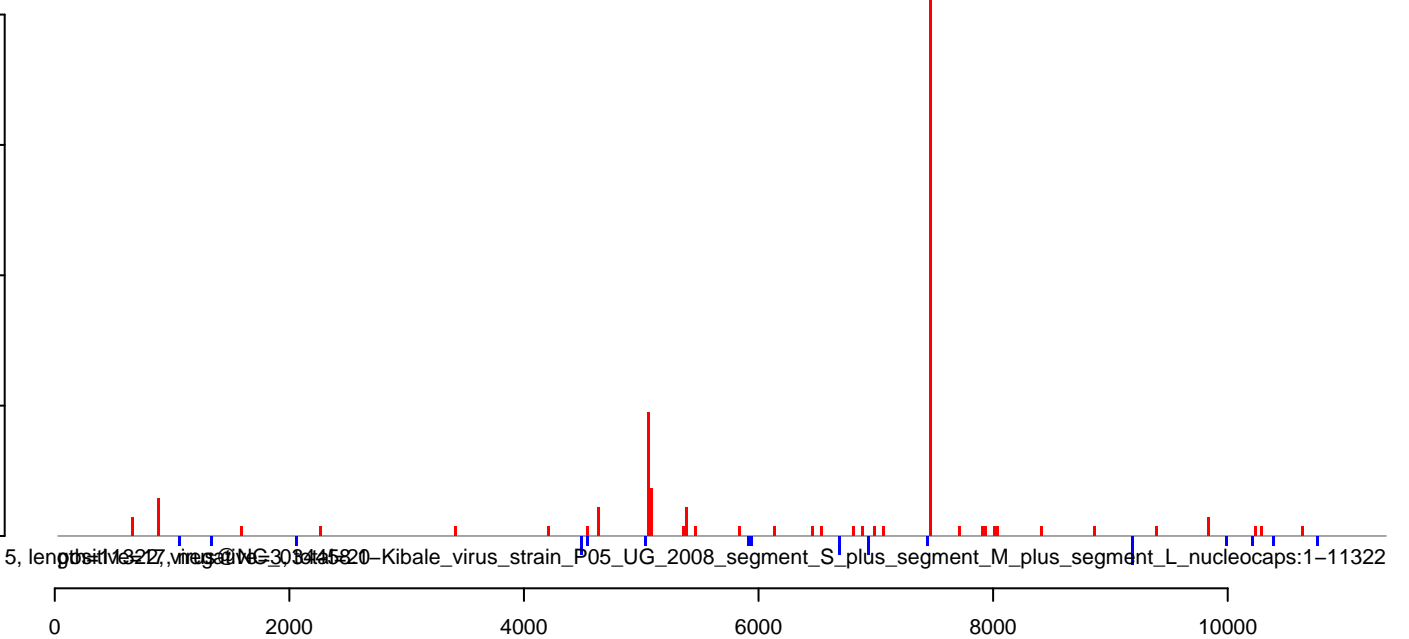
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



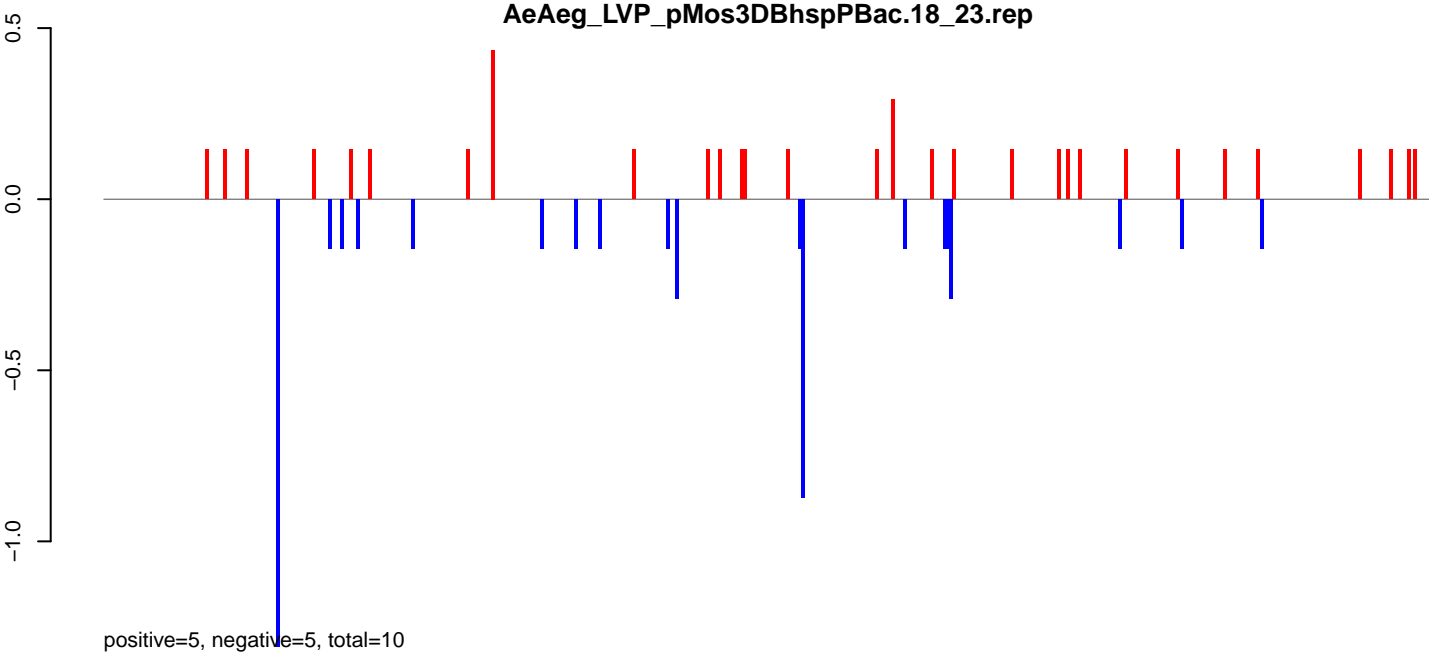
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



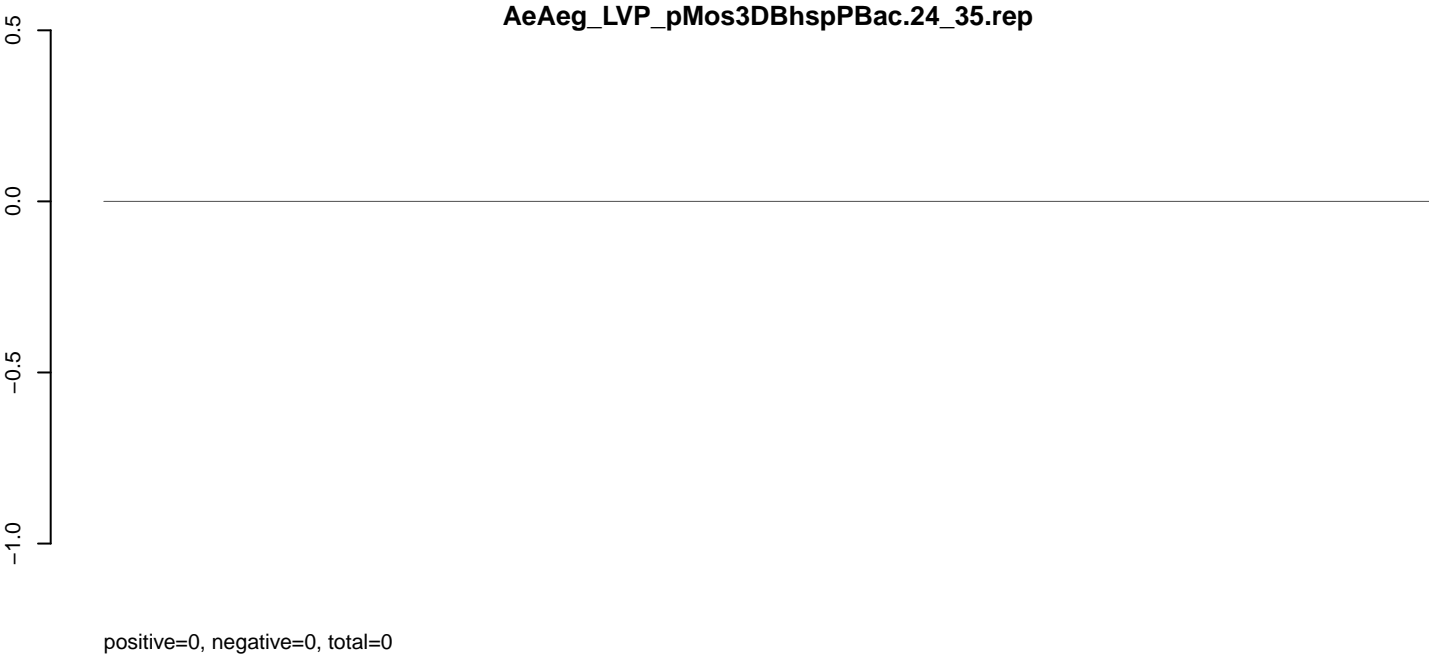
AeAeg_LVP_pMos3DBhspPBac.rep



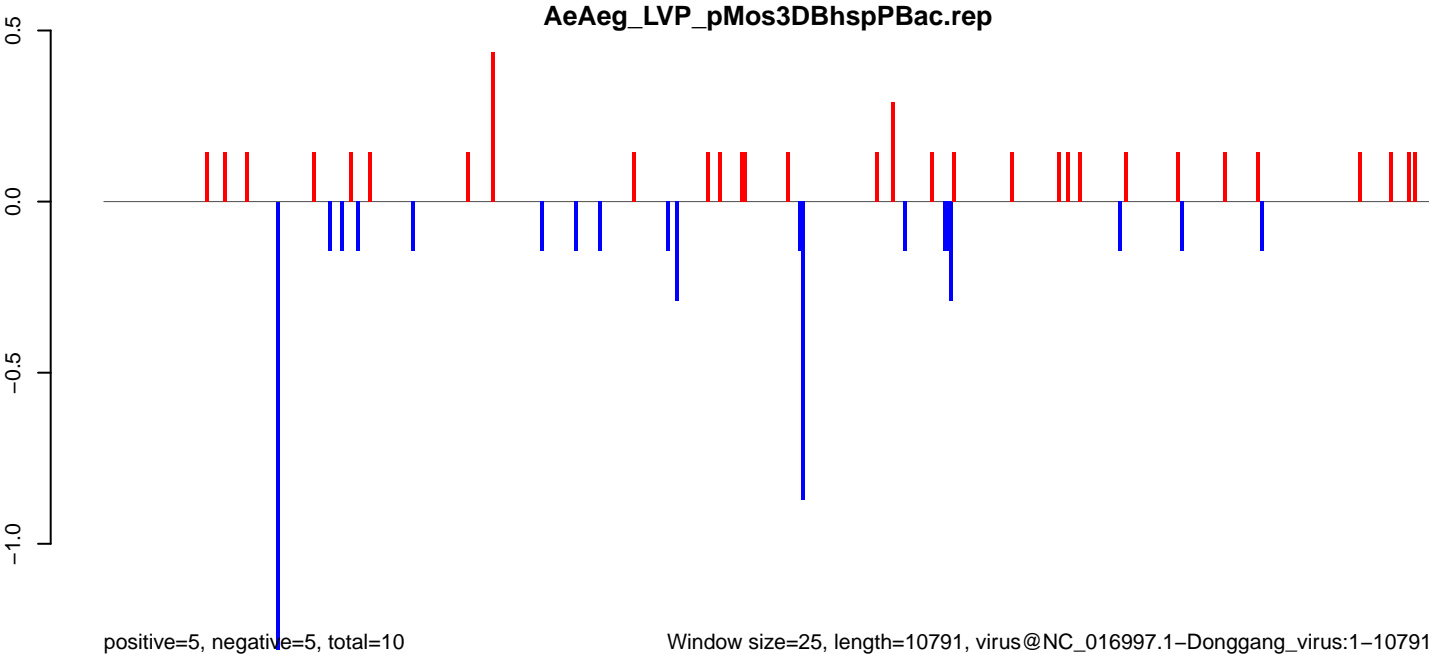
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



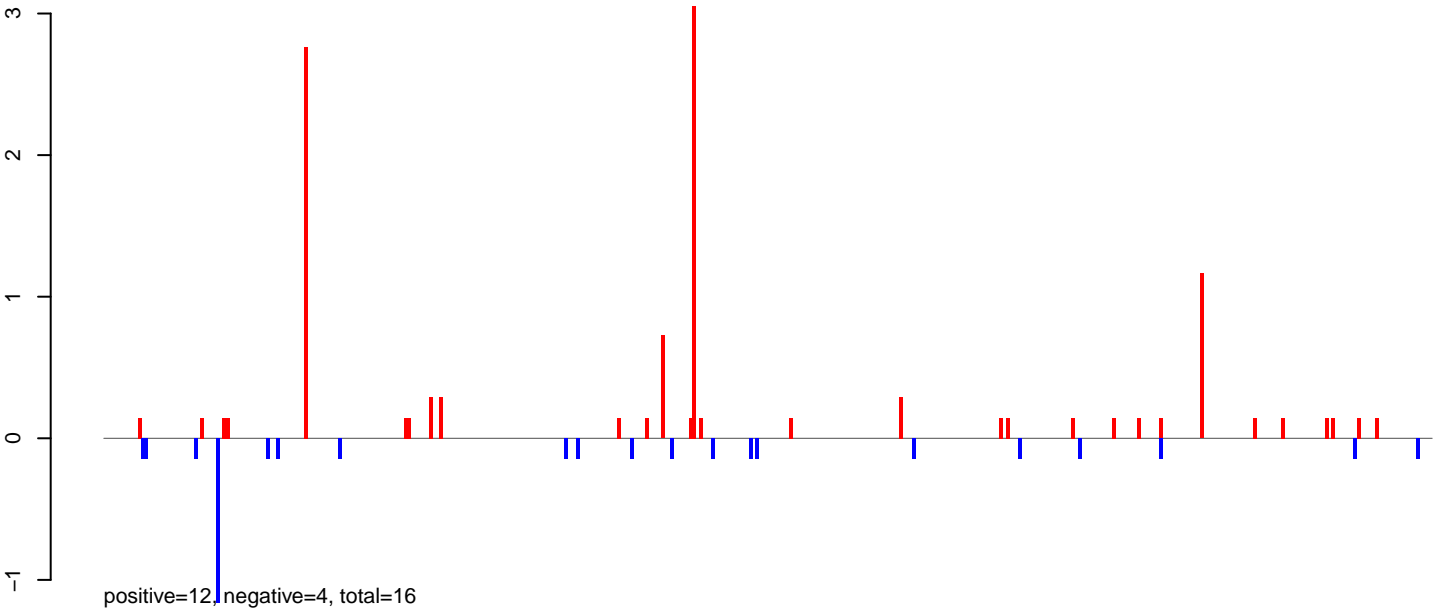
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



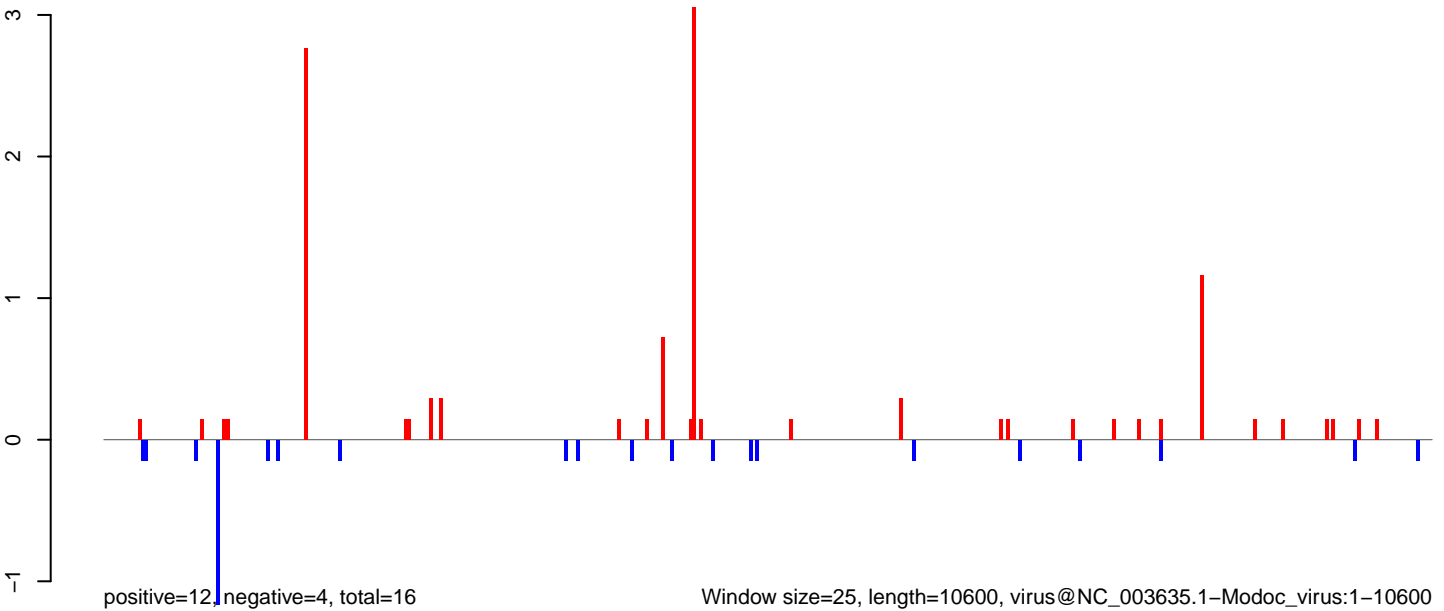
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



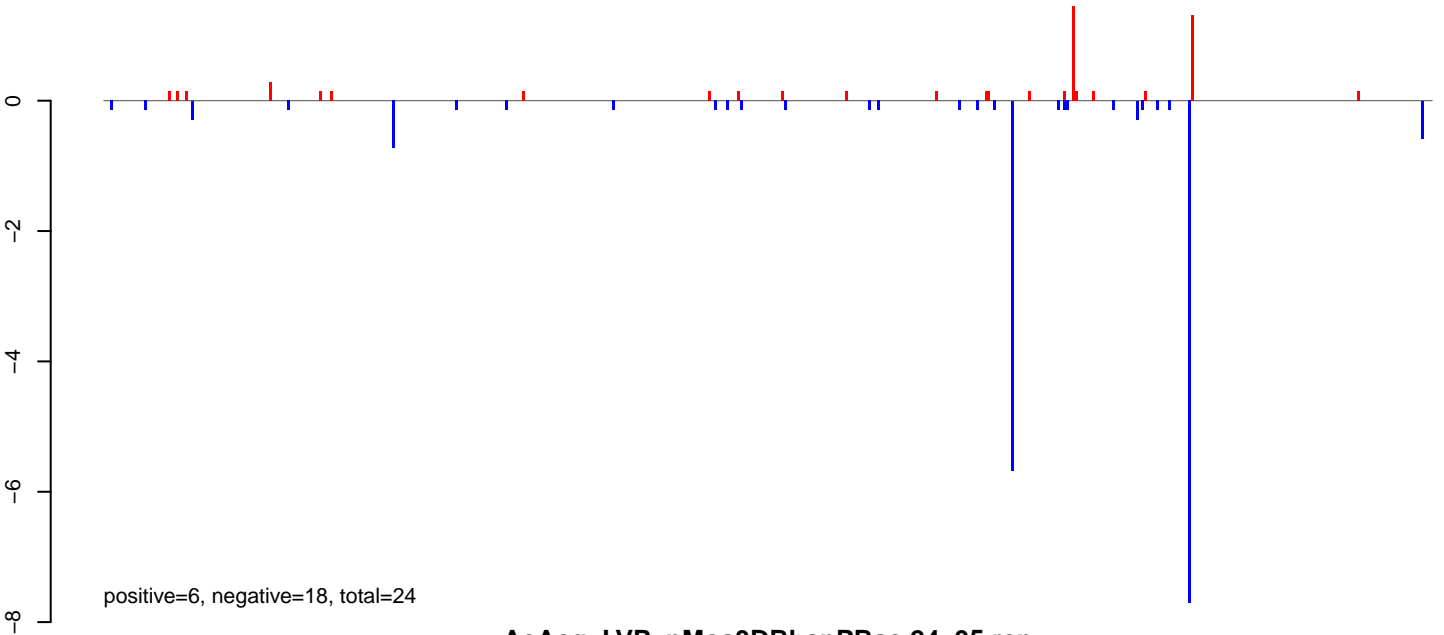
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



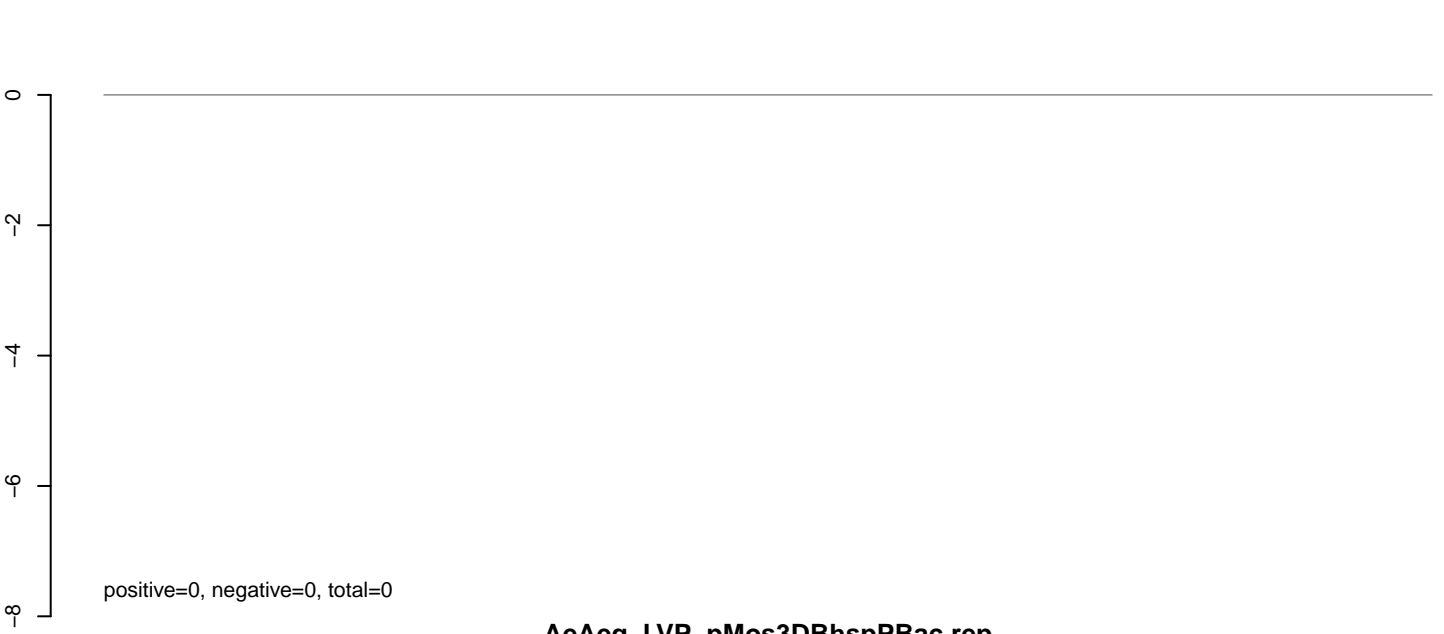
AeAeg_LVP_pMos3DBhspPBac.rep



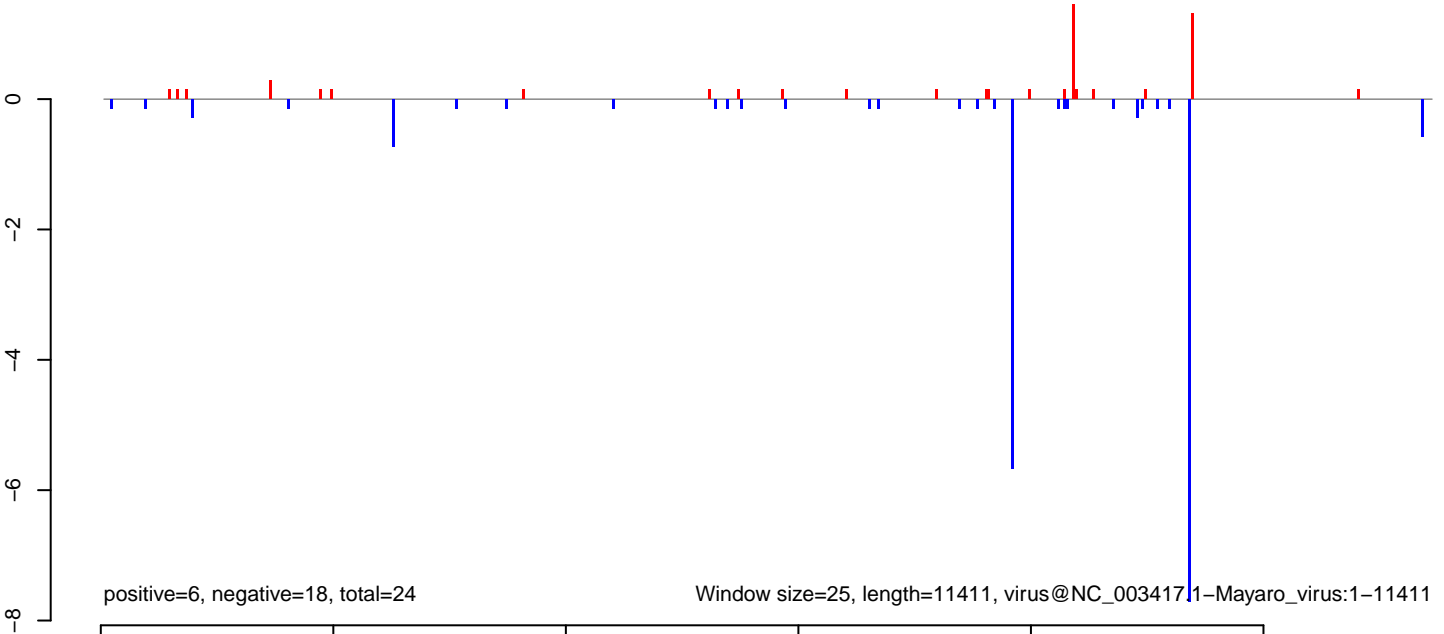
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



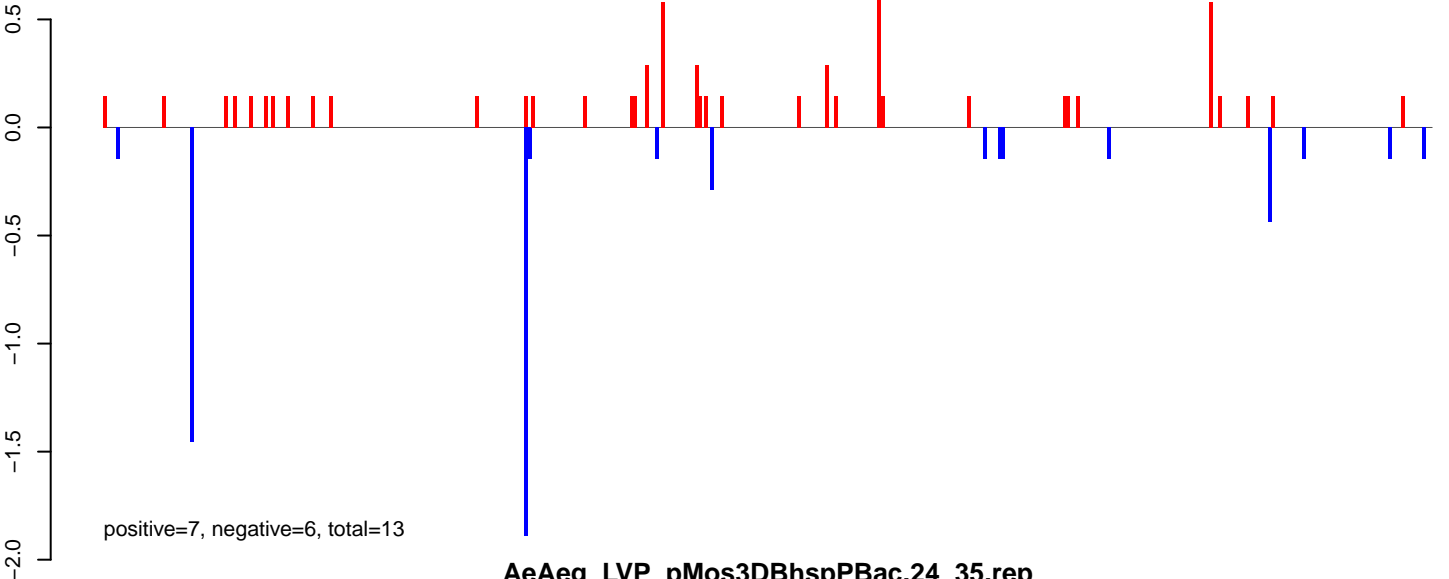
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



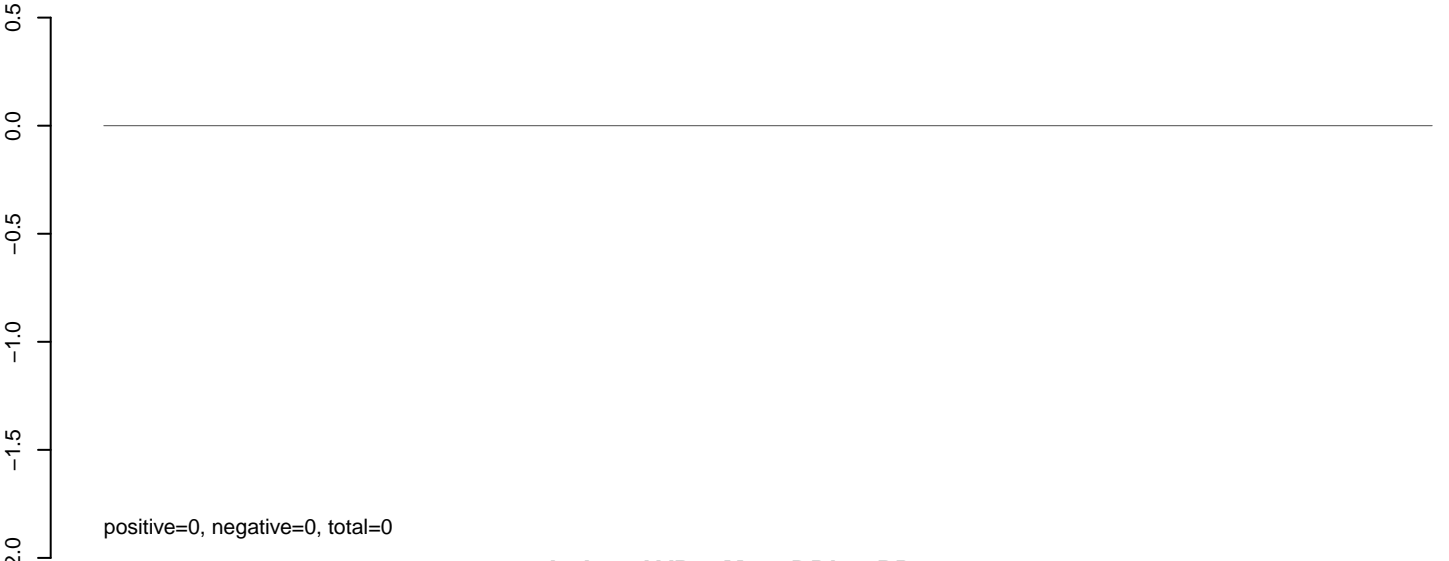
AeAeg_LVP_pMos3DBhspPBac.rep



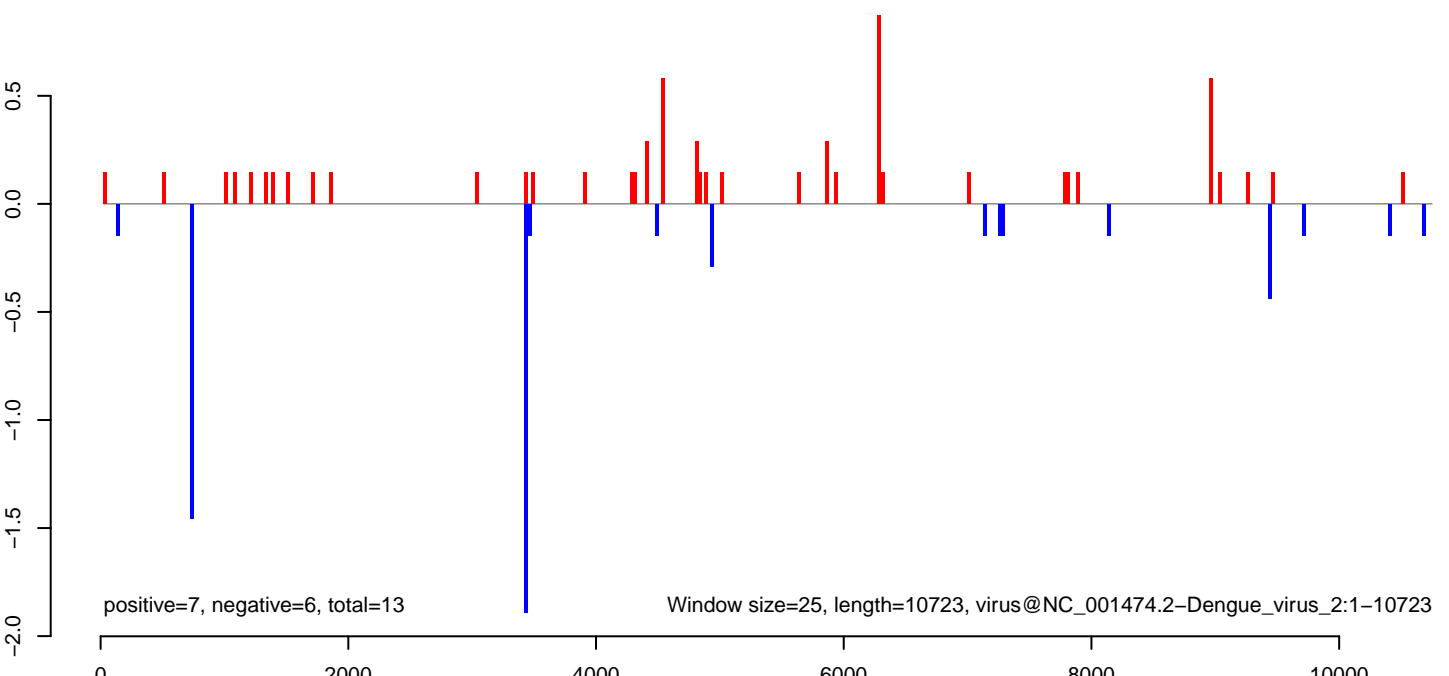
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



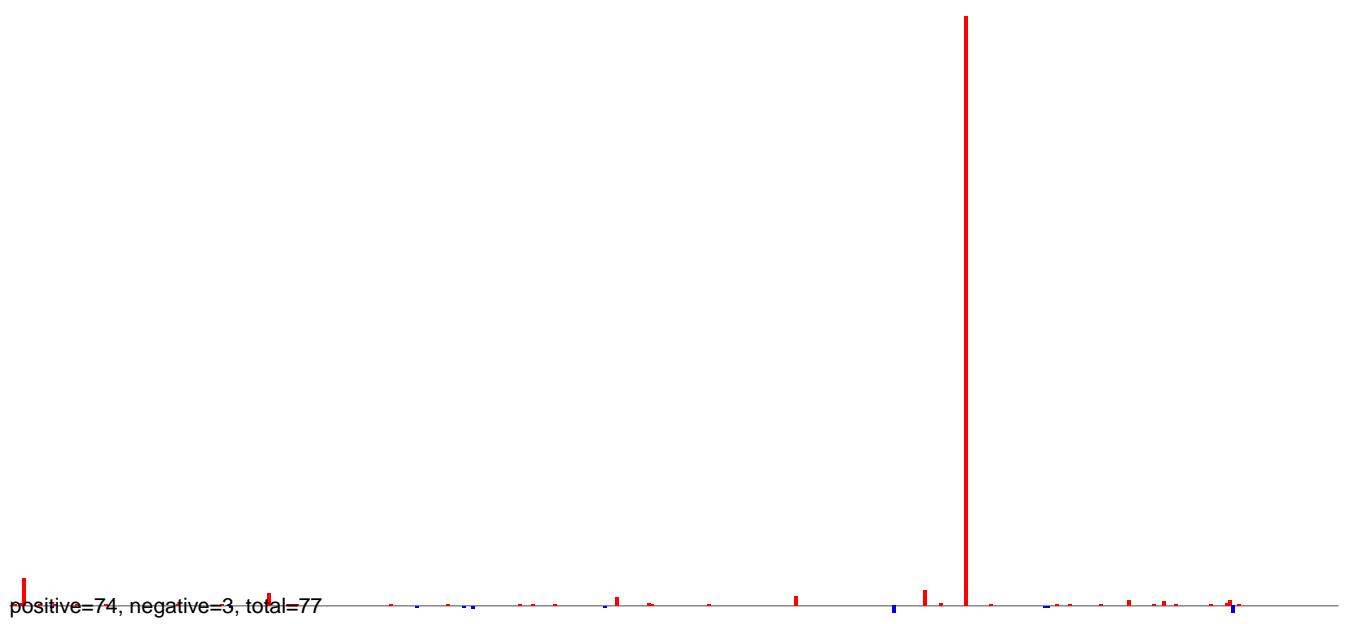
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



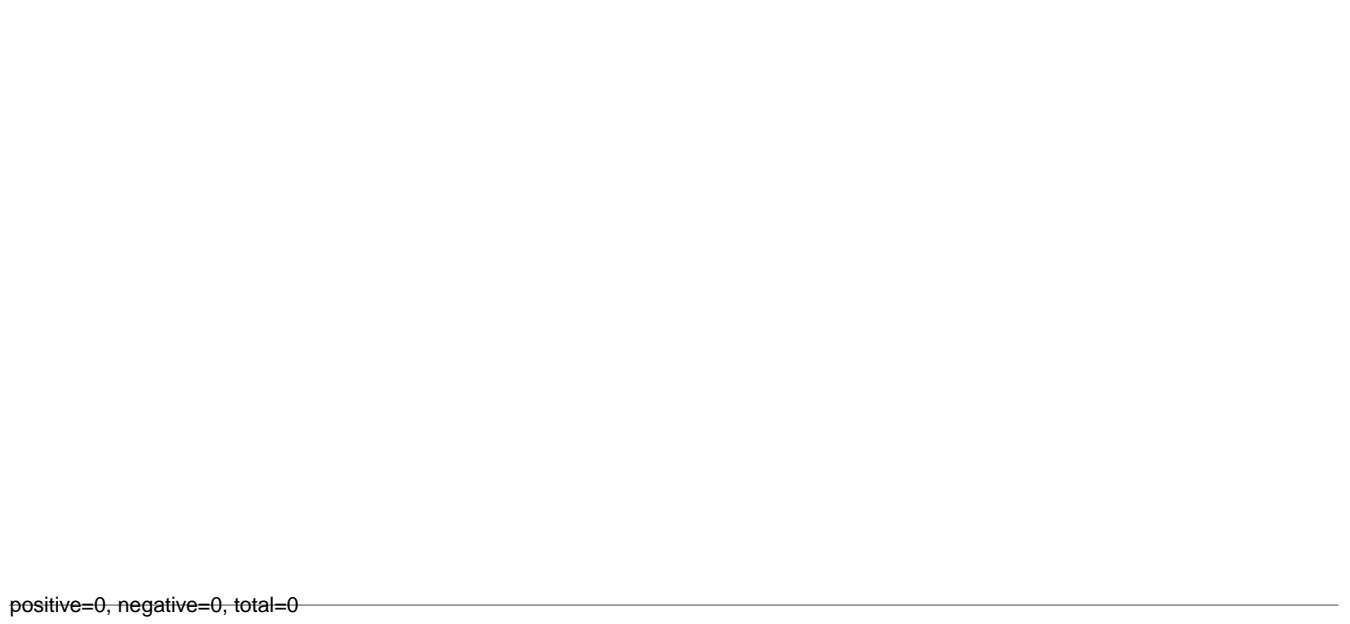
AeAeg_LVP_pMos3DBhspPBac.rep



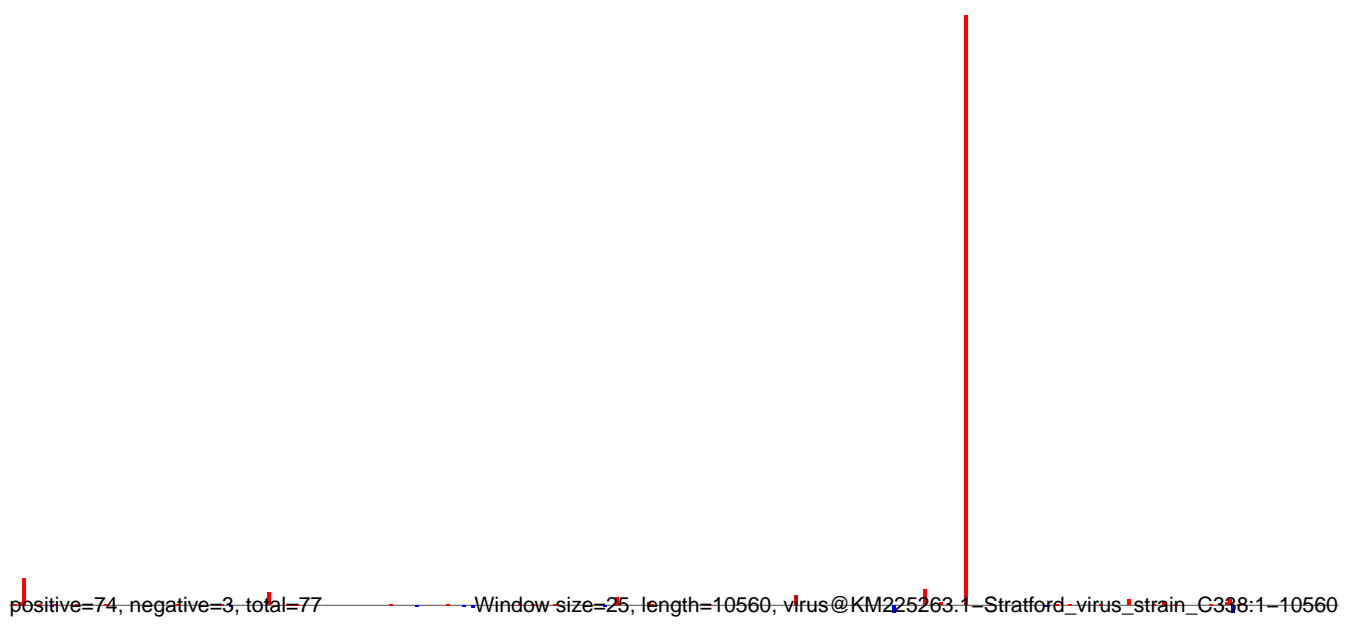
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



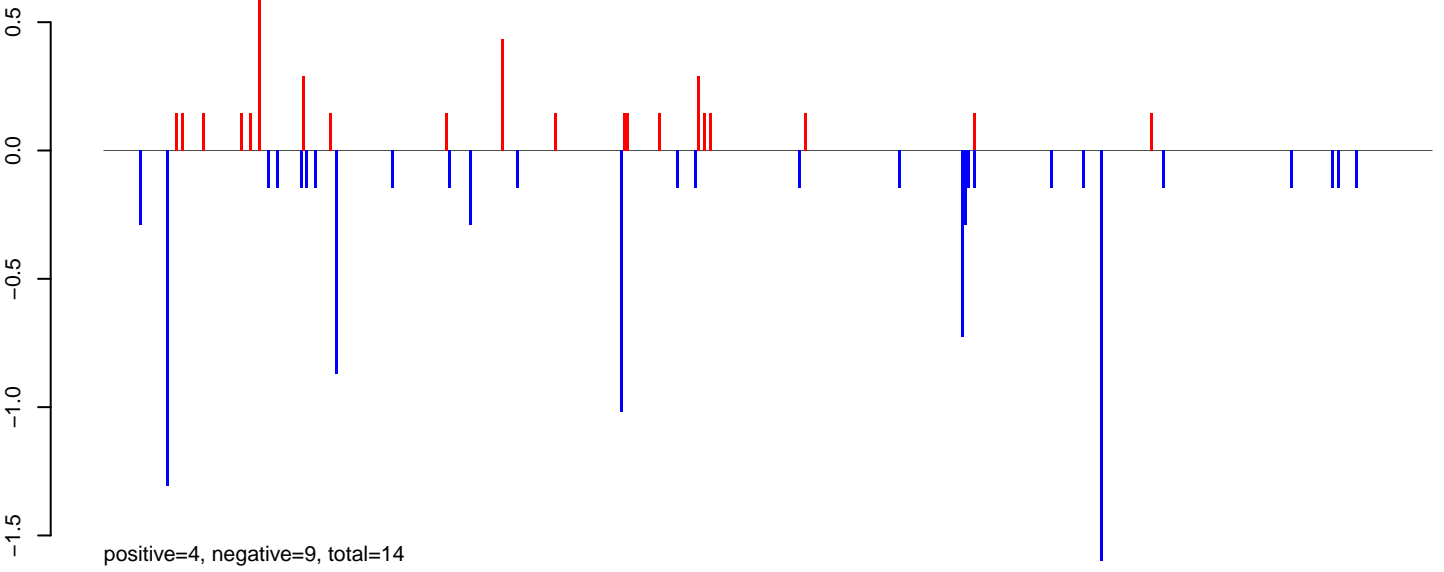
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



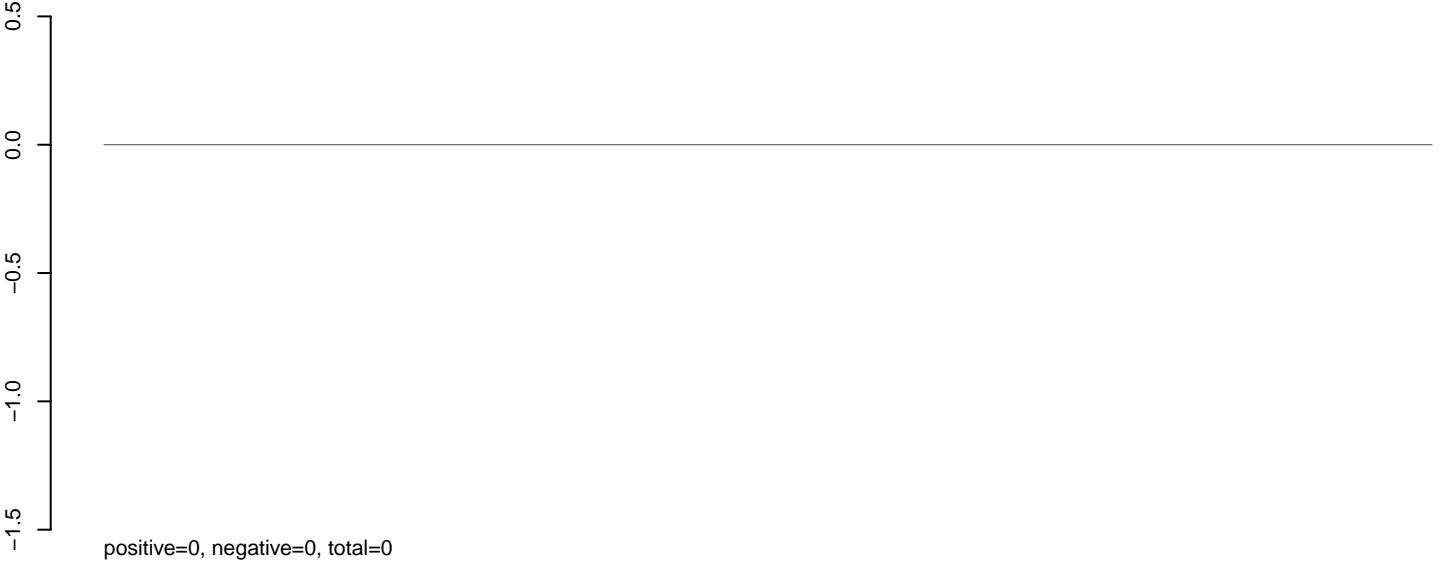
AeAeg_LVP_pMos3DBhspPBac.rep



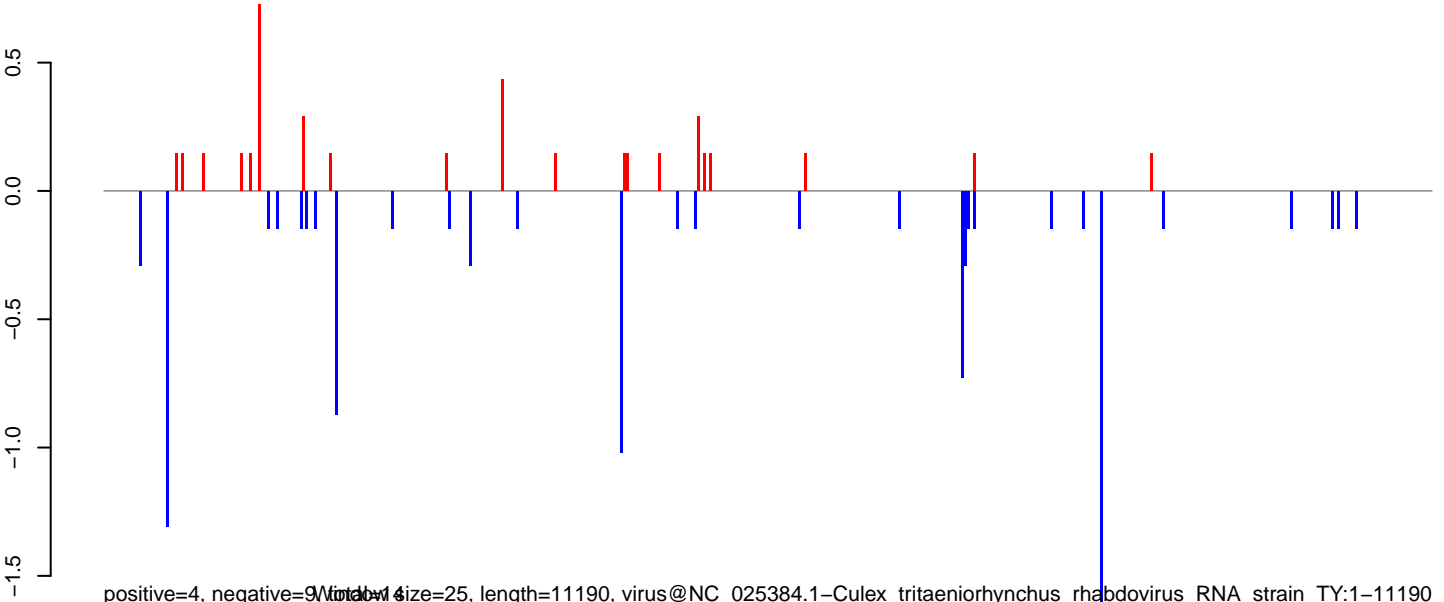
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



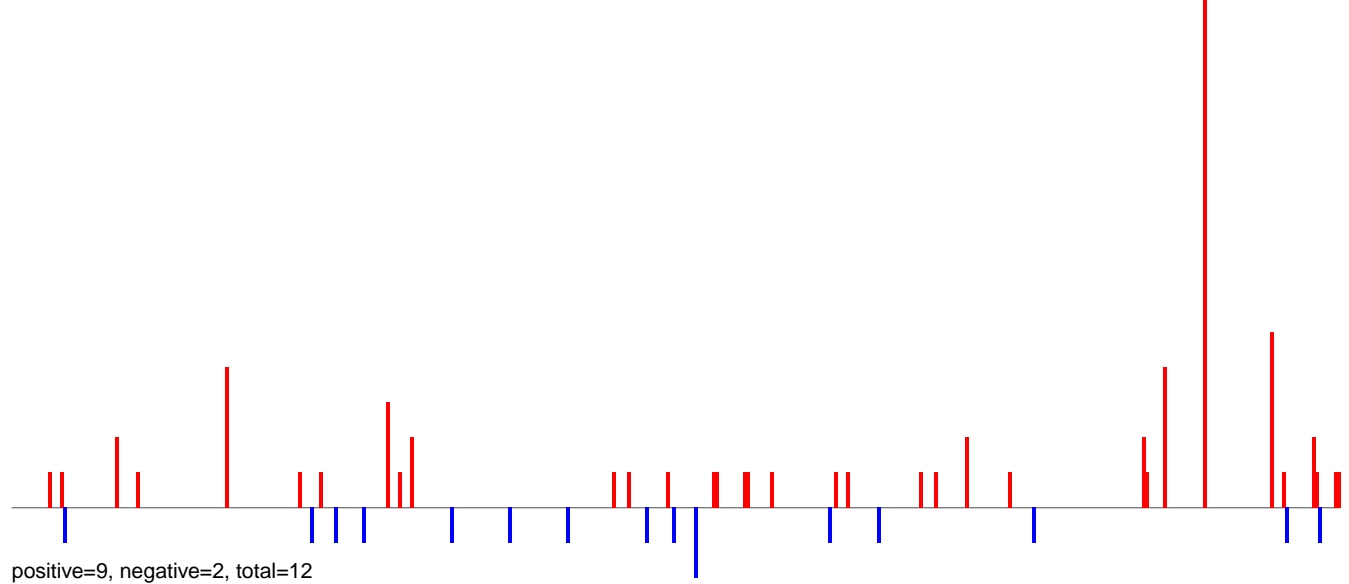
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



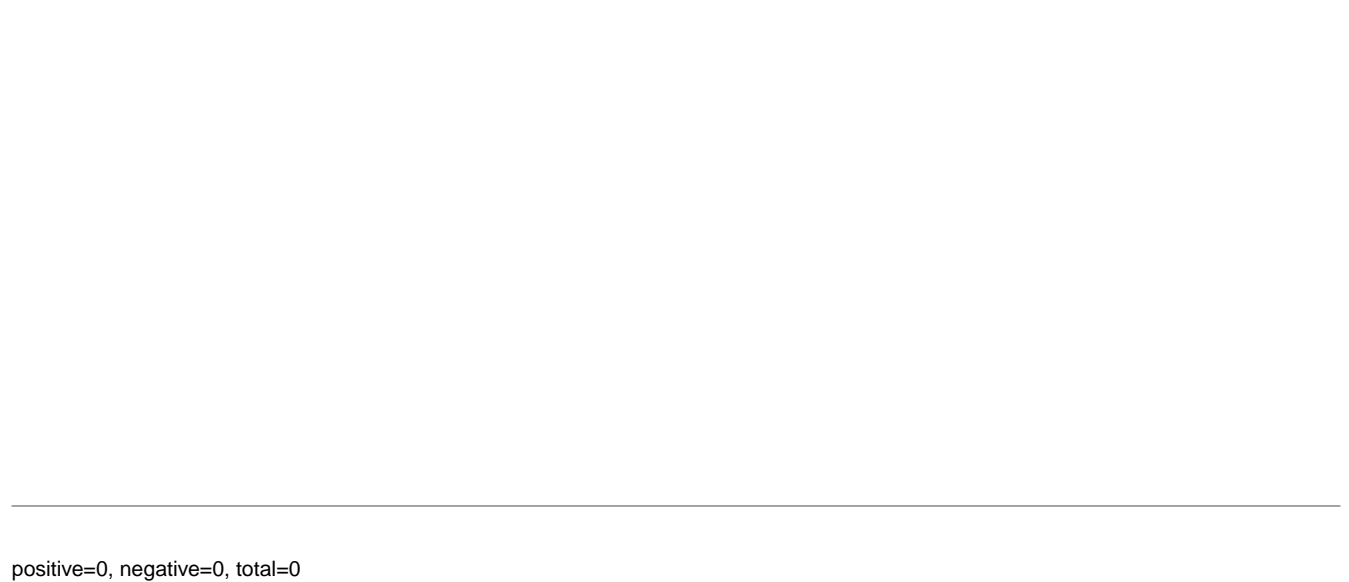
AeAeg_LVP_pMos3DBhspPBac.rep



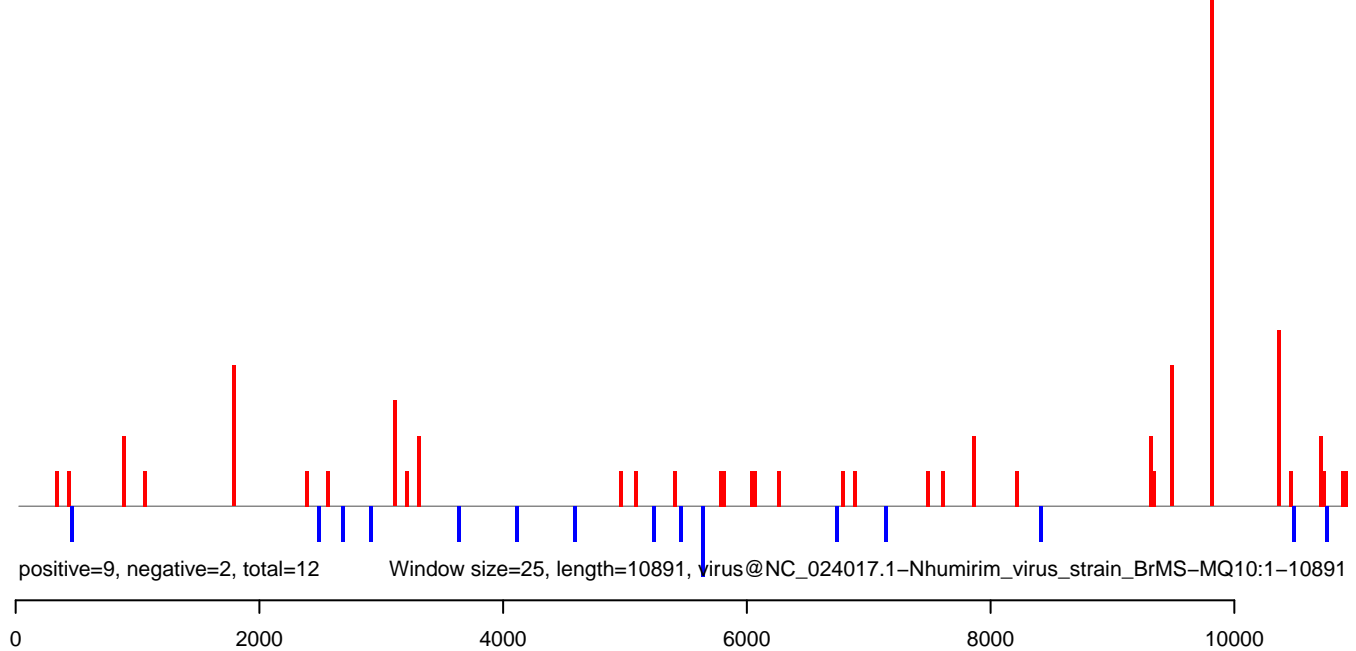
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



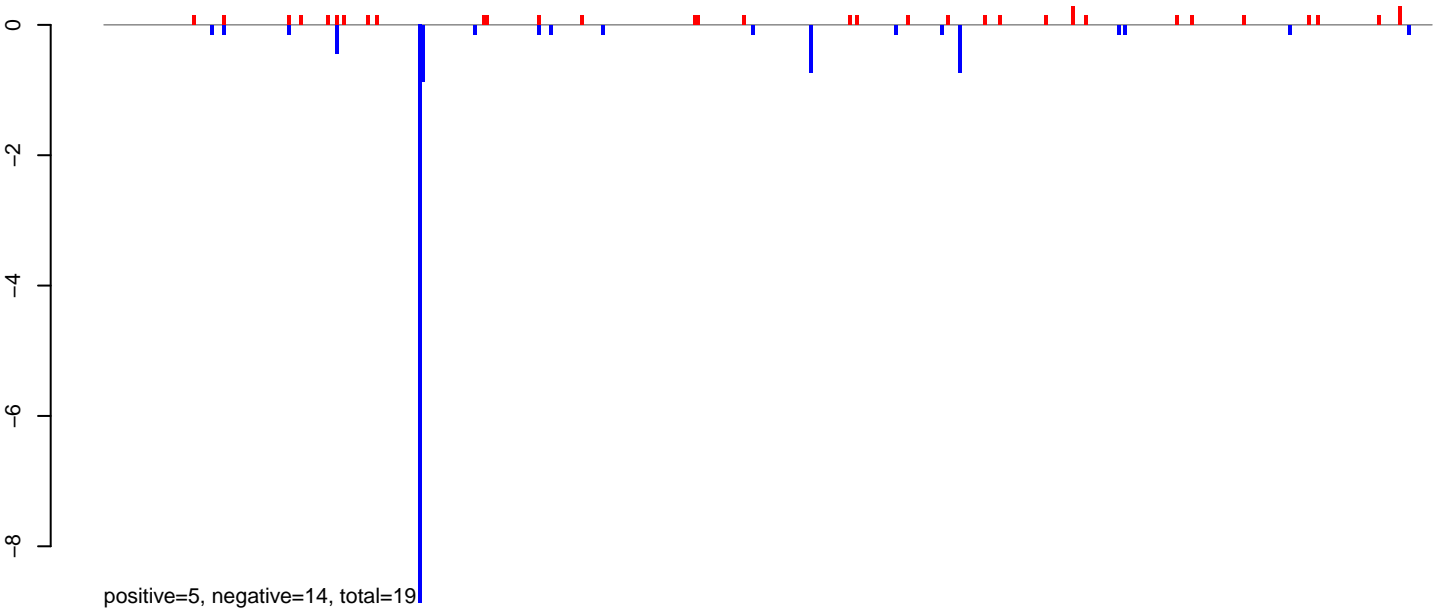
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



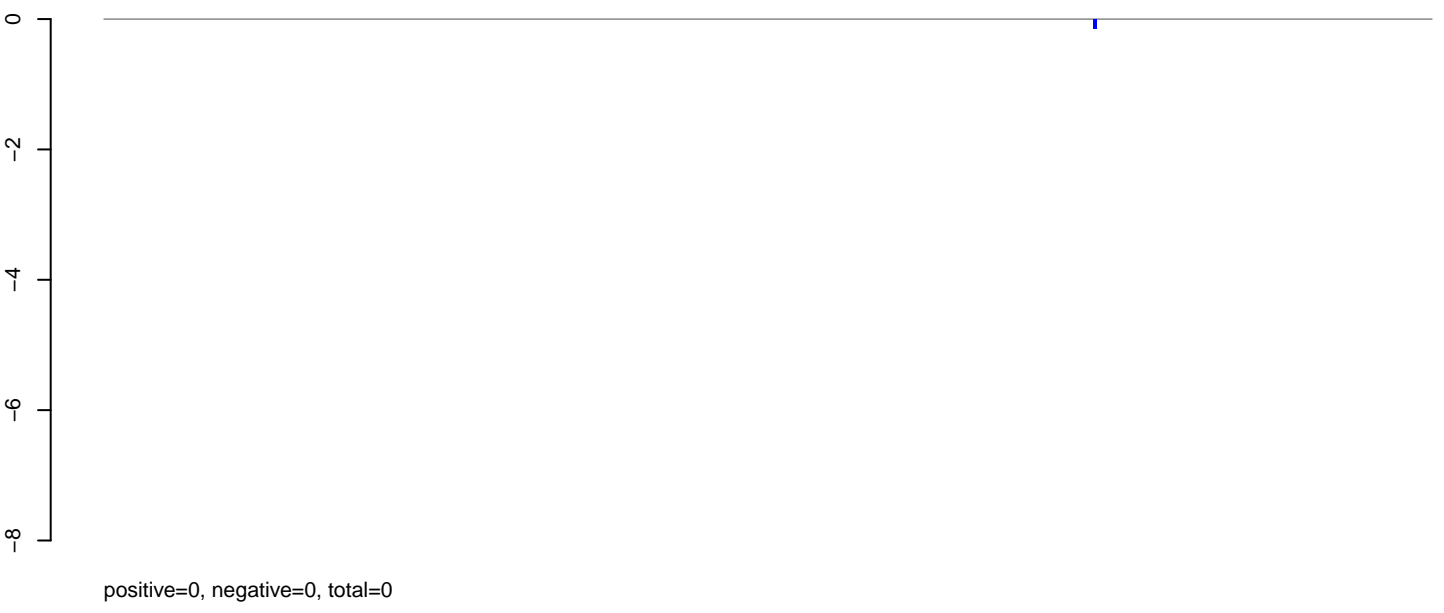
AeAeg_LVP_pMos3DBhspPBac.rep



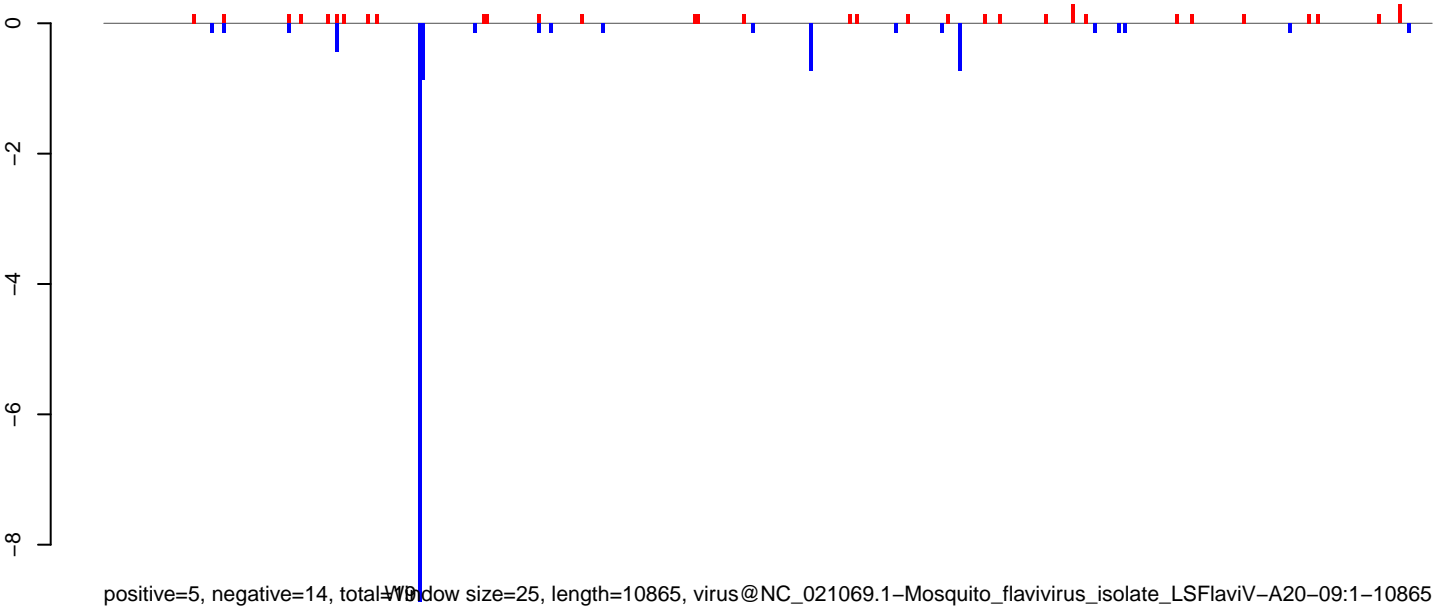
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



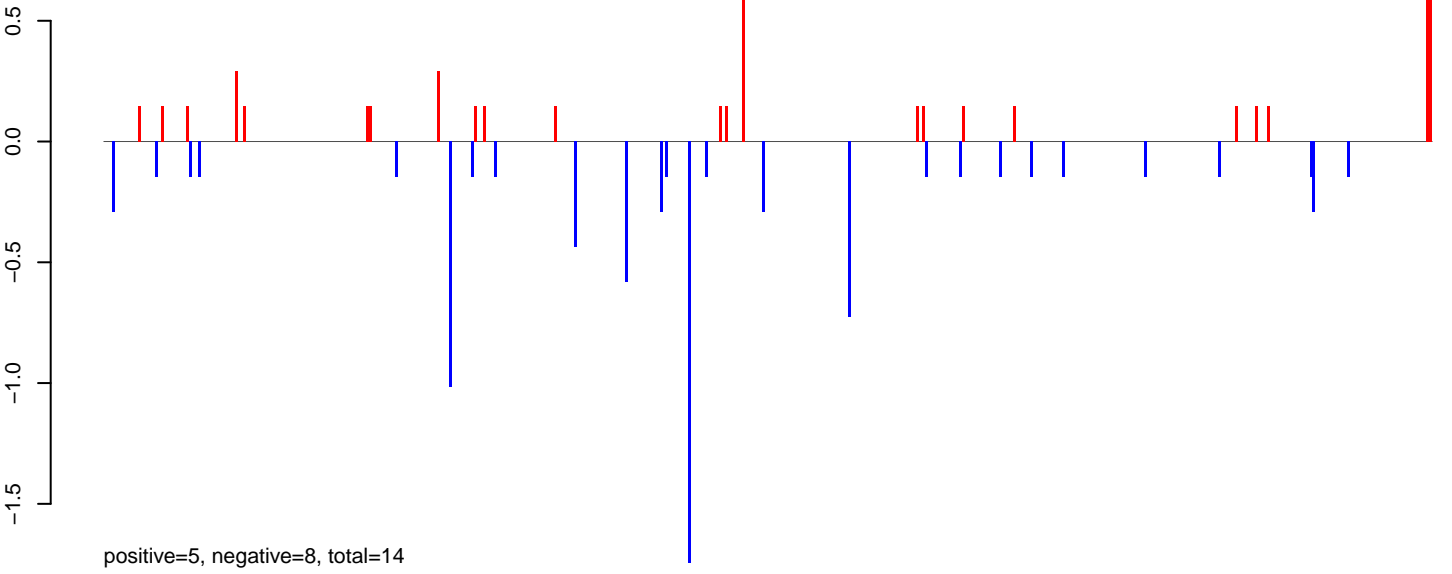
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



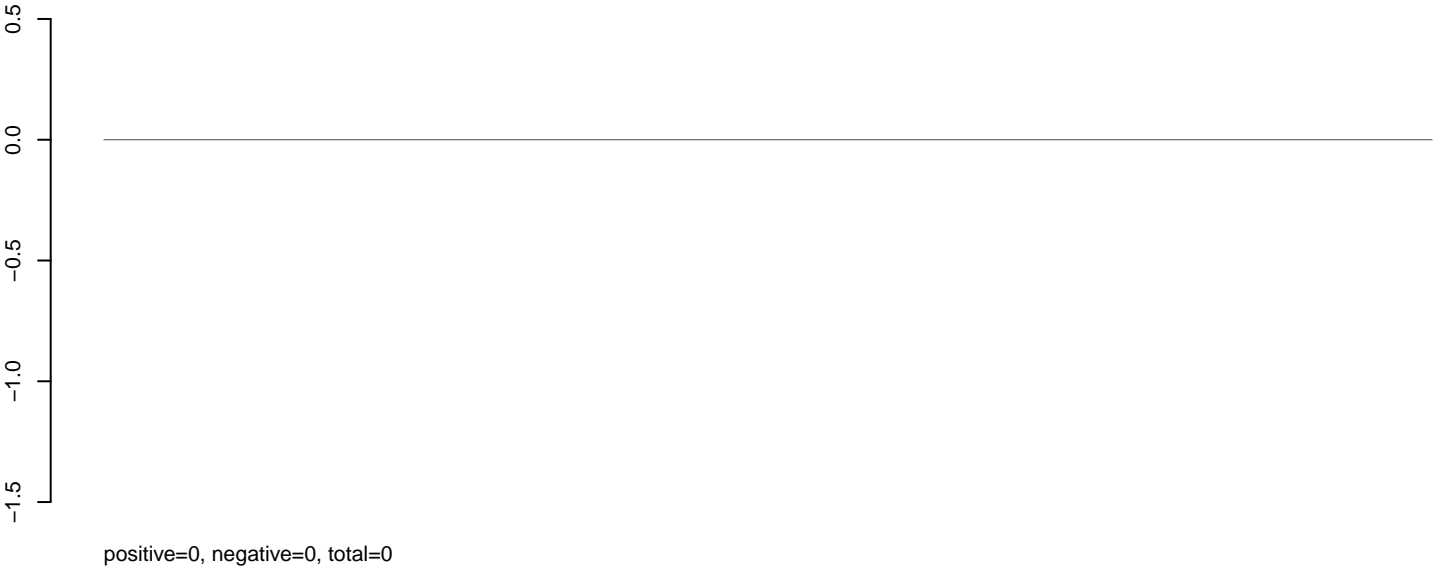
AeAeg_LVP_pMos3DBhspPBac.rep



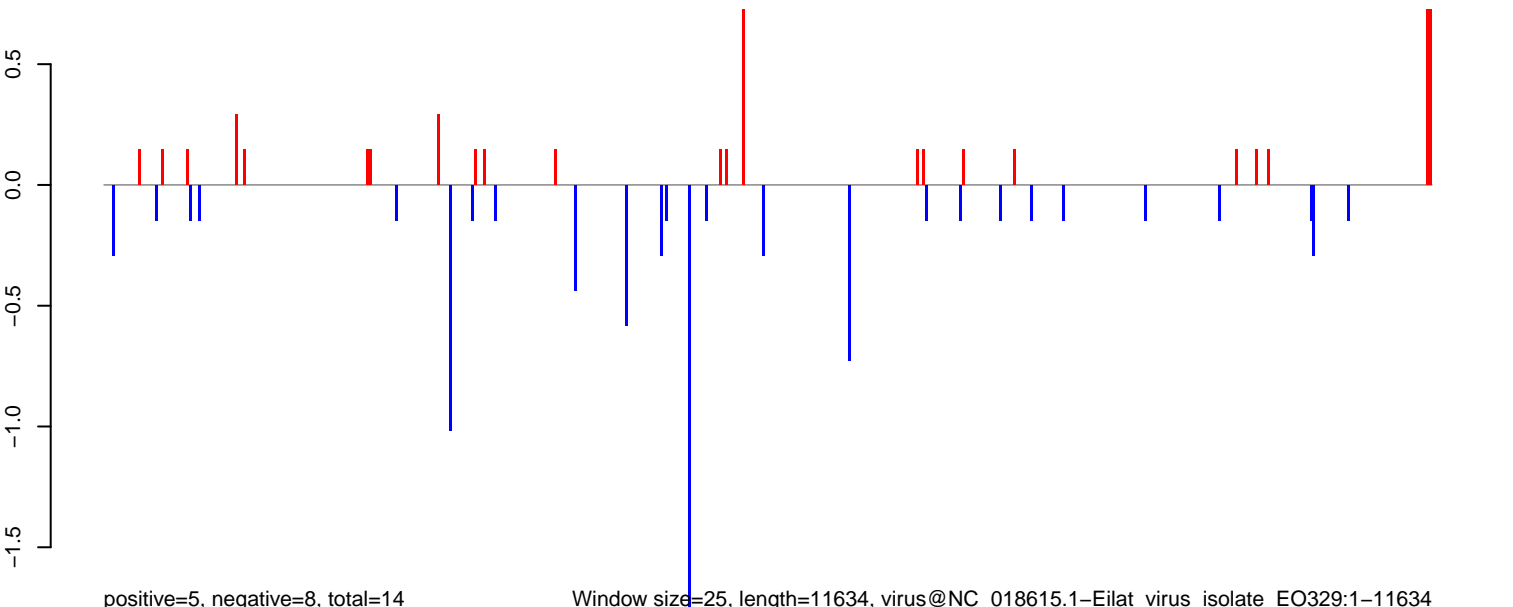
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



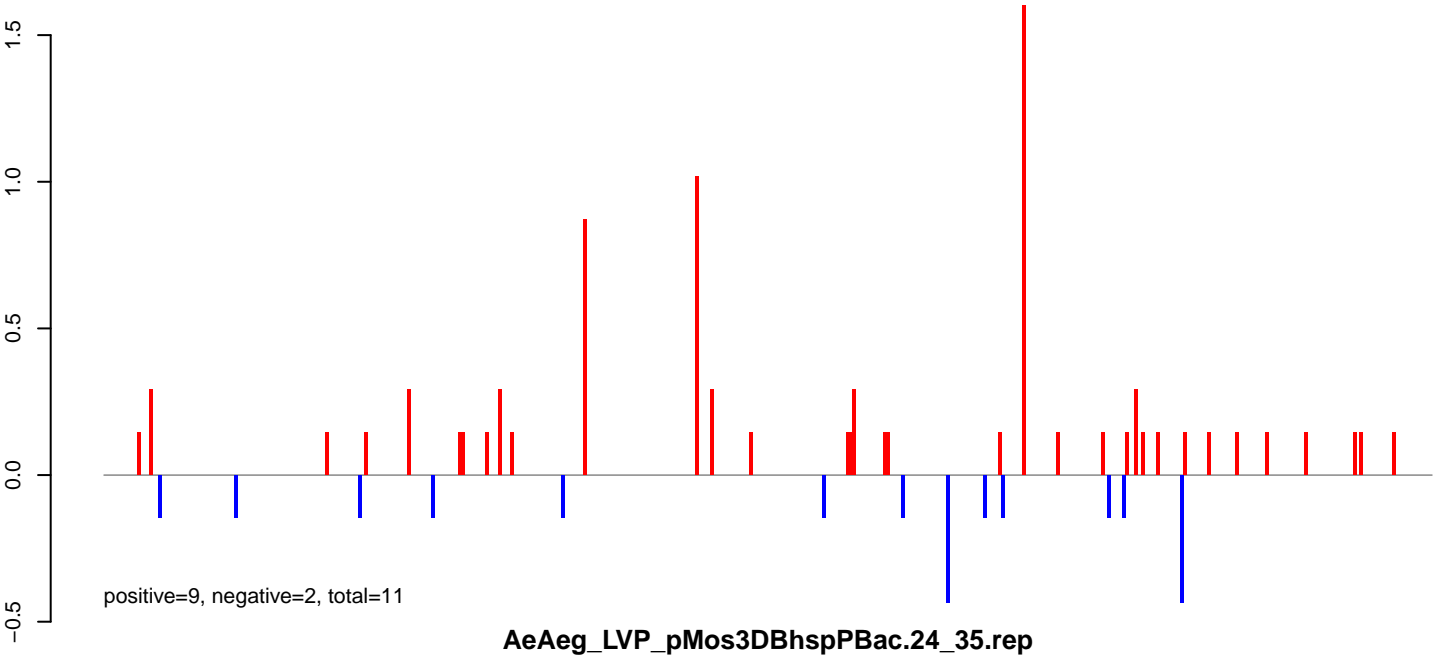
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



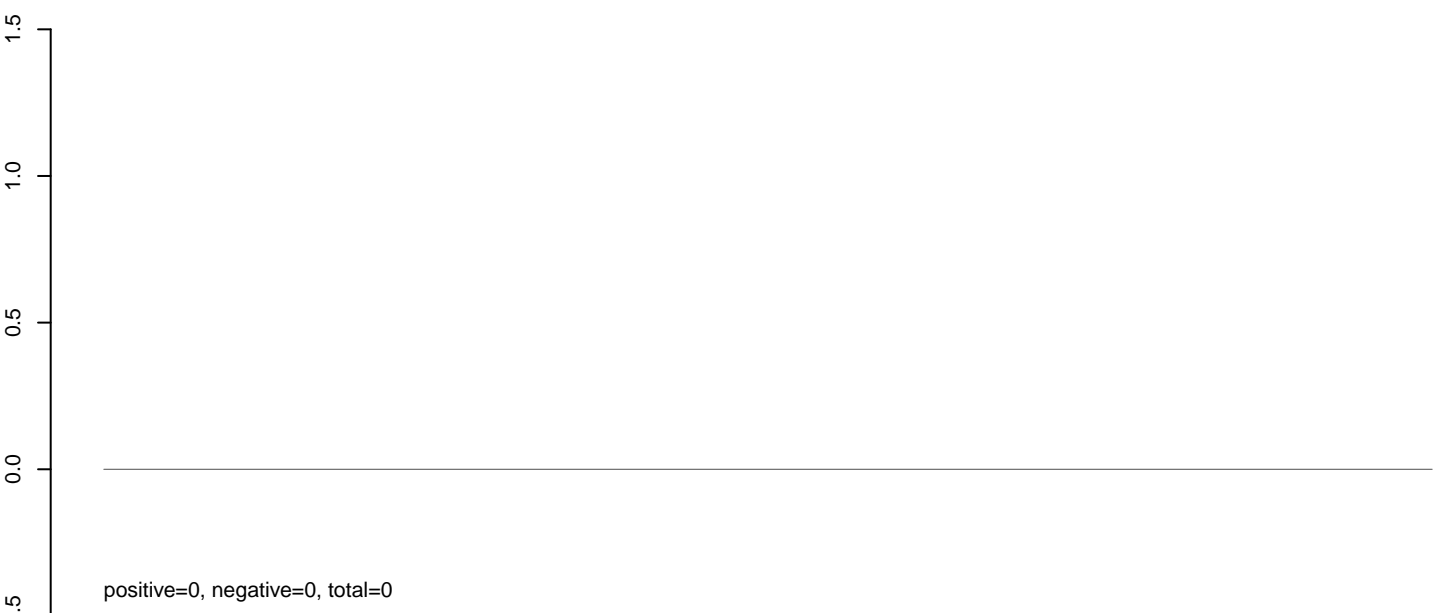
AeAeg_LVP_pMos3DBhspPBac.rep



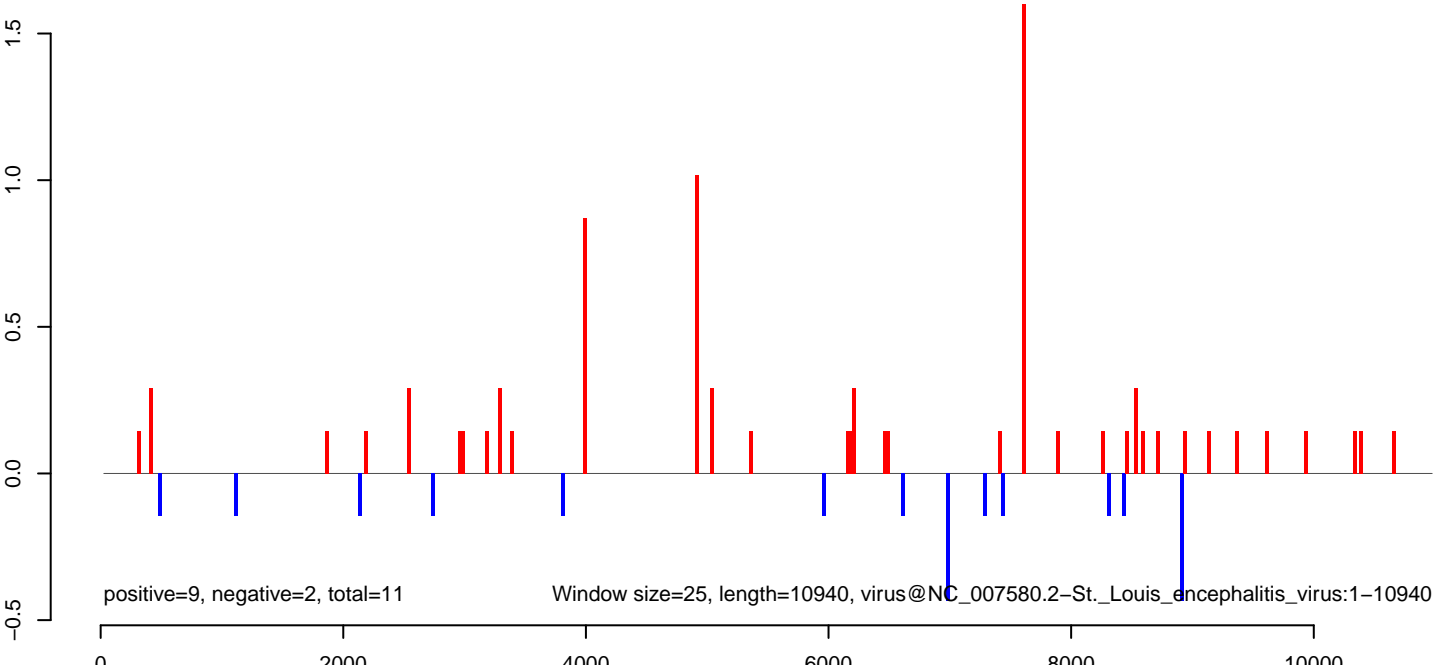
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



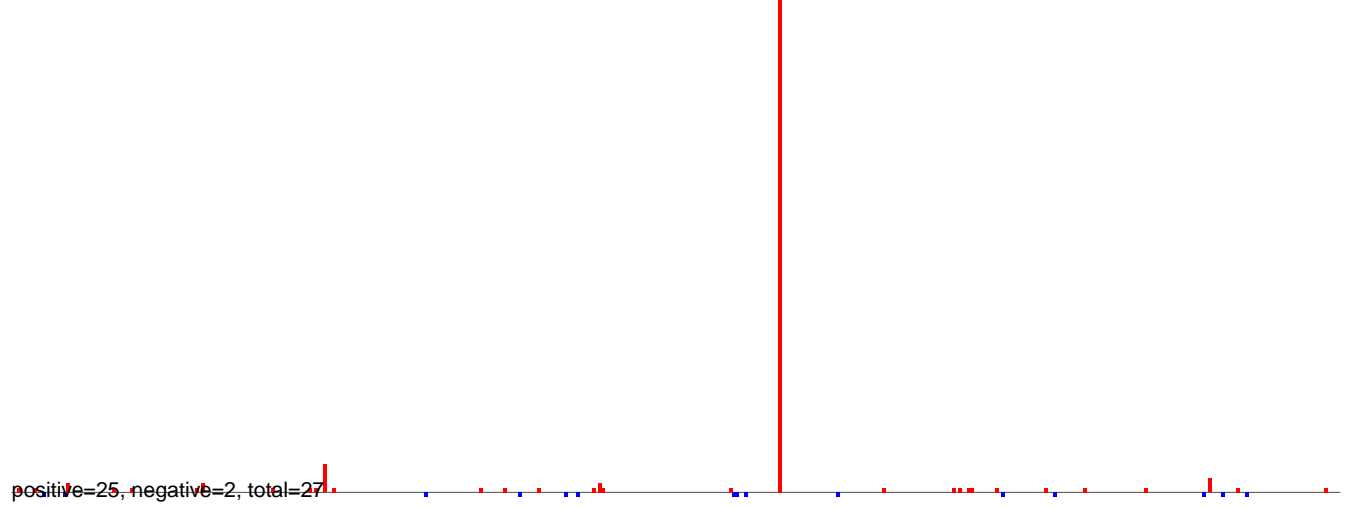
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



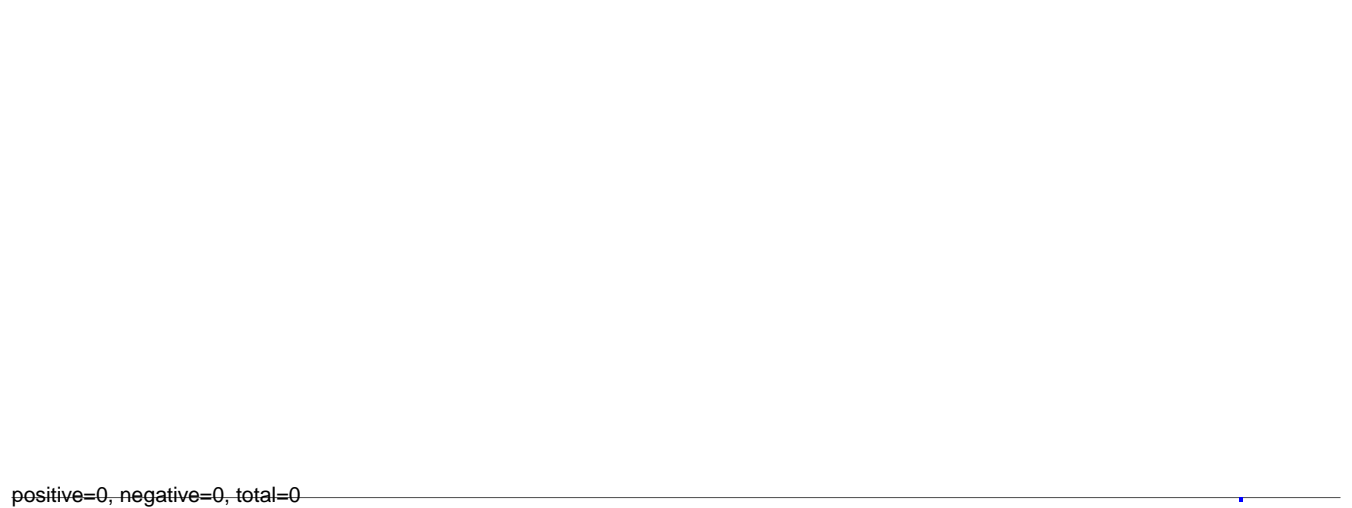
AeAeg_LVP_pMos3DBhspPBac.rep



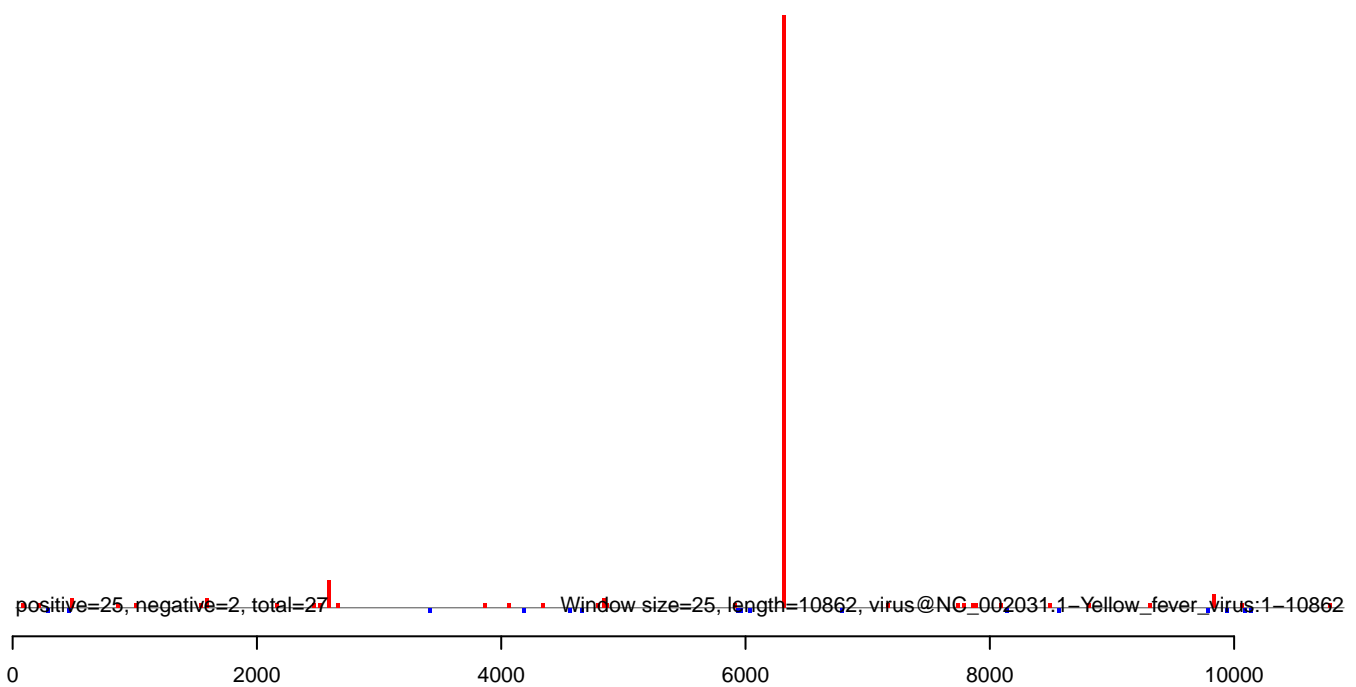
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



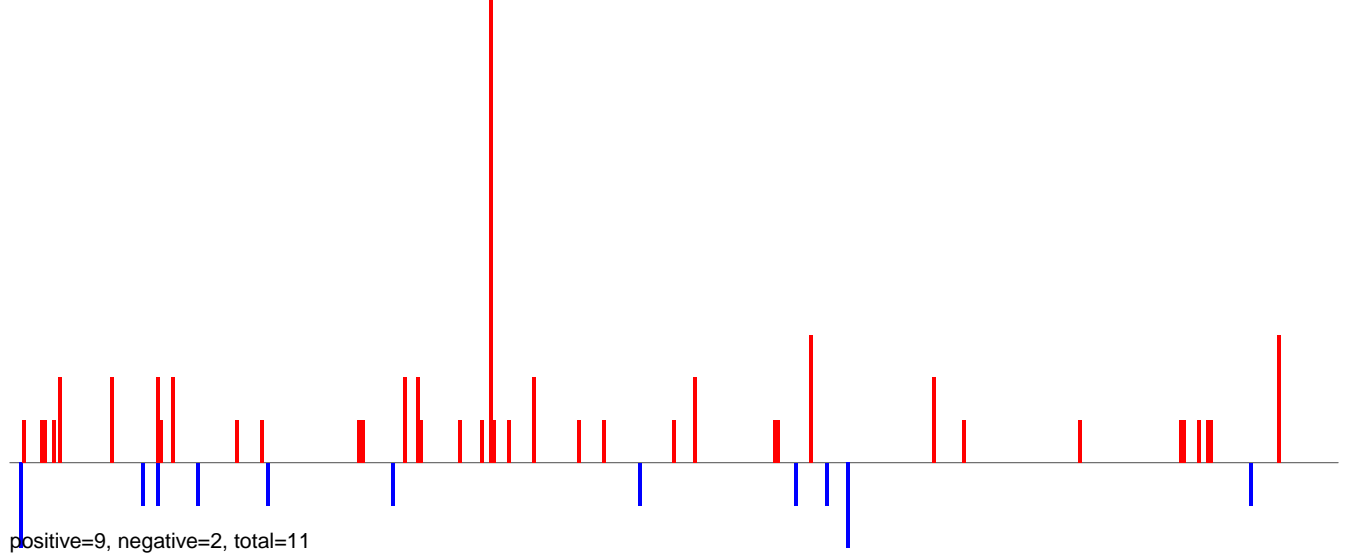
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



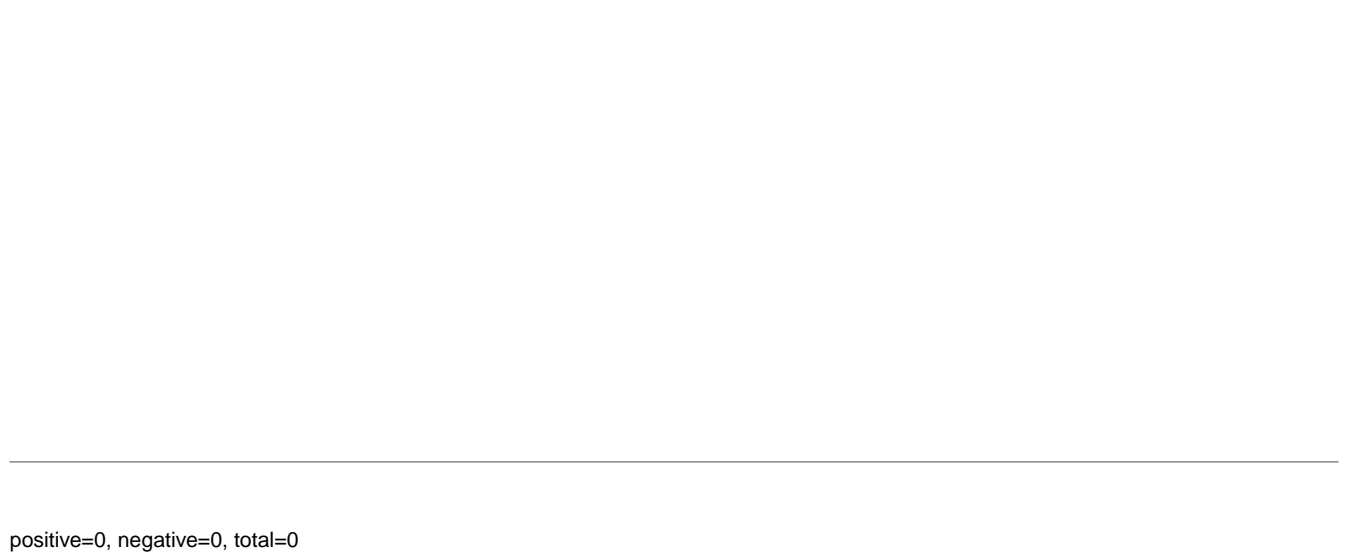
AeAeg_LVP_pMos3DBhspPBac.rep



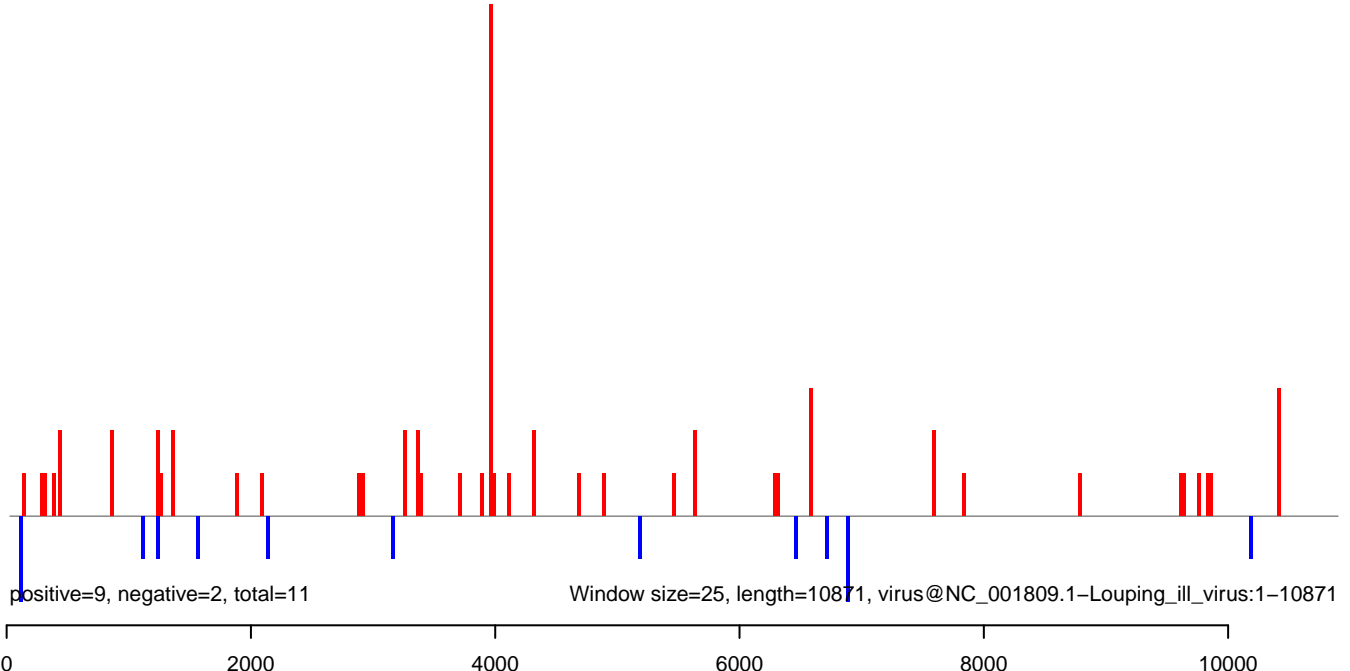
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



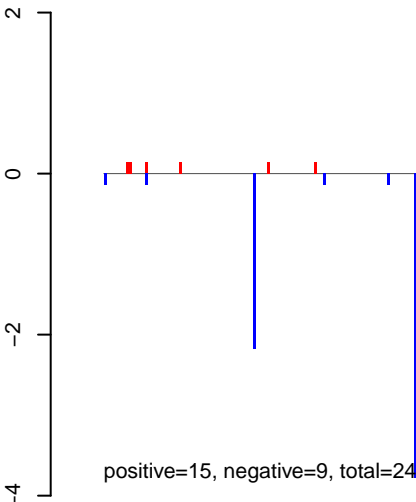
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



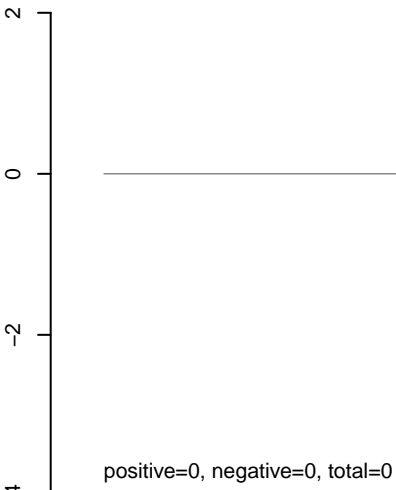
AeAeg_LVP_pMos3DBhspPBac.rep



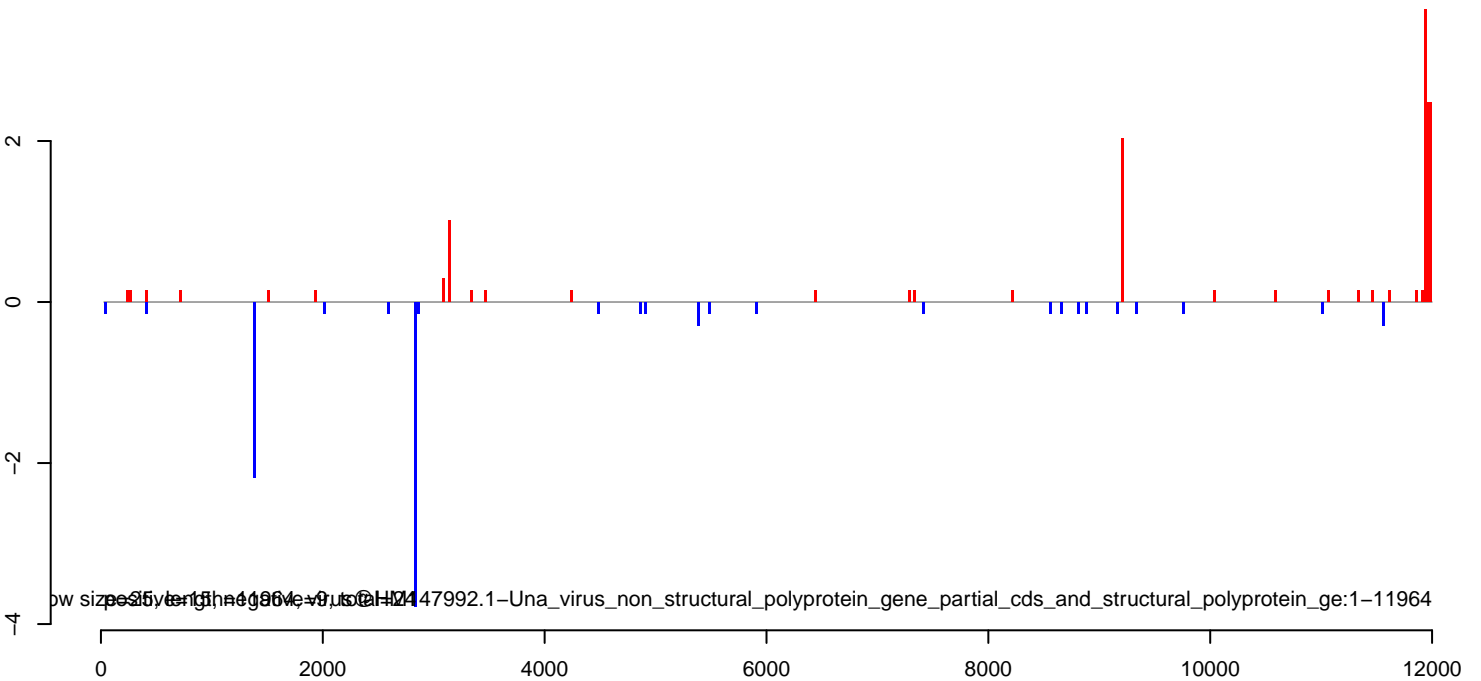
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



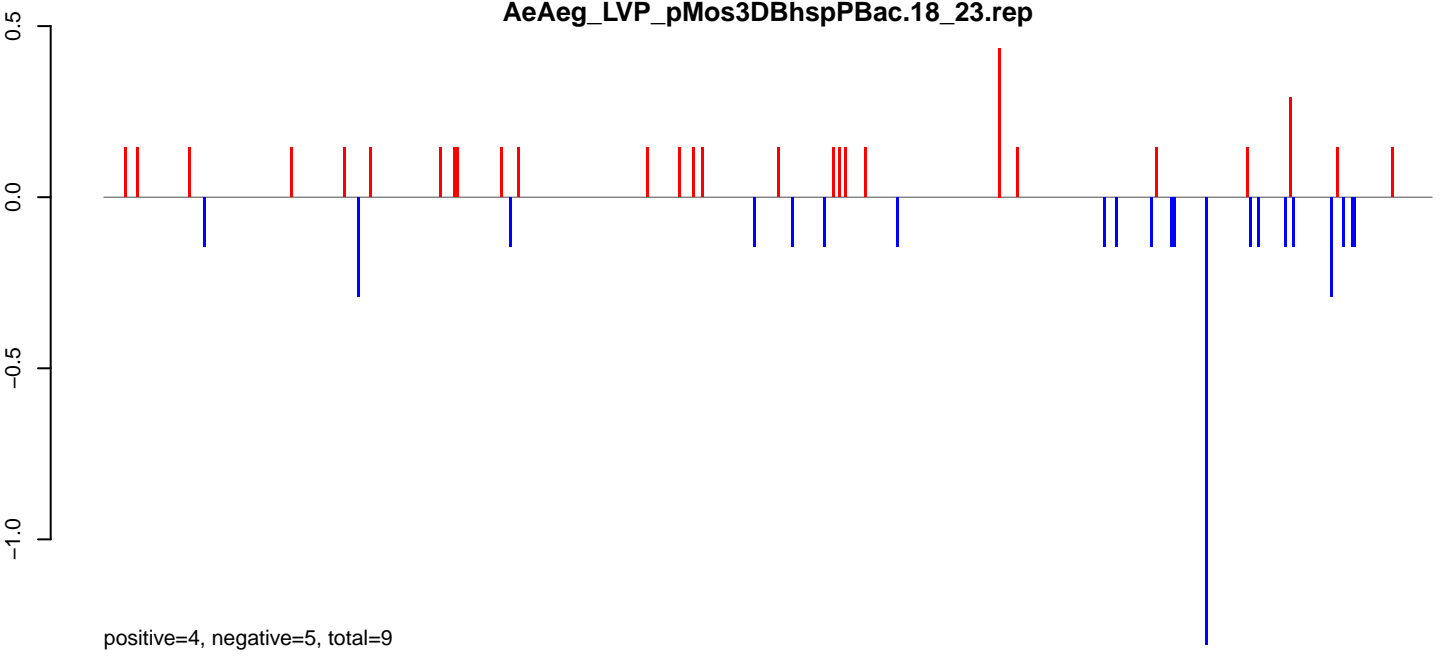
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



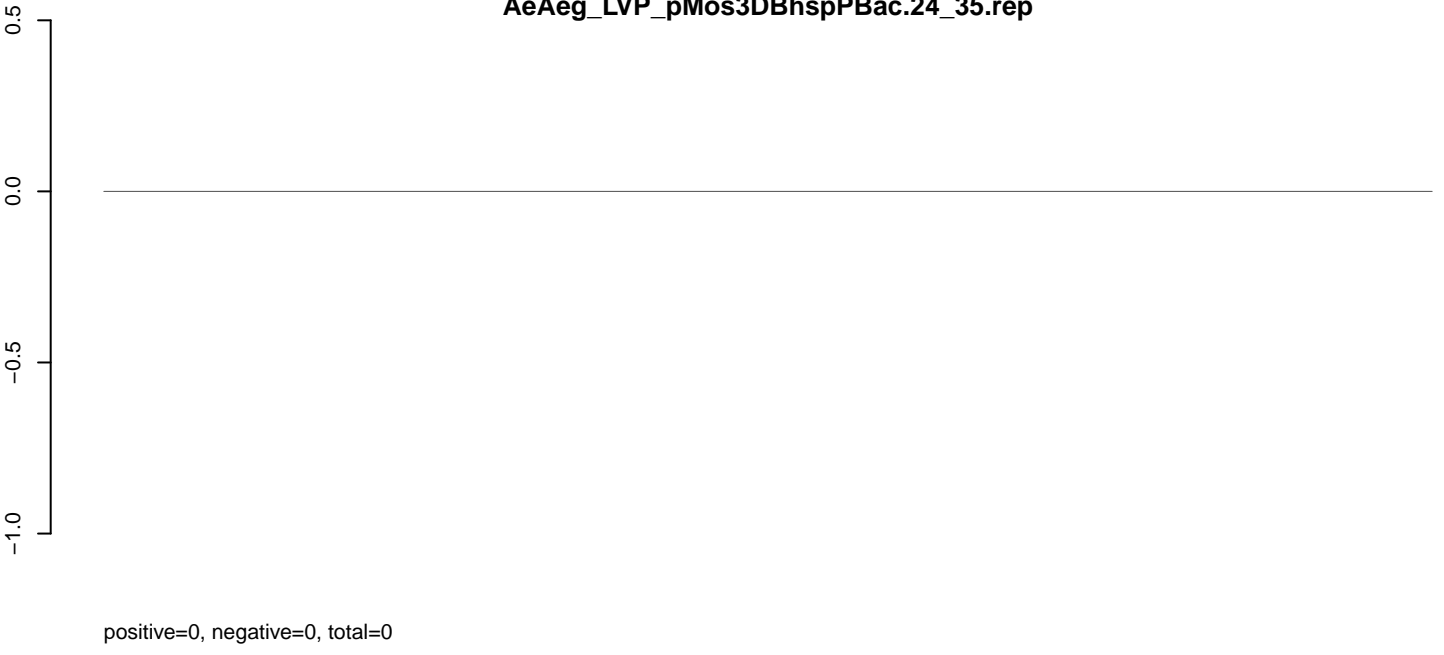
AeAeg_LVP_pMos3DBhspPBac.rep



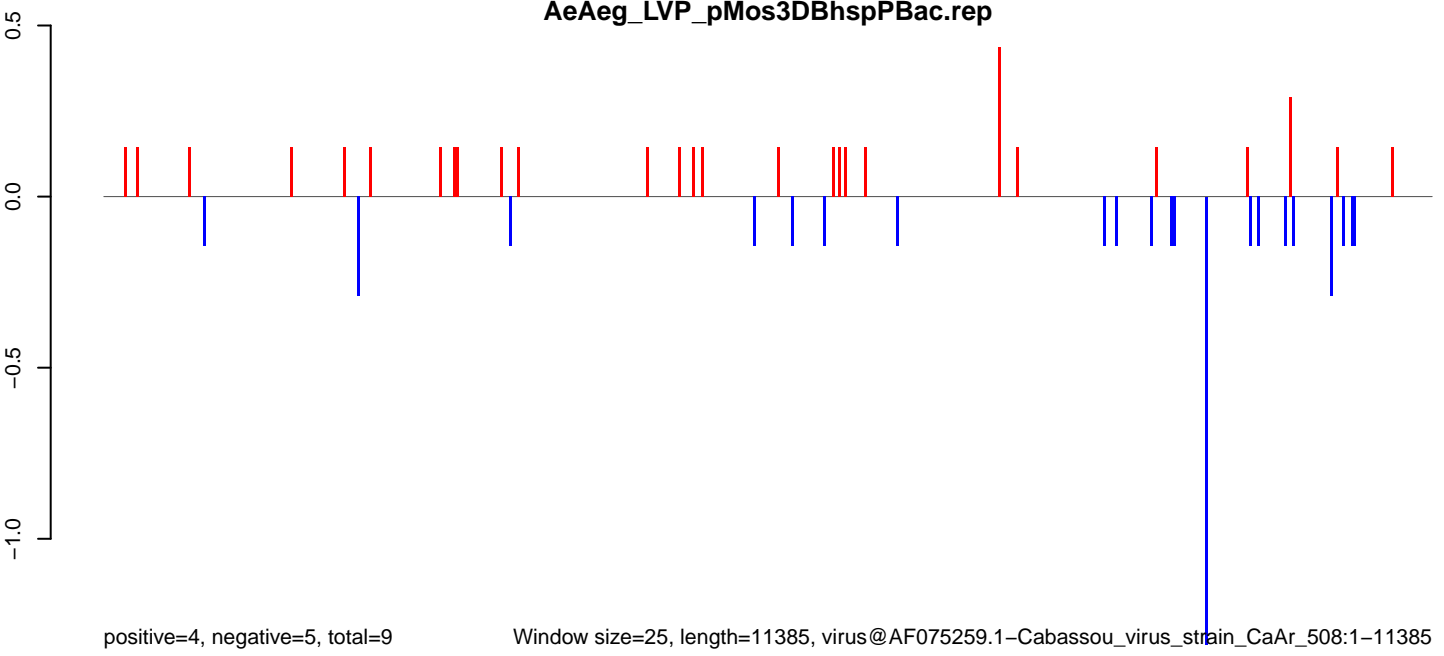
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



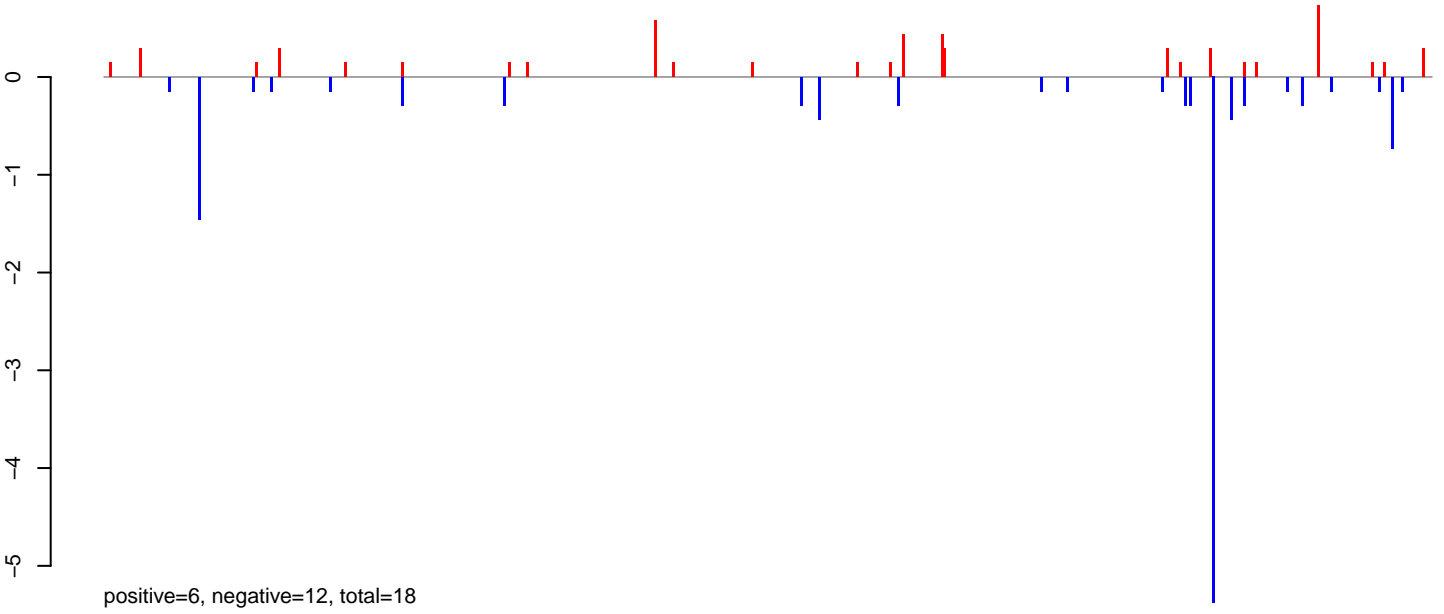
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



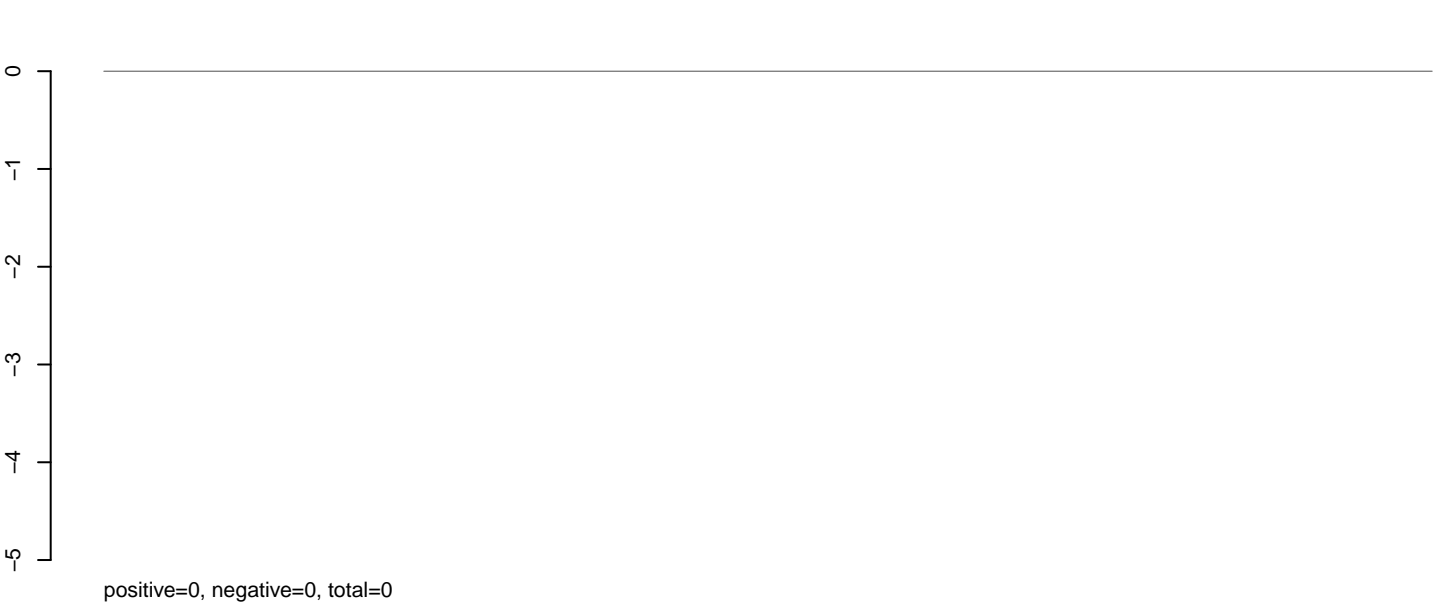
AeAeg_LVP_pMos3DBhspPBac.rep



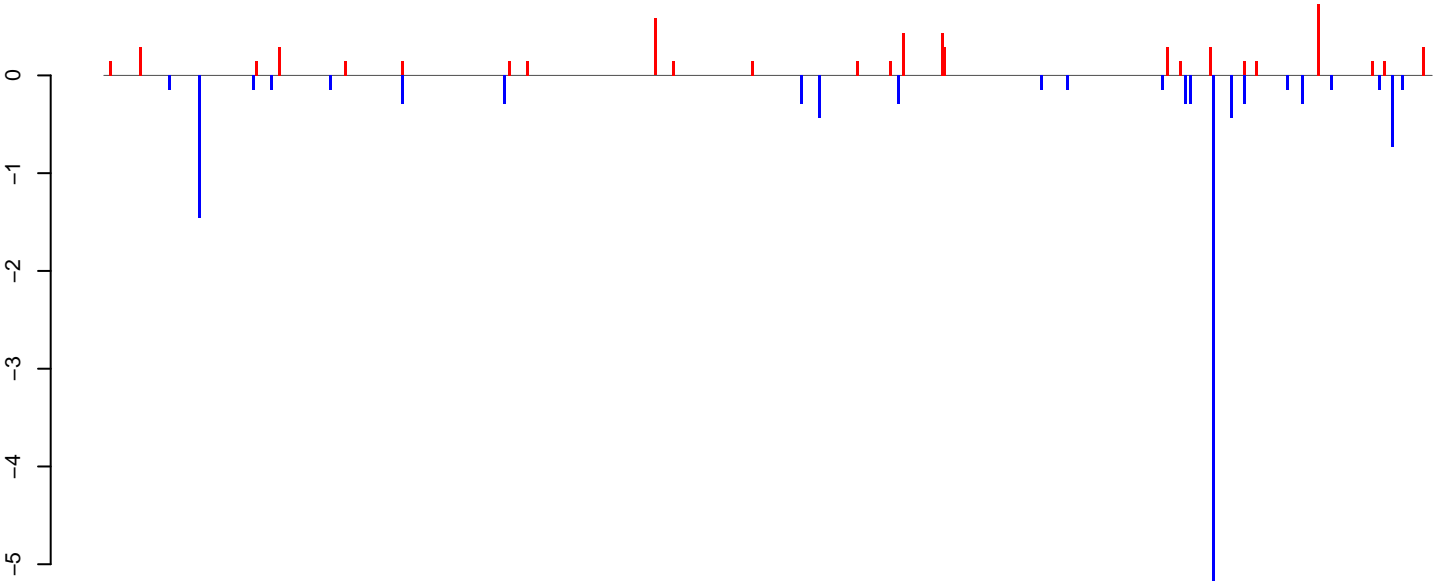
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



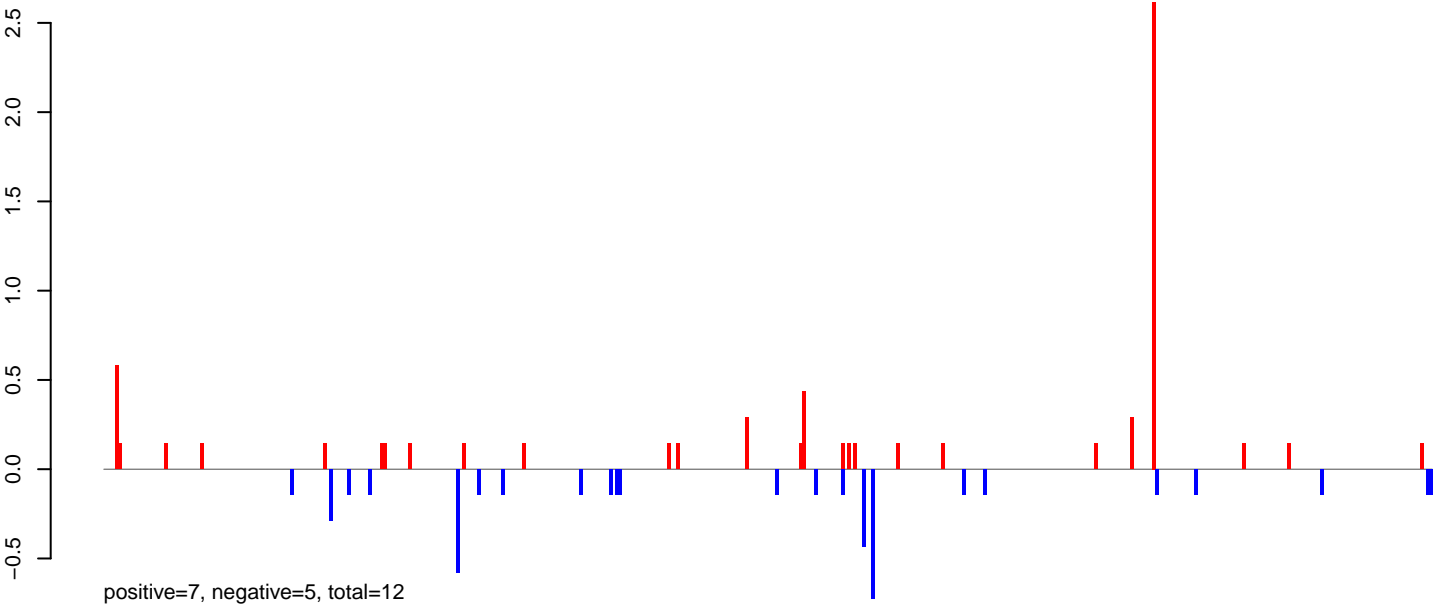
AeAeg_LVP_pMos3DBhspPBac.rep



w size=25, length=257, -12, score=1834448.1-Mosqueiro_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopro:1-12957

0 2000 4000 6000 8000 10000 12000

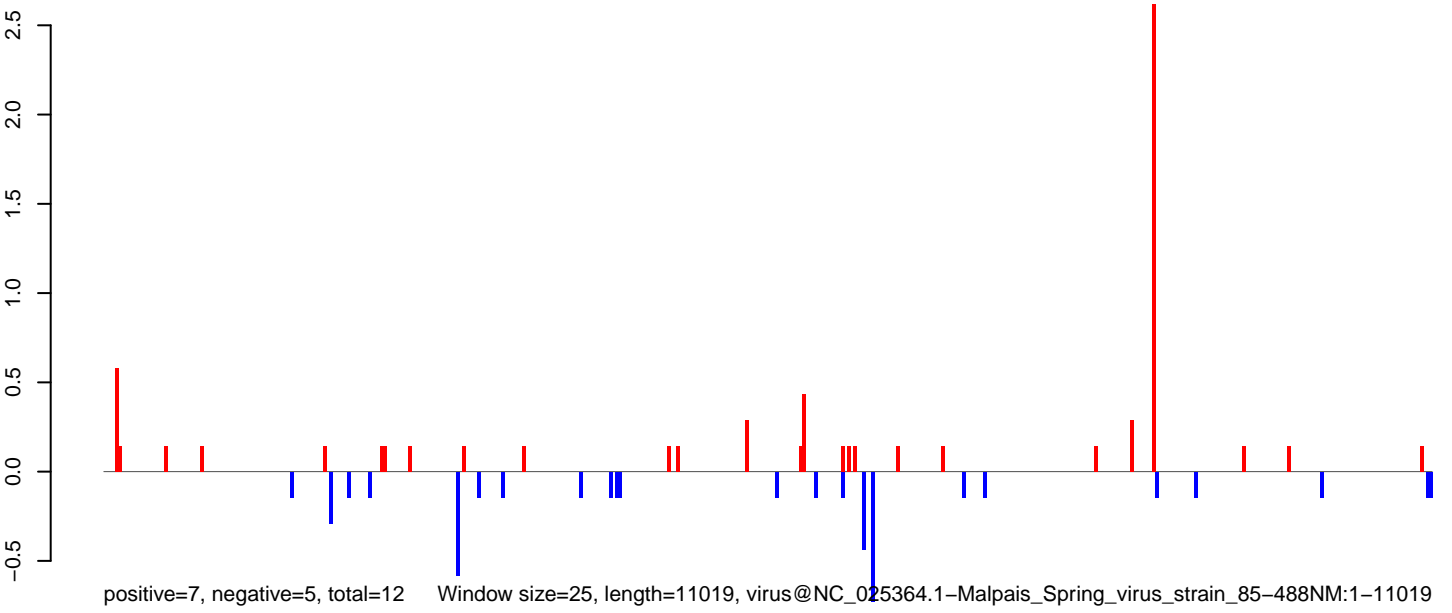
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



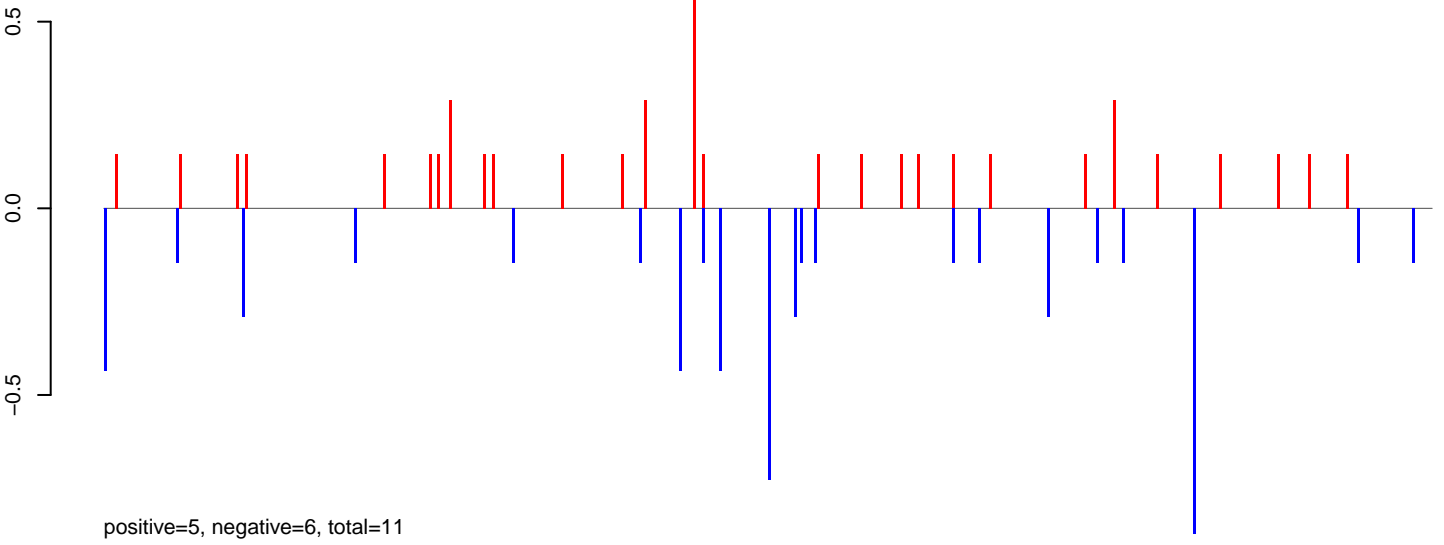
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



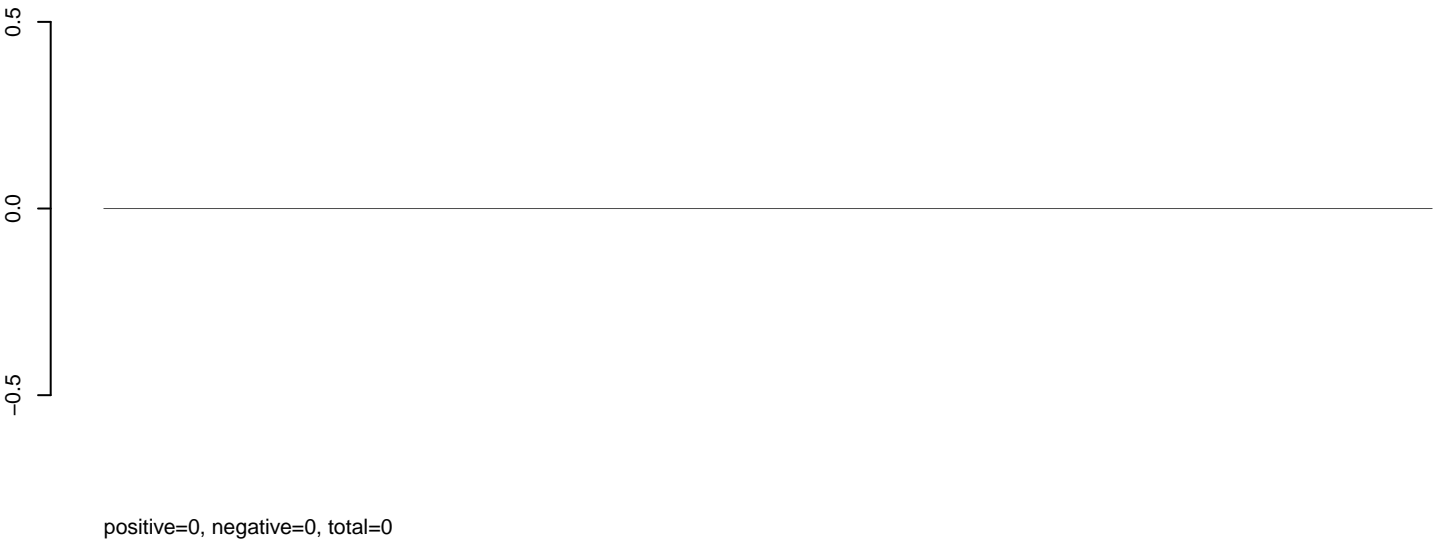
AeAeg_LVP_pMos3DBhspPBac.rep



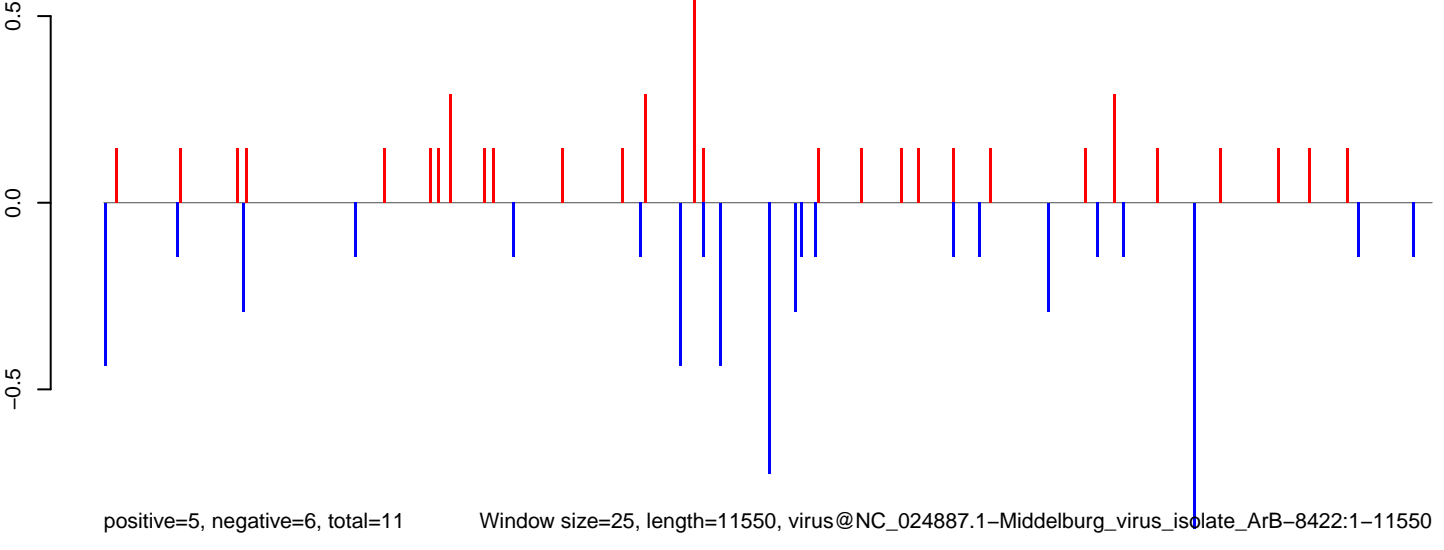
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

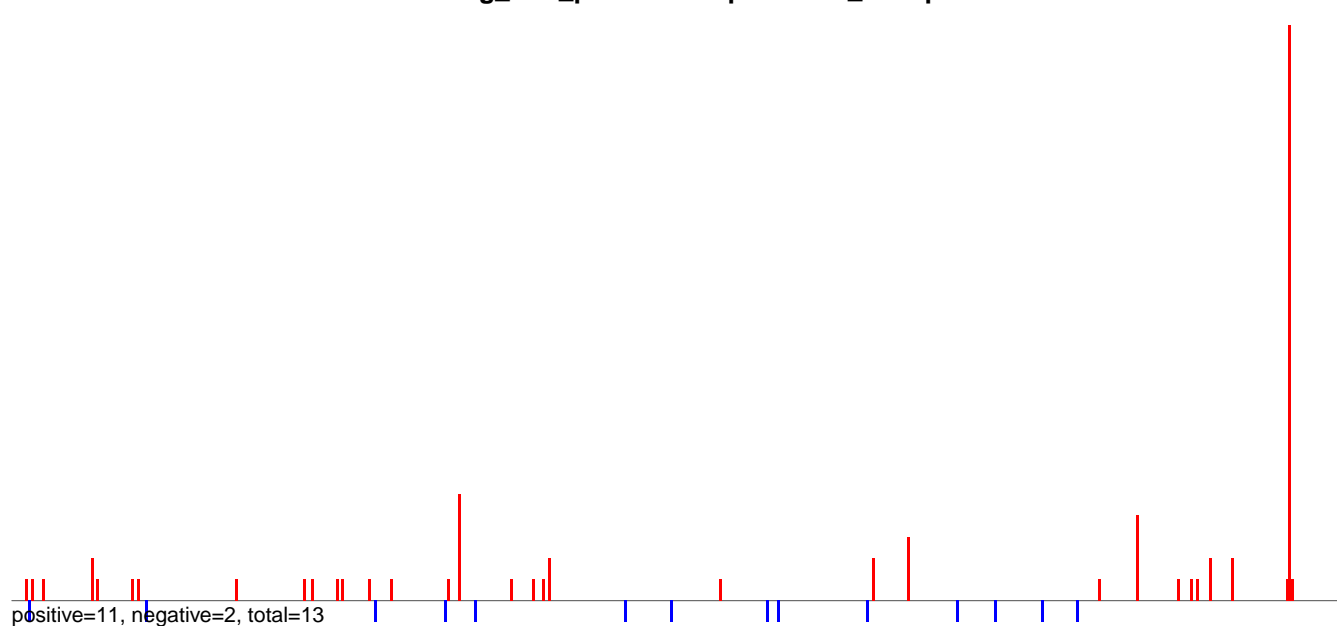


AeAeg_LVP_pMos3DBhspPBac.rep

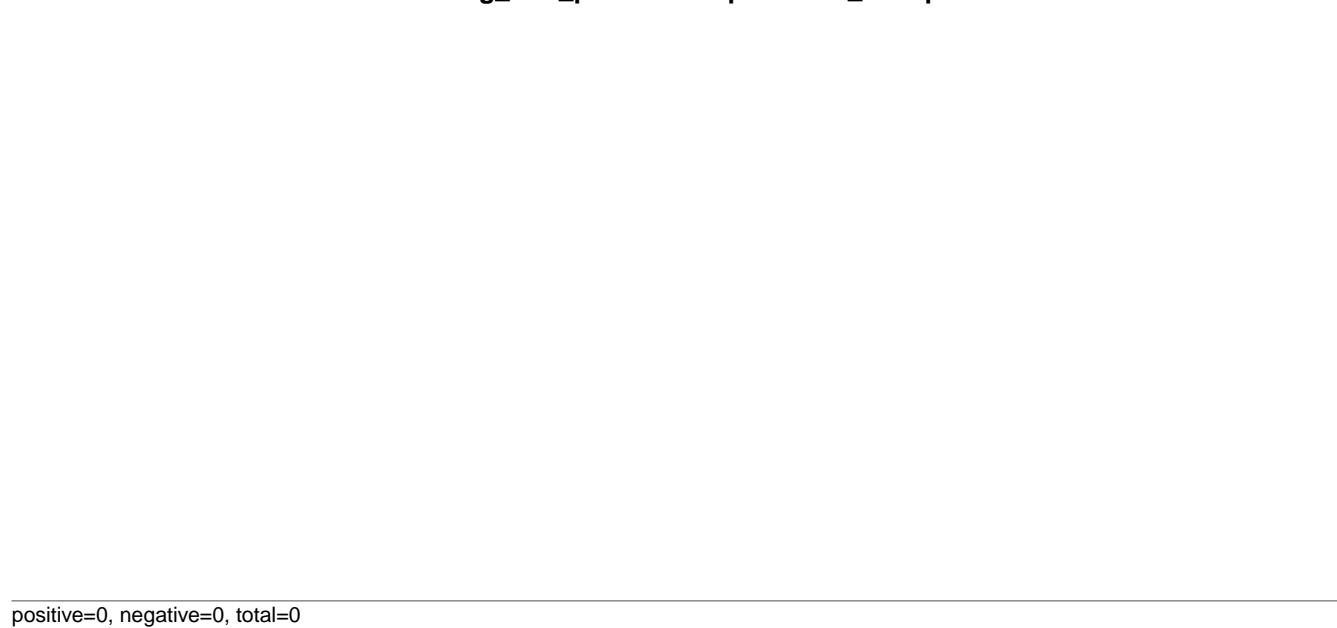


Window size=25, length=11550, virus@NC_024887.1-Middelburg_virus_isolate_ArB-8422:1-11550

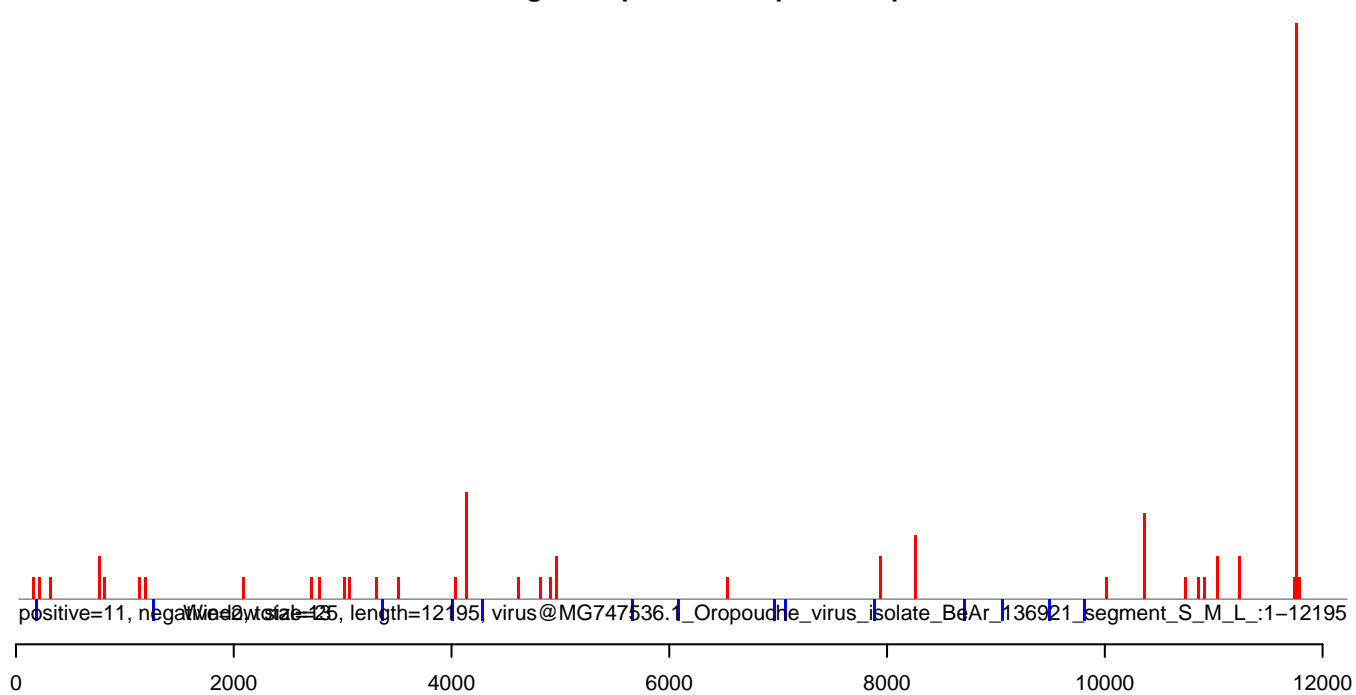
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



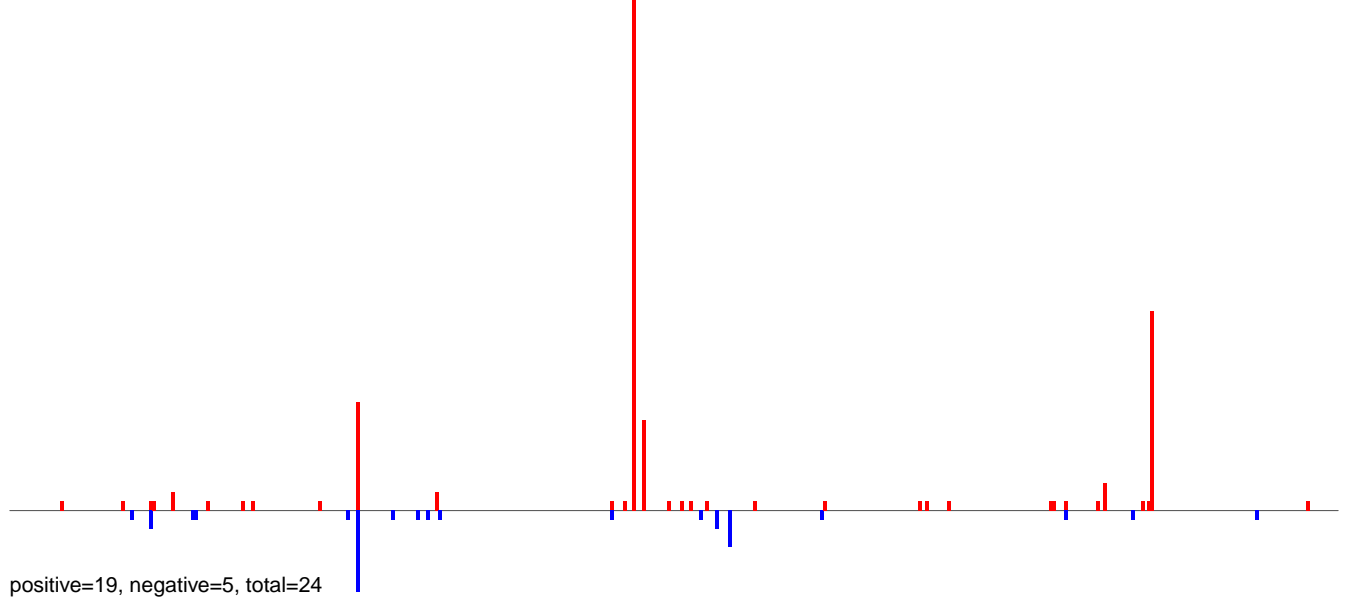
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



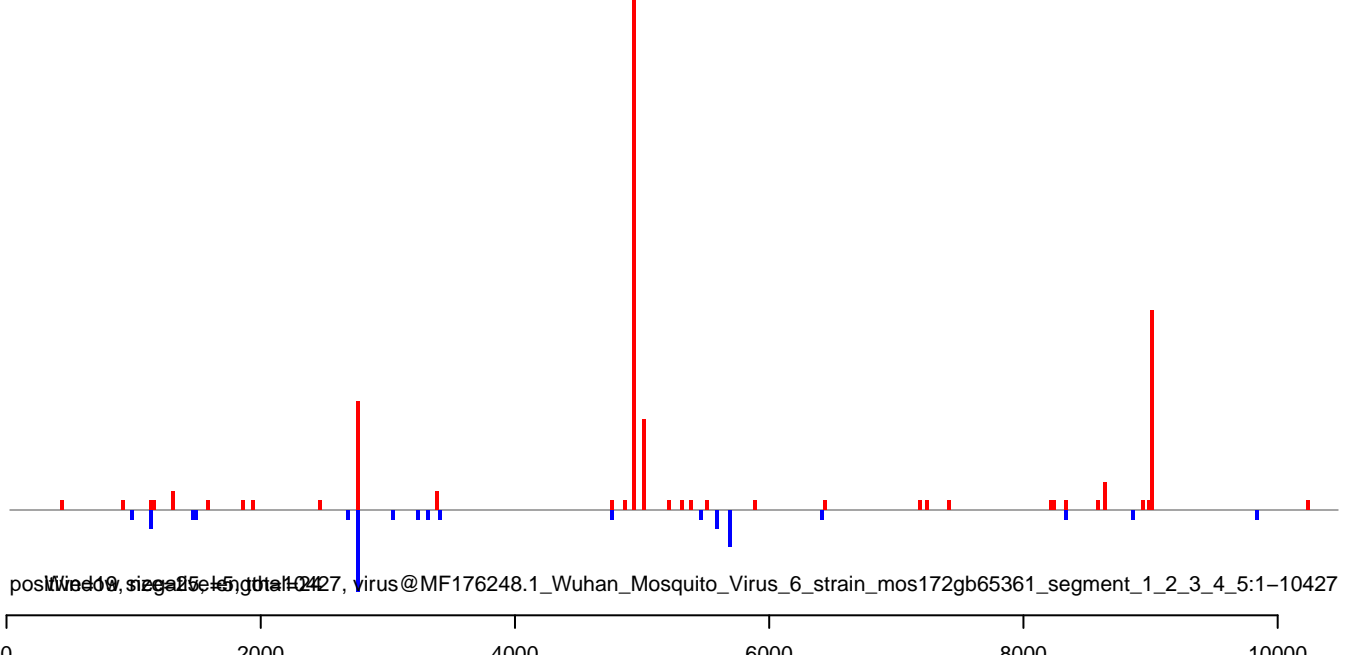
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



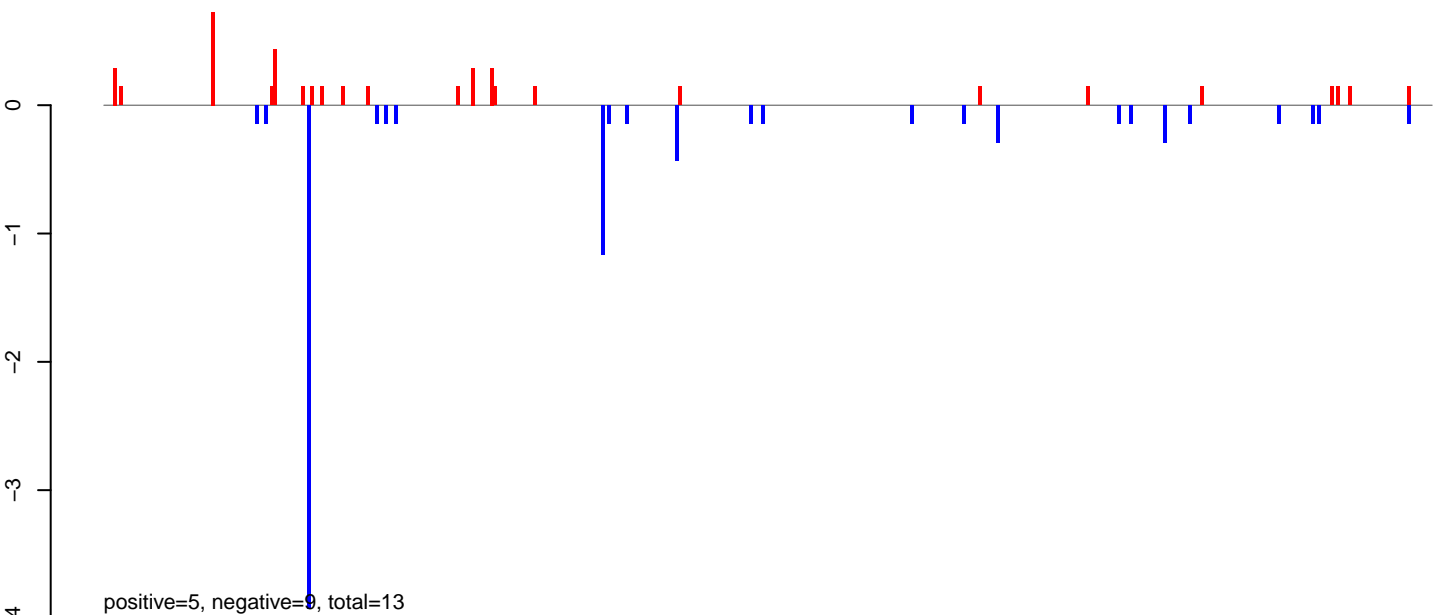
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



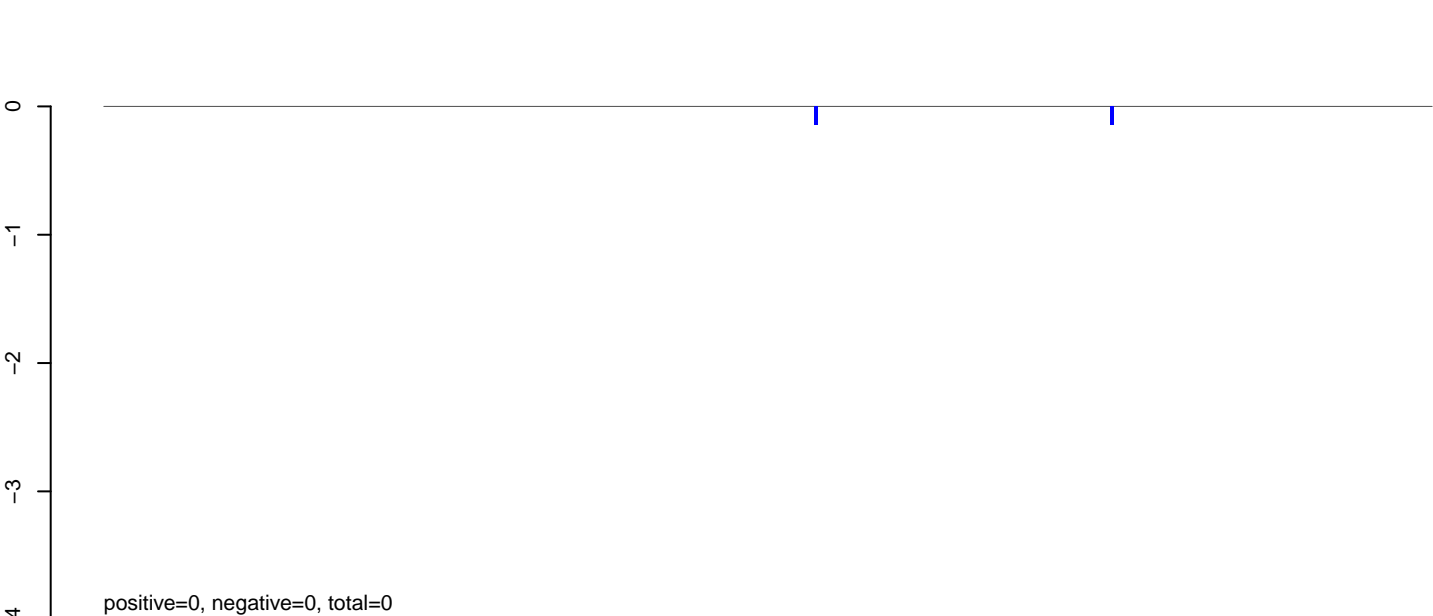
AeAeg_LVP_pMos3DBhspPBac.rep



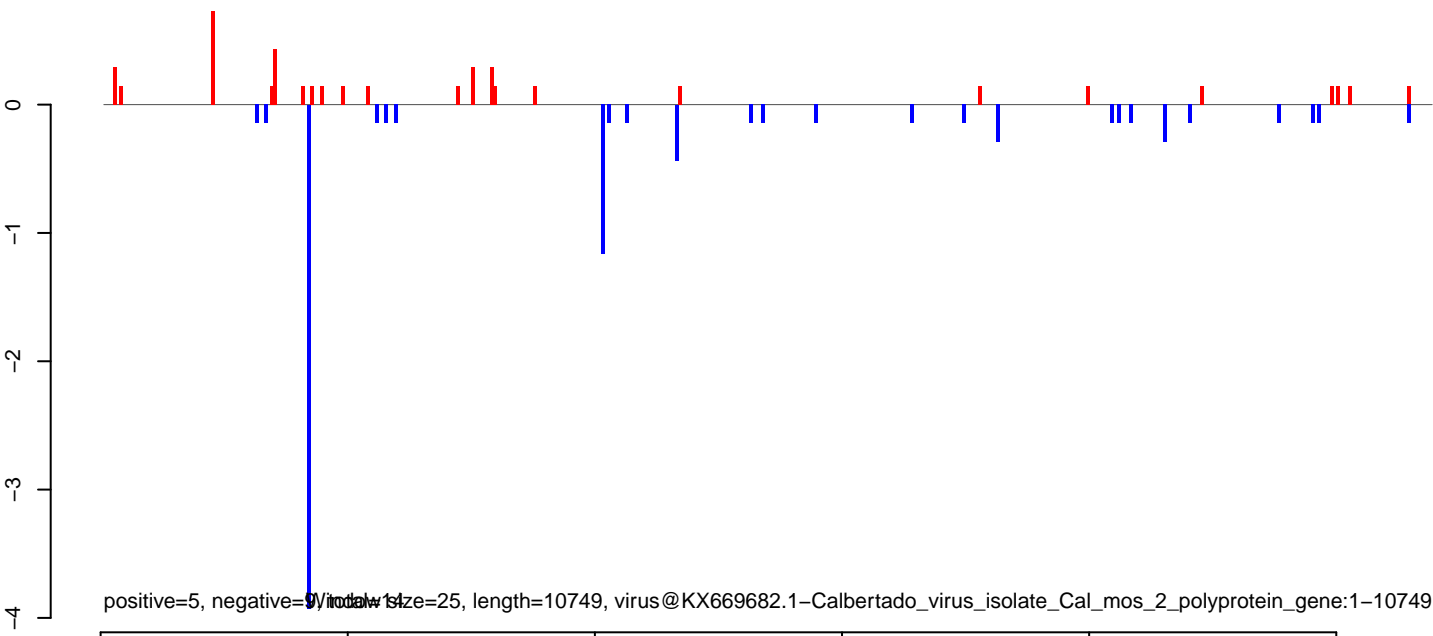
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



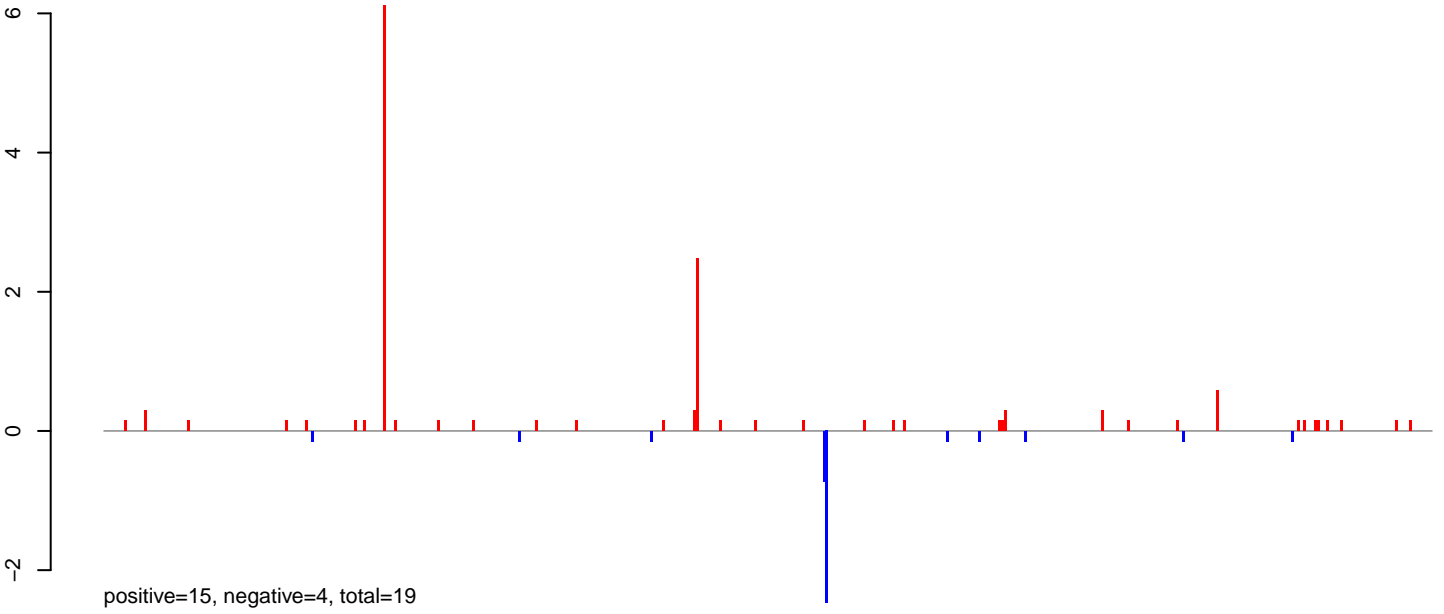
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



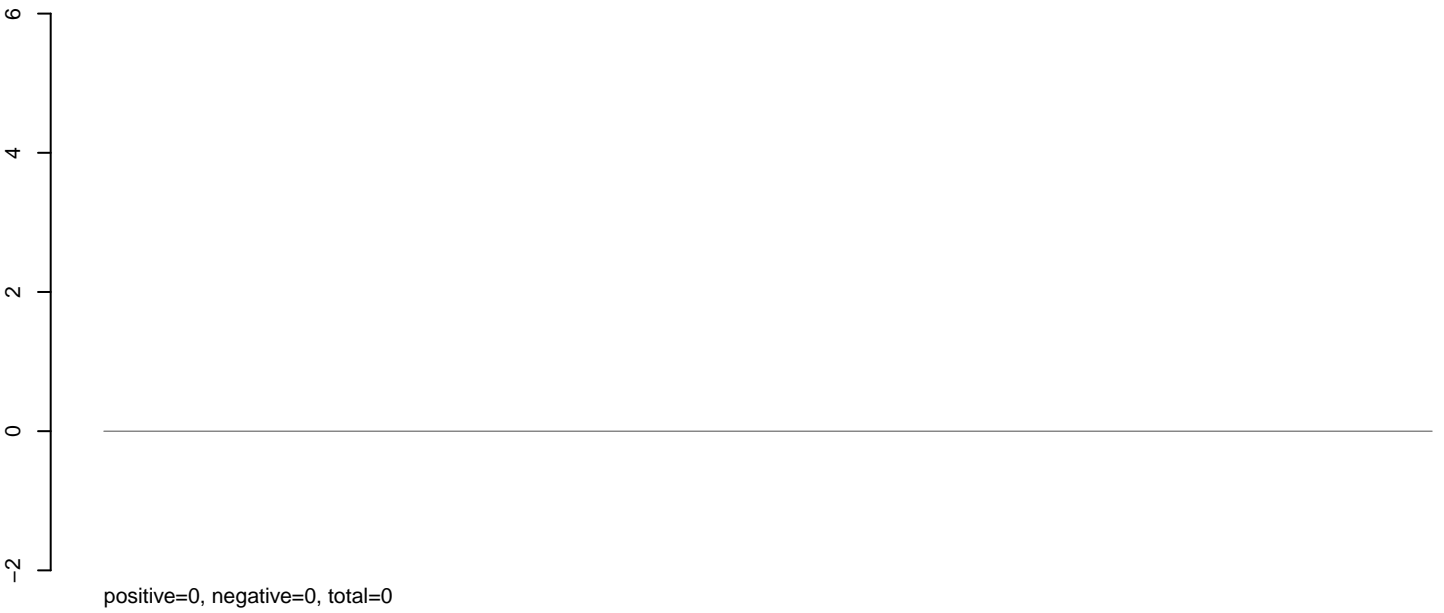
AeAeg_LVP_pMos3DBhspPBac.rep



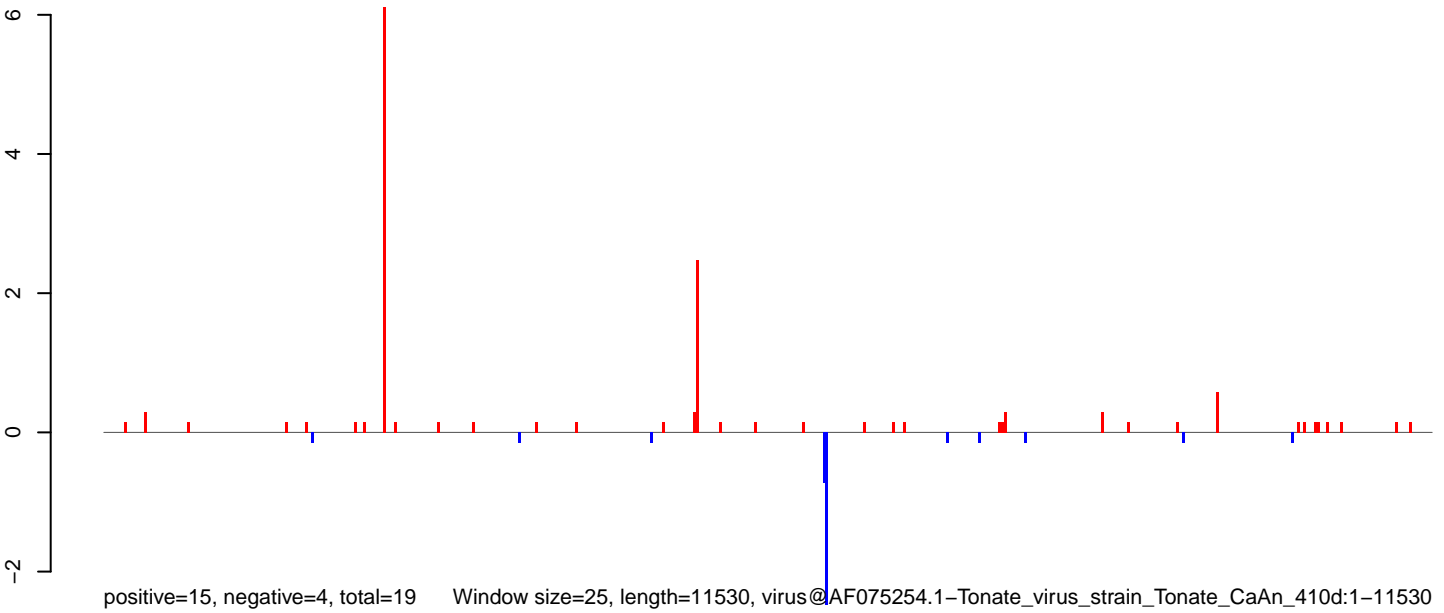
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



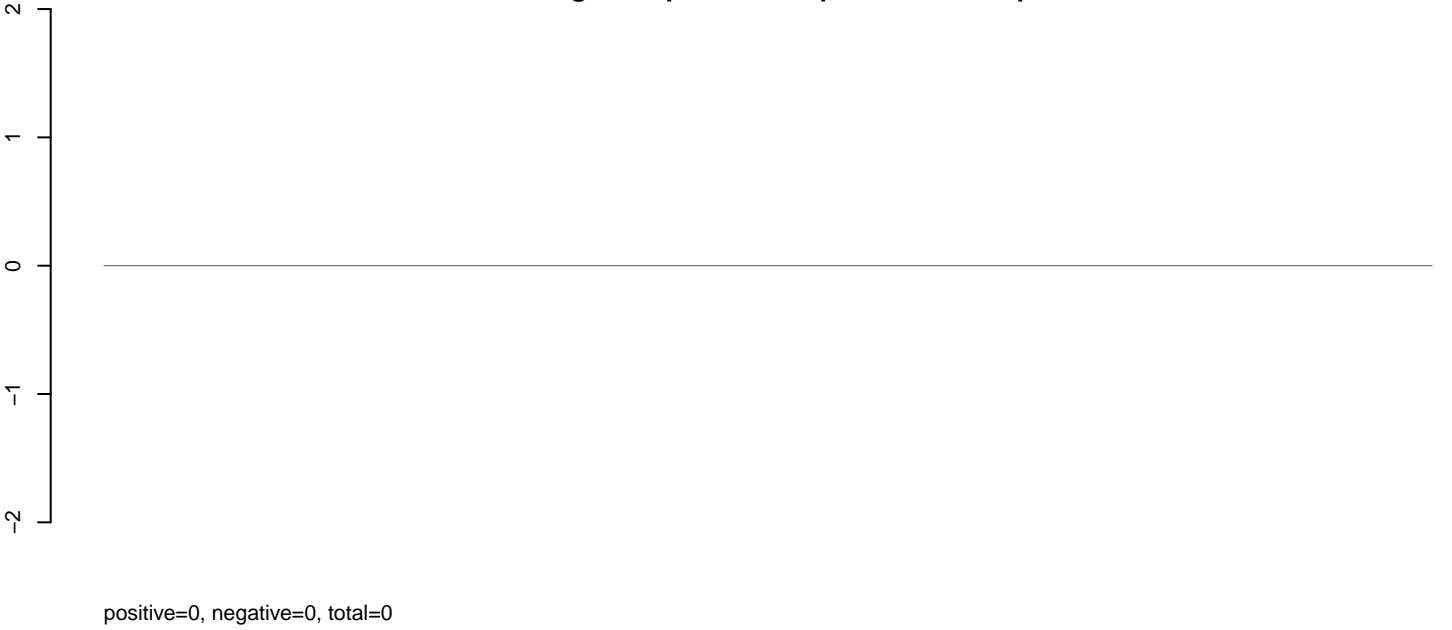
Window size=25, length=11530, virus@AF075254.1-Tonate_virus_strain_Tonate_CaAn_410d:1-11530

0 2000 4000 6000 8000 10000 12000

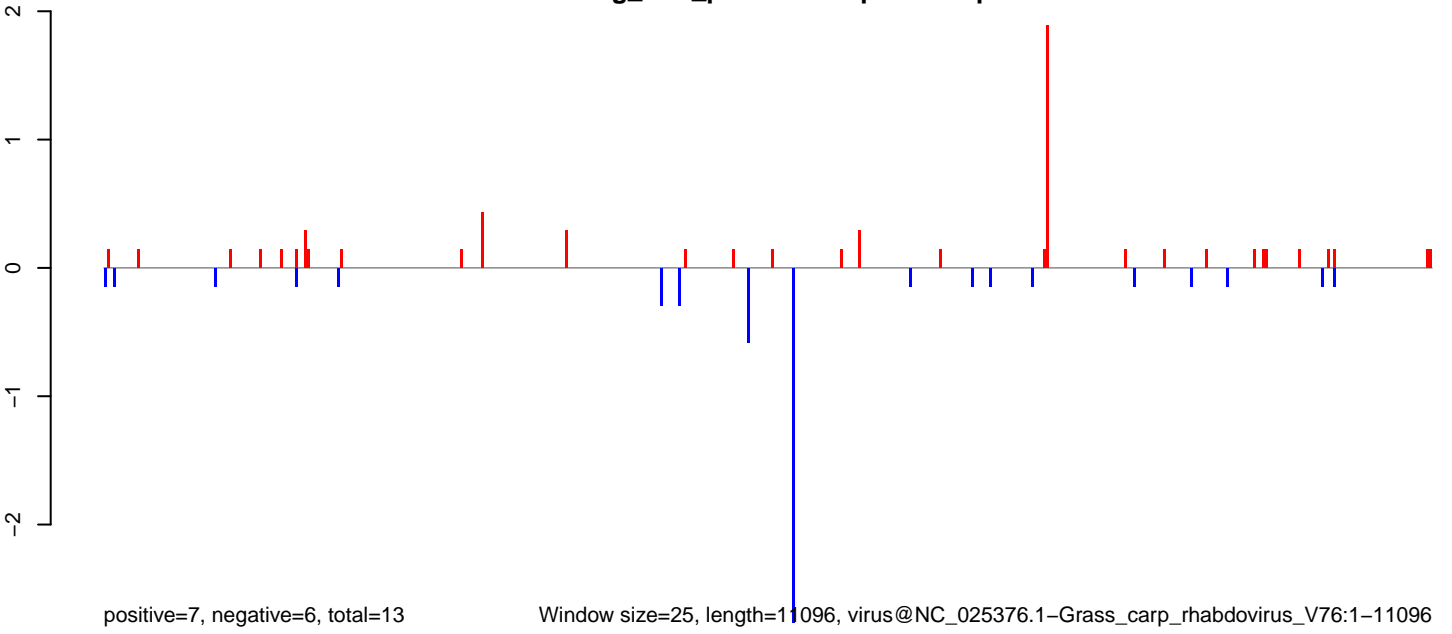
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

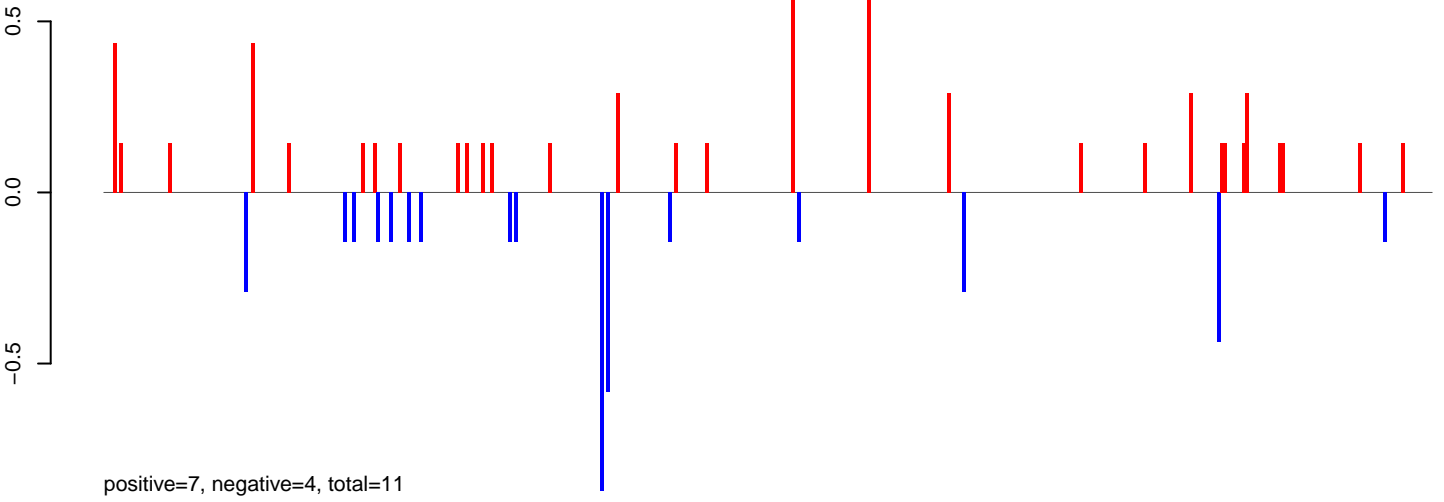


AeAeg_LVP_pMos3DBhspPBac.rep

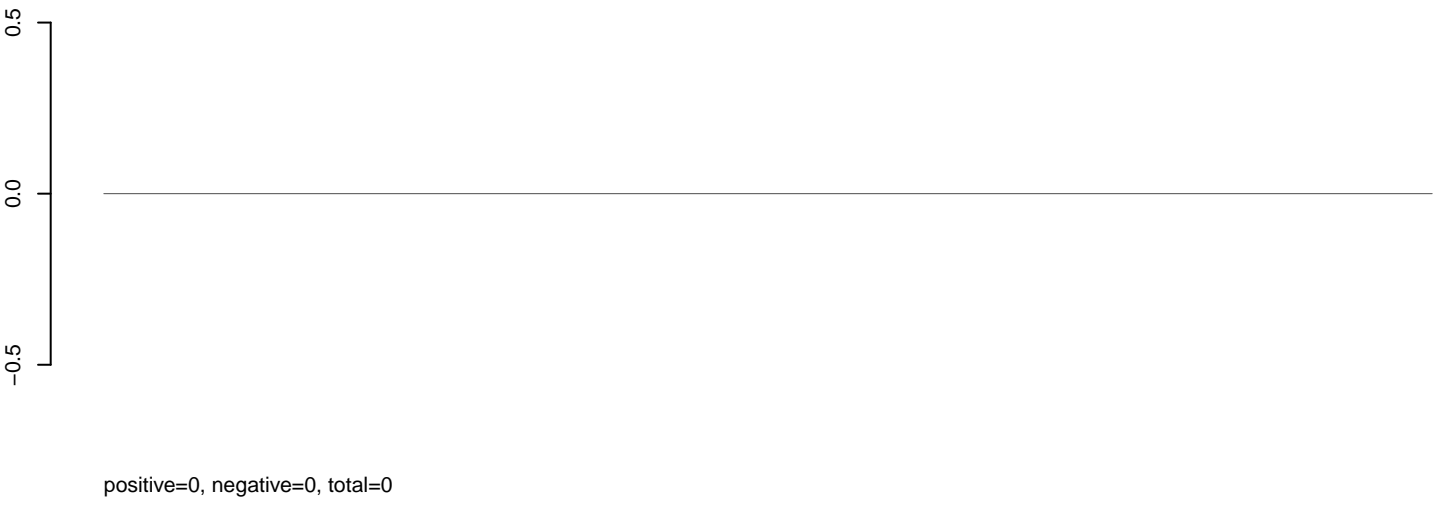


0 2000 4000 6000 8000 10000

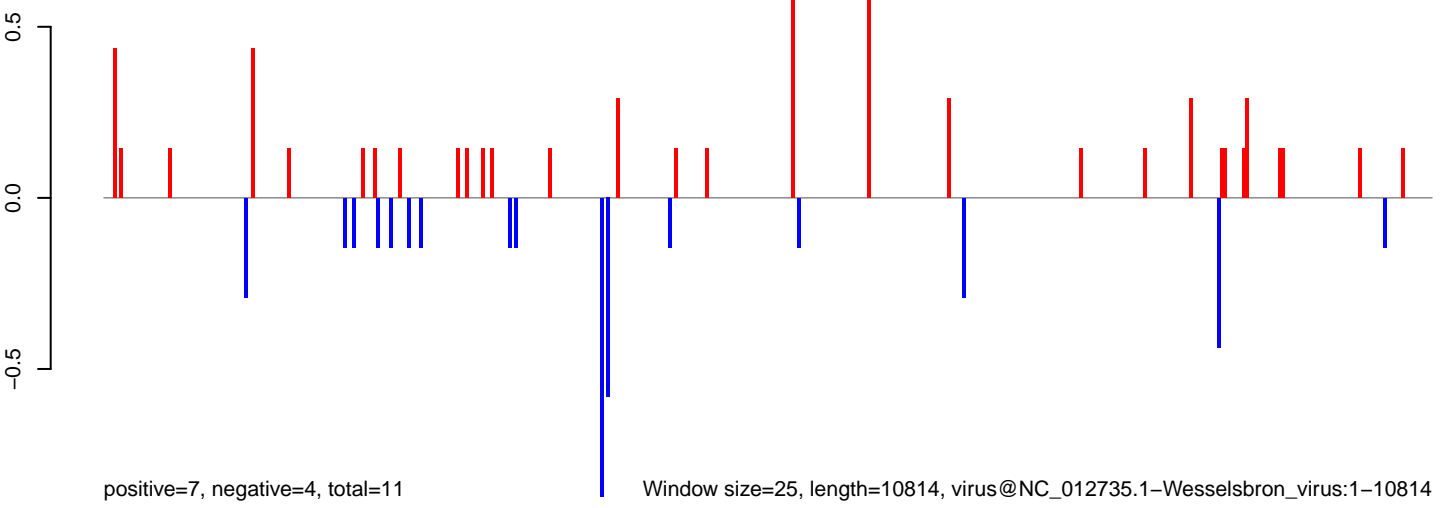
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

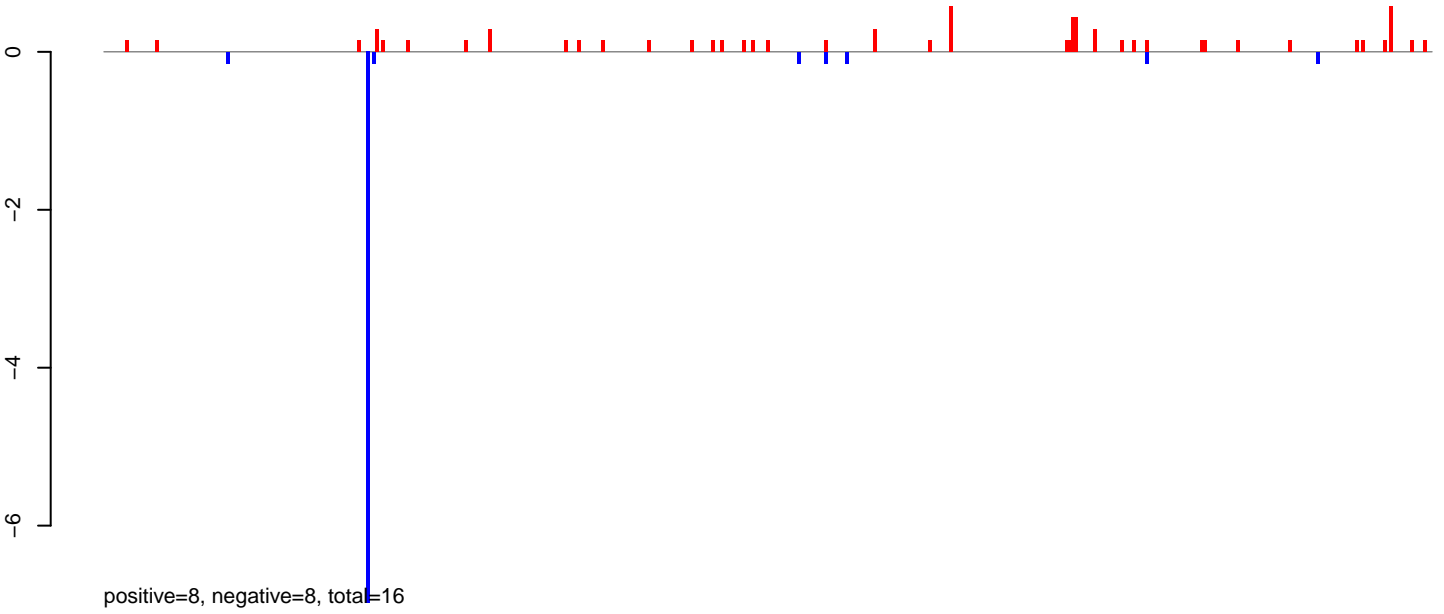


AeAeg_LVP_pMos3DBhspPBac.rep

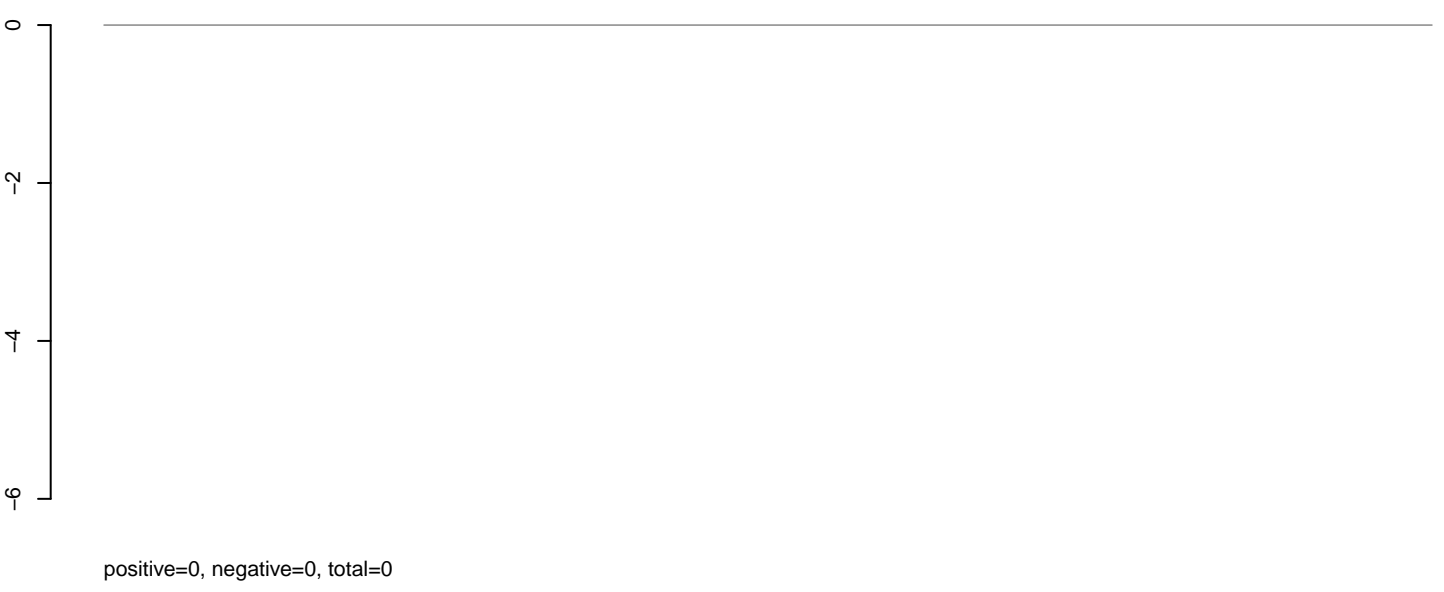


0 2000 4000 6000 8000 10000

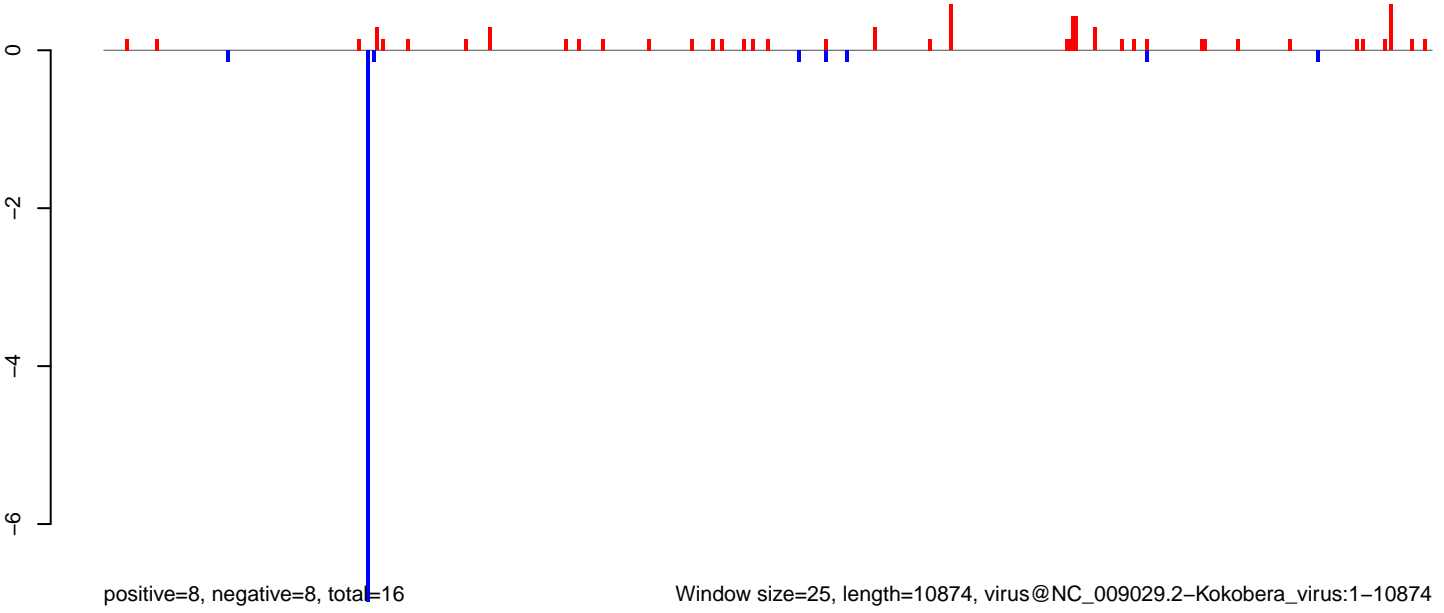
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



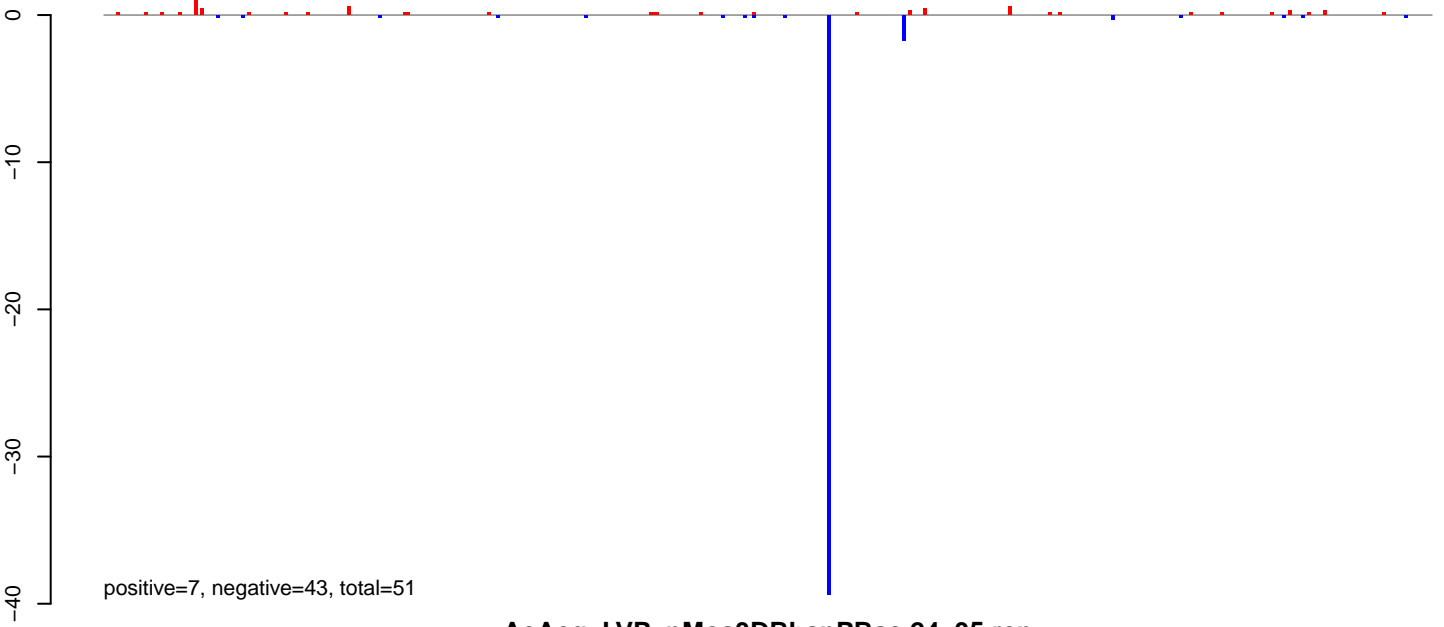
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



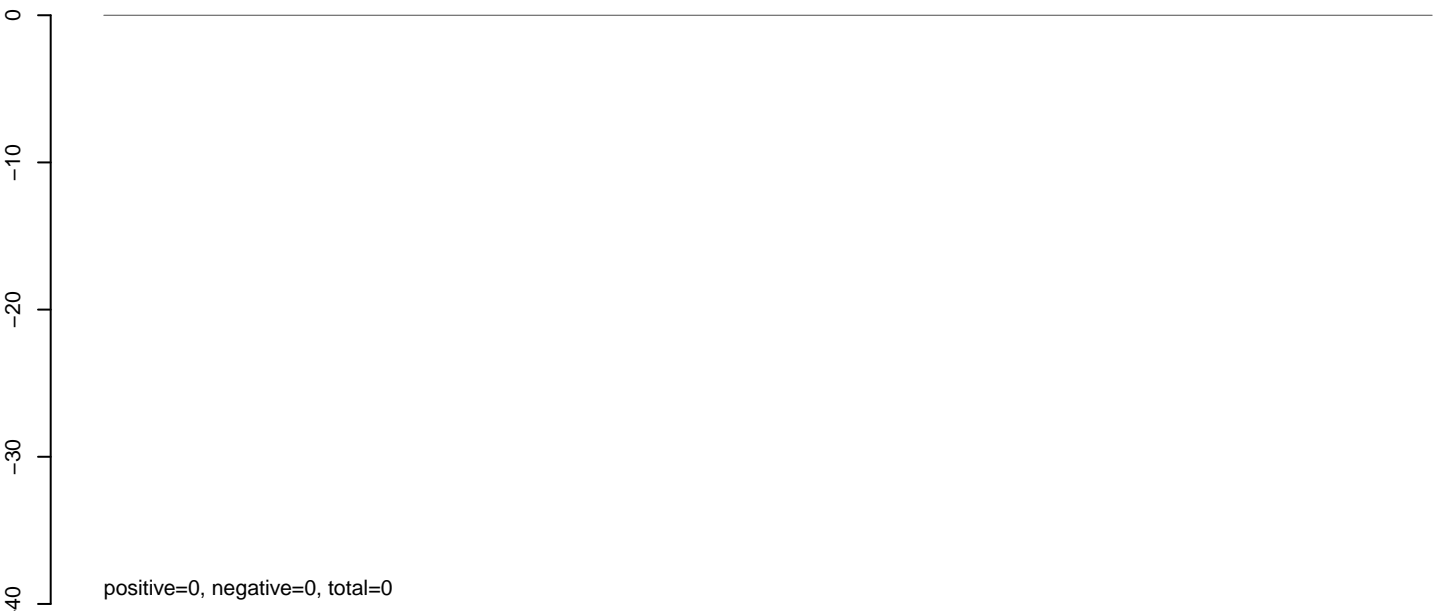
AeAeg_LVP_pMos3DBhspPBac.rep



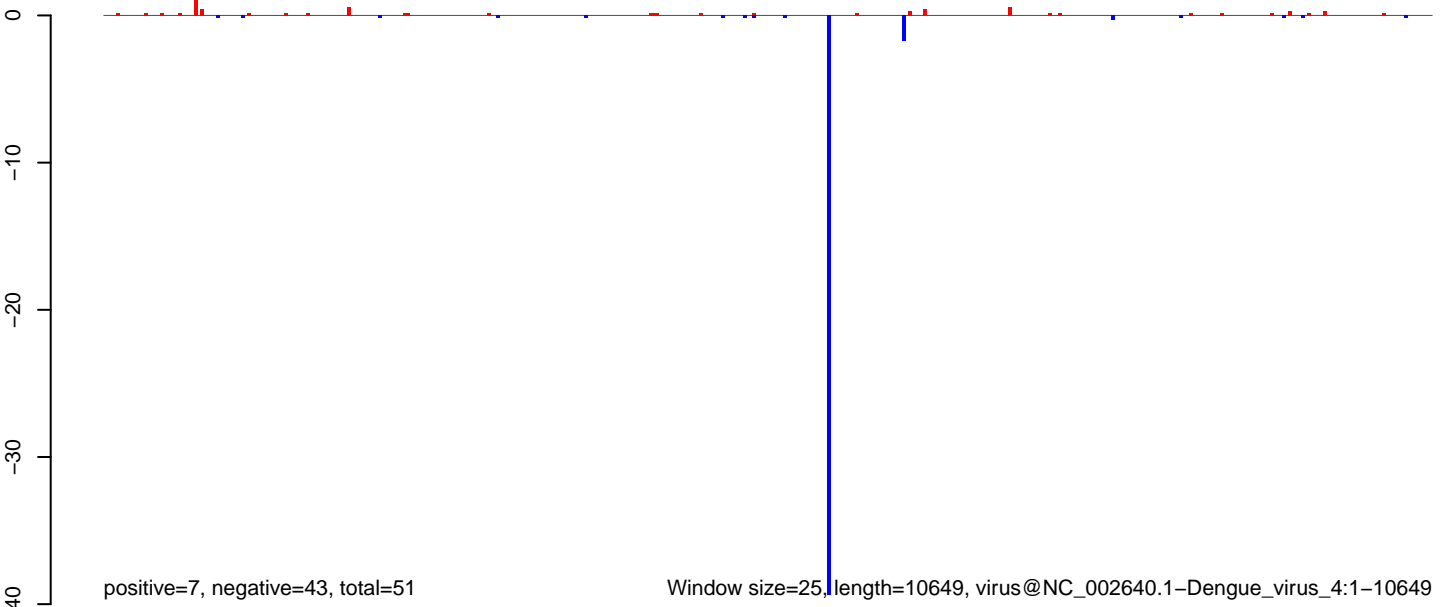
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



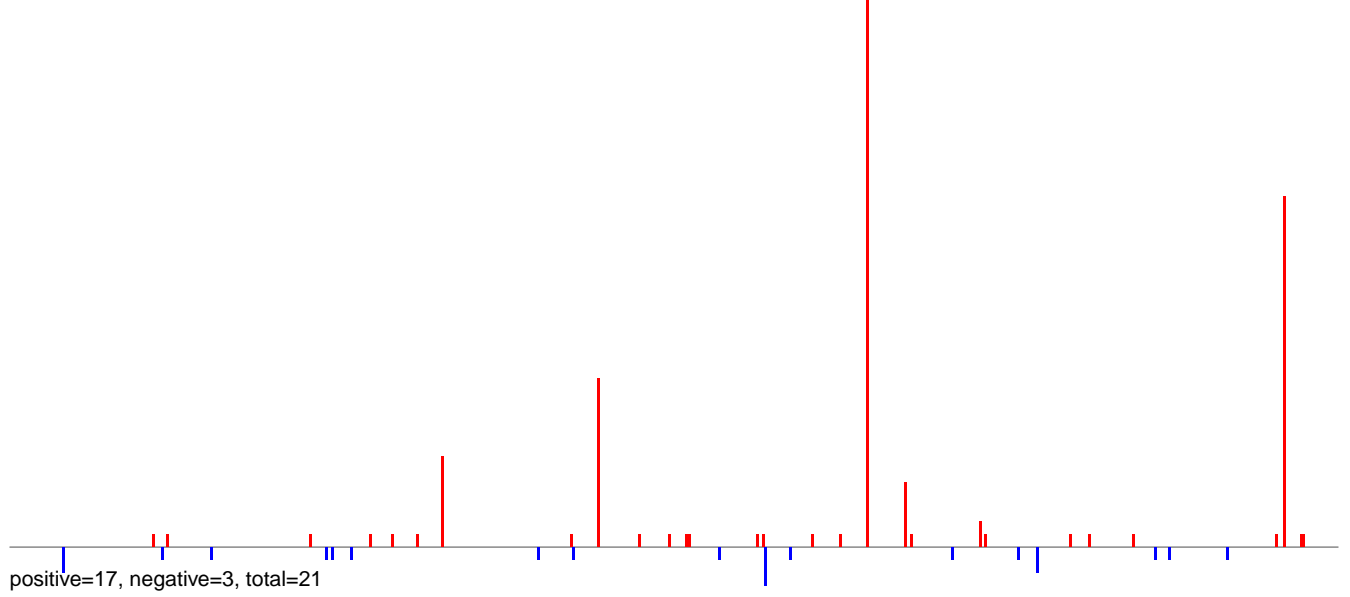
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



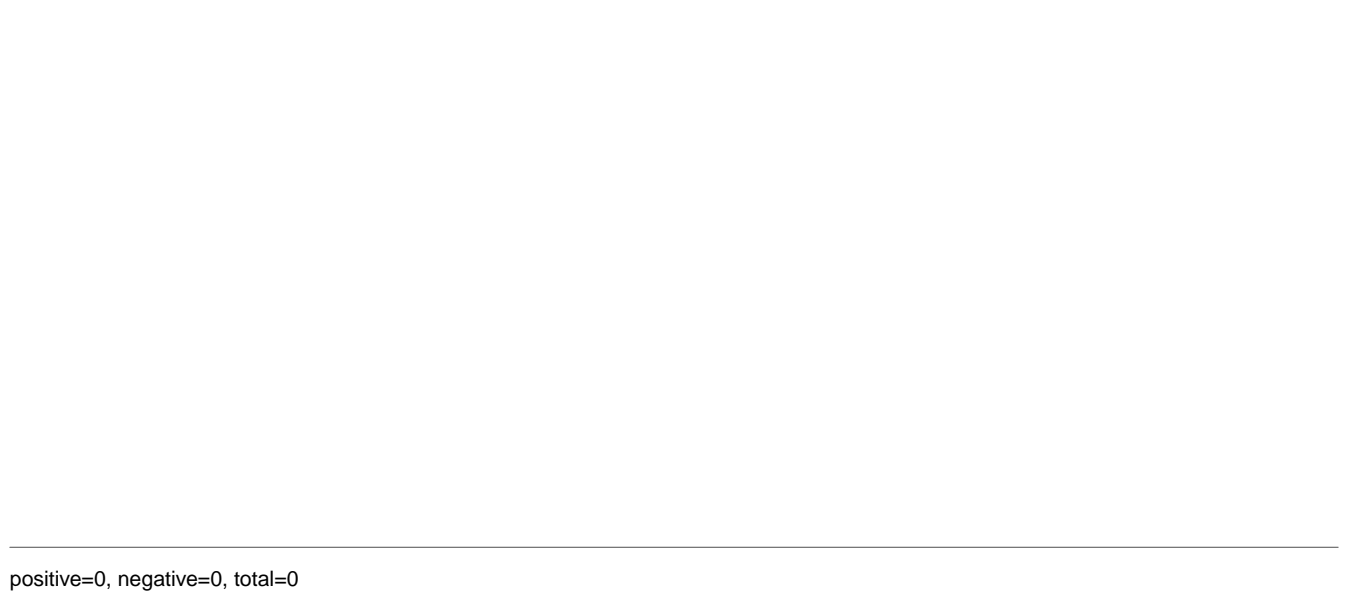
AeAeg_LVP_pMos3DBhspPBac.rep



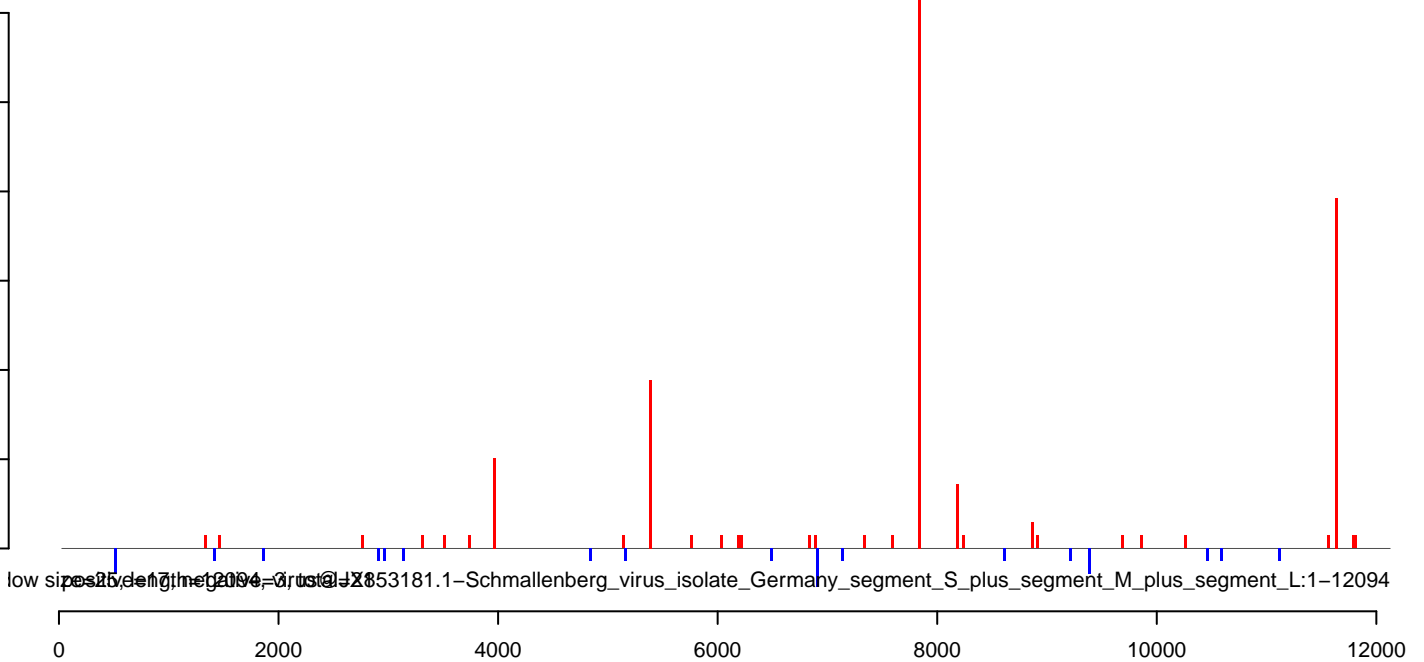
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



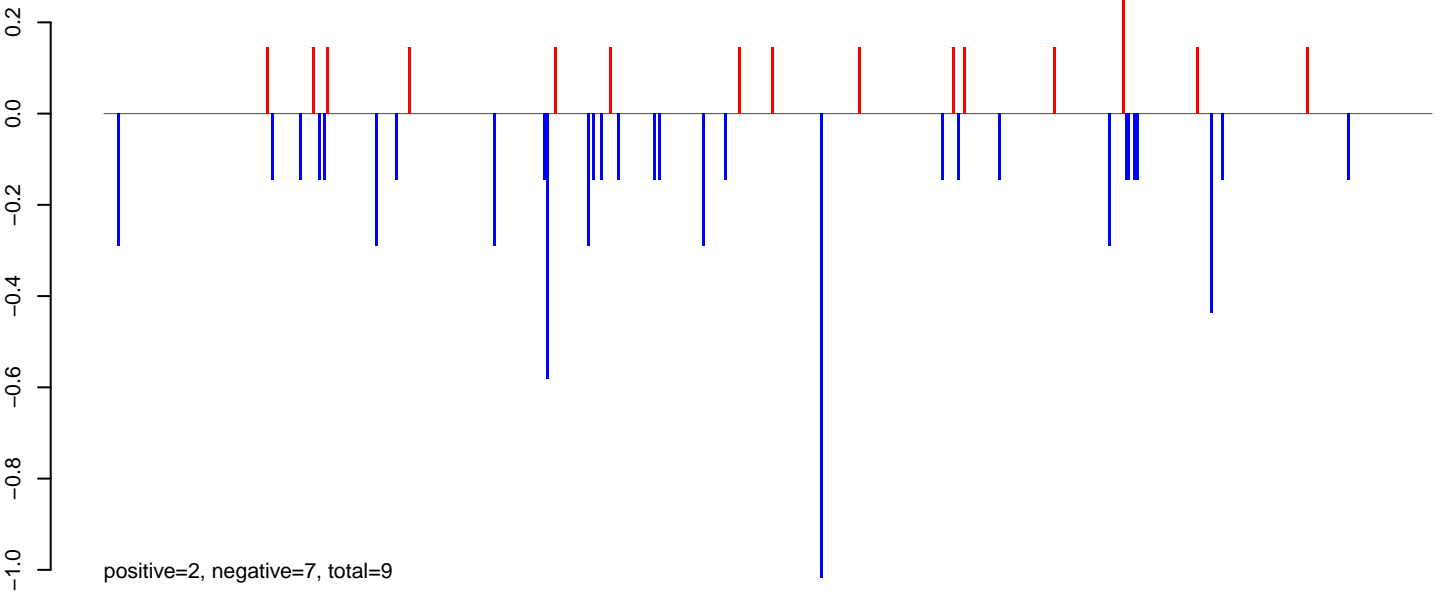
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



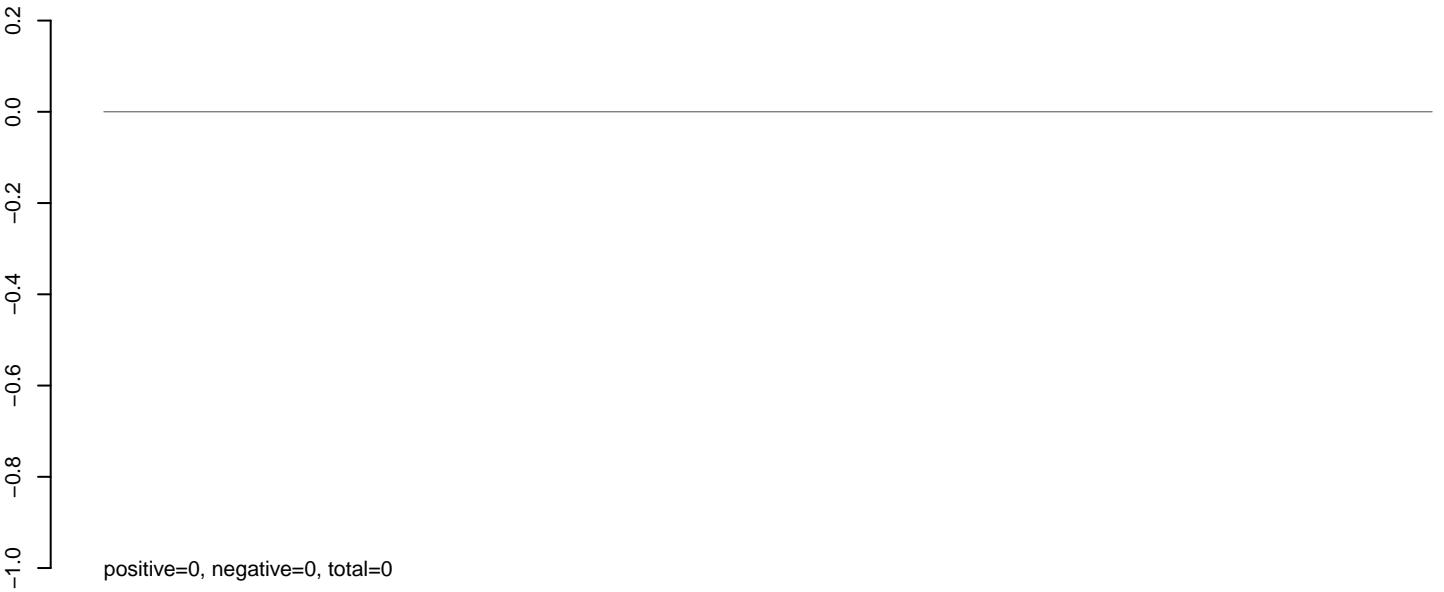
AeAeg_LVP_pMos3DBhspPBac.rep



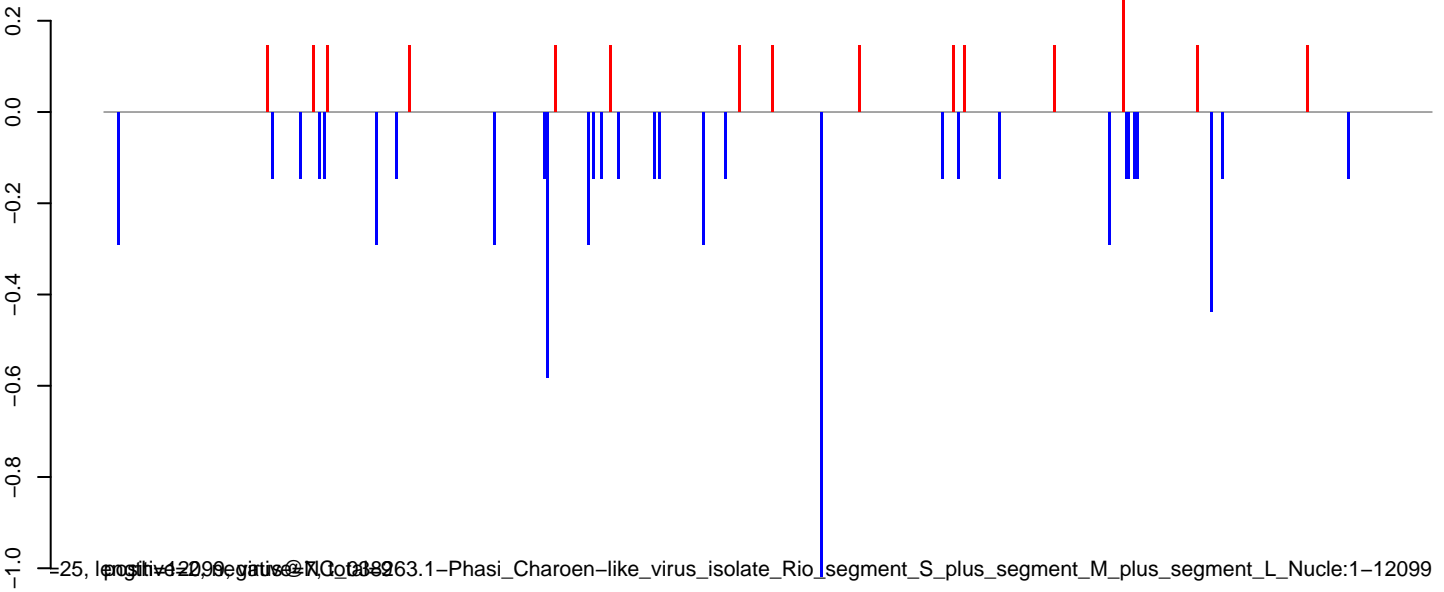
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



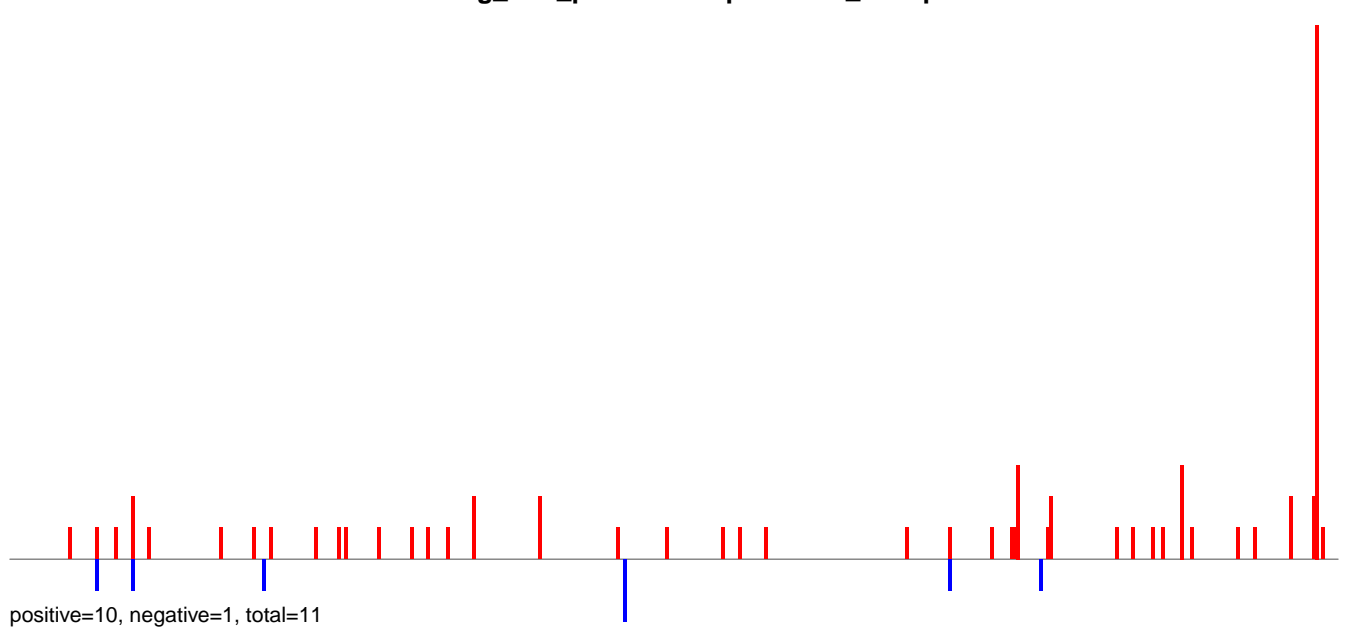
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



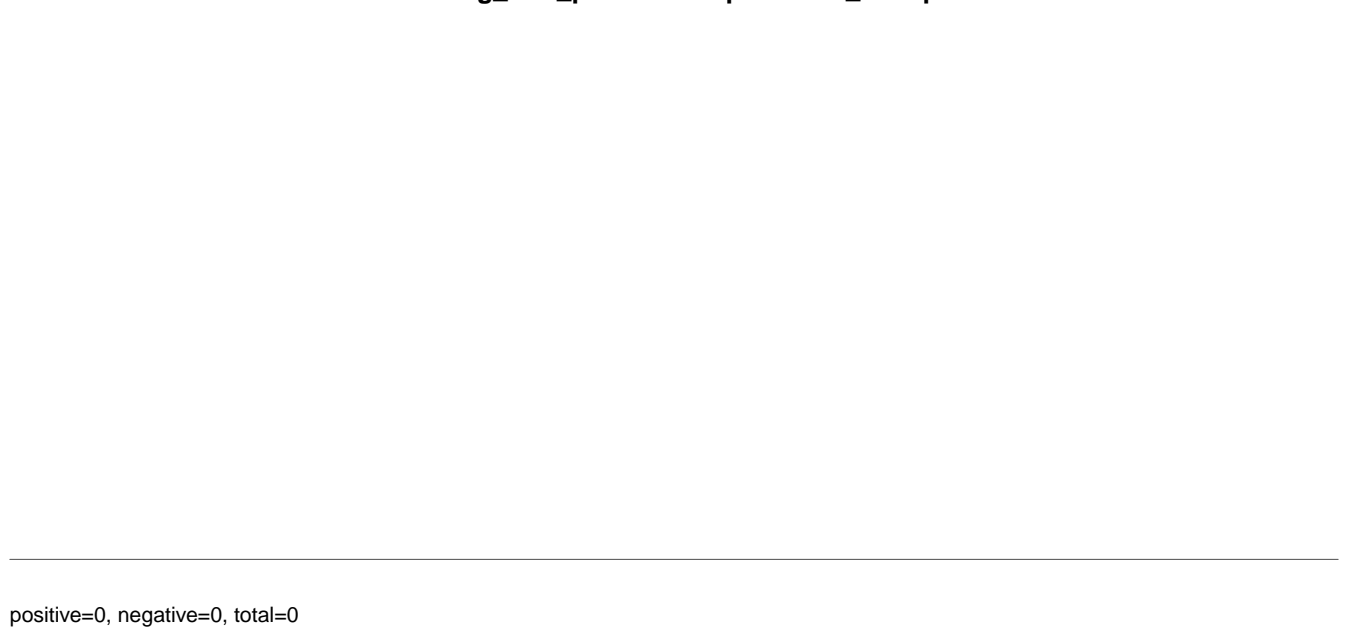
AeAeg_LVP_pMos3DBhspPBac.rep



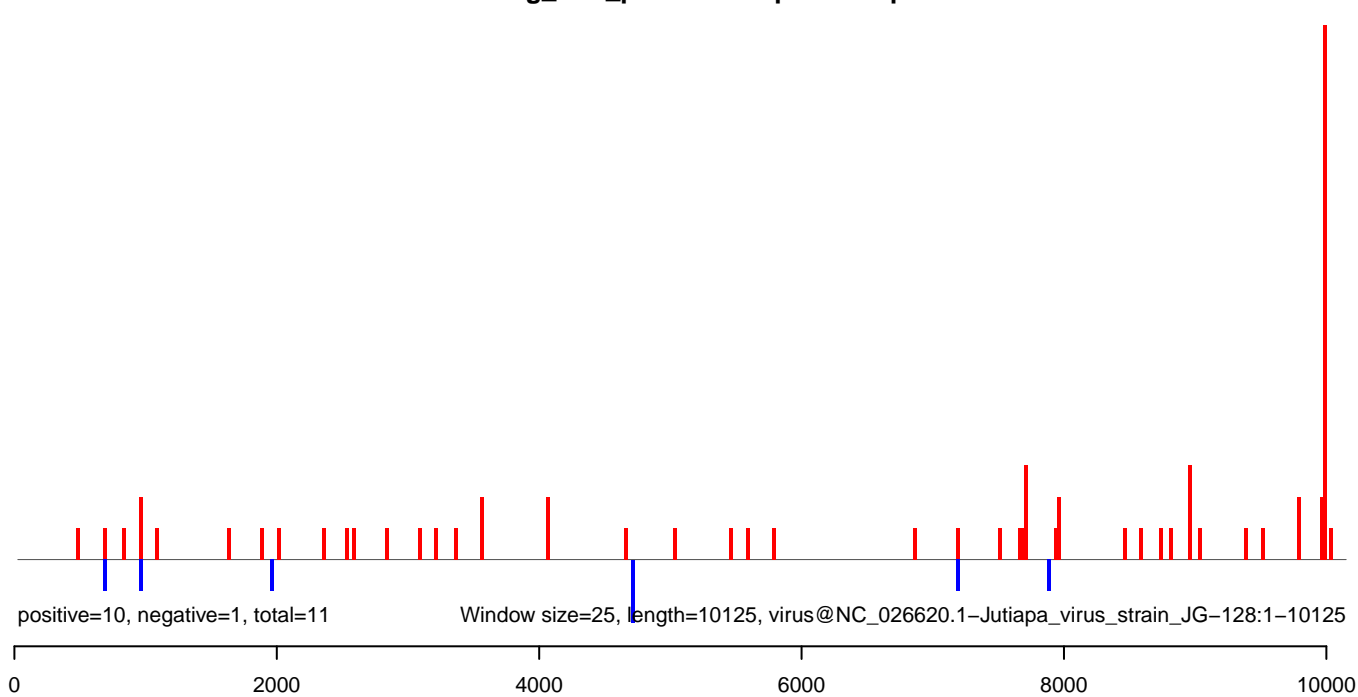
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



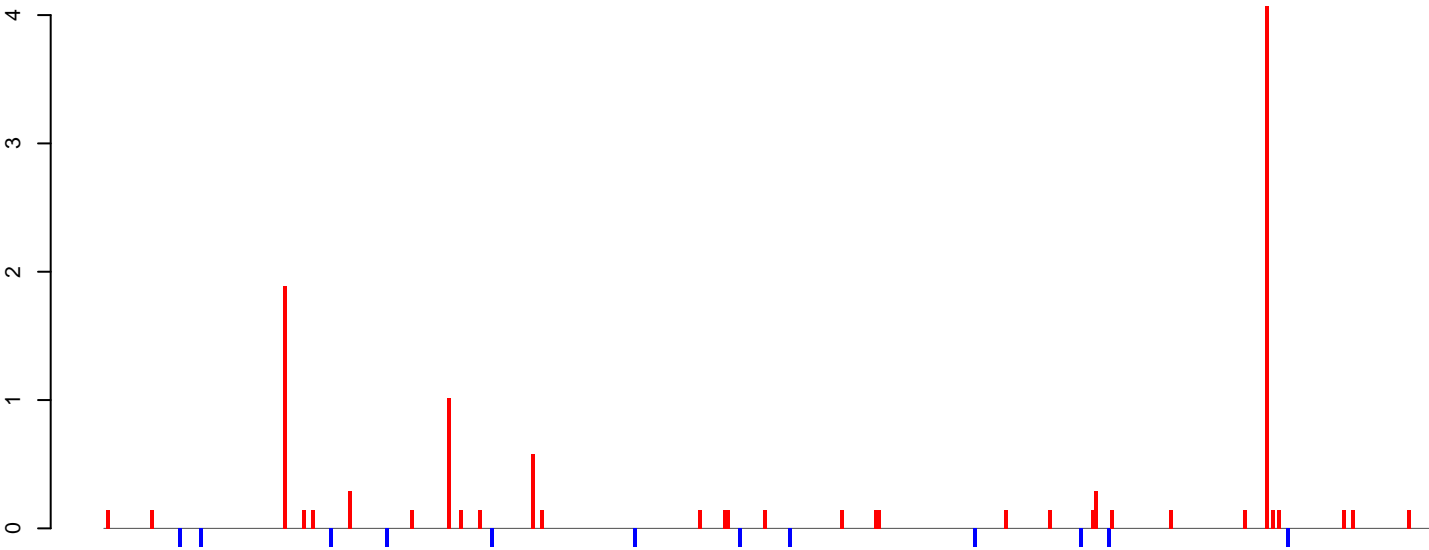
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep

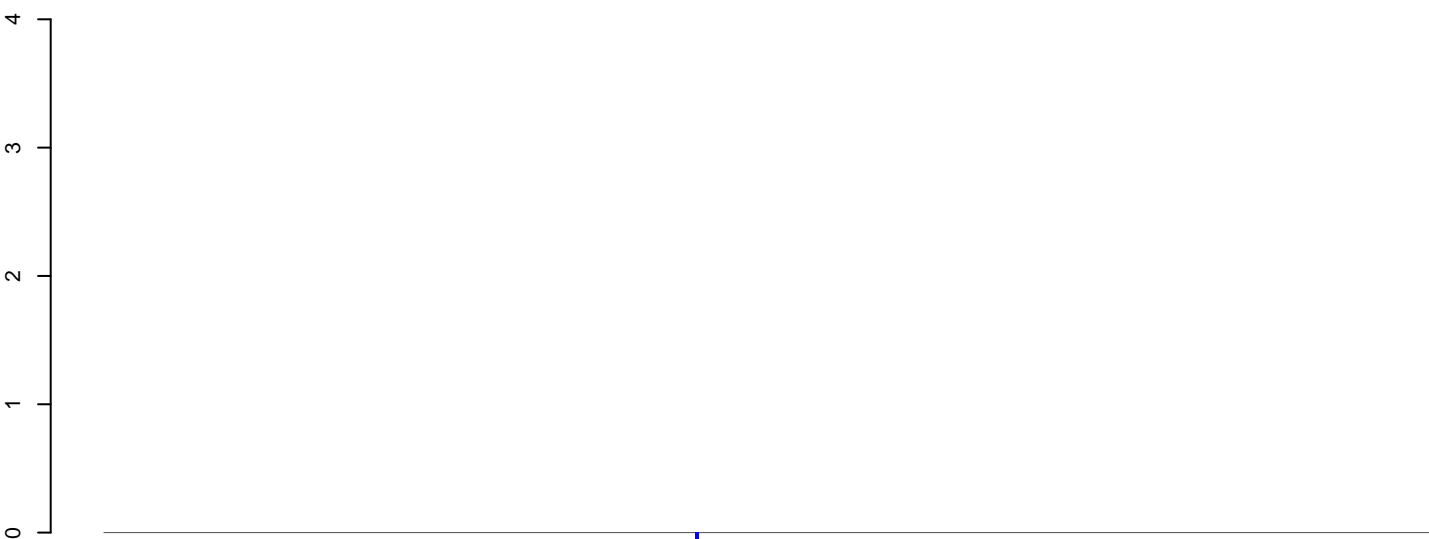


AeAeg_LVP_pMos3DBhspPBac.18_23.rep



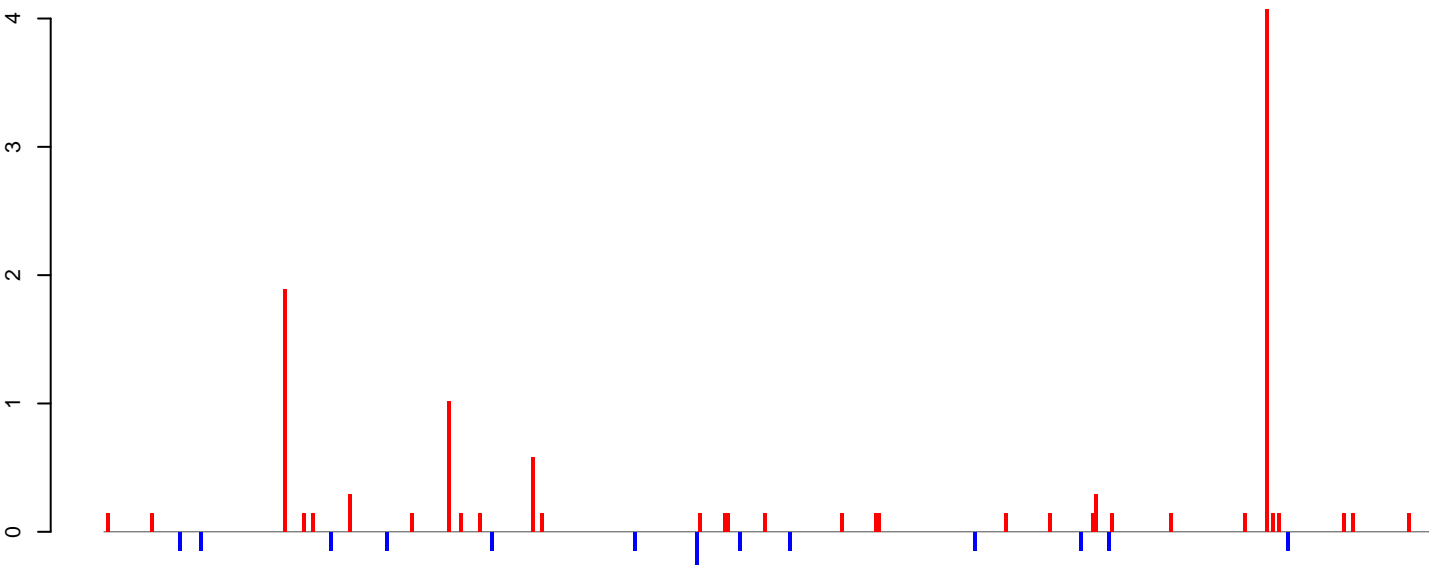
positive=12, negative=2, total=14

AeAeg_LVP_pMos3DBhspPBac.24_35.rep



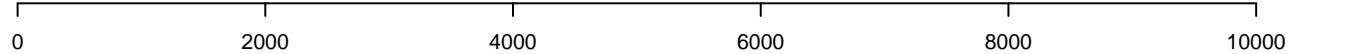
positive=0, negative=1, total=1

AeAeg_LVP_pMos3DBhspPBac.rep

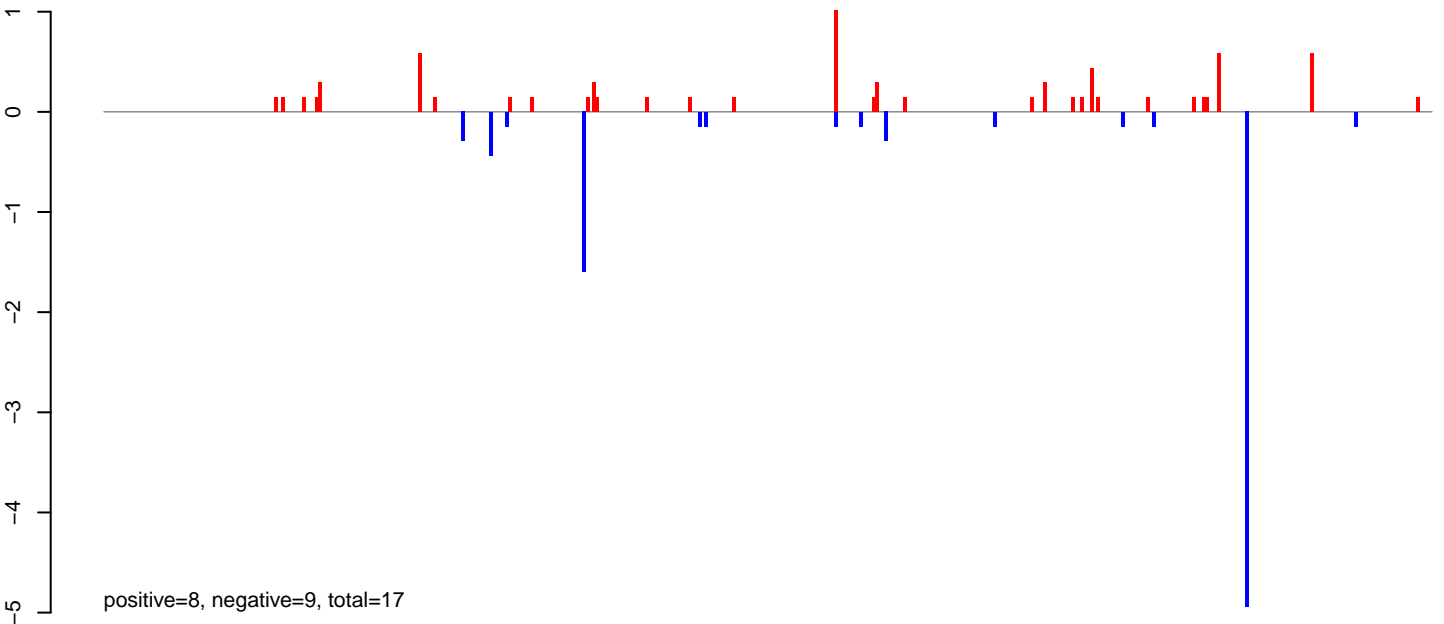


positive=12, negative=2, total=14

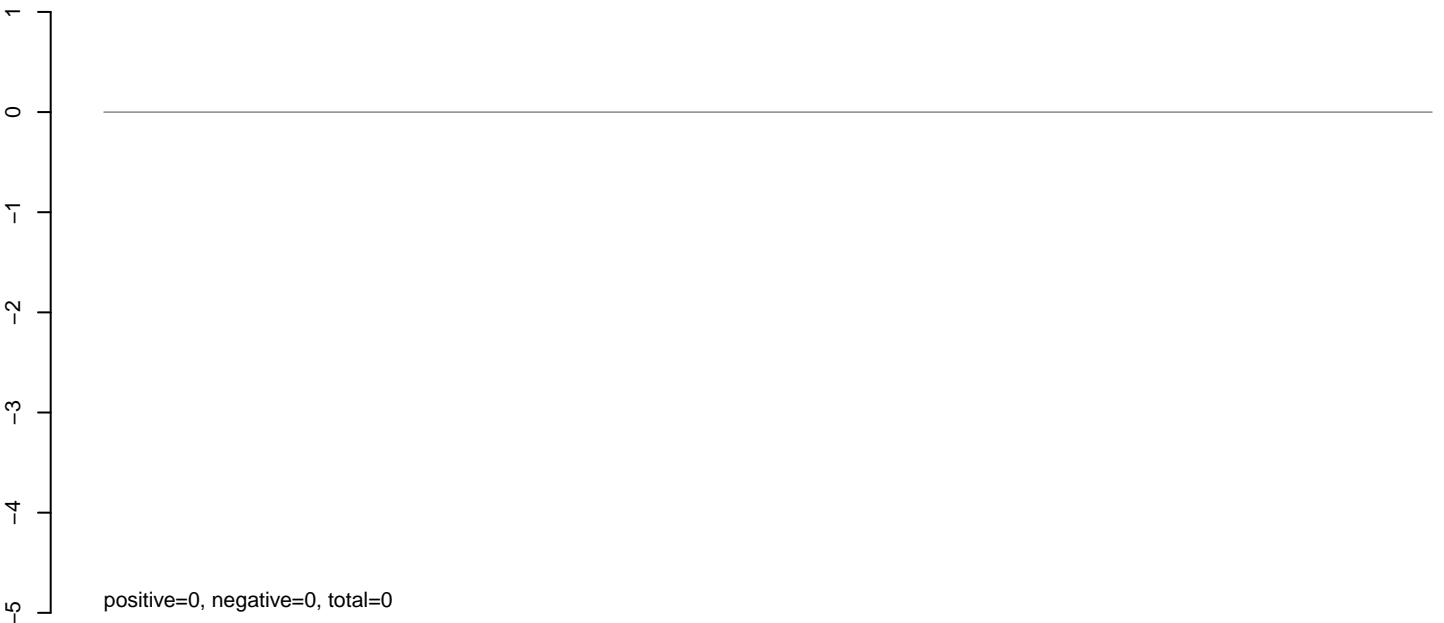
Window size=25, length=10723, virus@NC_012533.1-Kedougou_virus:1-10723



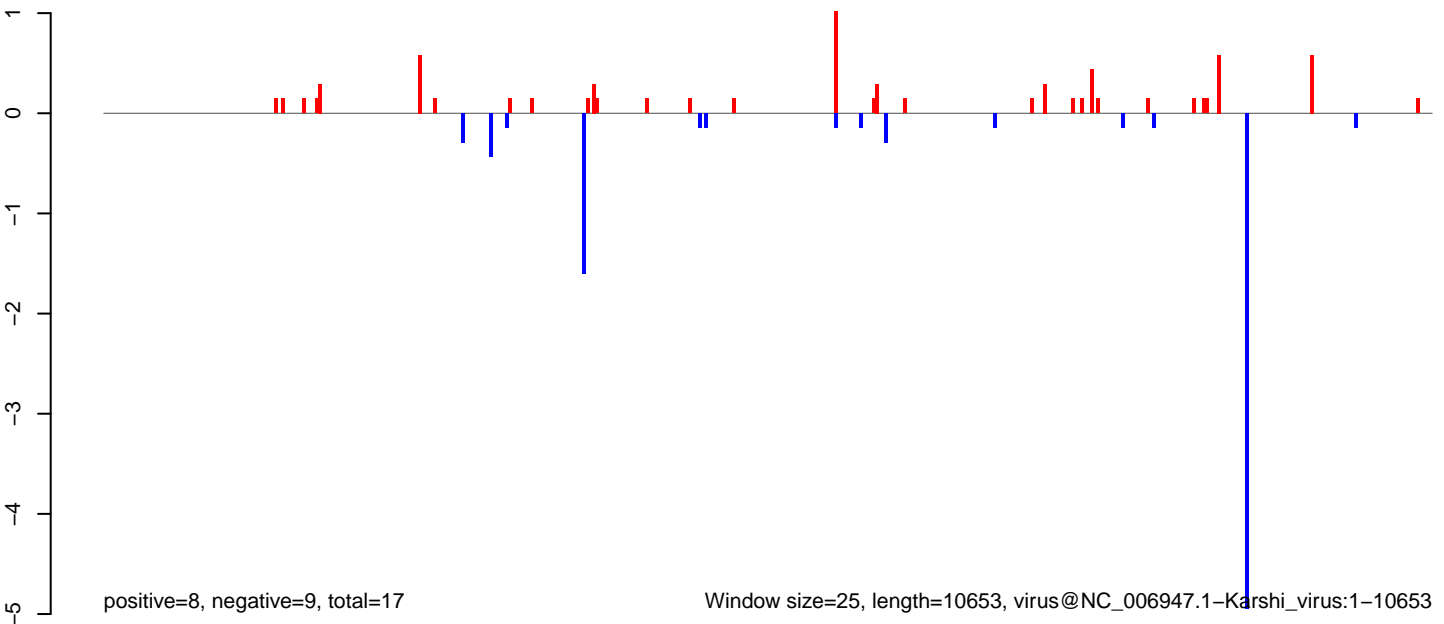
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



AeAeg_LVP_pMos3DBhspPBac.18_23.rep

0.4
0.2
0.0
-0.2

positive=6, negative=3, total=8

AeAeg_LVP_pMos3DBhspPBac.24_35.rep

0.4
0.2
0.0
-0.2

positive=0, negative=0, total=0

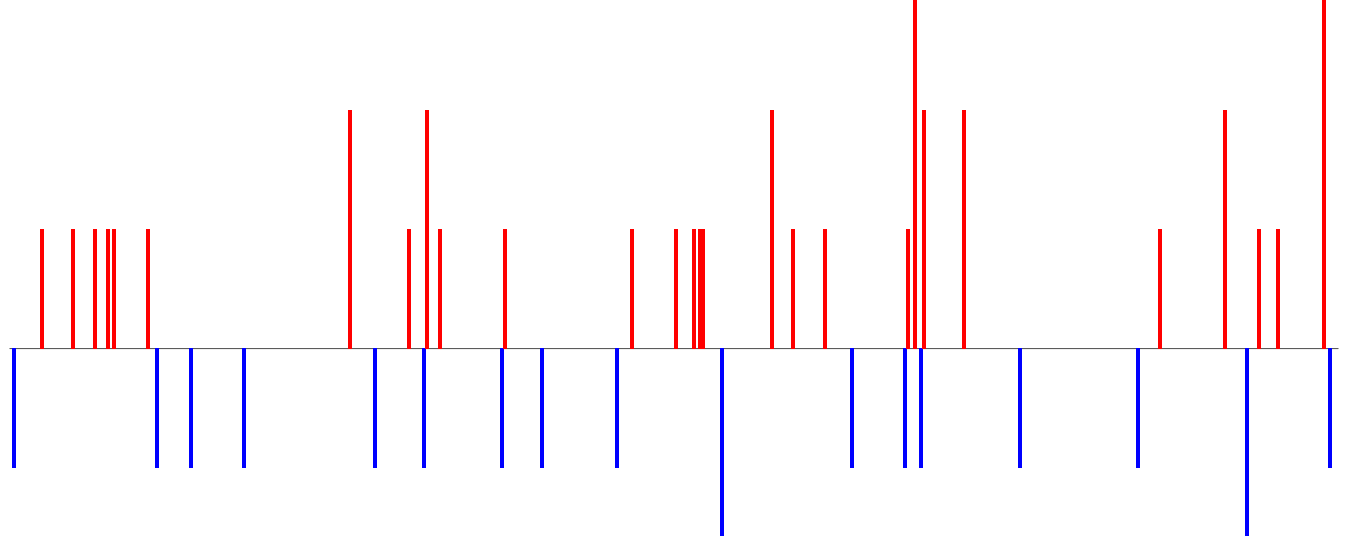
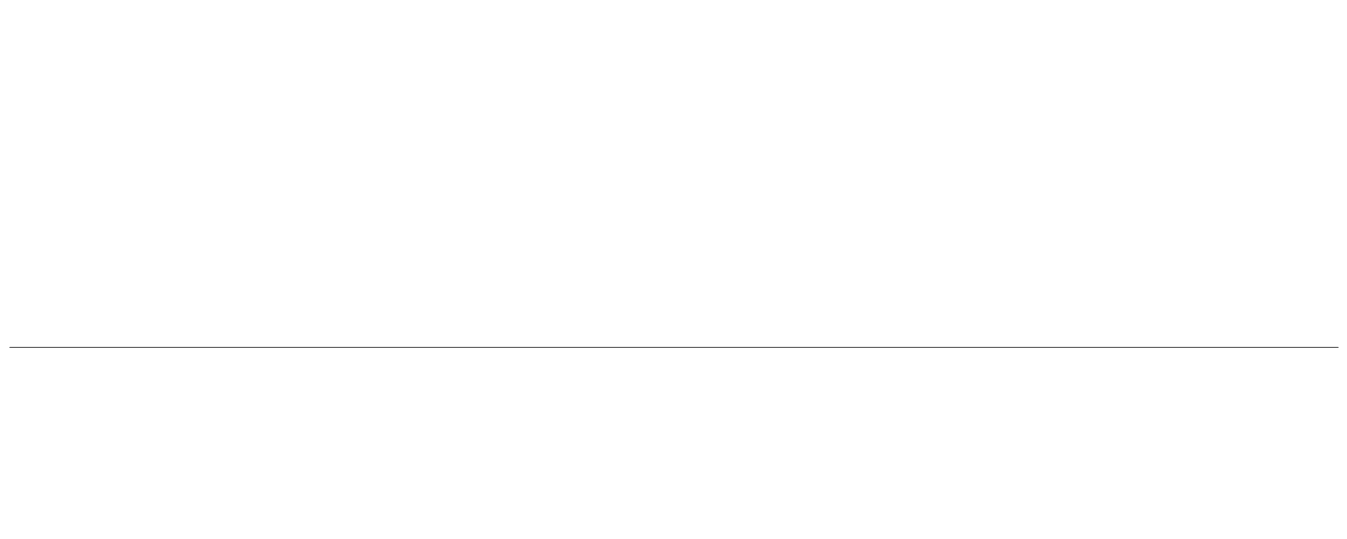
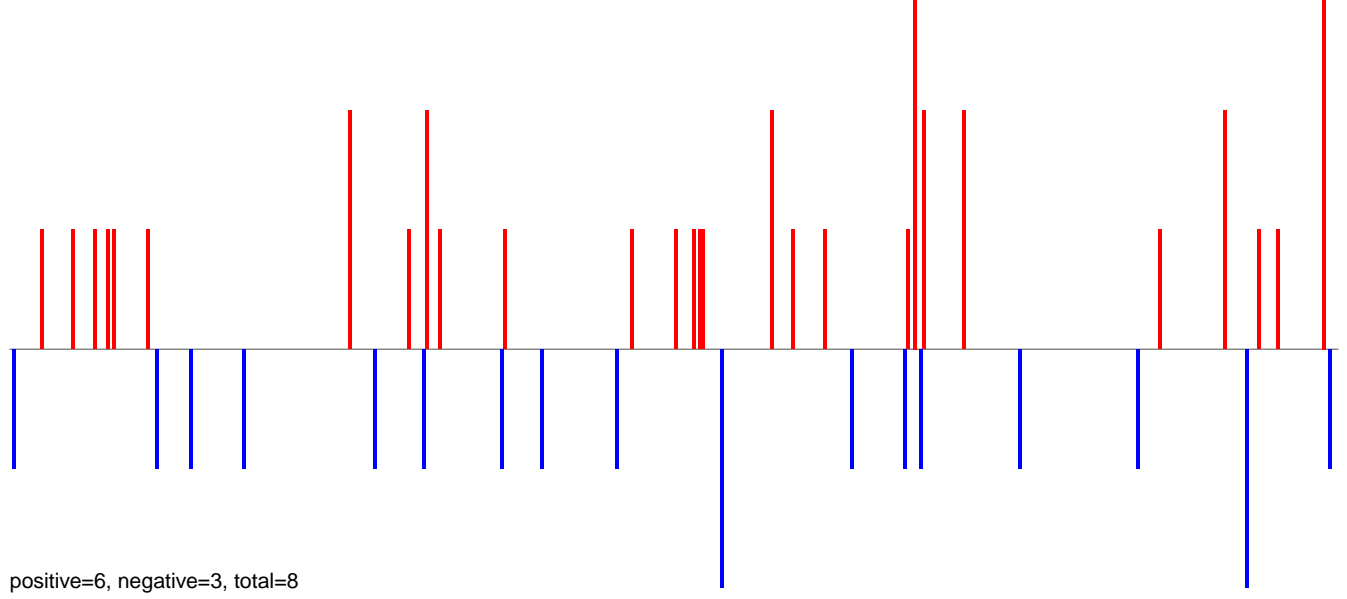
AeAeg_LVP_pMos3DBhspPBac.rep

0.4
0.2
0.0
-0.2

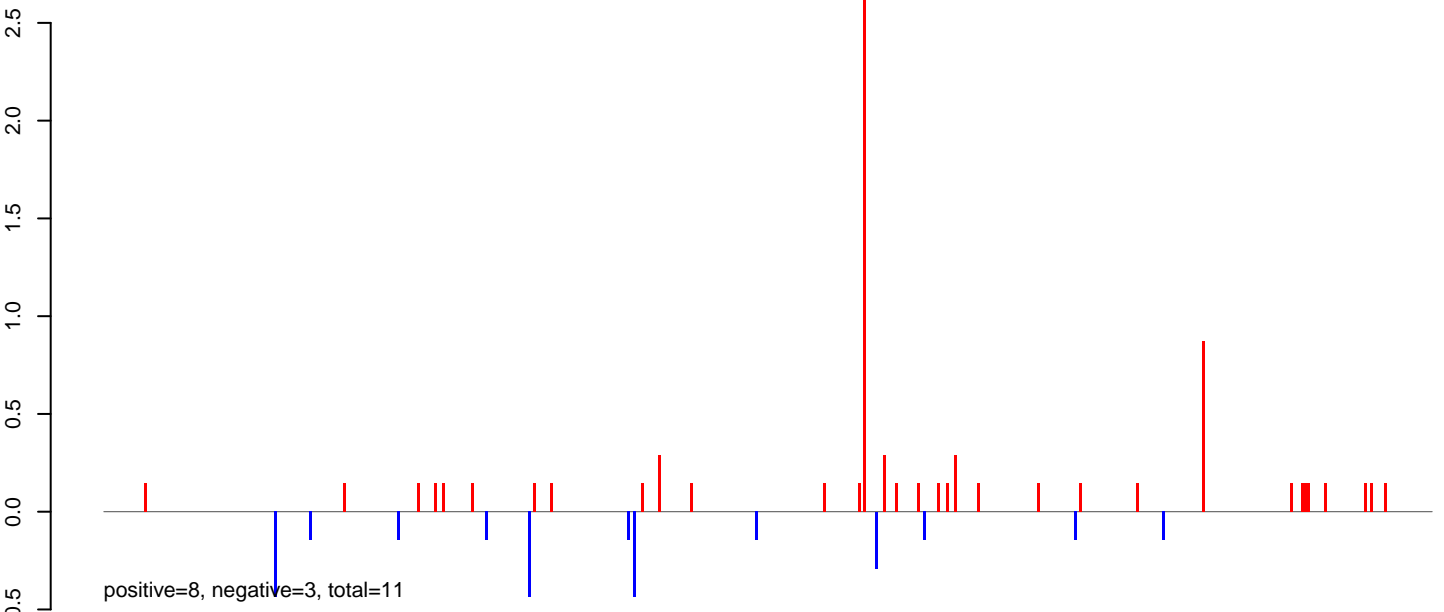
positive=6, negative=3, total=8

Window size=25, length=10685, virus@NC_004355.1-Alkhurma_virus:1-10685

0 2000 4000 6000 8000 10000



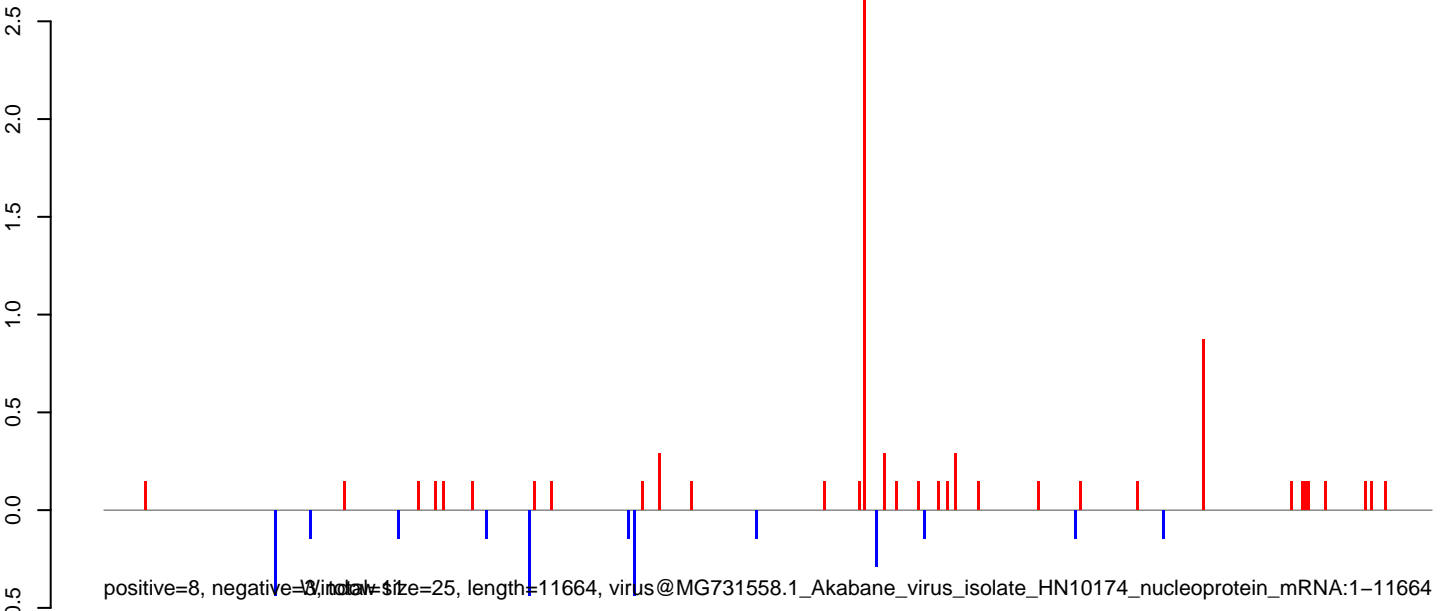
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



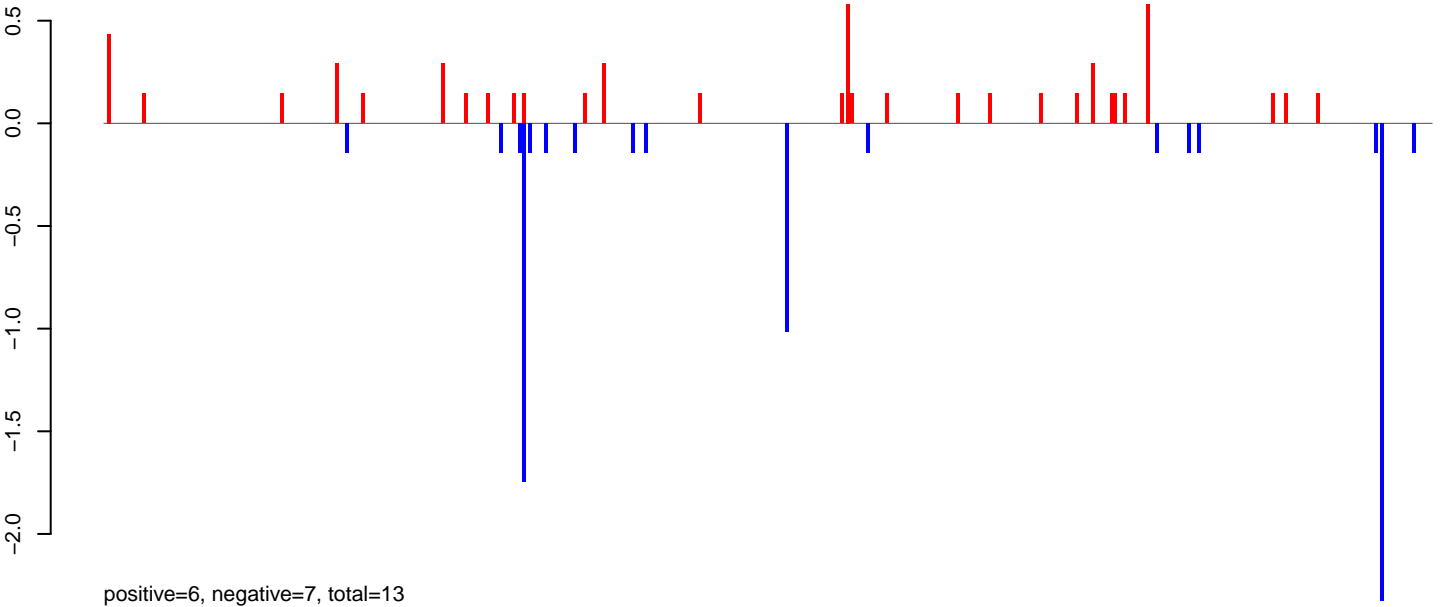
AeAeg_LVP_pMos3DBhspPBac.rep



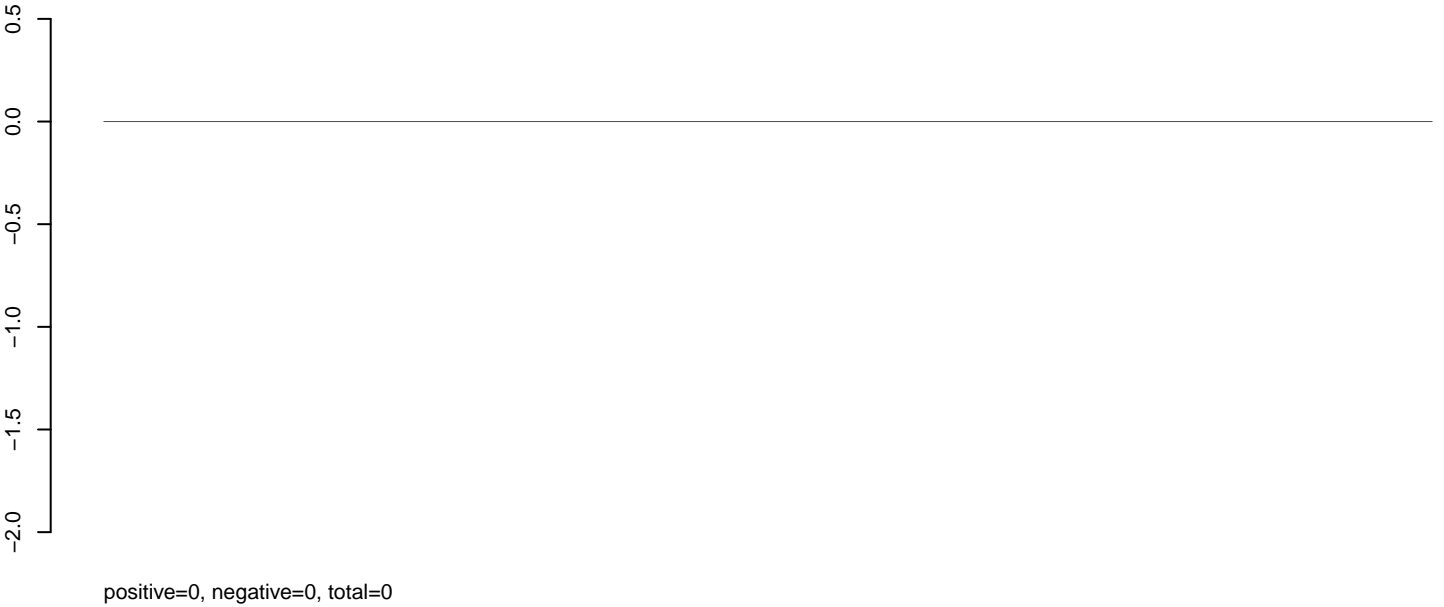
positive=8, negative=3, total=11, size=25, length=11664, virus@MG731558.1_Akabane_virus_isolate_HN10174_nucleoprotein_mRNA:1-11664

0 2000 4000 6000 8000 10000 12000

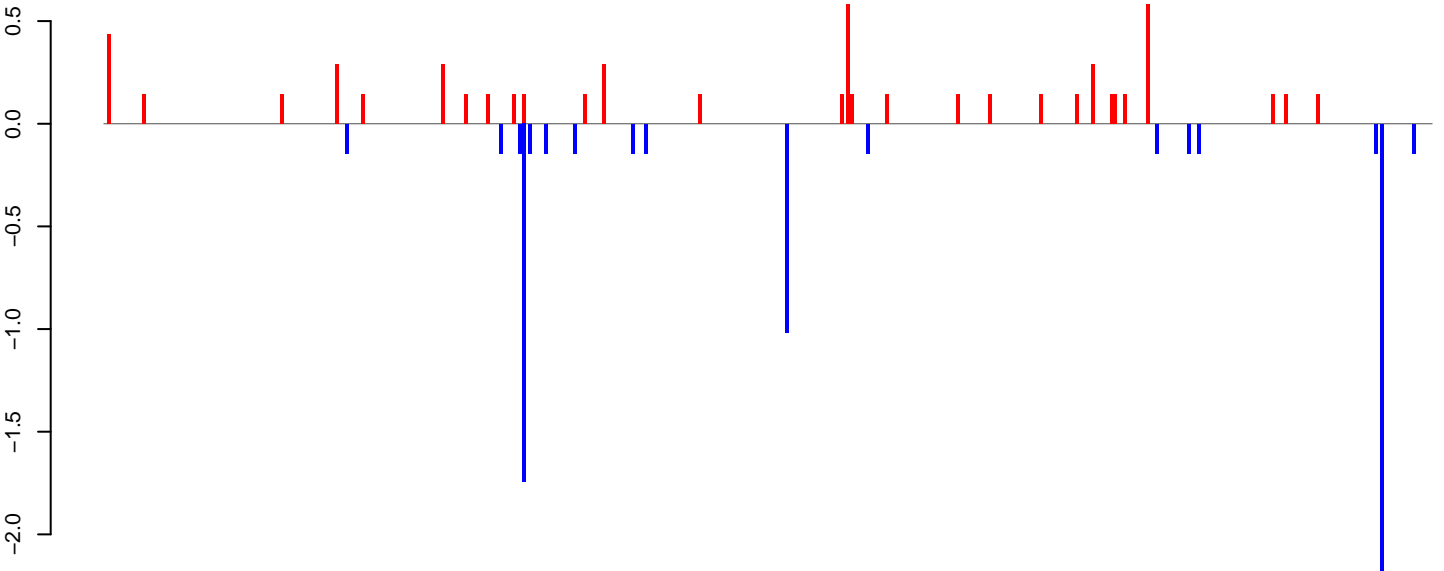
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



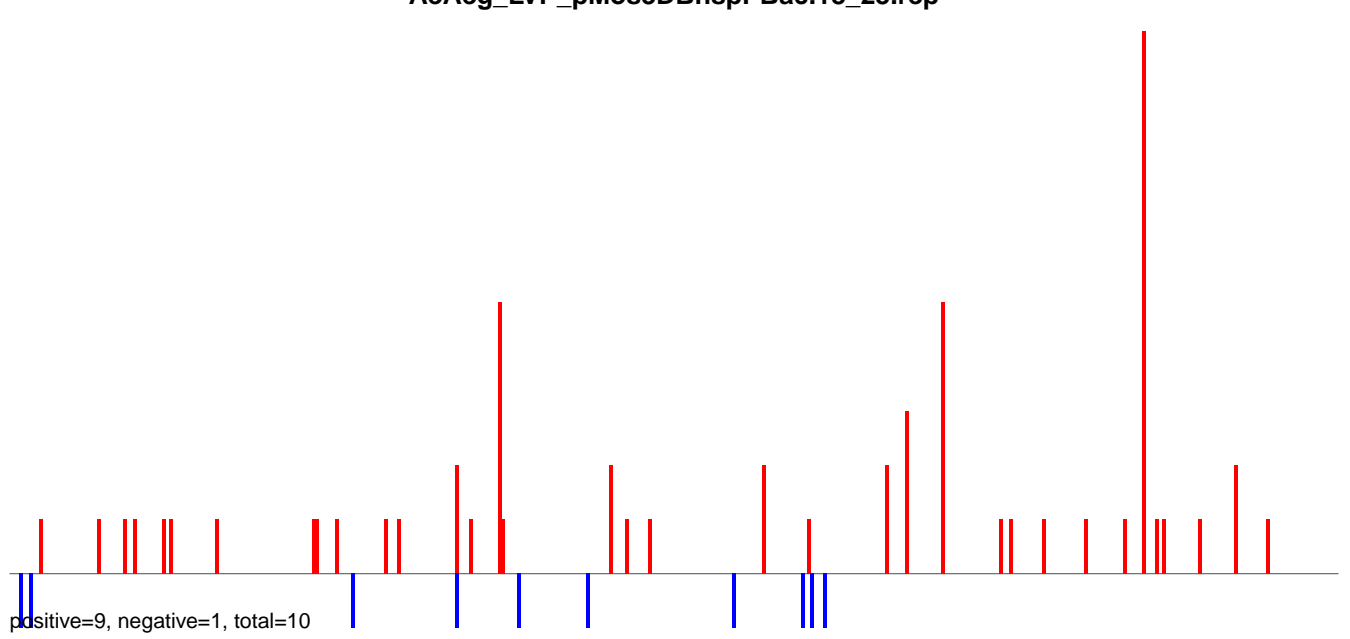
AeAeg_LVP_pMos3DBhspPBac.rep



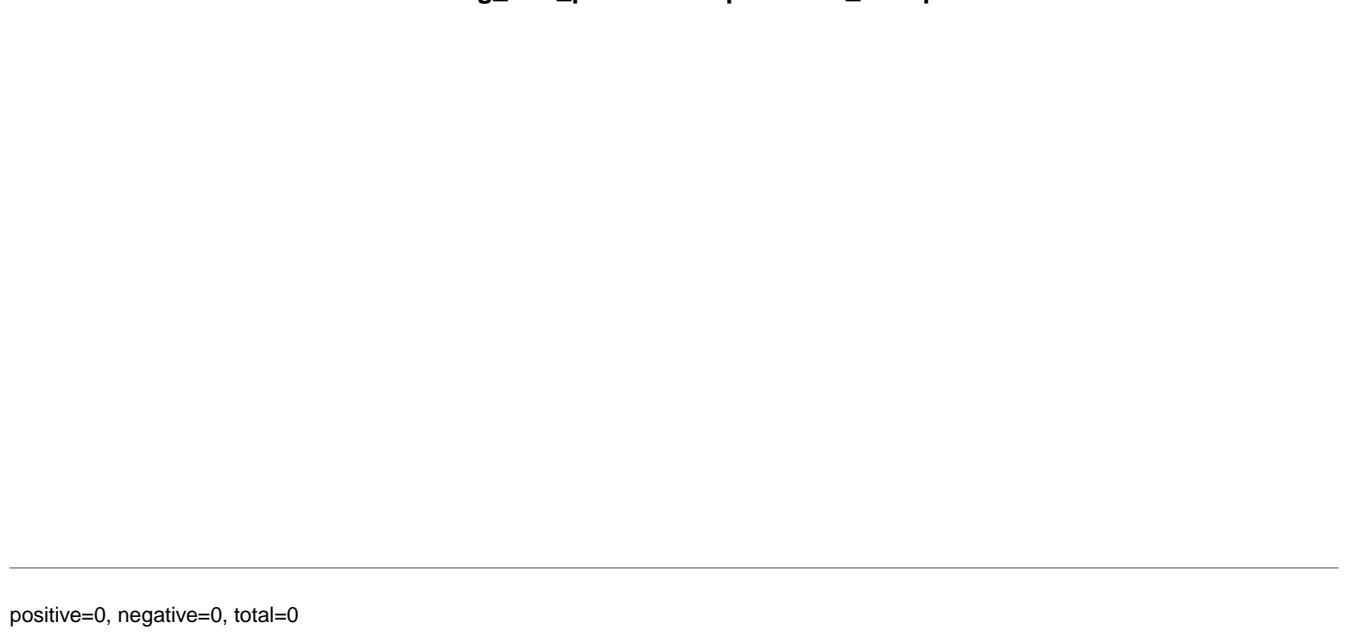
width=25, length=10302, virus@EU082200.2-Koutango_virus_strain_Dak_Ar_D_5443_polyprotein_gene:1-10302

0 2000 4000 6000 8000 10000

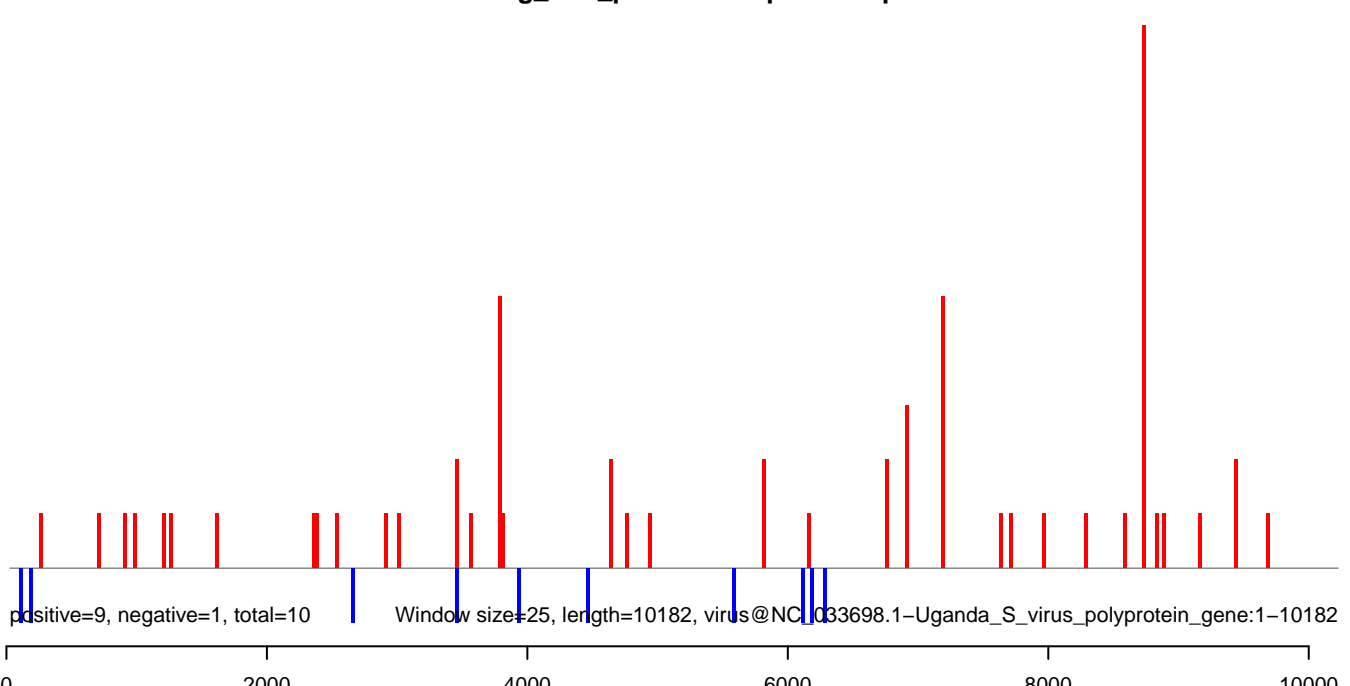
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



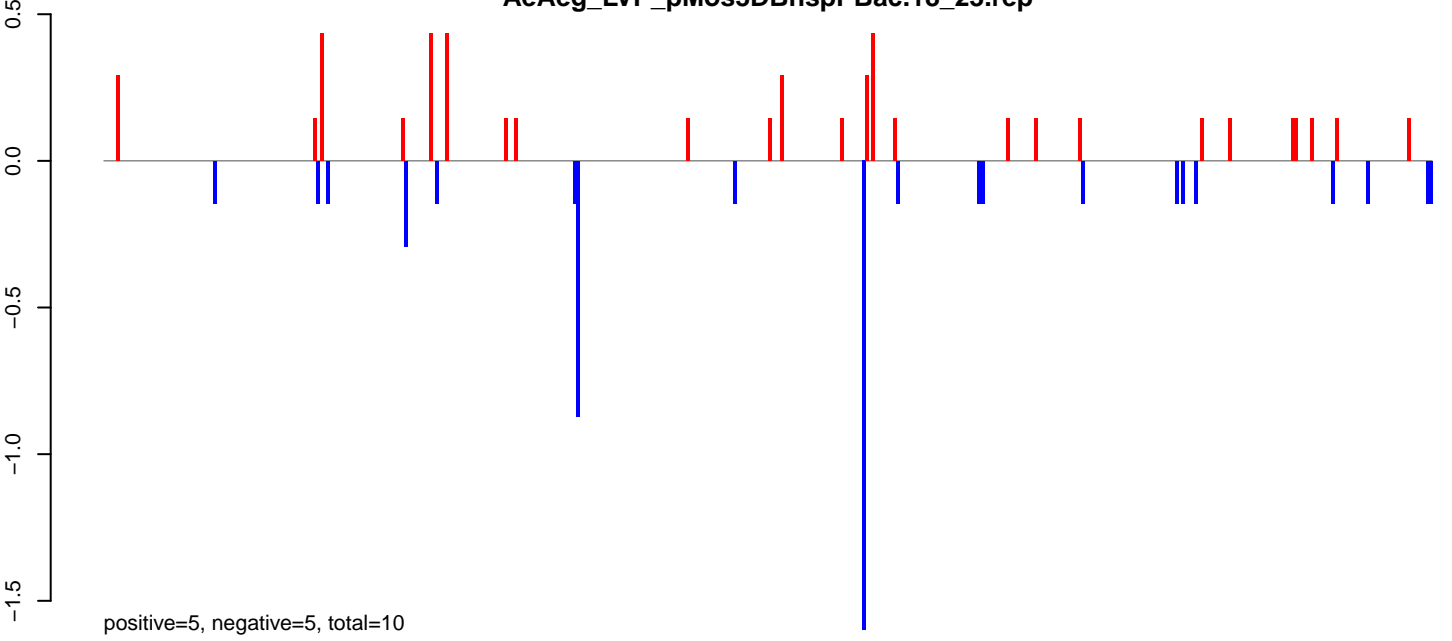
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



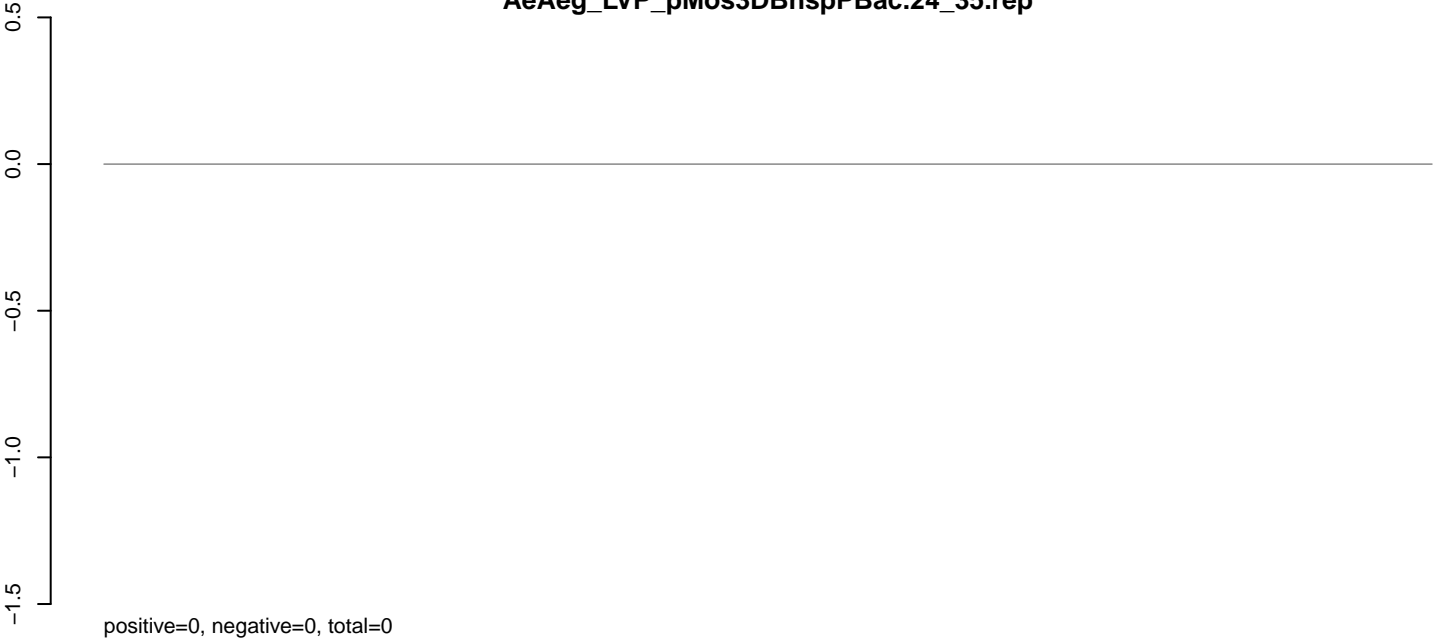
AeAeg_LVP_pMos3DBhspPBac.rep



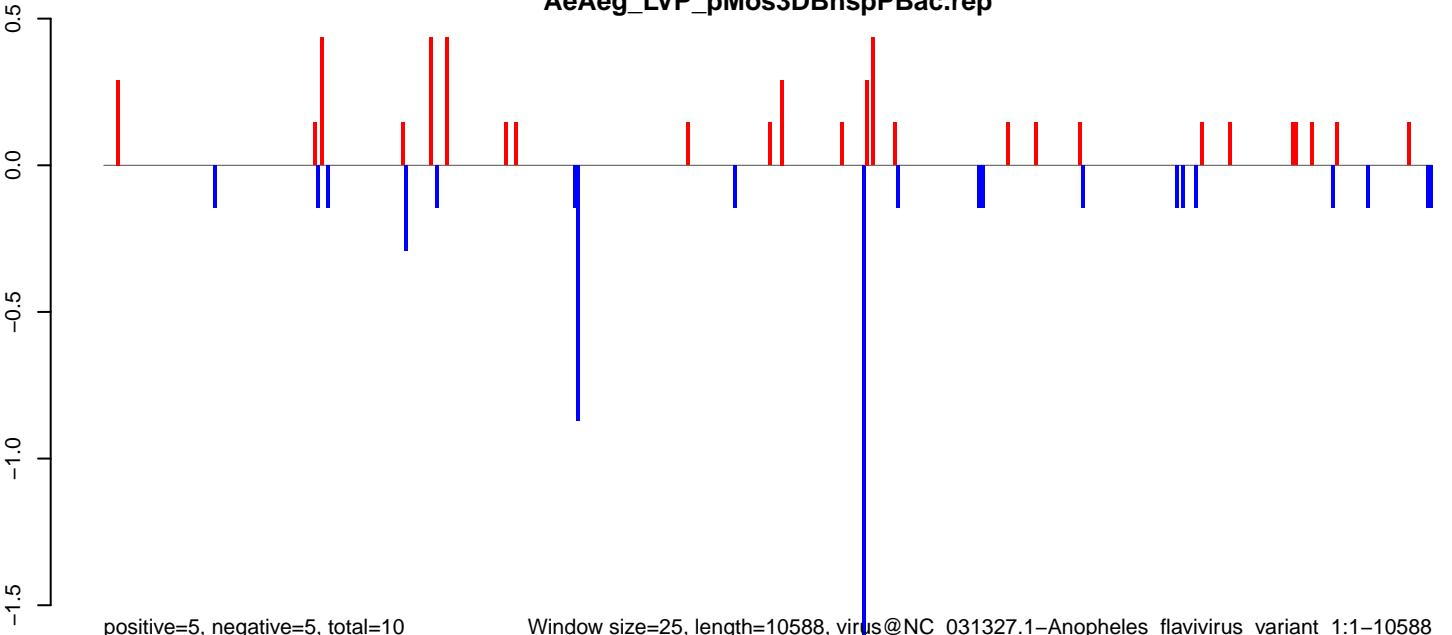
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



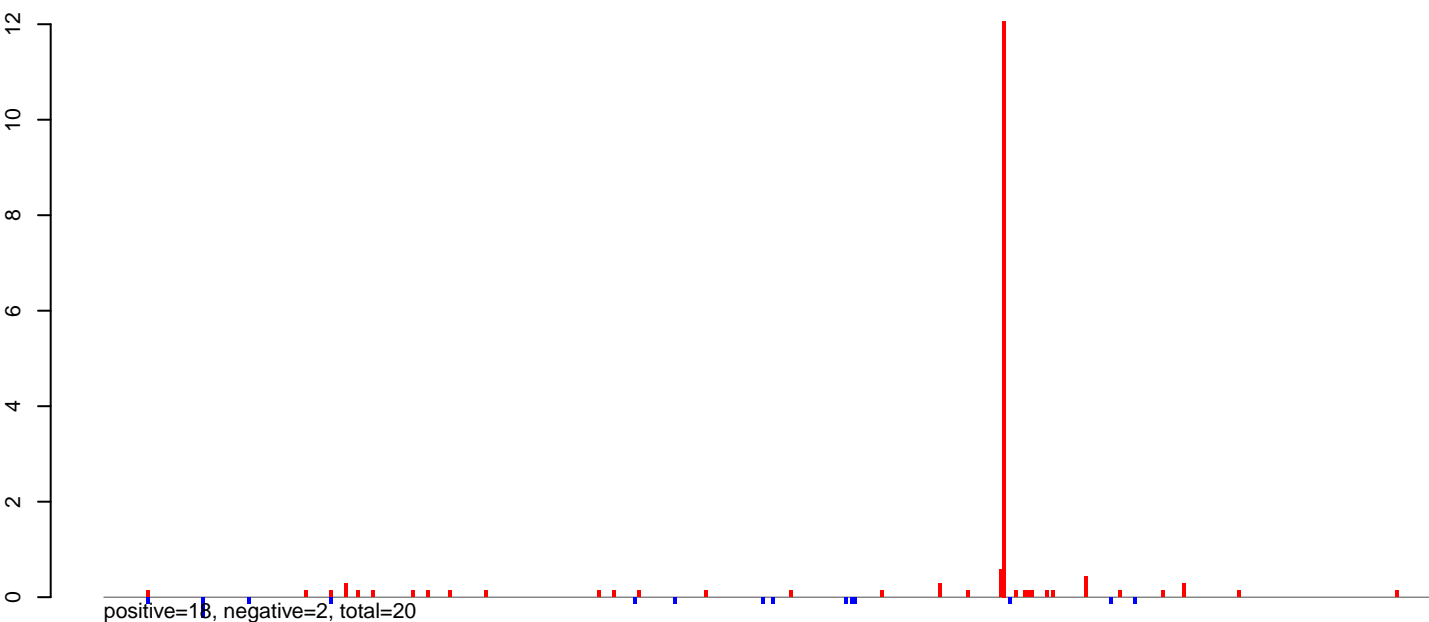
AeAeg_LVP_pMos3DBhspPBac.rep



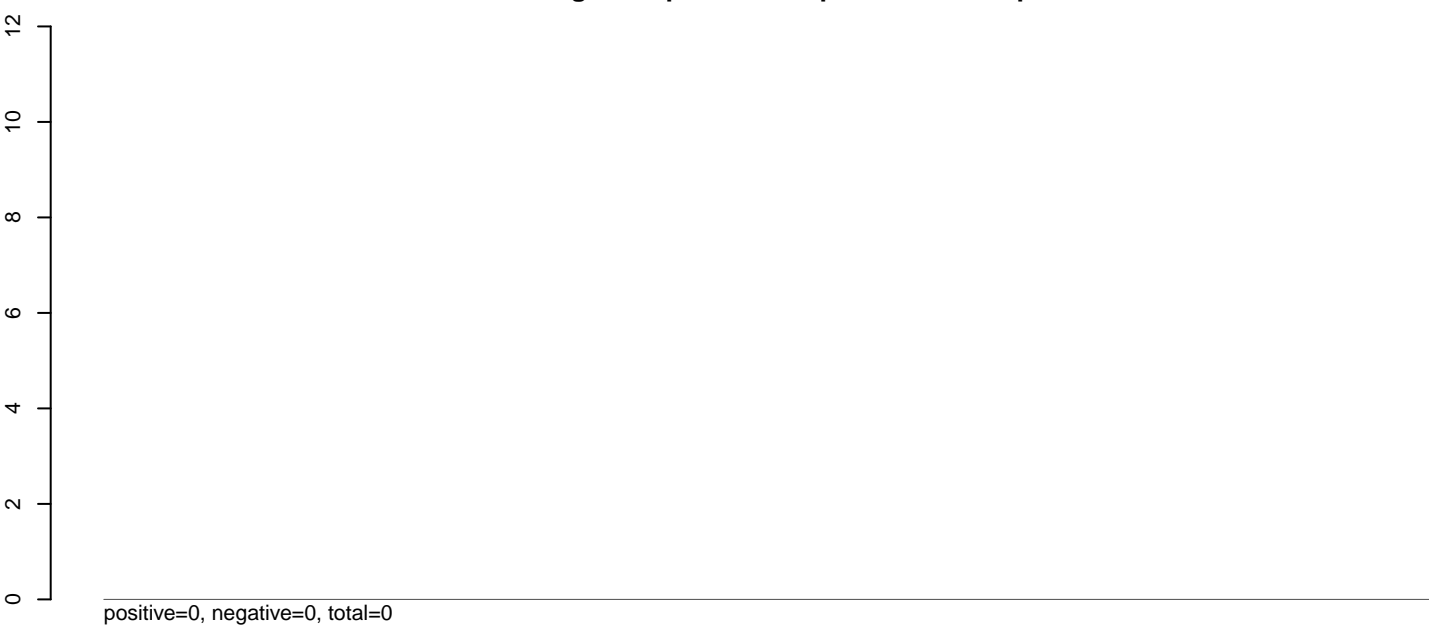
Window size=25, length=10588, virus@NC_031327.1-Anopheles_flavivirus_variant_1:1-10588

0 2000 4000 6000 8000 10000

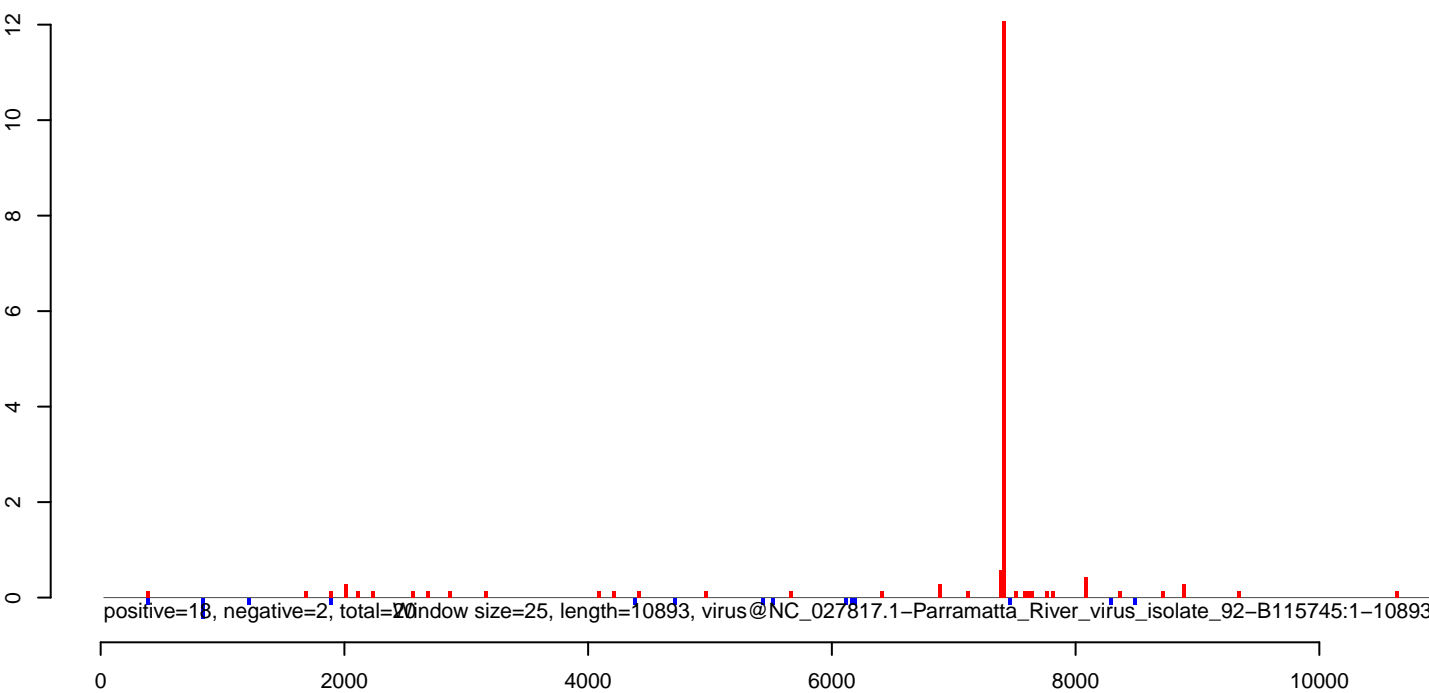
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



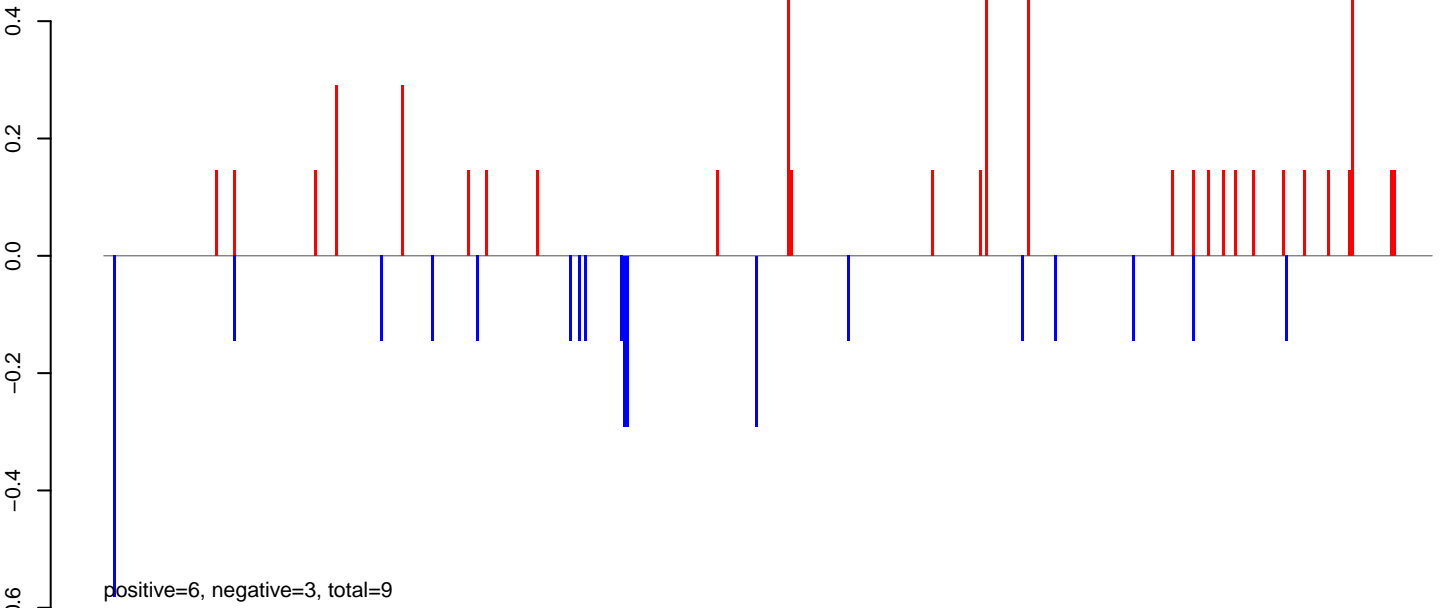
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



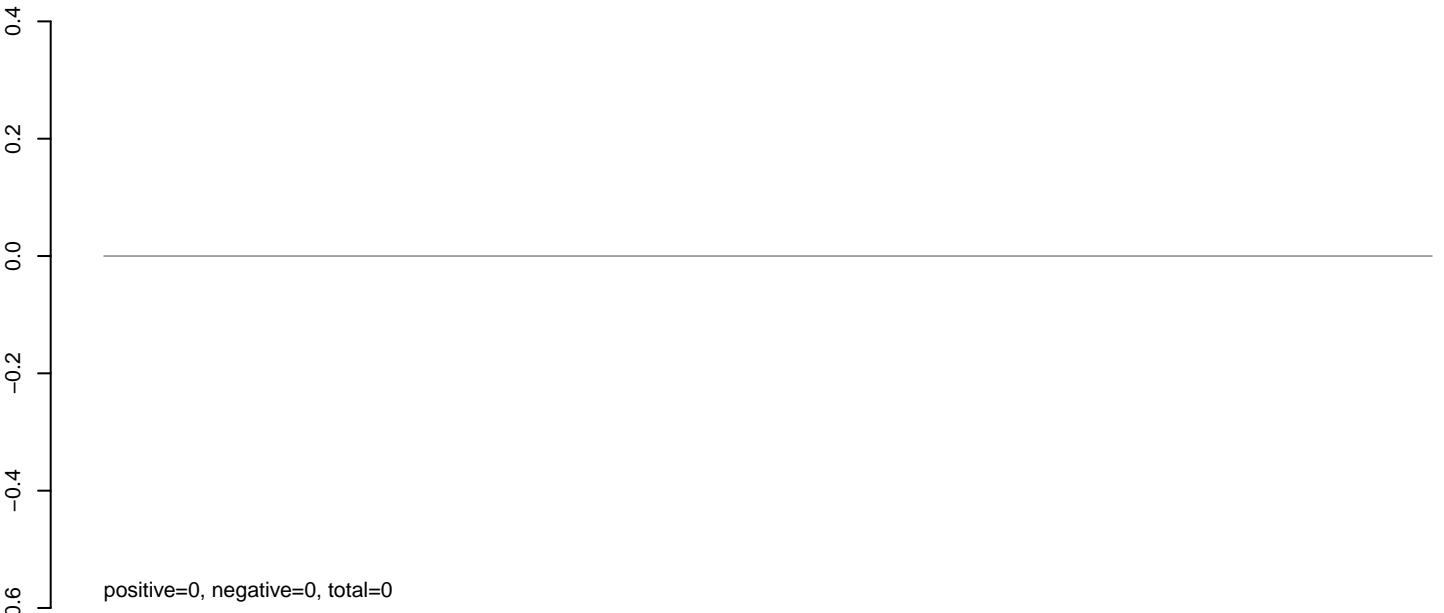
AeAeg_LVP_pMos3DBhspPBac.rep



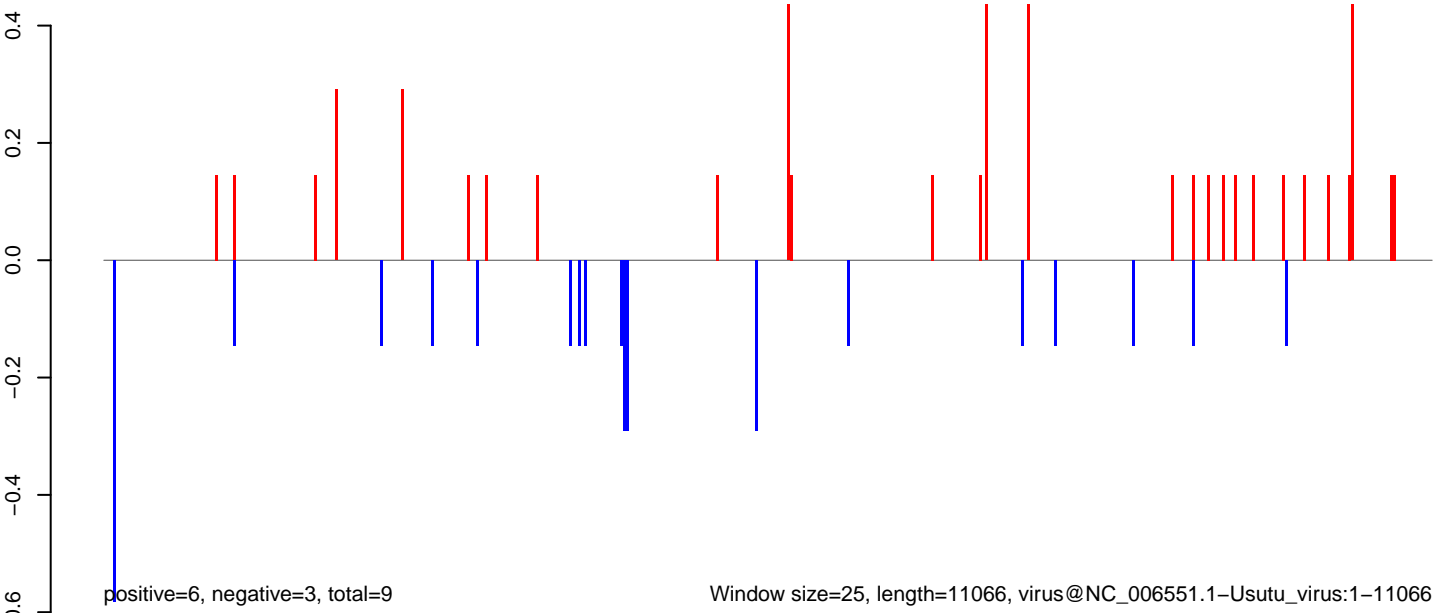
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



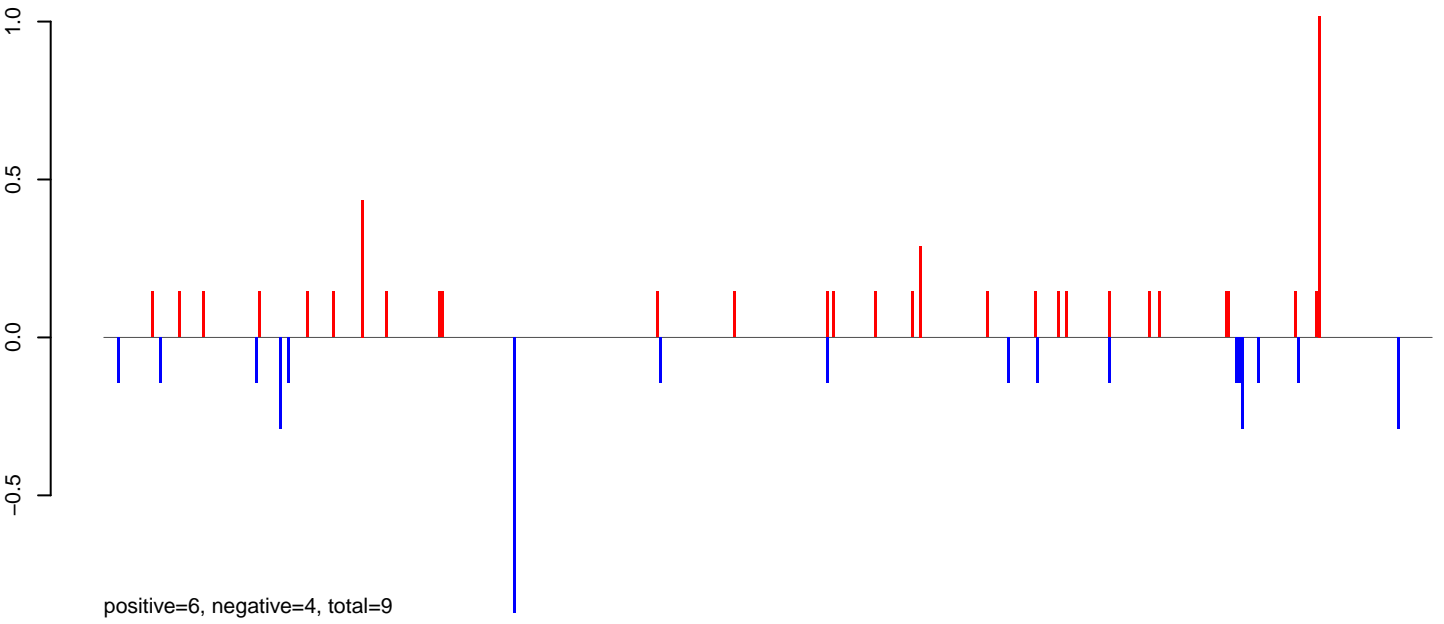
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



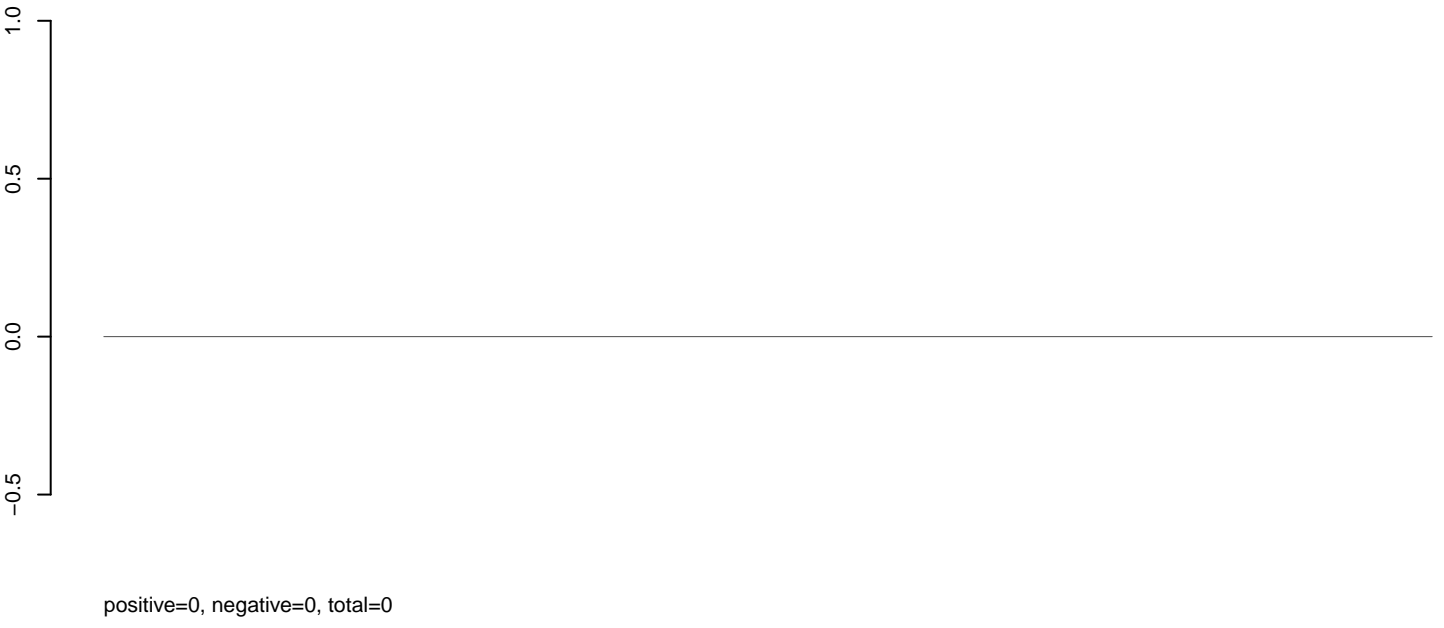
AeAeg_LVP_pMos3DBhspPBac.rep



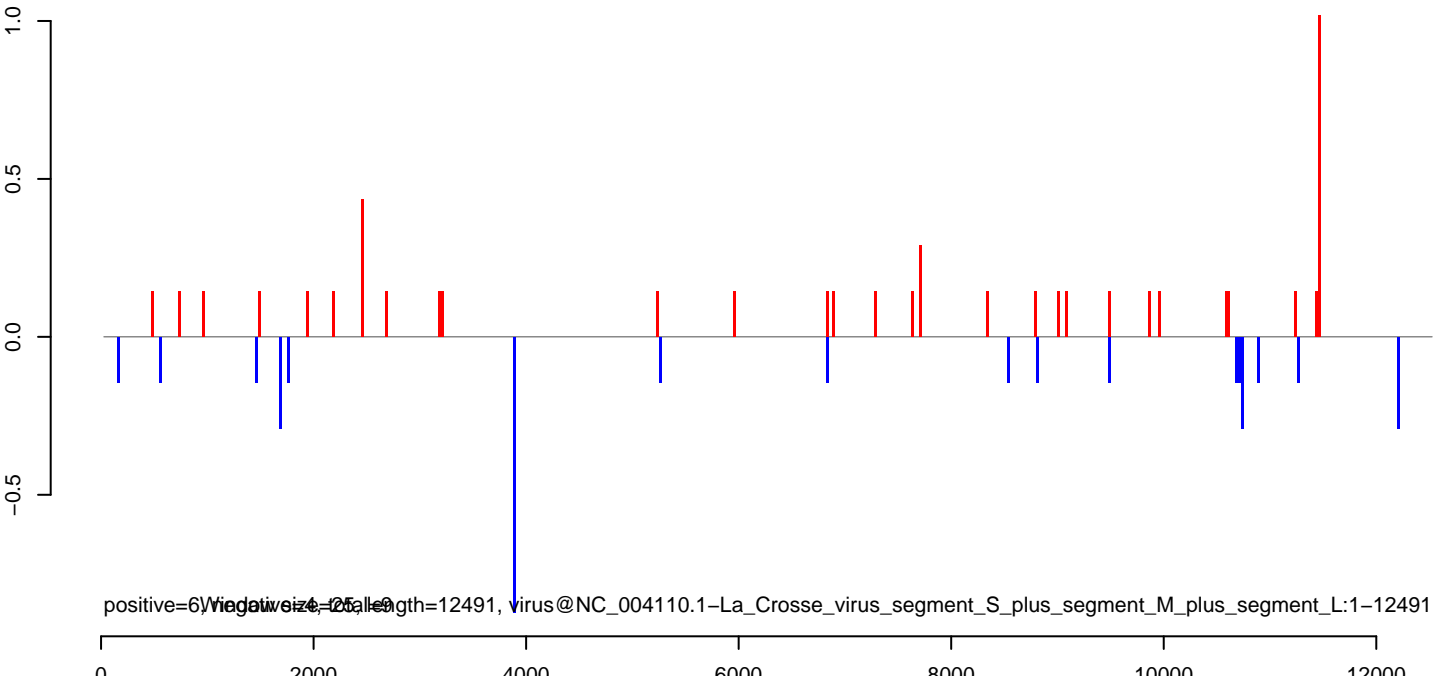
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



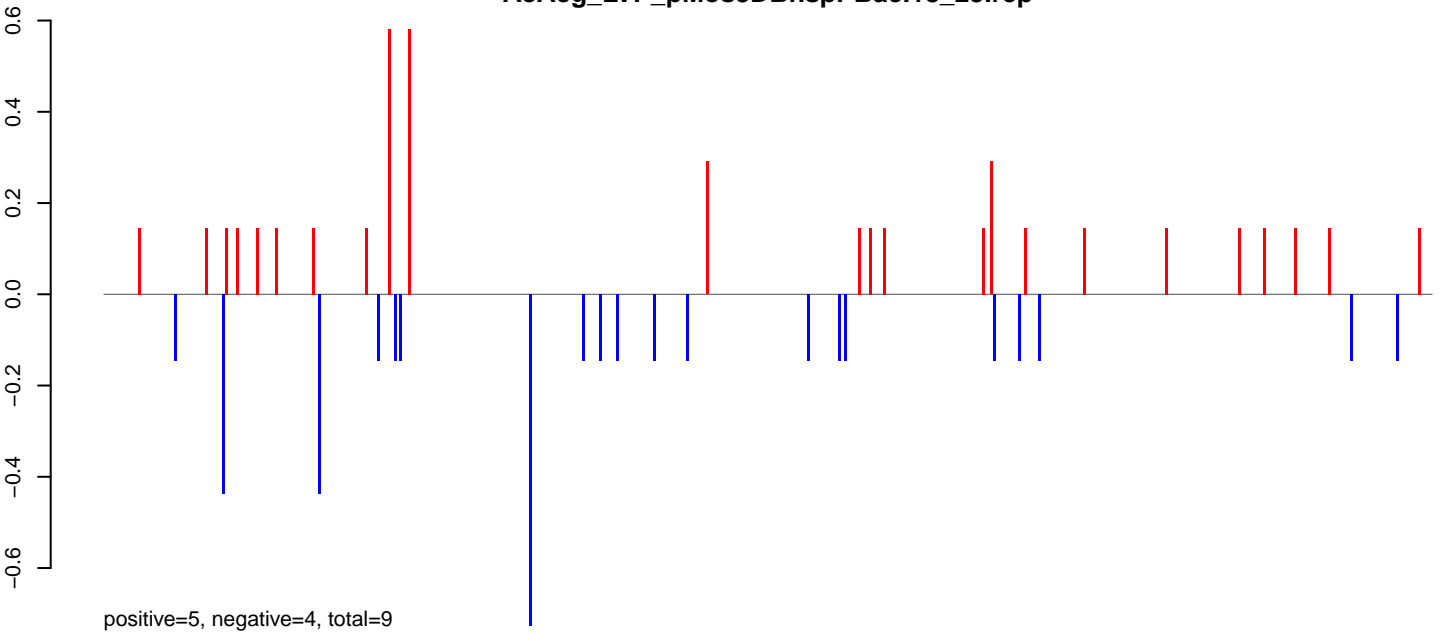
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



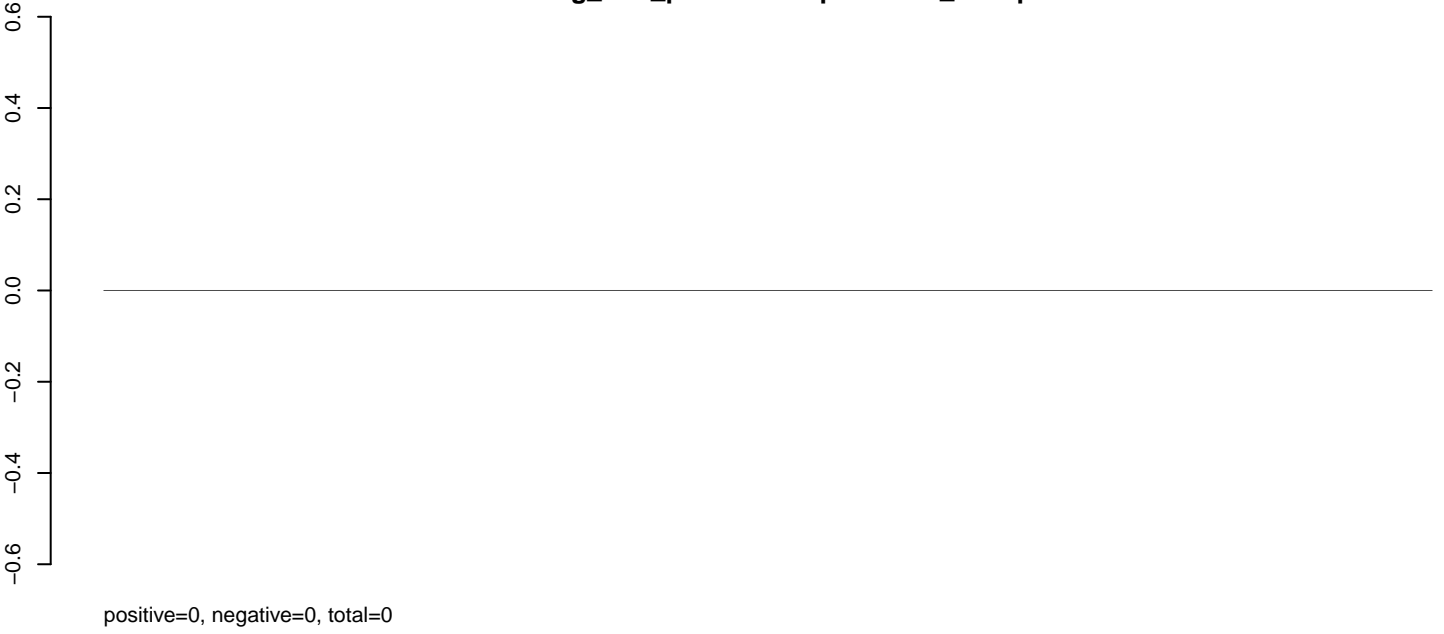
AeAeg_LVP_pMos3DBhspPBac.rep



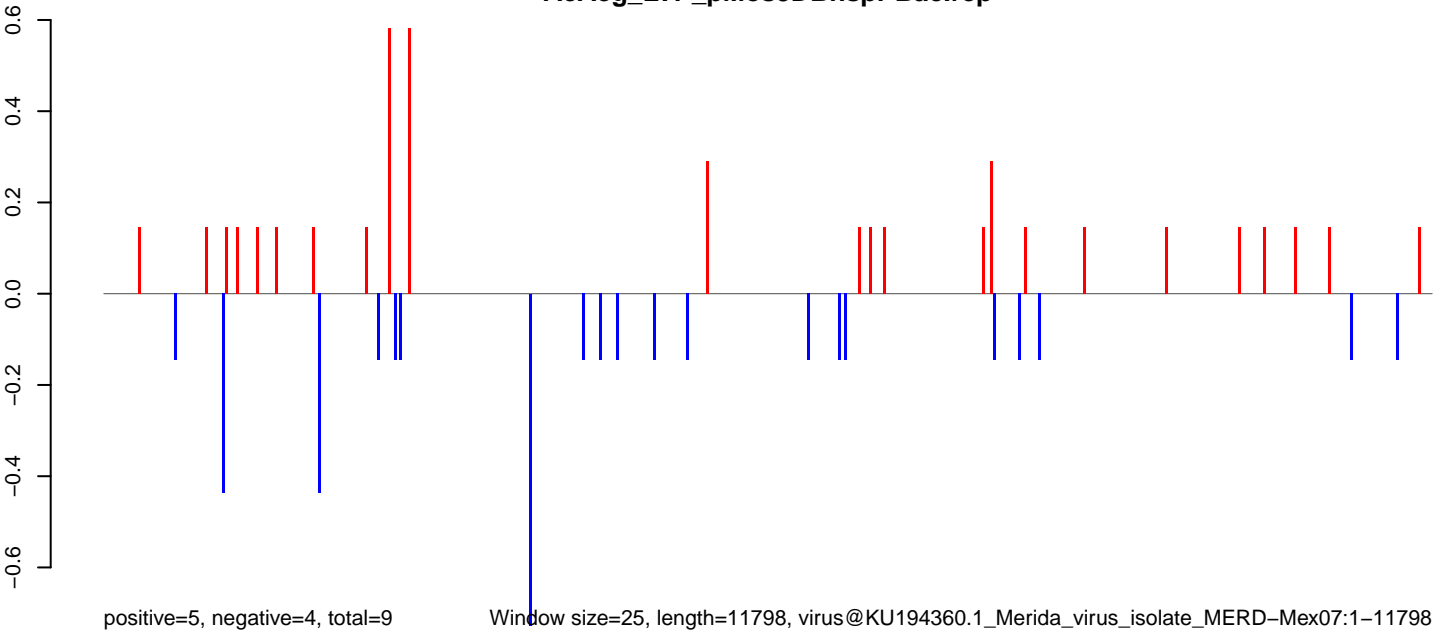
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

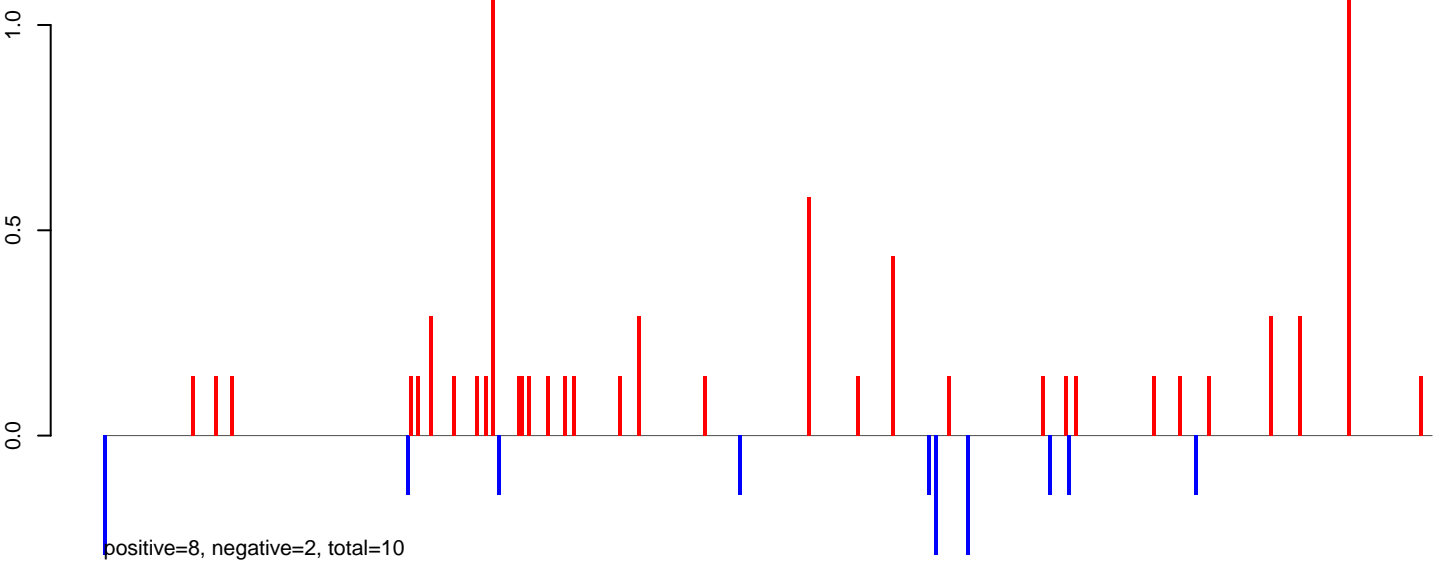


AeAeg_LVP_pMos3DBhspPBac.rep



0 2000 4000 6000 8000 10000 12000

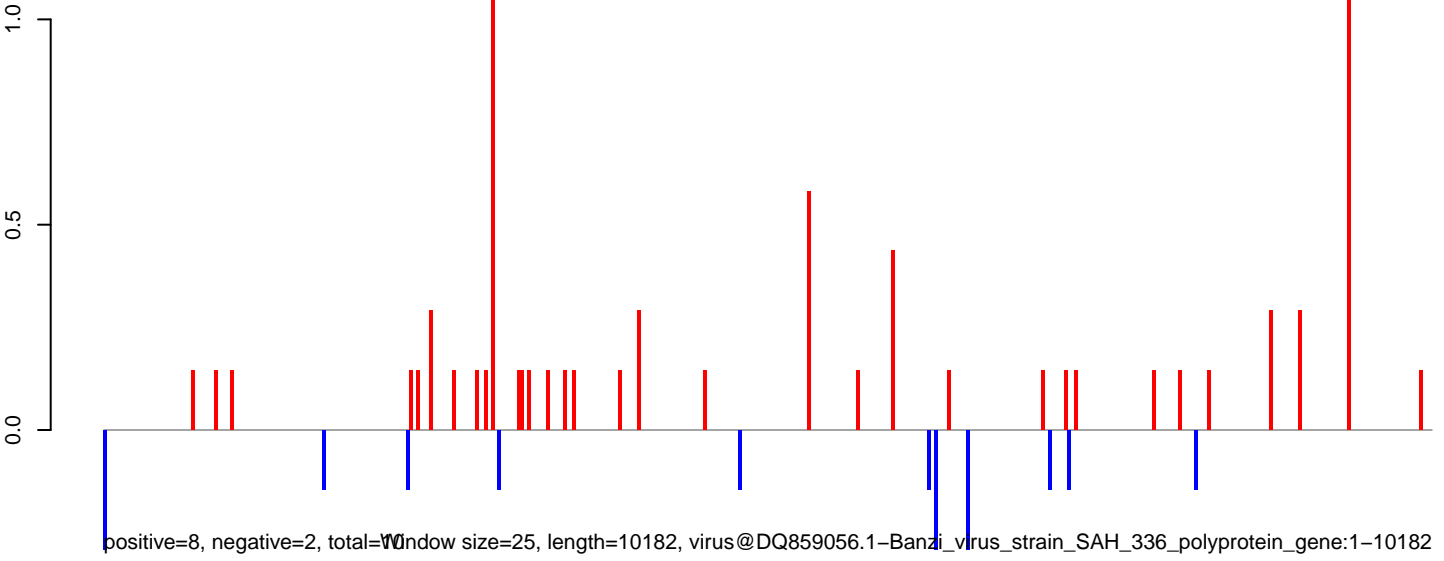
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



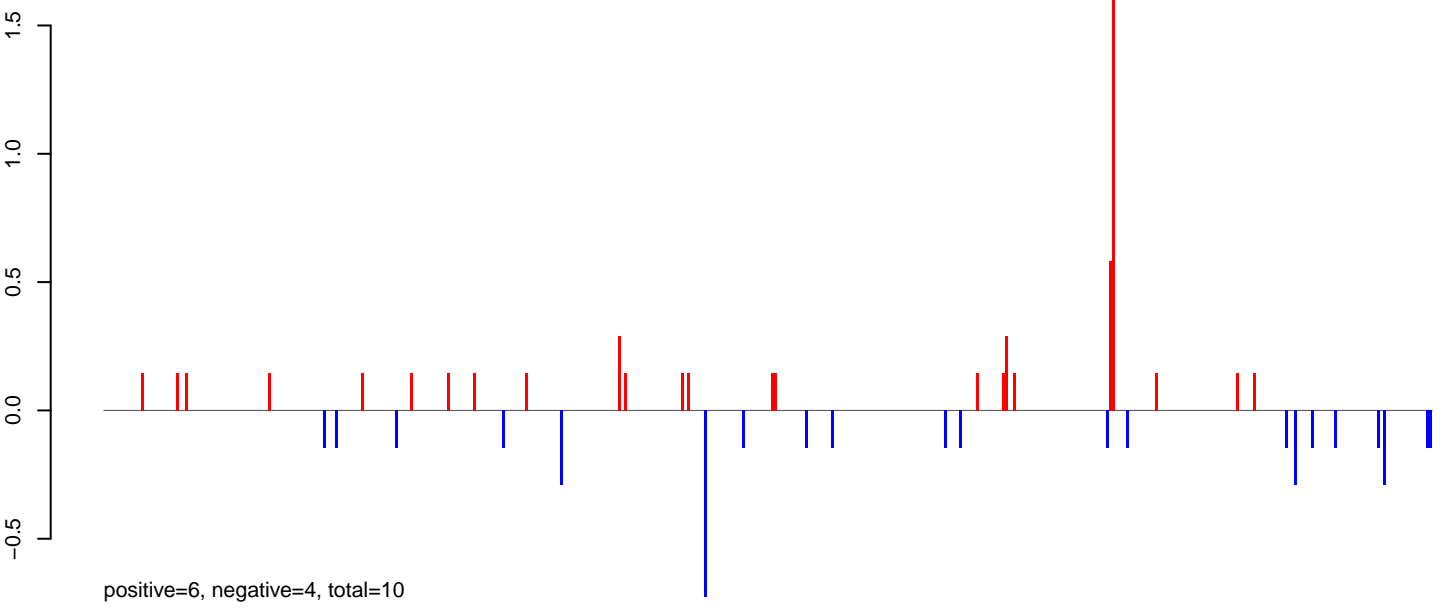
AeAeg_LVP_pMos3DBhspPBac.rep



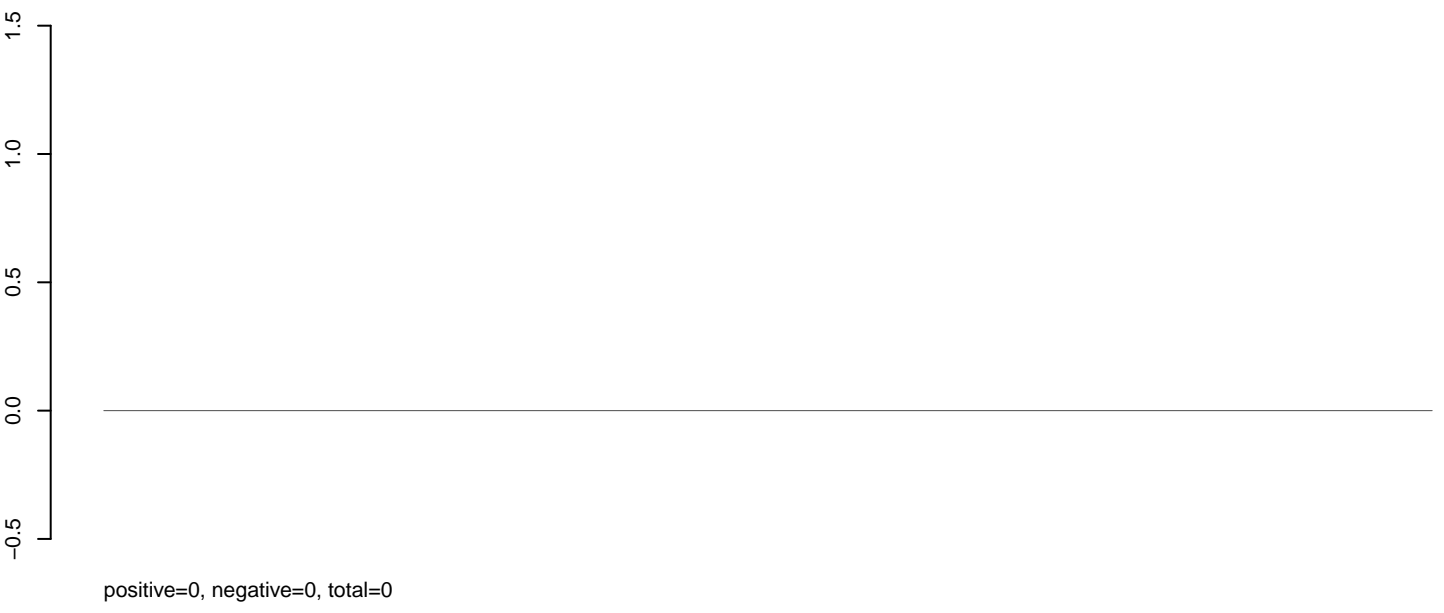
Window size=25, length=10182, virus@DQ859056.1-Banzi_virus_strain_SAH_336_polyprotein_gene:1-10182

0 2000 4000 6000 8000 10000

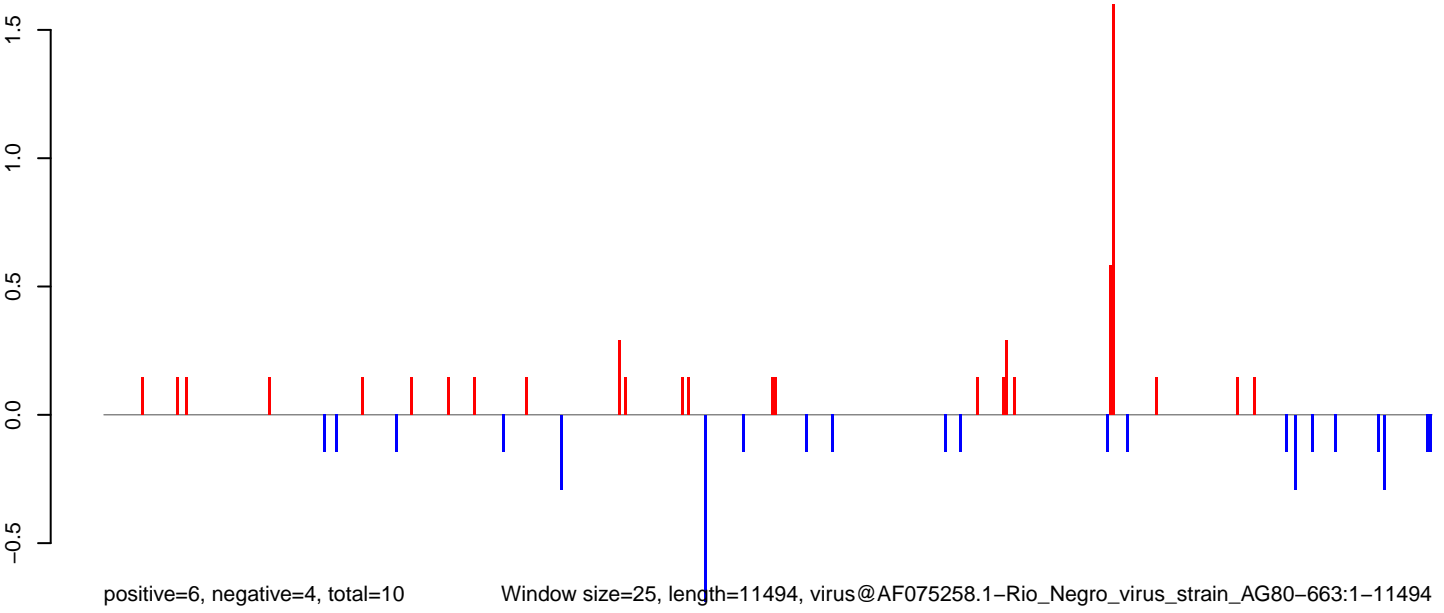
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



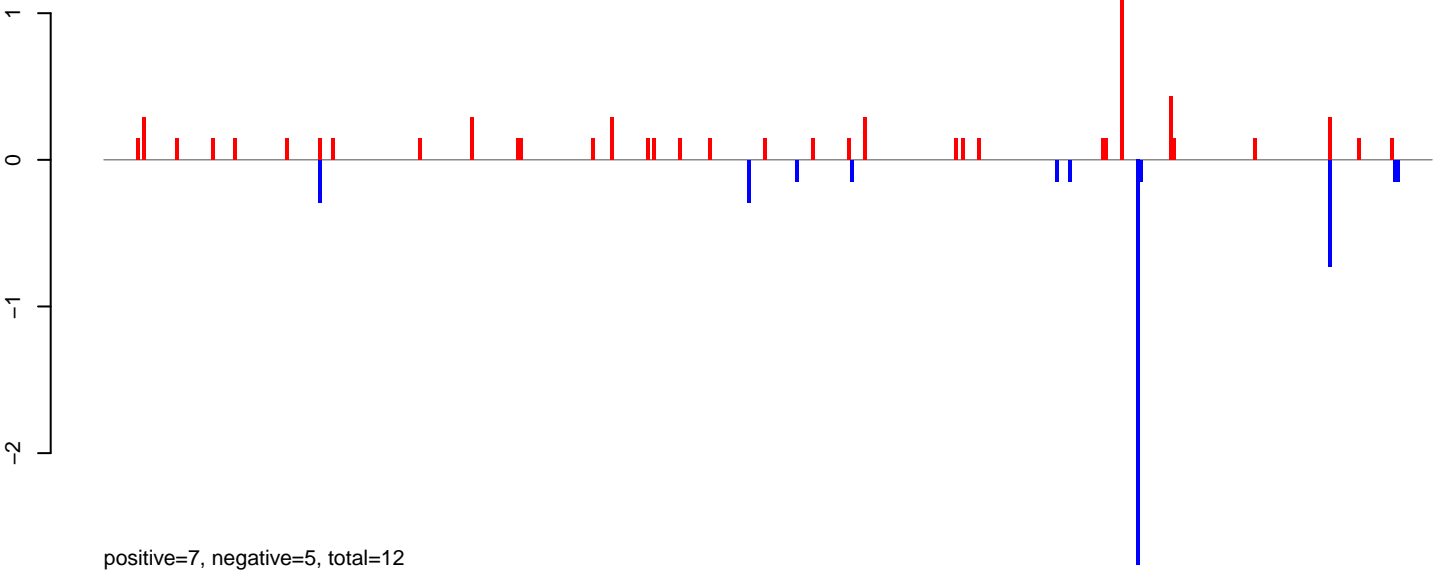
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



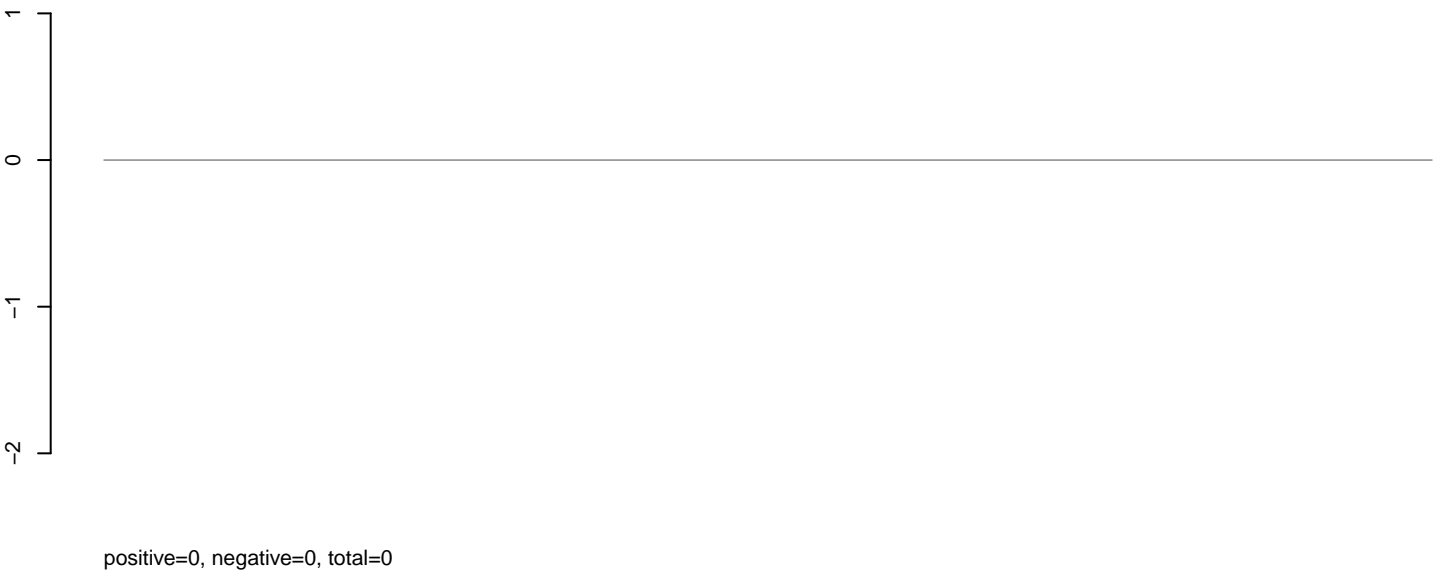
AeAeg_LVP_pMos3DBhspPBac.rep



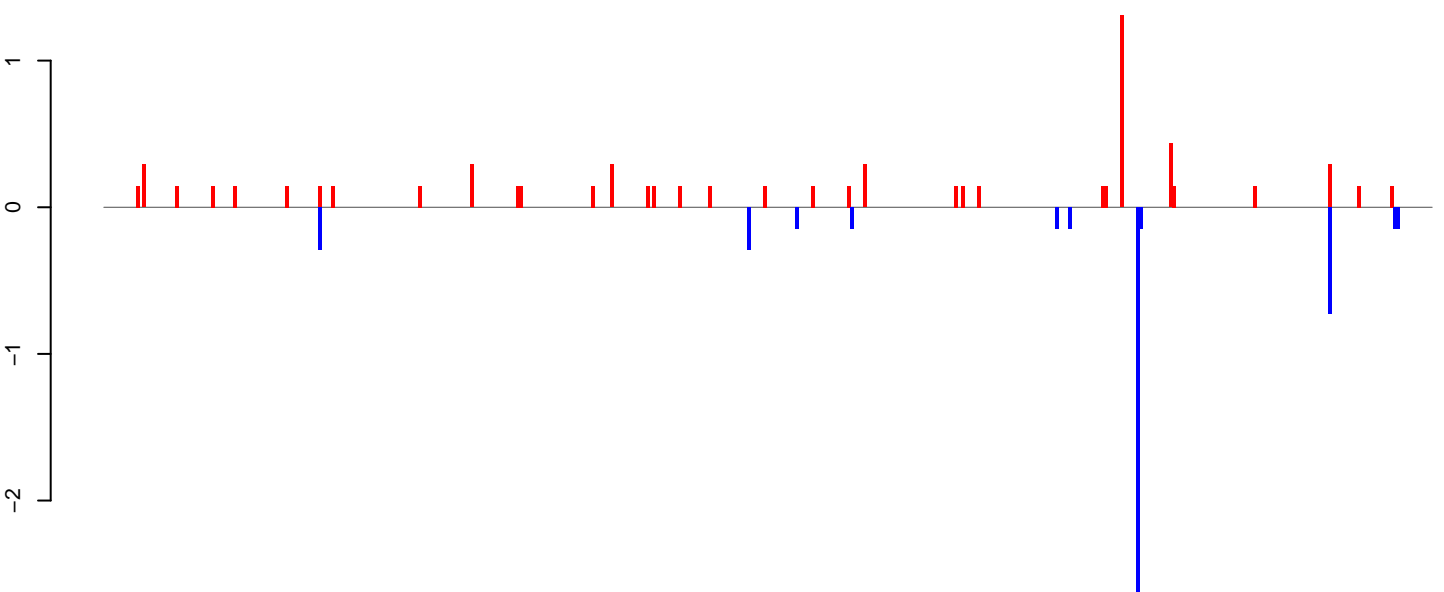
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



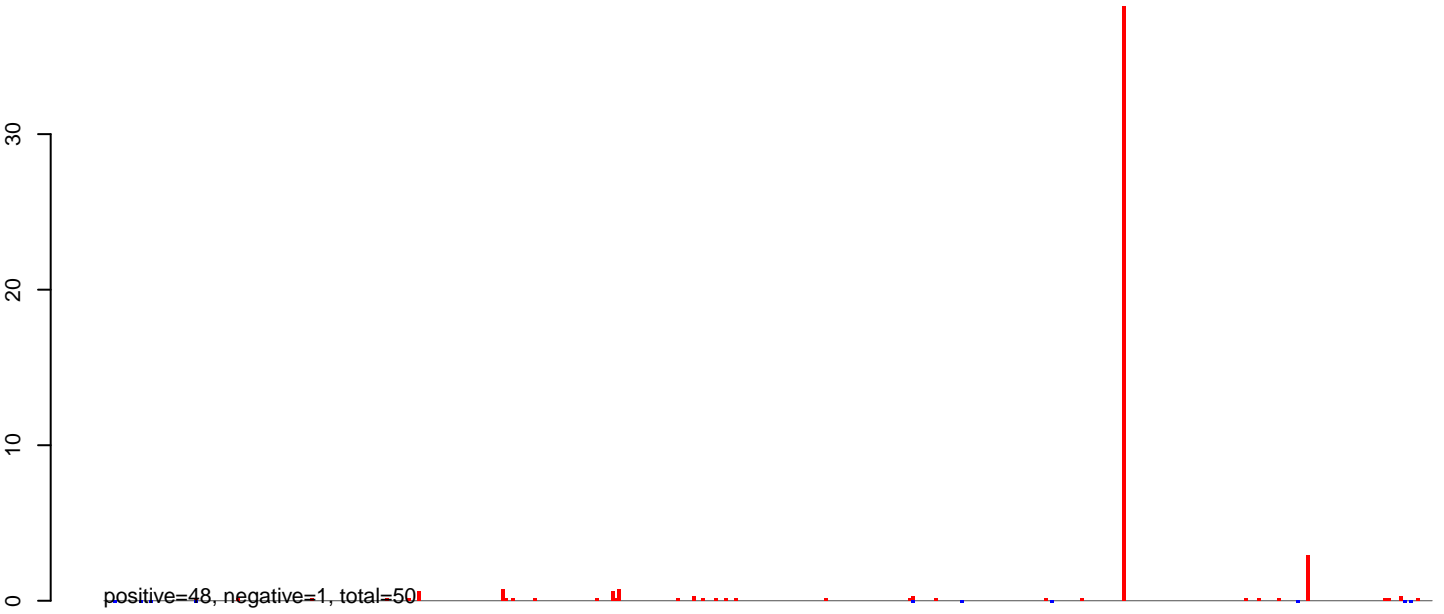
AeAeg_LVP_pMos3DBhspPBac.rep



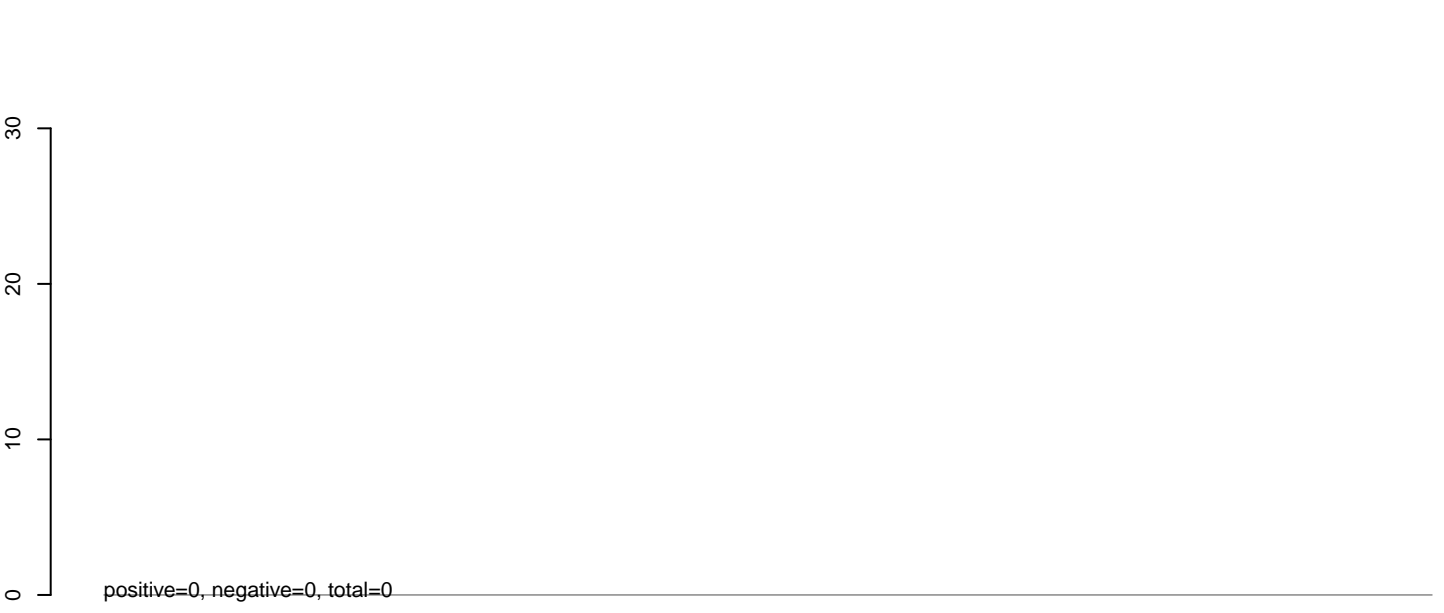
Window size=25, length=10215, virus@NC_033724.1-Kadam_virus_from_Uganda_polyprotein_gene:1-10215

0 2000 4000 6000 8000 10000

AeAeg_LVP_pMos3DBhspPBac.18_23.rep



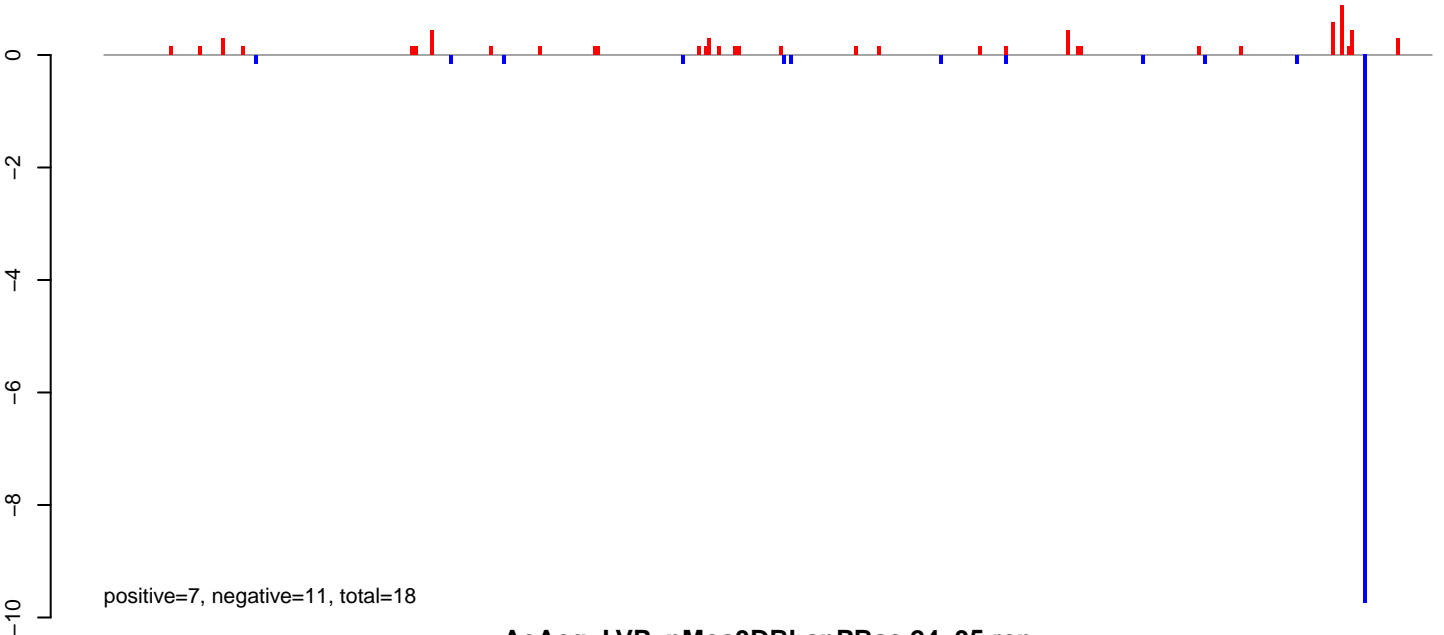
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



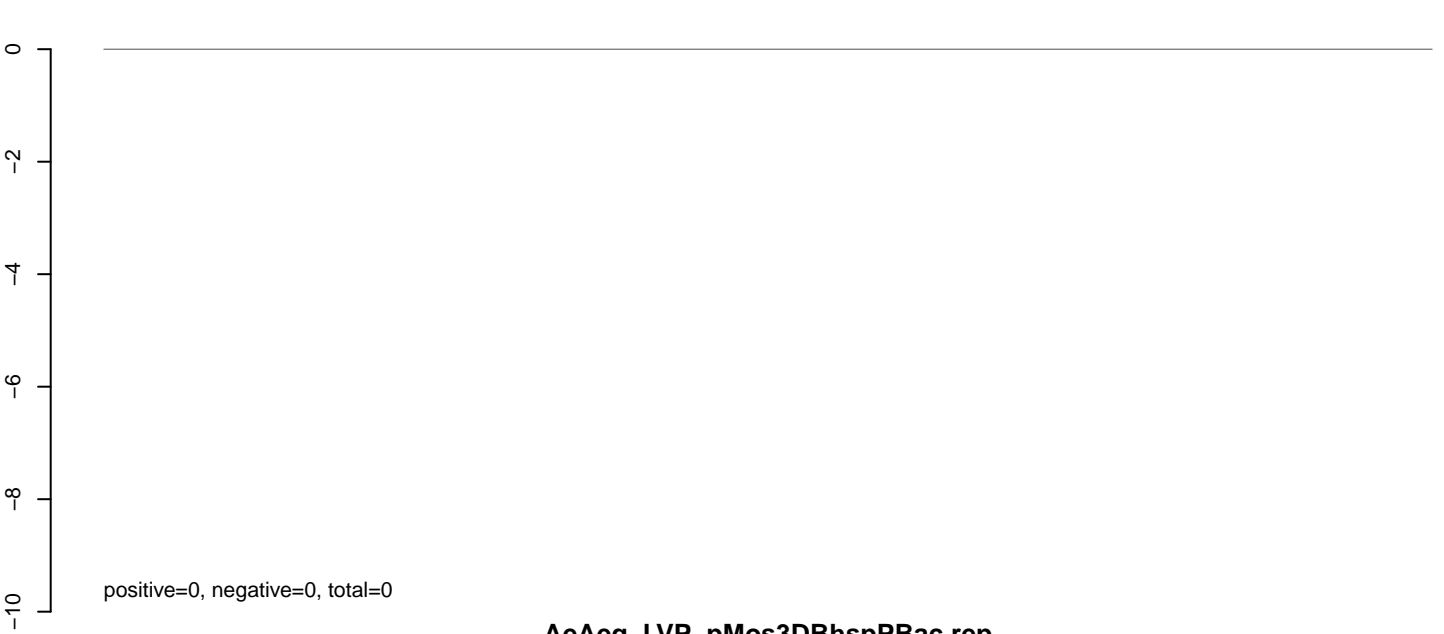
AeAeg_LVP_pMos3DBhspPBac.rep



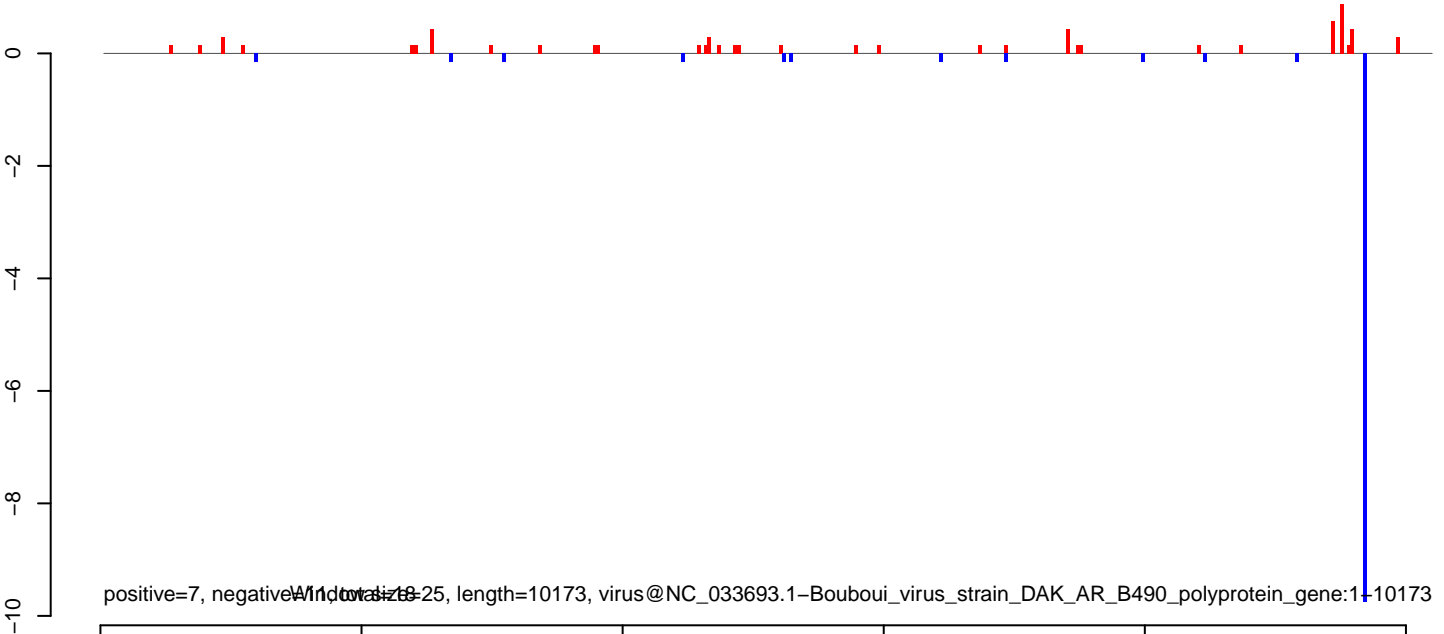
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

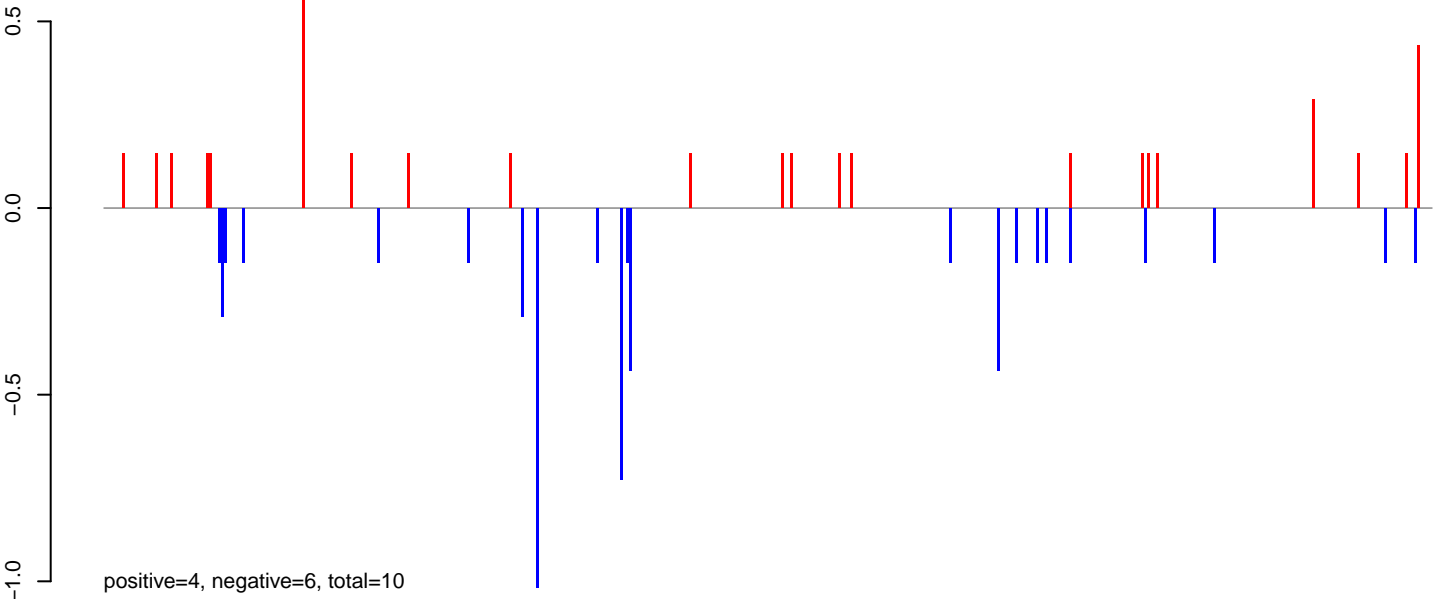


AeAeg_LVP_pMos3DBhspPBac.rep

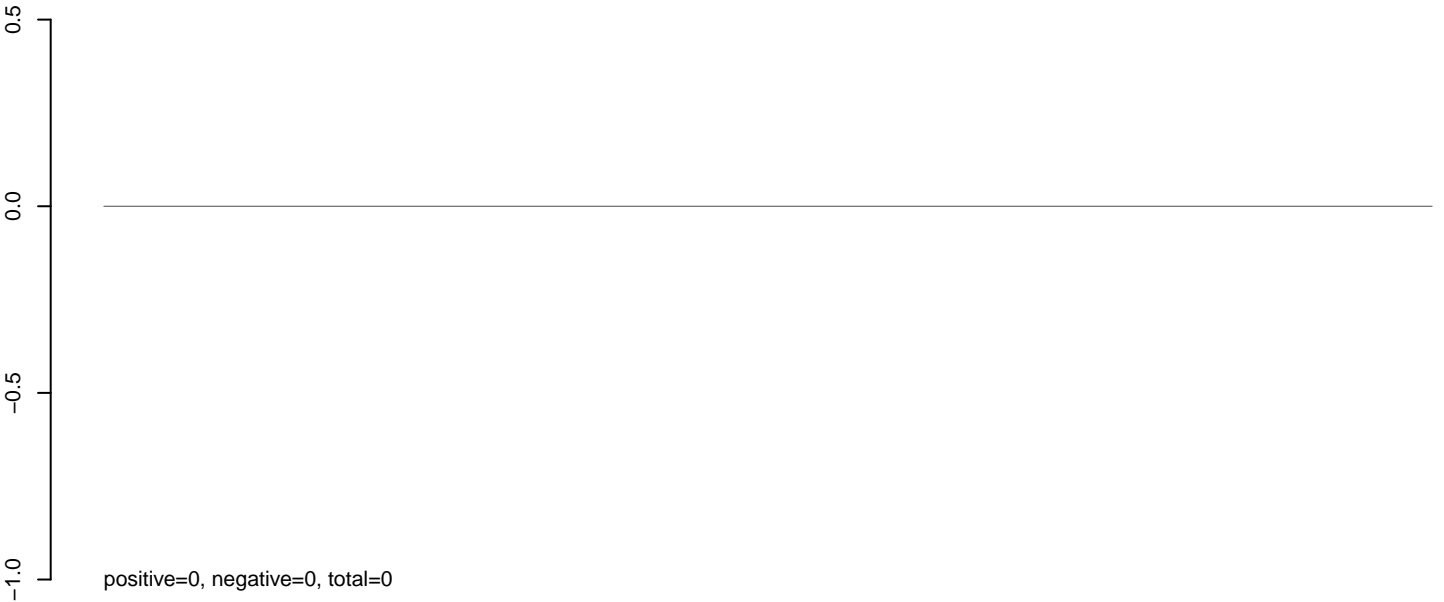


Window size=25, length=10173, virus@NC_033693.1-Bouboui_virus_strain_DAK_AR_B490_polyprotein_gene:1-10173

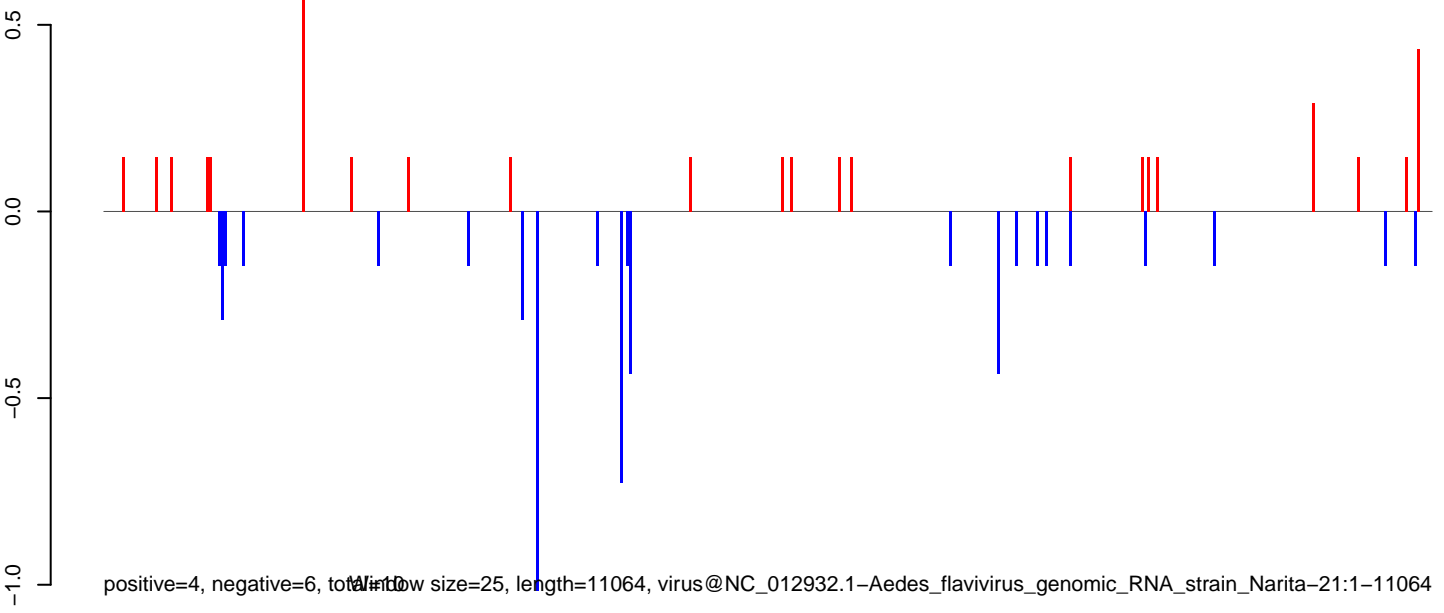
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



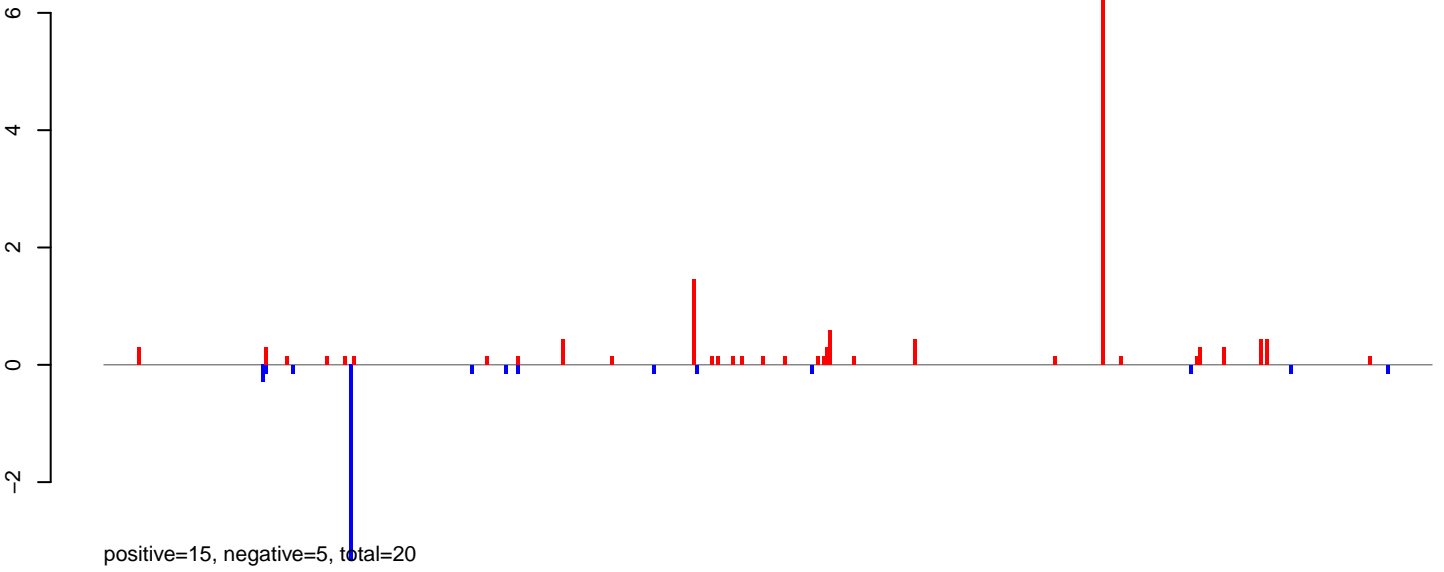
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



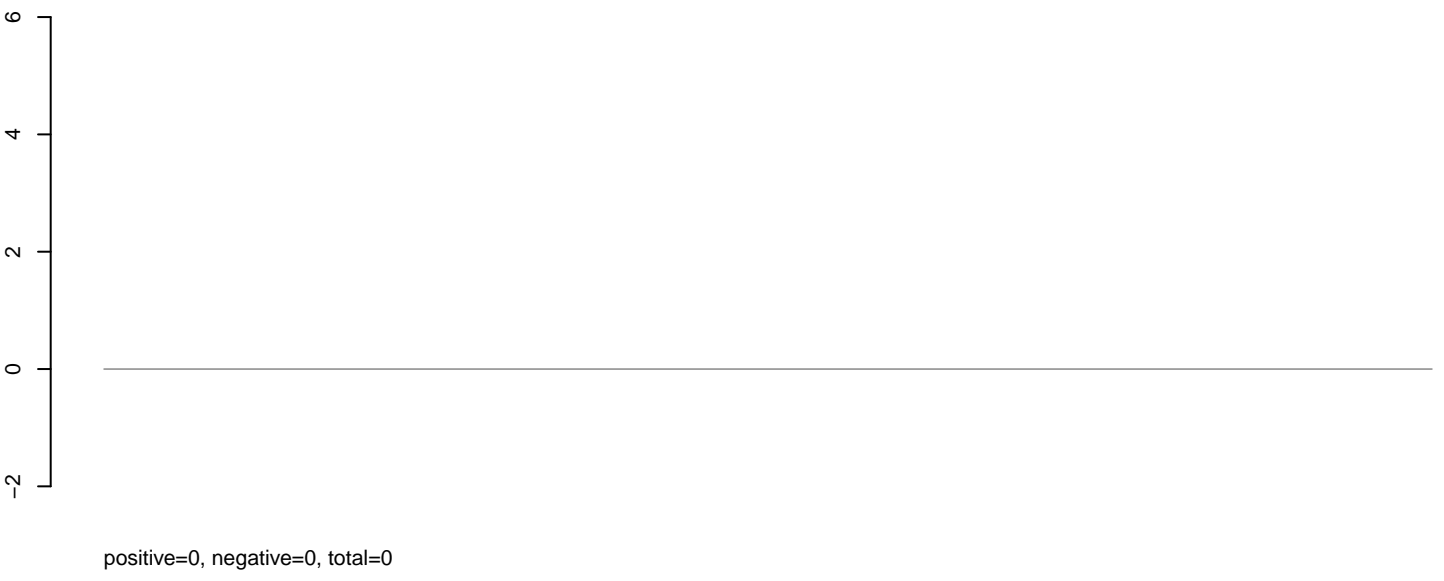
AeAeg_LVP_pMos3DBhspPBac.rep



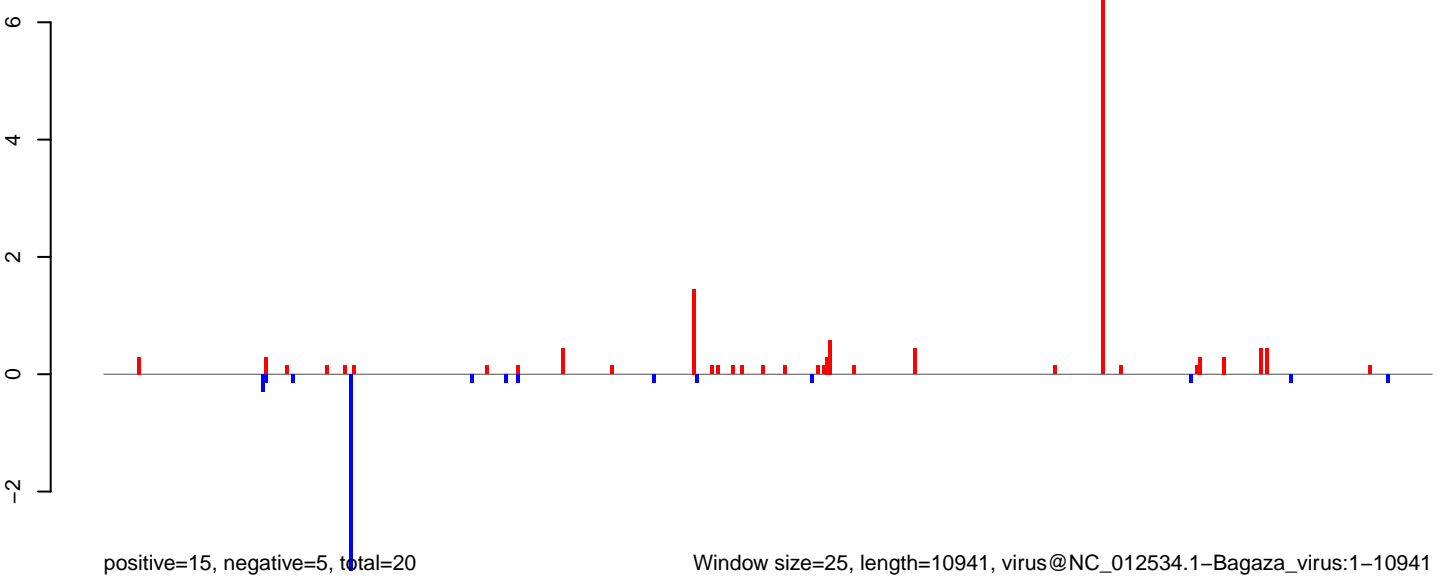
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



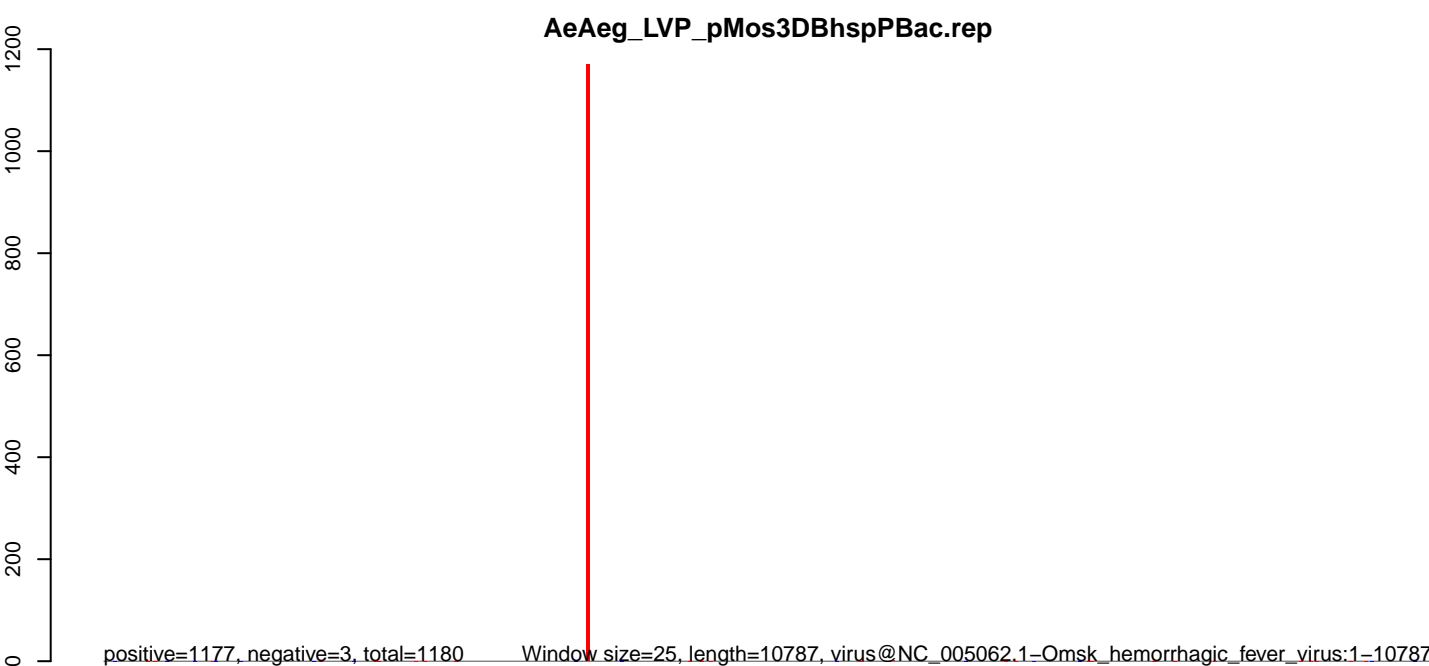
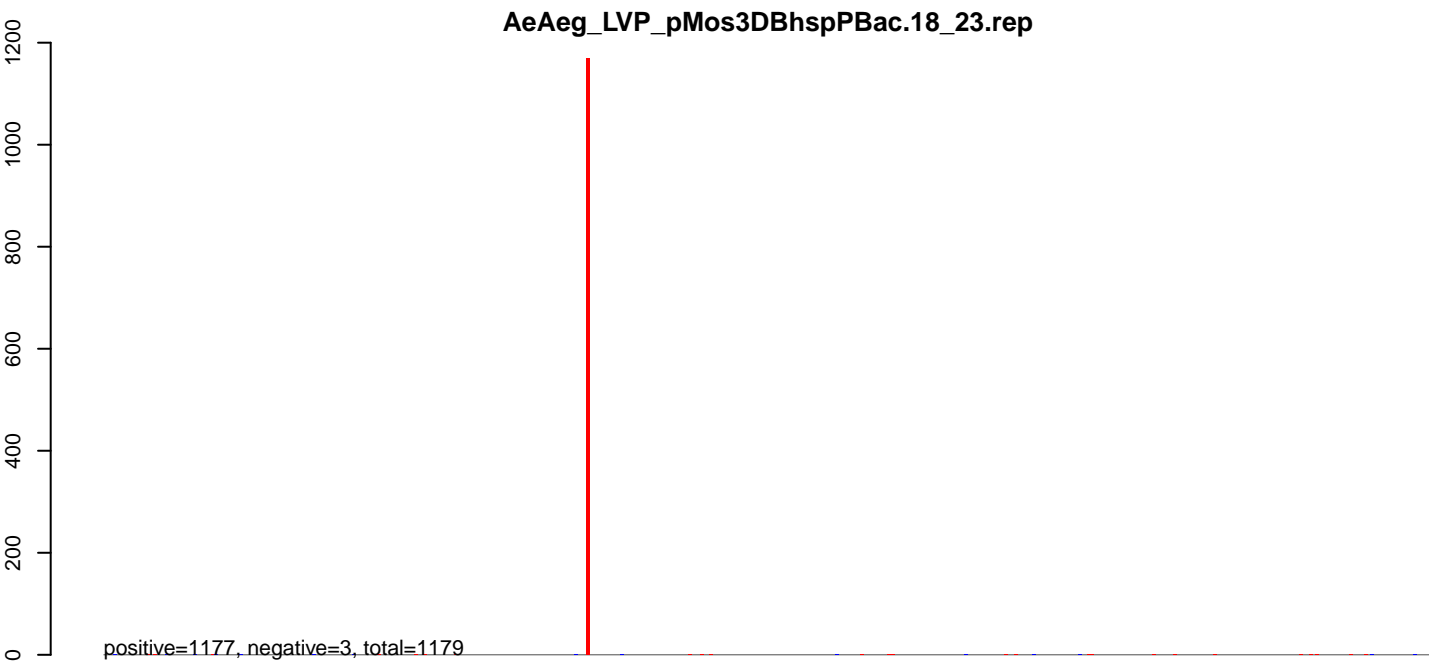
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



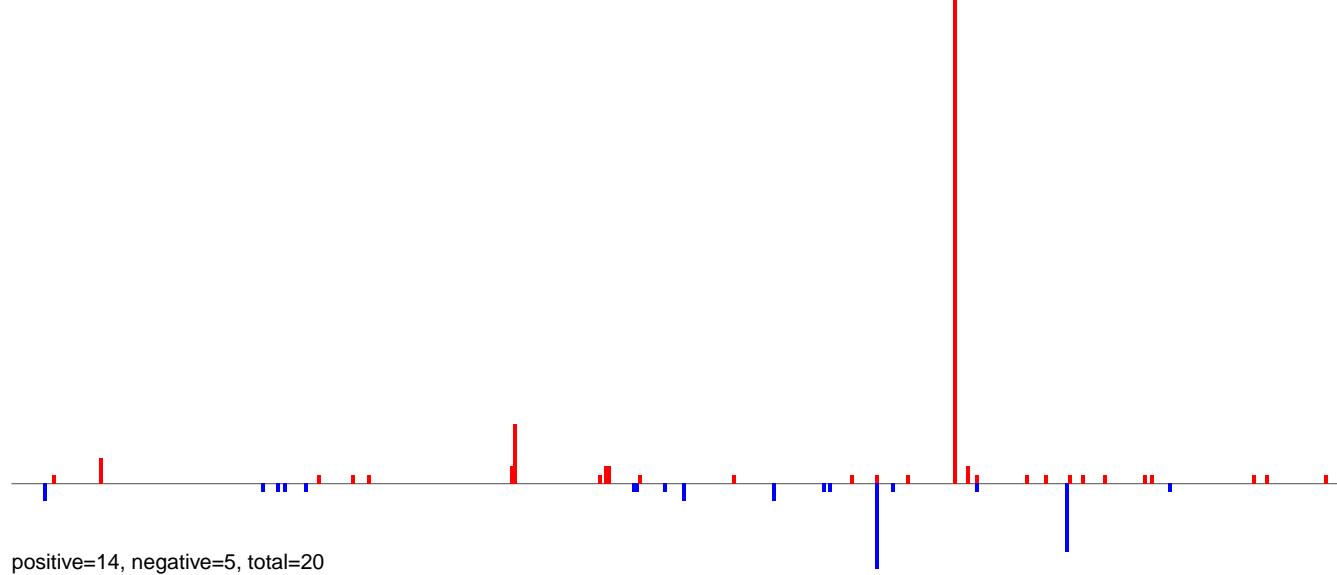
AeAeg_LVP_pMos3DBhspPBac.rep



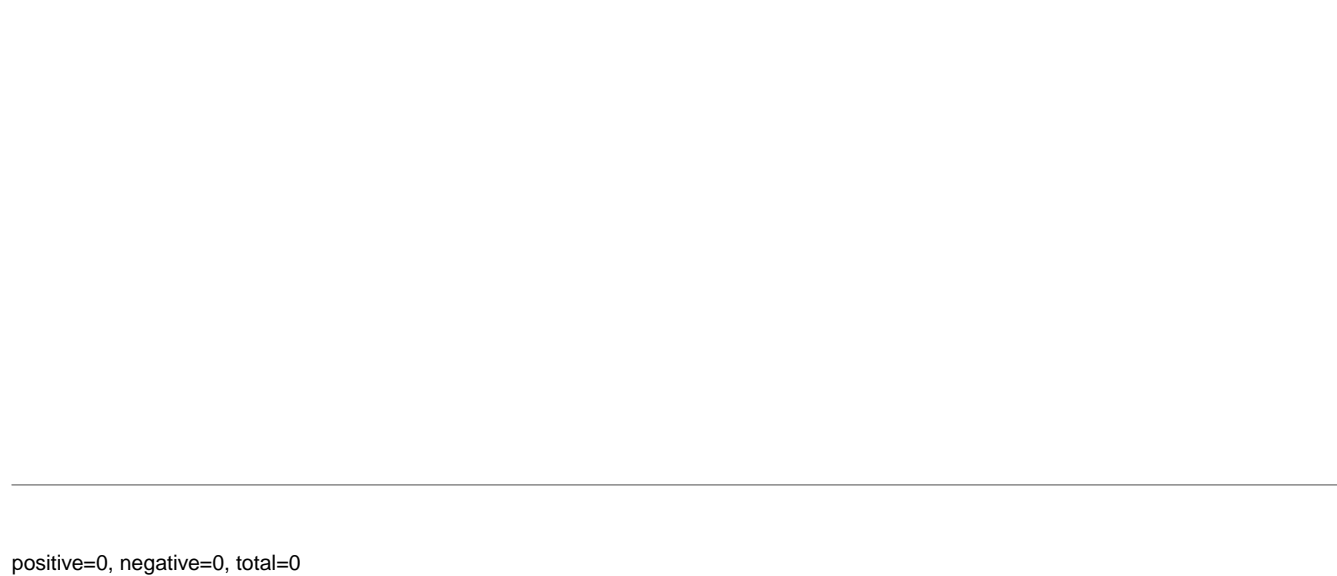
Window size=25, length=10941, virus@NC_012534.1-Bagaza_virus:1-10941



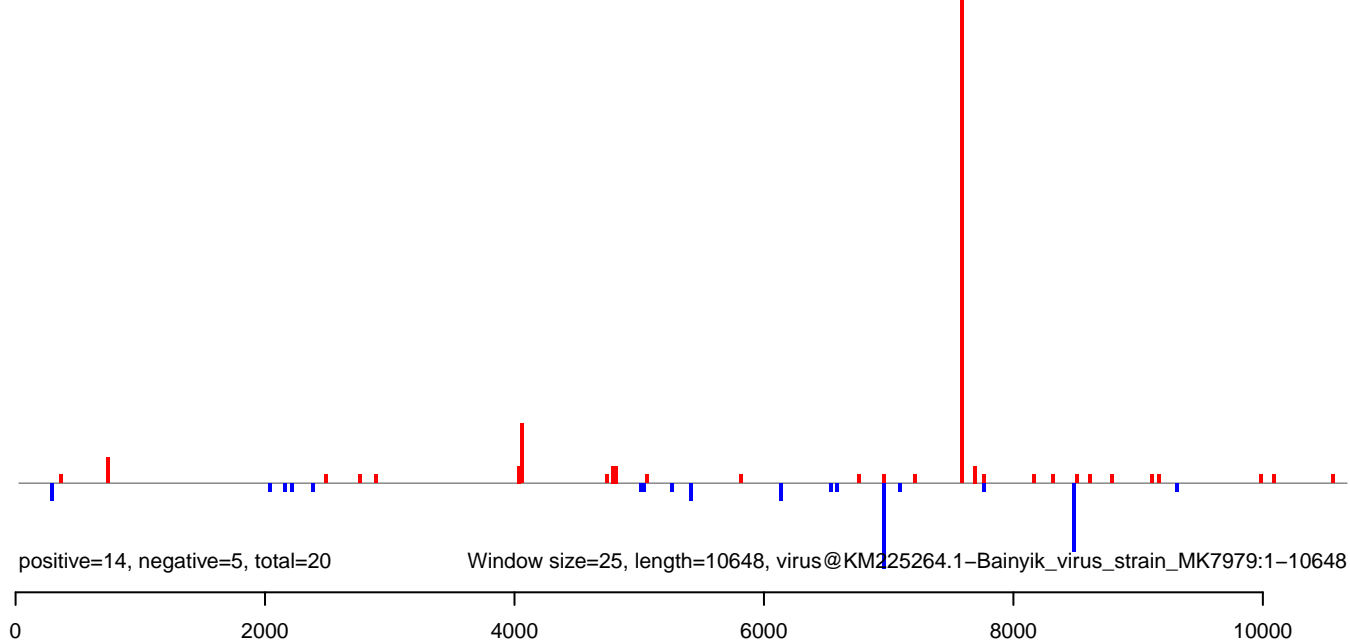
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



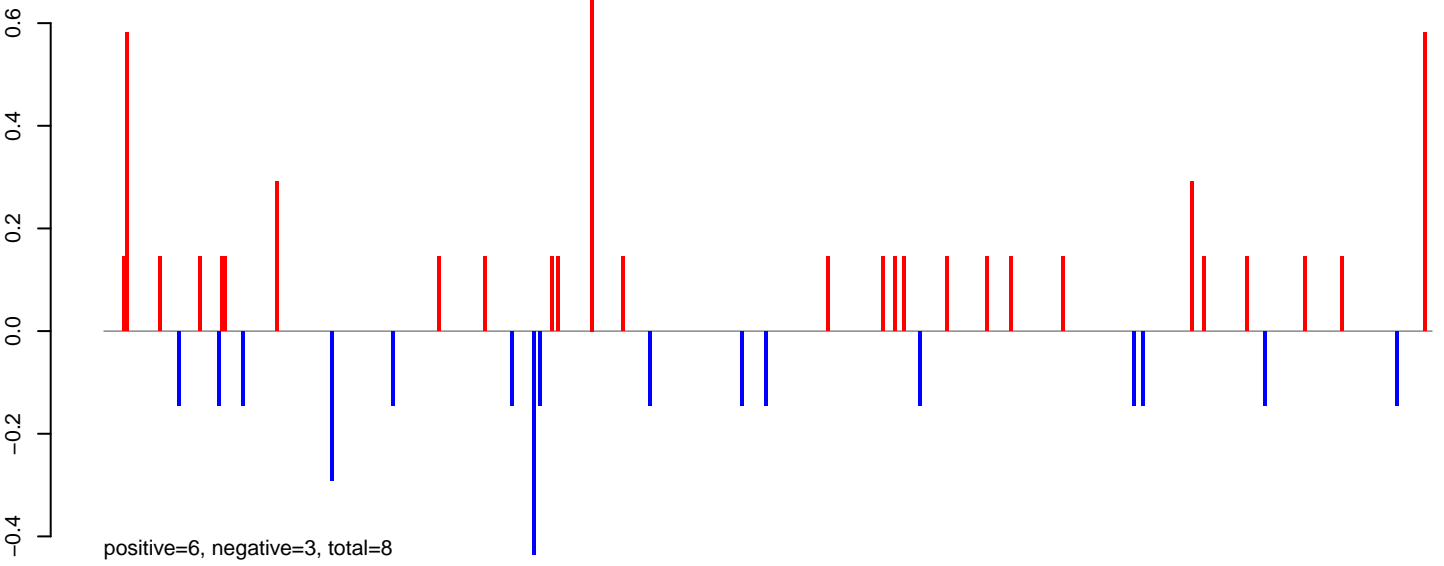
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



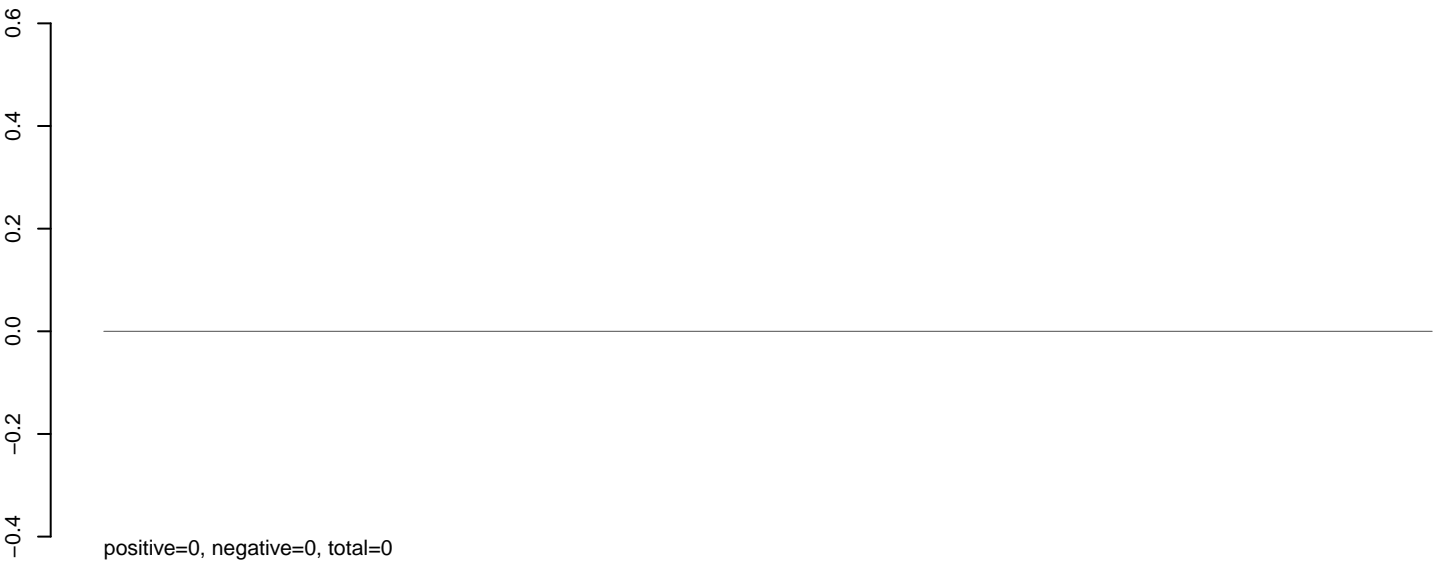
AeAeg_LVP_pMos3DBhspPBac.rep



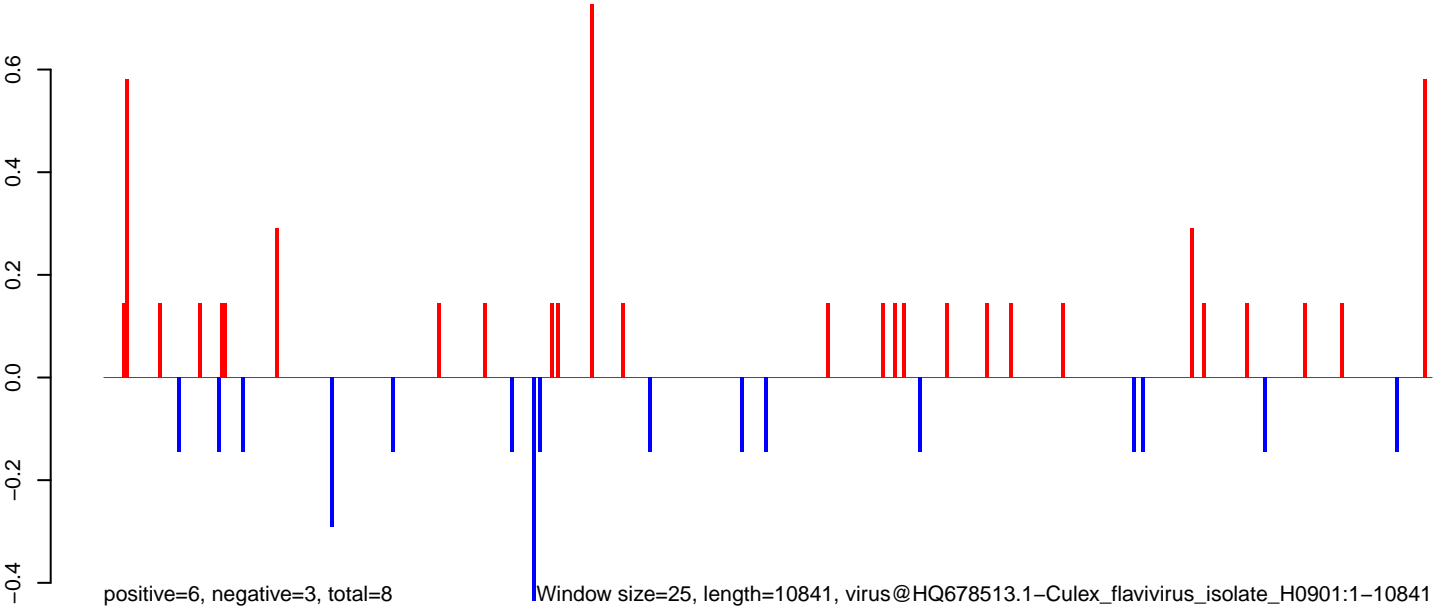
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

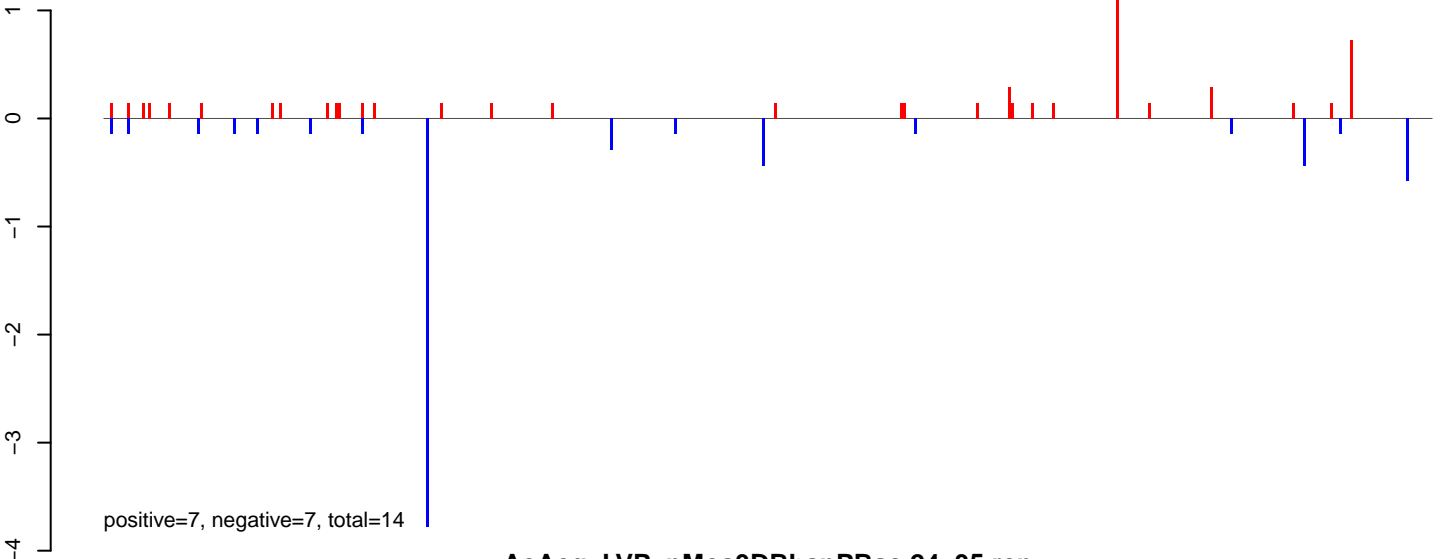


AeAeg_LVP_pMos3DBhspPBac.rep

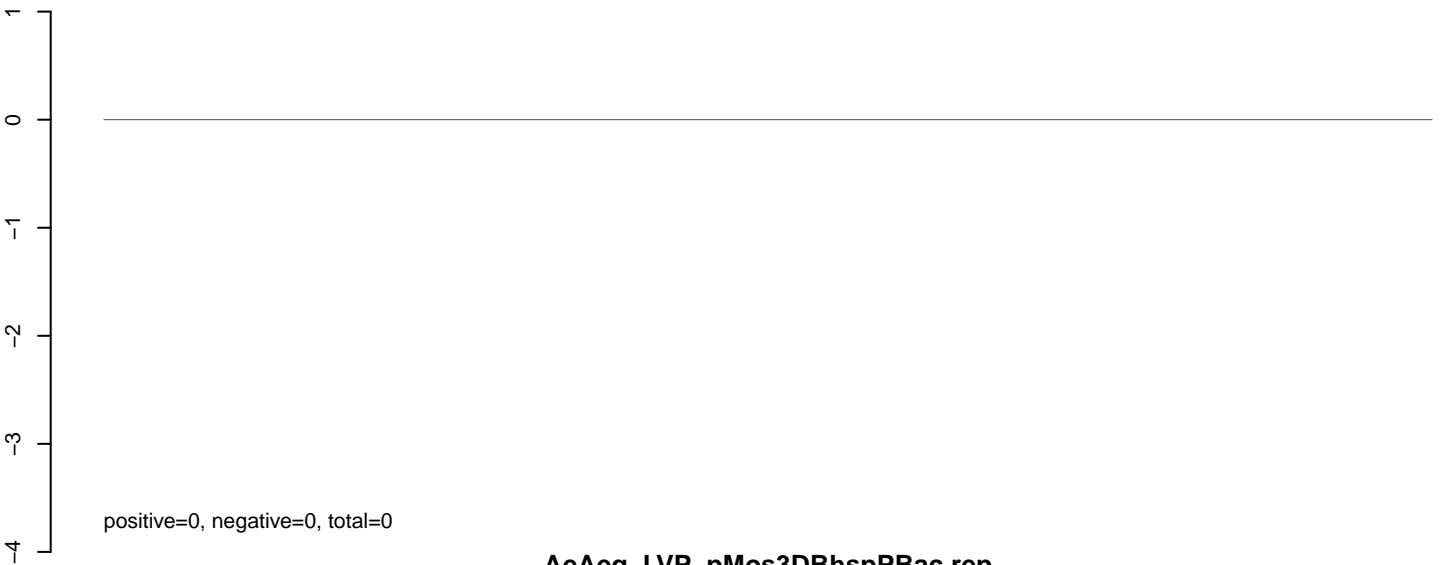


0 2000 4000 6000 8000 10000

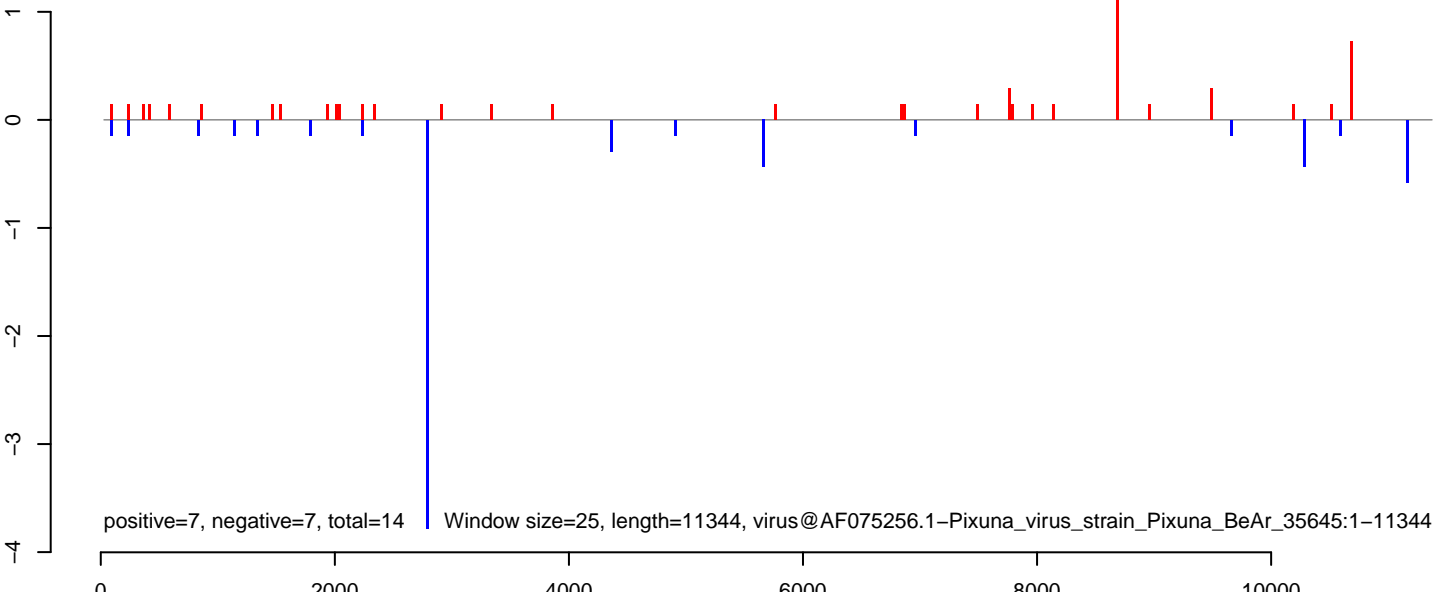
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



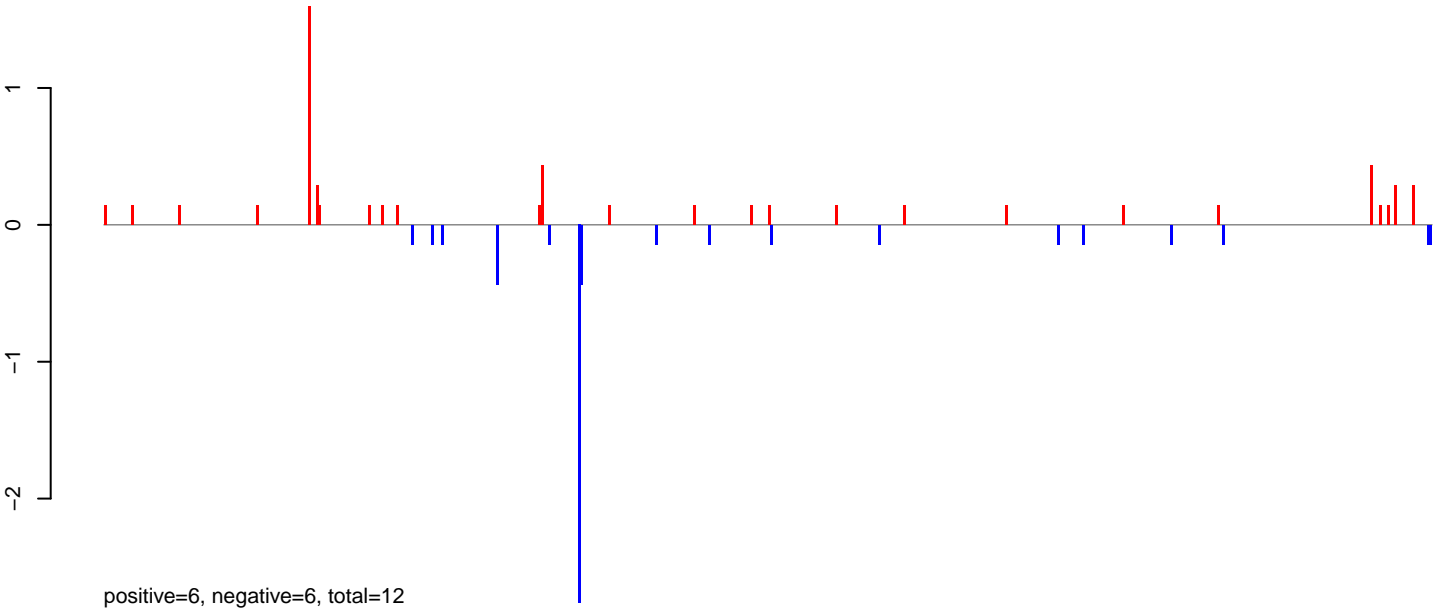
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



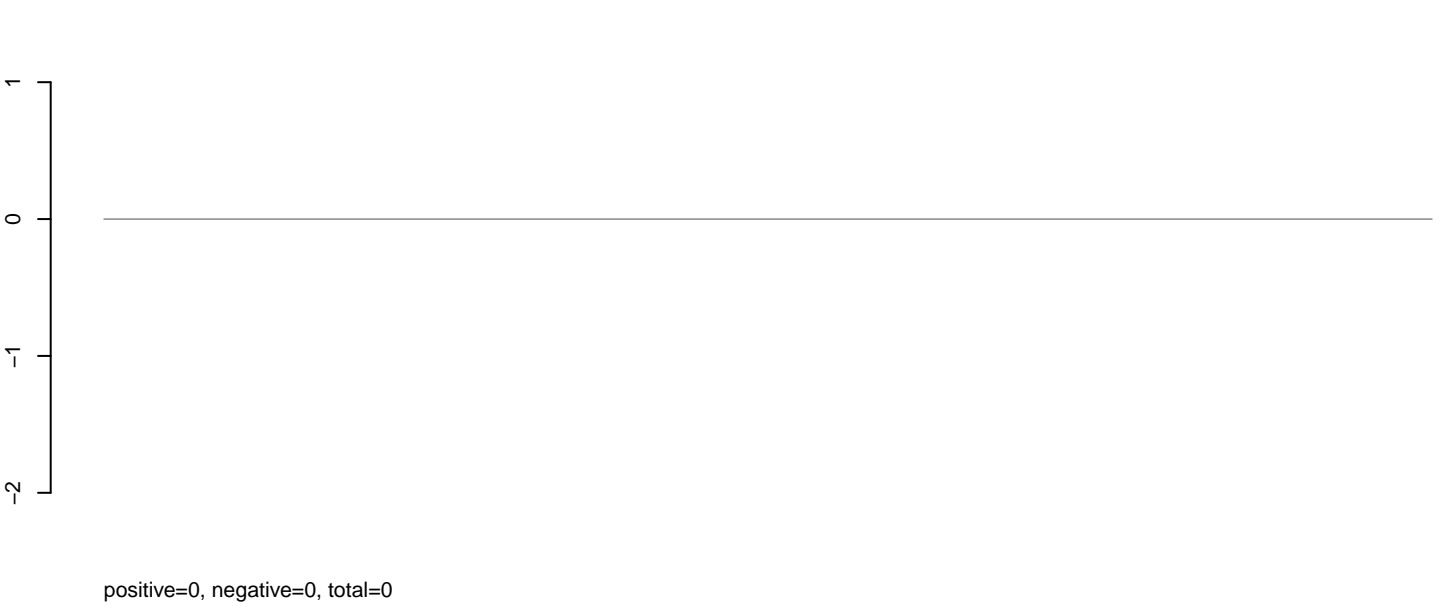
AeAeg_LVP_pMos3DBhspPBac.rep



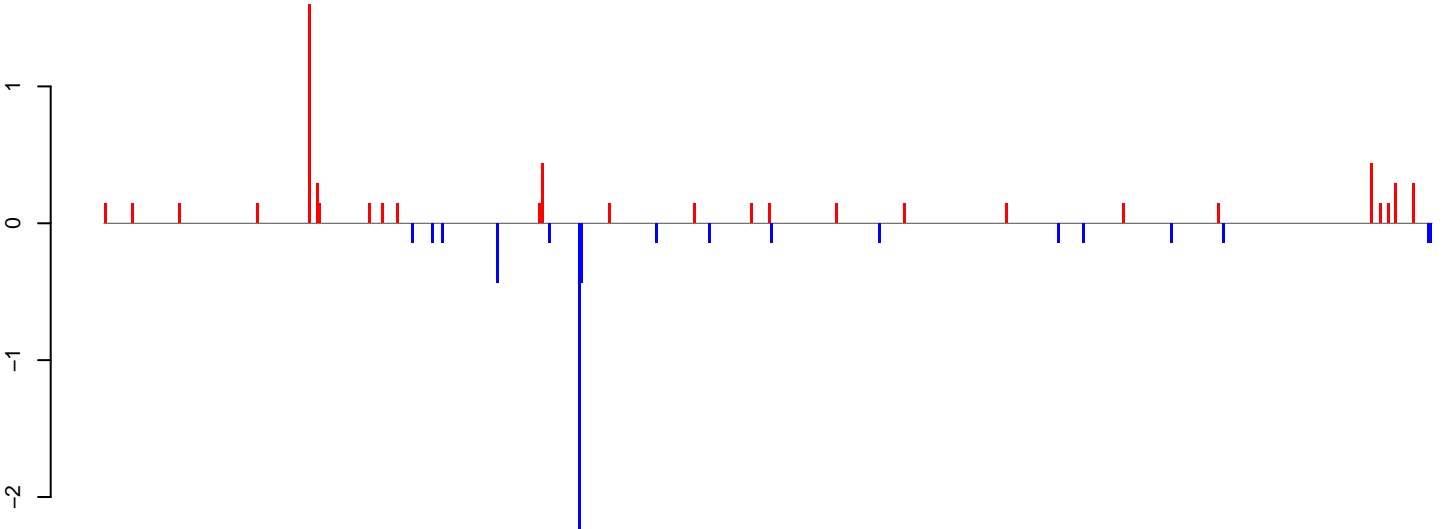
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



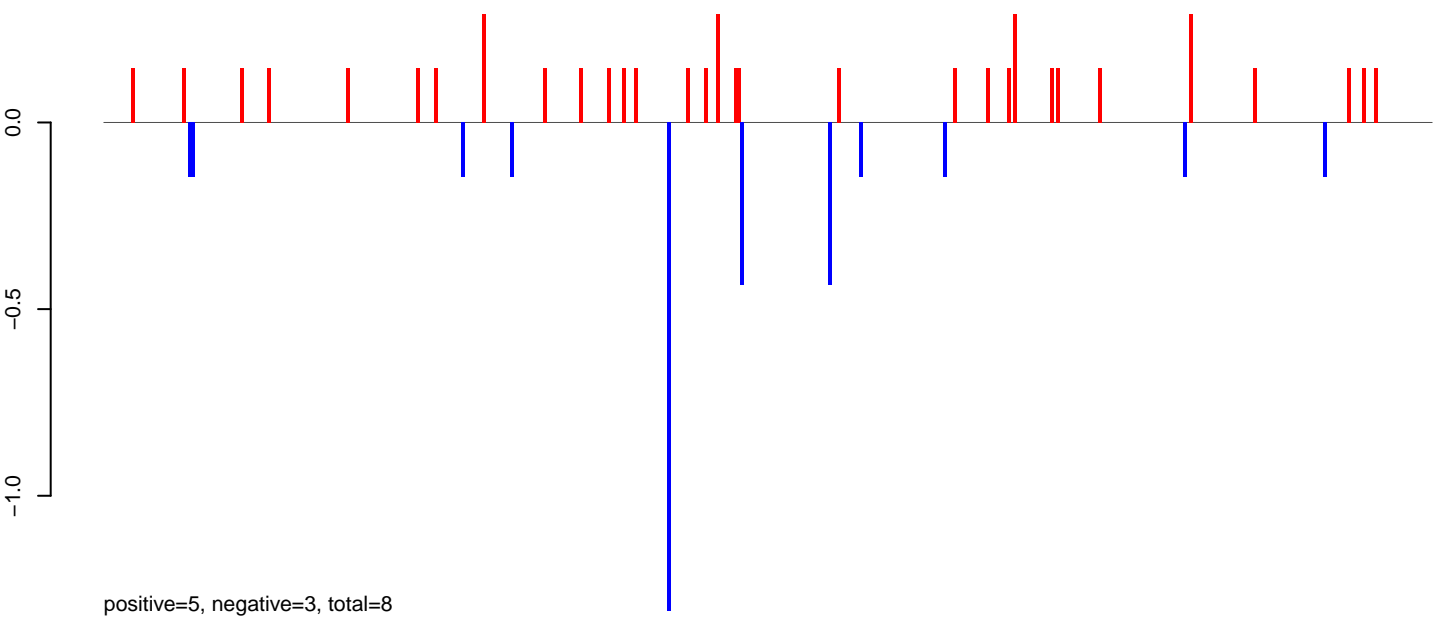
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



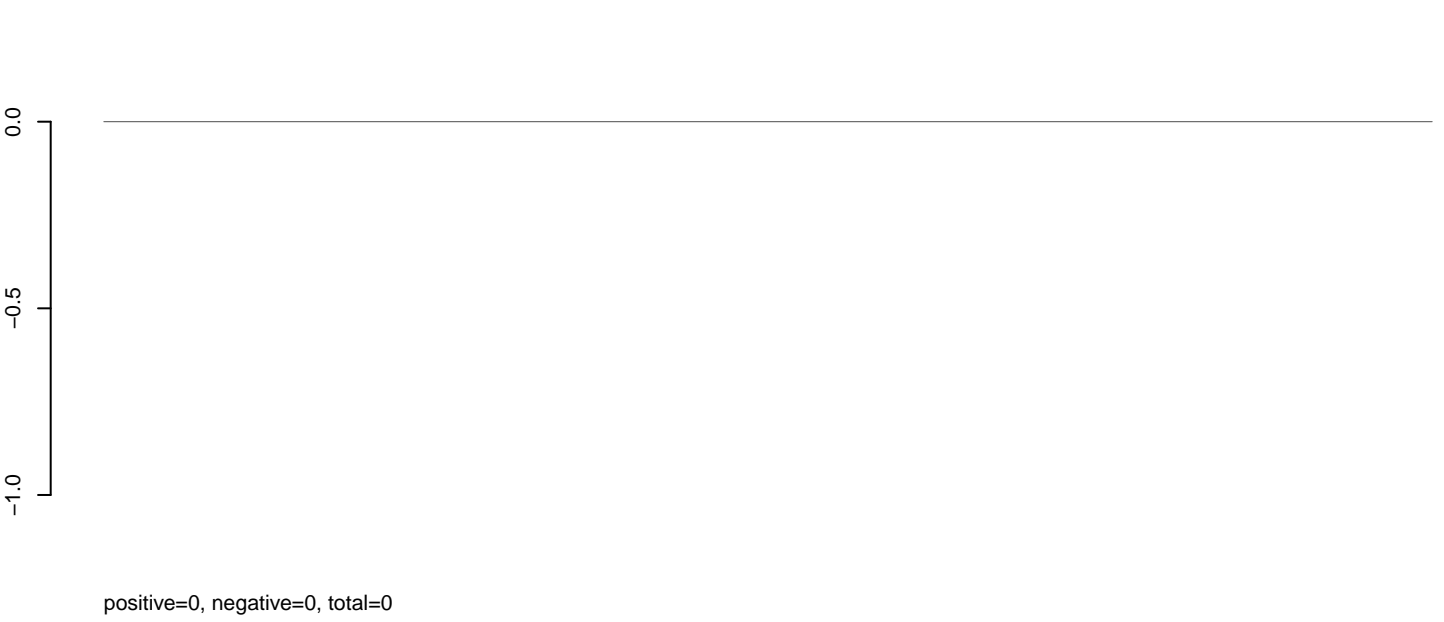
AeAeg_LVP_pMos3DBhspPBac.rep



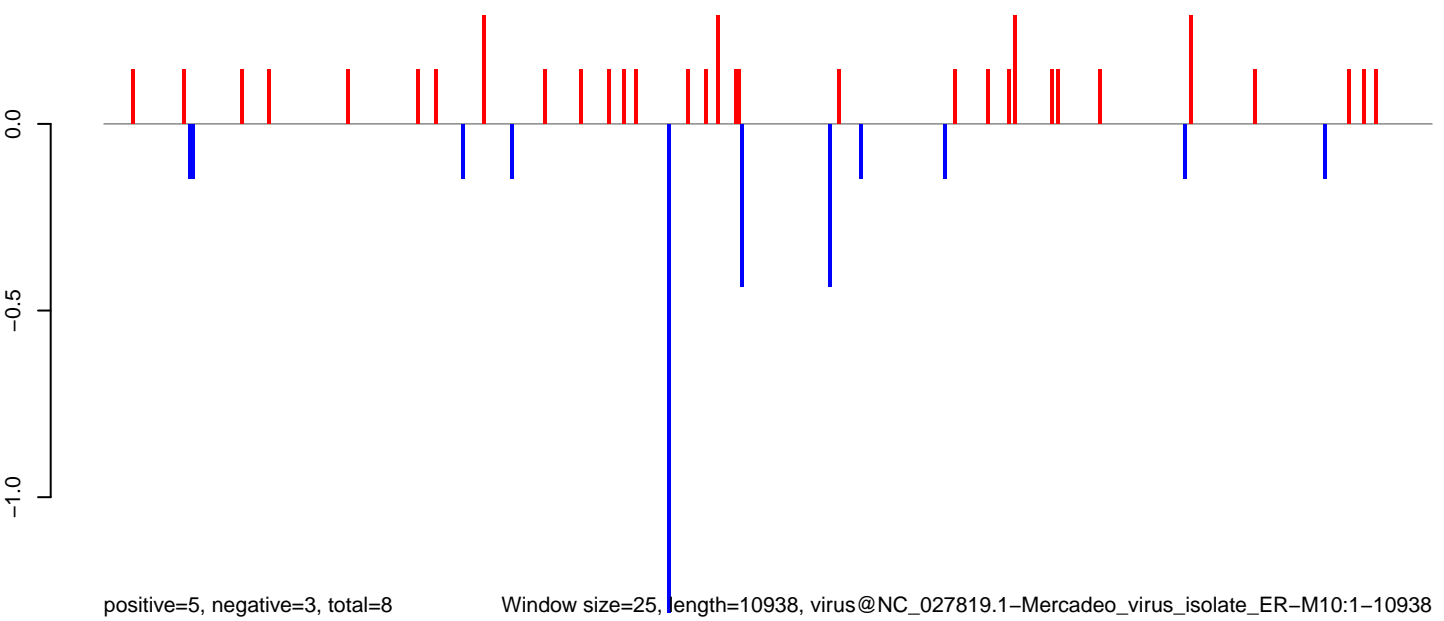
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



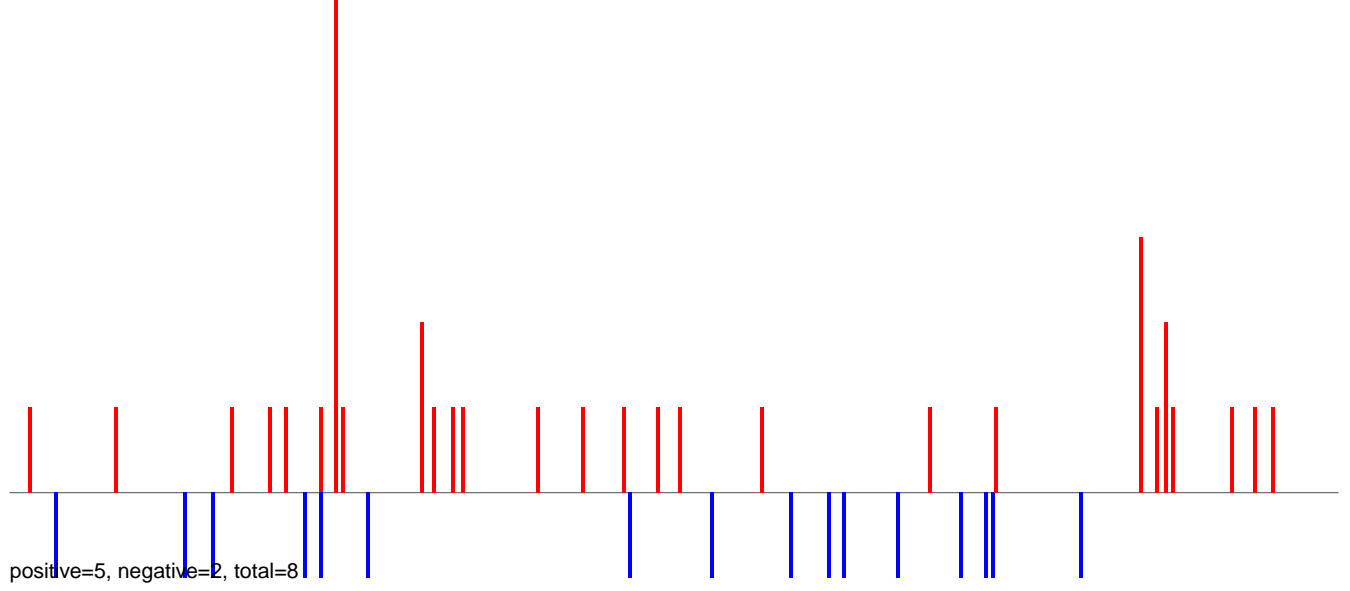
AeAeg_LVP_pMos3DBhspPBac.rep



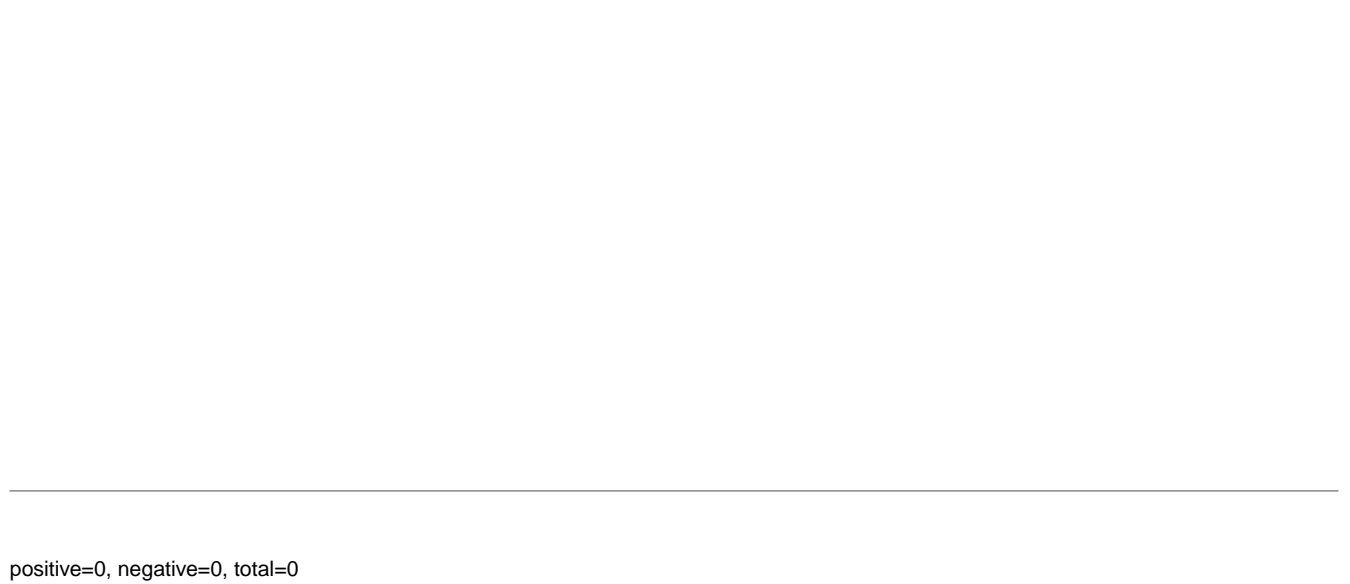
Window size=25, length=10938, virus@NC_027819.1-Mercadeo_virus_isolate_ER-M10:1-10938

0 2000 4000 6000 8000 10000

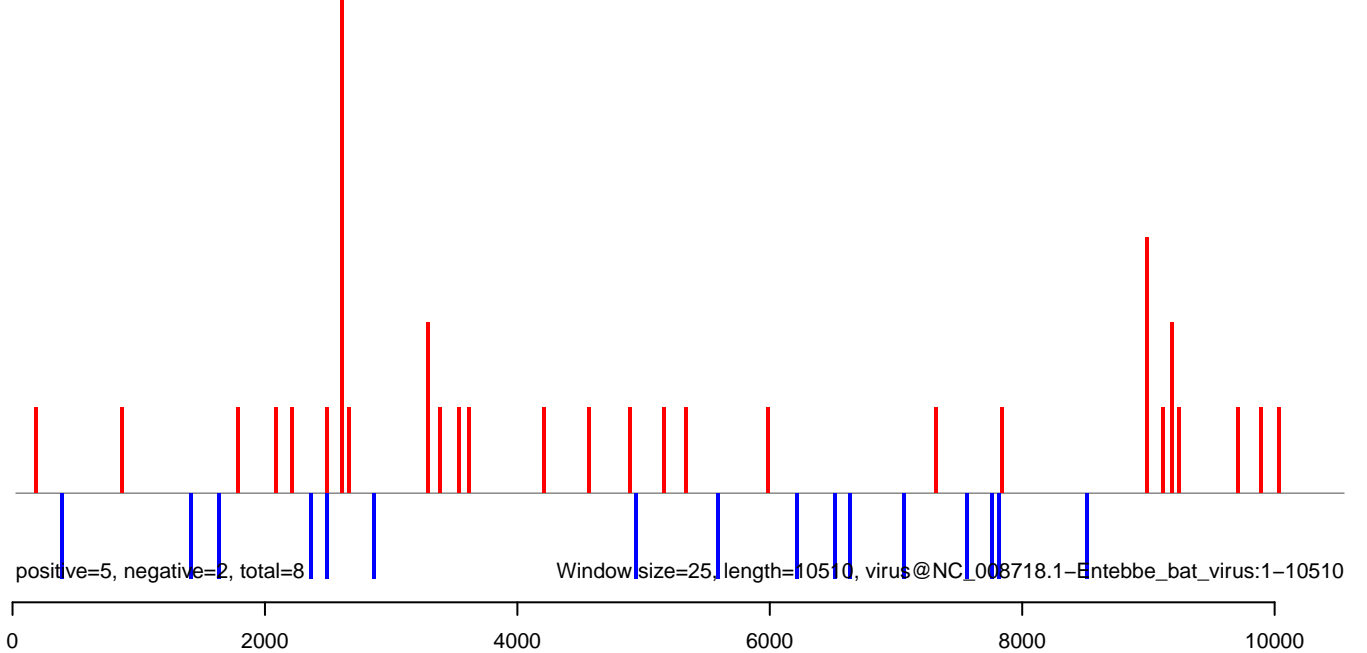
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



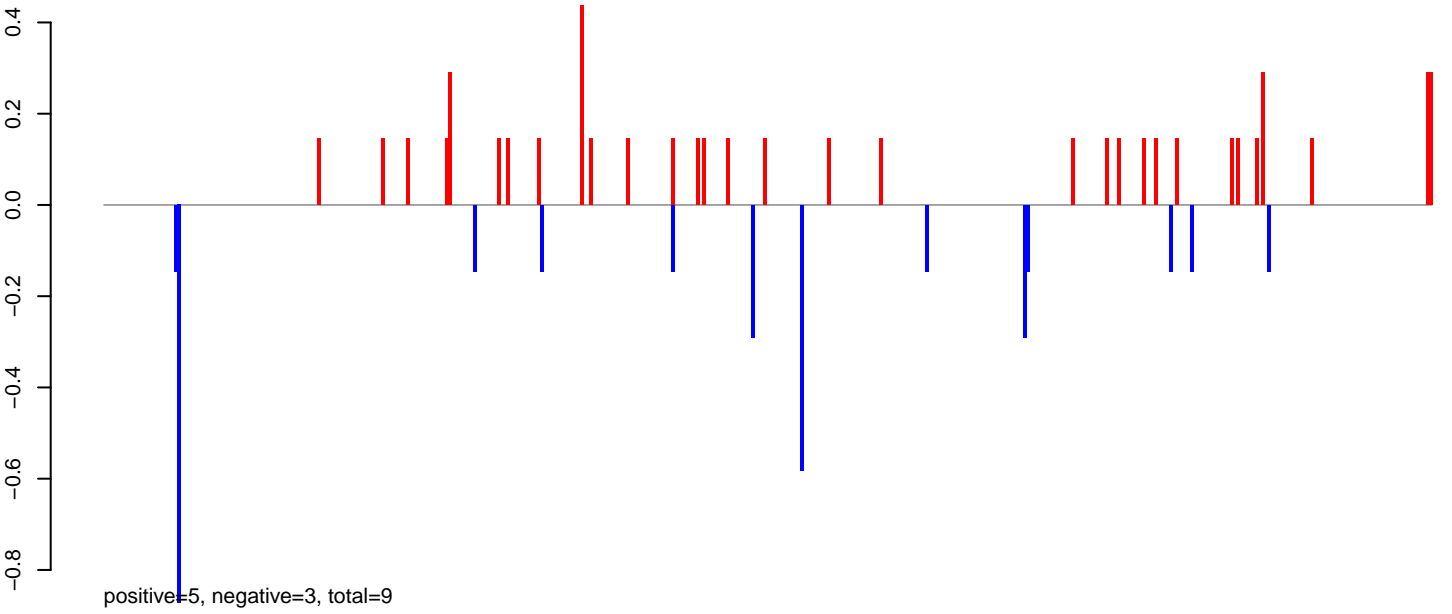
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



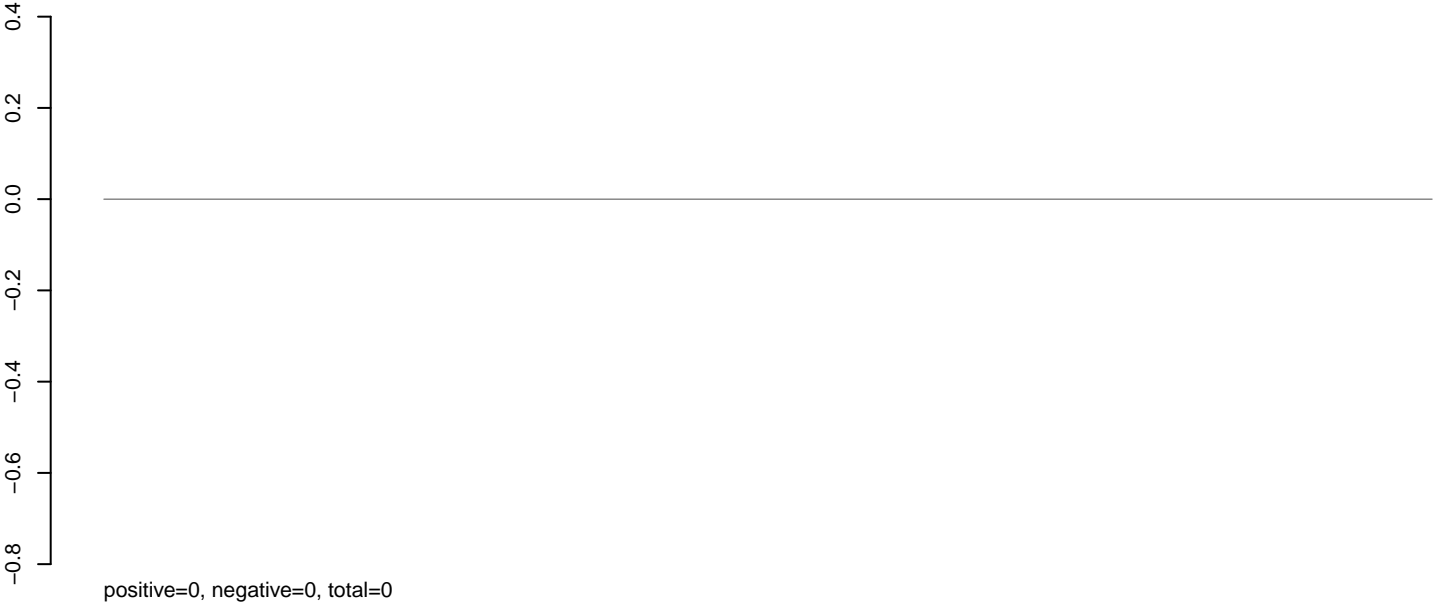
AeAeg_LVP_pMos3DBhspPBac.rep



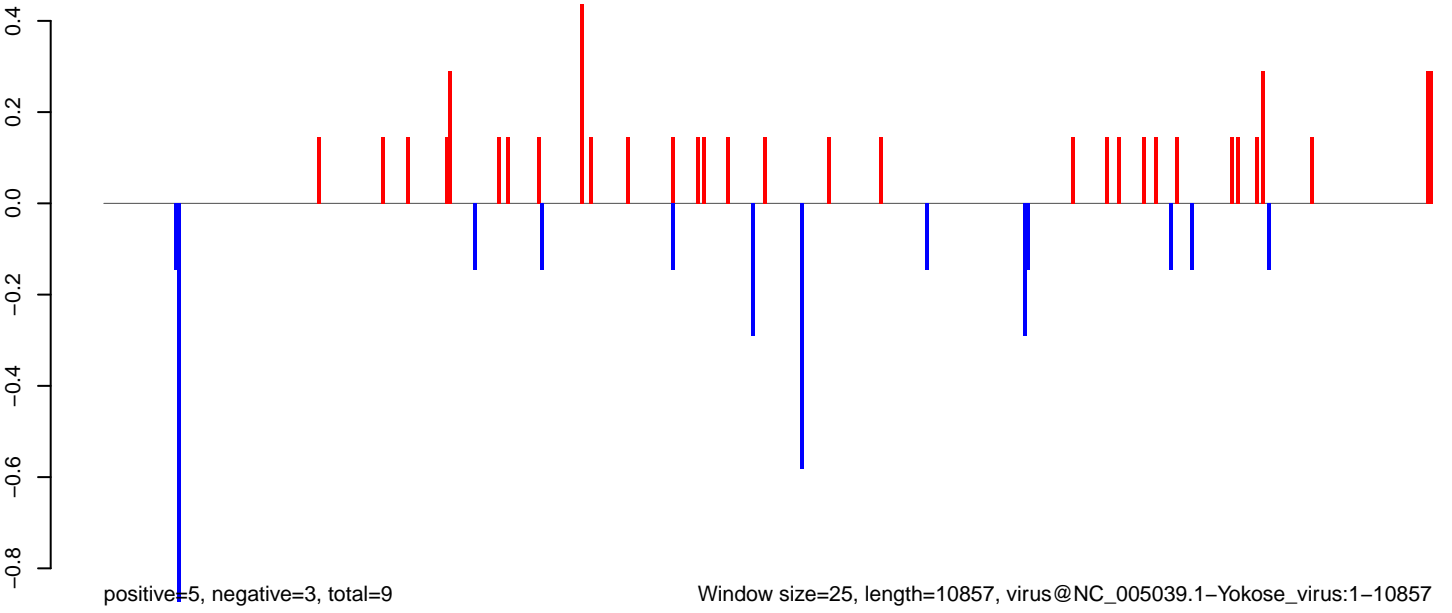
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



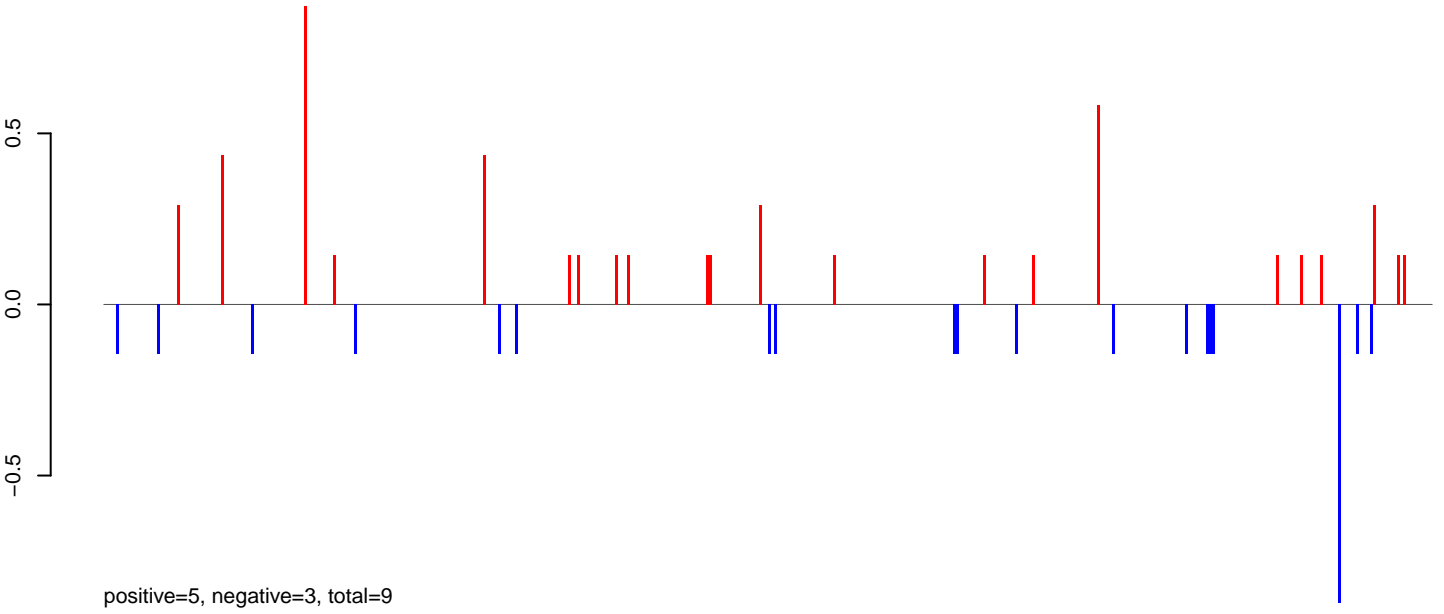
AeAeg_LVP_pMos3DBhspPBac.rep



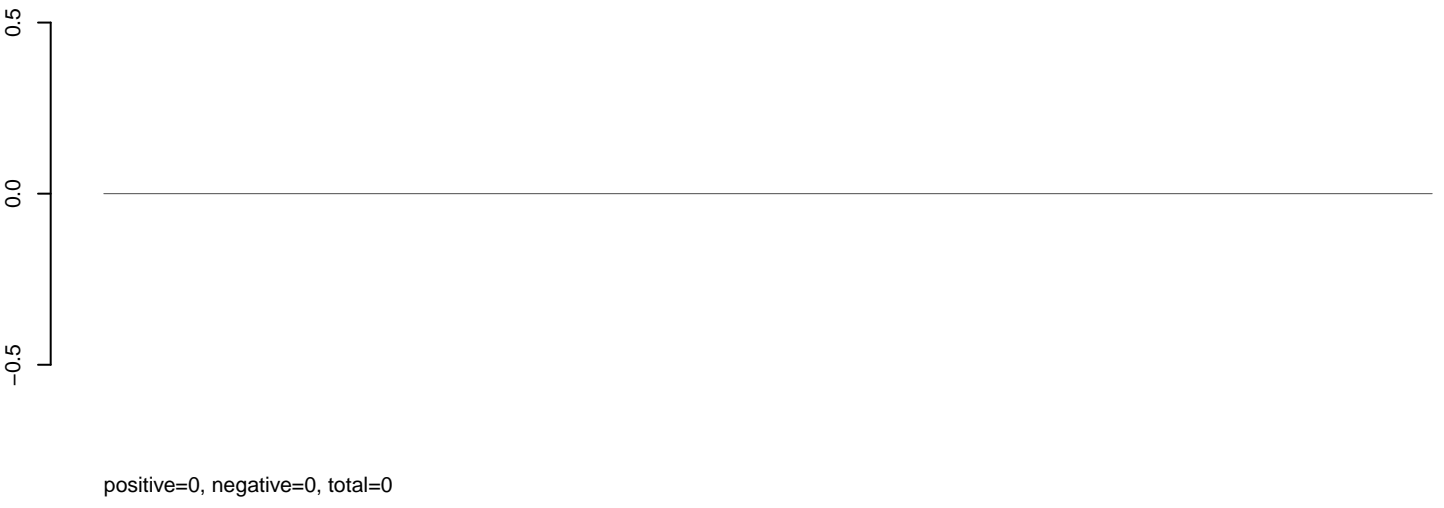
Window size=25, length=10857, virus@NC_005039.1-Yokose_virus:1-10857

0 2000 4000 6000 8000 10000

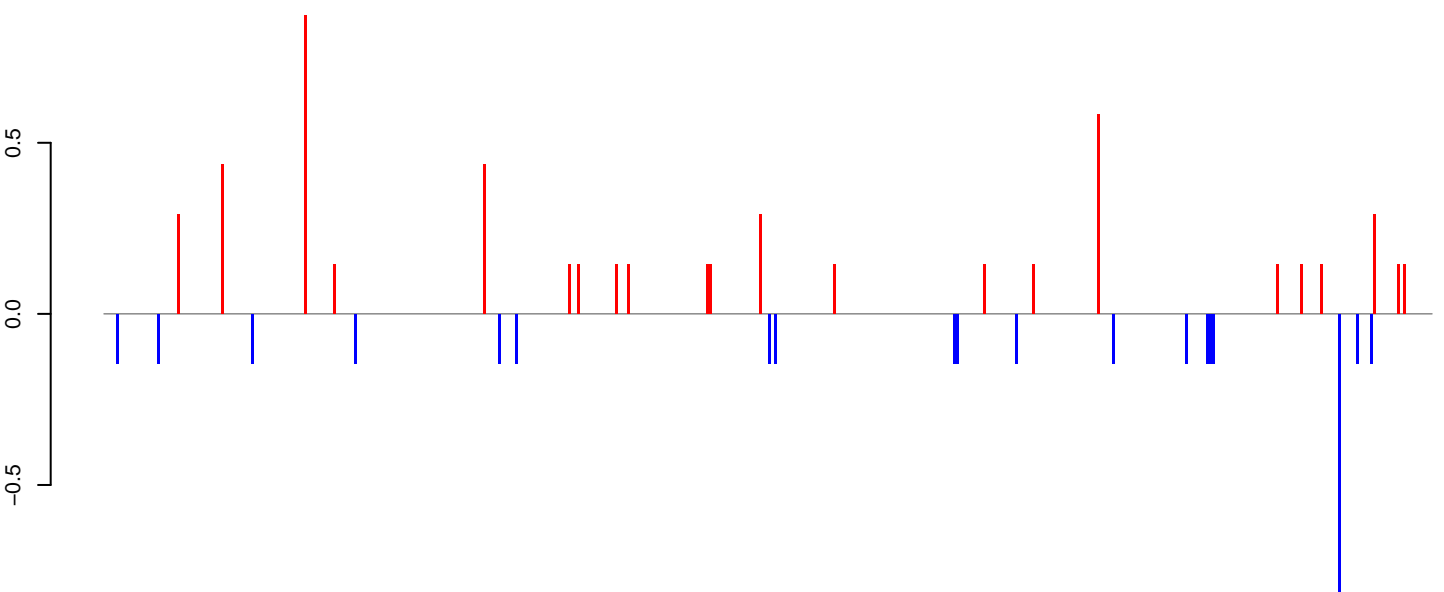
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



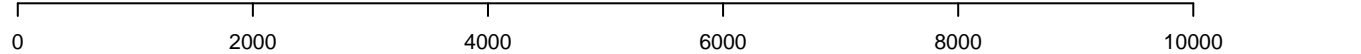
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



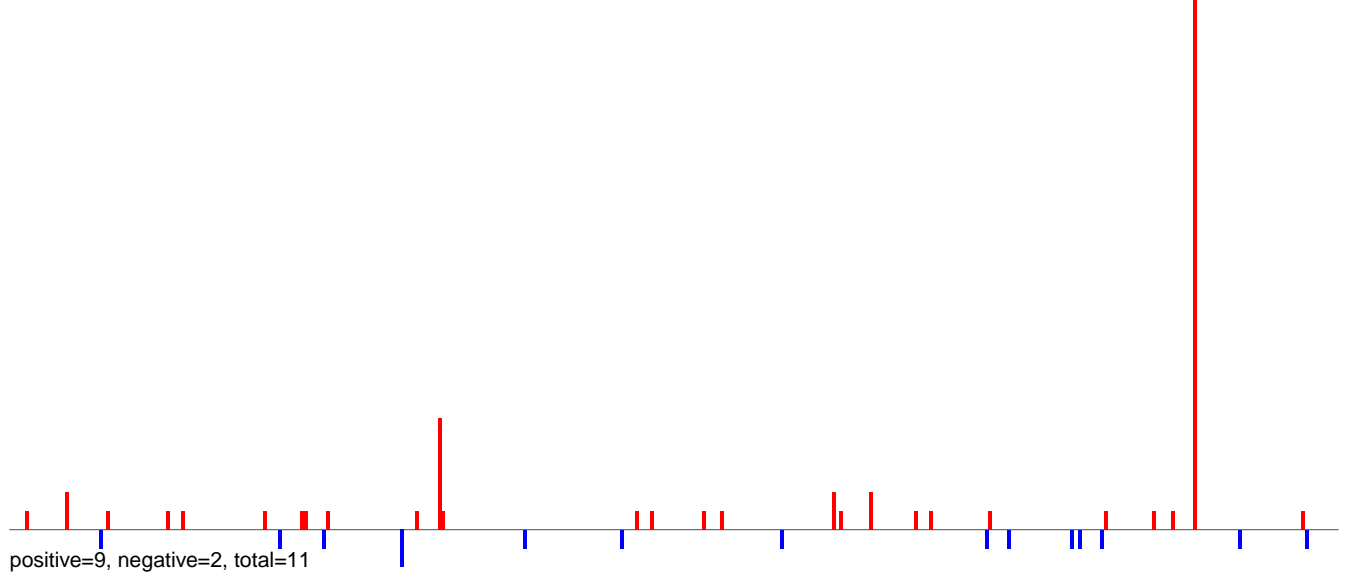
AeAeg_LVP_pMos3DBhspPBac.rep



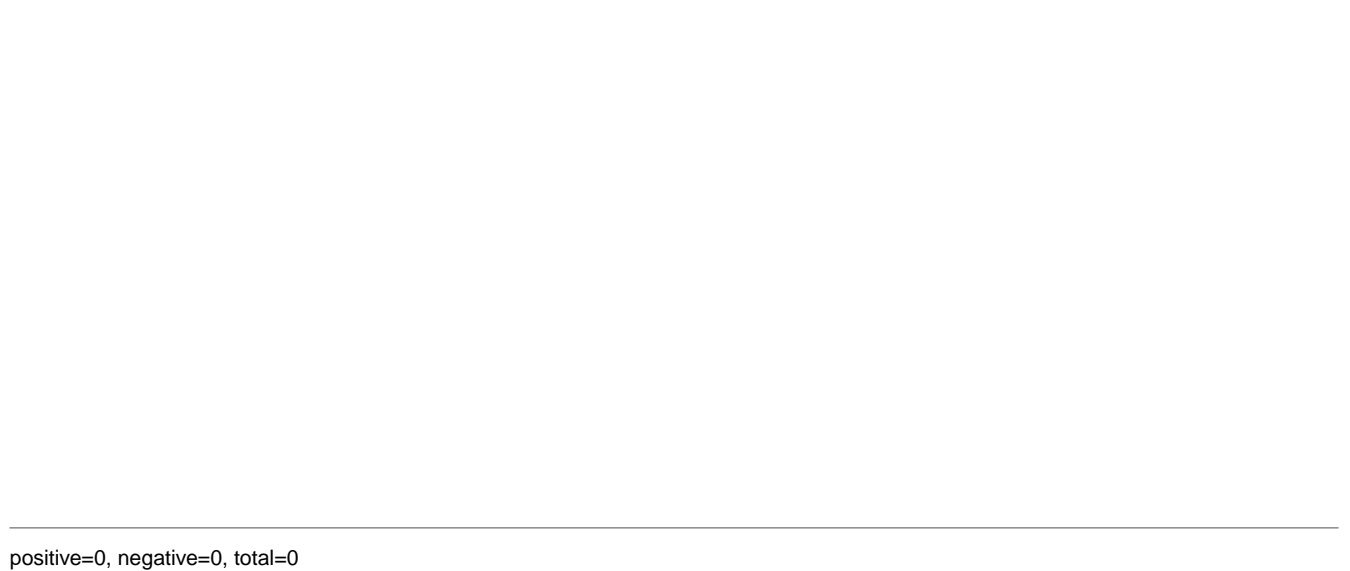
25, length=1280, negative=3, total=9, 15.1-Hubei_chuvirus-like_virus_3_strain_QTM26698_RNA-dependent_RNA_polymerase_putative_1-11280



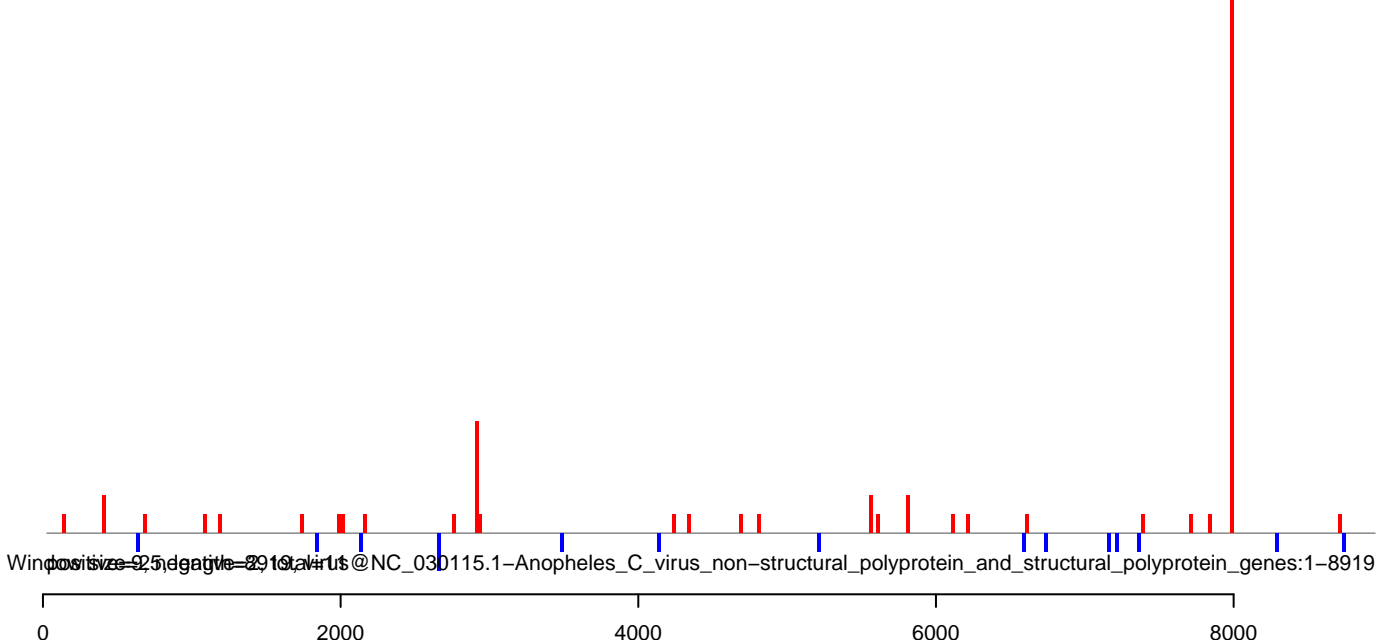
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



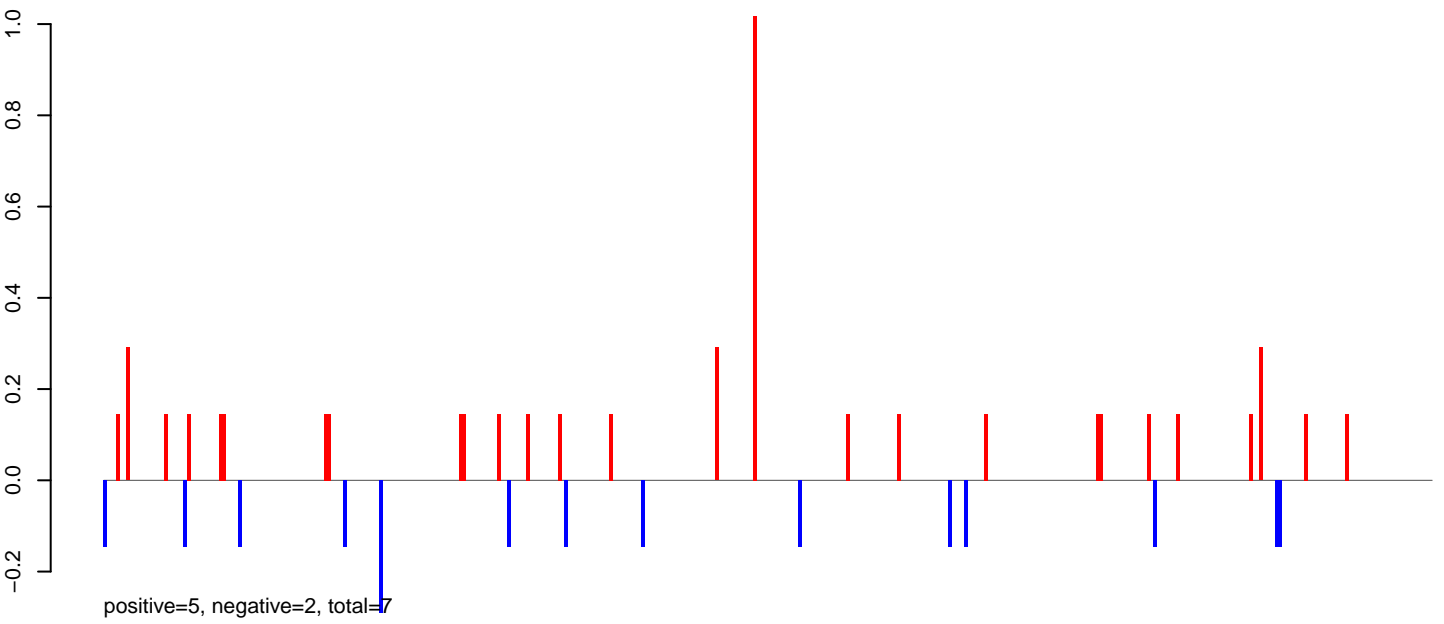
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



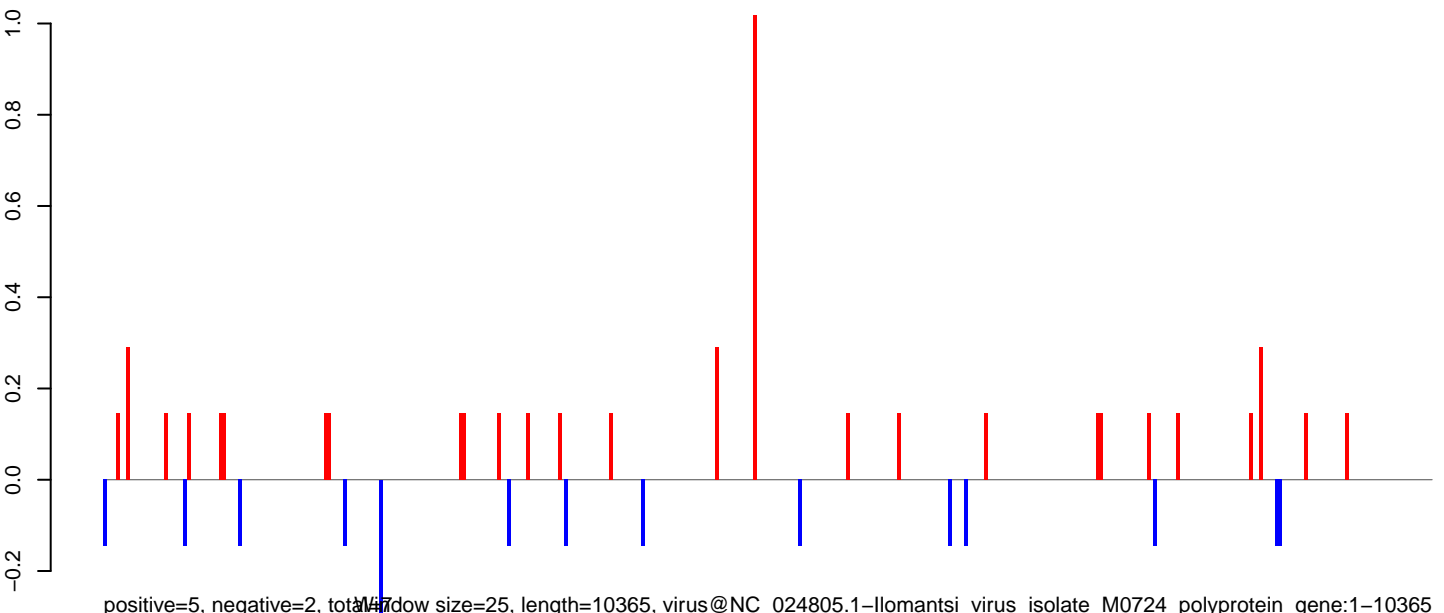
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



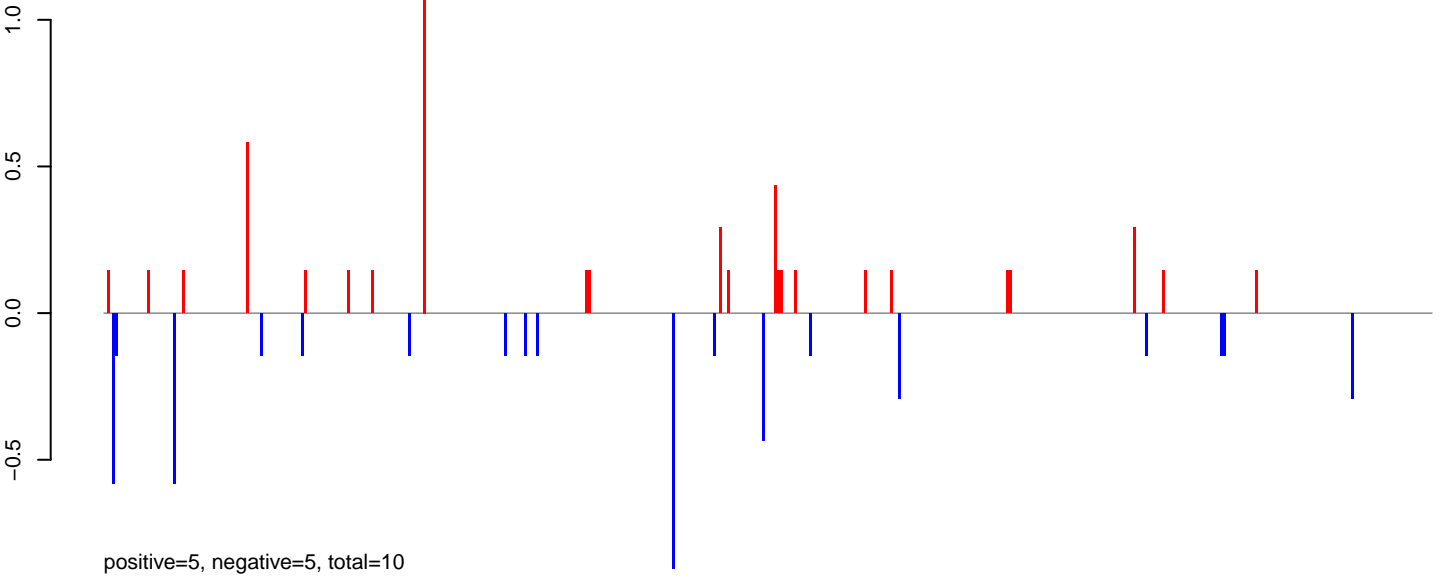
AeAeg_LVP_pMos3DBhspPBac.rep



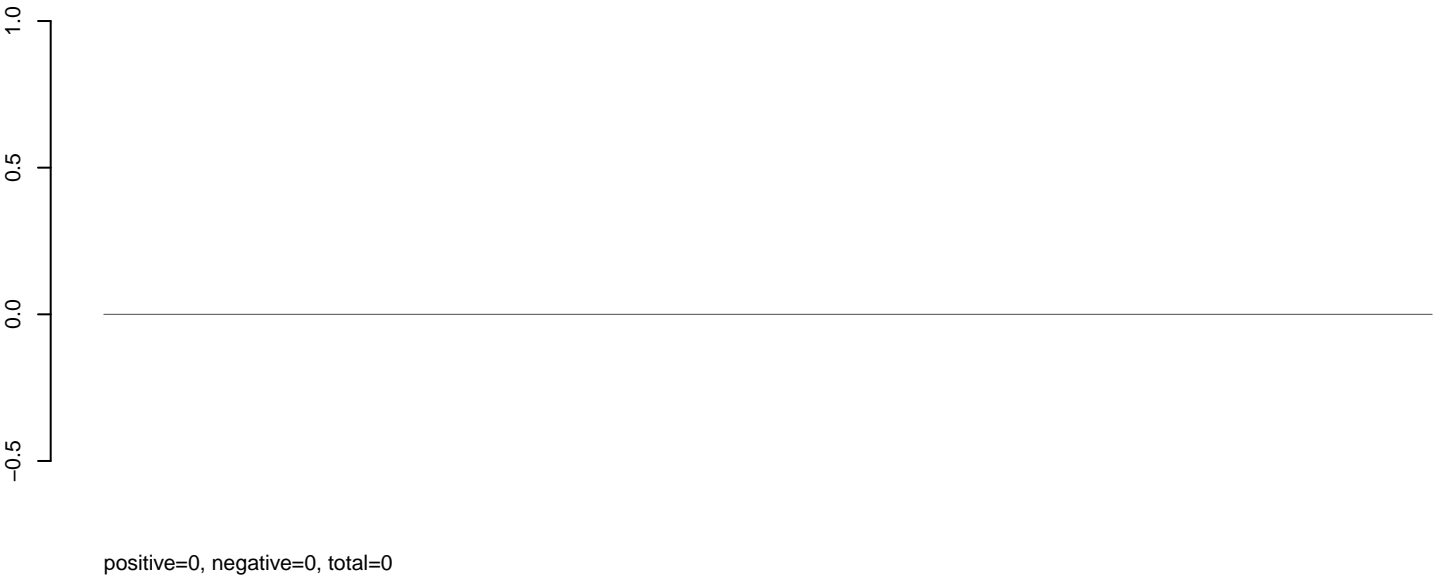
Window size=25, length=10365, virus@NC_024805.1-Ilomantsi_virus_isolate_M0724_polyprotein_gene:1-10365

0 2000 4000 6000 8000 10000

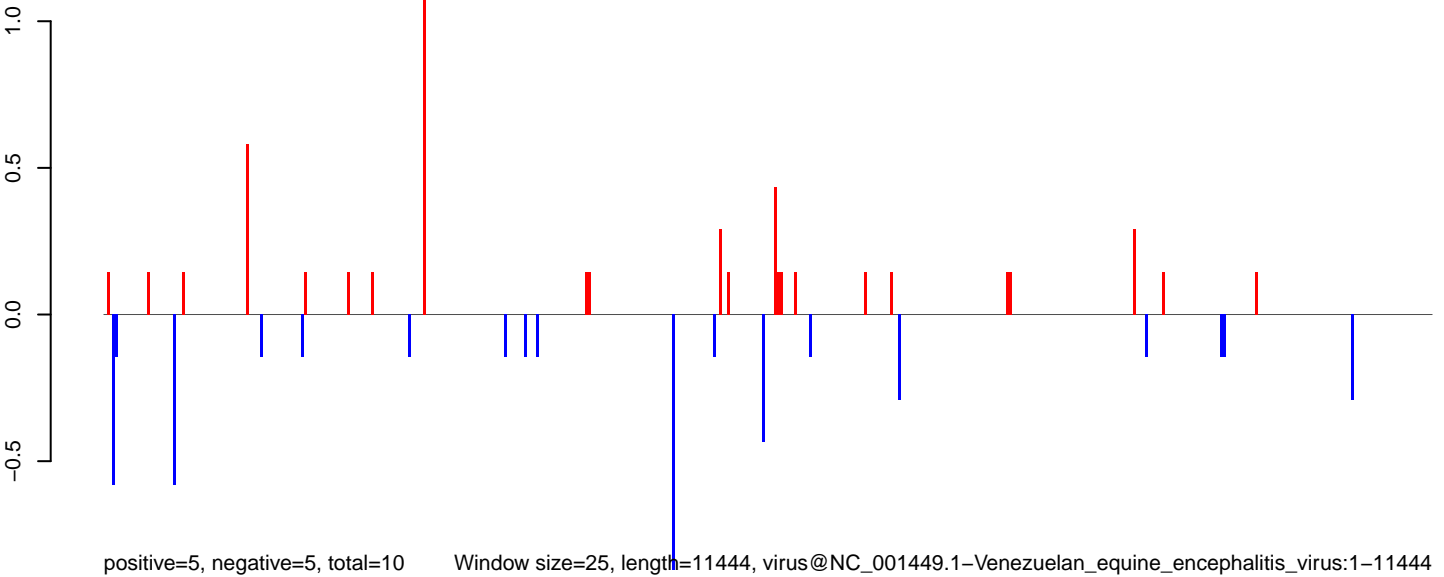
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

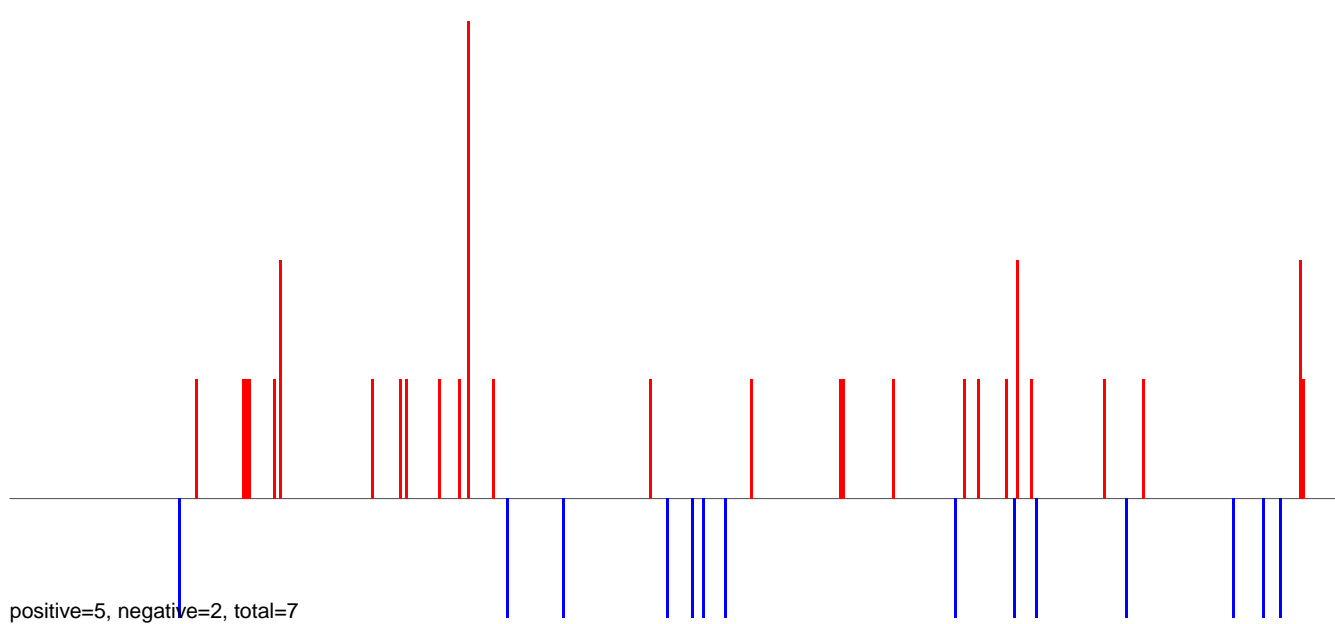


AeAeg_LVP_pMos3DBhspPBac.rep



0 2000 4000 6000 8000 10000

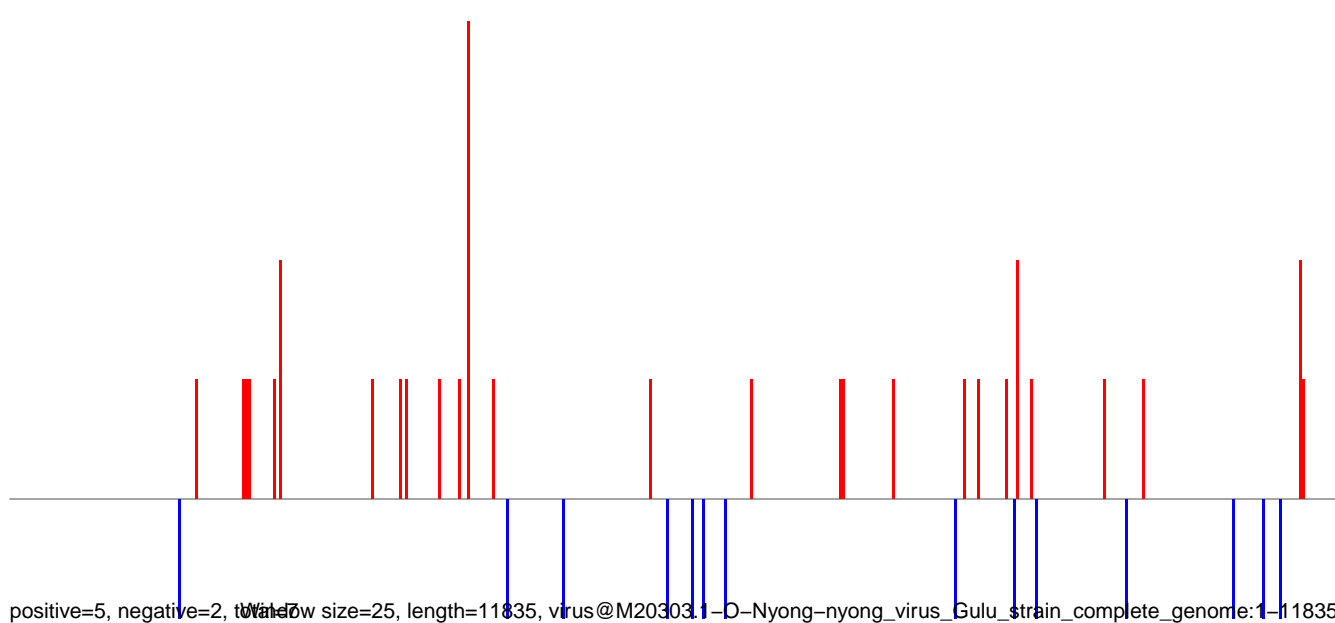
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



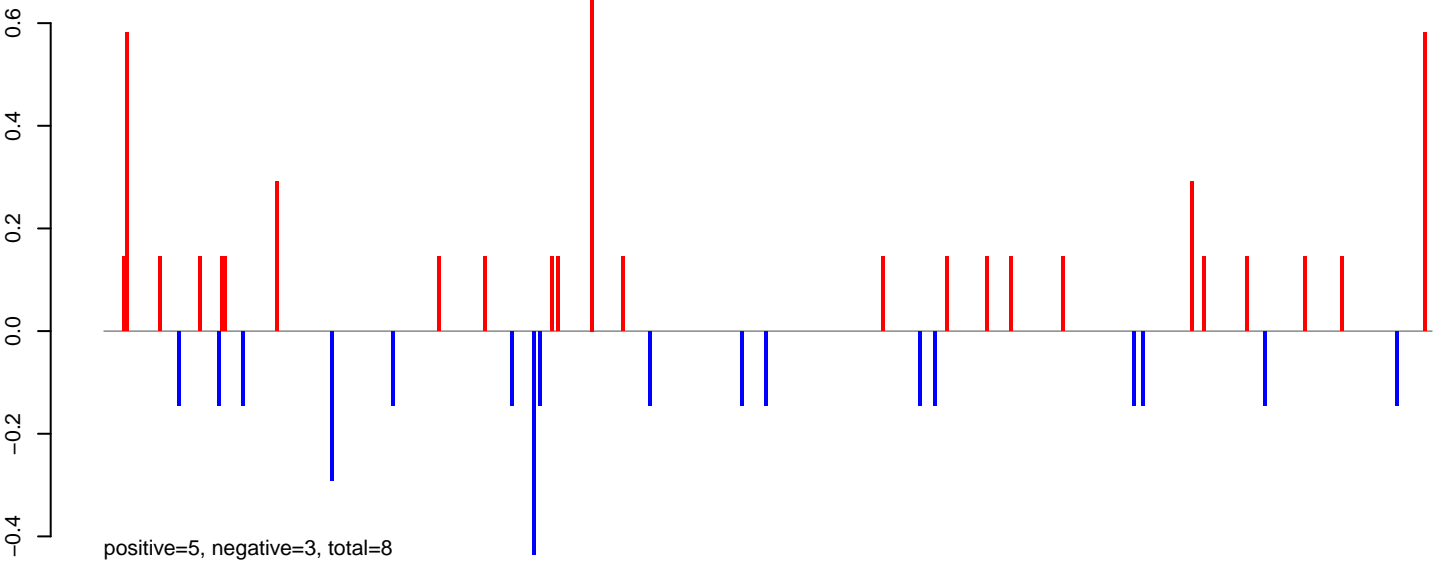
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



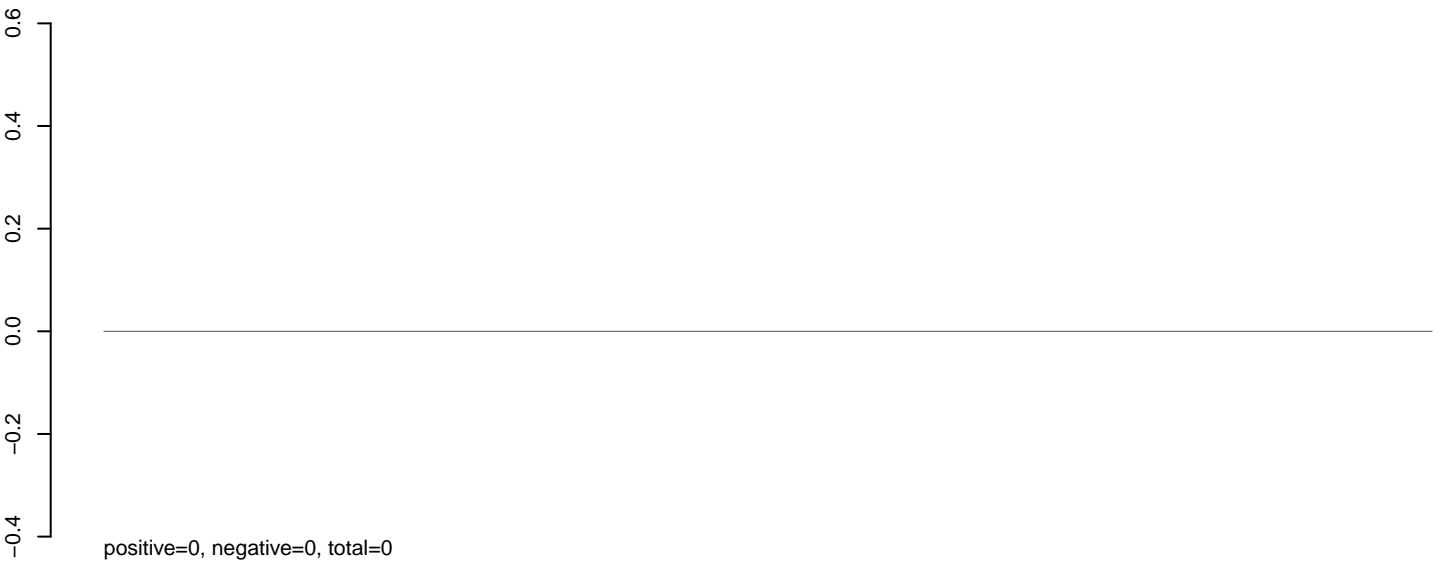
AeAeg_LVP_pMos3DBhspPBac.rep



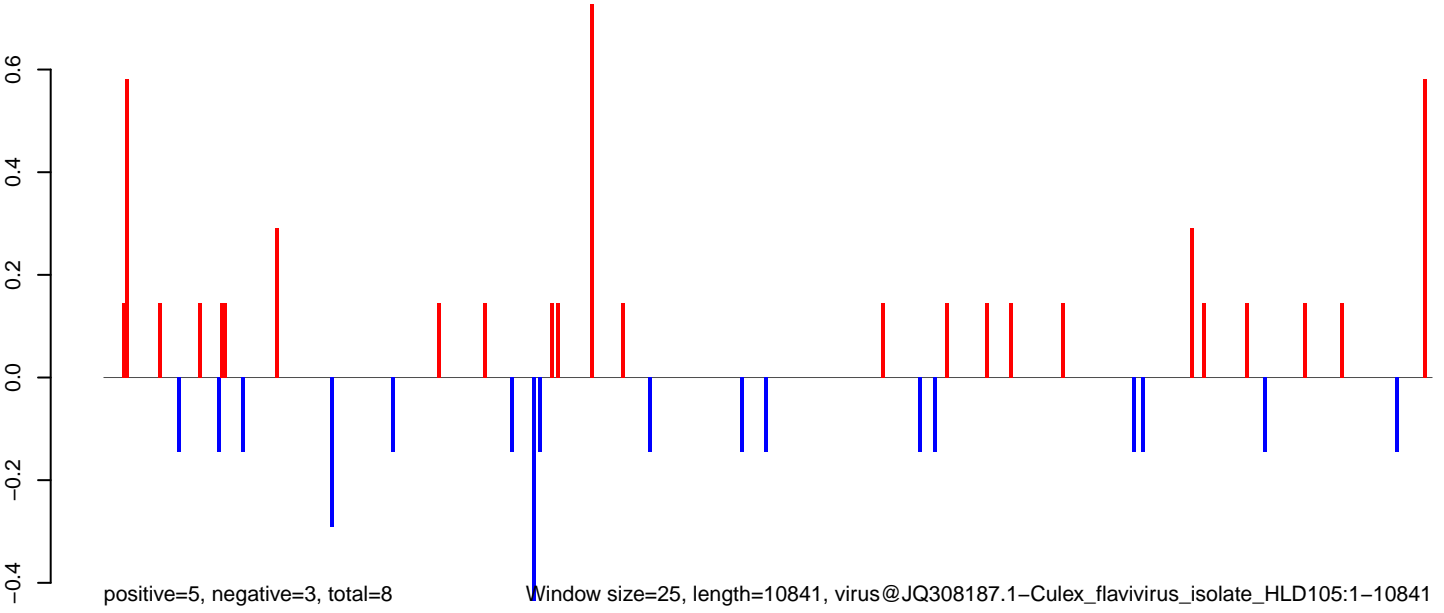
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

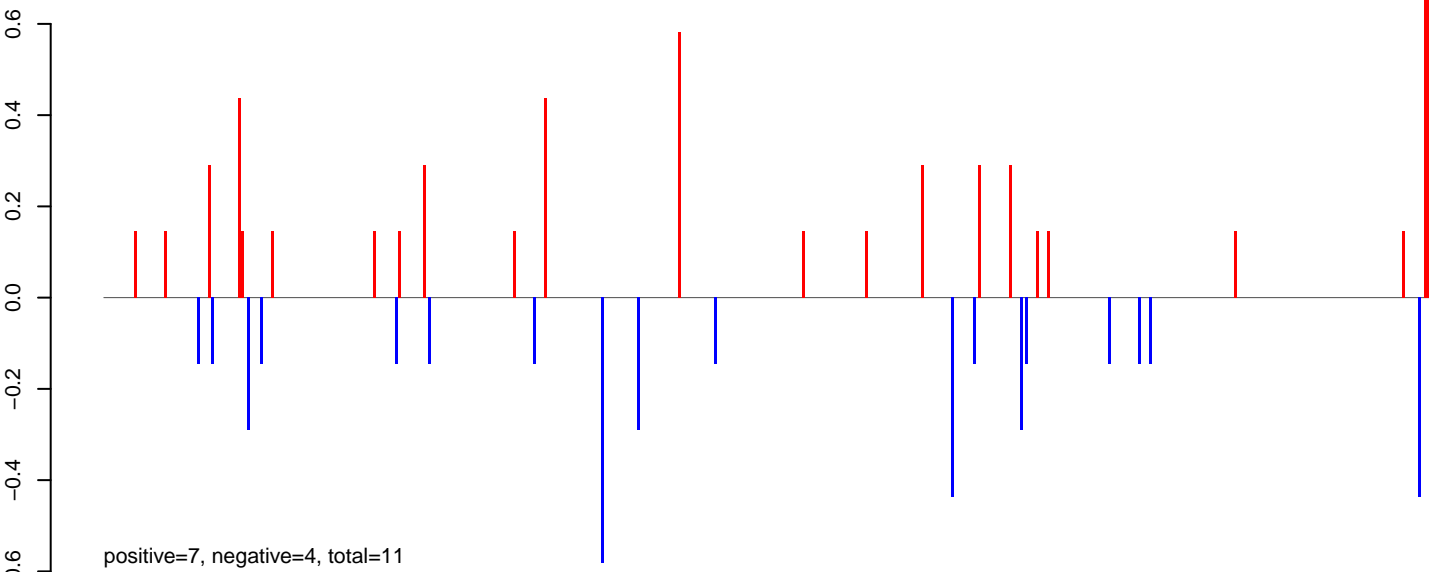


AeAeg_LVP_pMos3DBhspPBac.rep

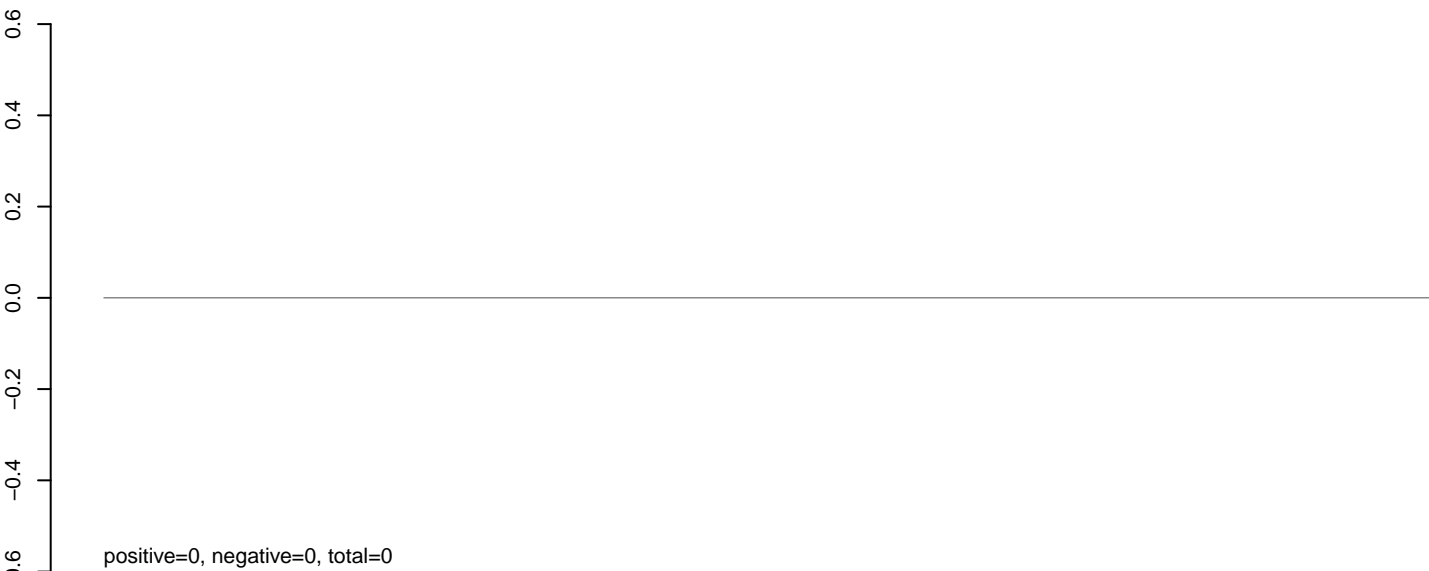


0 2000 4000 6000 8000 10000

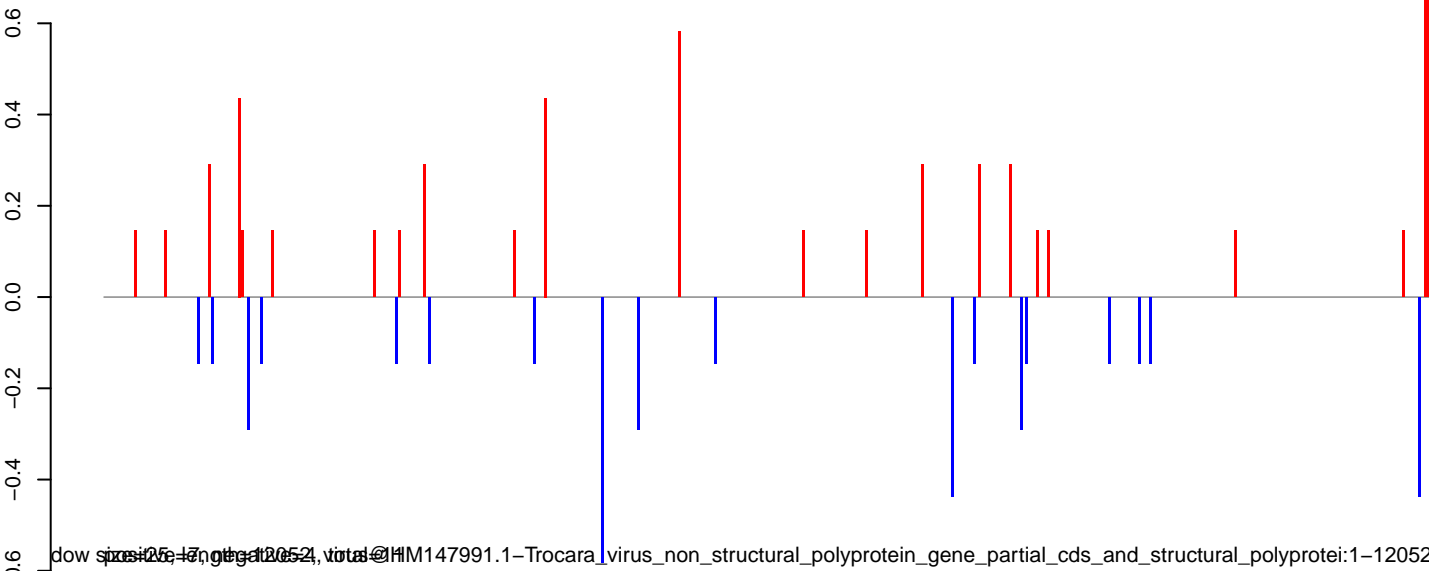
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

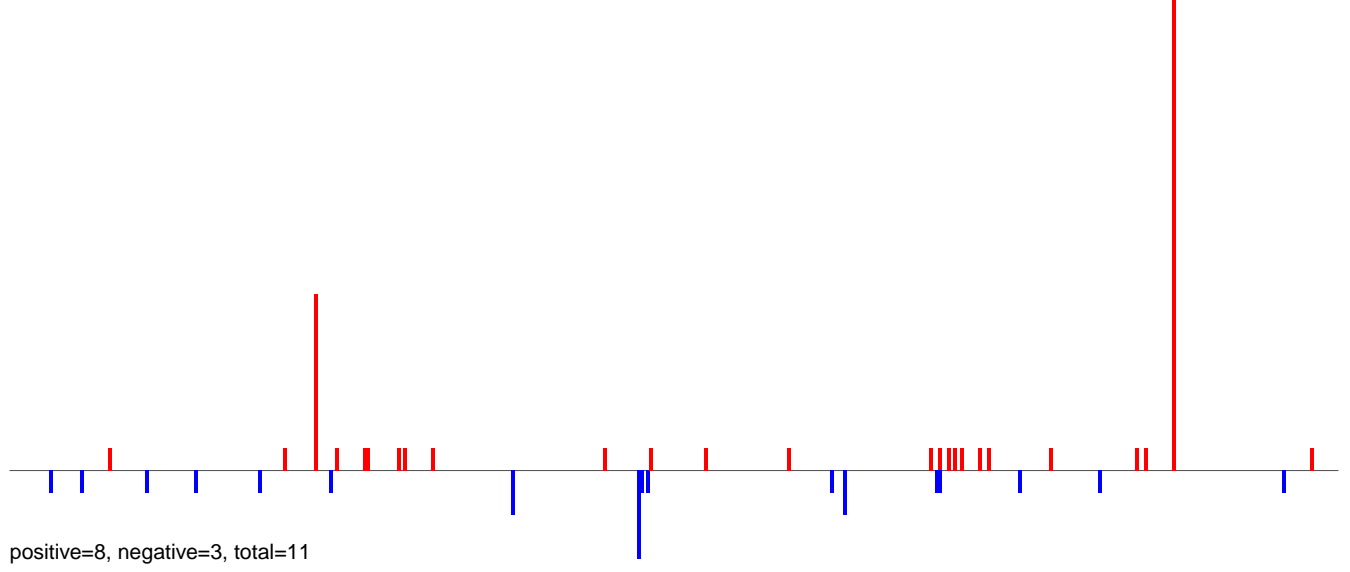


AeAeg_LVP_pMos3DBhspPBac.rep

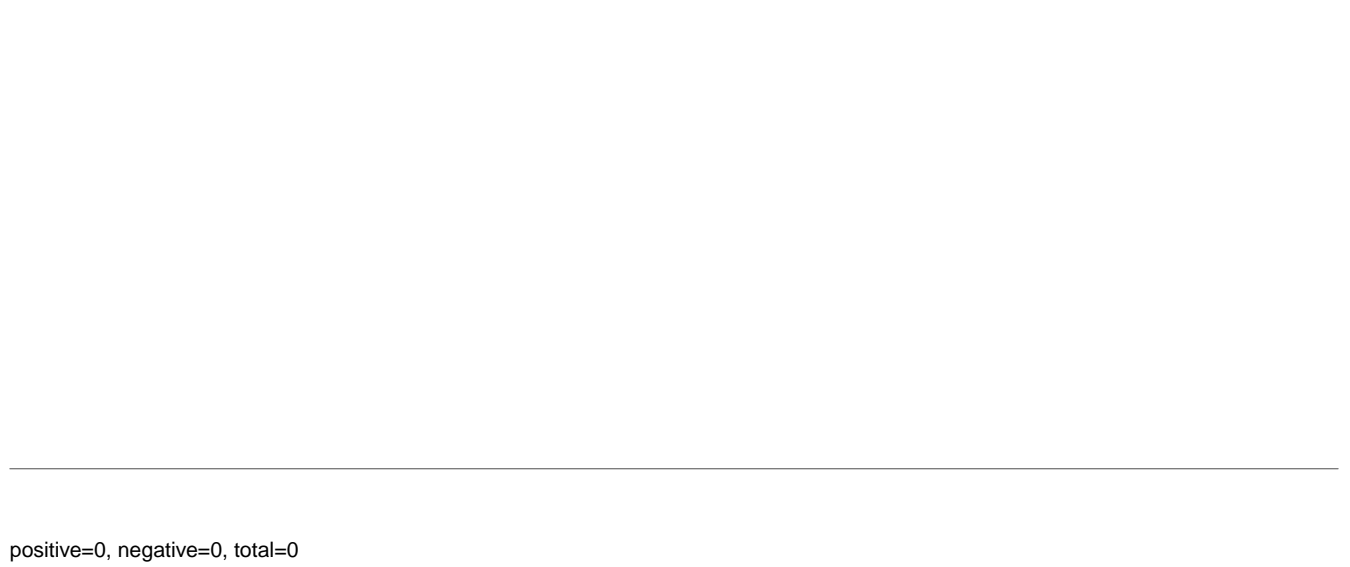


downloaded from <https://www.ncbi.nlm.nih.gov/nuccore/1147991.1> Trocara_virus_non_structural_polyprotein_gene_partial_cds_and_structural_polyprotei:1-12052

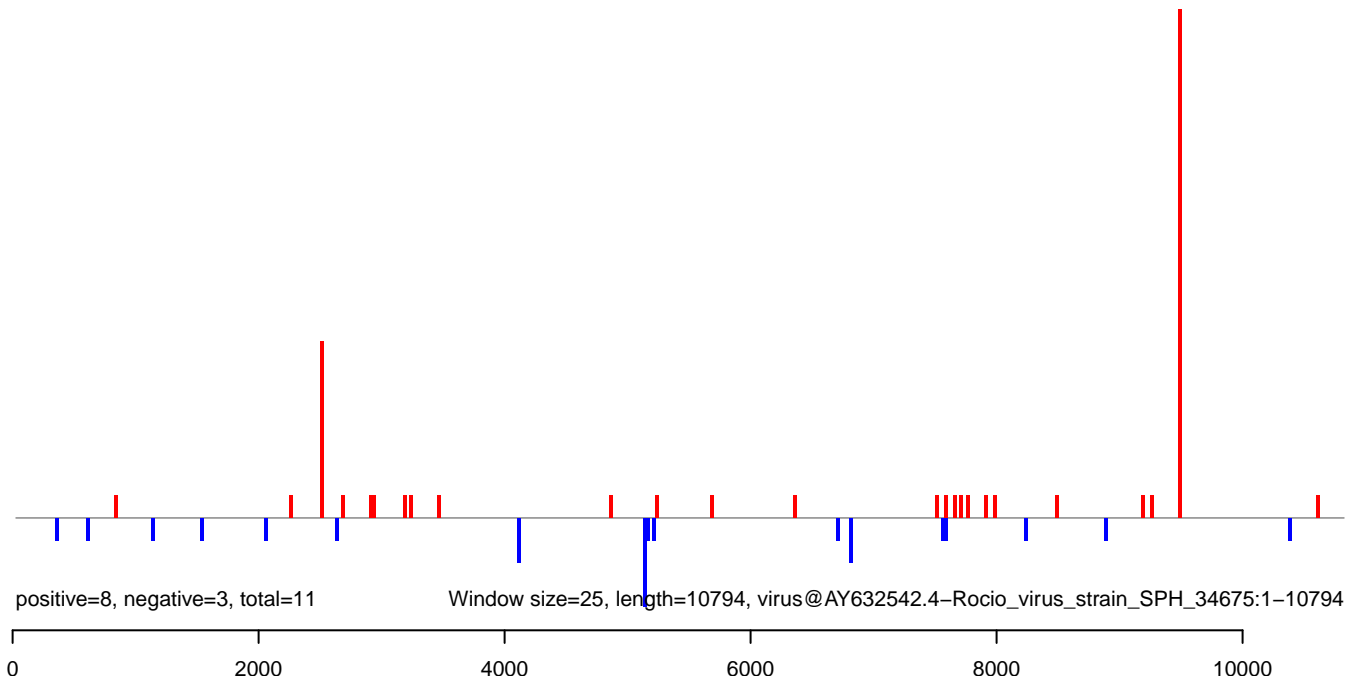
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



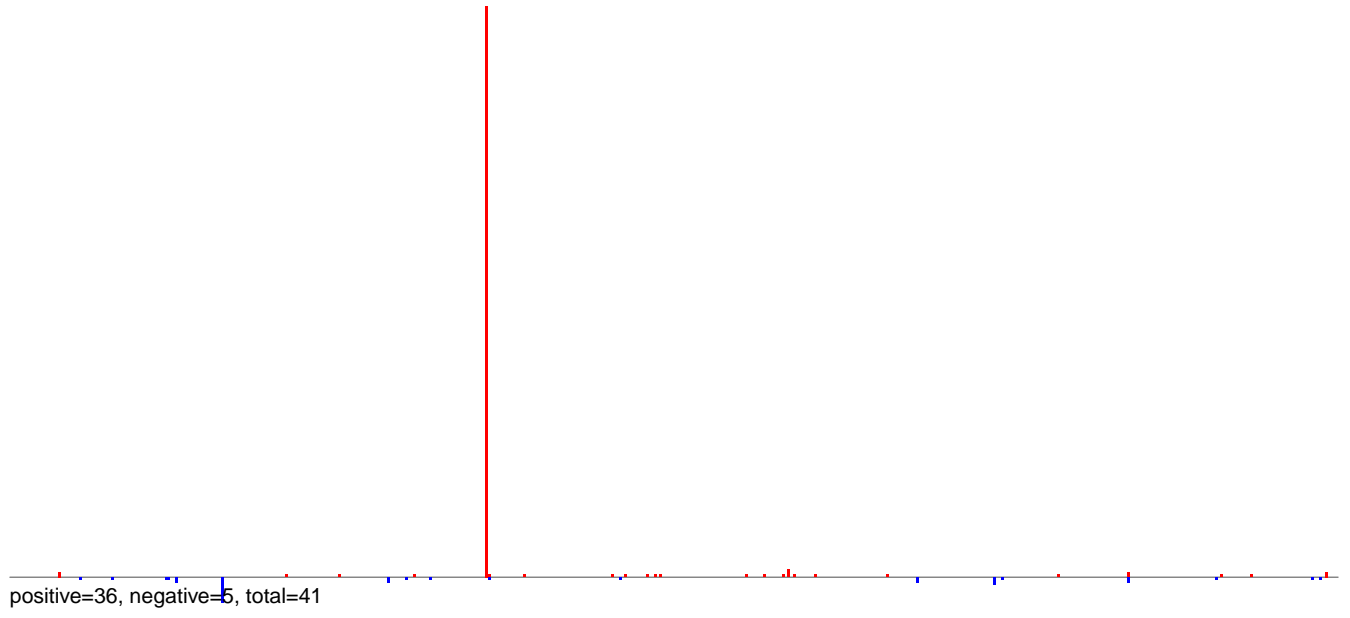
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



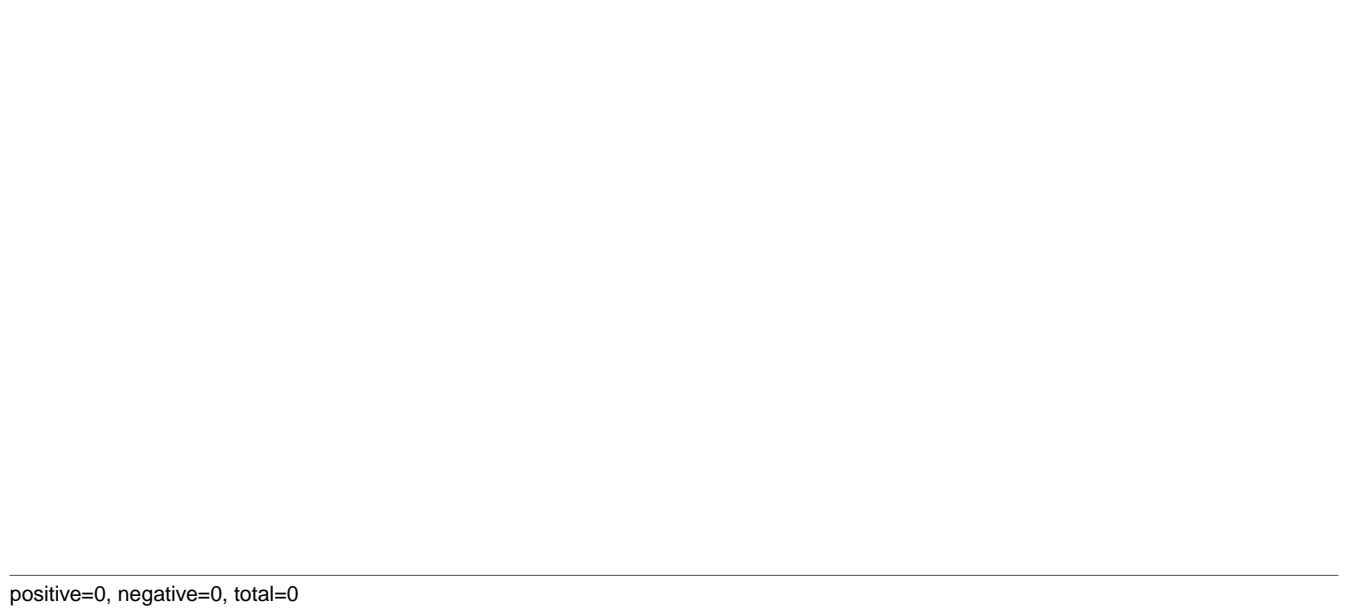
AeAeg_LVP_pMos3DBhspPBac.rep



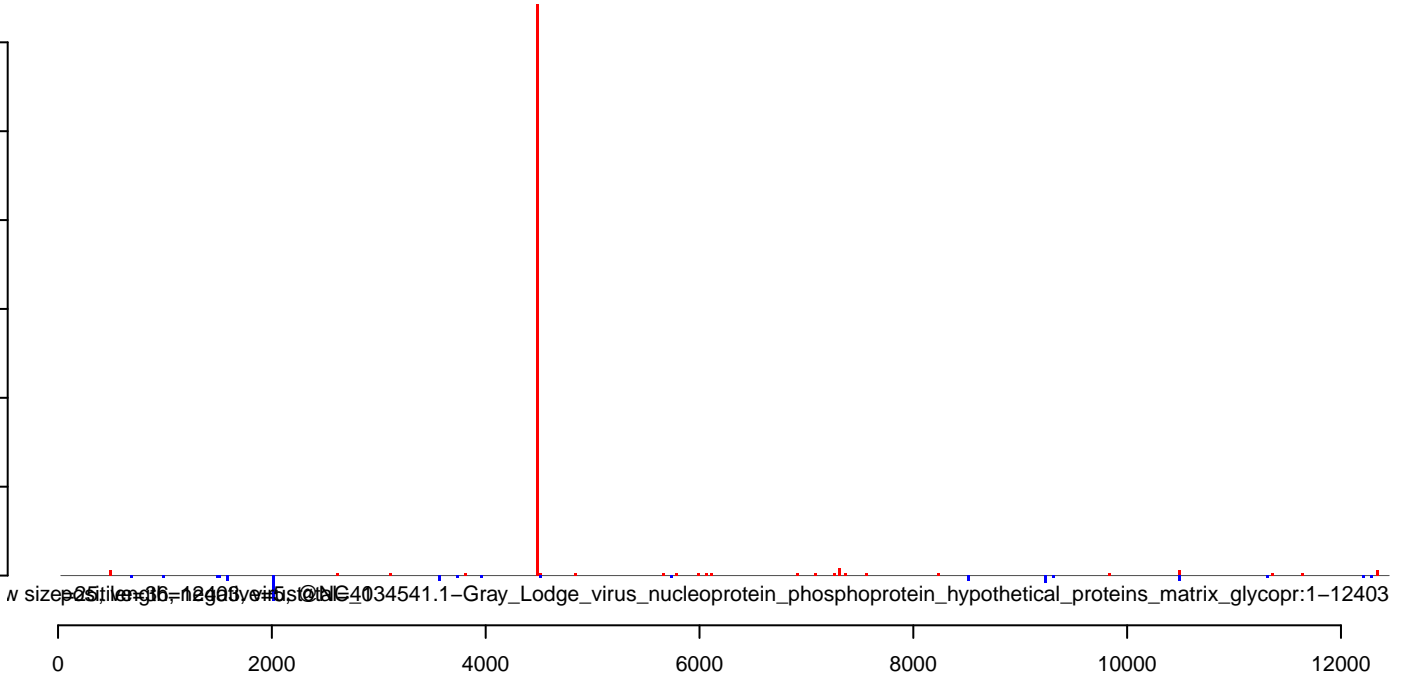
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

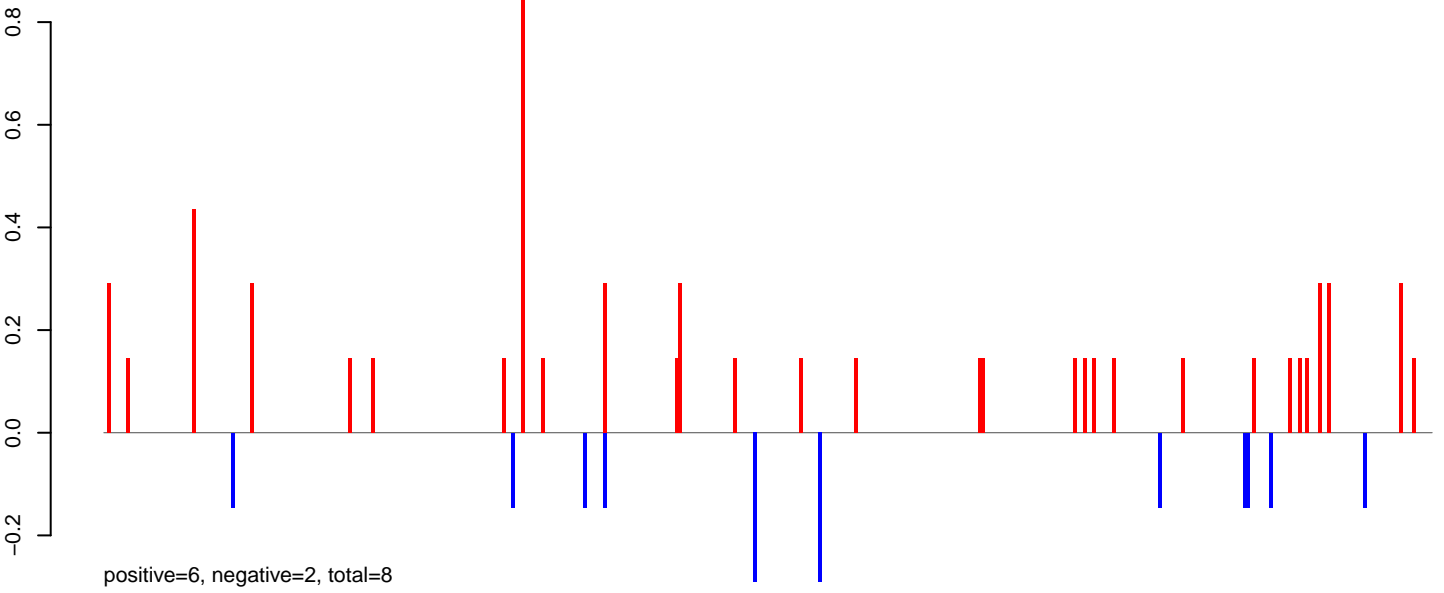


AeAeg_LVP_pMos3DBhspPBac.rep

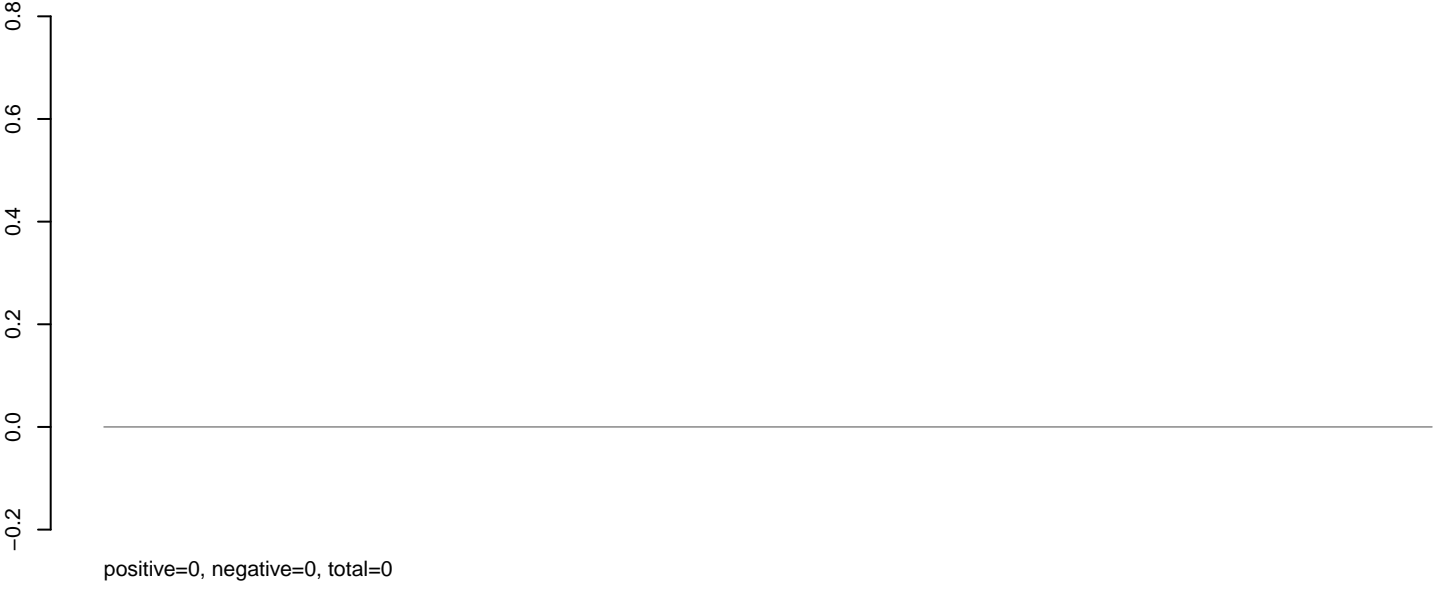


1-Gray_Lodge_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopr:1-12403

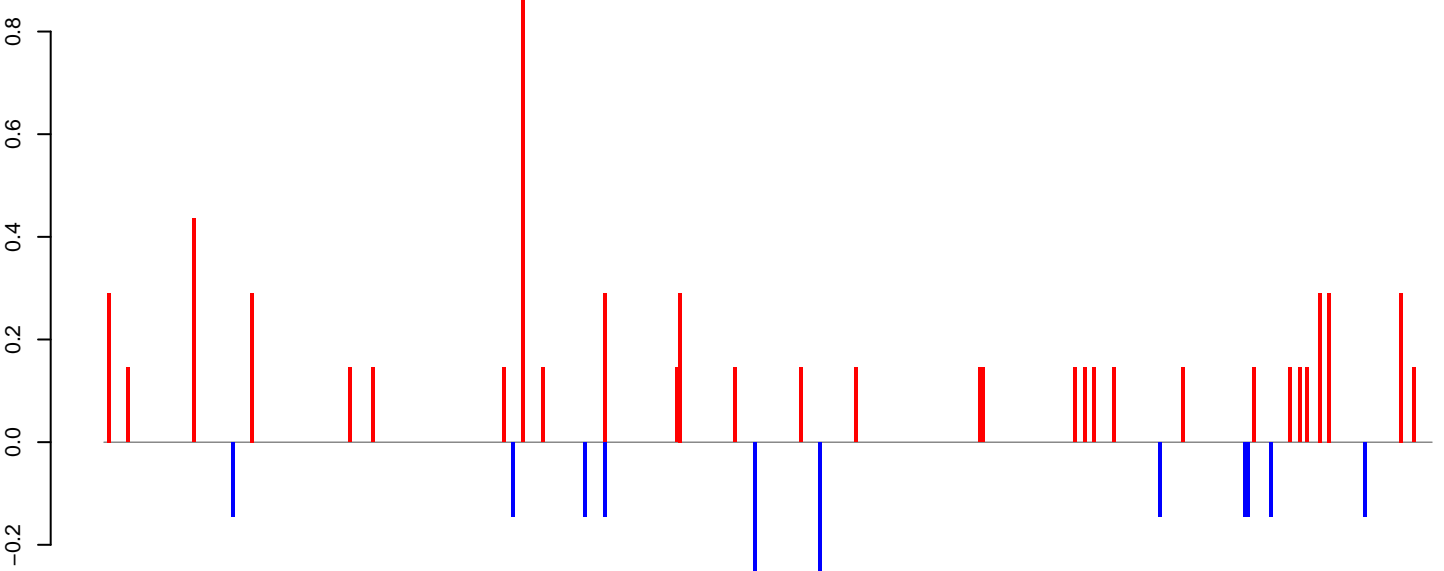
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



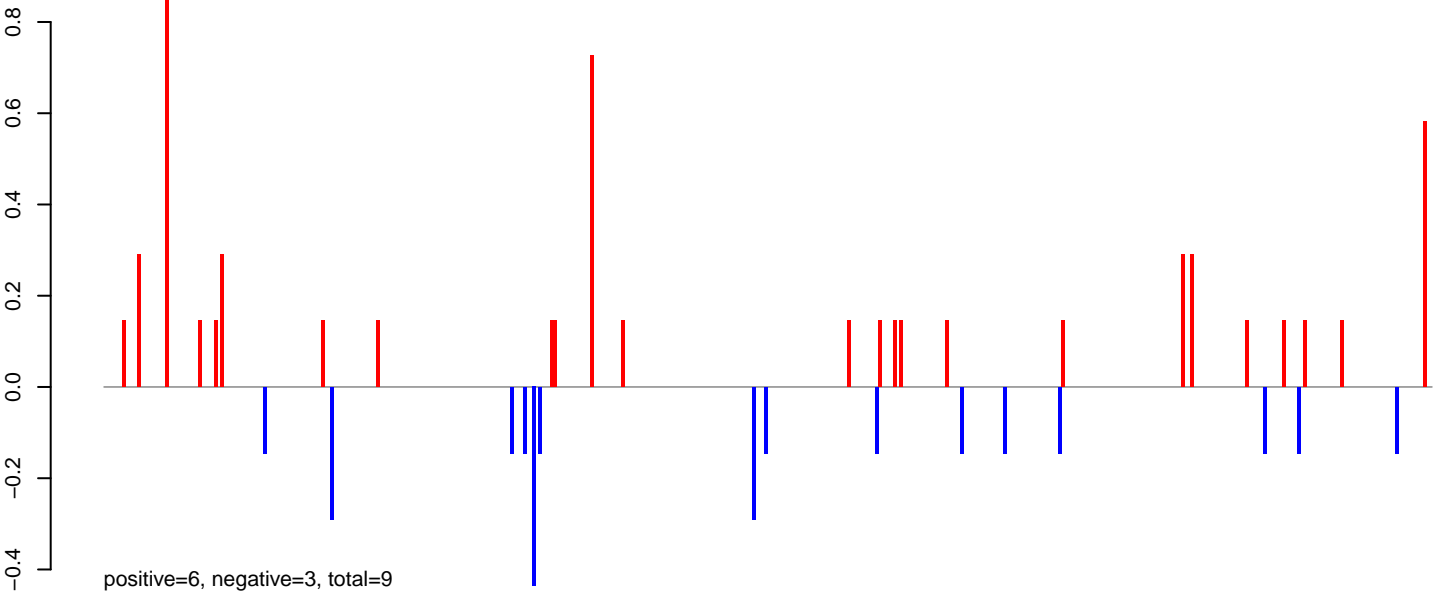
AeAeg_LVP_pMos3DBhspPBac.rep



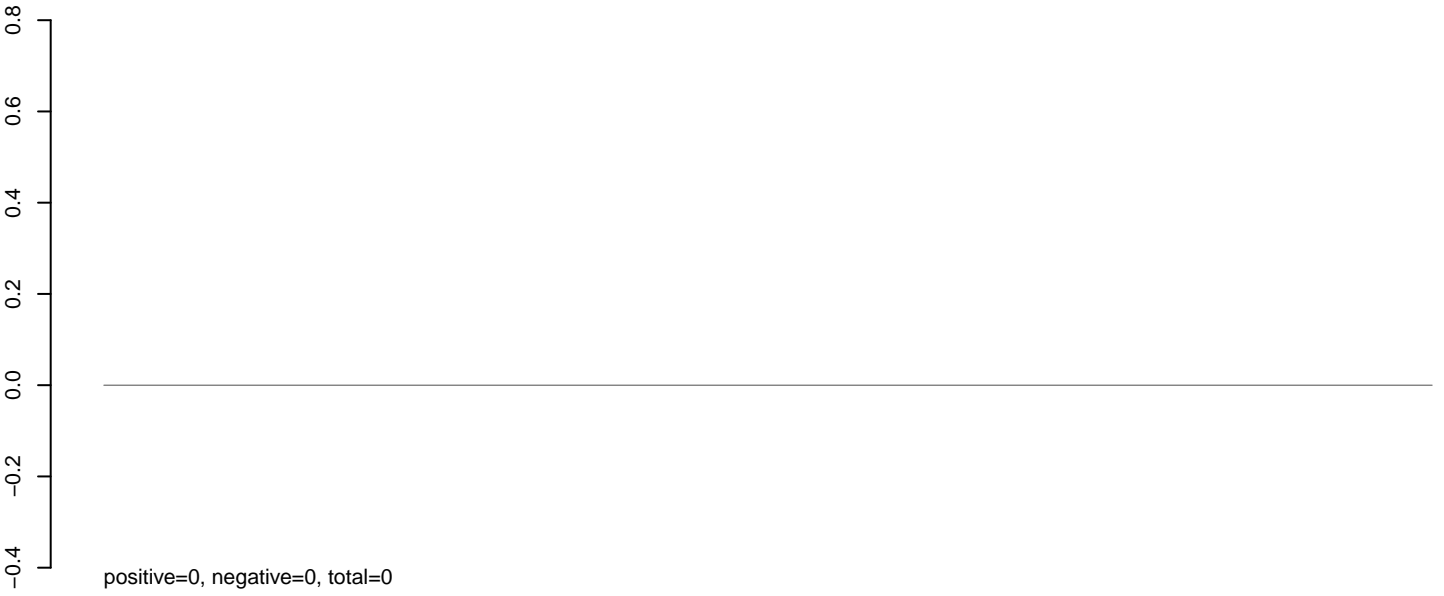
Window size=25, length=10173, virus@NC_033699.1-Jugra_virus_strain_P-9-314_polyprotein_gene:1-10173

0 2000 4000 6000 8000 10000

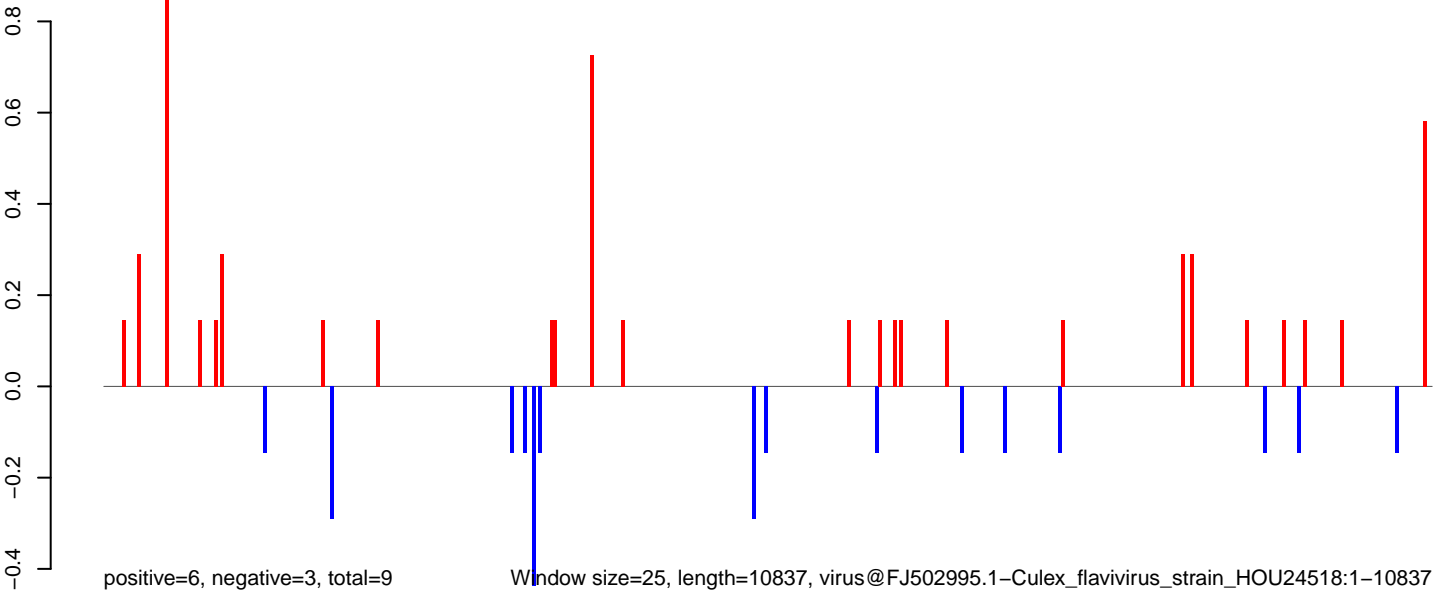
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

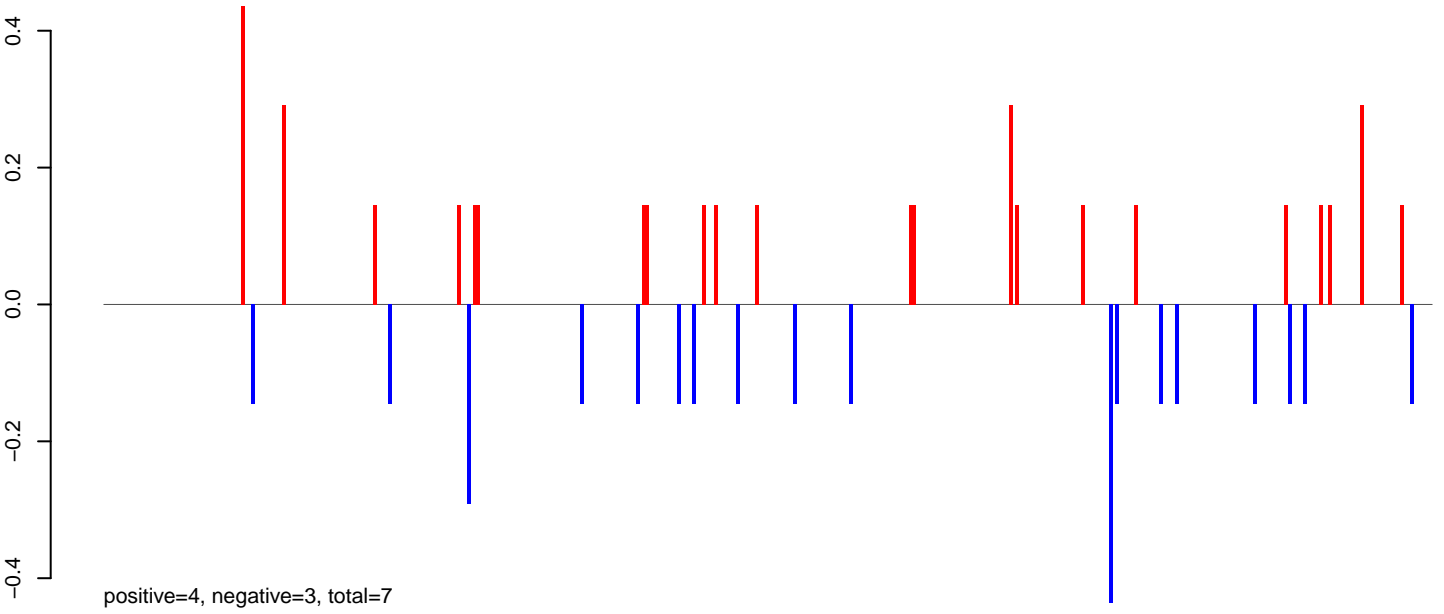


AeAeg_LVP_pMos3DBhspPBac.rep

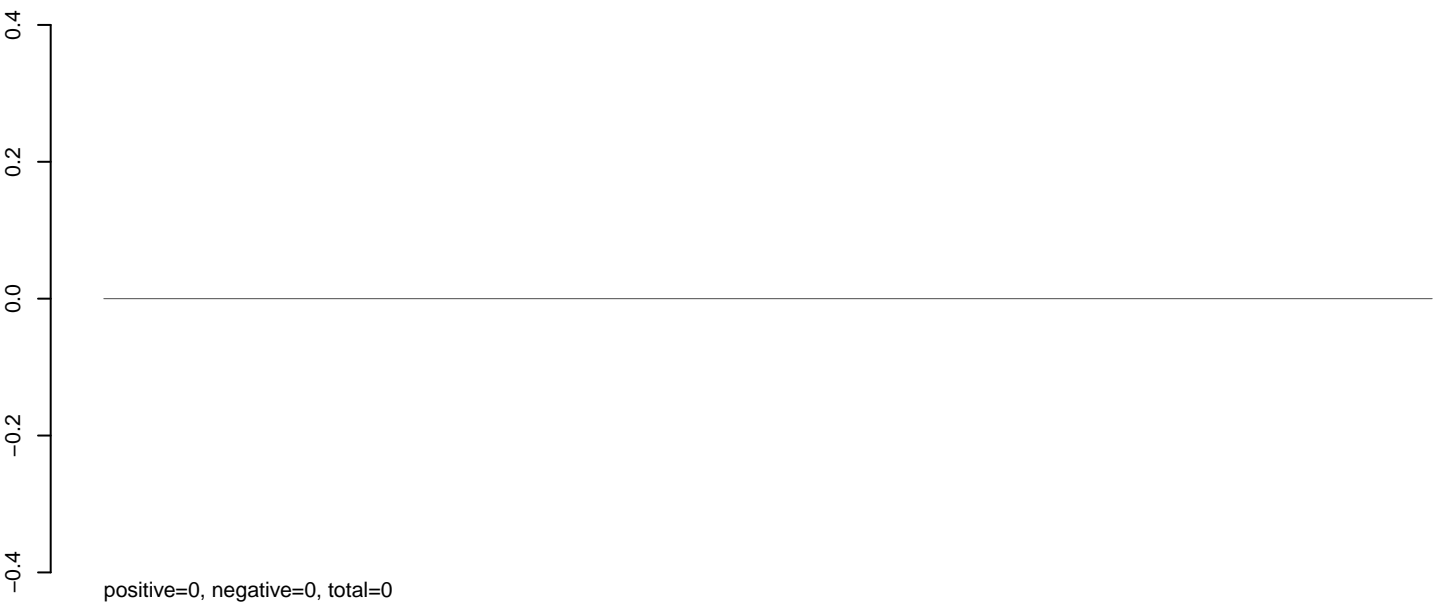


0 2000 4000 6000 8000 10000

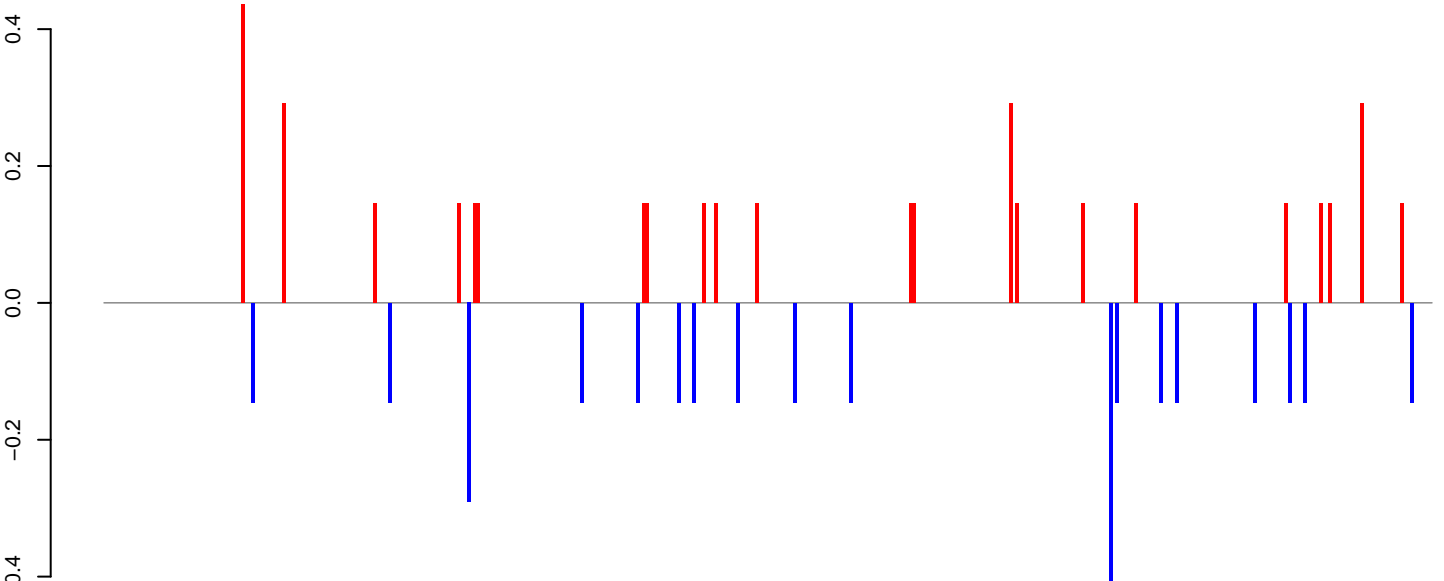
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

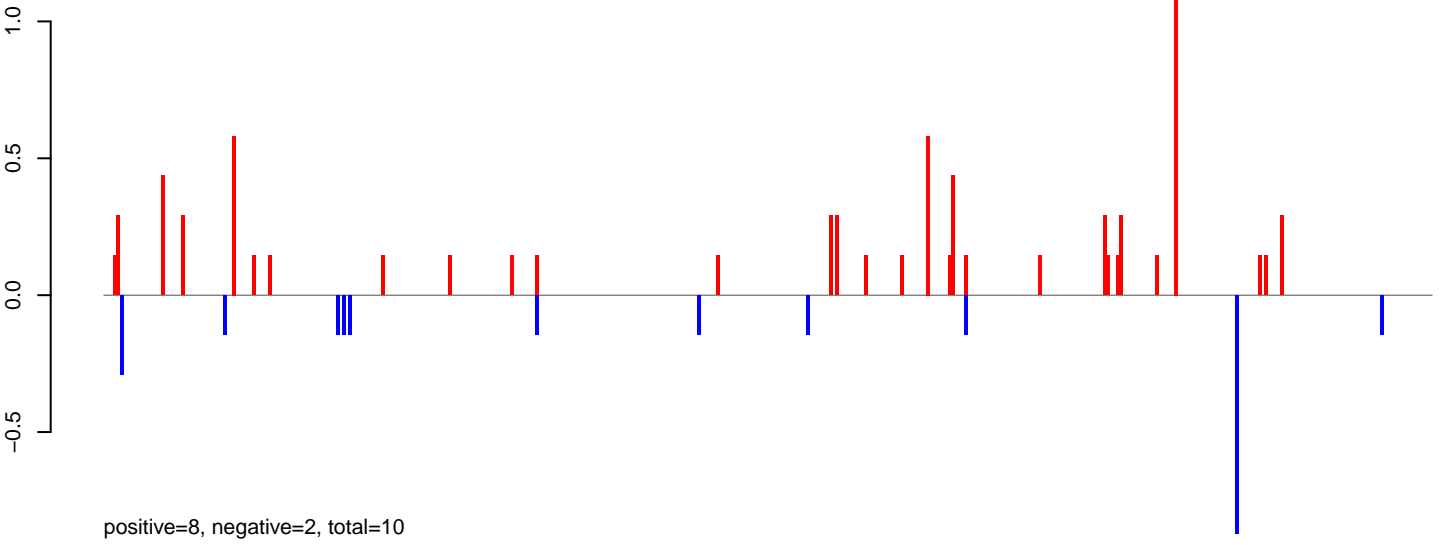


AeAeg_LVP_pMos3DBhspPBac.rep

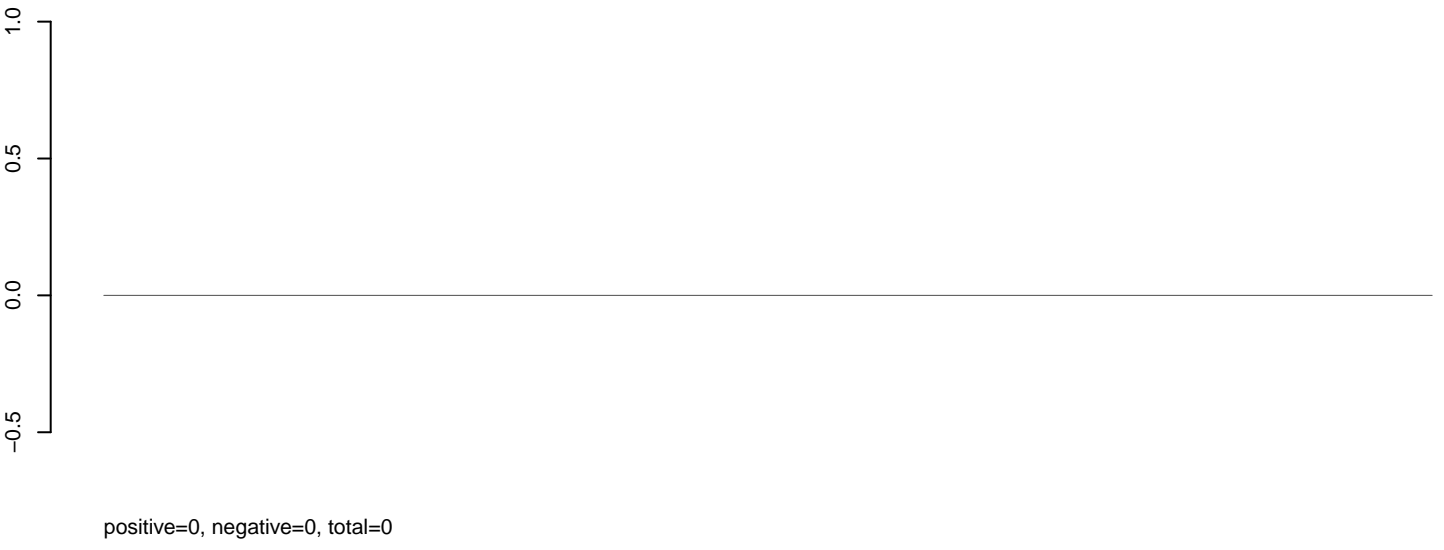


0 2000 4000 6000 8000 10000

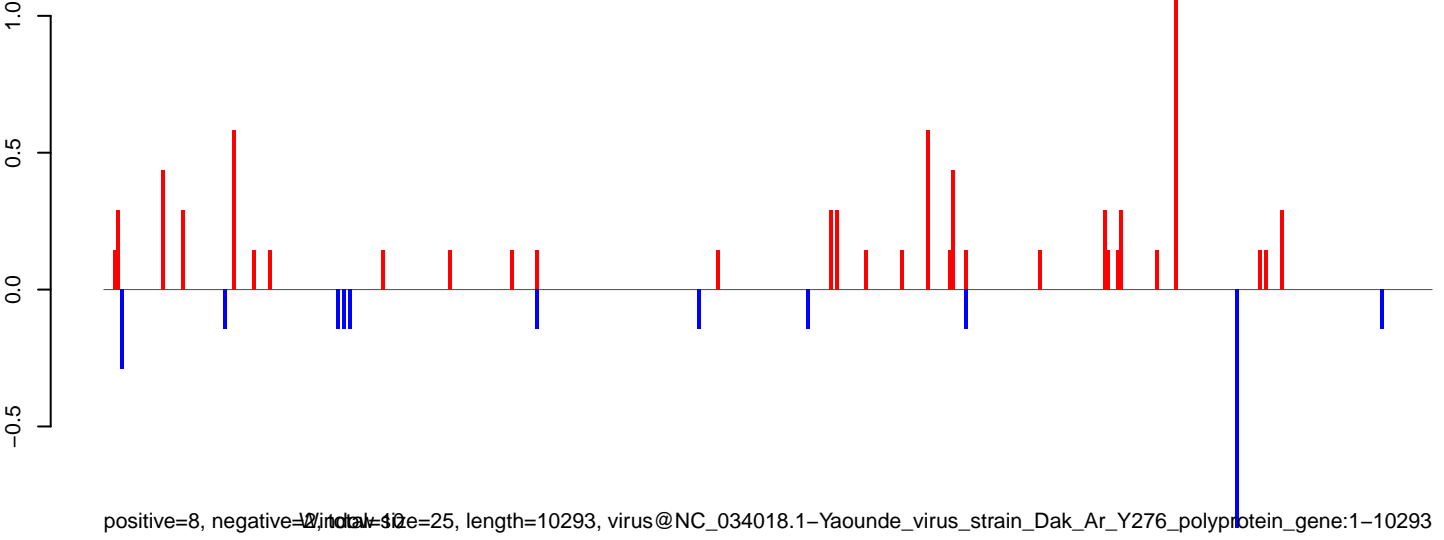
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

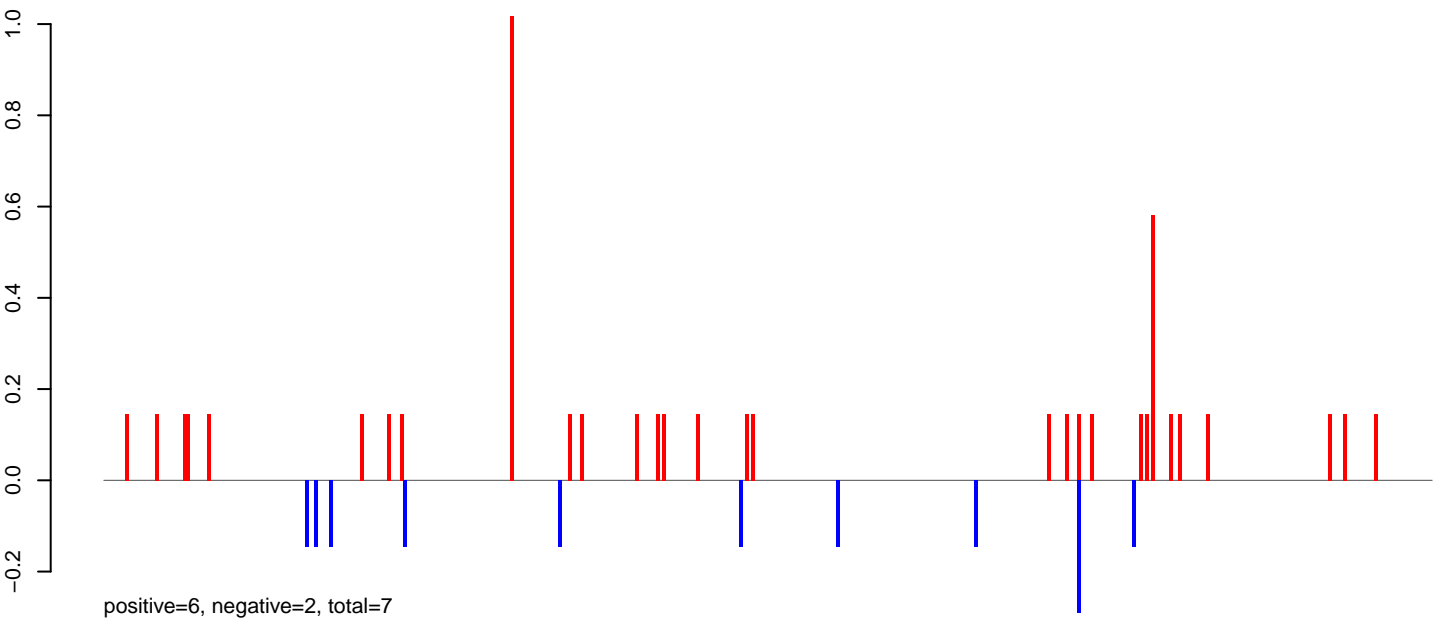


AeAeg_LVP_pMos3DBhspPBac.rep

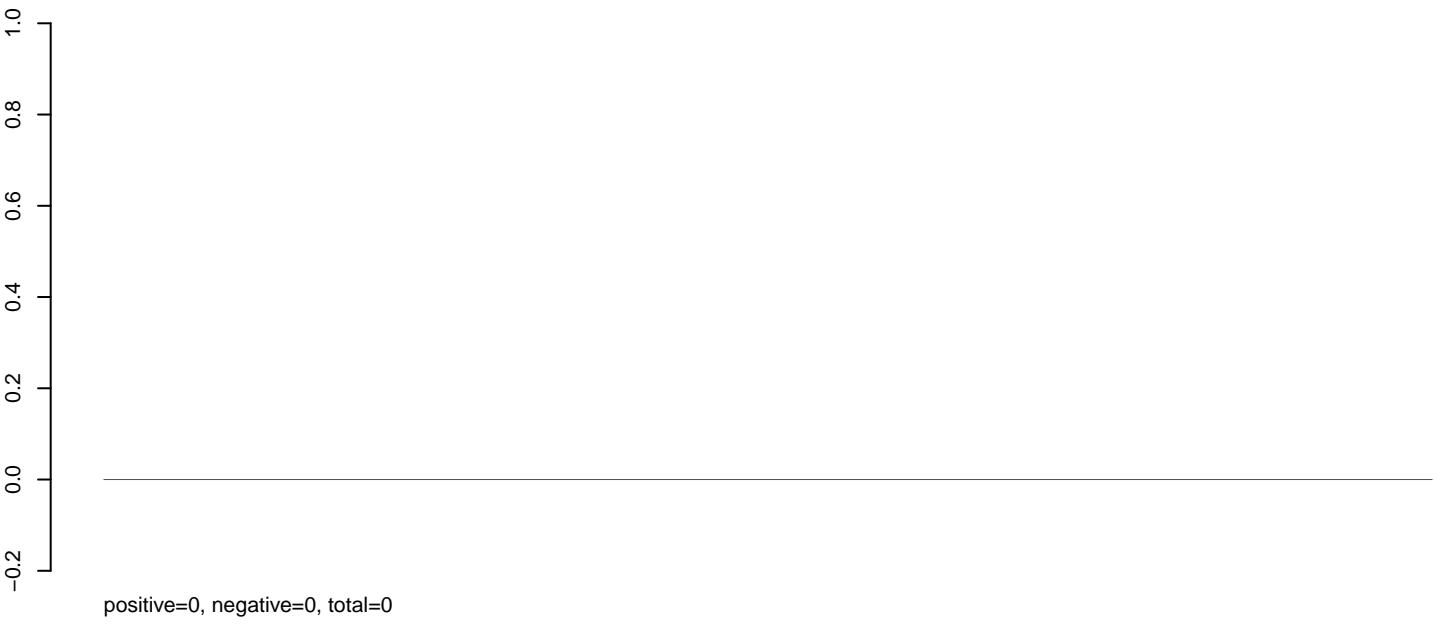


positive=8, negative=2, total=10, virus@NC_034018.1-Yaounde_virus_strain_Dak_Ar_Y276_polyprotein_gene:1-10293

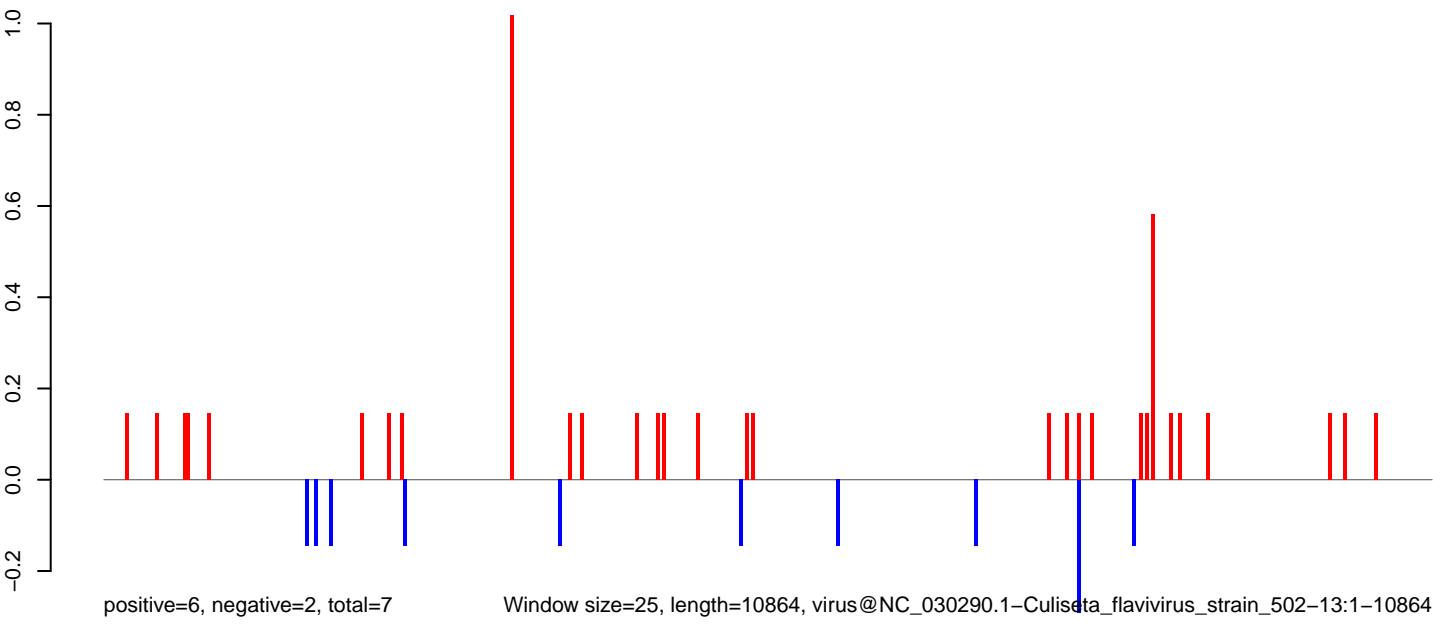
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



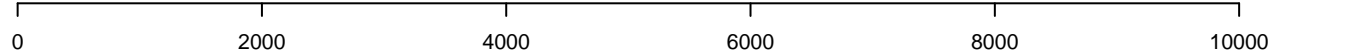
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



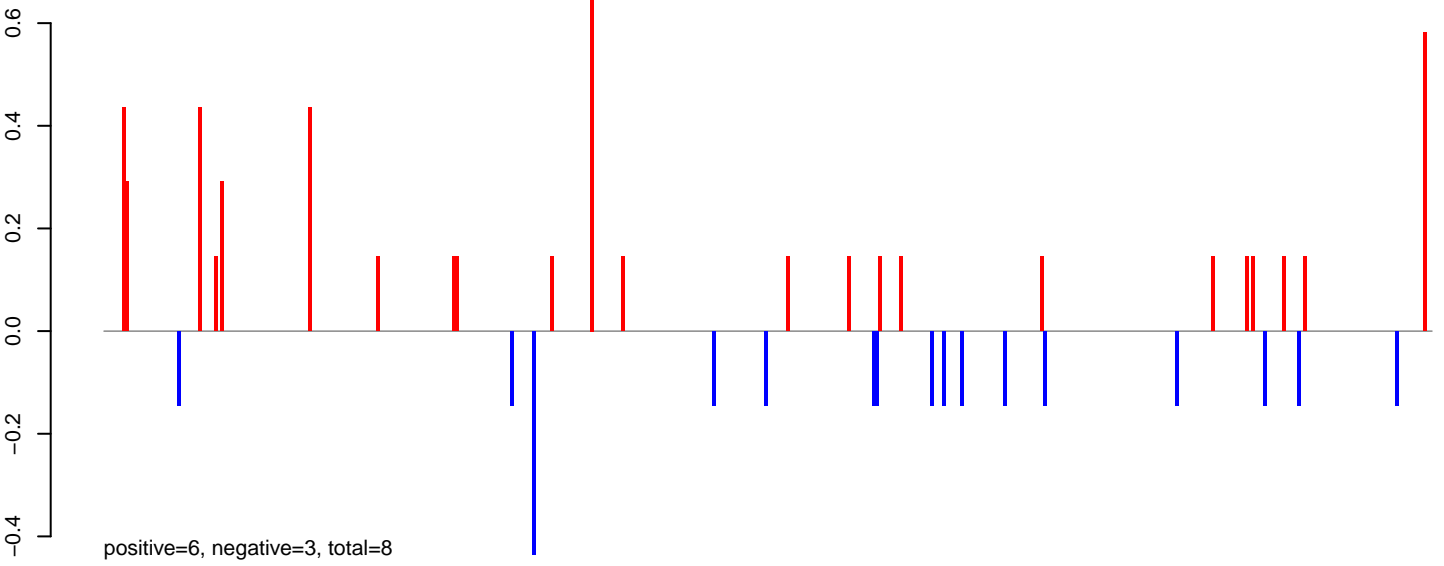
AeAeg_LVP_pMos3DBhspPBac.rep



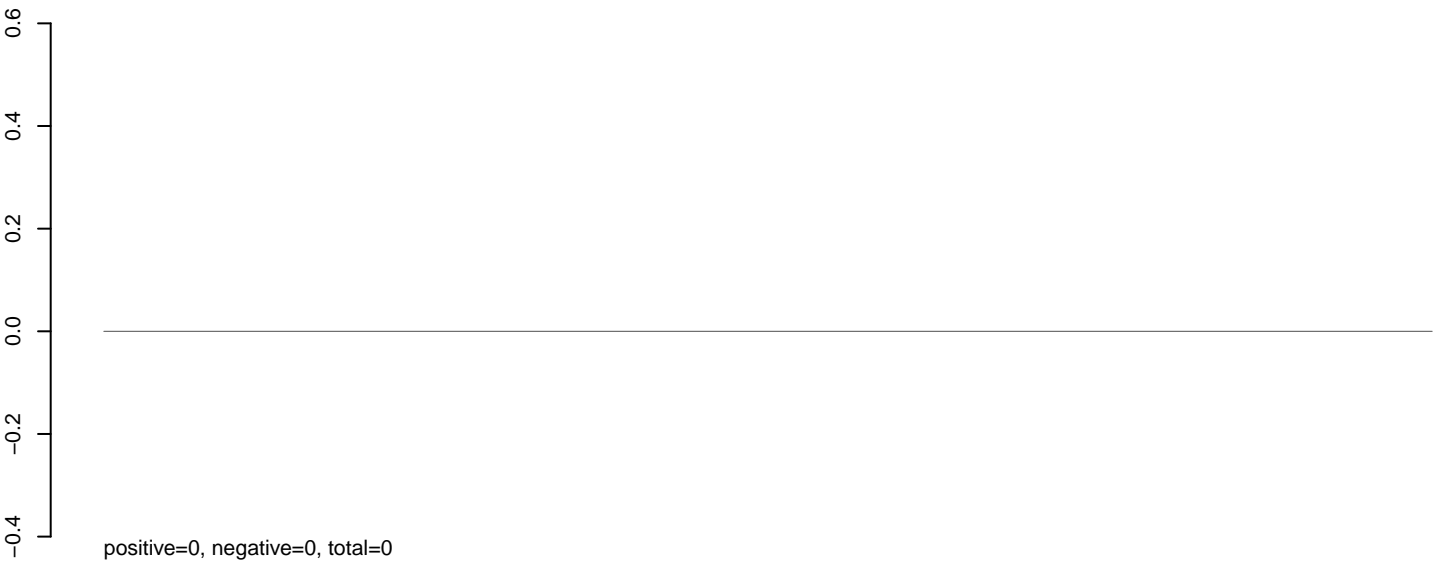
Window size=25, length=10864, virus@NC_030290.1-Culiseta_flavivirus_strain_502-13:1-10864



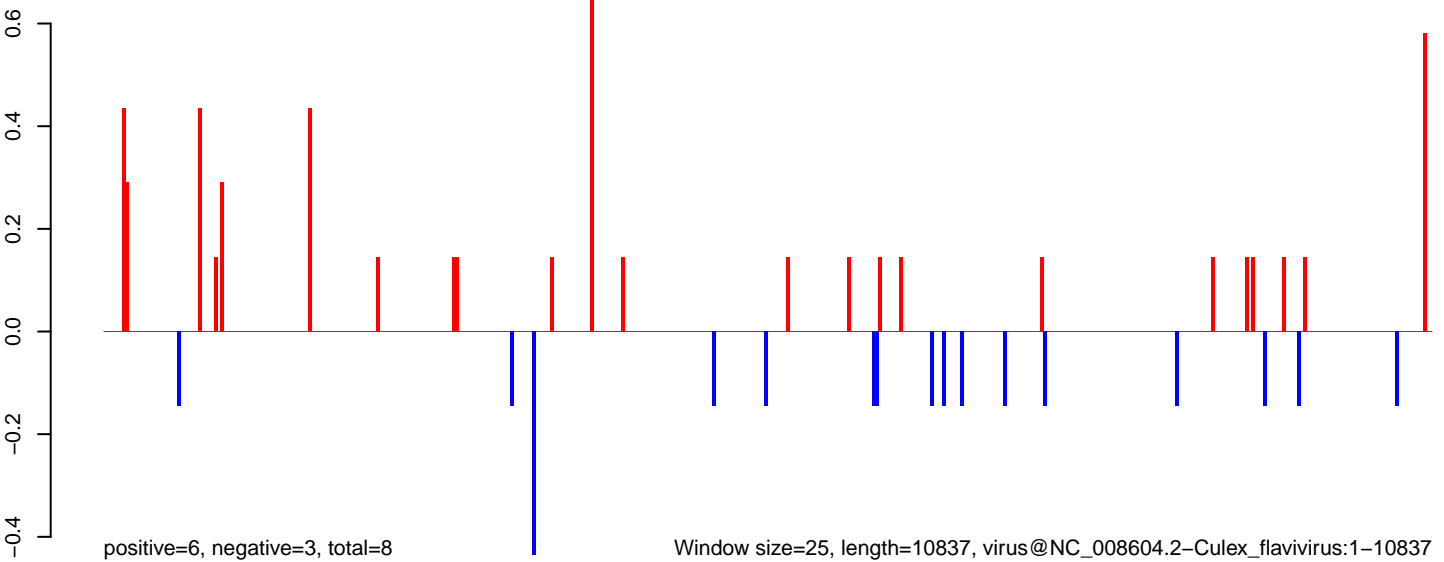
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



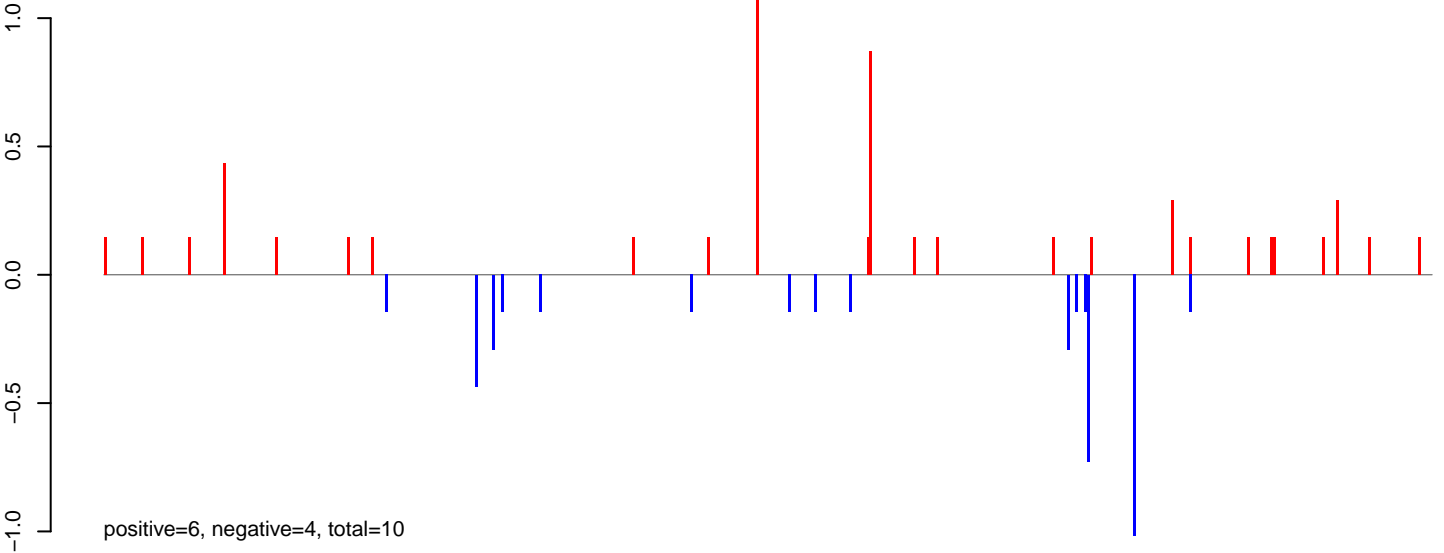
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



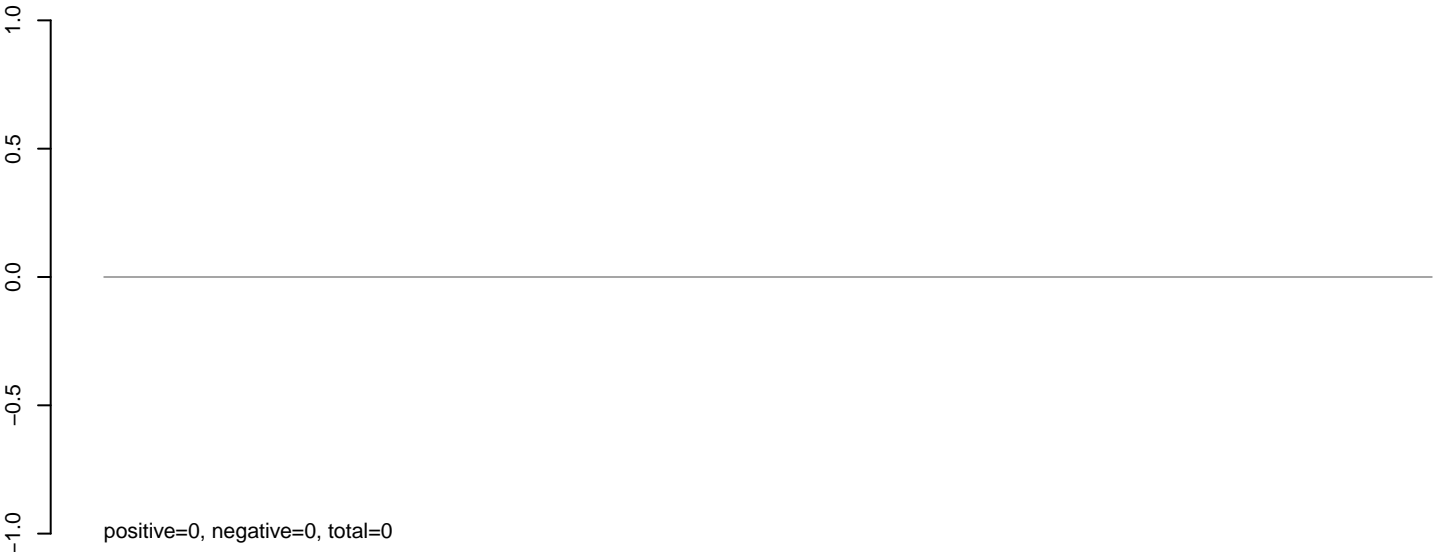
AeAeg_LVP_pMos3DBhspPBac.rep



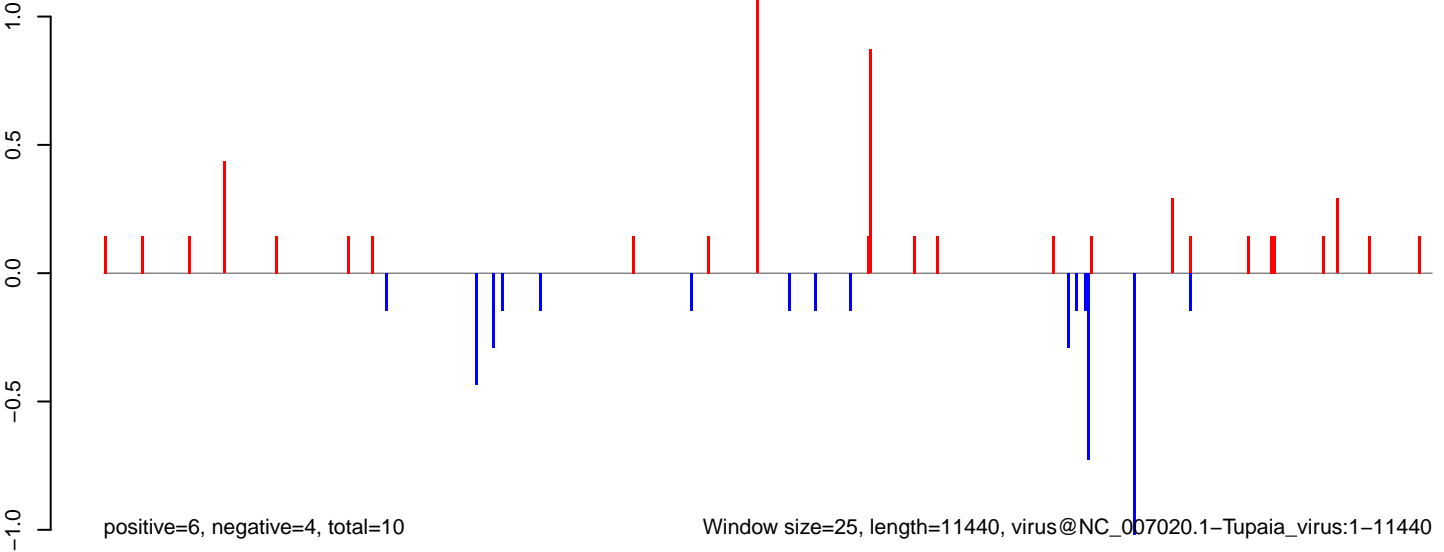
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



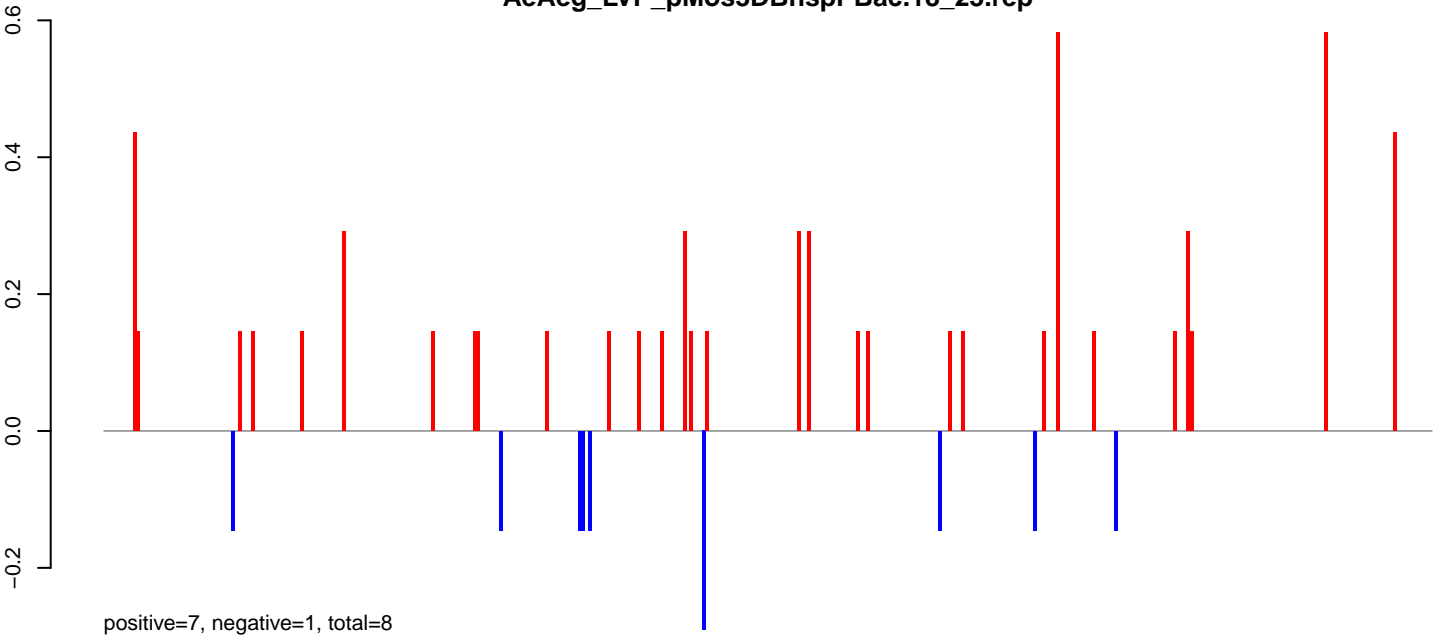
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



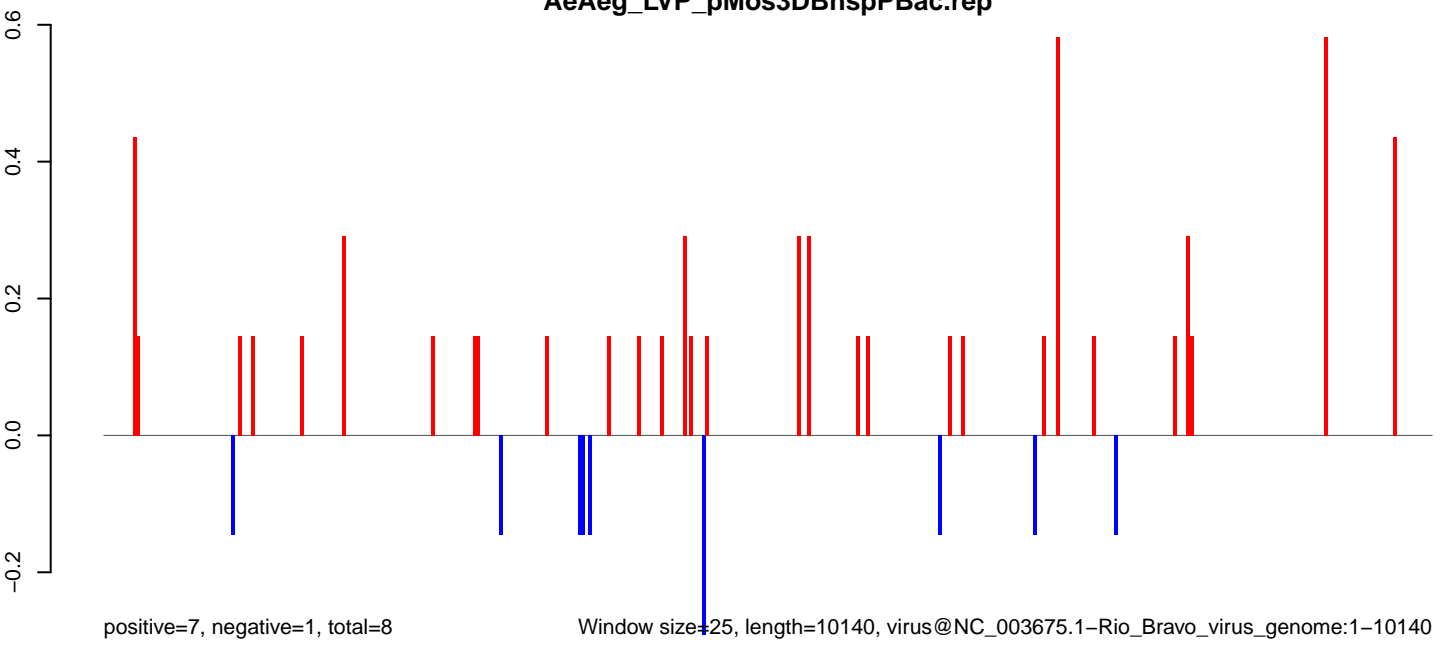
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

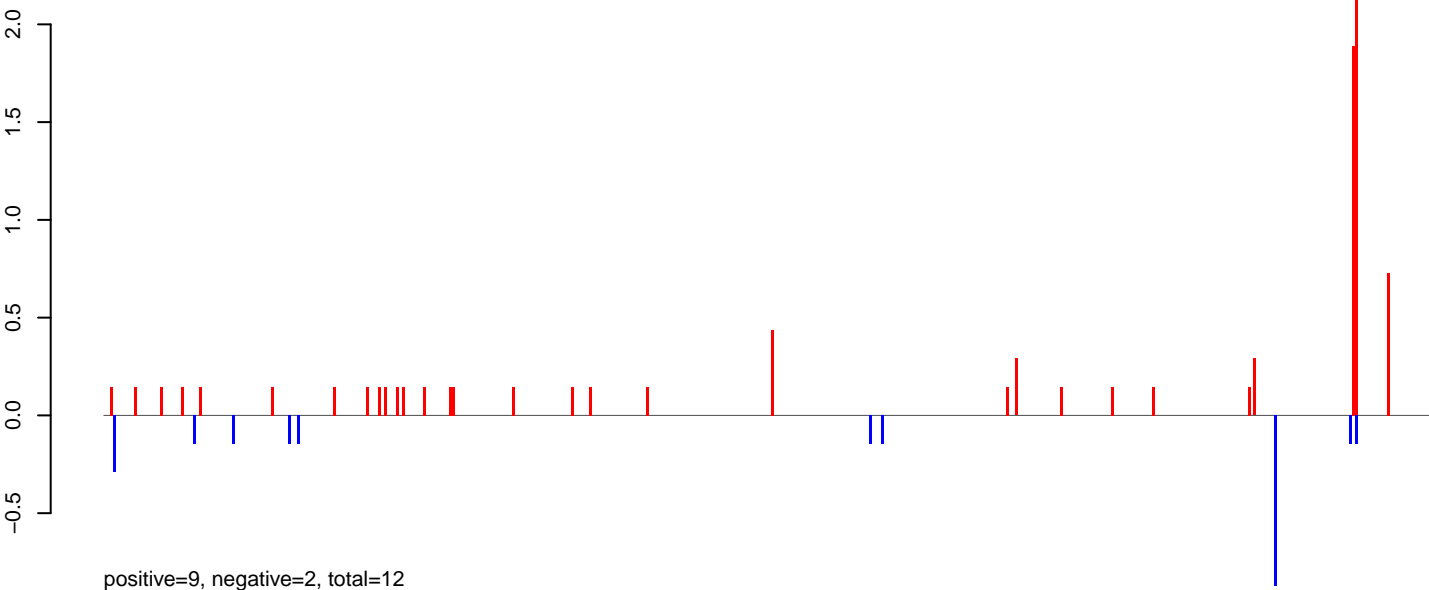


AeAeg_LVP_pMos3DBhspPBac.rep

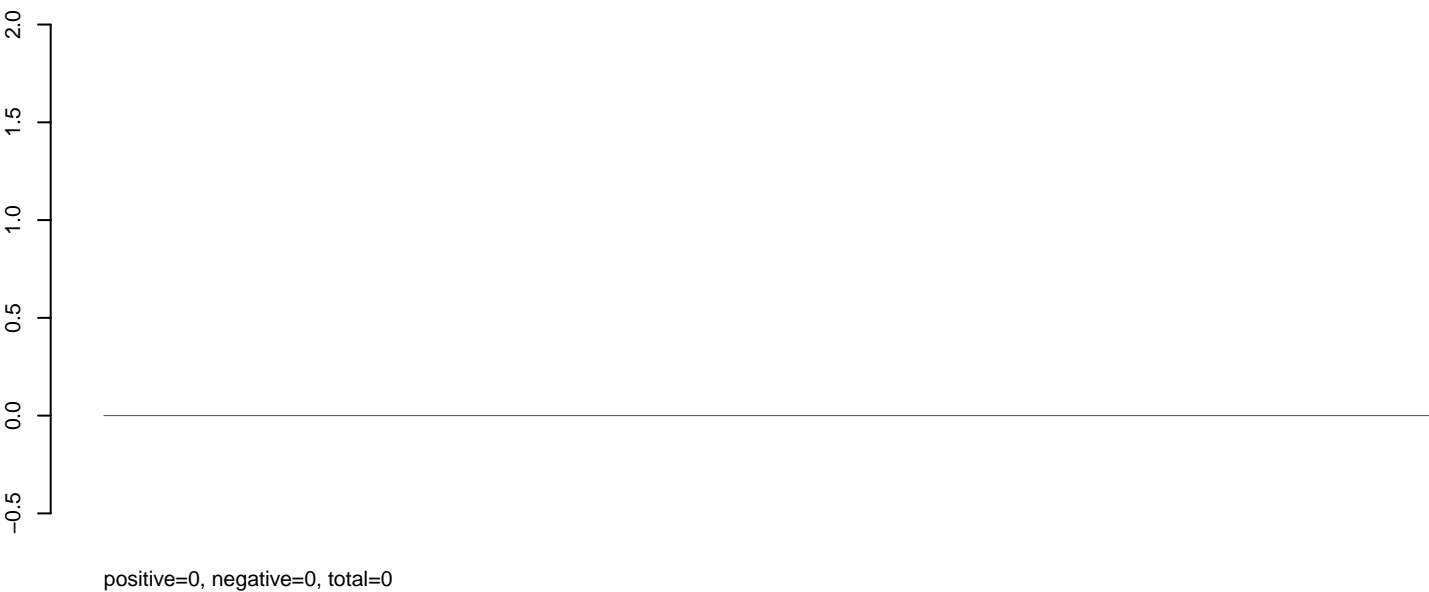


0 2000 4000 6000 8000 10000

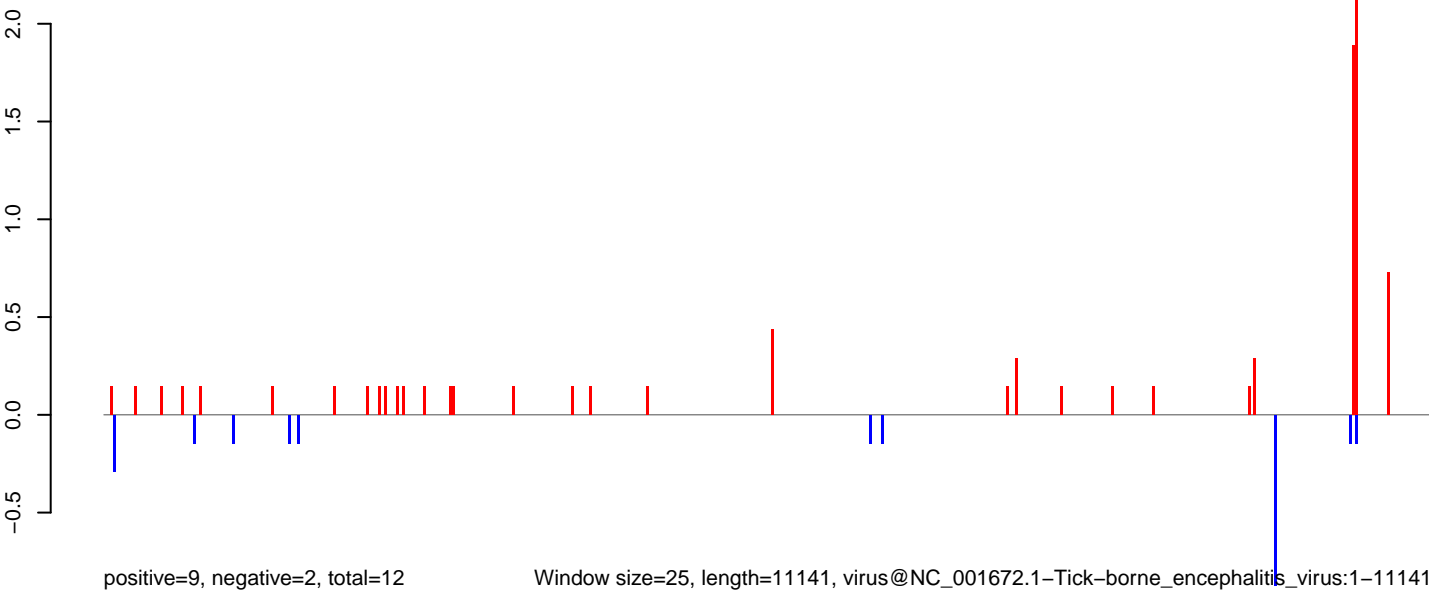
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

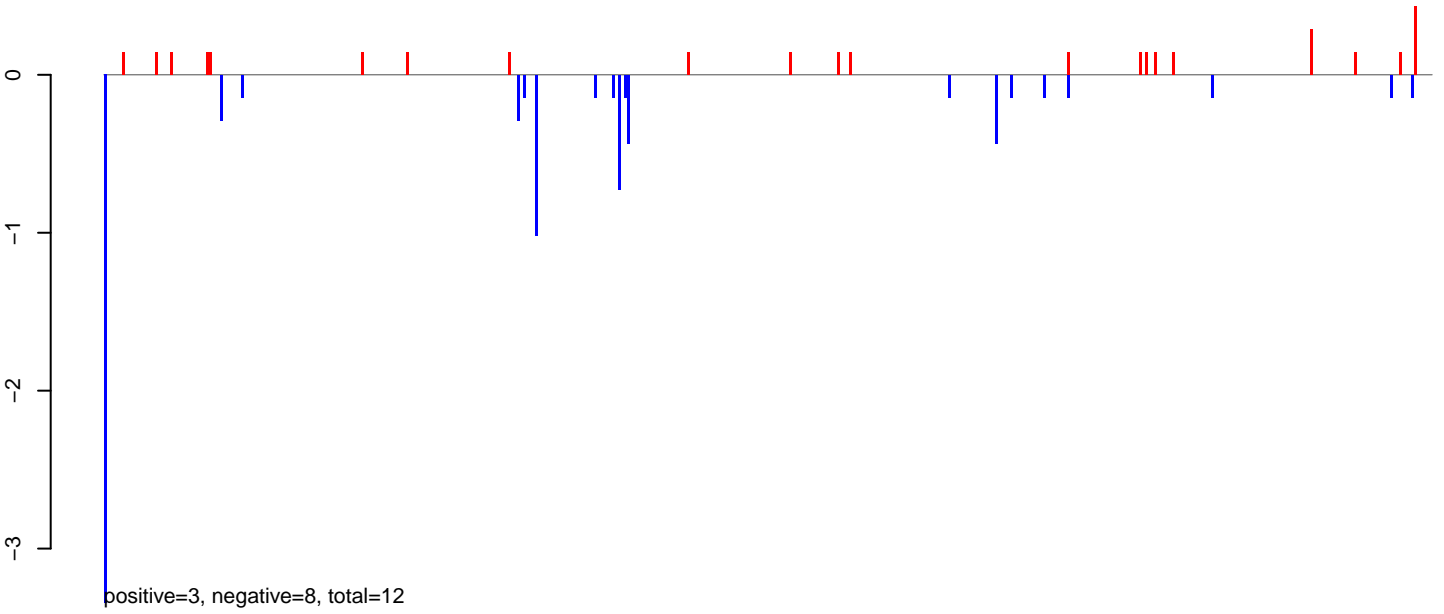


AeAeg_LVP_pMos3DBhspPBac.rep

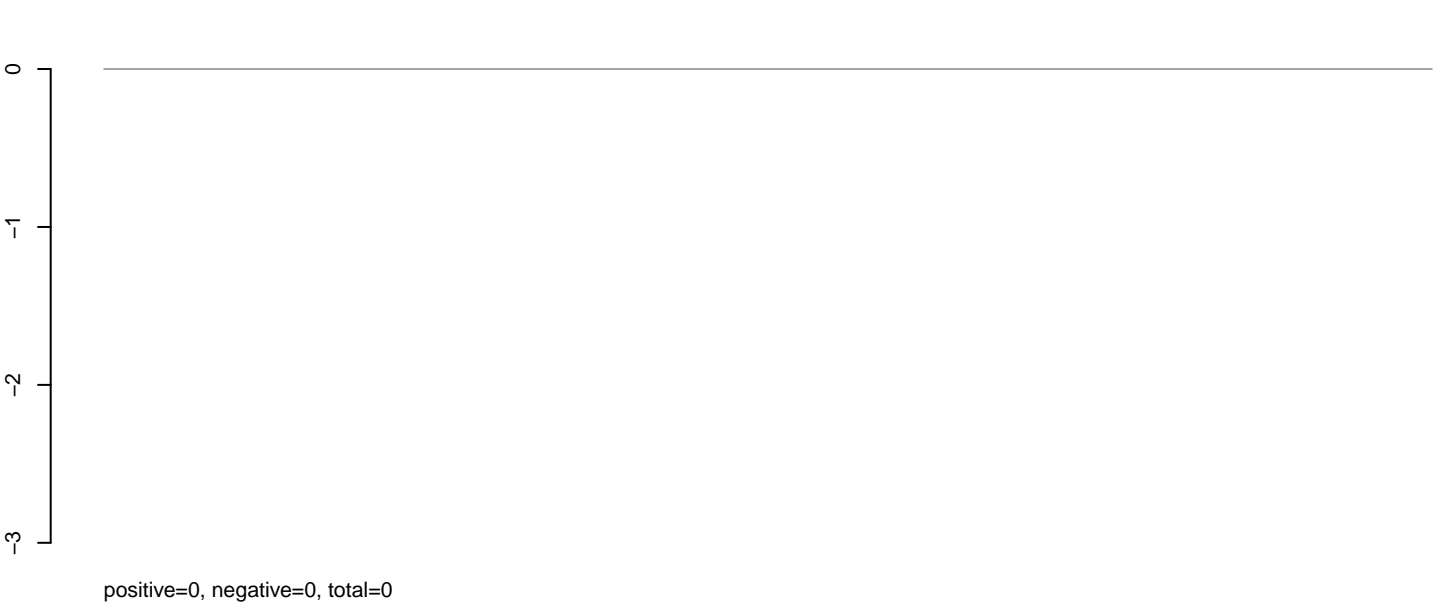


Window size=25, length=11141, virus@NC_001672.1-Tick-borne_encephalitis_virus:1-11141

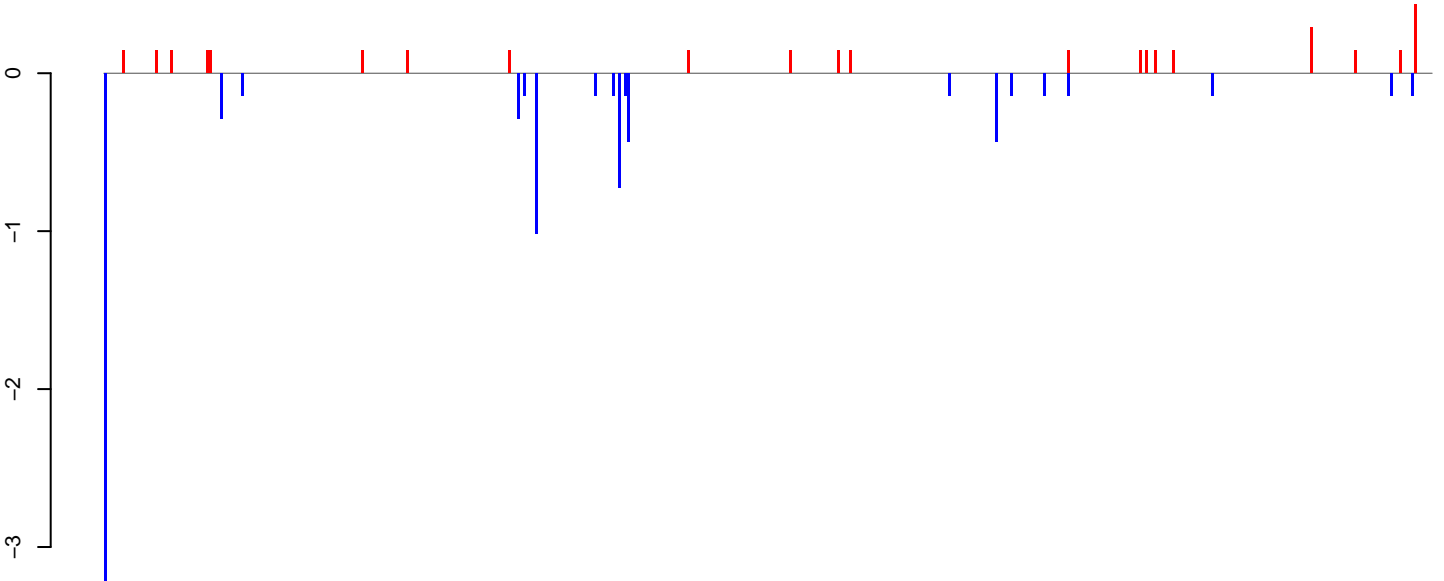
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



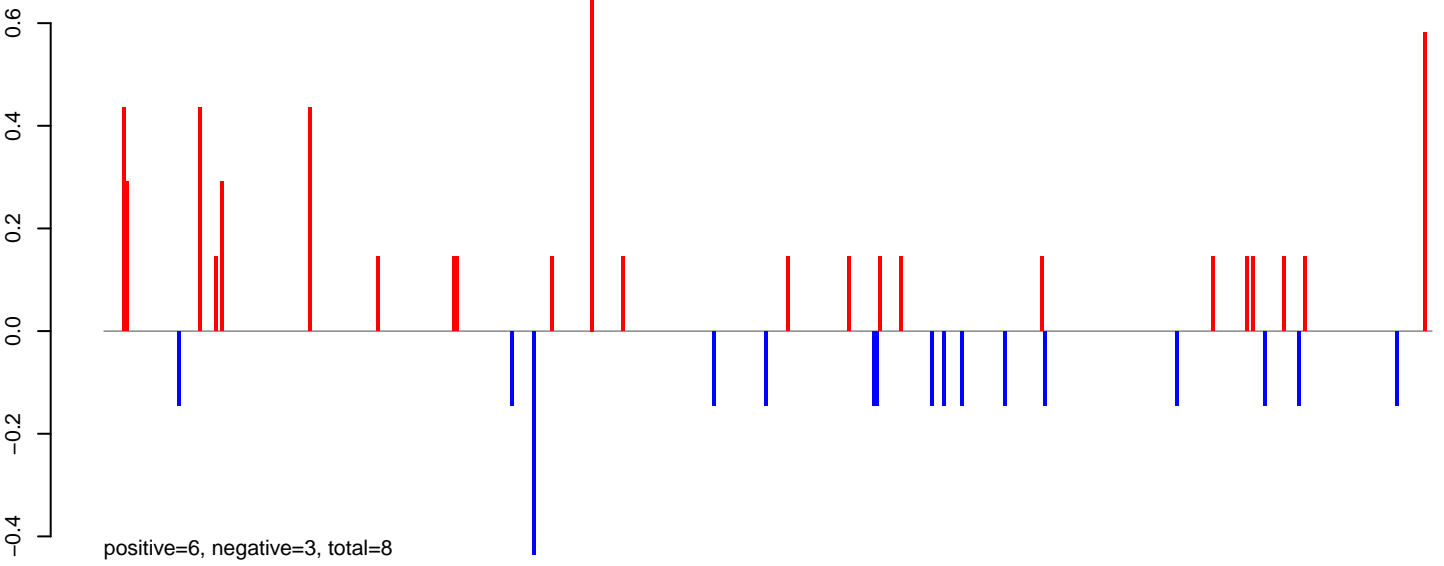
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



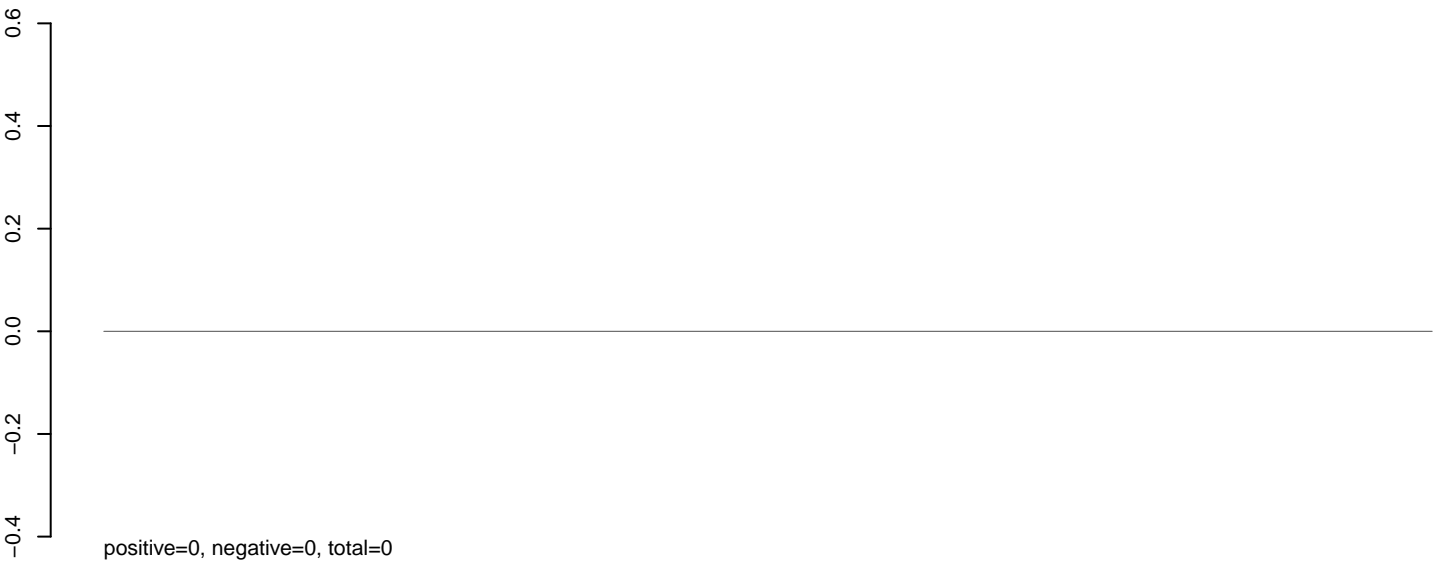
AeAeg_LVP_pMos3DBhspPBac.rep



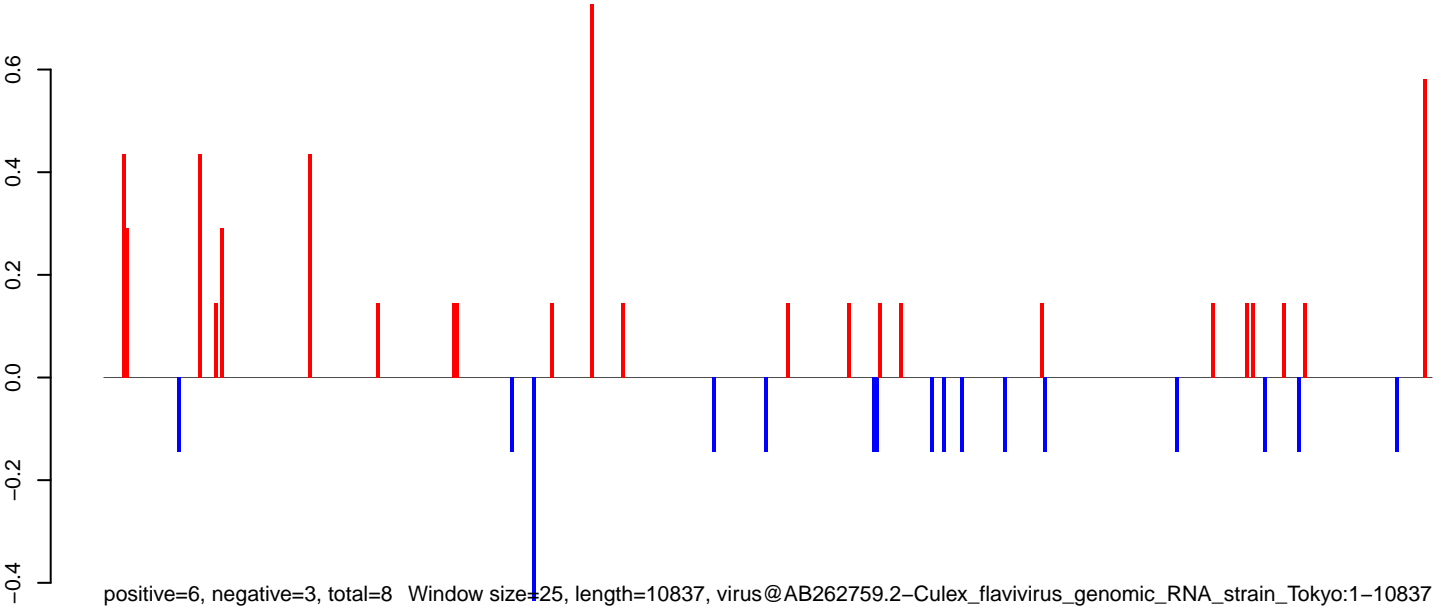
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

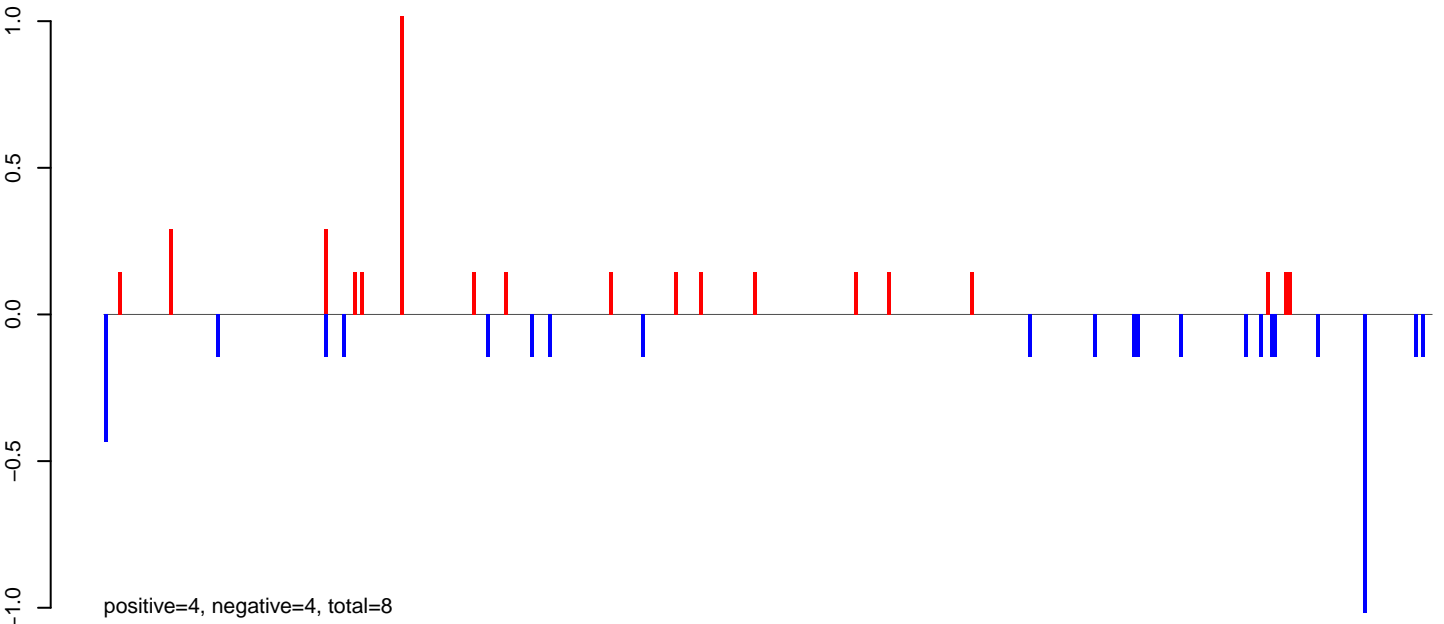


AeAeg_LVP_pMos3DBhspPBac.rep

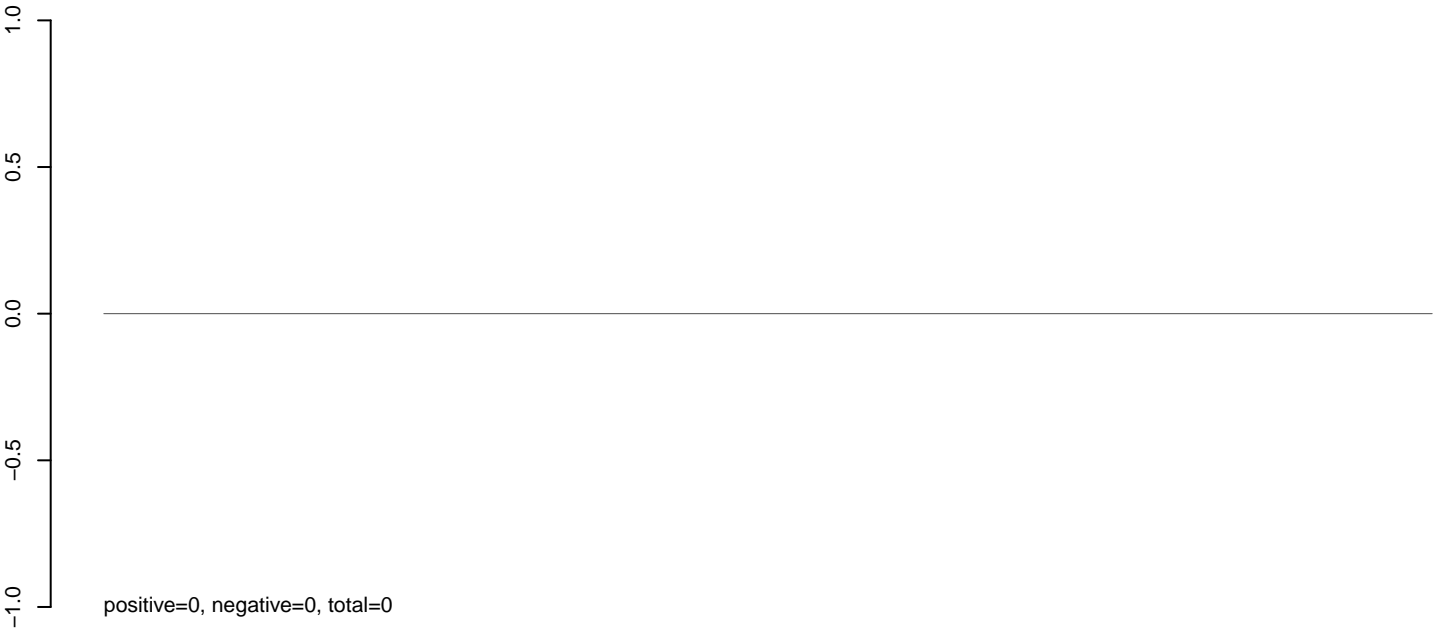


0 2000 4000 6000 8000 10000

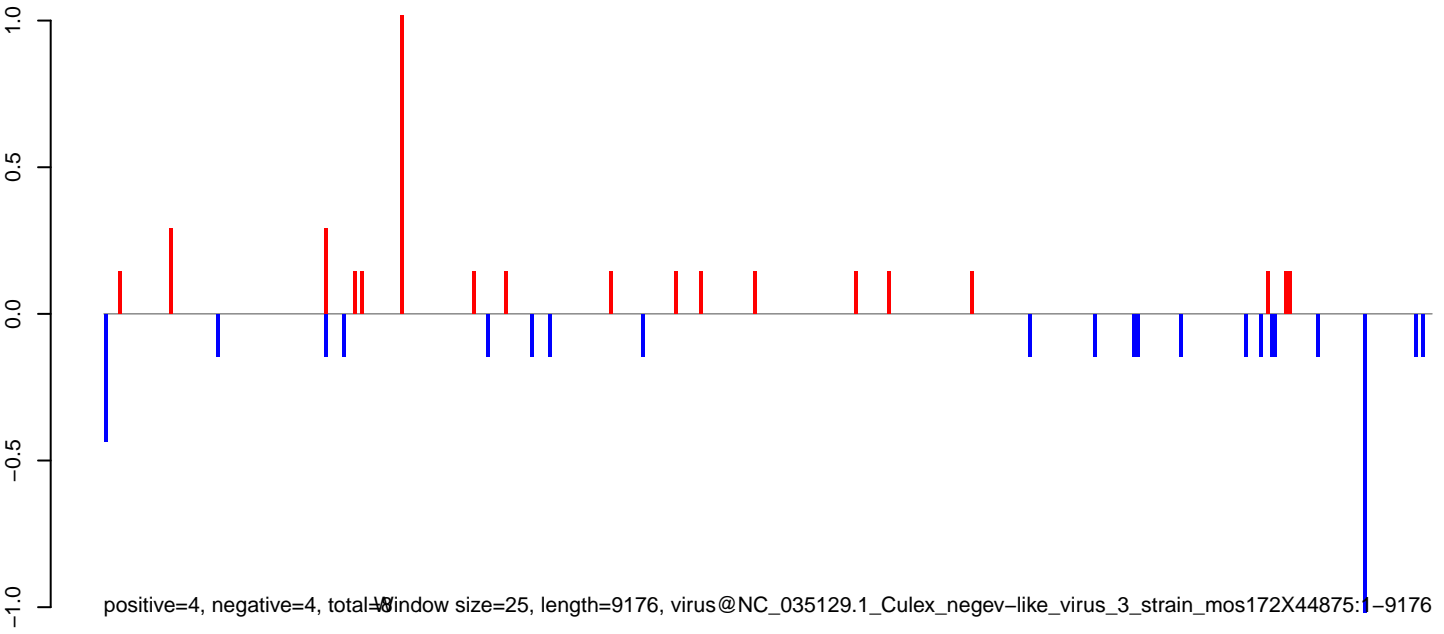
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

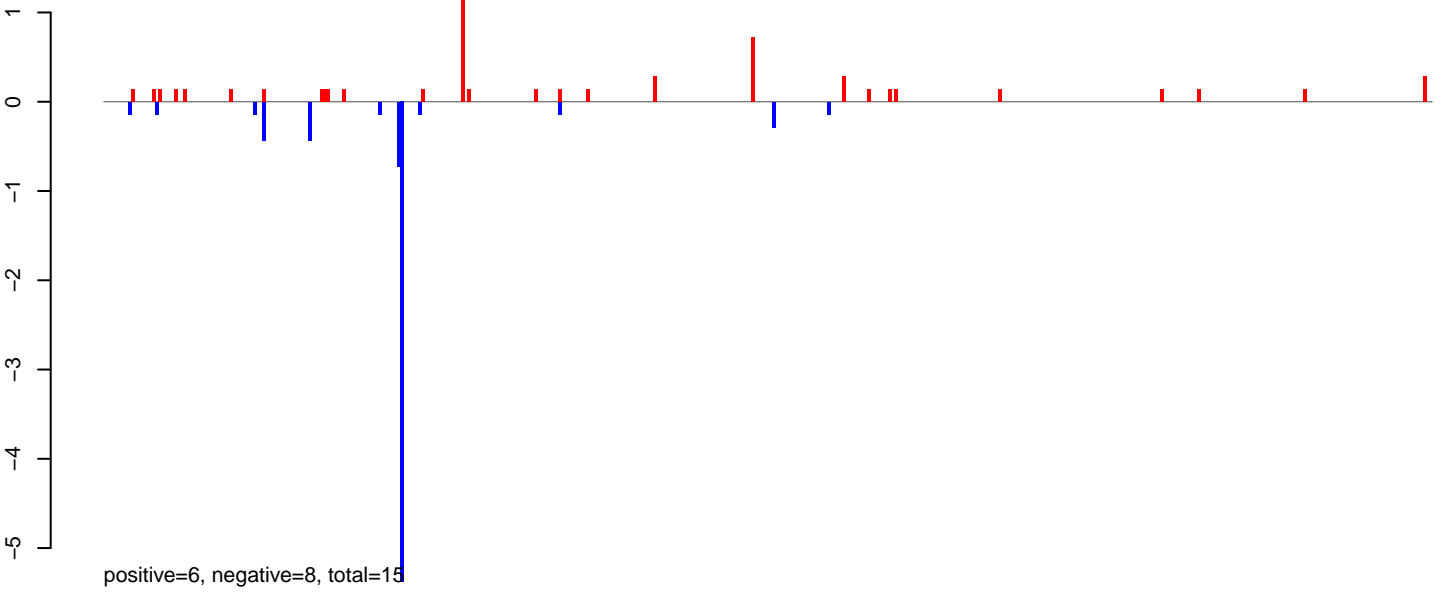


AeAeg_LVP_pMos3DBhspPBac.rep

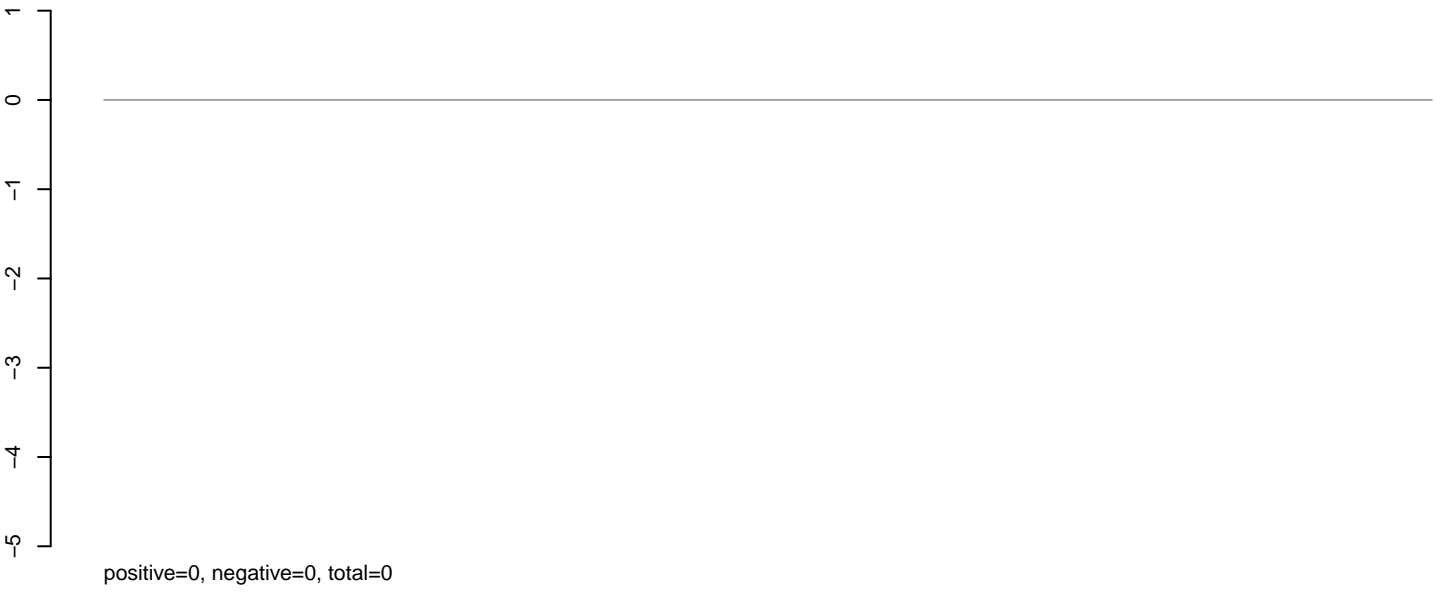


Window size=25, length=9176, virus@NC_035129.1_Culex_negev-like_virus_3_strain_mos172X44875:1-9176

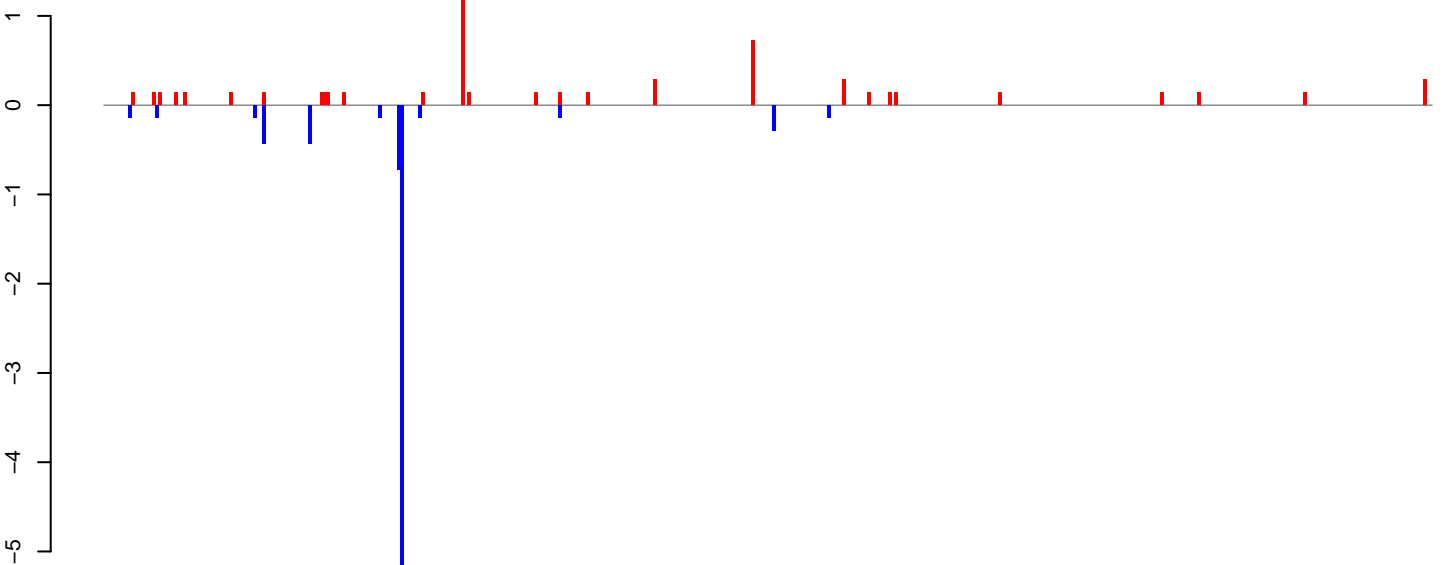
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



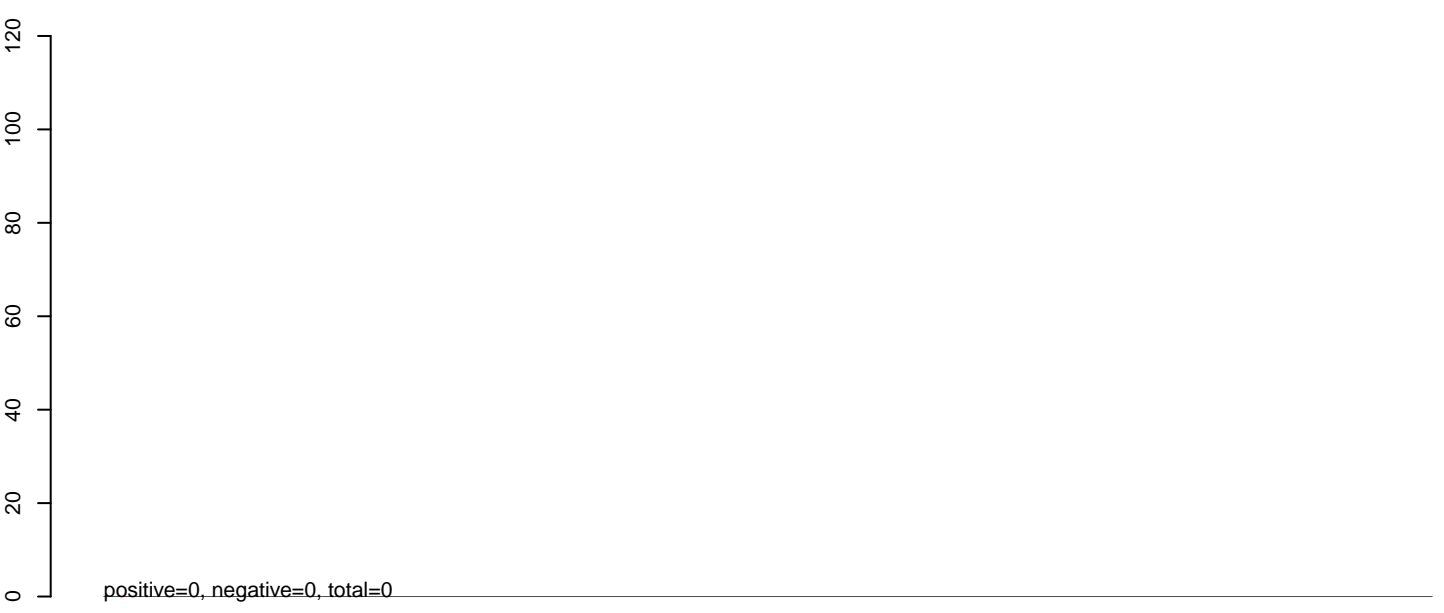
AeAeg_LVP_pMos3DBhspPBac.rep



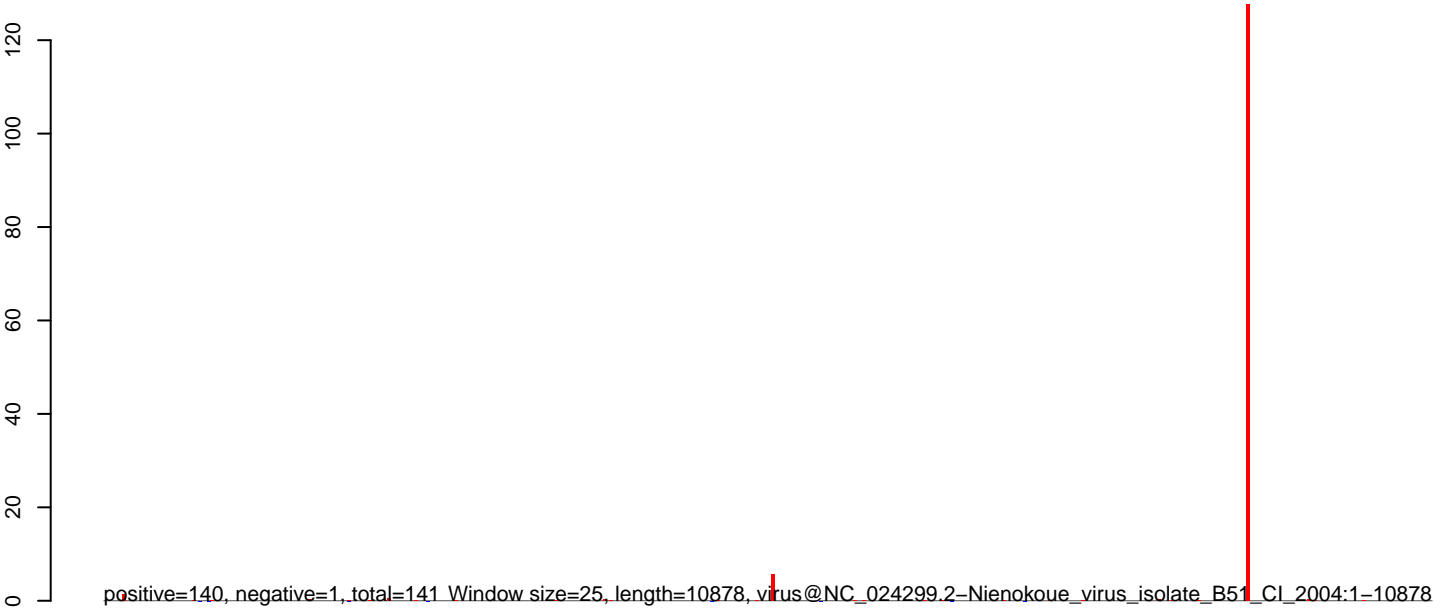
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

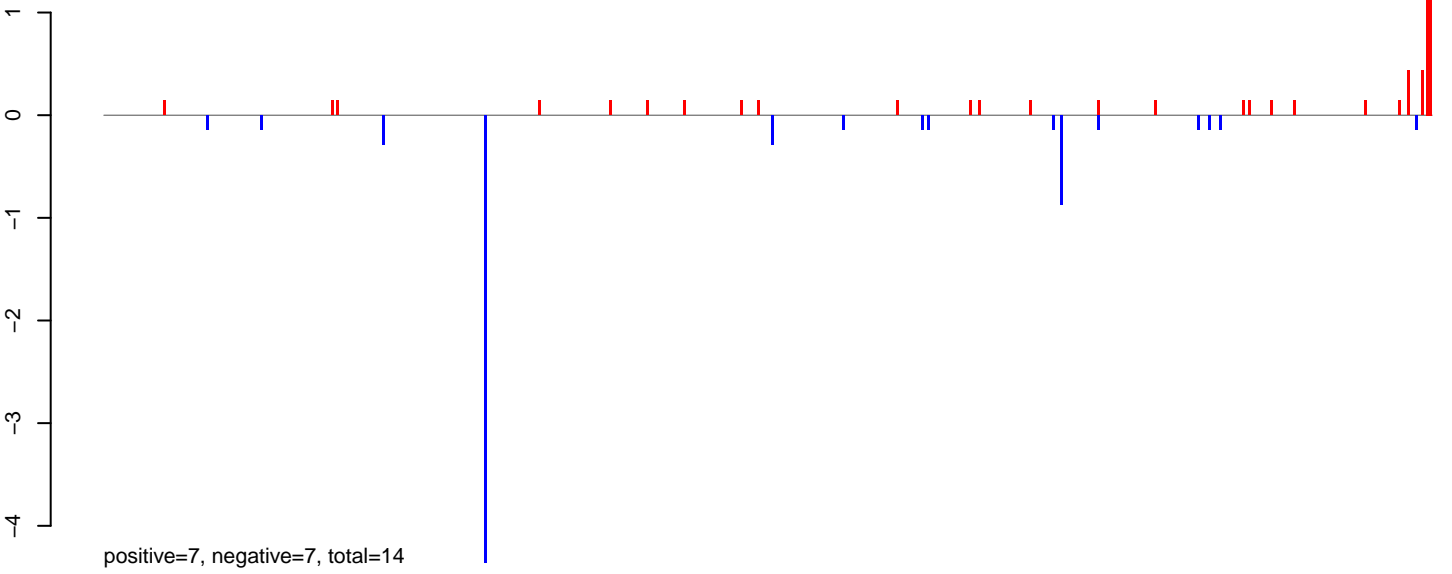


AeAeg_LVP_pMos3DBhspPBac.rep

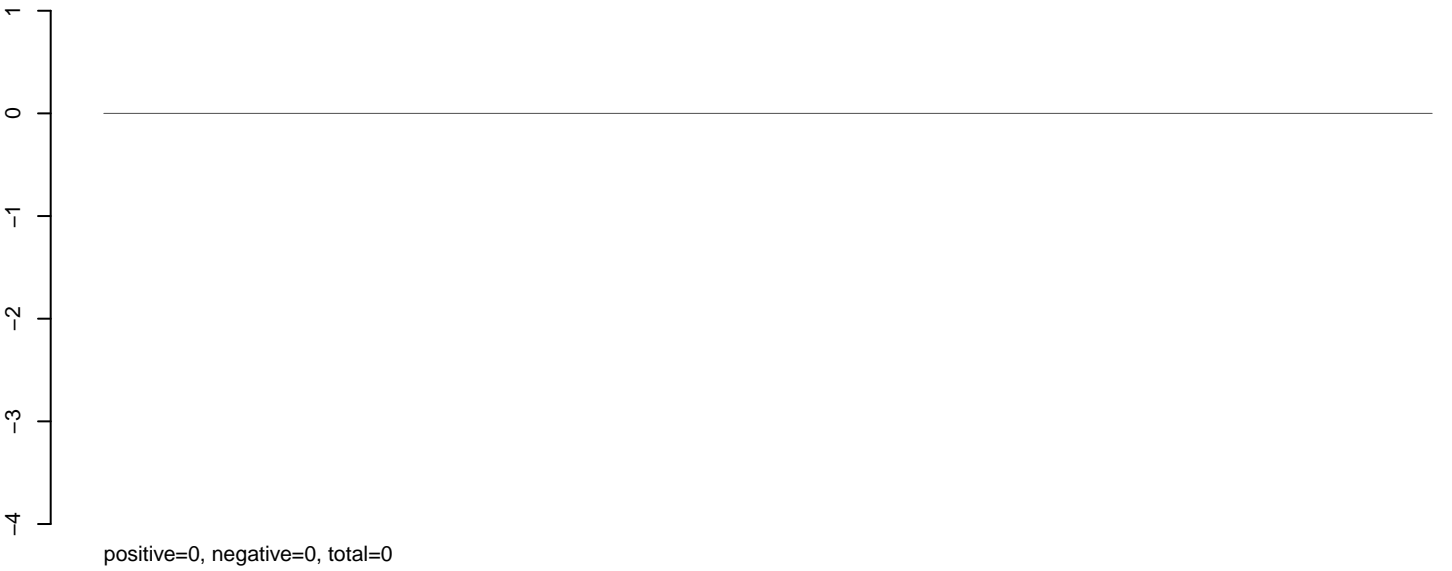


0 2000 4000 6000 8000 10000

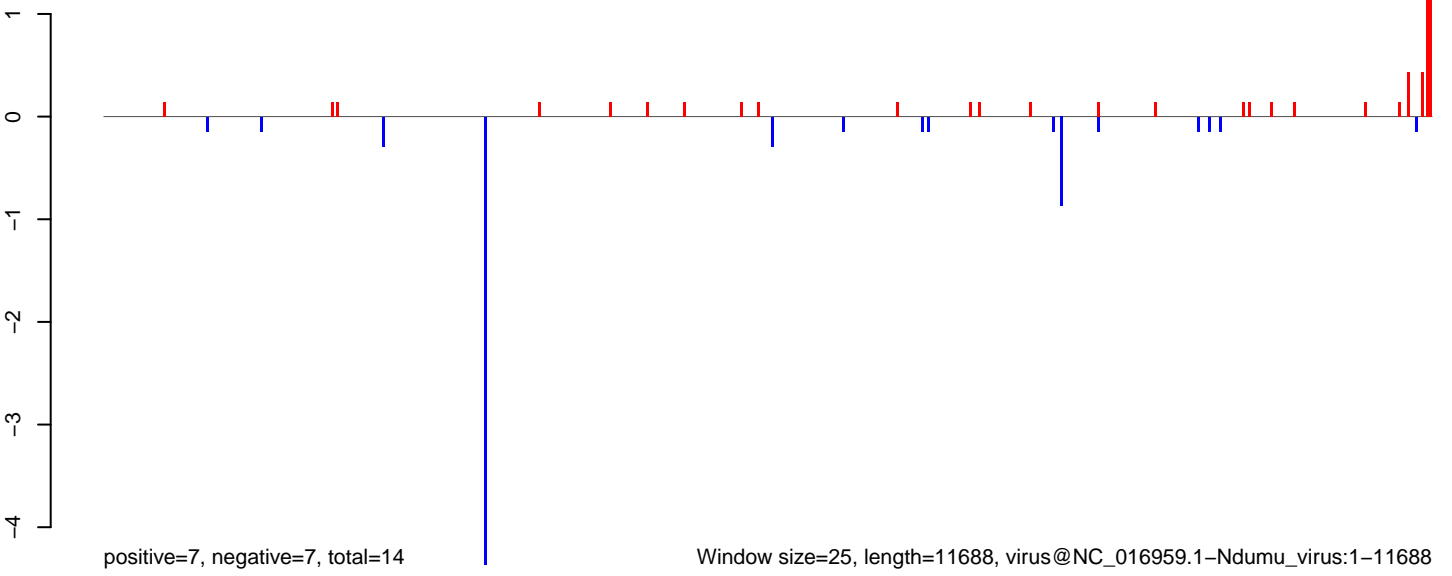
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



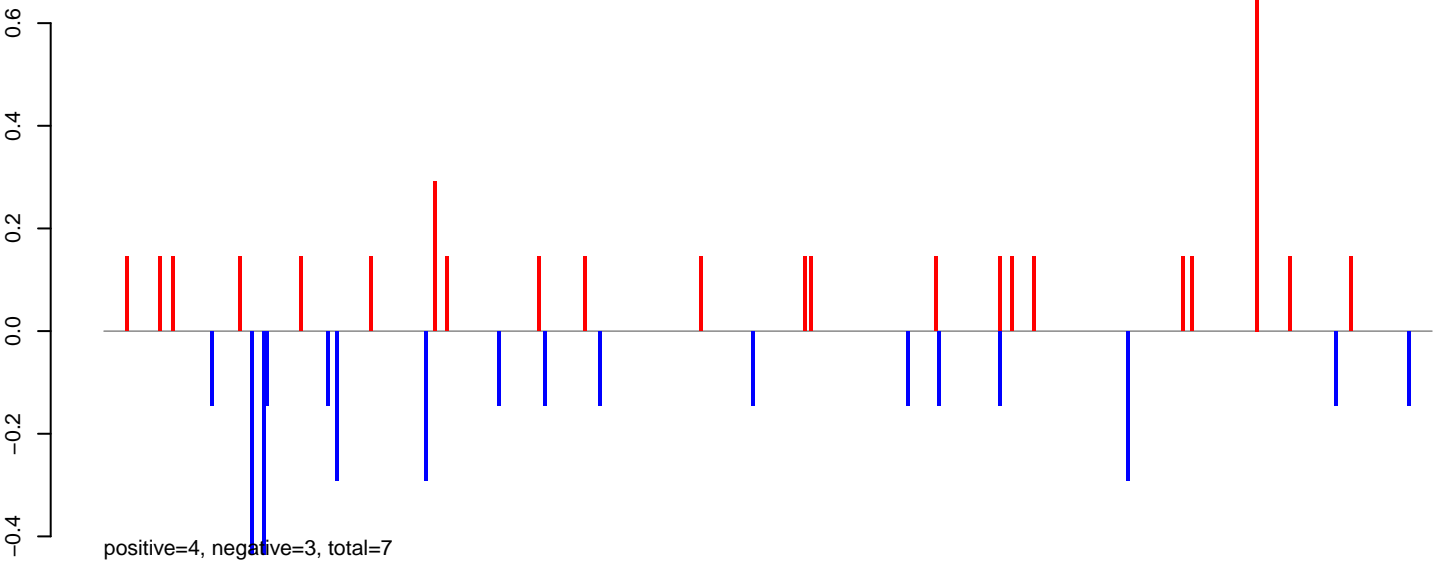
AeAeg_LVP_pMos3DBhspPBac.rep



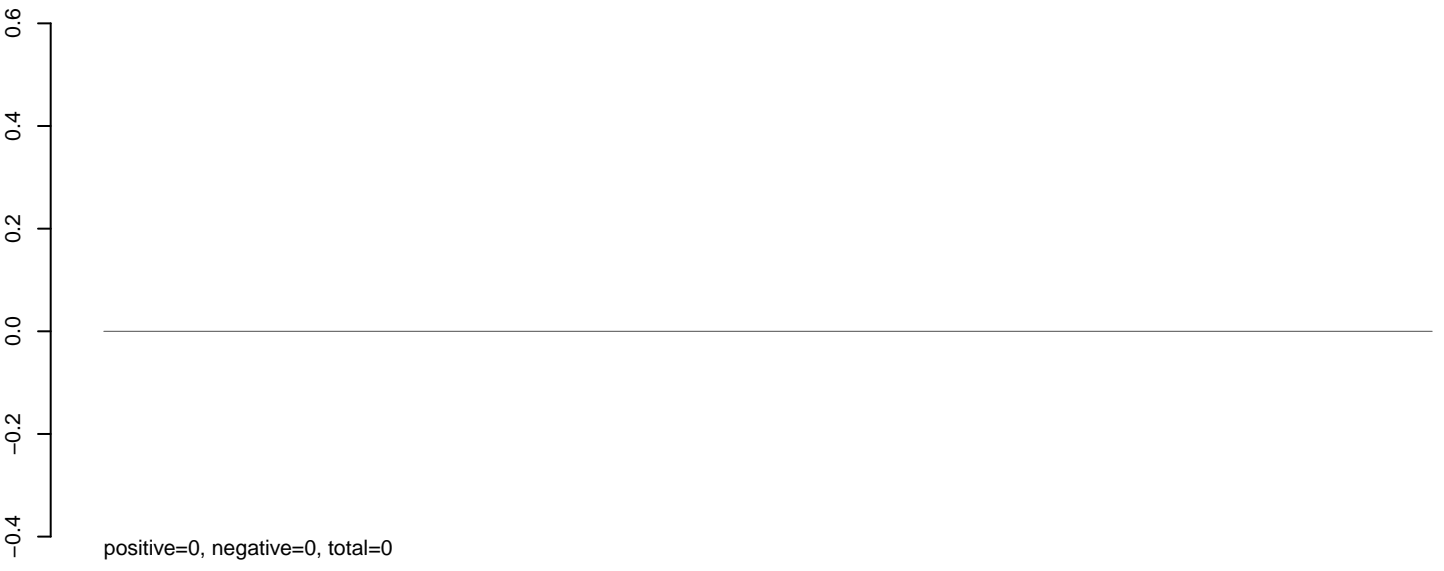
Window size=25, length=11688, virus@NC_016959.1-Ndumu_virus:1-11688

0 2000 4000 6000 8000 10000 12000

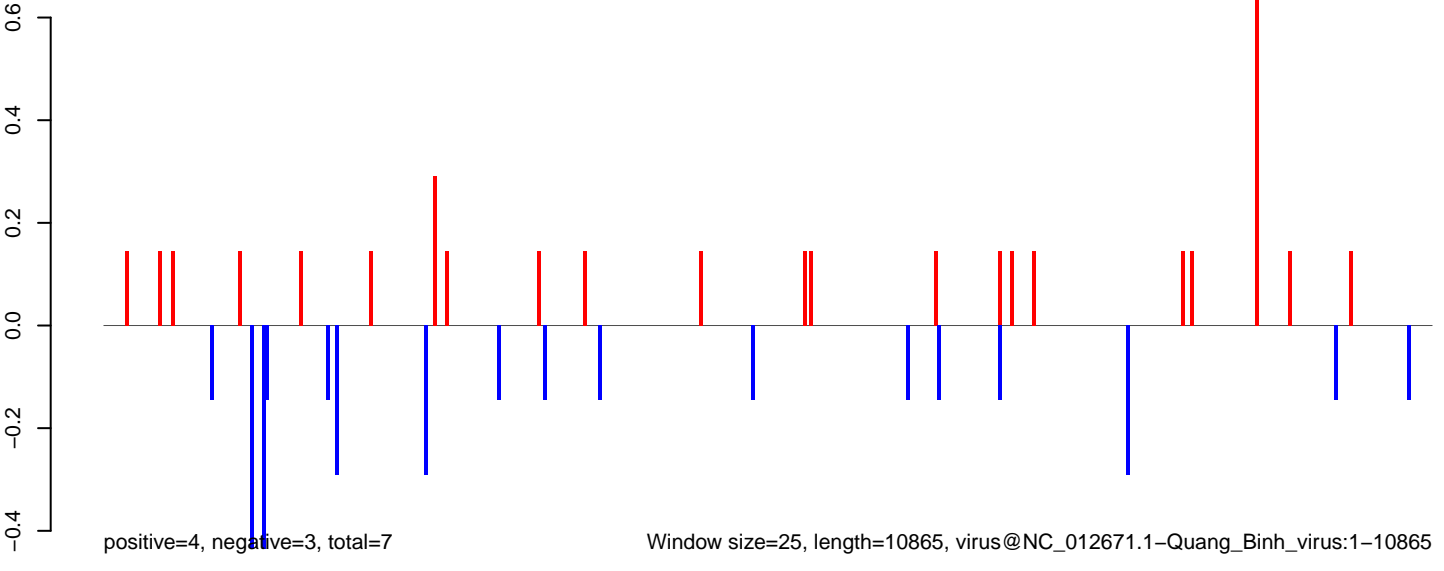
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



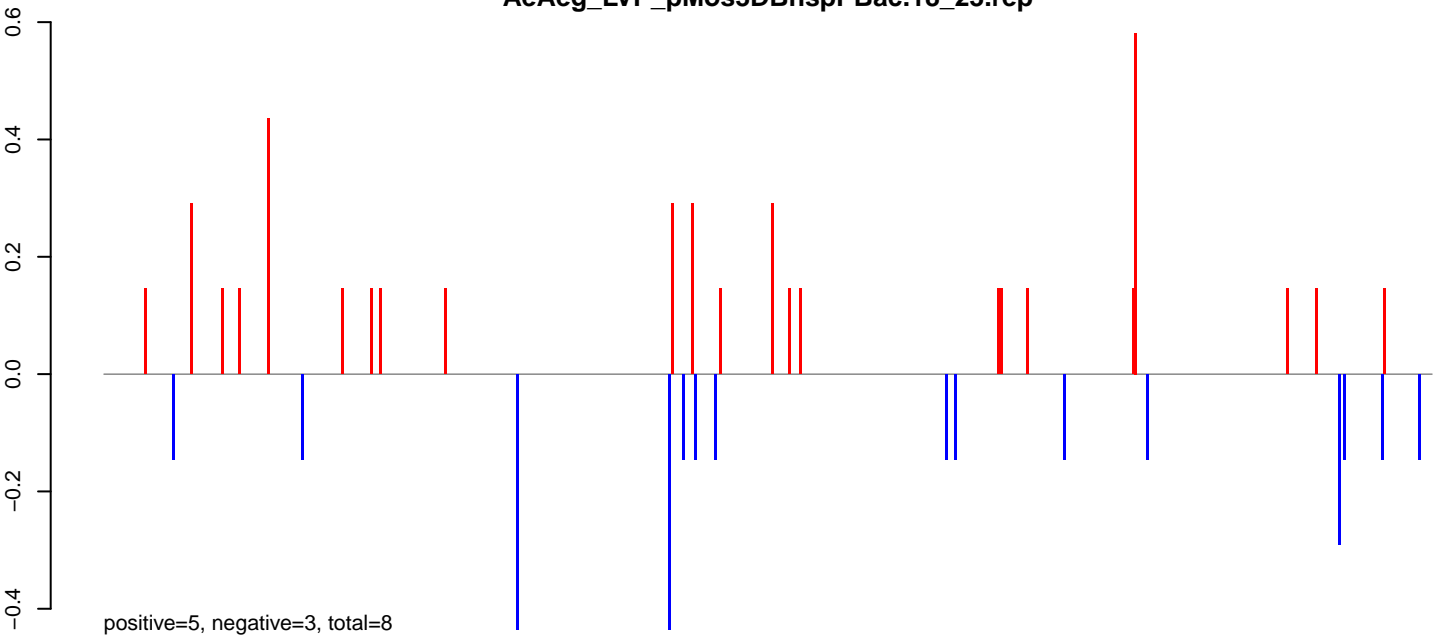
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



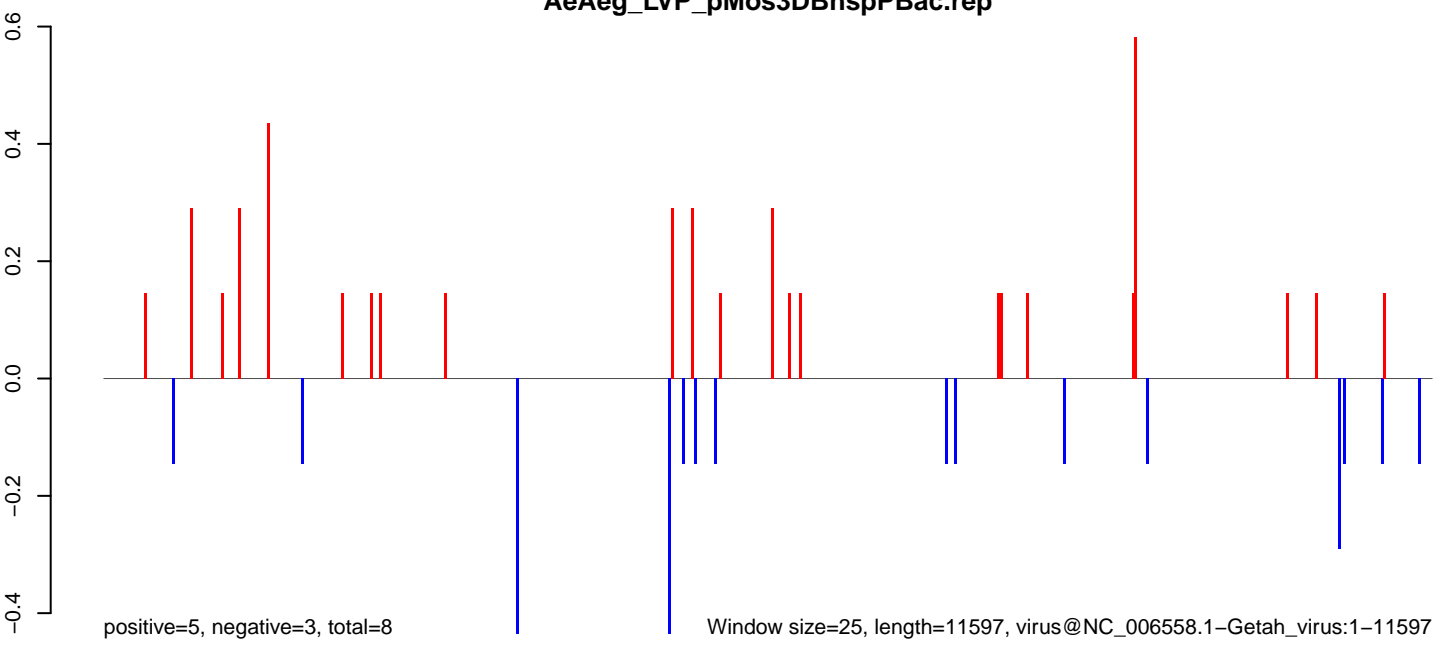
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

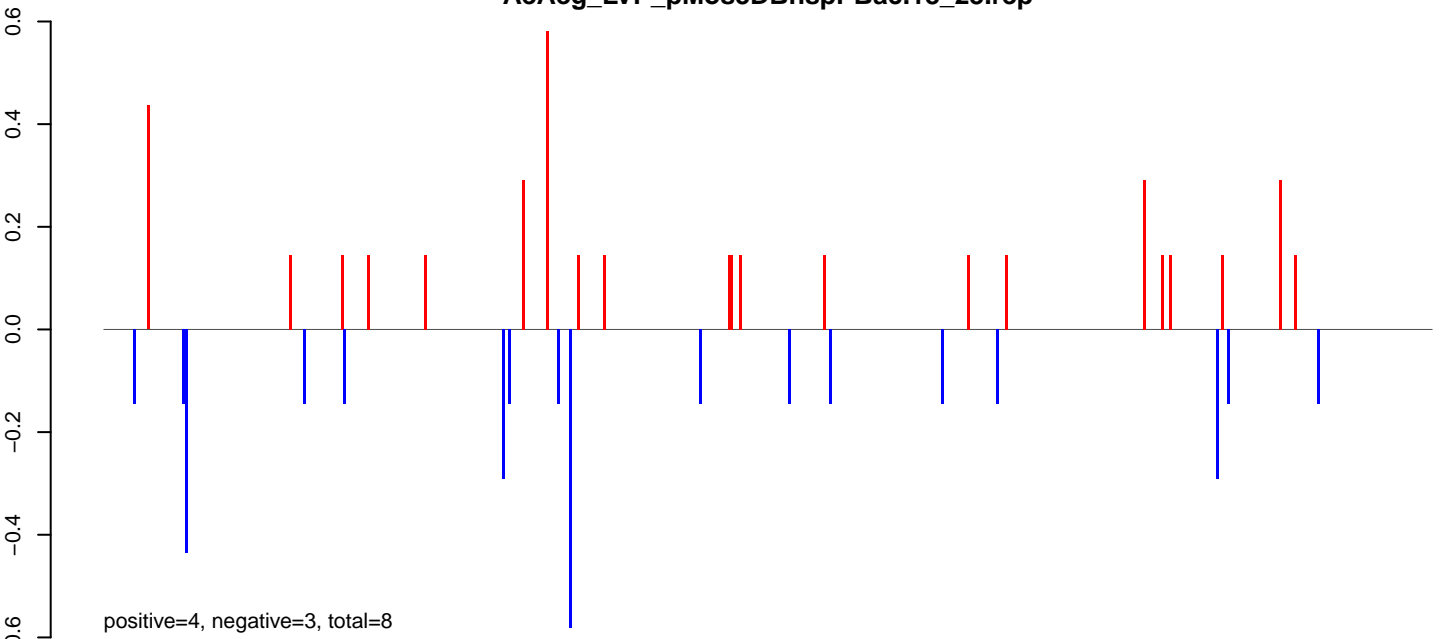


AeAeg_LVP_pMos3DBhspPBac.rep

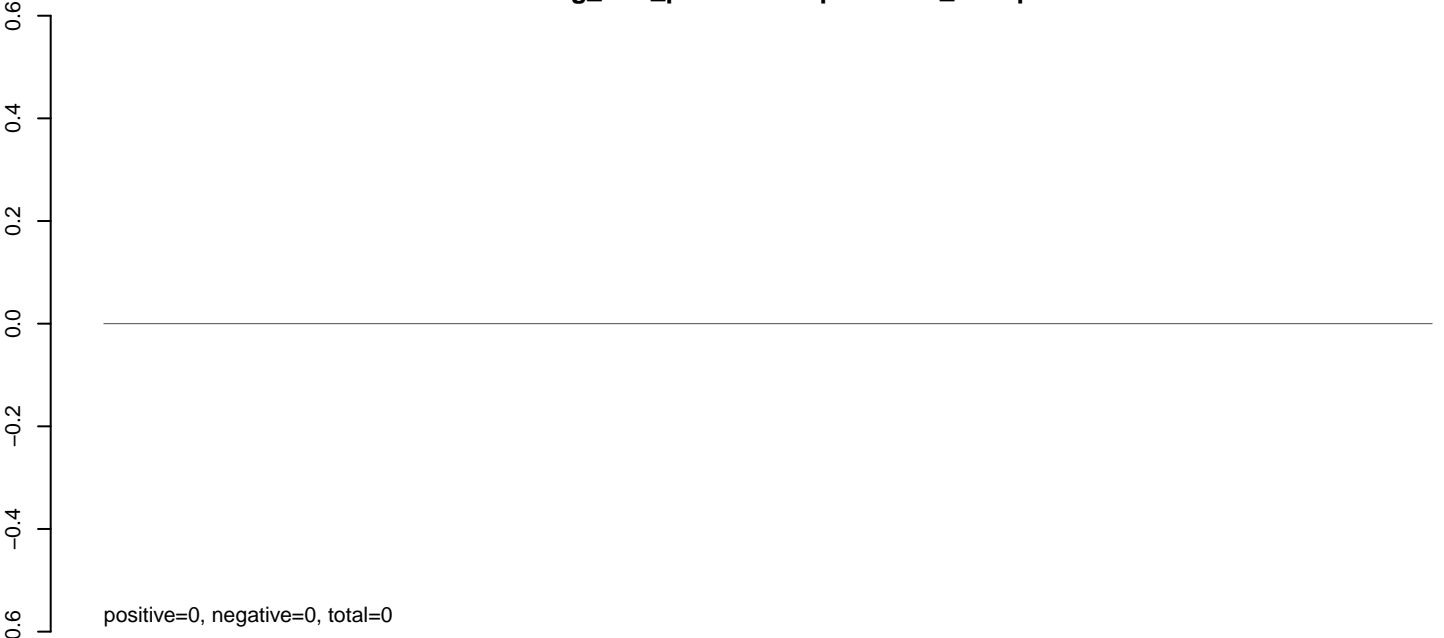


0 2000 4000 6000 8000 10000 12000

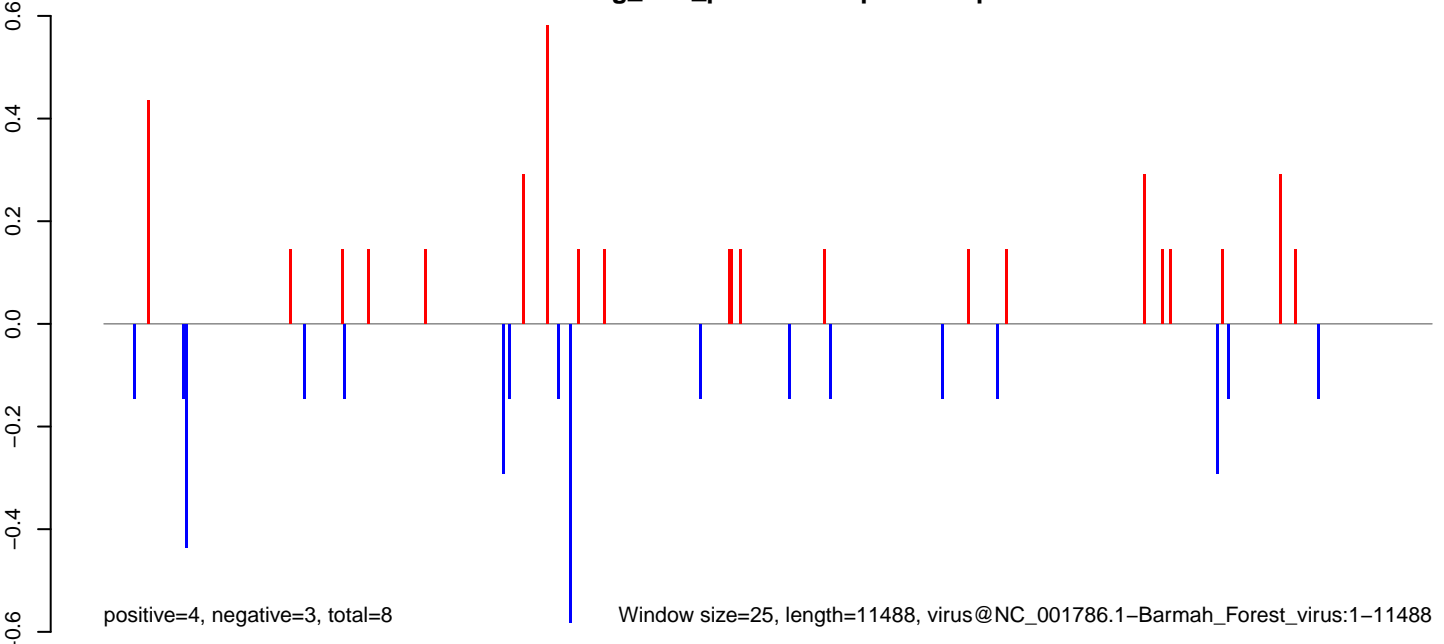
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

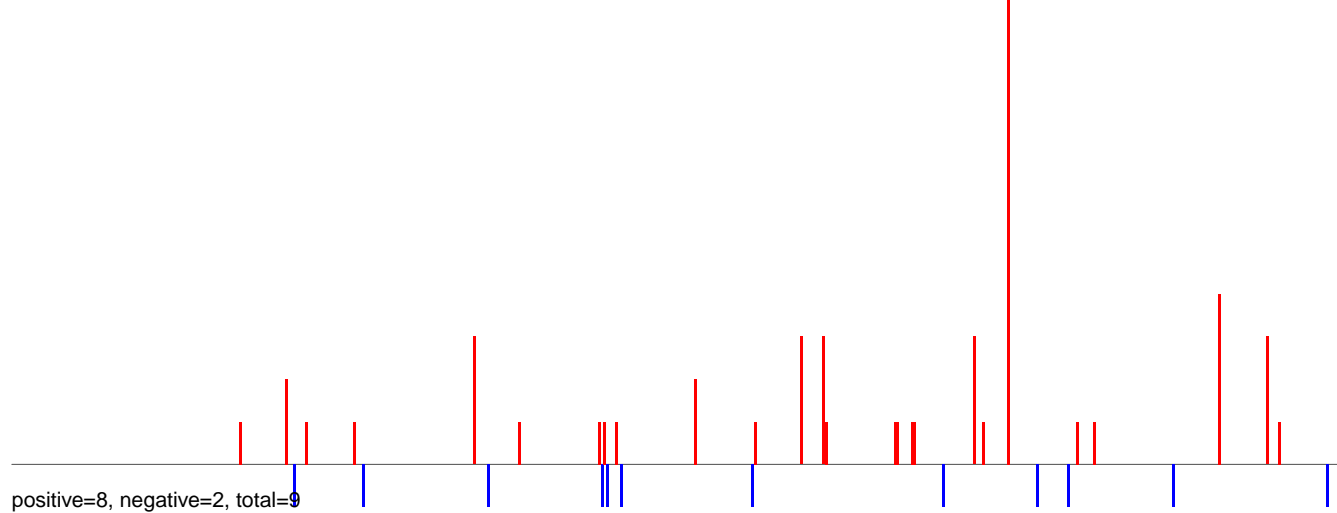


AeAeg_LVP_pMos3DBhspPBac.rep



0 2000 4000 6000 8000 10000

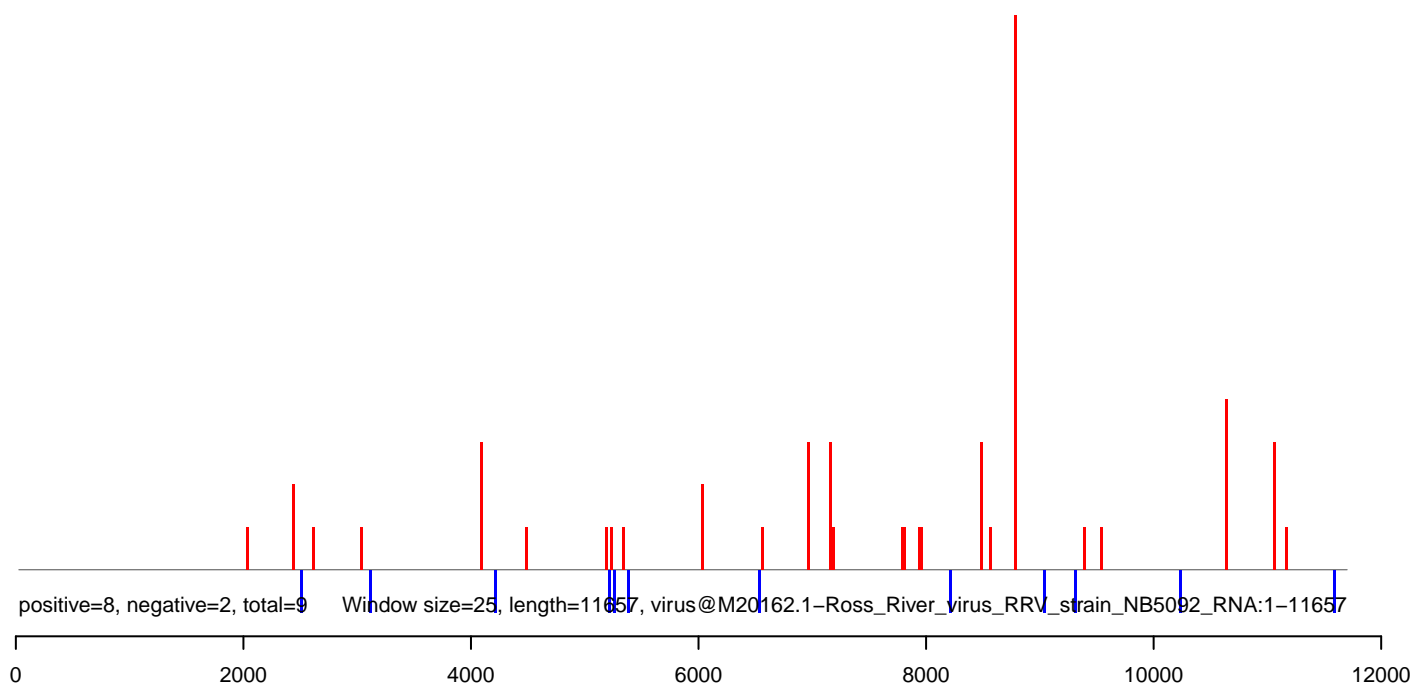
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



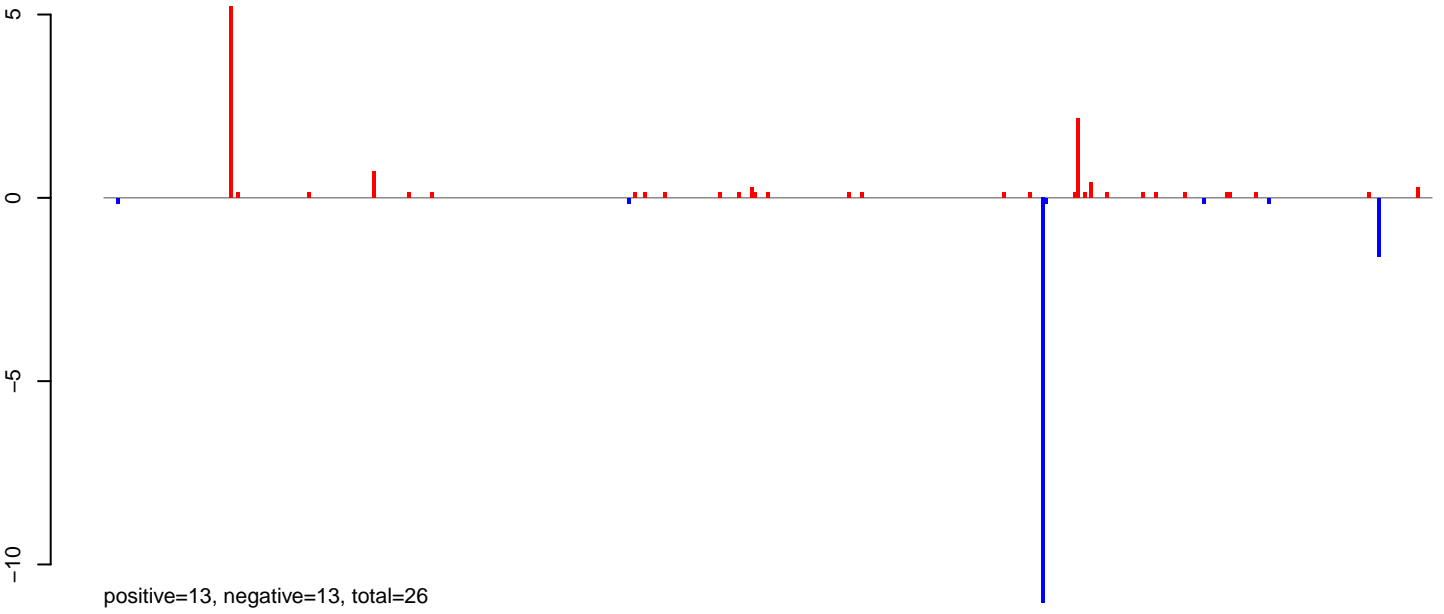
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



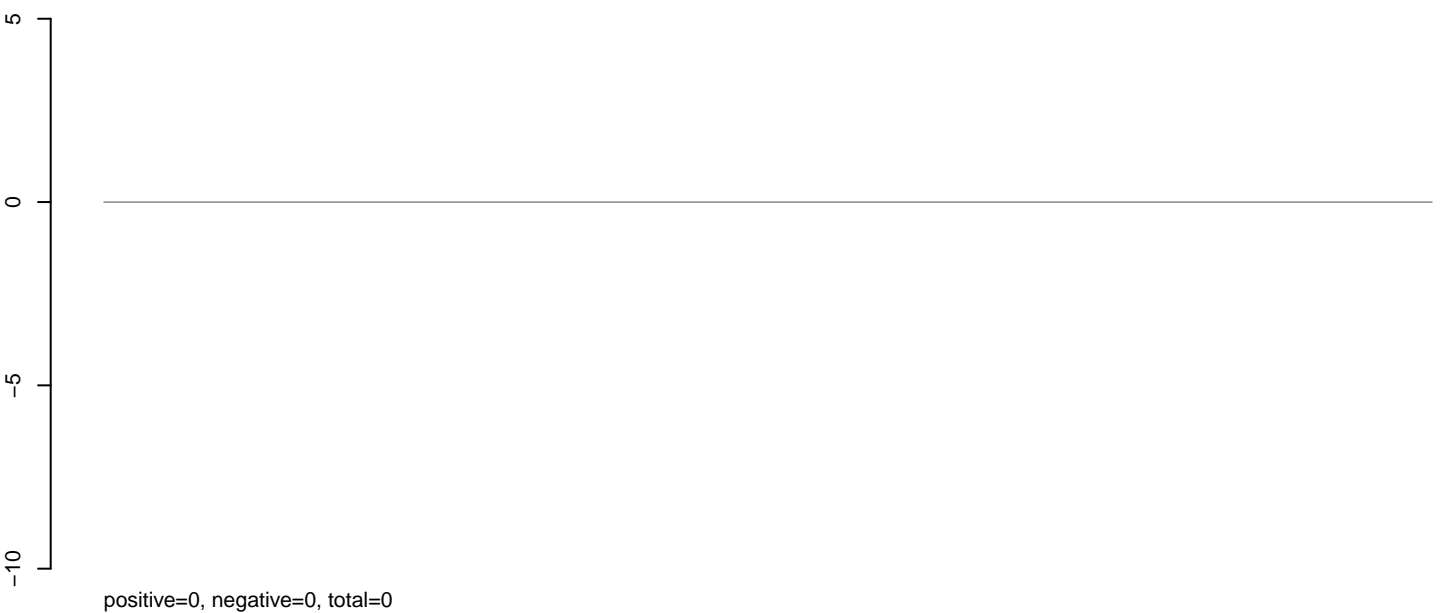
AeAeg_LVP_pMos3DBhspPBac.rep



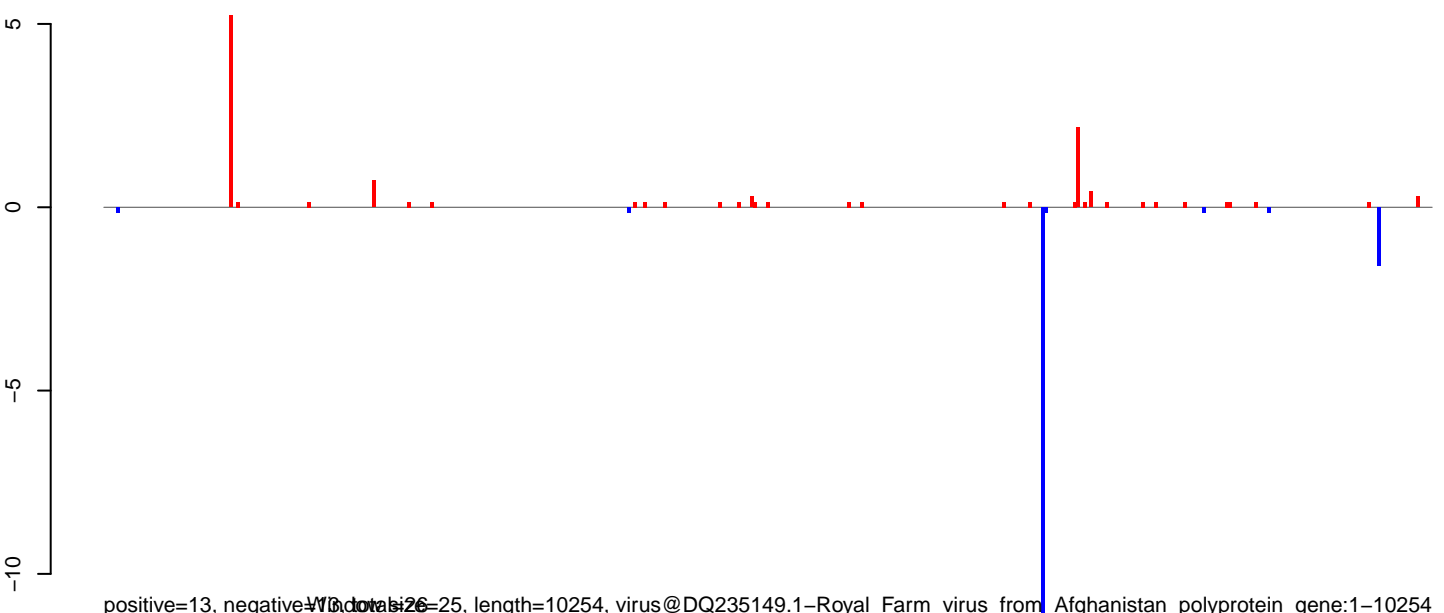
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

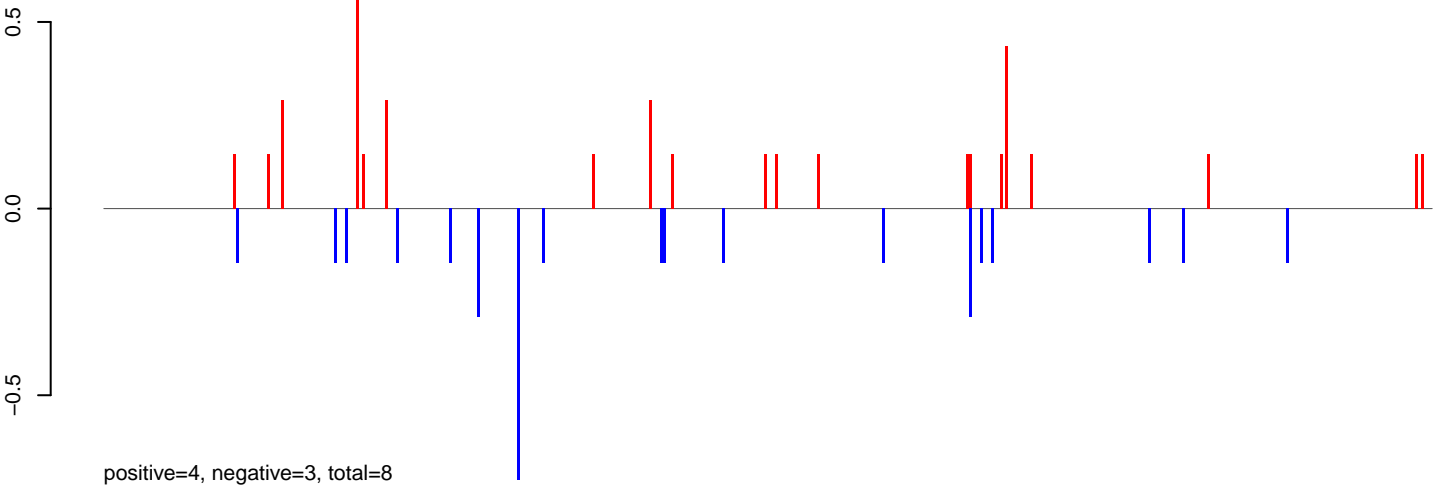


AeAeg_LVP_pMos3DBhspPBac.rep

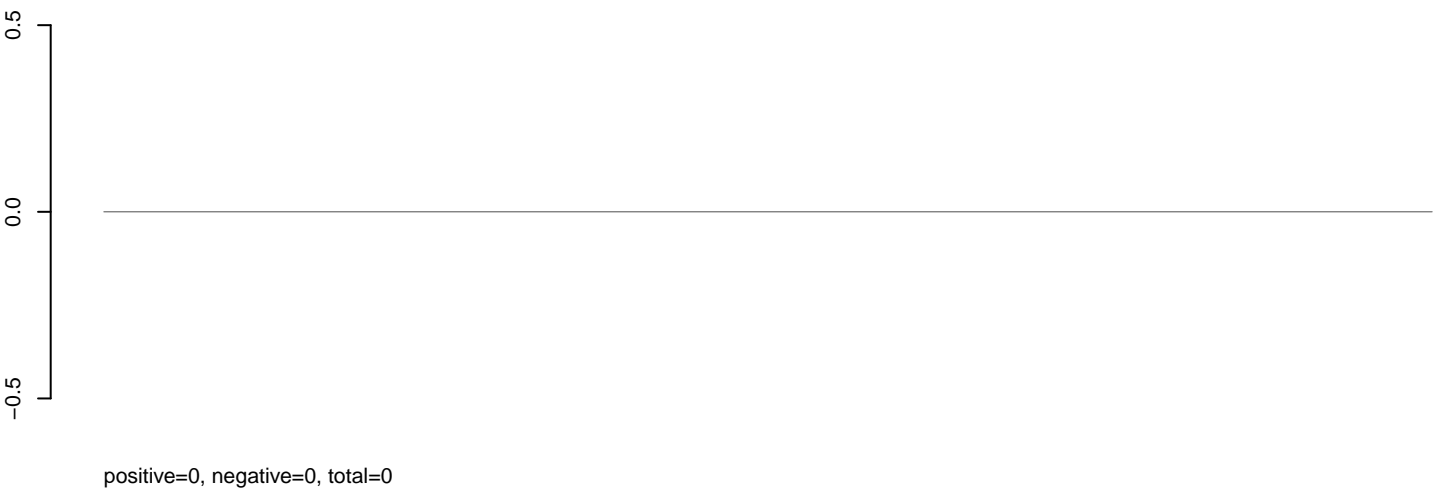


Window size=25, length=10254, virus@DQ235149.1-Royal_Farm_virus_from_Afghanistan_polyprotein_gene:1-10254

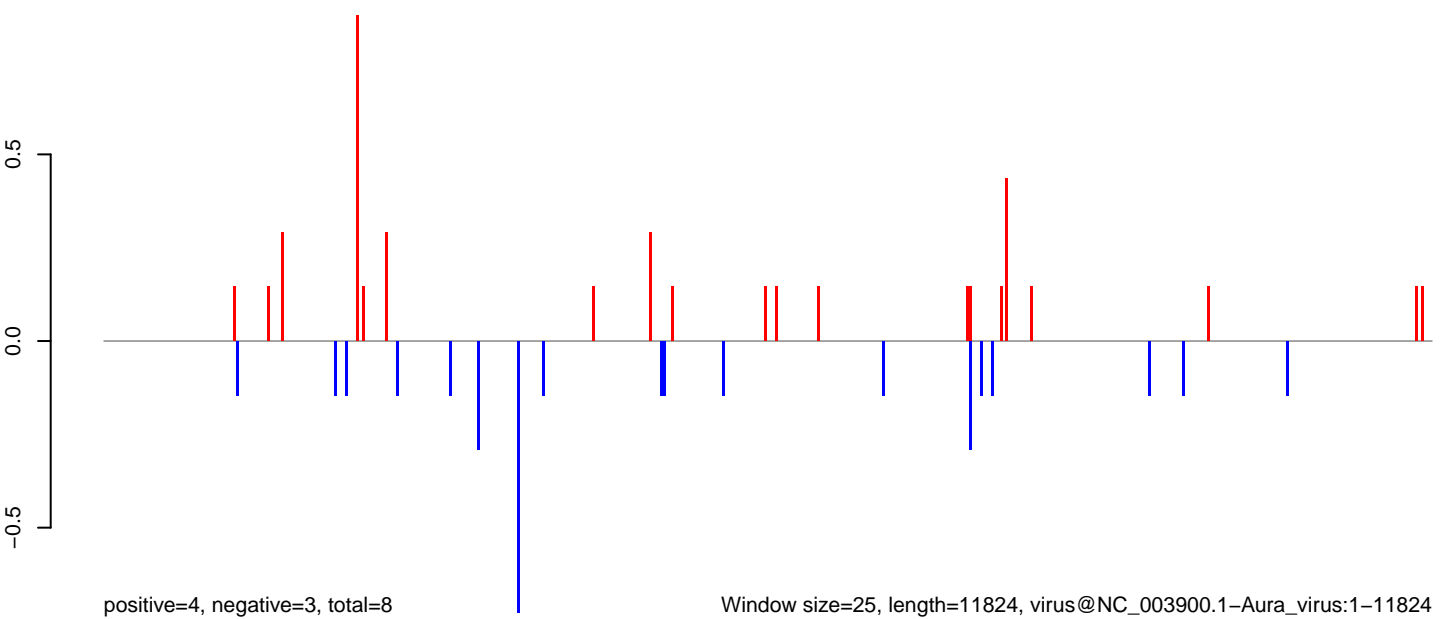
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



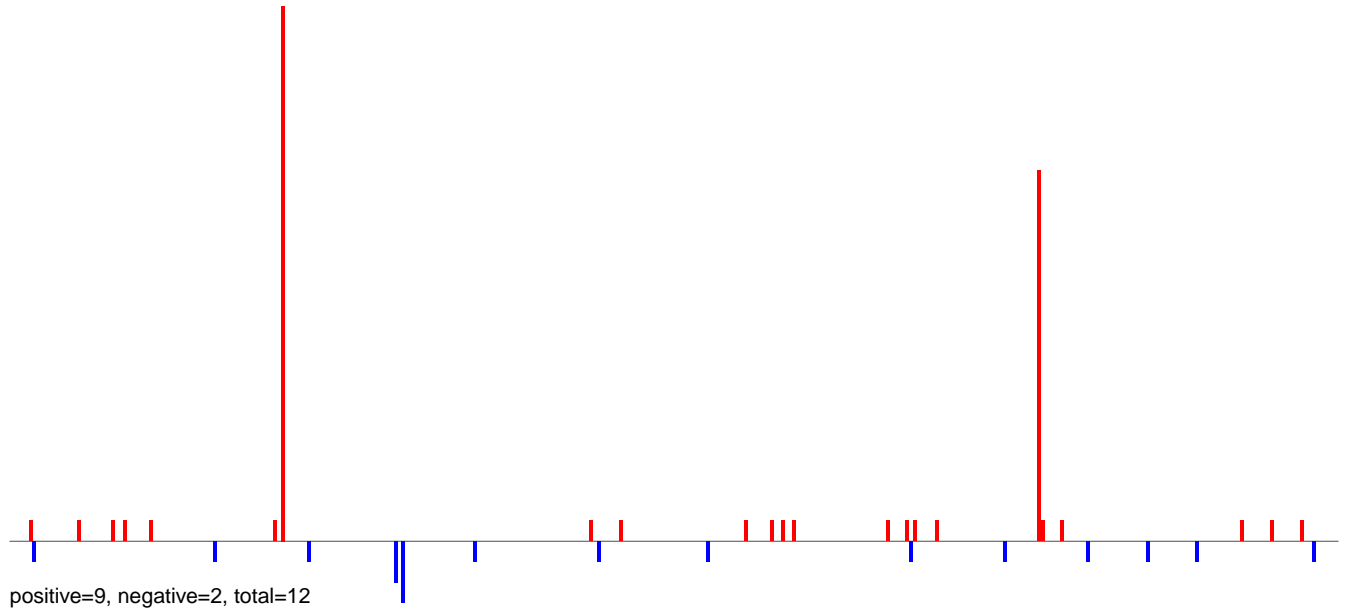
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



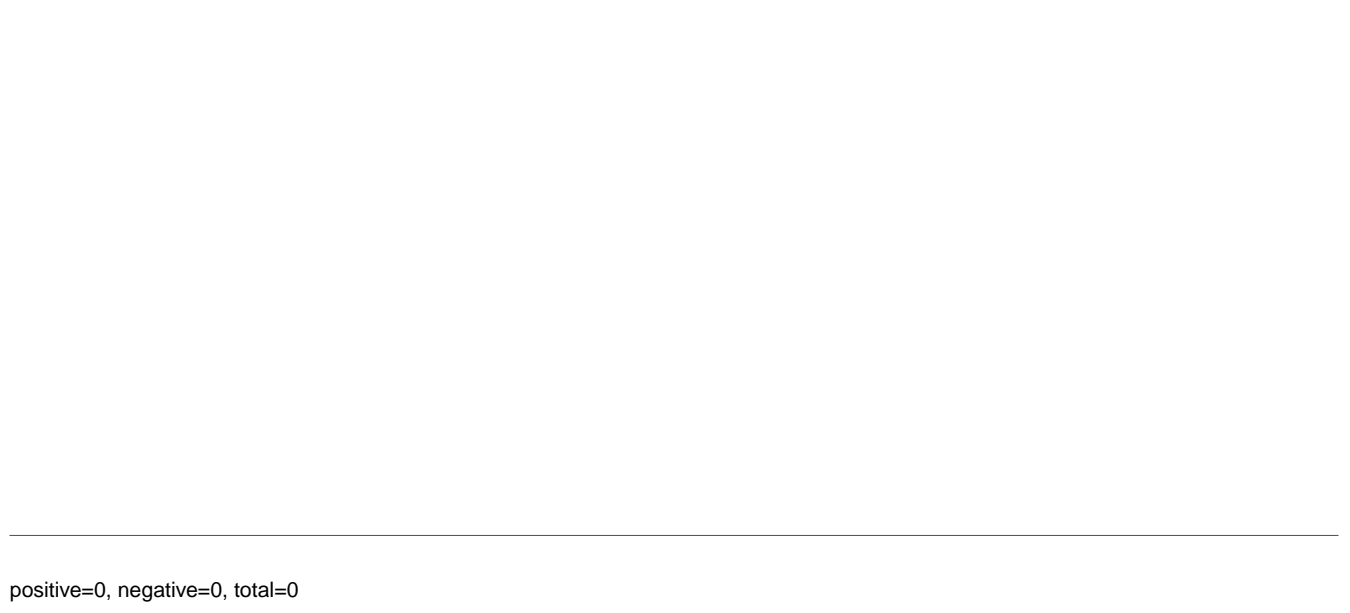
AeAeg_LVP_pMos3DBhspPBac.rep



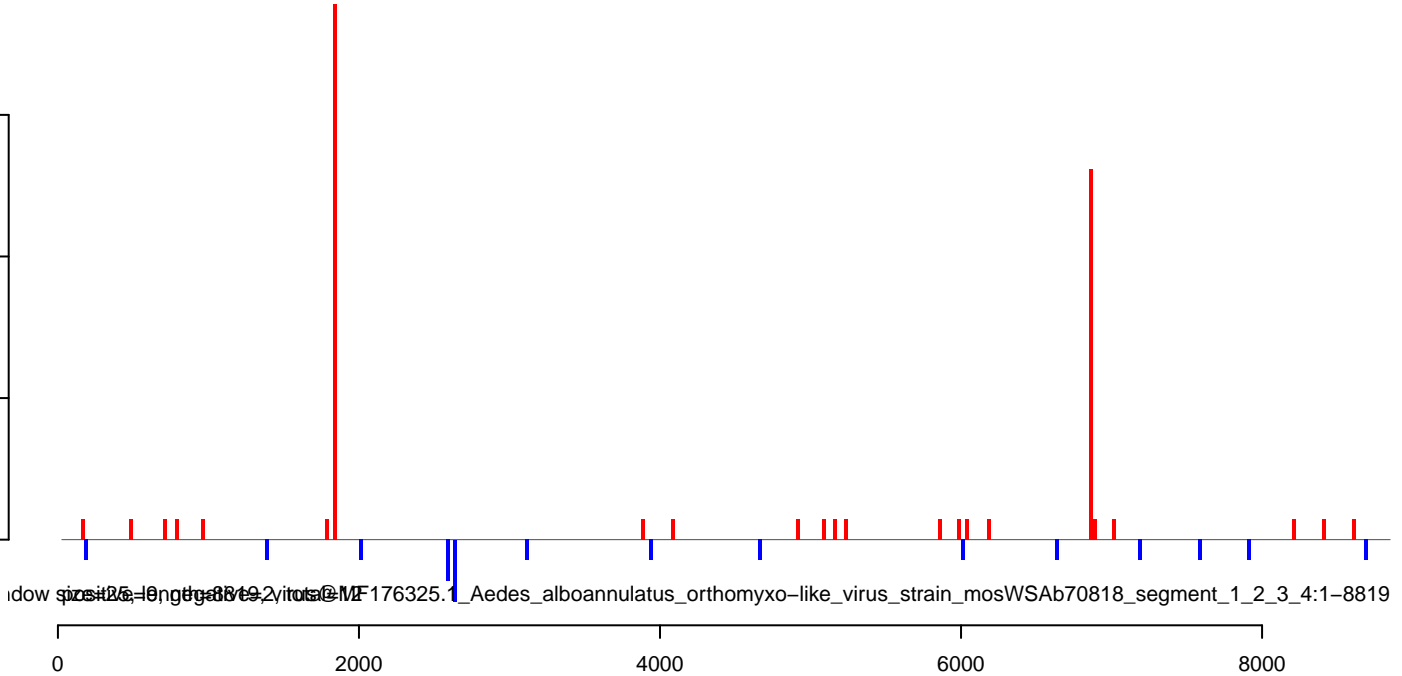
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



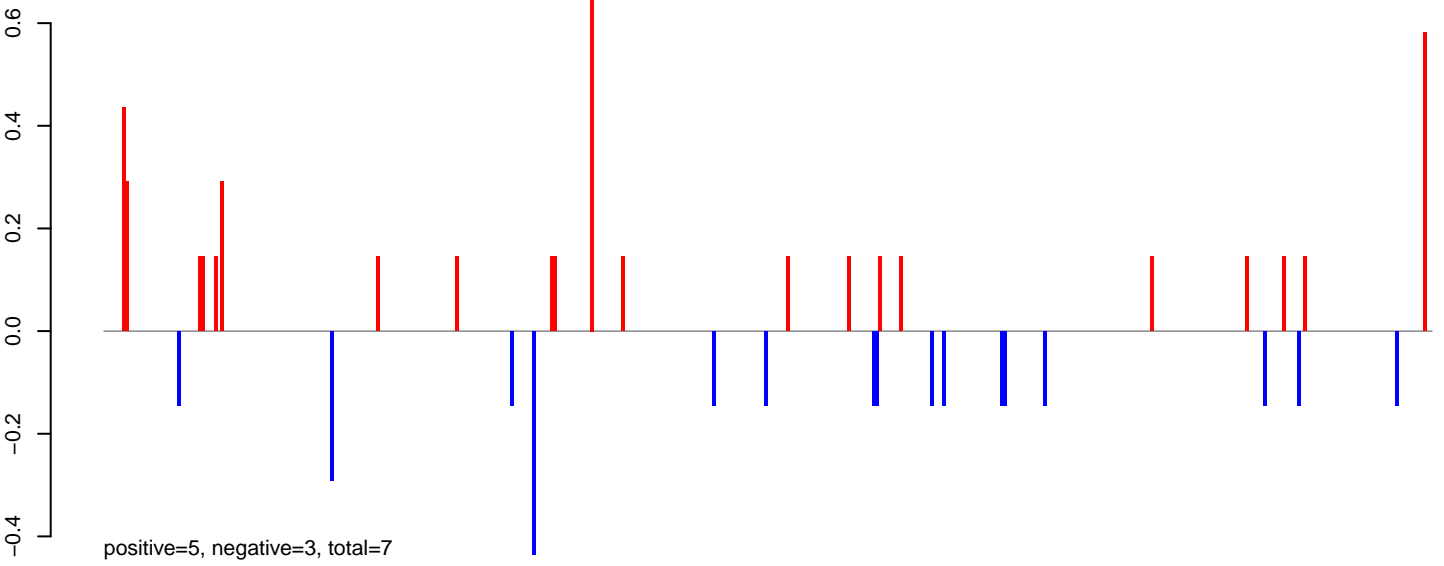
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



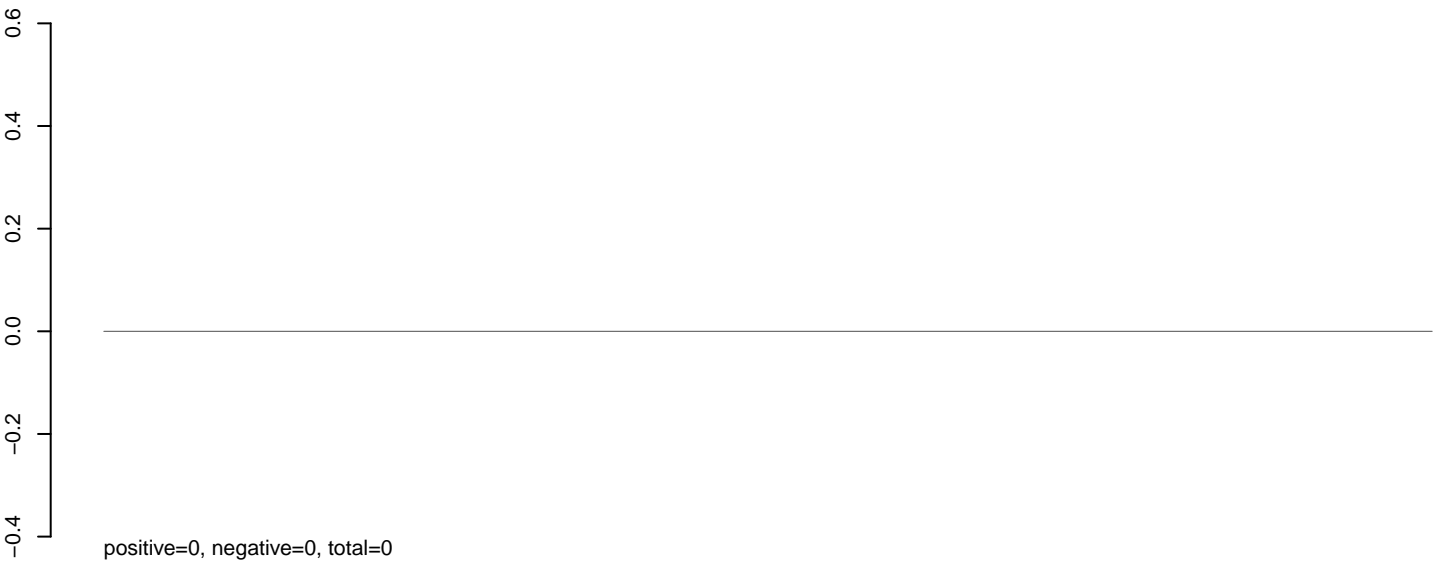
AeAeg_LVP_pMos3DBhspPBac.rep



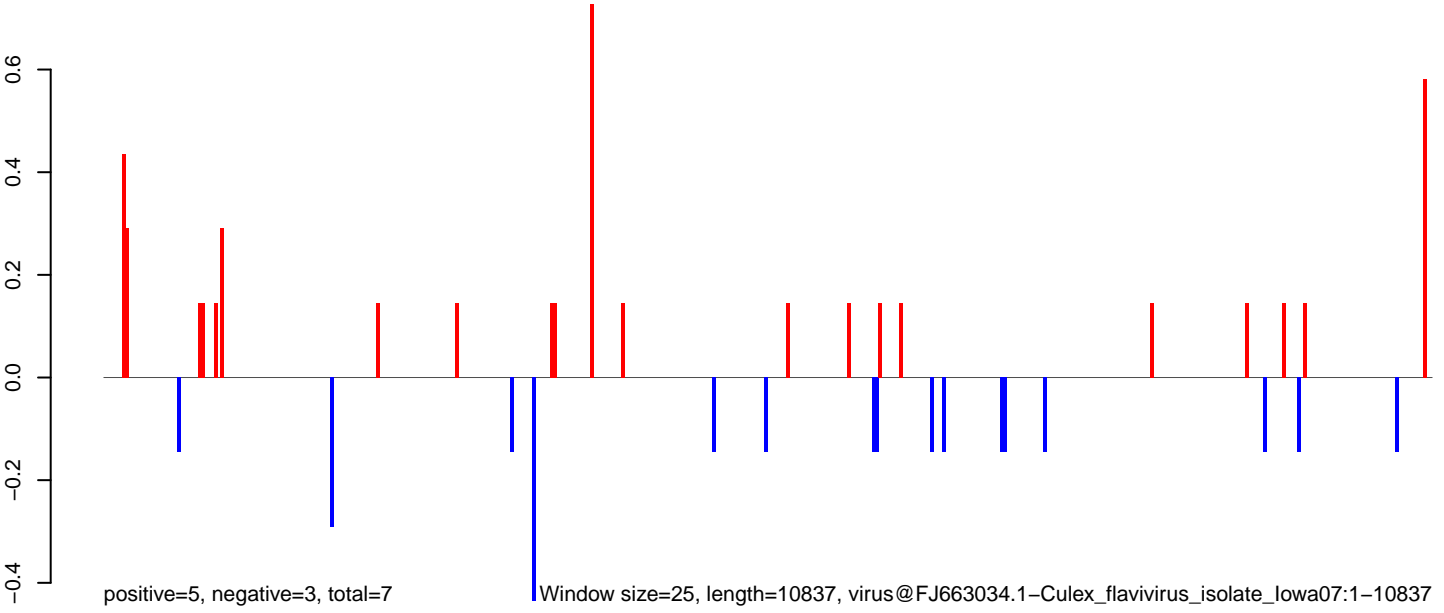
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

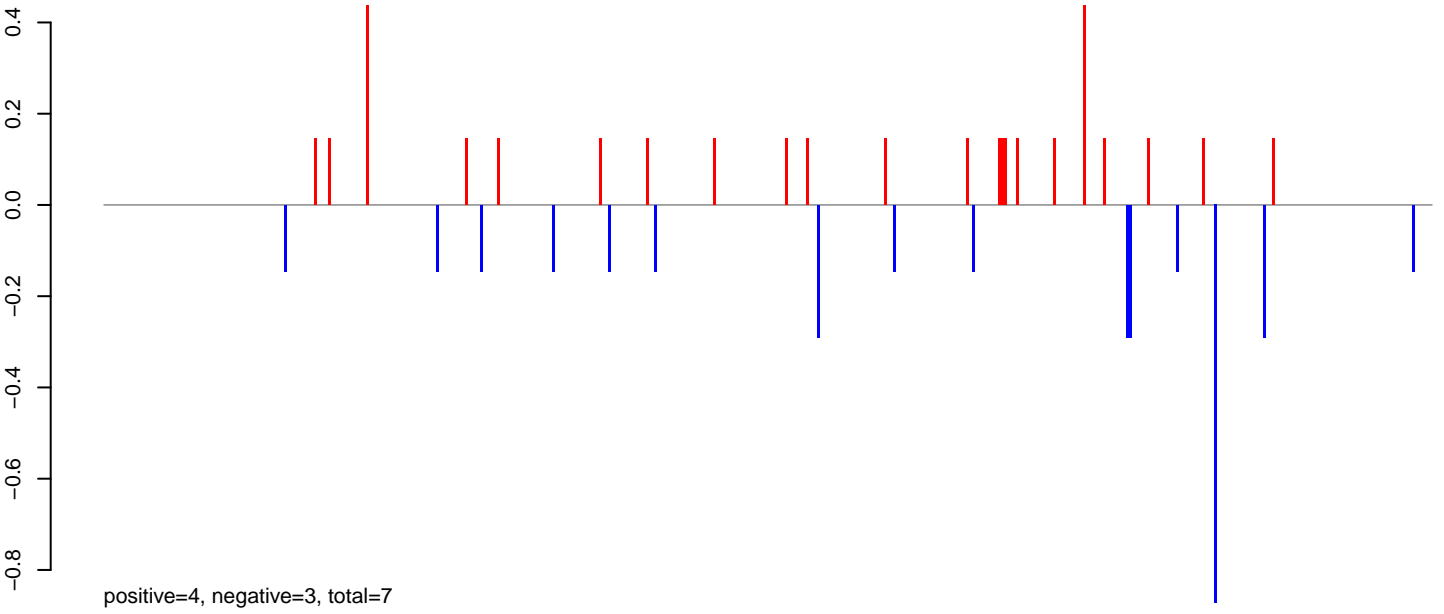


AeAeg_LVP_pMos3DBhspPBac.rep

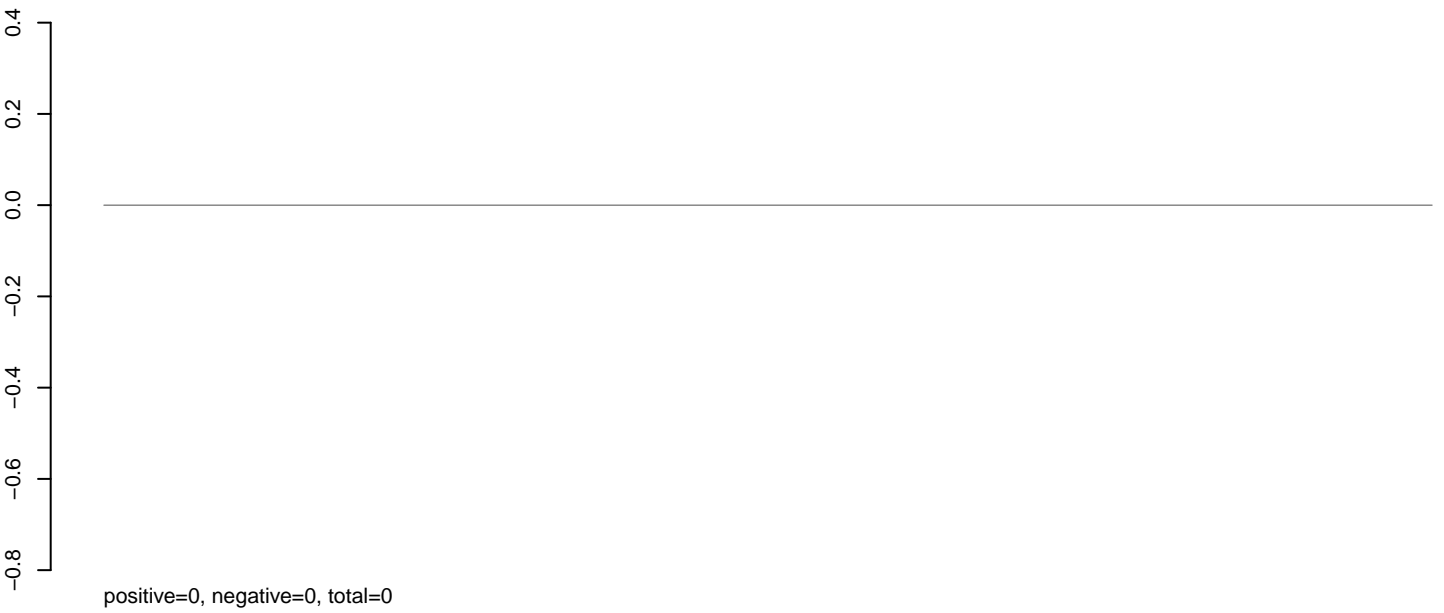


0 2000 4000 6000 8000 10000

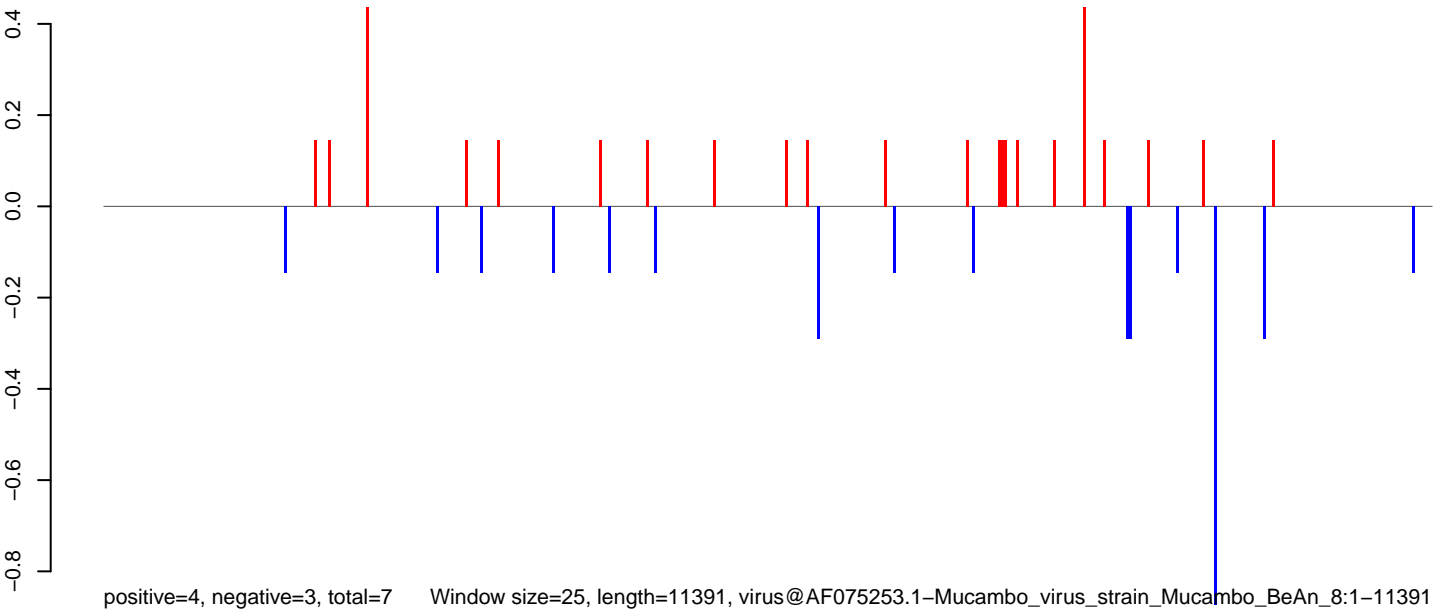
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



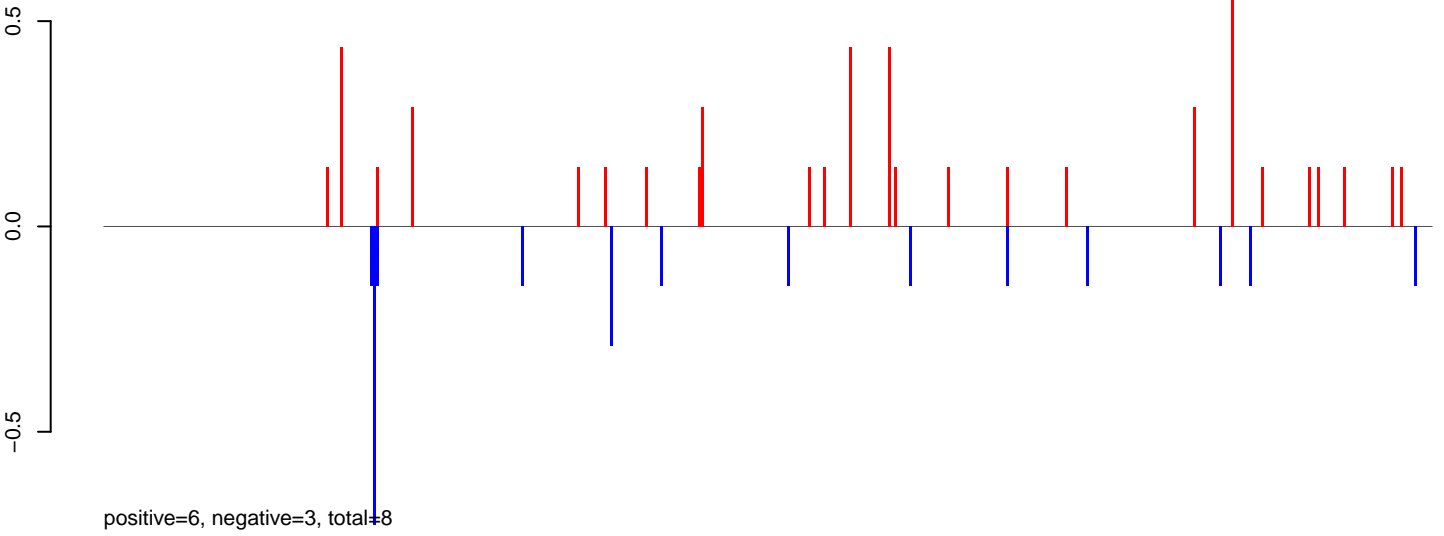
AeAeg_LVP_pMos3DBhspPBac.rep



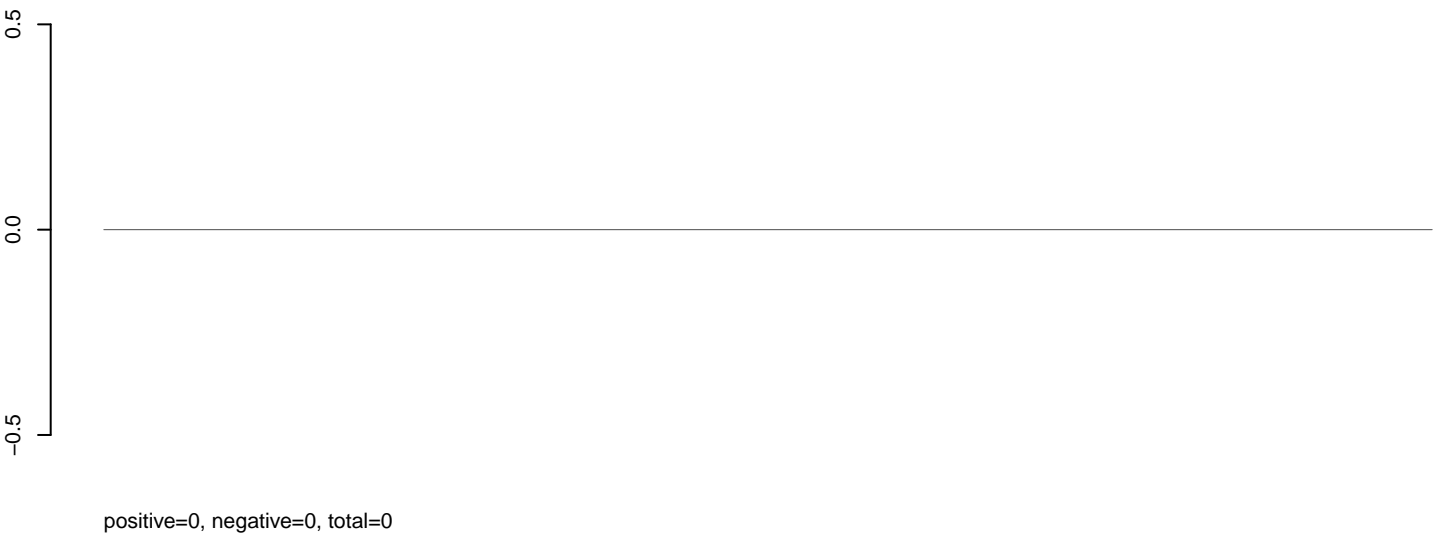
Window size=25, length=11391, virus@AF075253.1-Mucambo_virus_strain_Mucambo_BeAn_8:1-11391

0 2000 4000 6000 8000 10000

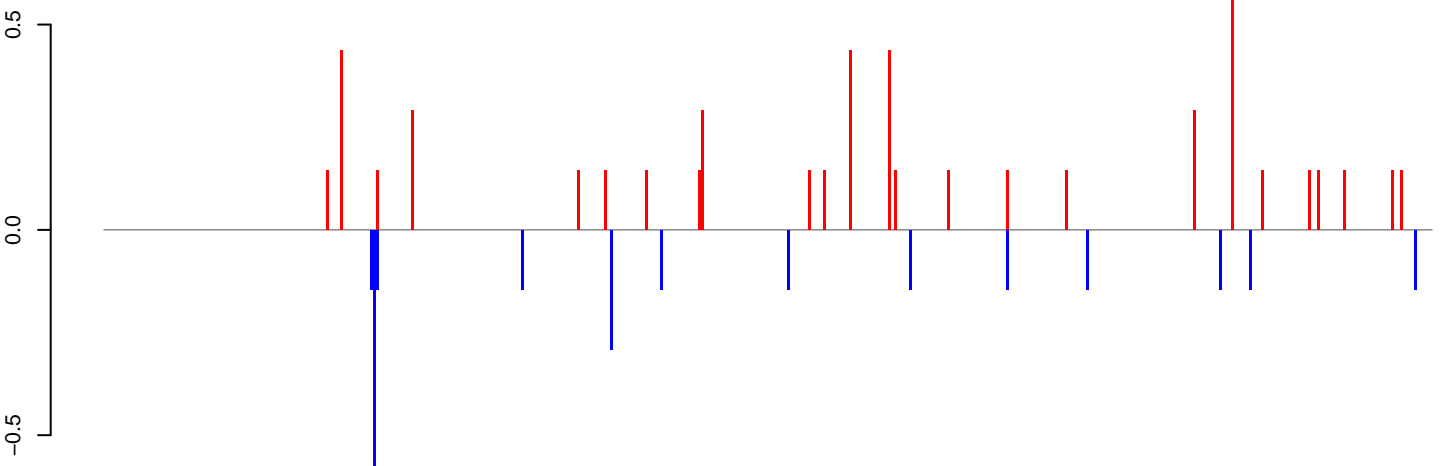
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



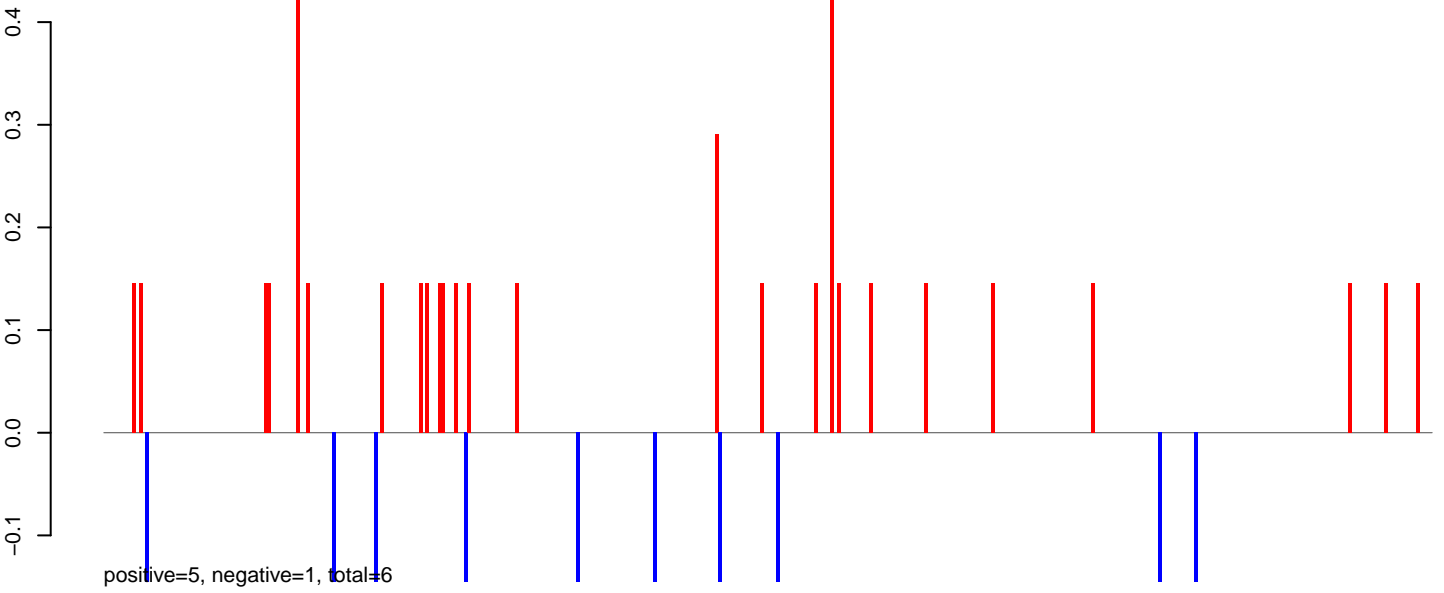
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



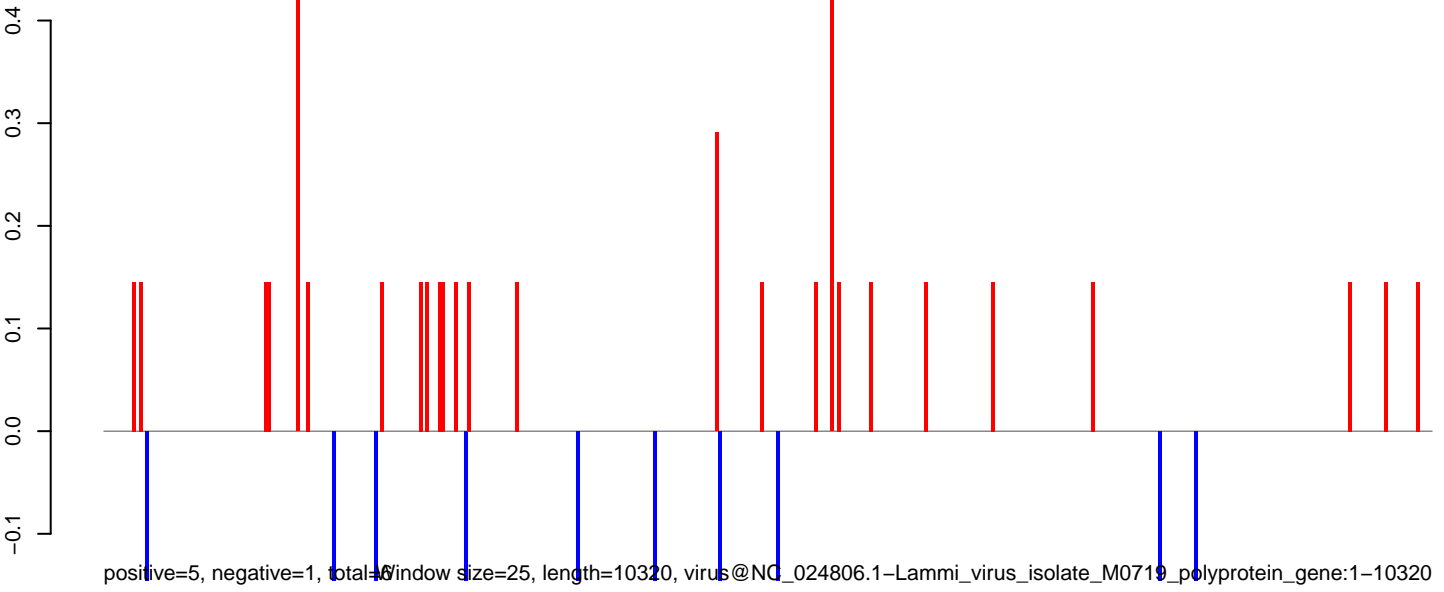
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



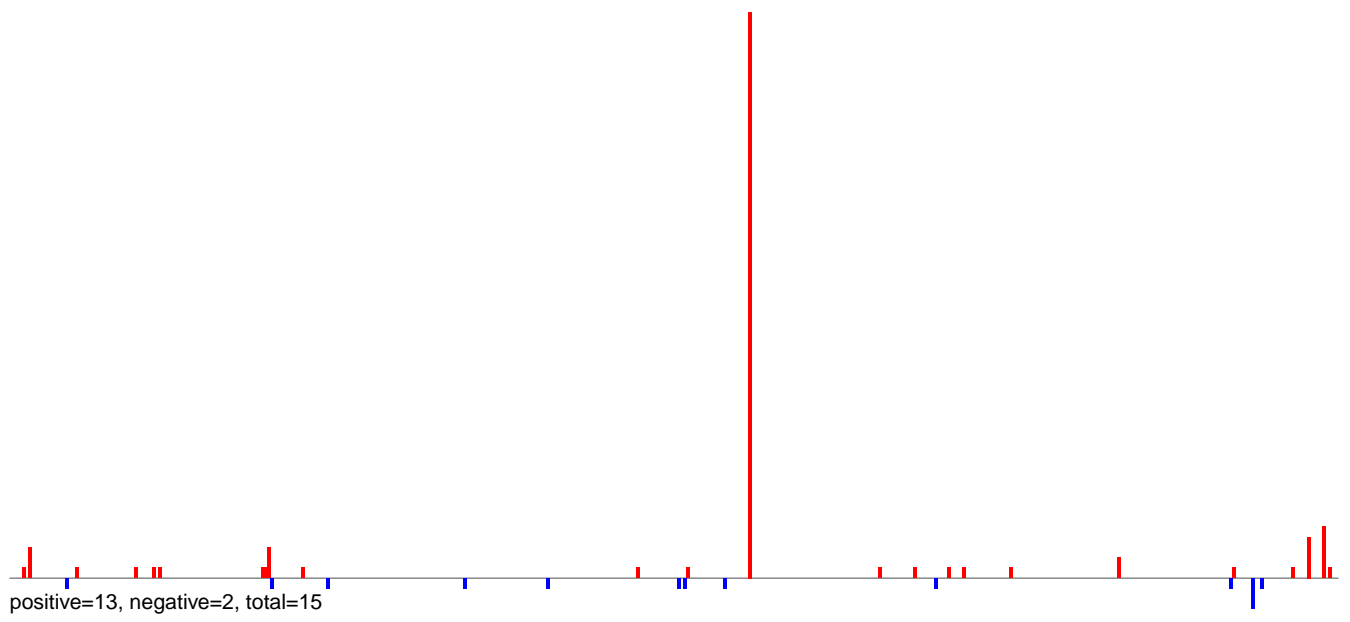
AeAeg_LVP_pMos3DBhspPBac.rep



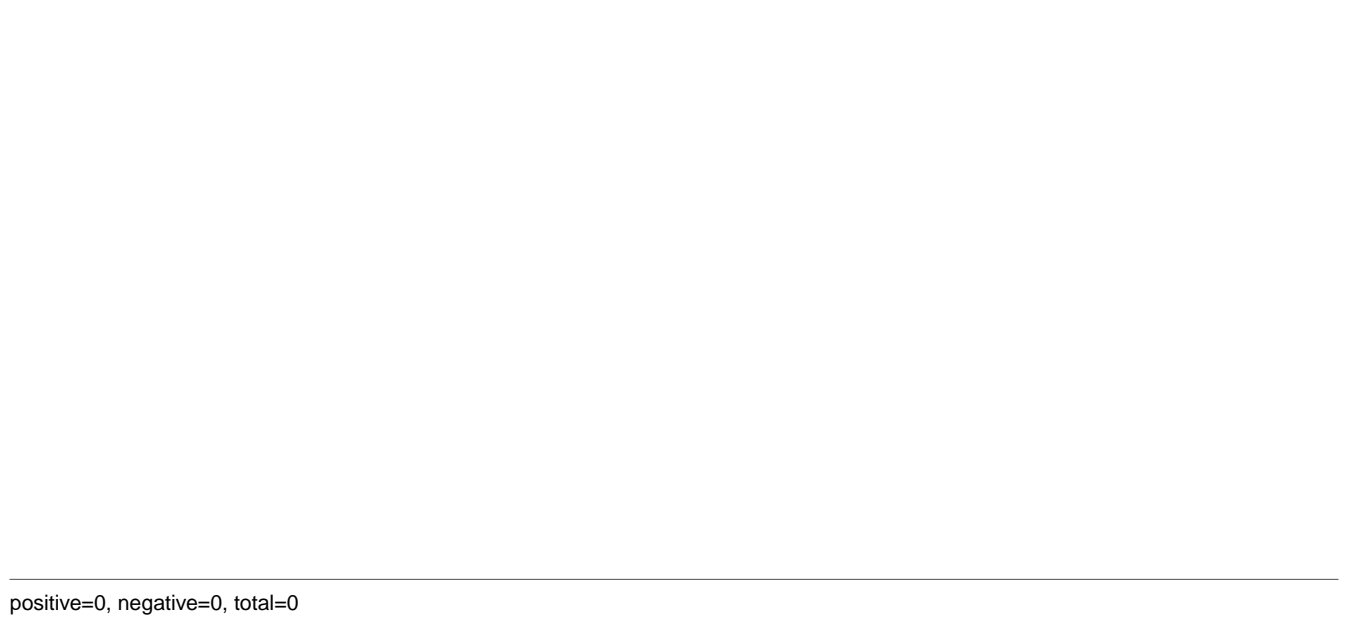
Window size=25, length=10320, virus@NC_024806.1-Lammi_virus_isolate_M0719_polyprotein_gene:1-10320

0 2000 4000 6000 8000 10000

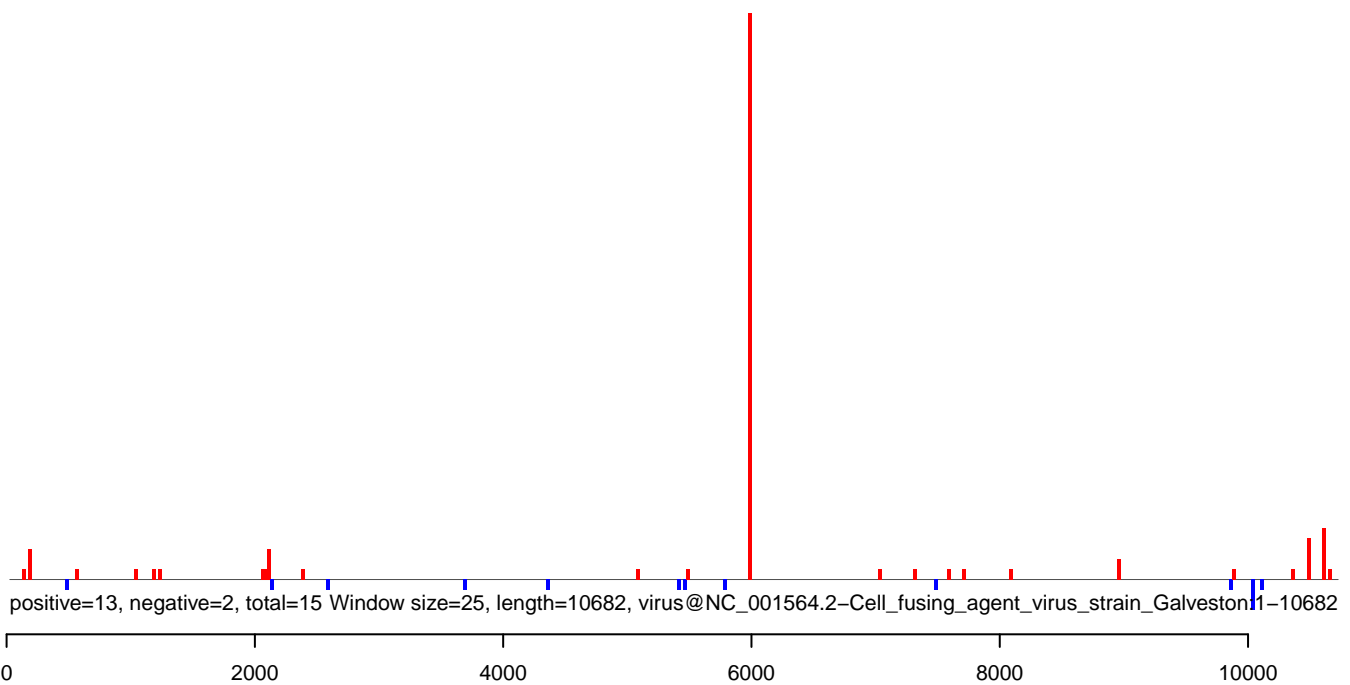
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



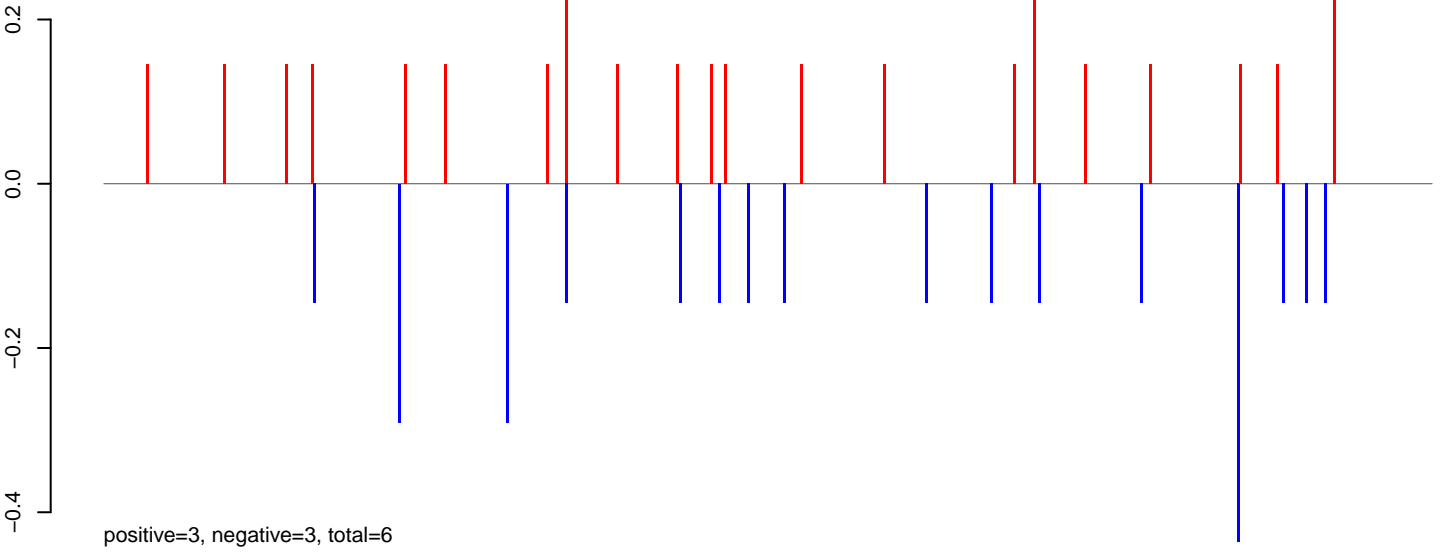
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



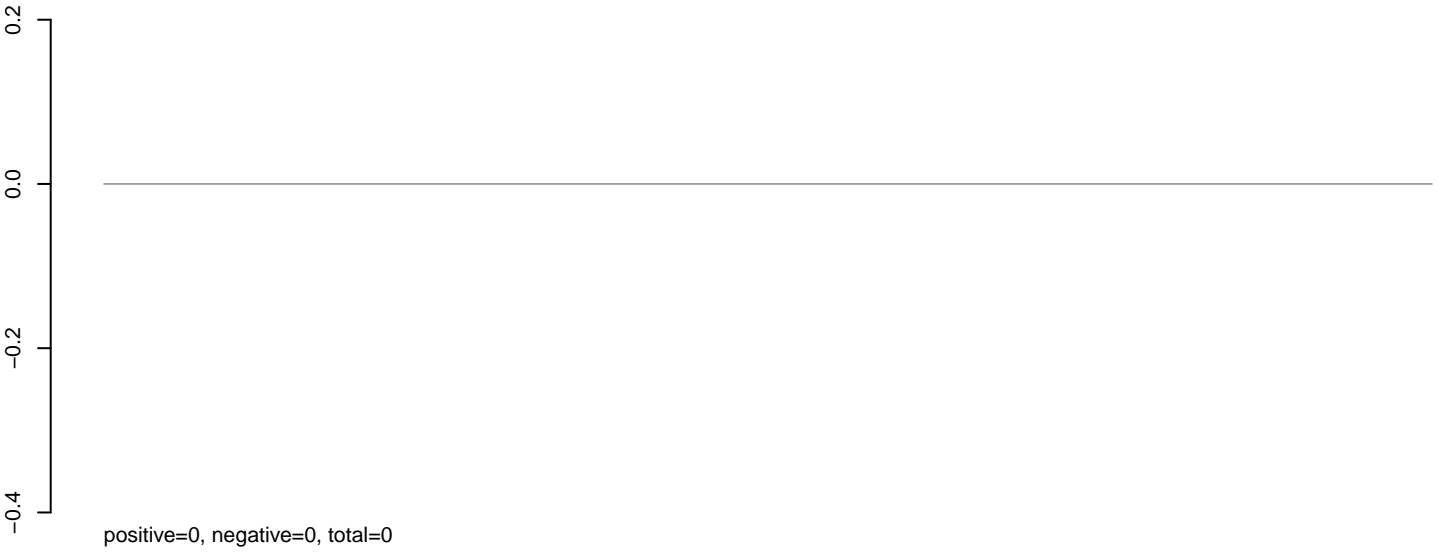
AeAeg_LVP_pMos3DBhspPBac.rep



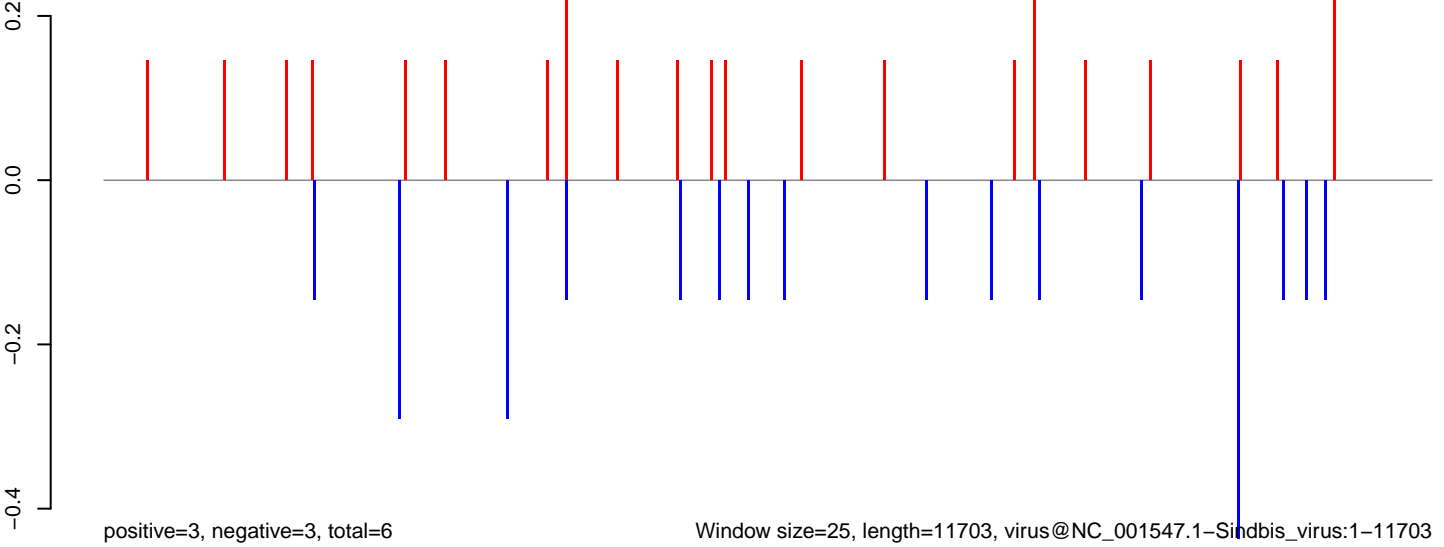
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



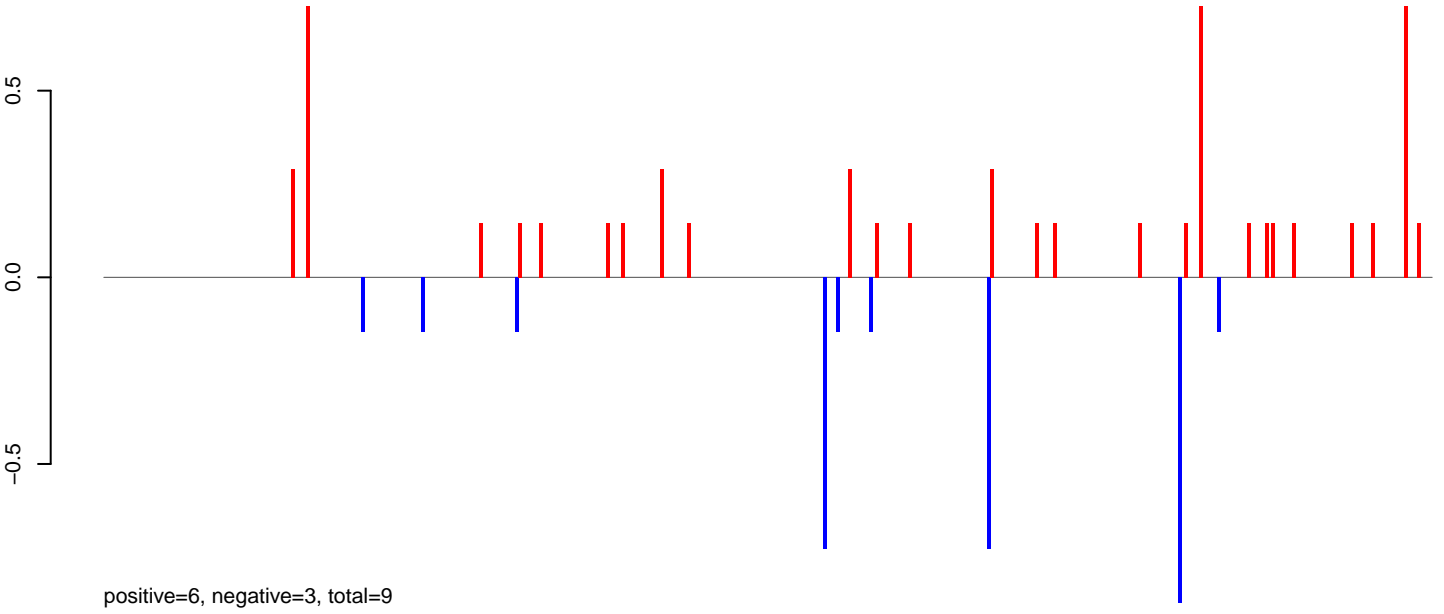
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



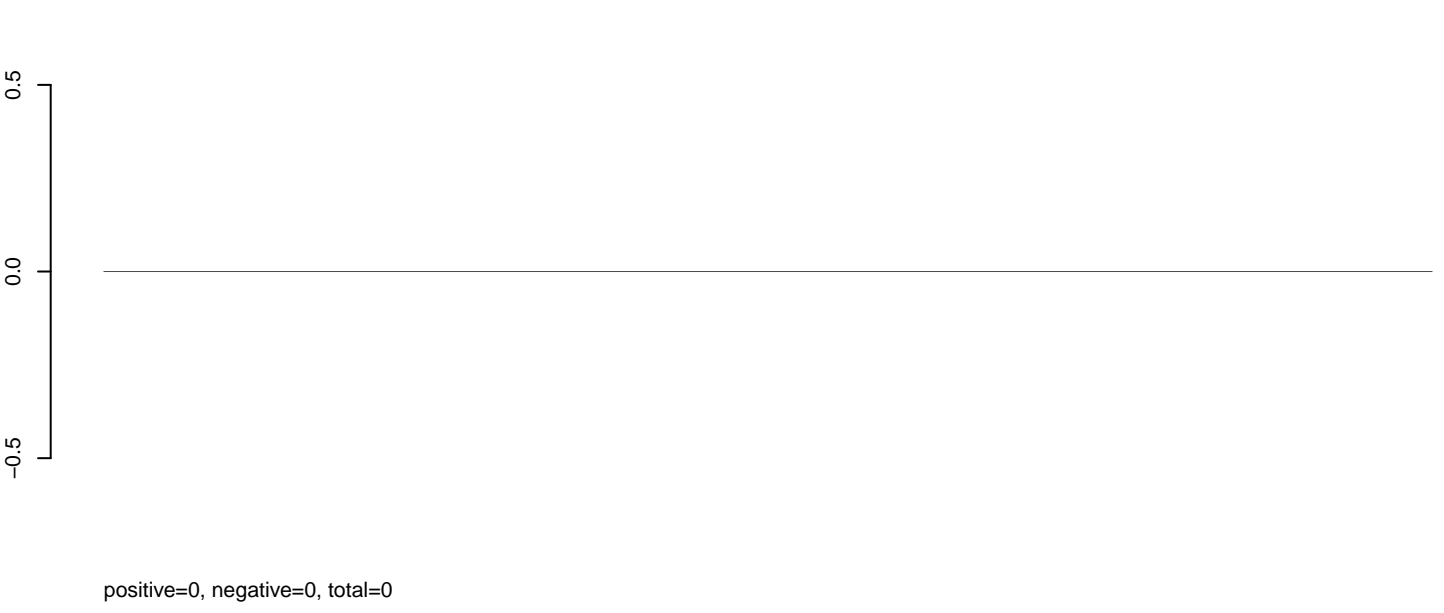
AeAeg_LVP_pMos3DBhspPBac.rep



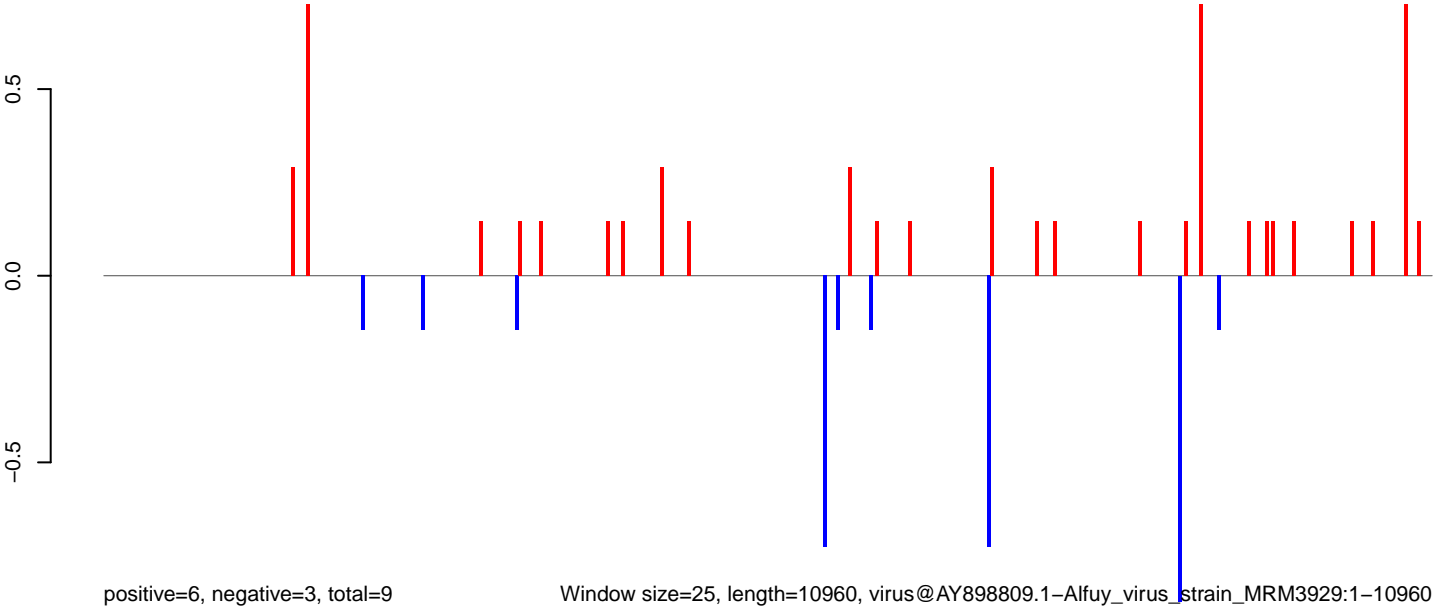
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



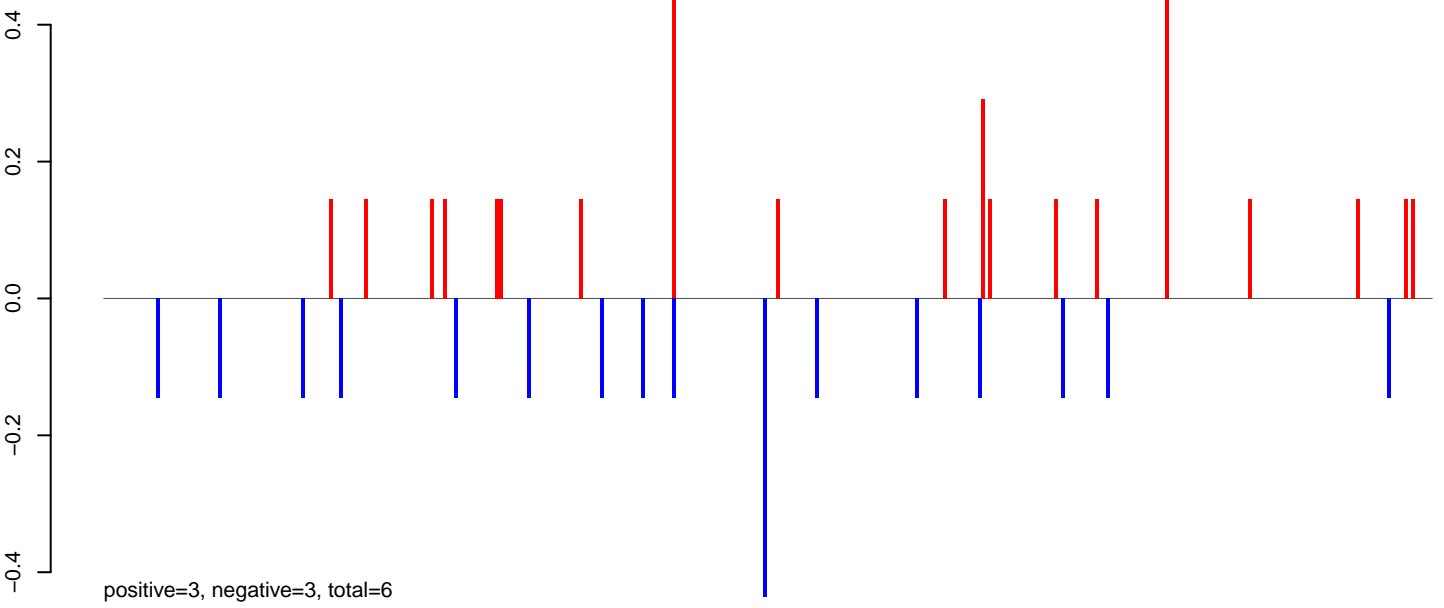
AeAeg_LVP_pMos3DBhspPBac.rep



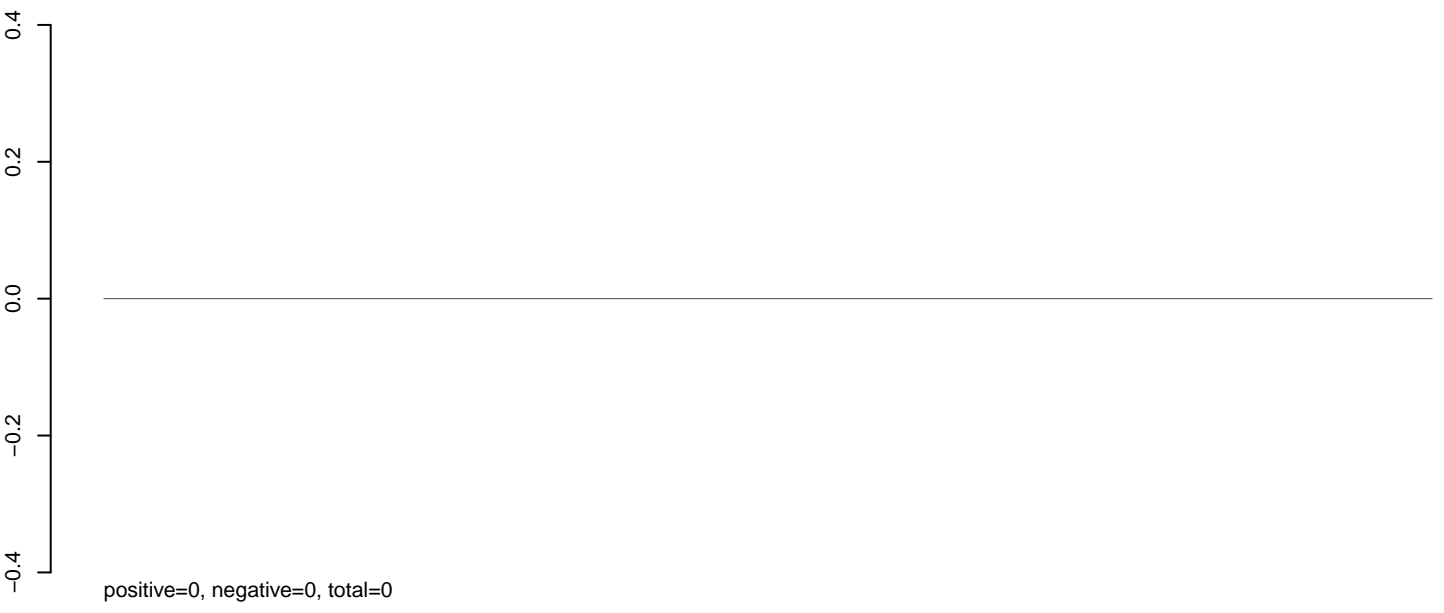
Window size=25, length=10960, virus@AY898809.1-Alfuy_virus_strain_MRM3929:1-10960

0 2000 4000 6000 8000 10000

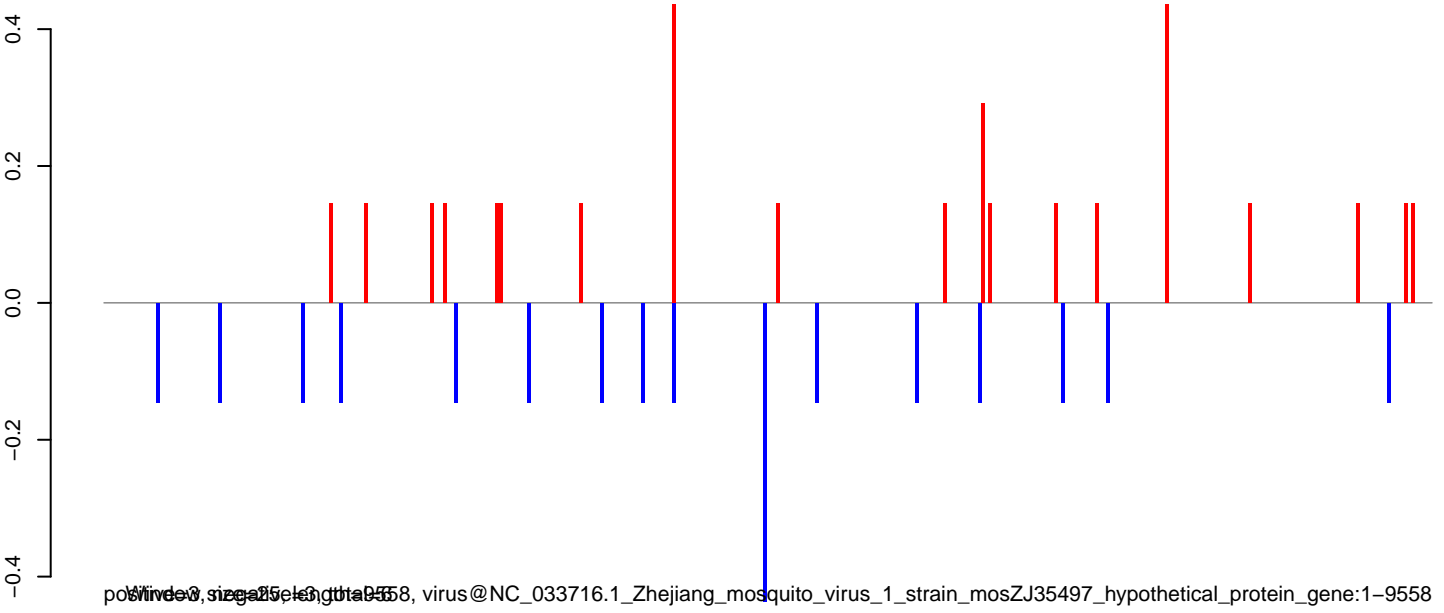
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

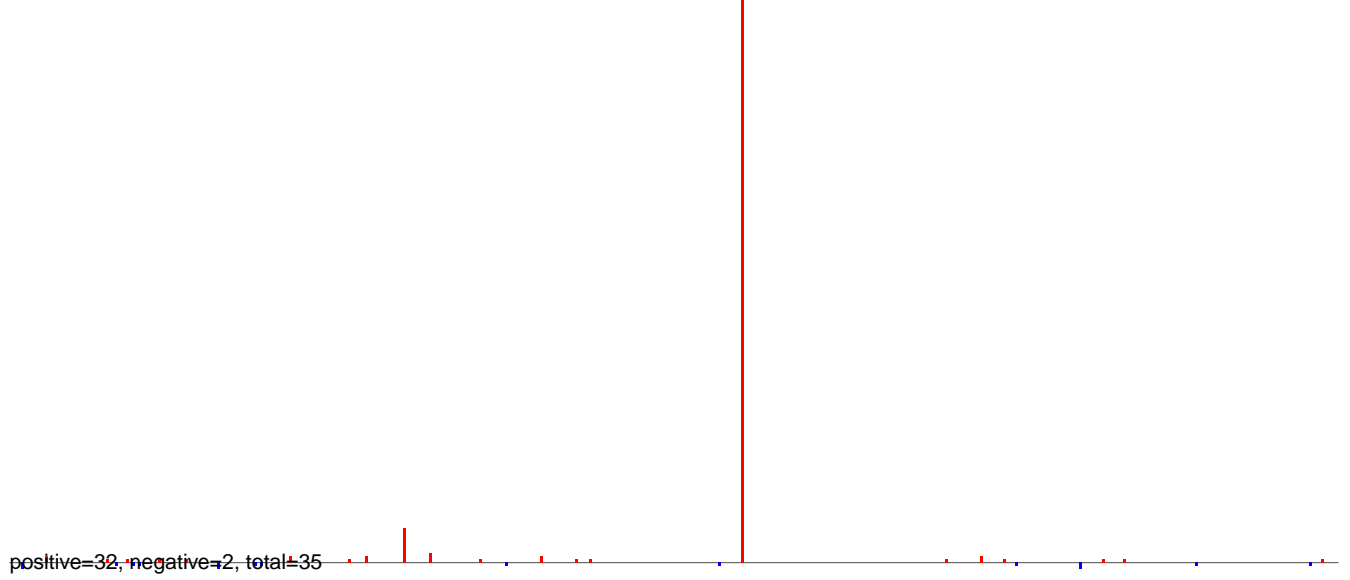


AeAeg_LVP_pMos3DBhspPBac.rep

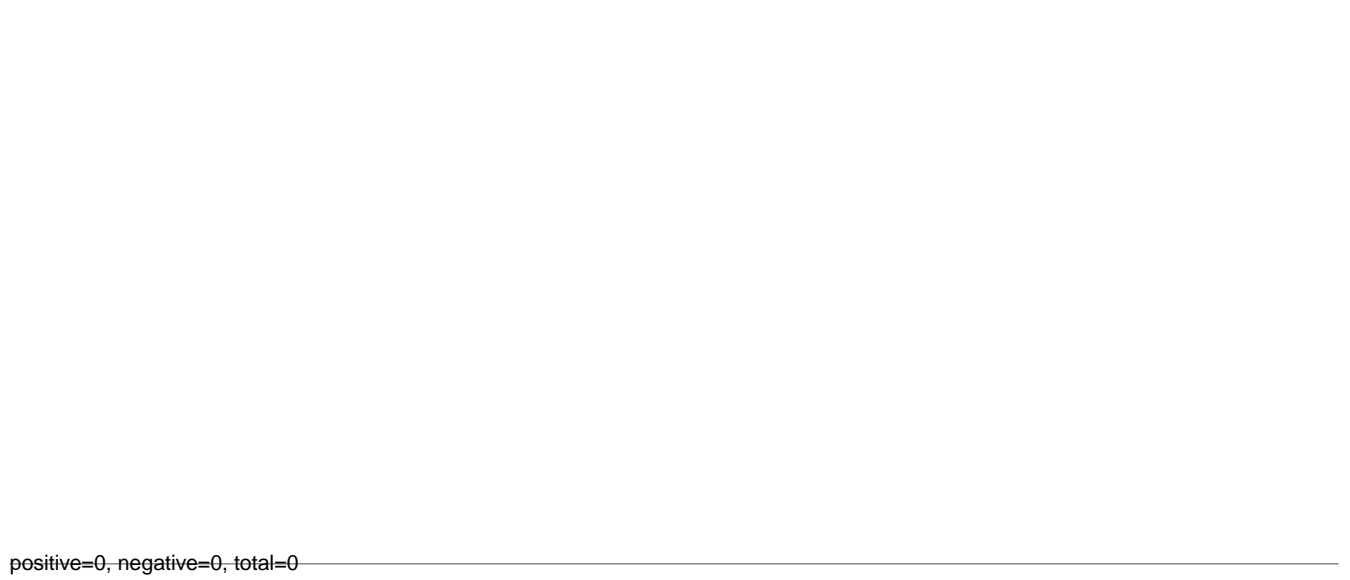


Window size=25, step=10, total=9558, virus@NC_033716.1_Zhejiang_mosquito_virus_1_strain_mosZJ35497_hypothetical_protein_gene:1-9558

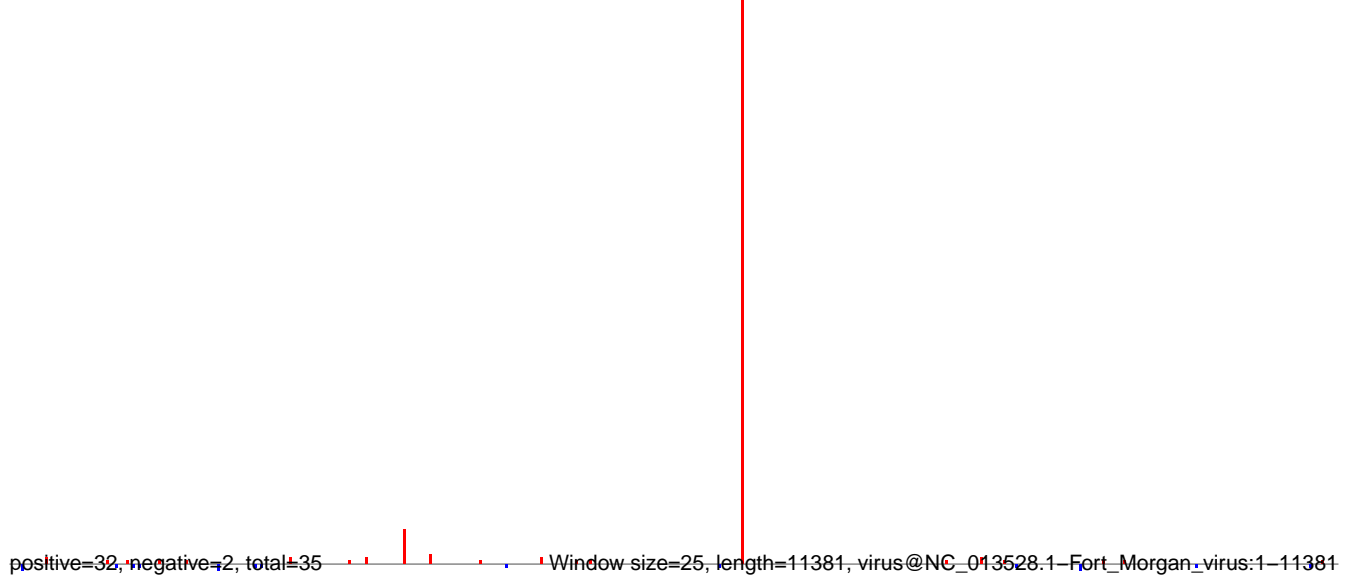
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



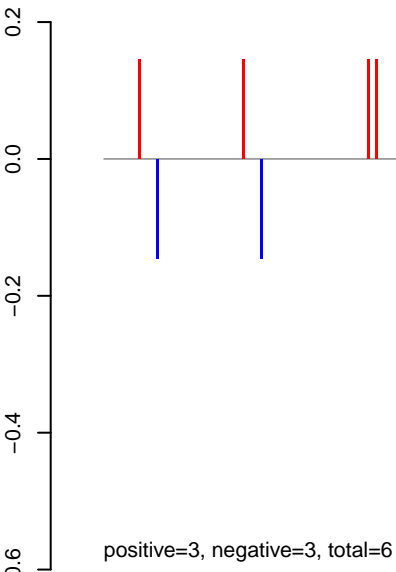
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



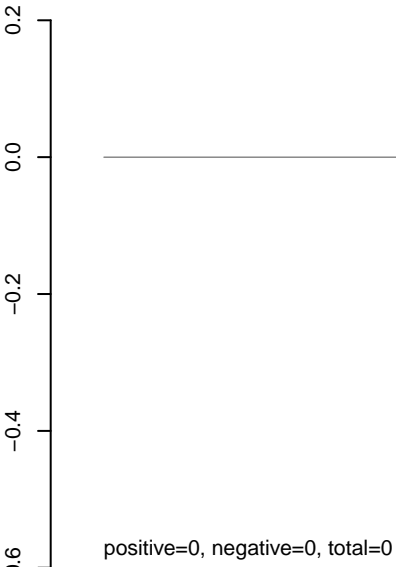
AeAeg_LVP_pMos3DBhspPBac.rep



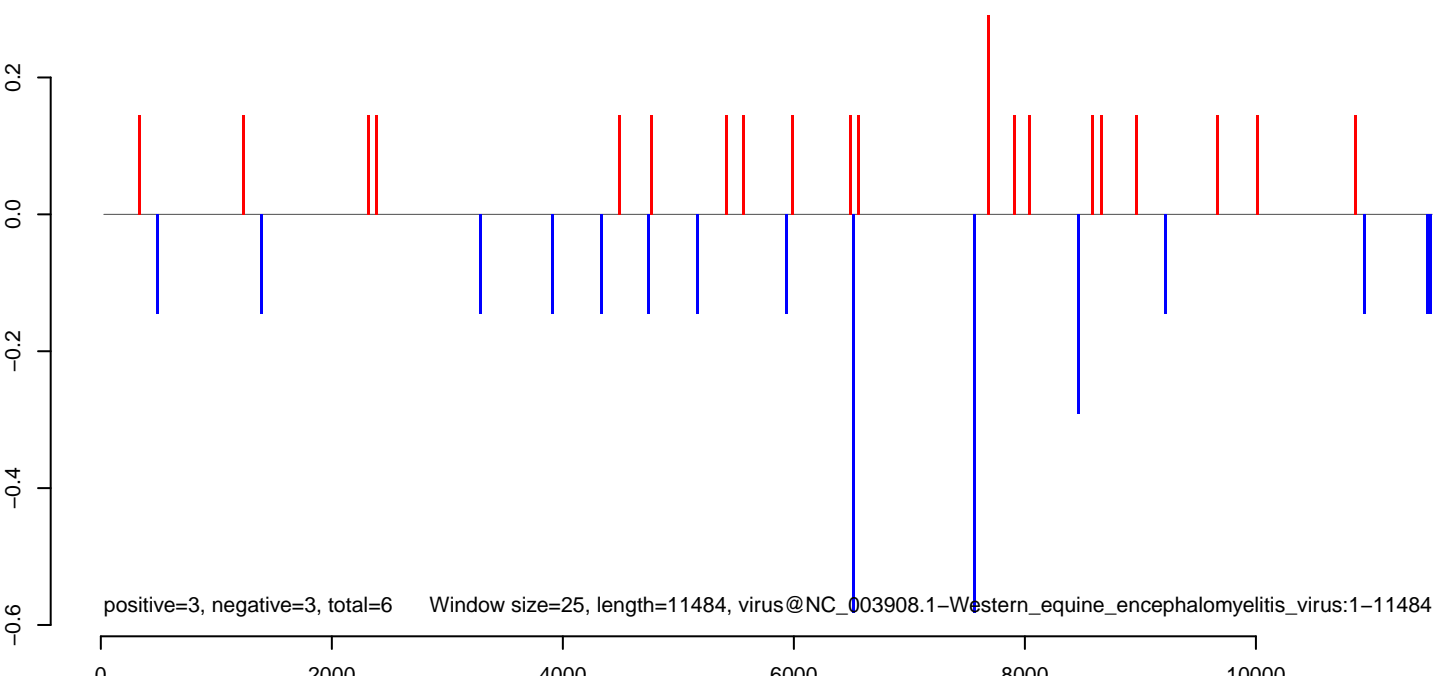
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



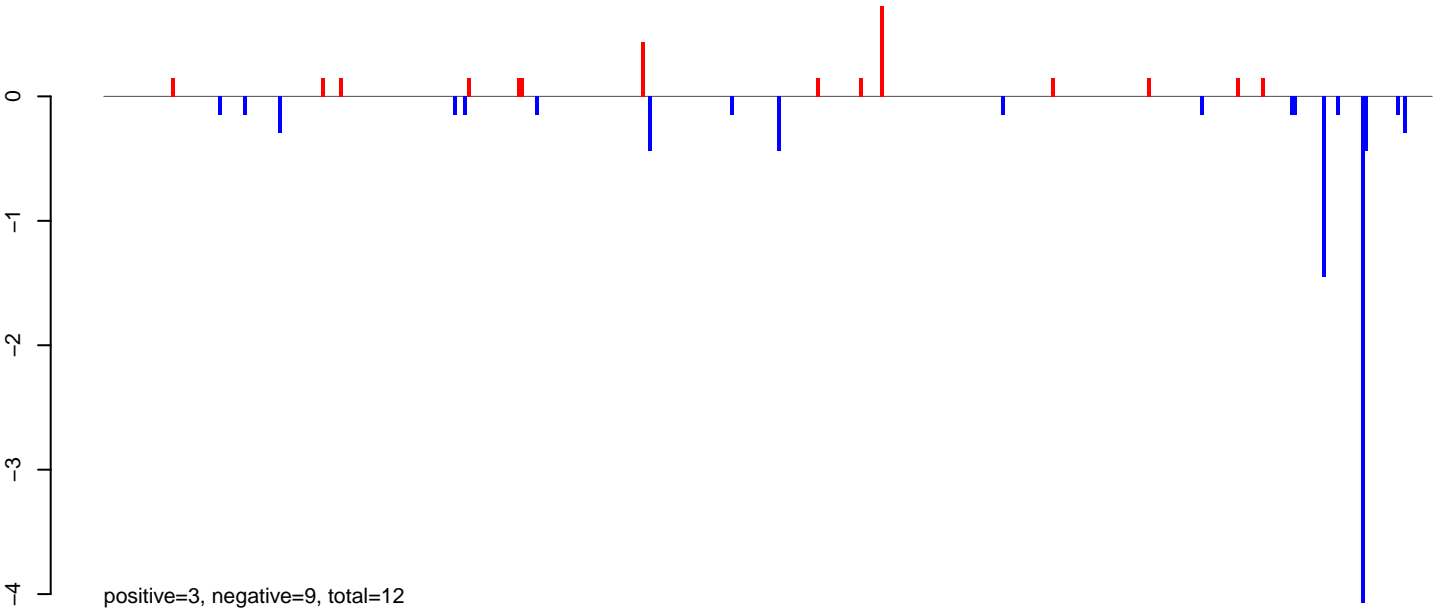
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



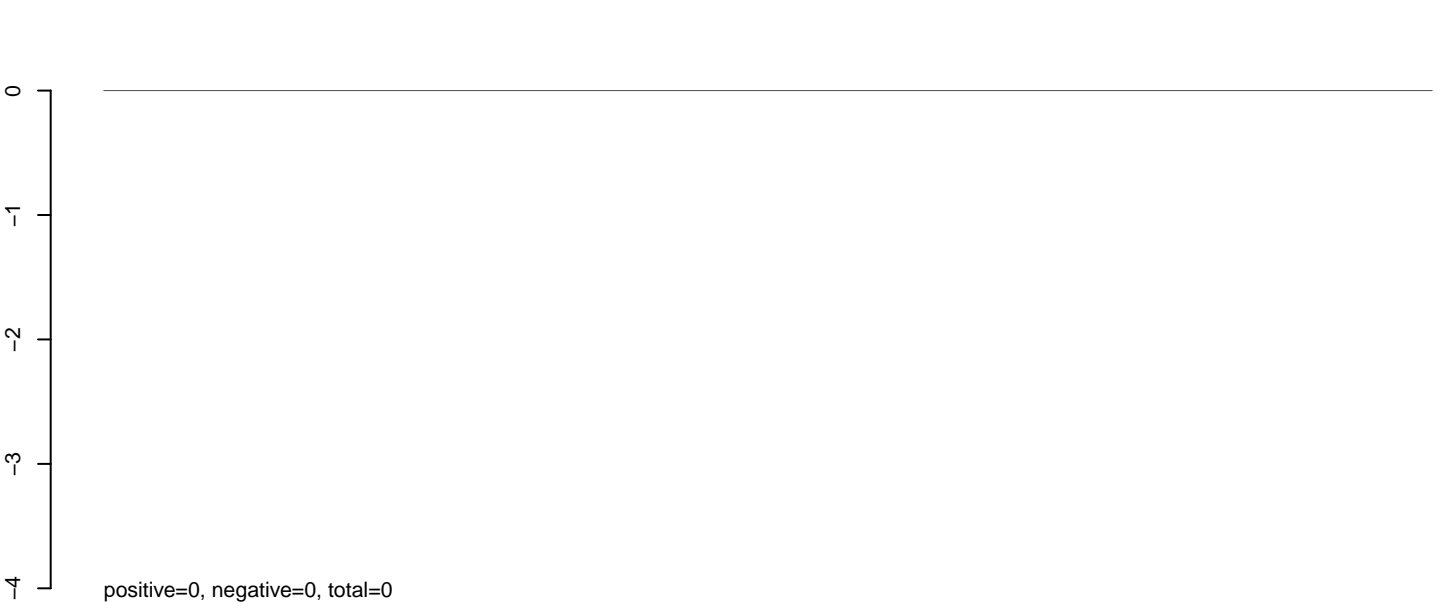
AeAeg_LVP_pMos3DBhspPBac.rep



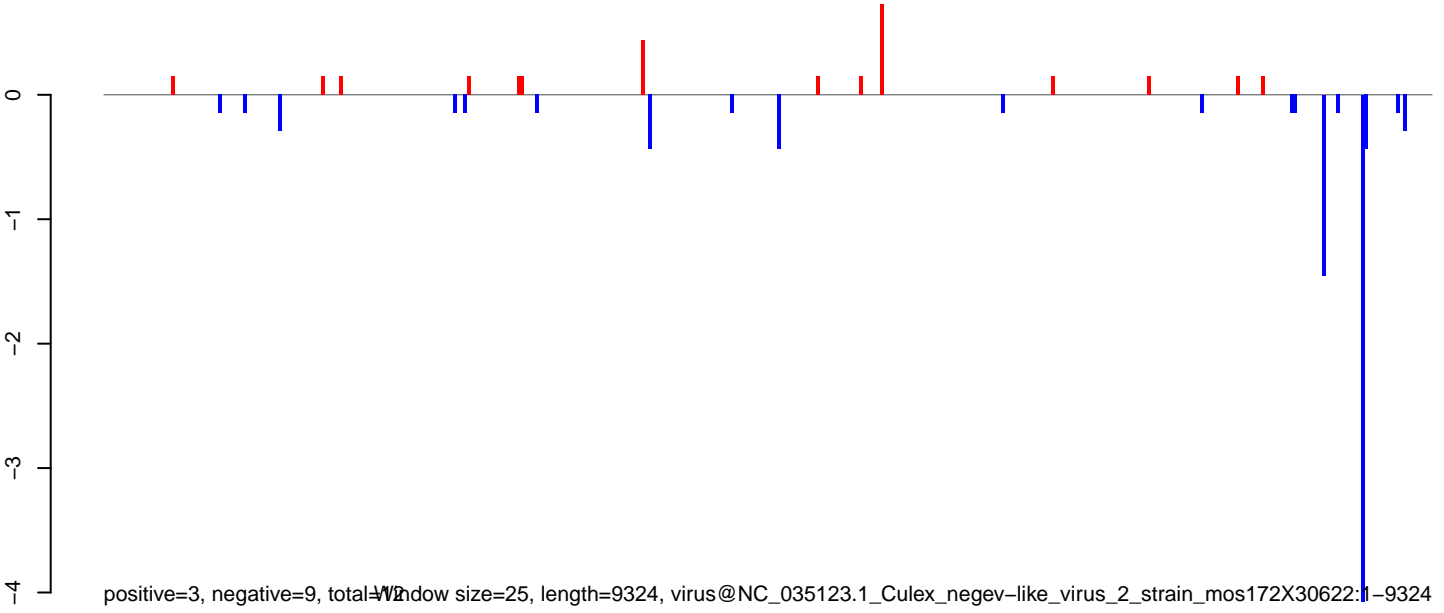
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



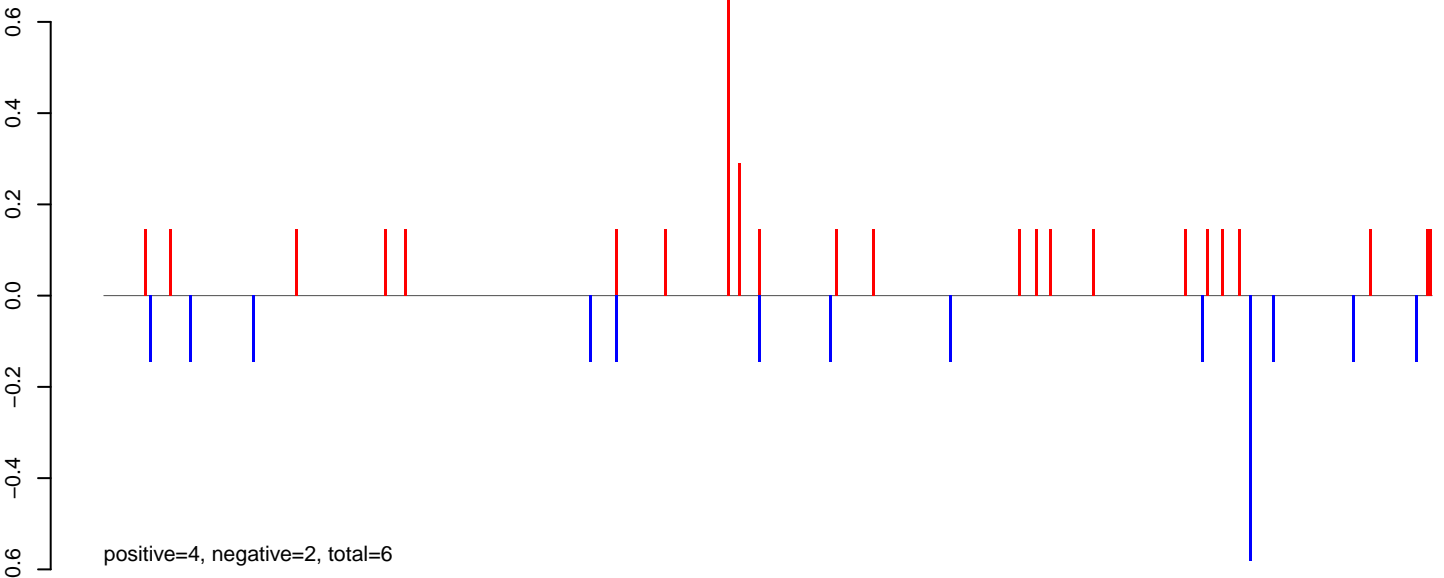
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



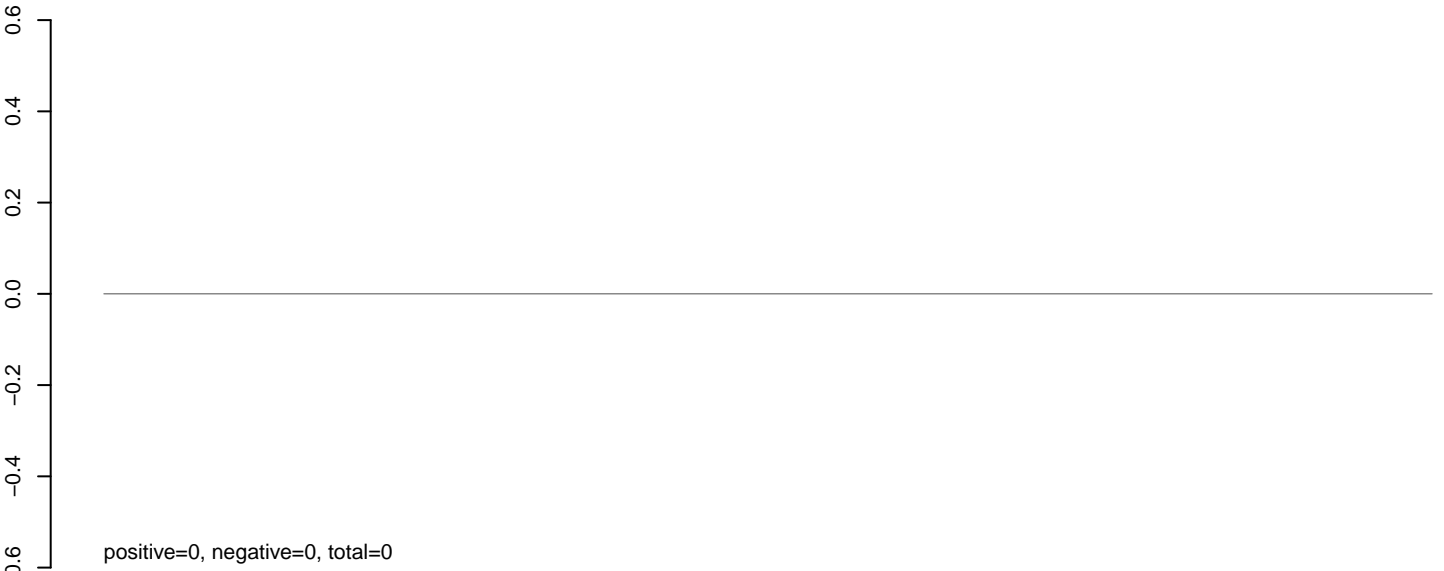
AeAeg_LVP_pMos3DBhspPBac.rep



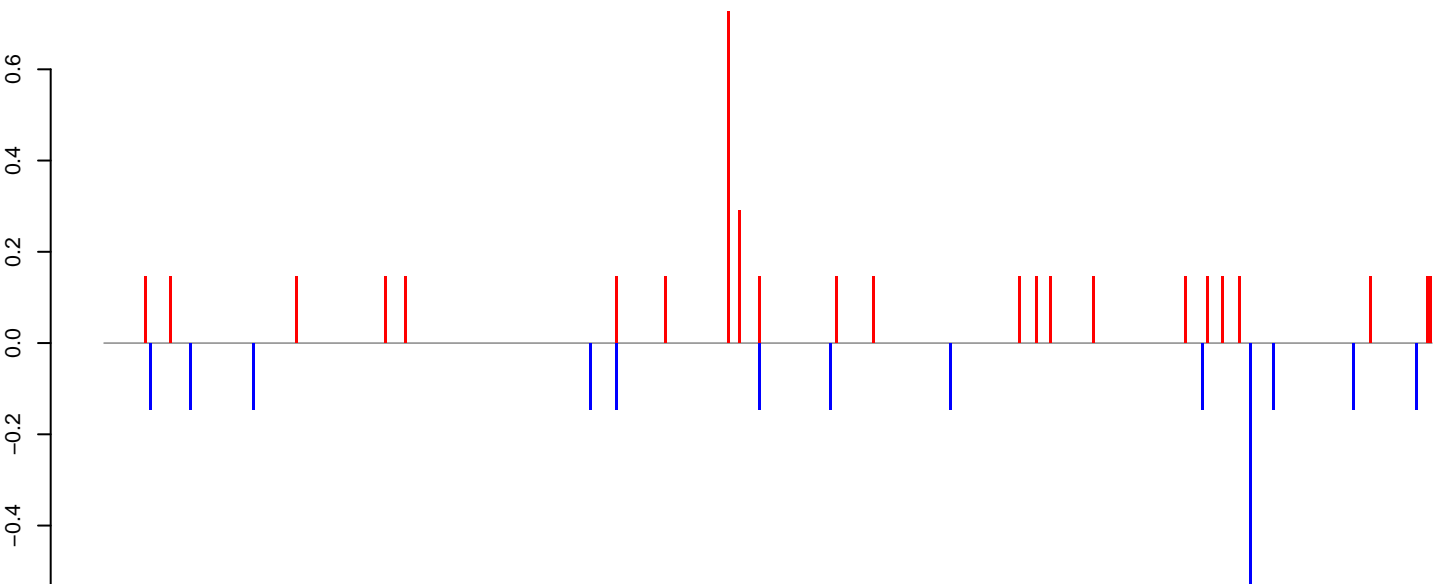
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



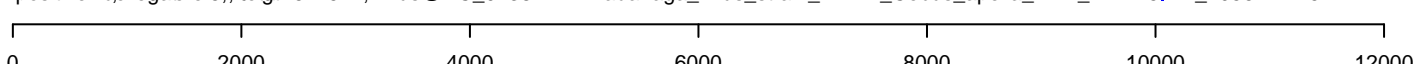
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



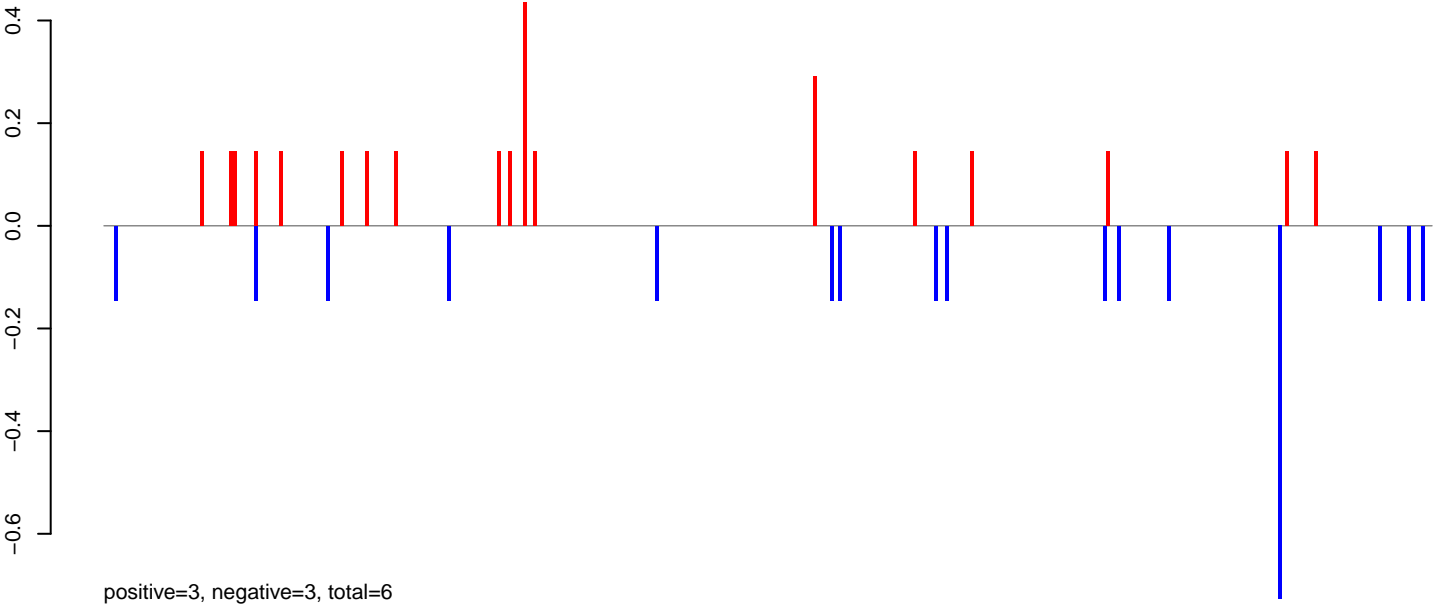
AeAeg_LVP_pMos3DBhspPBac.rep



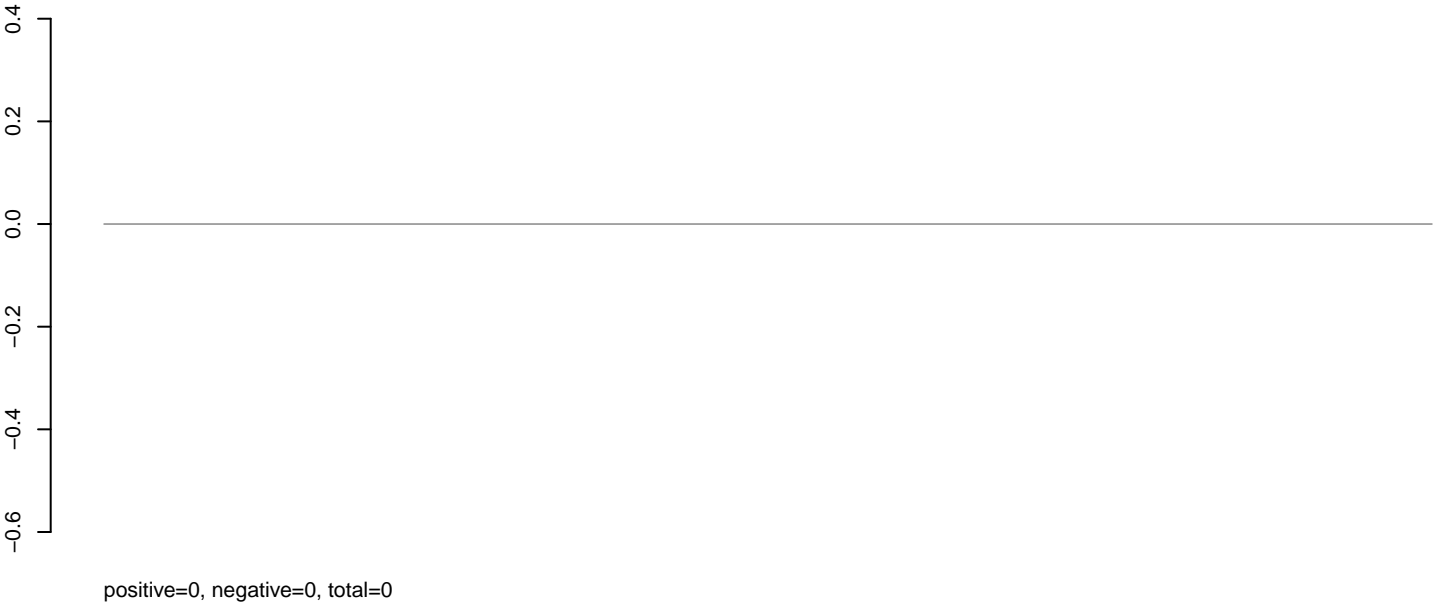
Viewed on <https://www.ncbi.nlm.nih.gov/trace/viz/11624>, virus@NC_023812.1-Madariaga_virus_strain_MADV_Cebus_apella_BRA_BEAN5122_1956:1-11624



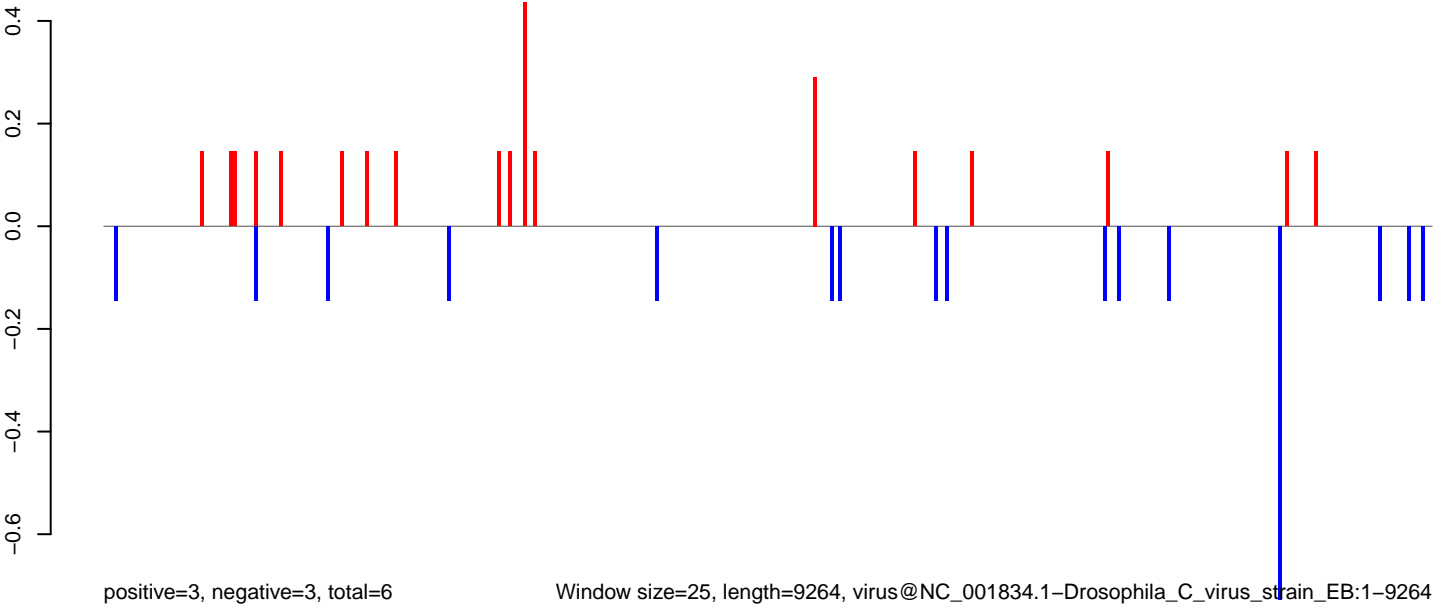
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



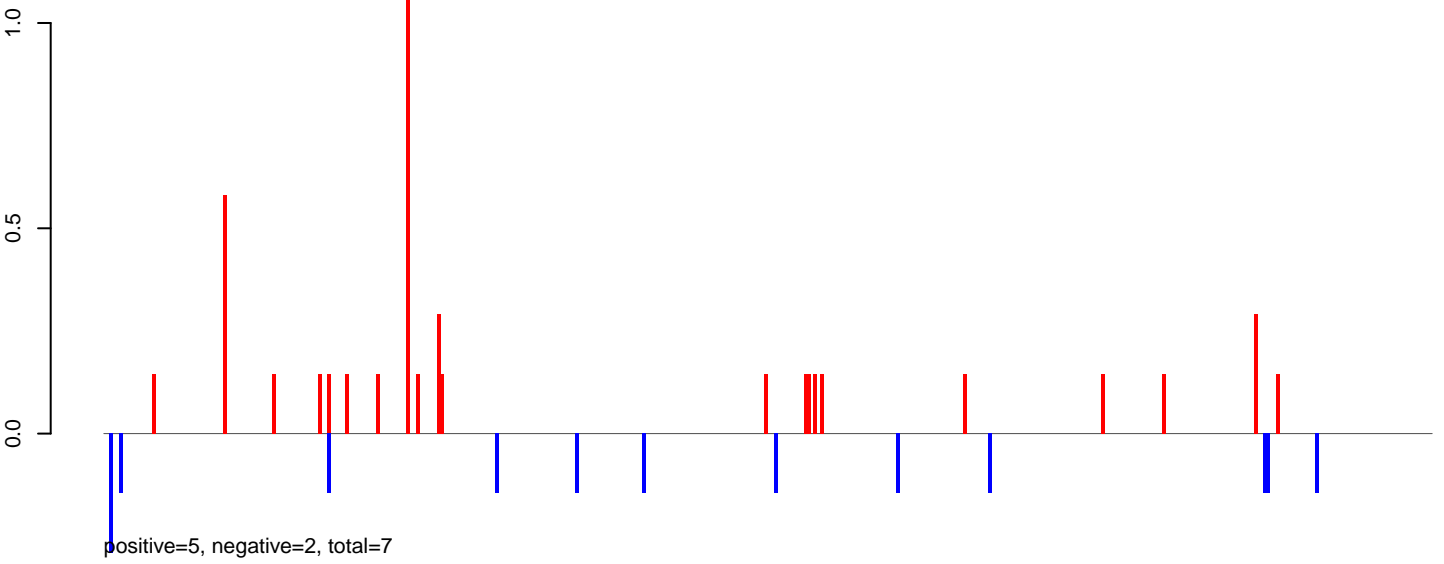
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



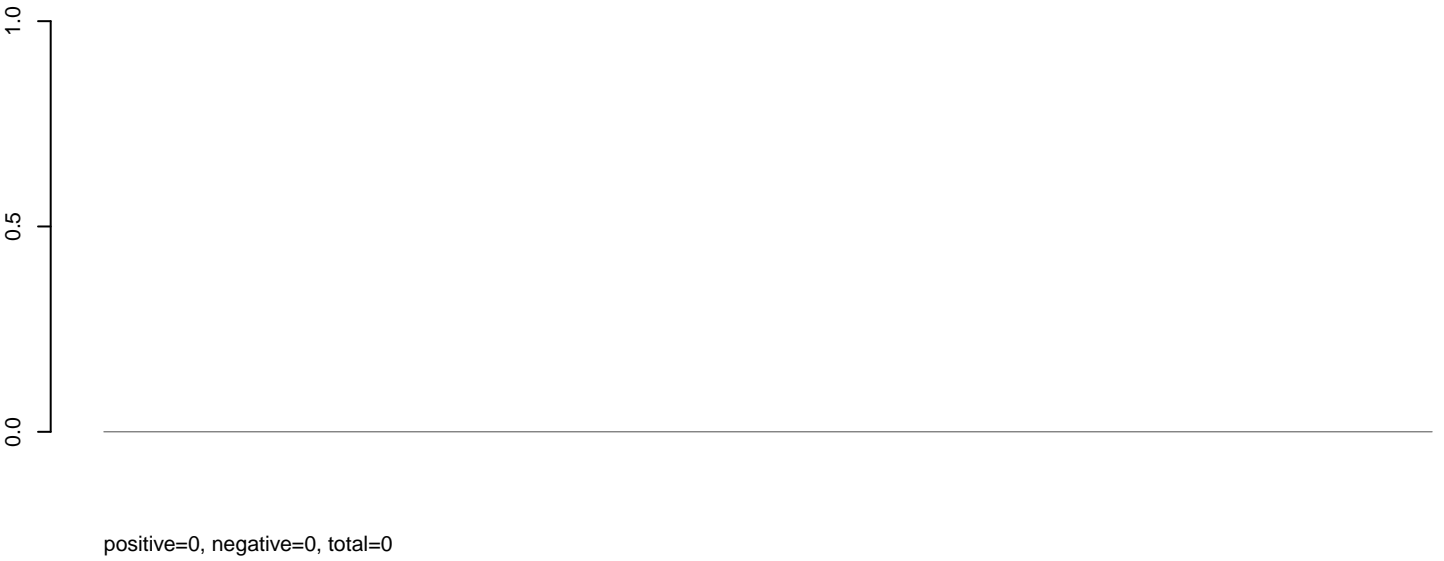
AeAeg_LVP_pMos3DBhspPBac.rep



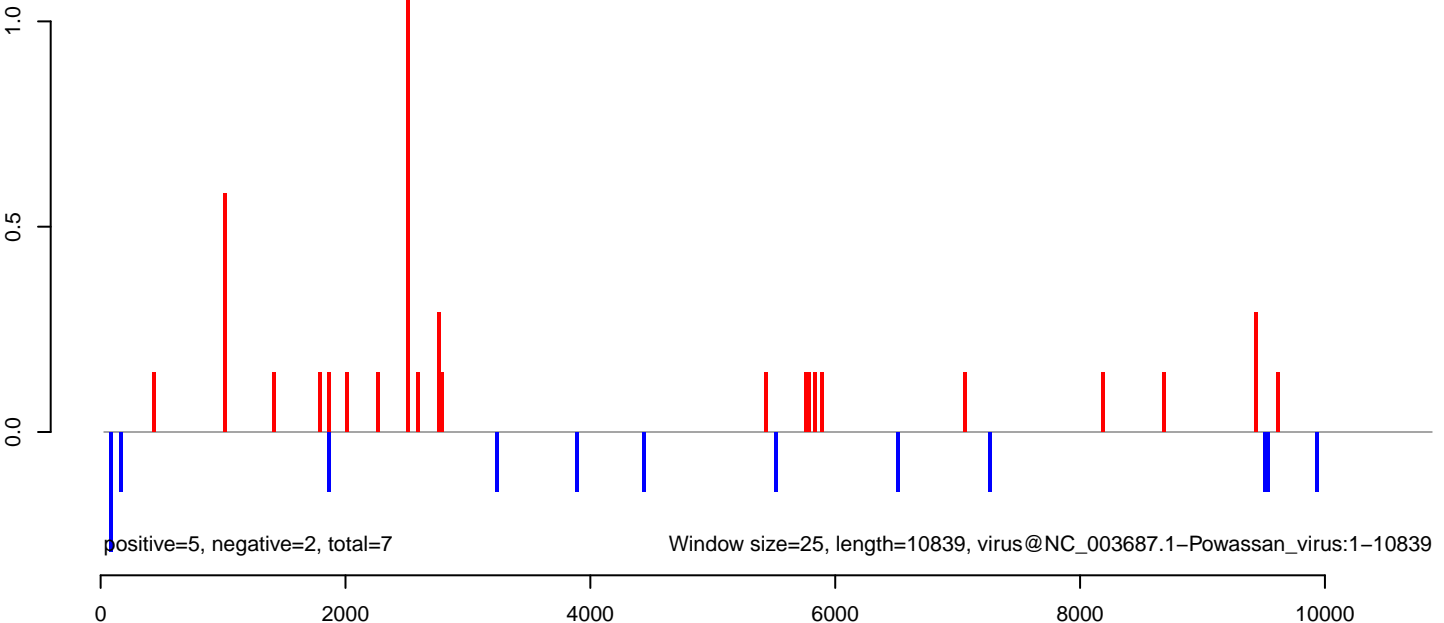
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



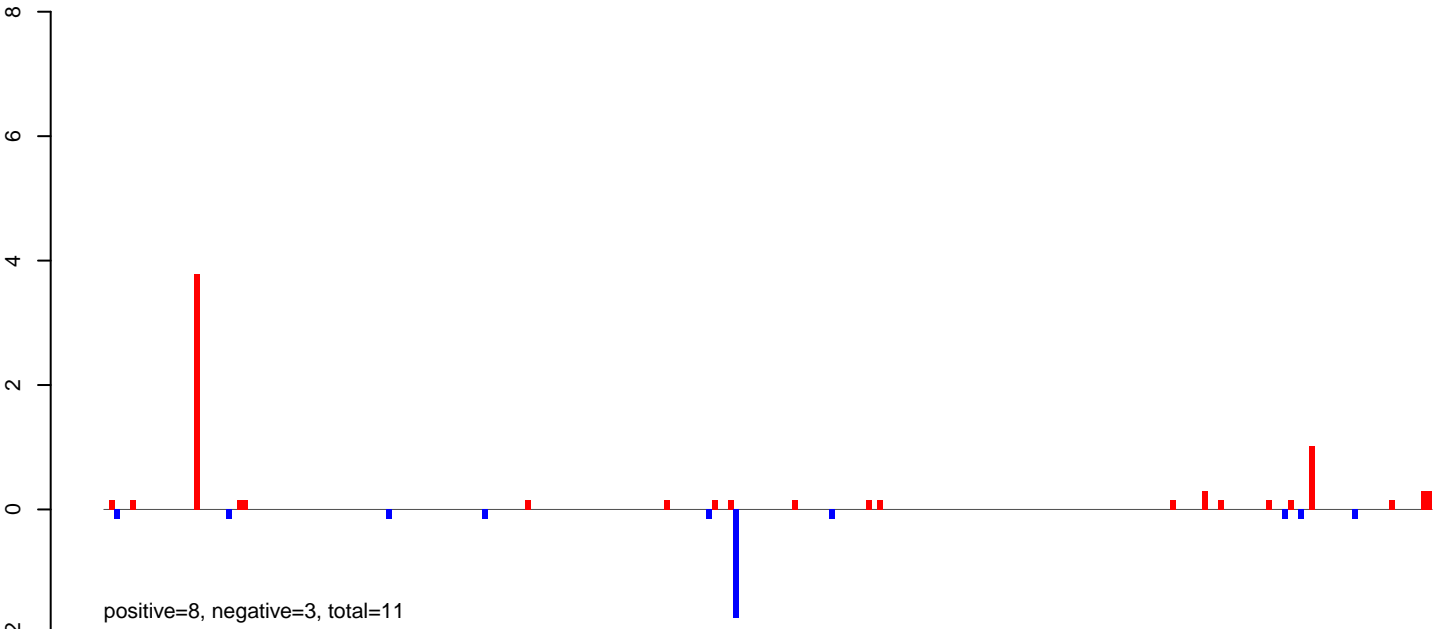
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



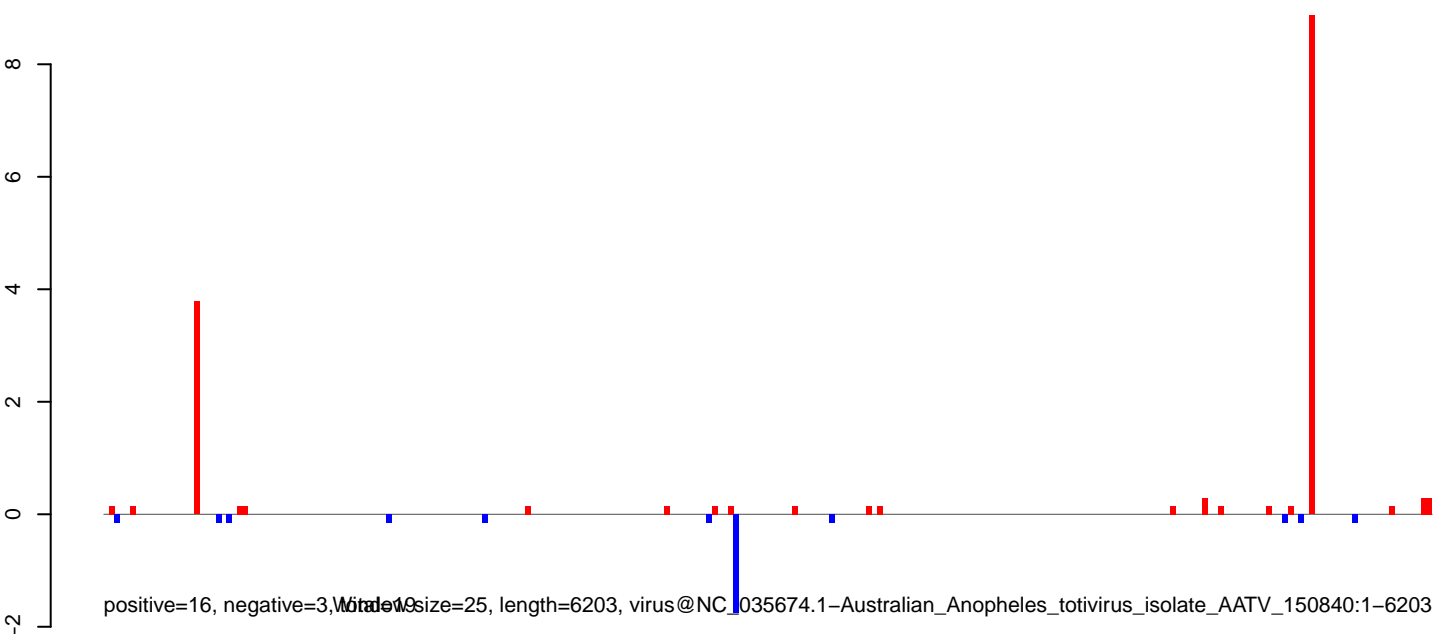
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



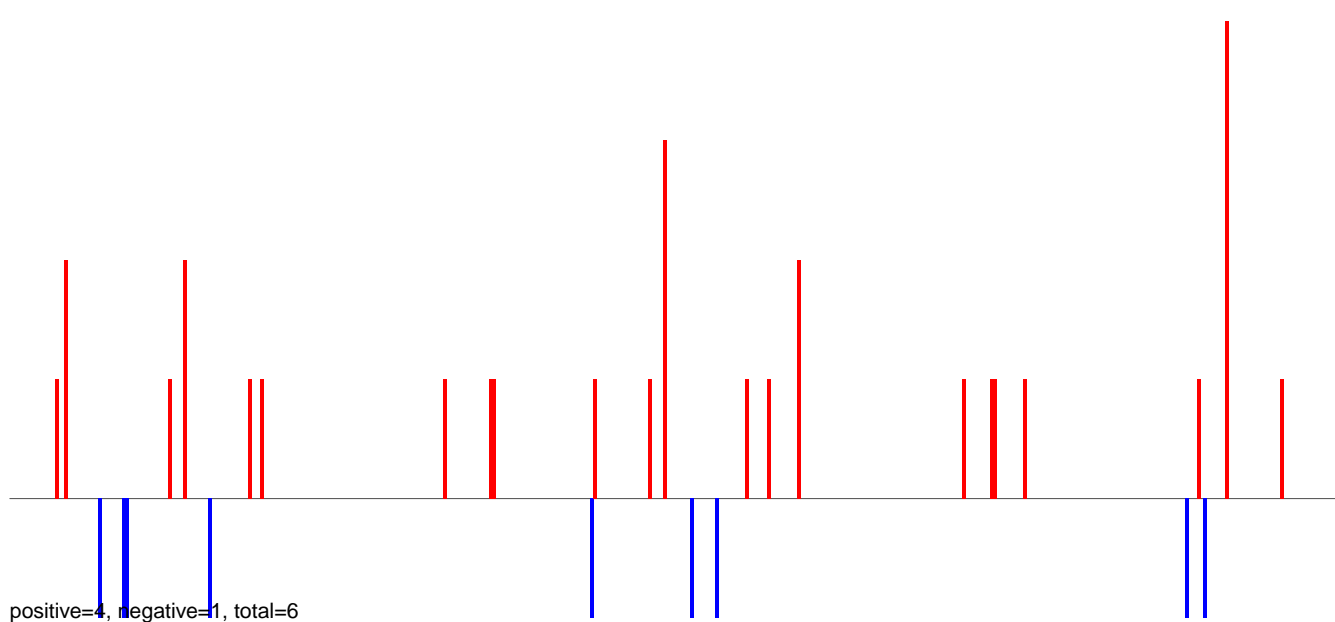
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



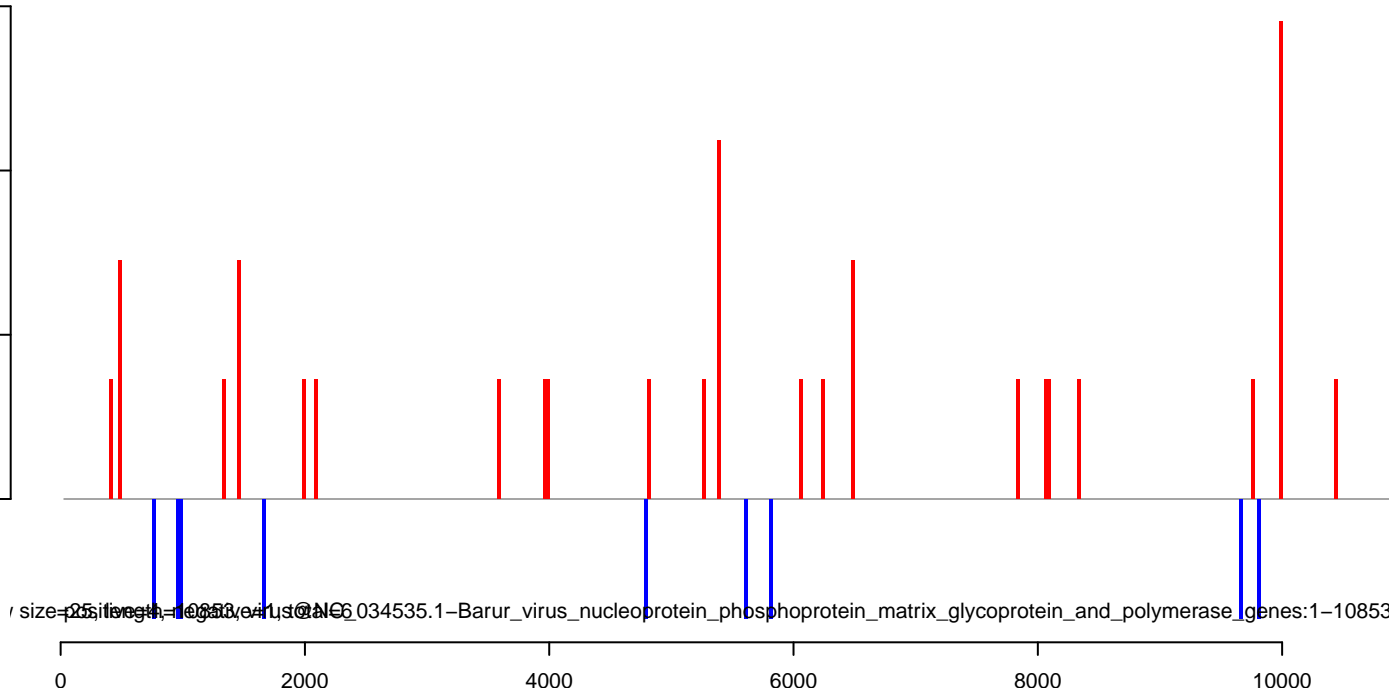
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



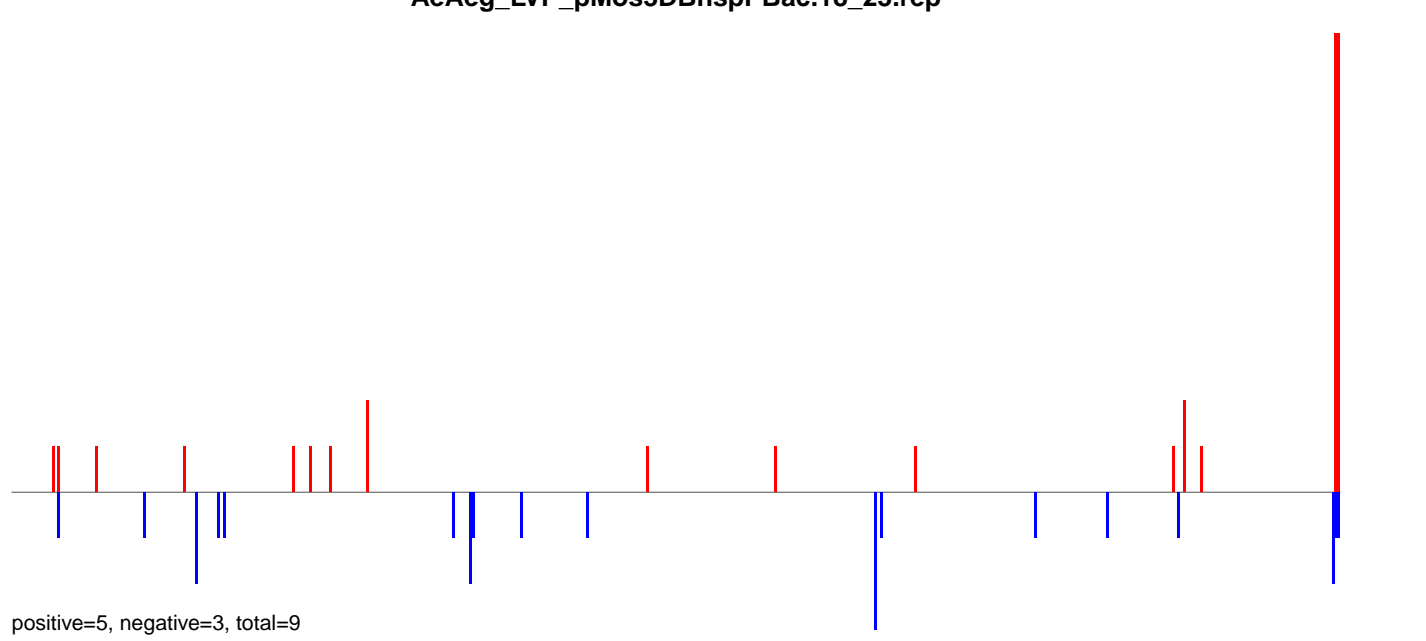
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



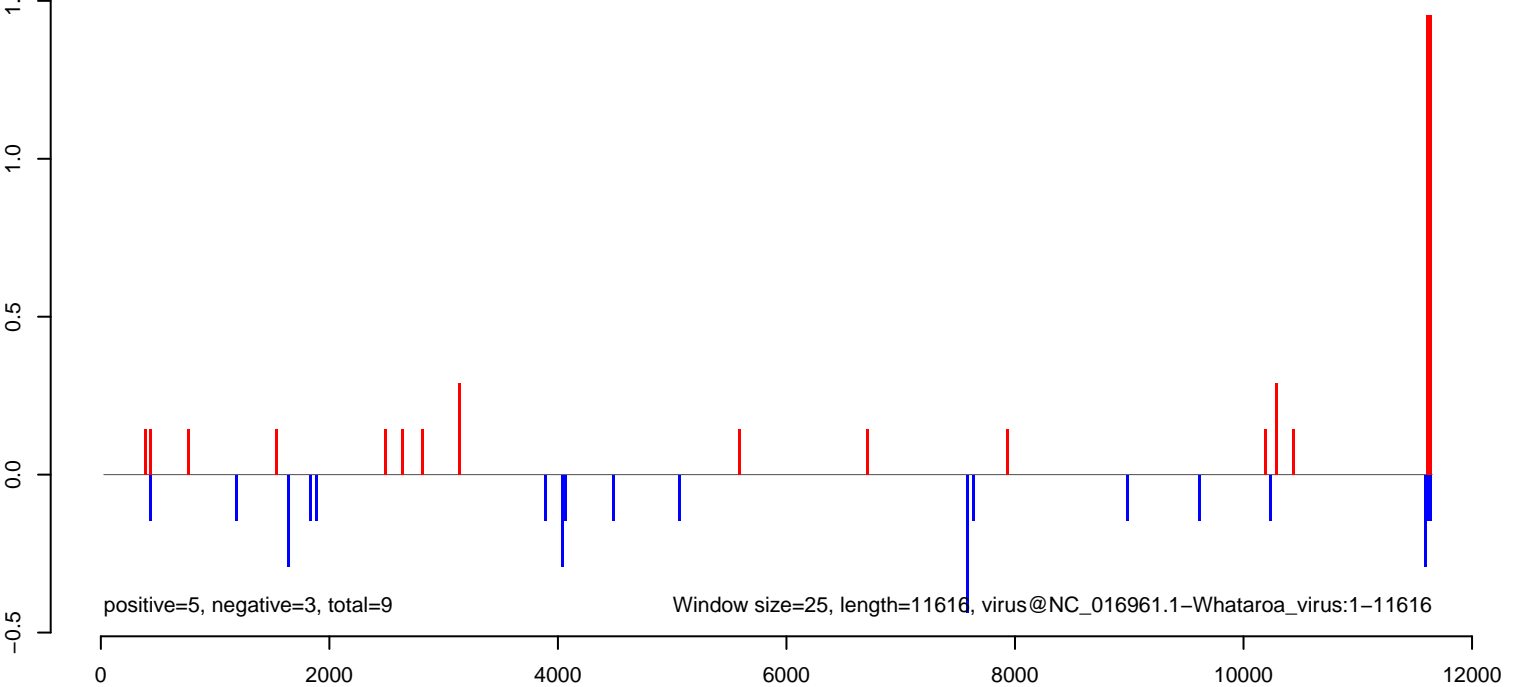
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



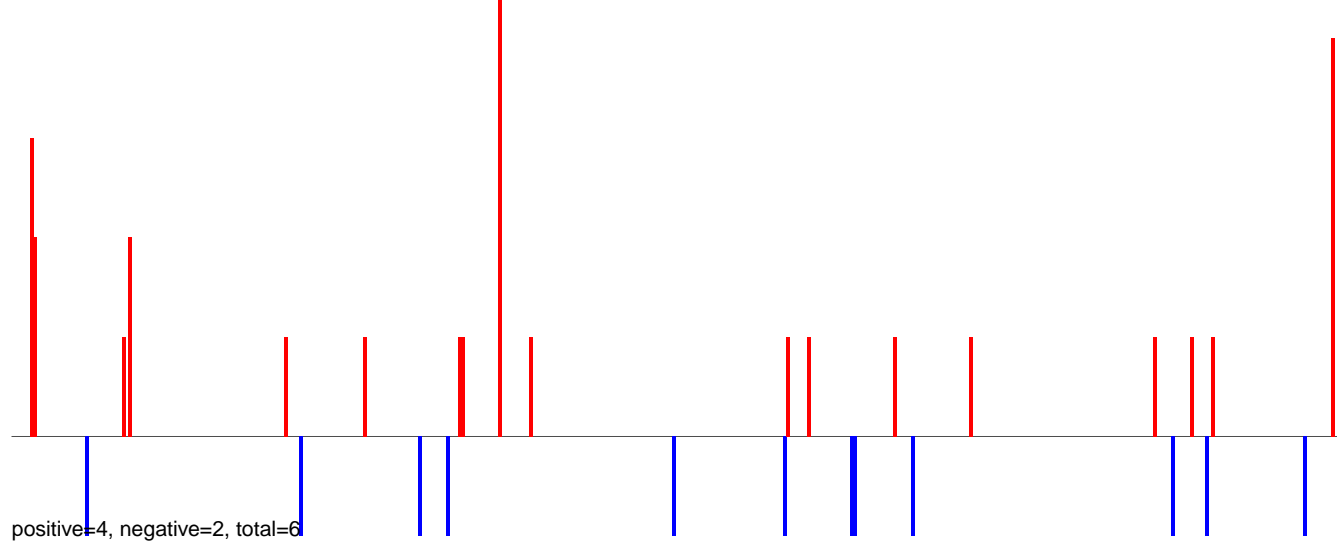
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



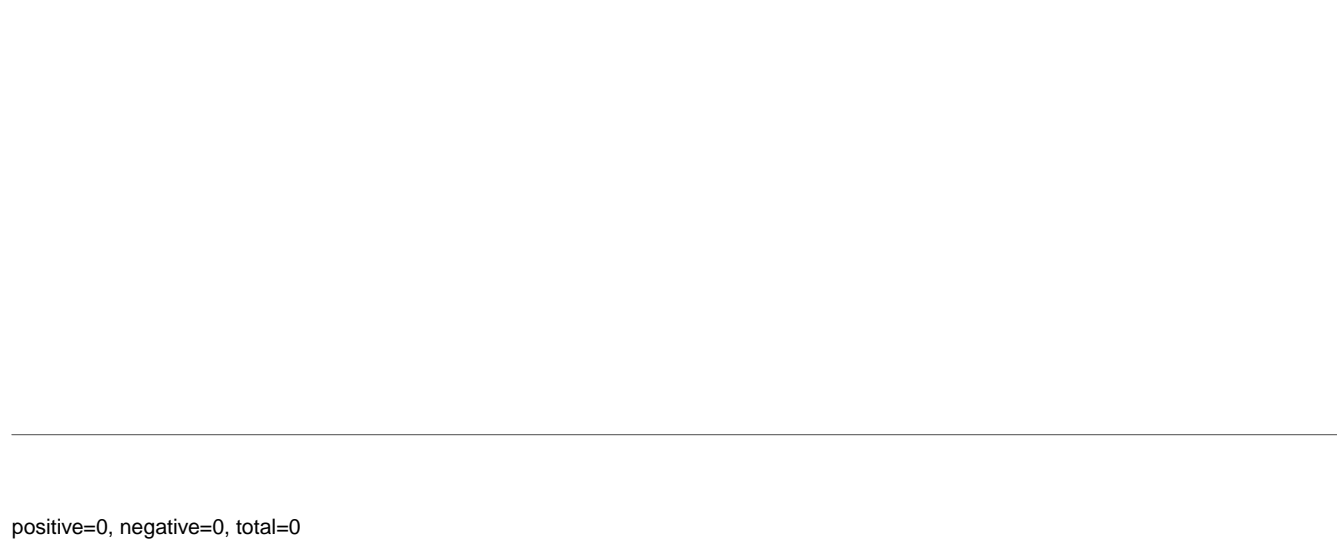
AeAeg_LVP_pMos3DBhspPBac.rep



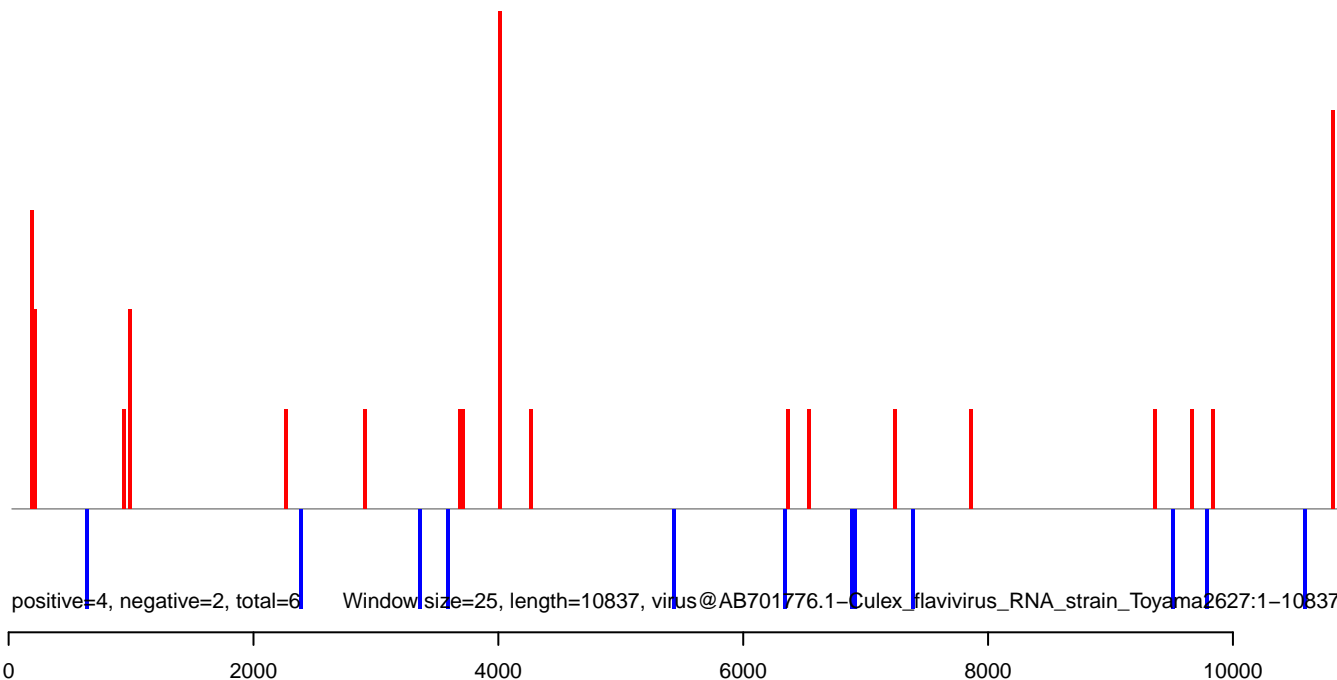
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



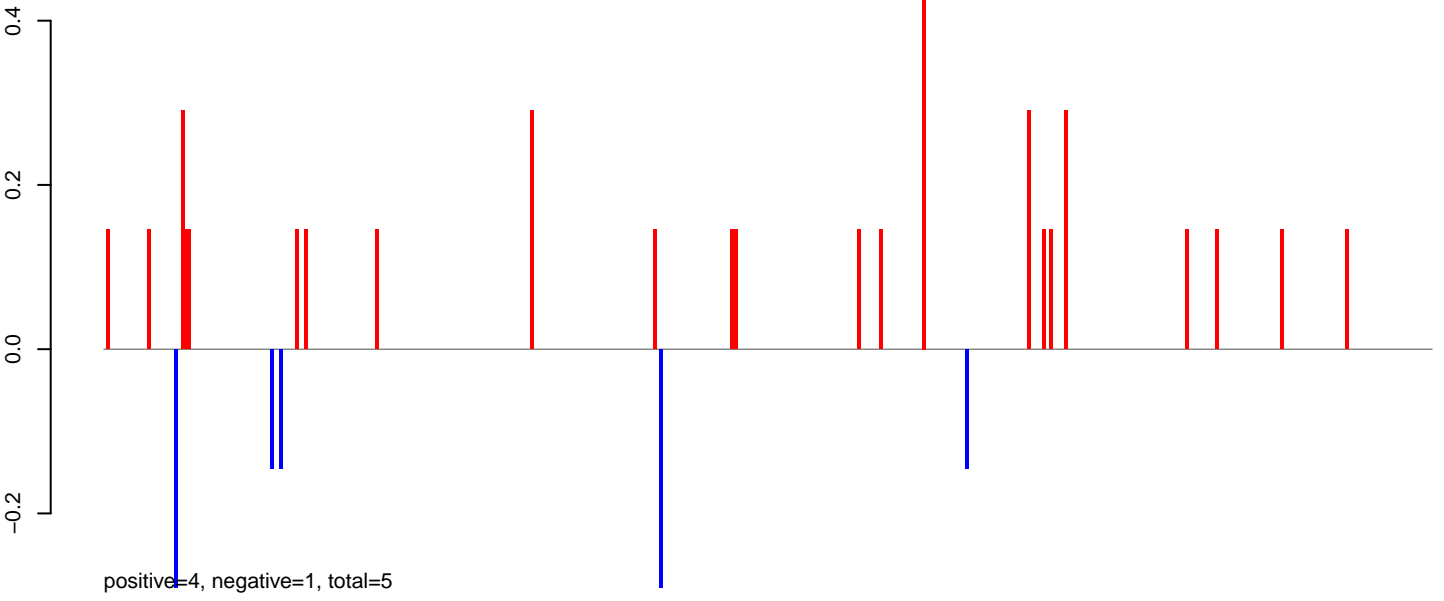
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



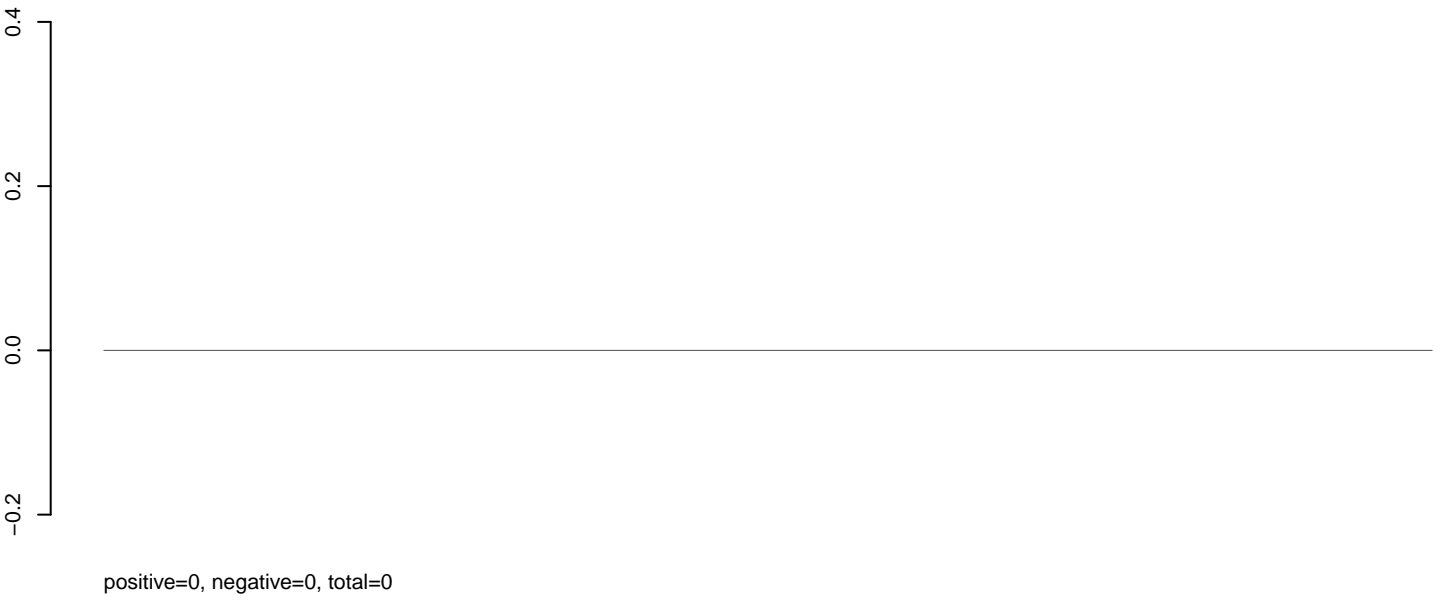
AeAeg_LVP_pMos3DBhspPBac.rep



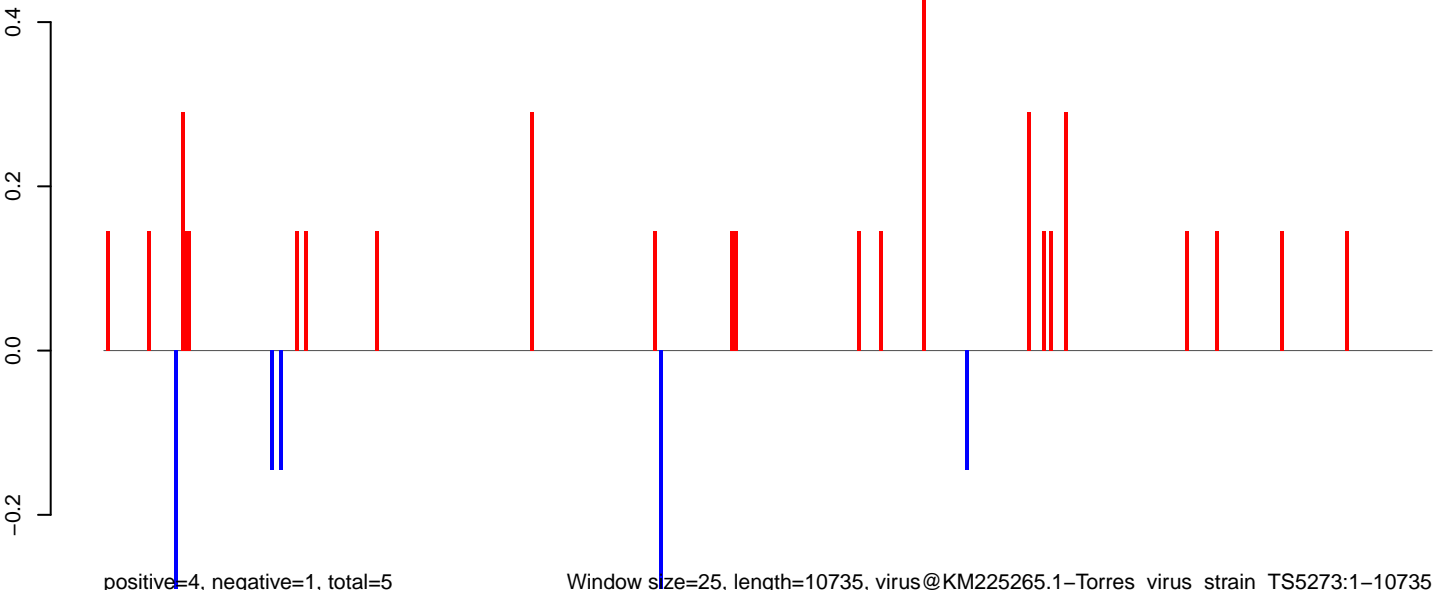
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

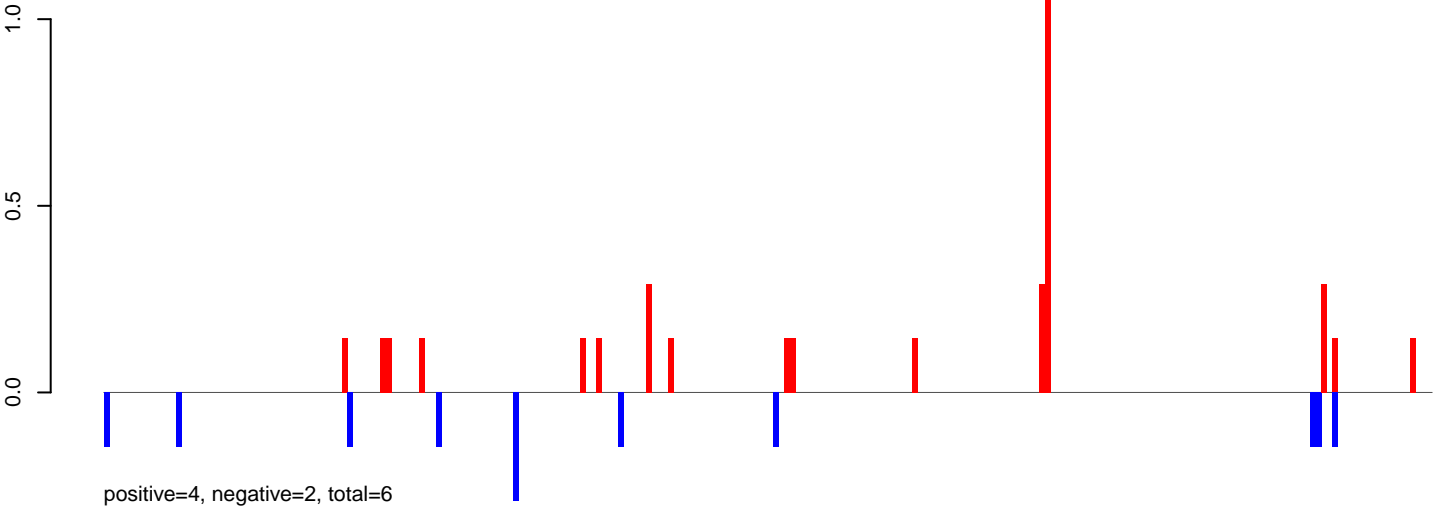


AeAeg_LVP_pMos3DBhspPBac.rep

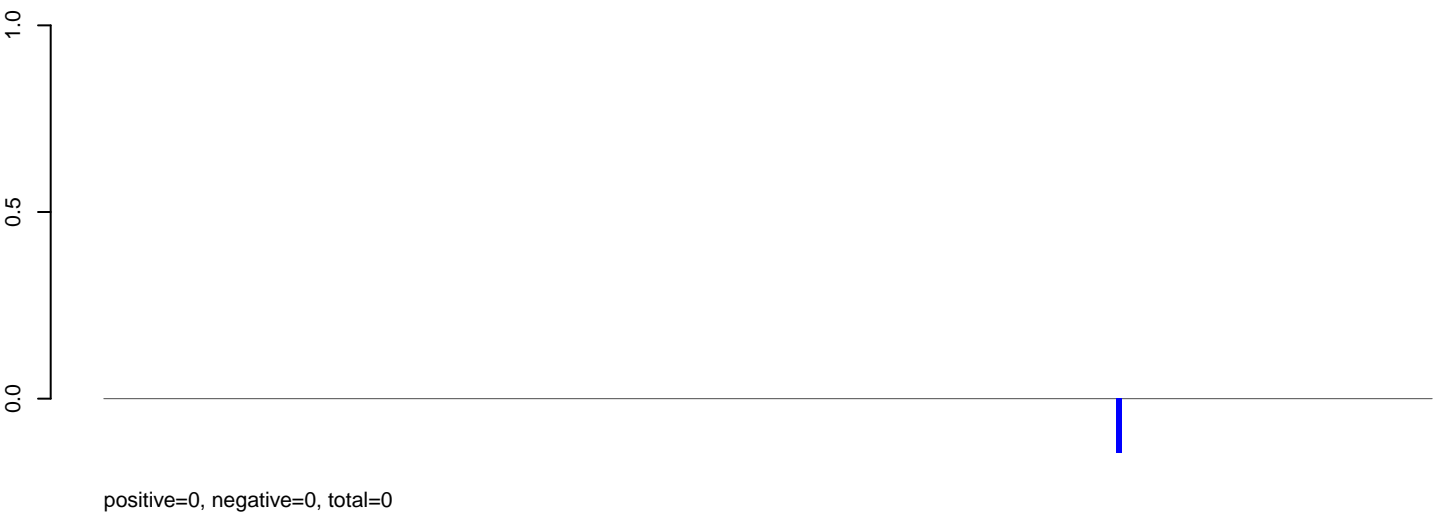


0 2000 4000 6000 8000 10000

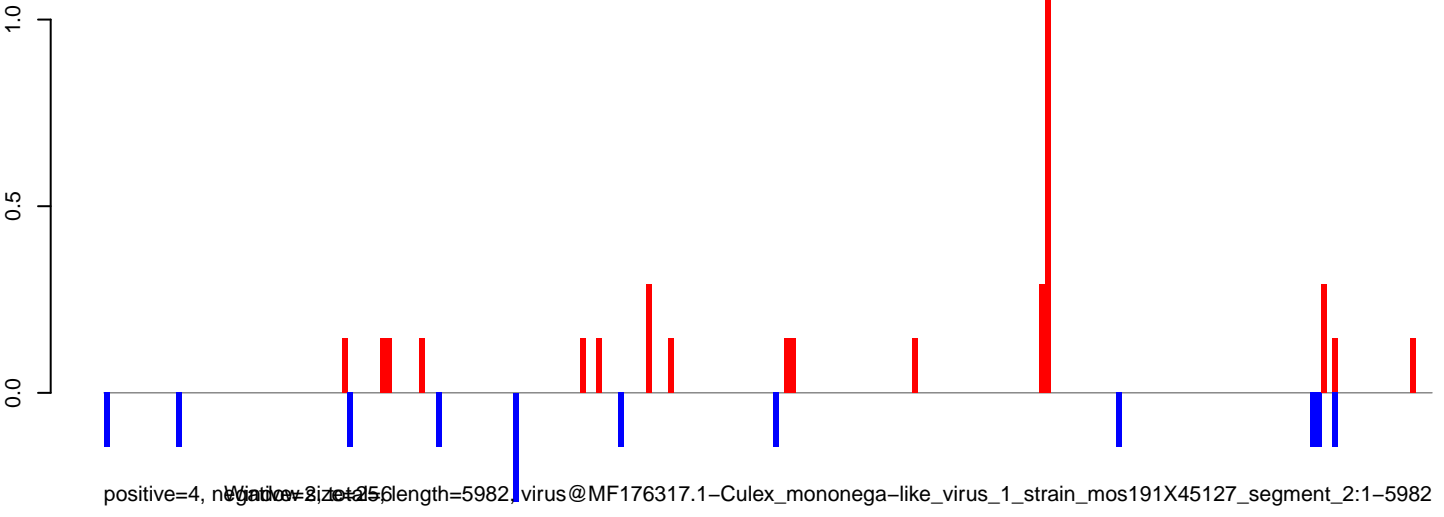
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

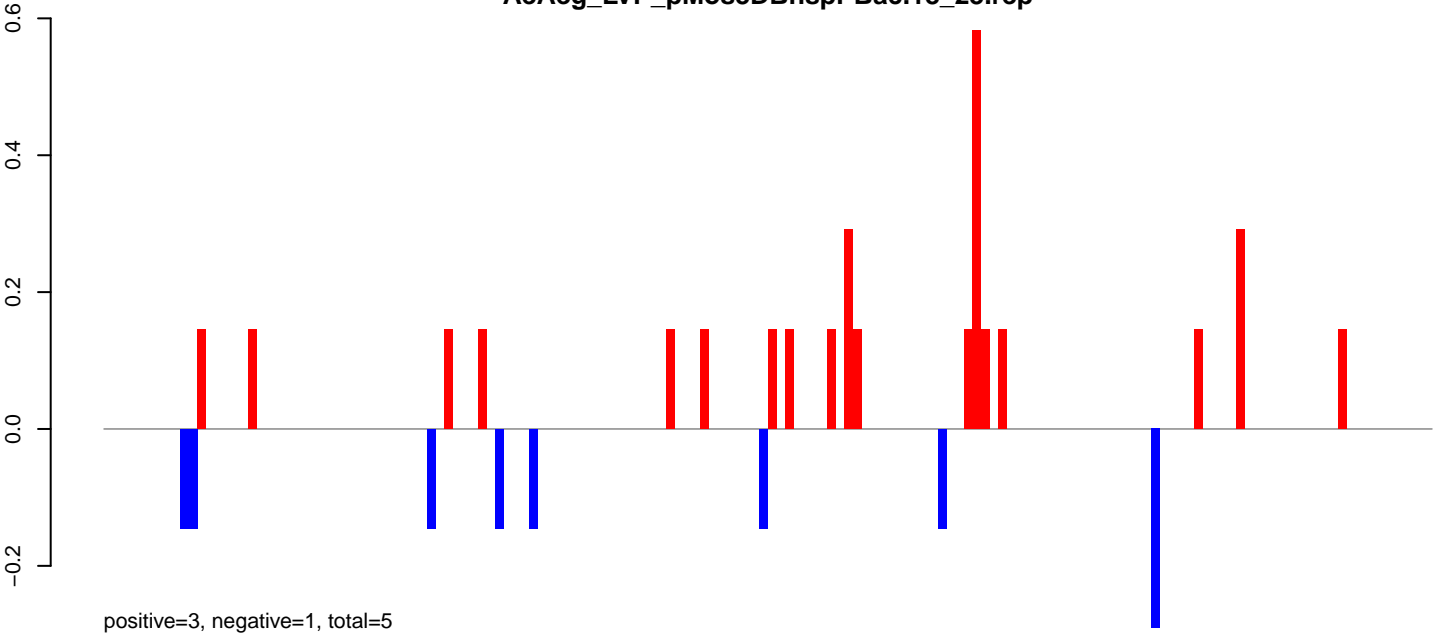


AeAeg_LVP_pMos3DBhspPBac.rep

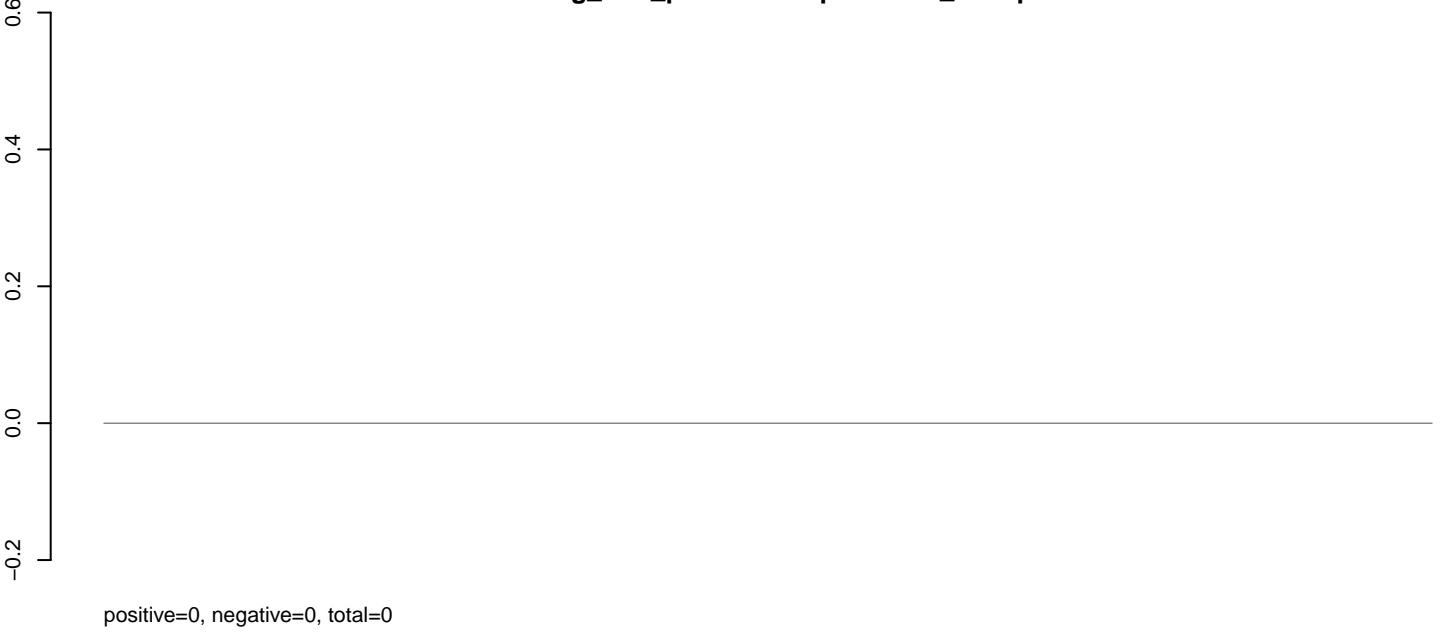


positive=4, negative=2, total=6, length=5982, virus@MF176317.1-Culex_mononega-like_virus_1_strain_mos191X45127_segment_2:1-5982

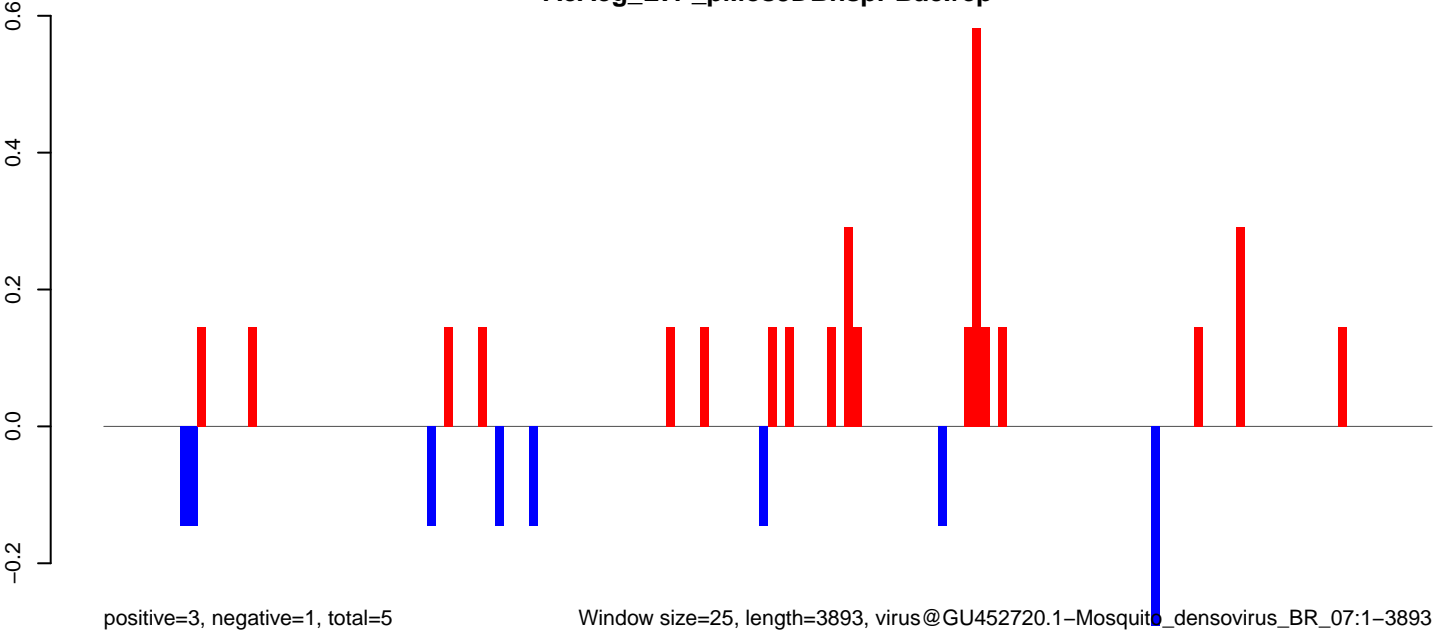
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

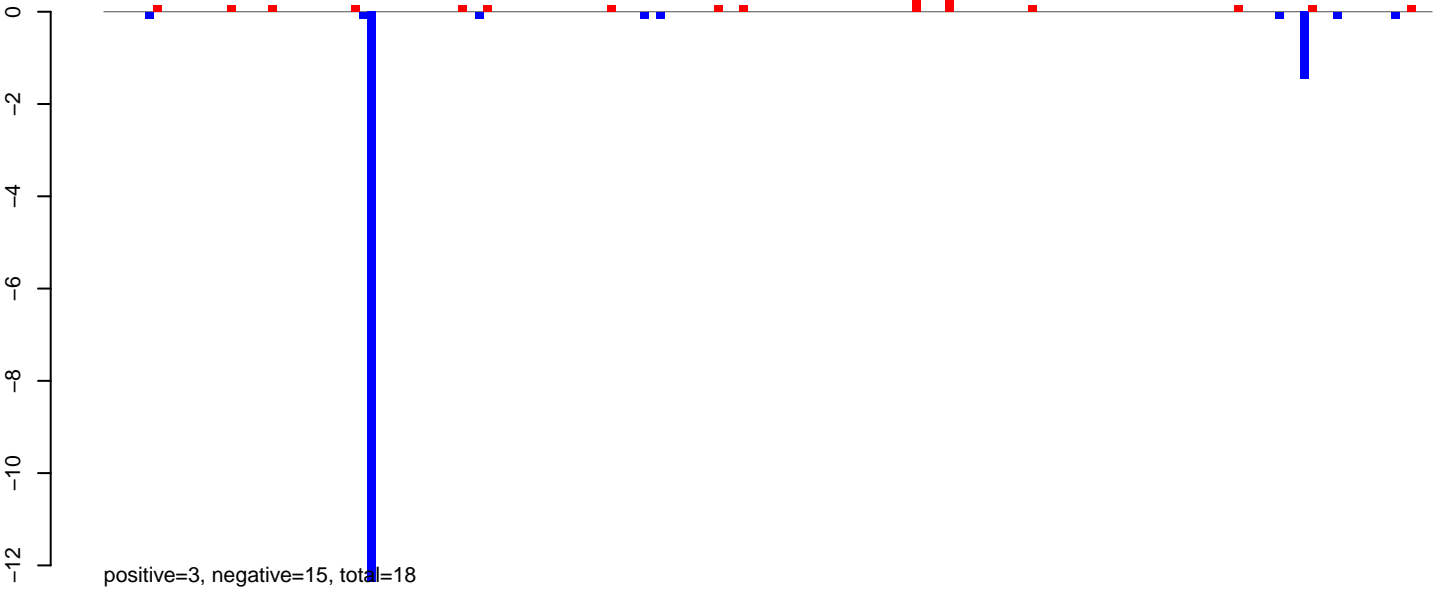


AeAeg_LVP_pMos3DBhspPBac.rep

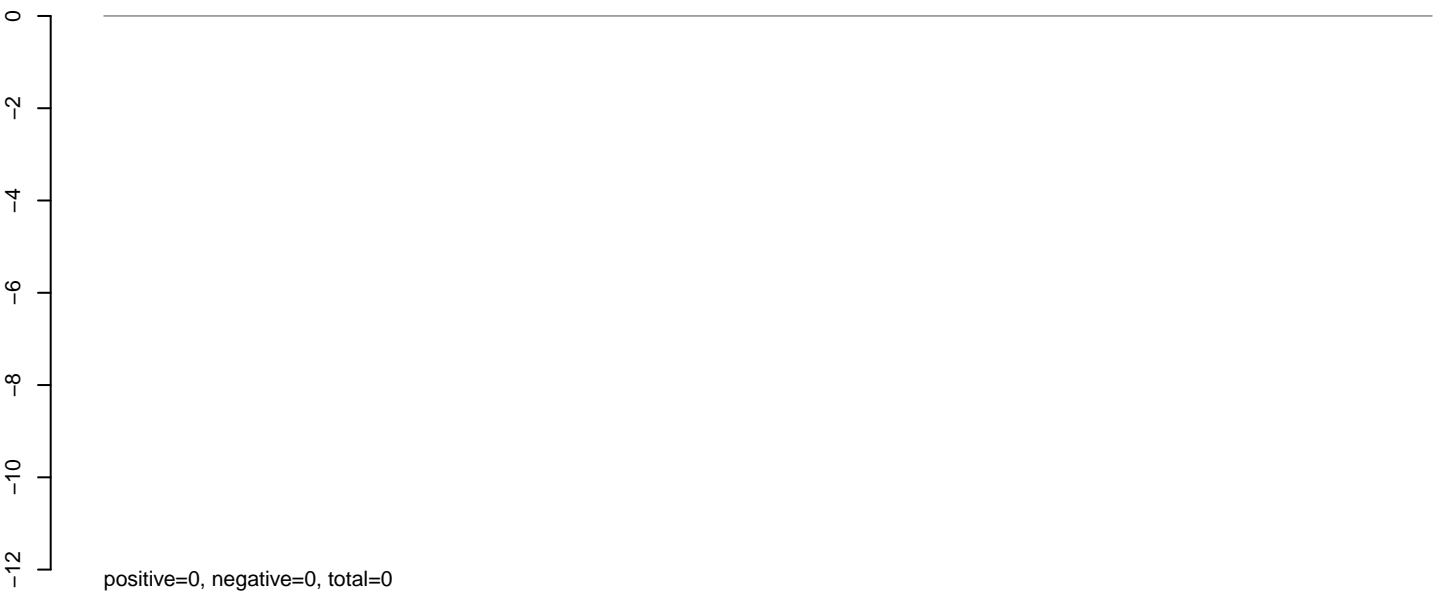


0 1000 2000 3000 4000

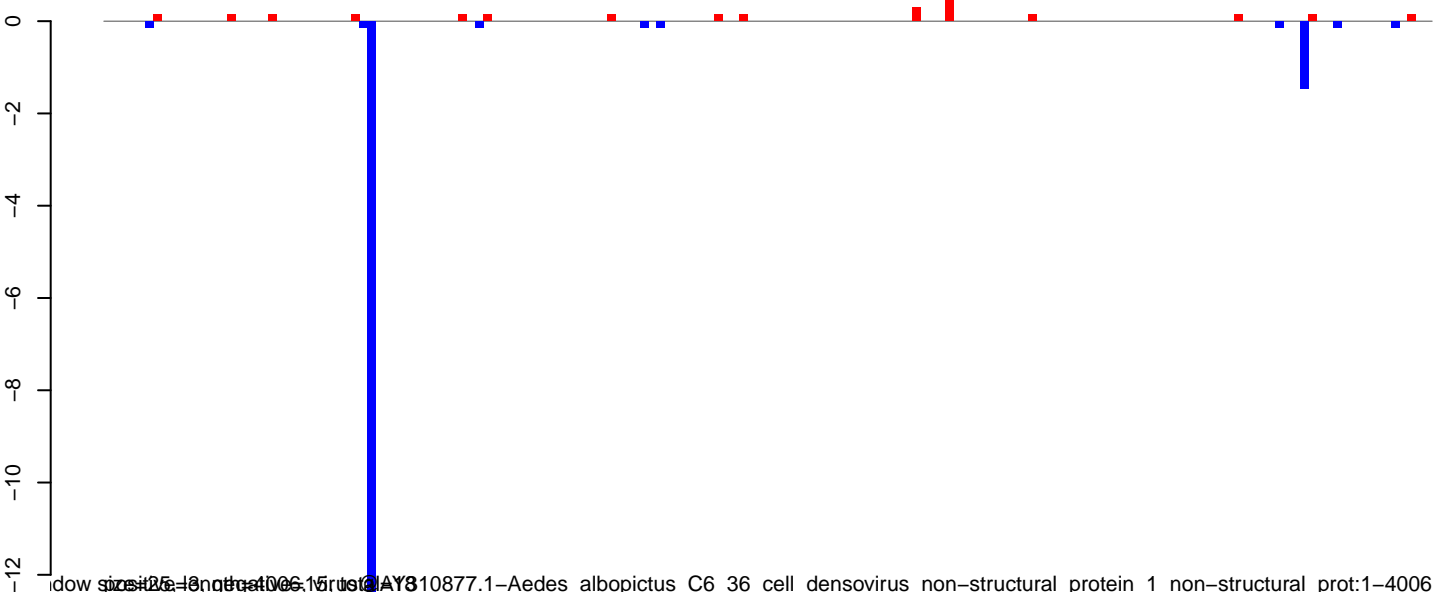
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



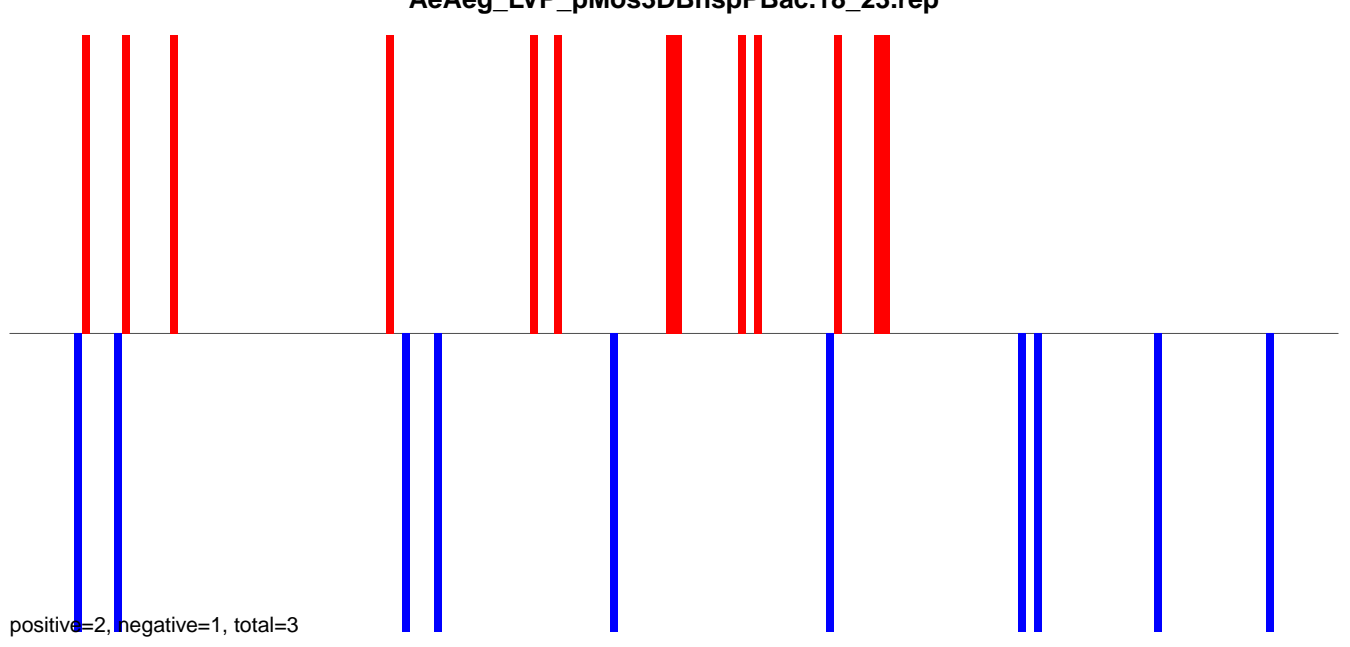
AeAeg_LVP_pMos3DBhspPBac.rep



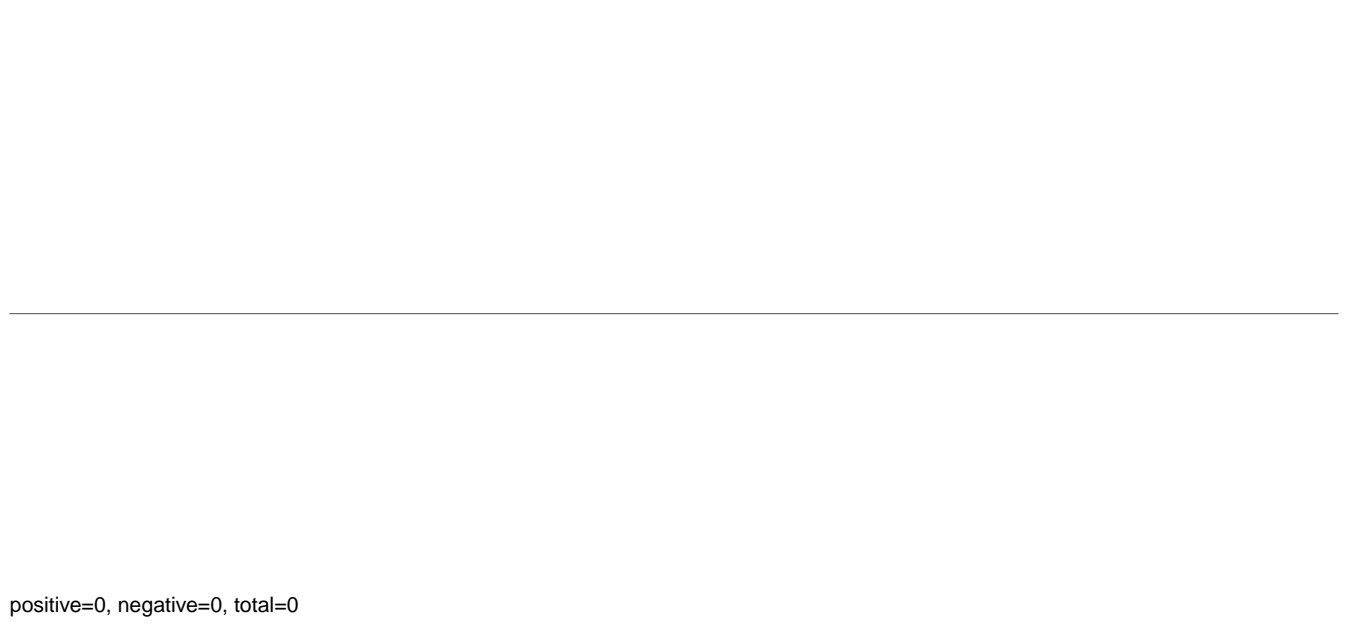
downloaded from <https://www.ncbi.nlm.nih.gov/nuclot/AY110877.1> Aedes albopictus C6_36_cell_densovirus_non-structural_protein_1_non-structural_prot:1-4006

0 1000 2000 3000 4000

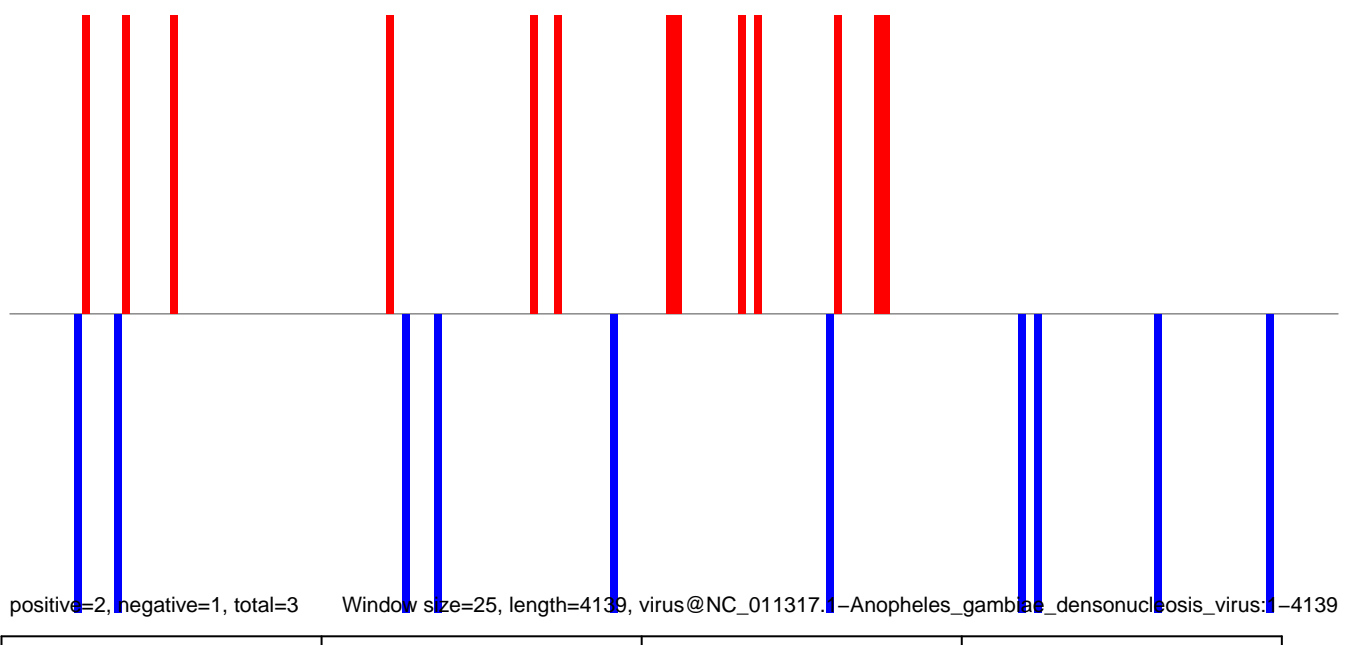
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



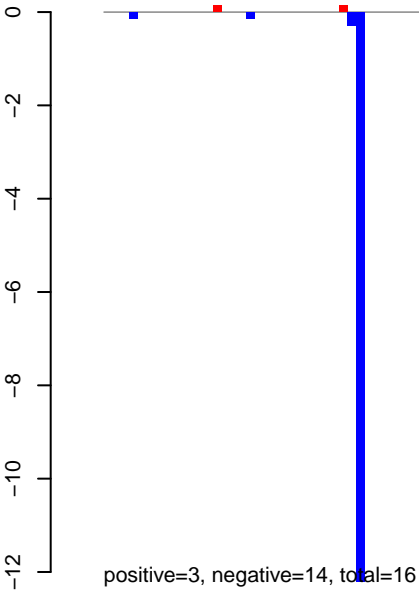
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



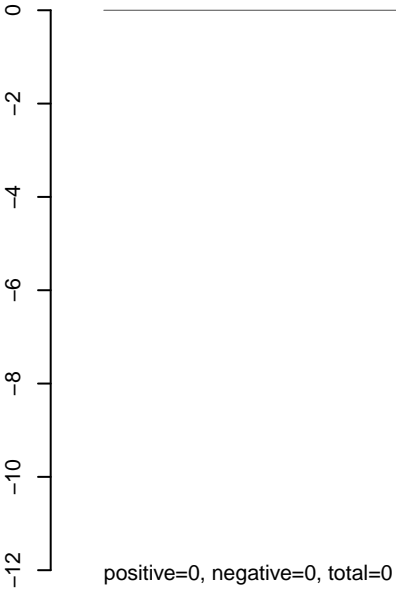
AeAeg_LVP_pMos3DBhspPBac.rep



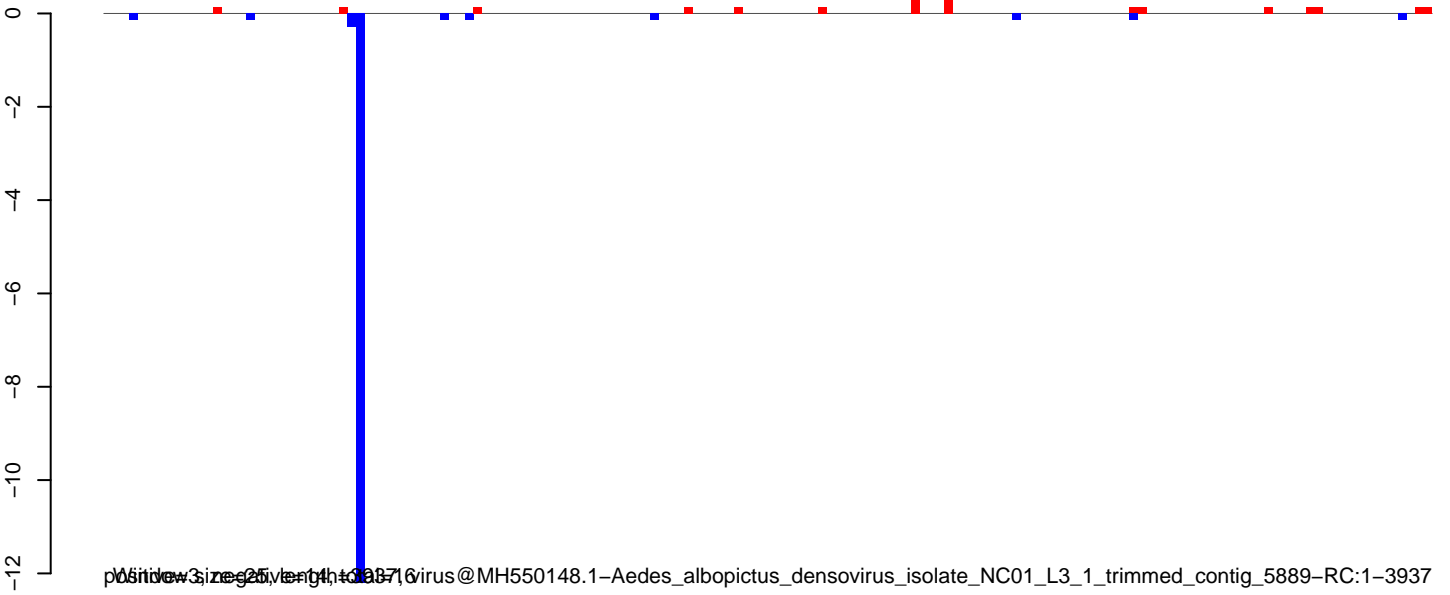
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



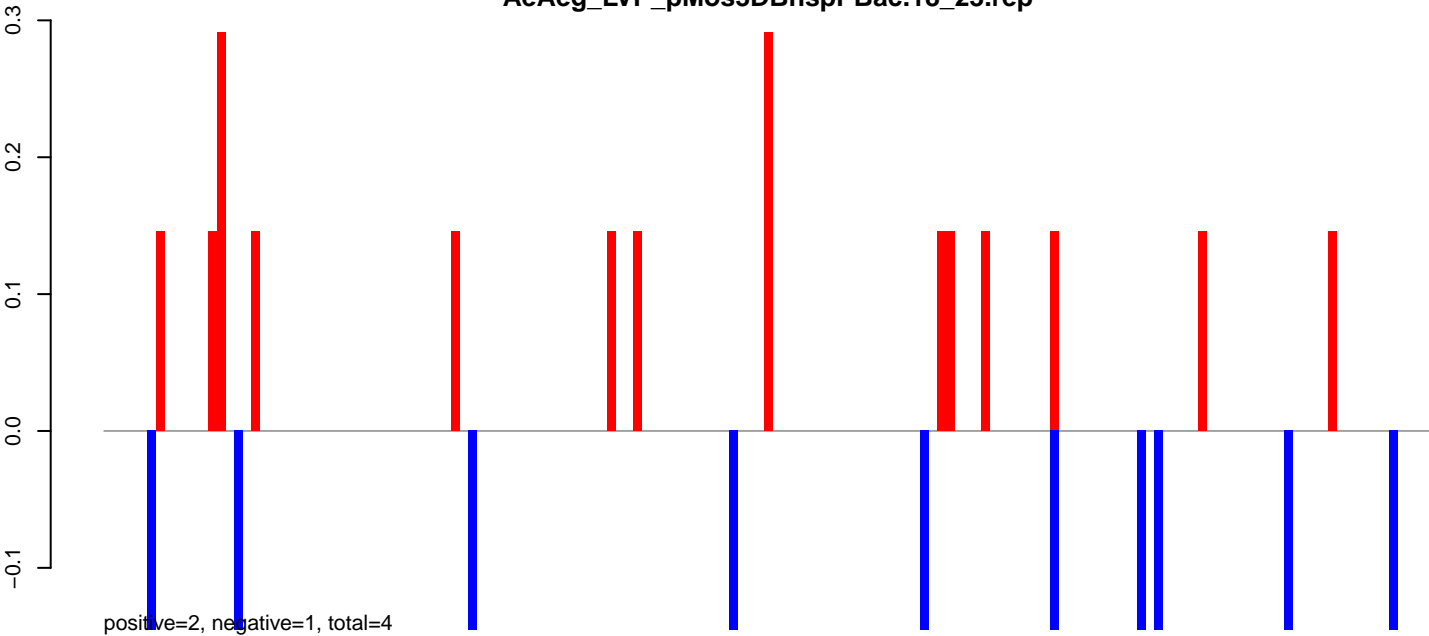
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



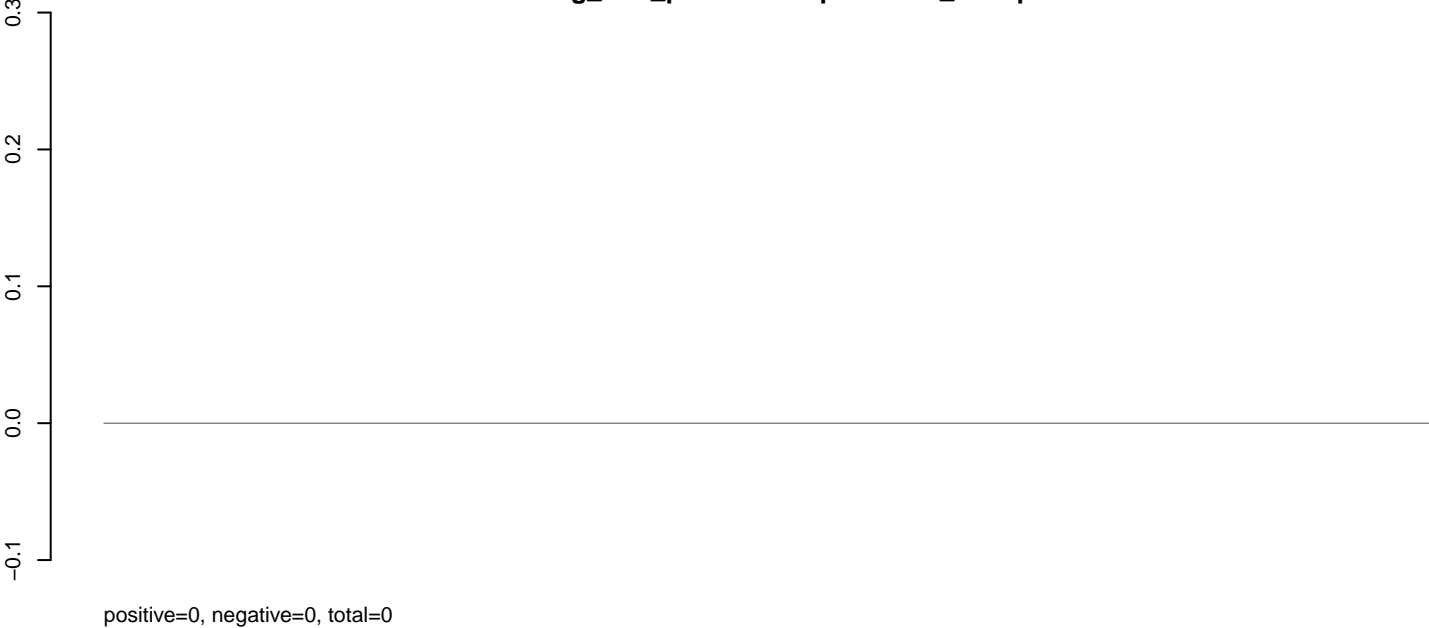
AeAeg_LVP_pMos3DBhspPBac.rep



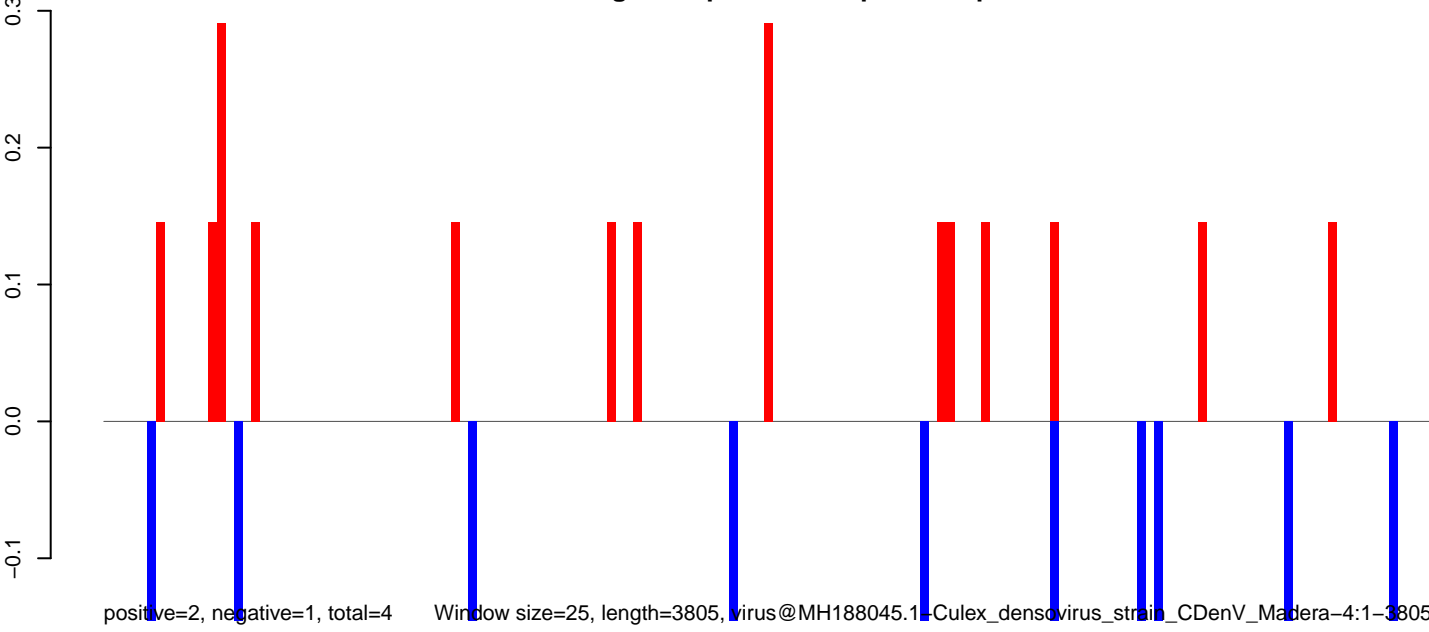
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

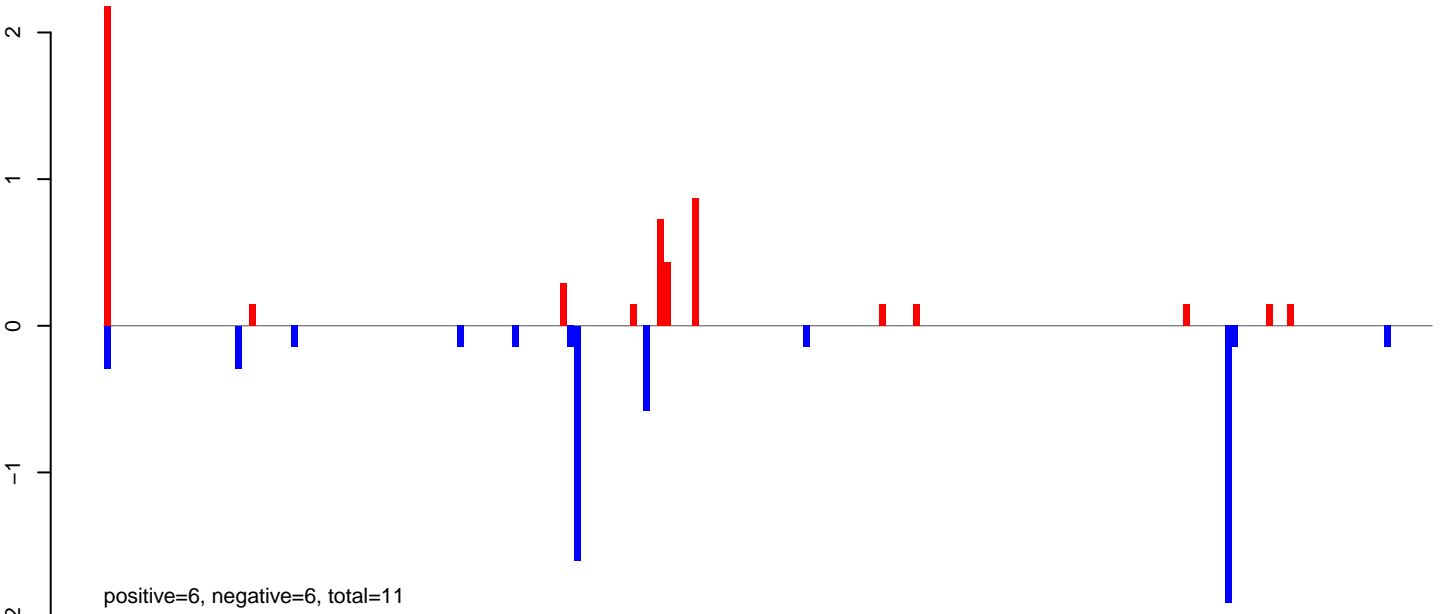


AeAeg_LVP_pMos3DBhspPBac.rep

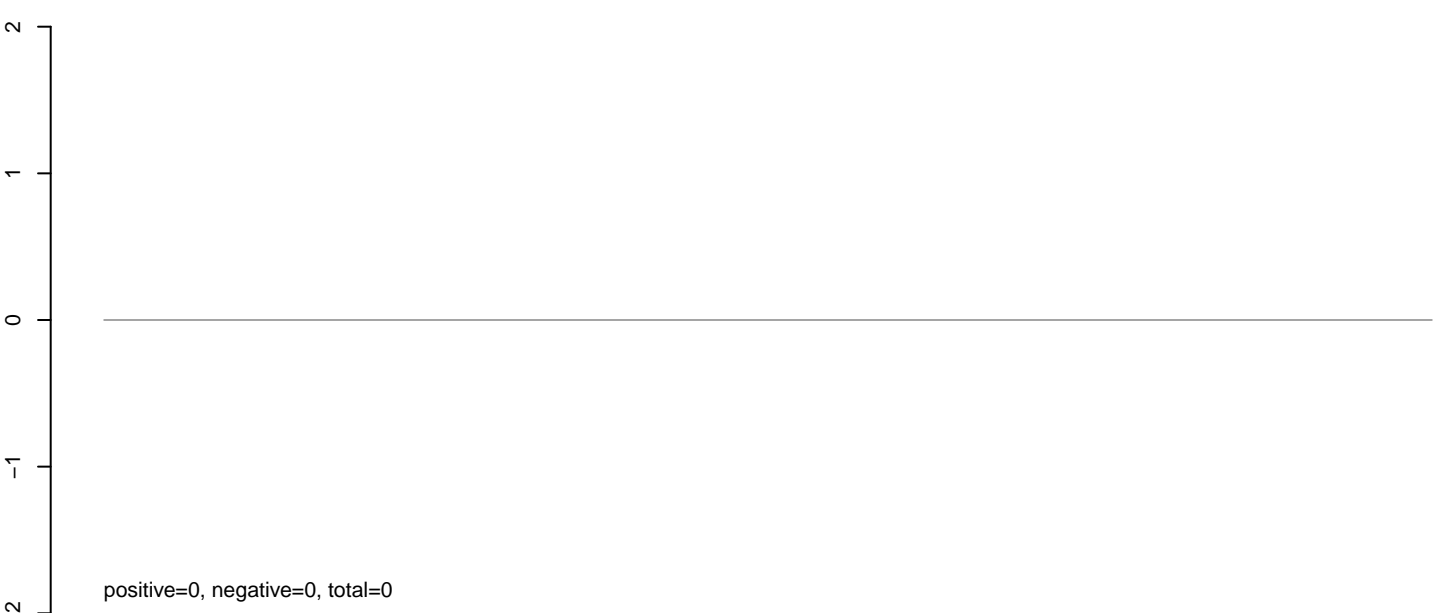


Window size=25, length=3805, virus@MH188045.1-Culex_densovirus_strain_CDENV_Madera-4:1-3805

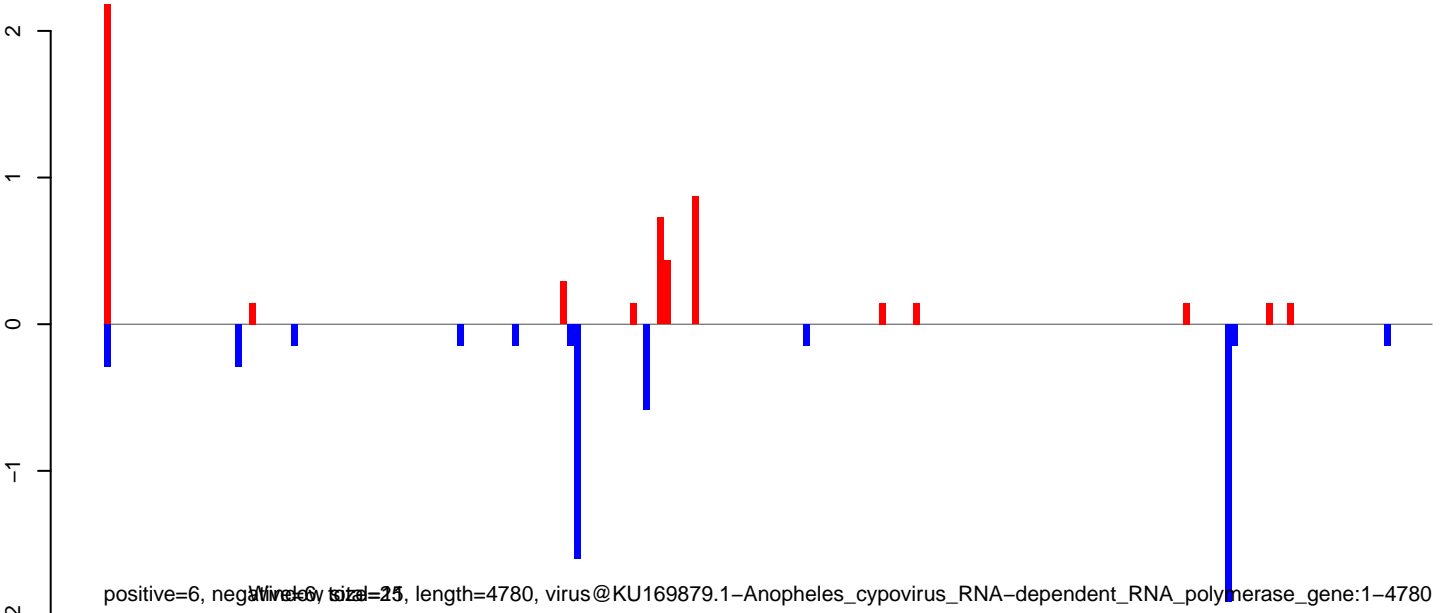
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



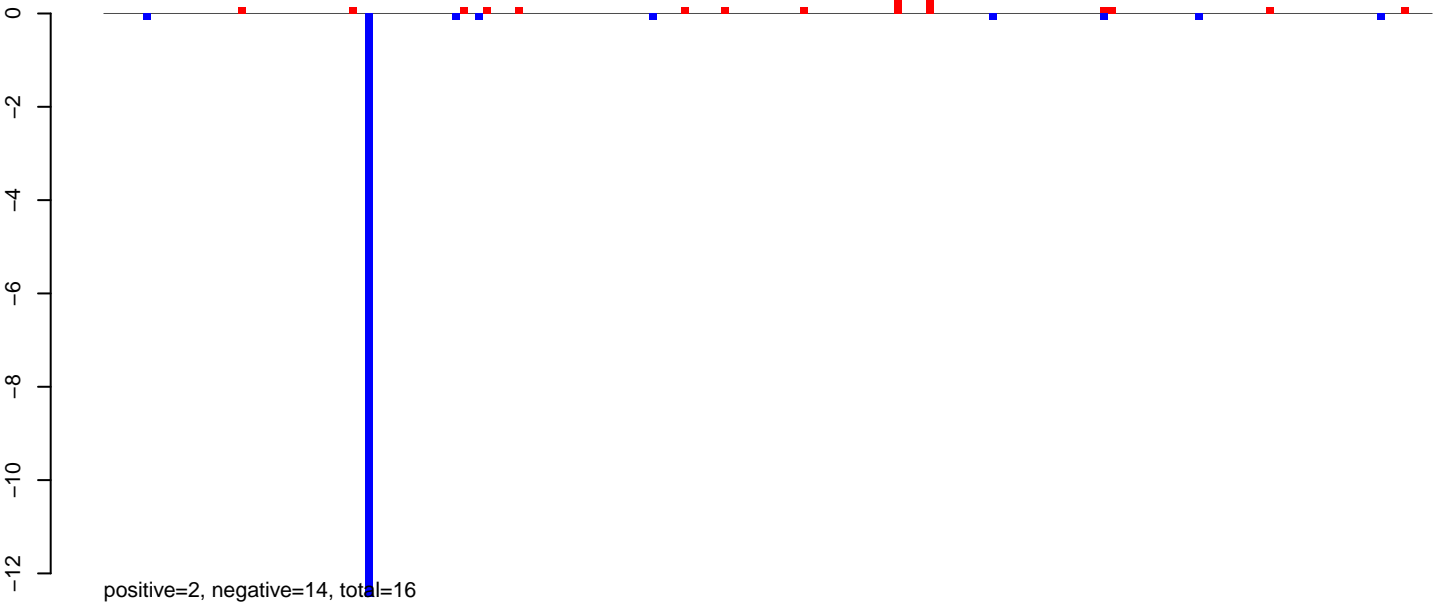
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



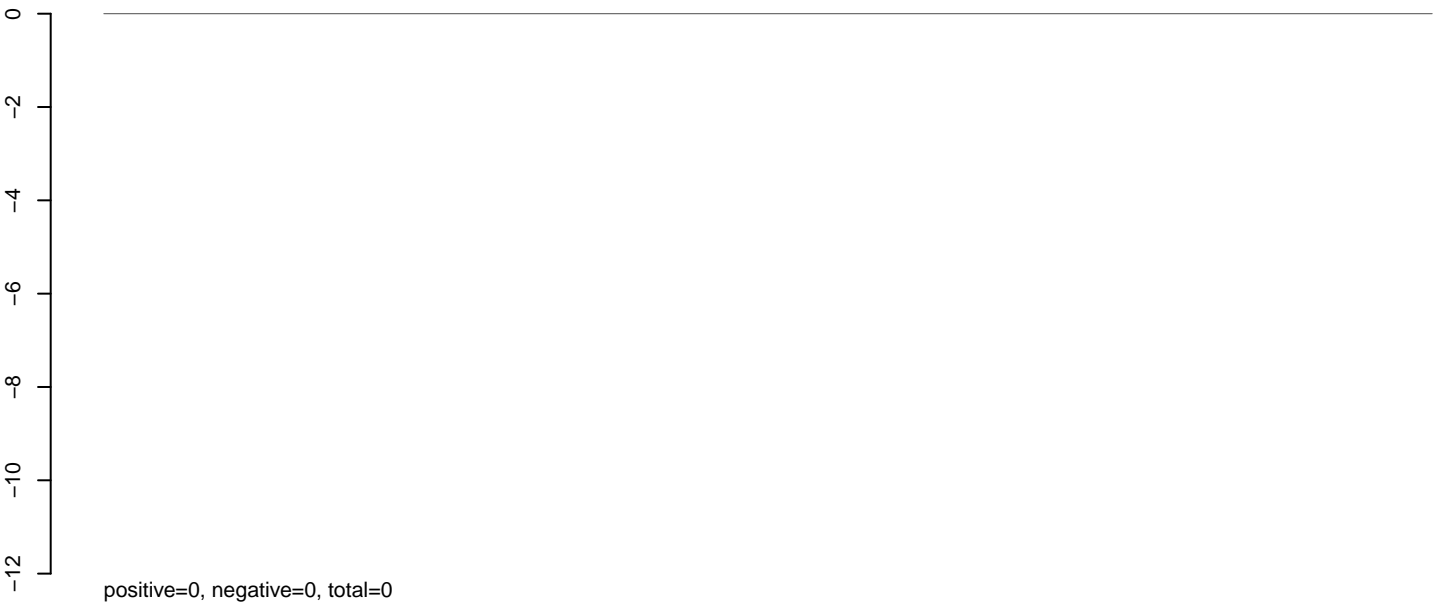
AeAeg_LVP_pMos3DBhspPBac.rep



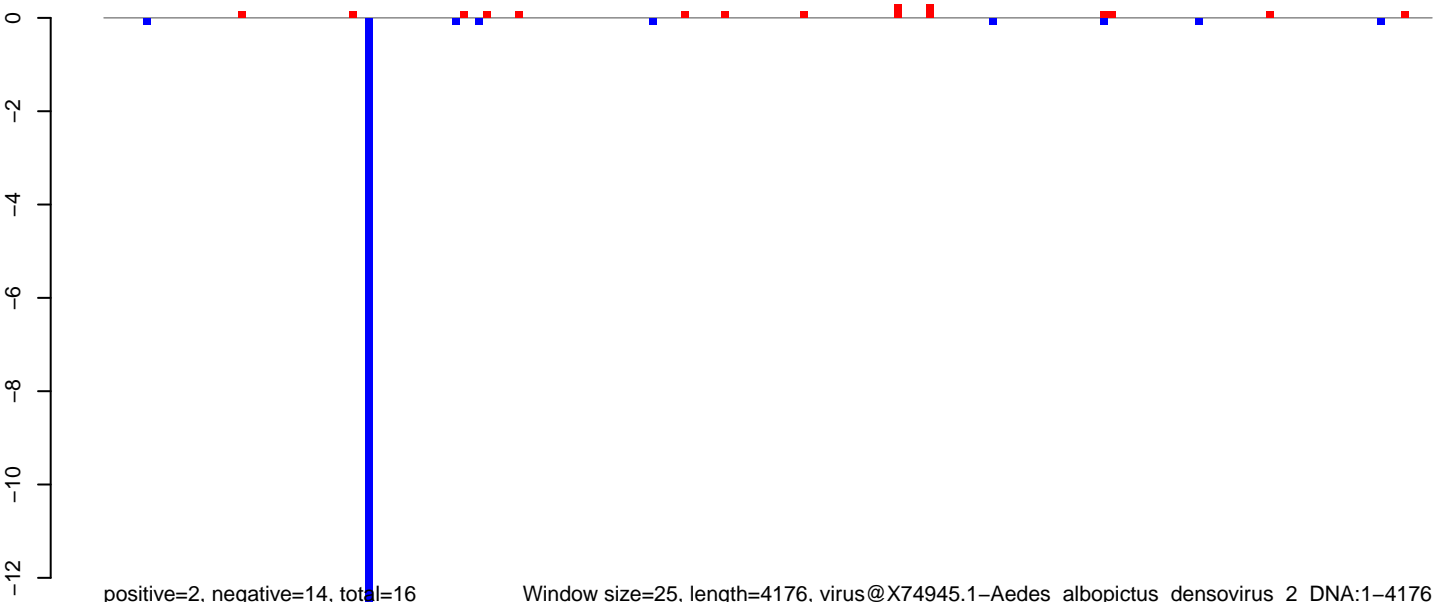
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



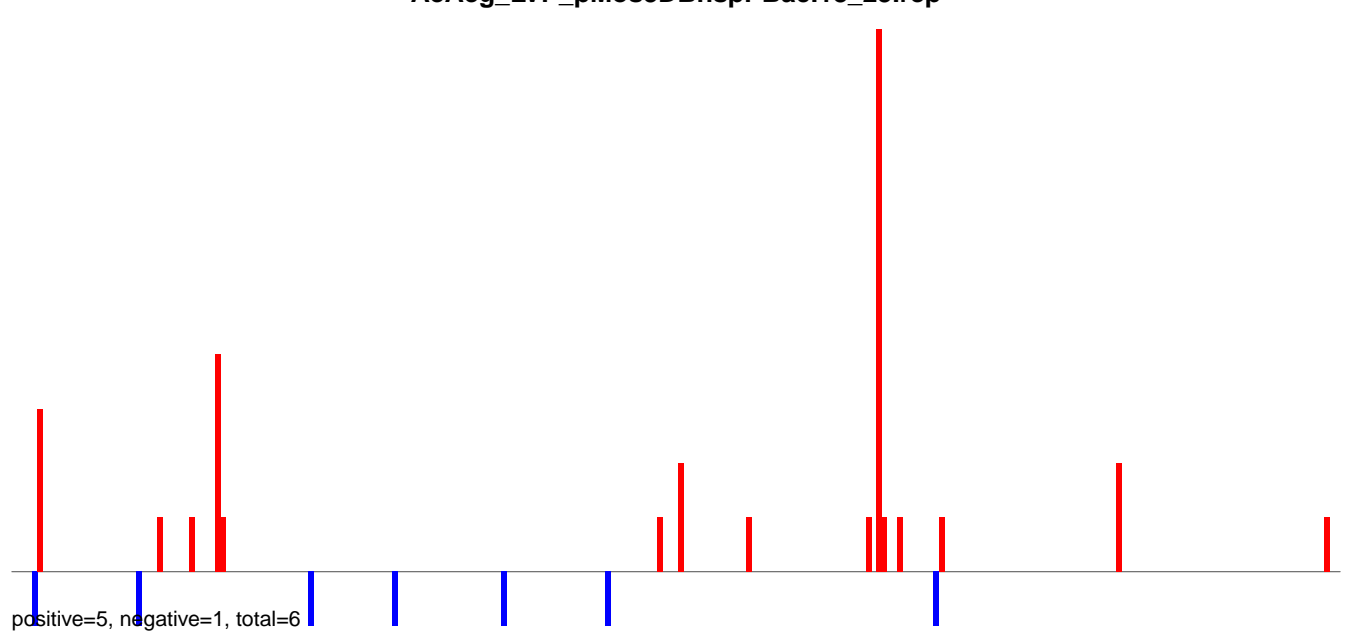
AeAeg_LVP_pMos3DBhspPBac.rep



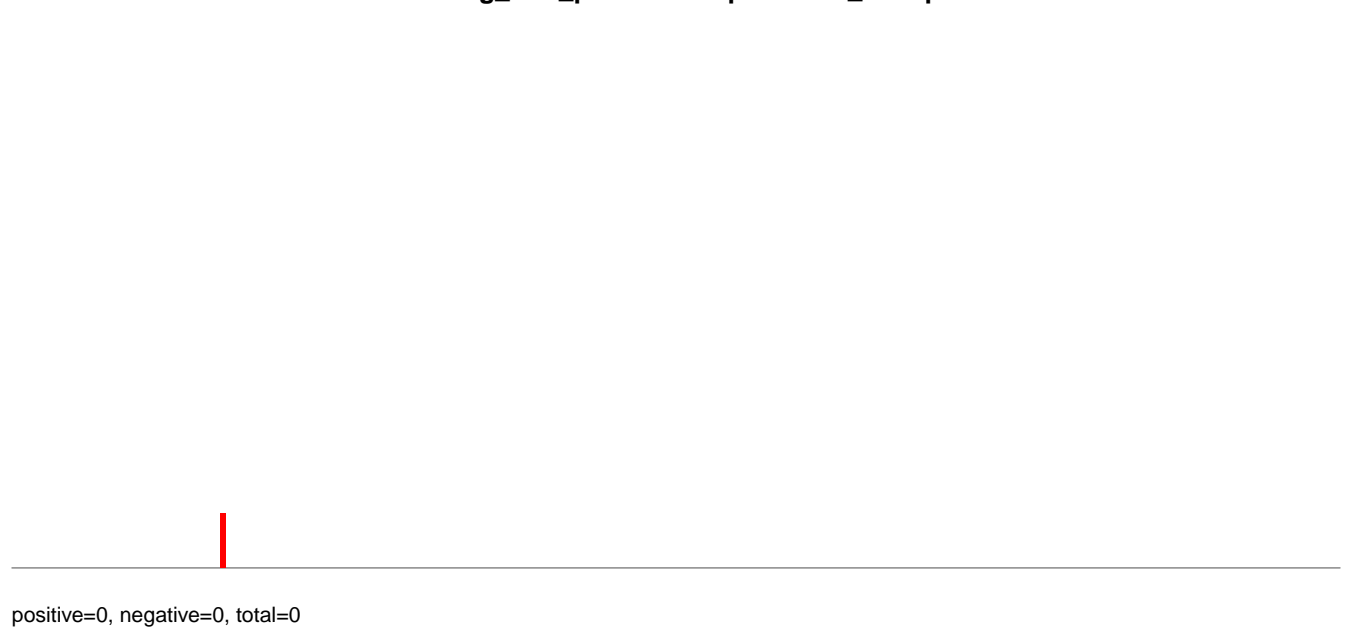
Window size=25, length=4176, virus@X74945.1-Aedes_albopictus_densovirus_2_DNA:1-4176

0 1000 2000 3000 4000

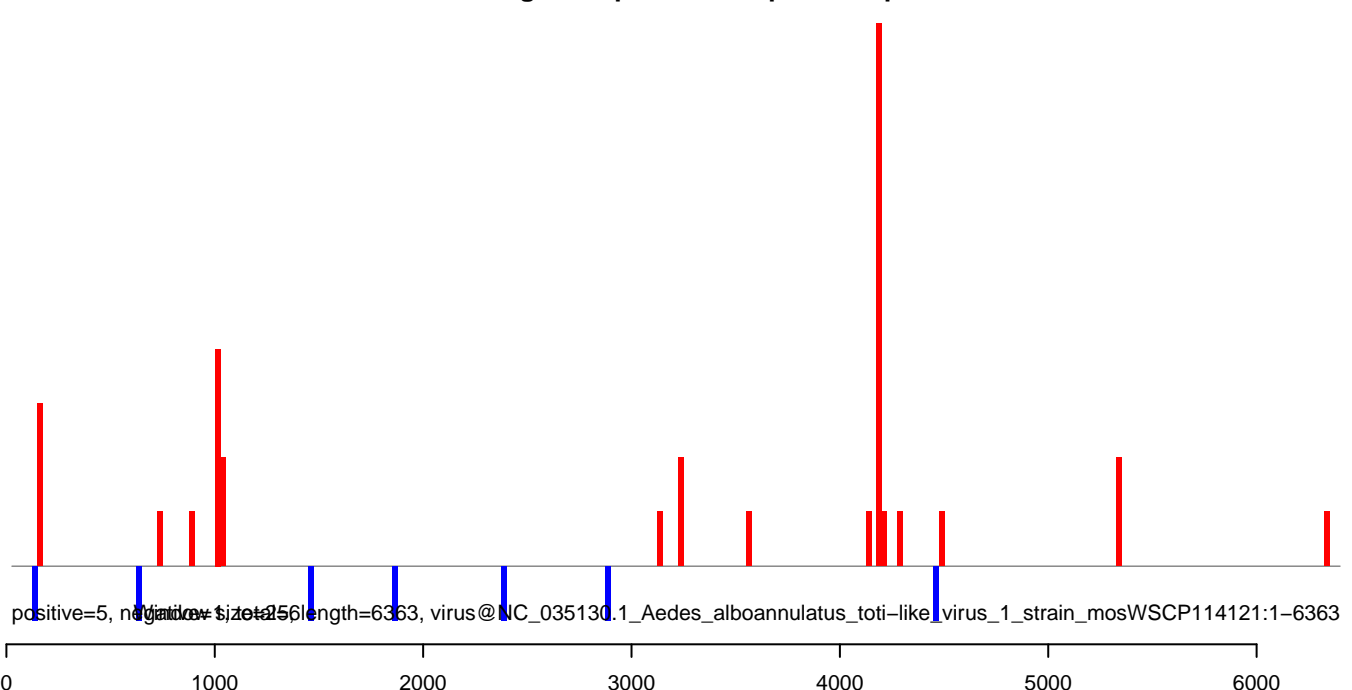
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



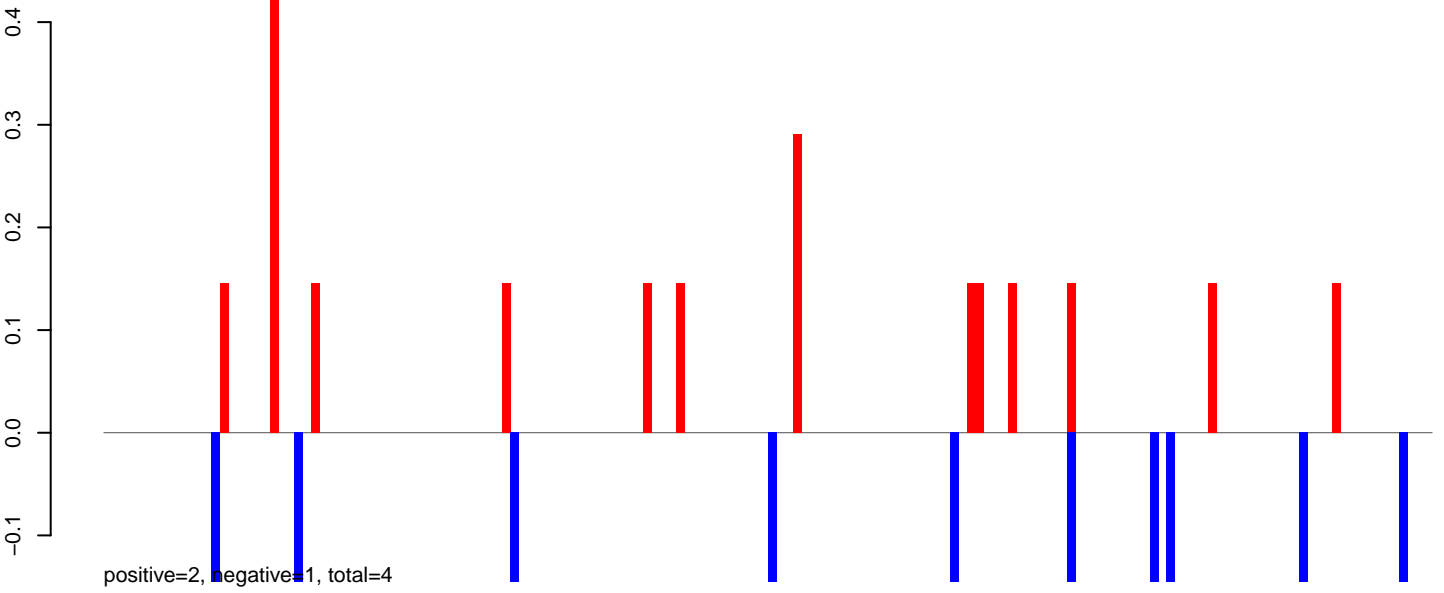
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



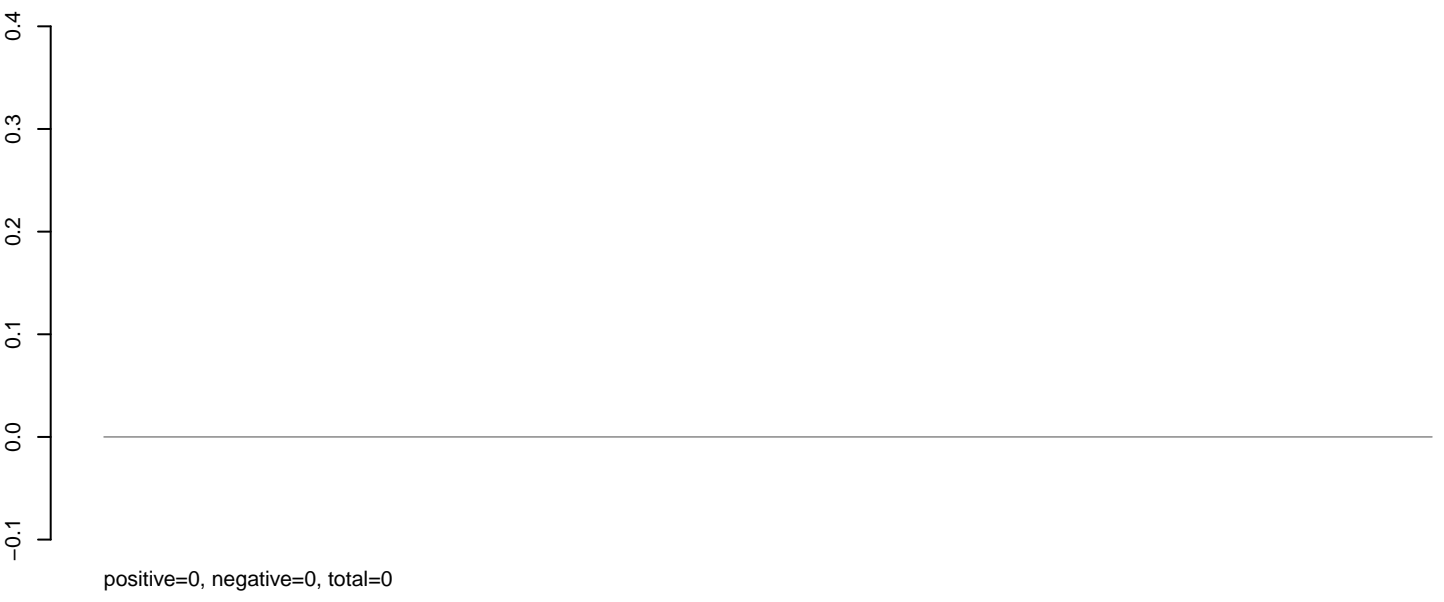
AeAeg_LVP_pMos3DBhspPBac.rep



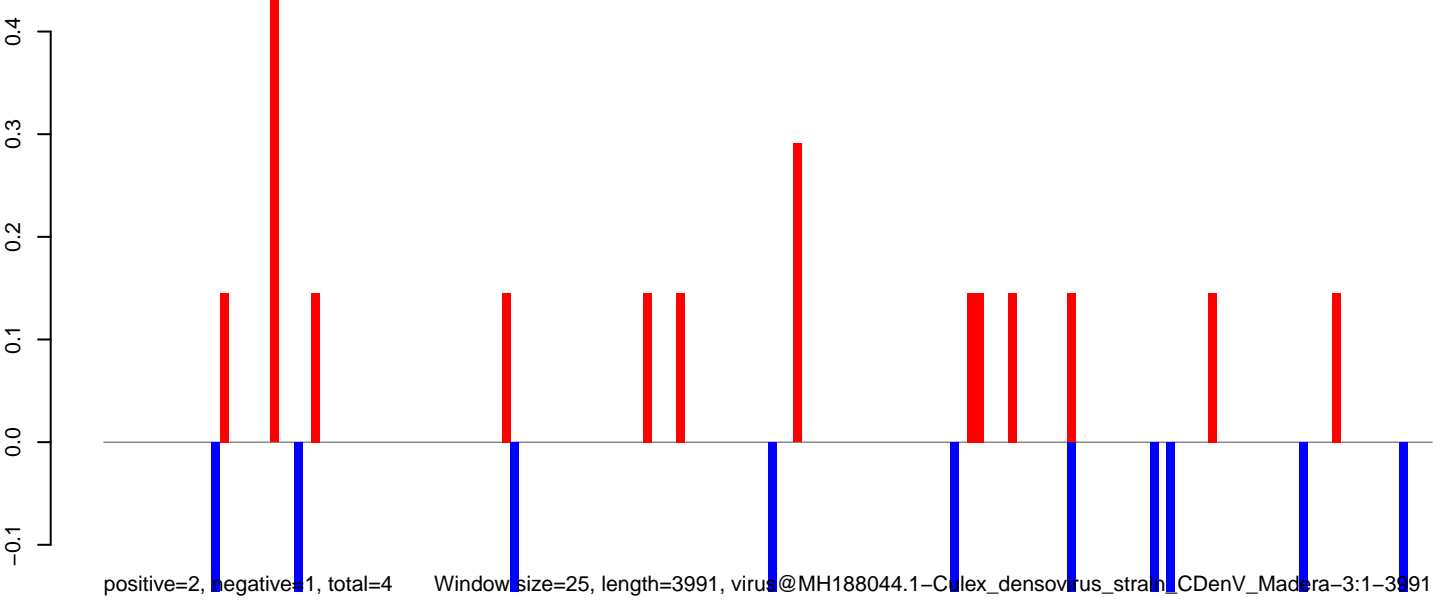
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



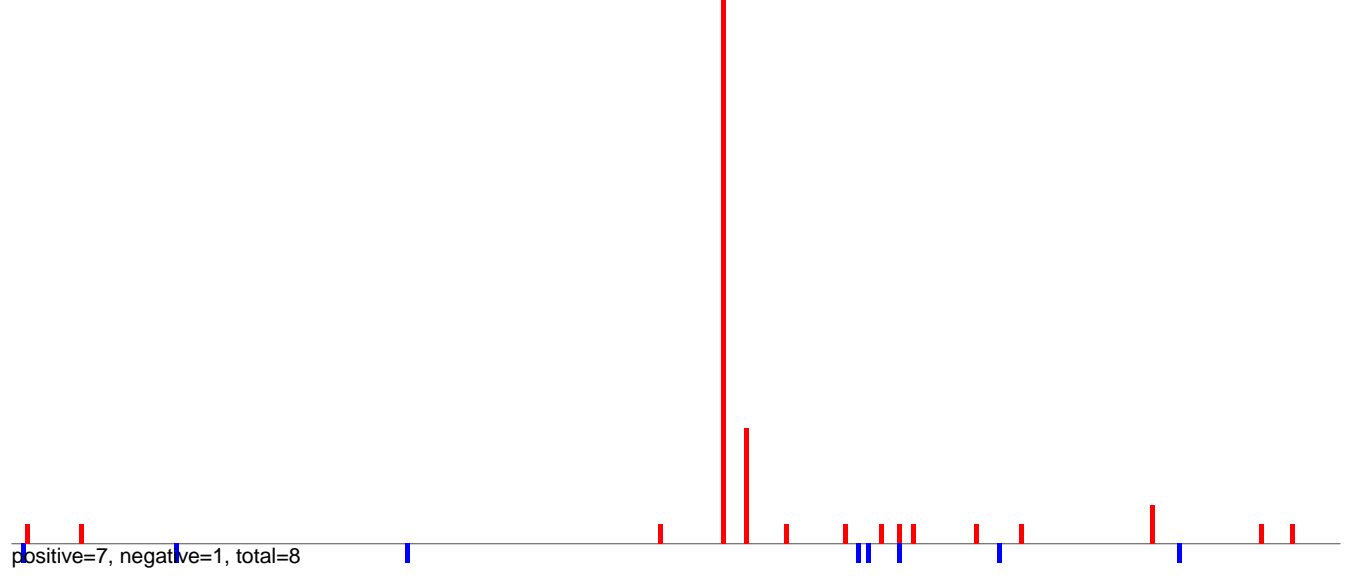
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



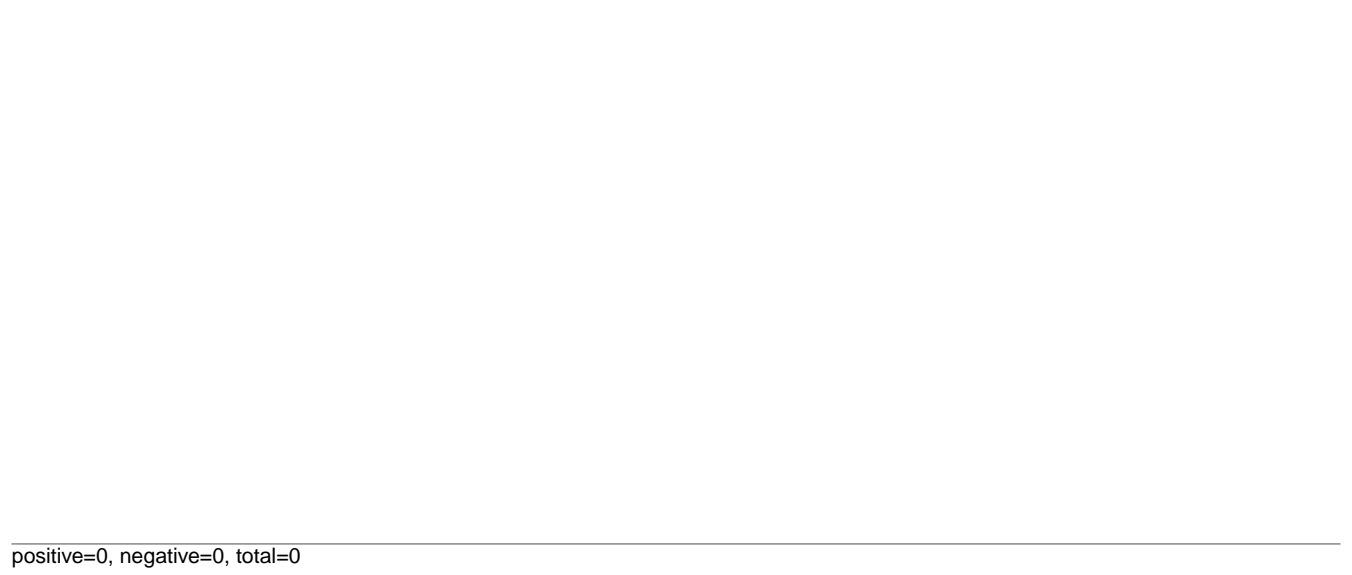
AeAeg_LVP_pMos3DBhspPBac.rep



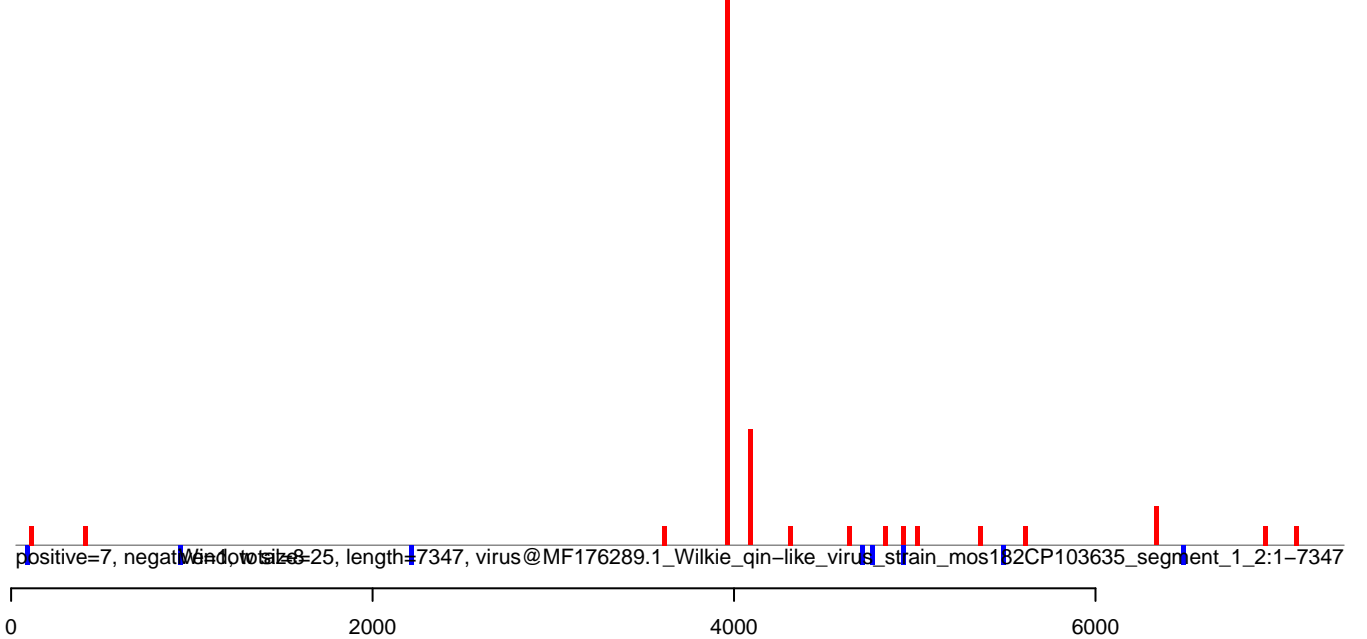
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



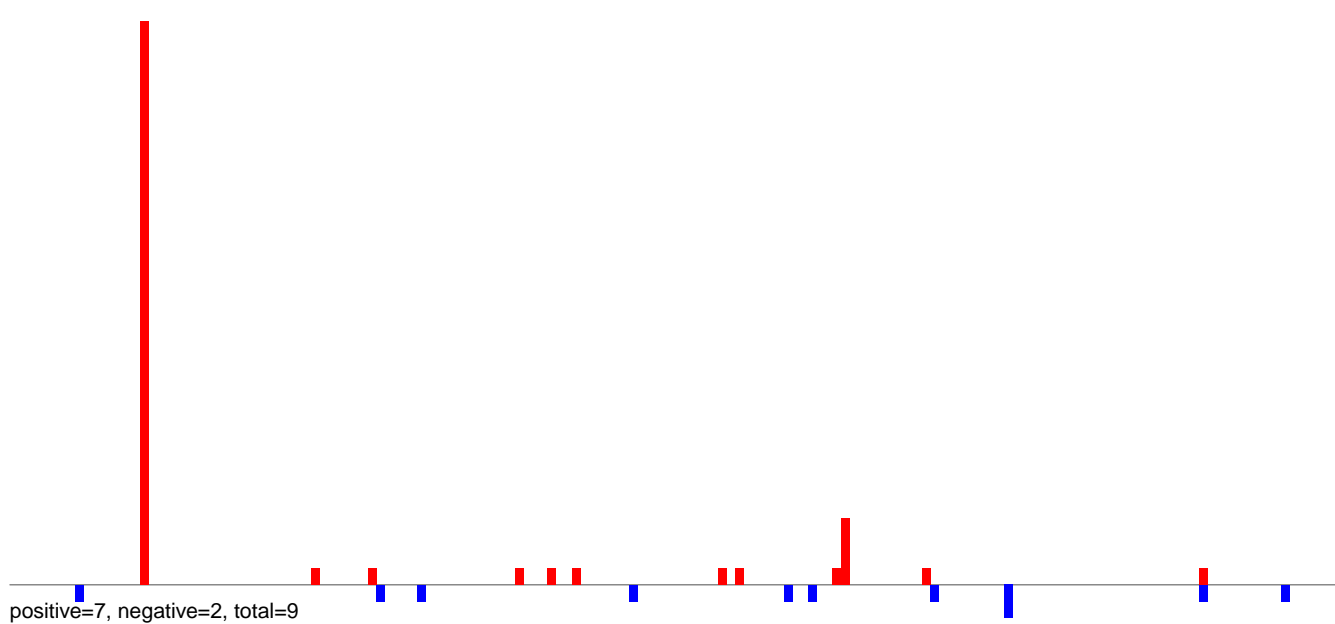
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



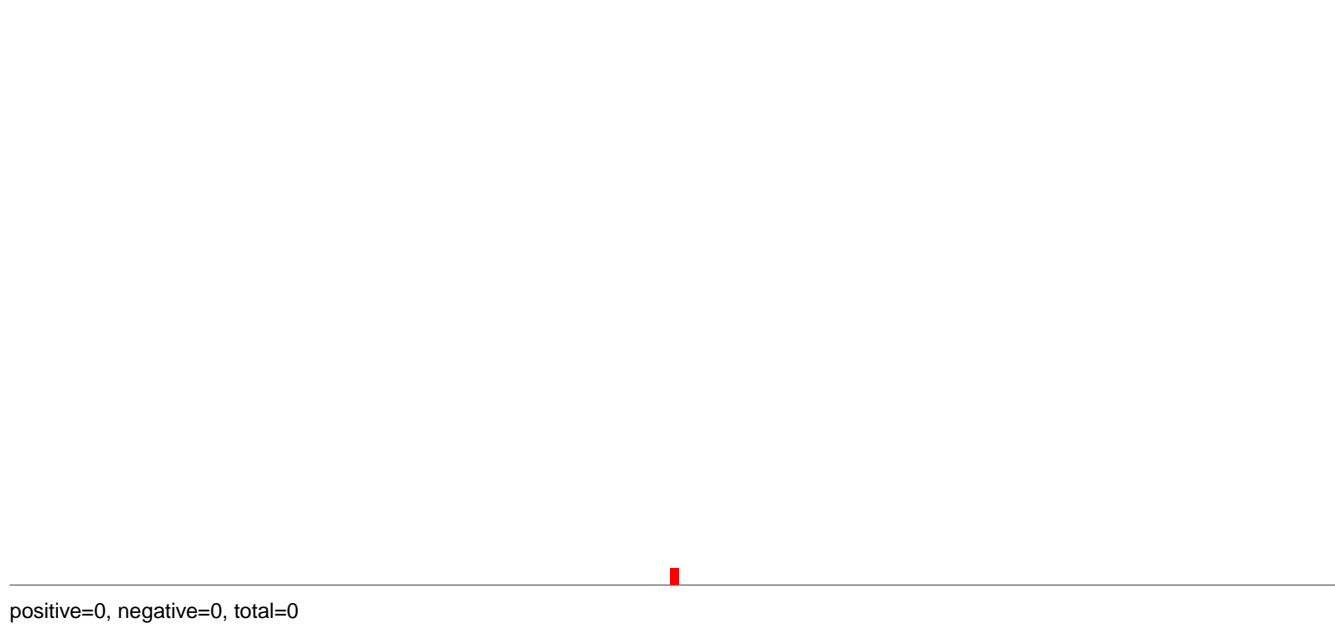
AeAeg_LVP_pMos3DBhspPBac.rep



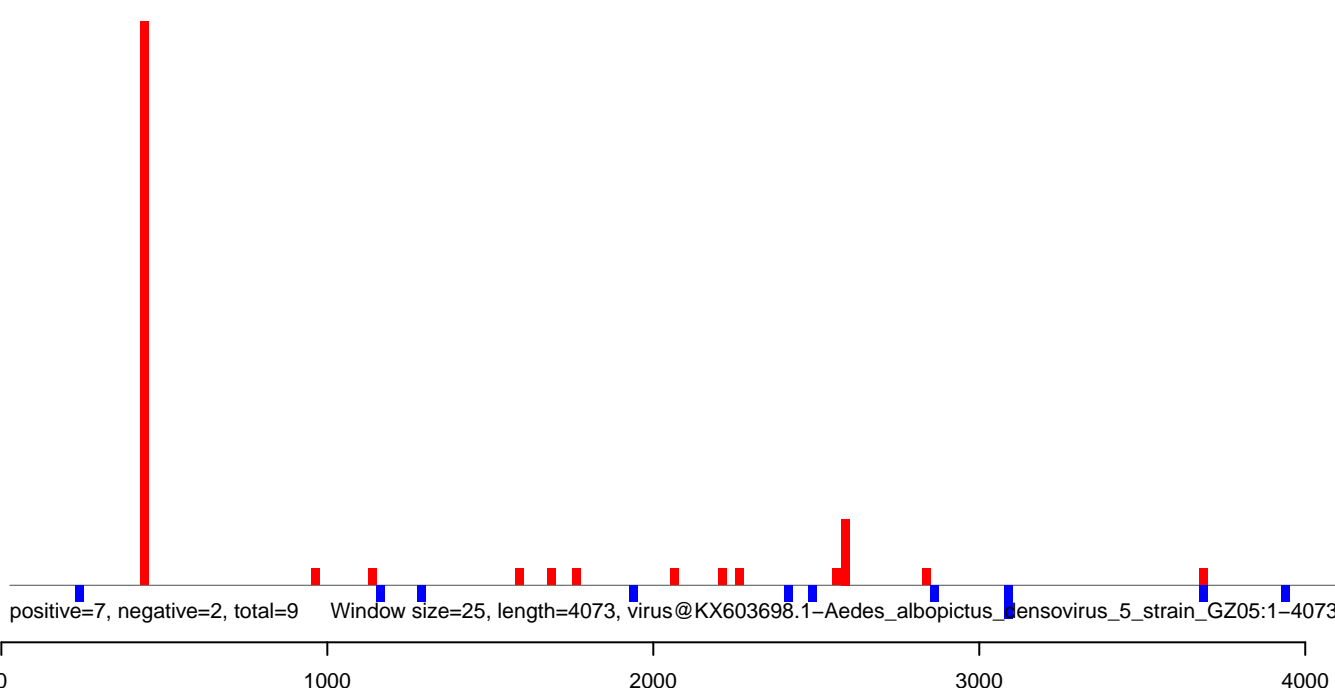
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



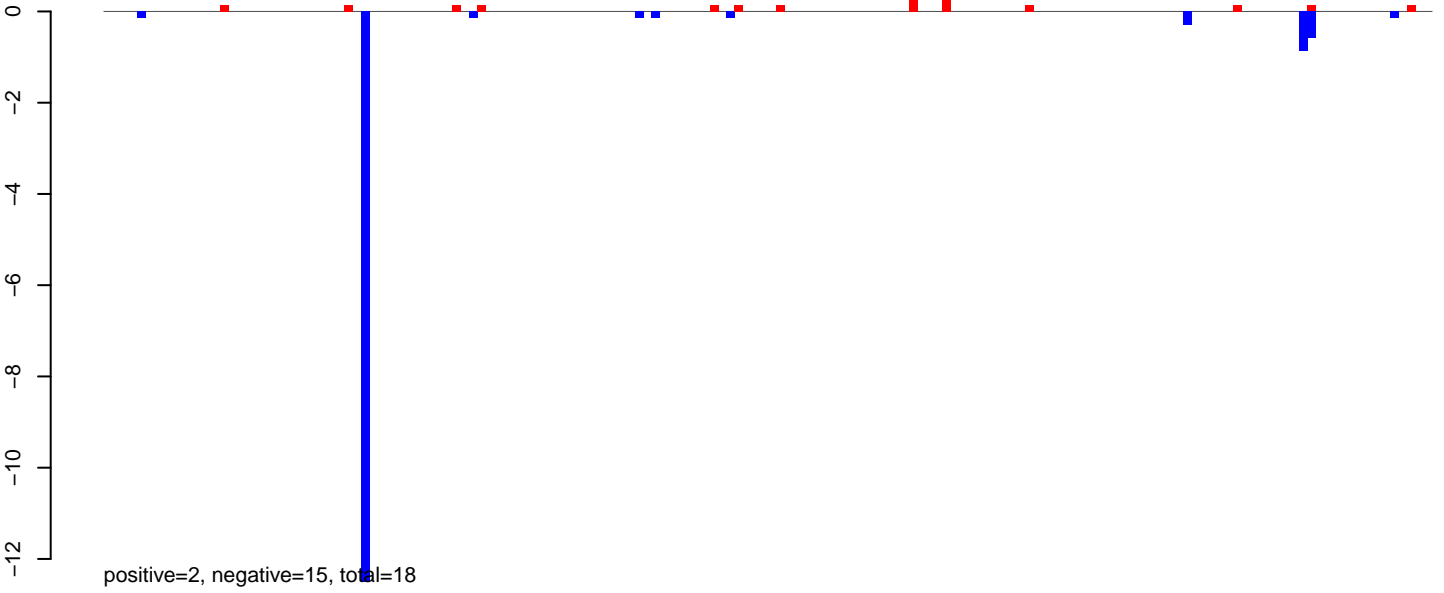
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



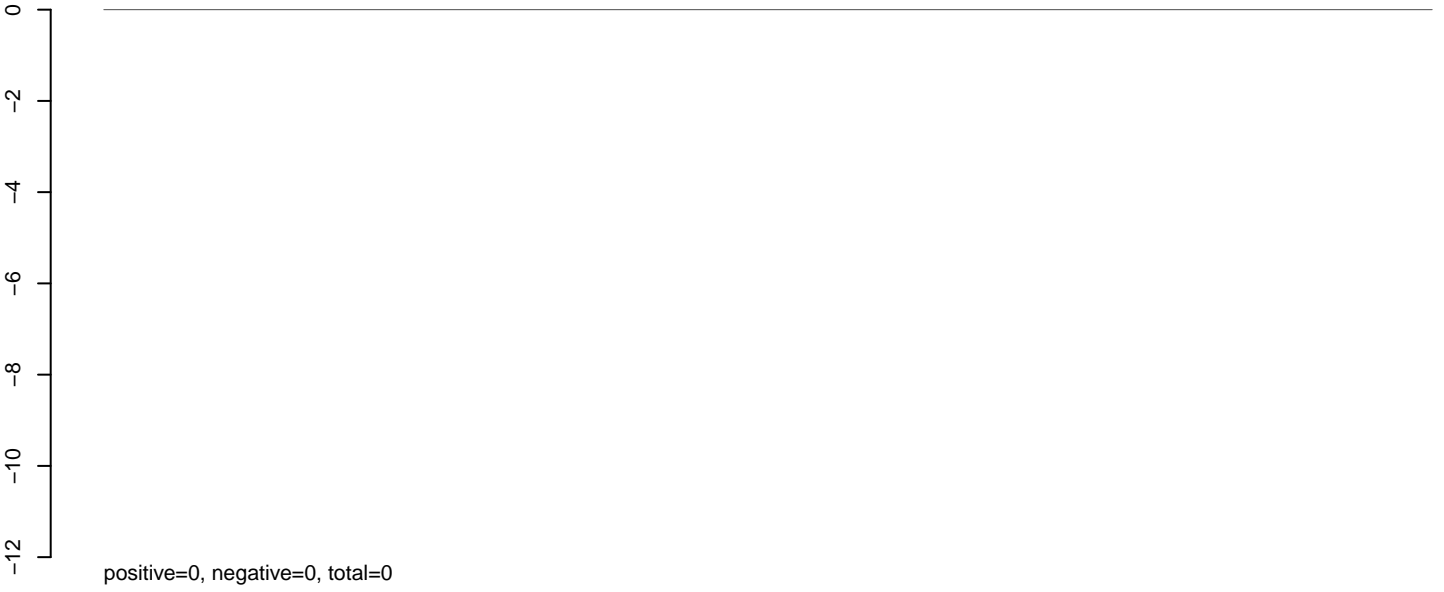
AeAeg_LVP_pMos3DBhspPBac.rep



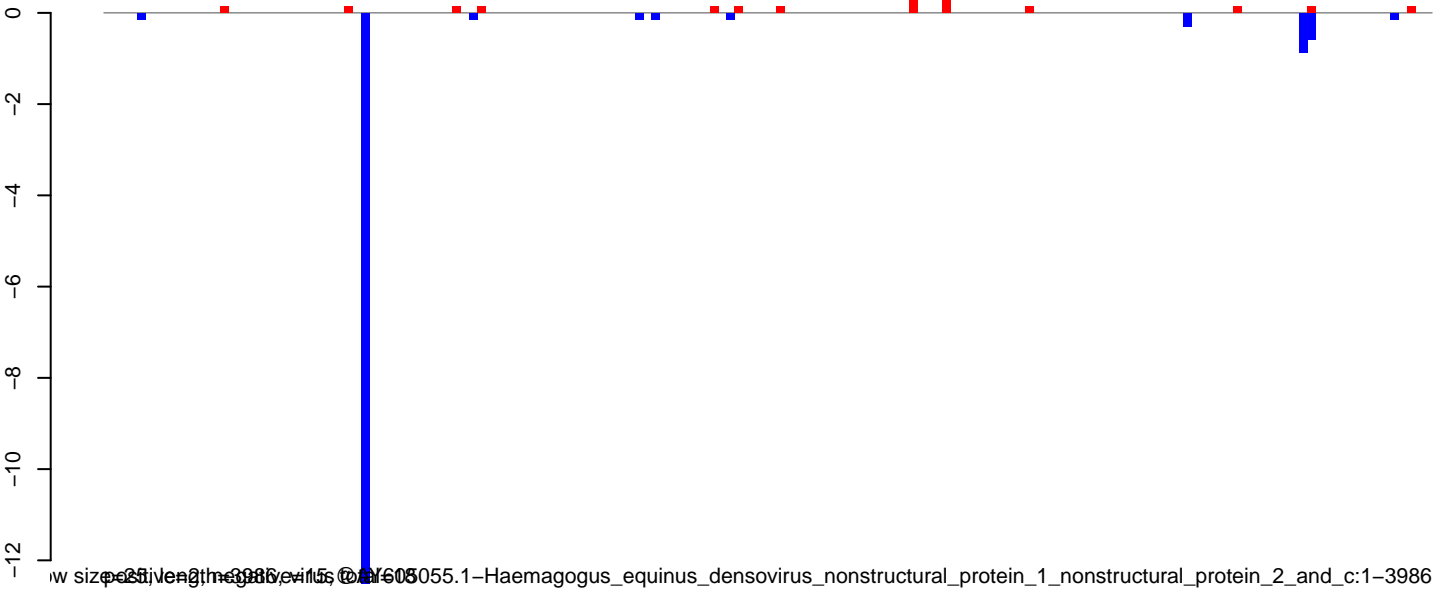
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



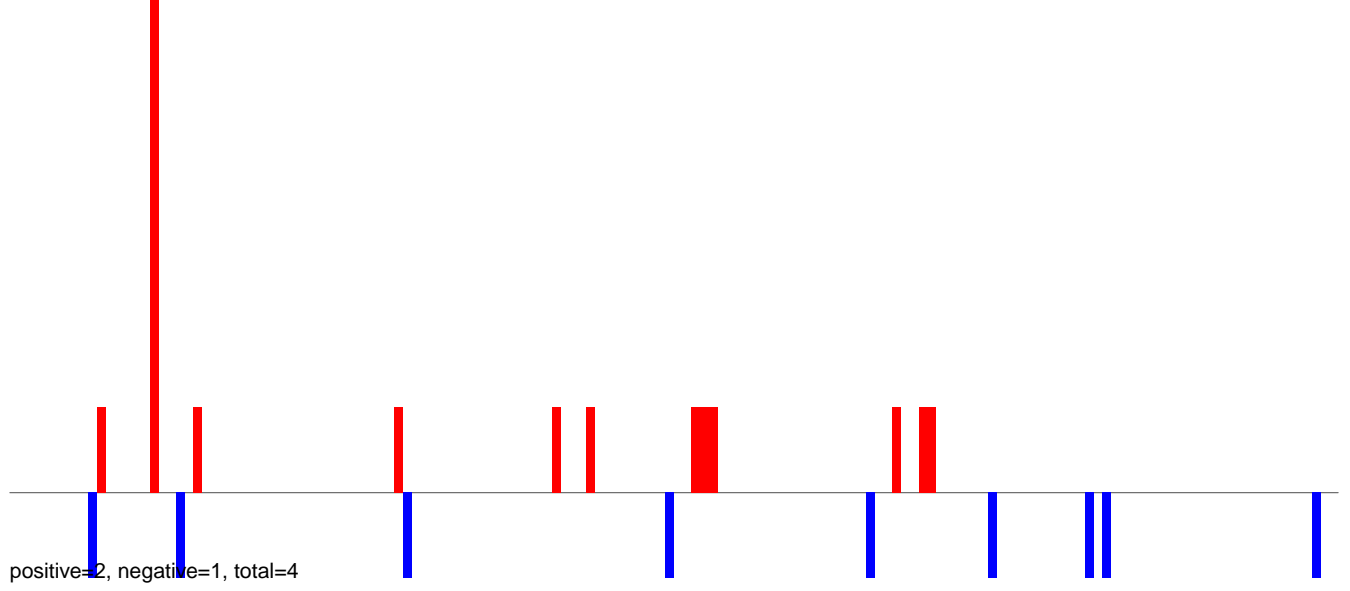
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



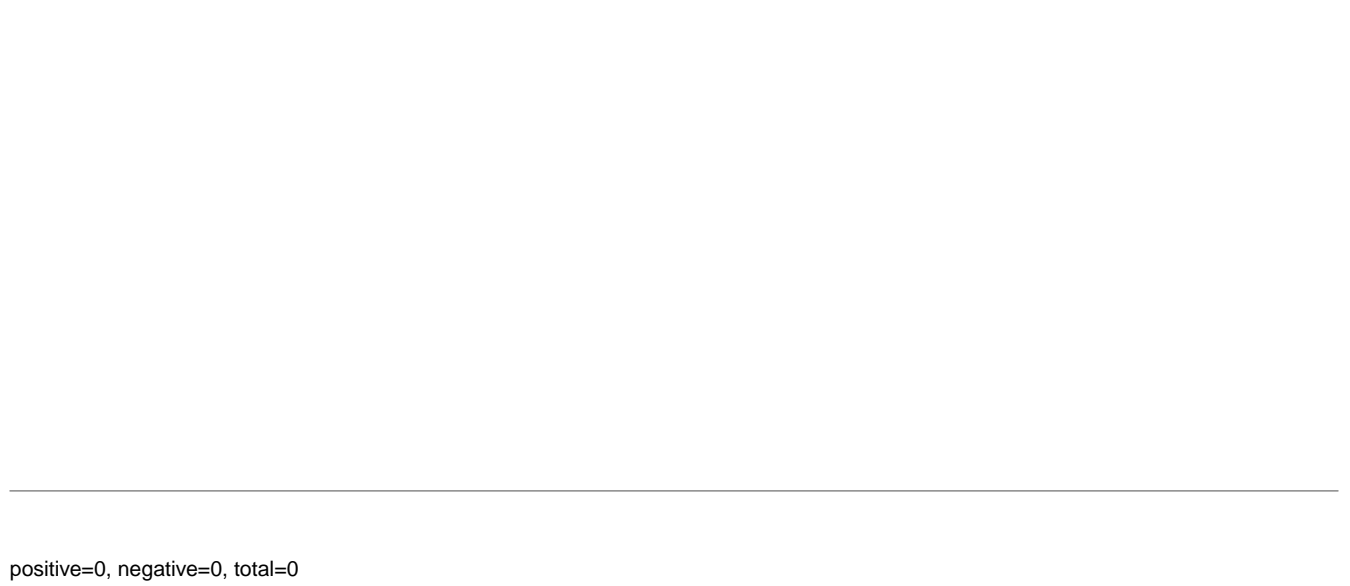
AeAeg_LVP_pMos3DBhspPBac.rep



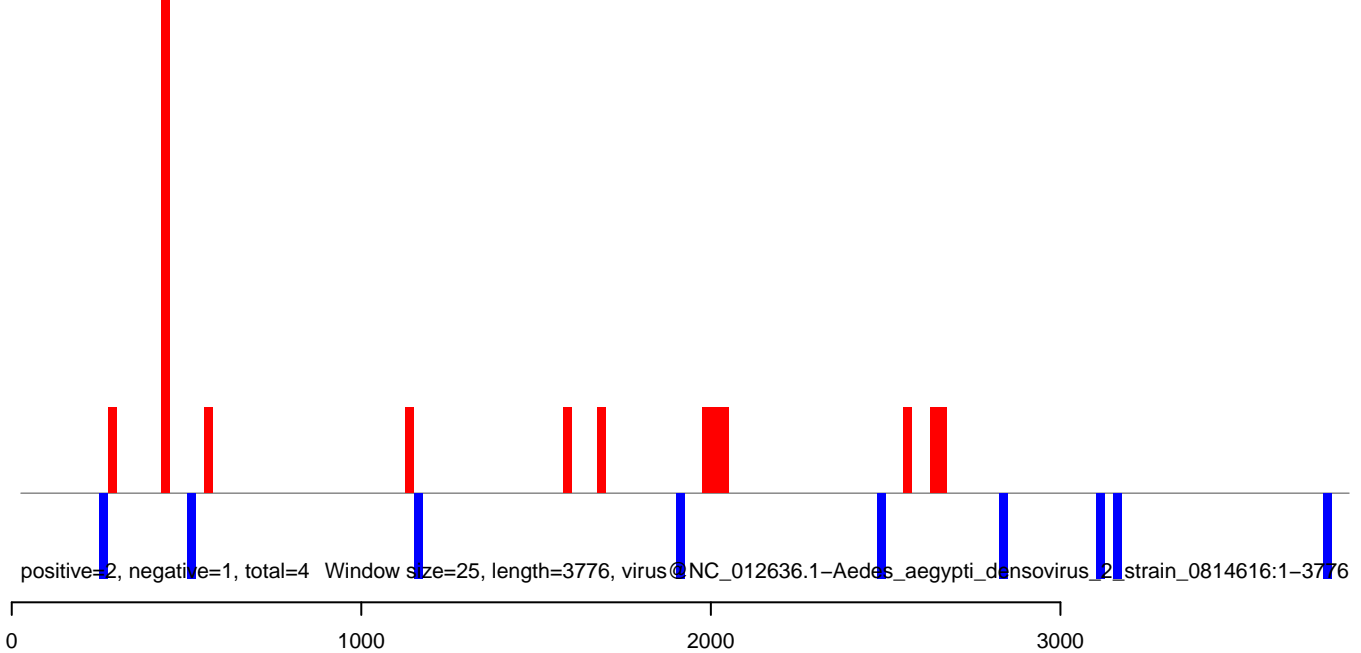
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



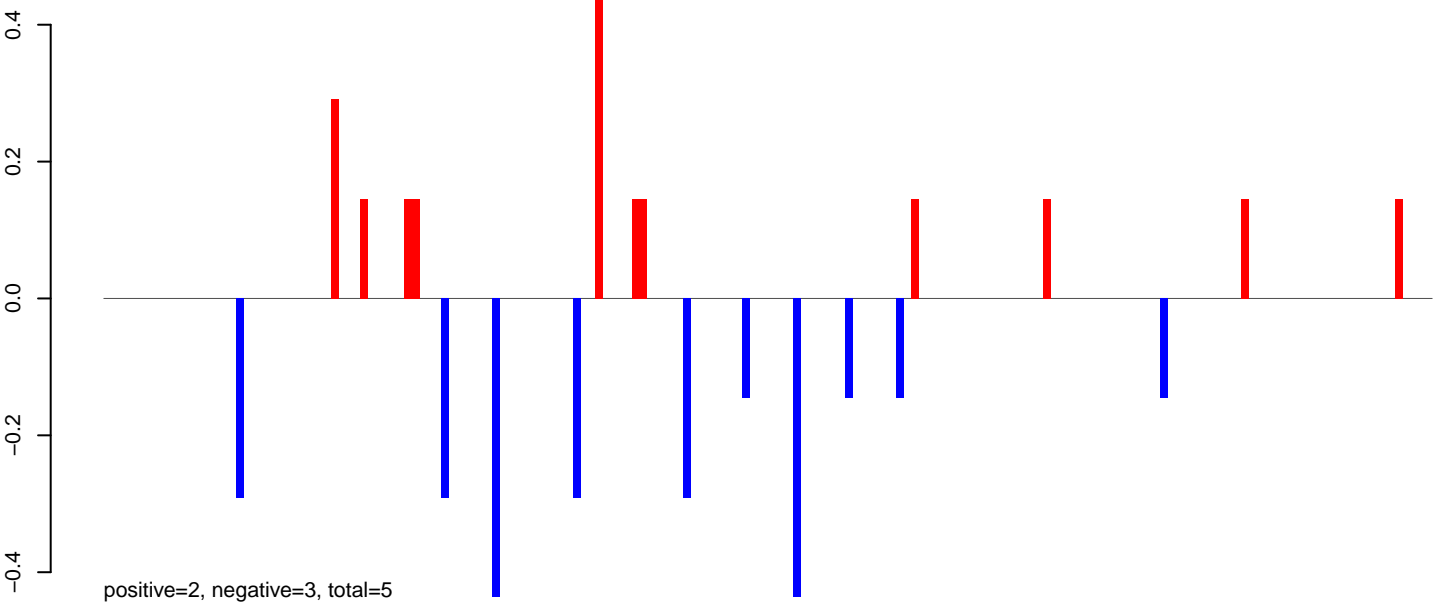
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



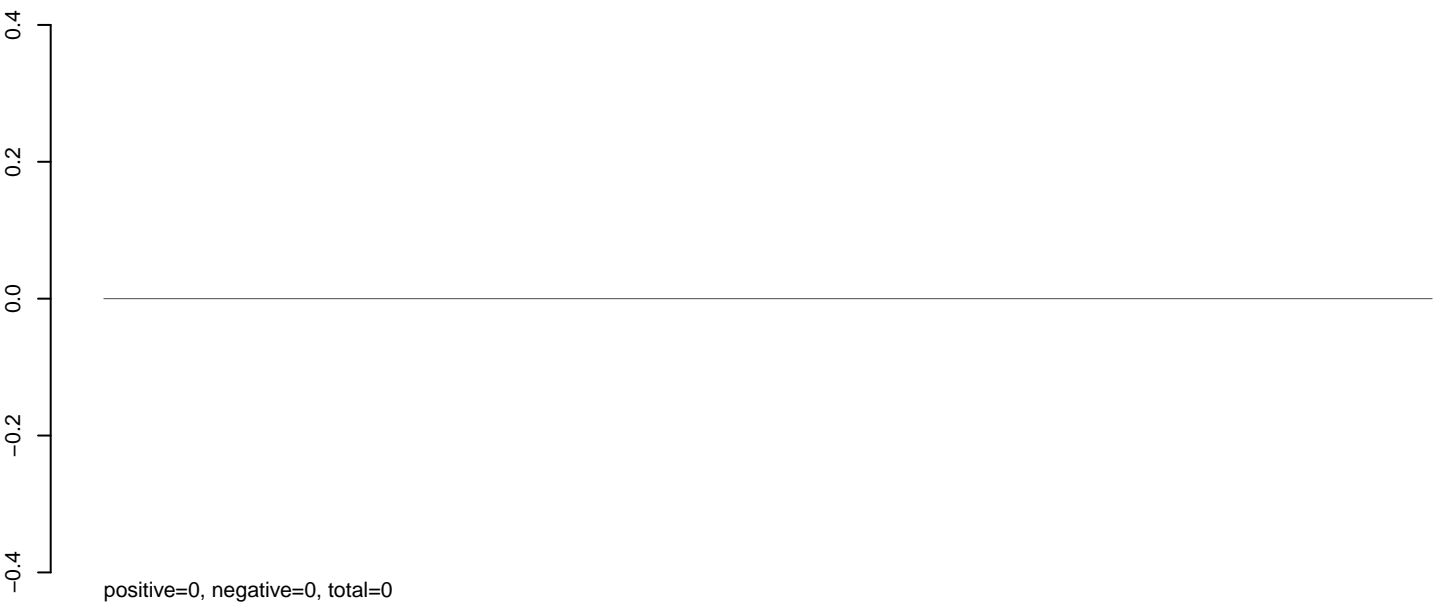
AeAeg_LVP_pMos3DBhspPBac.rep



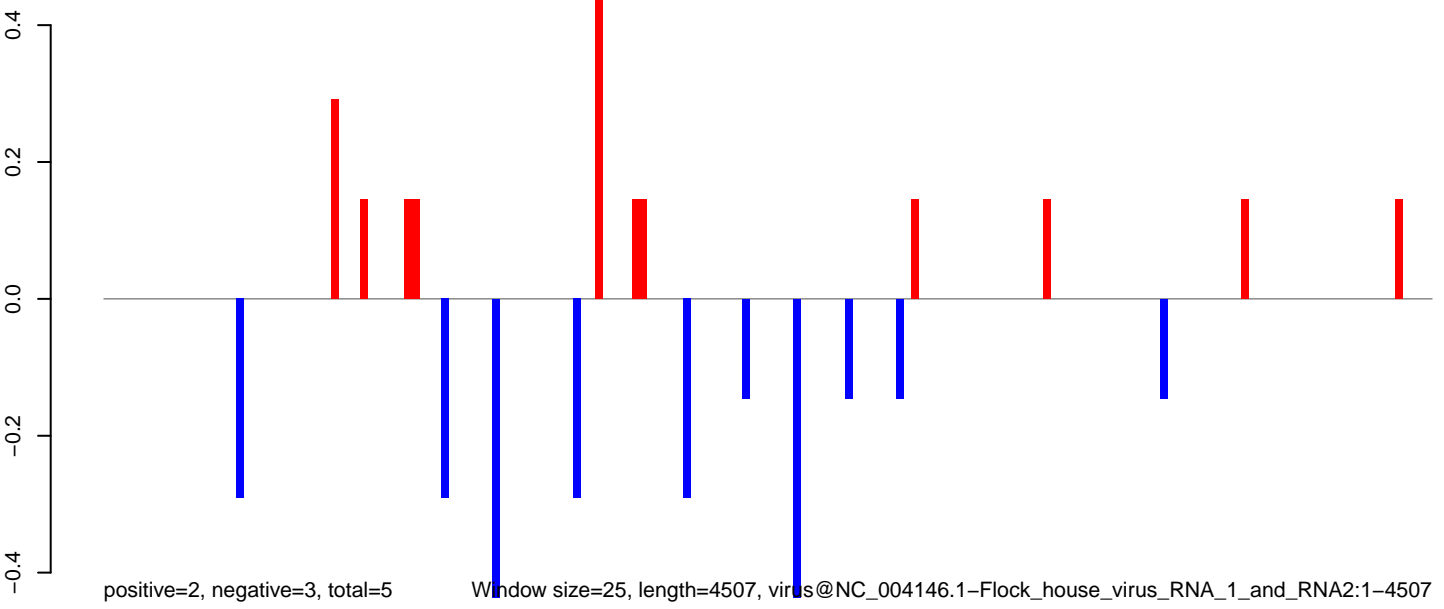
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

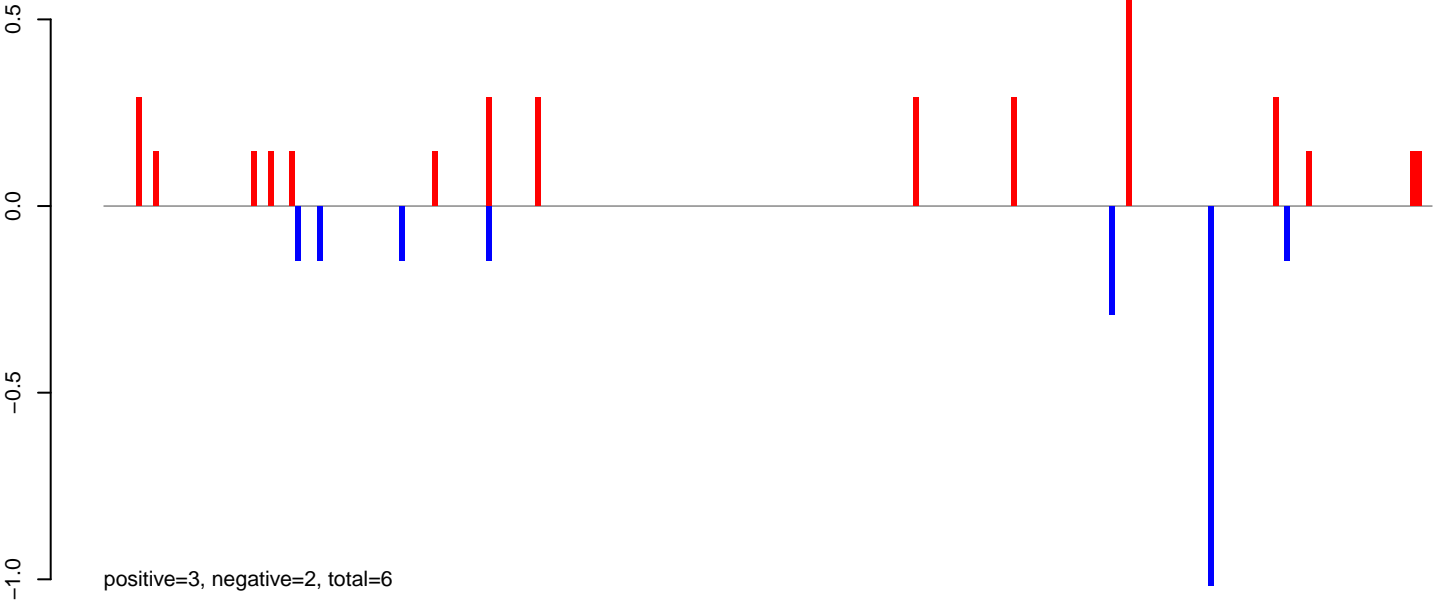


AeAeg_LVP_pMos3DBhspPBac.rep

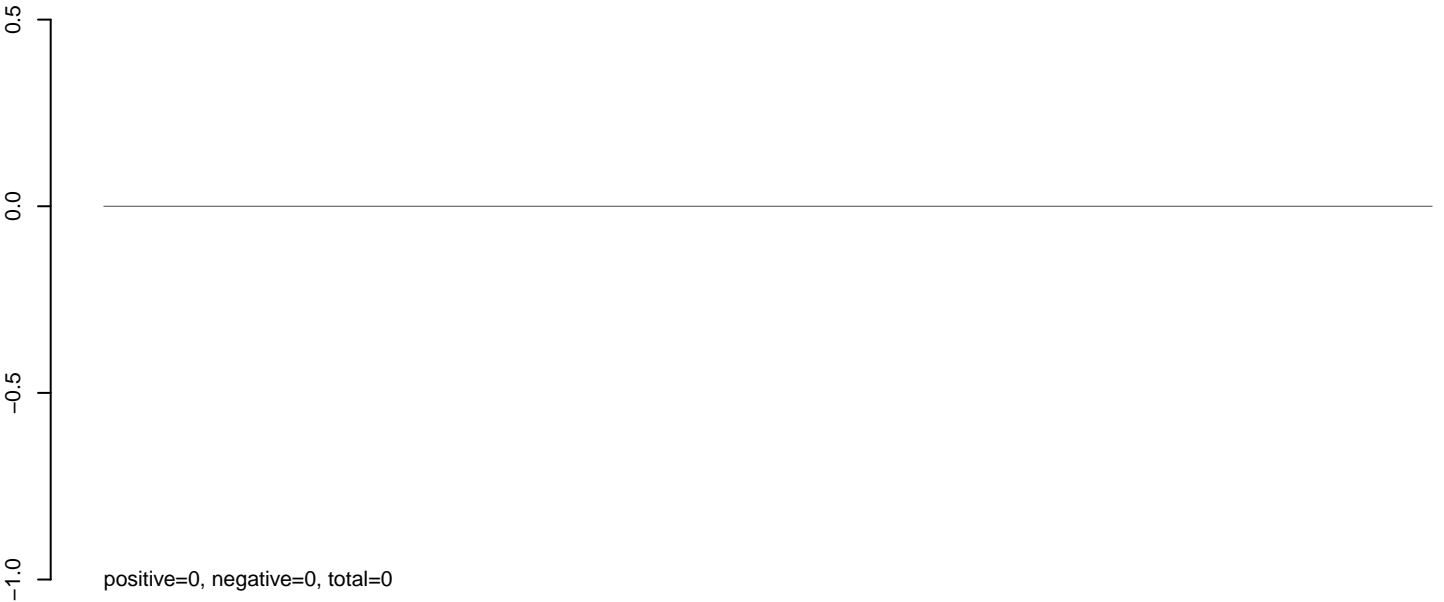


Window size=25, length=4507, virus@NC_004146.1-Flock_house_virus_RNA_1_and_RNA2:1-4507

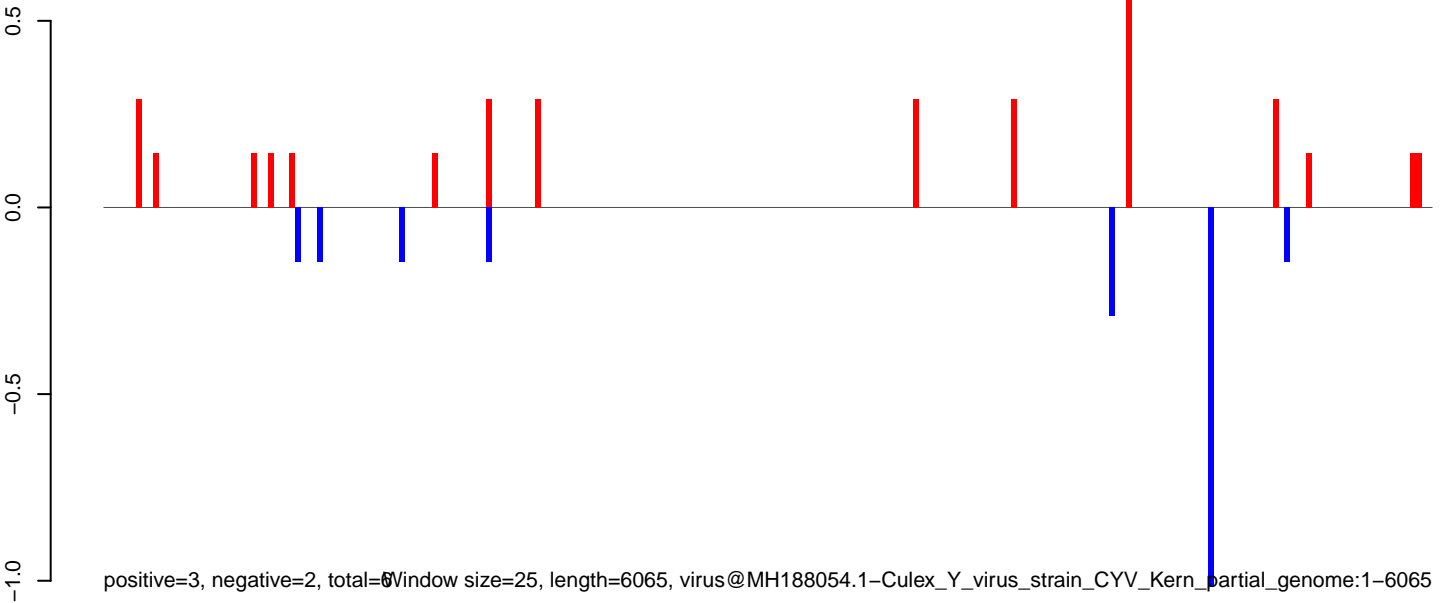
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



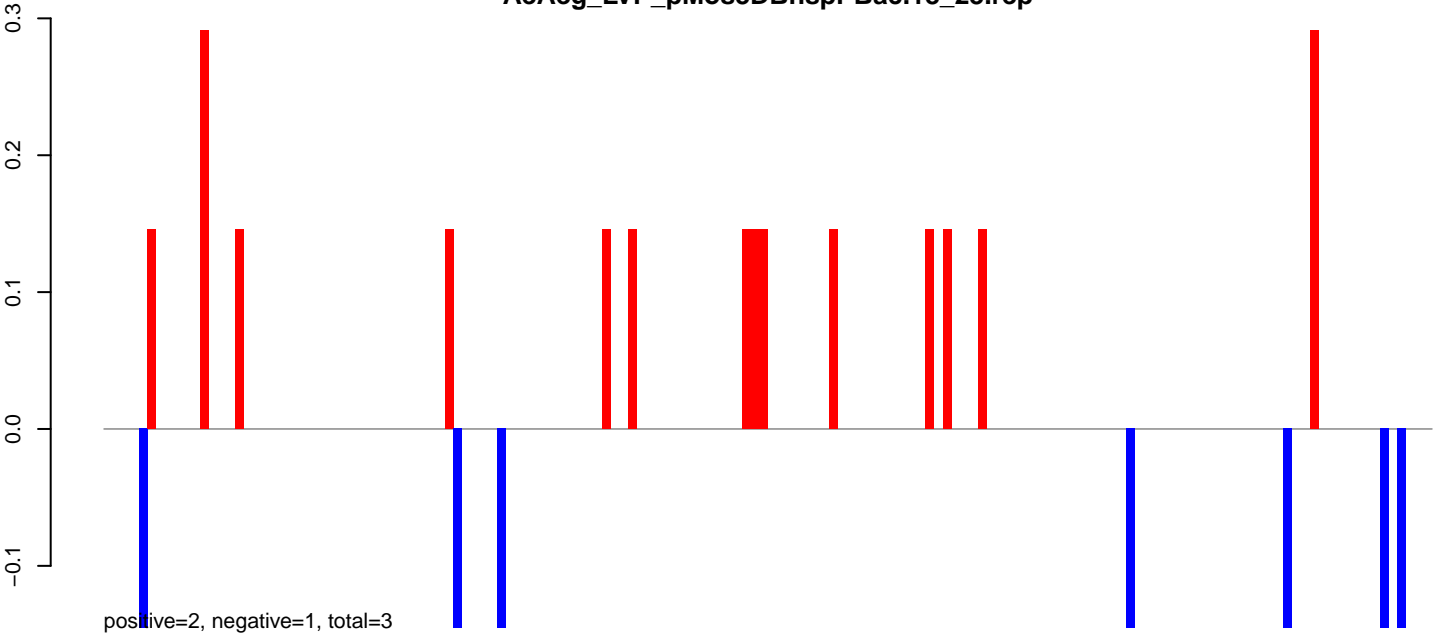
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



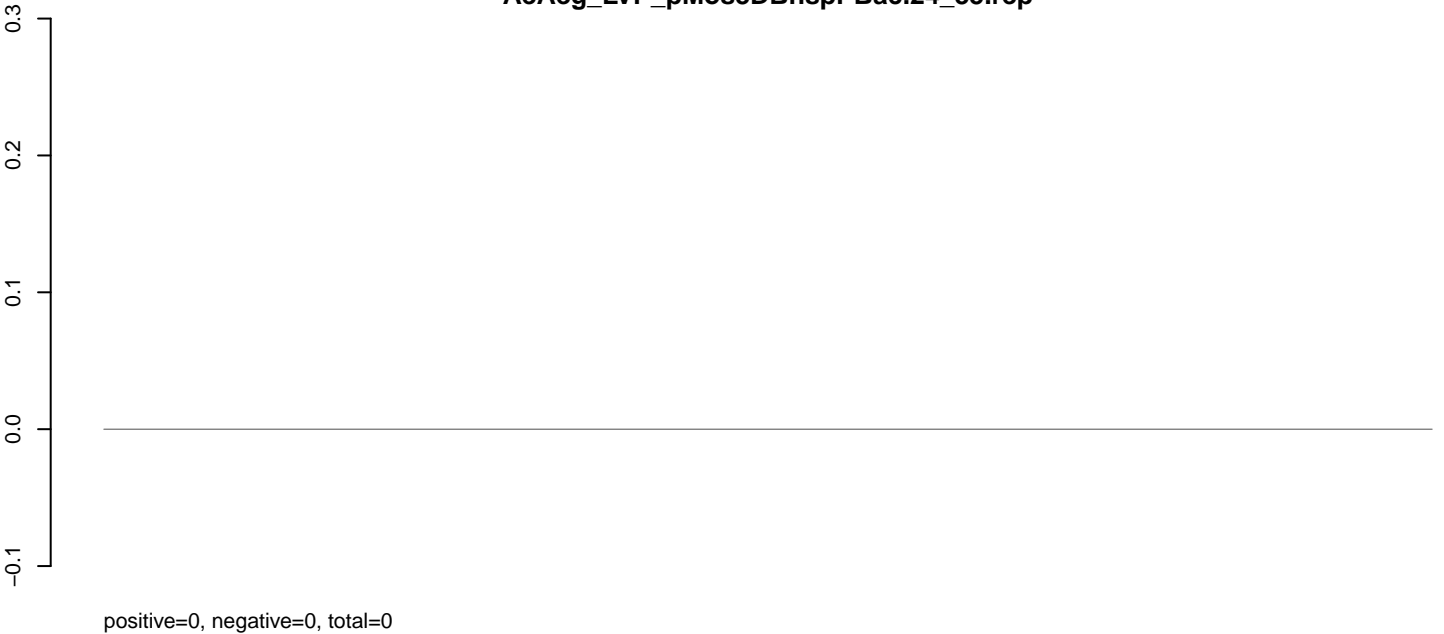
AeAeg_LVP_pMos3DBhspPBac.rep



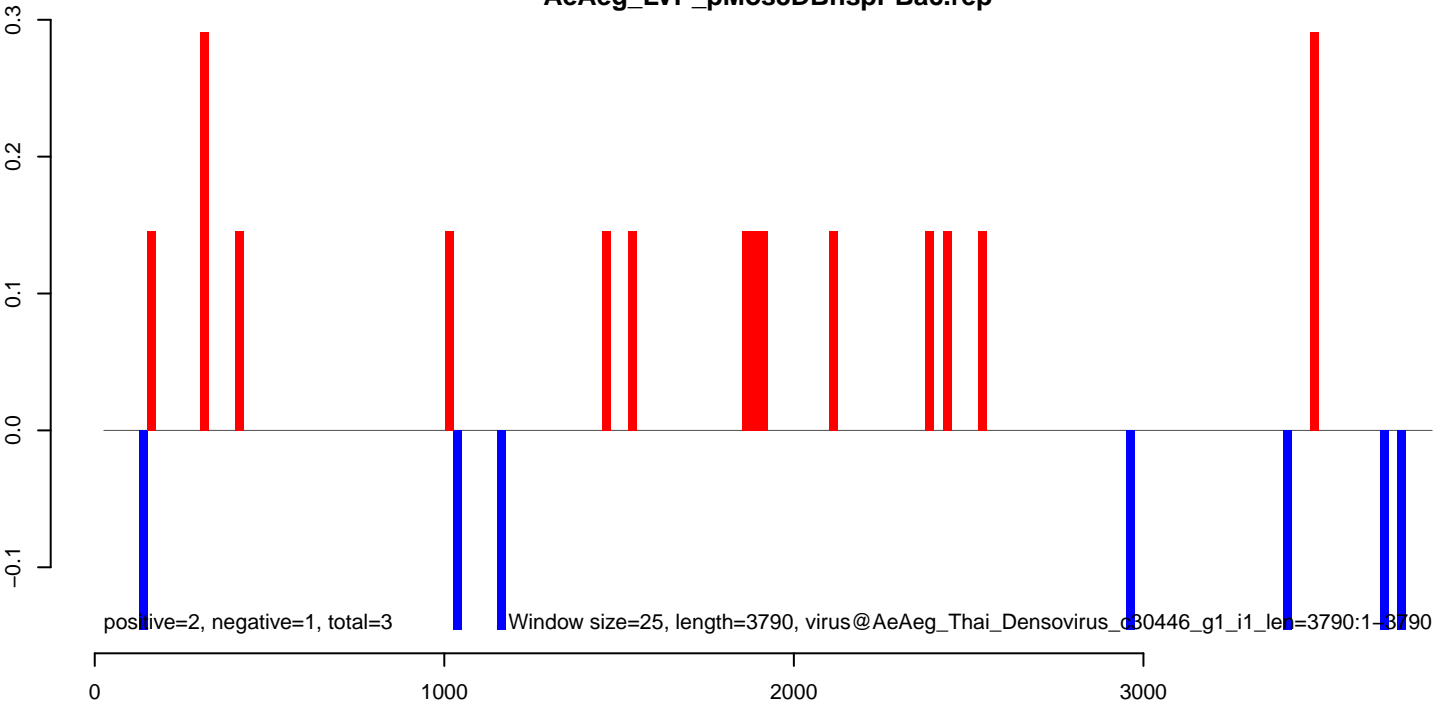
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



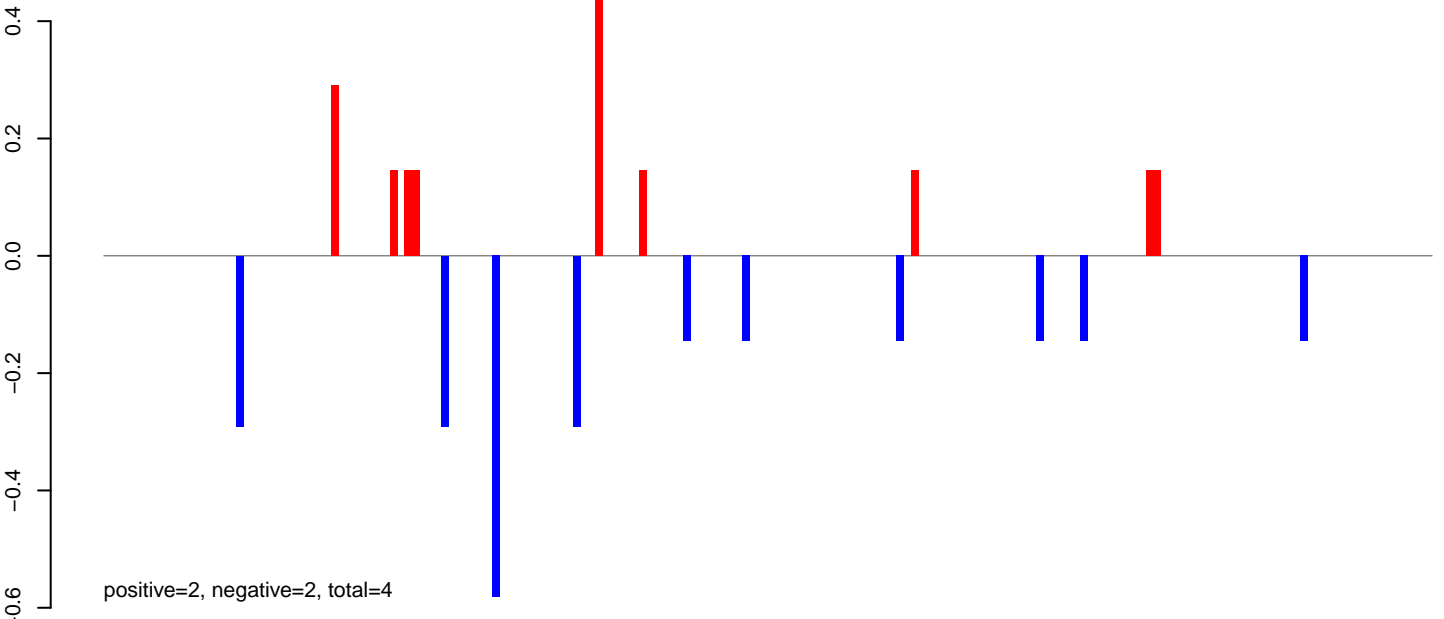
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



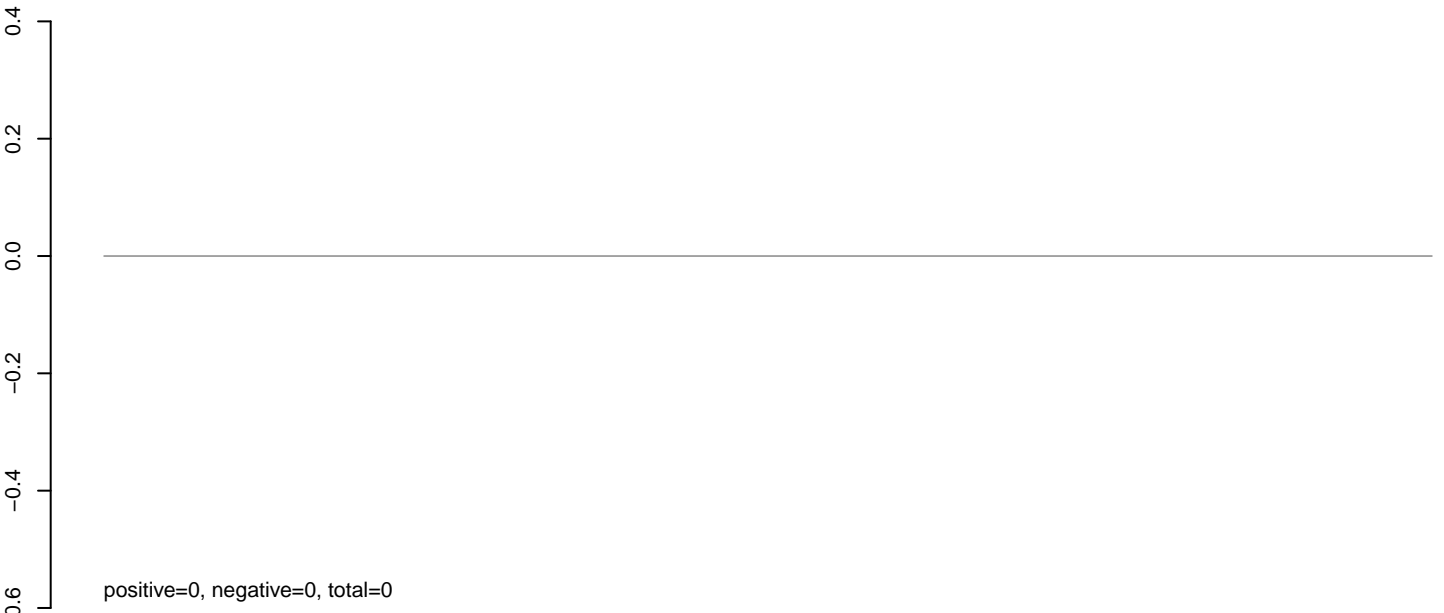
AeAeg_LVP_pMos3DBhspPBac.rep



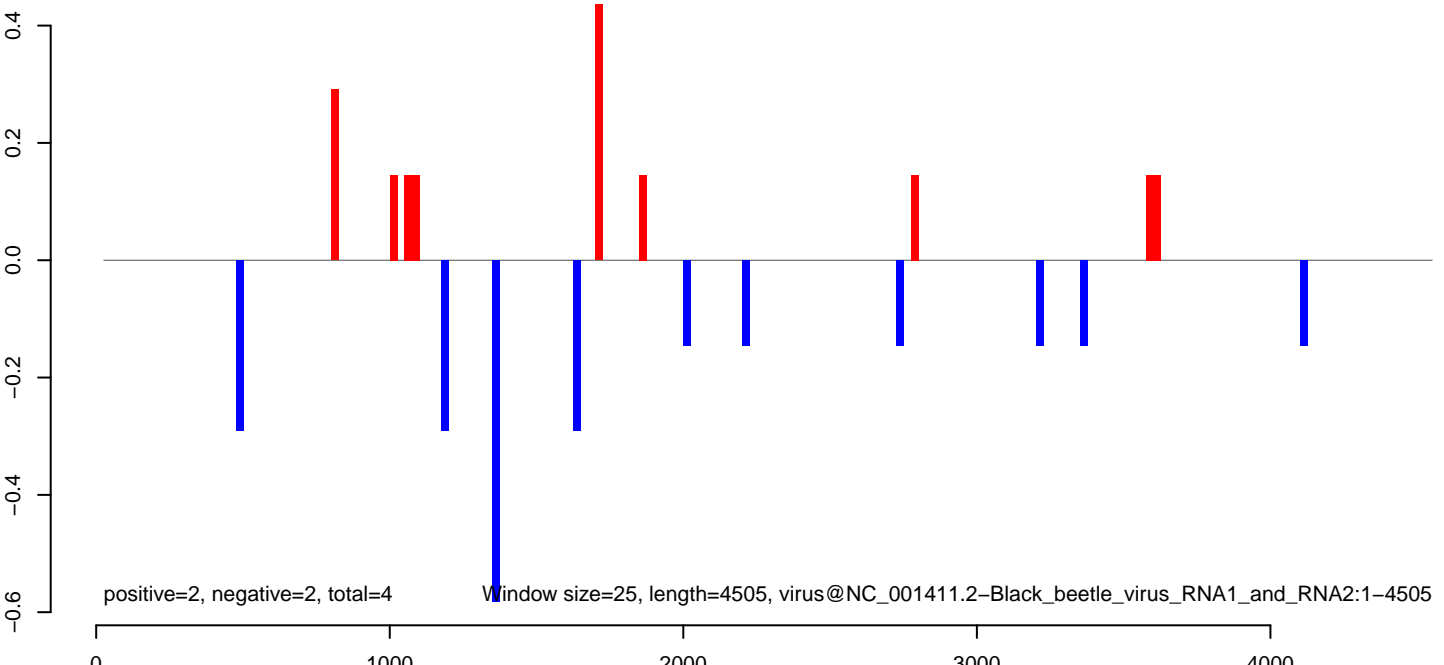
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



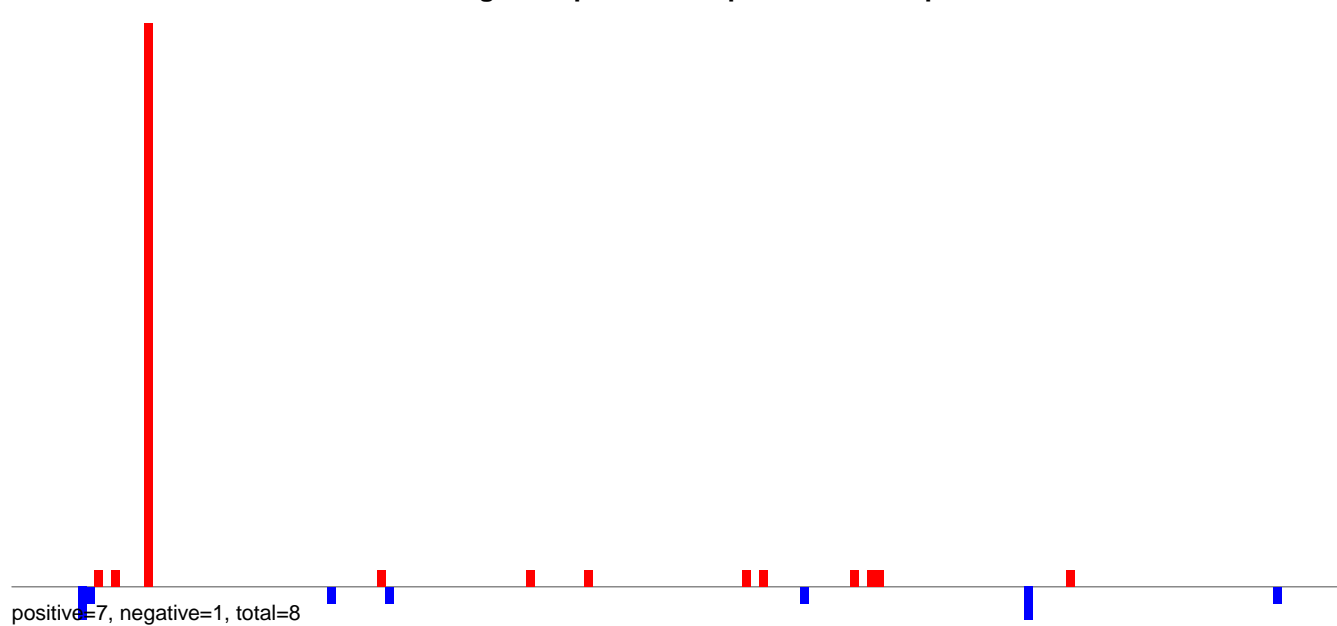
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



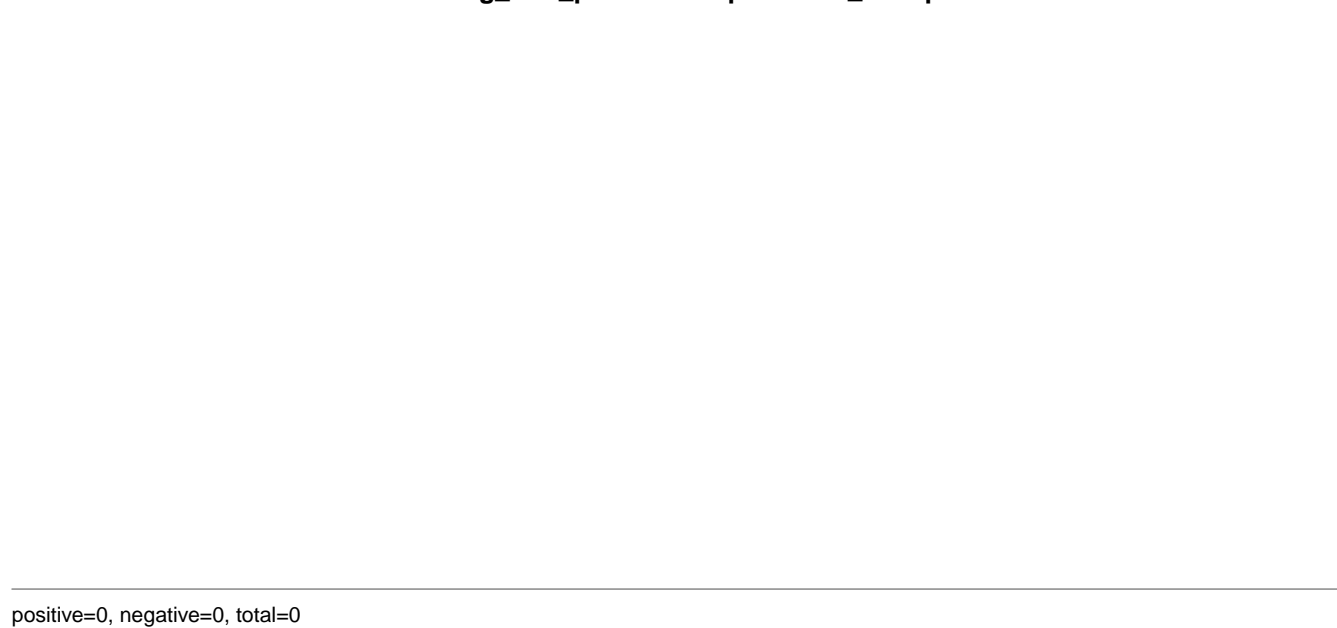
AeAeg_LVP_pMos3DBhspPBac.rep



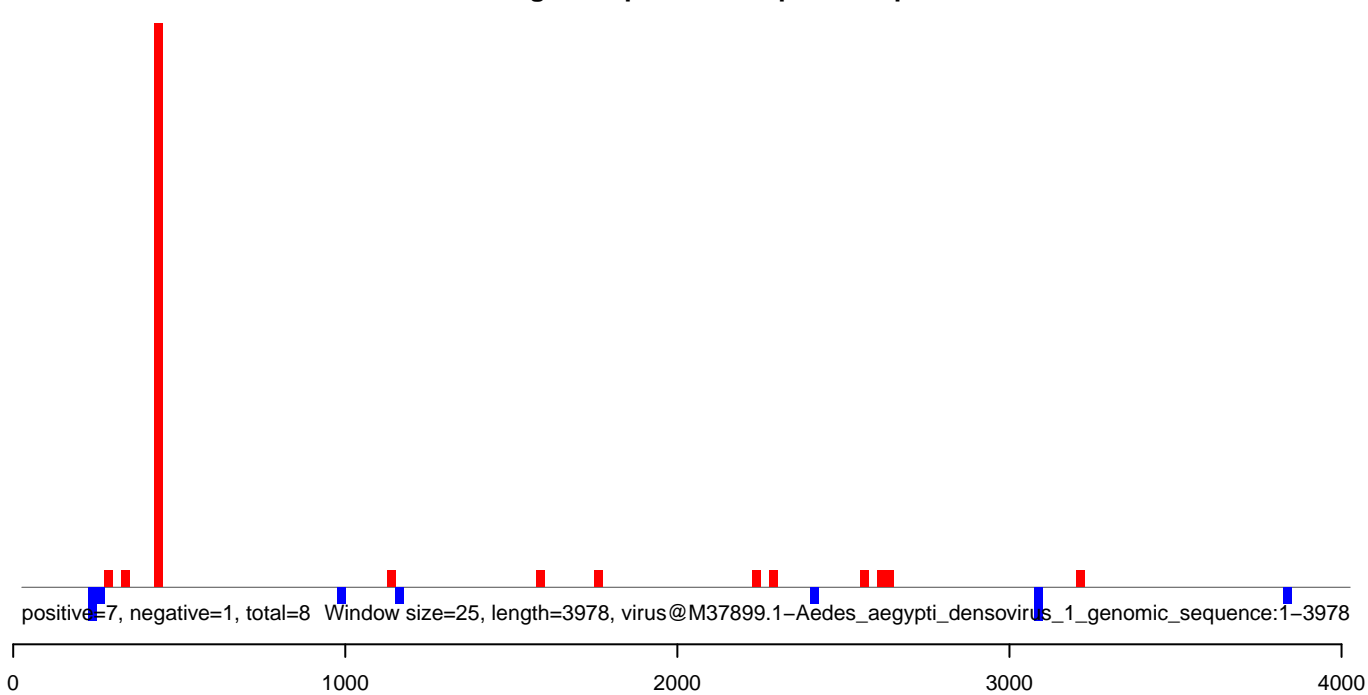
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



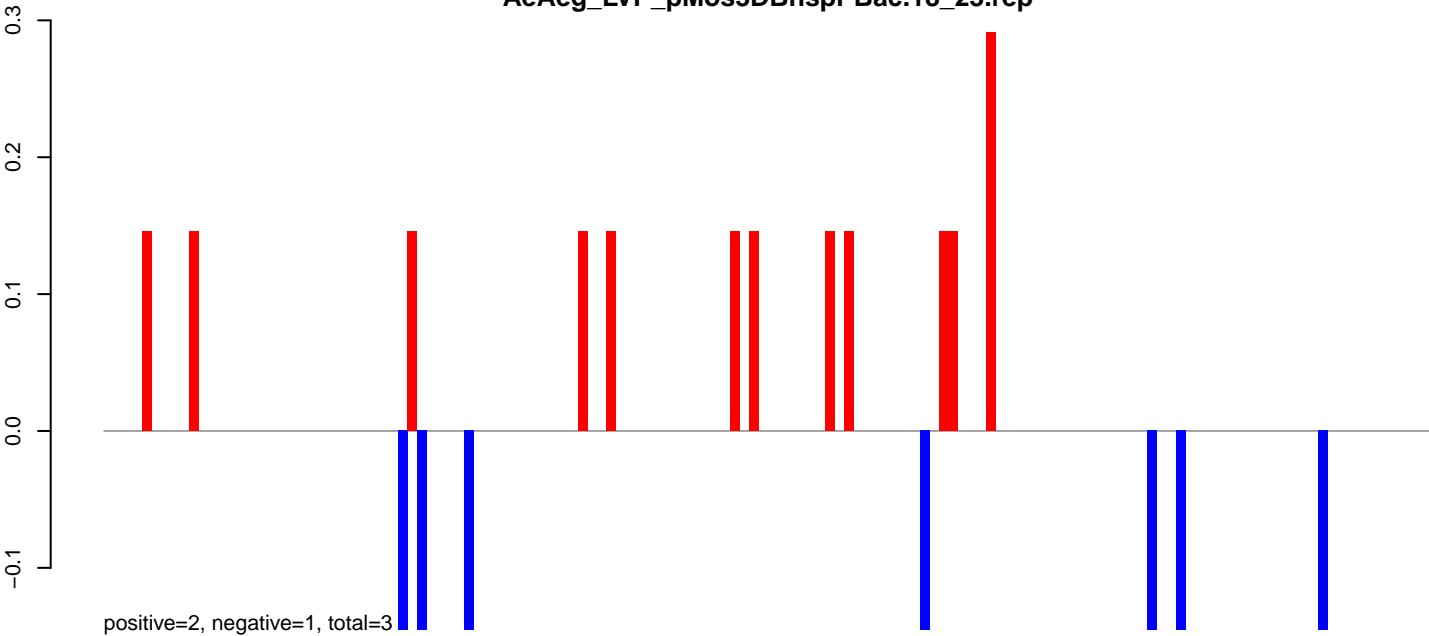
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



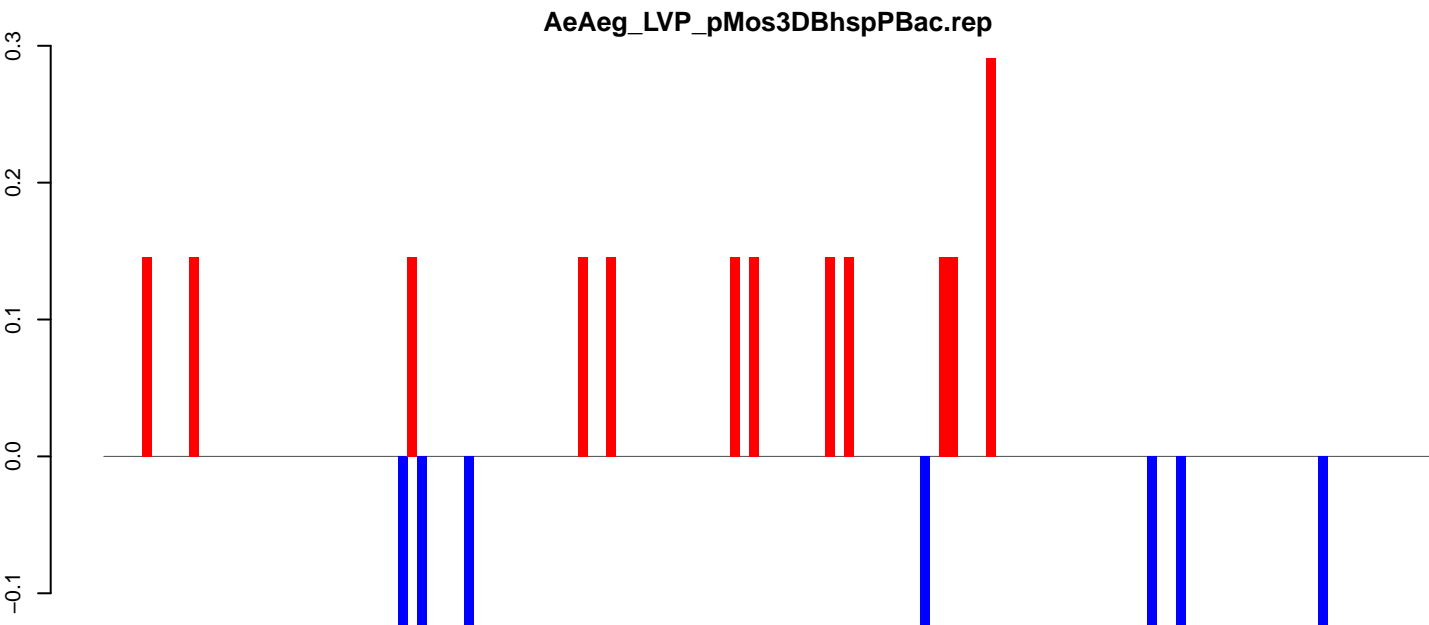
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



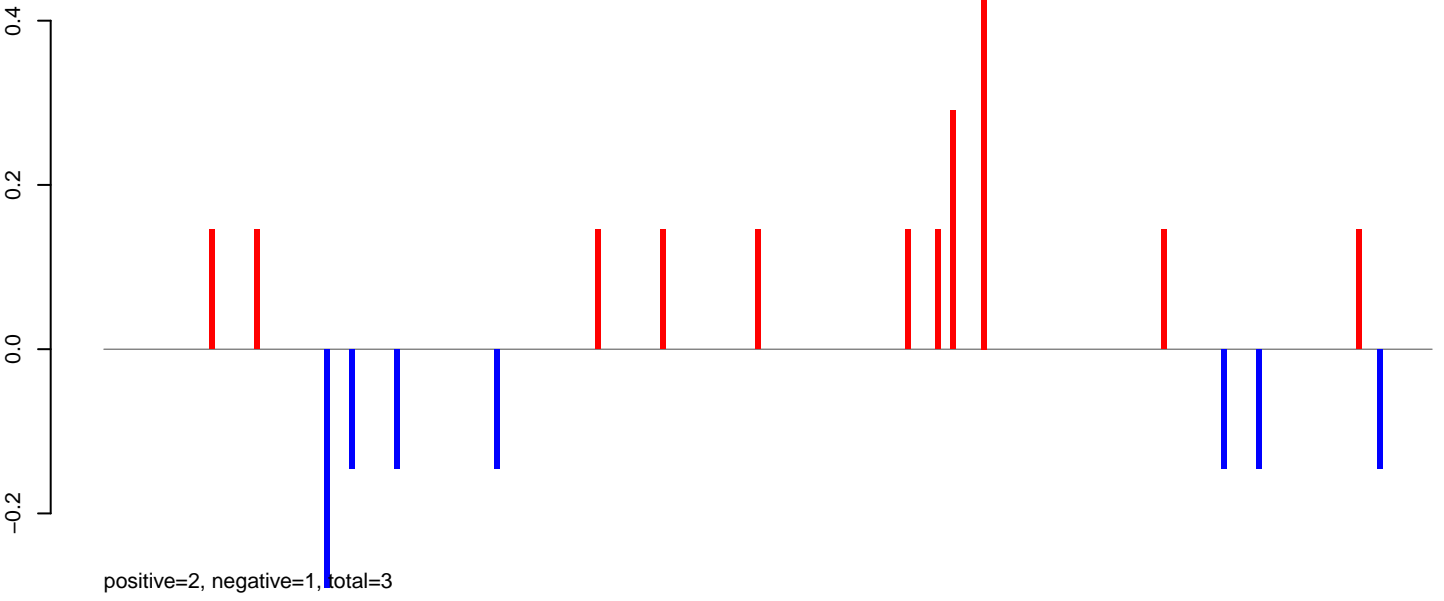
AeAeg_LVP_pMos3DBhspPBac.rep



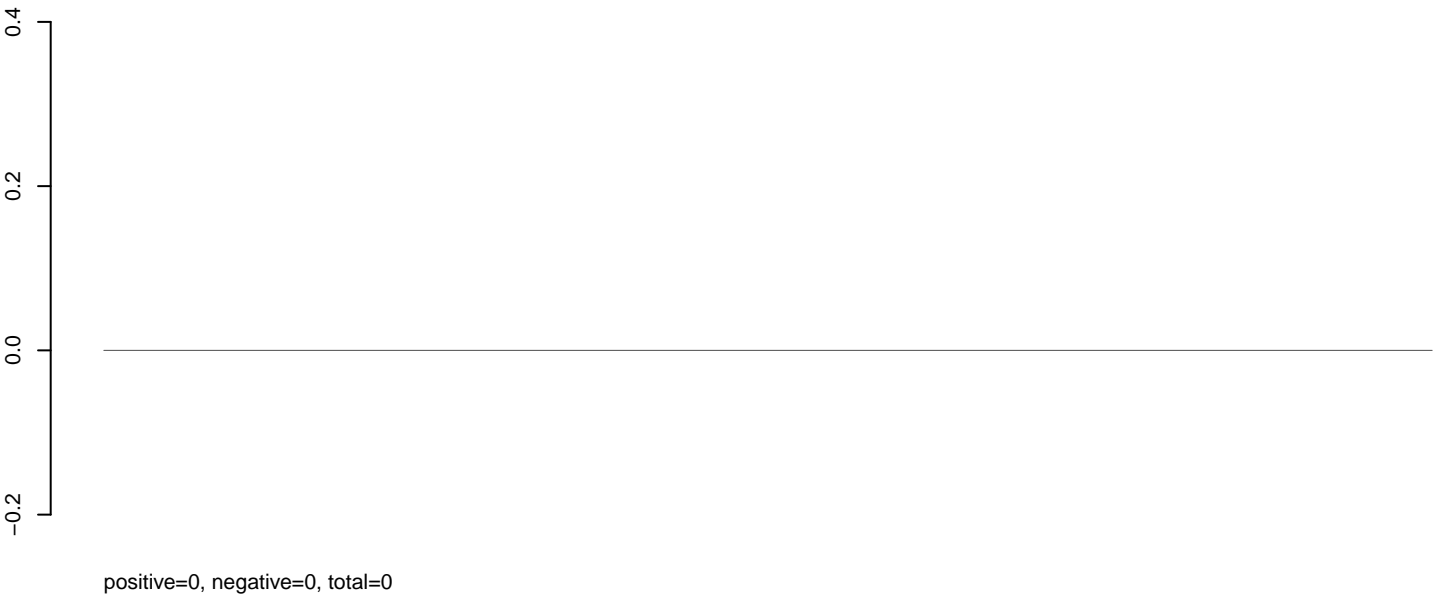
ow size=25, use=2, the=3498, -1, use=5707561-Culex_pipiens_pallens_densovirus_strain_JZ-16_non-structural_protein_1_NS1_non-stru:1-3498

0 500 1000 1500 2000 2500 3000 3500

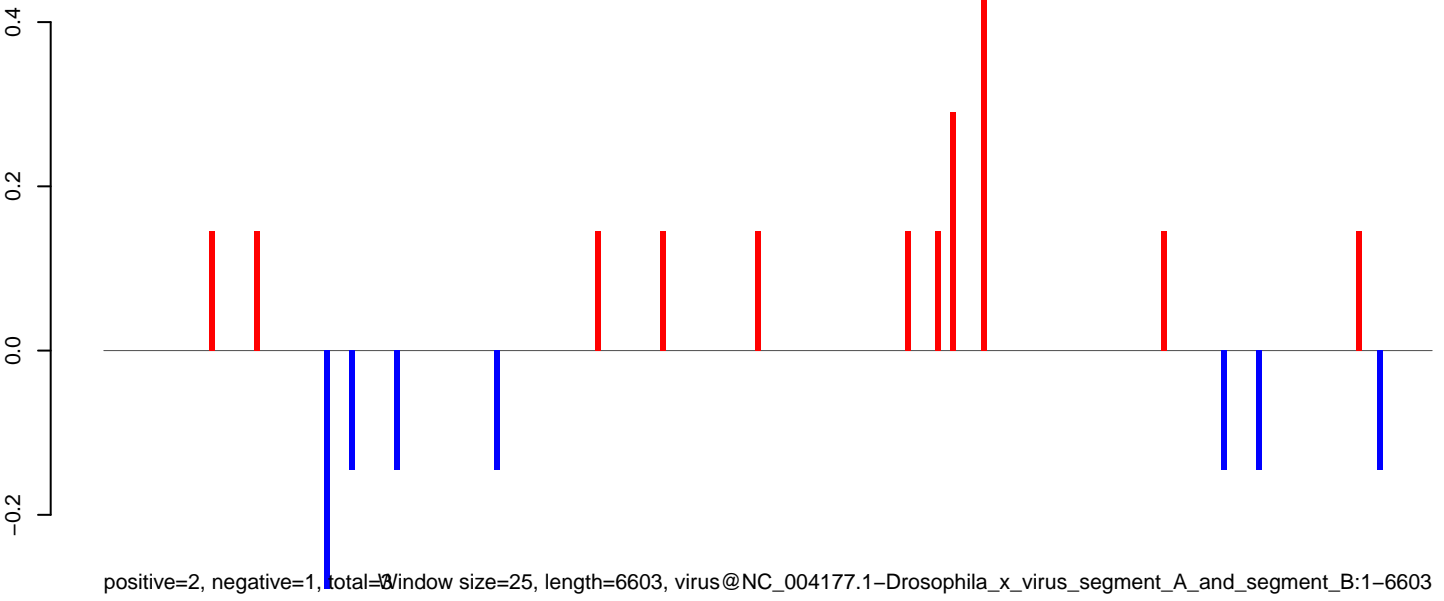
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

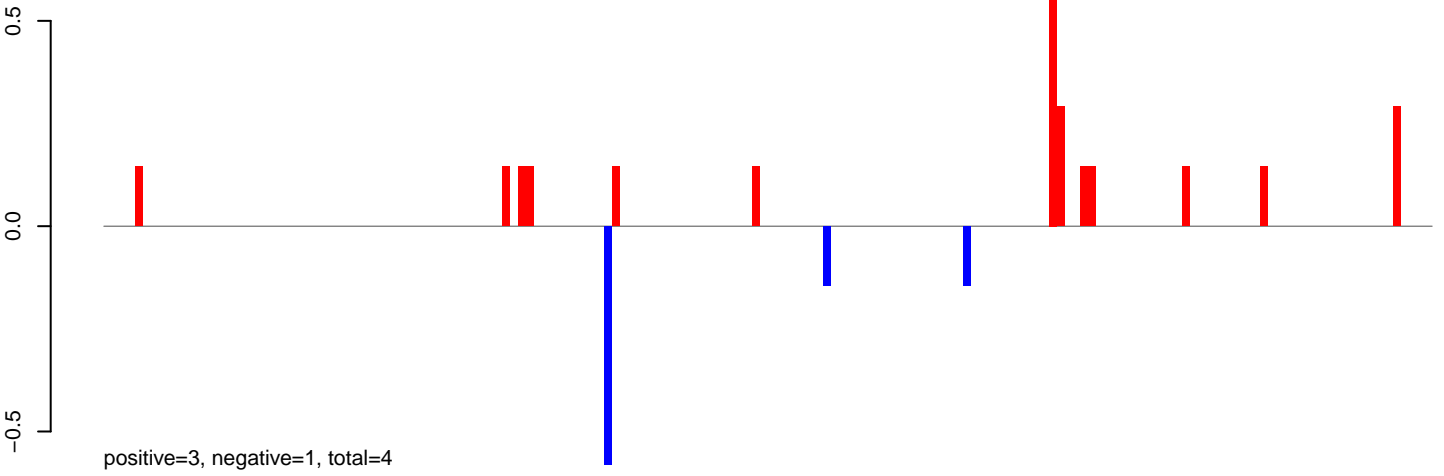


AeAeg_LVP_pMos3DBhspPBac.rep

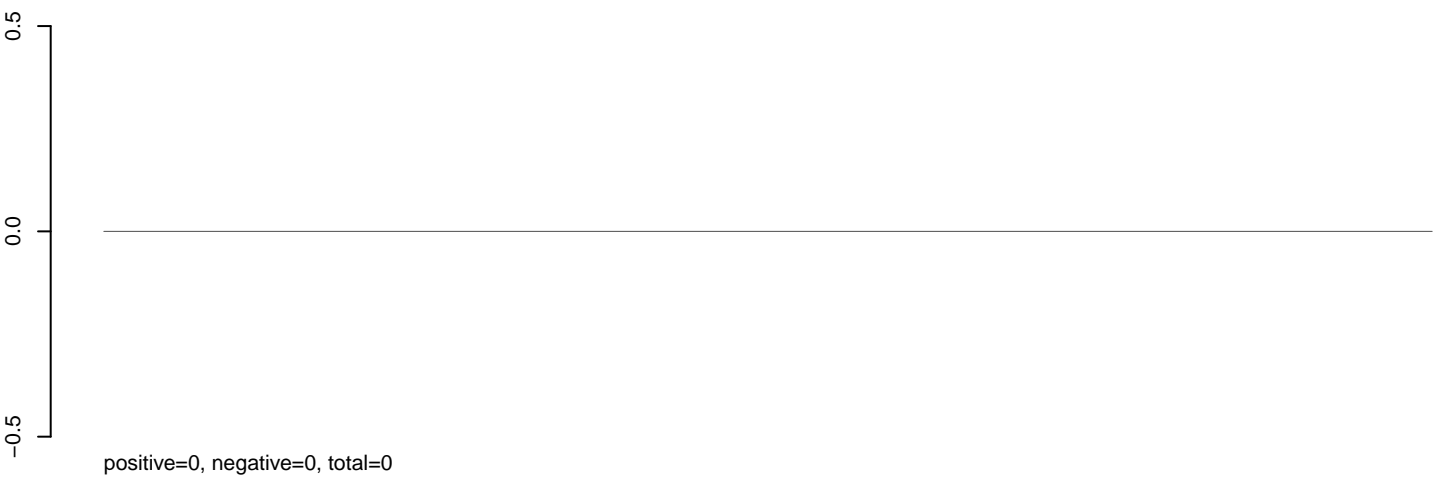


Window size=25, length=6603, virus@NC_004177.1-Drosophila_x_virus_segment_A_and_segment_B:1-6603

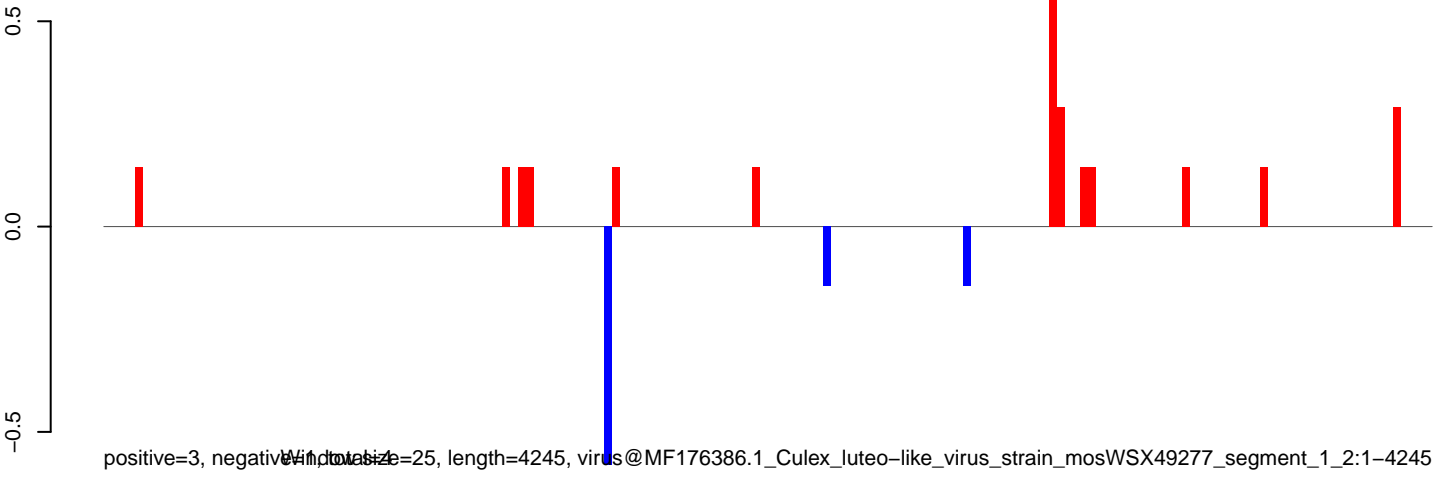
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

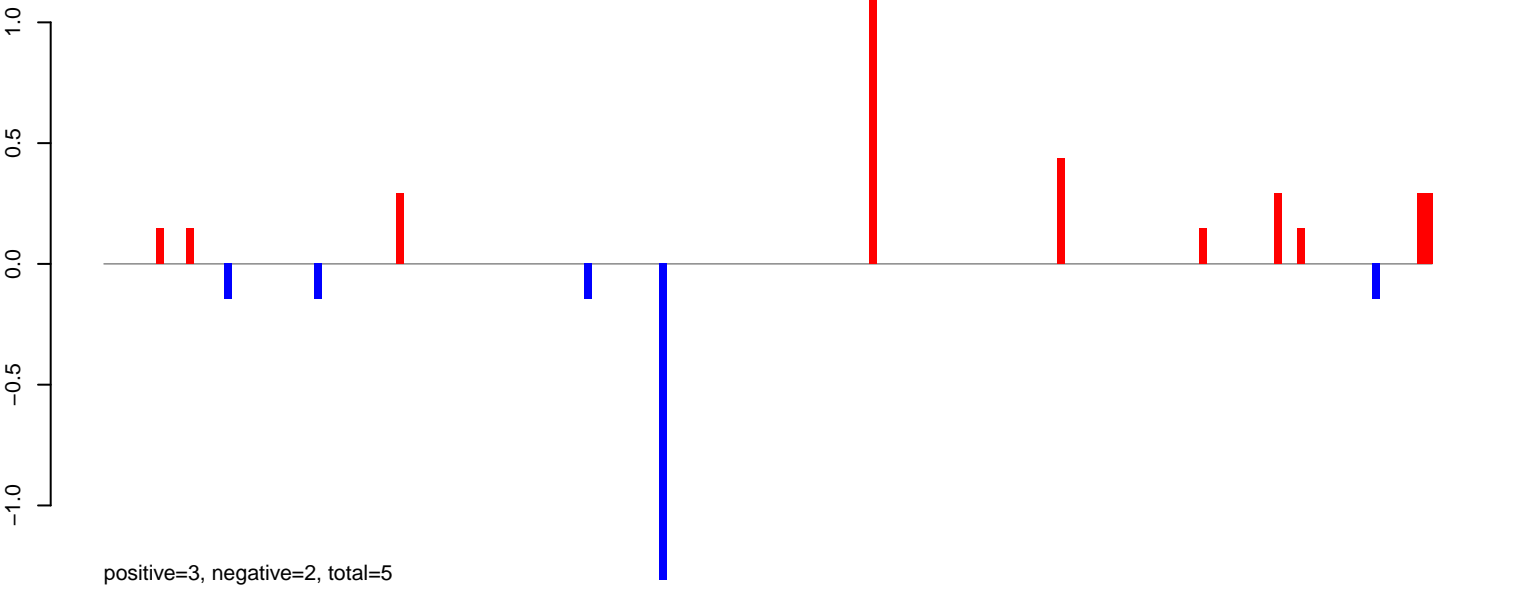


AeAeg_LVP_pMos3DBhspPBac.rep

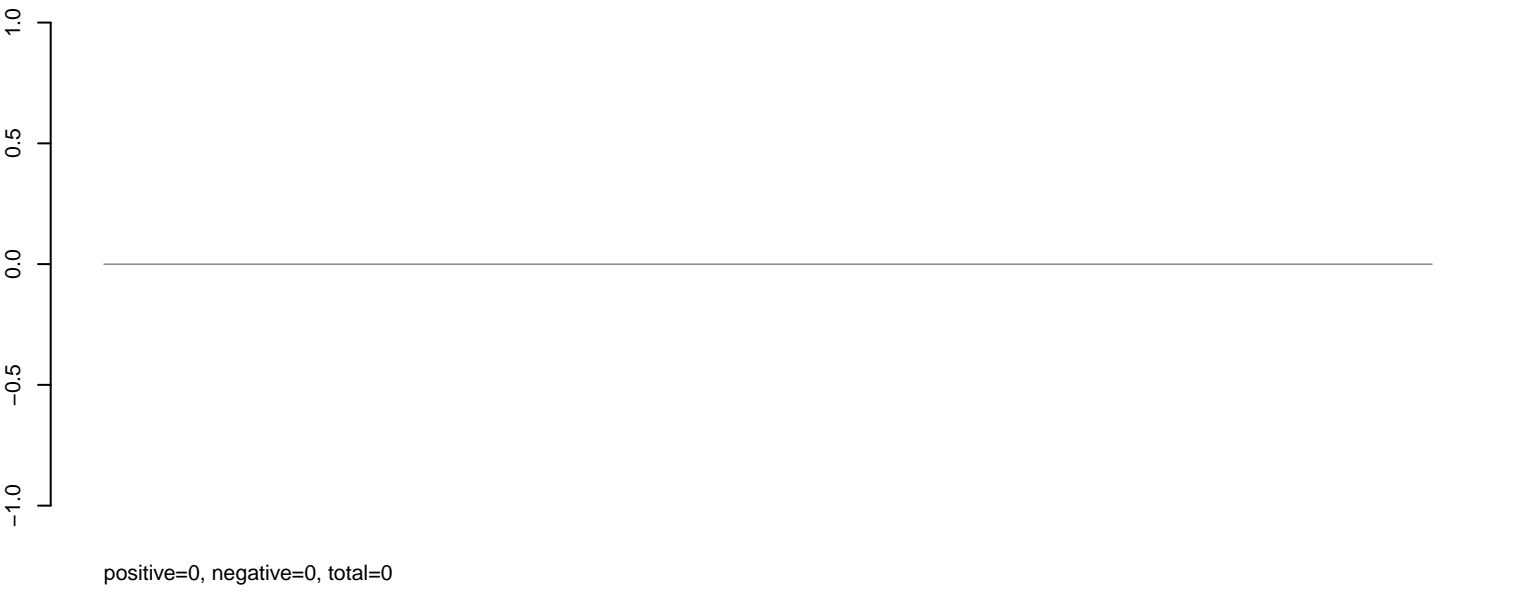


Window size=25, length=4245, virus@MF176386.1_Culex_luteo-like_virus_strain_mosWSX49277_segment_1_2:1-4245

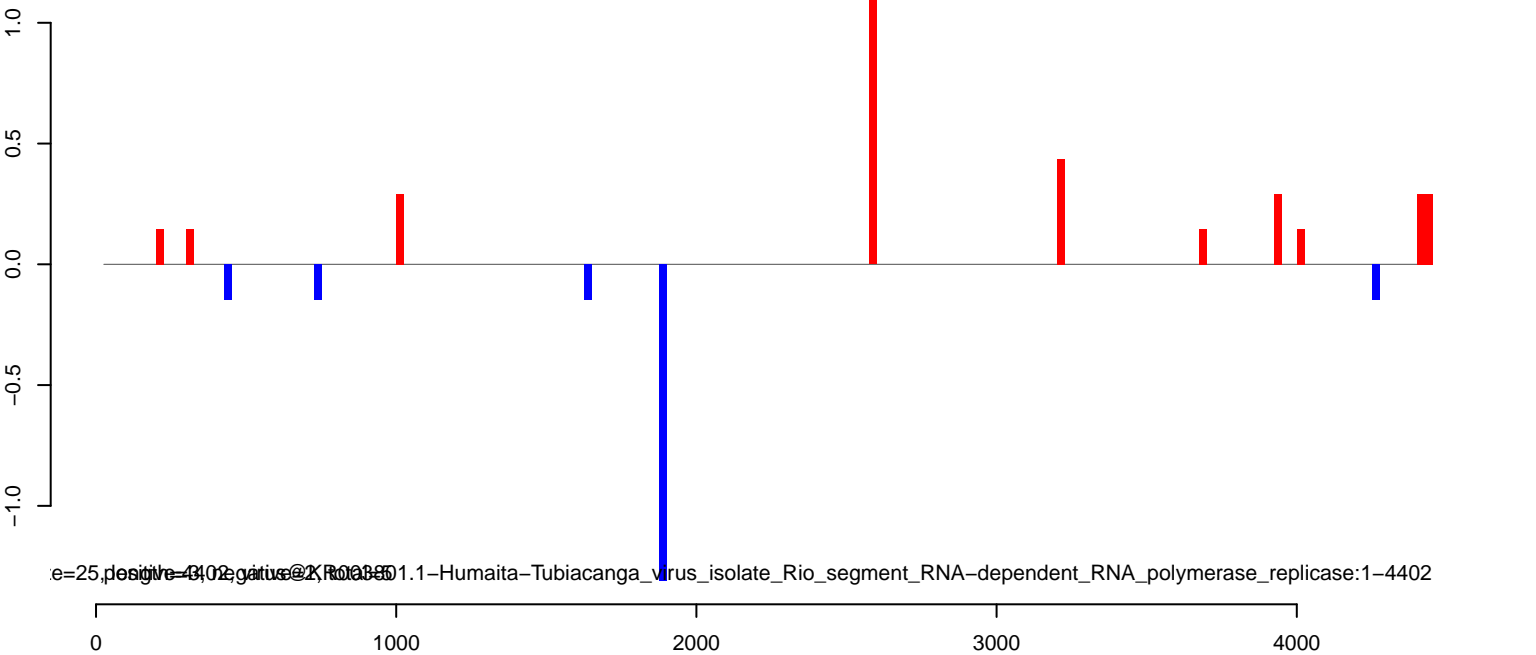
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



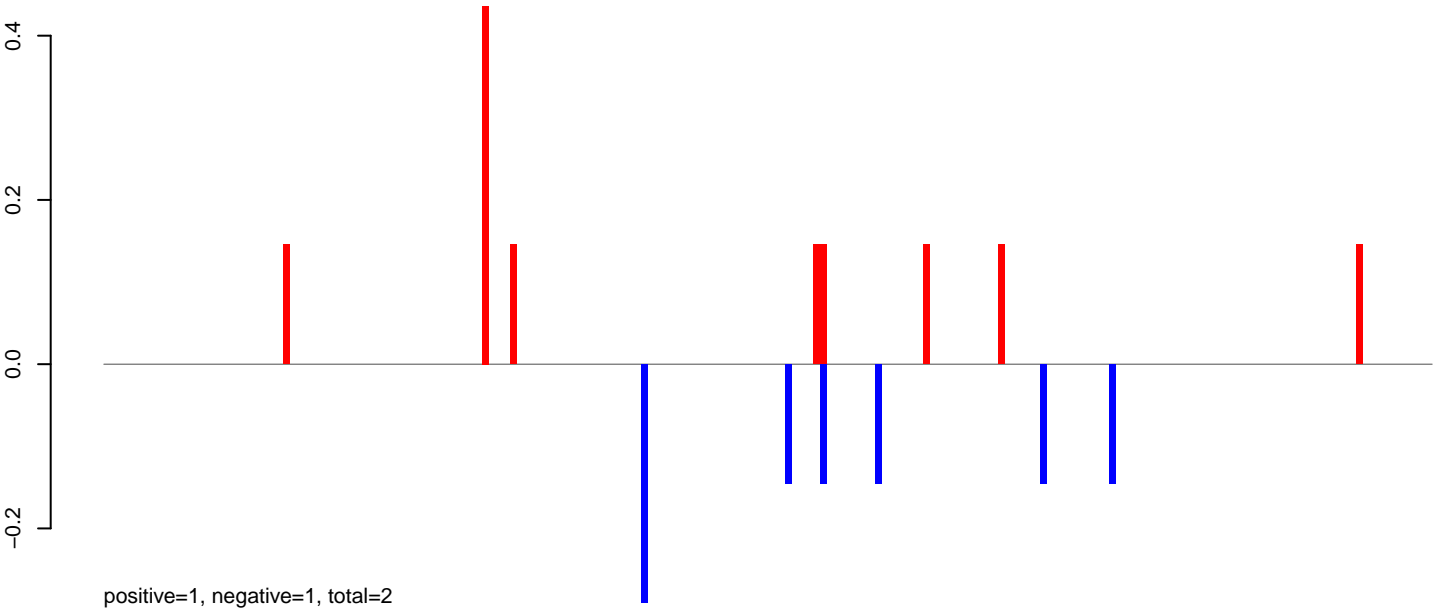
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



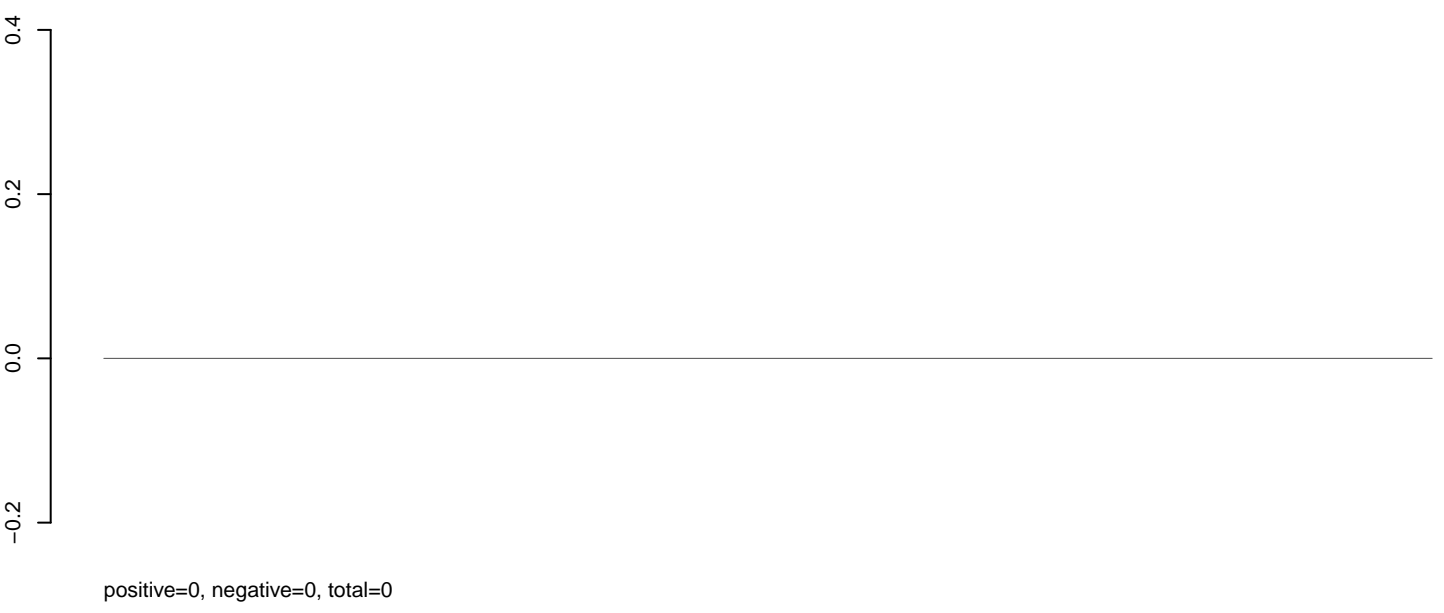
AeAeg_LVP_pMos3DBhspPBac.rep



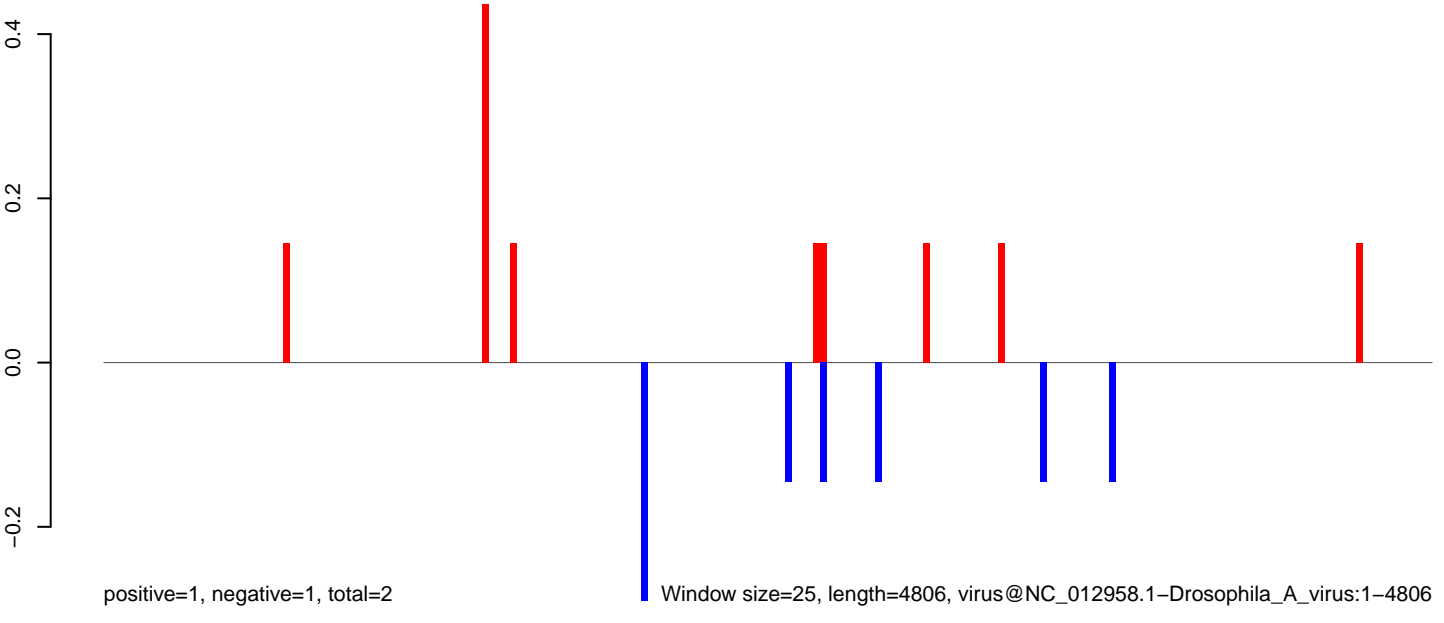
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



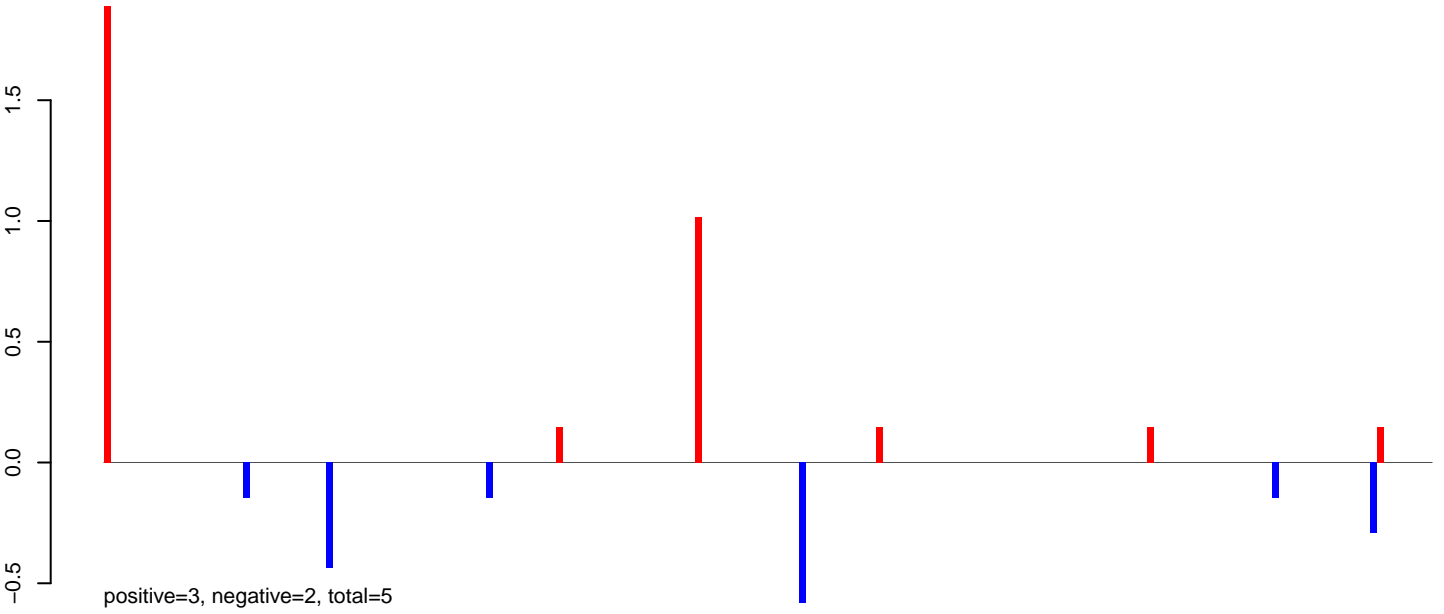
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



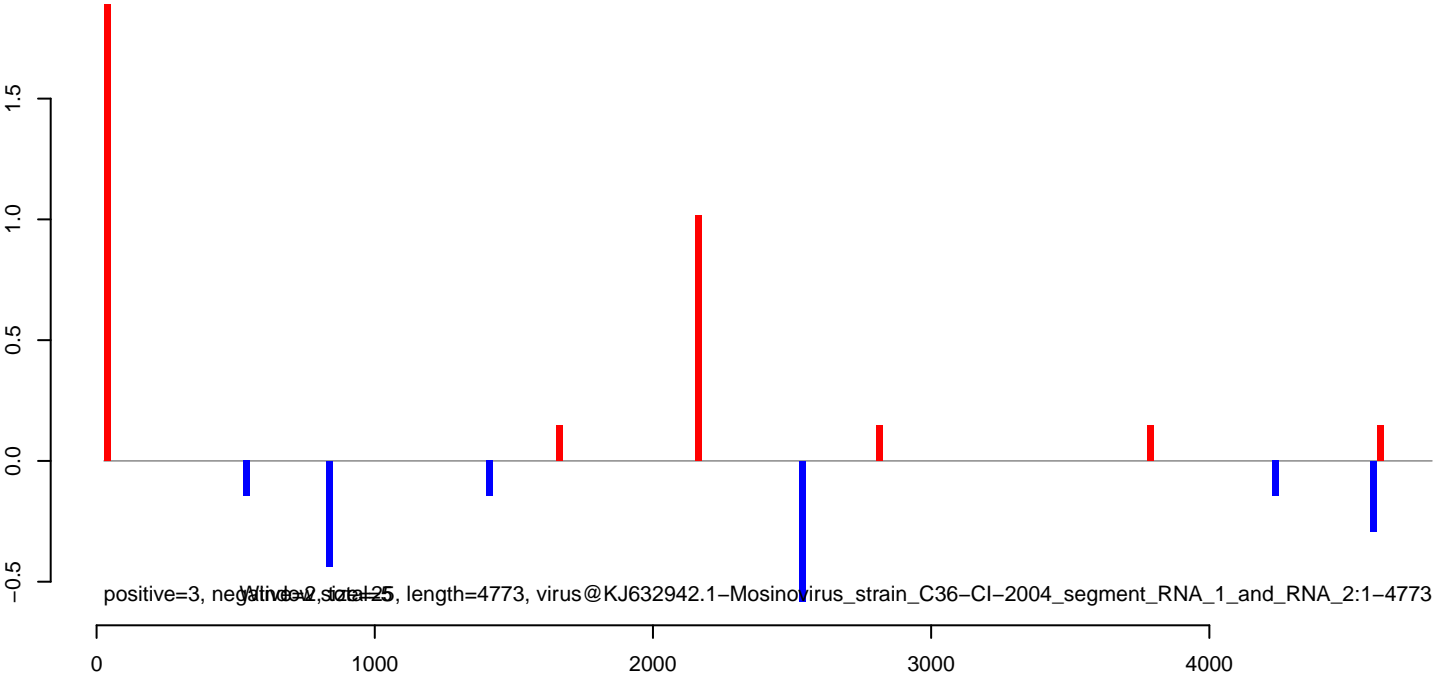
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



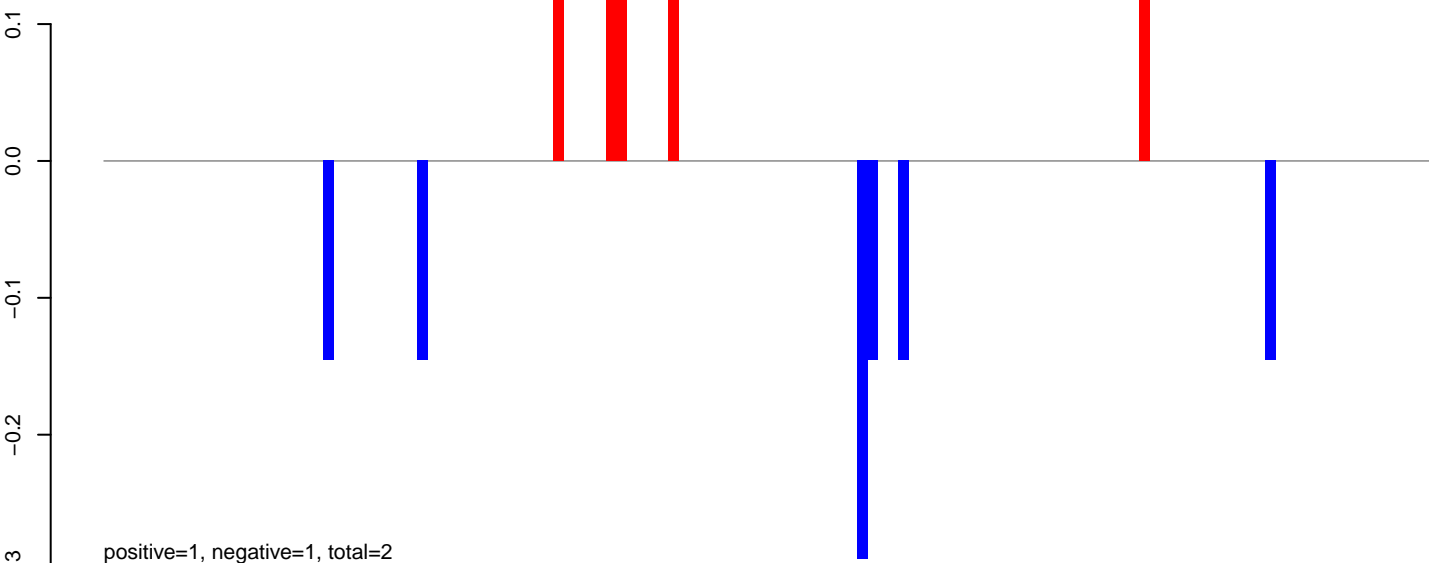
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



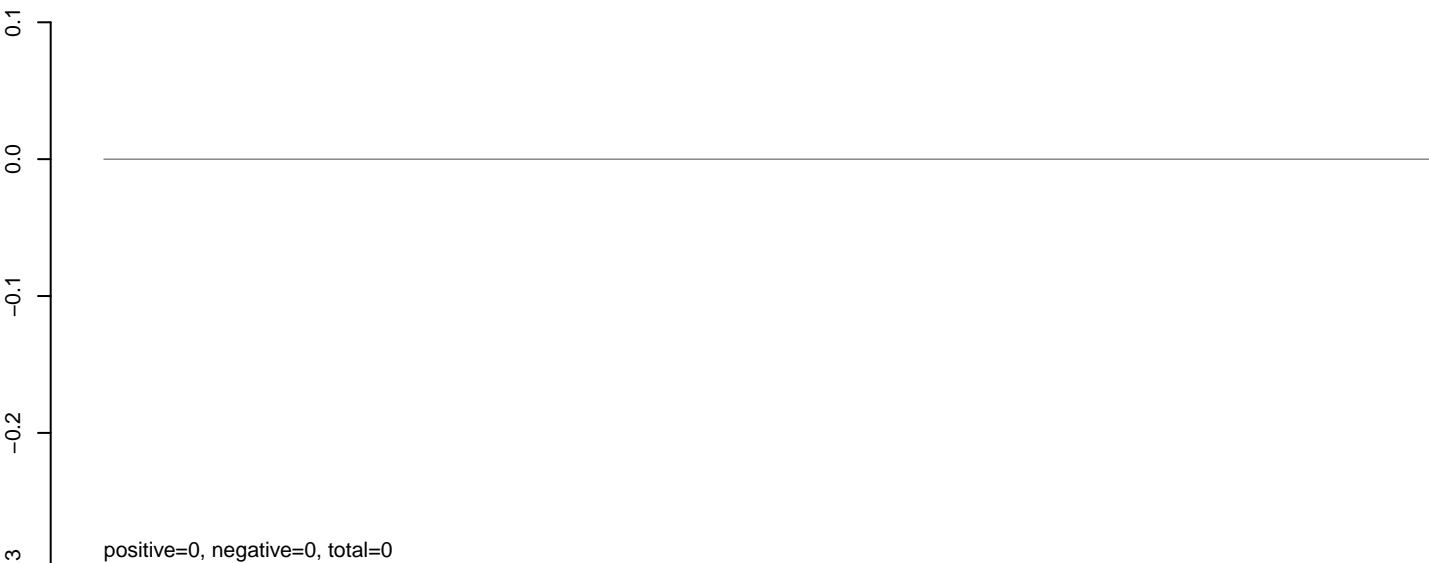
AeAeg_LVP_pMos3DBhspPBac.rep



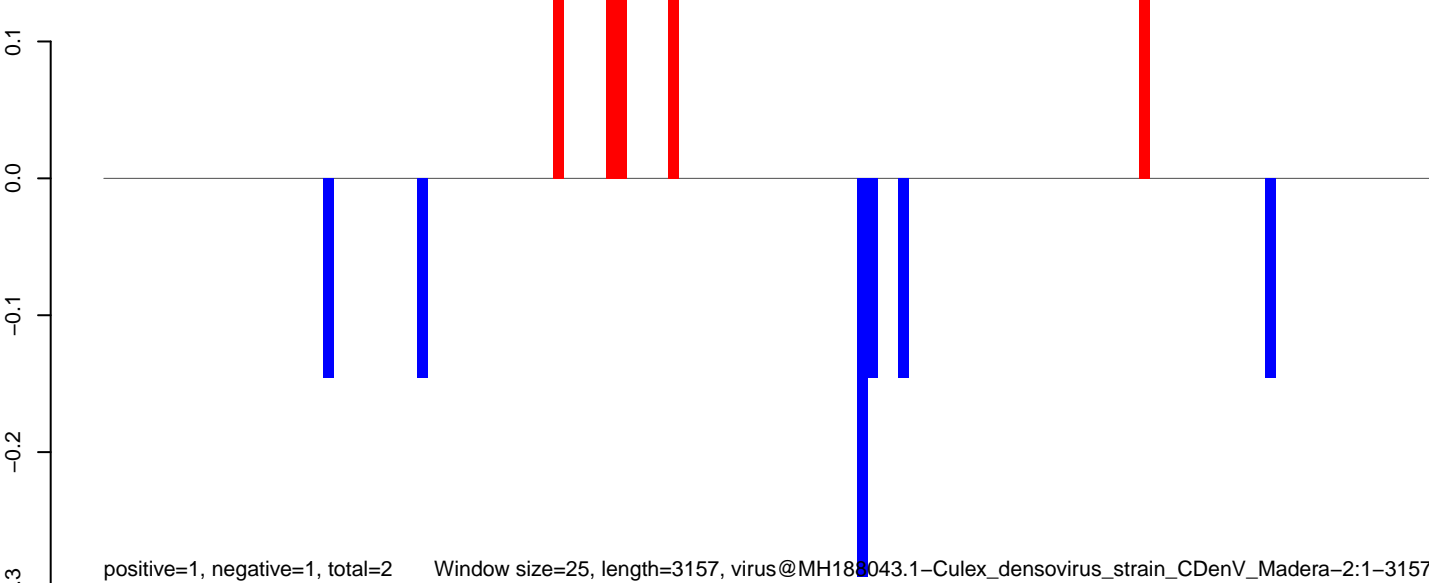
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep

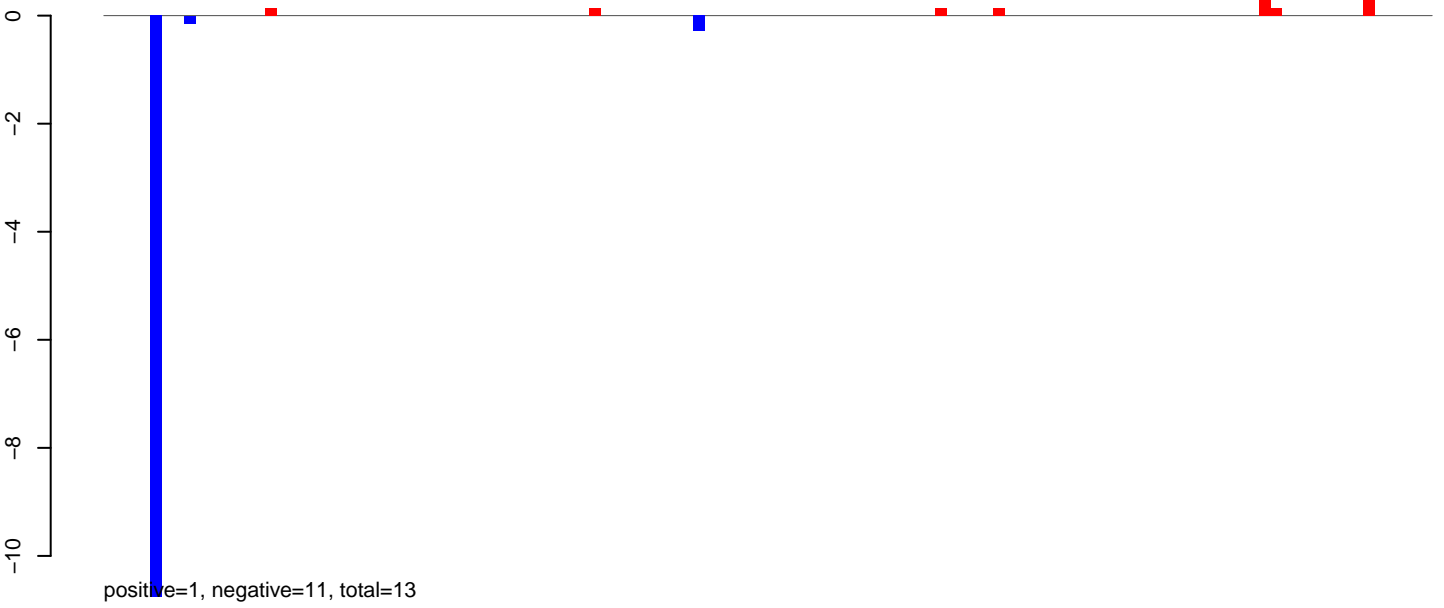


AeAeg_LVP_pMos3DBhspPBac.rep

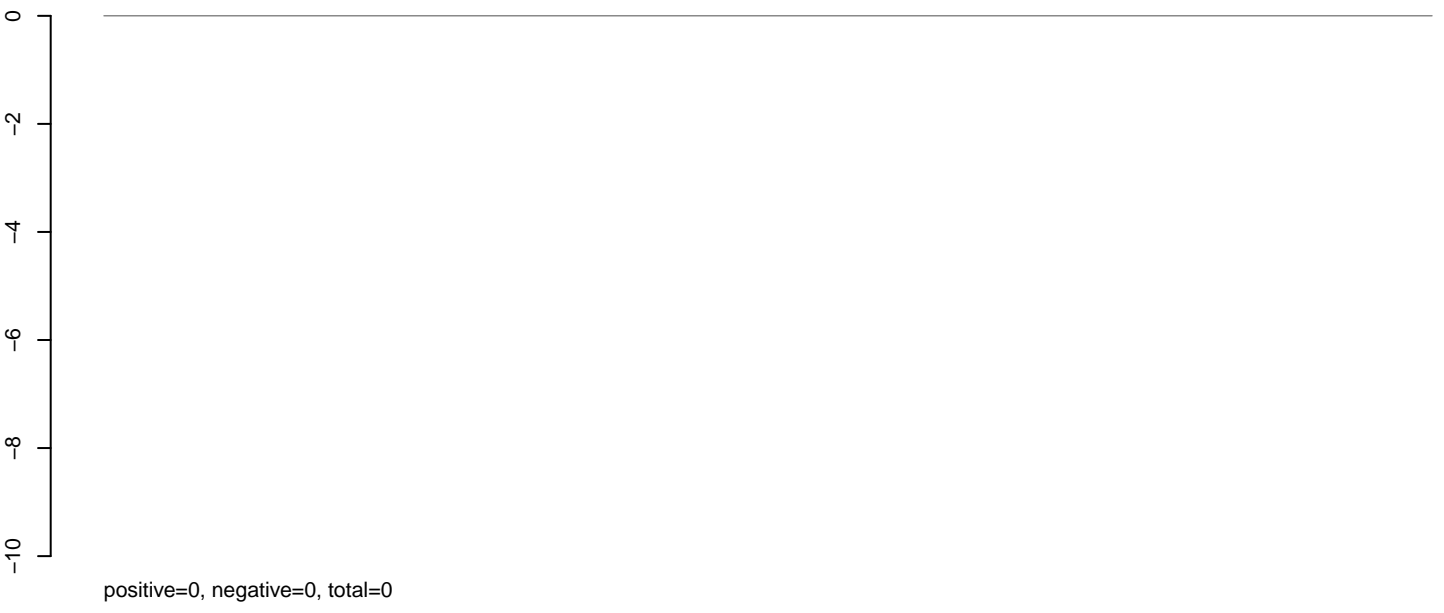


Window size=25, length=3157, virus@MH188043.1-Culex_densovirus_strain_CDENV_Madera-2:1-3157

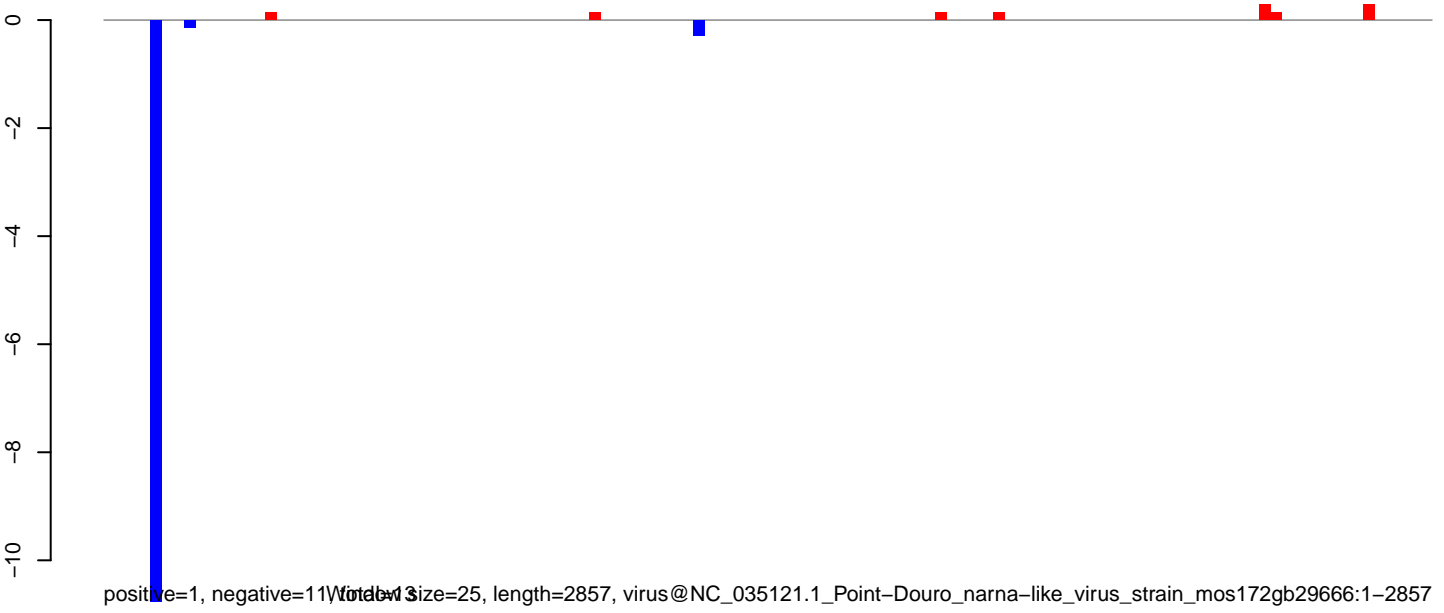
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



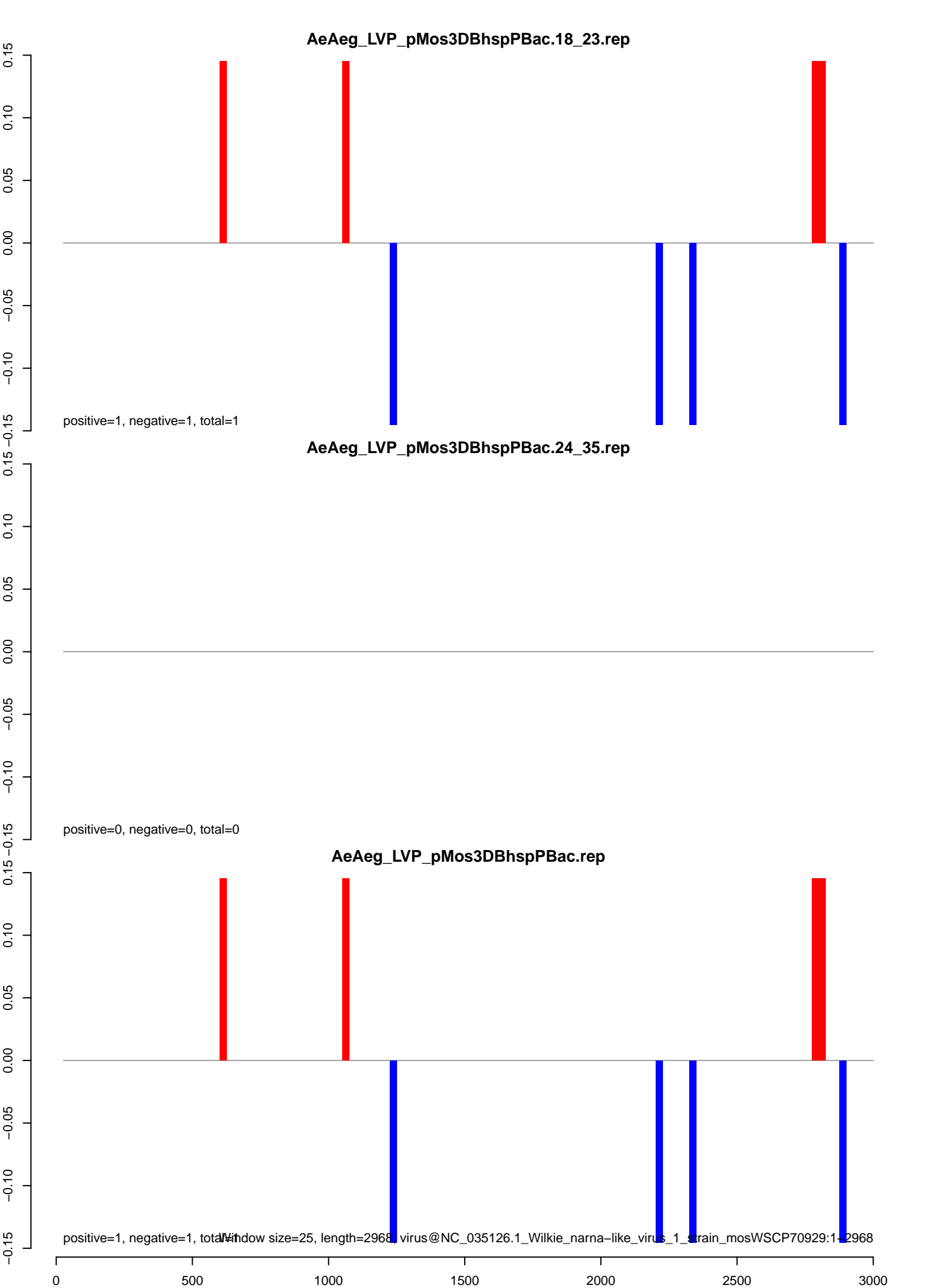
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



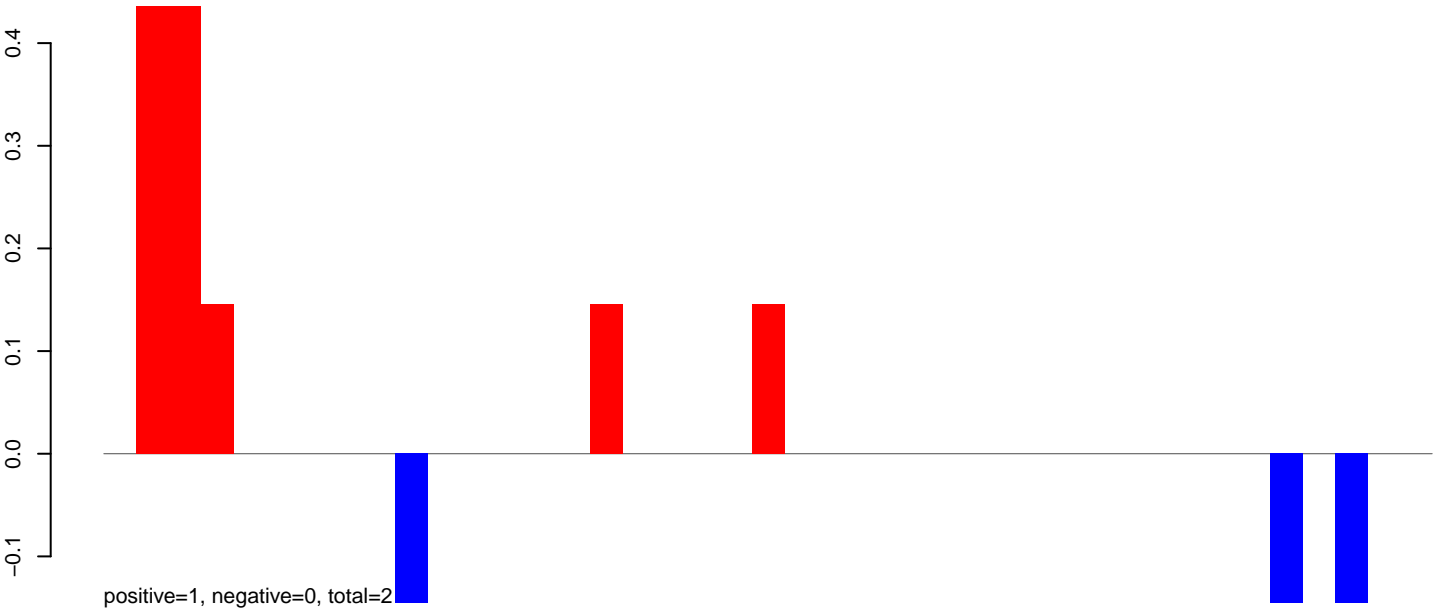
AeAeg_LVP_pMos3DBhspPBac.rep



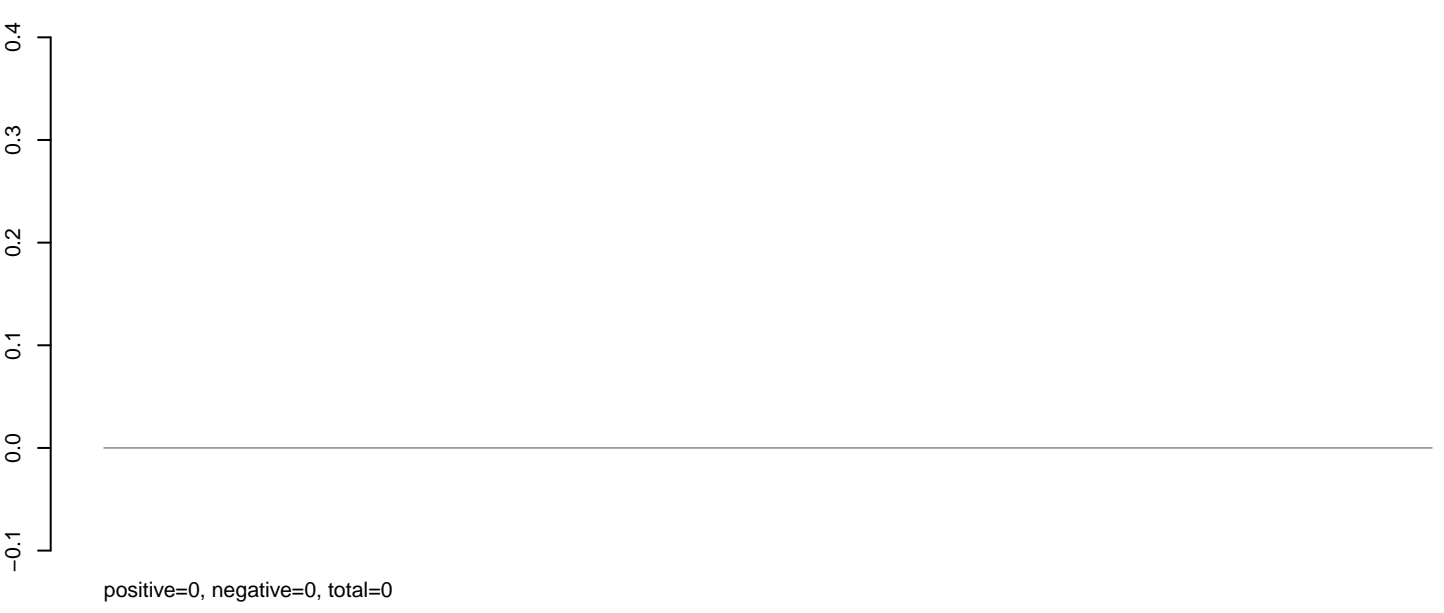
total size=25, length=2857, virus@NC_035121.1_Point-Douro_narna-like_virus_strain_mos172gb29666:1-2857



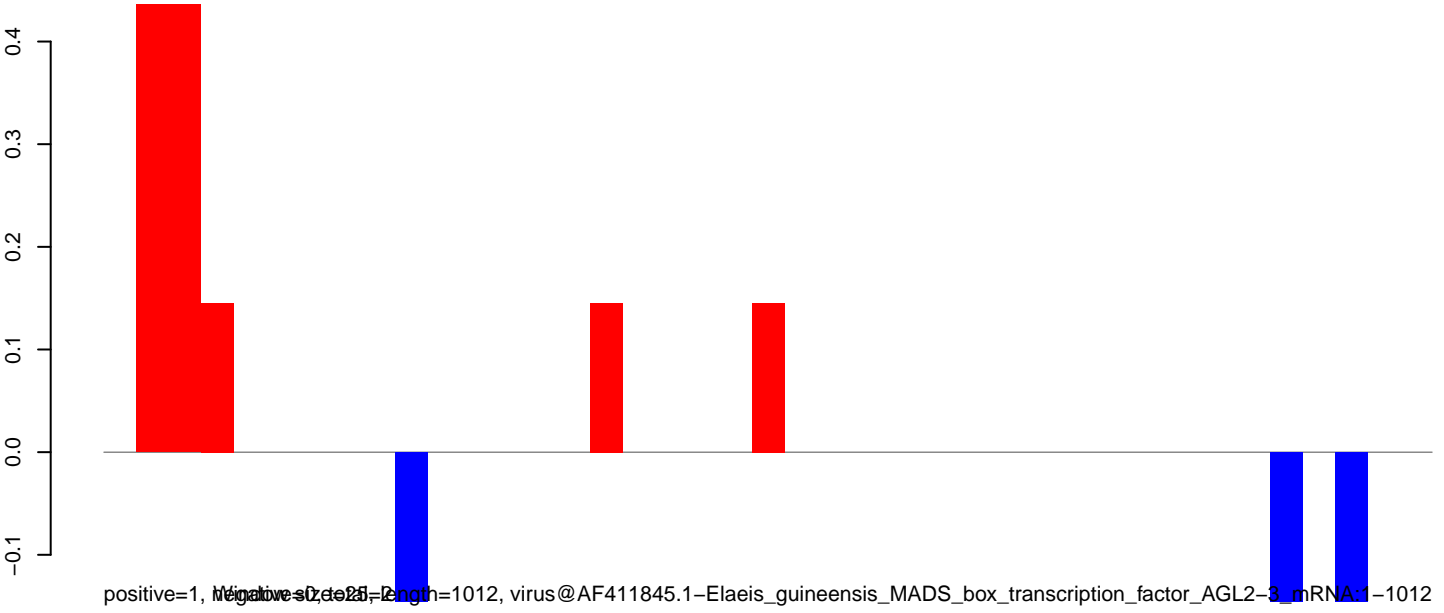
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



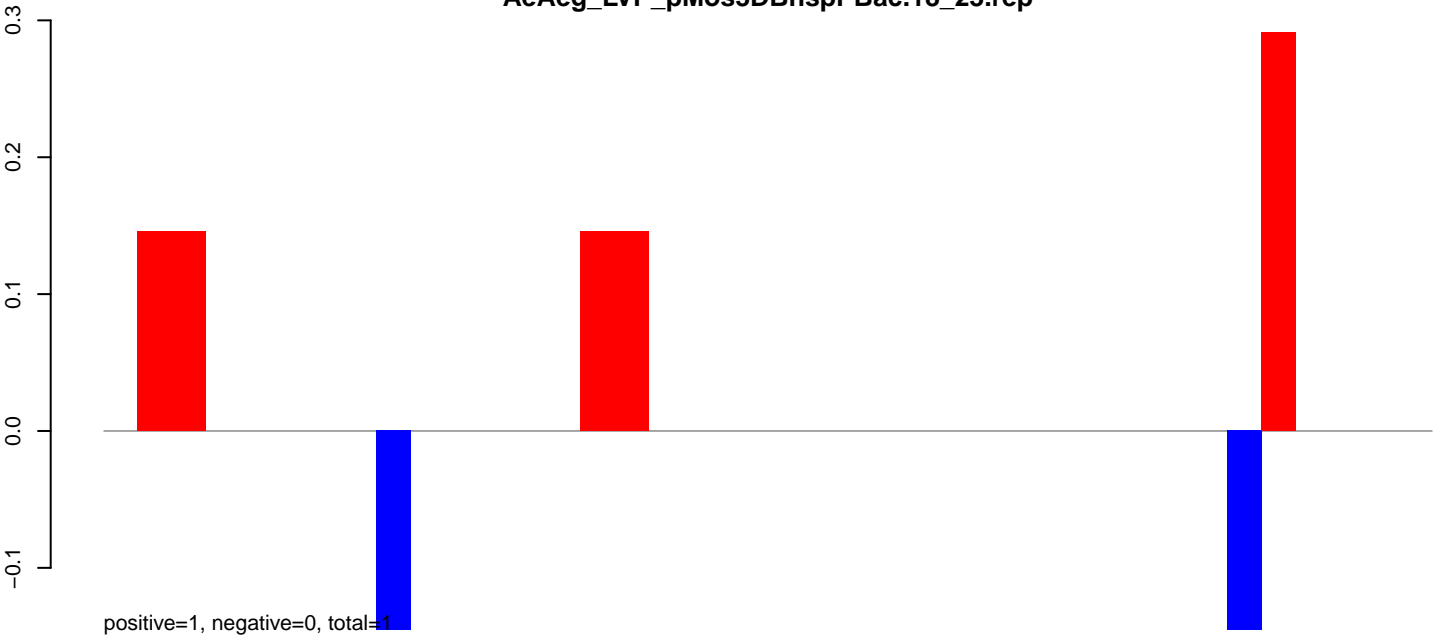
AeAeg_LVP_pMos3DBhspPBac.24_35.rep



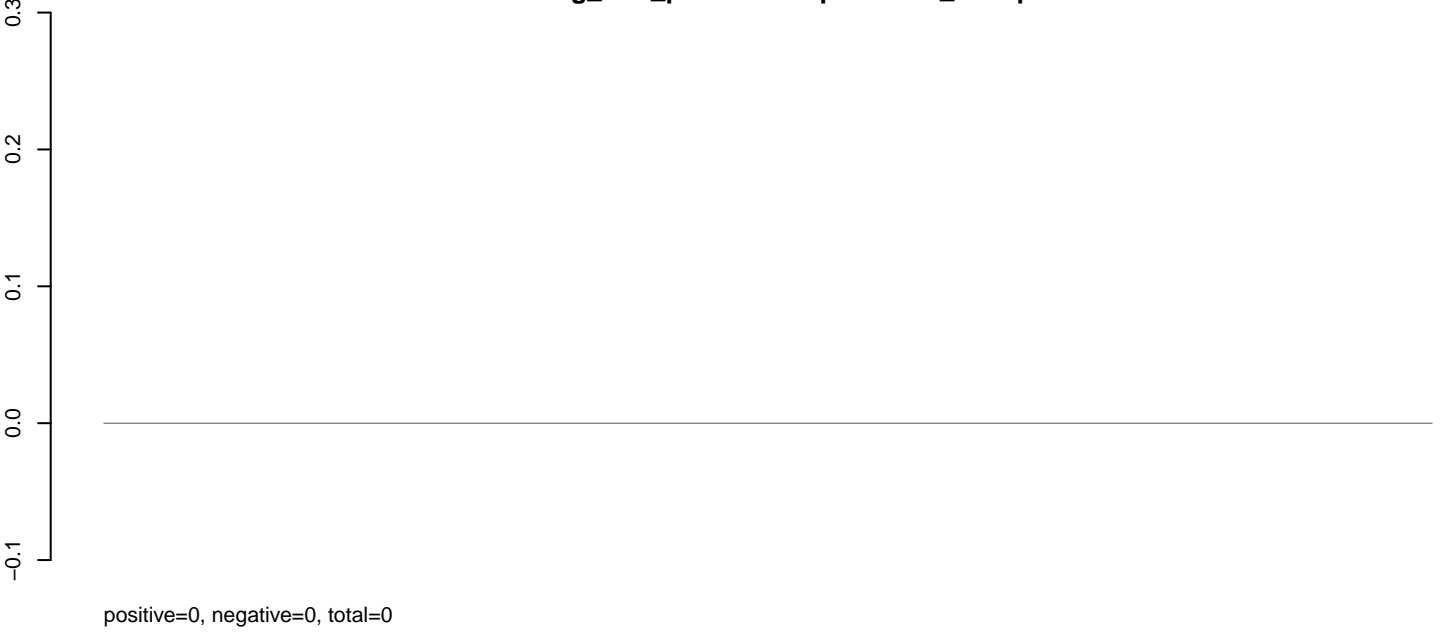
AeAeg_LVP_pMos3DBhspPBac.rep



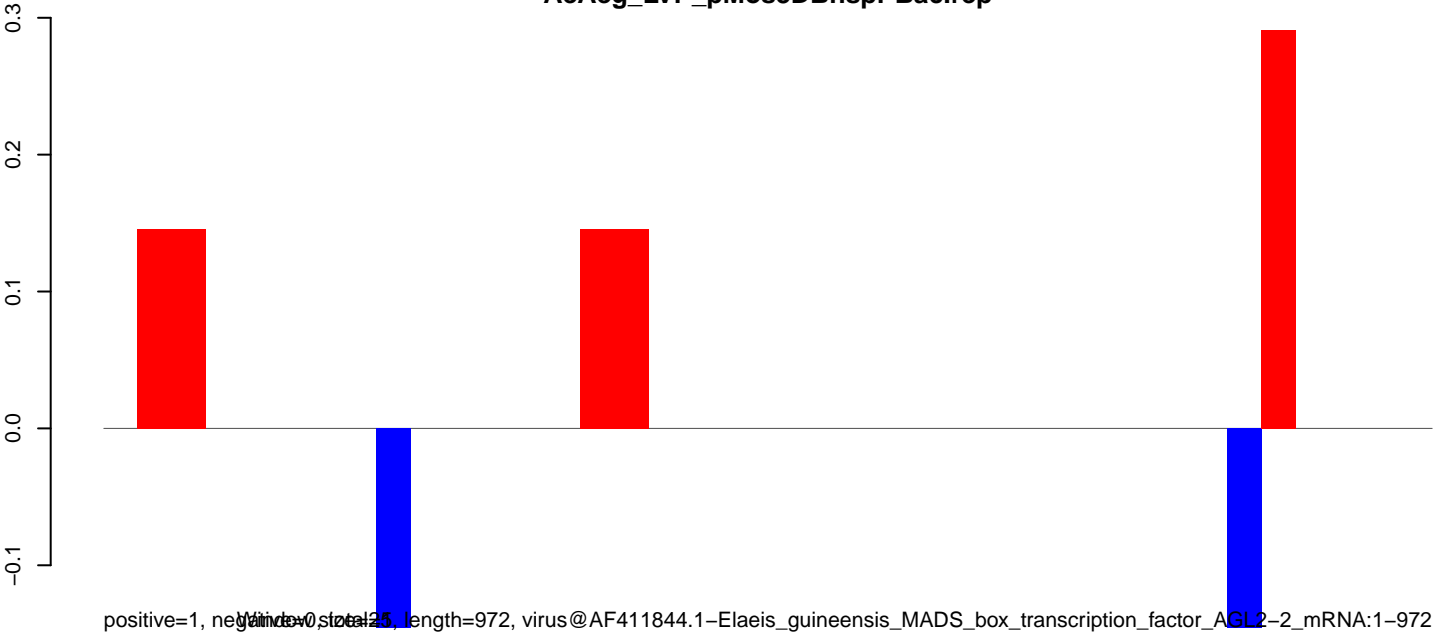
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



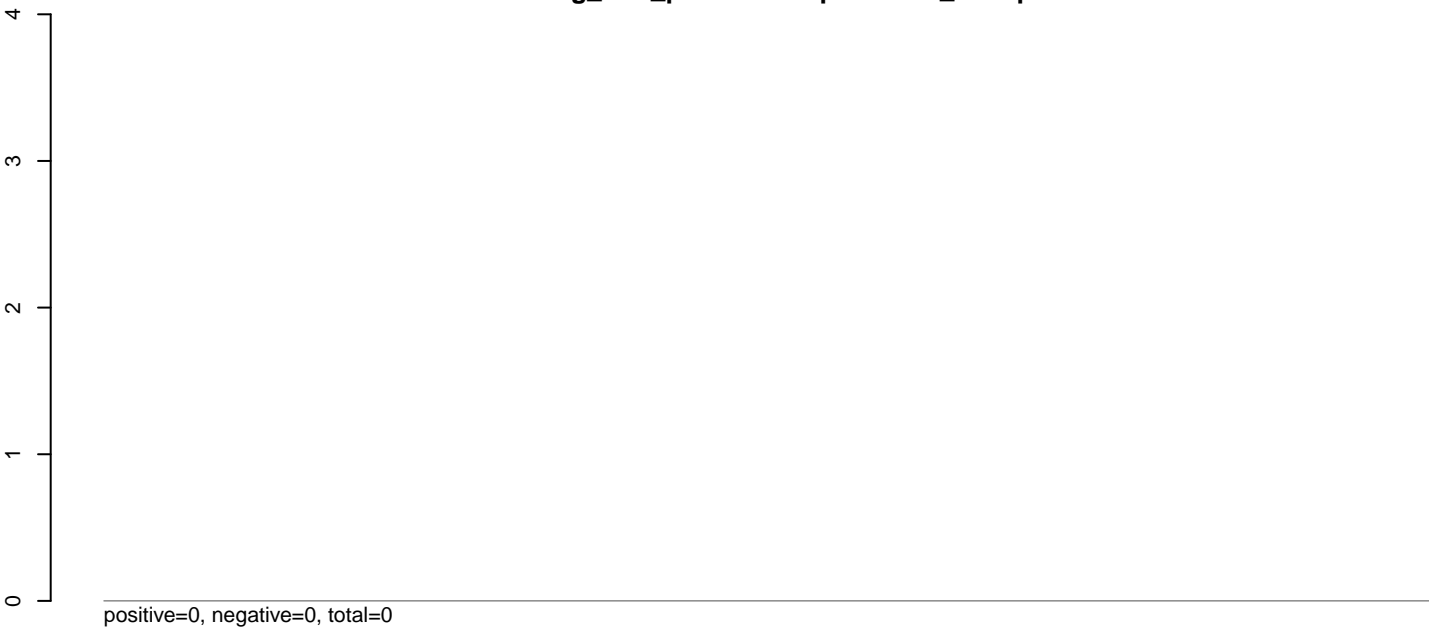
AeAeg_LVP_pMos3DBhspPBac.rep



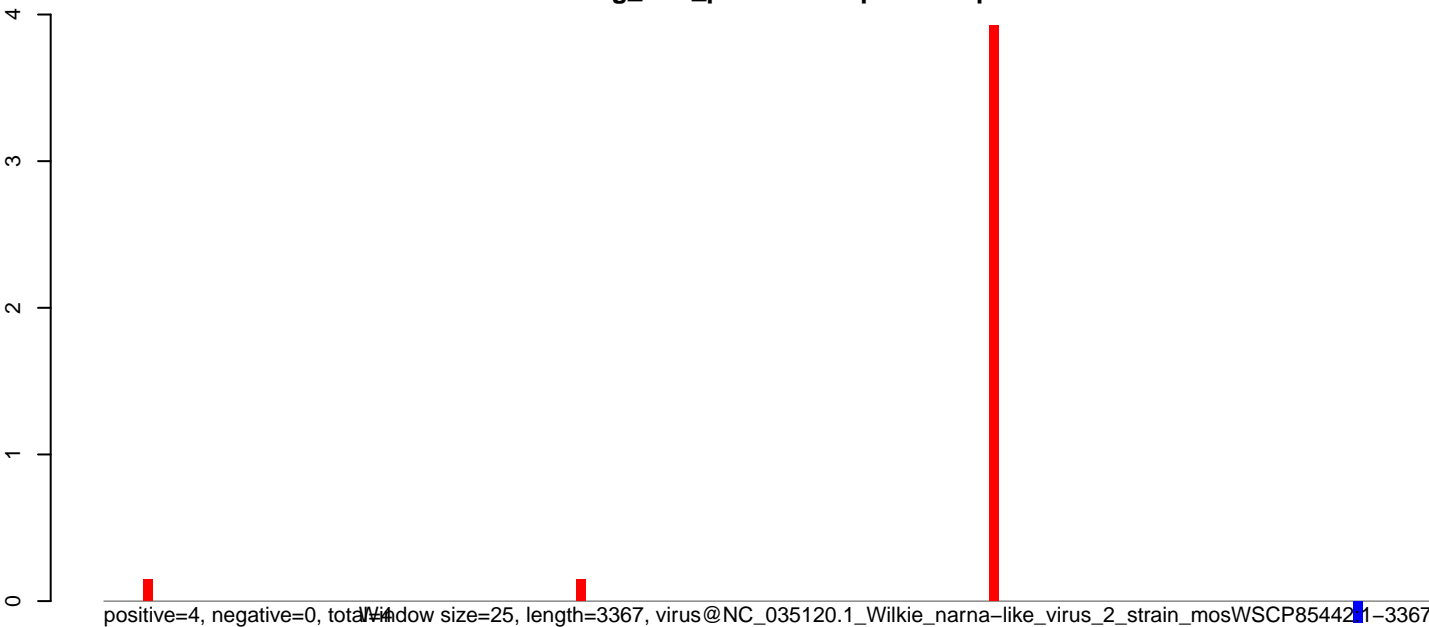
AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep



AeAeg_LVP_pMos3DBhspPBac.18_23.rep



AeAeg_LVP_pMos3DBhspPBac.24_35.rep



AeAeg_LVP_pMos3DBhspPBac.rep

