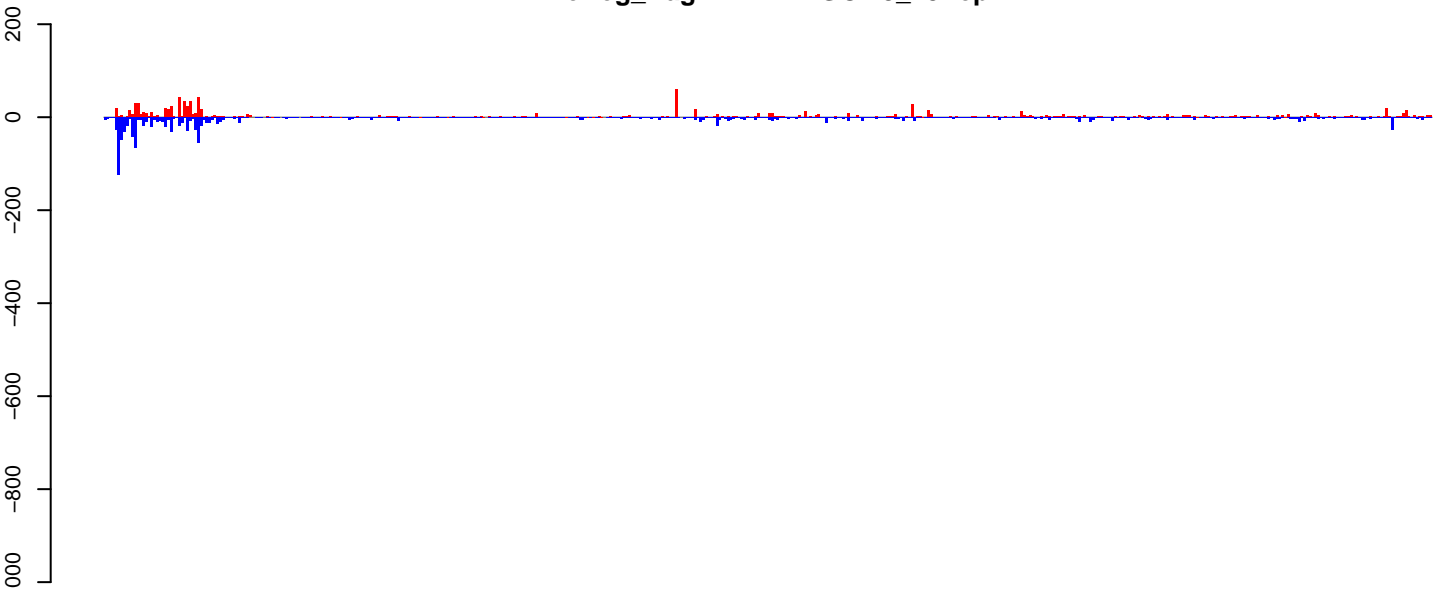
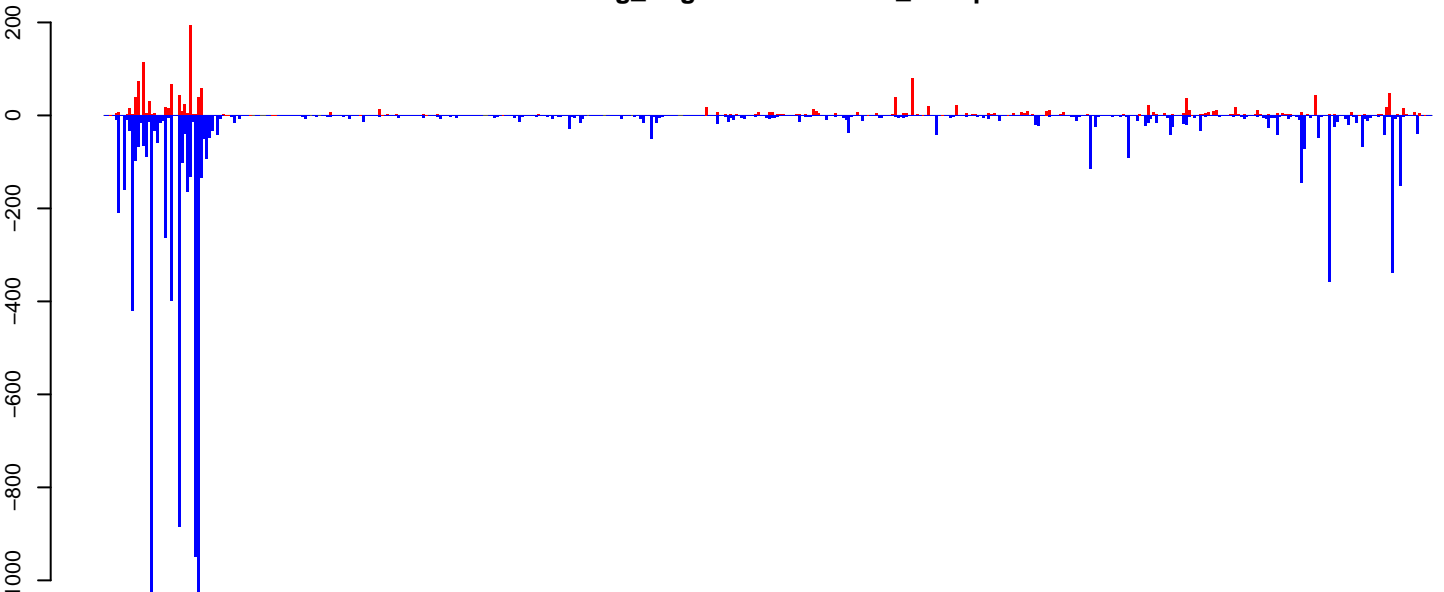


AeAeg_Aag2DENV2NGC.18_23.rep



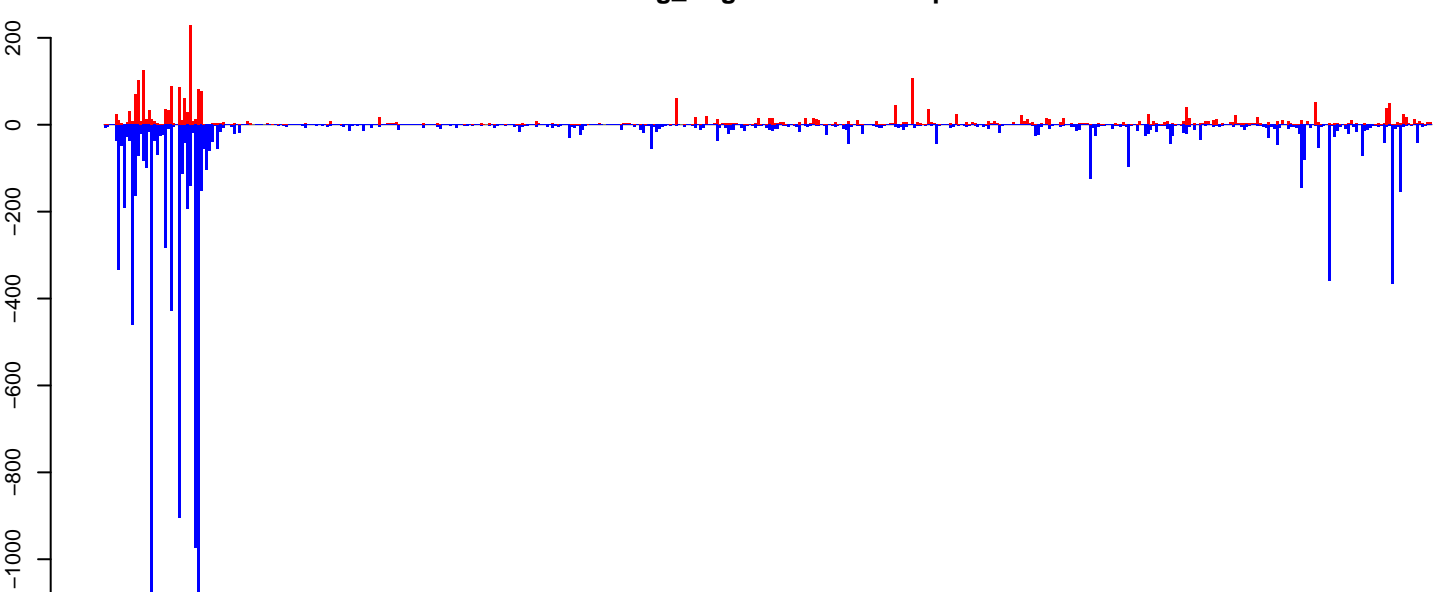
positive=1099, negative=1374, total=2473

AeAeg_Aag2DENV2NGC.24_35.rep



positive=1747, negative=9730, total=11477

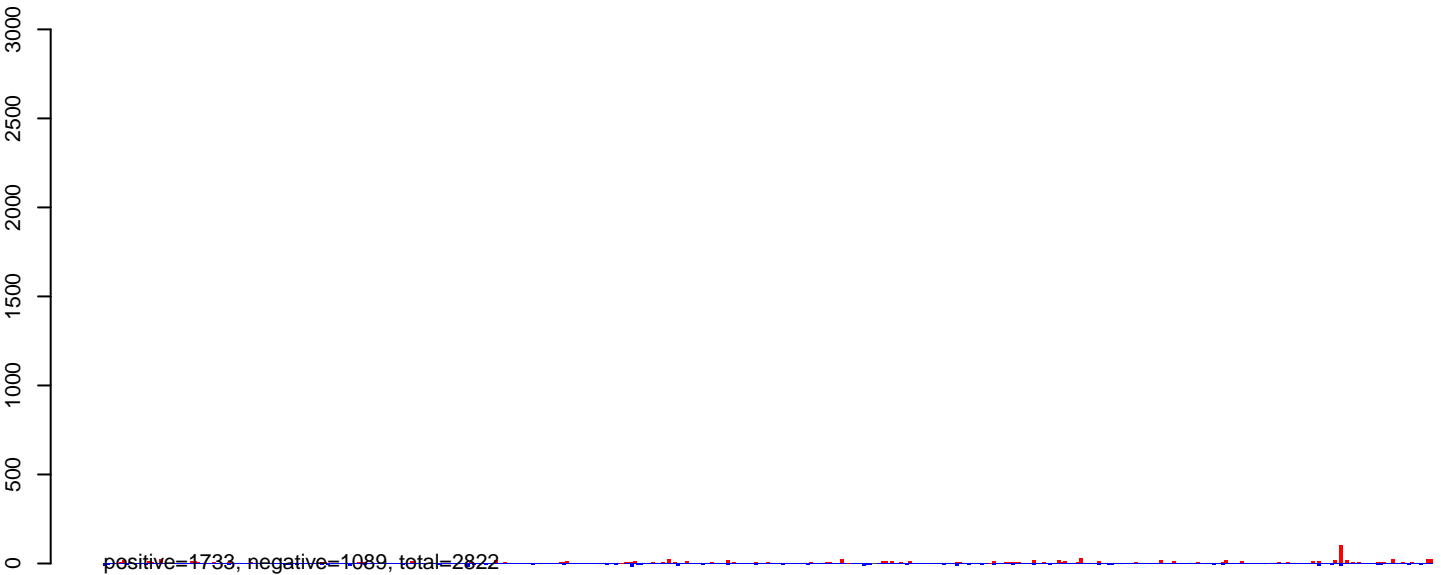
AeAeg_Aag2DENV2NGC.rep



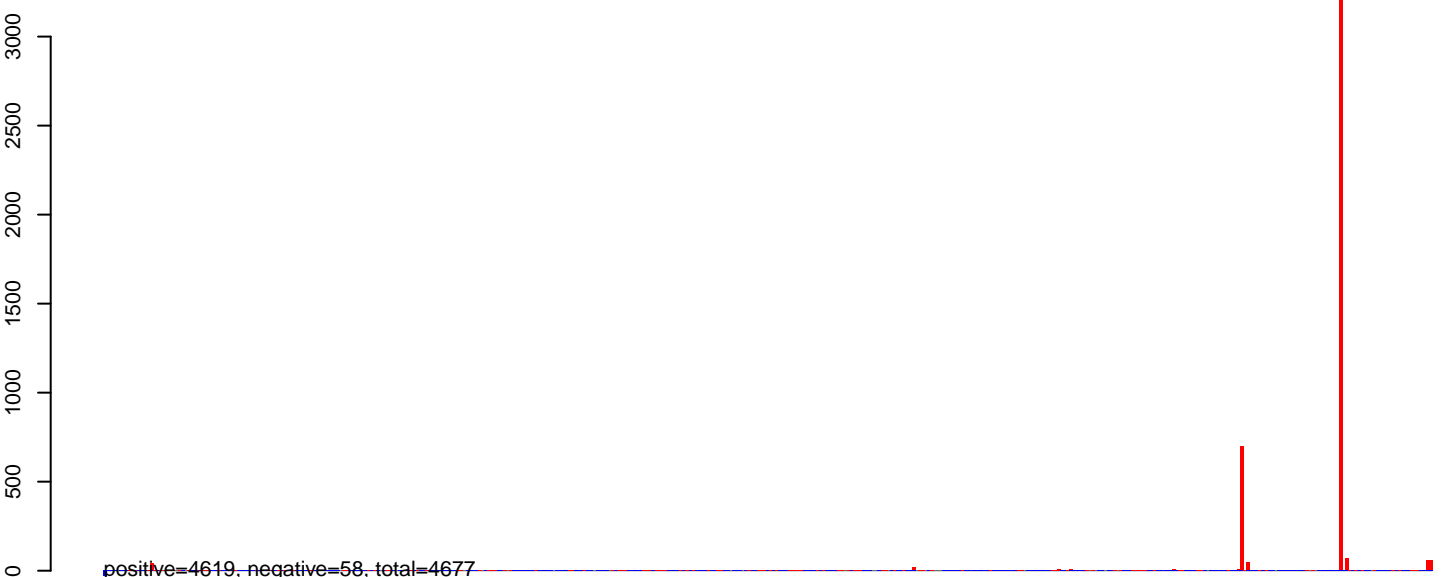
-25, length=12097, vireg@MC=1038263, total=11355 Charoen-like_virus_isolate_Rio_segment_S_plus_segment_M_plus_segment_L_Nucle:1-12099

0 2000 4000 6000 8000 10000 12000

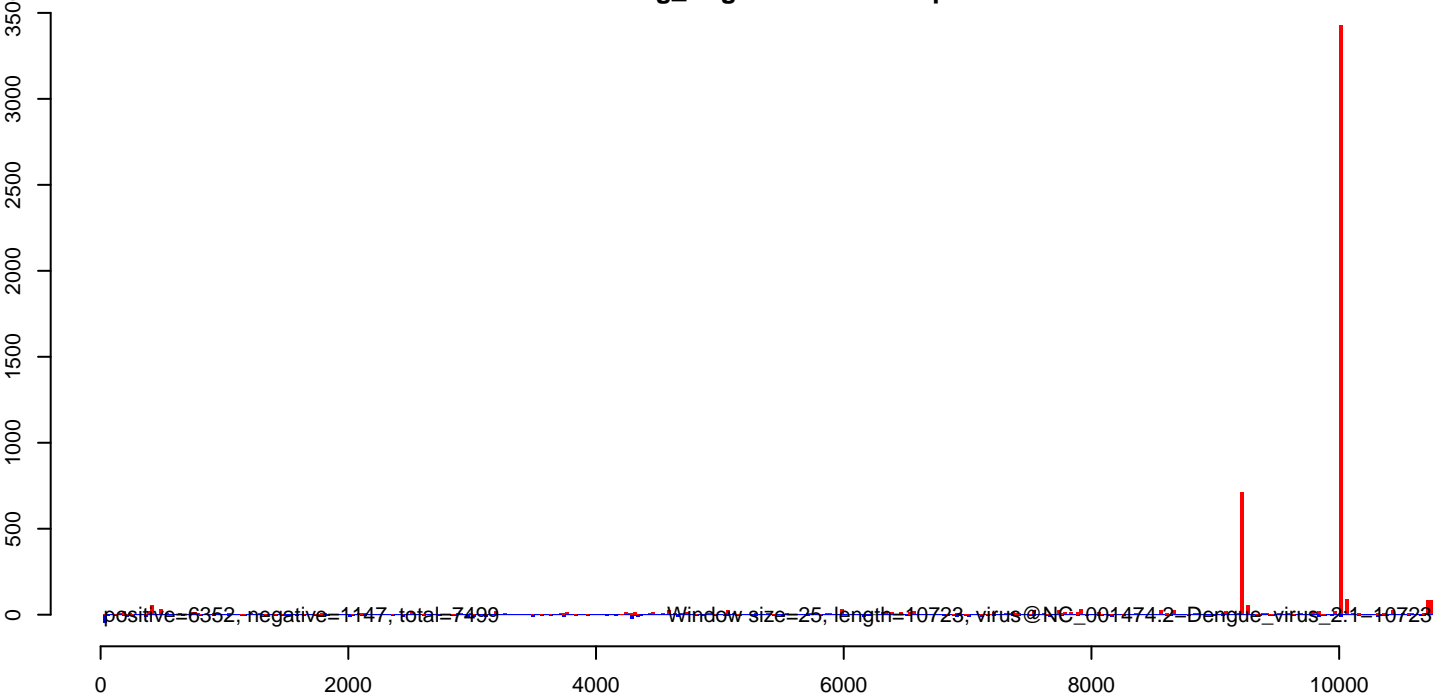
AeAeg_Aag2DENV2NGC.18_23.rep



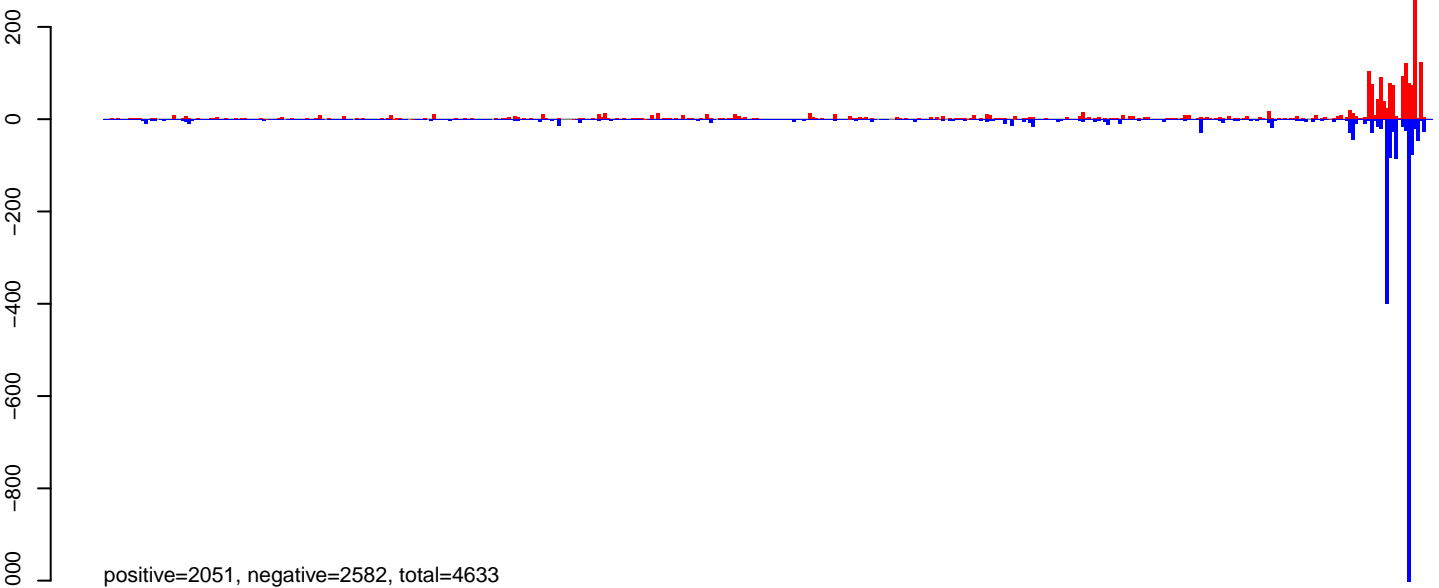
AeAeg_Aag2DENV2NGC.24_35.rep



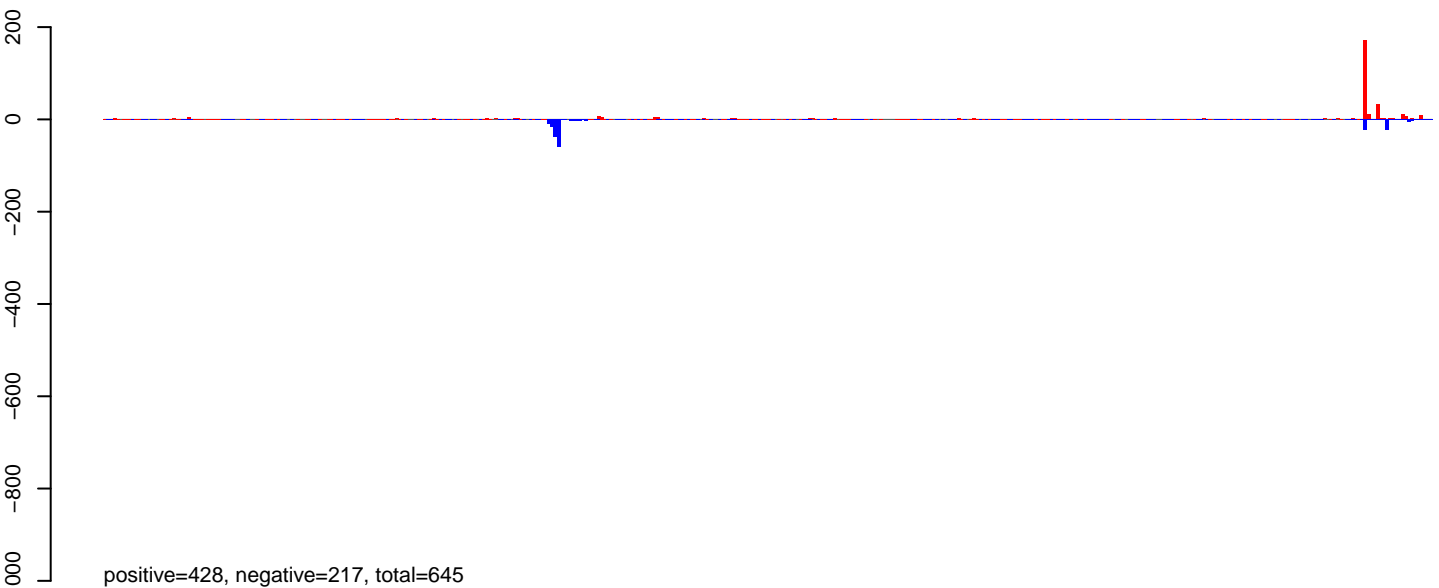
AeAeg_Aag2DENV2NGC.rep



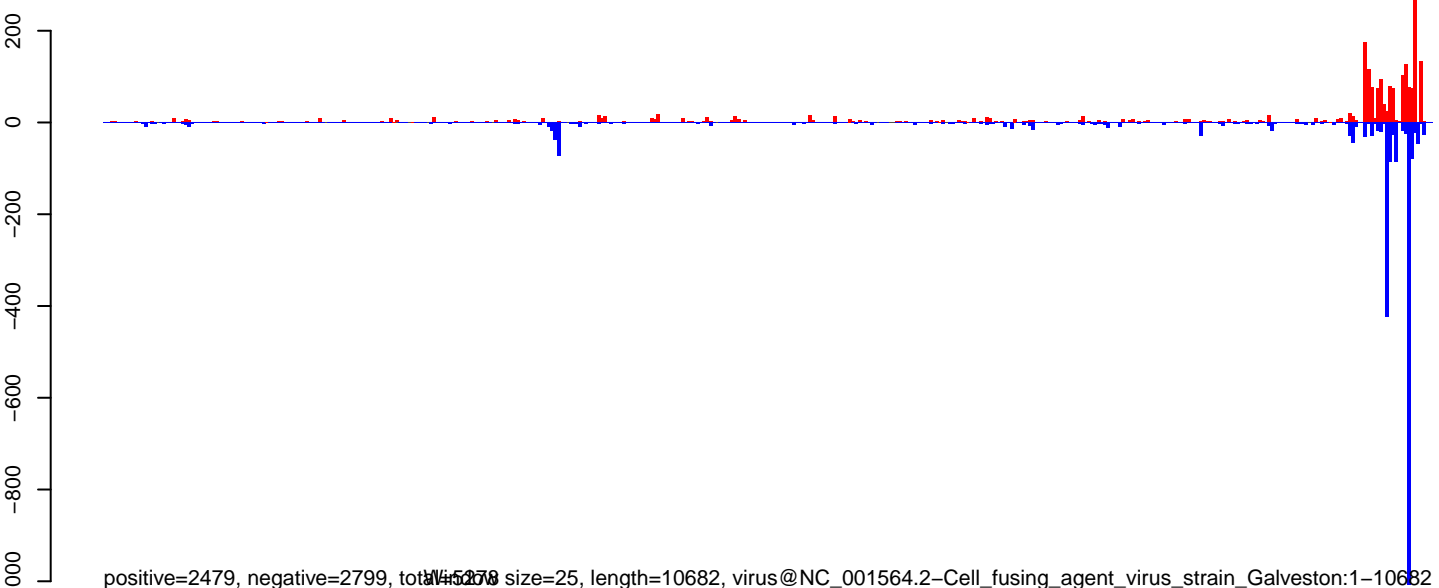
AeAeg_Aag2DENV2NGC.18_23.rep



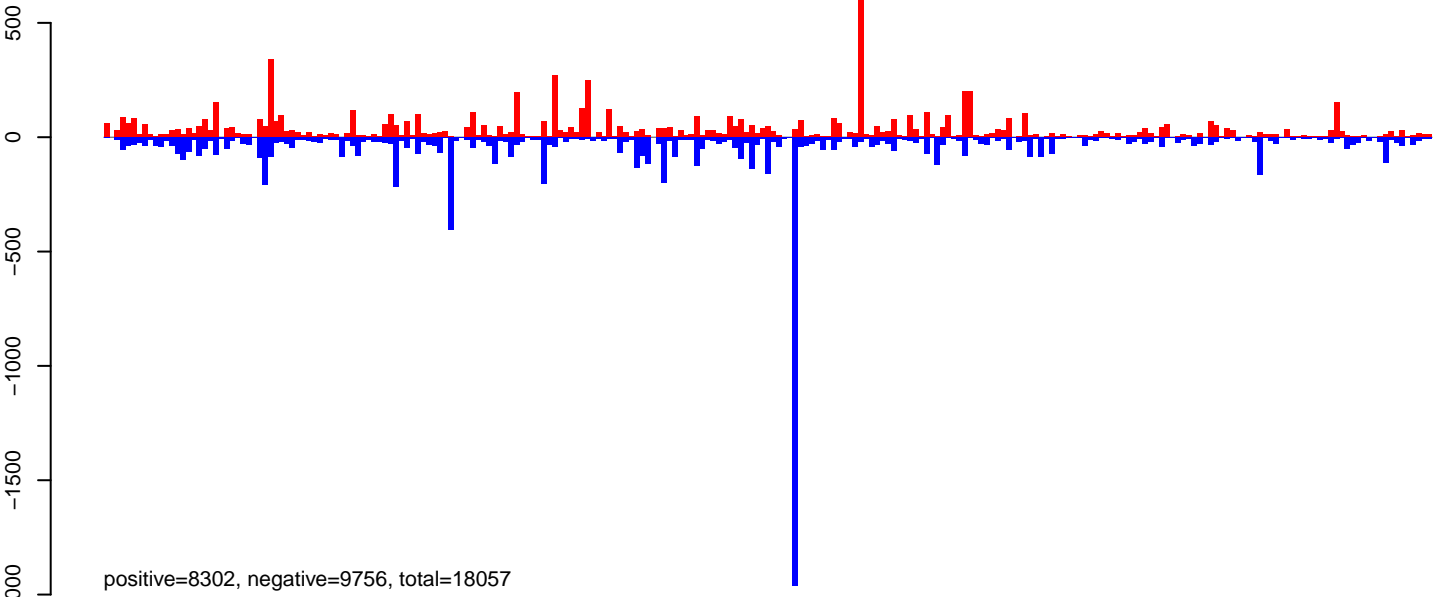
AeAeg_Aag2DENV2NGC.24_35.rep



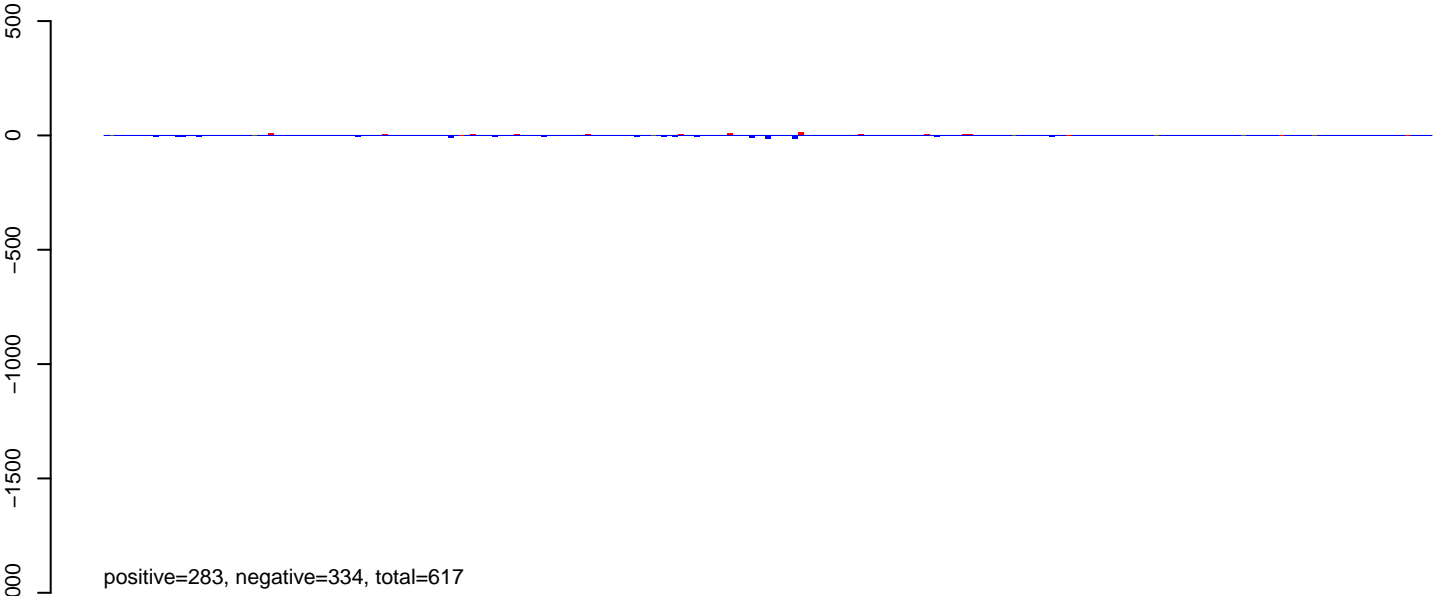
AeAeg_Aag2DENV2NGC.rep



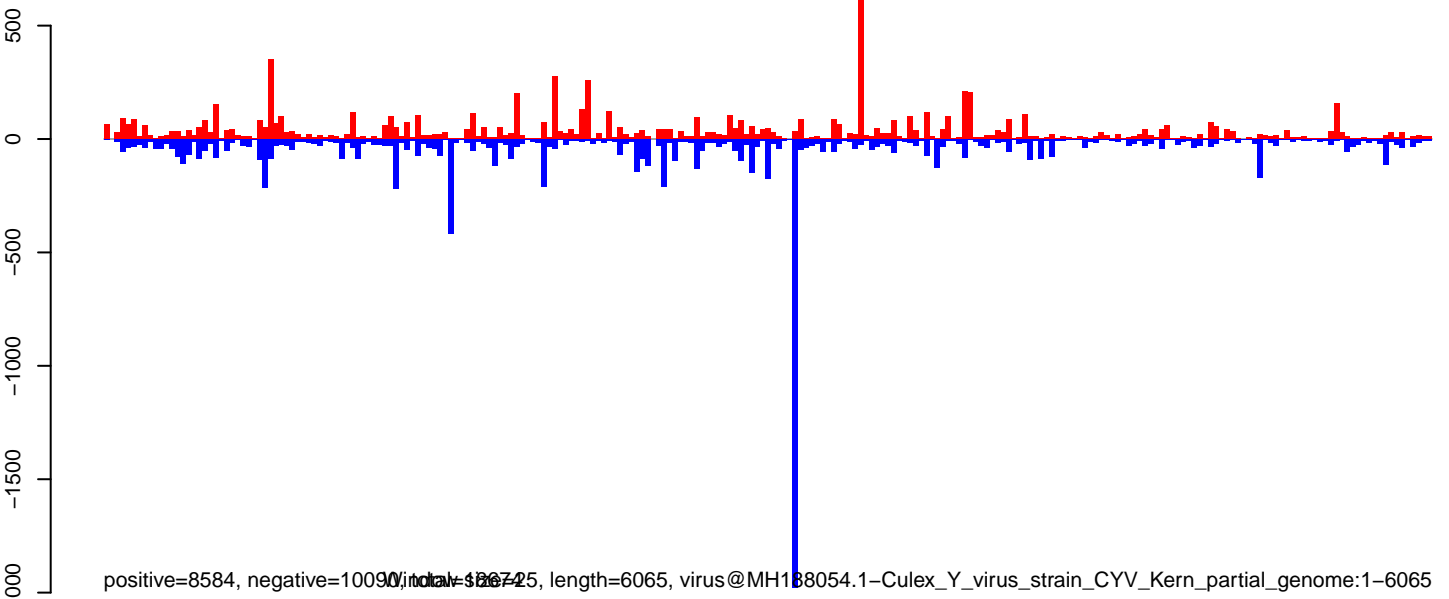
AeAeg_Aag2DENV2NGC.18_23.rep



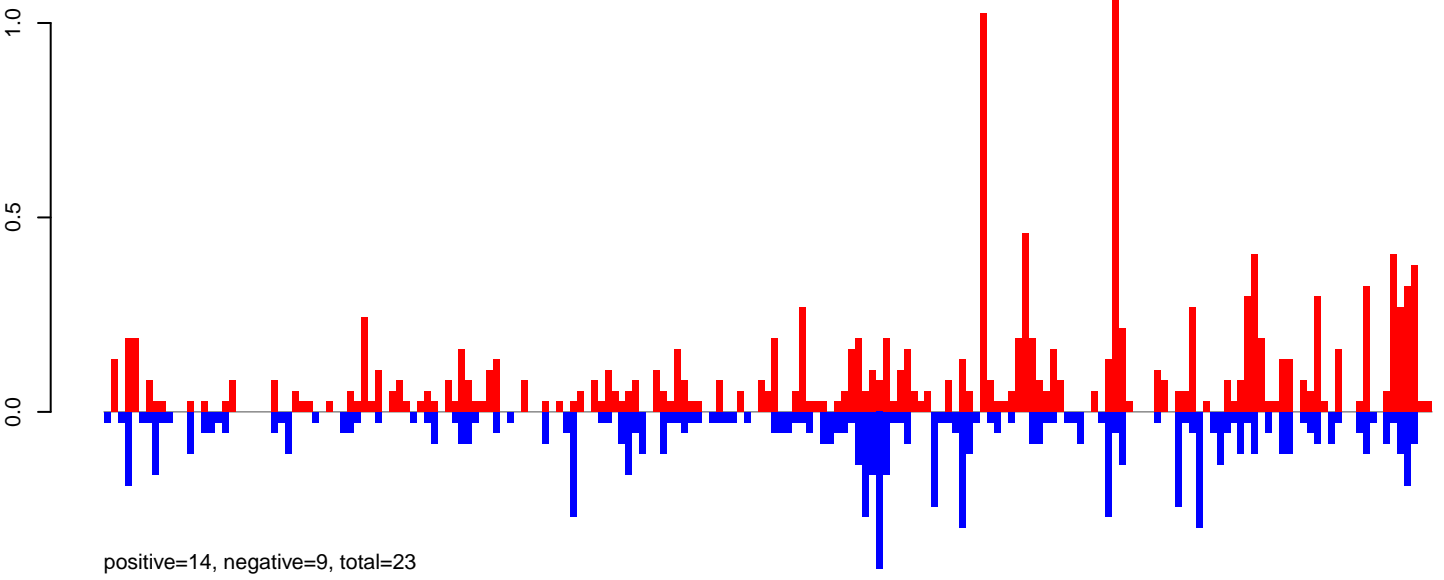
AeAeg_Aag2DENV2NGC.24_35.rep



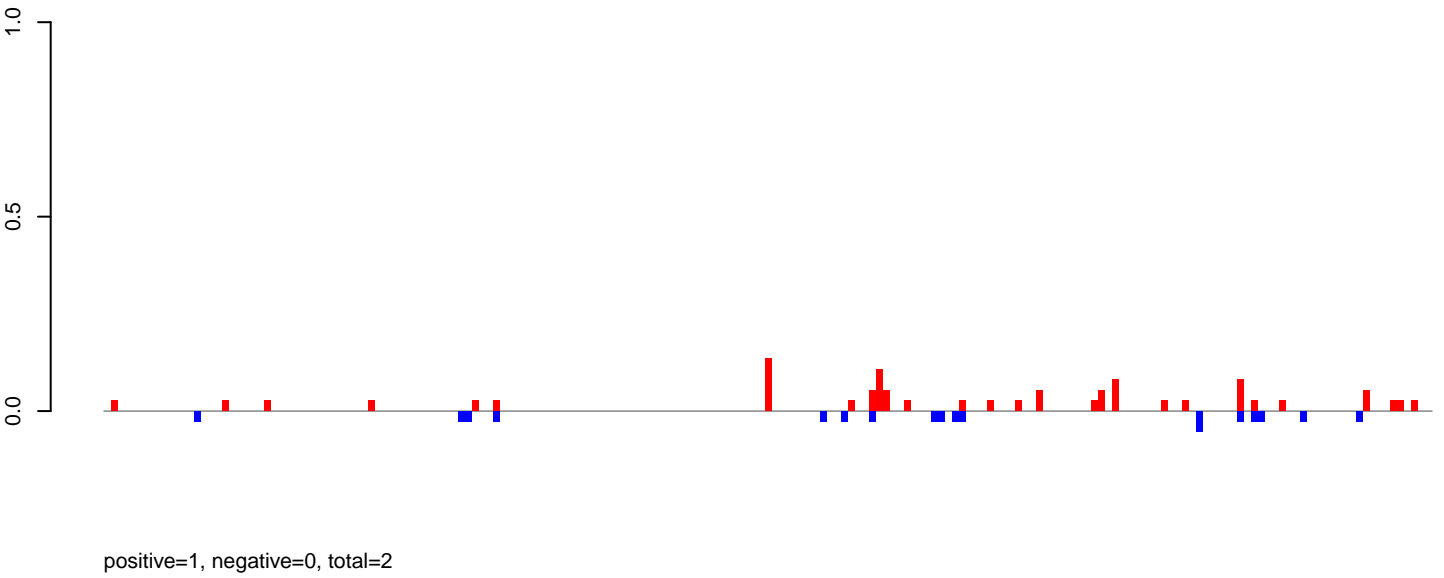
AeAeg_Aag2DENV2NGC.rep



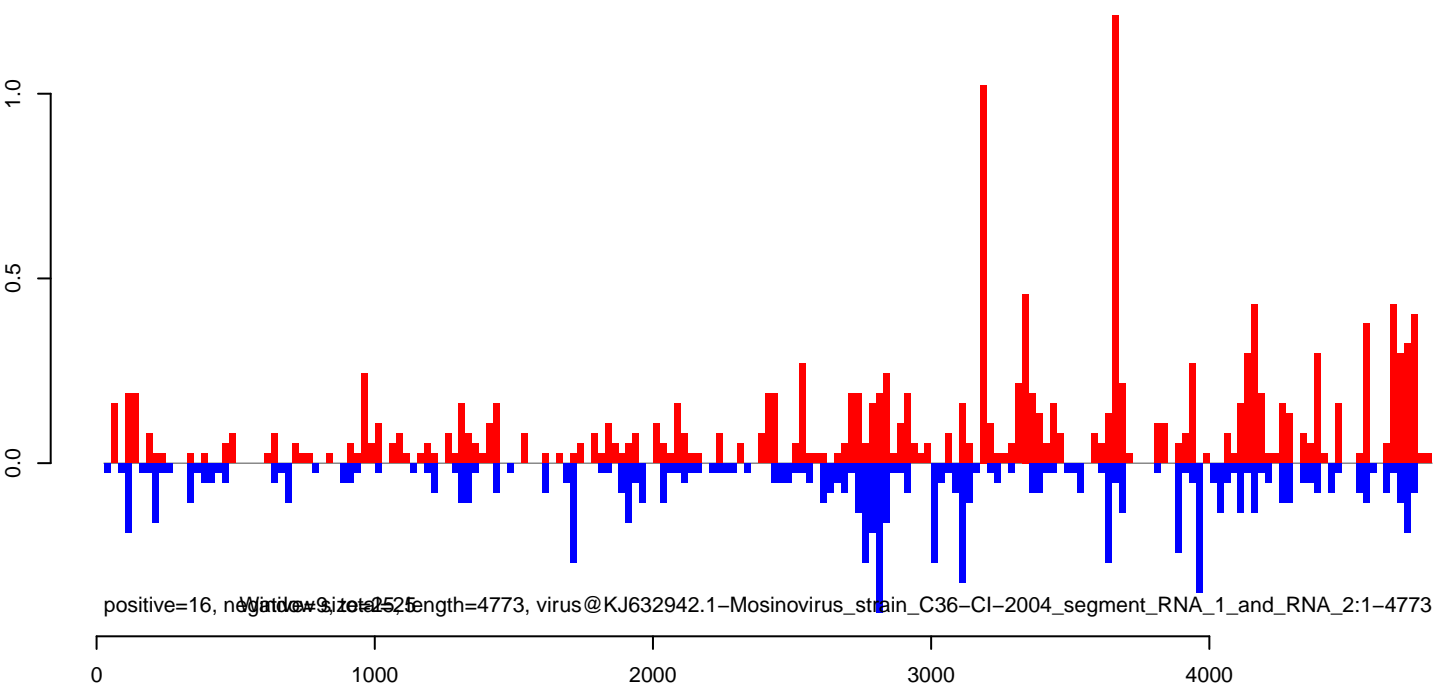
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

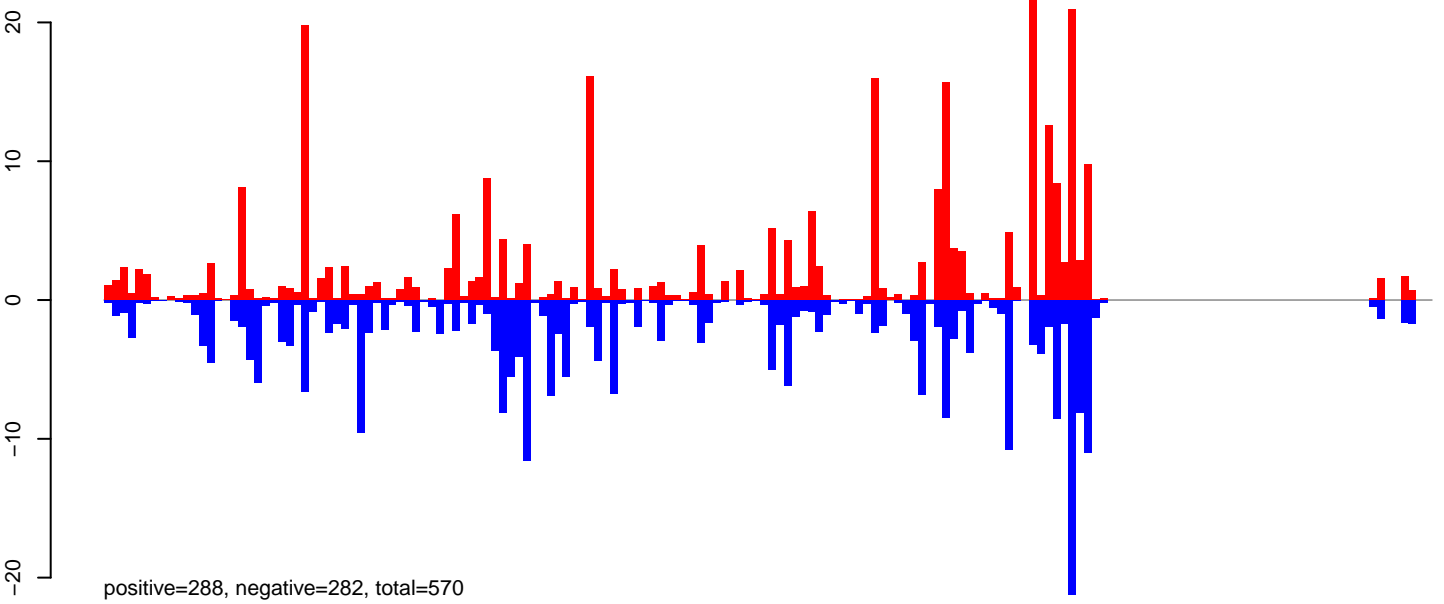


AeAeg_Aag2DENV2NGC.rep

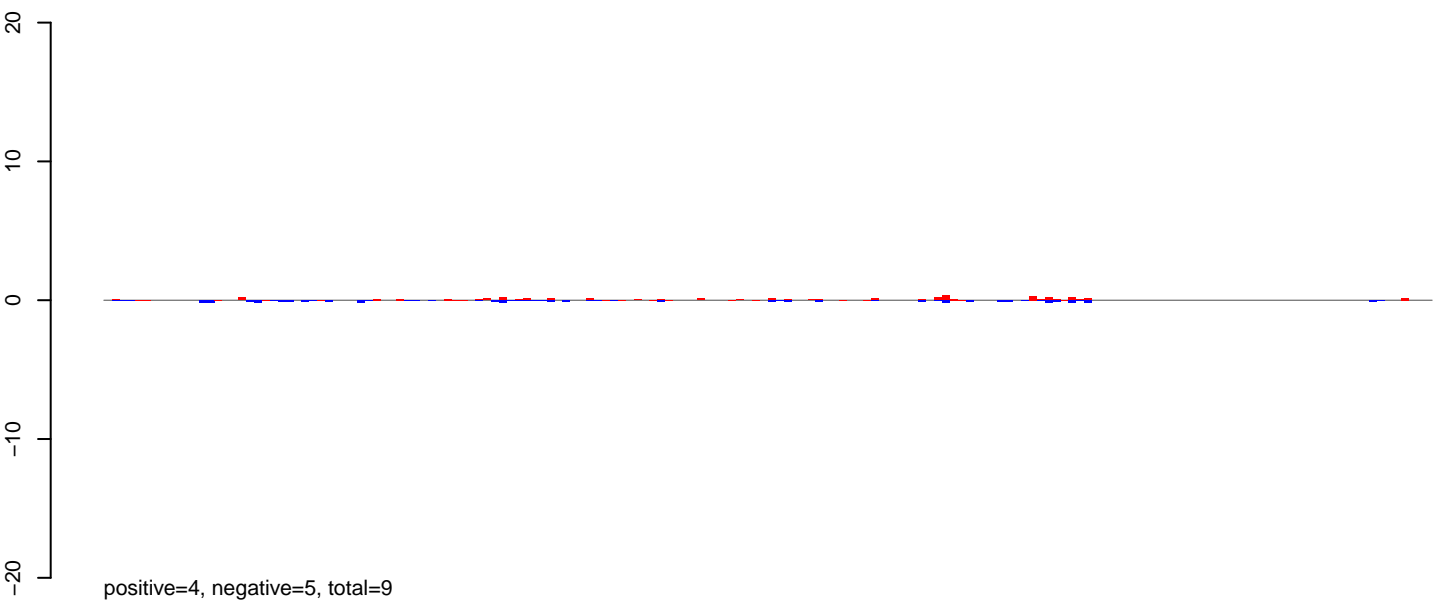


Window size=25, length=4773, virus@KJ632942.1-Mosinovirus_strain_C36-CI-2004_segment_RNA_1_and_RNA_2:1-4773

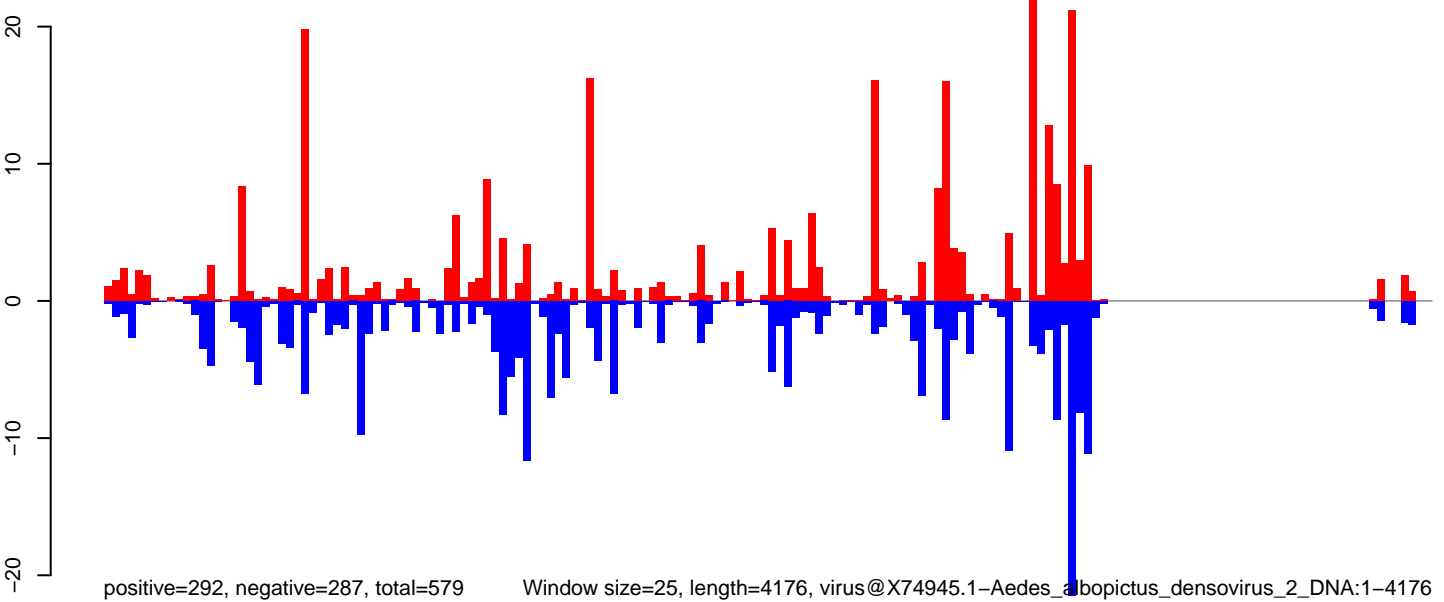
AeAeg_Aag2DENV2NGC.18_23.rep



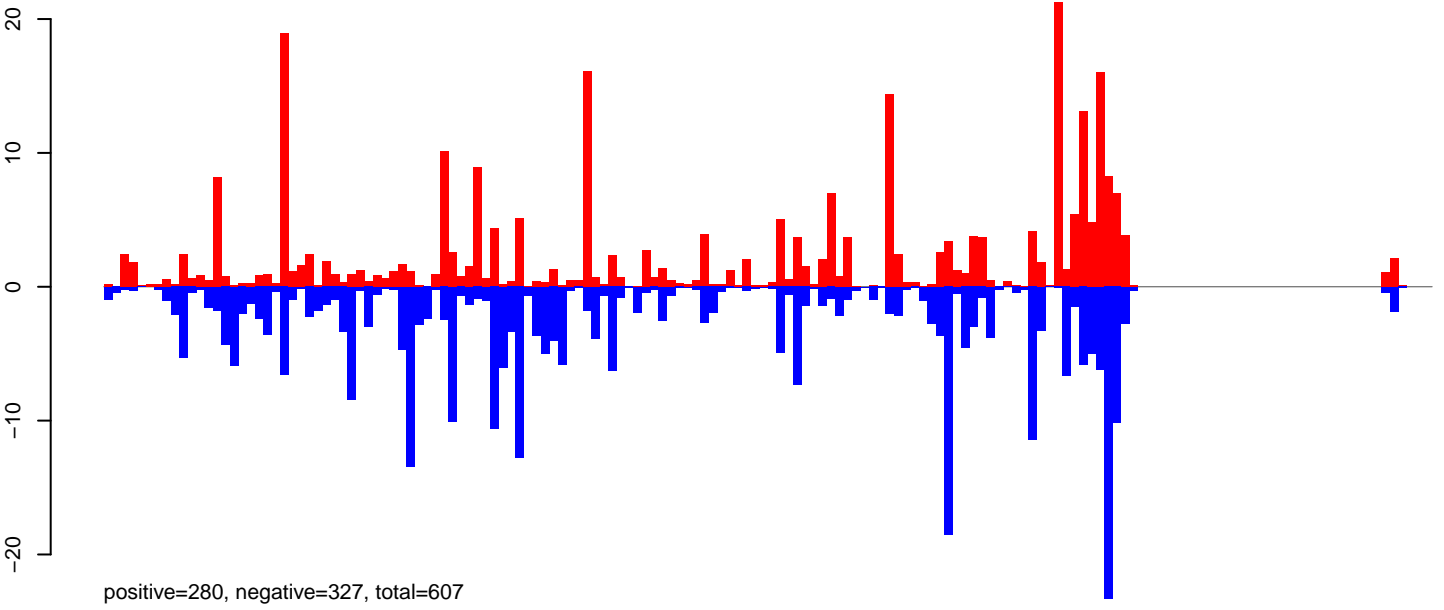
AeAeg_Aag2DENV2NGC.24_35.rep



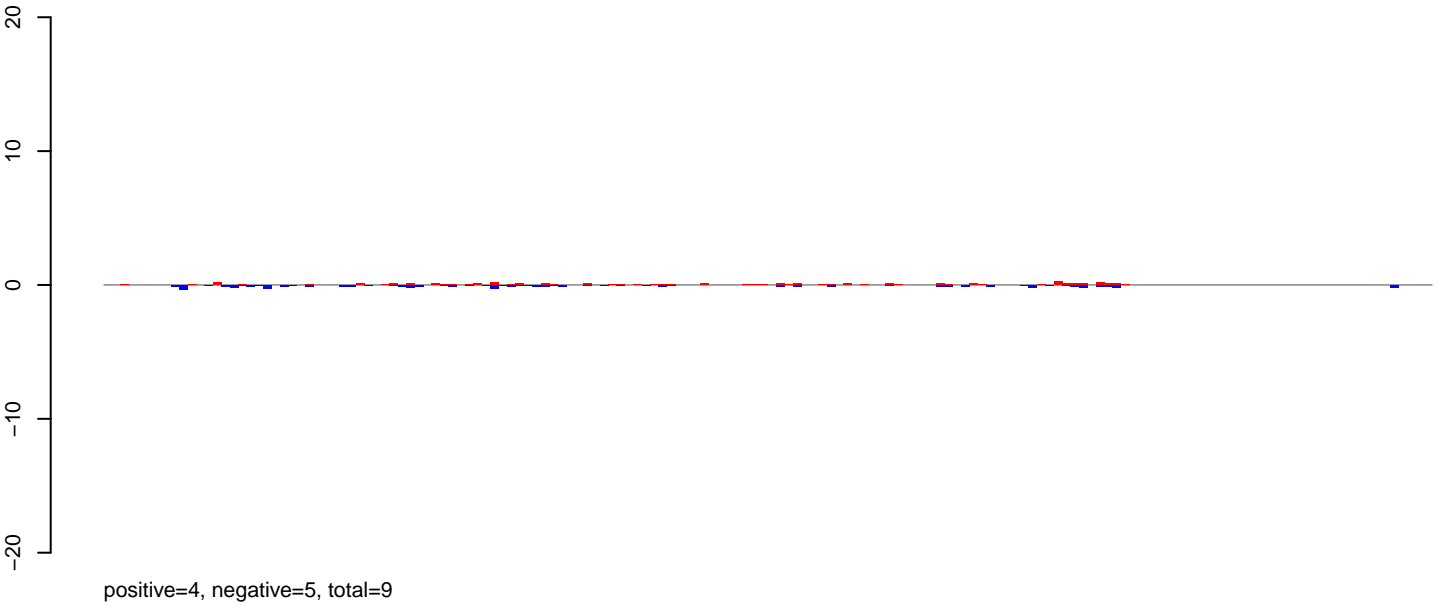
AeAeg_Aag2DENV2NGC.rep



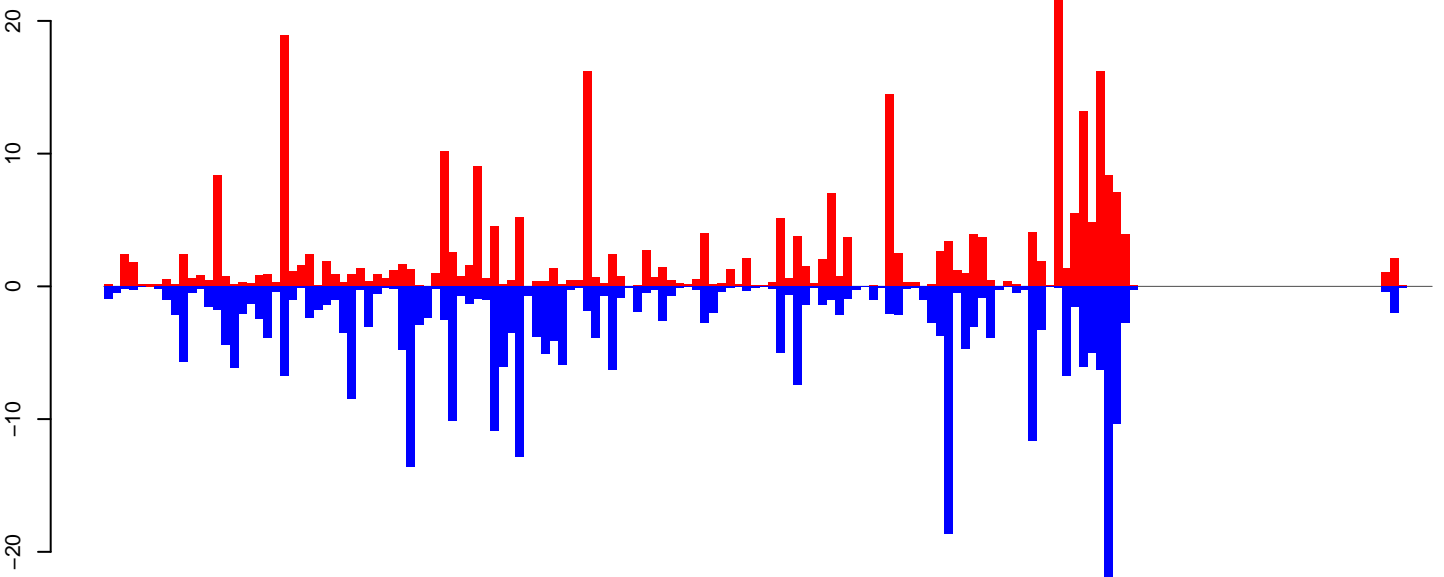
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep

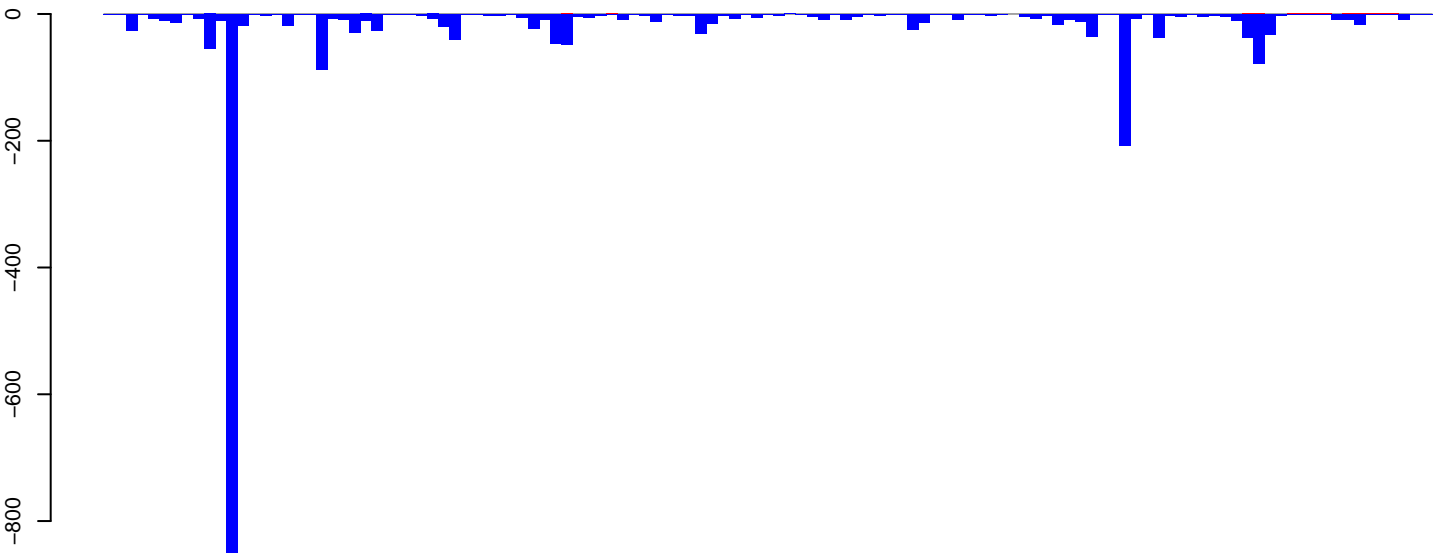


AeAeg_Aag2DENV2NGC.18_23.rep



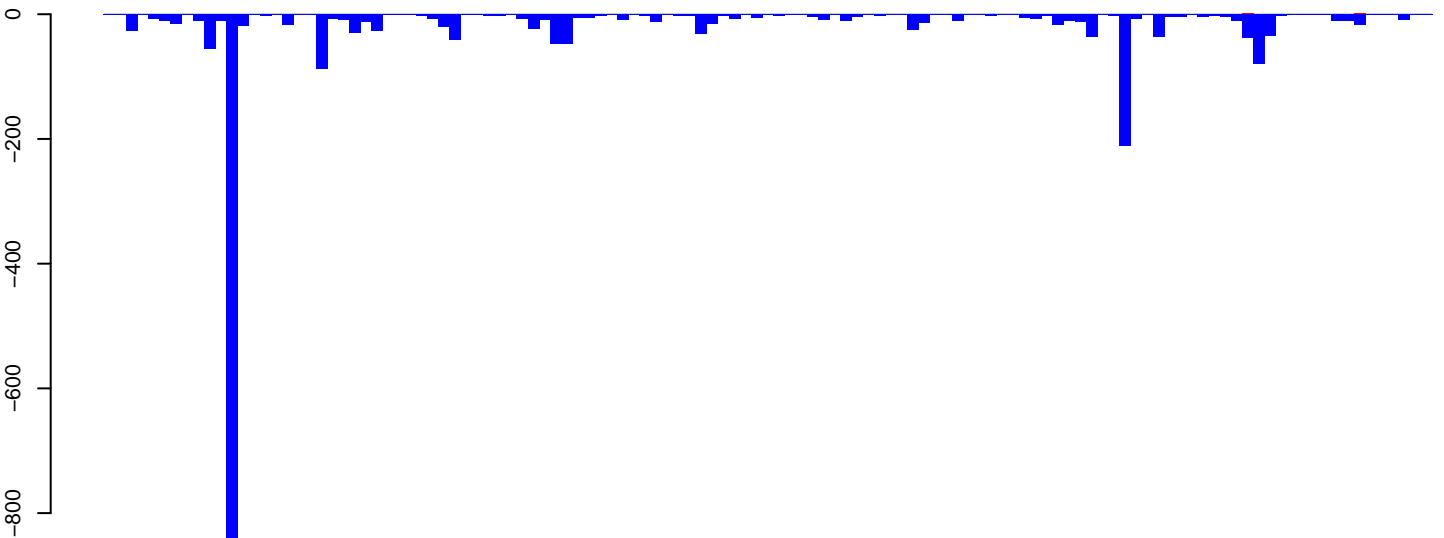
positive=1, negative=50, total=51

AeAeg_Aag2DENV2NGC.24_35.rep



positive=4, negative=2249, total=2252

AeAeg_Aag2DENV2NGC.rep

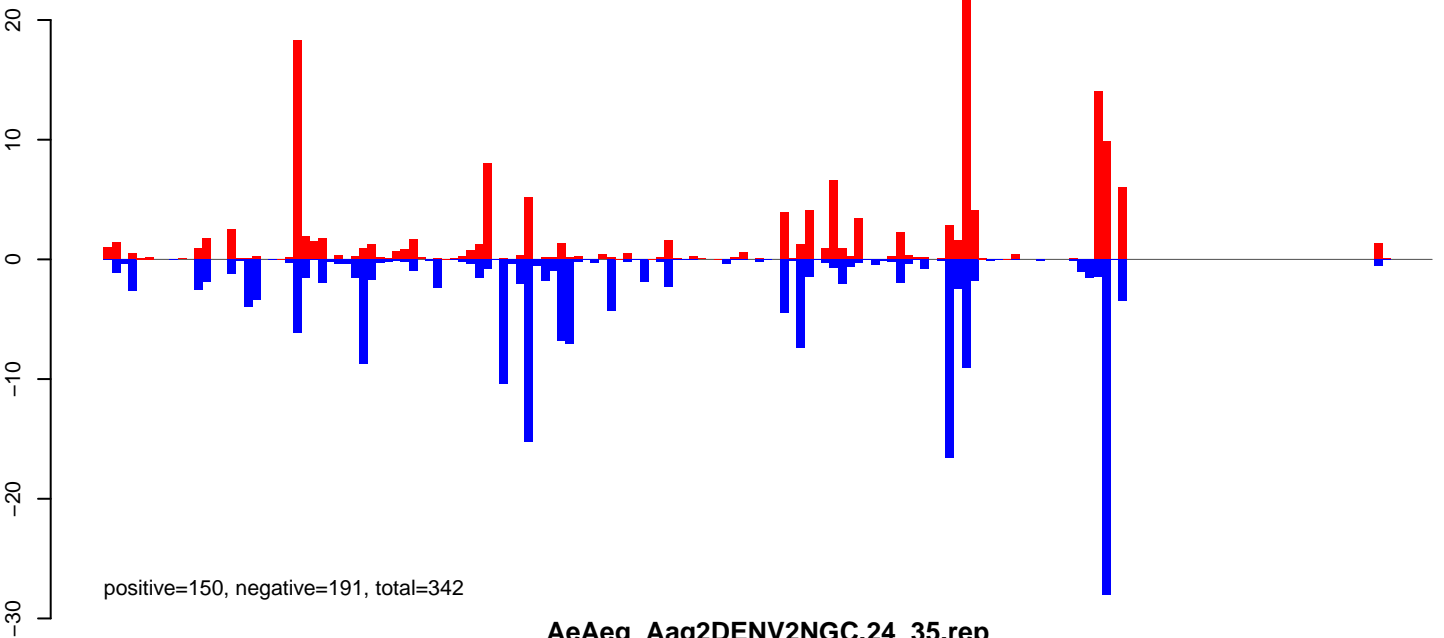


positive=5, negative=2298, total=2303

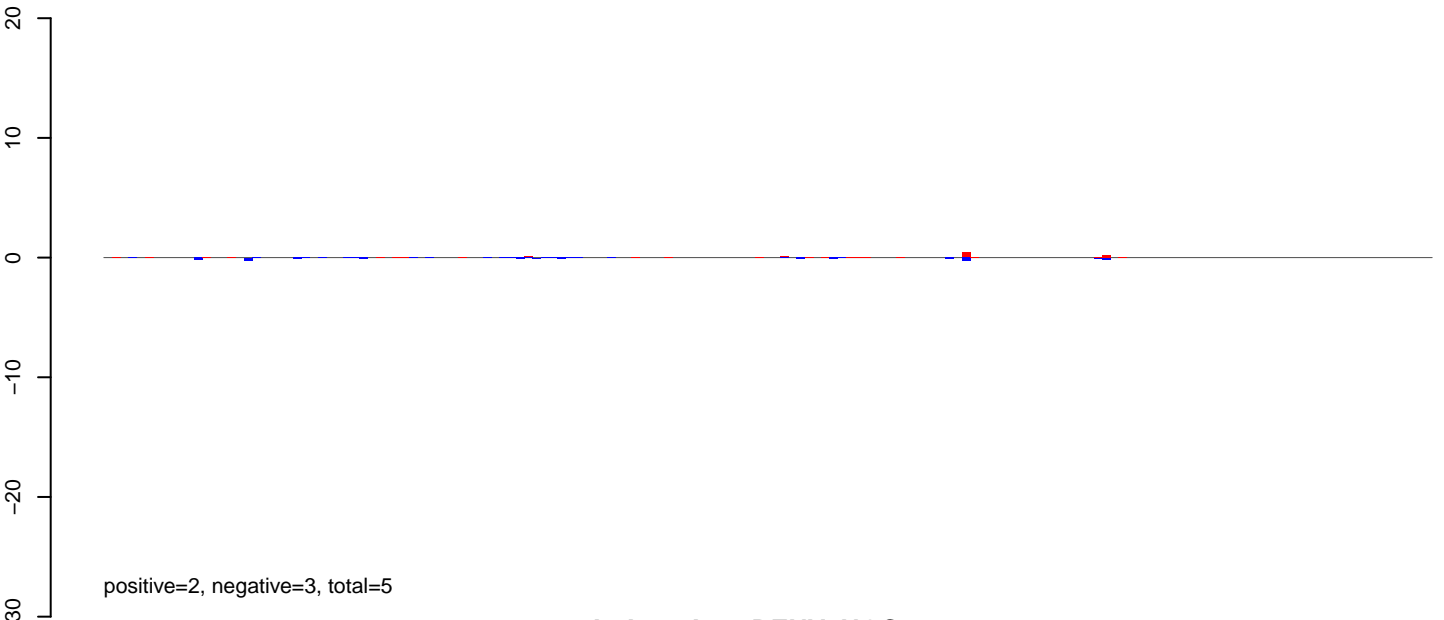
Window size=25, length=2965, virus@AY347953.2-Aedes_aegypti_genomic_sequence:1-2965

0 500 1000 1500 2000 2500 3000

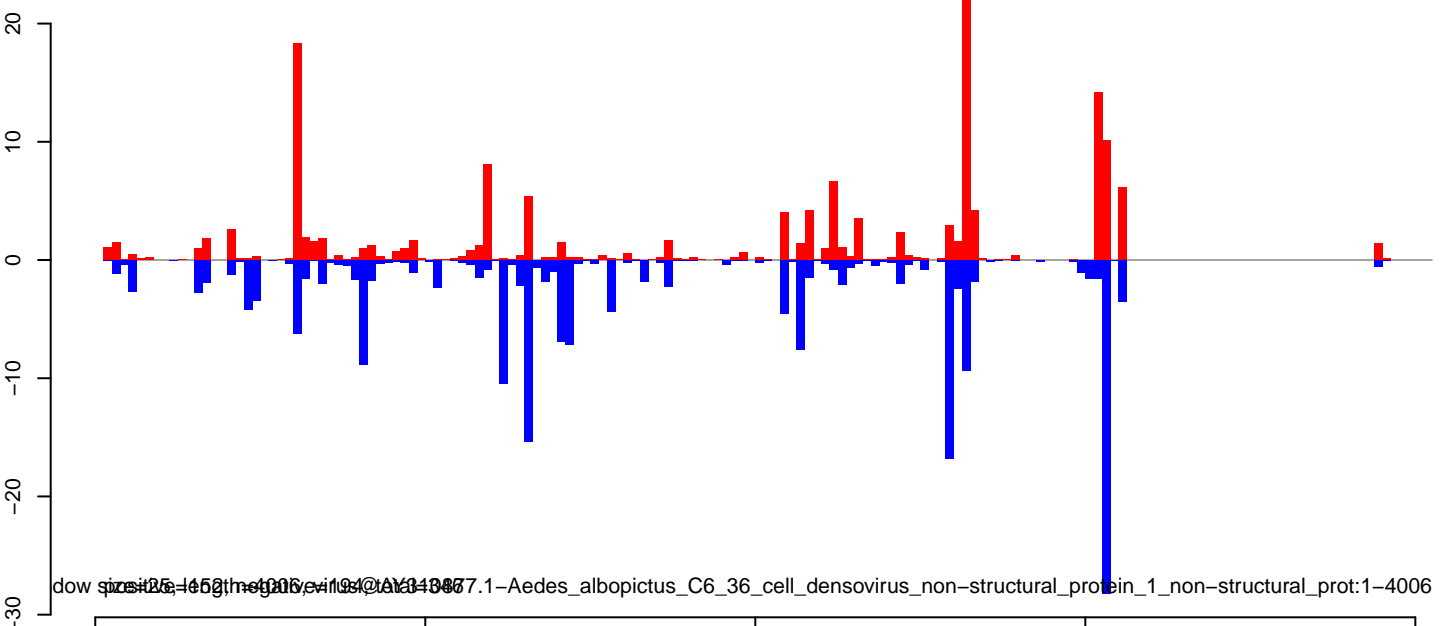
AeAeg_Aag2DENV2NGC.18_23.rep



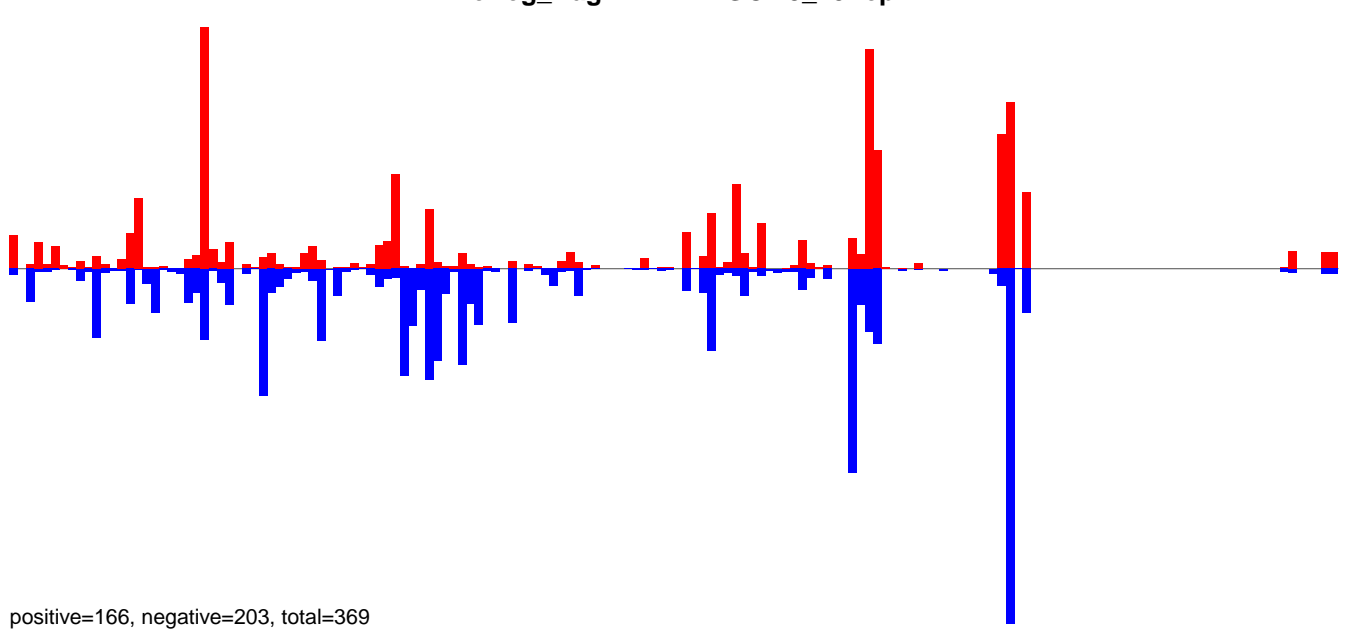
AeAeg_Aag2DENV2NGC.24_35.rep



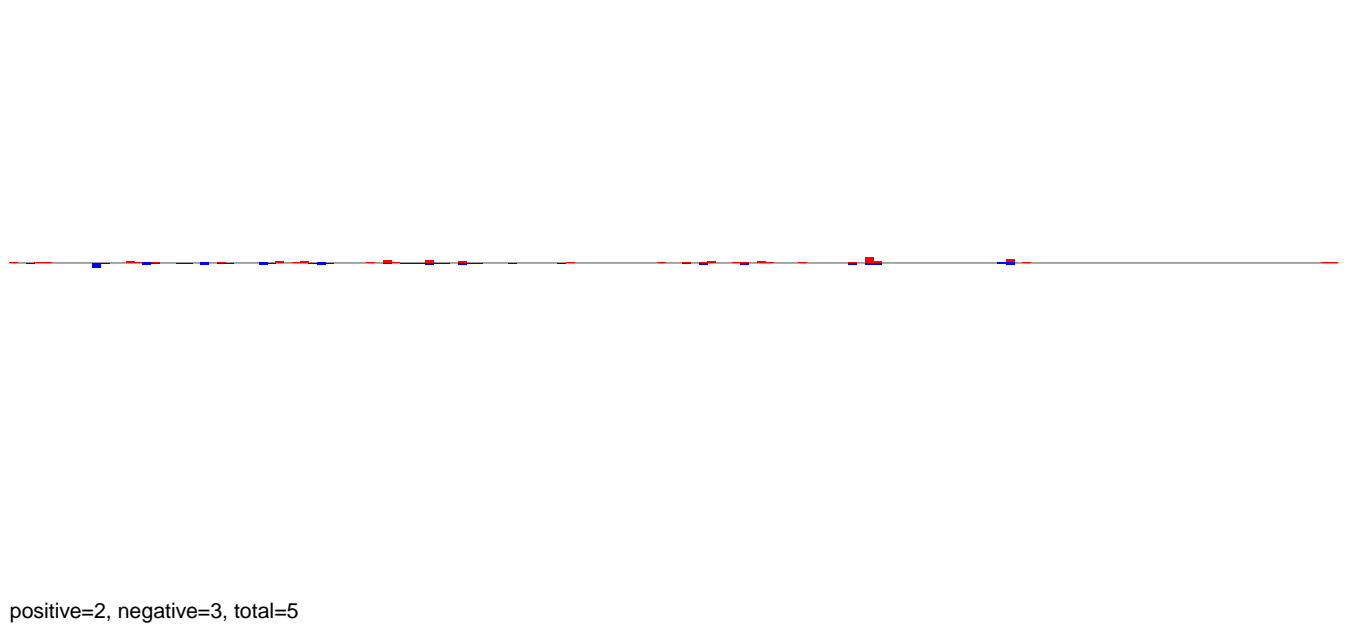
AeAeg_Aag2DENV2NGC.rep



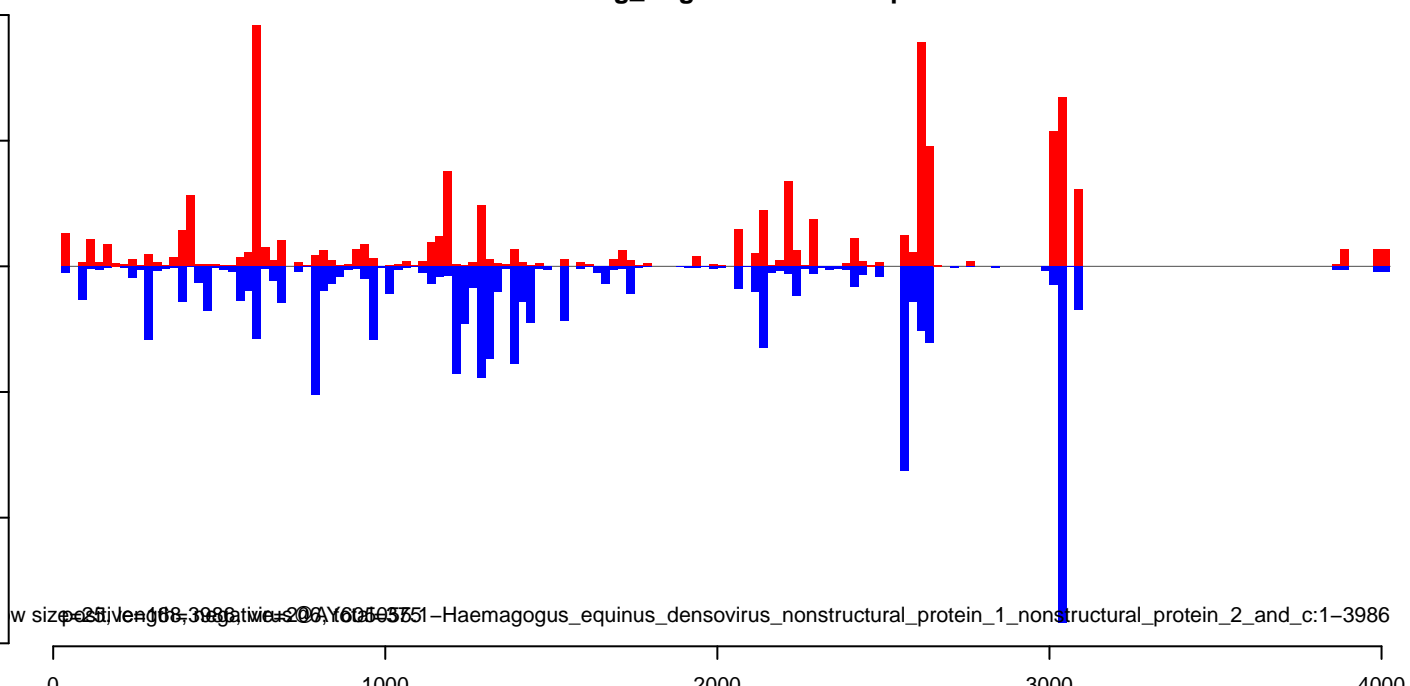
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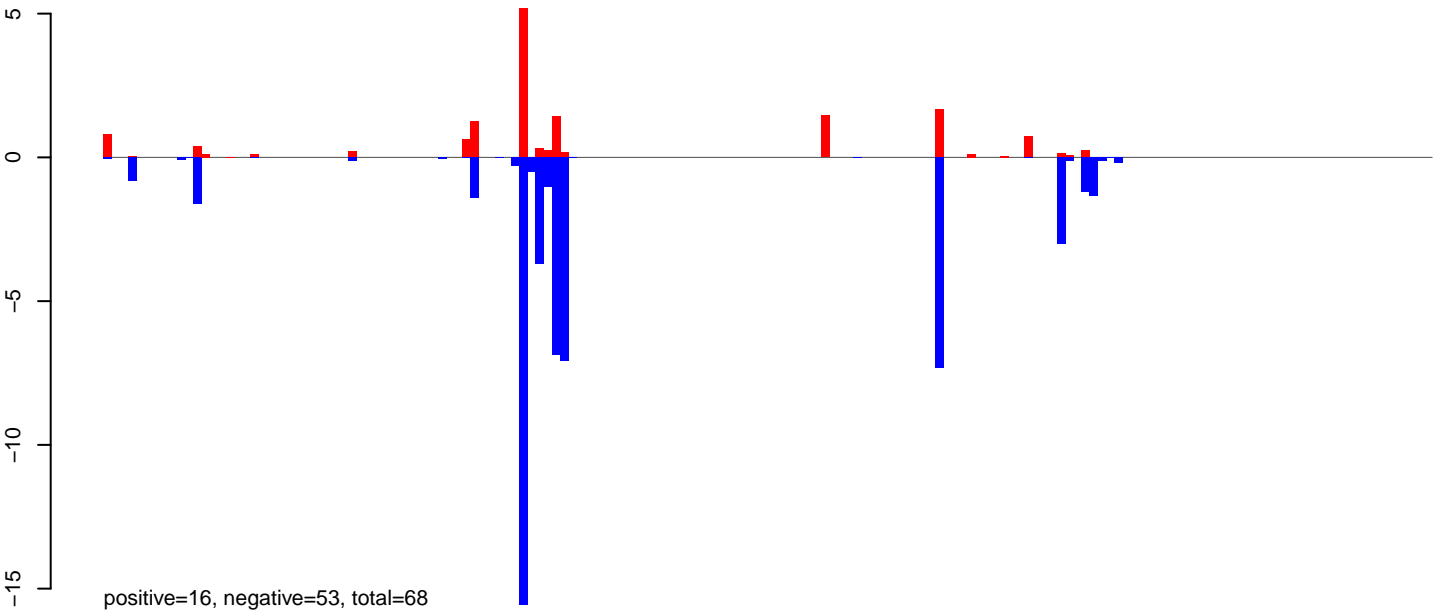
AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



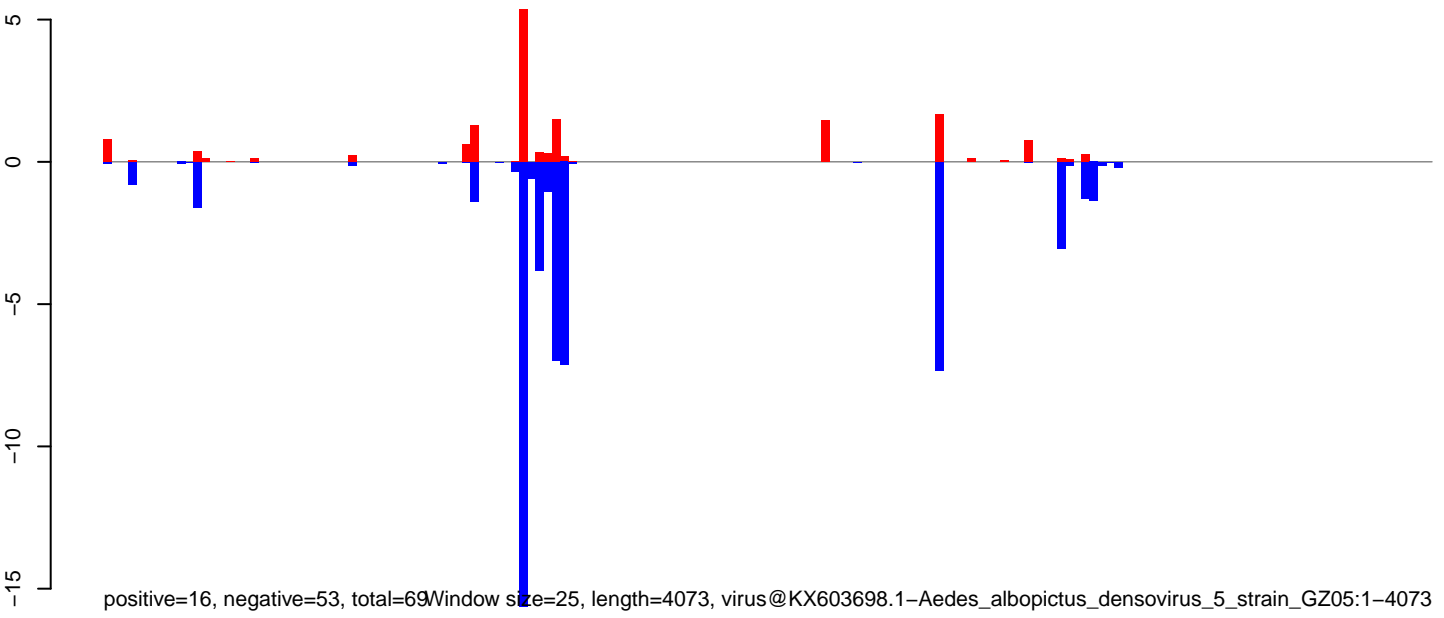
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

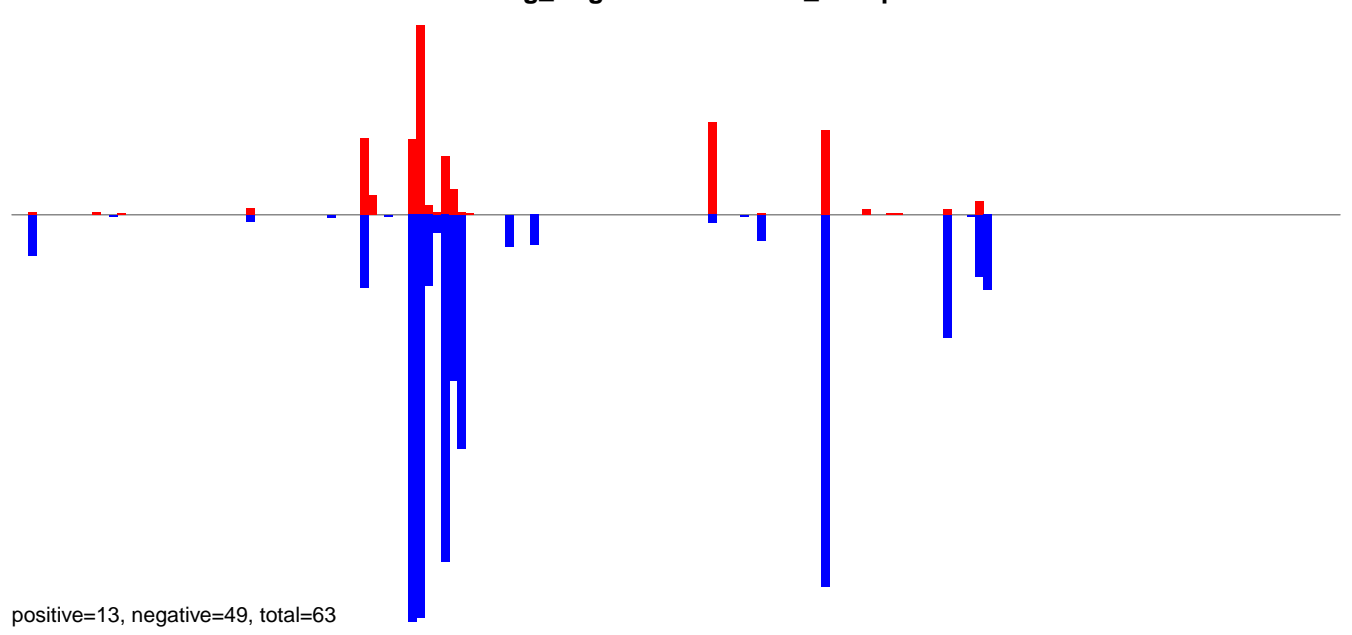


AeAeg_Aag2DENV2NGC.rep

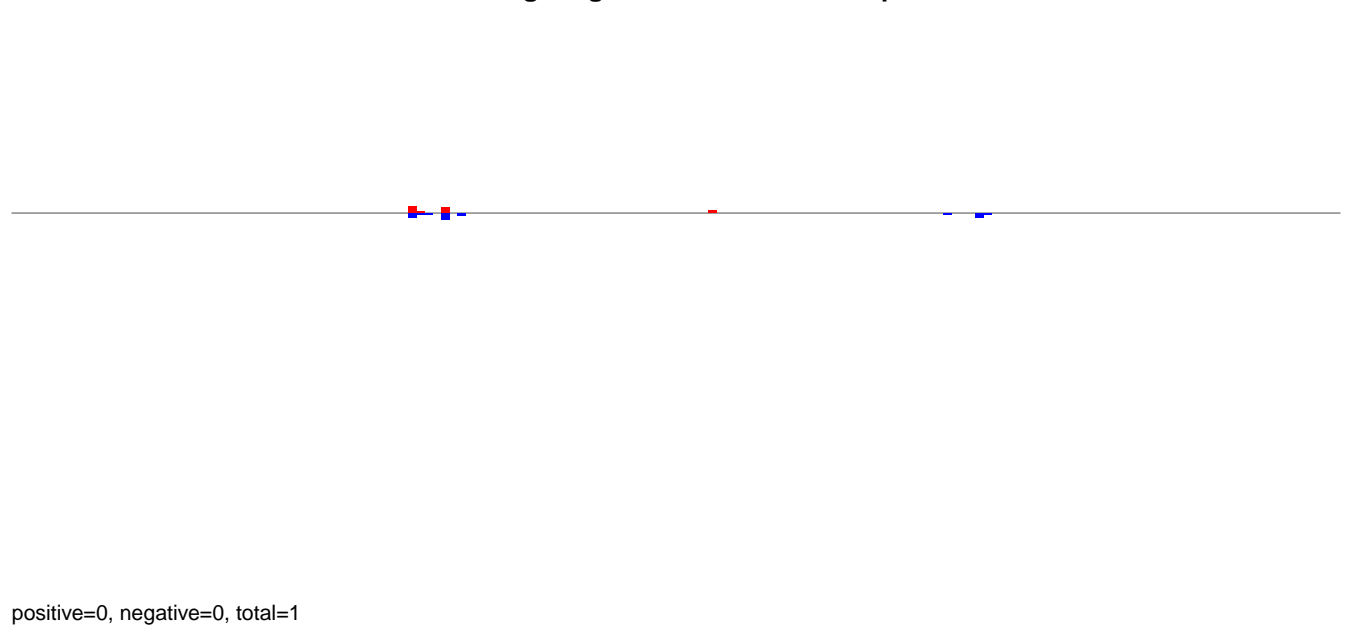


0 1000 2000 3000 4000

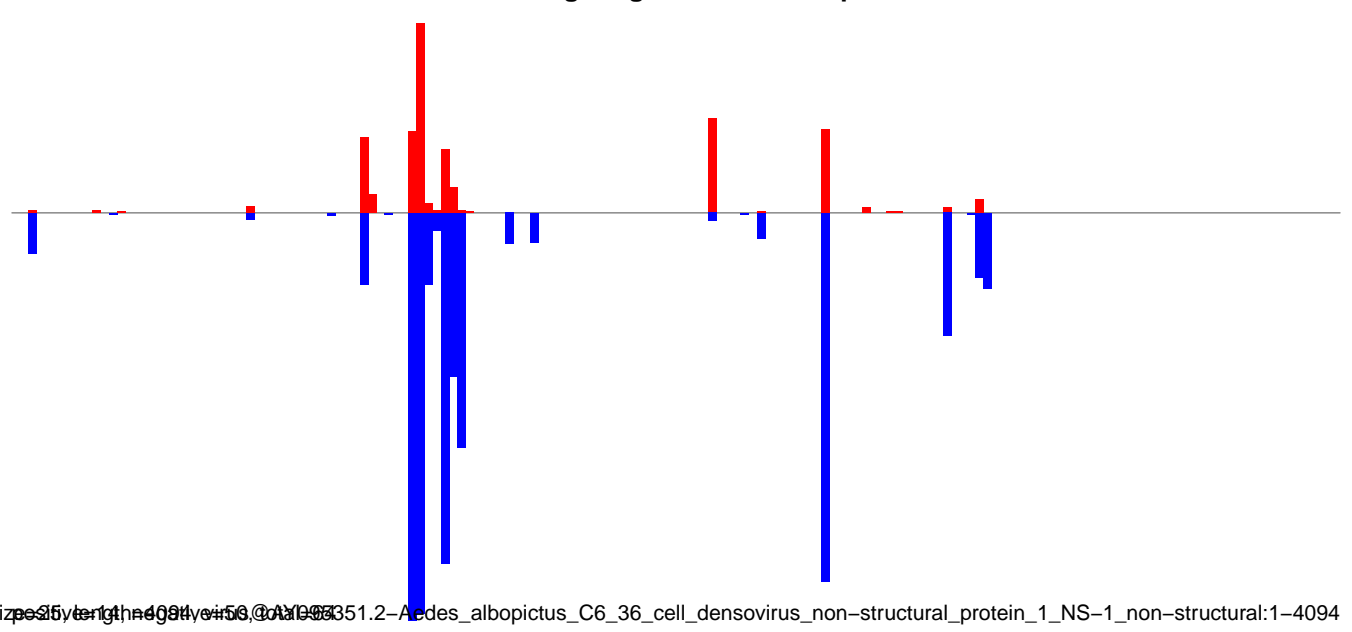
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

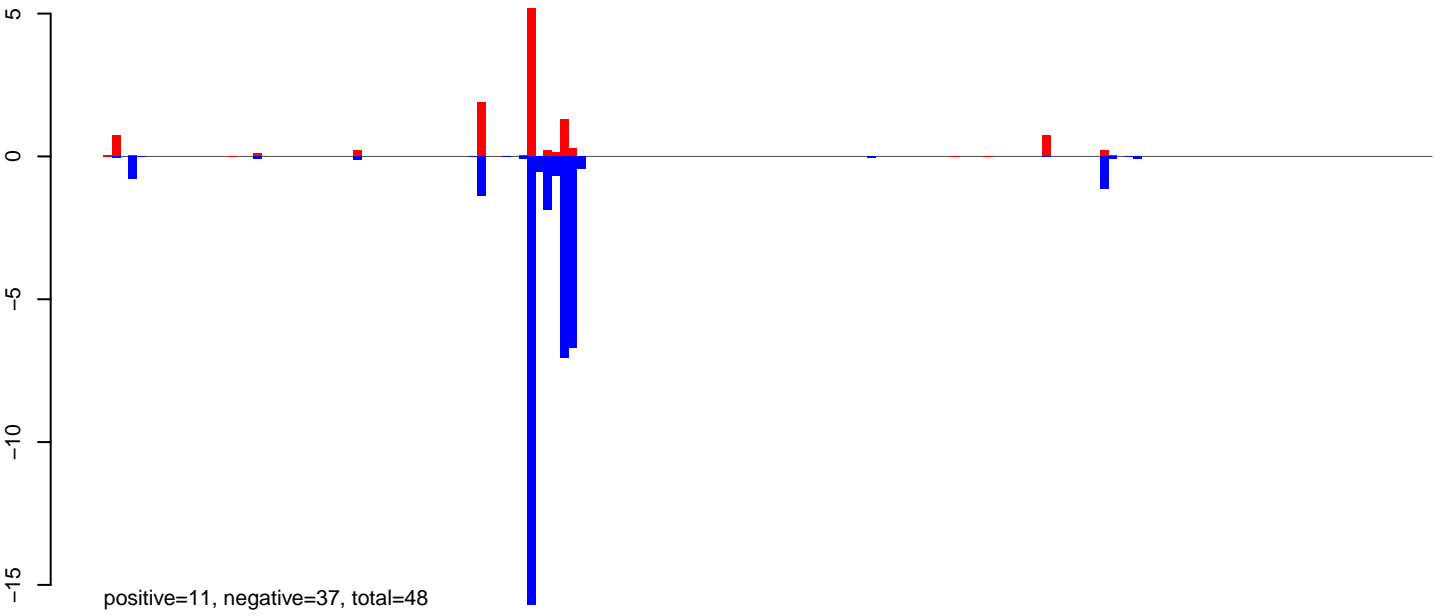


AeAeg_Aag2DENV2NGC.rep



bow size=15, length=4094, positive=56, negative=56, total=112
Aedes albopictus_C6_36_cell_densovirus_non-structural_protein_1_NS-1_non-structural:1-4094

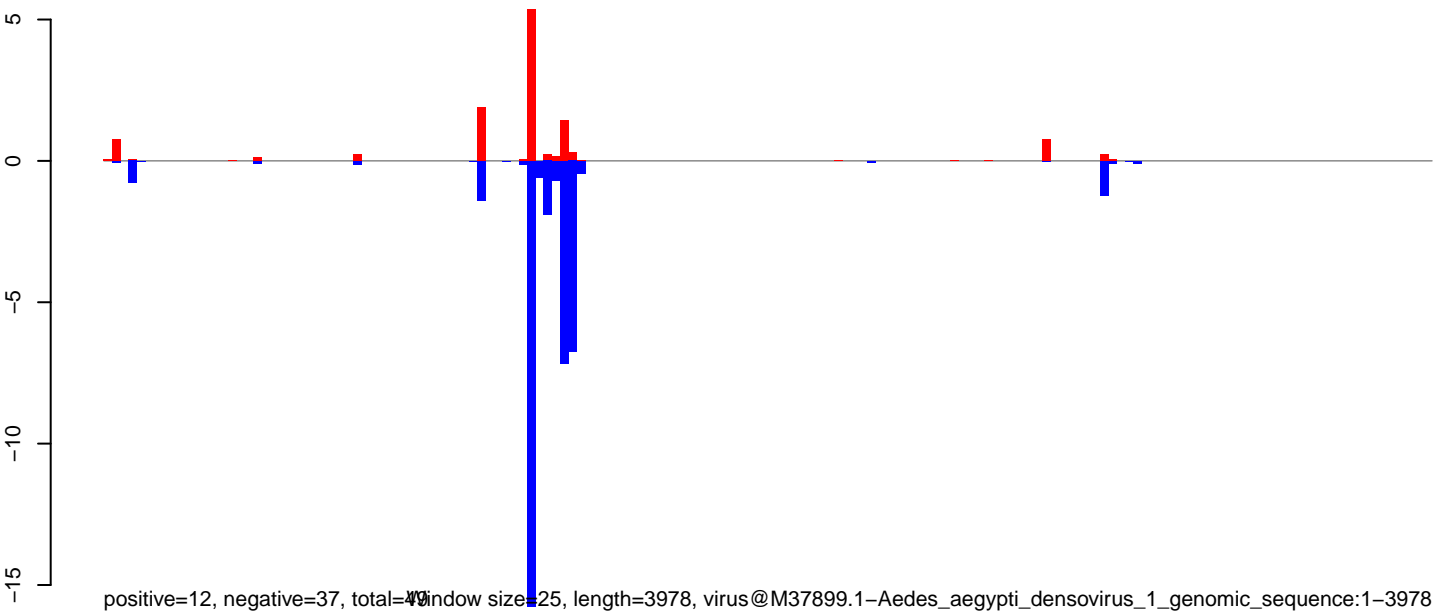
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

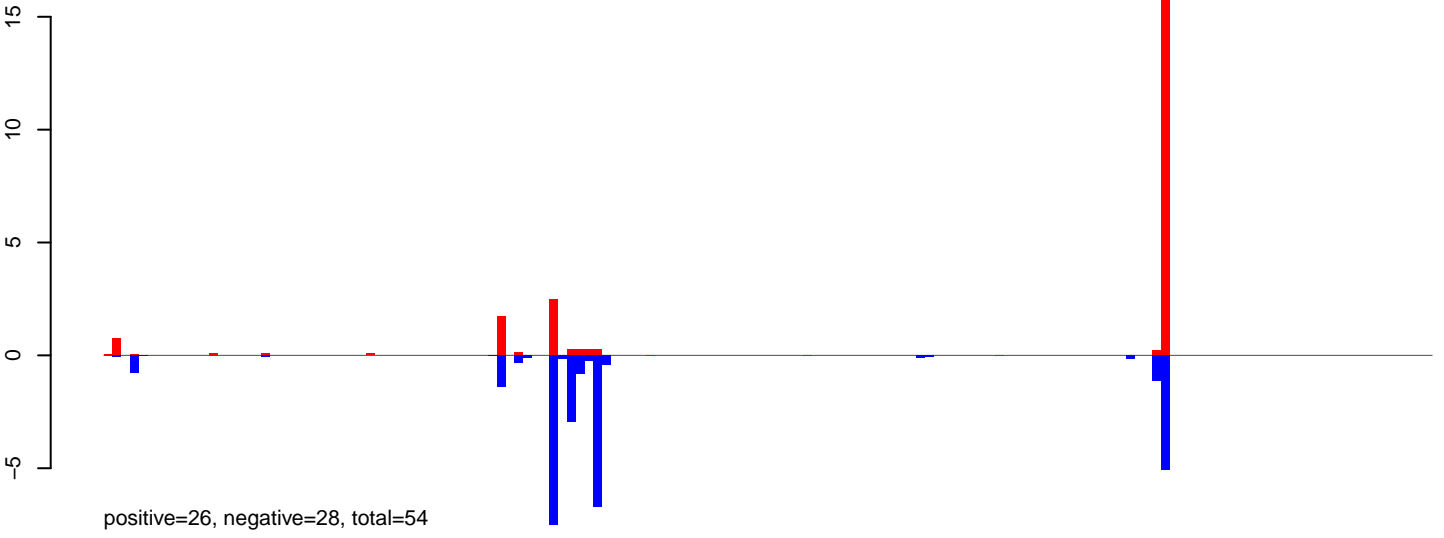


AeAeg_Aag2DENV2NGC.rep

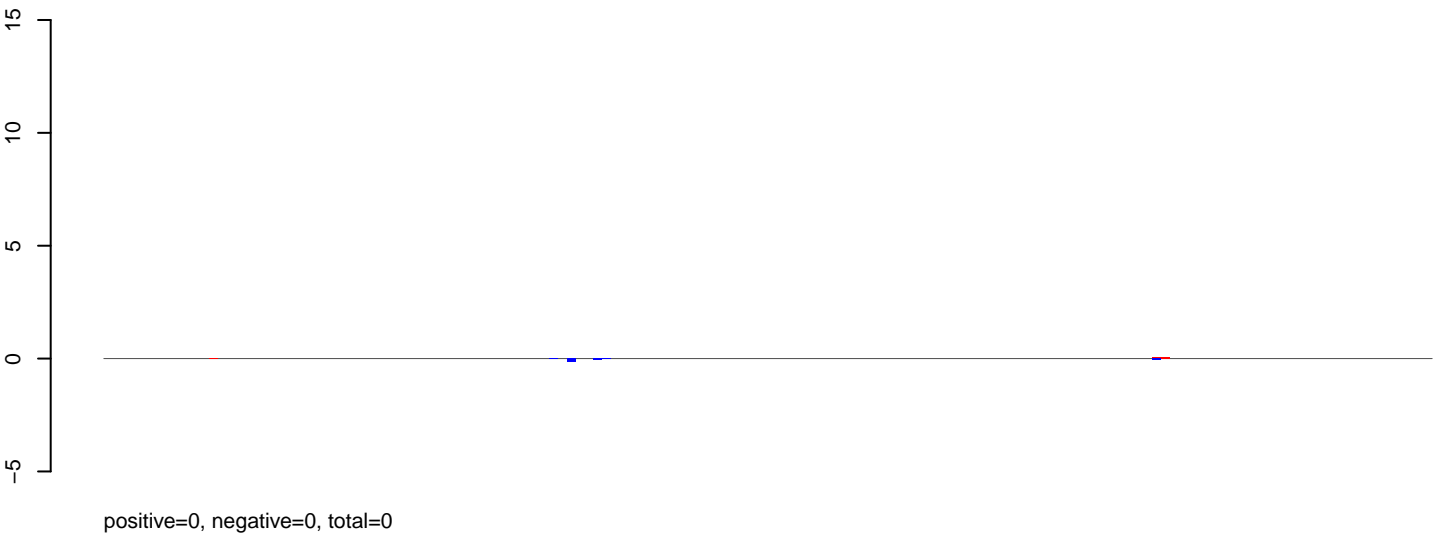


Window size=25, length=3978, virus@M37899.1-Aedes_aegypti_densovirus_1_genomic_sequence:1-3978

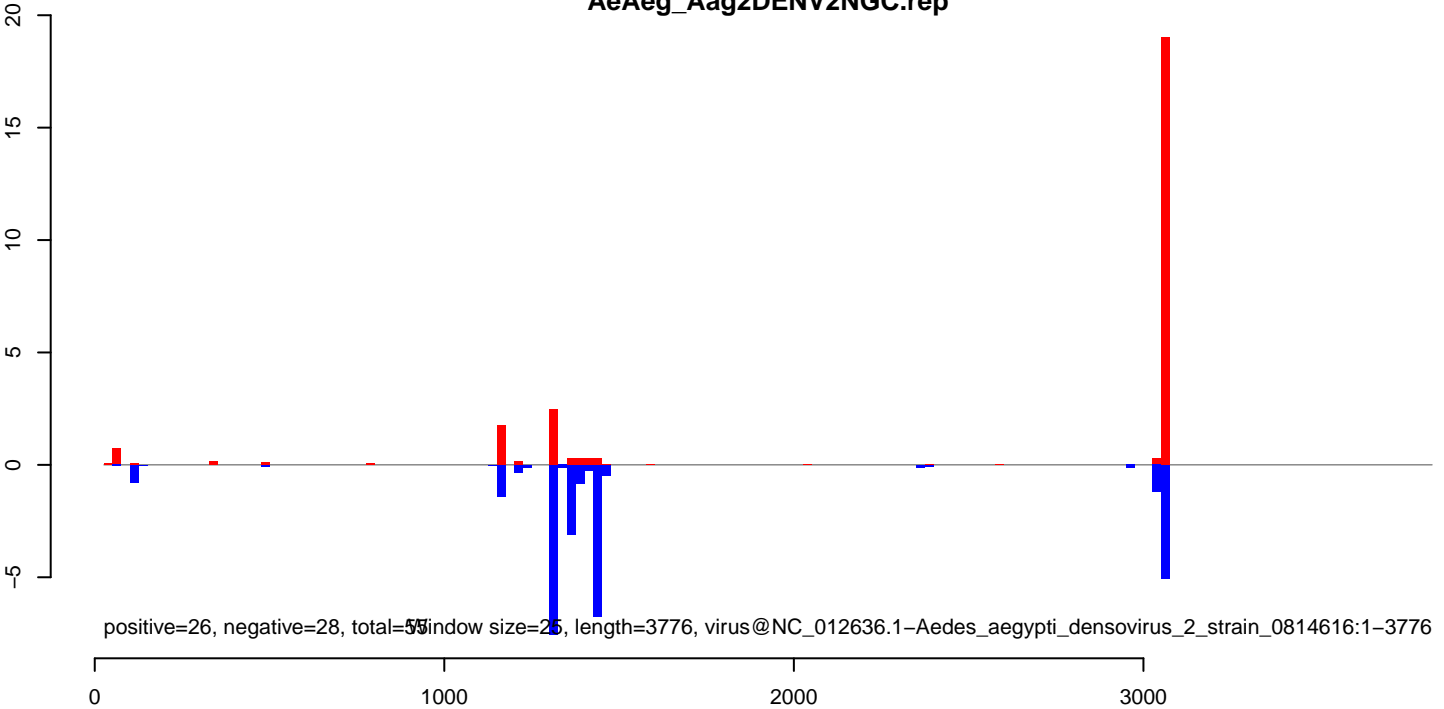
AeAeg_Aag2DENV2NGC.18_23.rep



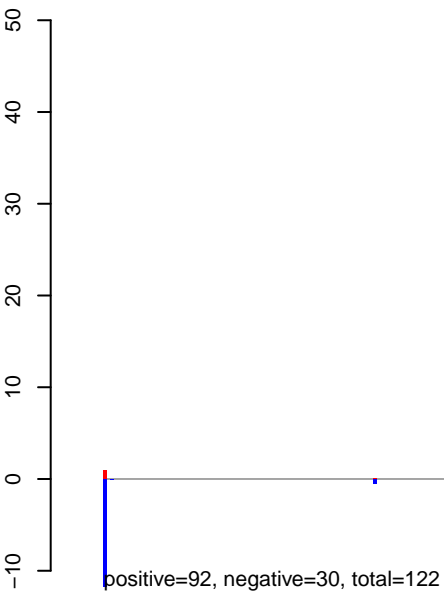
AeAeg_Aag2DENV2NGC.24_35.rep



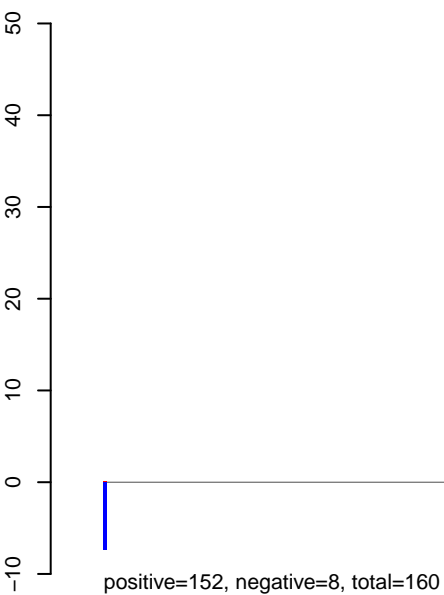
AeAeg_Aag2DENV2NGC.rep



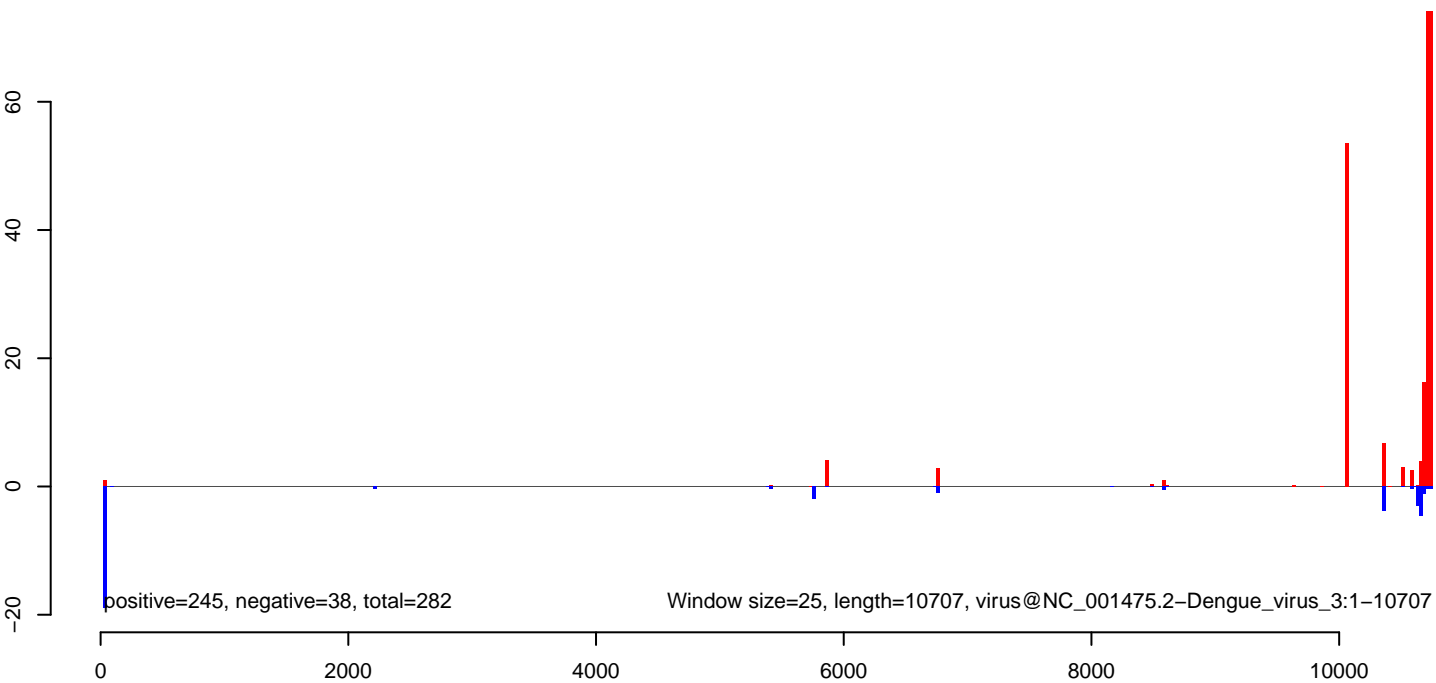
AeAeg_Aag2DENV2NGC.18_23.rep



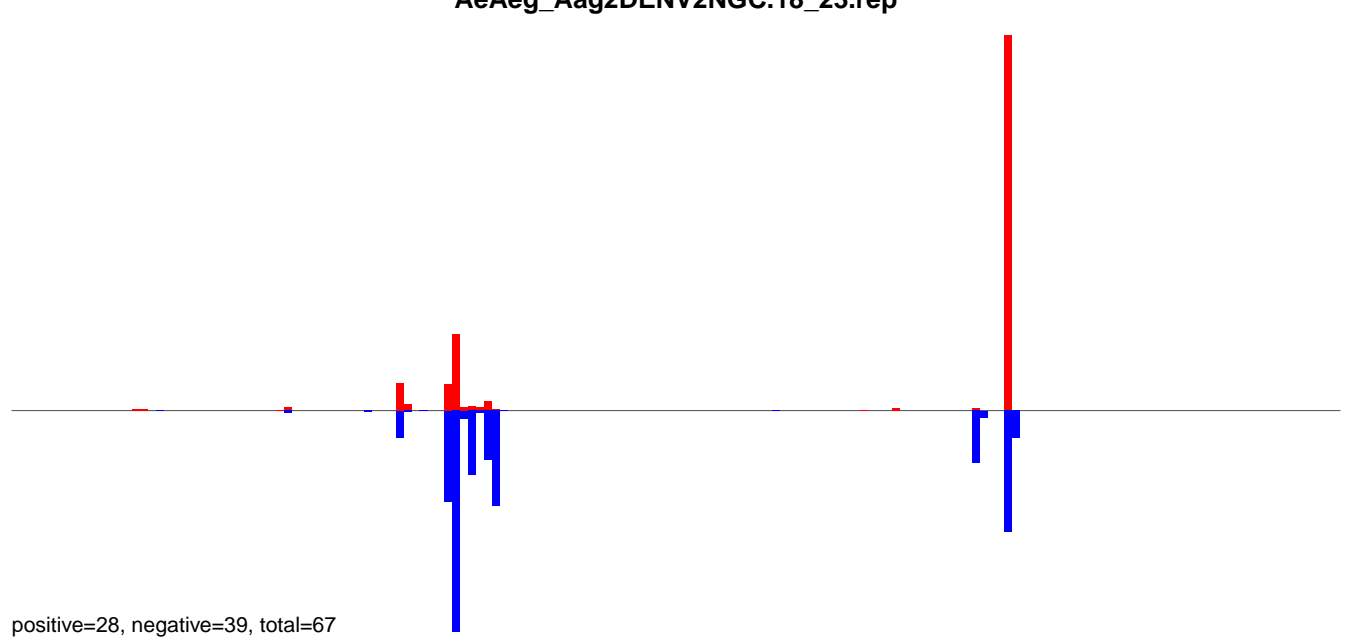
AeAeg_Aag2DENV2NGC.24_35.rep



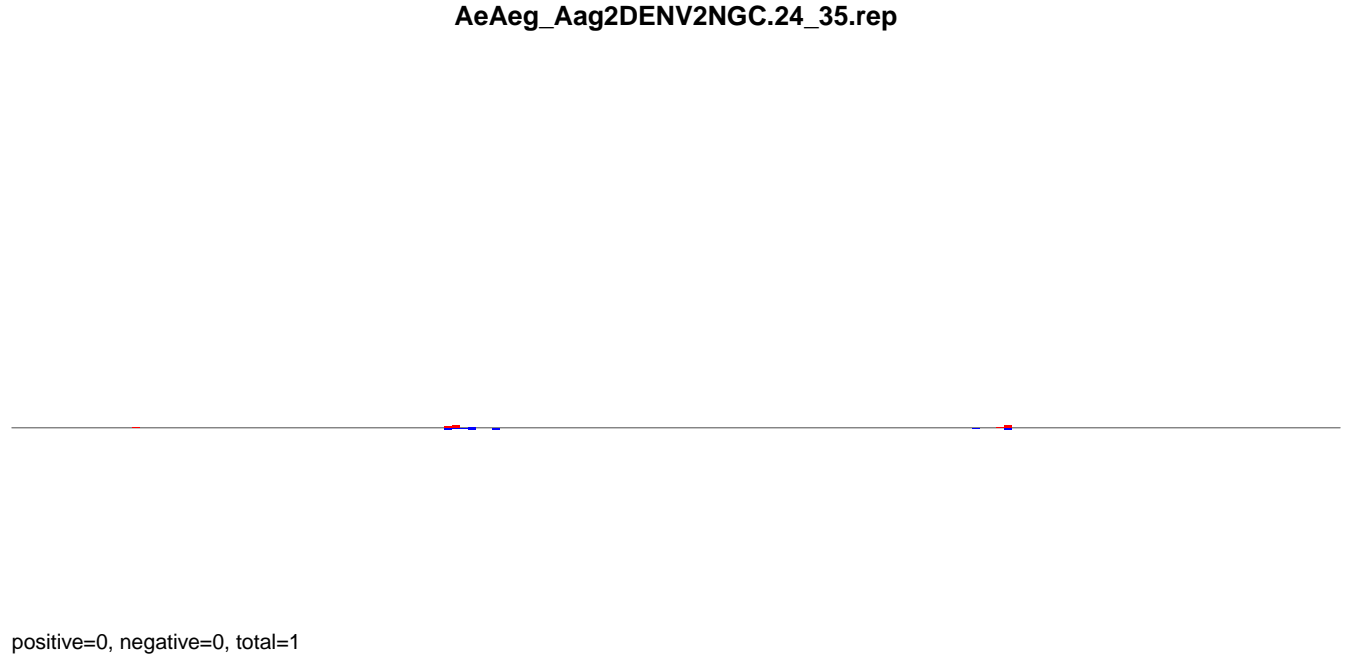
AeAeg_Aag2DENV2NGC.rep



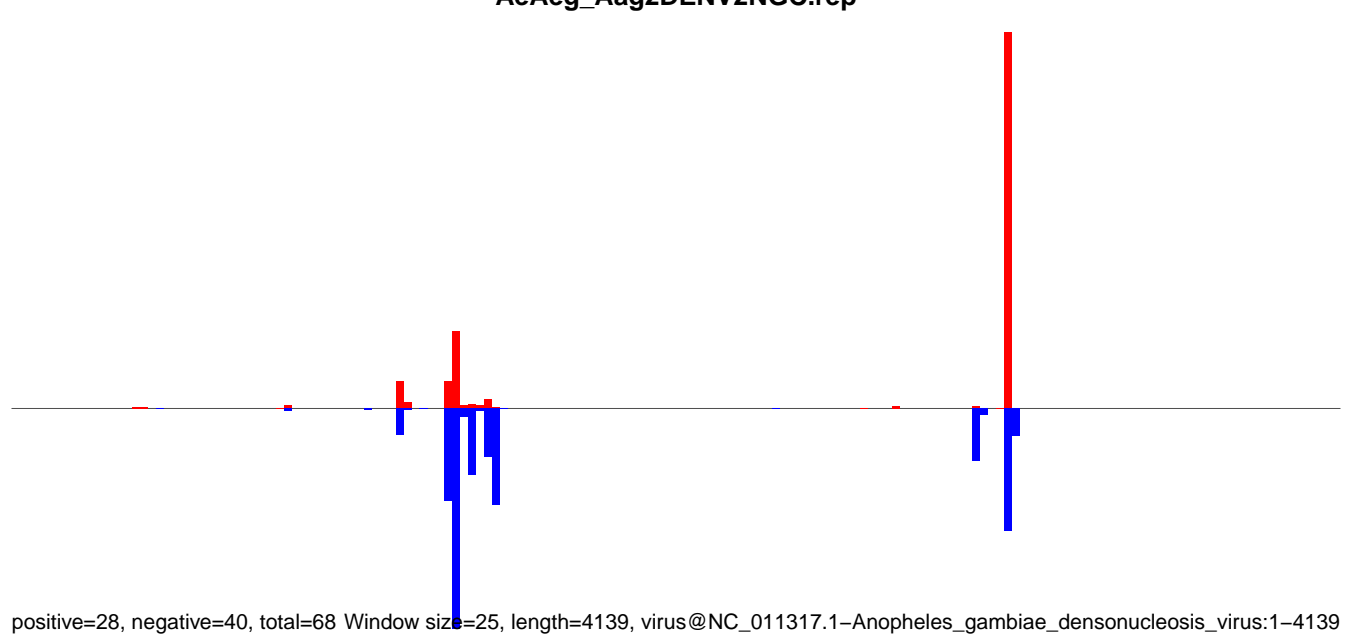
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

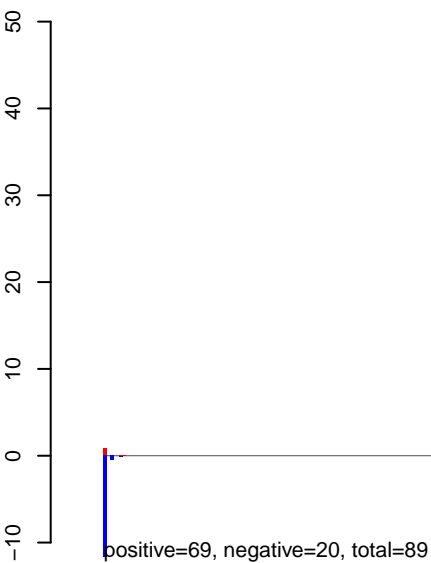


AeAeg_Aag2DENV2NGC.rep

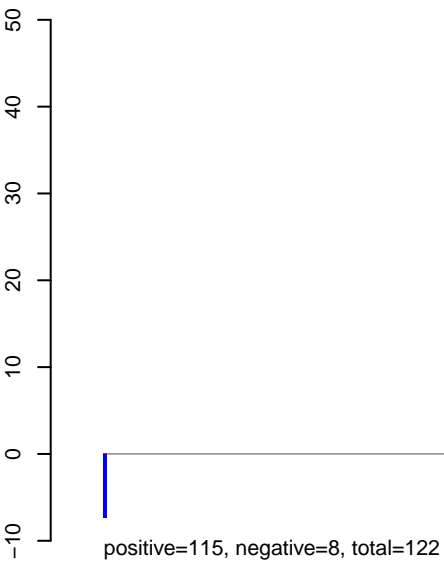


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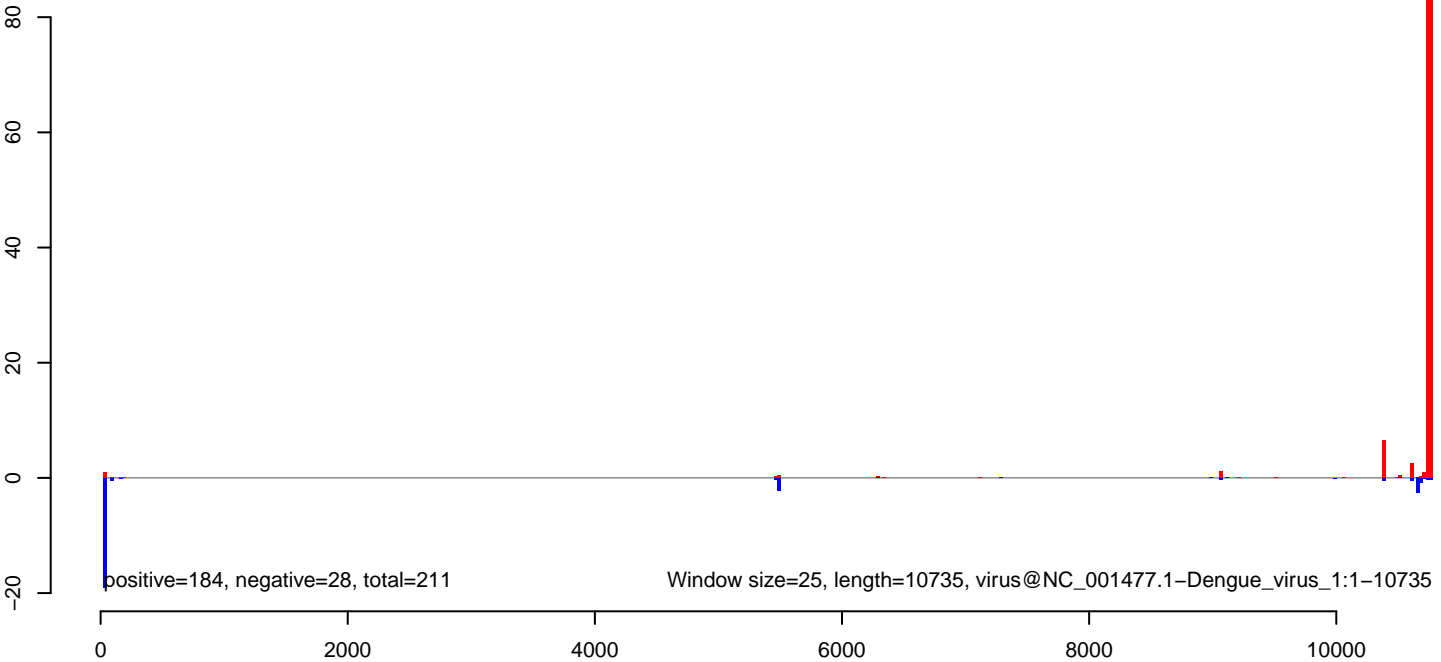
AeAeg_Aag2DENV2NGC.18_23.rep



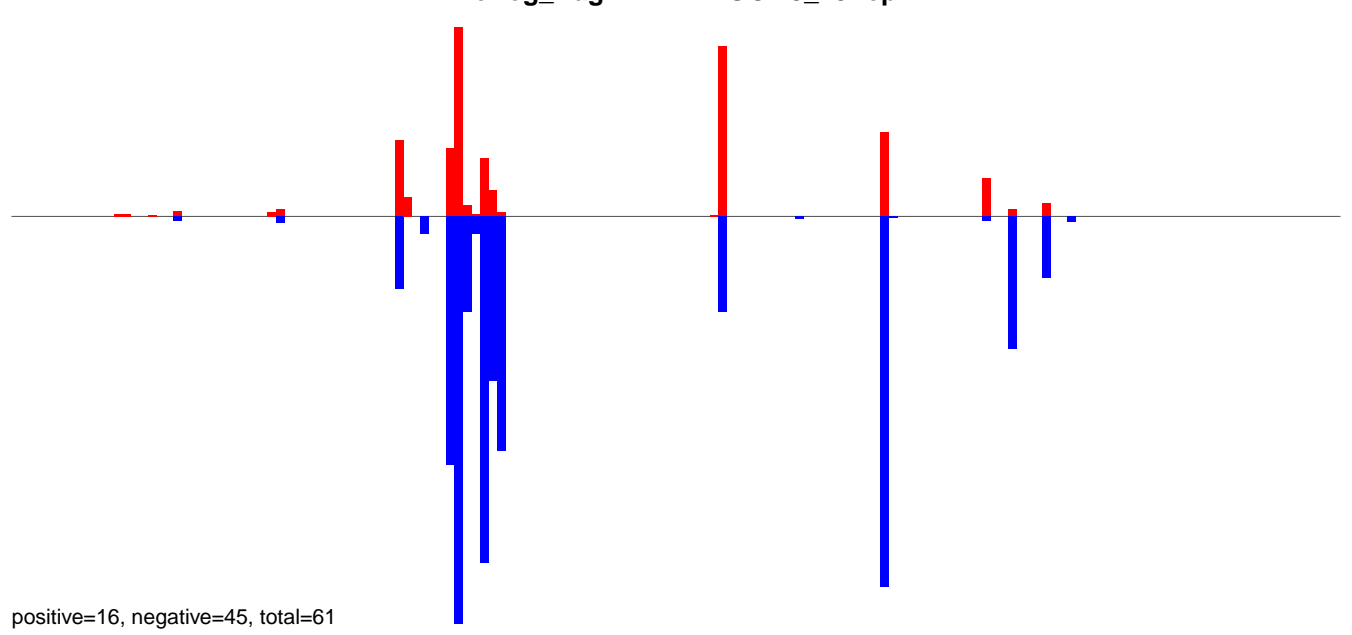
AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



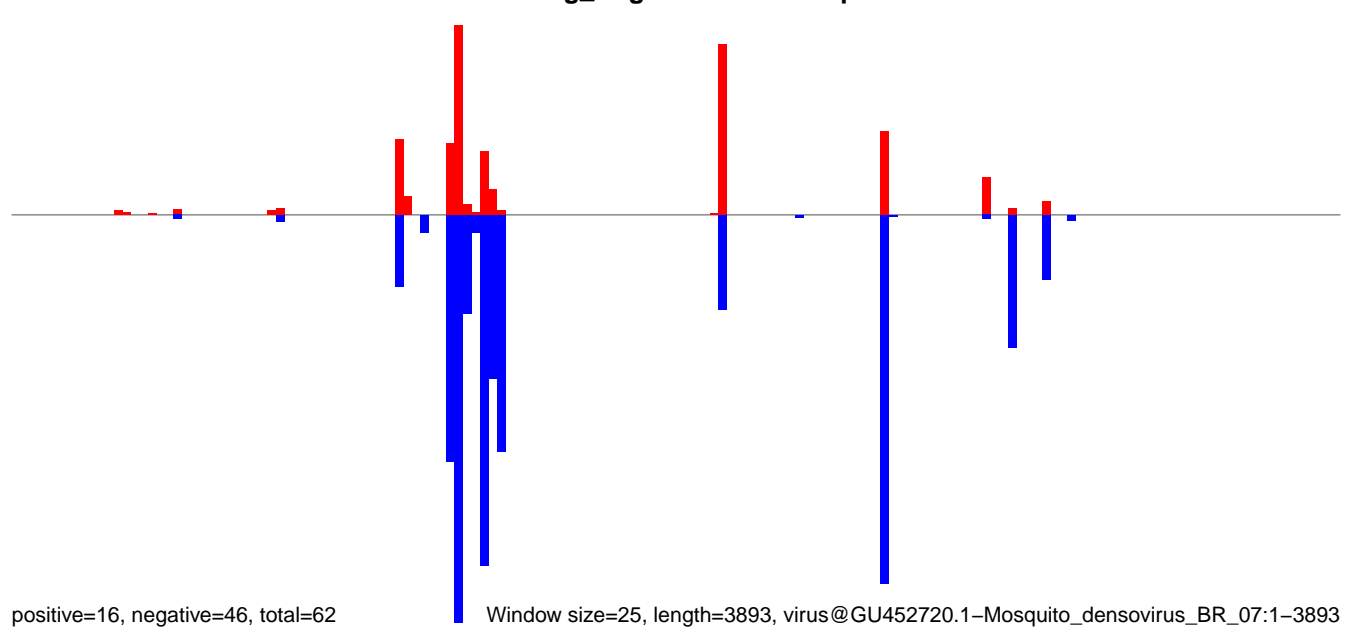
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



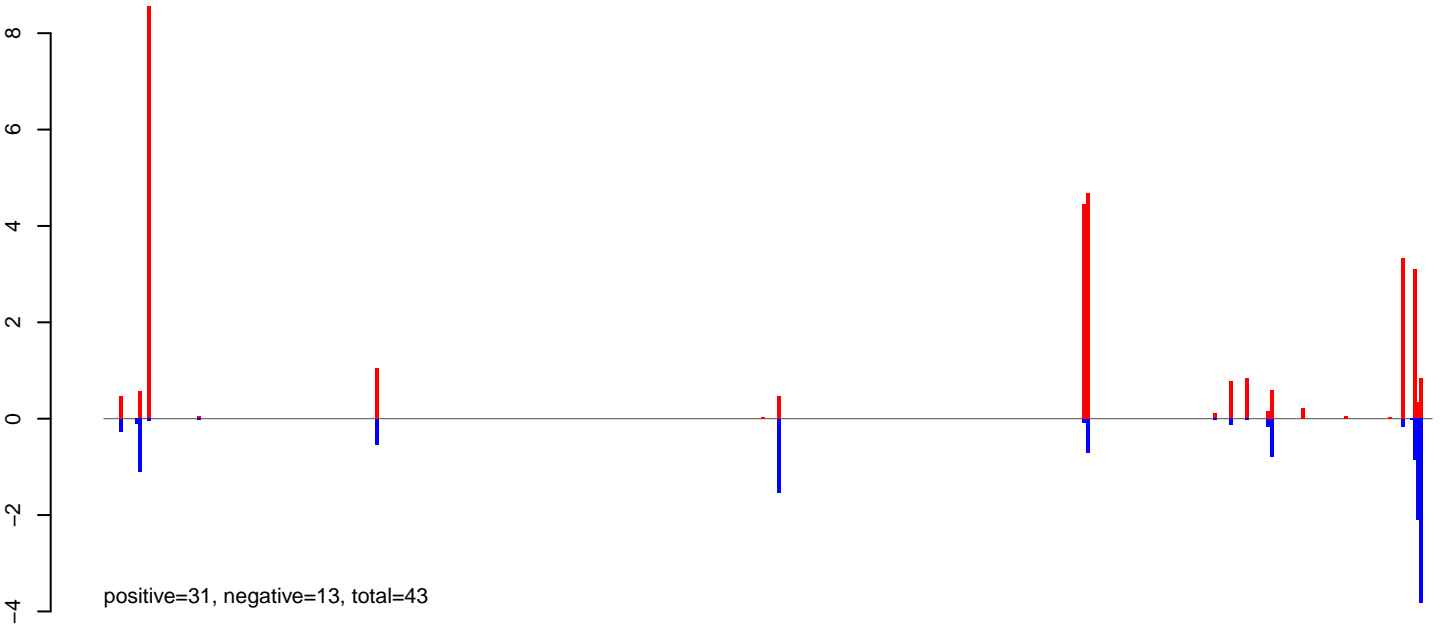
AeAeg_Aag2DENV2NGC.rep



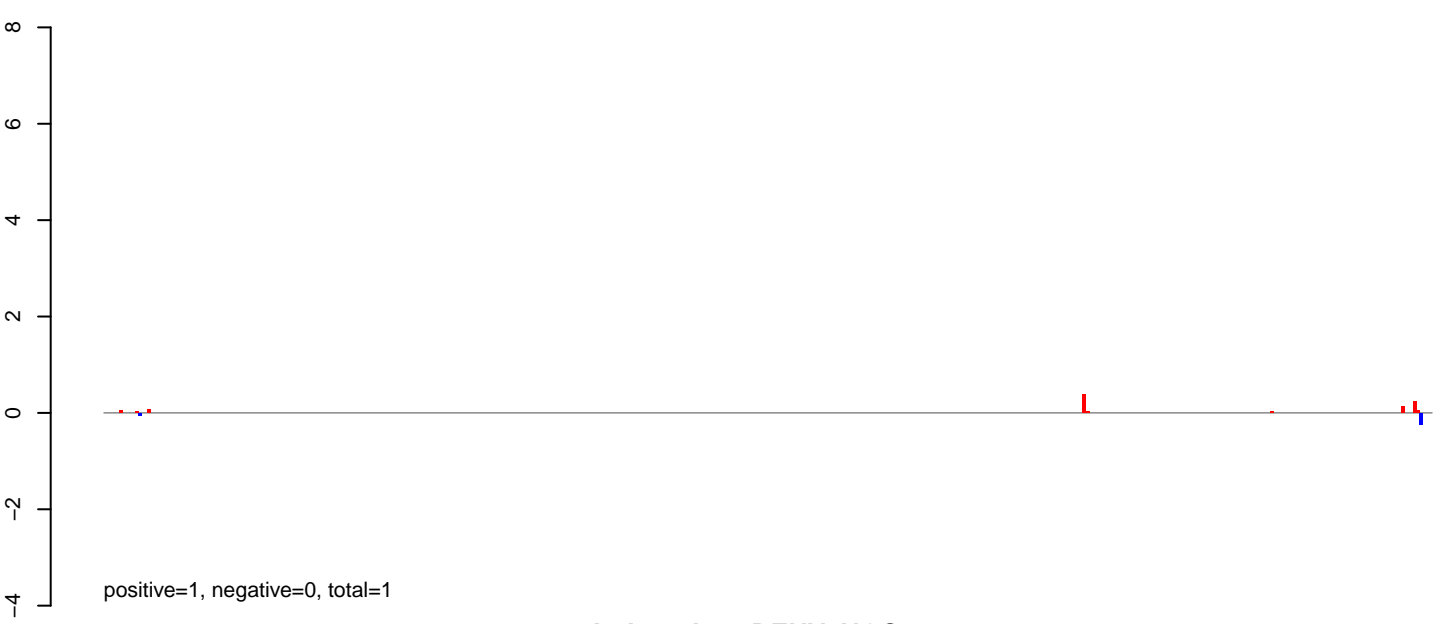
Window size=25, length=3893, virus@GU452720.1-Mosquito_densovirus_BR_07:1-3893

0 1000 2000 3000 4000

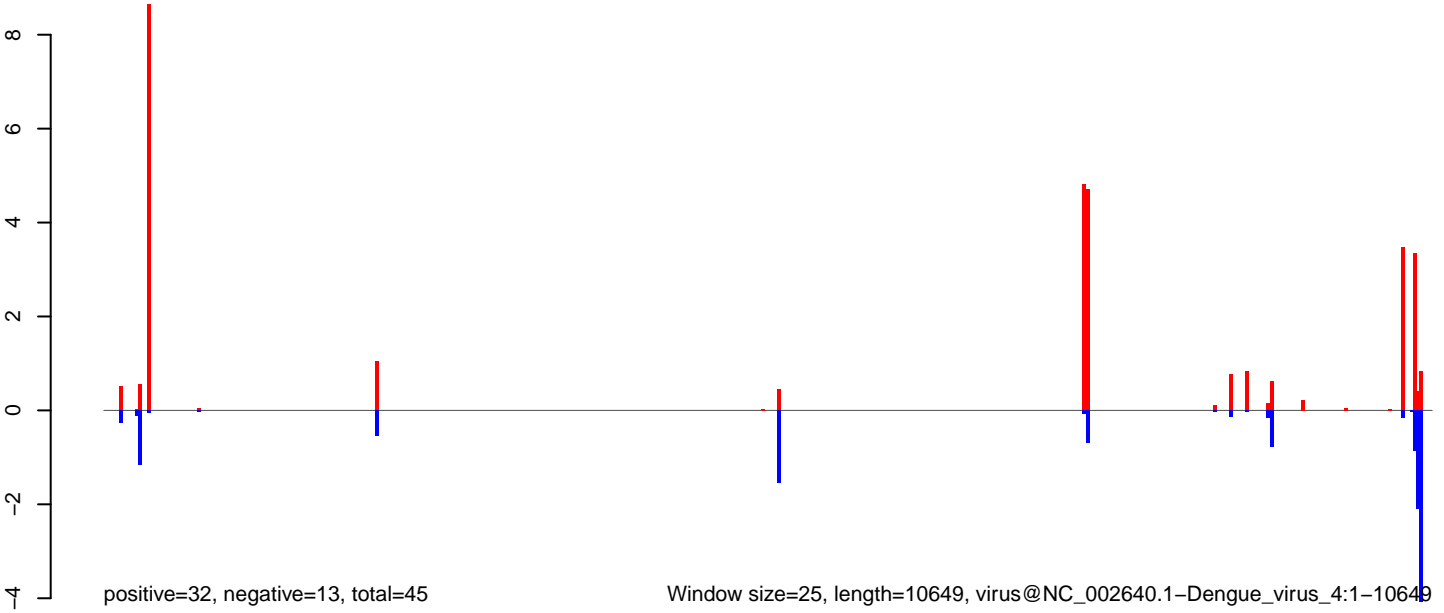
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

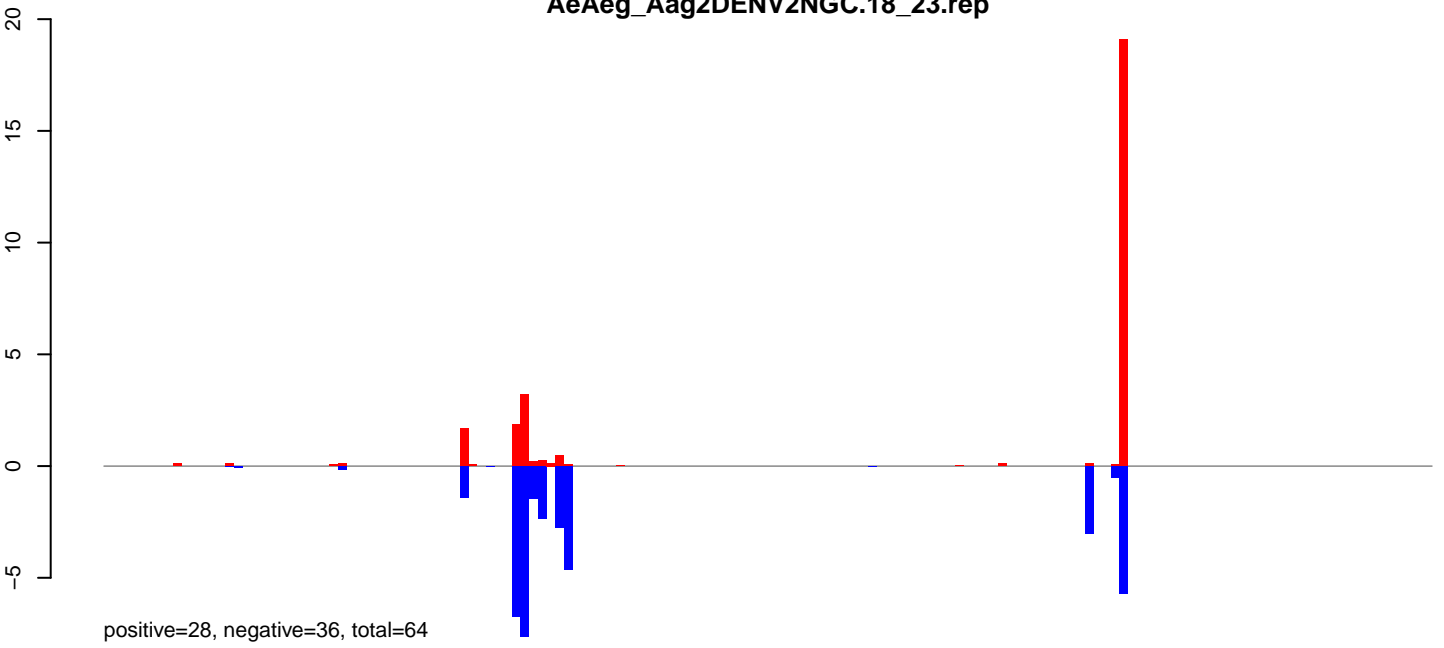


AeAeg_Aag2DENV2NGC.rep

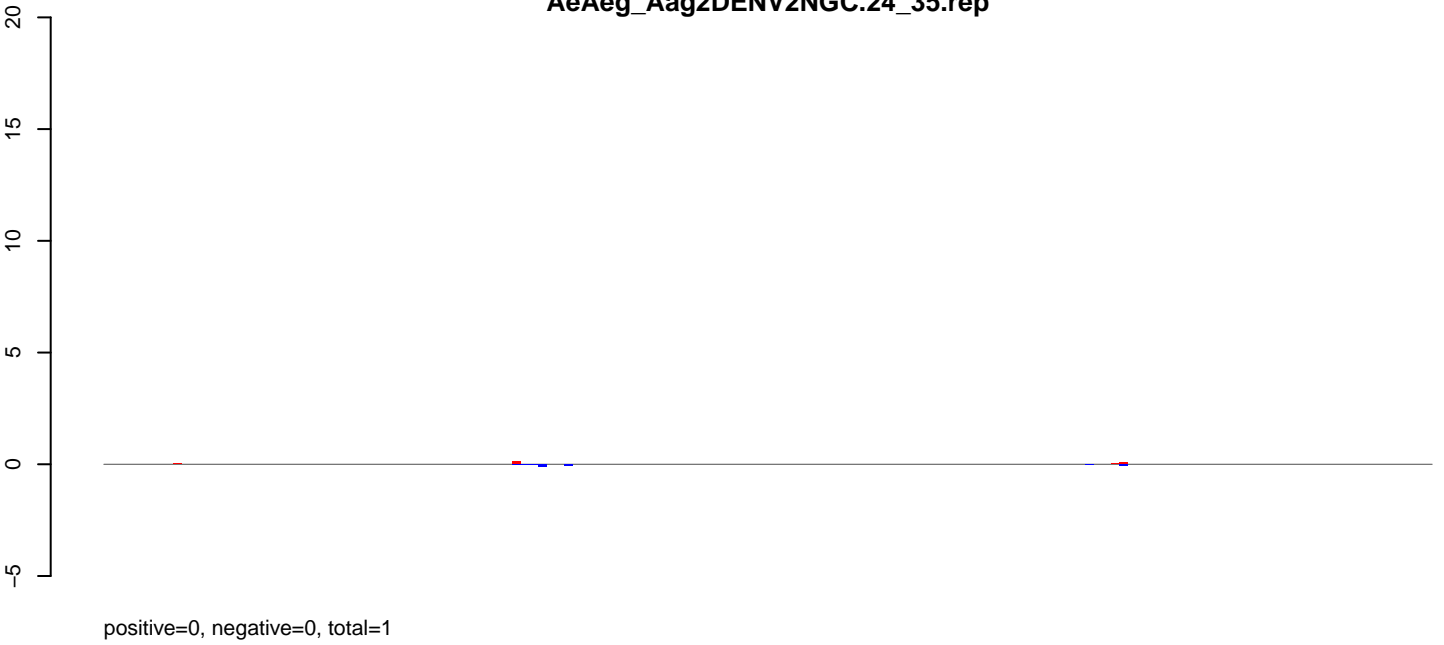


Window size=25, length=10649, virus@NC_002640.1-Dengue_virus_4:1-10649

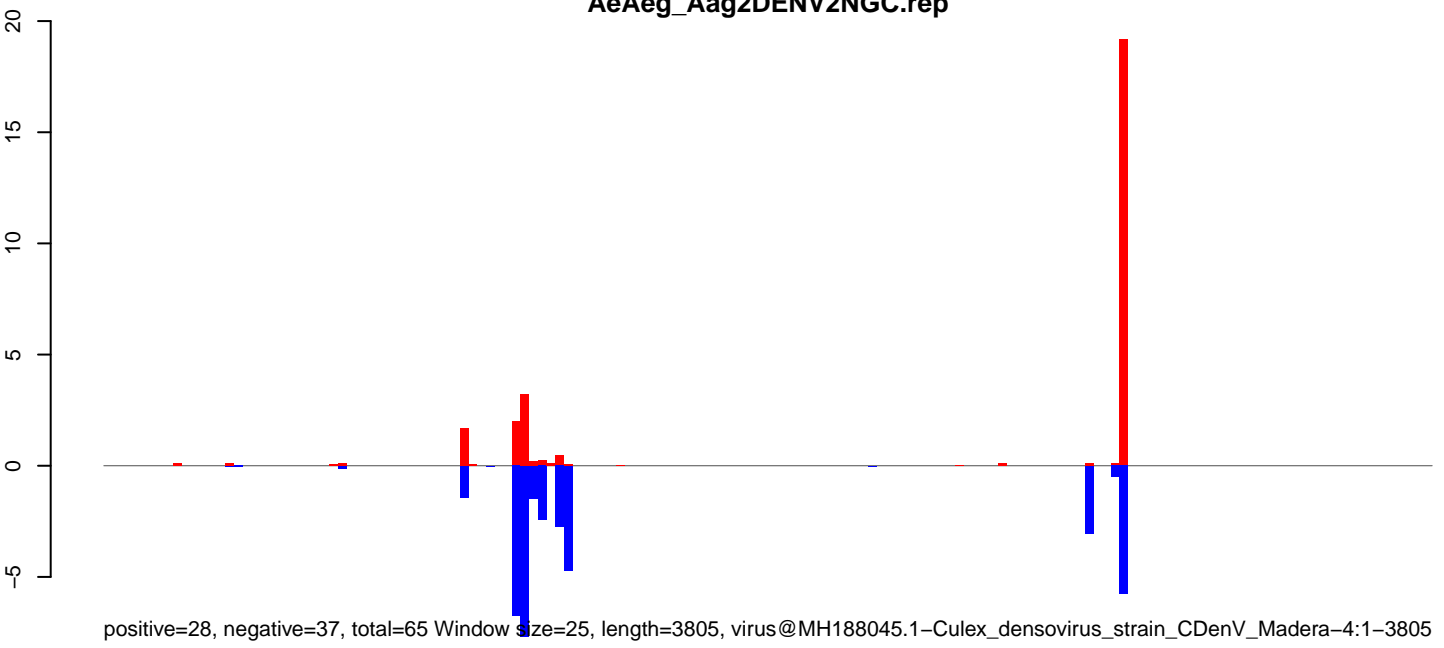
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep

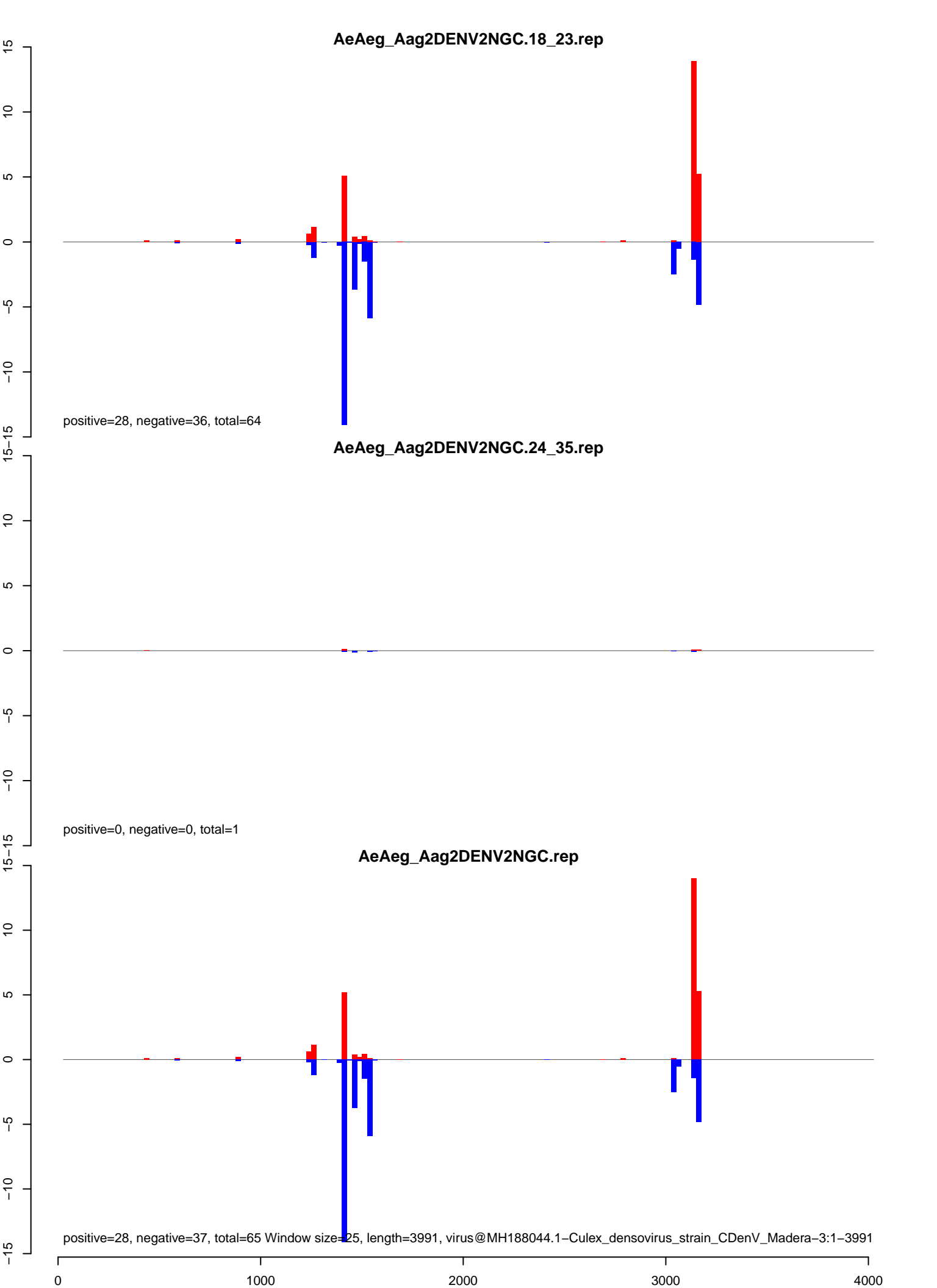
positive=28, negative=36, total=64

AeAeg_Aag2DENV2NGC.24_35.rep

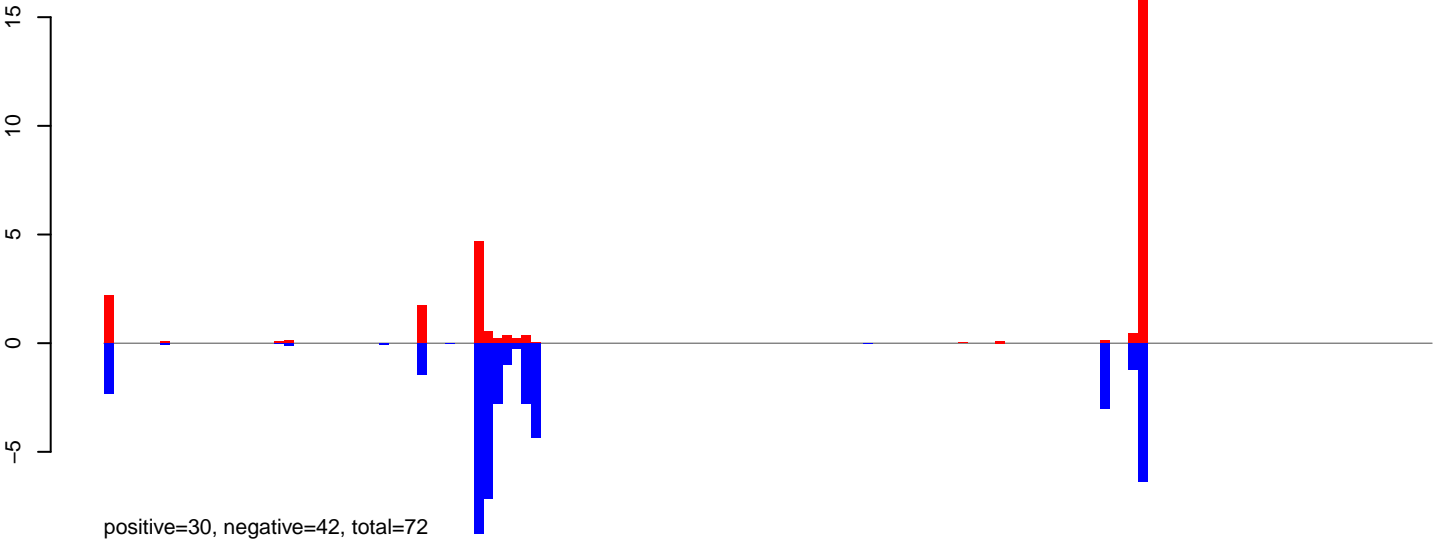
positive=0, negative=0, total=1

AeAeg_Aag2DENV2NGC.rep

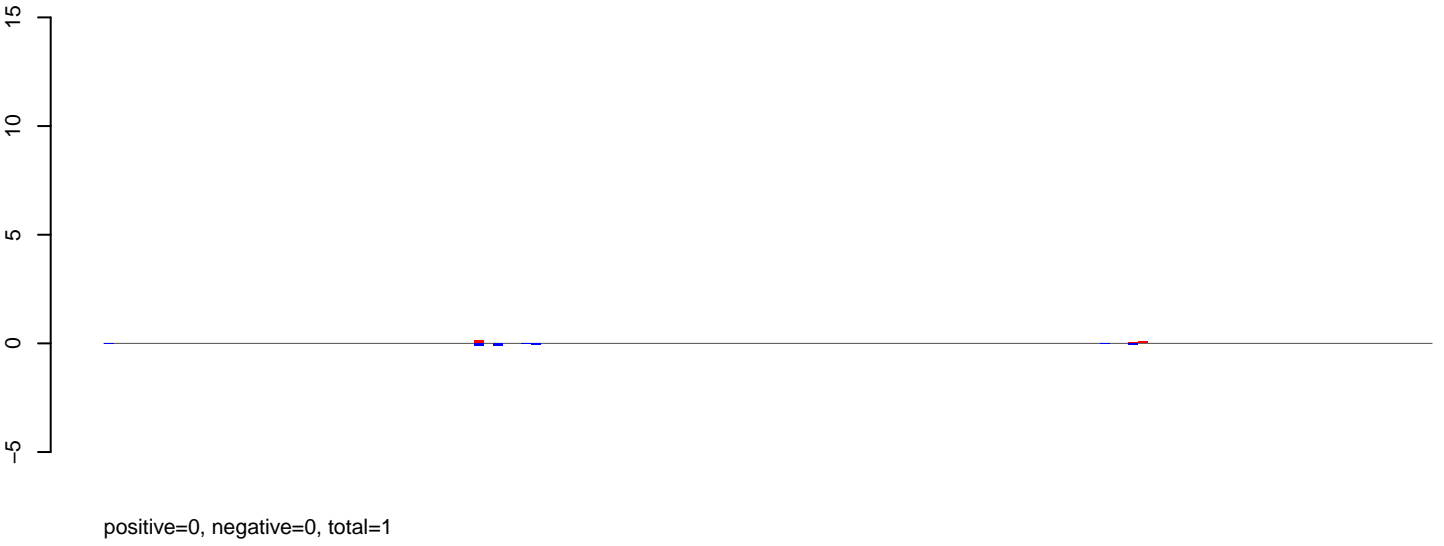
positive=28, negative=37, total=65 Window size=25, length=3991, virus@MH188044.1-Culex_densovirus_strain_CDenV_Madera-3:1-3991



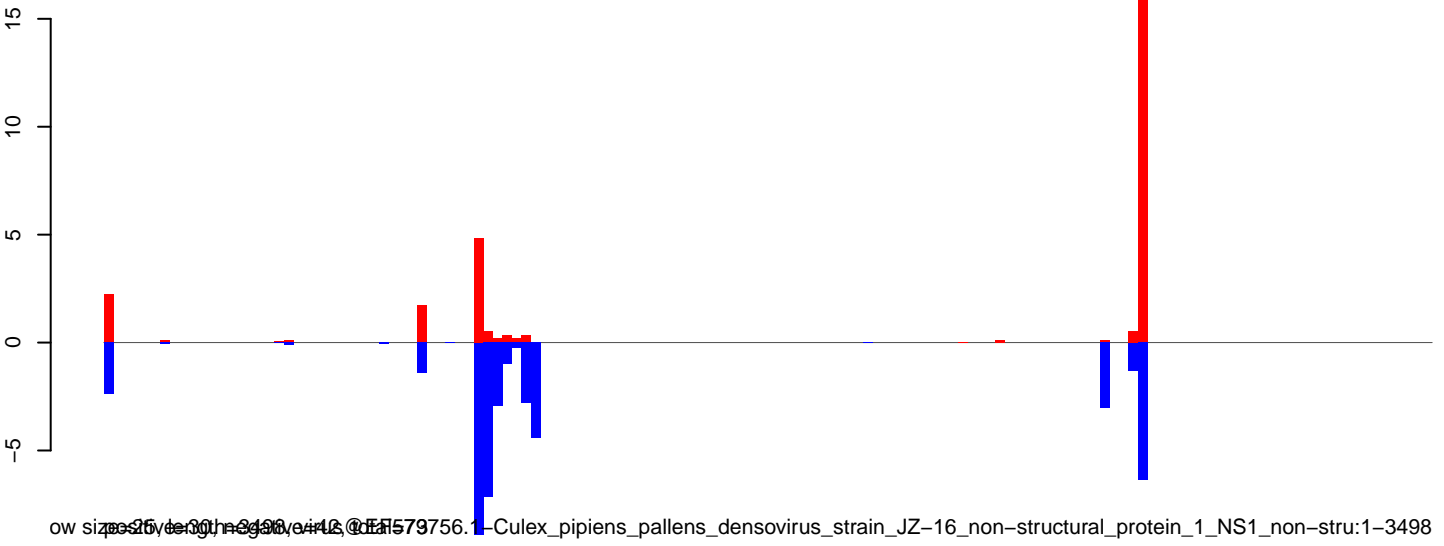
AeAeg_Aag2DENV2NGC.18_23.rep



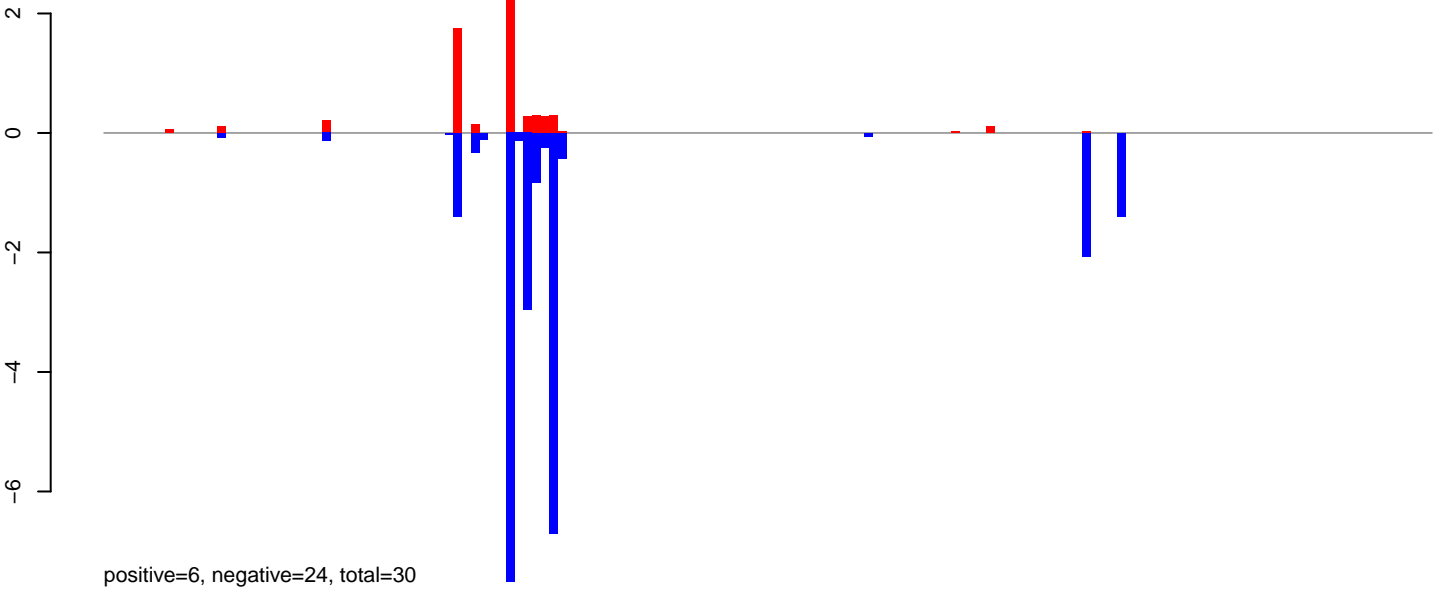
AeAeg_Aag2DENV2NGC.24_35.rep



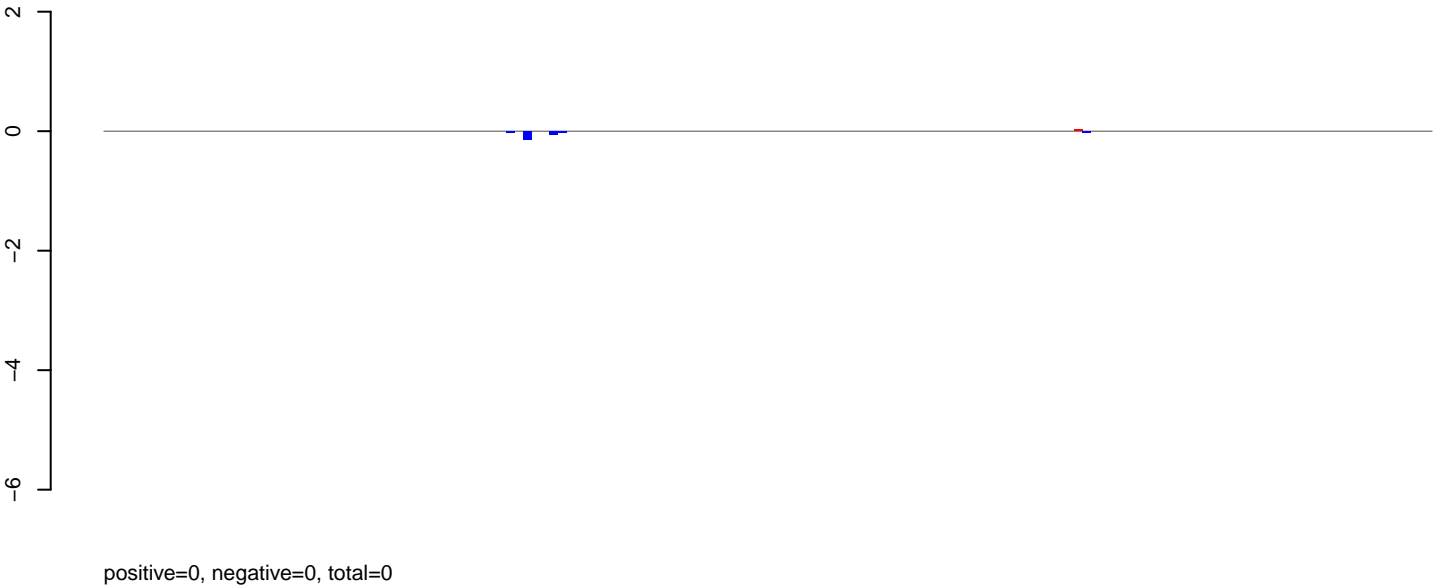
AeAeg_Aag2DENV2NGC.rep



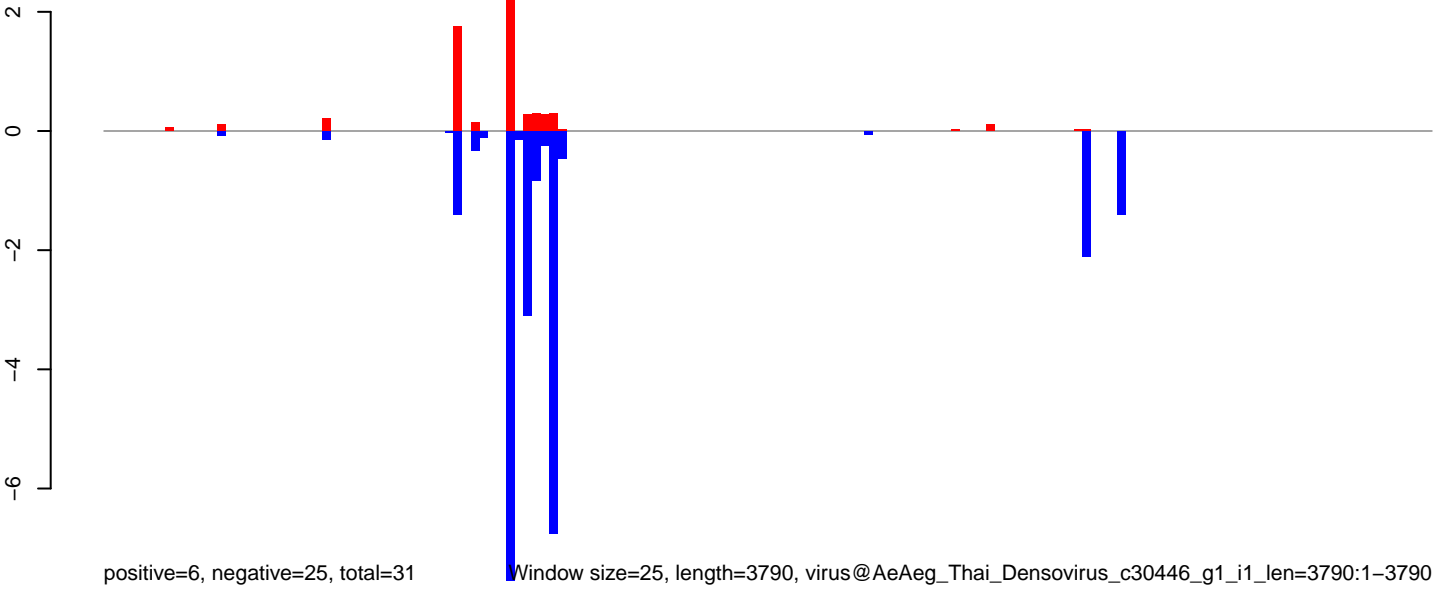
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



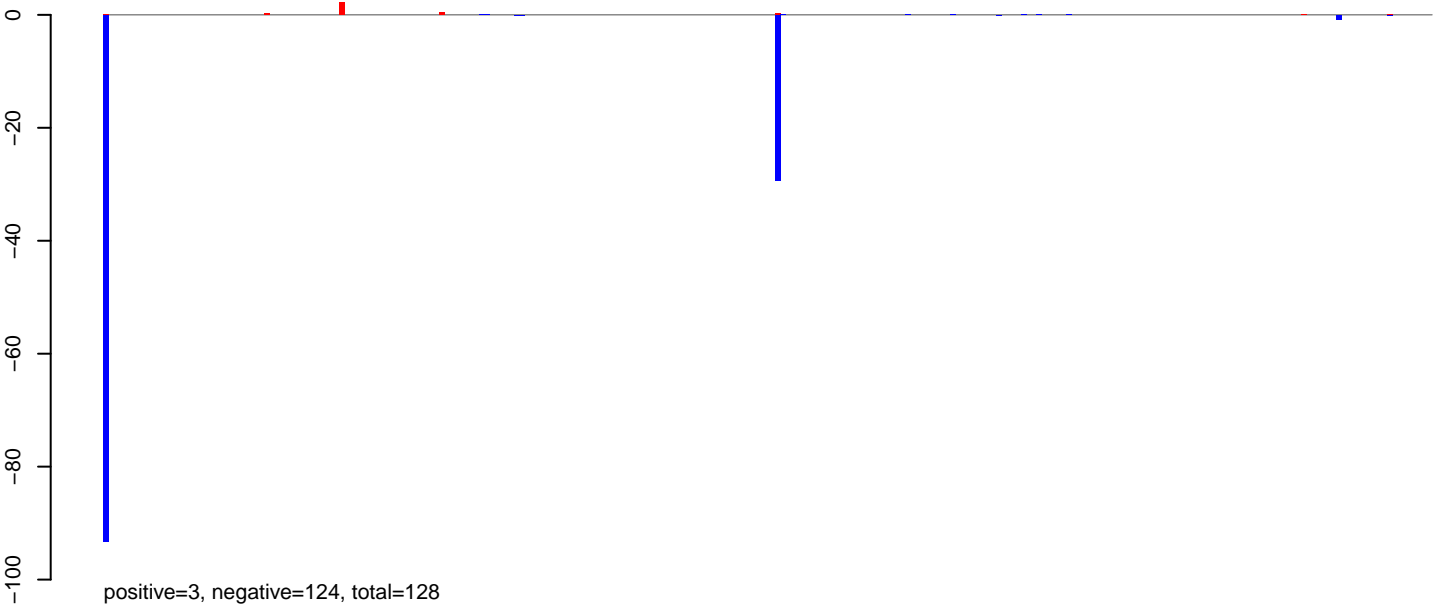
AeAeg_Aag2DENV2NGC.rep



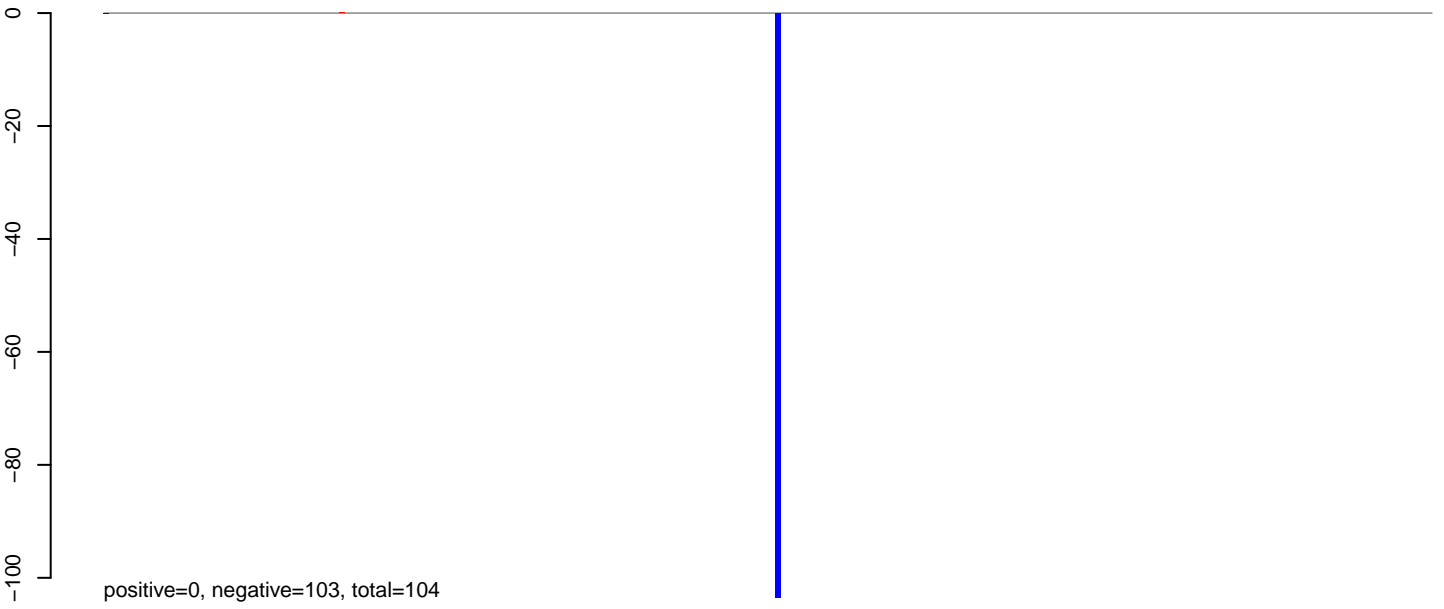
Window size=25, length=3790, virus@AeAeg_Thai_Densovirus_c30446_g1_i1_len=3790:1-3790

0 1000 2000 3000

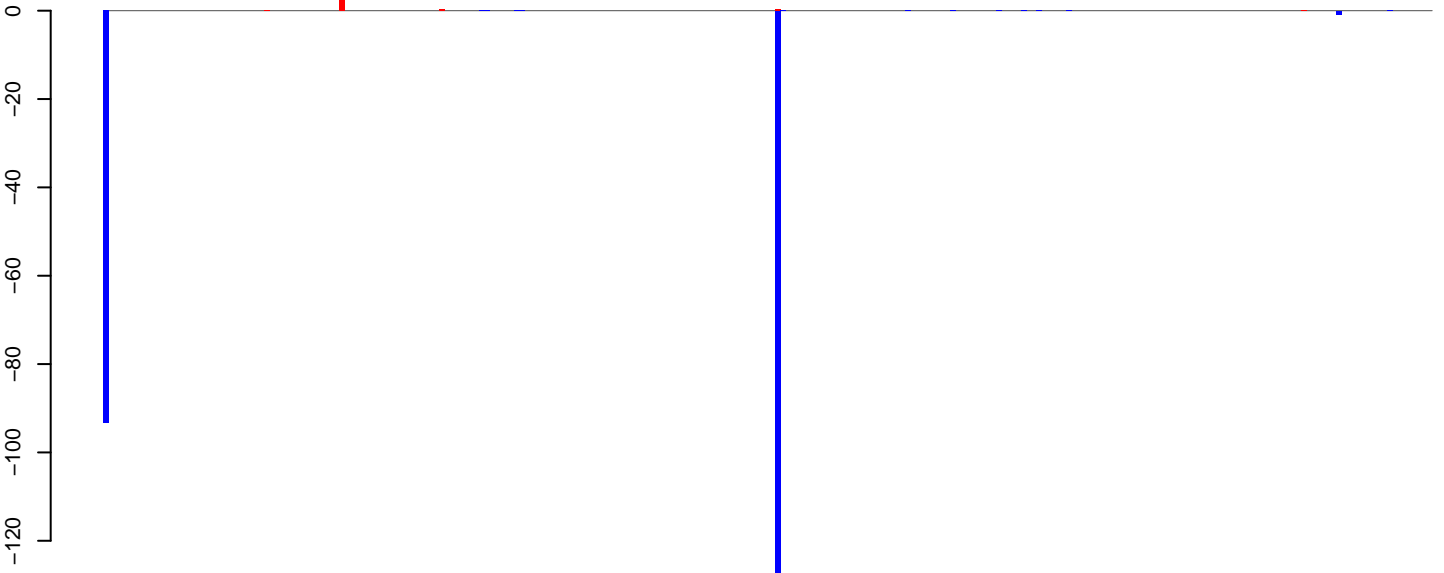
AeAeg_Aag2DENV2NGC.18_23.rep



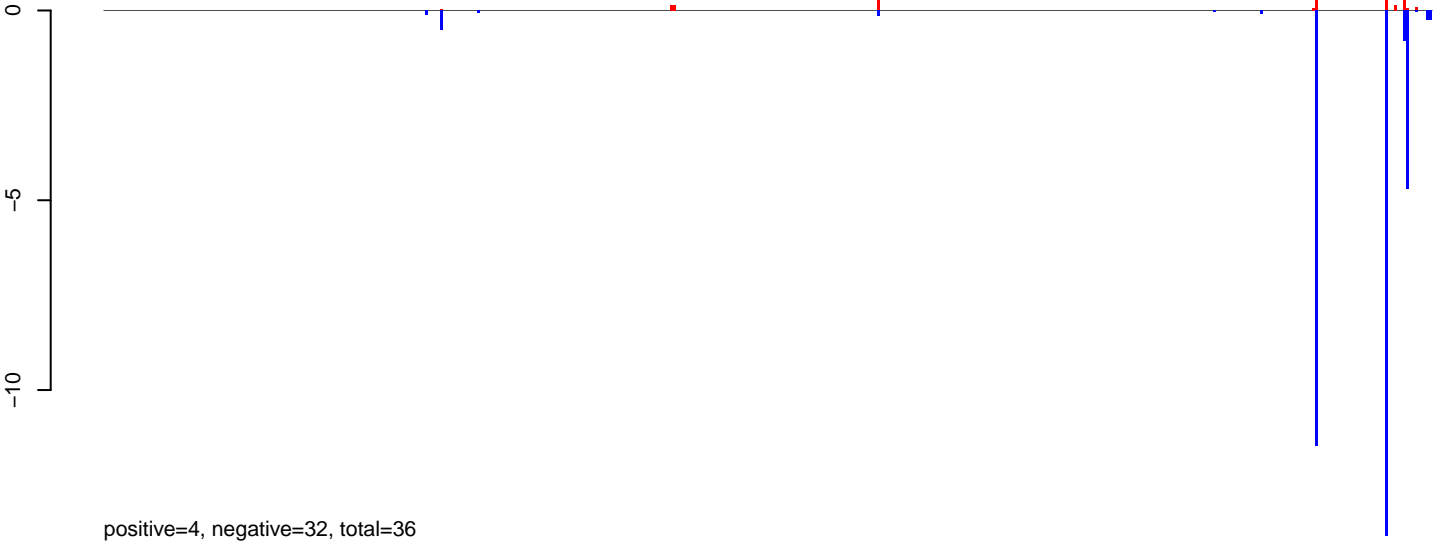
AeAeg_Aag2DENV2NGC.24_35.rep



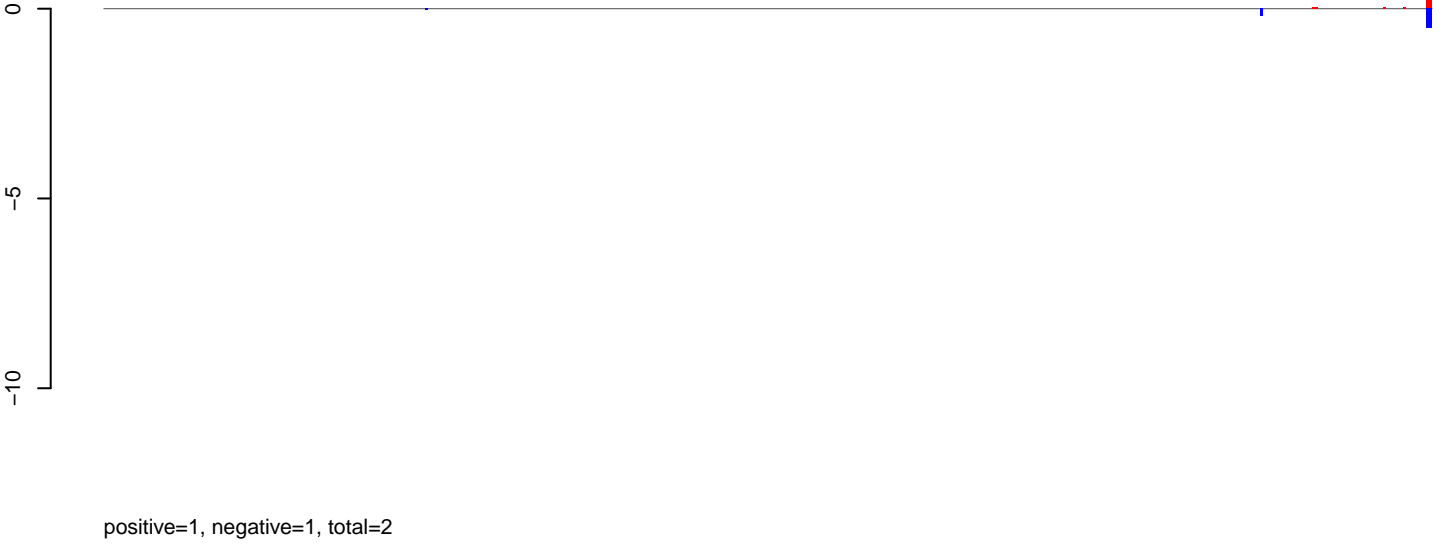
AeAeg_Aag2DENV2NGC.rep



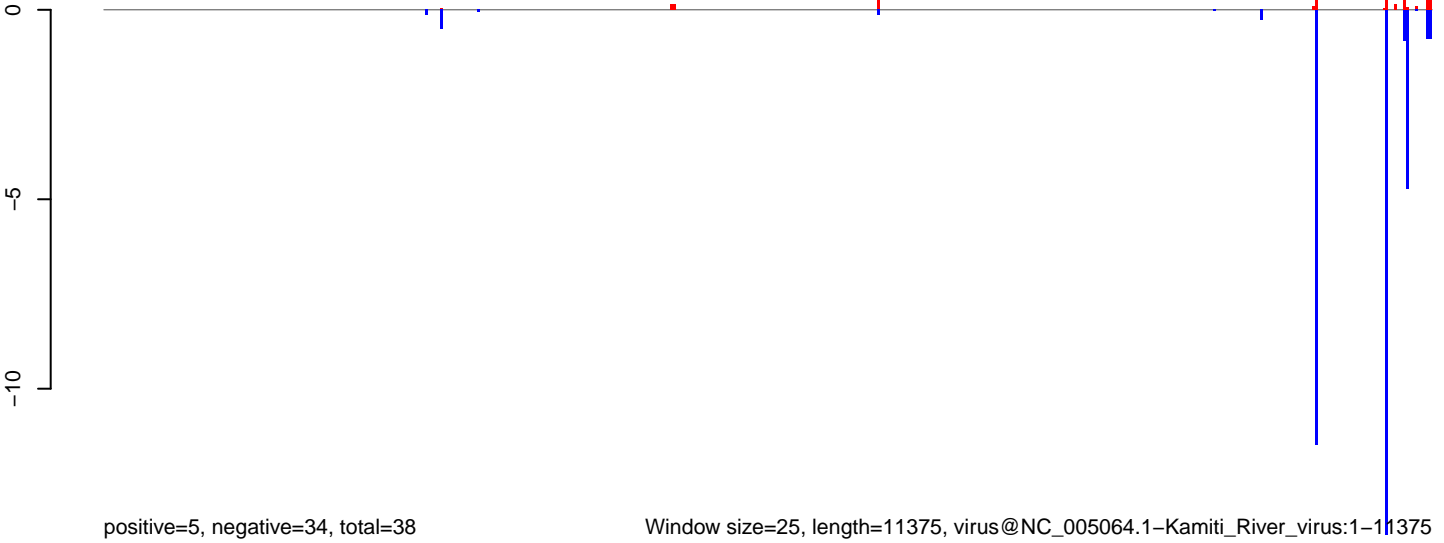
AeAeg_Aag2DENV2NGC.18_23.rep



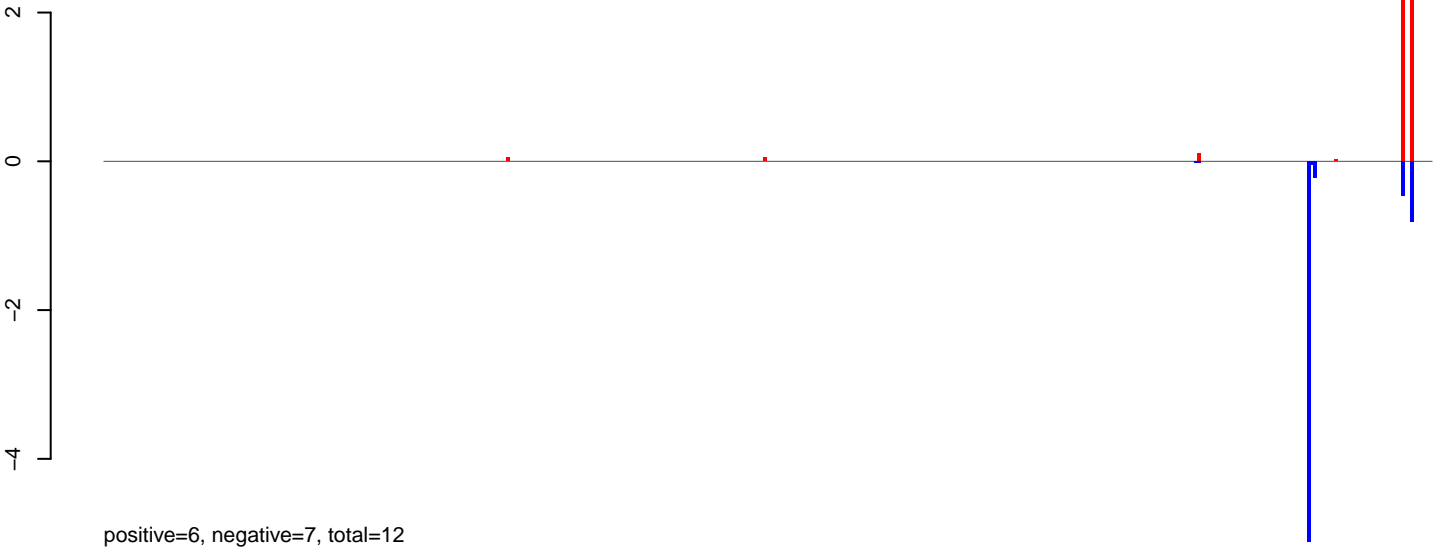
AeAeg_Aag2DENV2NGC.24_35.rep



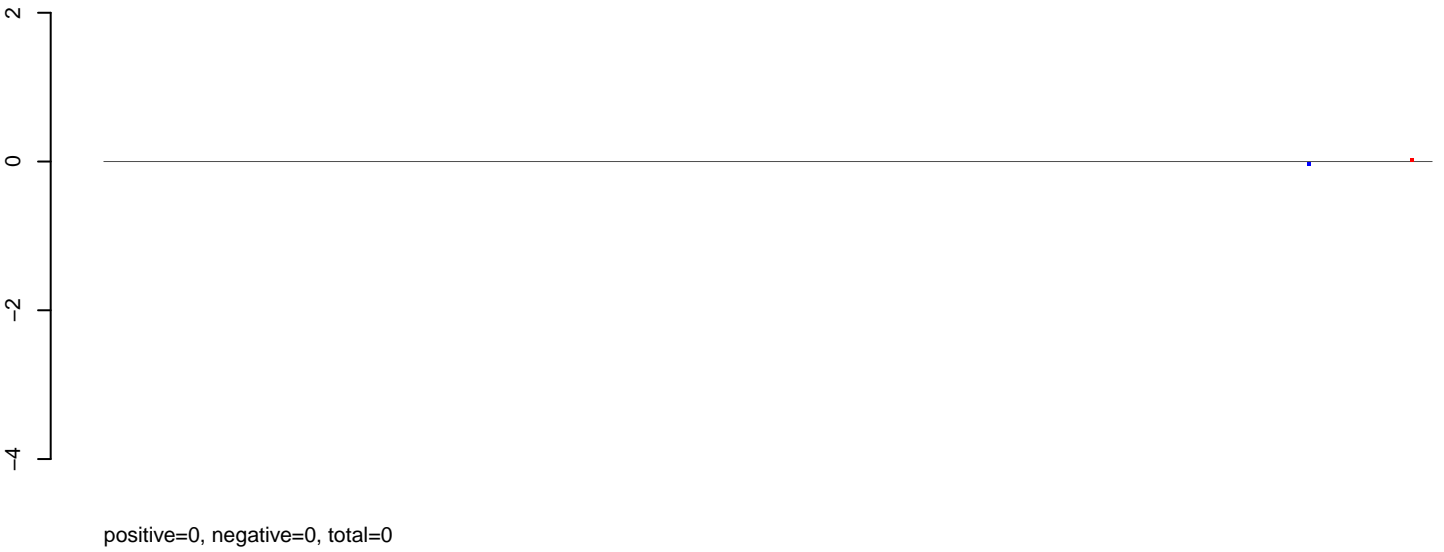
AeAeg_Aag2DENV2NGC.rep



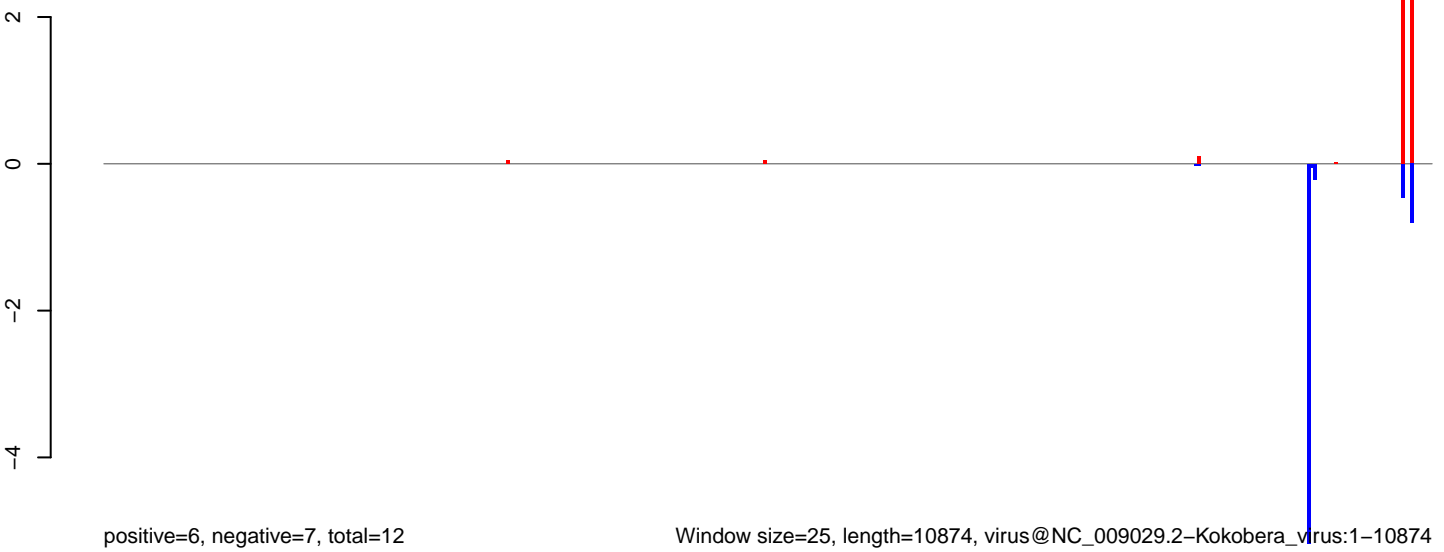
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep

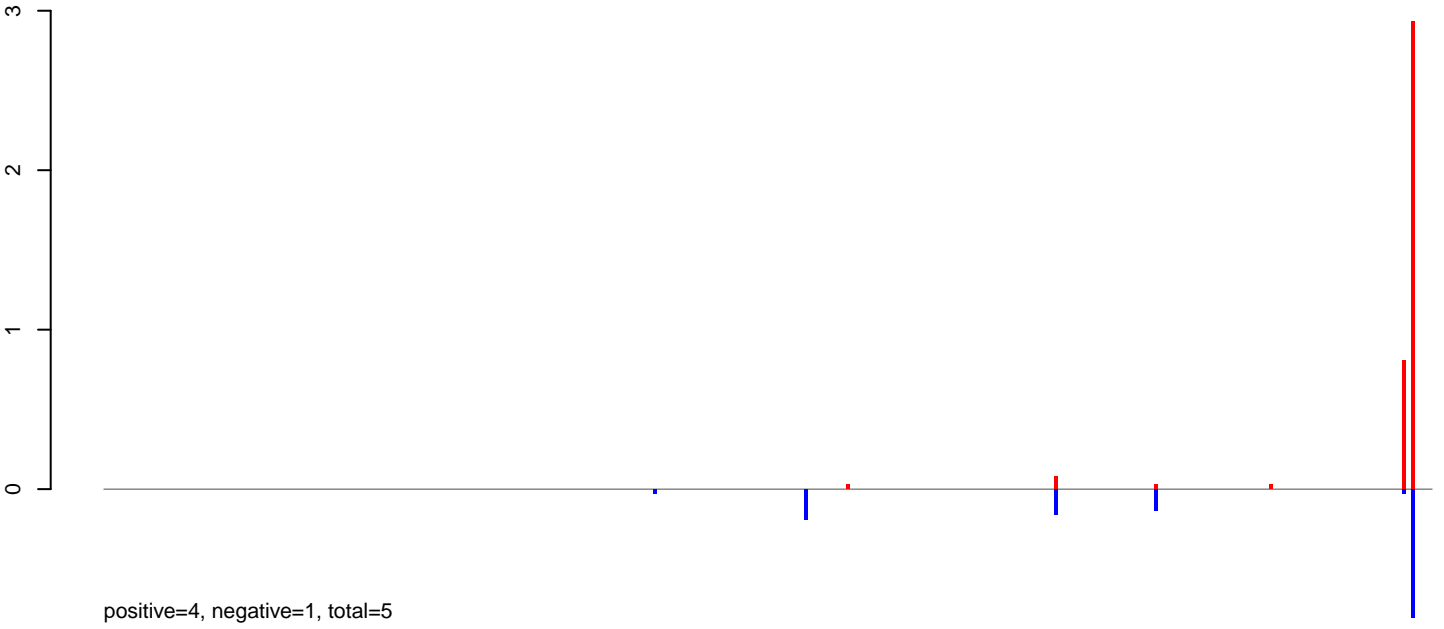


Window size=25, length=10874, virus@NC_009029.2-Kokobera_virus:1-10874

0 2000 4000 6000 8000 10000

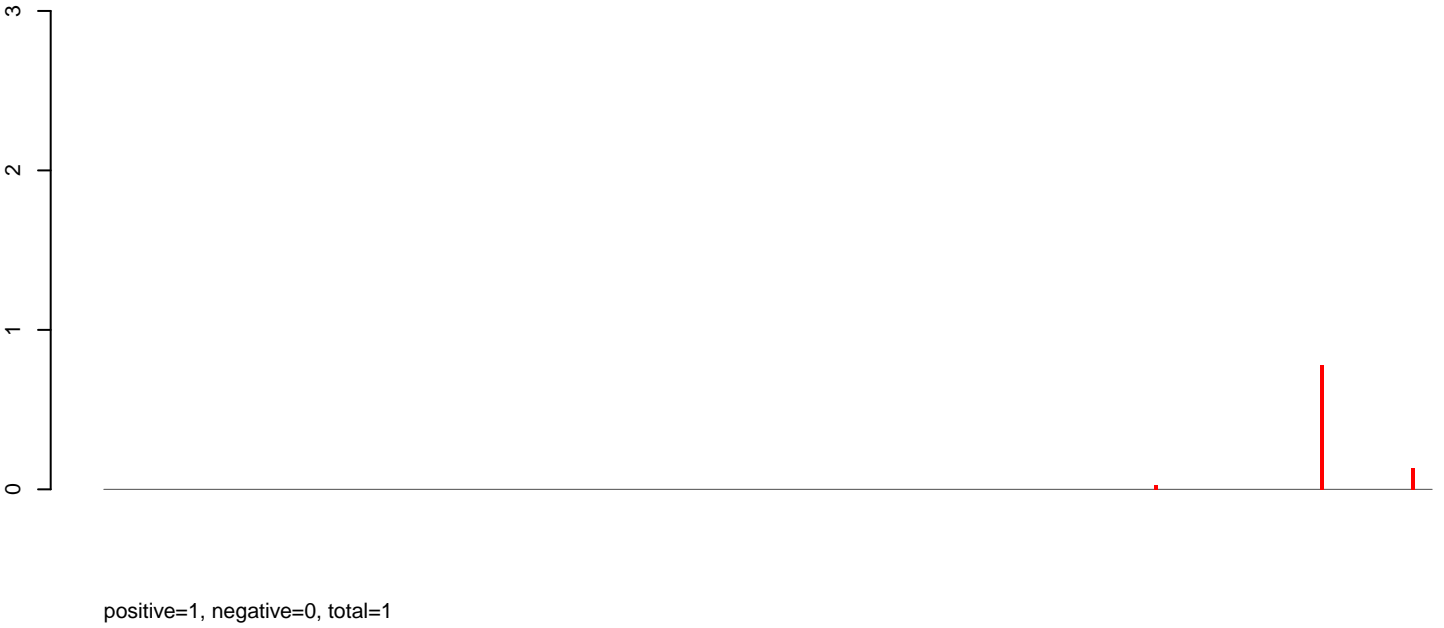
AeAeg_Aag2DENV2NGC.18_23.rep

positive=4, negative=1, total=5



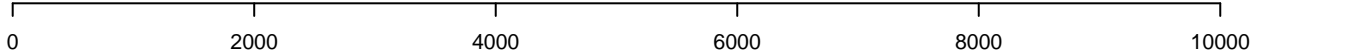
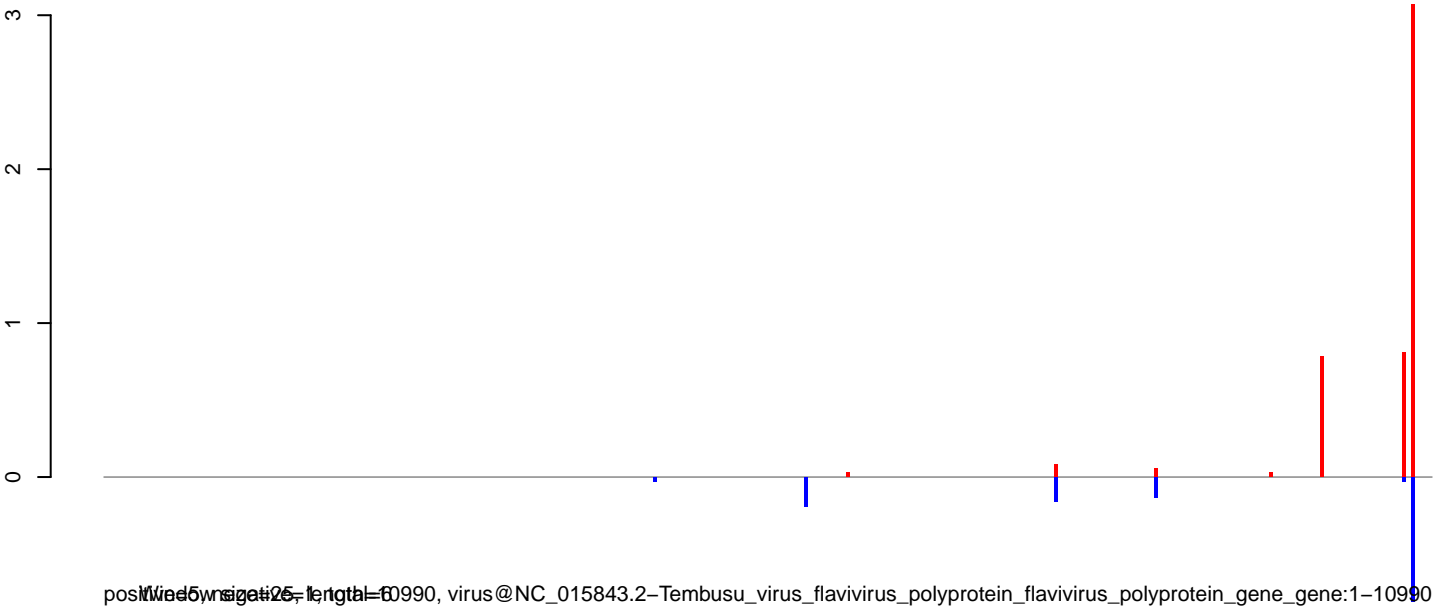
AeAeg_Aag2DENV2NGC.24_35.rep

positive=1, negative=0, total=1

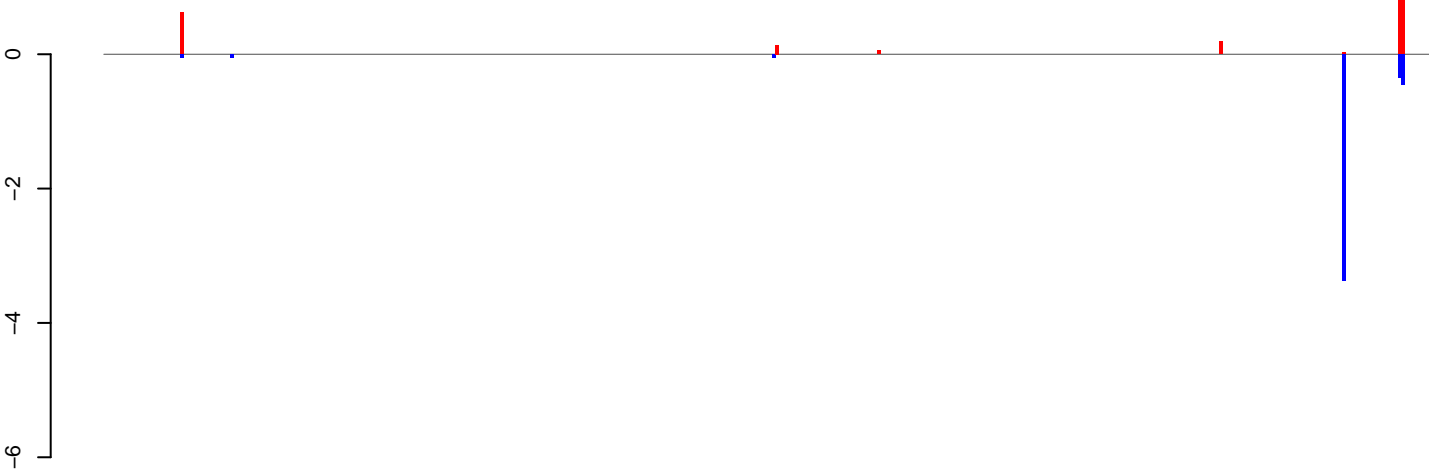


AeAeg_Aag2DENV2NGC.rep

positive=5, negative=1, total=6
width=25, length=60990, virus@NC_015843.2-Tembusu_virus_flavivirus_polyprotein_flavivirus_polyprotein_gene_gene:1-10990

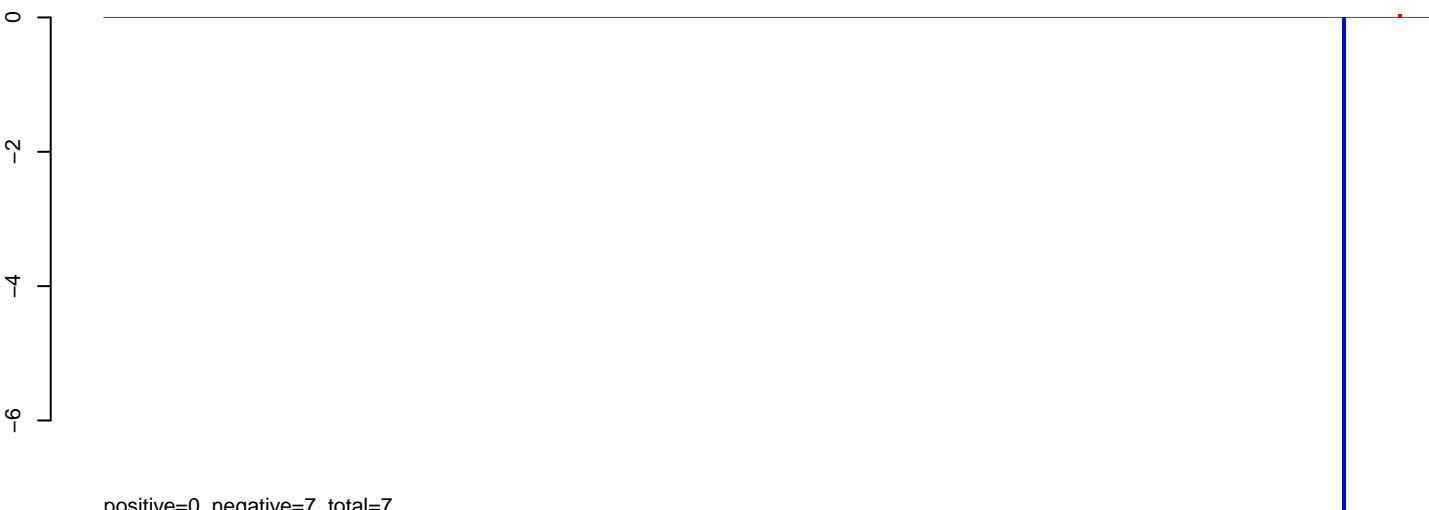


AeAeg_Aag2DENV2NGC.18_23.rep



positive=4, negative=4, total=8

AeAeg_Aag2DENV2NGC.24_35.rep



positive=0, negative=7, total=7

AeAeg_Aag2DENV2NGC.rep

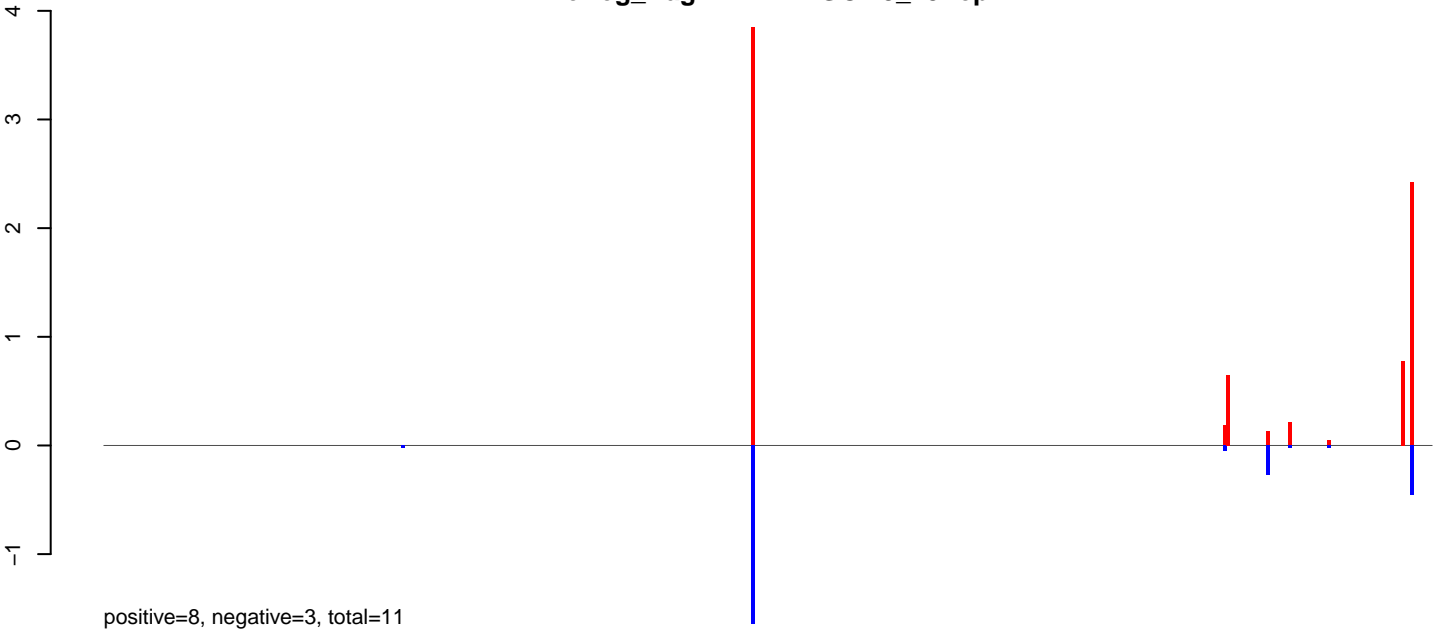


positive=4, negative=12, total=16

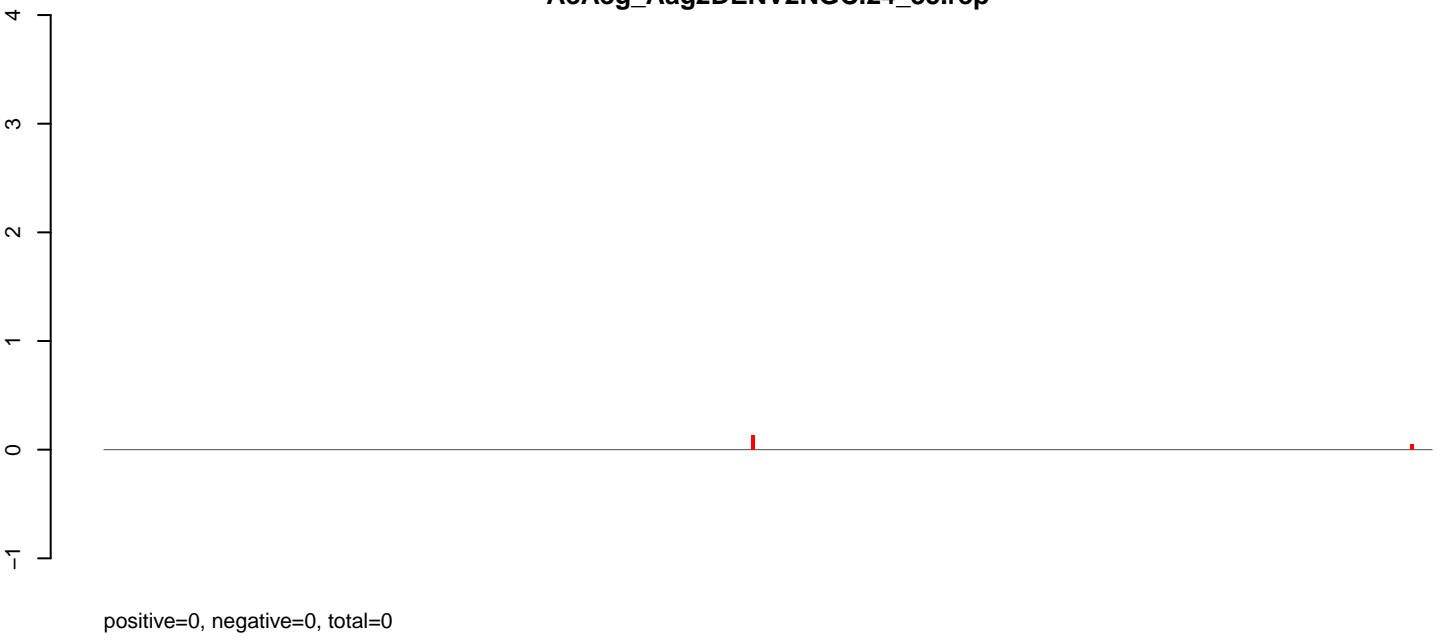
Window size=25, length=10755, virus@NC_009028.2-Ilheus_virus1-10755

0 2000 4000 6000 8000 10000

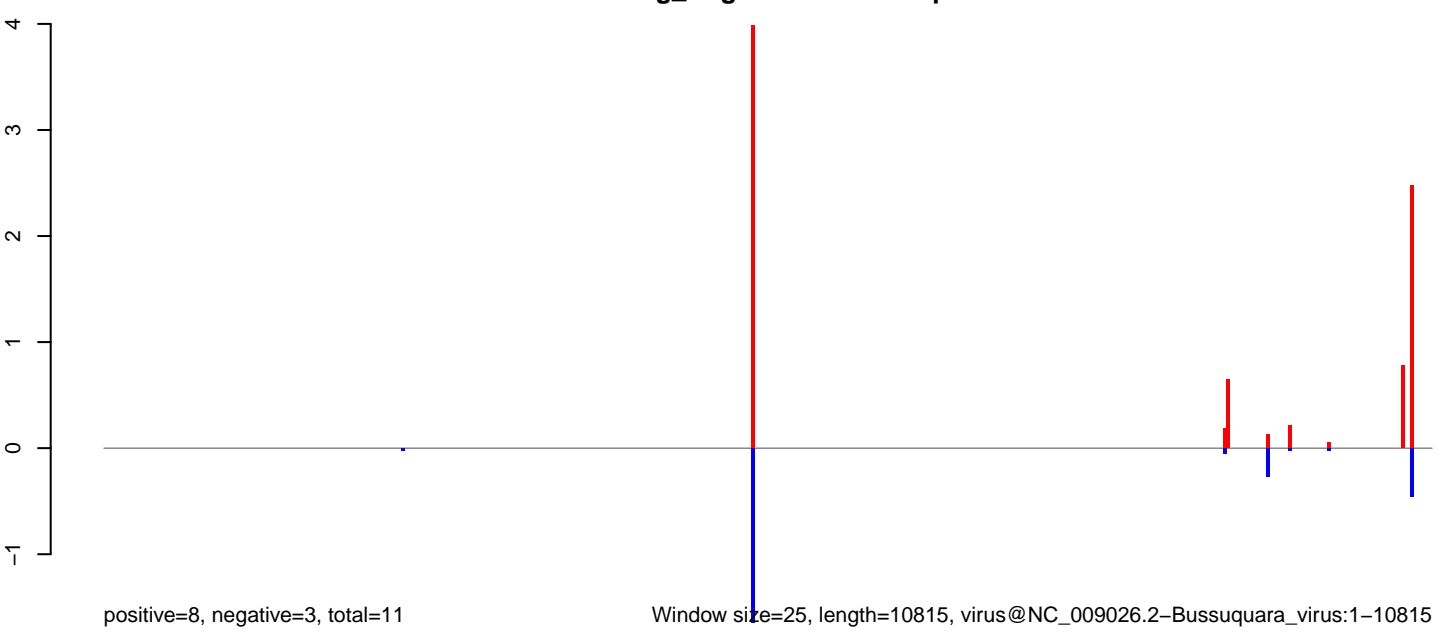
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

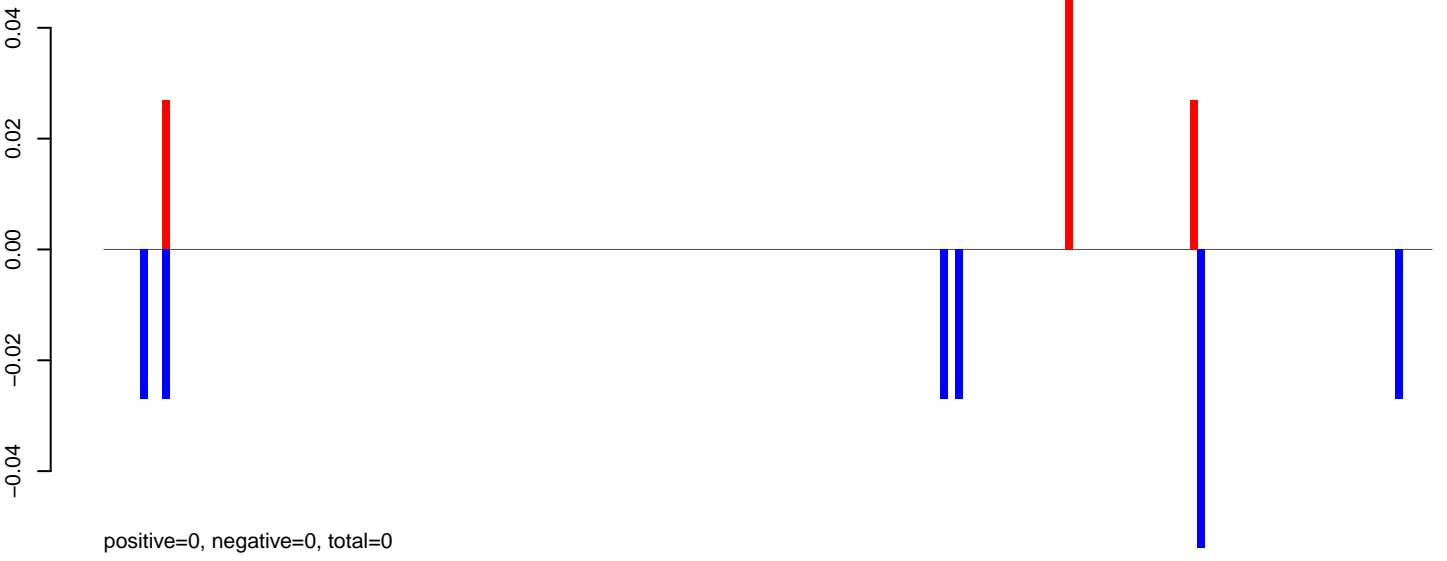


AeAeg_Aag2DENV2NGC.rep

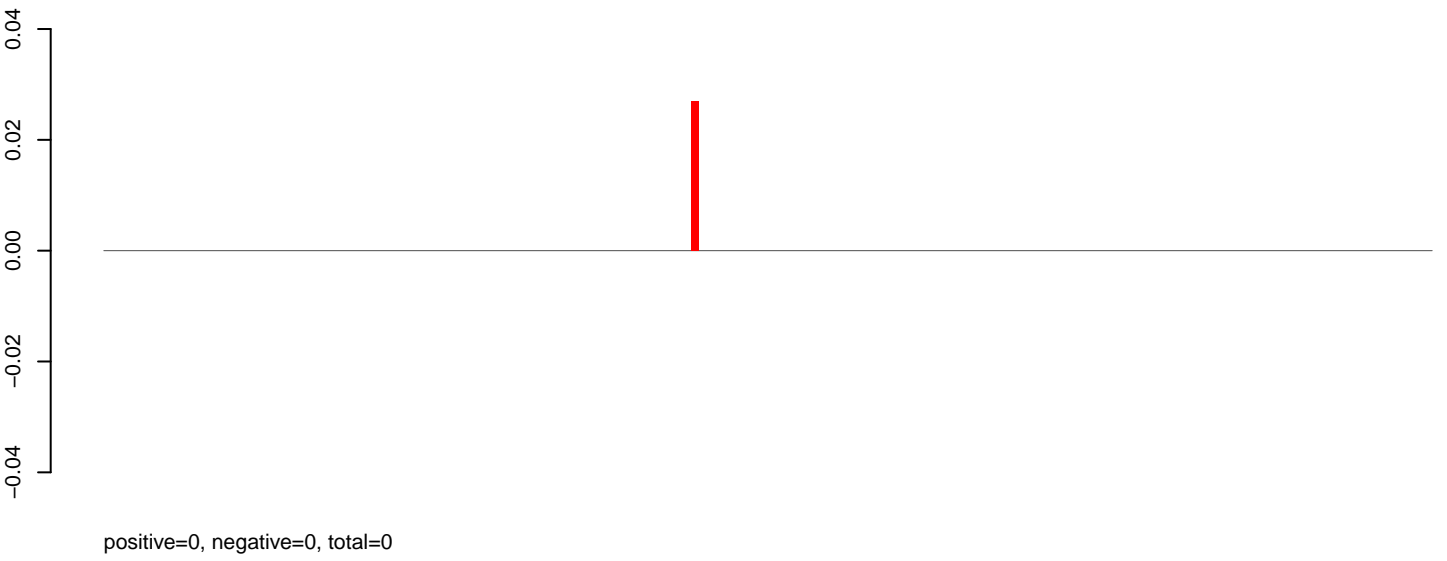


Window size=25, length=10815, virus@NC_009026.2-Bussuquara_virus:1-10815

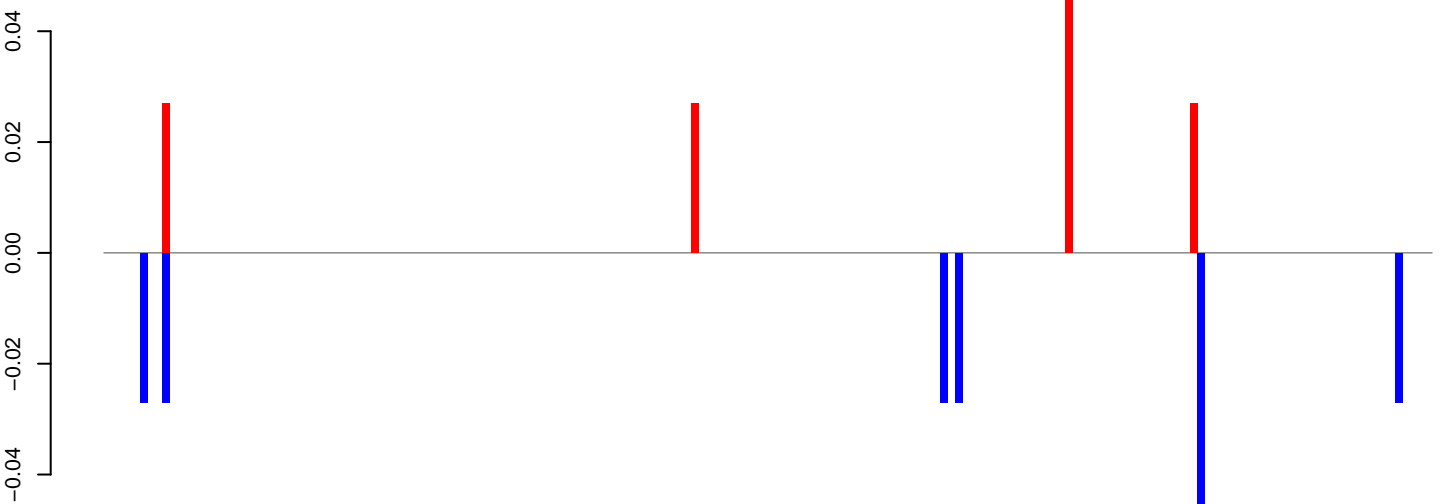
AeAeg_Aag2DENV2NGC.18_23.rep



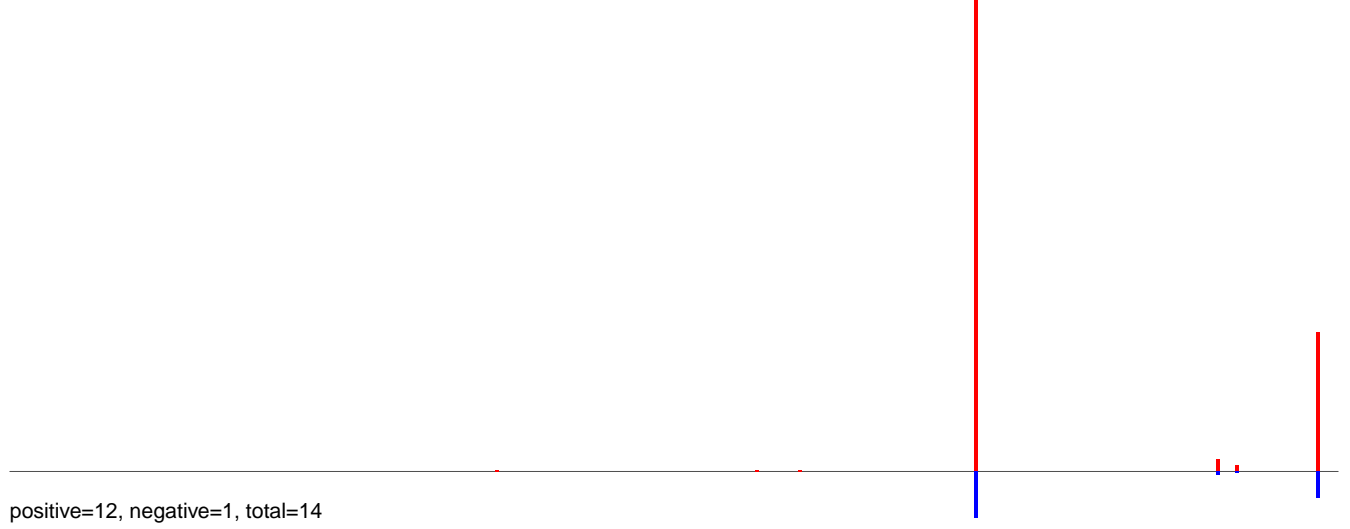
AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



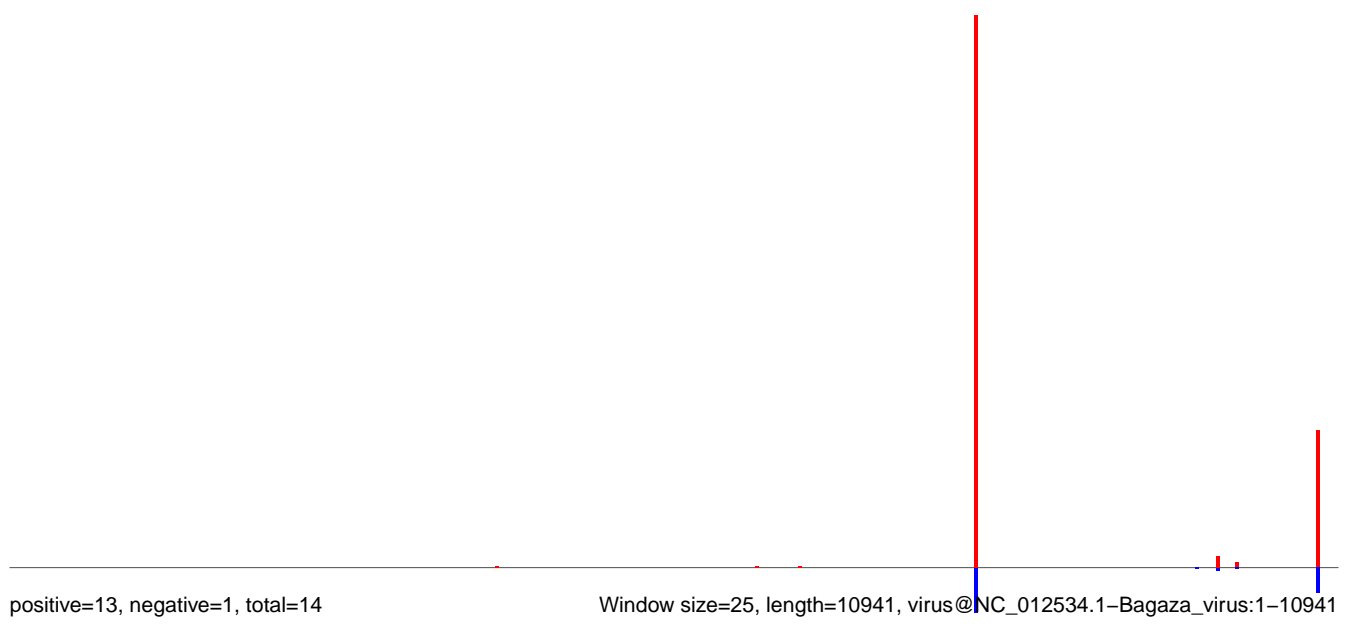
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



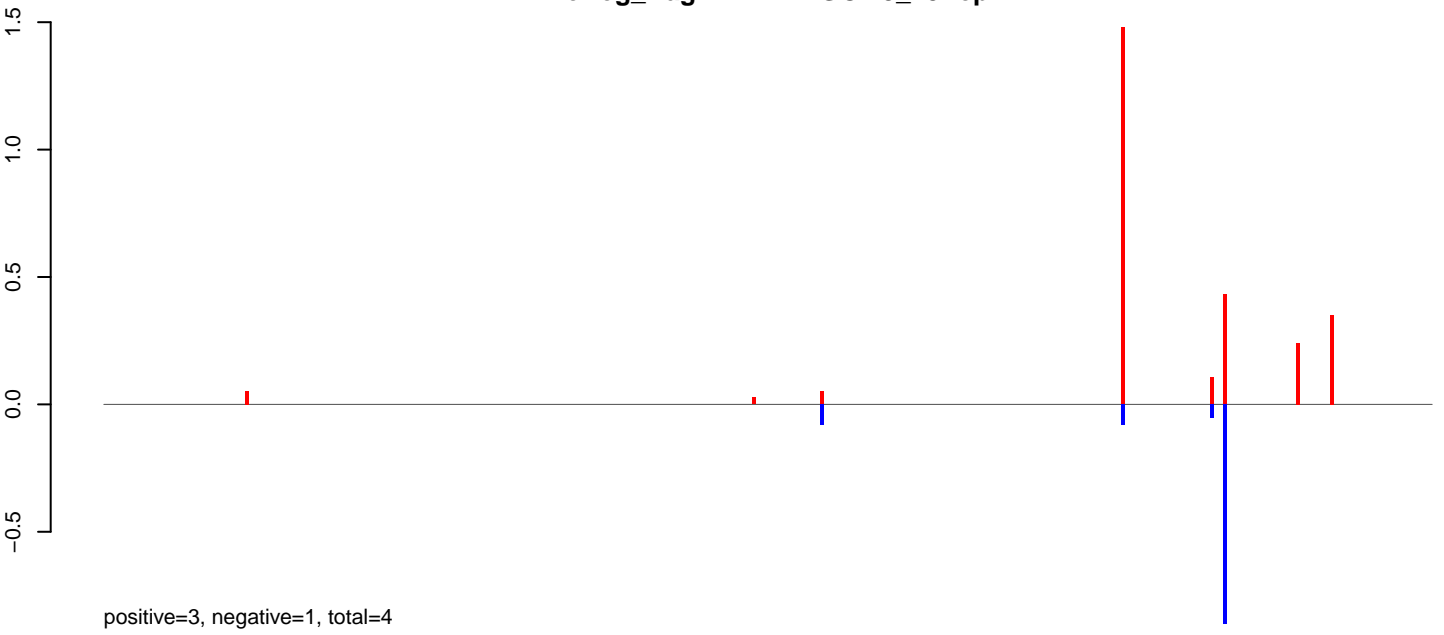
AeAeg_Aag2DENV2NGC.rep



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

0 2000 4000 6000 8000 10000

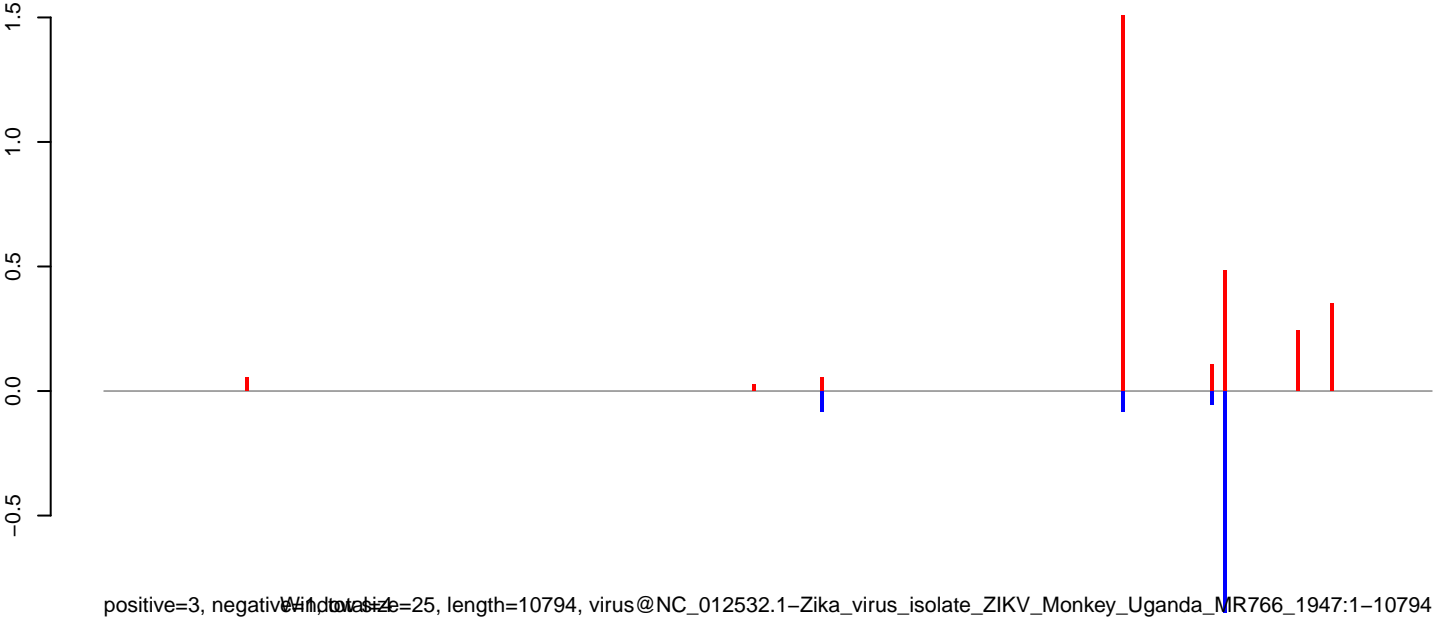
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



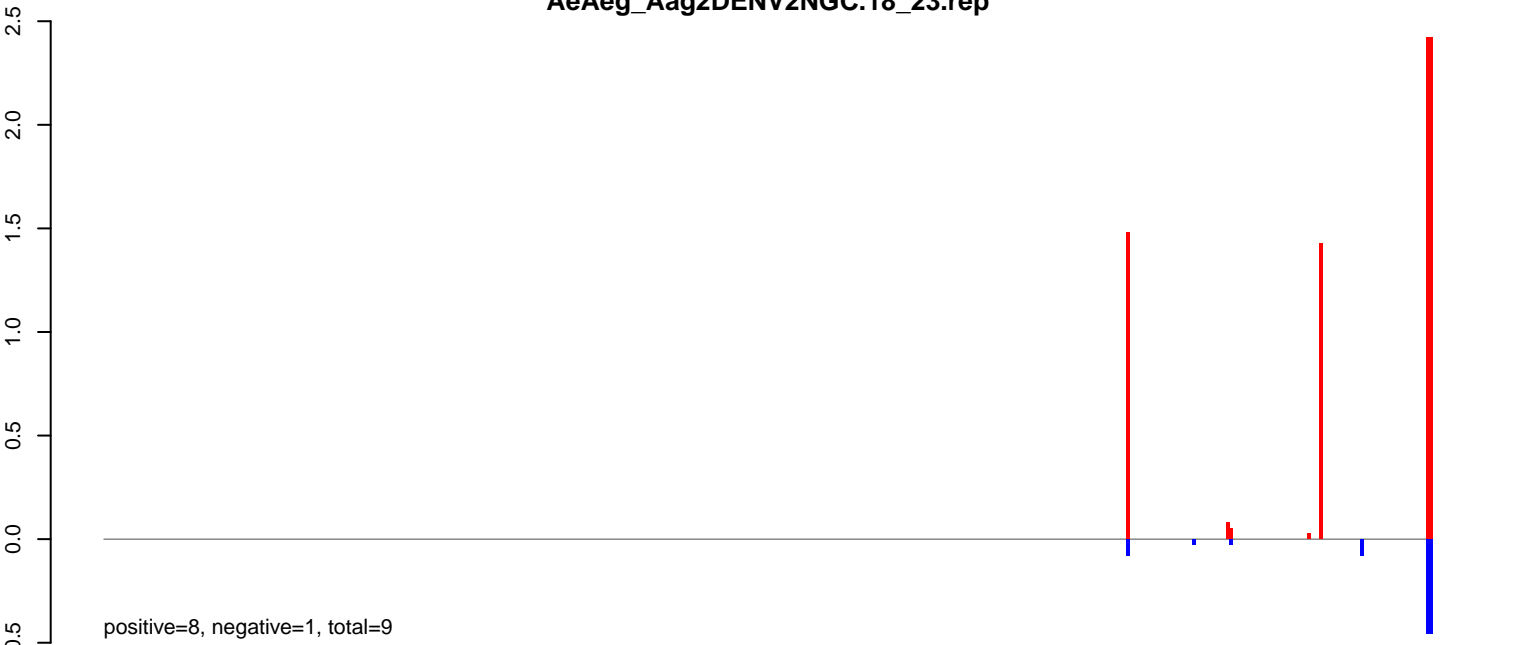
AeAeg_Aag2DENV2NGC.rep



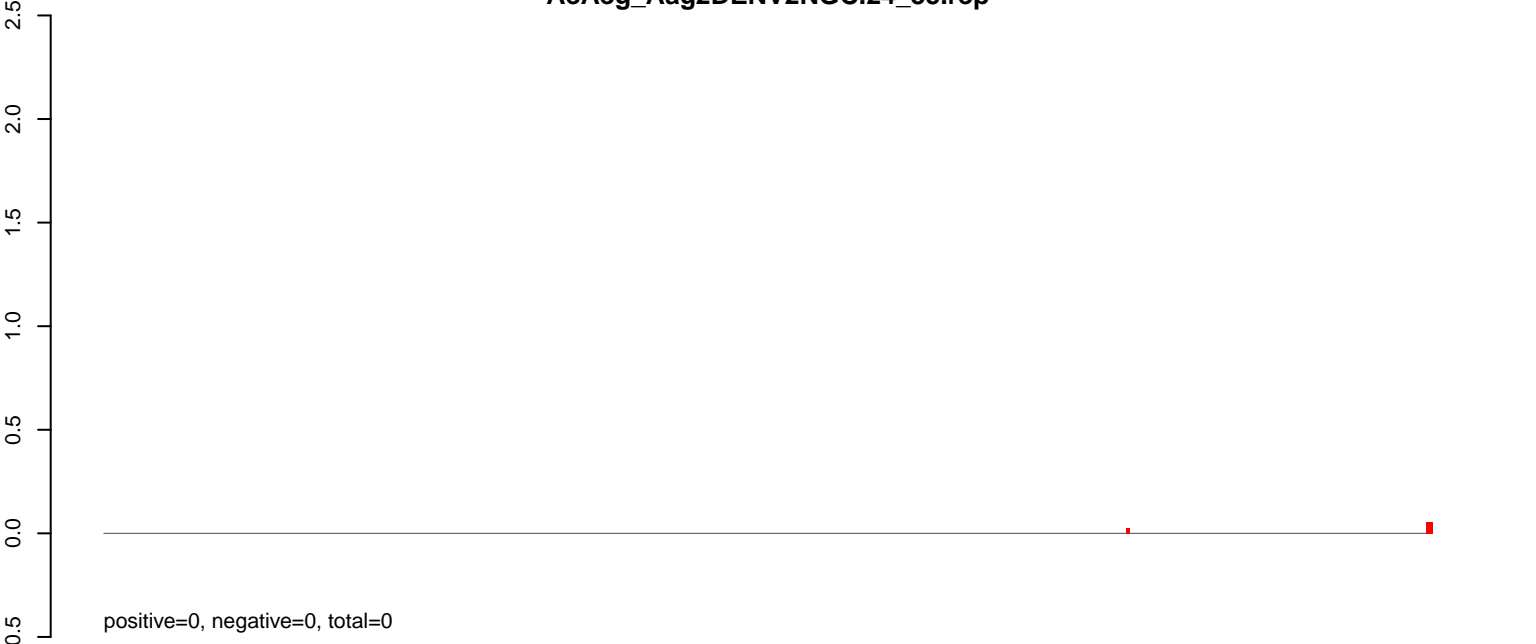
Window size=25, length=10794, virus@NC_012532.1-Zika_virus_isolate_ZIKV_Monkey_Uganda_MR766_1947:1-10794

0 2000 4000 6000 8000 10000

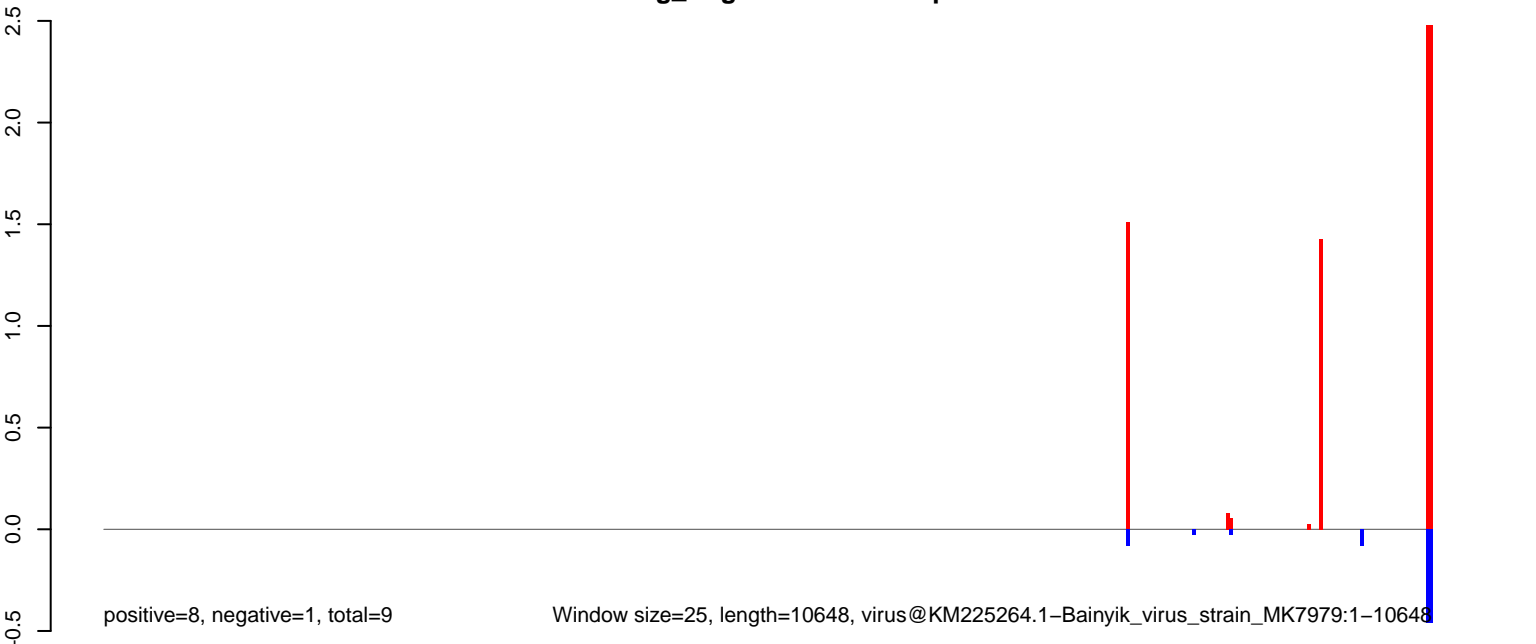
AeAeg_Aag2DENV2NGC.18_23.rep



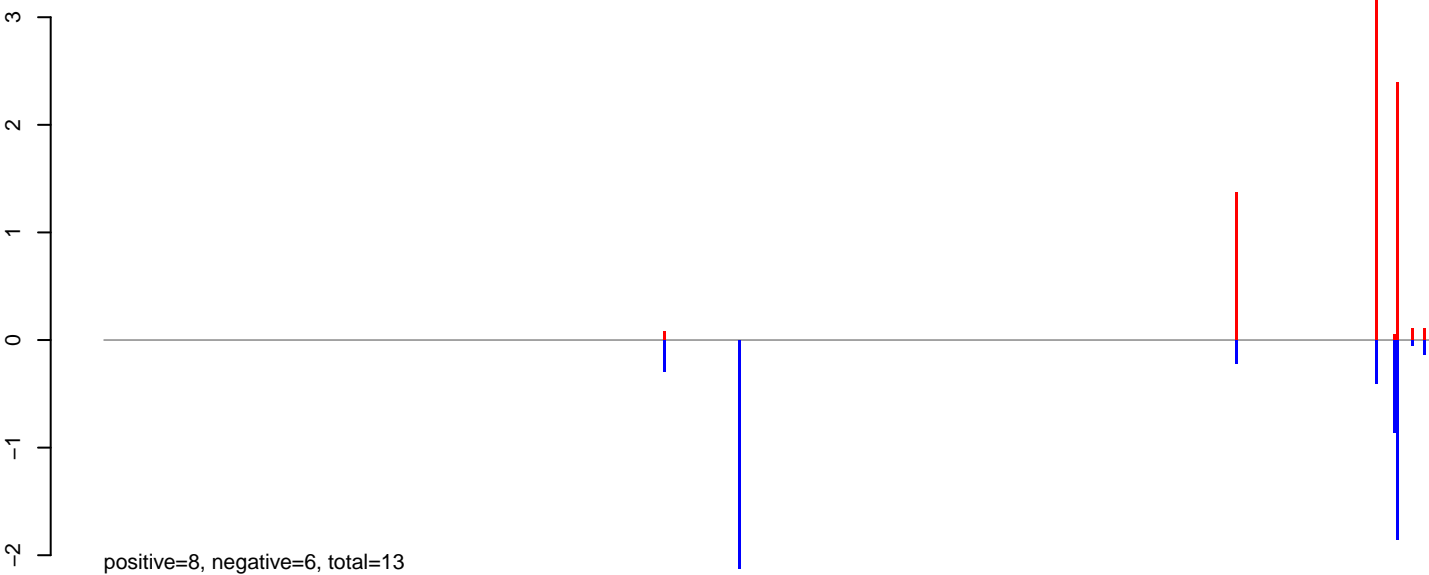
AeAeg_Aag2DENV2NGC.24_35.rep



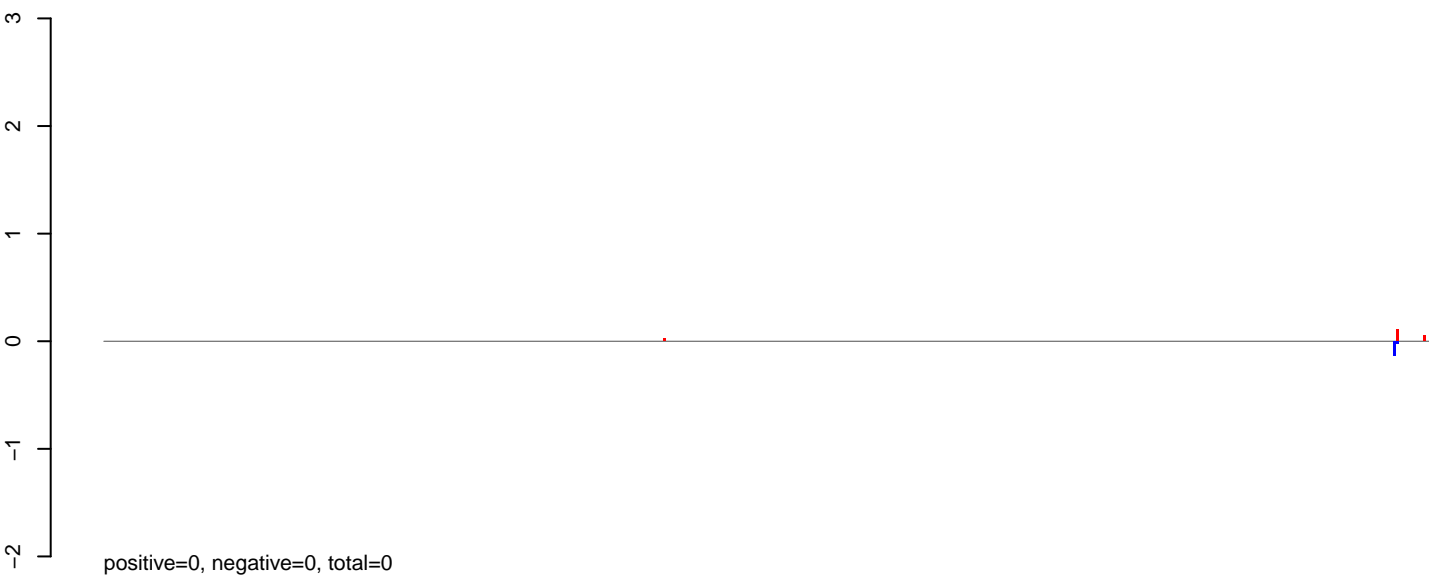
AeAeg_Aag2DENV2NGC.rep



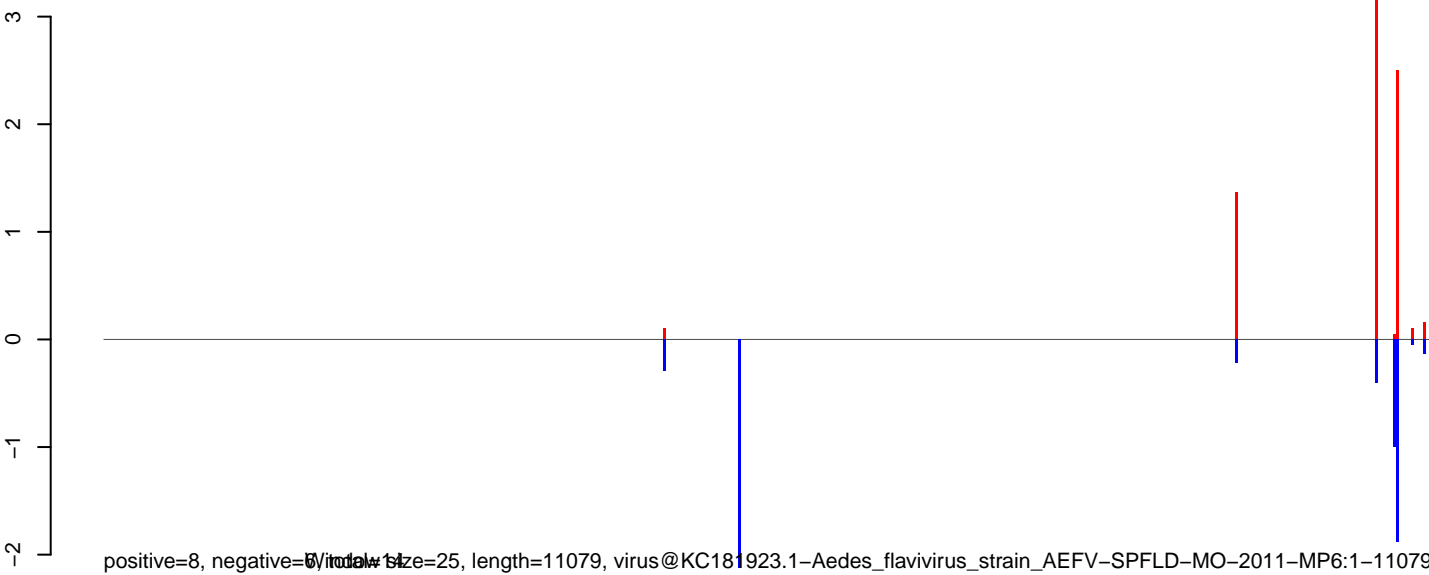
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



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

0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep

positive=3, negative=2, total=5

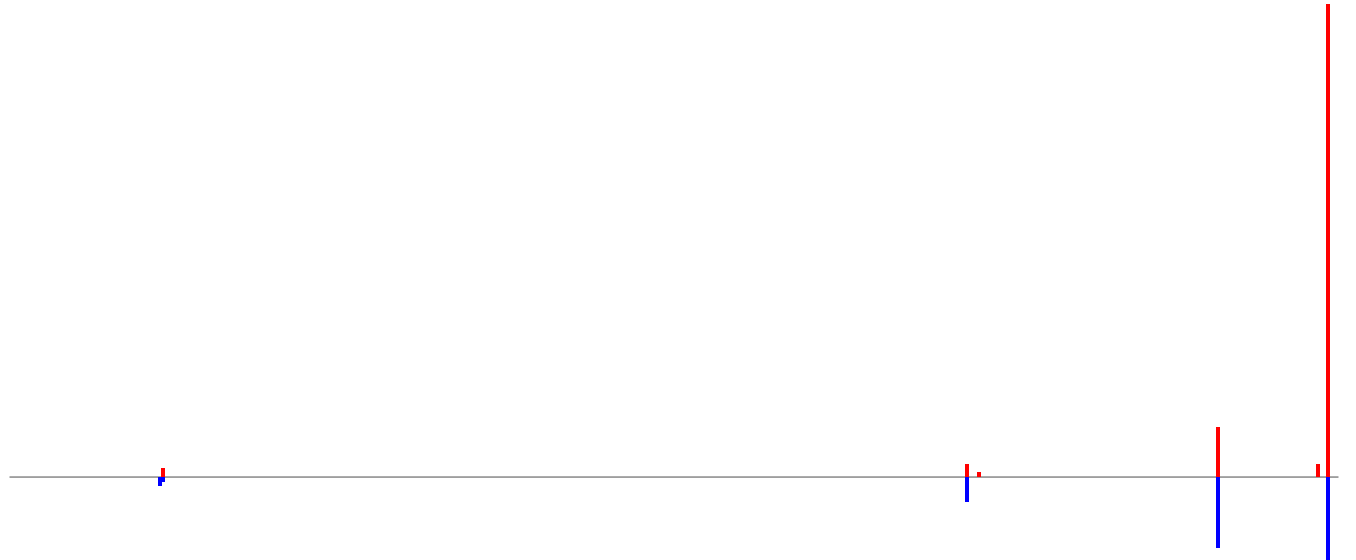
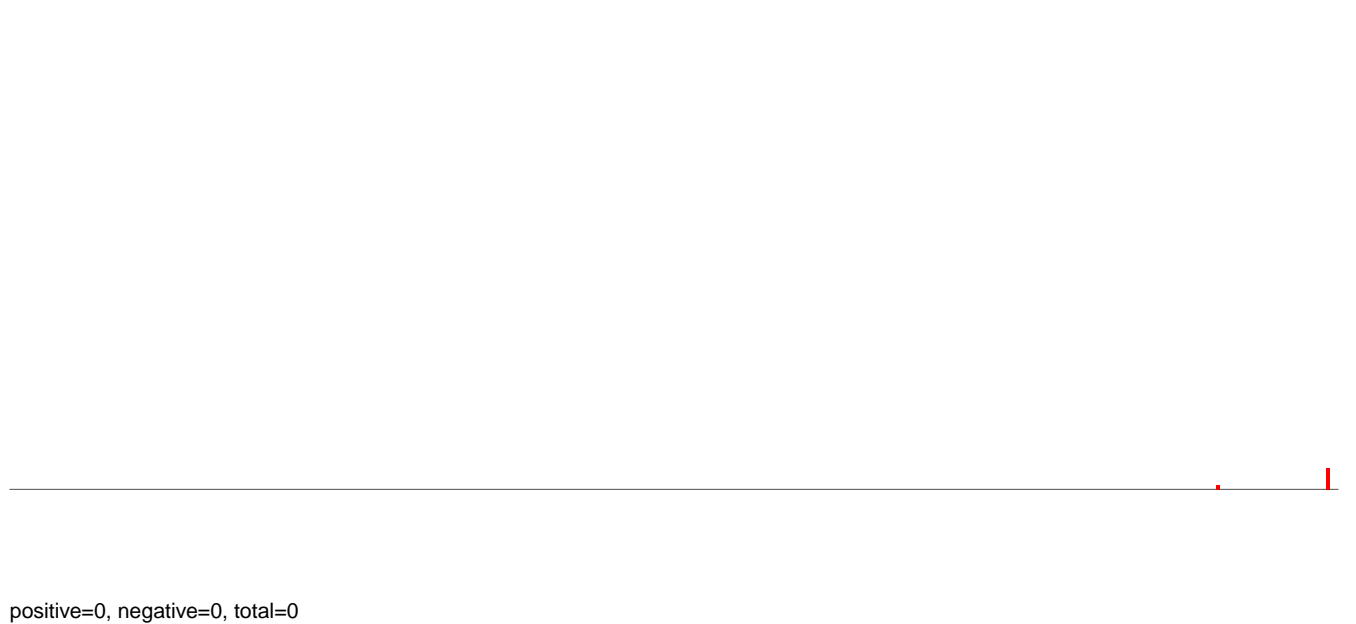
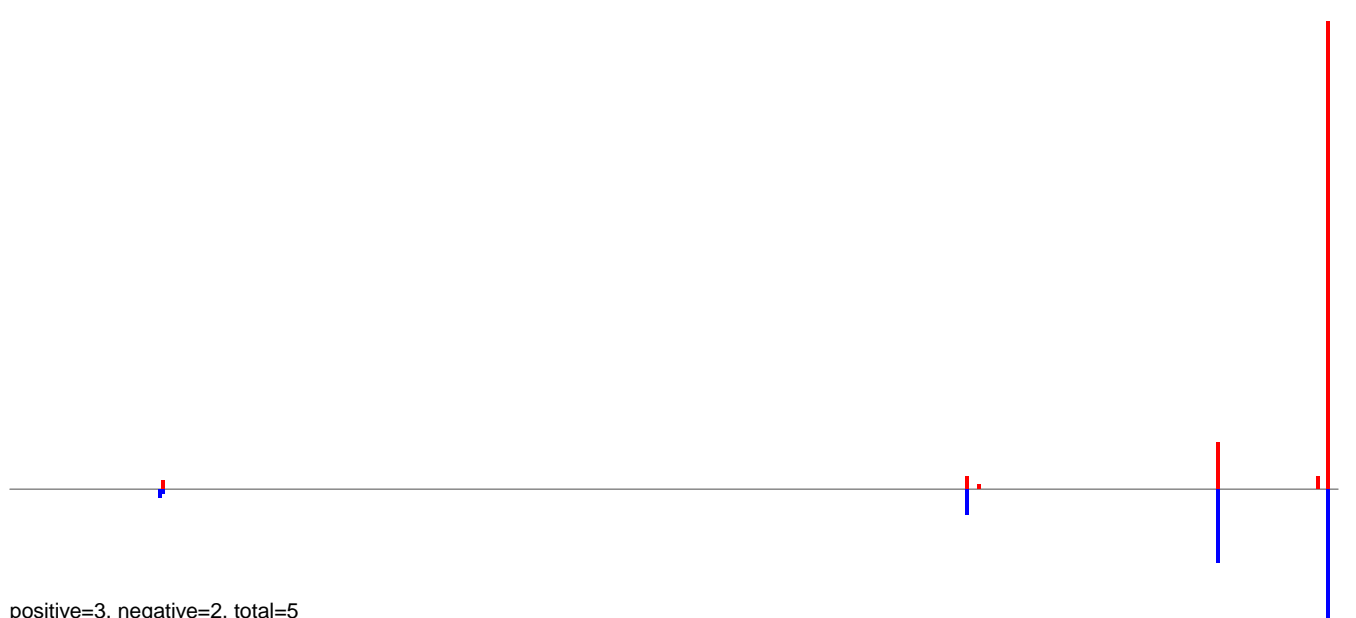
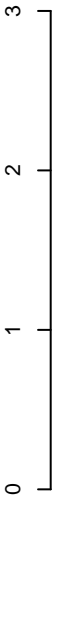
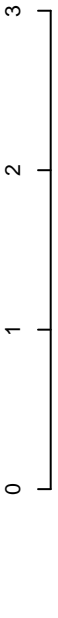
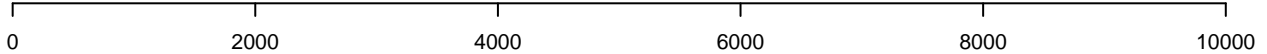
AeAeg_Aag2DENV2NGC.24_35.rep

positive=0, negative=0, total=0

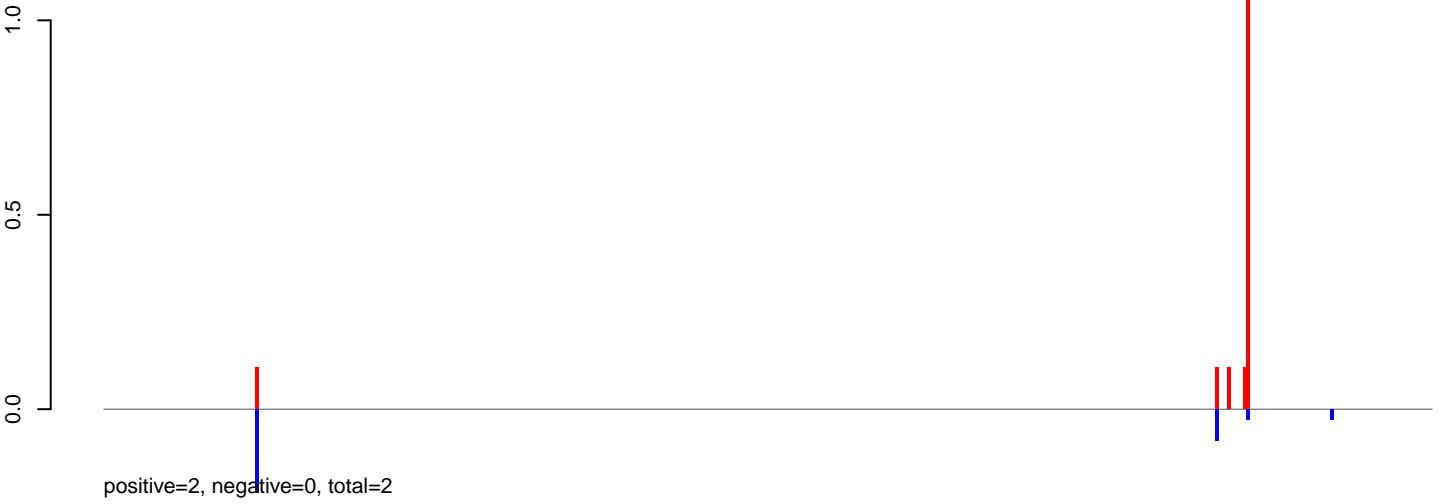
AeAeg_Aag2DENV2NGC.rep

positive=4, negative=2, total=5

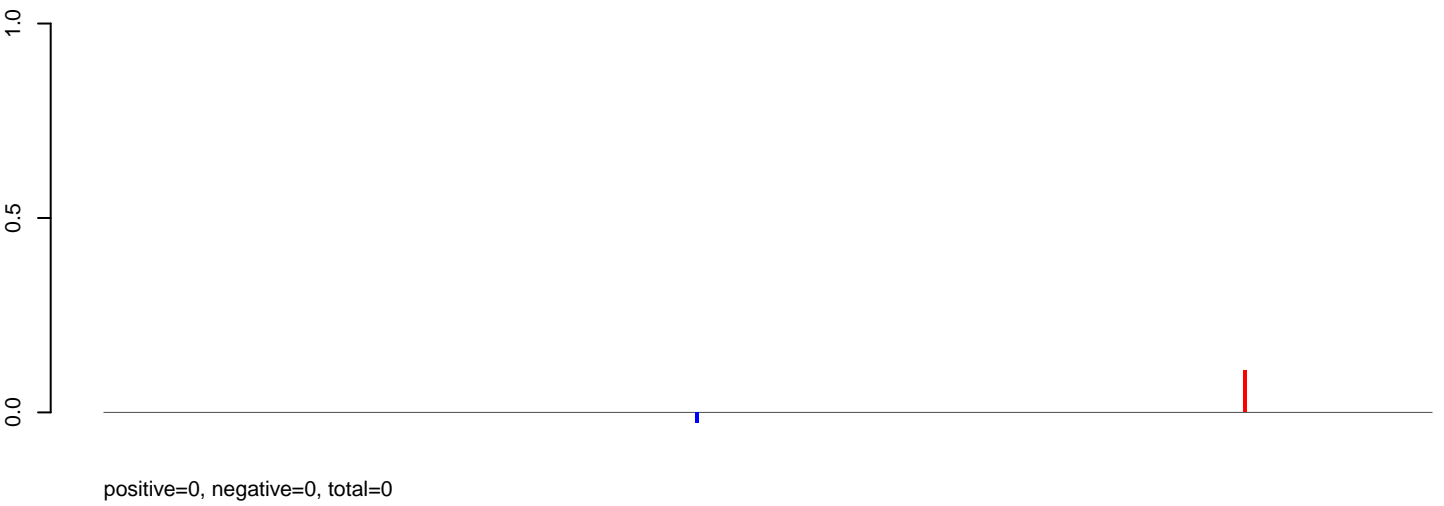
Window size=25, length=10943, virus@NC_018705.3-Ntaya_virus:1-10943



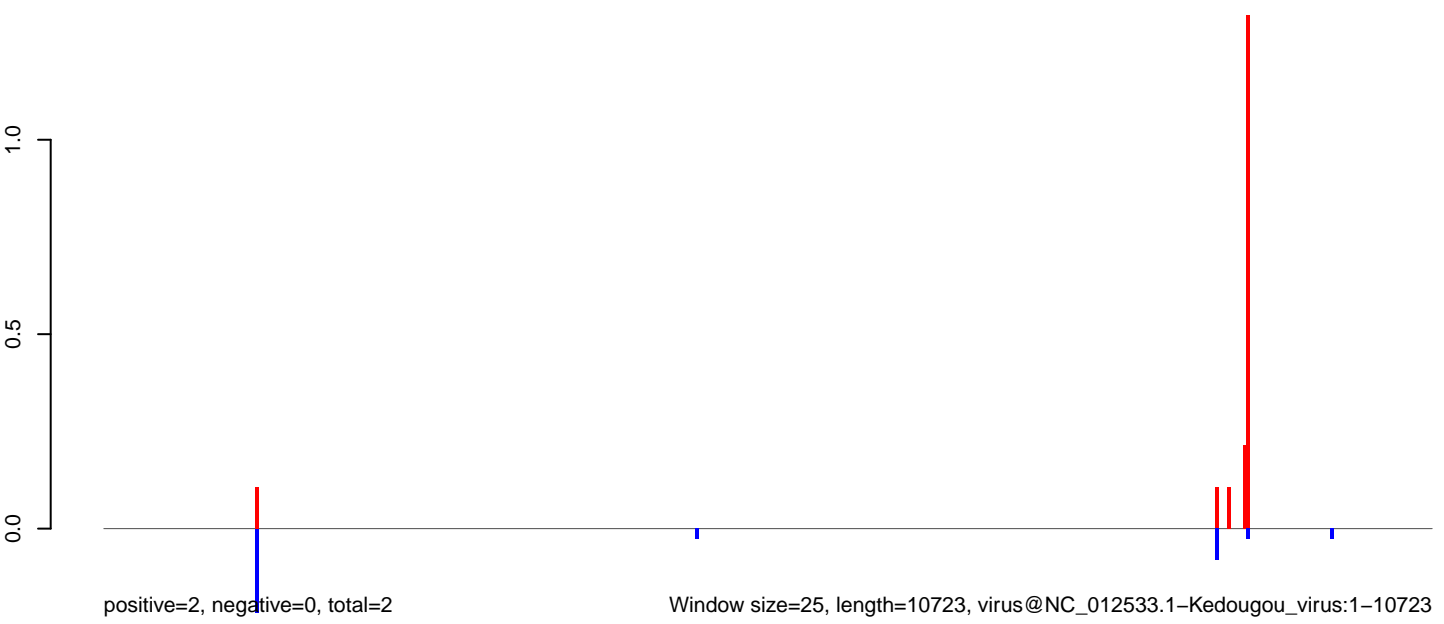
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



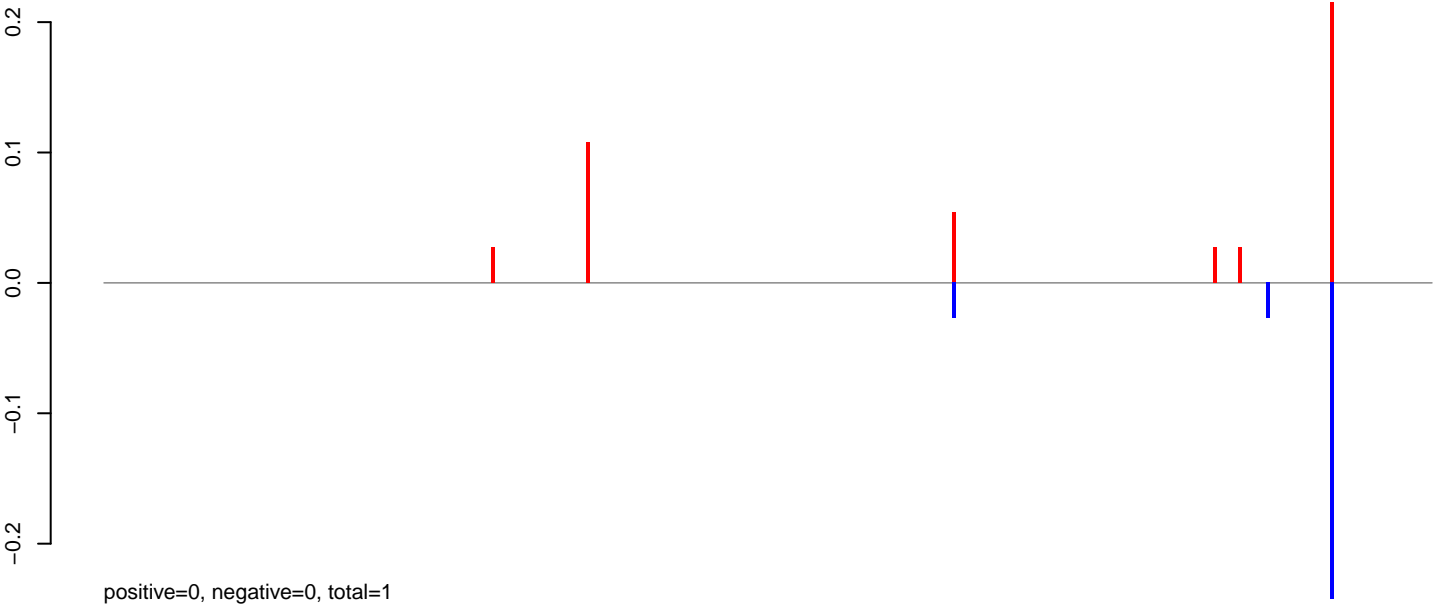
AeAeg_Aag2DENV2NGC.rep



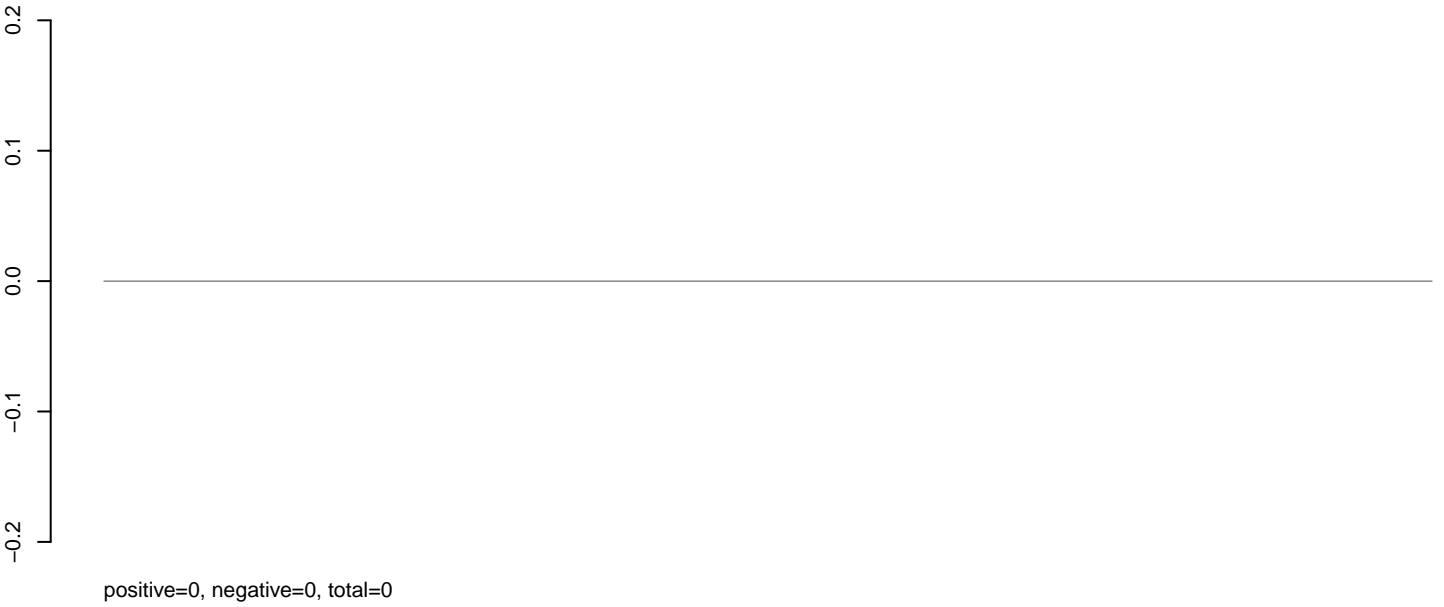
Window size=25, length=10723, virus@NC_012533.1-Kedougou_virus:1-10723

0 2000 4000 6000 8000 10000

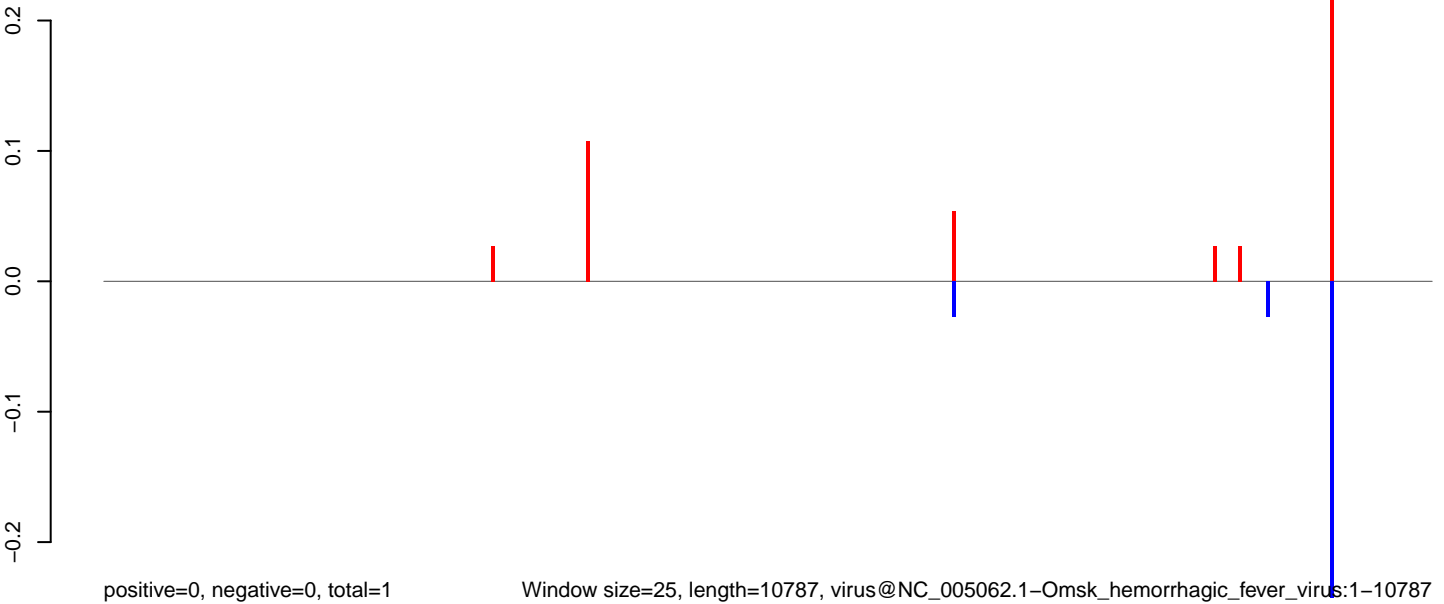
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



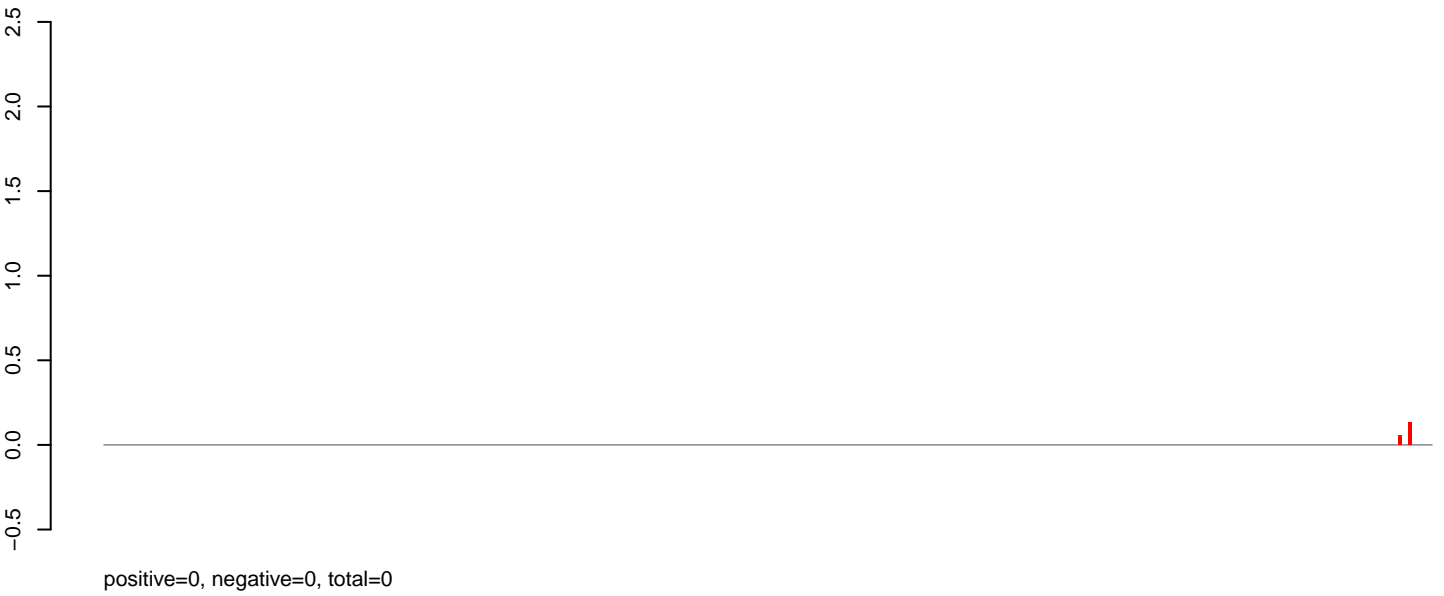
AeAeg_Aag2DENV2NGC.rep



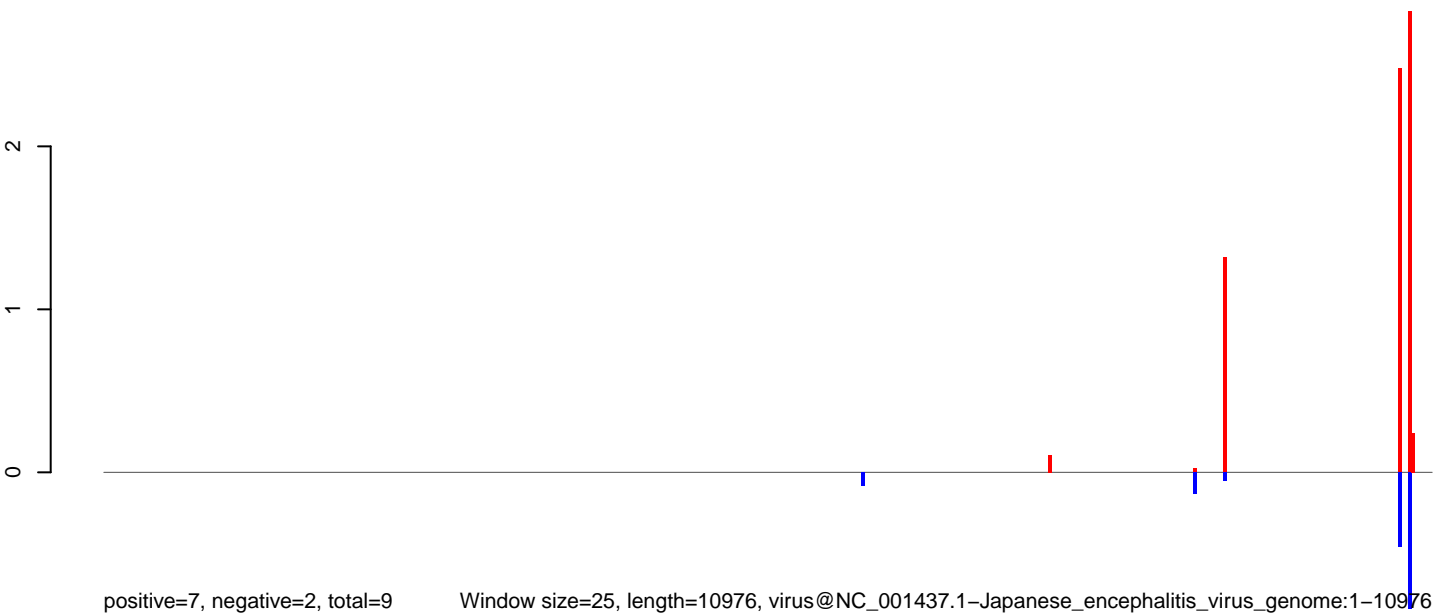
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



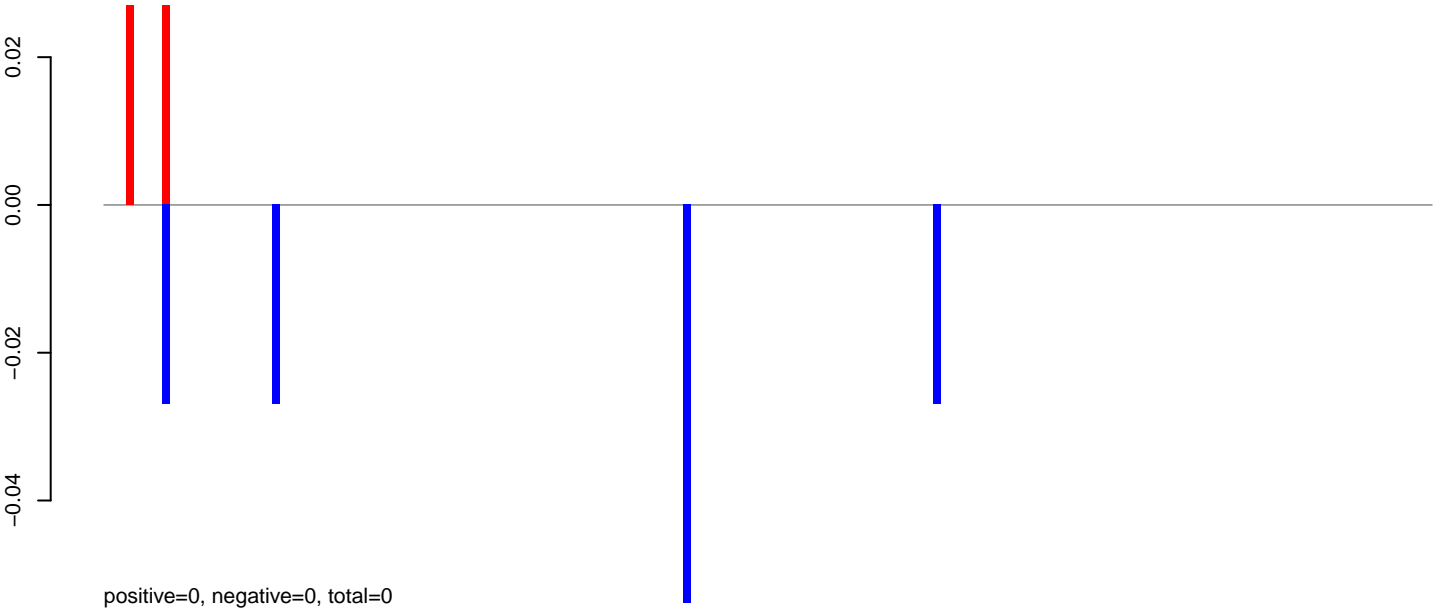
AeAeg_Aag2DENV2NGC.rep



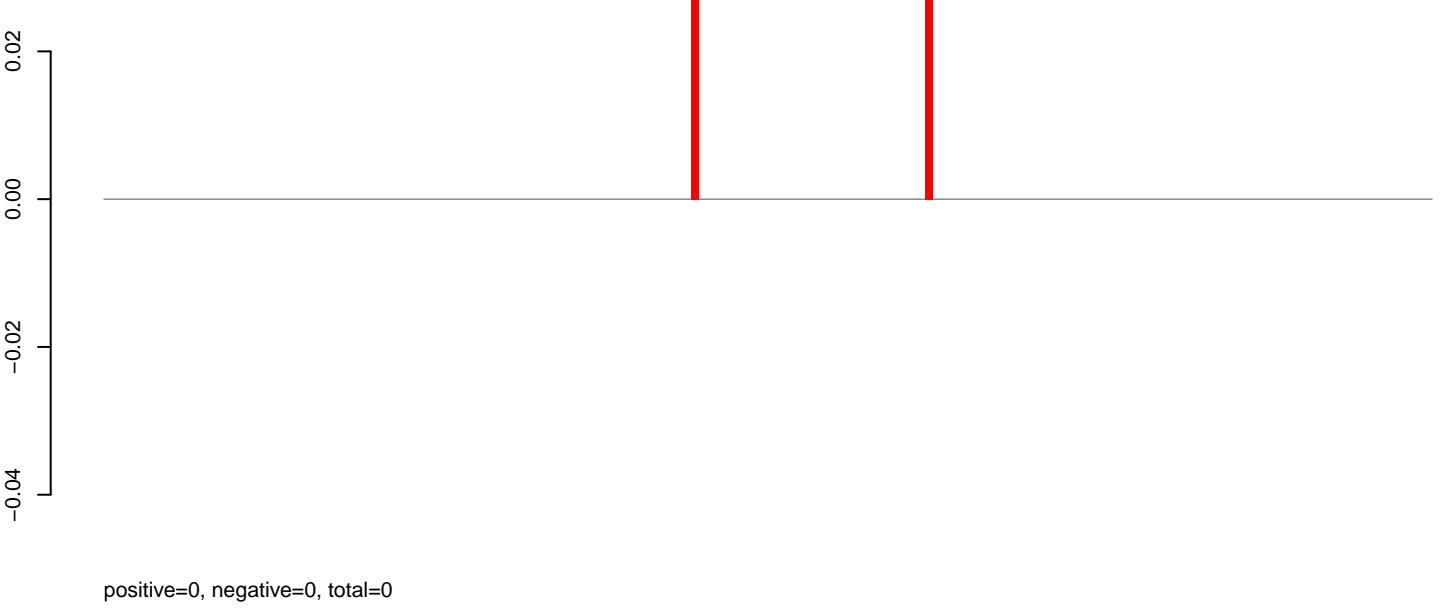
Window size=25, length=10976, virus@NC_001437.1-Japanese_encephalitis_virus_genome:1-10976

0 2000 4000 6000 8000 10000

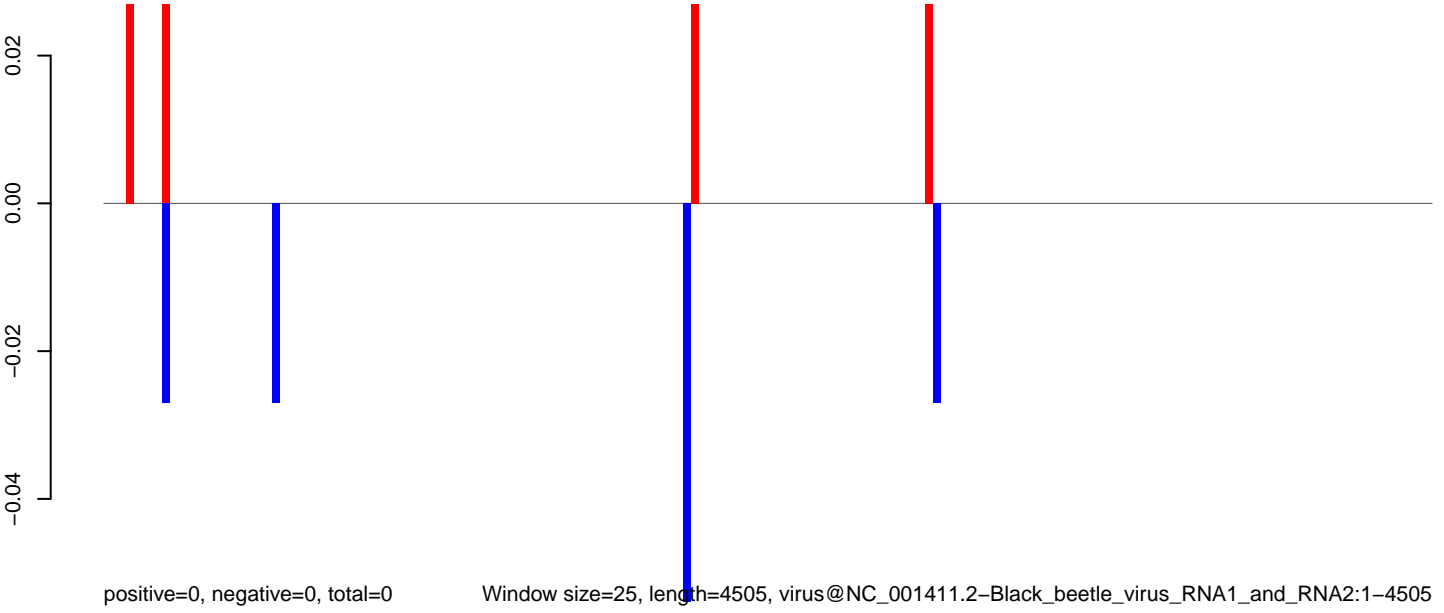
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



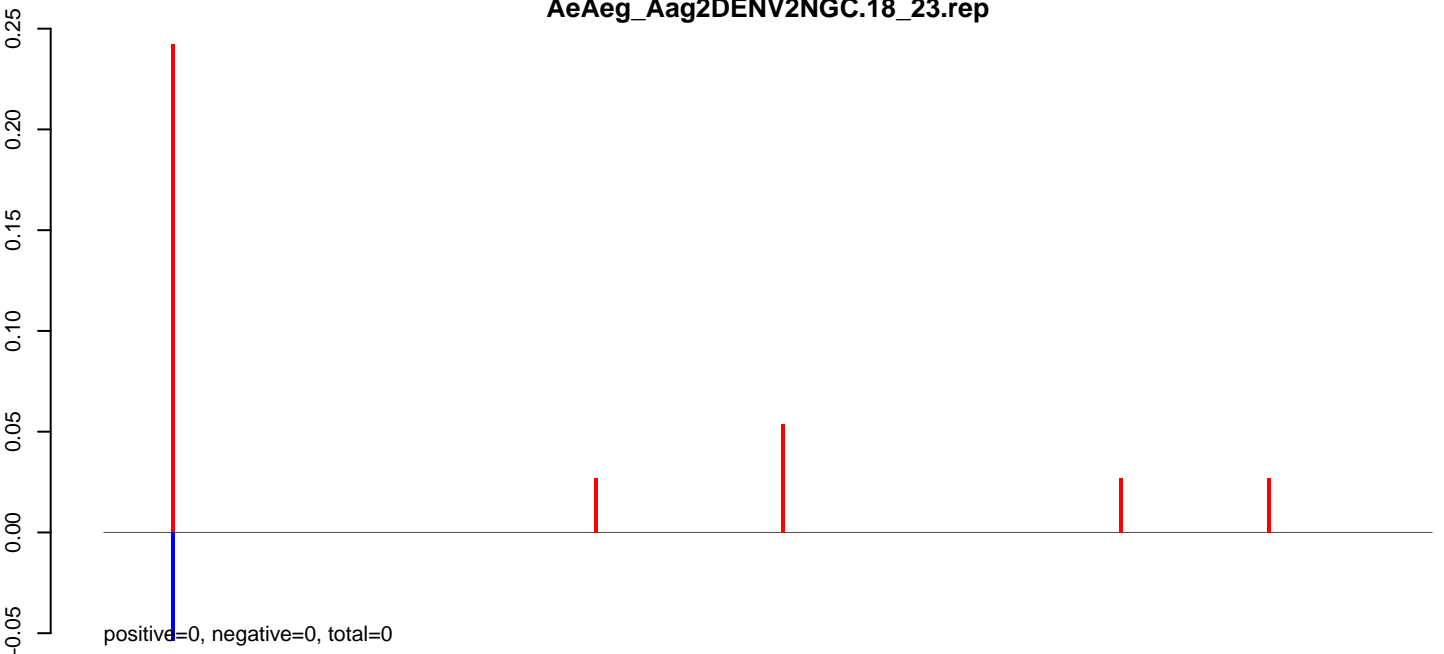
AeAeg_Aag2DENV2NGC.rep



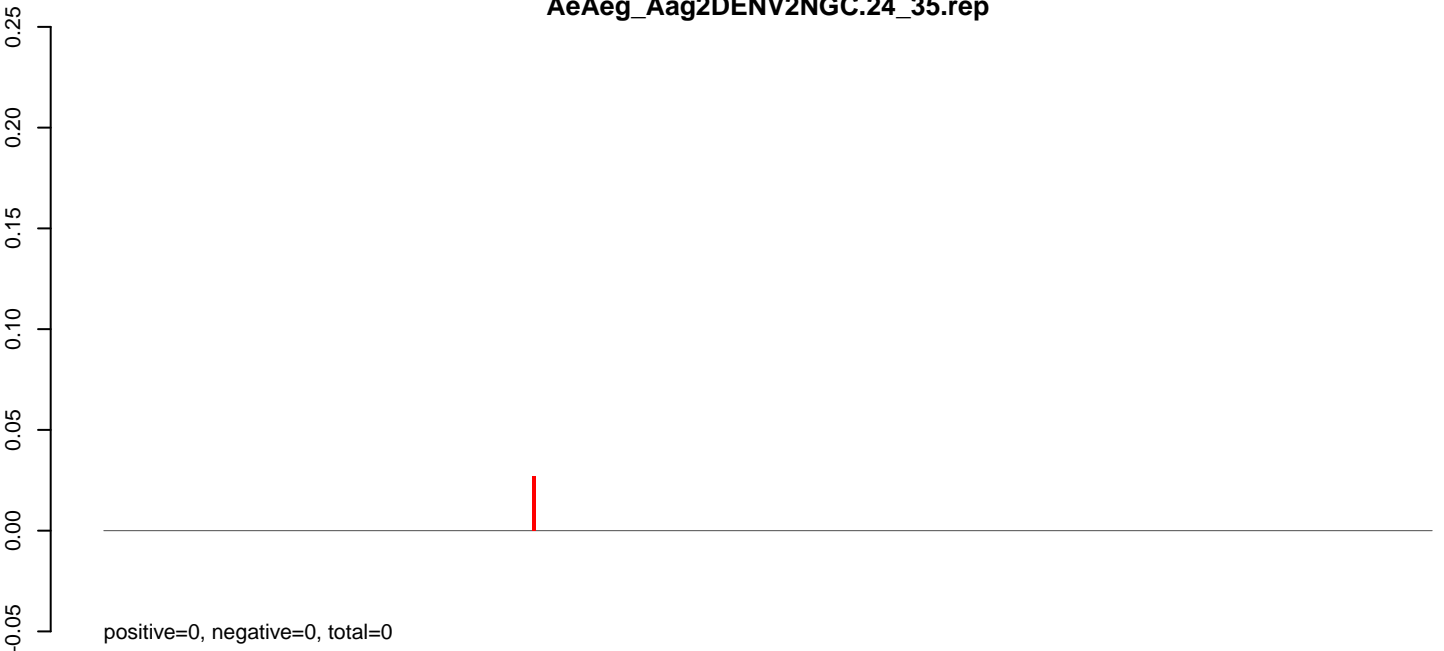
Window size=25, length=4505, virus@NC_001411.2-Black_beetle_virus_RNA1_and_RNA2:1-4505

0 1000 2000 3000 4000

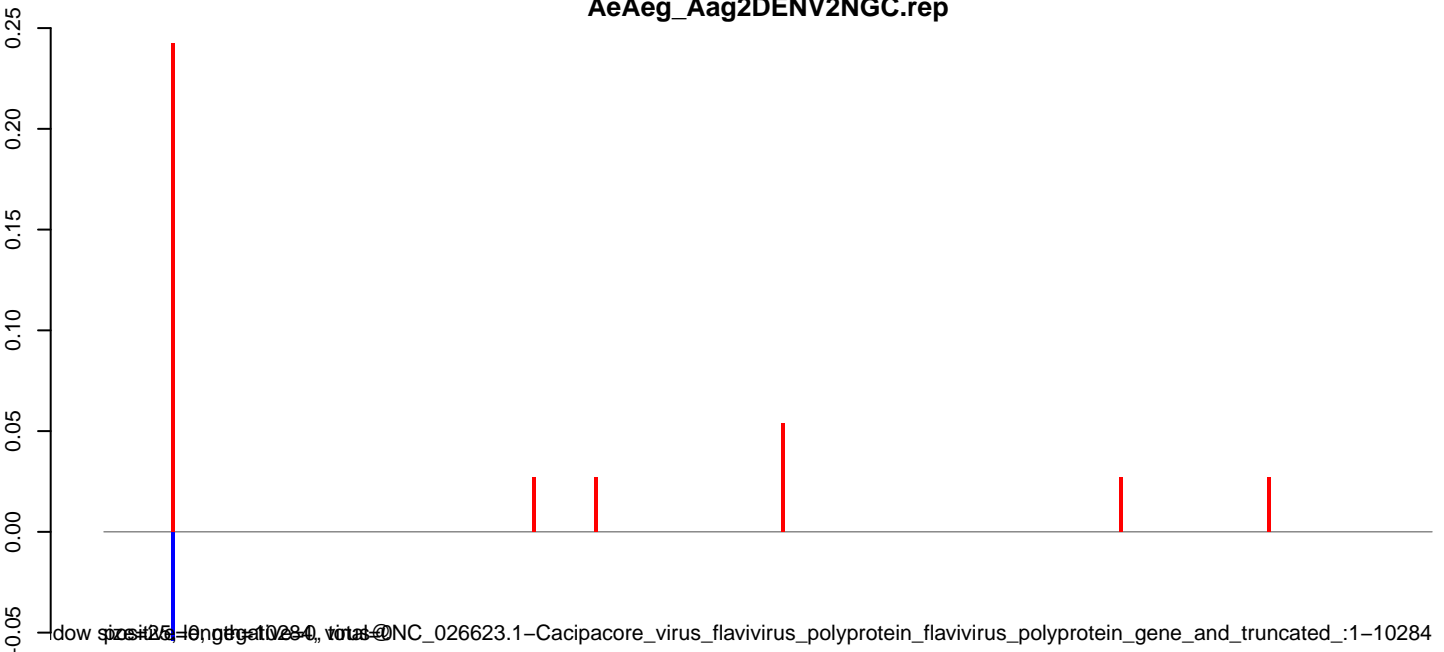
AeAeg_Aag2DENV2NGC.18_23.rep



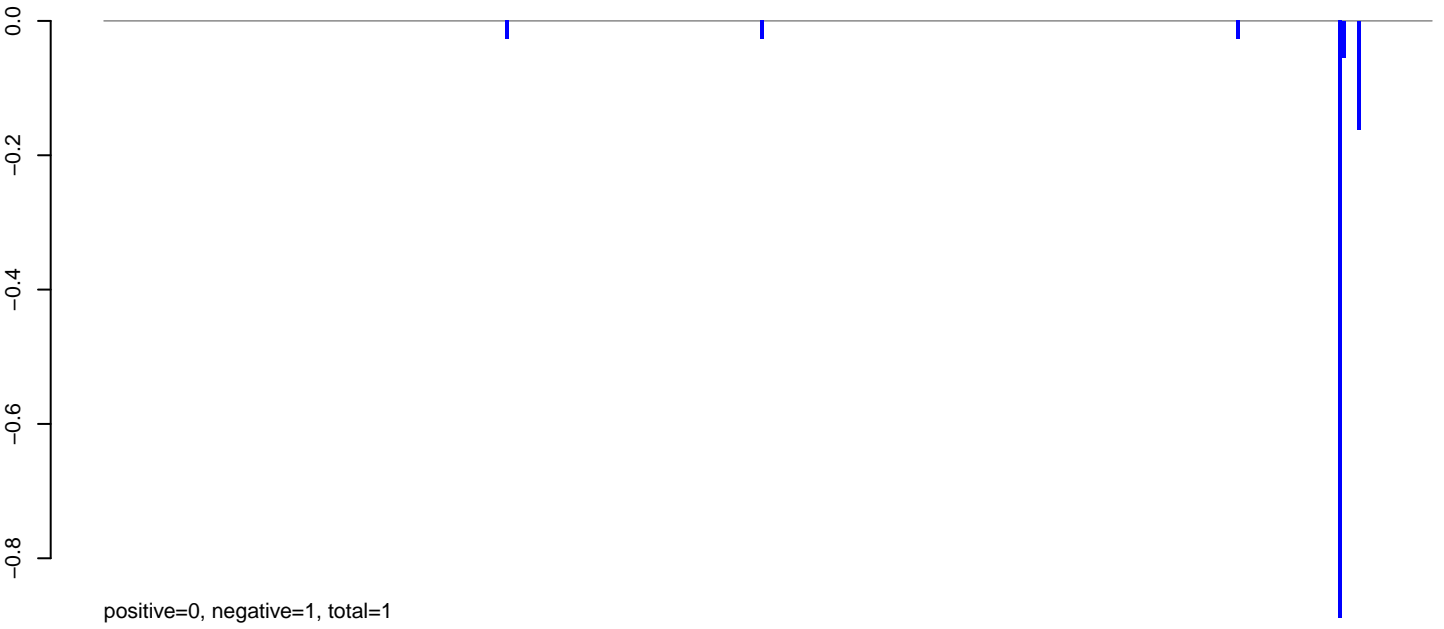
AeAeg_Aag2DENV2NGC.24_35.rep



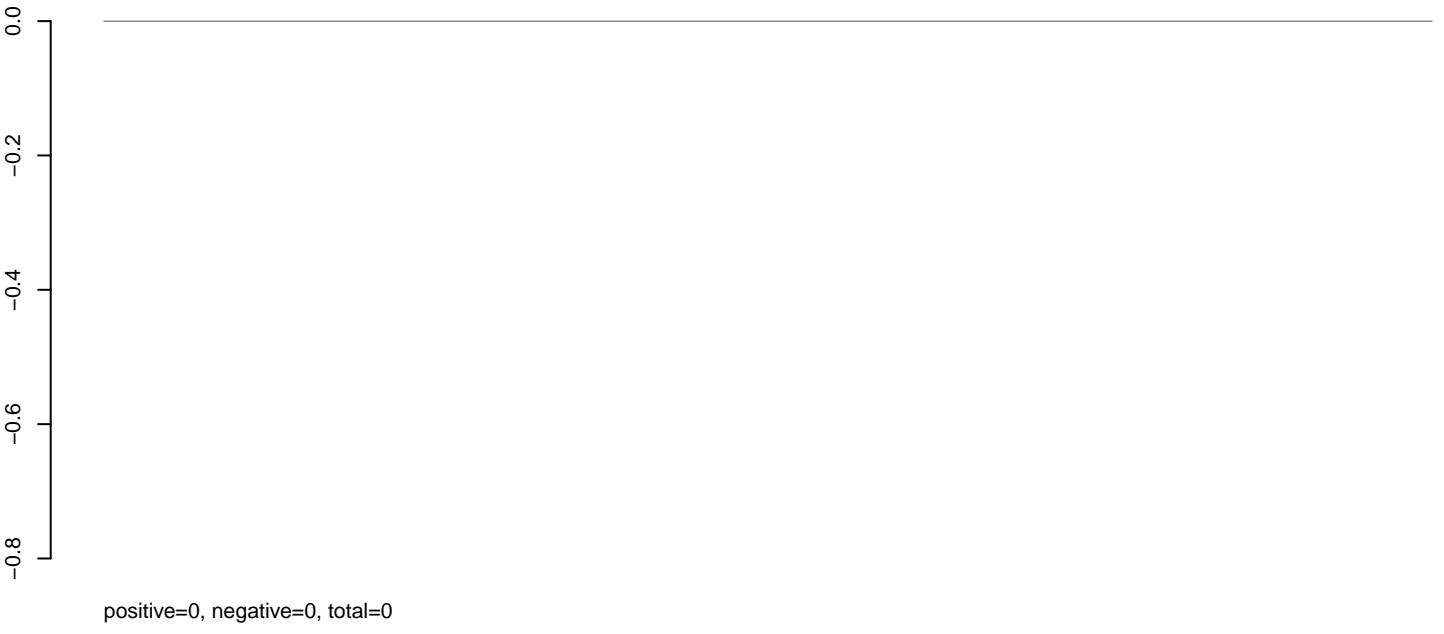
AeAeg_Aag2DENV2NGC.rep



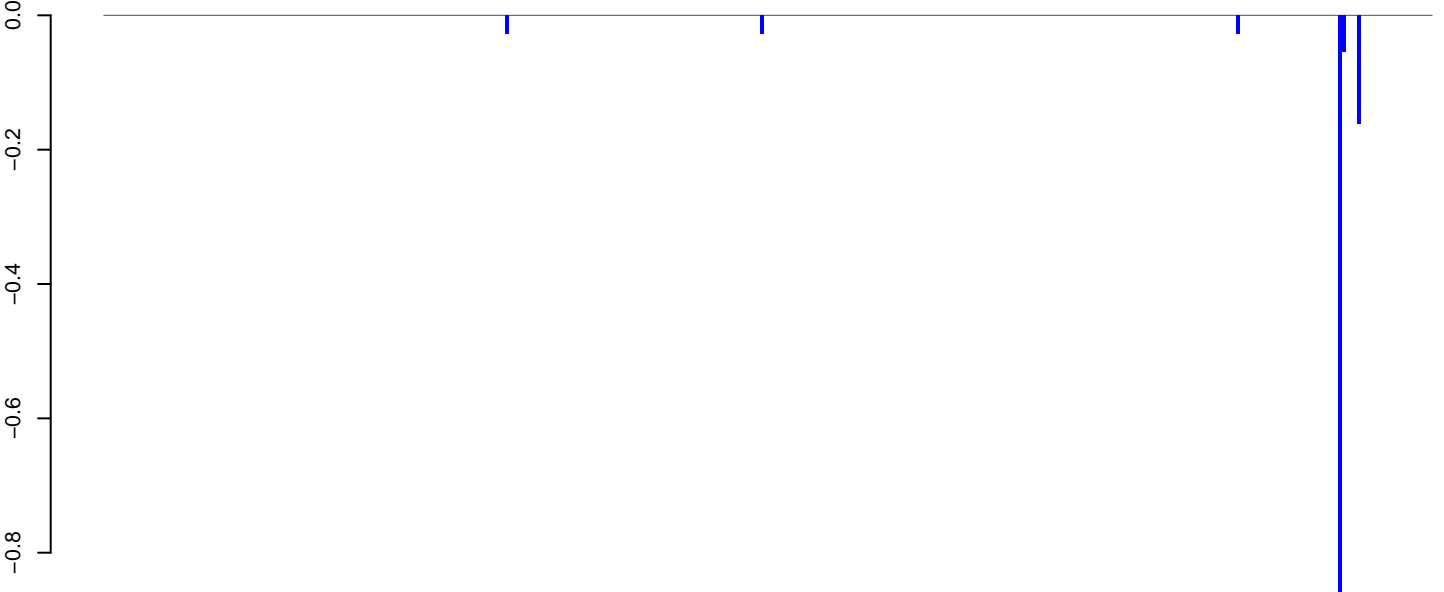
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



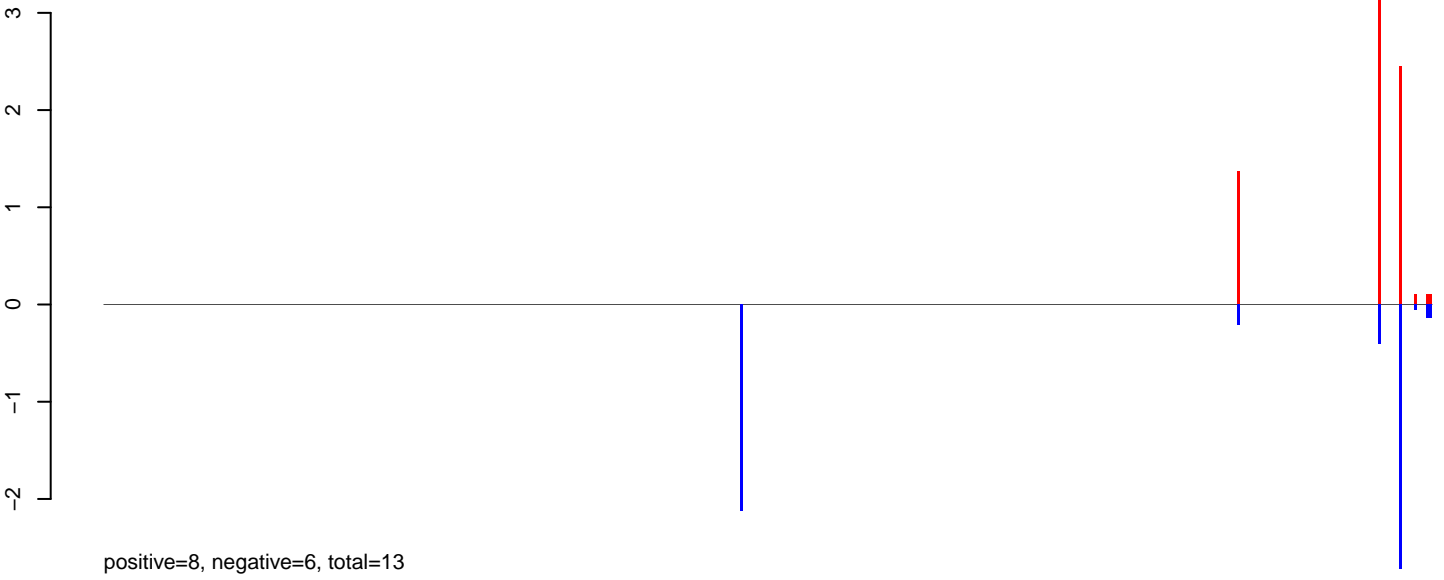
AeAeg_Aag2DENV2NGC.rep



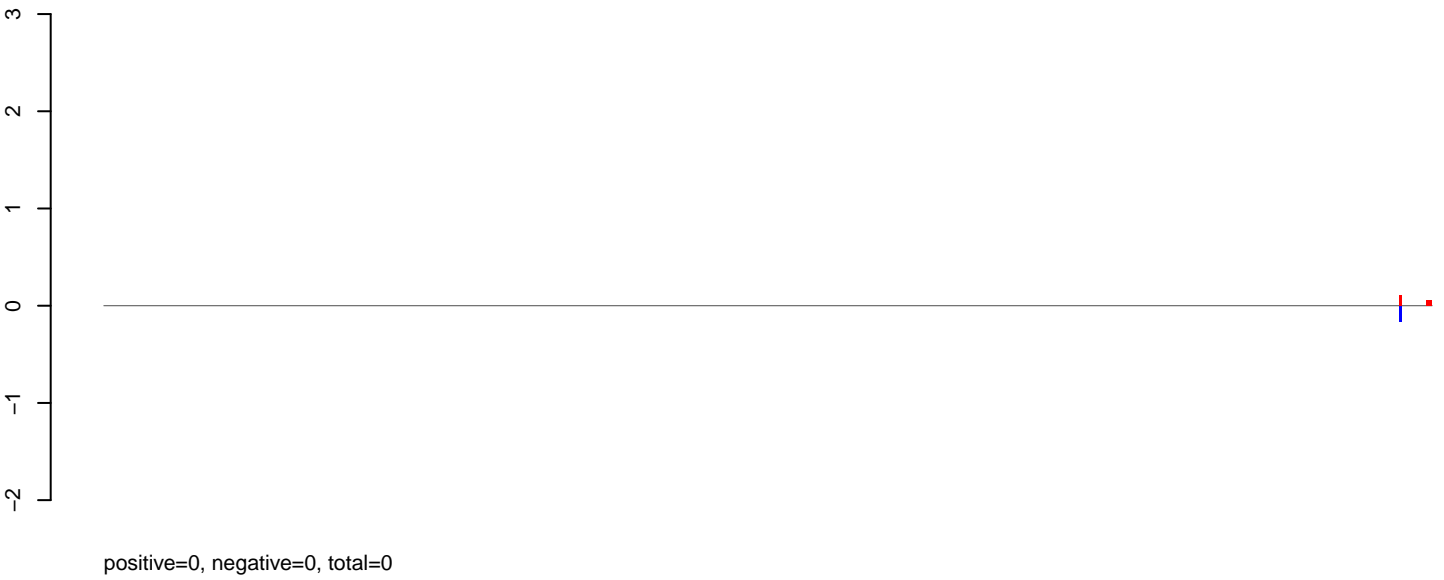
Window size=25, length=10673, virus@NC_023424.1-Tyuleniy_virus_strain_LEIV-6C_polyprotein_gene:1-10673

0 2000 4000 6000 8000 10000

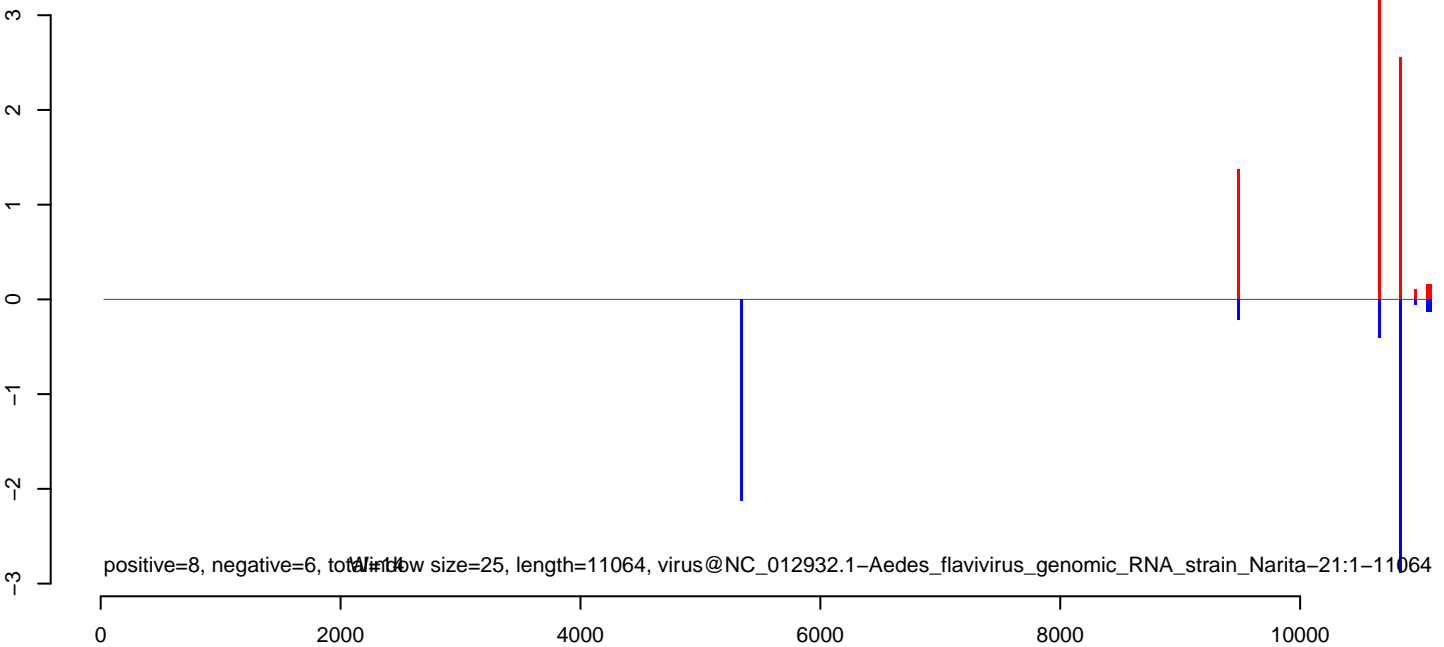
AeAeg_Aag2DENV2NGC.18_23.rep



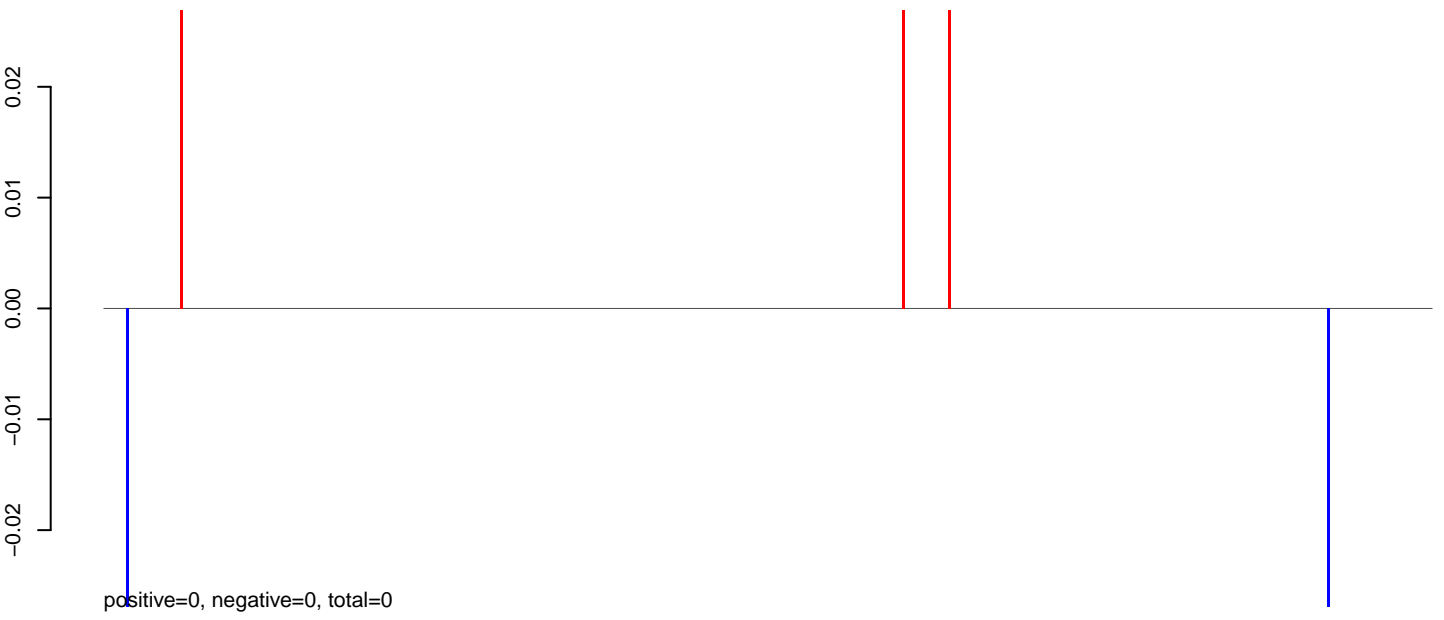
AeAeg_Aag2DENV2NGC.24_35.rep



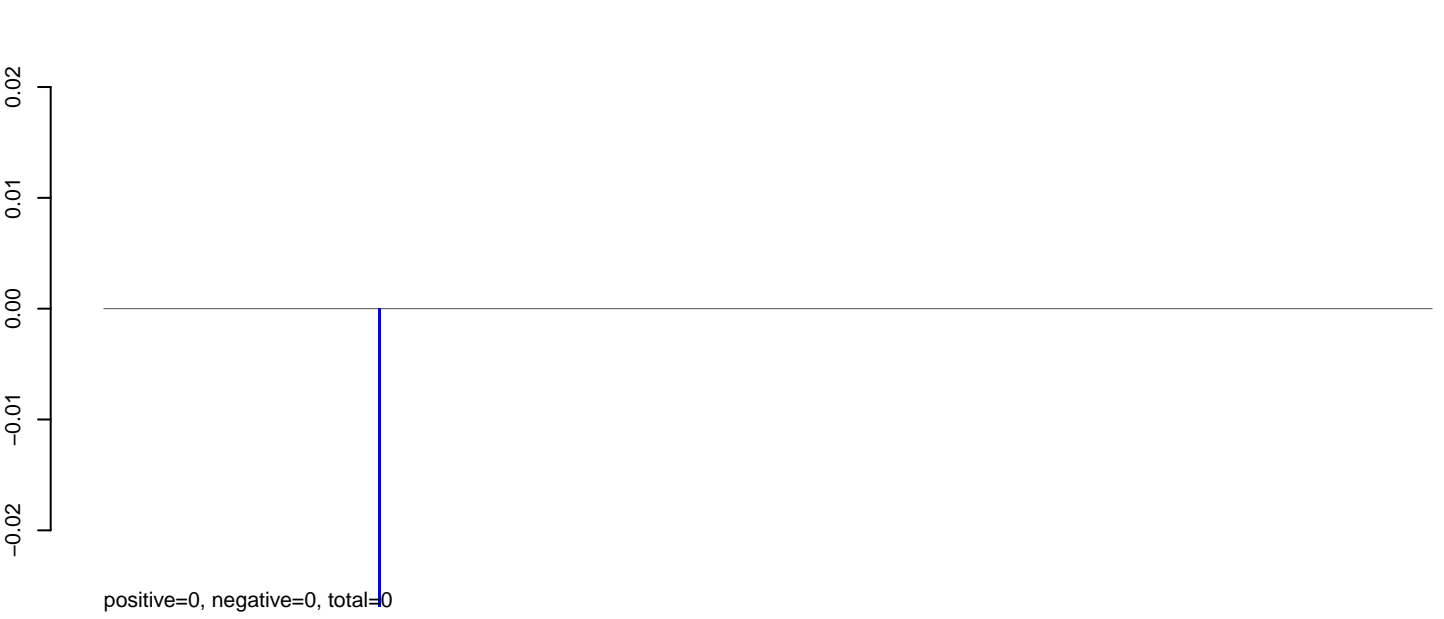
AeAeg_Aag2DENV2NGC.rep



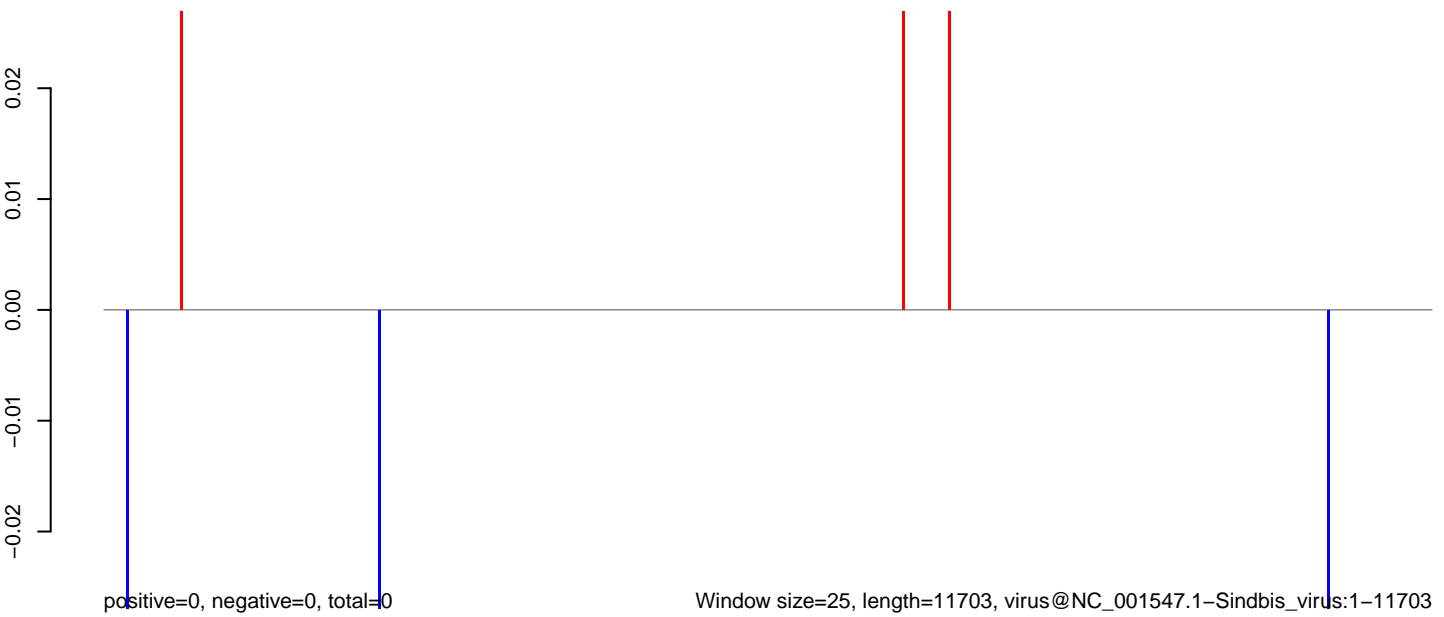
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



Window size=25, length=11703, virus@NC_001547.1-Sindbis_virus:1-11703

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2DENV2NGC.18_23.rep

positive=4, negative=1, total=5

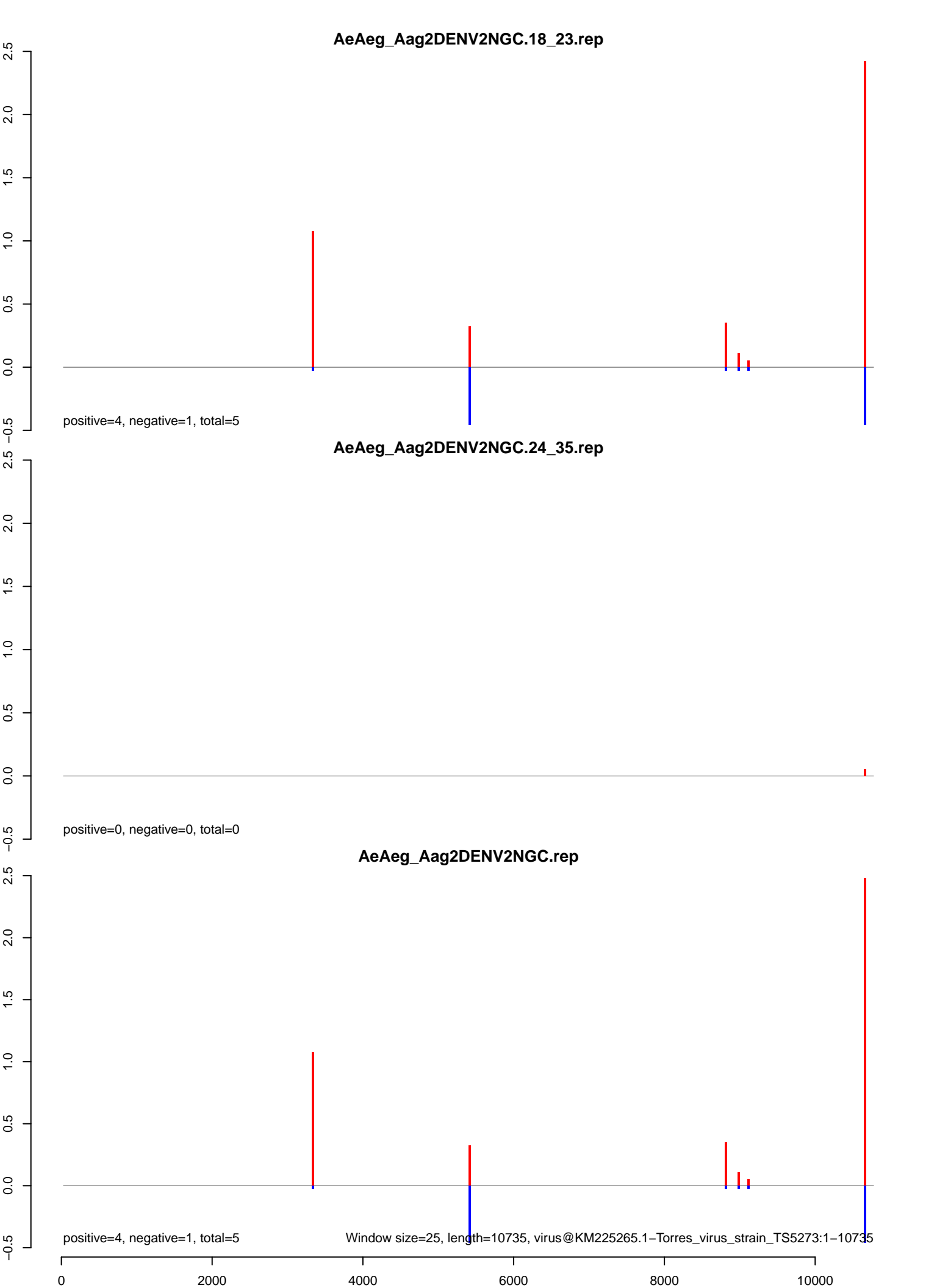
AeAeg_Aag2DENV2NGC.24_35.rep

positive=0, negative=0, total=0

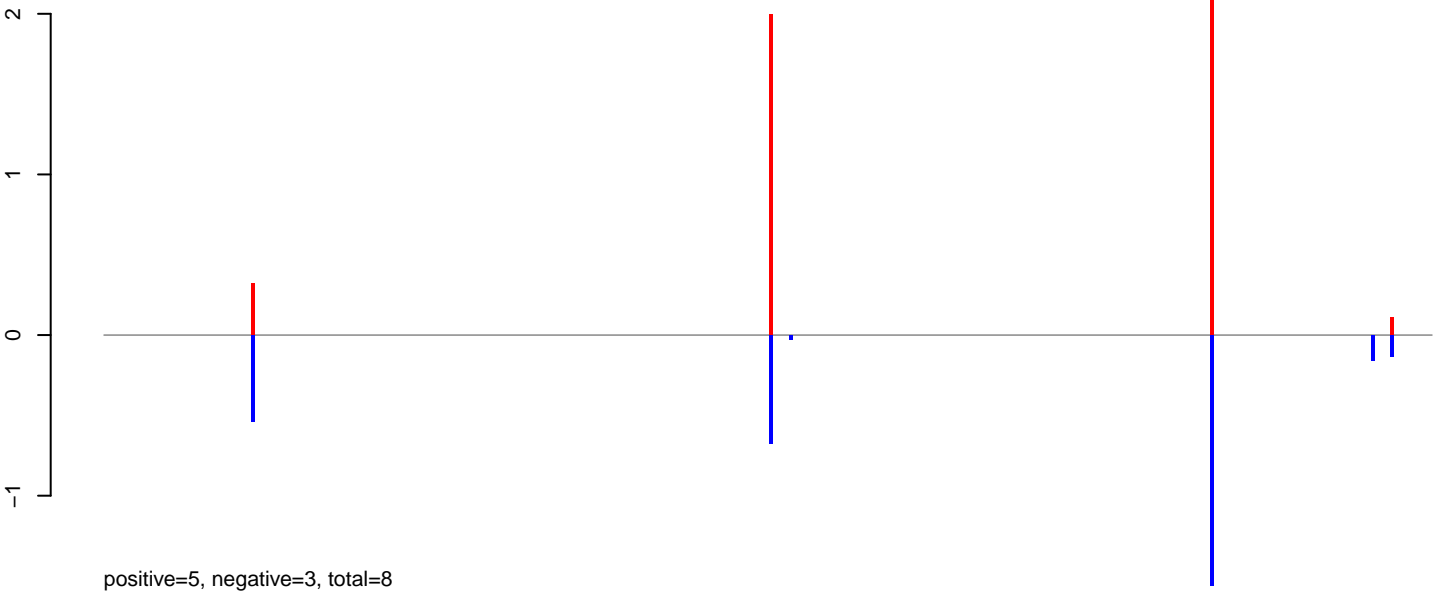
AeAeg_Aag2DENV2NGC.rep

positive=4, negative=1, total=5

Window size=25, length=10735, virus@KM225265.1-Torres_virus_strain_TS5273:1-10735



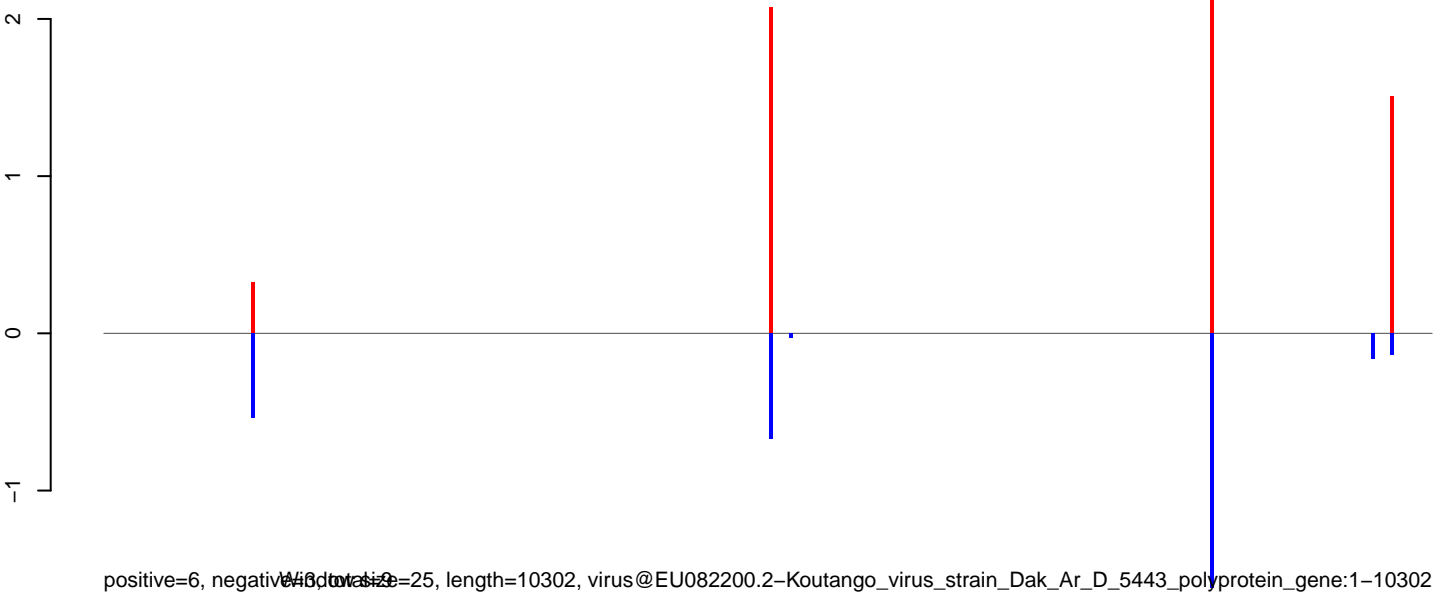
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



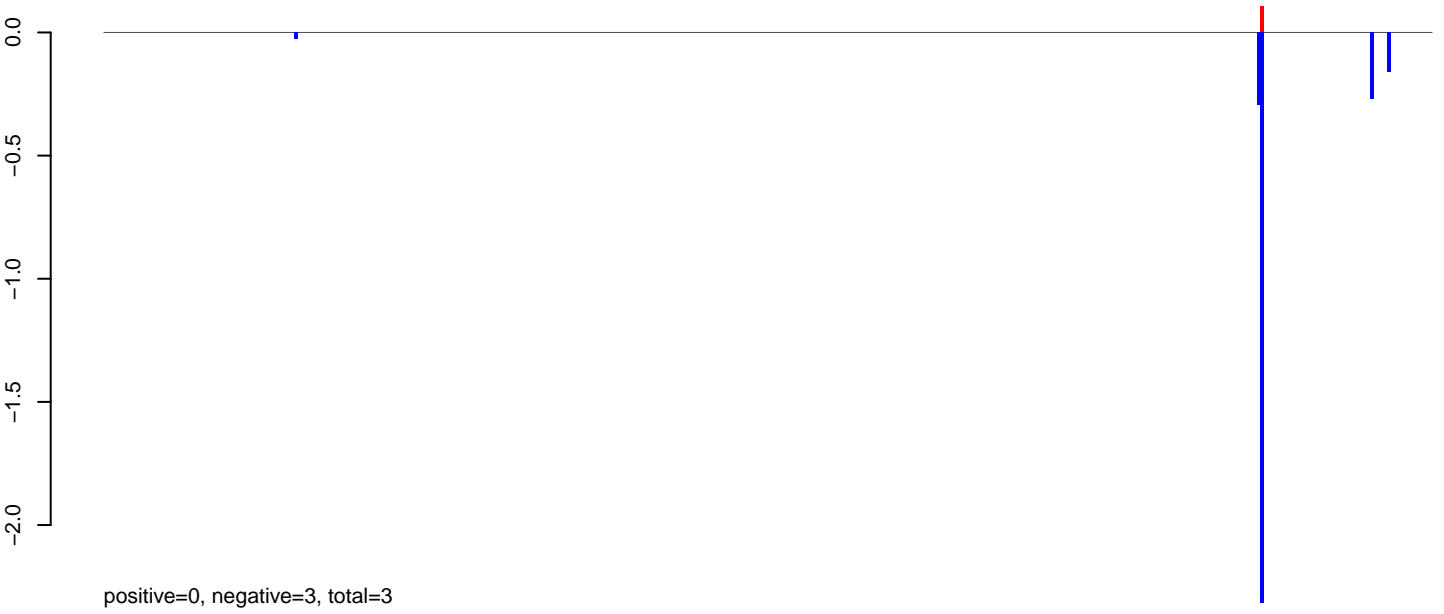
AeAeg_Aag2DENV2NGC.rep



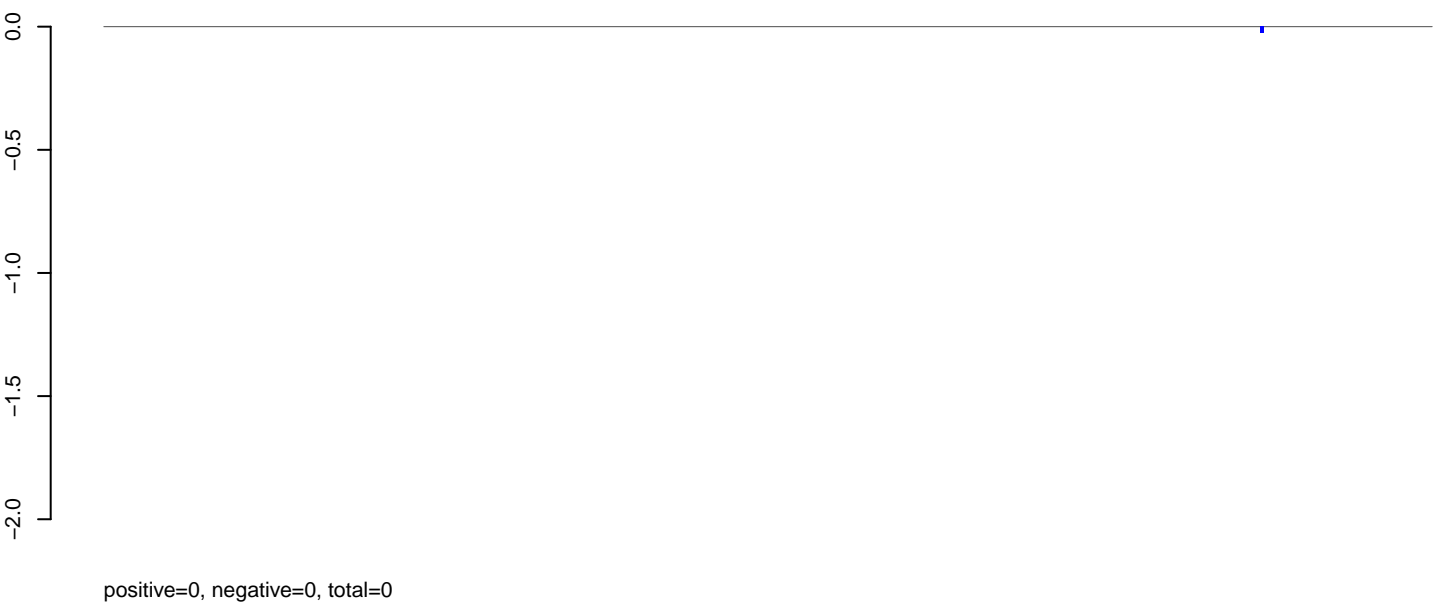
width=25, length=10302, virus@EU082200.2-Koutango_virus_strain_Dak_Ar_D_5443_polyprotein_gene:1-10302

0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



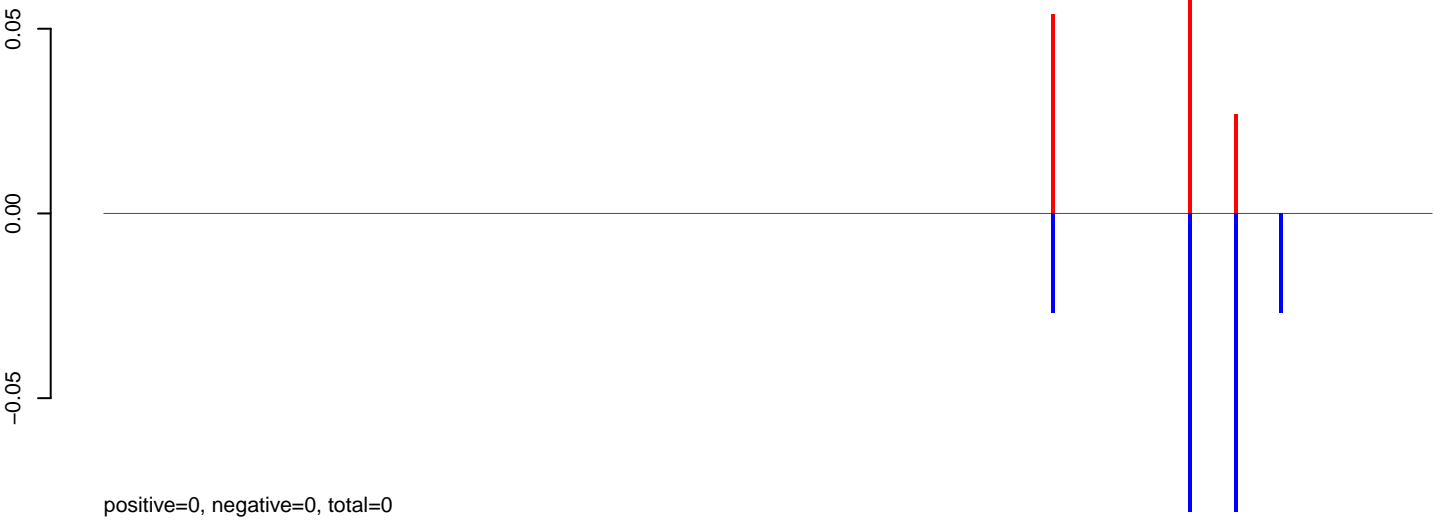
AeAeg_Aag2DENV2NGC.rep



positive=0, negative=3, total=3, size=25, length=10251, virus@NC_033723.1-Gadgets_Gully_virus_from_Australia_polyprotein_gene:1-10251

0 2000 4000 6000 8000 10000

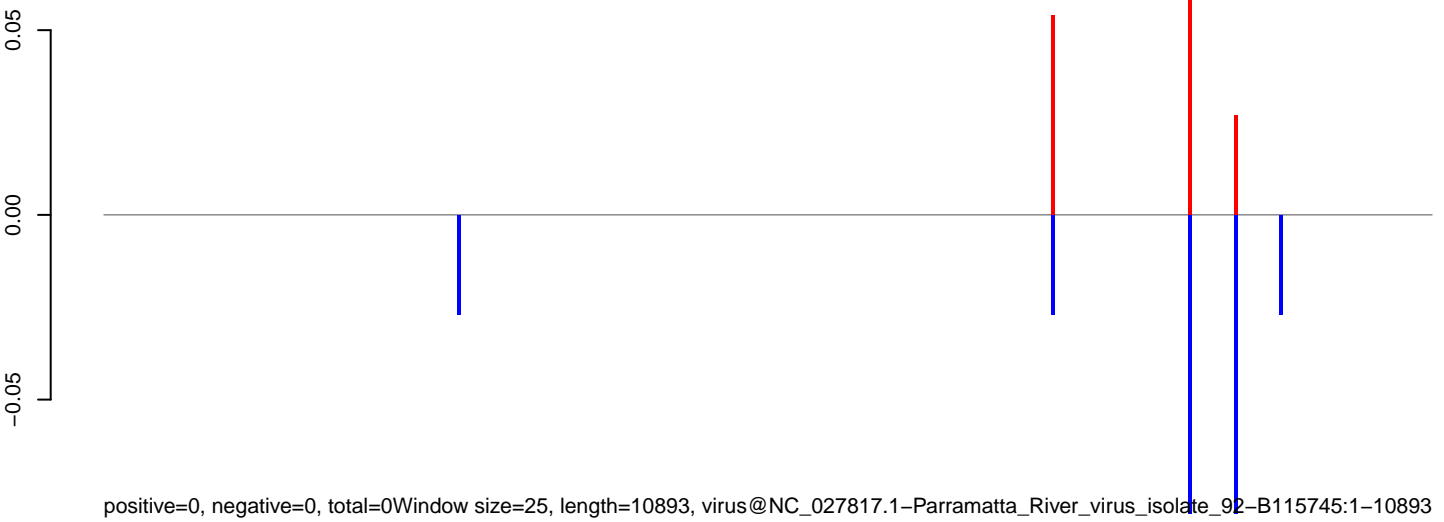
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



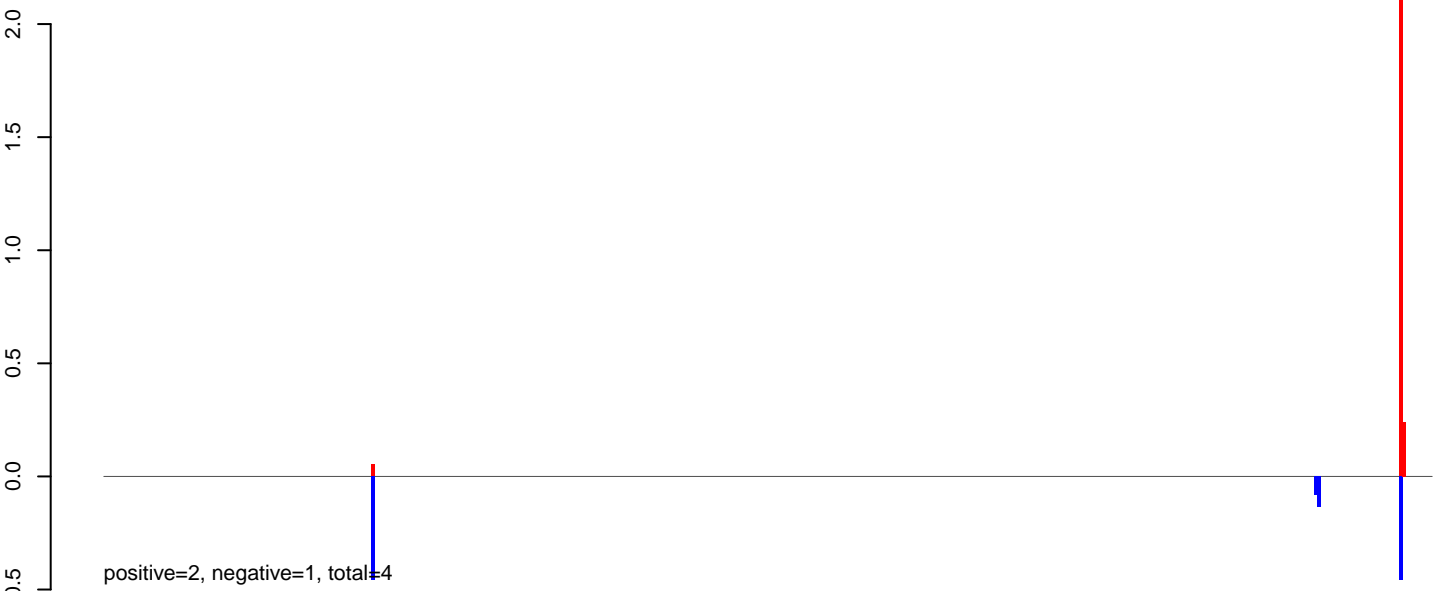
AeAeg_Aag2DENV2NGC.rep



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

0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

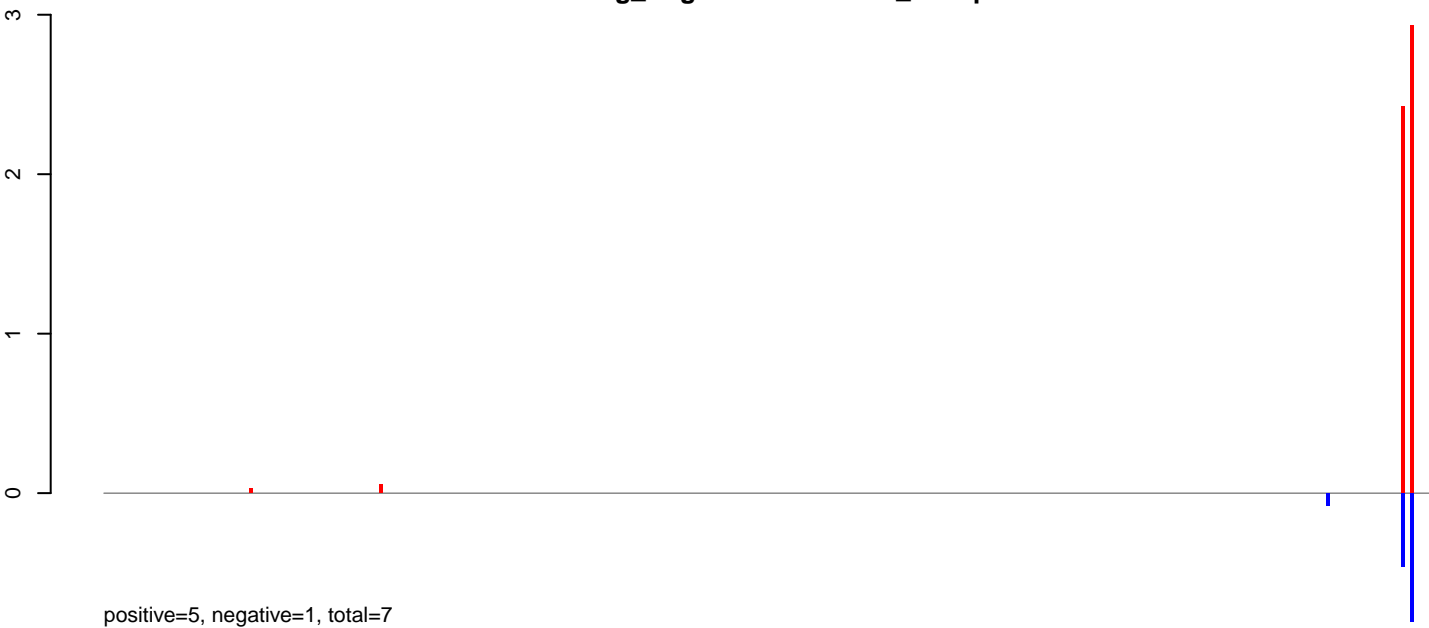


AeAeg_Aag2DENV2NGC.rep

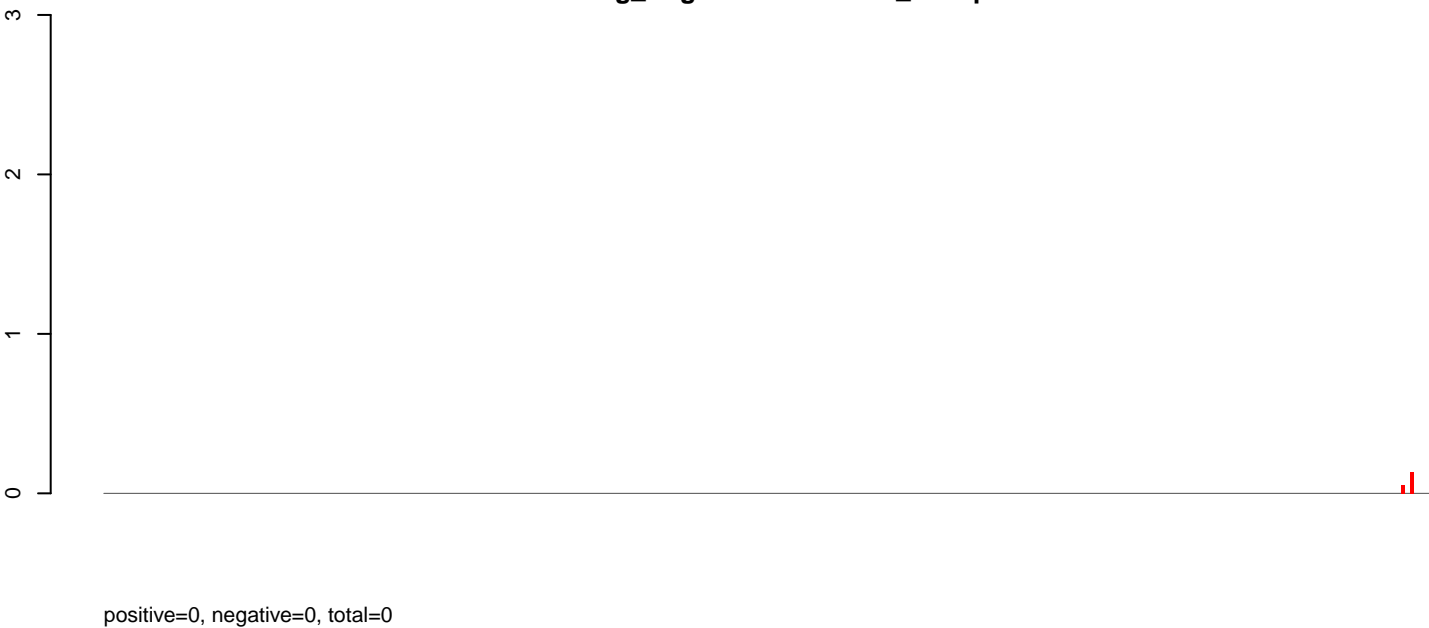


0 2000 4000 6000 8000 10000

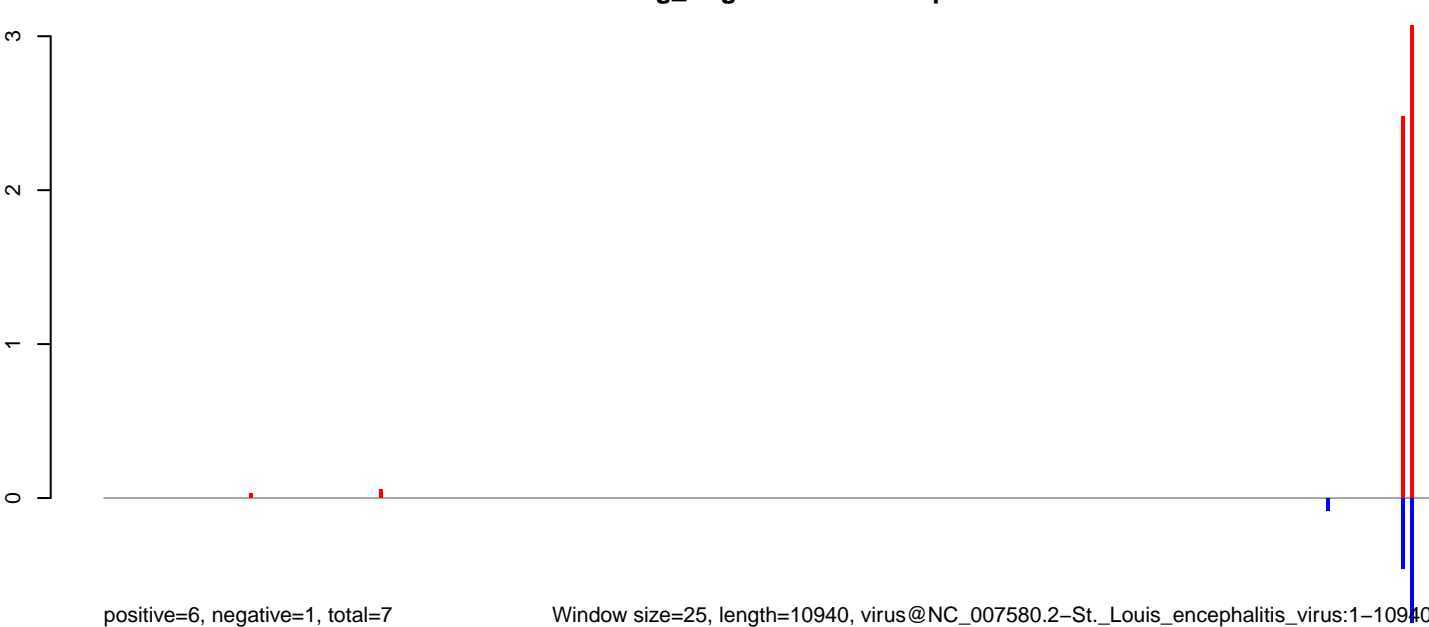
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



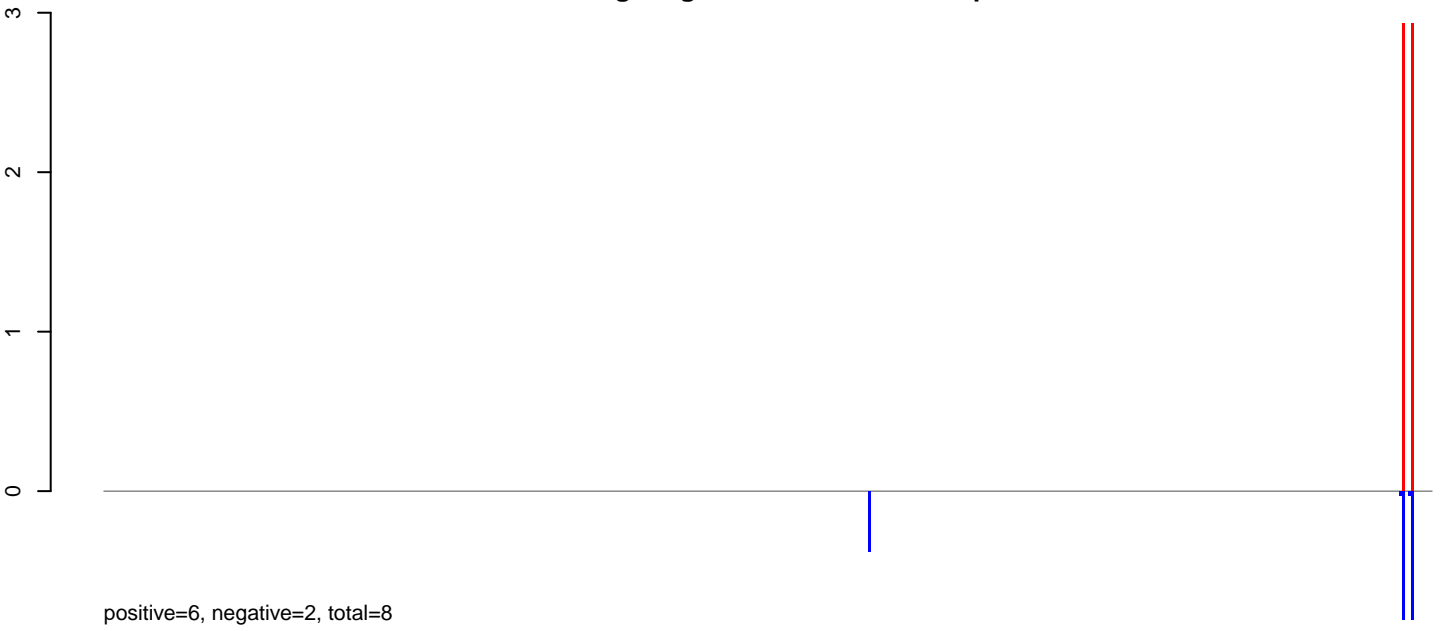
AeAeg_Aag2DENV2NGC.rep



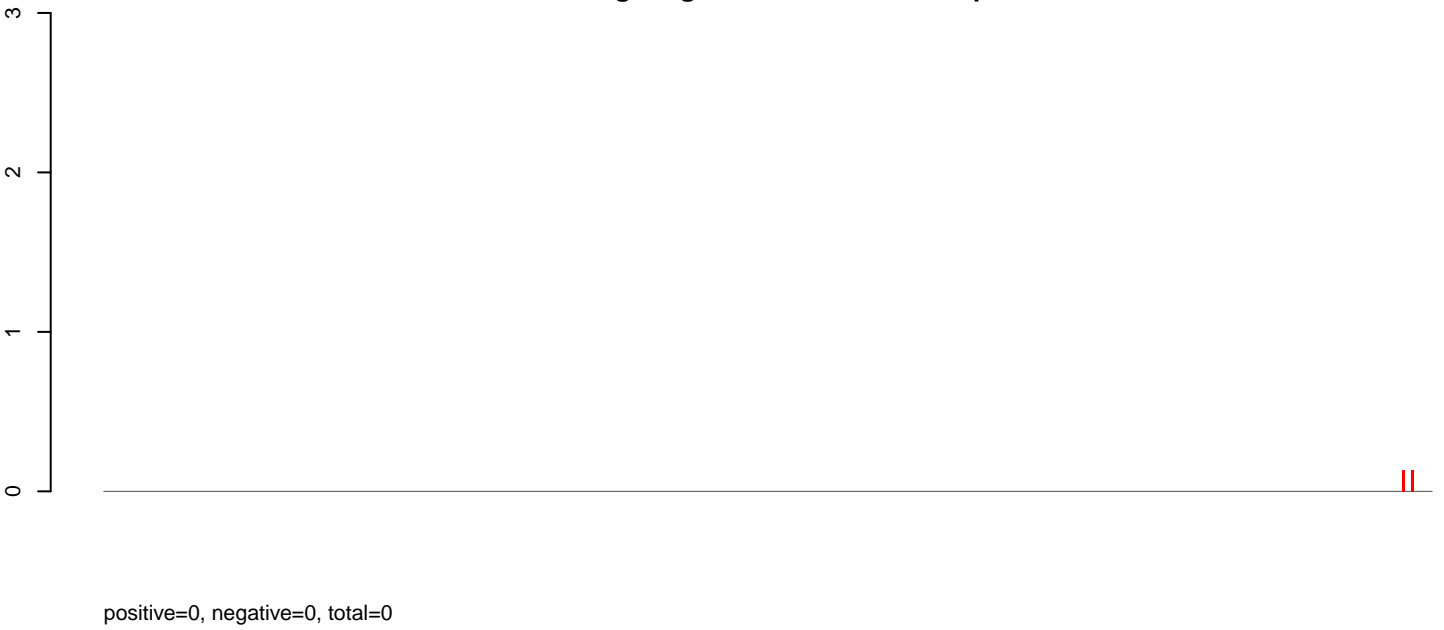
Window size=25, length=10940, virus@NC_007580.2-St._Louis_encephalitis_virus:1-10940

0 2000 4000 6000 8000 10000

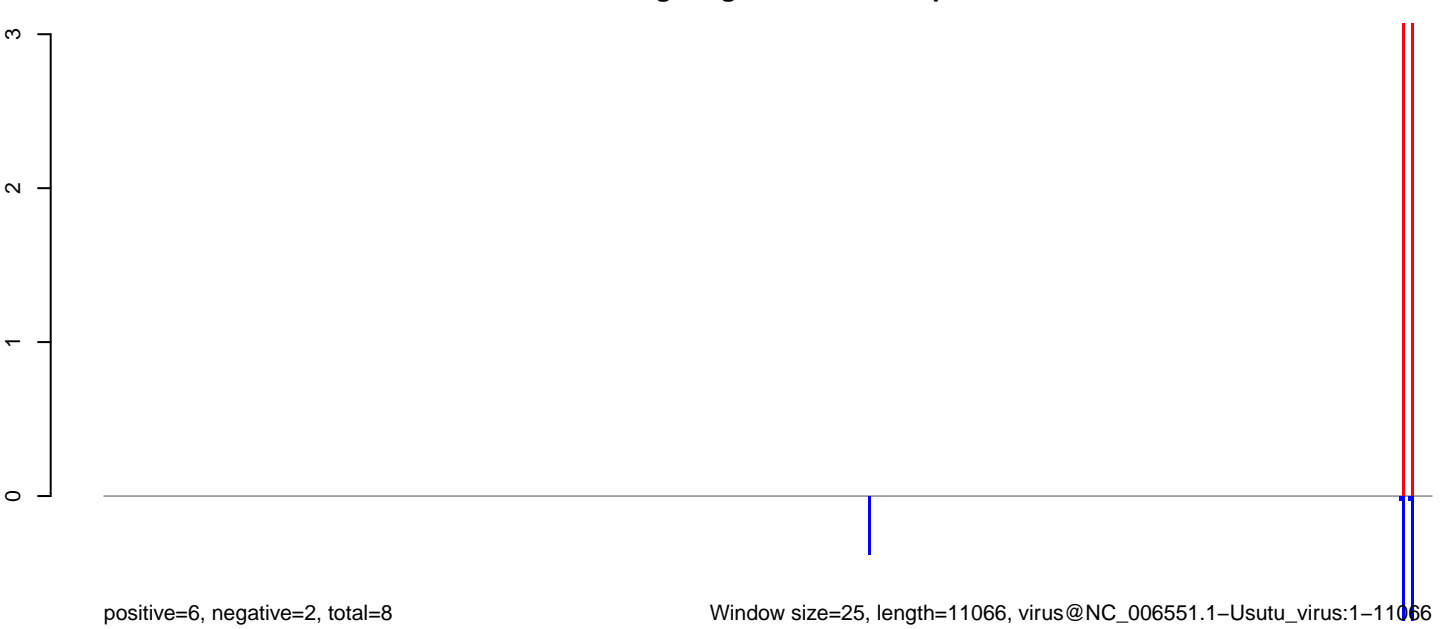
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



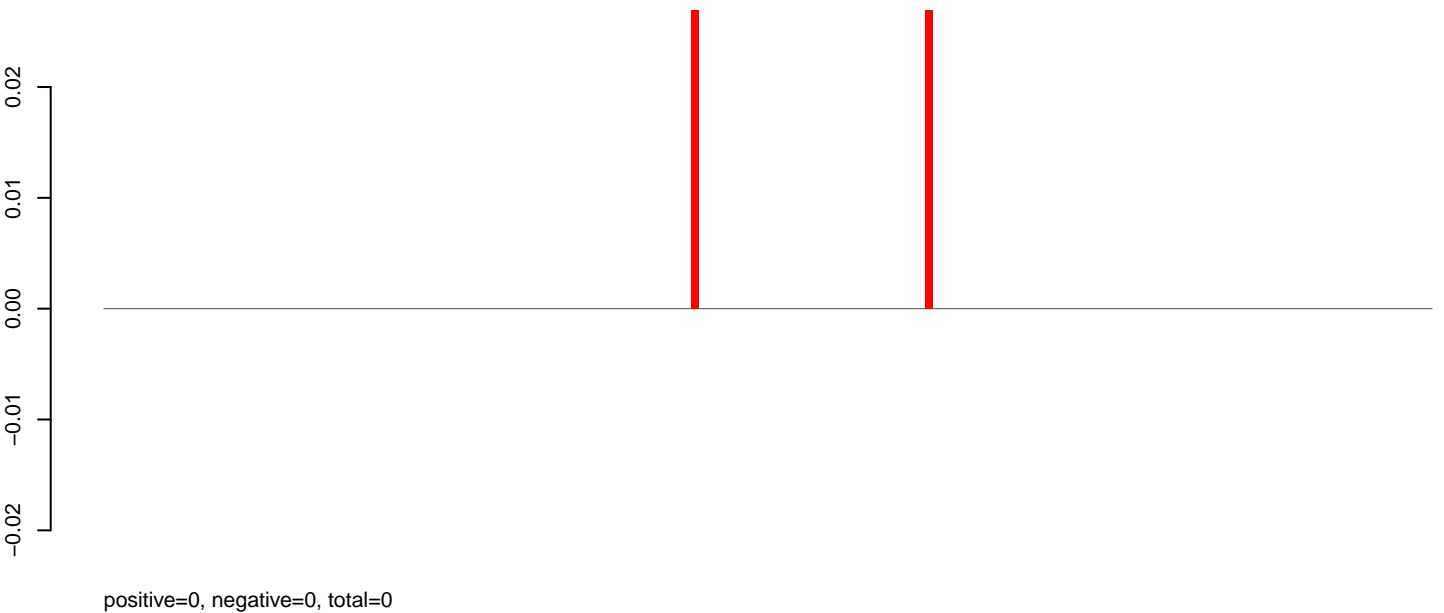
Window size=25, length=11066, virus@NC_006551.1-Uсутu_virus:1-11066

0 2000 4000 6000 8000 10000

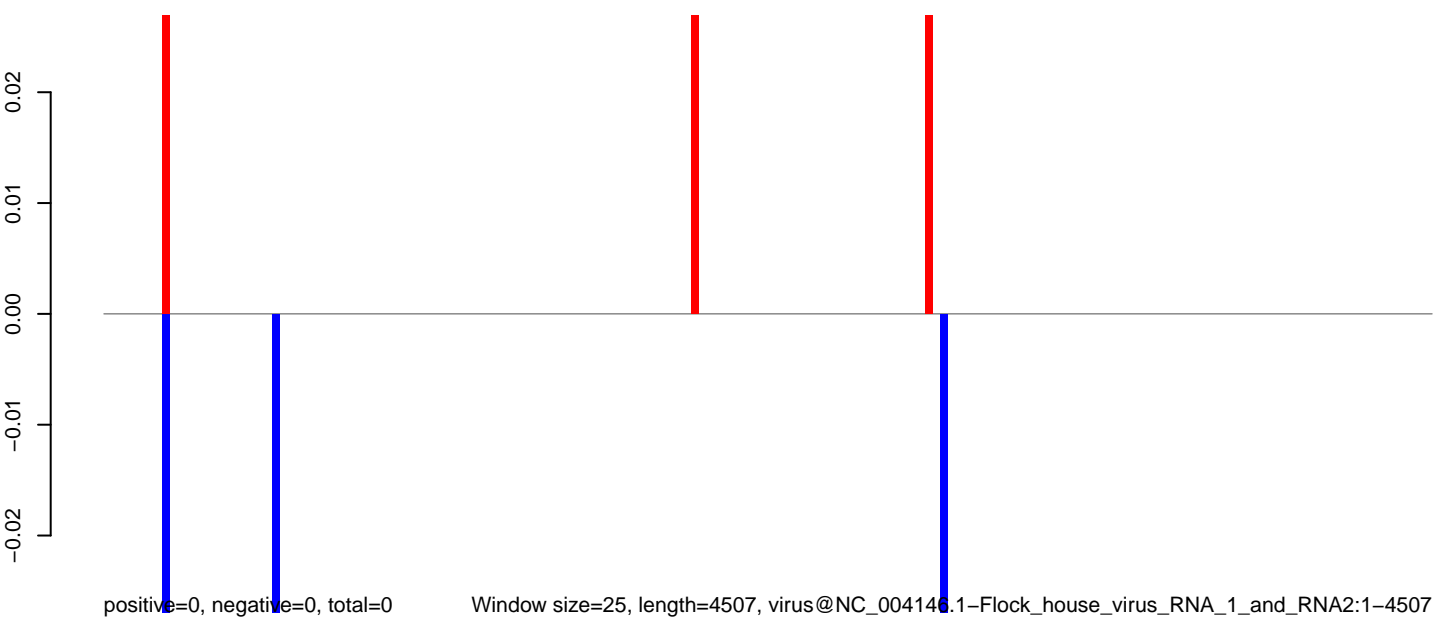
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



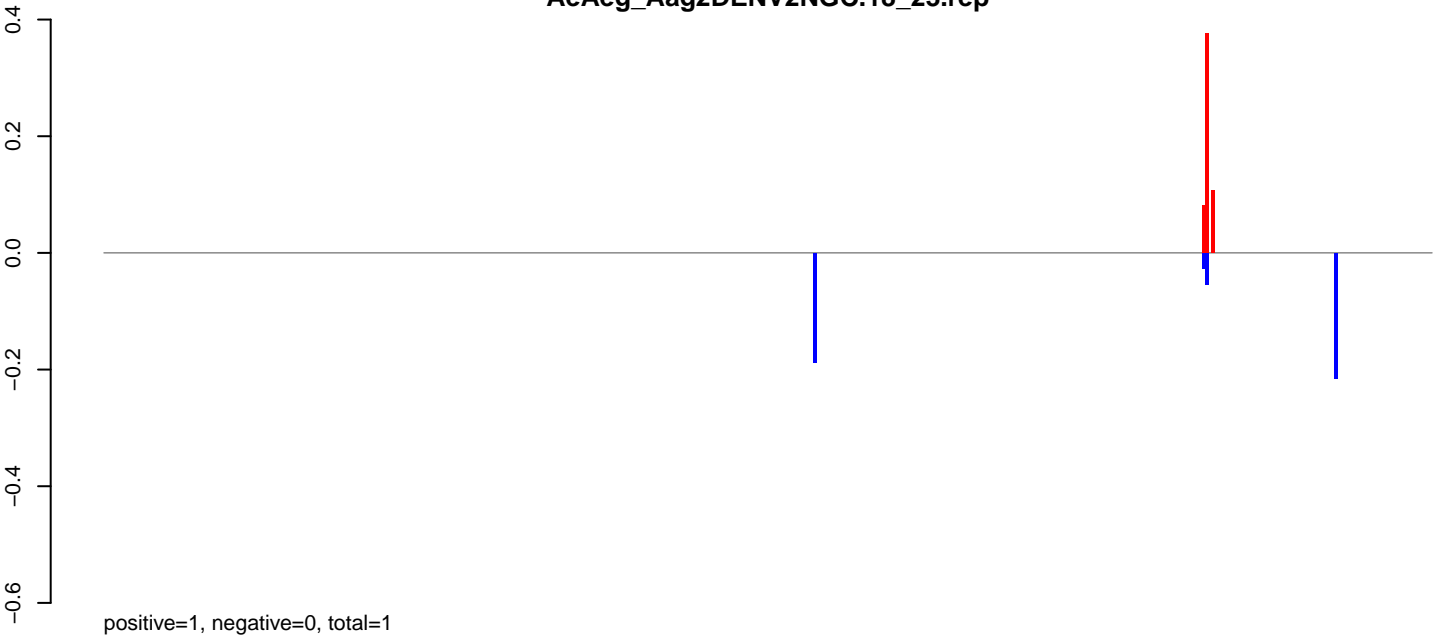
AeAeg_Aag2DENV2NGC.rep



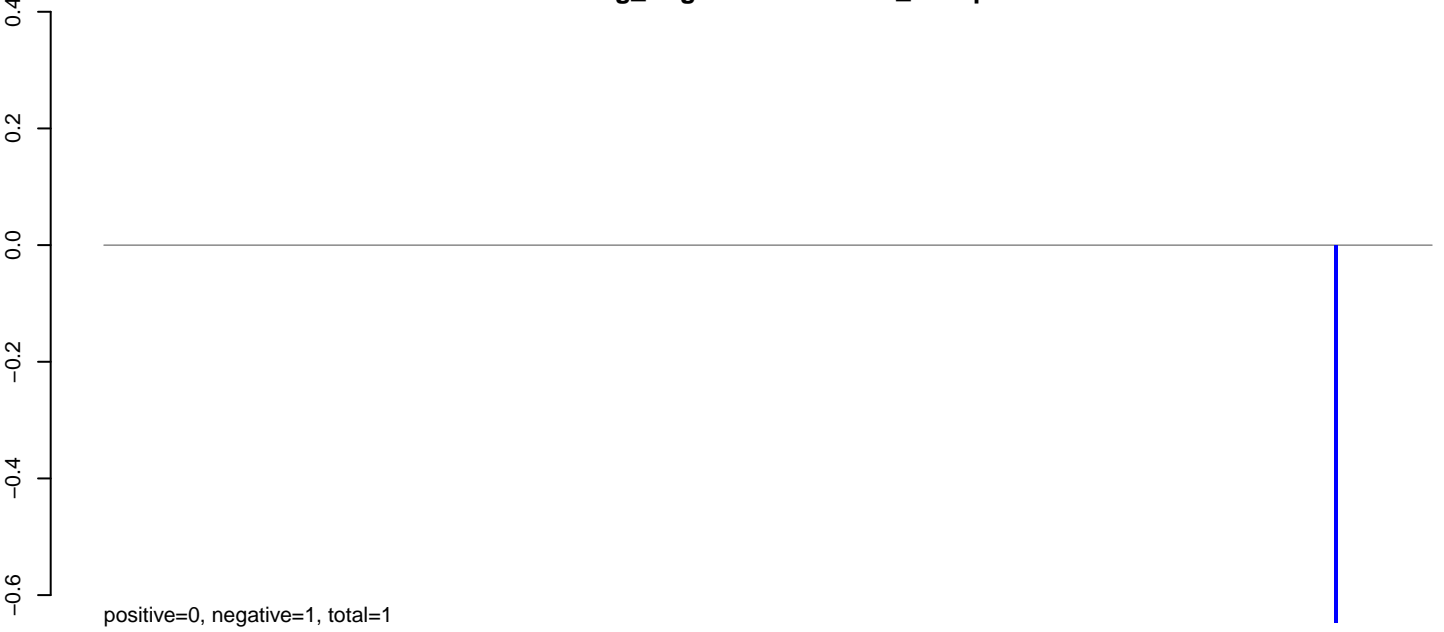
Window size=25, length=4507, virus@NC_004146.1-Flock_house_virus_RNA_1_and_RNA2:1-4507

0 1000 2000 3000 4000

AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



Window size=25, length=10839, virus@NC_003687.1-Powassan_virus:1-10839

0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep

positive=5, negative=5, total=9

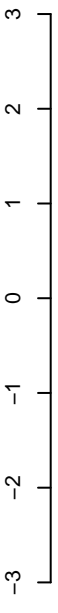
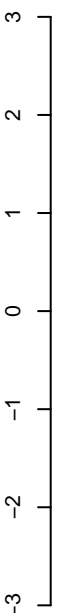
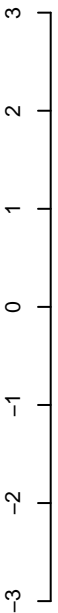
AeAeg_Aag2DENV2NGC.24_35.rep

positive=0, negative=0, total=0

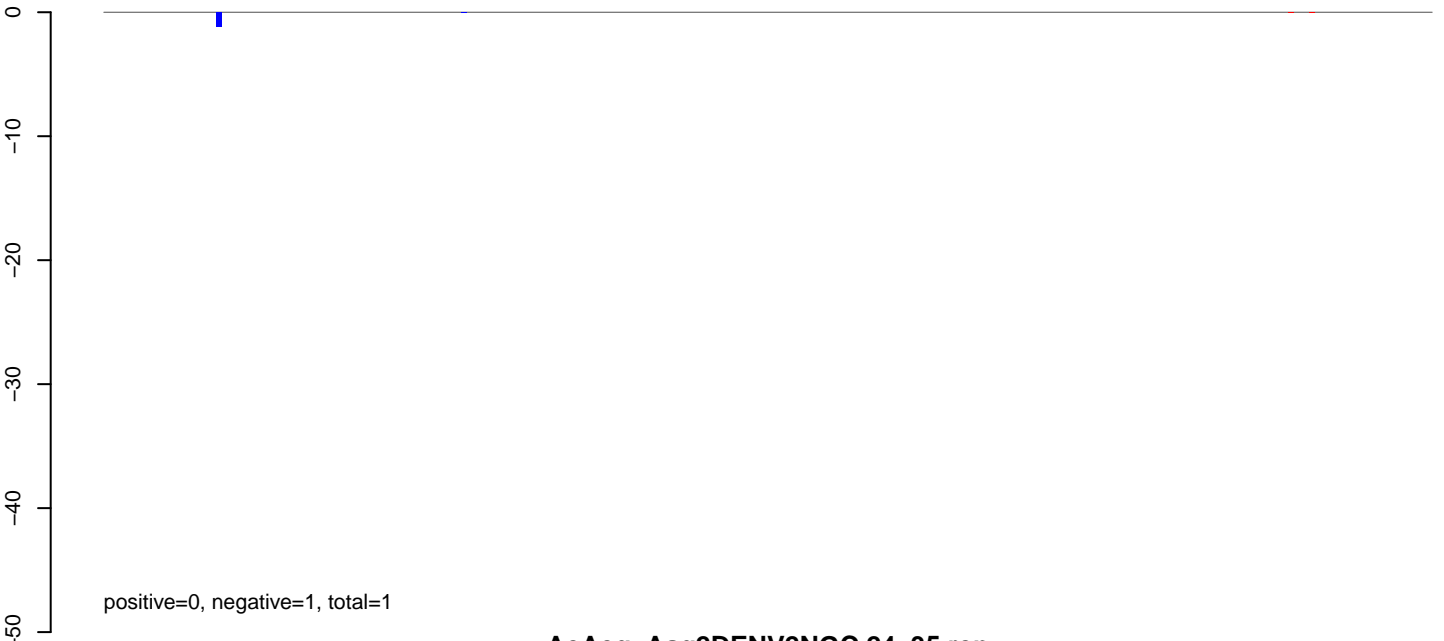
AeAeg_Aag2DENV2NGC.rep

positive=5, negative=5, total=9

Window size=25, length=10794, virus@AY632542.4-Rocio_virus_strain_SPH_34675:1-10794



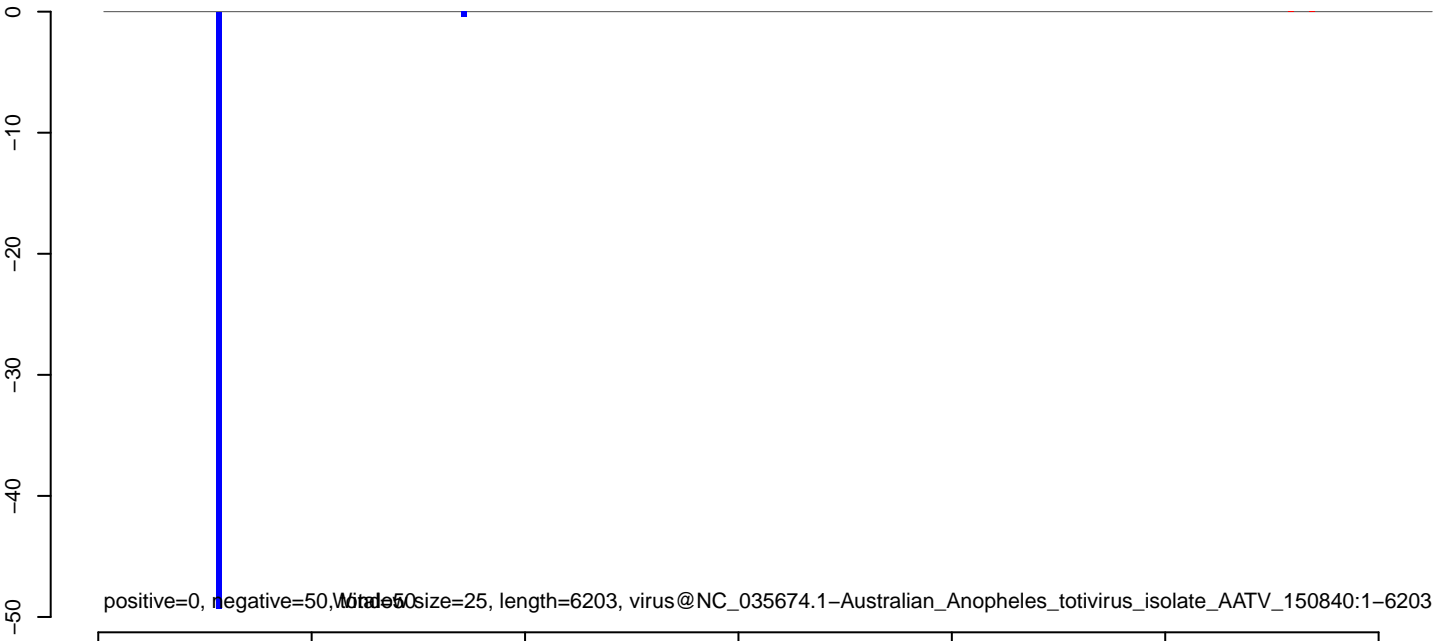
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



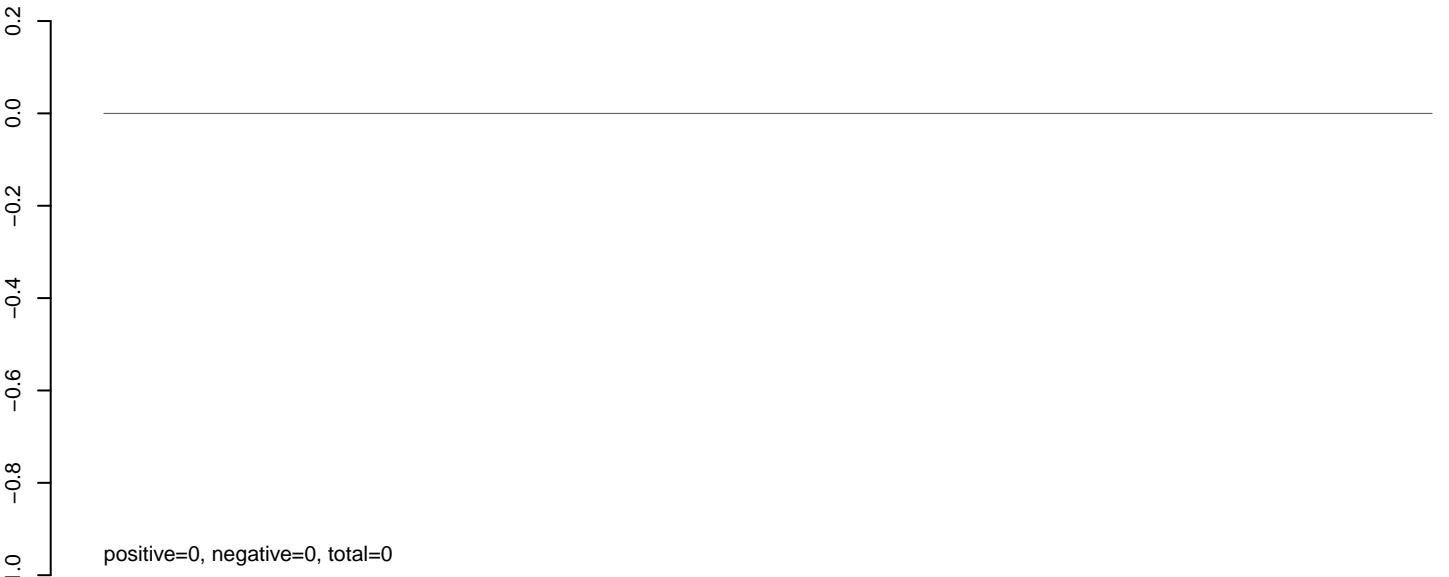
AeAeg_Aag2DENV2NGC.rep



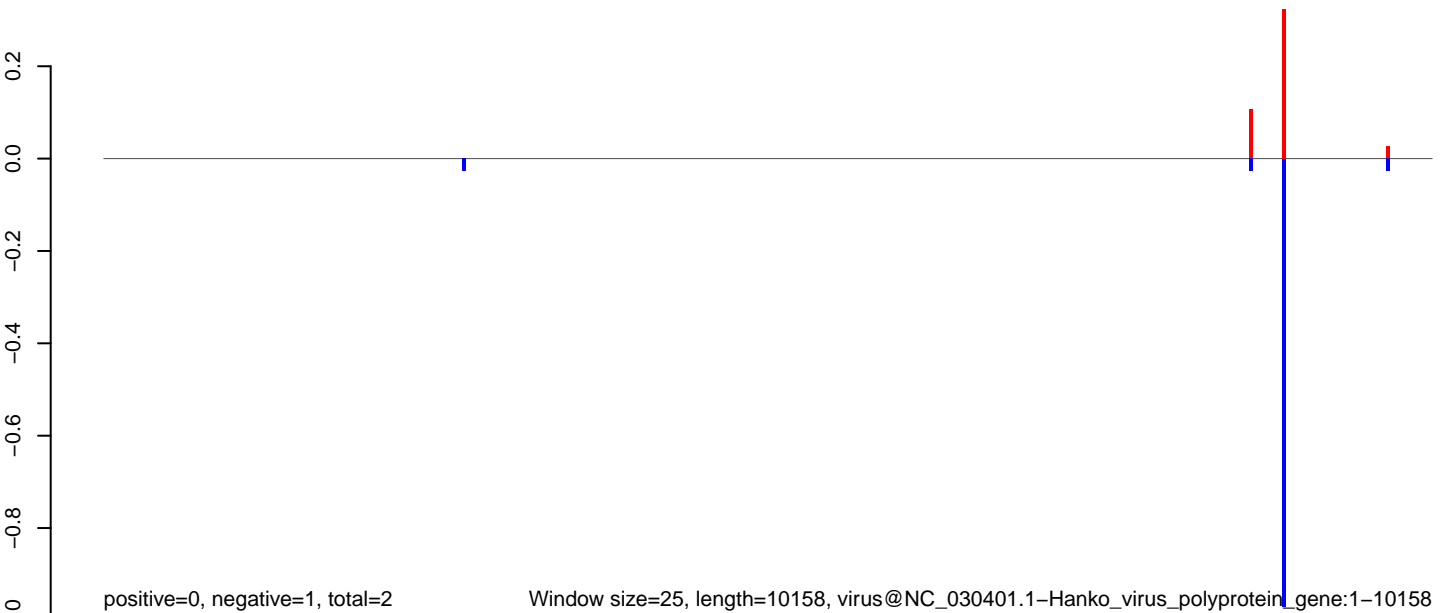
AeAeg_Aag2DENV2NGC.18_23.rep



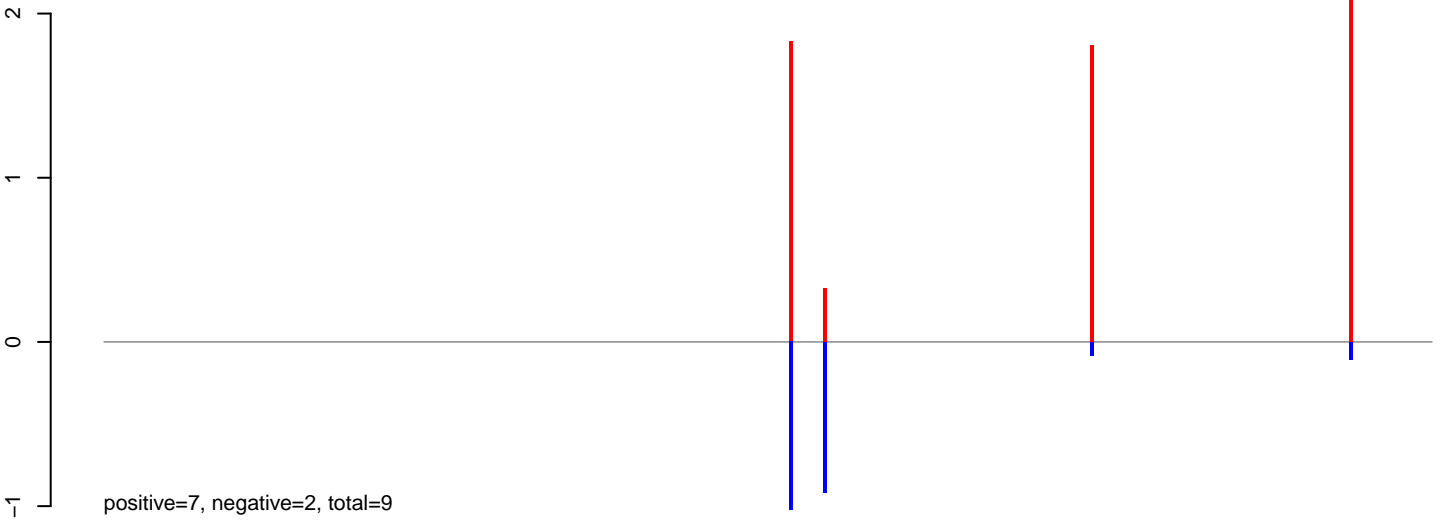
AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



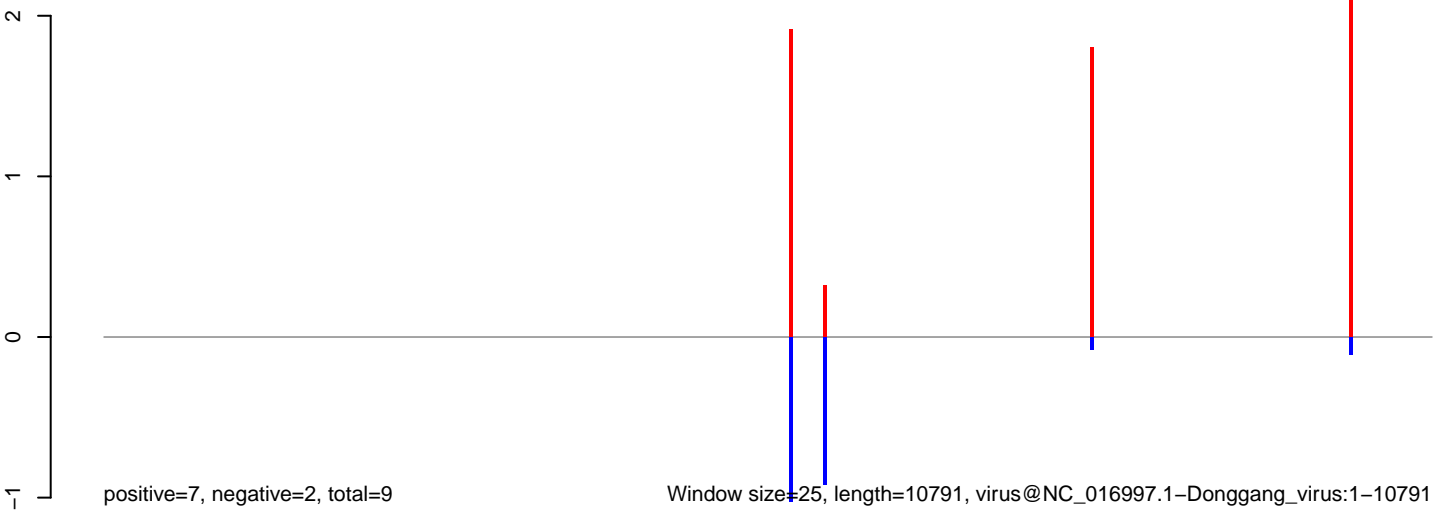
AeAeg_Aag2DENV2NGC.18_23.rep



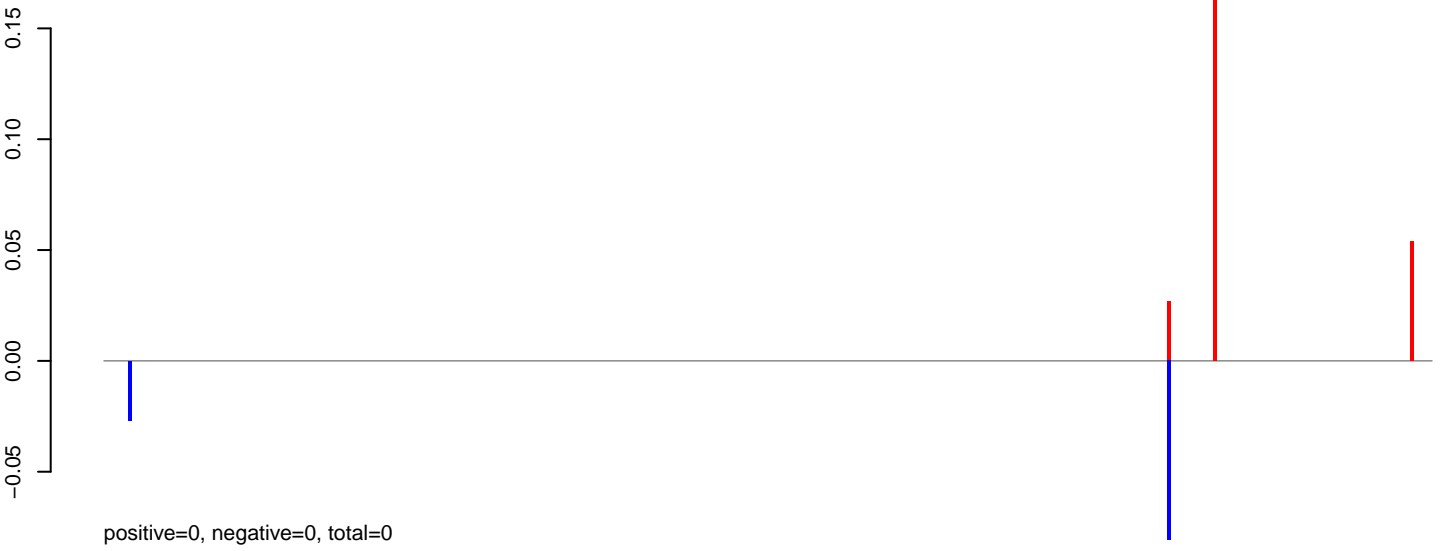
AeAeg_Aag2DENV2NGC.24_35.rep



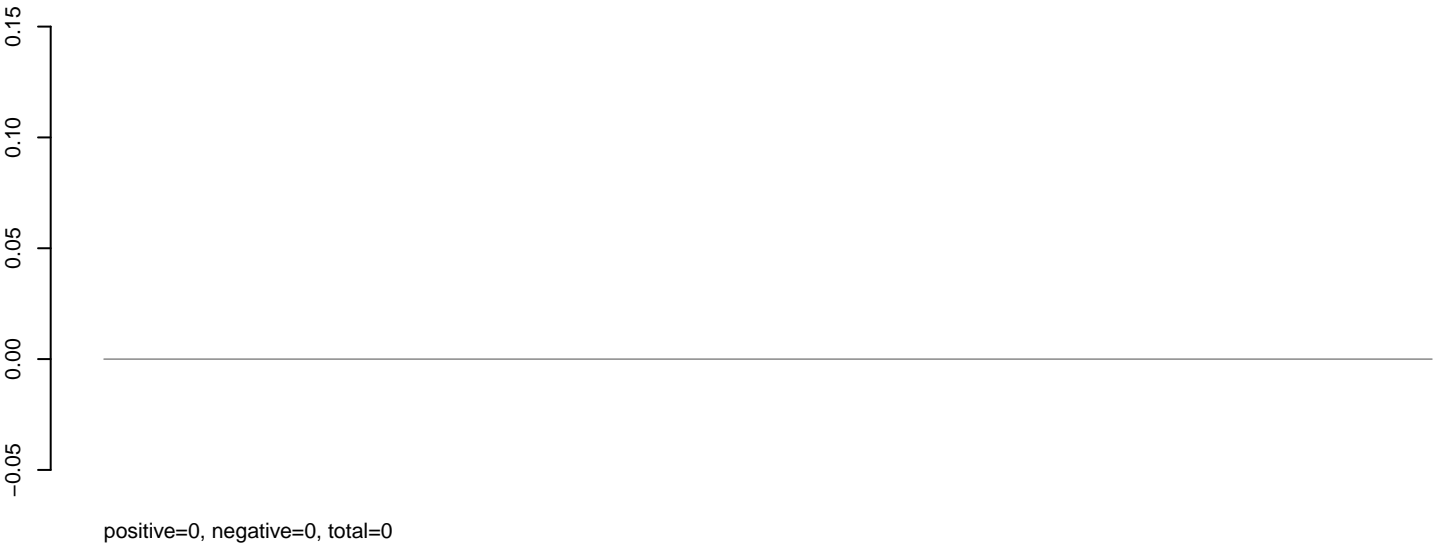
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



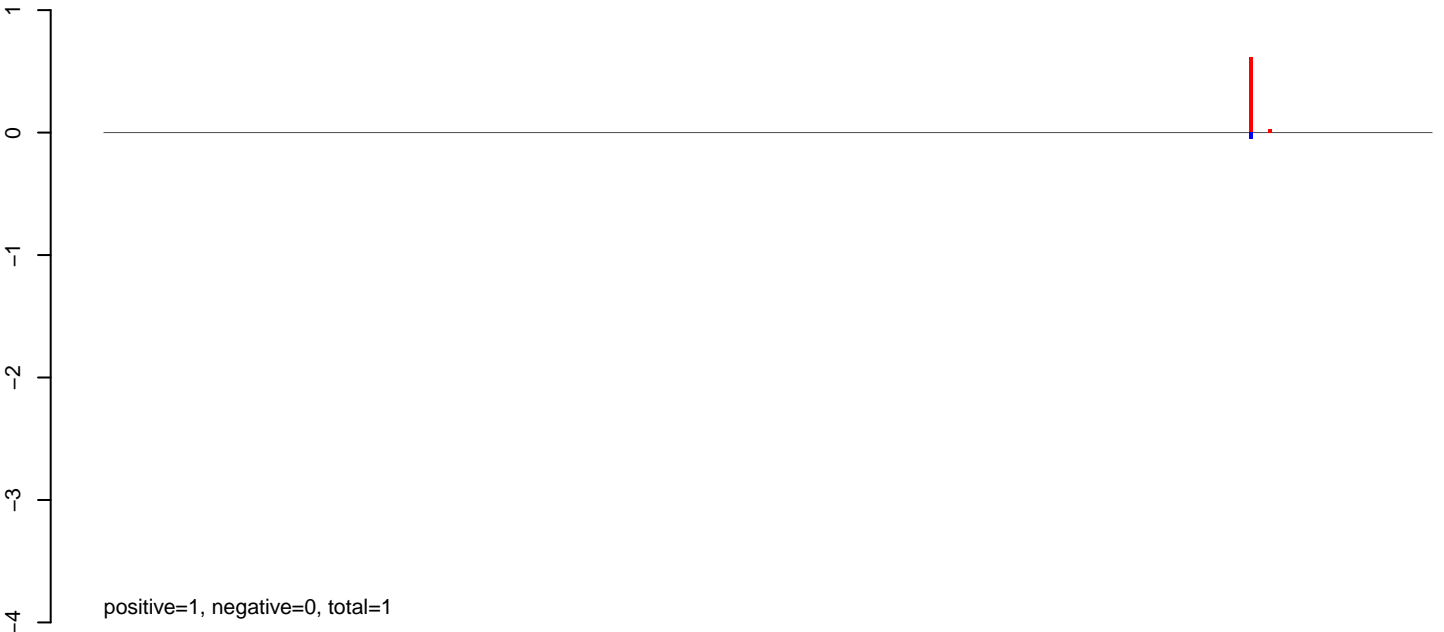
Window size=25, length=10793, virus@NC_008719.1-Sepik_virus:1-10793

0 2000 4000 6000 8000 10000

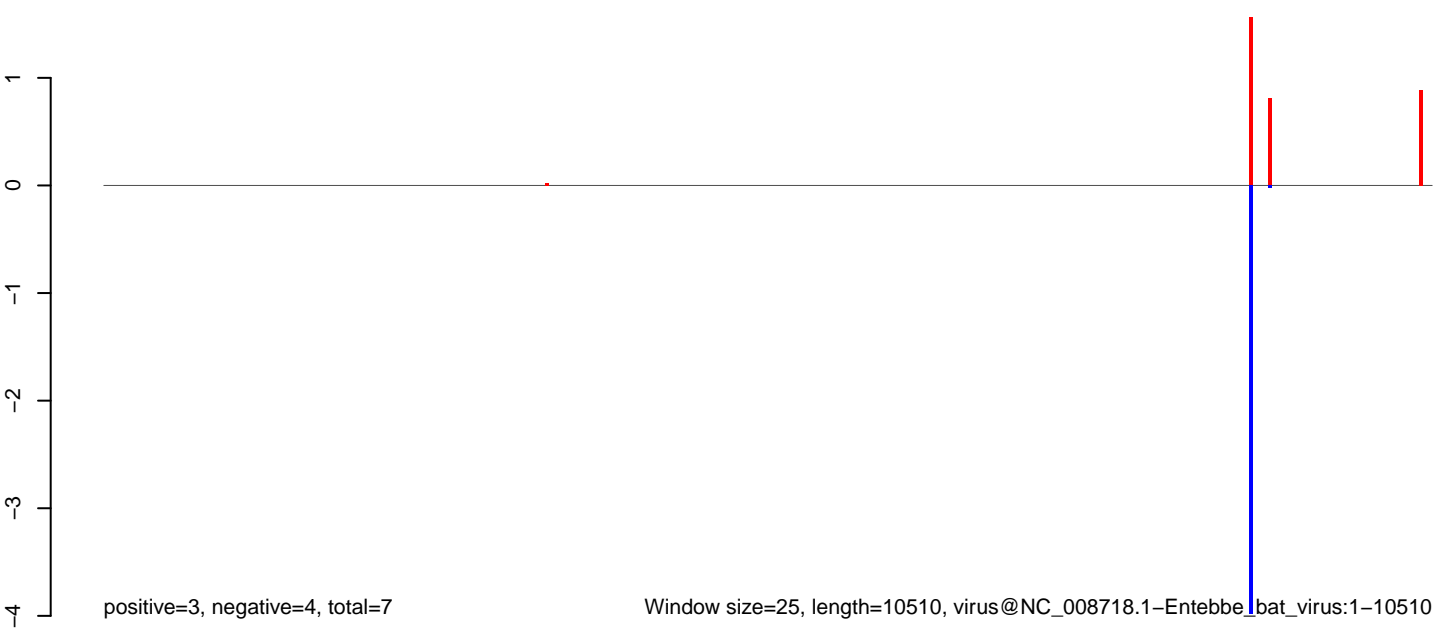
AeAeg_Aag2DENV2NGC.18_23.rep



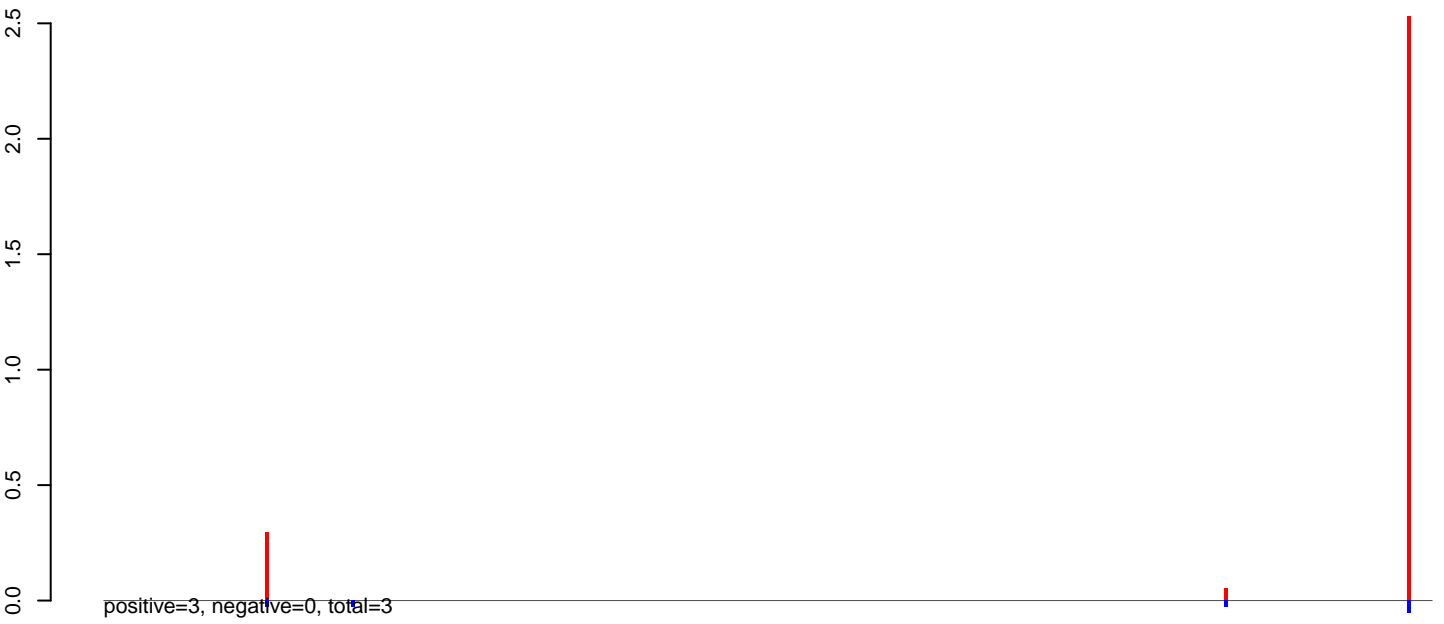
AeAeg_Aag2DENV2NGC.24_35.rep



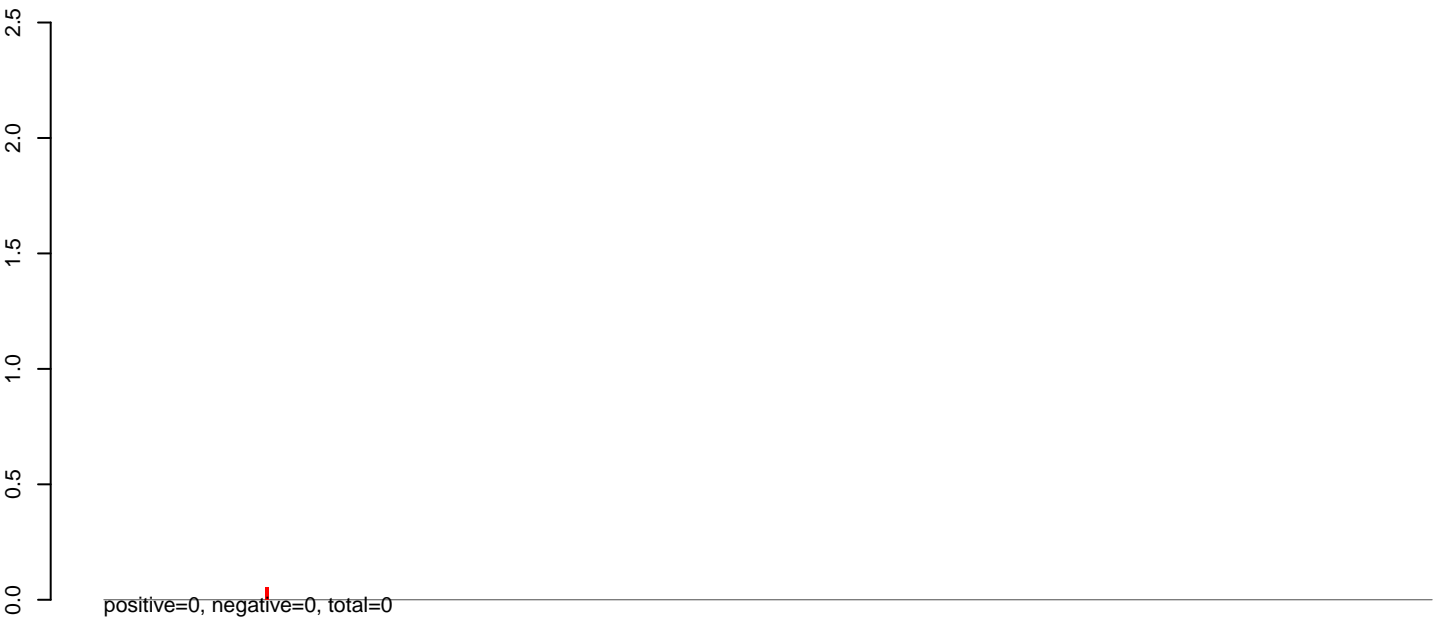
AeAeg_Aag2DENV2NGC.rep



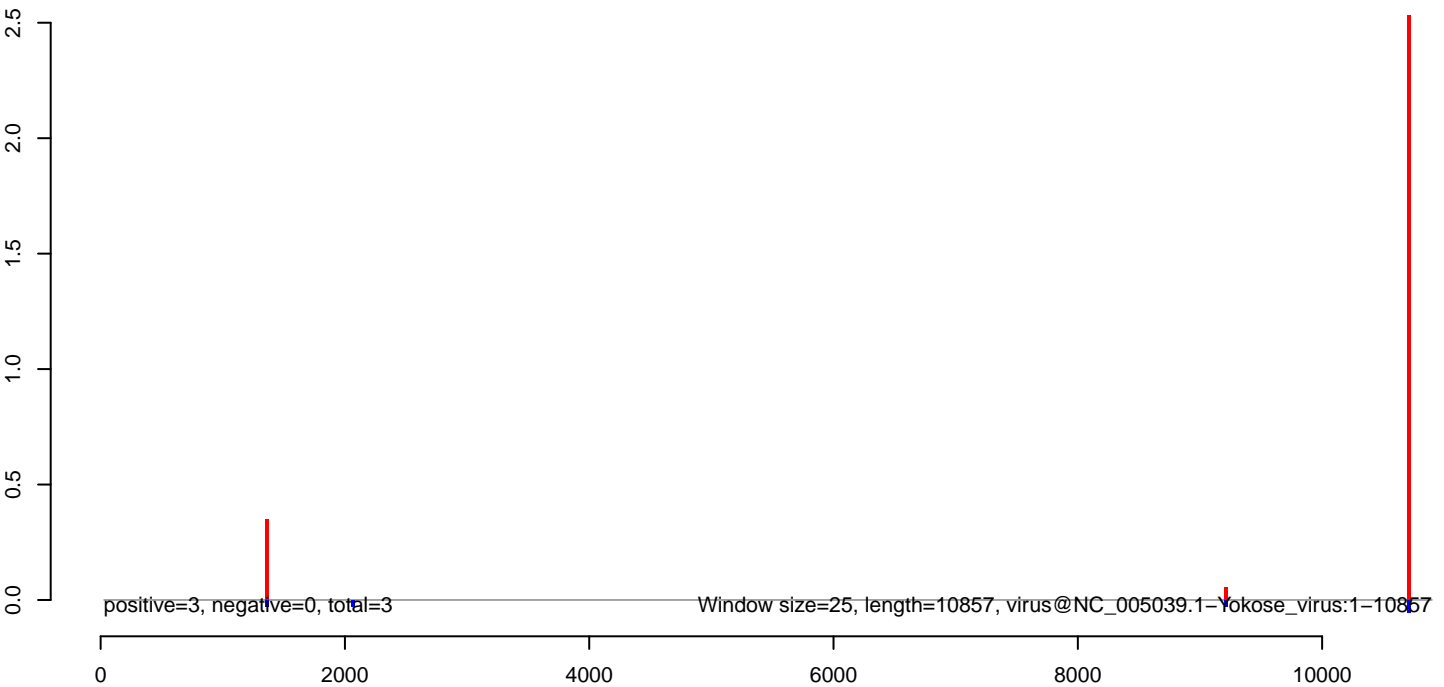
AeAeg_Aag2DENV2NGC.18_23.rep



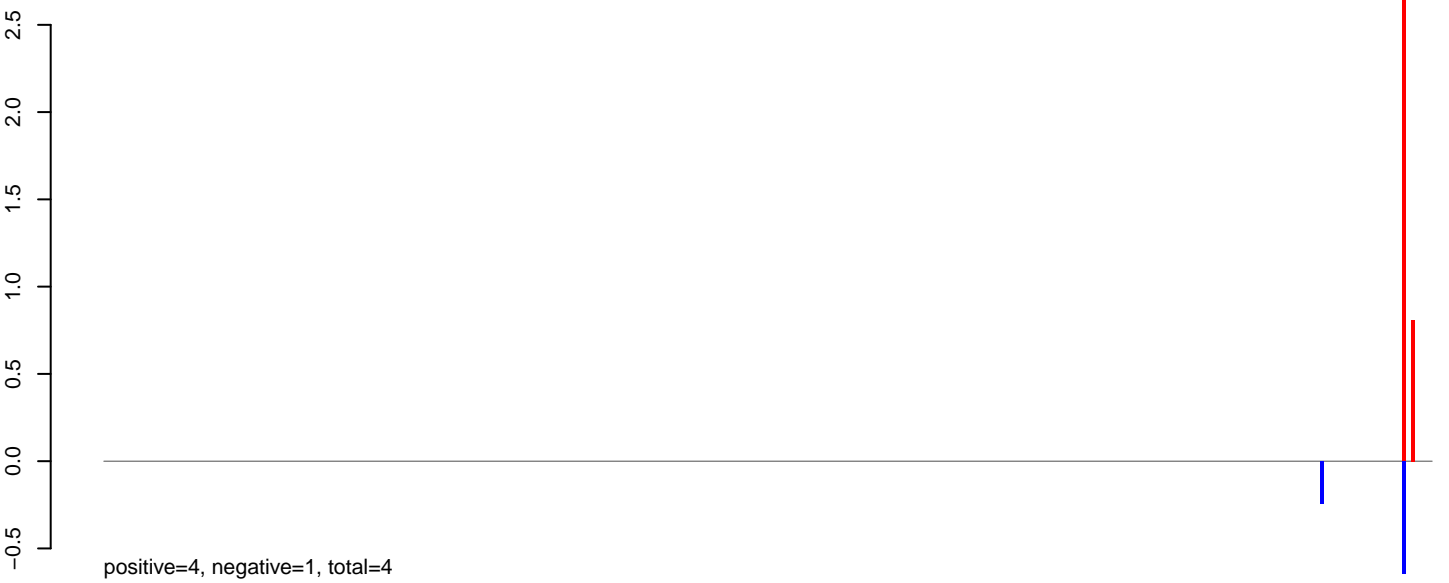
AeAeg_Aag2DENV2NGC.24_35.rep



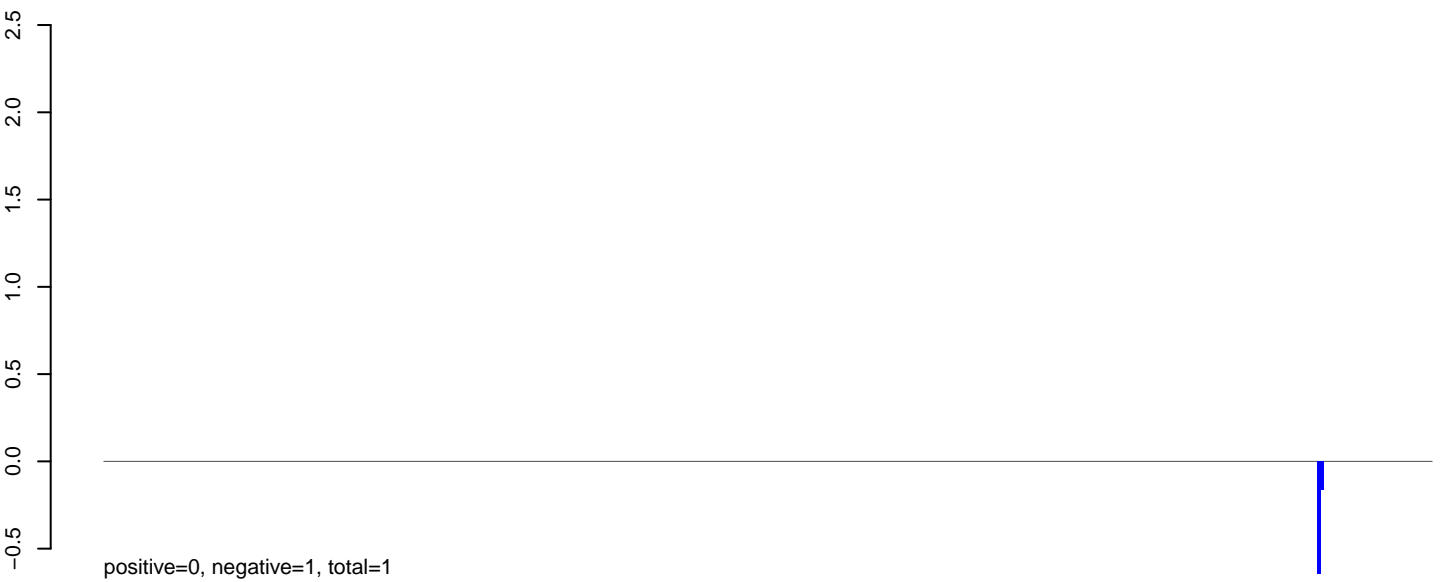
AeAeg_Aag2DENV2NGC.rep



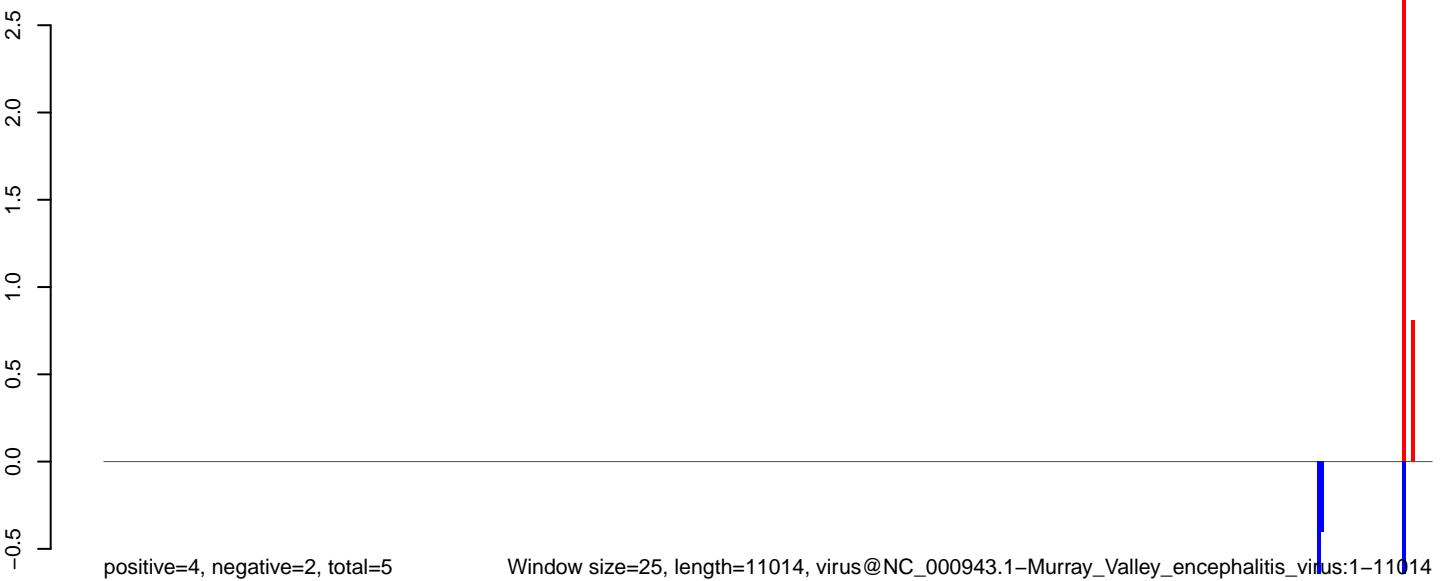
AeAeg_Aag2DENV2NGC.18_23.rep



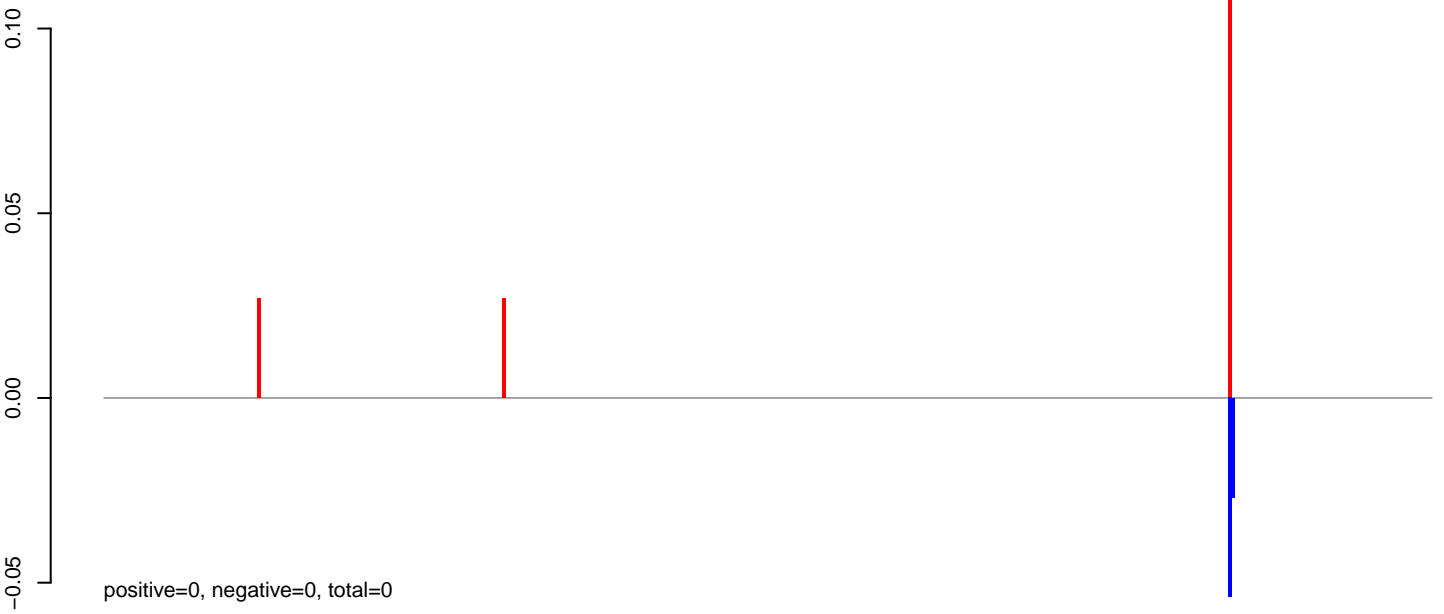
AeAeg_Aag2DENV2NGC.24_35.rep



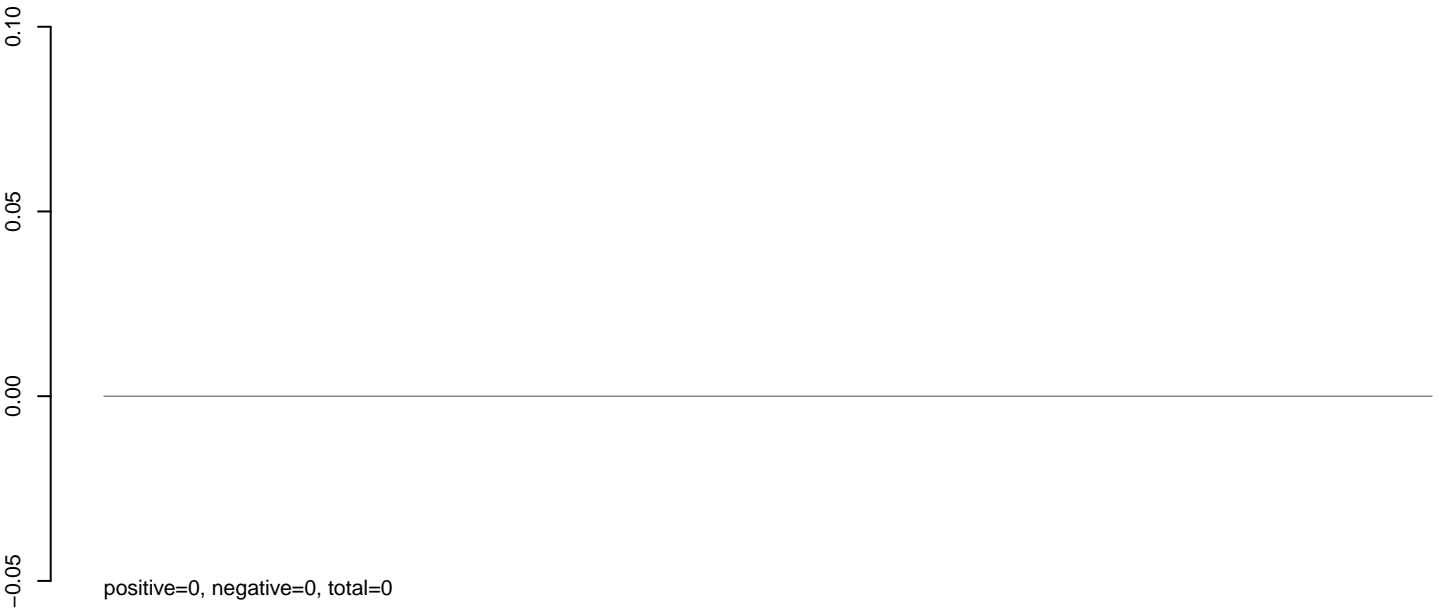
AeAeg_Aag2DENV2NGC.rep



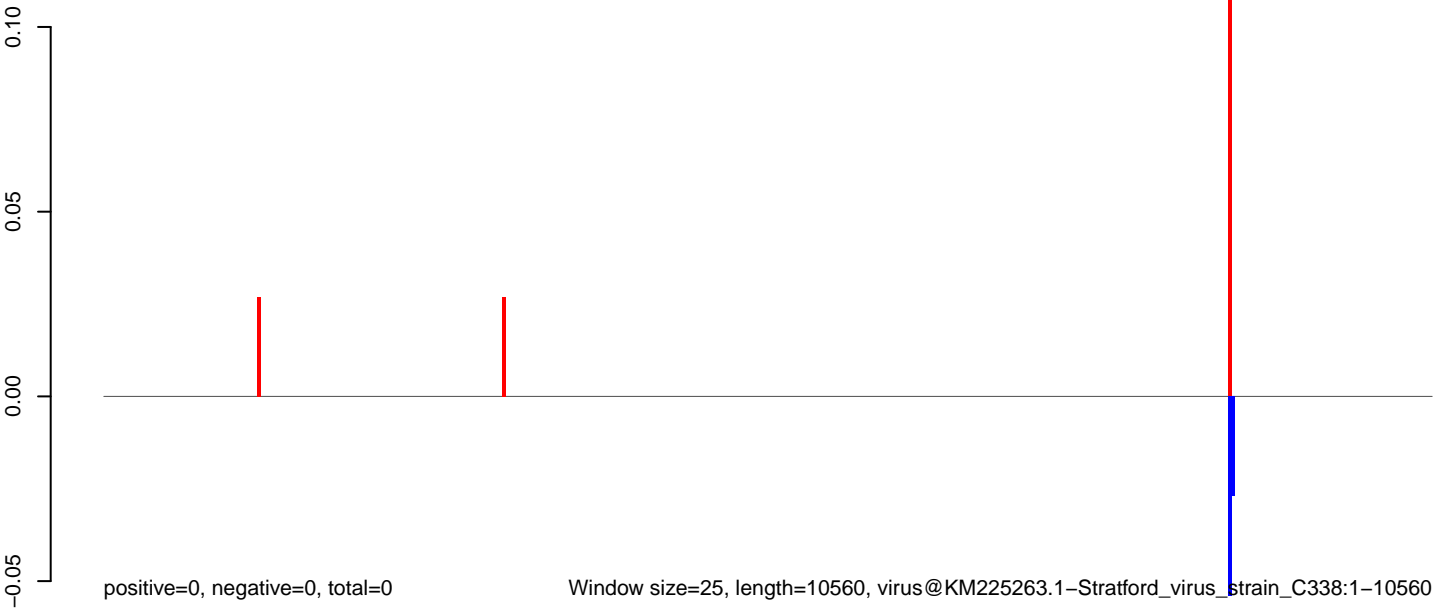
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



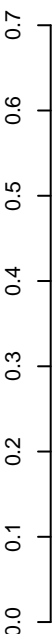
AeAeg_Aag2DENV2NGC.rep



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

0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



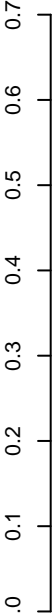
positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.24_35.rep

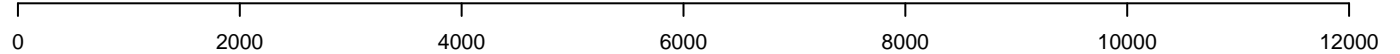


positive=1, negative=0, total=1

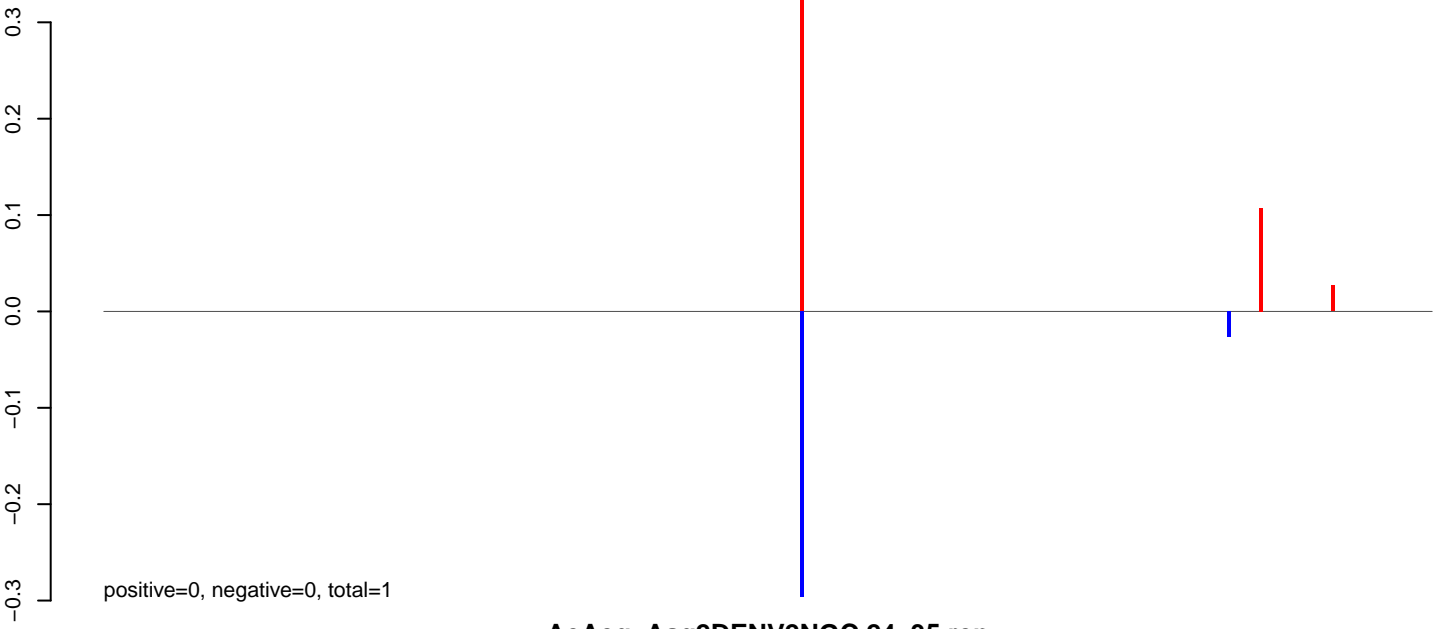
AeAeg_Aag2DENV2NGC.rep



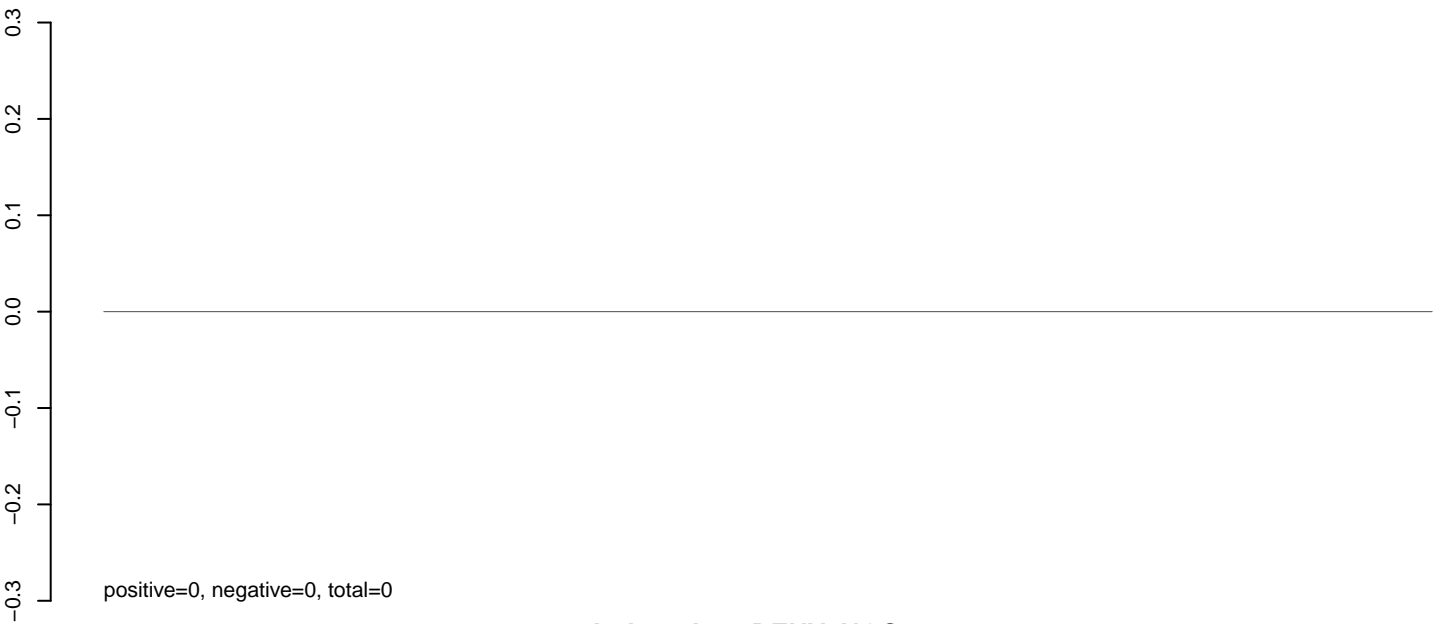
bow size=15, style=2, the1964-0, insa=12, M147992.1-Una_virus_non_structural_polyprotein_gene_partial_cds_and_structural_polyprotein_ge:1-11964



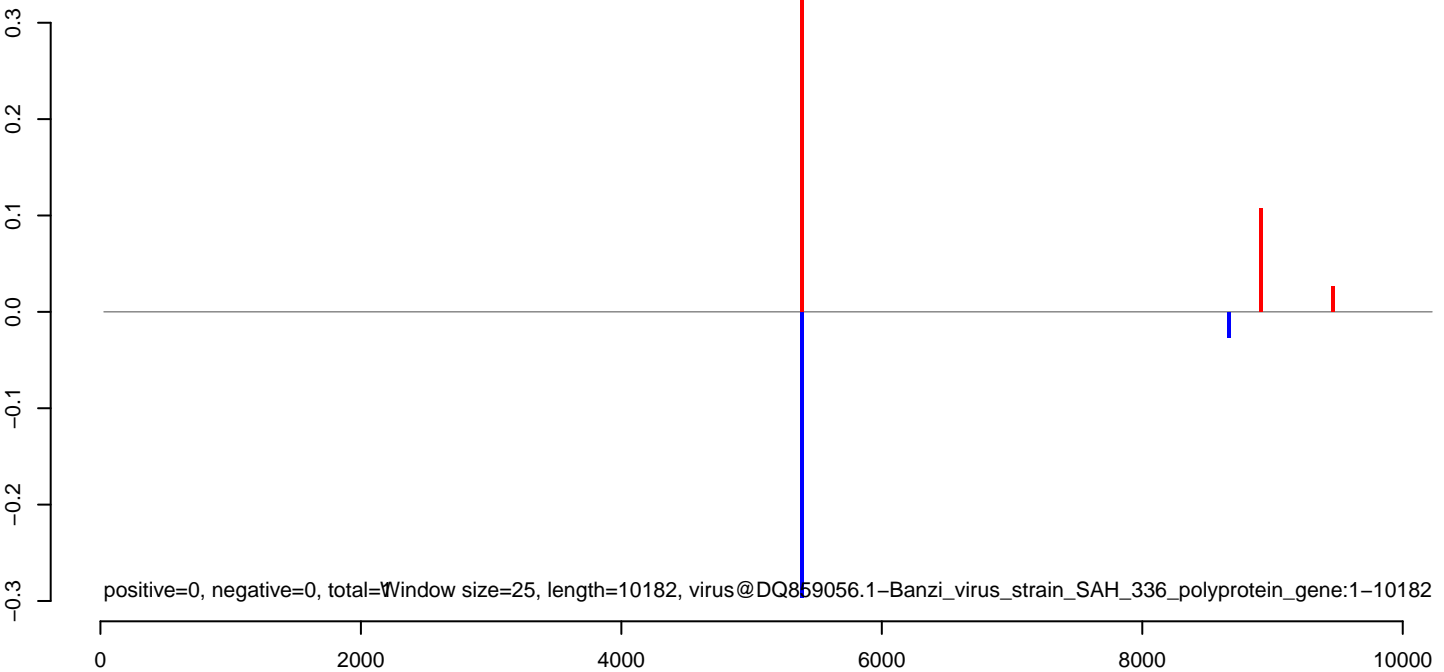
AeAeg_Aag2DENV2NGC.18_23.rep



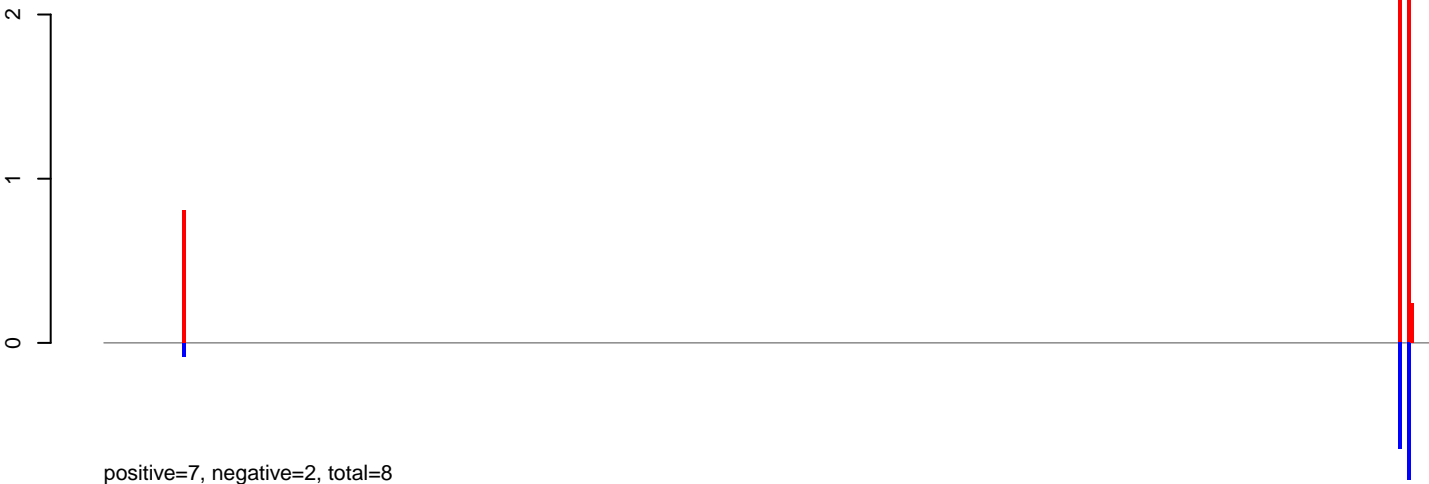
AeAeg_Aag2DENV2NGC.24_35.rep



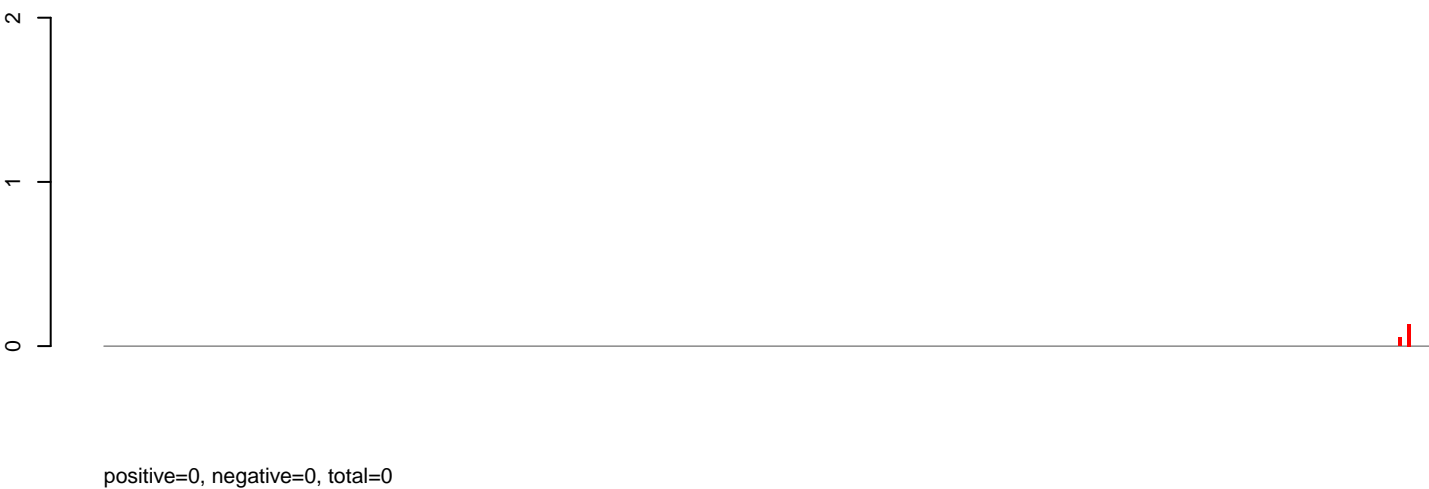
AeAeg_Aag2DENV2NGC.rep



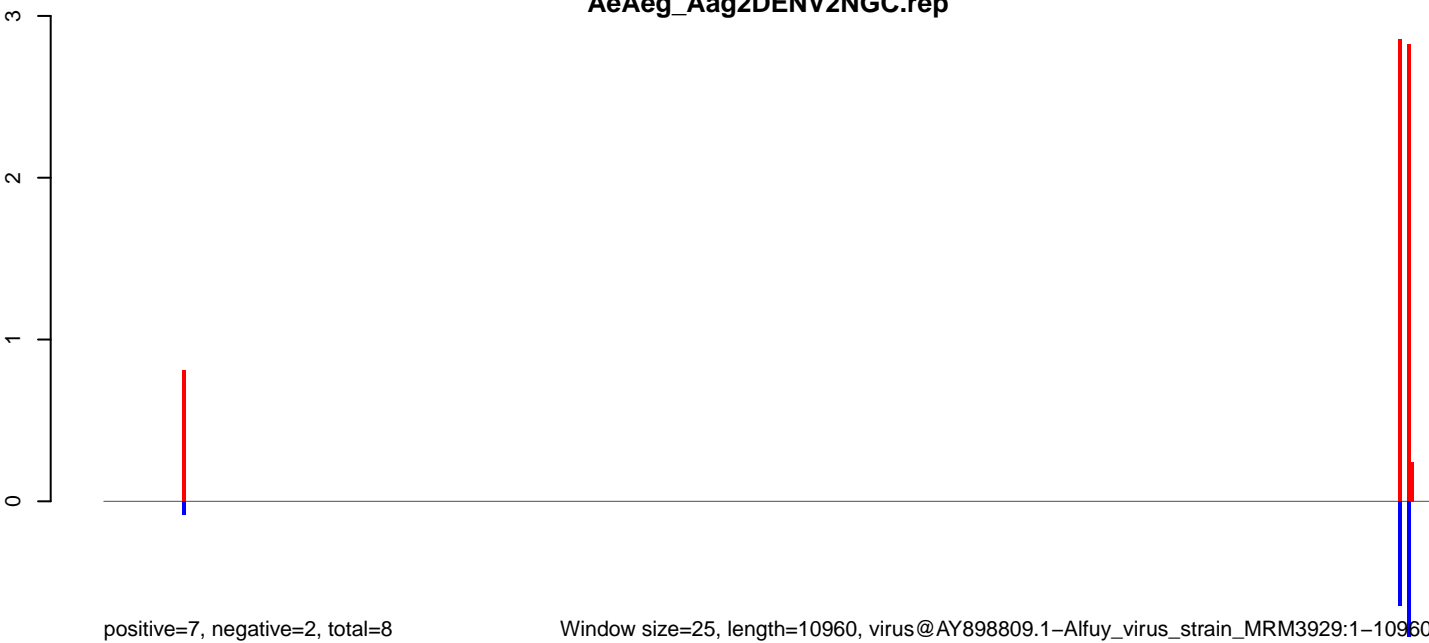
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



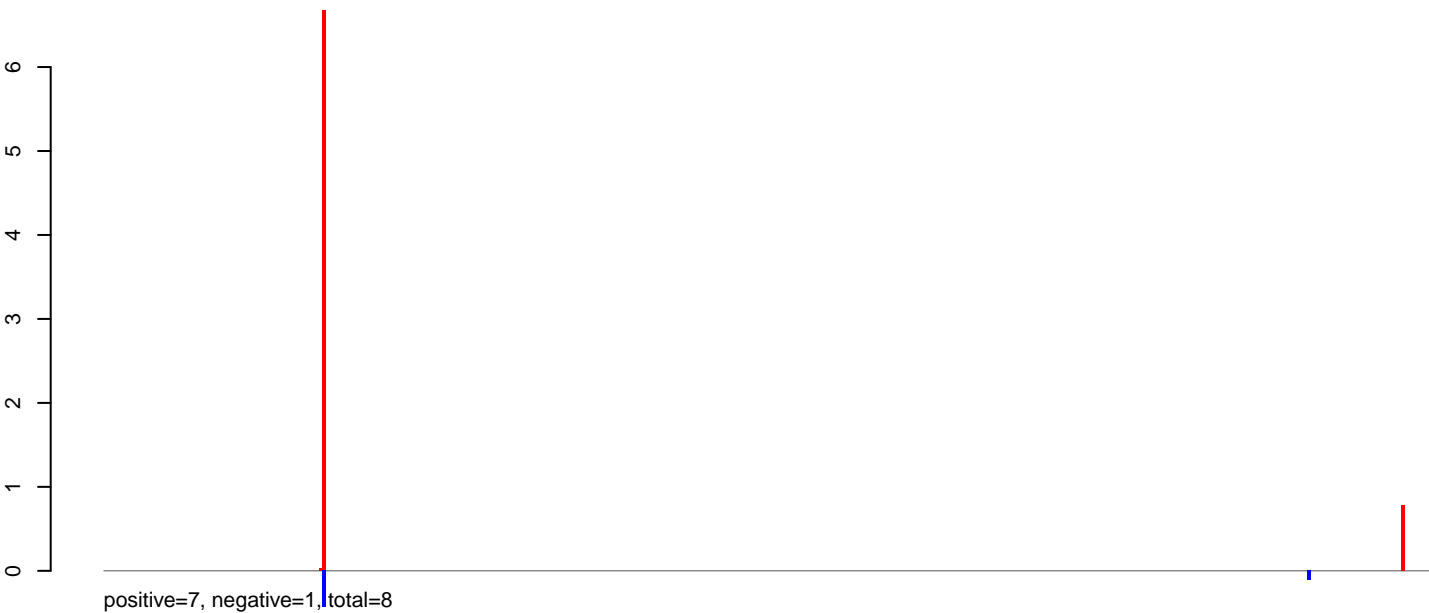
AeAeg_Aag2DENV2NGC.rep



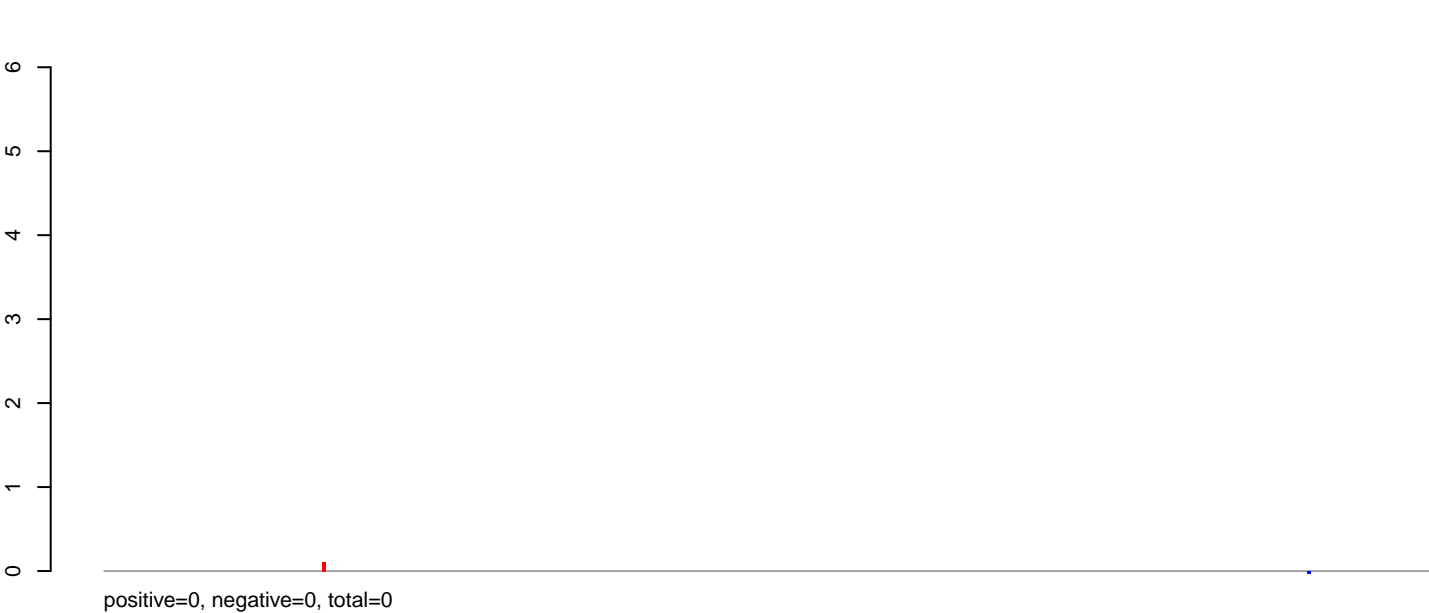
Window size=25, length=10960, virus@AY898809.1-Alfuy_virus_strain_MRM3929:1-10960

0 2000 4000 6000 8000 10000

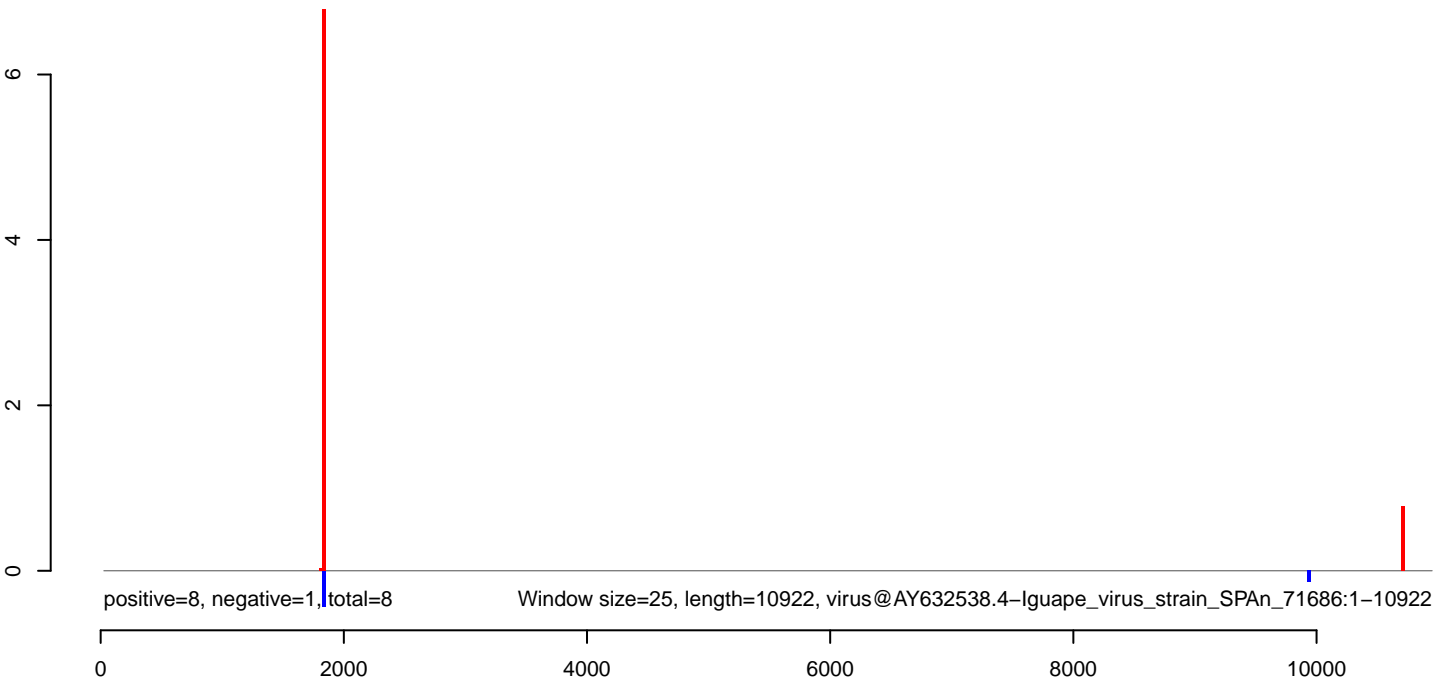
AeAeg_Aag2DENV2NGC.18_23.rep



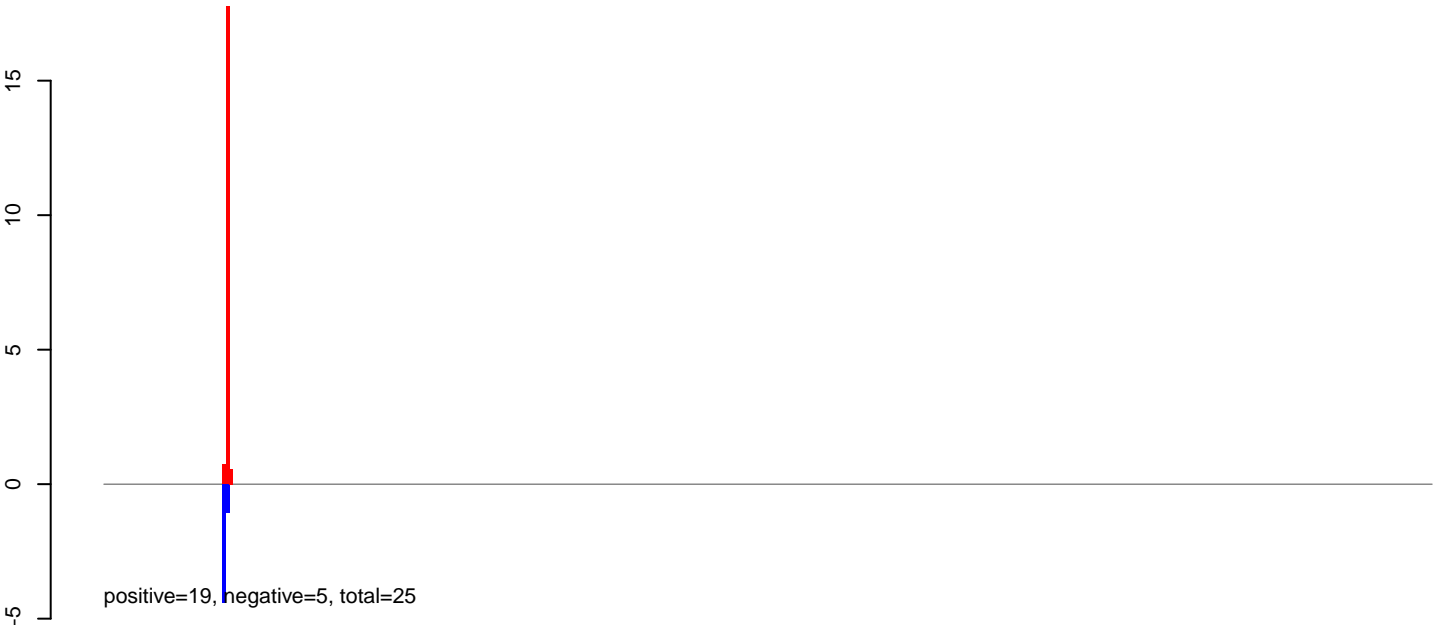
AeAeg_Aag2DENV2NGC.24_35.rep



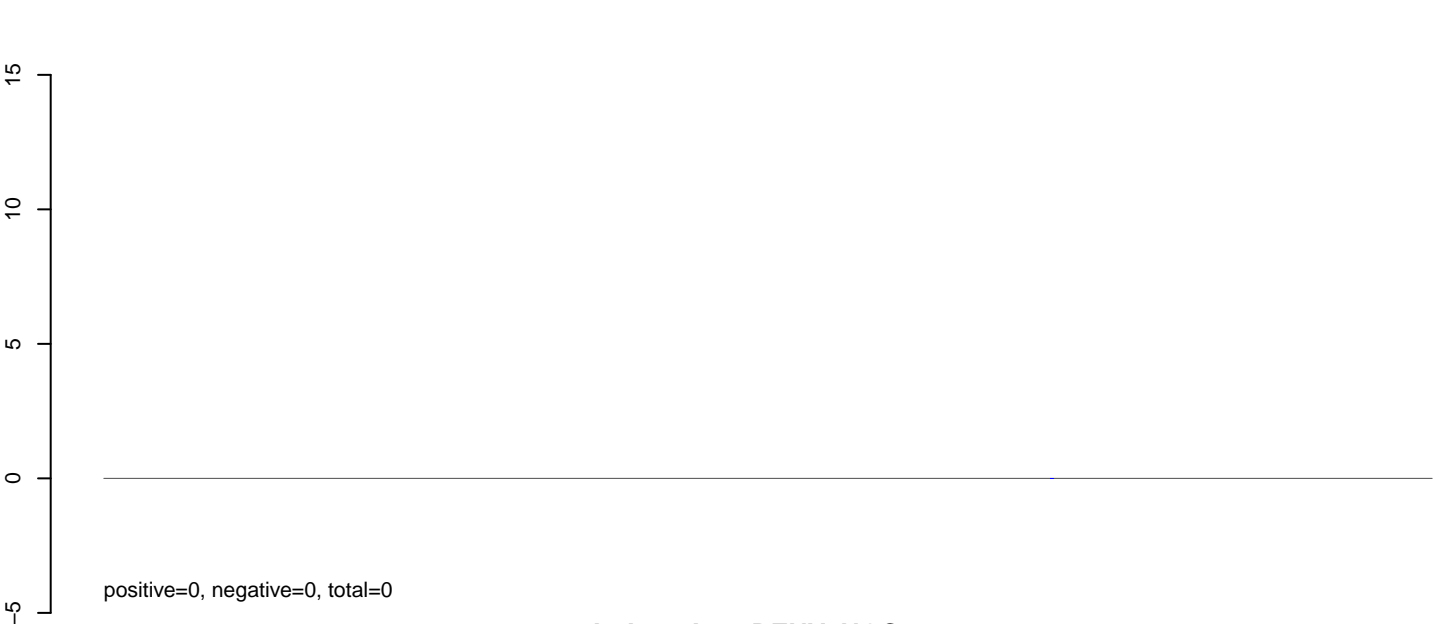
AeAeg_Aag2DENV2NGC.rep



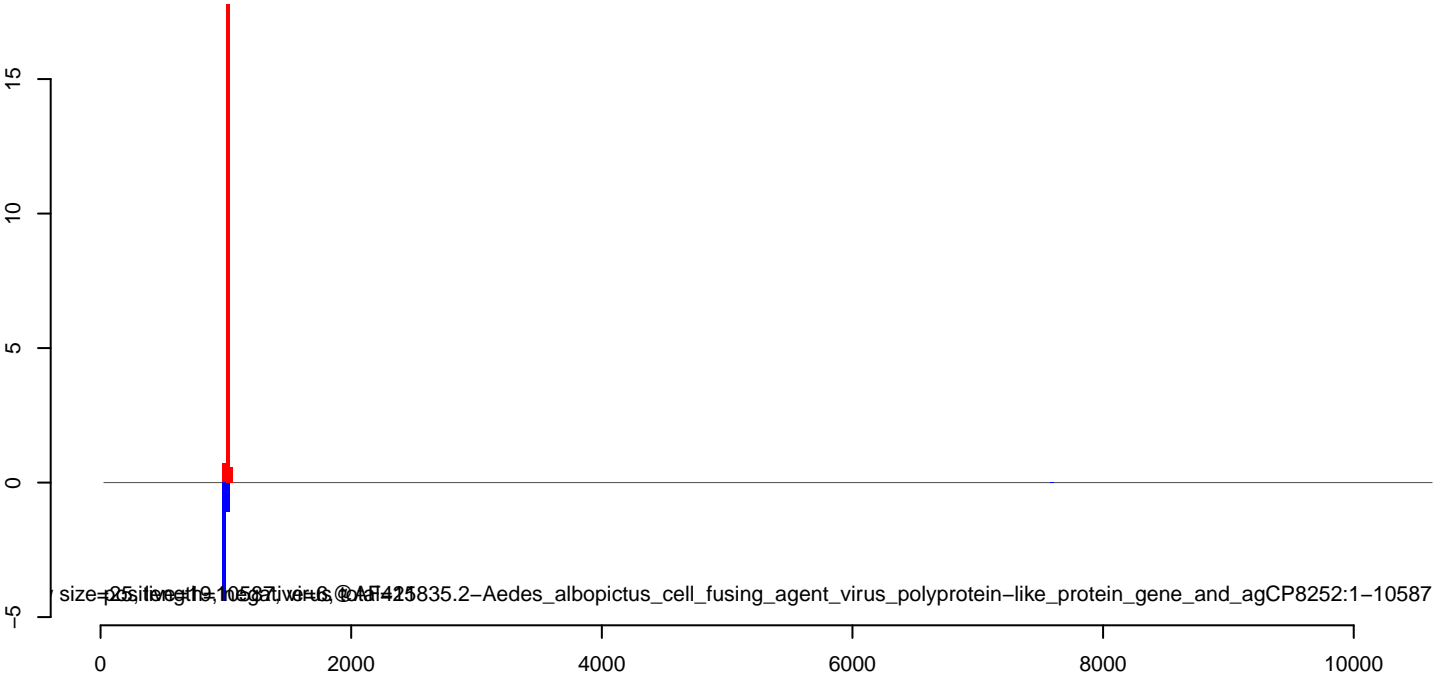
AeAeg_Aag2DENV2NGC.18_23.rep



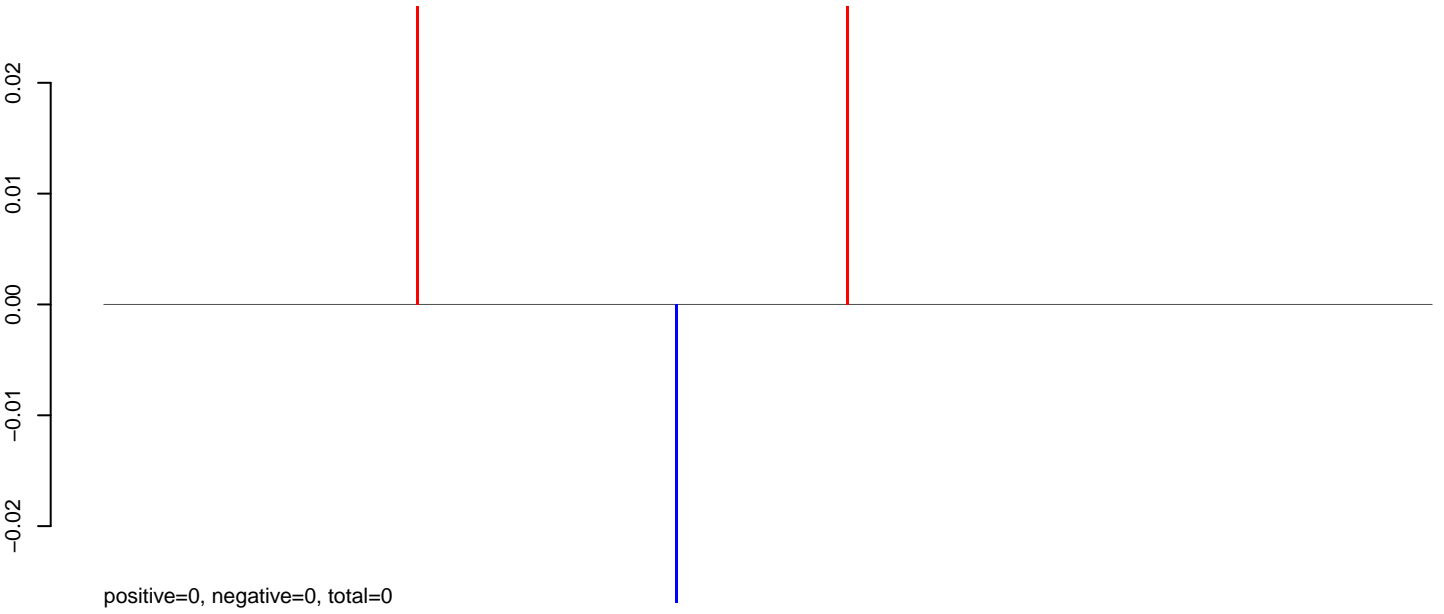
AeAeg_Aag2DENV2NGC.24_35.rep



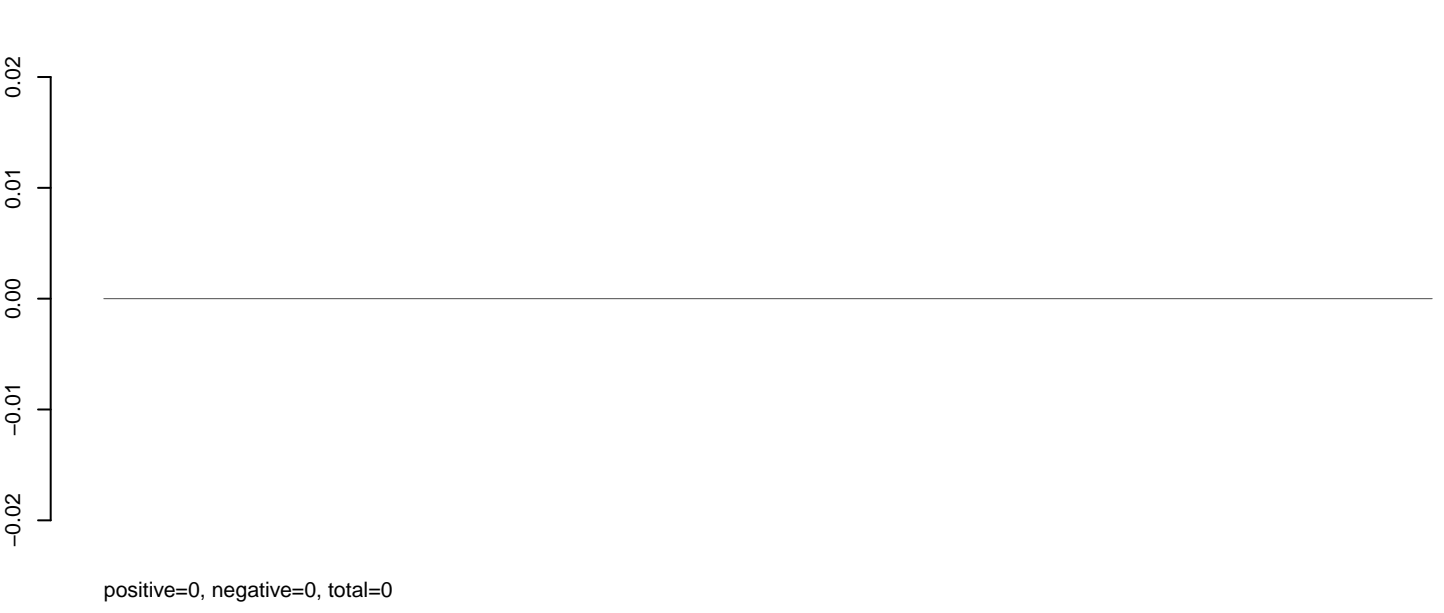
AeAeg_Aag2DENV2NGC.rep



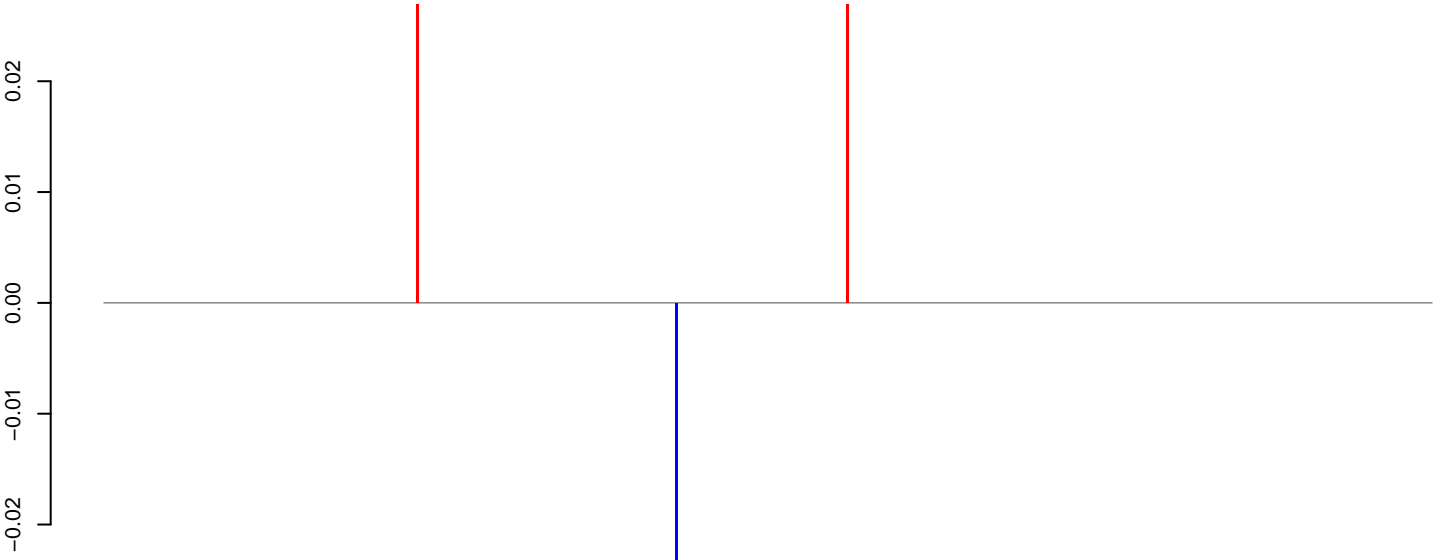
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



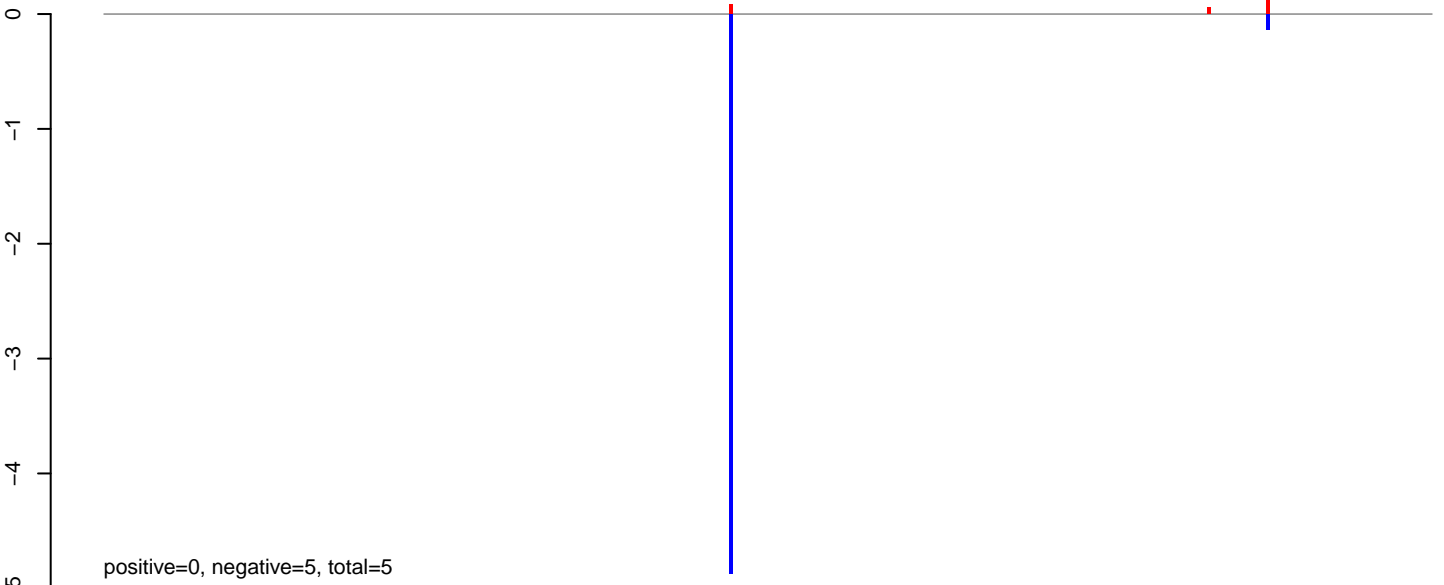
AeAeg_Aag2DENV2NGC.rep



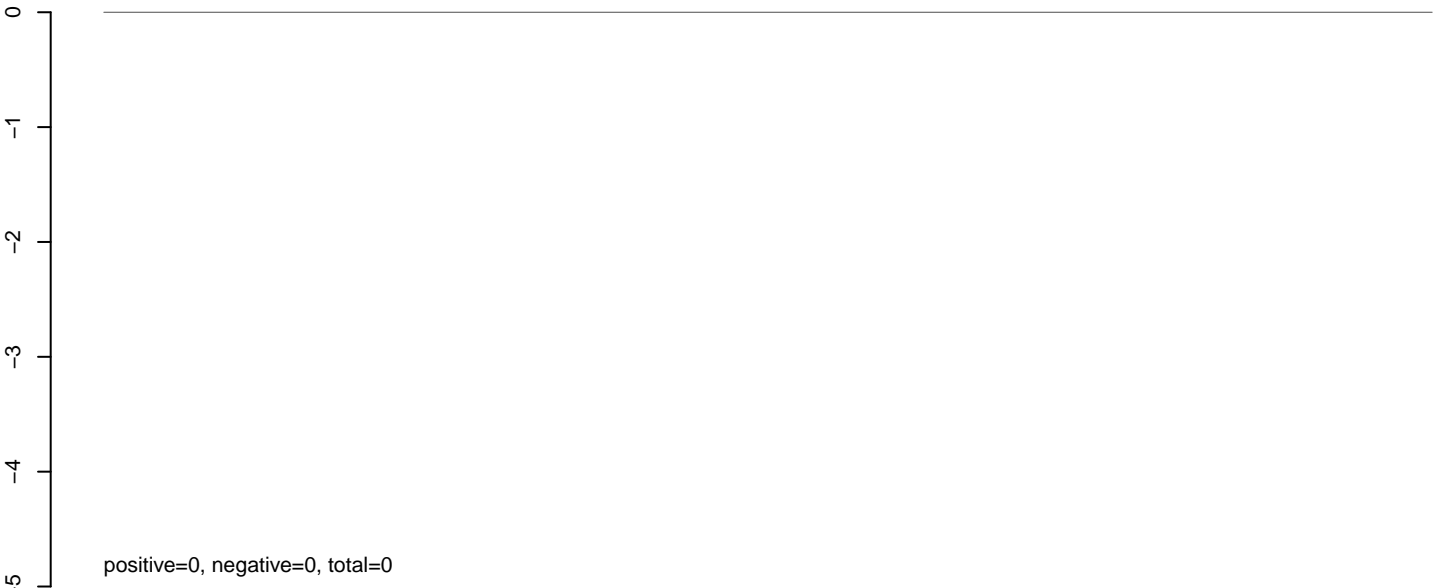
NC_003453.1-Rochambeau_virus_nucleoprotein_phosphoprotein_matrix_hypothetical_proteins_glycopr:1-13593

0 2000 4000 6000 8000 10000 12000 14000

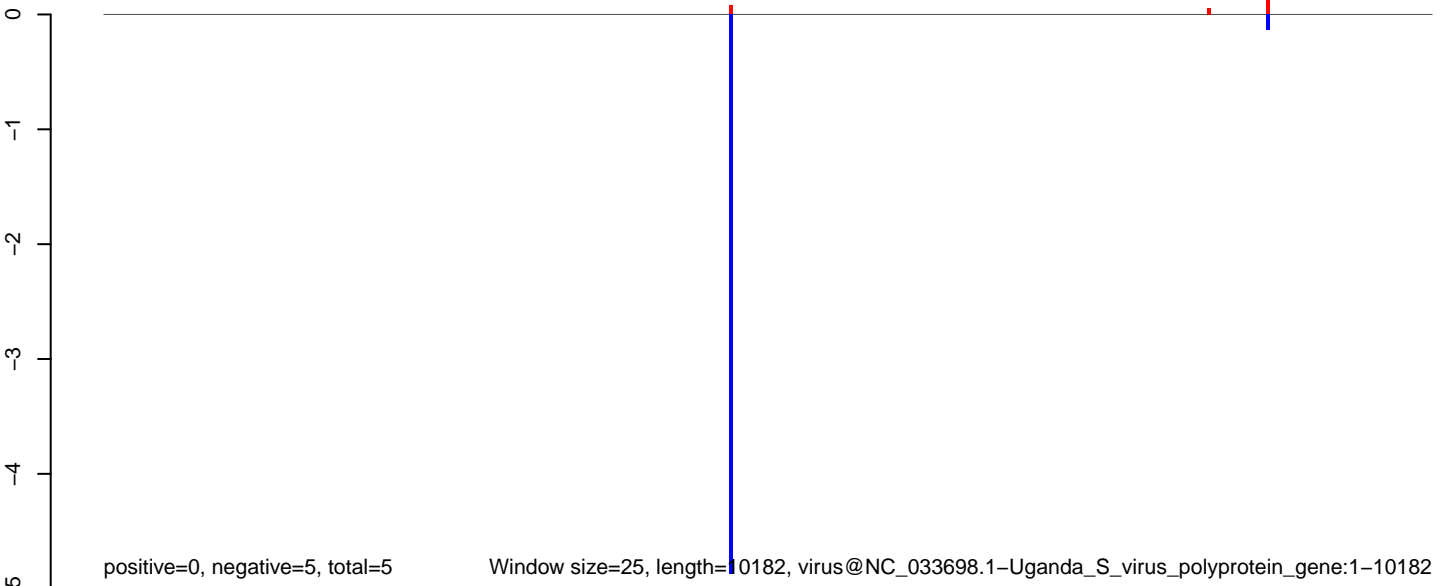
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



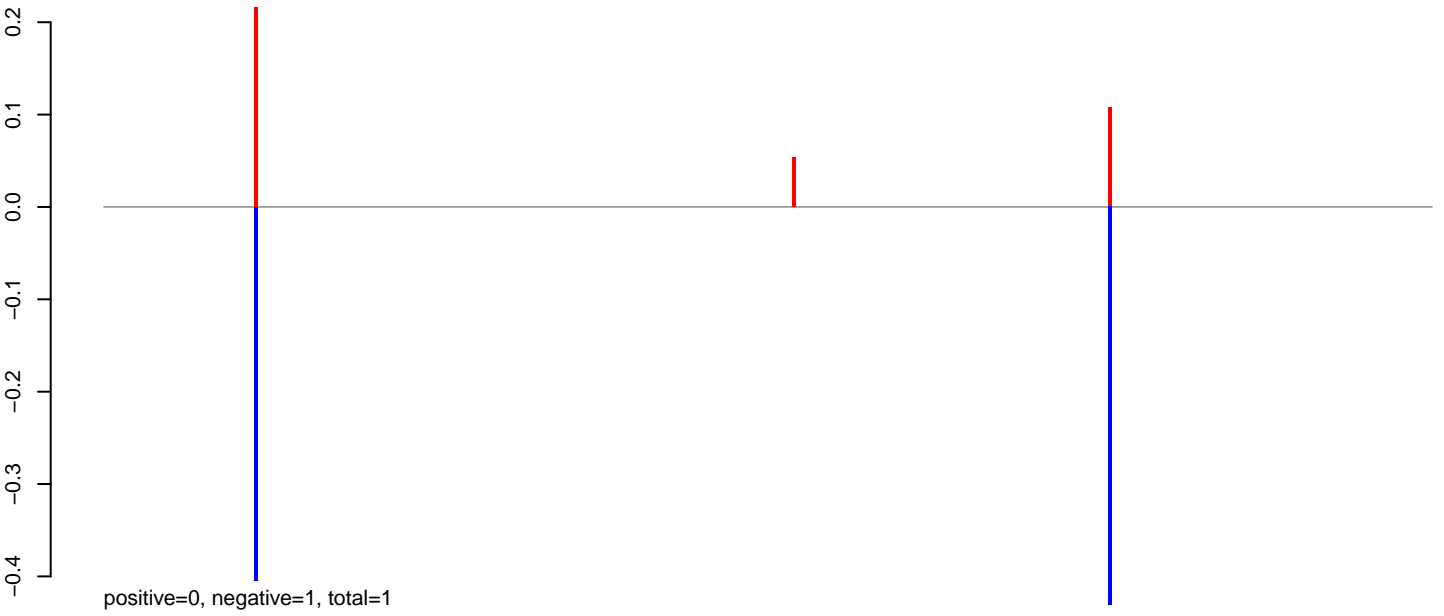
AeAeg_Aag2DENV2NGC.rep



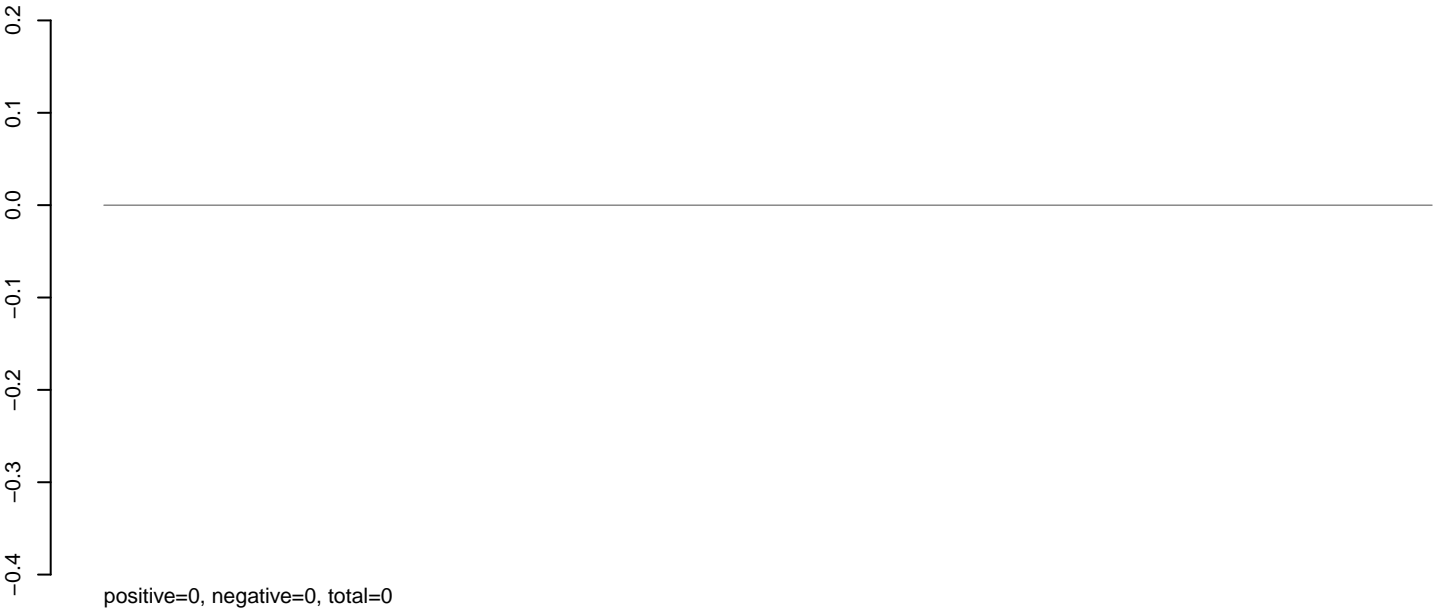
Window size=25, length=10182, virus@NC_033698.1-Uganda_S_virus_polyprotein_gene:1-10182

0 2000 4000 6000 8000 10000

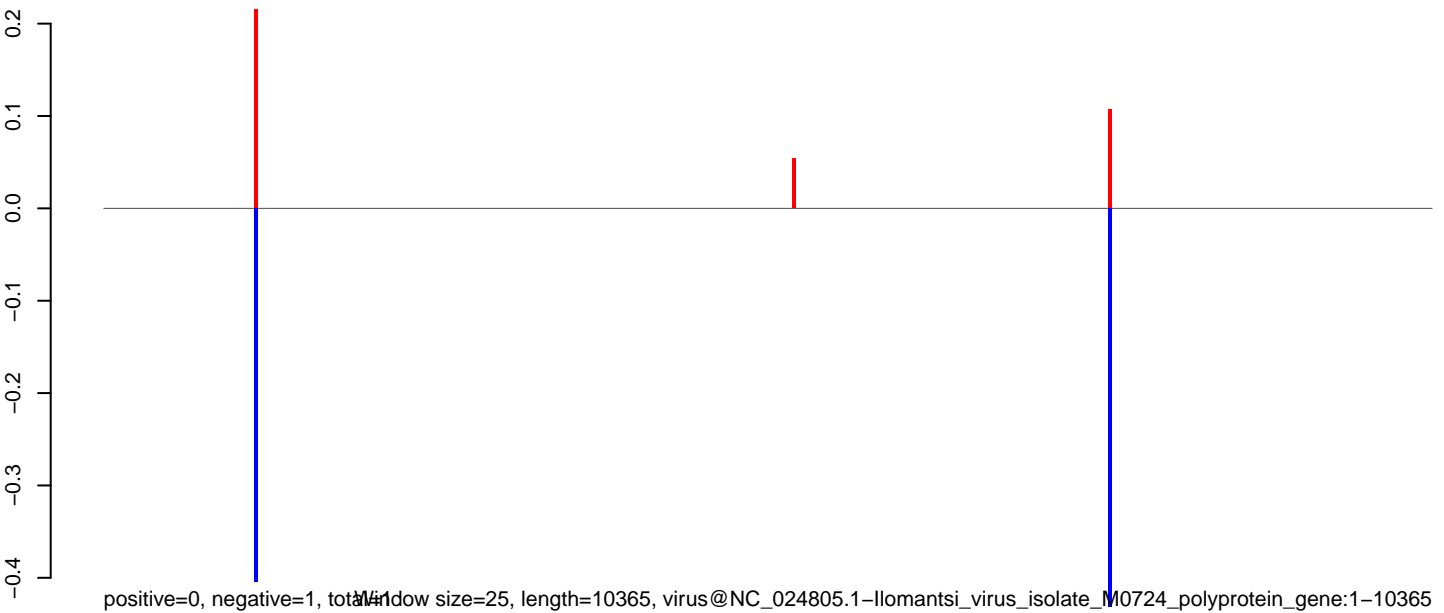
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



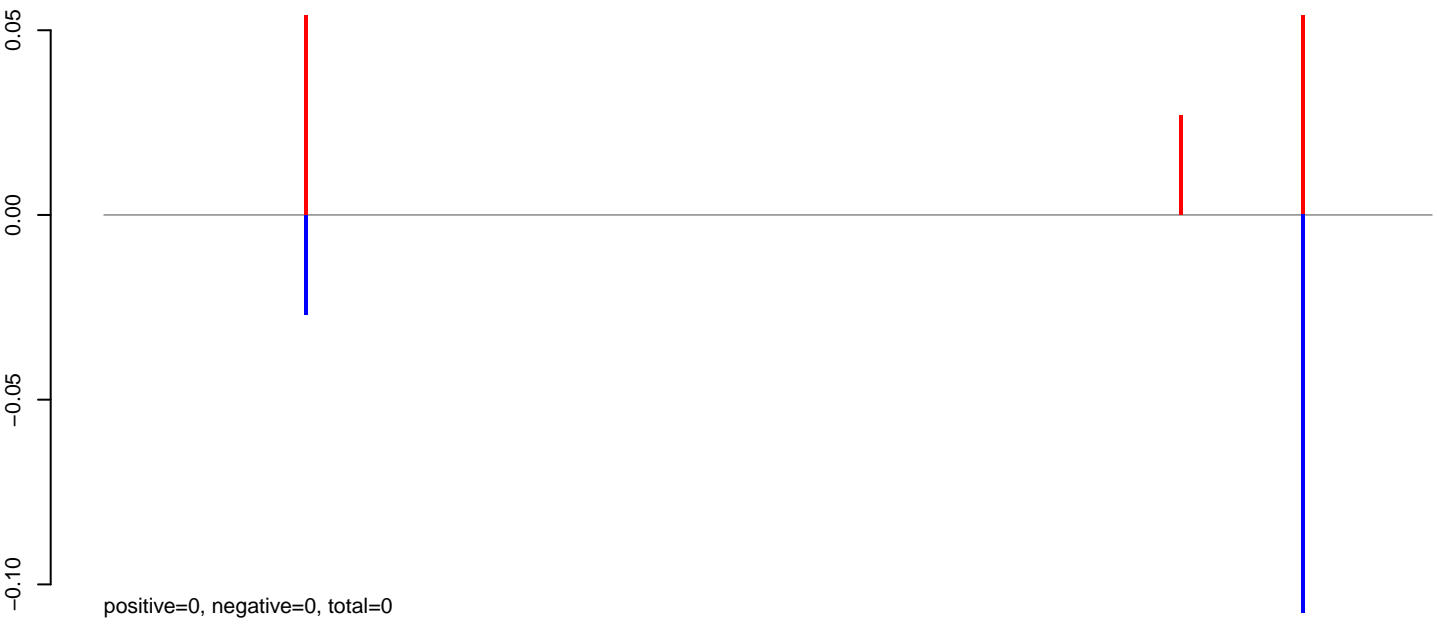
AeAeg_Aag2DENV2NGC.rep



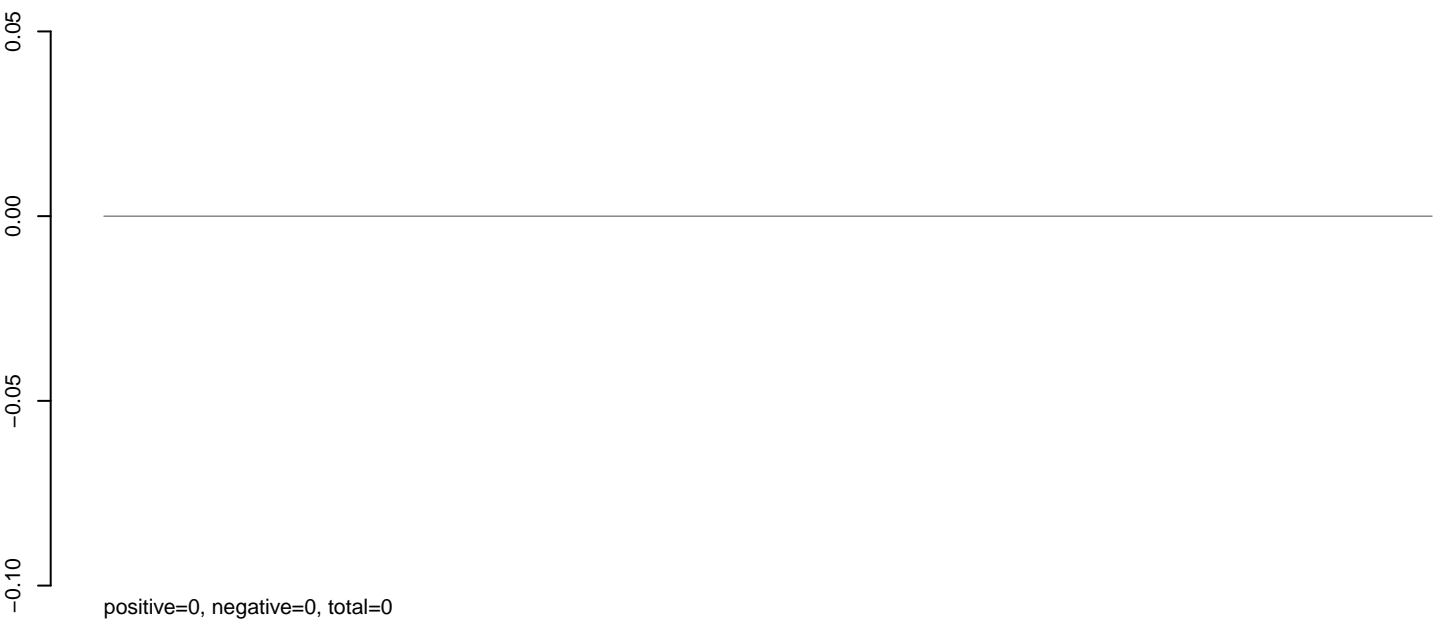
Window size=25, length=10365, virus@NC_024805.1-Ilomantsi_virus_isolate_M0724_polyprotein_gene:1-10365

0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



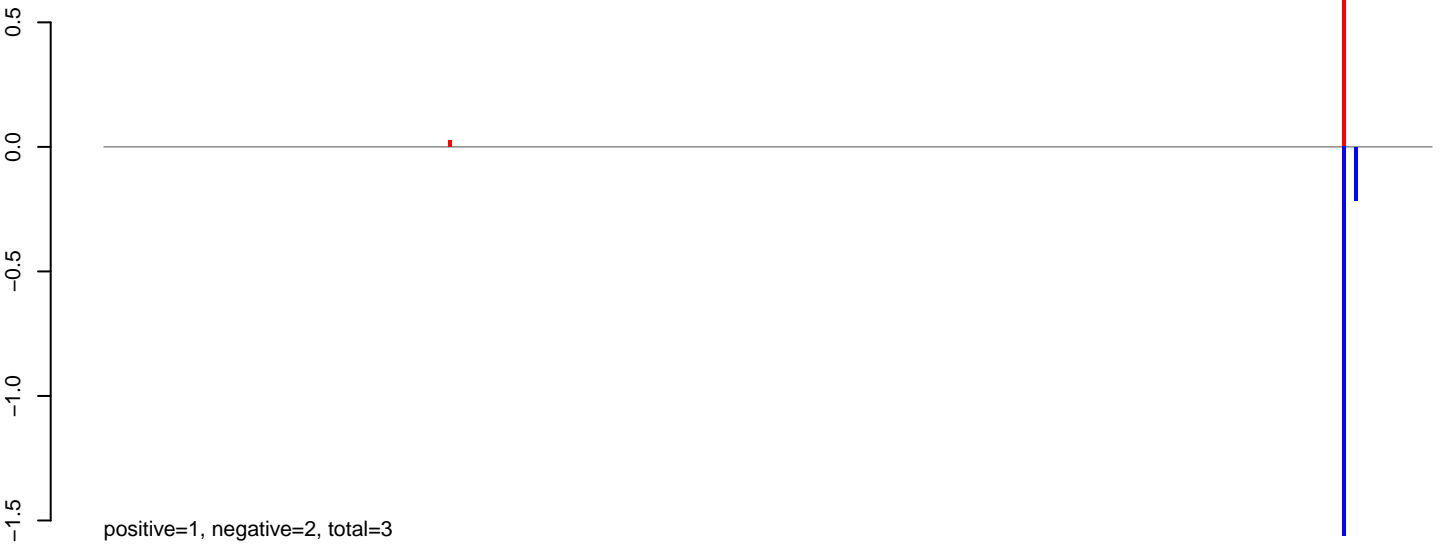
AeAeg_Aag2DENV2NGC.24_35.rep



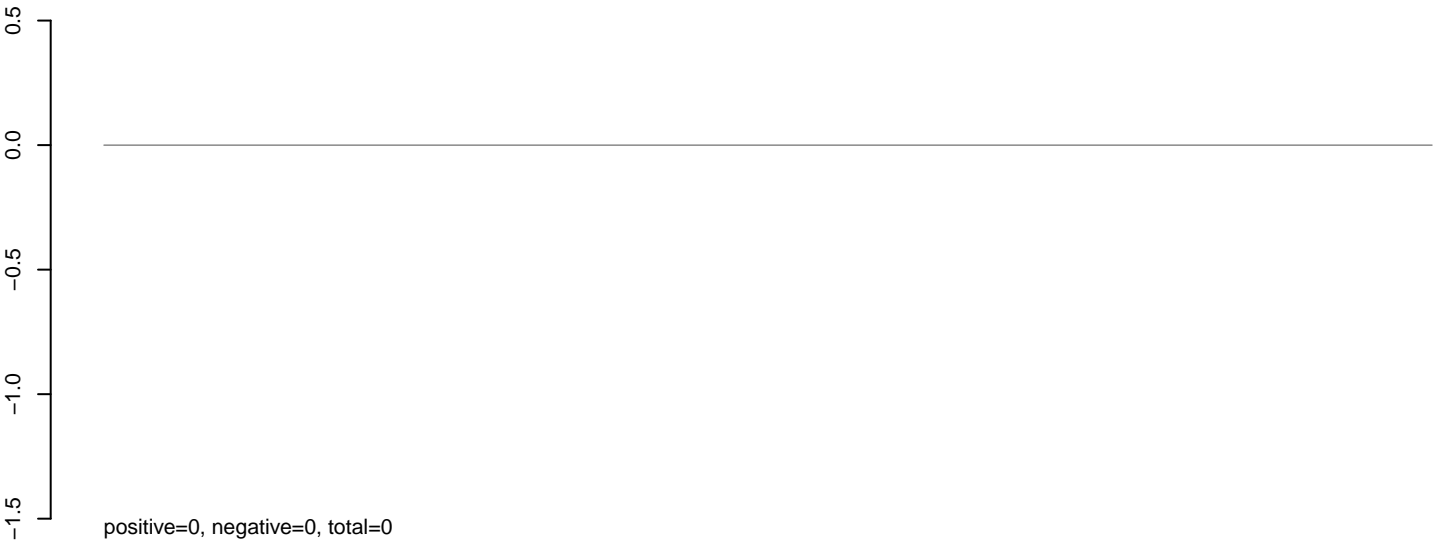
AeAeg_Aag2DENV2NGC.rep



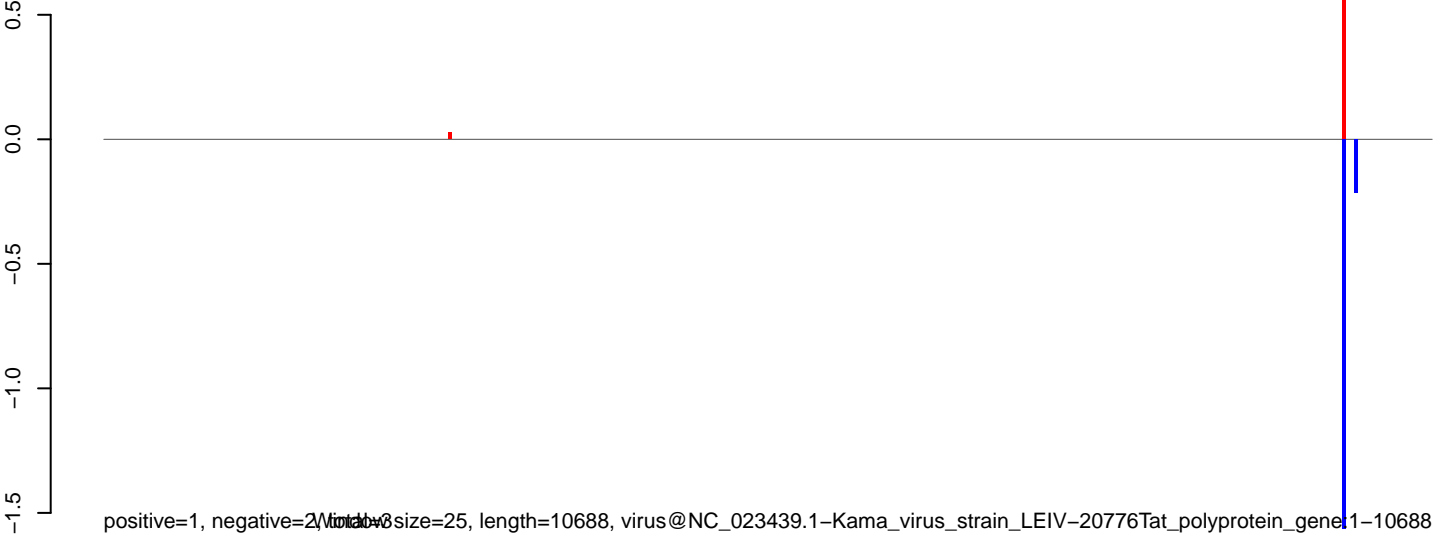
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



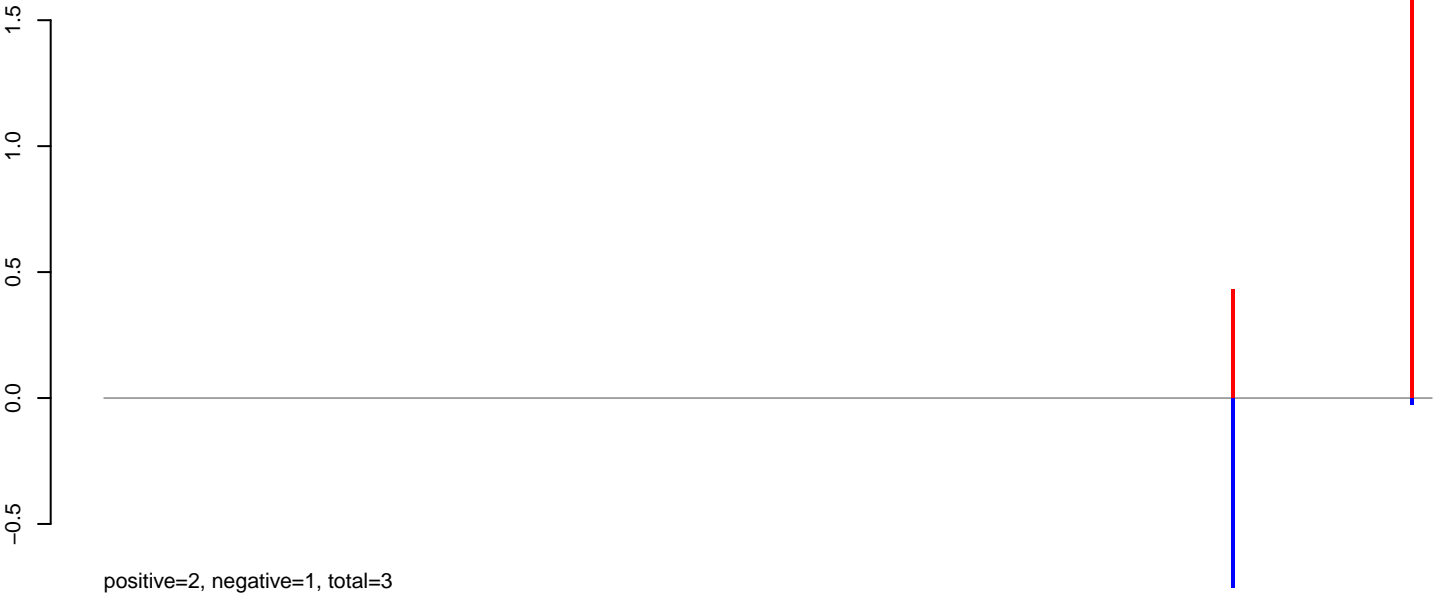
AeAeg_Aag2DENV2NGC.rep



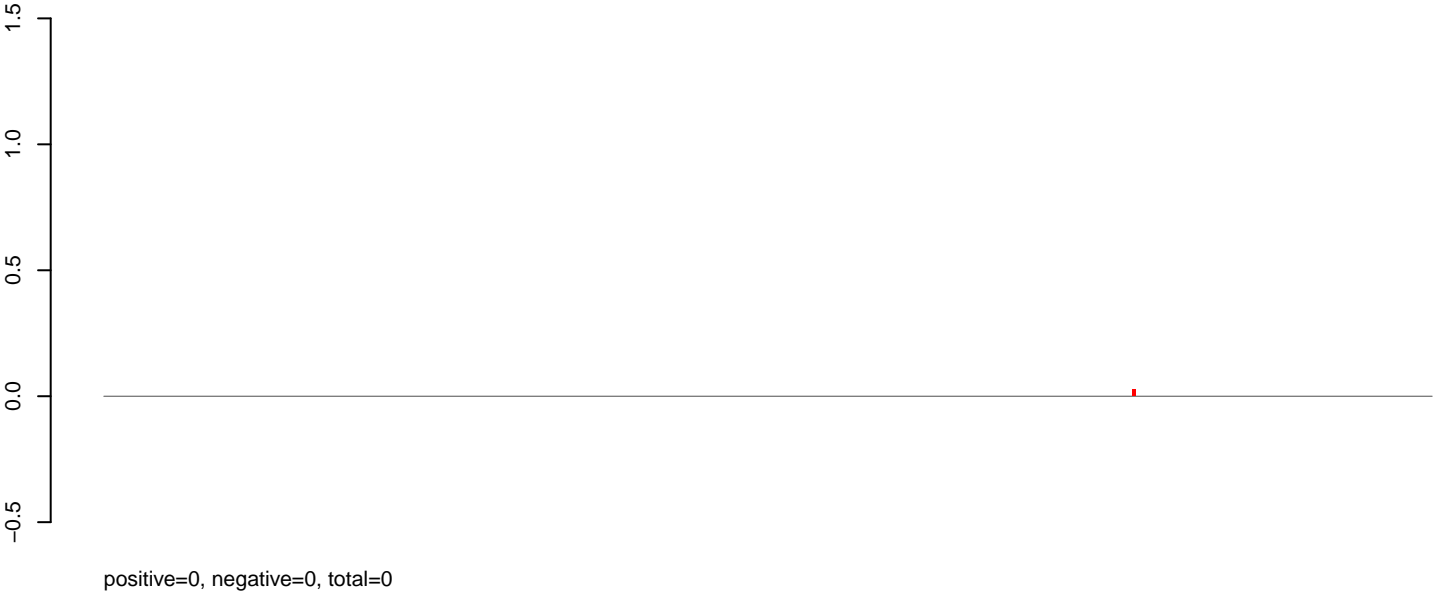
width=8 size=25, length=10688, virus@NC_023439.1-Kama_virus_strain_LEIV-20776Tat_polyprotein_gene1-10688

0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



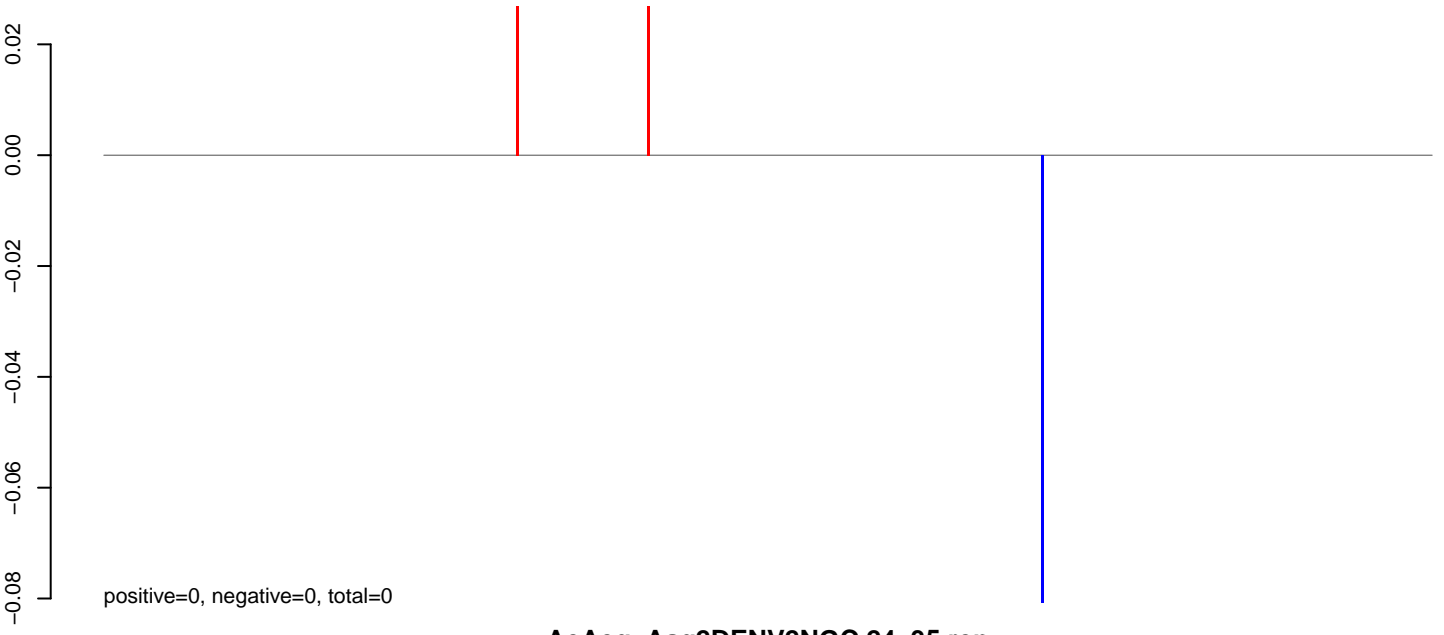
AeAeg_Aag2DENV2NGC.rep



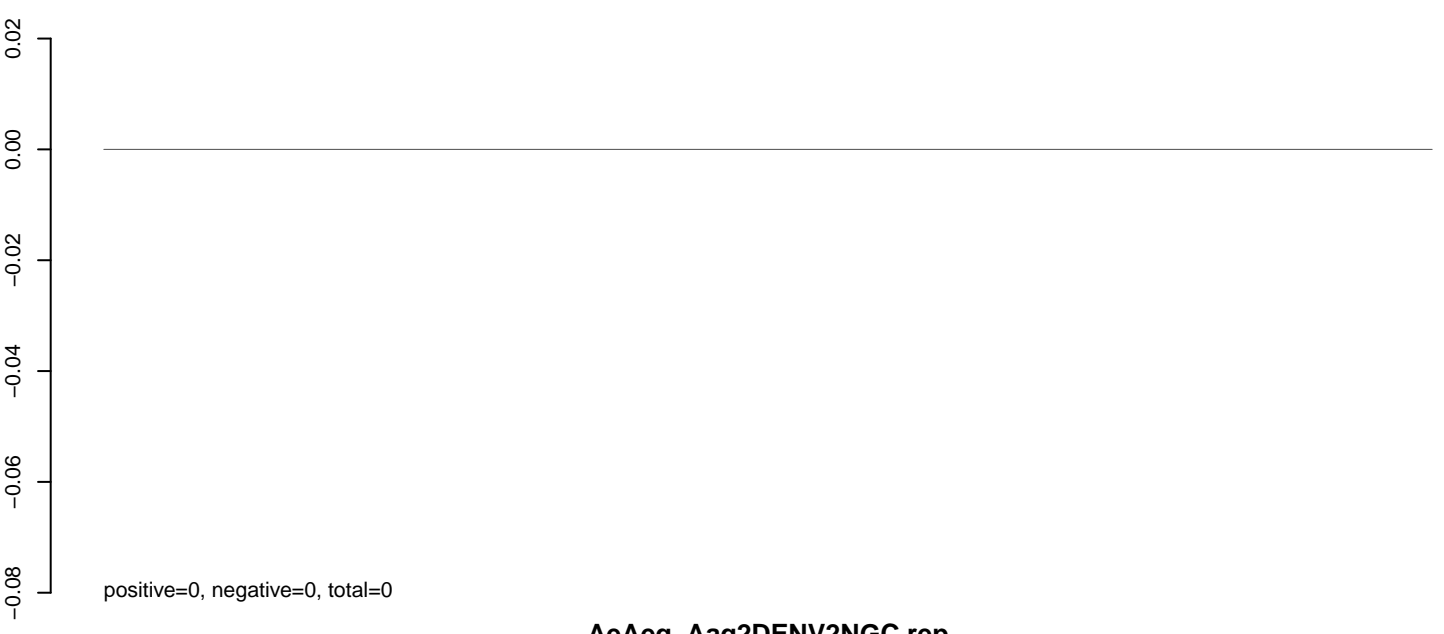
Window size=25, length=10733, virus@NC_017086.1-Chaoyang_virus:1-10733

0 2000 4000 6000 8000 10000

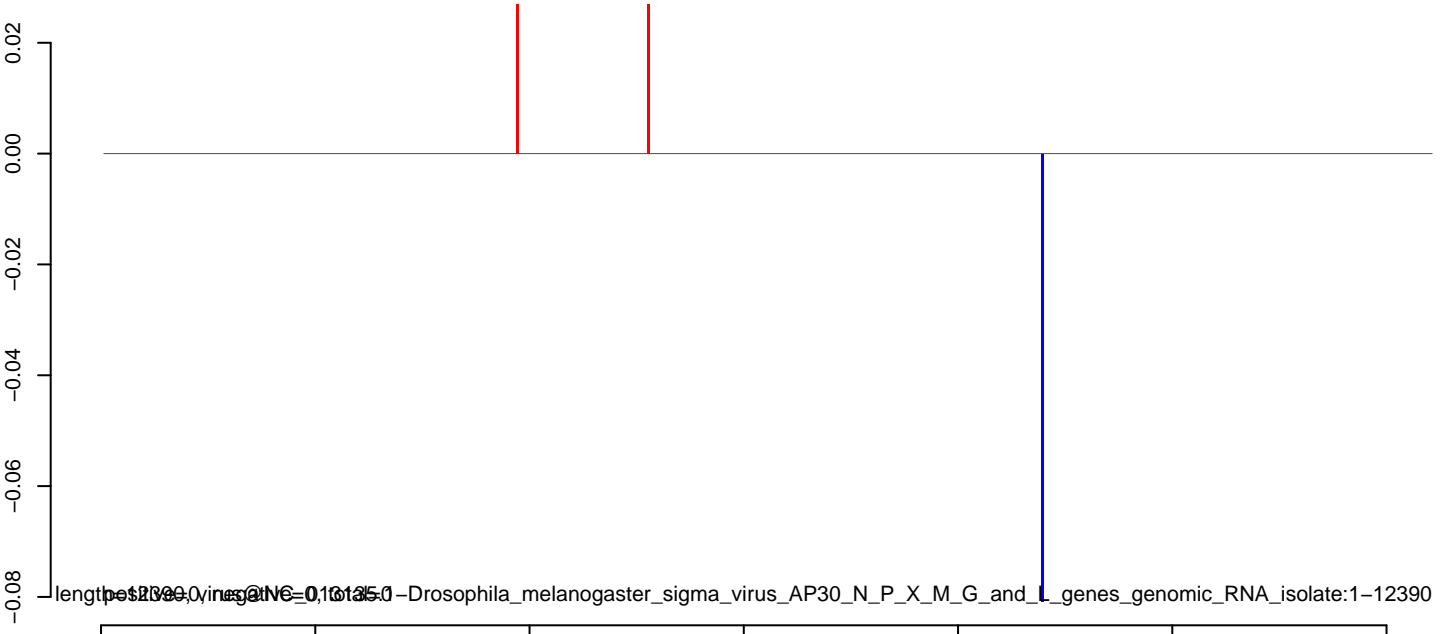
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



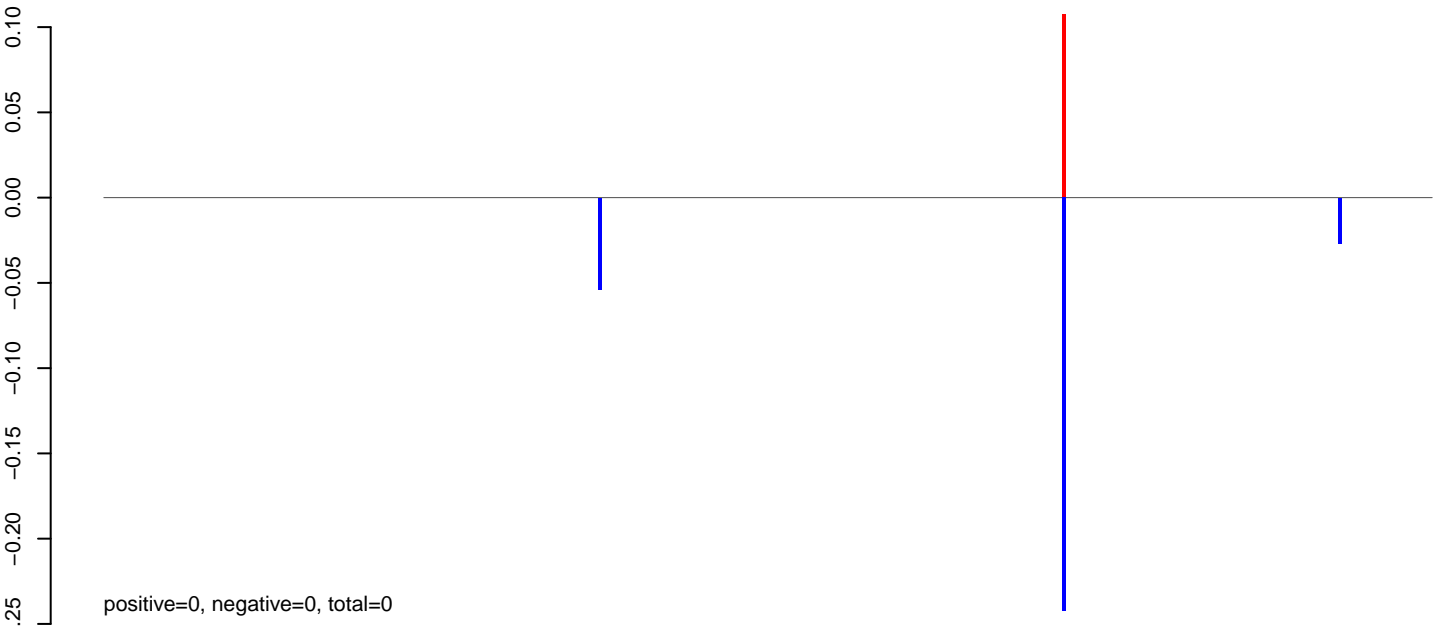
AeAeg_Aag2DENV2NGC.rep



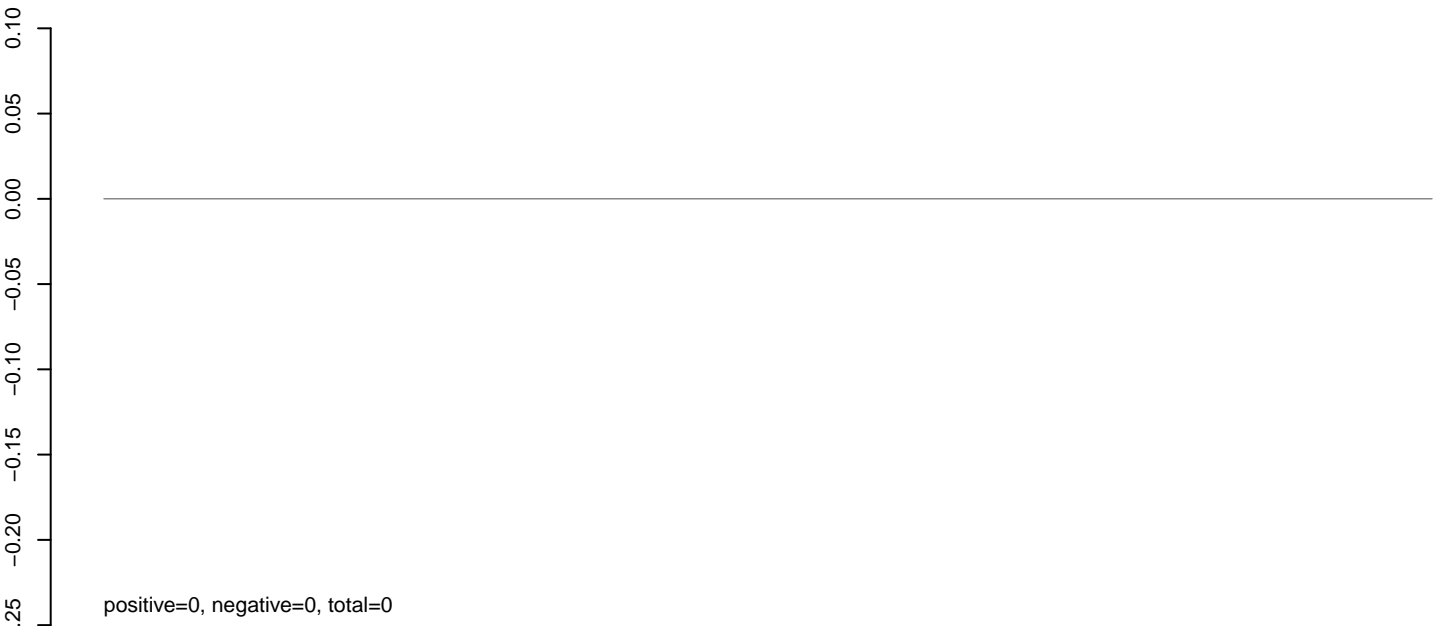
length=12390, positive=0, negative=0, total=0 -Drosophila_melanogaster_sigma_virus_AP30_N_P_X_M_G_and_L_genes_genomic_RNA_isolate:1-12390

0 2000 4000 6000 8000 10000 12000

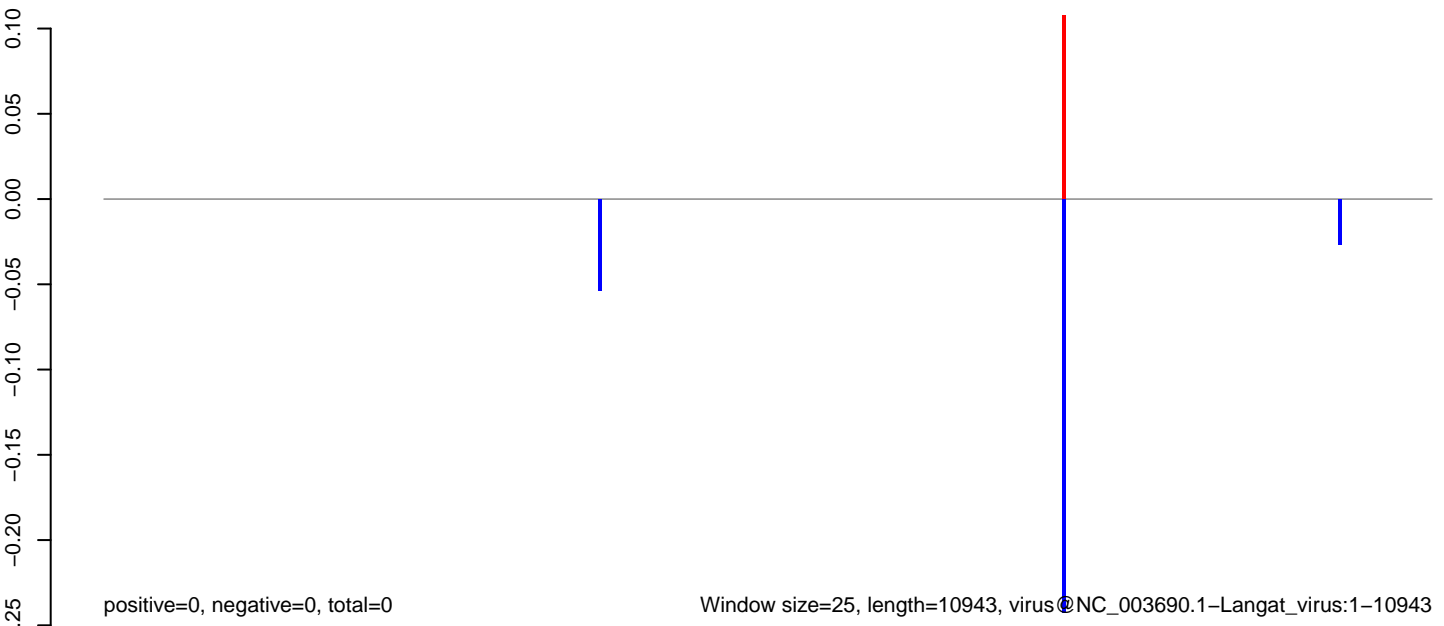
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

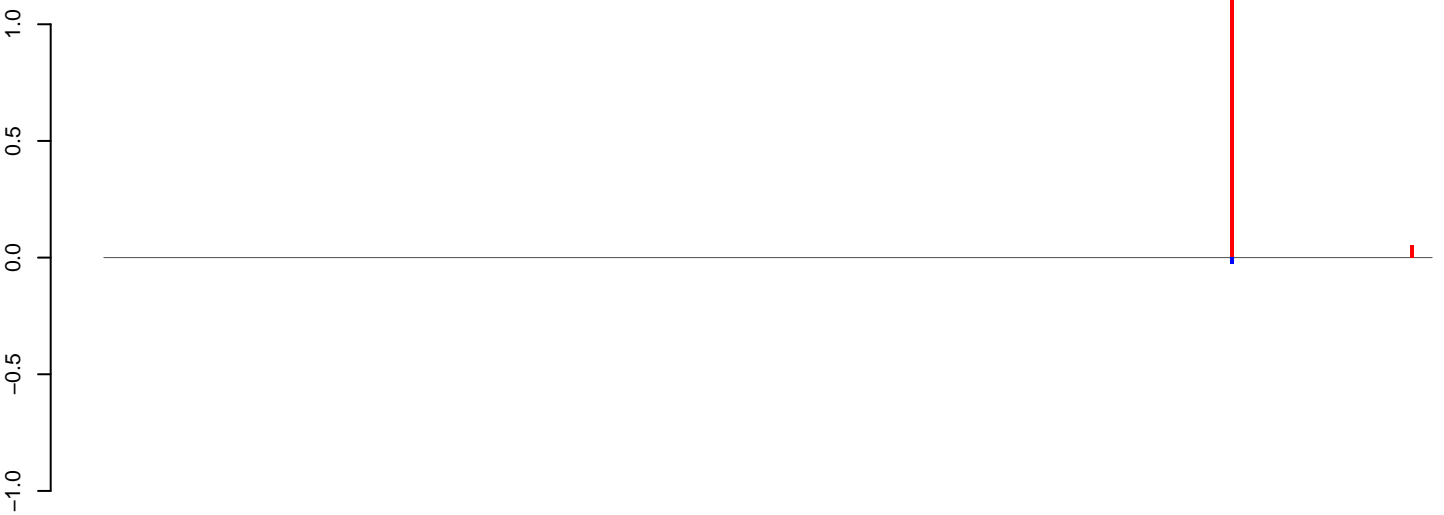


AeAeg_Aag2DENV2NGC.rep



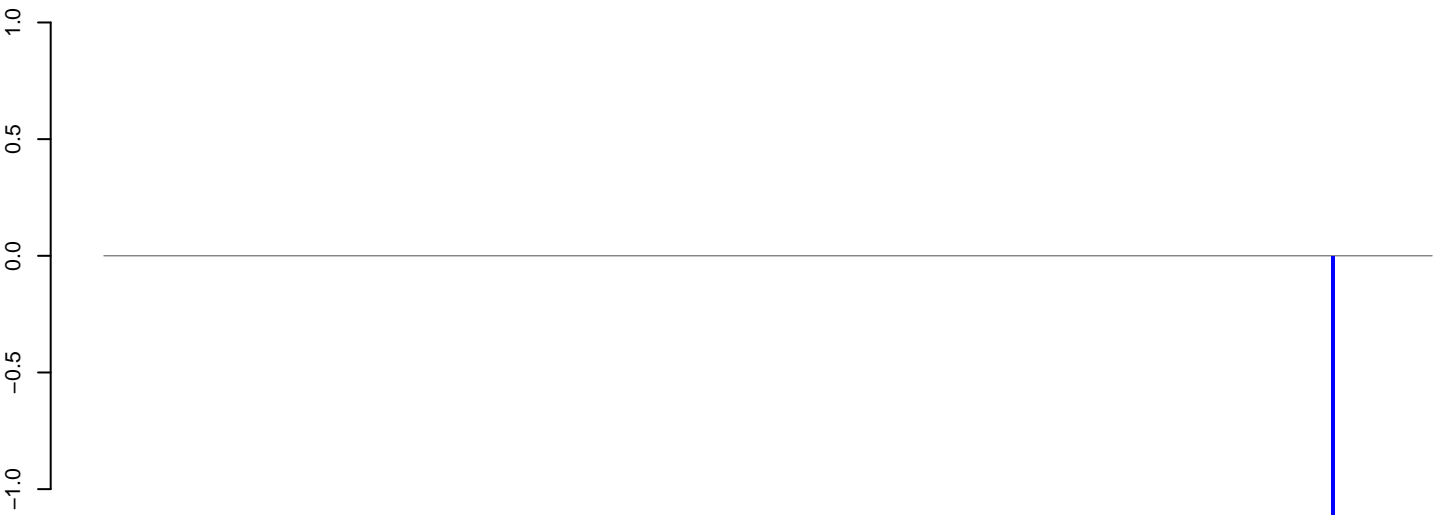
0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



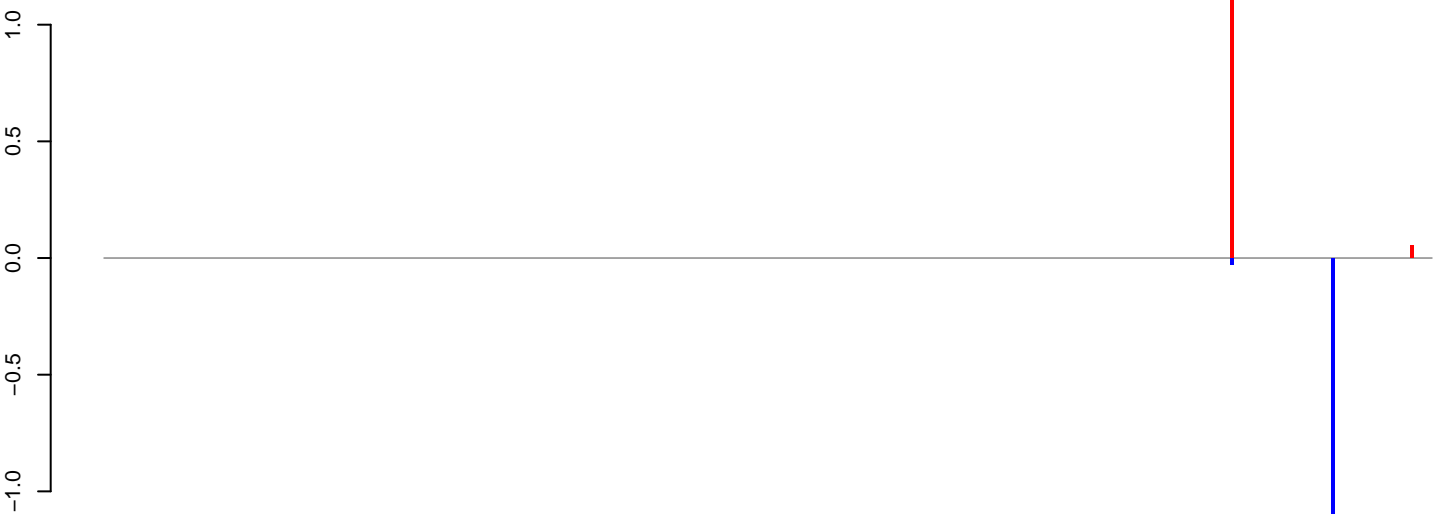
positive=1, negative=0, total=1

AeAeg_Aag2DENV2NGC.24_35.rep



positive=0, negative=1, total=1

AeAeg_Aag2DENV2NGC.rep

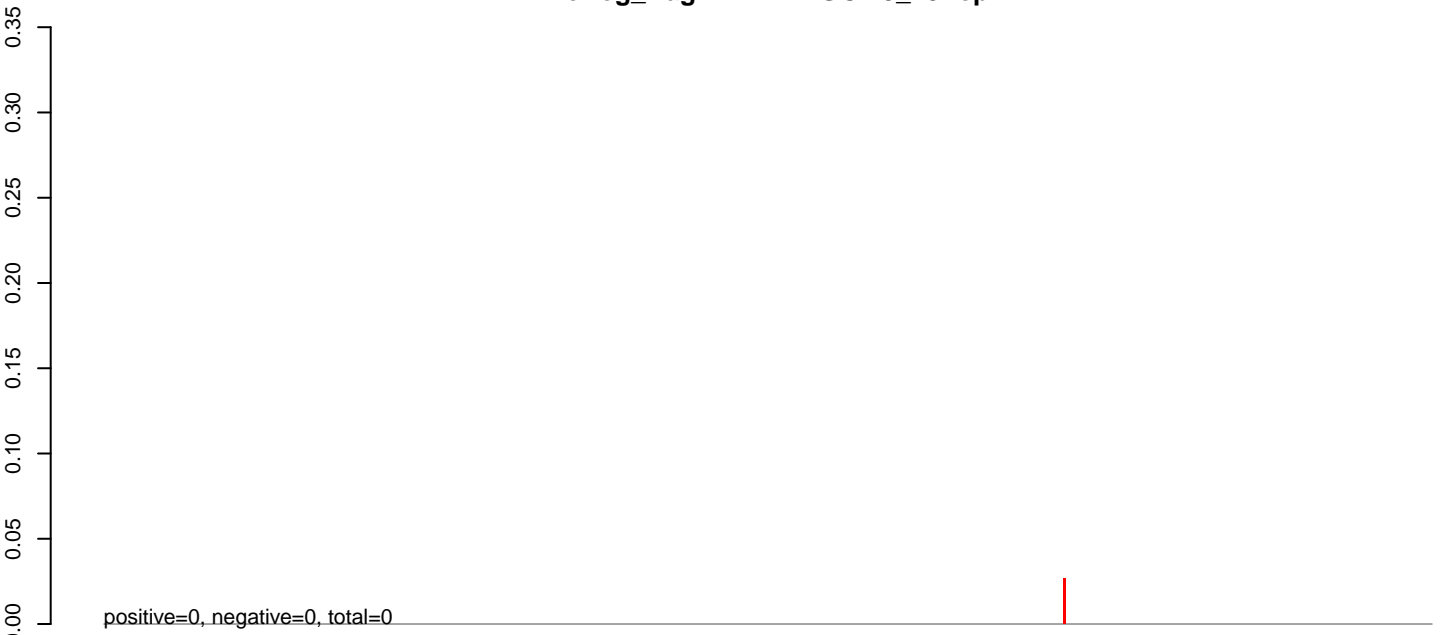


positive=1, negative=1, total=3

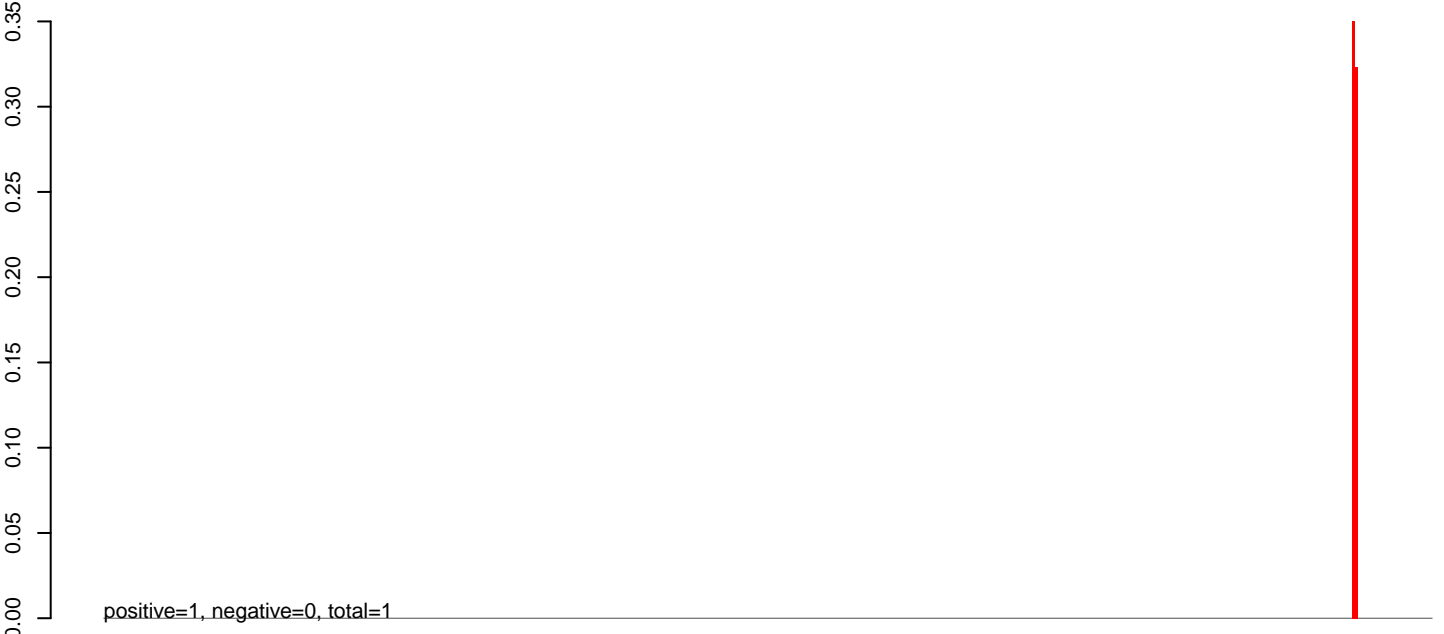
Window size=25, length=10862, virus@NC_002031.1-Yellow_fever_virus:1-10862

0 2000 4000 6000 8000 10000

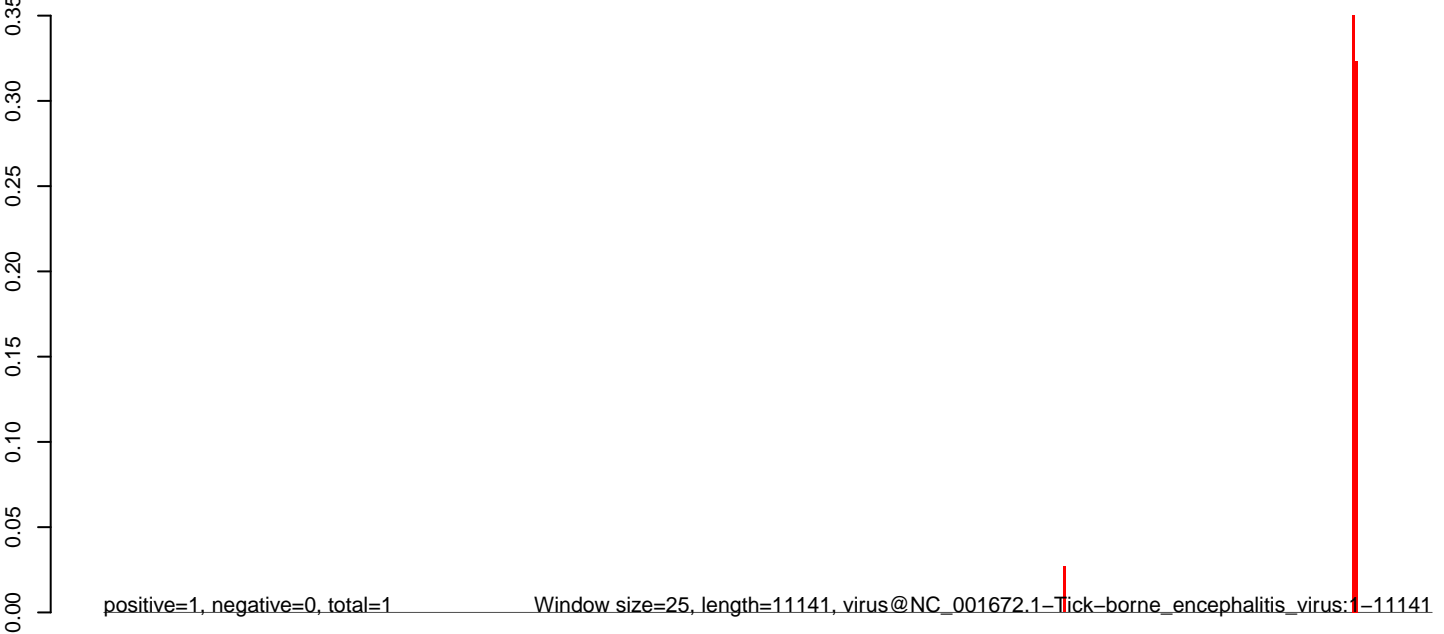
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



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

0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep

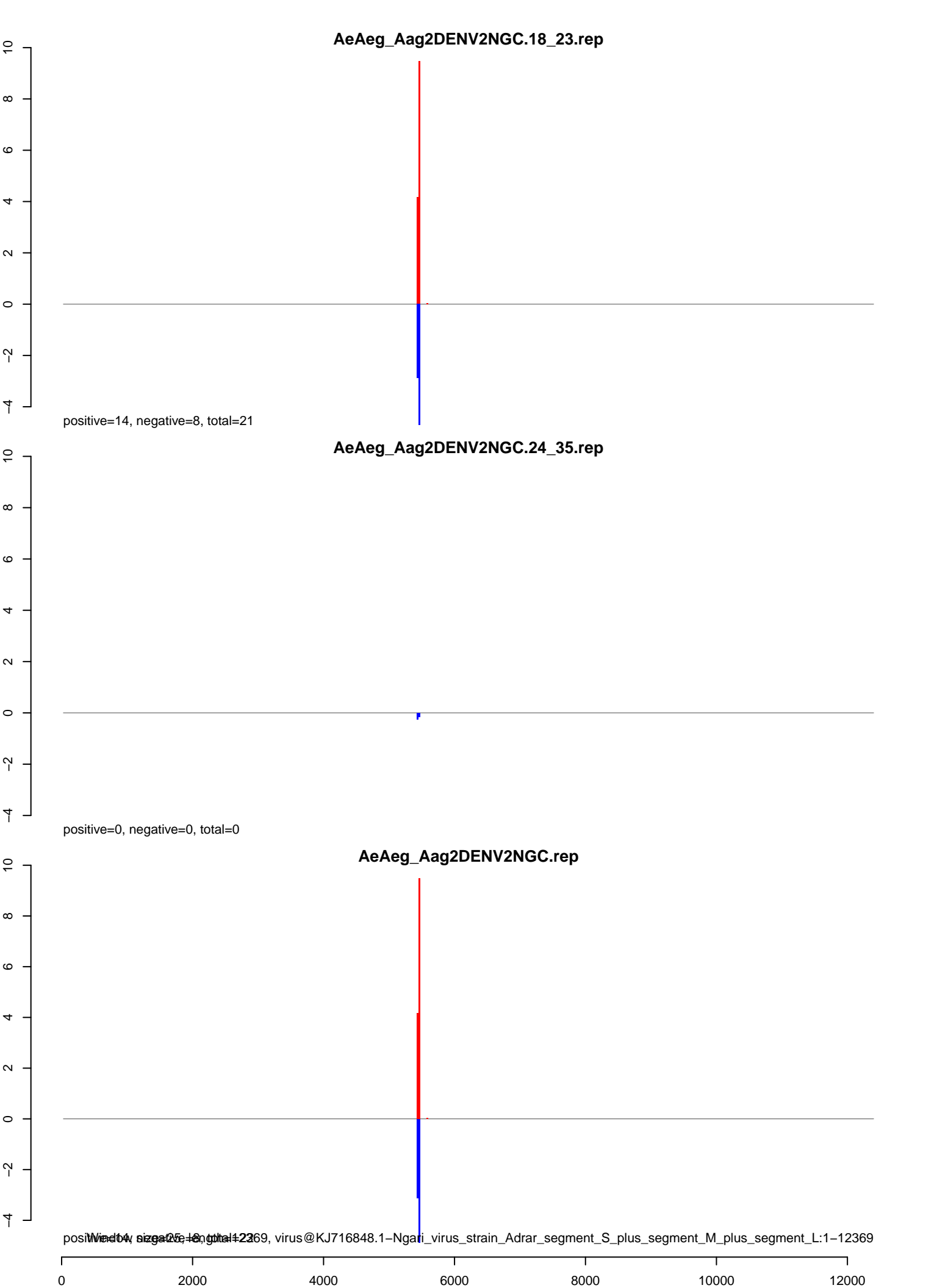
positive=14, negative=8, total=21

AeAeg_Aag2DENV2NGC.24_35.rep

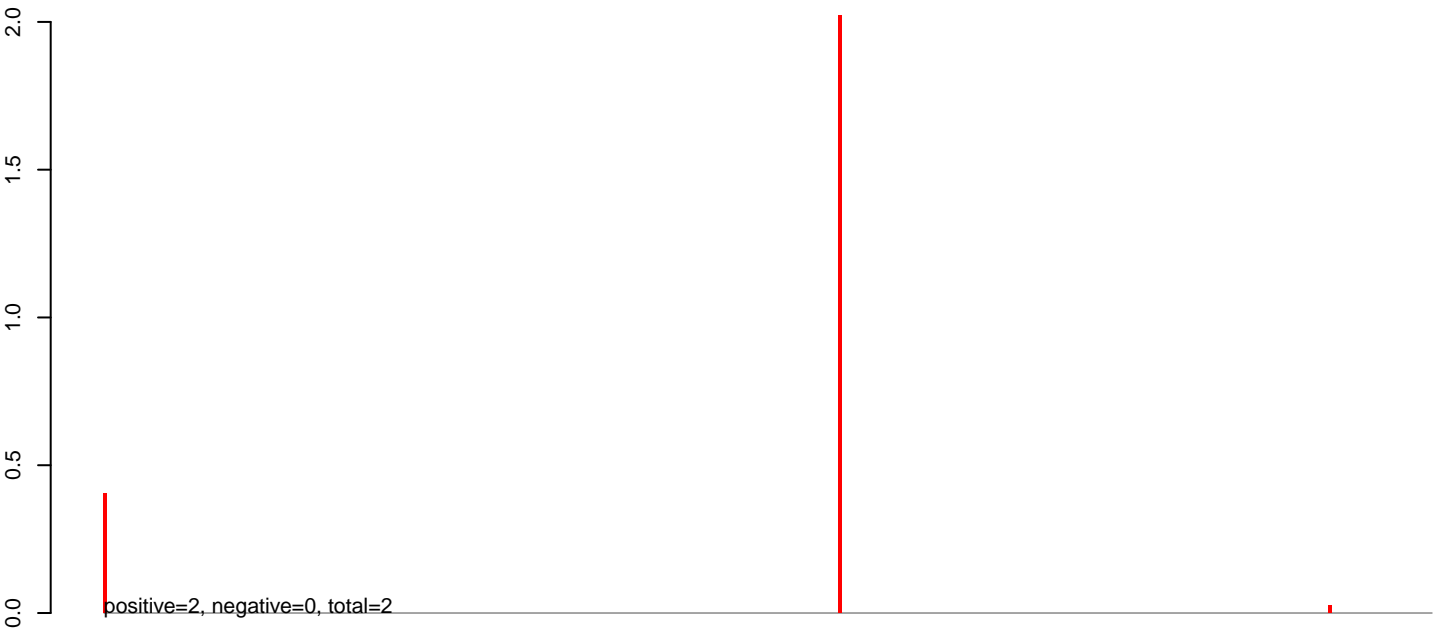
positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.rep

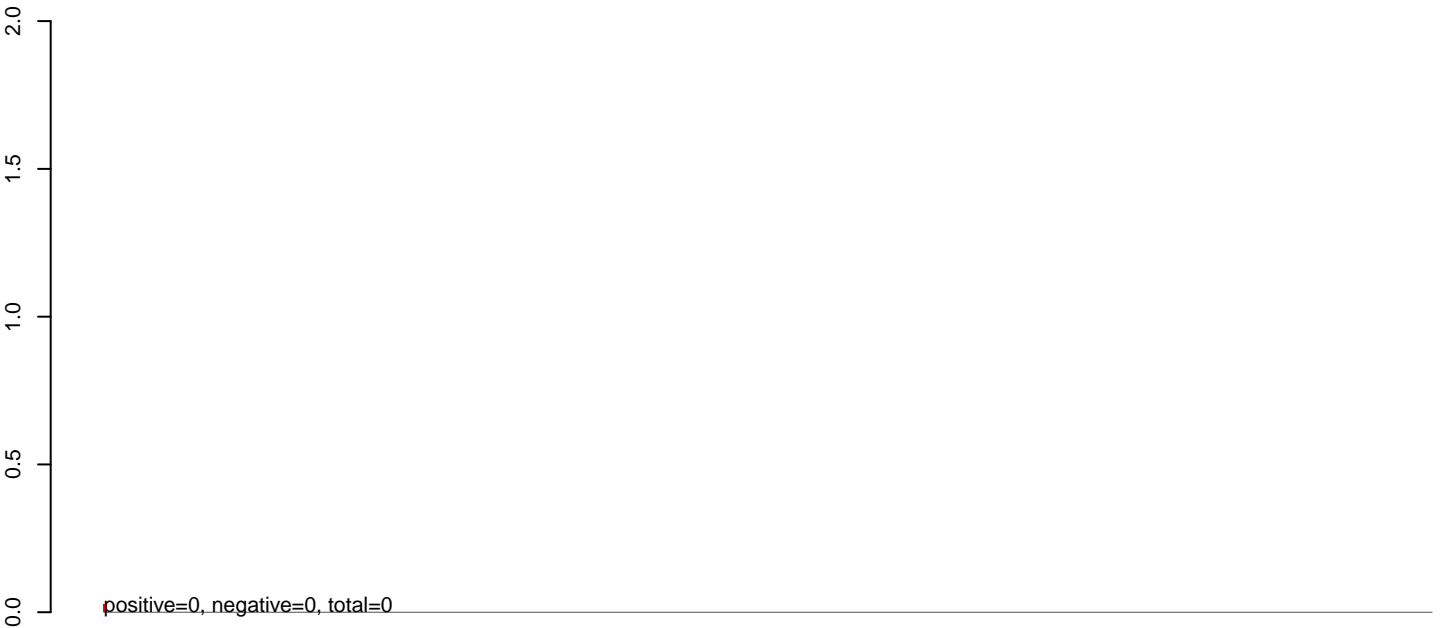
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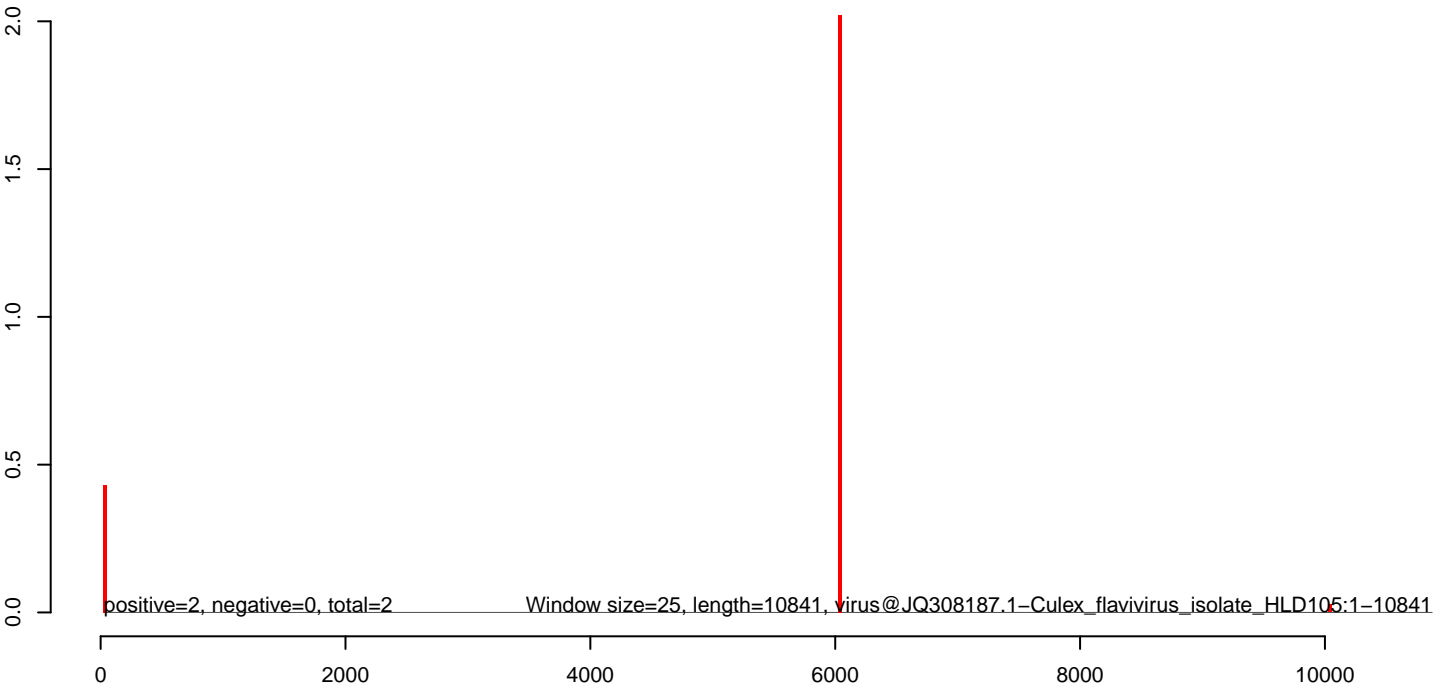
AeAeg_Aag2DENV2NGC.18_23.rep



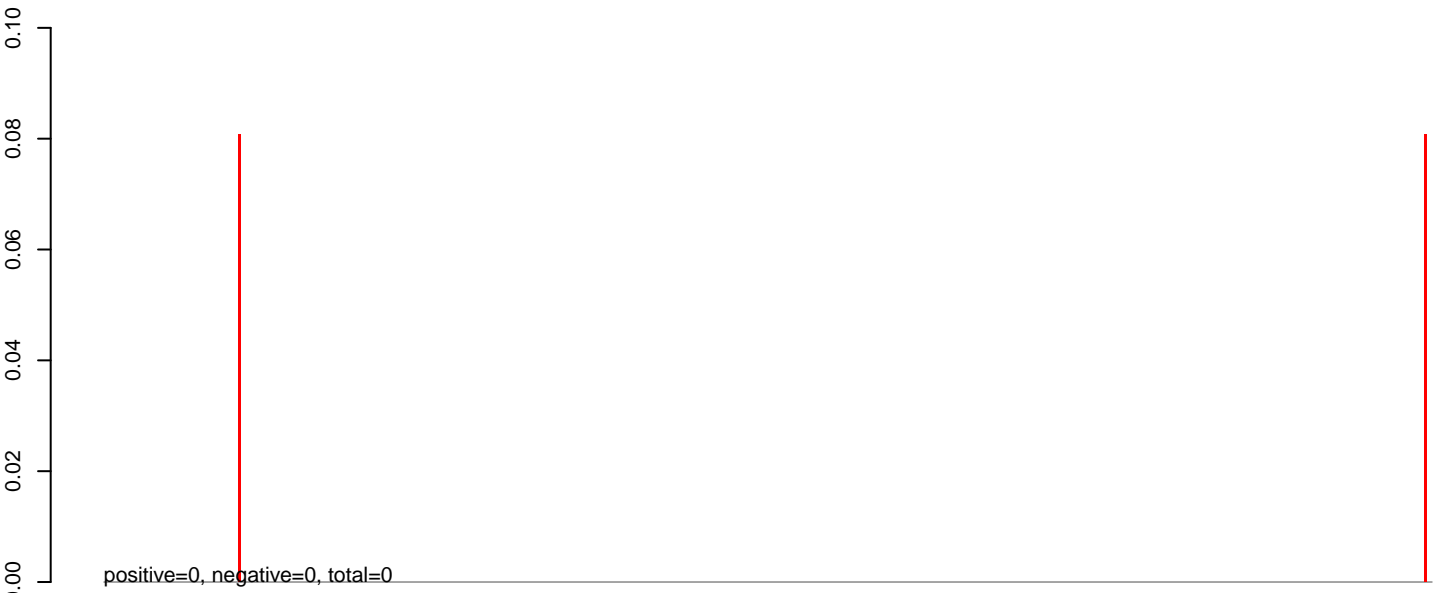
AeAeg_Aag2DENV2NGC.24_35.rep



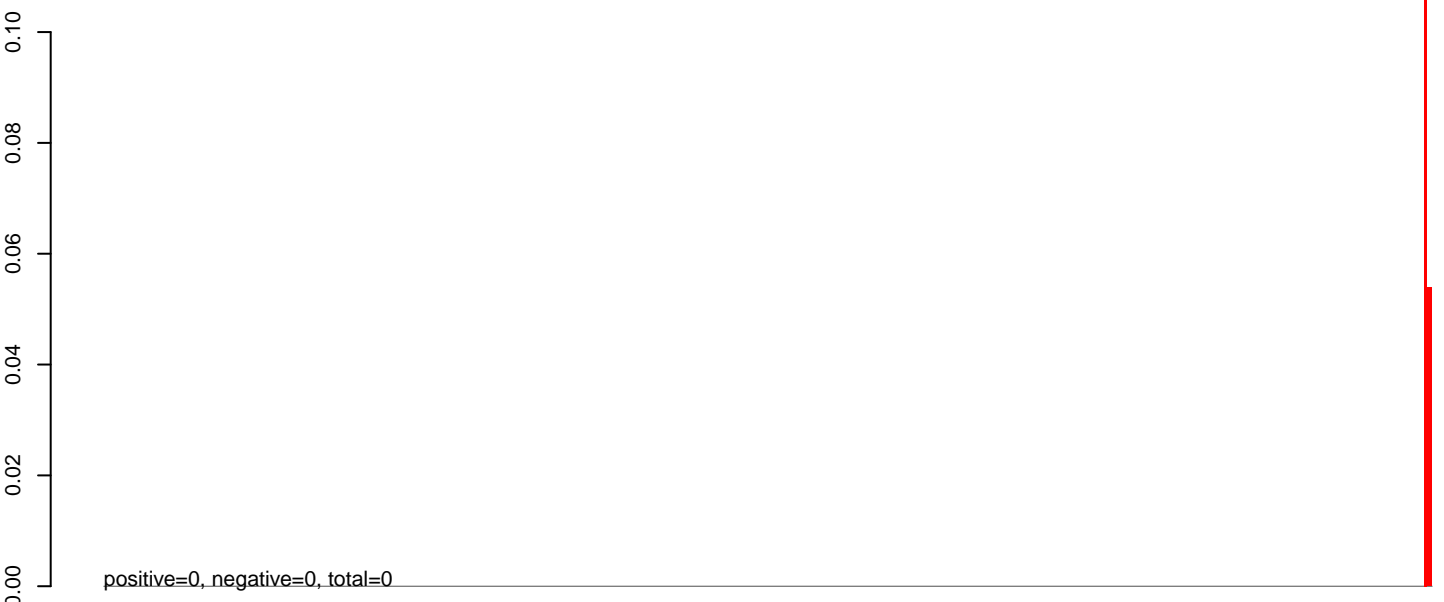
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



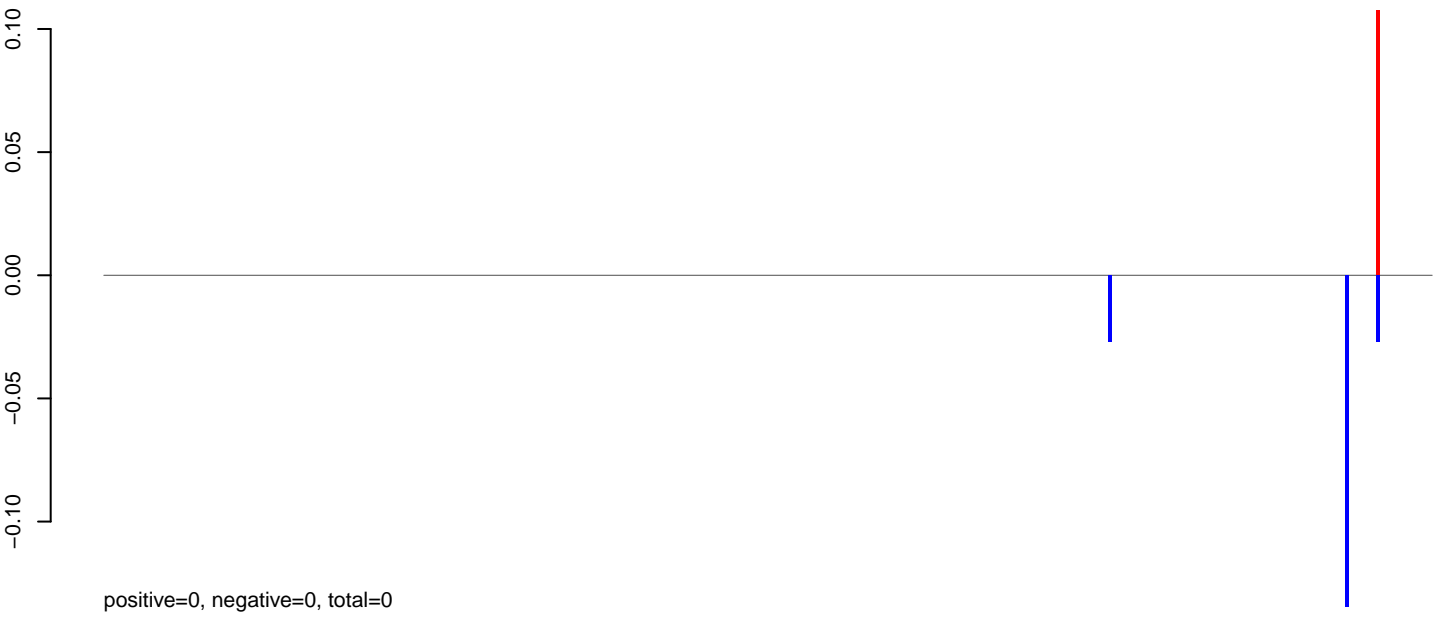
AeAeg_Aag2DENV2NGC.24_35.rep



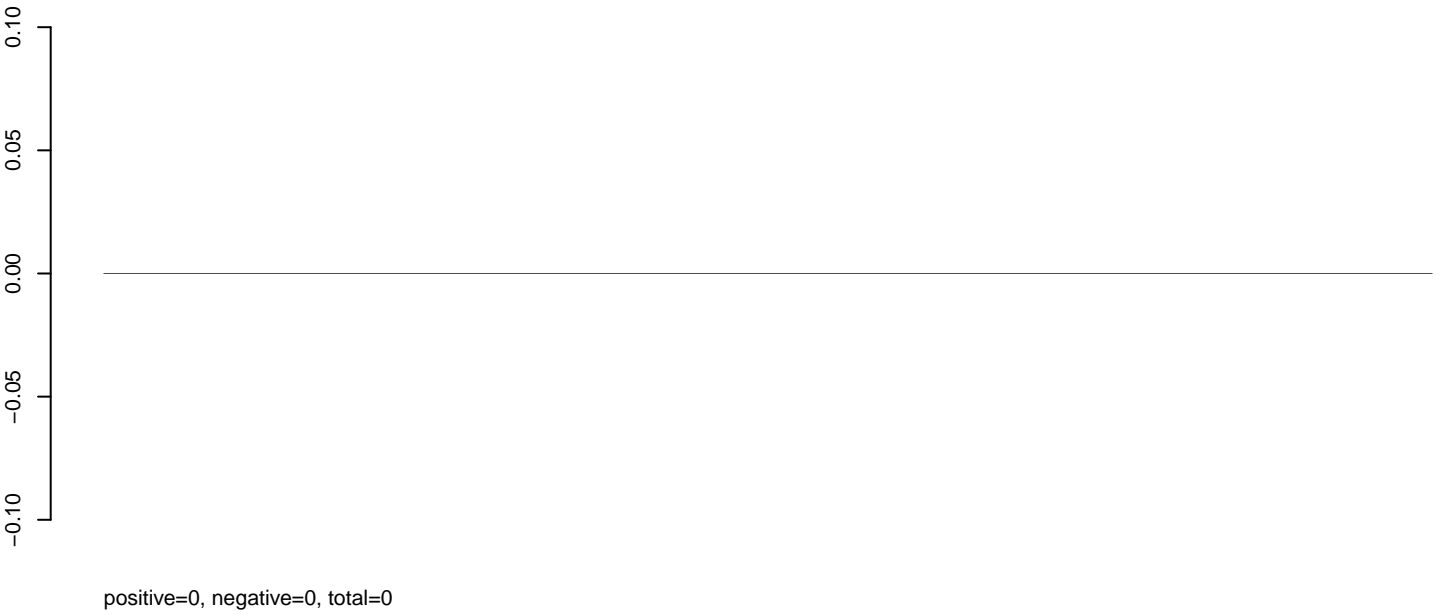
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



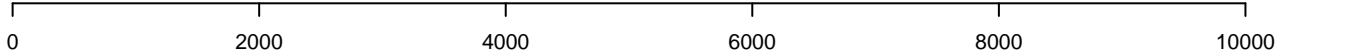
AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



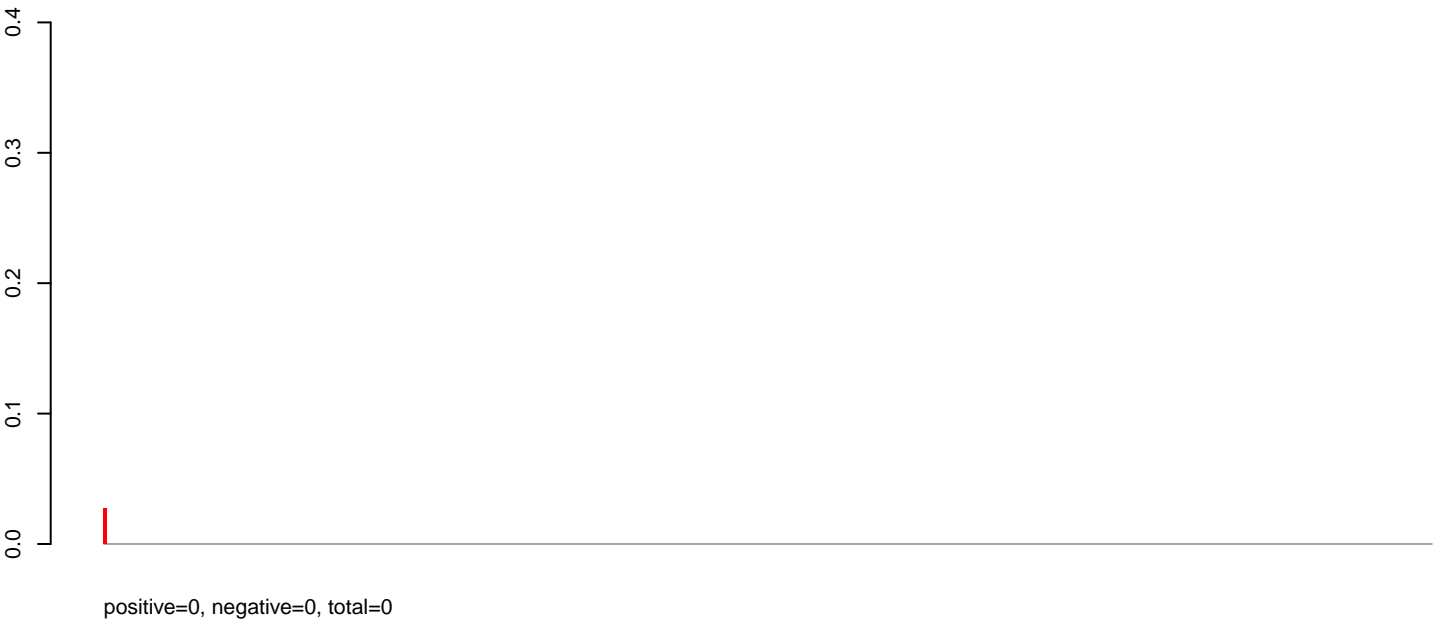
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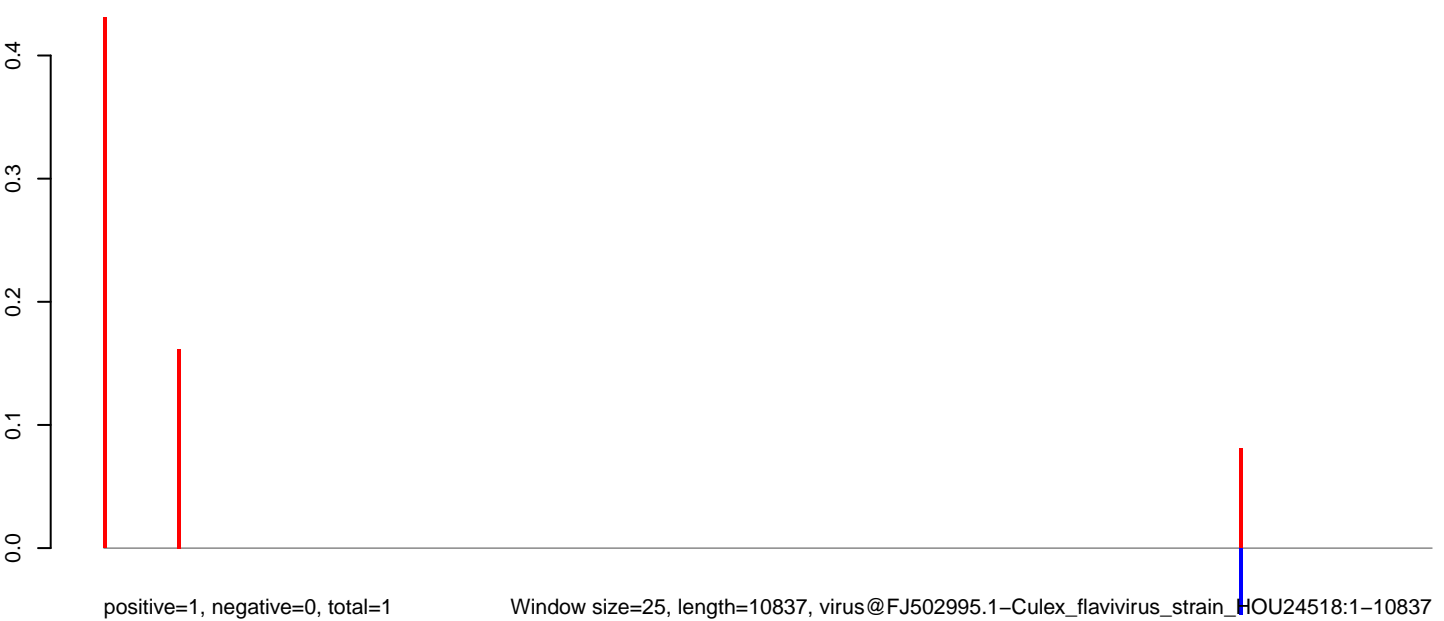
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



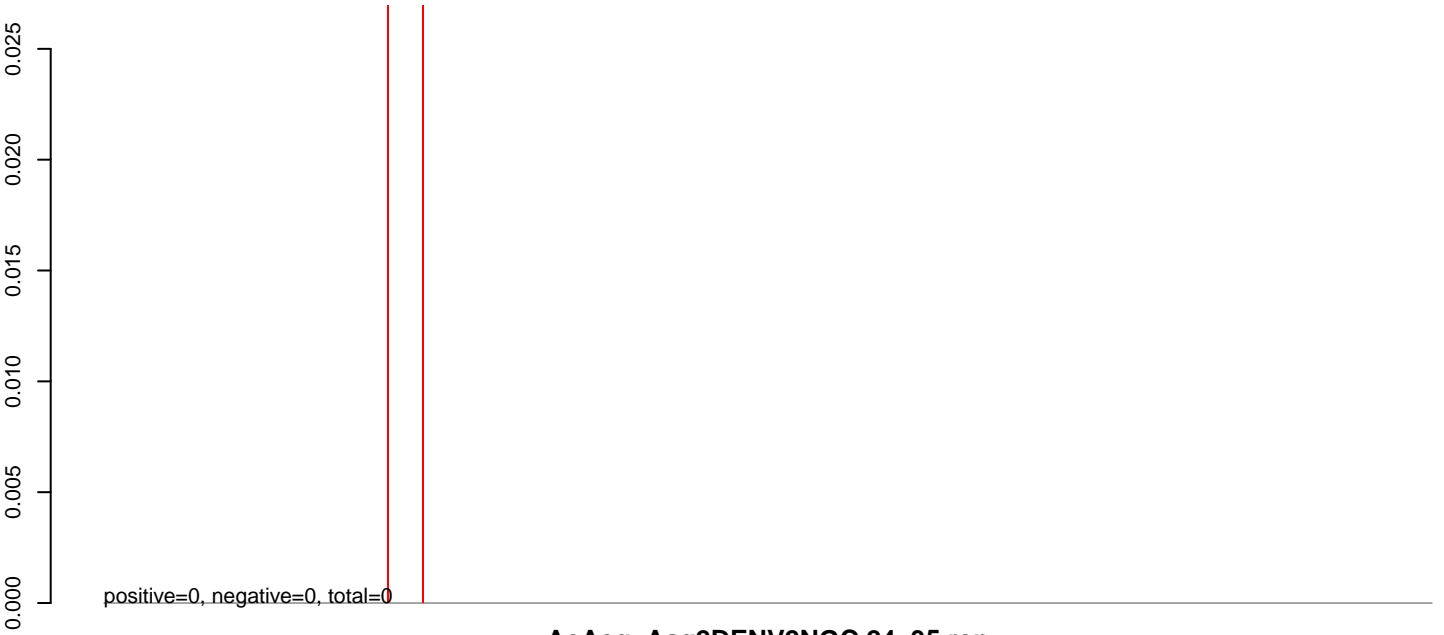
AeAeg_Aag2DENV2NGC.rep



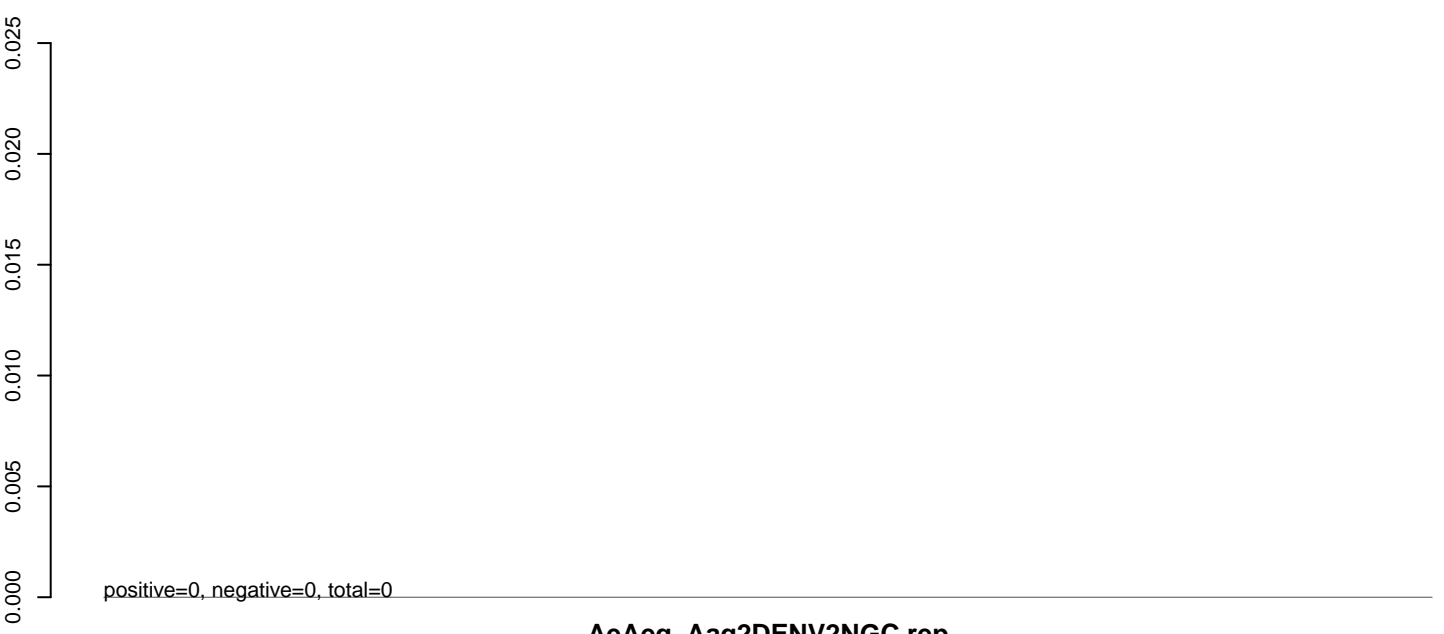
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0 2000 4000 6000 8000 10000

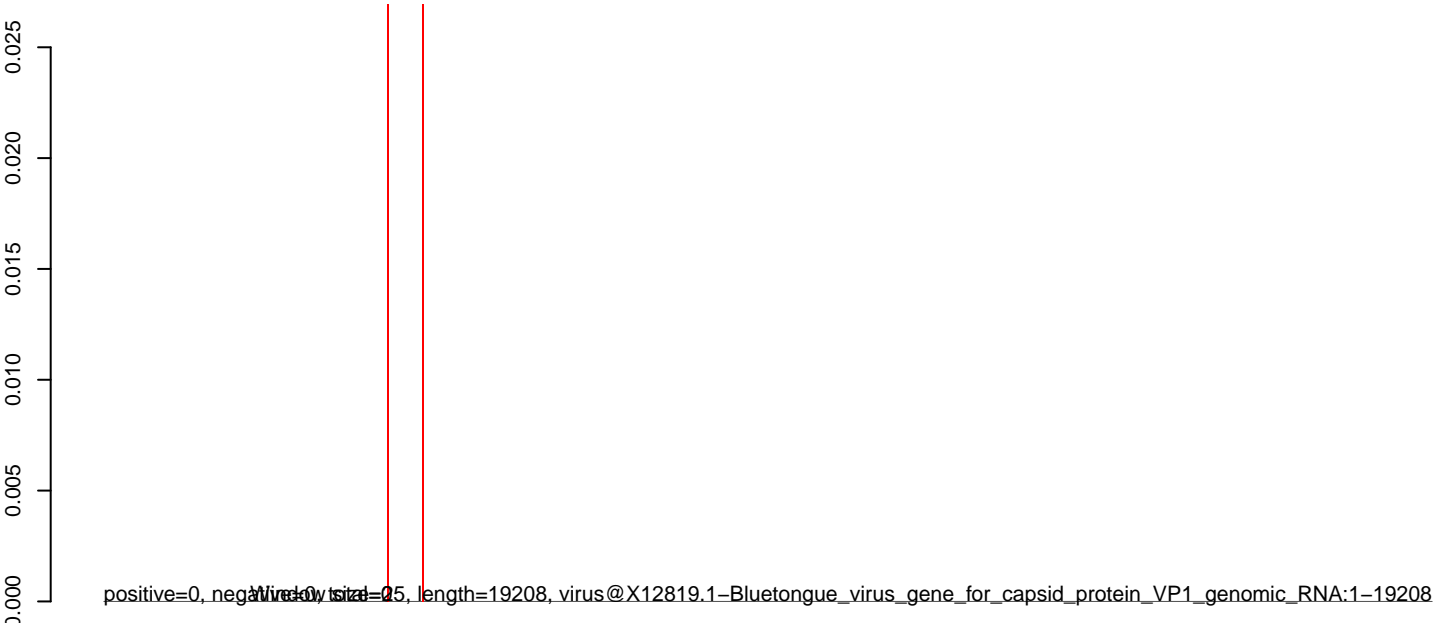
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

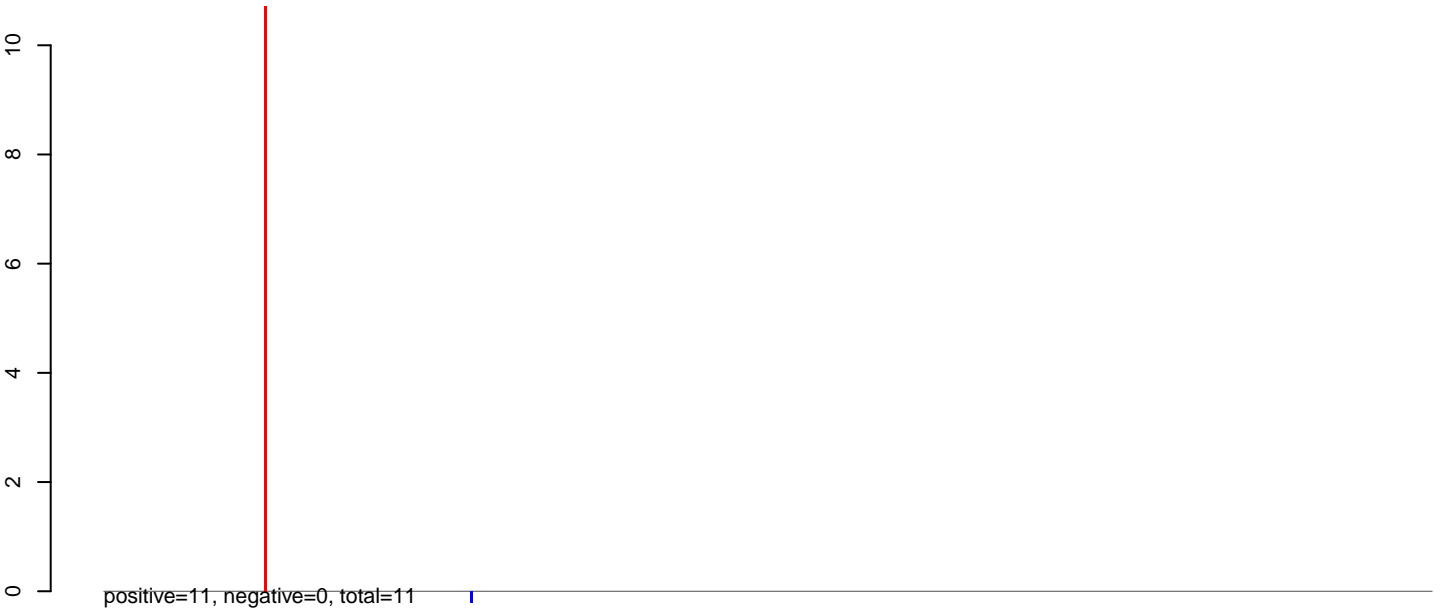


AeAeg_Aag2DENV2NGC.rep

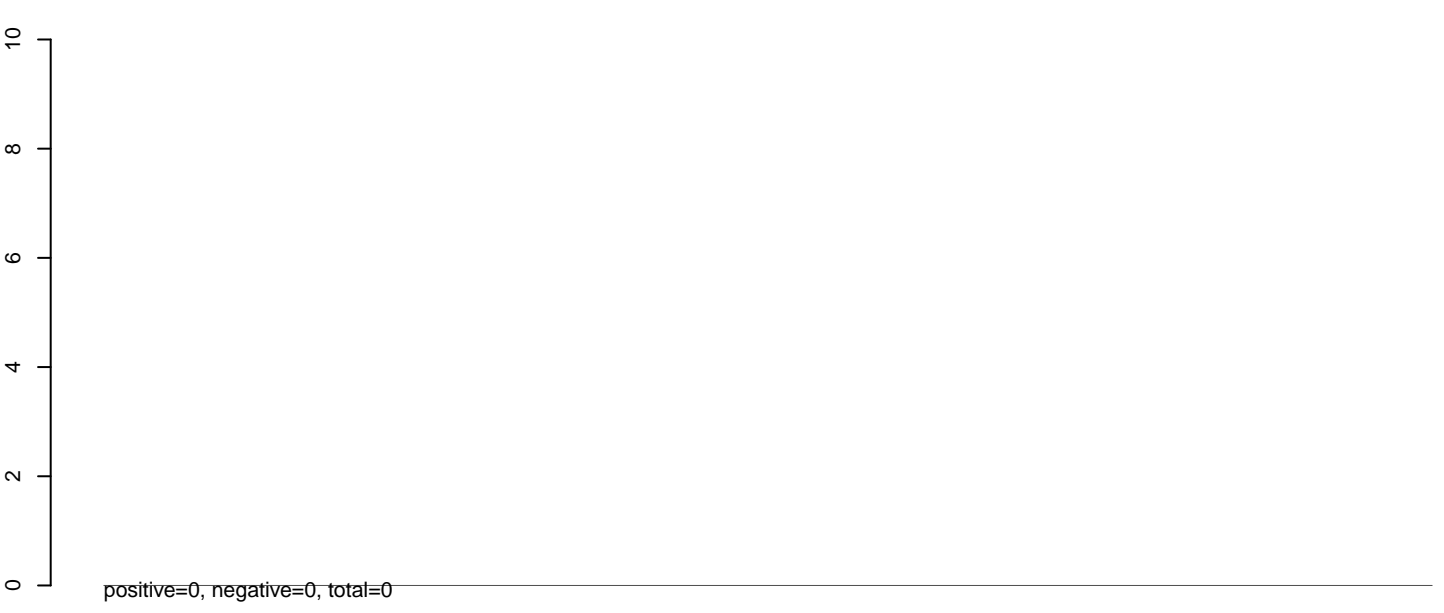


0 5000 10000 15000 20000

AeAeg_Aag2DENV2NGC.18_23.rep



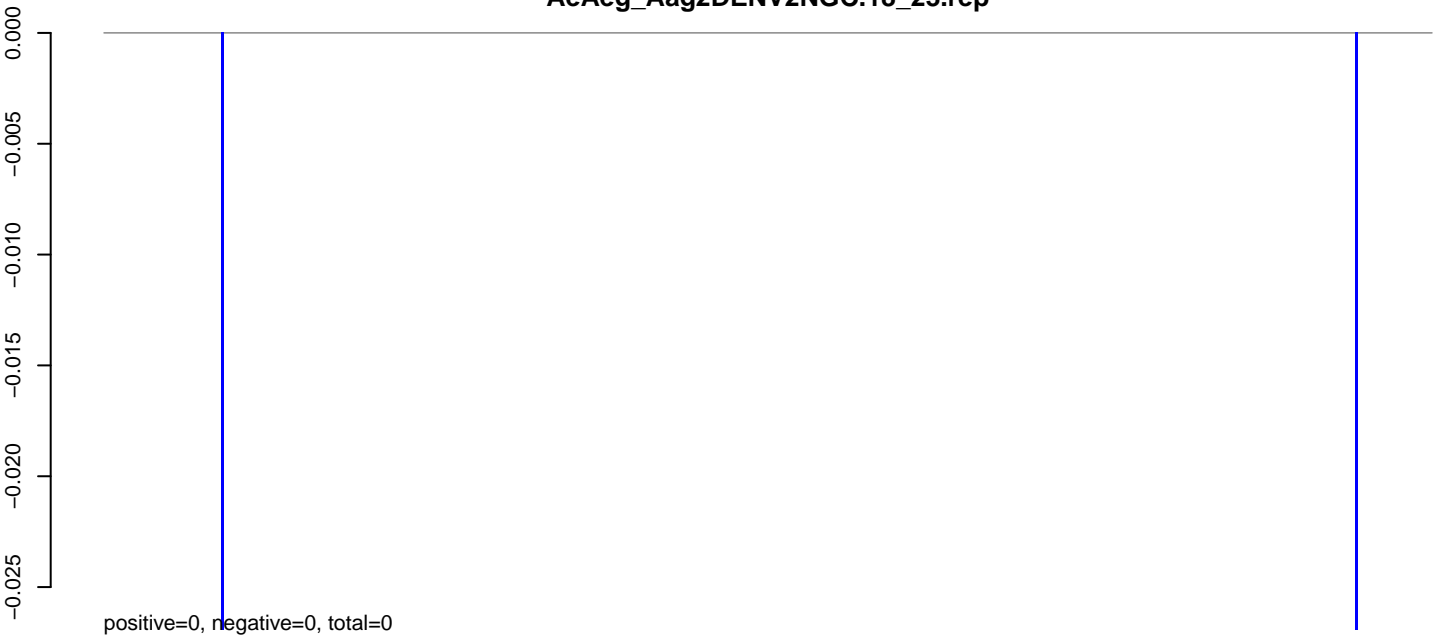
AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



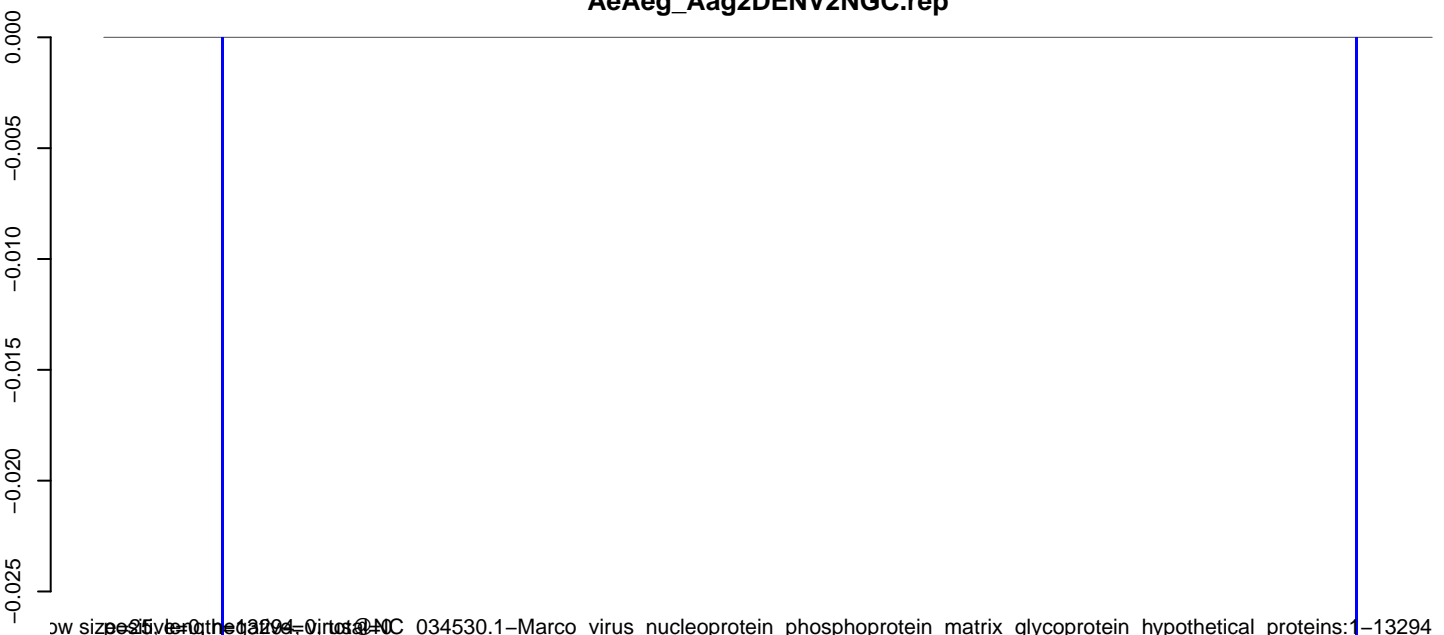
AeAeg_Aag2DENV2NGC.18_23.rep



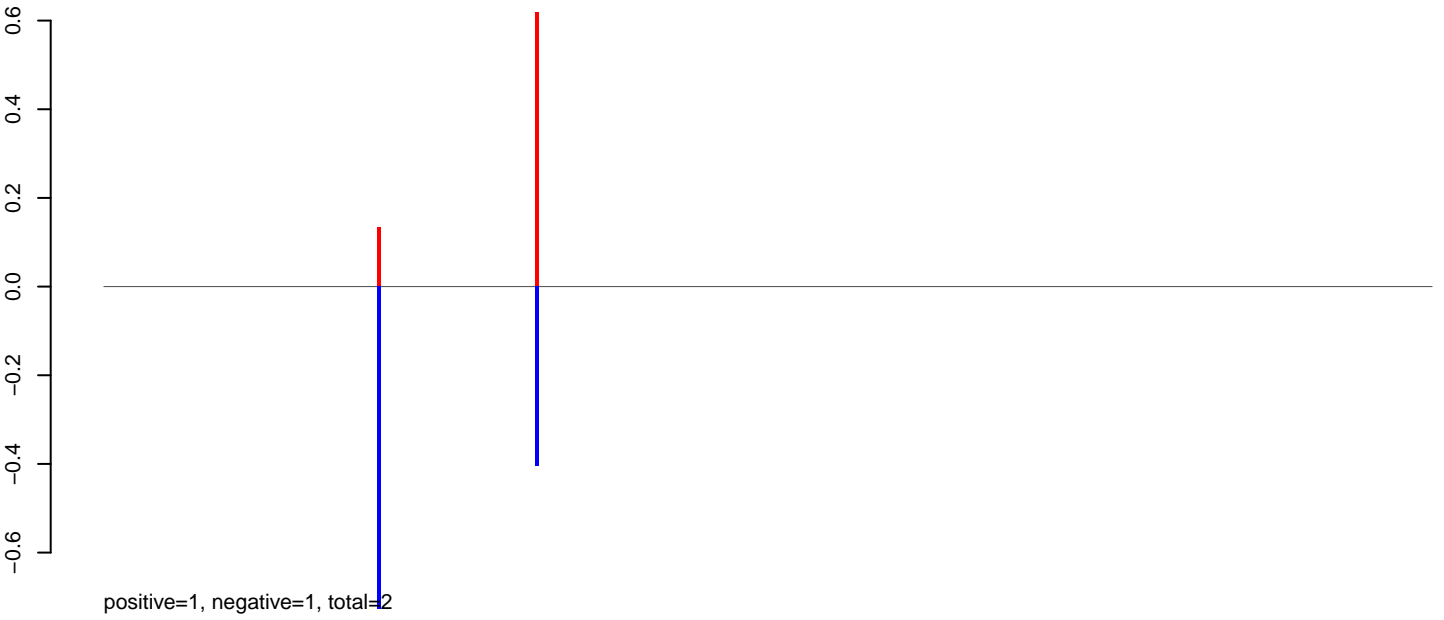
AeAeg_Aag2DENV2NGC.24_35.rep



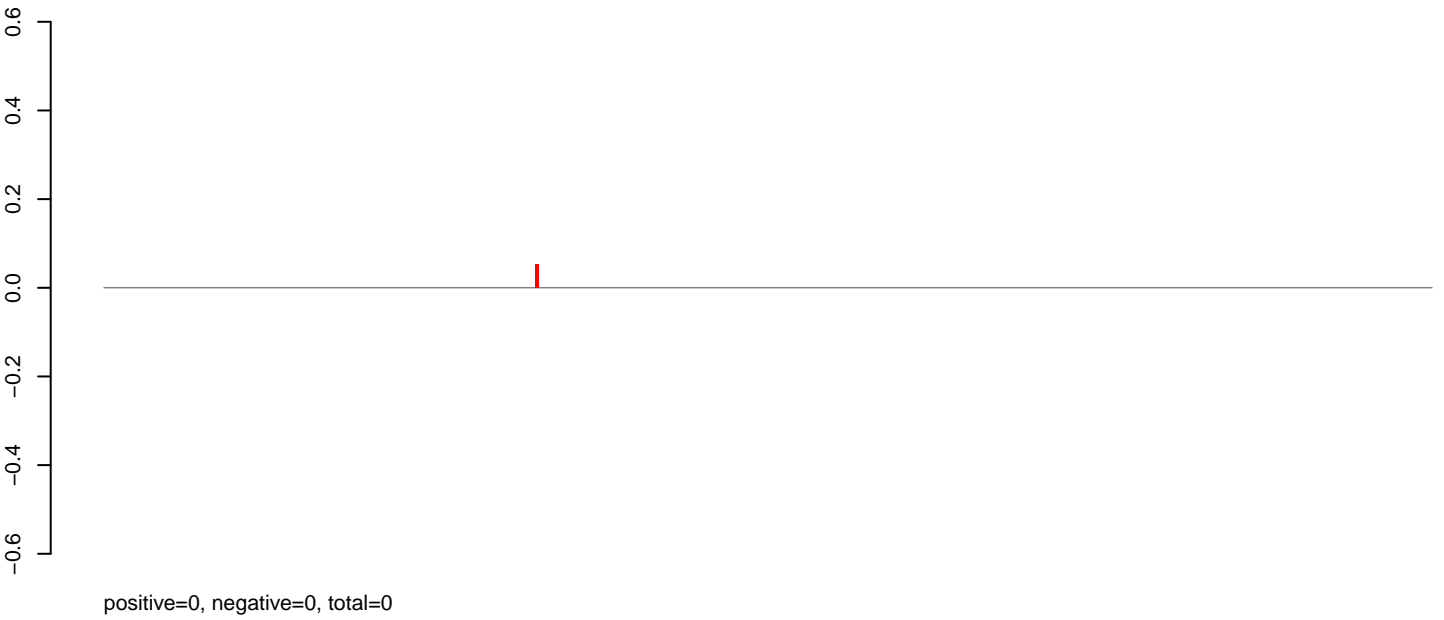
AeAeg_Aag2DENV2NGC.rep



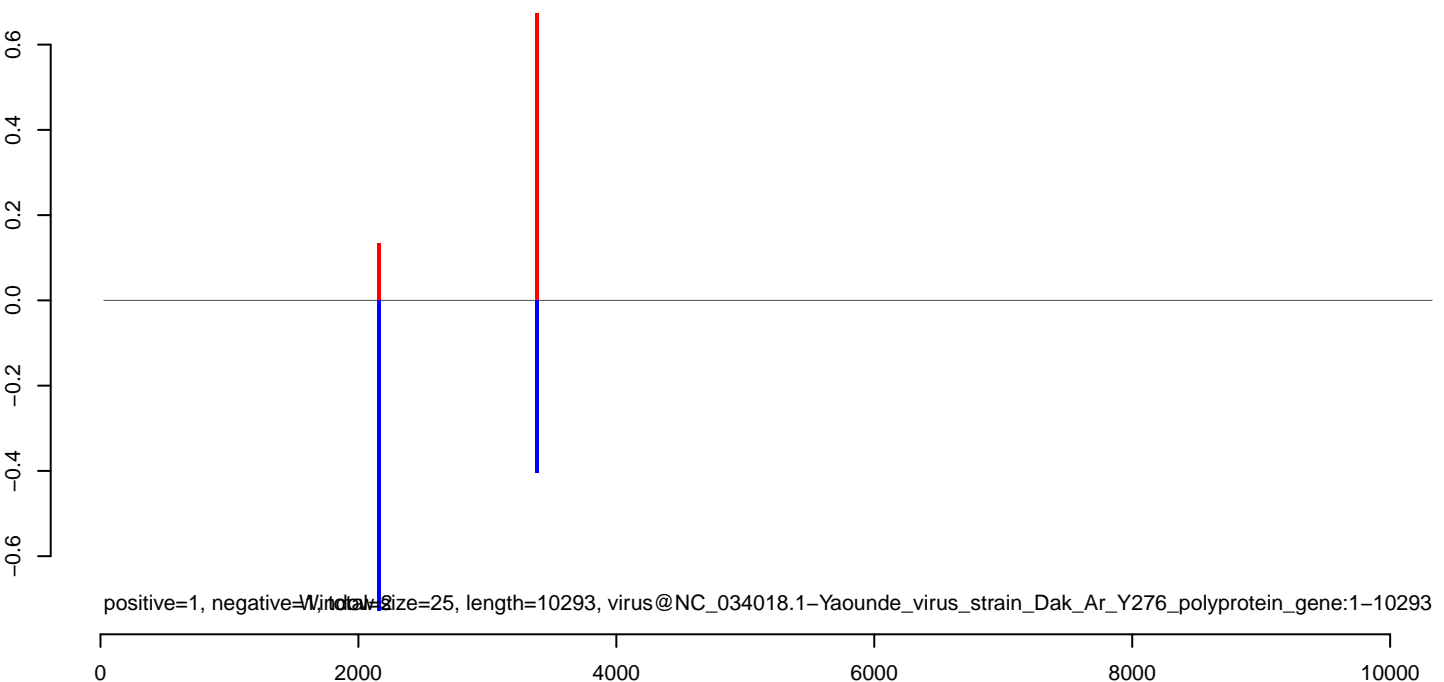
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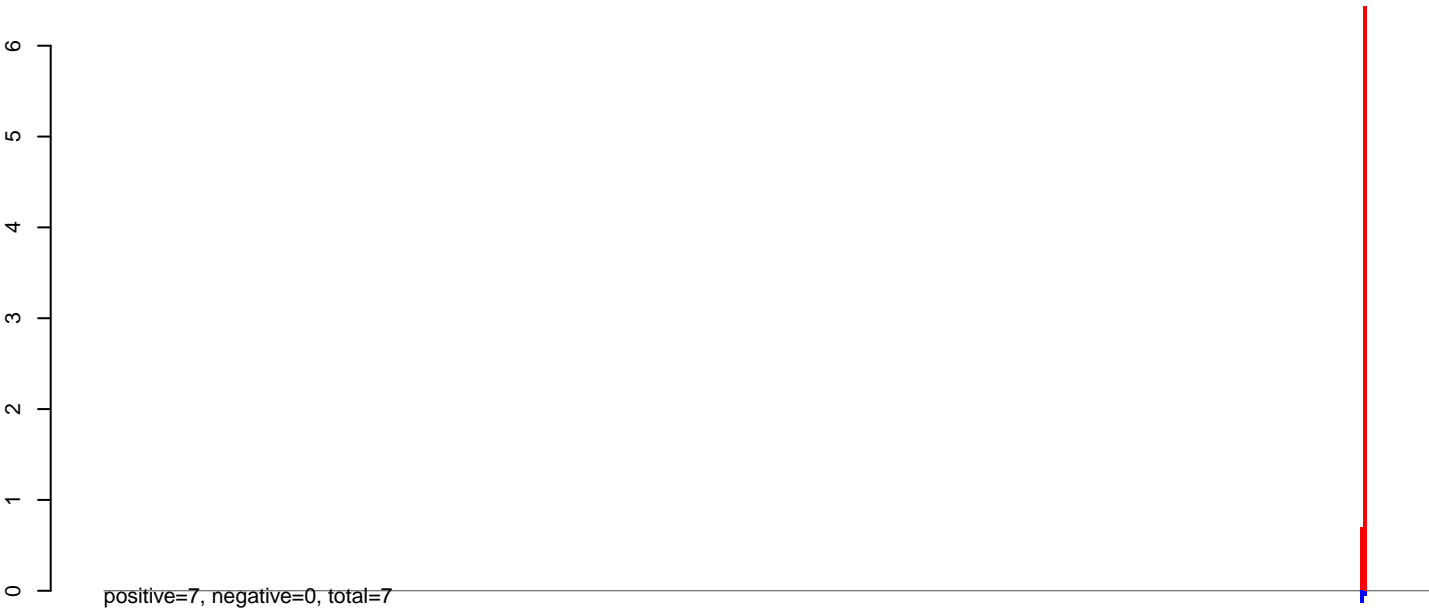
AeAeg_Aag2DENV2NGC.24_35.rep



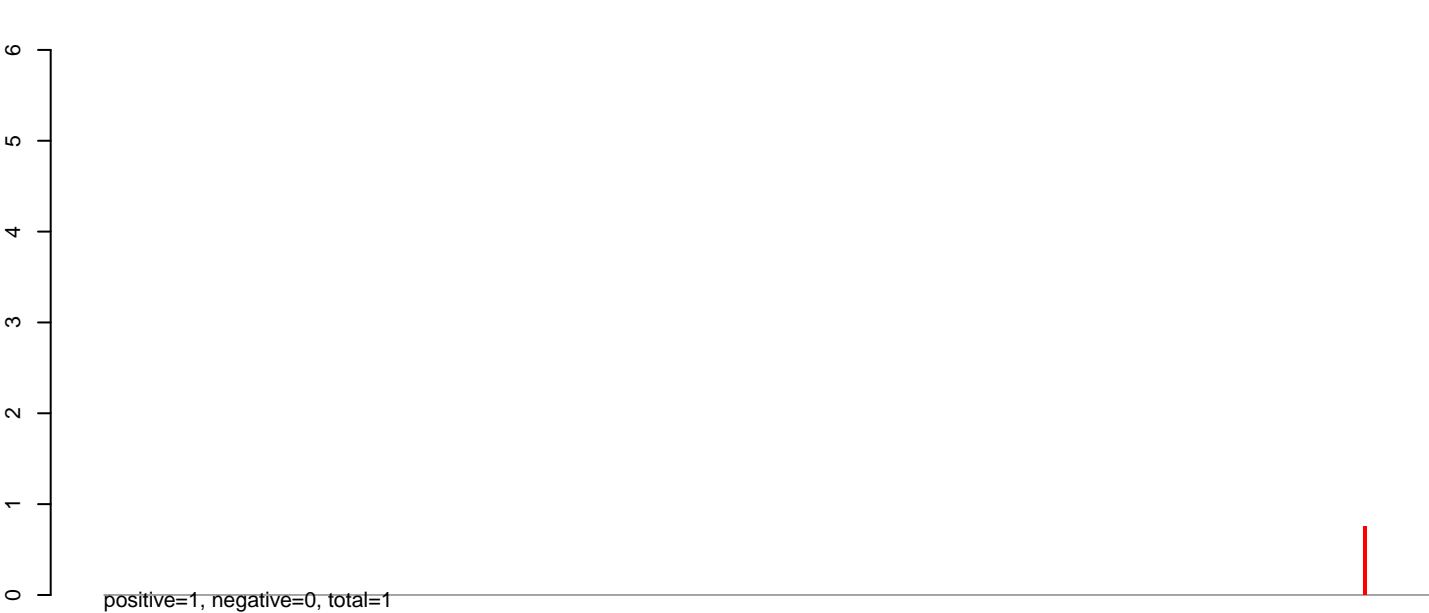
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



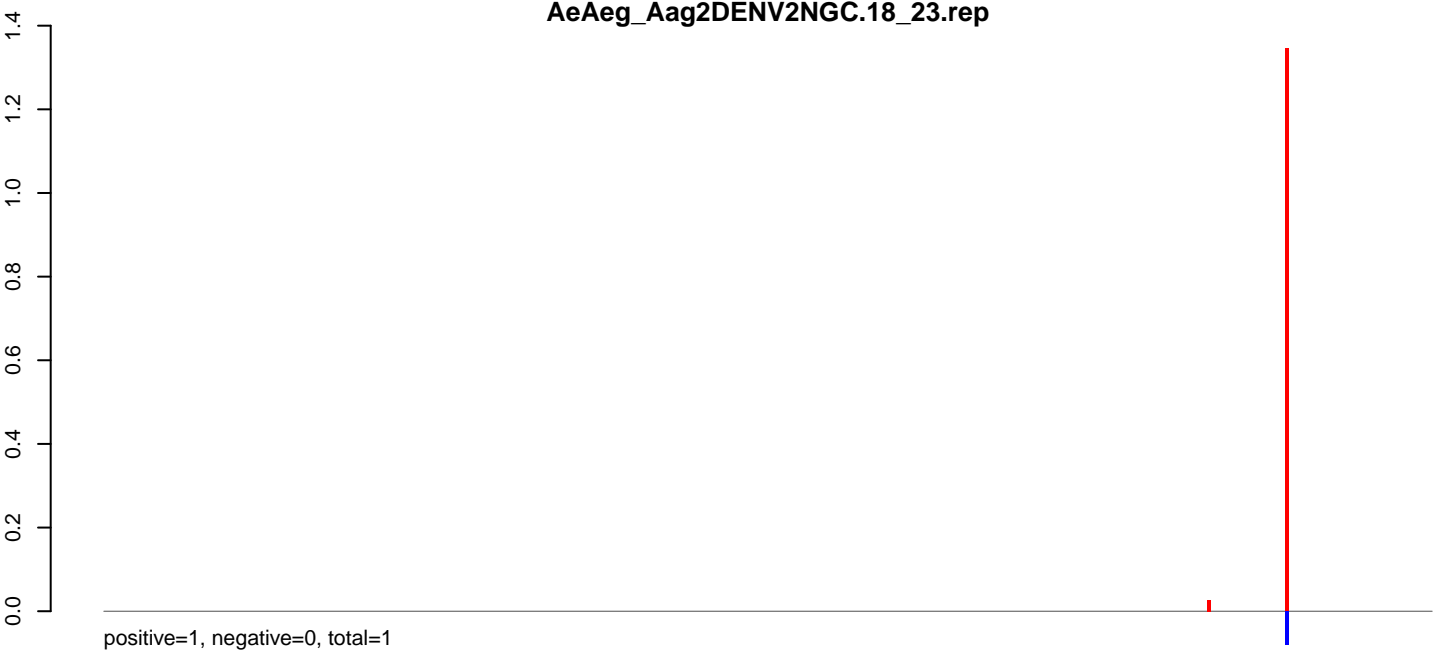
AeAeg_Aag2DENV2NGC.24_35.rep



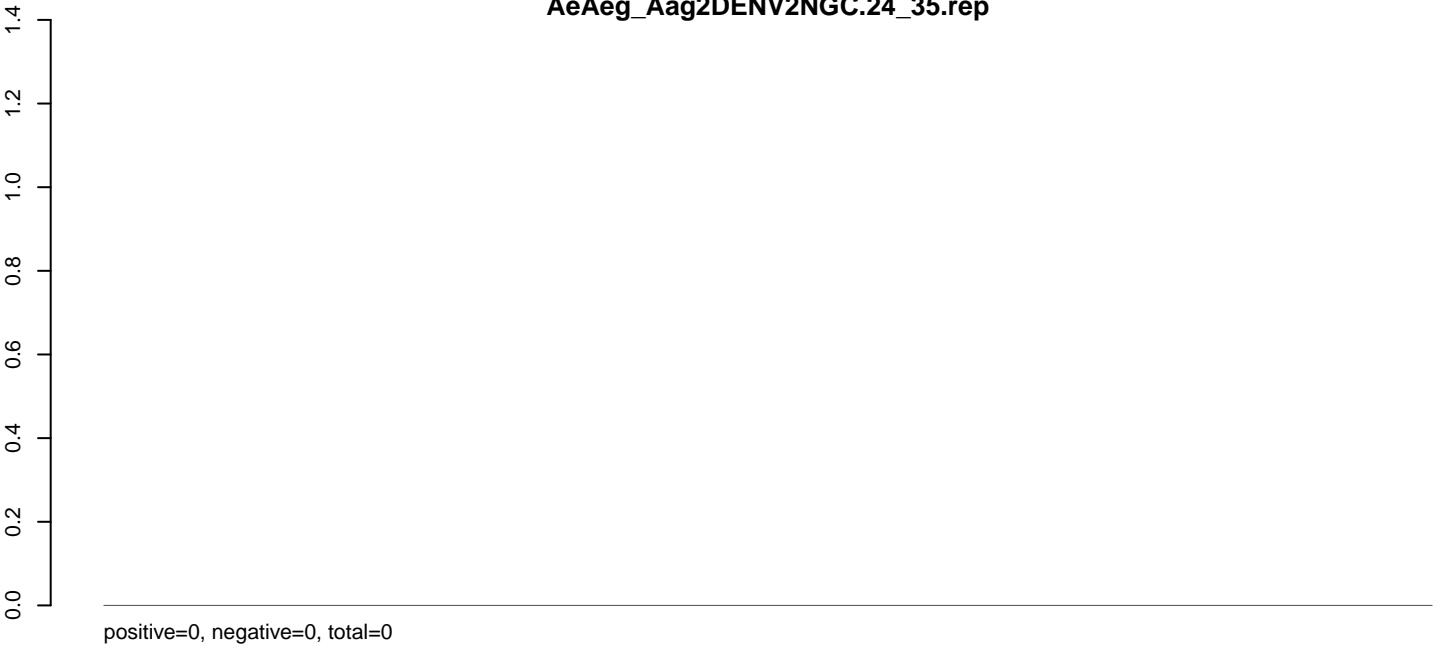
AeAeg_Aag2DENV2NGC.rep



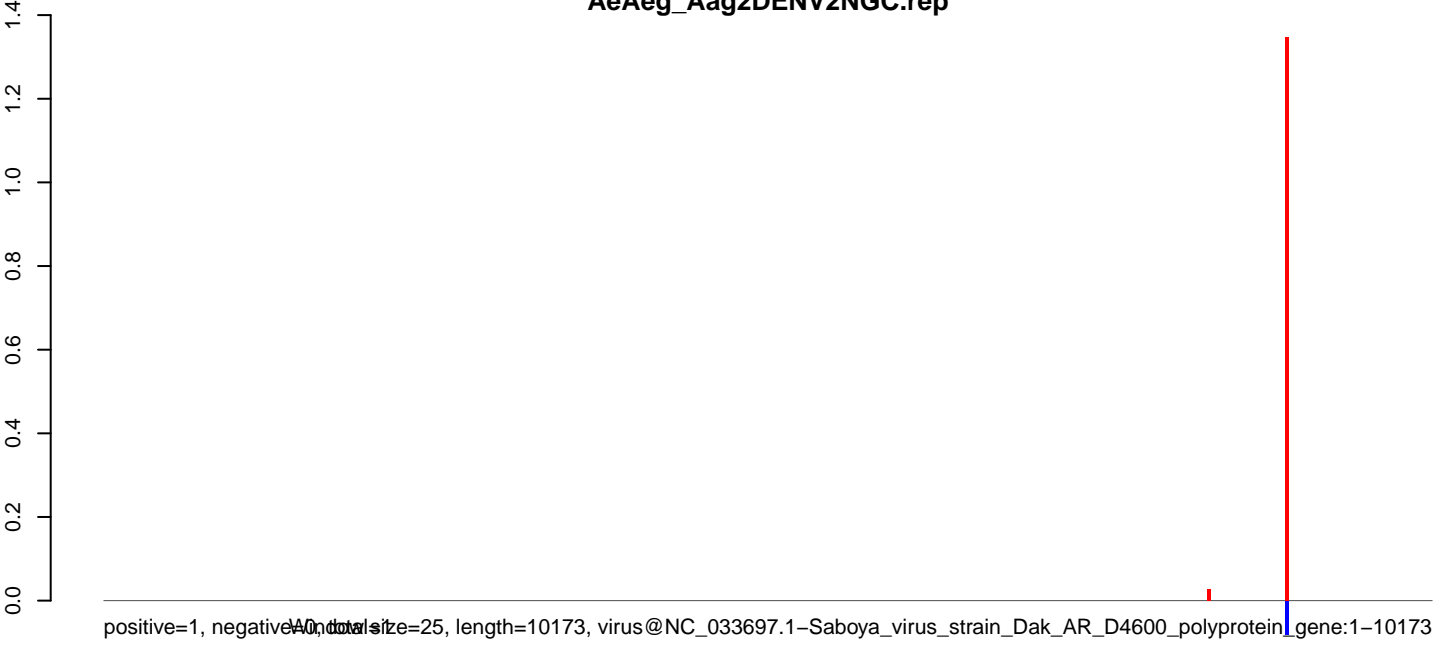
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



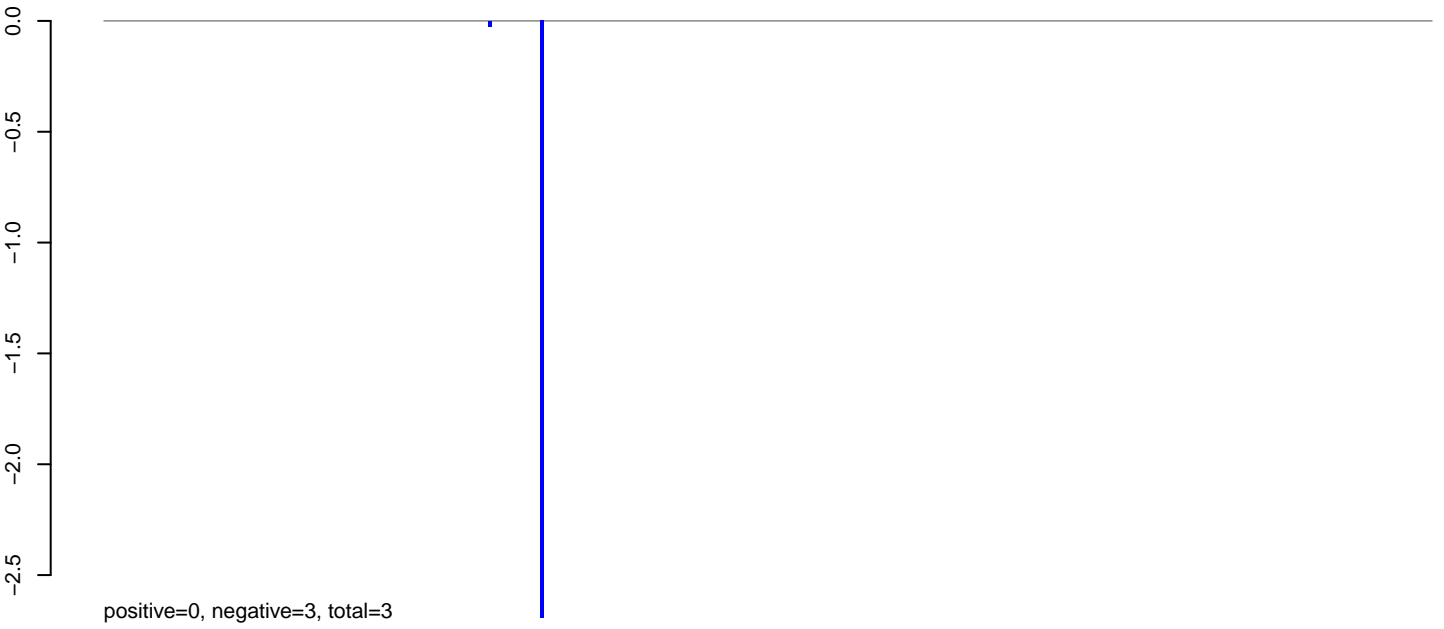
AeAeg_Aag2DENV2NGC.rep



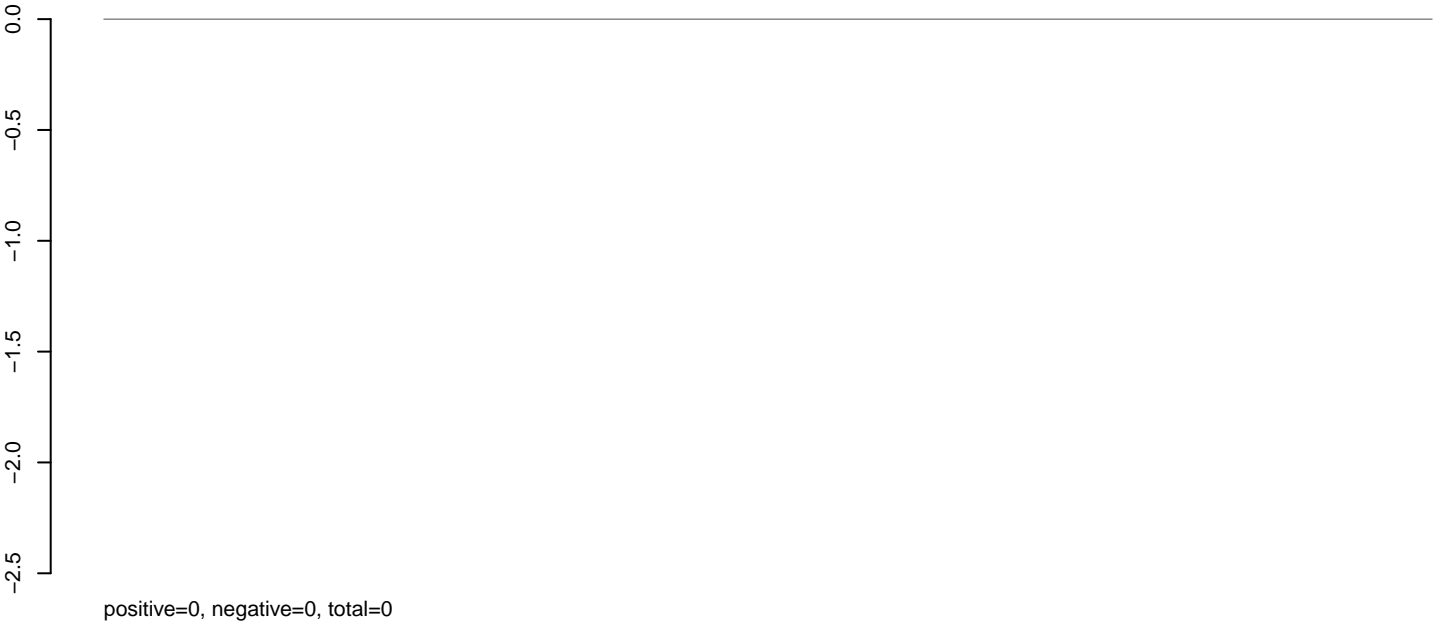
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0 2000 4000 6000 8000 10000

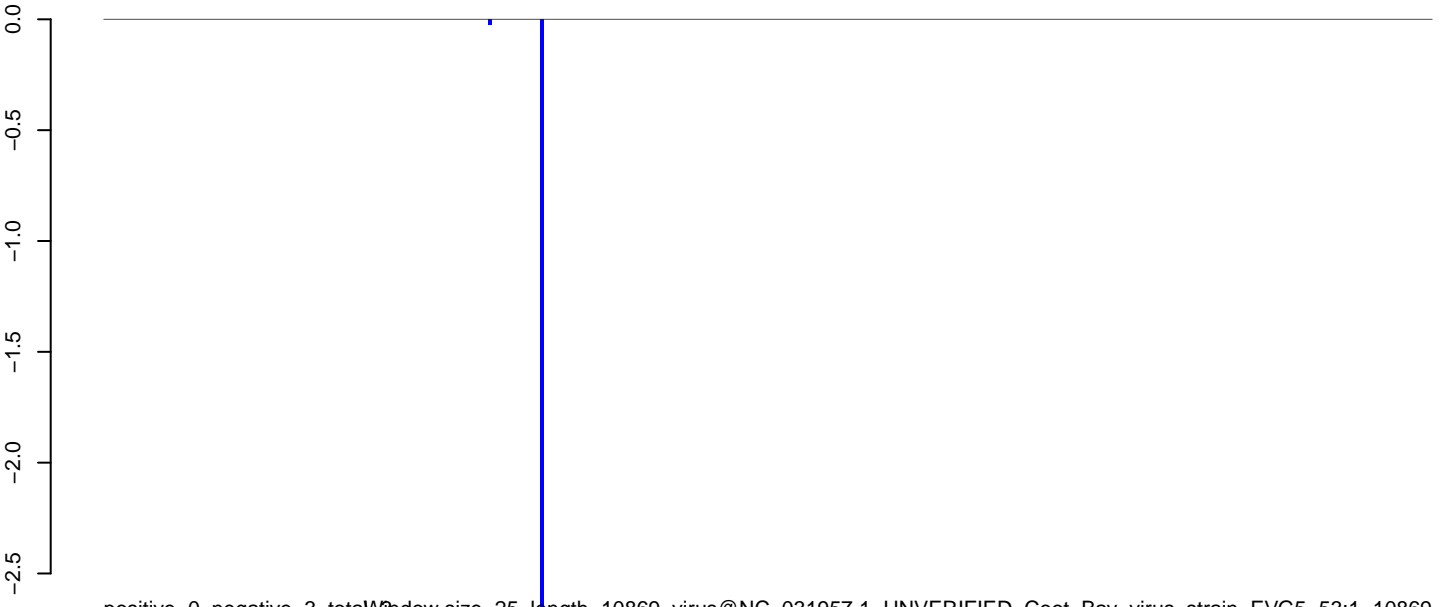
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AeAeg_Aag2DENV2NGC.24_35.rep

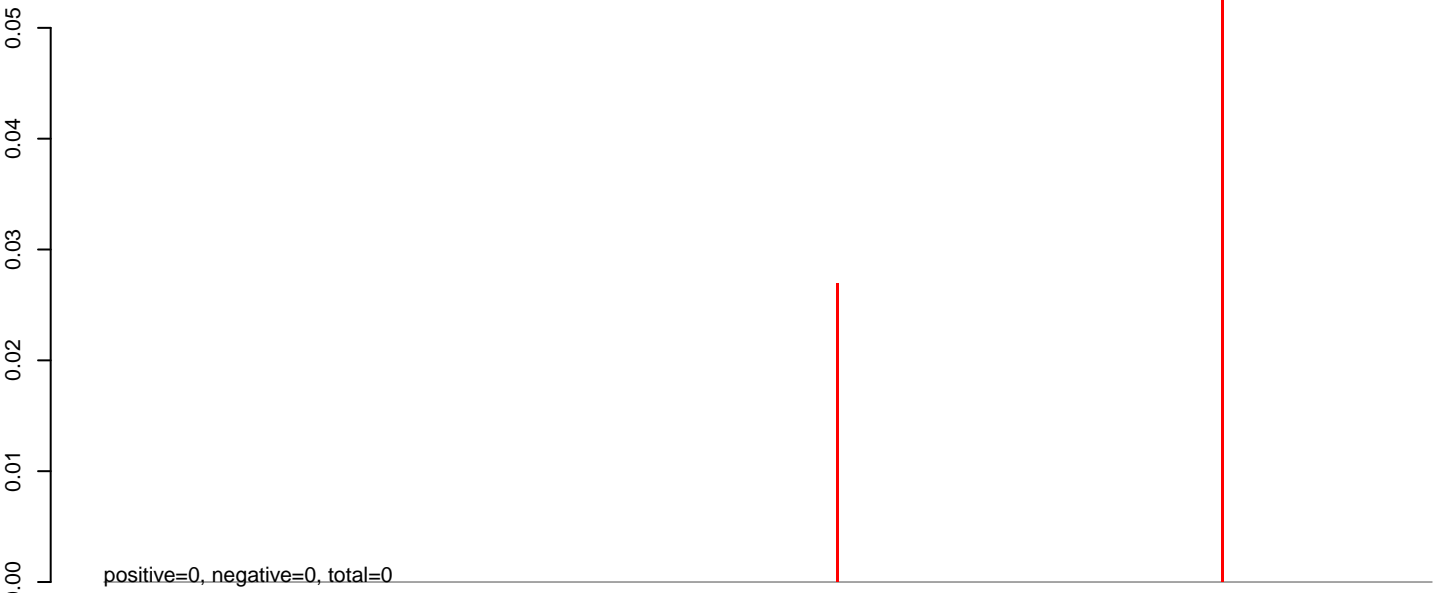


AeAeg_Aag2DENV2NGC.rep

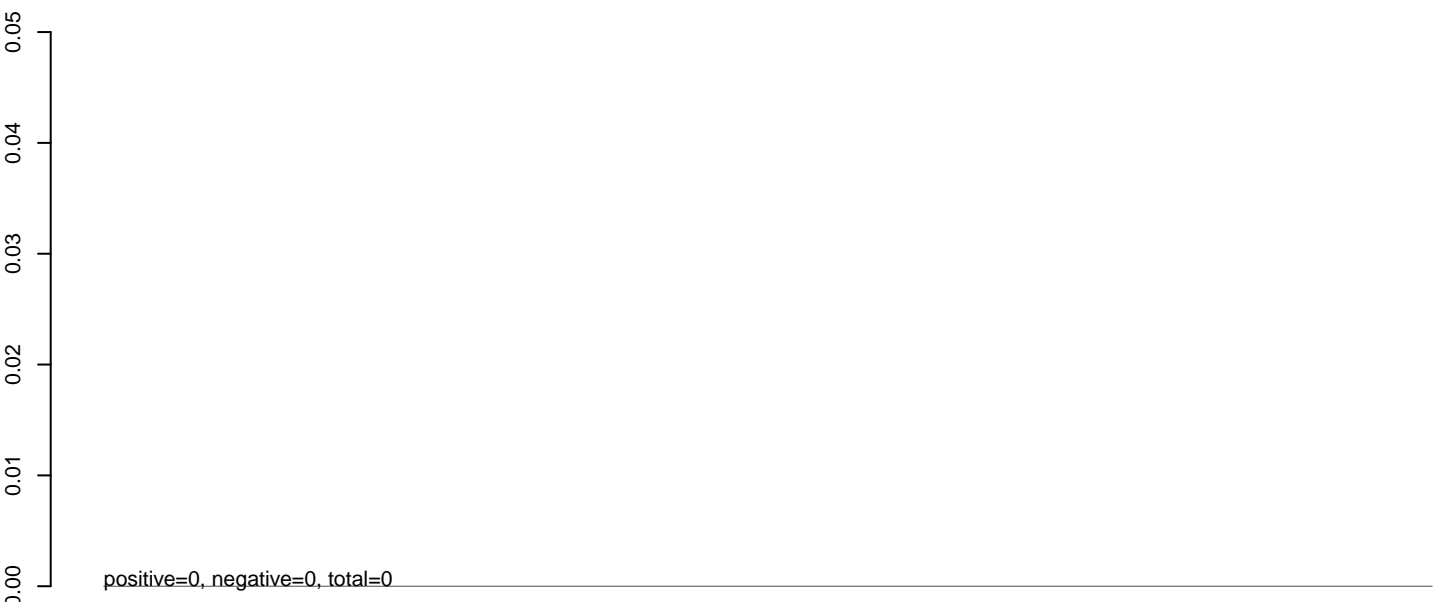


Window size=25, length=10869, virus@NC_031957.1-UNVERIFIED_Coot_Bay_virus_strain_EVG5-53:1-10869

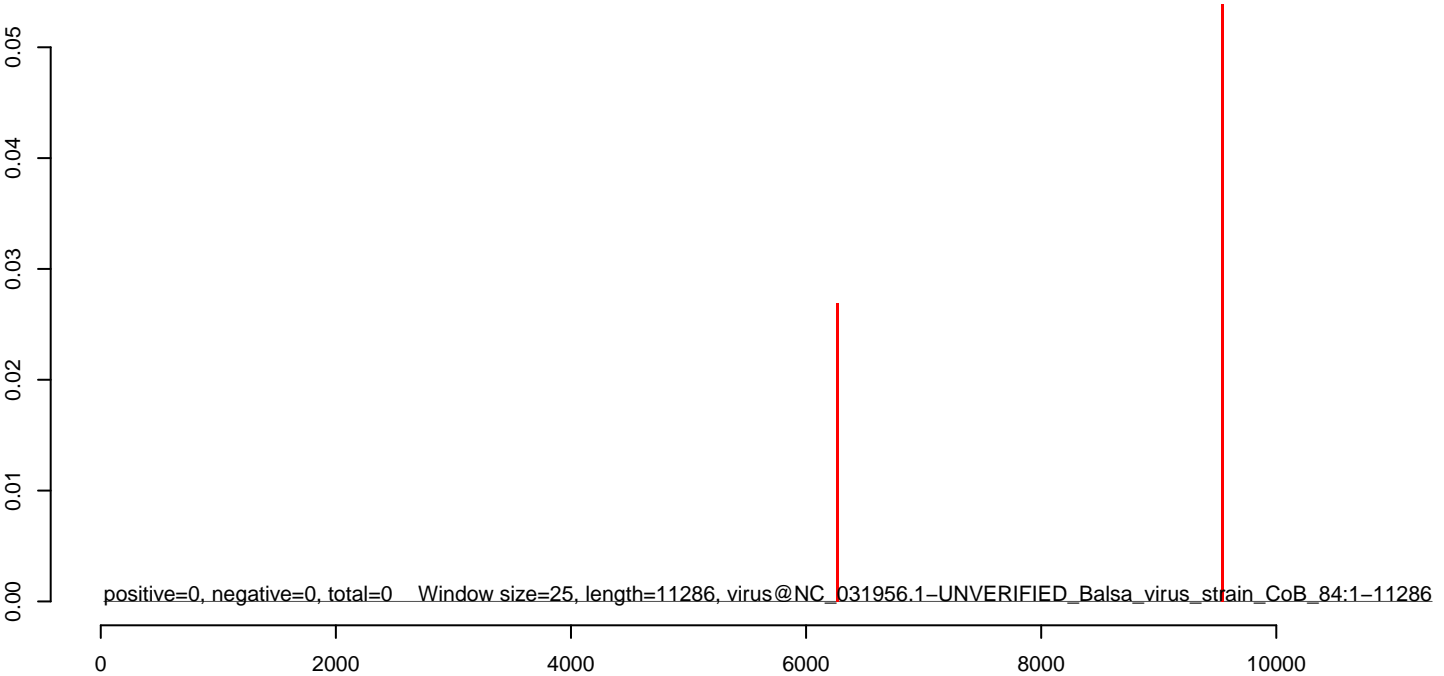
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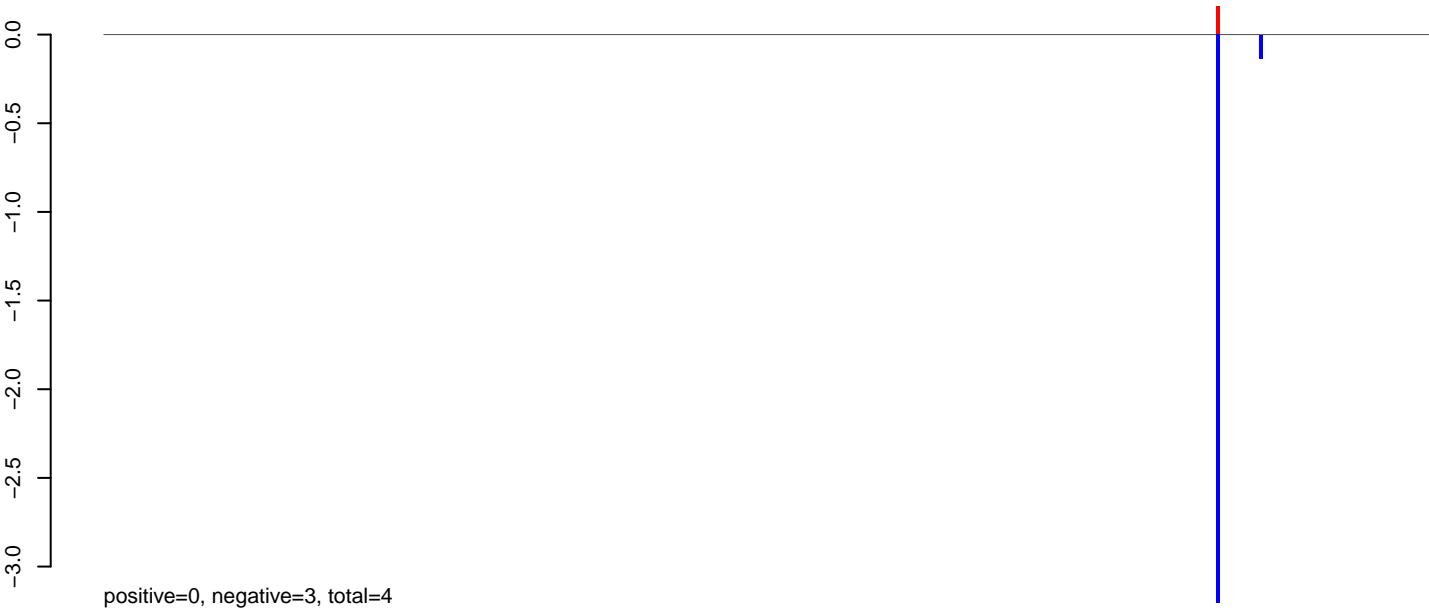
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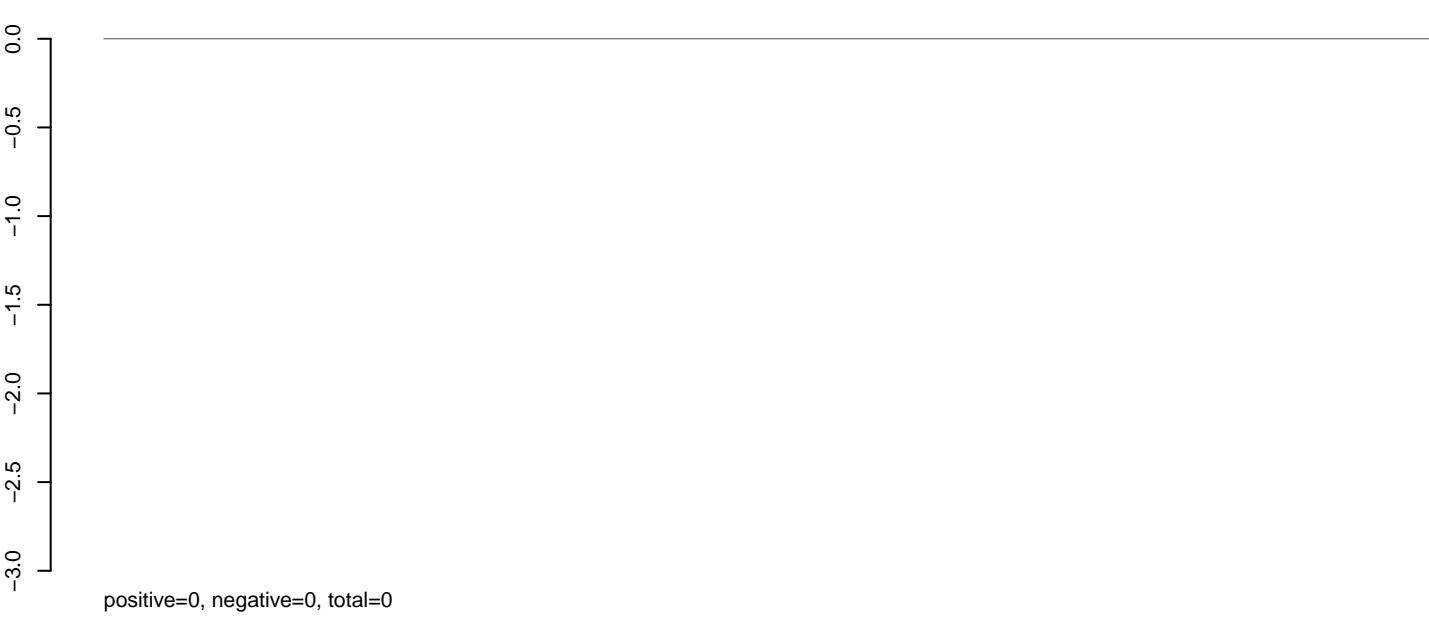
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



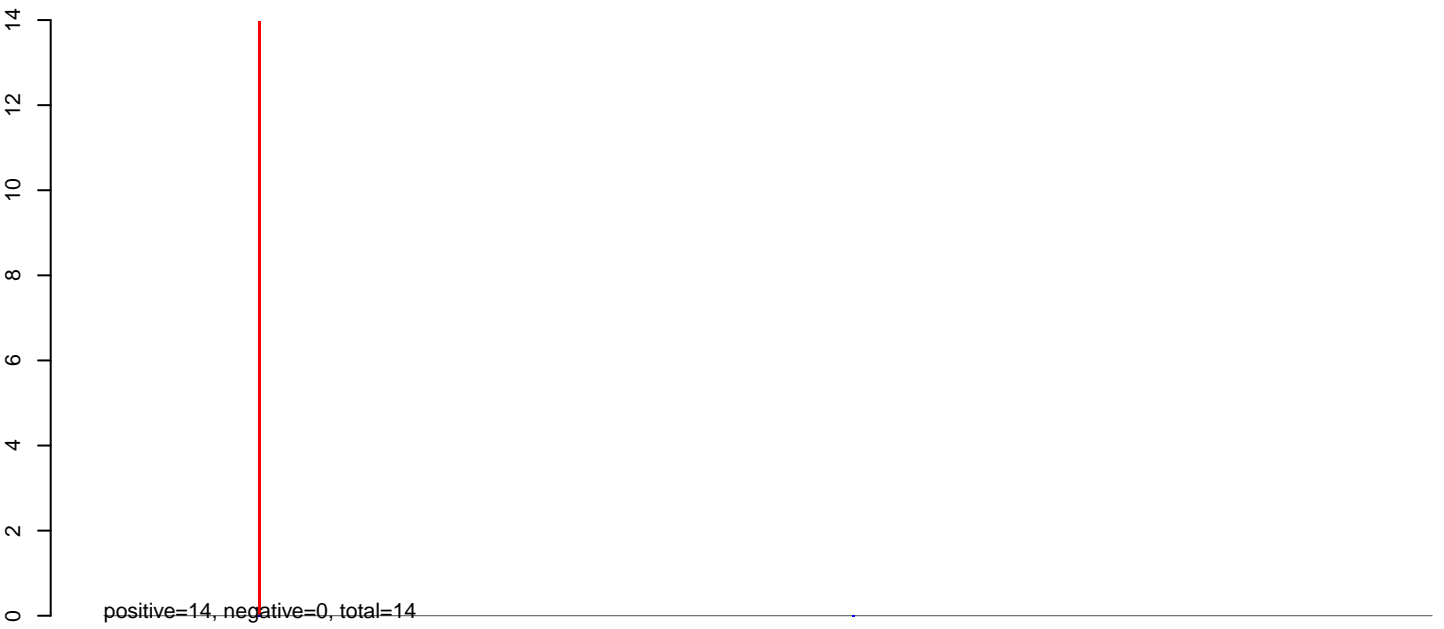
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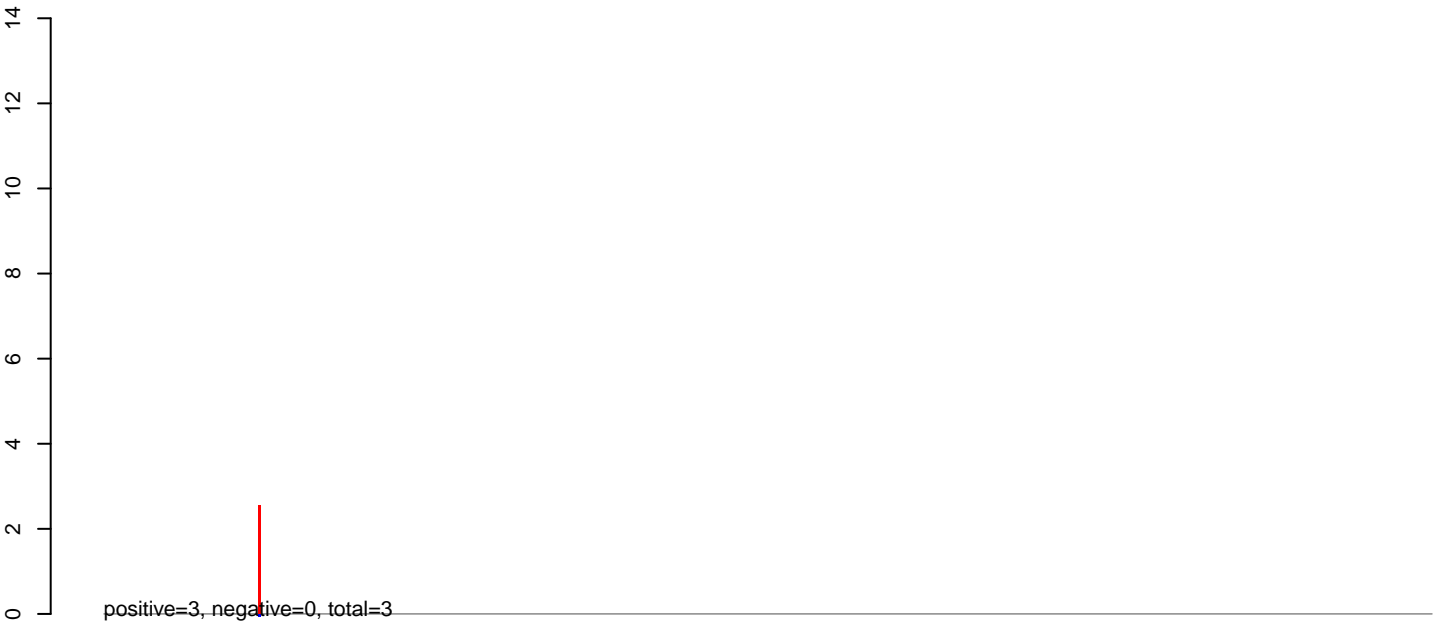
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0 2000 4000 6000 8000 10000

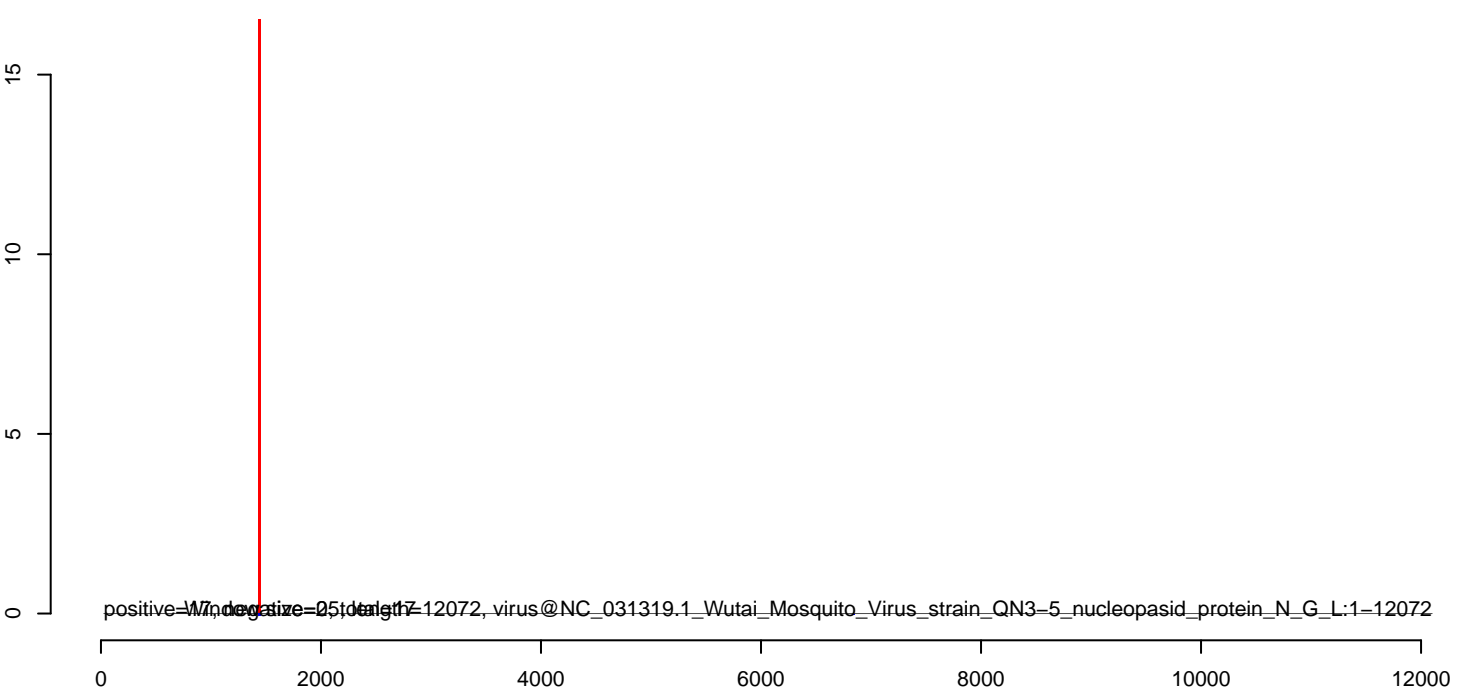
AeAeg_Aag2DENV2NGC.18_23.rep



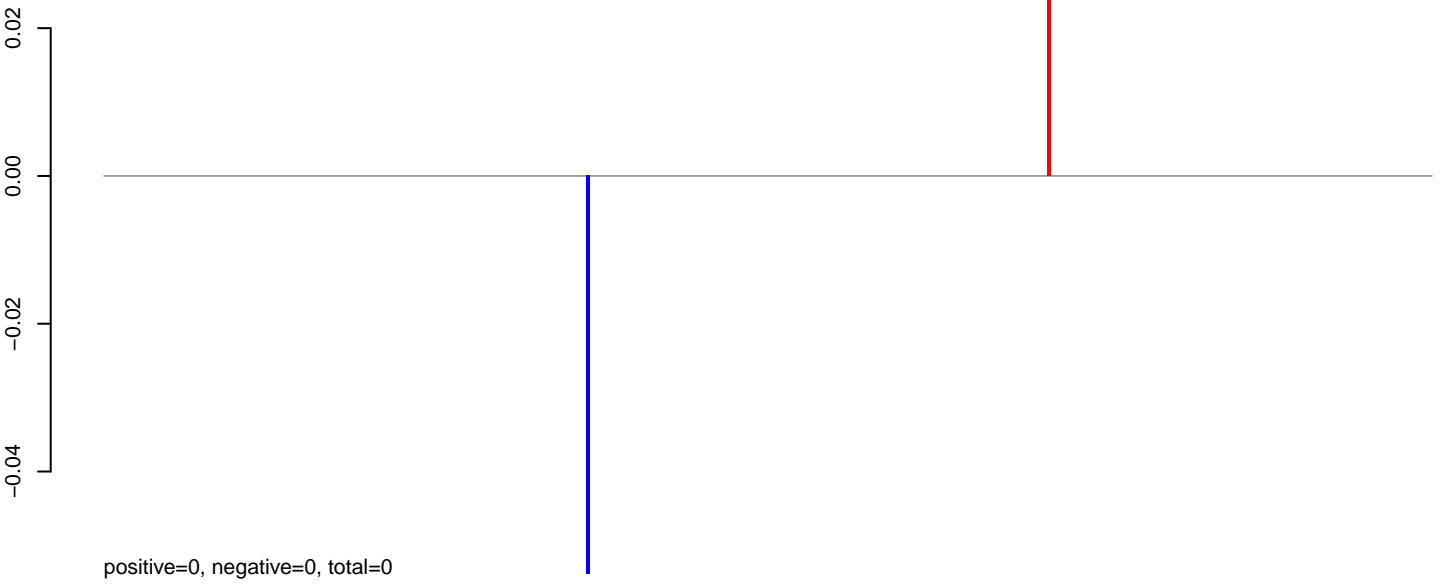
AeAeg_Aag2DENV2NGC.24_35.rep



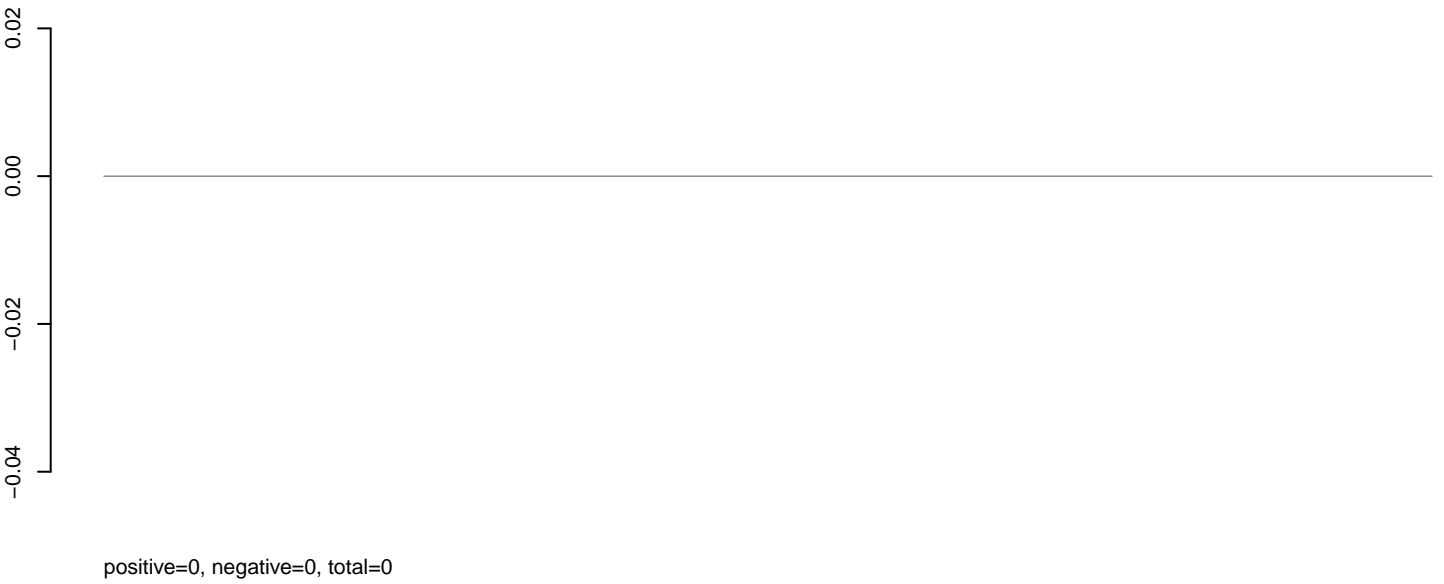
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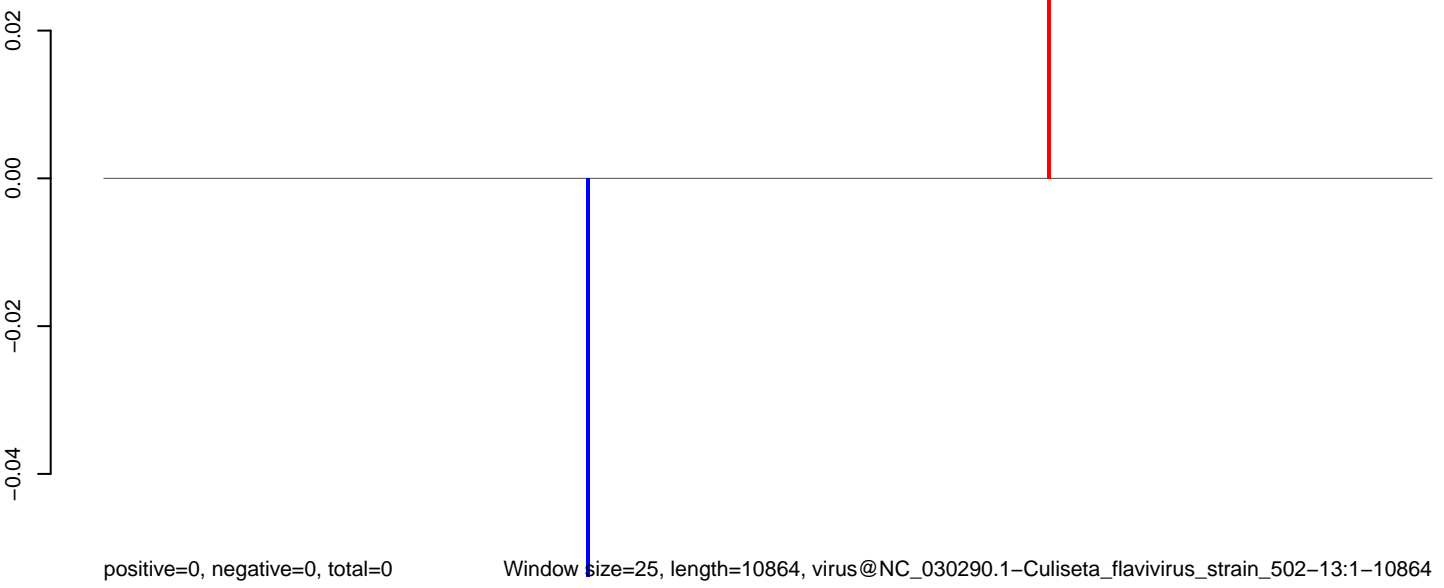
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AeAeg_Aag2DENV2NGC.24_35.rep



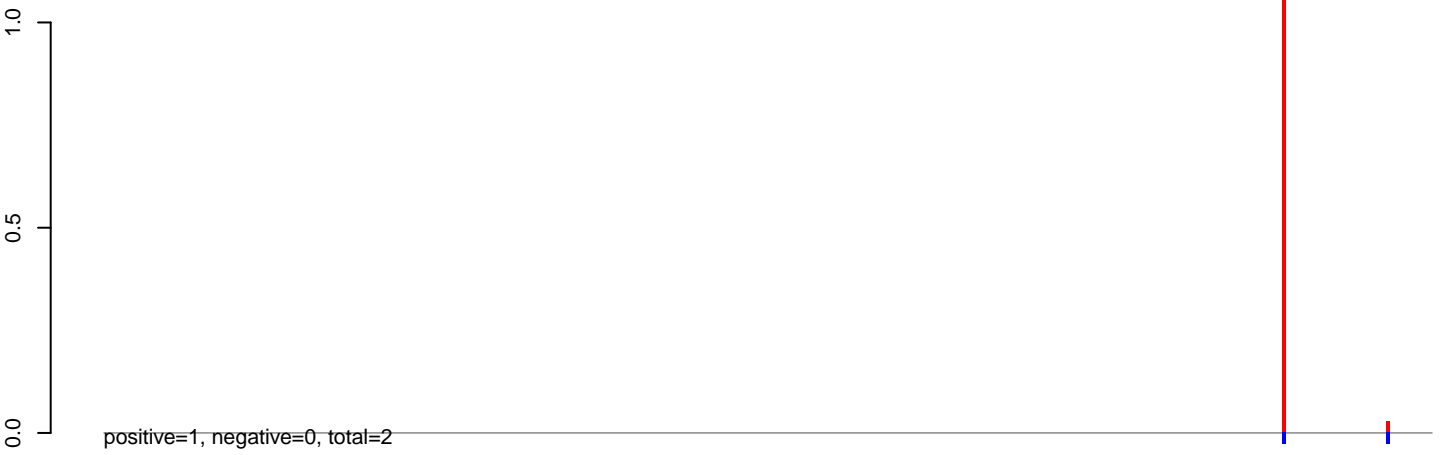
AeAeg_Aag2DENV2NGC.rep



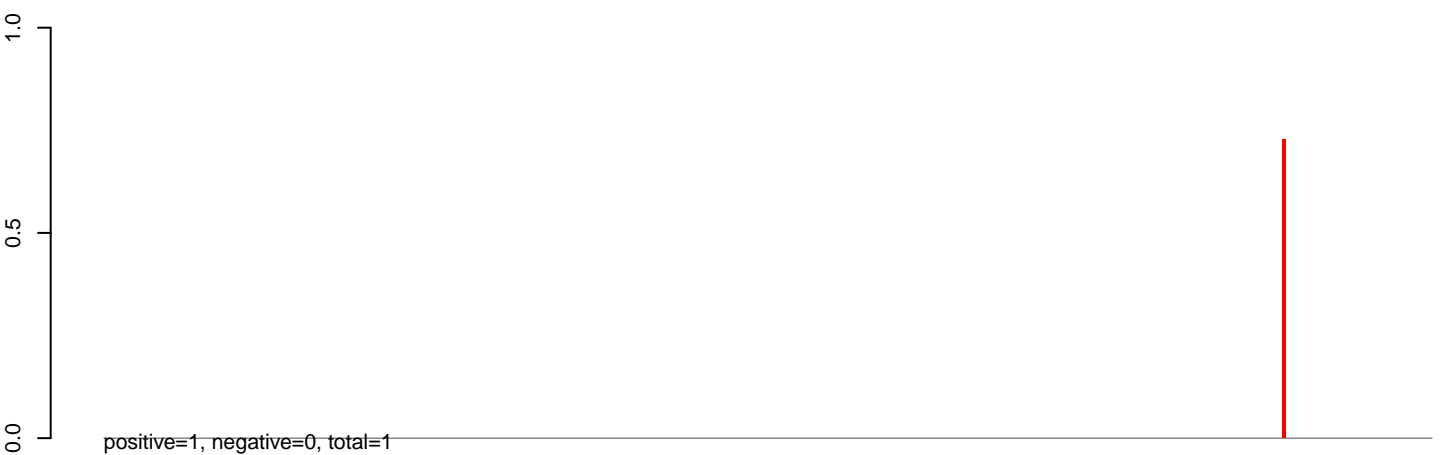
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0 2000 4000 6000 8000 10000

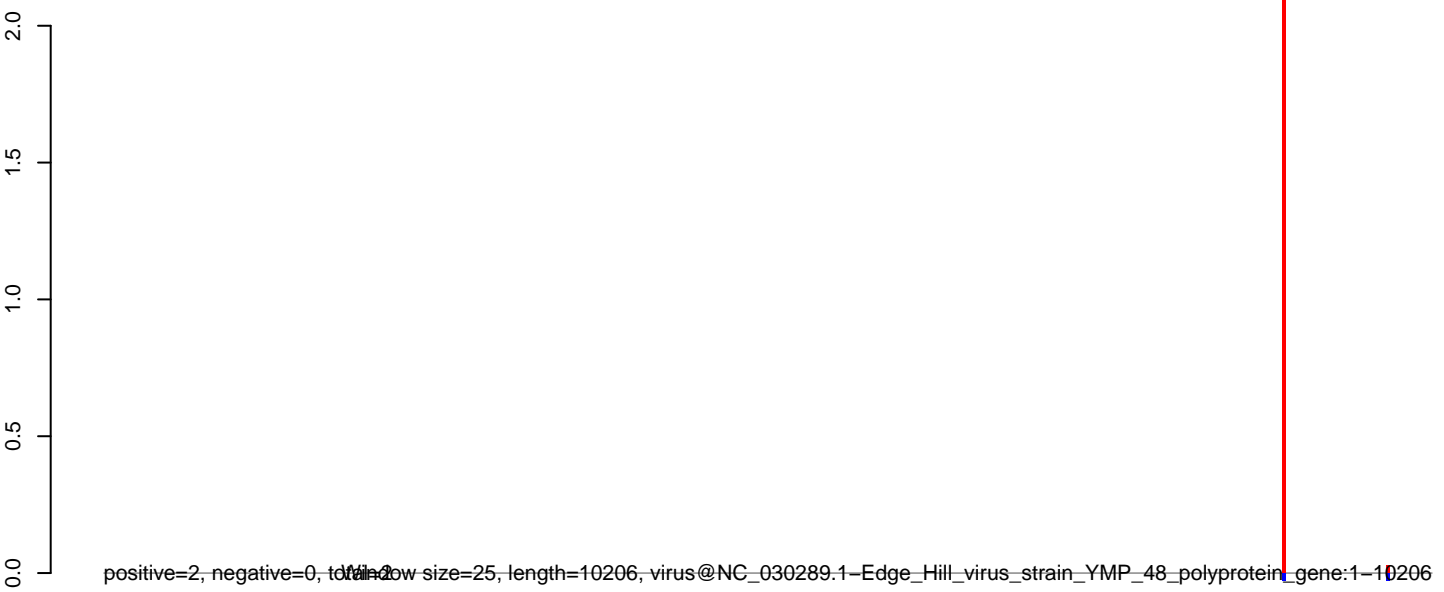
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



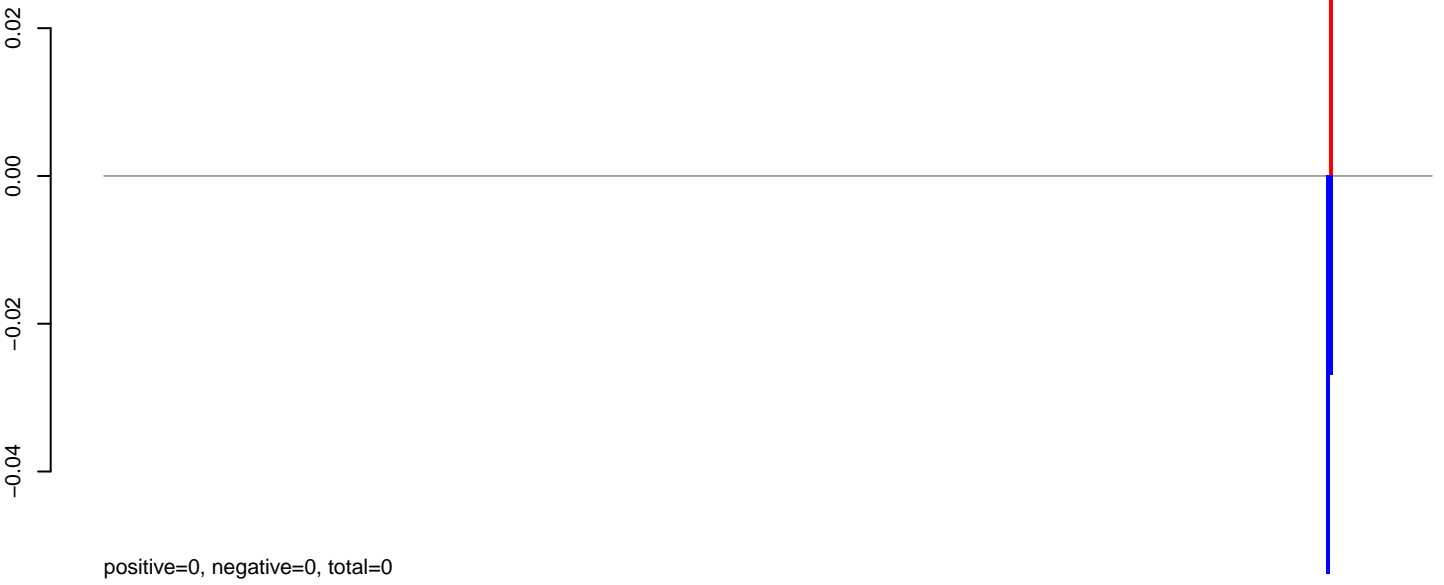
AeAeg_Aag2DENV2NGC.rep



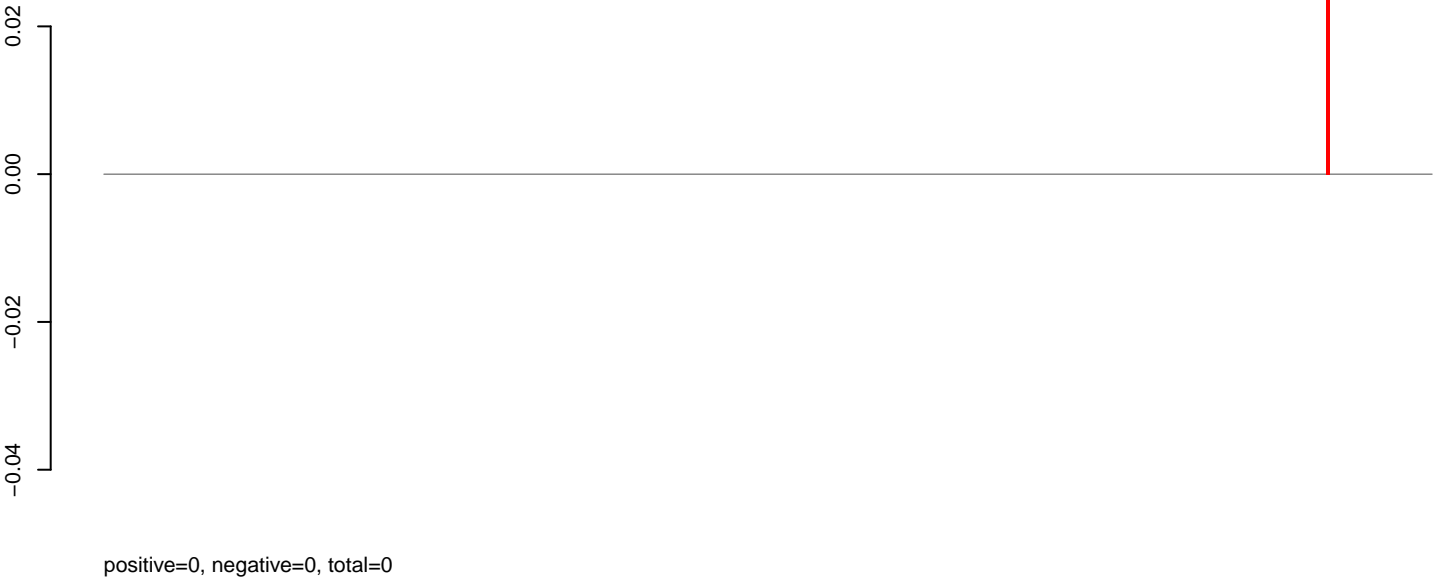
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0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



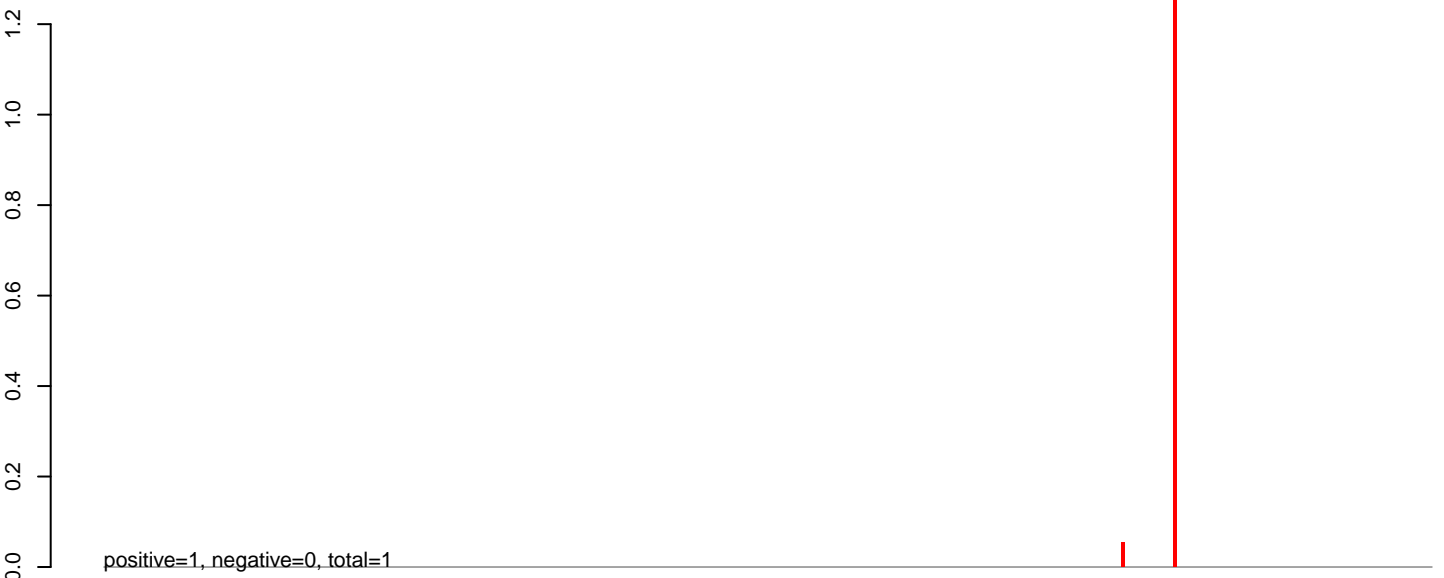
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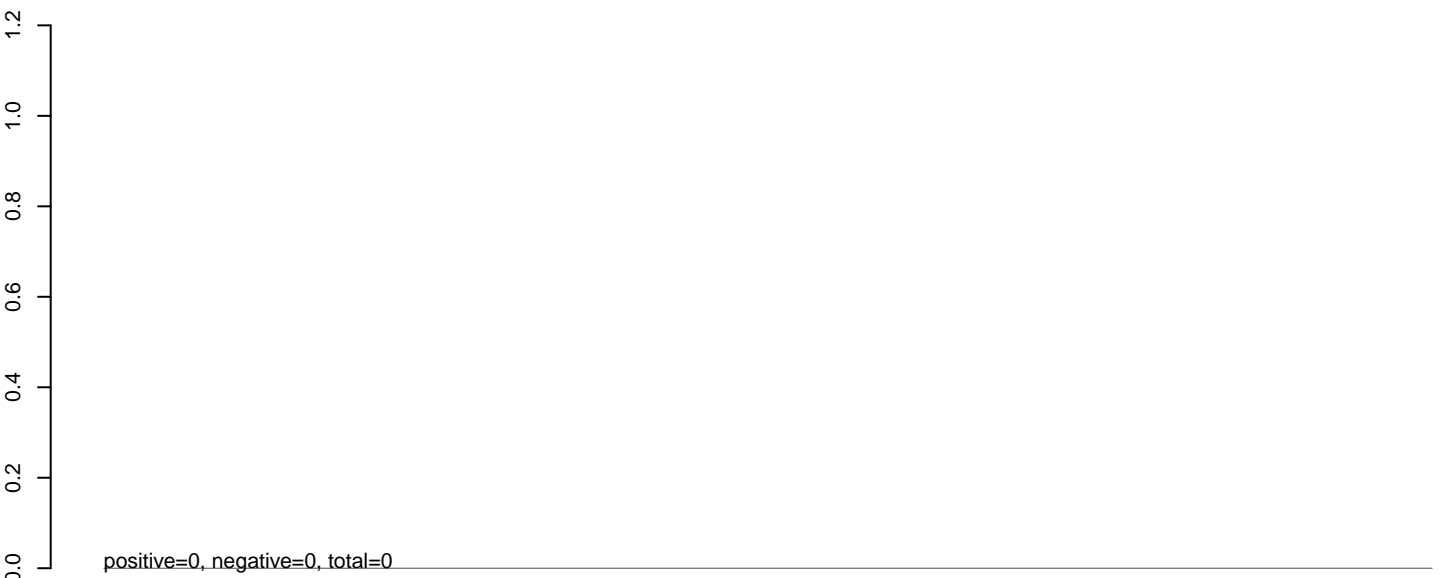
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0 2000 4000 6000 8000 10000

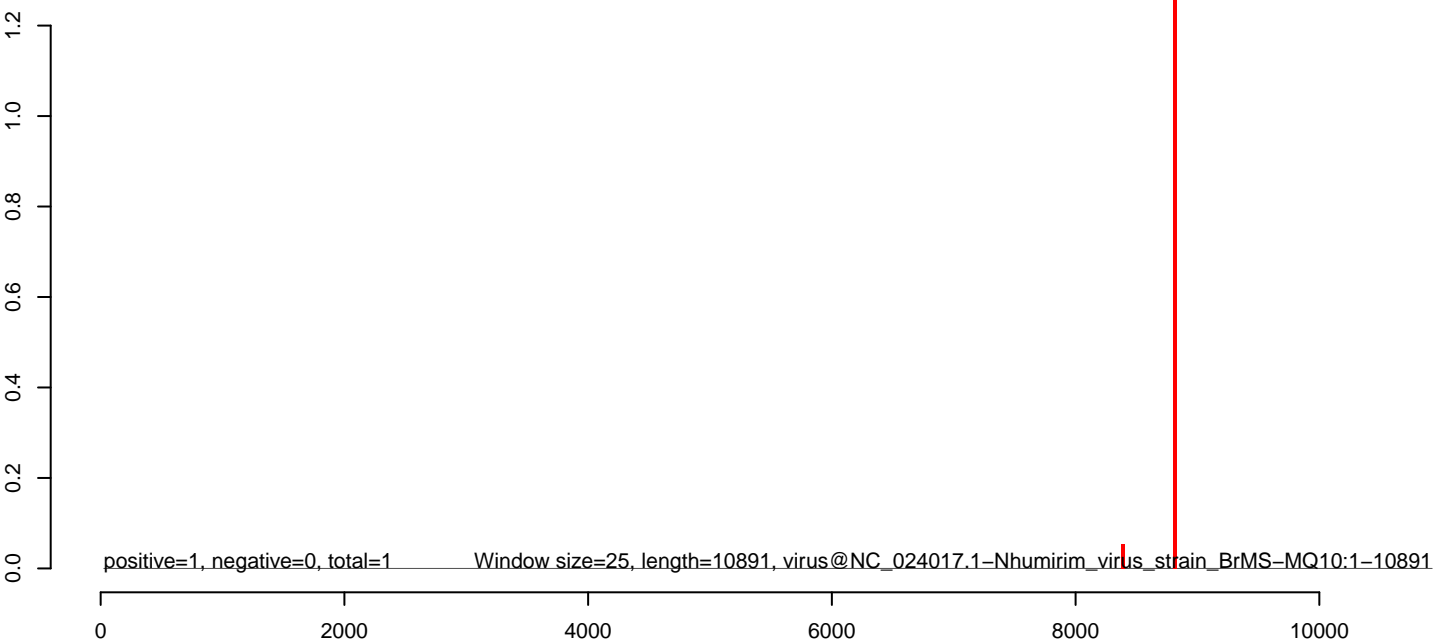
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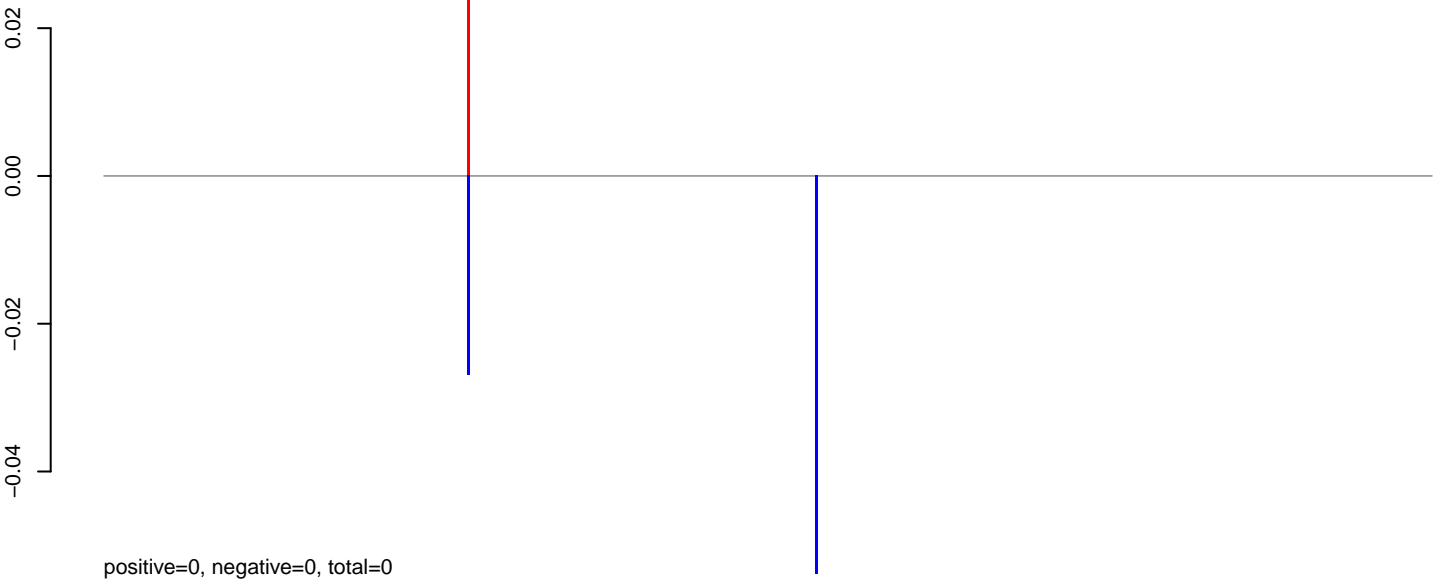
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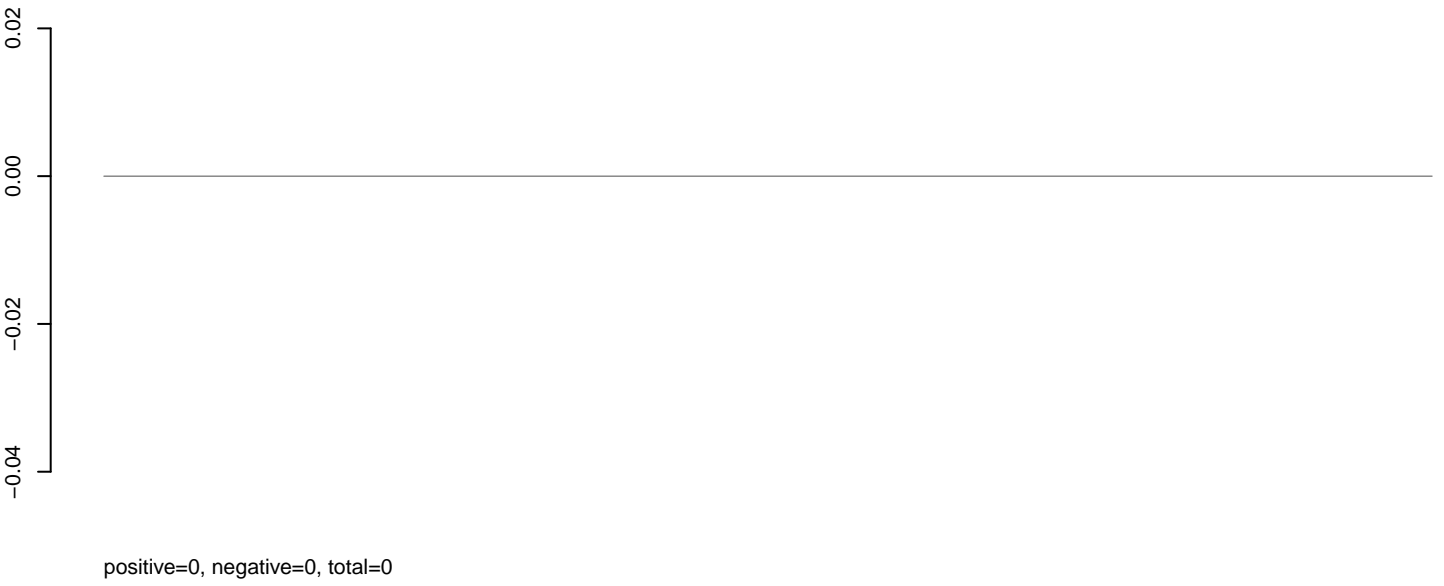
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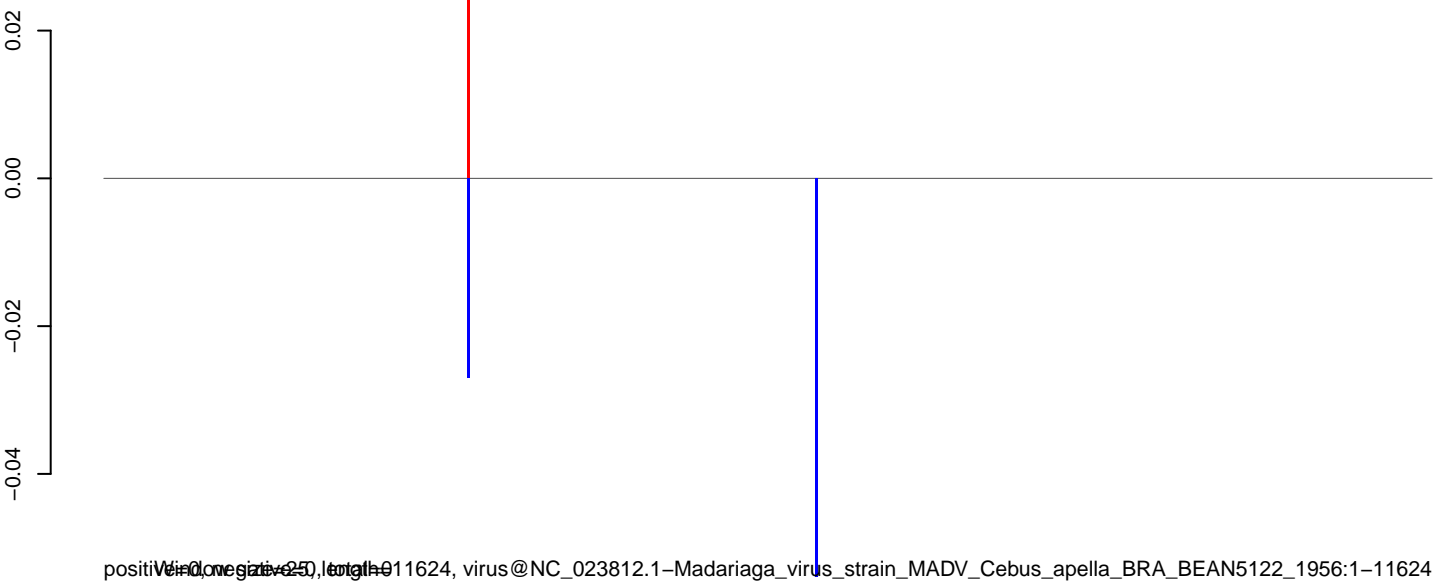
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AeAeg_Aag2DENV2NGC.24_35.rep

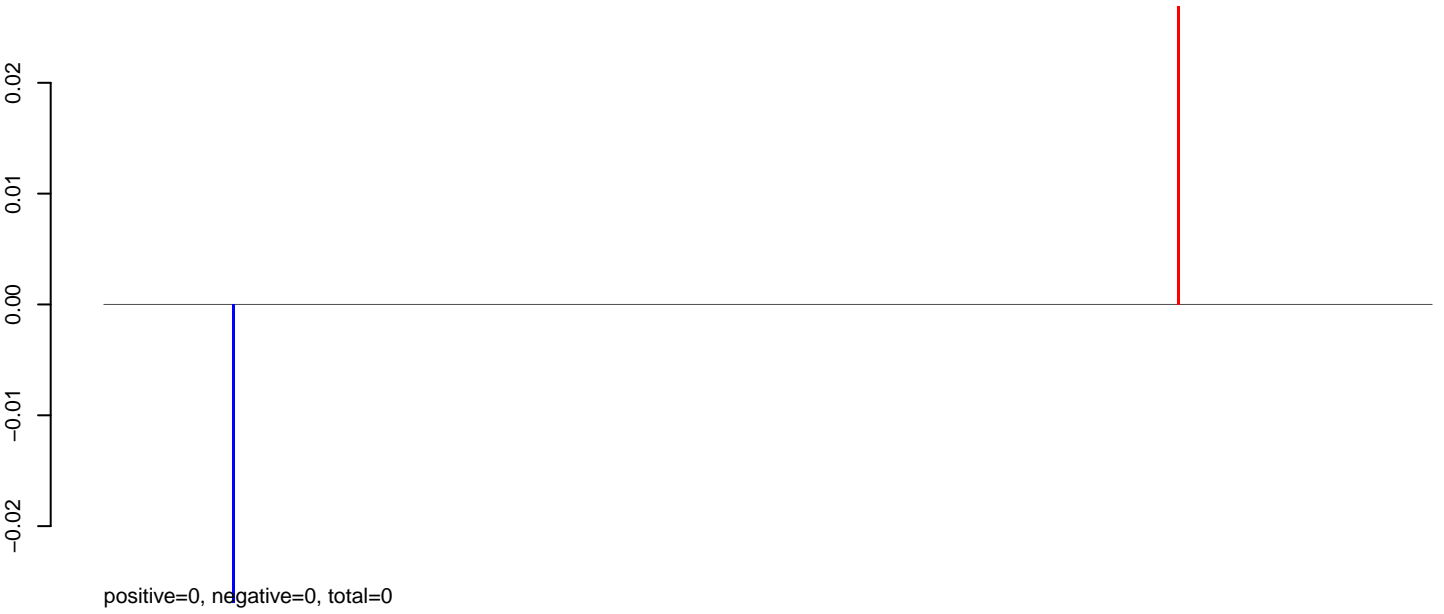


AeAeg_Aag2DENV2NGC.rep

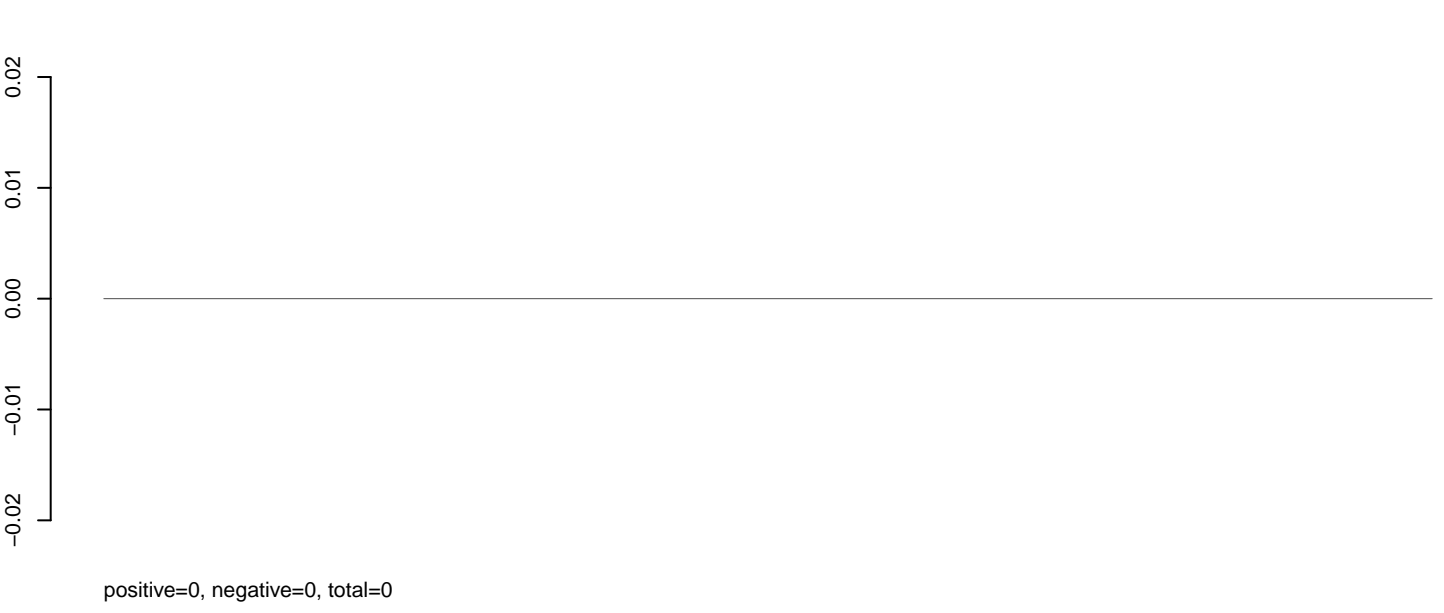


0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2DENV2NGC.18_23.rep



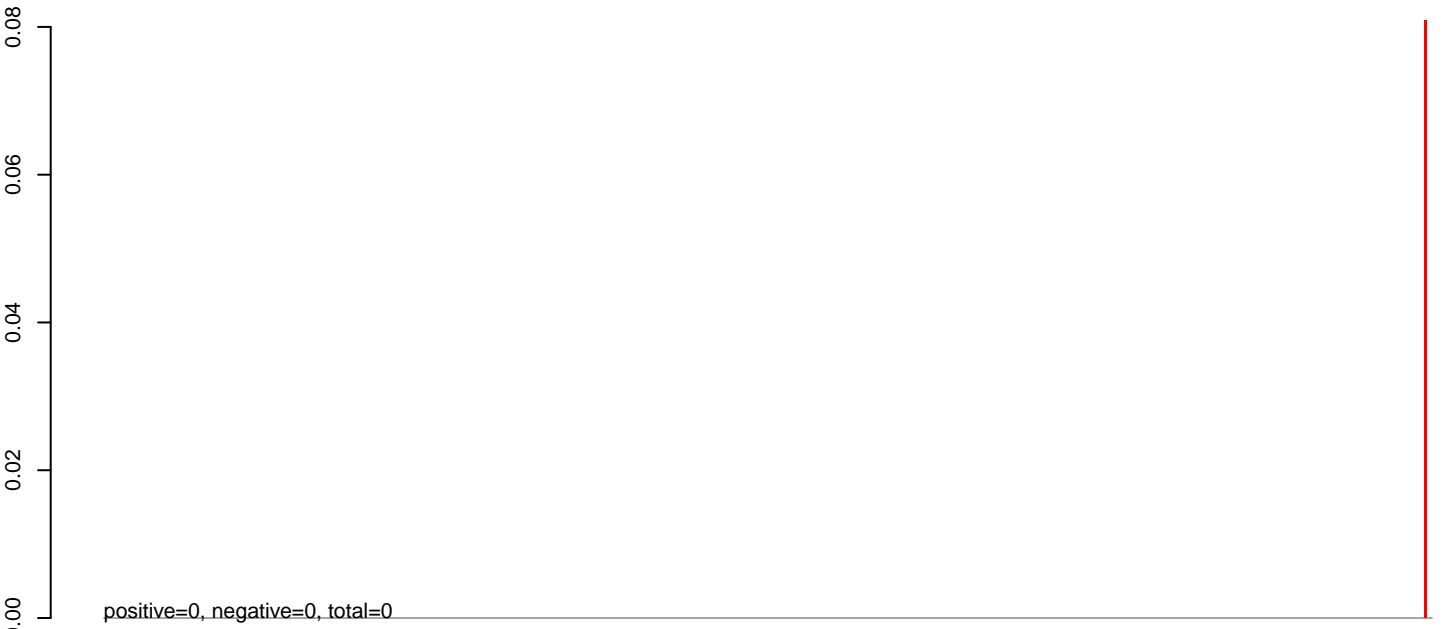
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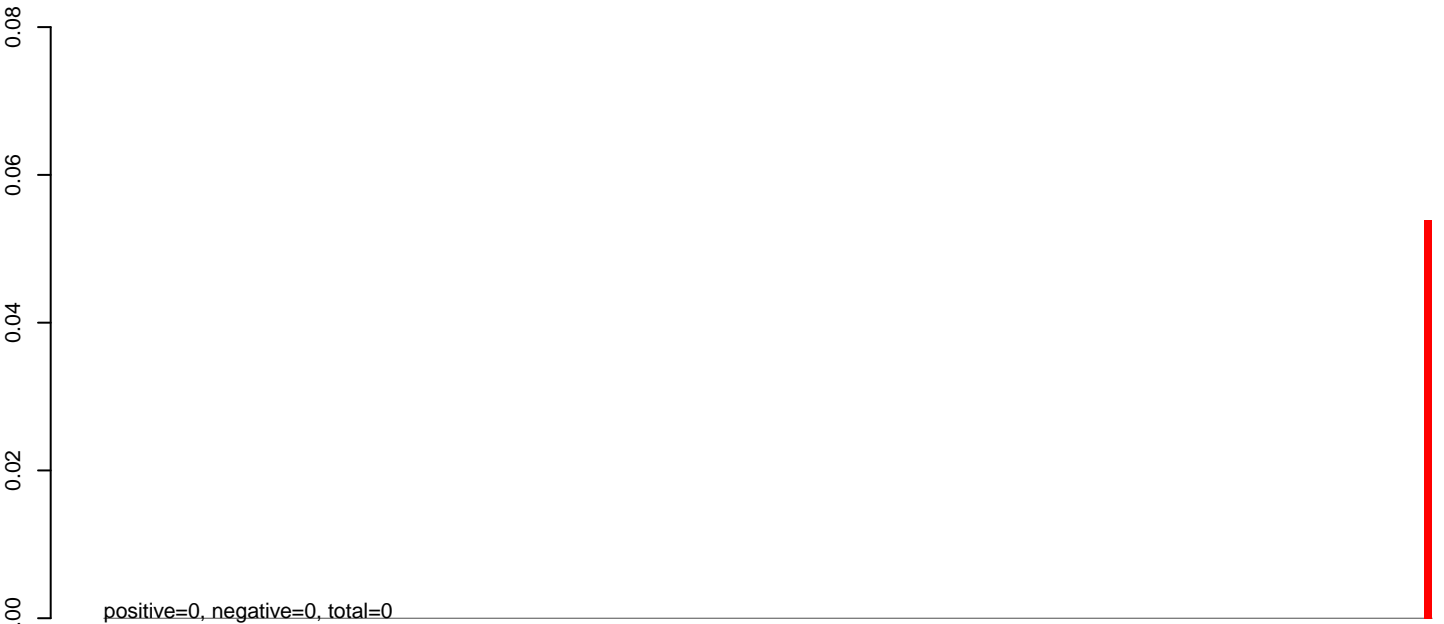
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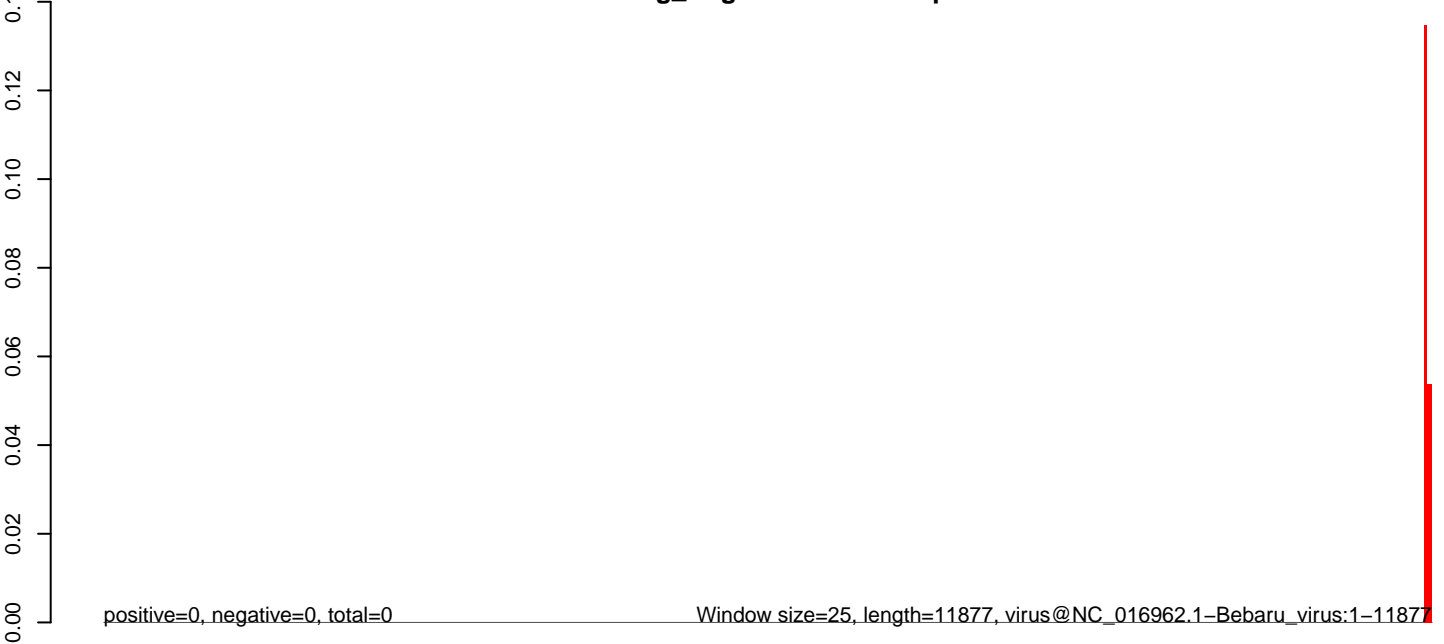
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AeAeg_Aag2DENV2NGC.24_35.rep

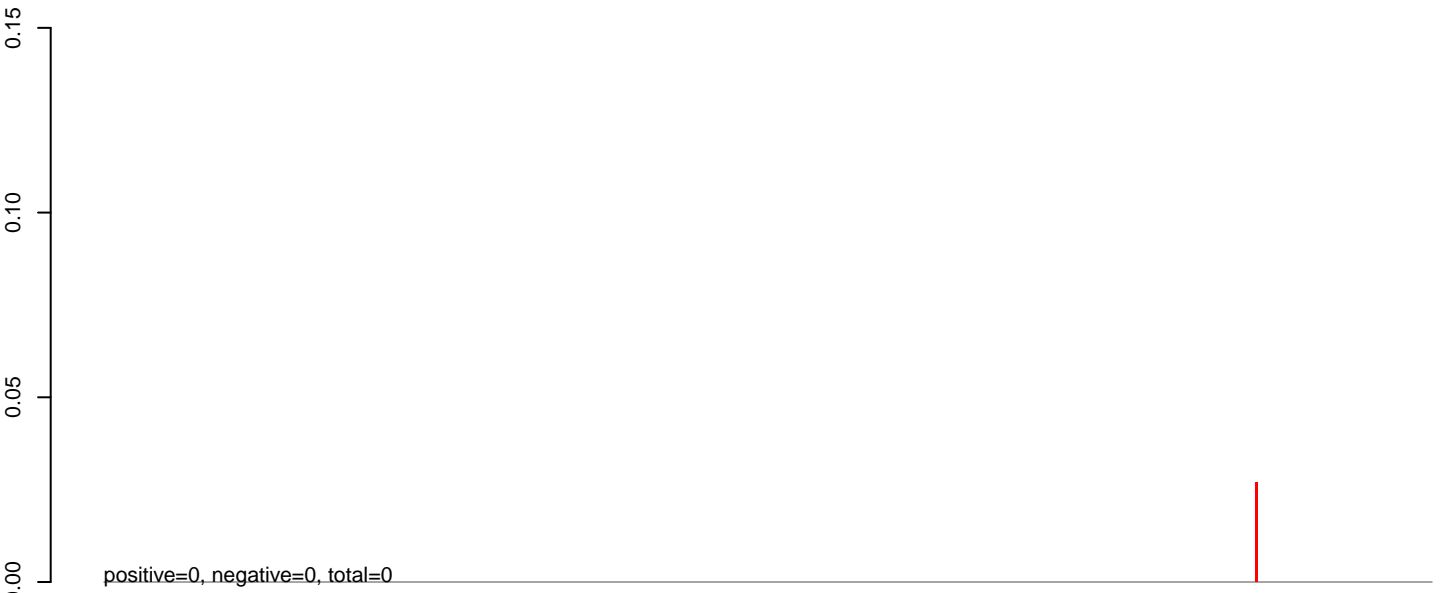


AeAeg_Aag2DENV2NGC.rep

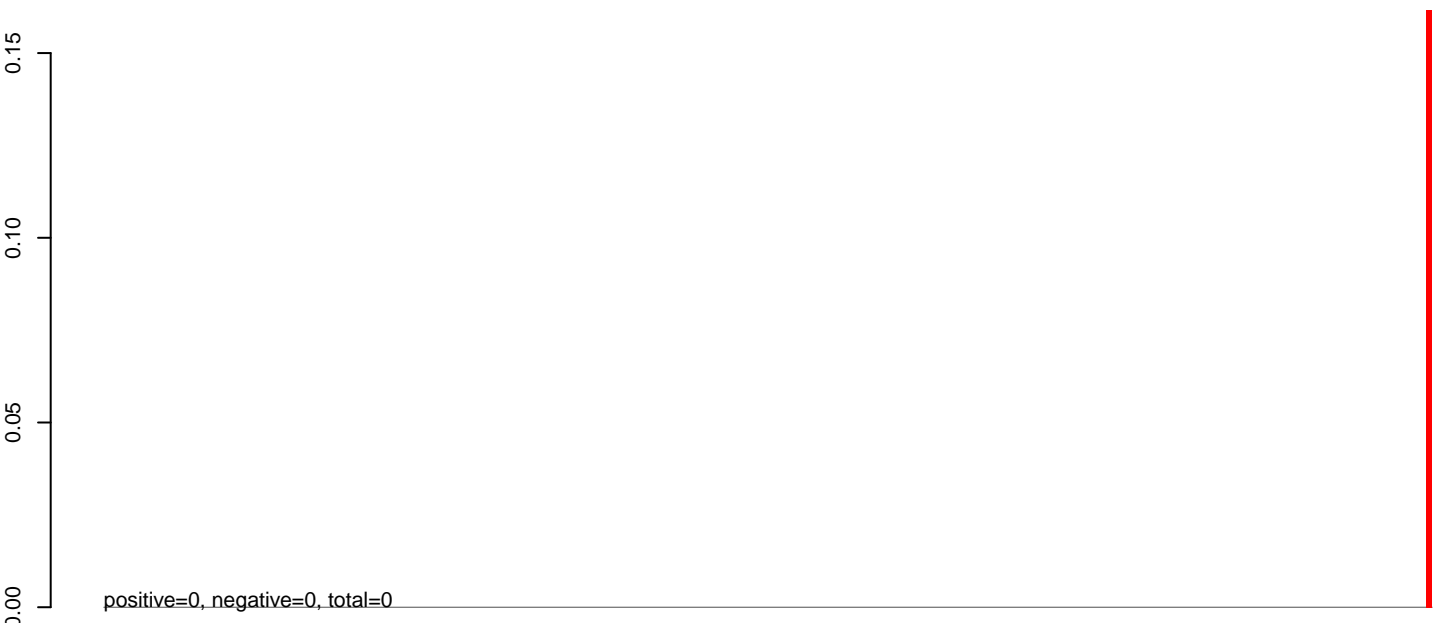


0 2000 4000 6000 8000 10000 12000

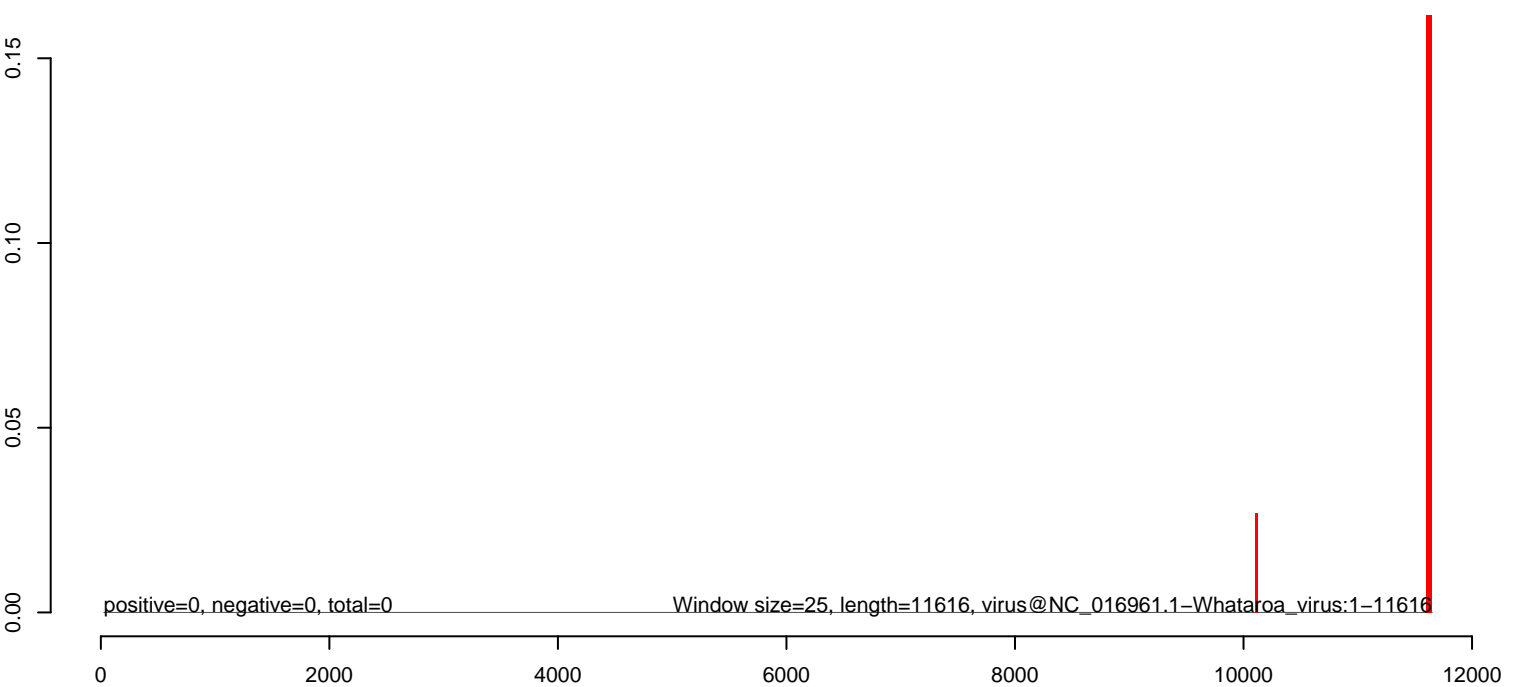
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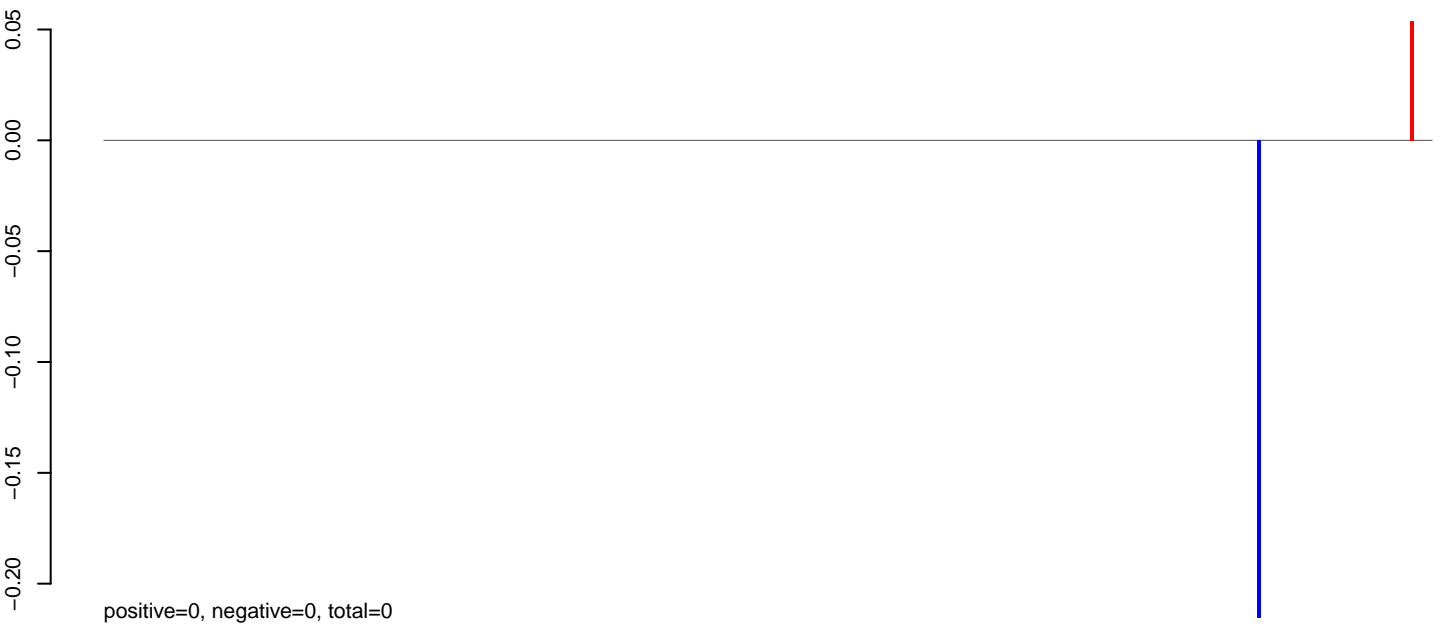
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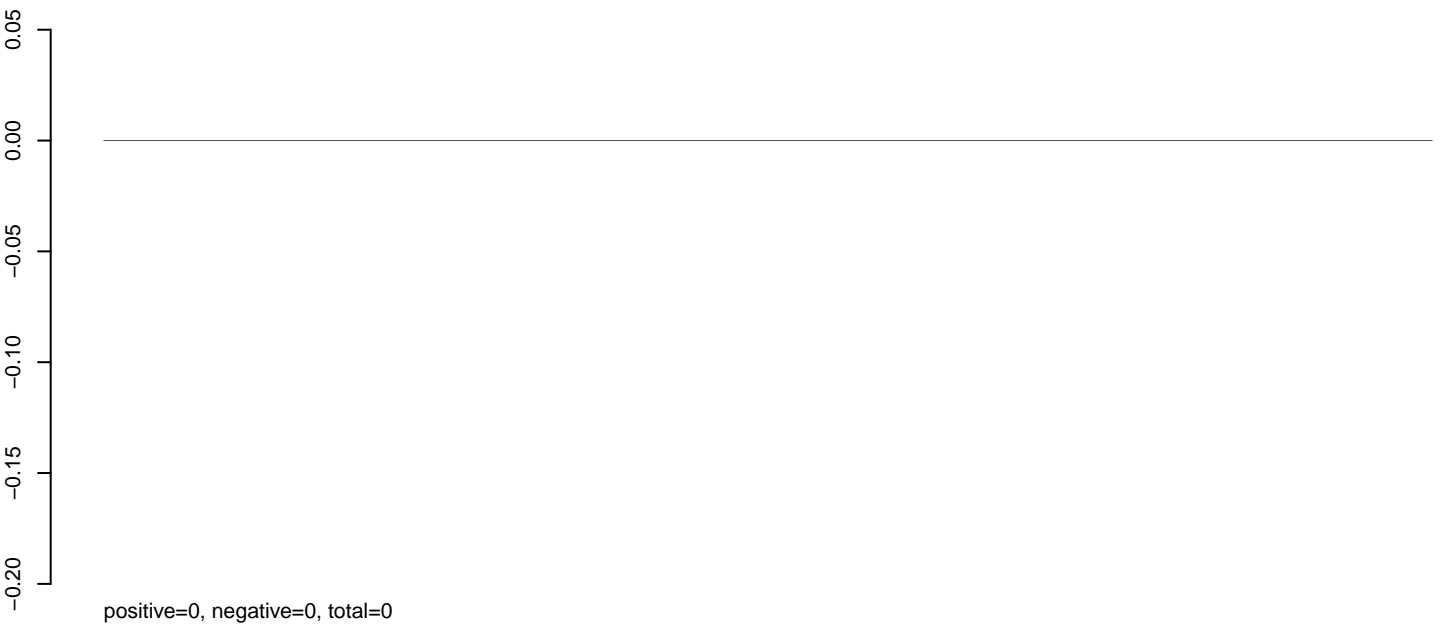
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



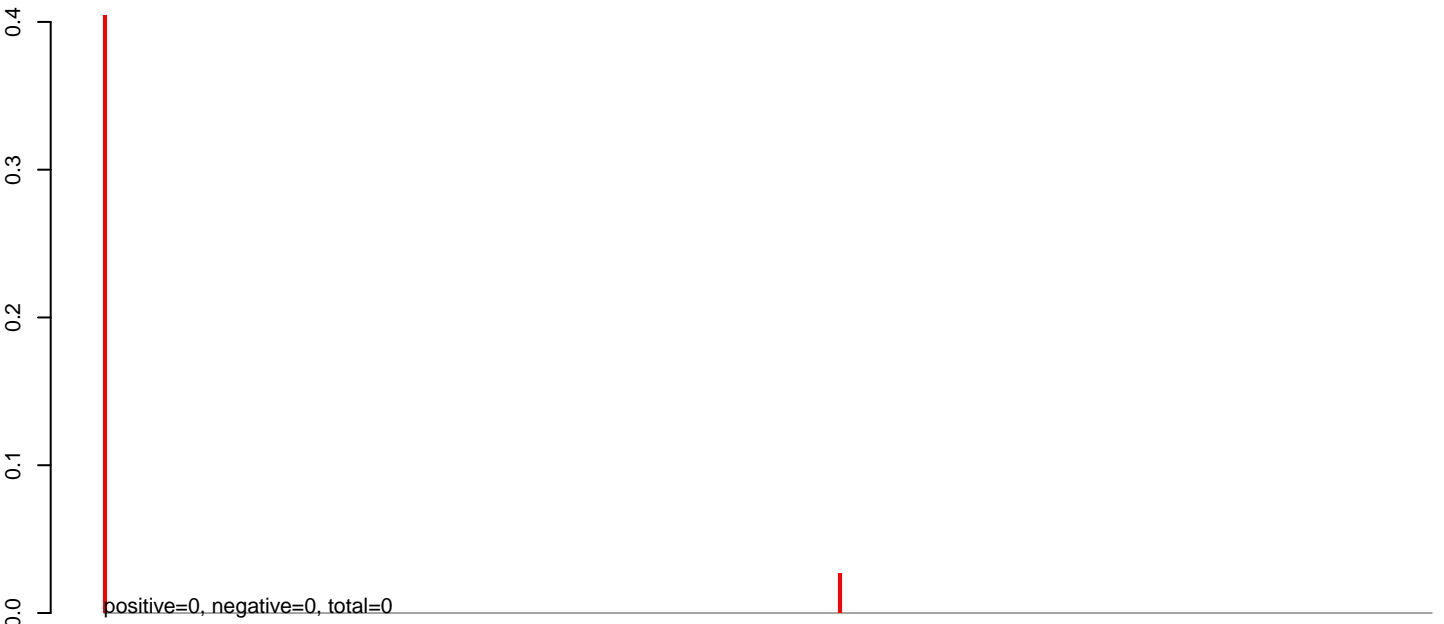
AeAeg_Aag2DENV2NGC.rep



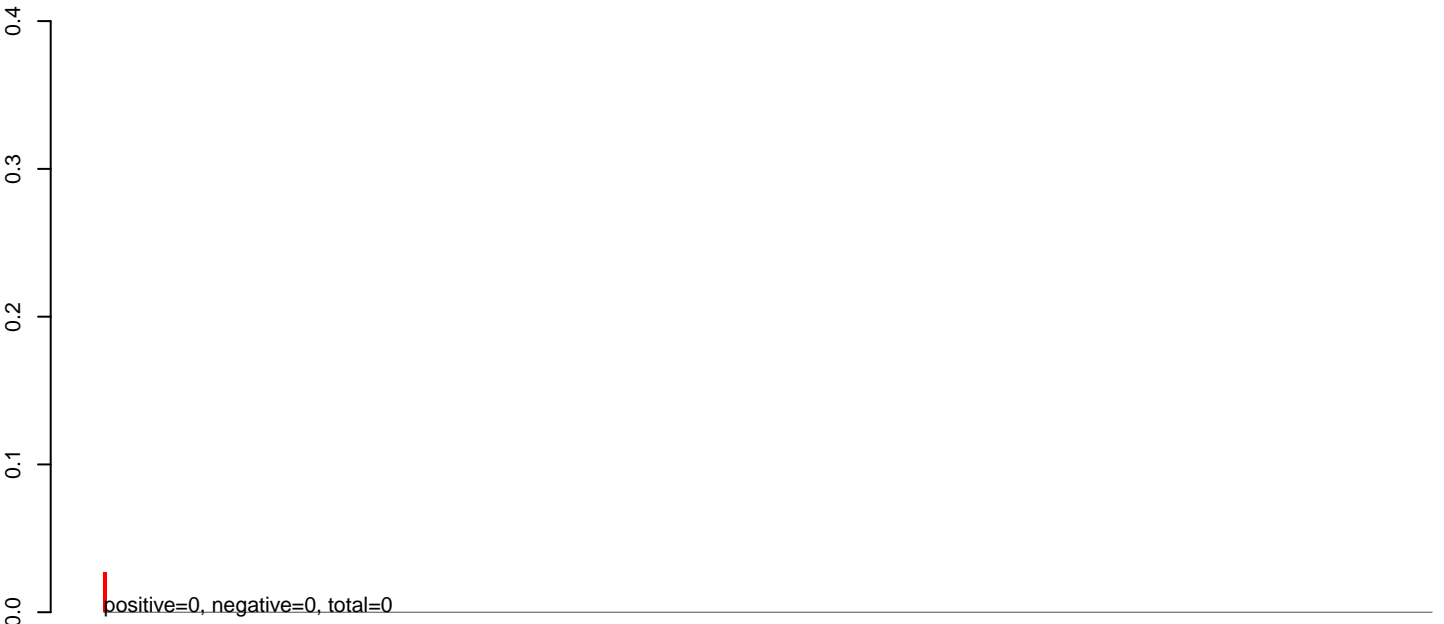
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0 2000 4000 6000 8000 10000

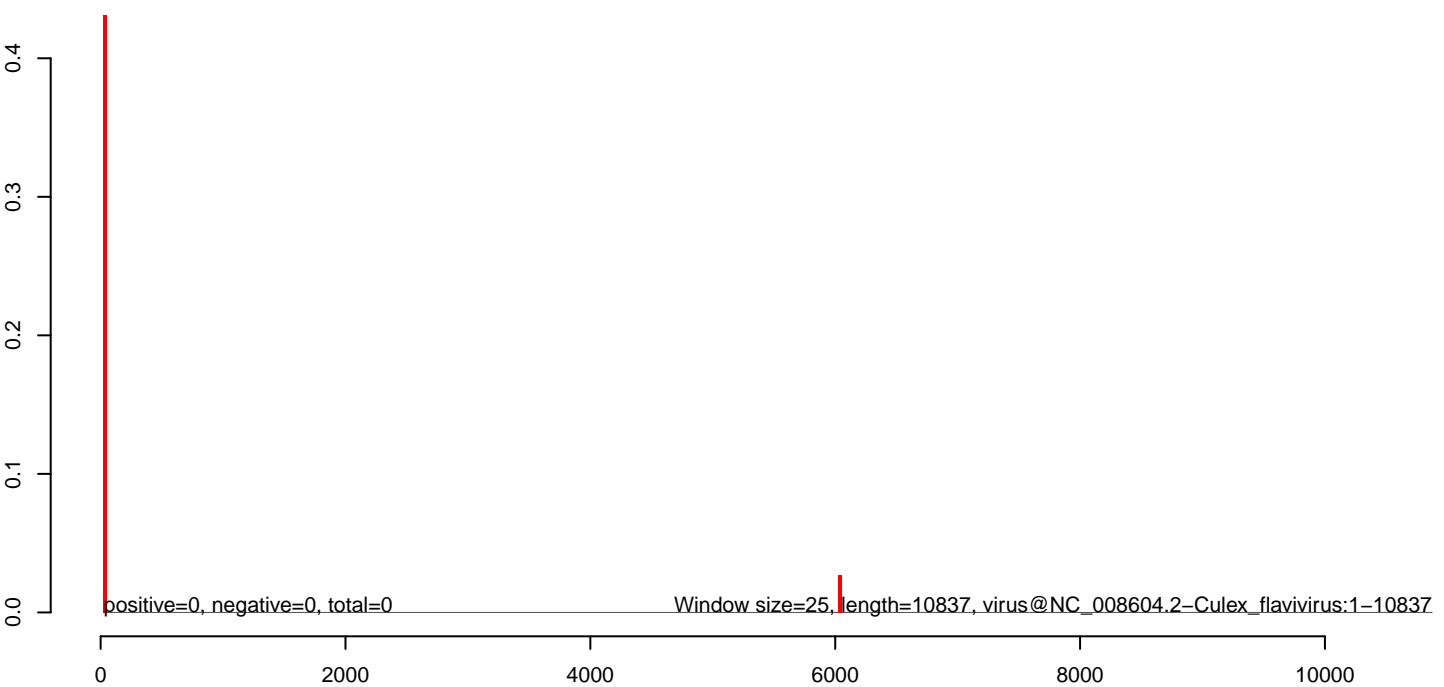
AeAeg_Aag2DENV2NGC.18_23.rep



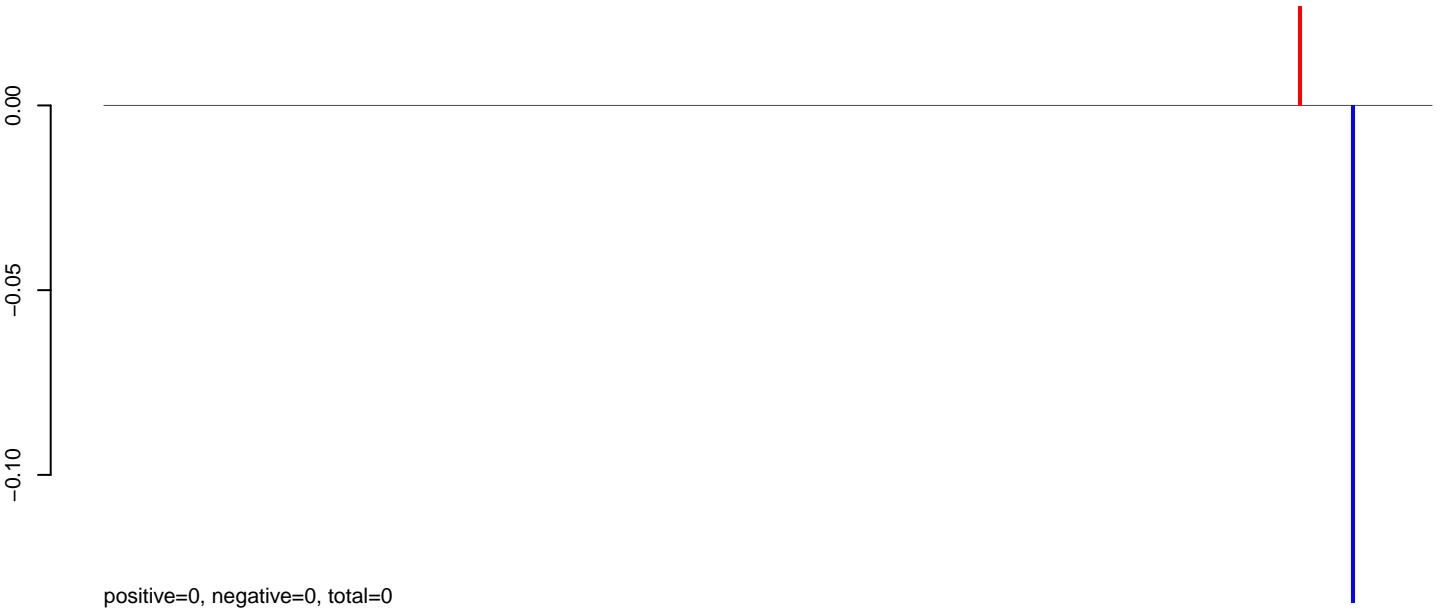
AeAeg_Aag2DENV2NGC.24_35.rep



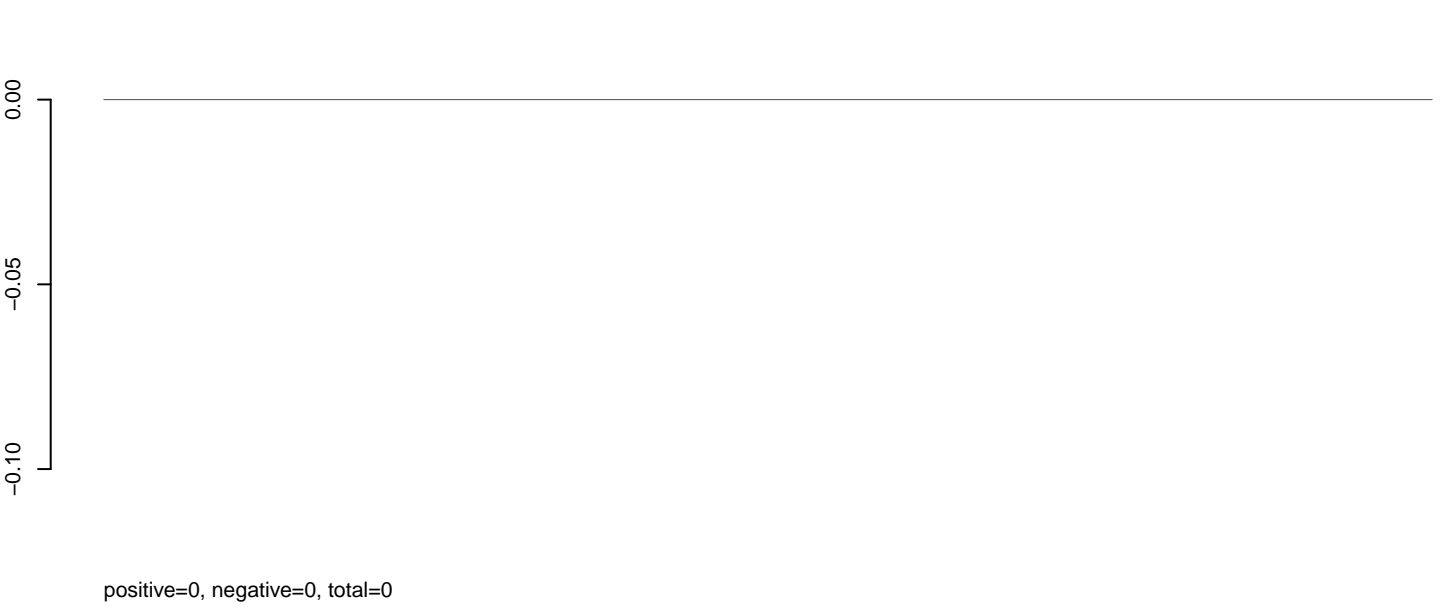
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



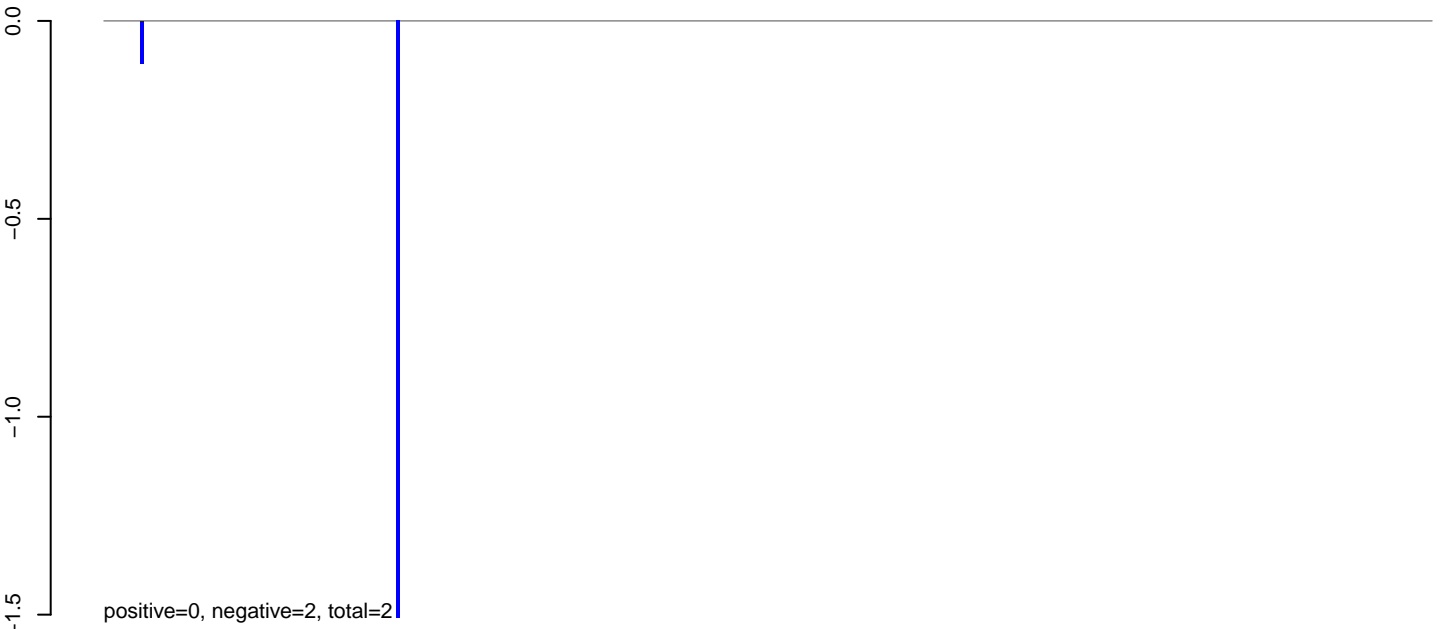
AeAeg_Aag2DENV2NGC.24_35.rep



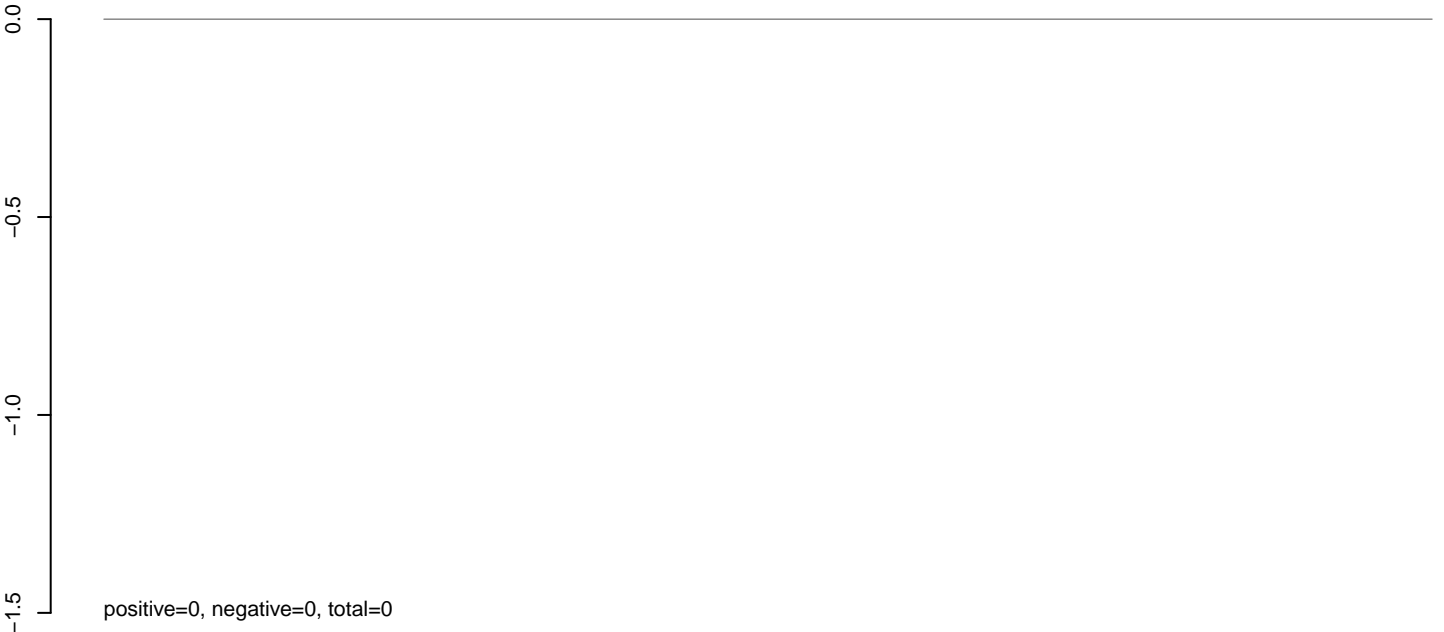
AeAeg_Aag2DENV2NGC.rep



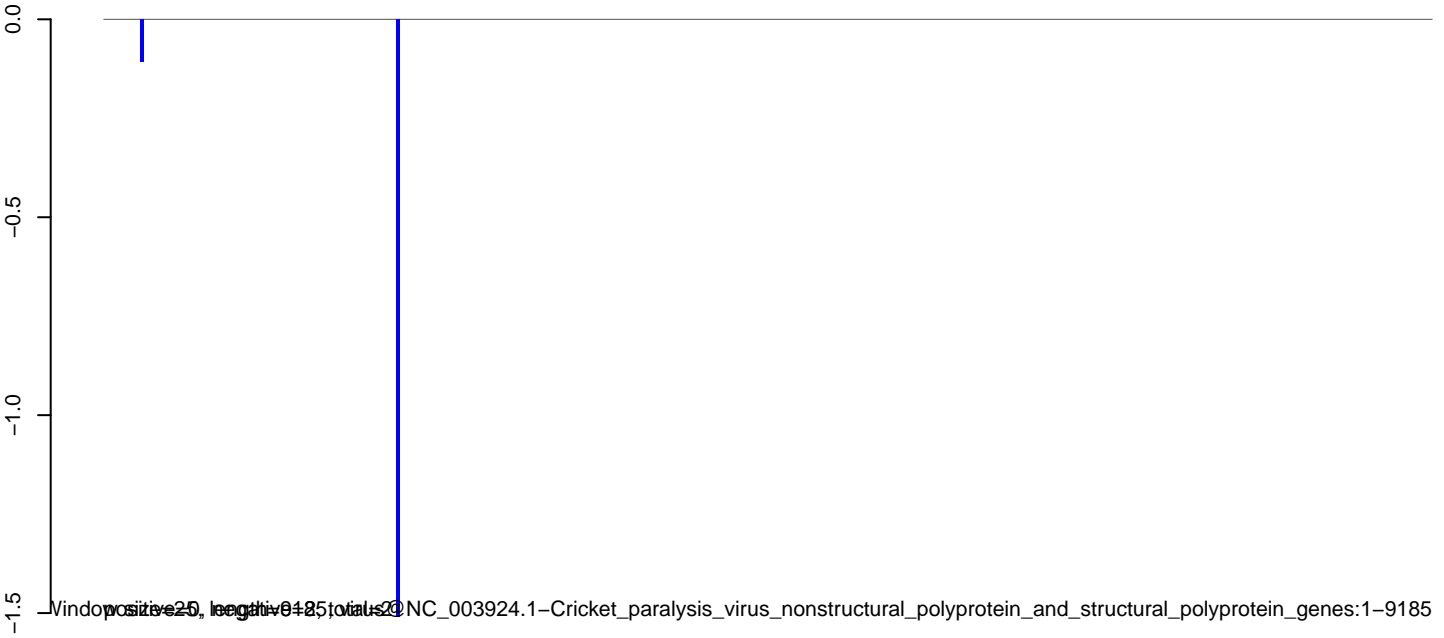
AeAeg_Aag2DENV2NGC.18_23.rep



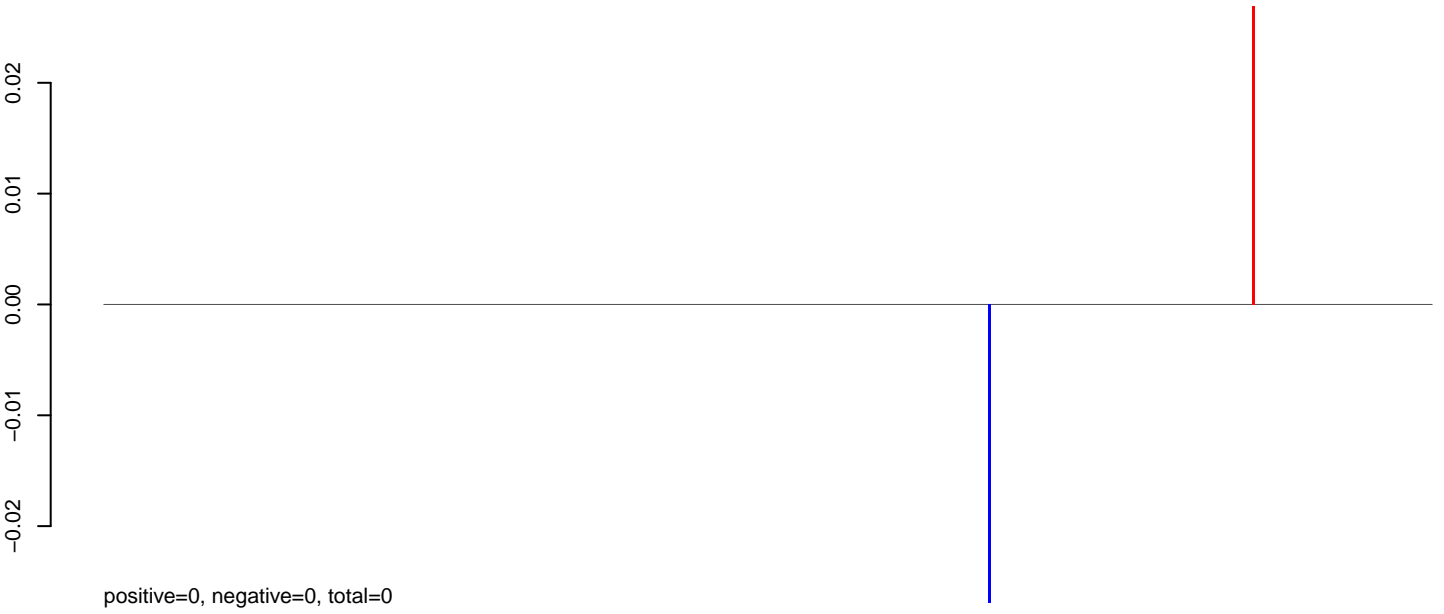
AeAeg_Aag2DENV2NGC.24_35.rep



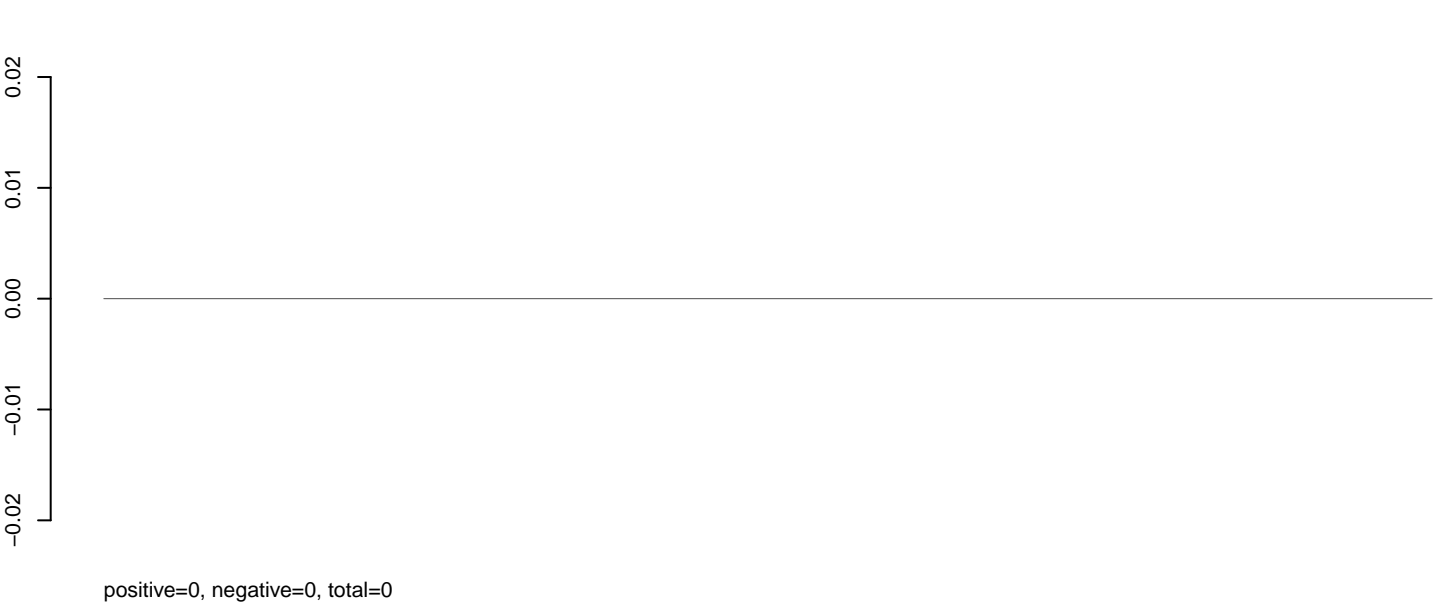
AeAeg_Aag2DENV2NGC.rep



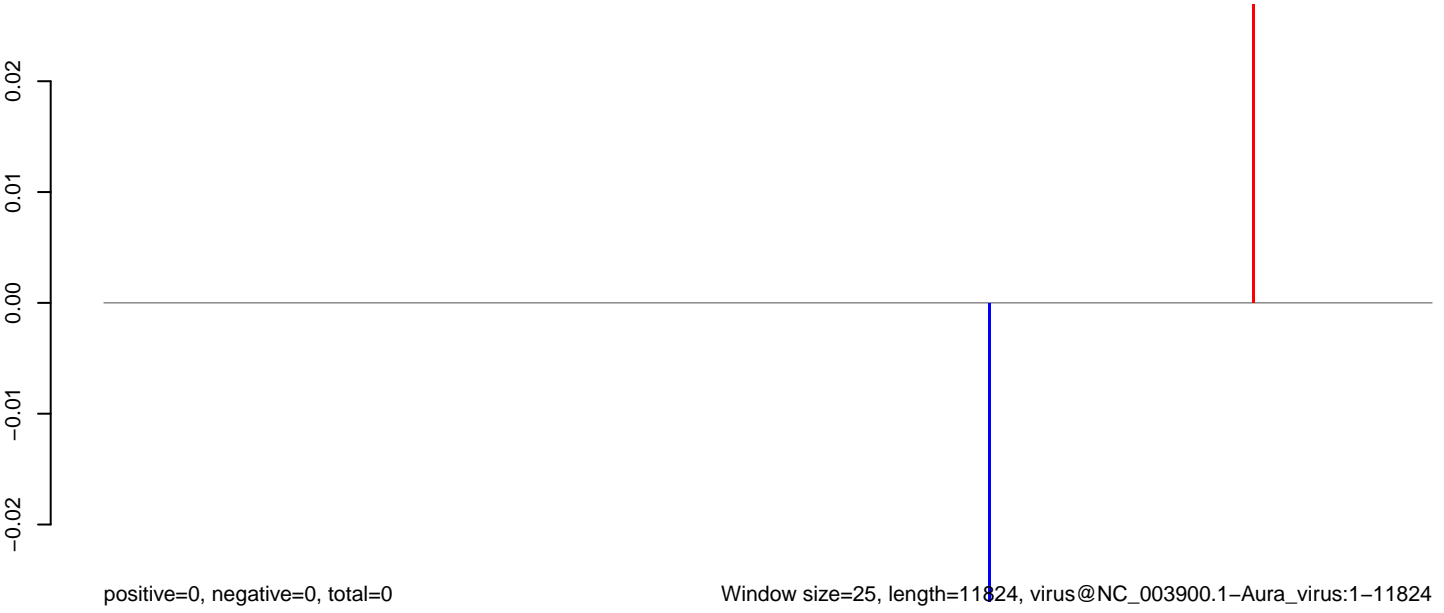
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



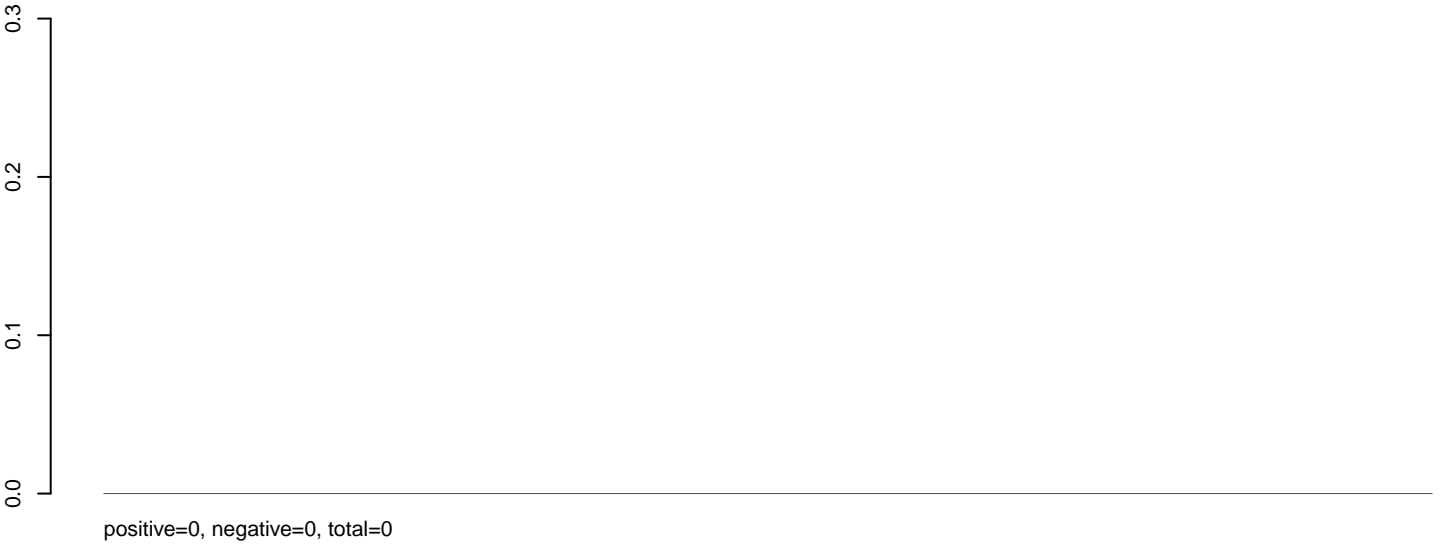
Window size=25, length=11824, virus@NC_003900.1-Aura_virus:1-11824

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

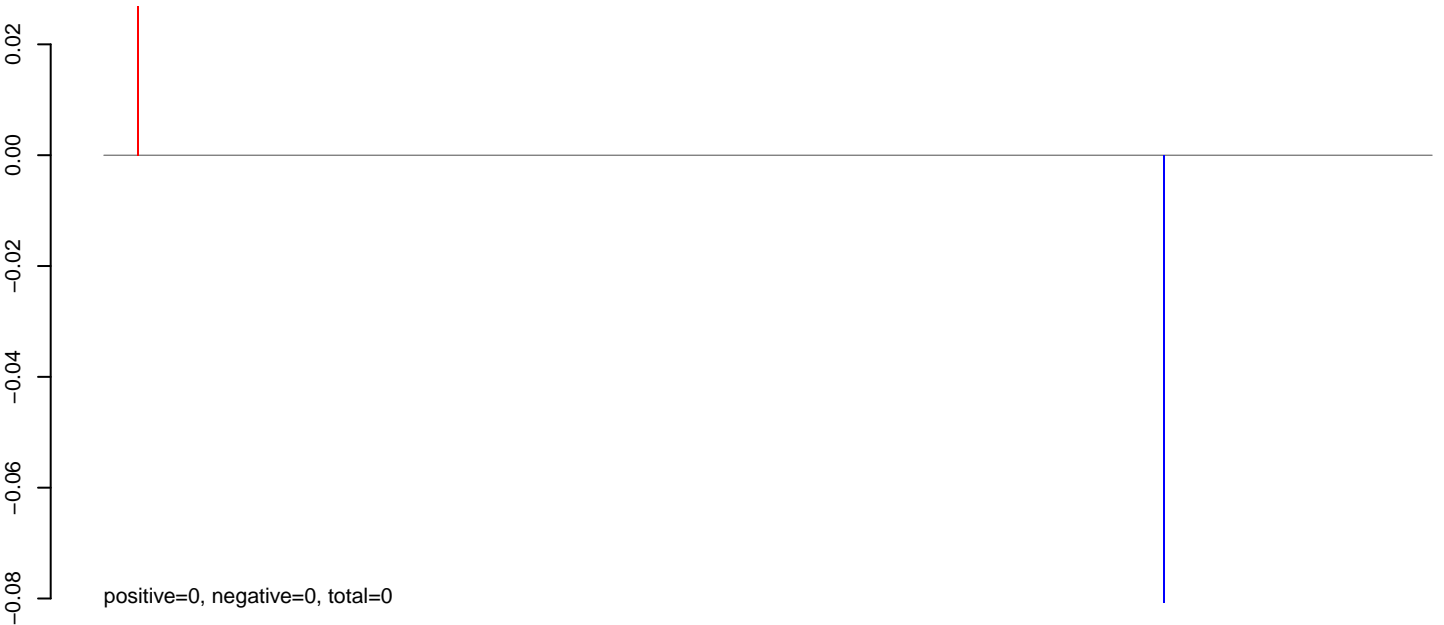


AeAeg_Aag2DENV2NGC.rep

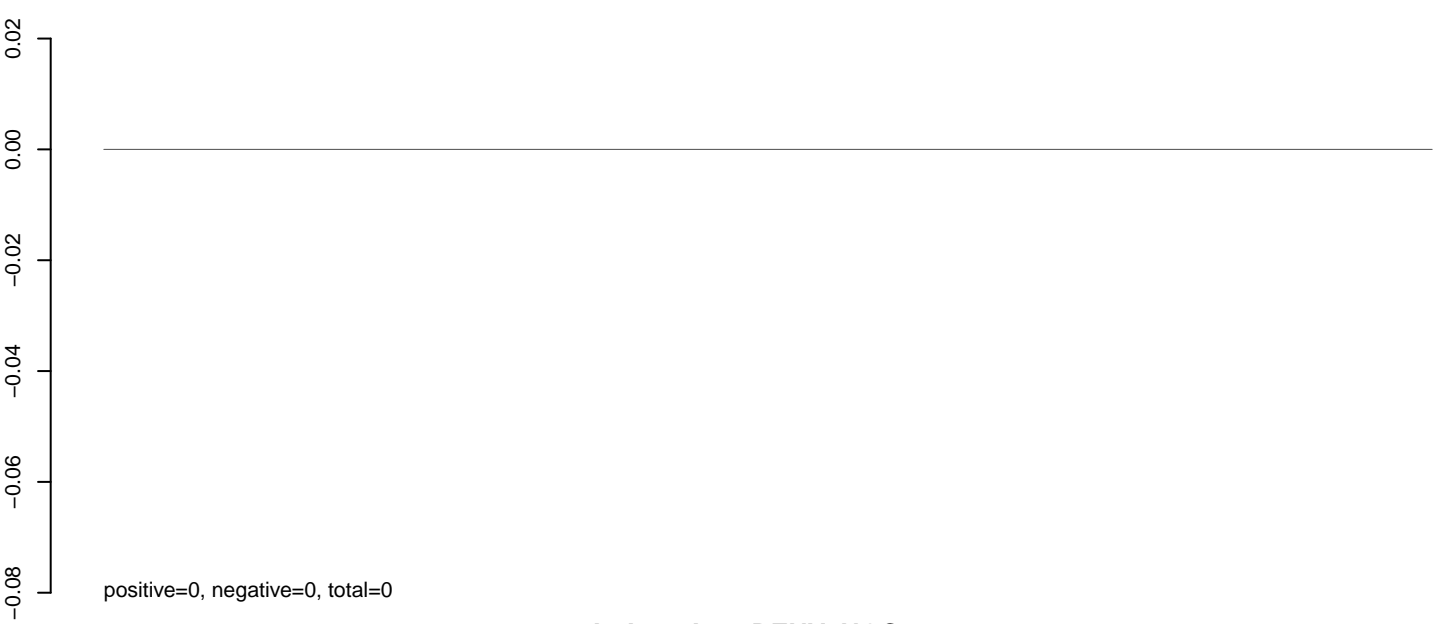


0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



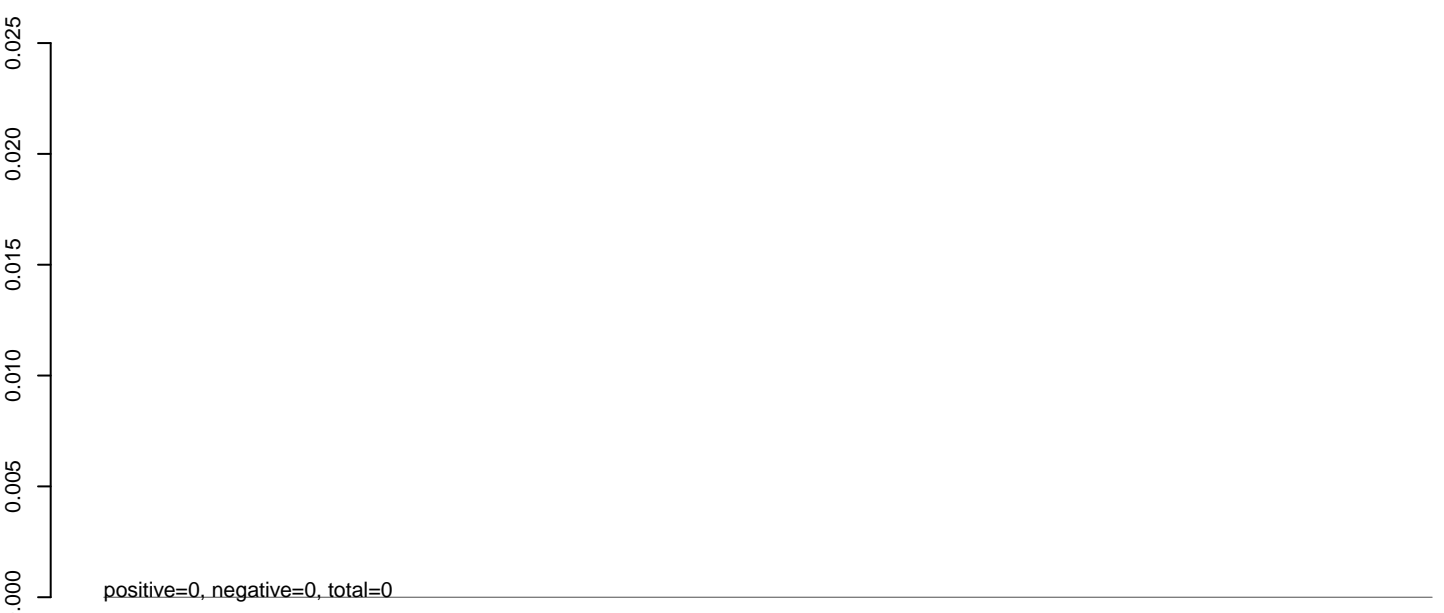
AeAeg_Aag2DENV2NGC.rep



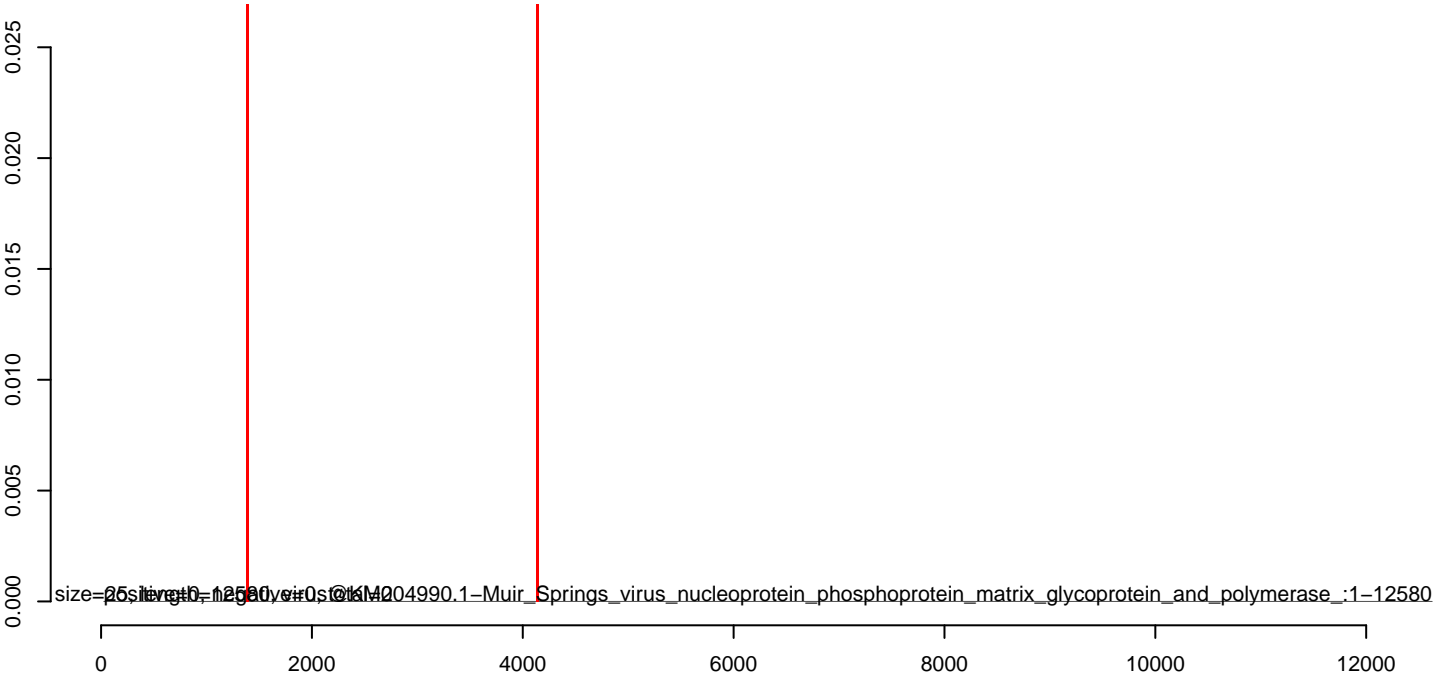
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



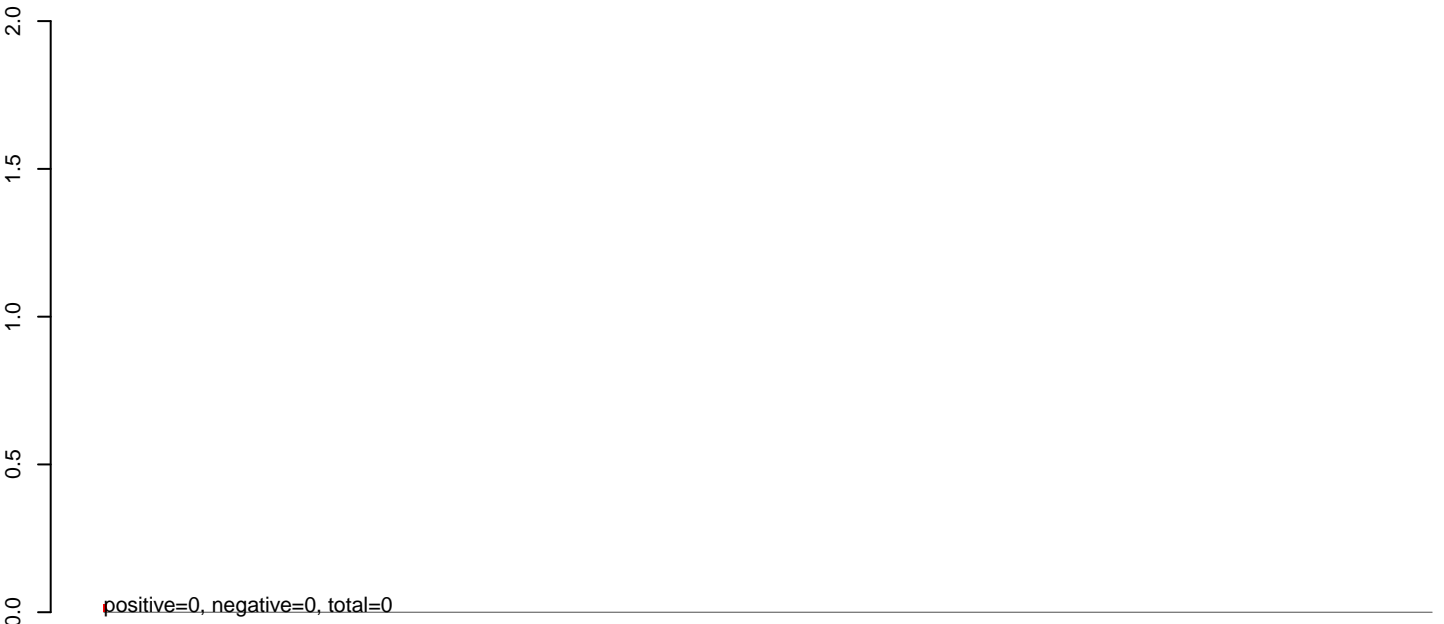
AeAeg_Aag2DENV2NGC.rep



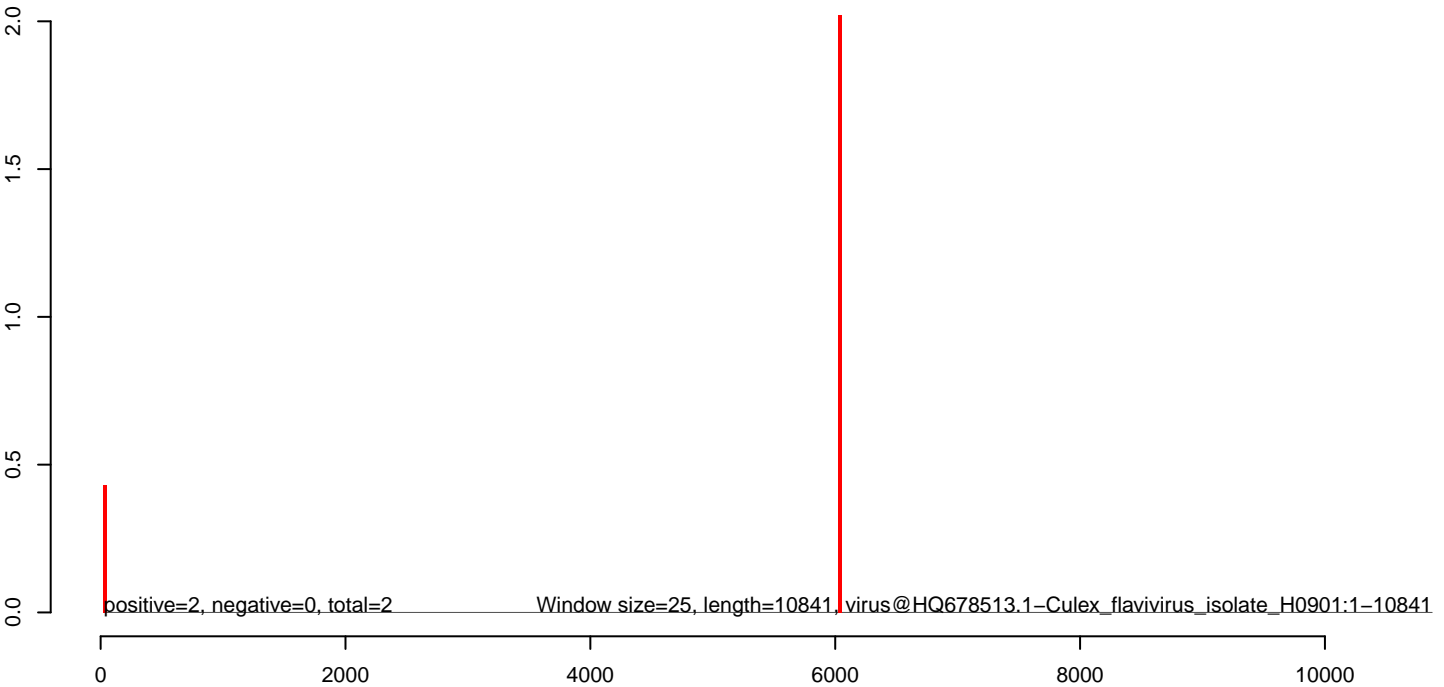
AeAeg_Aag2DENV2NGC.18_23.rep



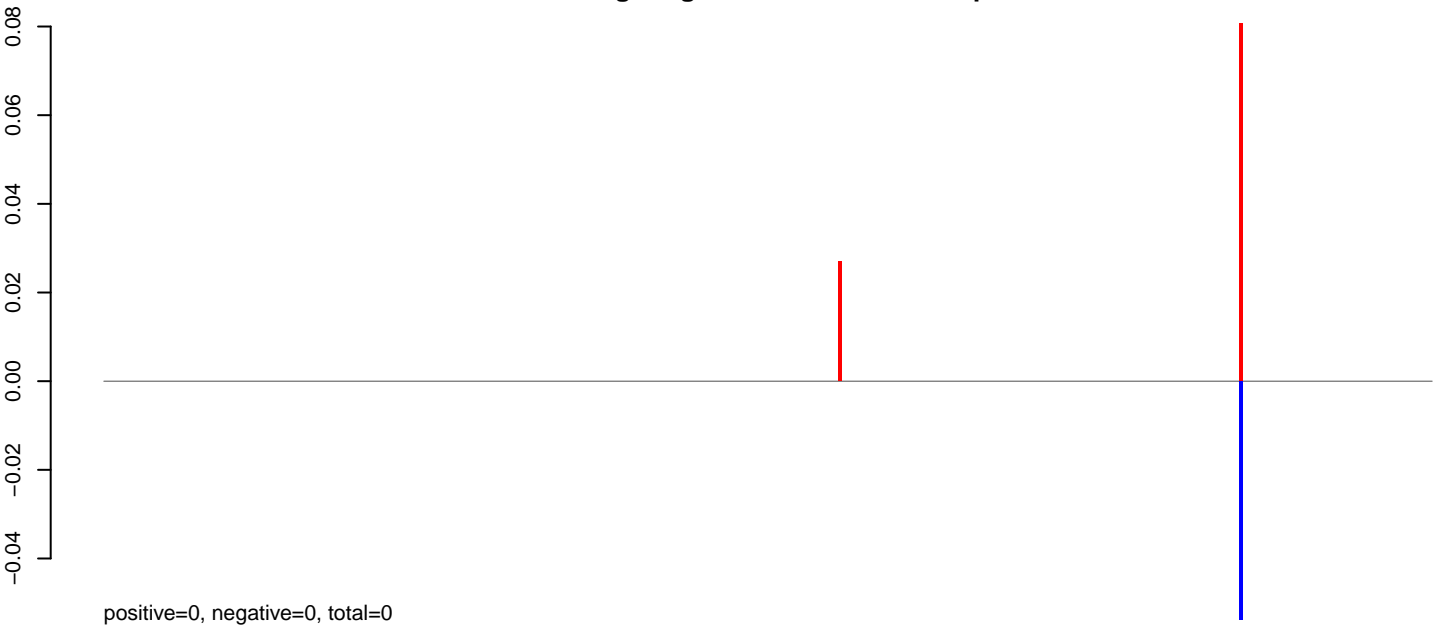
AeAeg_Aag2DENV2NGC.24_35.rep



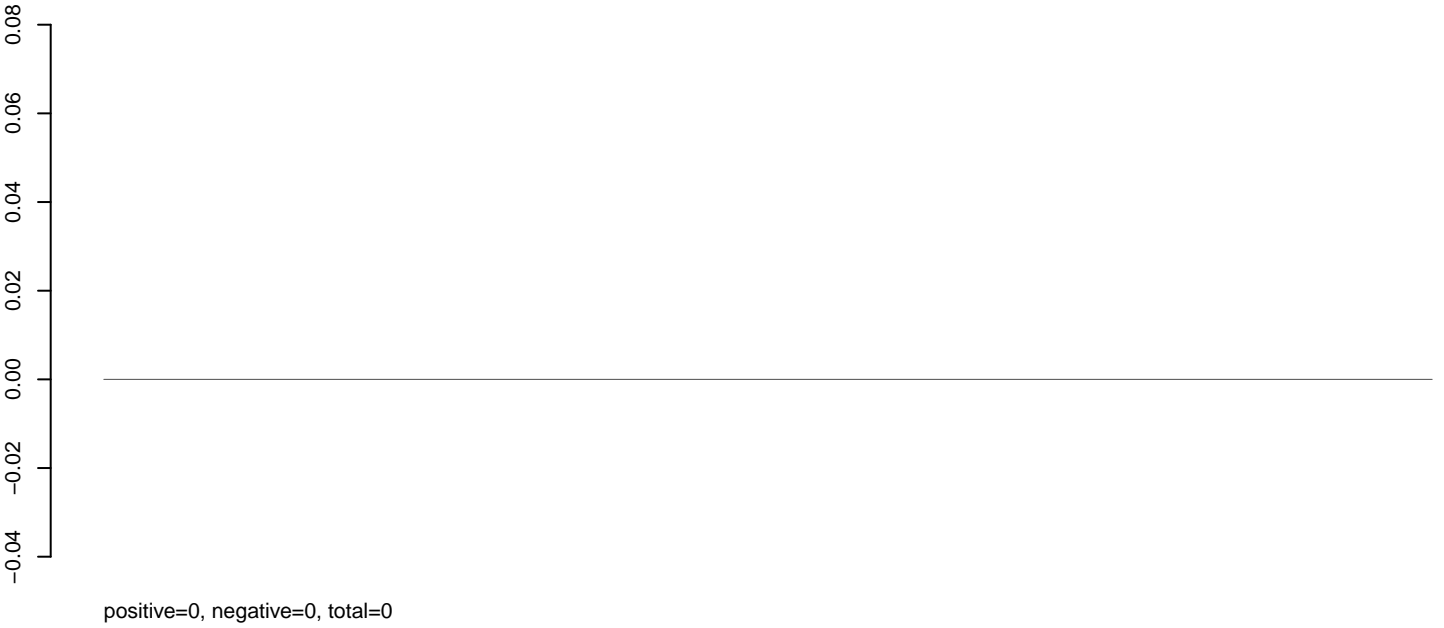
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



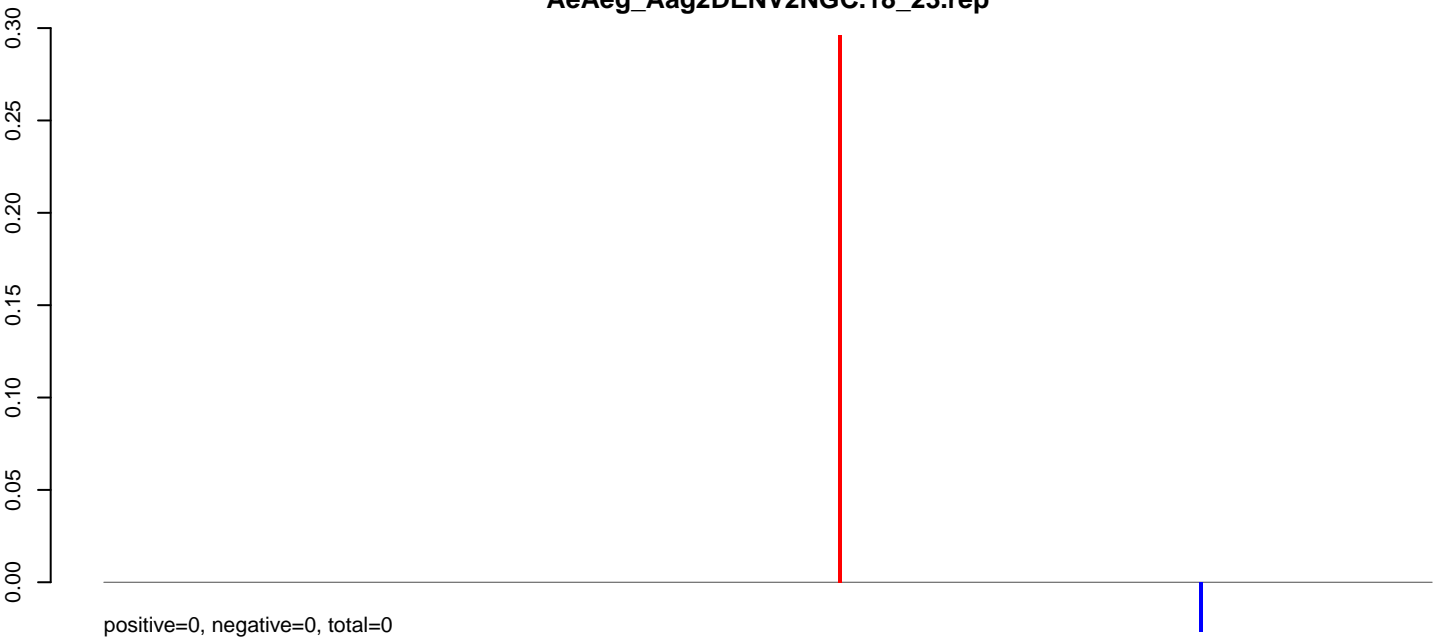
AeAeg_Aag2DENV2NGC.rep



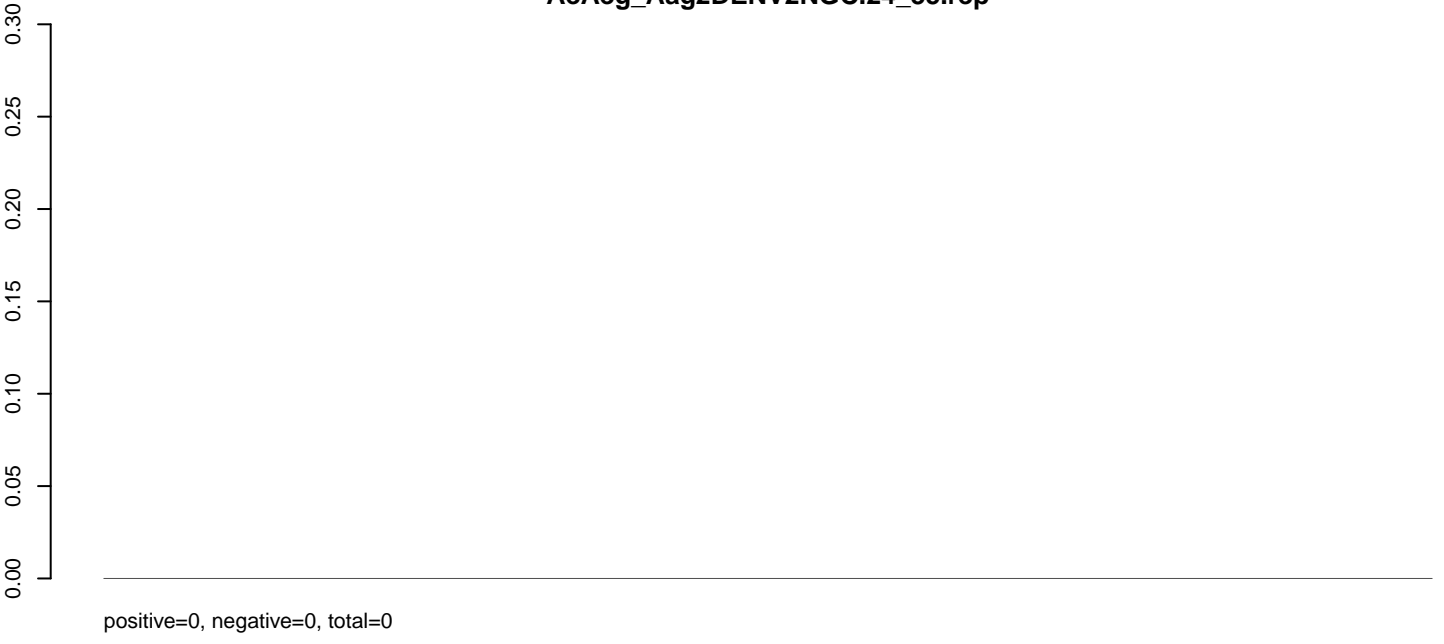
Window size=25, length=10837, virus@FJ663034.1-Culex_flavivirus_isolate_lowa07:1-10837

0 2000 4000 6000 8000 10000

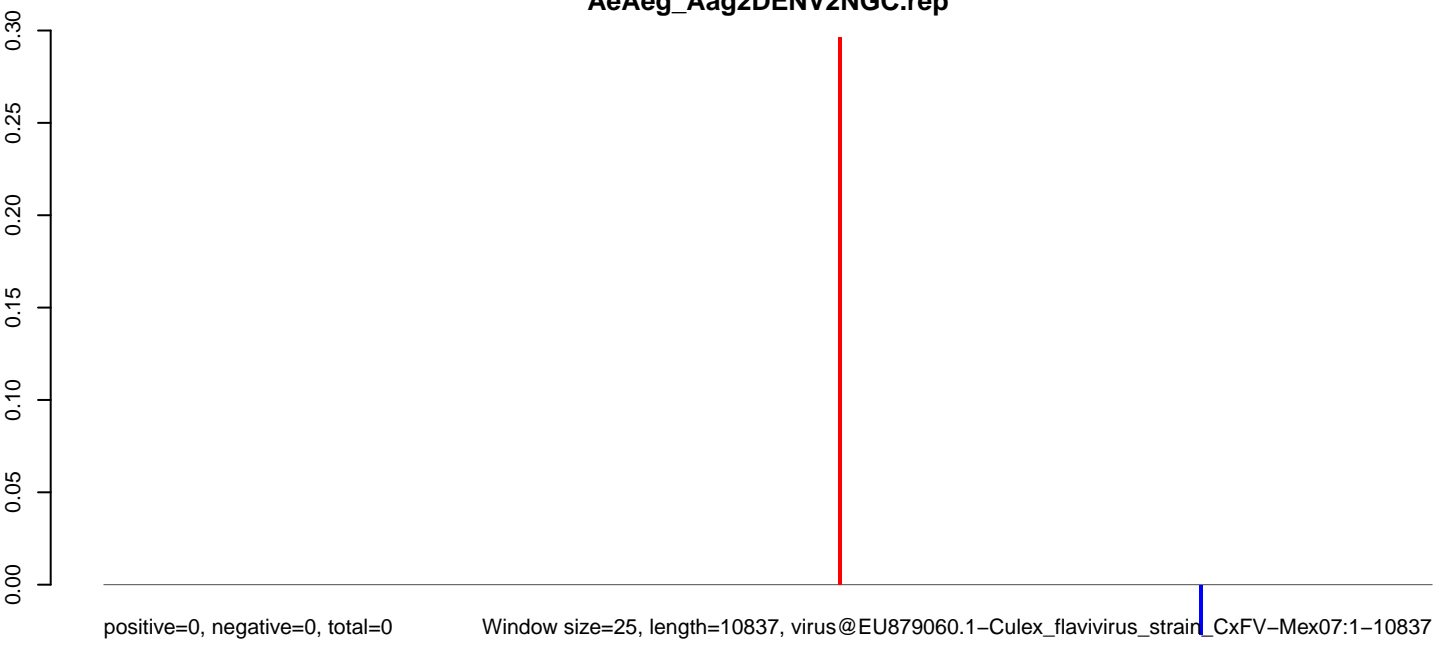
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



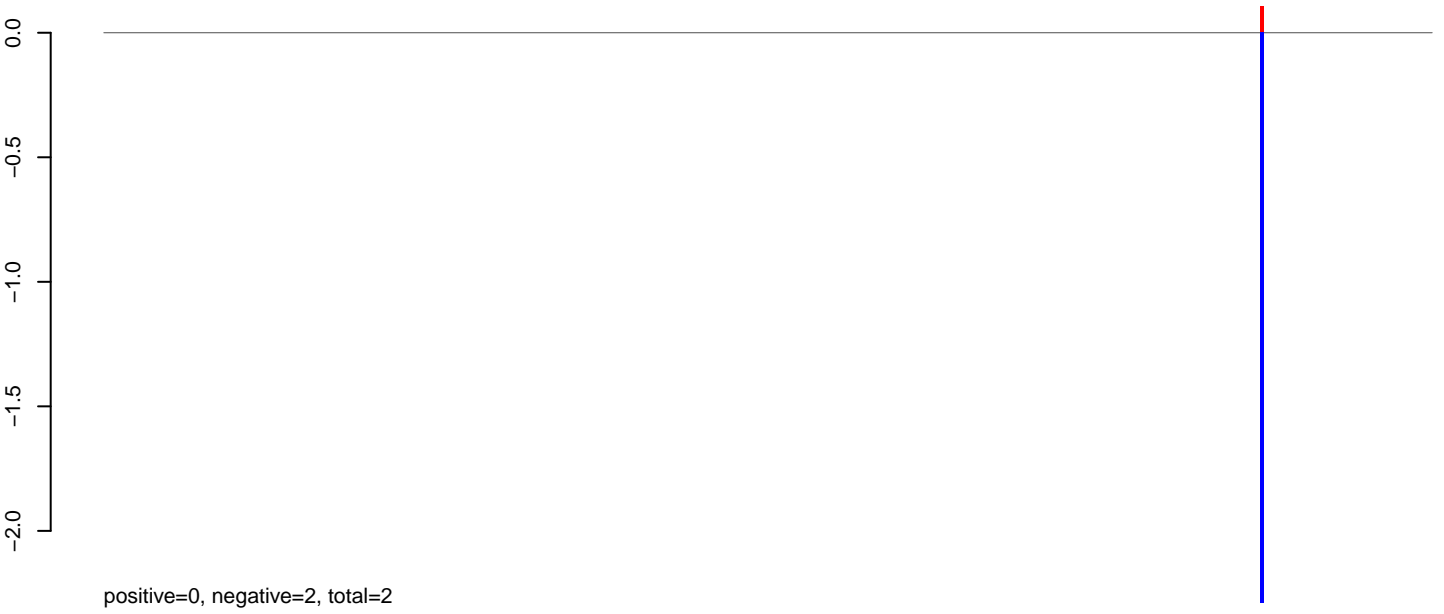
AeAeg_Aag2DENV2NGC.rep



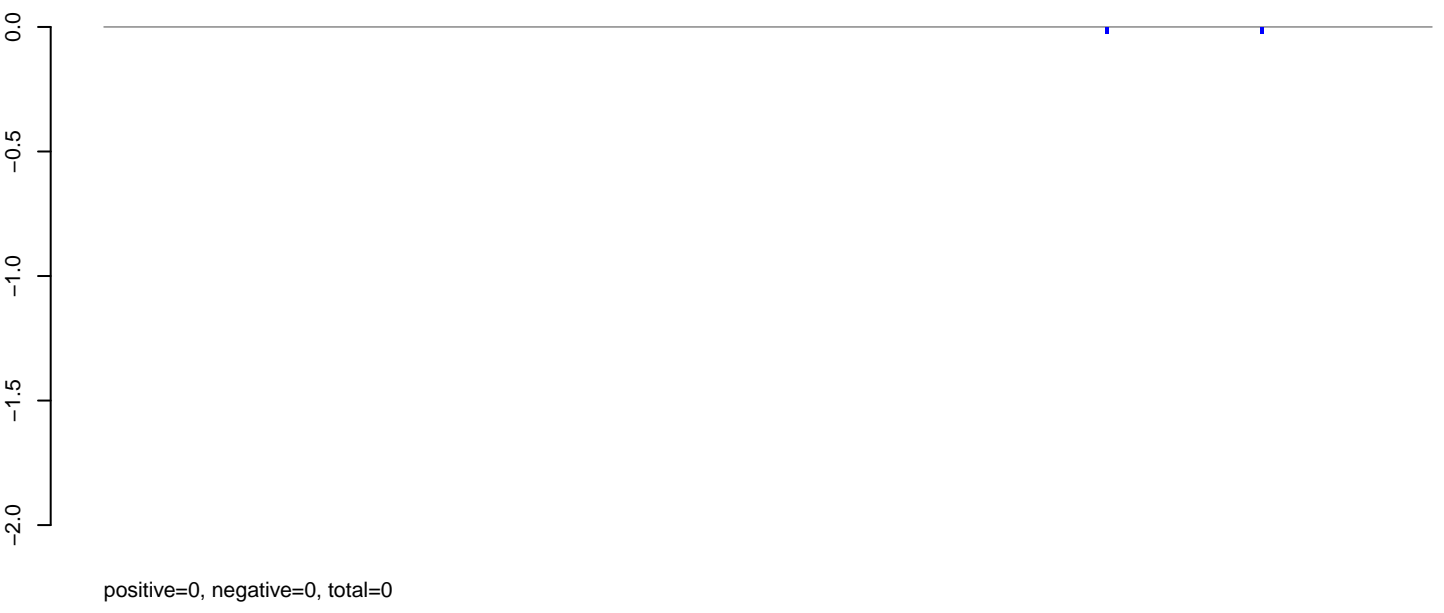
Window size=25, length=10837, virus@EU879060.1-Culex_flavivirus_strain_CxFV-Mex07:1-10837

0 2000 4000 6000 8000 10000

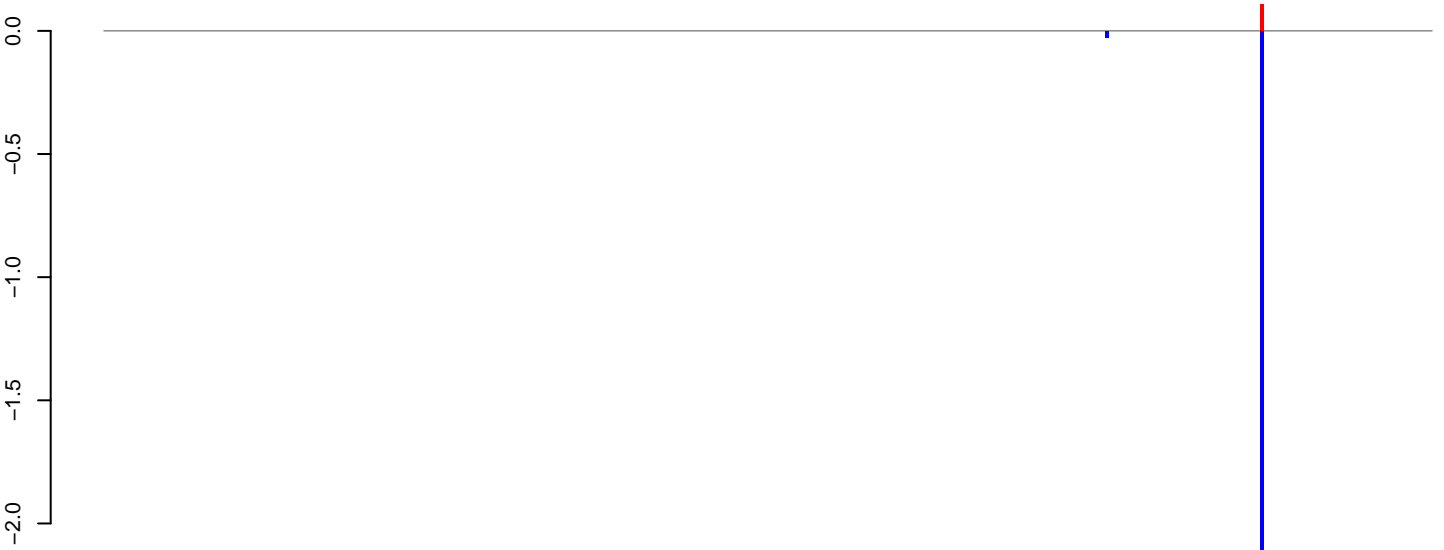
AeAeg_Aag2DENV2NGC.18_23.rep



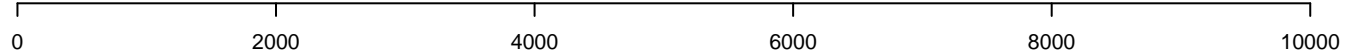
AeAeg_Aag2DENV2NGC.24_35.rep



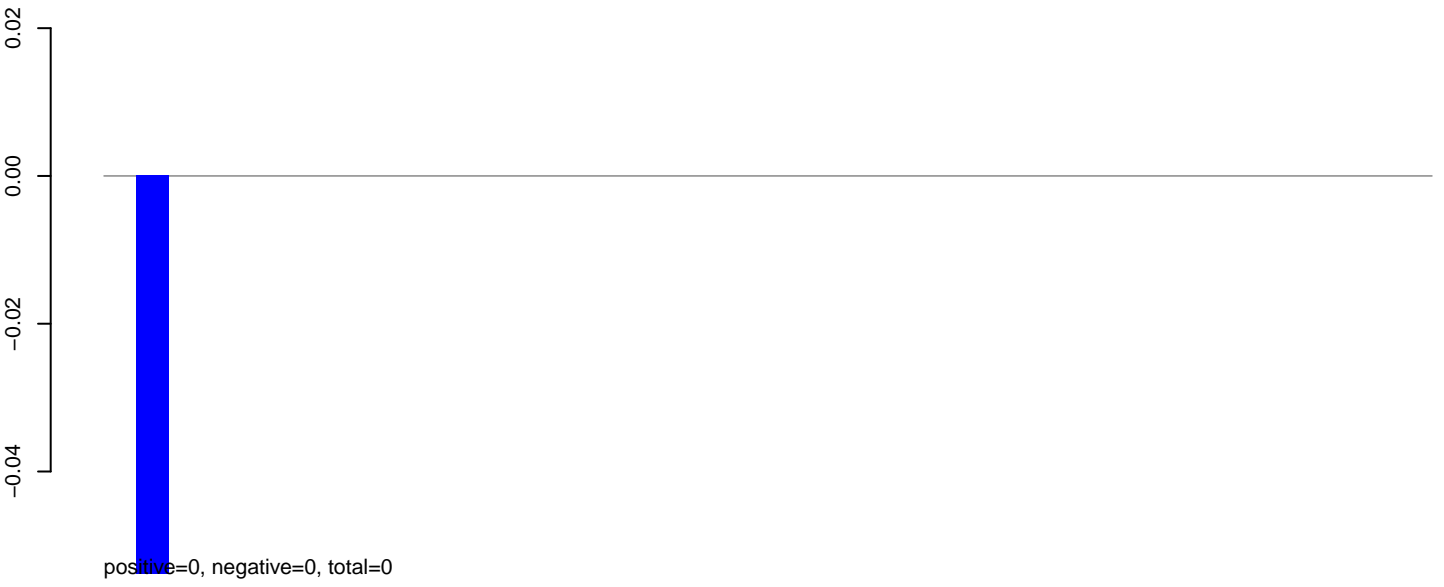
AeAeg_Aag2DENV2NGC.rep



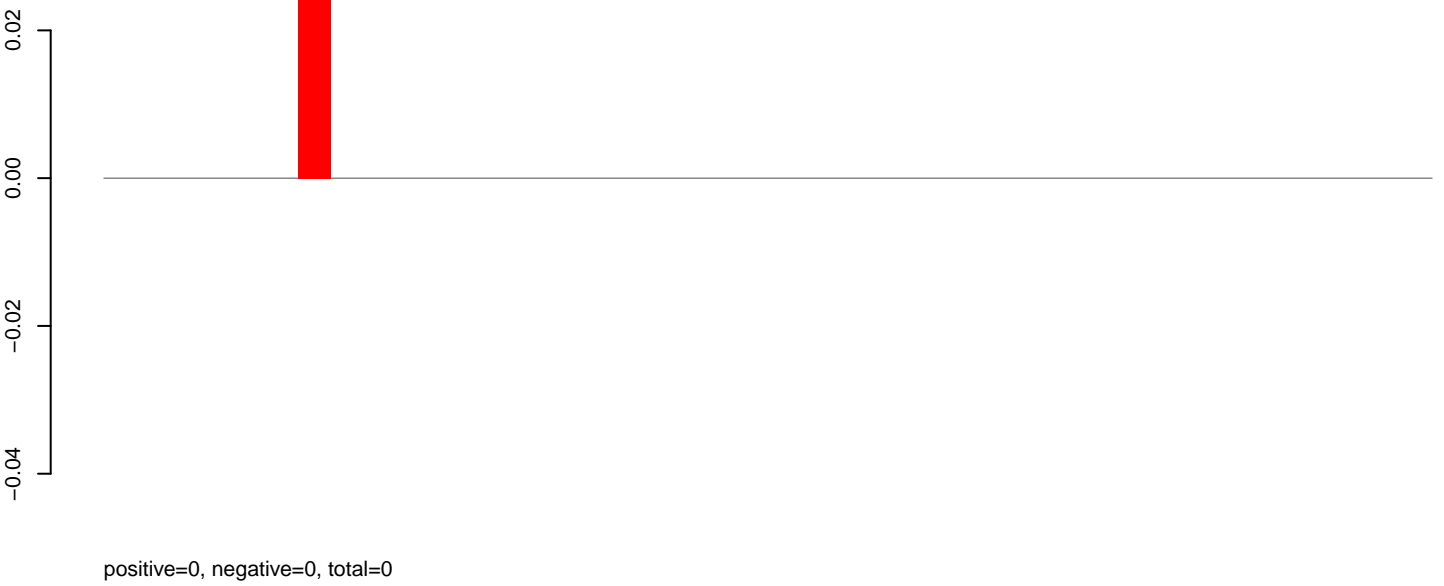
Window size=25, length=10254, virus@DQ235149.1-Royal_Farm_virus_from_Afghanistan_polyprotein_gene:1-10254



AeAeg_Aag2DENV2NGC.18_23.rep



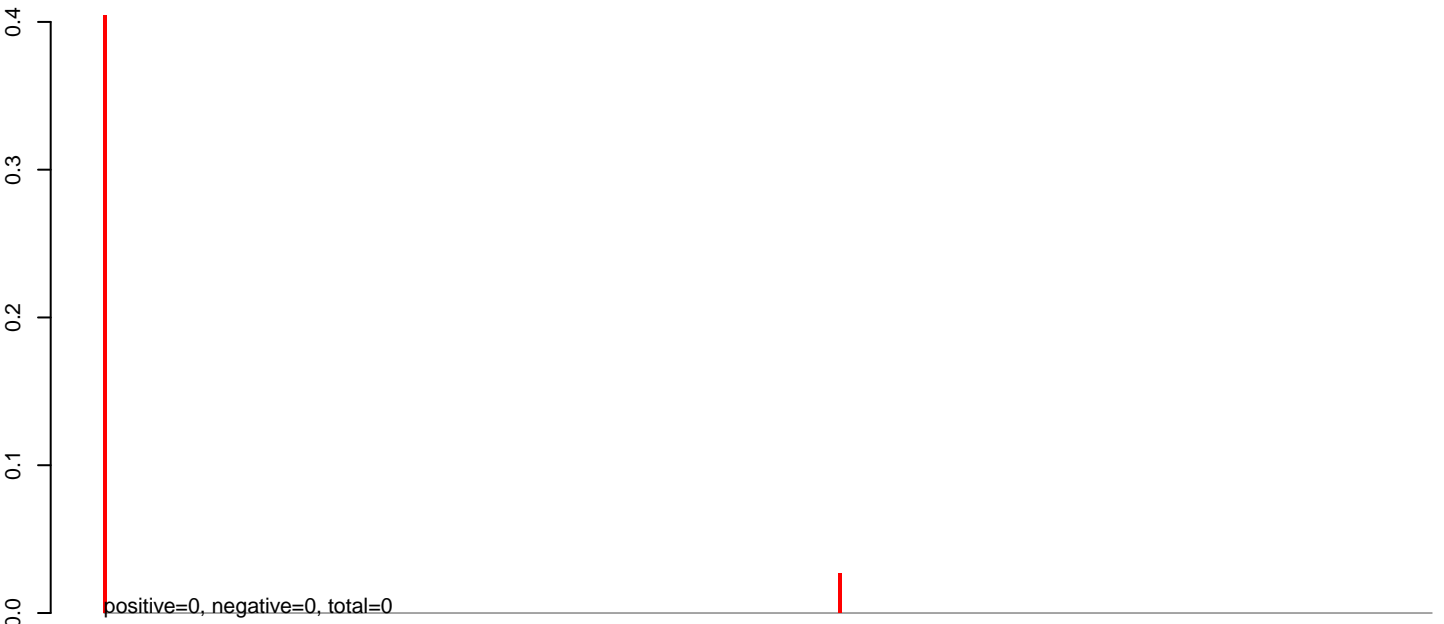
AeAeg_Aag2DENV2NGC.24_35.rep



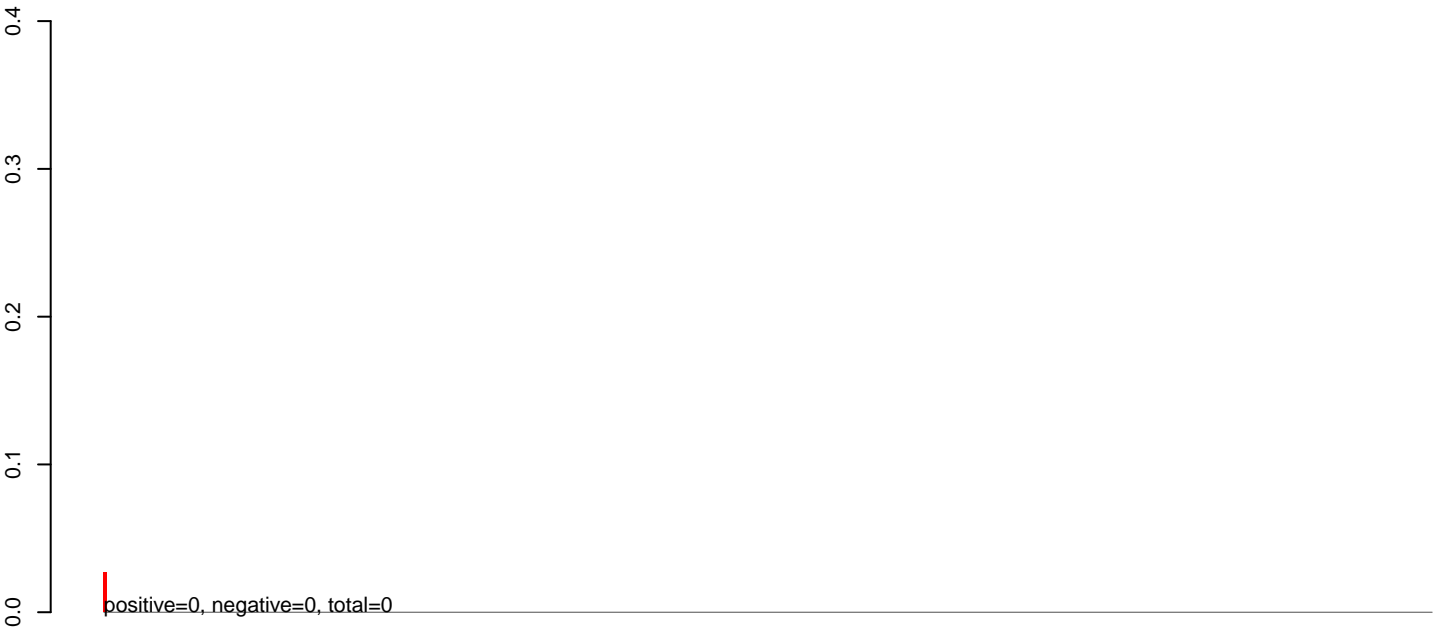
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



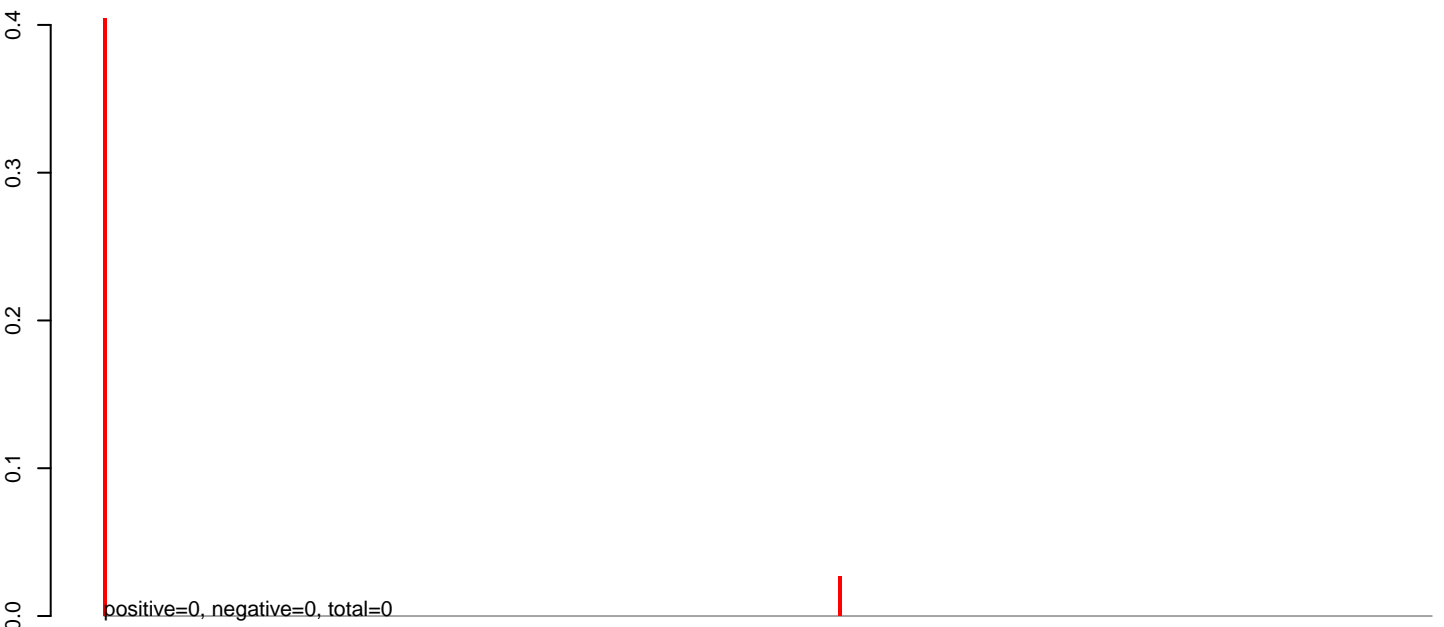
AeAeg_Aag2DENV2NGC.24_35.rep



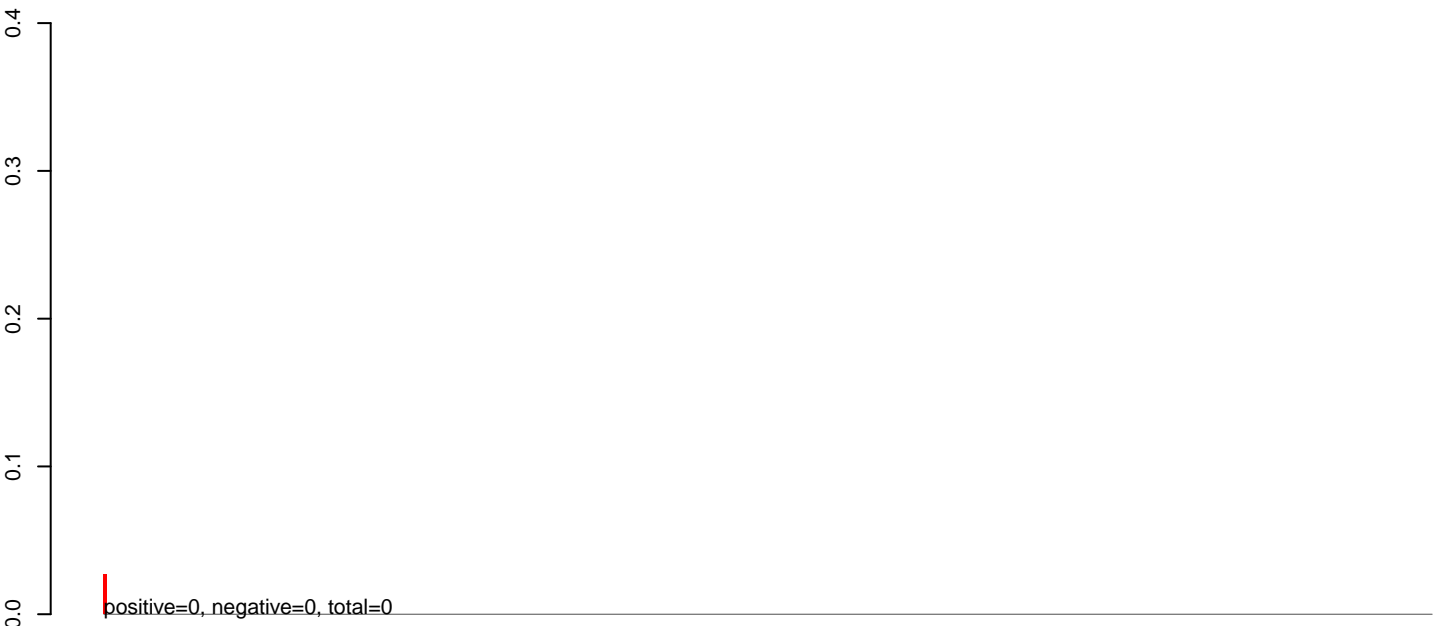
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



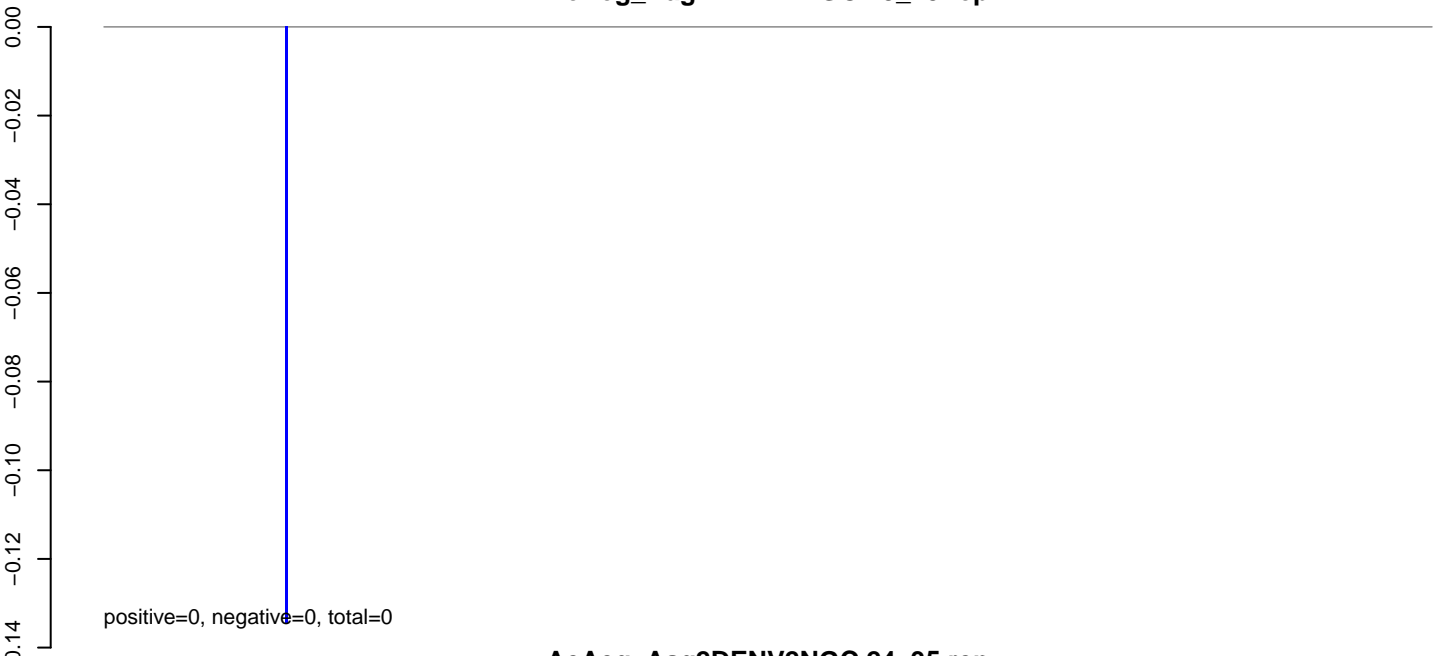
AeAeg_Aag2DENV2NGC.24_35.rep



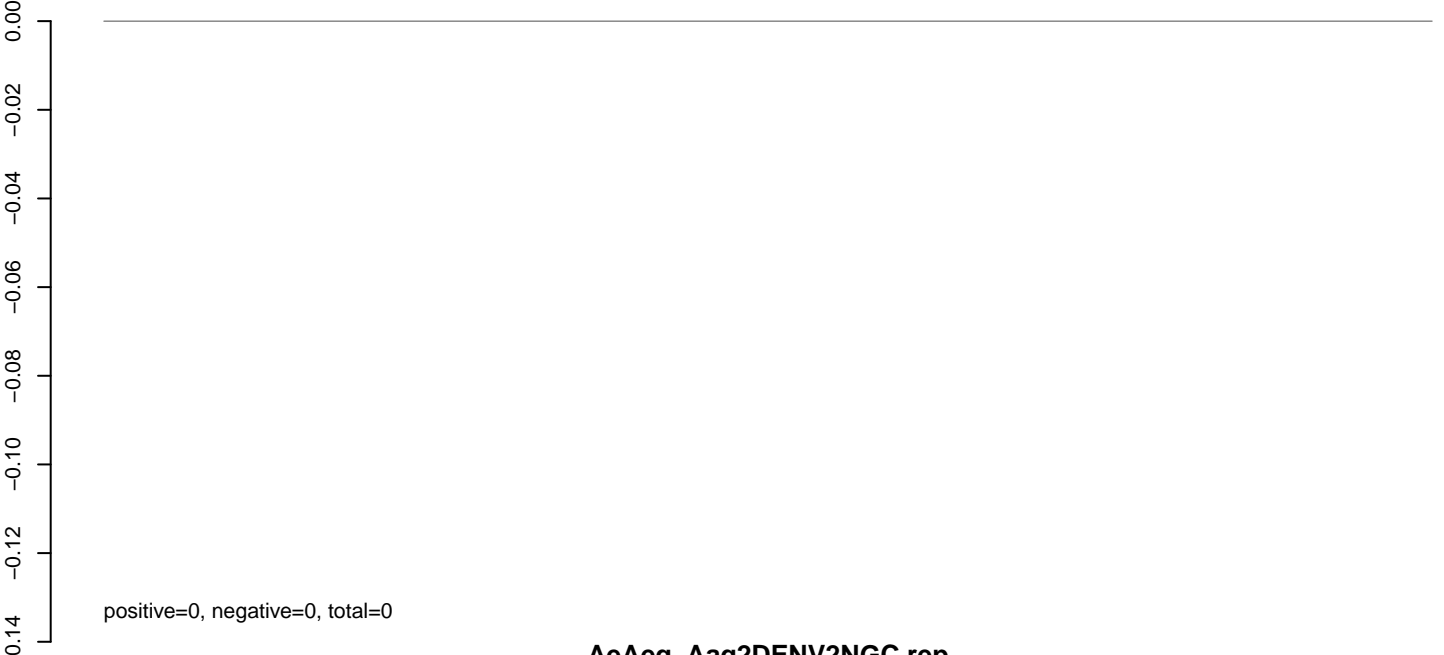
AeAeg_Aag2DENV2NGC.rep



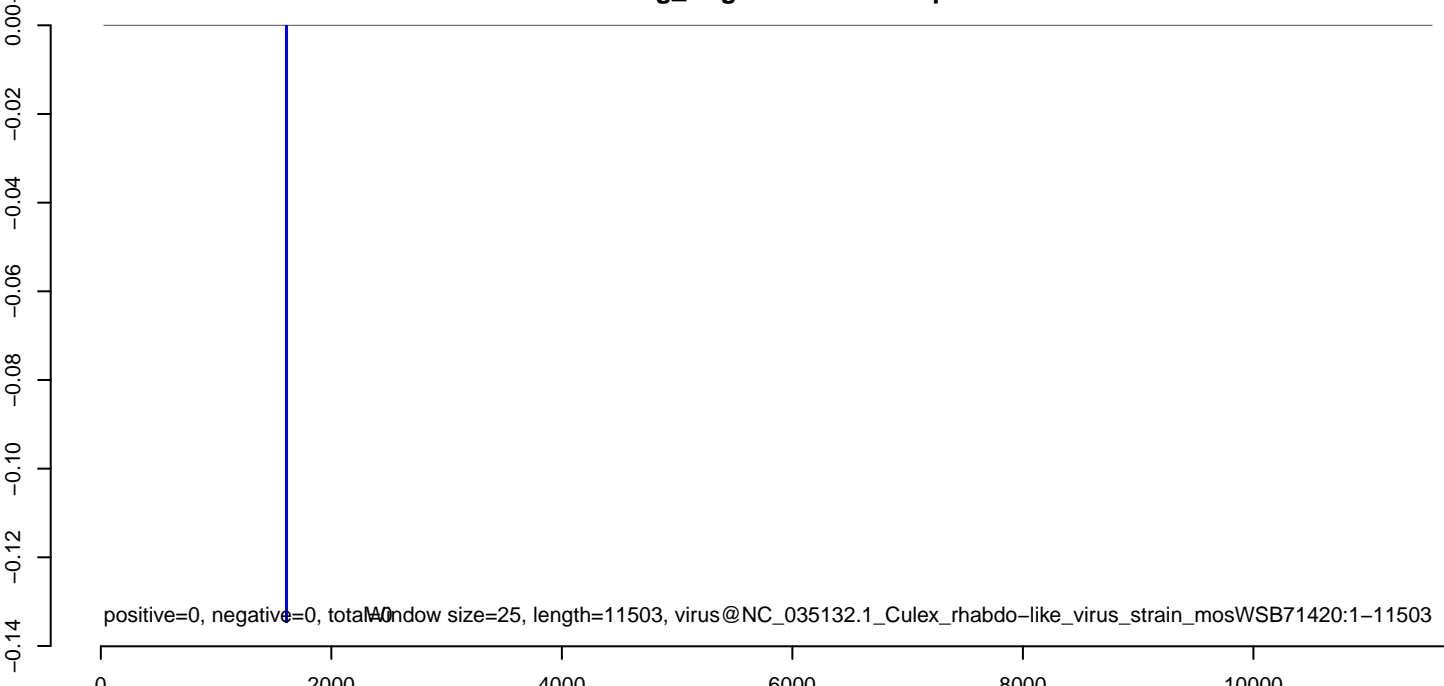
AeAeg_Aag2DENV2NGC.18_23.rep



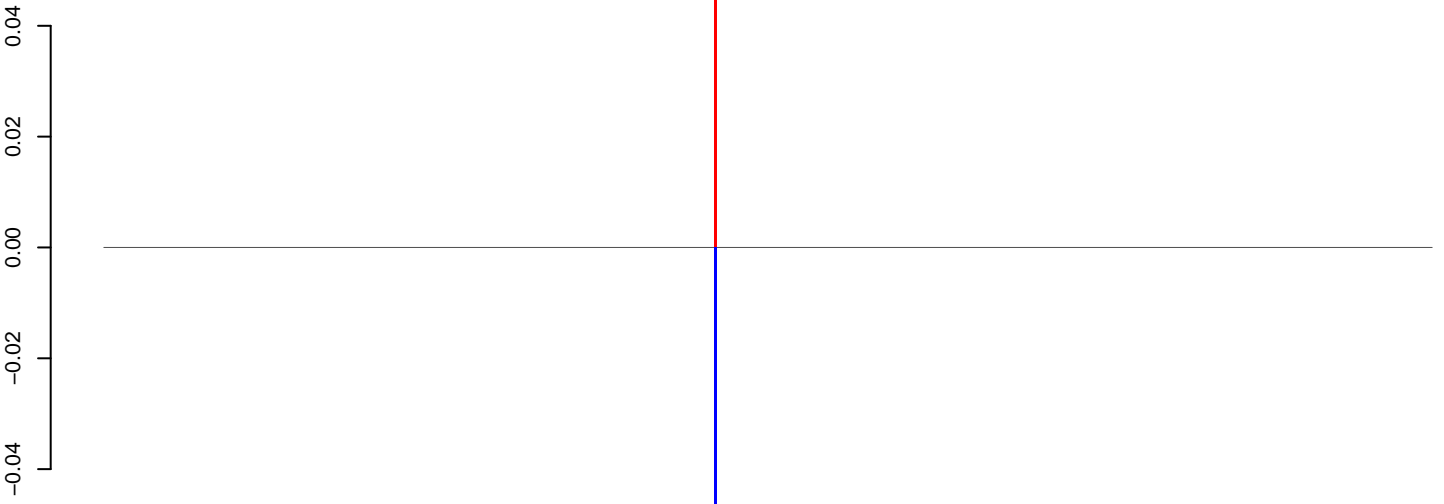
AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep

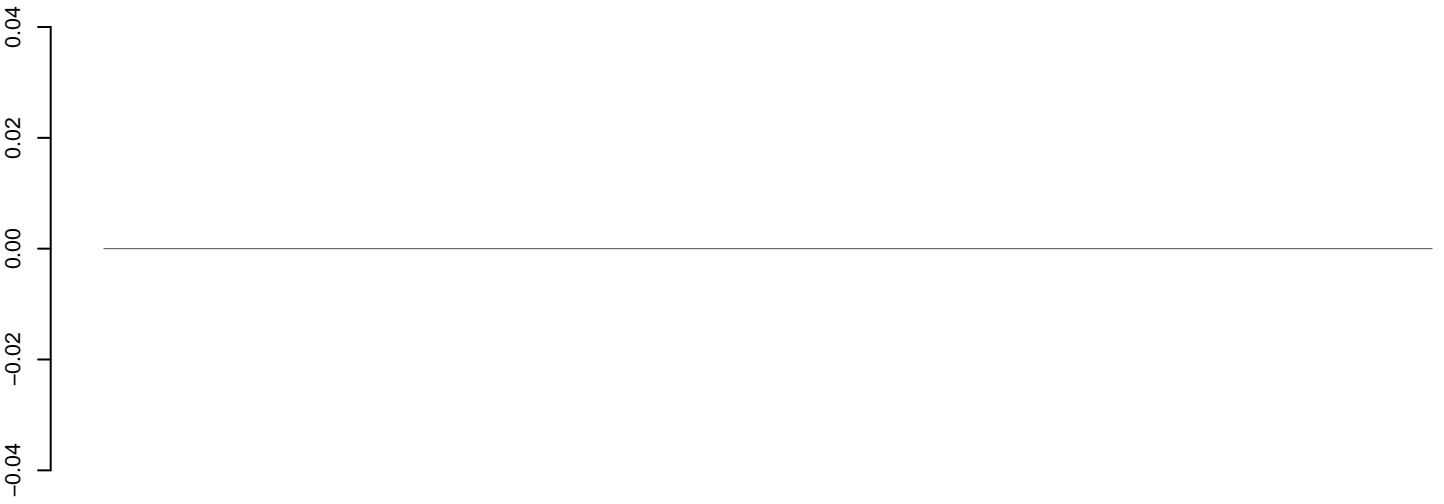


AeAeg_Aag2DENV2NGC.18_23.rep



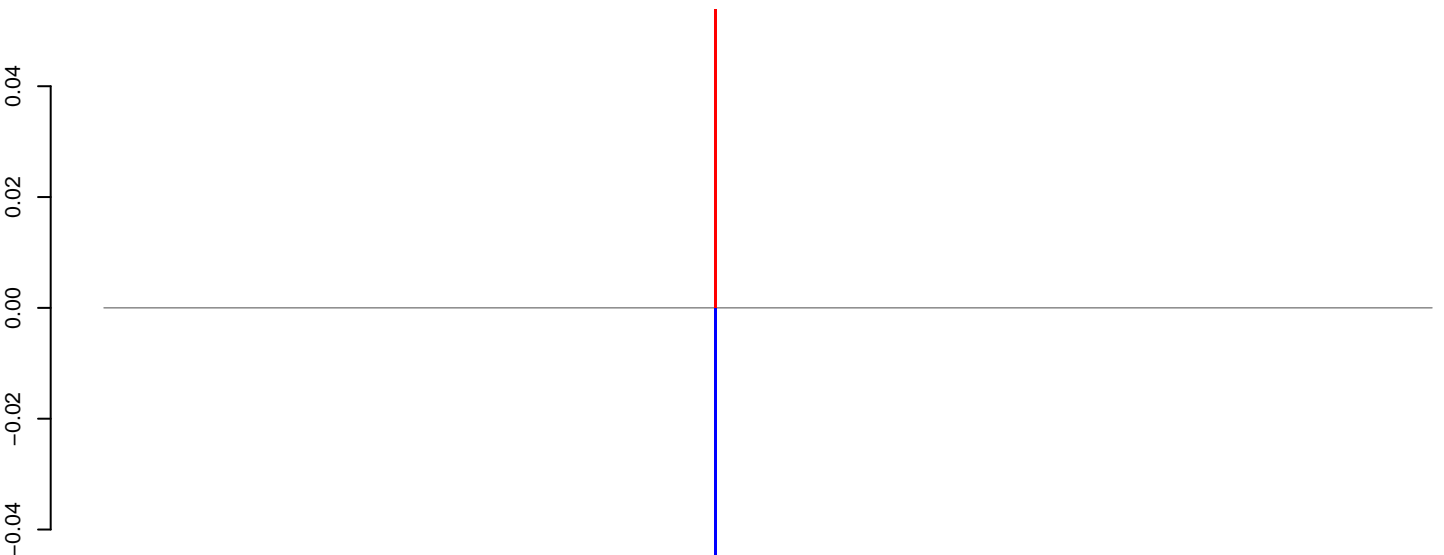
positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.rep



positive=0, negative=0, total=0, virus@NC_035128.1_Aedes_camptorhynchus_negev-like_virus_strain_mos191CP47896:1-11470

0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



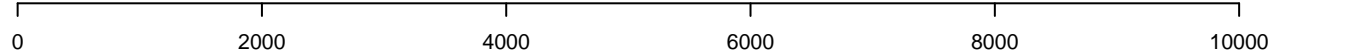
AeAeg_Aag2DENV2NGC.24_35.rep



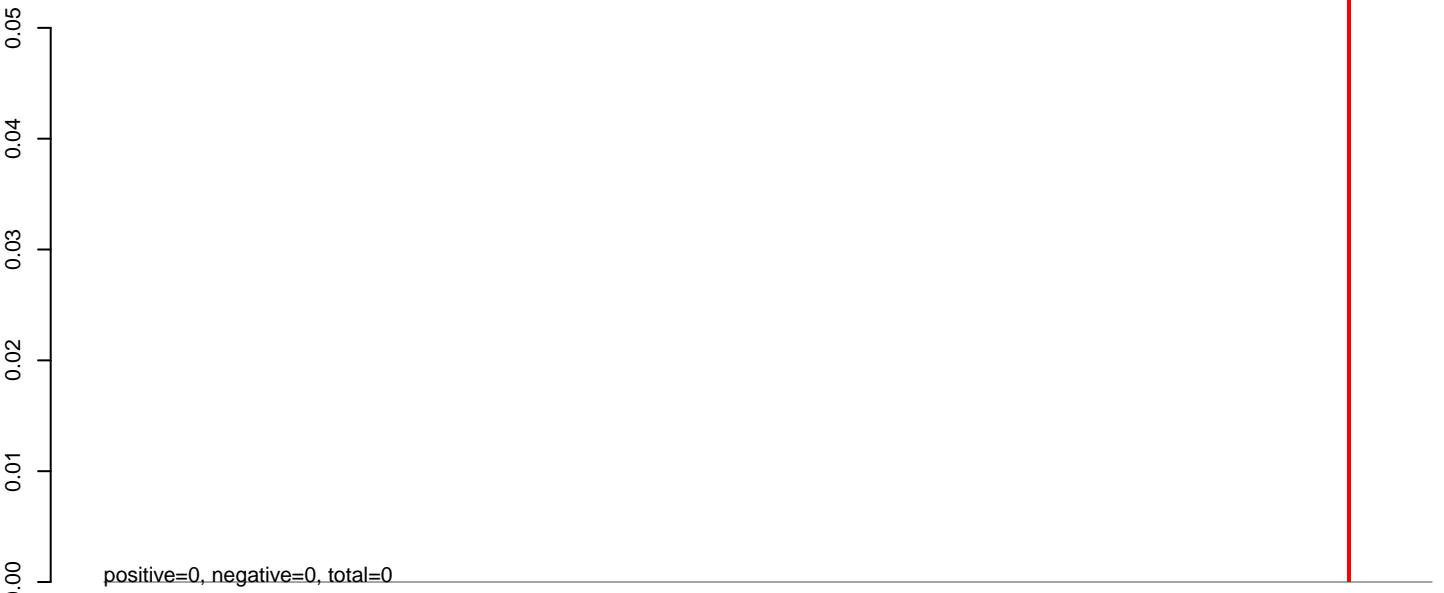
AeAeg_Aag2DENV2NGC.rep



Window size=25, length=10867, virus@NC_035124.1_Culex_negev-like_virus_1_strain_mosWSX37710:1-10867

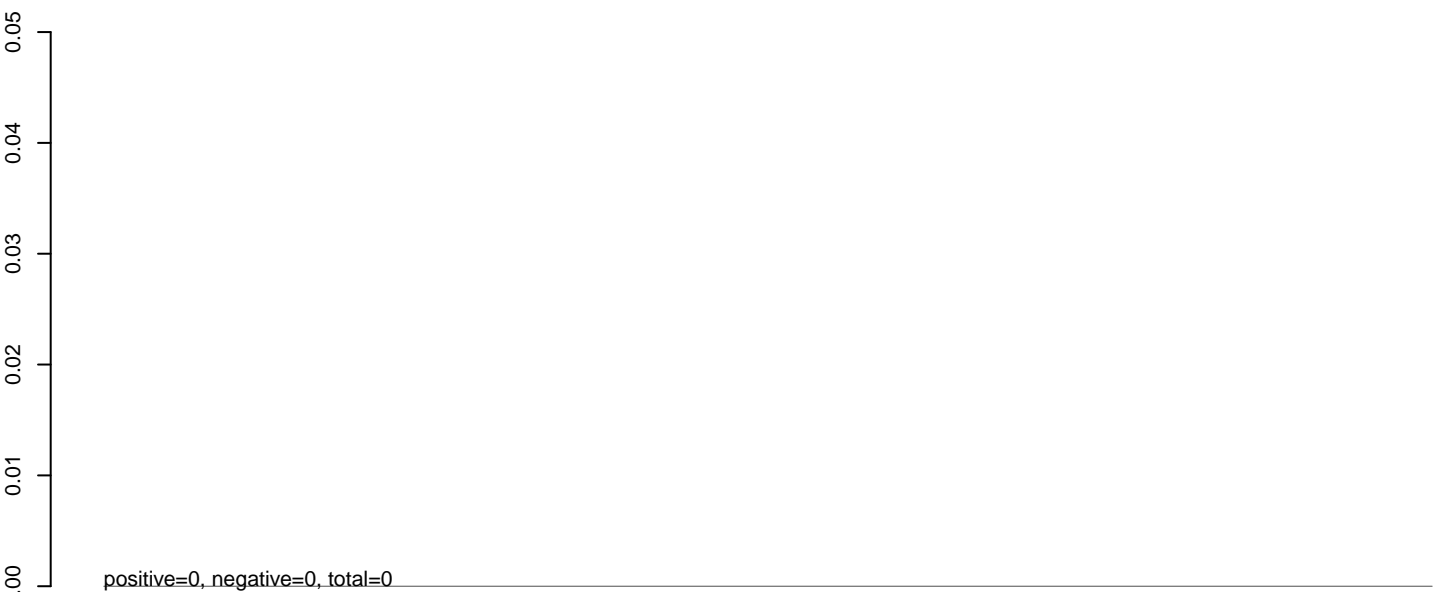


AeAeg_Aag2DENV2NGC.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.24_35.rep



positive=0, negative=0, total=0

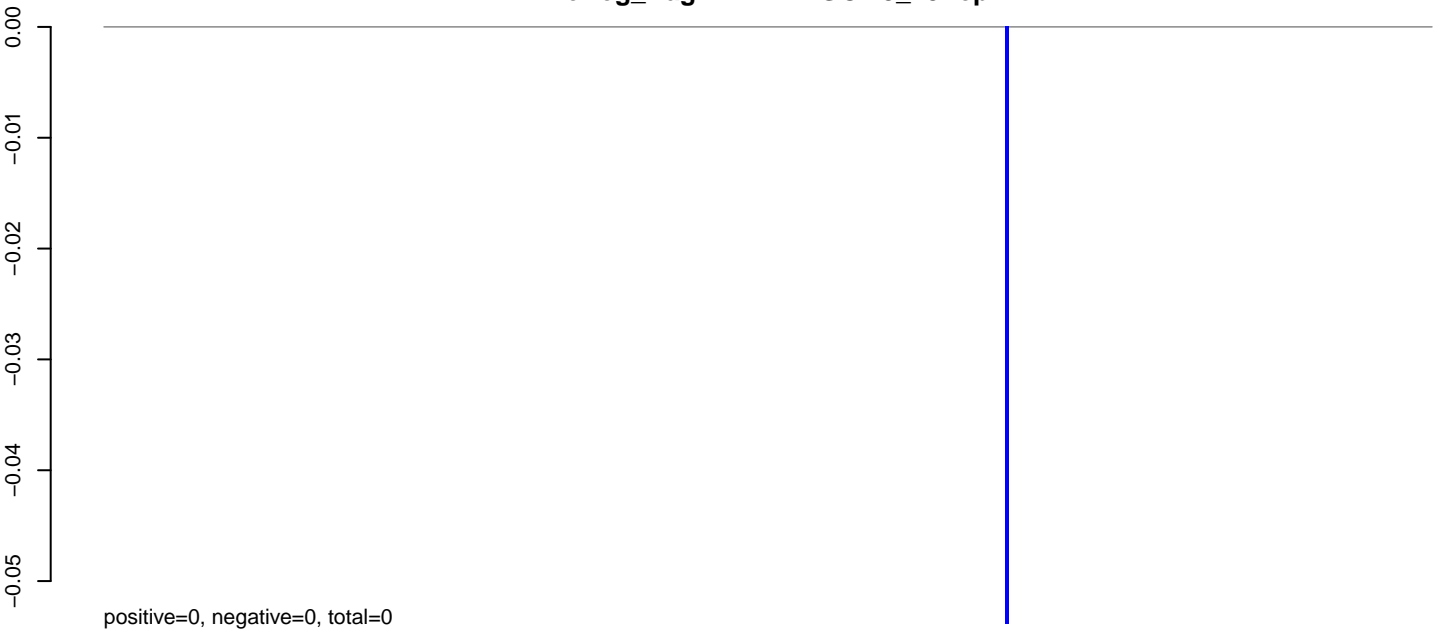
AeAeg_Aag2DENV2NGC.rep



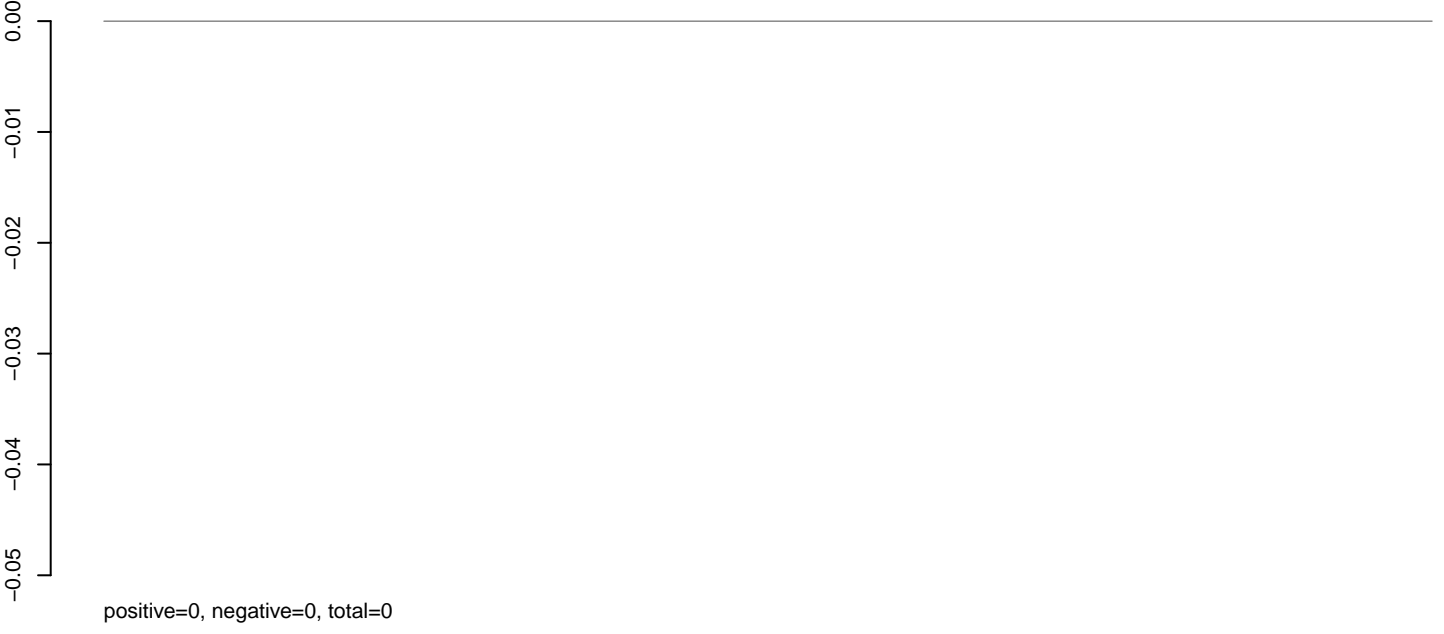
positive=0, negative=0, total=0, window size=25, length=9324, virus@NC_035123.1_Culex_negev-like_virus_2_strain_mos172X30622:1-9324

0 2000 4000 6000 8000

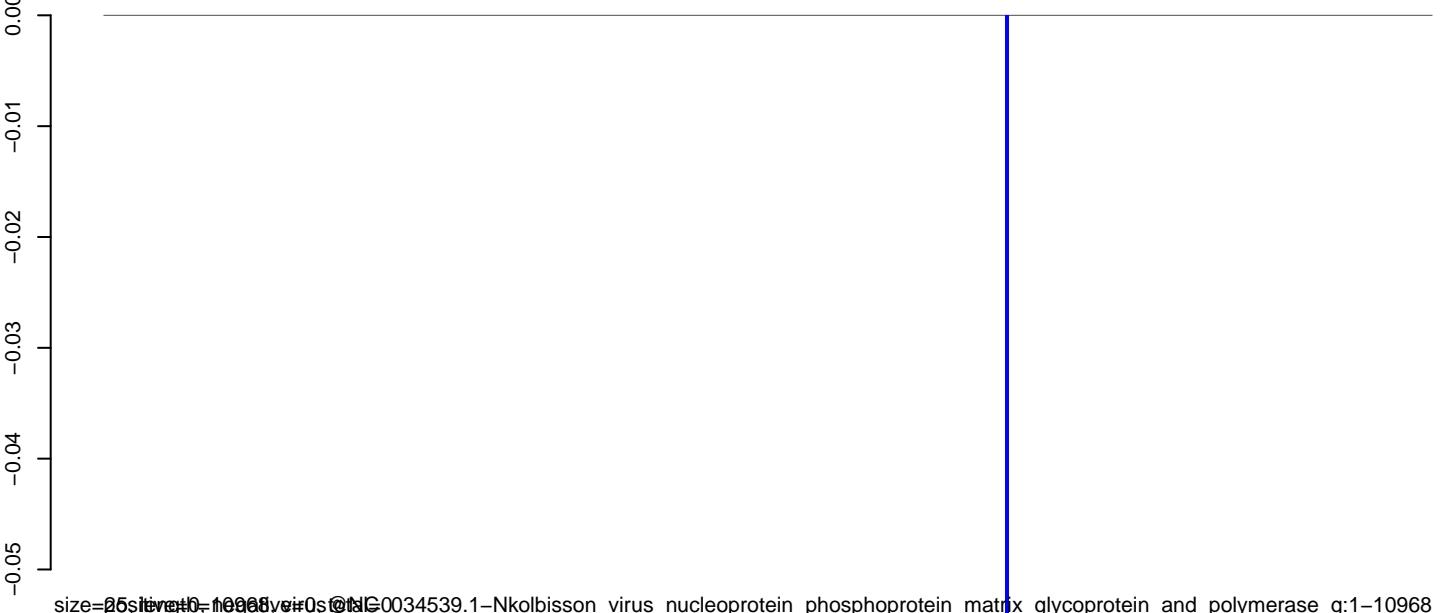
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

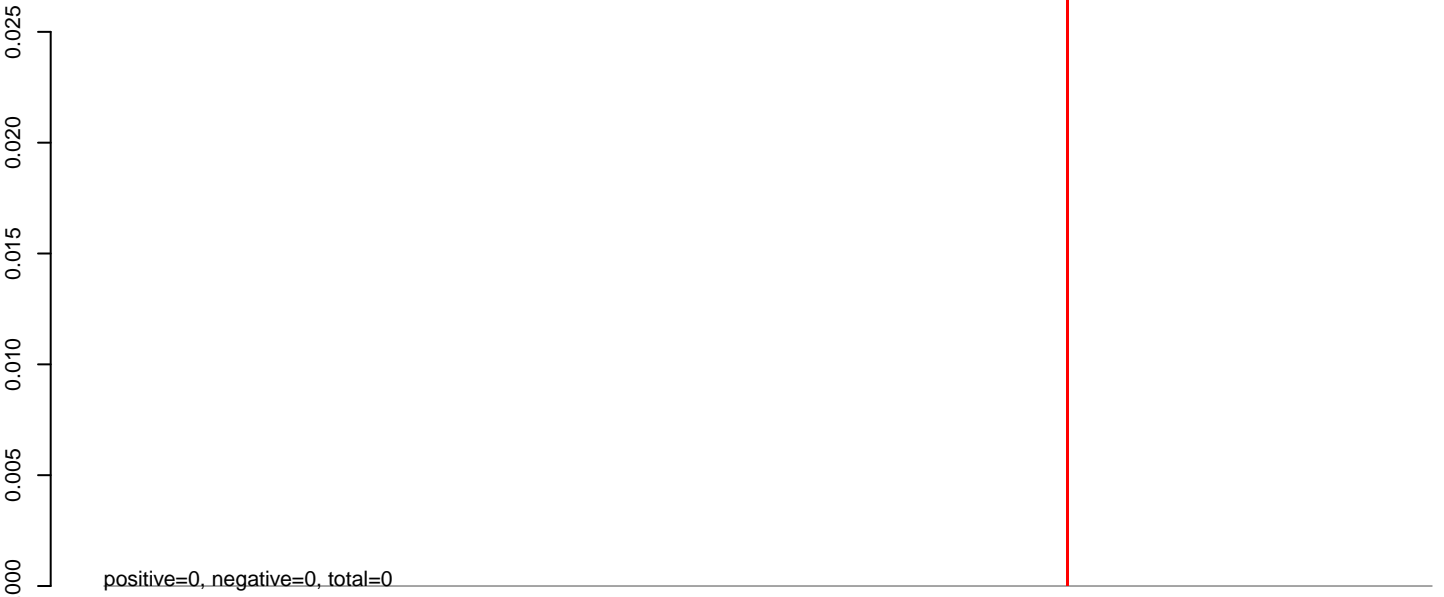


AeAeg_Aag2DENV2NGC.rep

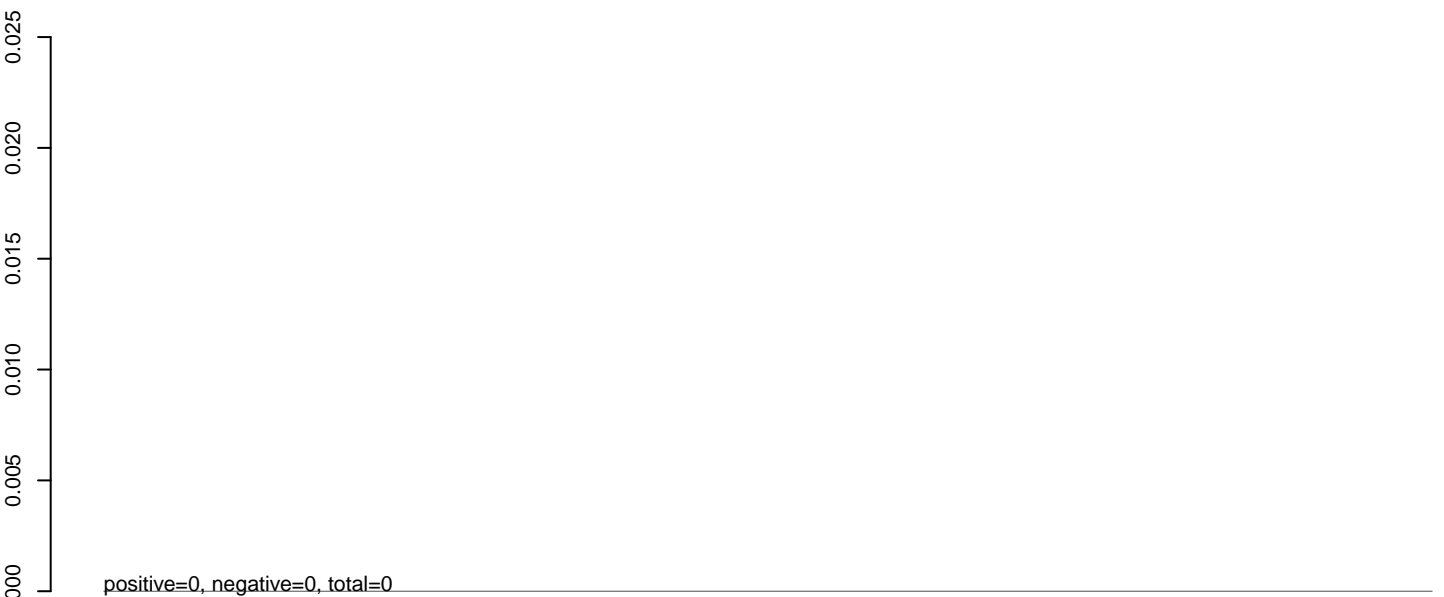


0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



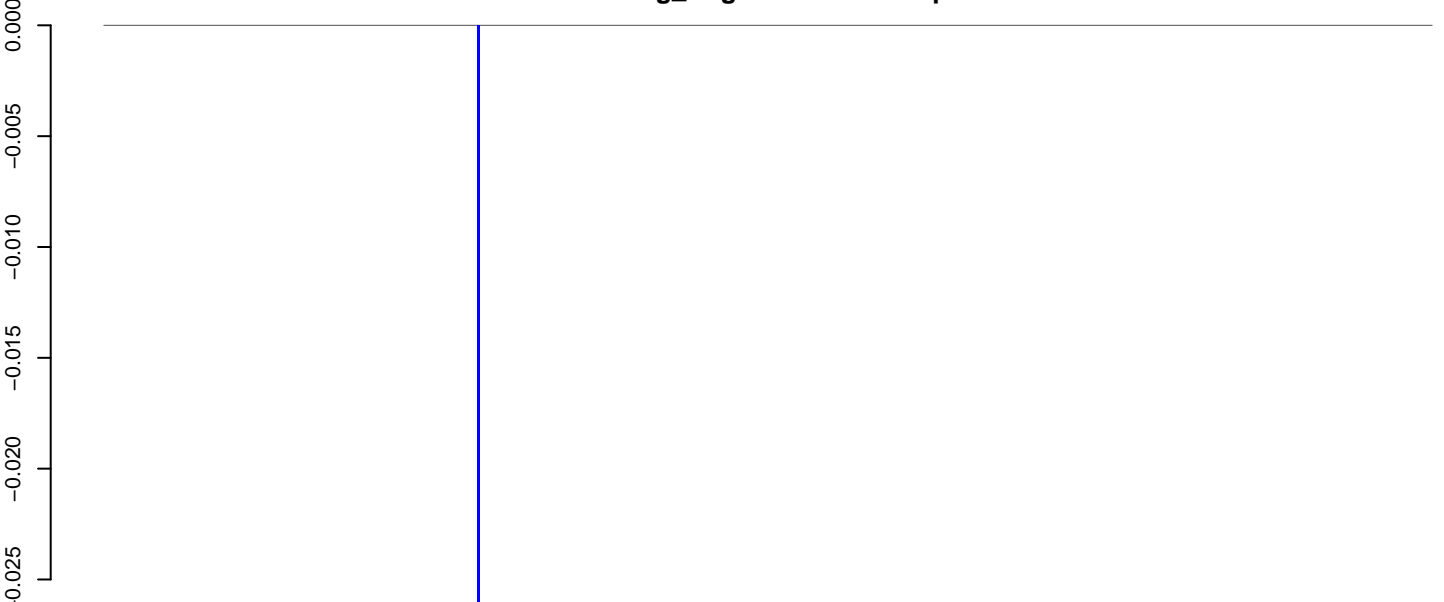
AeAeg_Aag2DENV2NGC.18_23.rep



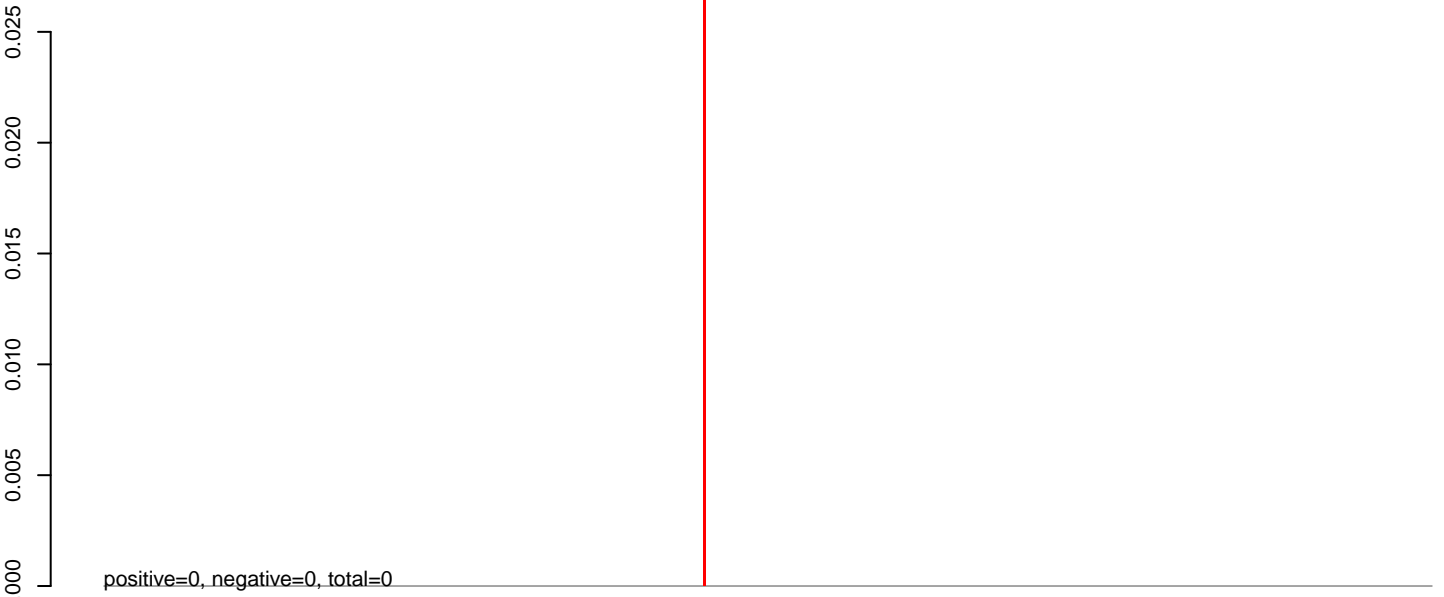
AeAeg_Aag2DENV2NGC.24_35.rep



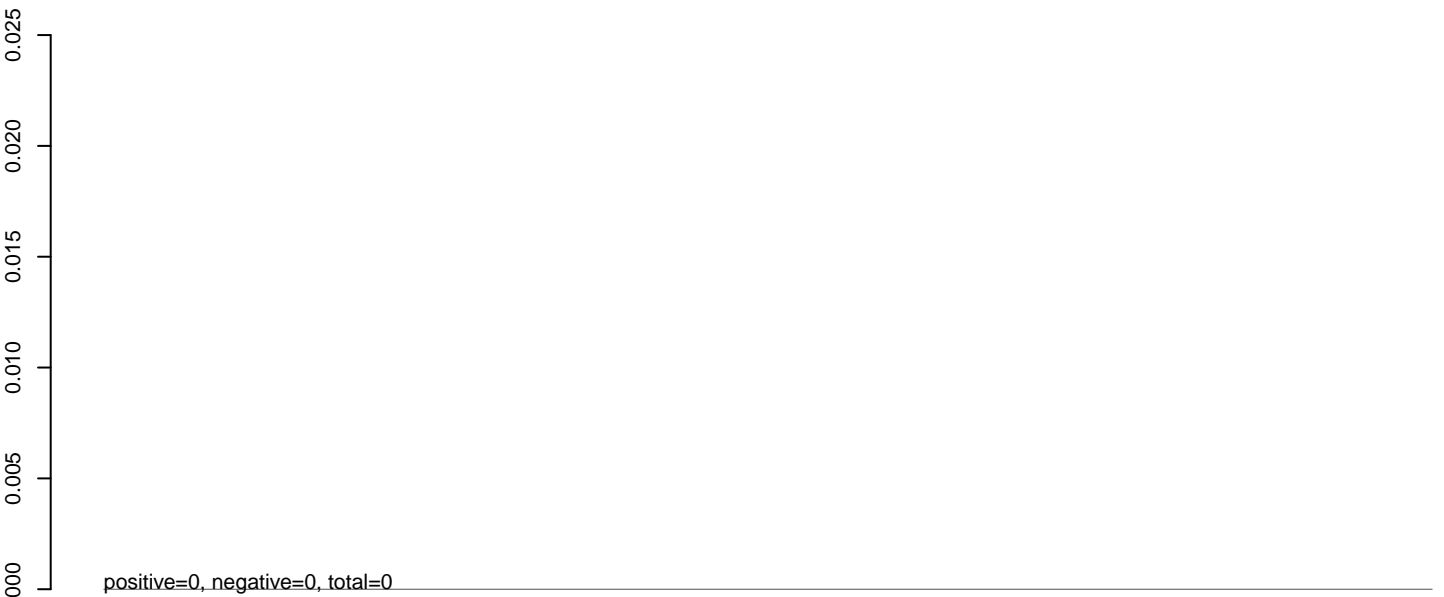
AeAeg_Aag2DENV2NGC.rep



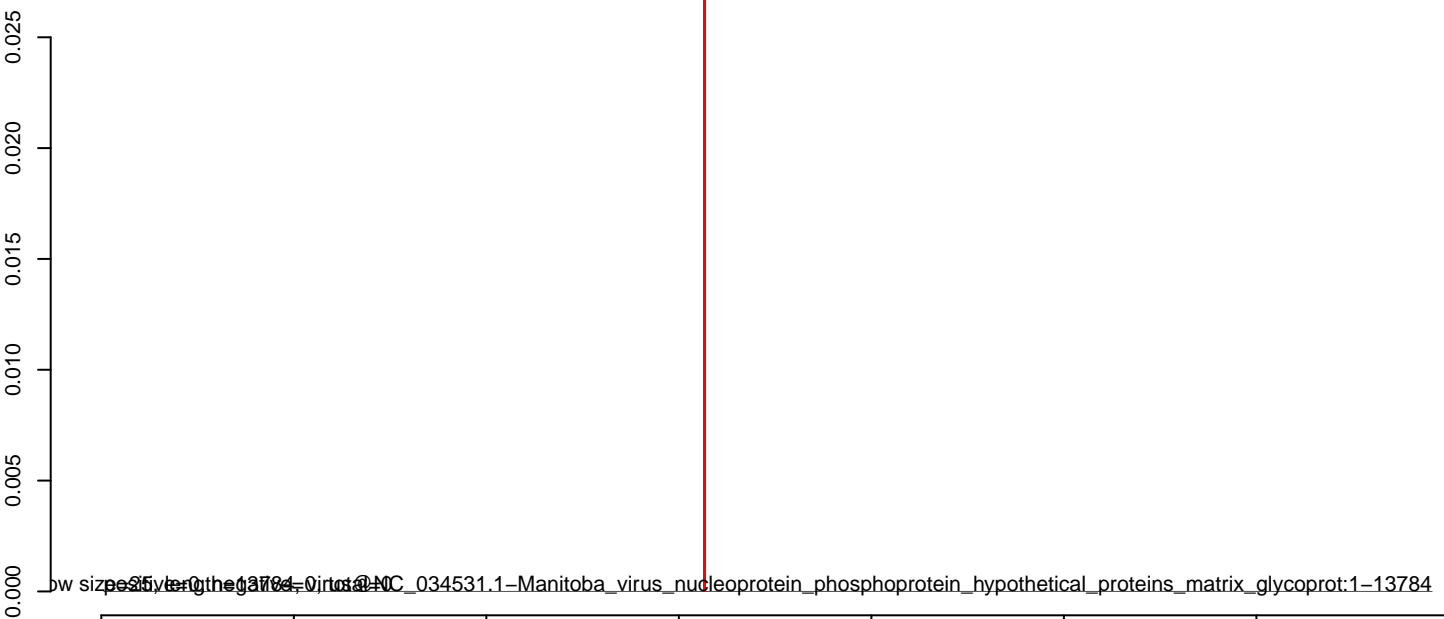
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

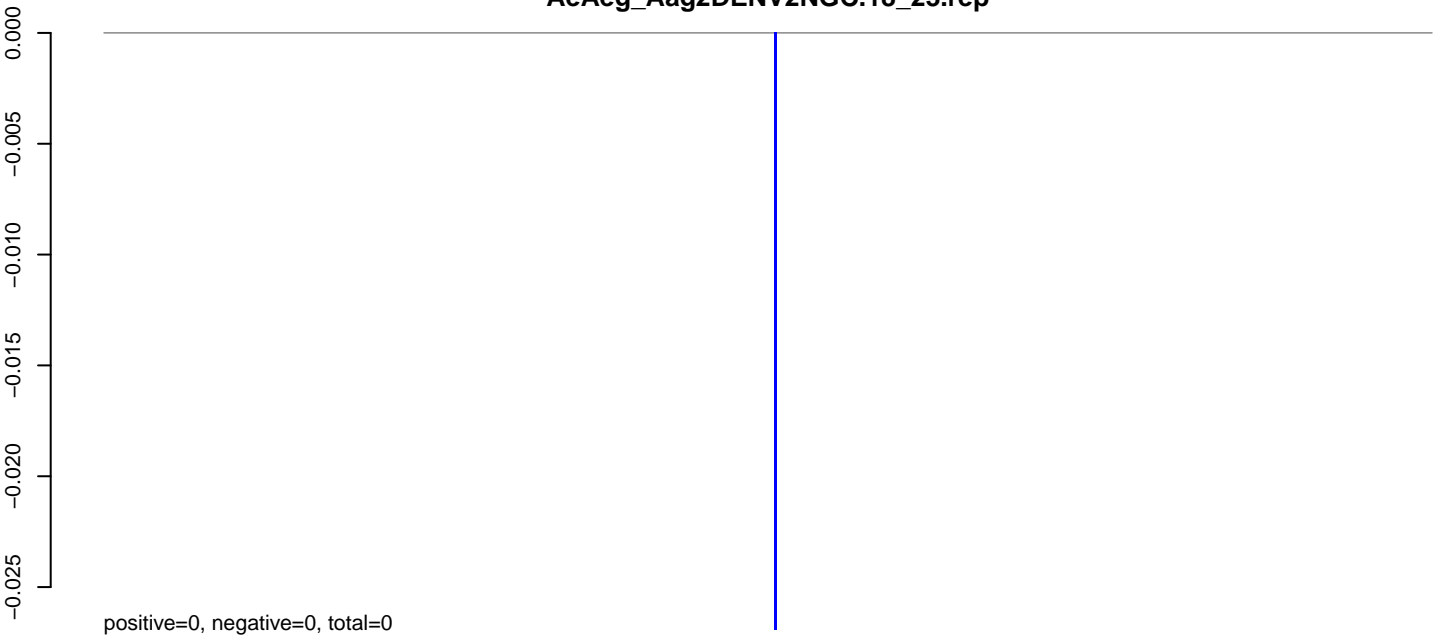


AeAeg_Aag2DENV2NGC.rep



bow size 25, length 13784, int 0, NC_034531.1-Manitoba_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycoprot:1-13784

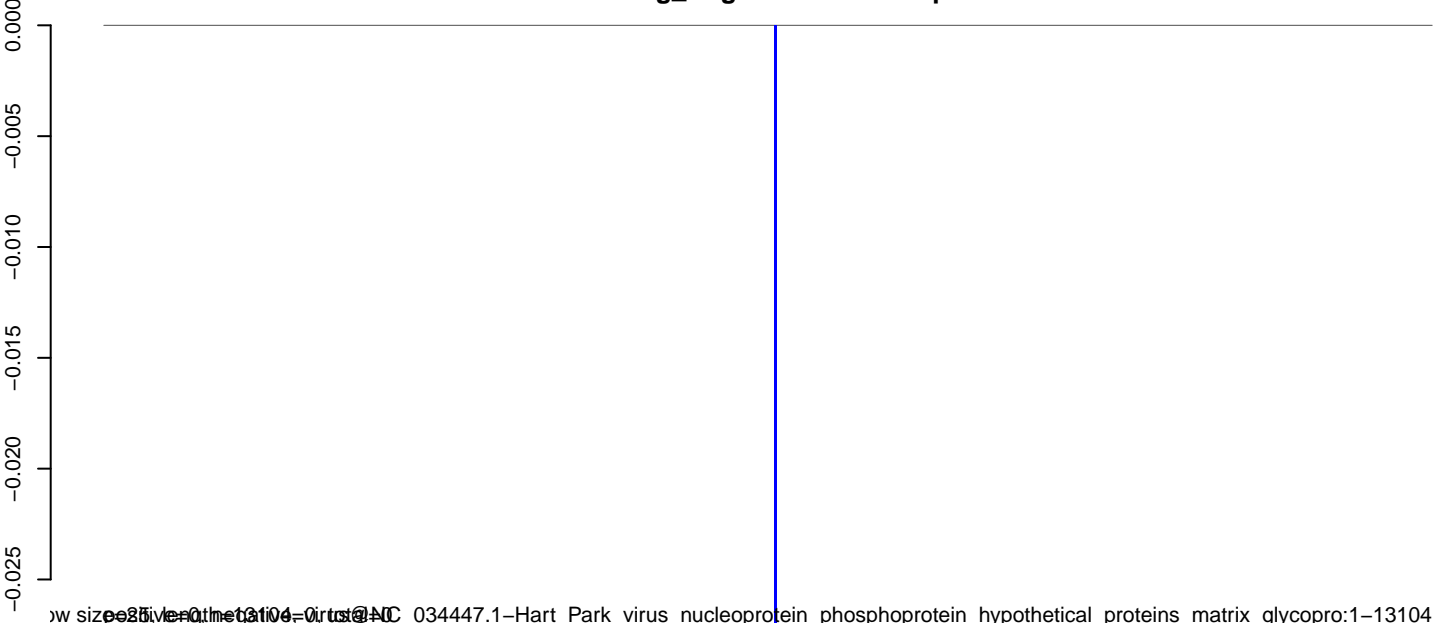
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



ww size 13104, length 13104, virus NC_034447.1-Hart_Park_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopro:1-13104

0 2000 4000 6000 8000 10000 12000

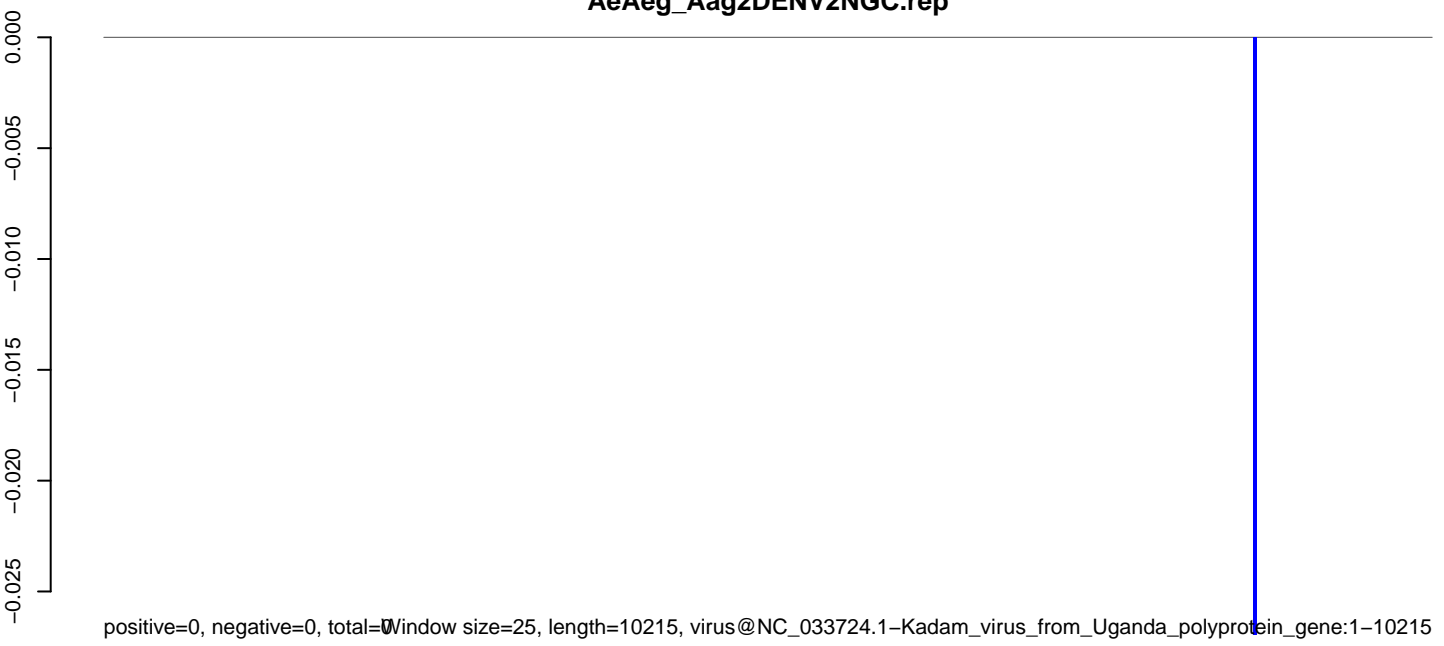
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



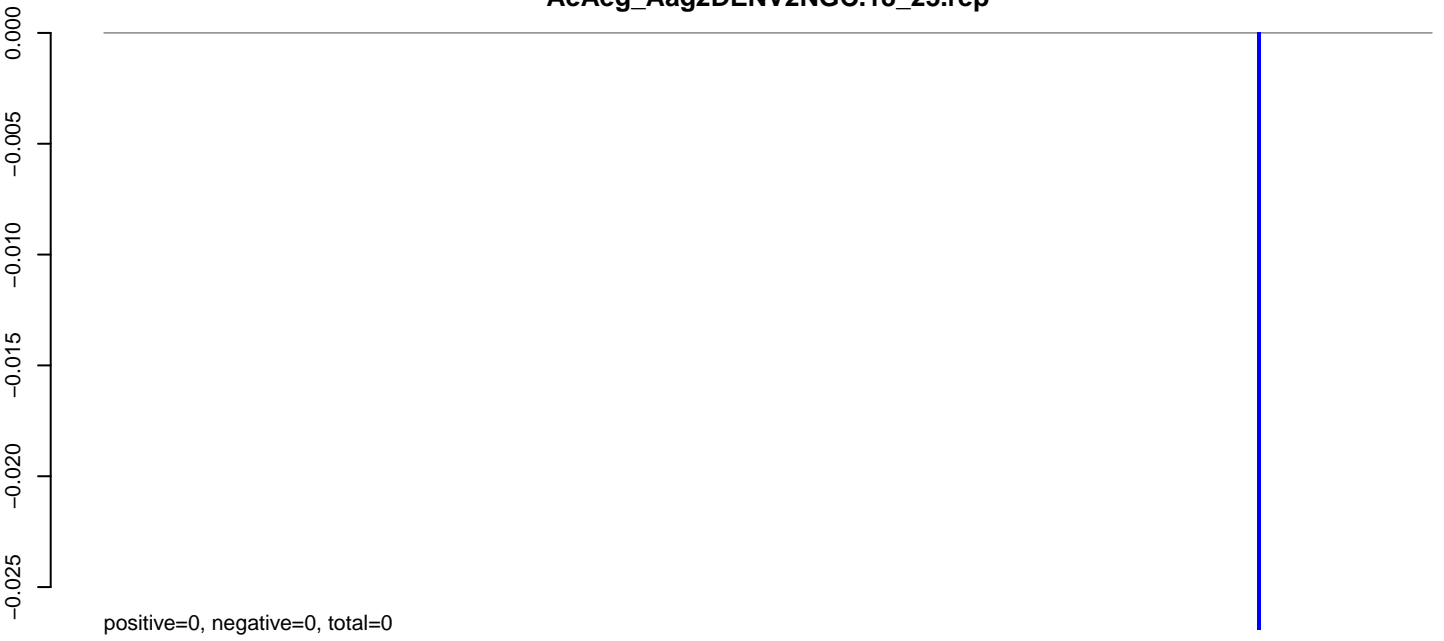
AeAeg_Aag2DENV2NGC.rep



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

0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



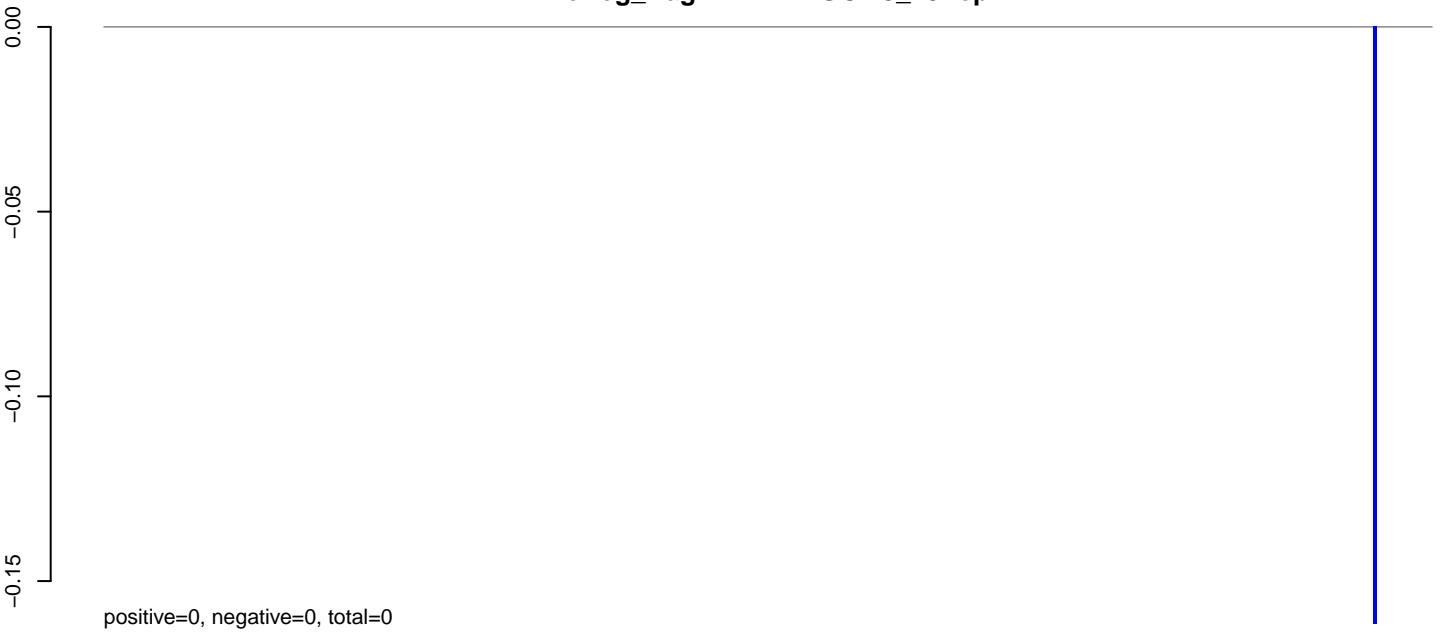
AeAeg_Aag2DENV2NGC.rep



Window size=25, length=10266, virus@NC_033721.1-Meaban_virus_from_France_polyprotein_gene:1-10266

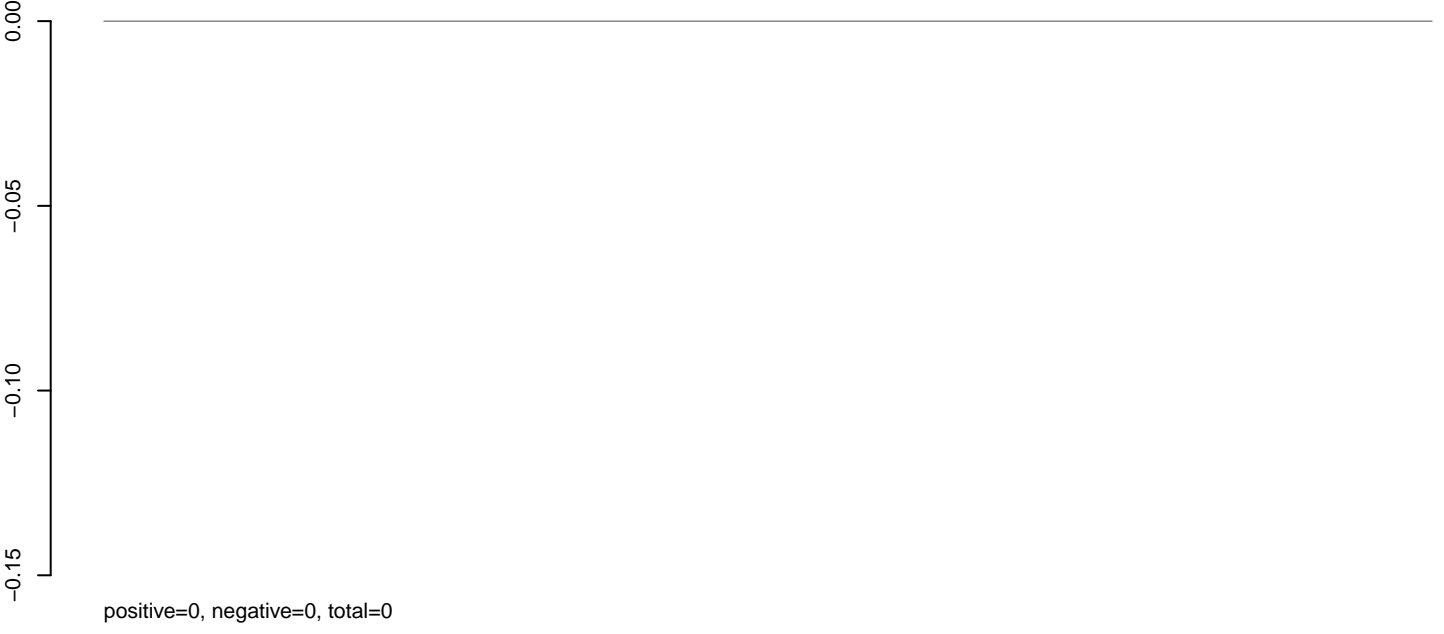
0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



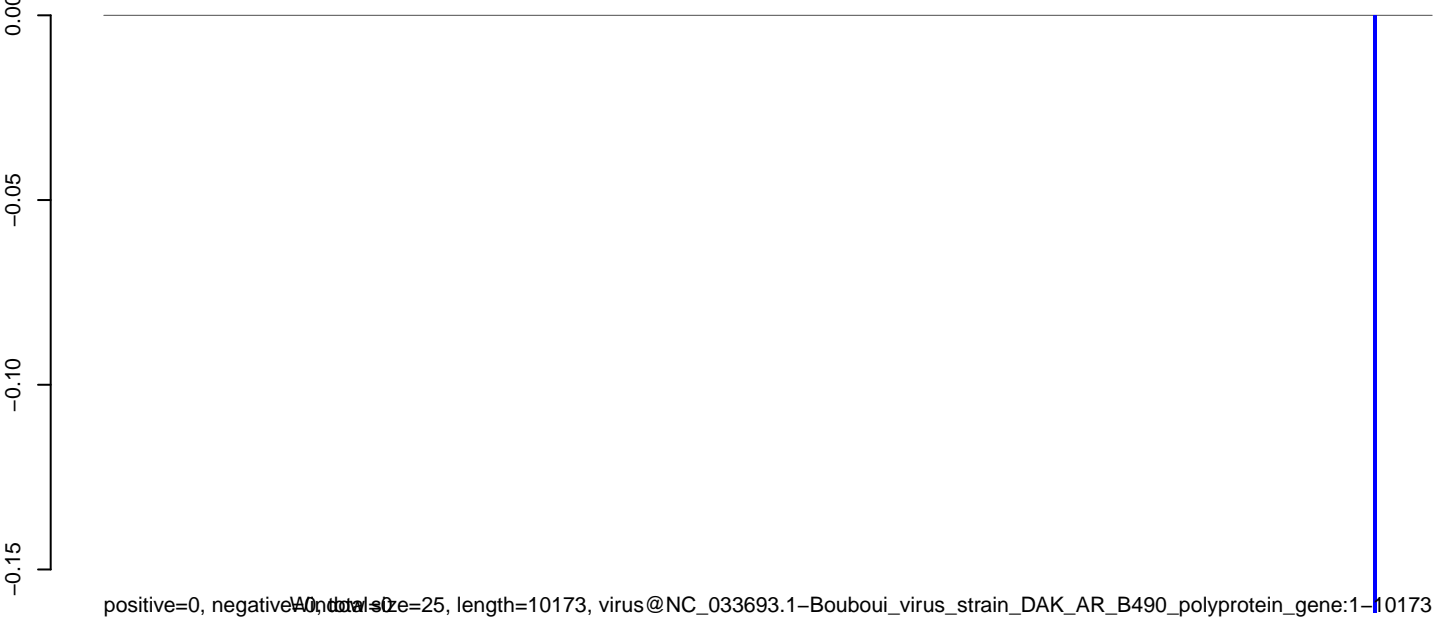
positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.24_35.rep



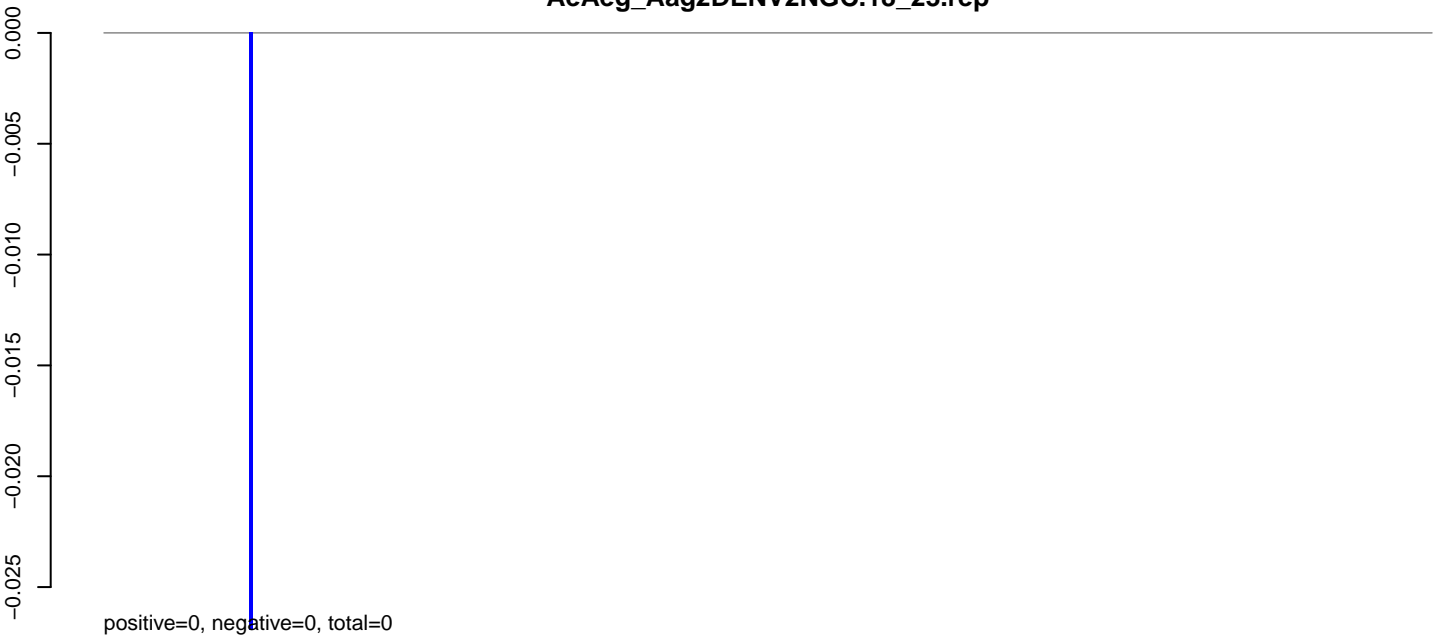
positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.rep



positive=0, negative=0, total=0, n=25, length=10173, virus@NC_033693.1-Bouboui_virus_strain_DAK_AR_B490_polyprotein_gene:1-10173

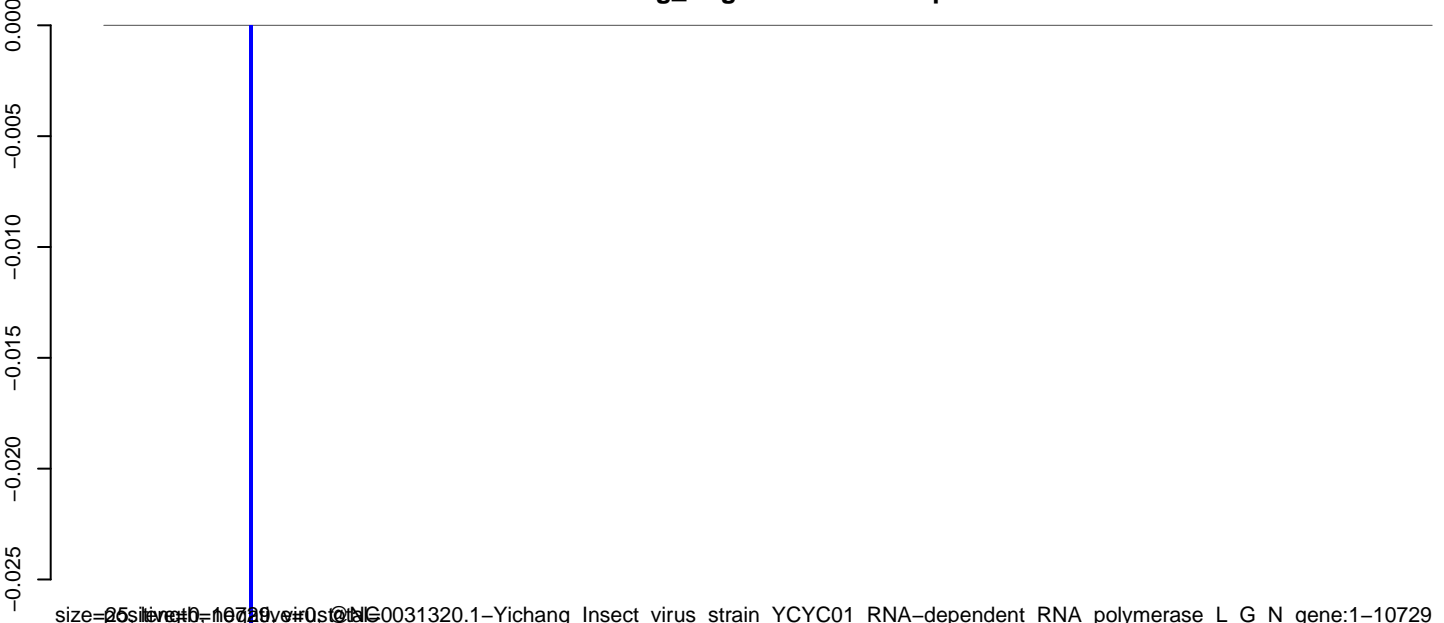
AeAeg_Aag2DENV2NGC.18_23.rep



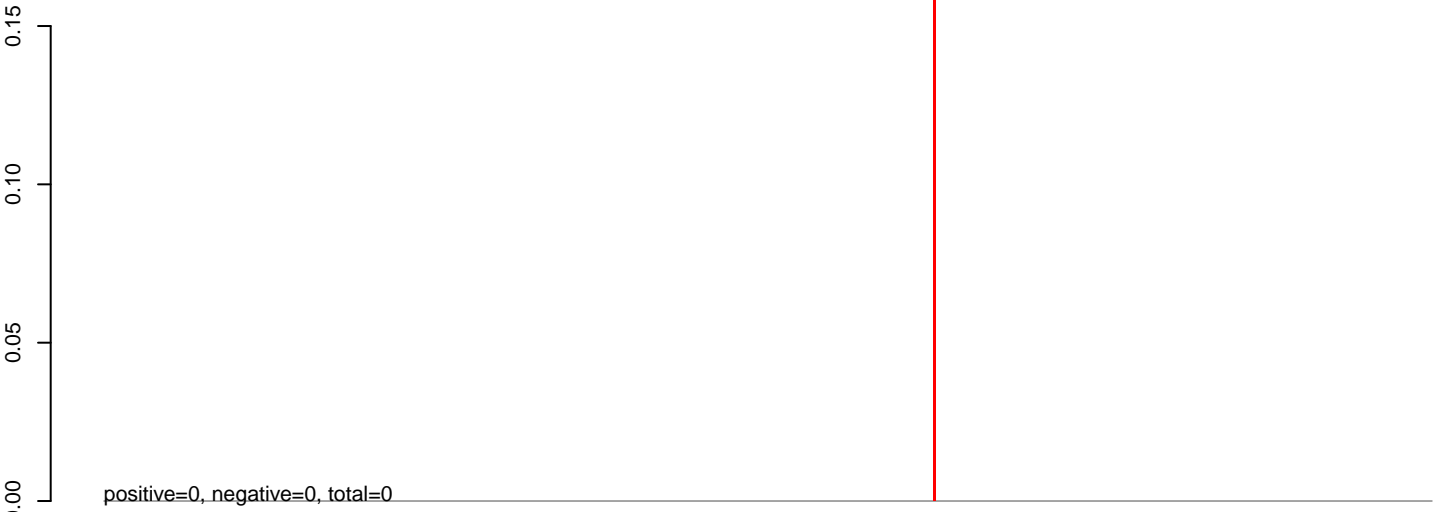
AeAeg_Aag2DENV2NGC.24_35.rep



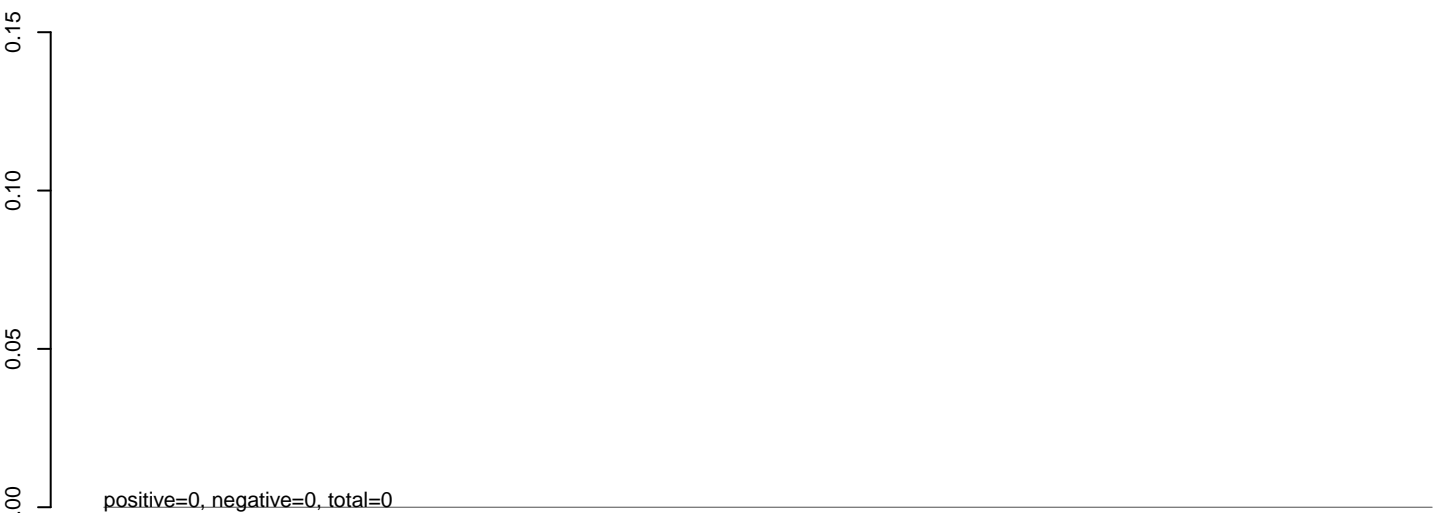
AeAeg_Aag2DENV2NGC.rep



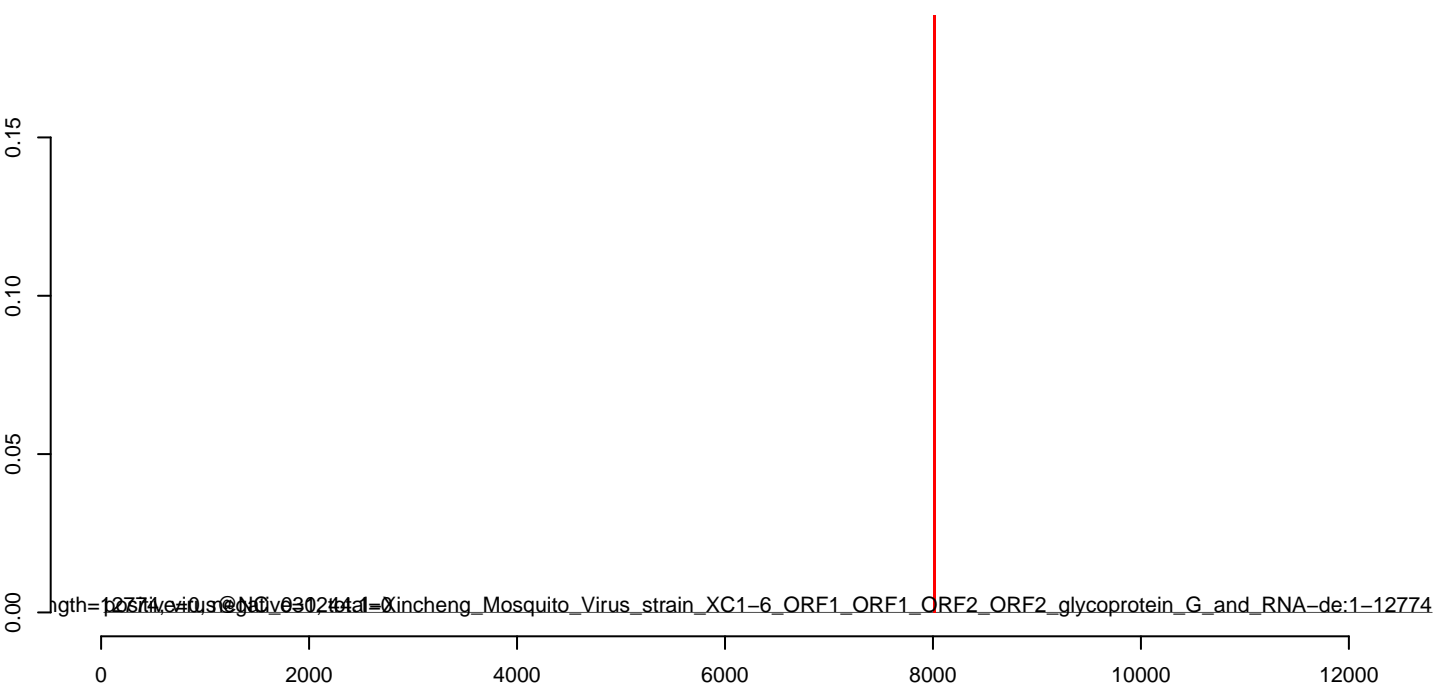
AeAeg_Aag2DENV2NGC.18_23.rep



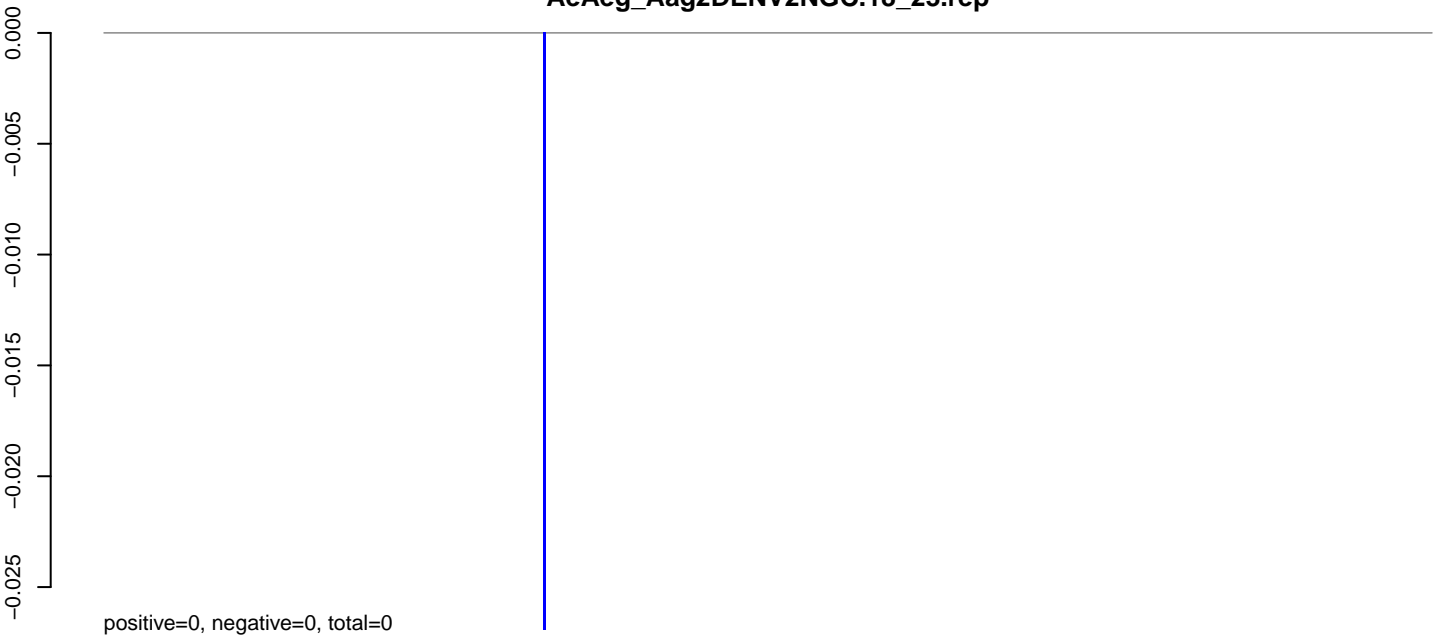
AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



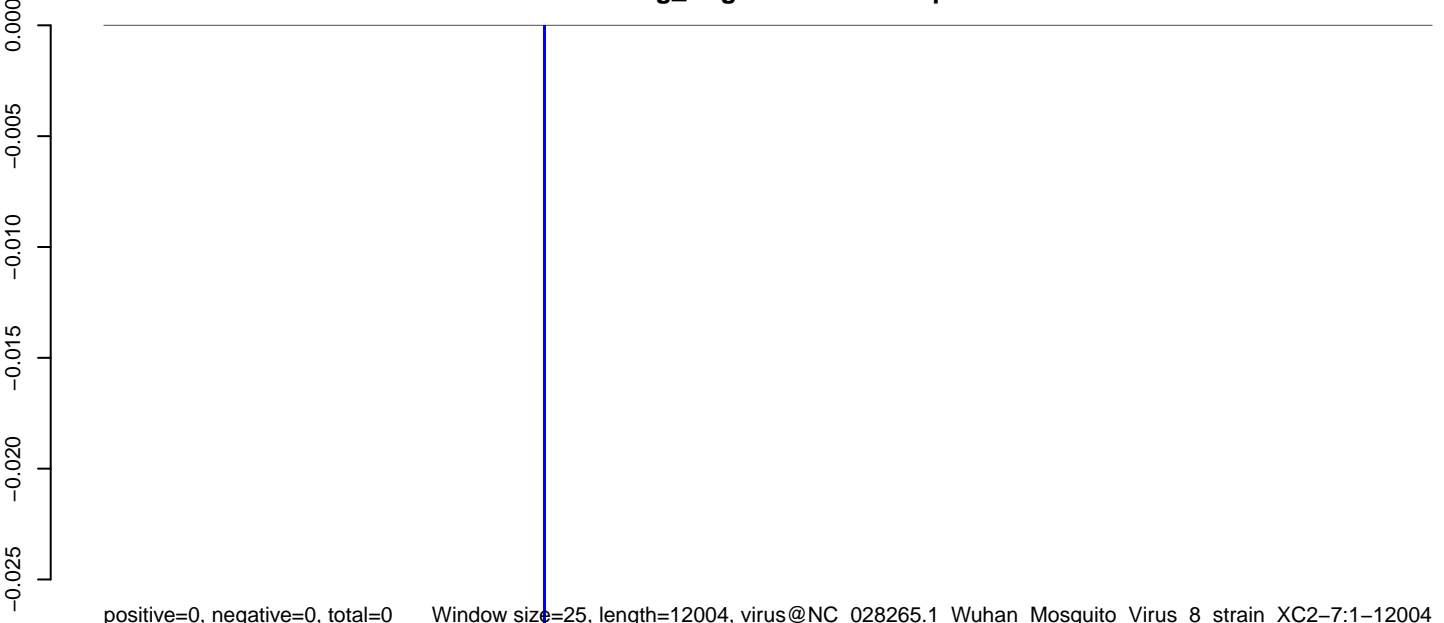
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



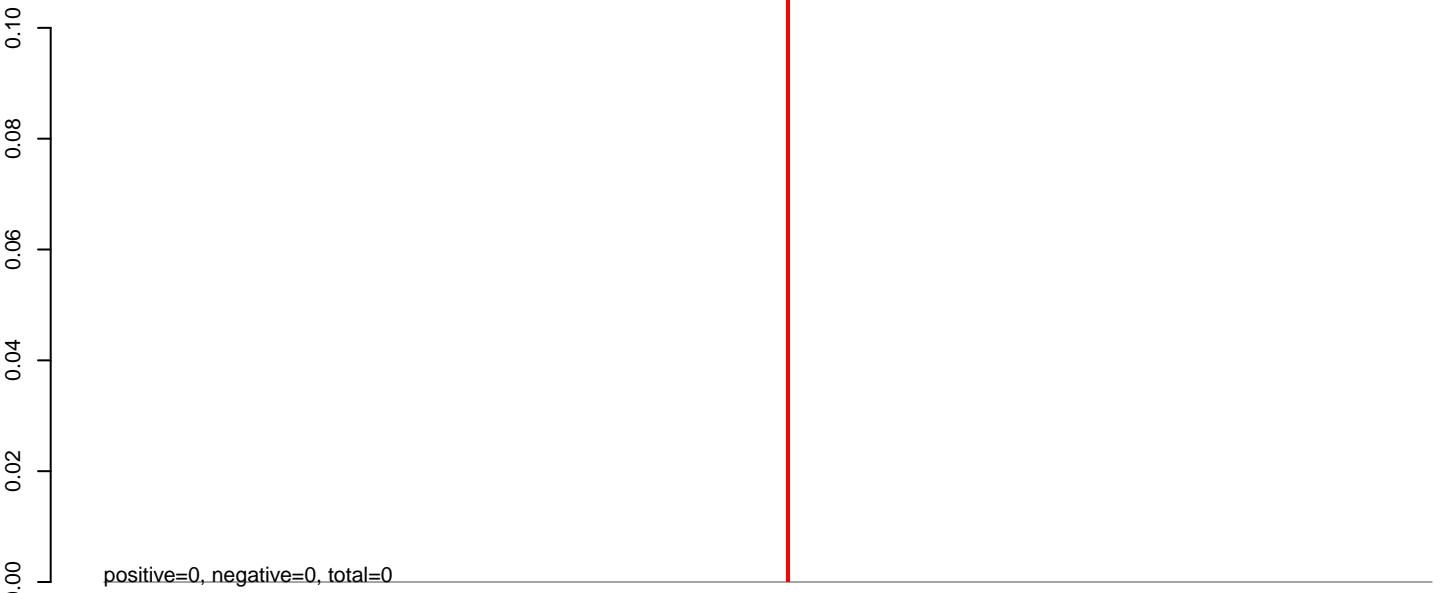
AeAeg_Aag2DENV2NGC.rep



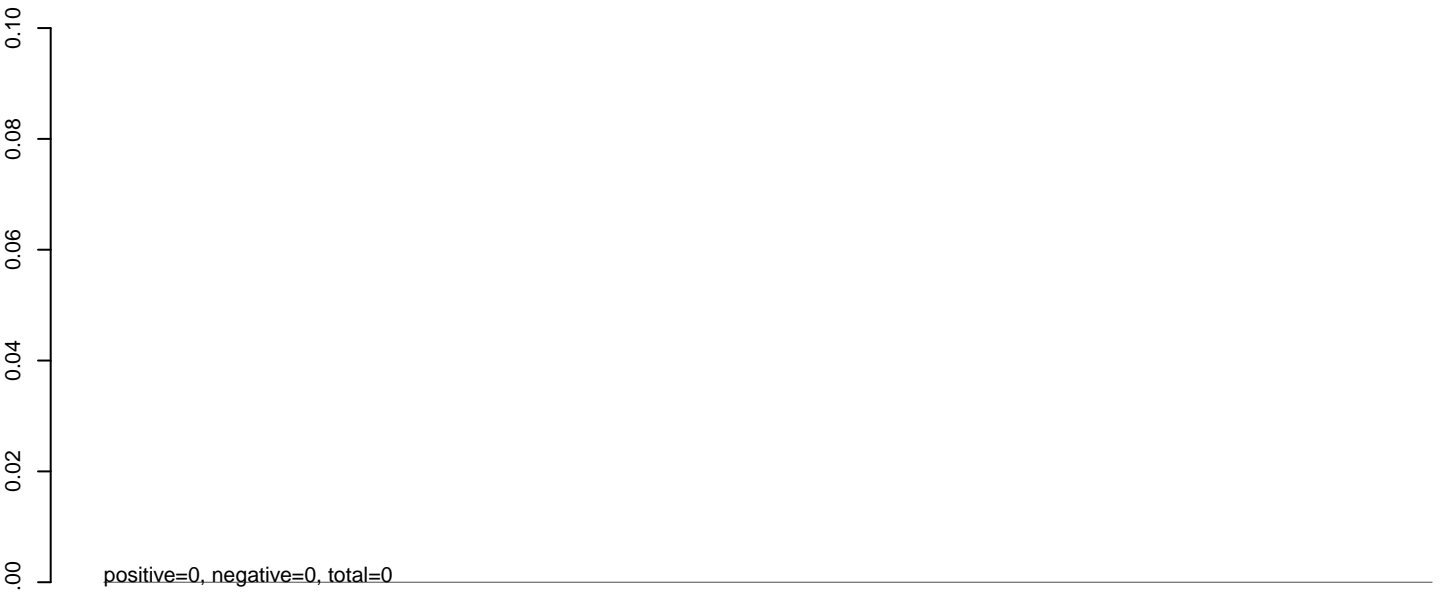
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0 2000 4000 6000 8000 10000 12000

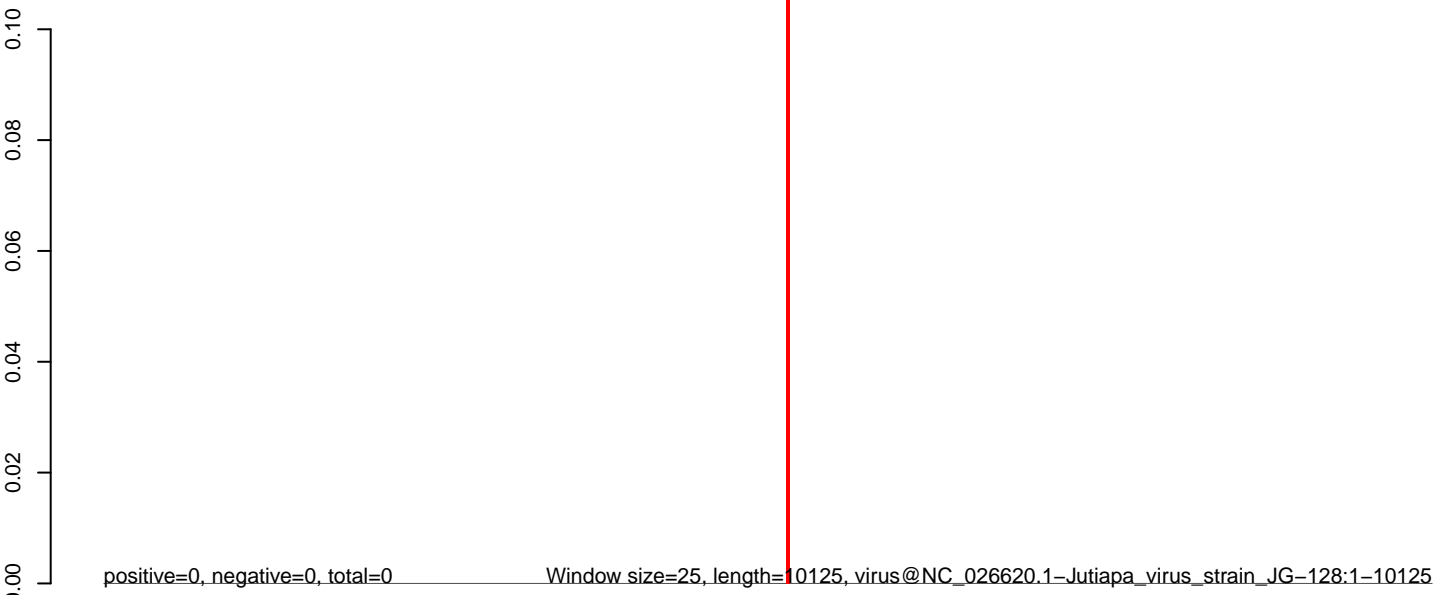
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep

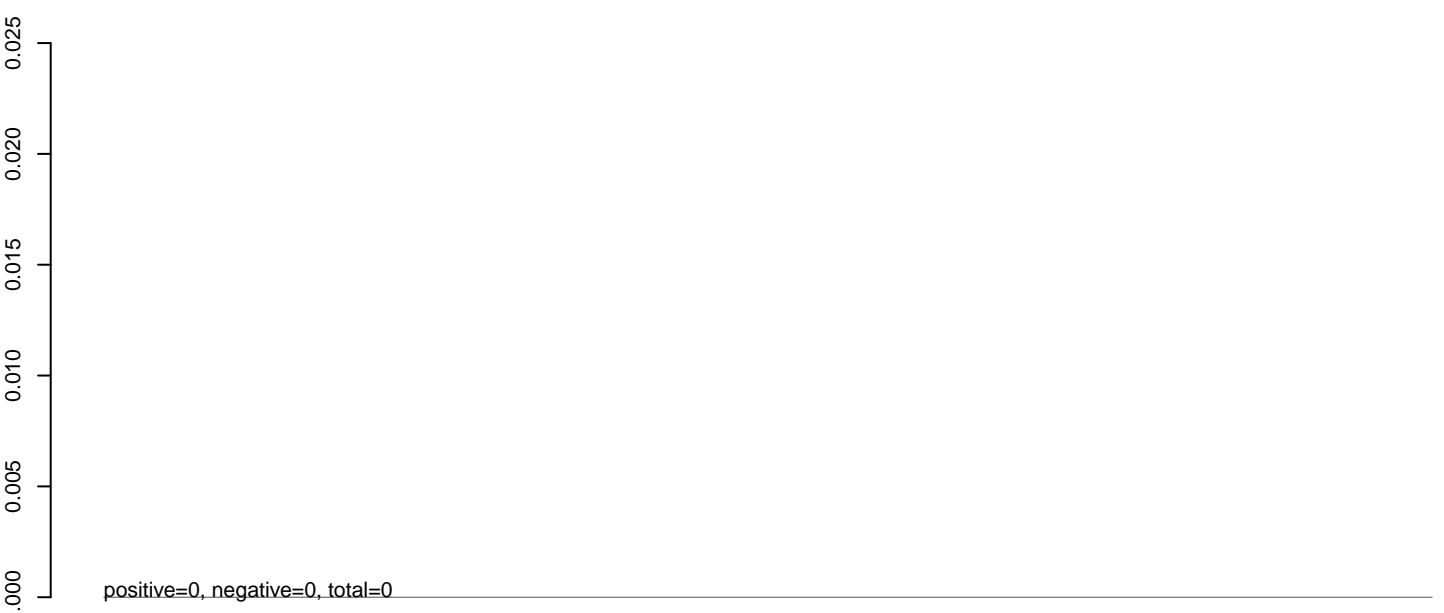


0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



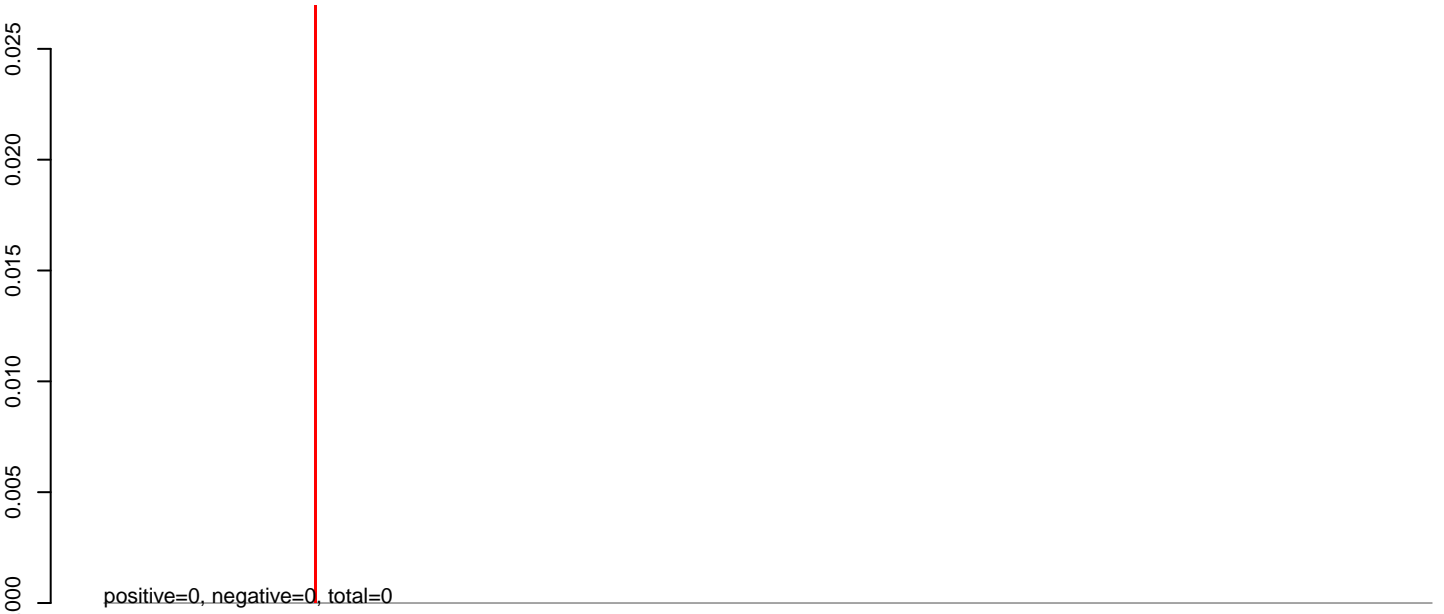
AeAeg_Aag2DENV2NGC.rep



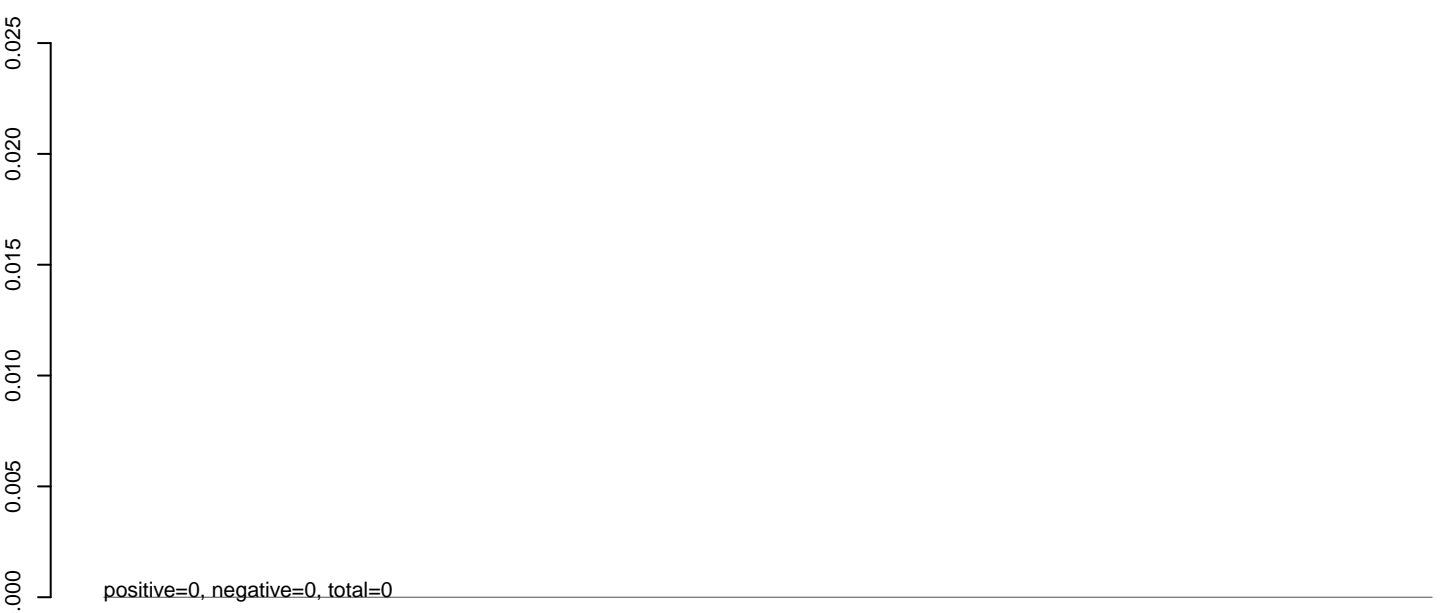
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0 2000 4000 6000 8000 10000 12000

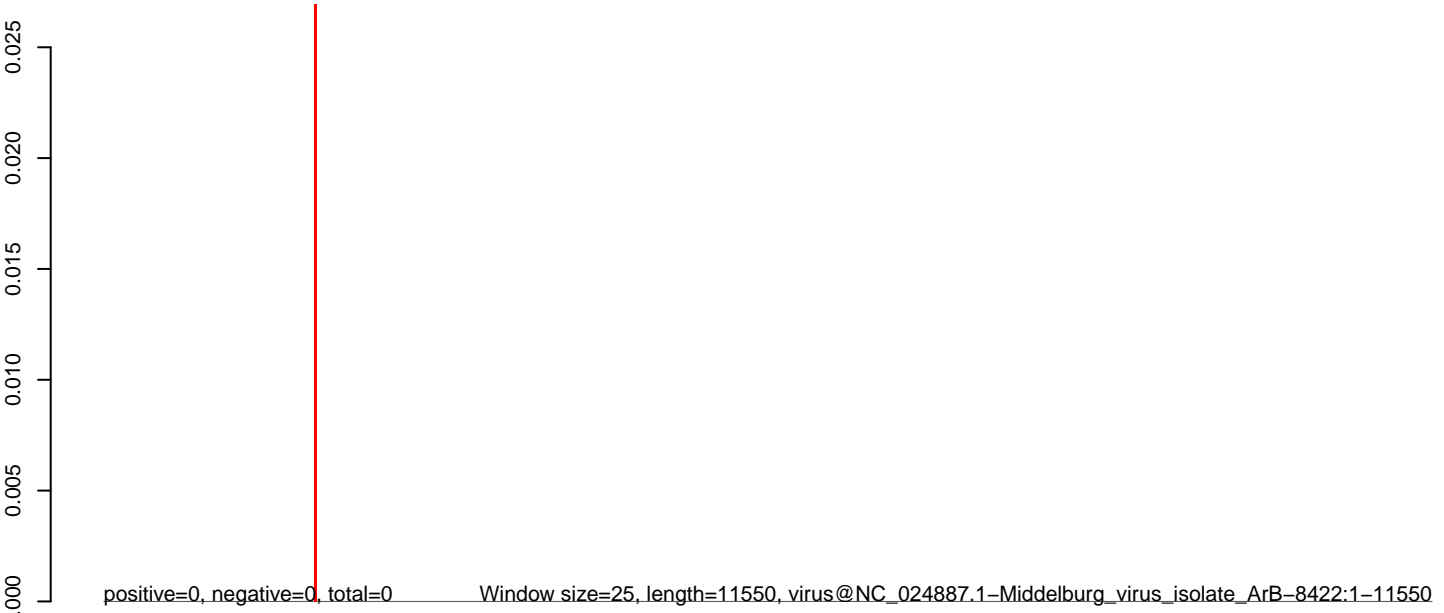
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

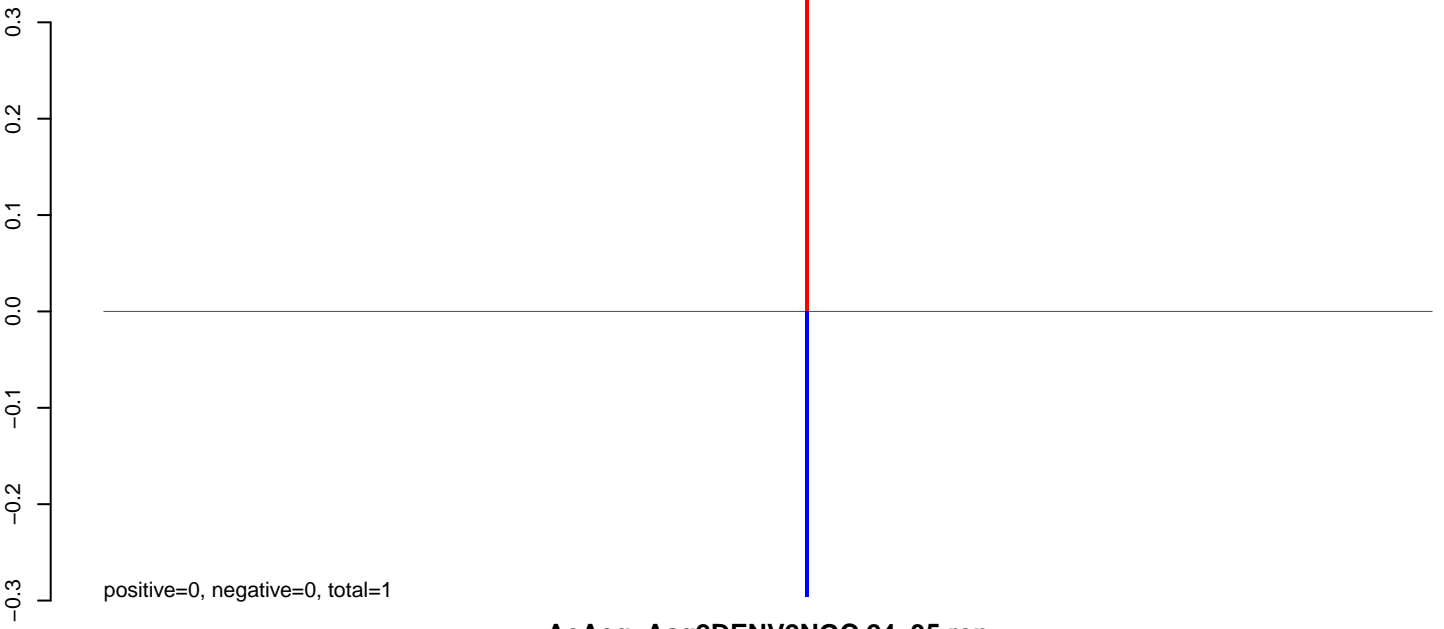


AeAeg_Aag2DENV2NGC.rep

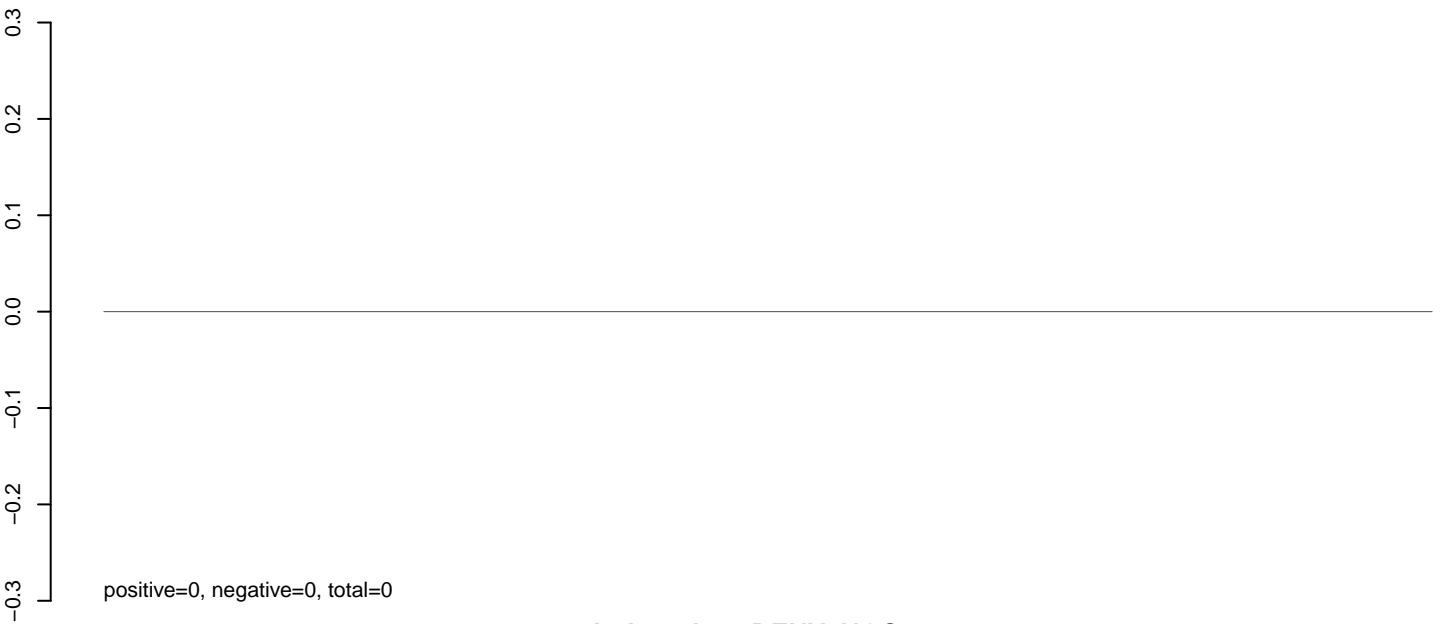


0 2000 4000 6000 8000 10000 12000

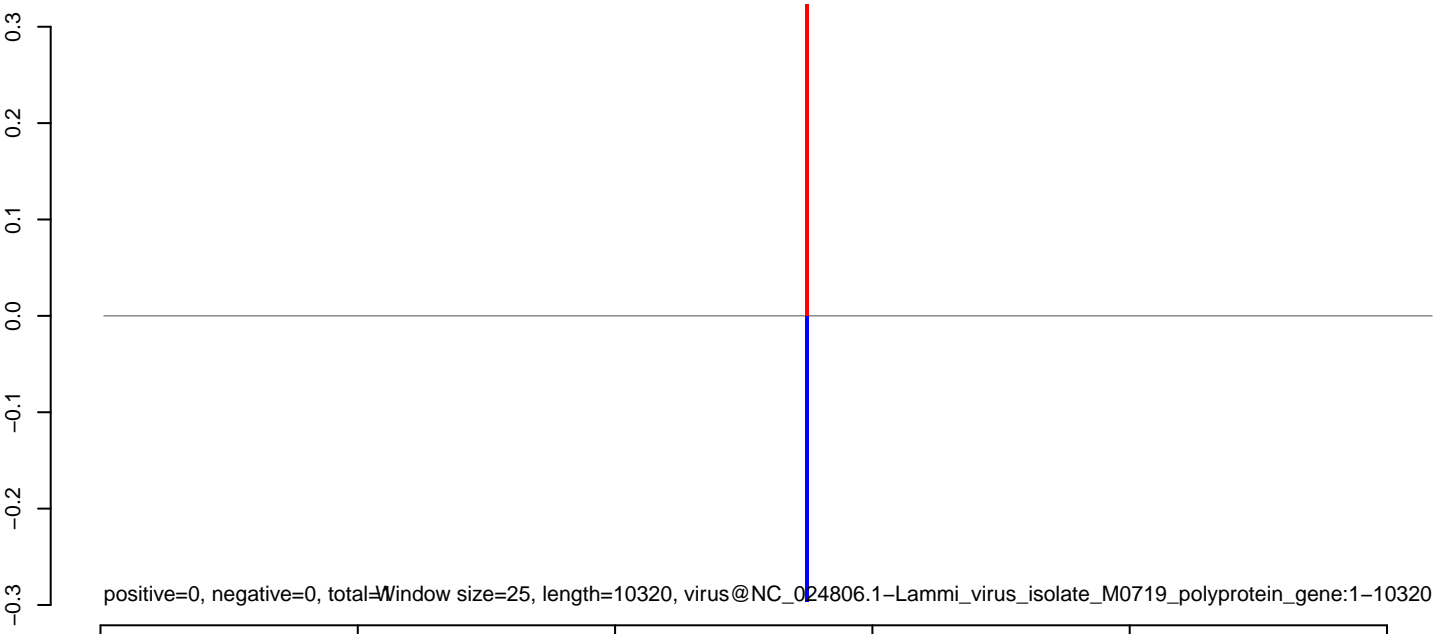
AeAeg_Aag2DENV2NGC.18_23.rep



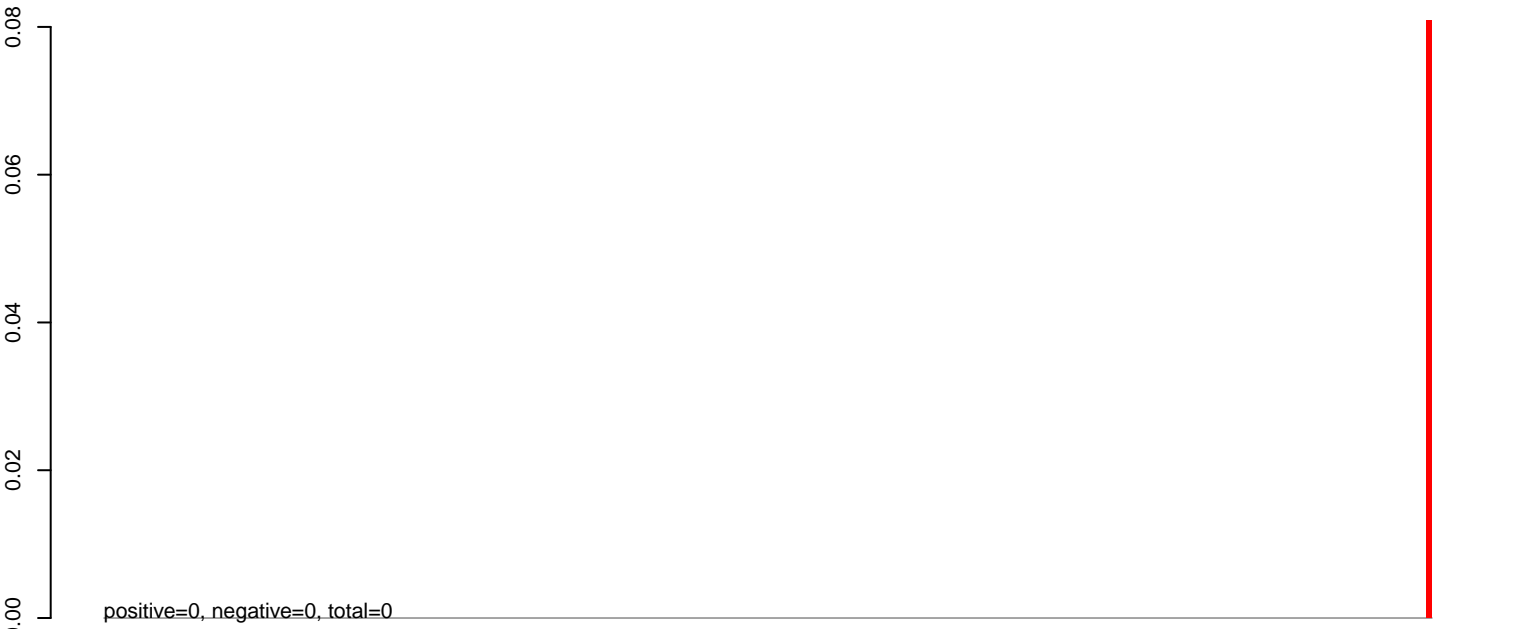
AeAeg_Aag2DENV2NGC.24_35.rep



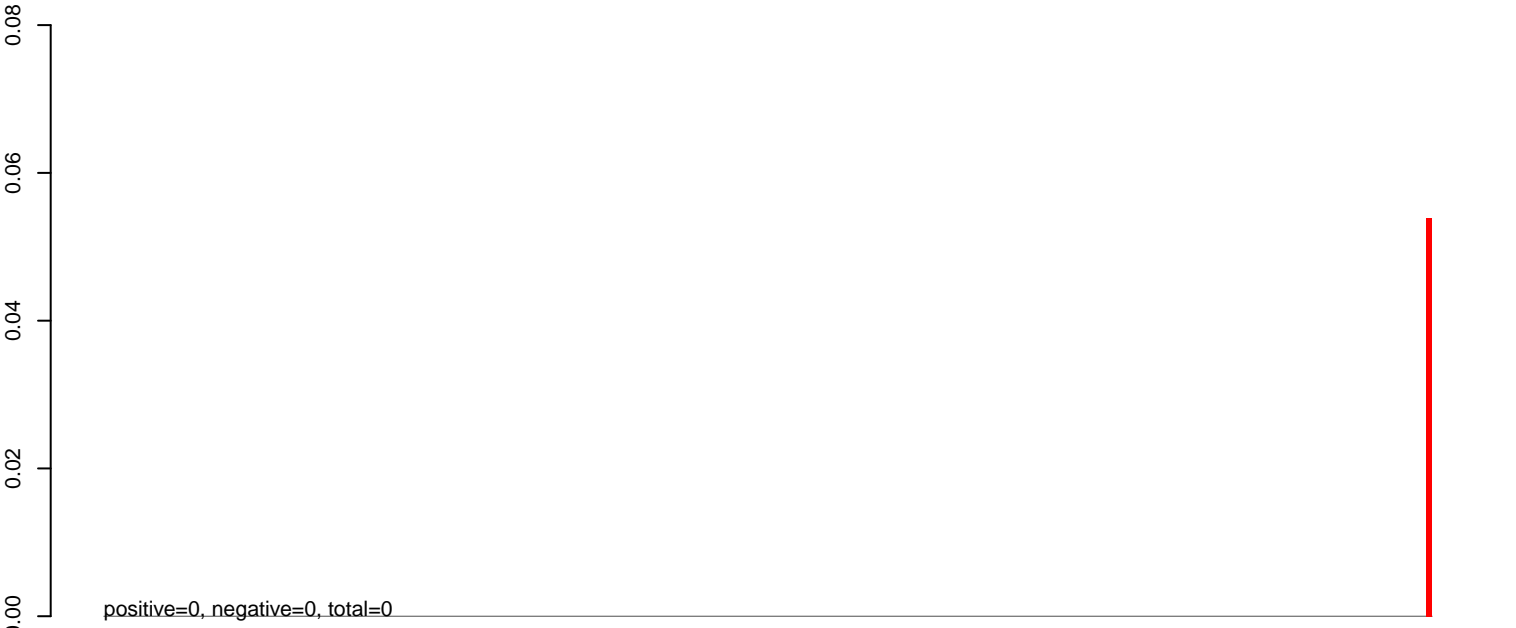
AeAeg_Aag2DENV2NGC.rep



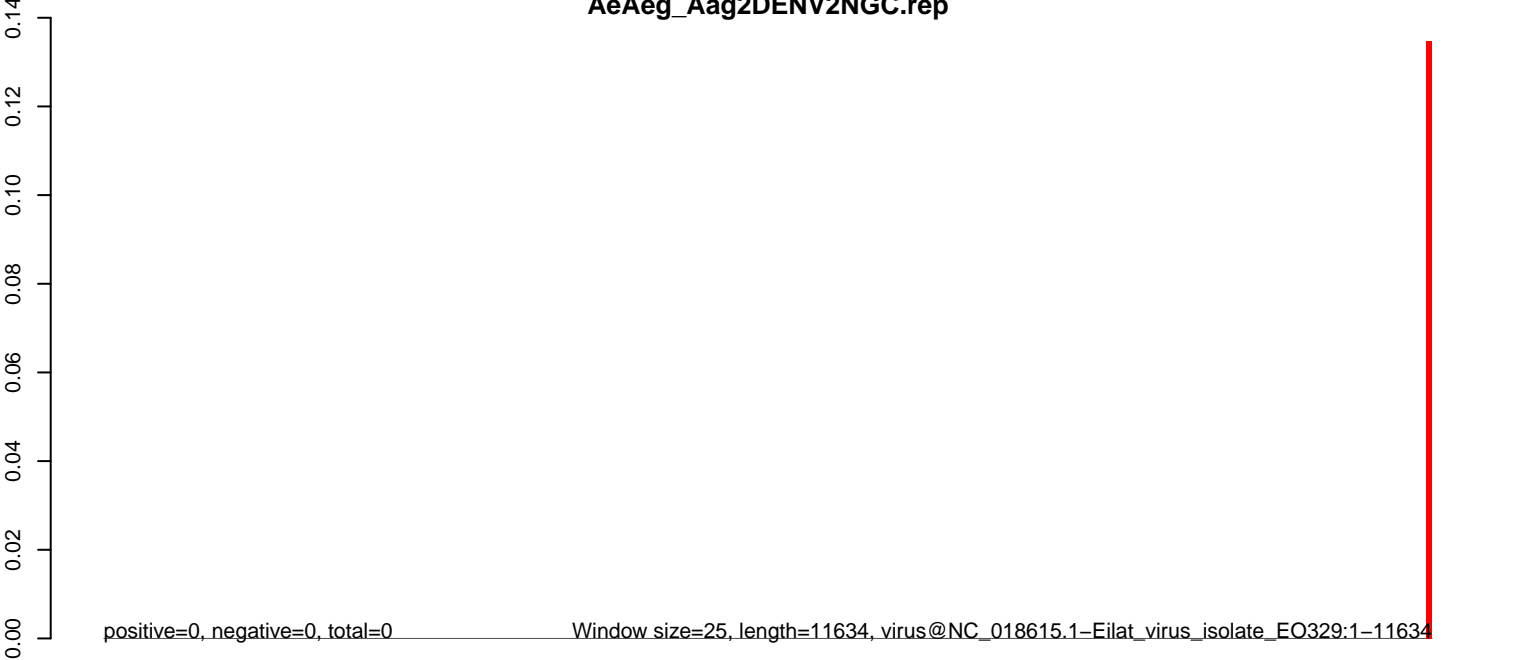
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2DENV2NGC.18_23.rep

0.15
0.10
0.05
0.00

positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.24_35.rep

0.15
0.10
0.05
0.00

positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.rep

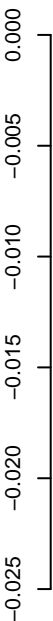
0.15
0.10
0.05
0.00

positive=0, negative=0, total=0

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

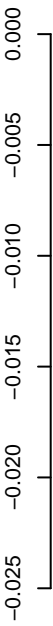
0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2DENV2NGC.18_23.rep



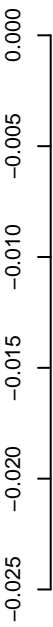
positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.24_35.rep



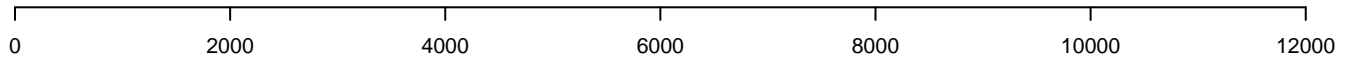
positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.rep

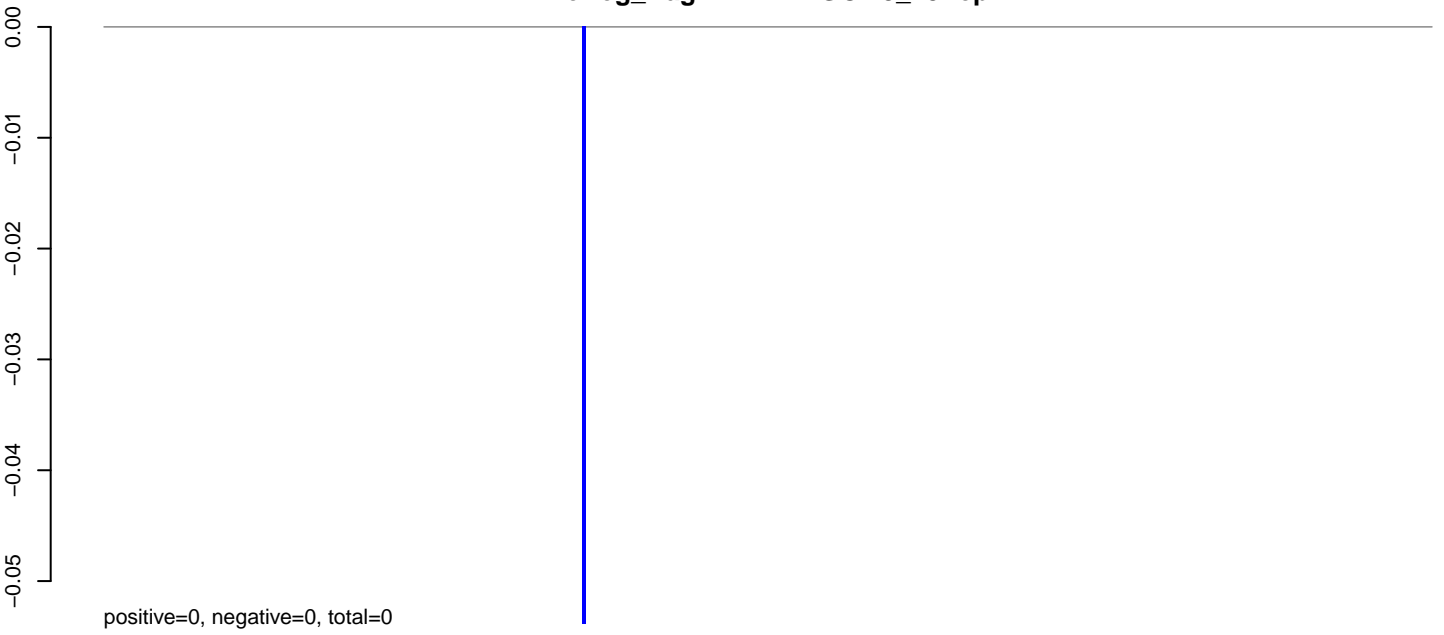


positive=0, negative=0, total=0

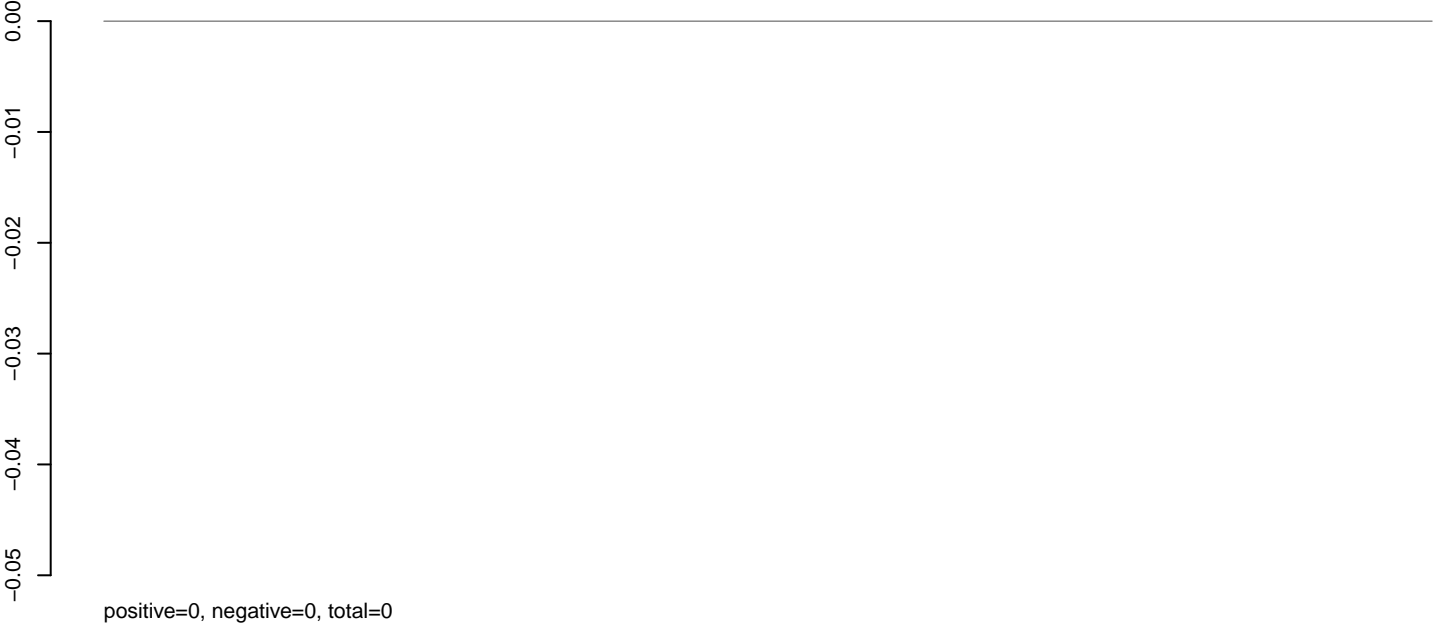
Window size=25, length=12333, virus@NC_007919.3-Nora_virus:1-12333



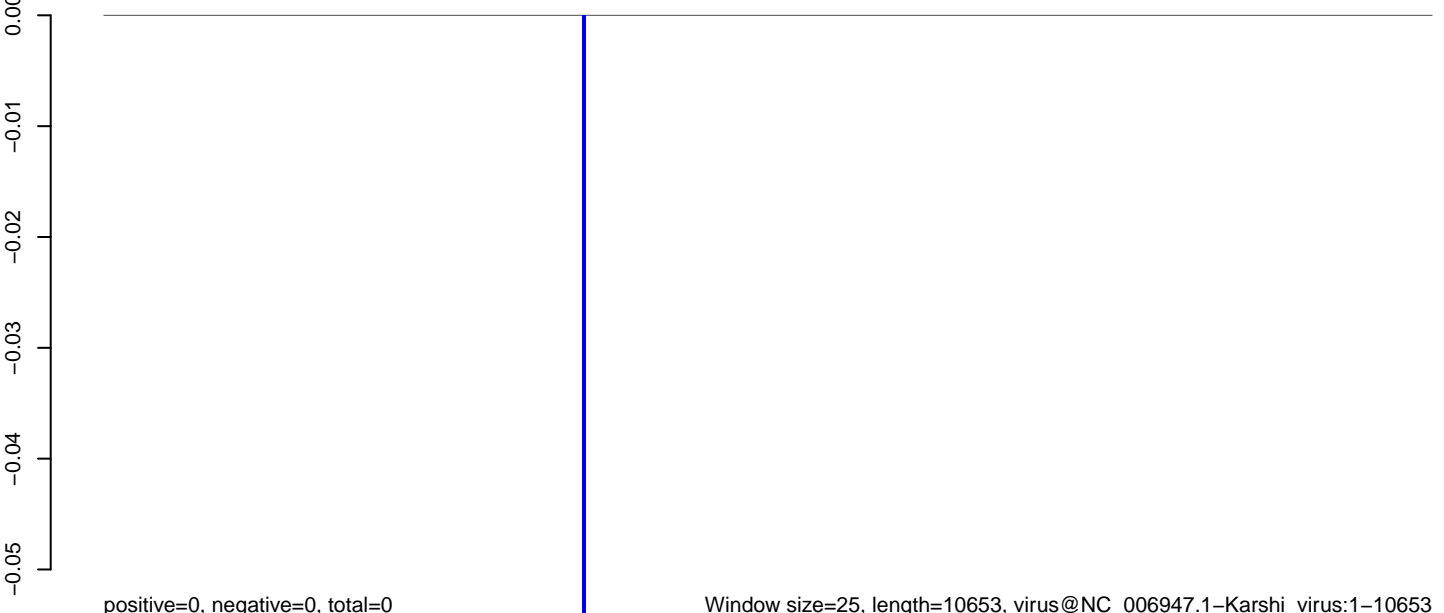
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



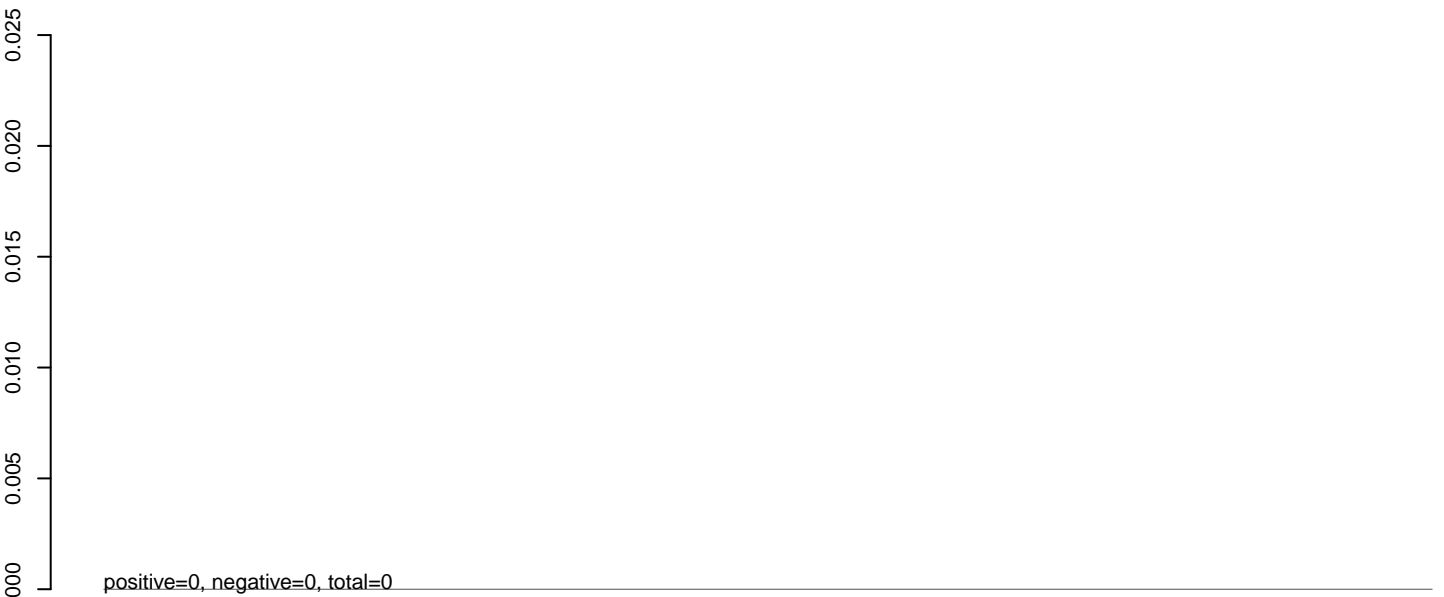
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

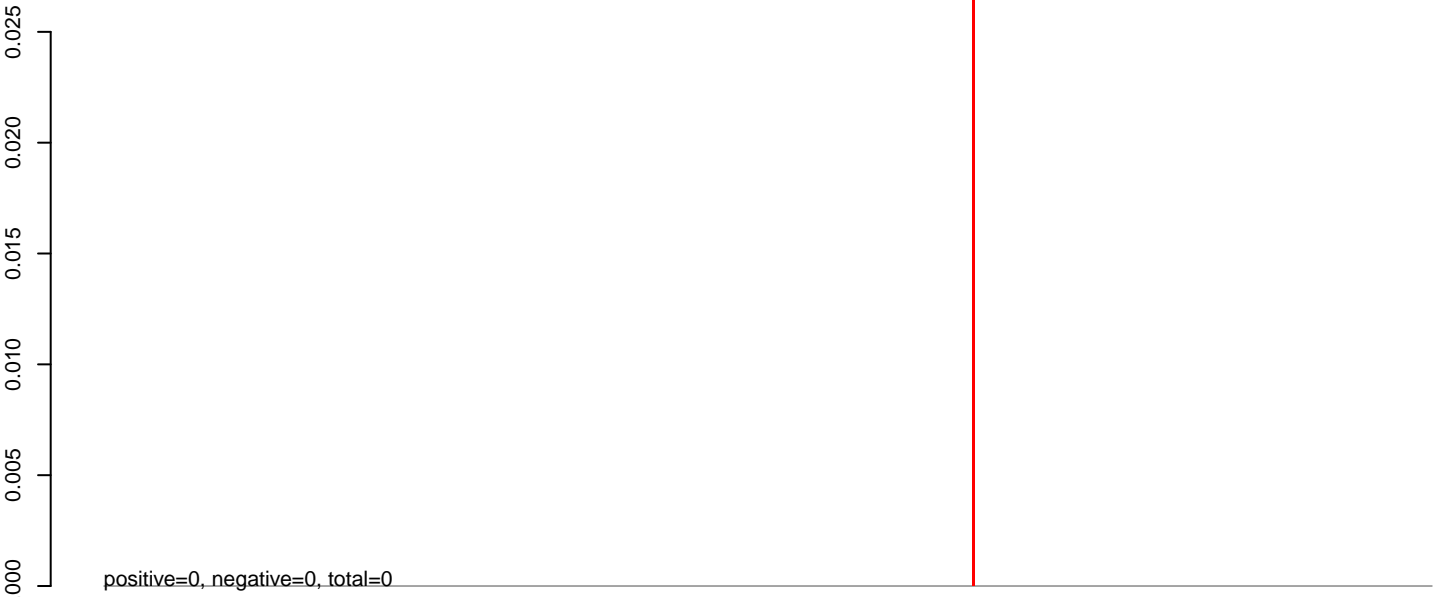


AeAeg_Aag2DENV2NGC.rep

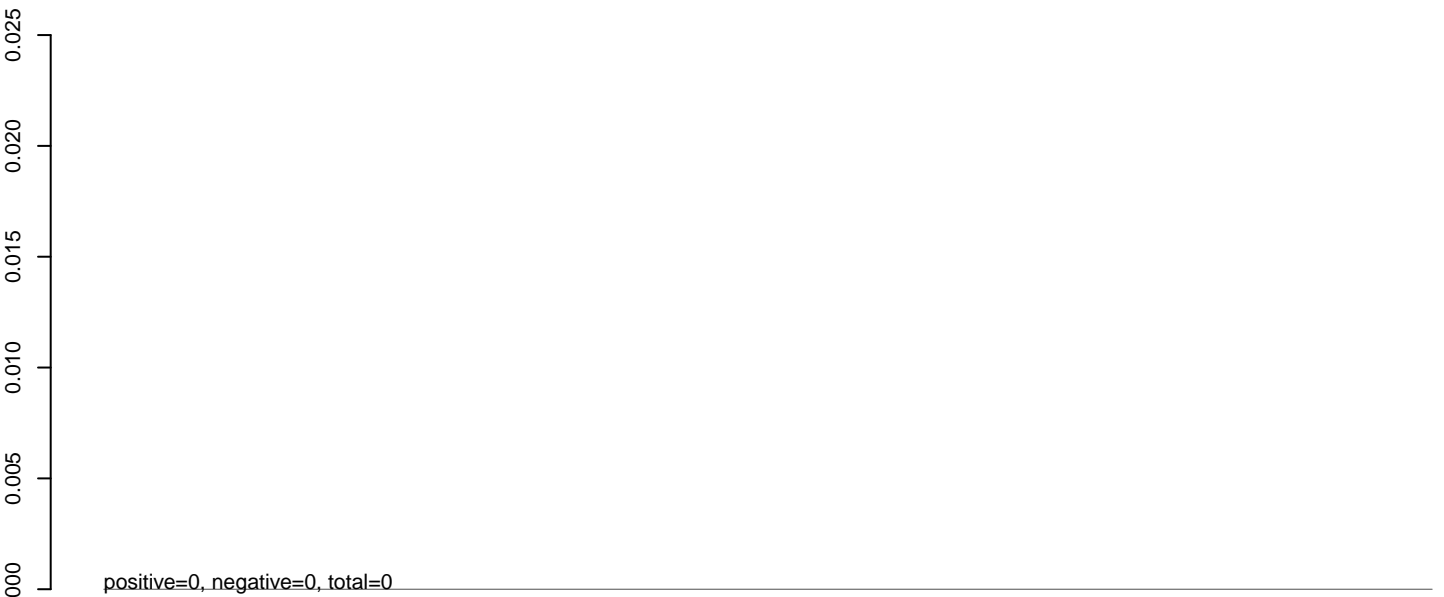


0 2000 4000 6000 8000 10000 12000

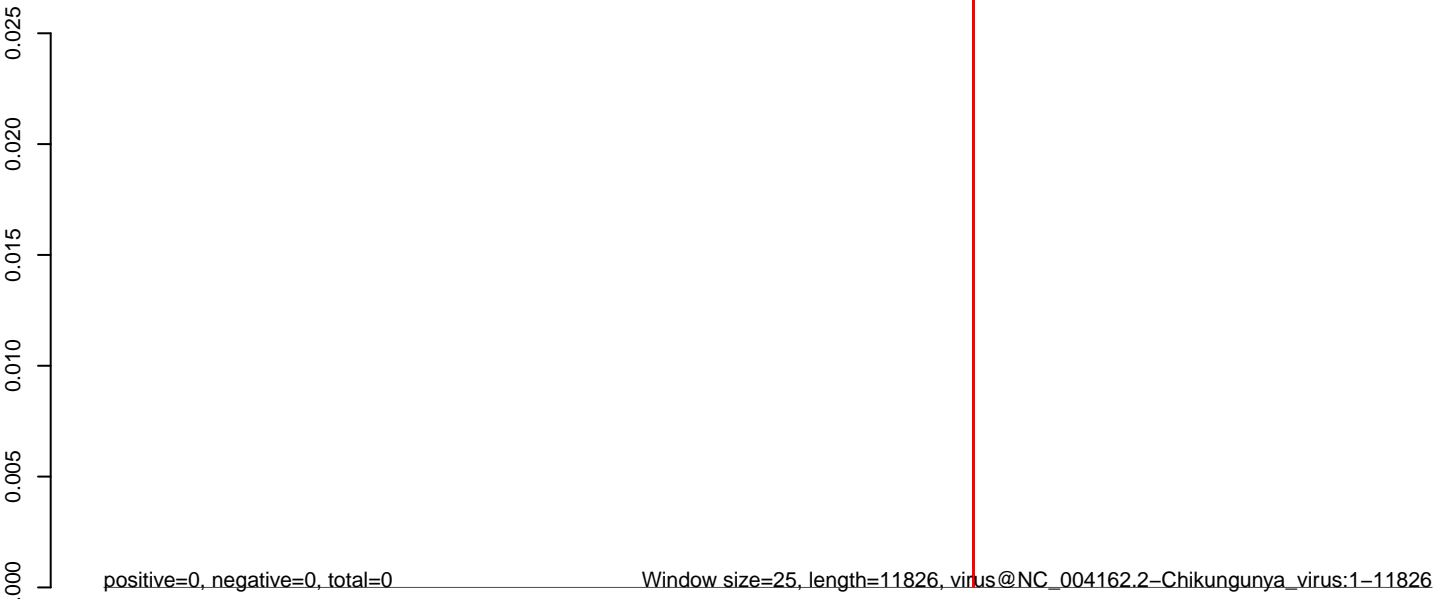
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

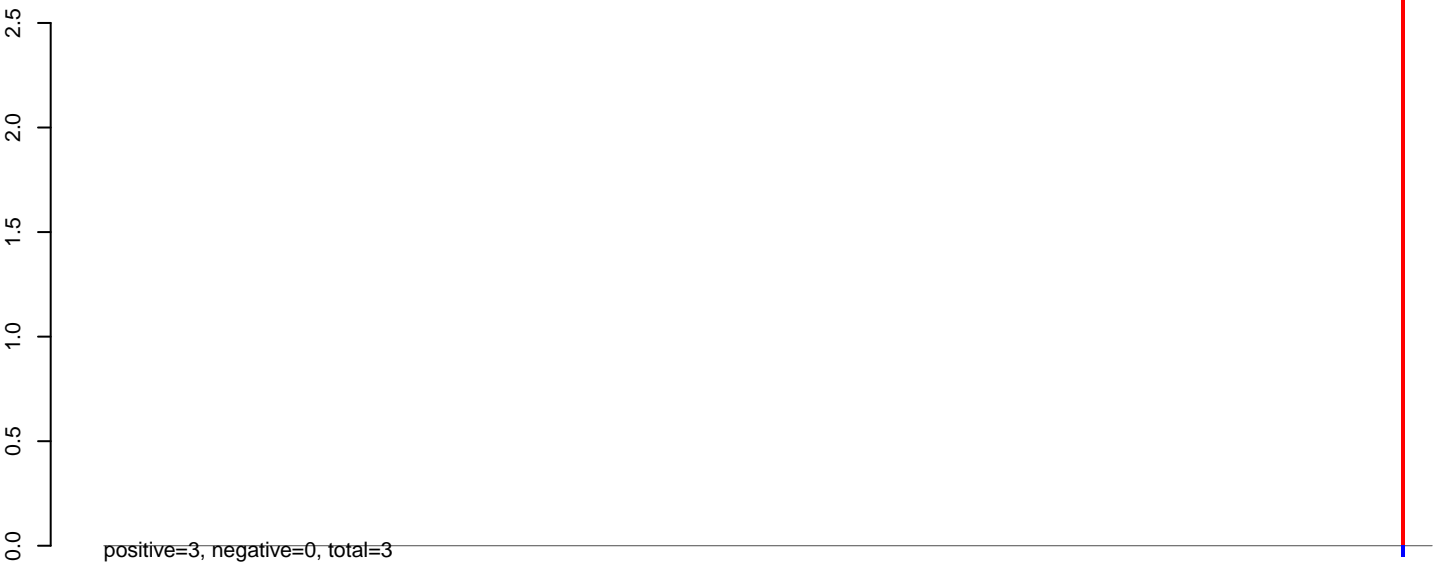


AeAeg_Aag2DENV2NGC.rep

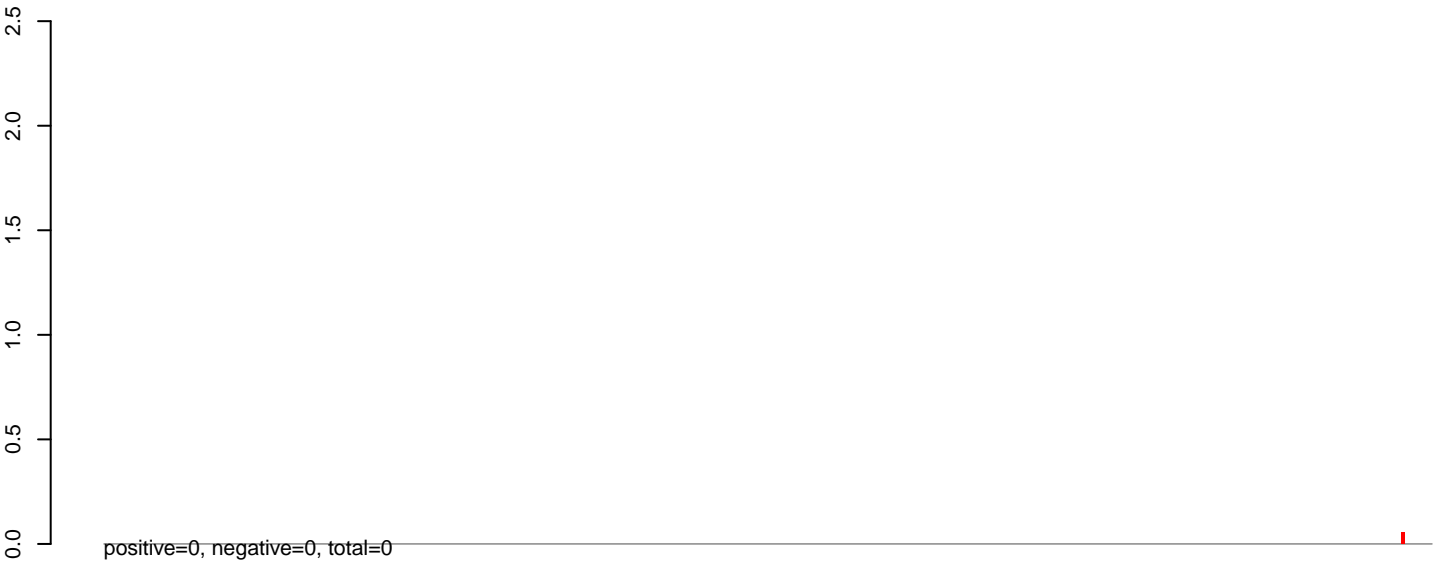


0 2000 4000 6000 8000 10000 12000

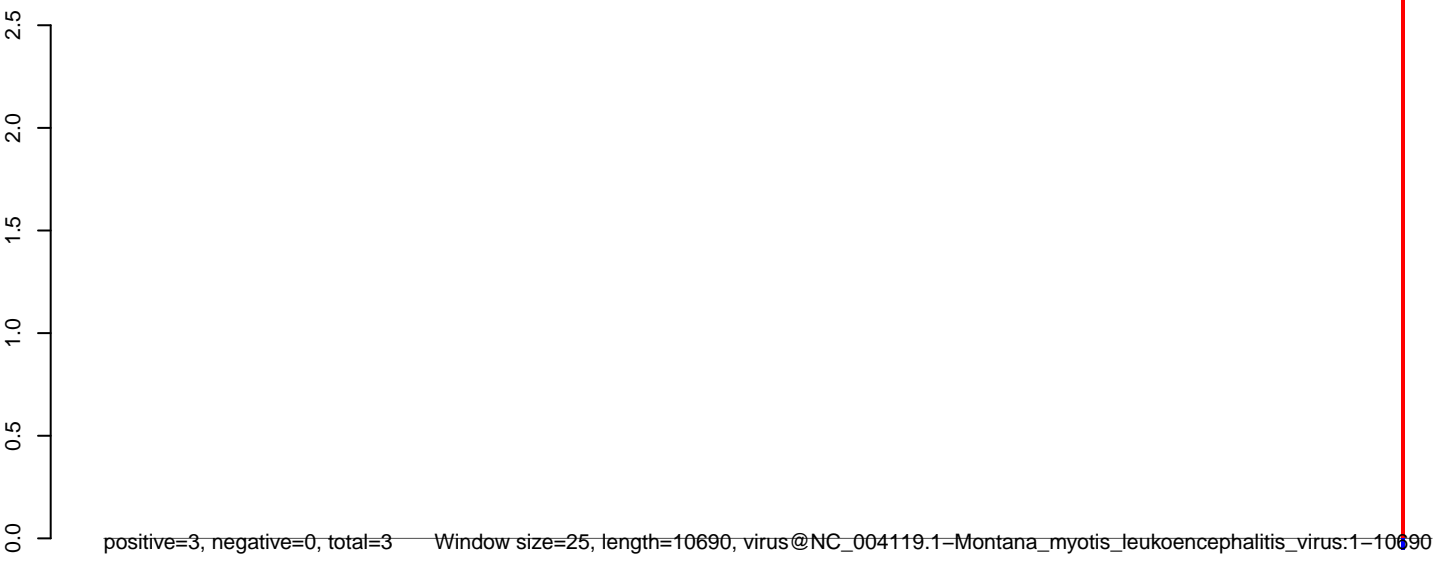
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

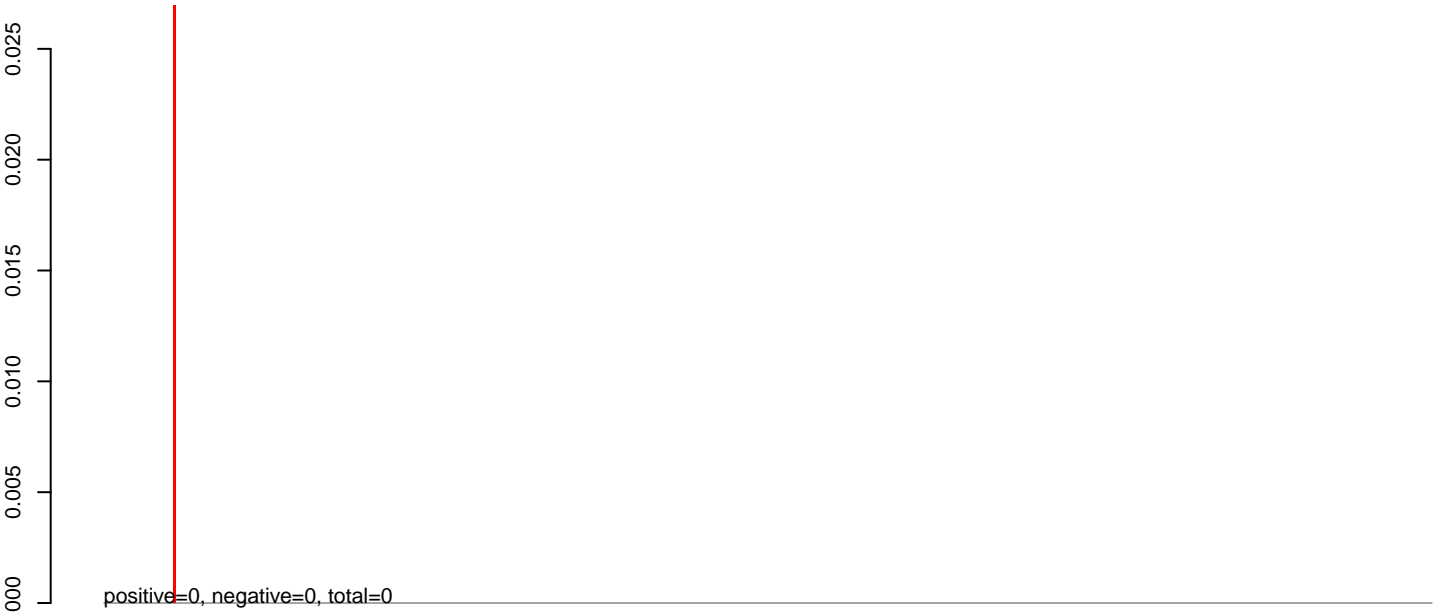


AeAeg_Aag2DENV2NGC.rep

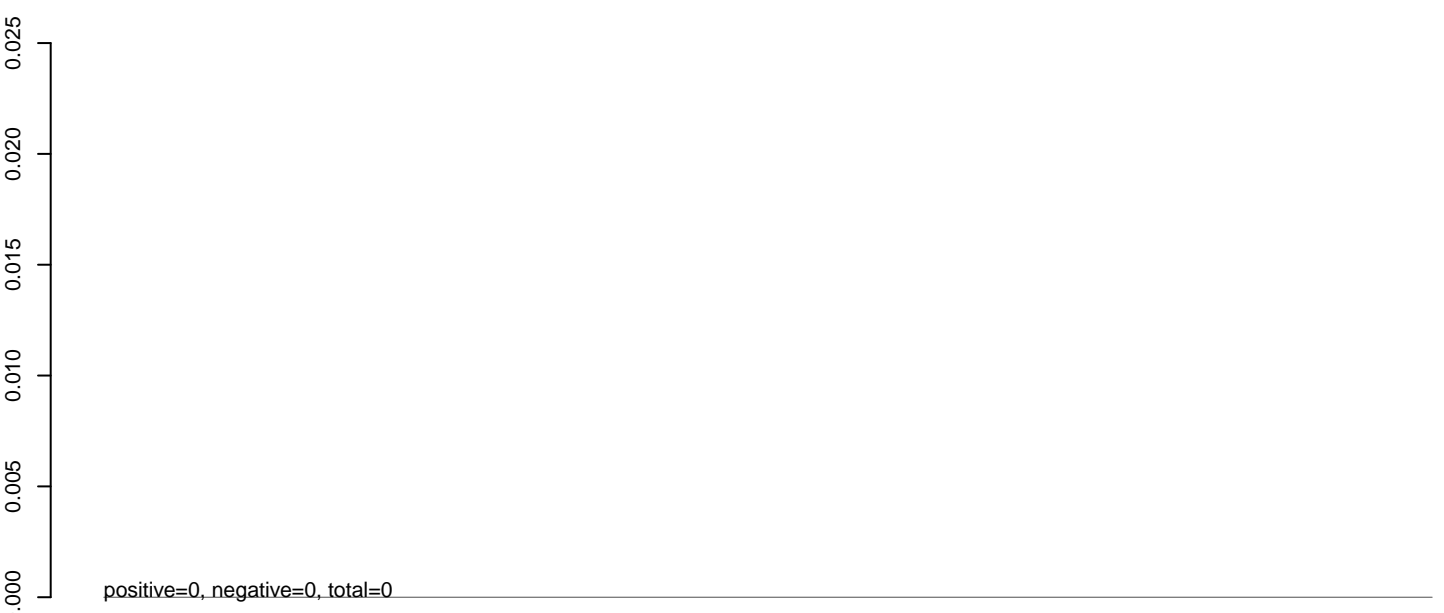


0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep



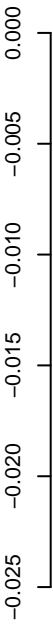
AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep

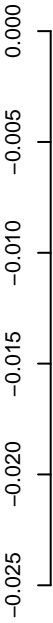


AeAeg_Aag2DENV2NGC.18_23.rep



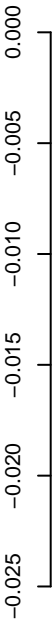
positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.24_35.rep



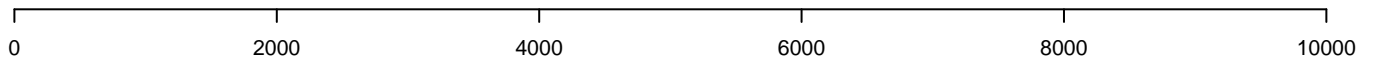
positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.rep

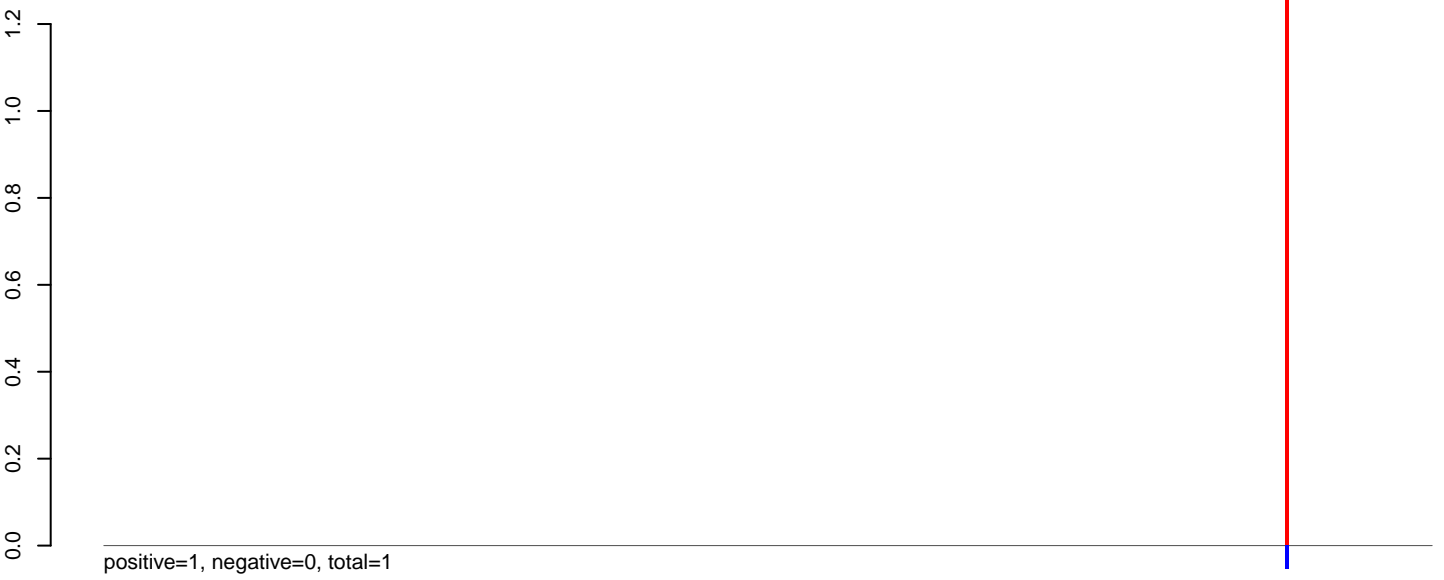


positive=0, negative=0, total=0

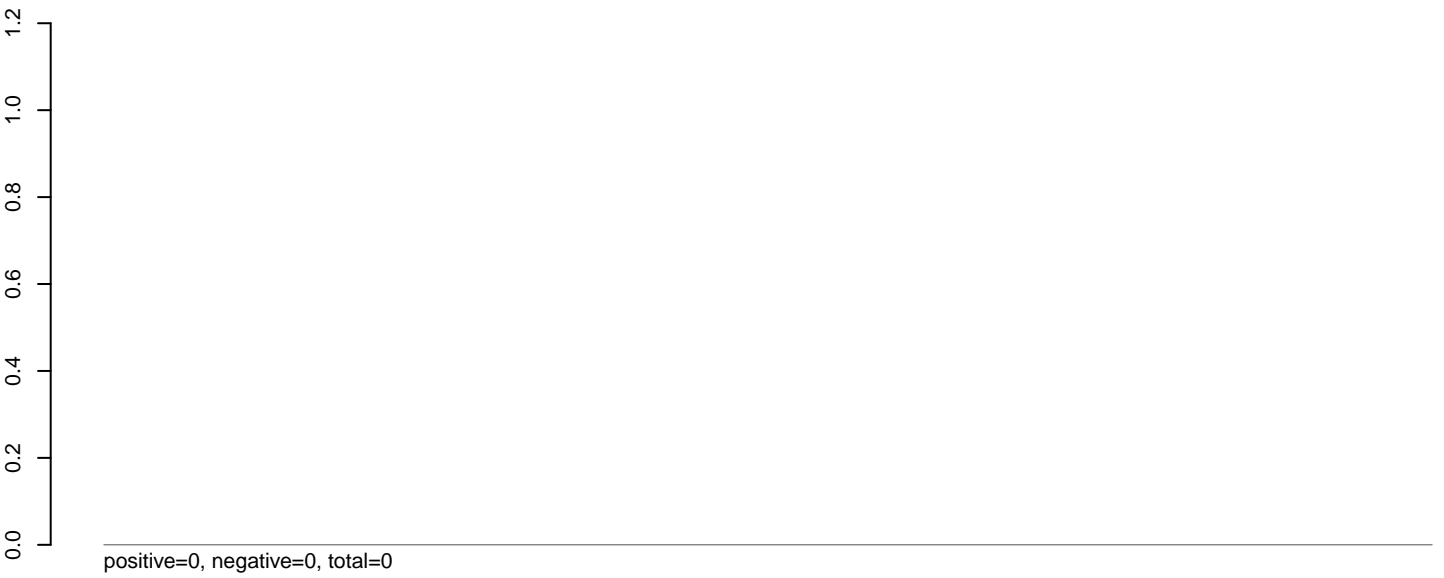
Window size=25, length=10116, virus@NC_003676.1-Apoi_virus_genome:1-10116



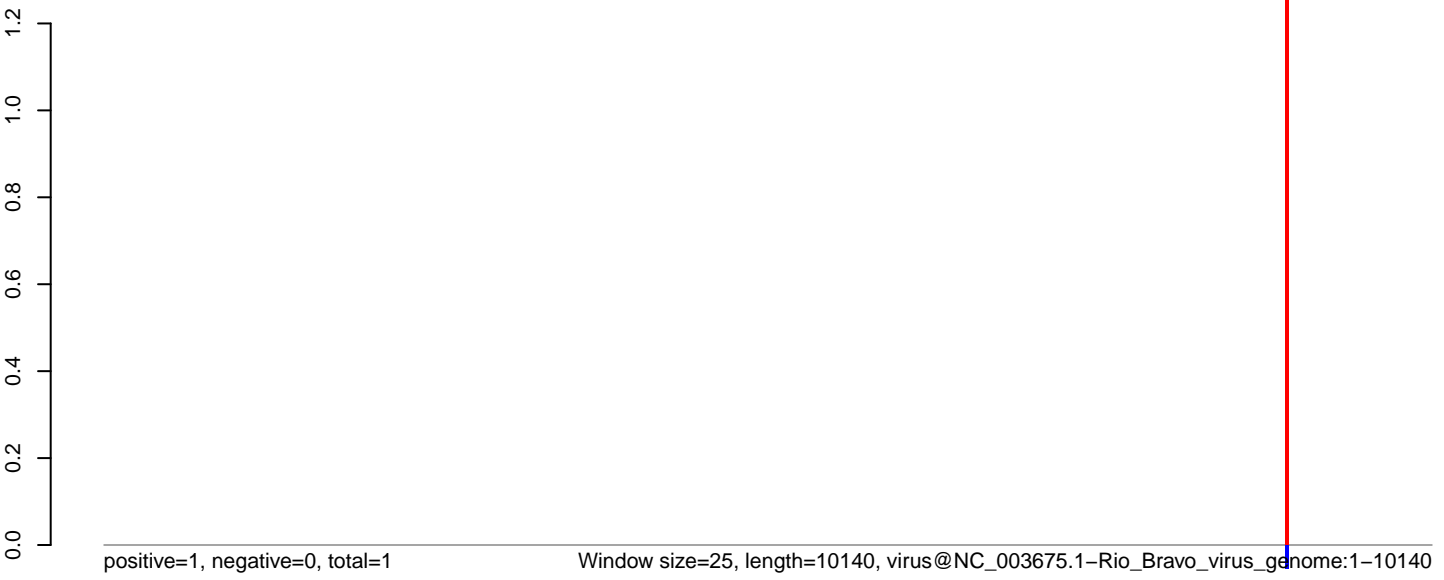
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



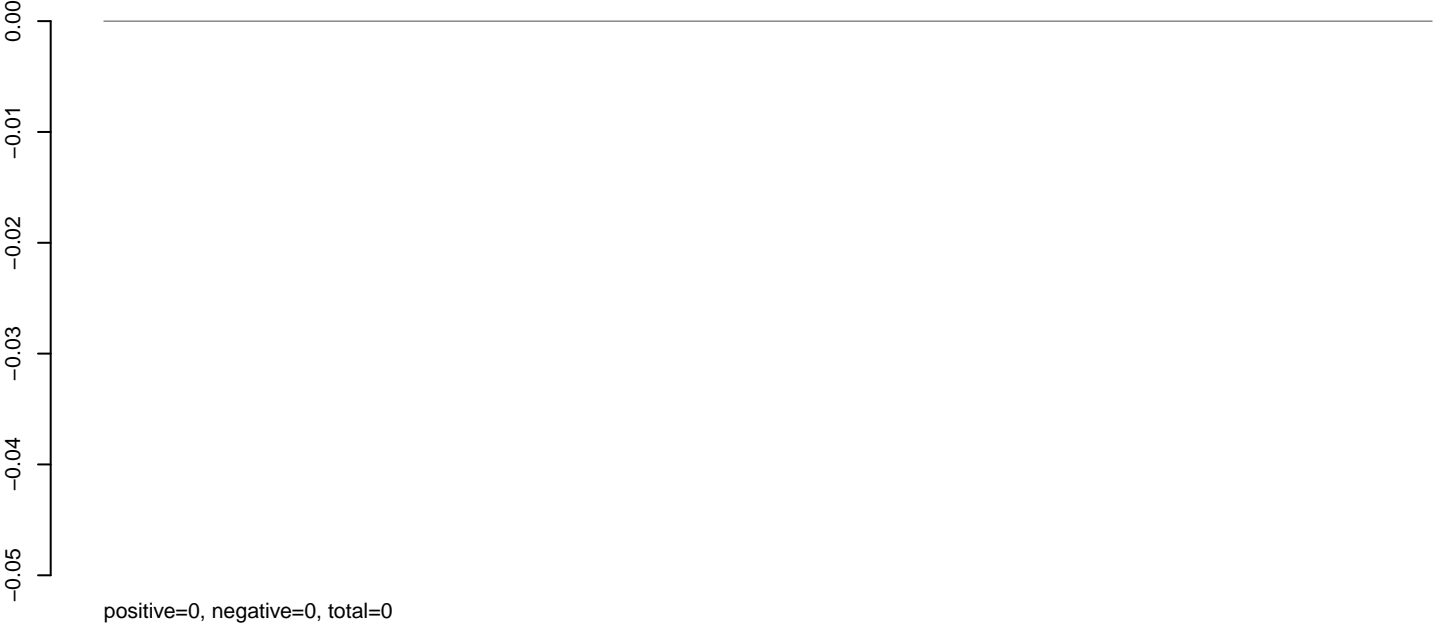
Window size=25, length=10140, virus@NC_003675.1-Rio_Bravo_virus_genome:1-10140

0 2000 4000 6000 8000 10000

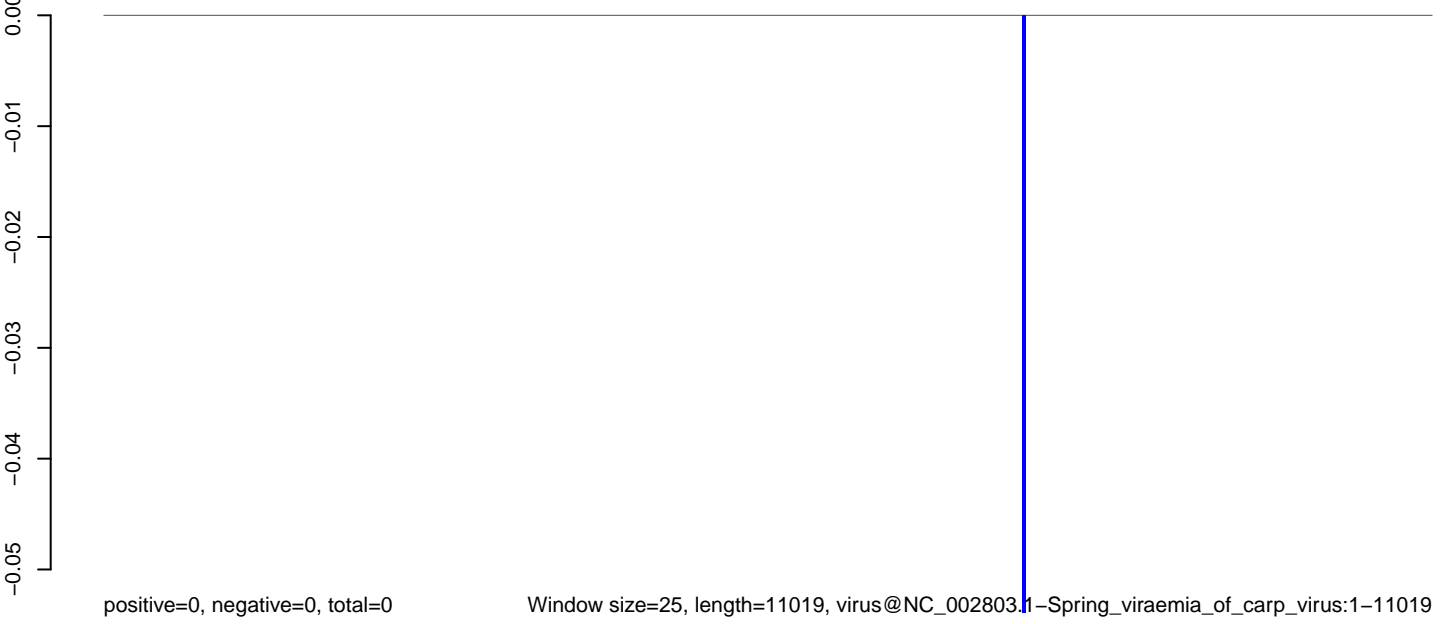
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



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

0 2000 4000 6000 8000 10000

AeAeg_Aag2DENV2NGC.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

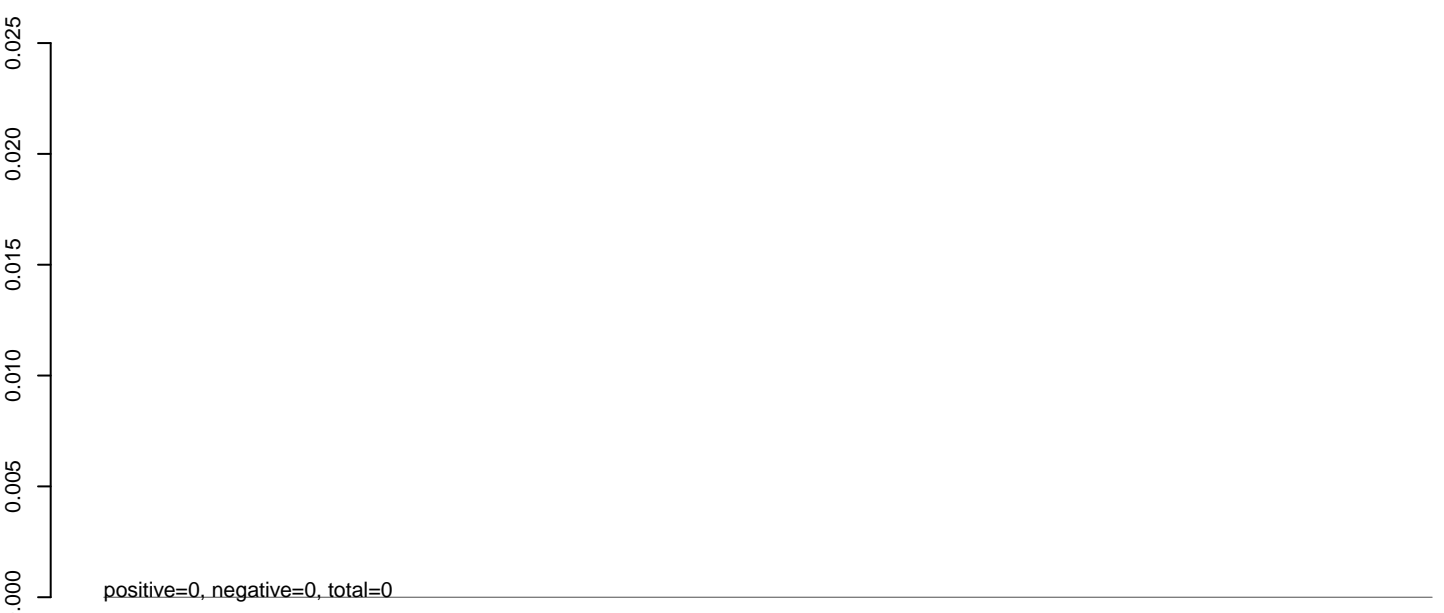
positive=0, negative=0, total=0
Window size=25, length=12294, virus@NC_001927.1-Bunyamwera_virus_segment_S_plus_segment_M_plus_segment_L:1-12294

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

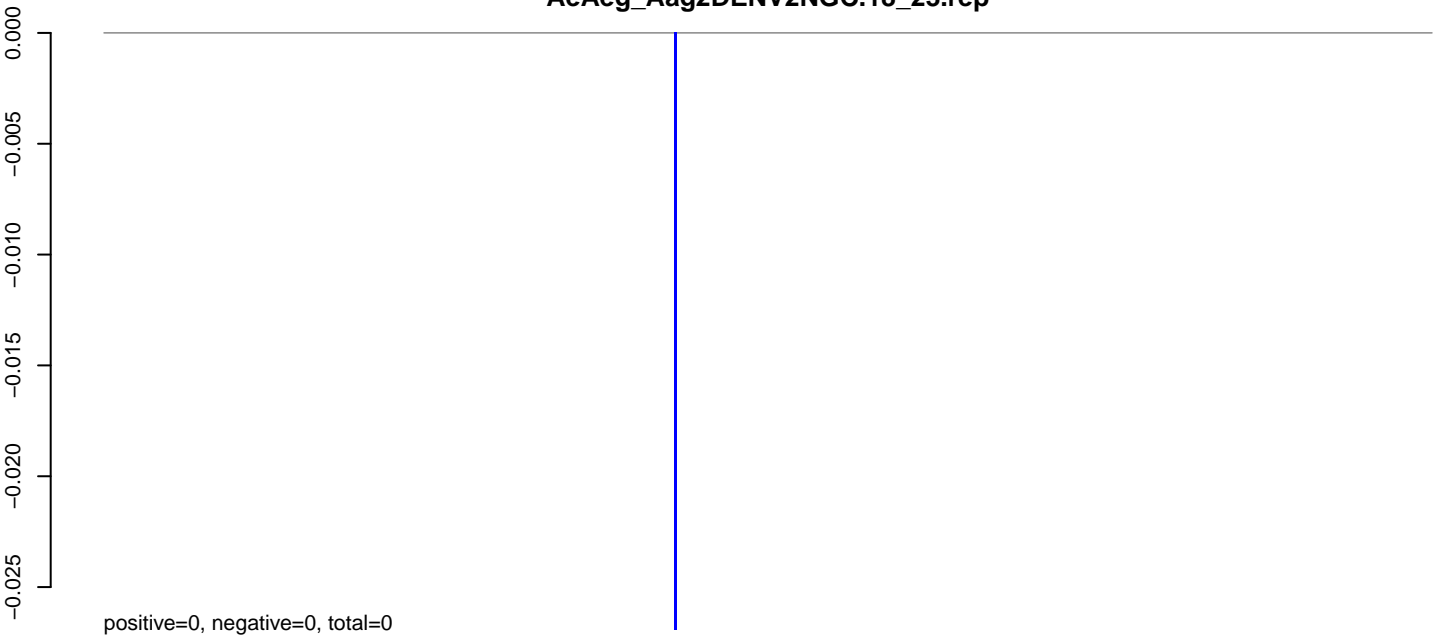


AeAeg_Aag2DENV2NGC.rep



Window size=25, length=11444, virus@NC_001449.1-Venezuelan_equine_encephalitis_virus:1-11444

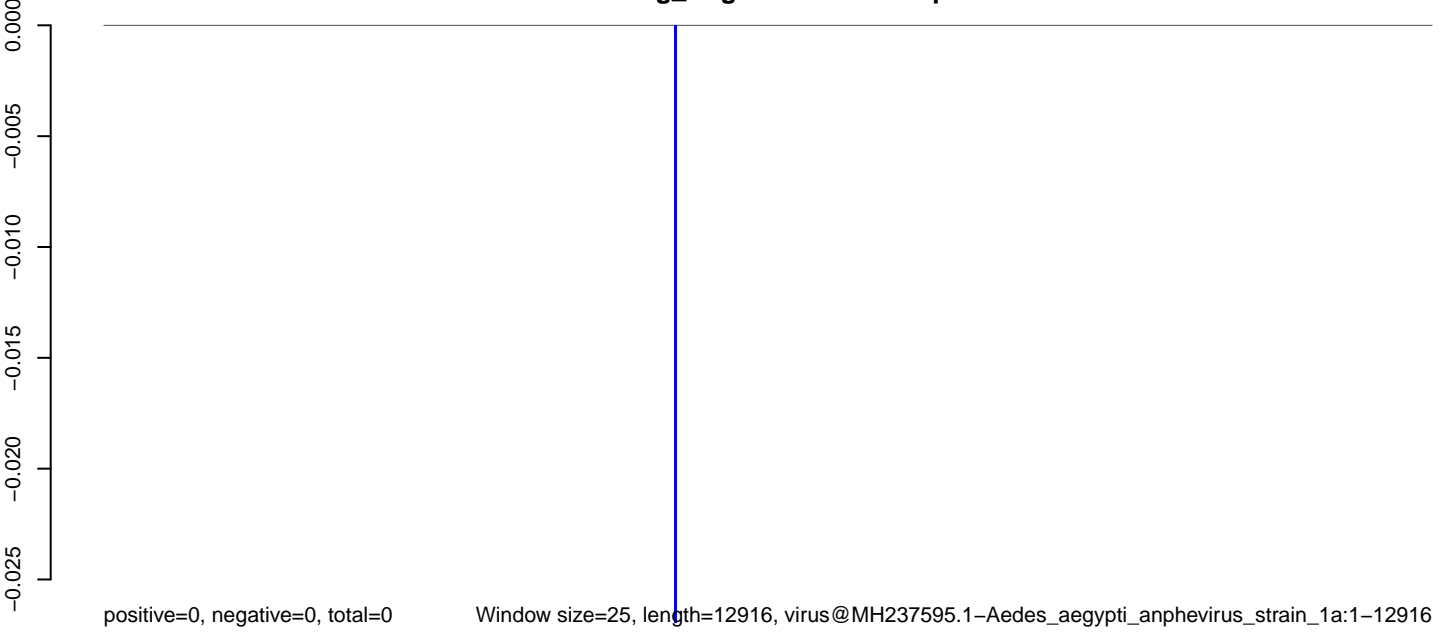
AeAeg_Aag2DENV2NGC.18_23.rep



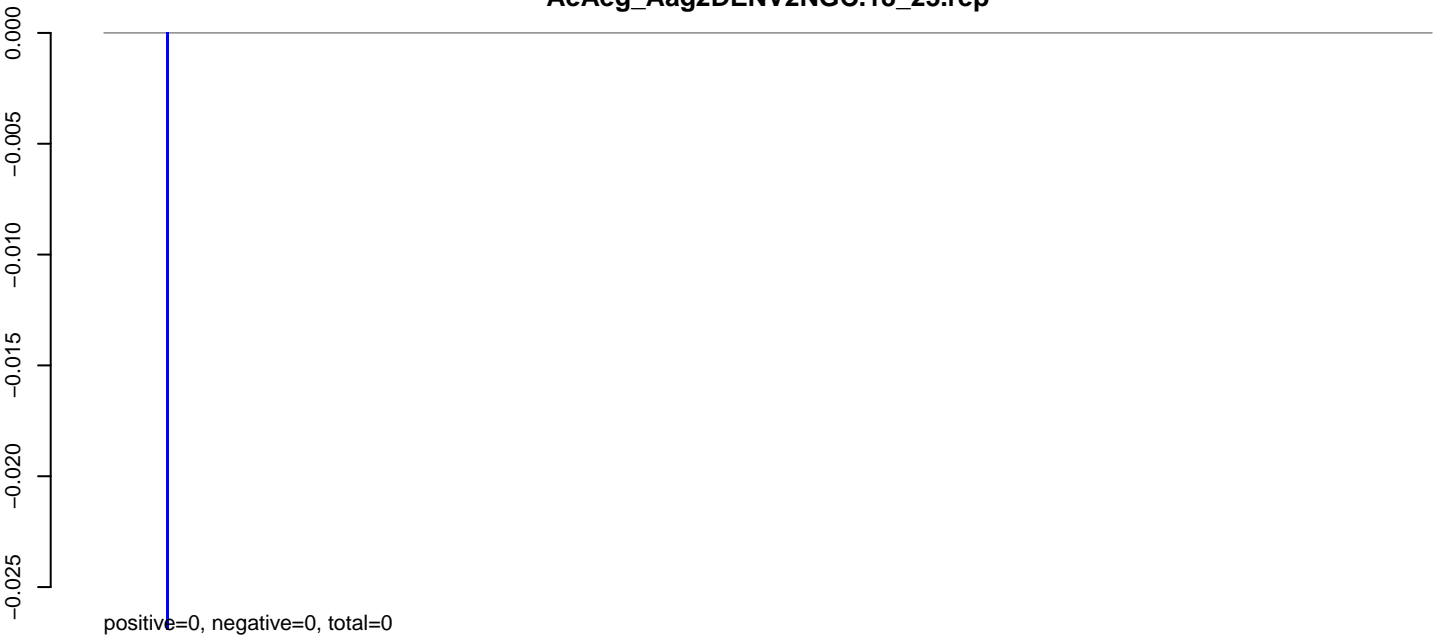
AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



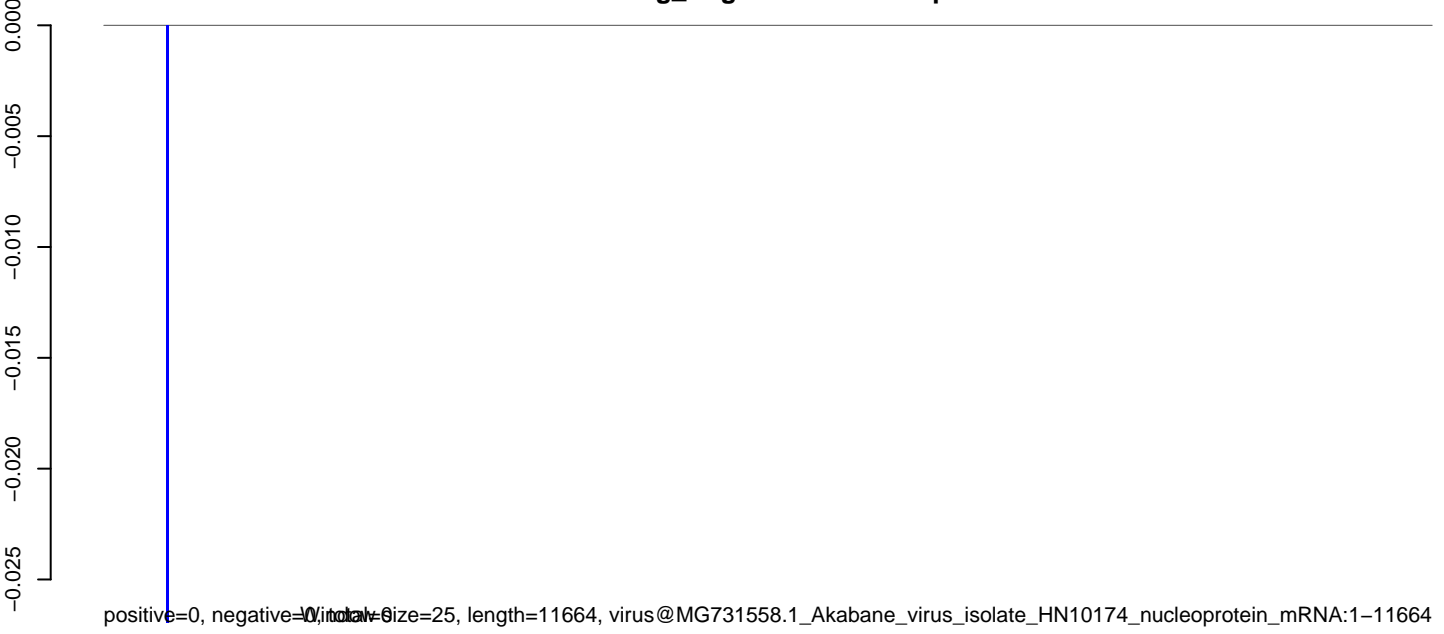
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



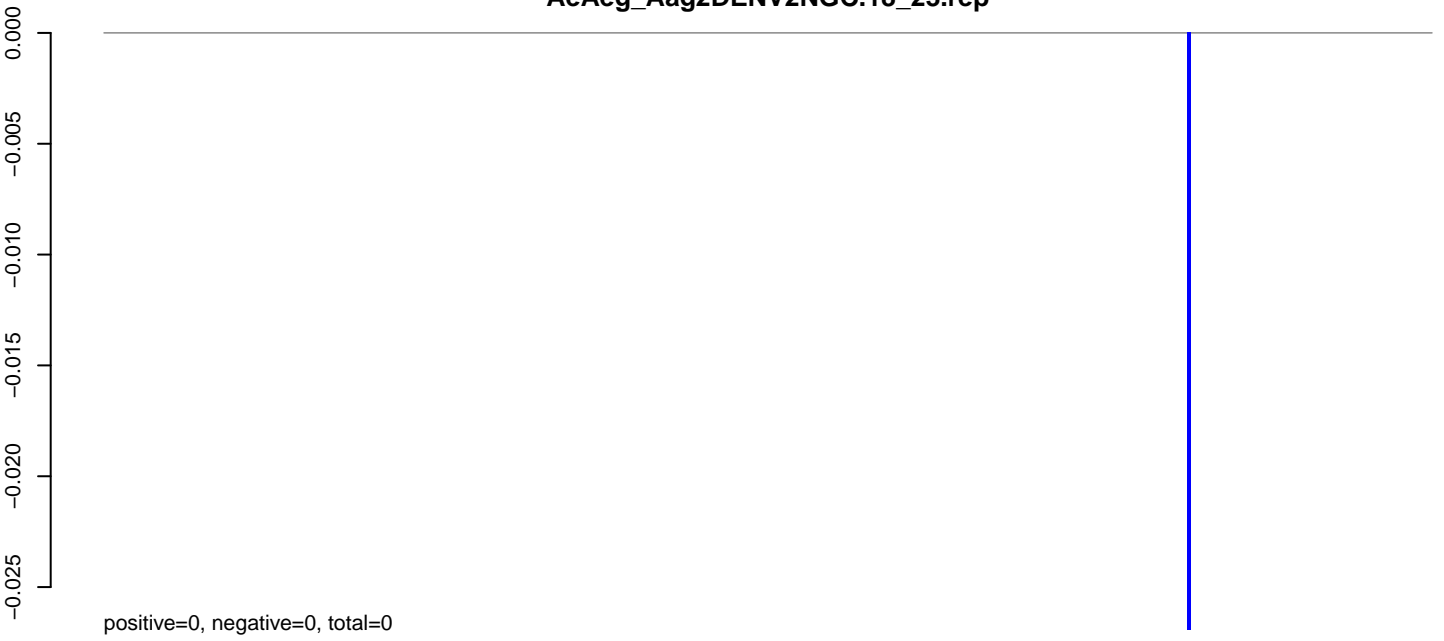
AeAeg_Aag2DENV2NGC.rep



positive=0, negative=0, total=6, size=25, length=11664, virus@MG731558.1_Akabane_virus_isolate_HN10174_nucleoprotein_mRNA:1-11664

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2DENV2NGC.18_23.rep



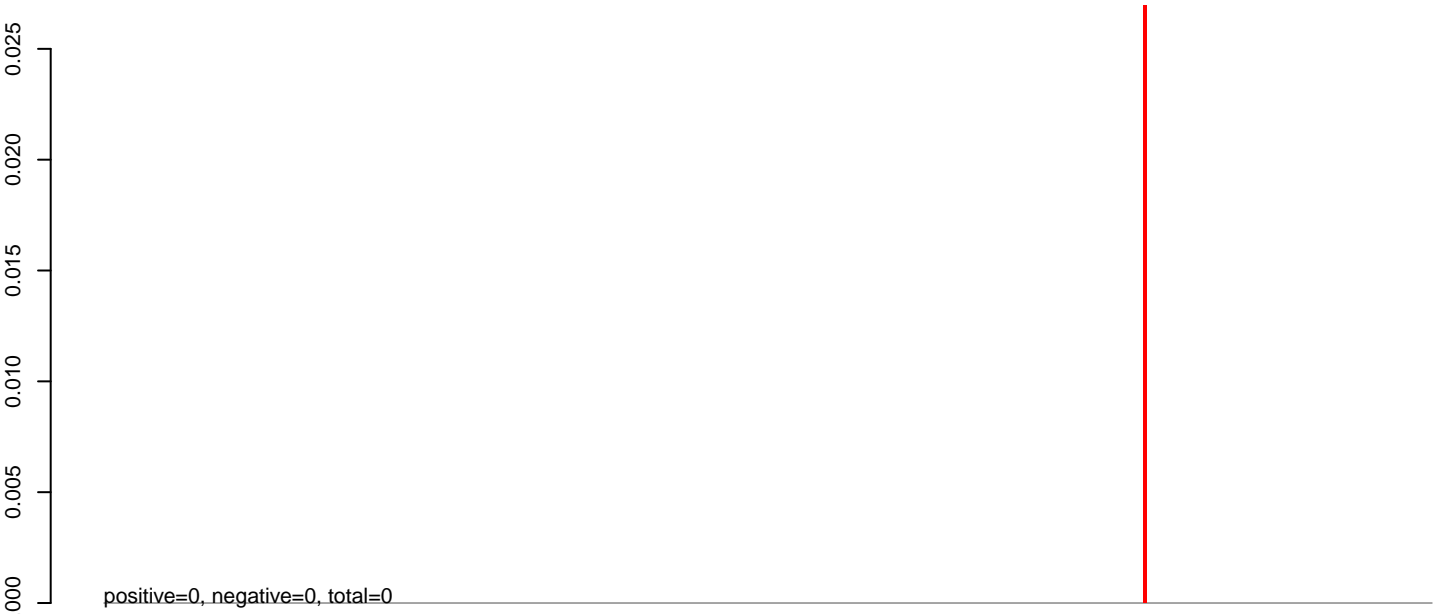
AeAeg_Aag2DENV2NGC.24_35.rep



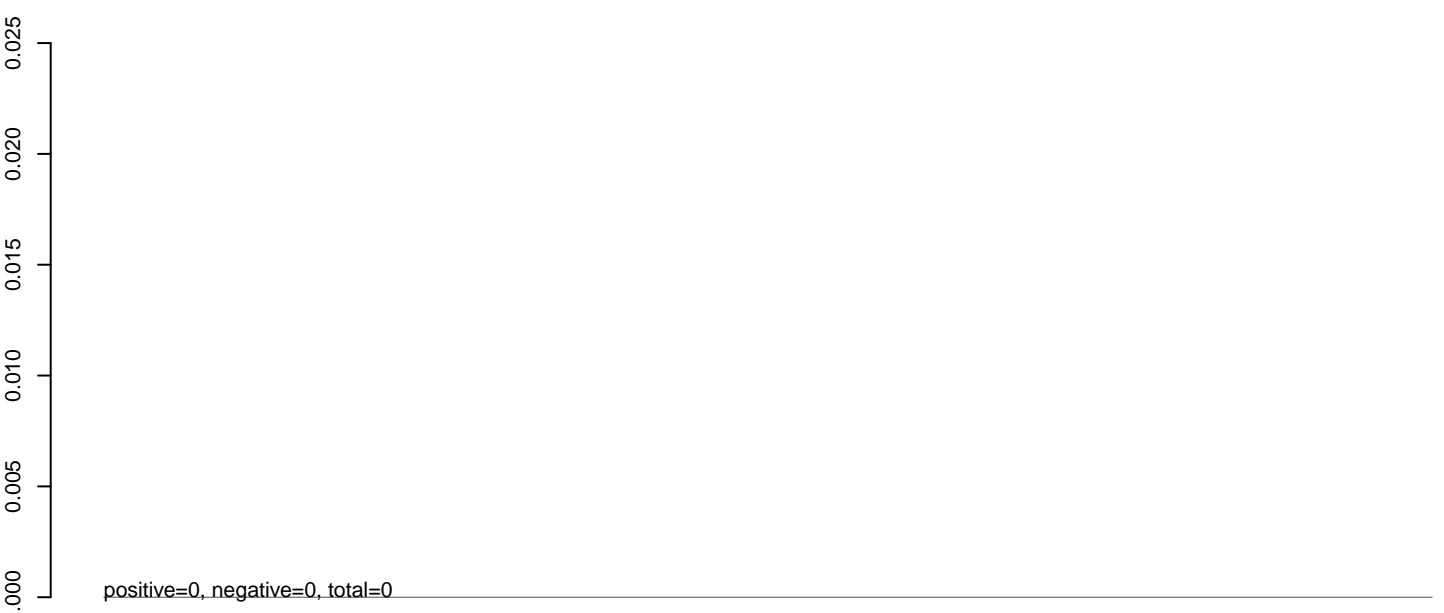
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep

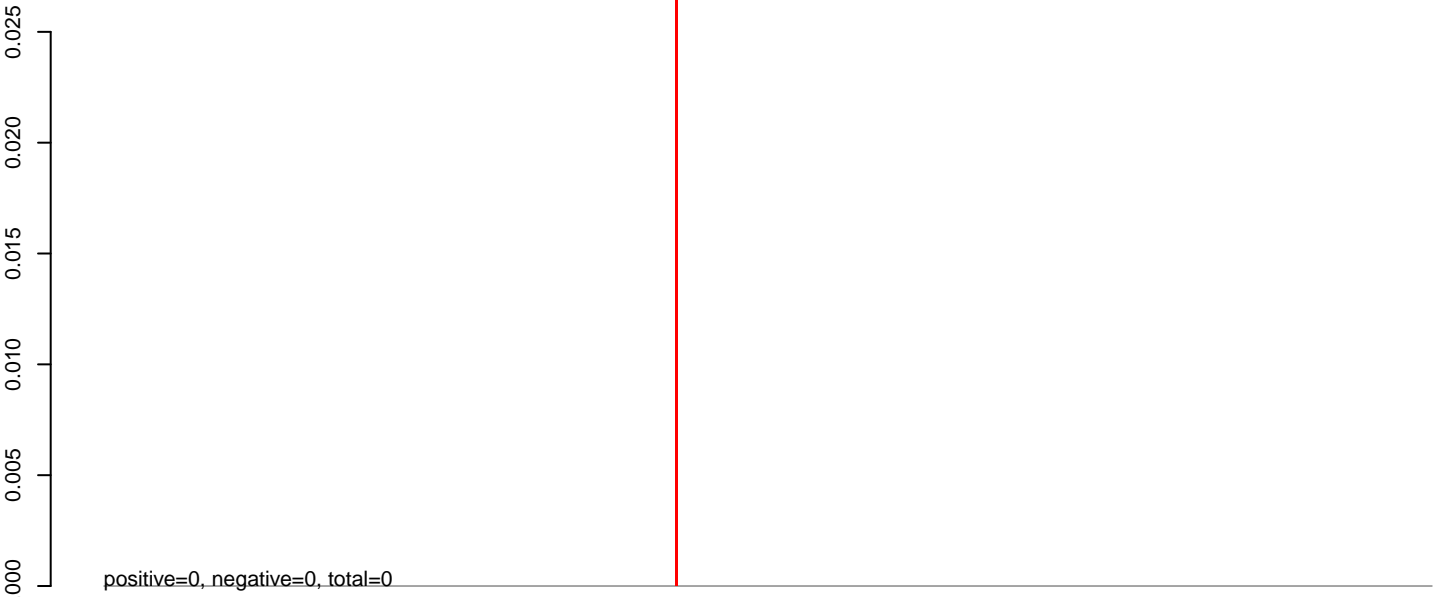


AeAeg_Aag2DENV2NGC.rep

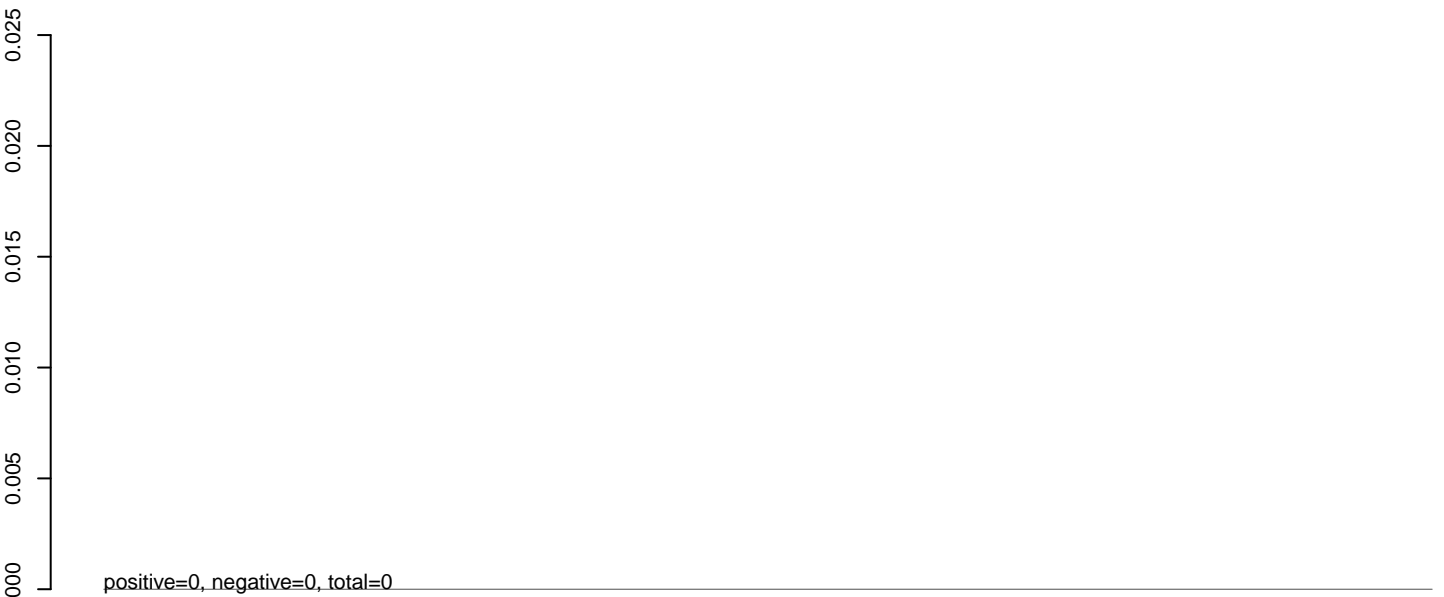


0 2000 4000 6000 8000 10000

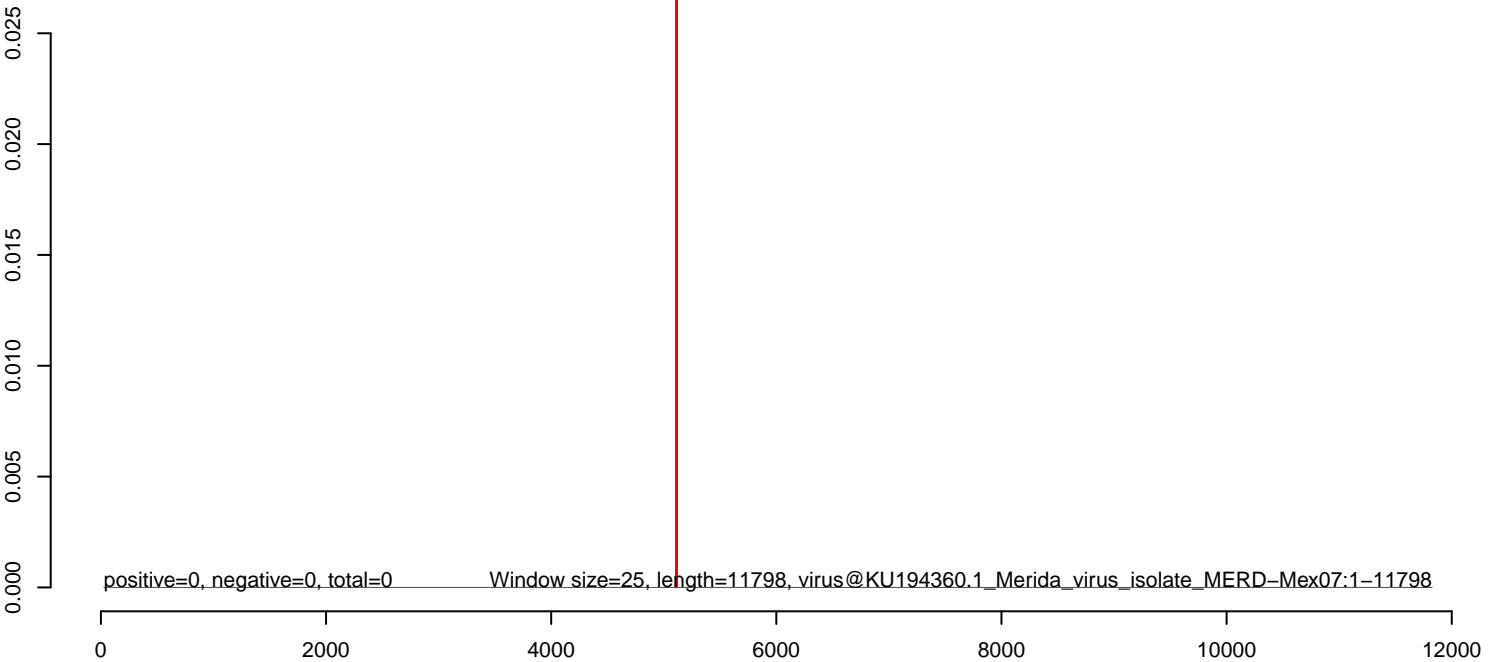
AeAeg_Aag2DENV2NGC.18_23.rep



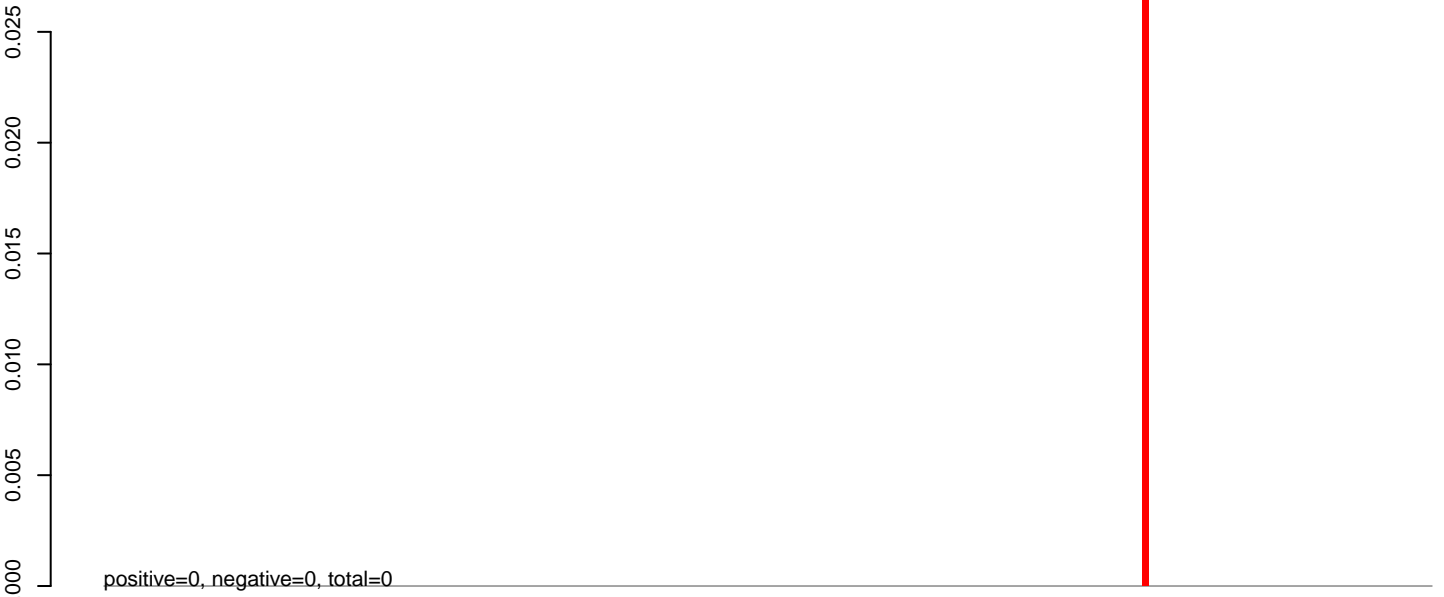
AeAeg_Aag2DENV2NGC.24_35.rep



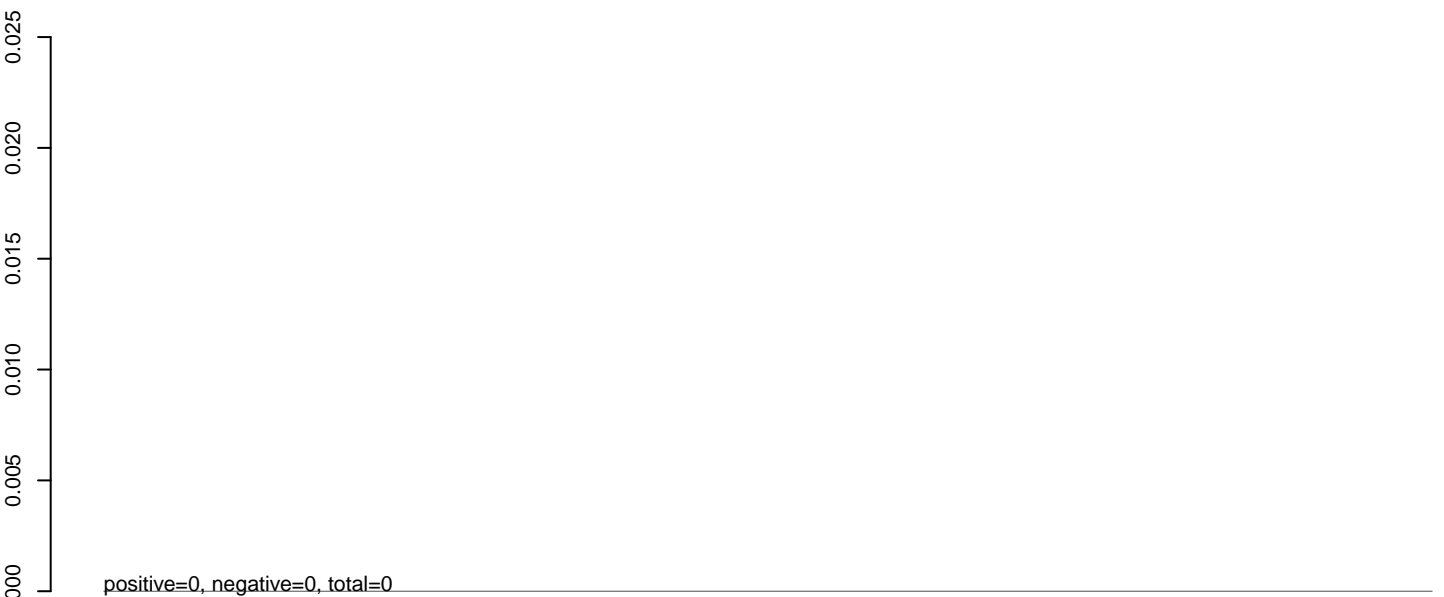
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



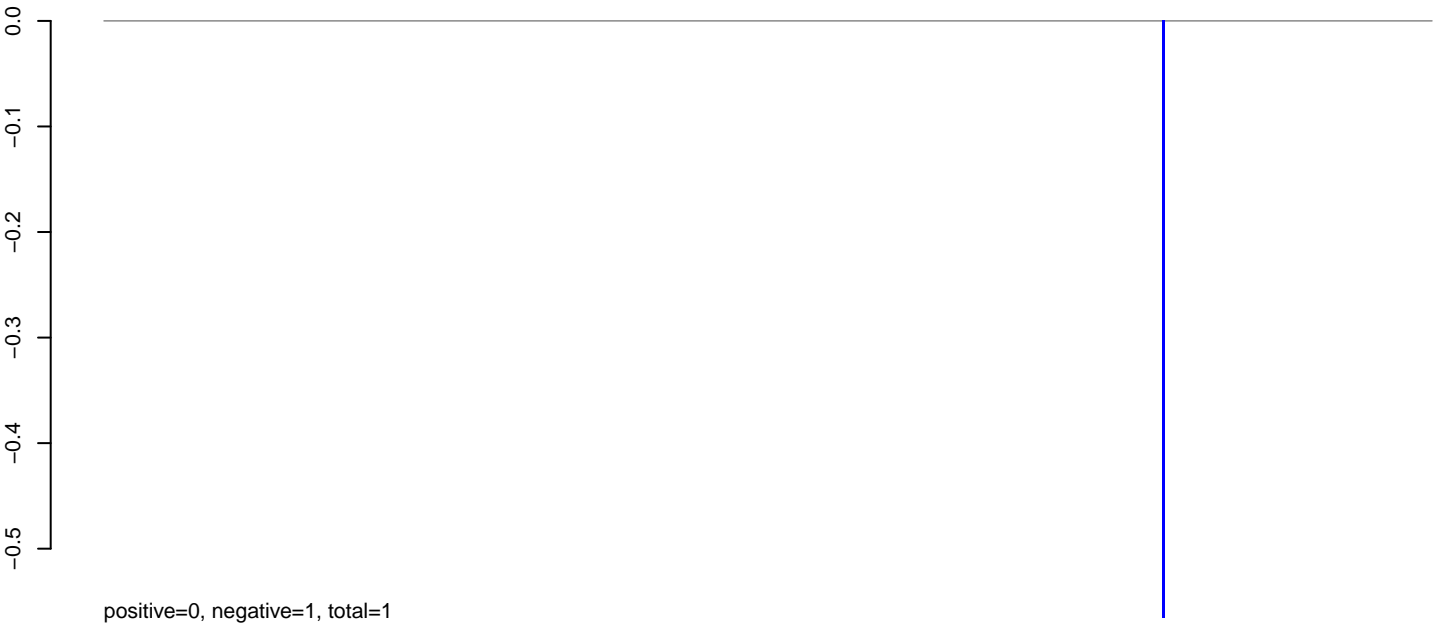
AeAeg_Aag2DENV2NGC.24_35.rep



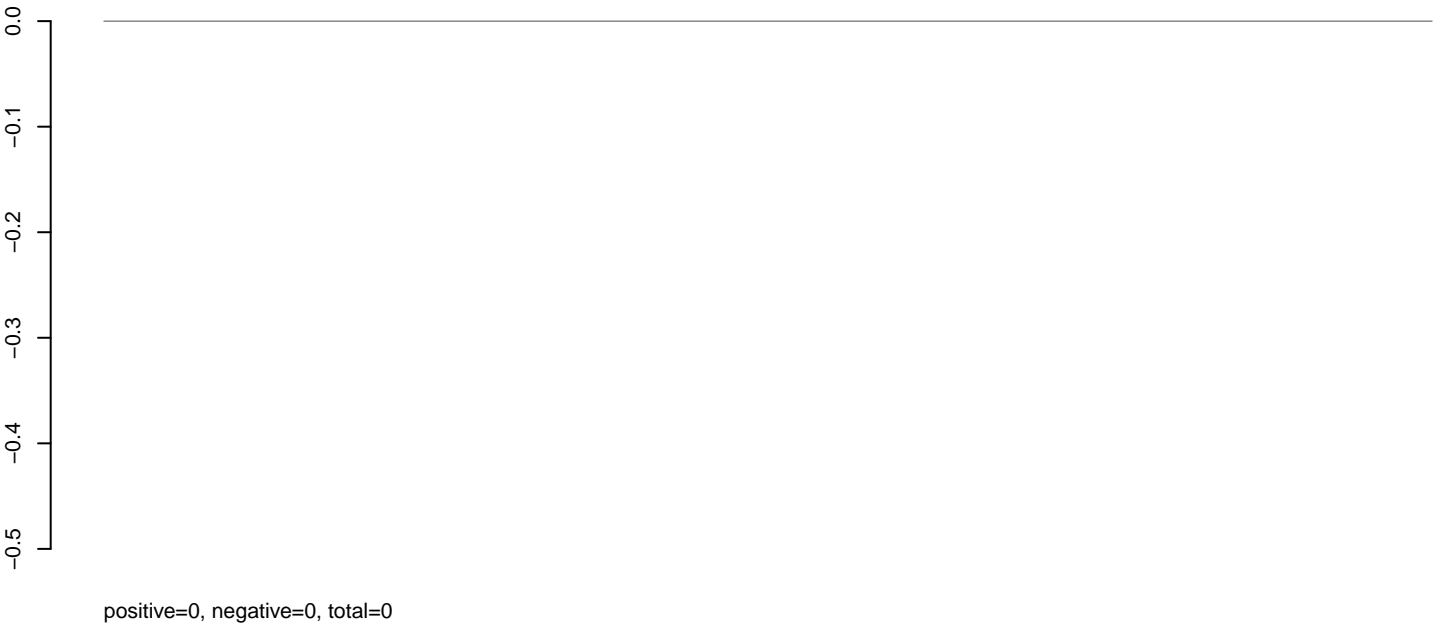
AeAeg_Aag2DENV2NGC.rep



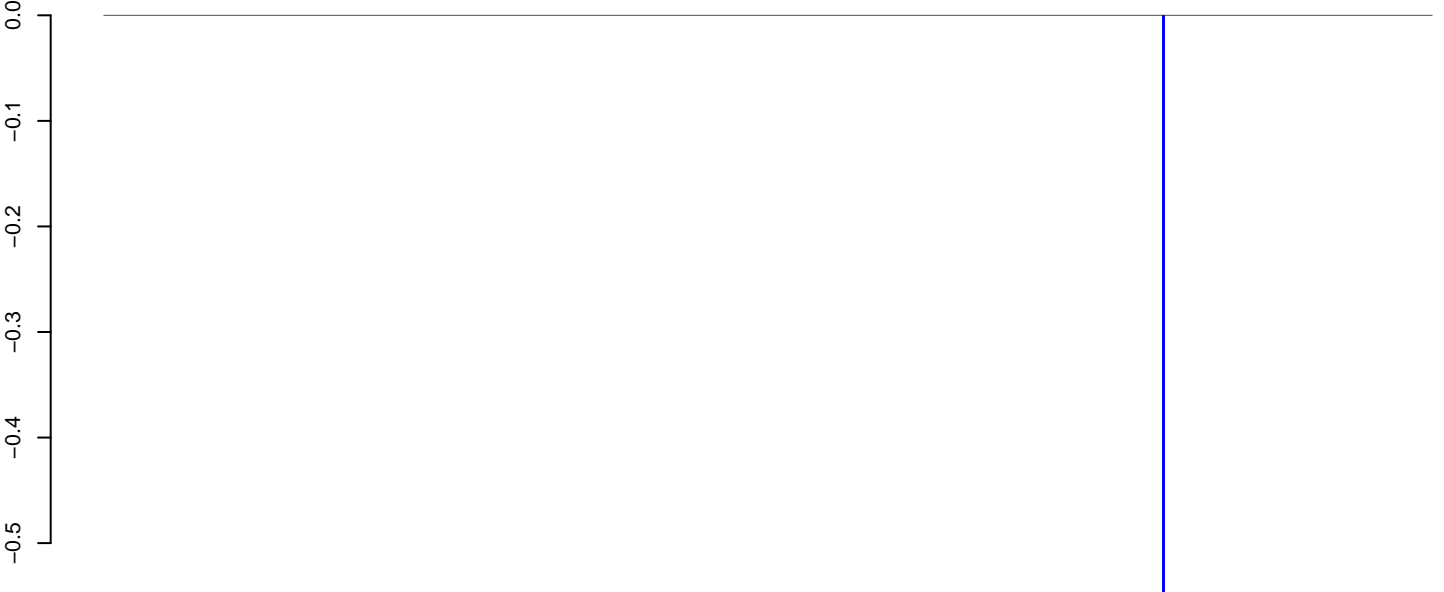
AeAeg_Aag2DENV2NGC.18_23.rep



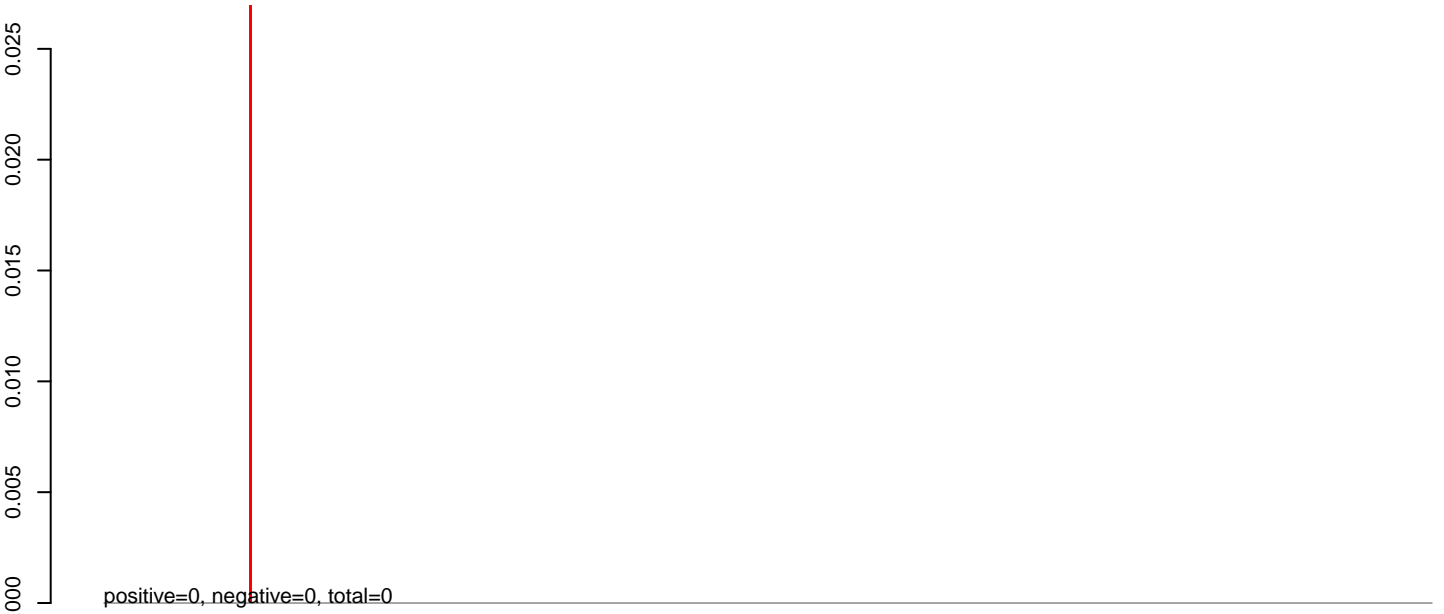
AeAeg_Aag2DENV2NGC.24_35.rep



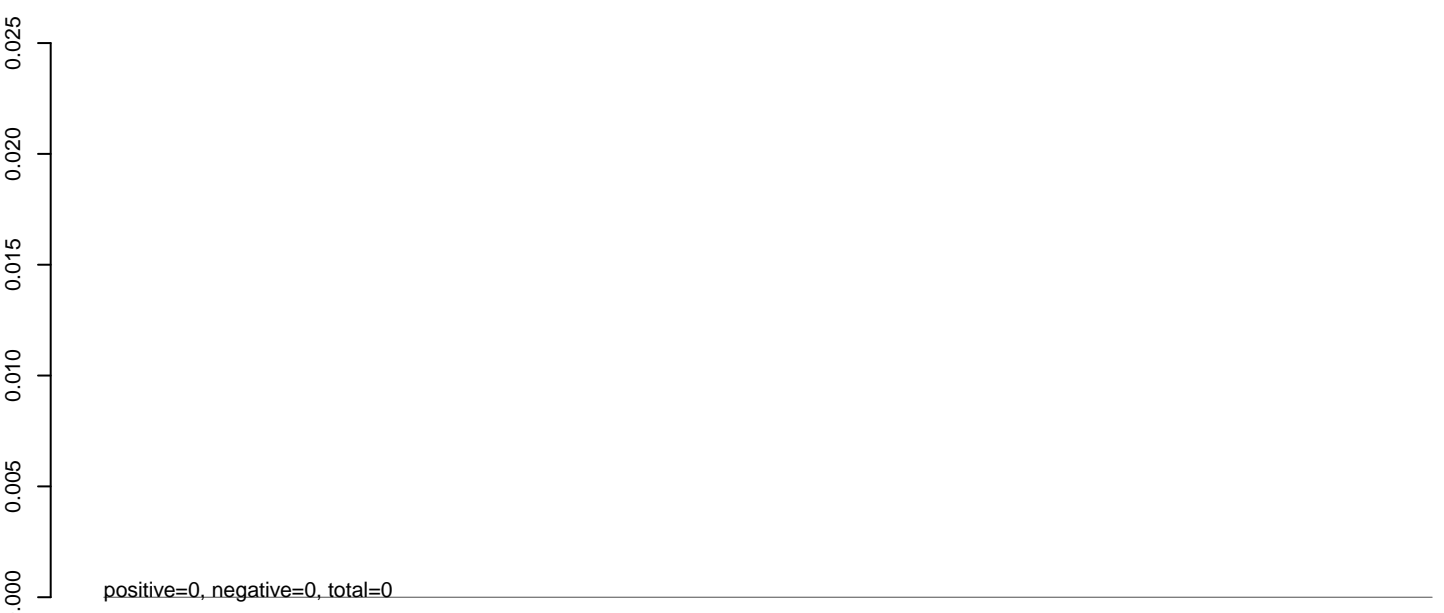
AeAeg_Aag2DENV2NGC.rep



AeAeg_Aag2DENV2NGC.18_23.rep



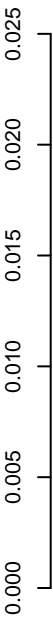
AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep

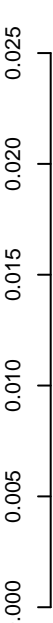


AeAeg_Aag2DENV2NGC.18_23.rep



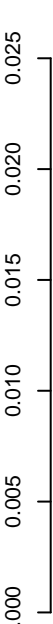
positive=0, negative=0, total=0

AeAeg_Aag2DENV2NGC.24_35.rep

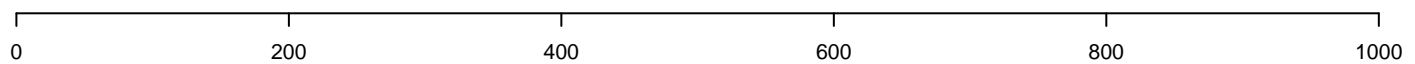


positive=0, negative=0, total=0

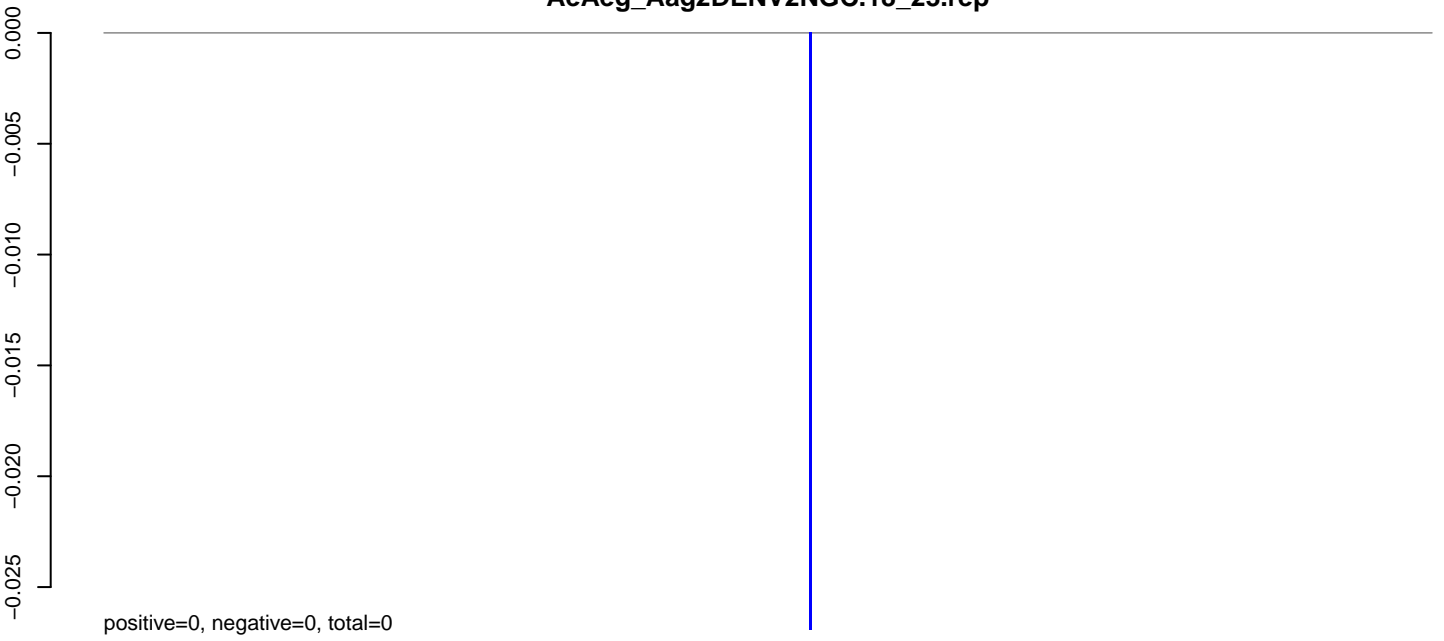
AeAeg_Aag2DENV2NGC.rep



positive=0, negative=0, total=26, length=972, virus@AF411844.1-Elaeis guineensis_MADS_box_transcription_factor_AGL2-2_mRNA:1-972



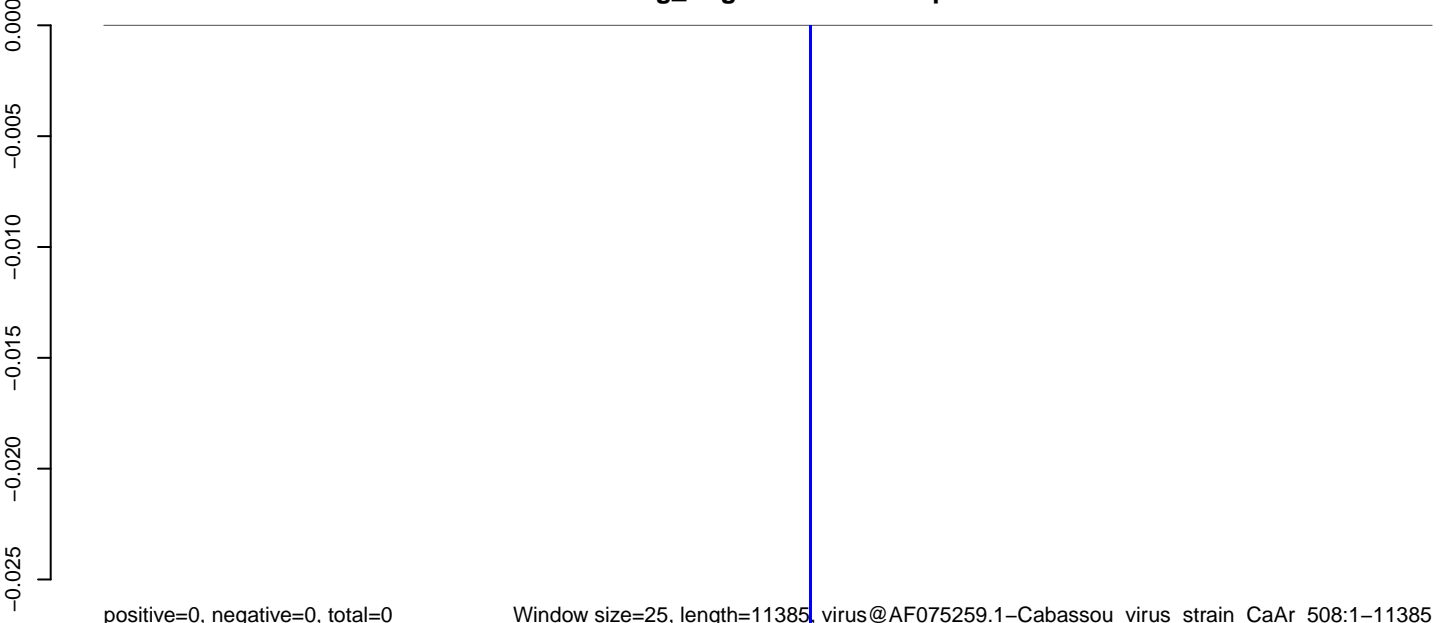
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



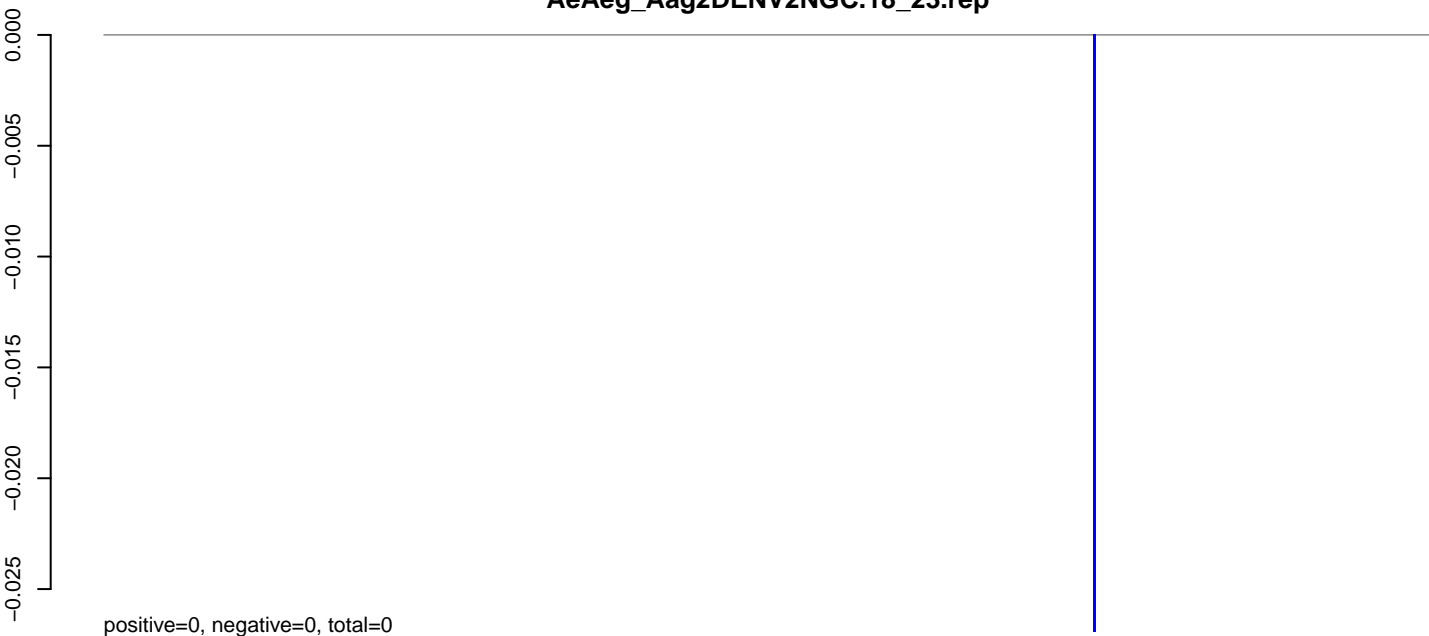
AeAeg_Aag2DENV2NGC.rep



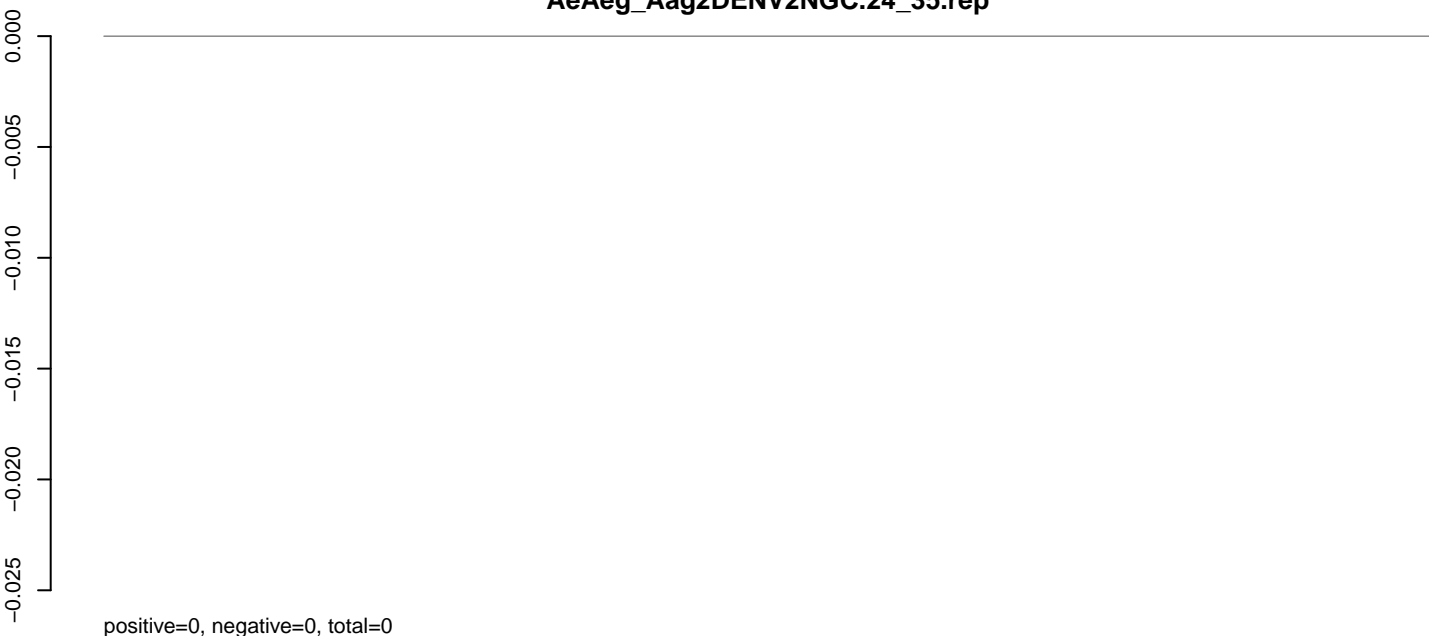
Window size=25, length=11385, virus@AF075259.1-Cabassou_virus_strain_CaAr_508:1-11385

0 2000 4000 6000 8000 10000

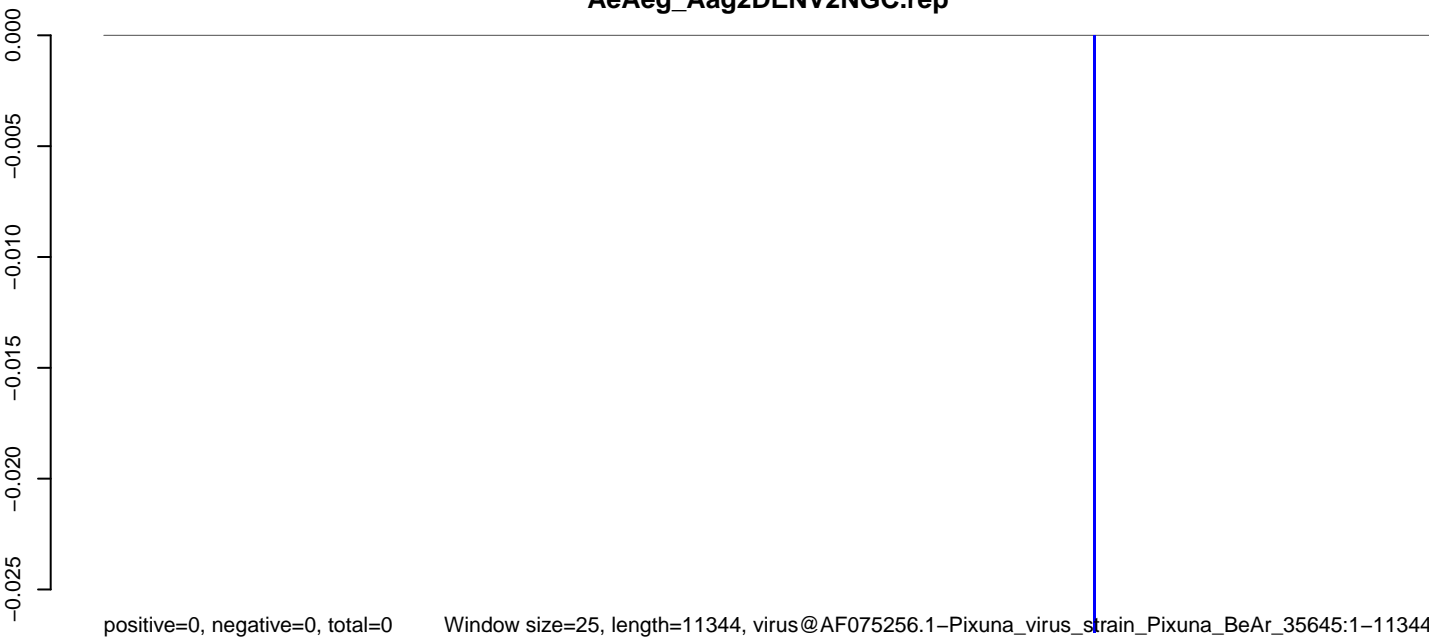
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



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

0 2000 4000 6000 8000 10000

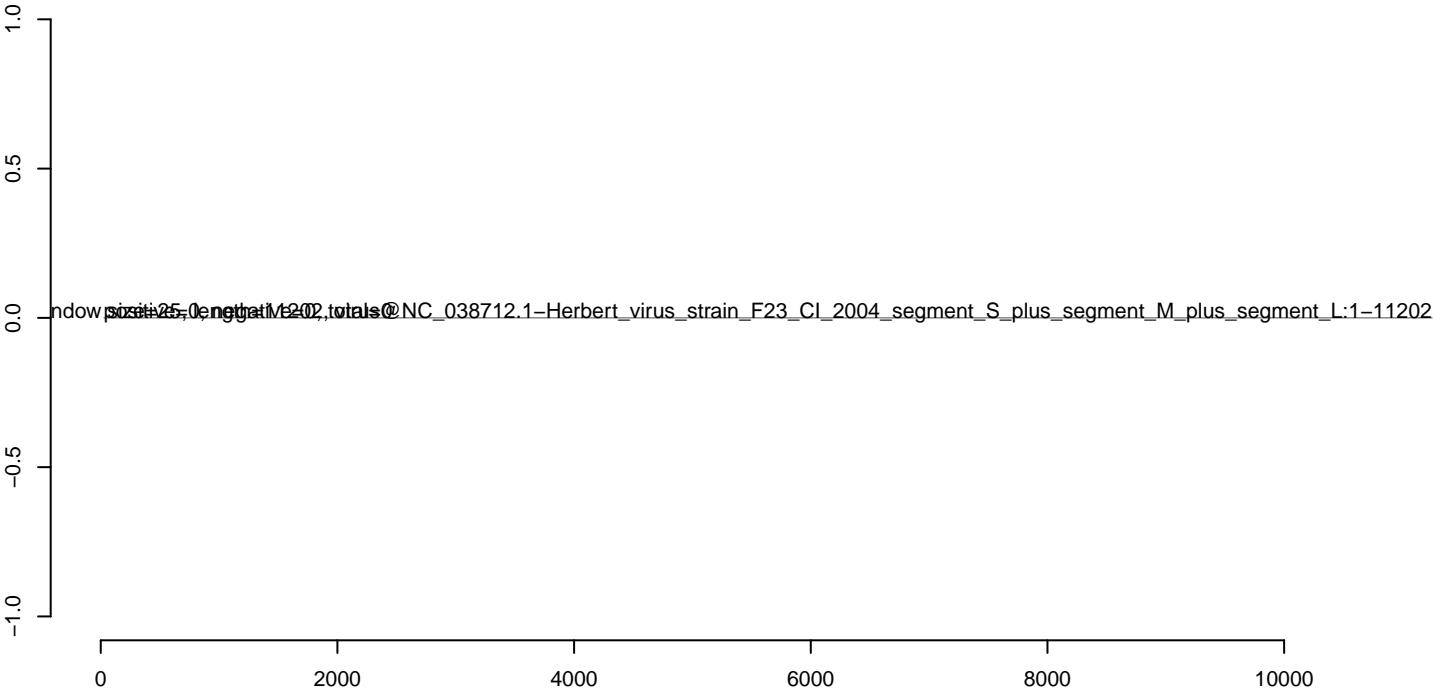
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



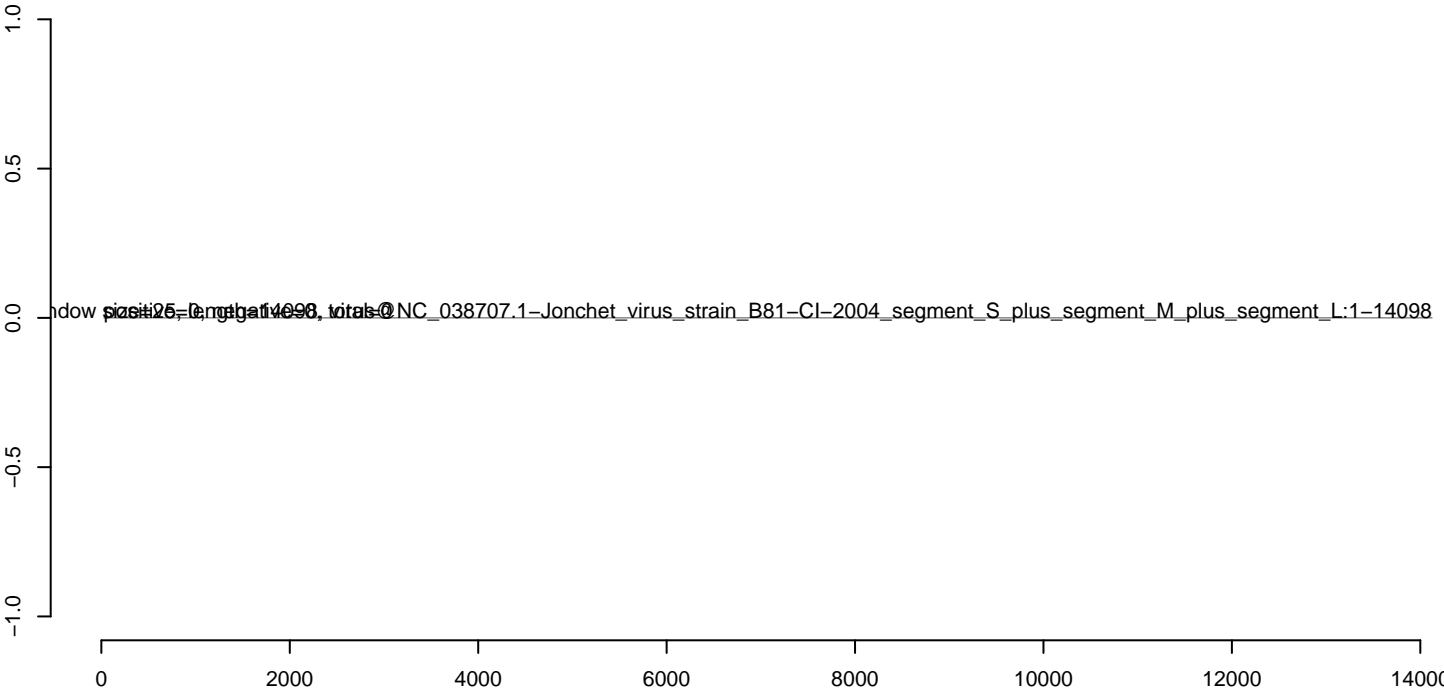
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



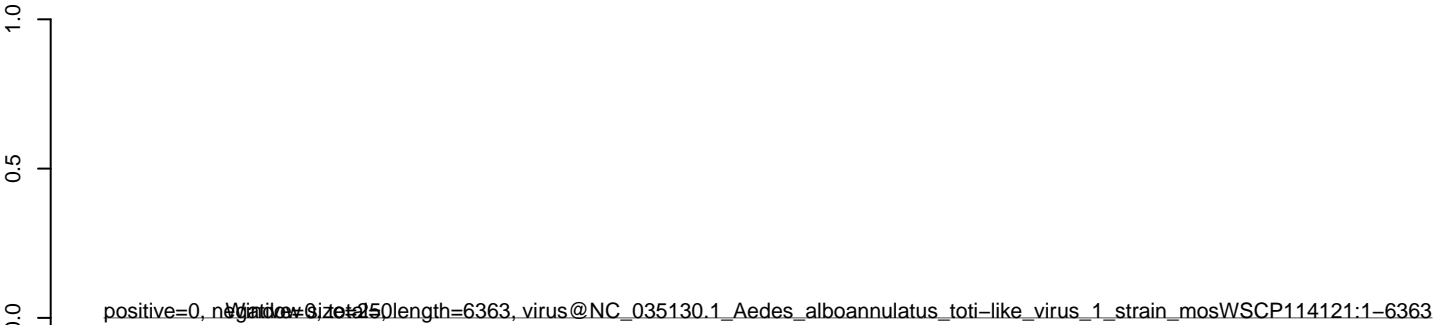
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



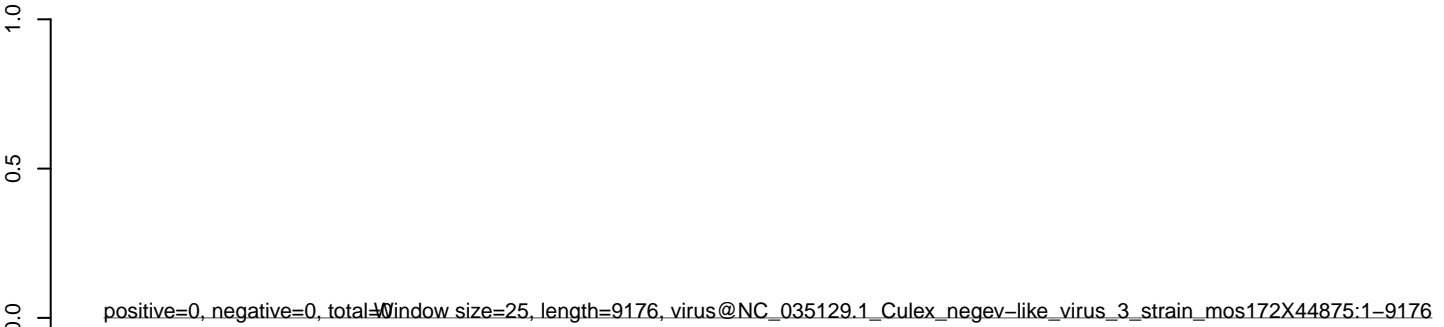
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



0 2000 4000 6000 8000

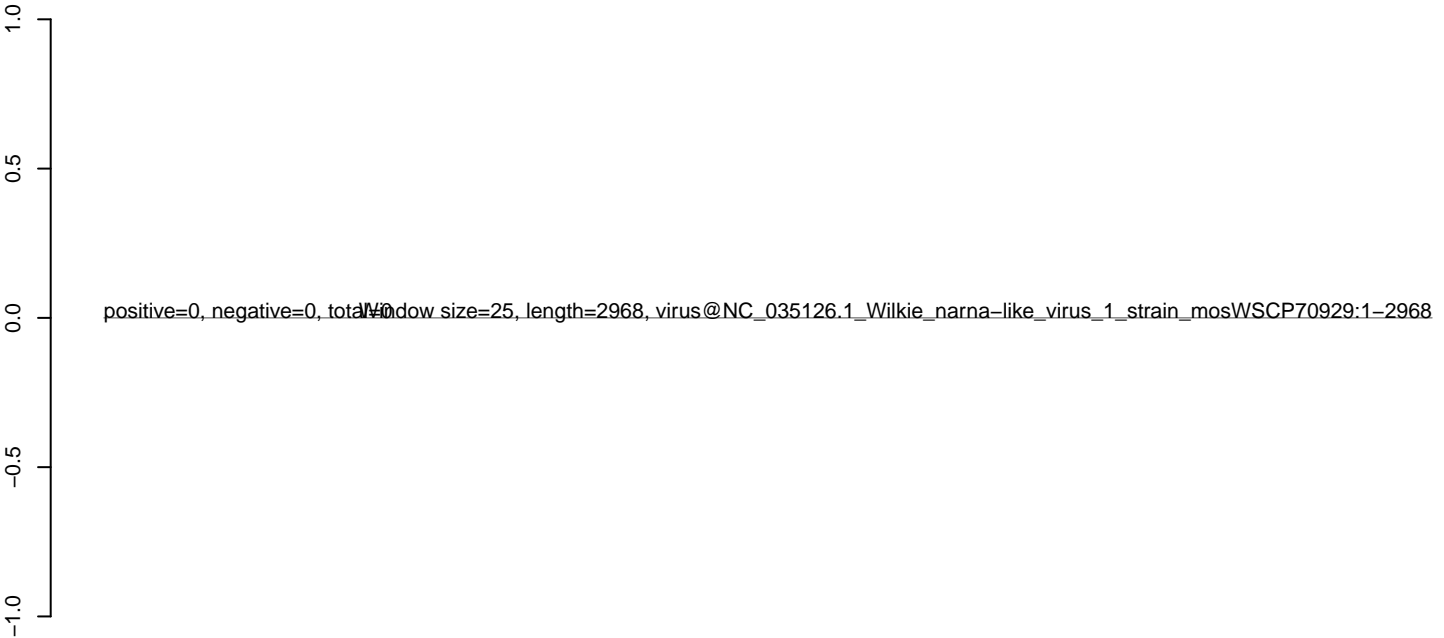
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



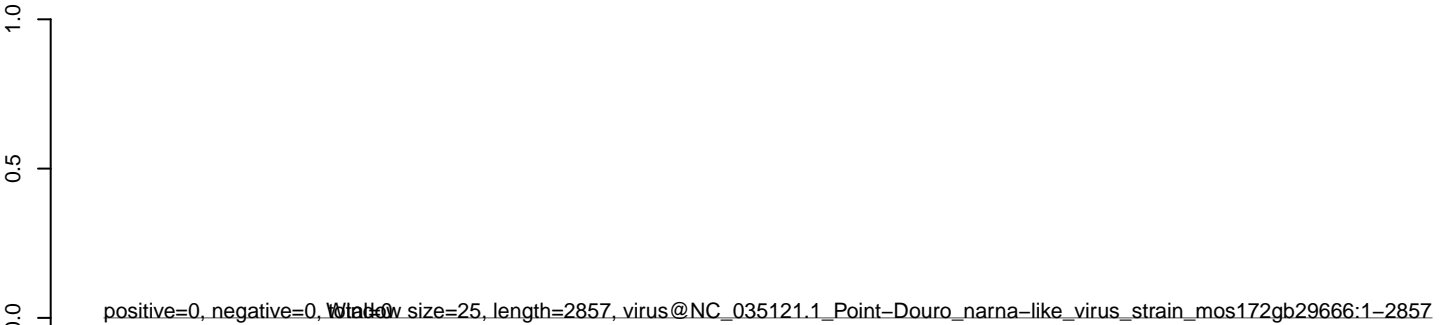
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



0 500 1000 1500 2000 2500 3000

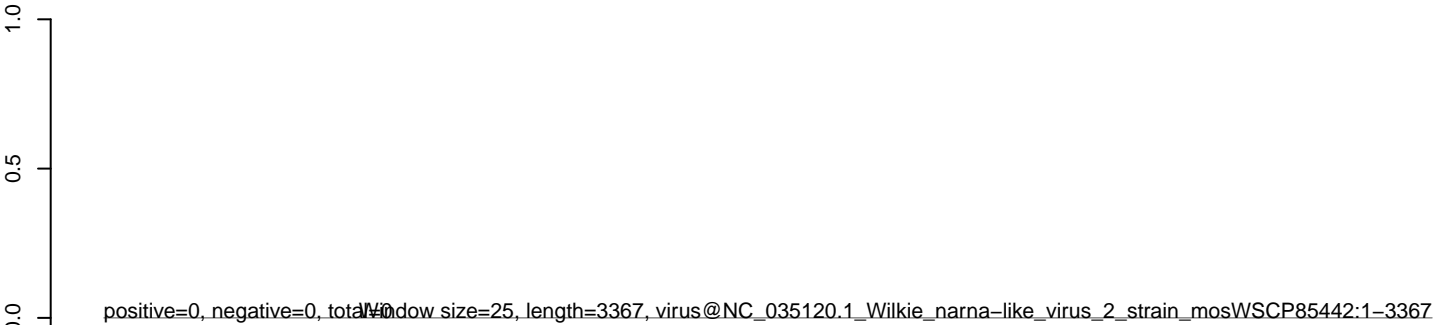
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



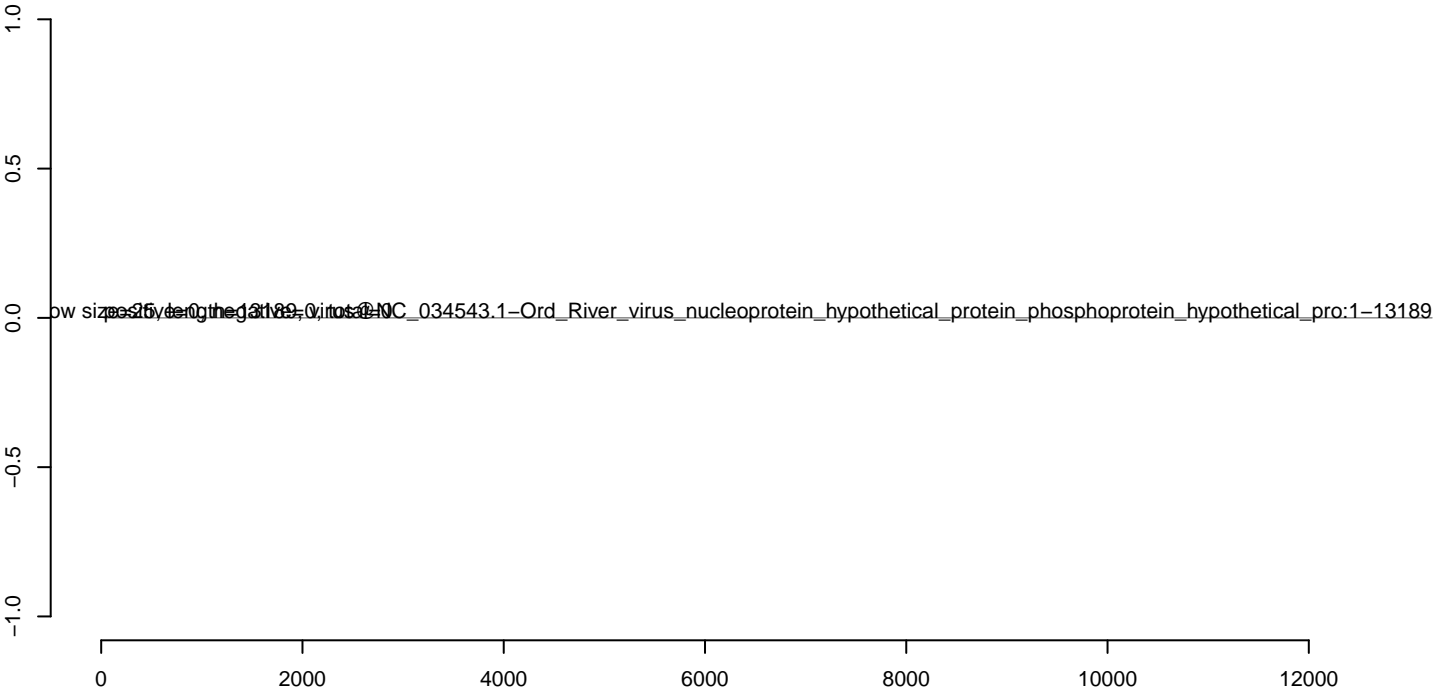
AeAeg_Aag2DENV2NGC.18_23.rep



AeAeg_Aag2DENV2NGC.24_35.rep



AeAeg_Aag2DENV2NGC.rep



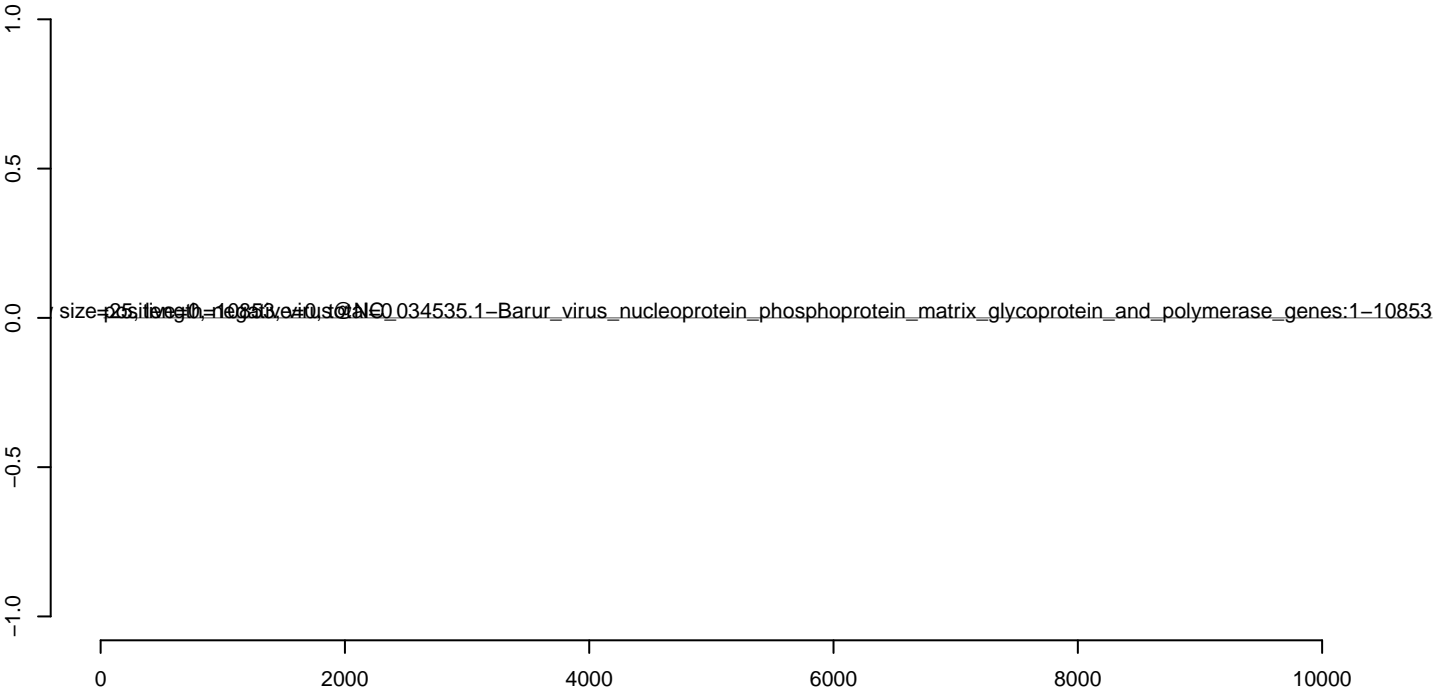
AeAeg_Aag2DENV2NGC.18_23.rep



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AeAeg_Aag2DENV2NGC.rep



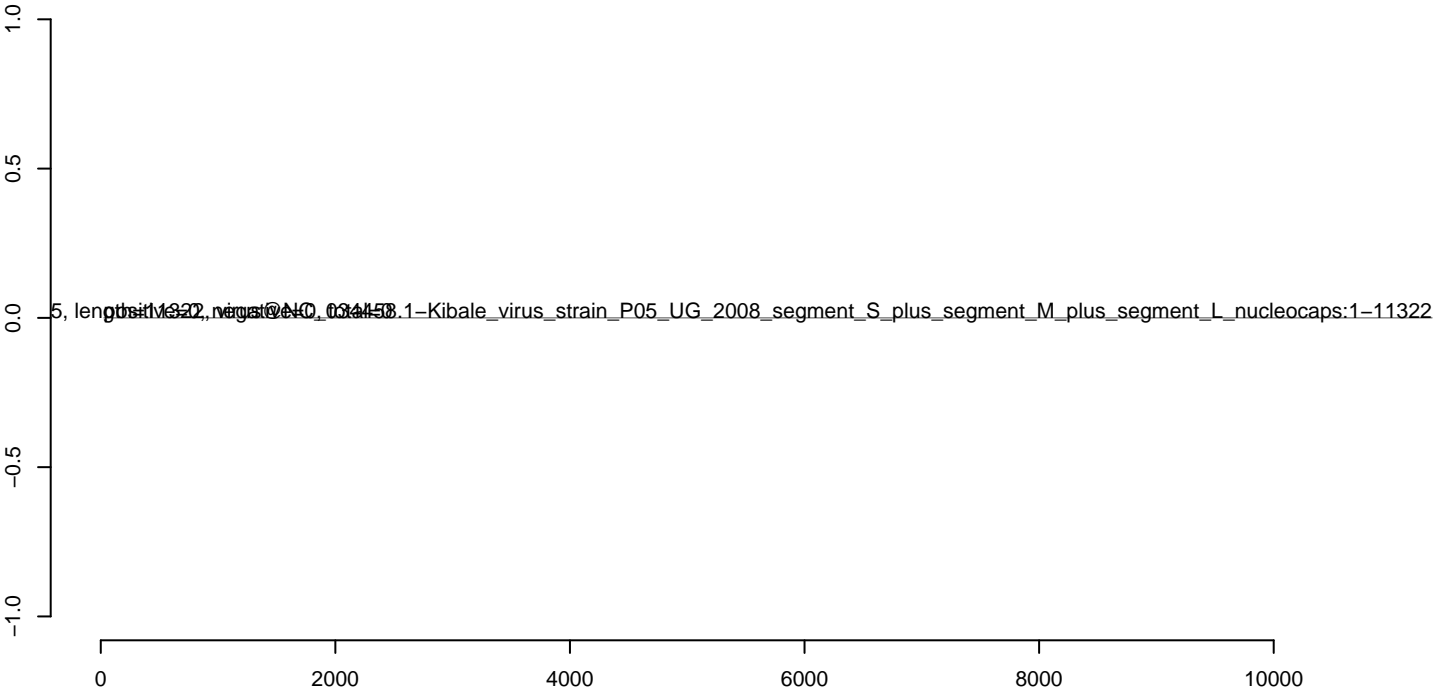
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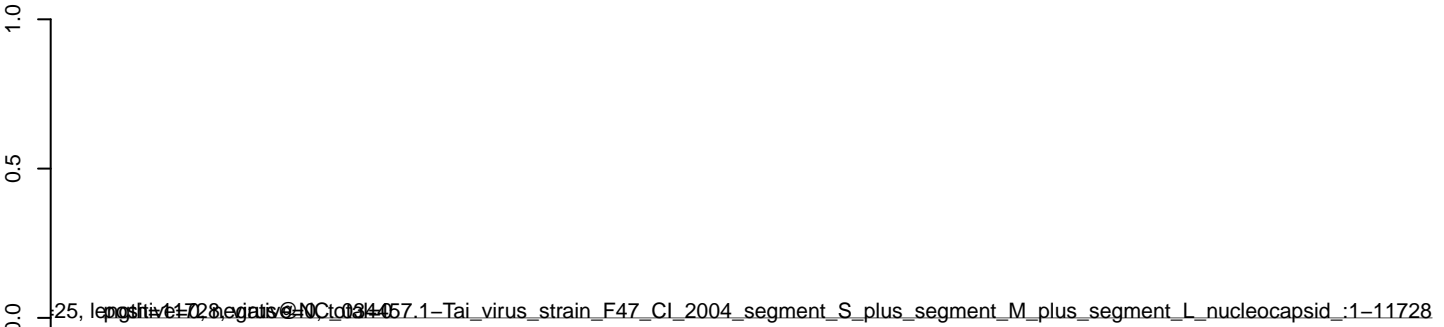
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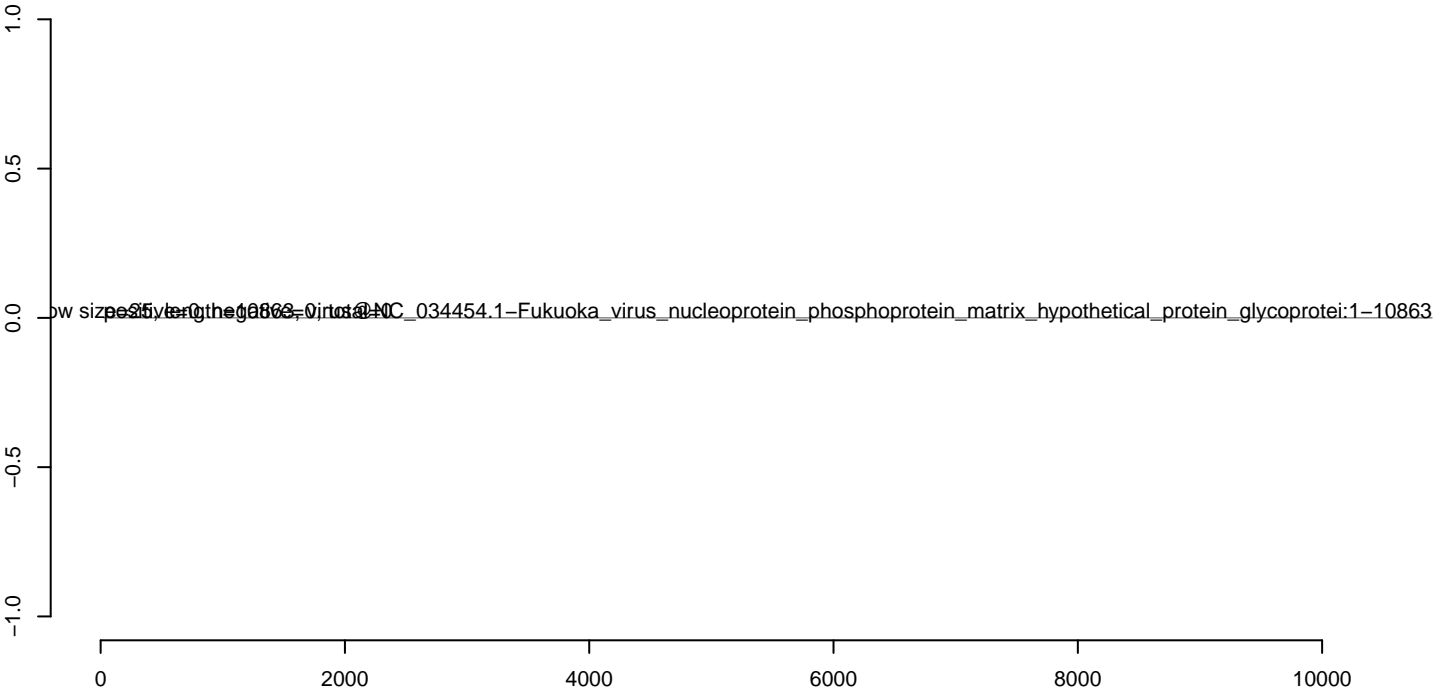
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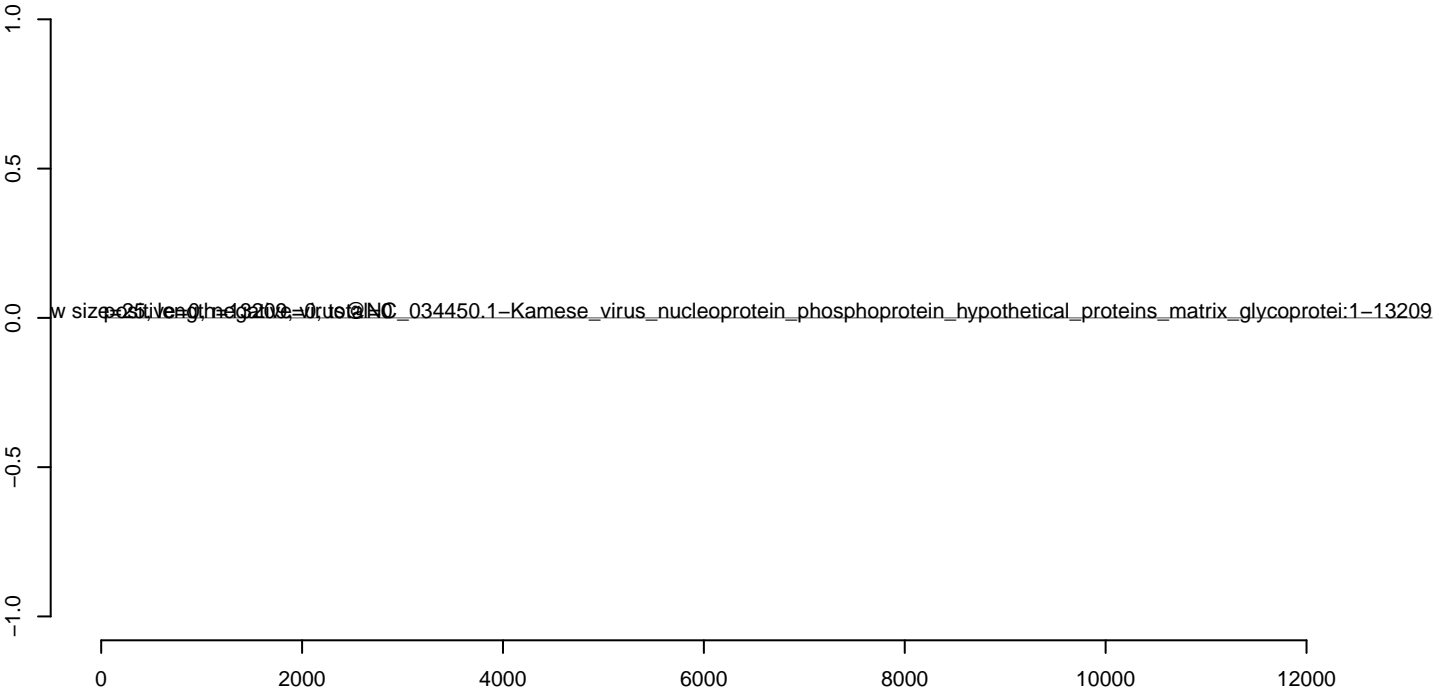
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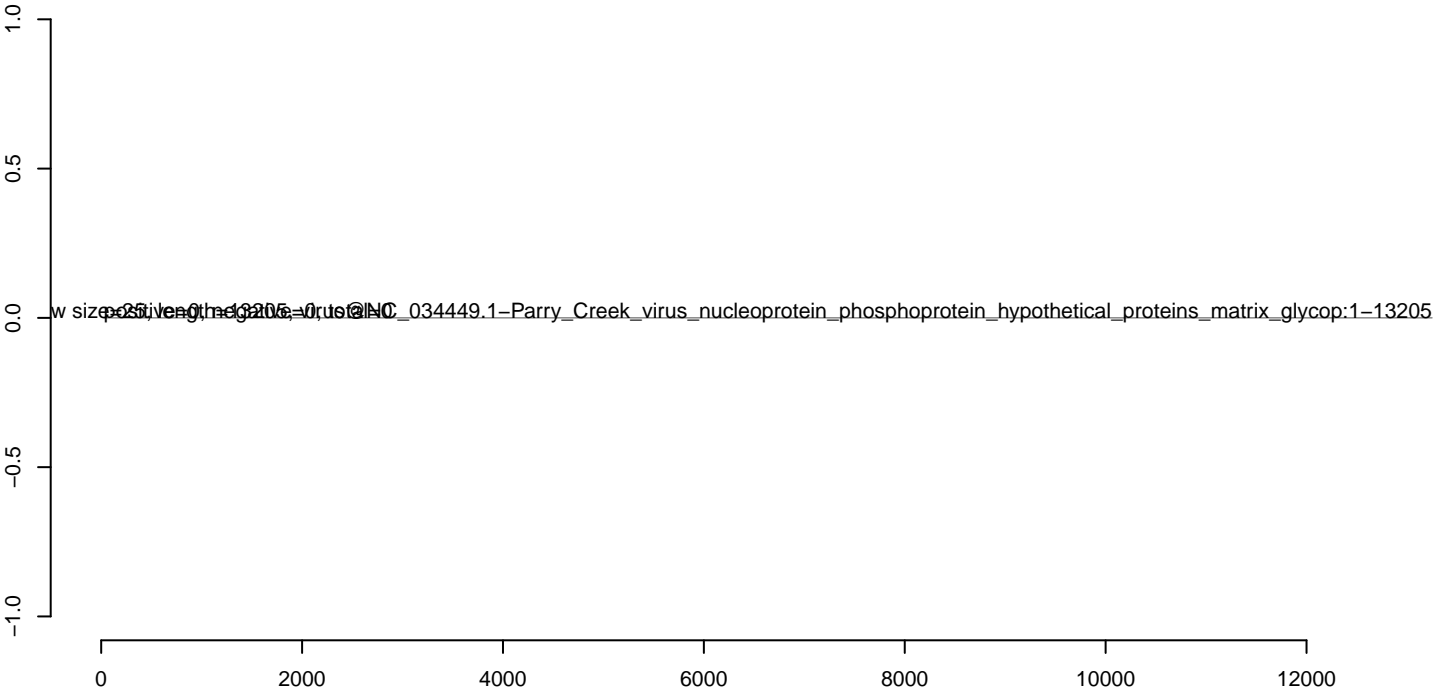
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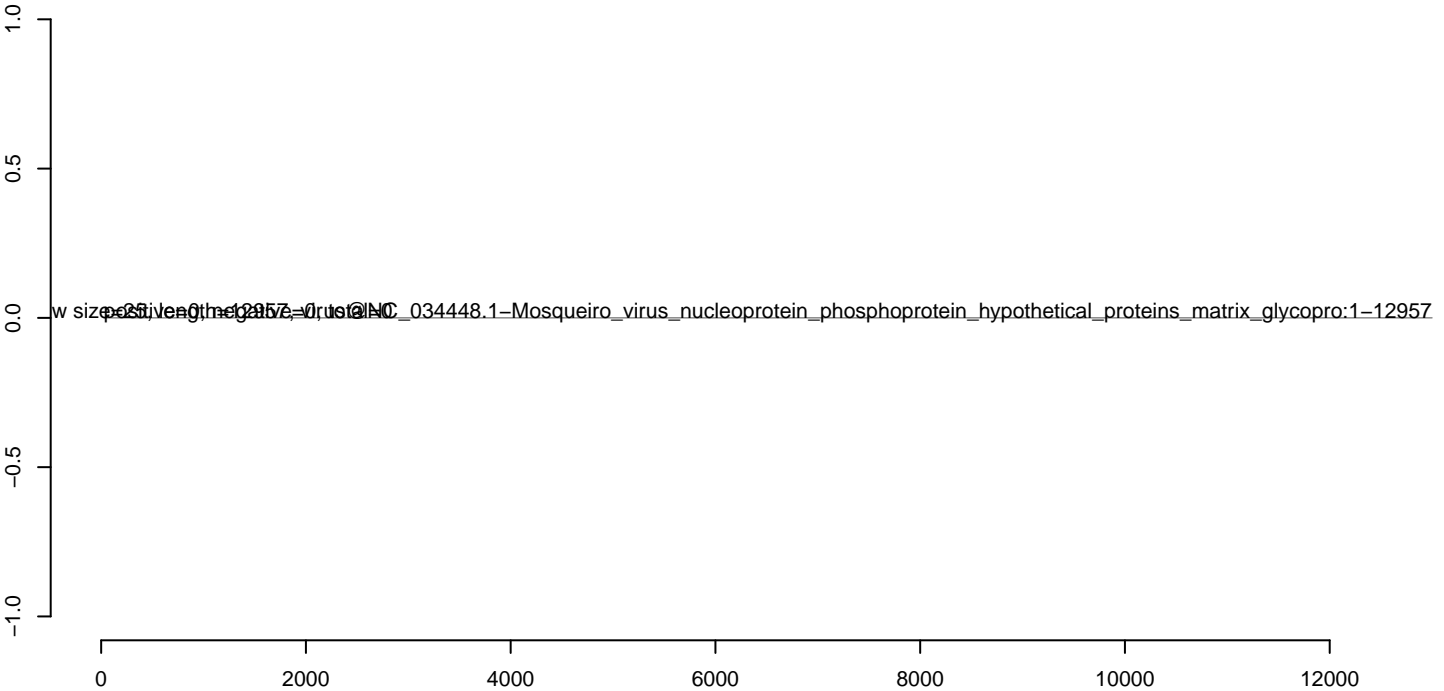
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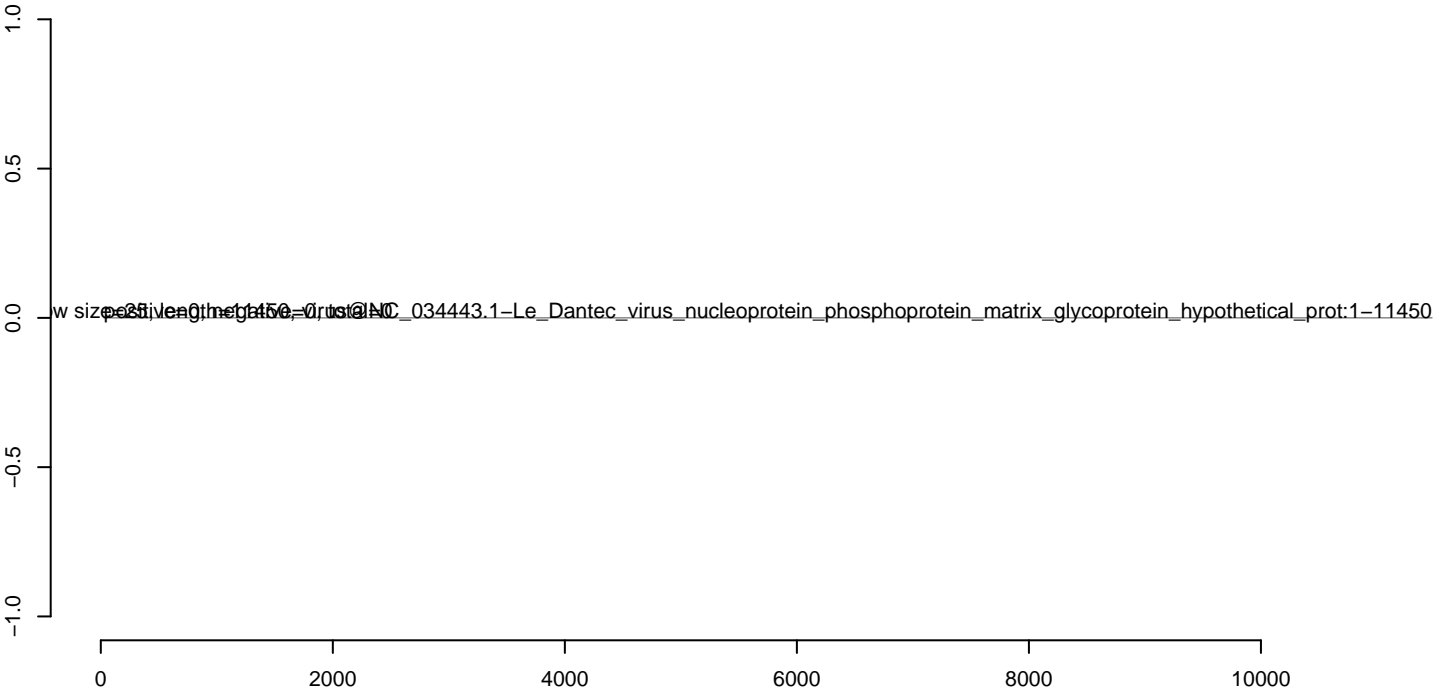
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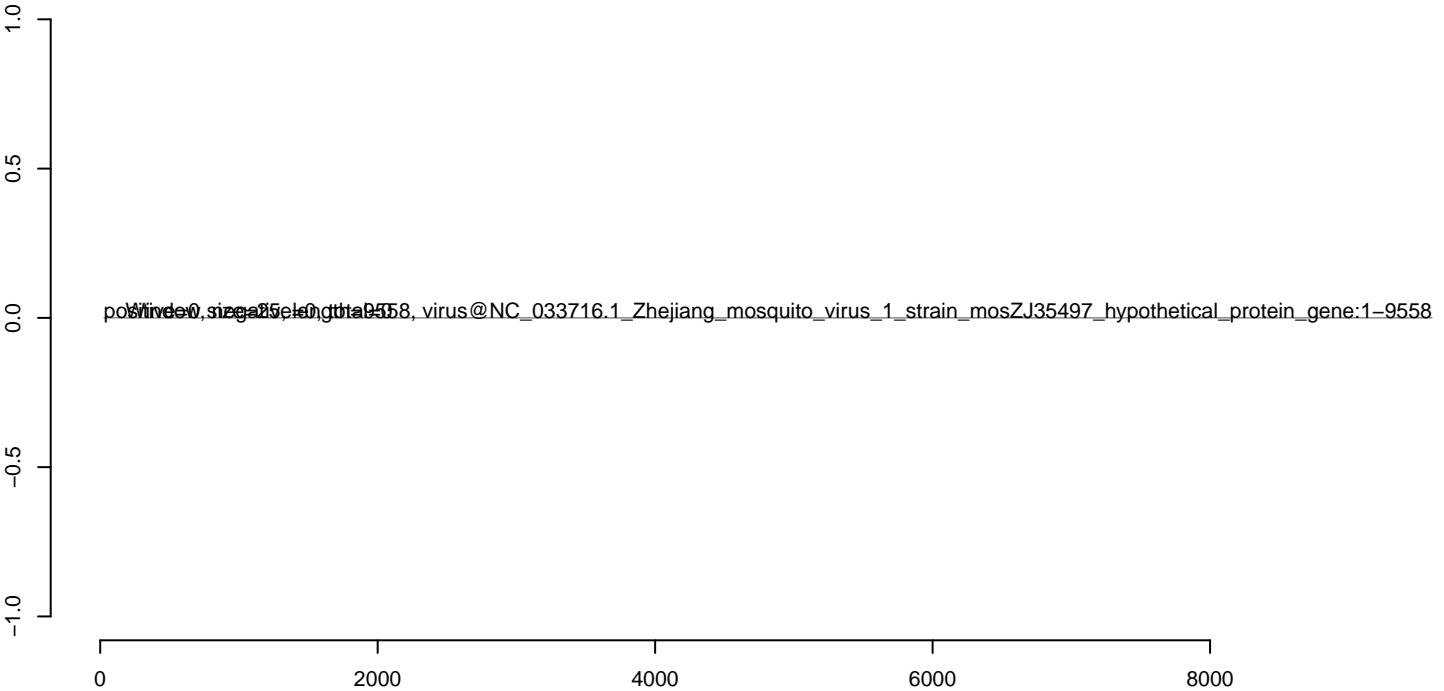
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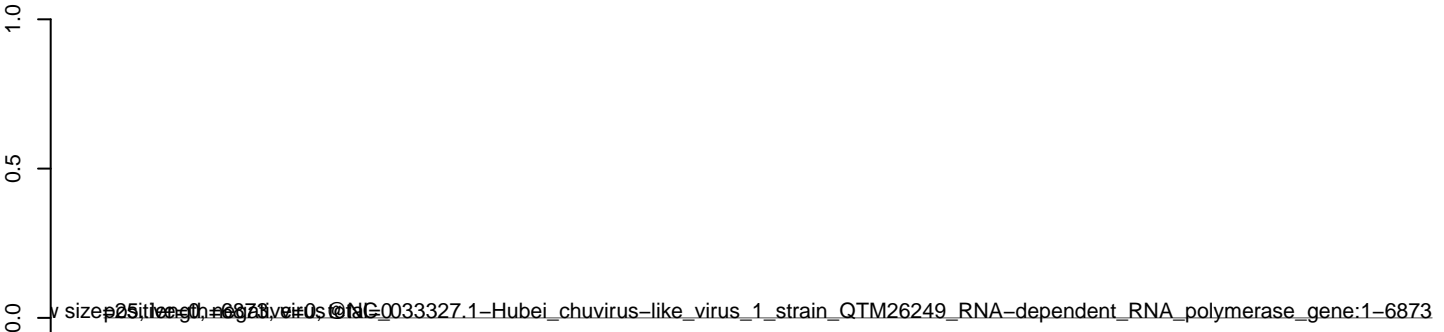
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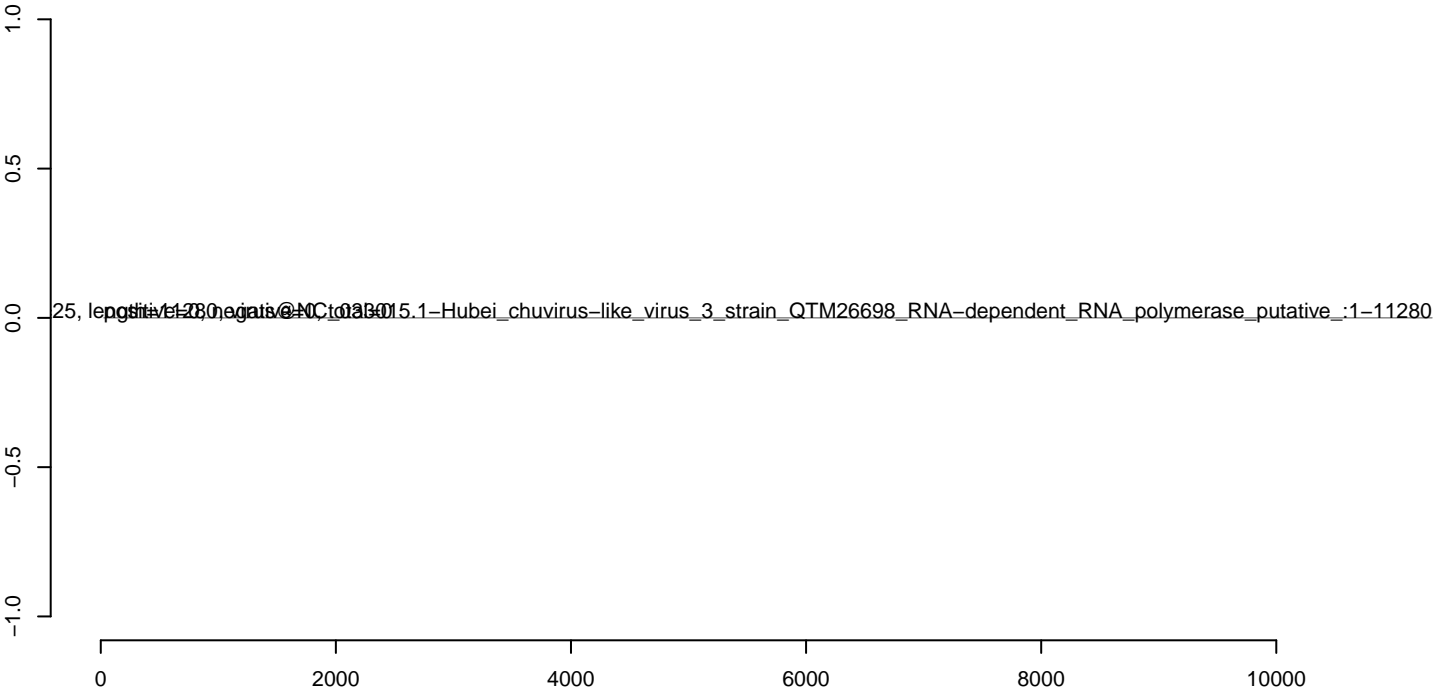
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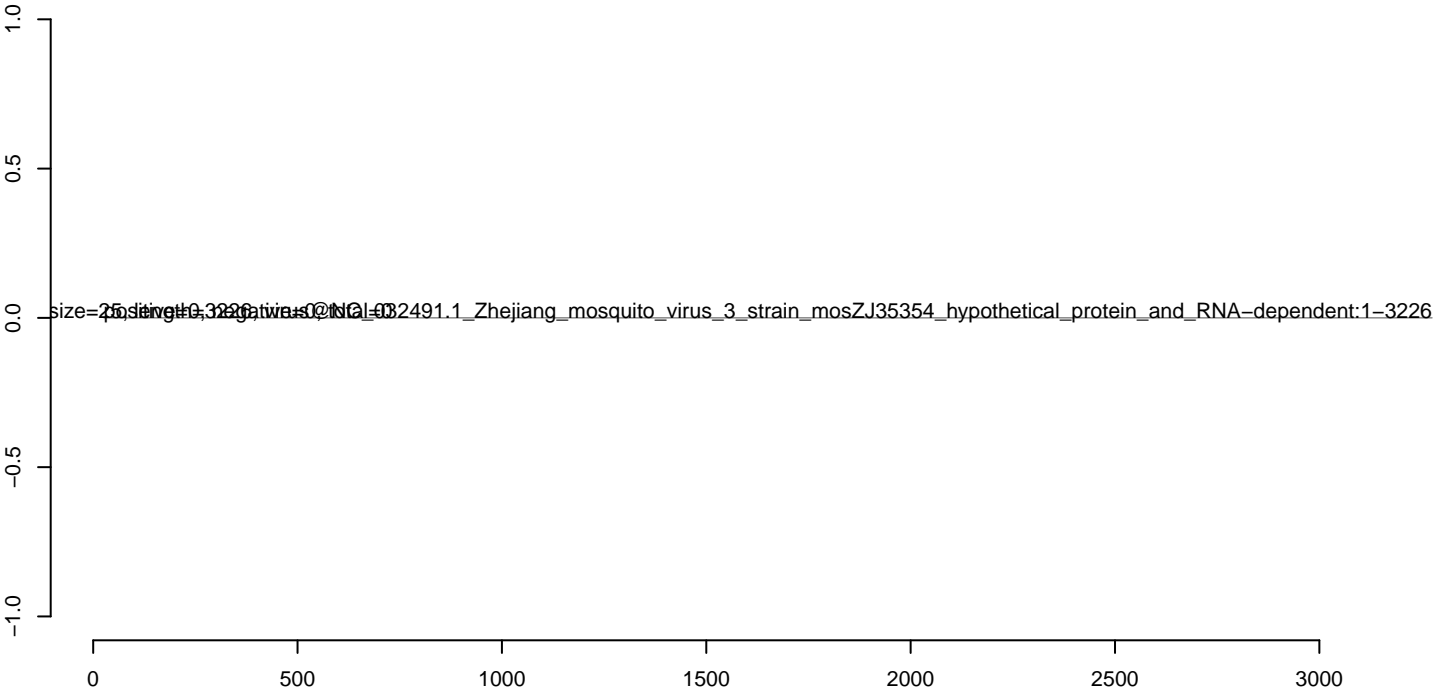
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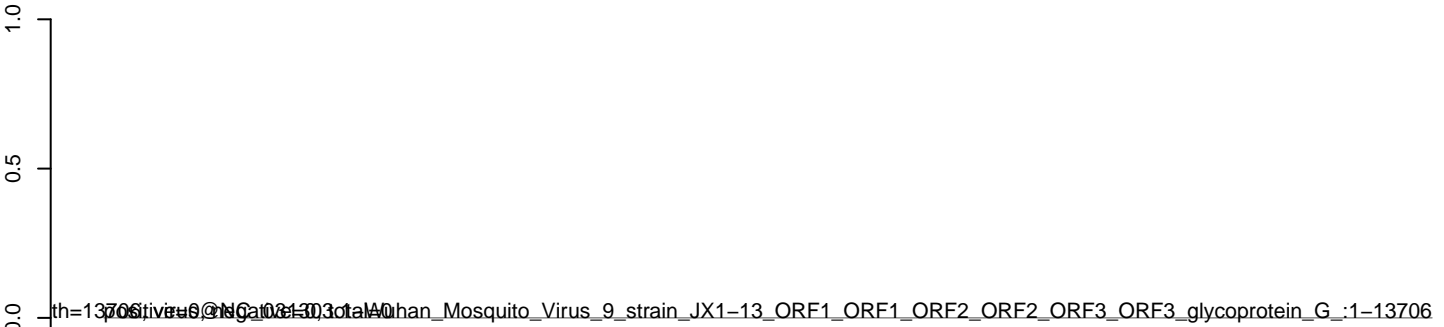
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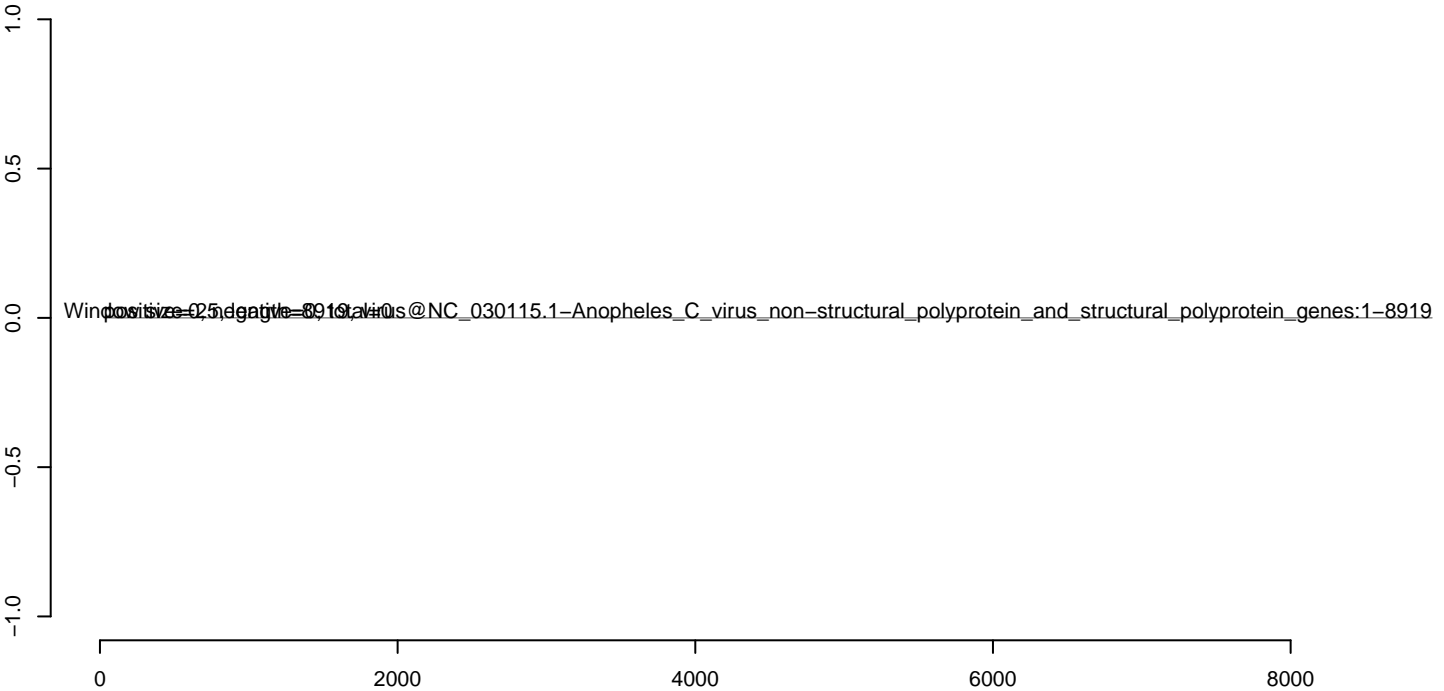
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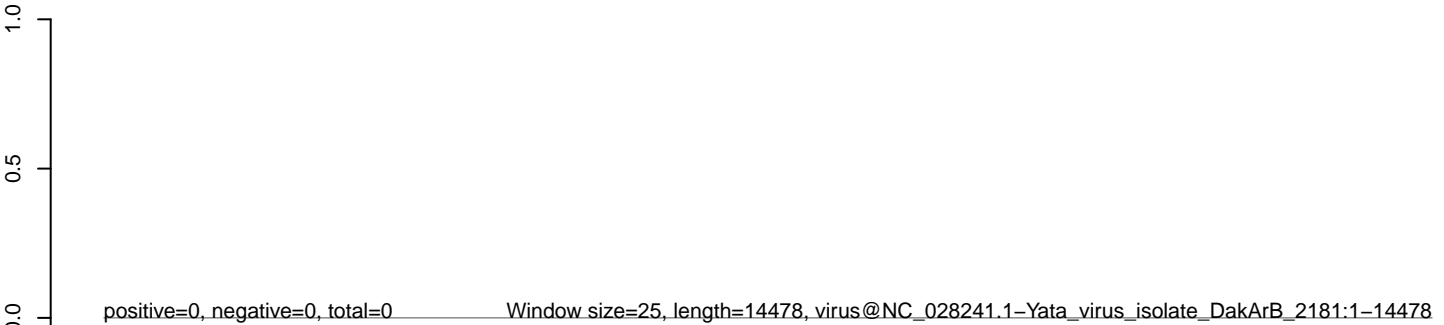
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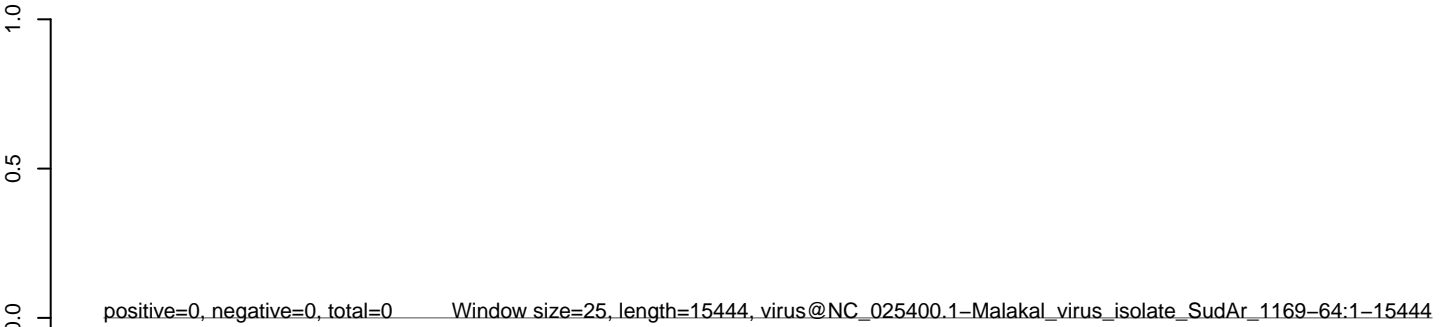
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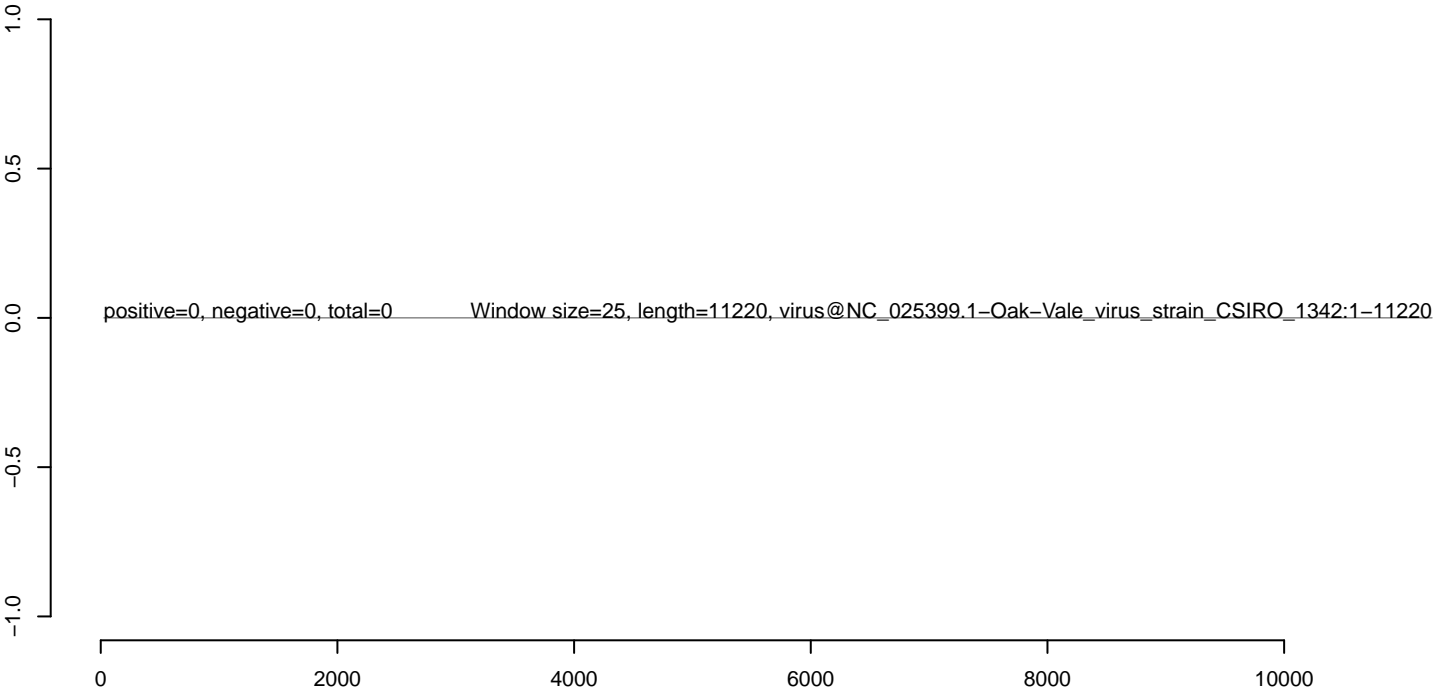
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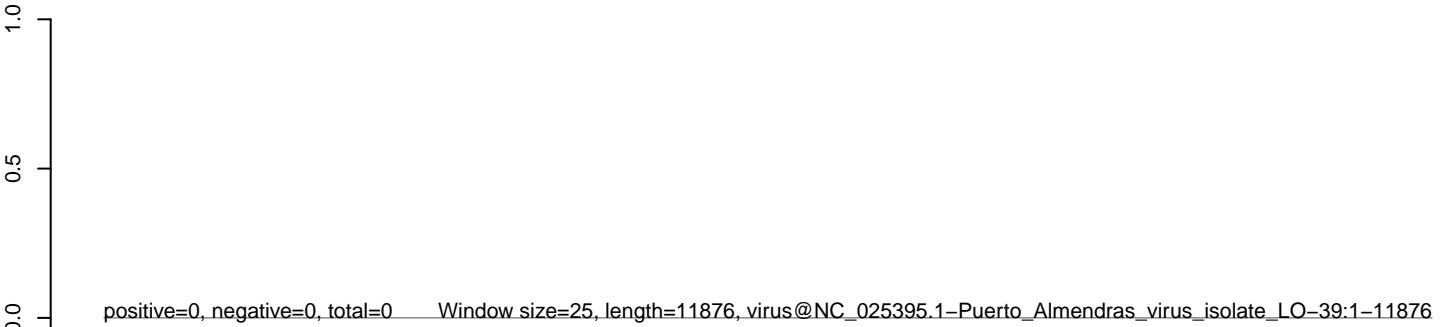
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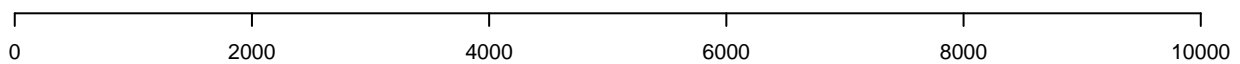
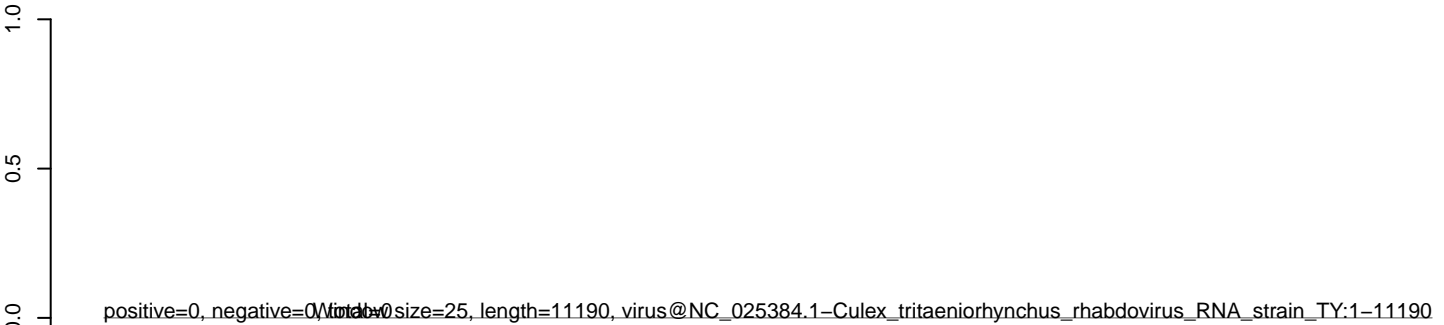
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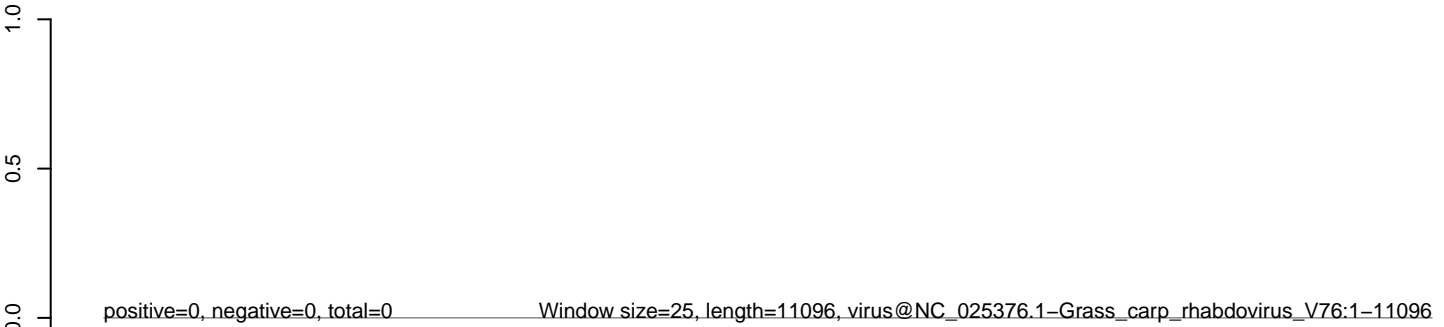
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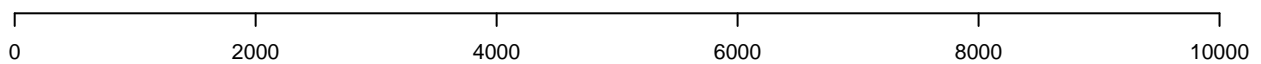
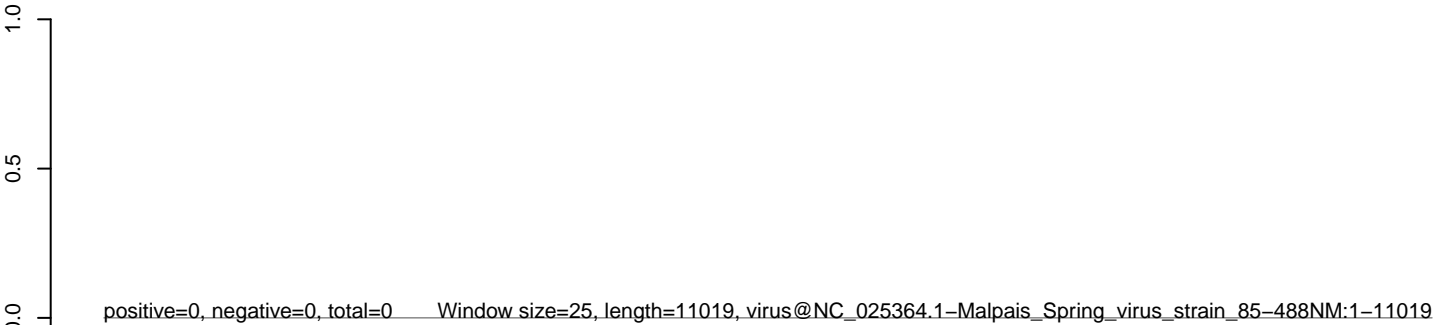
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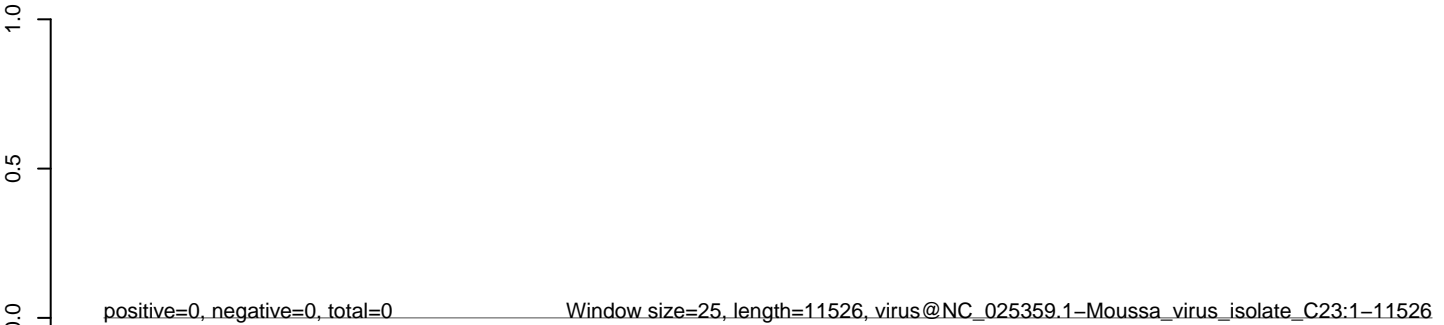
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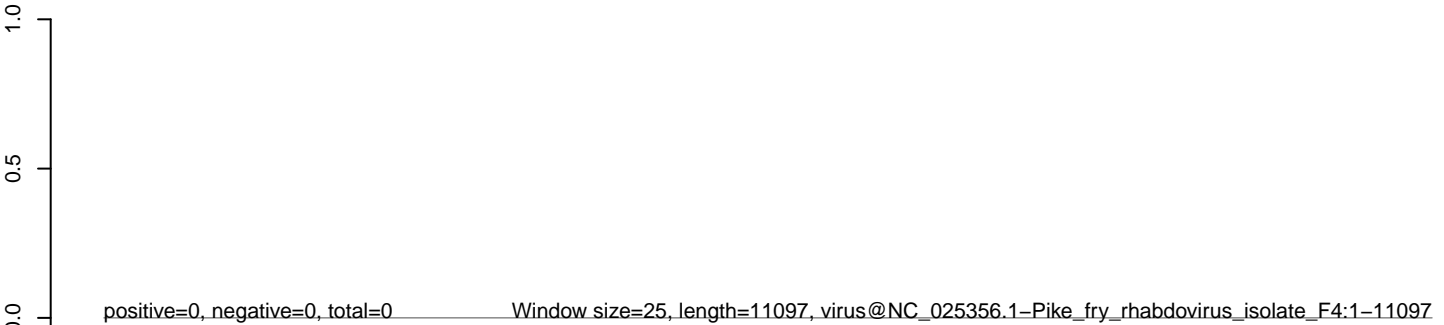
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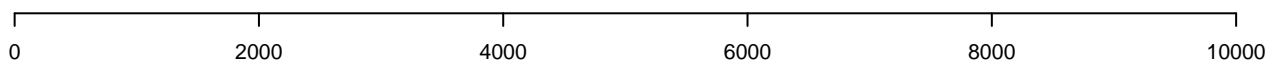
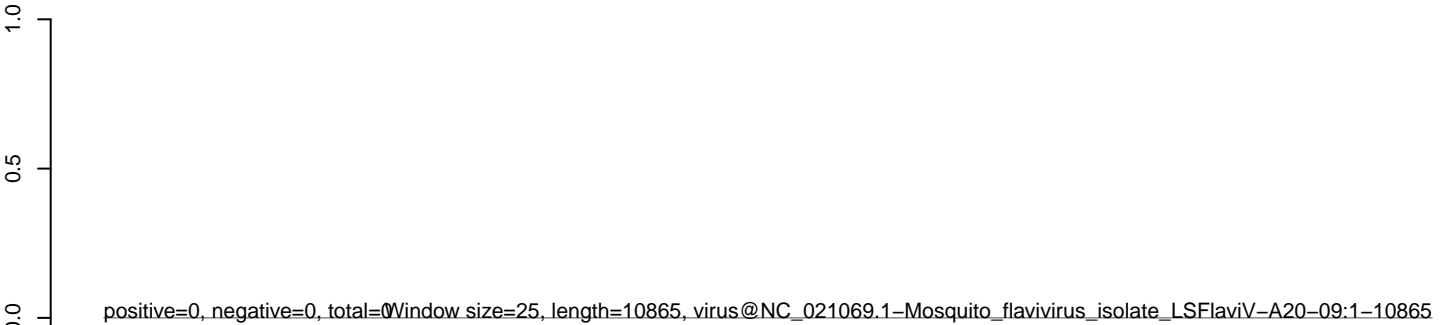
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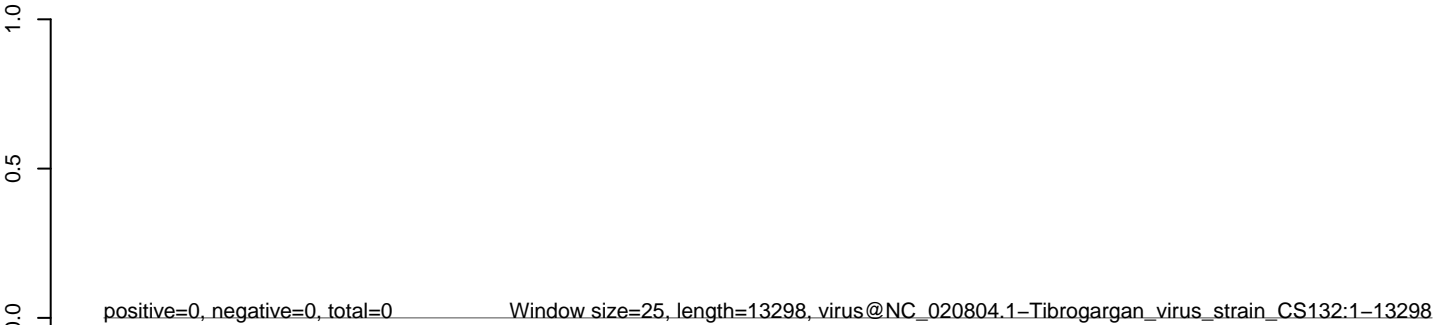
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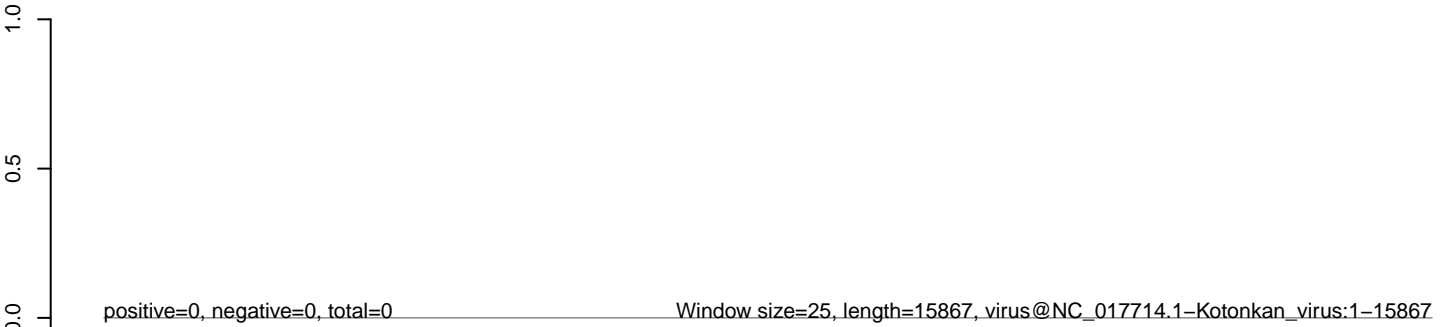
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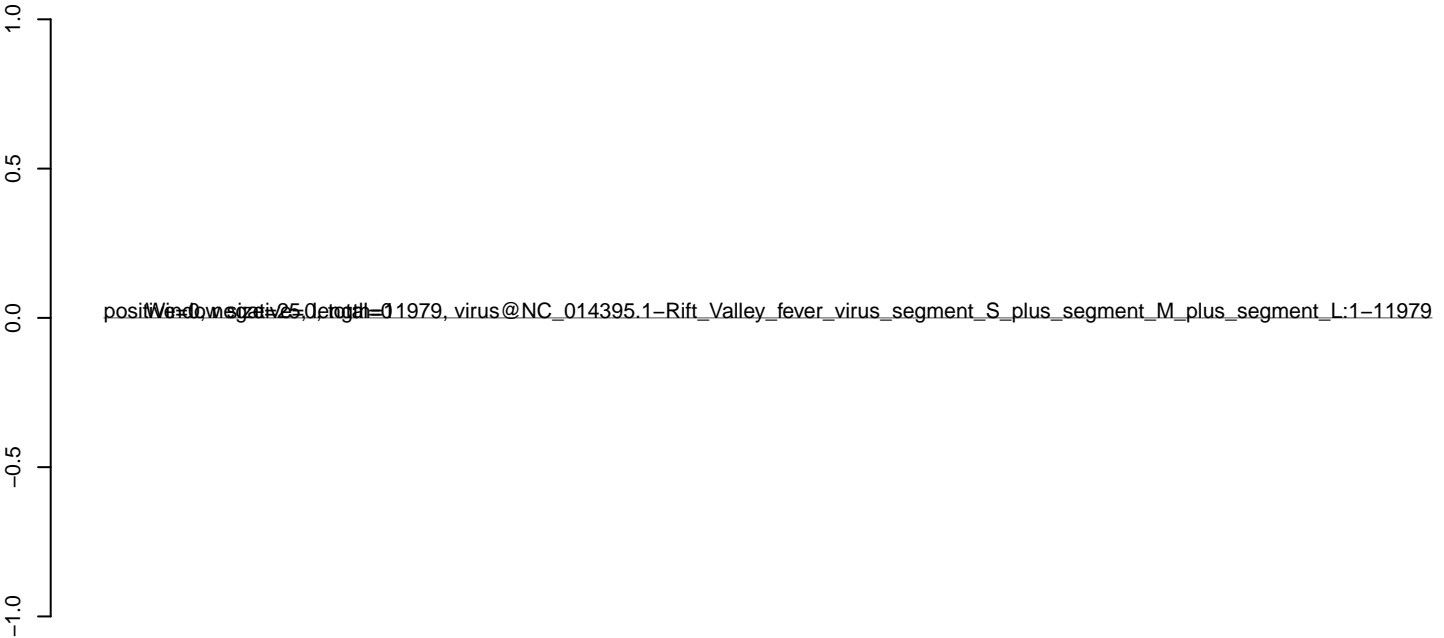
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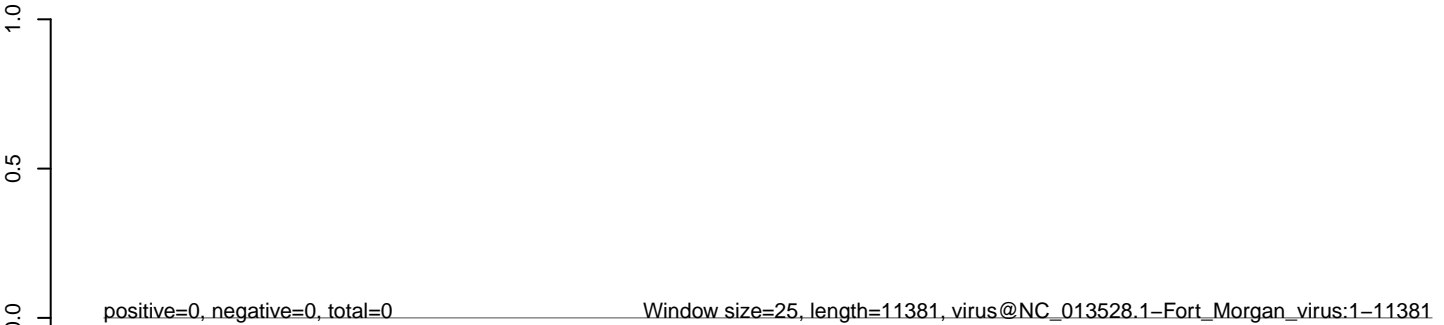
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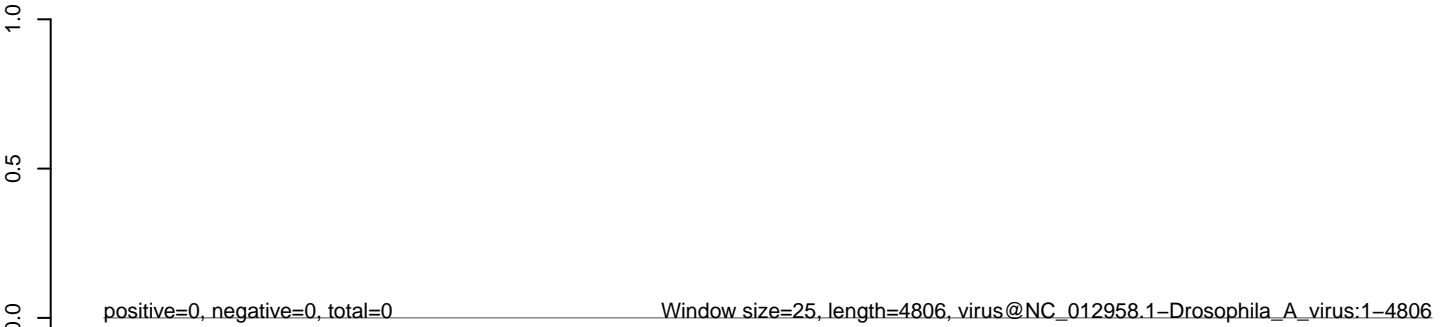
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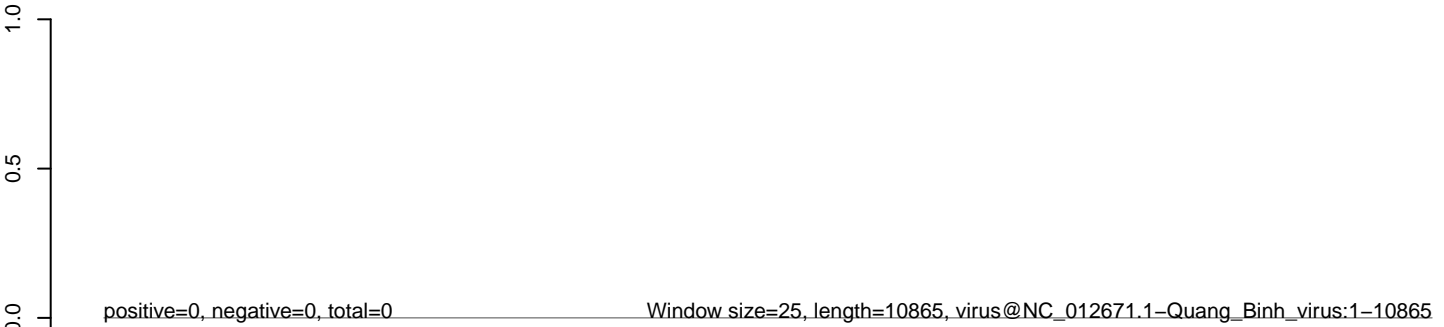
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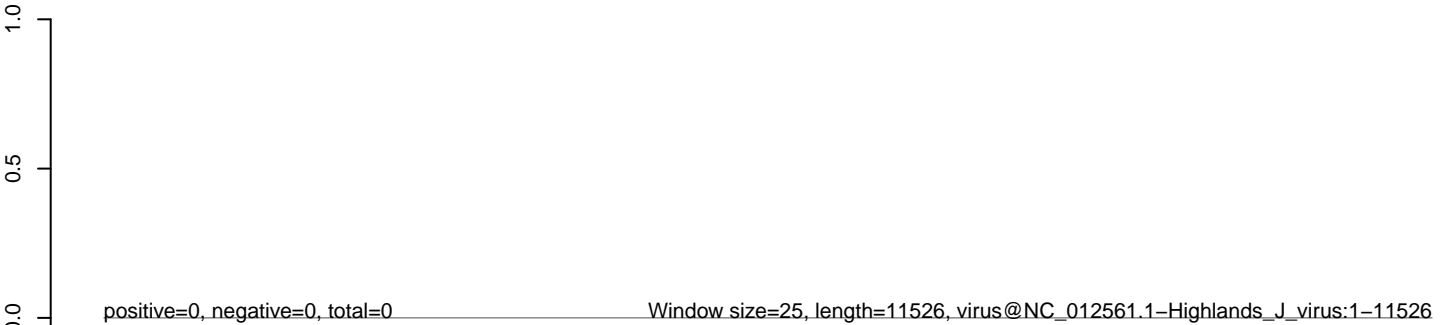
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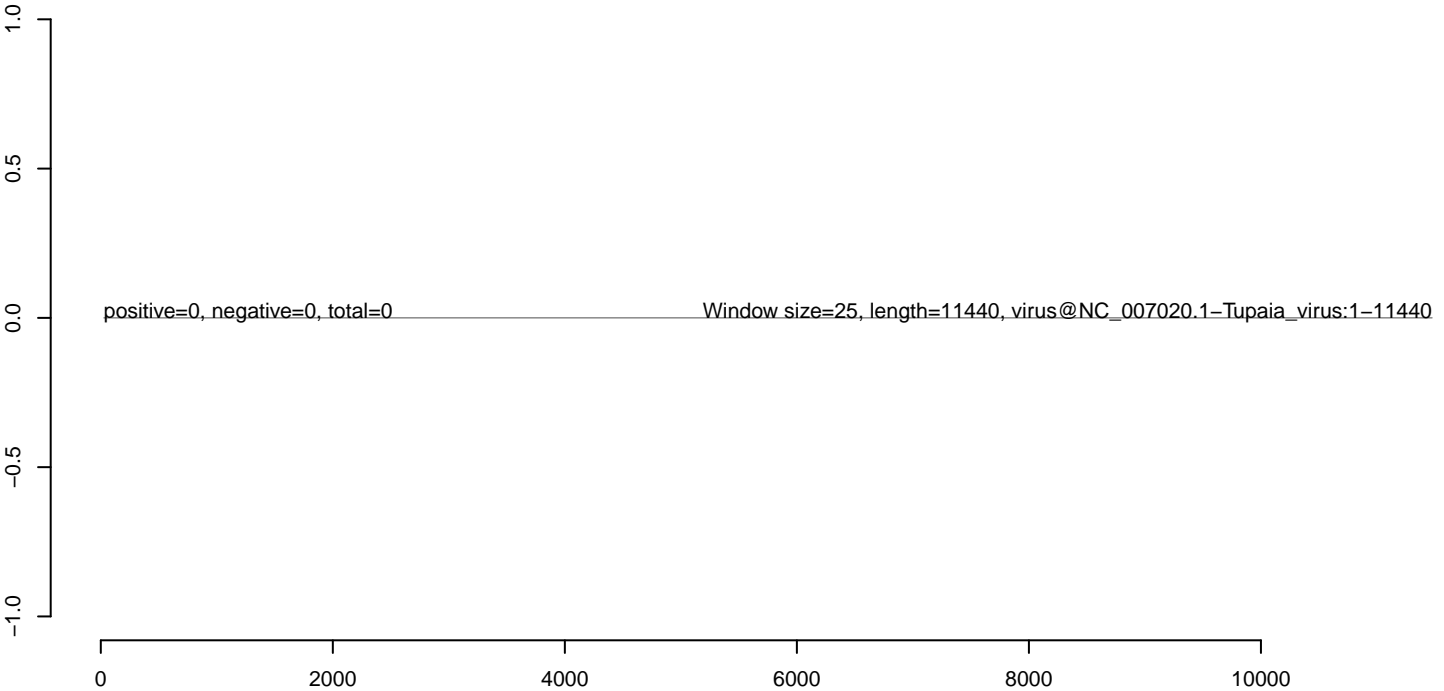
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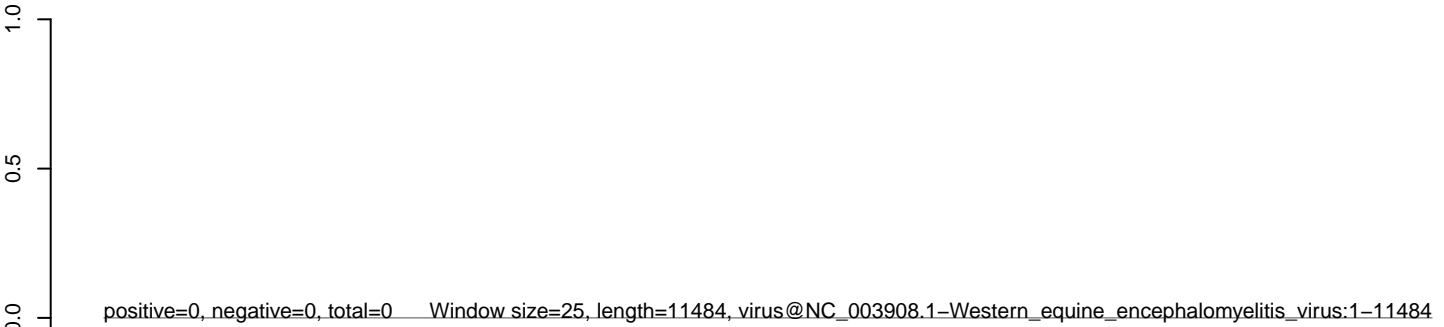
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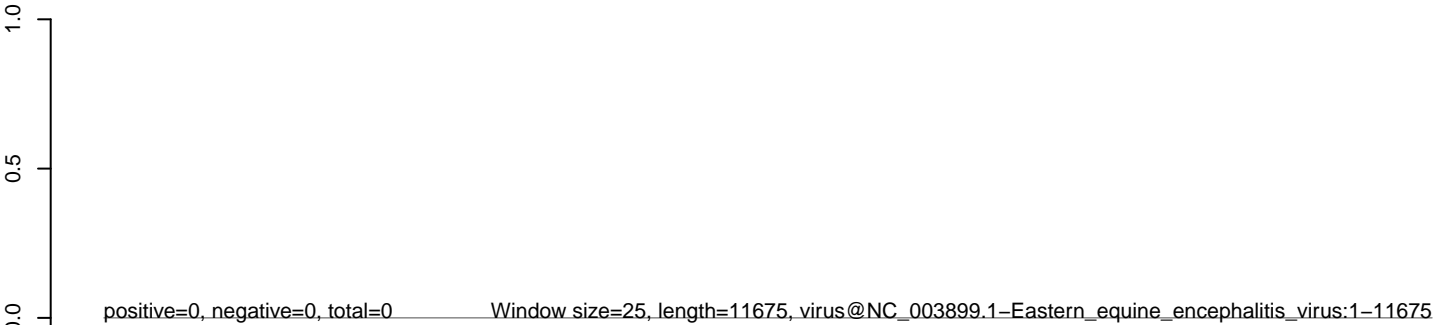
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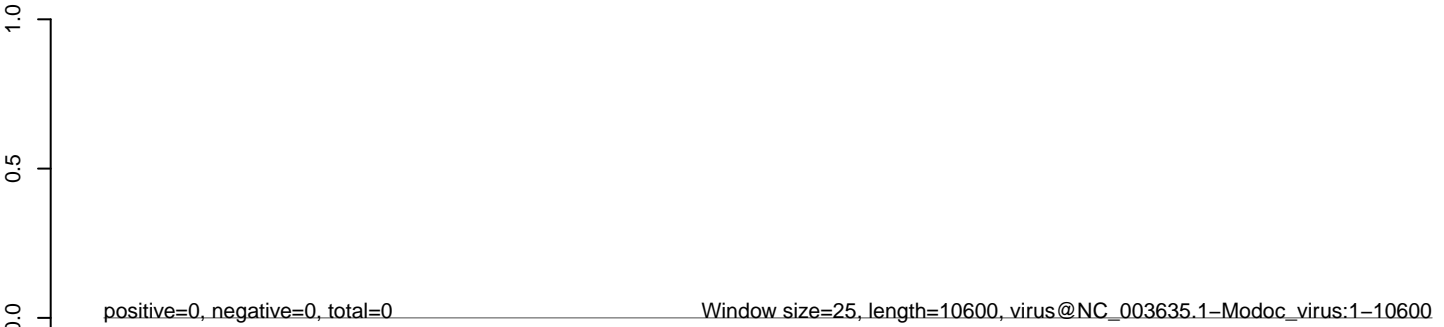
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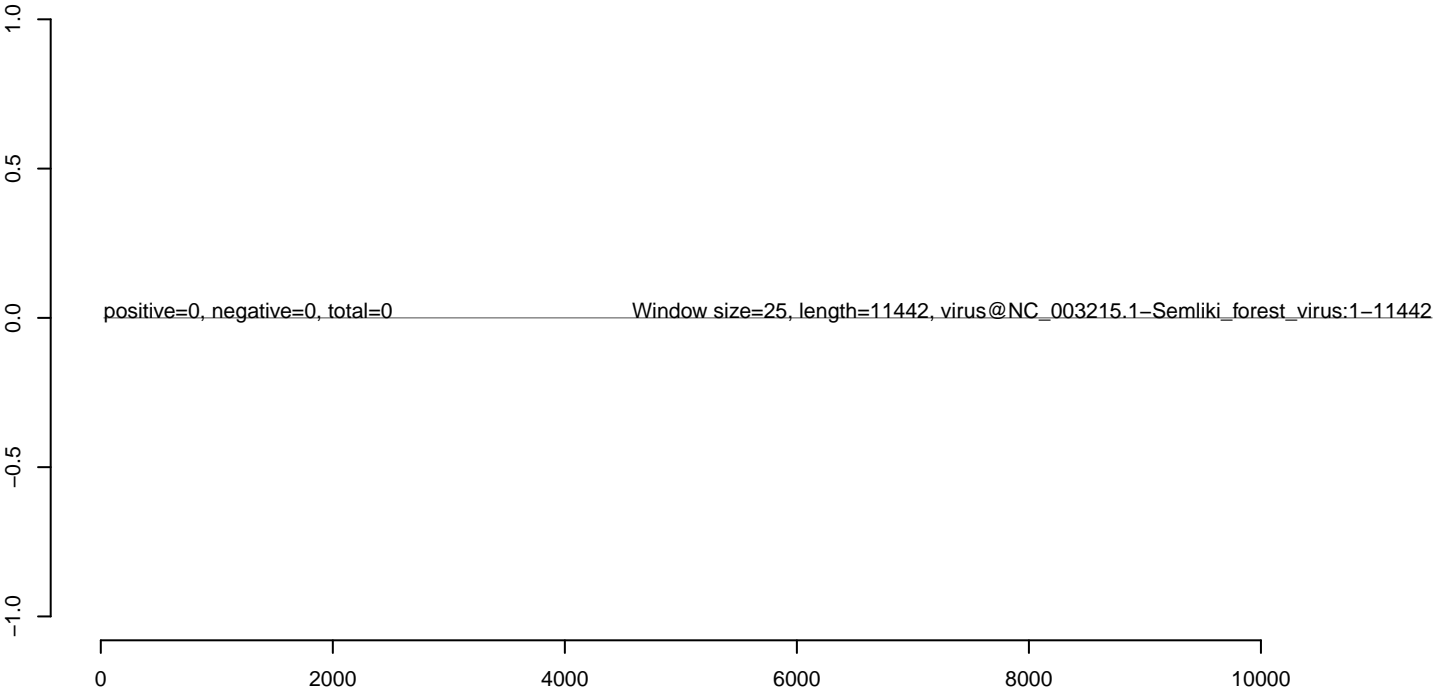
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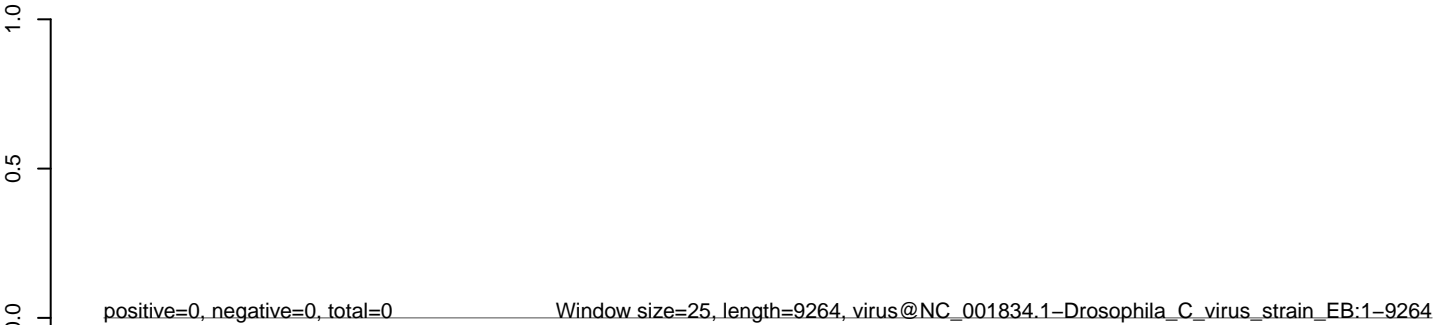
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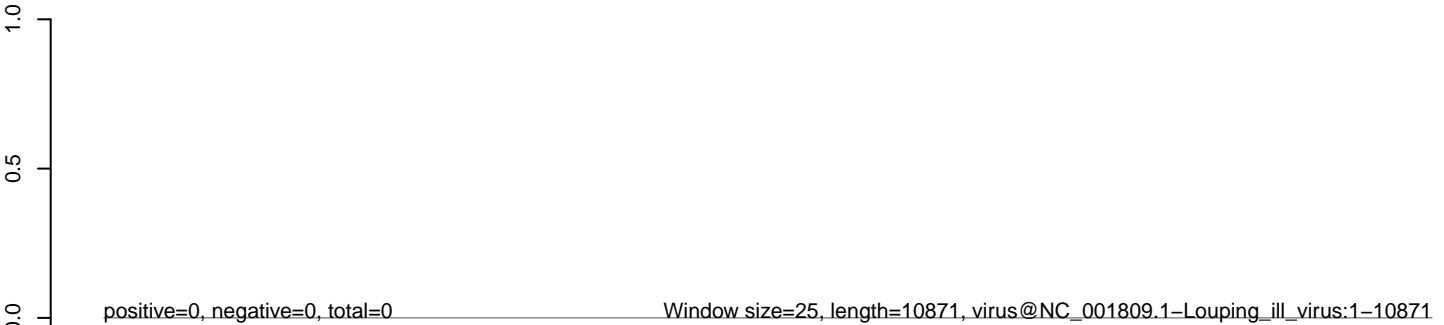
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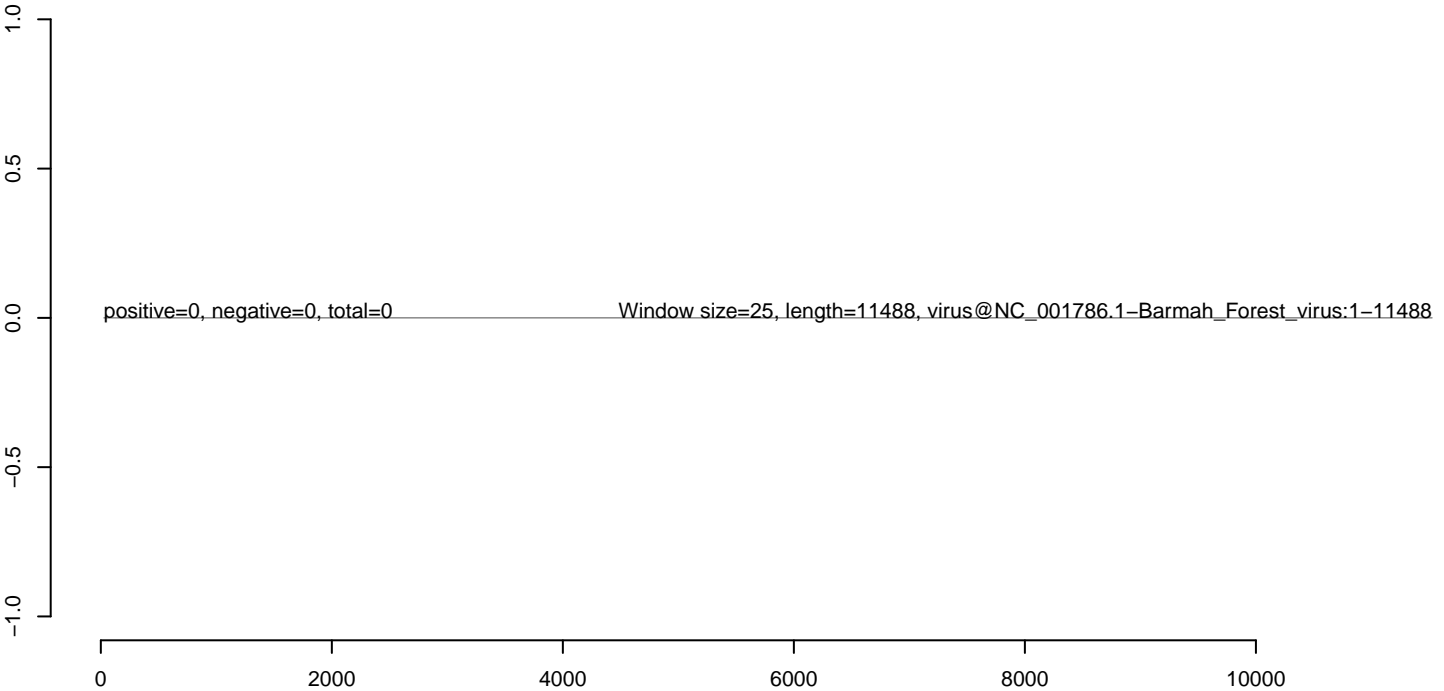
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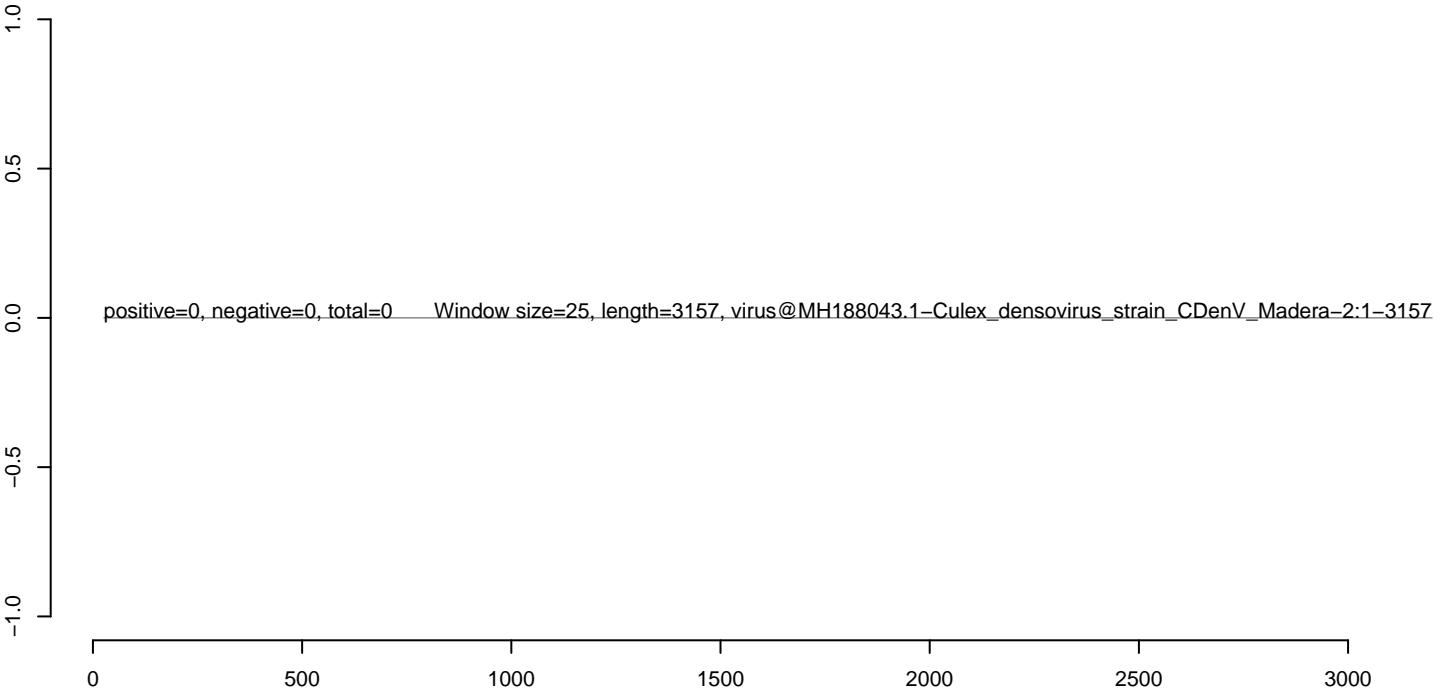
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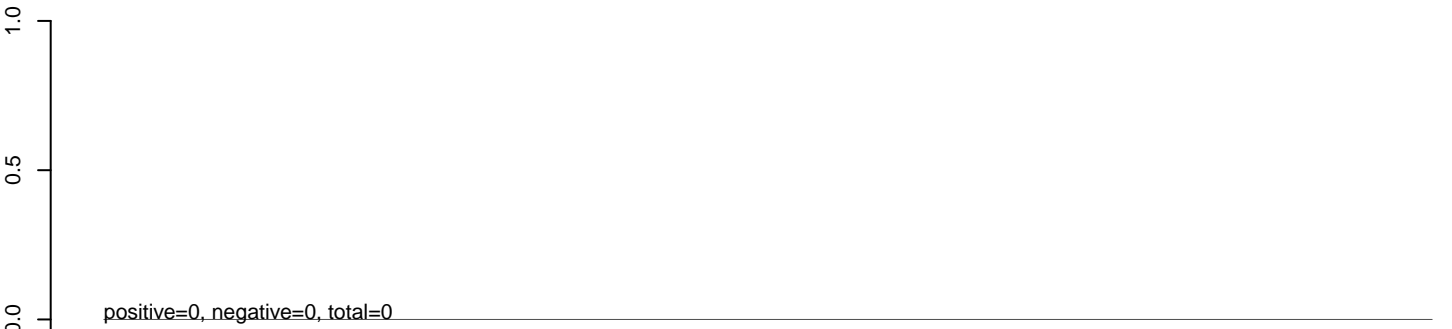
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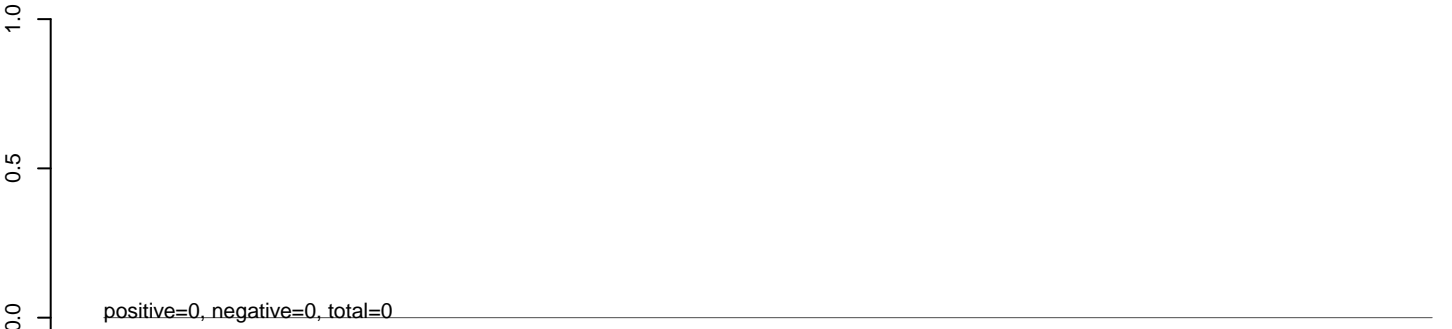
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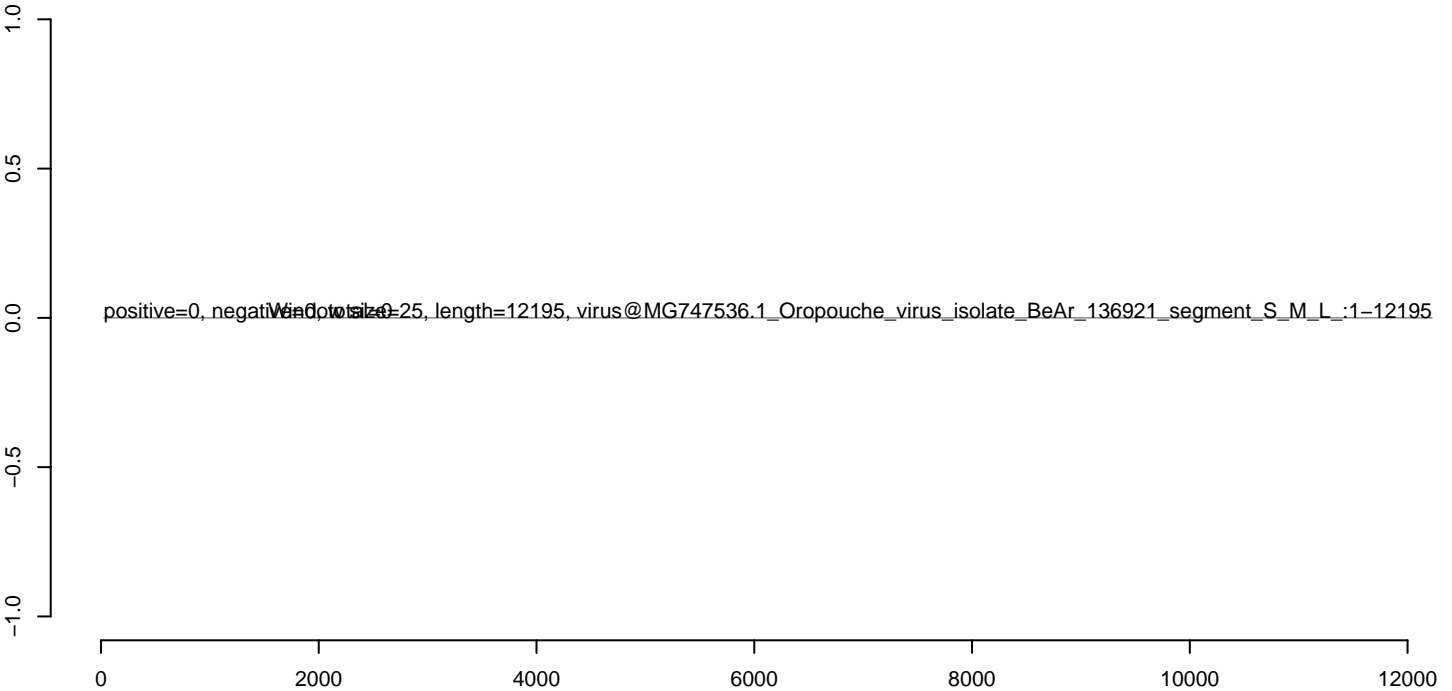
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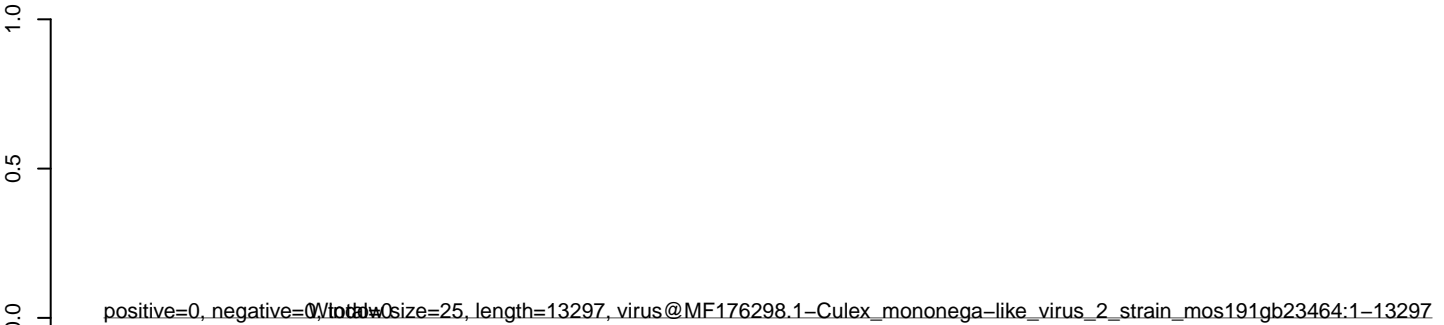
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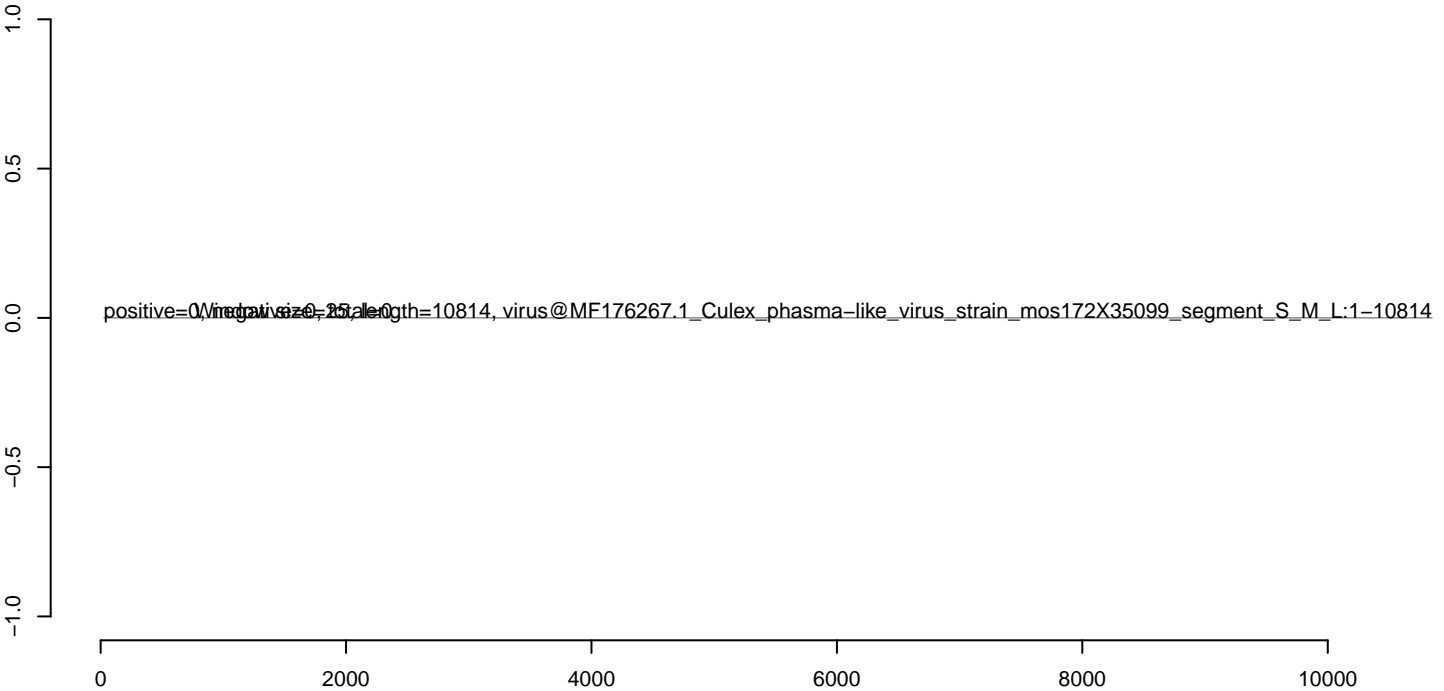
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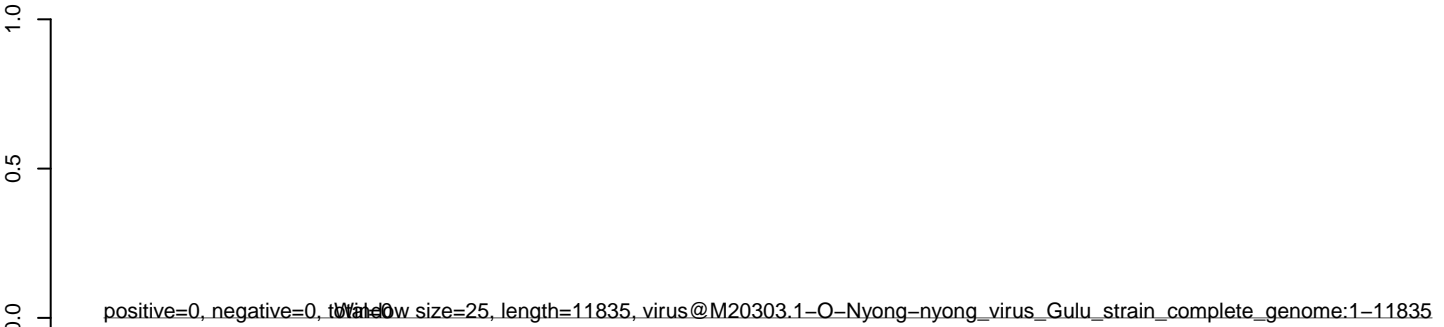
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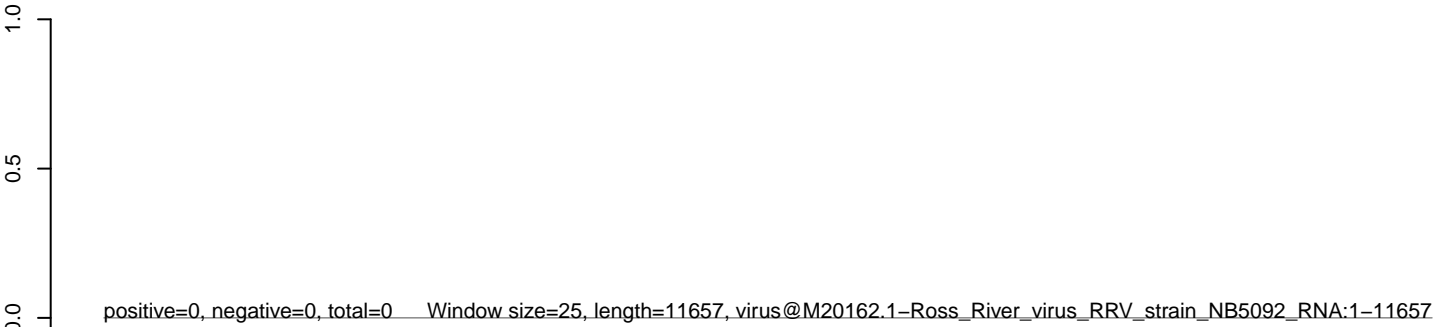
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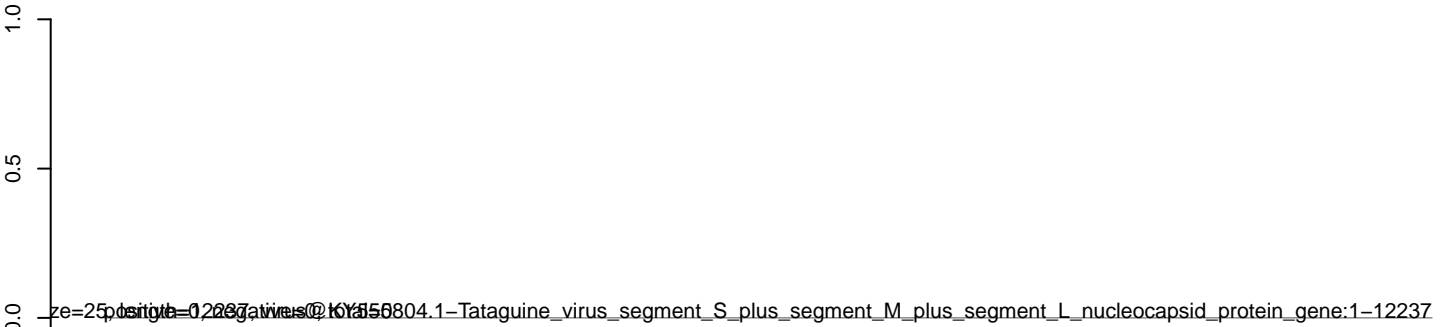
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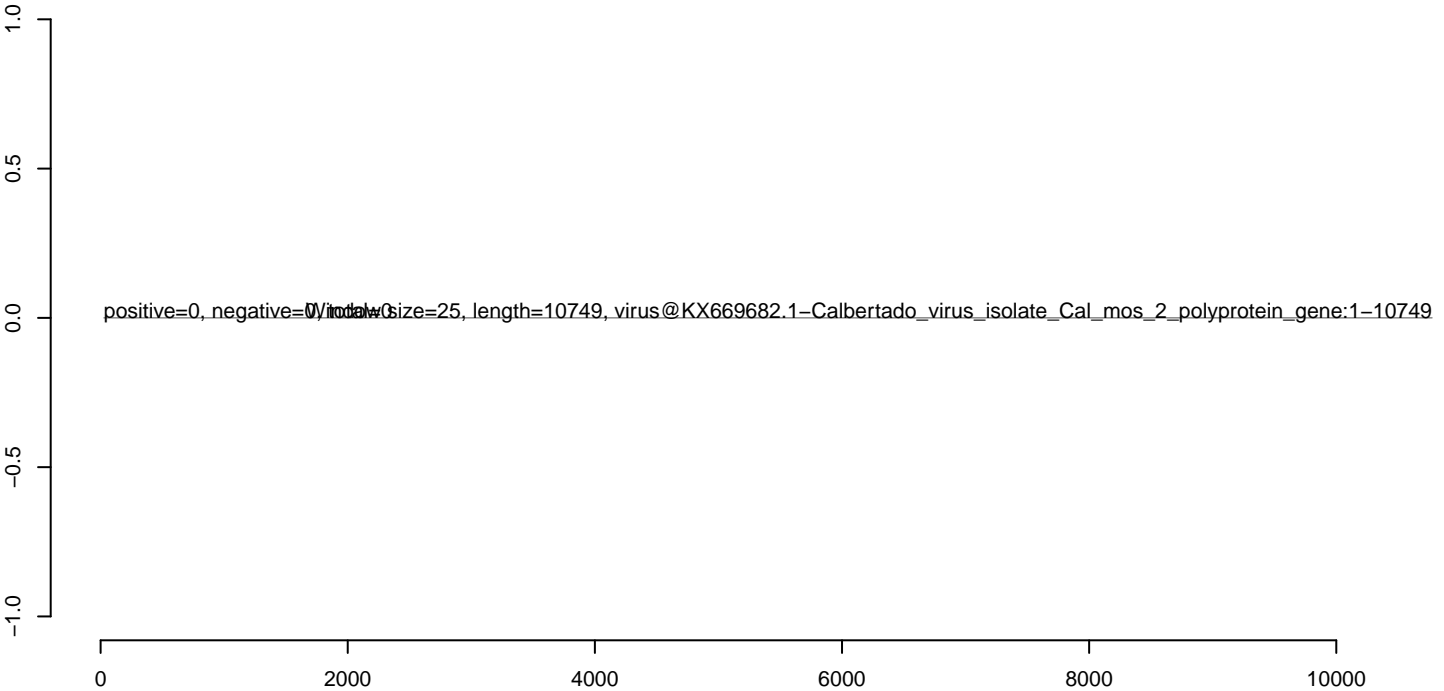
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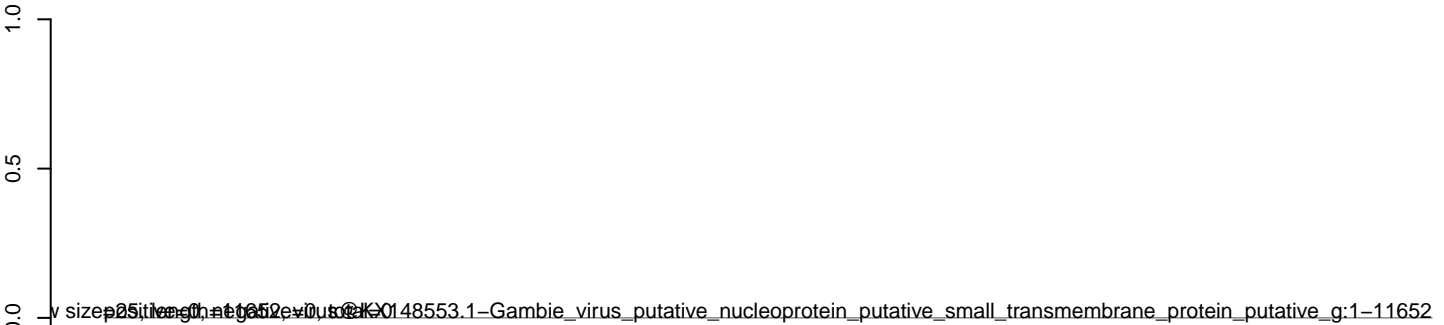
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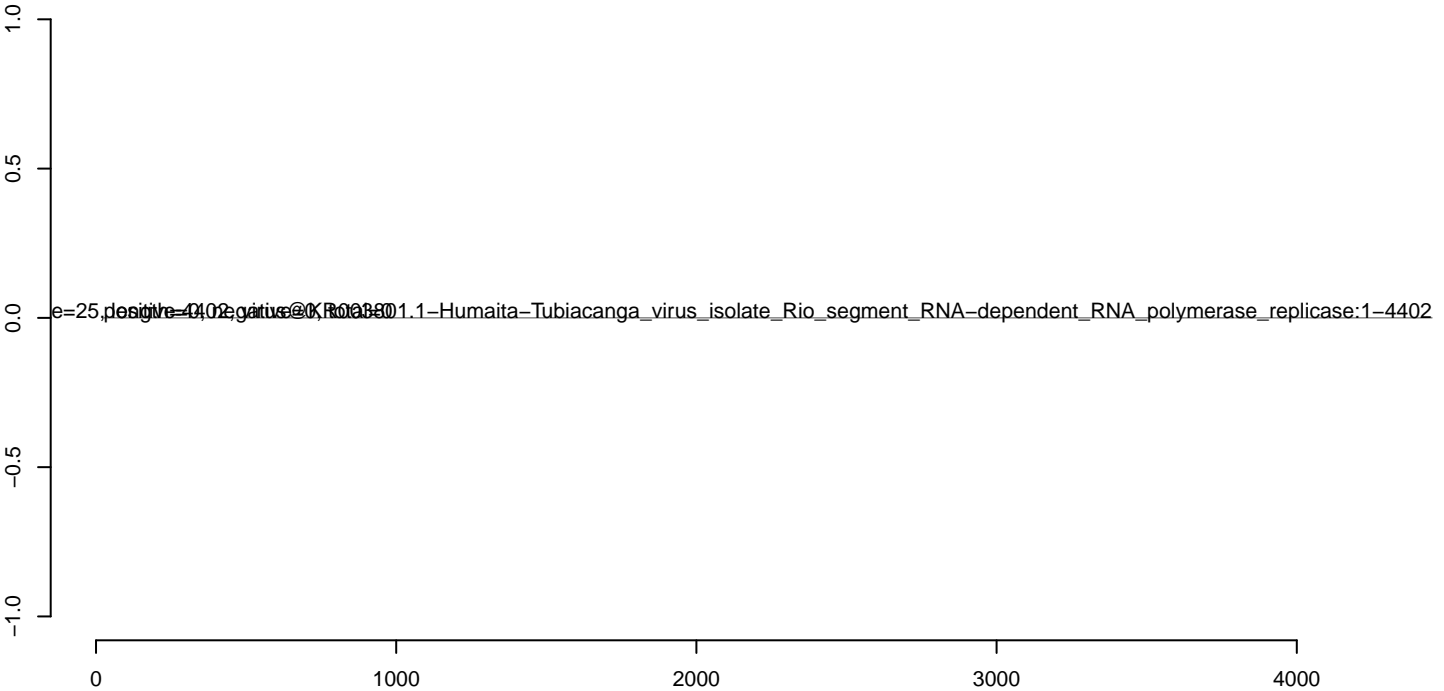
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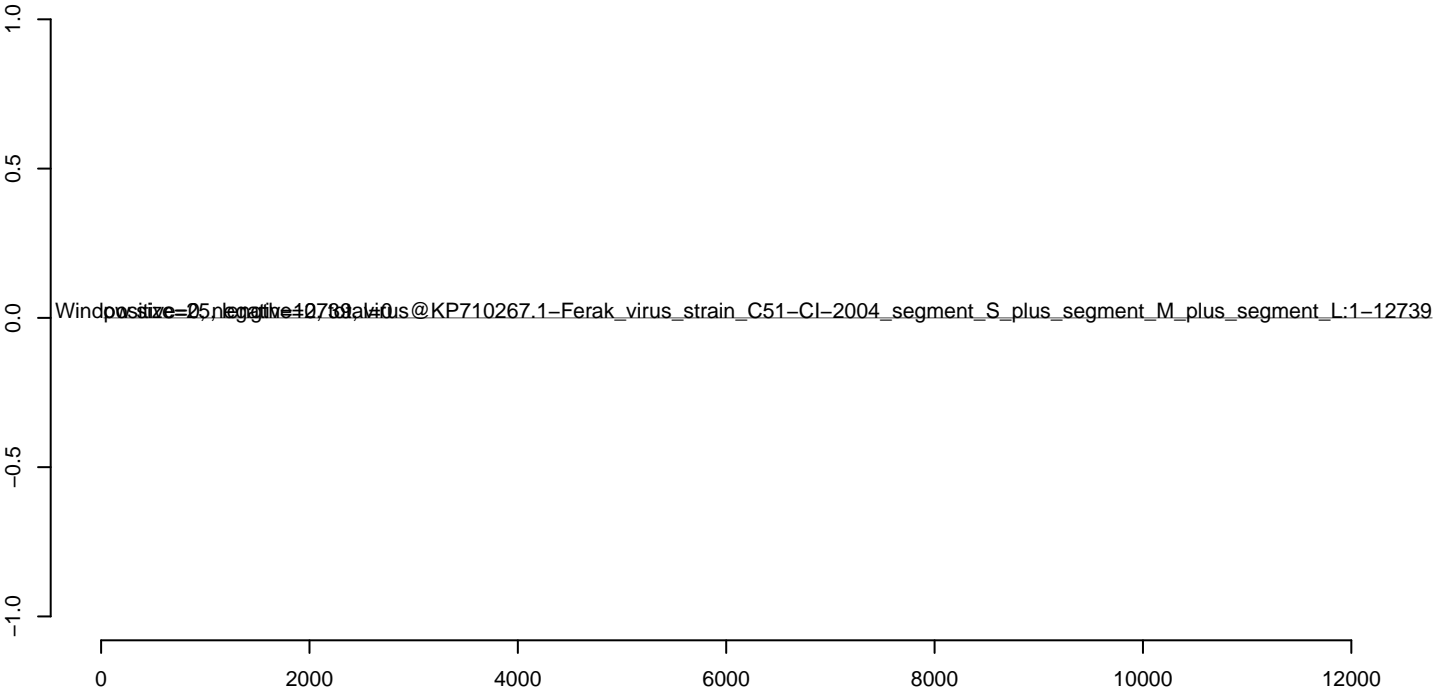
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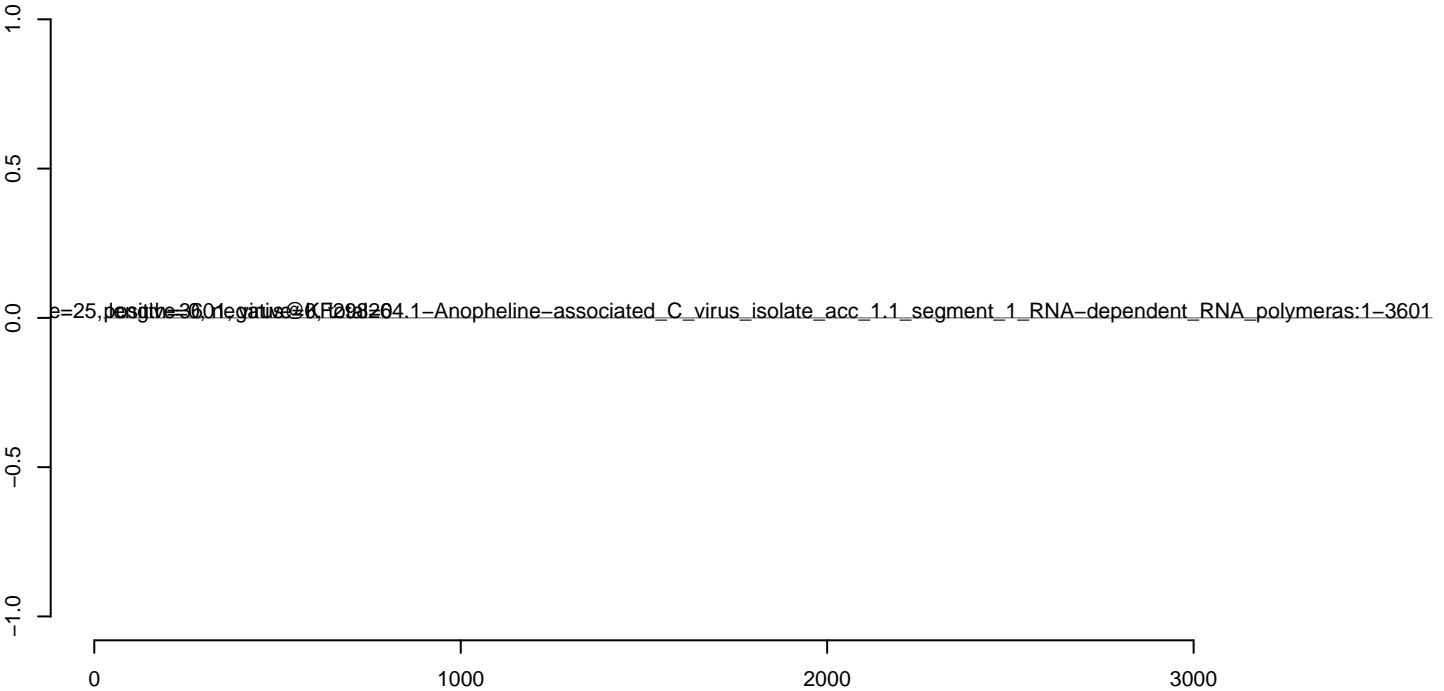
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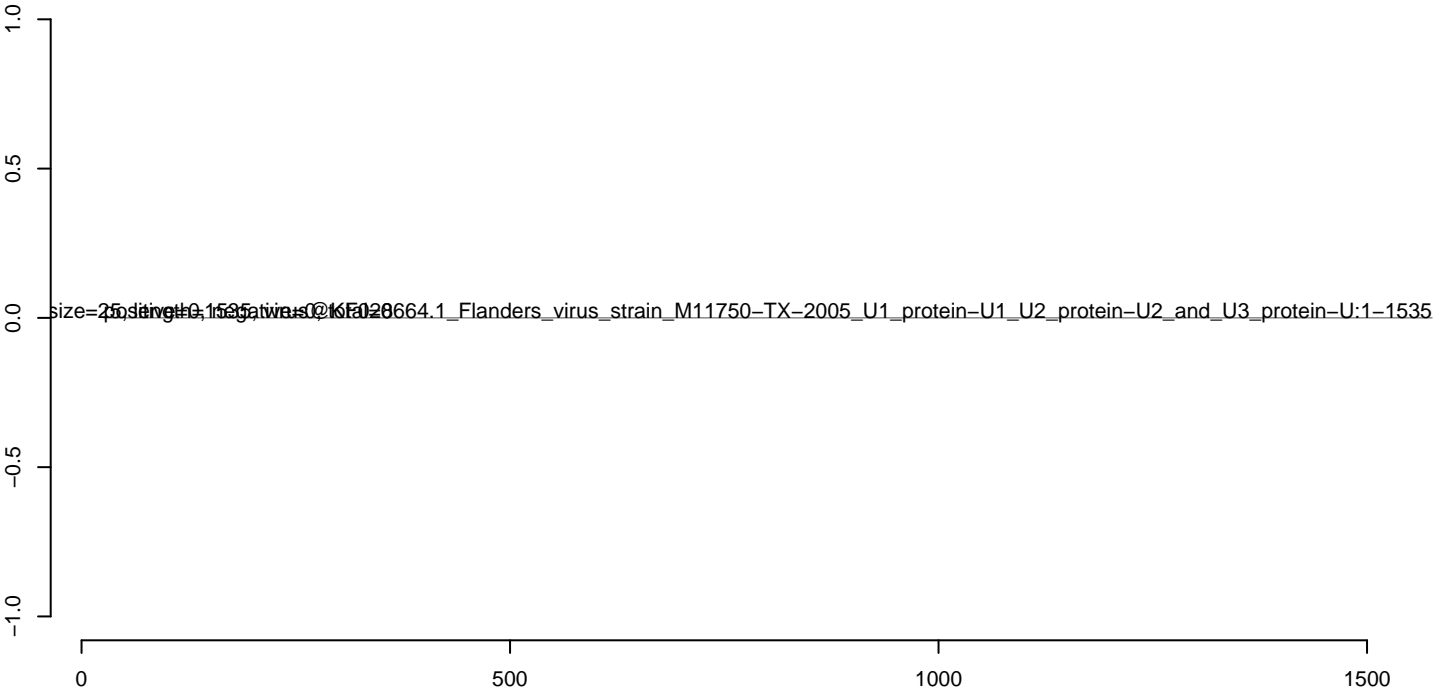
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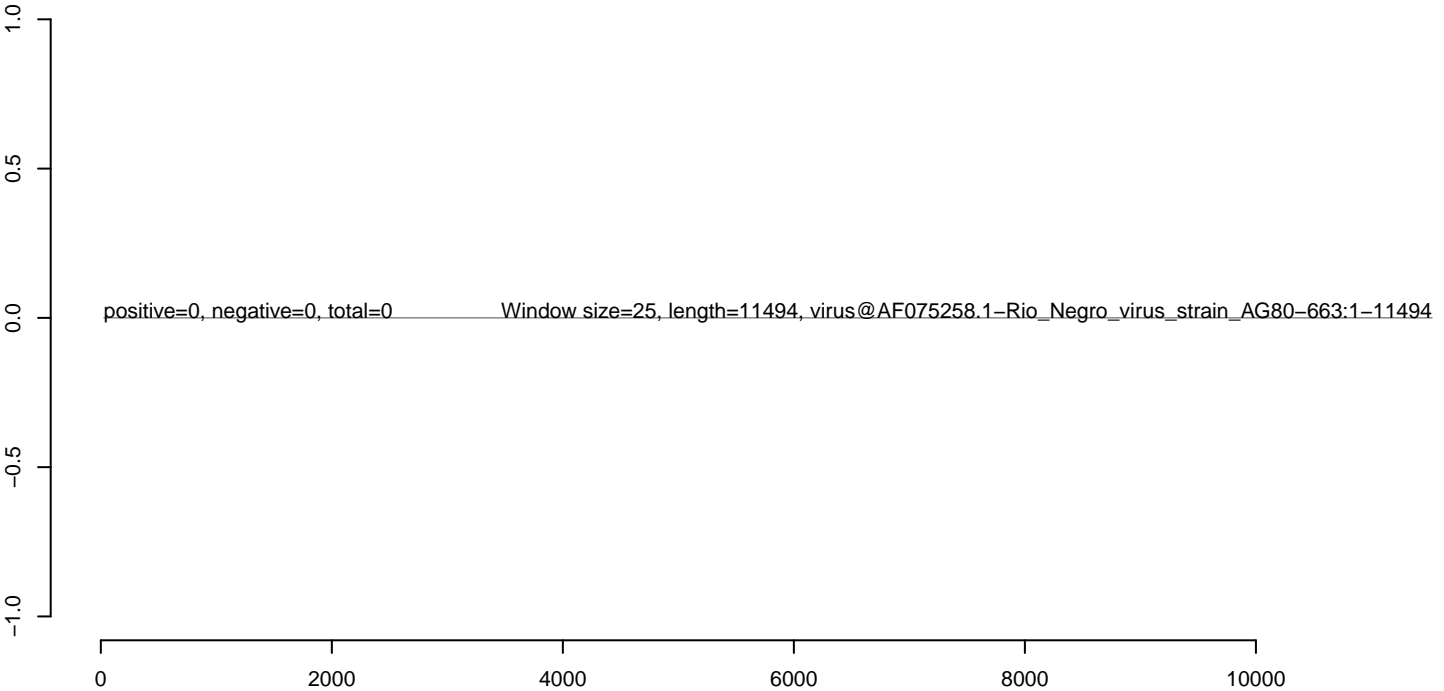
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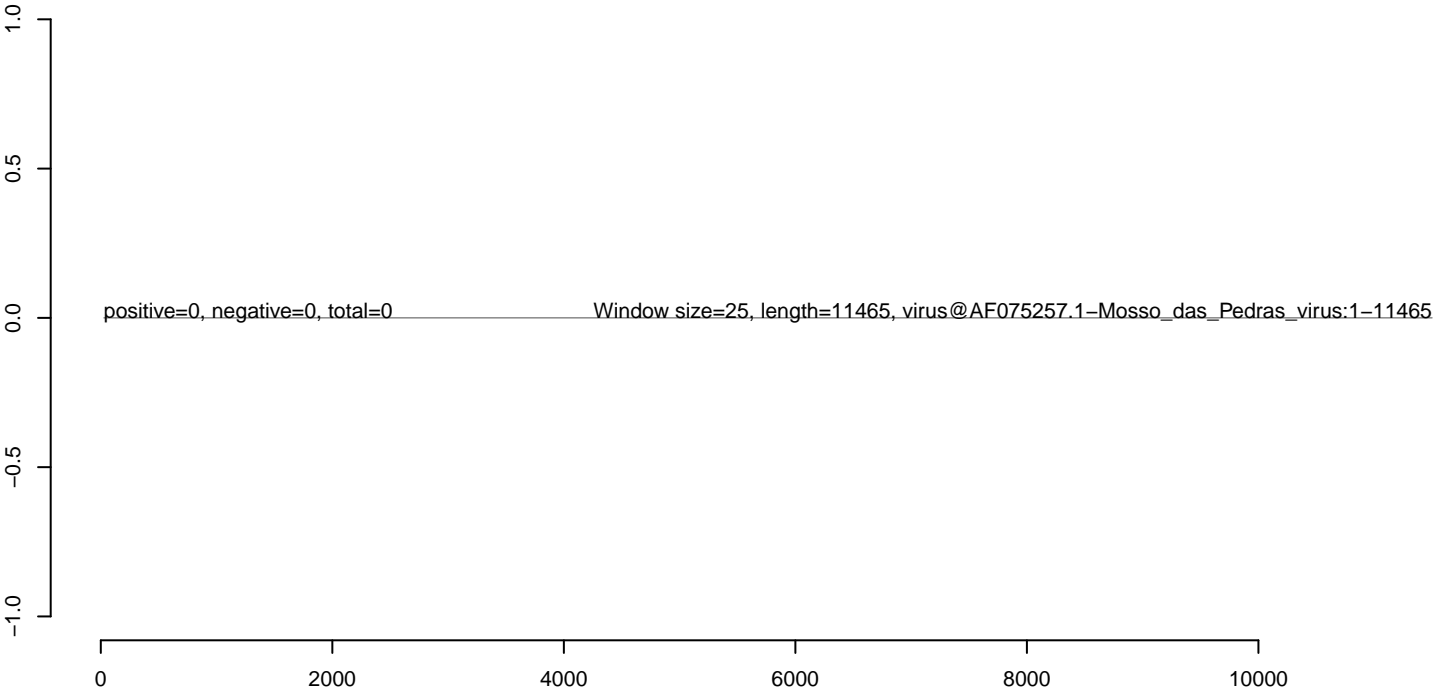
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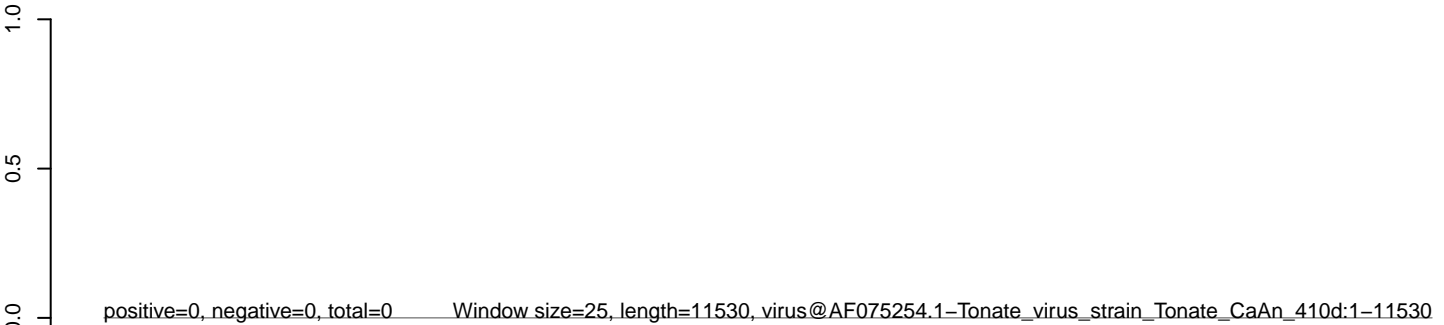
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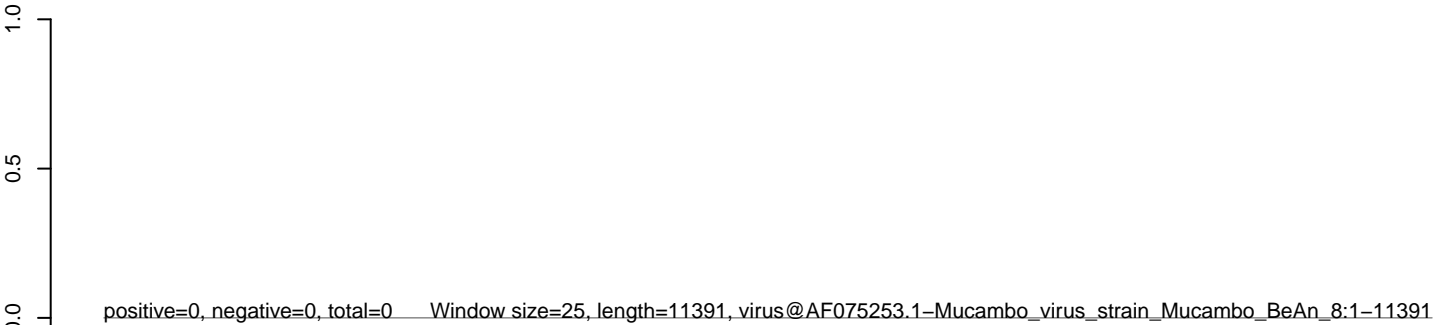
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AeAeg_Aag2DENV2NGC.rep



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