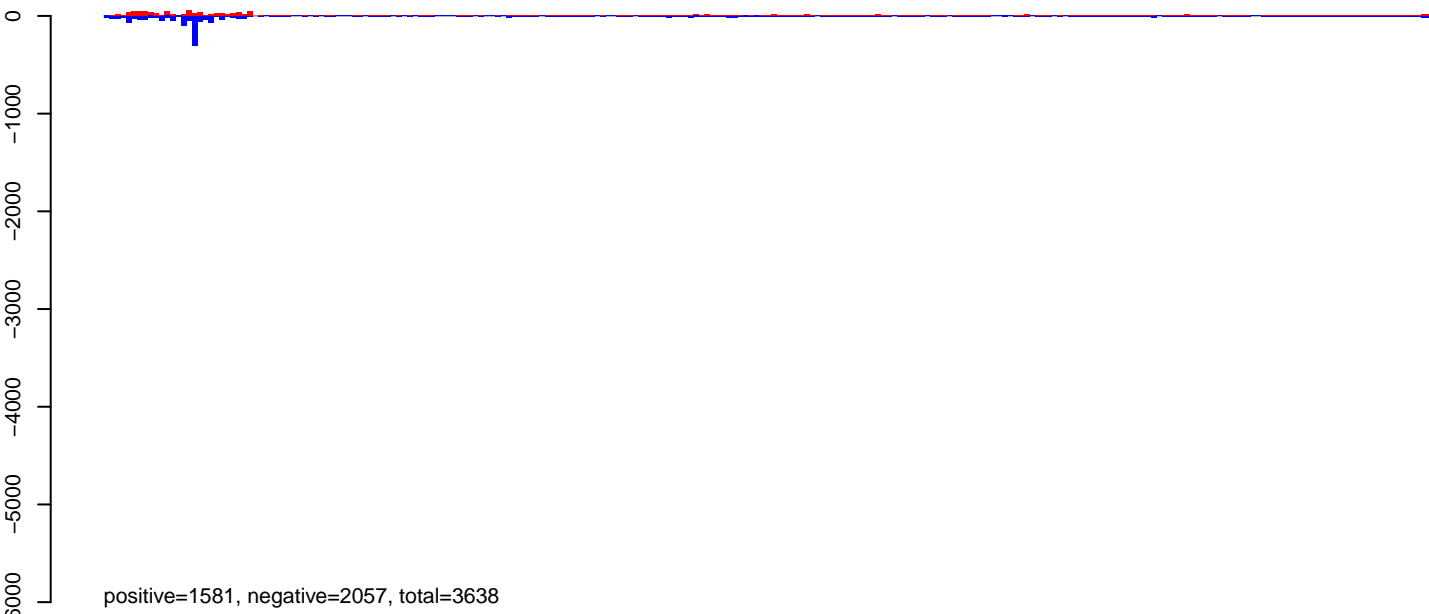
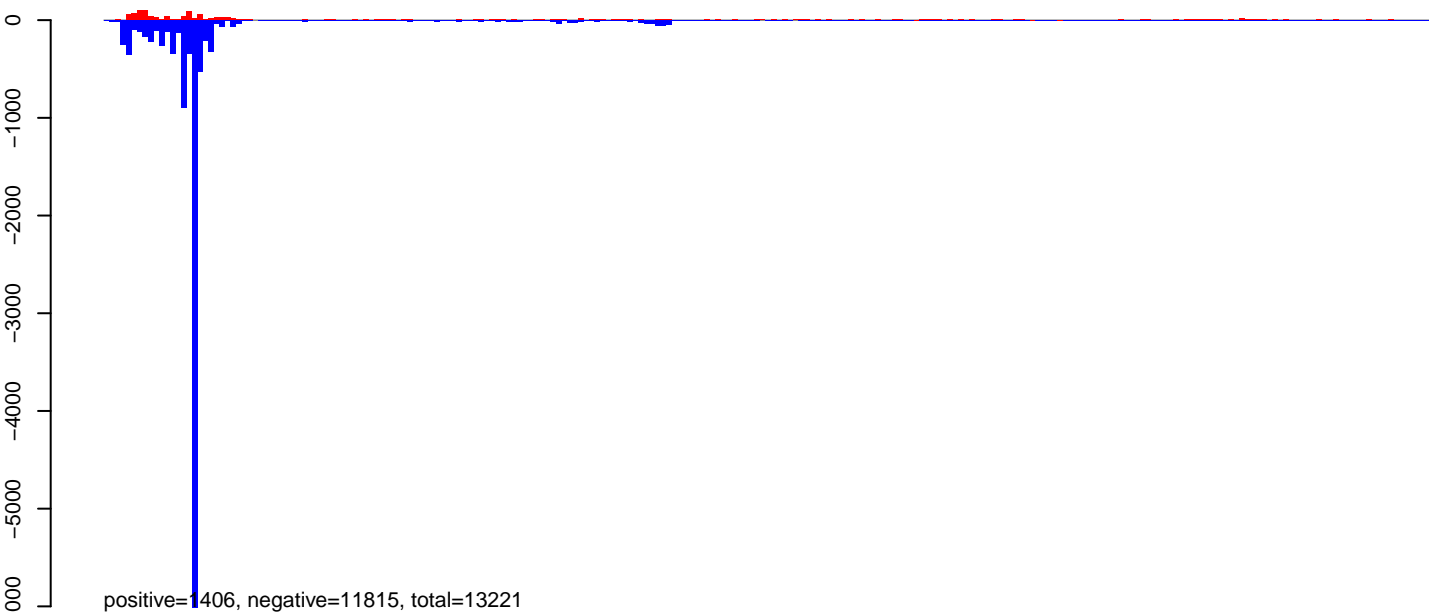


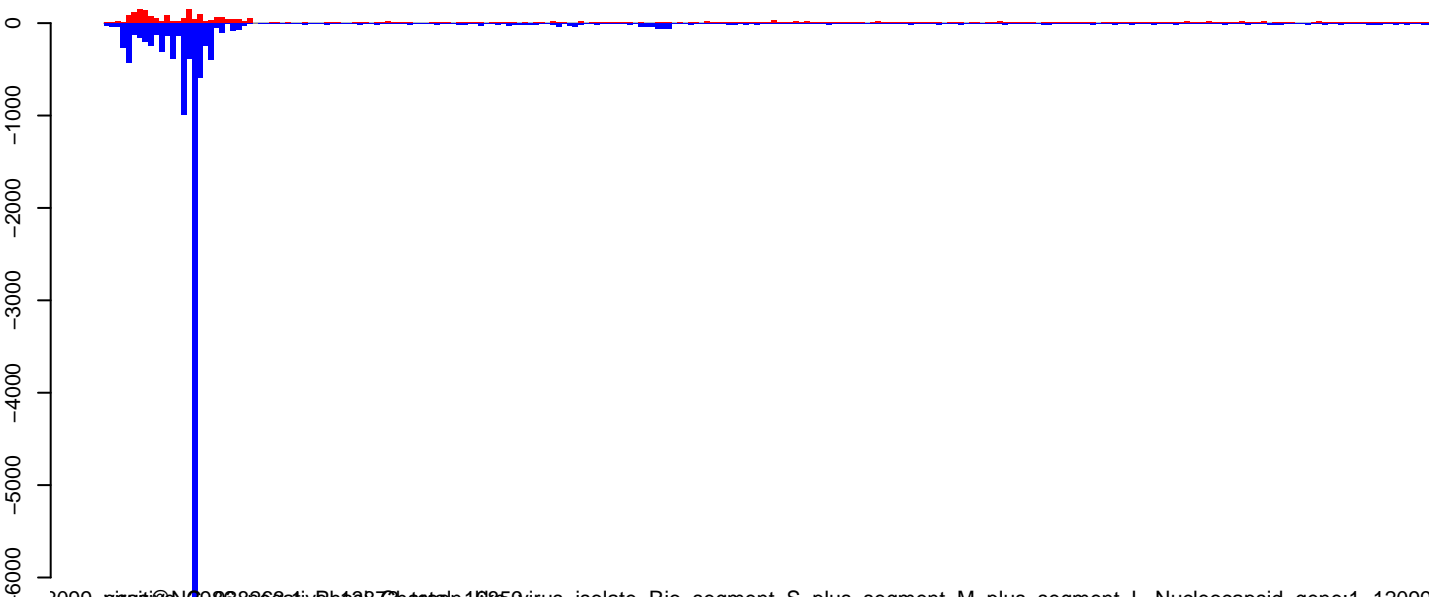
AeAeg_Aag2_DENV4.18_23.rep



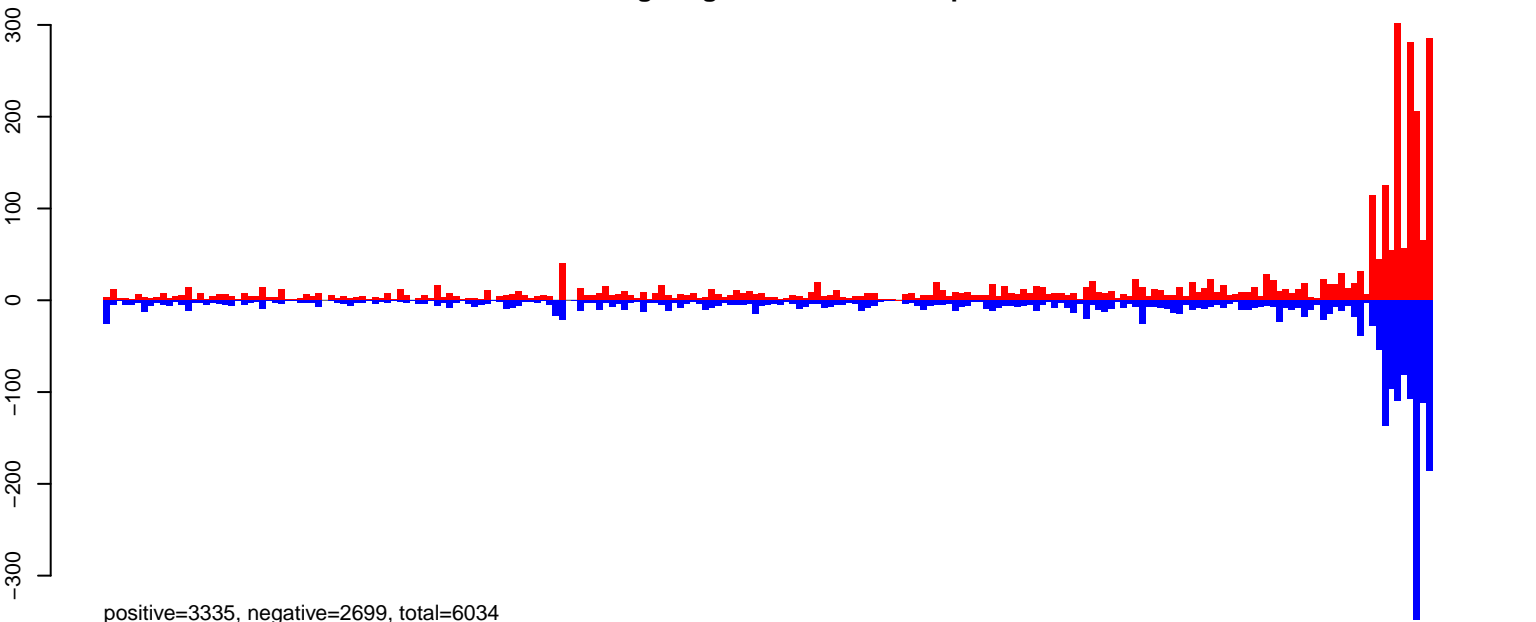
AeAeg_Aag2_DENV4.24_35.rep



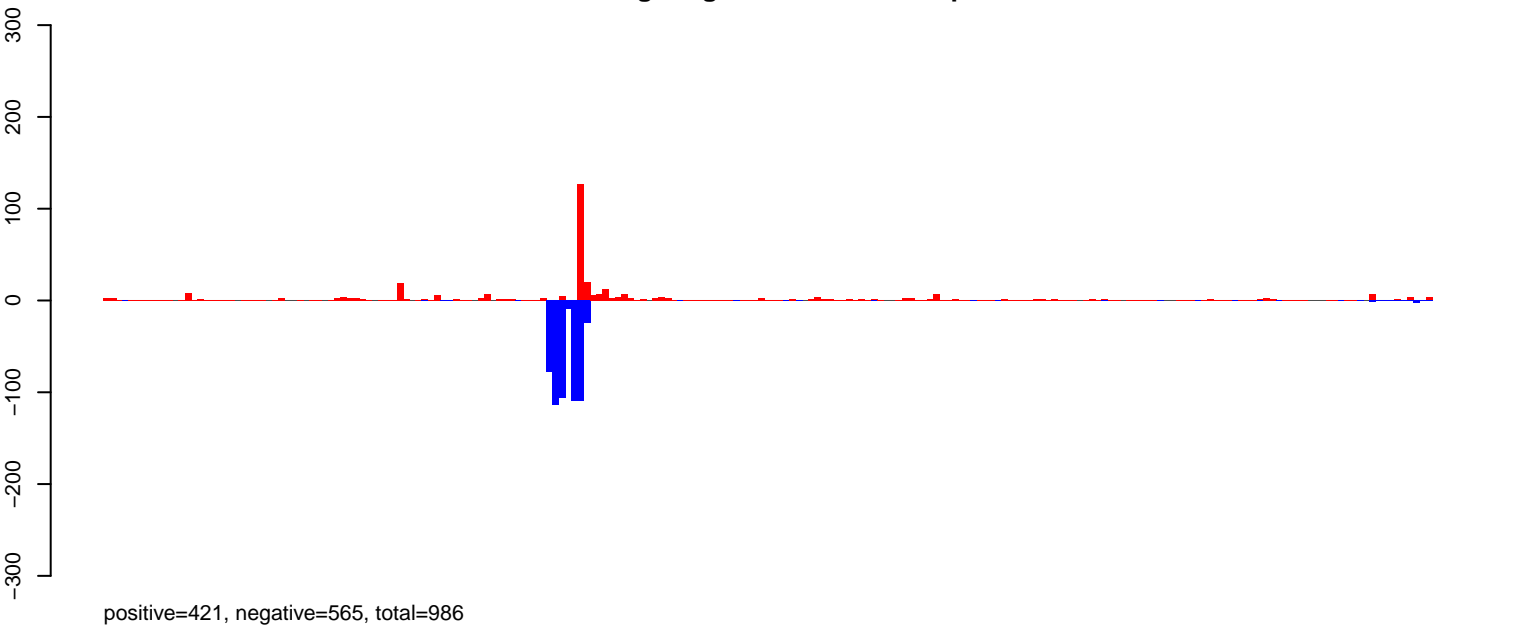
AeAeg_Aag2_DENV4.rep



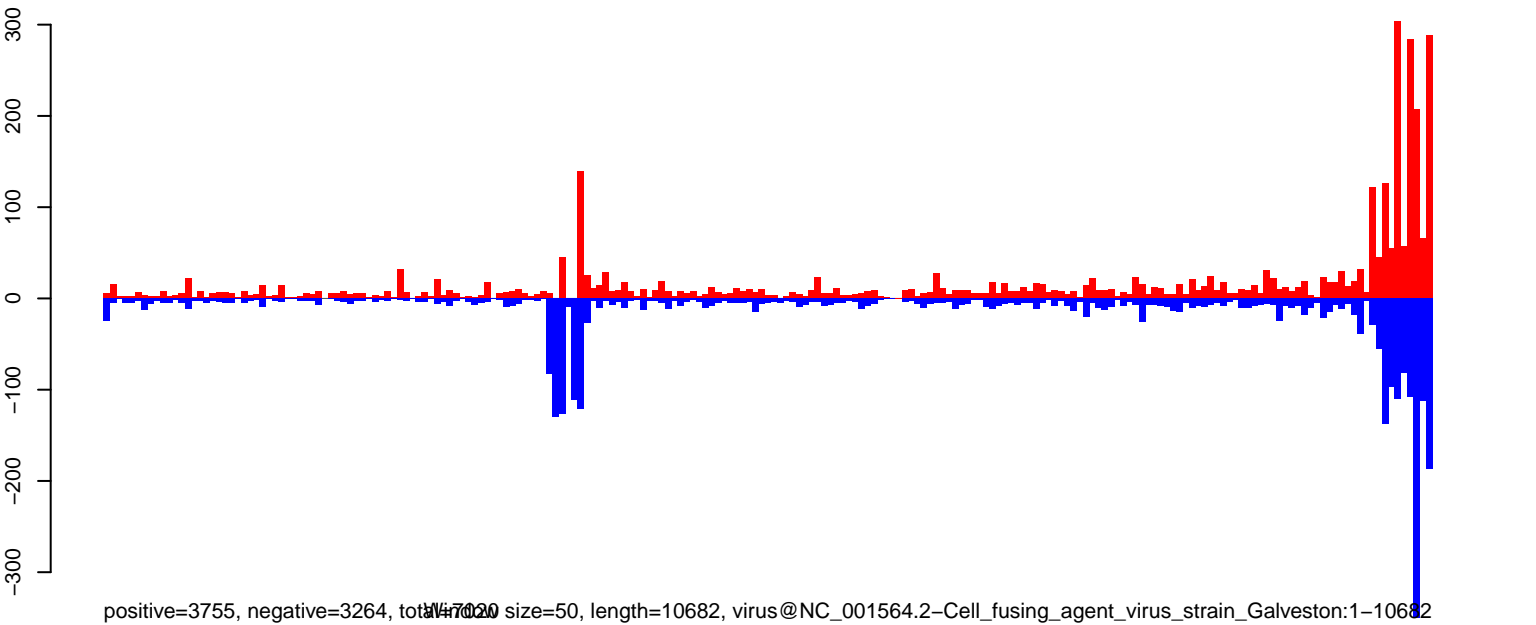
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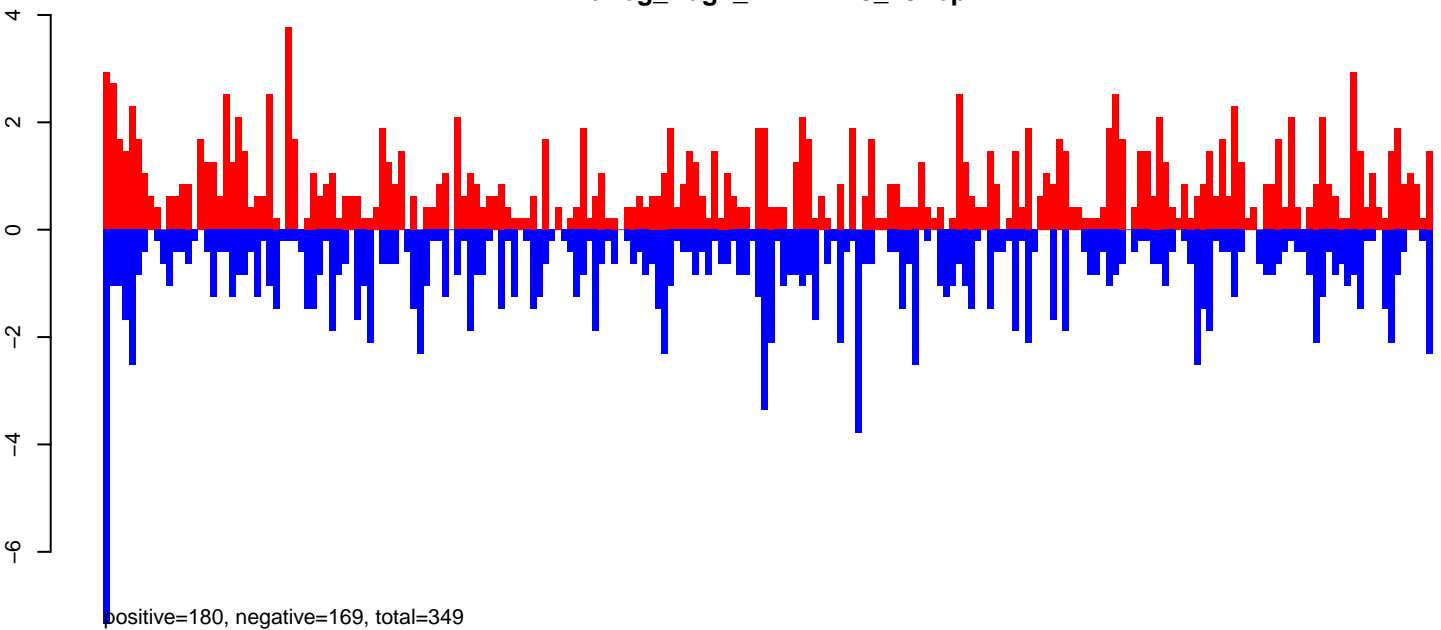
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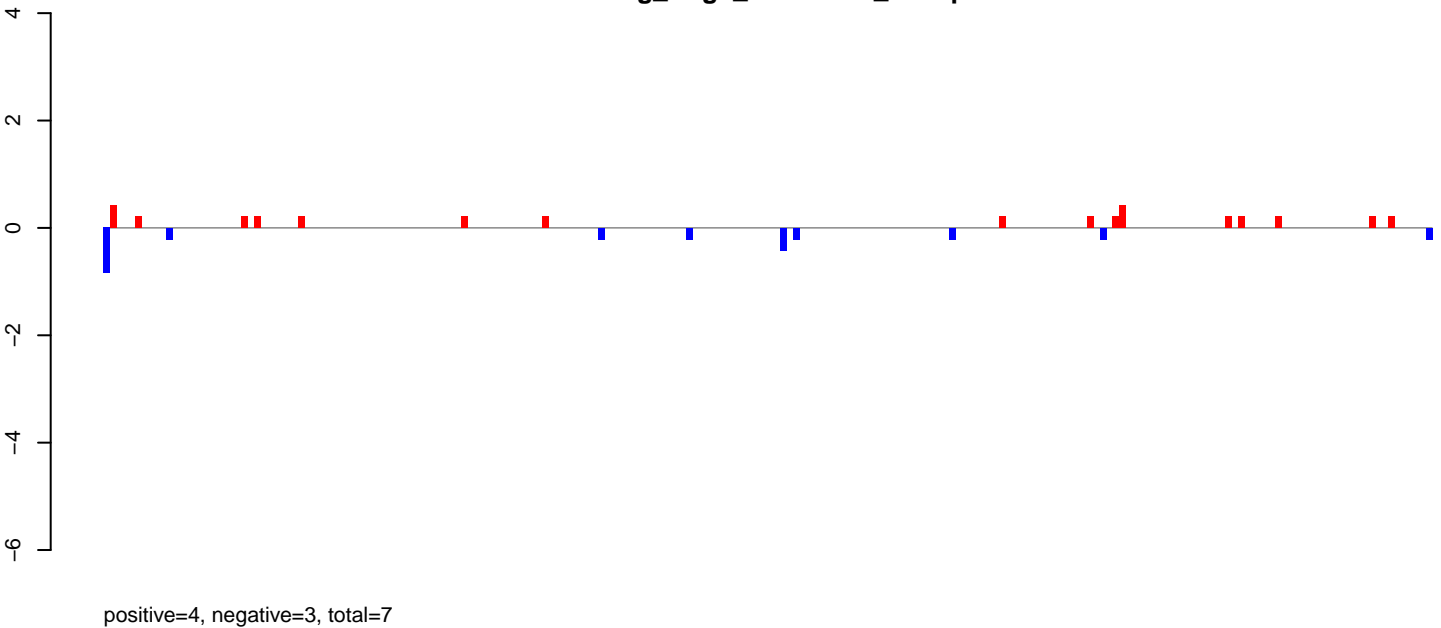
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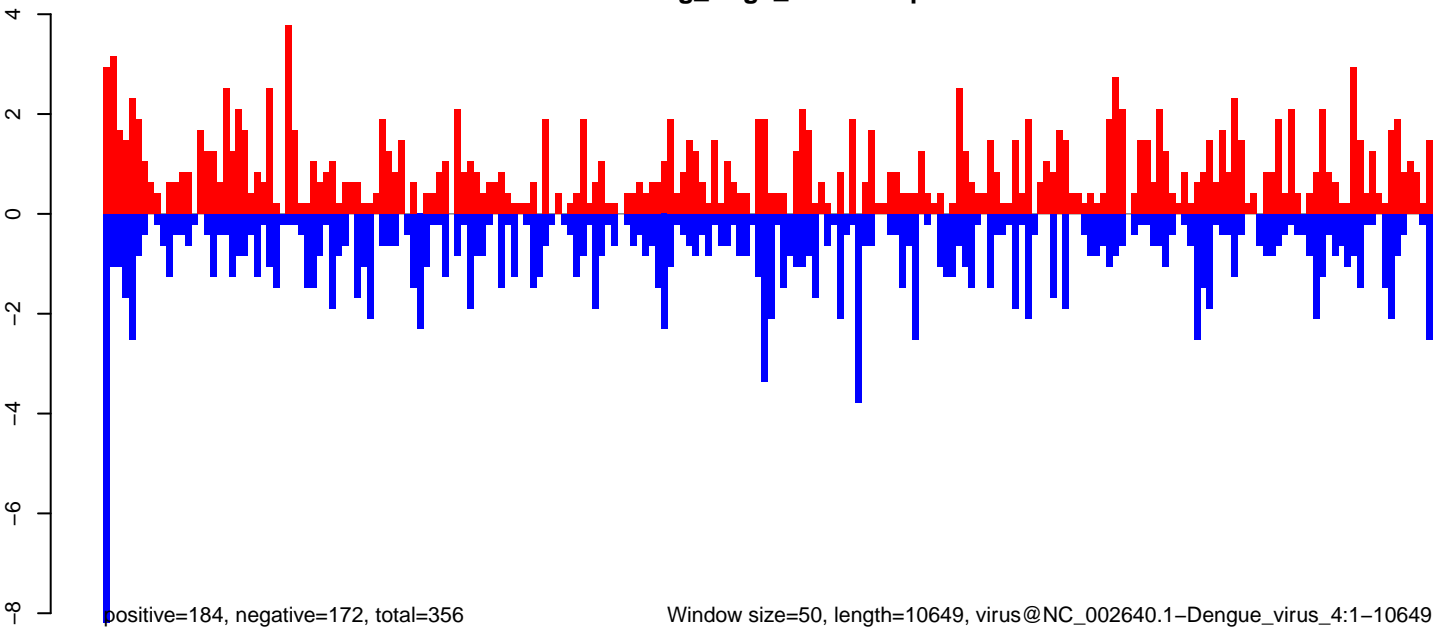
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

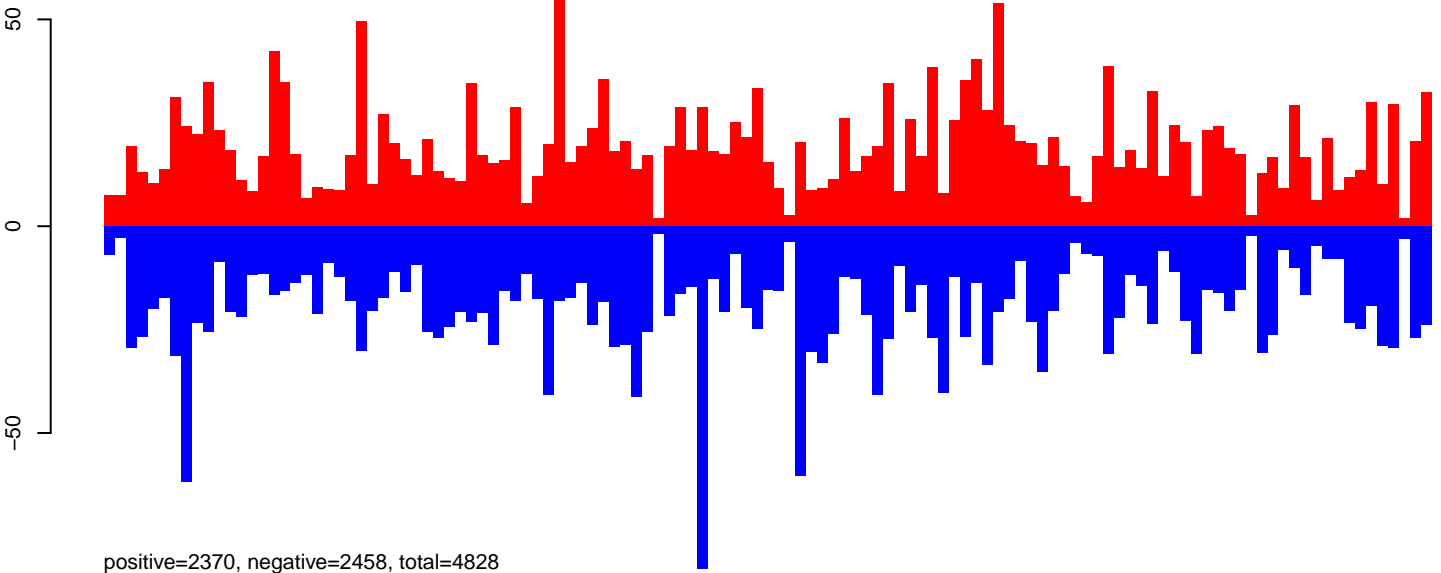


AeAeg_Aag2_DENV4.rep

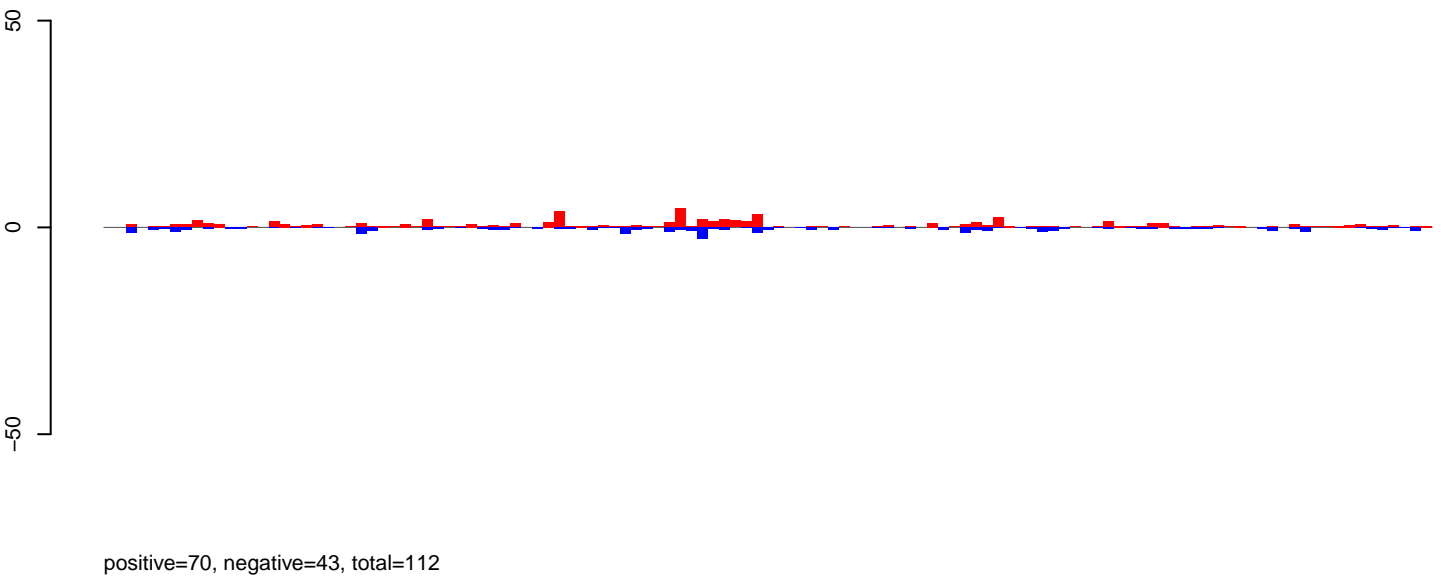


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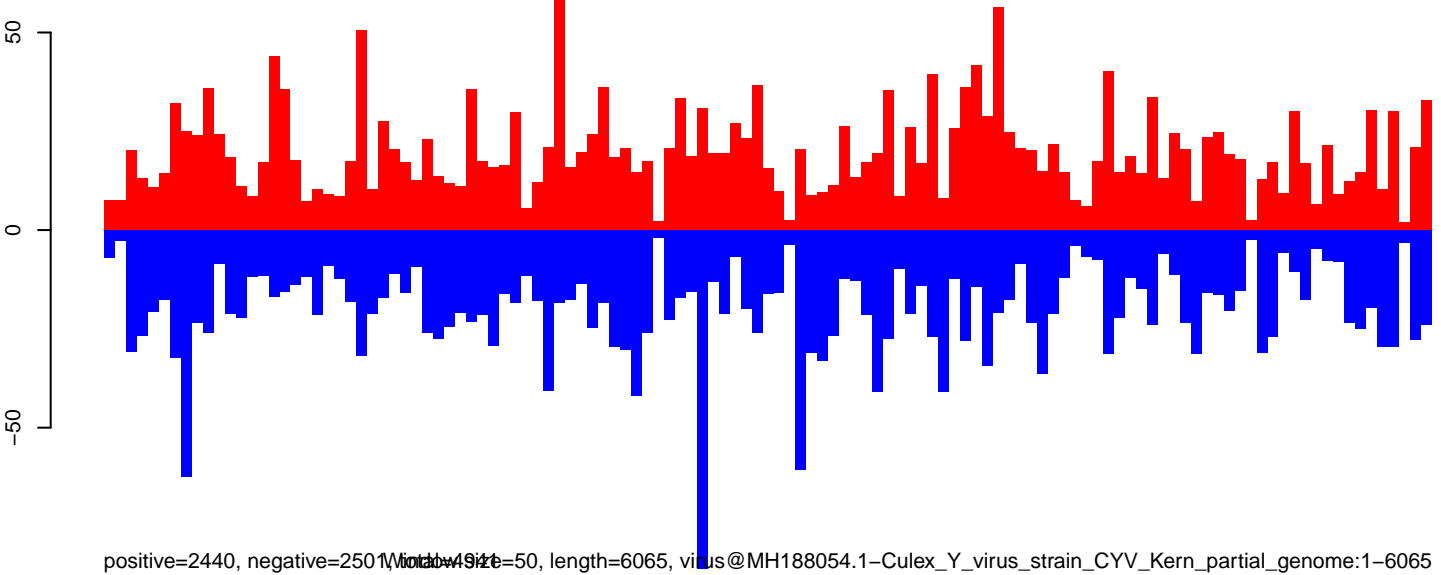
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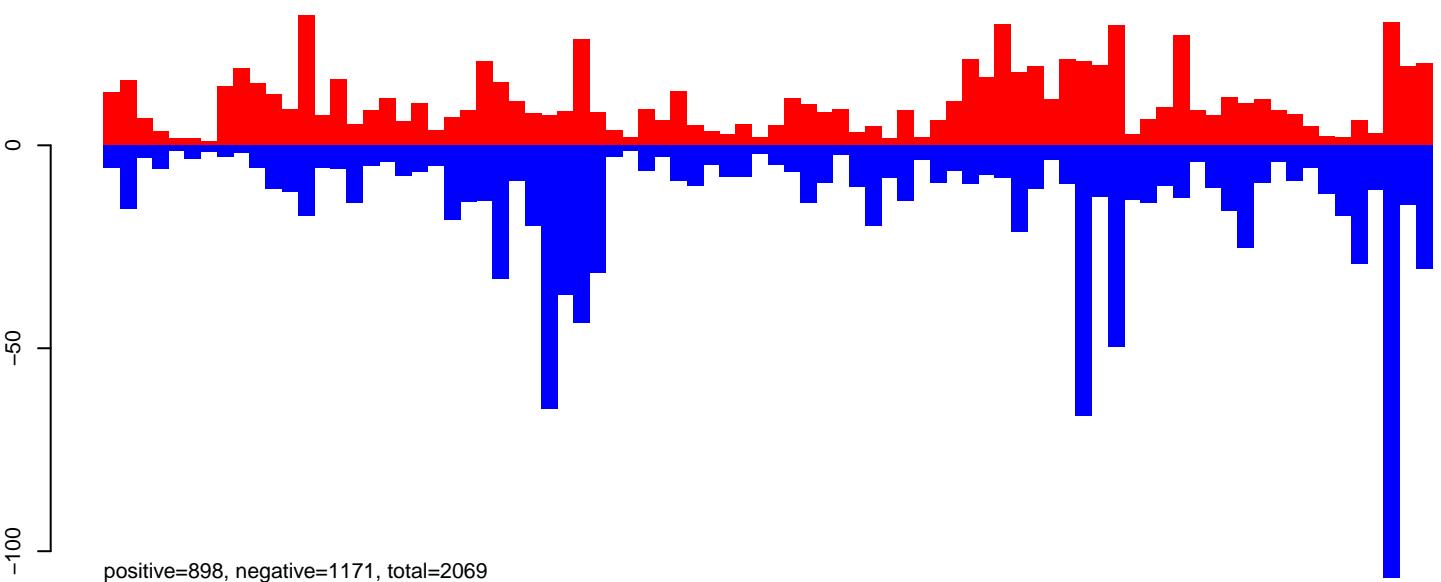
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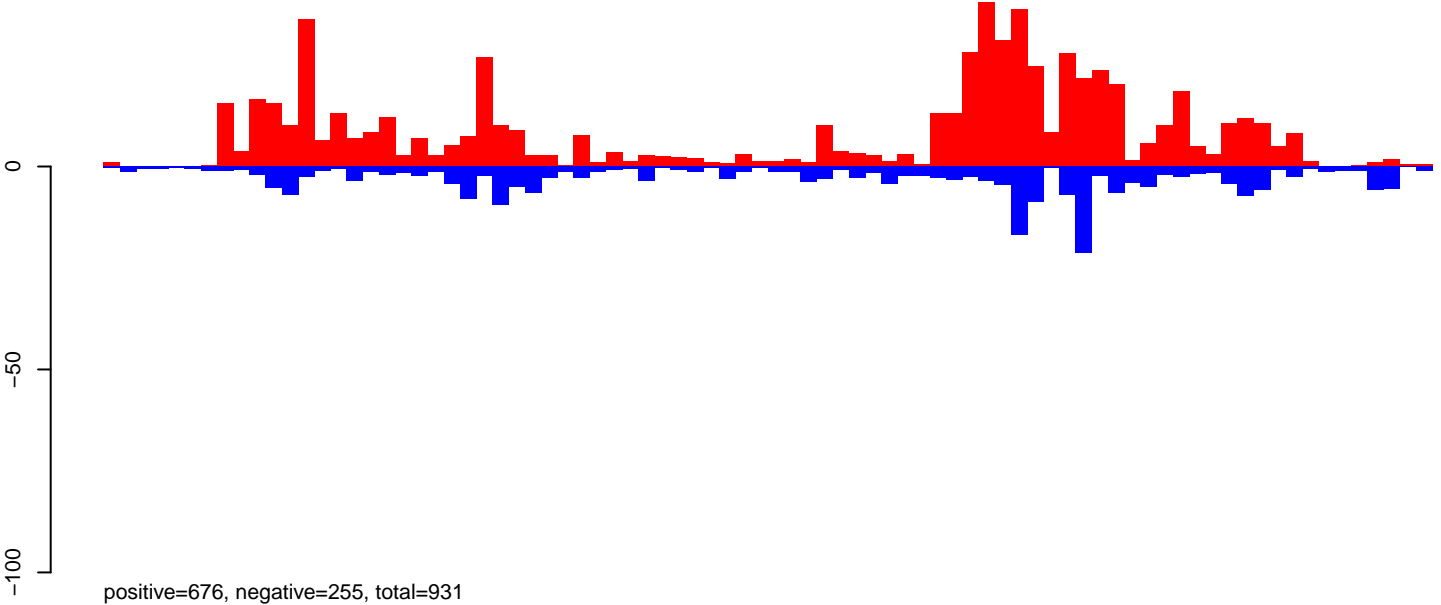
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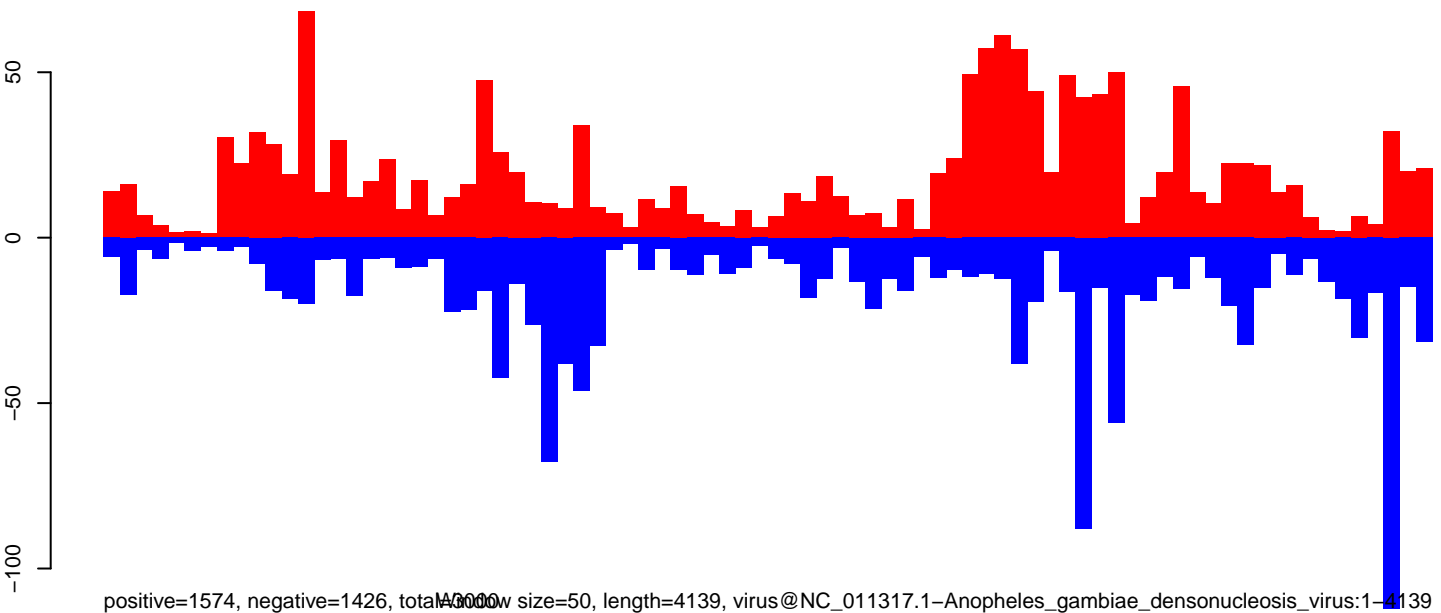
AeAeg_Aag2_DENV4.18_23.rep



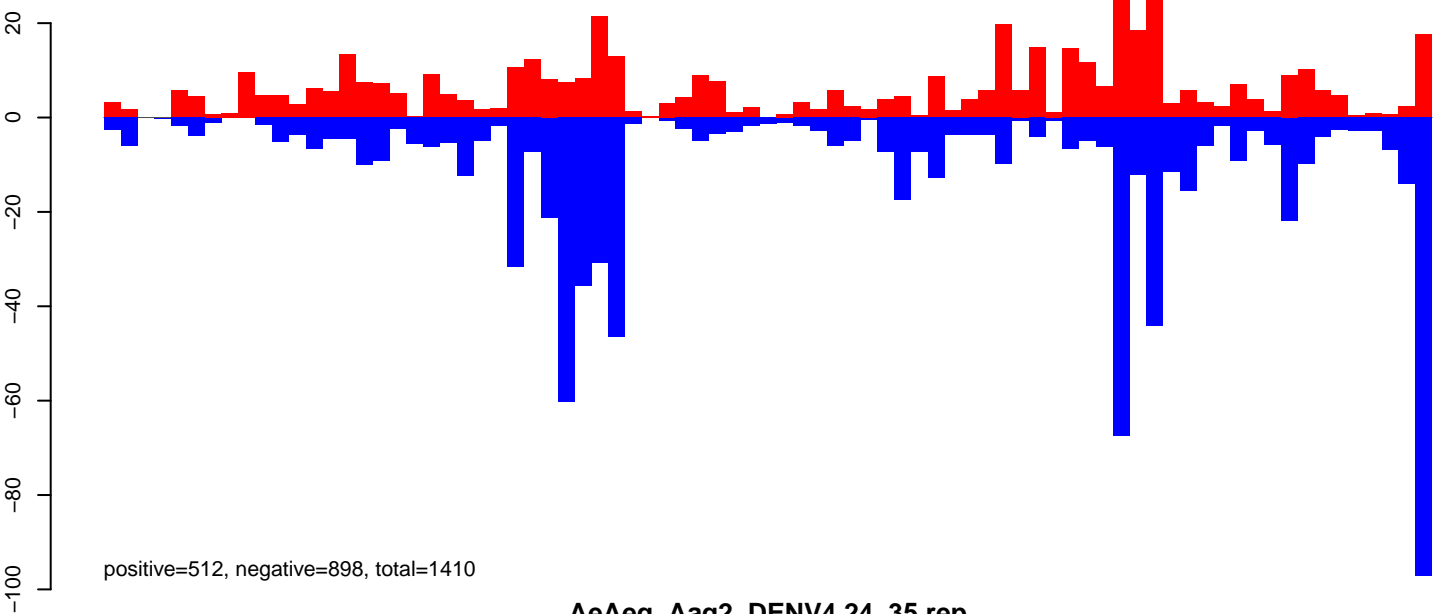
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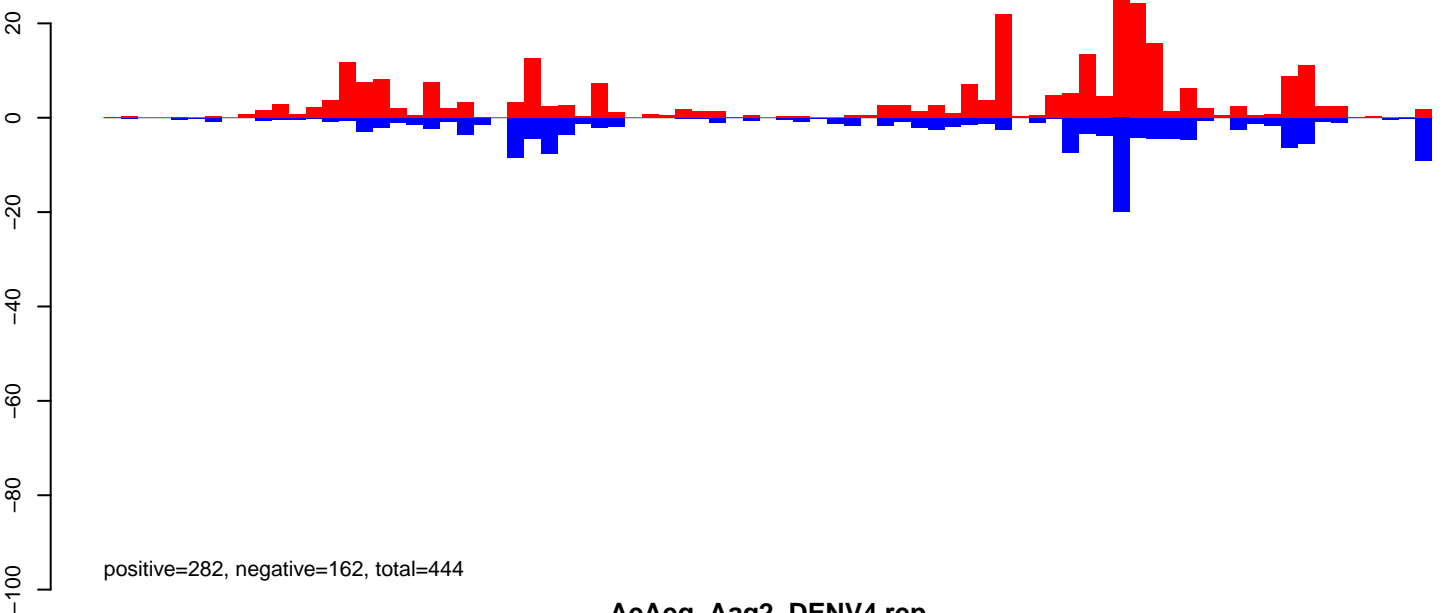
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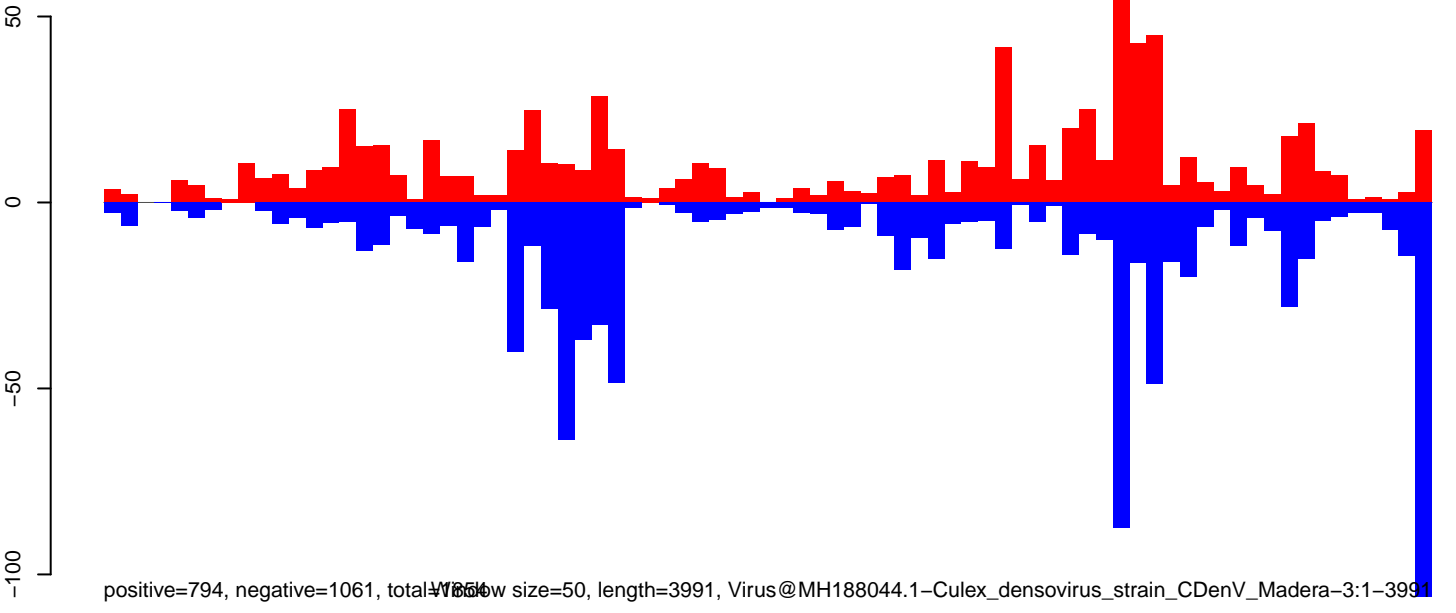
AeAeg_Aag2_DENV4.18_23.rep



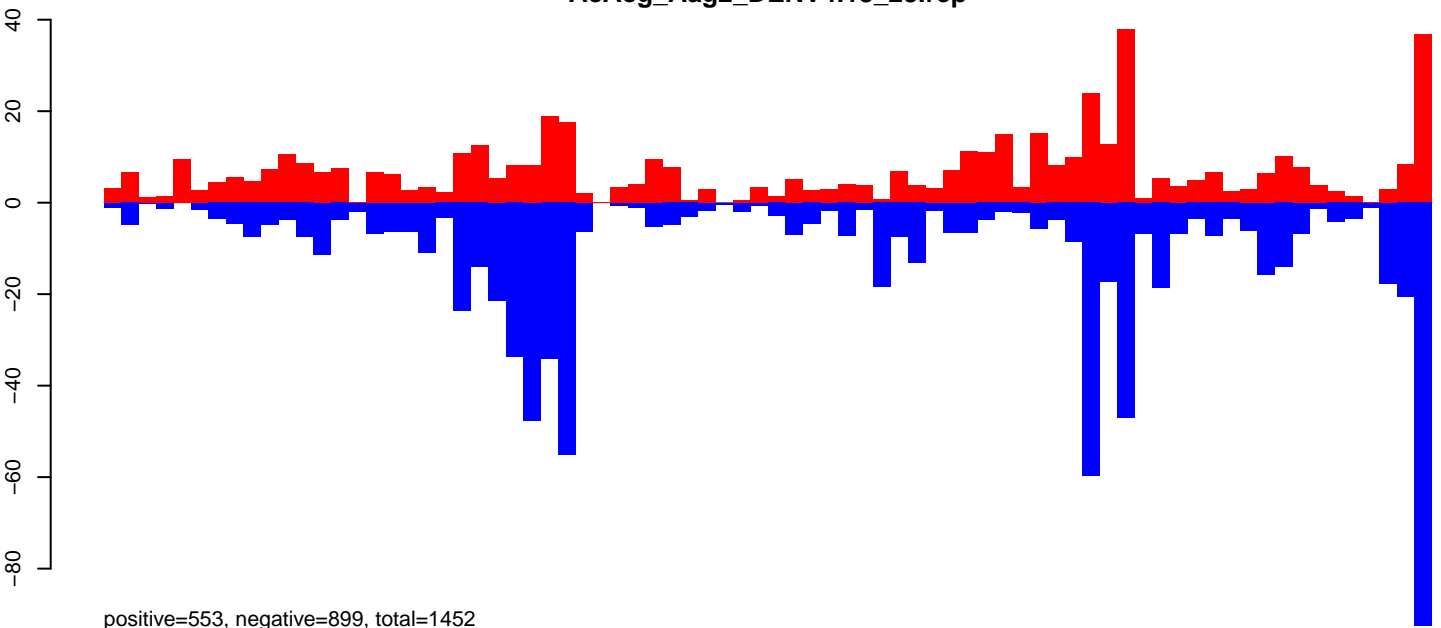
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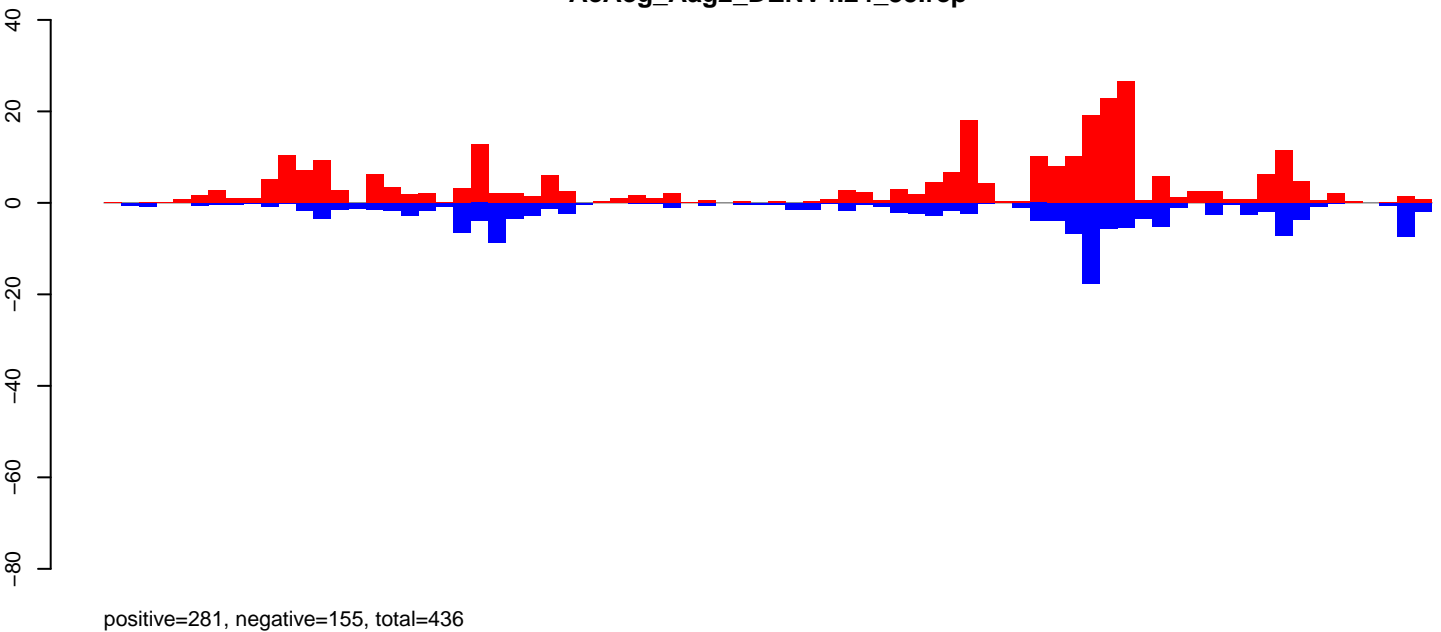
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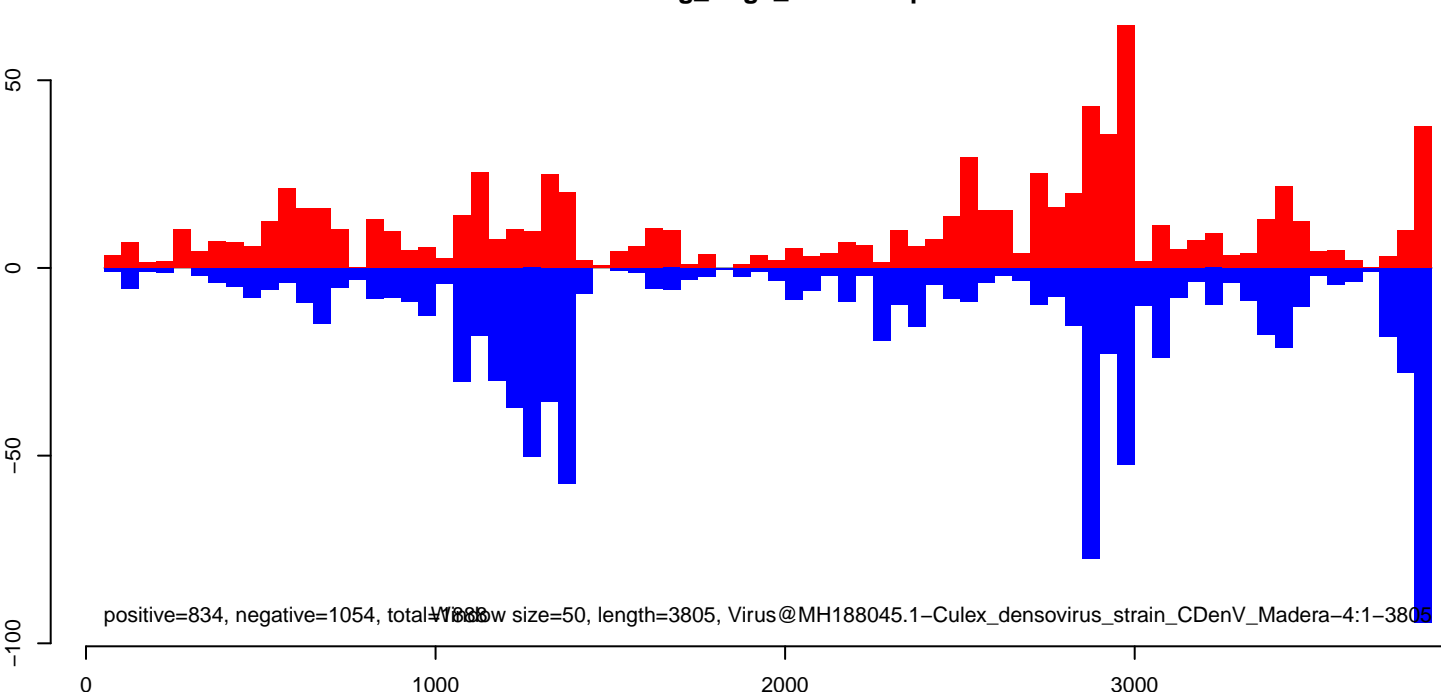
AeAeg_Aag2_DENV4.18_23.rep



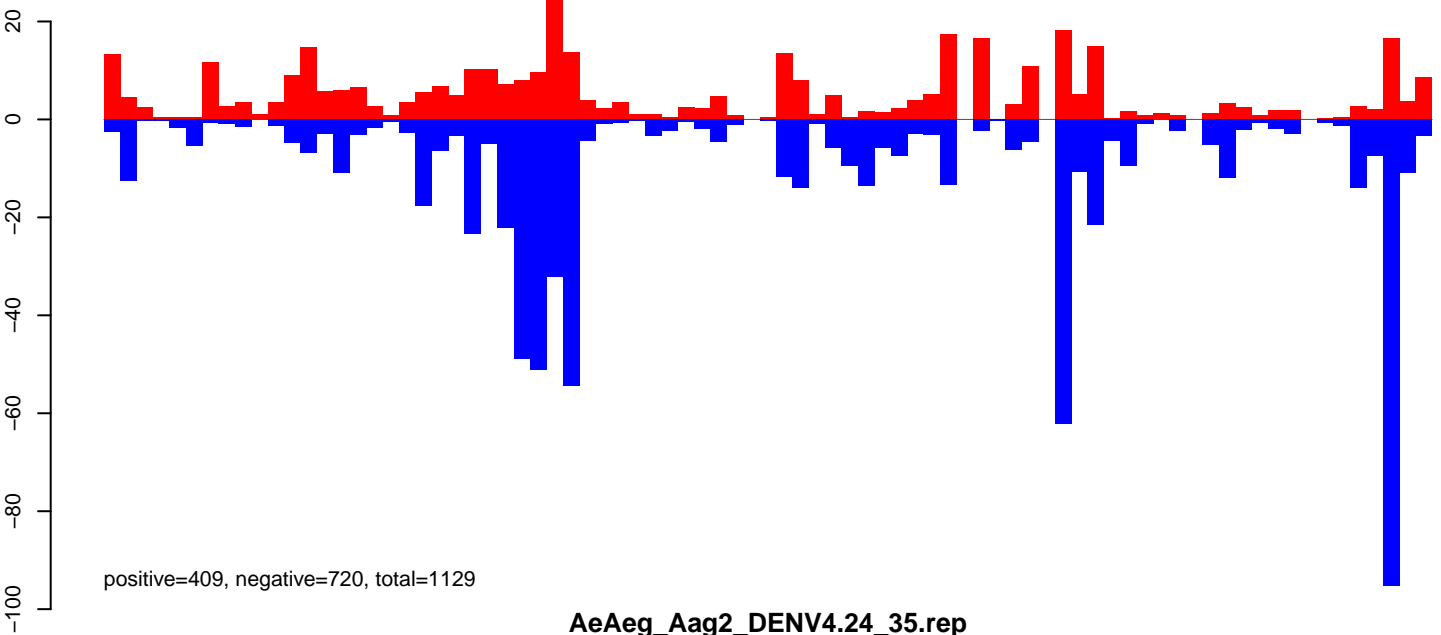
AeAeg_Aag2_DENV4.24_35.rep



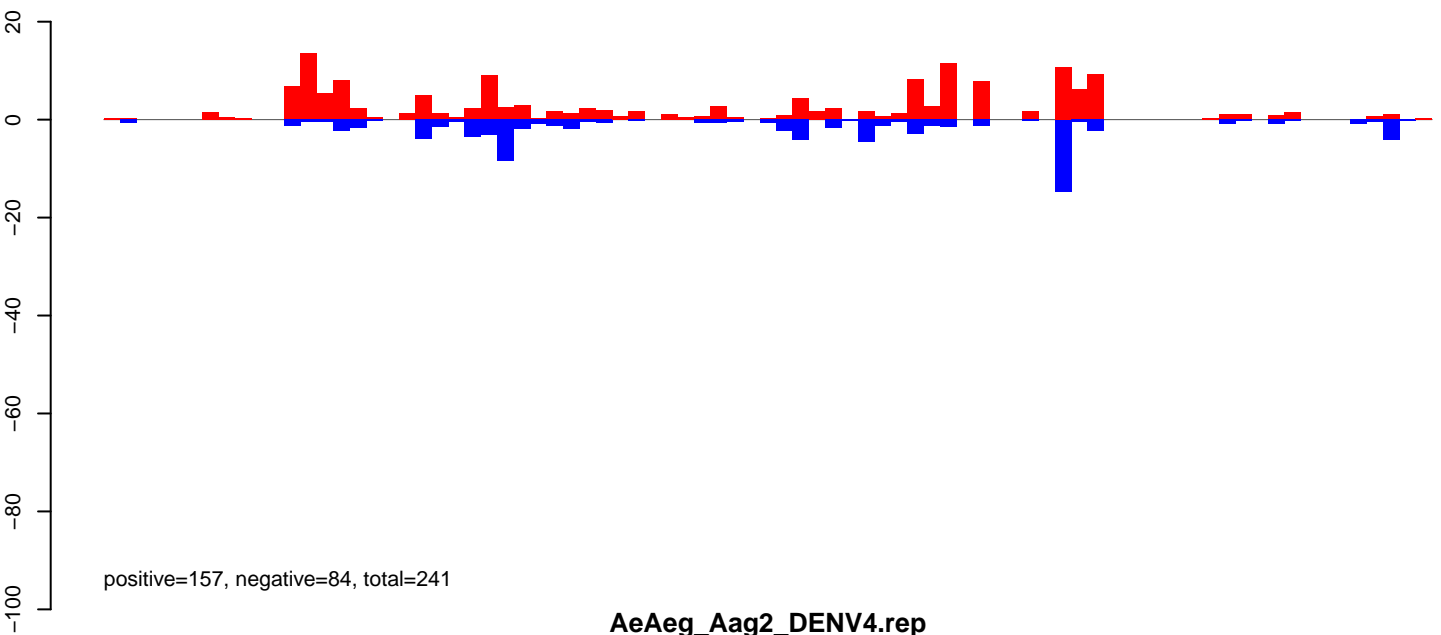
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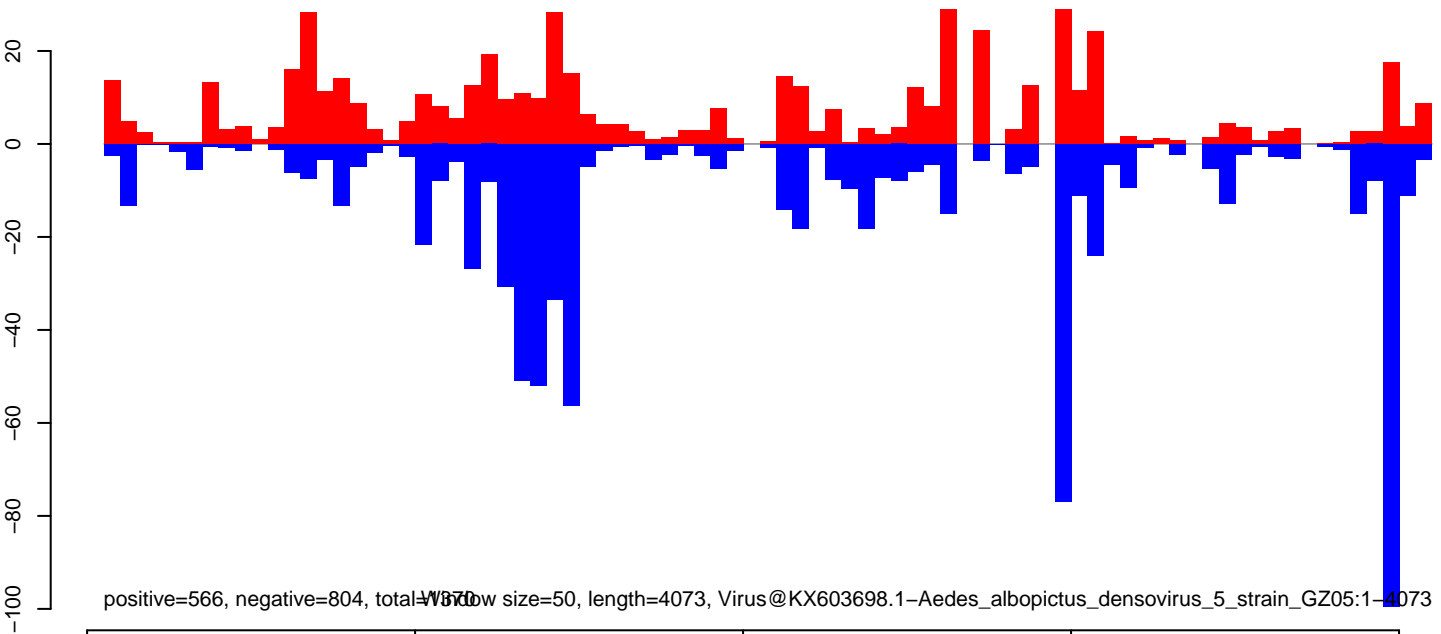
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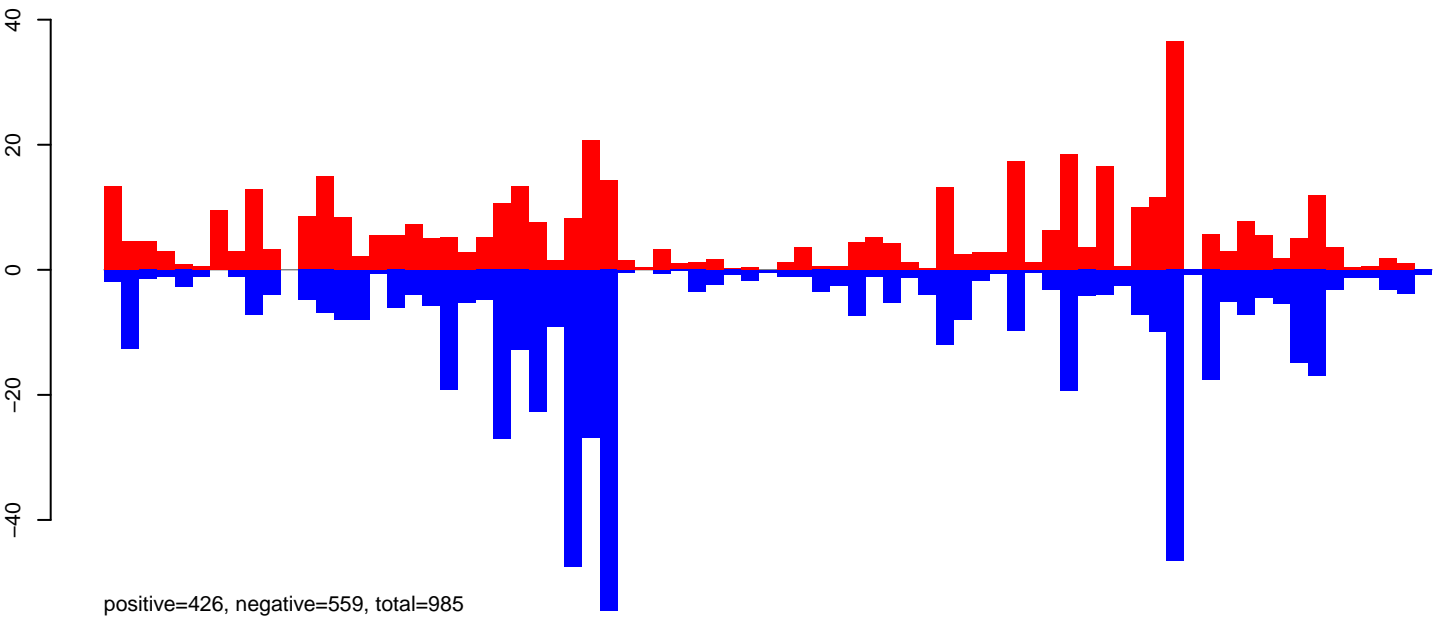
AeAeg_Aag2_DENV4.24_35.rep



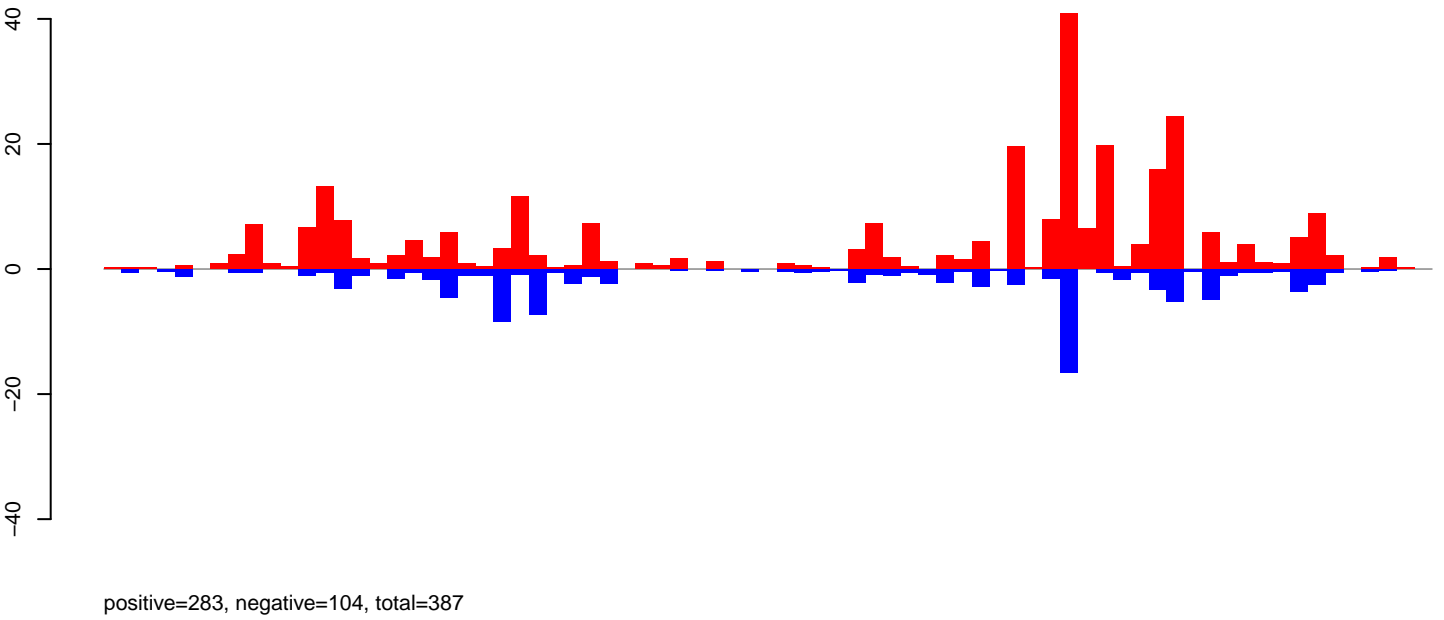
AeAeg_Aag2_DENV4.rep



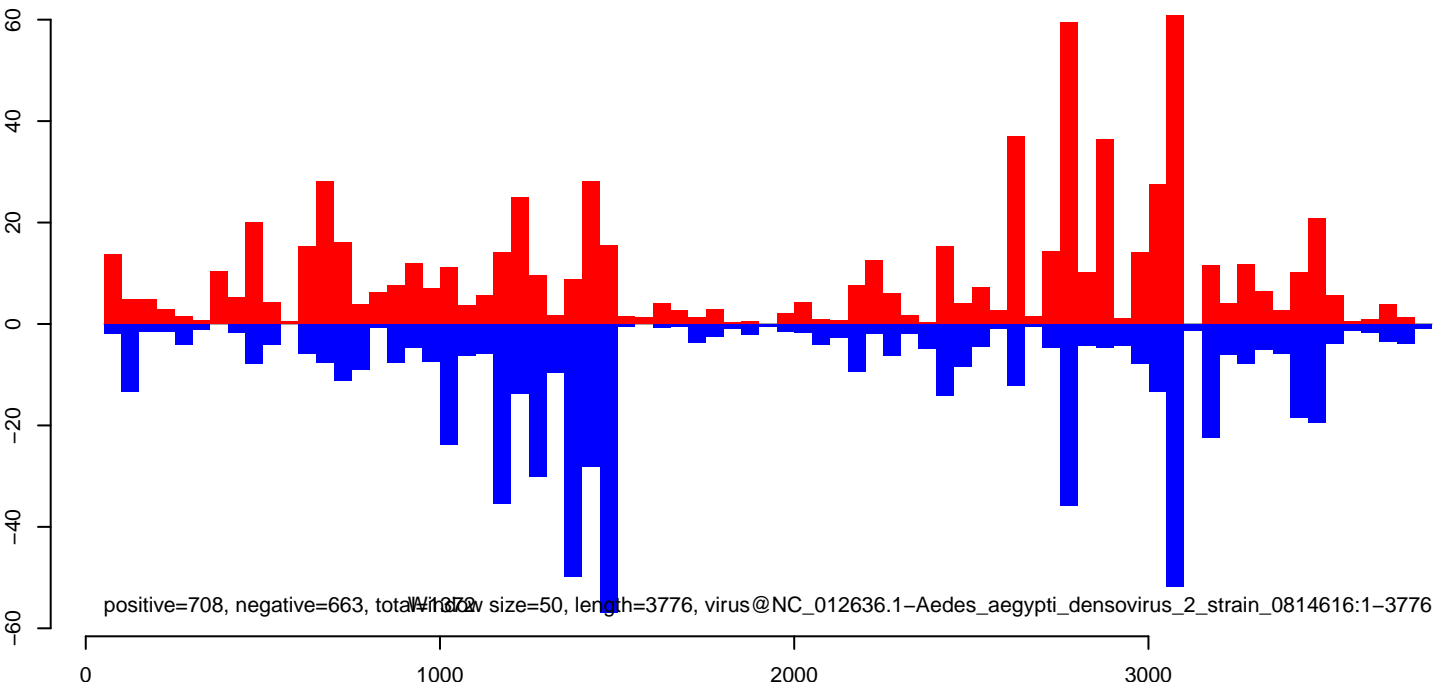
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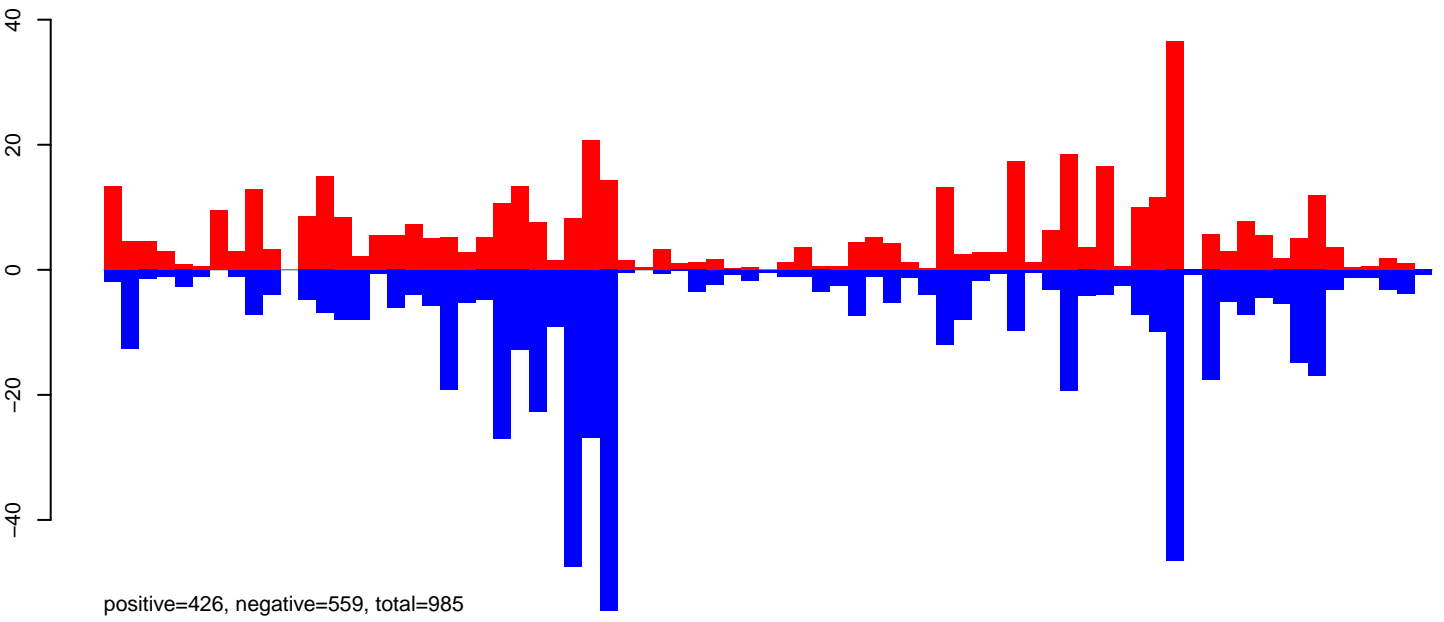
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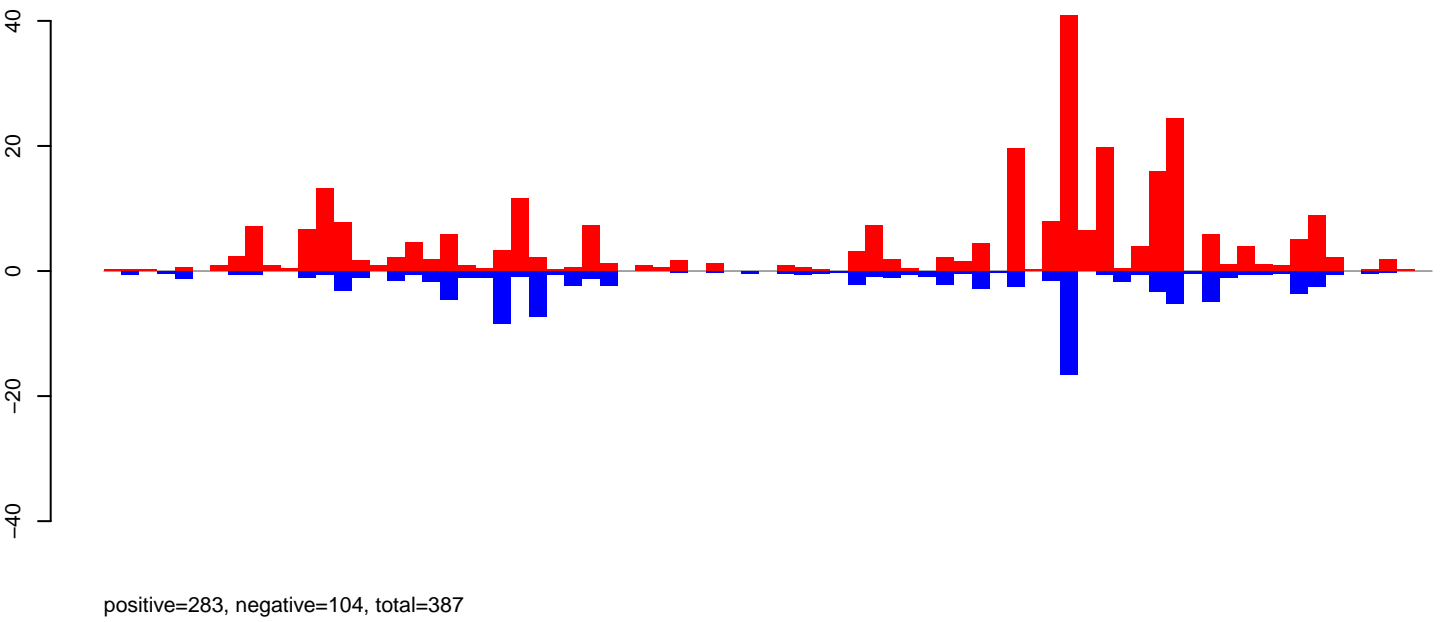
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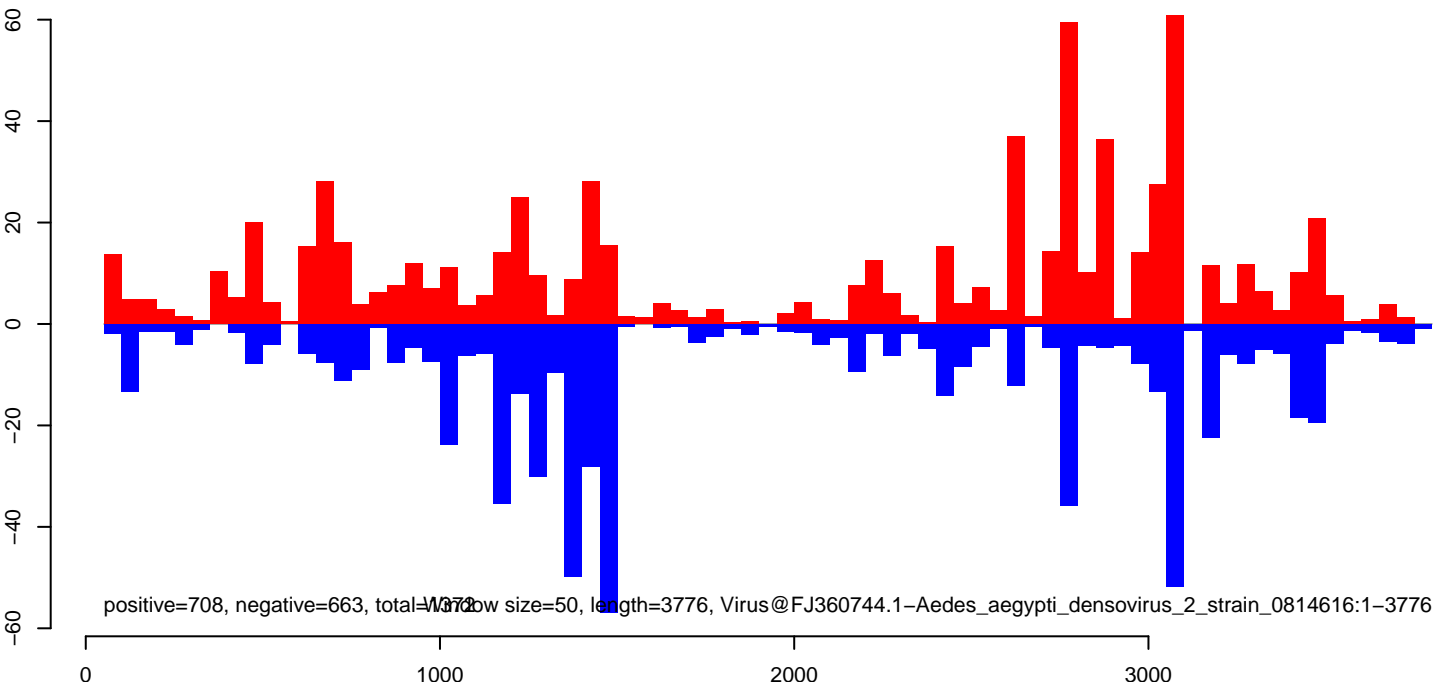
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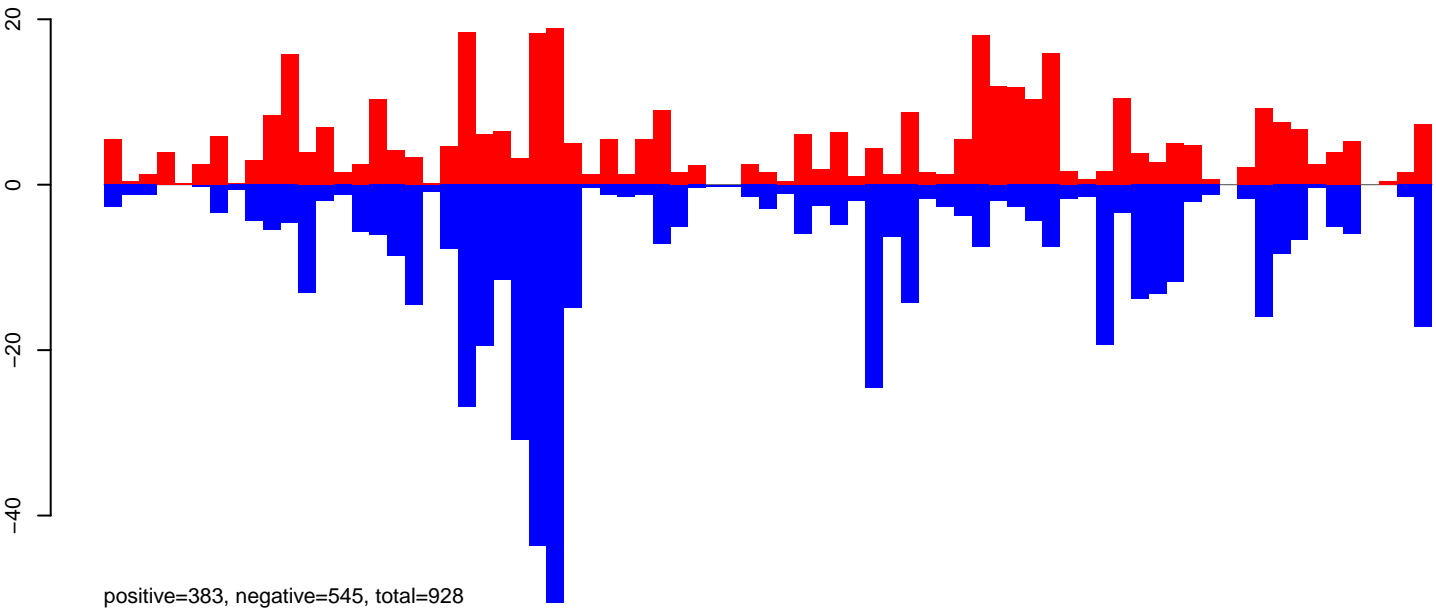
AeAeg_Aag2_DENV4.24_35.rep



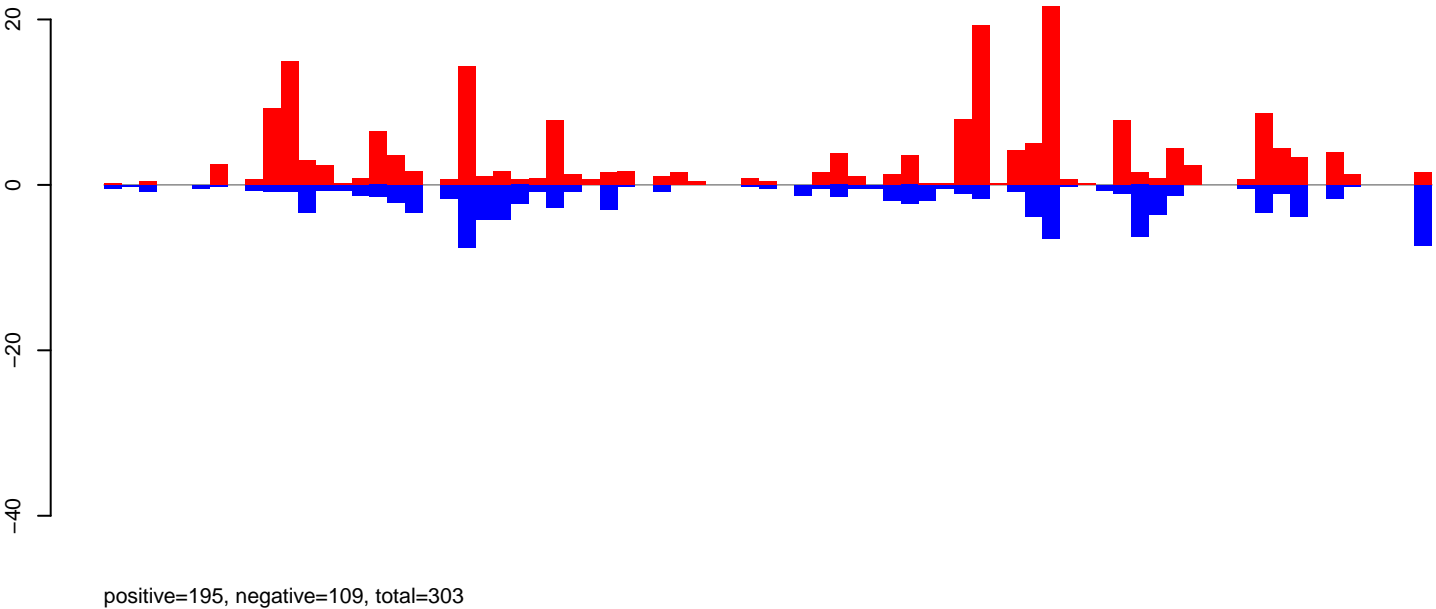
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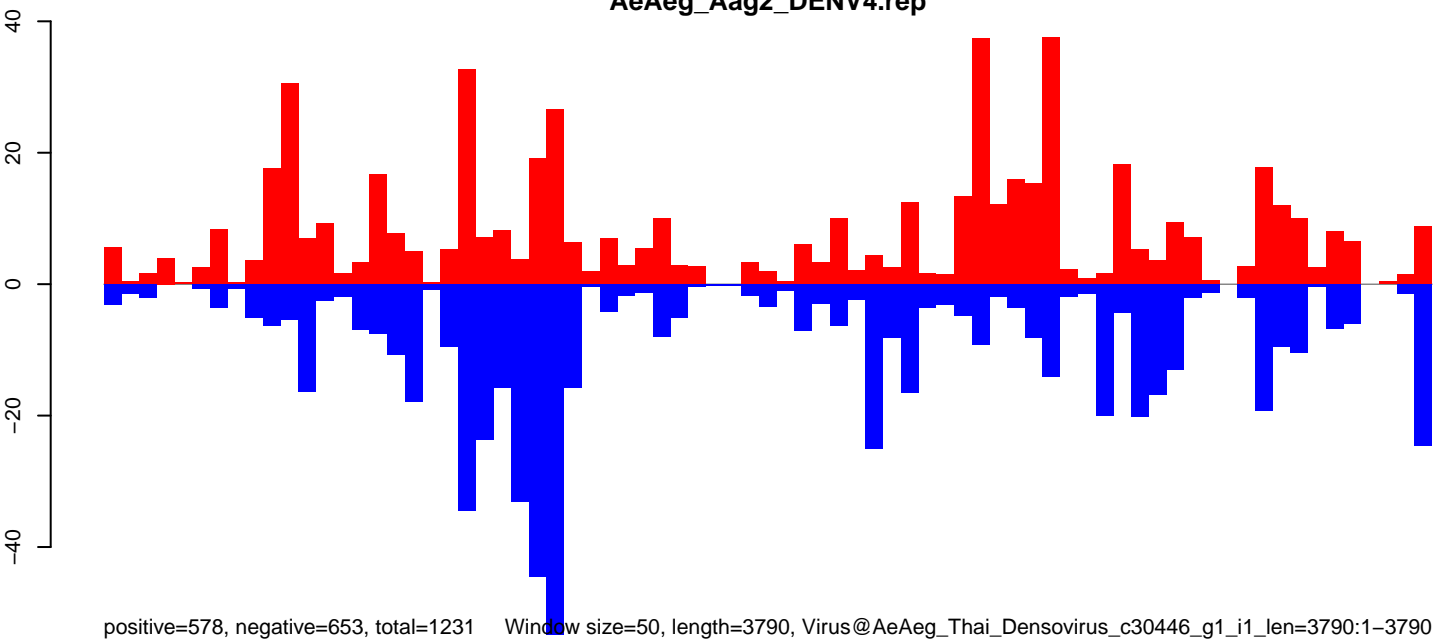
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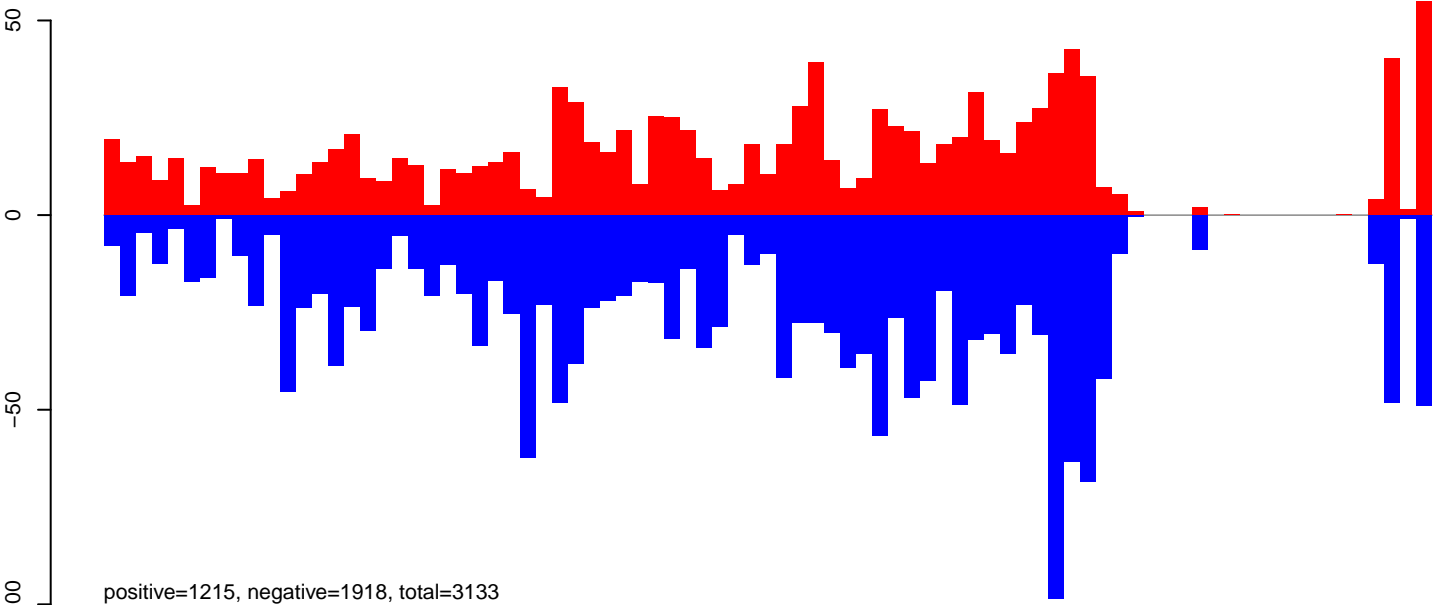
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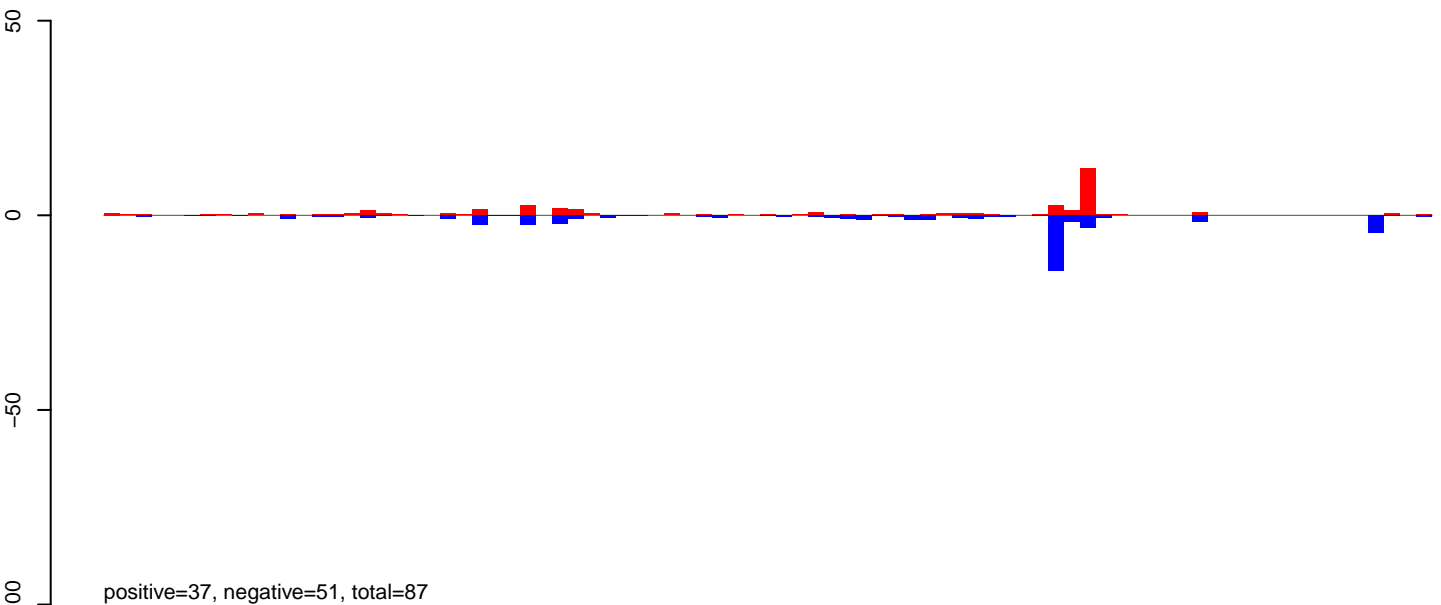
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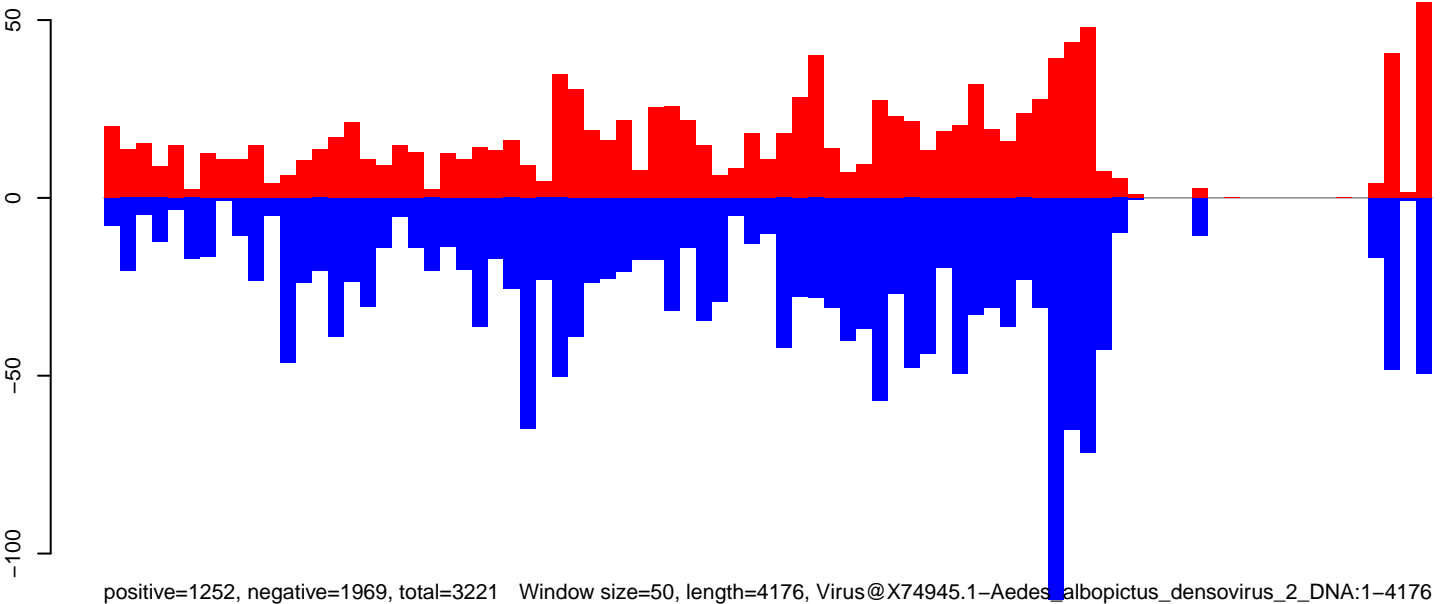
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

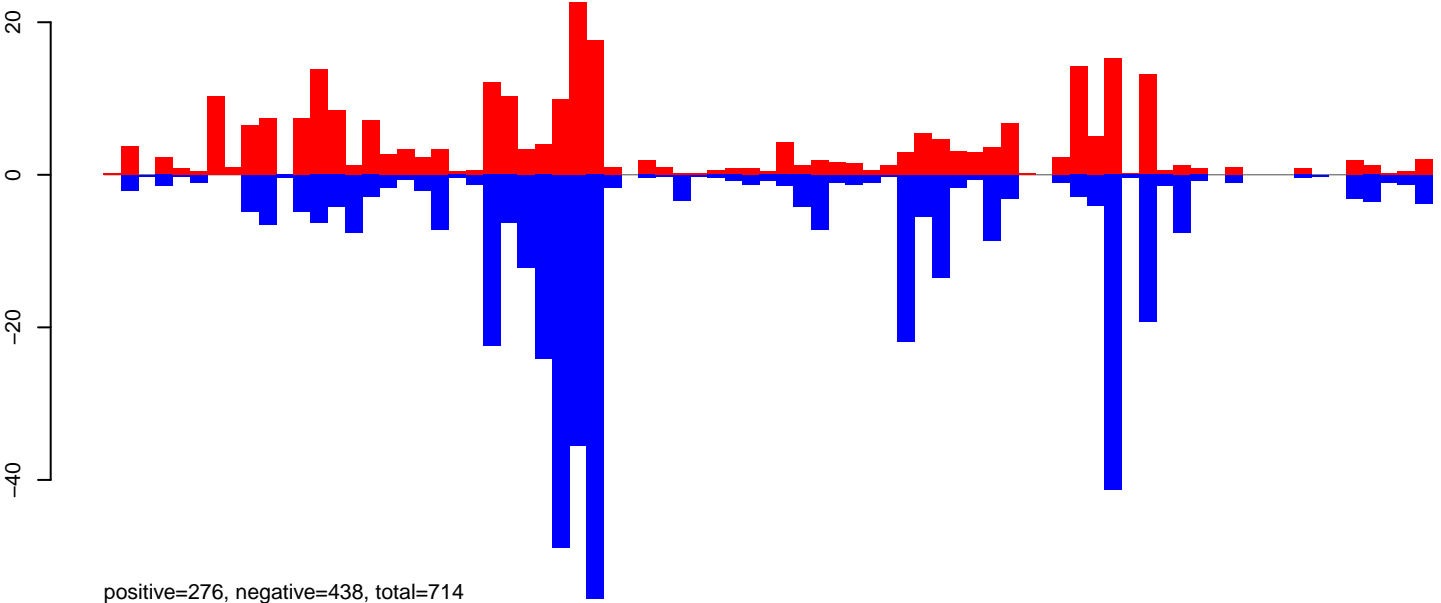


AeAeg_Aag2_DENV4.rep

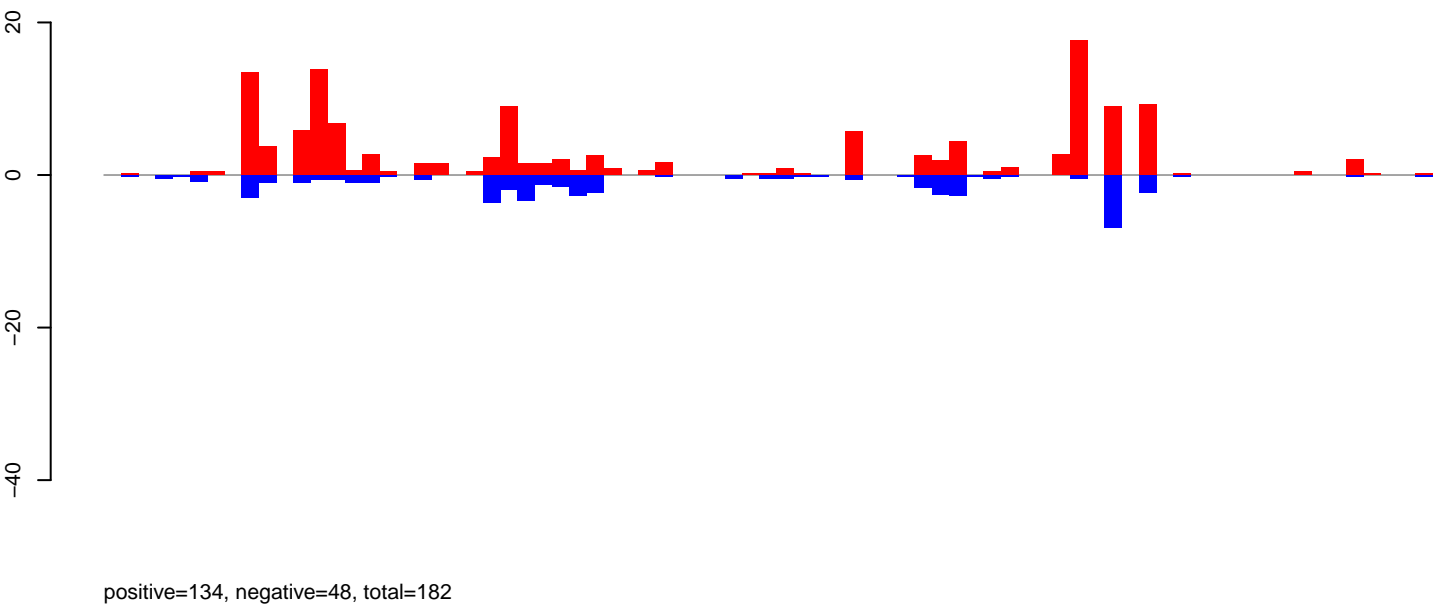


Window size=50, length=4176, Virus@X74945.1-Aedes albopictus_densovirus_2_DNA:1-4176

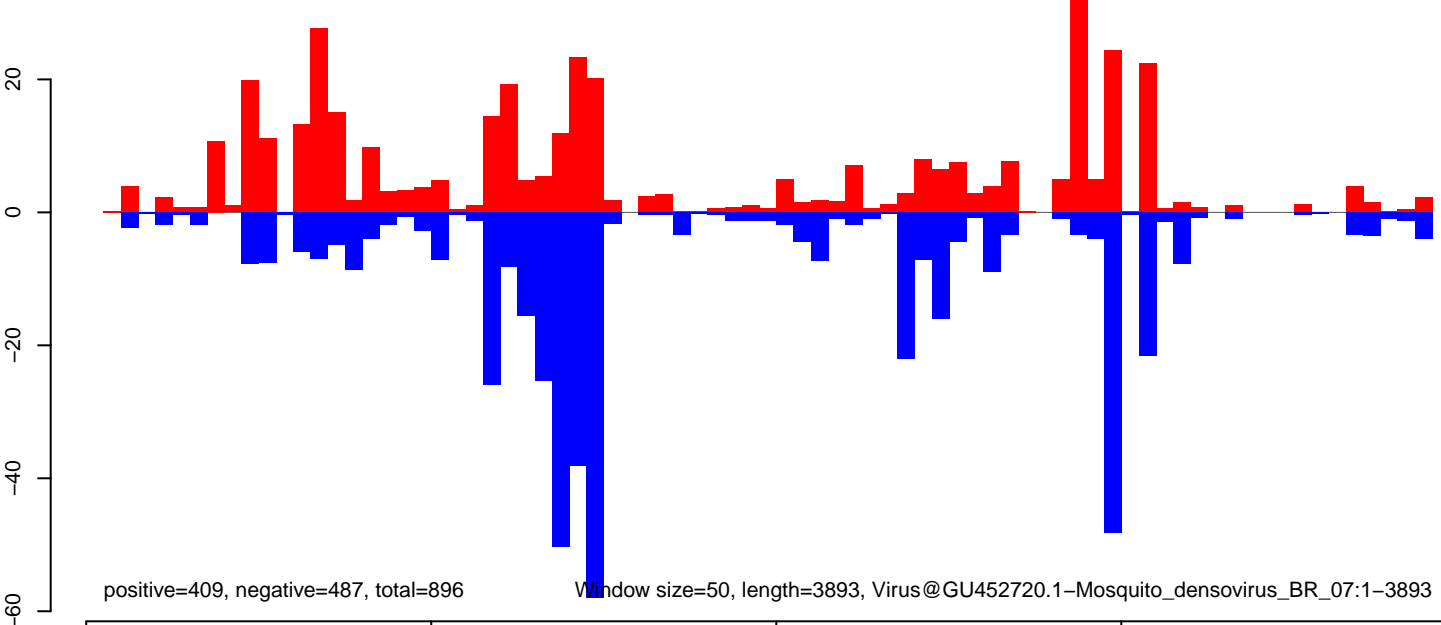
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

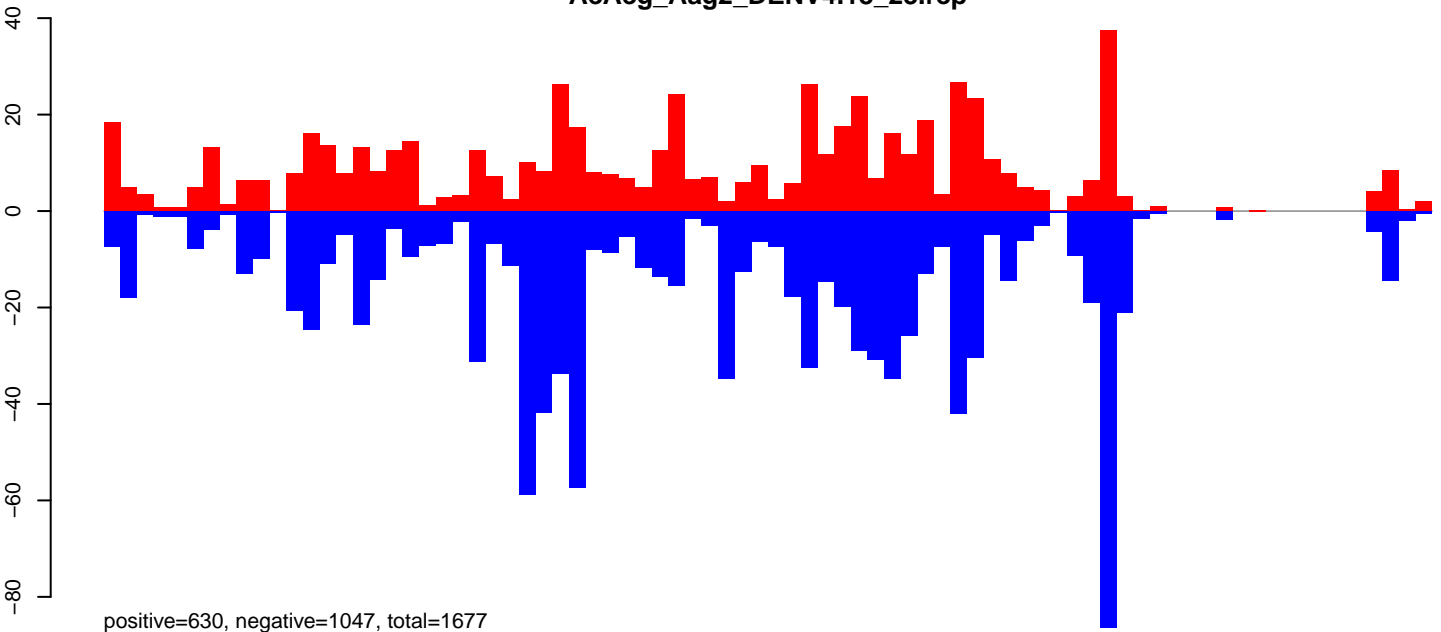


AeAeg_Aag2_DENV4.rep

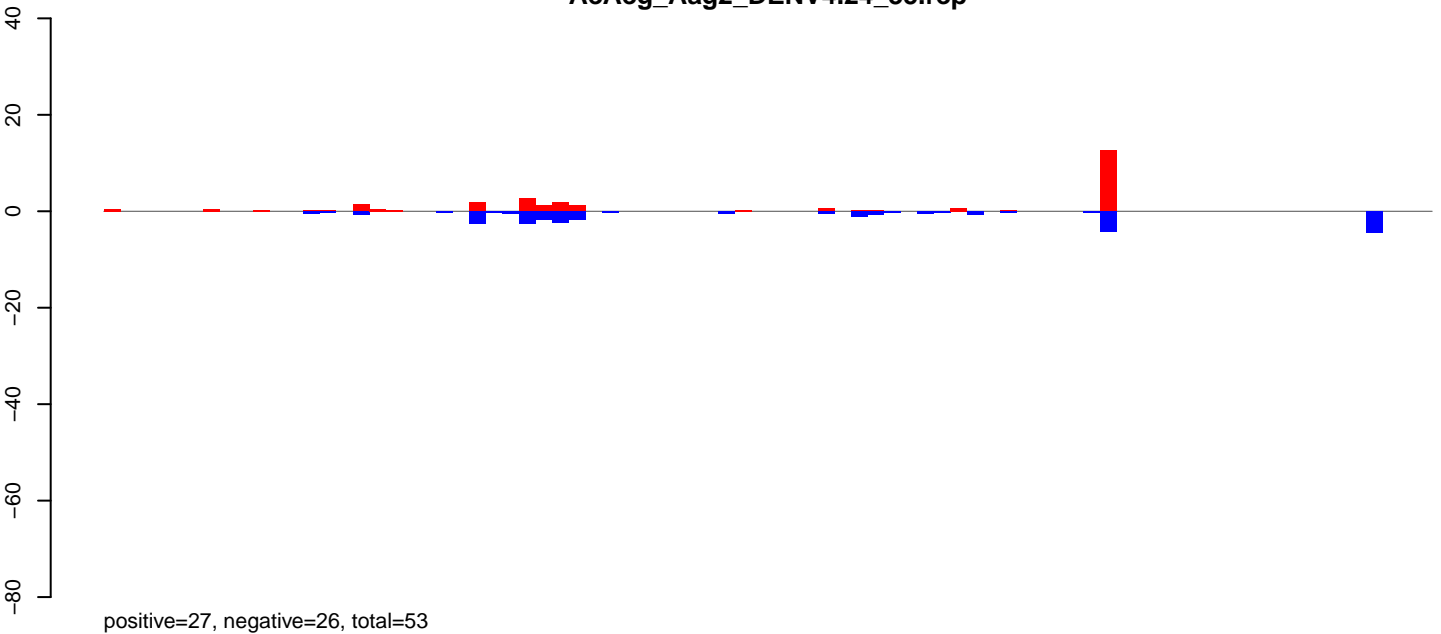


Window size=50, length=3893, Virus@GU452720.1-Mosquito_densovirus_BR_07:1-3893

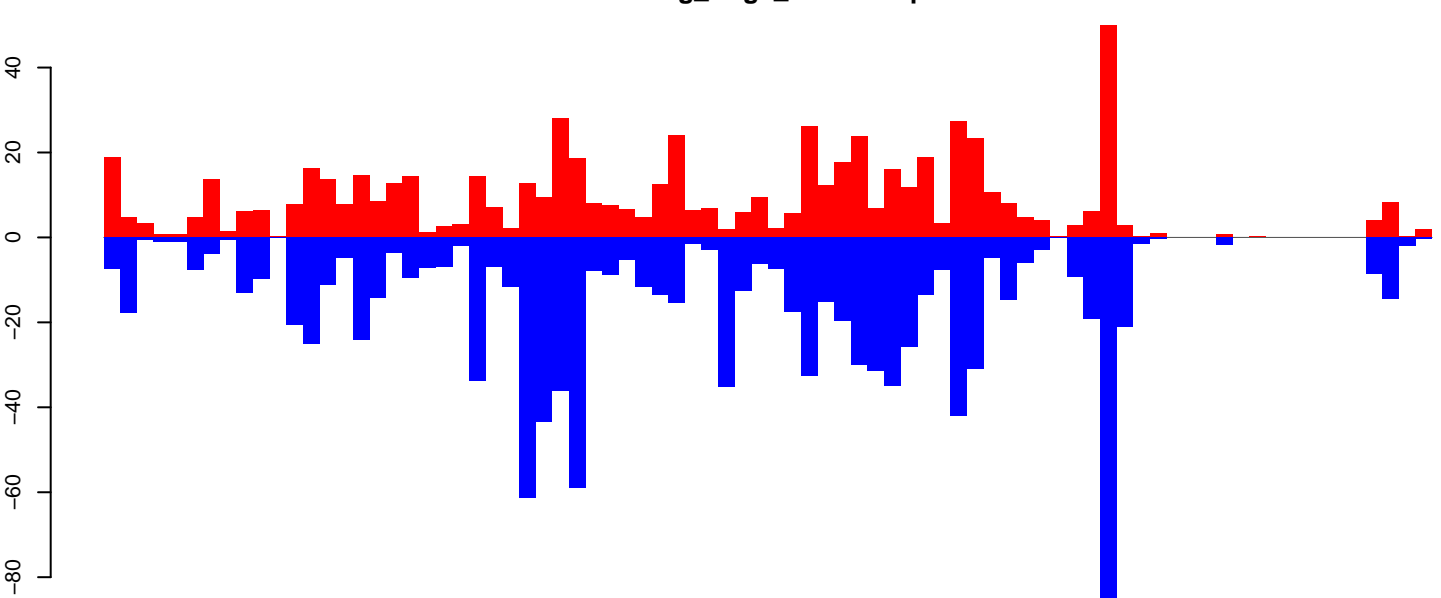
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



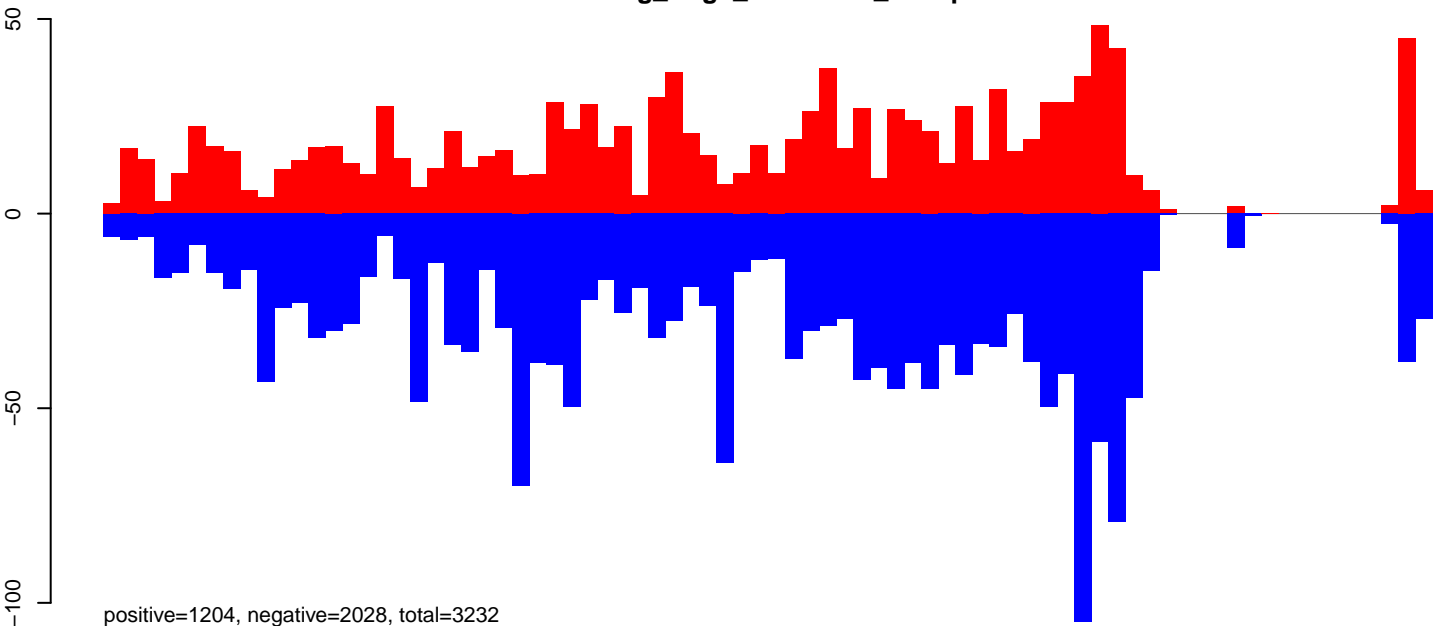
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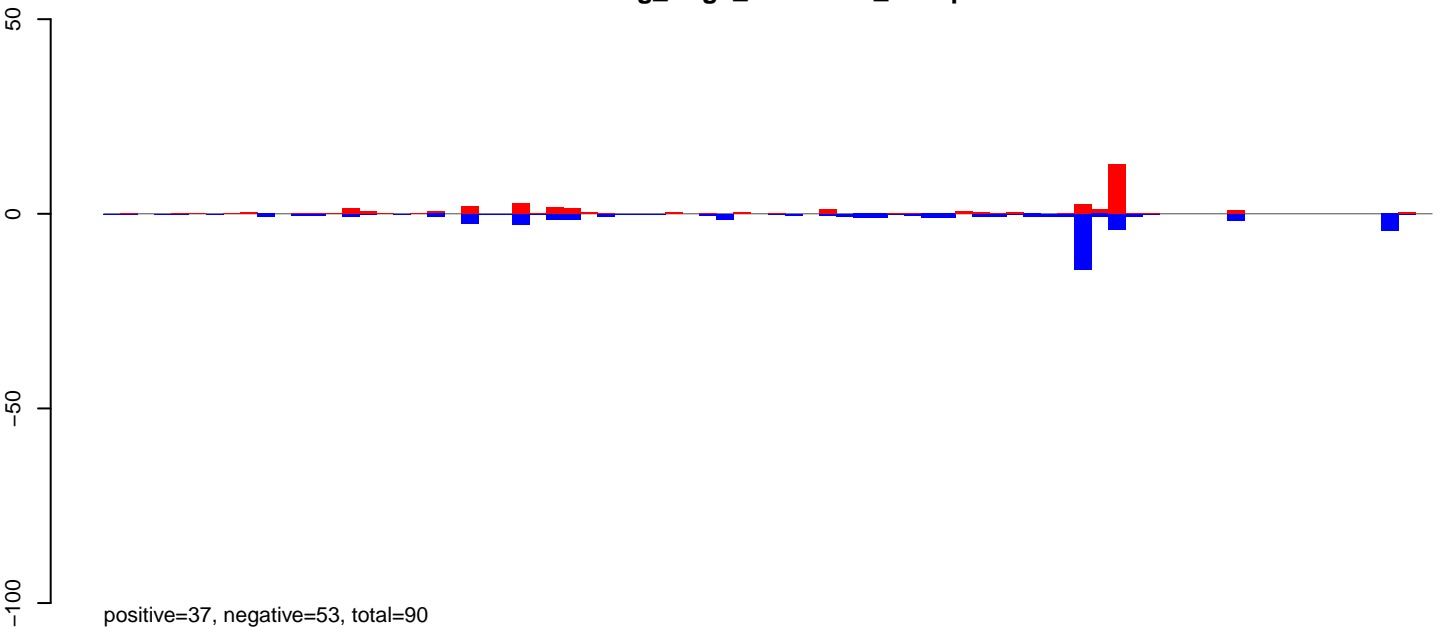
AY310874=557, segment1=1673, 006=1730, dengovirus_non-structural_protein_1_non-structural_protein_2_and_capsid_protein_genes:1-4006

0 1000 2000 3000 4000

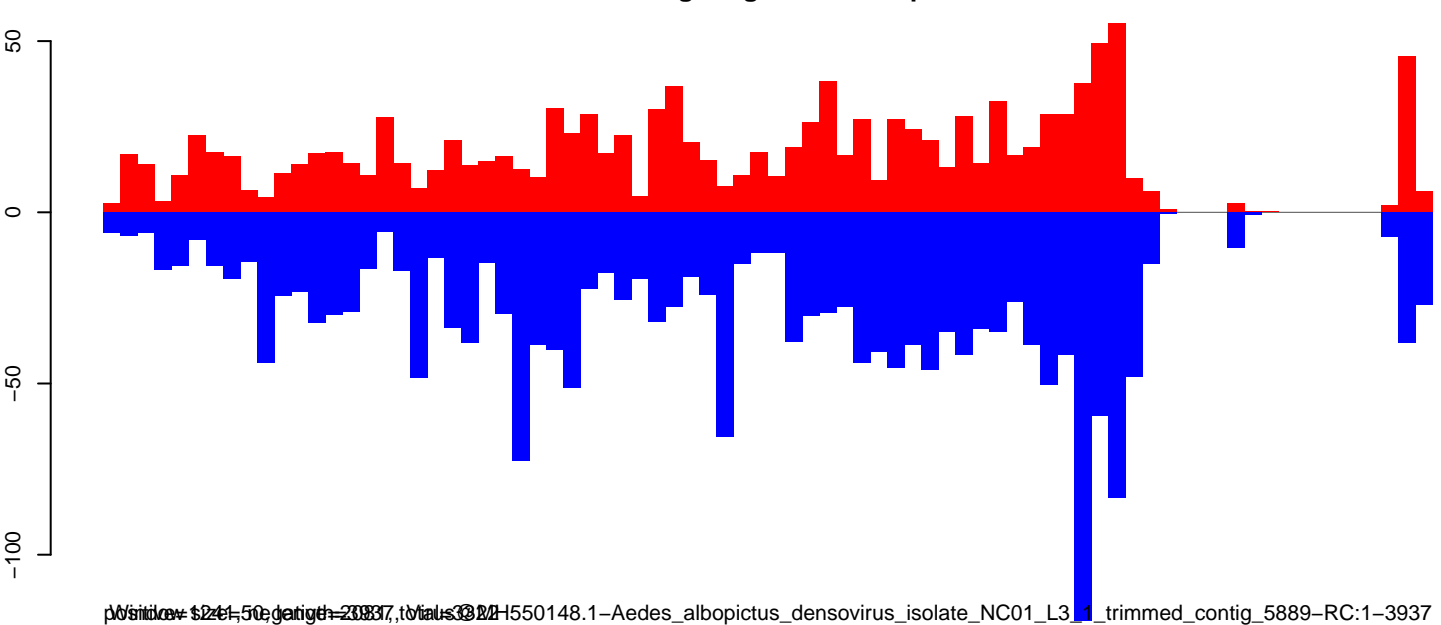
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



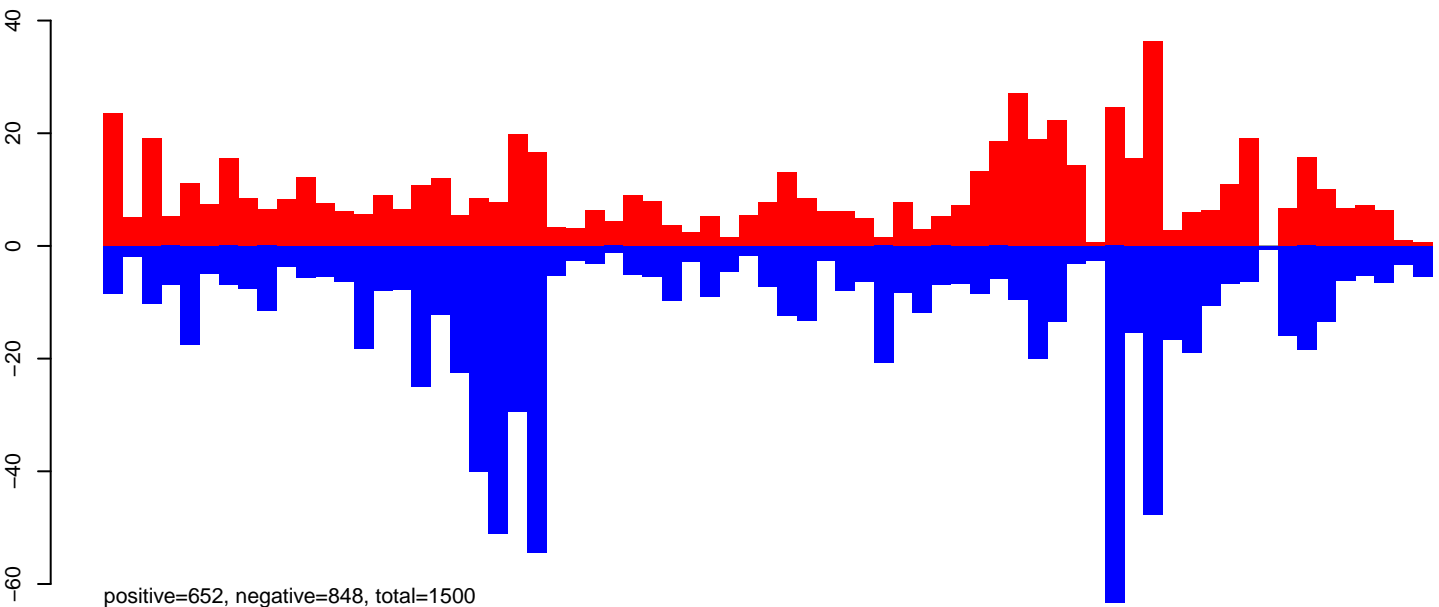
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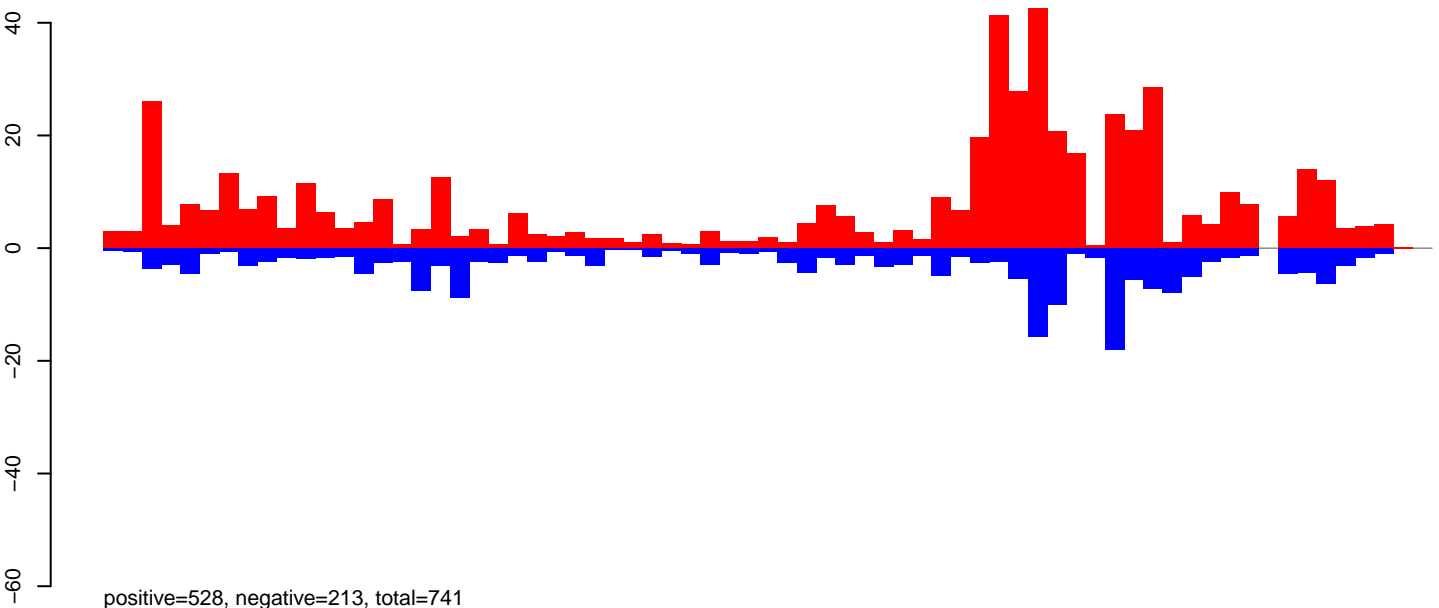
NC1550148.1-Aedes_albopictus_densovirus_isolate_NC01_L3_1_trimmed_contig_5889-RC:1-3937

0 1000 2000 3000 4000

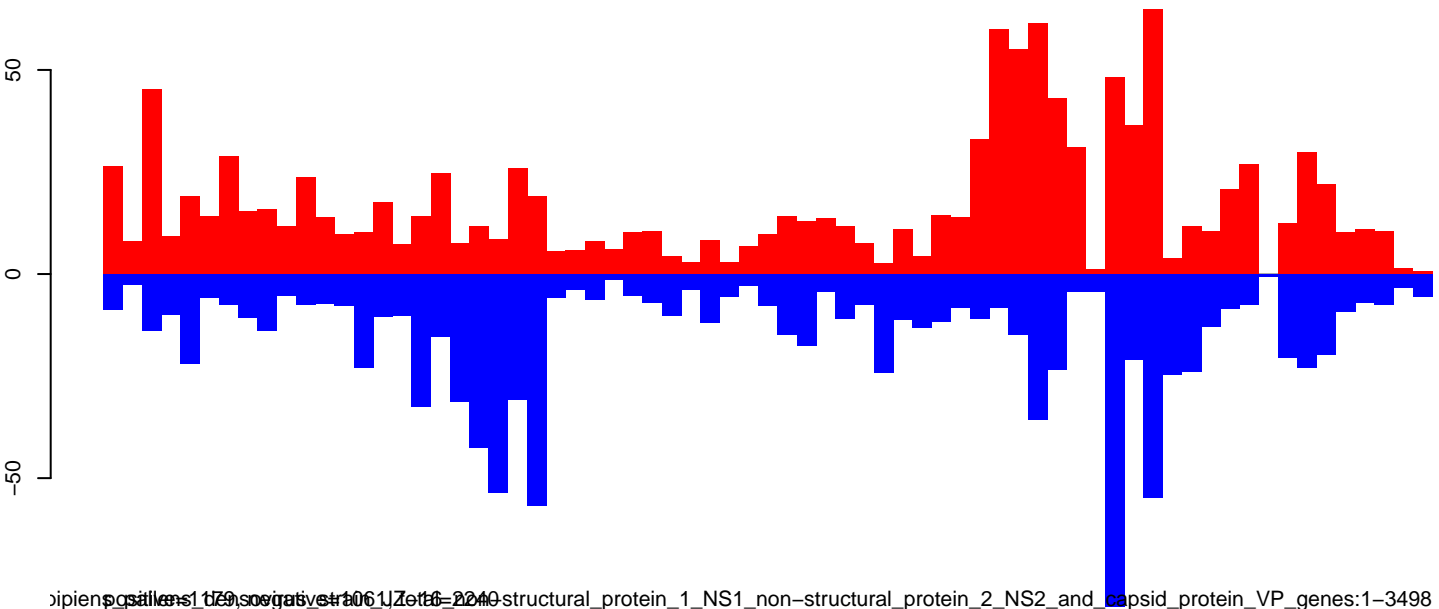
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



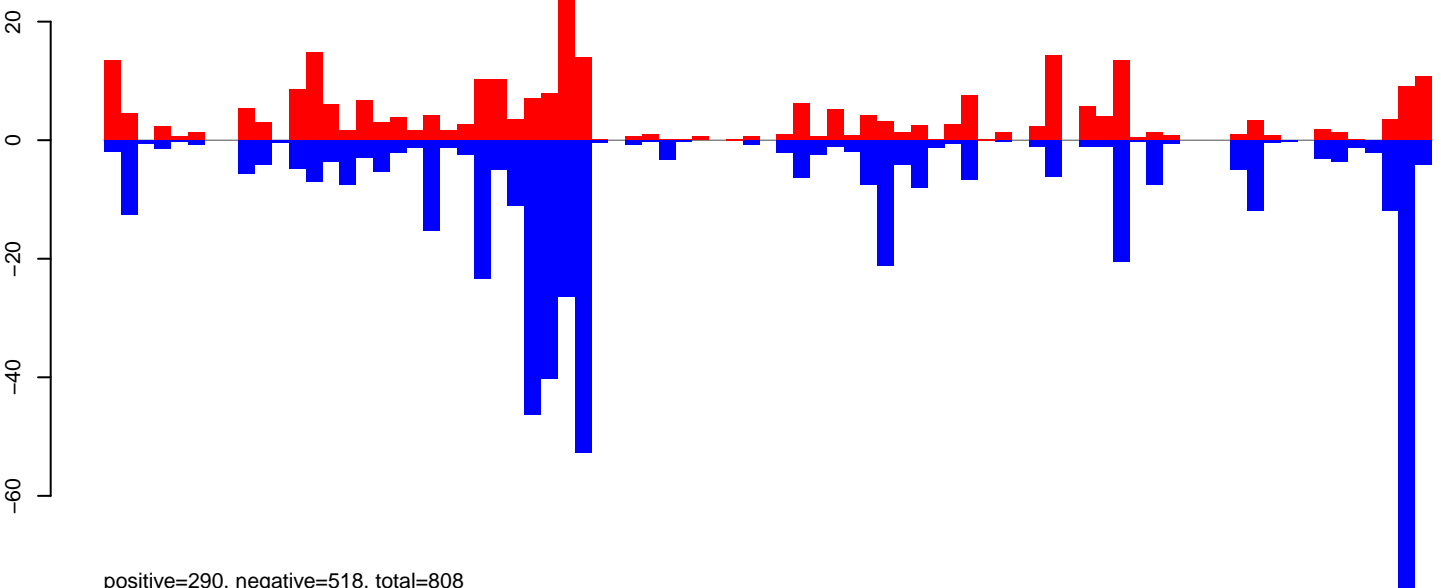
AeAeg_Aag2_DENV4.rep



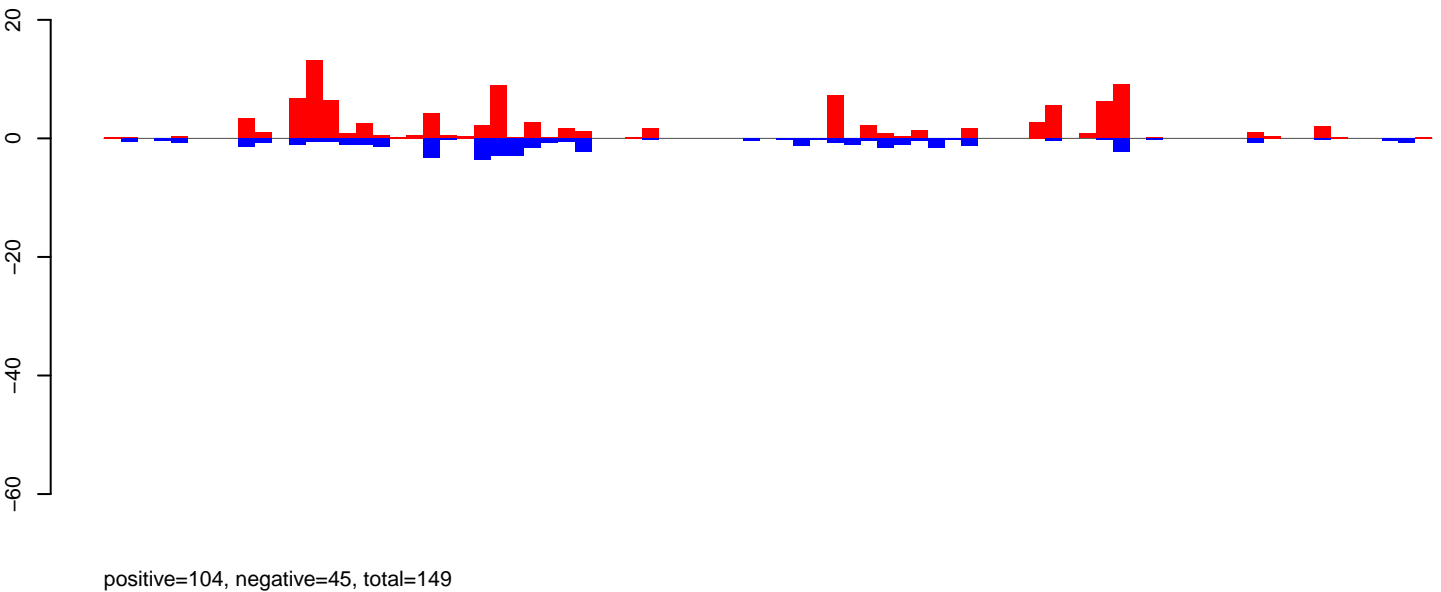
pienspiens1179, positive=416, total=1595 structural_protein_1_NS1_non-structural_protein_2_NS2_and_capsid_protein_VP_genes:1-3498

0 500 1000 1500 2000 2500 3000 3500

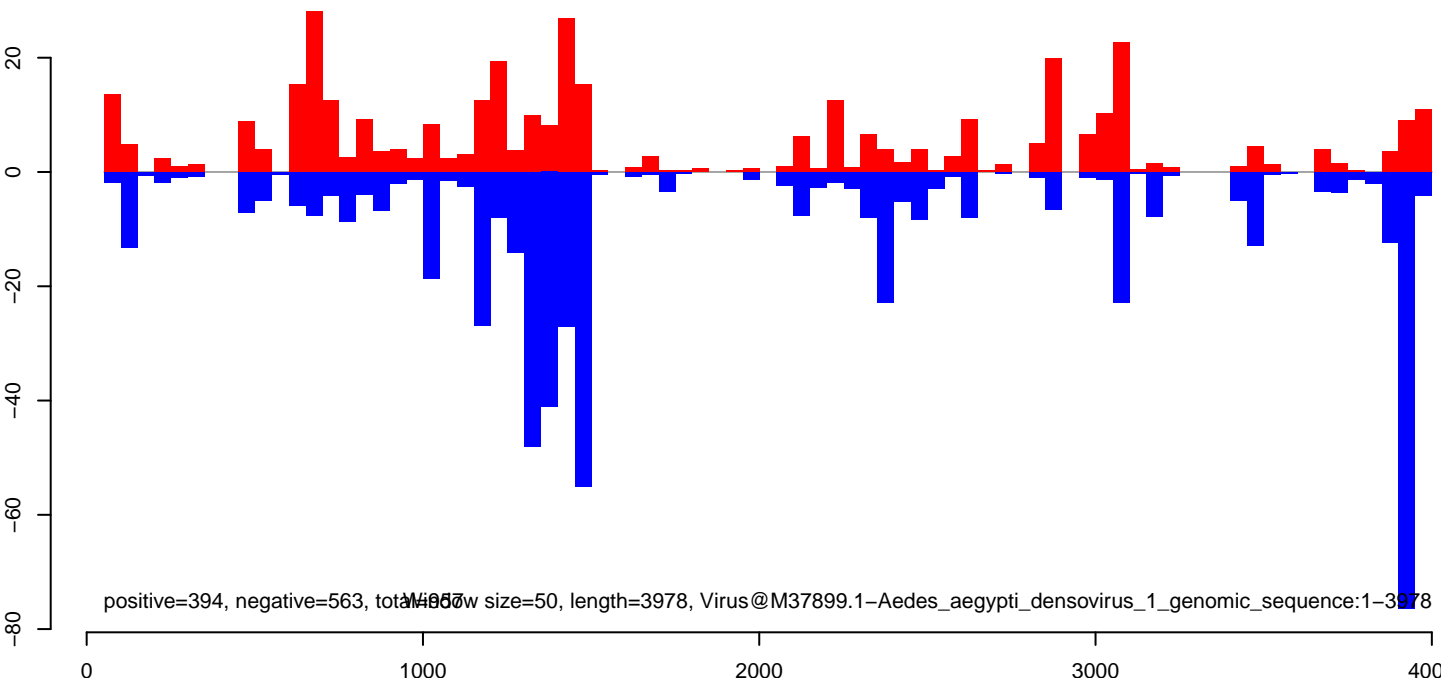
AeAeg_Aag2_DENV4.18_23.rep



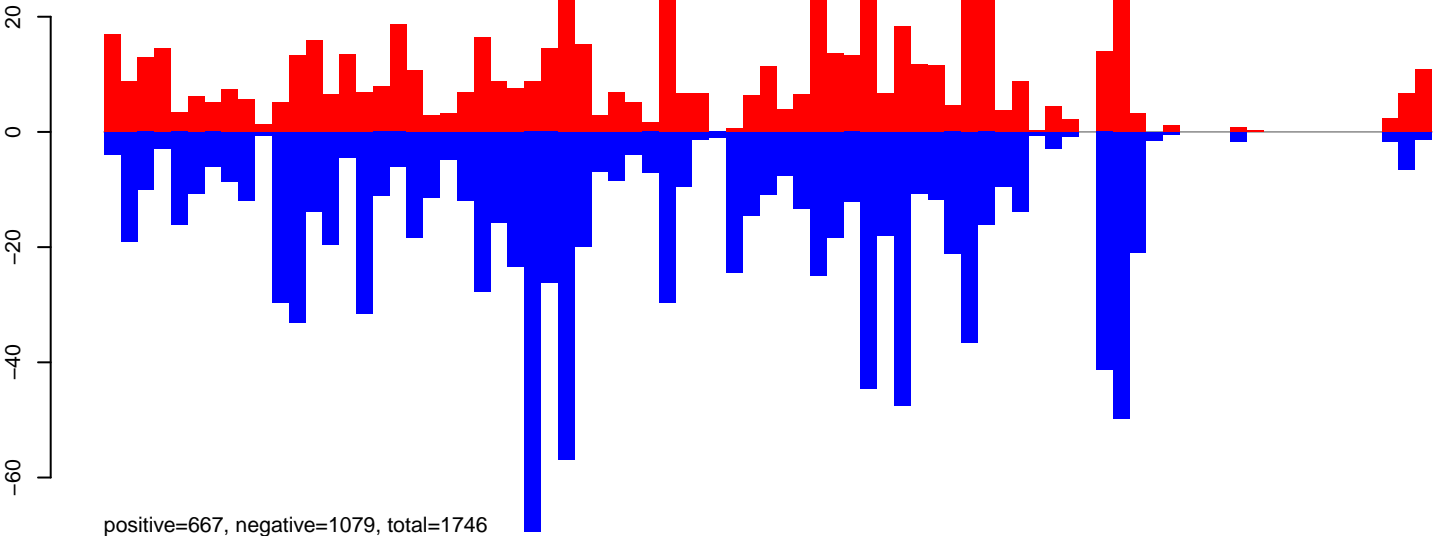
AeAeg_Aag2_DENV4.24_35.rep



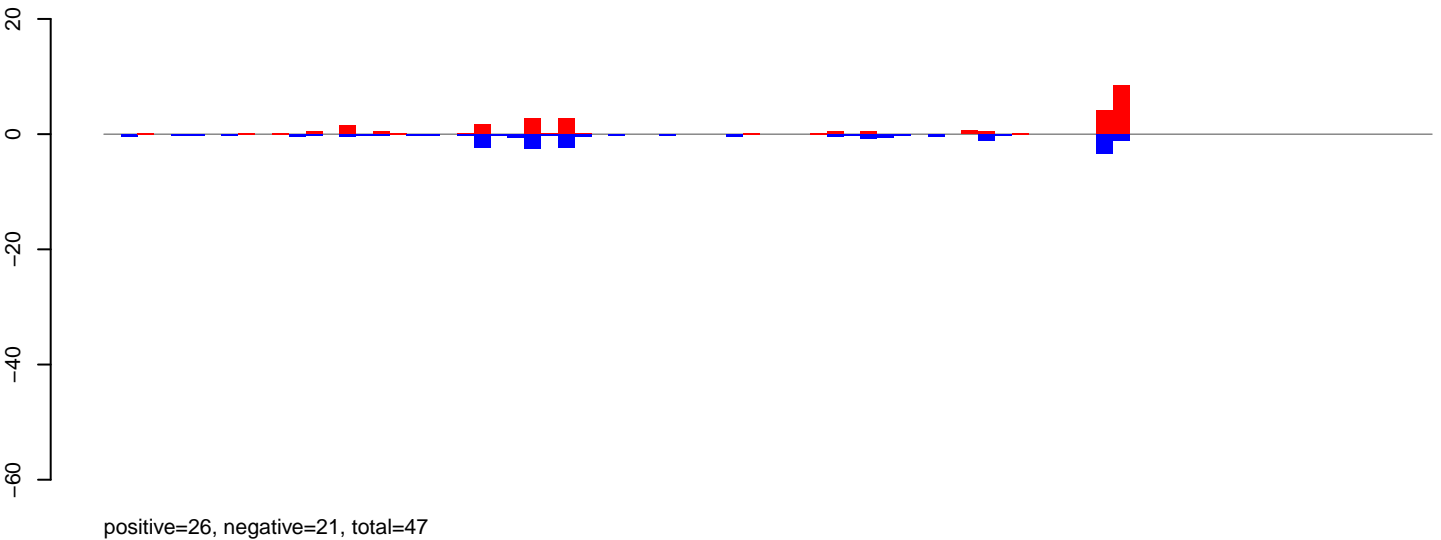
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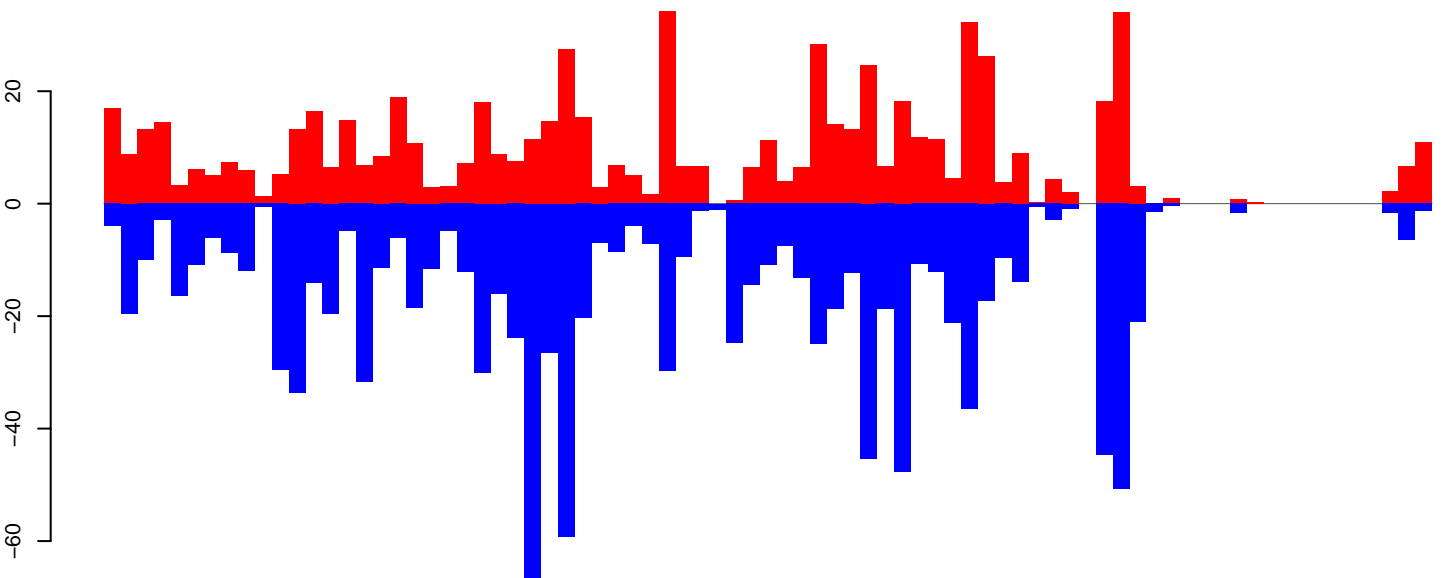
AeAeg_Aag2_DENV4.18_23.rep



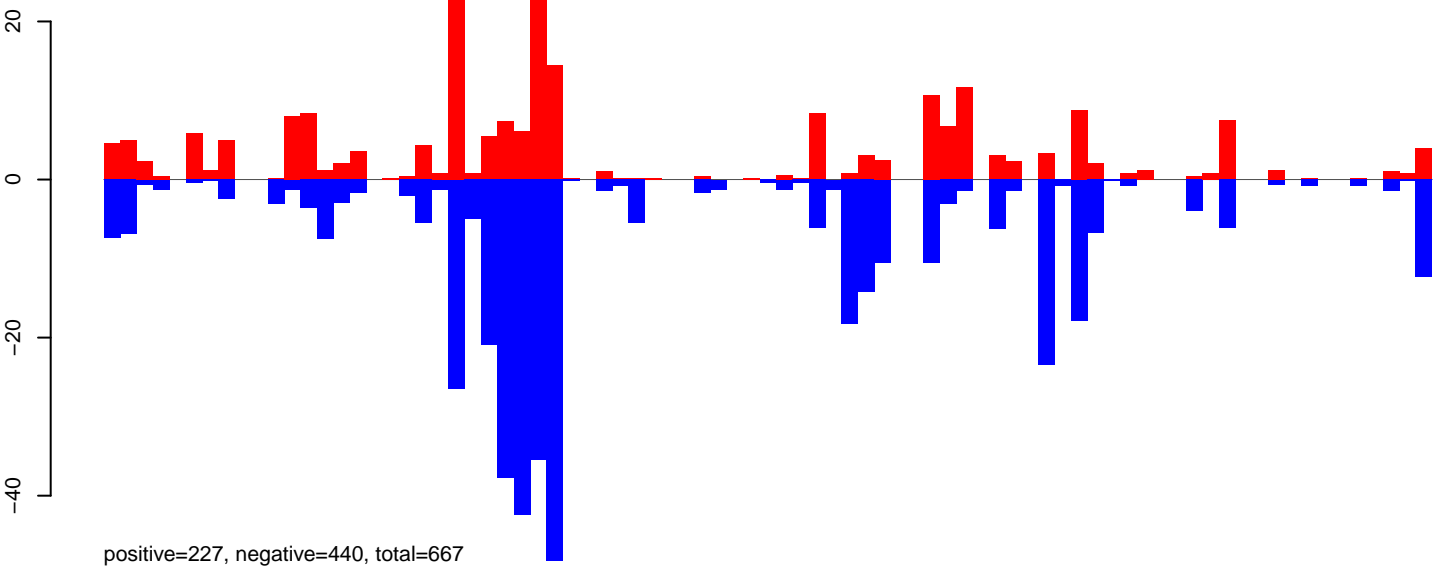
AeAeg_Aag2_DENV4.24_35.rep



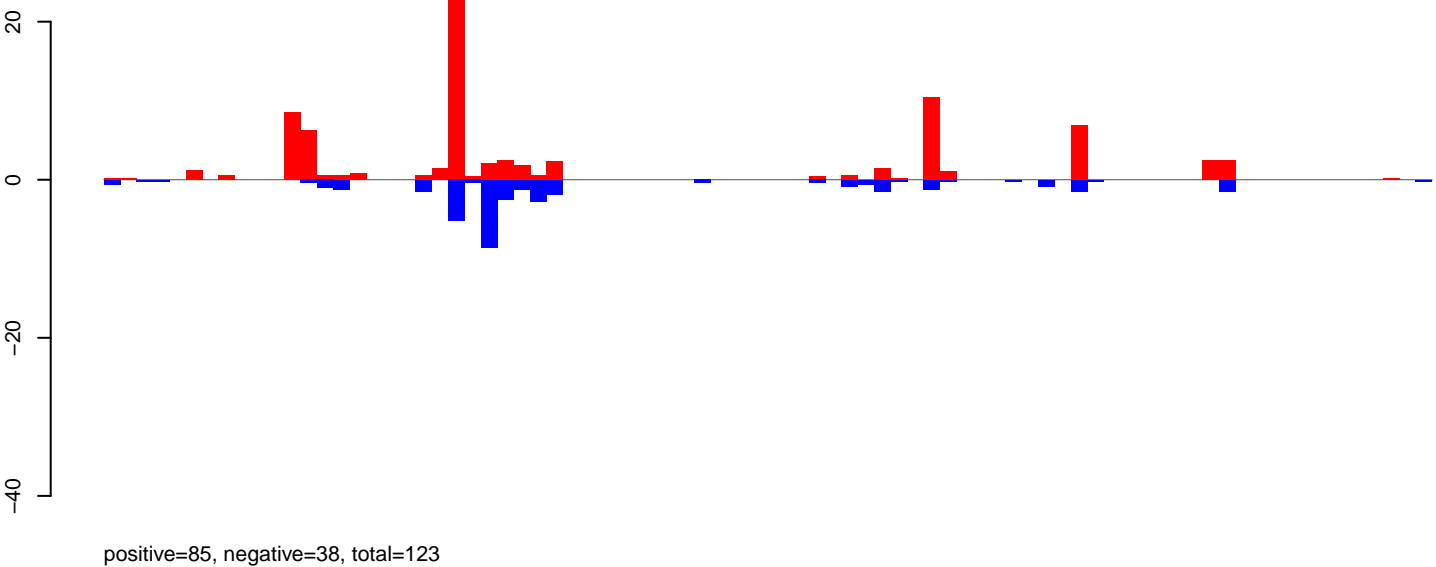
AeAeg_Aag2_DENV4.rep



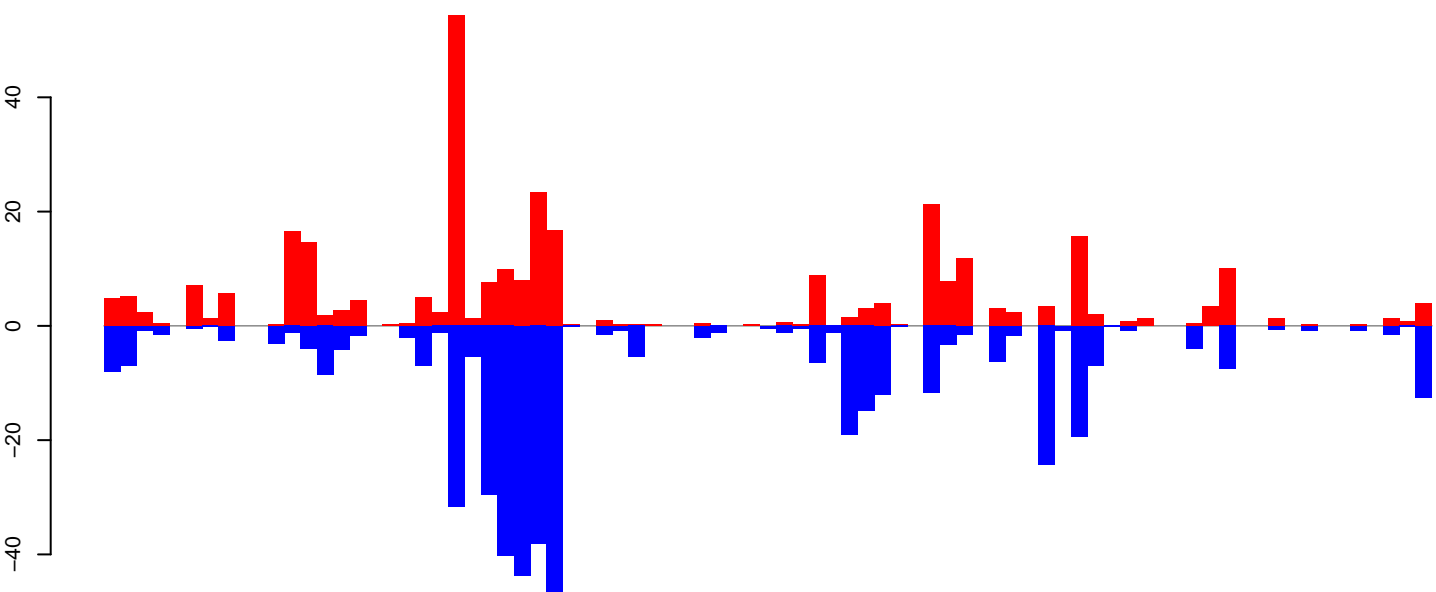
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



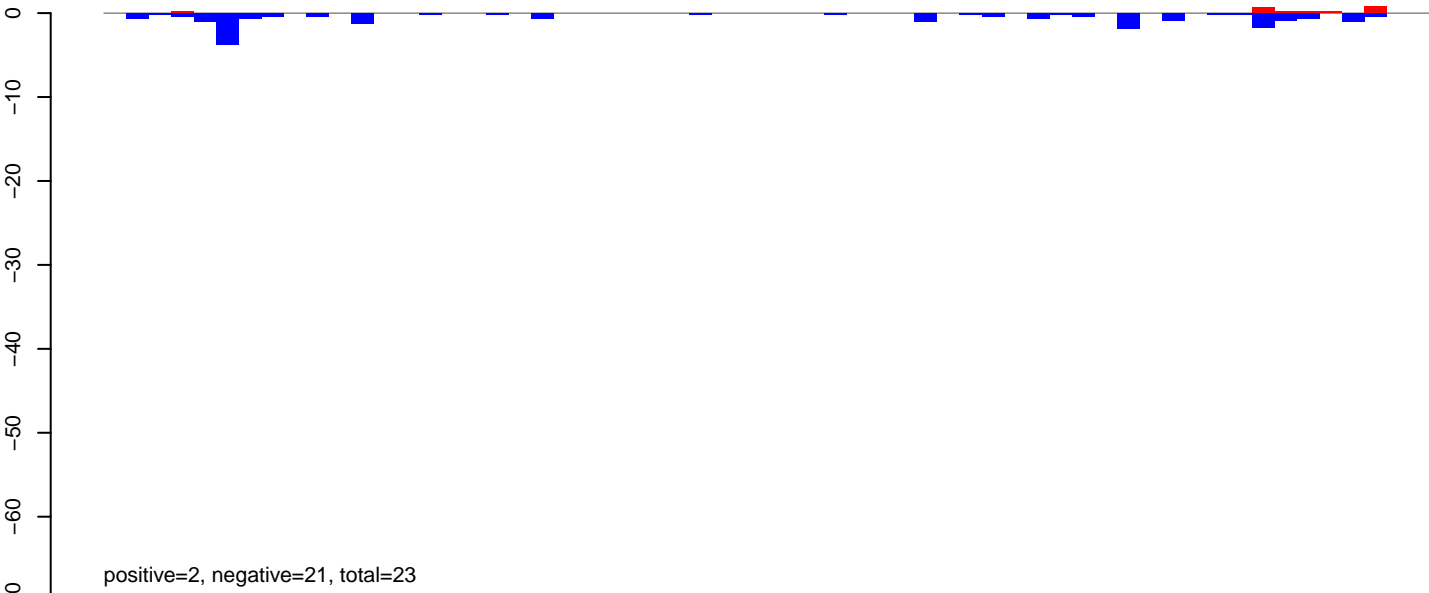
AeAeg_Aag2_DENV4.rep



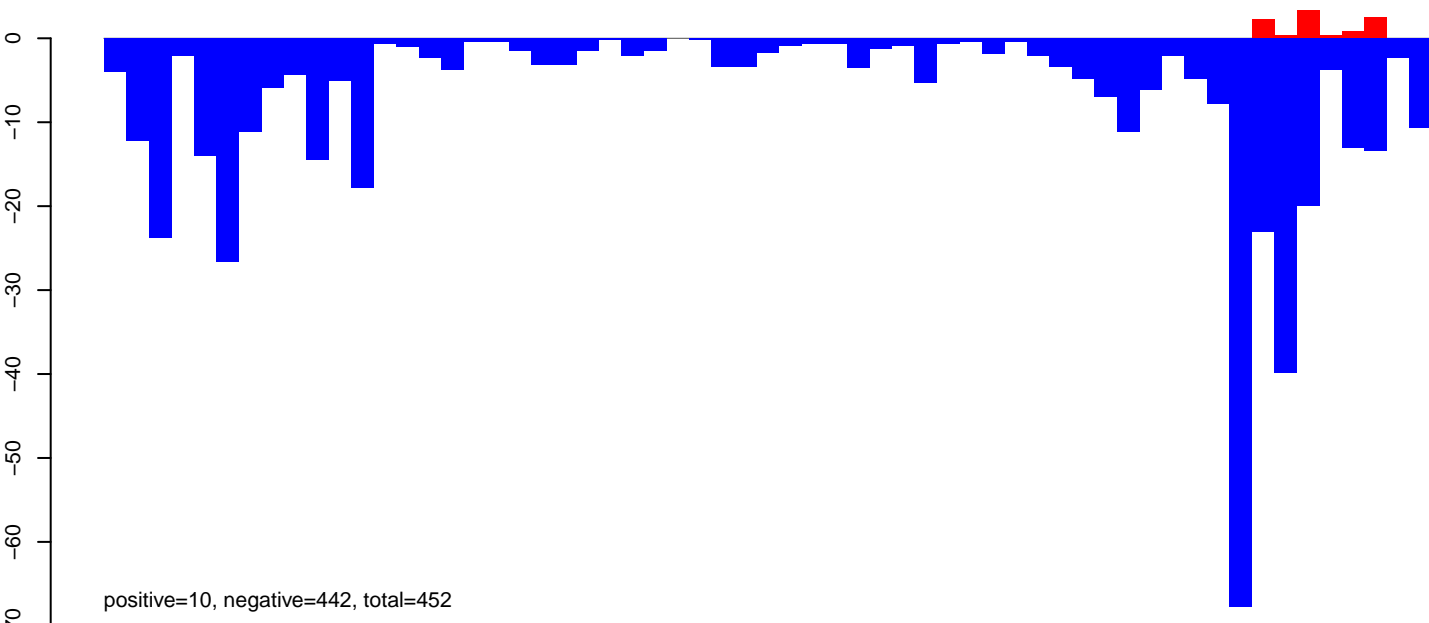
les_alp..._332_86..._478..._760-structural_protein_1_NS-1_non-structural_protein_2_NS-2_and_capsid_protein_VP_genes:1-4094

0 1000 2000 3000 4000

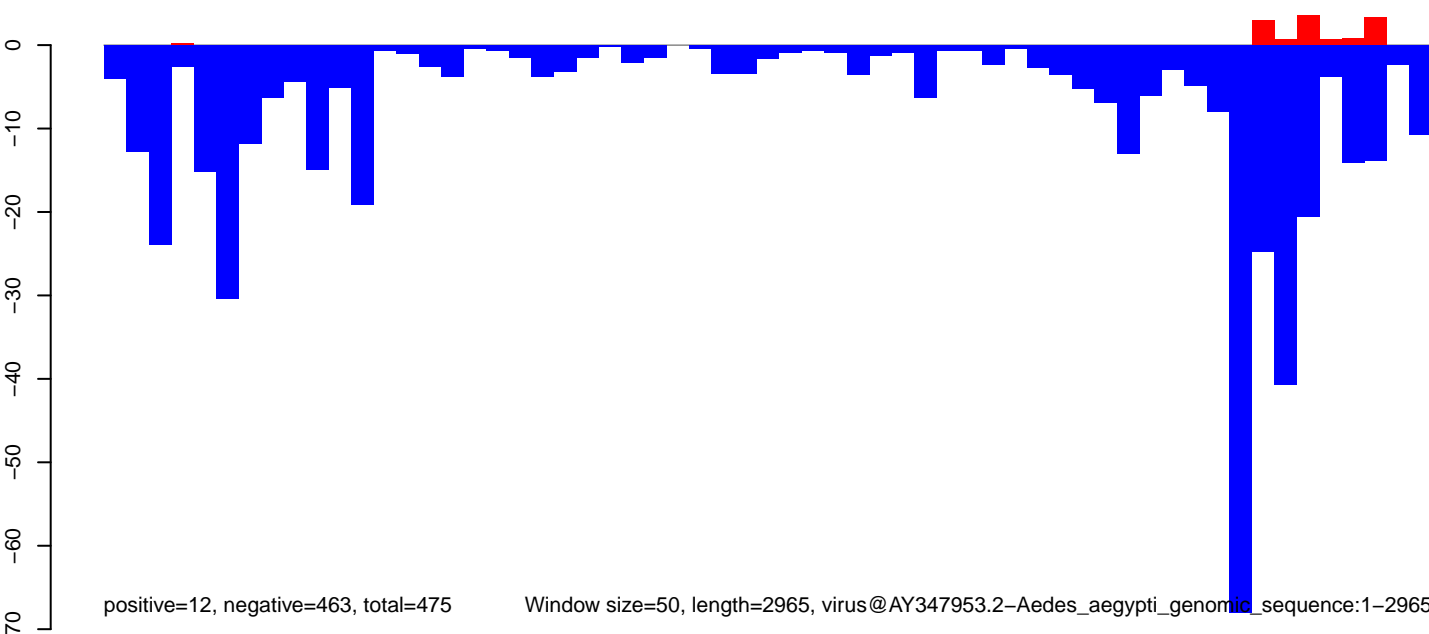
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

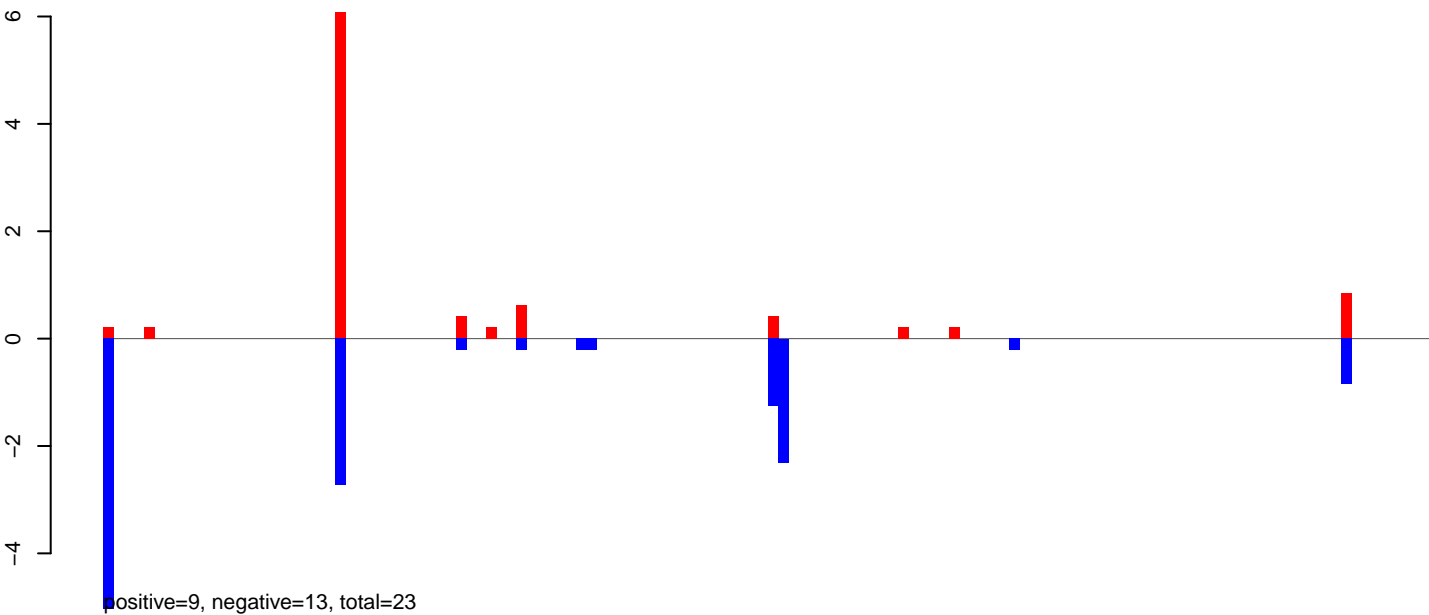


AeAeg_Aag2_DENV4.rep

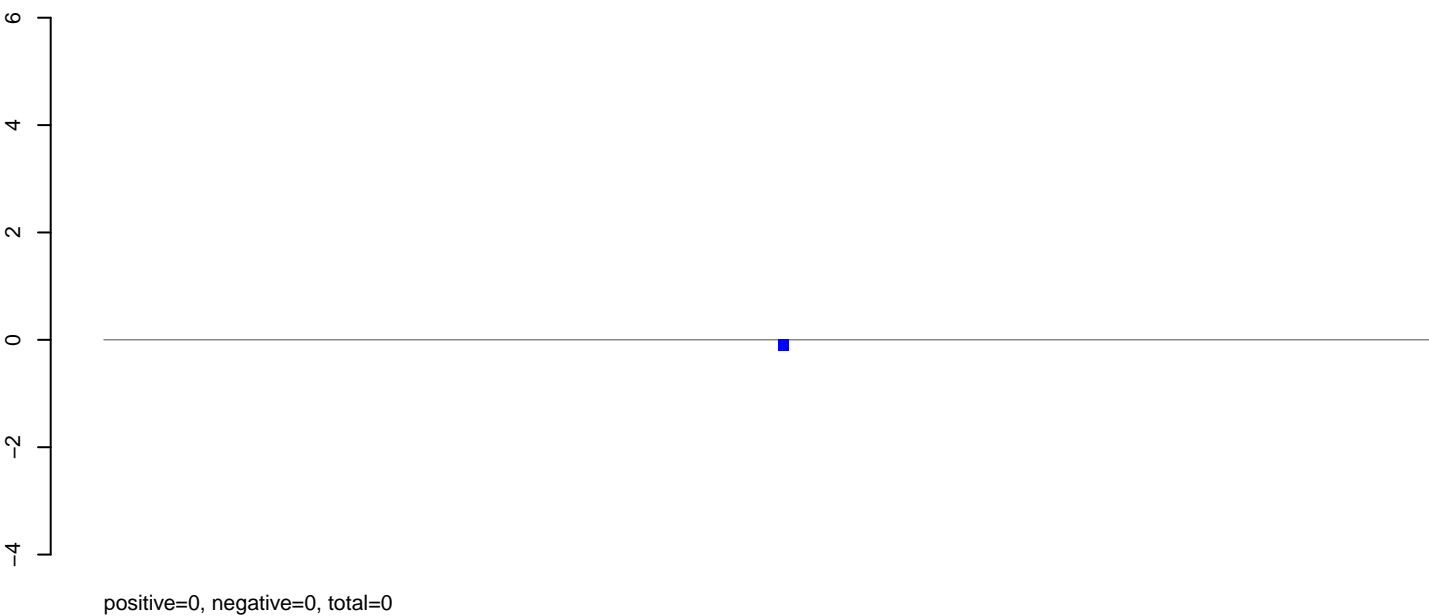


Window size=50, length=2965, virus@AY347953.2-Aedes_aegypti_genomic_sequence:1-2965

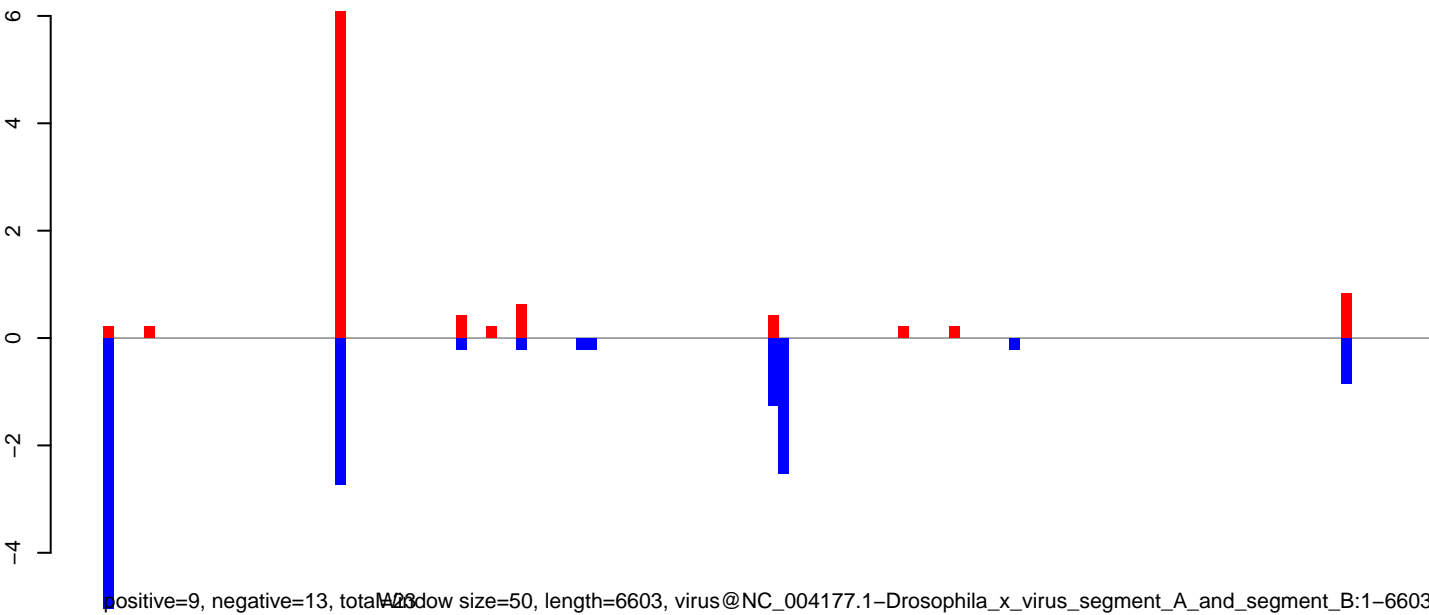
AeAeg_Aag2_DENV4.18_23.rep



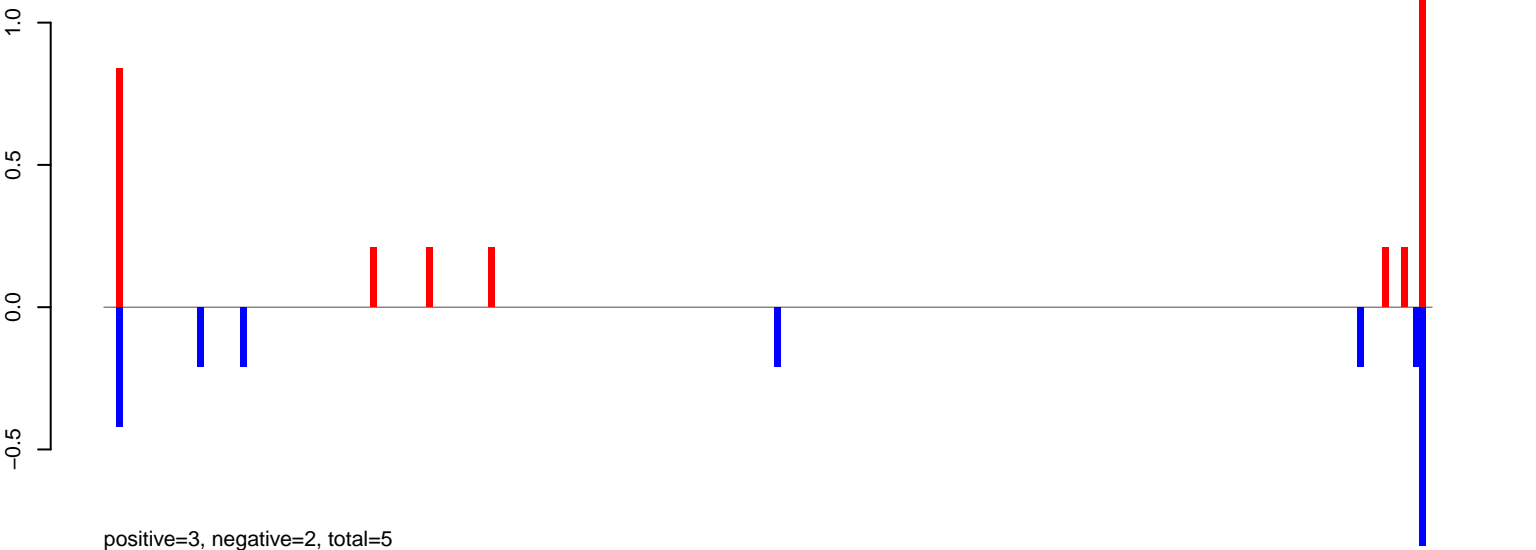
AeAeg_Aag2_DENV4.24_35.rep



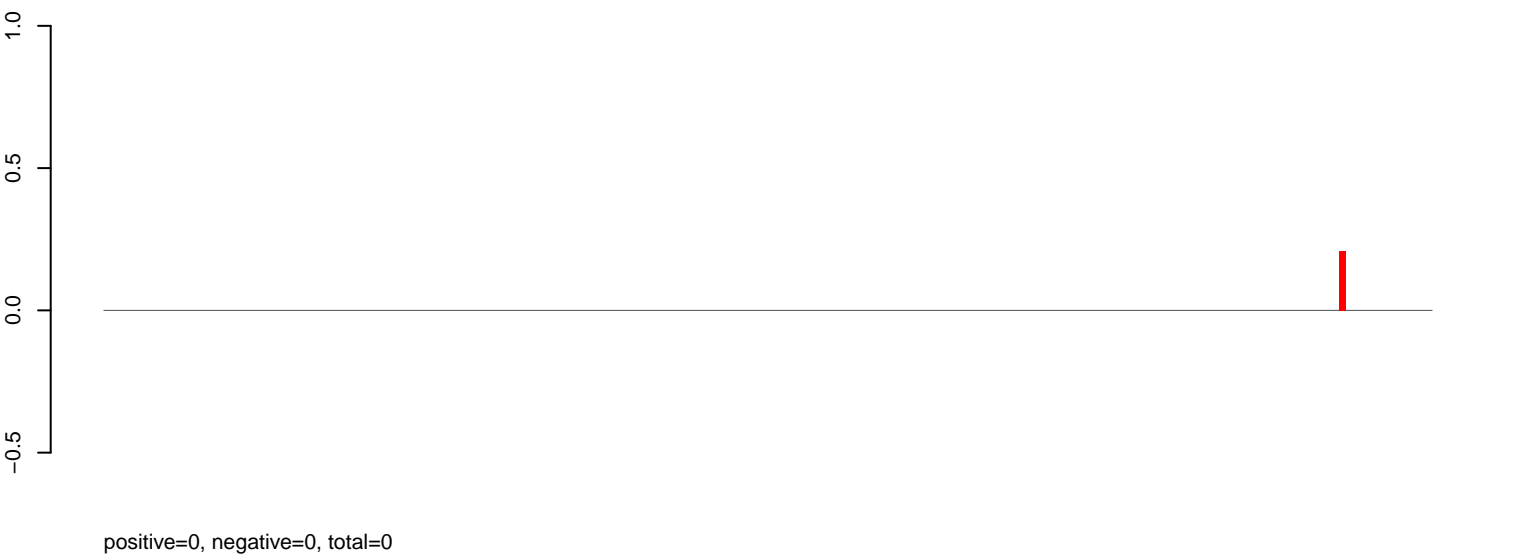
AeAeg_Aag2_DENV4.rep



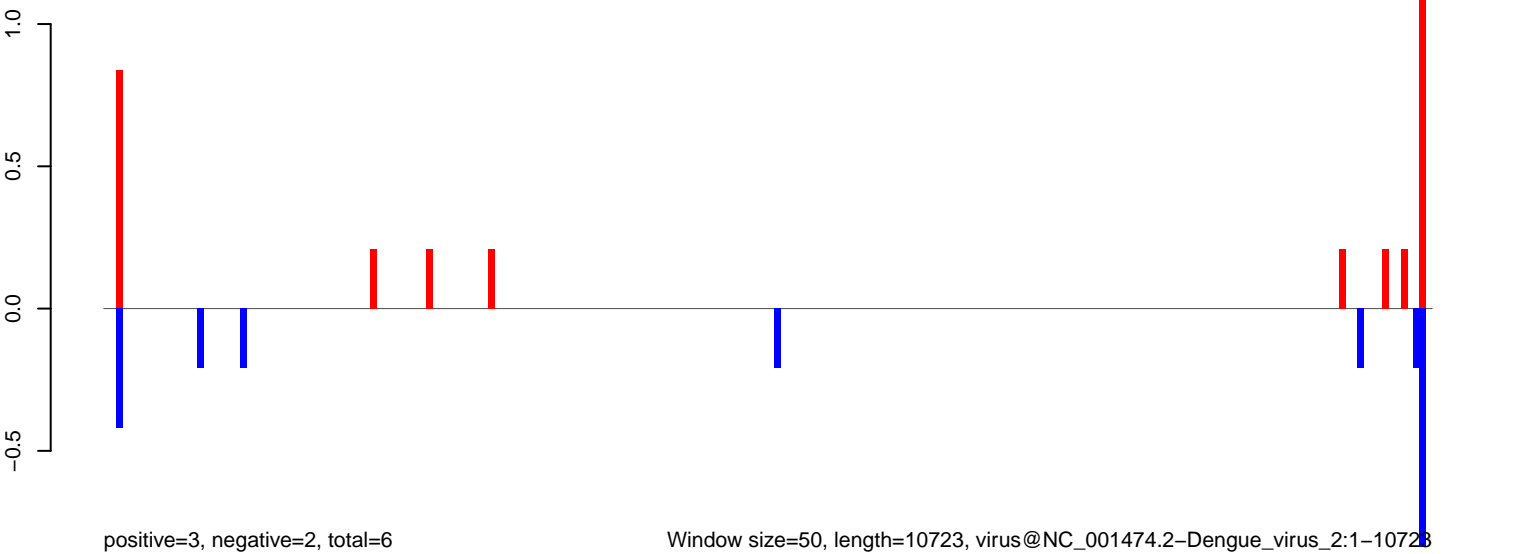
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

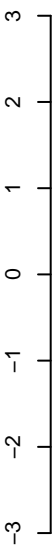


AeAeg_Aag2_DENV4.rep



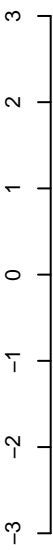
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AeAeg_Aag2_DENV4.18_23.rep



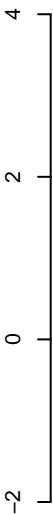
positive=10, negative=14, total=24

AeAeg_Aag2_DENV4.24_35.rep



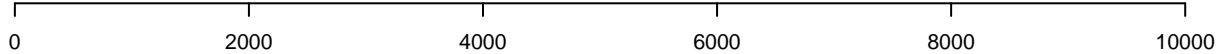
positive=7, negative=1, total=8

AeAeg_Aag2_DENV4.rep



positive=17, negative=15, total=32

Window size=50, length=11375, virus@NC_005064.1-Kamiti_River_virus:1-11375



AeAeg_Aag2_DENV4.18_23.rep

30
20
10
0
-10
-20

positive=63, negative=28, total=90

AeAeg_Aag2_DENV4.24_35.rep

30
20
10
0
-10
-20

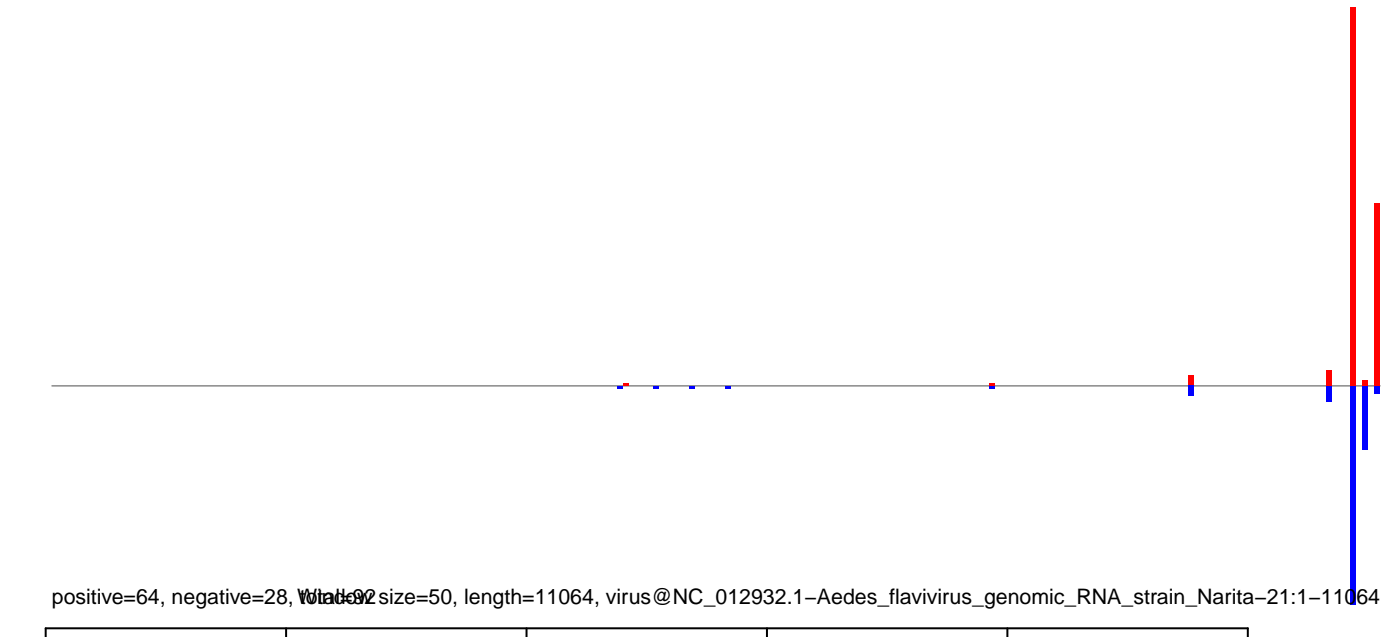
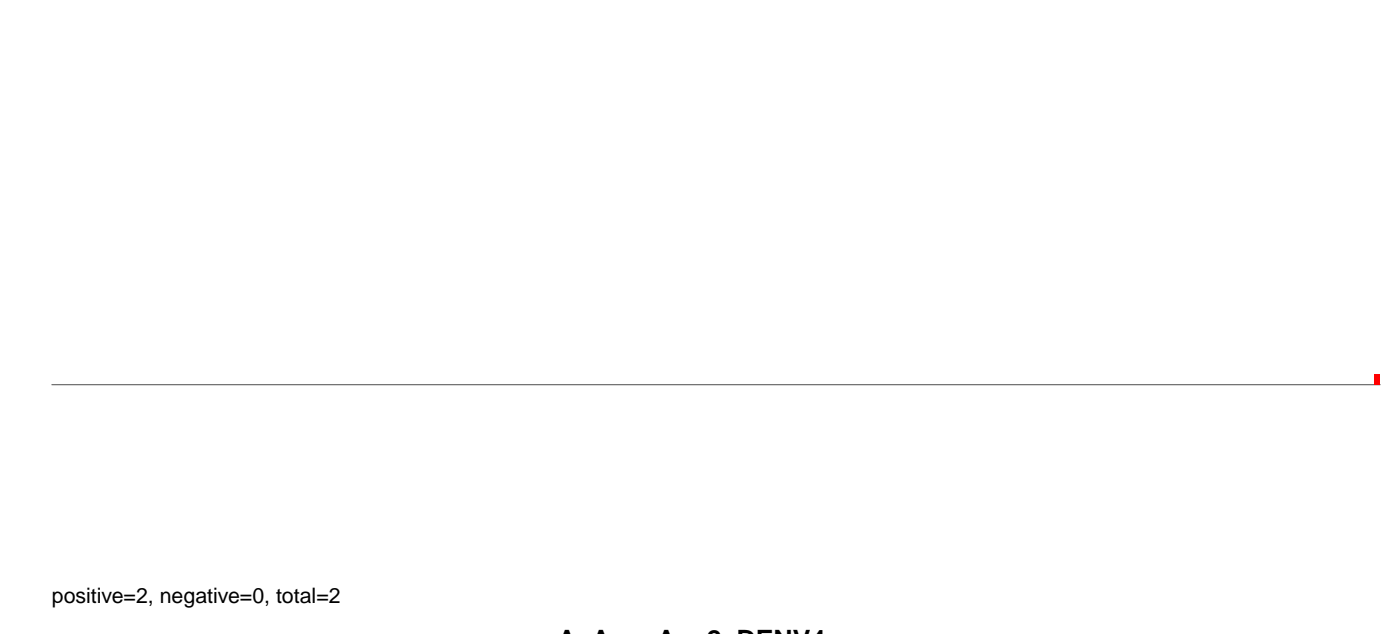
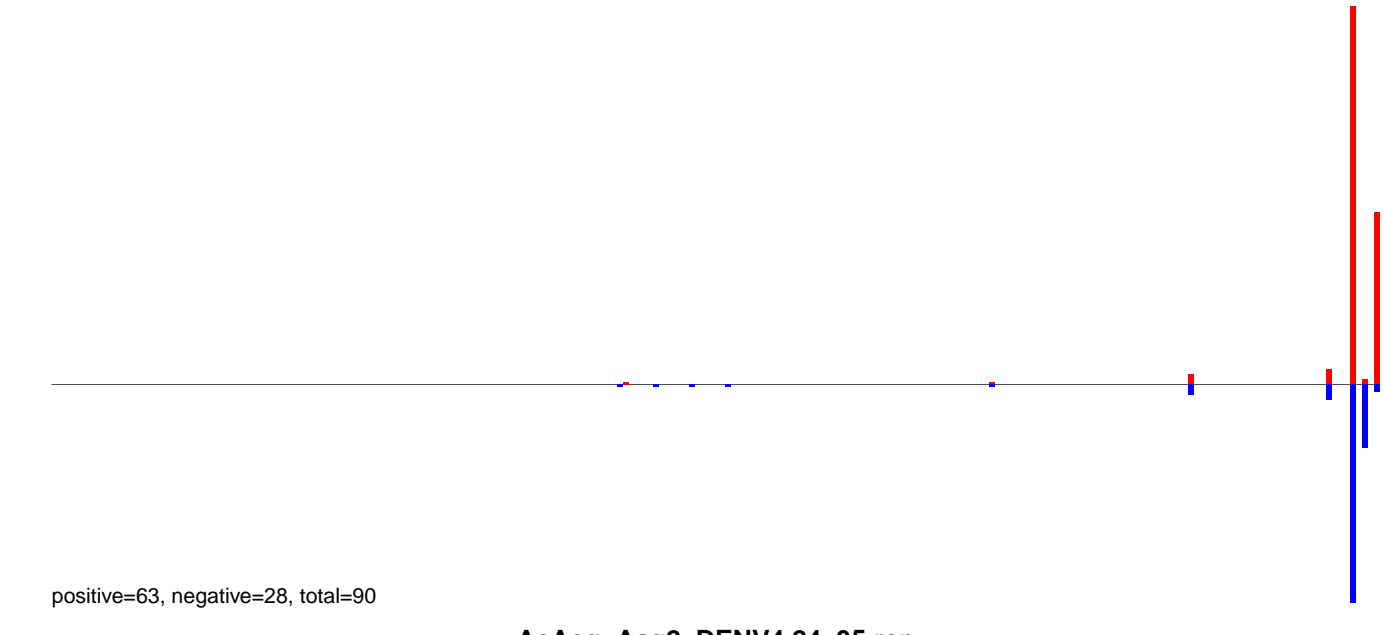
positive=2, negative=0, total=2

AeAeg_Aag2_DENV4.rep

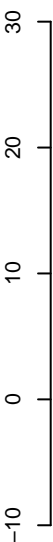
30
20
10
0
-10
-20

positive=64, negative=28, window size=50, length=11064, virus@NC_012932.1-Aedes_flavivirus_genomic_RNA_strain_Narita-21:1-11064

0 2000 4000 6000 8000 10000

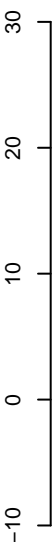


AeAeg_Aag2_DENV4.18_23.rep



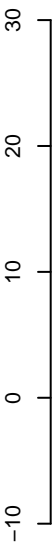
positive=64, negative=28, total=92

AeAeg_Aag2_DENV4.24_35.rep

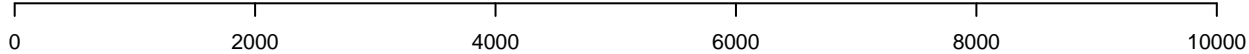


positive=2, negative=0, total=2

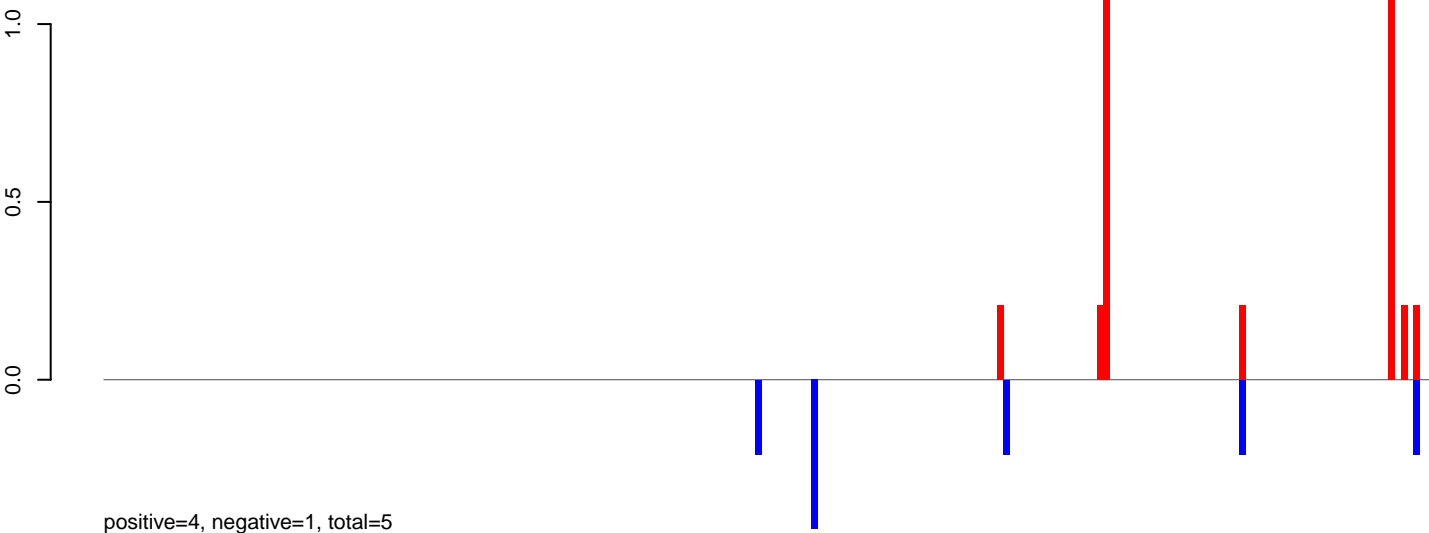
AeAeg_Aag2_DENV4.rep



positive=66, negative=26, total=92, length=11079, virus@KC181923.1-Aedes_flavivirus_strain_AEFV-SPFLD-MO-2011-MP6:1-11079



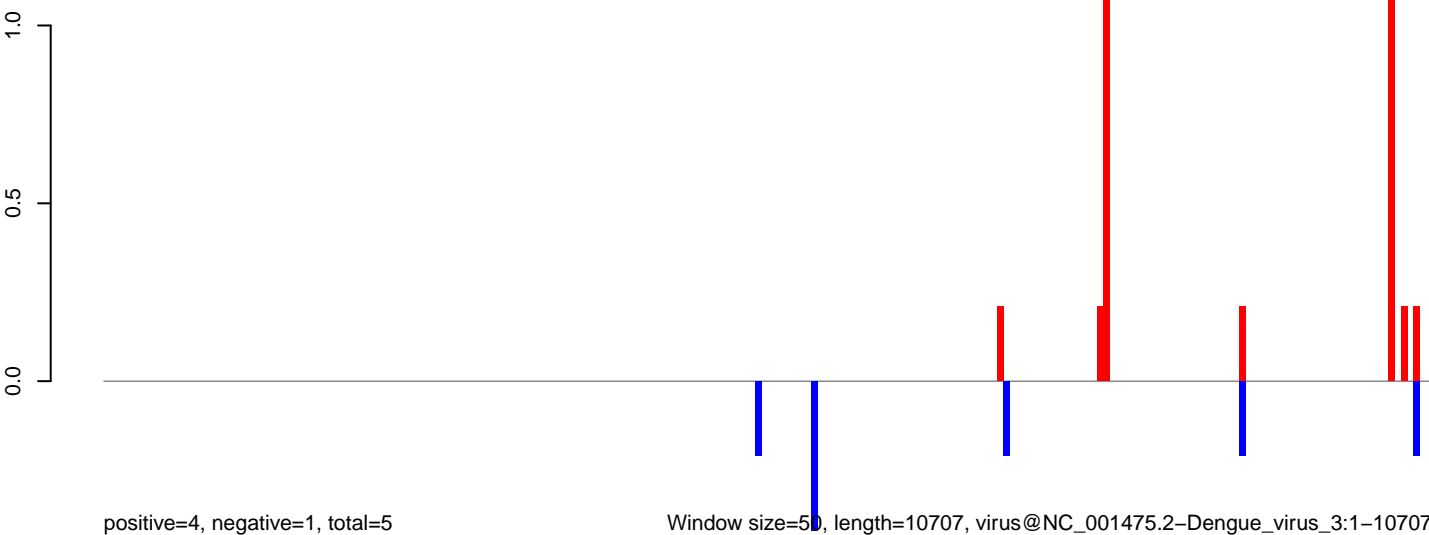
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

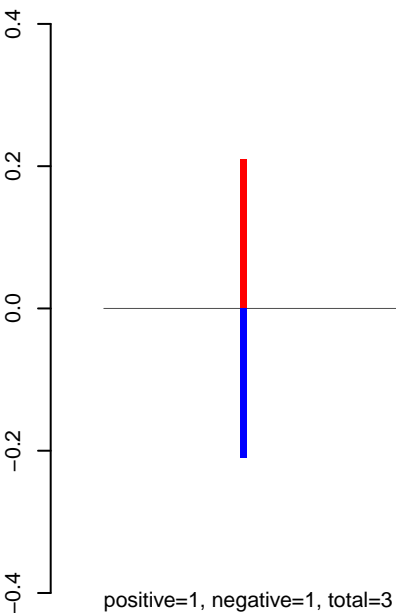


AeAeg_Aag2_DENV4.rep

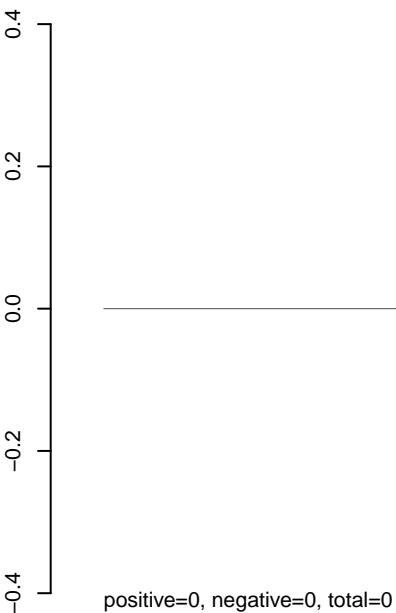


0 2000 4000 6000 8000 10000

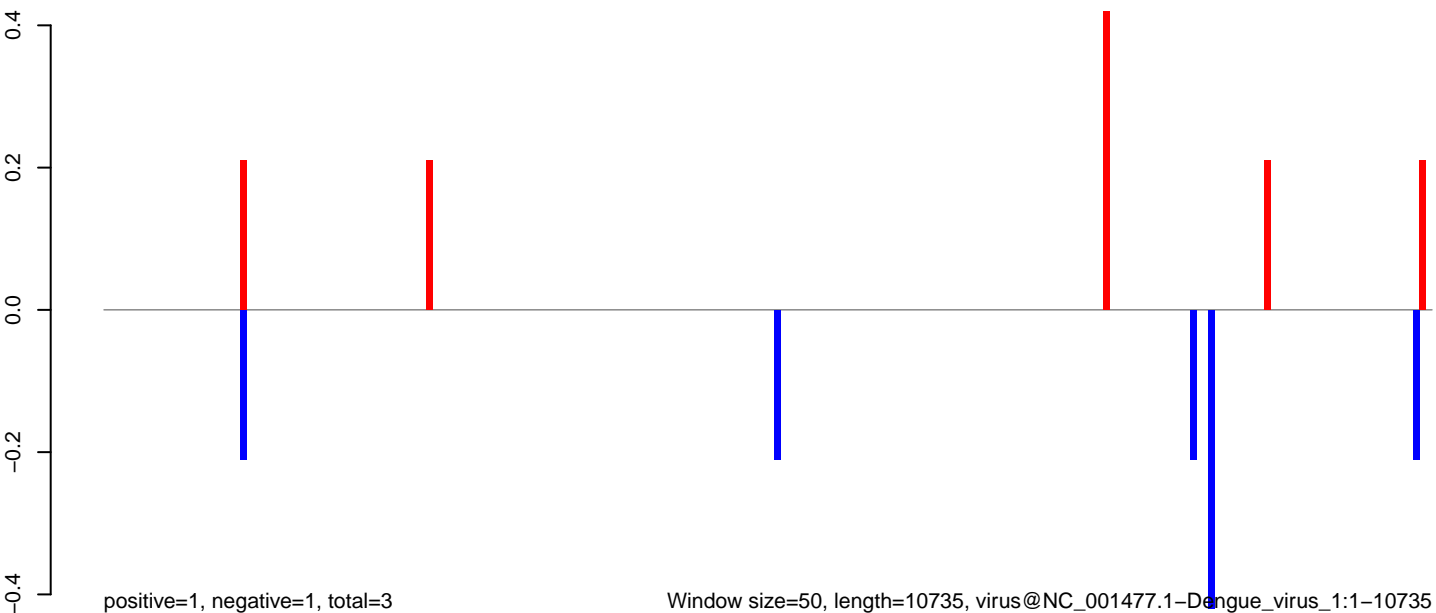
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep

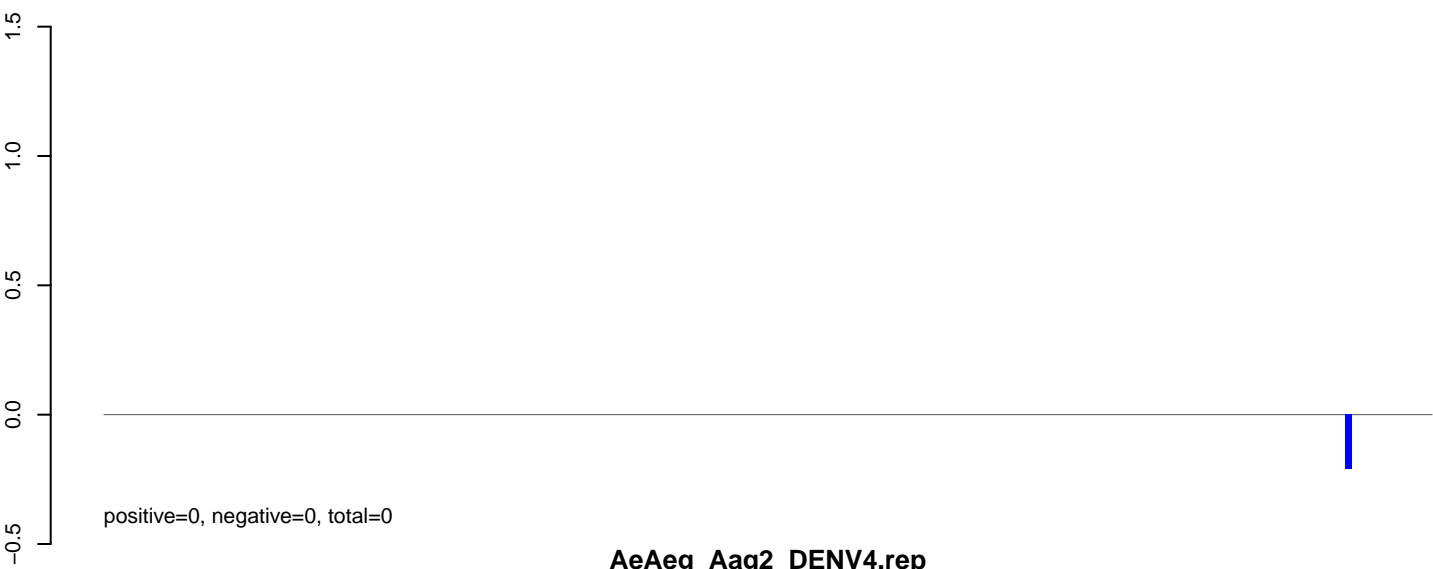


0 2000 4000 6000 8000 10000

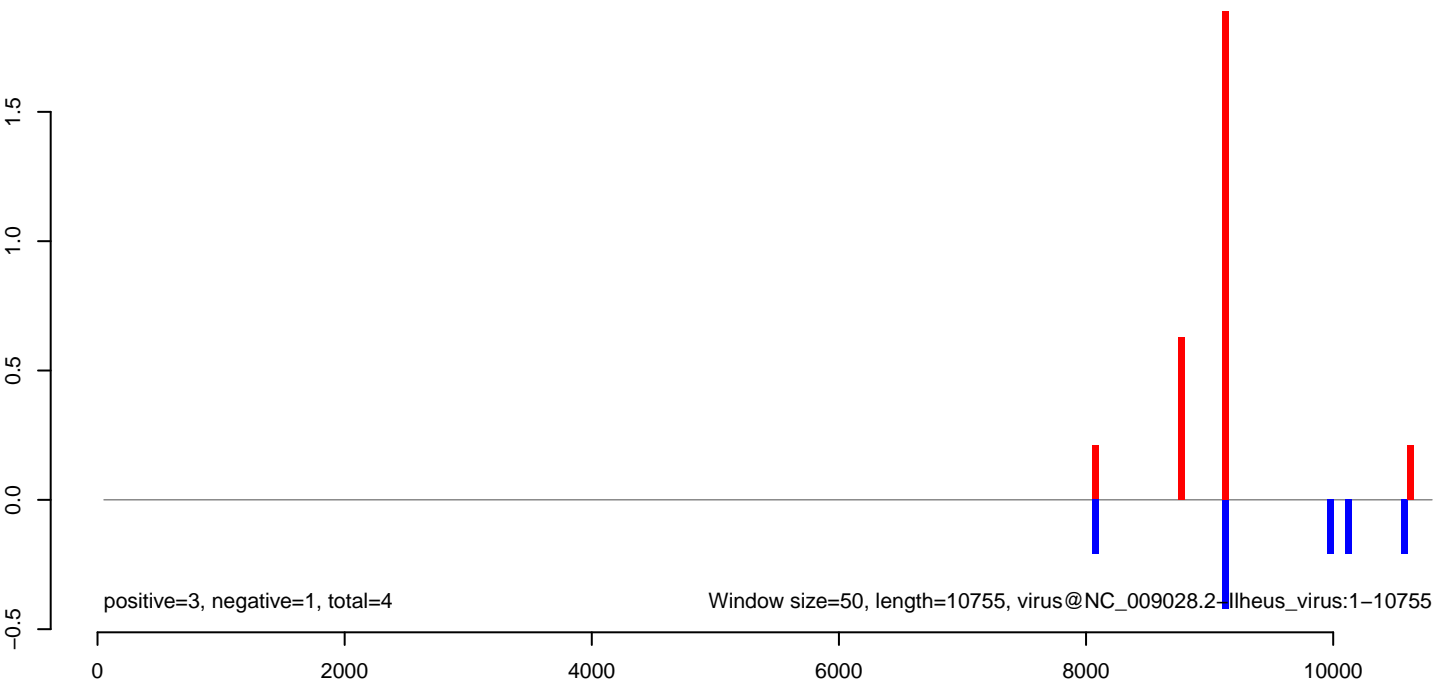
AeAeg_Aag2_DENV4.18_23.rep



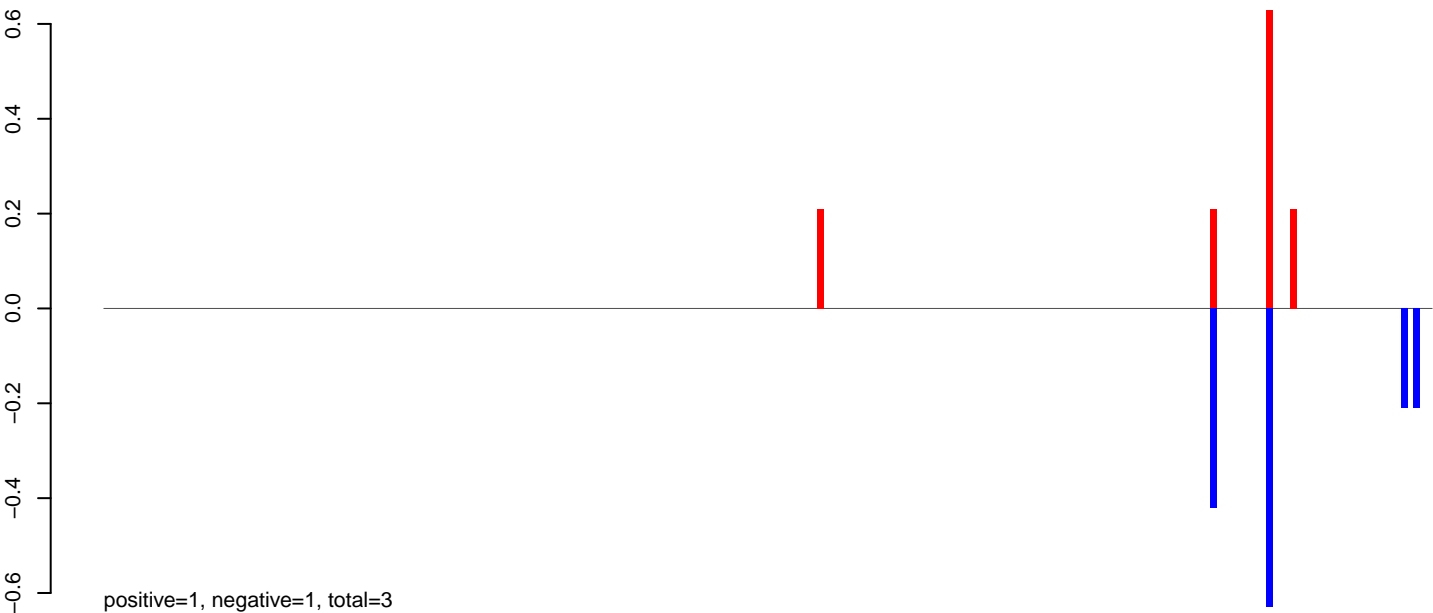
AeAeg_Aag2_DENV4.24_35.rep



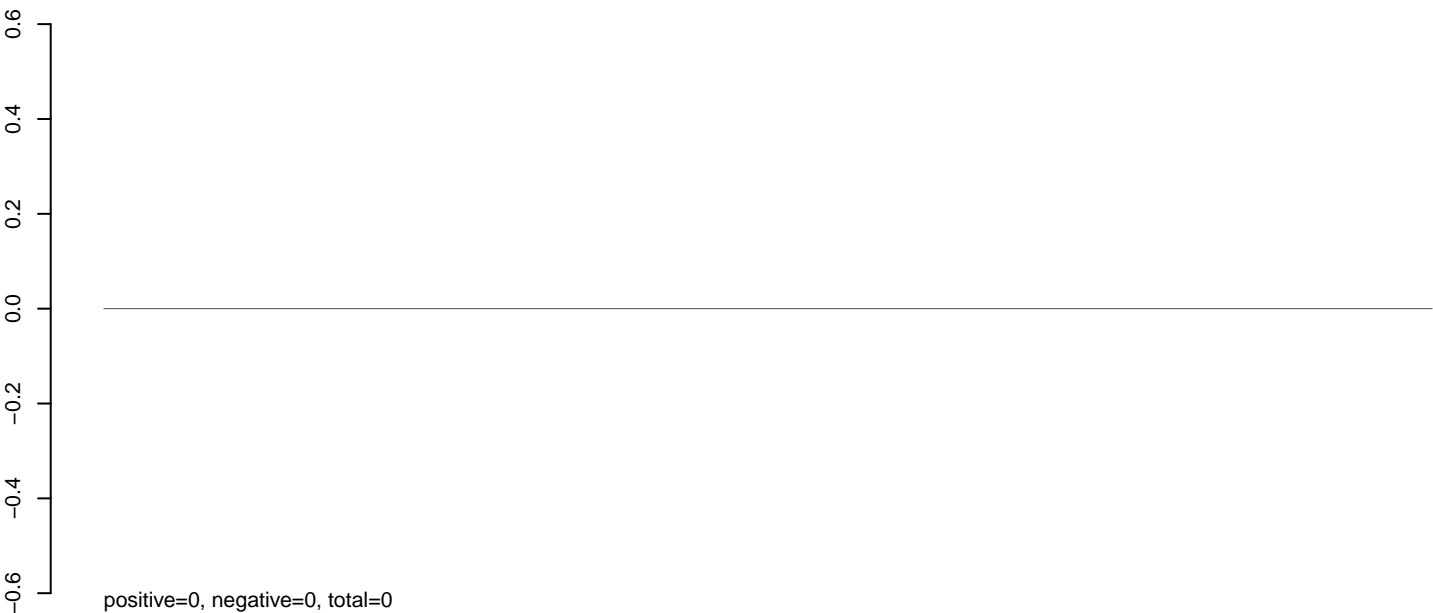
AeAeg_Aag2_DENV4.rep



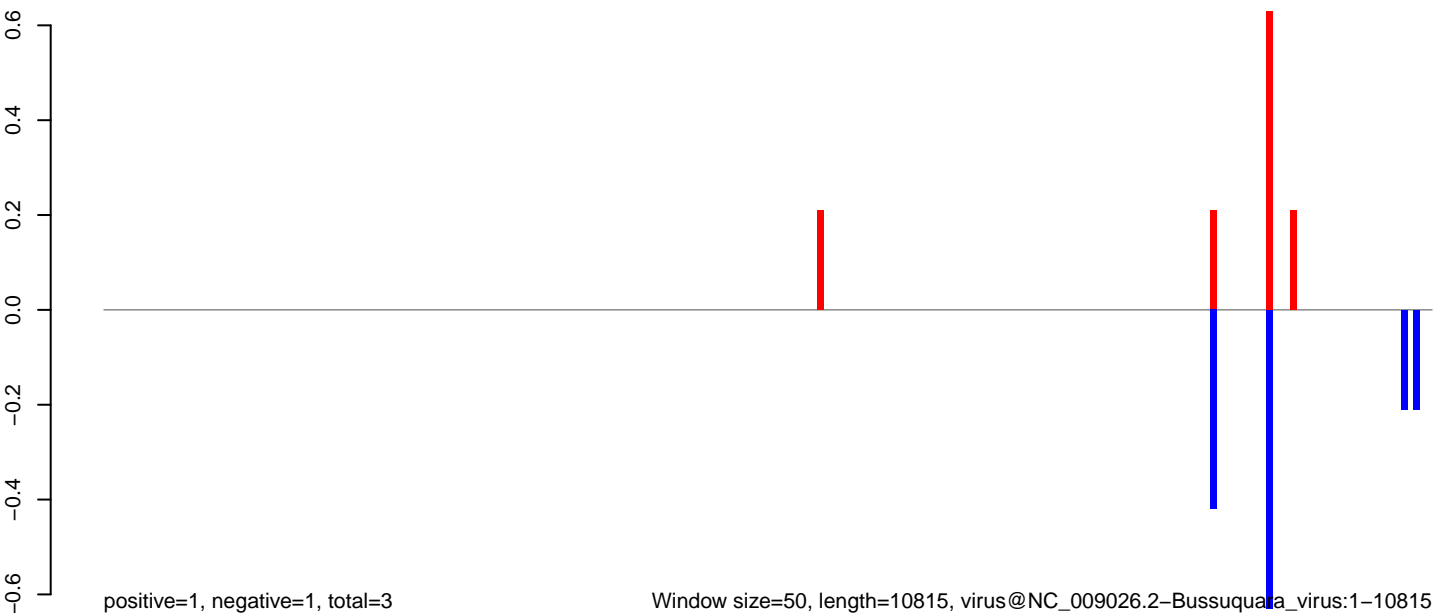
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



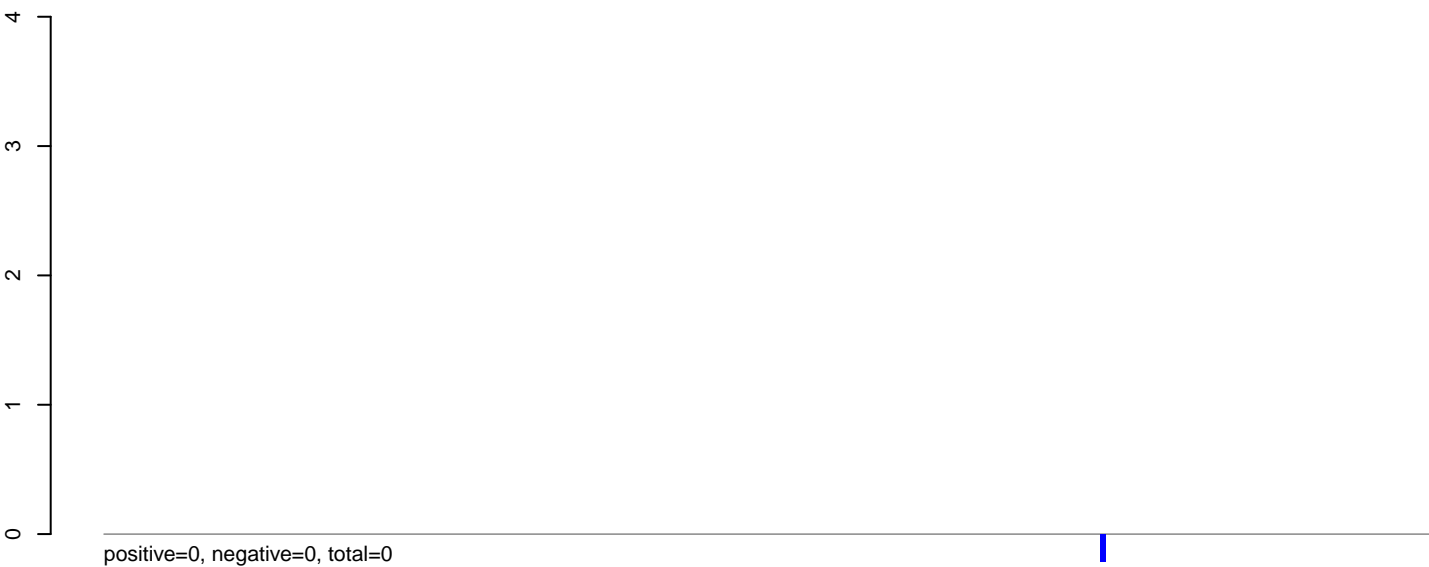
Window size=50, length=10815, virus@NC_009026.2-Bussuquara_virus:1-10815

0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



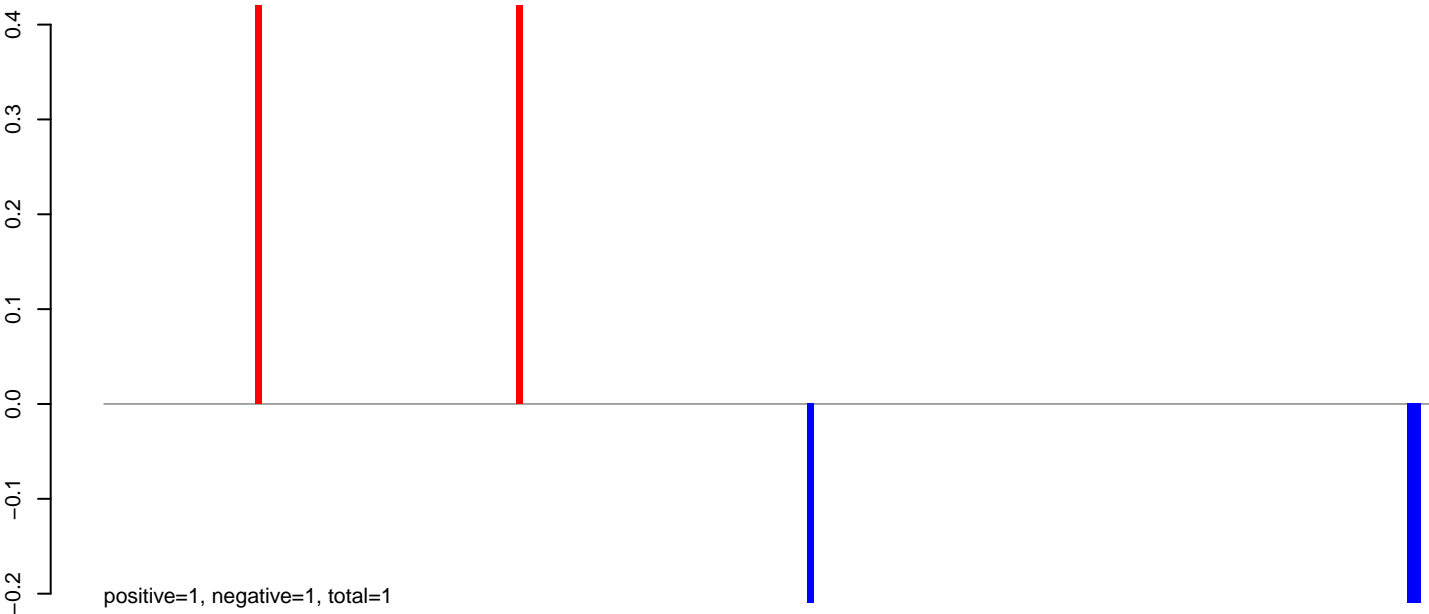
AeAeg_Aag2_DENV4.rep



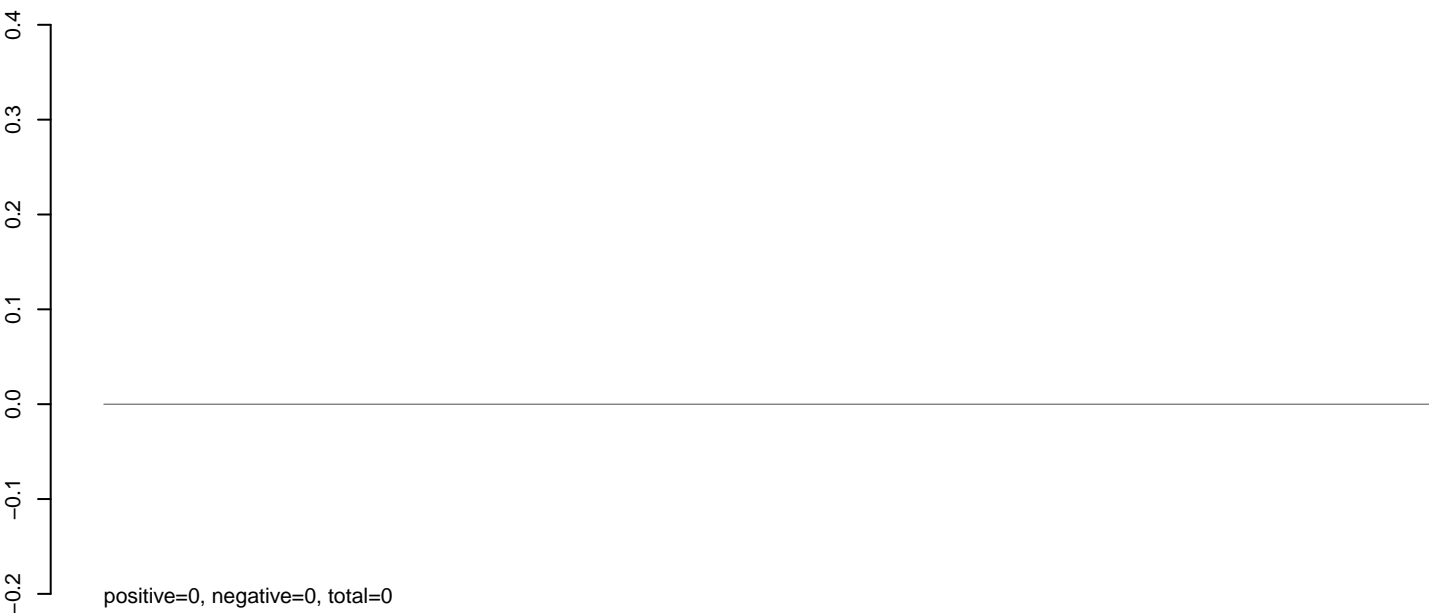
positive=5, negative=0, total=5, virus@NC_038259.1-Badu_virus_isolate_TS6347_segment_S_plus_segment_M_plus_segment_L:1-12247

0 2000 4000 6000 8000 10000 12000

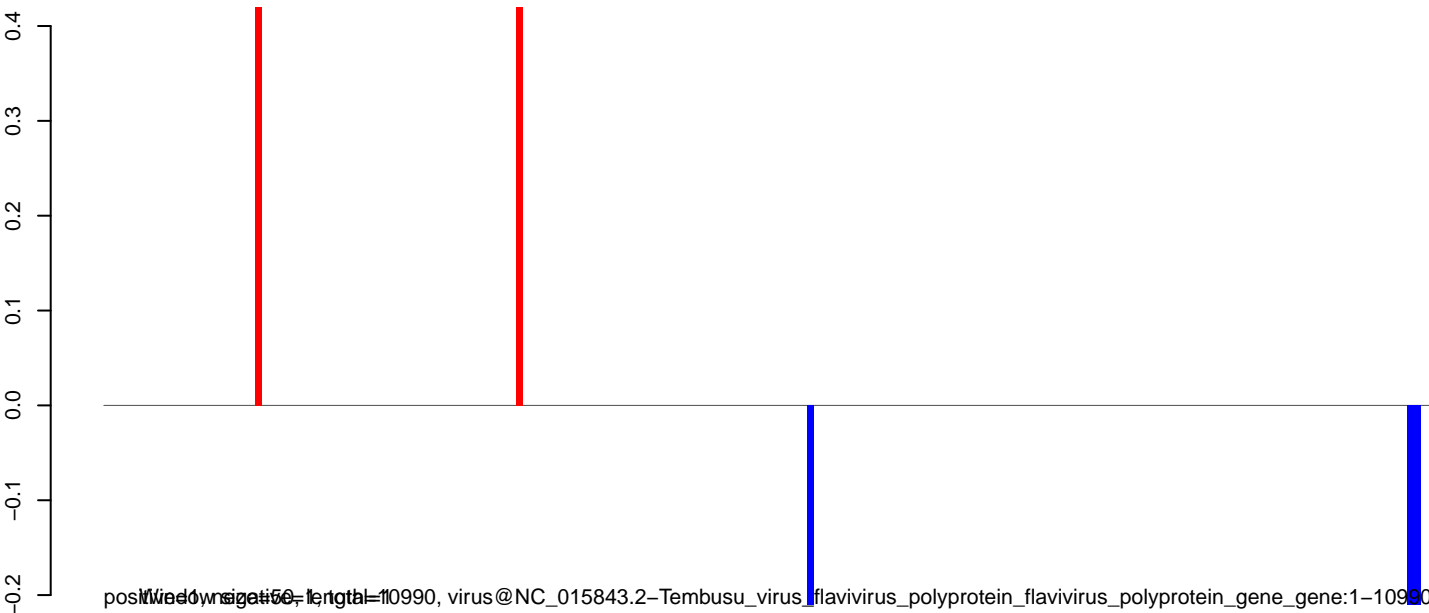
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



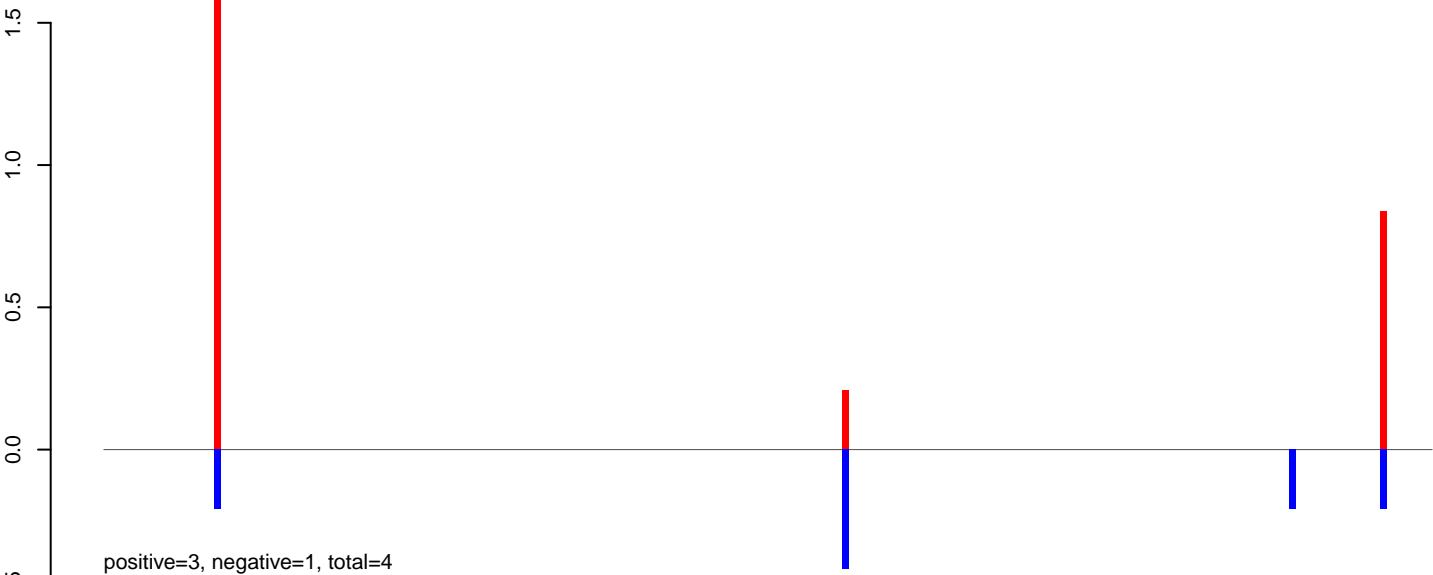
AeAeg_Aag2_DENV4.rep



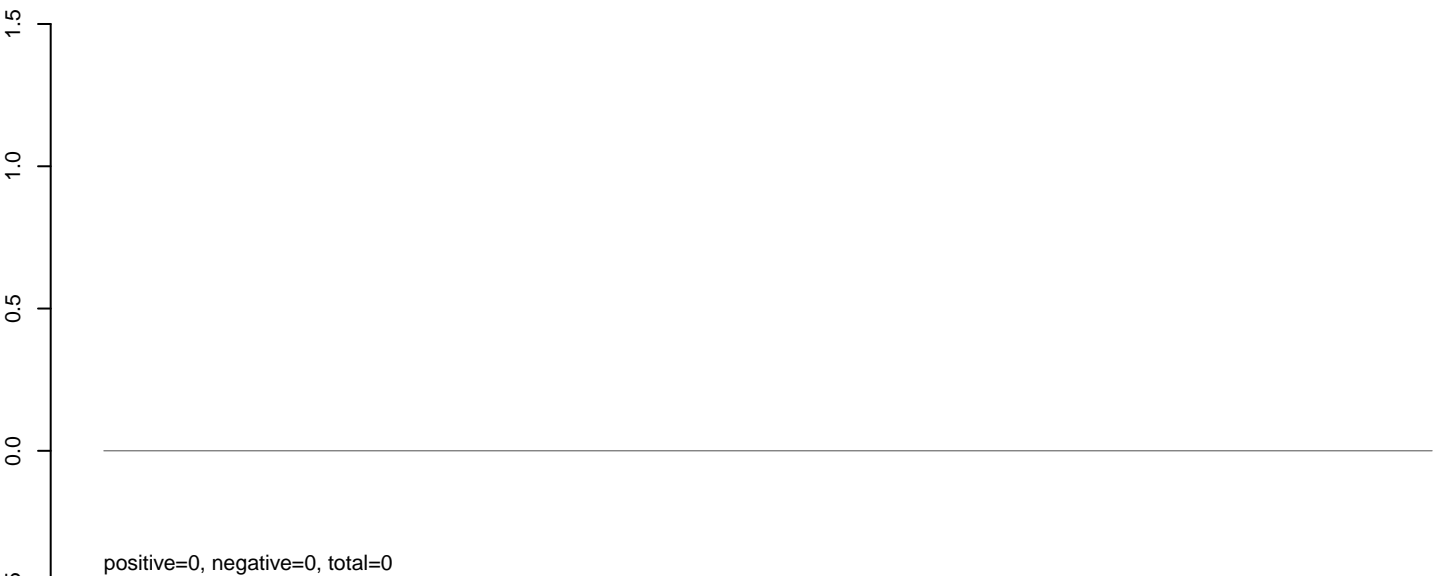
Windows size=50, length=10990, virus@NC_015843.2-Tembusu_virus_flavivirus_polyprotein_flavivirus_polyprotein_gene_gene:1-10990

0 2000 4000 6000 8000 10000

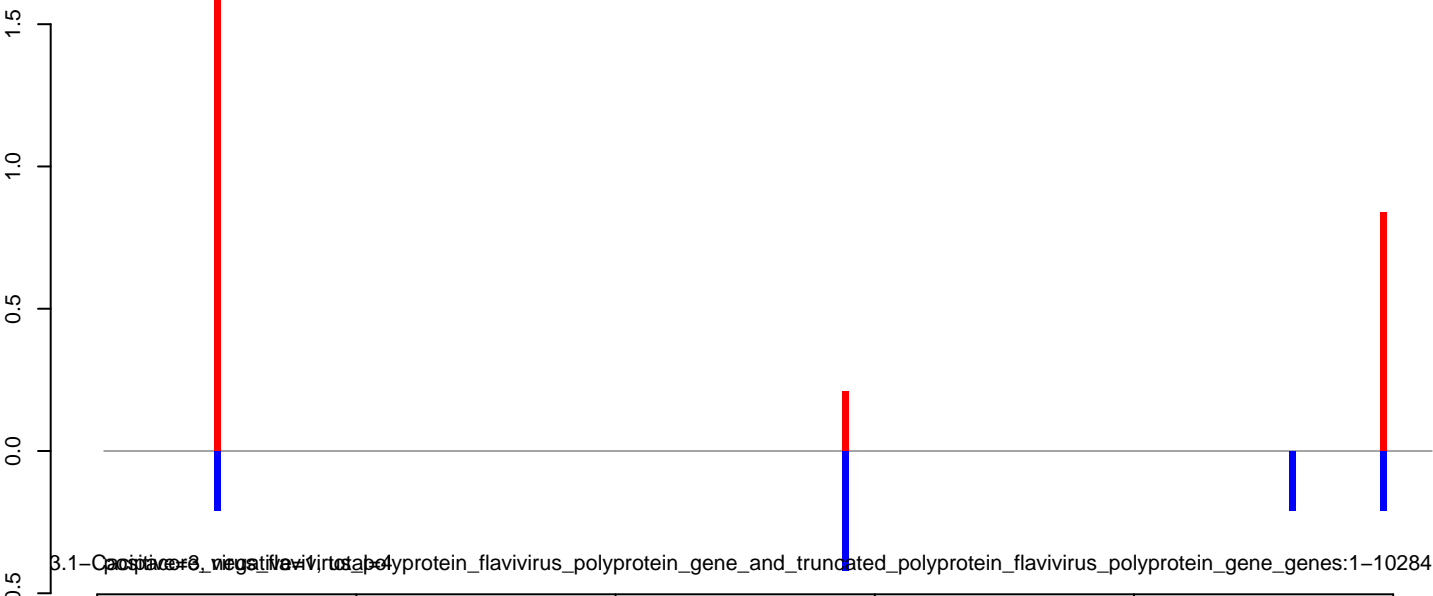
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



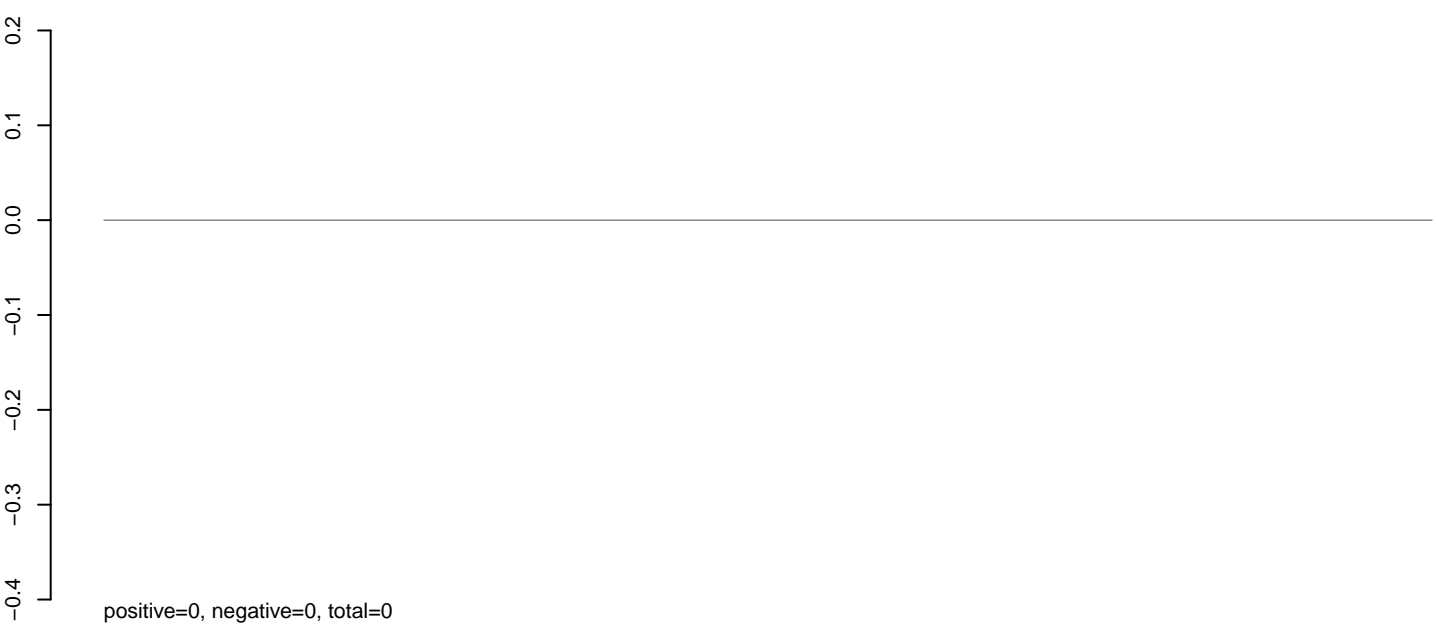
AeAeg_Aag2_DENV4.rep



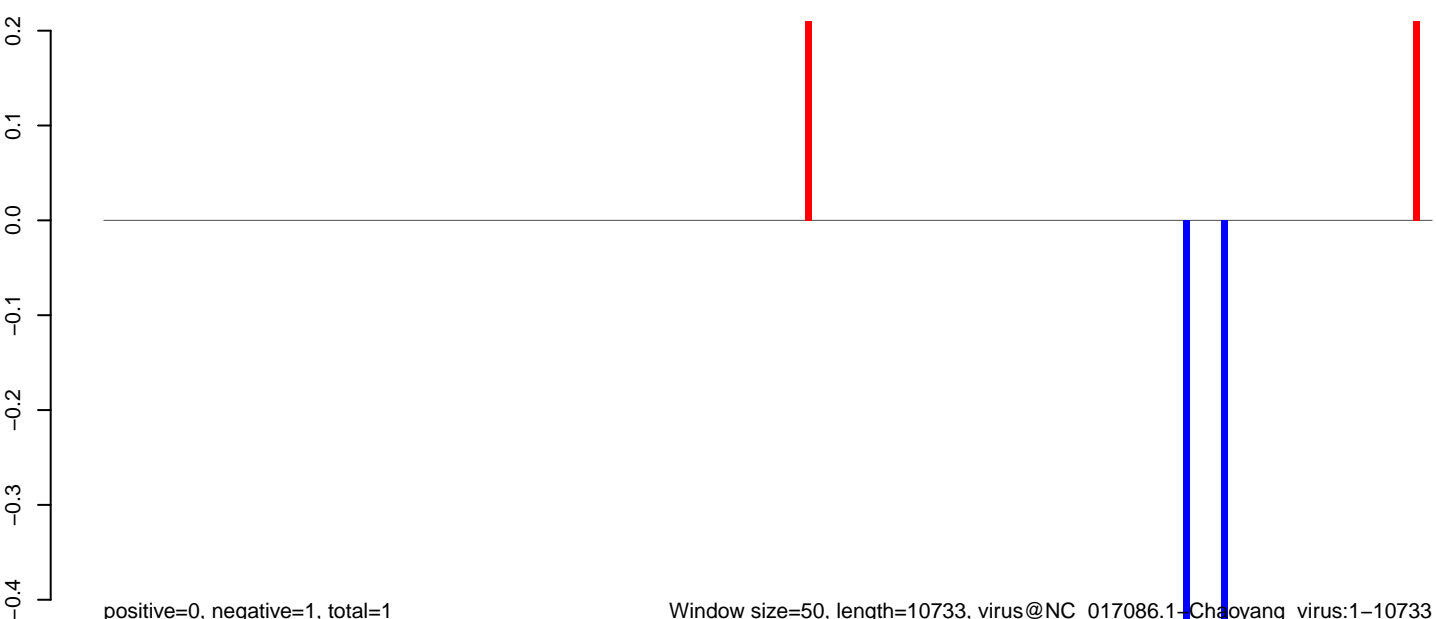
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



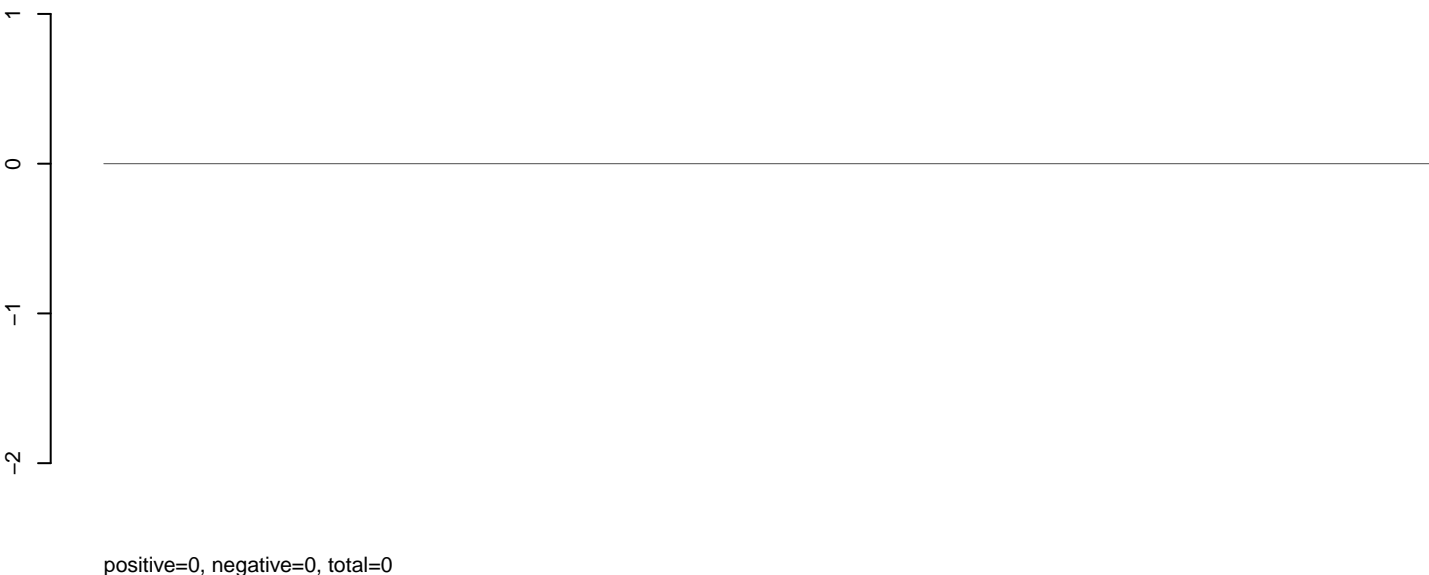
Window size=50, length=10733, virus@NC_017086.1-Chaoyang_virus:1-10733

0 2000 4000 6000 8000 10000

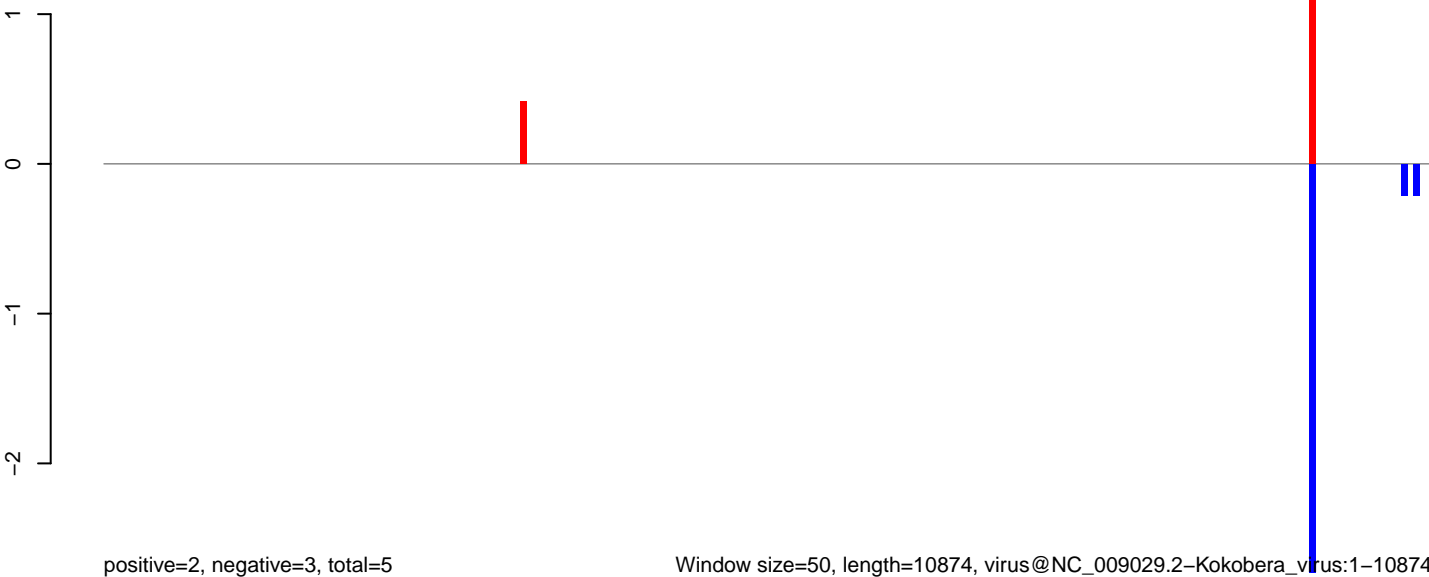
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

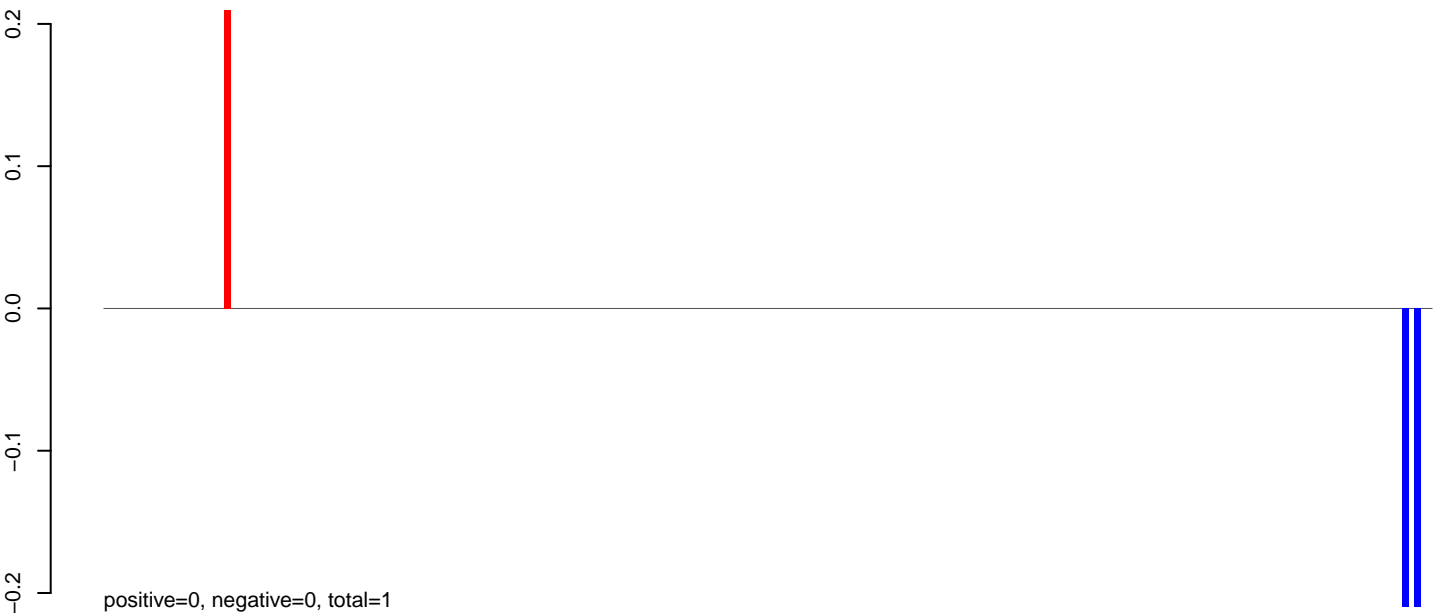


AeAeg_Aag2_DENV4.rep

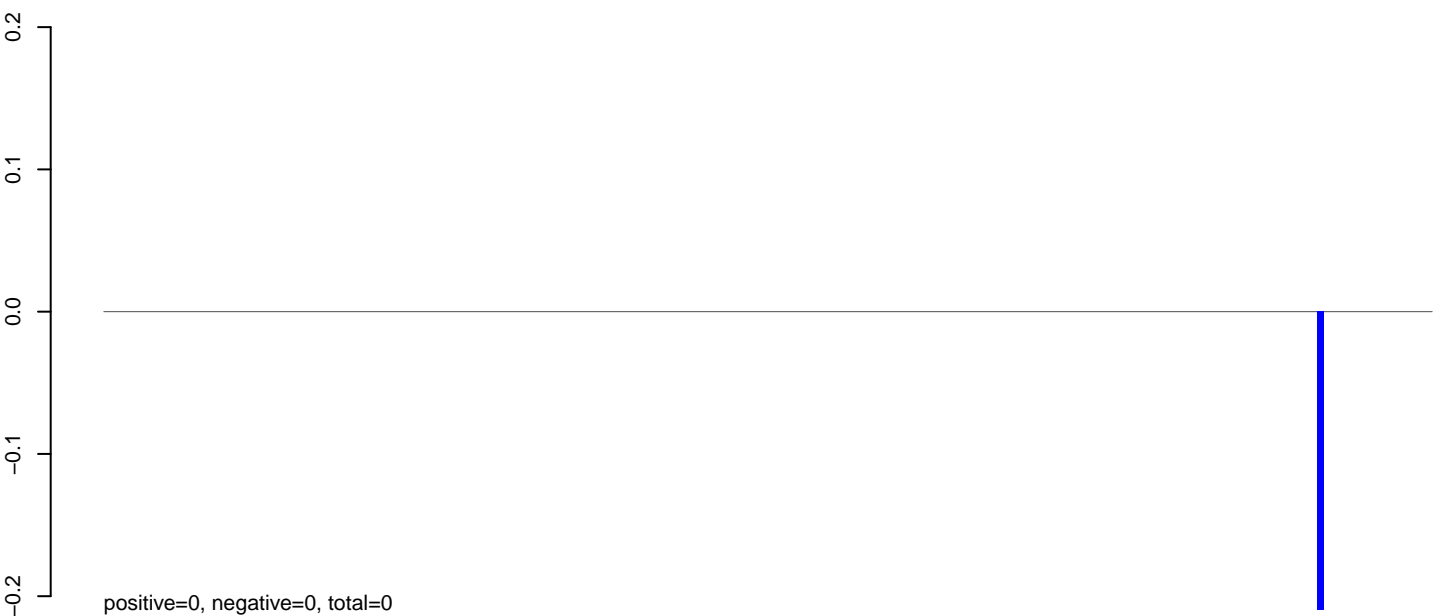


Window size=50, length=11141, virus@NC_001672.1-Tick-borne-encephalitis_virus:1-11141

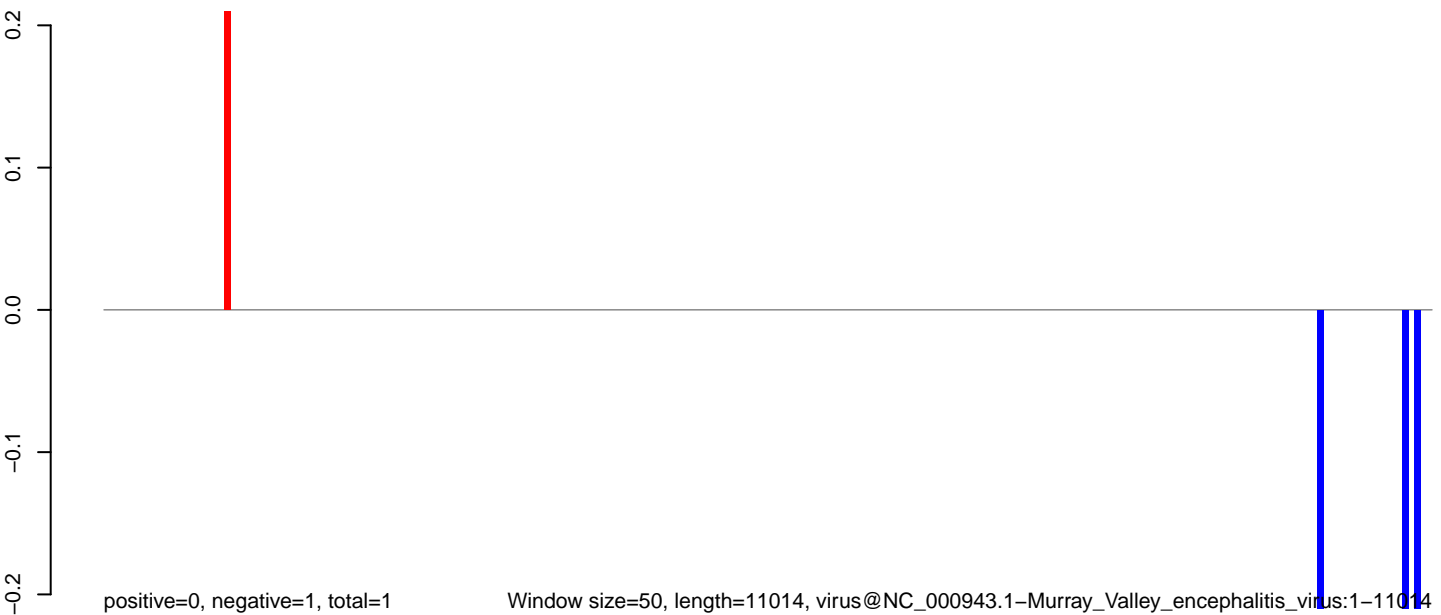
AeAeg_Aag2_DENV4.18_23.rep



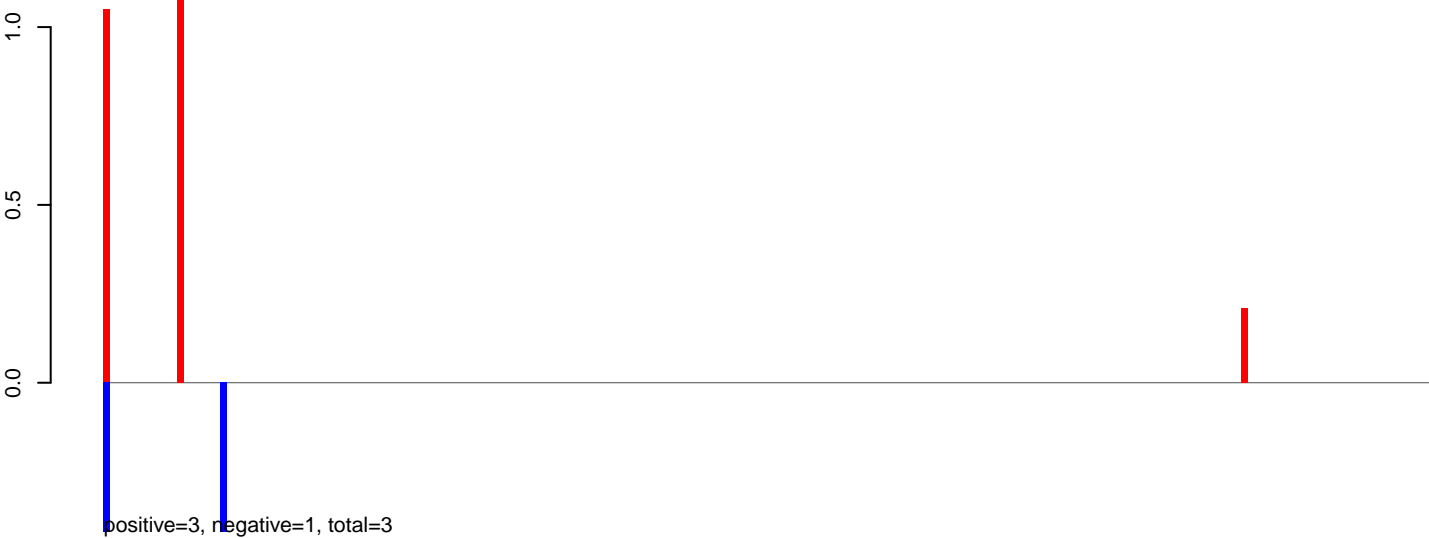
AeAeg_Aag2_DENV4.24_35.rep



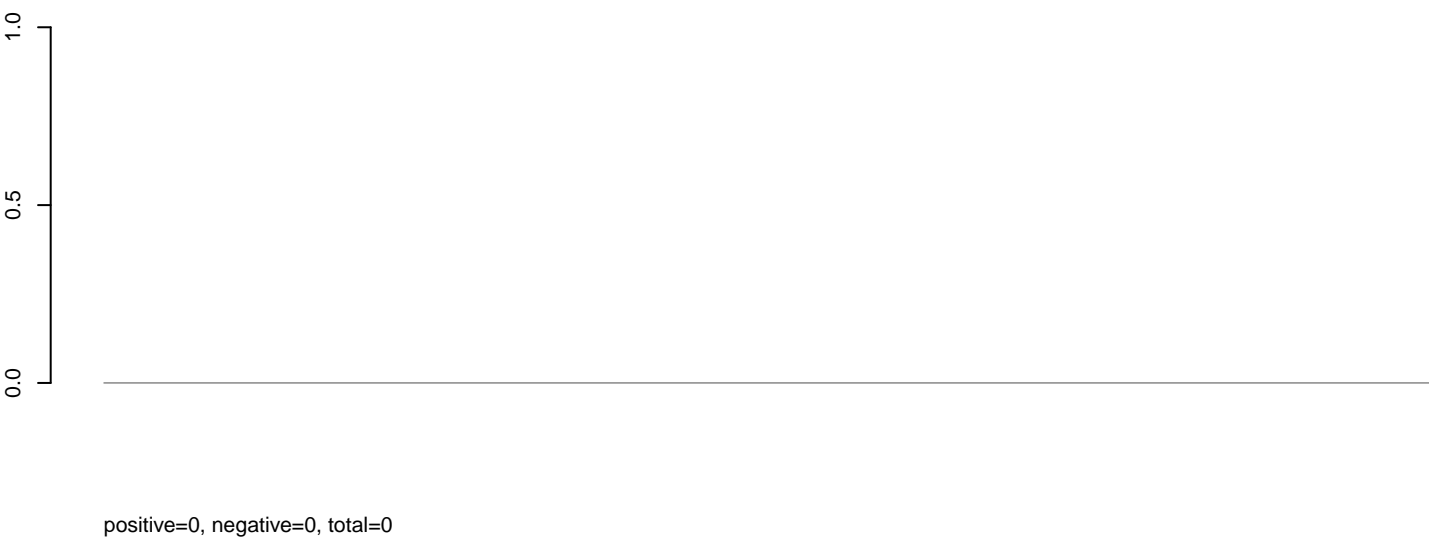
AeAeg_Aag2_DENV4.rep



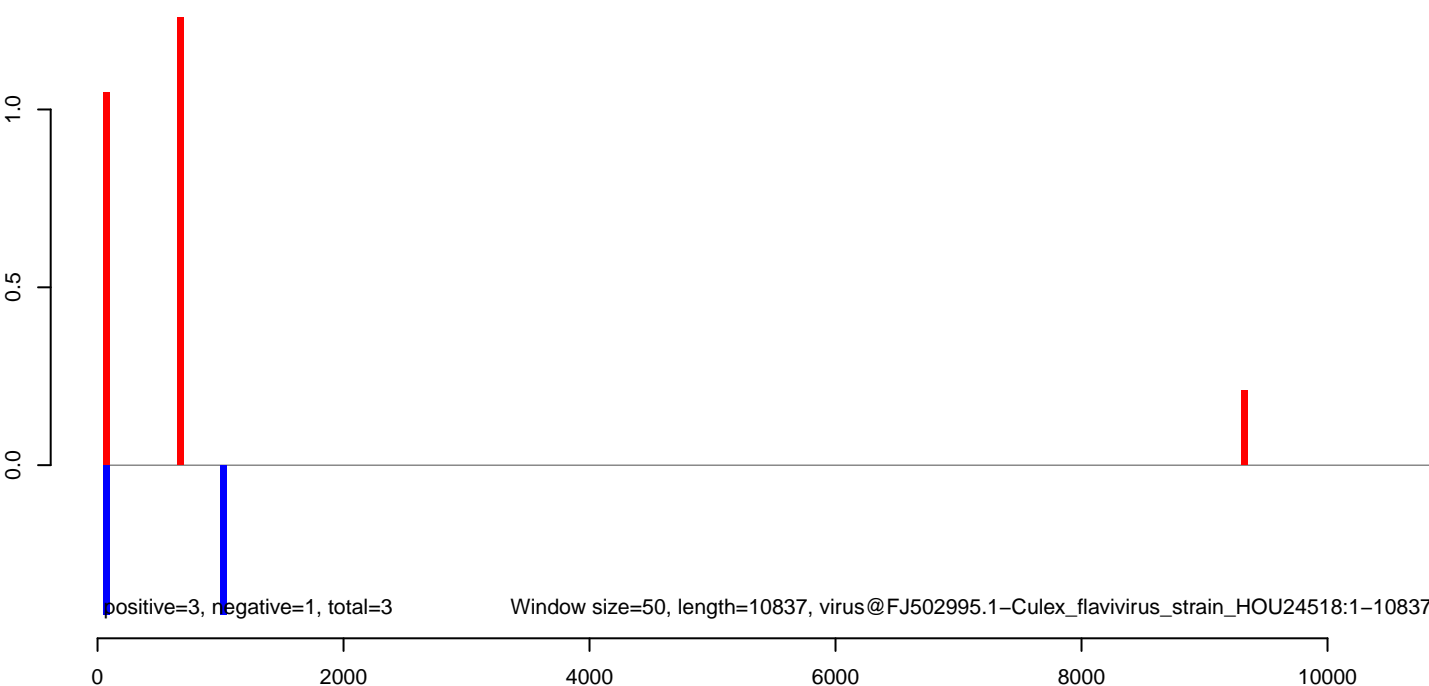
AeAeg_Aag2_DENV4.18_23.rep



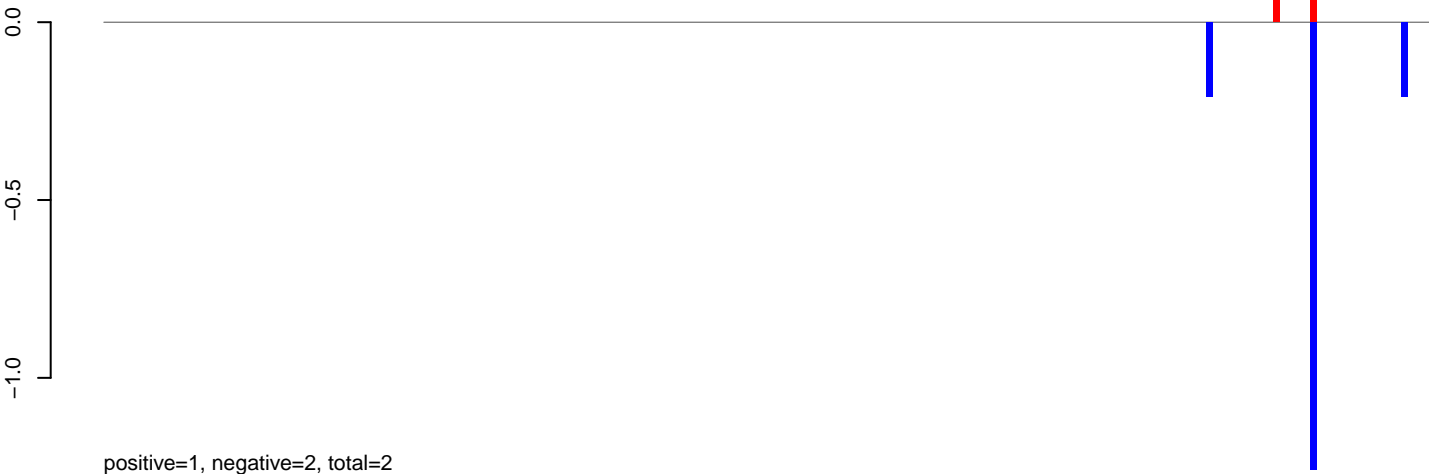
AeAeg_Aag2_DENV4.24_35.rep



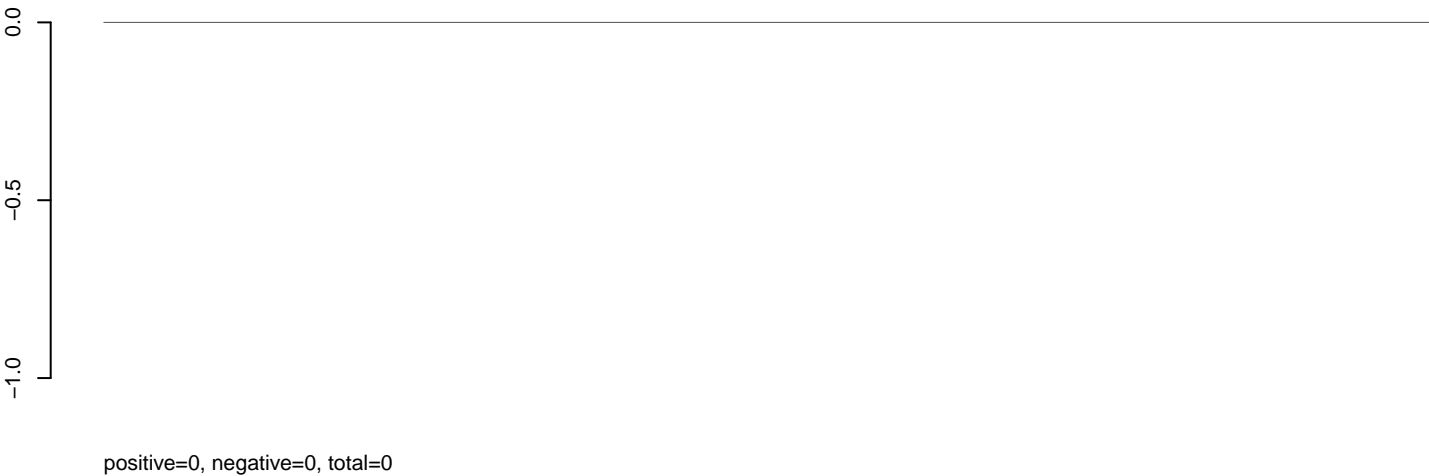
AeAeg_Aag2_DENV4.rep



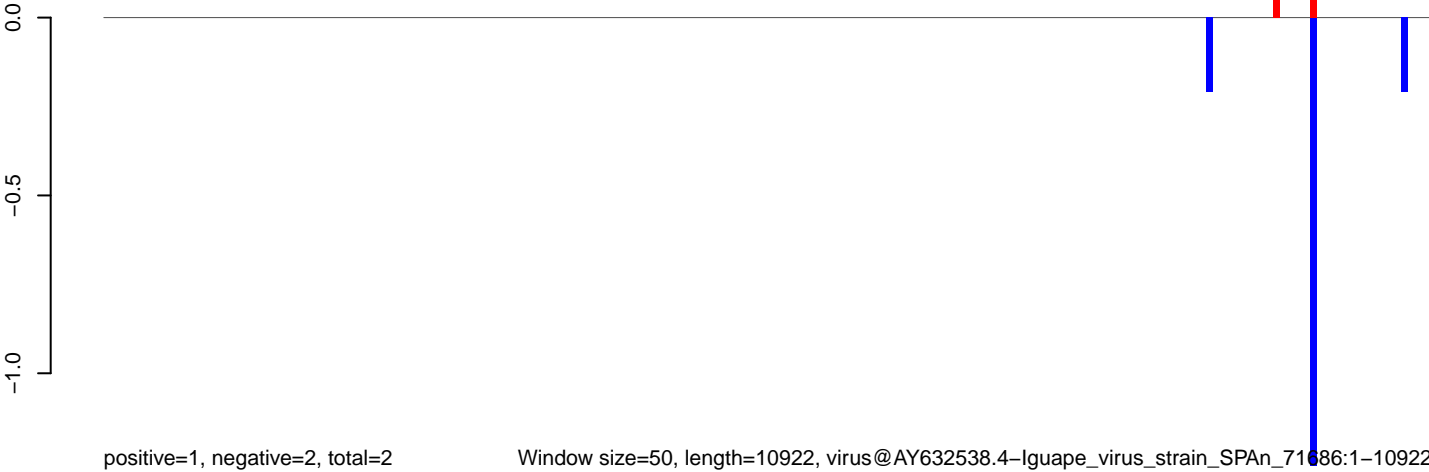
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

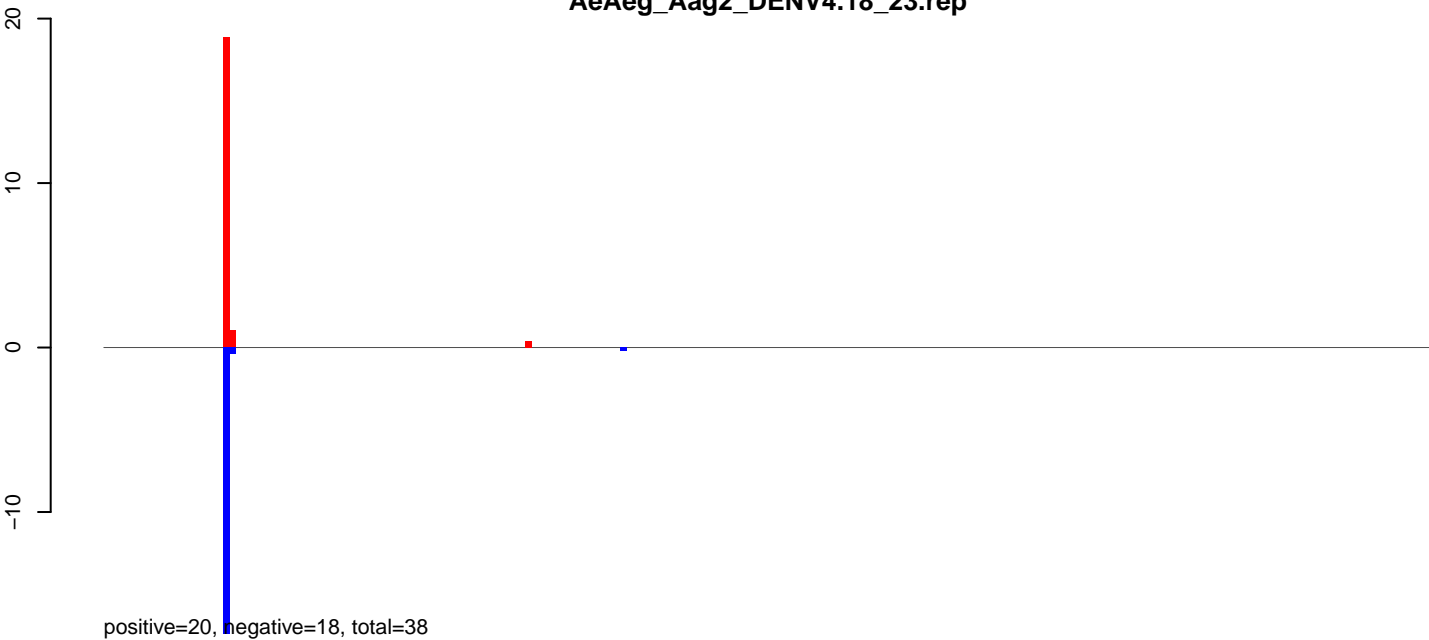


AeAeg_Aag2_DENV4.rep

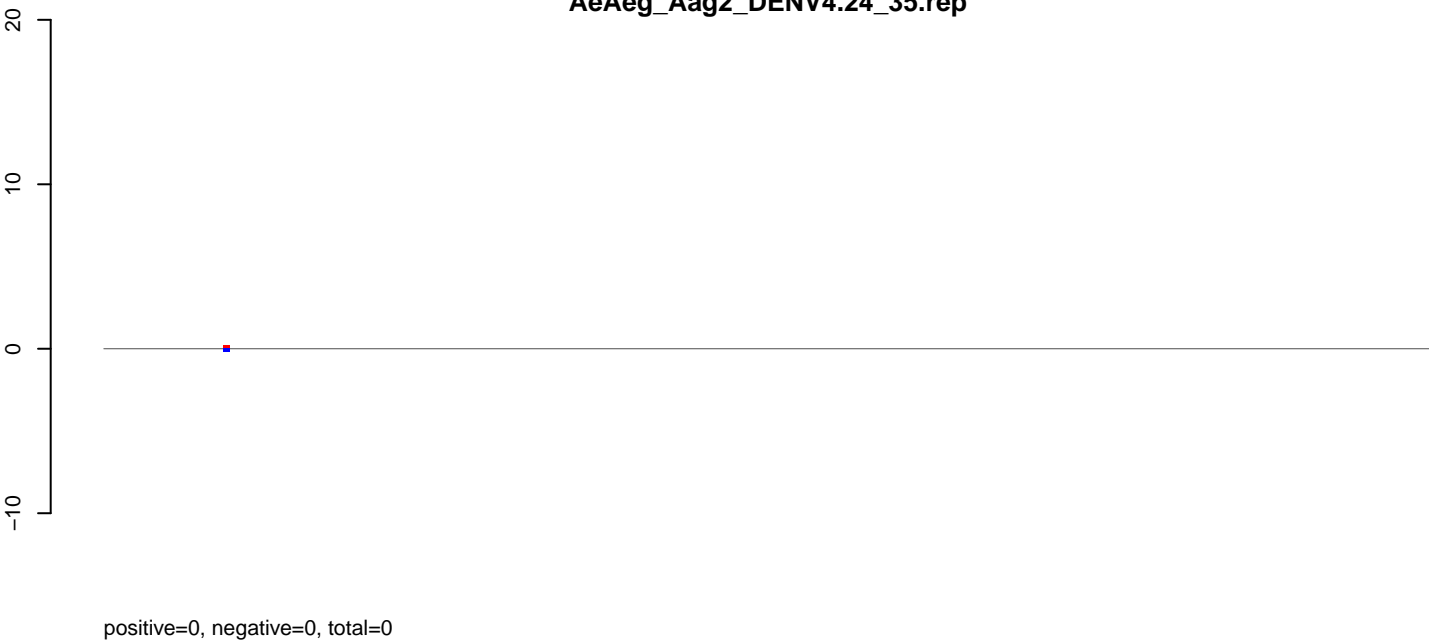


Window size=50, length=10922, virus@AY632538.4-Iguape_virus_strain_SPAn_71686:1-10922

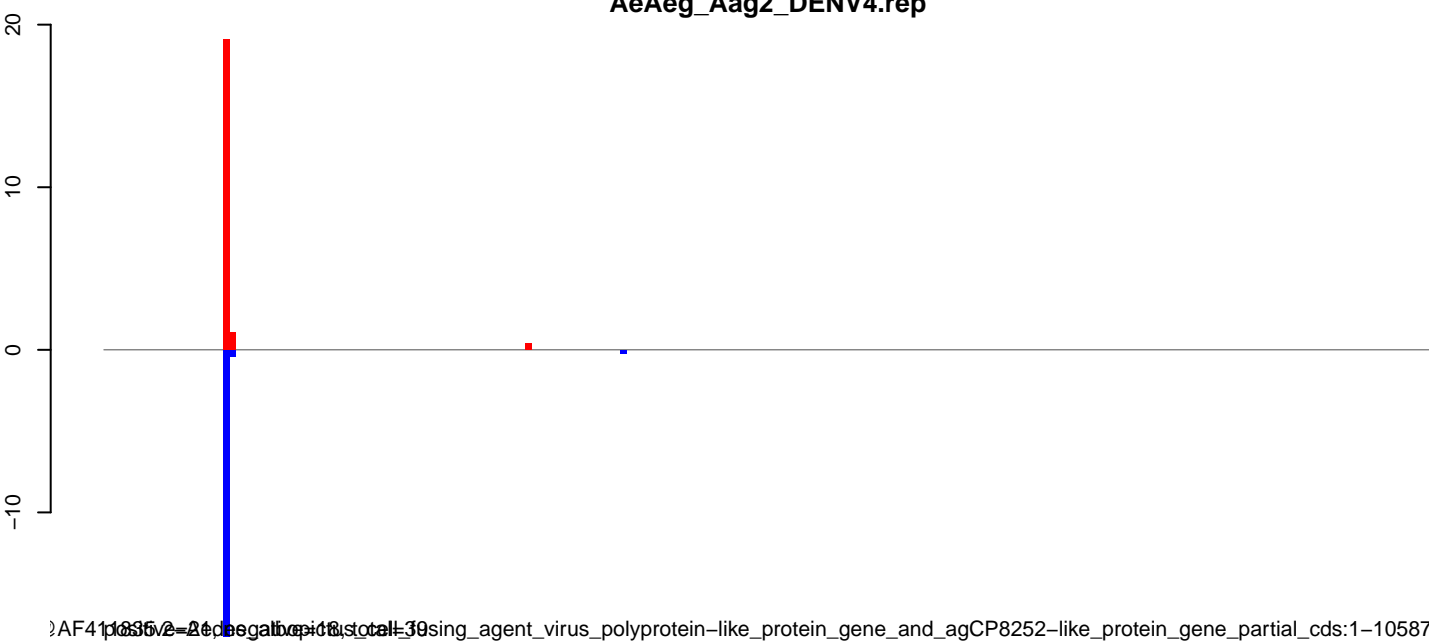
AeAeg_Aag2_DENV4.18_23.rep



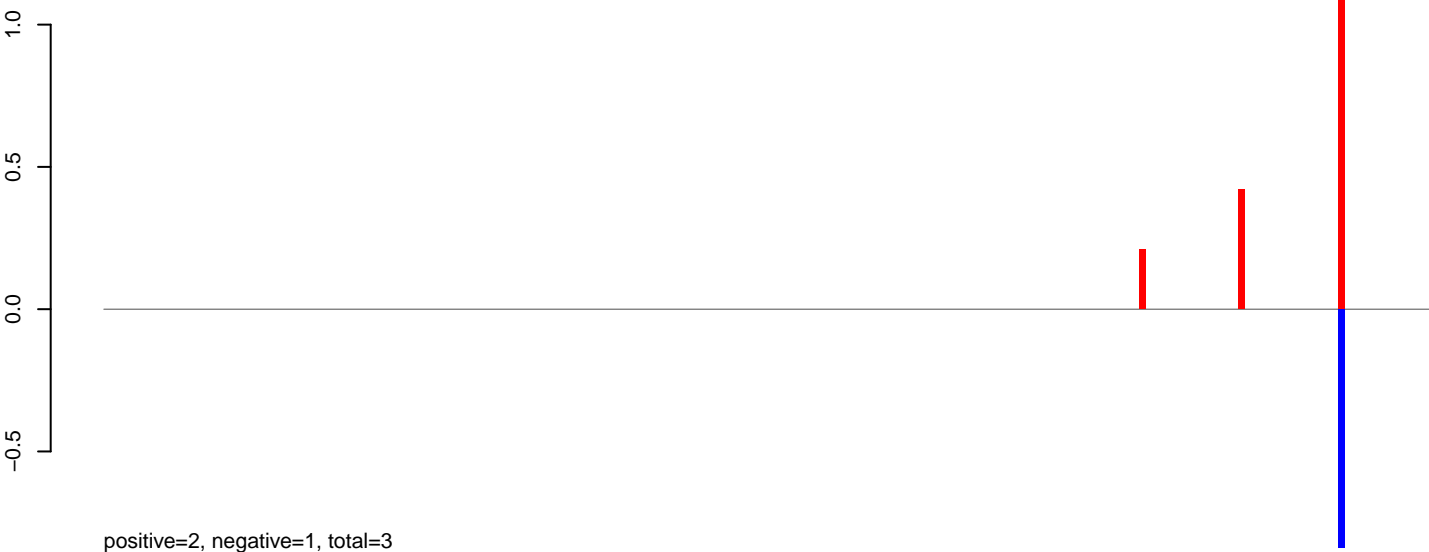
AeAeg_Aag2_DENV4.24_35.rep



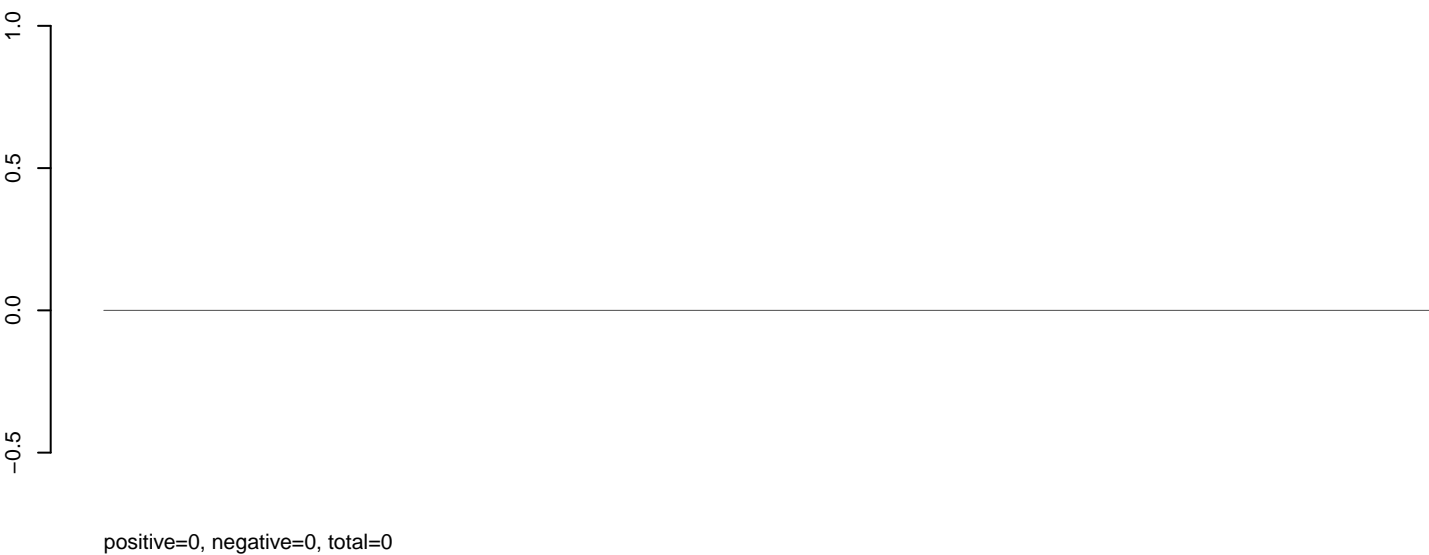
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



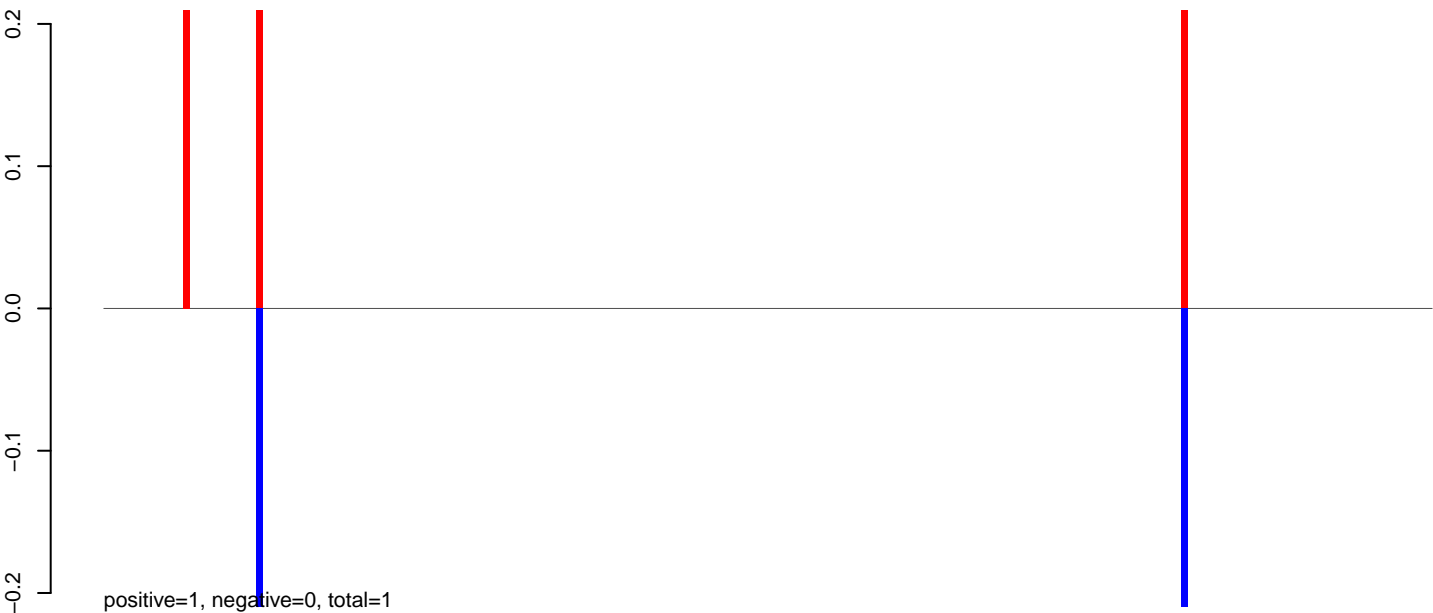
AeAeg_Aag2_DENV4.24_35.rep



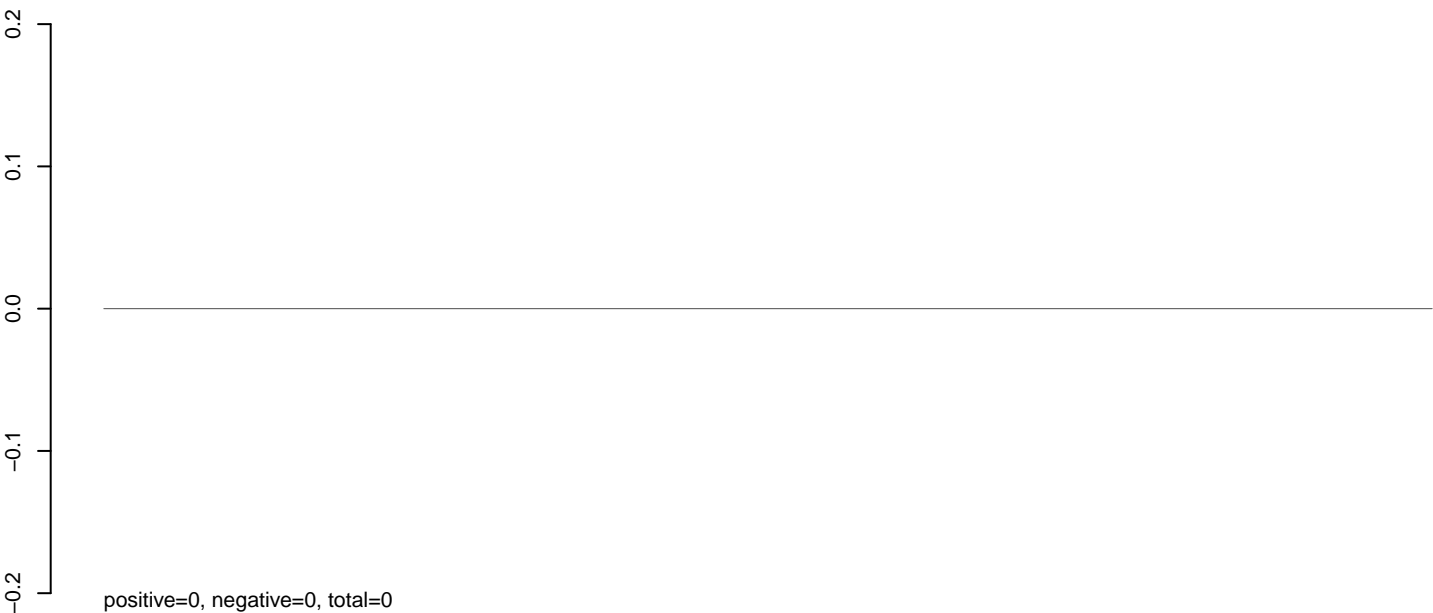
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



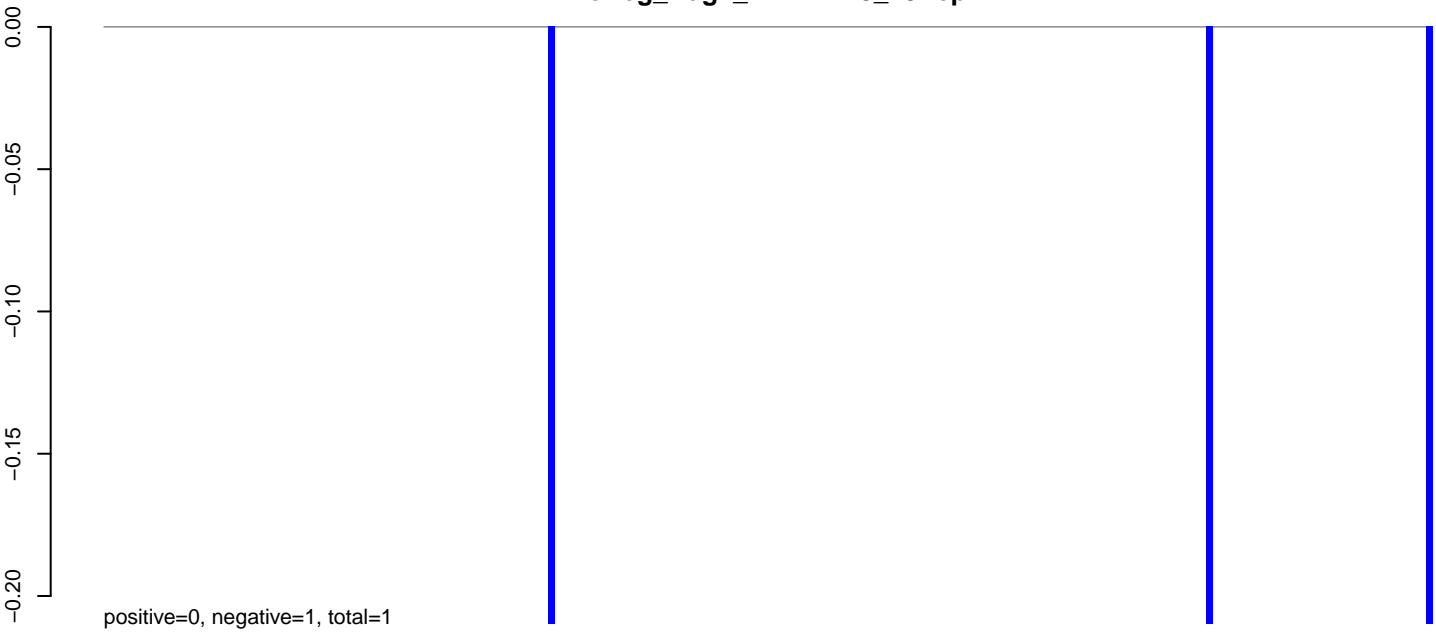
AeAeg_Aag2_DENV4.24_35.rep



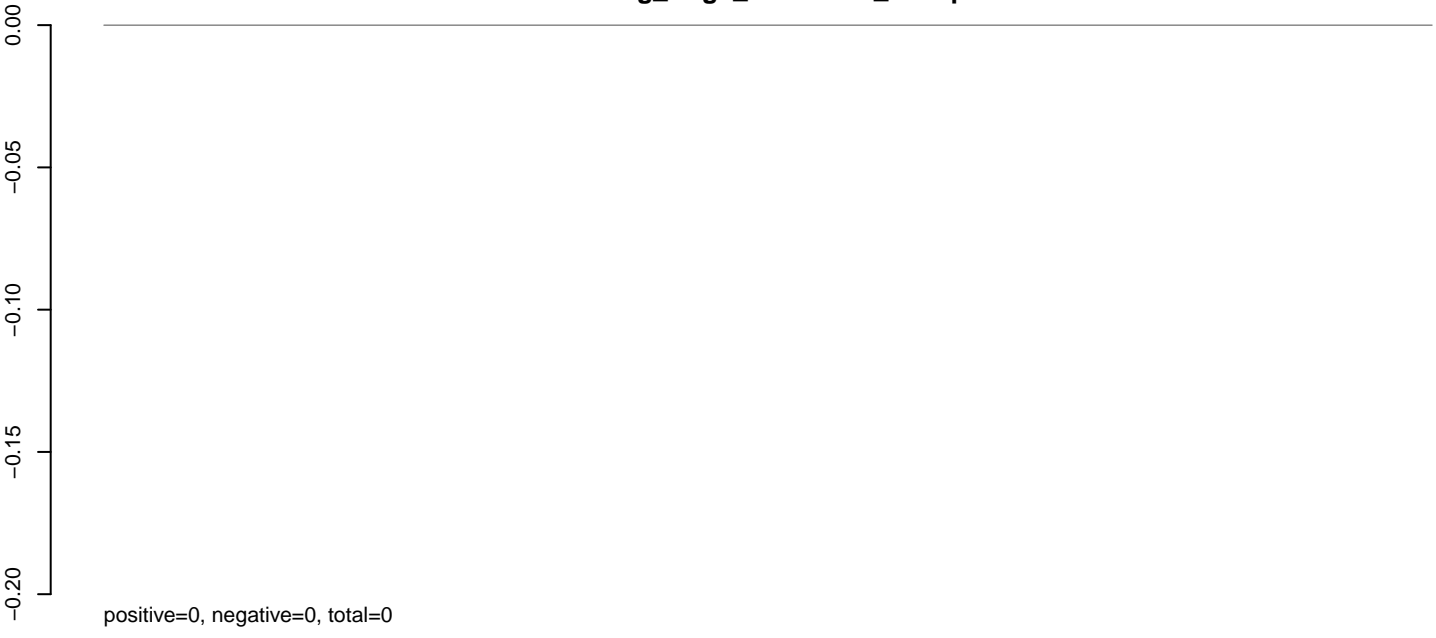
AeAeg_Aag2_DENV4.rep



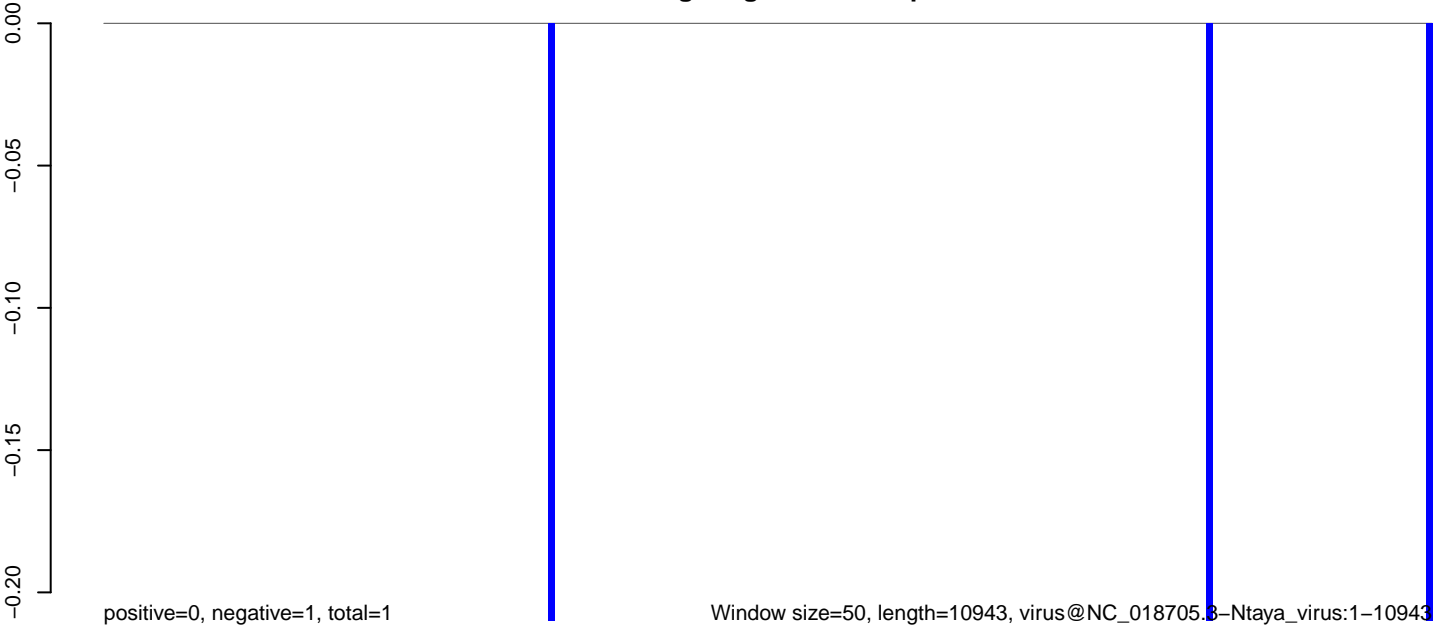
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

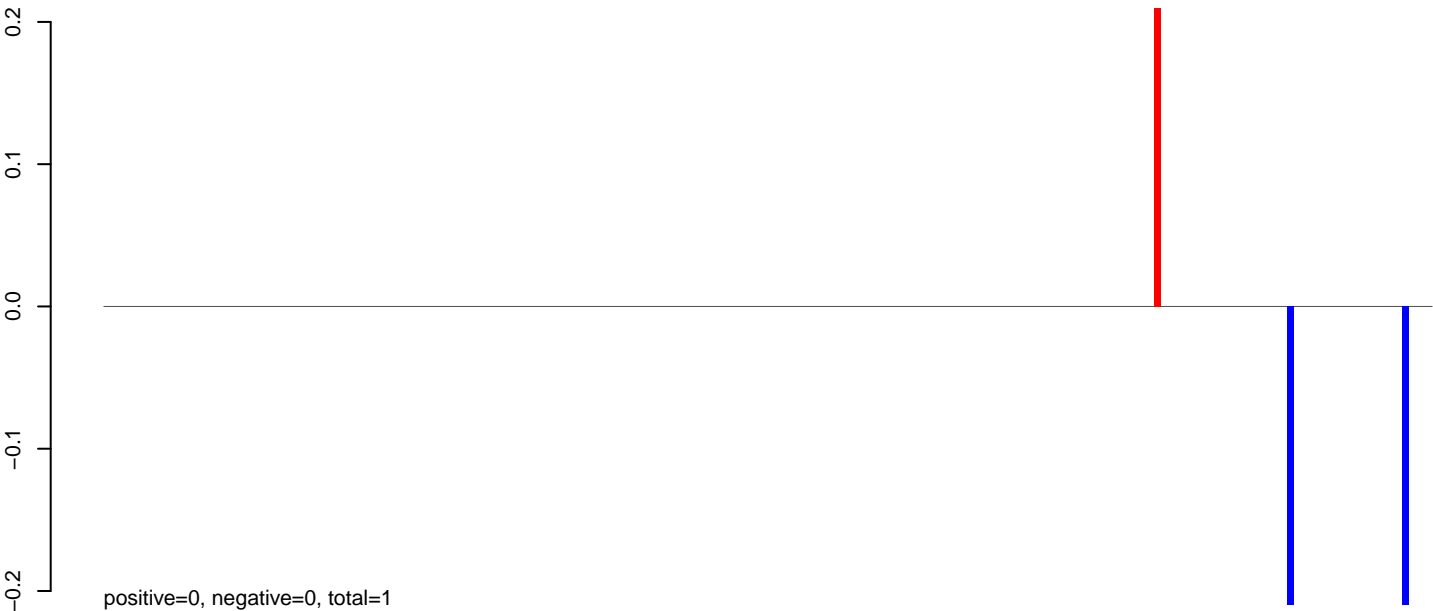


AeAeg_Aag2_DENV4.rep

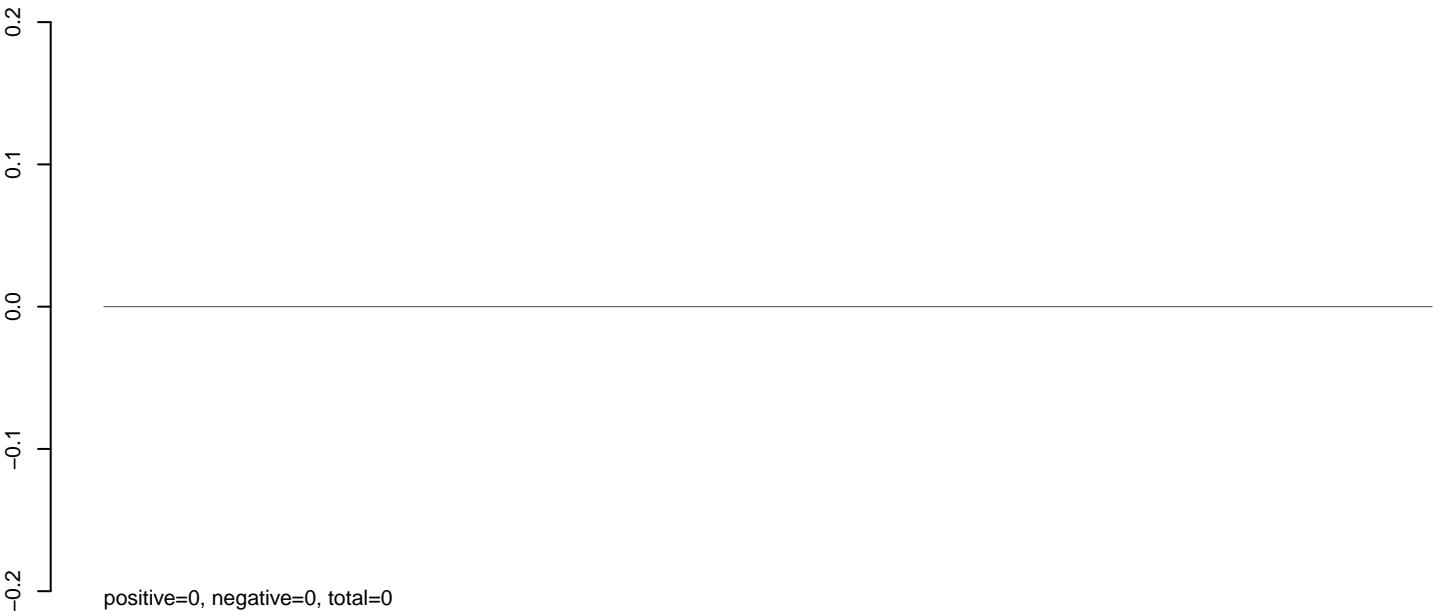


0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



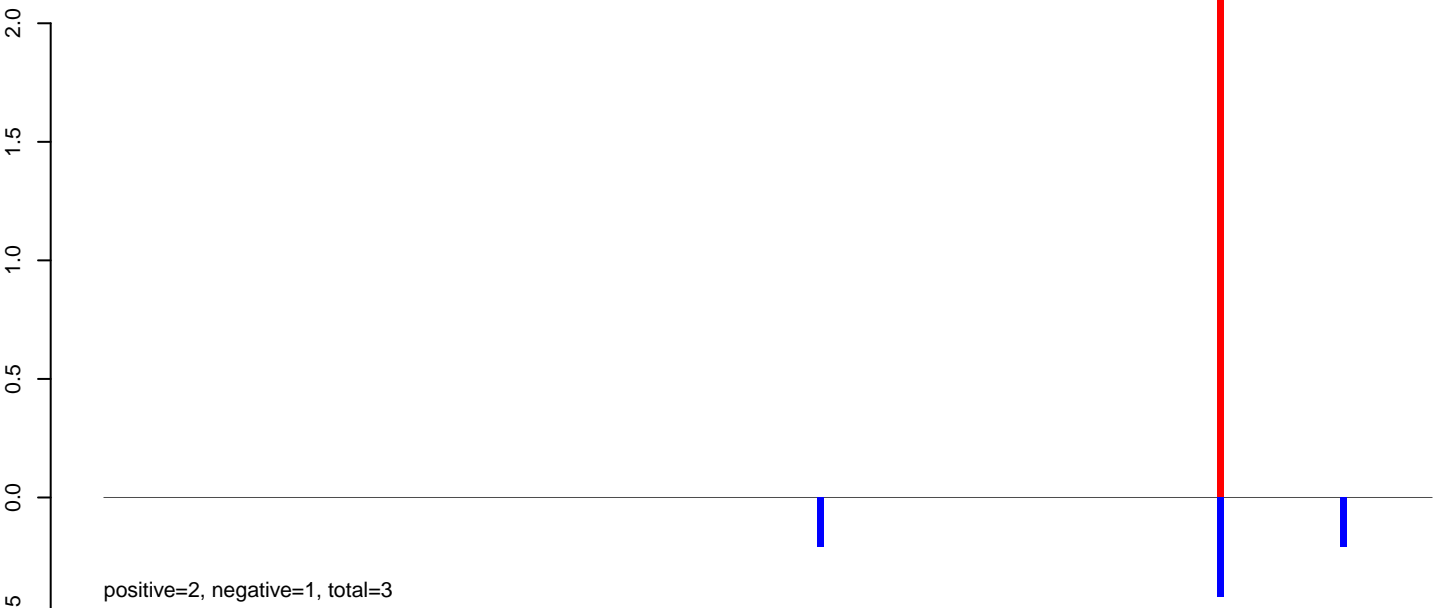
AeAeg_Aag2_DENV4.rep



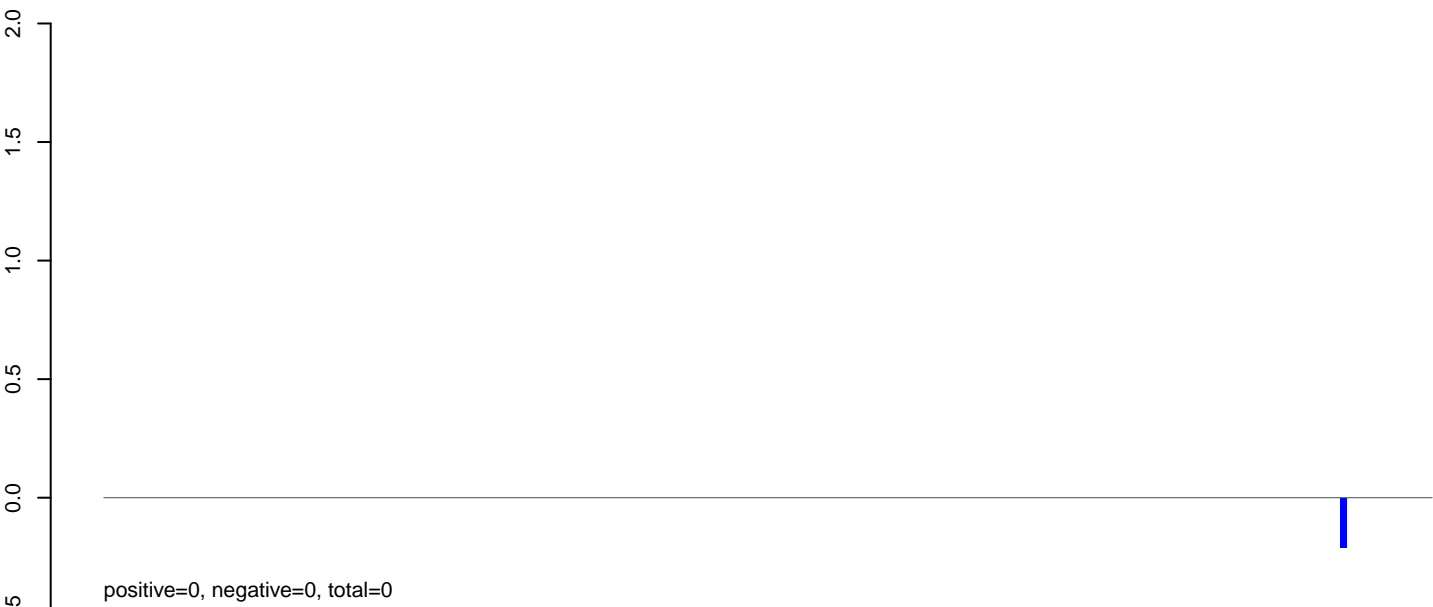
Window size=50, length=11029, virus@NC_009942.1-West_Nile_virus_lineage_1:1-11029

0 2000 4000 6000 8000 10000

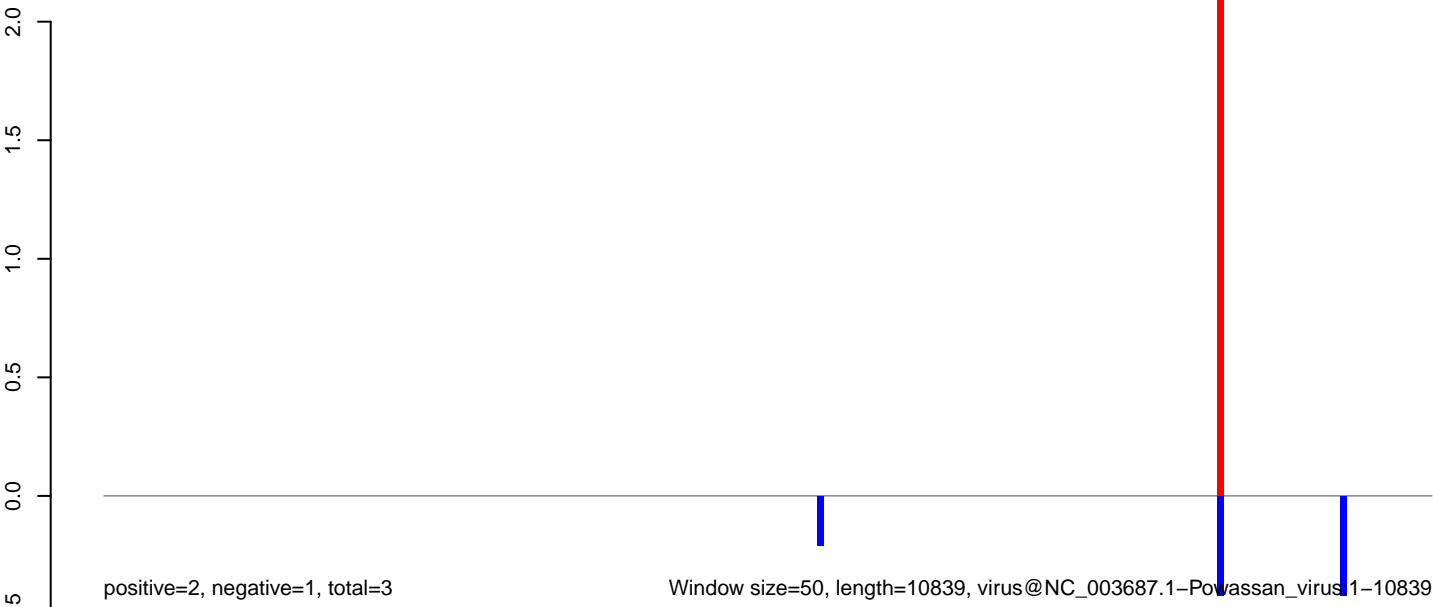
AeAeg_Aag2_DENV4.18_23.rep



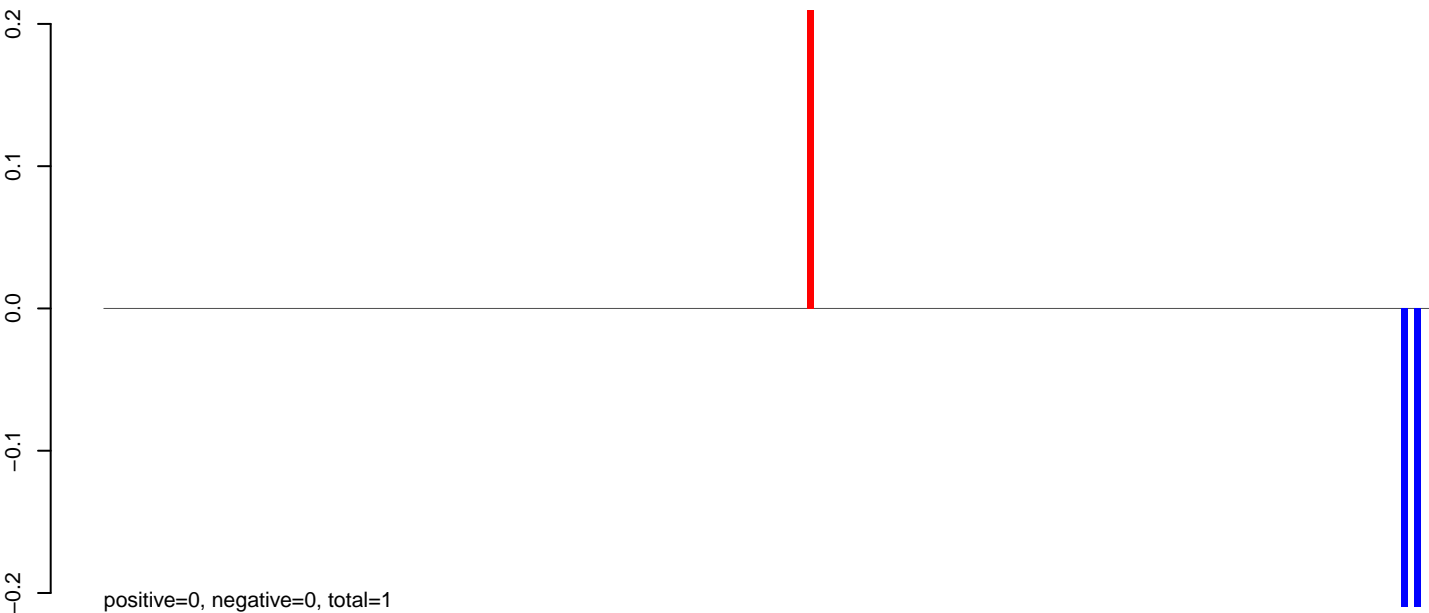
AeAeg_Aag2_DENV4.24_35.rep



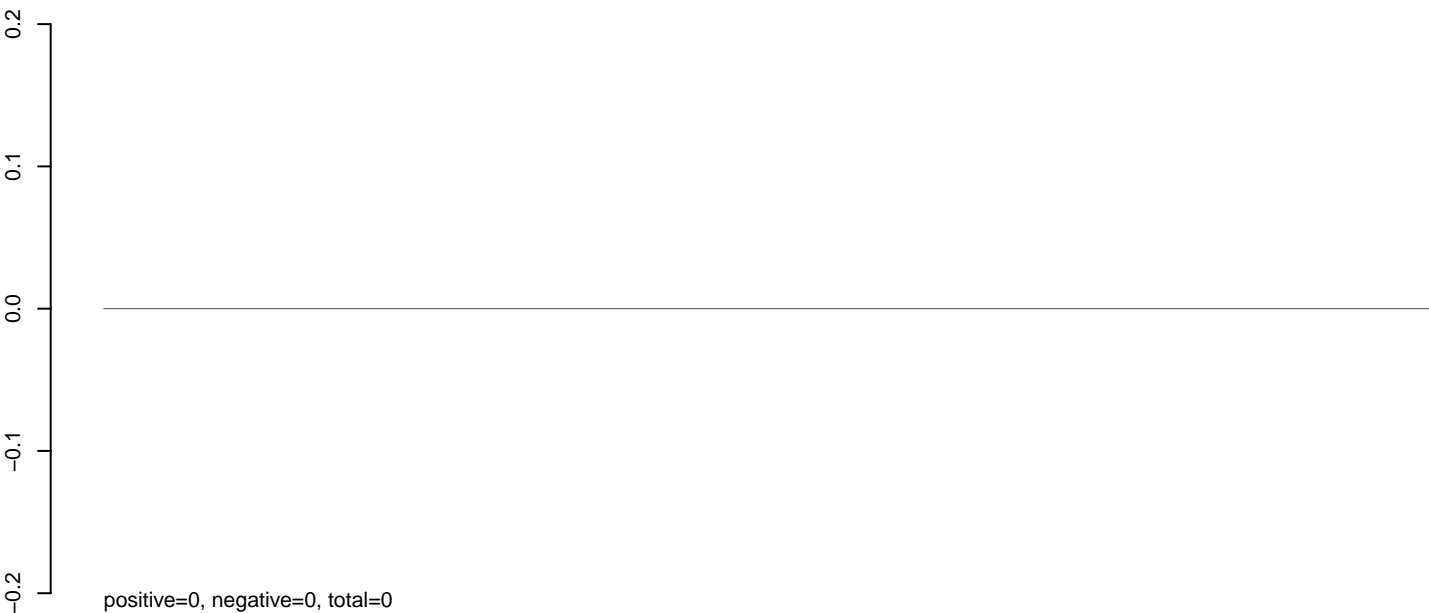
AeAeg_Aag2_DENV4.rep



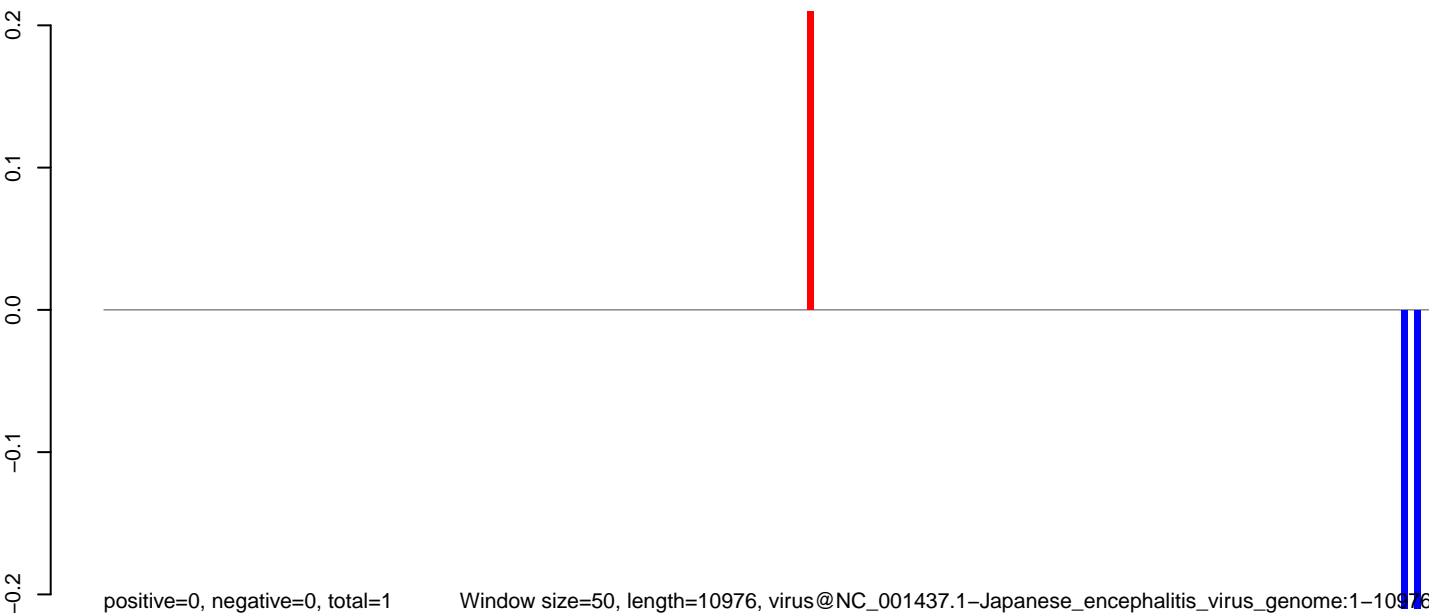
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep

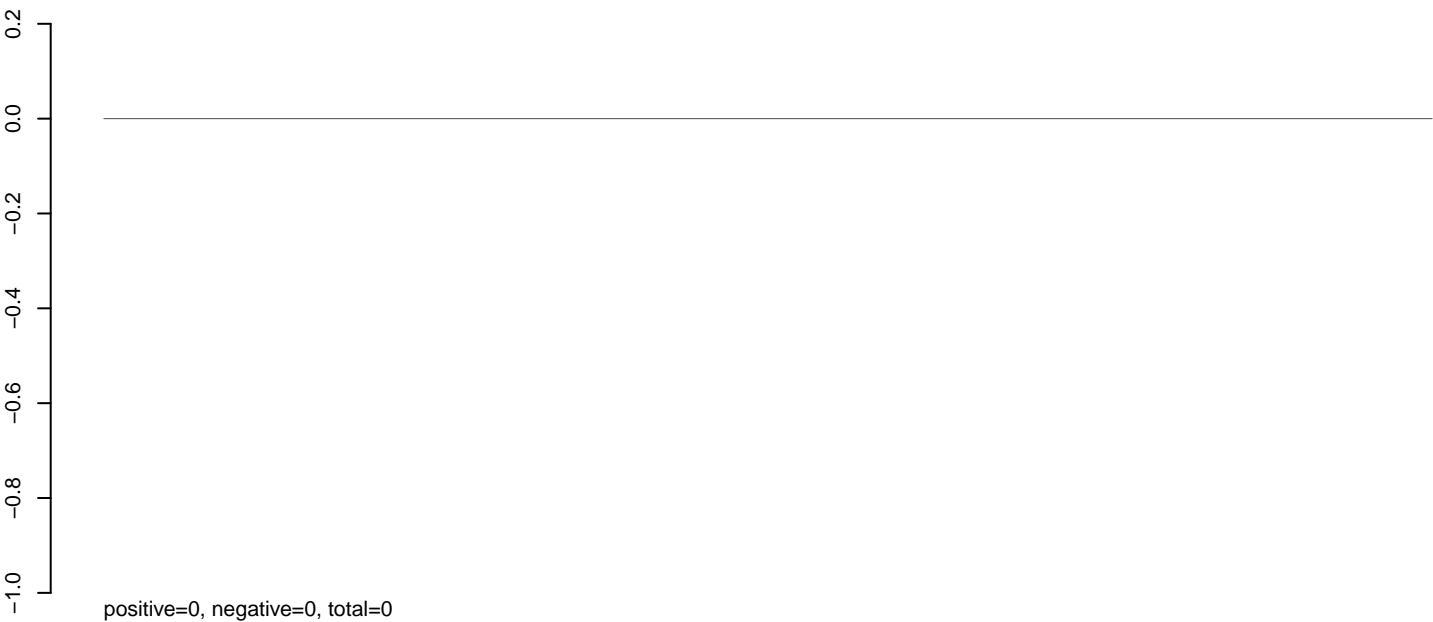


0 2000 4000 6000 8000 10000

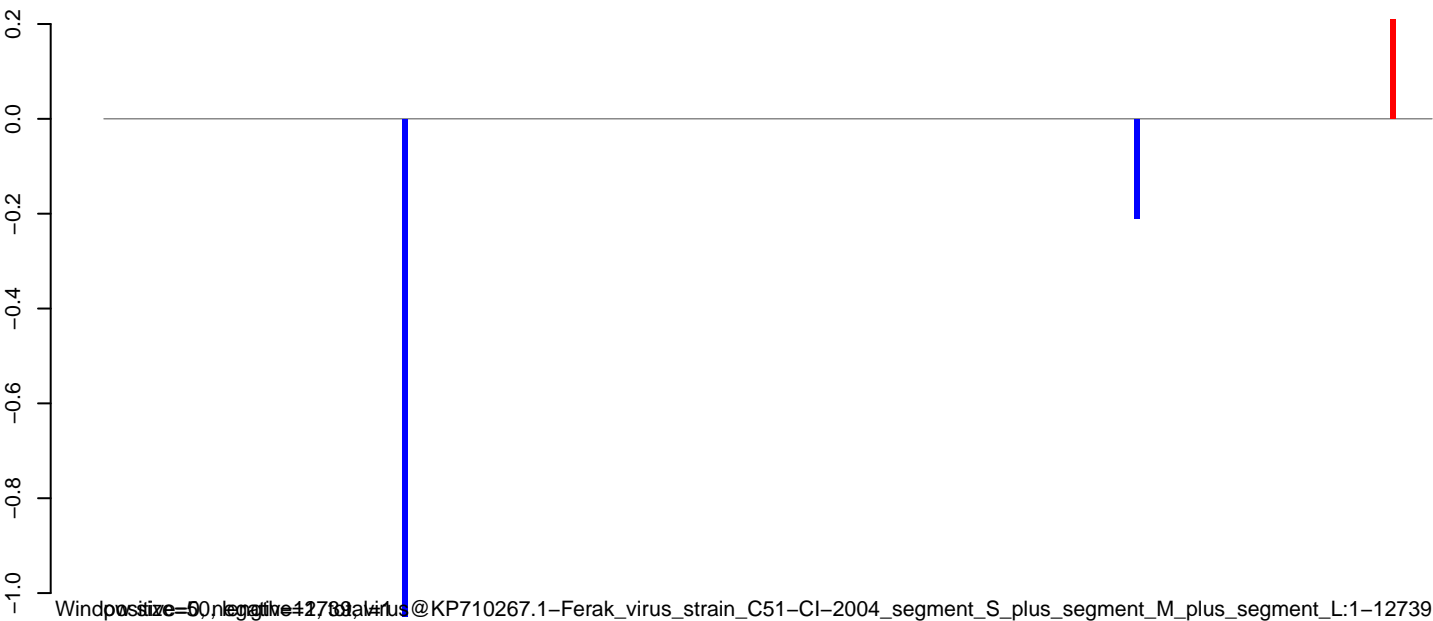
AeAeg_Aag2_DENV4.18_23.rep



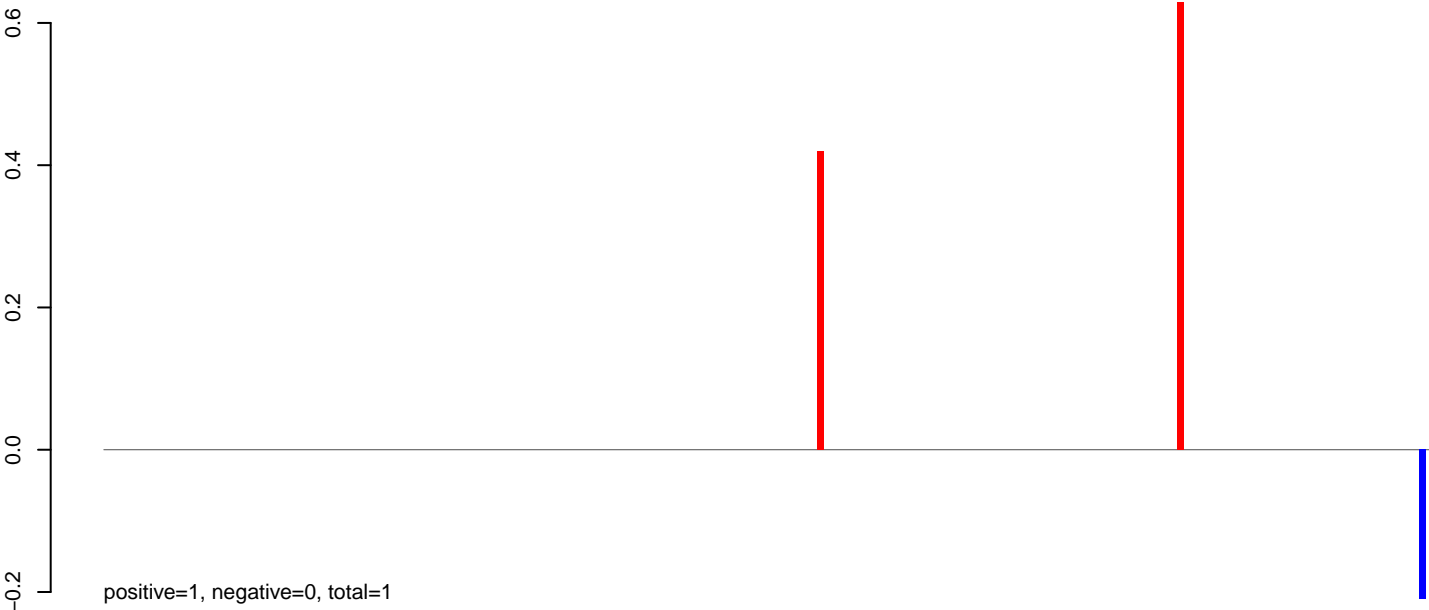
AeAeg_Aag2_DENV4.24_35.rep



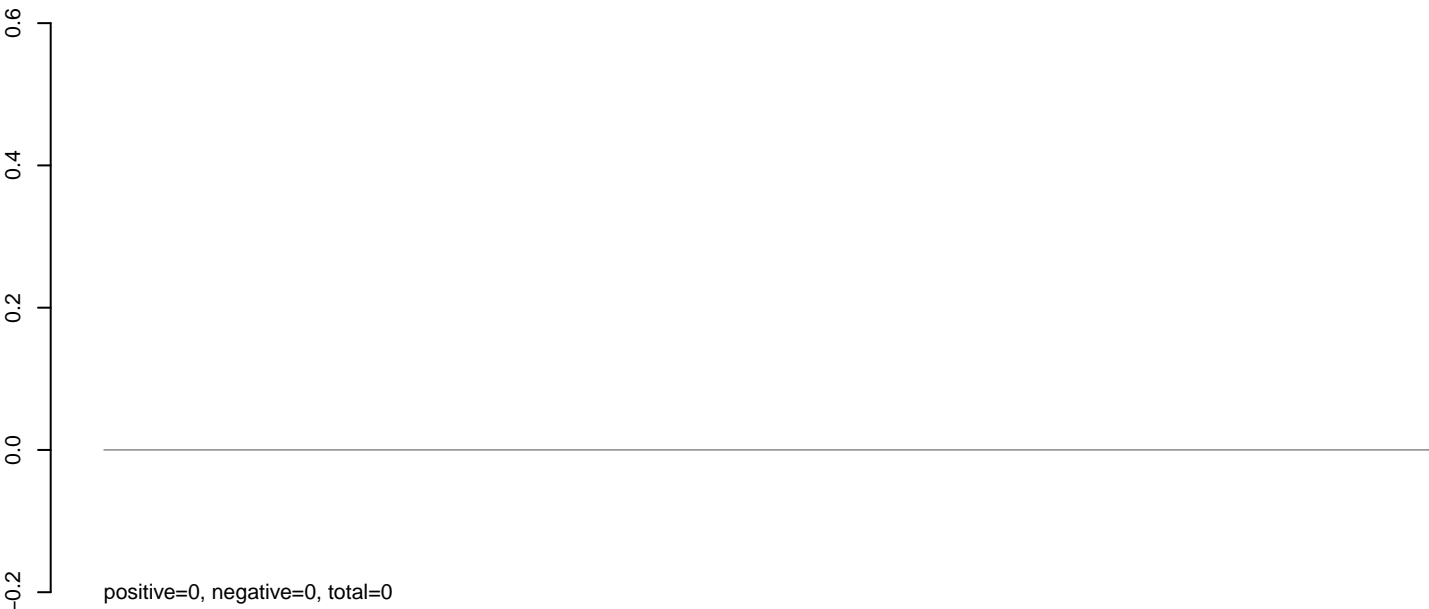
AeAeg_Aag2_DENV4.rep



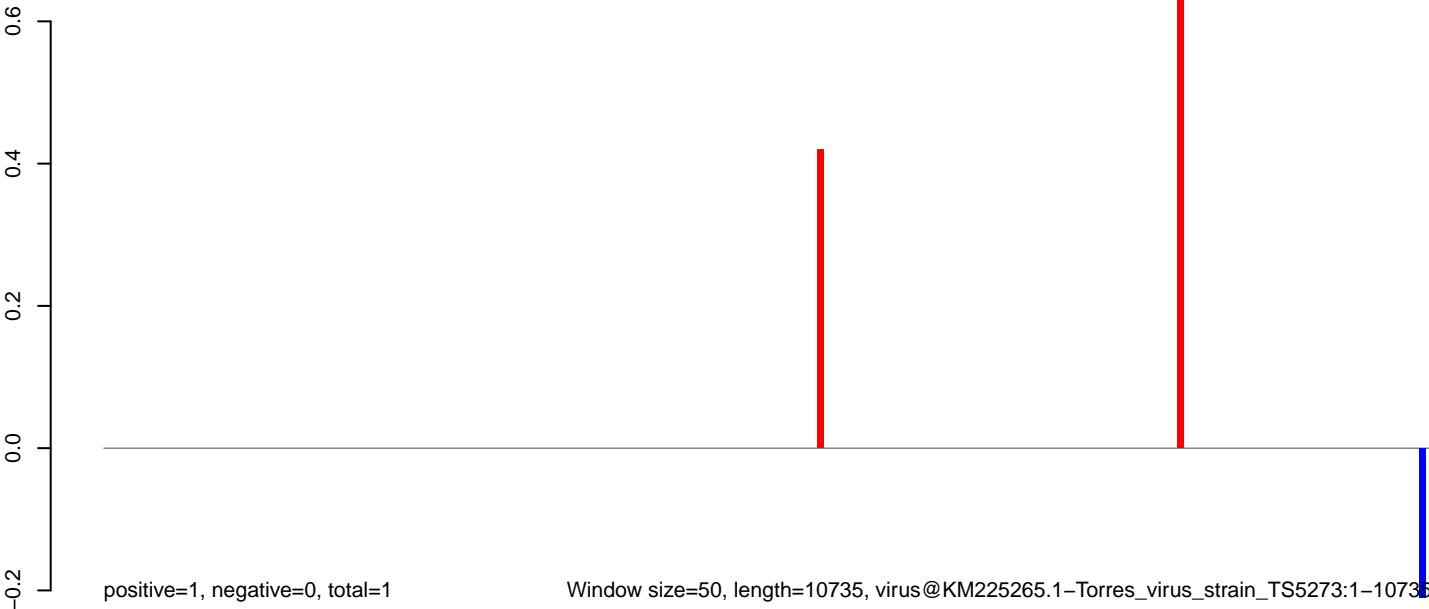
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



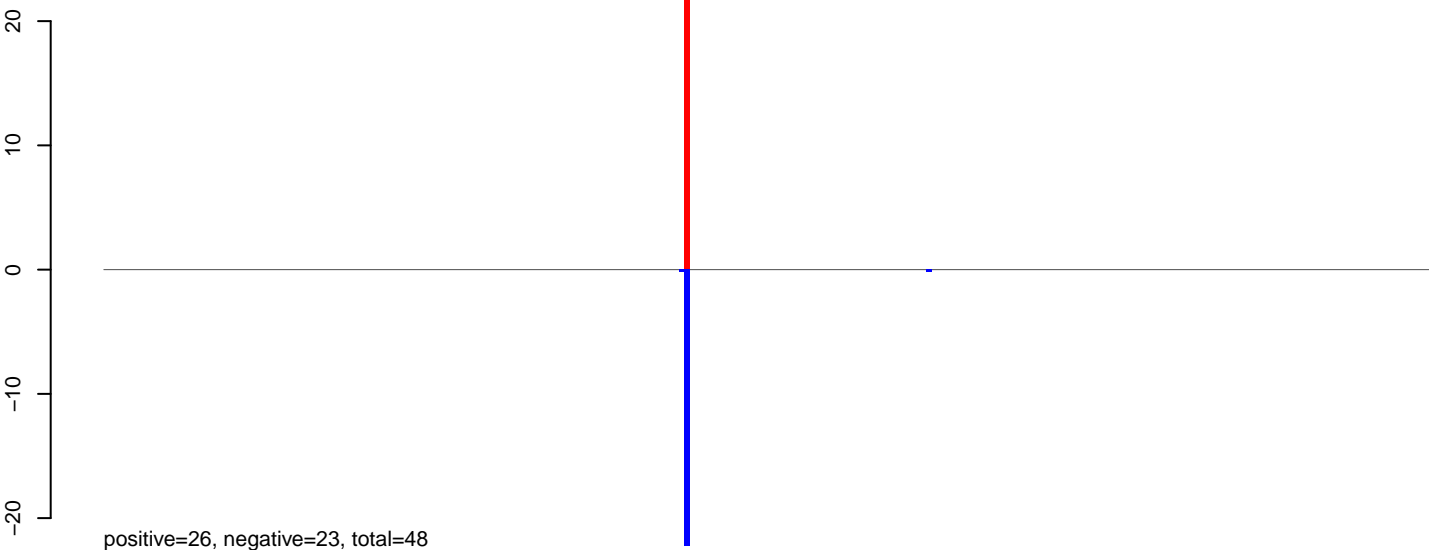
AeAeg_Aag2_DENV4.rep



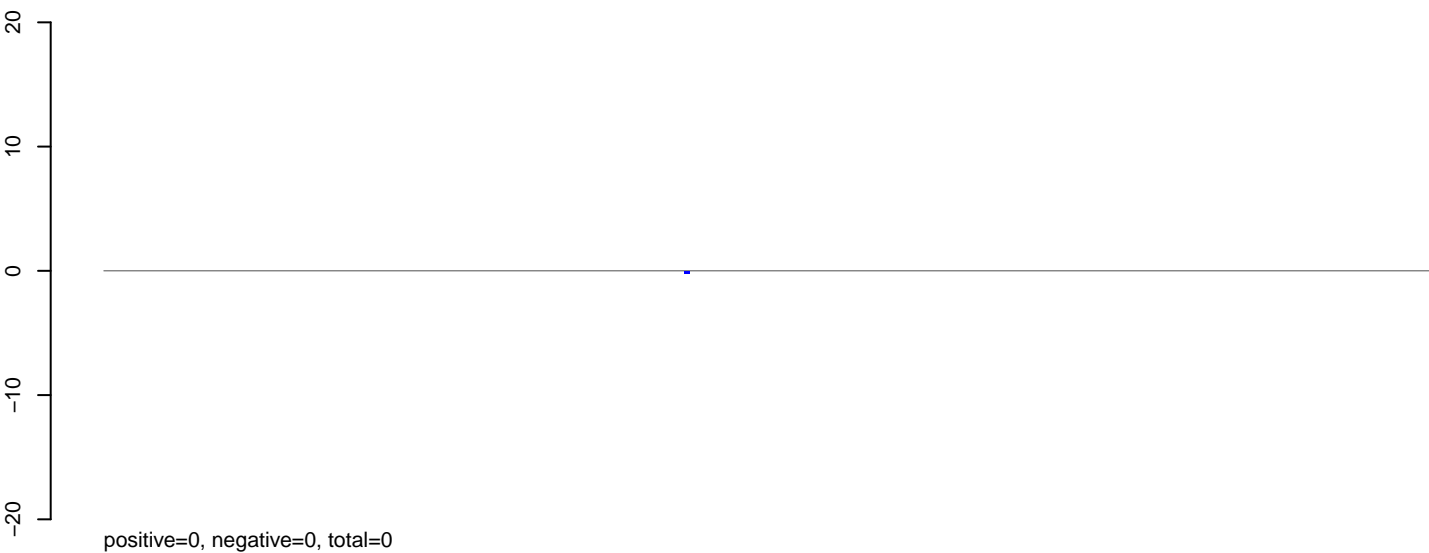
Window size=50, length=10735, virus@KM225265.1-Torres_virus_strain_TS5273:1-10735

0 2000 4000 6000 8000 10000

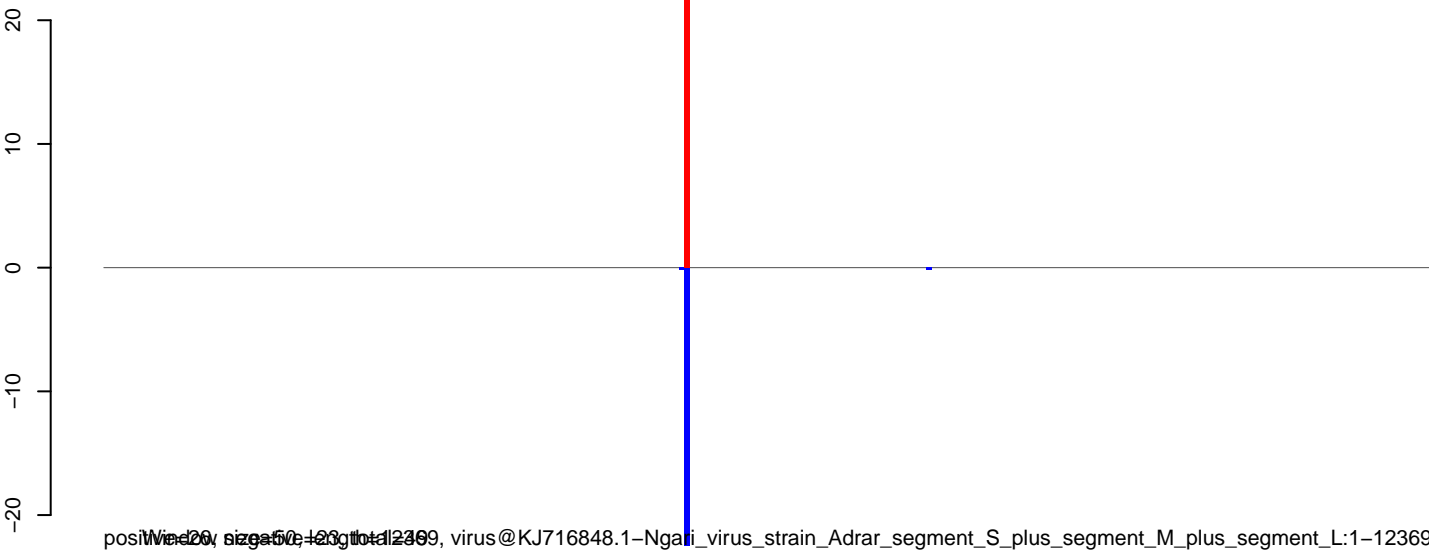
AeAeg_Aag2_DENV4.18_23.rep



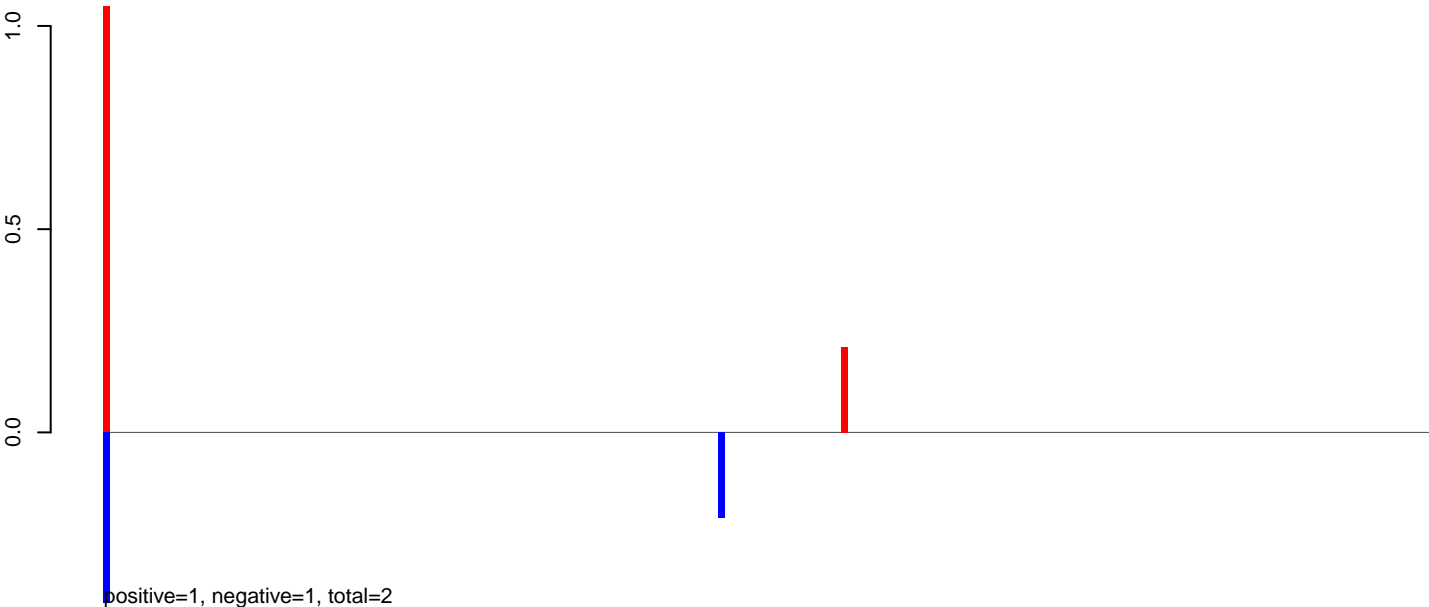
AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



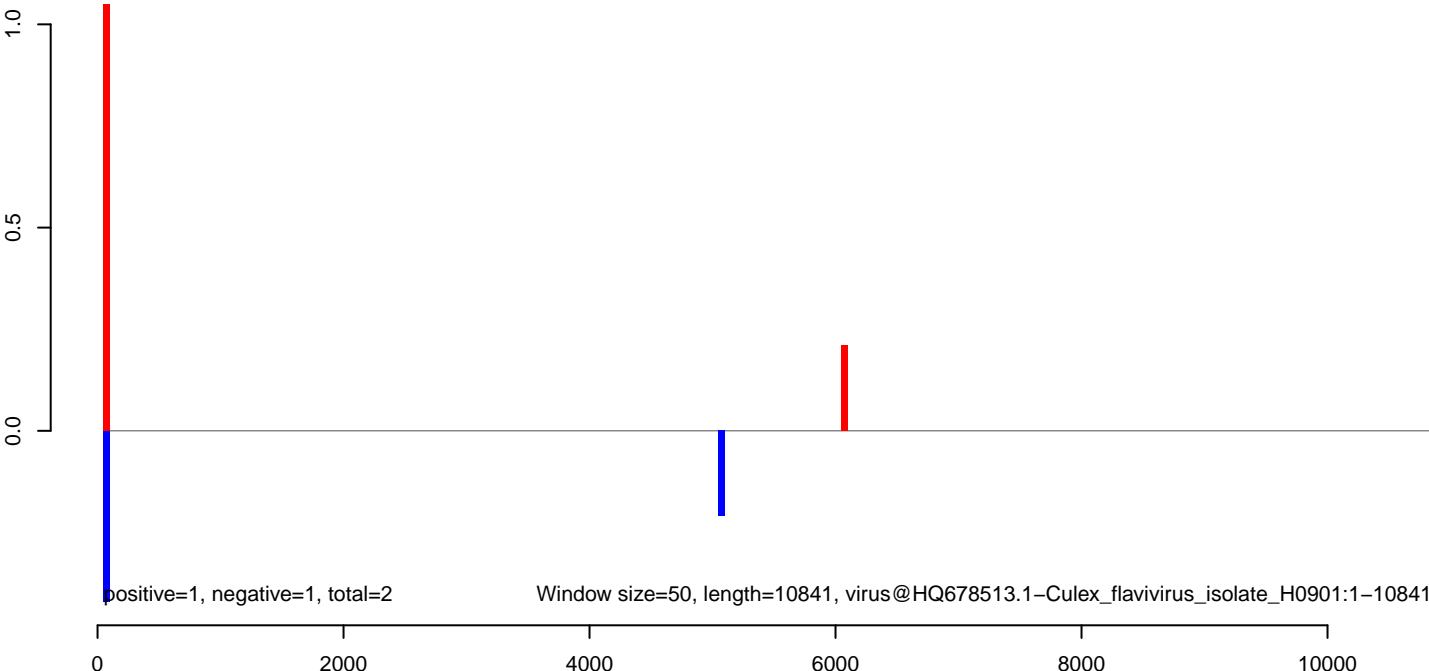
AeAeg_Aag2_DENV4.18_23.rep



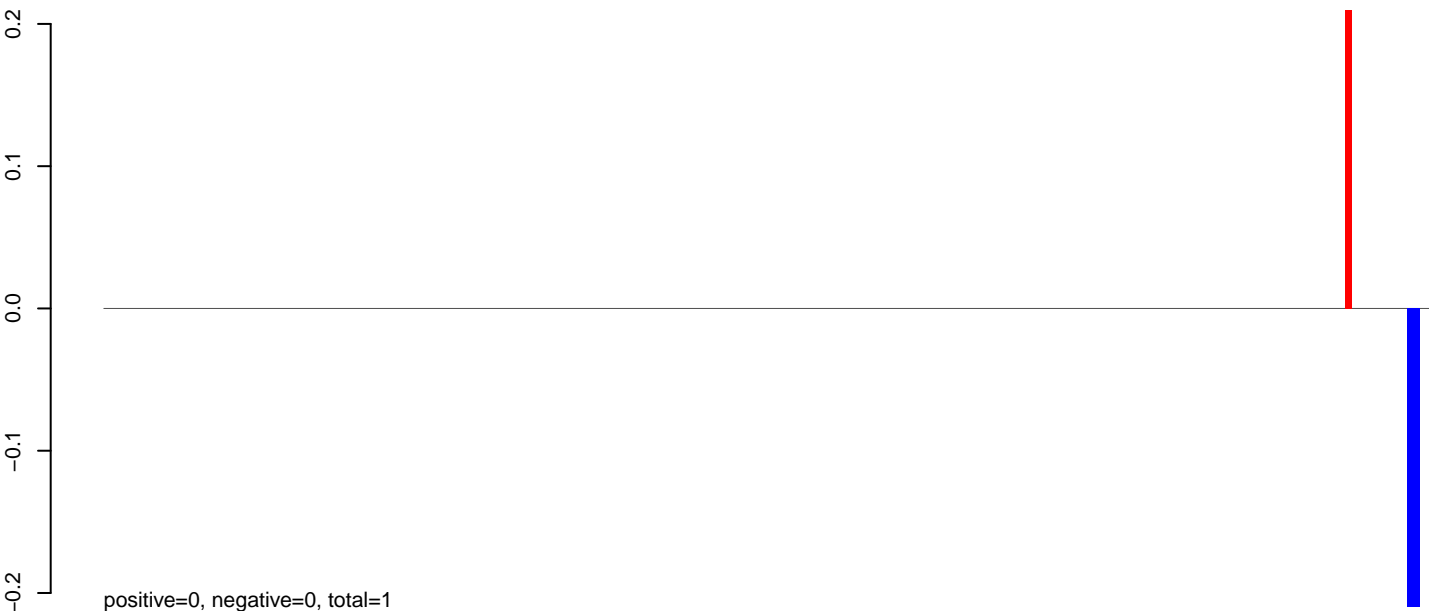
AeAeg_Aag2_DENV4.24_35.rep



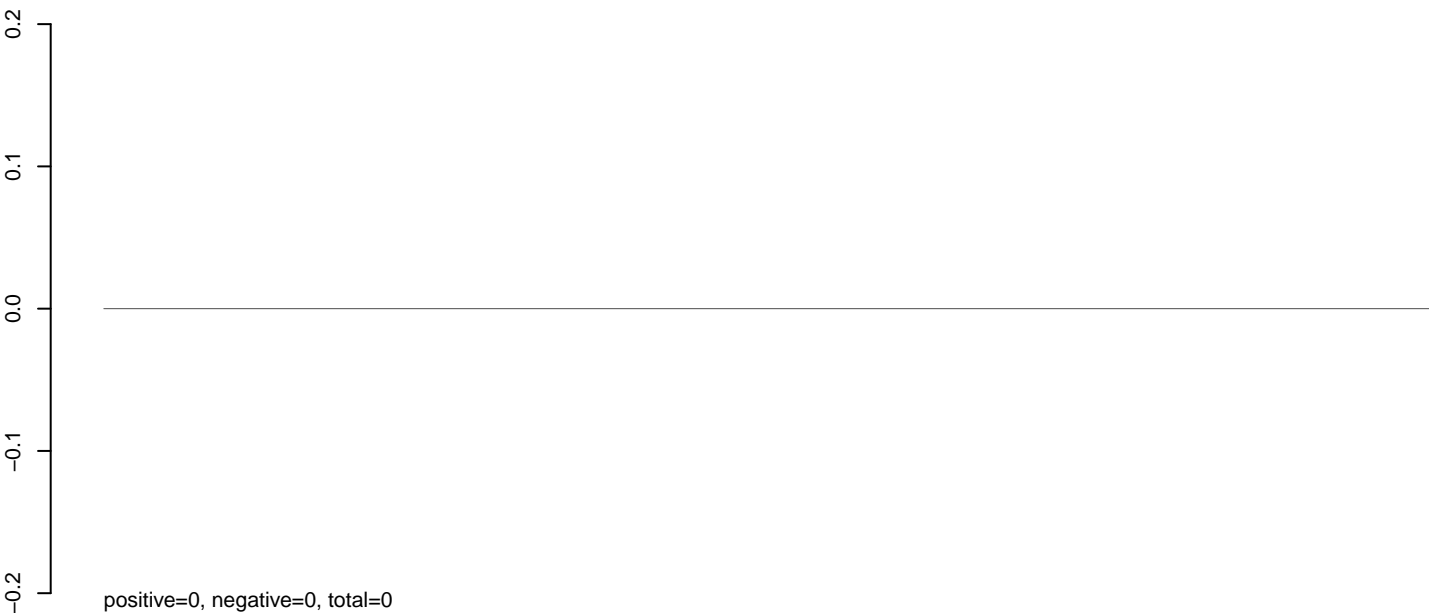
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



Window size=50, length=10794, virus@AY632542.4-Rocio_virus_strain_SPH_34675:1-10794

0 2000 4000 6000 8000 10000

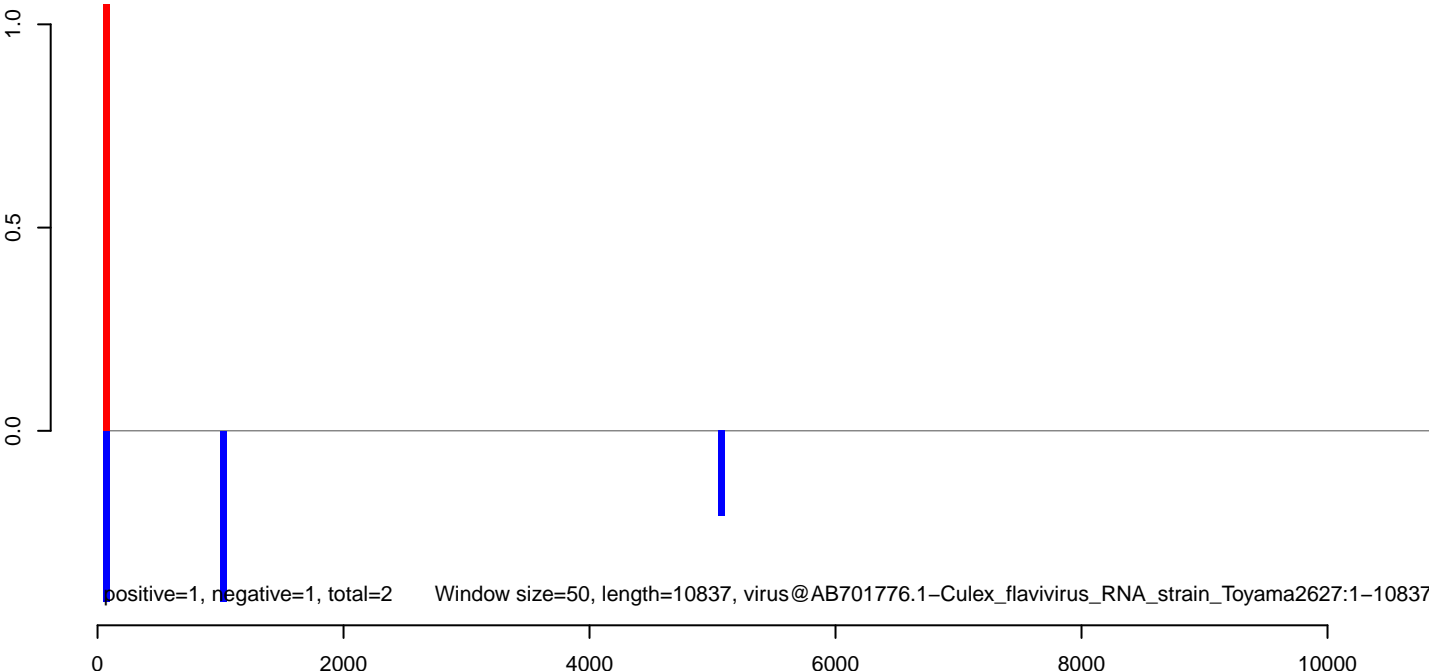
AeAeg_Aag2_DENV4.18_23.rep



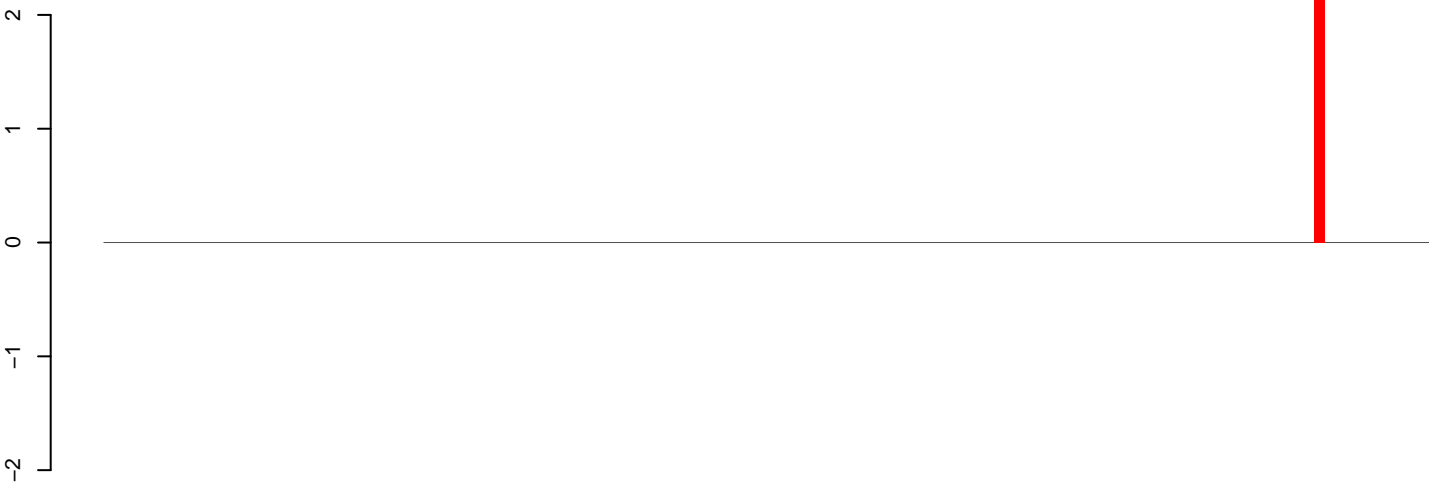
AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep

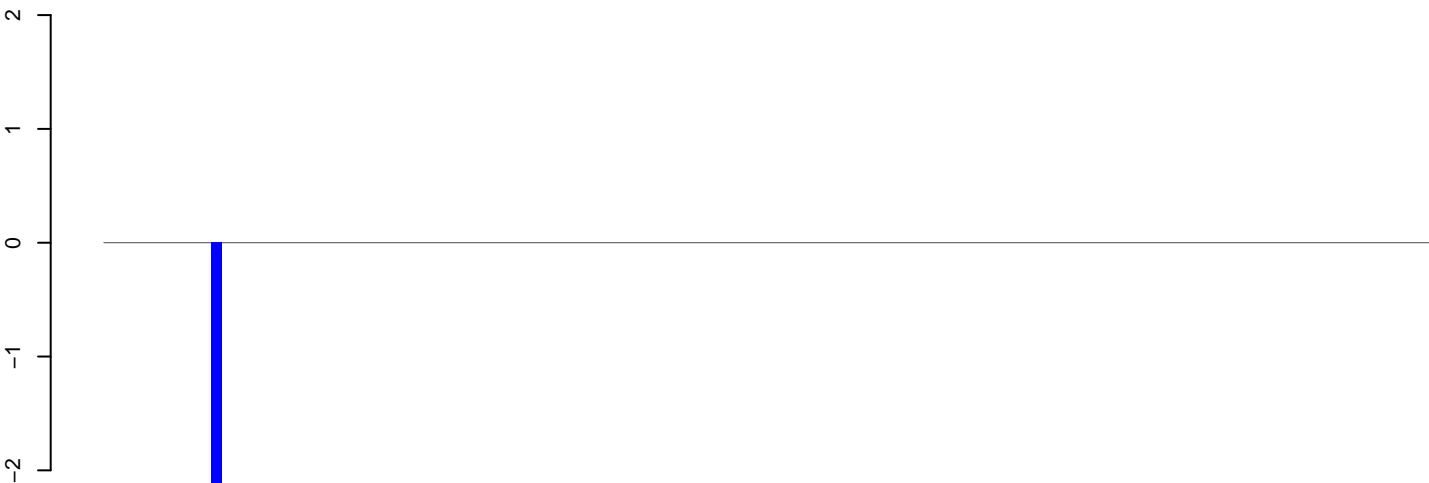


AeAeg_Aag2_DENV4.18_23.rep



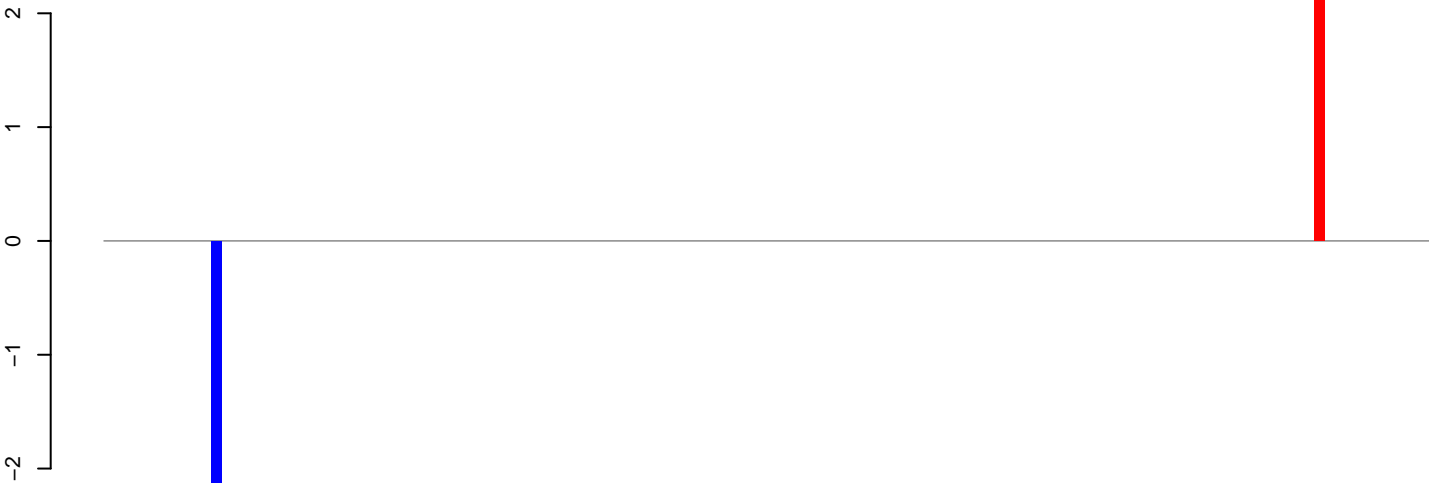
positive=3, negative=0, total=3

AeAeg_Aag2_DENV4.24_35.rep



positive=0, negative=3, total=3

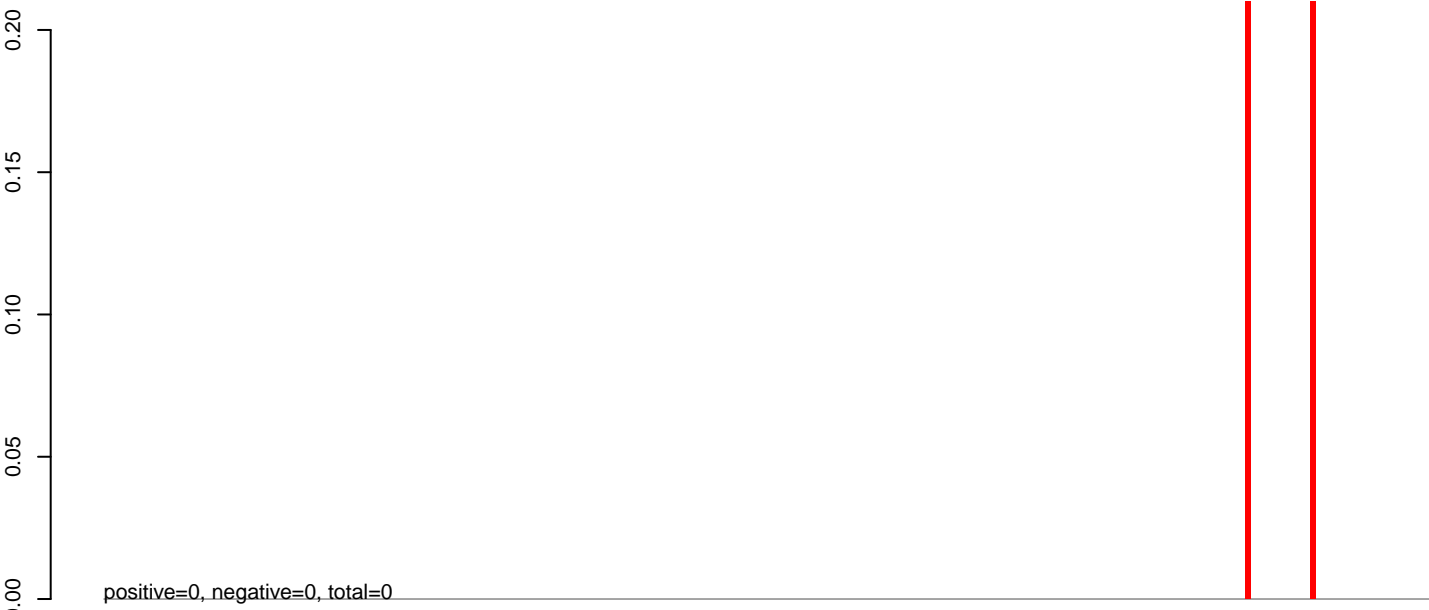
AeAeg_Aag2_DENV4.rep



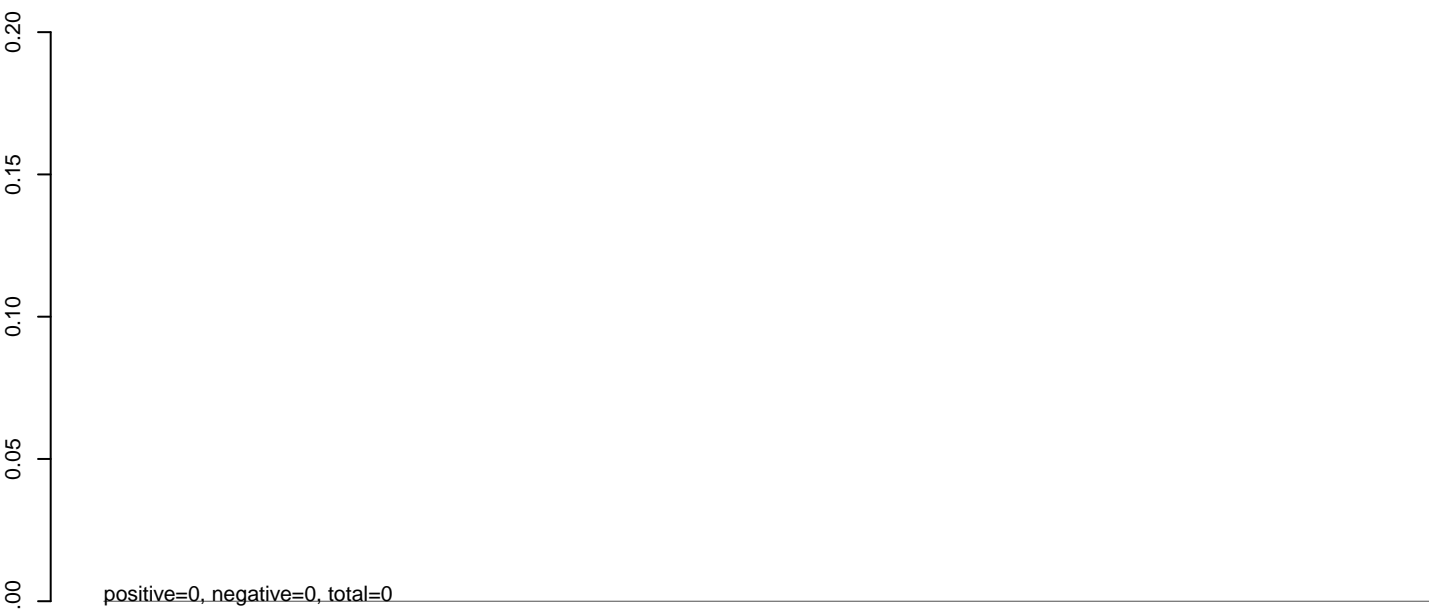
positive=3, negative=3, total=6, window size=50, length=6203, virus@NC_035674.1-Australian_Anopheles_totivirus_isolate_AATV_150840:1-6203

0 1000 2000 3000 4000 5000 6000

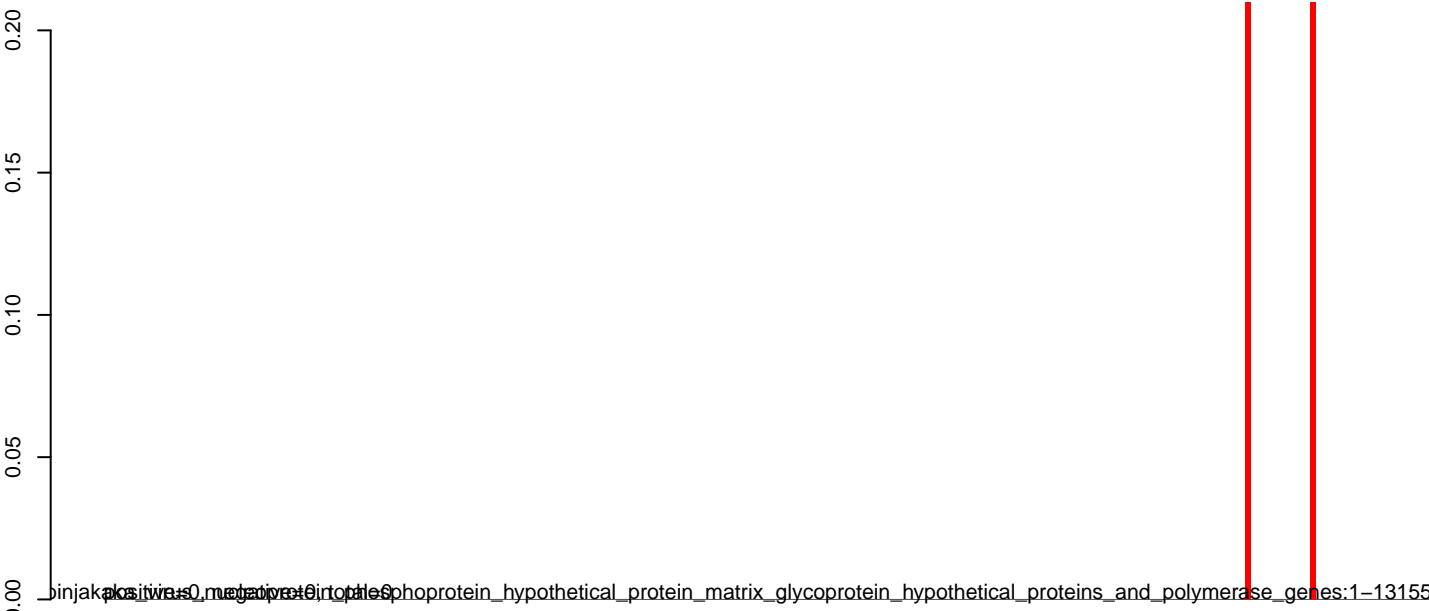
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



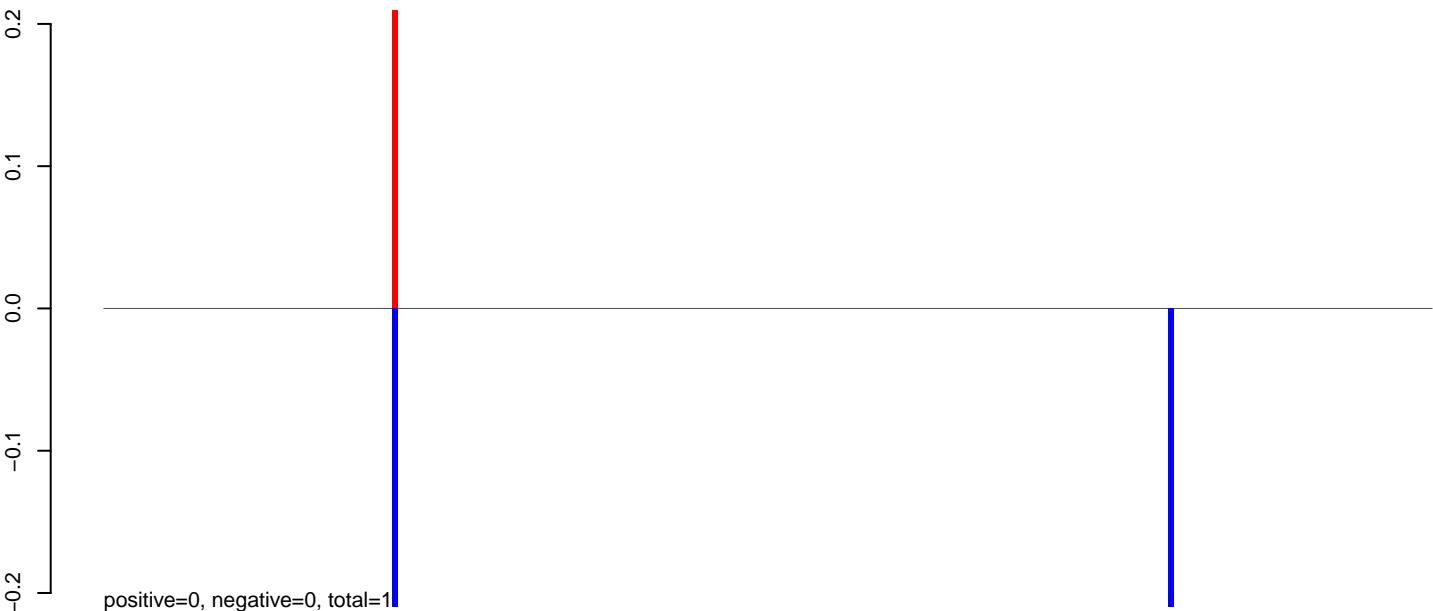
AeAeg_Aag2_DENV4.rep



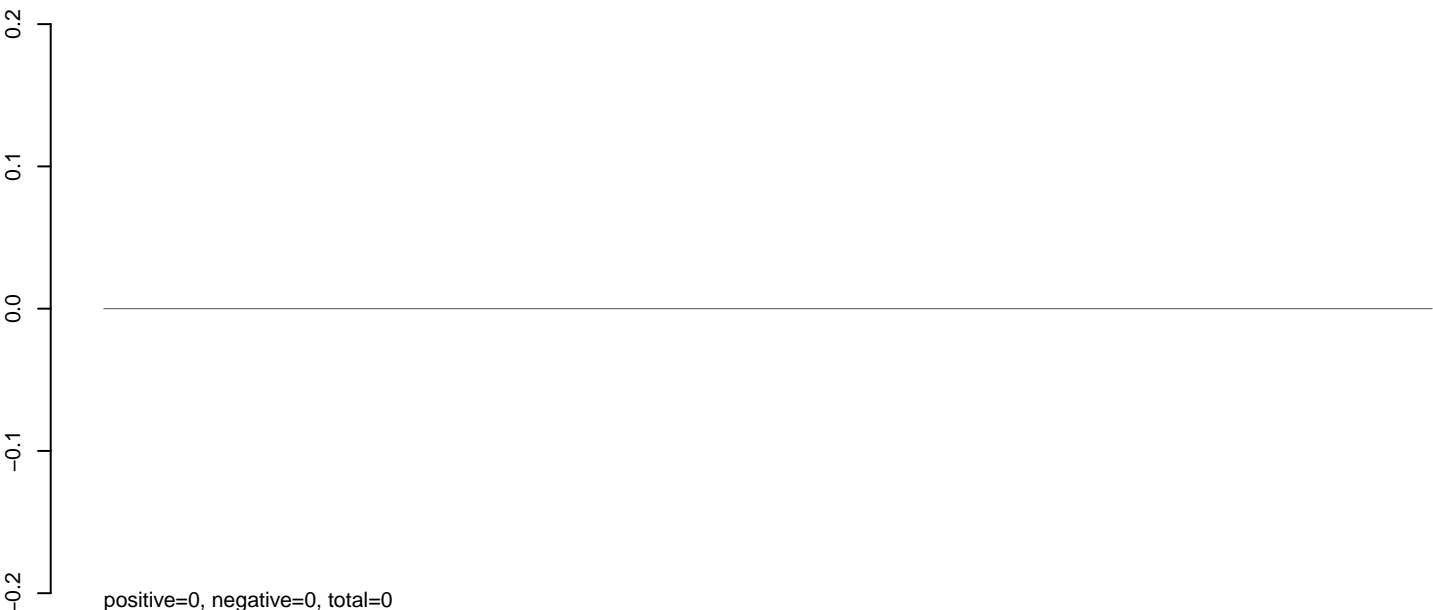
binjakas positive negative total phosphoprotein_hypothetical_protein_matrix_glycoprotein_hypothetical_proteins_and_polymerase_genes:1-13155

0 2000 4000 6000 8000 10000 12000

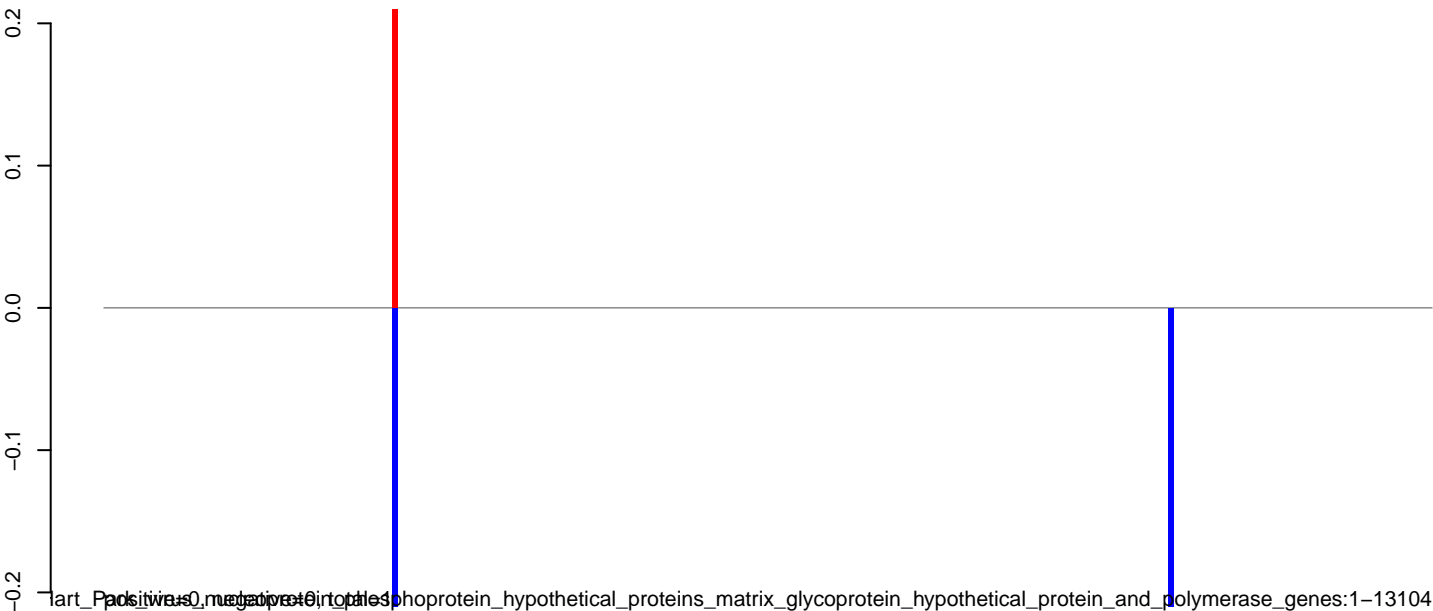
AeAeg_Aag2_DENV4.18_23.rep



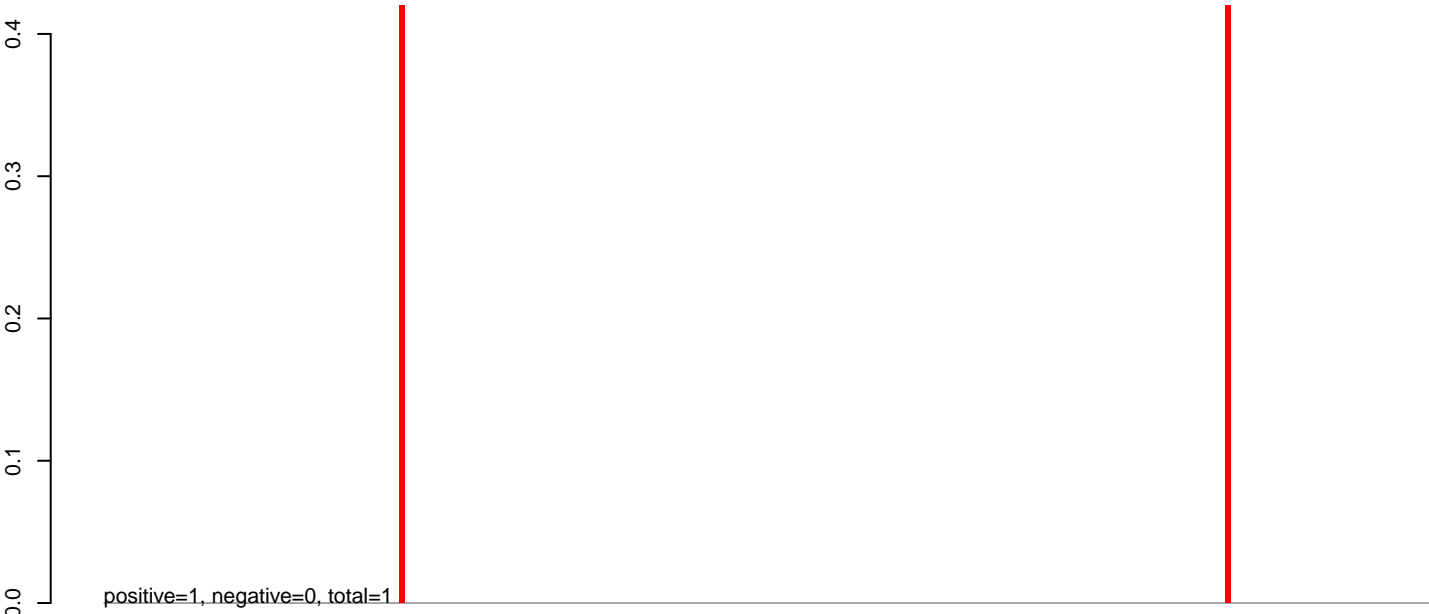
AeAeg_Aag2_DENV4.24_35.rep



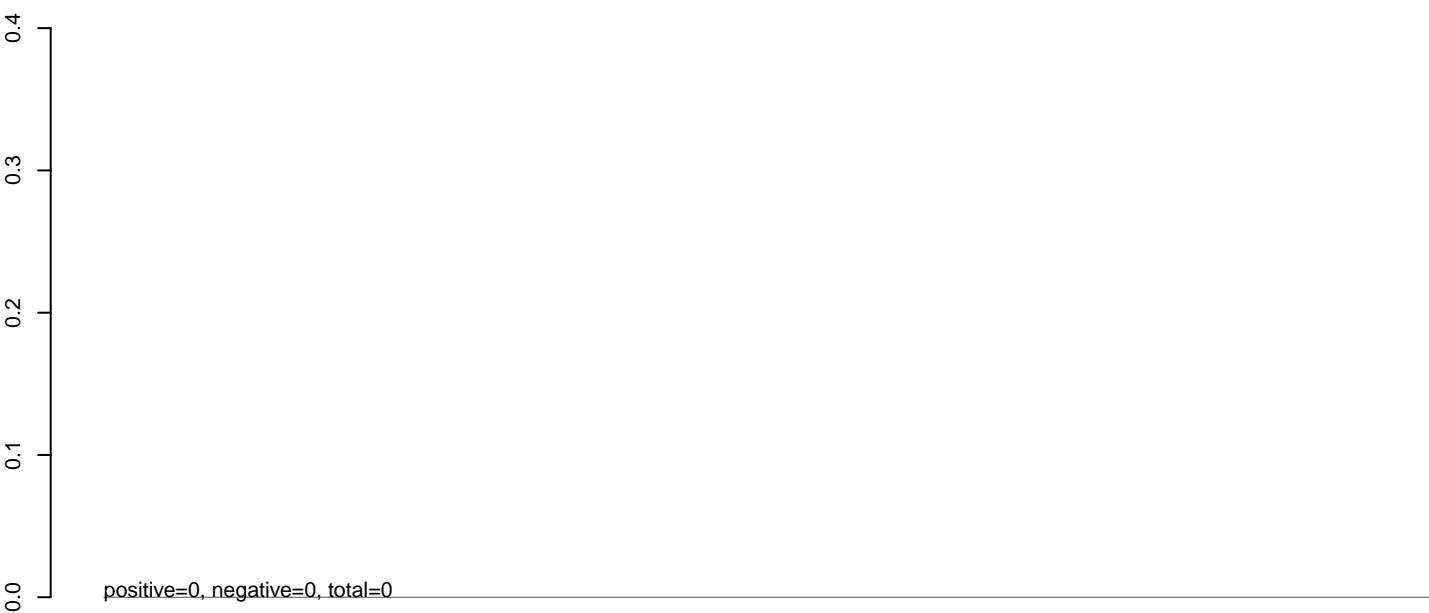
AeAeg_Aag2_DENV4.rep



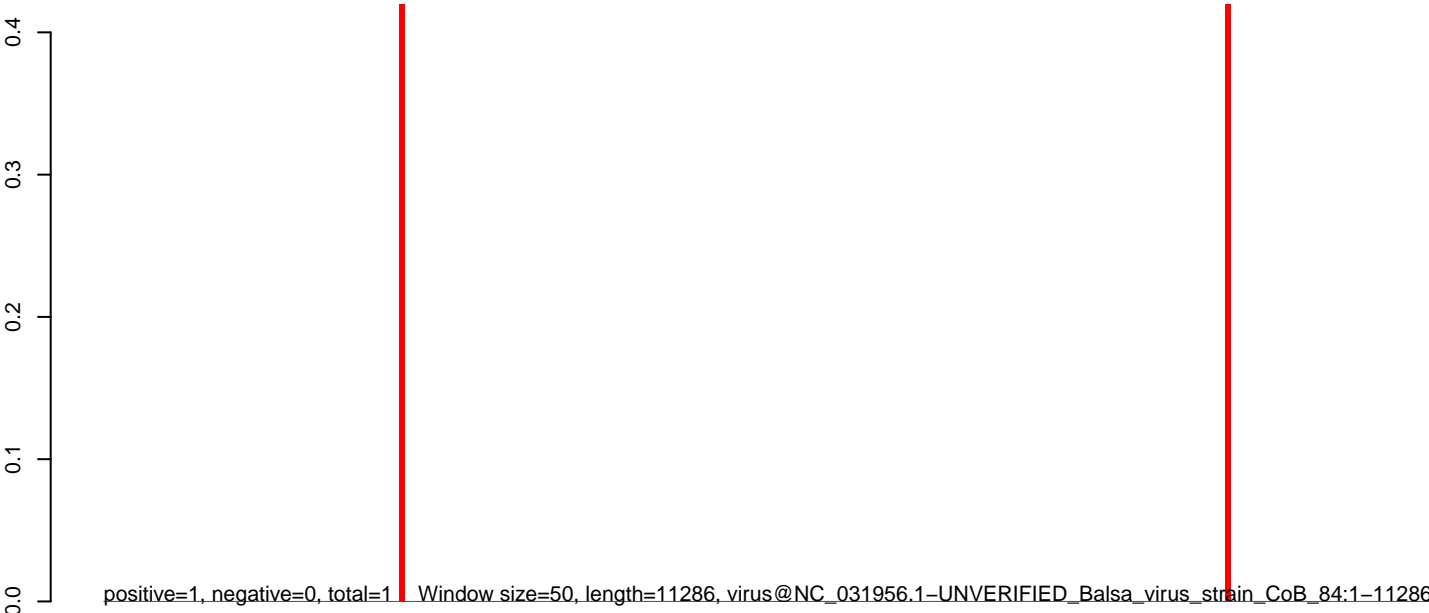
AeAeg_Aag2_DENV4.18_23.rep



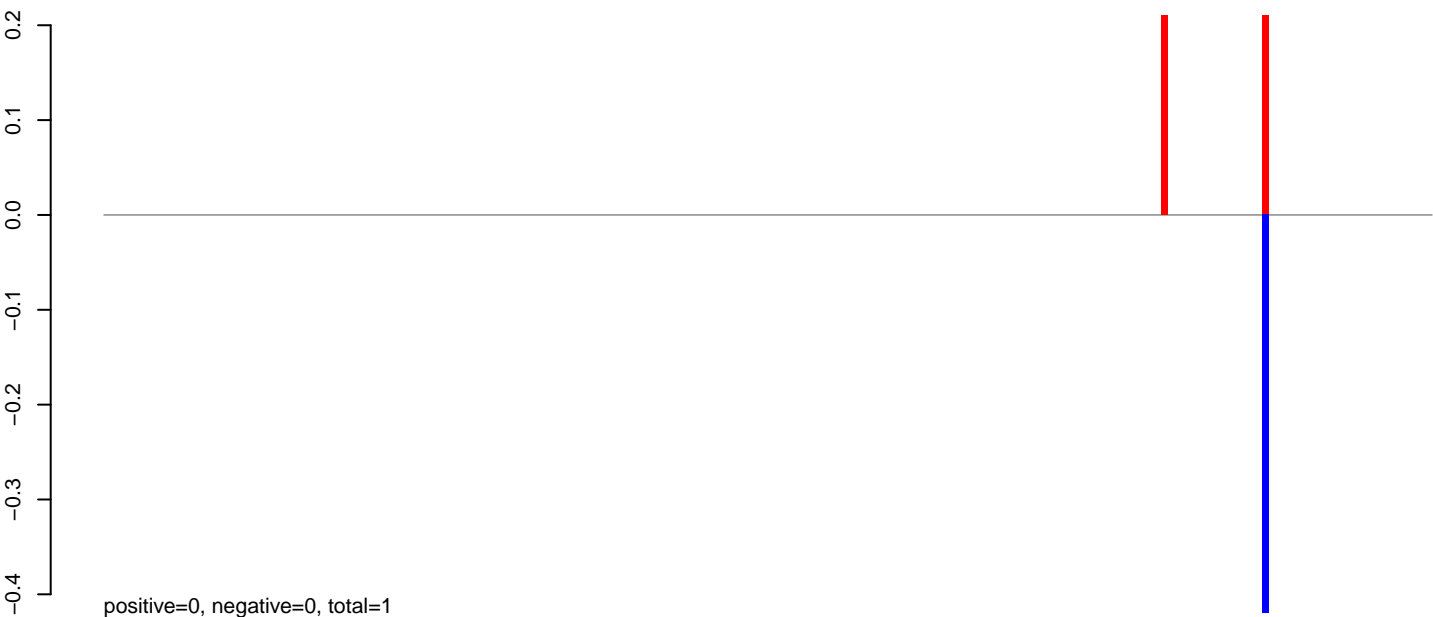
AeAeg_Aag2_DENV4.24_35.rep



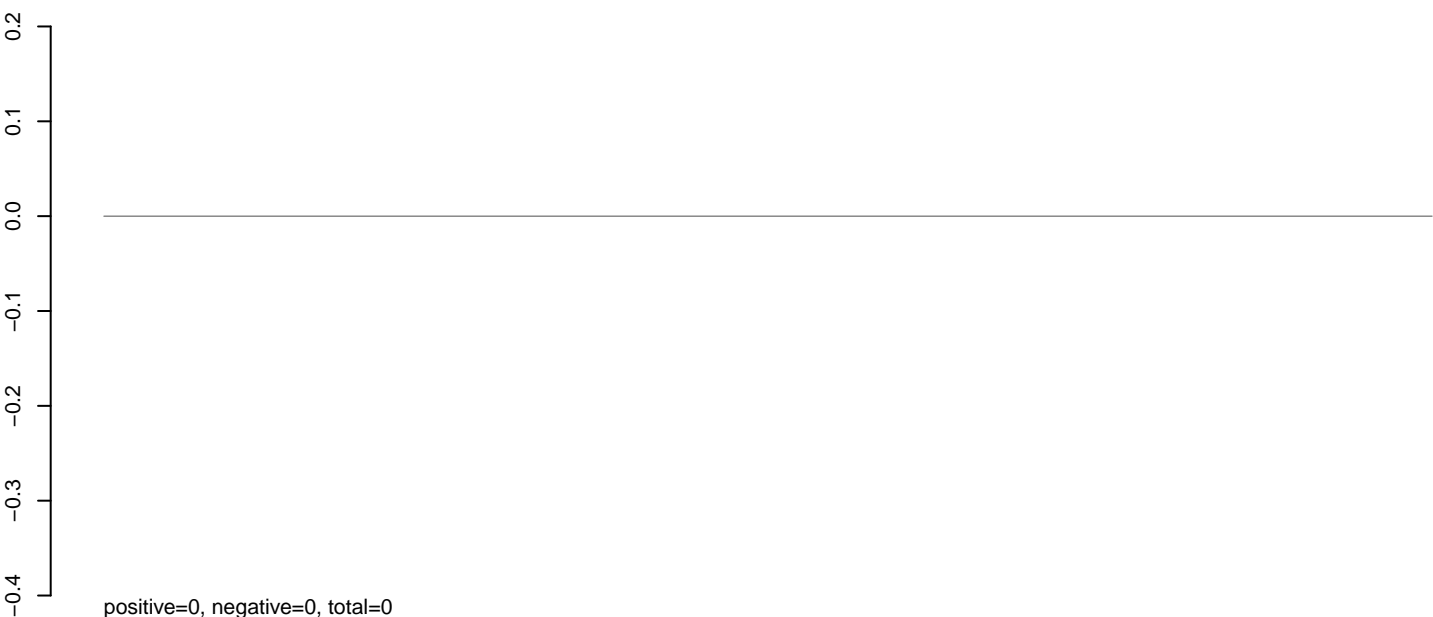
AeAeg_Aag2_DENV4.rep



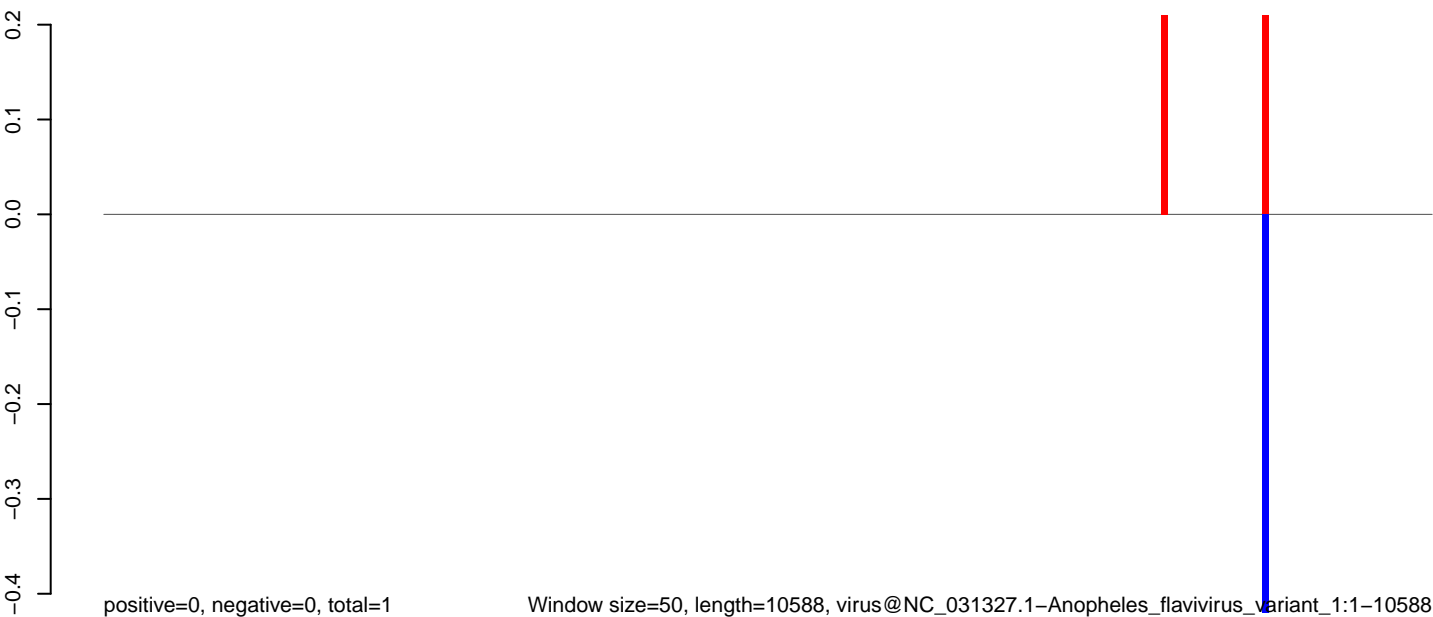
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



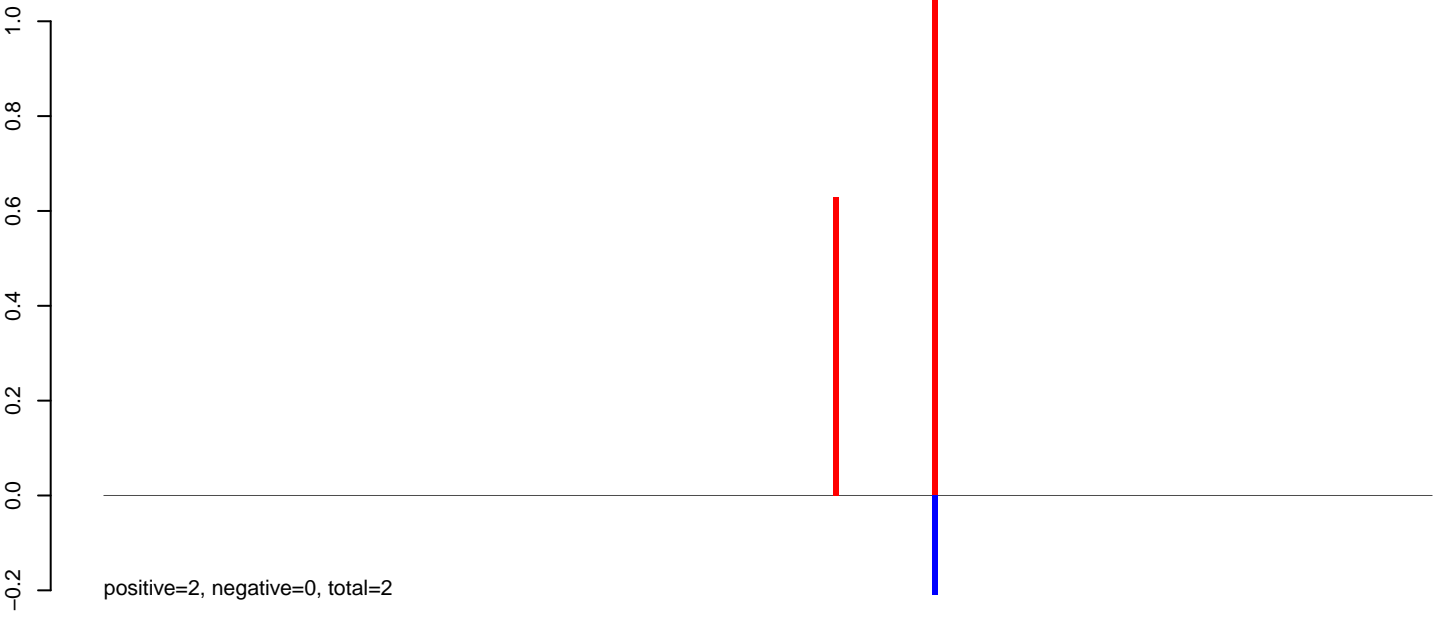
AeAeg_Aag2_DENV4.rep



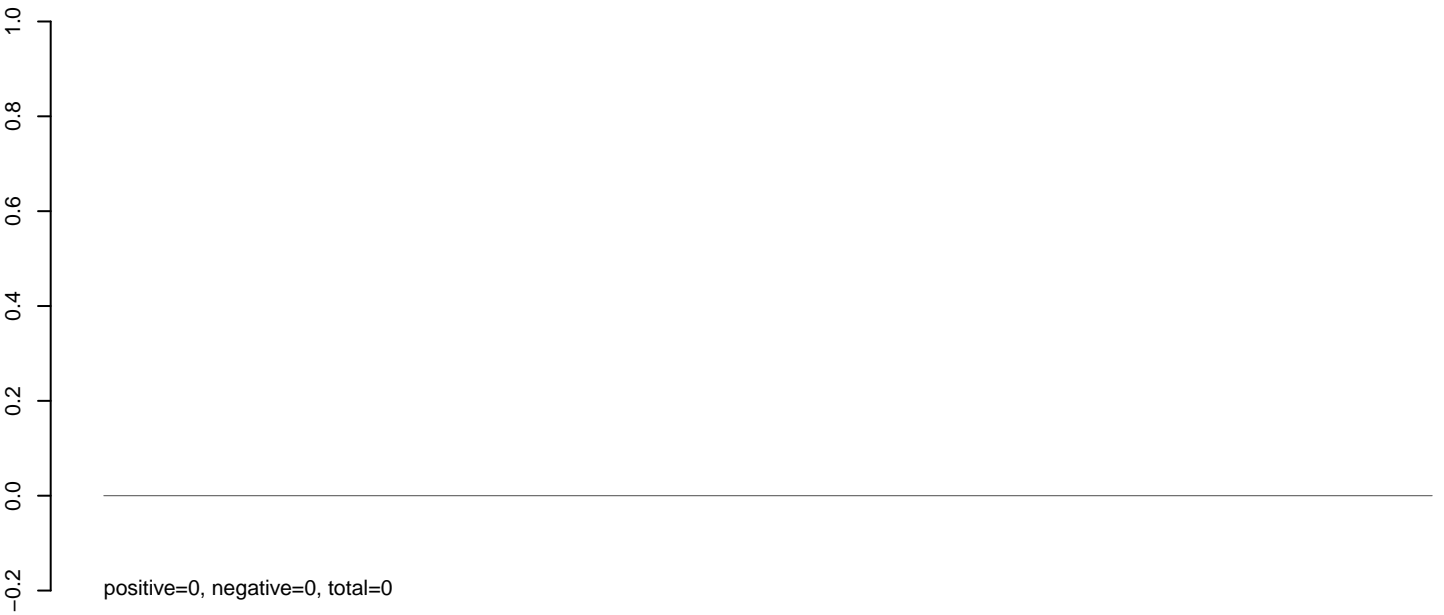
Window size=50, length=10588, virus@NC_031327.1-Anopheles_flavivirus_variant_1:1-10588

0 2000 4000 6000 8000 10000

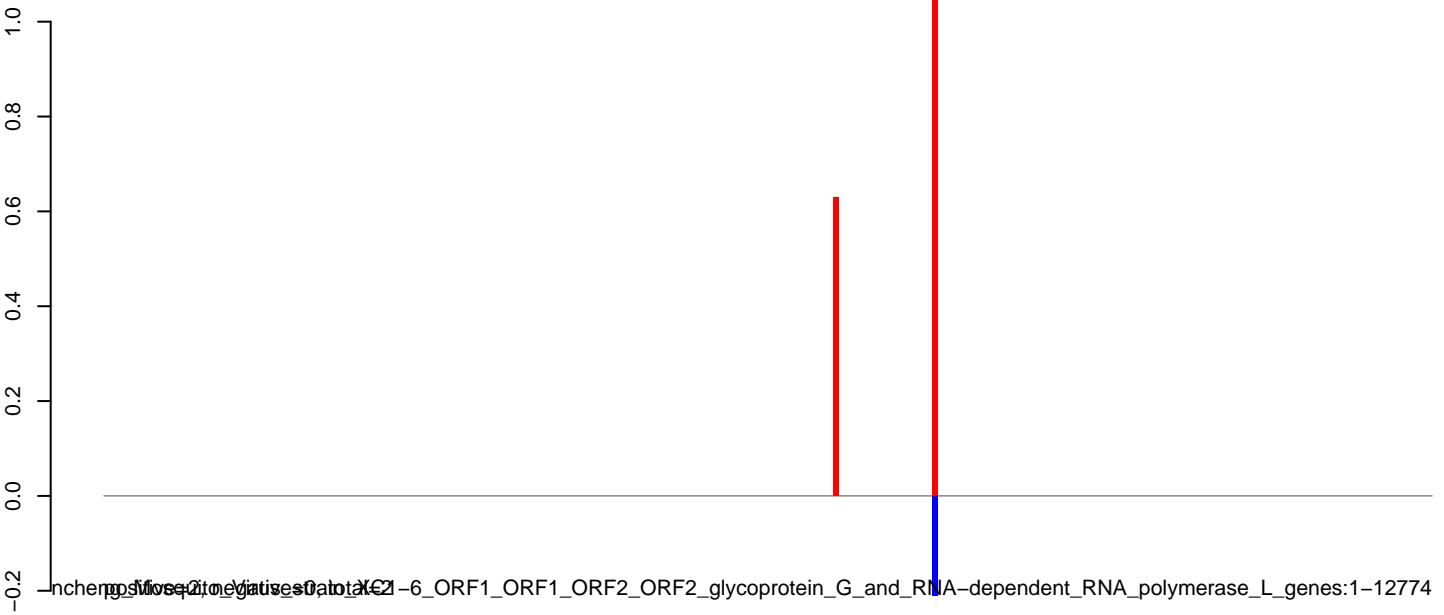
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

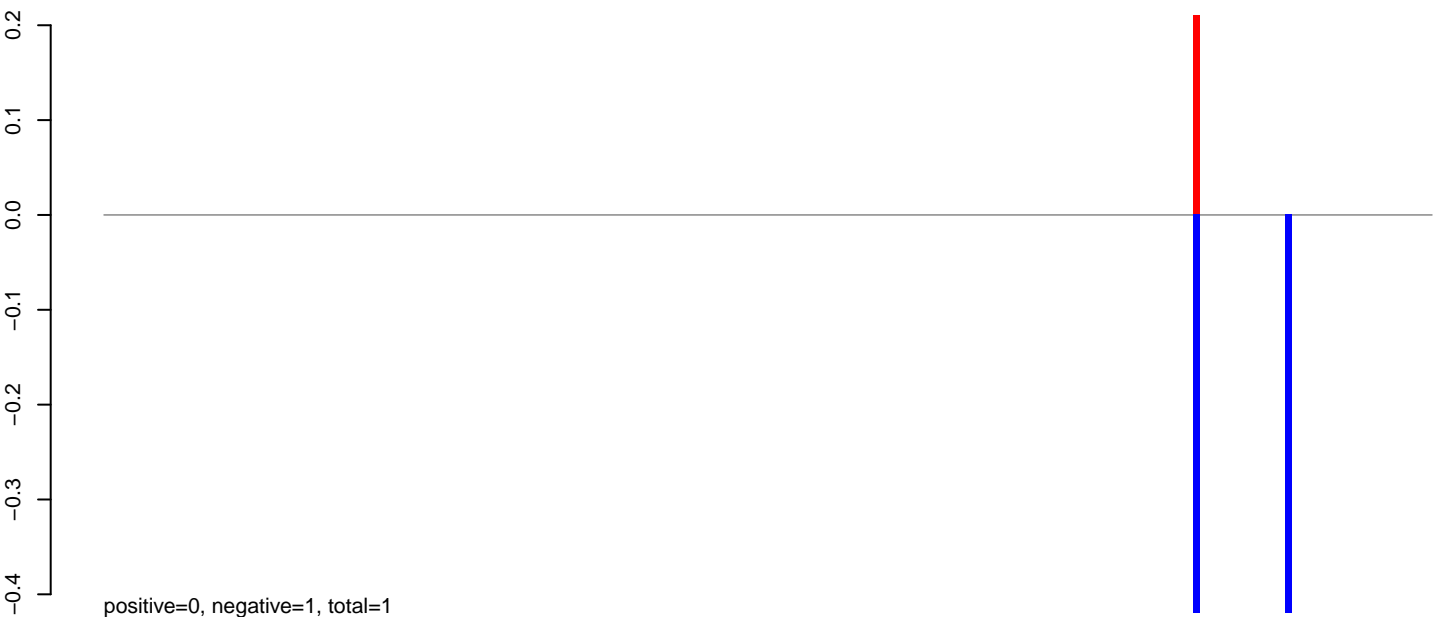


AeAeg_Aag2_DENV4.rep

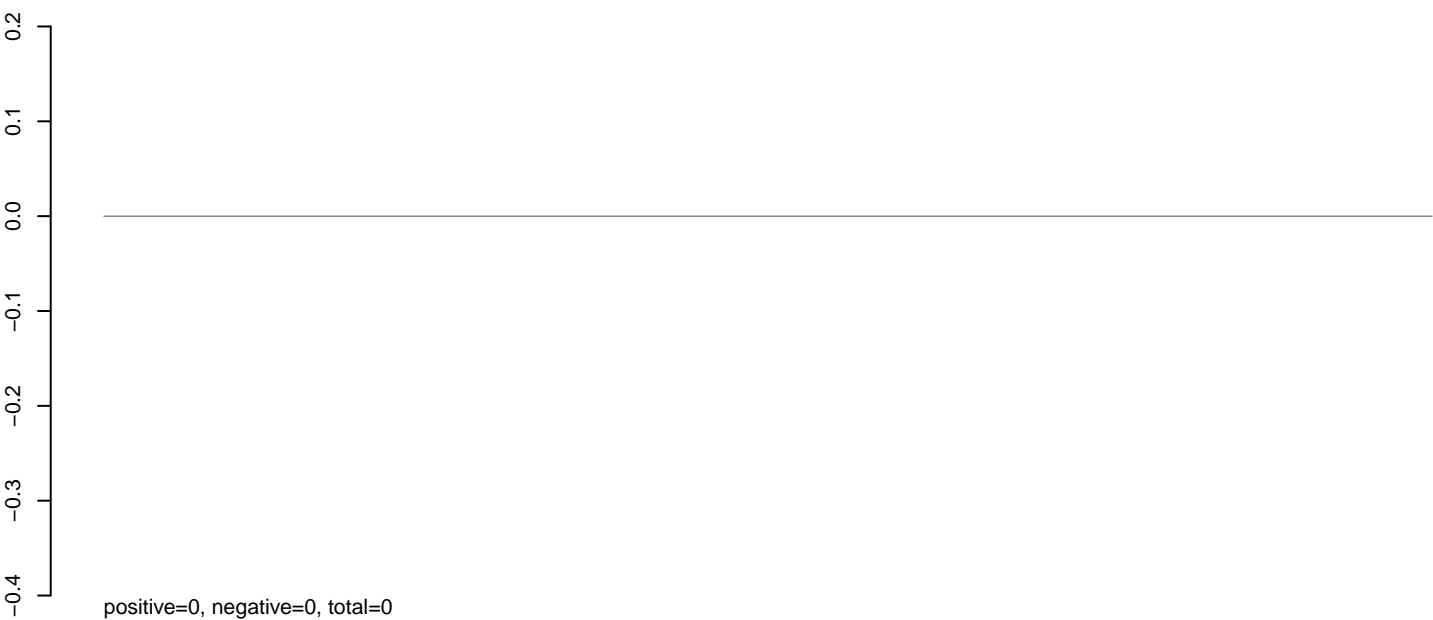


inches; size 2; meg; use 0; total 62 -6_ORF1_ORF1_ORF2_ORF2_glycoprotein_G_and_RNA-dependent_RNA_polymerase_L_genes:1-12774

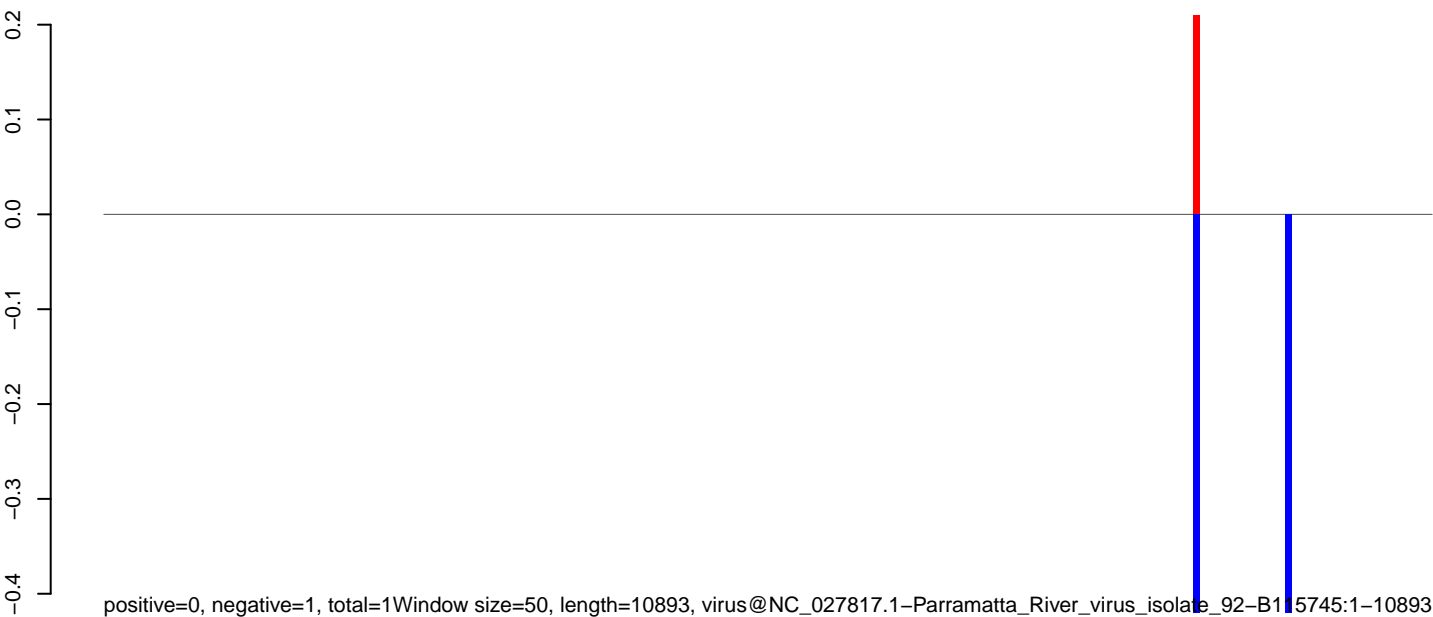
AeAeg_Aag2_DENV4.18_23.rep



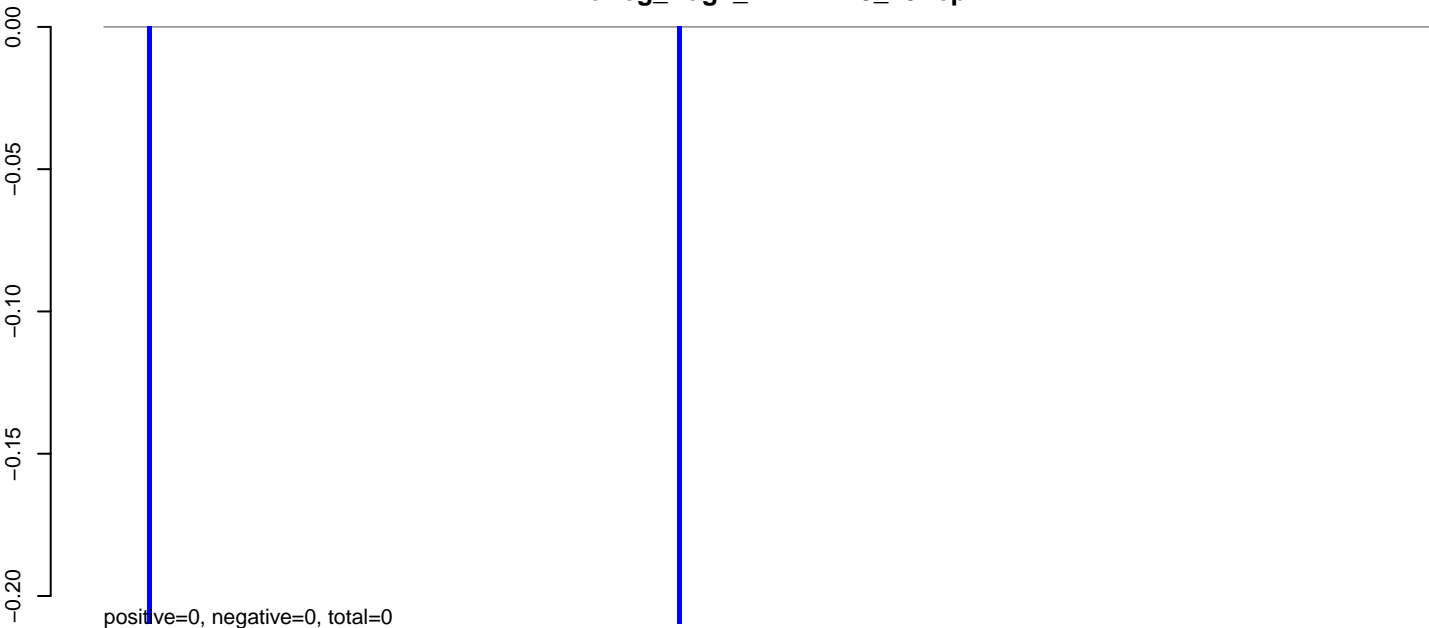
AeAeg_Aag2_DENV4.24_35.rep



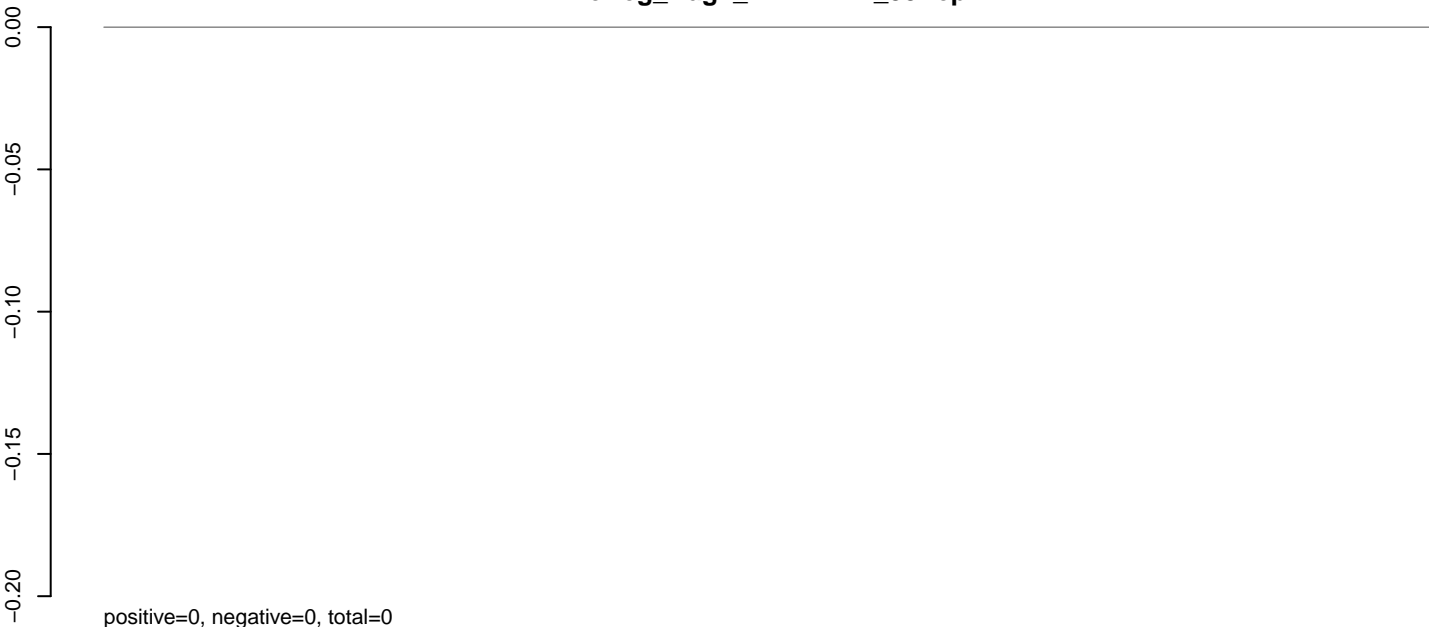
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



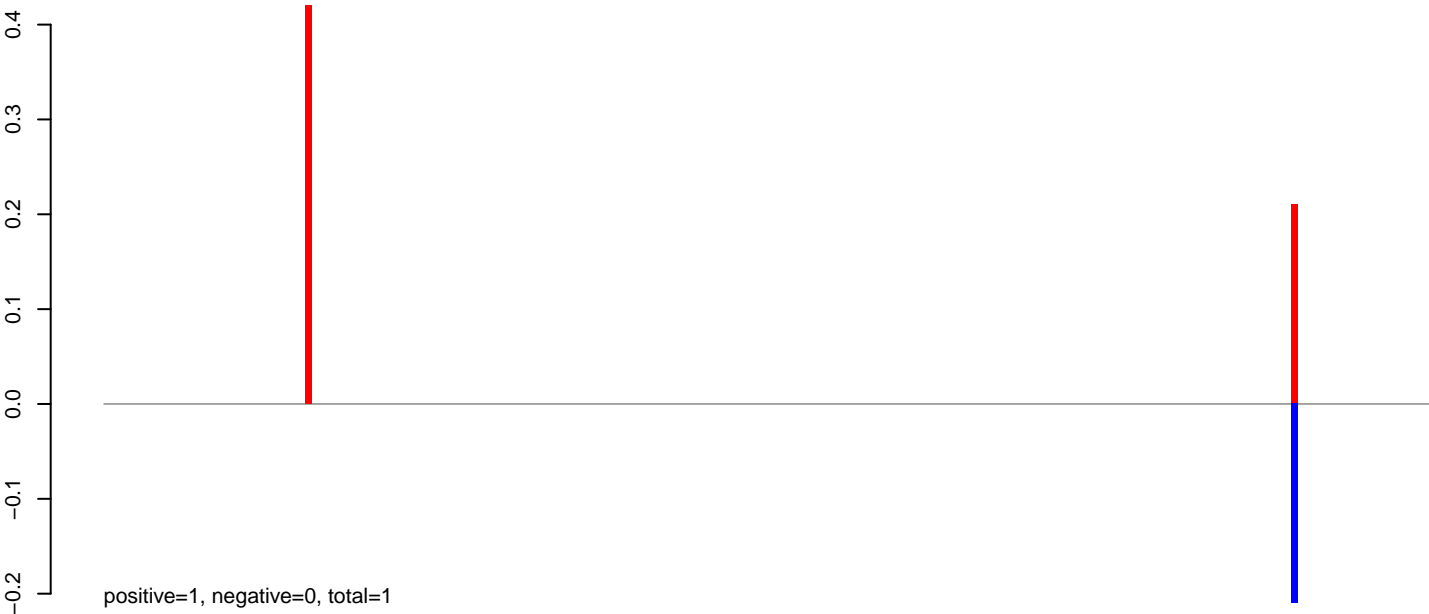
AeAeg_Aag2_DENV4.24_35.rep



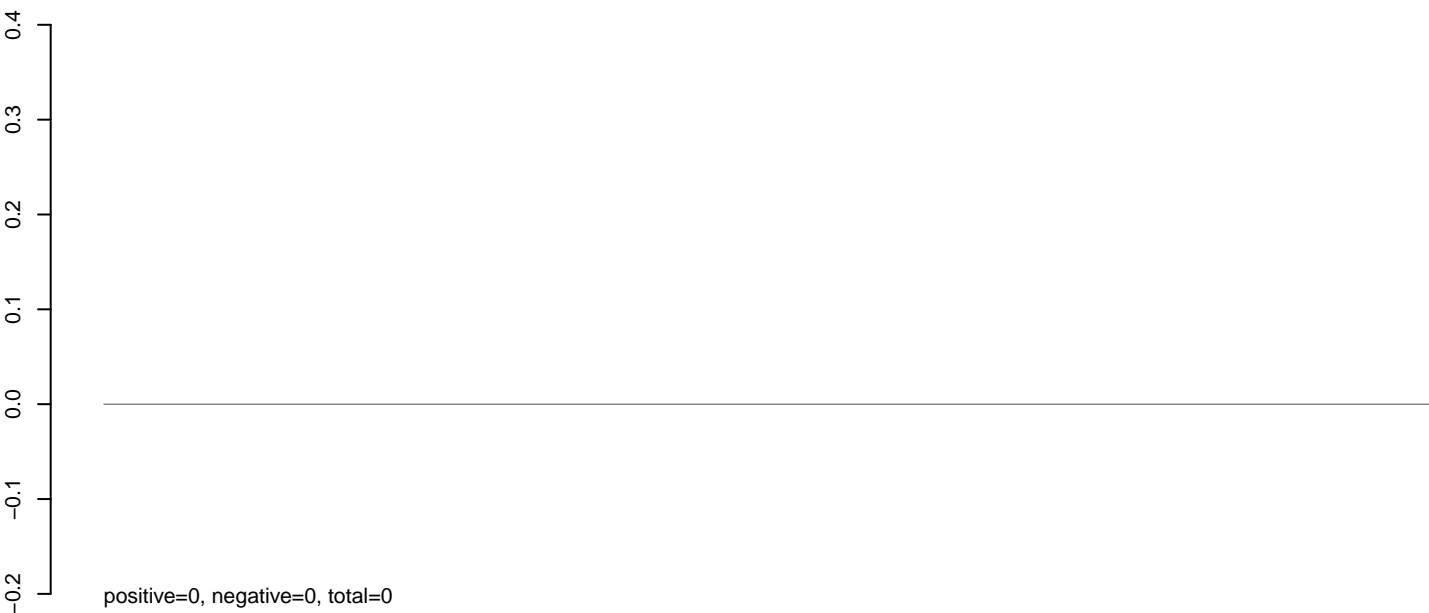
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



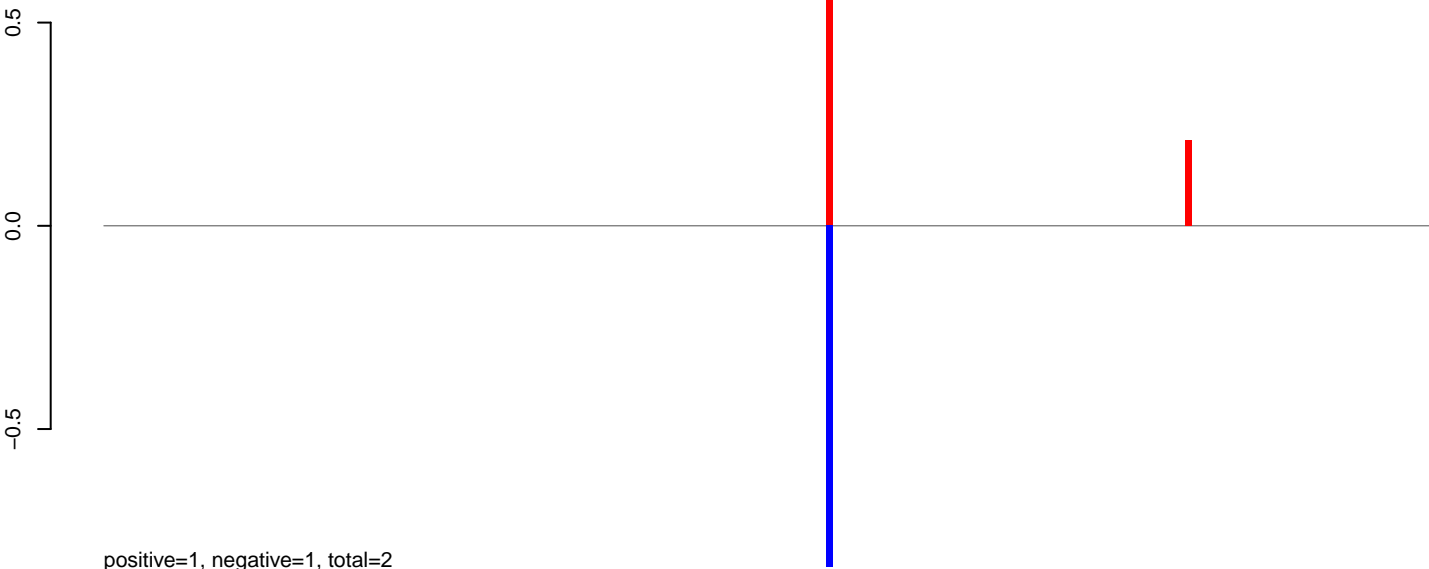
AeAeg_Aag2_DENV4.rep



Window size=50, length=10878, virus@NC_024299.2-Nienokoue_virus_isolate_B51_CI_2004:1-10878

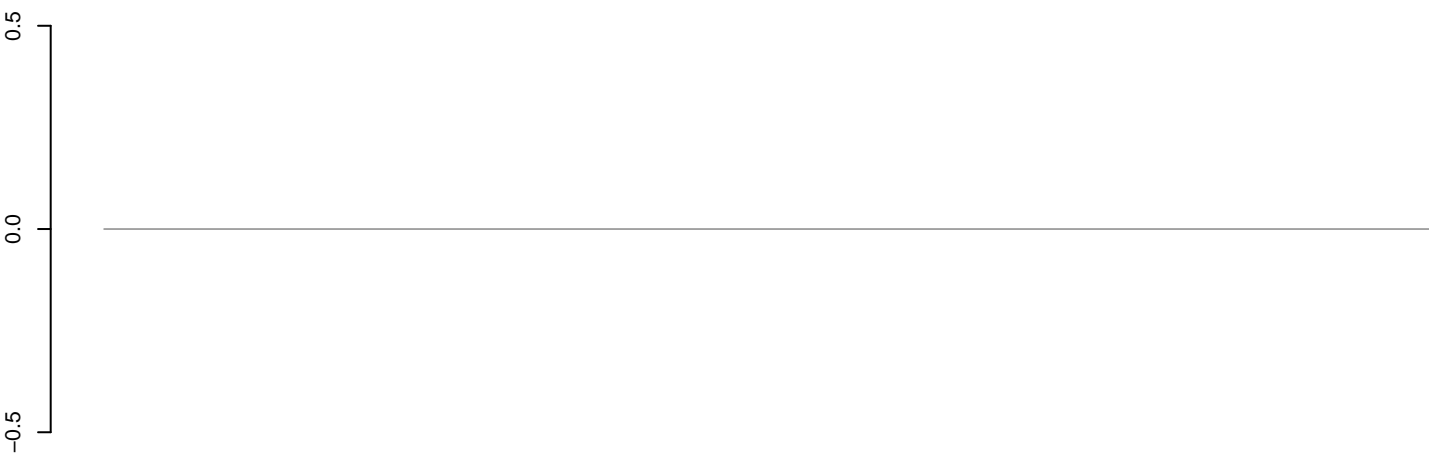
0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV4.18_23.rep



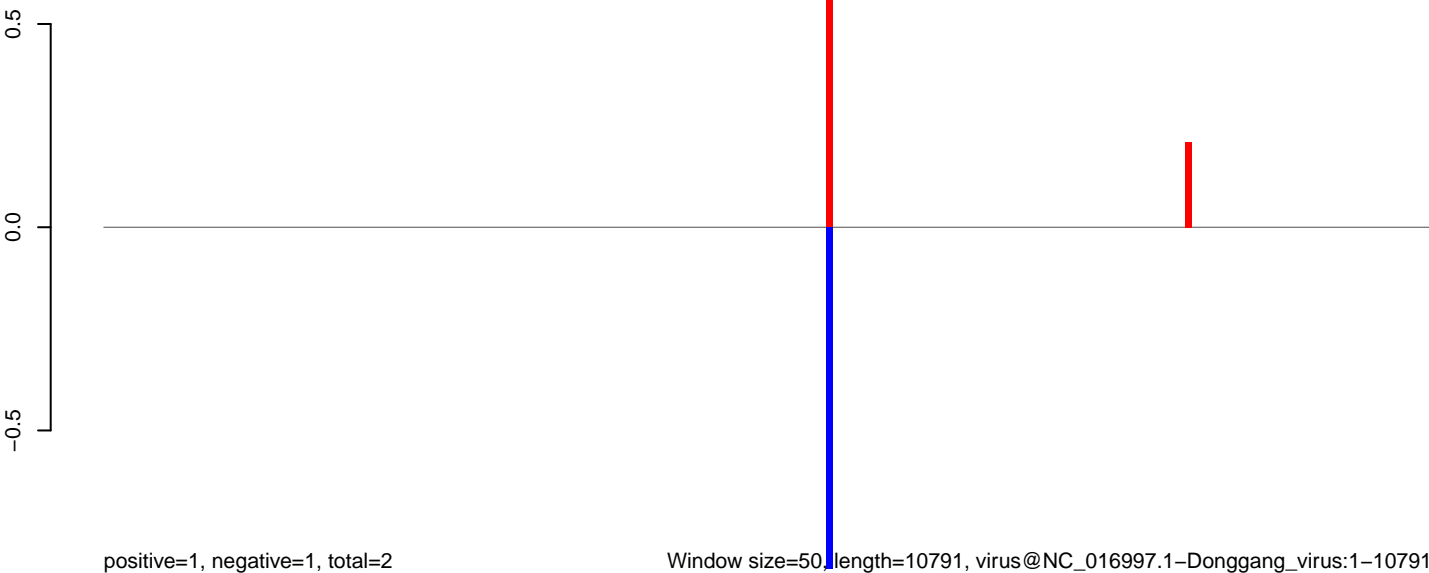
positive=1, negative=1, total=2

AeAeg_Aag2_DENV4.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_DENV4.rep

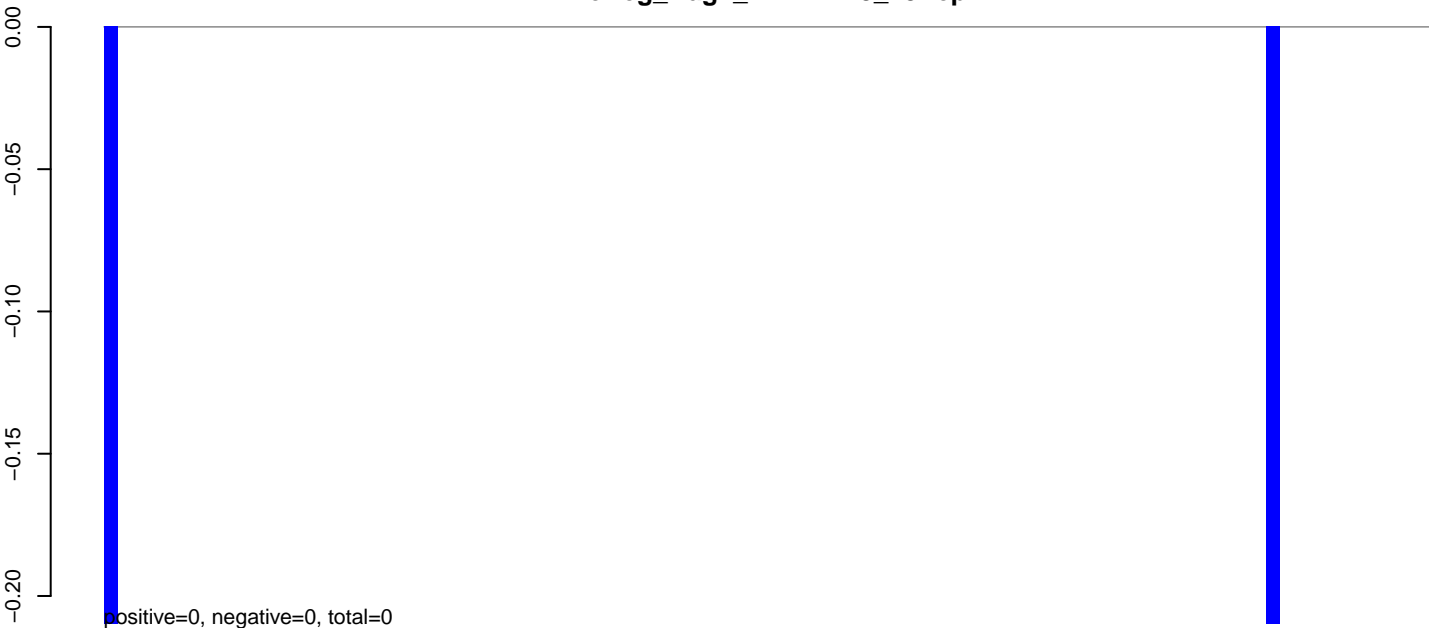


positive=1, negative=1, total=2

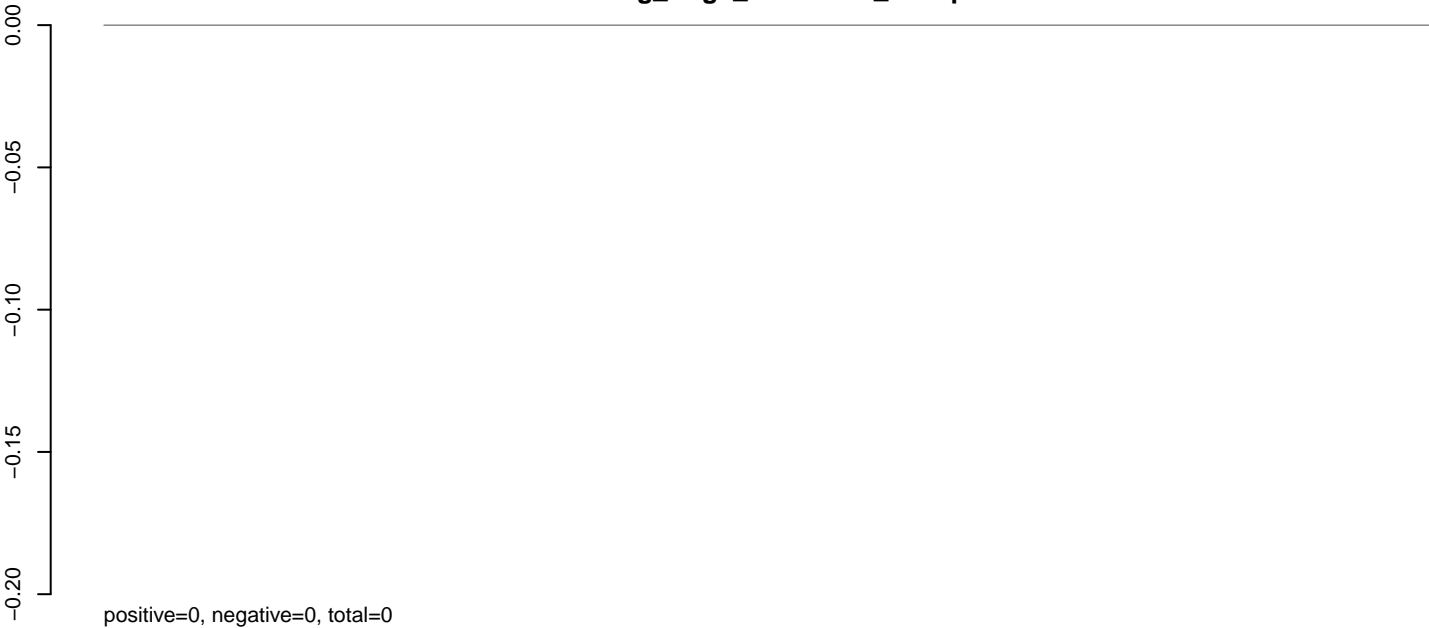
Window size=50, length=10791, virus@NC_016997.1-Donggang_virus:1-10791

0 2000 4000 6000 8000 10000

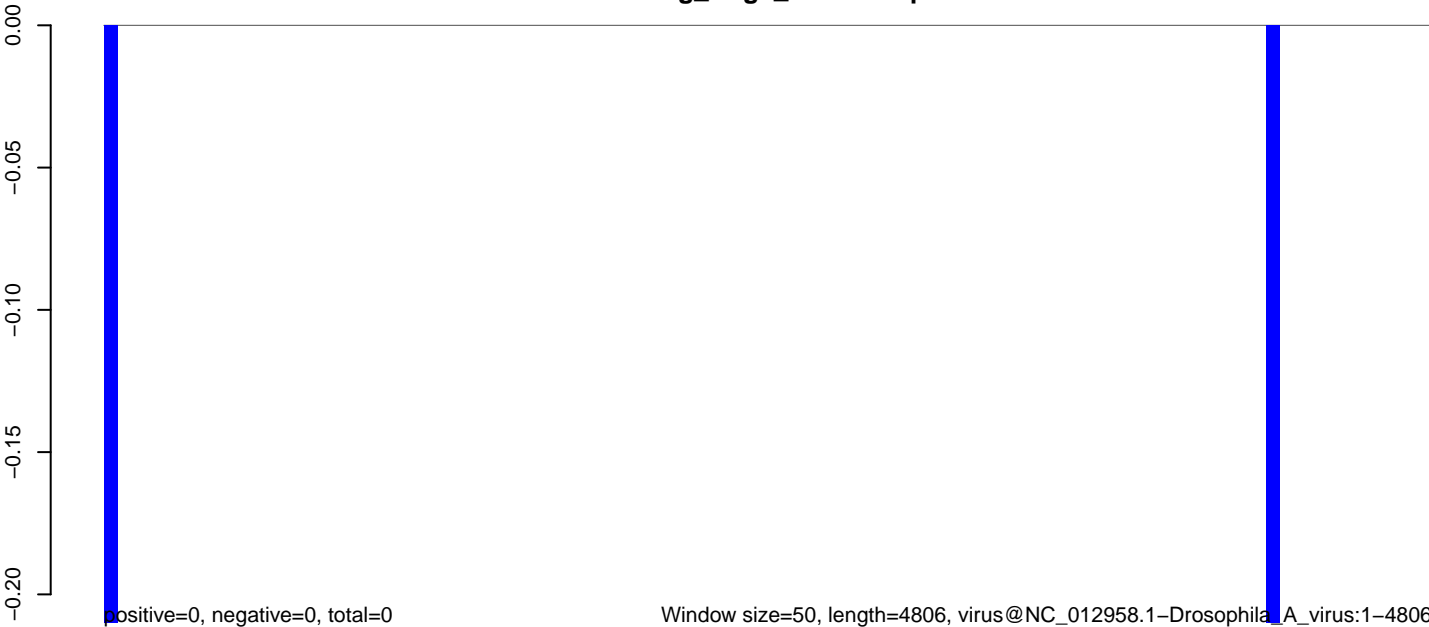
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



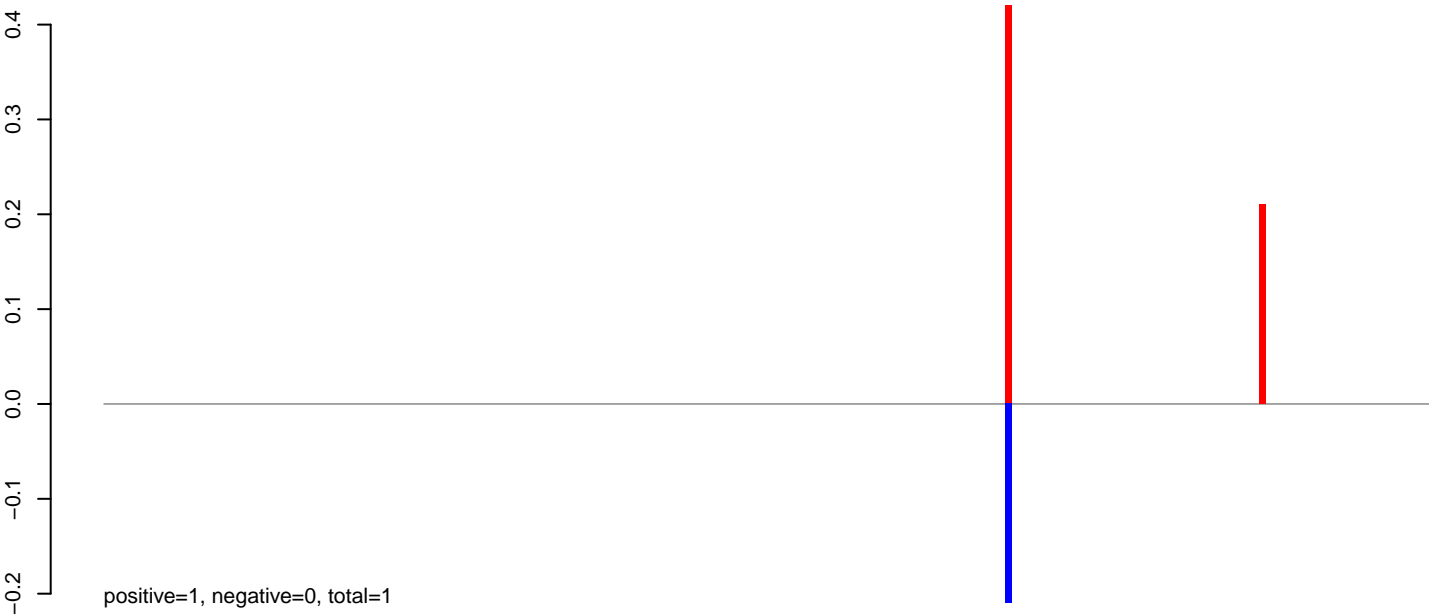
AeAeg_Aag2_DENV4.rep



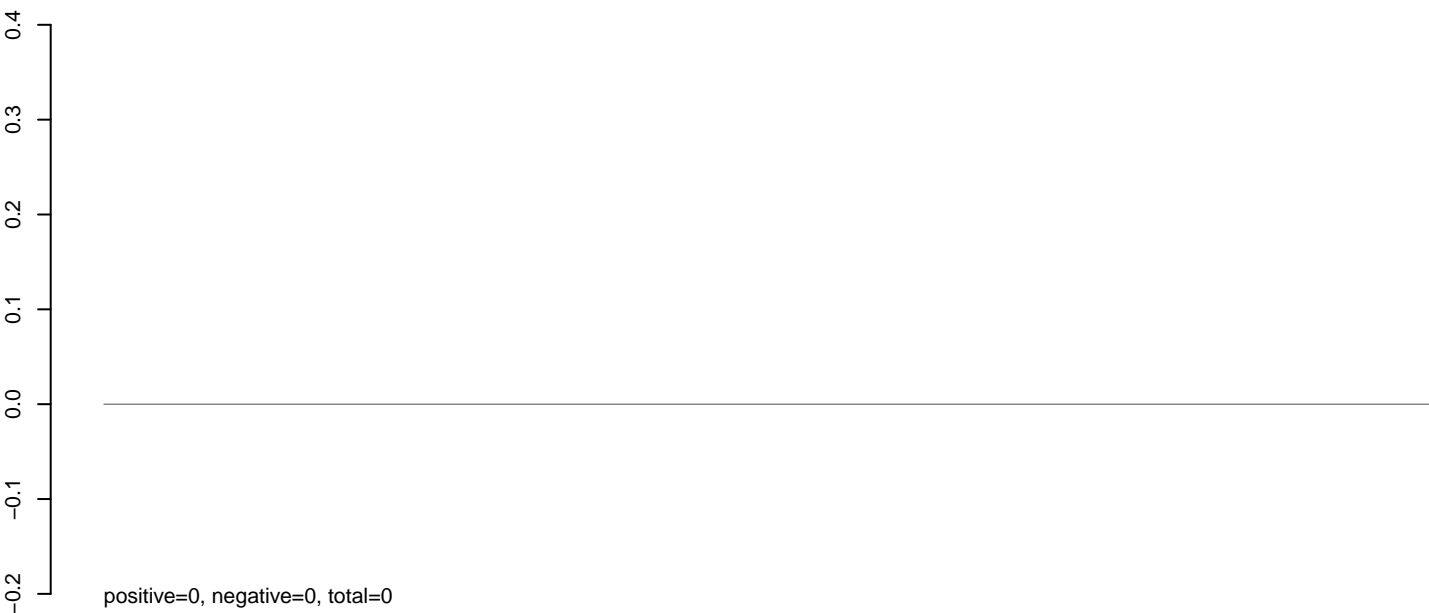
Window size=50, length=4806, virus@NC_012958.1-Drosophila_A_virus:1-4806

0 1000 2000 3000 4000 5000

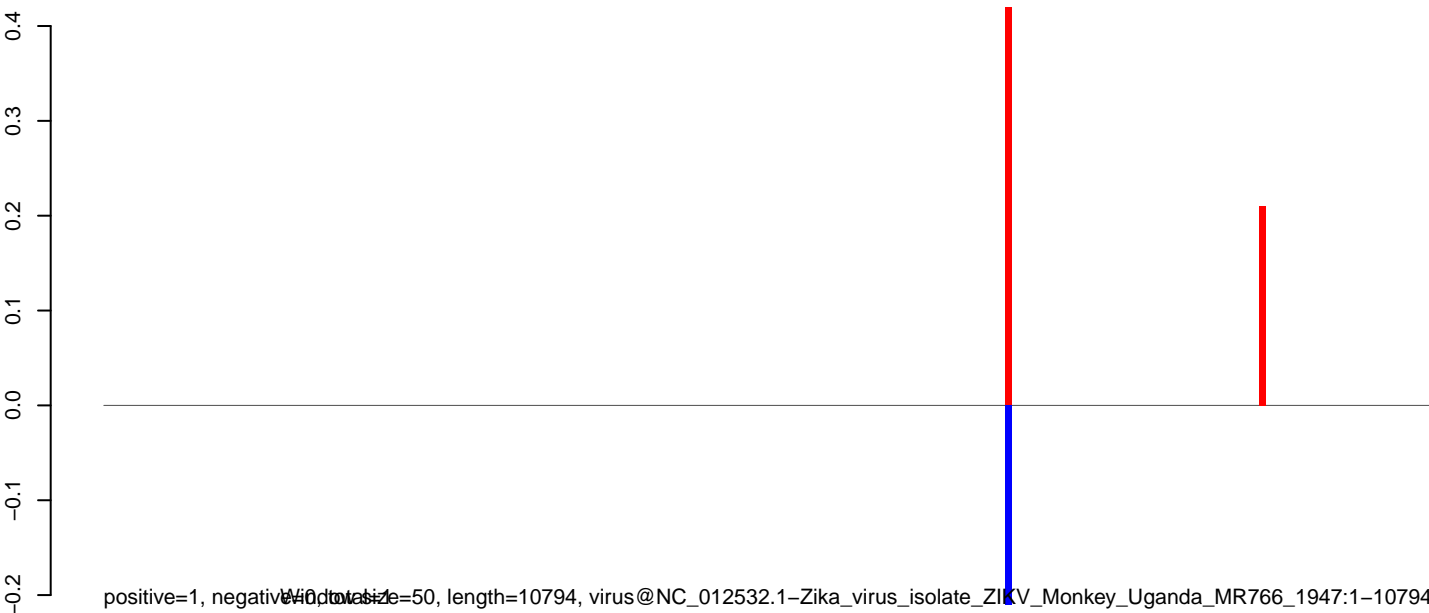
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



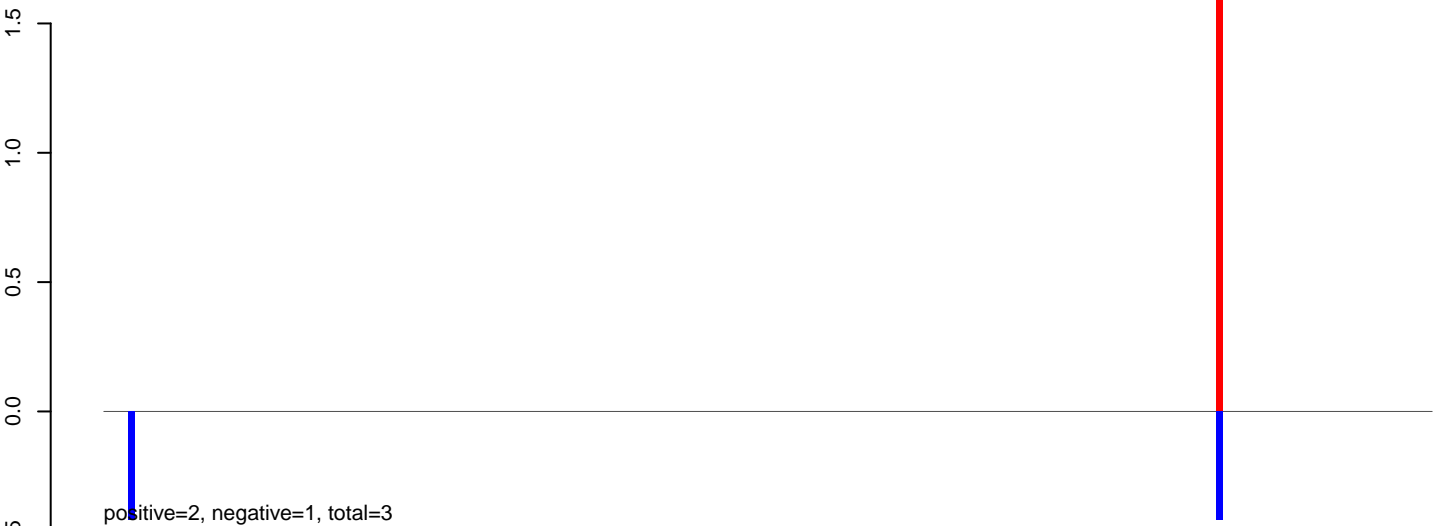
AeAeg_Aag2_DENV4.rep



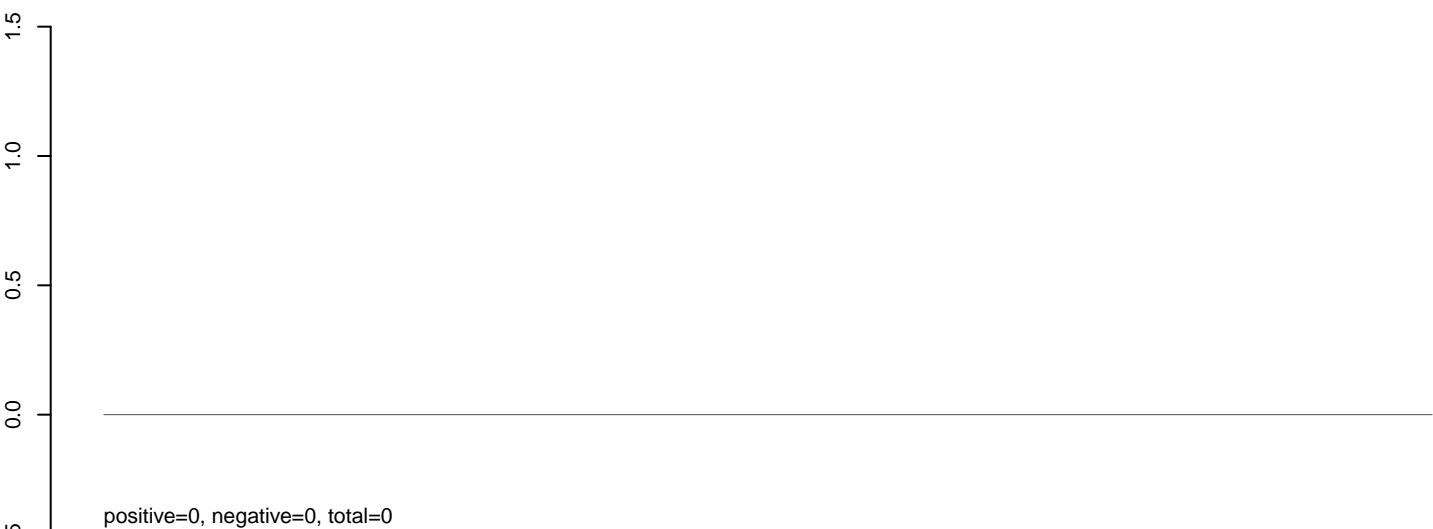
Window size=50, length=10794, virus@NC_012532.1-Zika_virus_isolate_ZIKV_Monkey_Uganda_MR766_1947:1-10794

0 2000 4000 6000 8000 10000

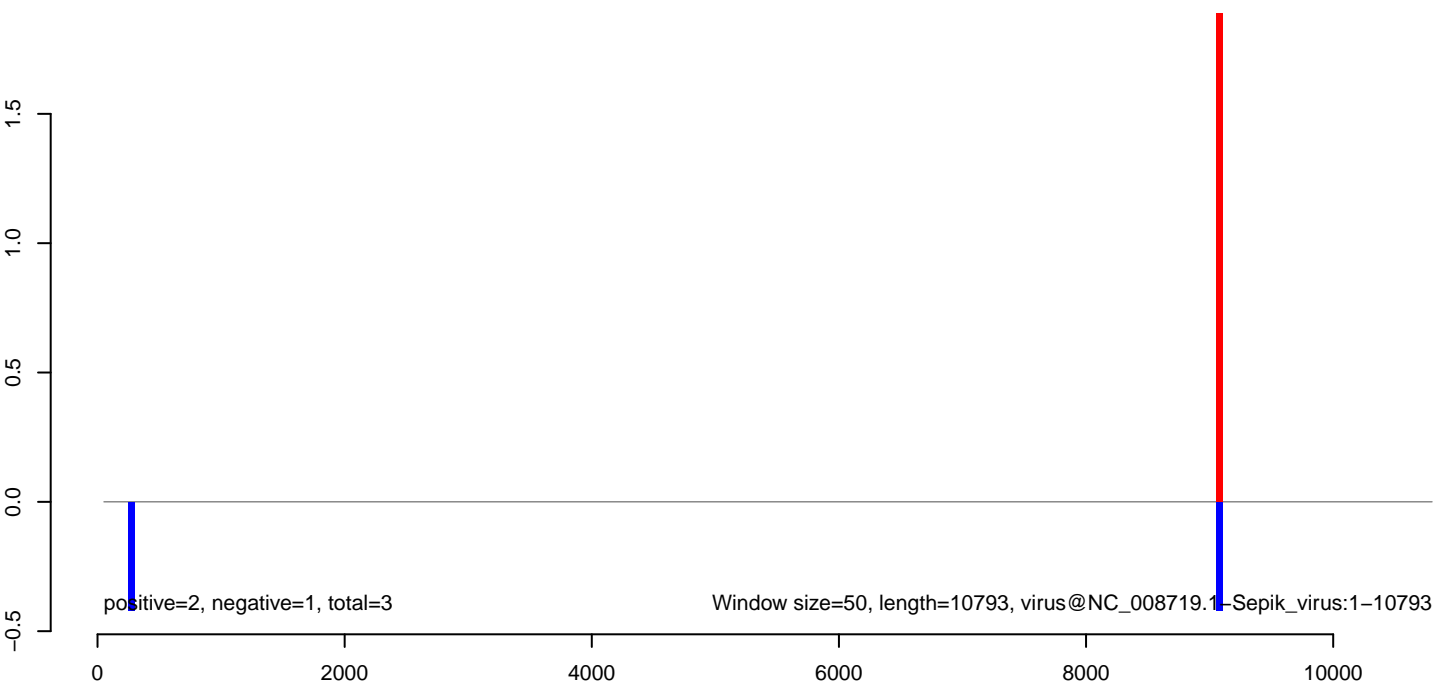
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



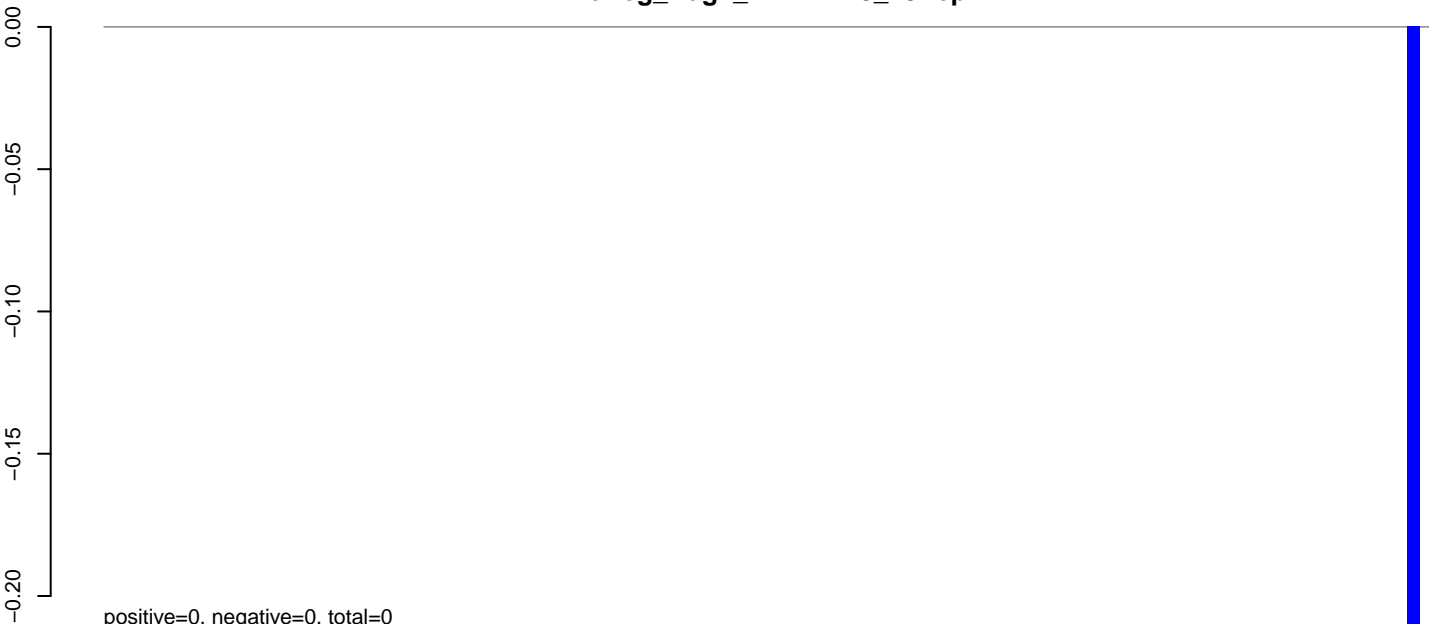
AeAeg_Aag2_DENV4.24_35.rep



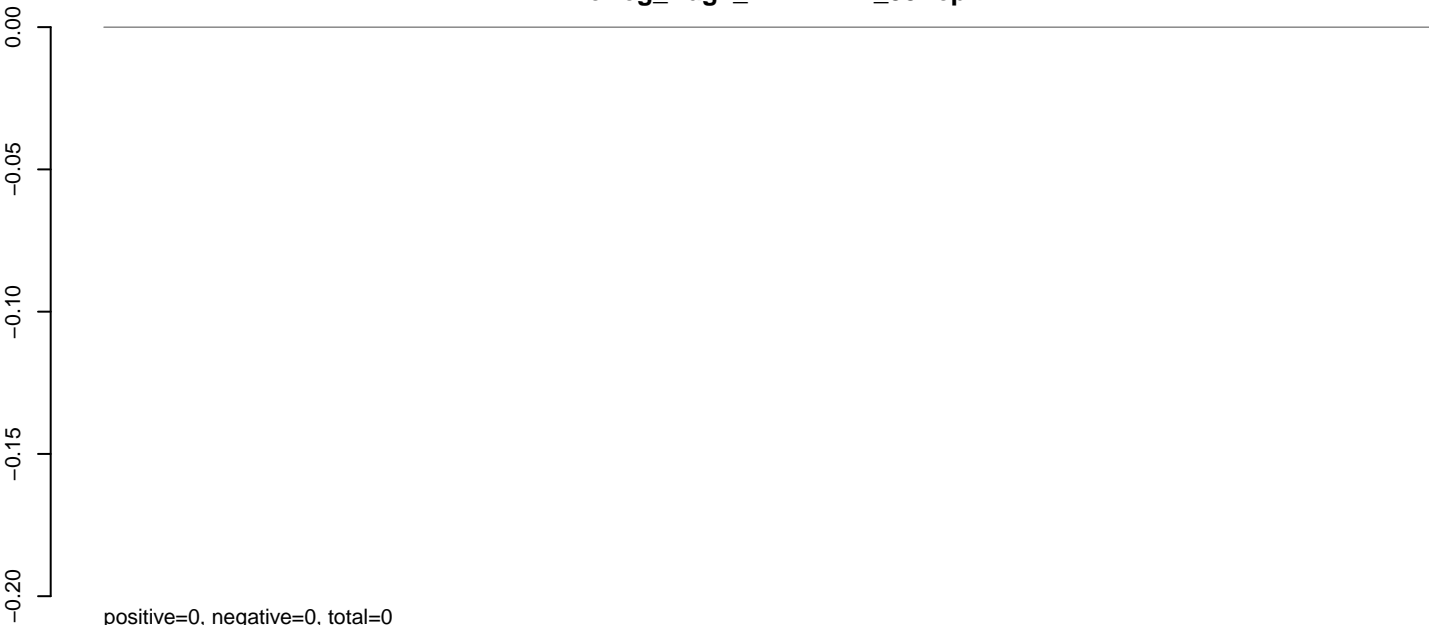
AeAeg_Aag2_DENV4.rep



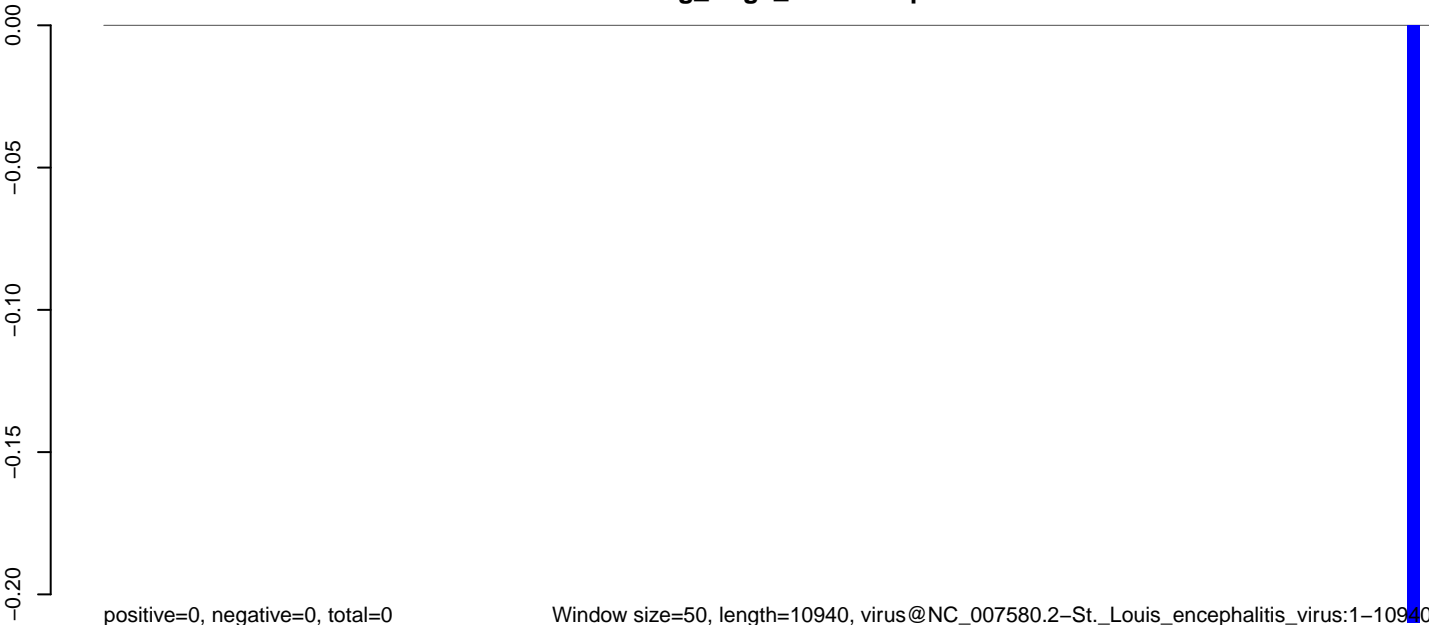
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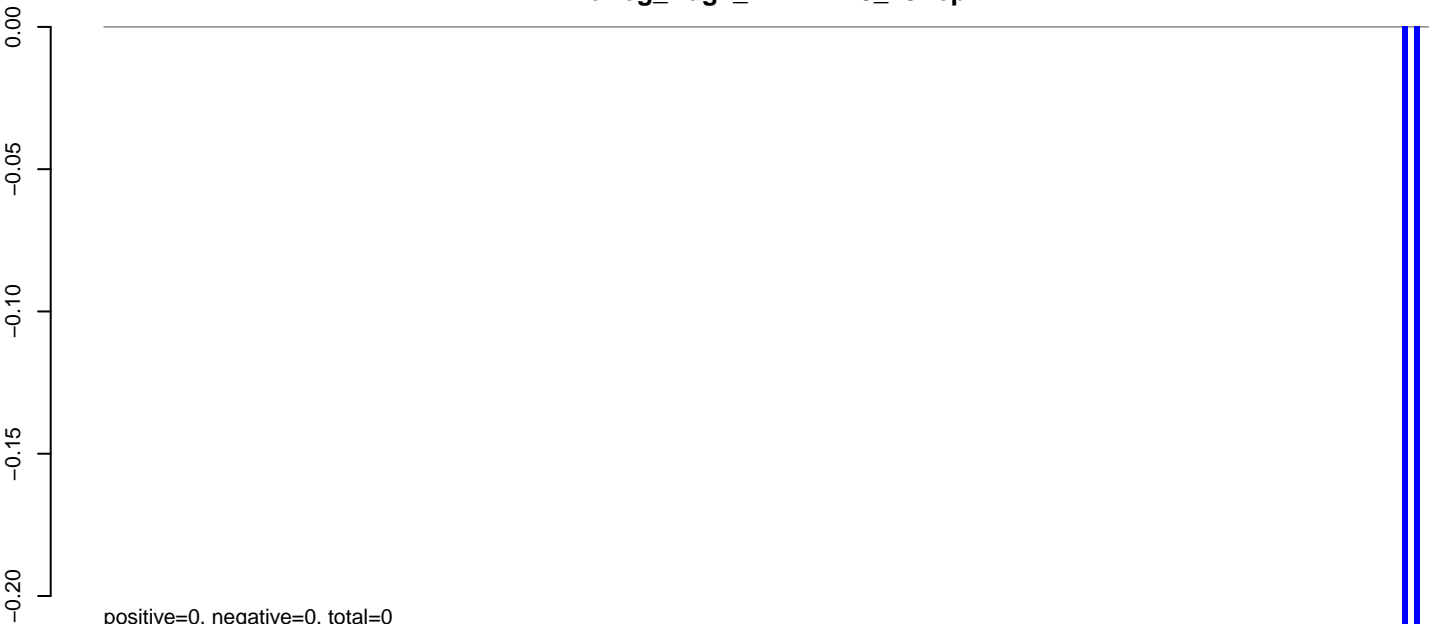
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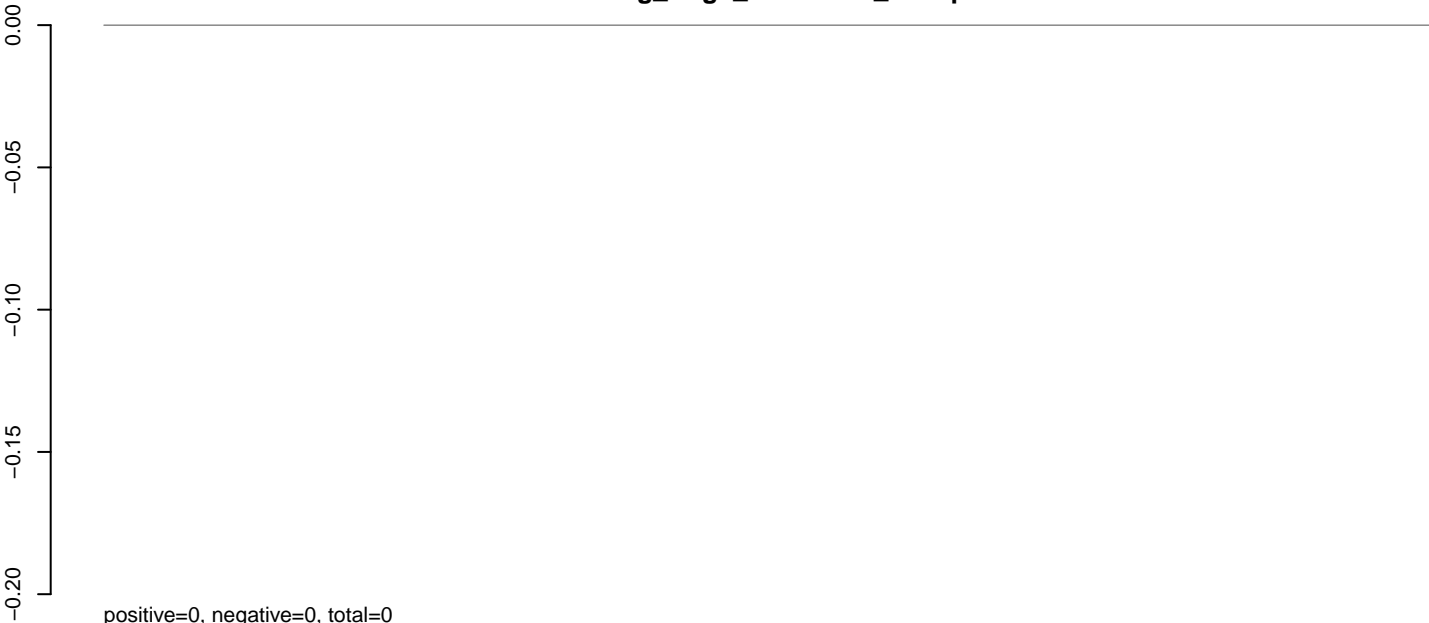
AeAeg_Aag2_DENV4.rep



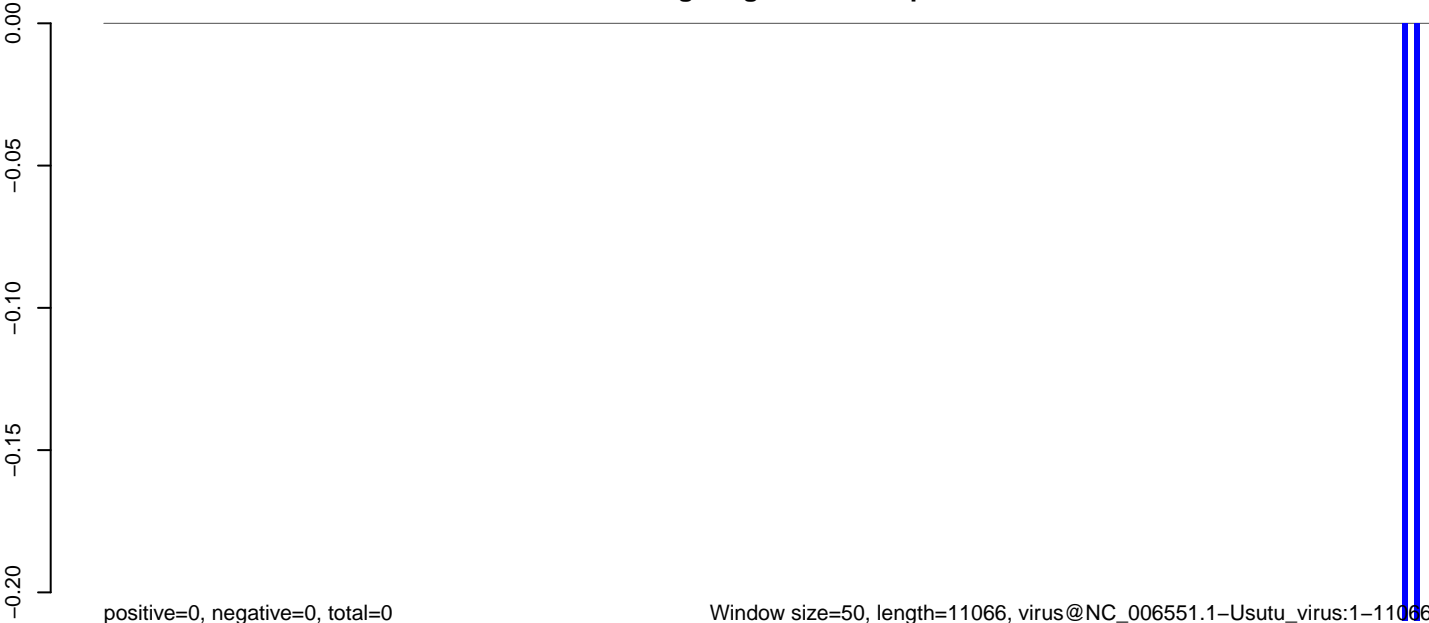
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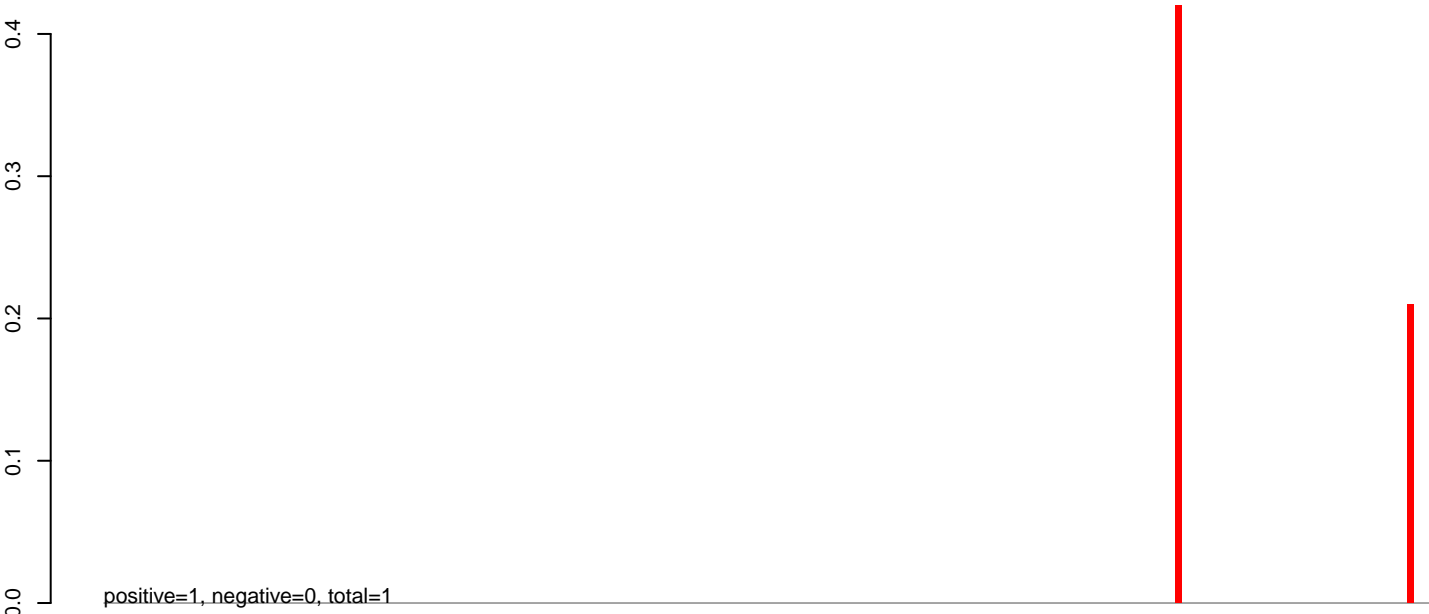
AeAeg_Aag2_DENV4.24_35.rep



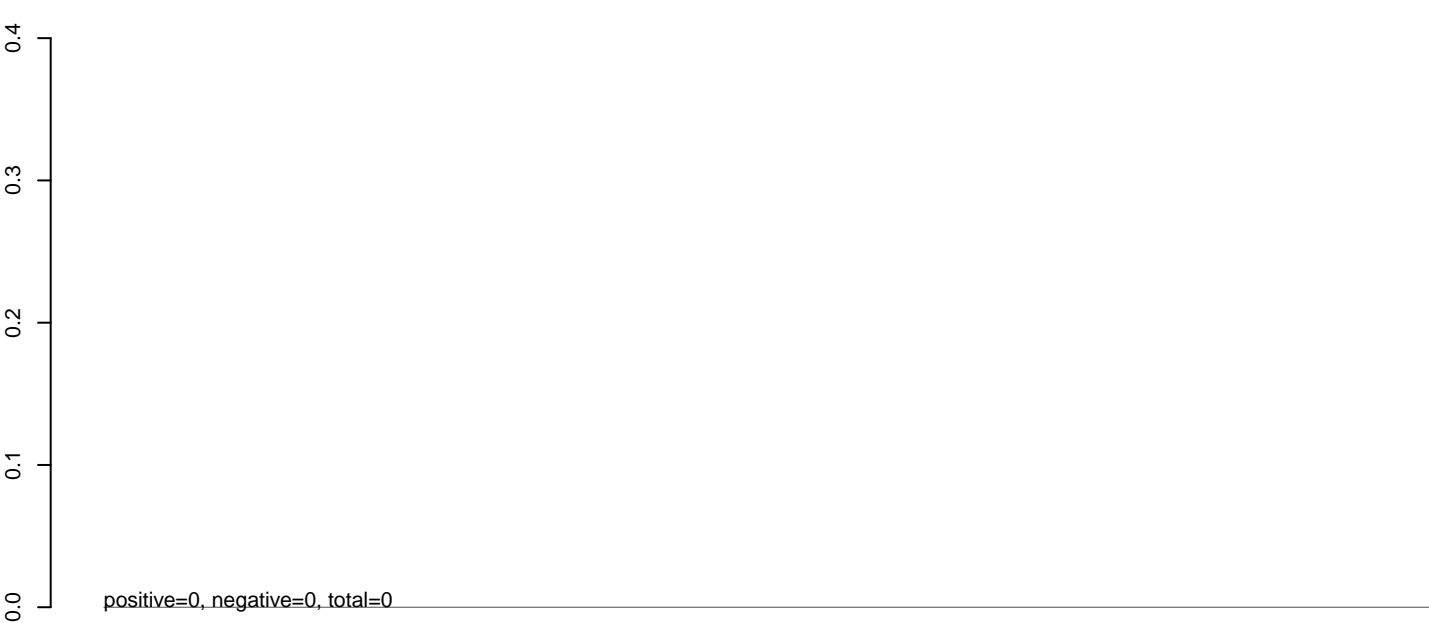
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

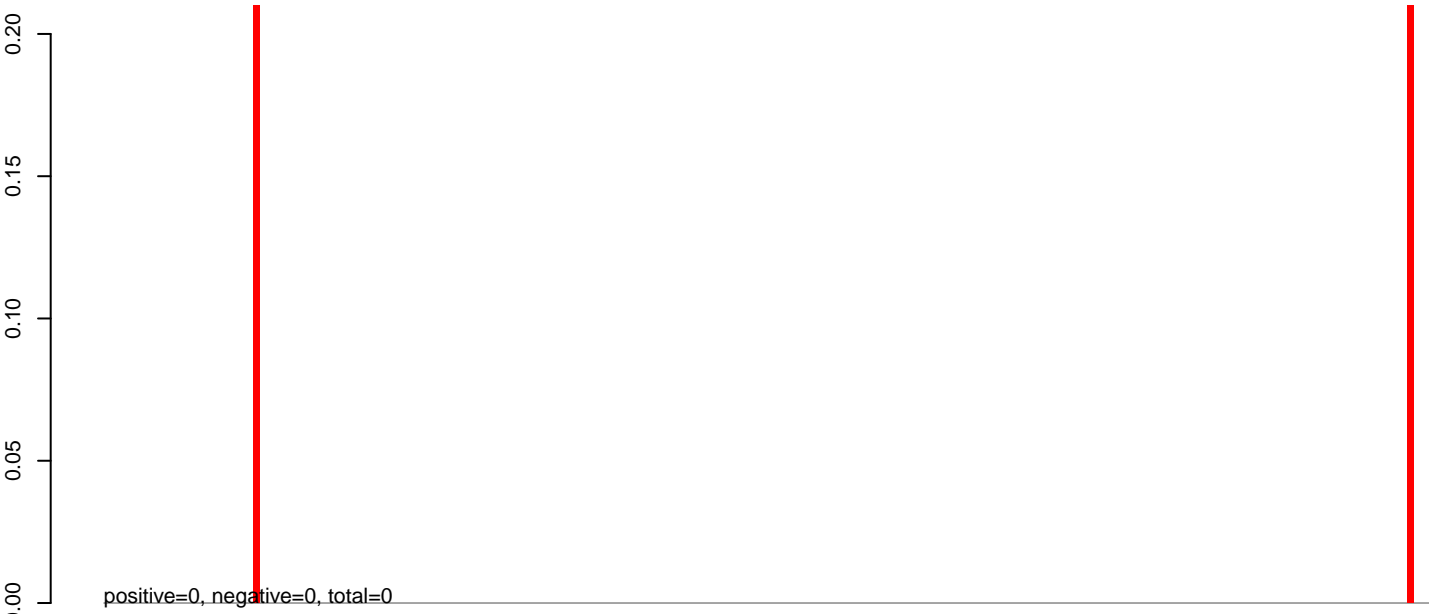


AeAeg_Aag2_DENV4.rep

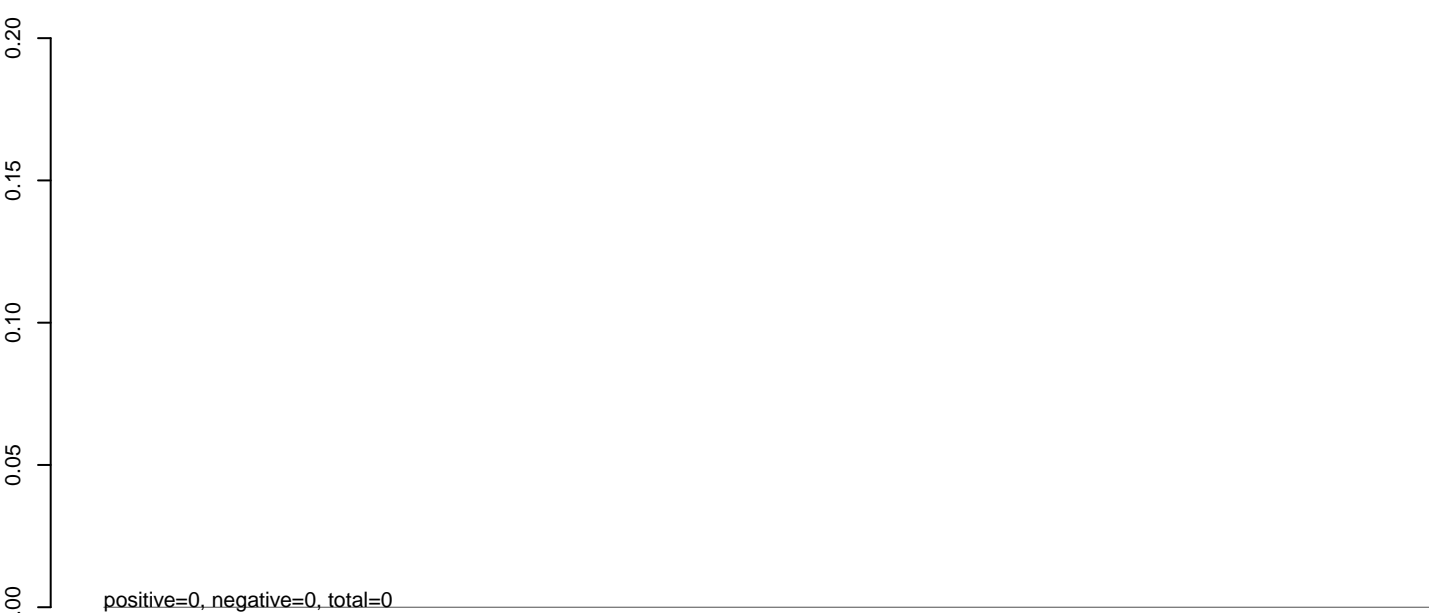


0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

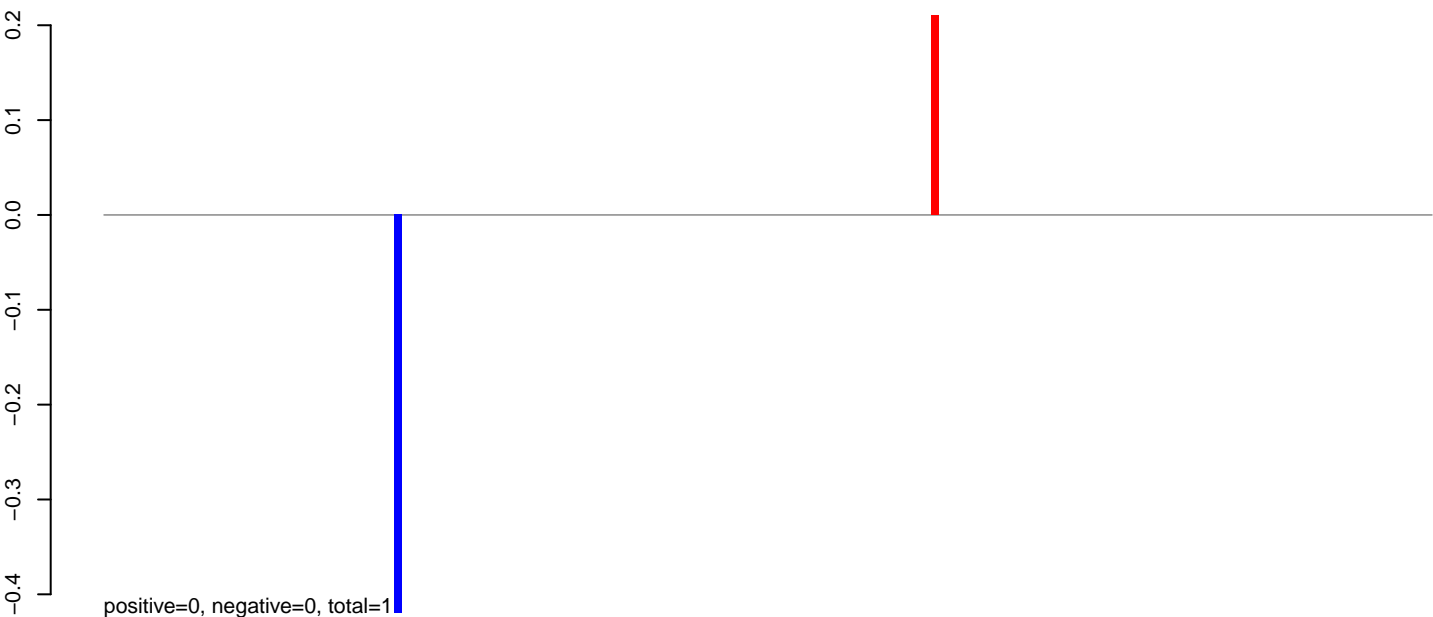


AeAeg_Aag2_DENV4.rep

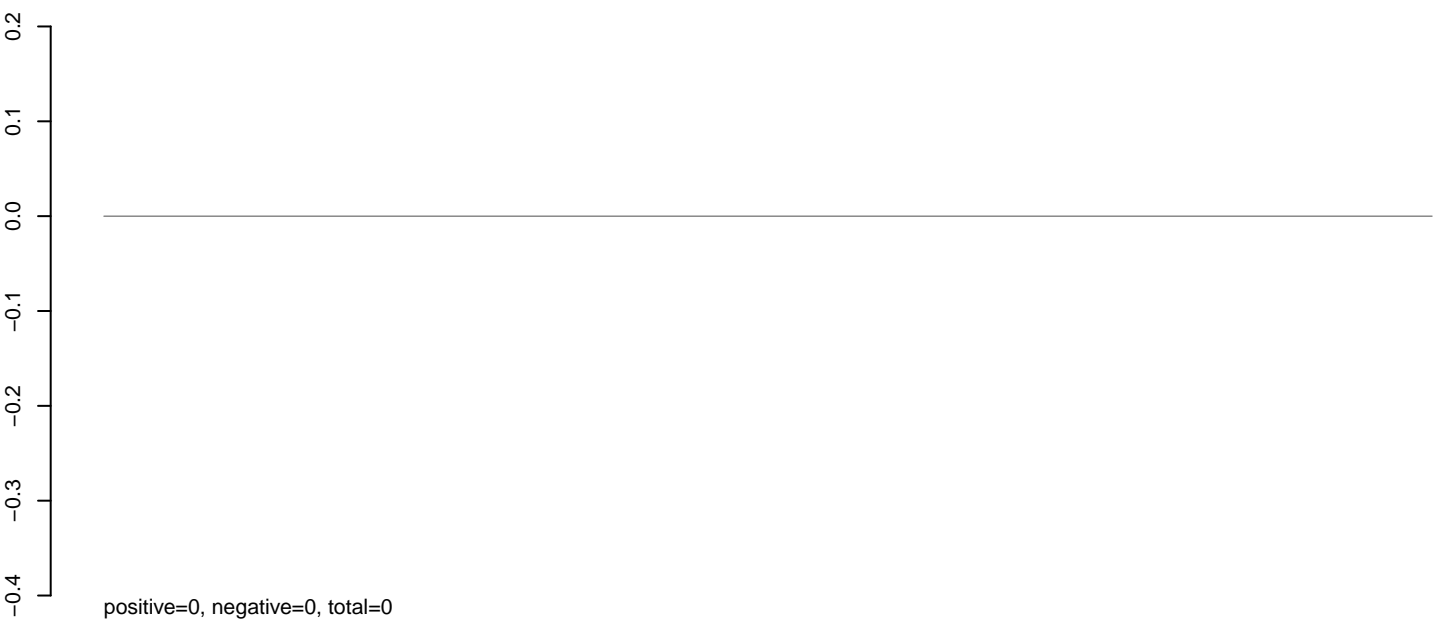


0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV4.18_23.rep



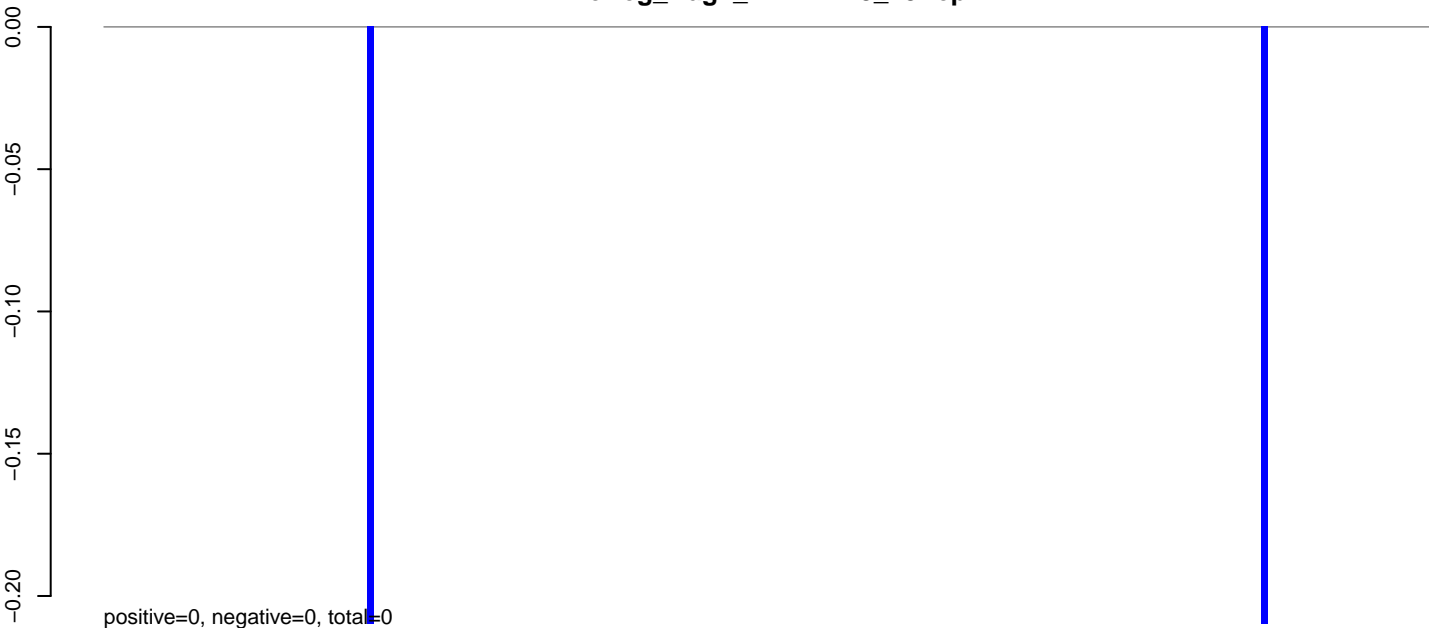
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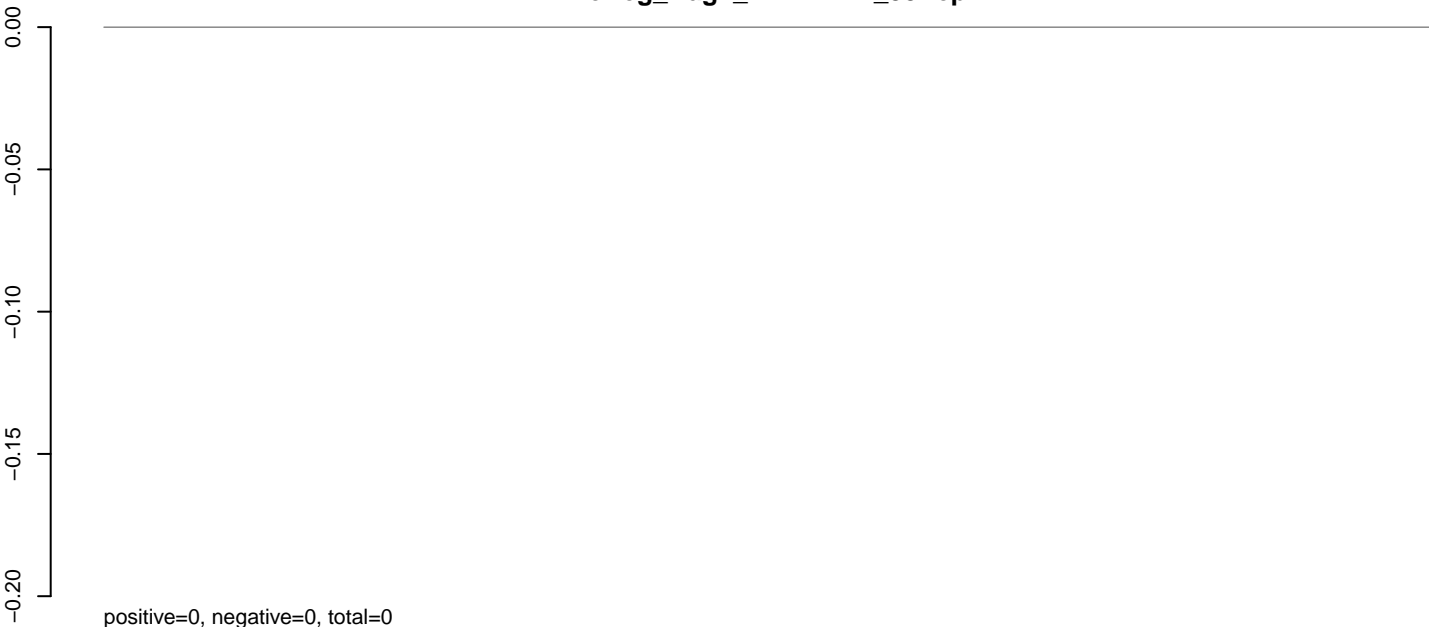
AeAeg_Aag2_DENV4.rep



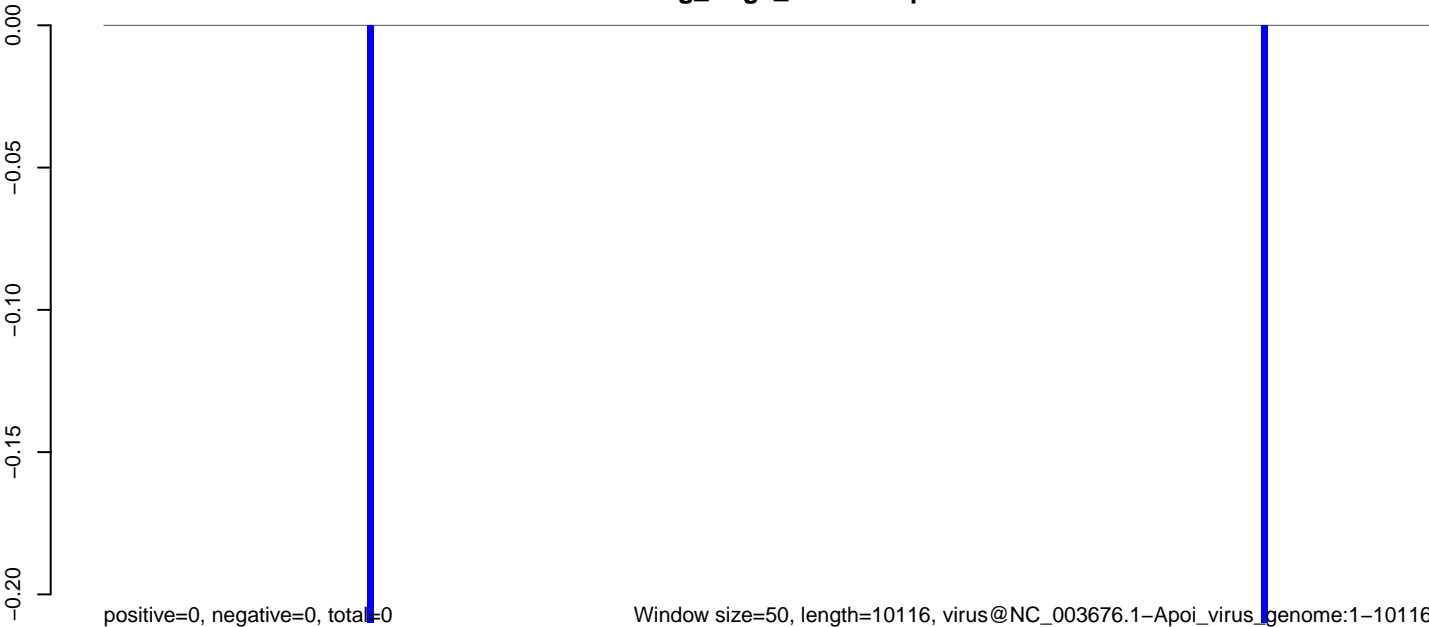
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



0 2000 4000 6000 8000 10000

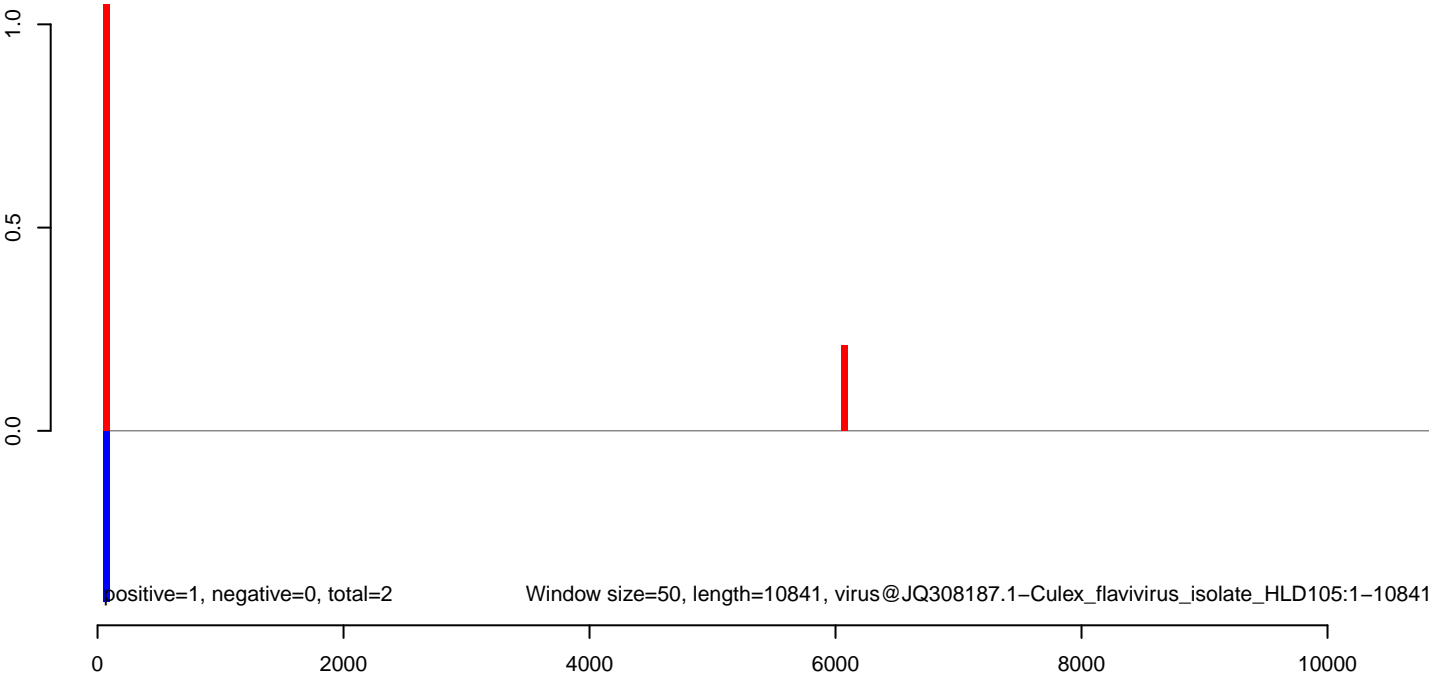
AeAeg_Aag2_DENV4.18_23.rep



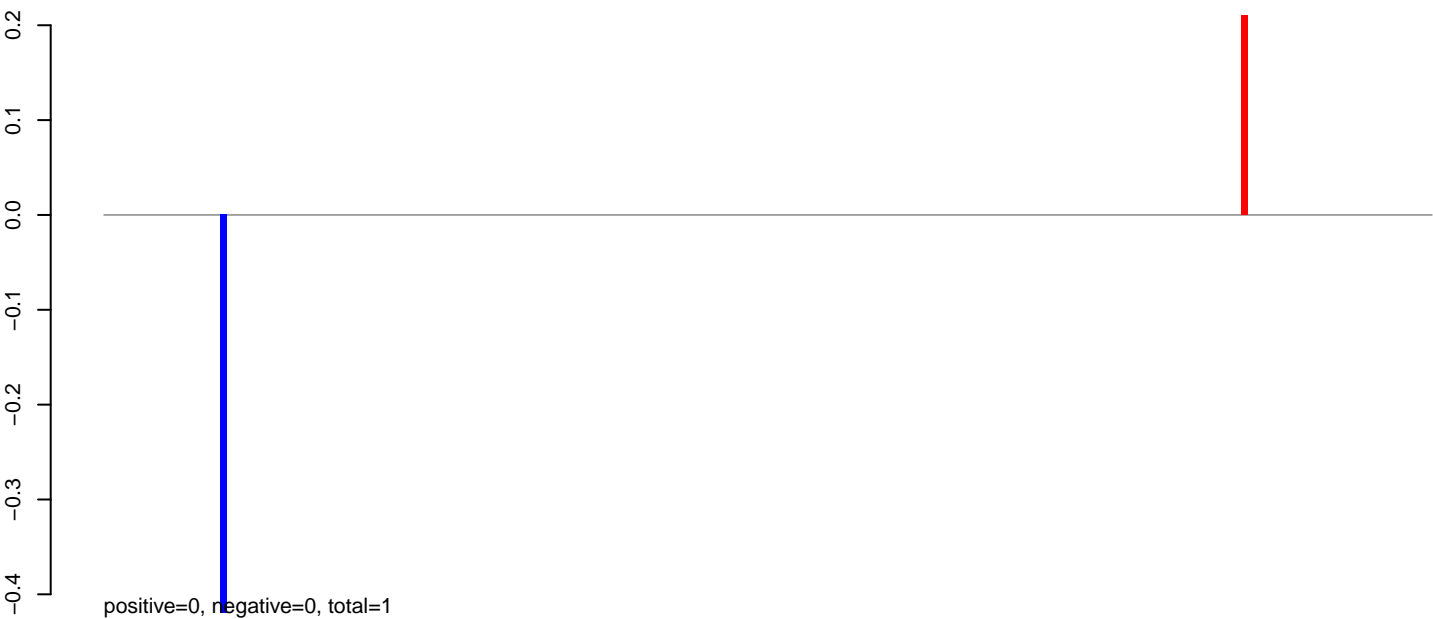
AeAeg_Aag2_DENV4.24_35.rep



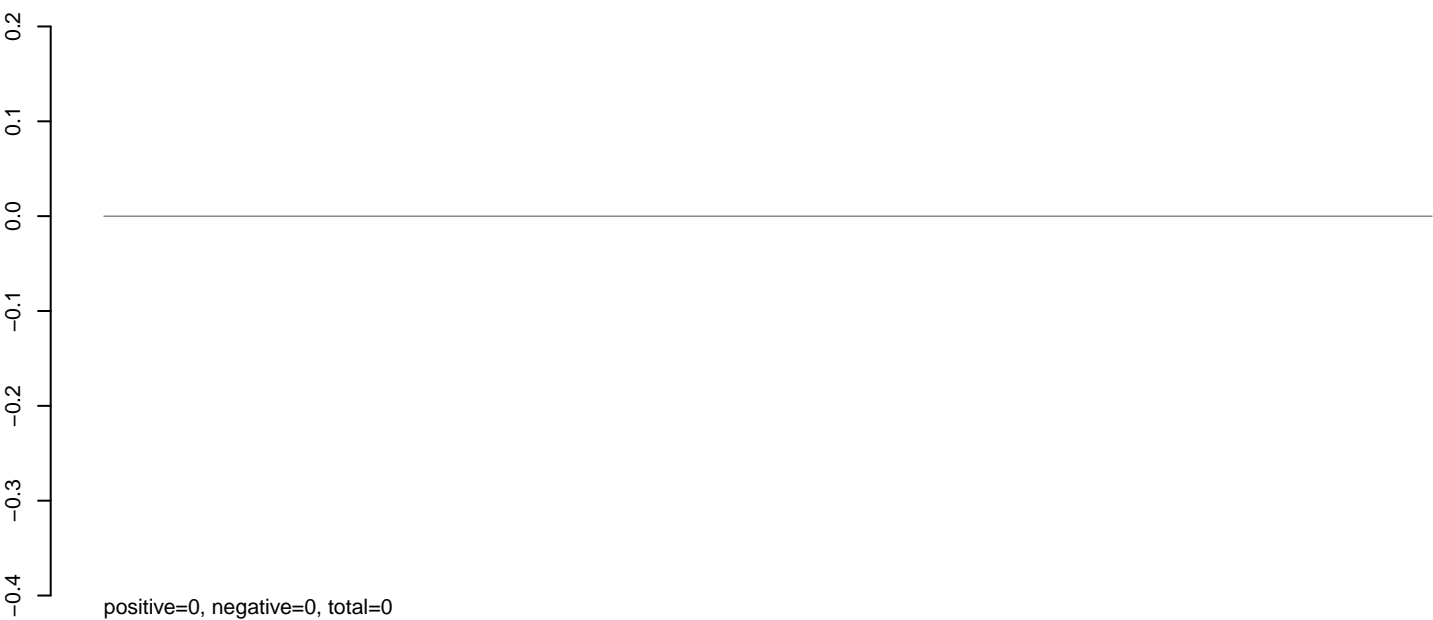
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



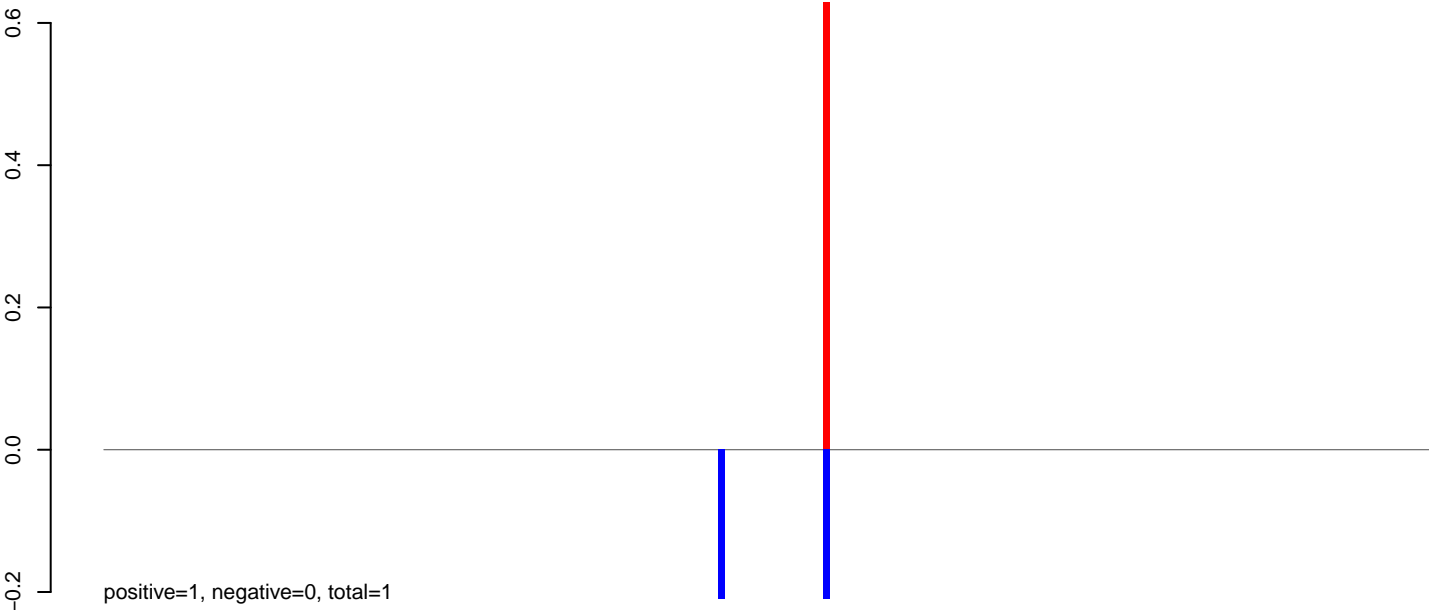
AeAeg_Aag2_DENV4.rep



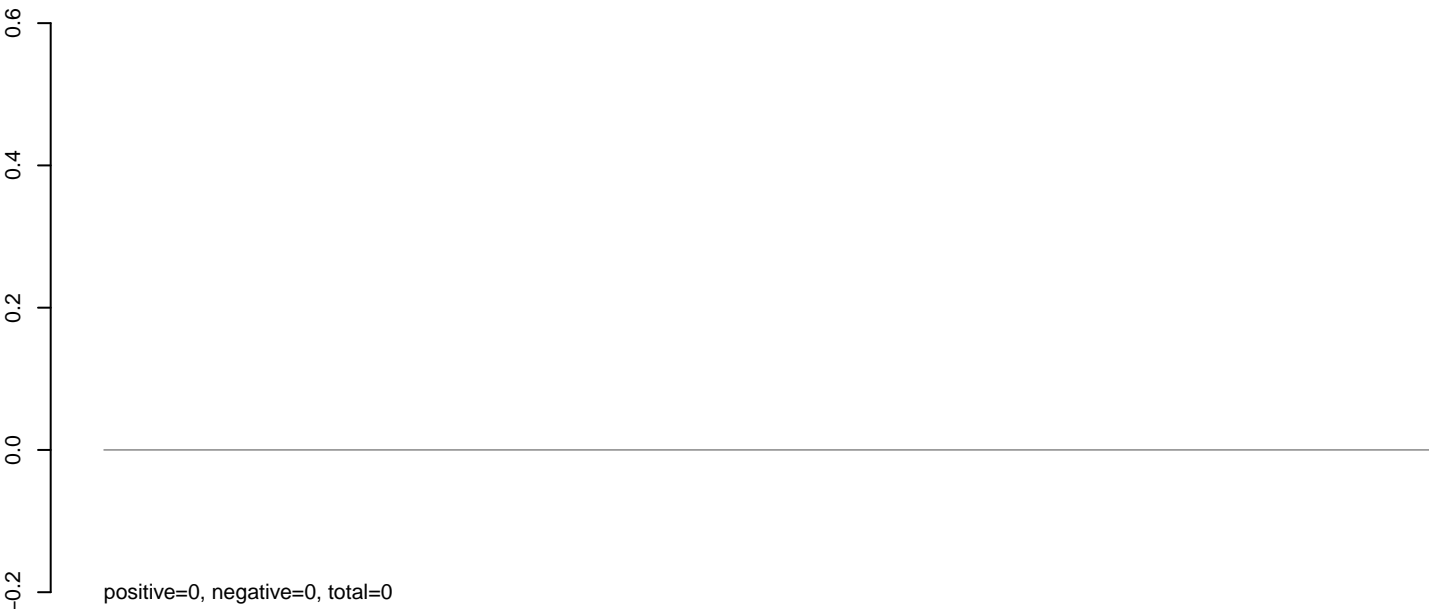
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0 2000 4000 6000 8000 10000

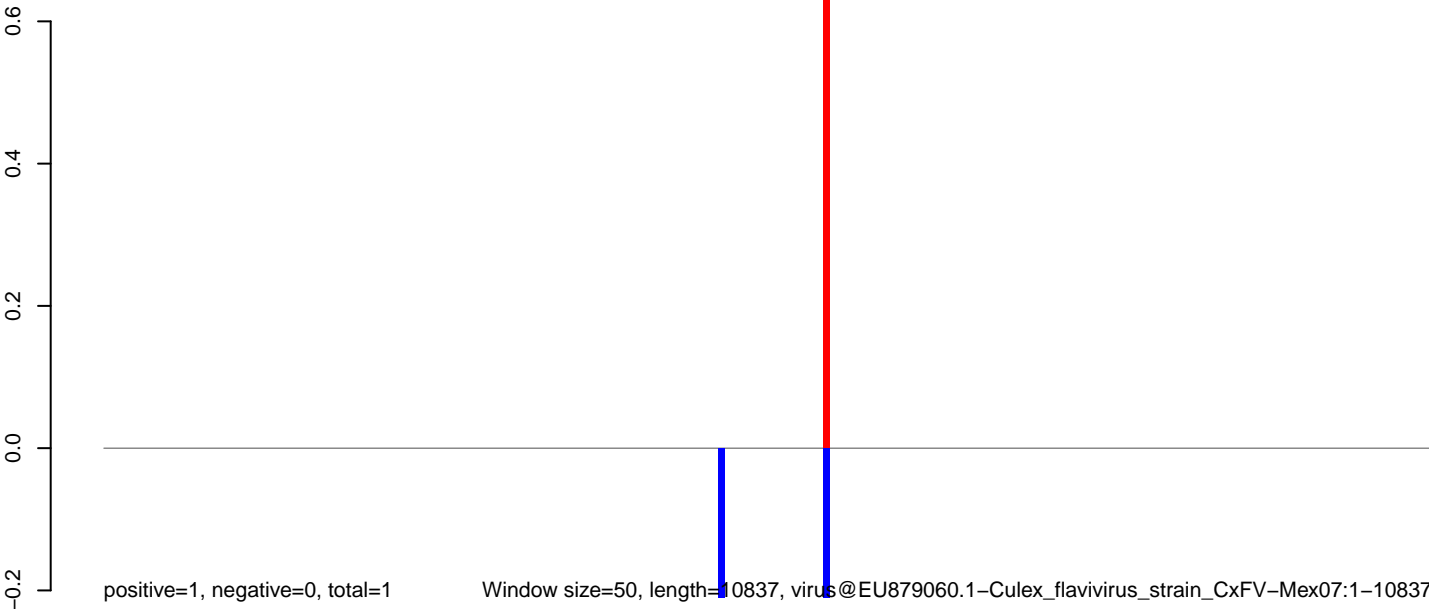
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

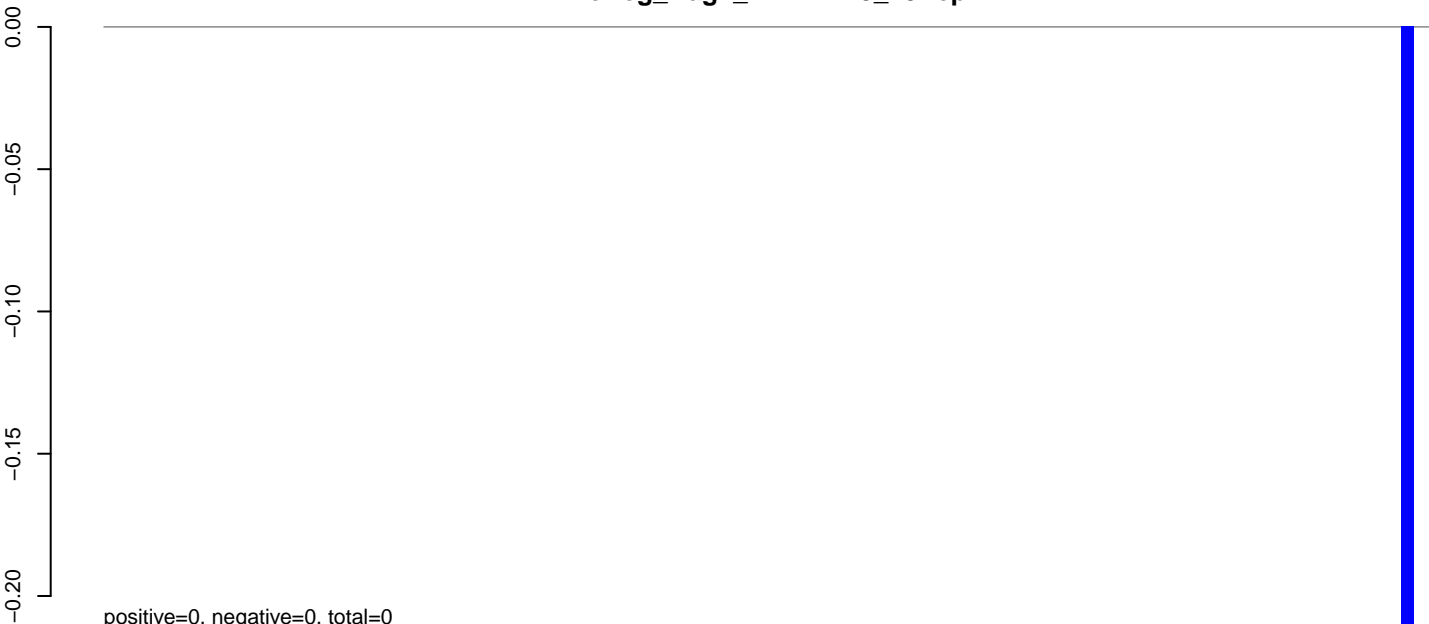


AeAeg_Aag2_DENV4.rep



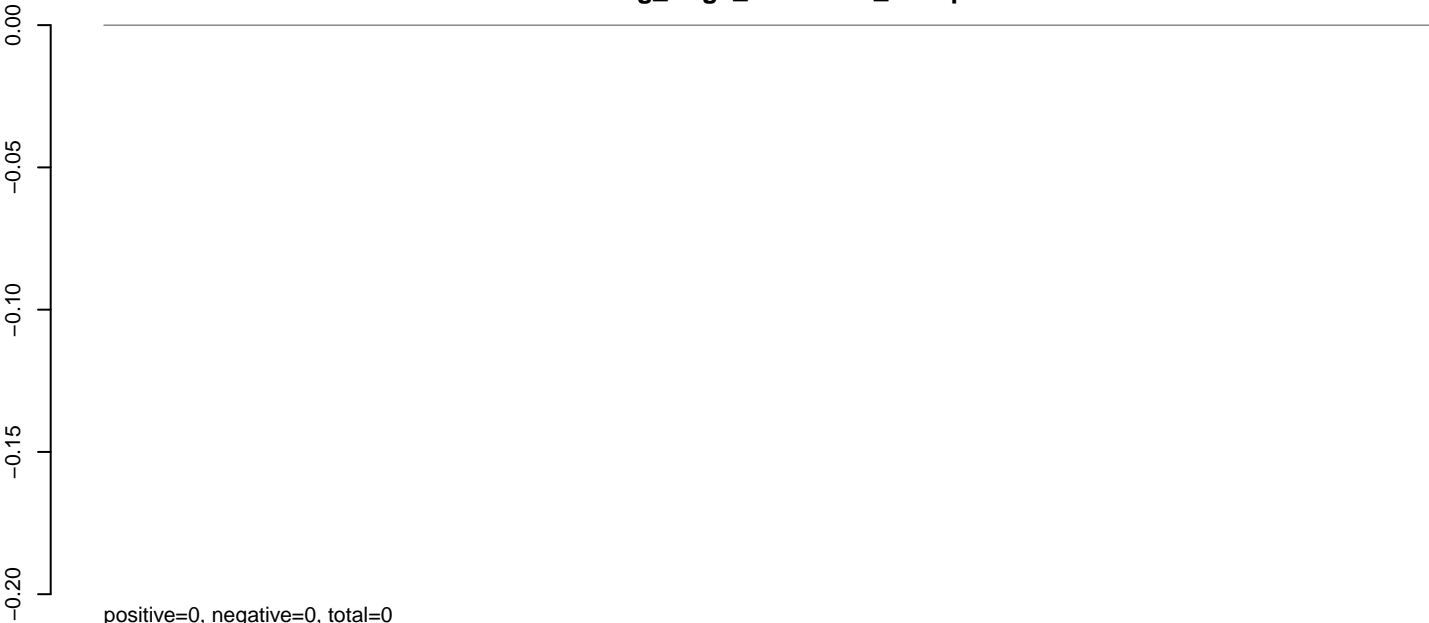
0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV4.18_23.rep



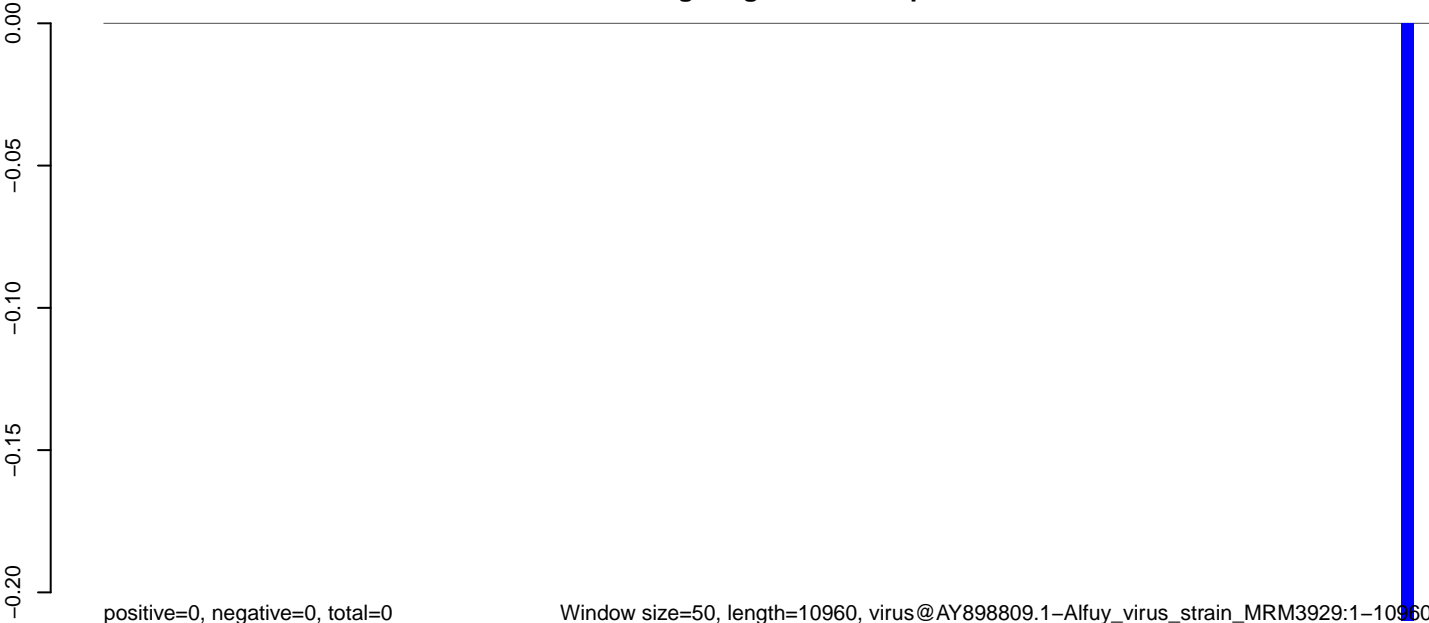
positive=0, negative=0, total=0

AeAeg_Aag2_DENV4.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_DENV4.rep



positive=0, negative=0, total=0

Window size=50, length=10960, virus@AY898809.1-Alfuy_virus_strain_MRM3929:1-10960

0 2000 4000 6000 8000 10000

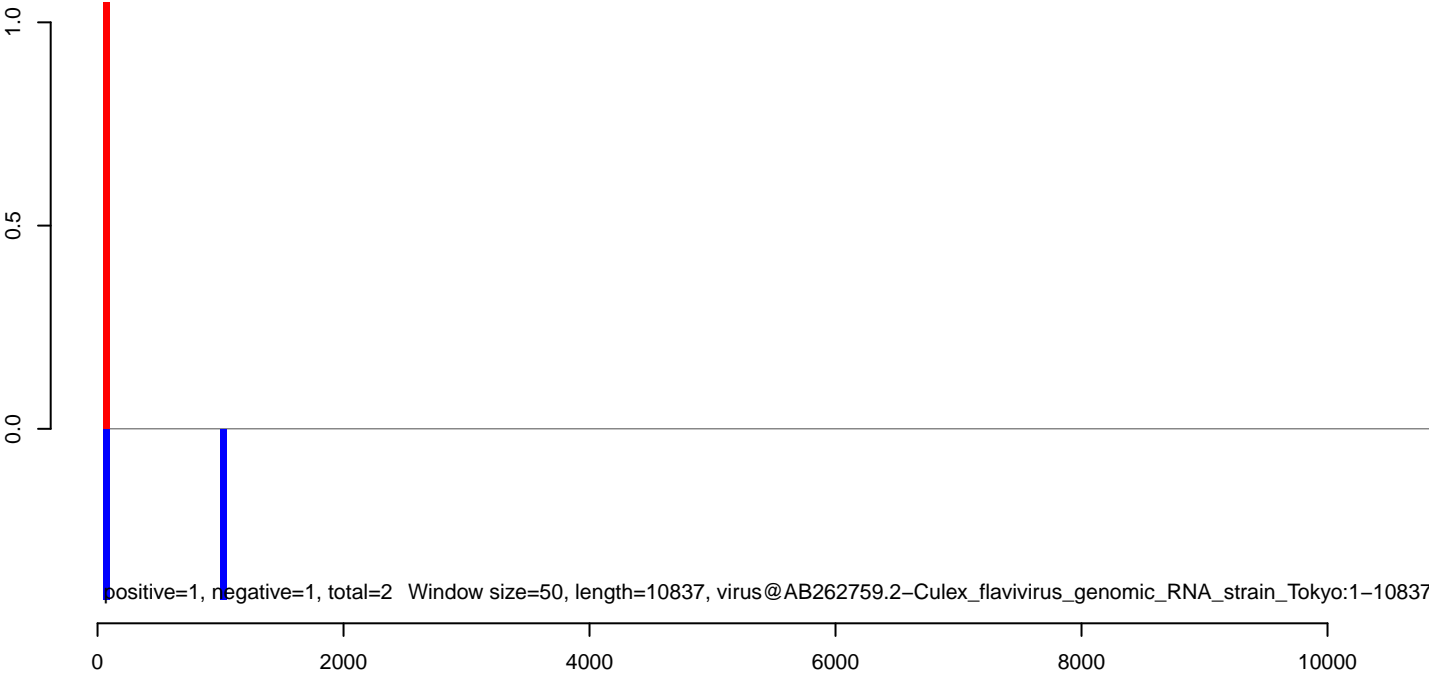
AeAeg_Aag2_DENV4.18_23.rep



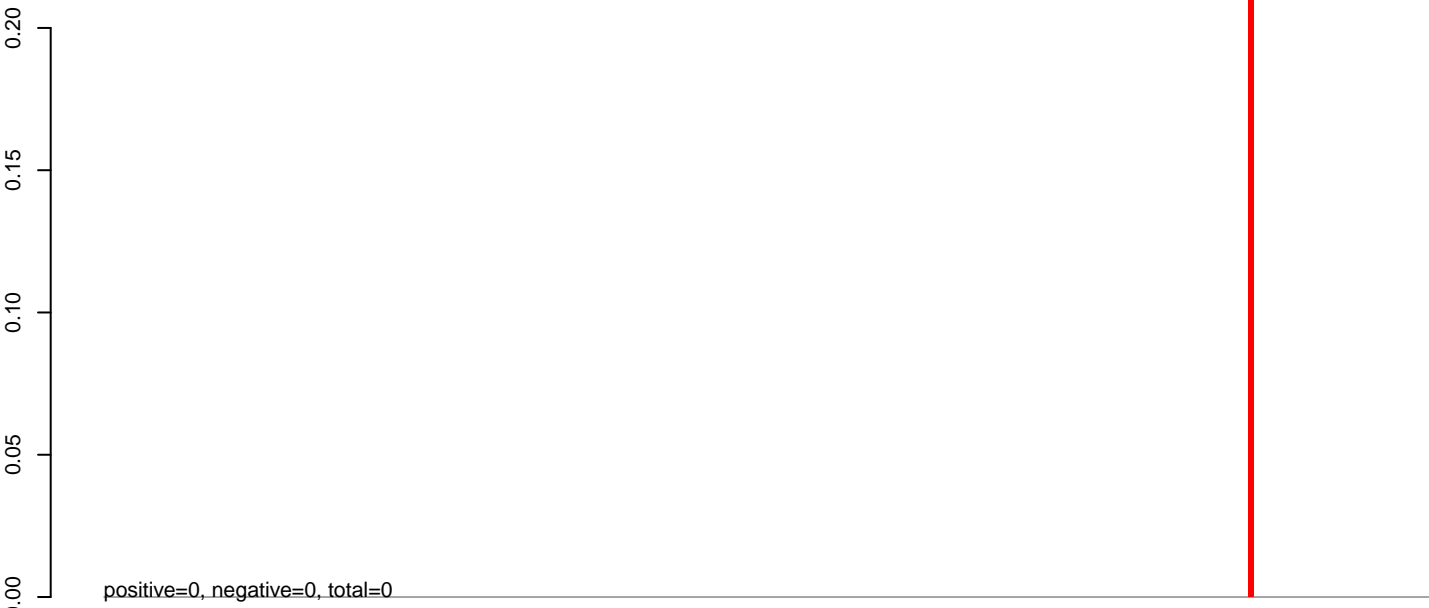
AeAeg_Aag2_DENV4.24_35.rep



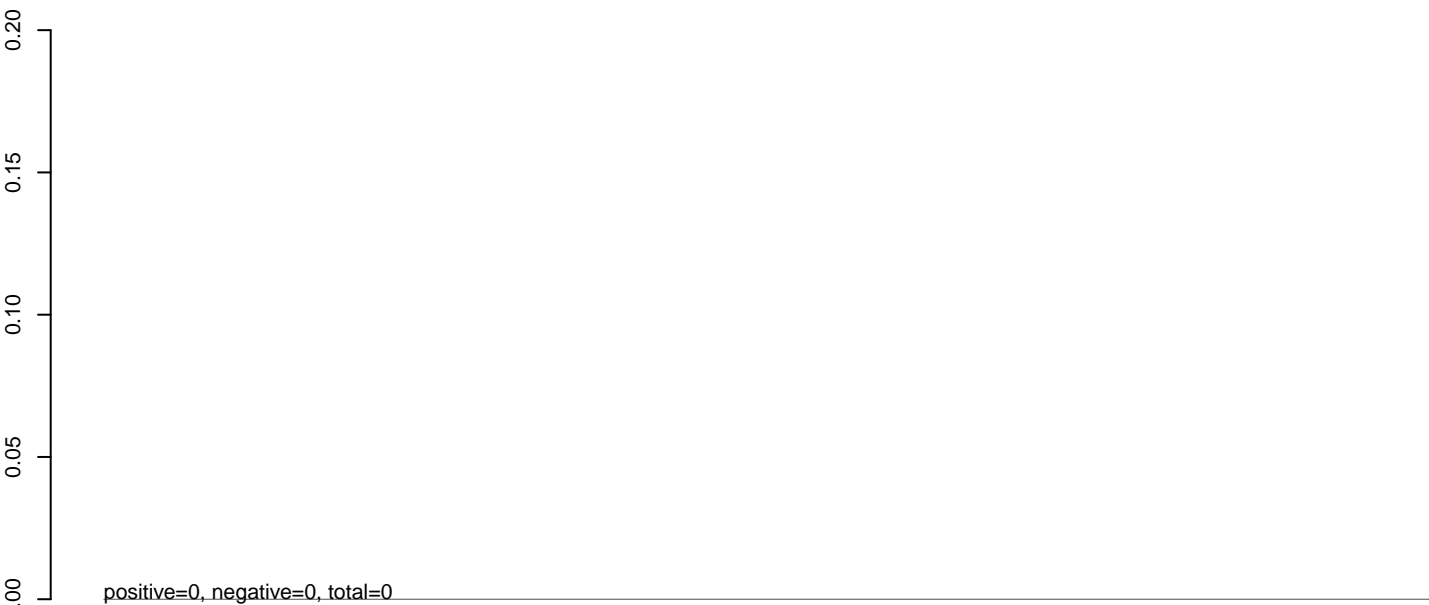
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



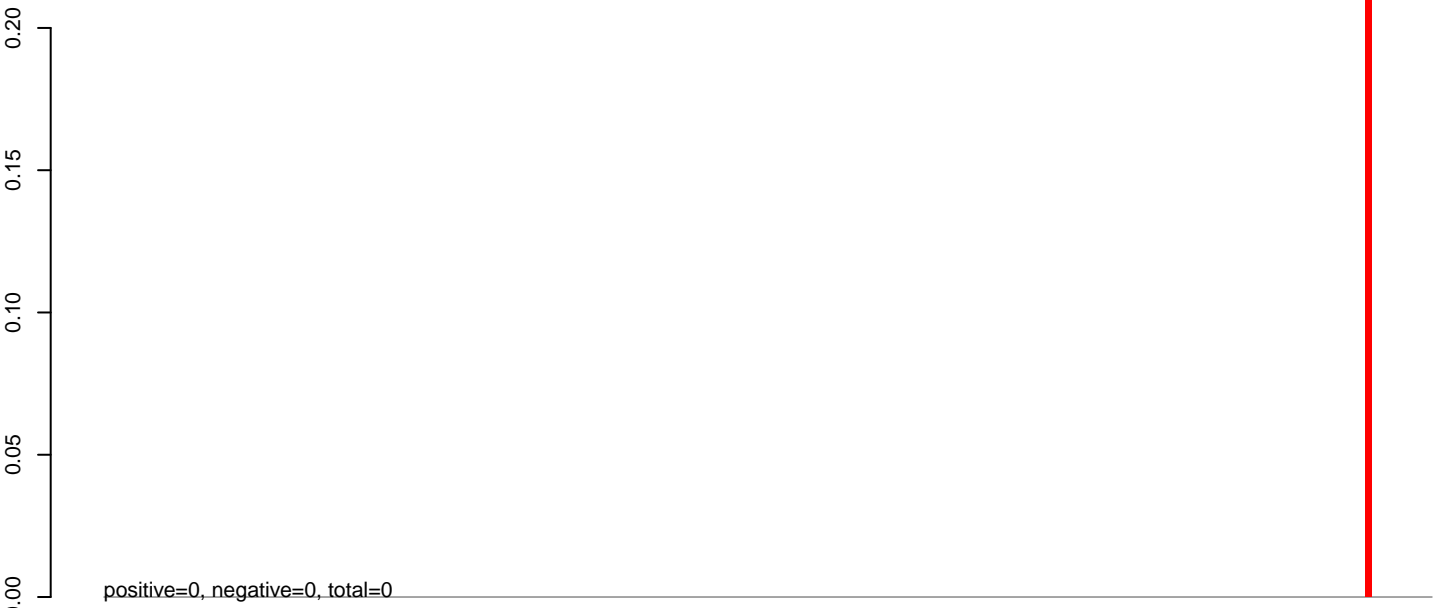
AeAeg_Aag2_DENV4.24_35.rep



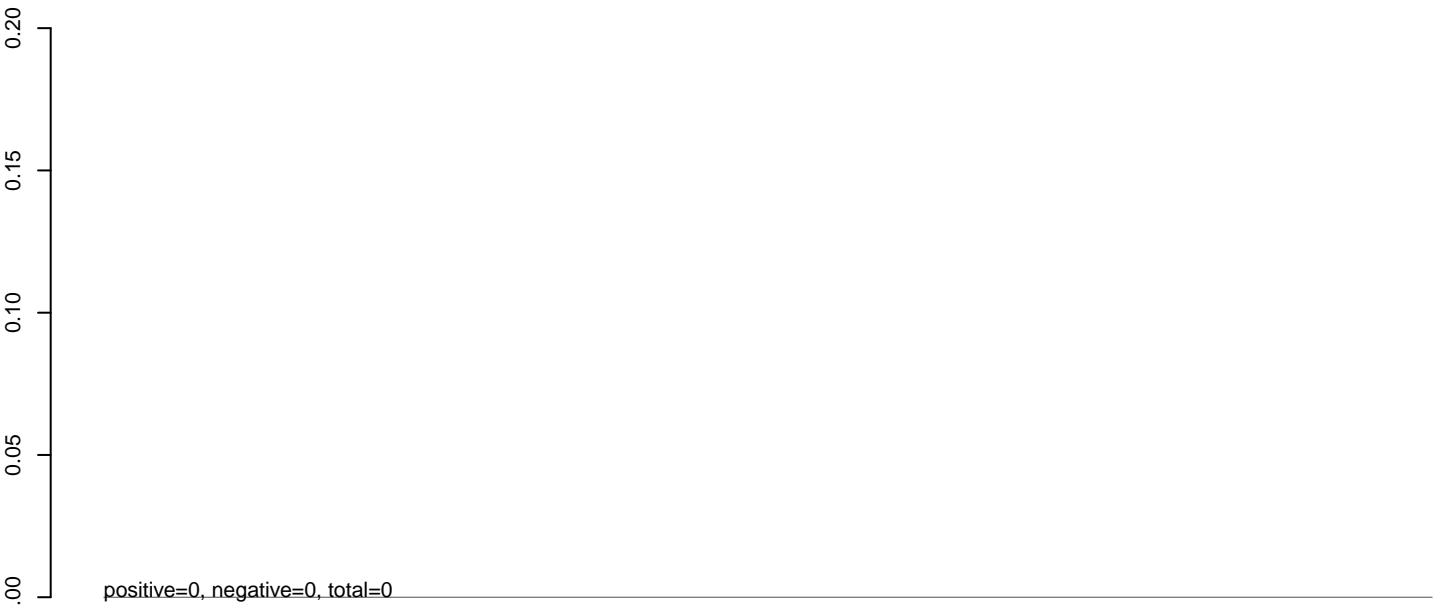
AeAeg_Aag2_DENV4.rep



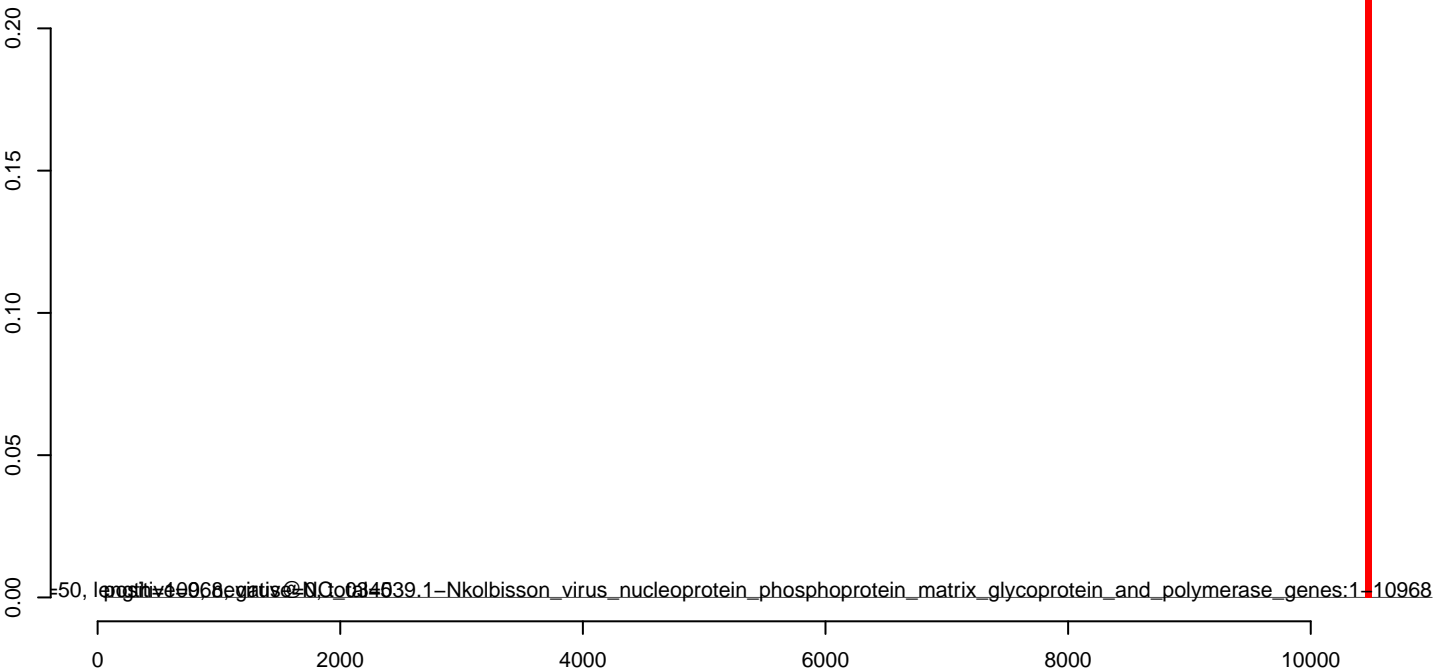
AeAeg_Aag2_DENV4.18_23.rep



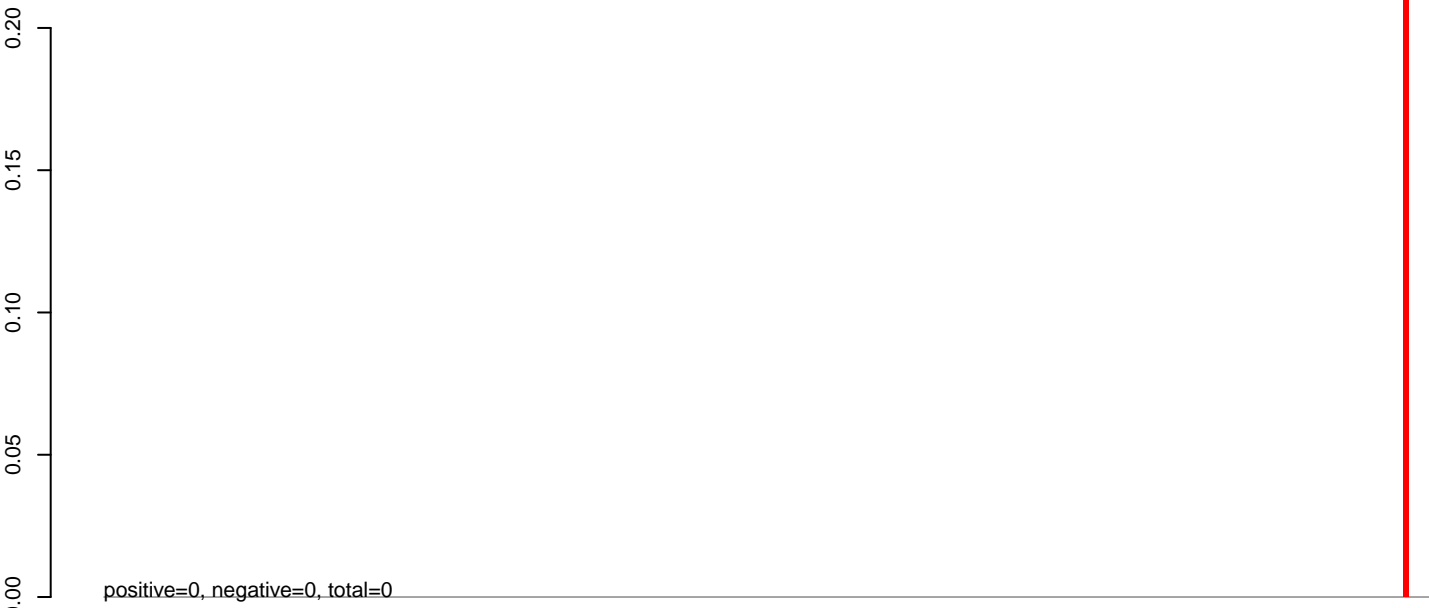
AeAeg_Aag2_DENV4.24_35.rep



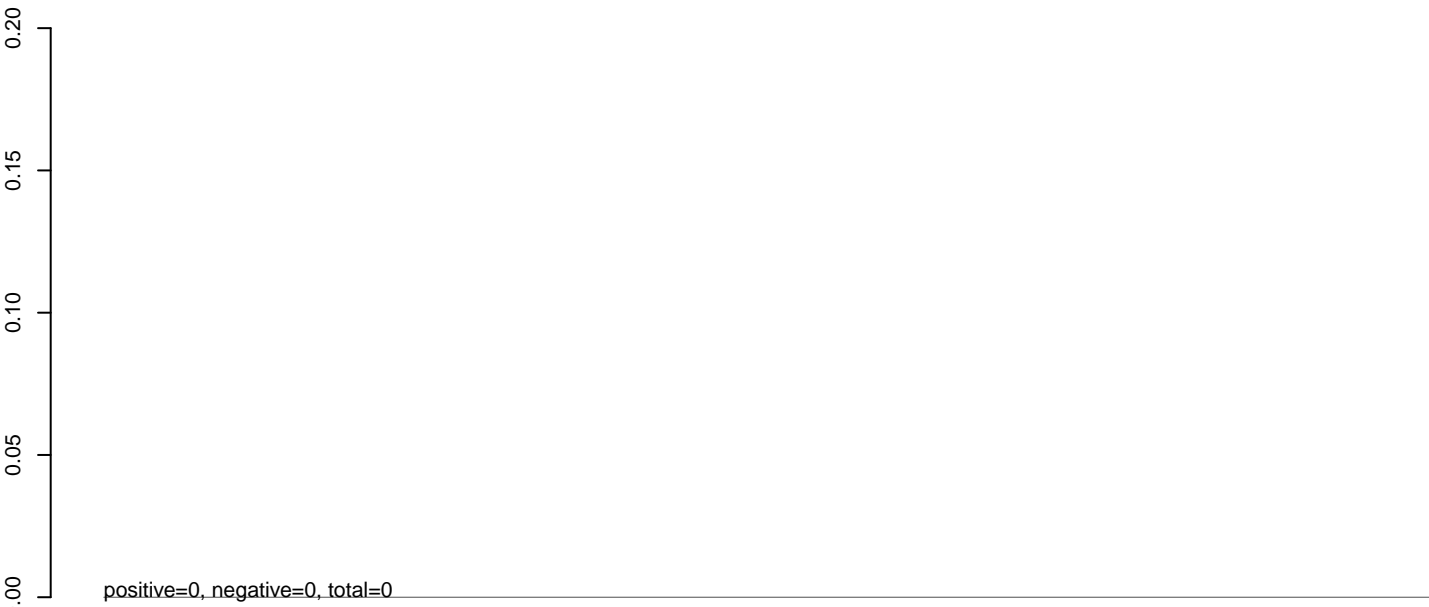
AeAeg_Aag2_DENV4.rep



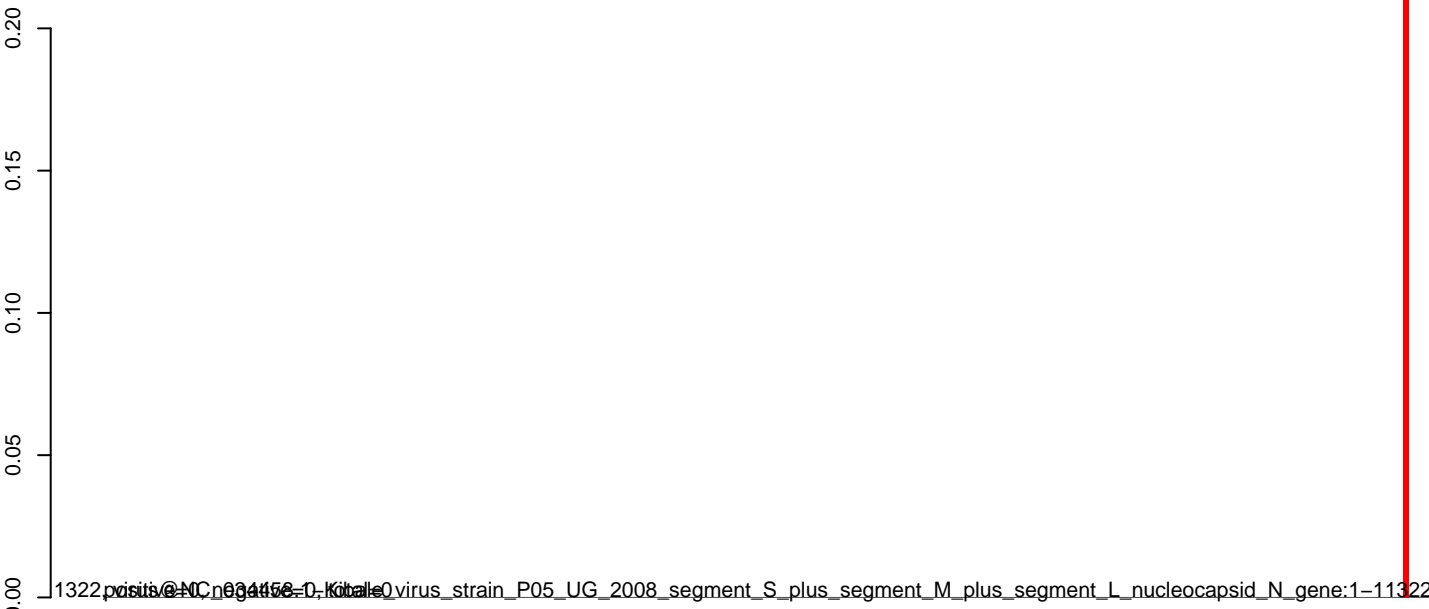
AeAeg_Aag2_DENV4.18_23.rep



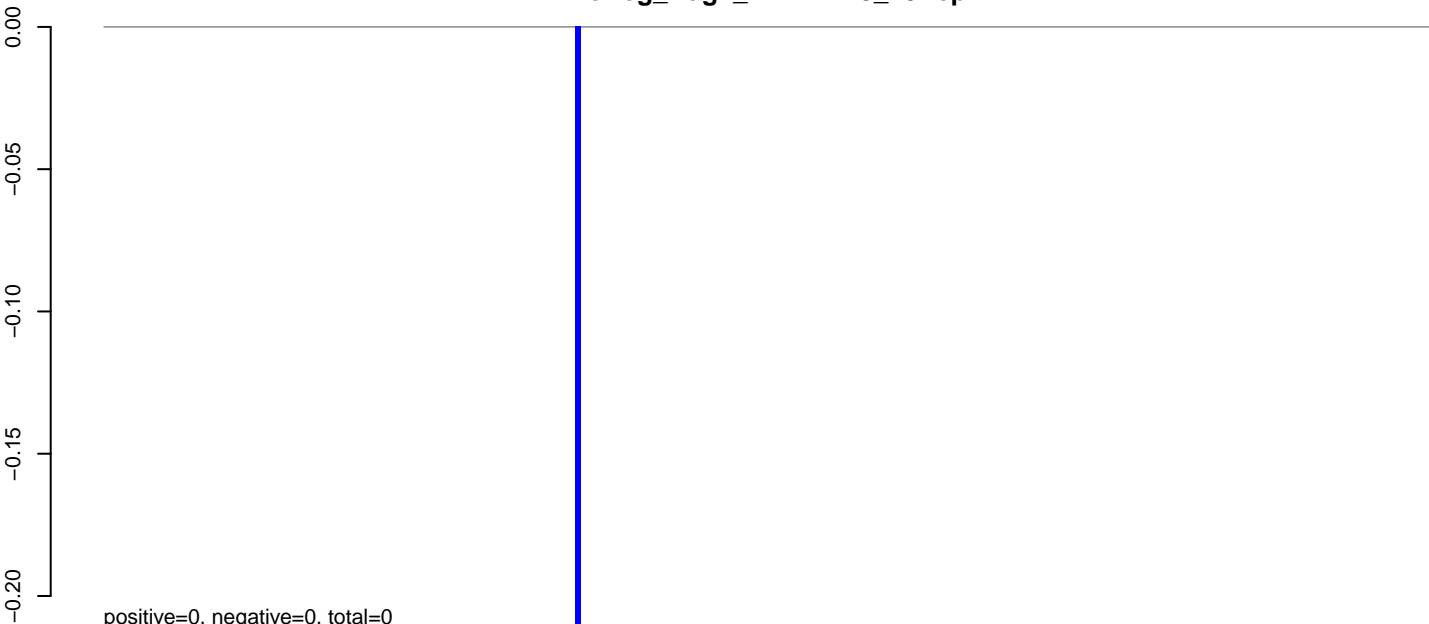
AeAeg_Aag2_DENV4.24_35.rep



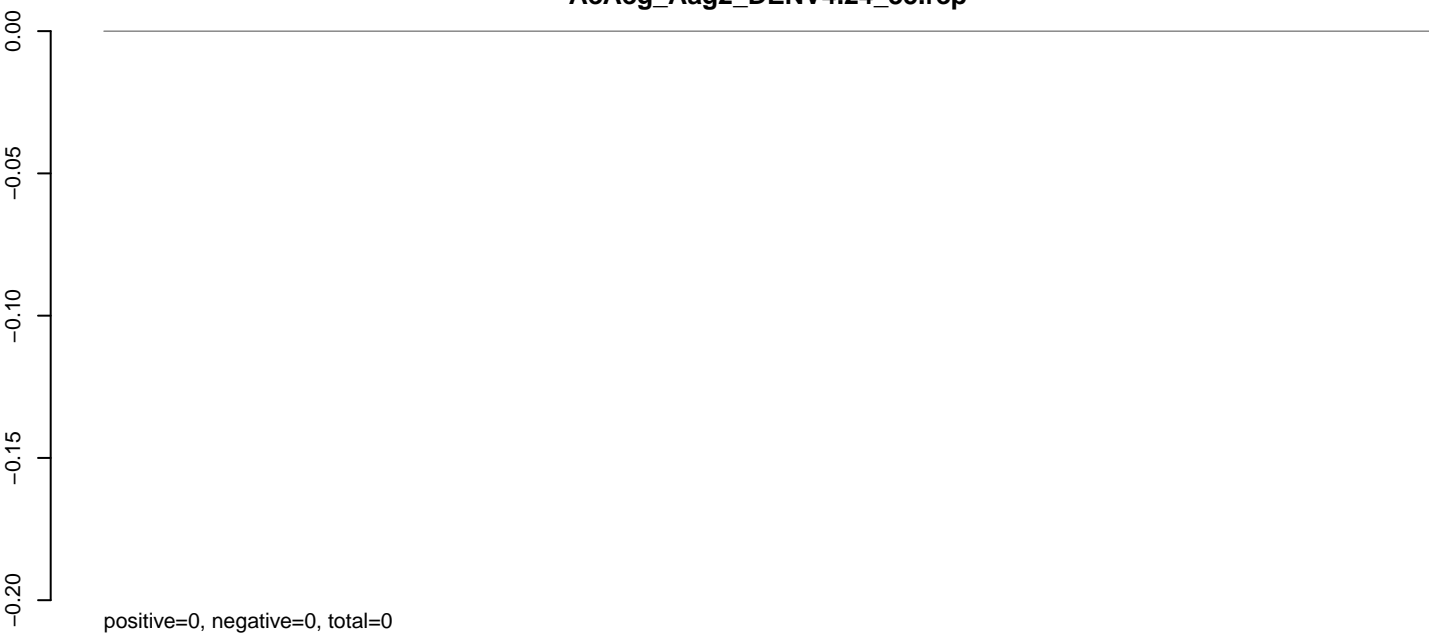
AeAeg_Aag2_DENV4.rep



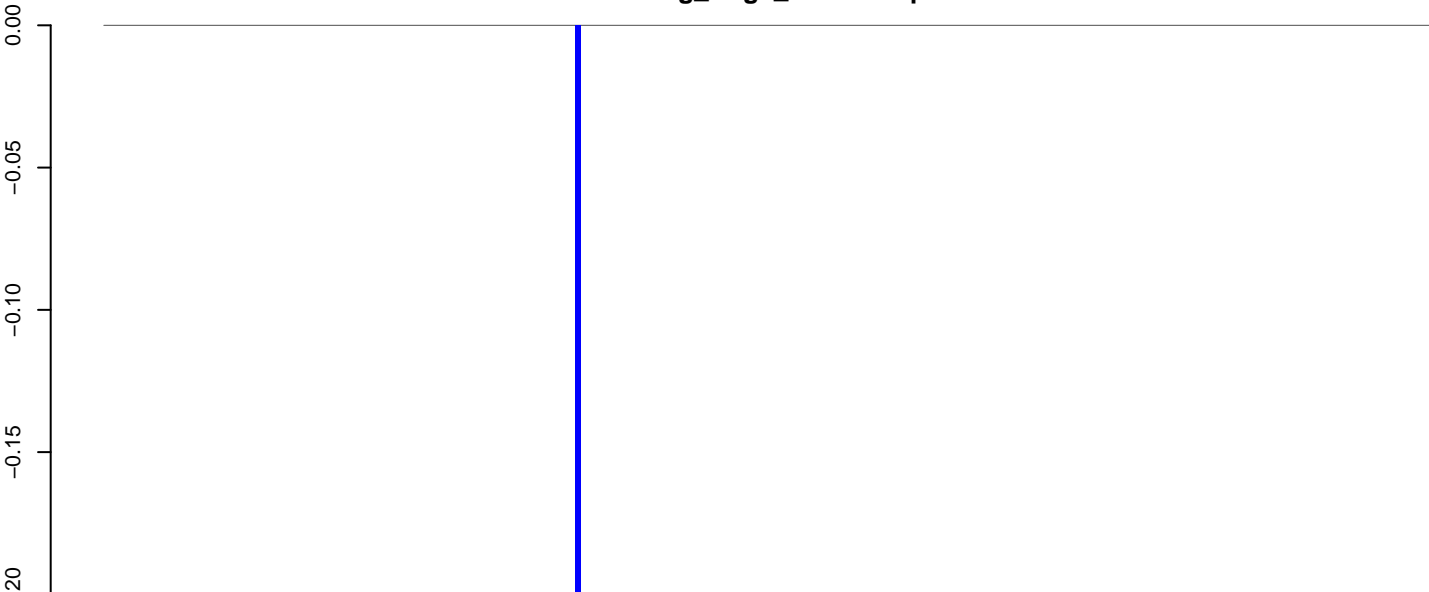
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



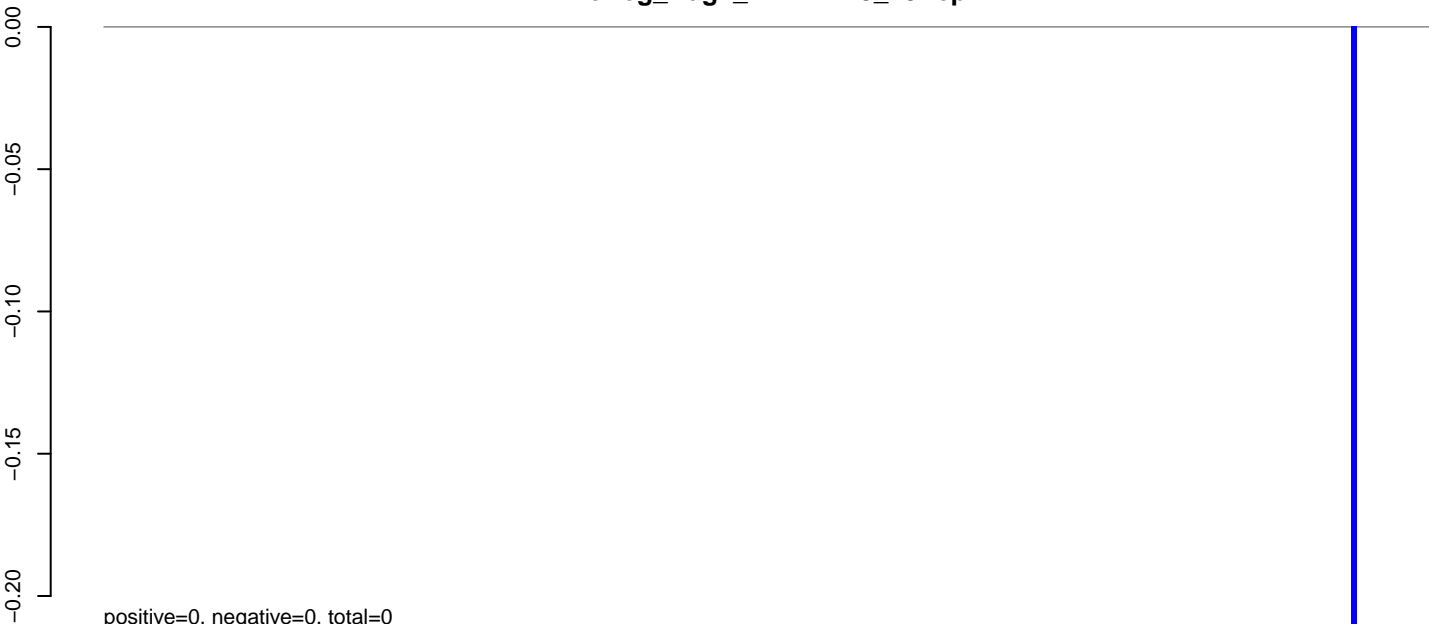
AeAeg_Aag2_DENV4.rep



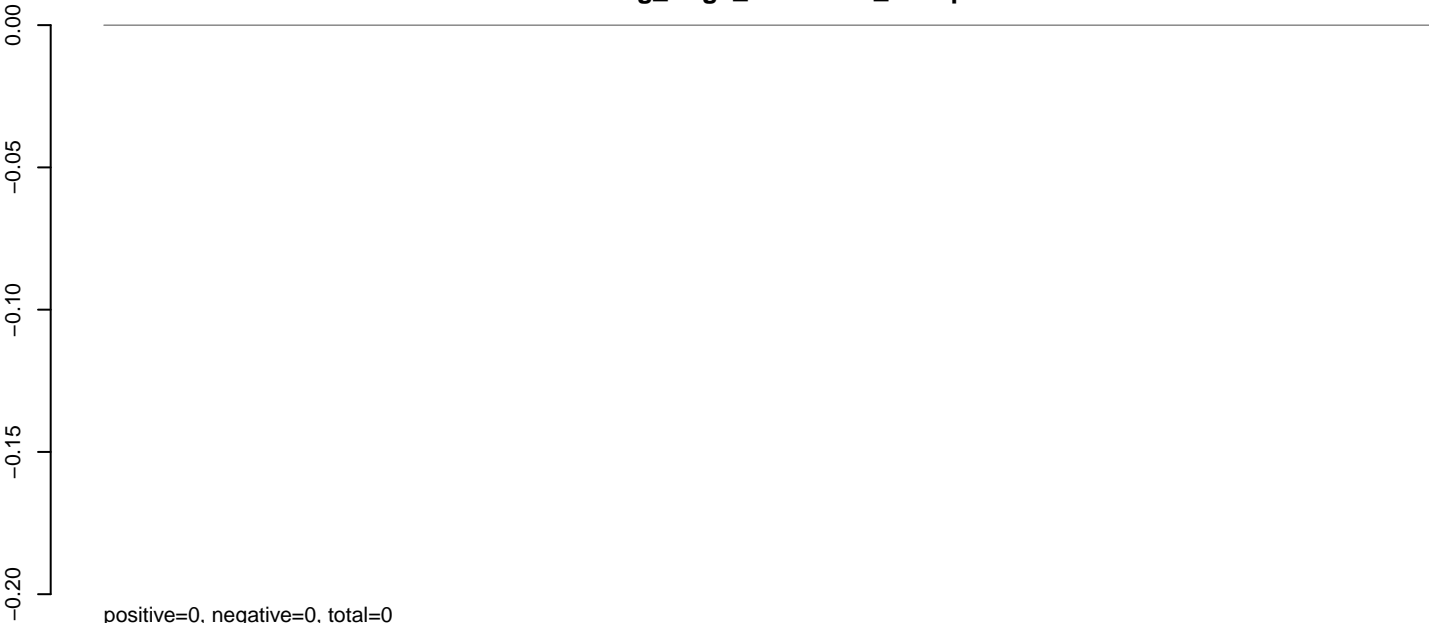
path=11728_virus_0103405701_virus_strain_F47_CI_2004_segment_S_plus_segment_M_plus_segment_L_nucleocapsid_N_gene:1-11728

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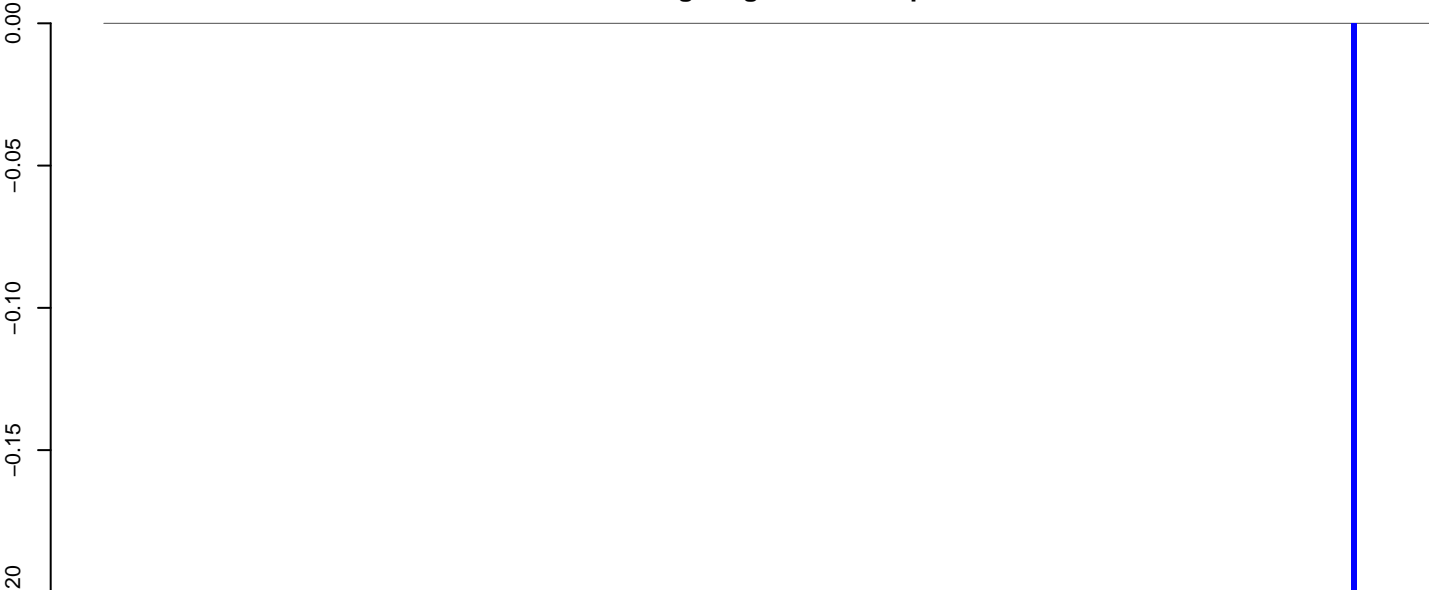
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



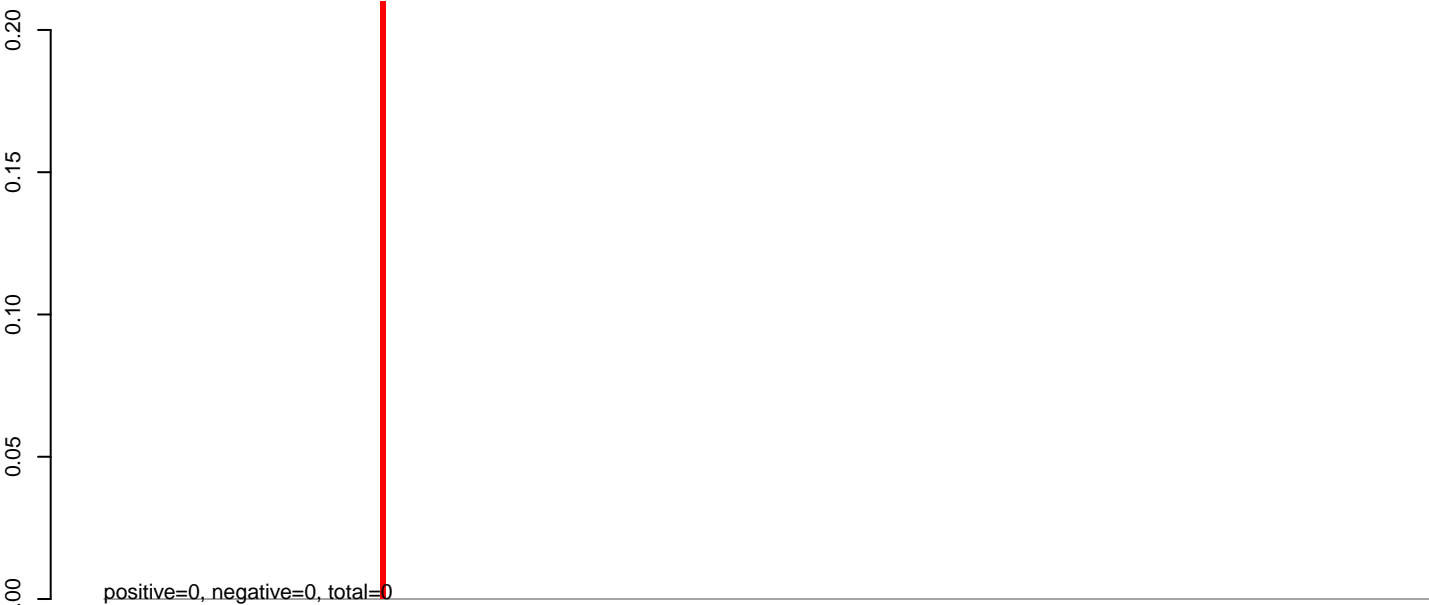
AeAeg_Aag2_DENV4.rep



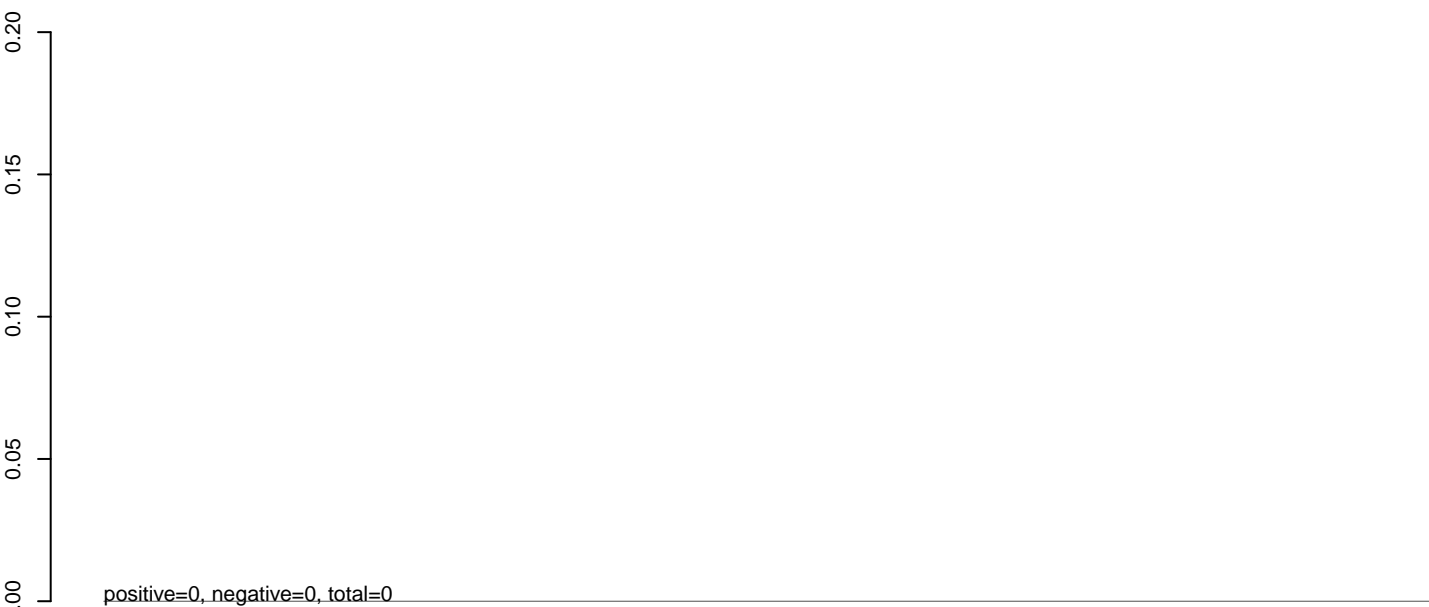
-Kamperlingus_megaprotiles phosphoprotein_hypothetical_proteins_matrix_glycoprotein_hypothetical_protein_and_polymerase_genes:1-13209

0 2000 4000 6000 8000 10000 12000

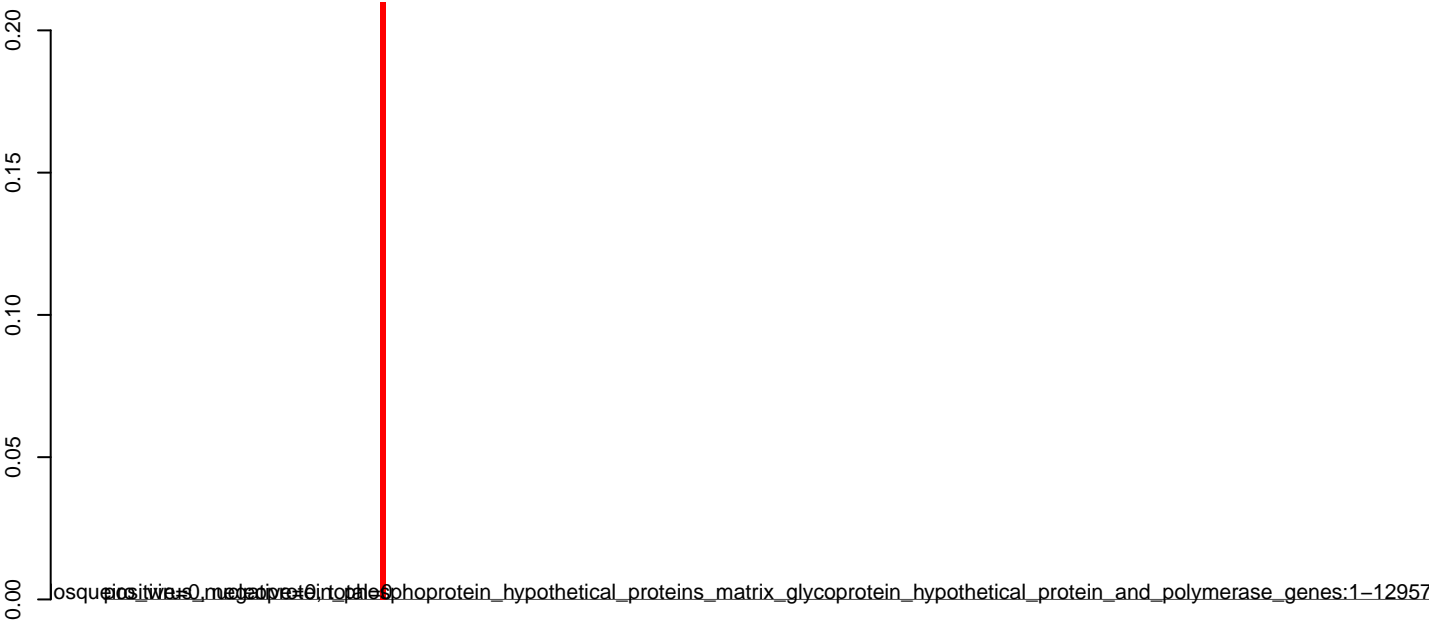
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



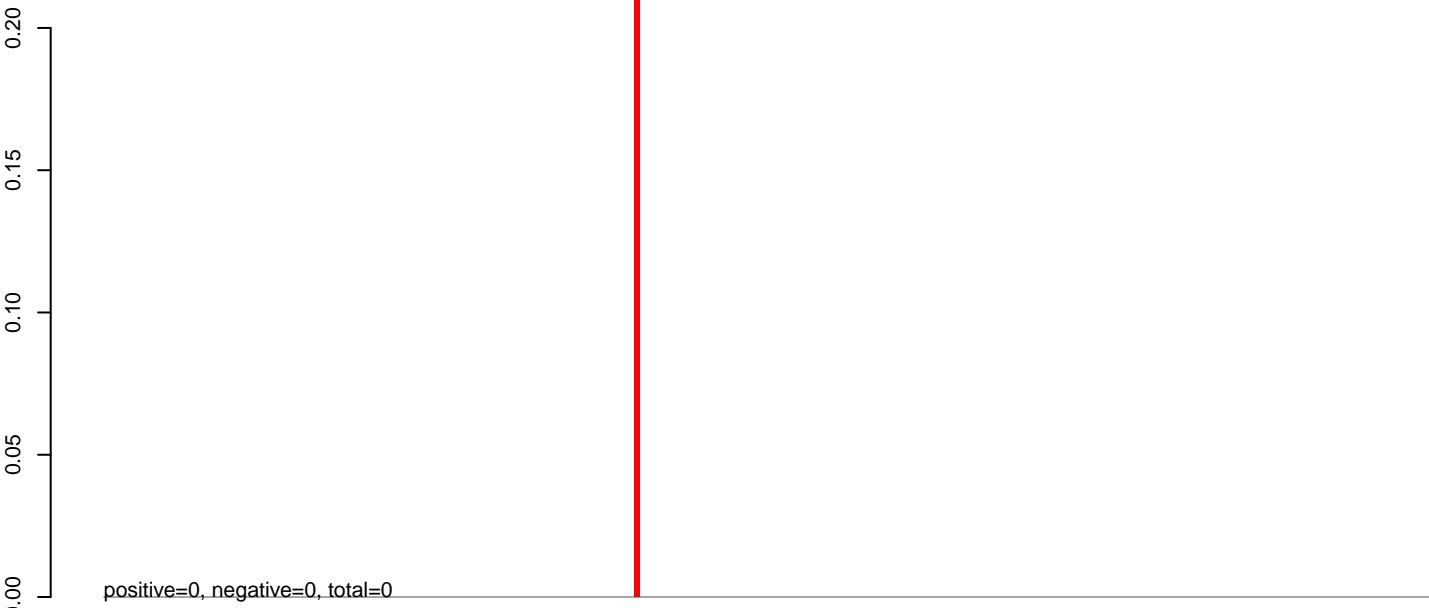
AeAeg_Aag2_DENV4.rep



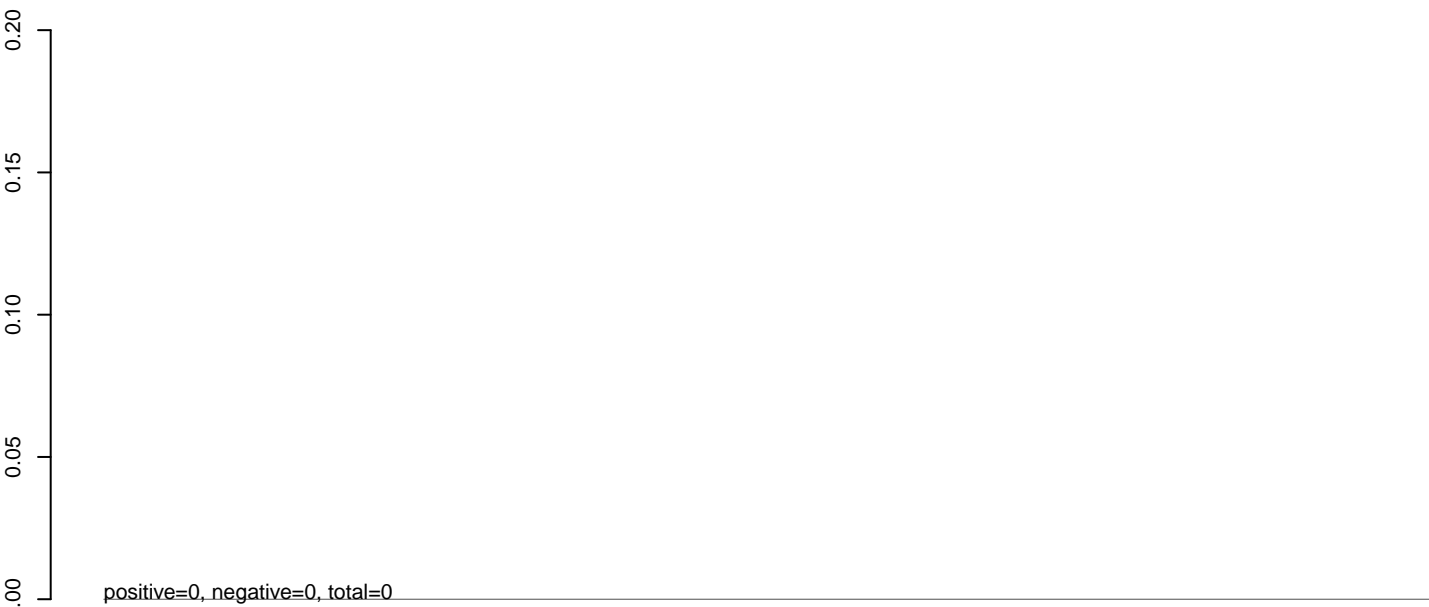
osquepinos, megalopods, and other crustaceans; phosphoprotein_hypothetical_proteins_matrix_glycoprotein_hypothetical_protein_and_polymerase_genes:1-12957

0 2000 4000 6000 8000 10000 12000

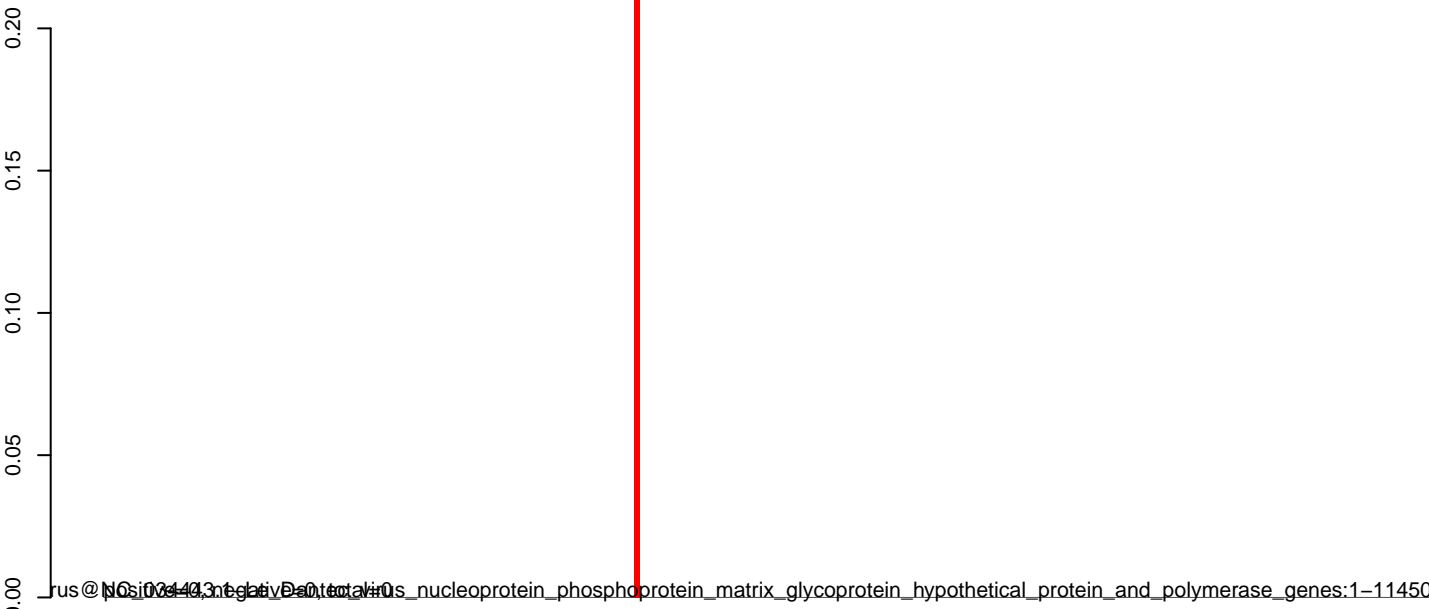
AeAeg_Aag2_DENV4.18_23.rep



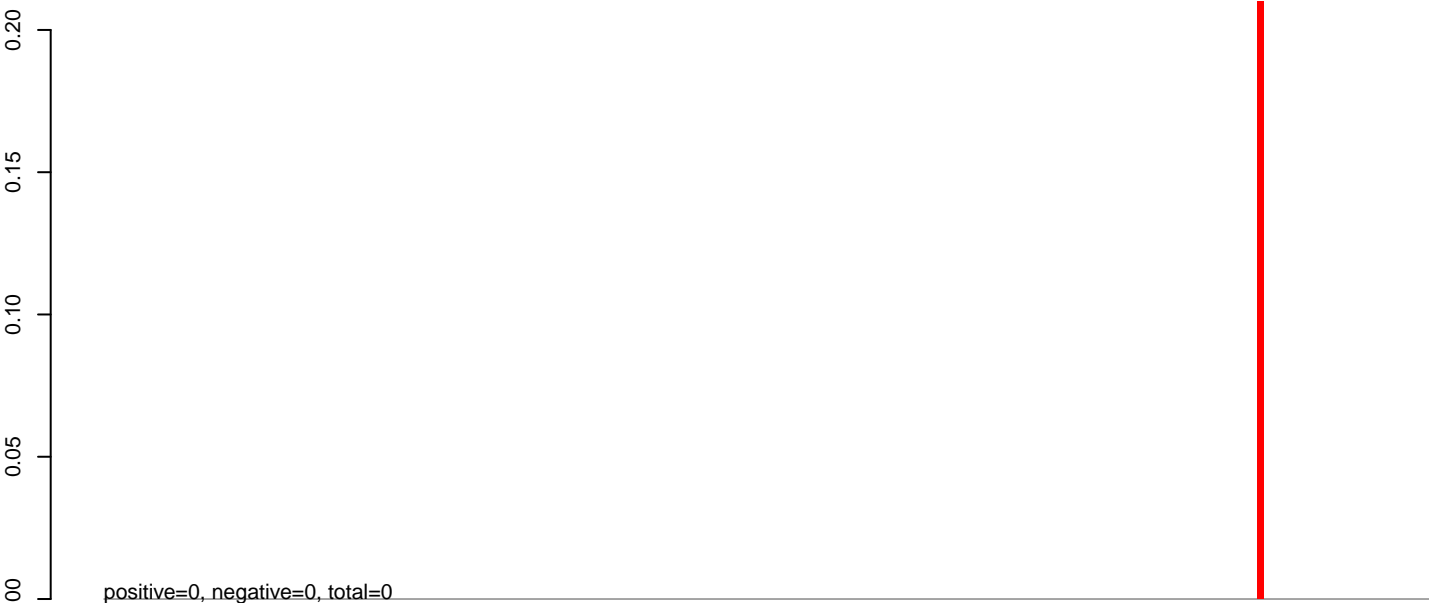
AeAeg_Aag2_DENV4.24_35.rep



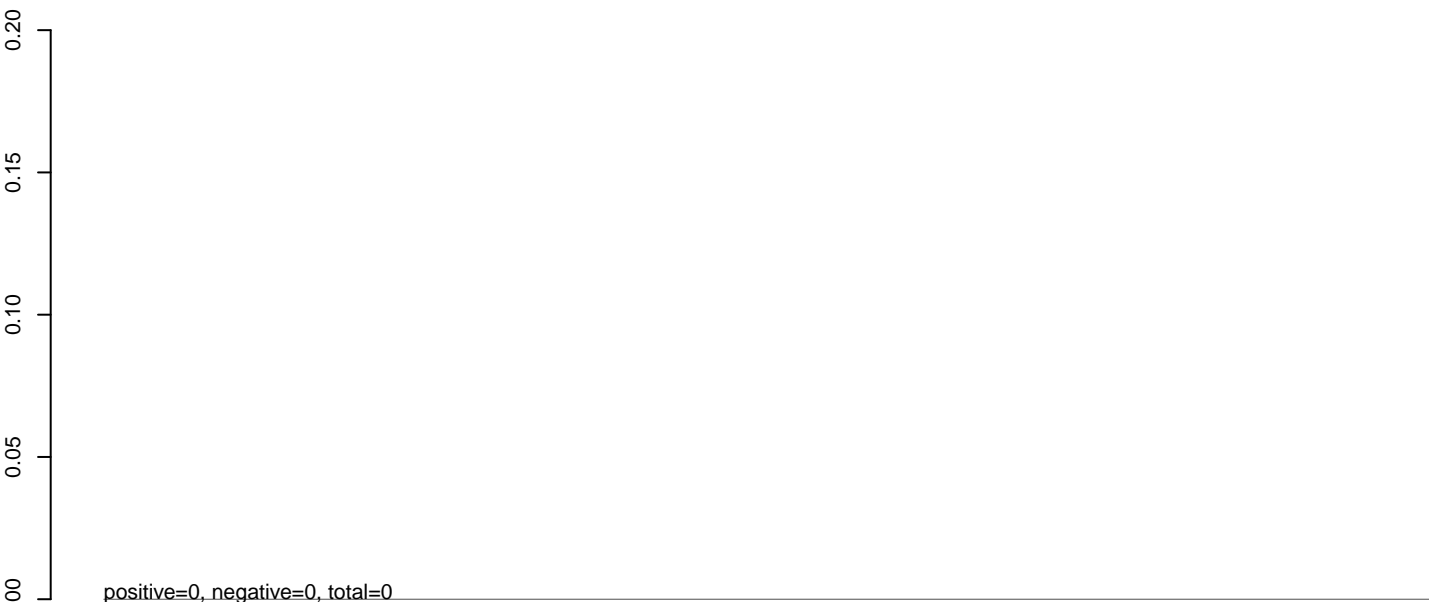
AeAeg_Aag2_DENV4.rep



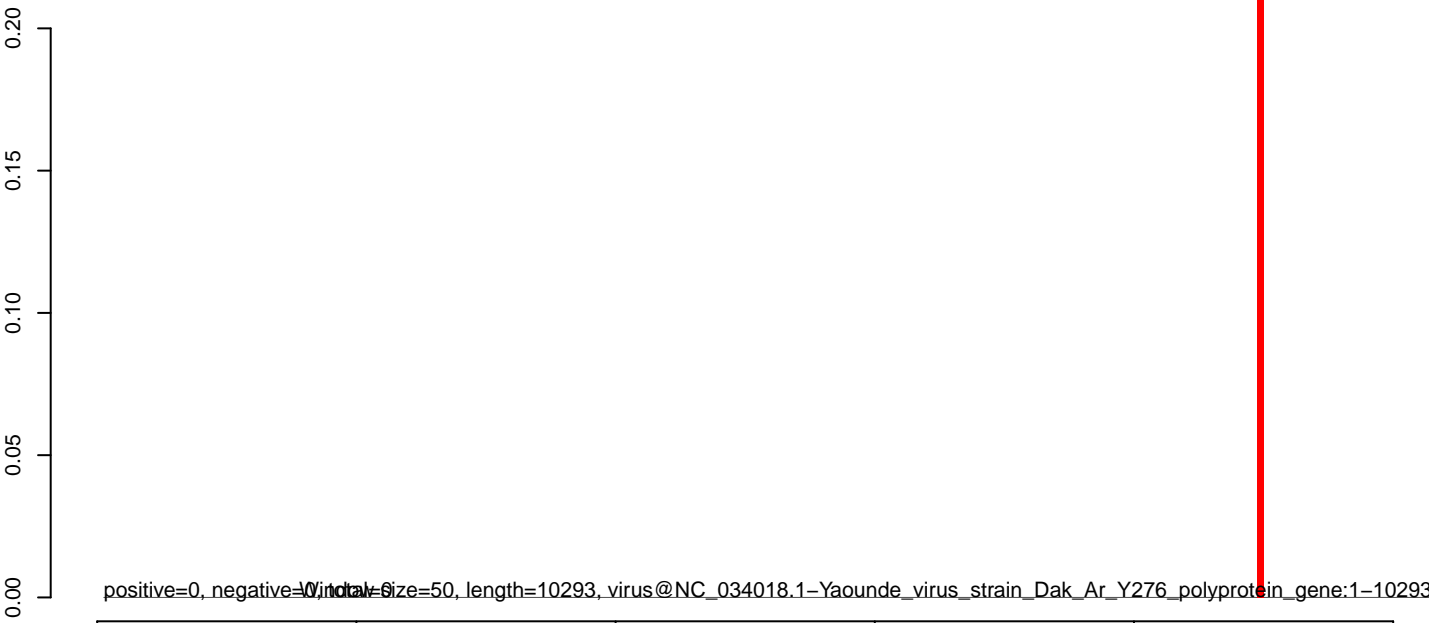
AeAeg_Aag2_DENV4.18_23.rep



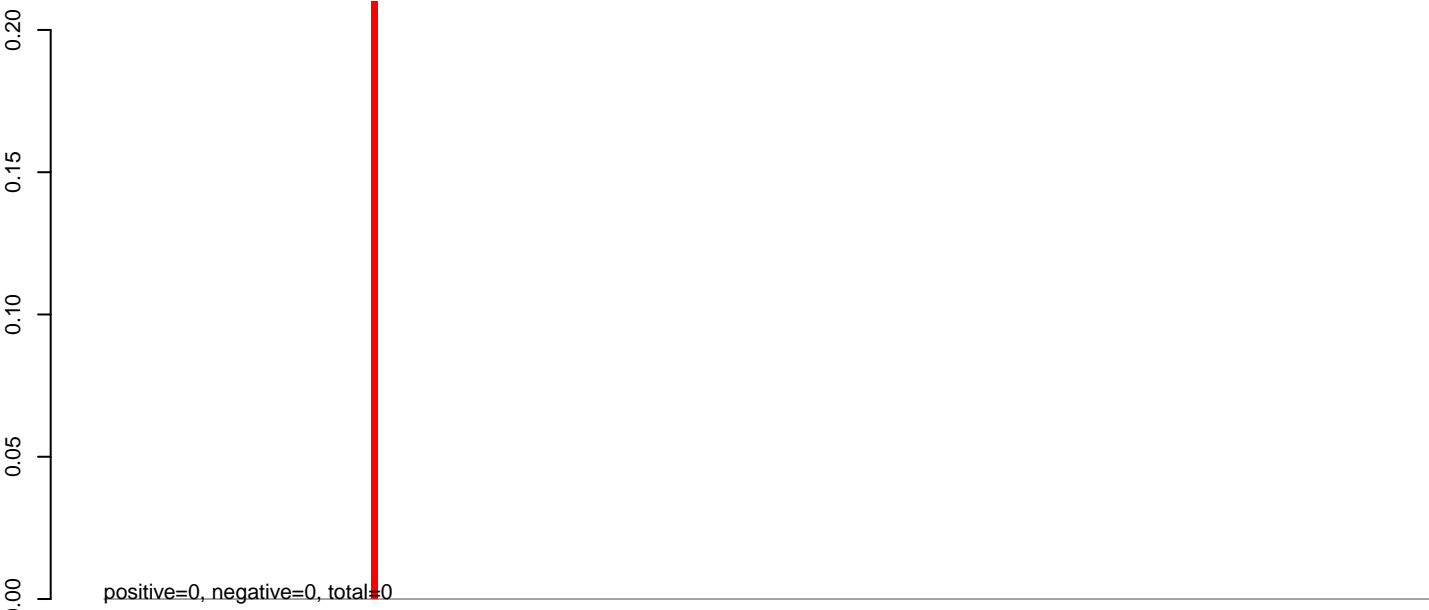
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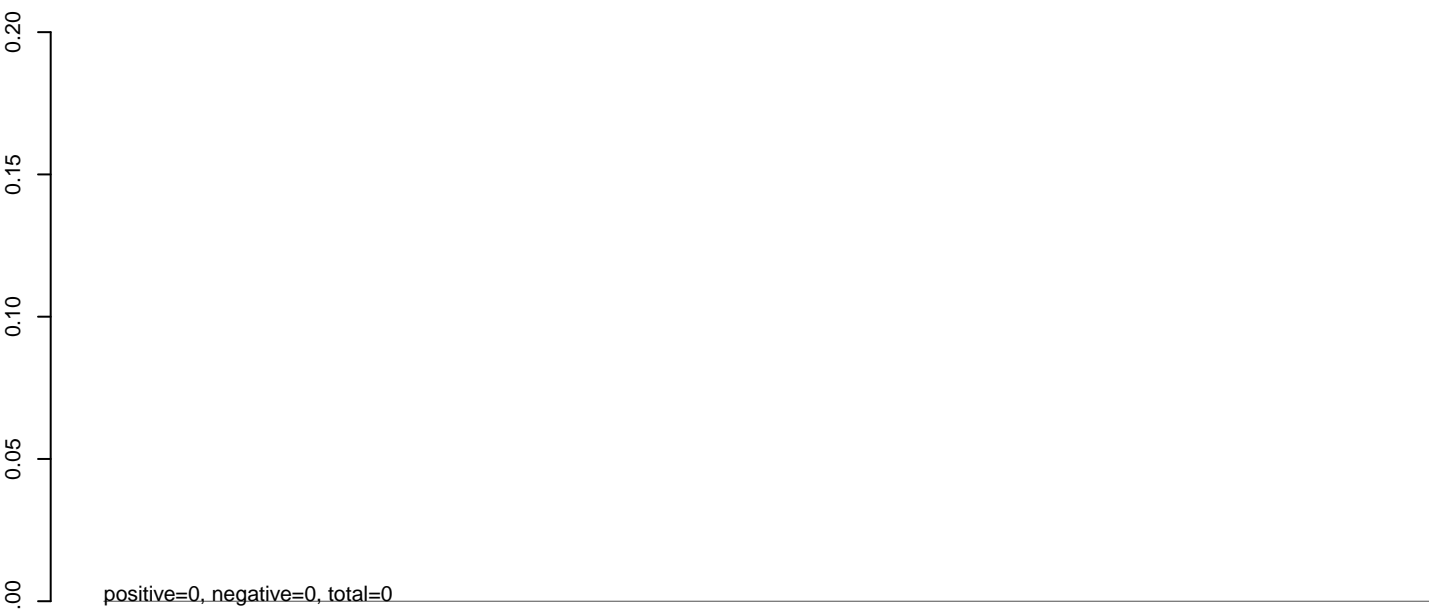
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



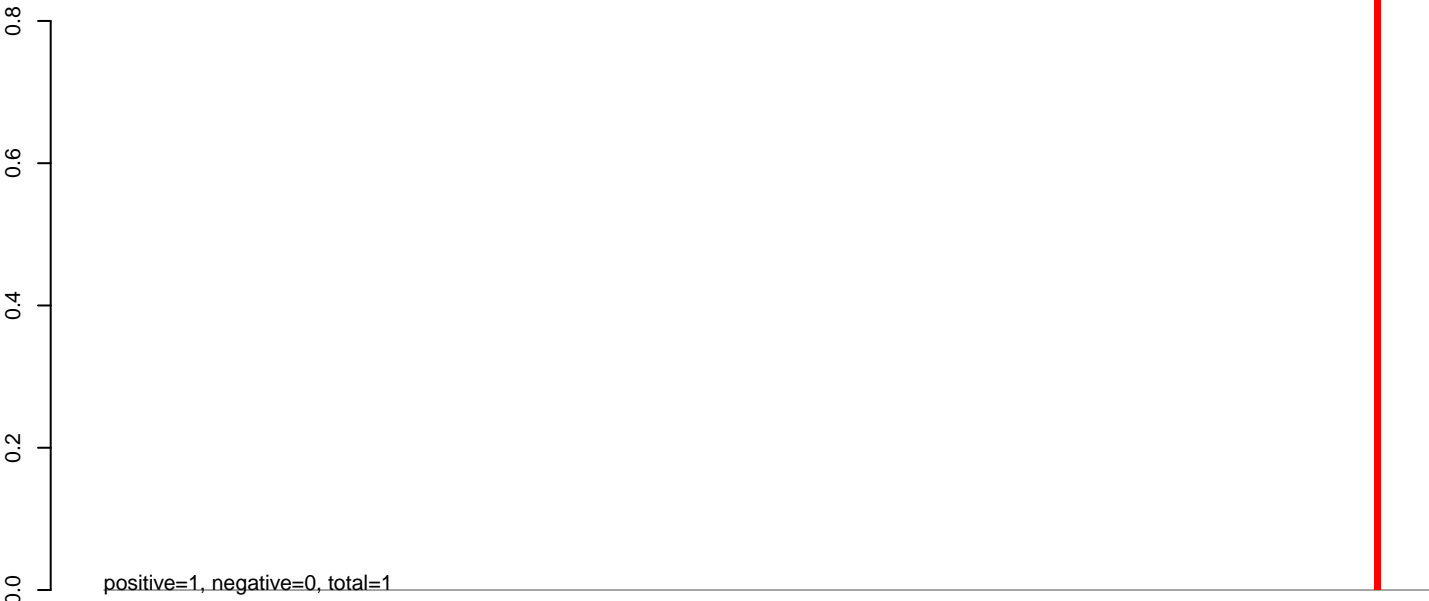
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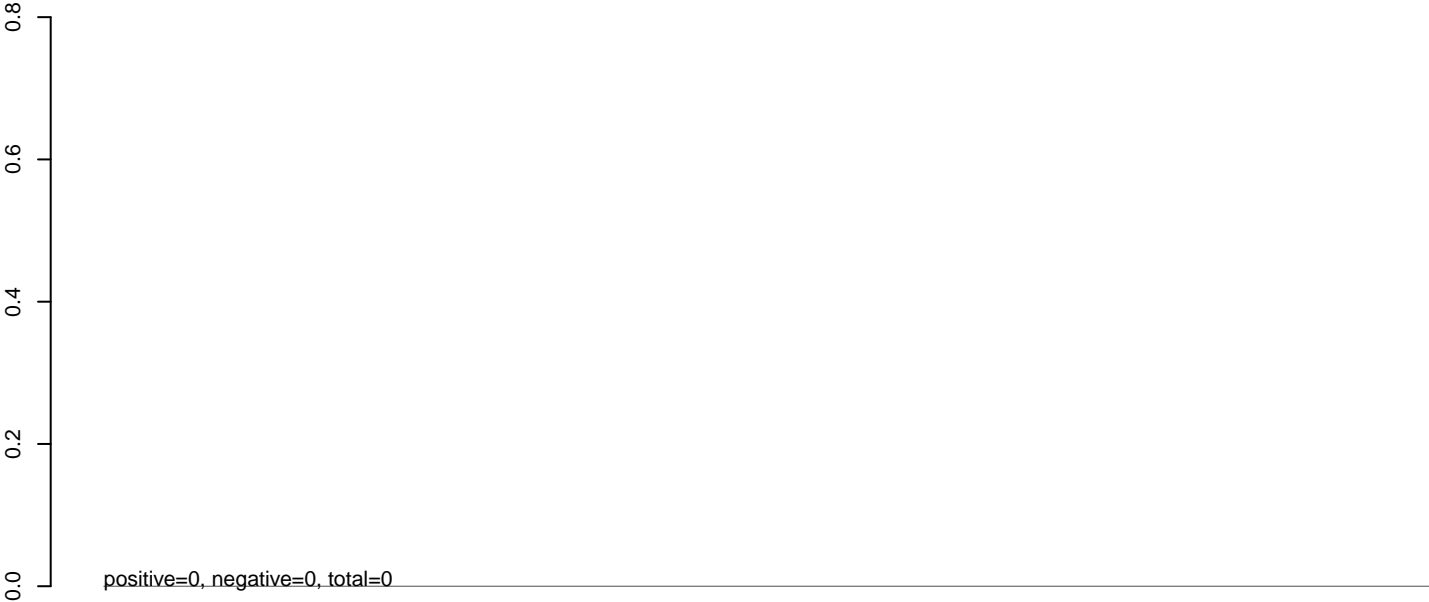
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



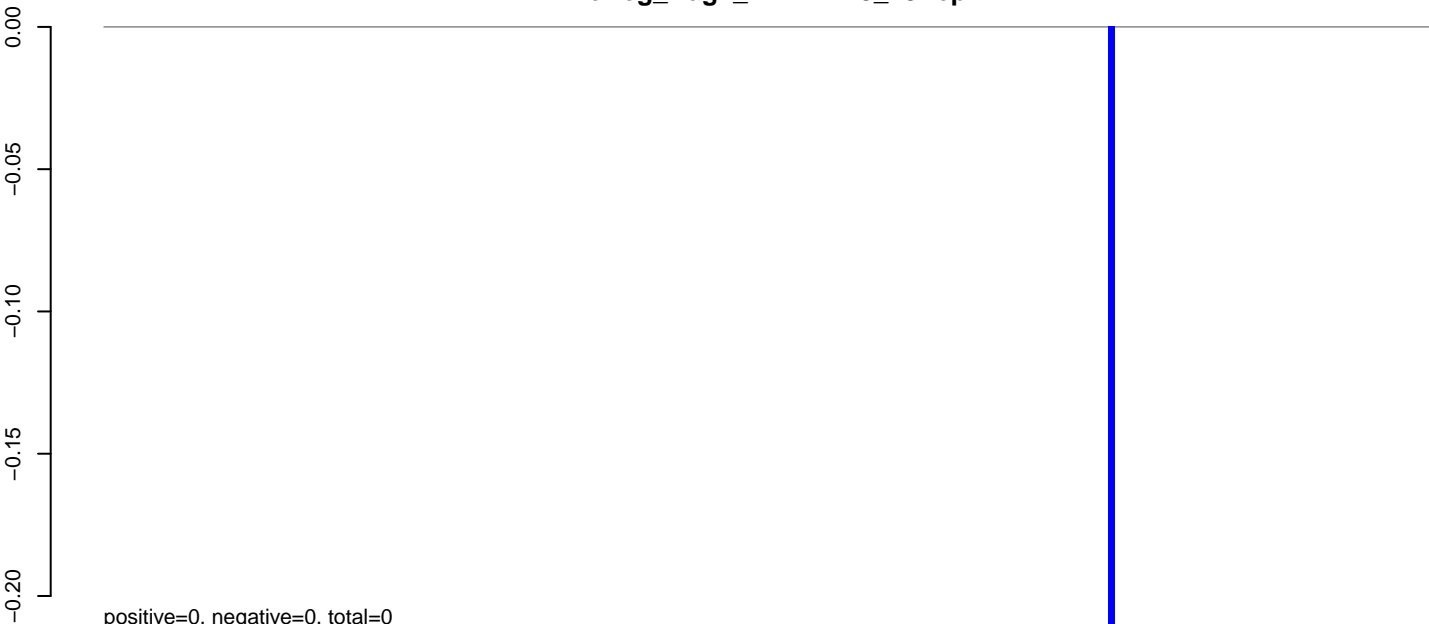
AeAeg_Aag2_DENV4.24_35.rep



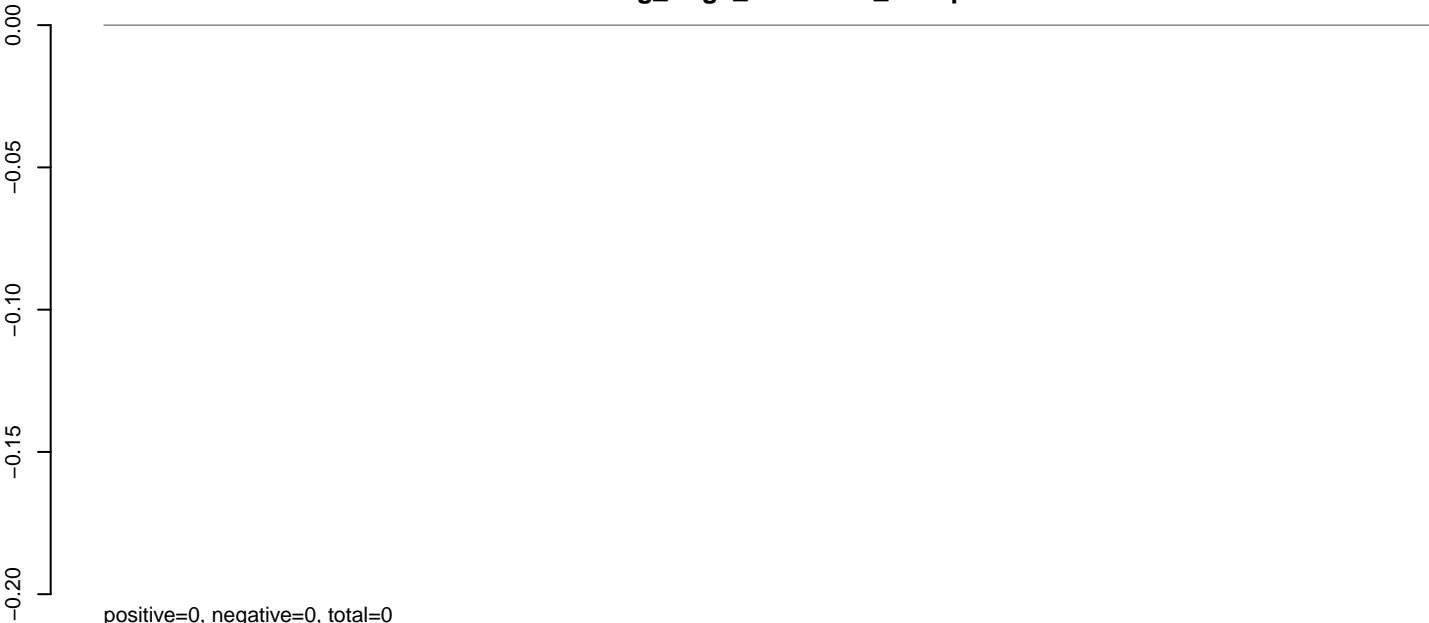
AeAeg_Aag2_DENV4.rep



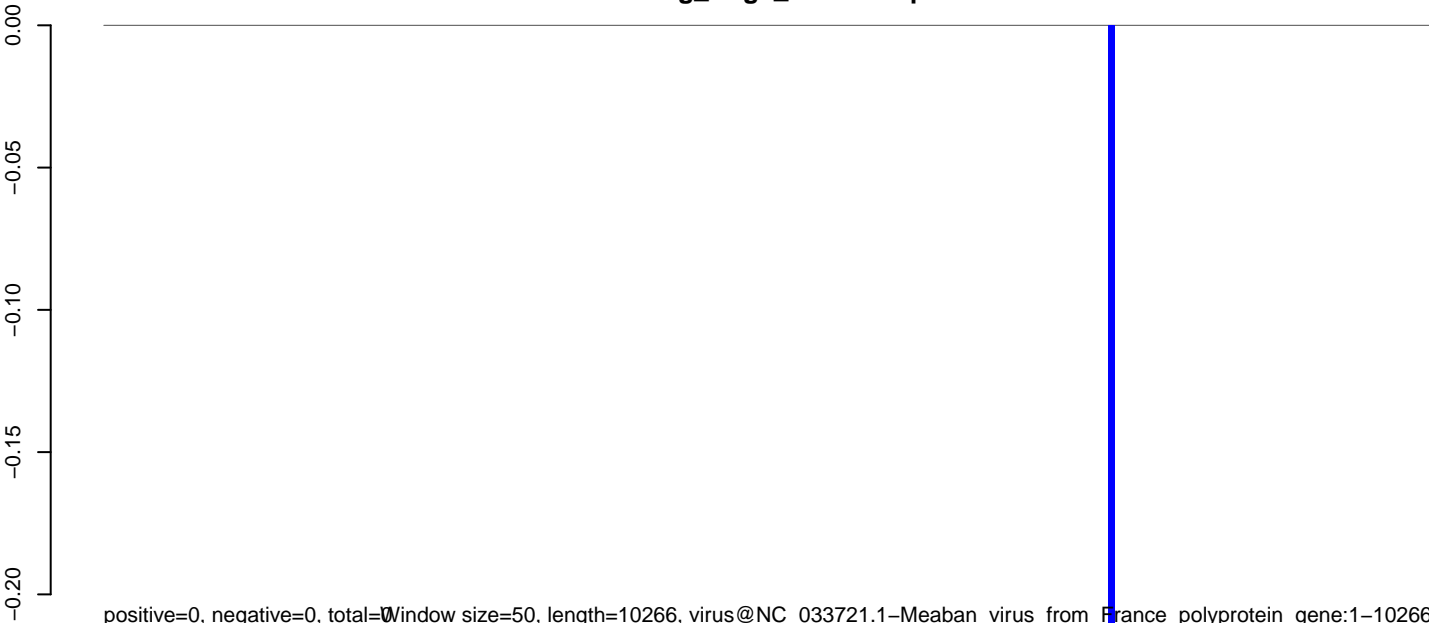
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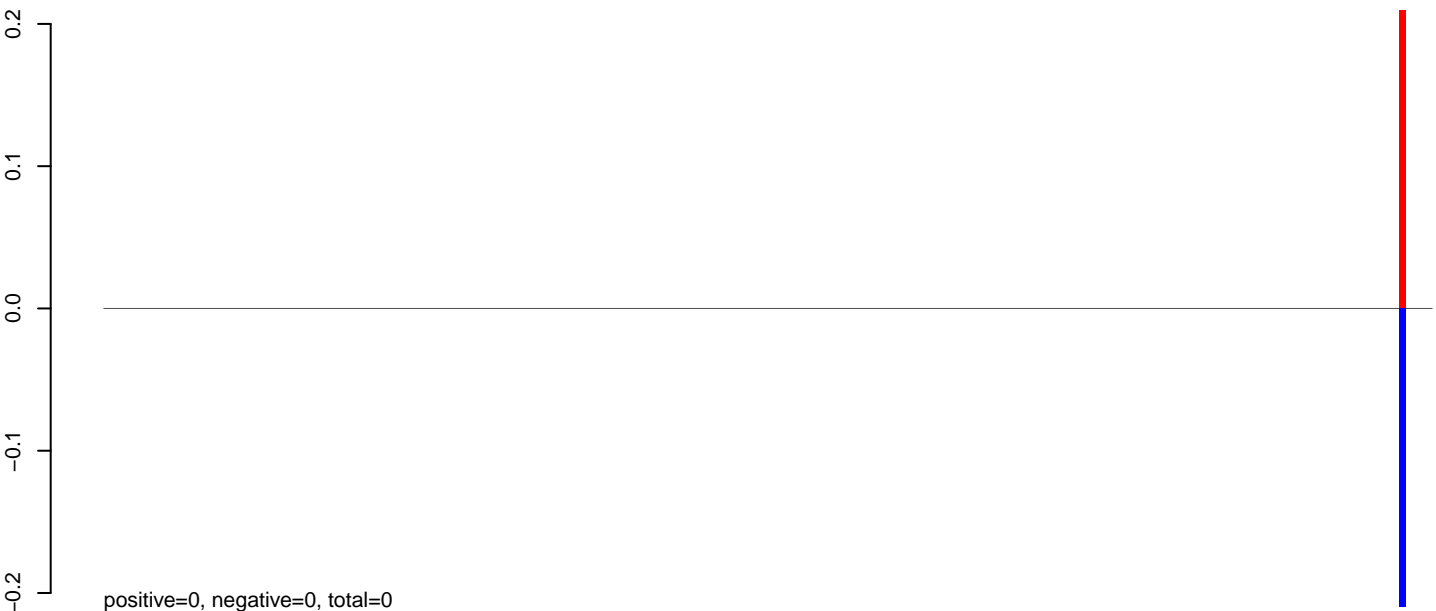
AeAeg_Aag2_DENV4.24_35.rep



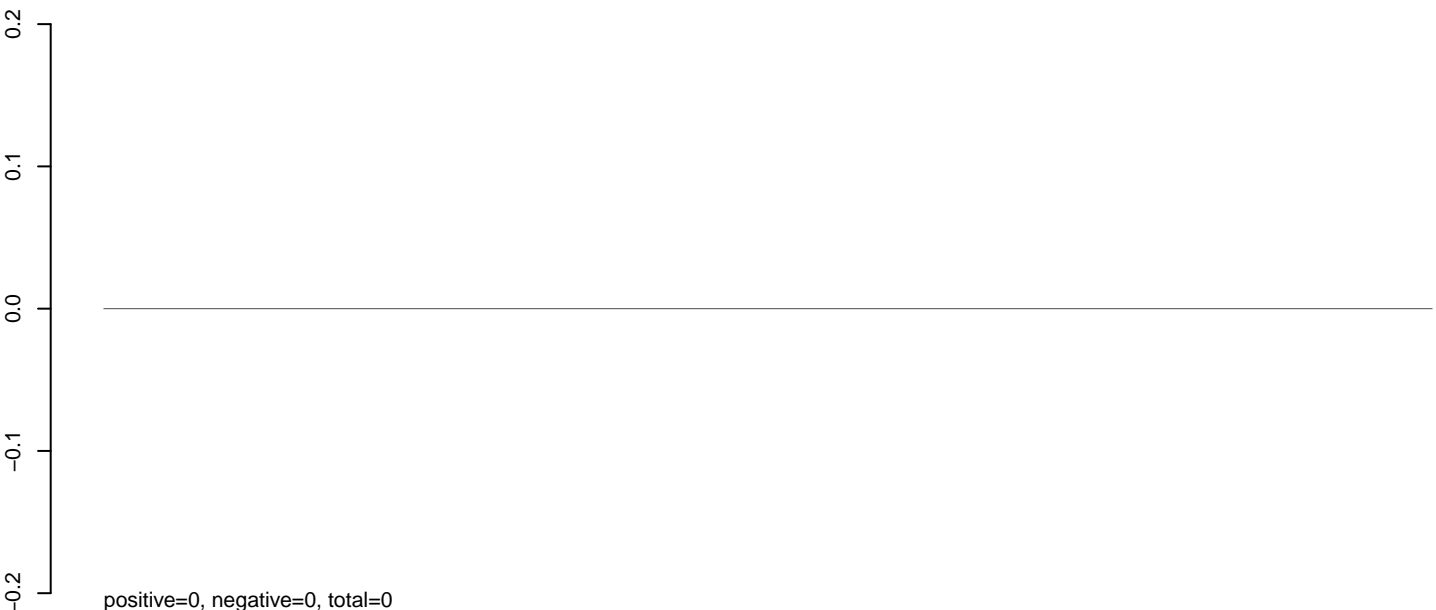
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

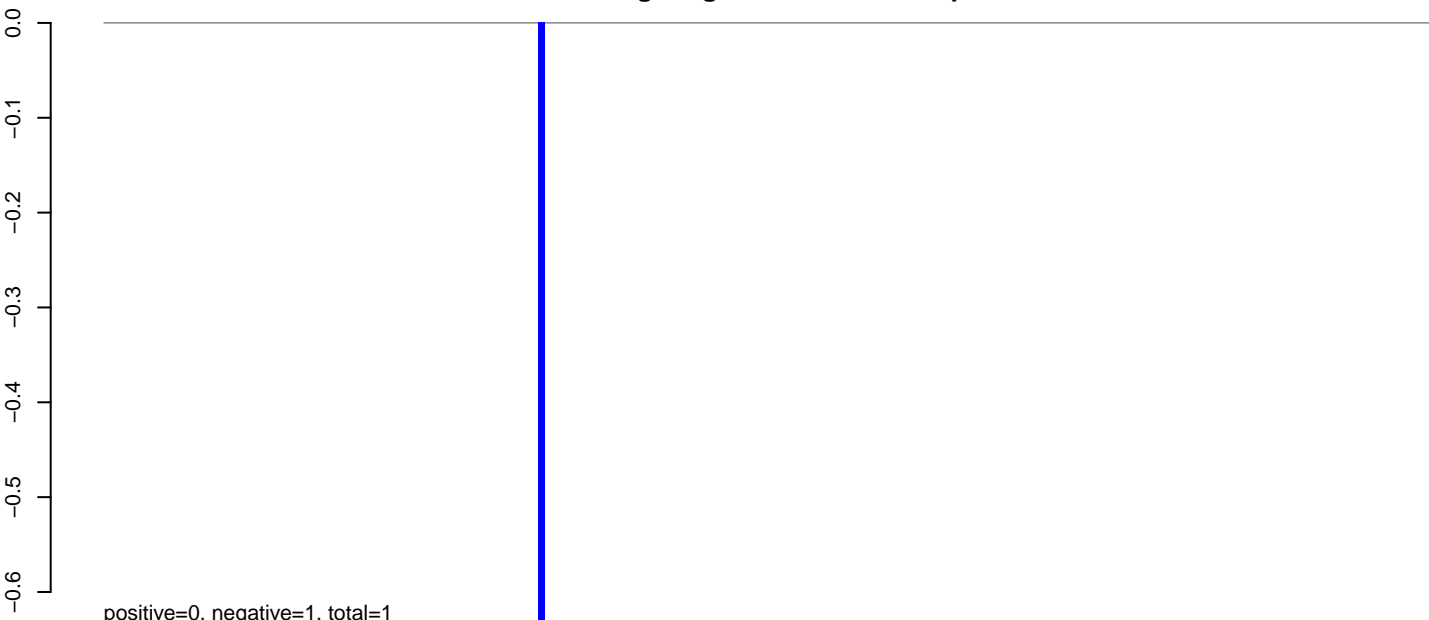


AeAeg_Aag2_DENV4.rep

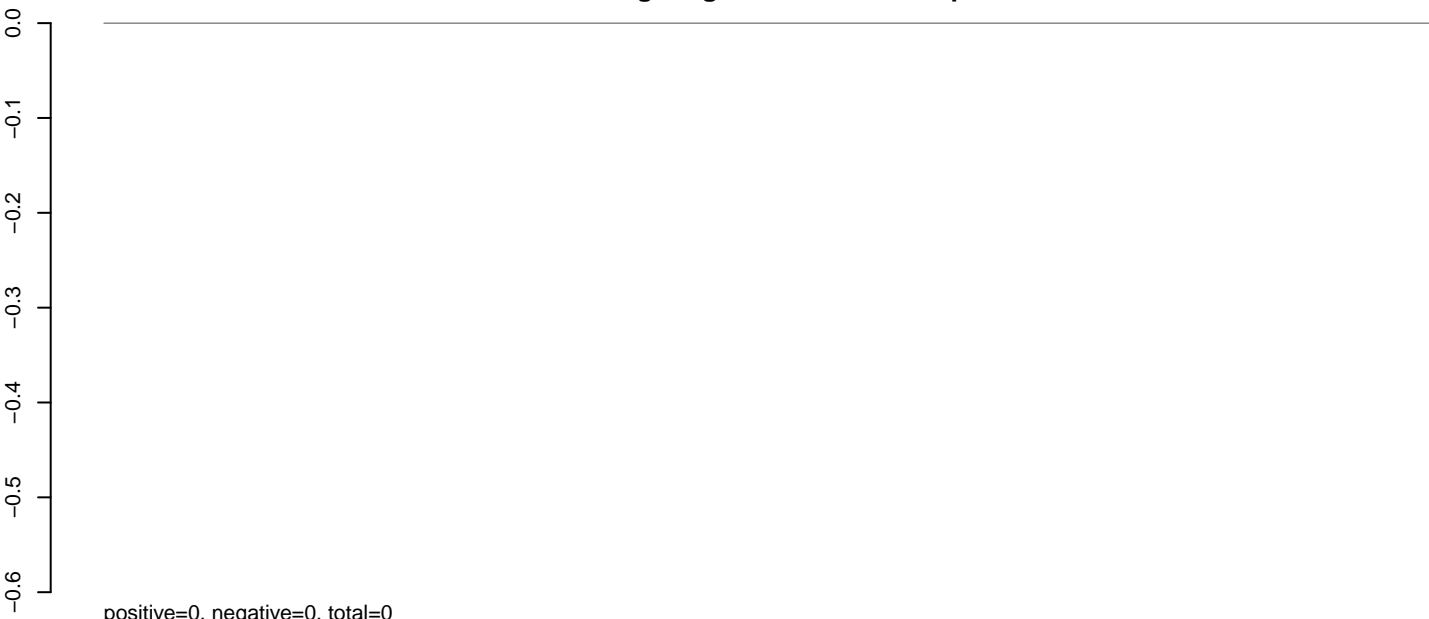


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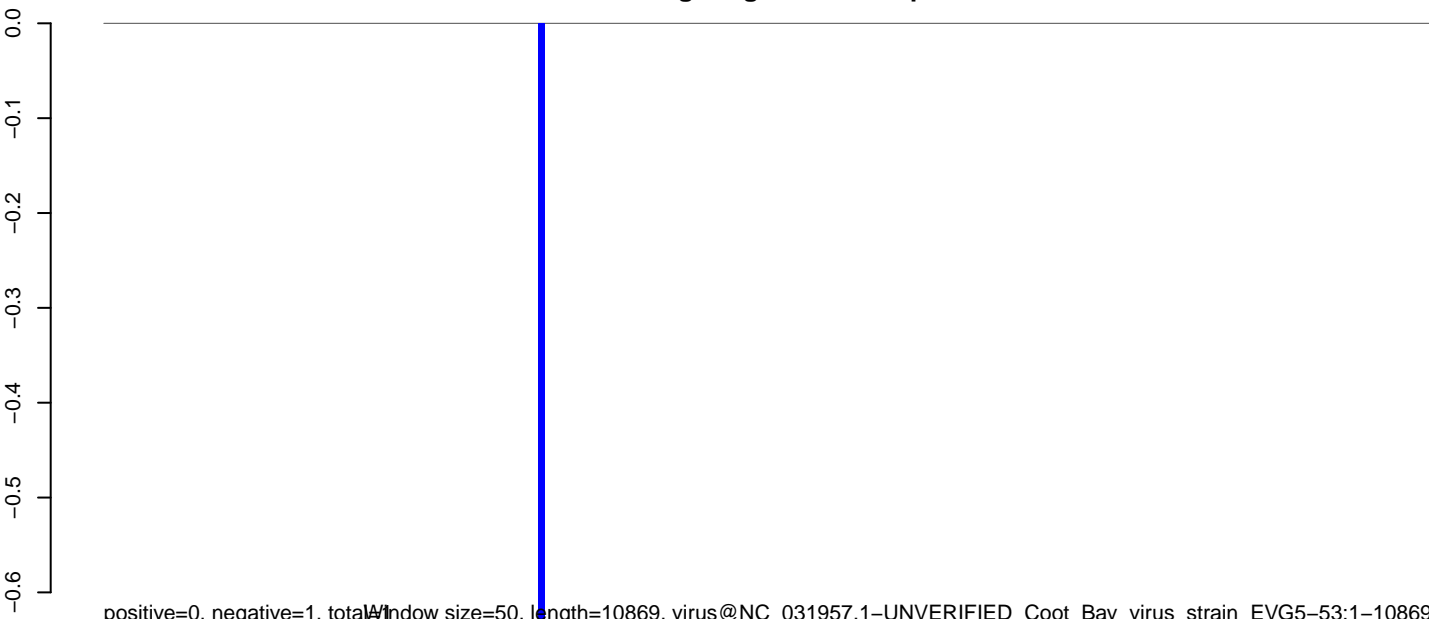
AeAeg_Aag2_DENV4.18_23.rep



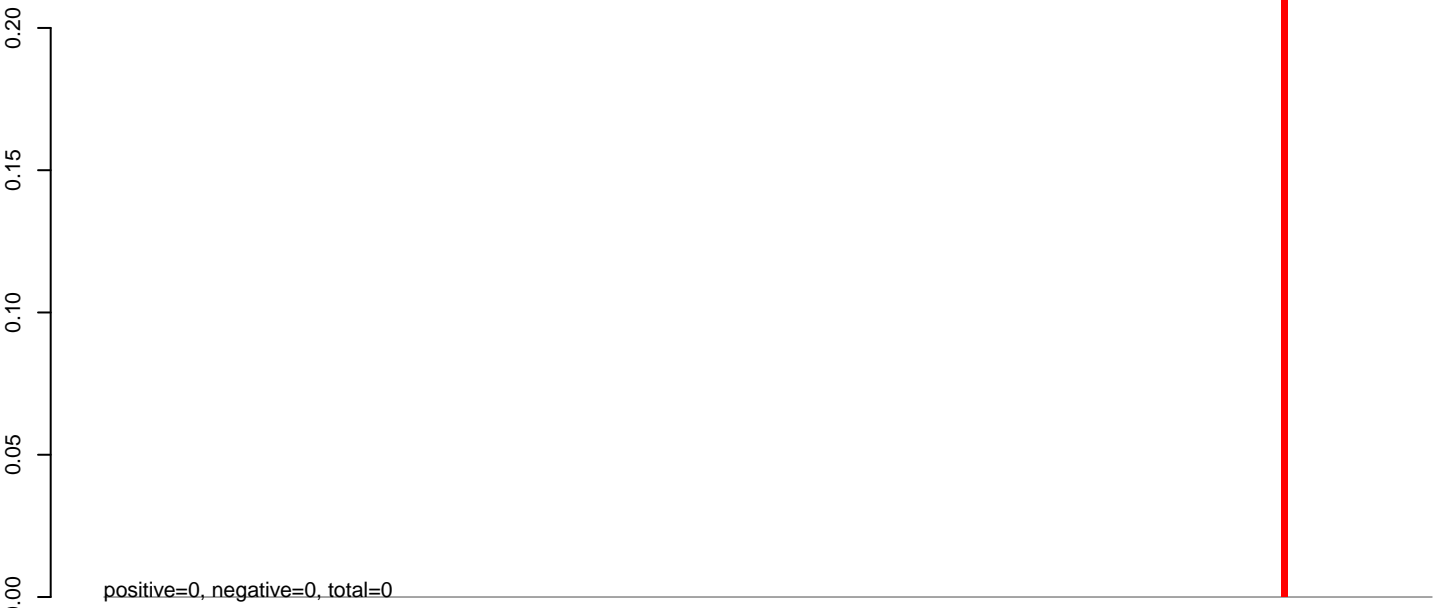
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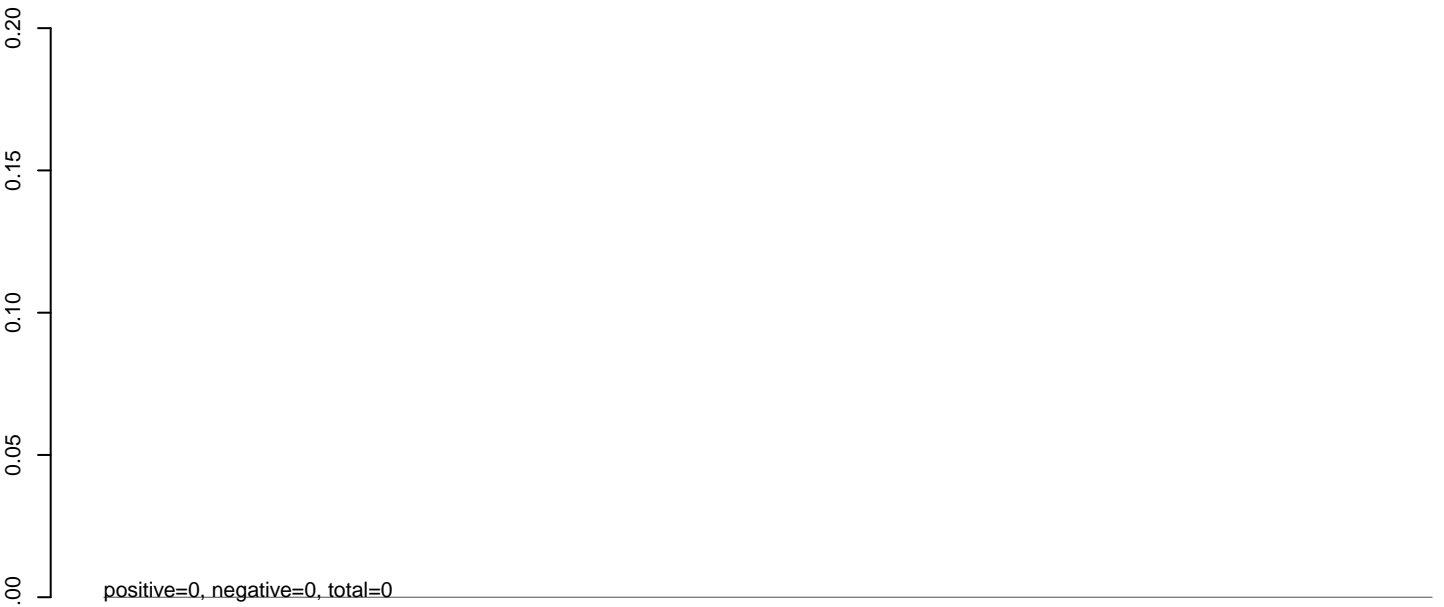
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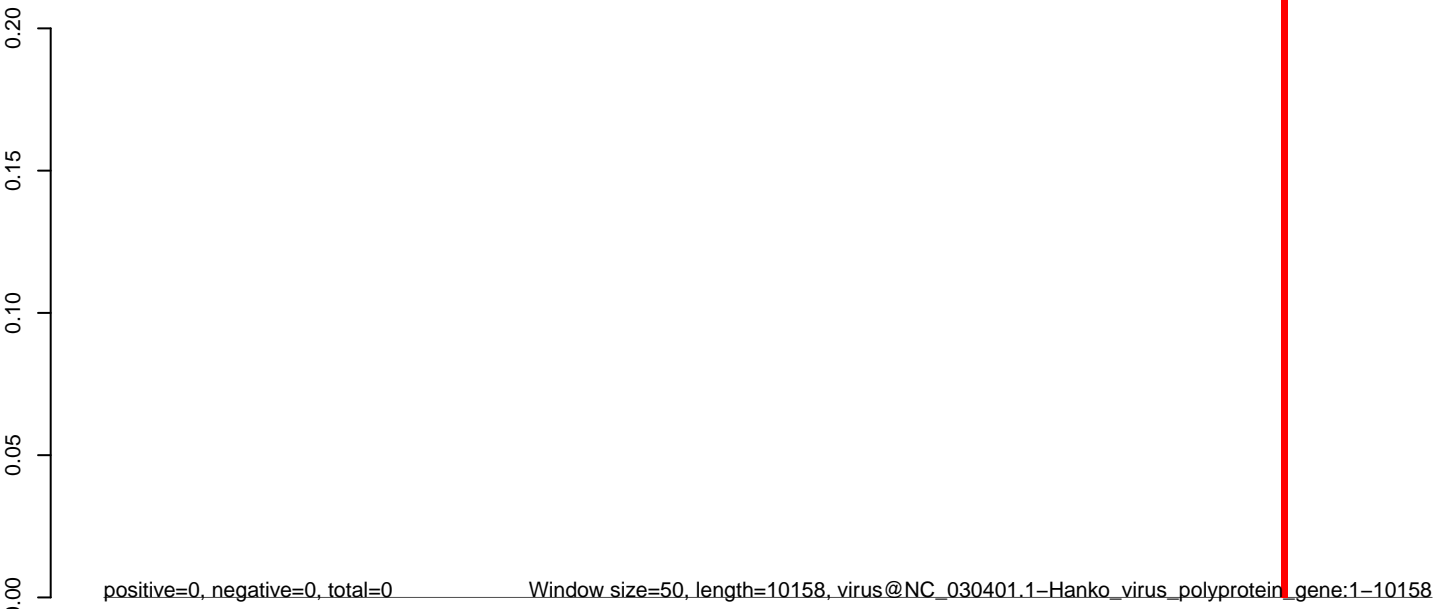
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

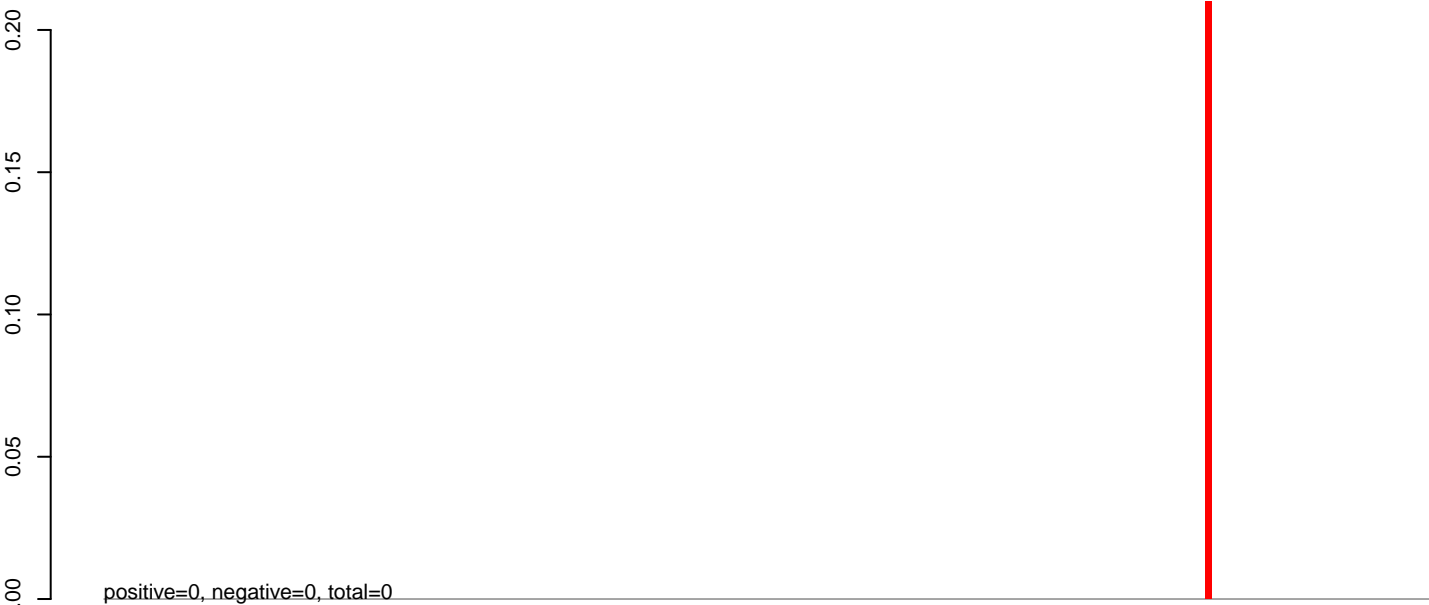


AeAeg_Aag2_DENV4.rep



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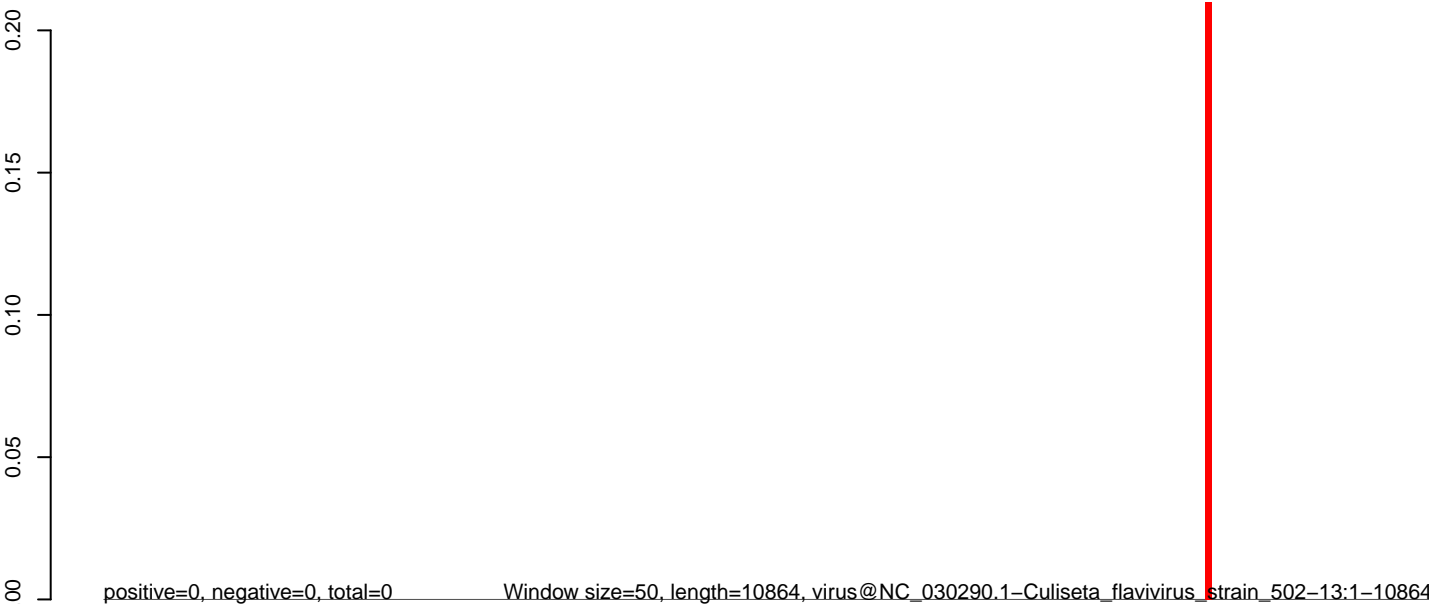
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

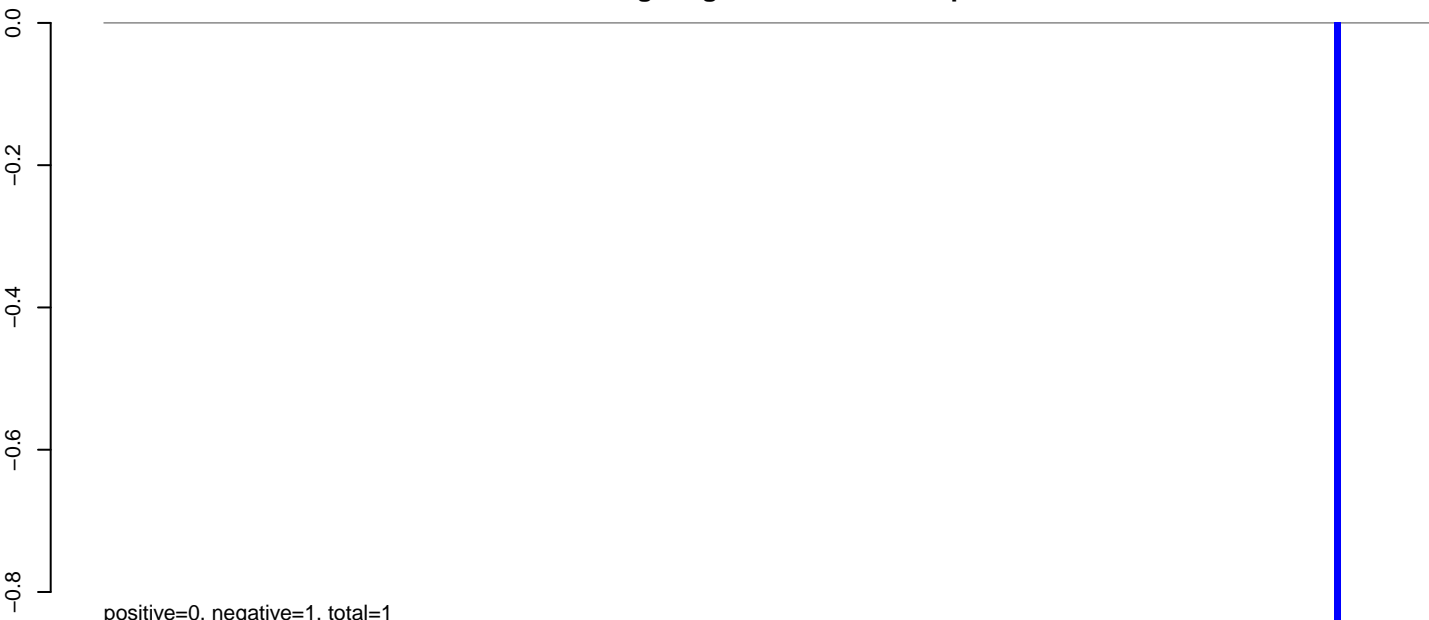


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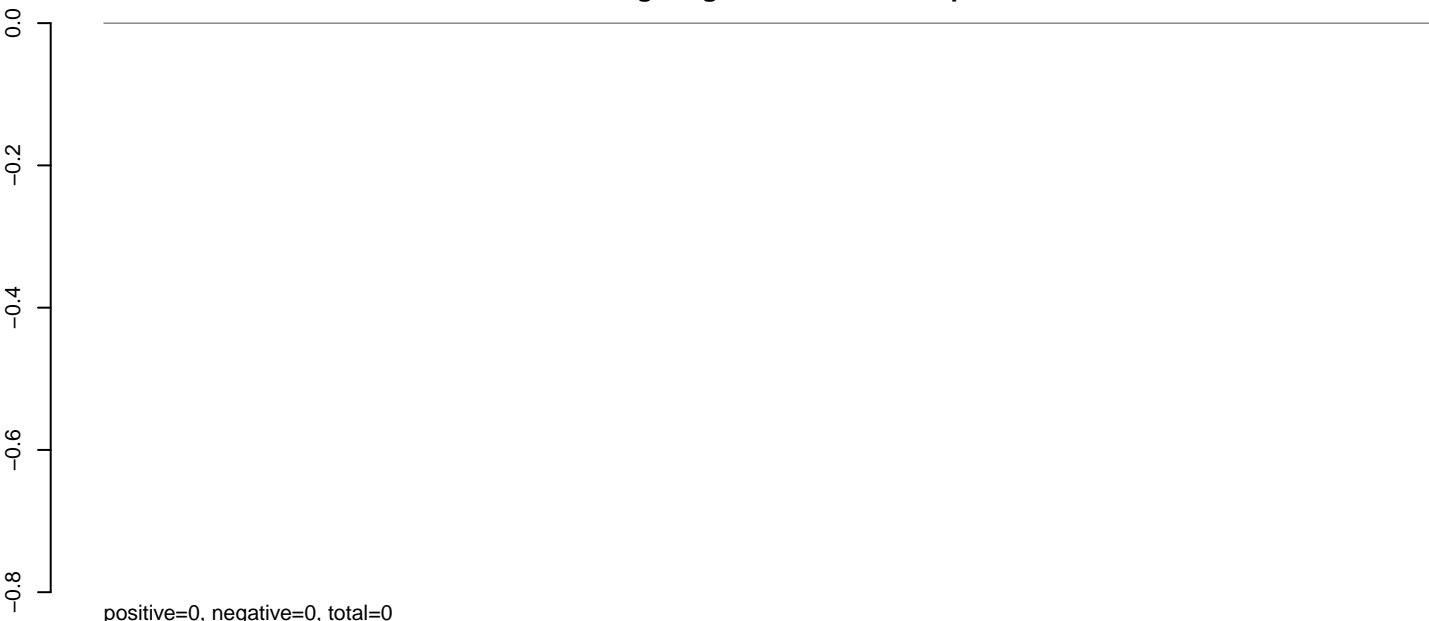


0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



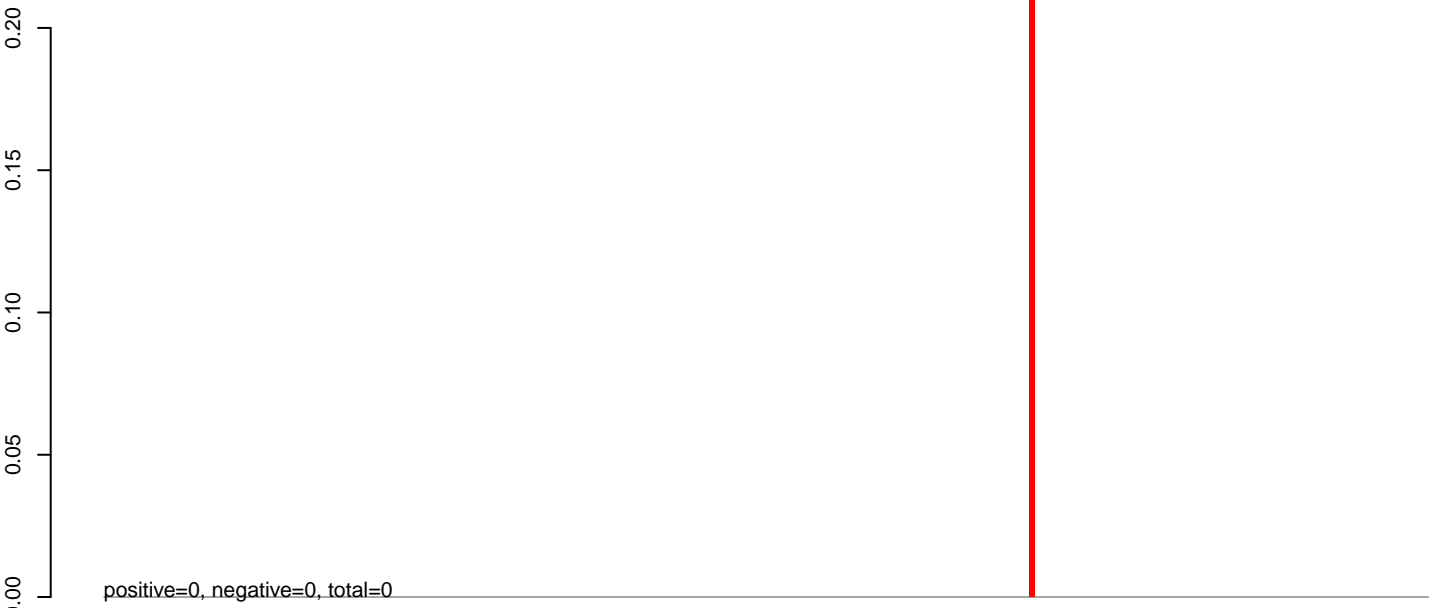
AeAeg_Aag2_DENV4.rep



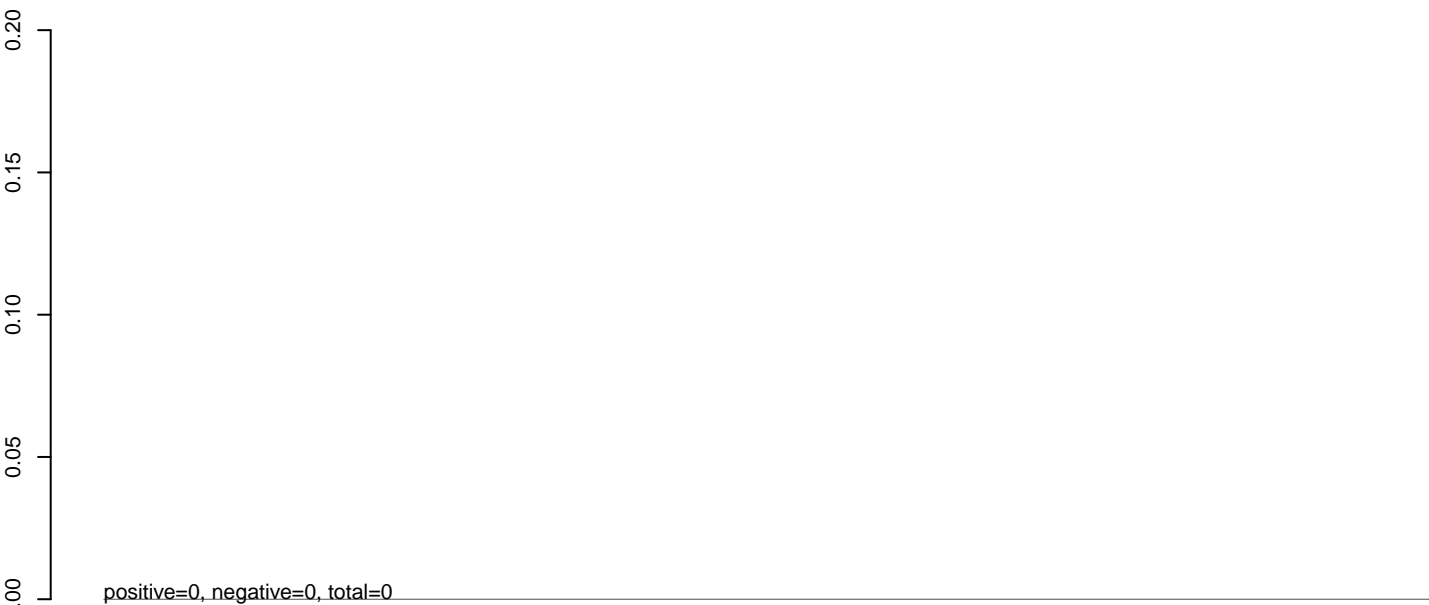
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0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV4.18_23.rep



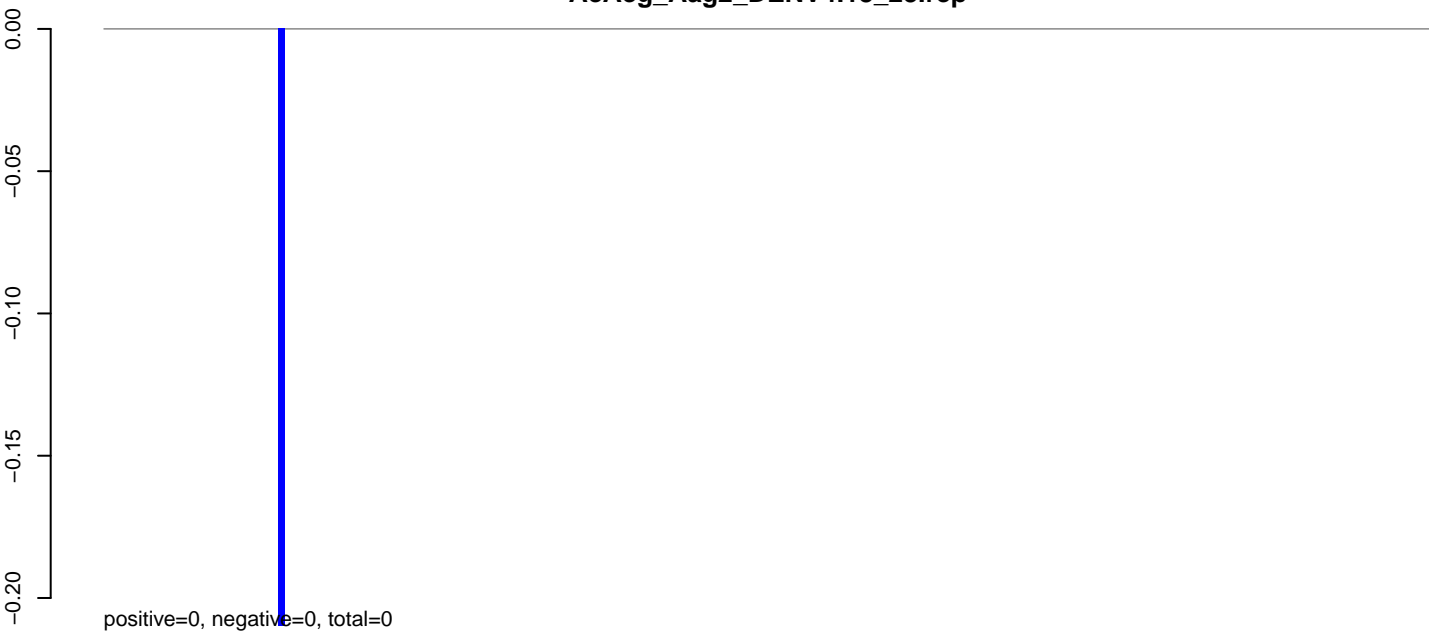
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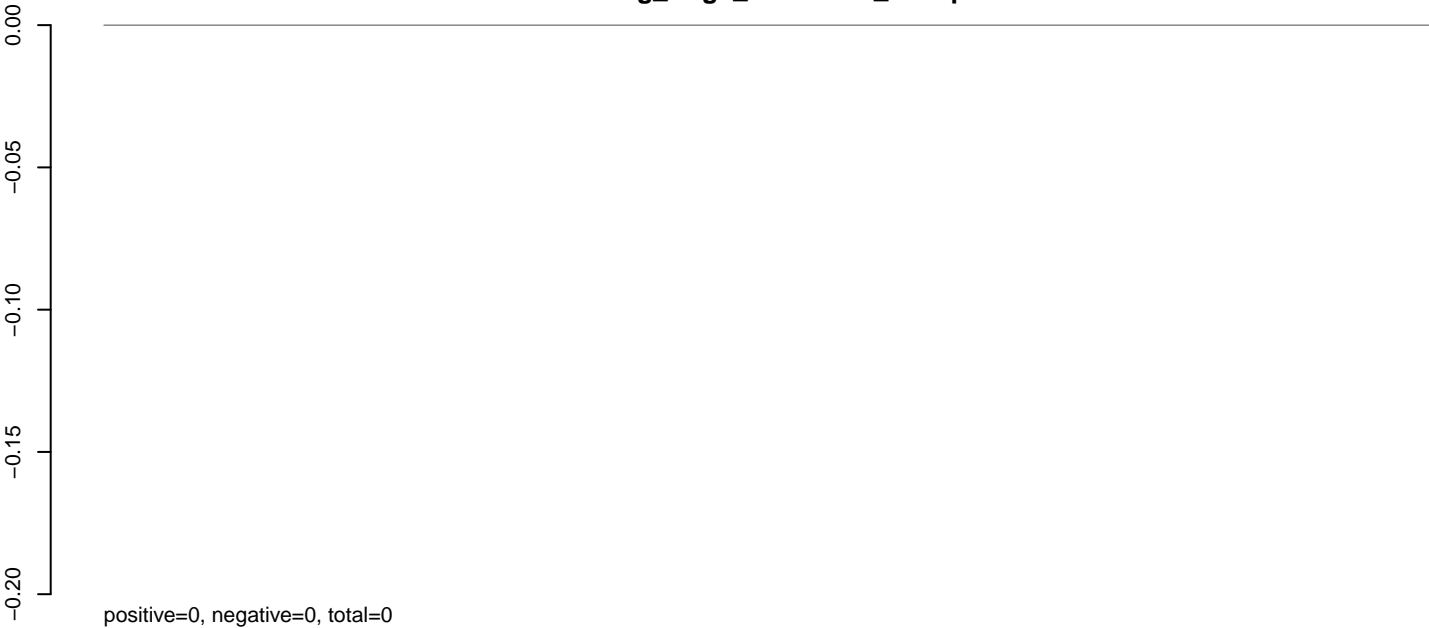
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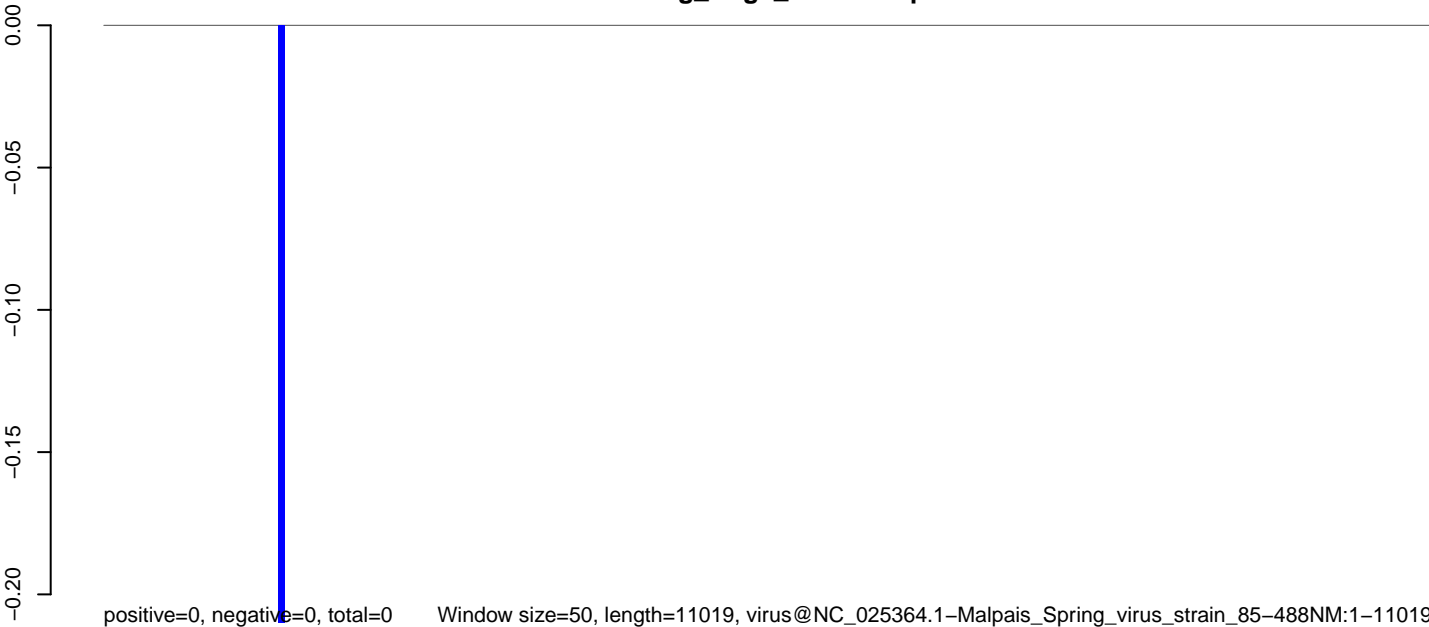
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



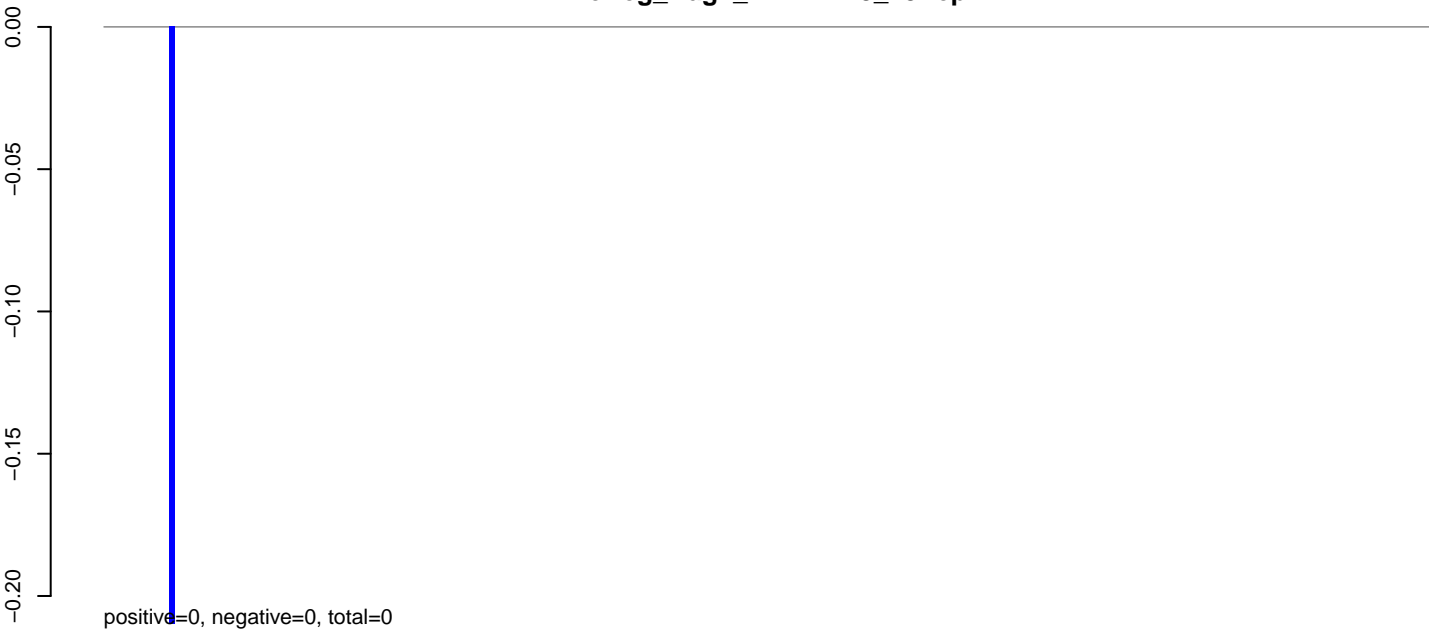
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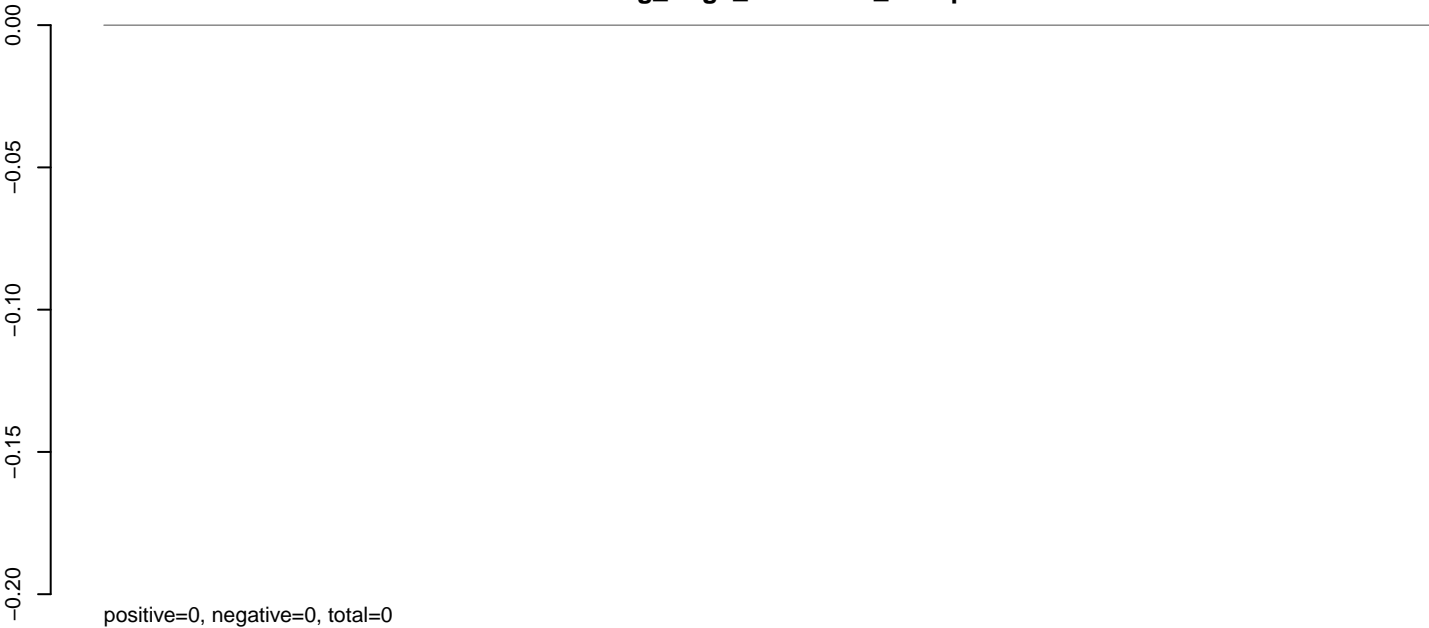
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0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV4.18_23.rep



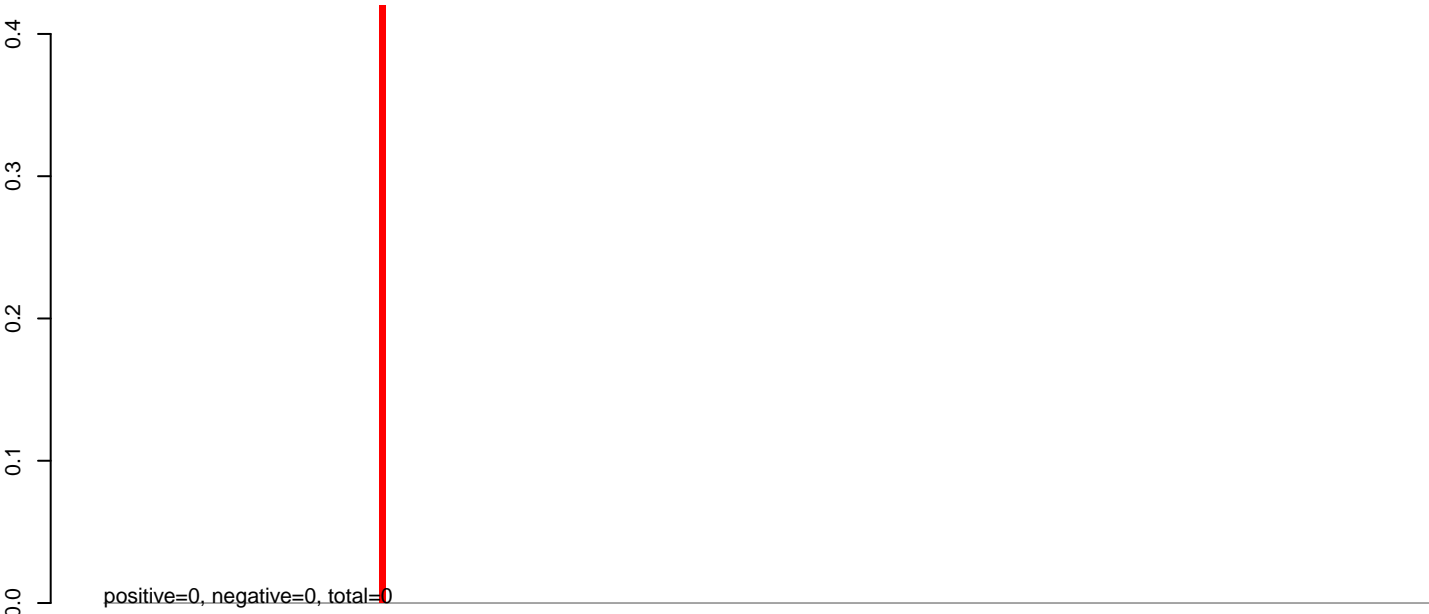
AeAeg_Aag2_DENV4.24_35.rep



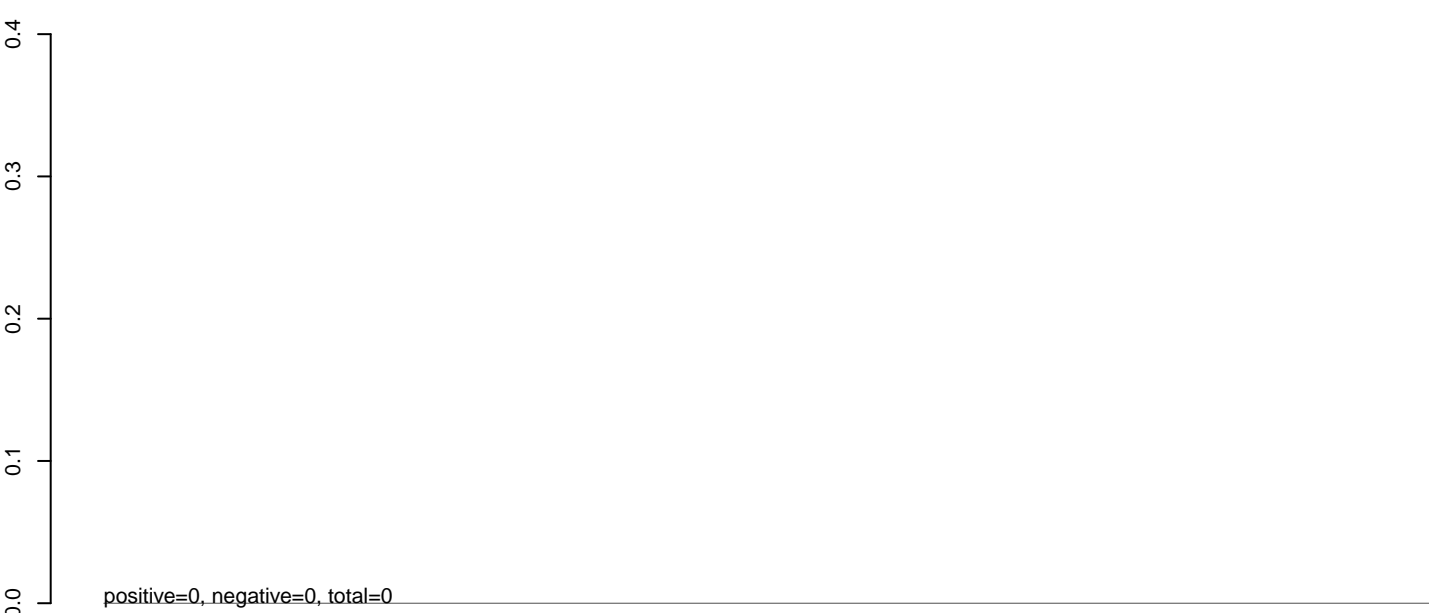
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

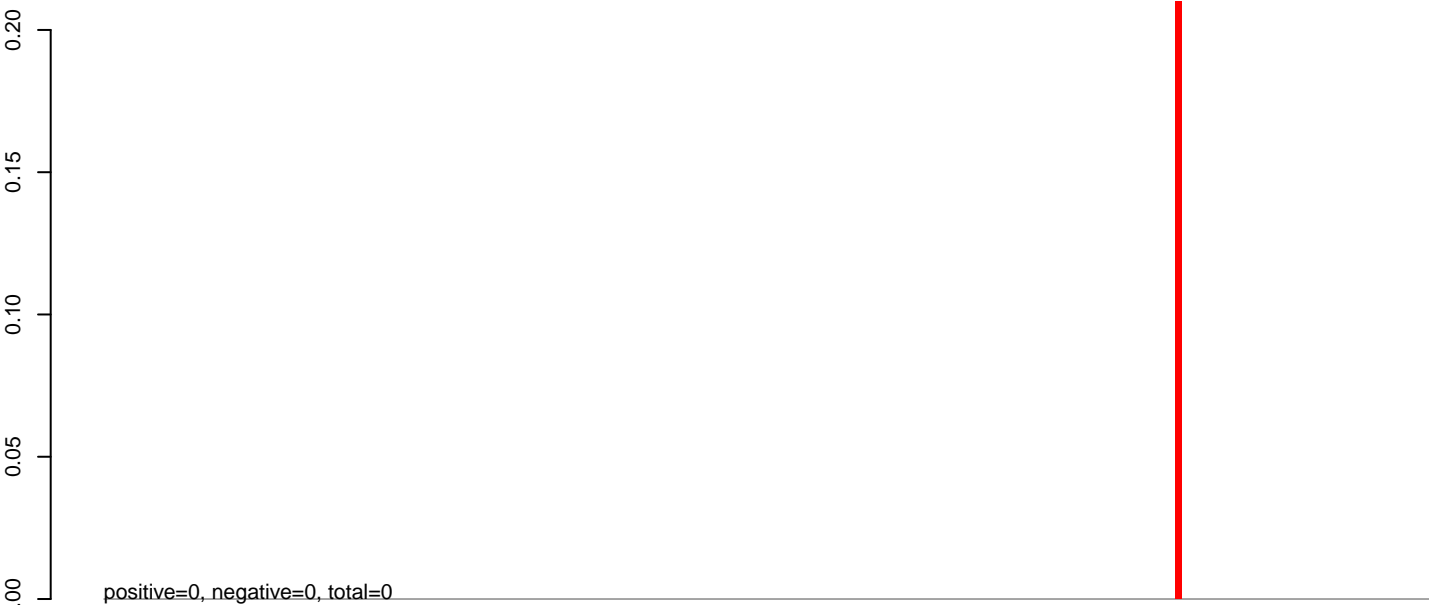


AeAeg_Aag2_DENV4.rep

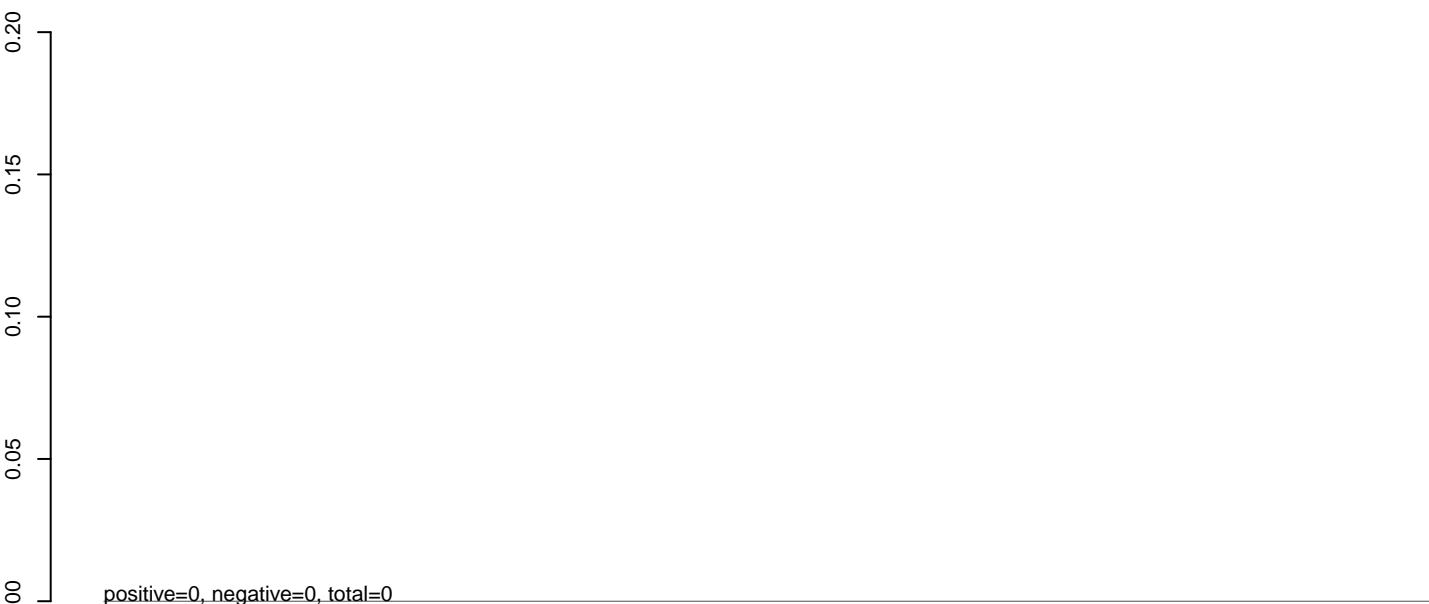


0 2000 4000 6000 8000 10000
window size=50, length=10365, virus@NC_024805.1-Ilomantsi_virus_isolate_M0724_polyprotein_gene:1-10365

AeAeg_Aag2_DENV4.18_23.rep



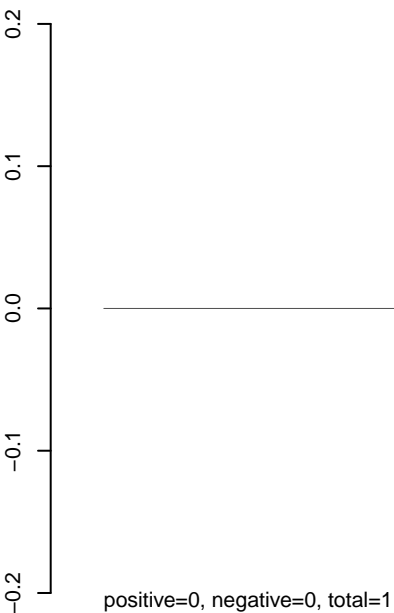
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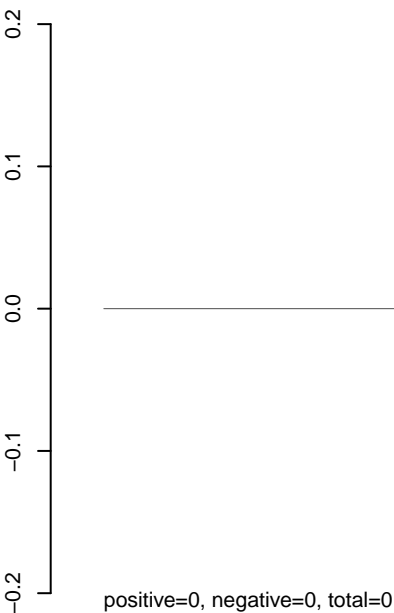
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

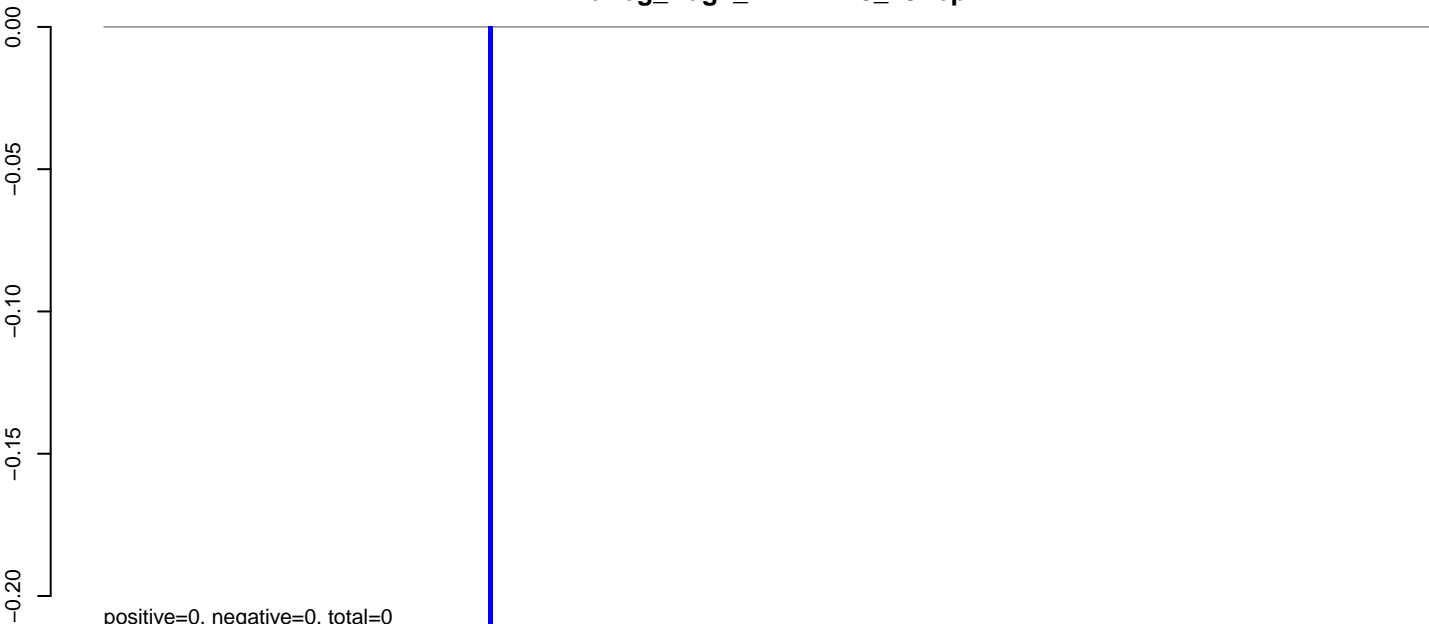


AeAeg_Aag2_DENV4.rep

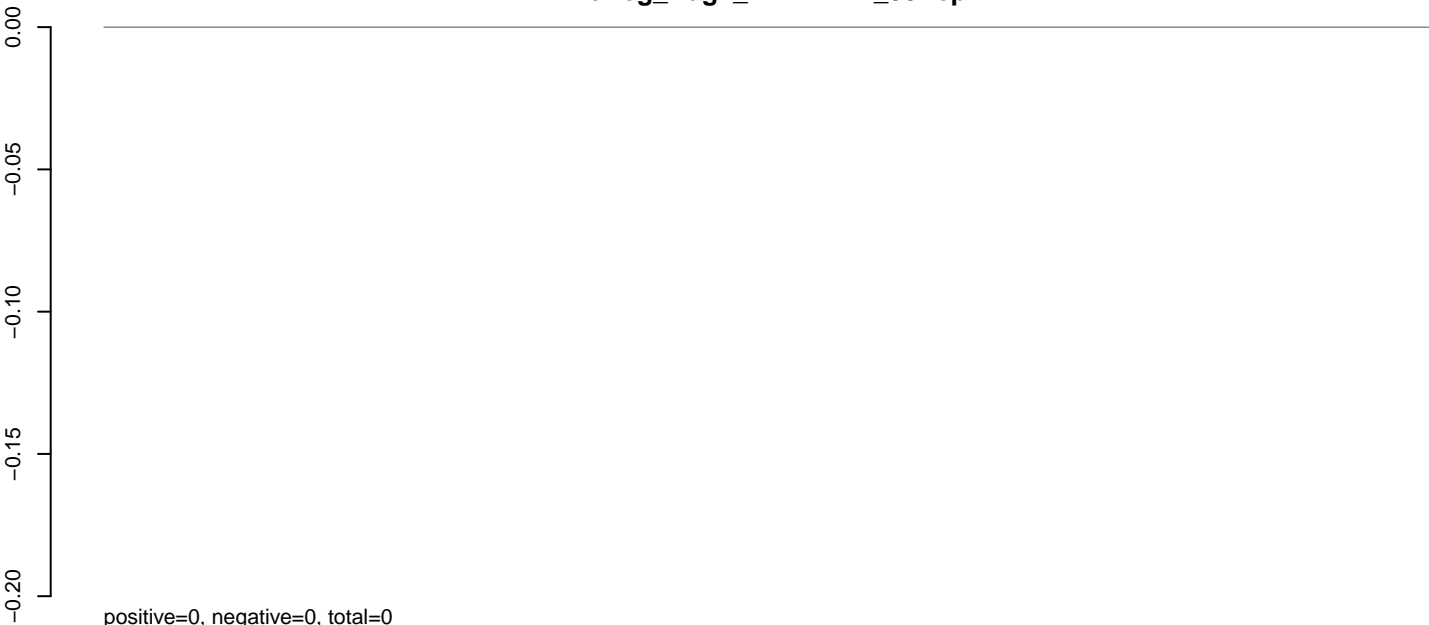


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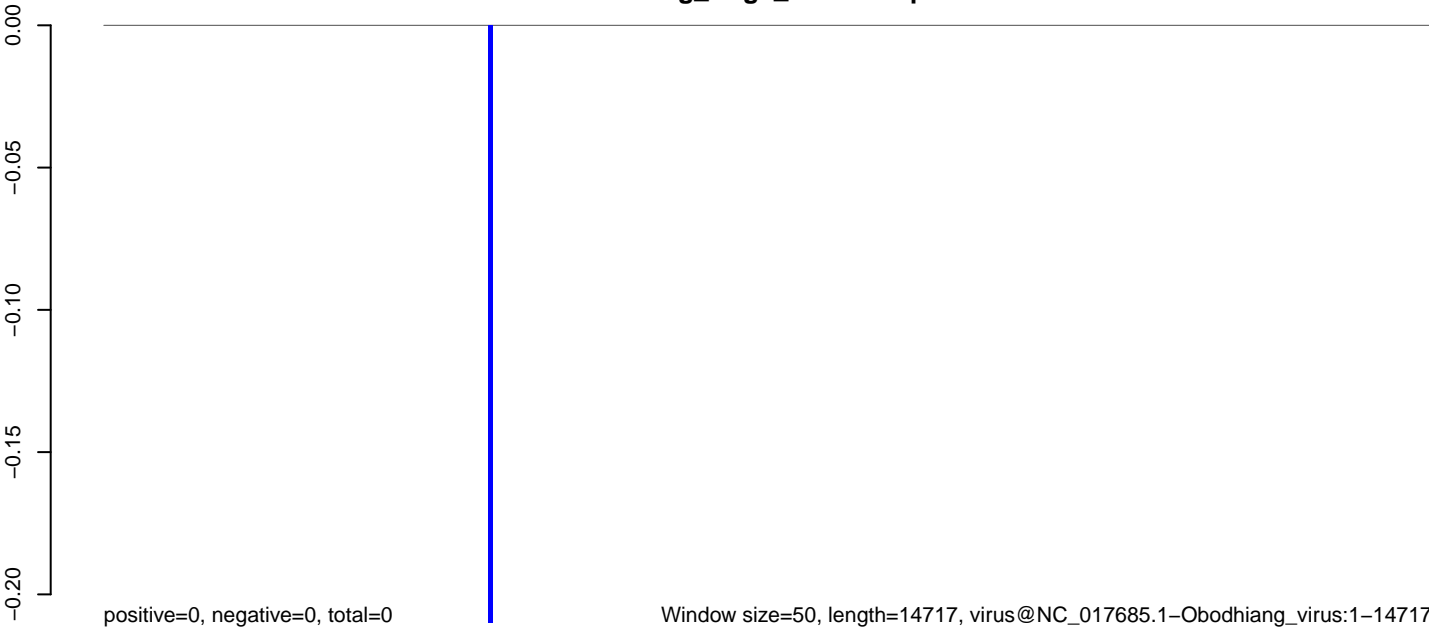
AeAeg_Aag2_DENV4.18_23.rep



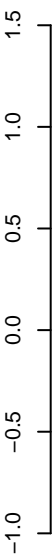
AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep

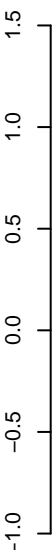


AeAeg_Aag2_DENV4.18_23.rep



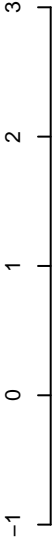
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AeAeg_Aag2_DENV4.24_35.rep



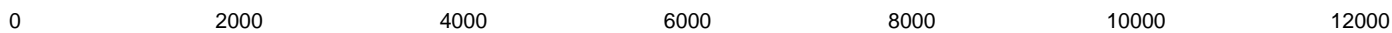
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AeAeg_Aag2_DENV4.rep

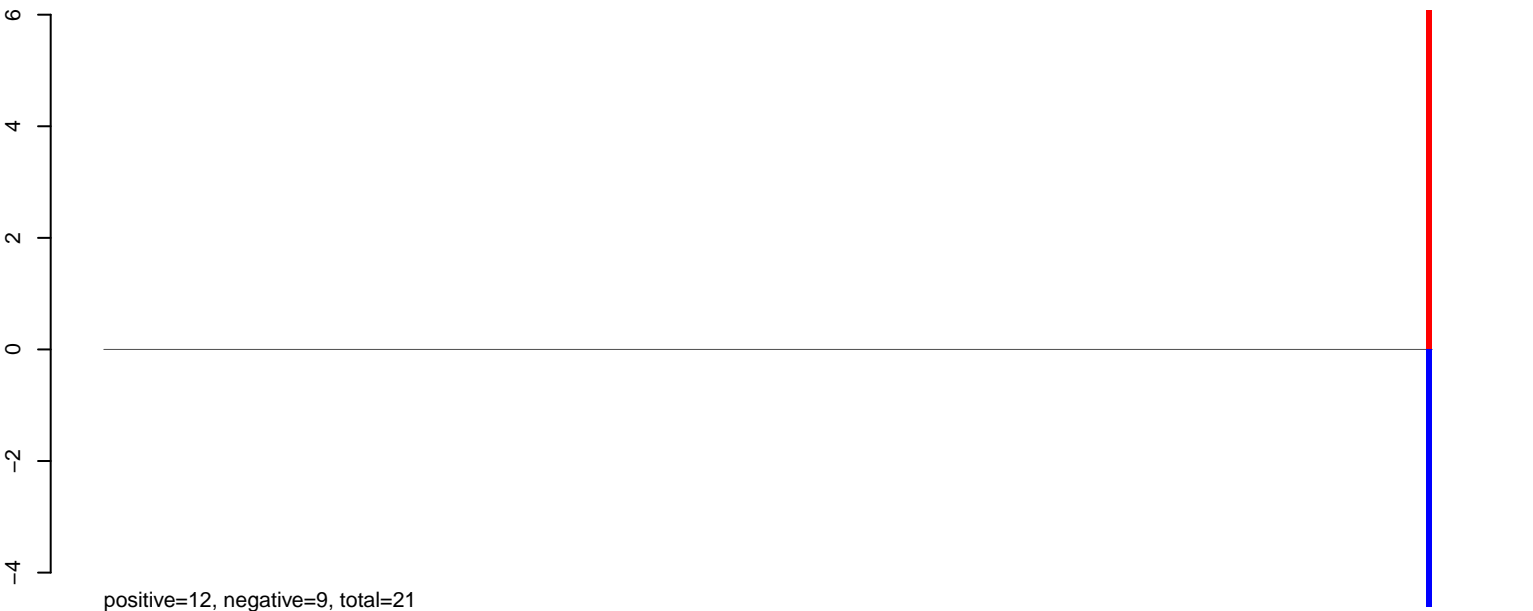


positive=6, negative=3, total=9

Window size=50, length=11877, virus@NC_016962.1-Bebaru_virus:1-11877



AeAeg_Aag2_DENV4.18_23.rep



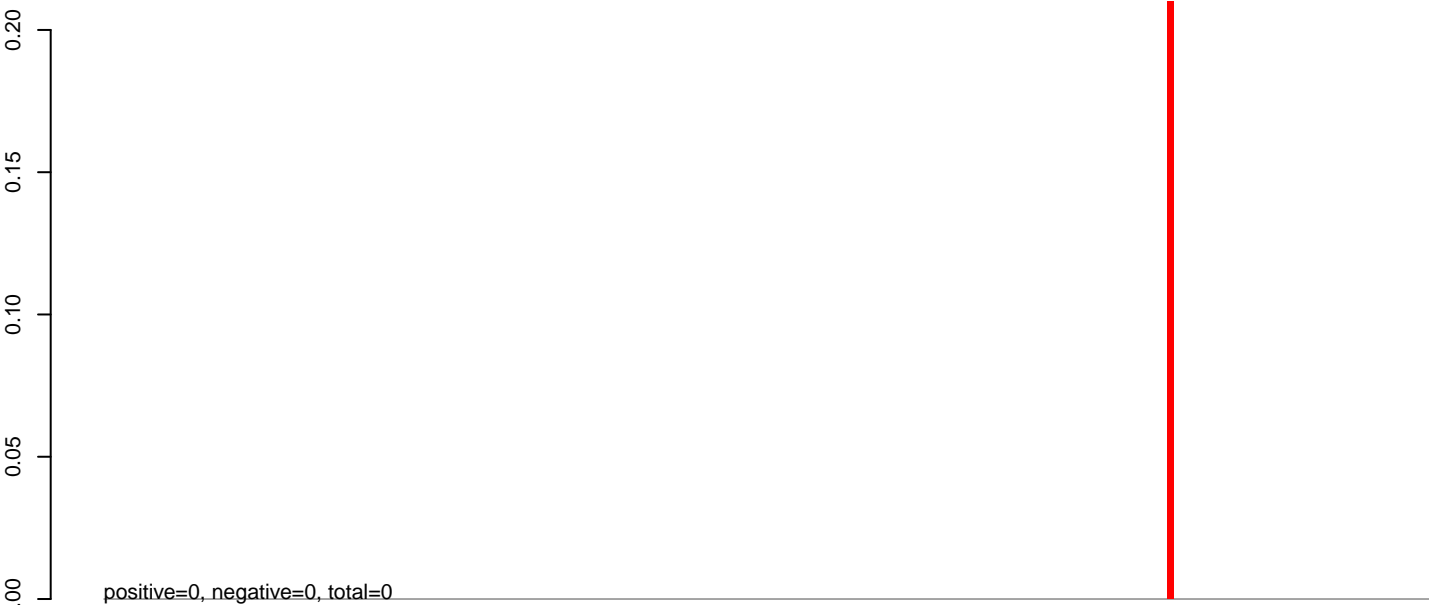
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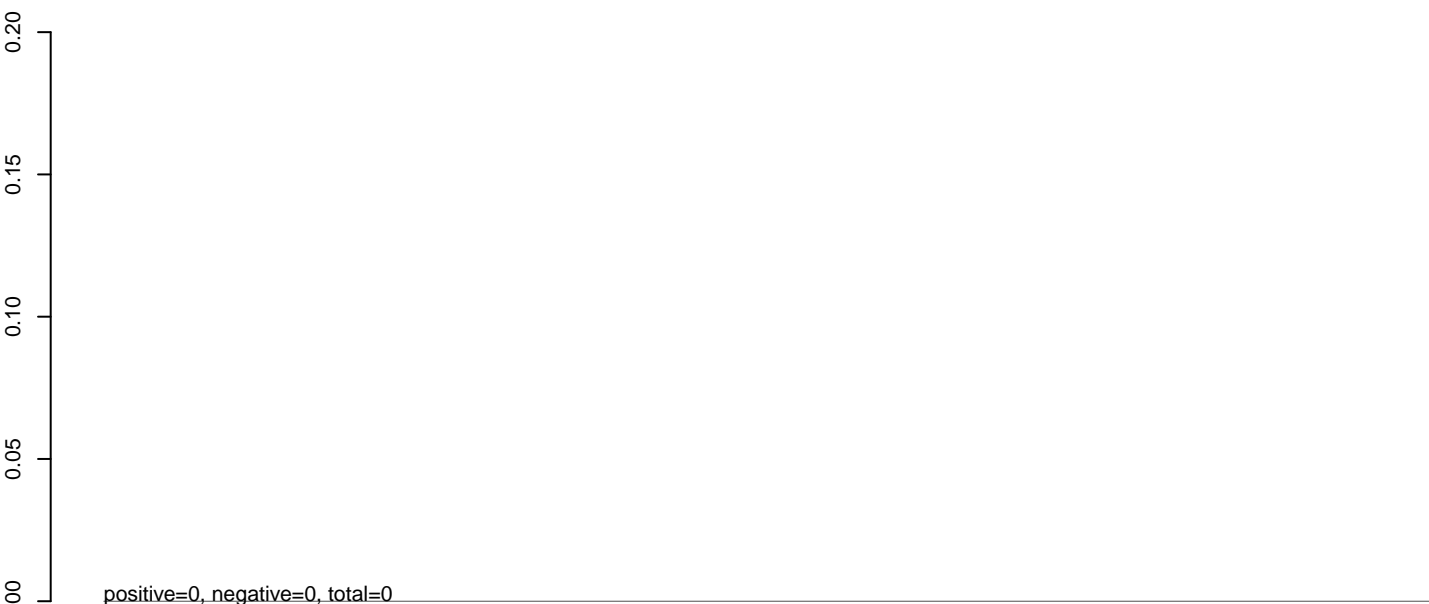
AeAeg_Aag2_DENV4.rep



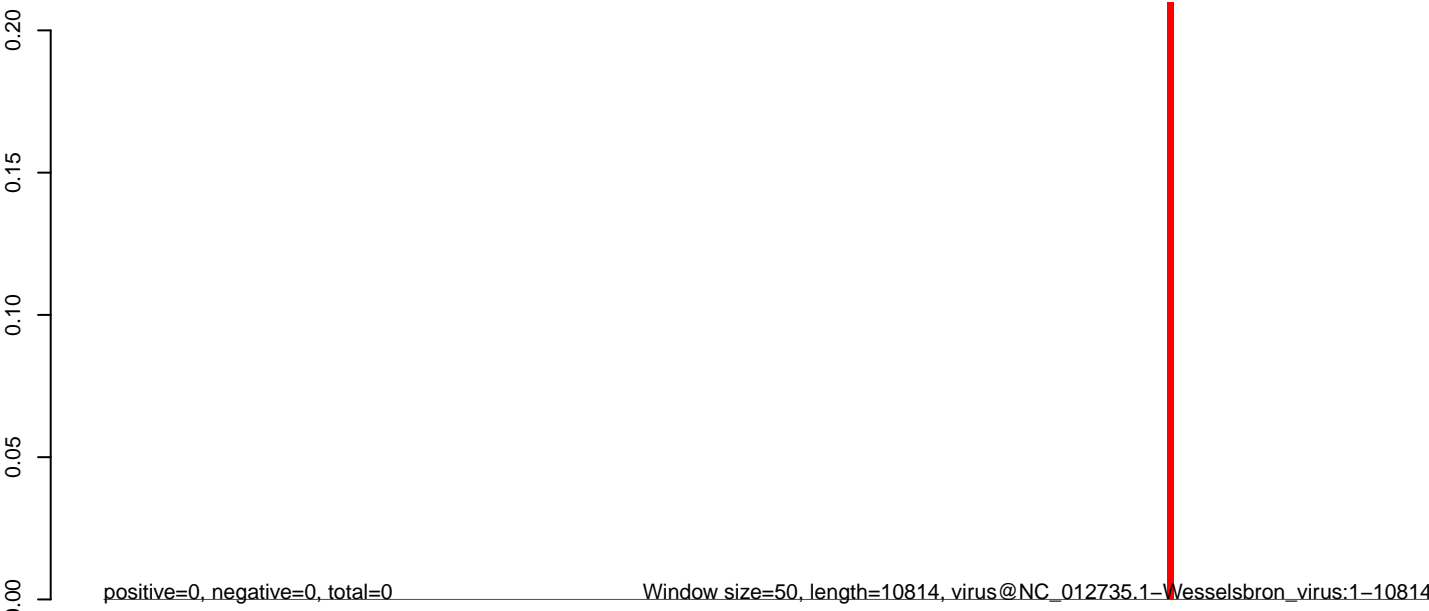
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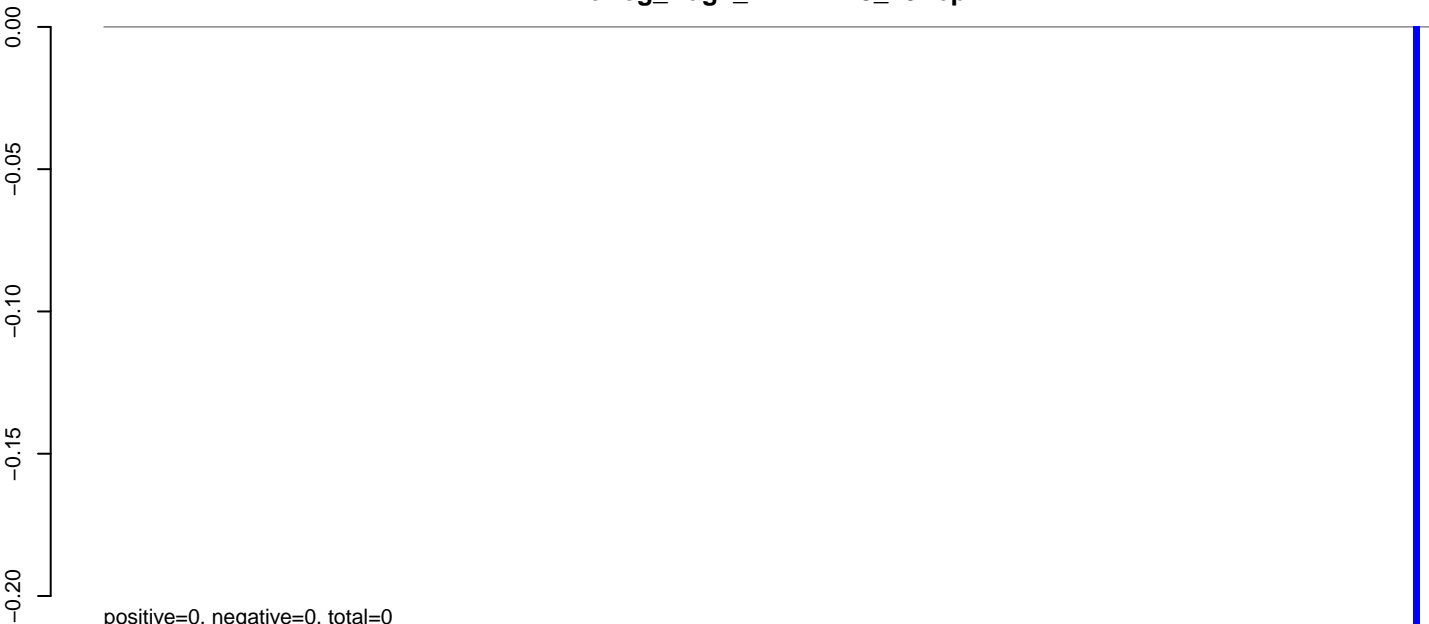
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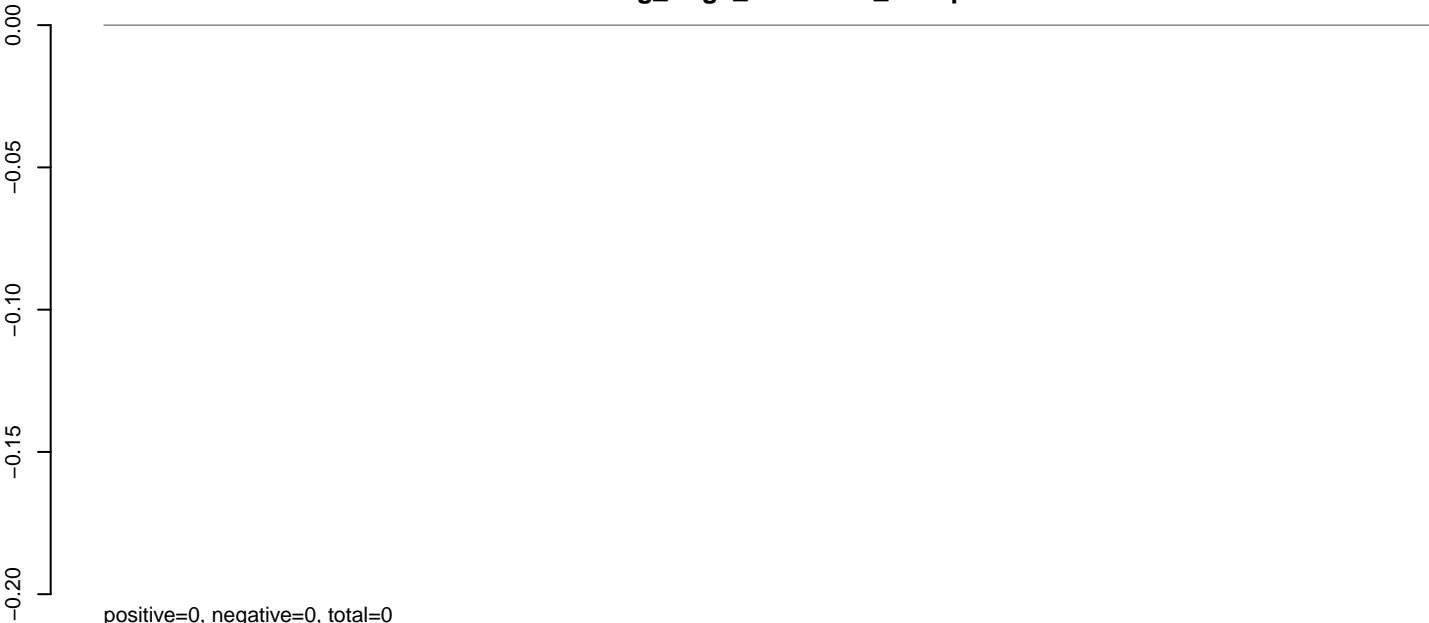
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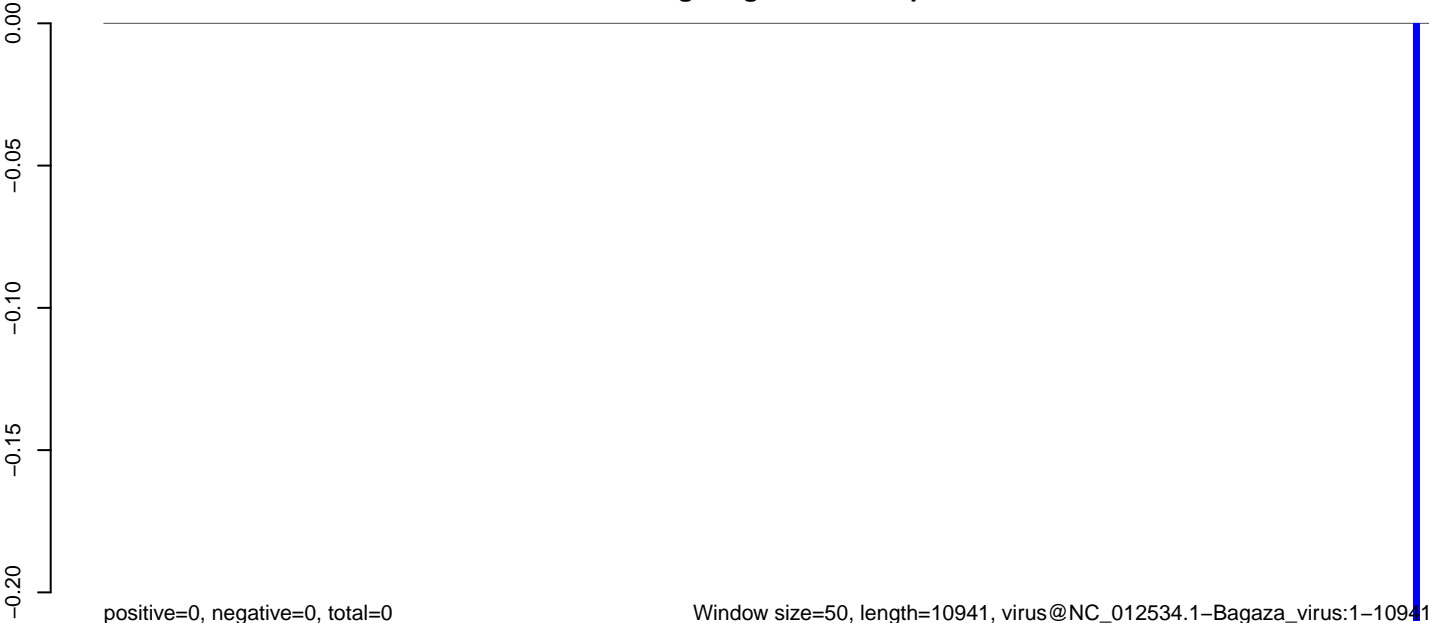
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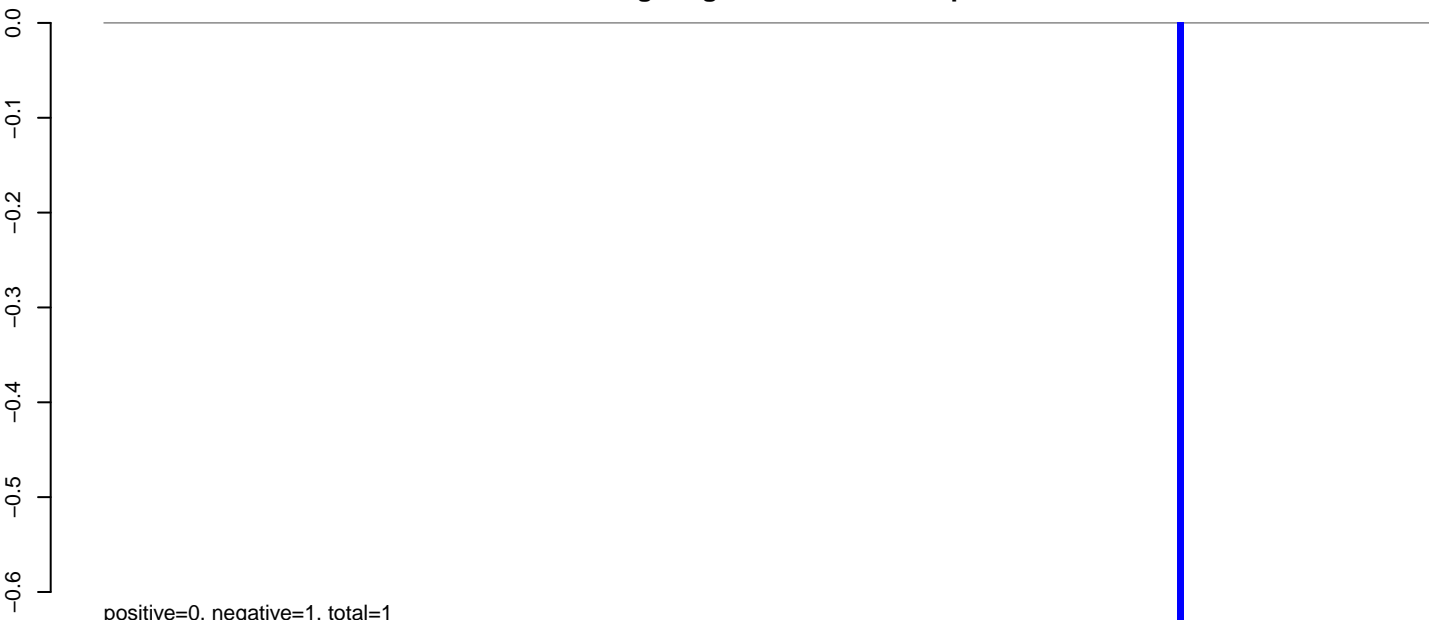
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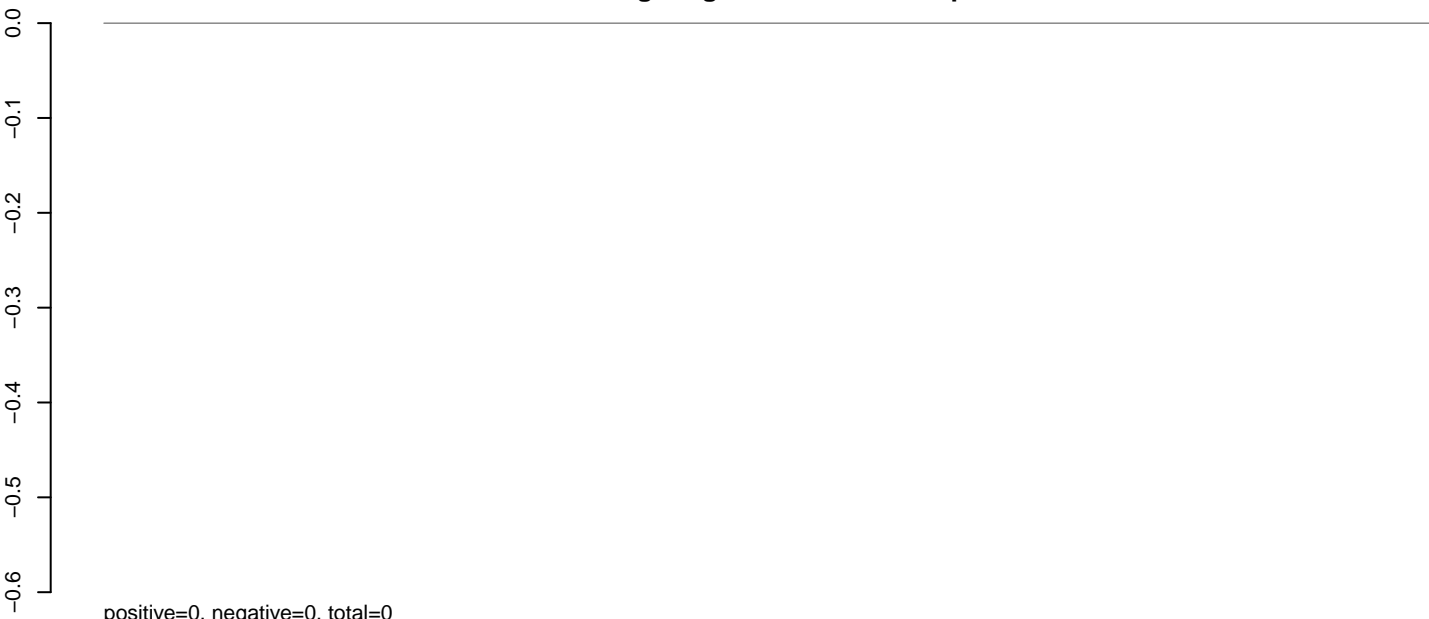
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



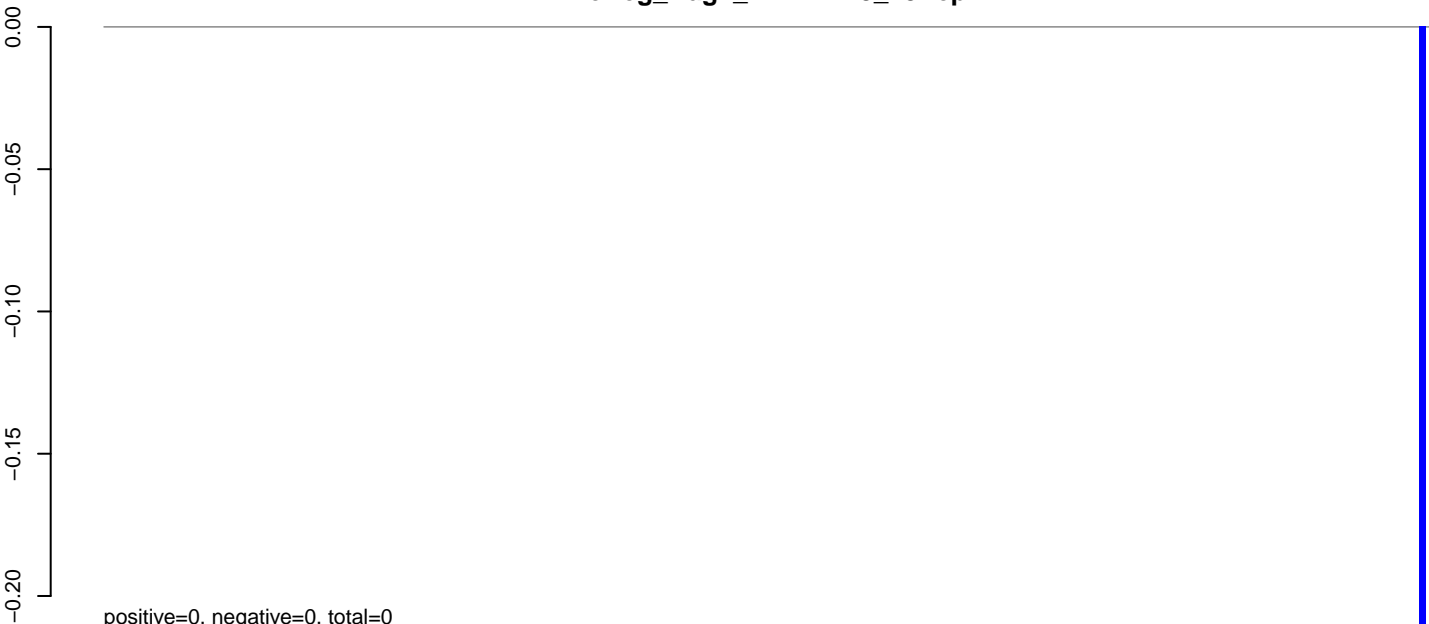
AeAeg_Aag2_DENV4.rep



Window size=50, length=10723, virus@NC_012533.1-Kedougou_virus:1-10723

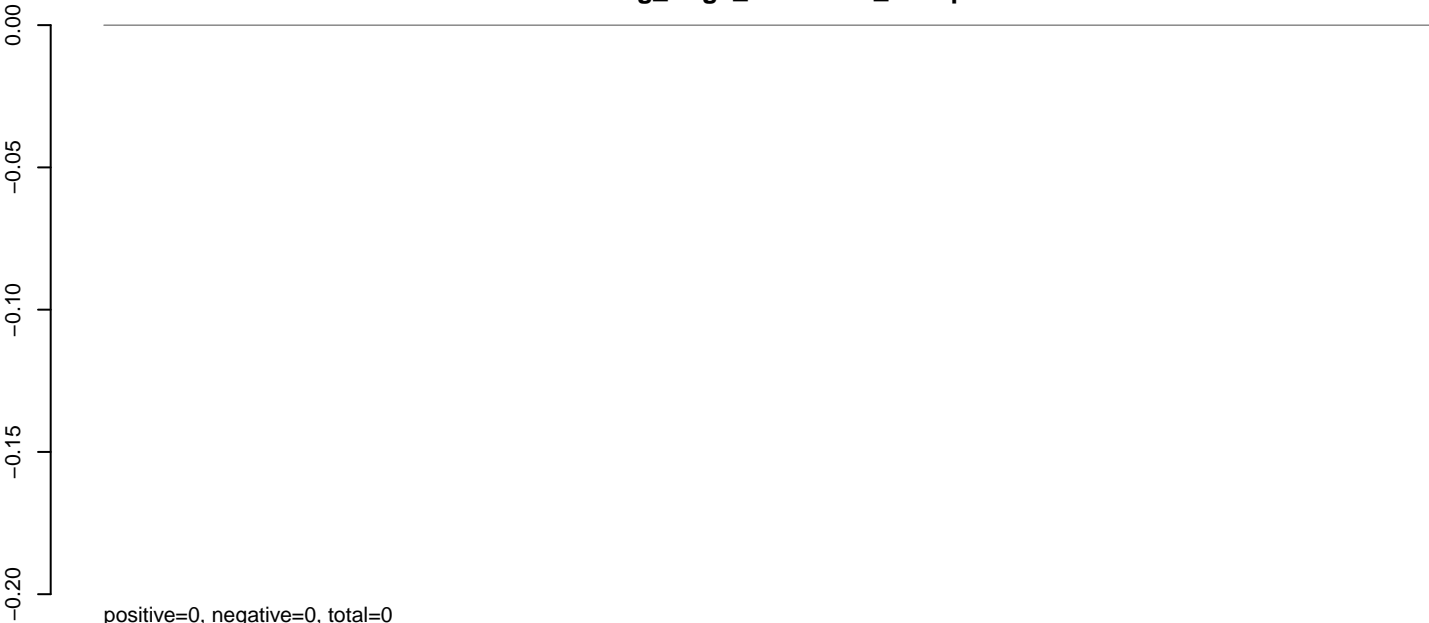
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AeAeg_Aag2_DENV4.18_23.rep



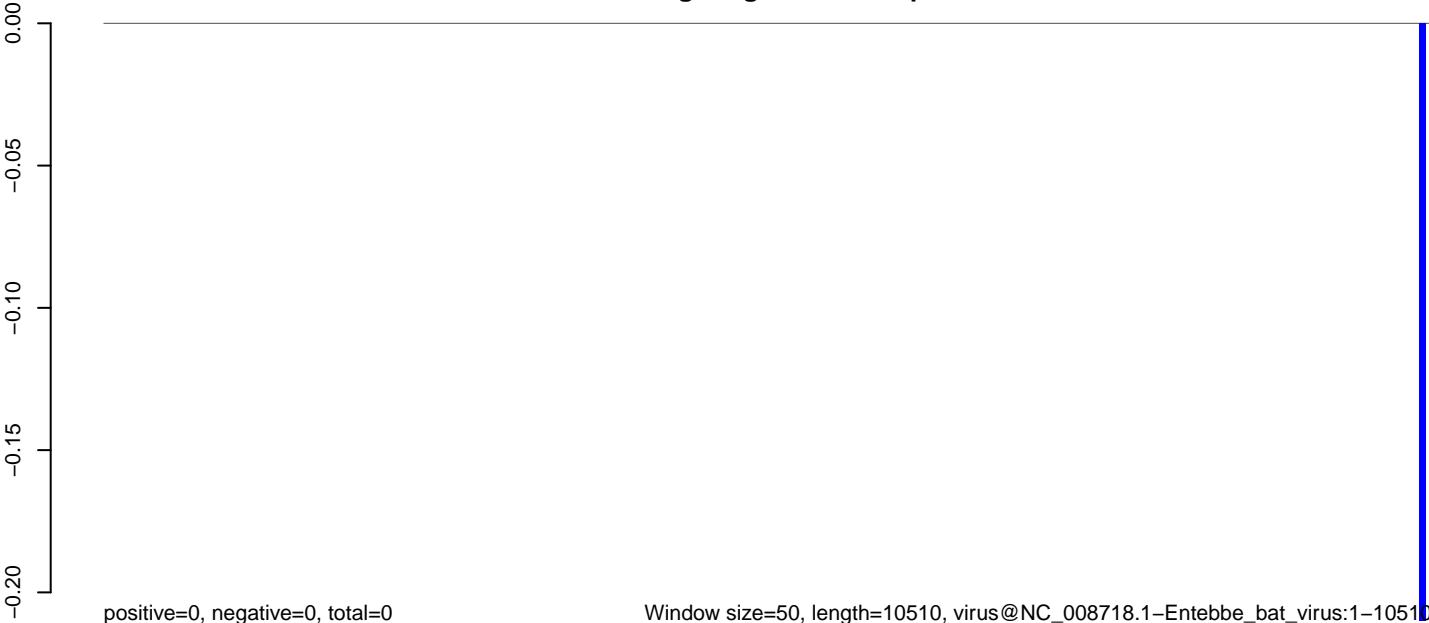
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AeAeg_Aag2_DENV4.24_35.rep



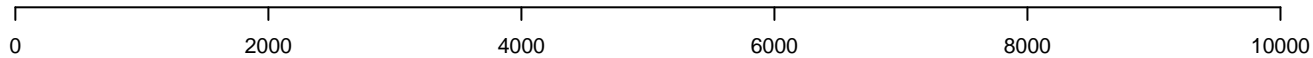
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AeAeg_Aag2_DENV4.rep

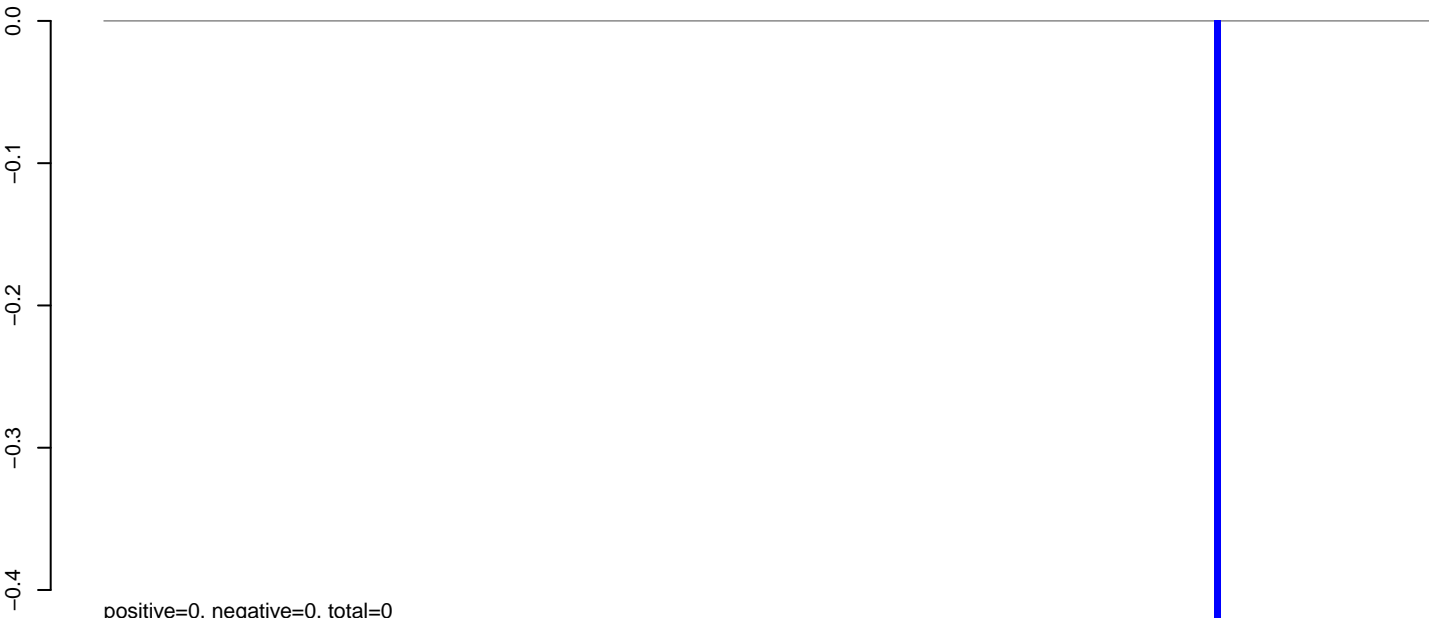


positive=0, negative=0, total=0

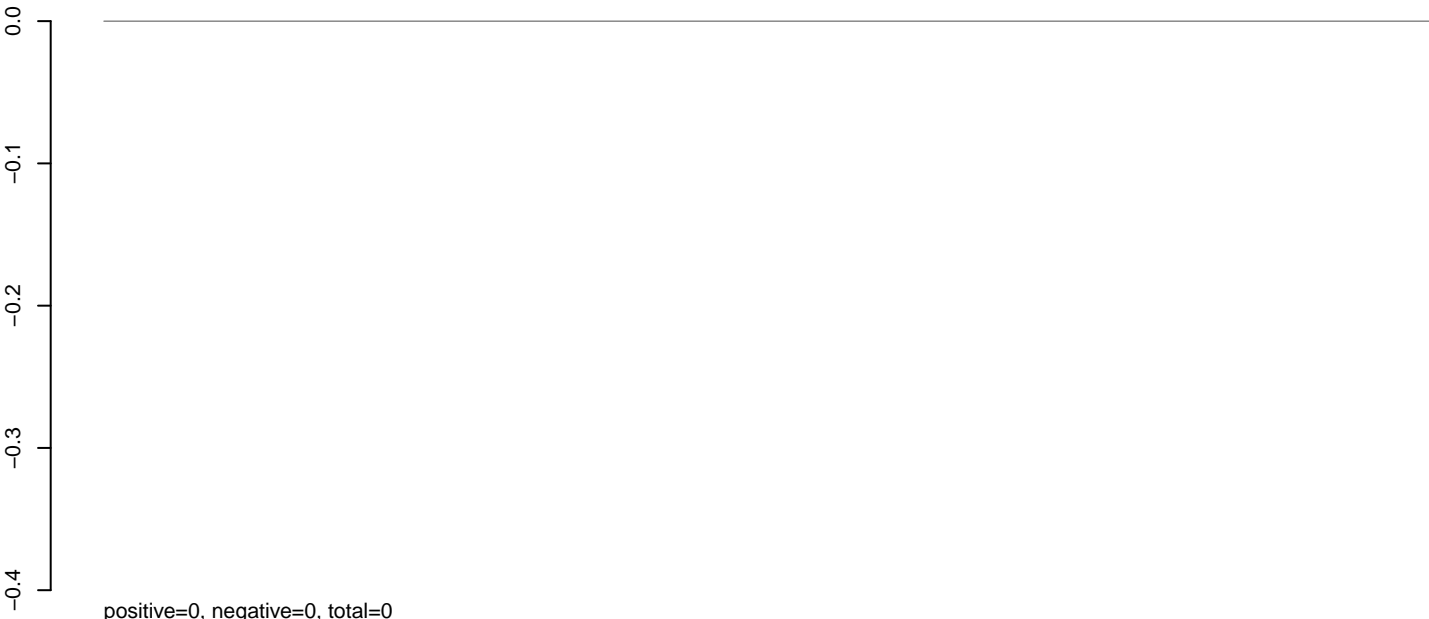
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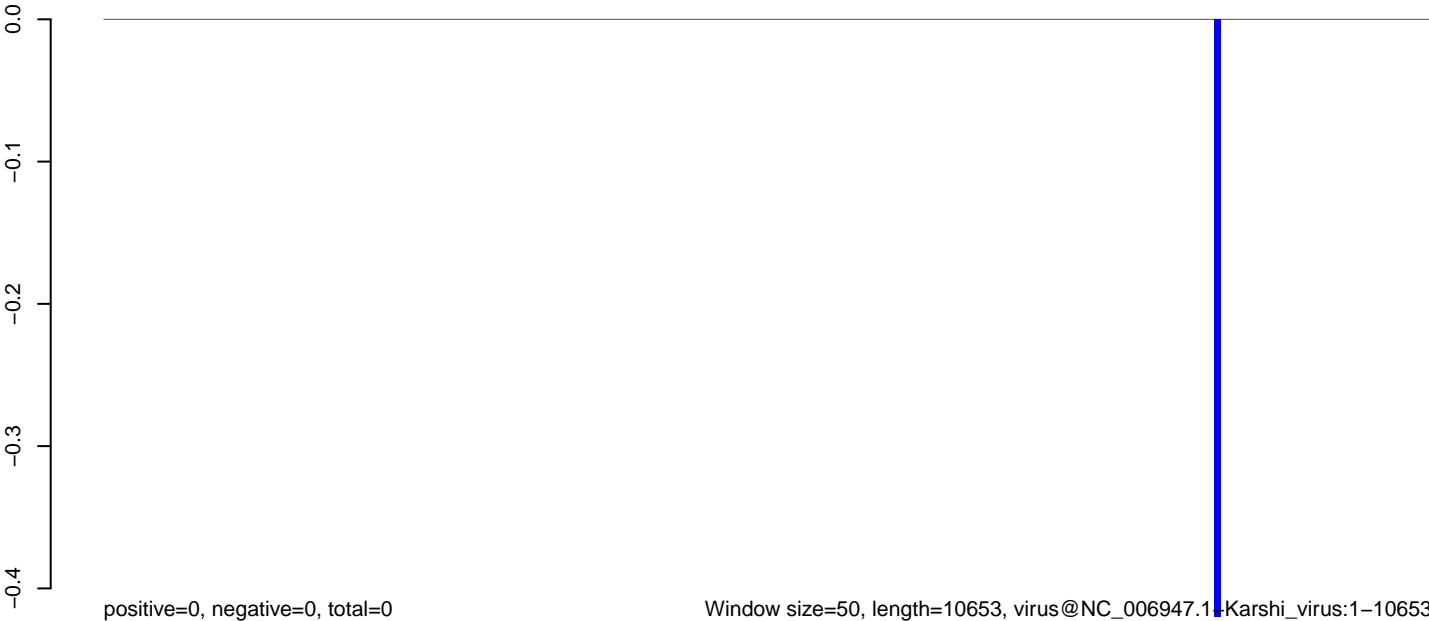
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



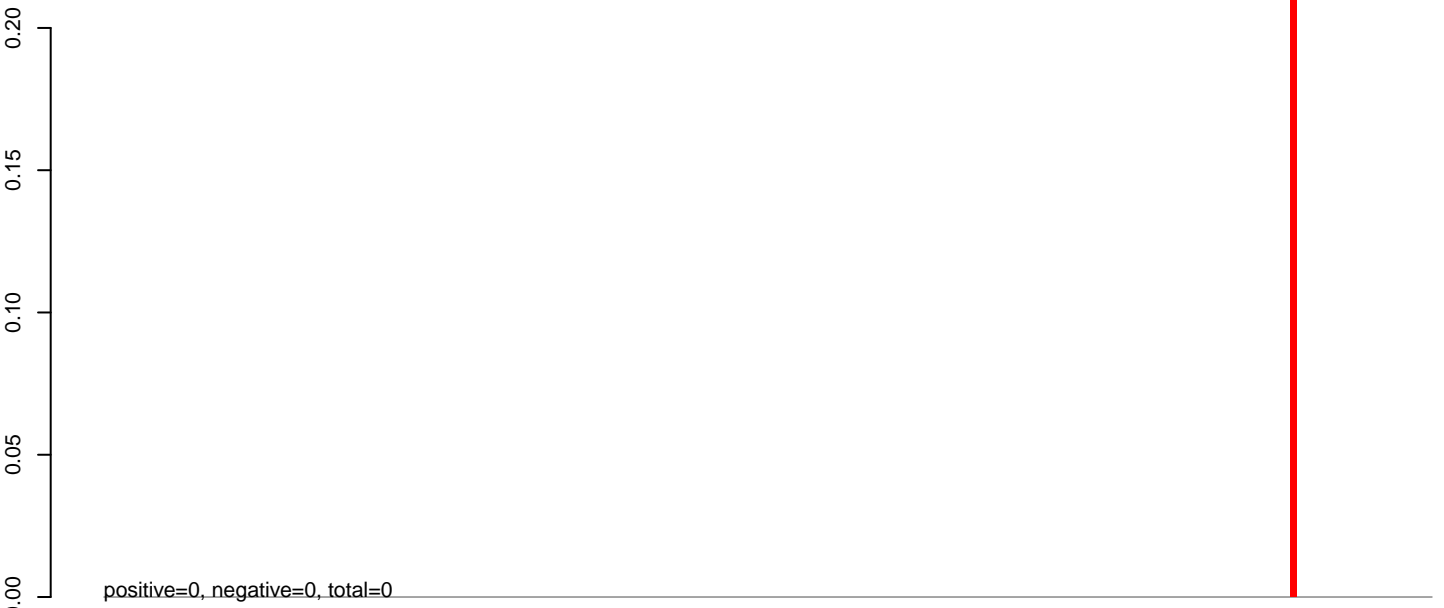
AeAeg_Aag2_DENV4.rep



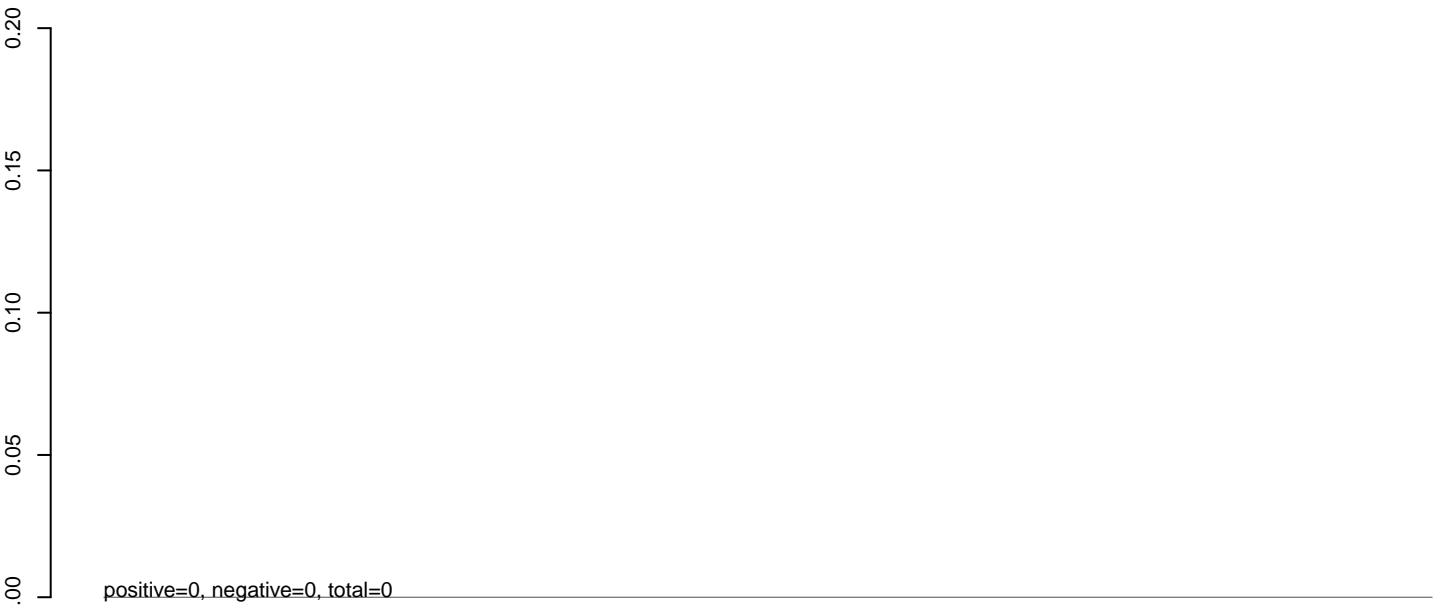
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0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV4.18_23.rep



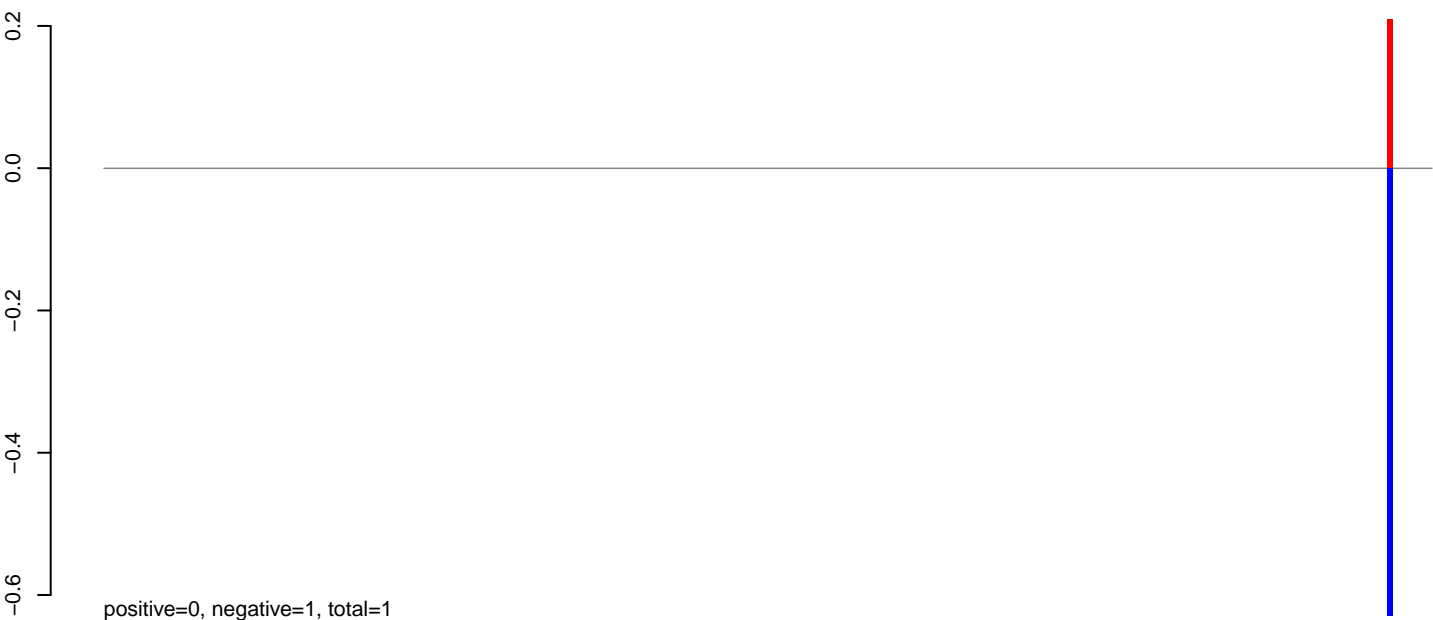
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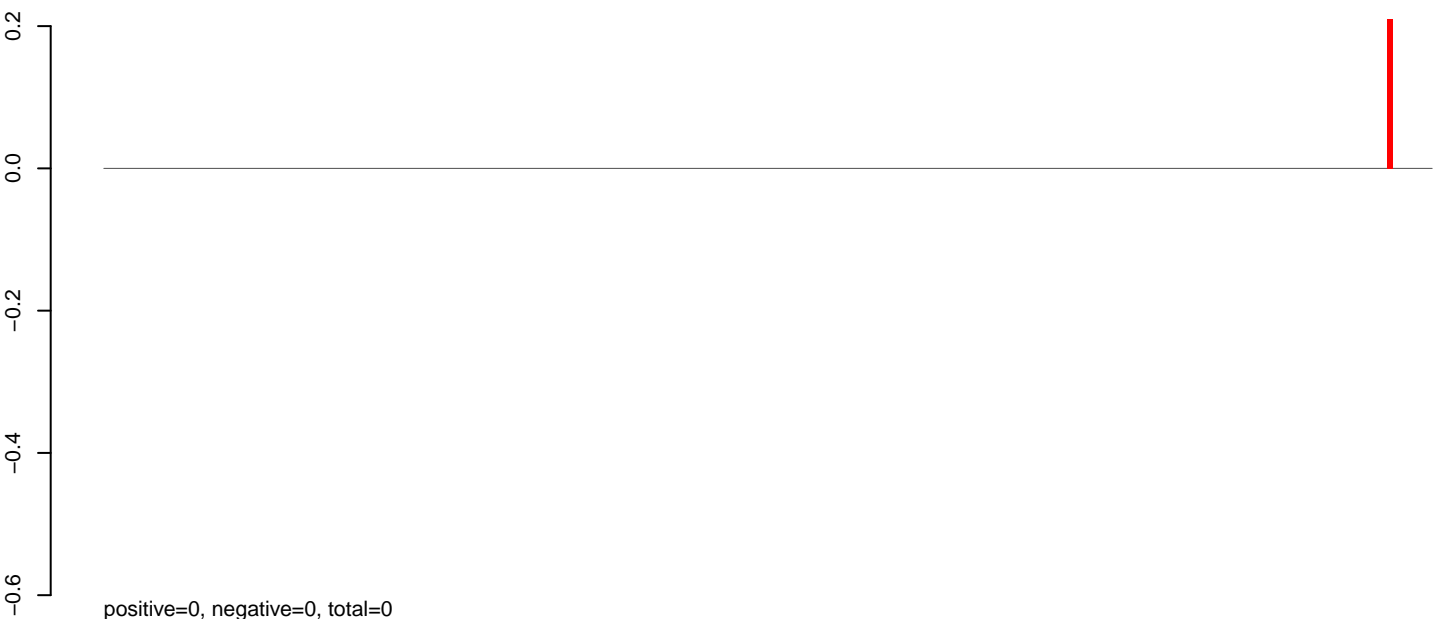
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



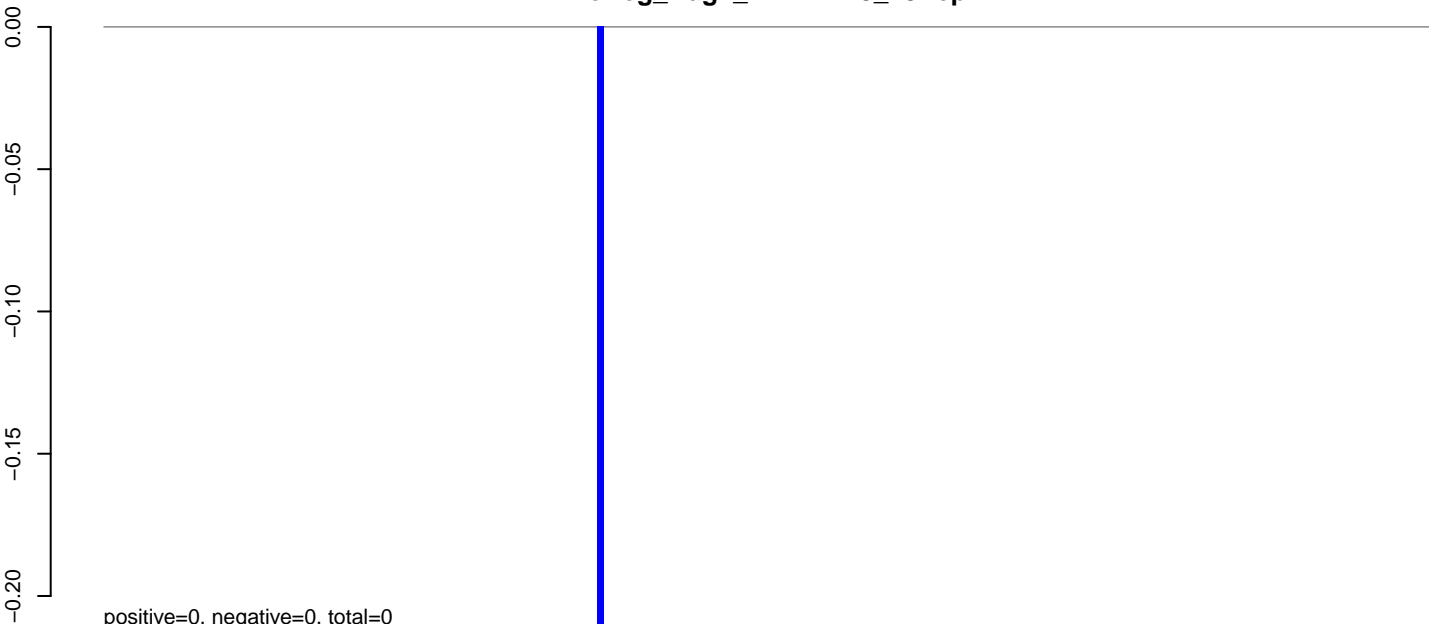
AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



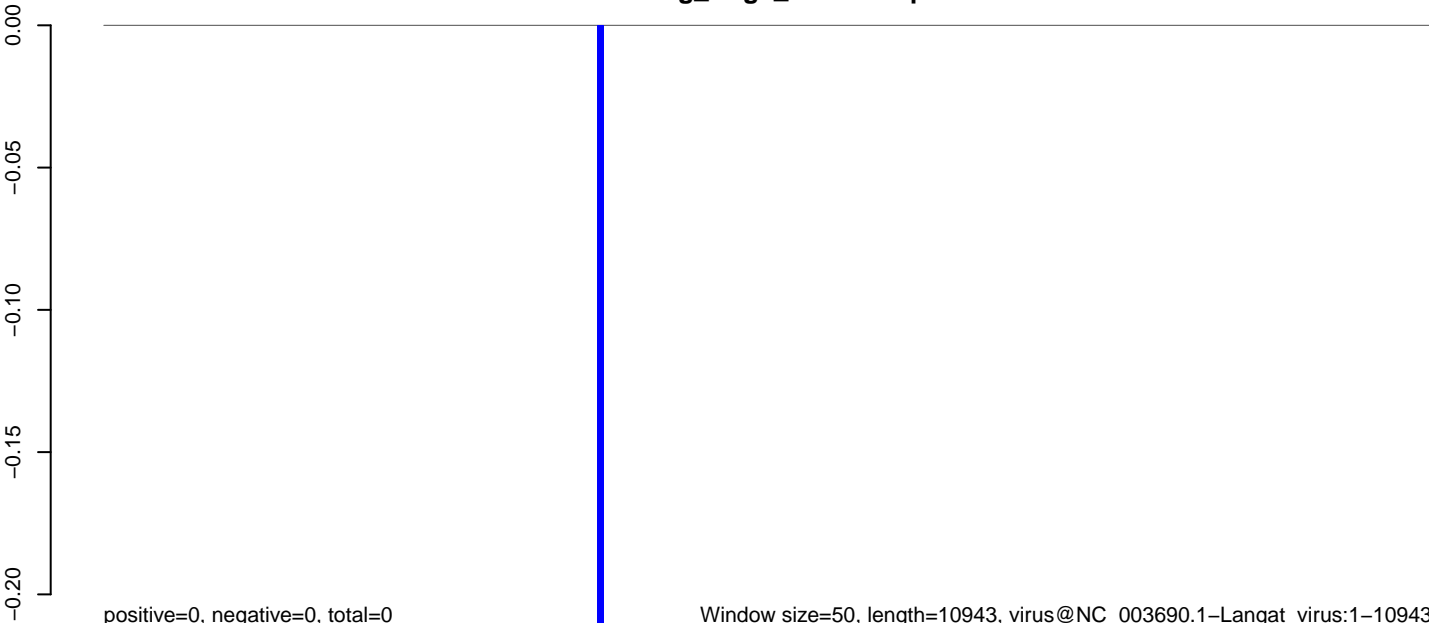
positive=0, negative=0, total=0

AeAeg_Aag2_DENV4.24_35.rep



positive=0, negative=0, total=0

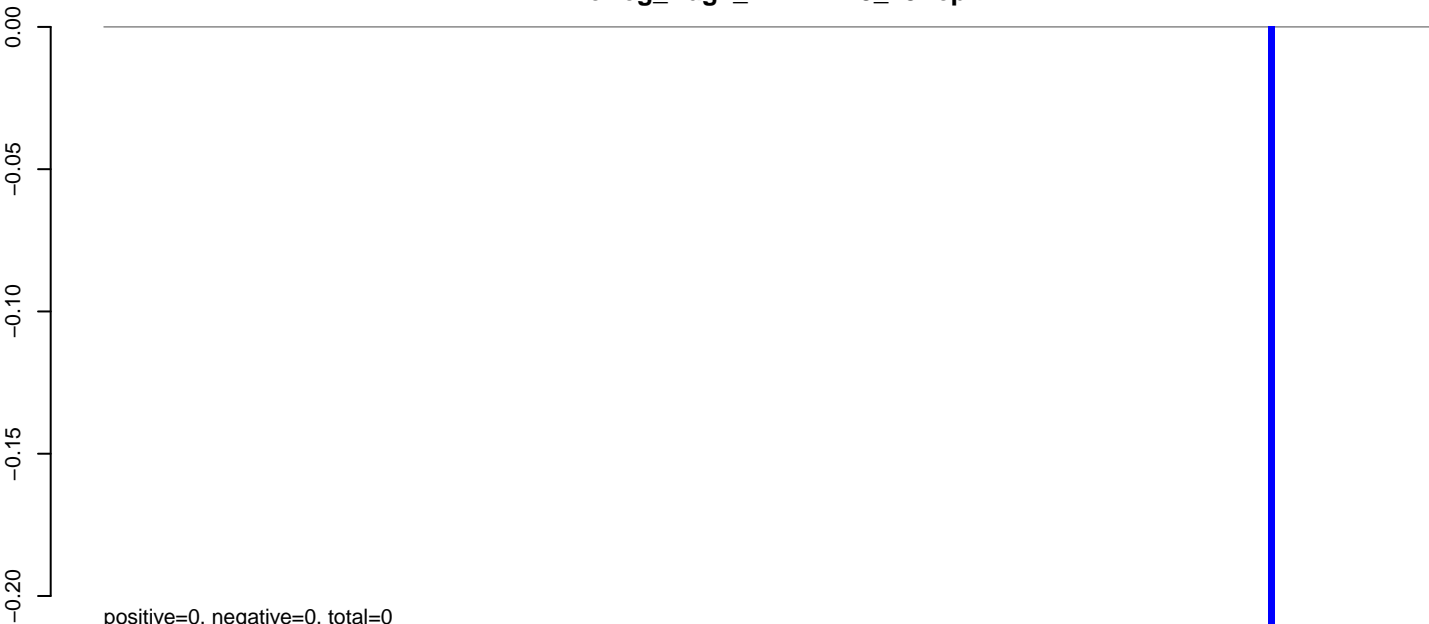
AeAeg_Aag2_DENV4.rep



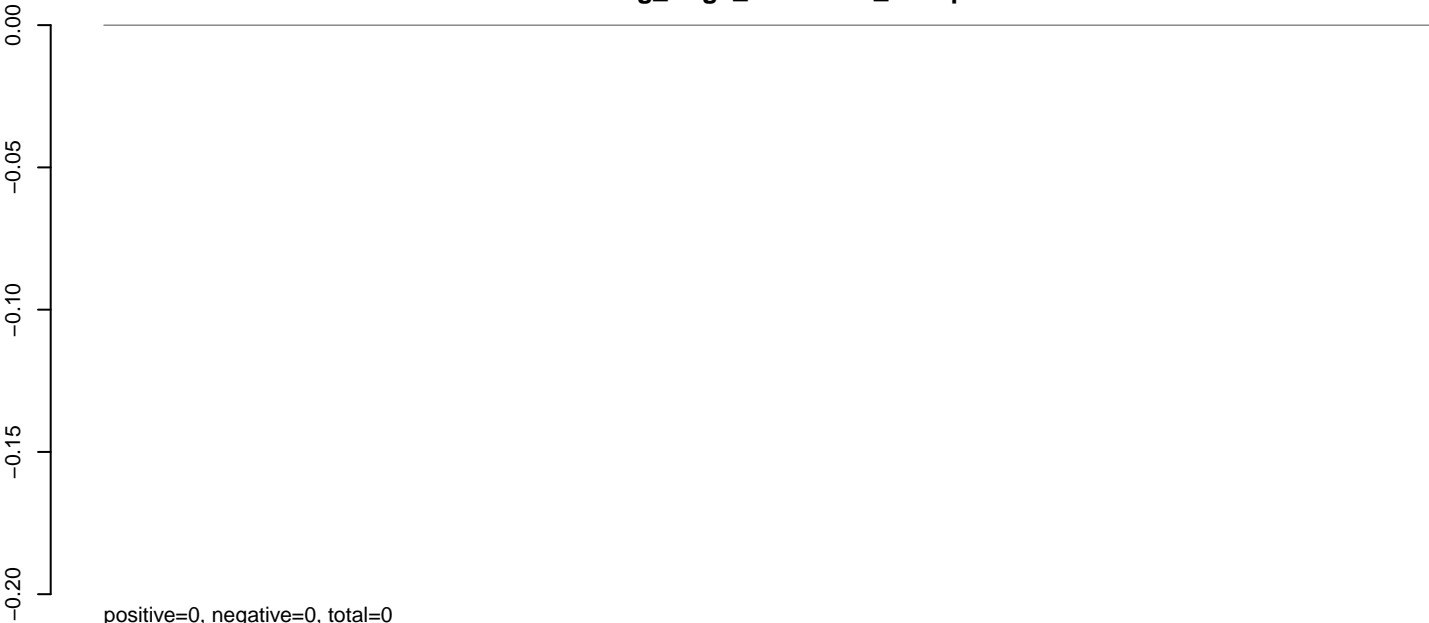
positive=0, negative=0, total=0

Window size=50, length=10943, virus@NC_003690.1-Langat_virus:1-10943

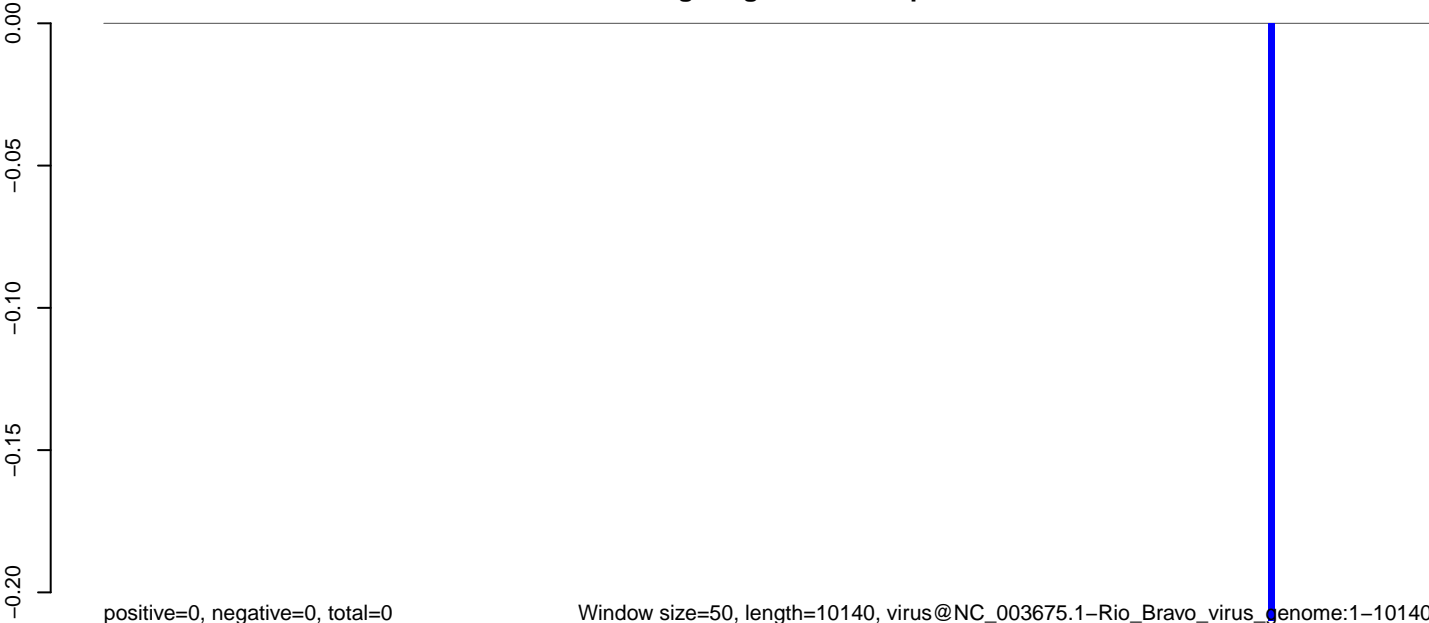
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



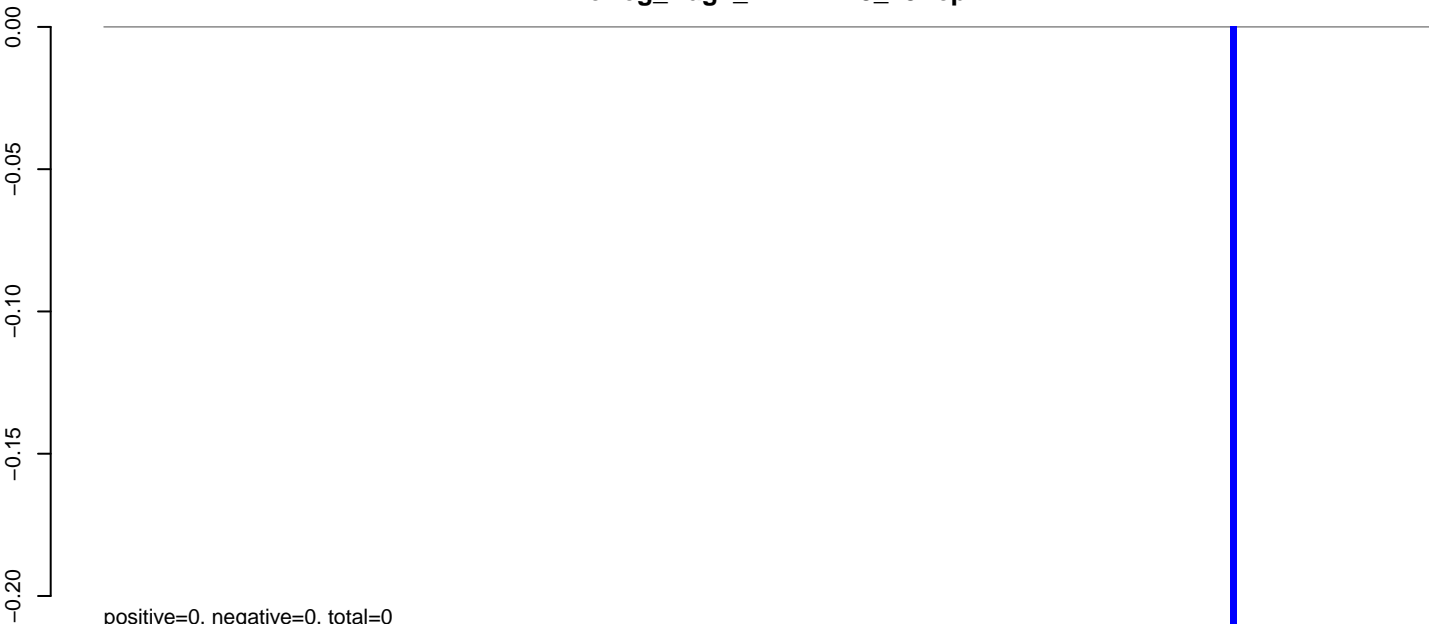
AeAeg_Aag2_DENV4.rep



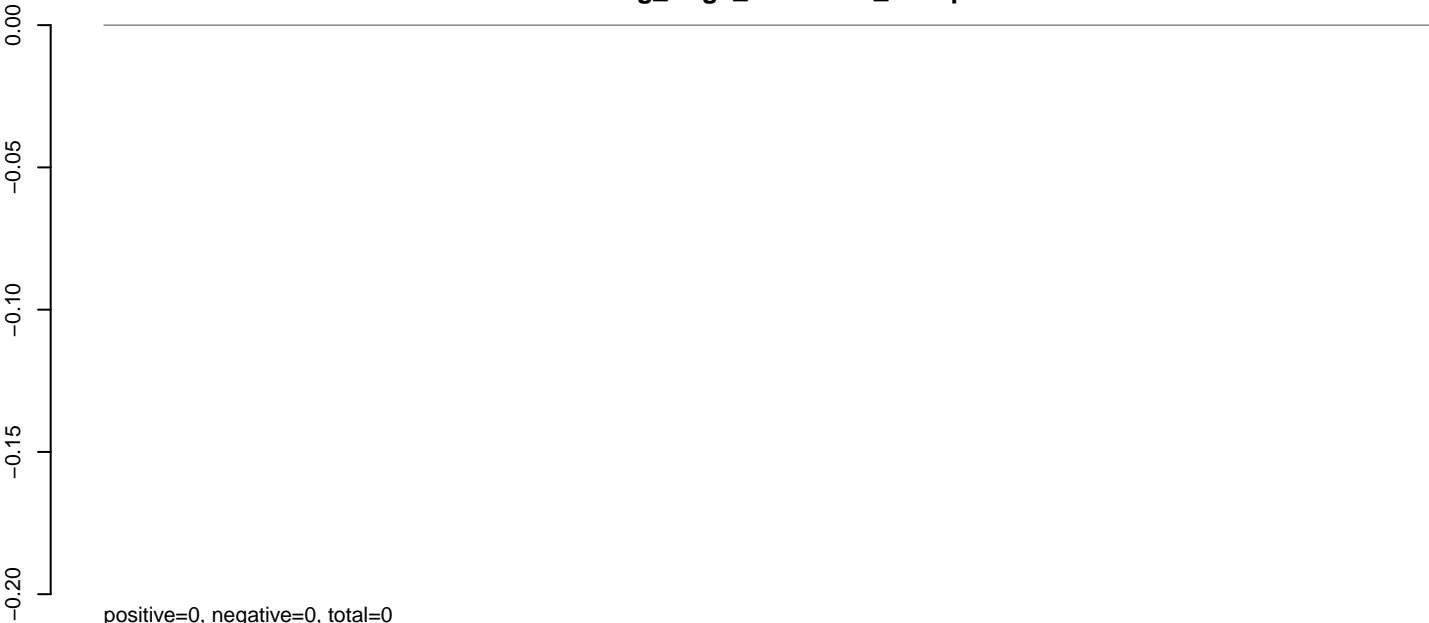
Window size=50, length=10140, virus@NC_003675.1-Rio_Bravo_virus_genome:1-10140

0 2000 4000 6000 8000 10000

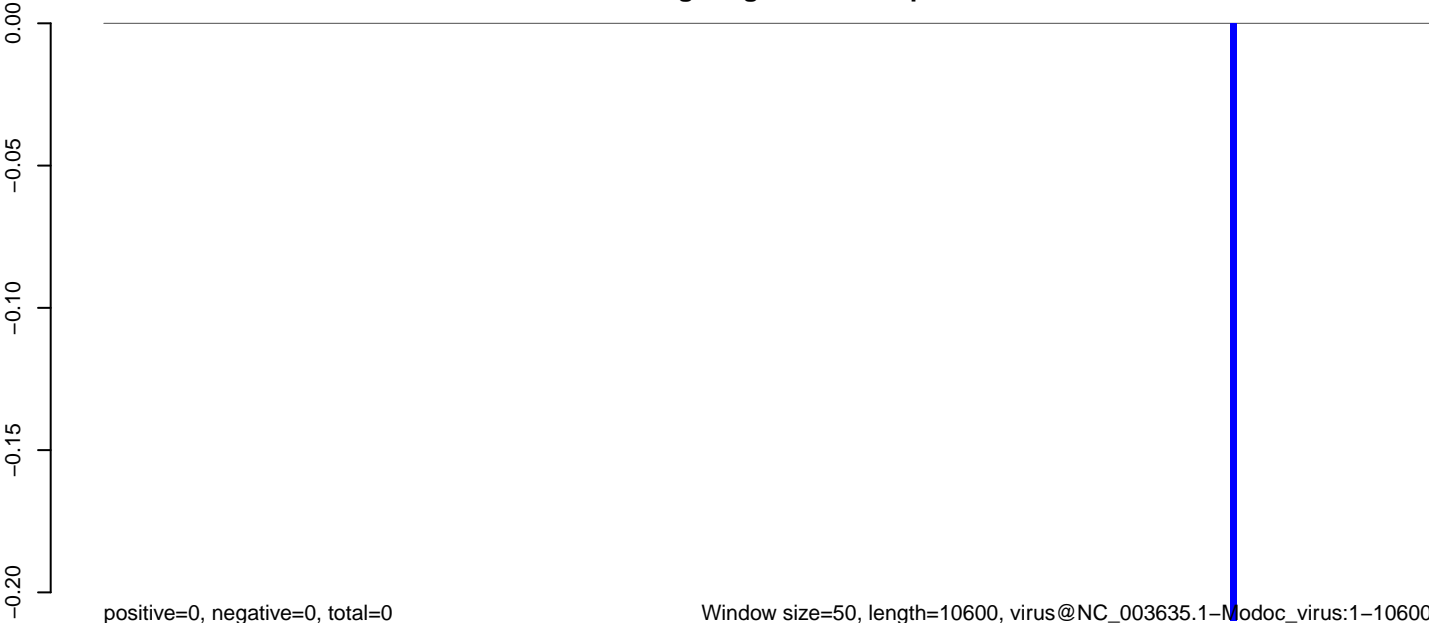
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

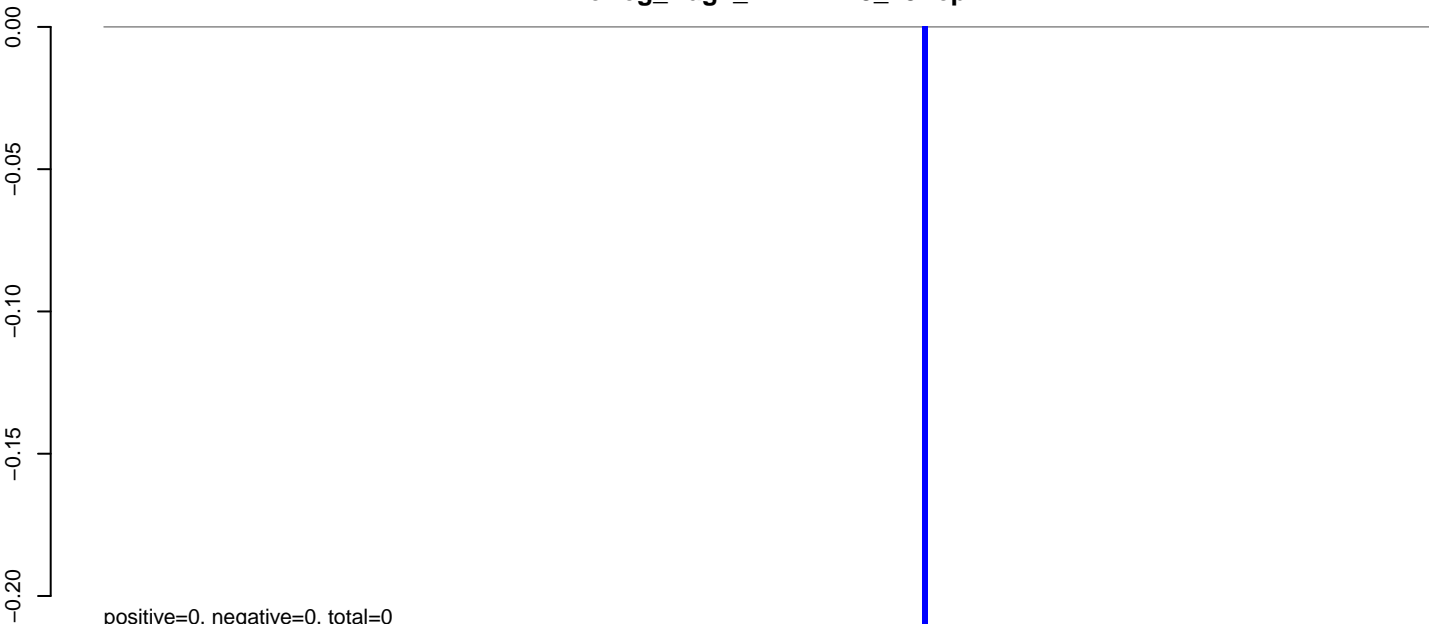


AeAeg_Aag2_DENV4.rep

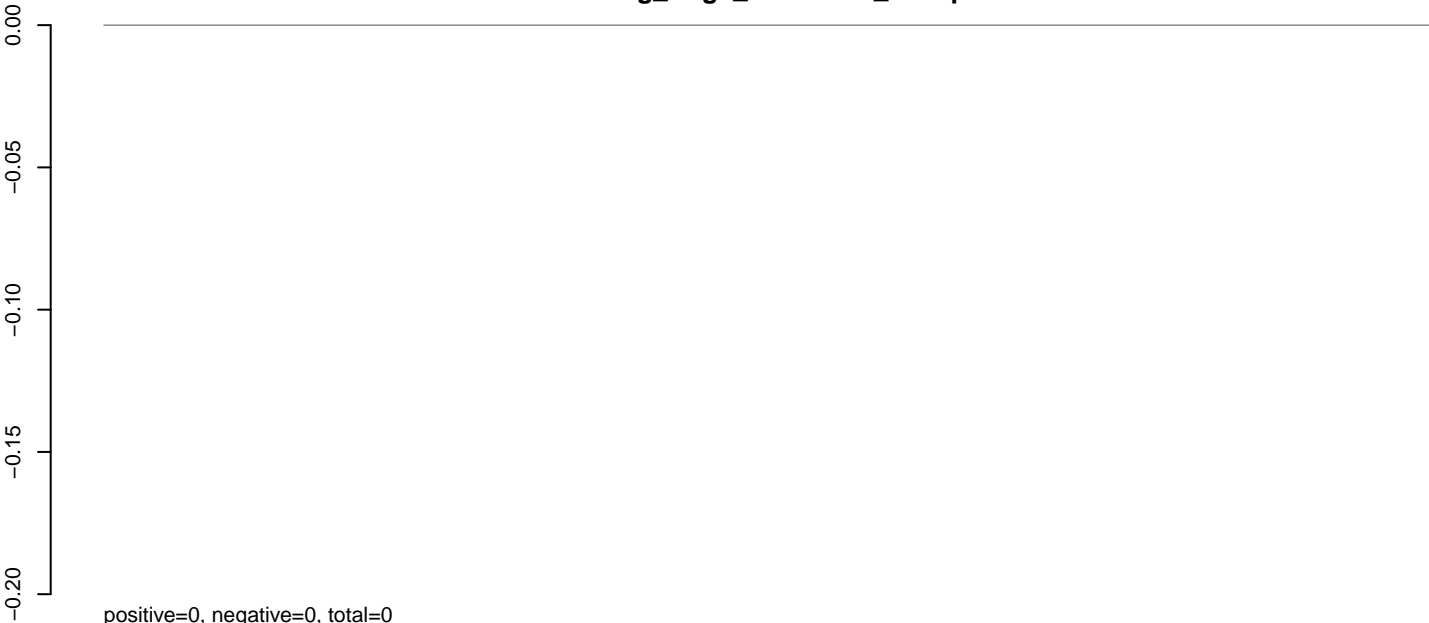


0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV4.18_23.rep



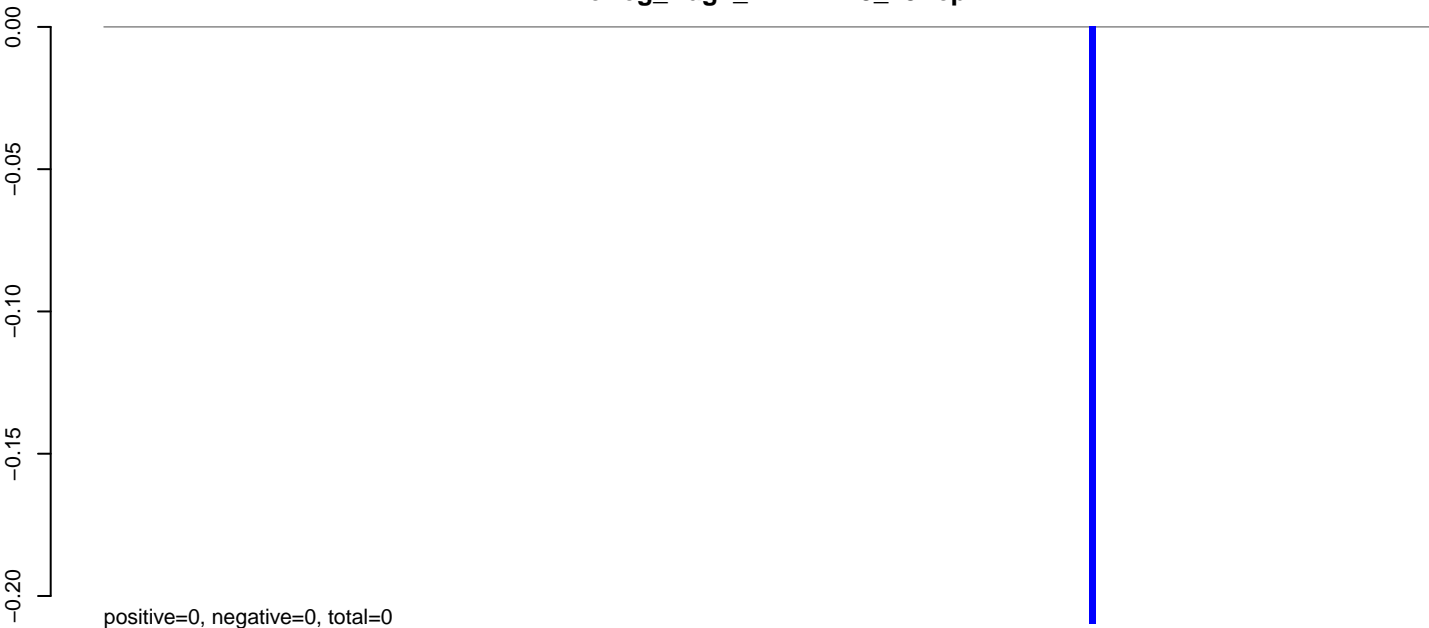
AeAeg_Aag2_DENV4.24_35.rep



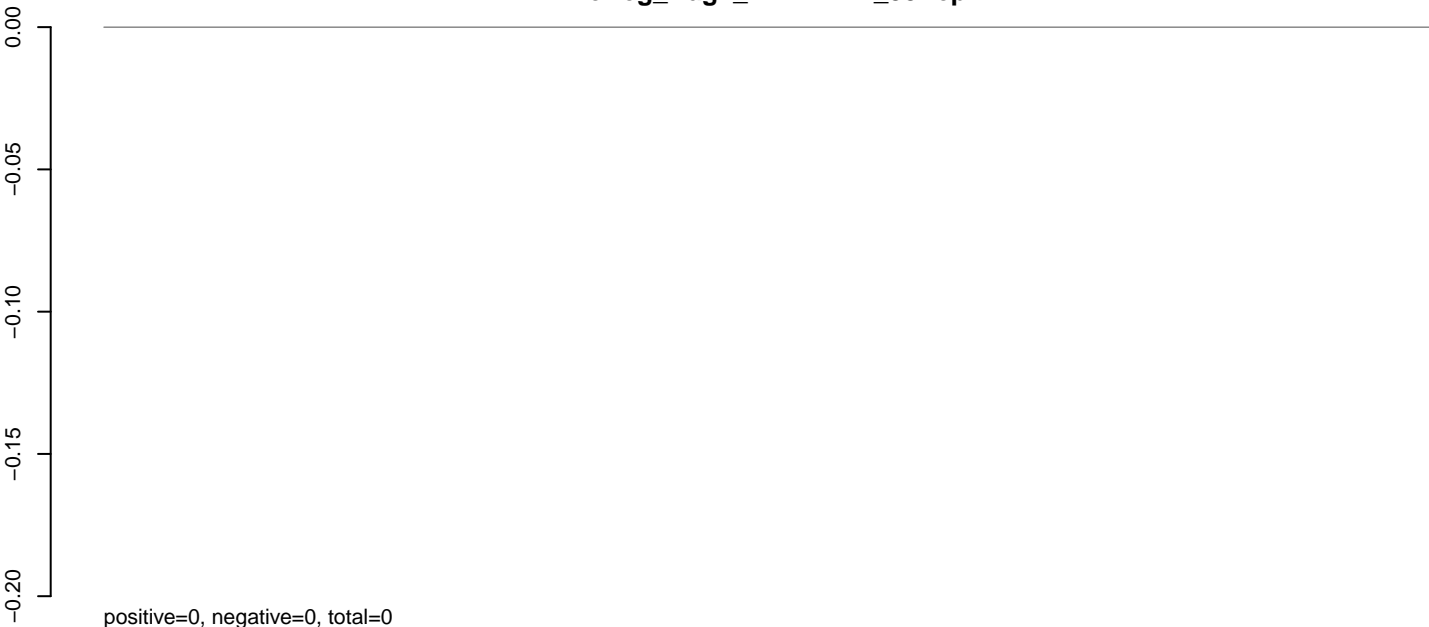
AeAeg_Aag2_DENV4.rep



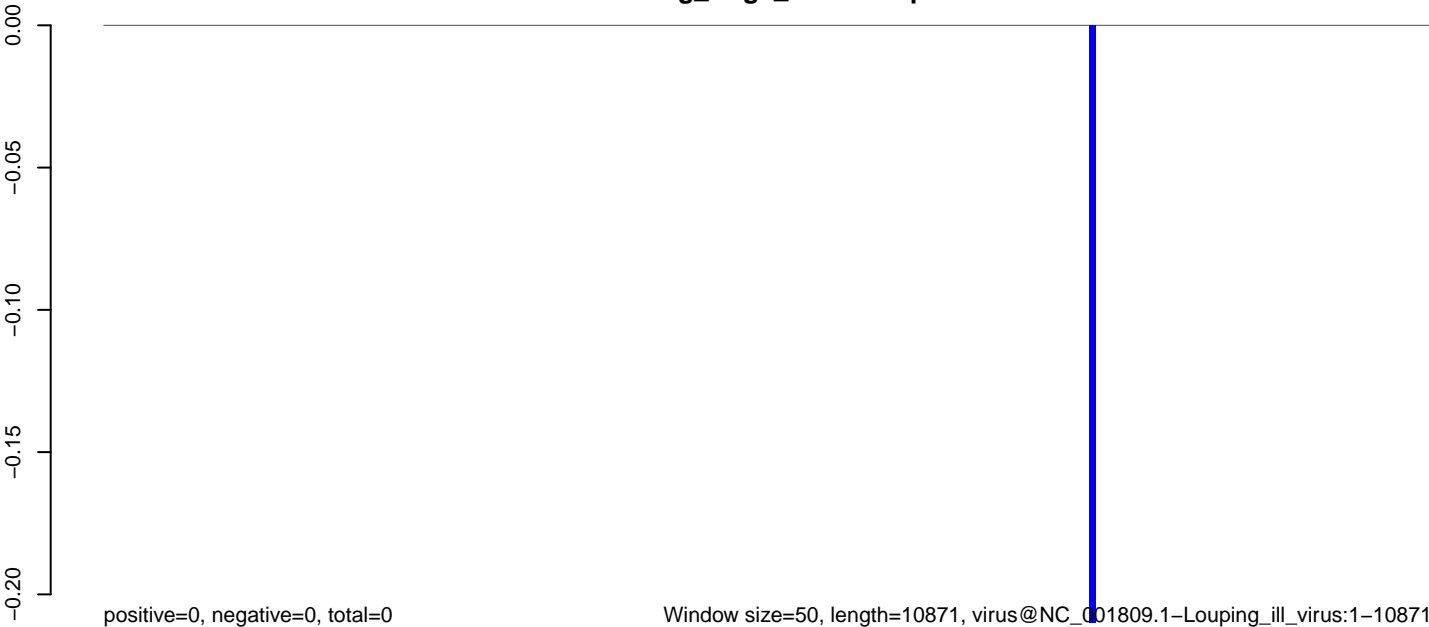
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

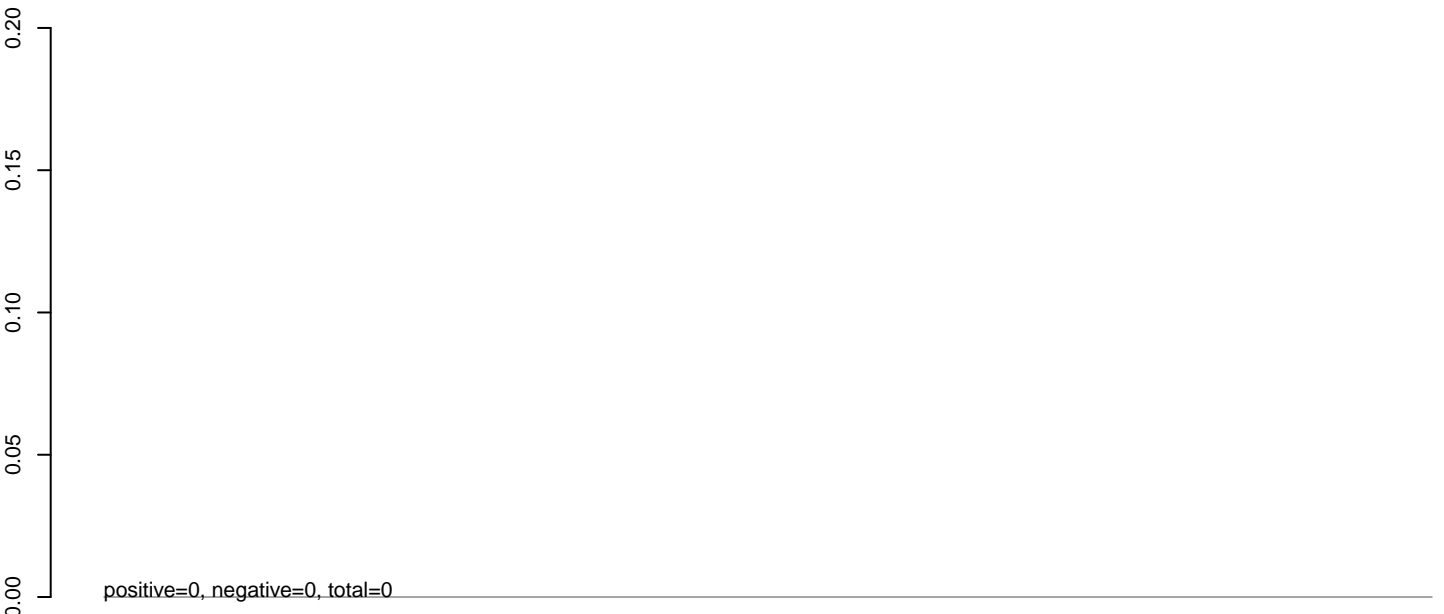


AeAeg_Aag2_DENV4.rep

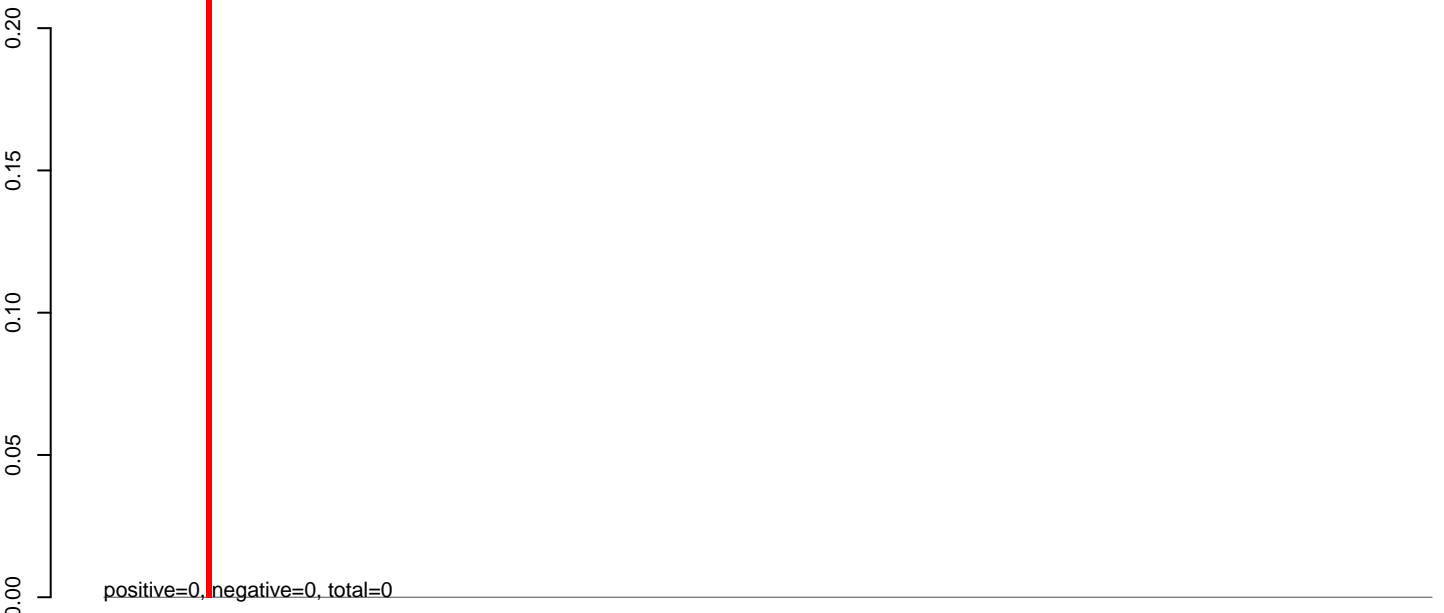


0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

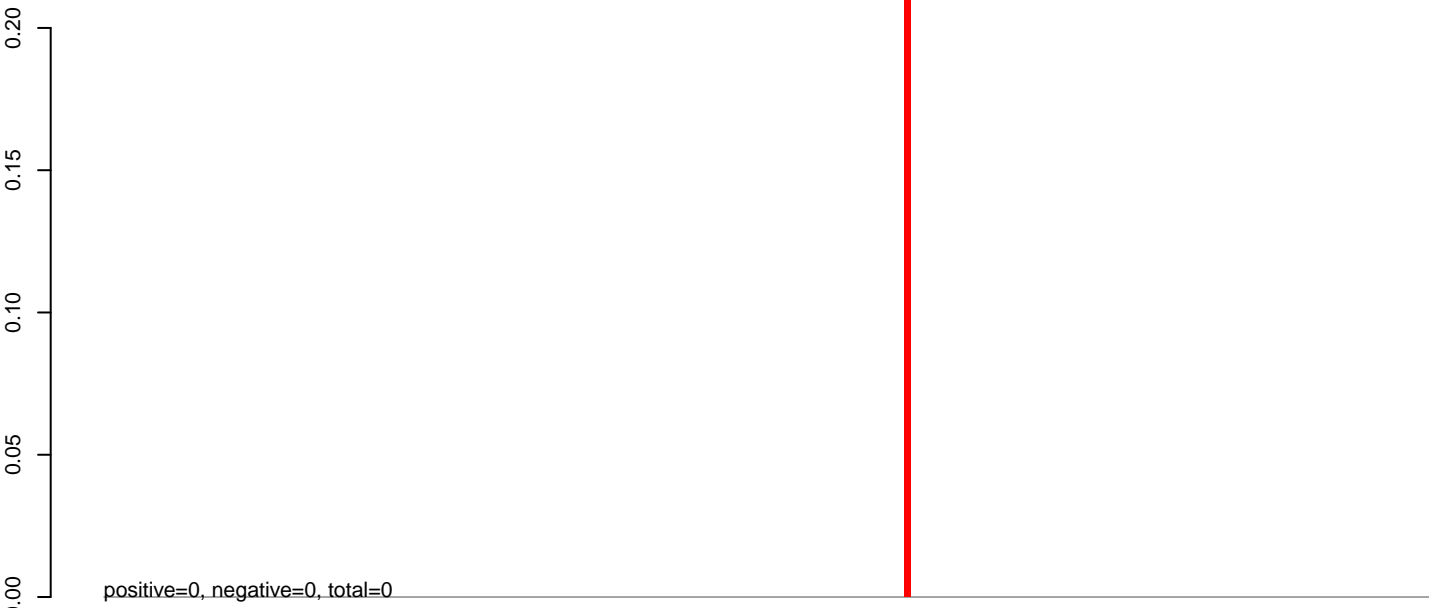


AeAeg_Aag2_DENV4.rep

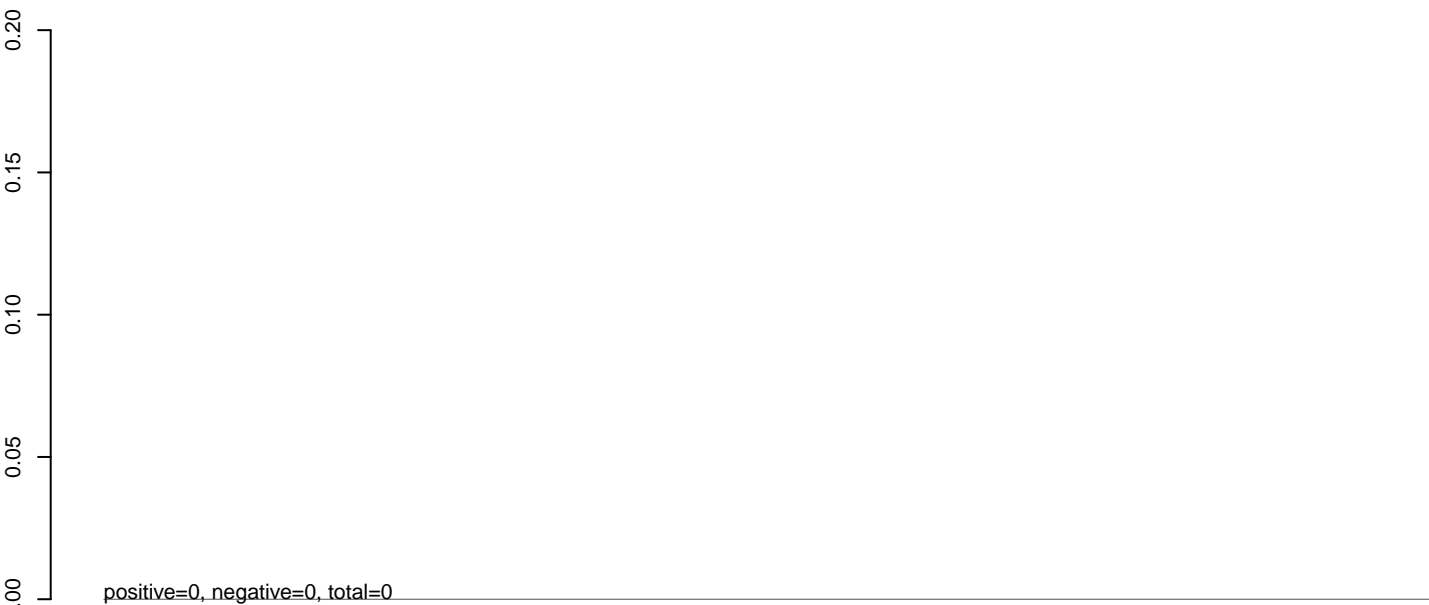


Window size=50, length=12916, virus@MH237595.1-Aedes_aegypti_anphevirus_strain_1a:1-12916

AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

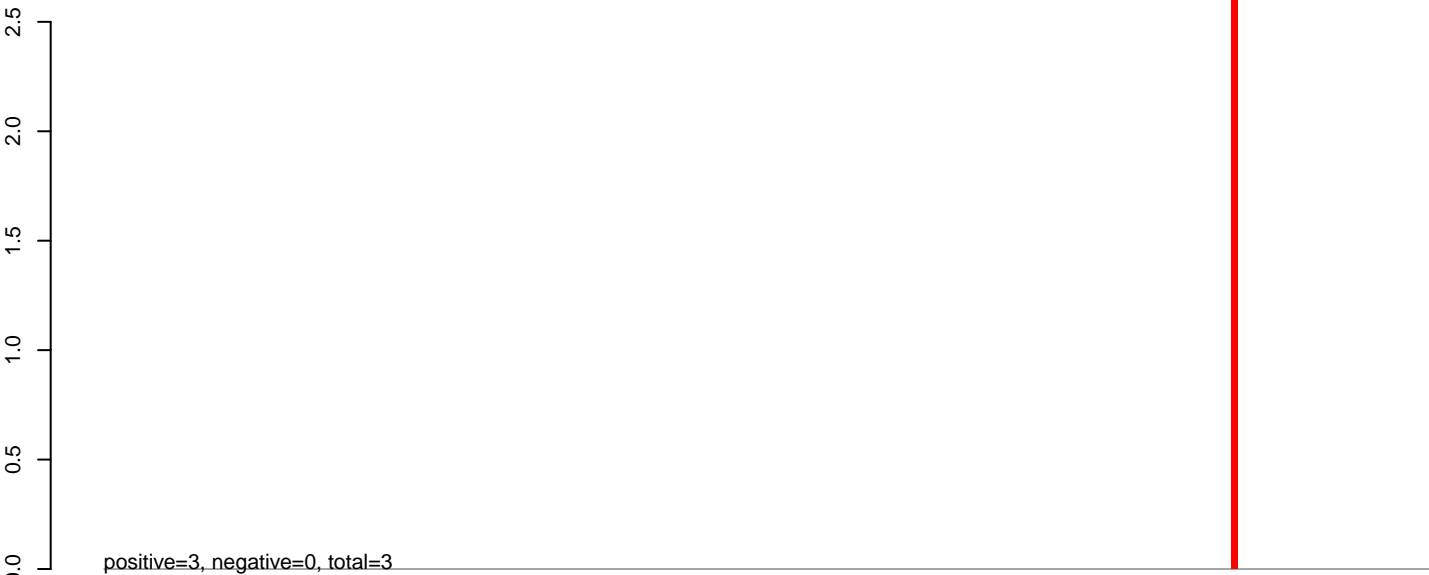


AeAeg_Aag2_DENV4.rep

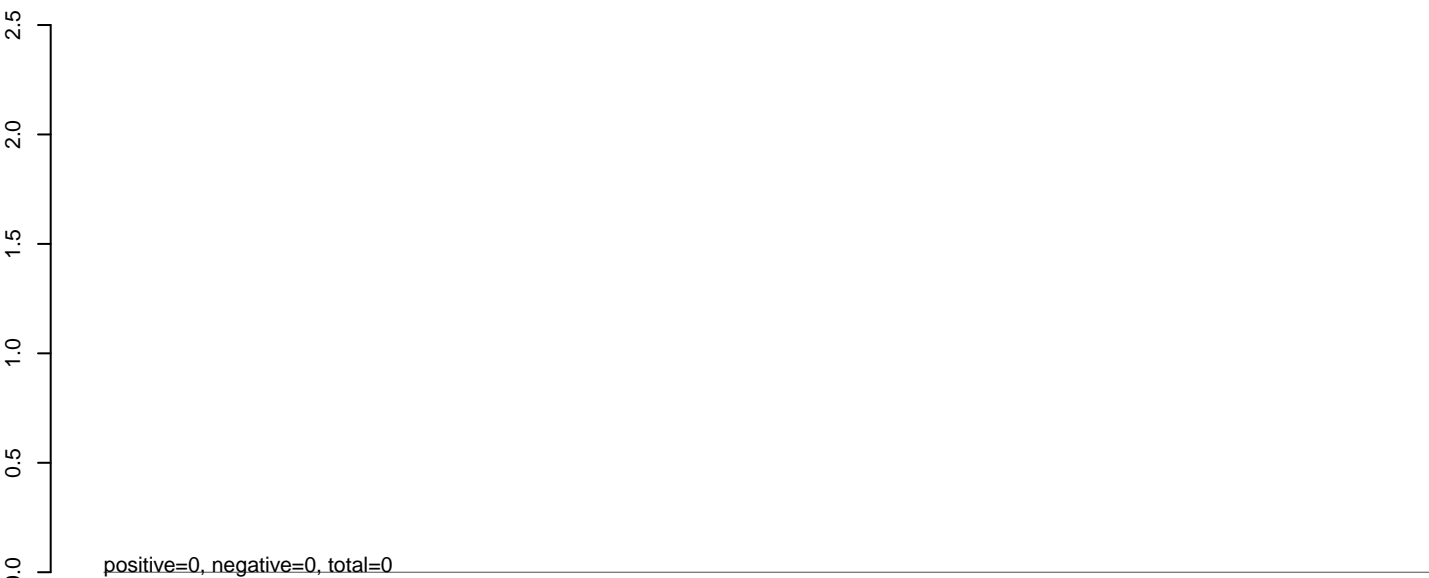


Window size=50, length=10749, virus@KX669682.1-Calbertado_virus_isolate_Cal_mos_2_polyprotein_gene:1-10749

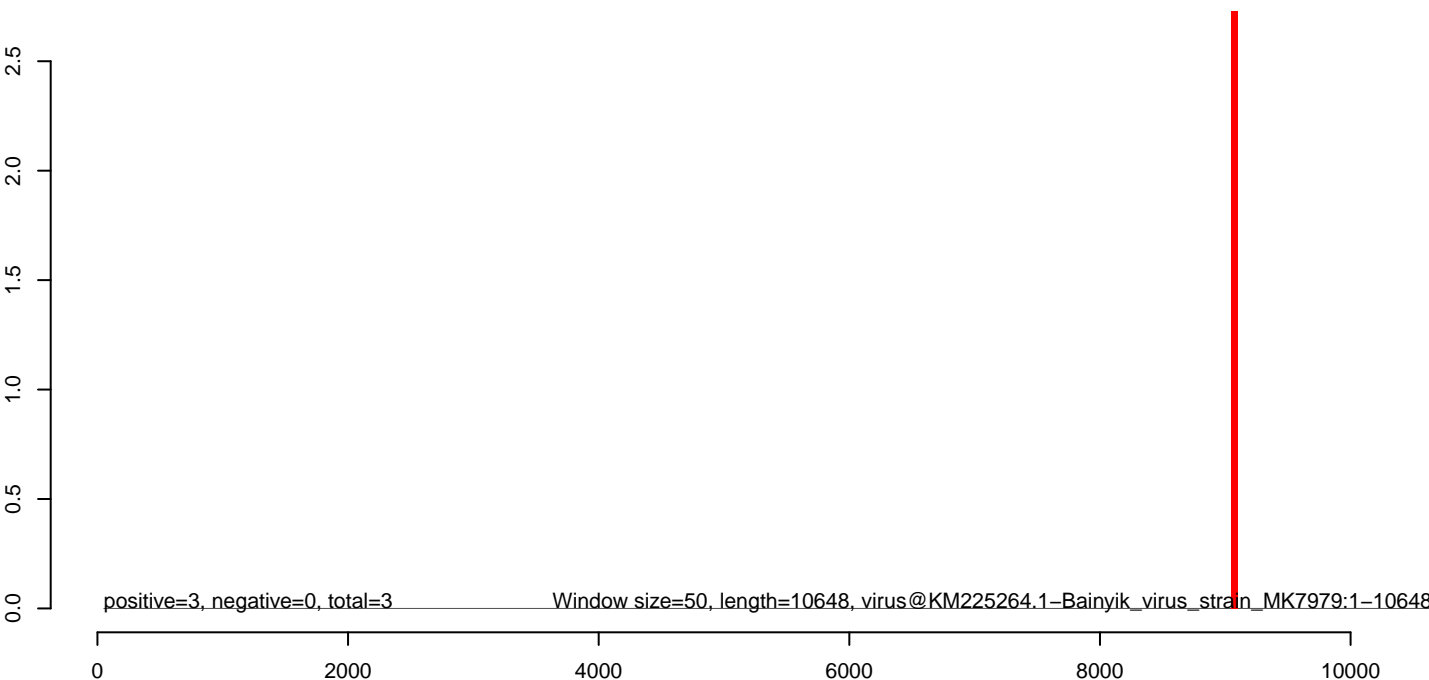
AeAeg_Aag2_DENV4.18_23.rep



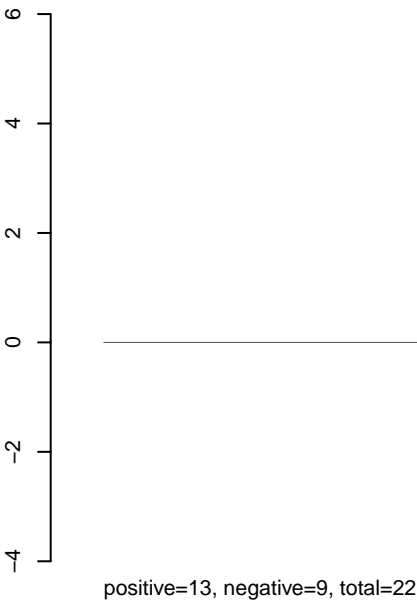
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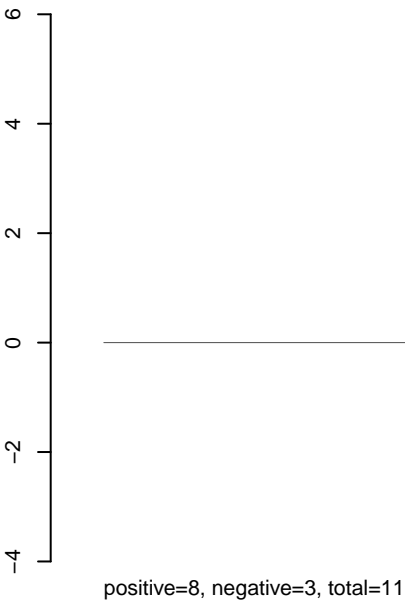
AeAeg_Aag2_DENV4.rep



AeAeg_Aag2_DENV4.18_23.rep



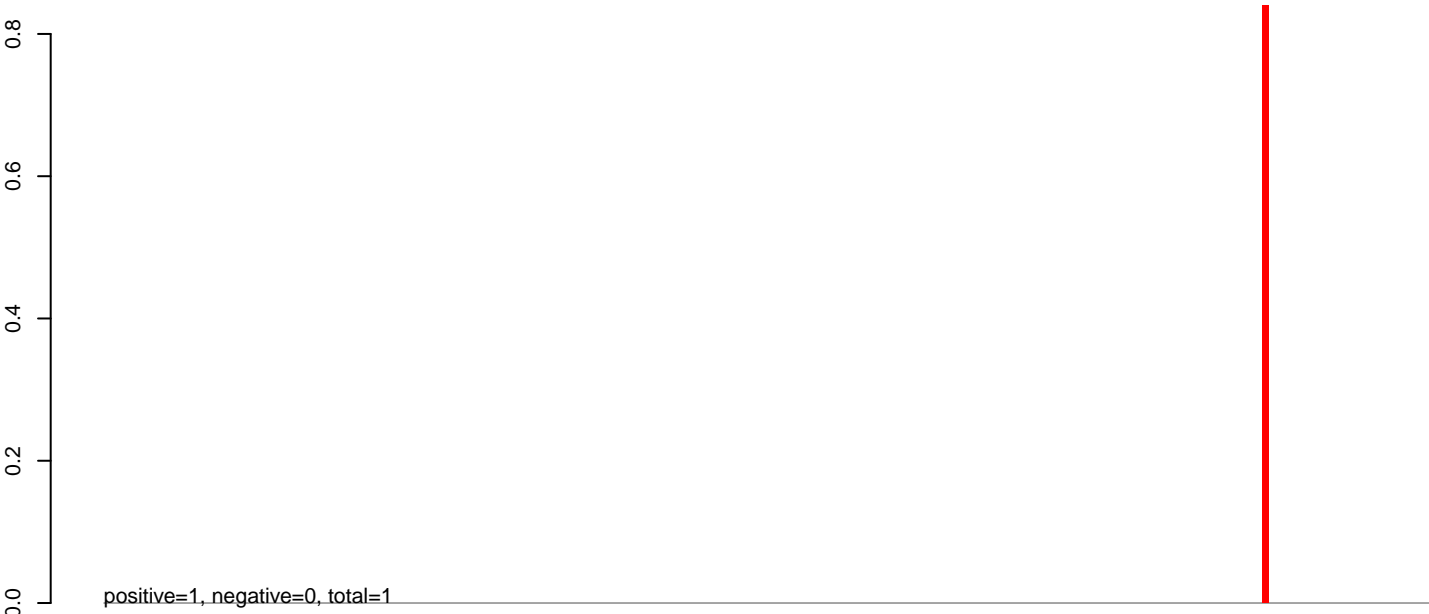
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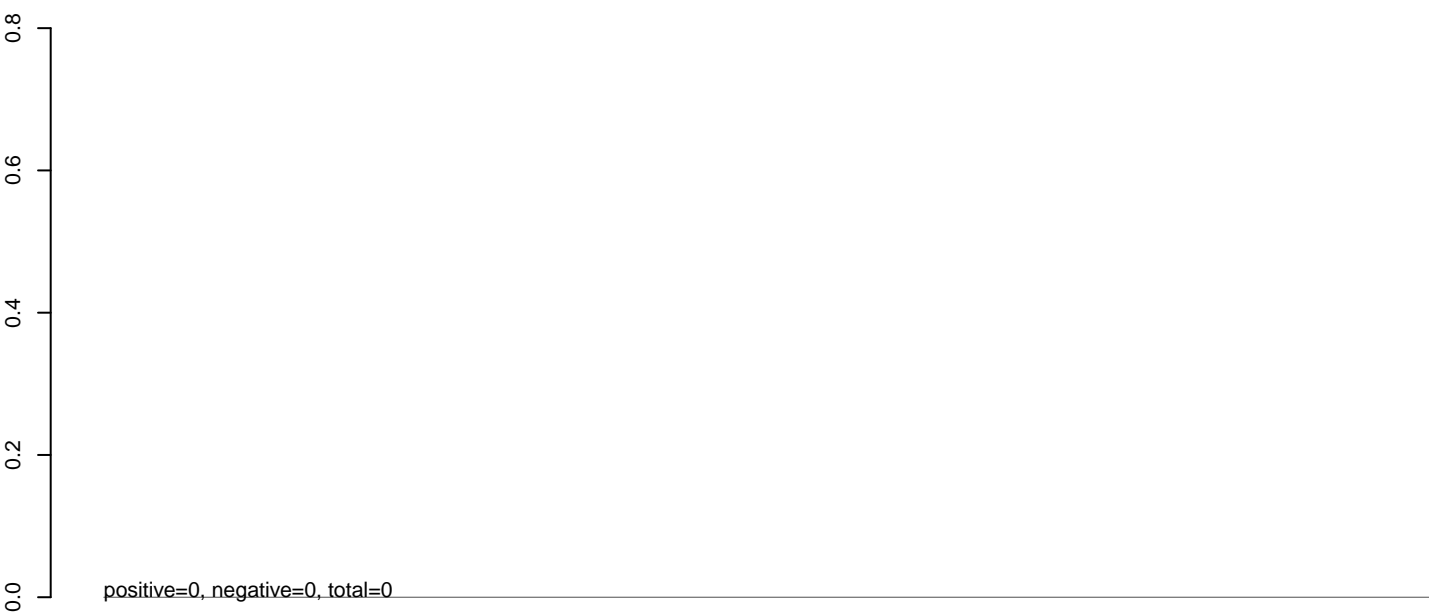
AeAeg_Aag2_DENV4.rep



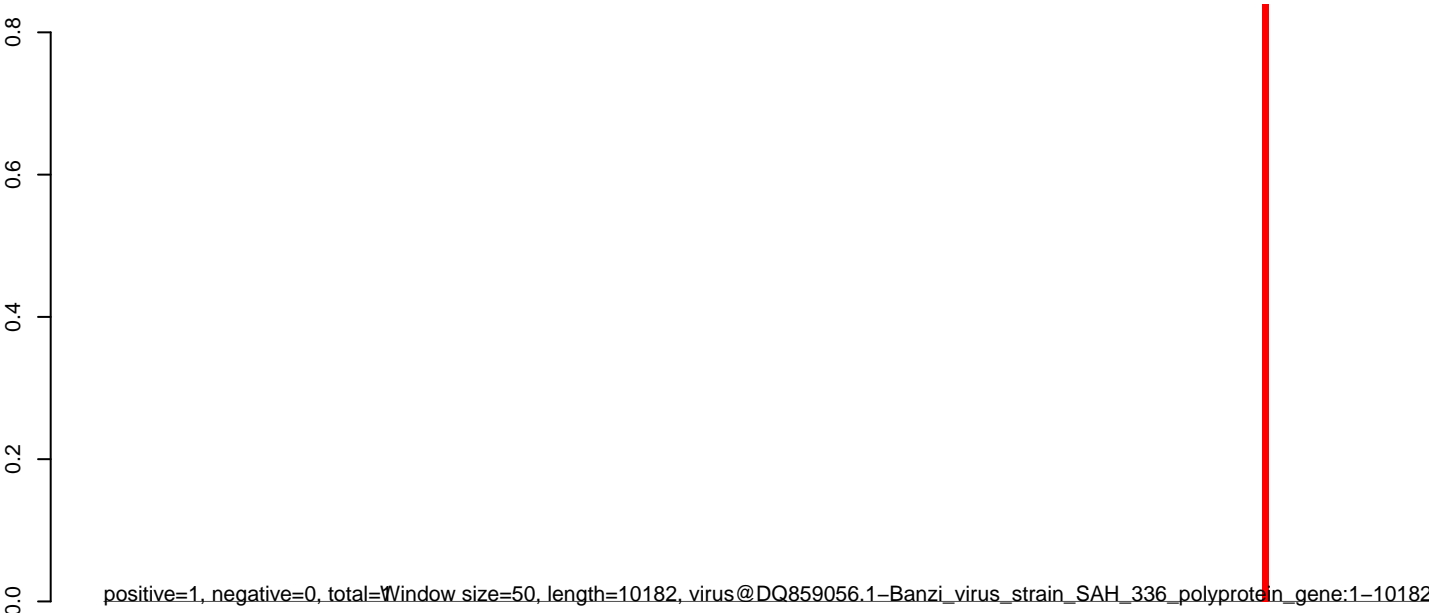
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AeAeg_Aag2_DENV4.24_35.rep

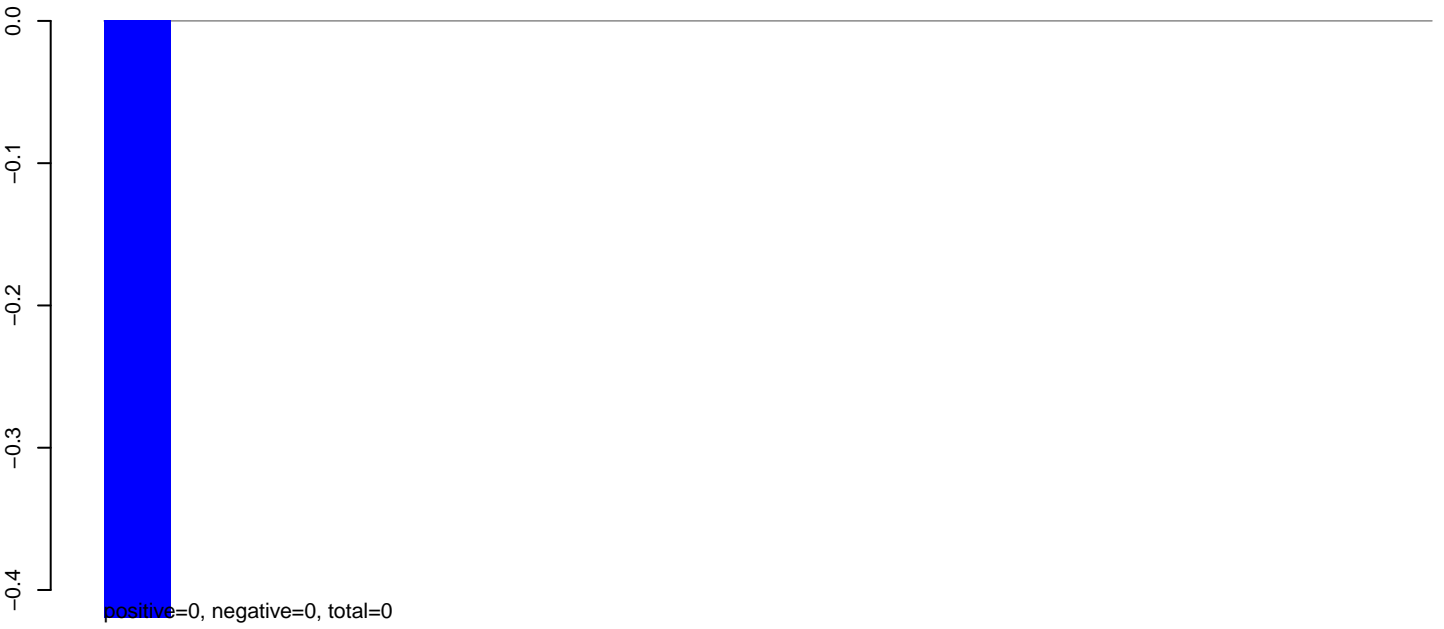


AeAeg_Aag2_DENV4.rep

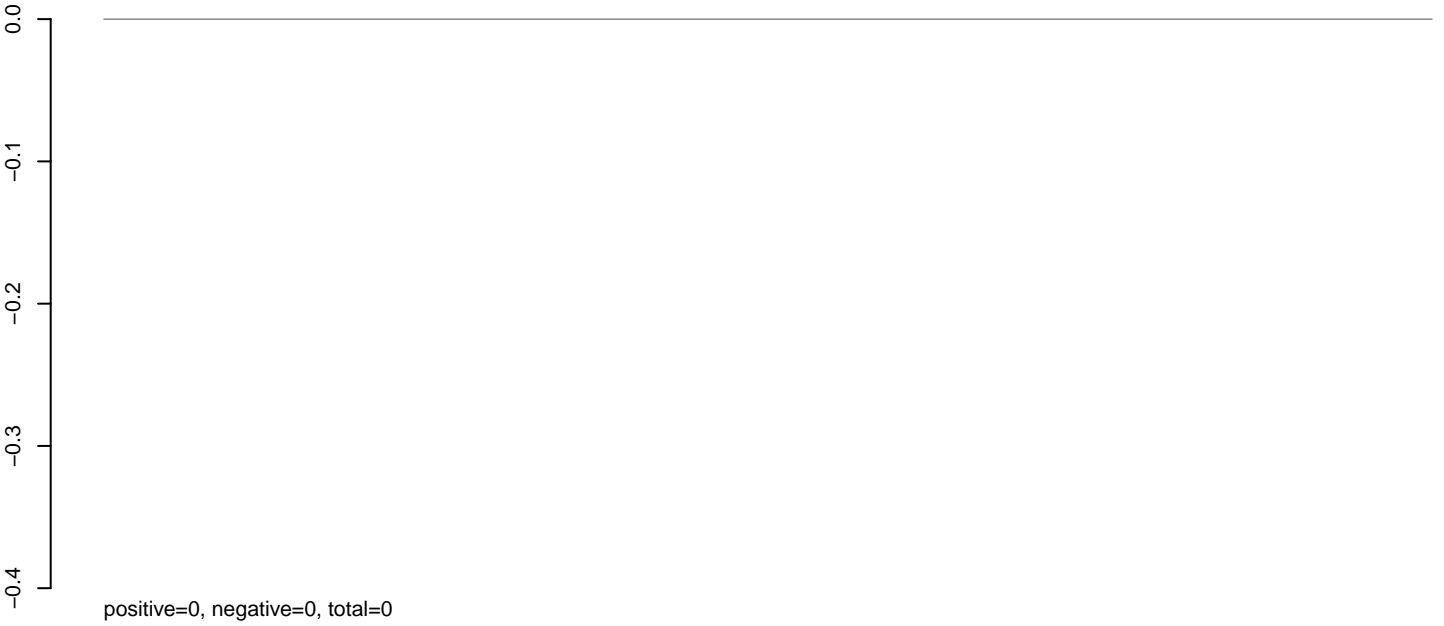


0 2000 4000 6000 8000 10000

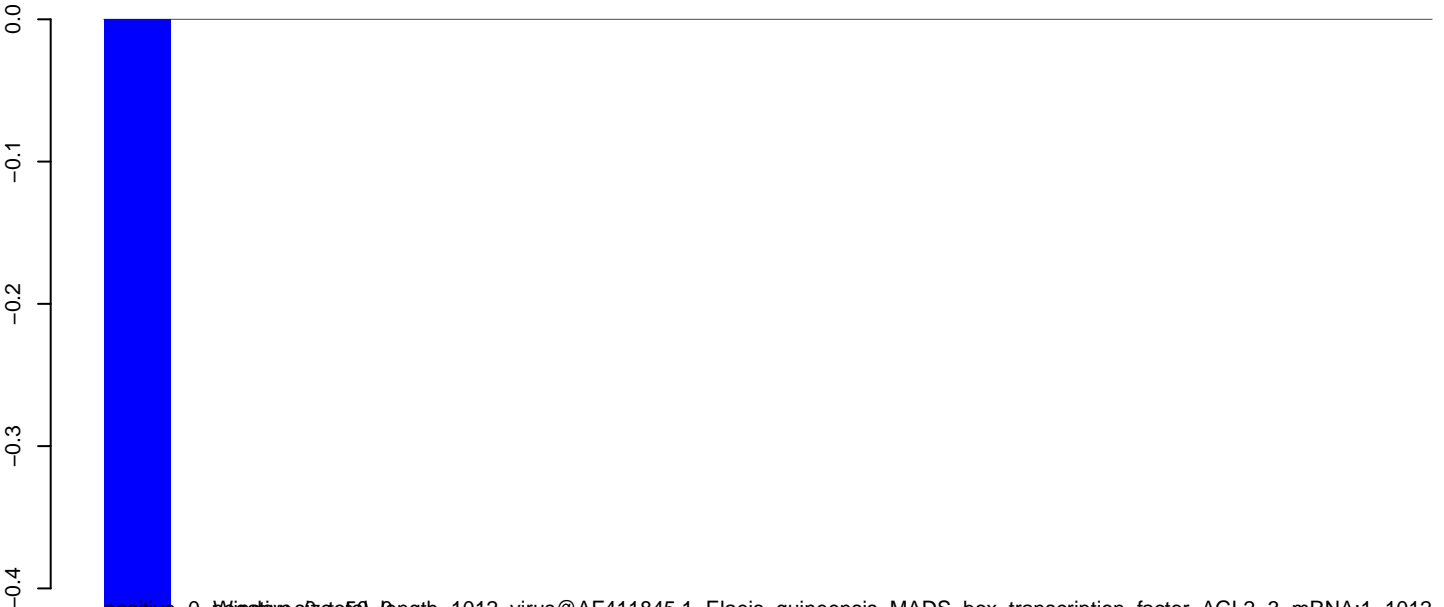
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



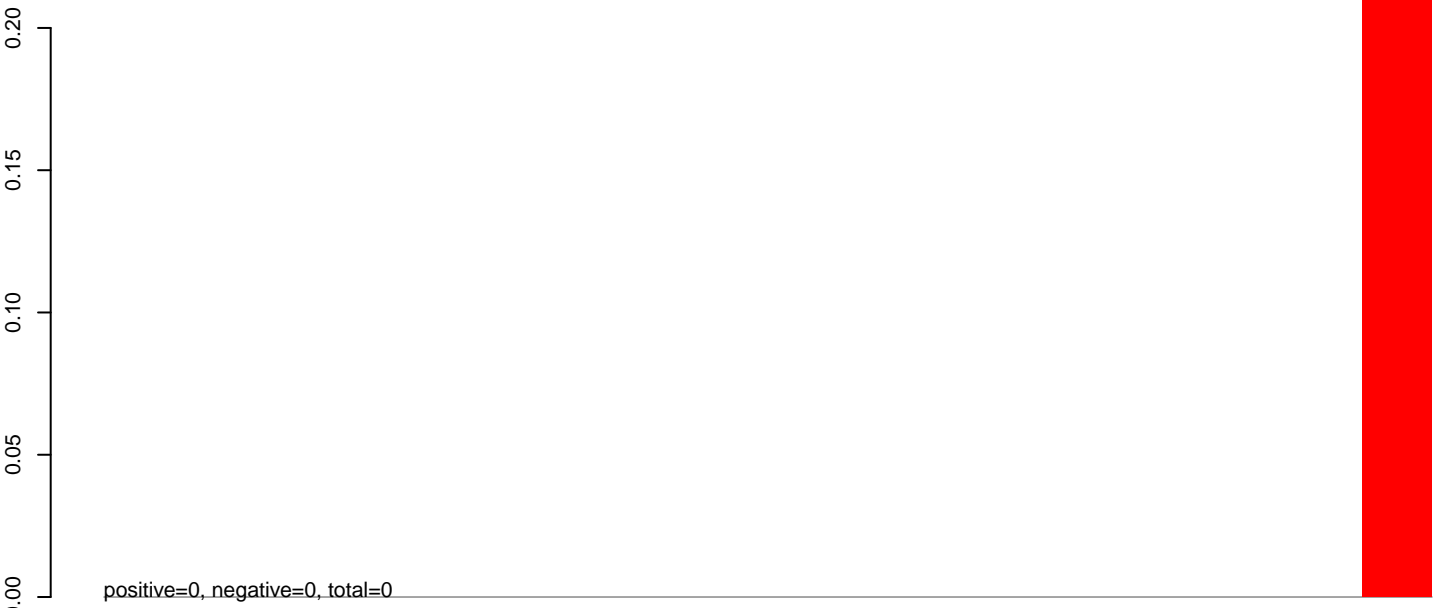
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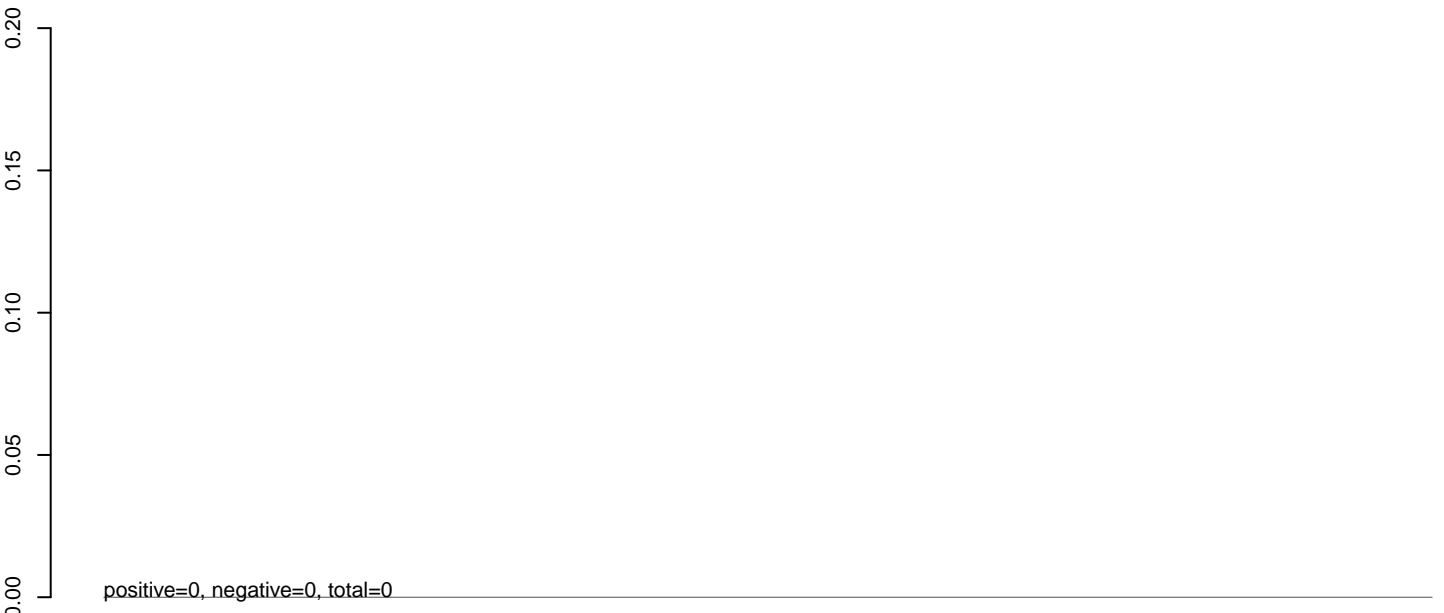
200 400 600 800 1000

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

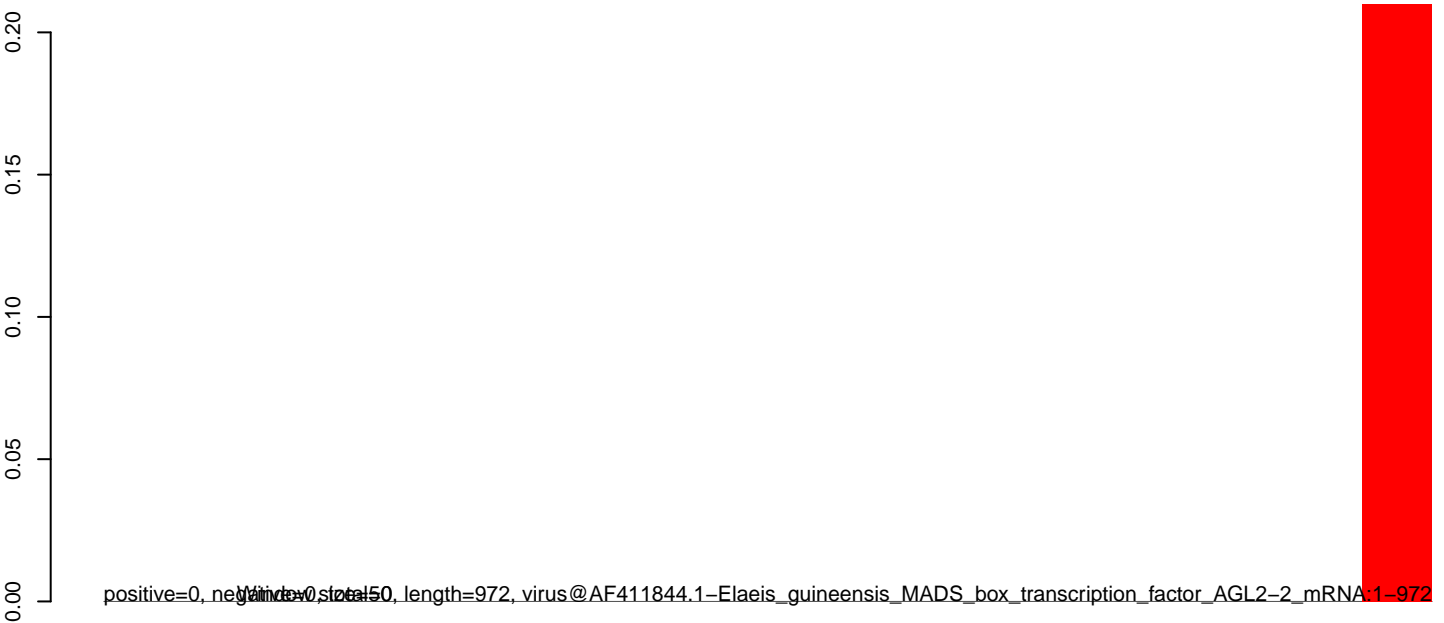
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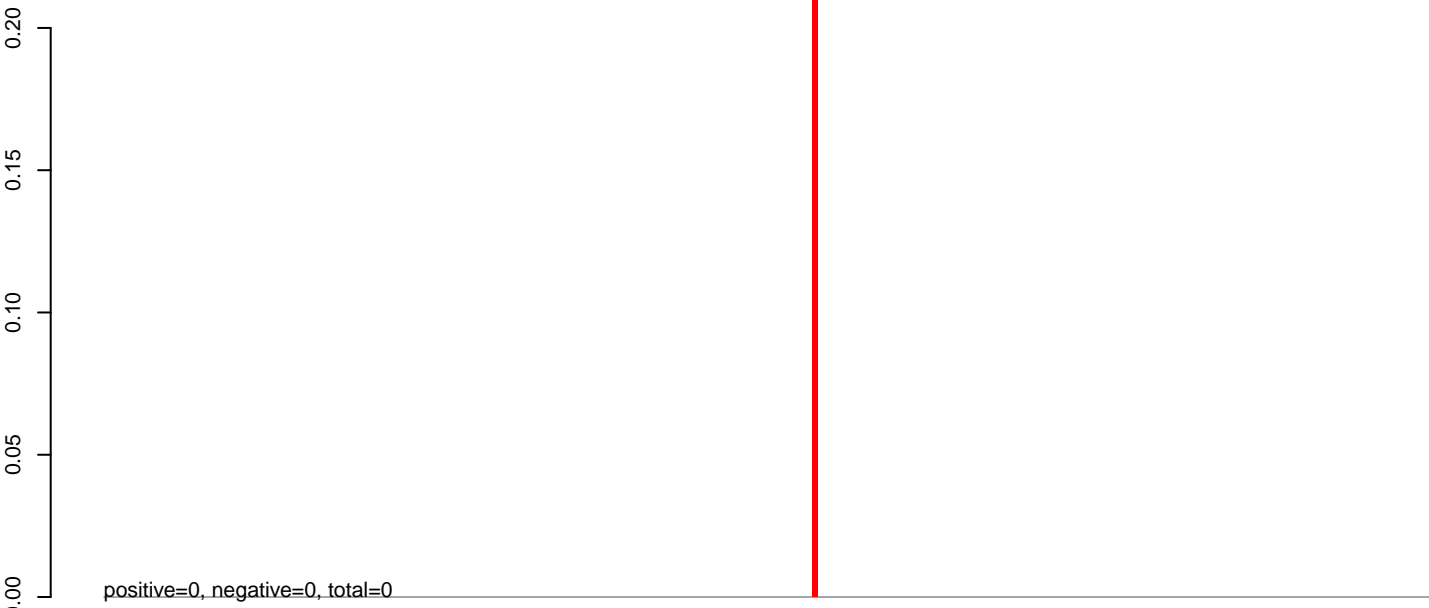
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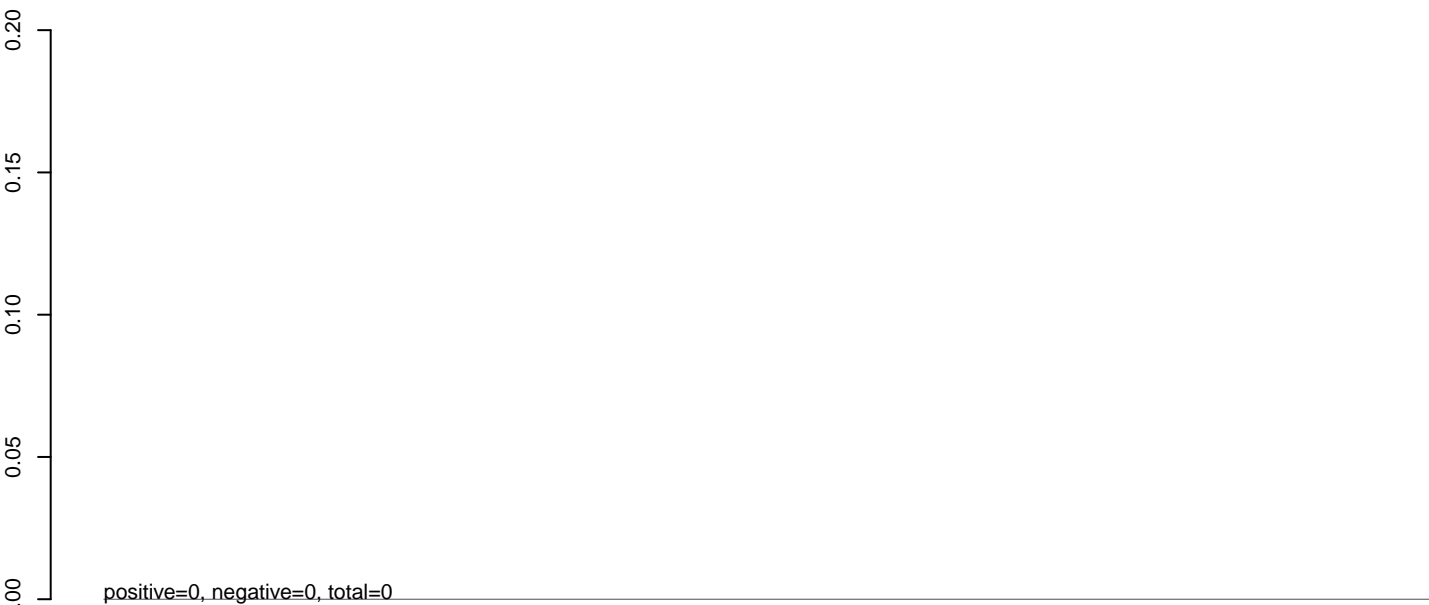
AeAeg_Aag2_DENV4.rep



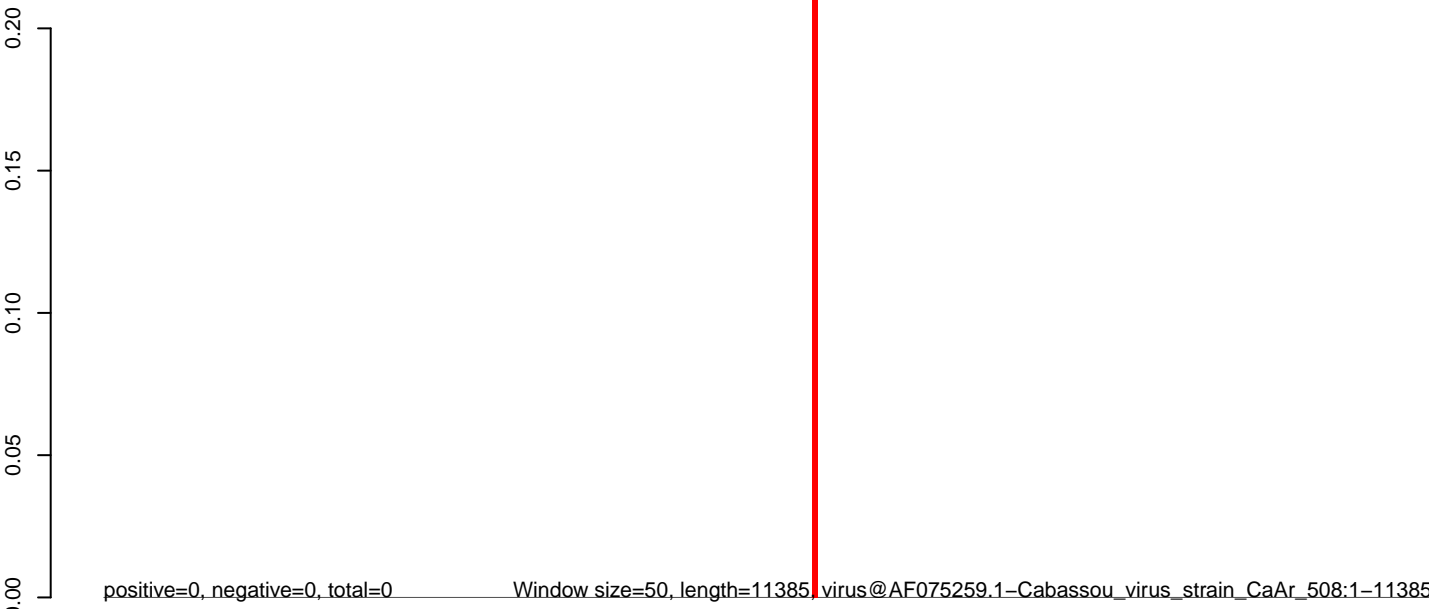
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



0 2000 4000 6000 8000 10000

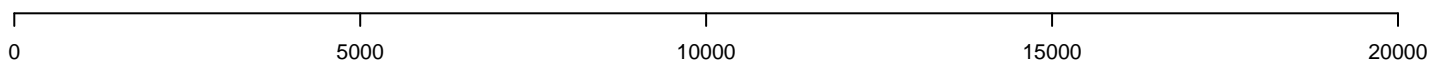
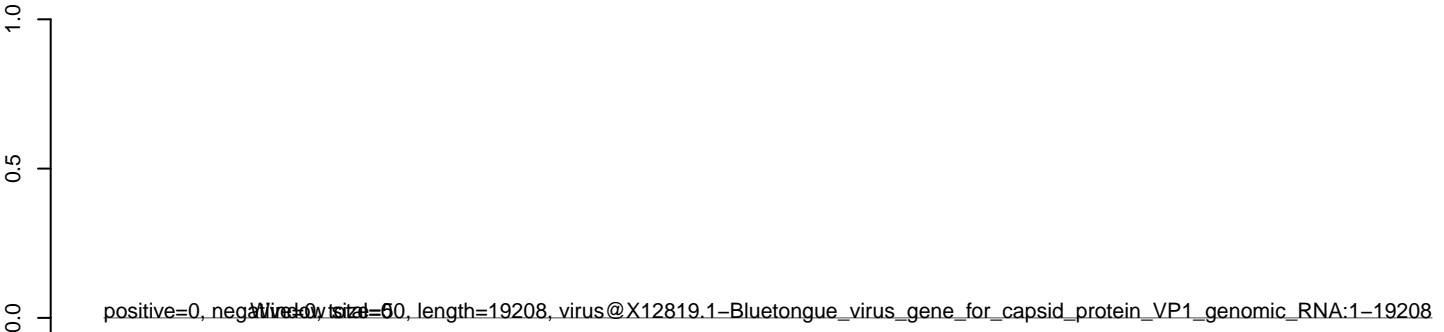
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



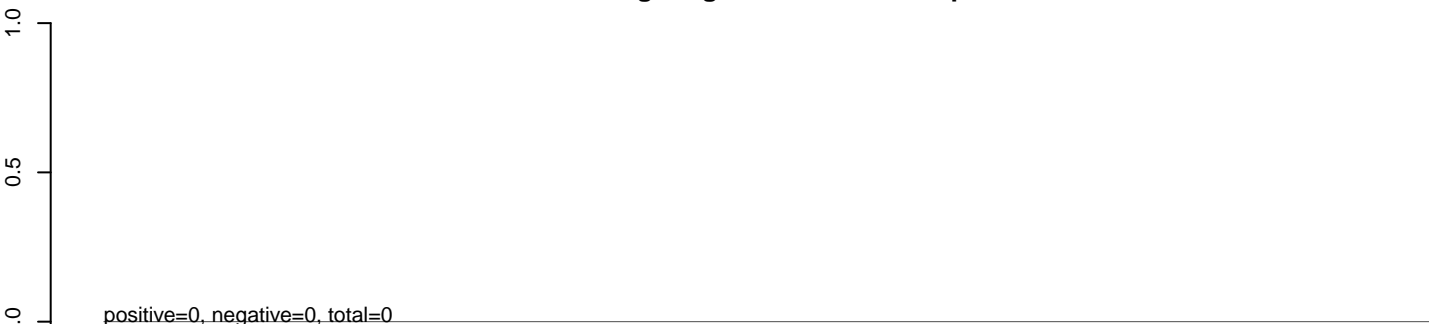
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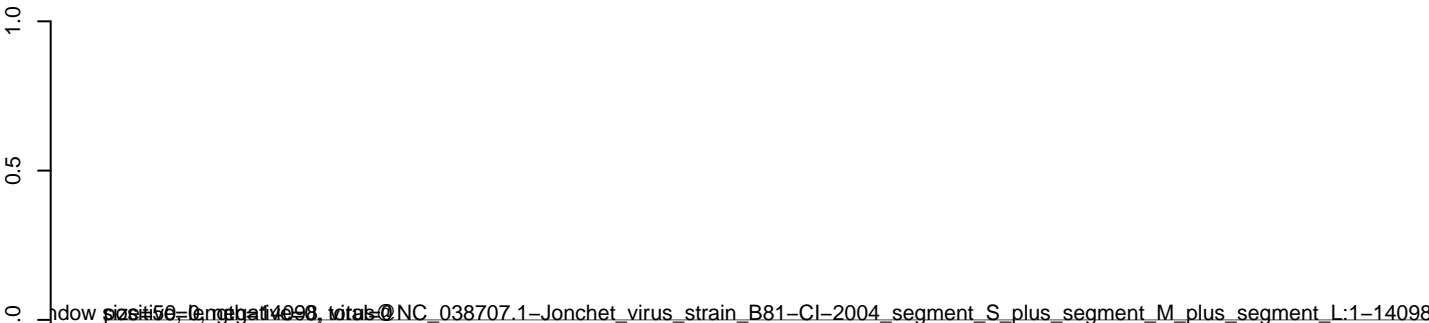
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

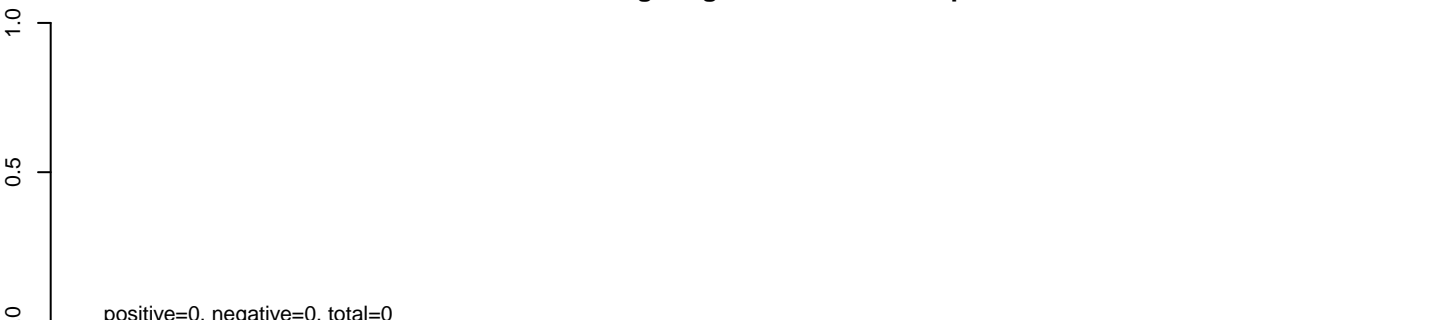


AeAeg_Aag2_DENV4.rep

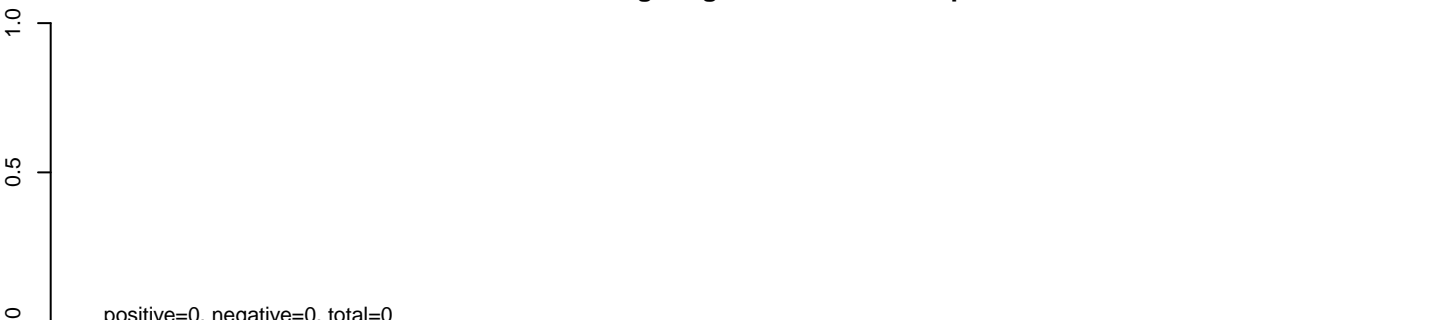


0 2000 4000 6000 8000 10000 12000 14000

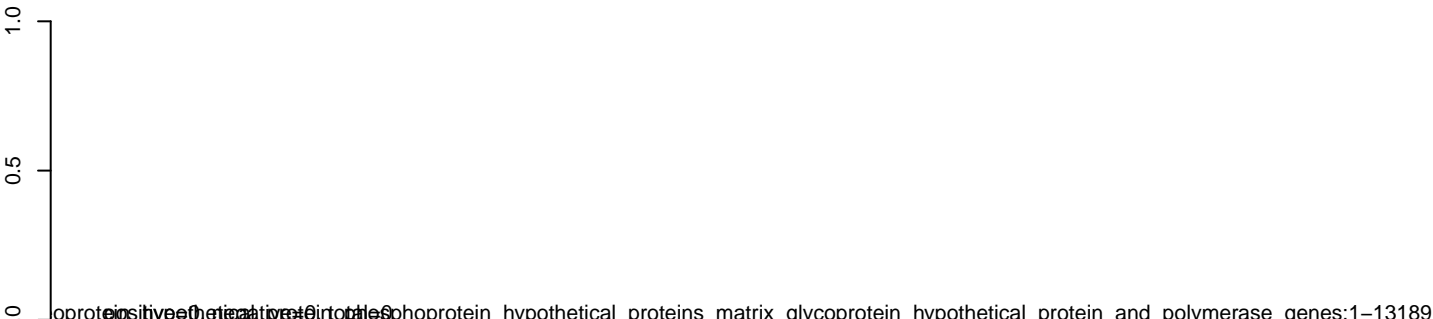
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



0 2000 4000 6000 8000 10000 12000

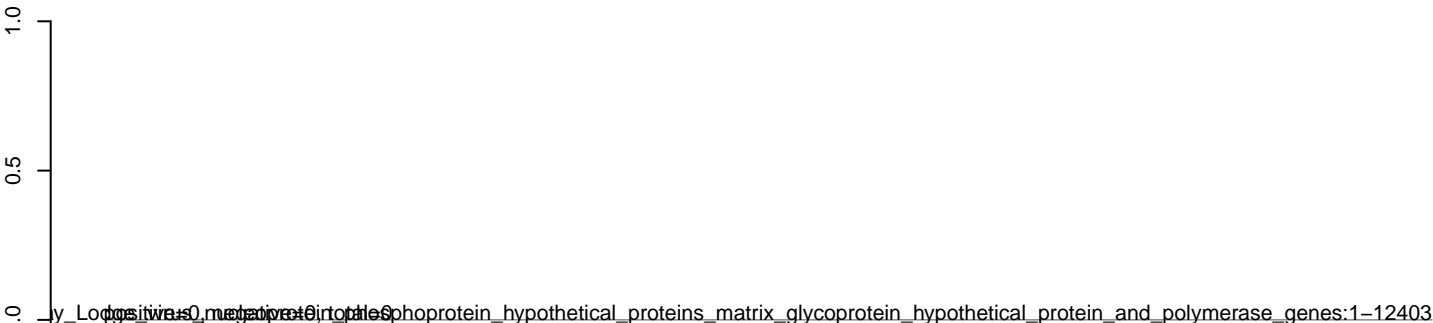
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep

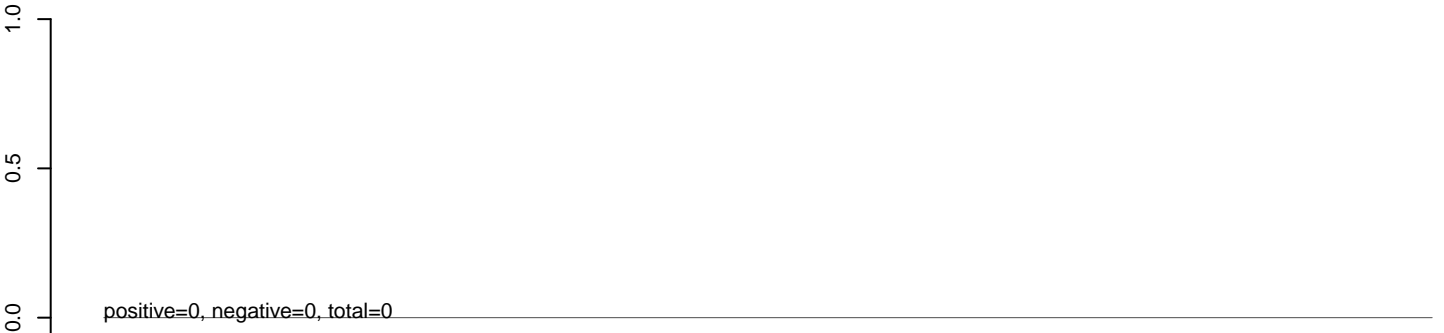


0 2000 4000 6000 8000 10000 12000

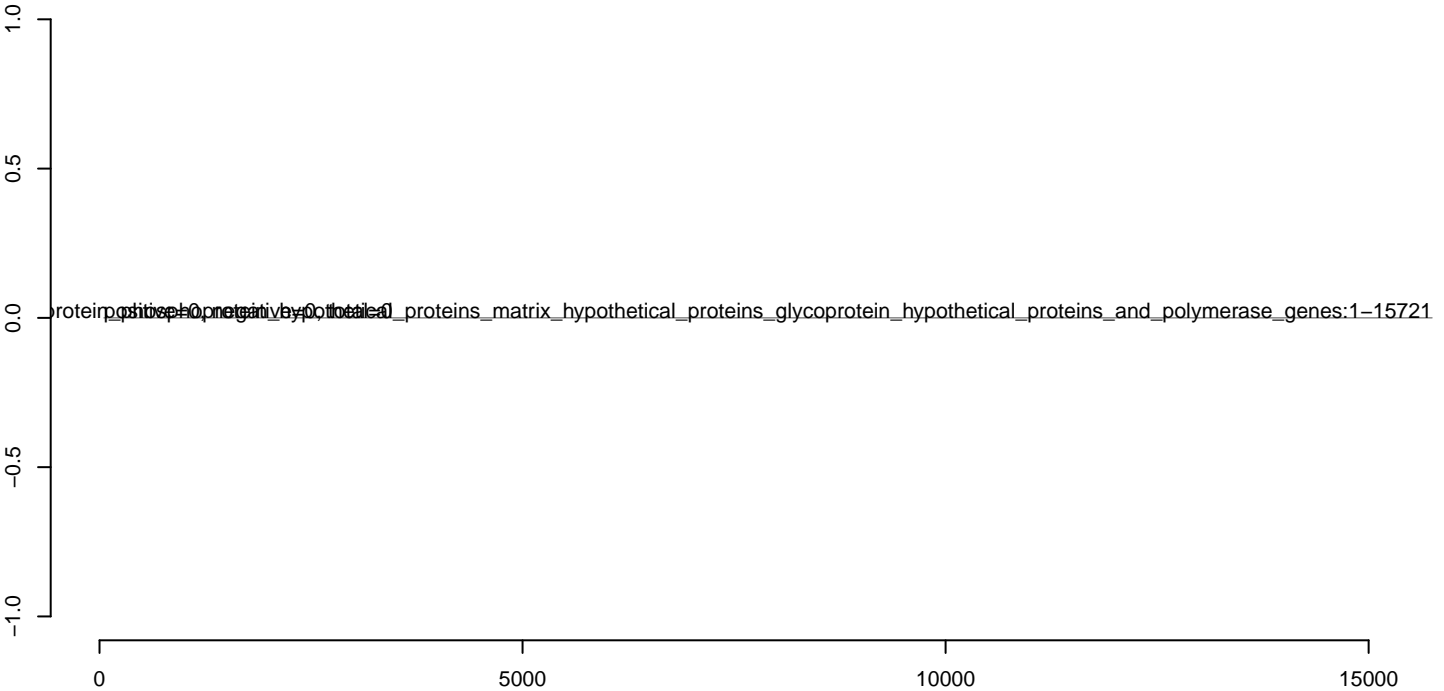
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



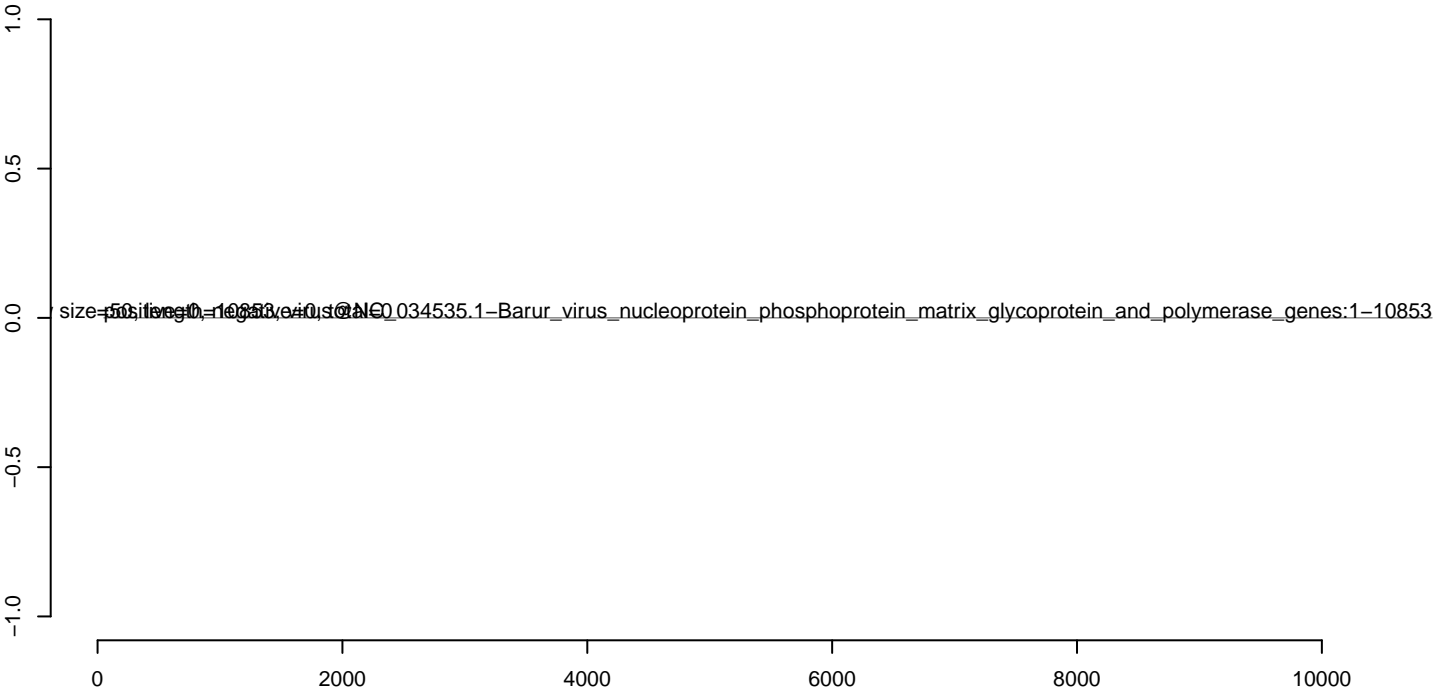
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



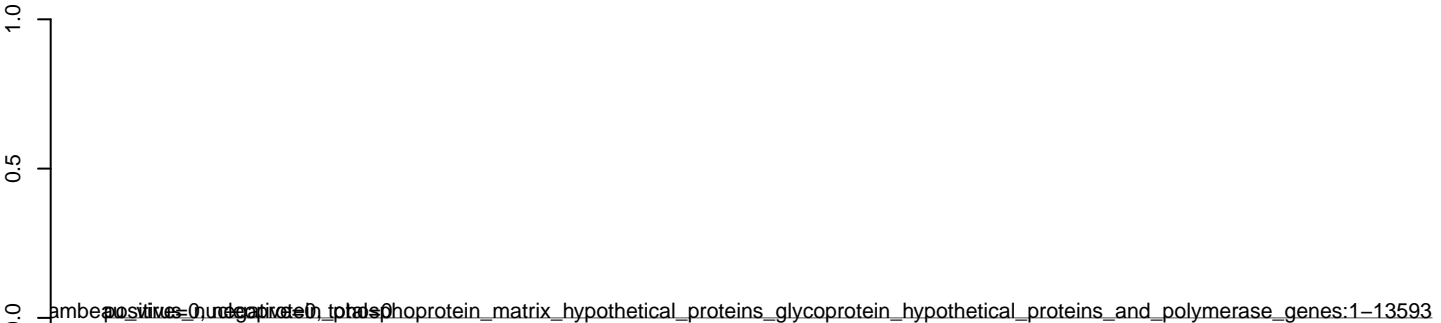
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AeAeg_Aag2_DENV4.24_35.rep

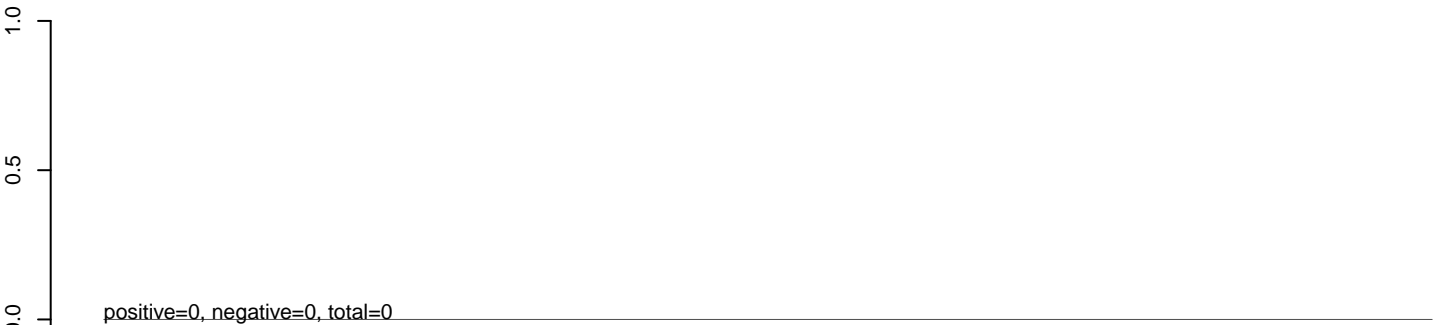


AeAeg_Aag2_DENV4.rep



0 2000 4000 6000 8000 10000 12000 14000

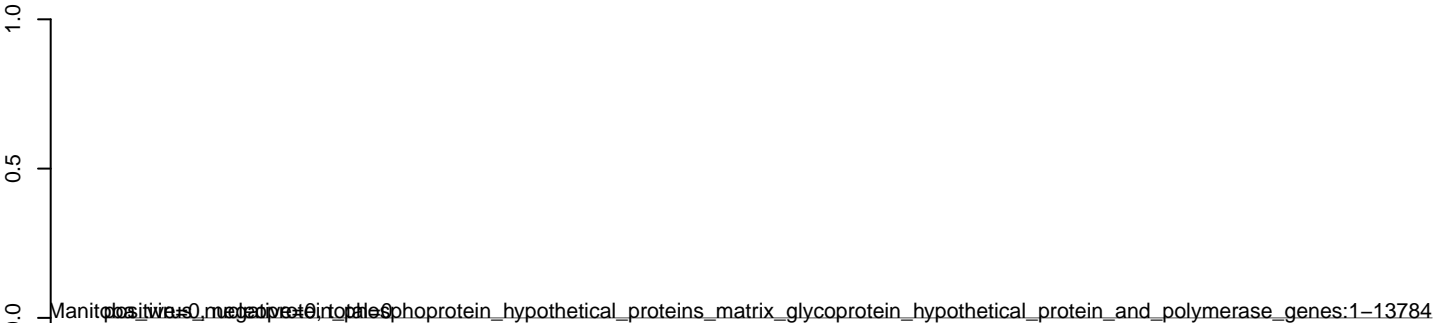
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

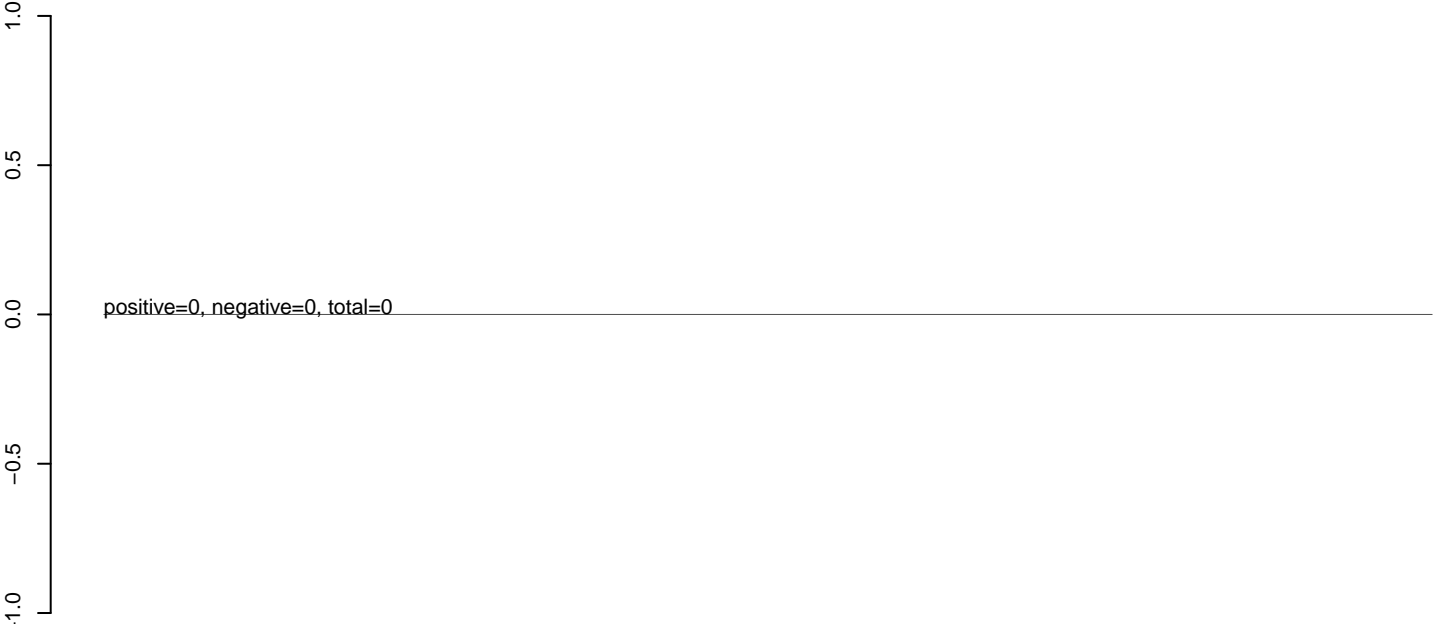


AeAeg_Aag2_DENV4.rep

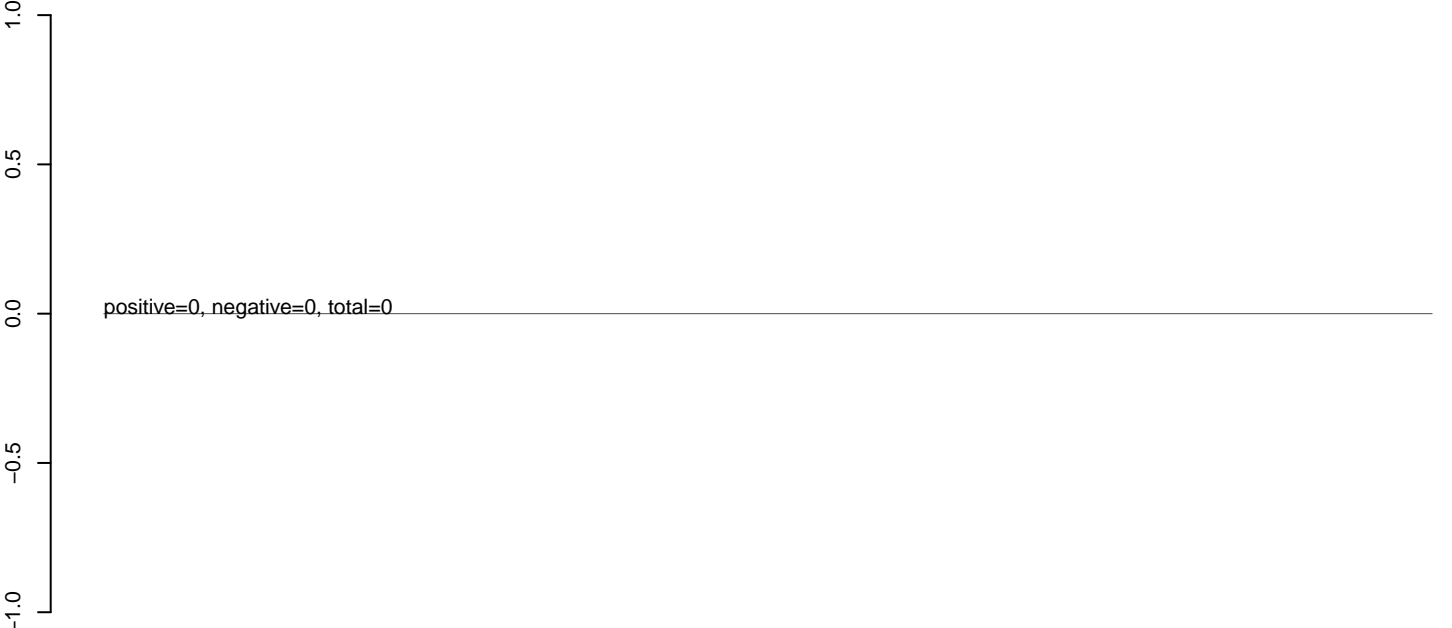


0 2000 4000 6000 8000 10000 12000 14000

AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep

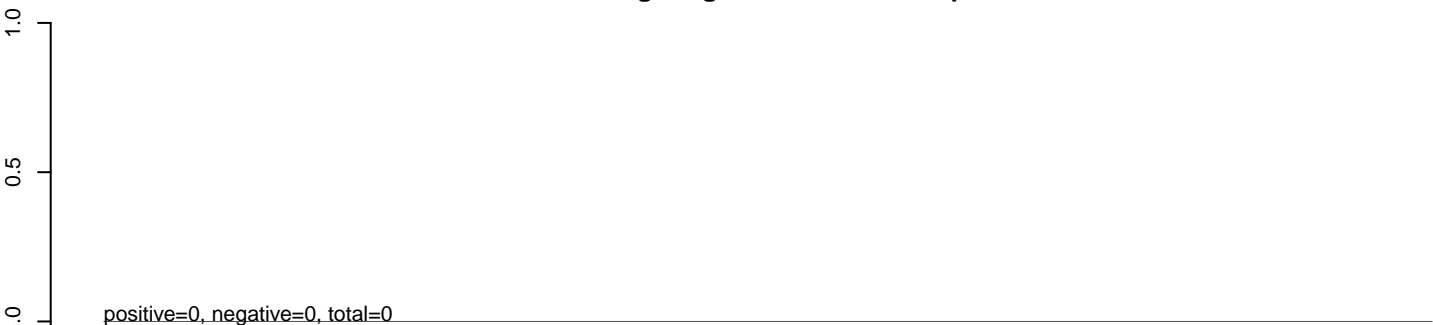


AeAeg_Aag2_DENV4.rep

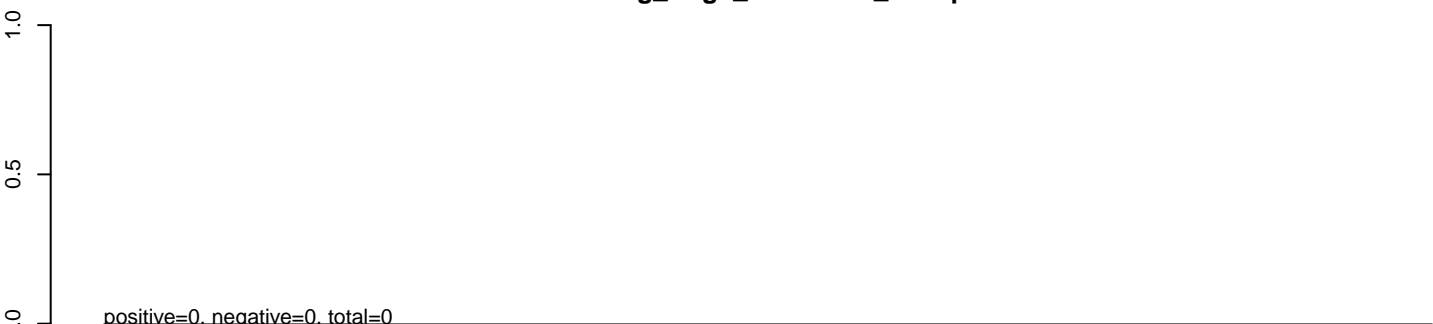


GenBank:AF146161.1, virus:G04582, negative:Matrix_6_nucleoprotein_phosphoprotein_matrix_glycoprotein_hypothetical_proteins_and_polymerase_genes:1-13294

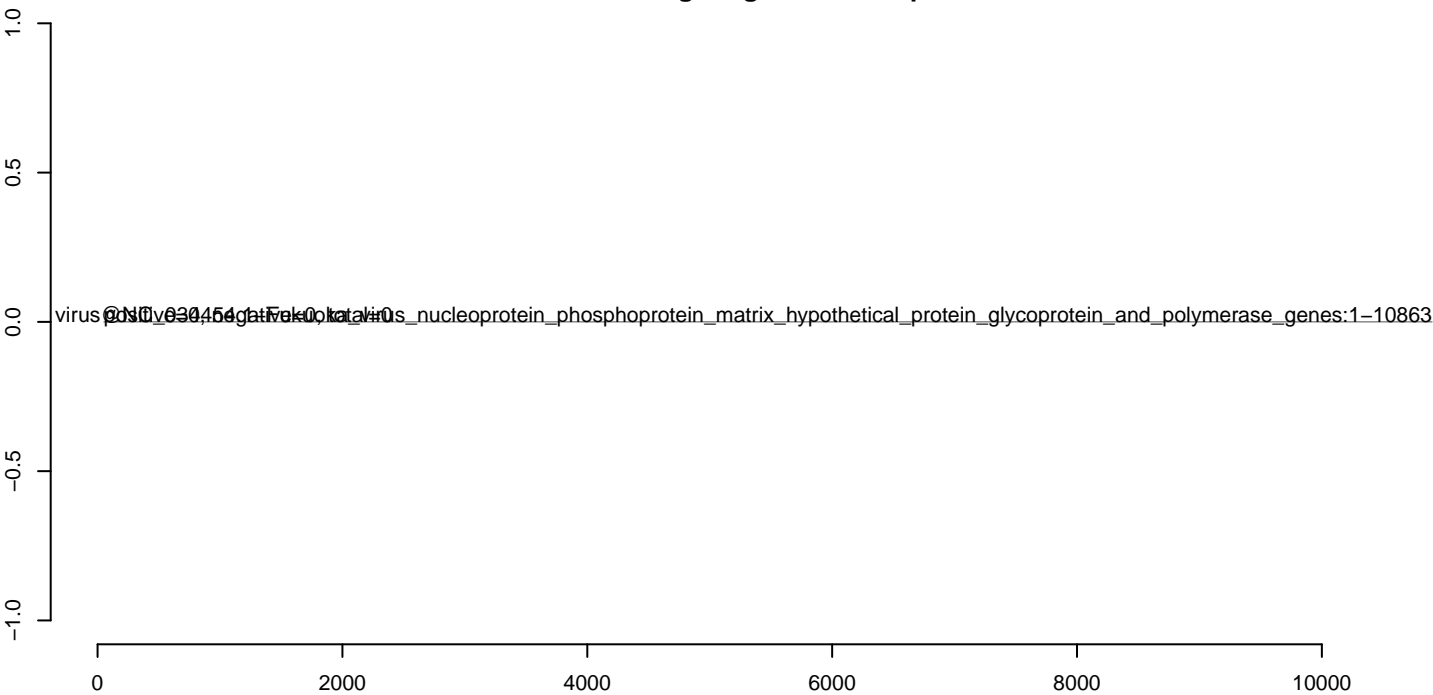
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



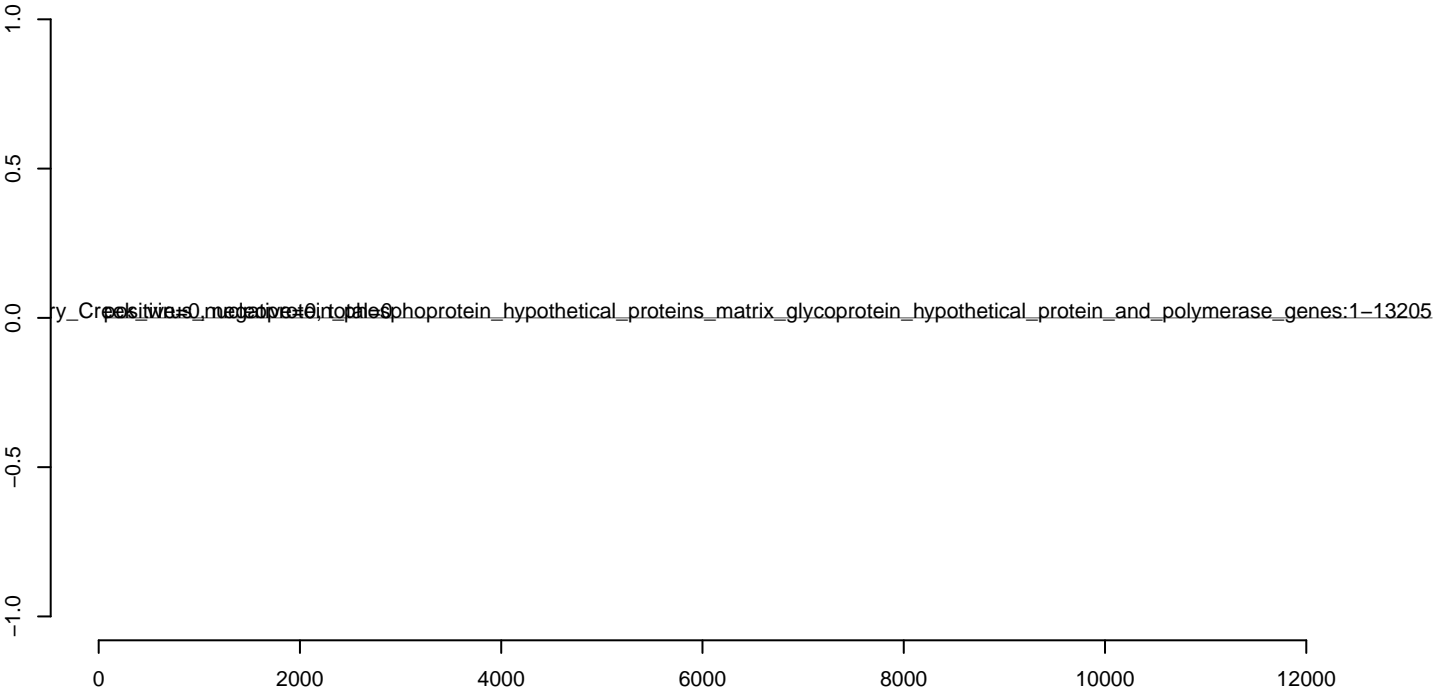
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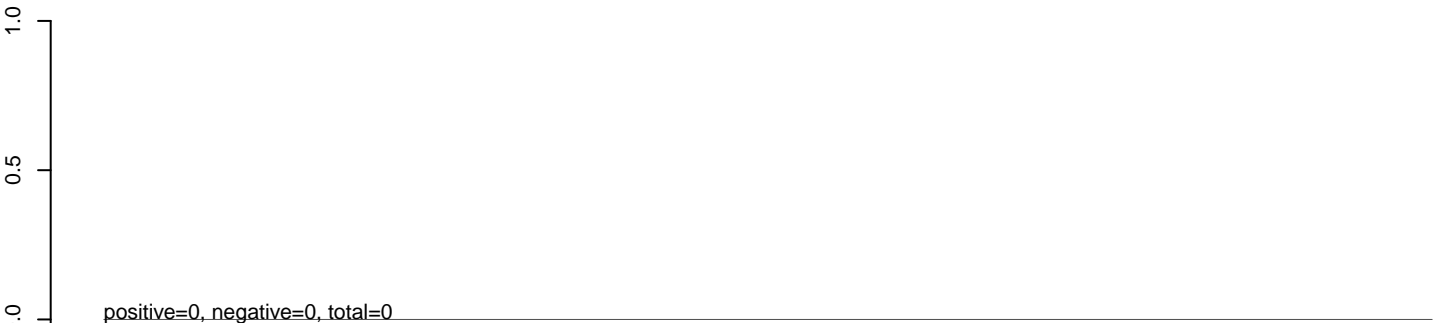
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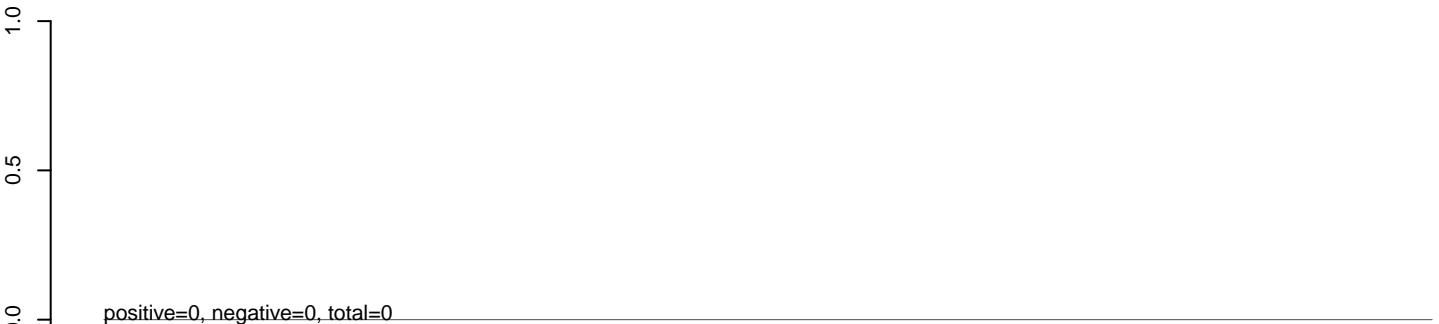
AeAeg_Aag2_DENV4.rep



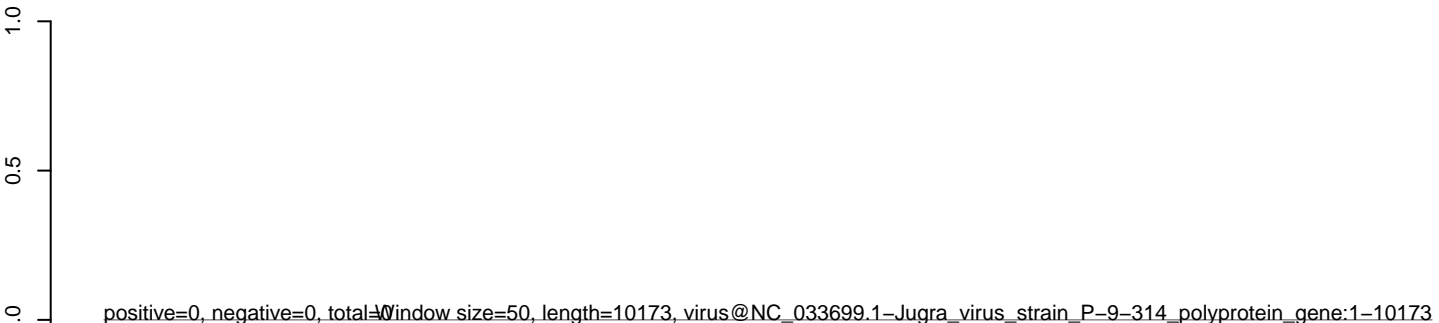
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



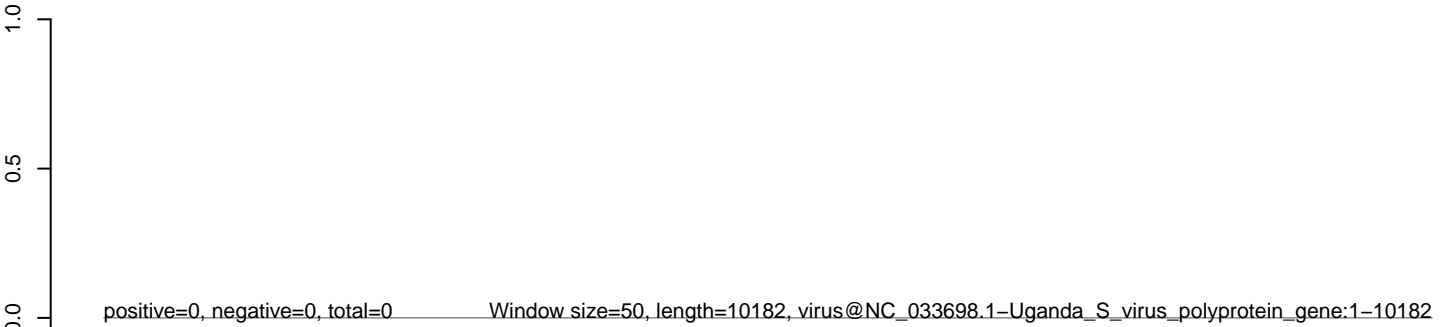
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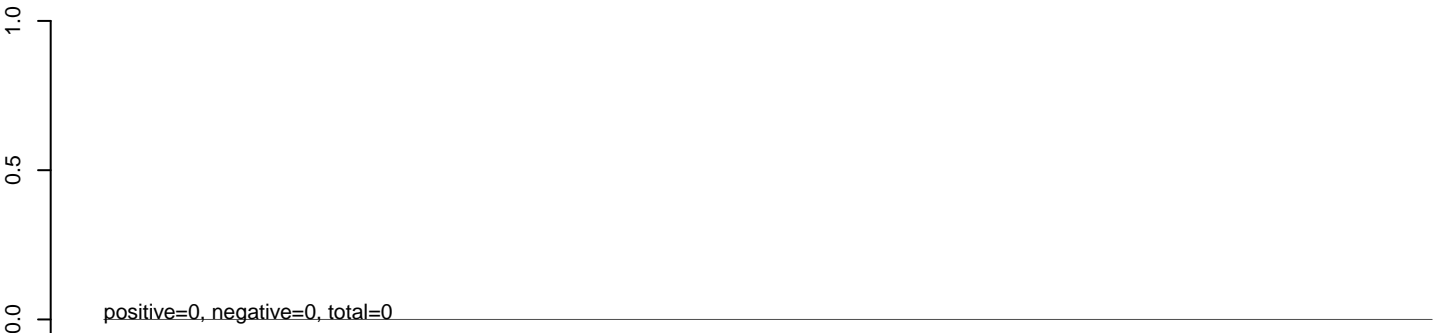
AeAeg_Aag2_DENV4.24_35.rep



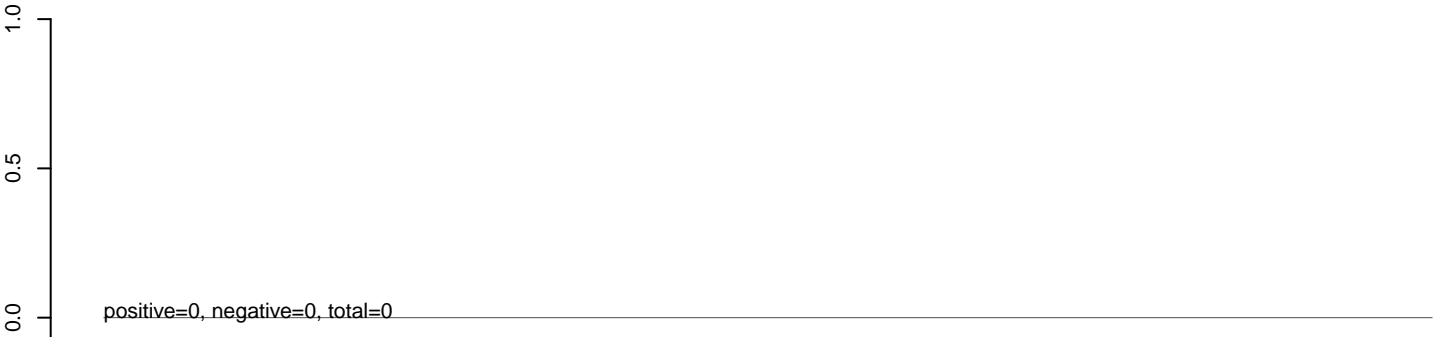
AeAeg_Aag2_DENV4.rep



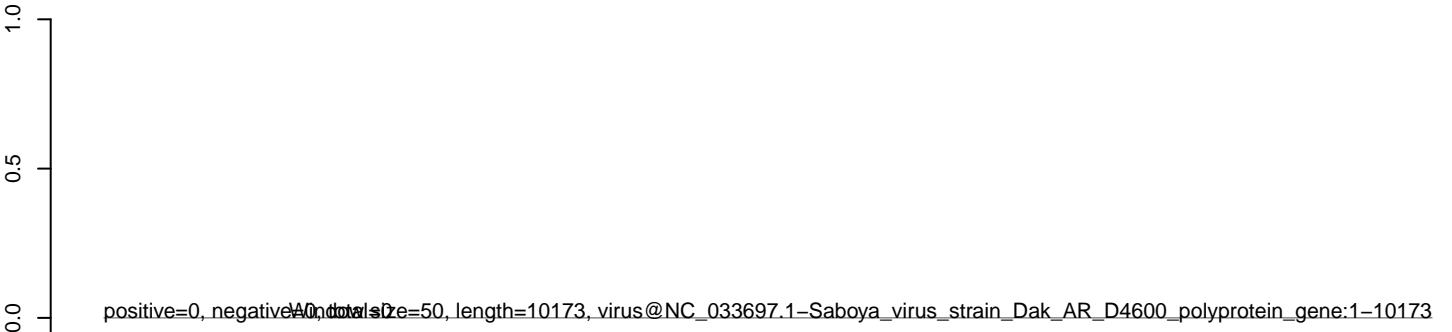
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



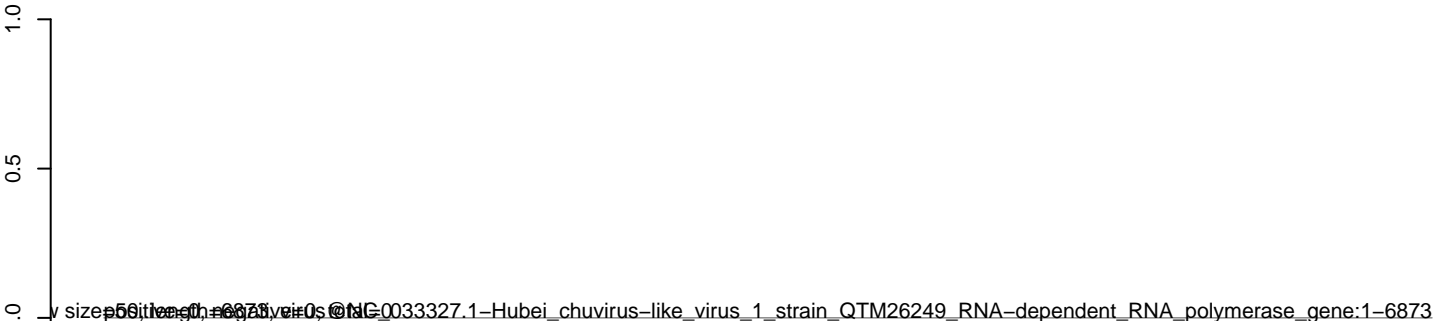
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



0 1000 2000 3000 4000 5000 6000 7000

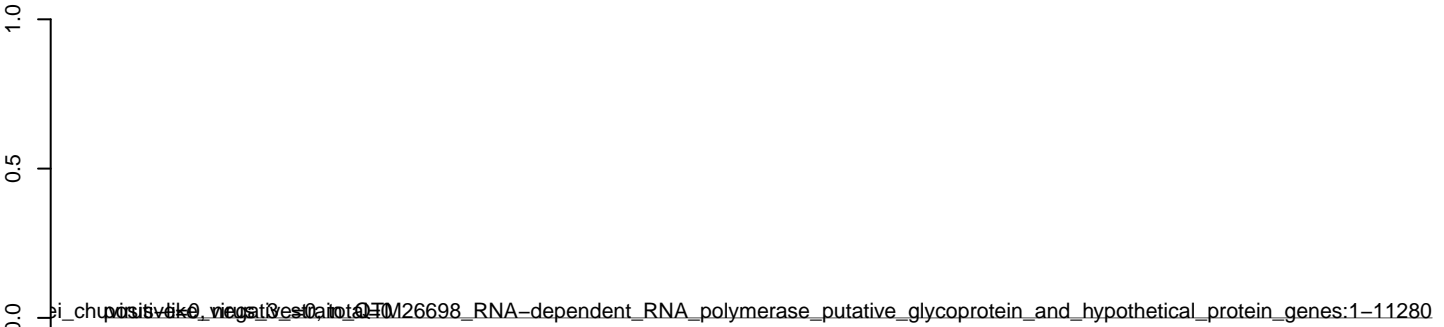
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep



0 2000 4000 6000 8000 10000

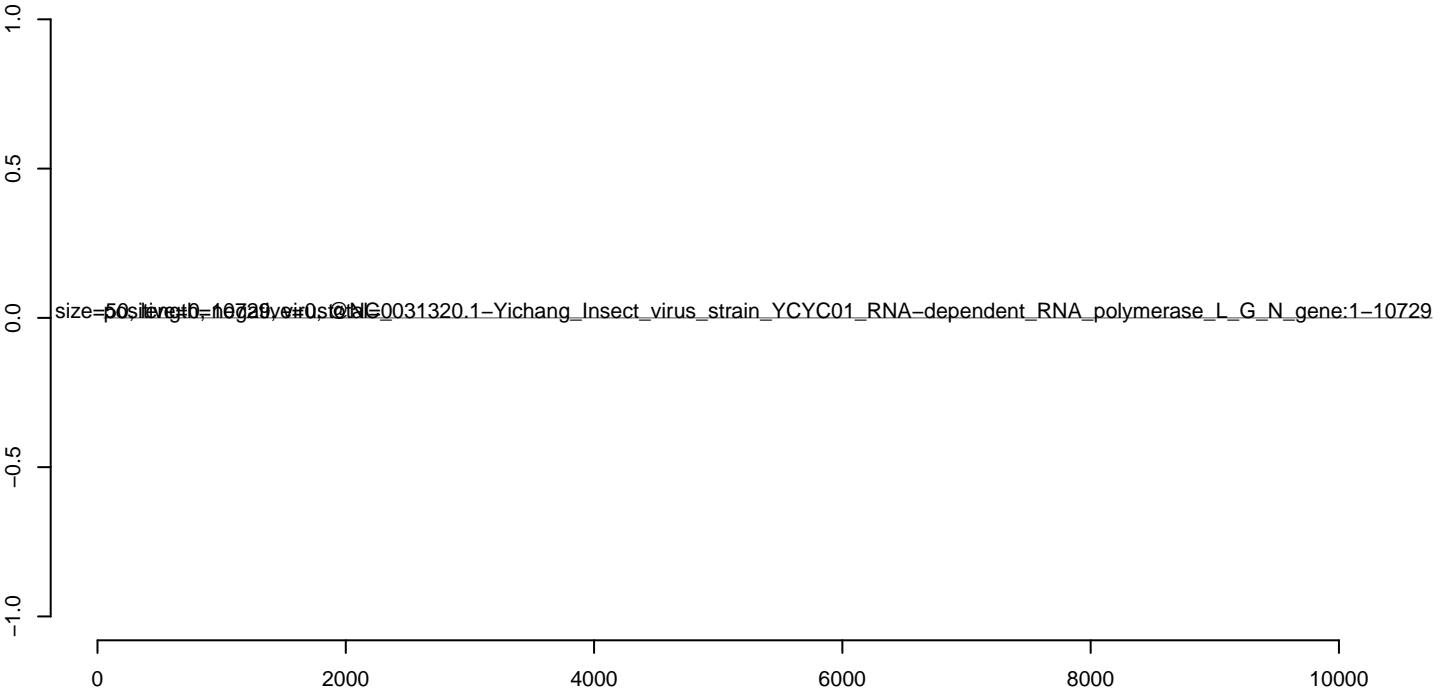
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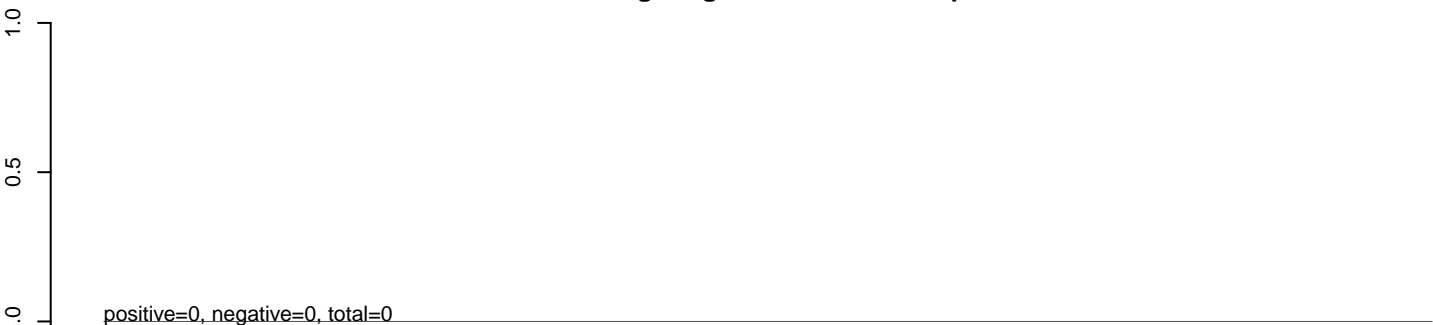
AeAeg_Aag2_DENV4.24_35.rep



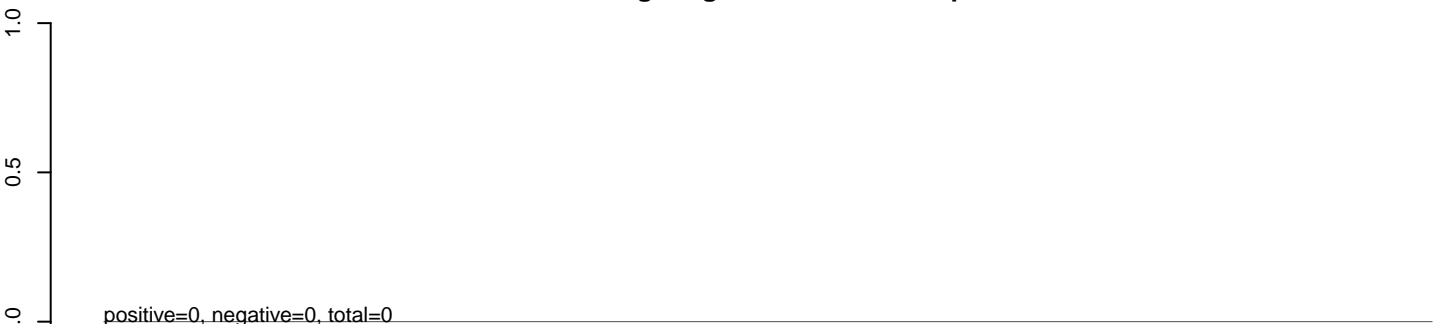
AeAeg_Aag2_DENV4.rep



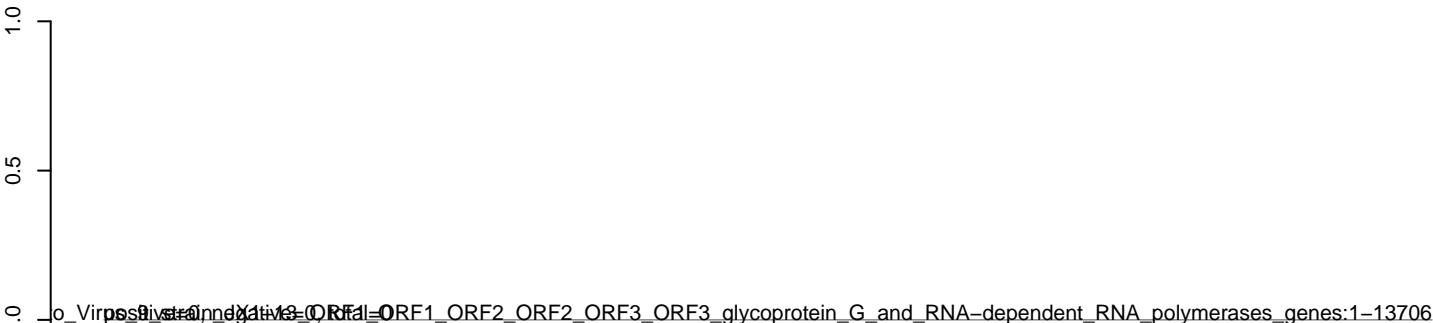
AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep

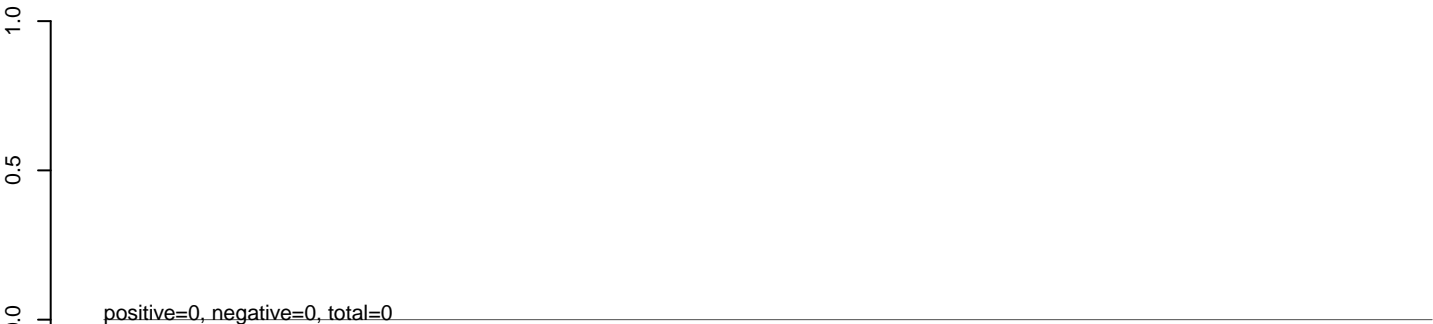


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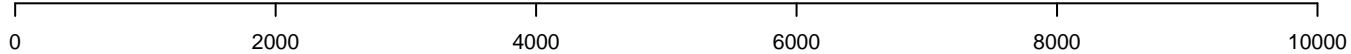
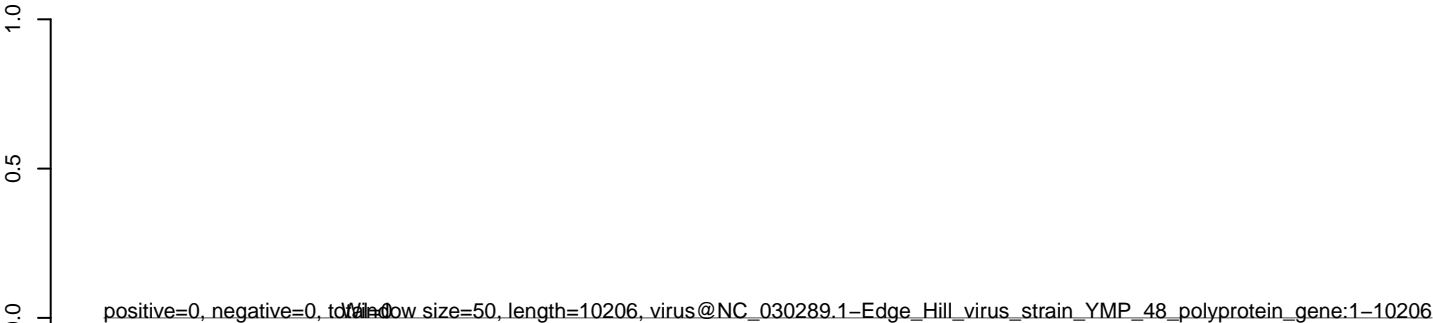
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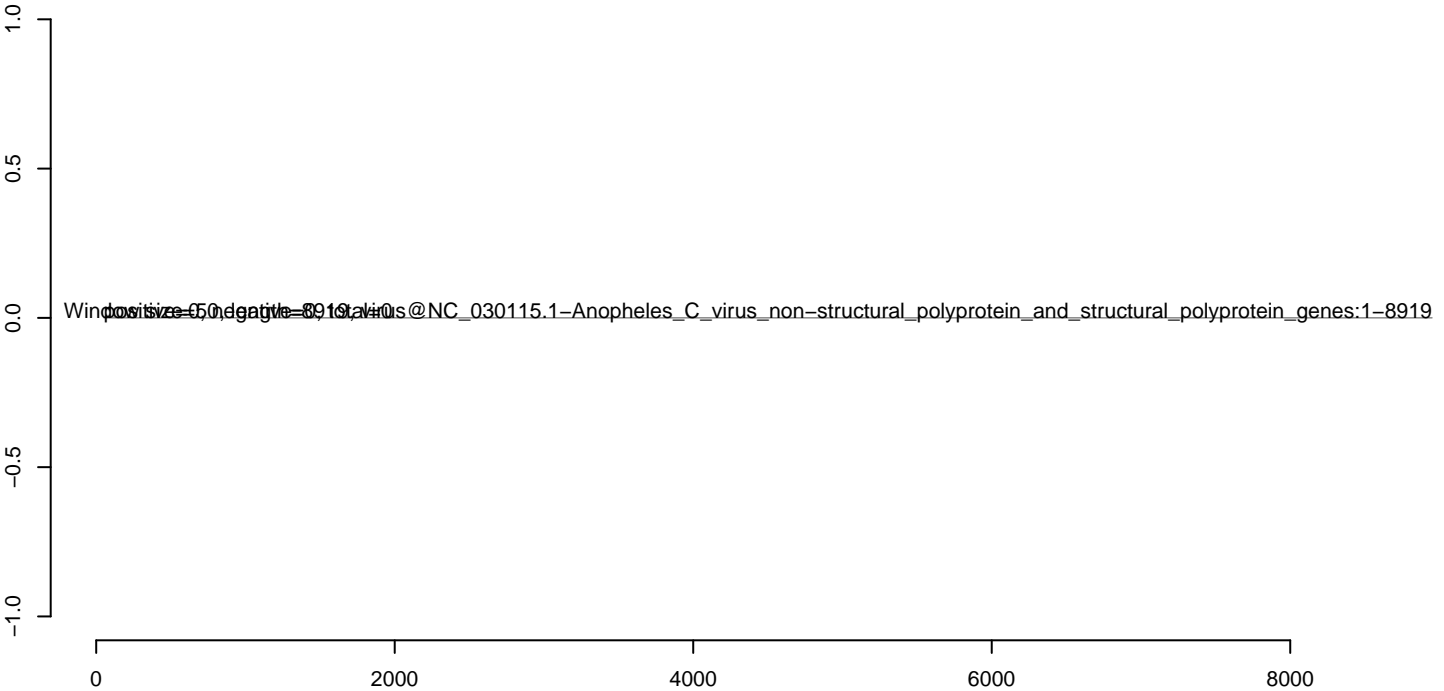
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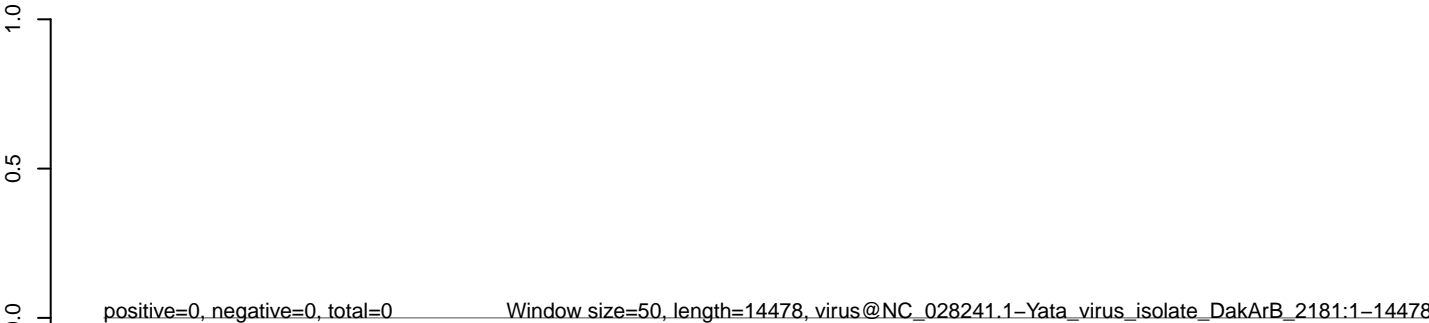
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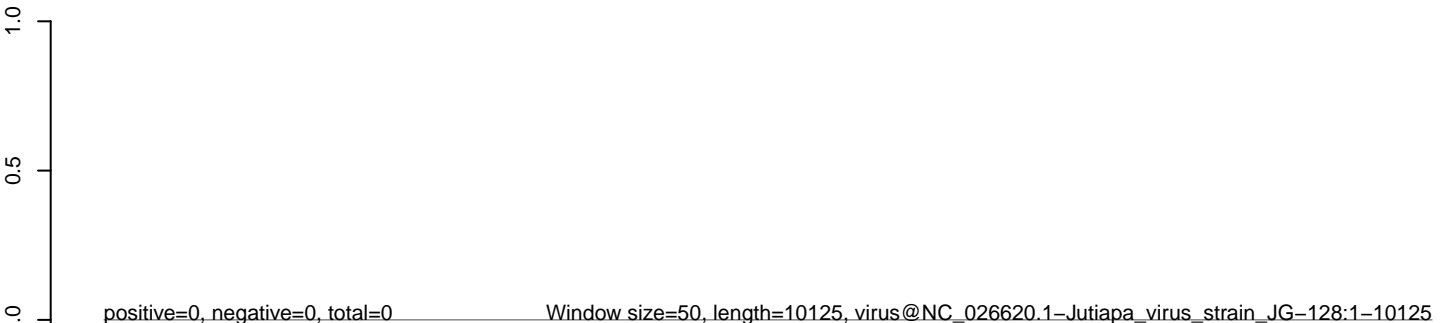
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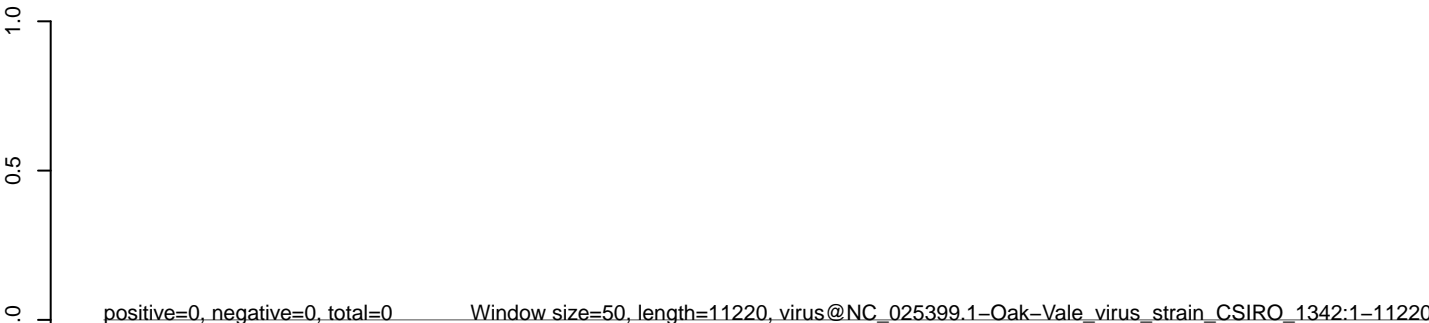
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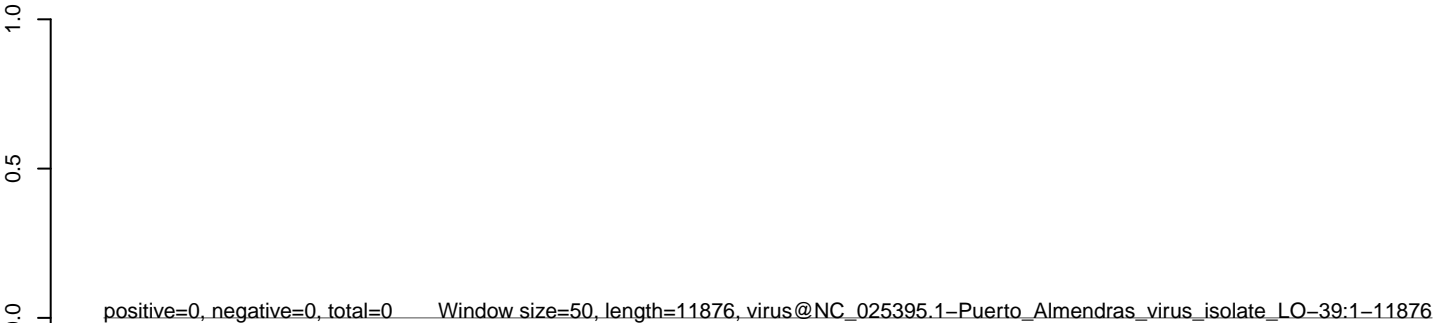
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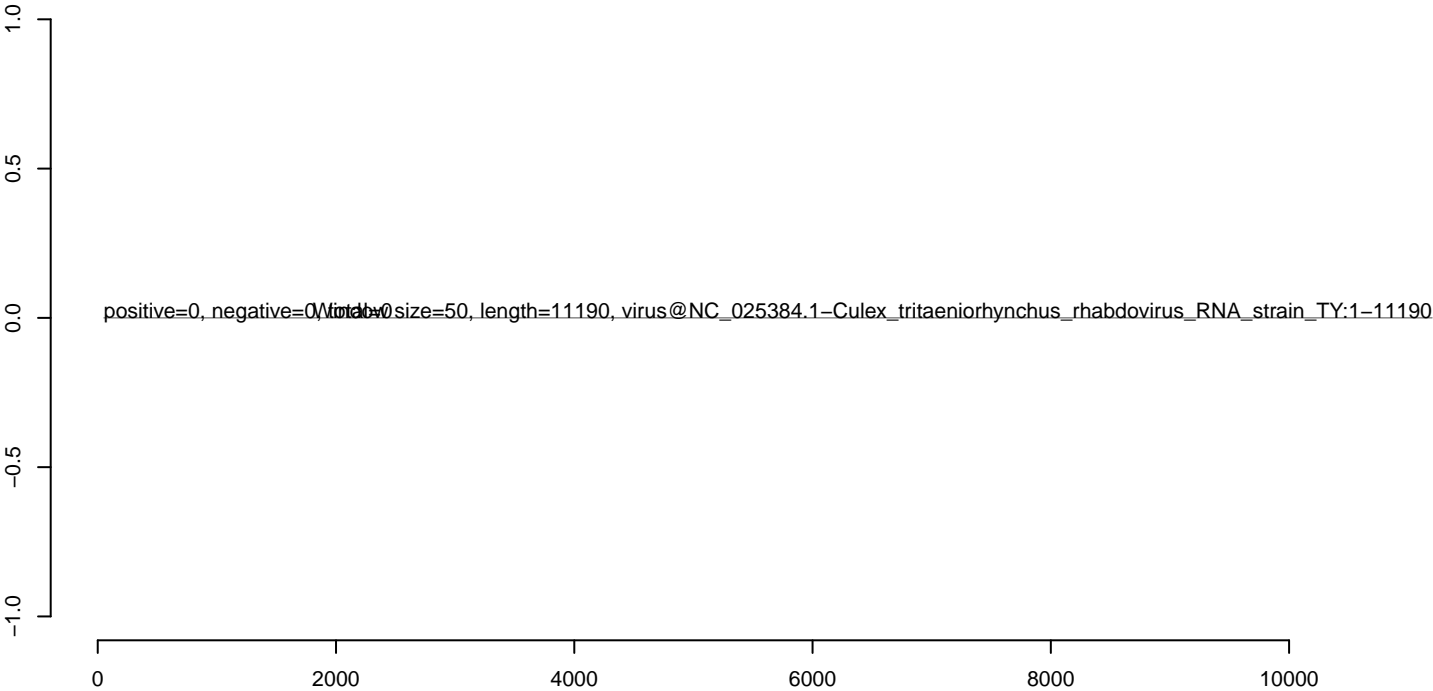
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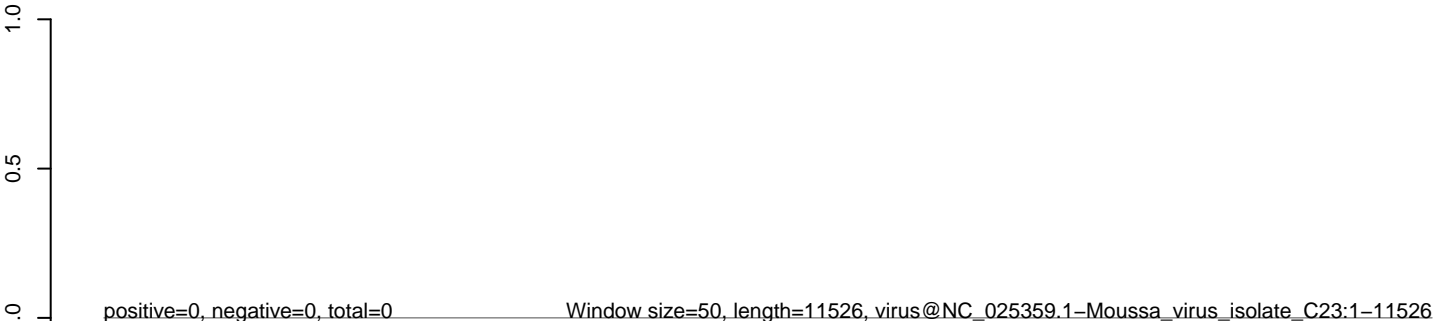
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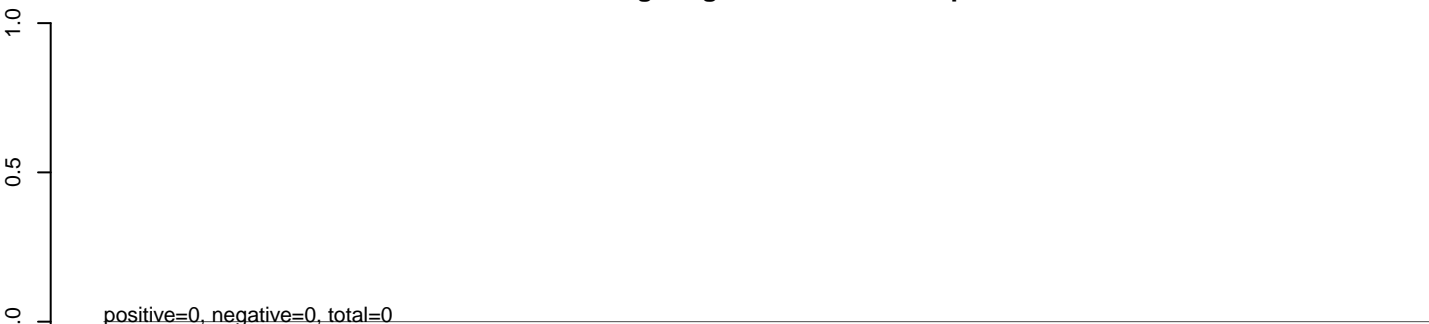


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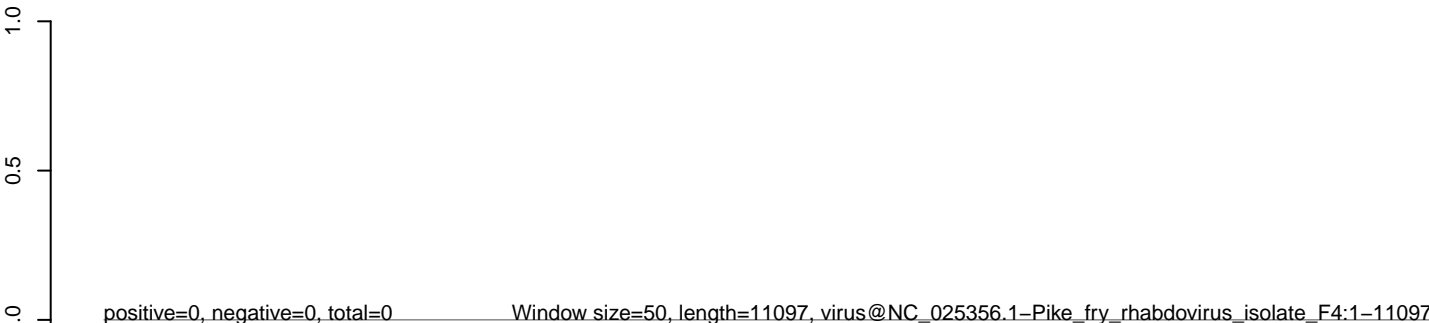
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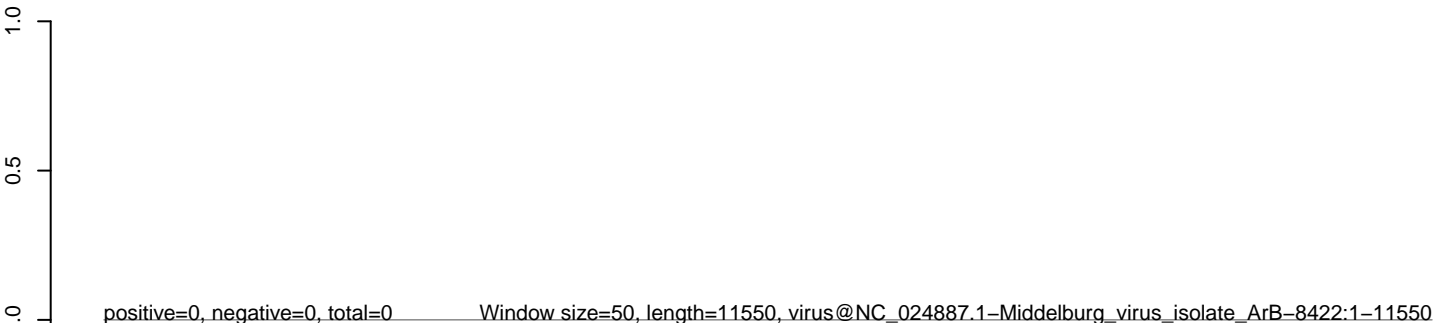
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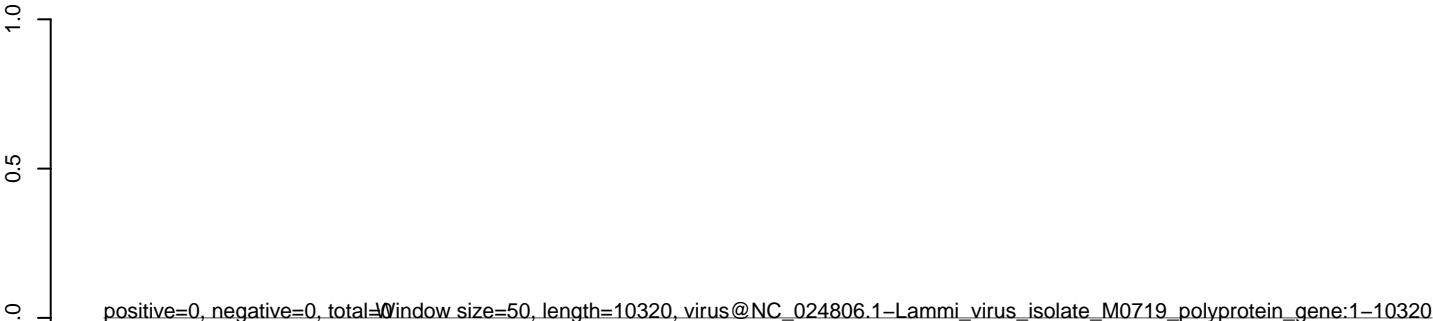
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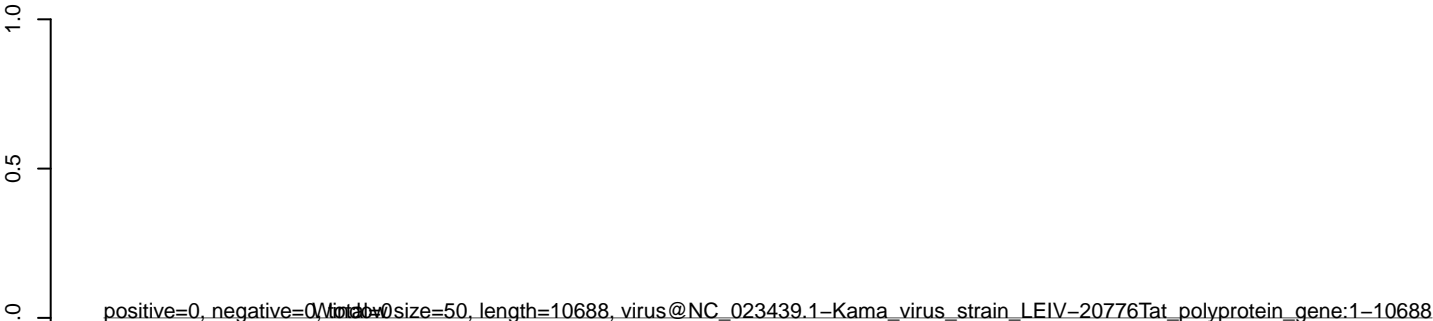
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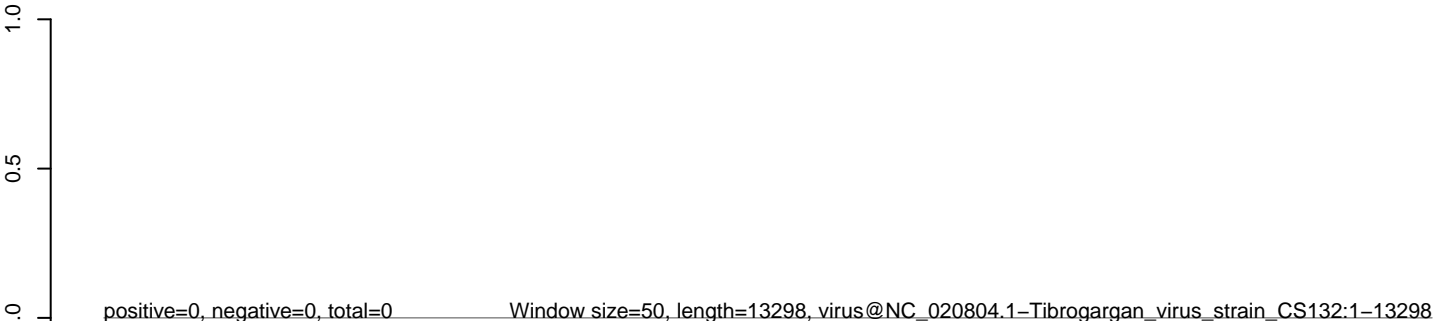
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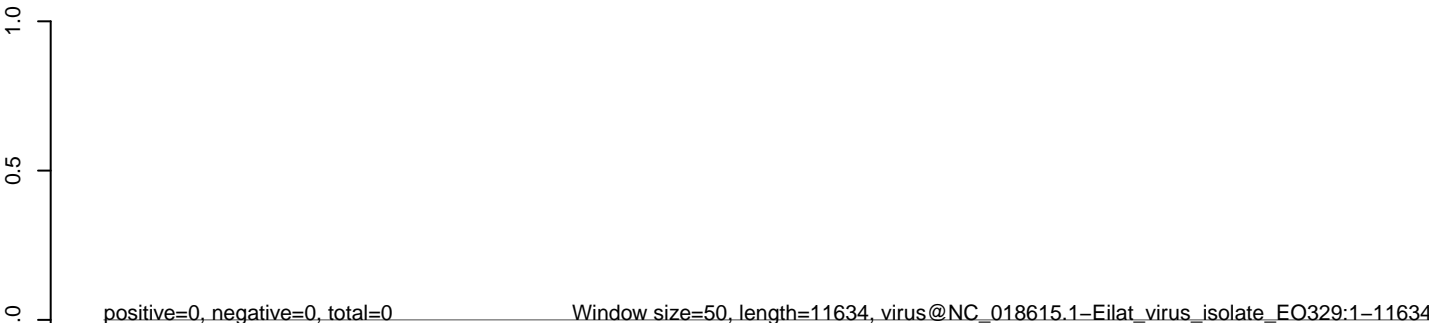
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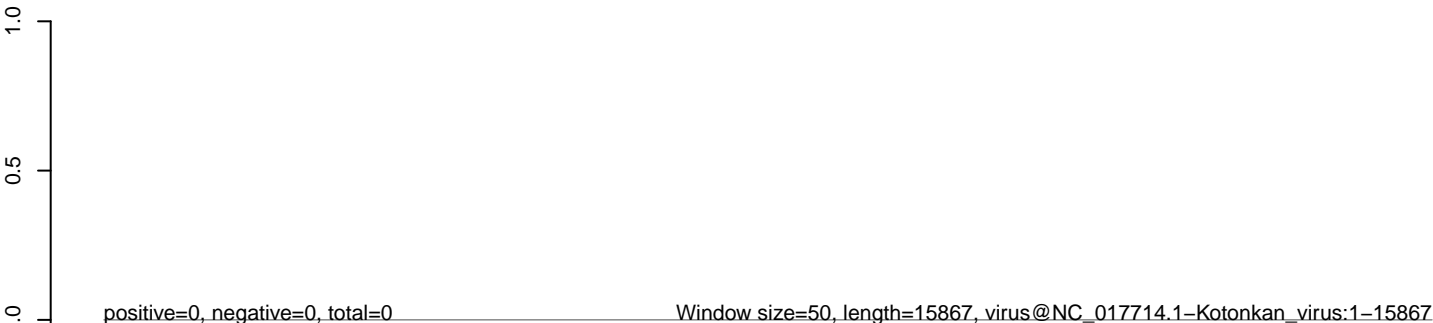
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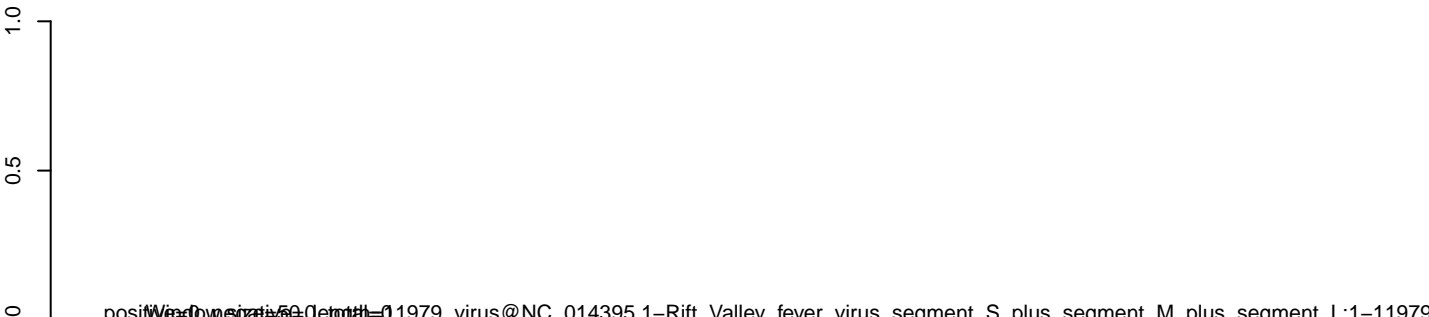
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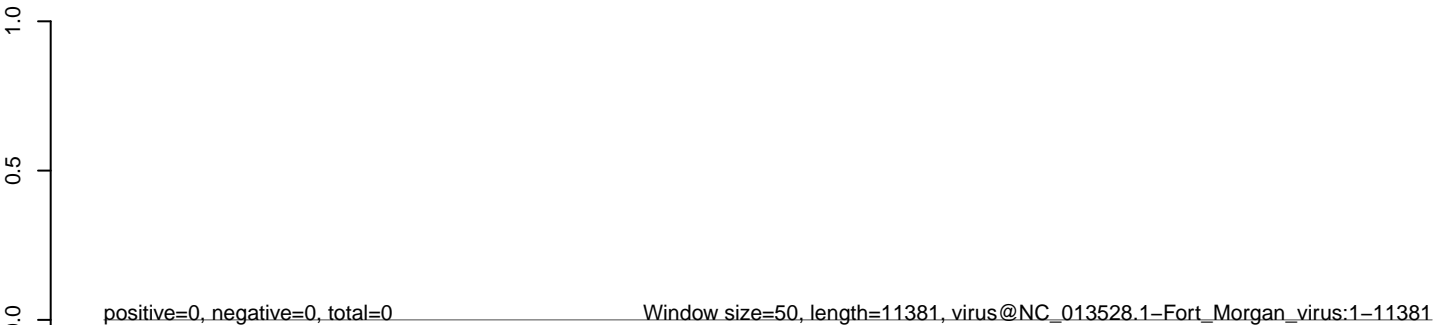
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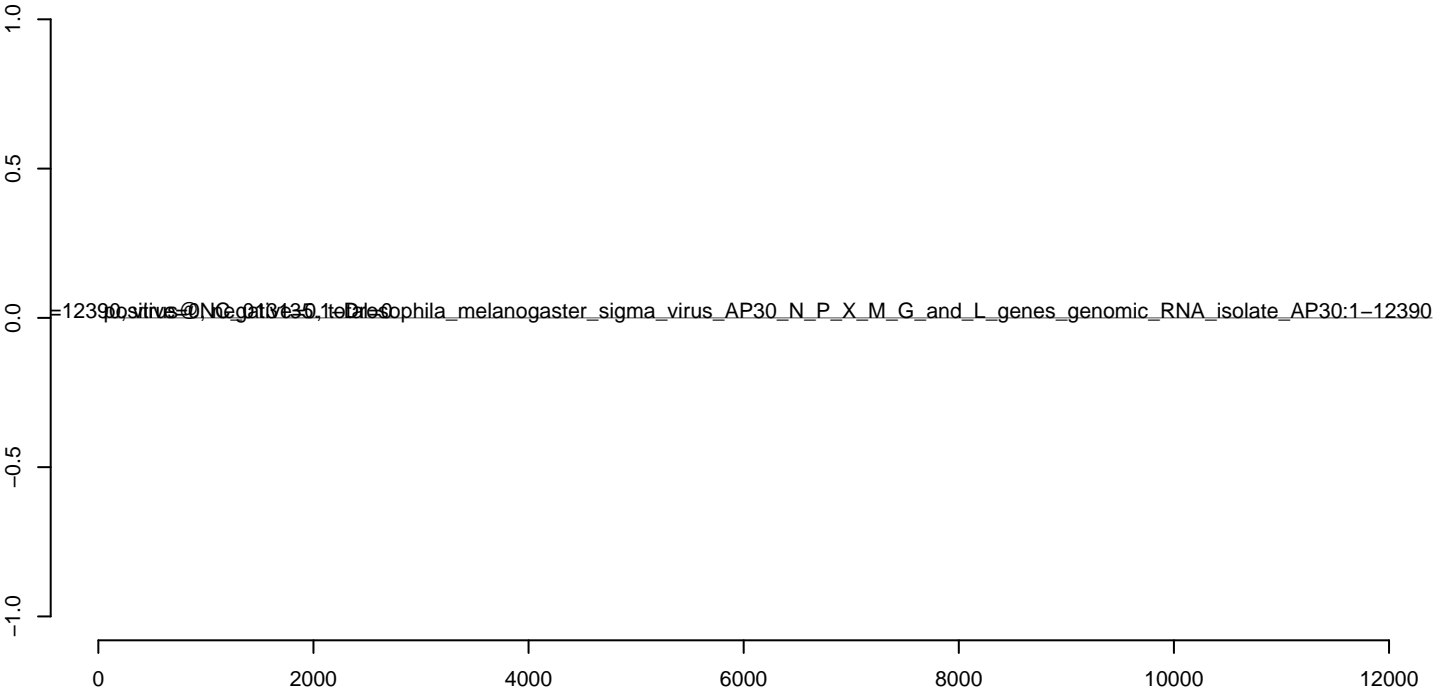
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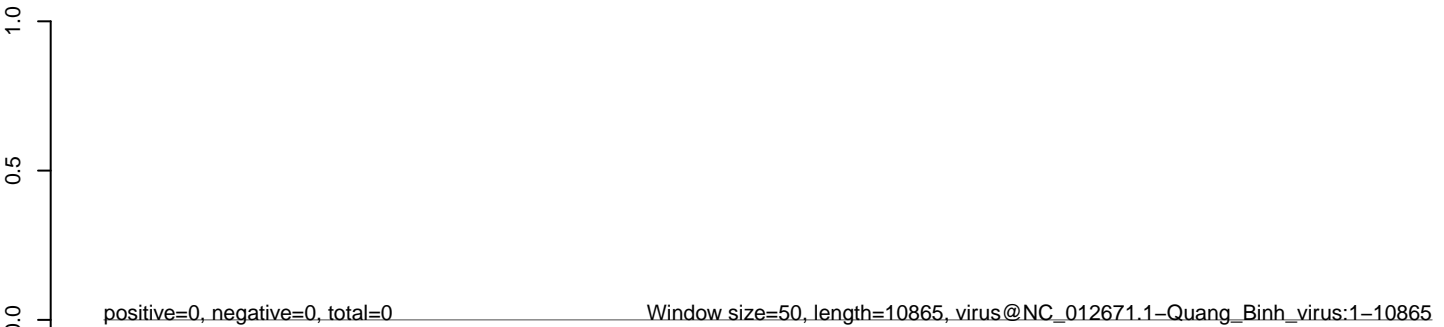
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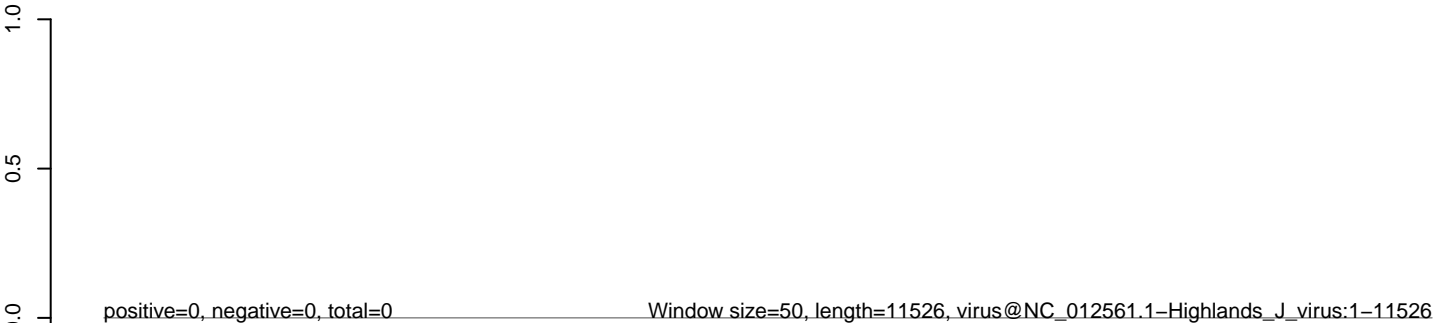
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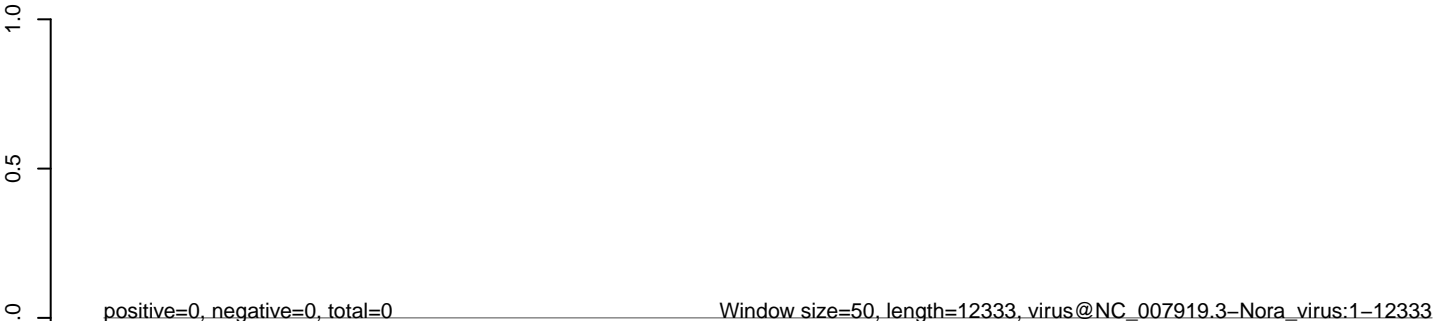
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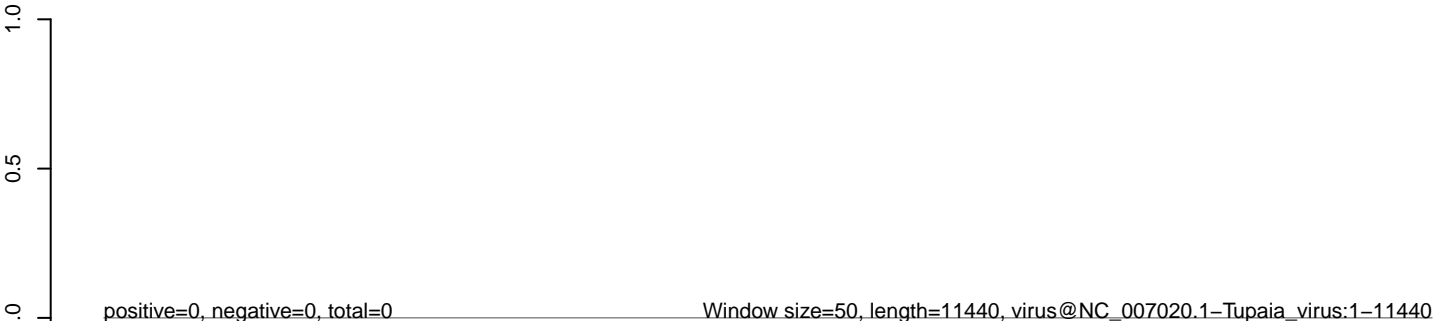
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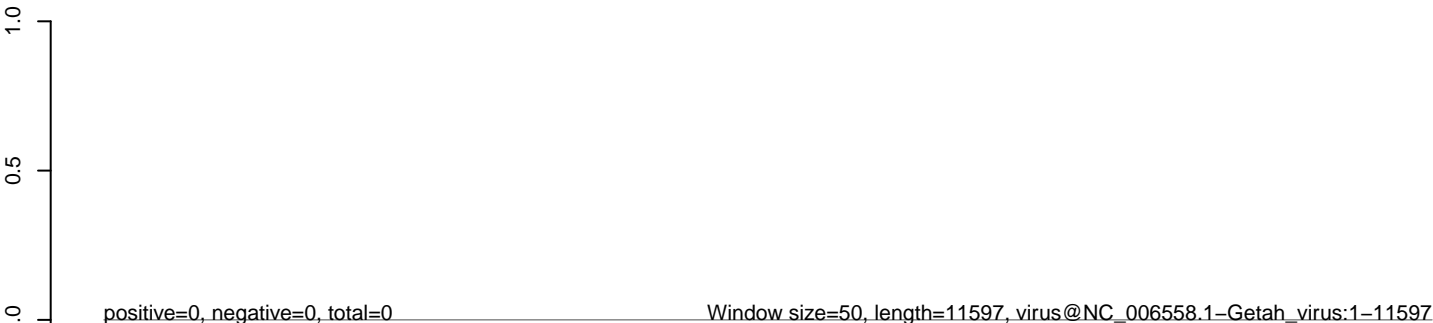
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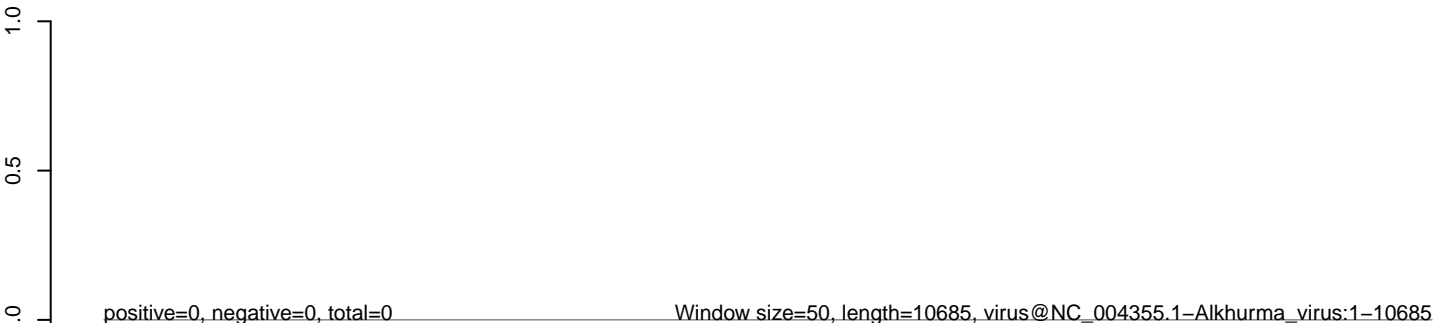
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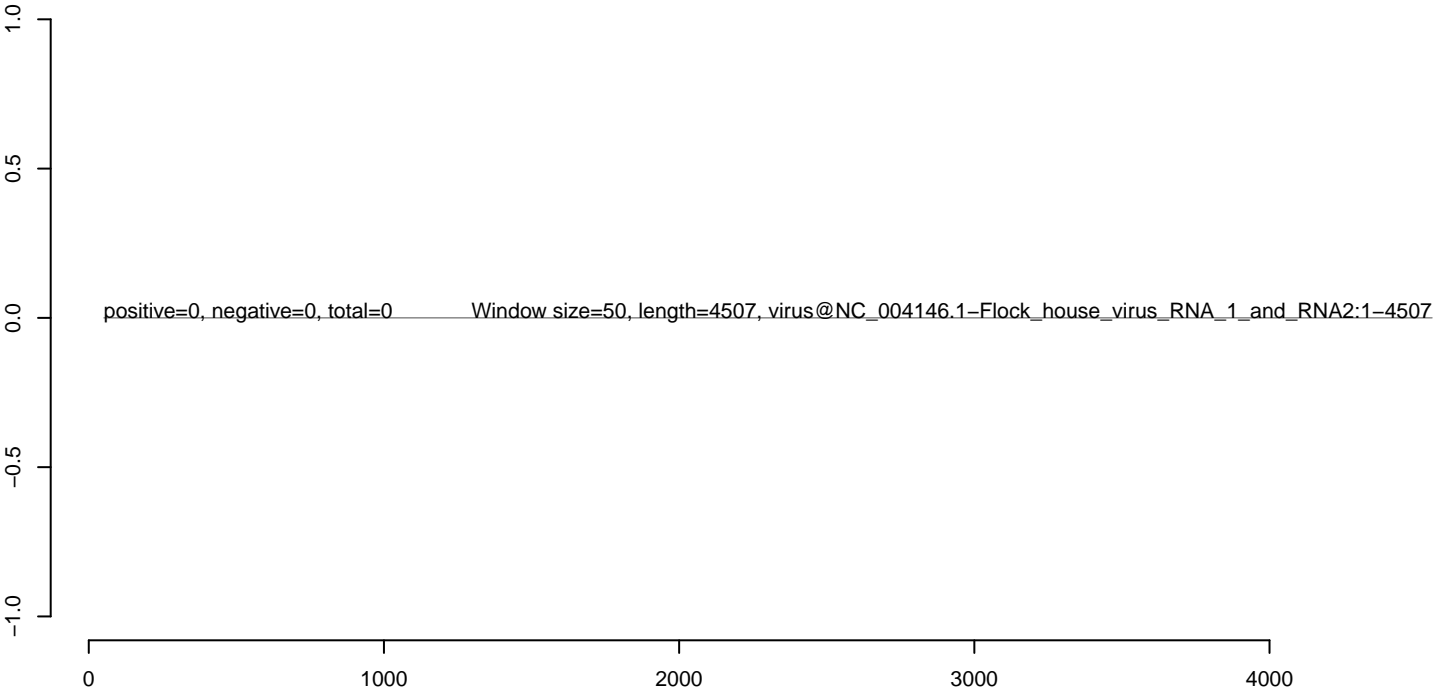
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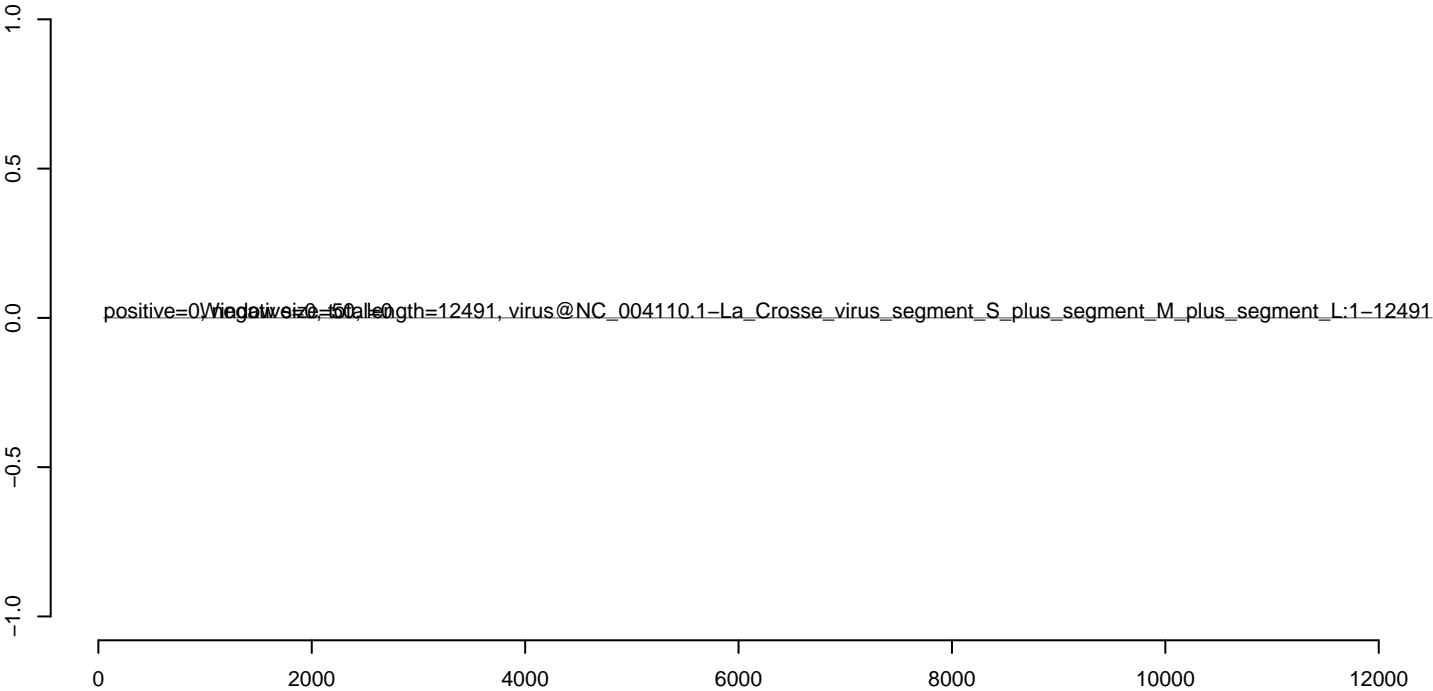
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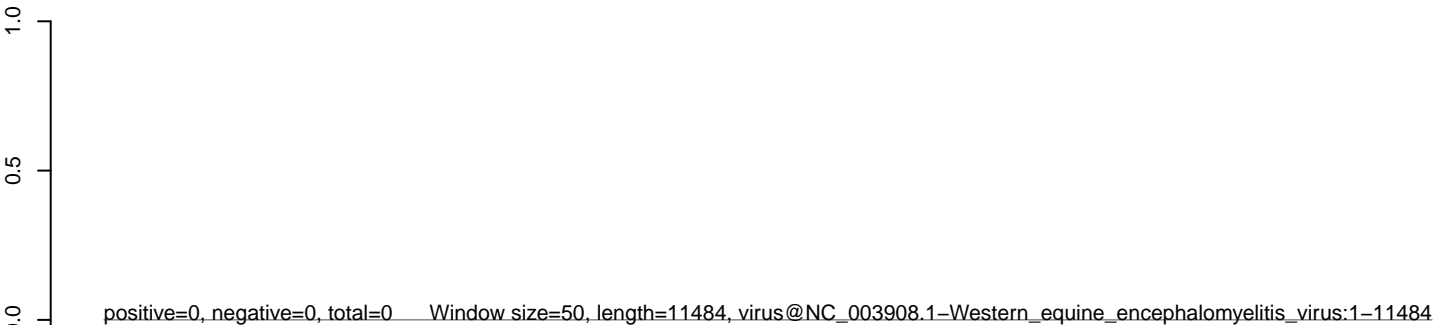
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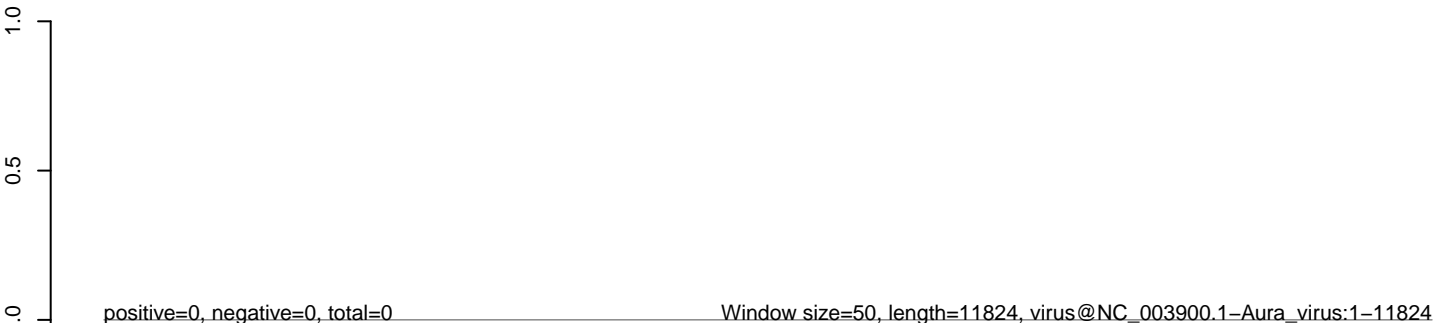
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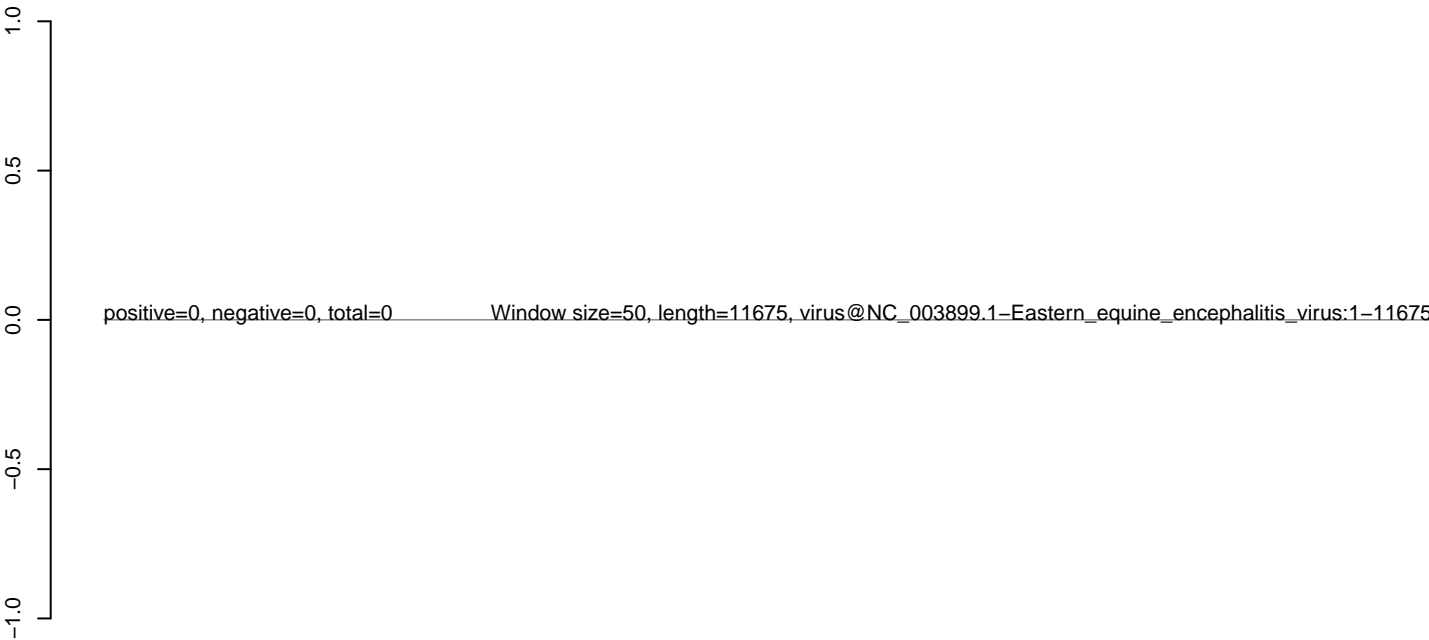
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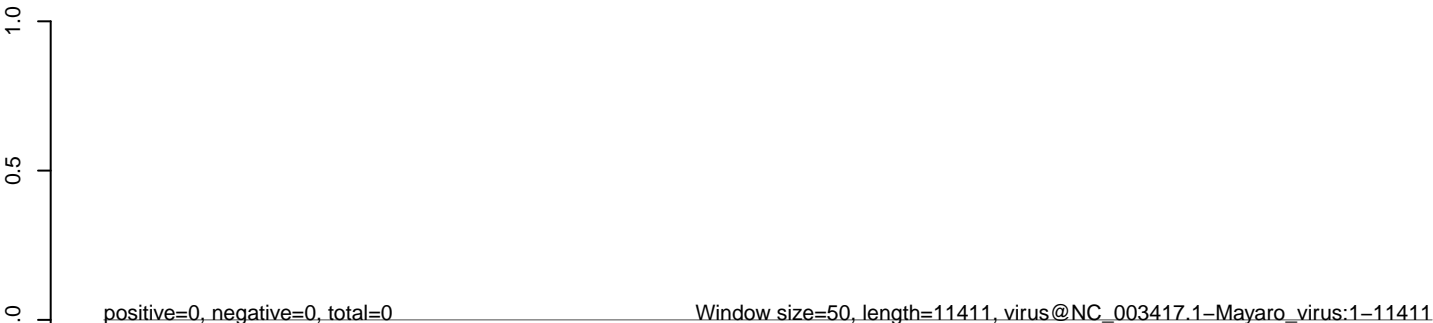
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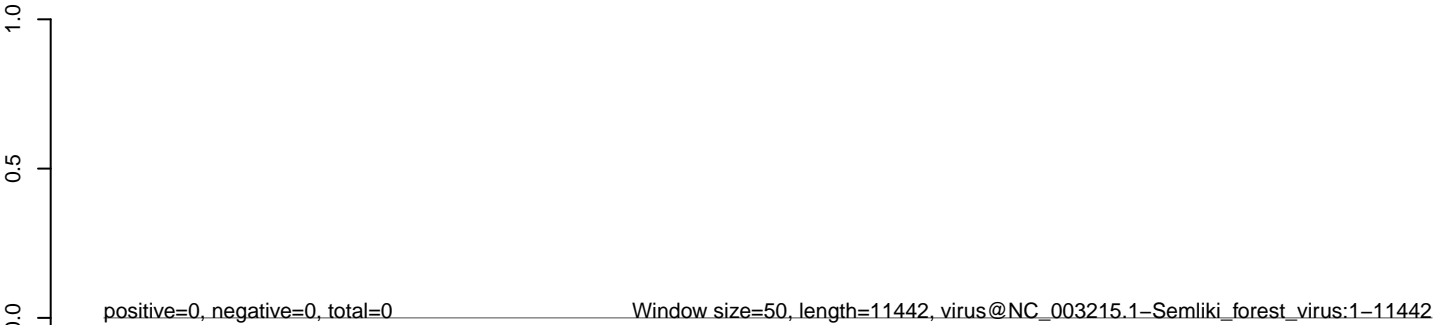
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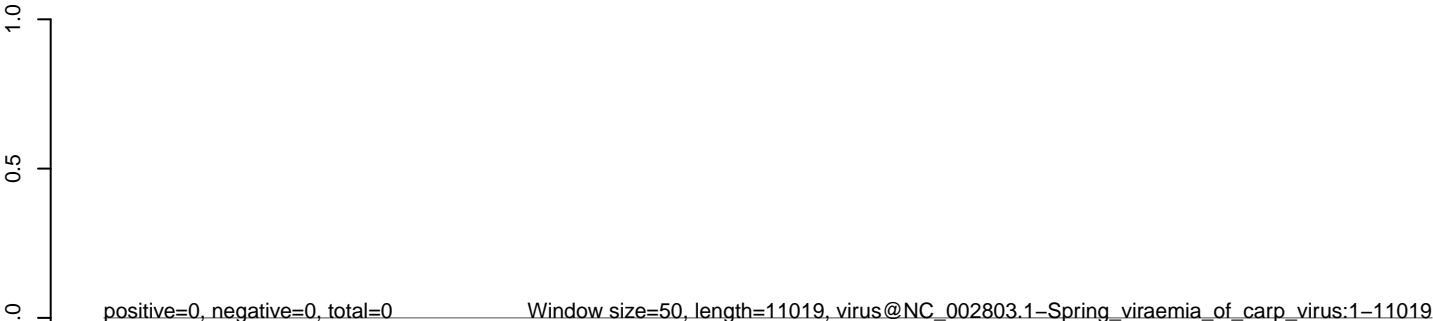
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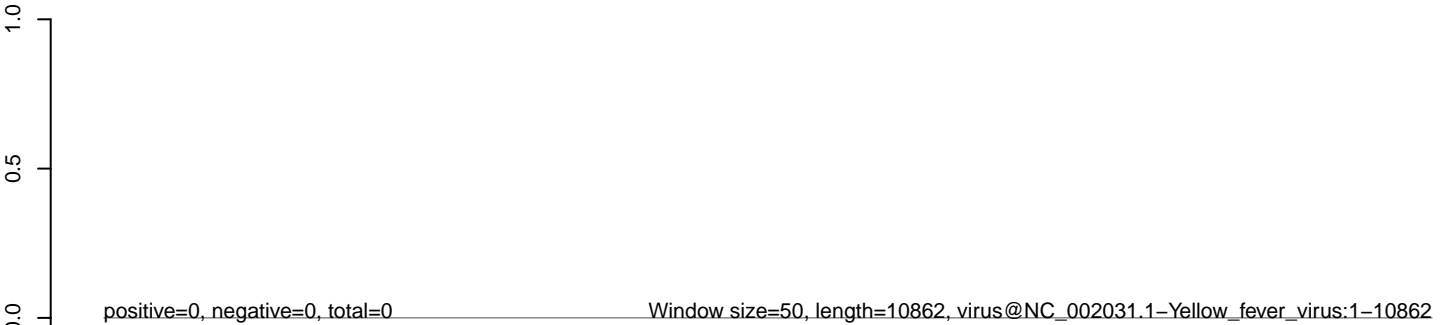
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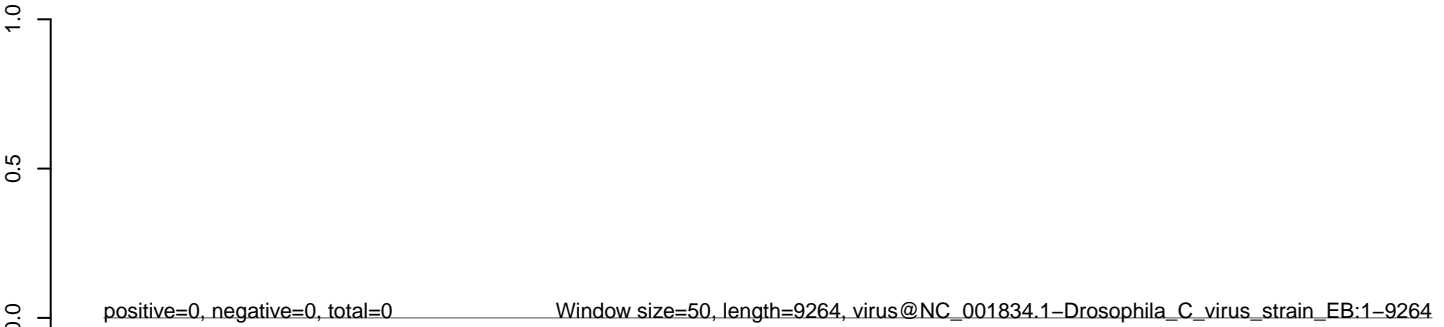
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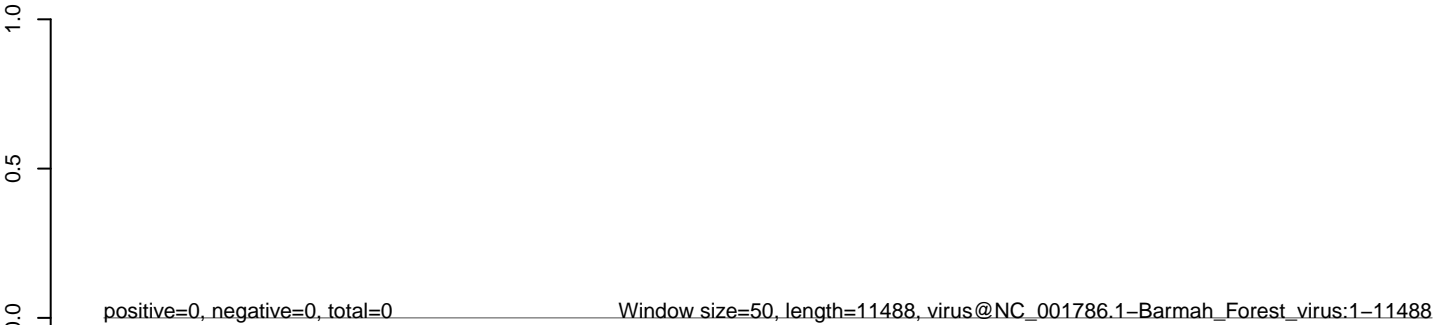
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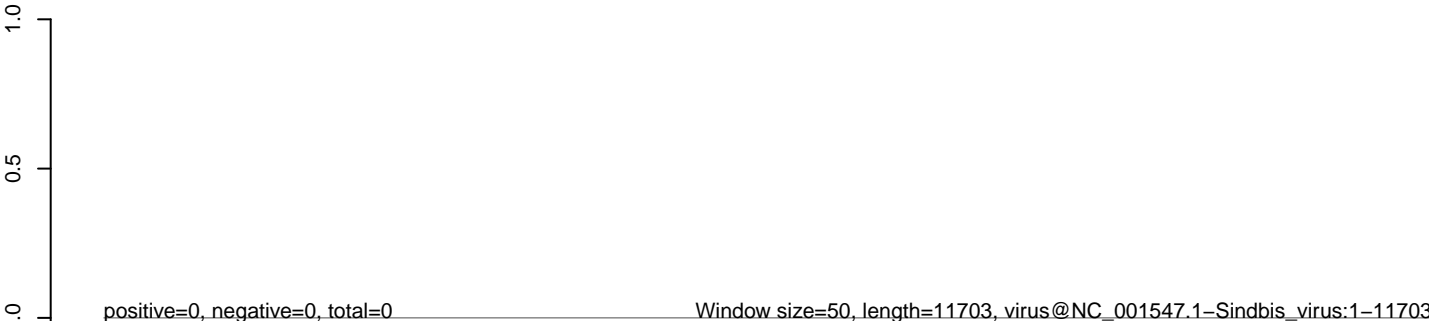
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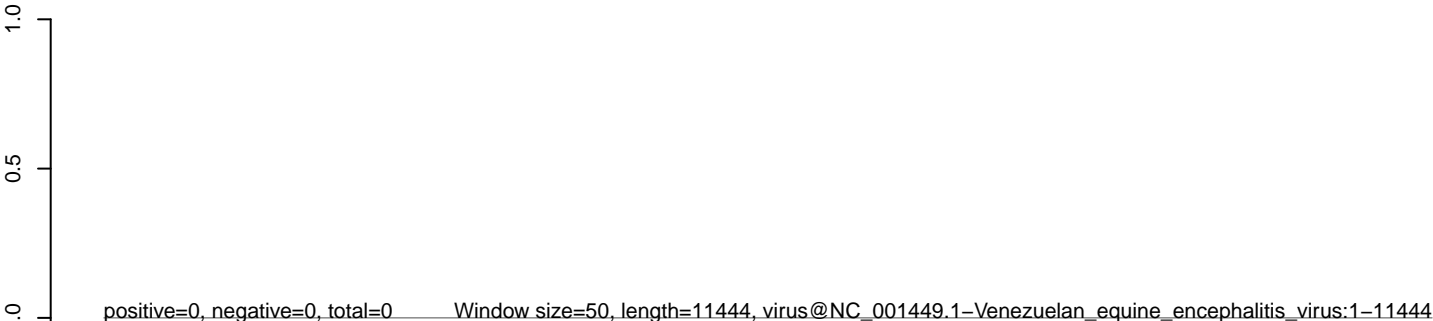
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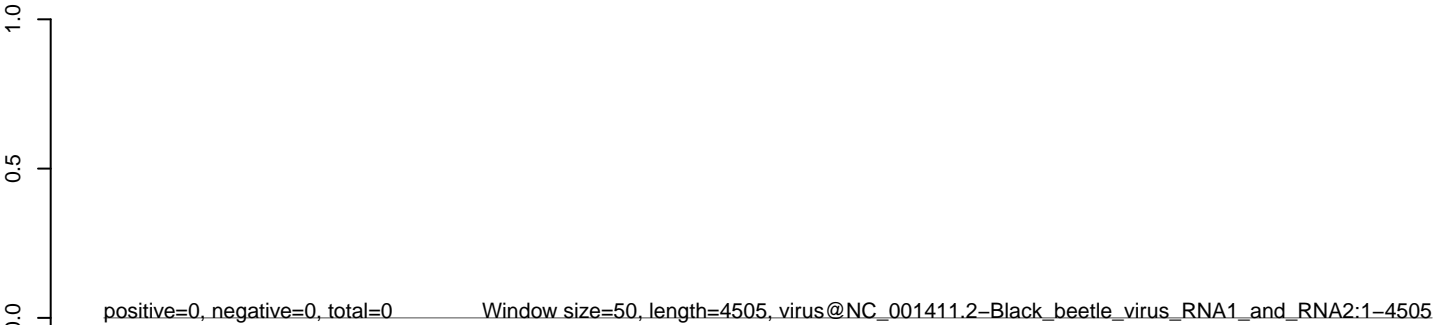
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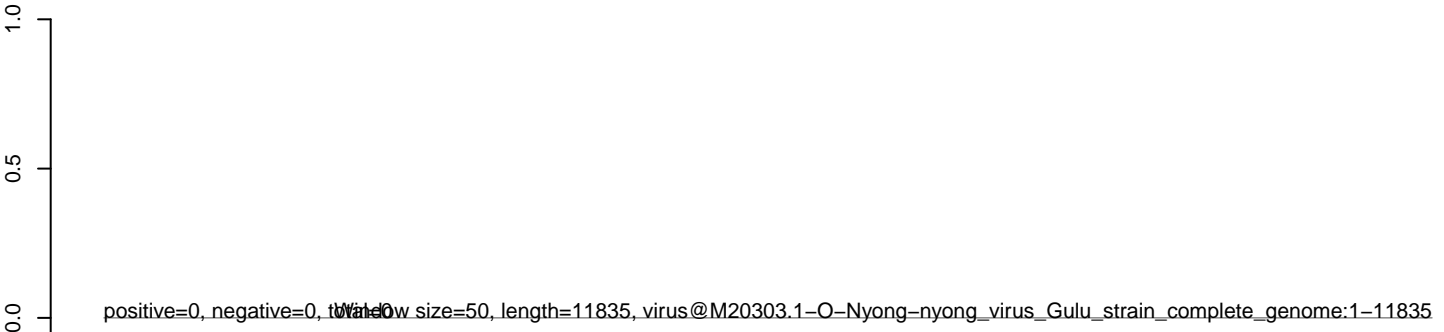
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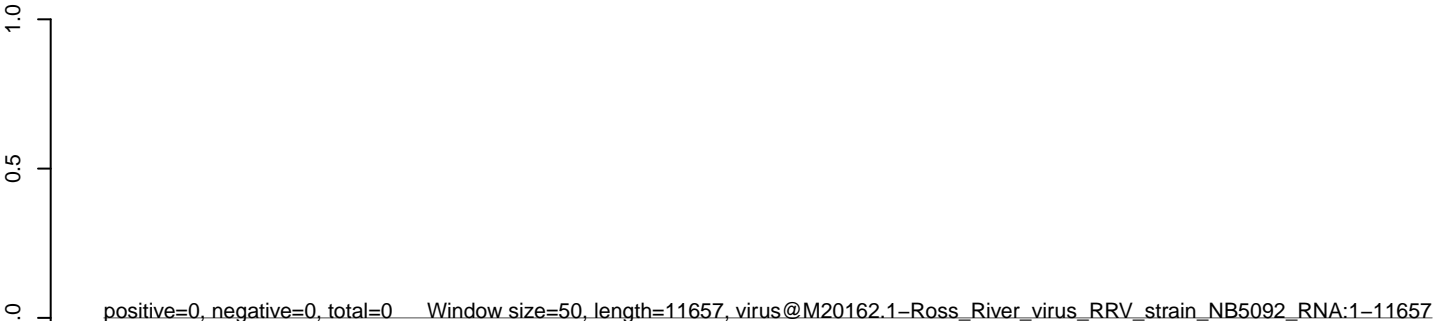
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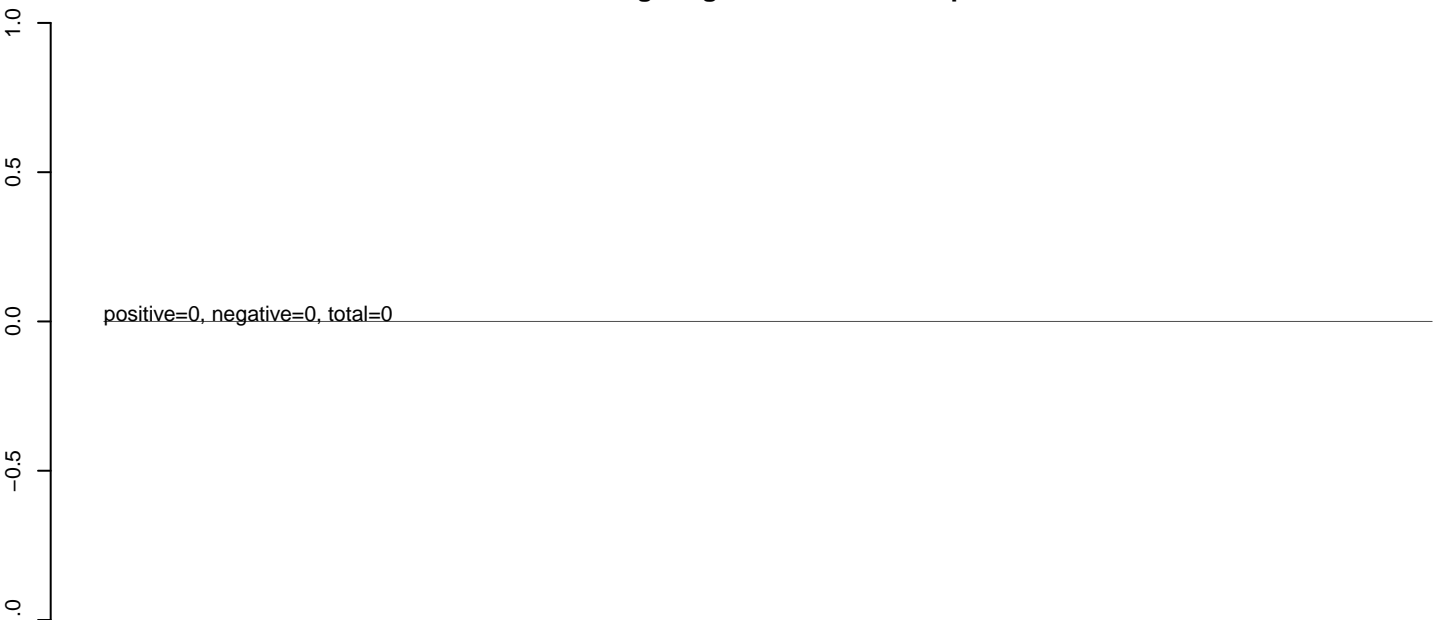
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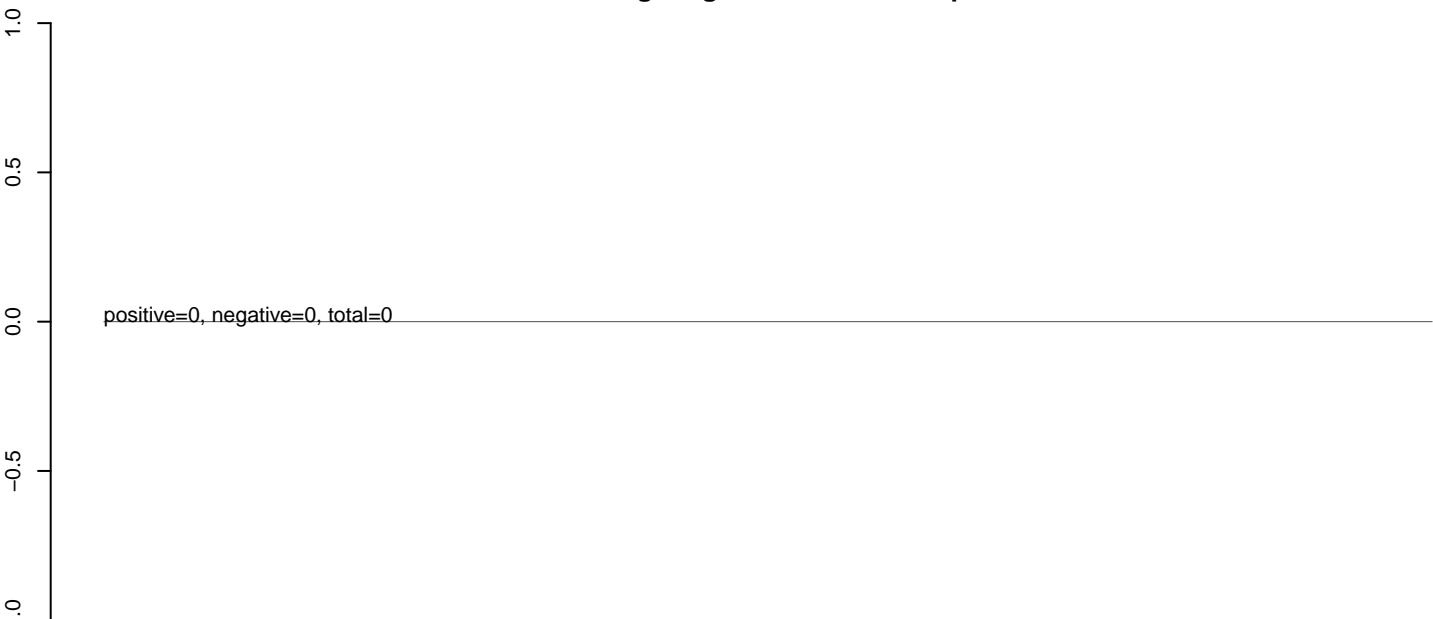
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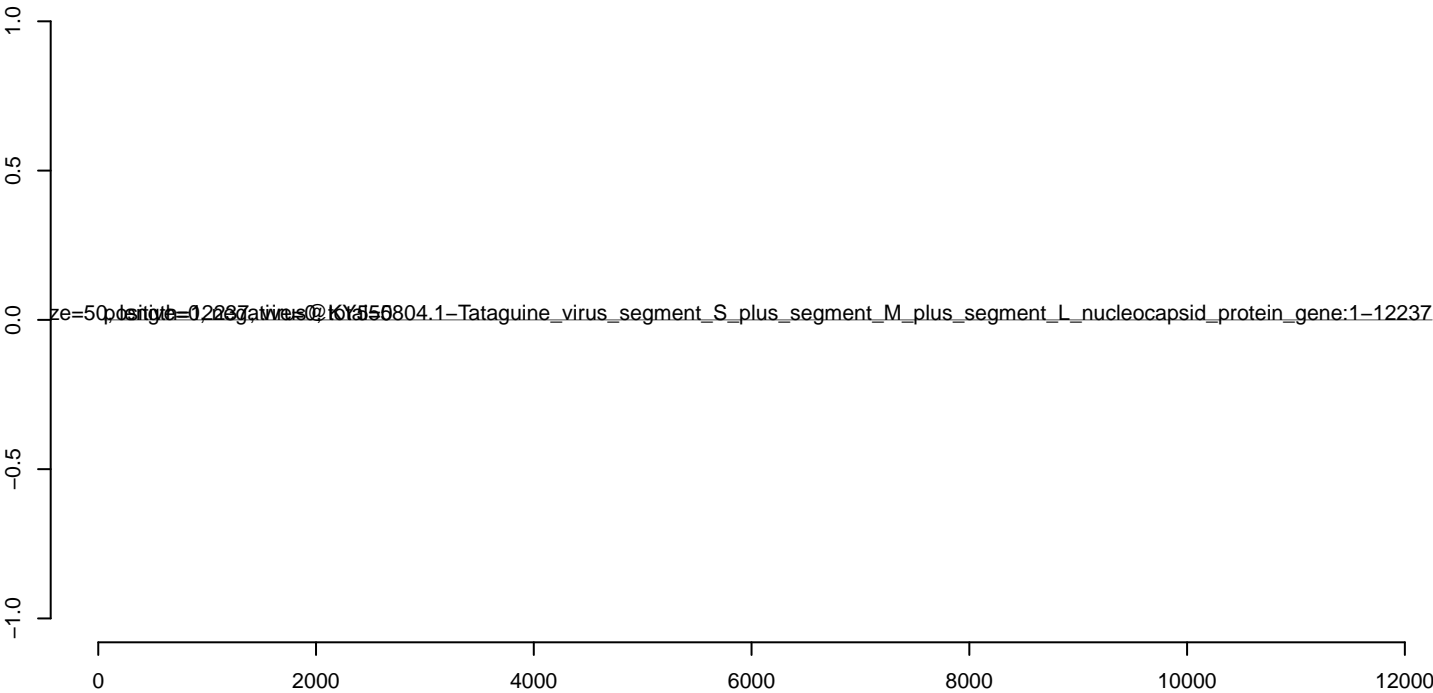
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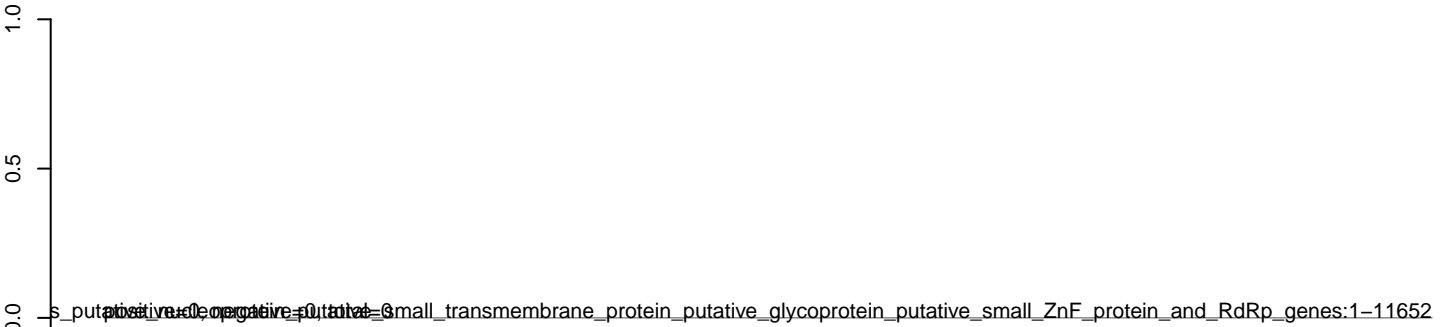
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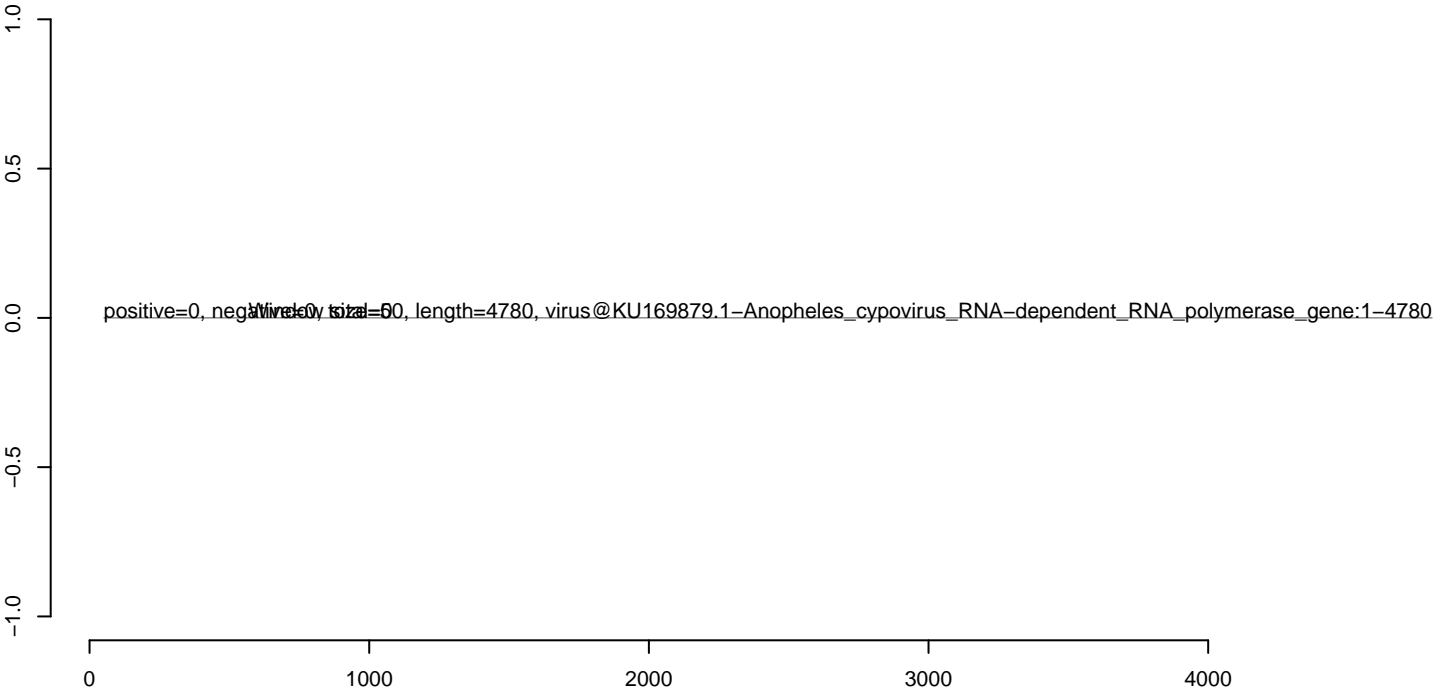
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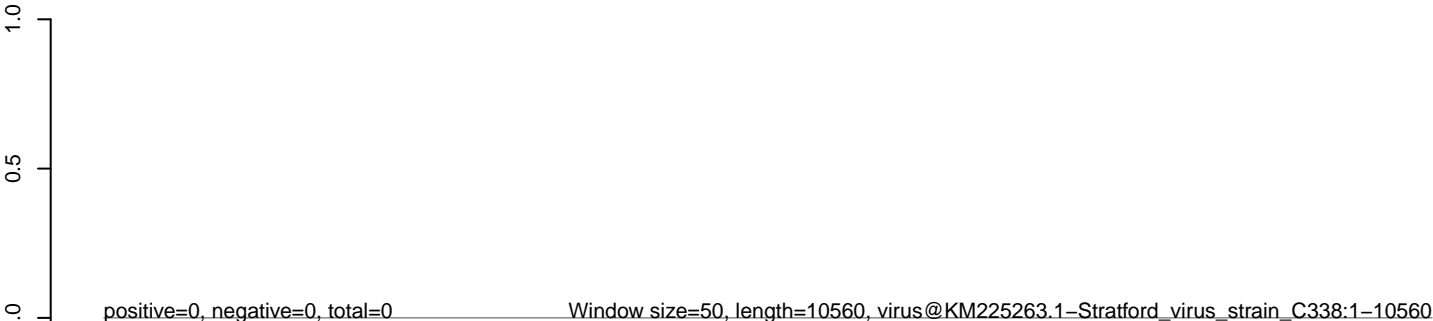
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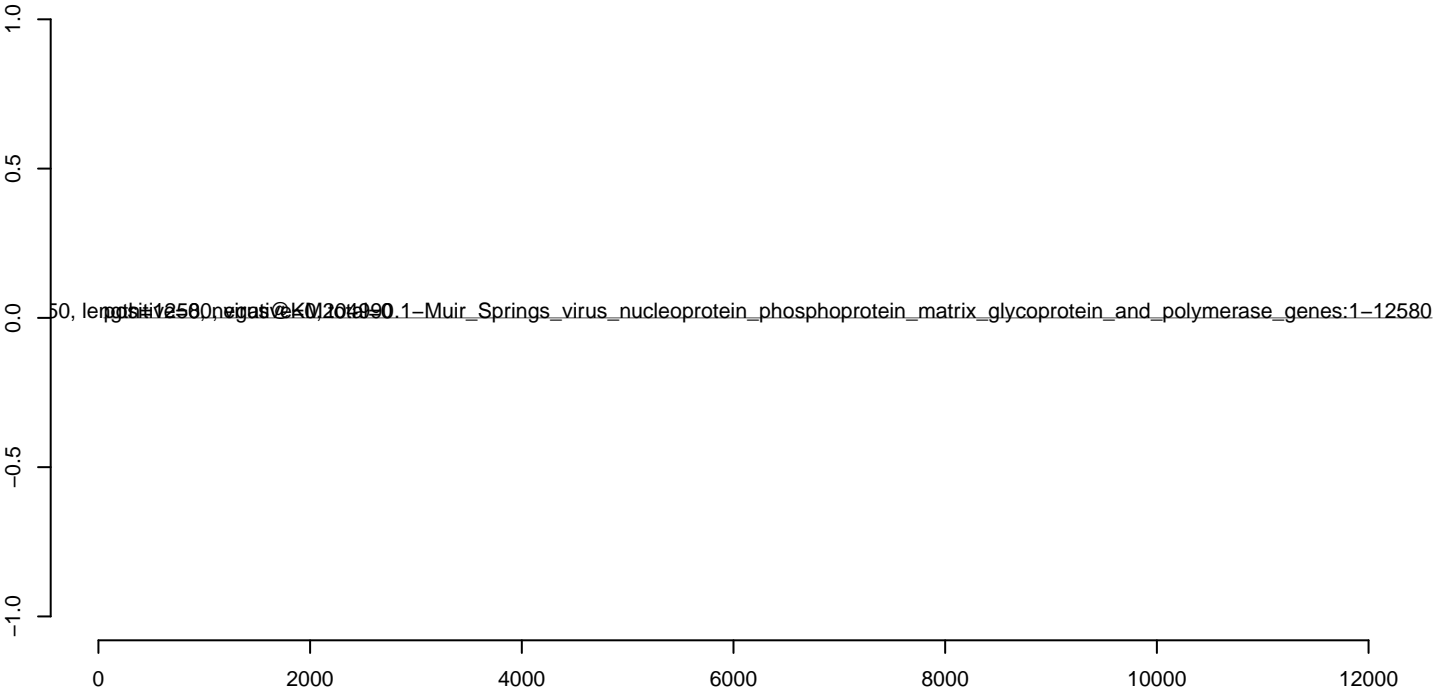
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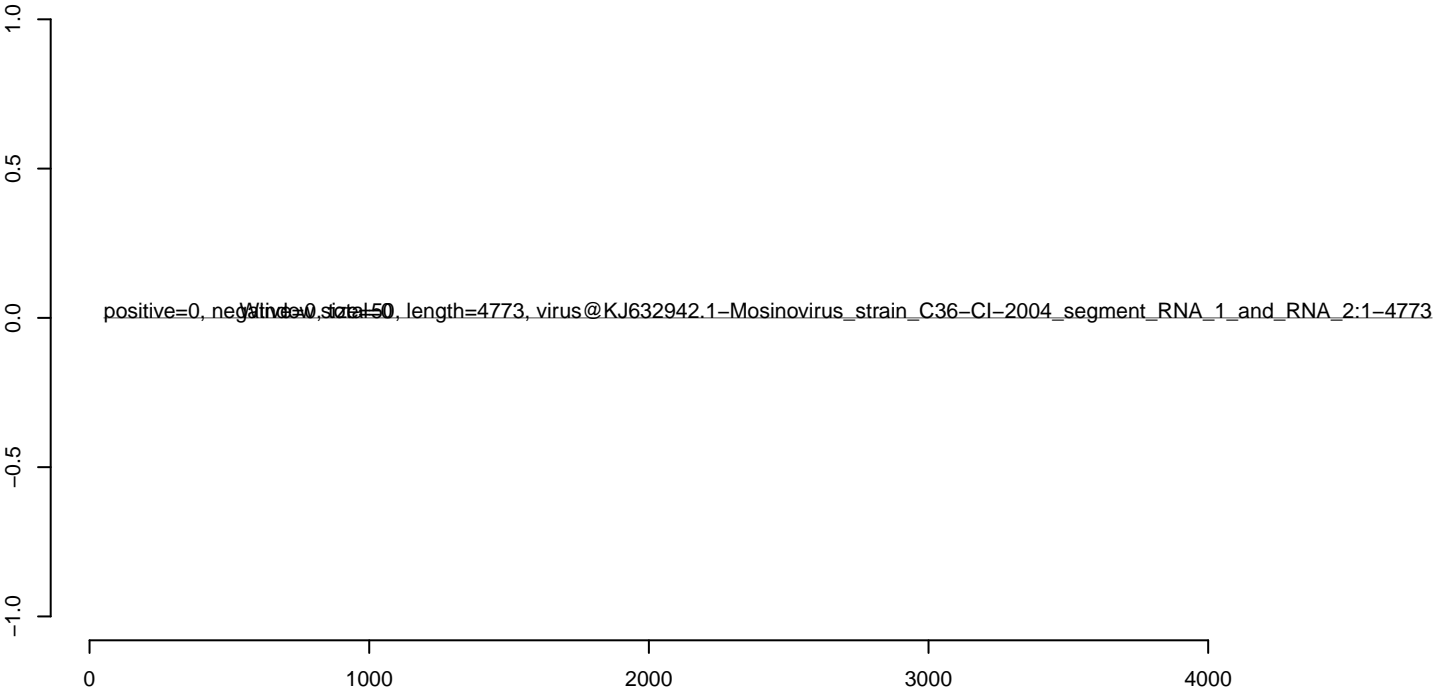
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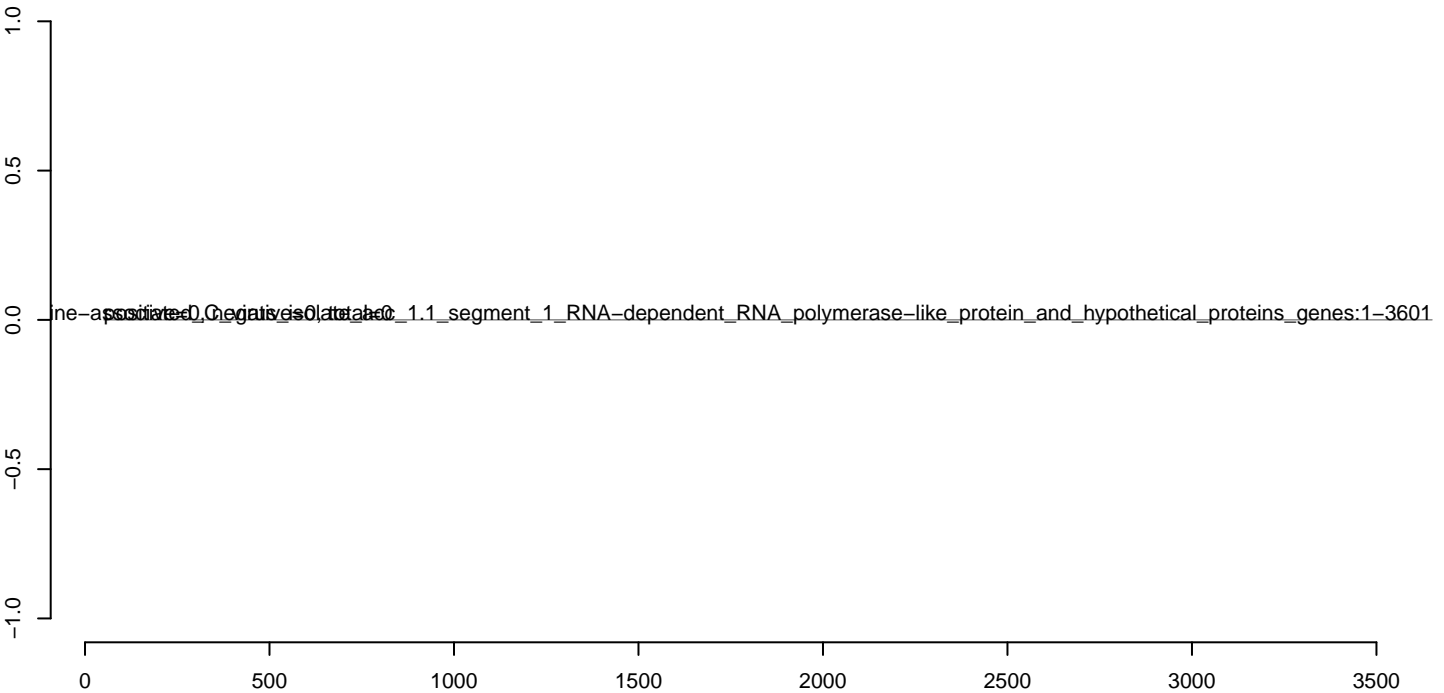
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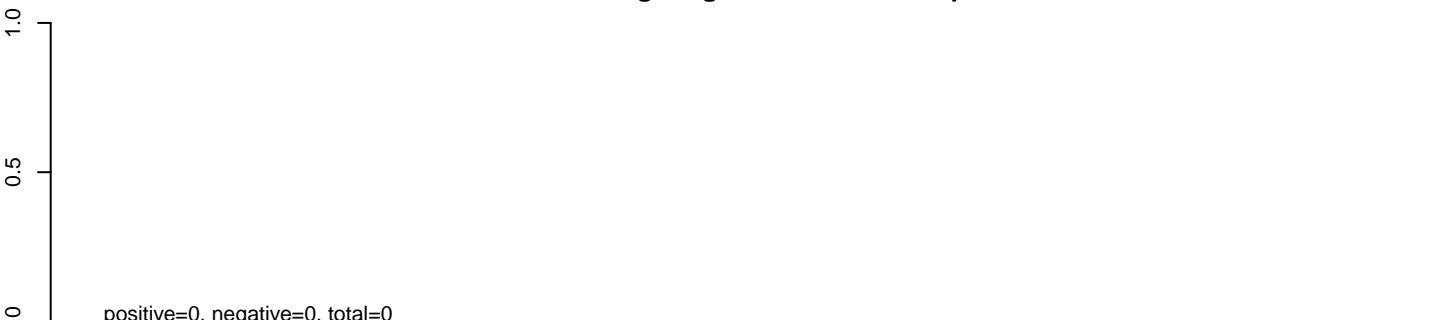
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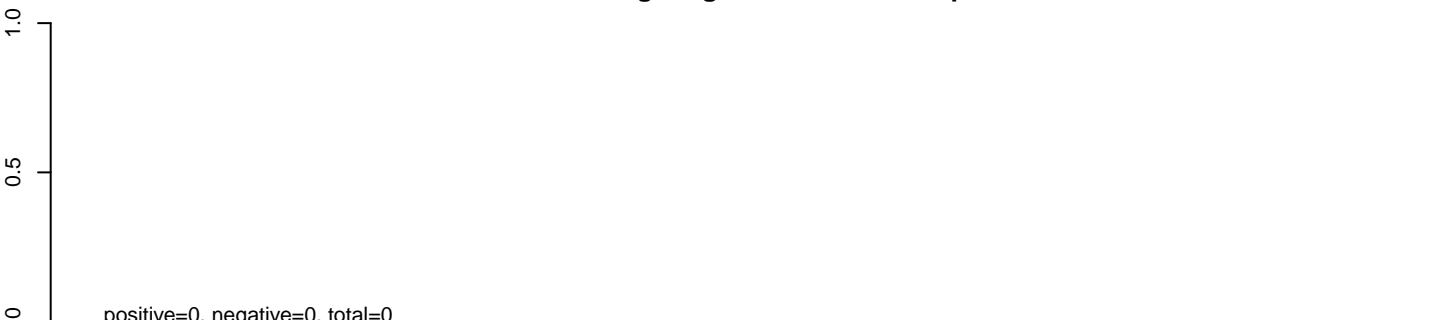
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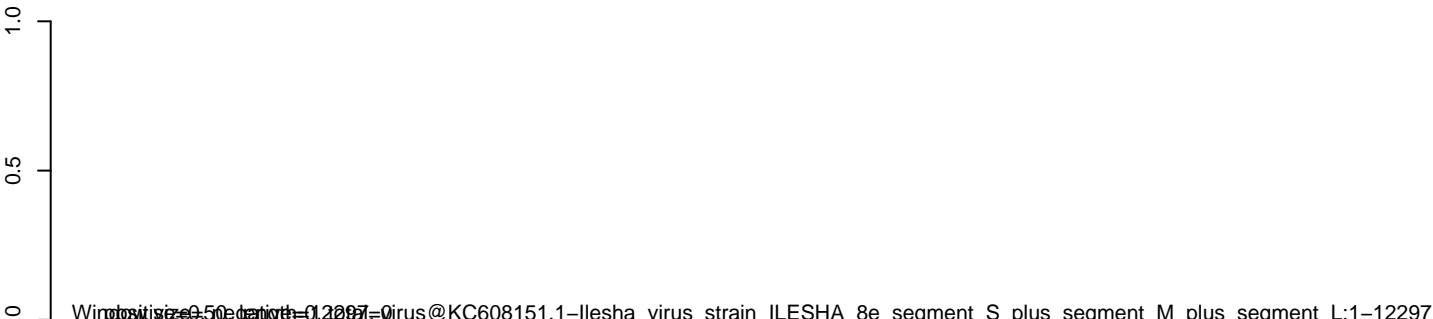
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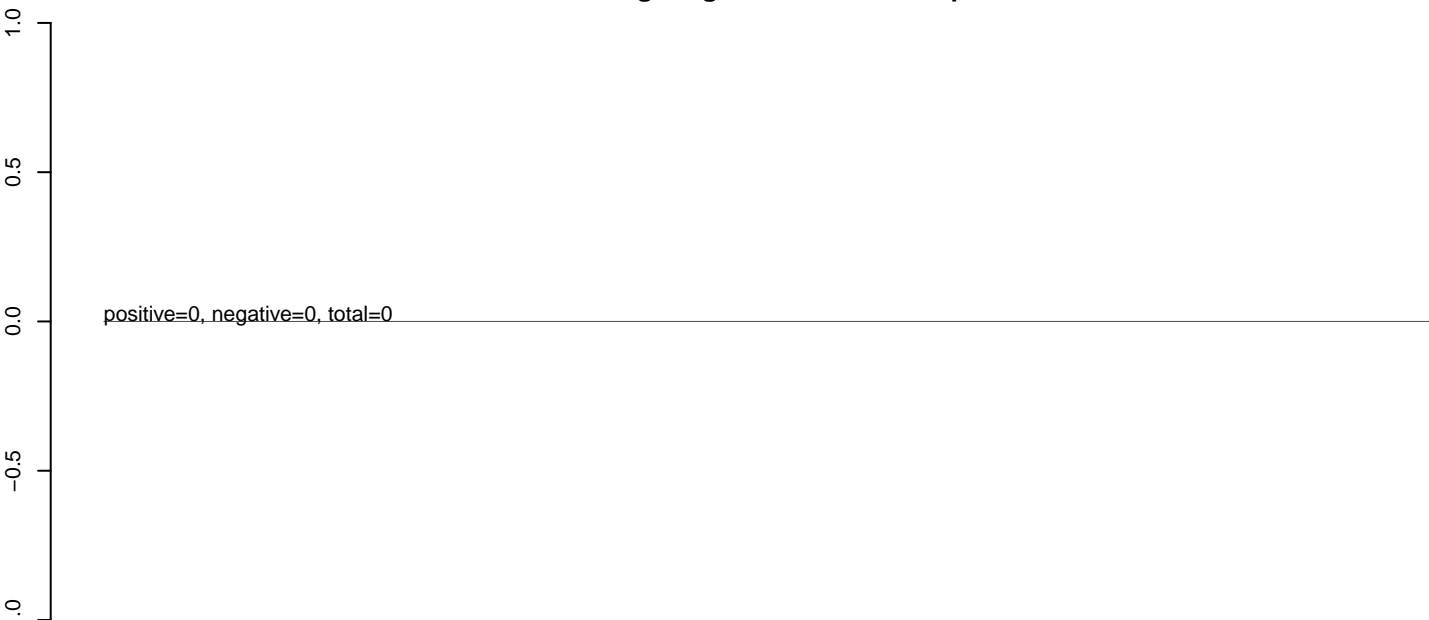


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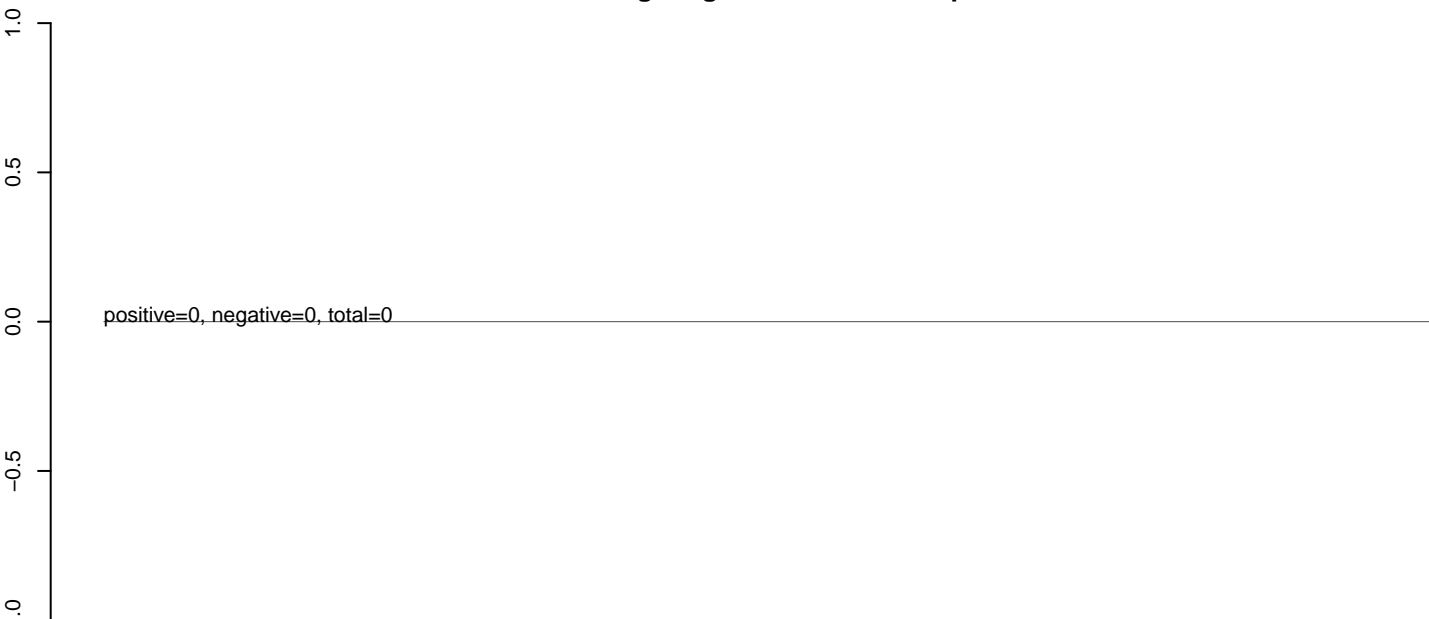


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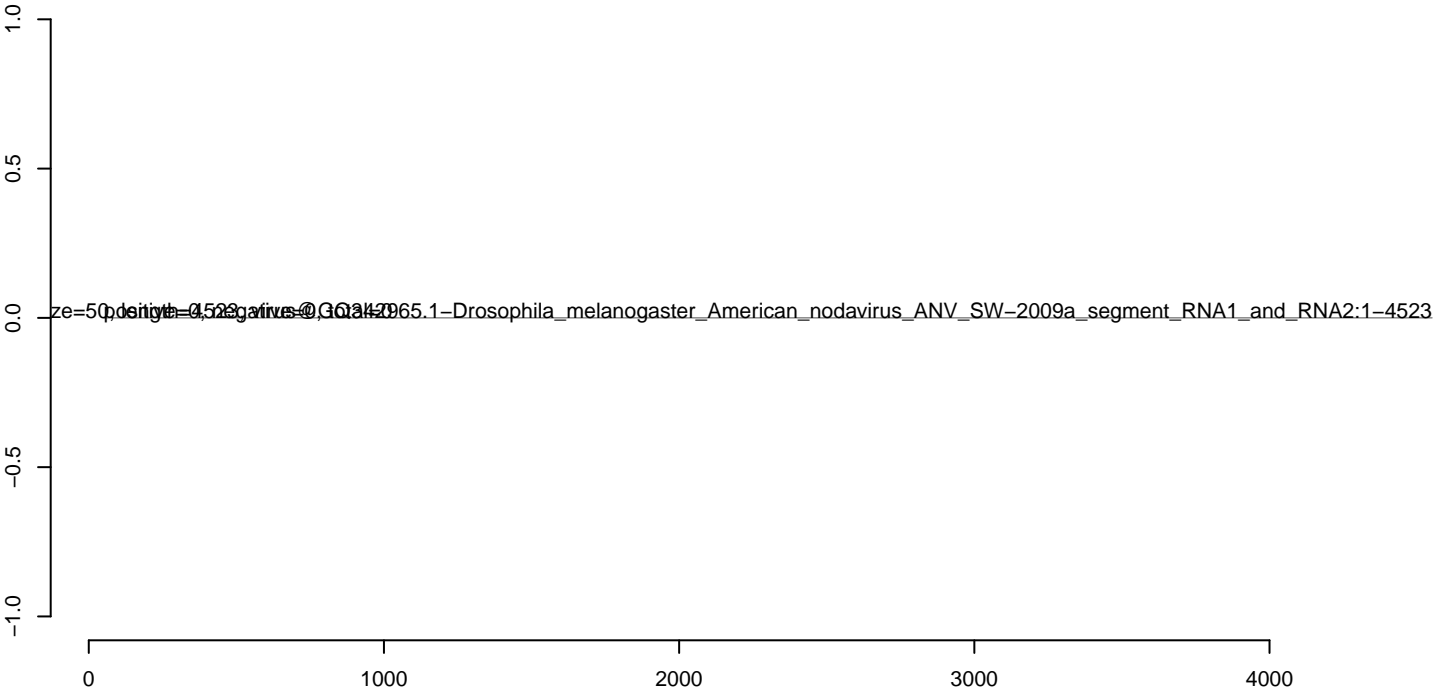
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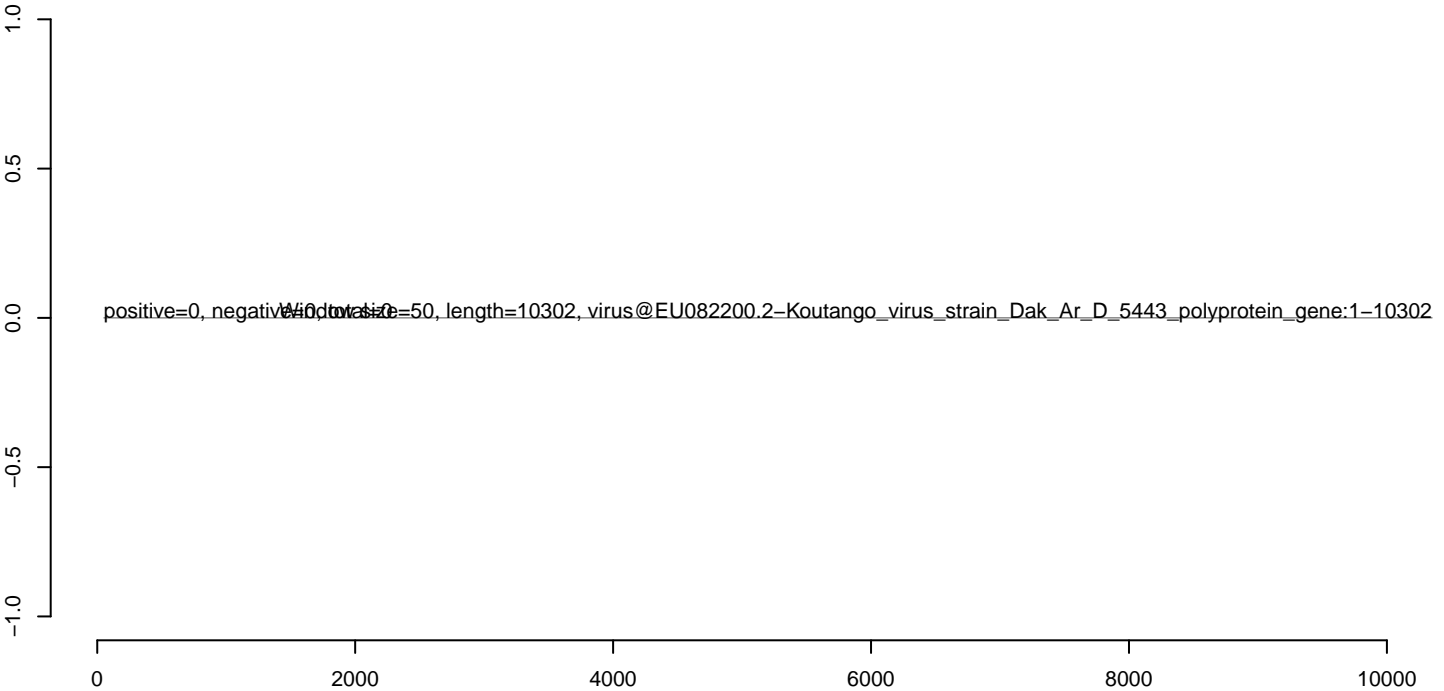
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AeAeg_Aag2_DENV4.rep



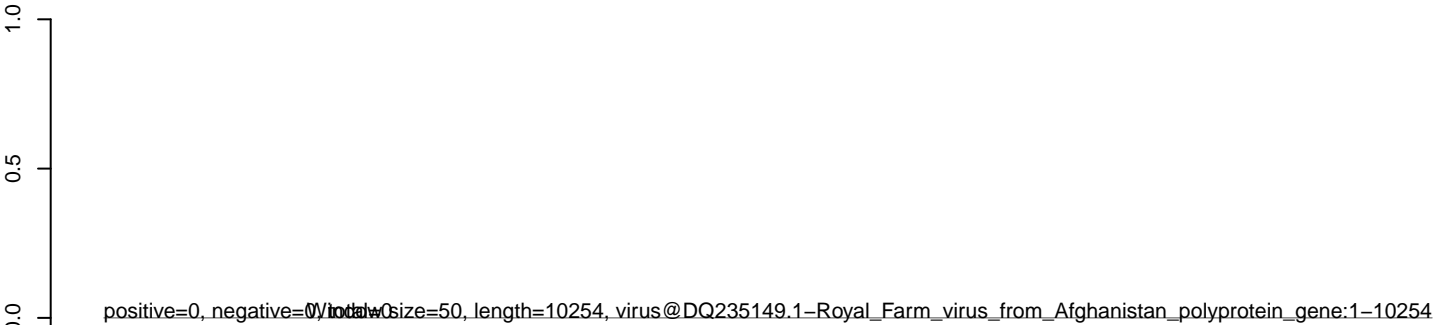
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AeAeg_Aag2_DENV4.rep



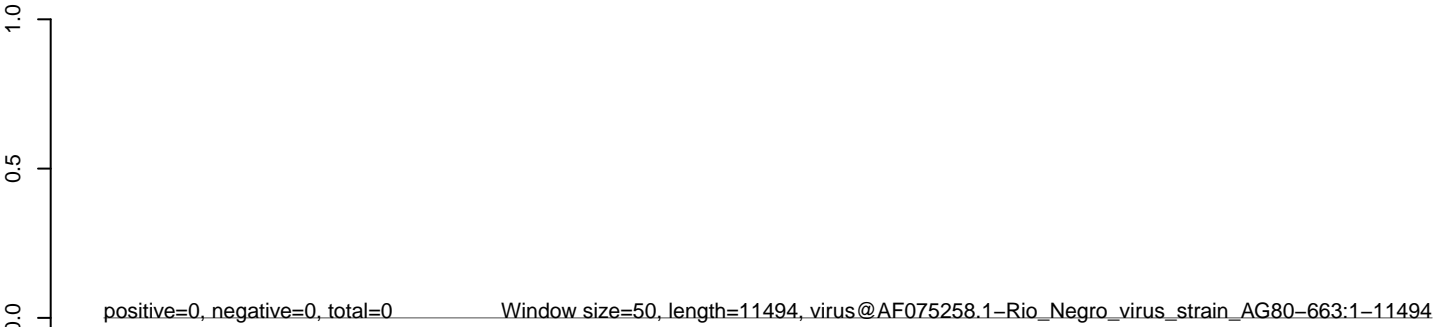
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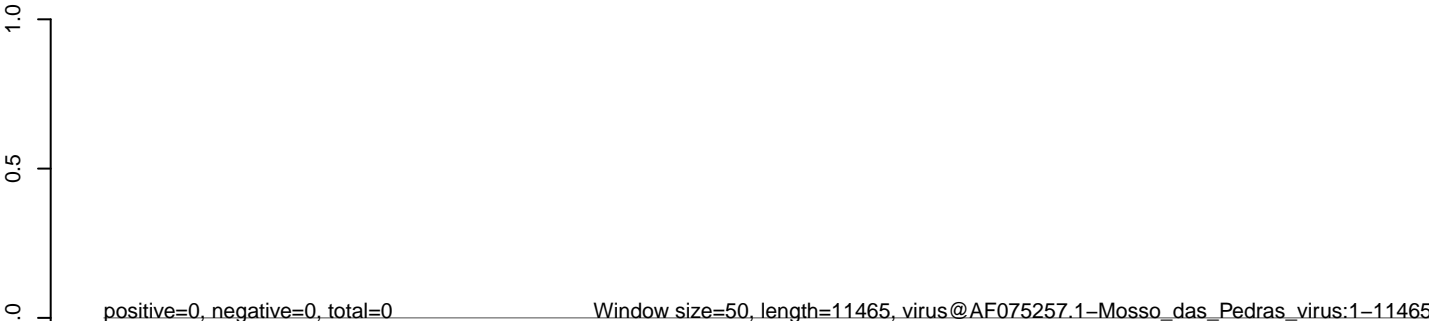
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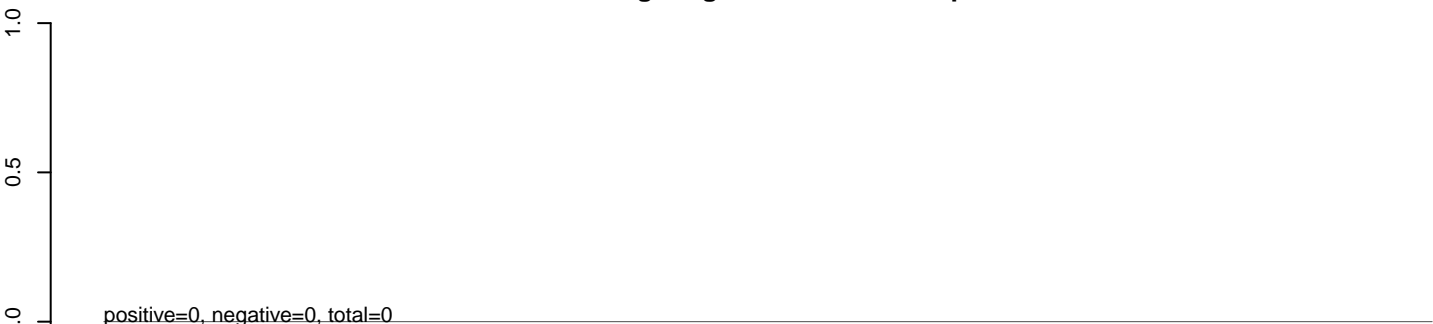


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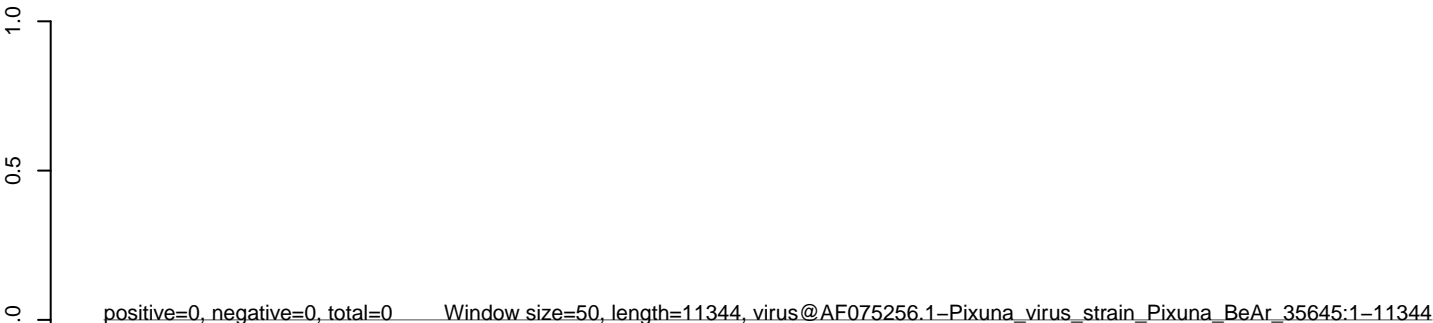
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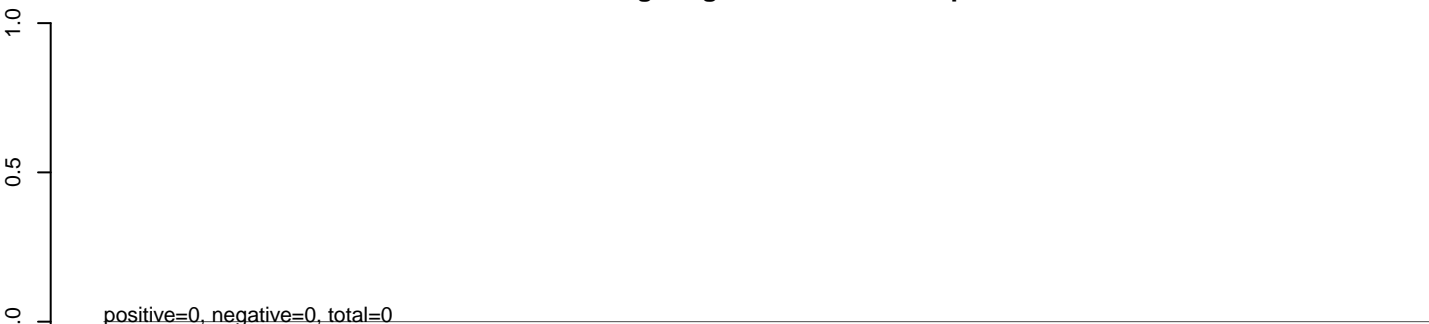


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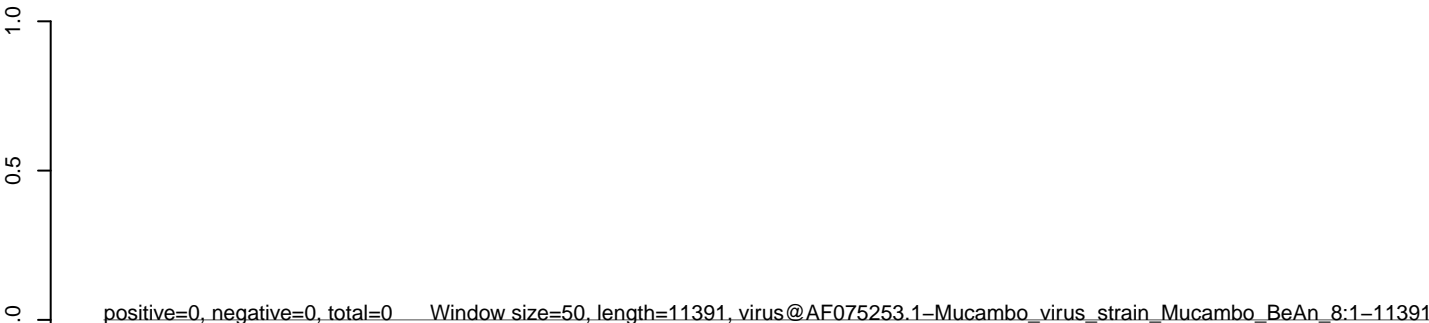
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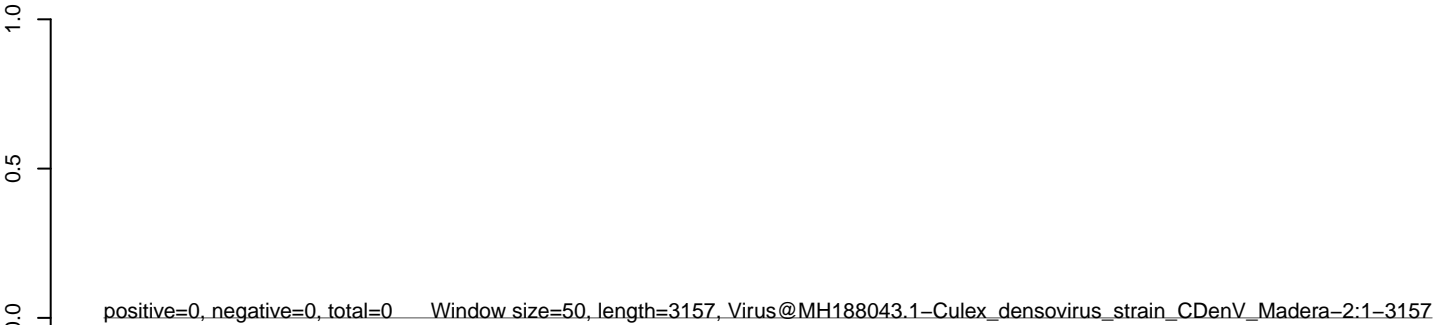
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AeAeg_Aag2_DENV4.rep



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AeAeg_Aag2_DENV4.18_23.rep



AeAeg_Aag2_DENV4.24_35.rep



AeAeg_Aag2_DENV4.rep

