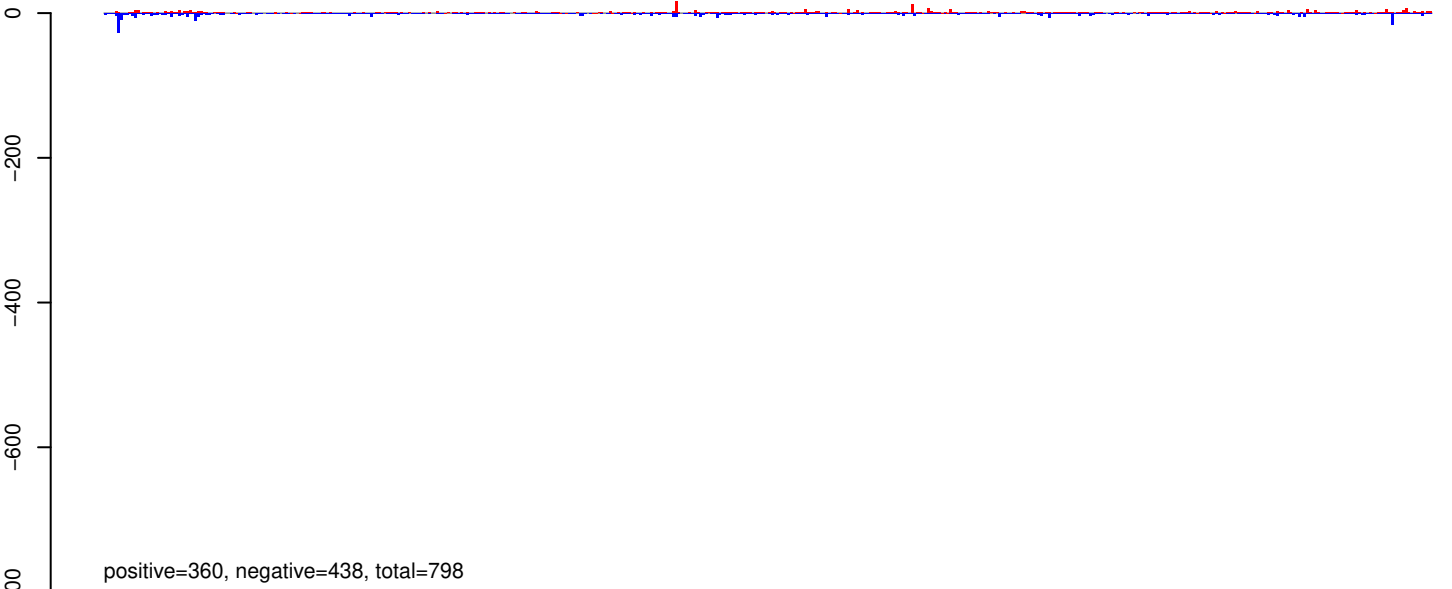
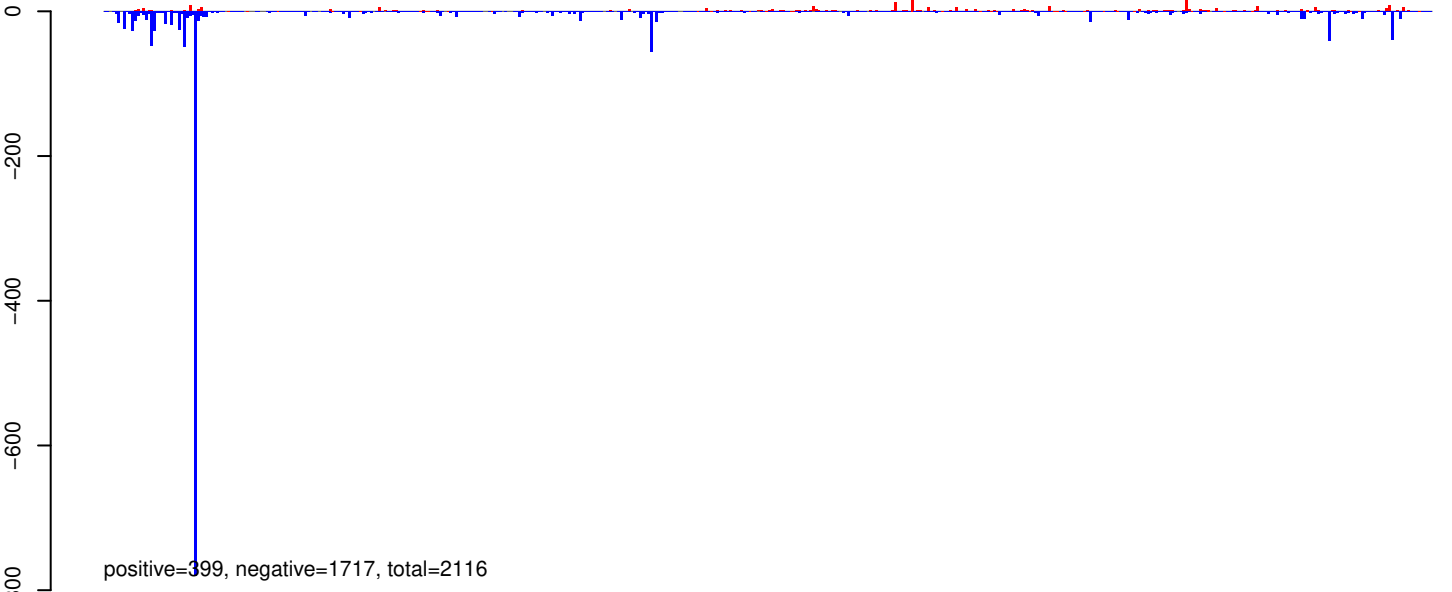


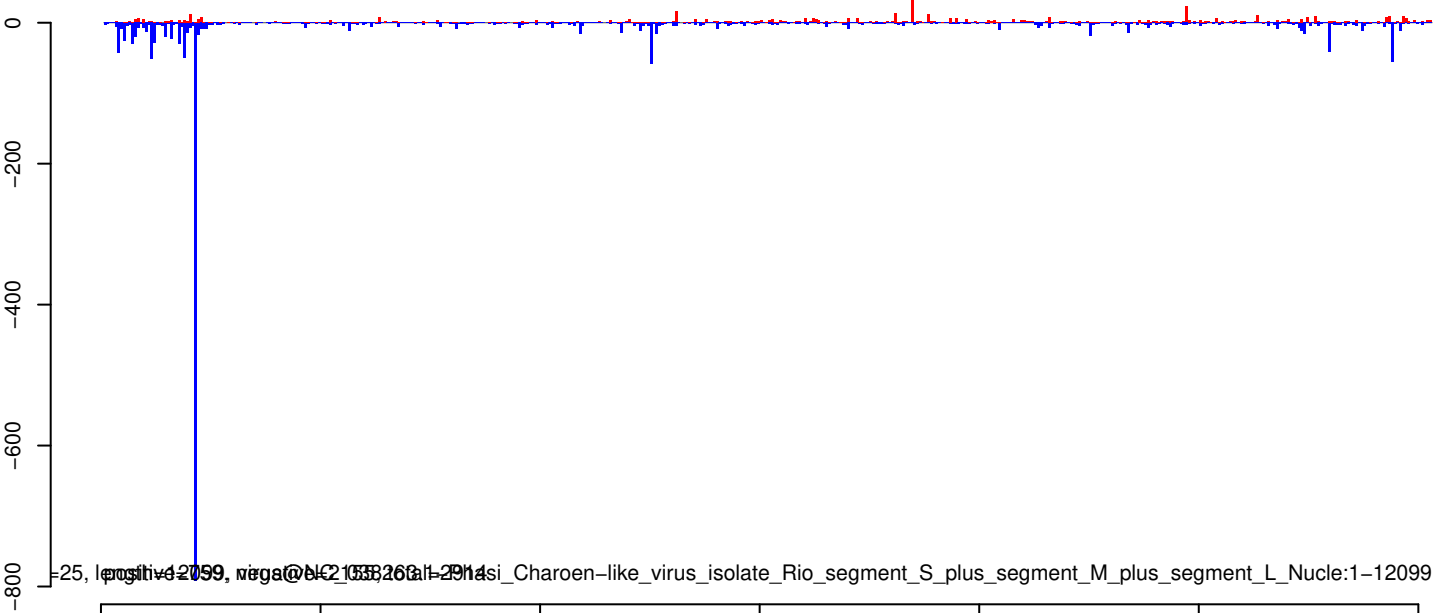
AeAeg_Aag2_BTV2_AK.18_23.rep



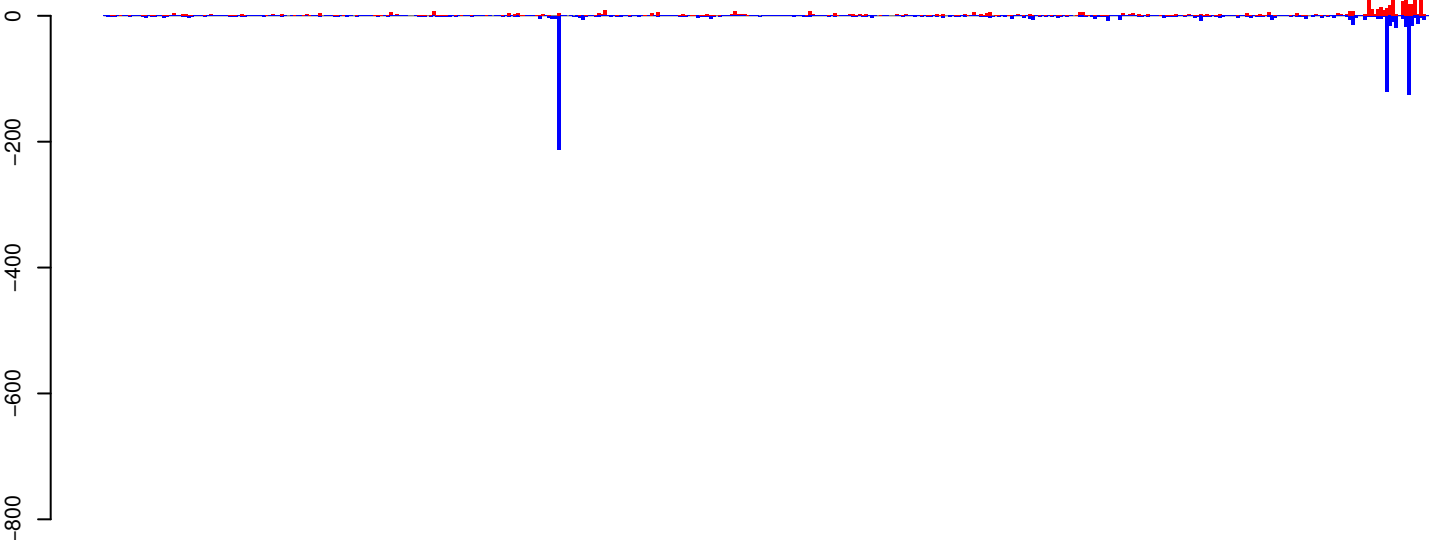
AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep

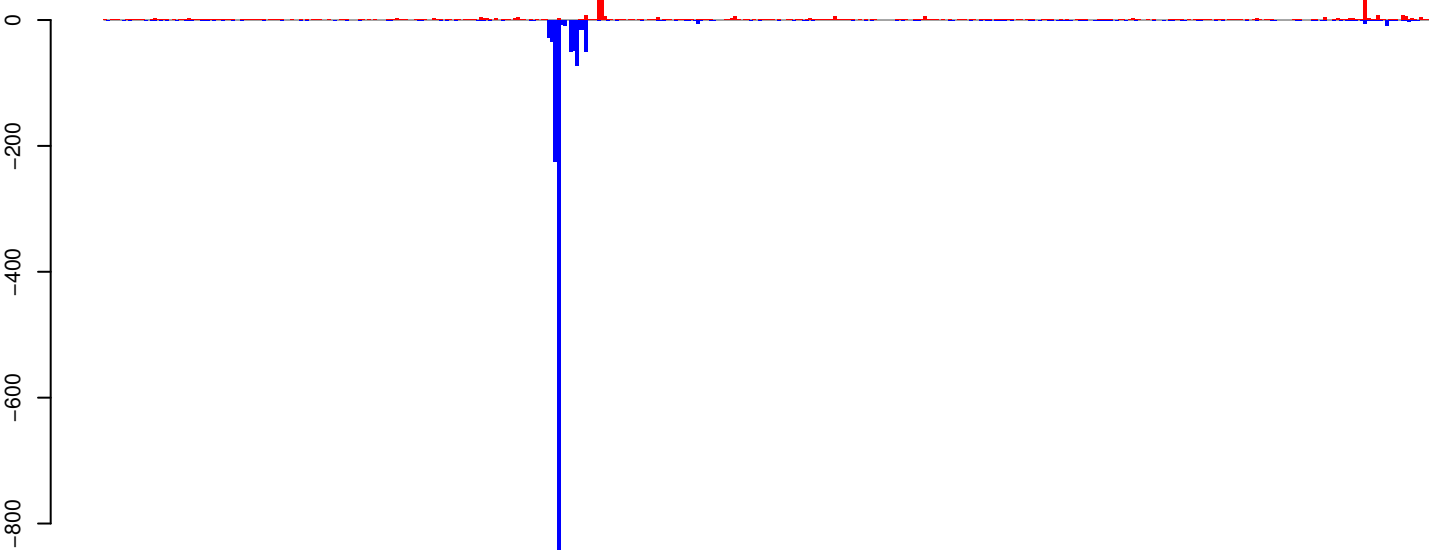


AeAeg_Aag2_BTV2_AK.18_23.rep



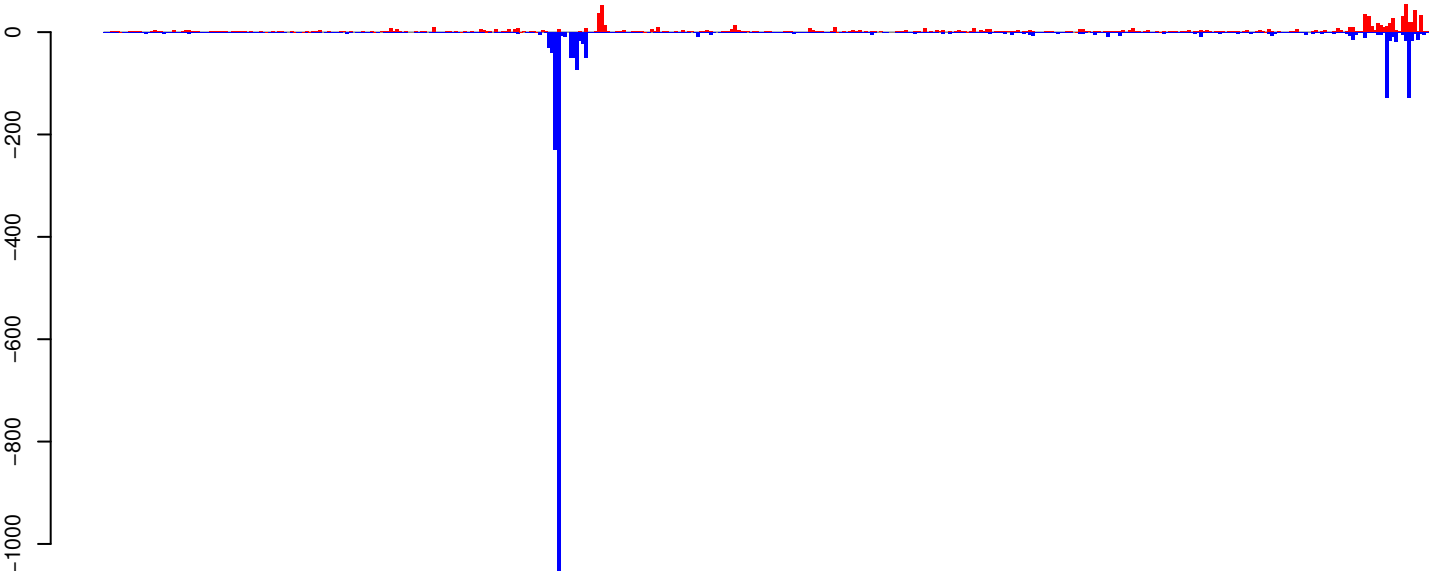
positive=707, negative=942, total=1649

AeAeg_Aag2_BTV2_AK.24_35.rep



positive=302, negative=1496, total=1798

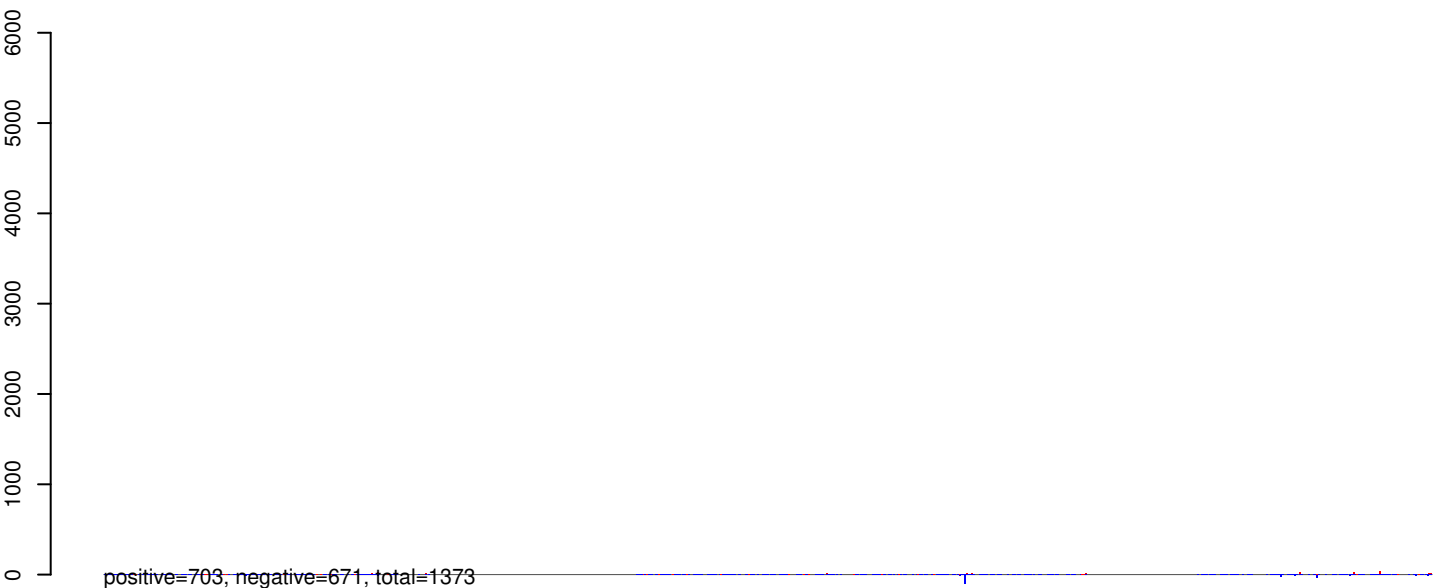
AeAeg_Aag2_BTV2_AK.rep



positive=1009, negative=2438, total=3447, size=25, length=10682, virus@NC_001564.2-Cell_fusing_agent_virus_strain_Galveston:1-10682

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

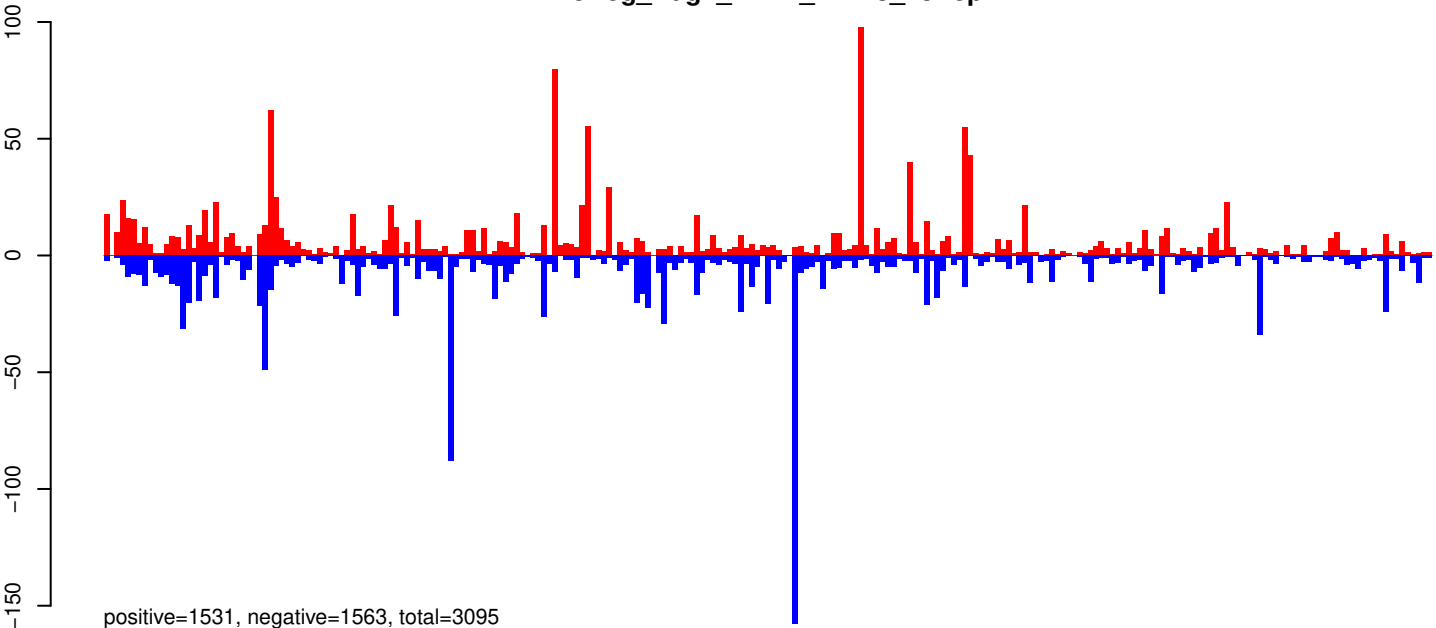


AeAeg_Aag2_BTV2_AK.rep

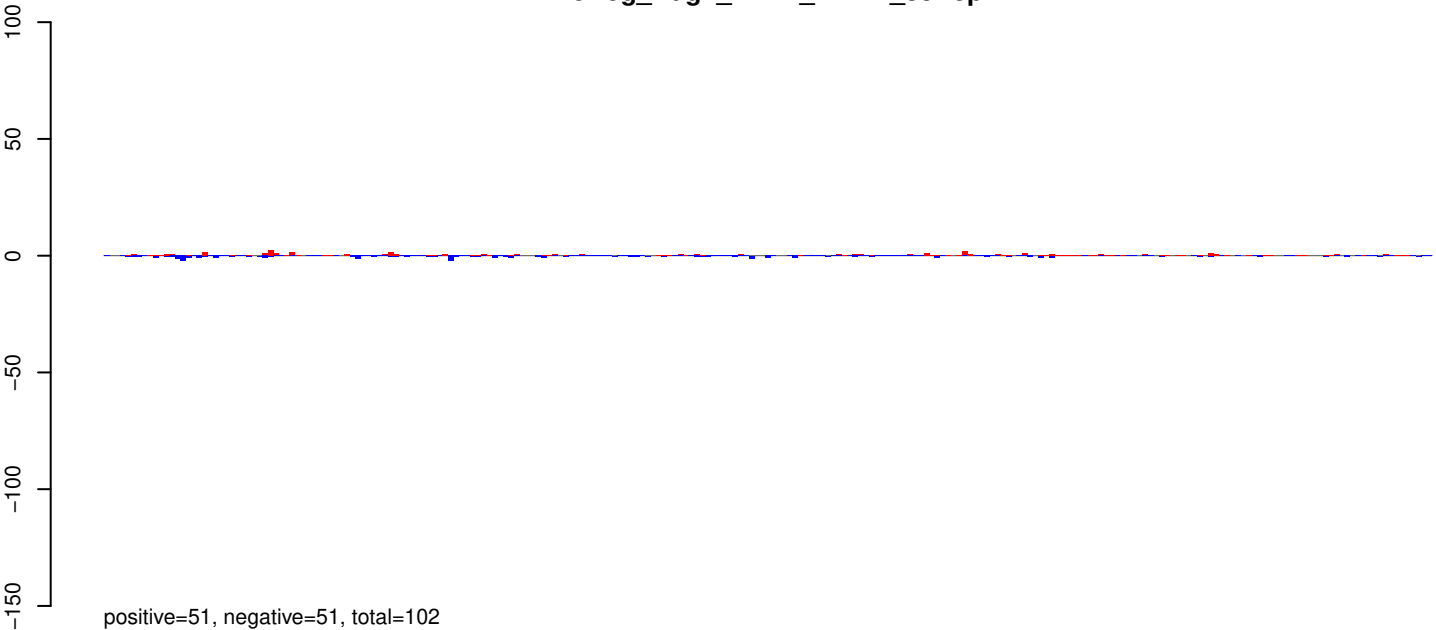


0 5000 10000 15000 20000

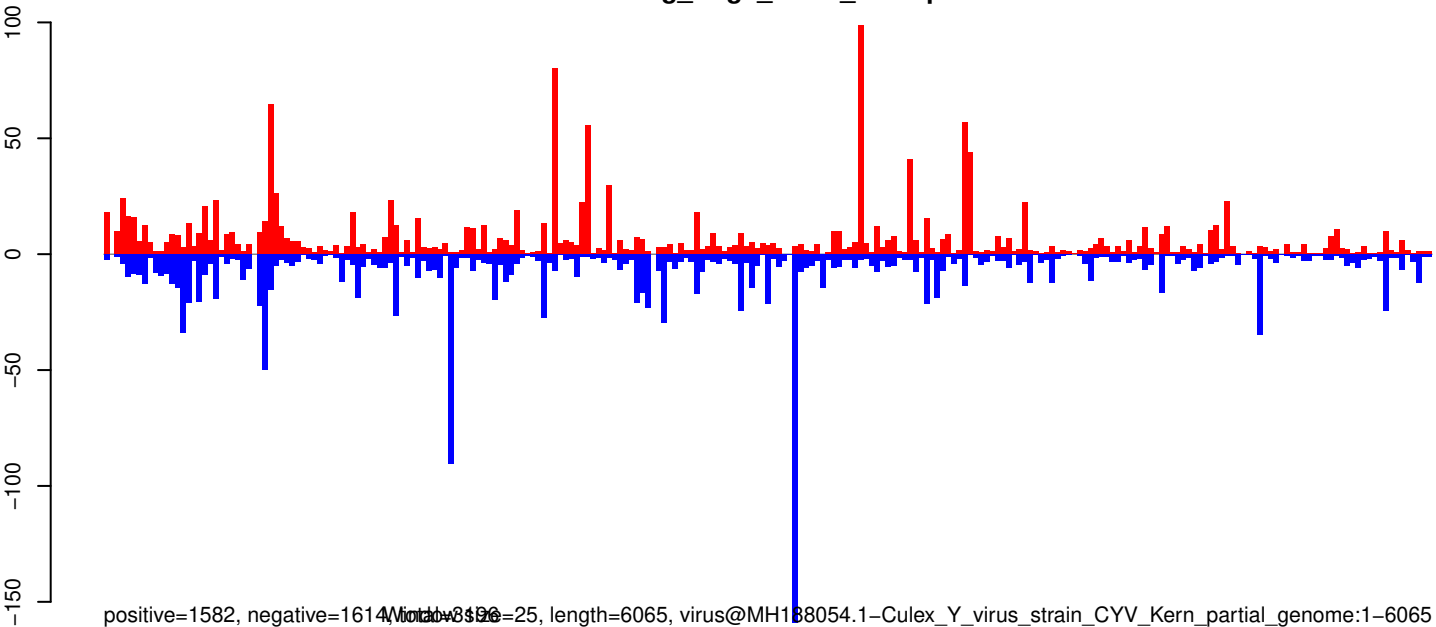
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

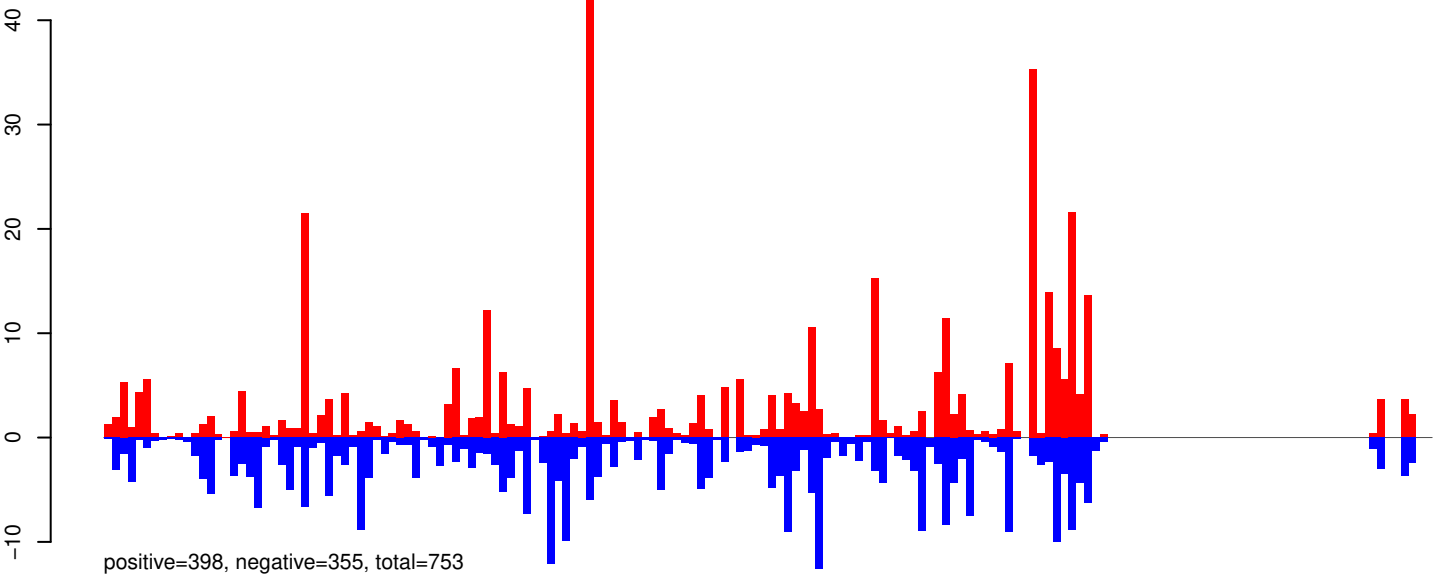


AeAeg_Aag2_BTV2_AK.rep

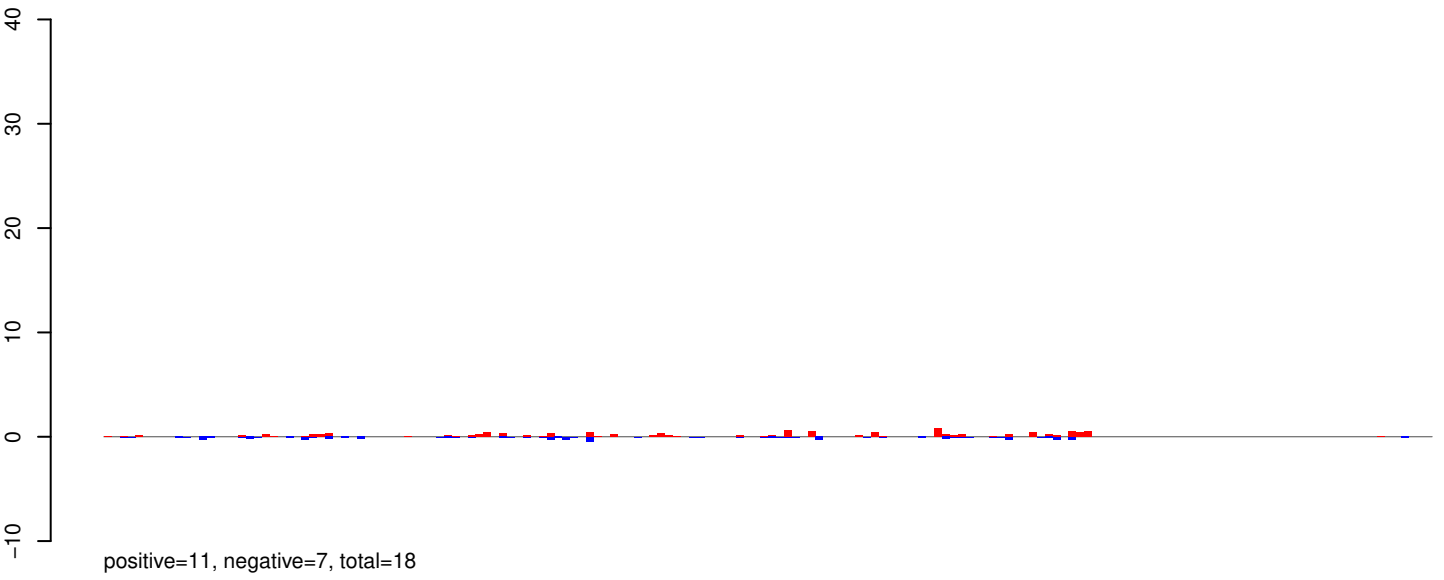


width=326, length=6065, virus@MH188054.1-Culex_Y_virus_strain_CYV_Kern_partial_genome:1-6065

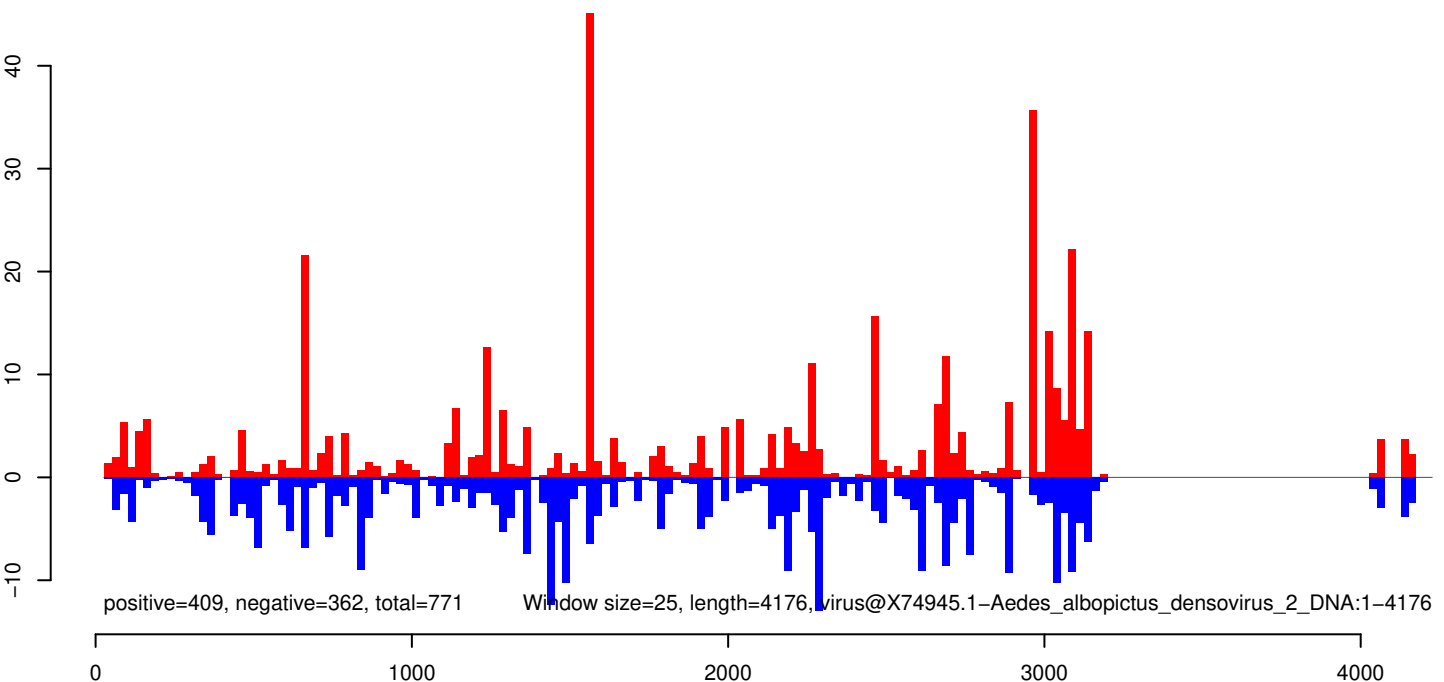
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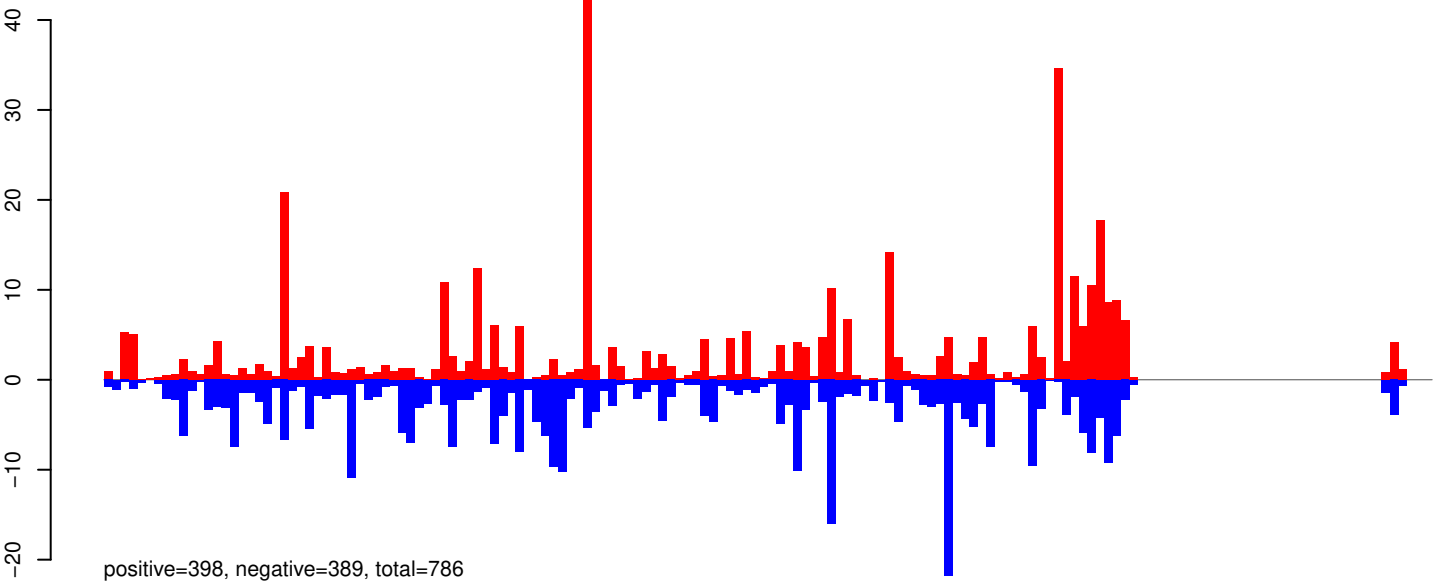
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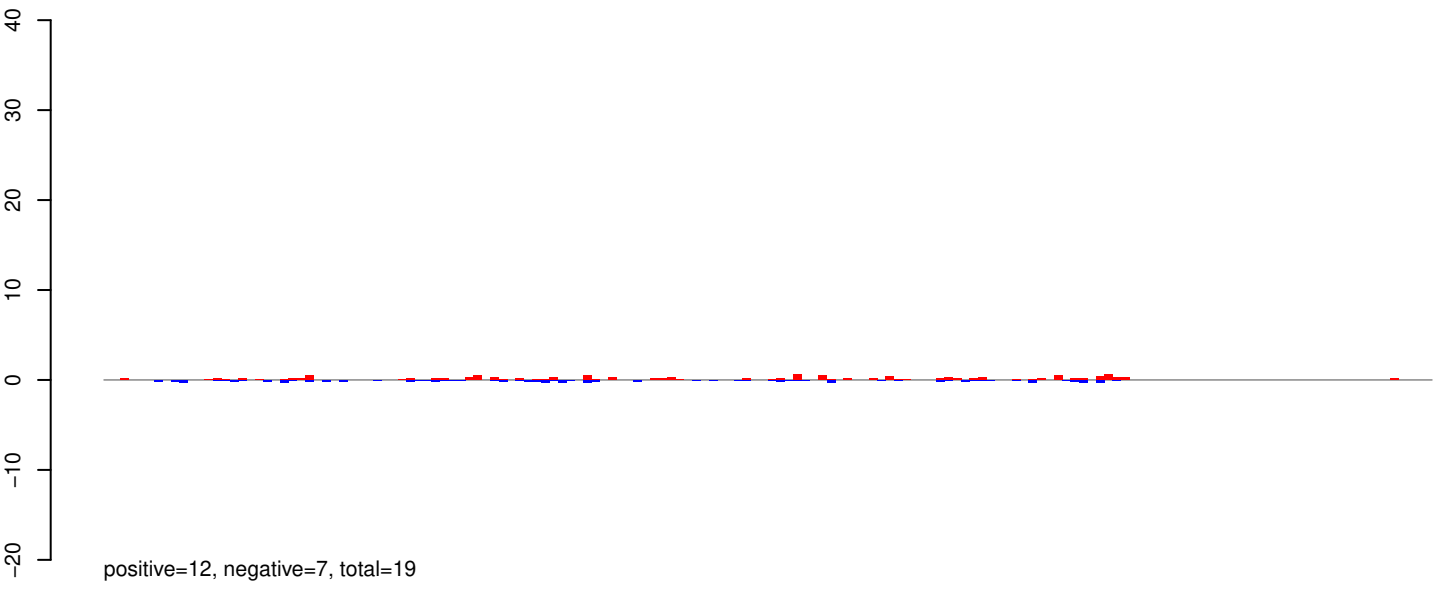
AeAeg_Aag2_BTV2_AK.rep



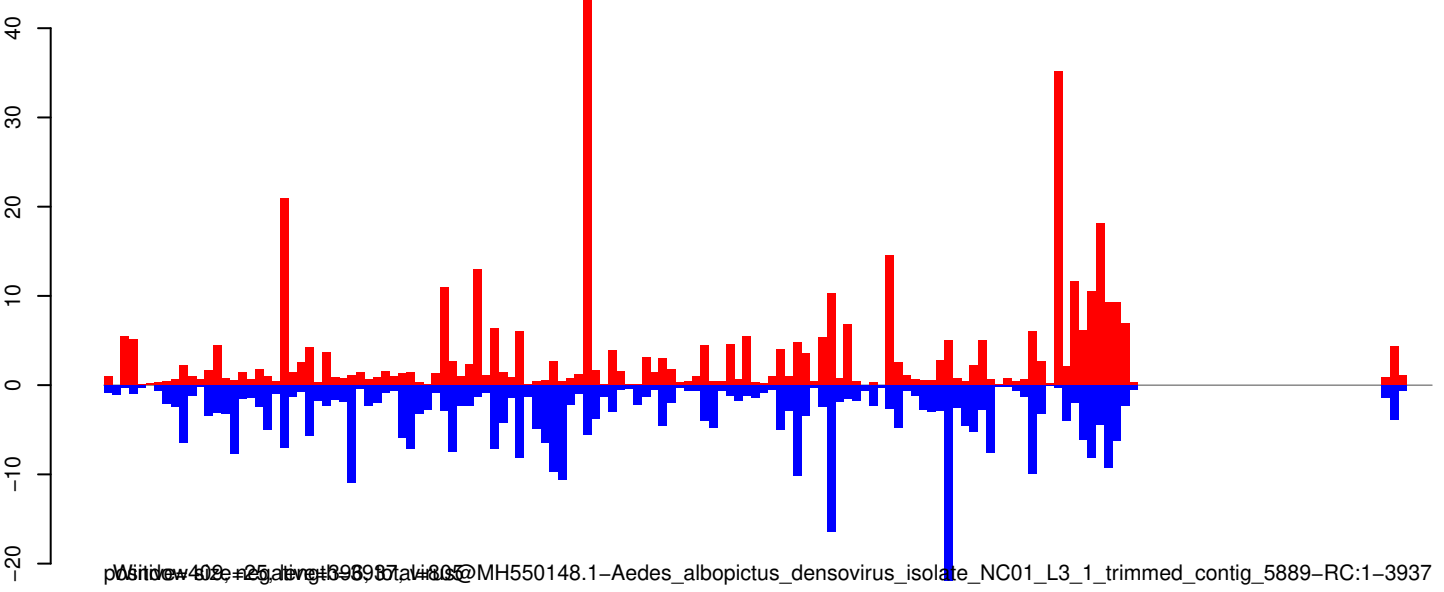
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



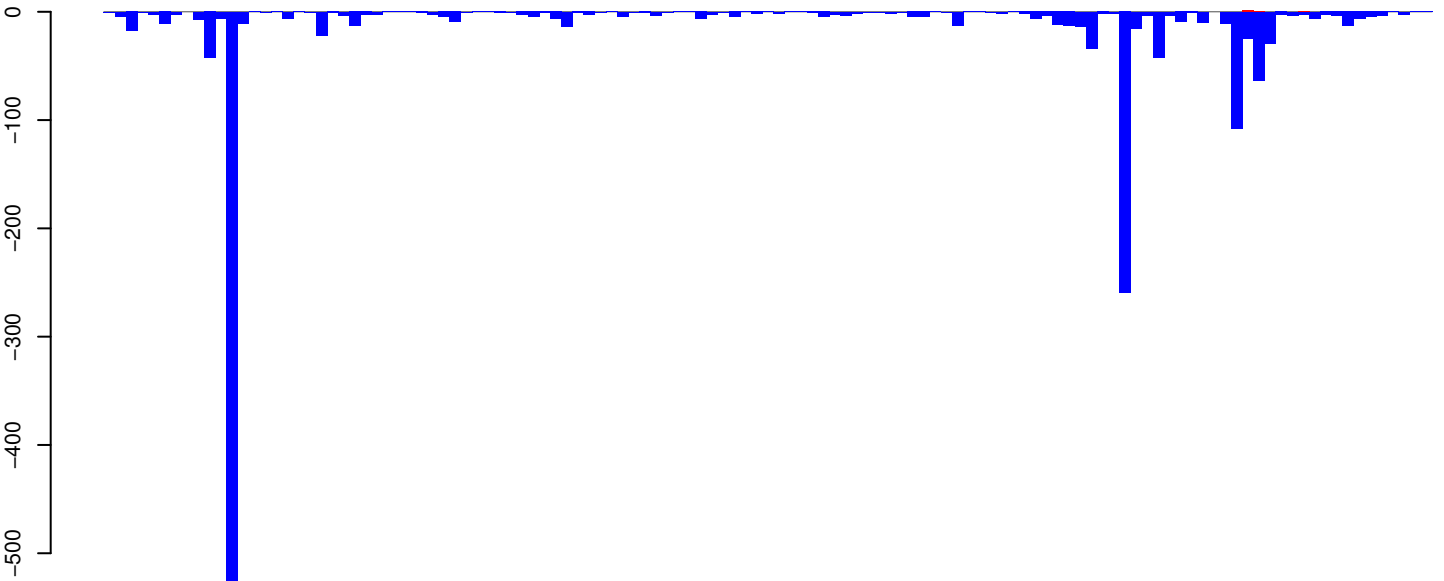
0 1000 2000 3000 4000

AeAeg_Aag2_BTV2_AK.18_23.rep



positive=1, negative=38, total=38

AeAeg_Aag2_BTV2_AK.24_35.rep



positive=3, negative=1557, total=1561

AeAeg_Aag2_BTV2_AK.rep

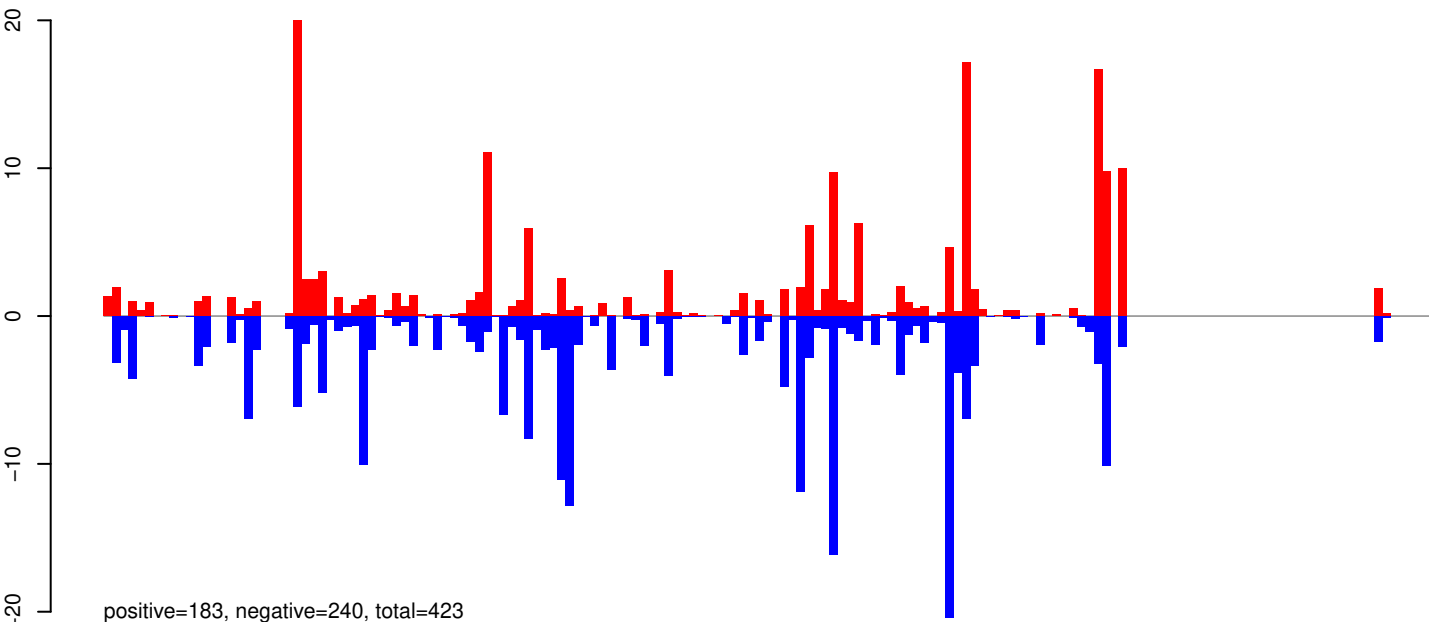


positive=4, negative=1595, total=1599

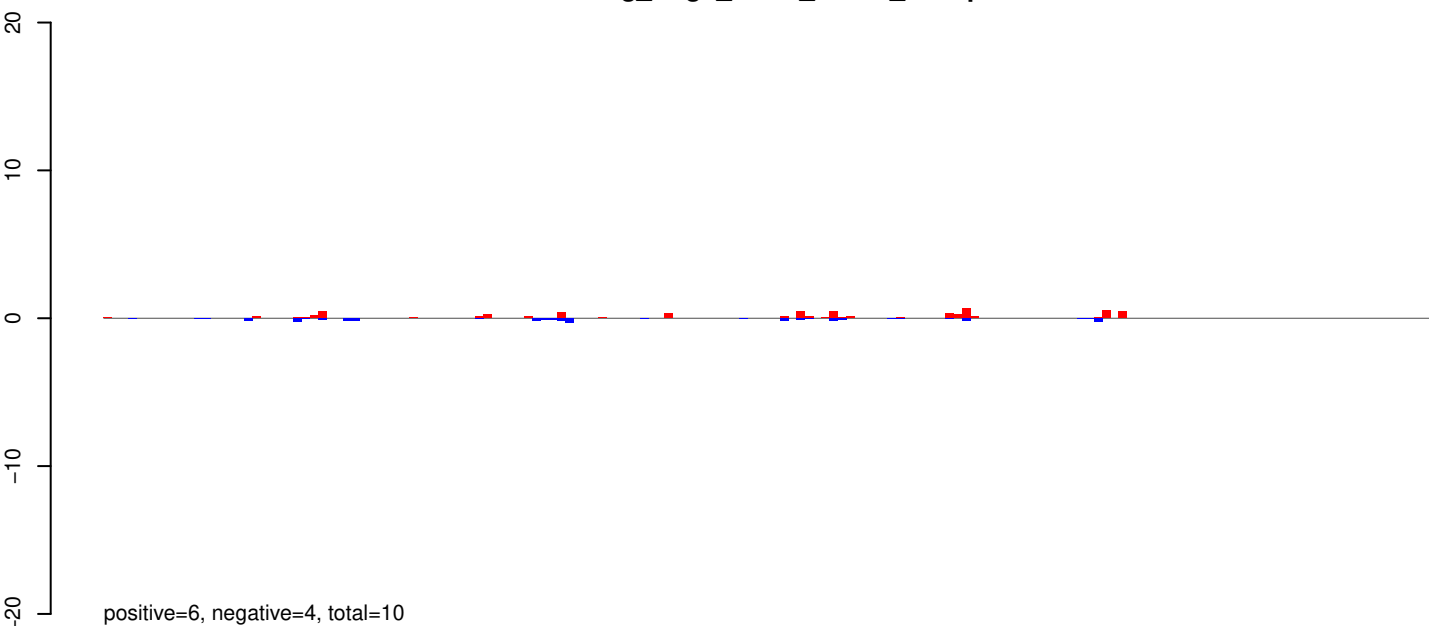
Window size=25, length=2965, virus@AY347953.2-Aedes_aegypti_genomic_sequence:1-2965

0 500 1000 1500 2000 2500 3000

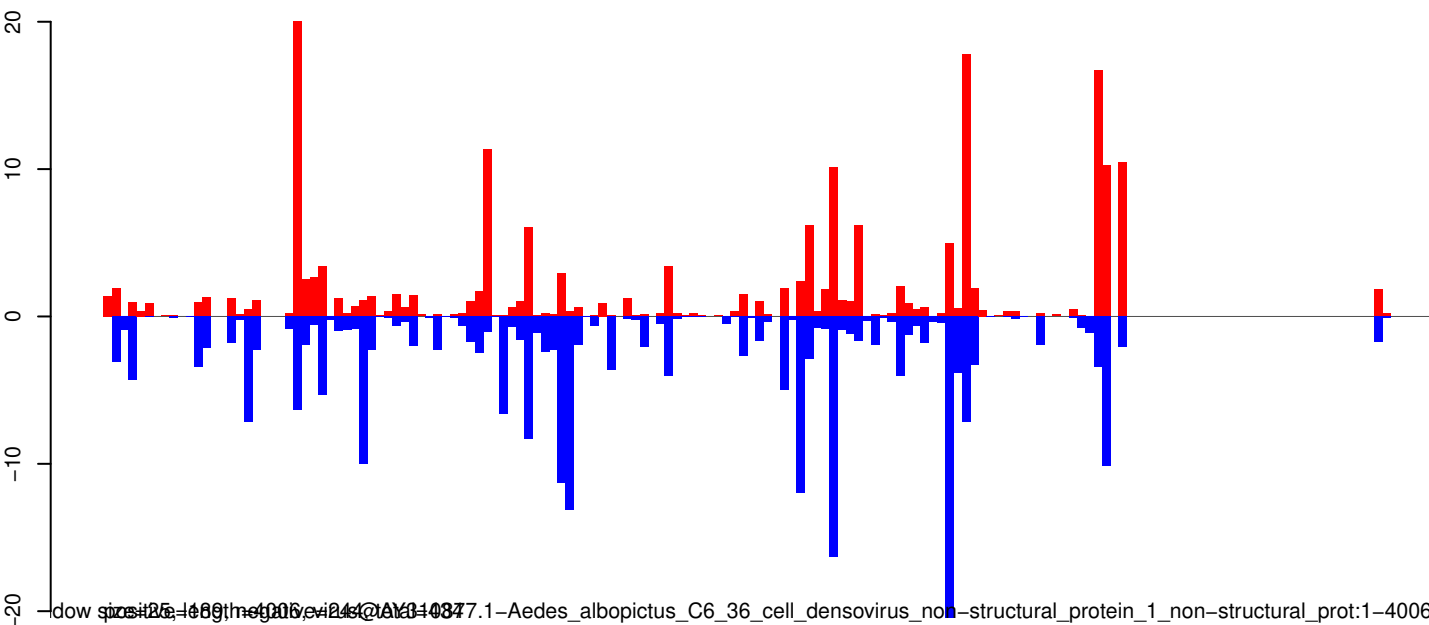
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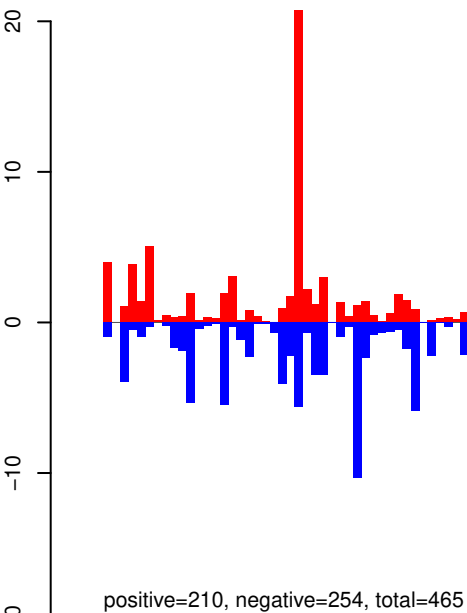
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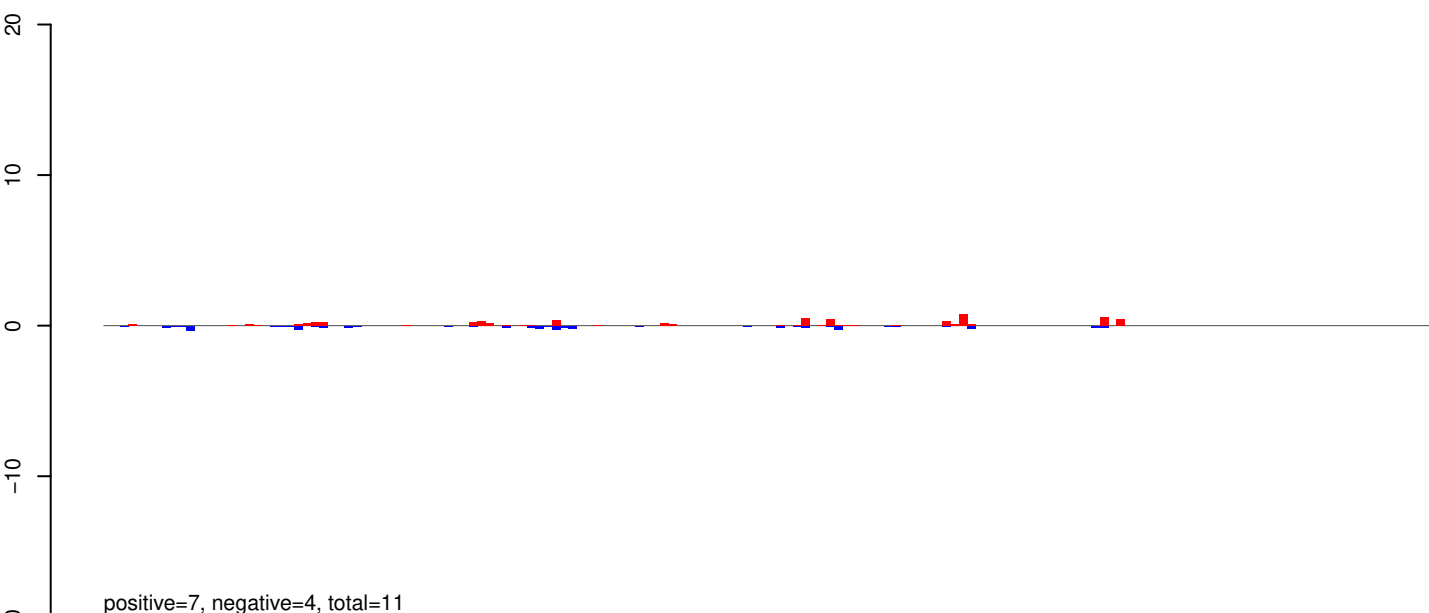
AeAeg_Aag2_BTV2_AK.rep



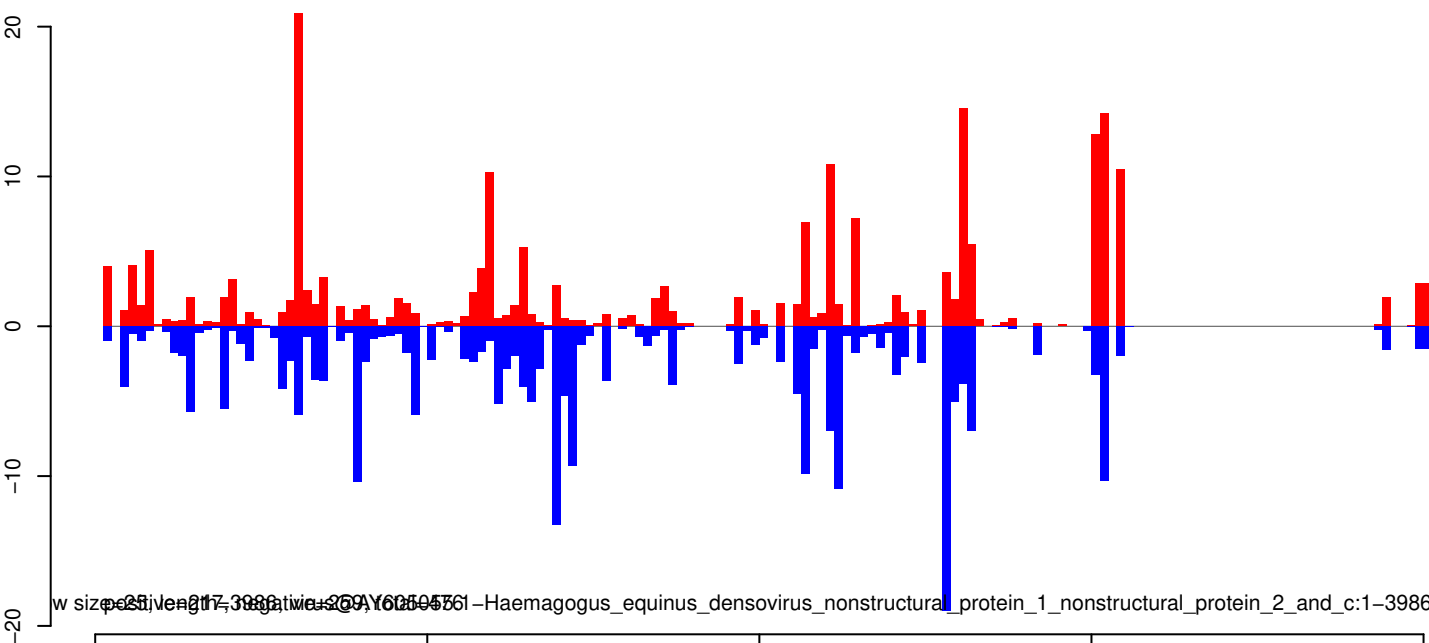
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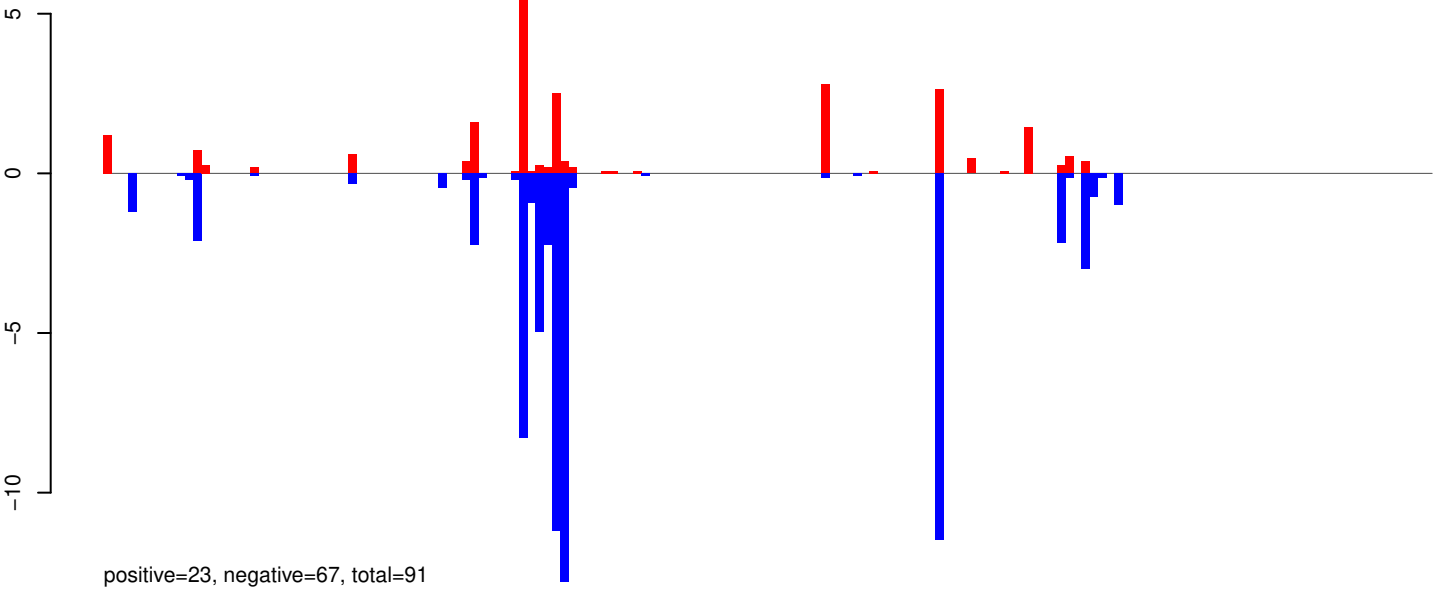
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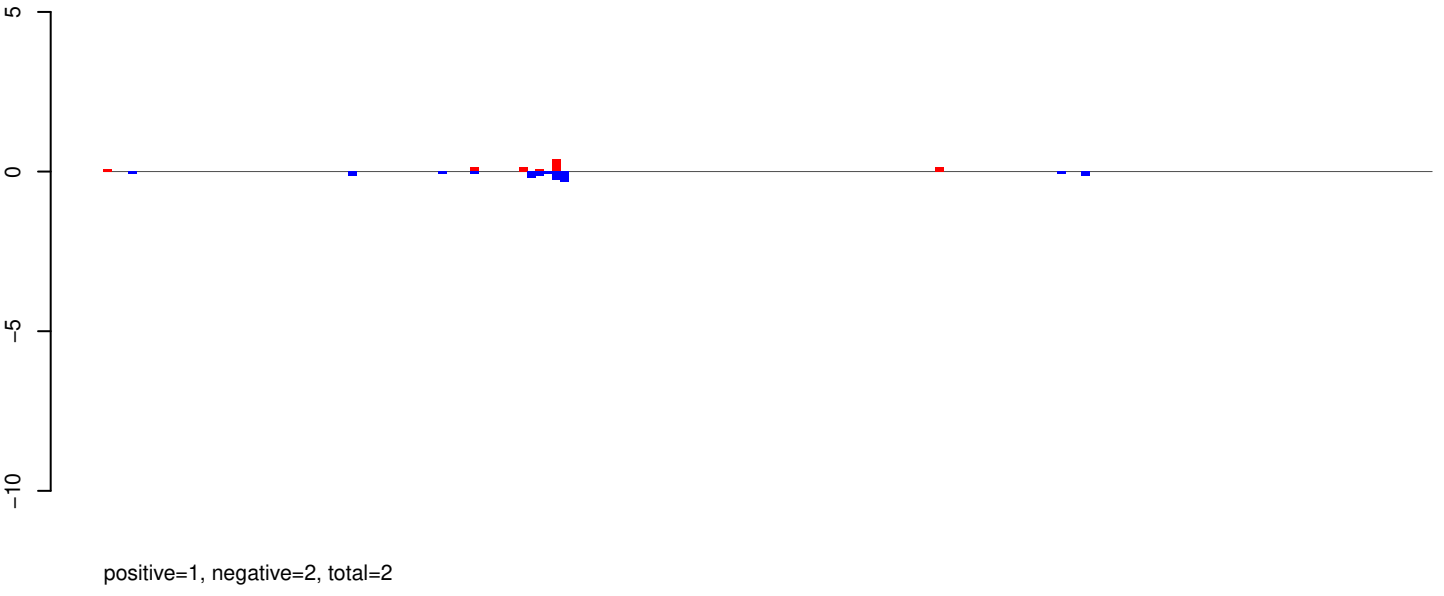
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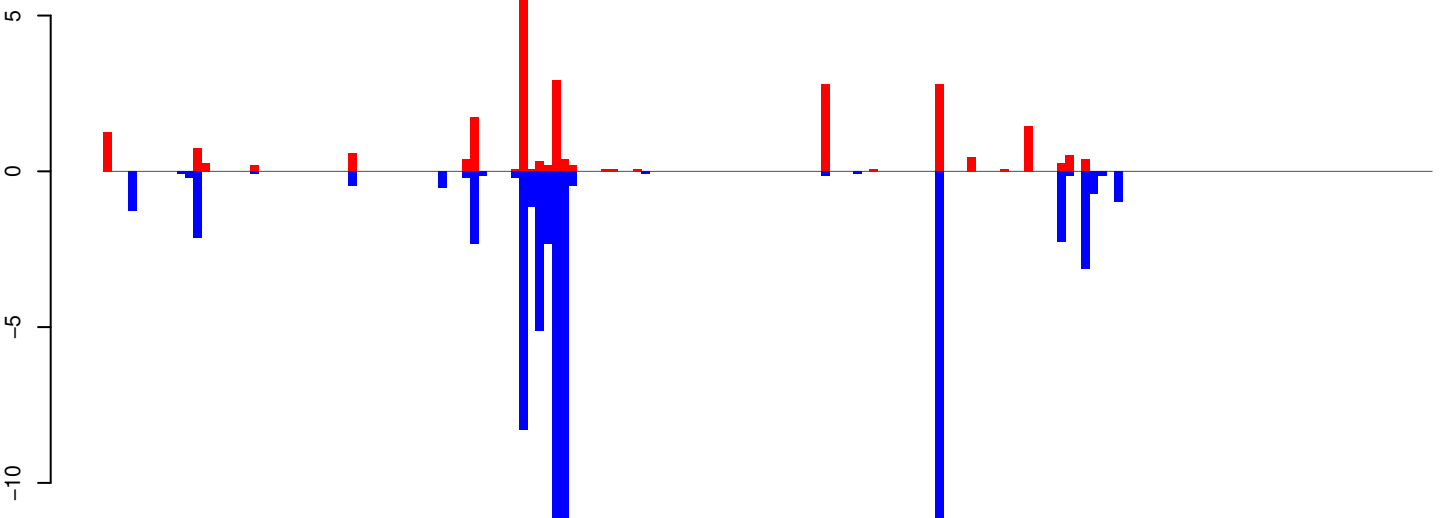
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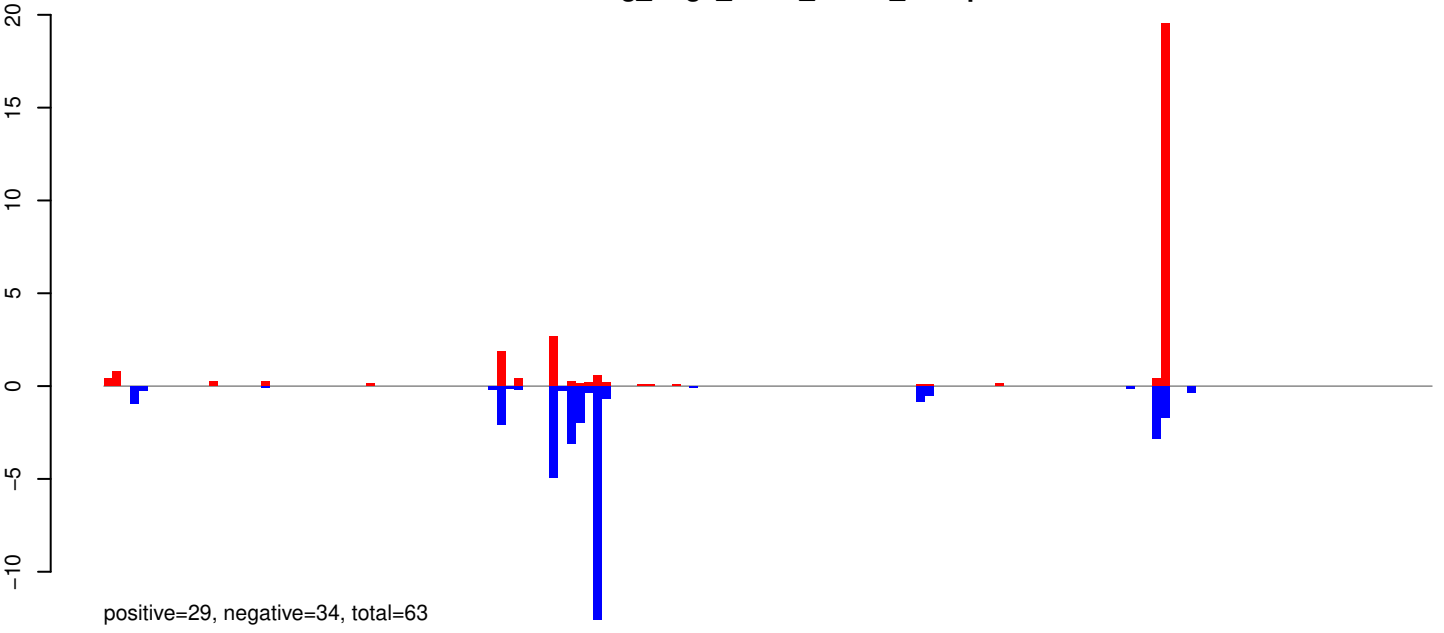
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AeAeg_Aag2_BTV2_AK.rep



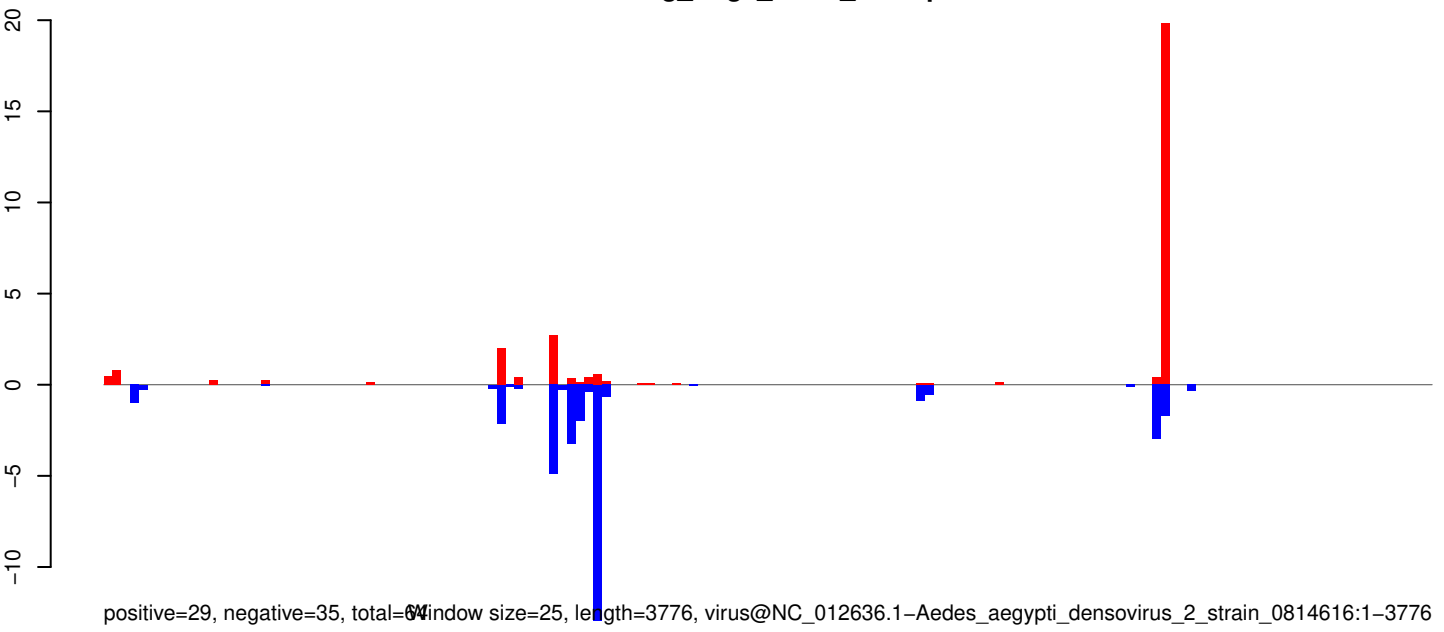
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

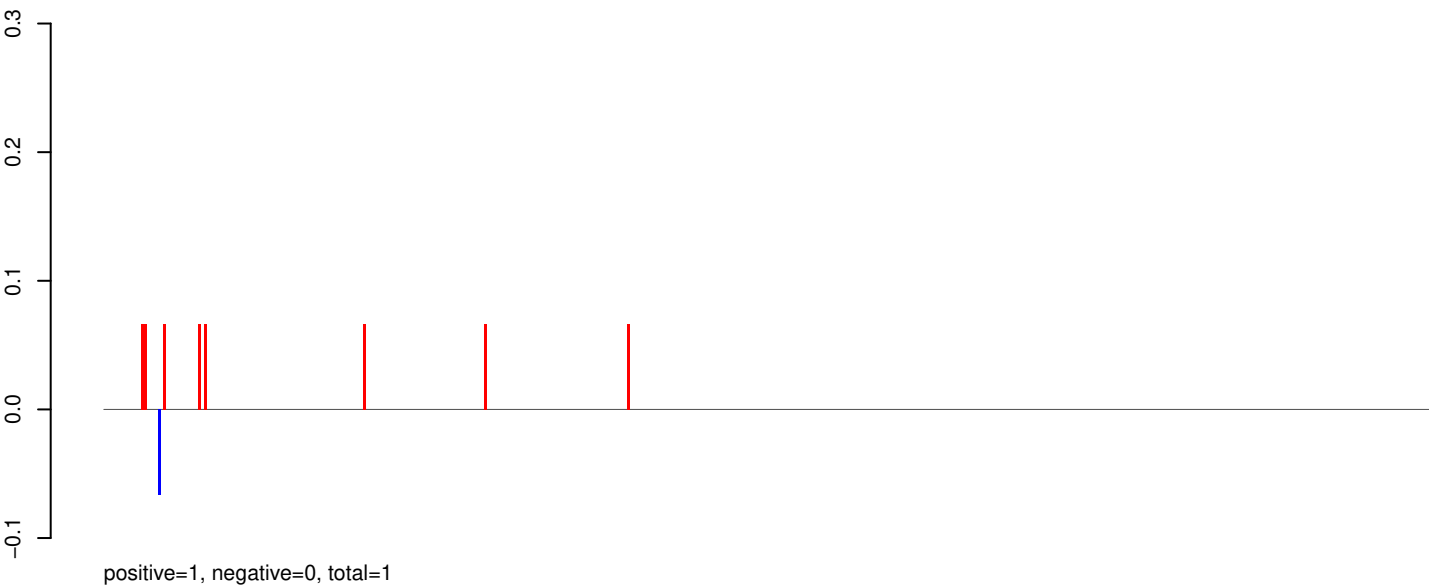


AeAeg_Aag2_BTV2_AK.rep

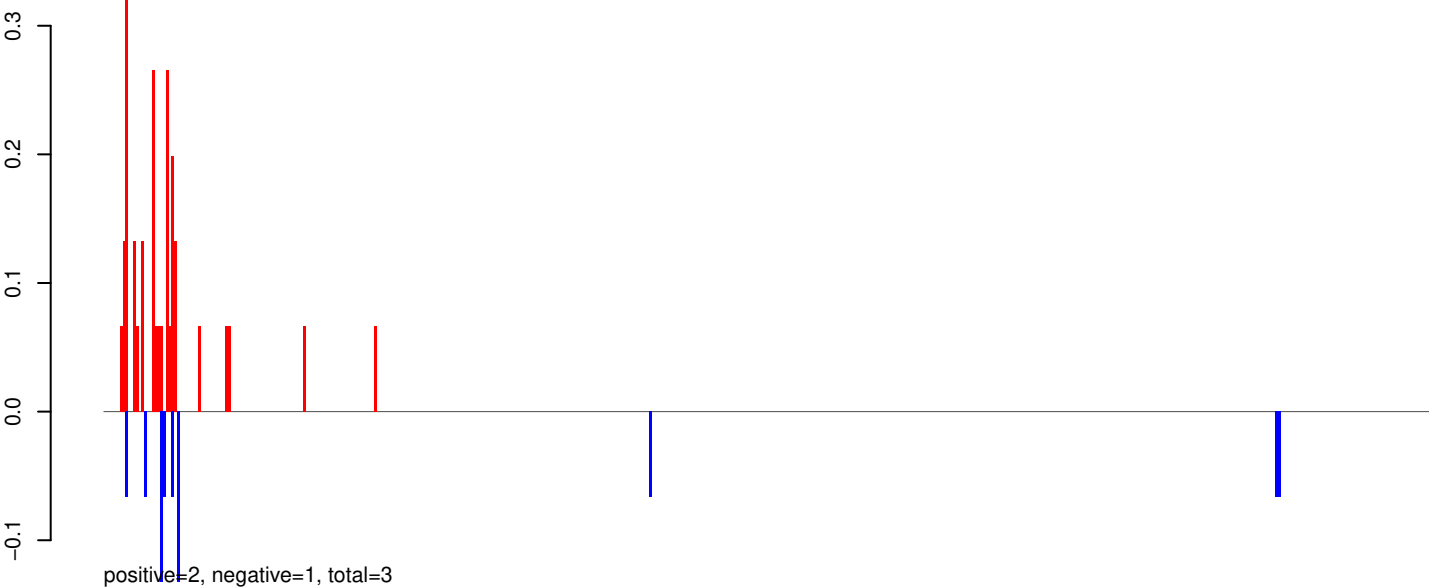


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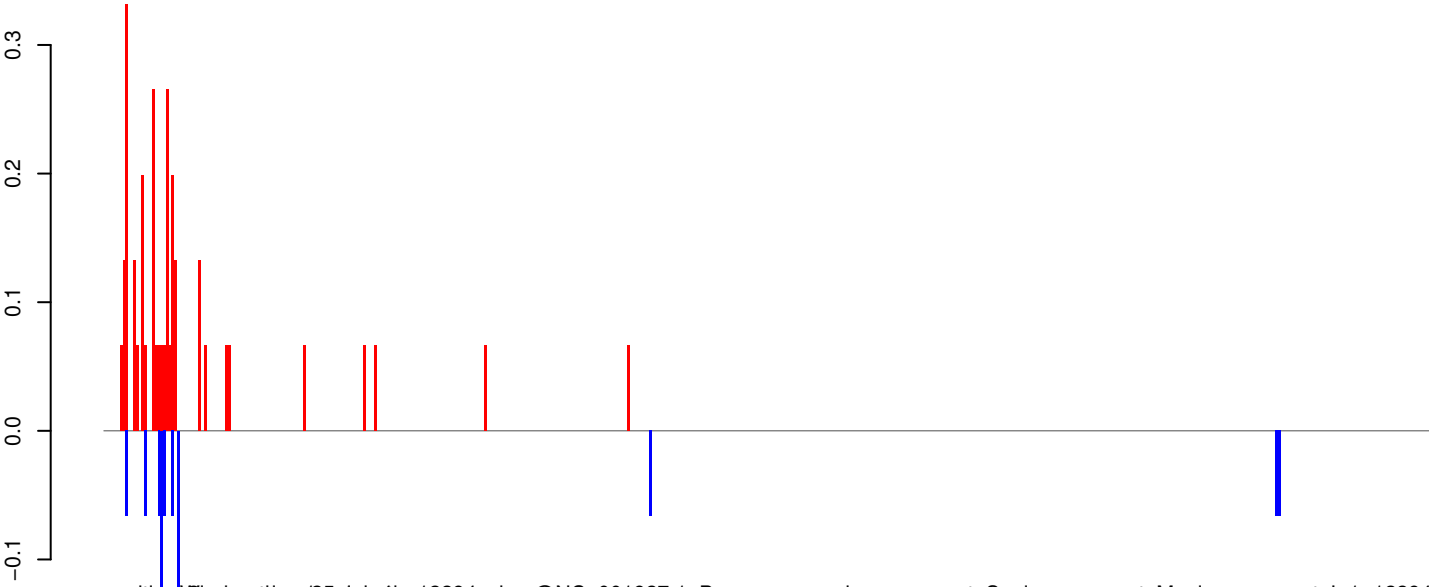
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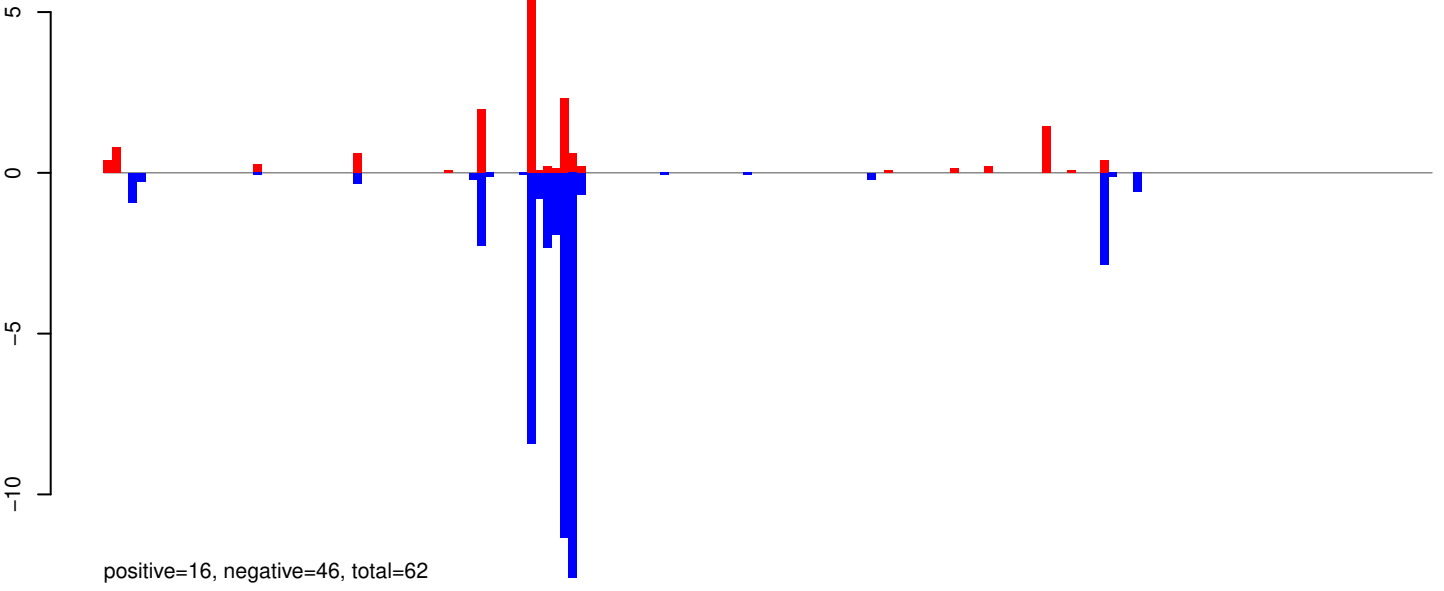
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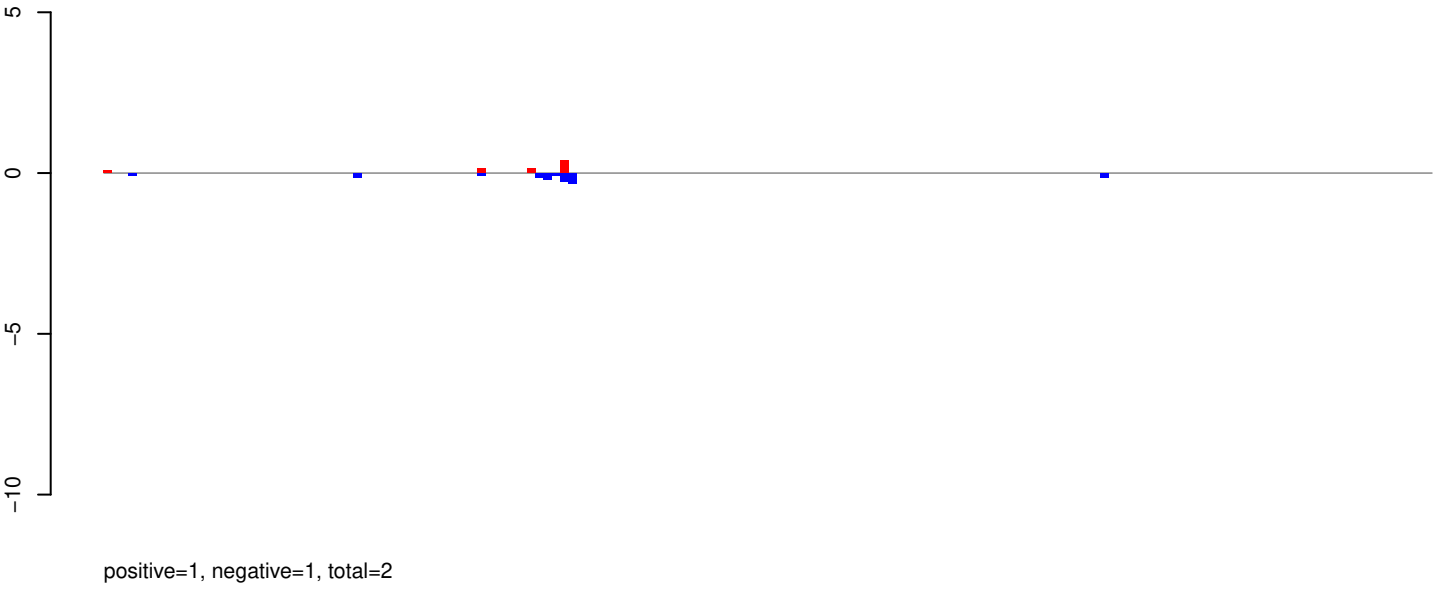
AeAeg_Aag2_BTV2_AK.rep



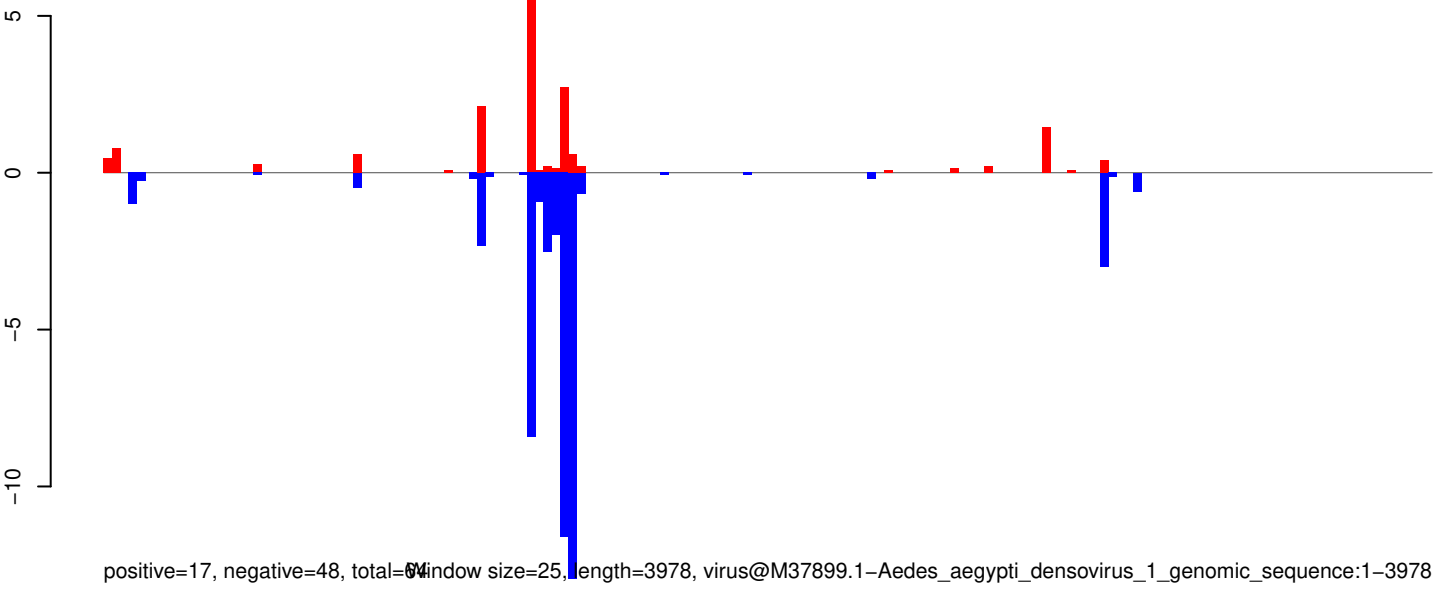
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AeAeg_Aag2_BTV2_AK.24_35.rep

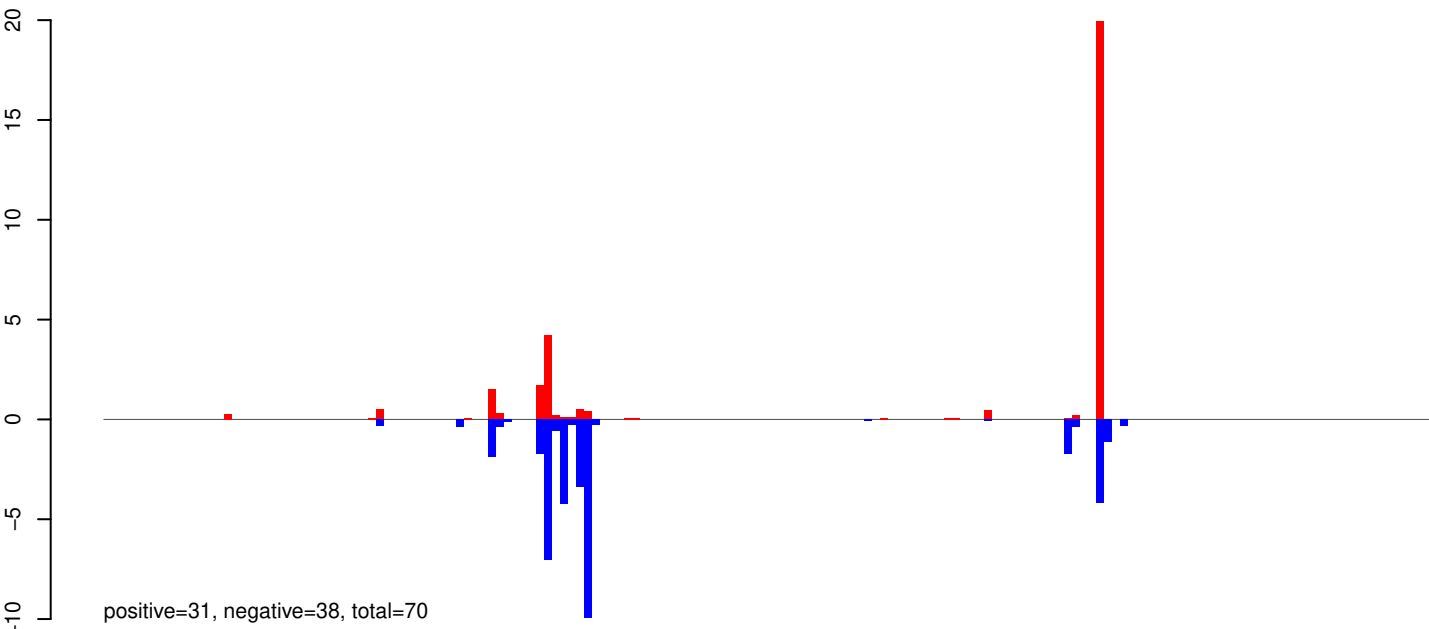


AeAeg_Aag2_BTV2_AK.rep

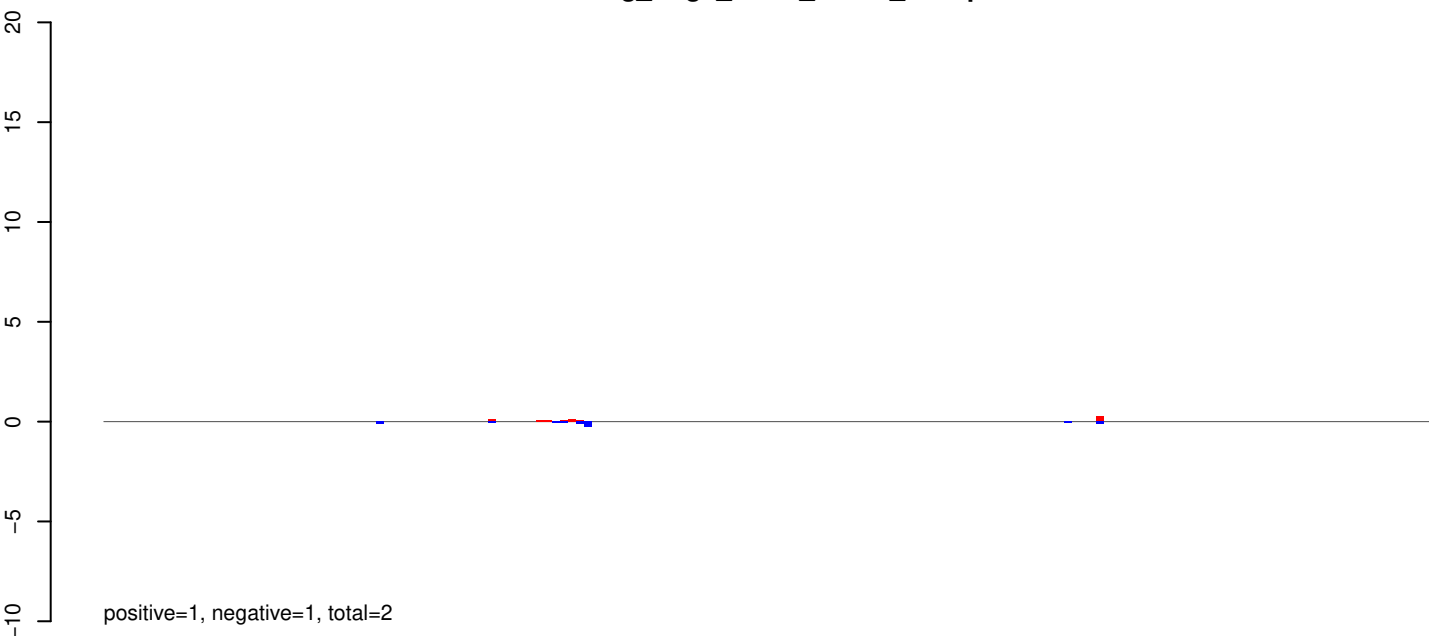


Window size=25, length=3978, virus@M37899.1-Aedes_aegypti_densovirus_1_genomic_sequence:1-3978

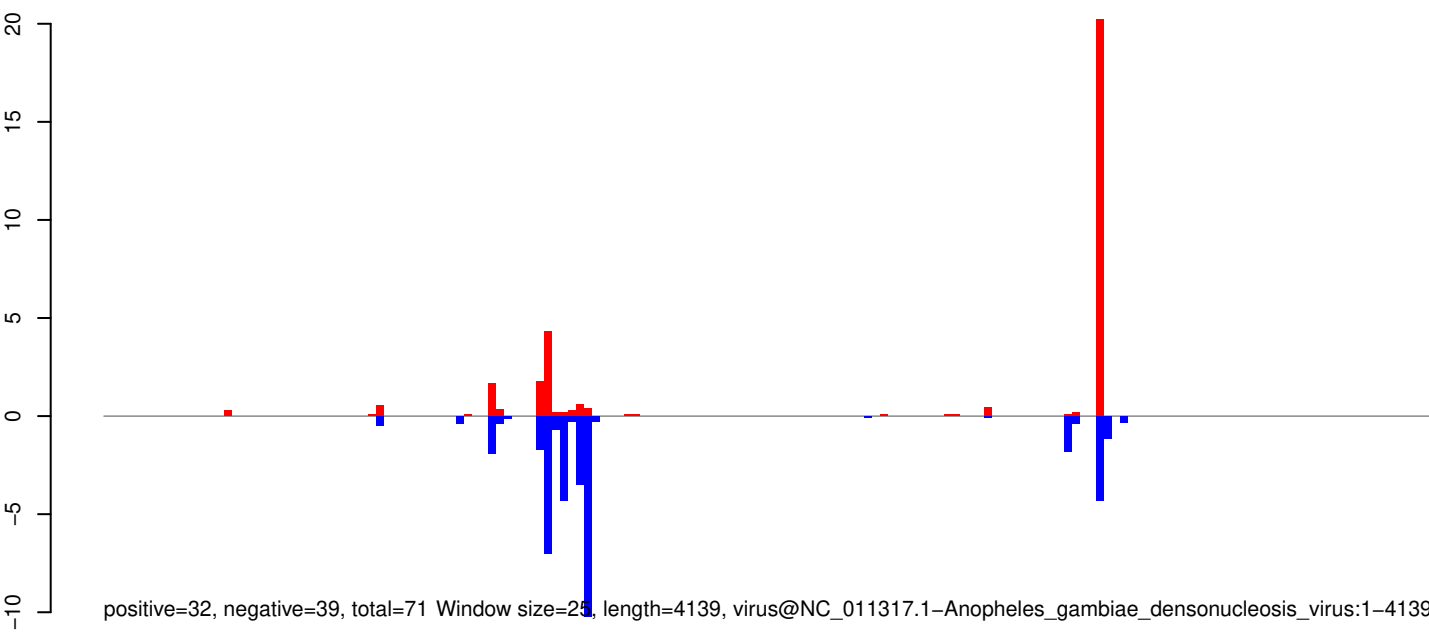
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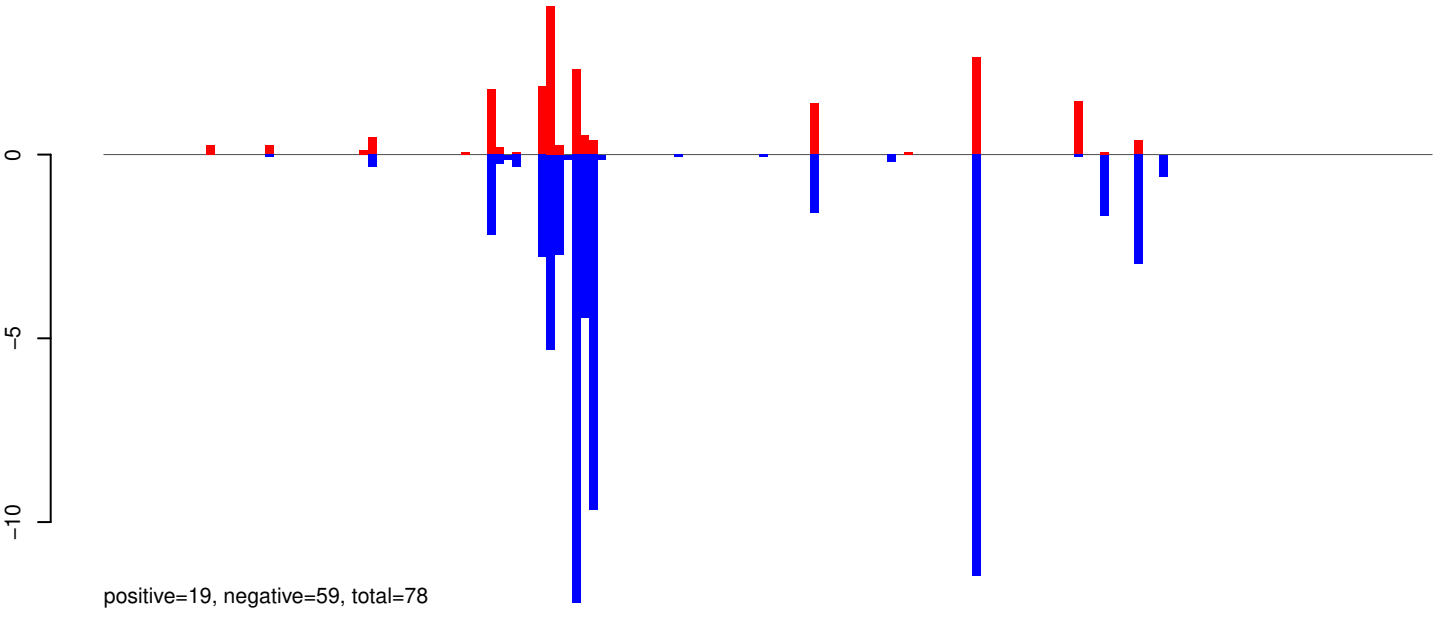
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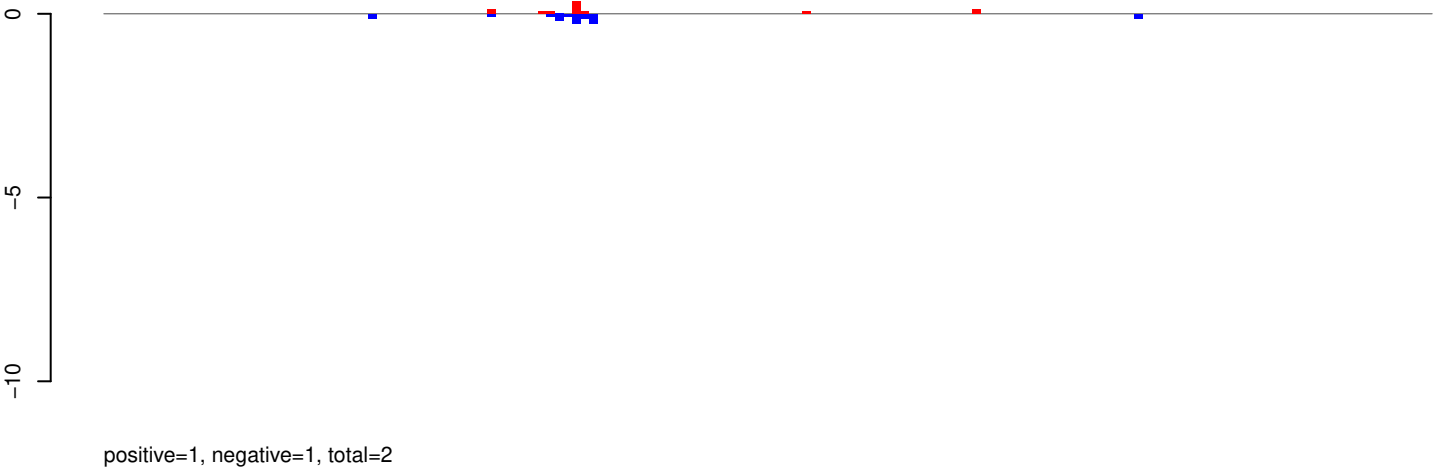
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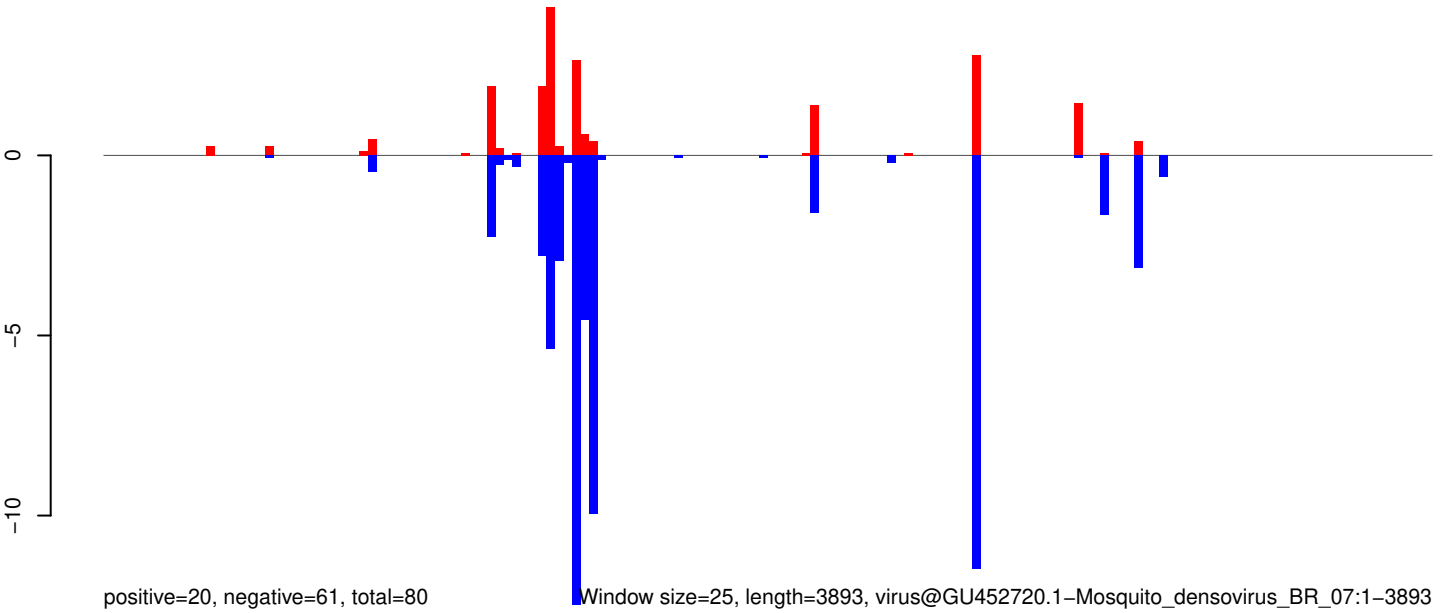
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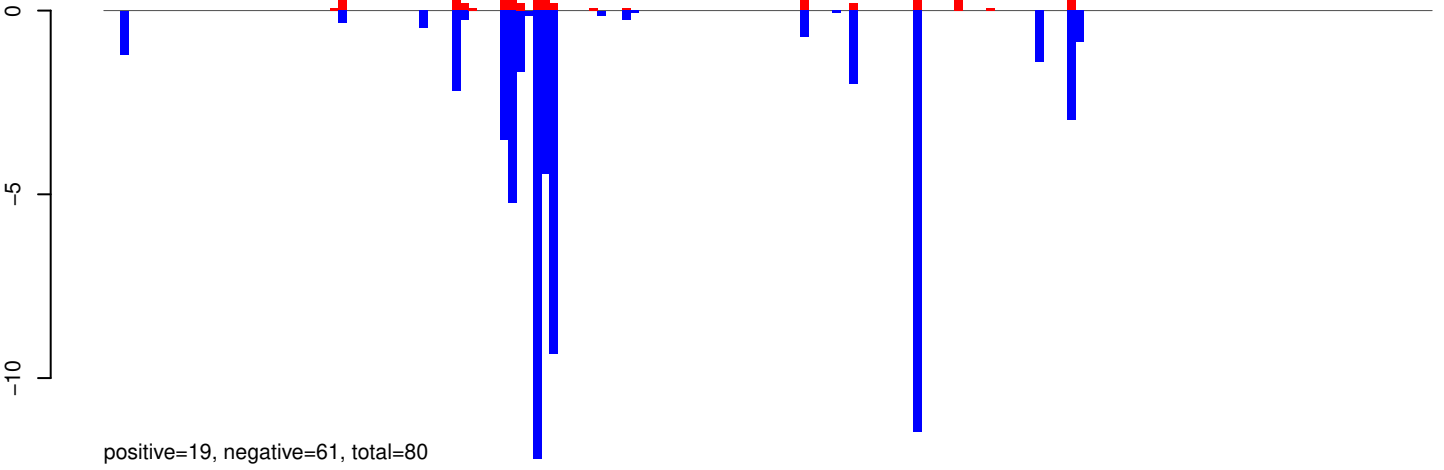
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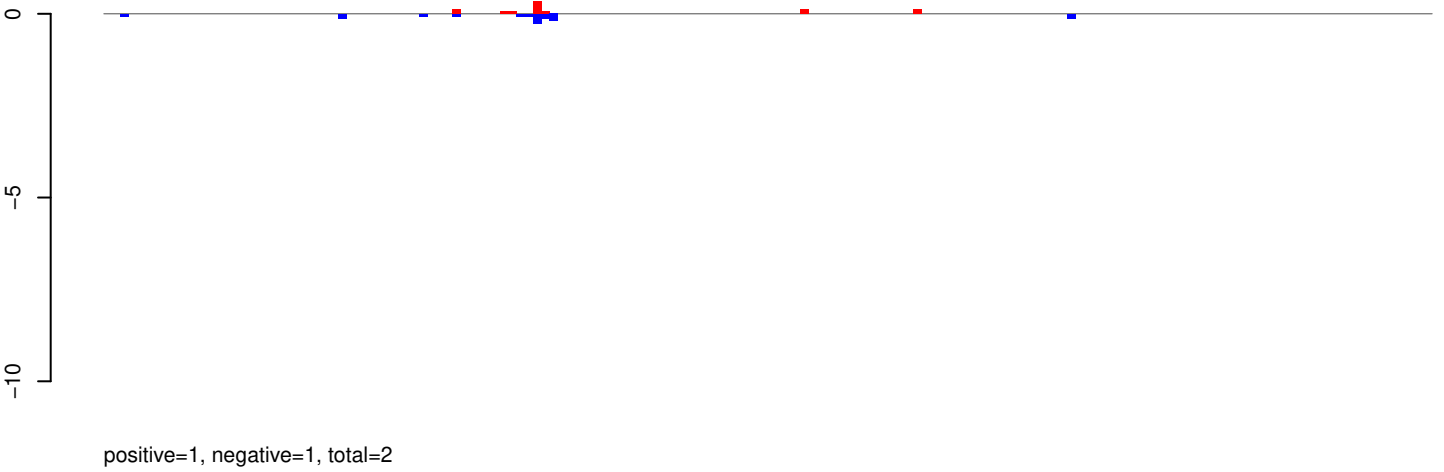
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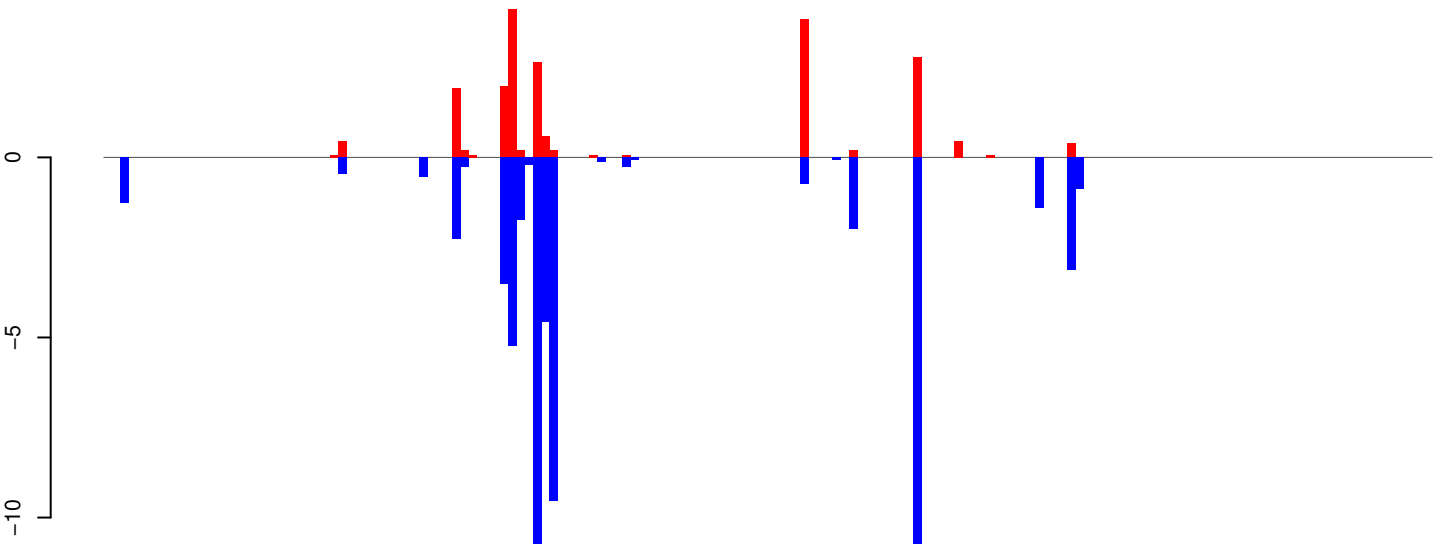
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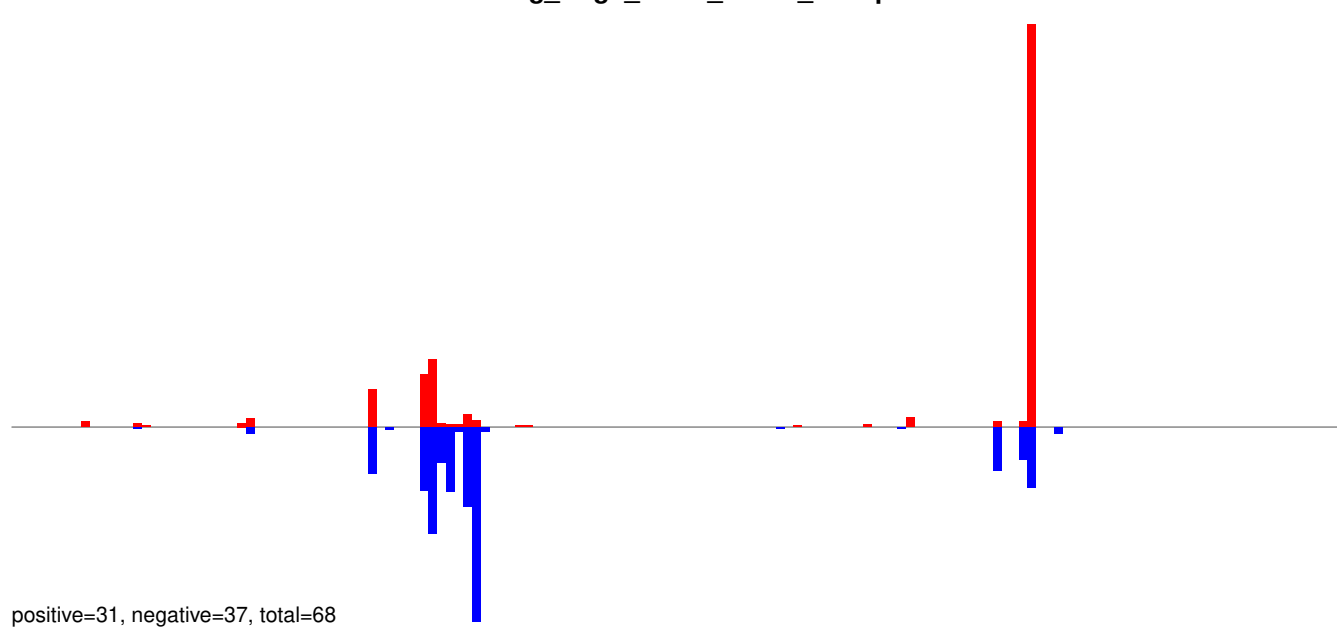
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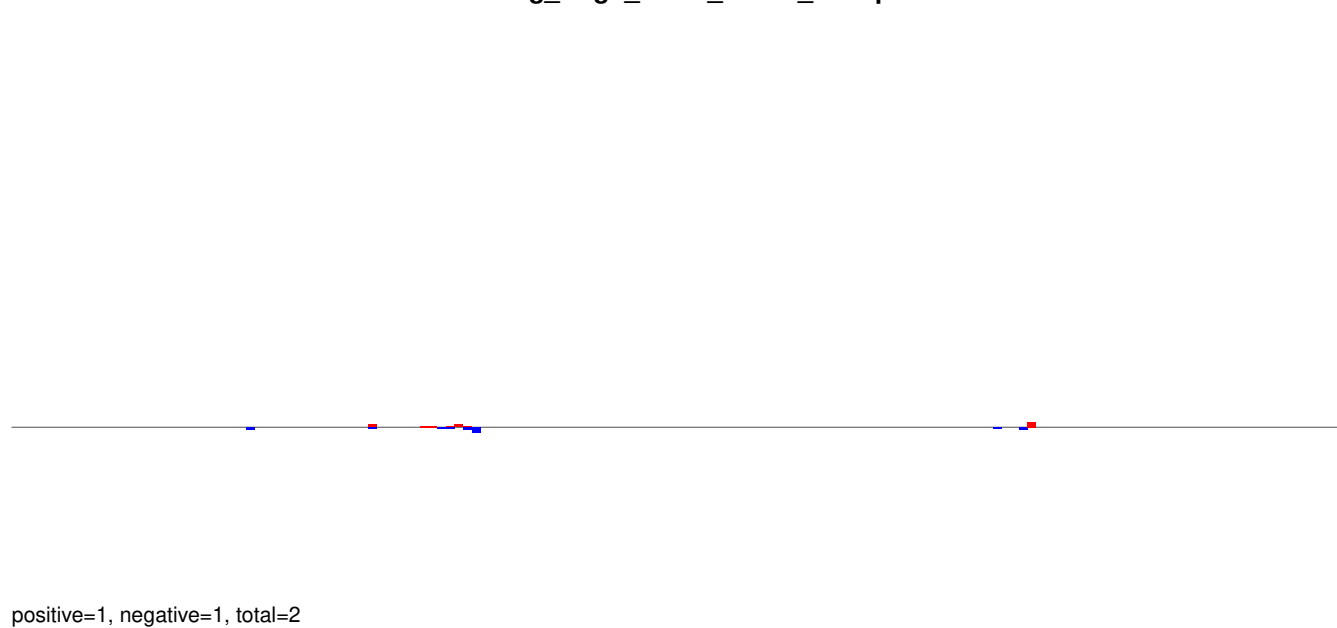
AeAeg_Aag2_BTV2_AK.rep



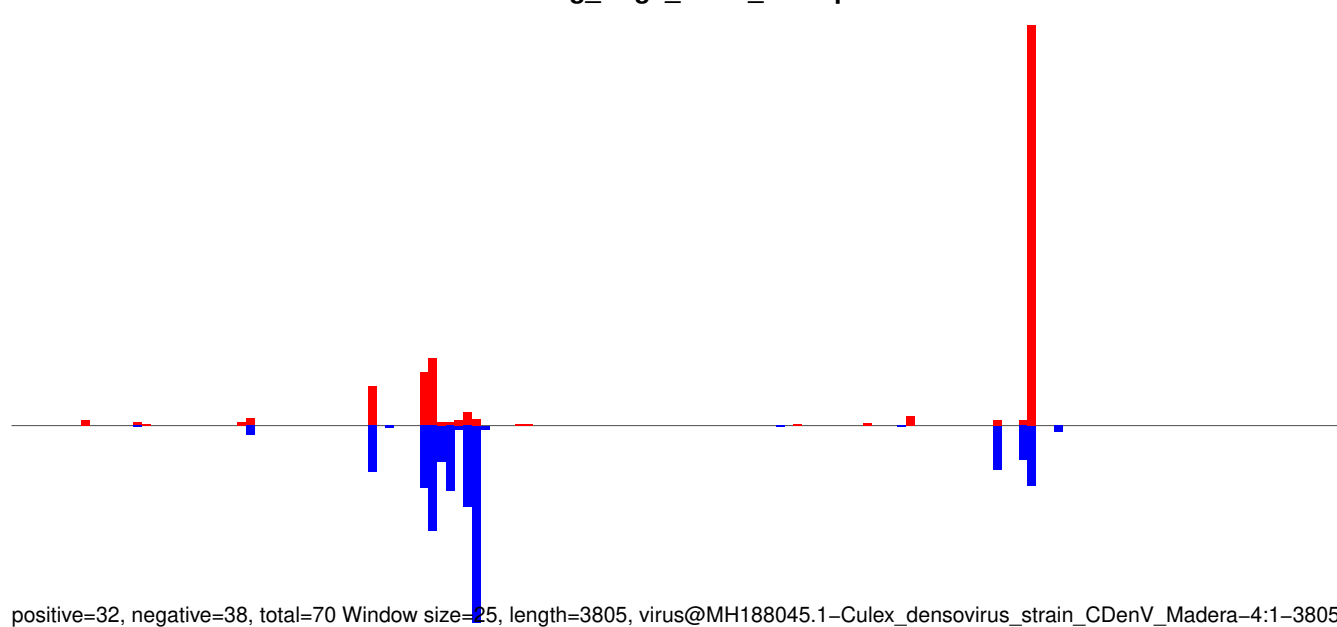
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

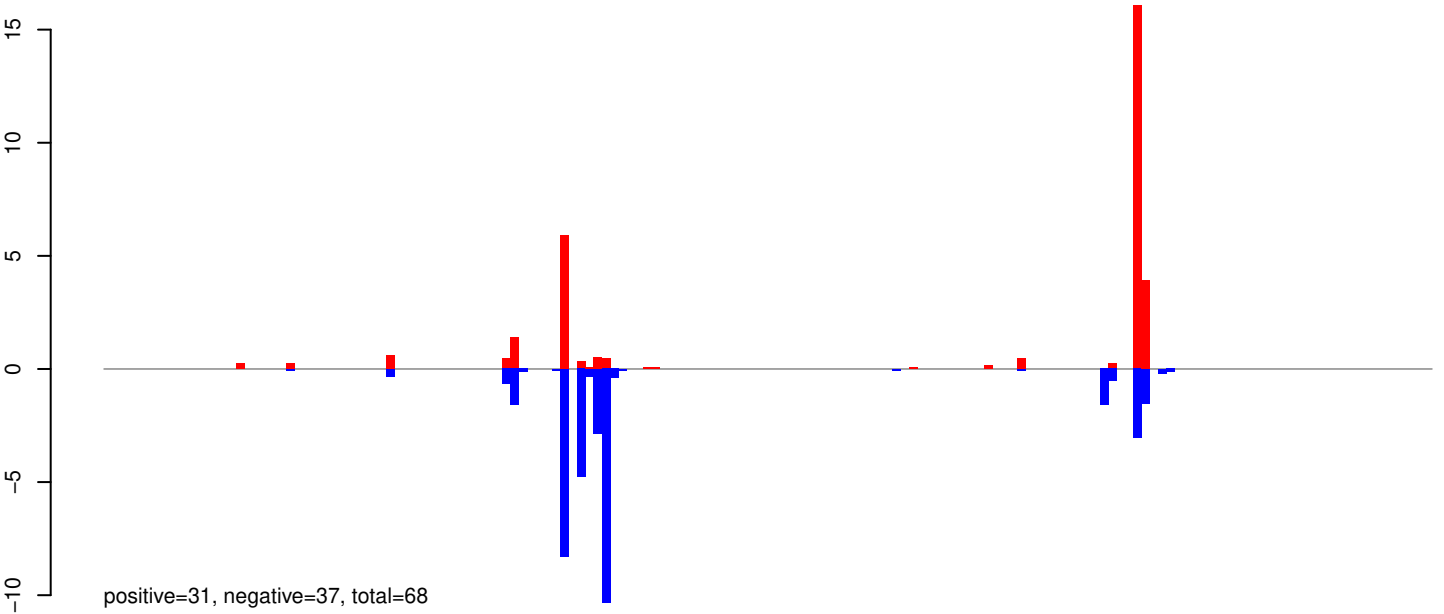


AeAeg_Aag2_BTV2_AK.rep

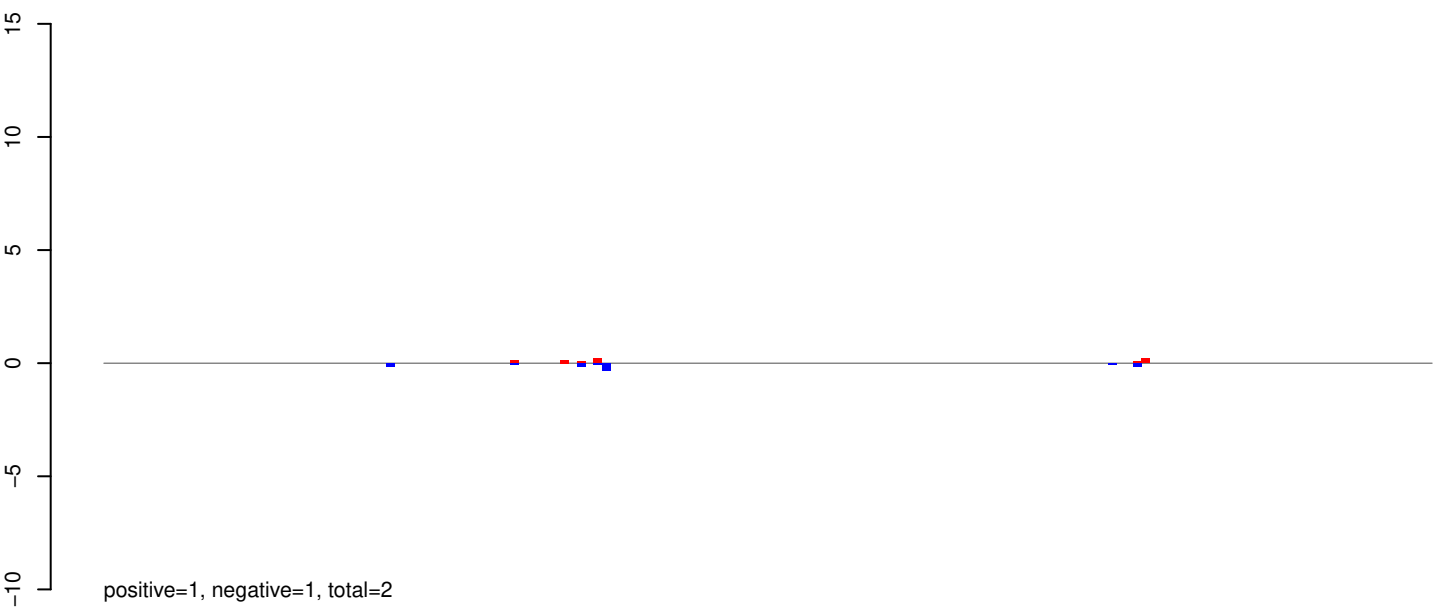


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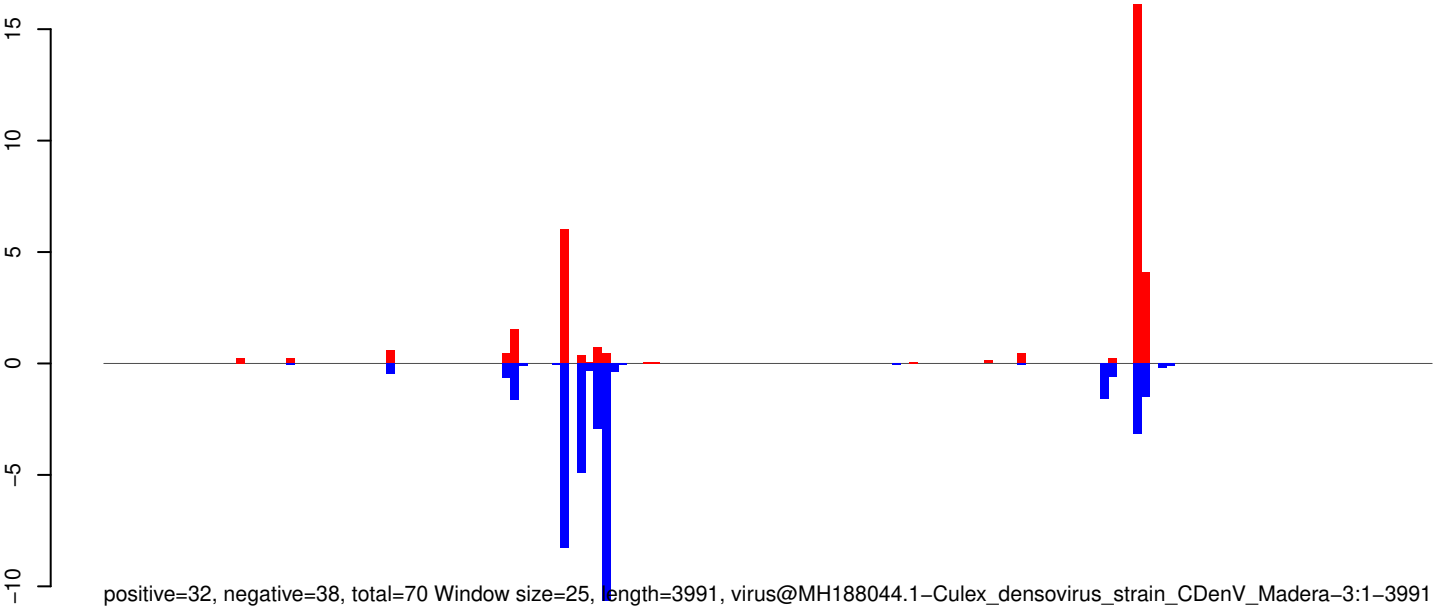
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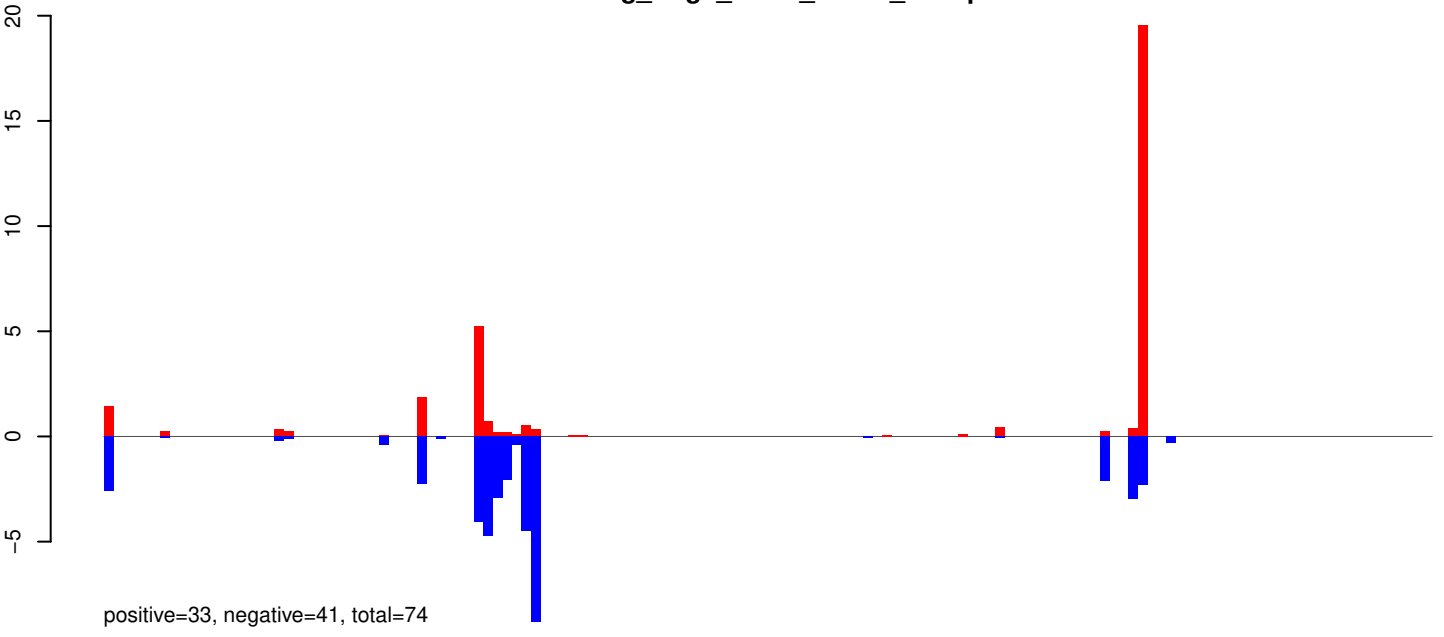
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AeAeg_Aag2_BTV2_AK.rep



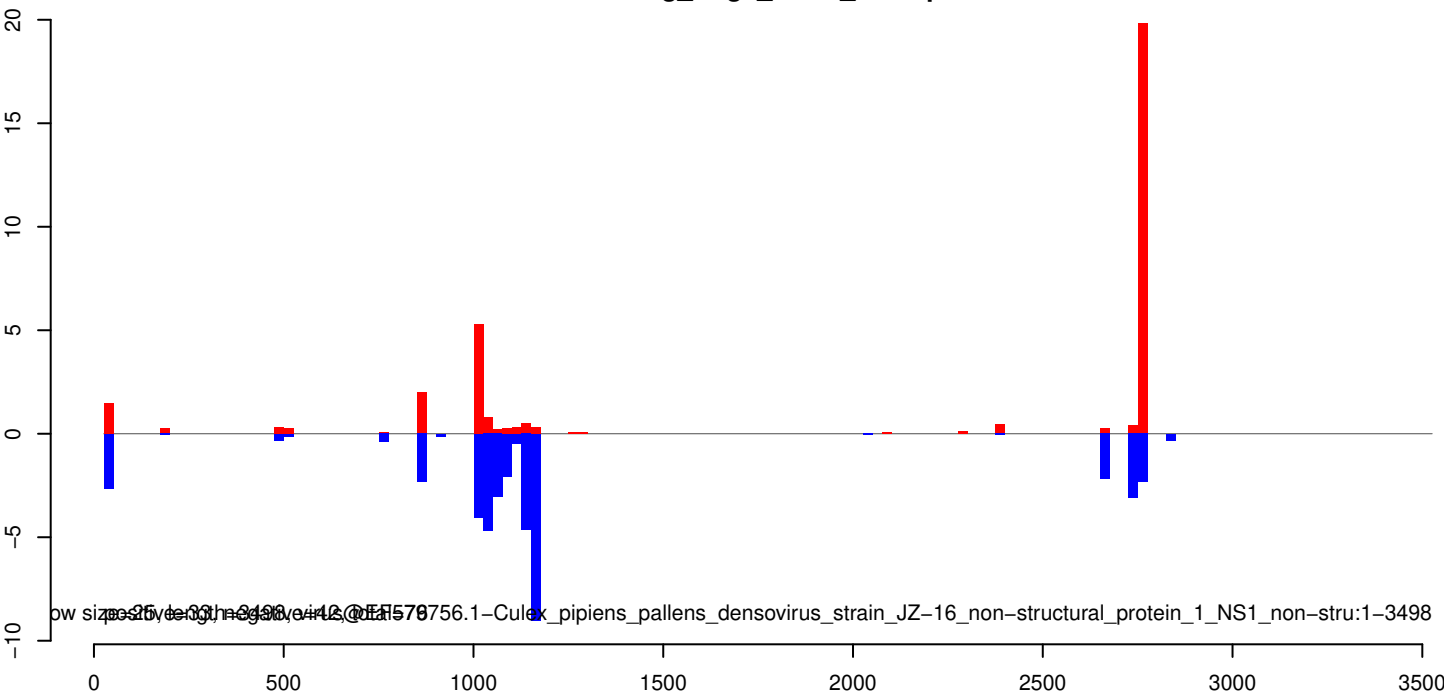
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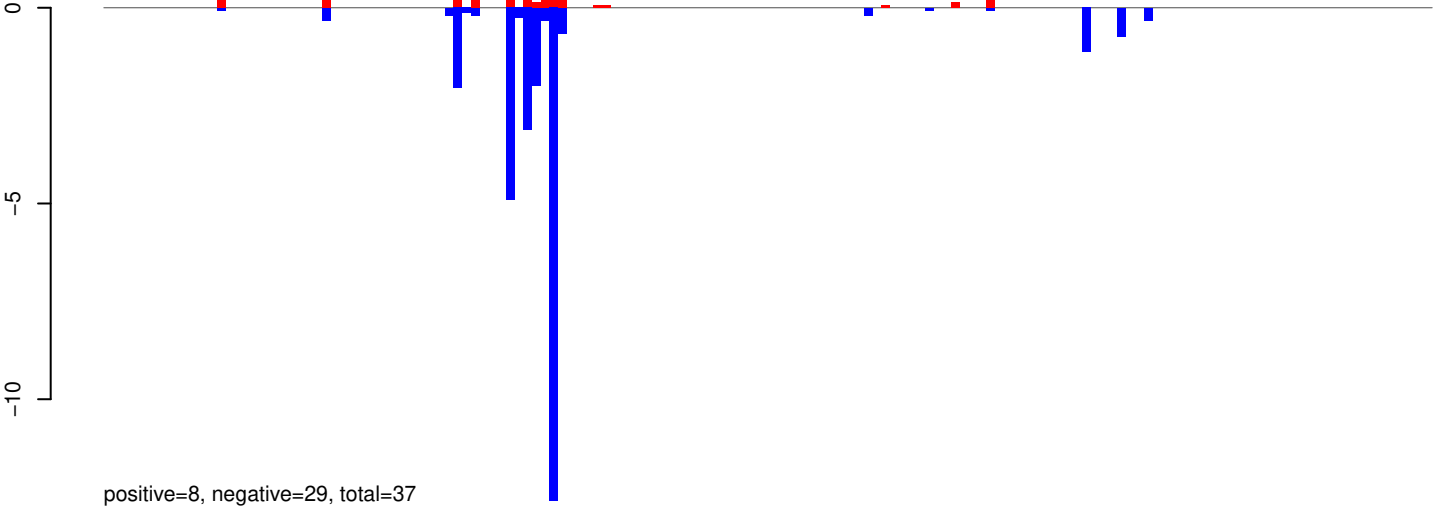
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AeAeg_Aag2_BTV2_AK.rep



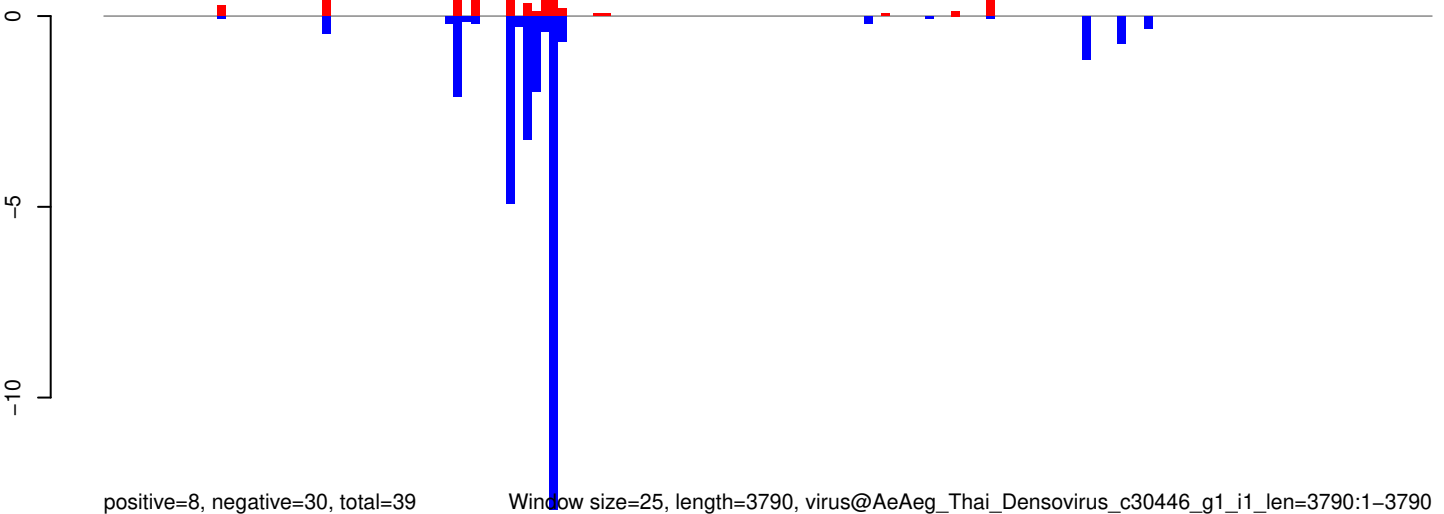
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AeAeg_Aag2_BTV2_AK.24_35.rep

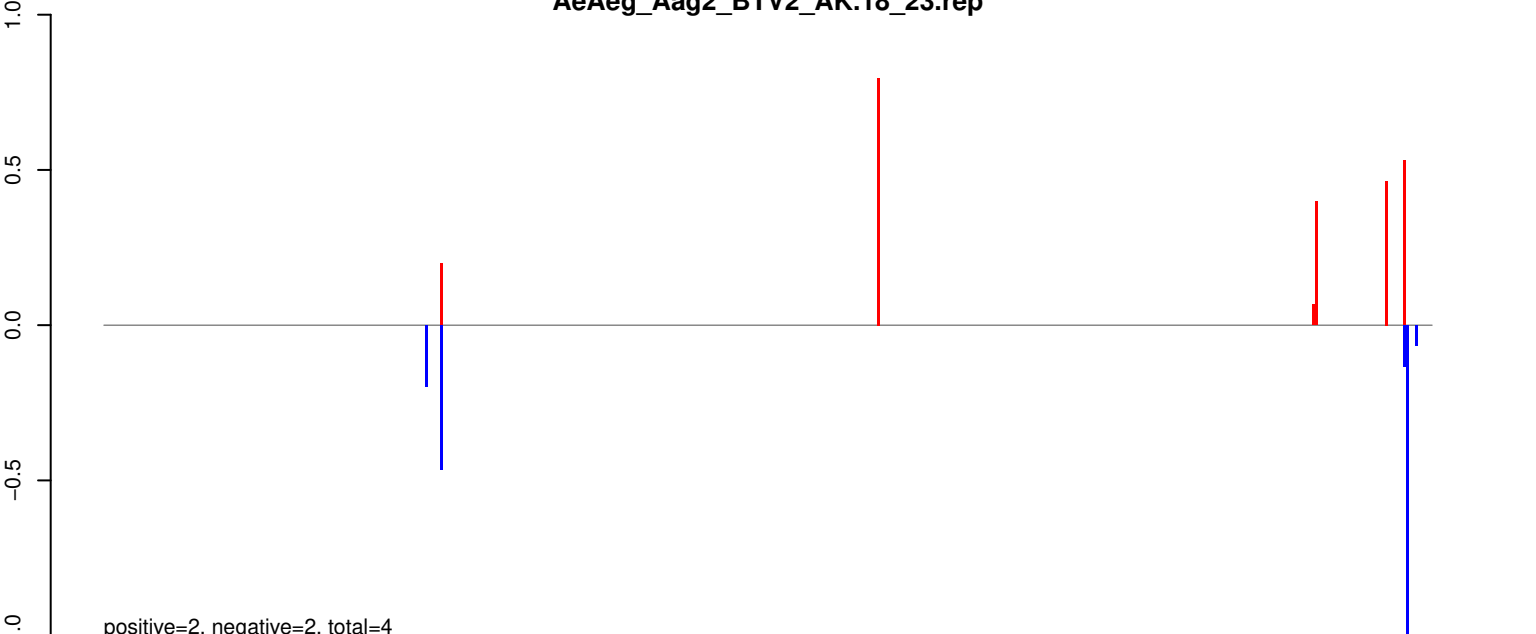


AeAeg_Aag2_BTV2_AK.rep



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

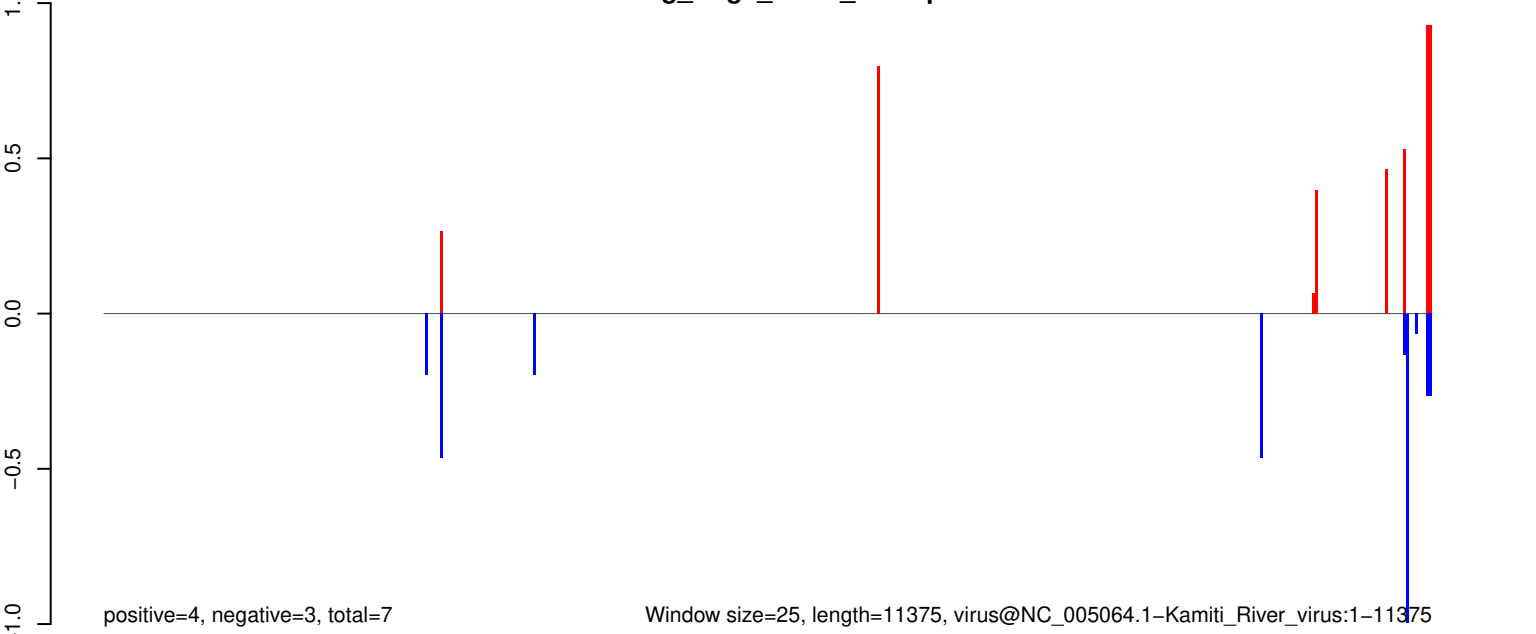
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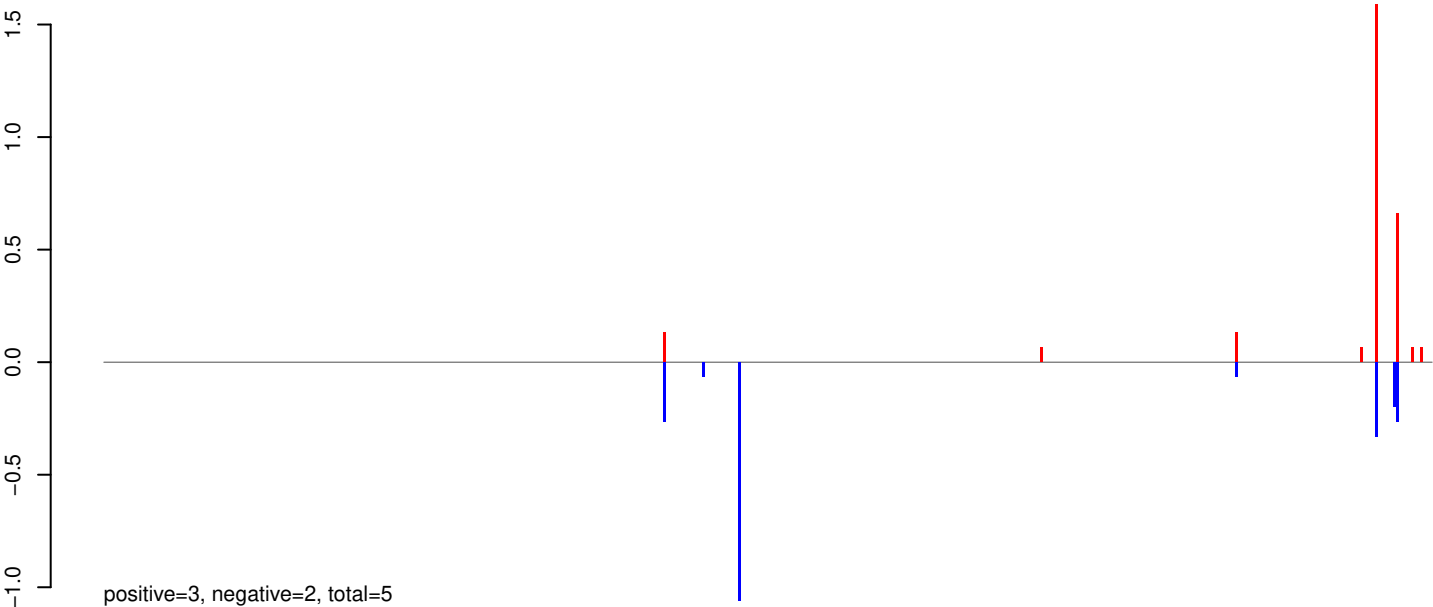
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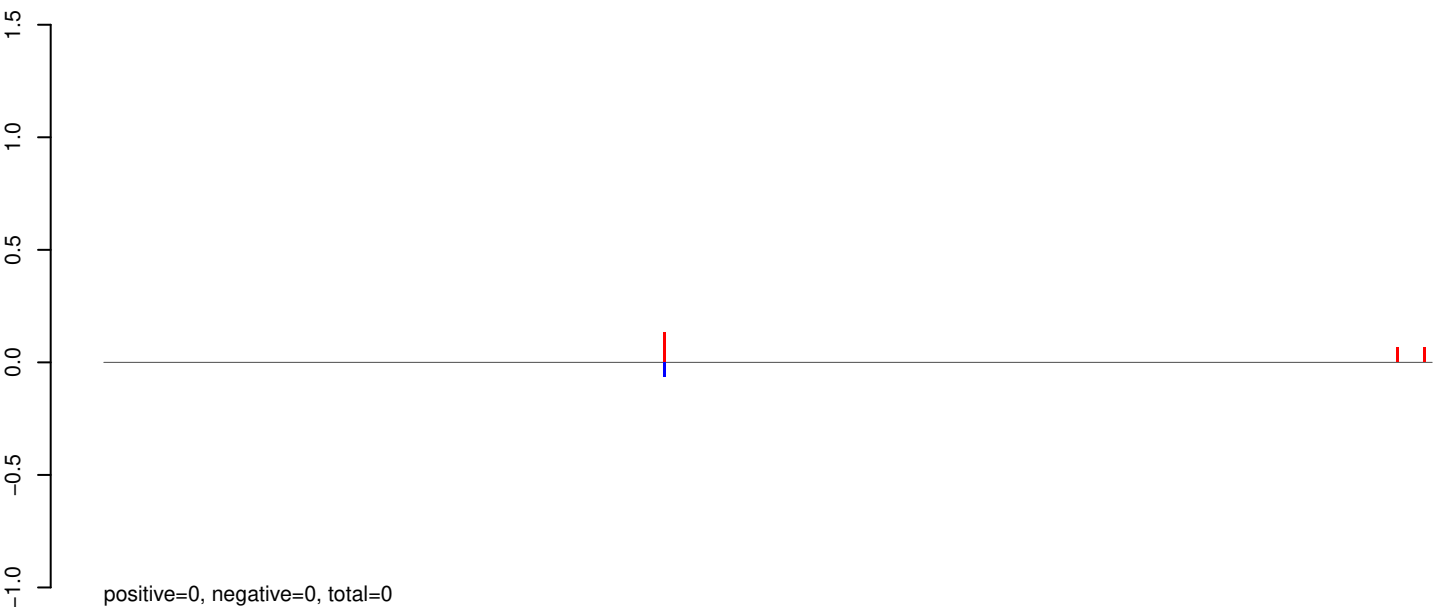
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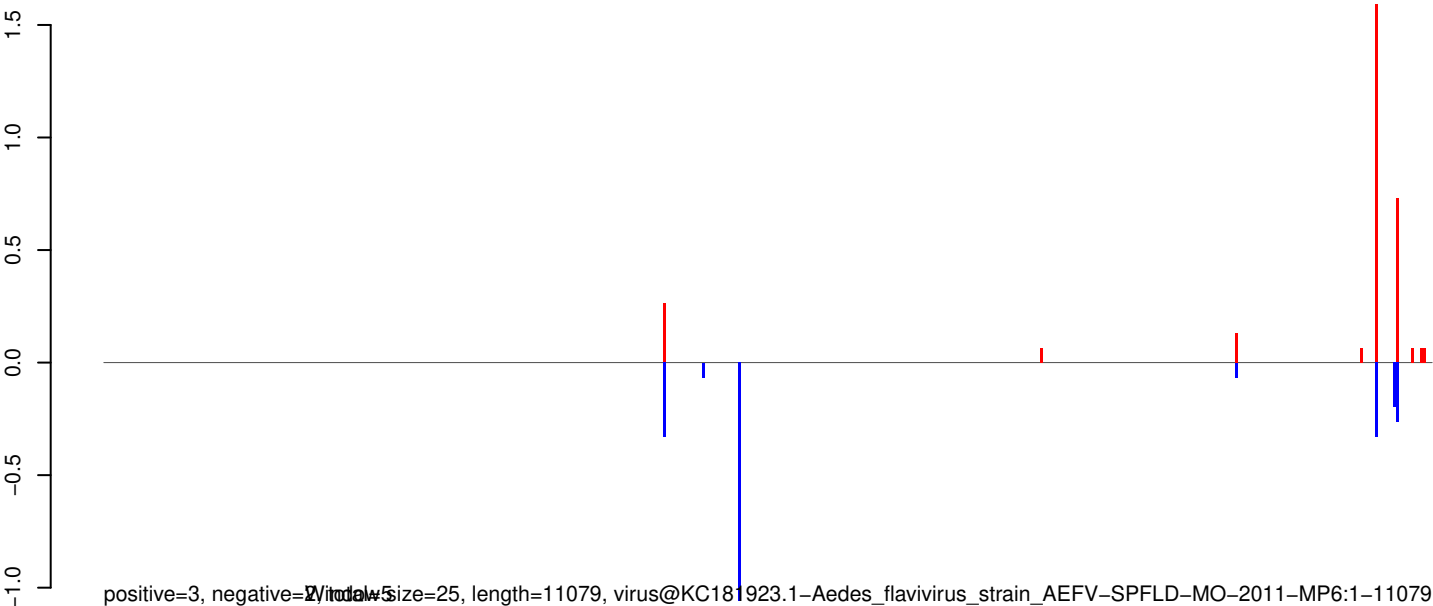
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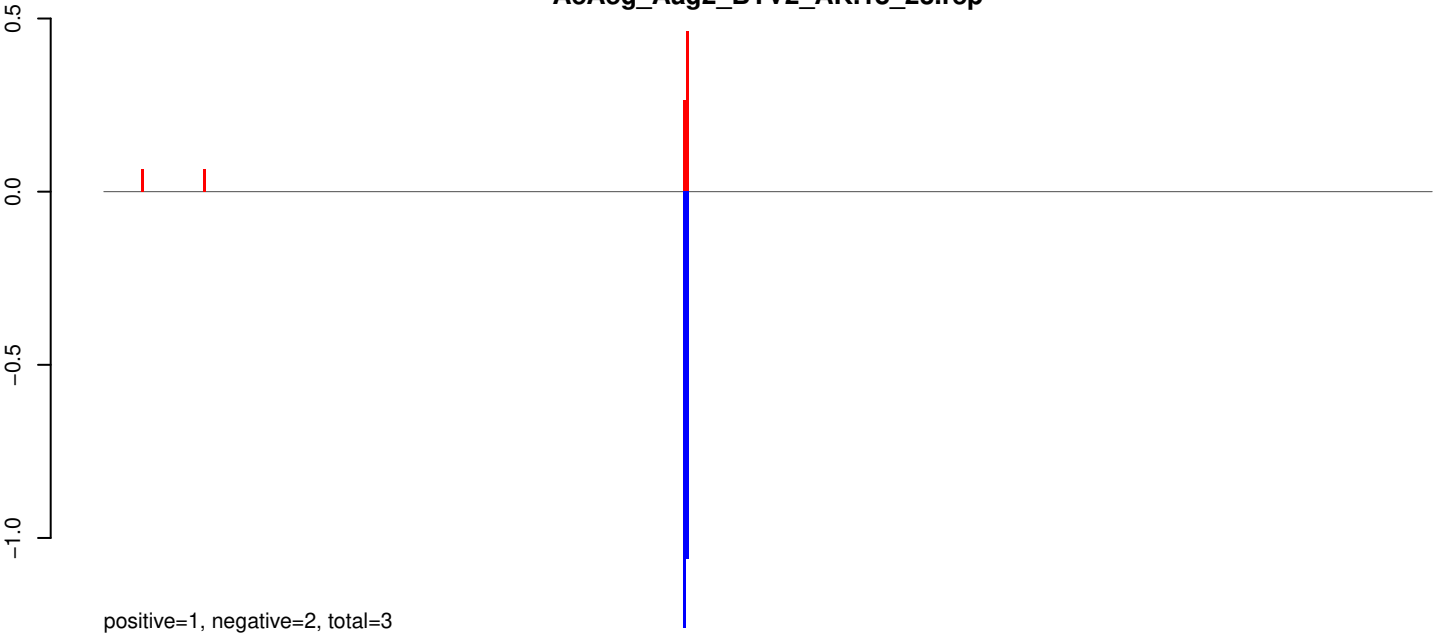
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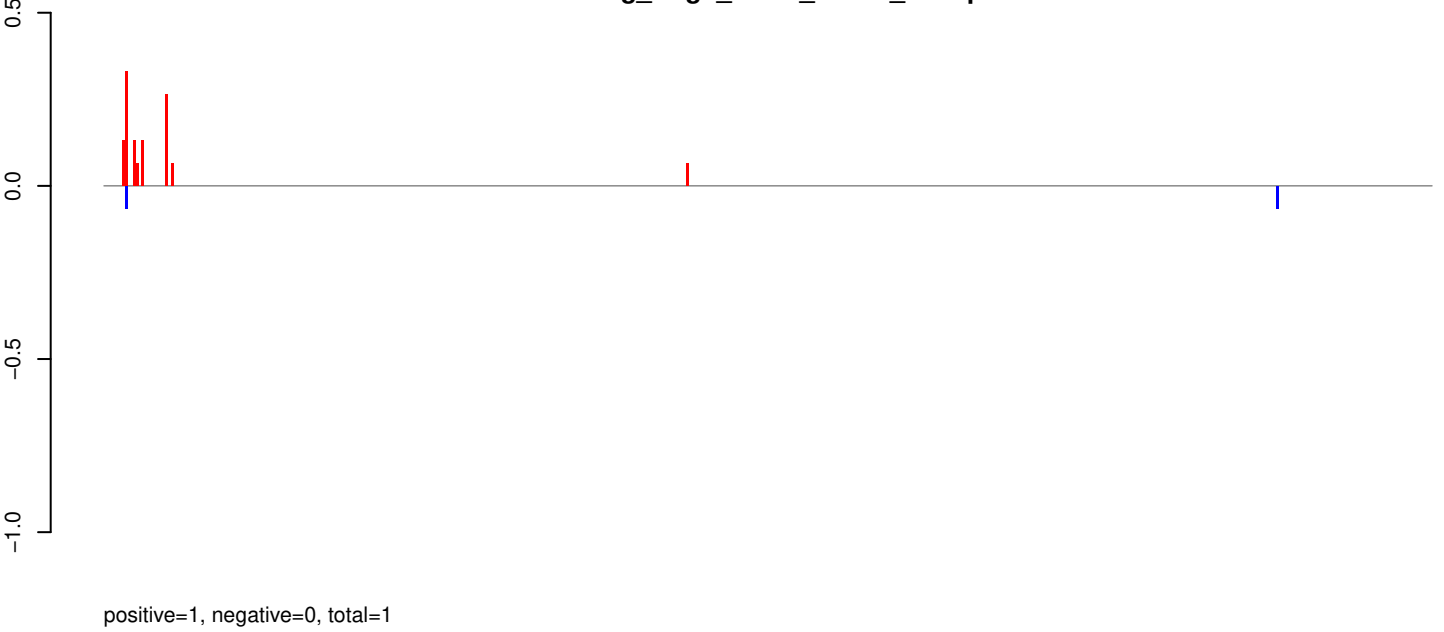
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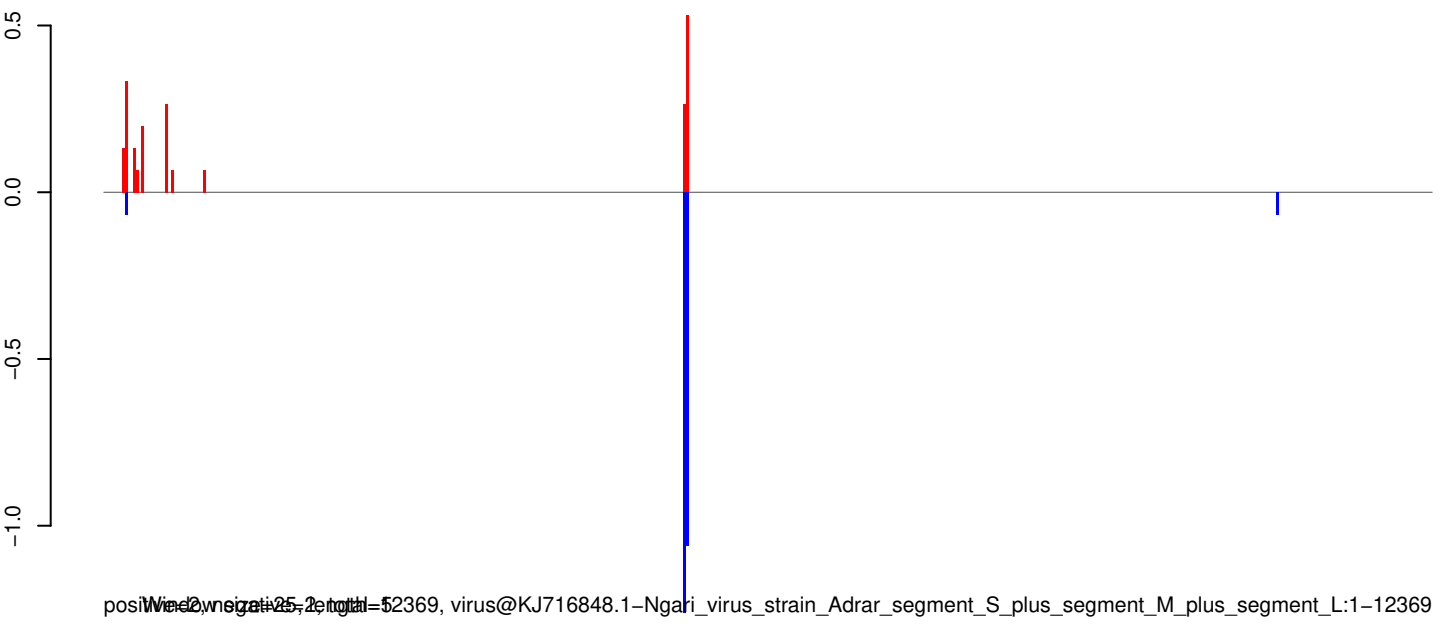
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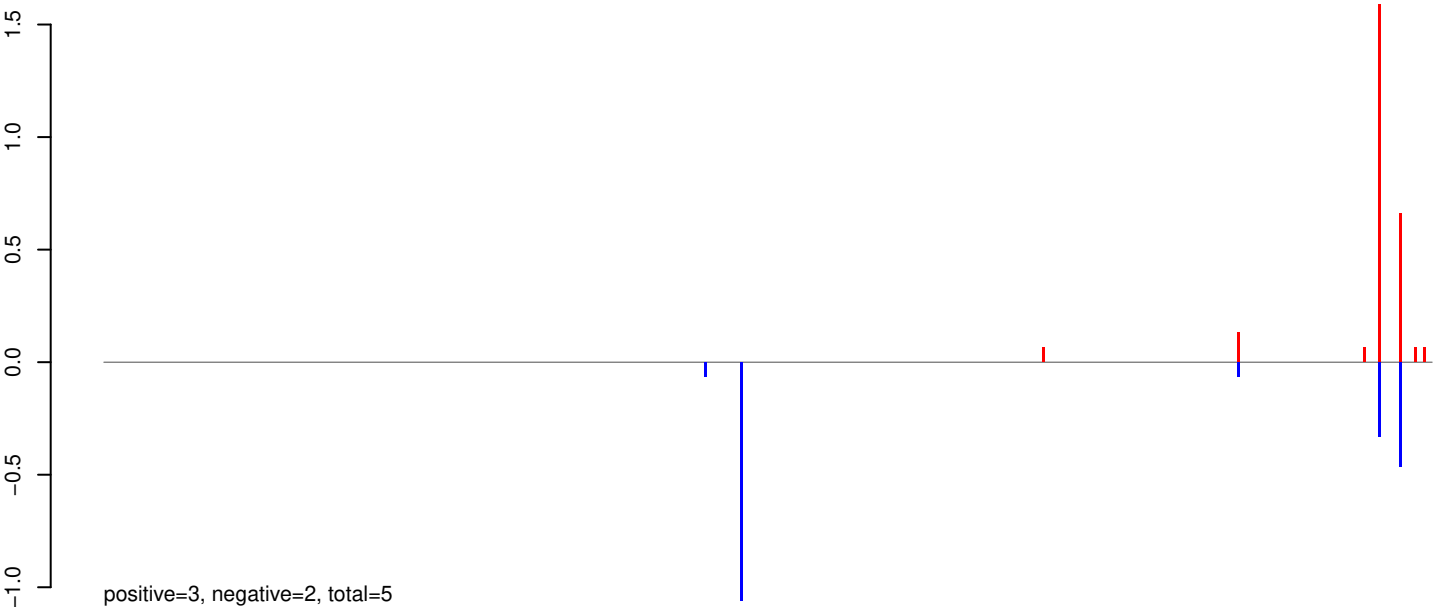
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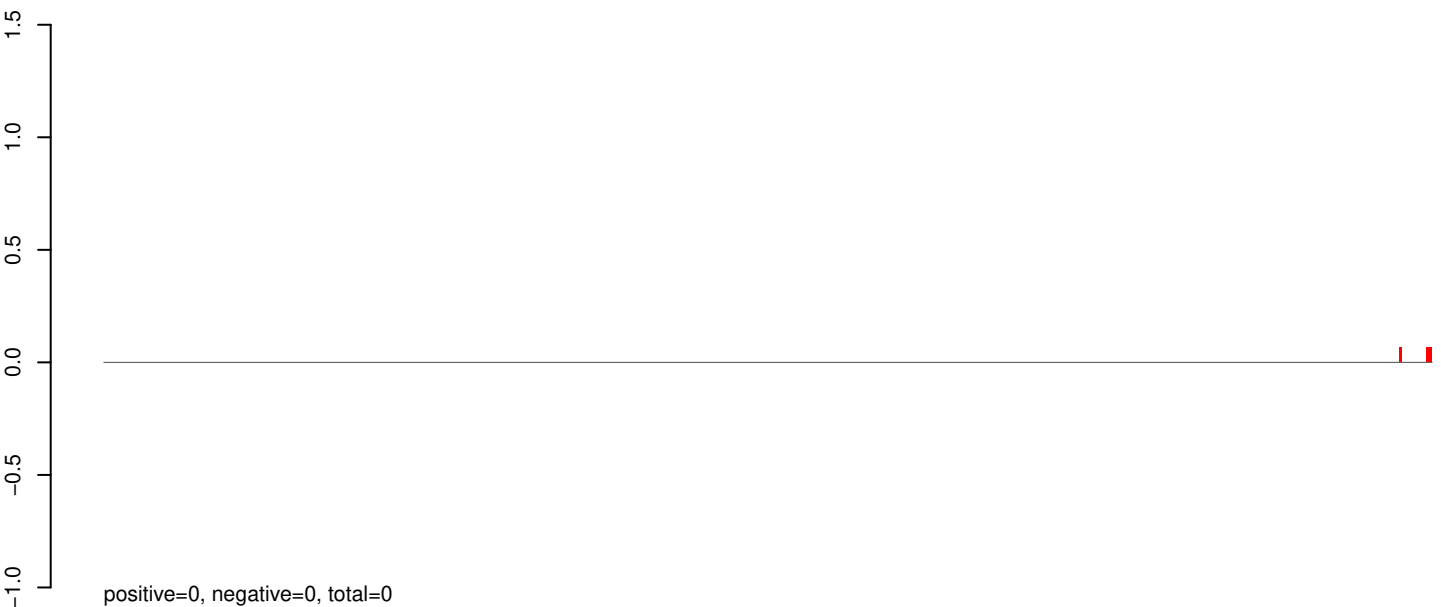
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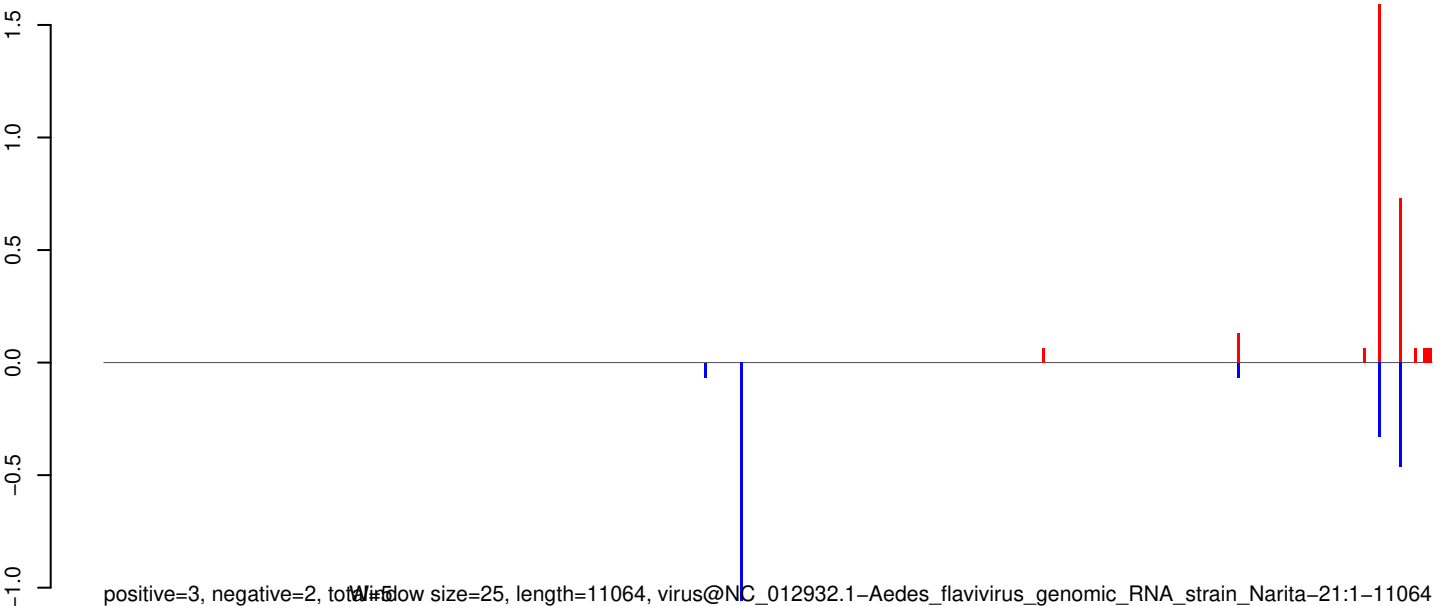
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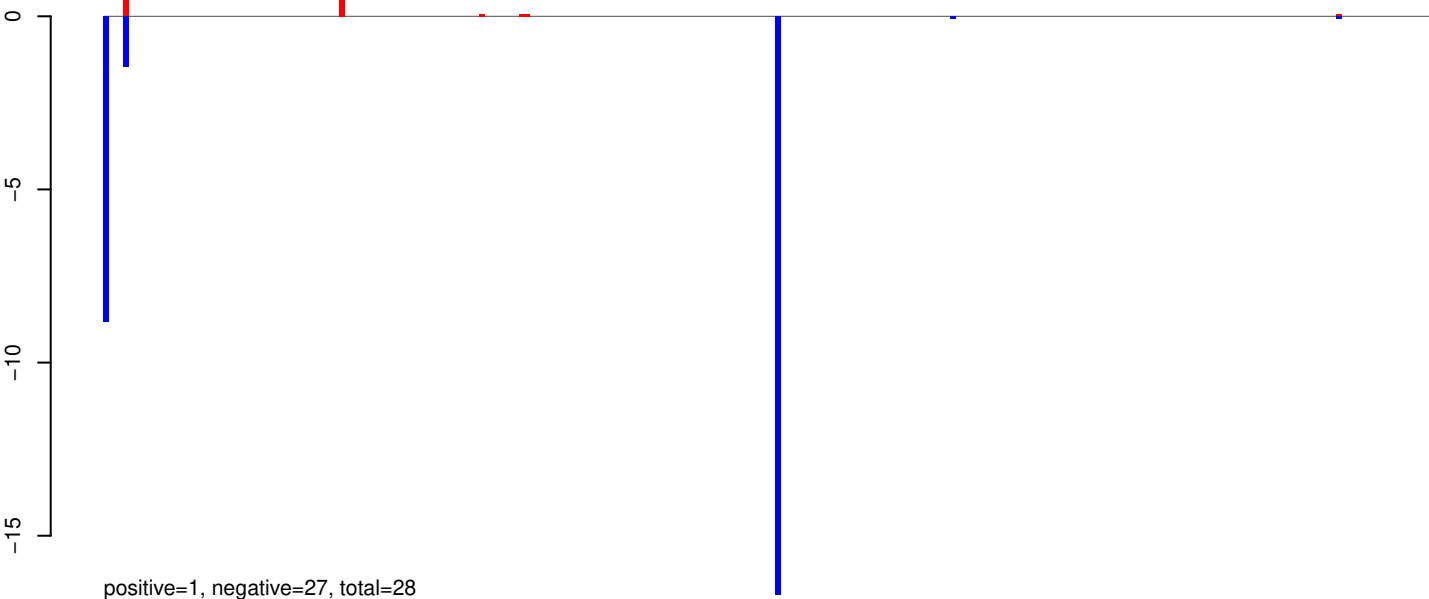
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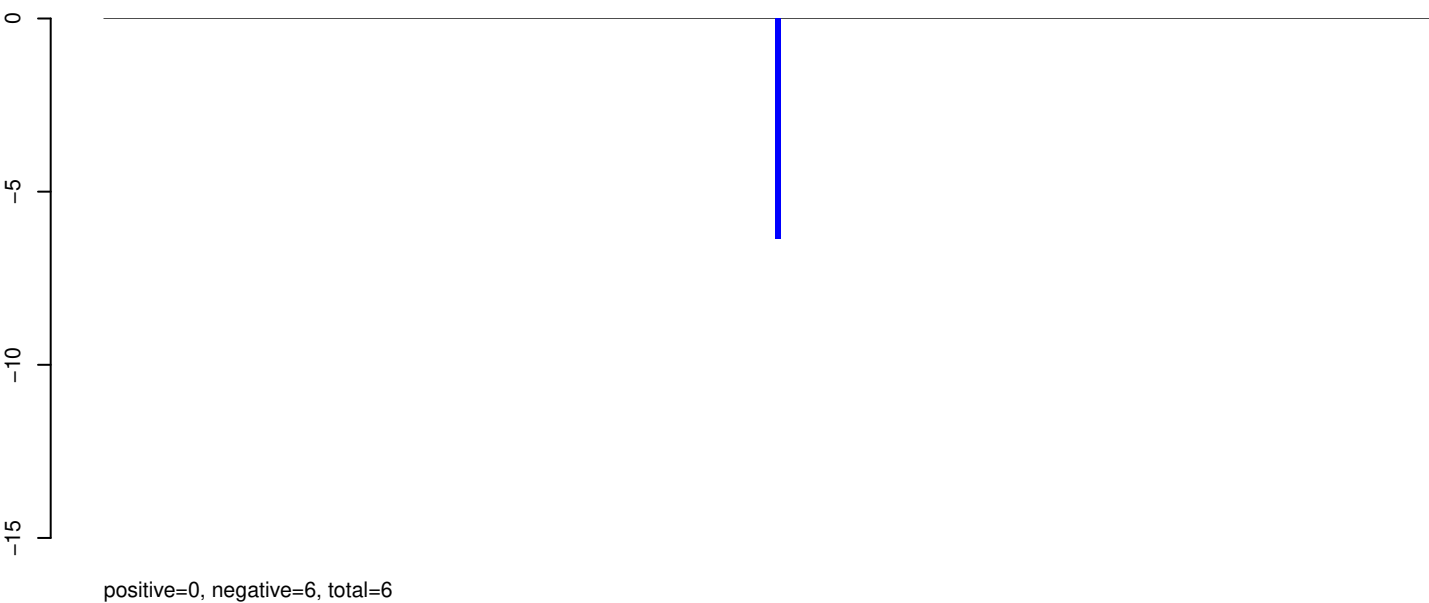
AeAeg_Aag2_BTV2_AK.rep



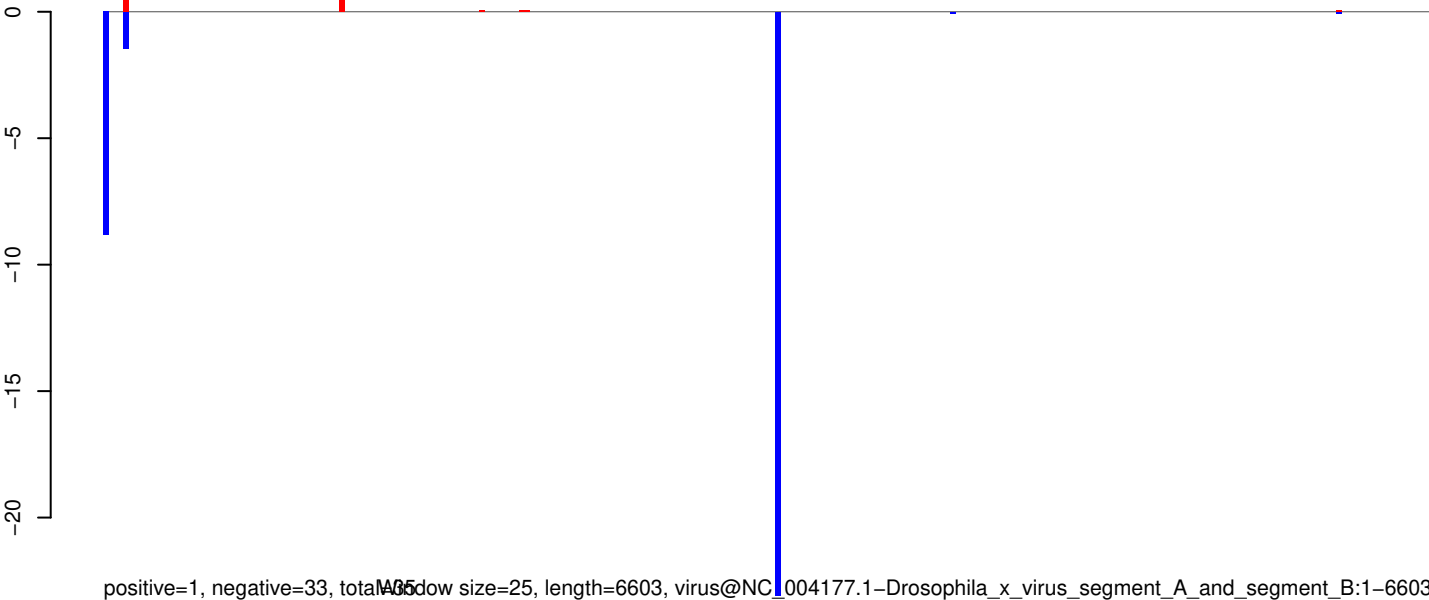
AeAeg_Aag2_BTV2_AK.18_23.rep



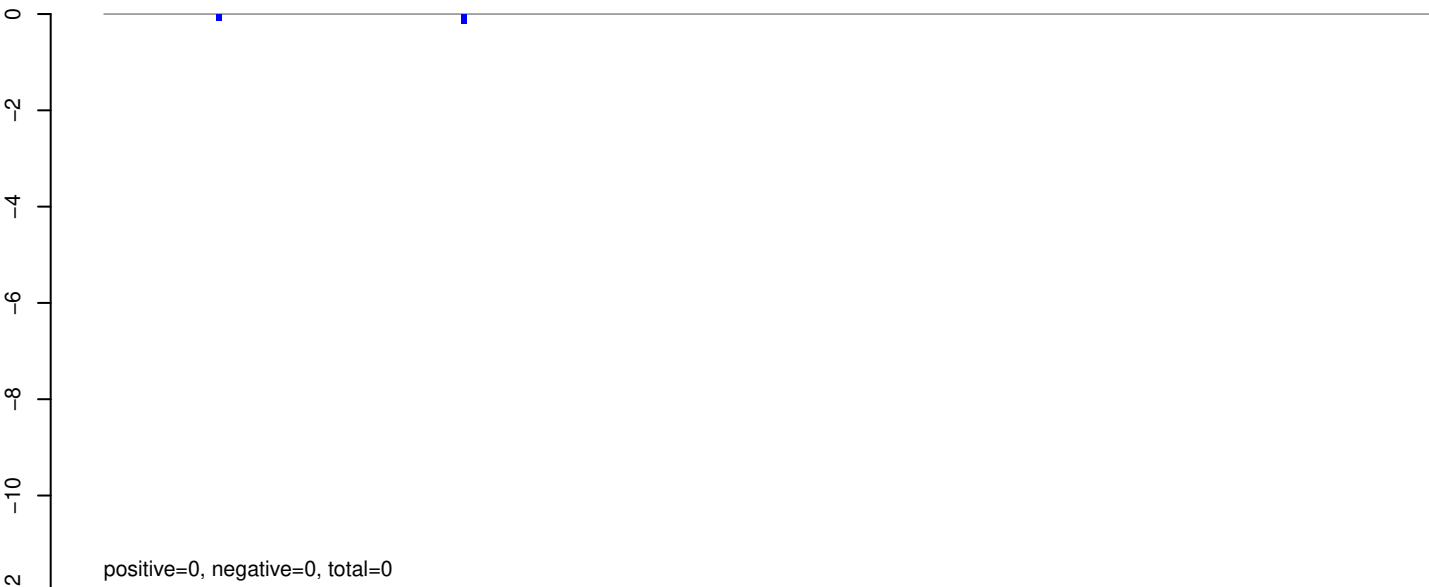
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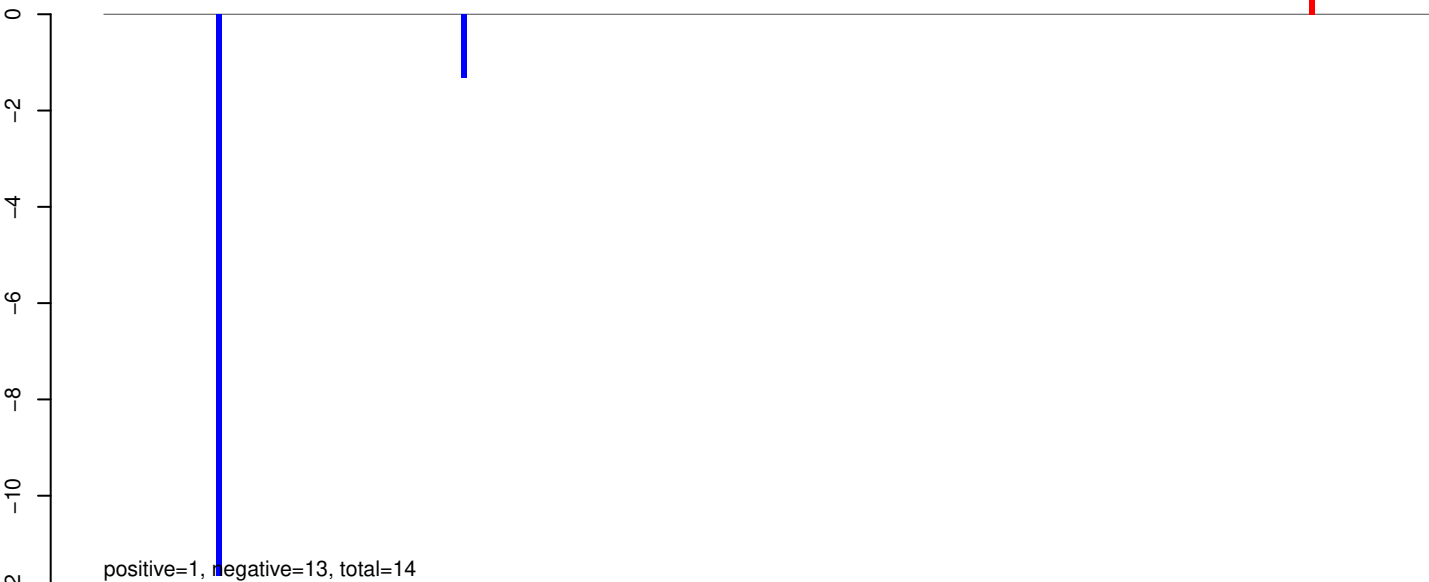
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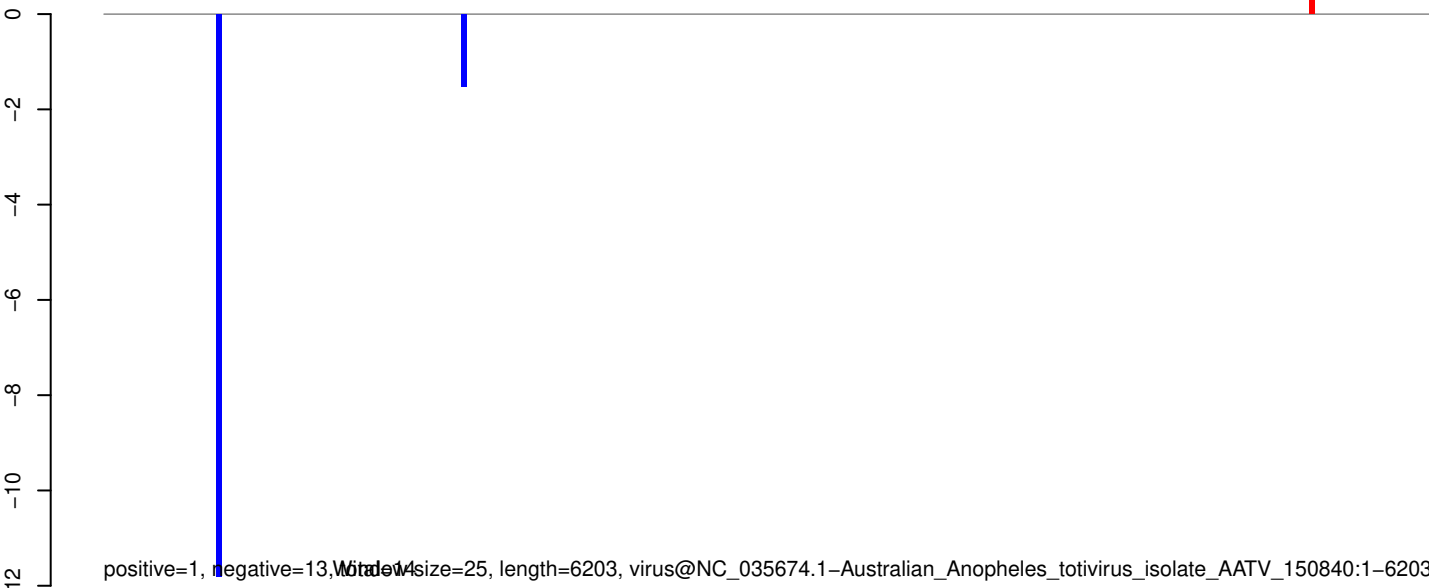
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AeAeg_Aag2_BTV2_AK.24_35.rep

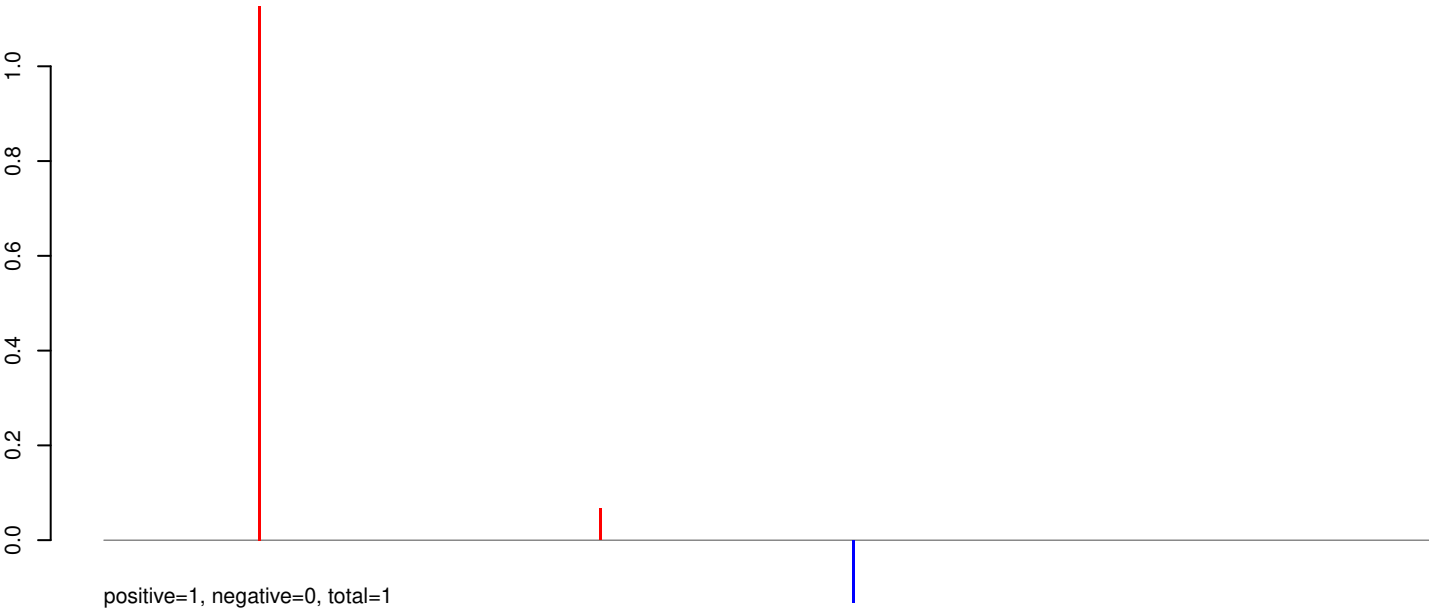


AeAeg_Aag2_BTV2_AK.rep

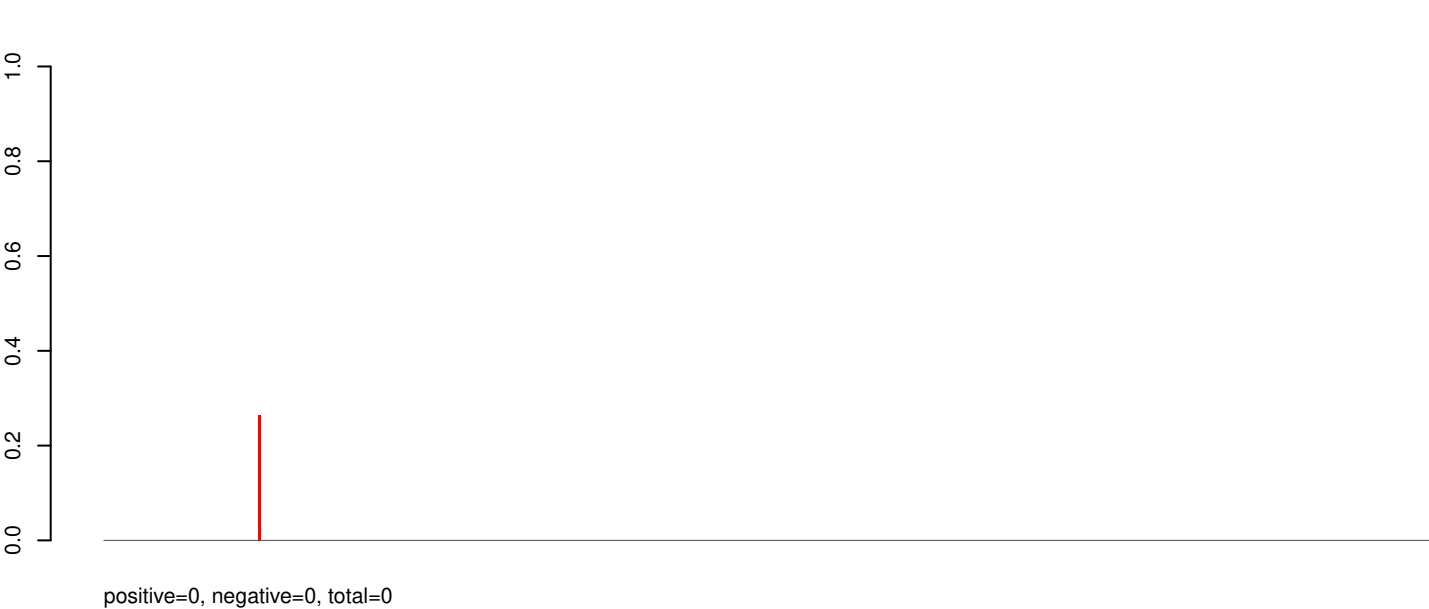


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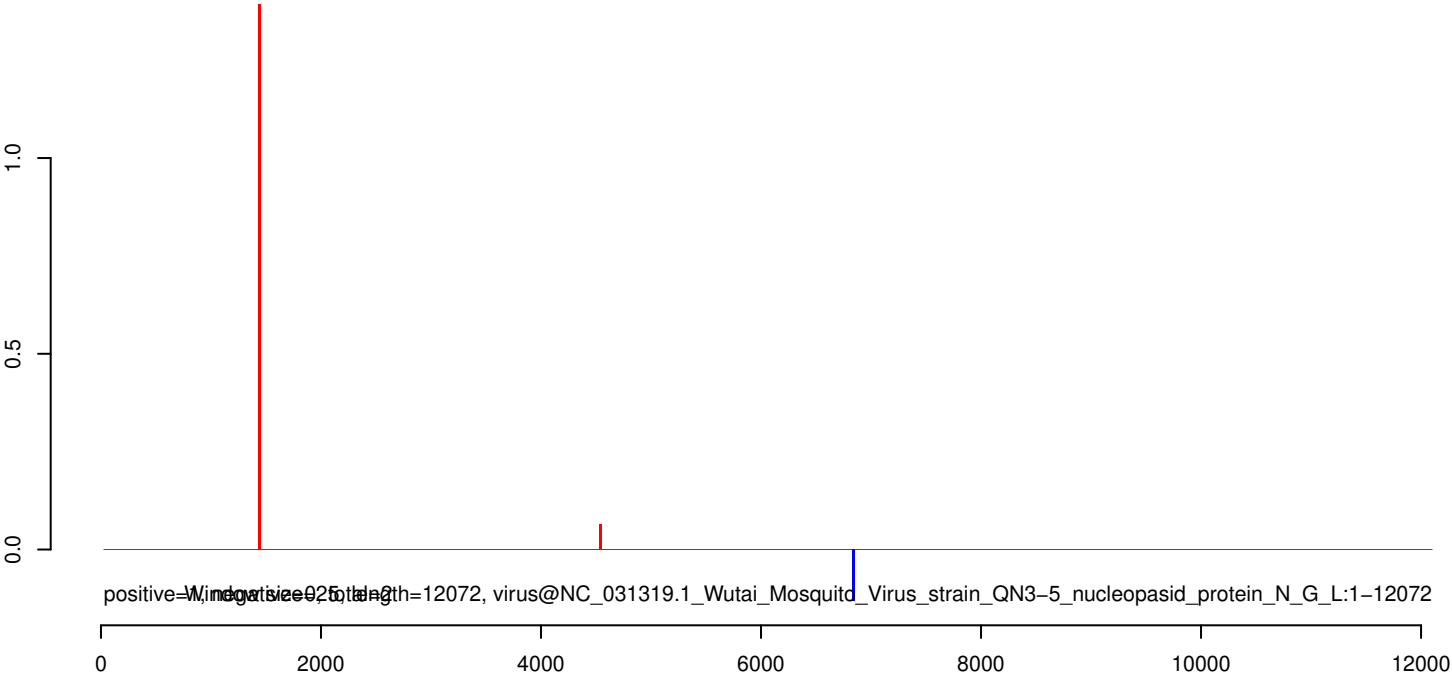
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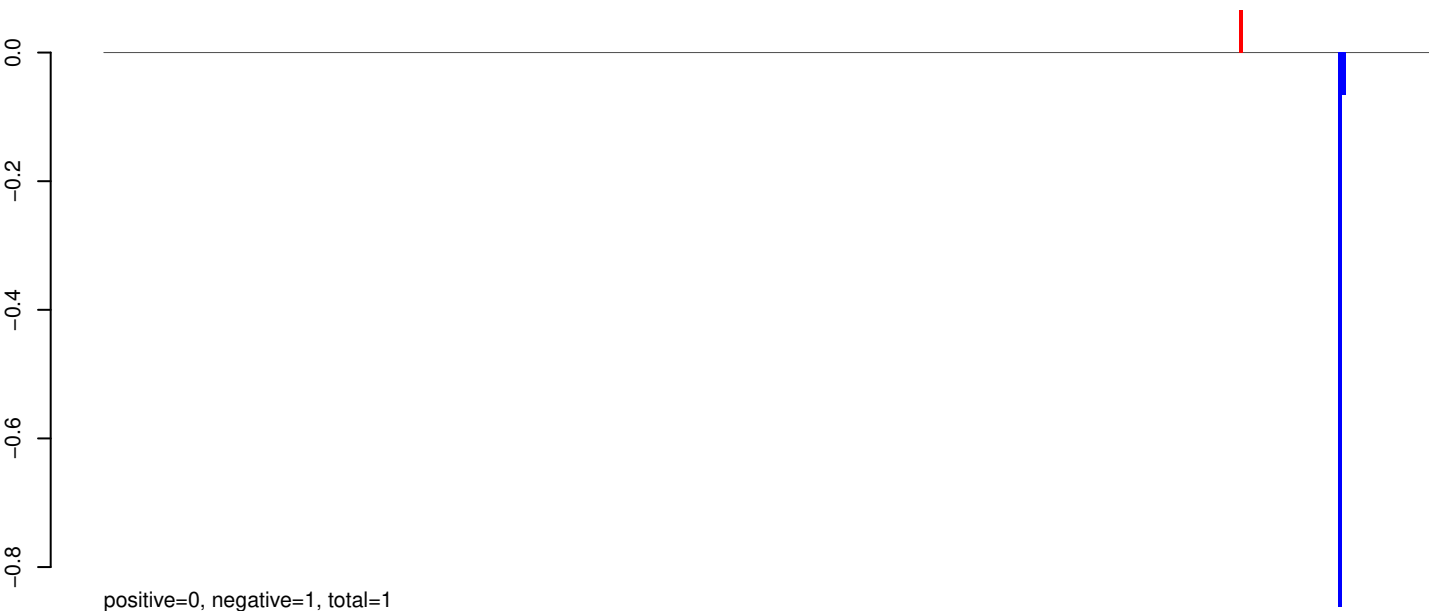
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AeAeg_Aag2_BTV2_AK.rep

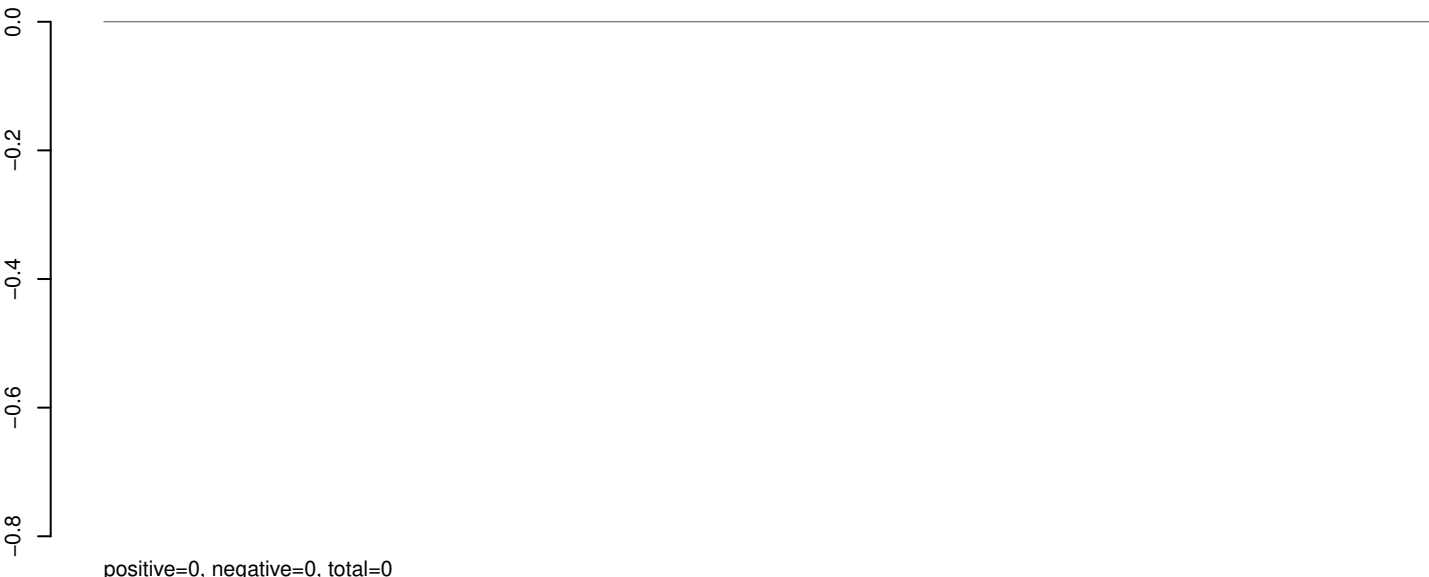


AeAeg_Aag2_BTV2_AK.18_23.rep



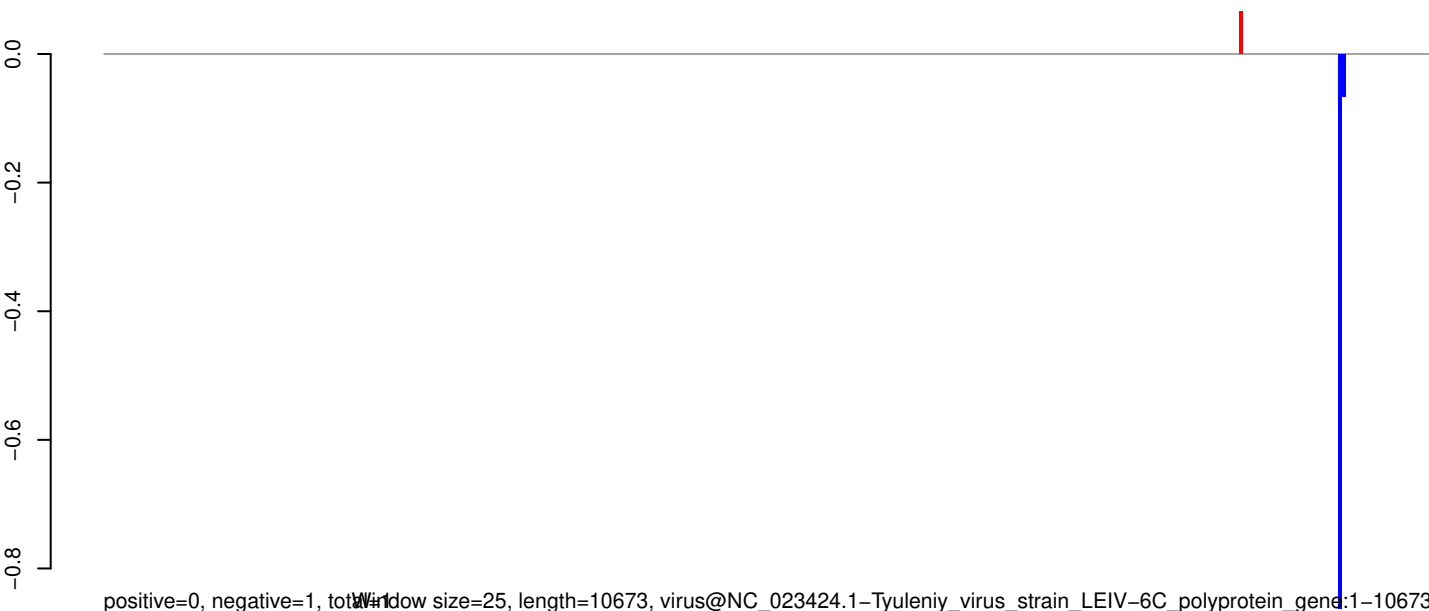
positive=0, negative=1, total=1

AeAeg_Aag2_BTV2_AK.24_35.rep



positive=0, negative=0, total=0

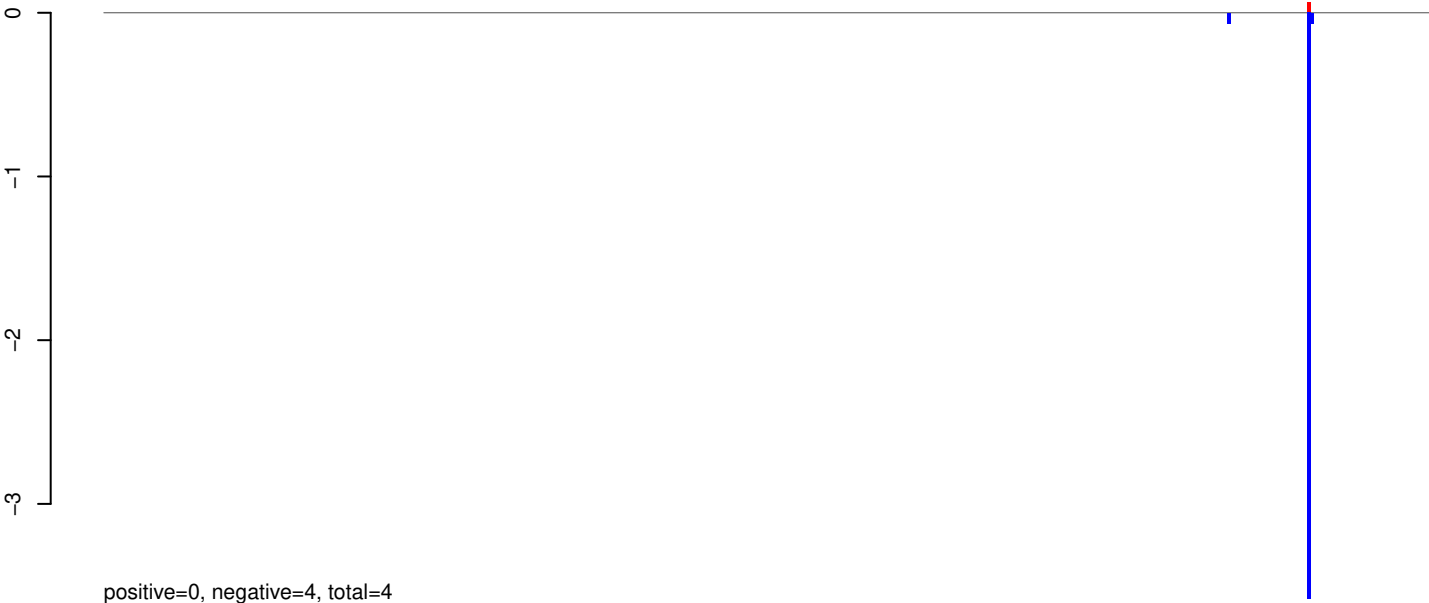
AeAeg_Aag2_BTV2_AK.rep



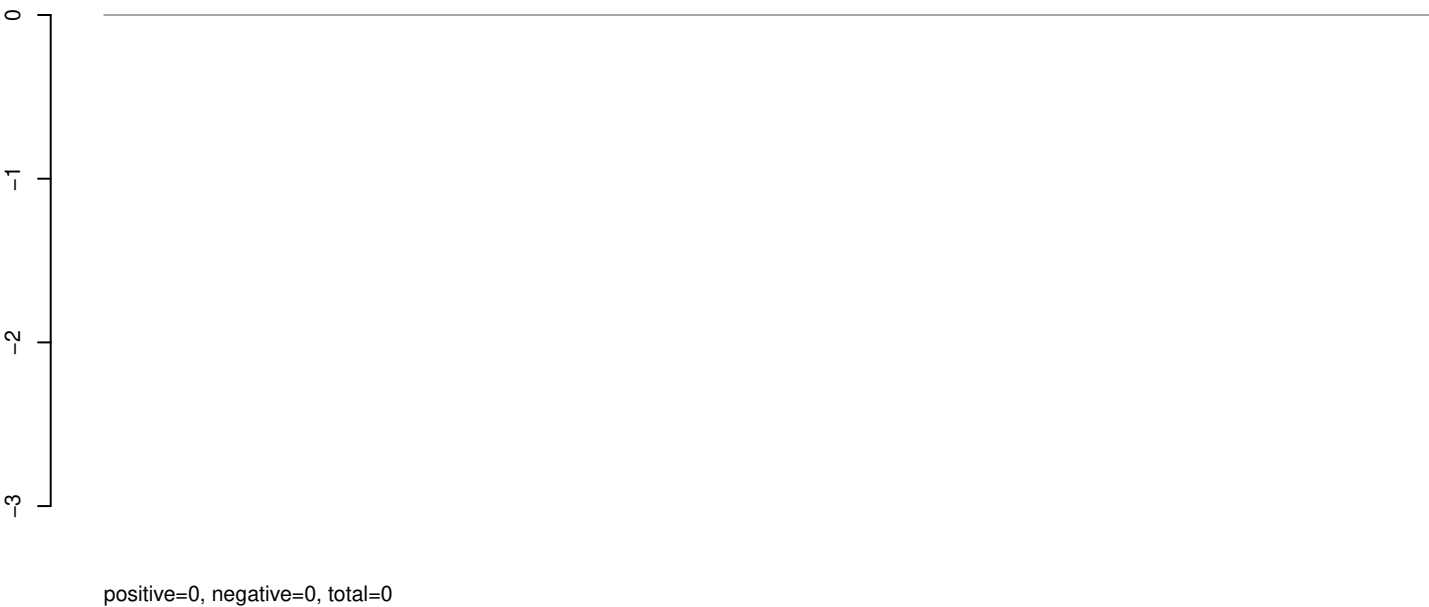
positive=0, negative=1, total=1 window size=25, length=10673, virus@NC_023424.1-Tyuleniy_virus_strain_LEIV-6C_polyprotein_gene:1-10673

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



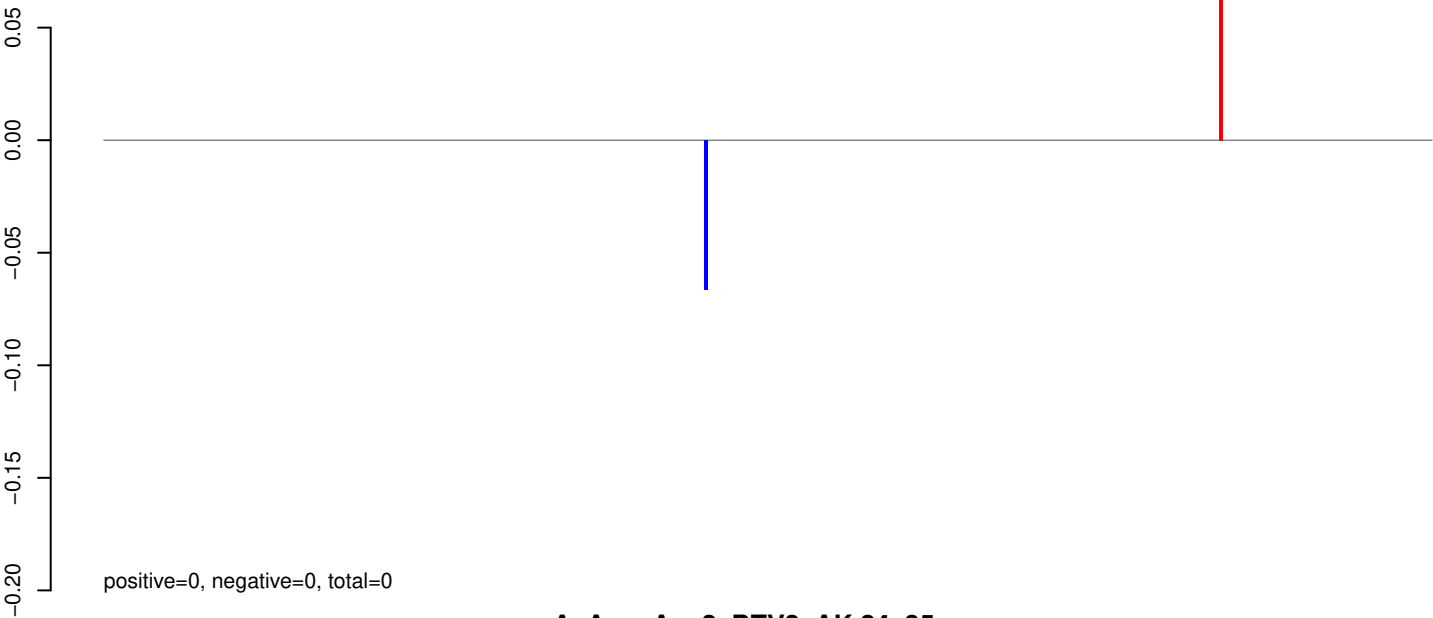
AeAeg_Aag2_BTV2_AK.rep



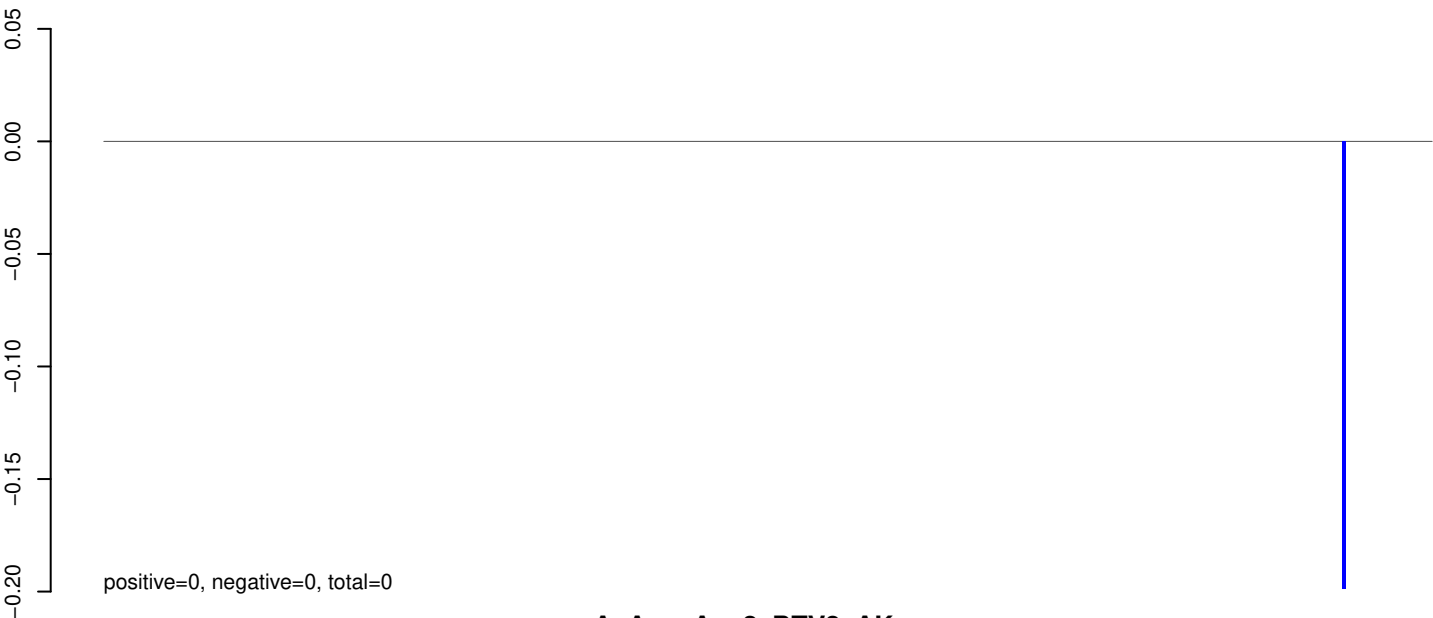
Window size=25, length=10874, virus@NC_009029.2-Kokobera_virus:1-10874

0 2000 4000 6000 8000 10000

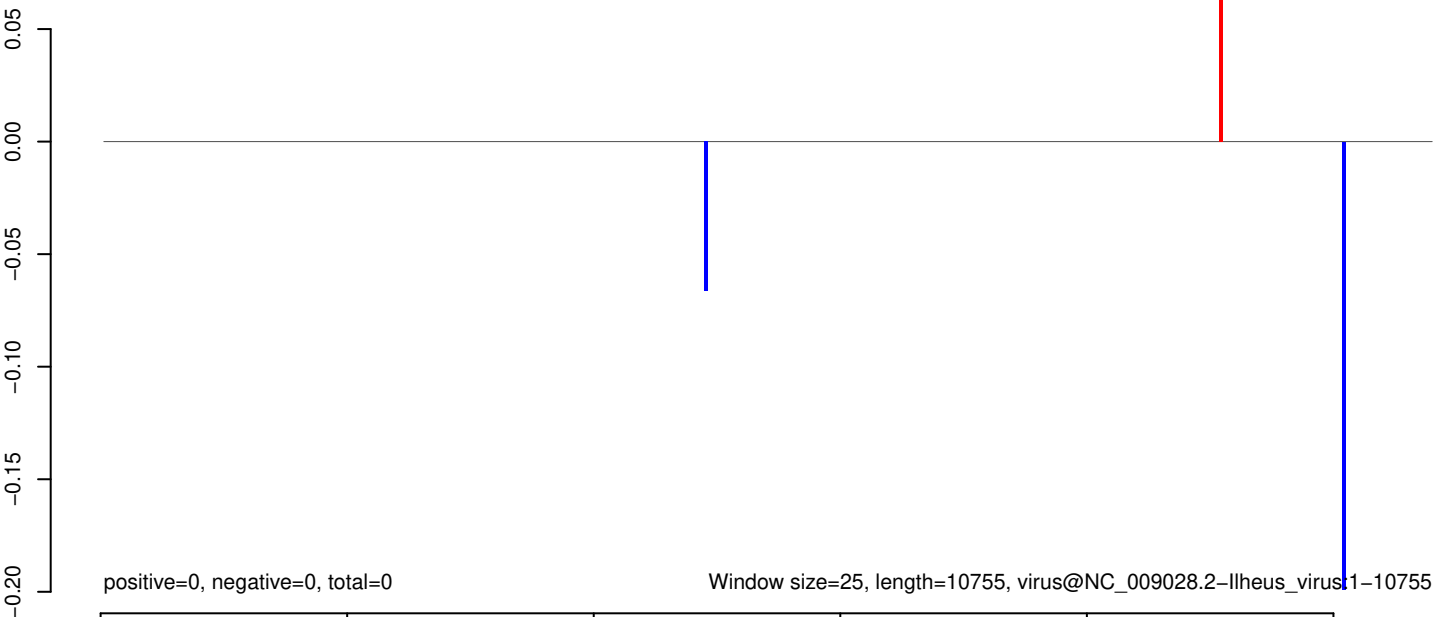
AeAeg_Aag2_BTV2_AK.18_23.rep



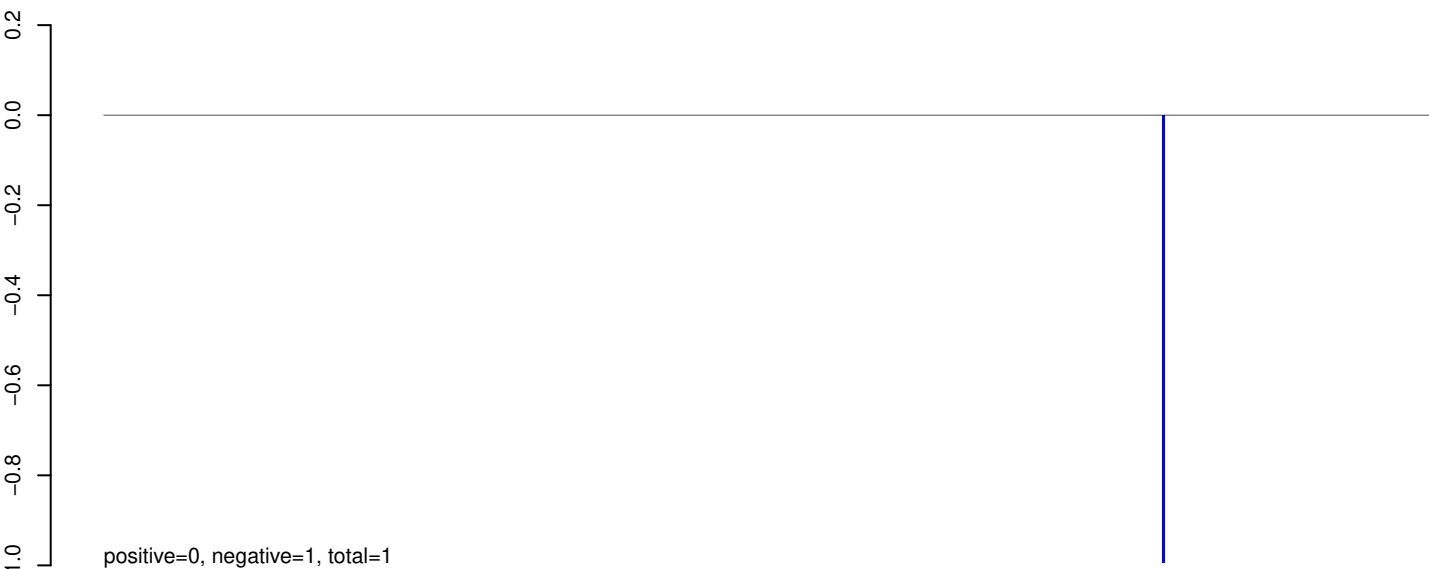
AeAeg_Aag2_BTV2_AK.24_35.rep



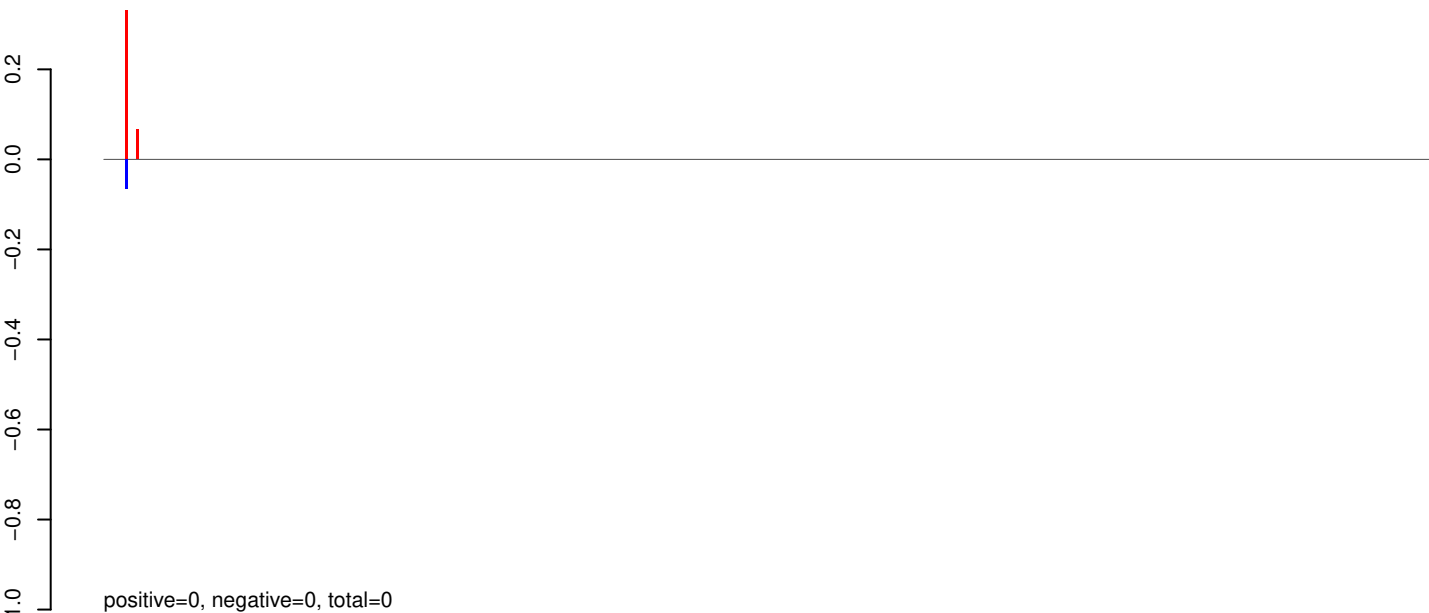
AeAeg_Aag2_BTV2_AK.rep



AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



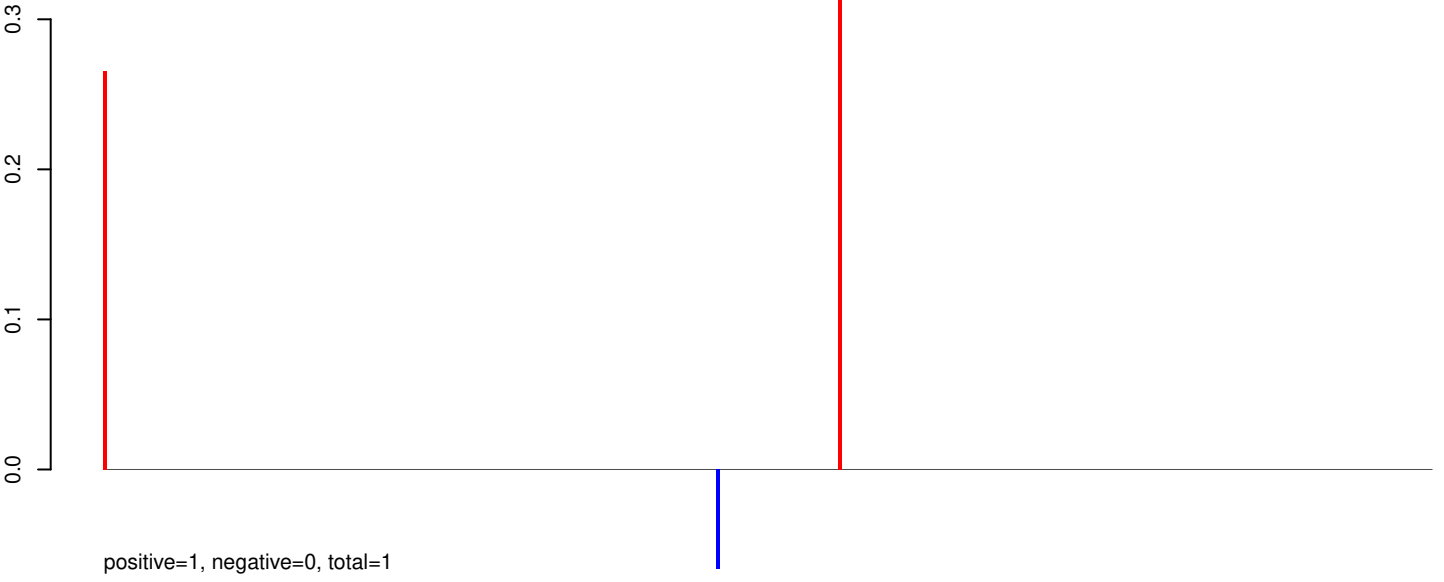
AeAeg_Aag2_BTV2_AK.rep



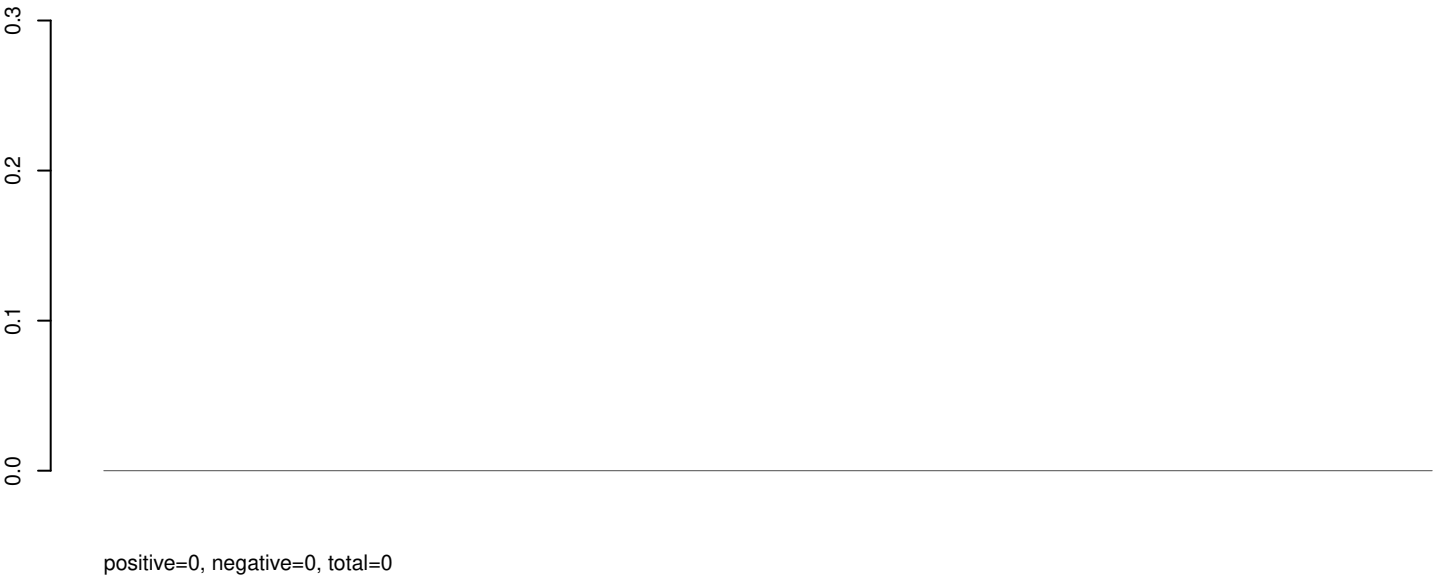
W:\pos\size=0.25, length=1,2897-Virus@KC608151.1-Ilesha_virus_strain_ILESHA_8e_segment_S_plus_segment_M_plus_segment_L:1-12297

0 2000 4000 6000 8000 10000 12000

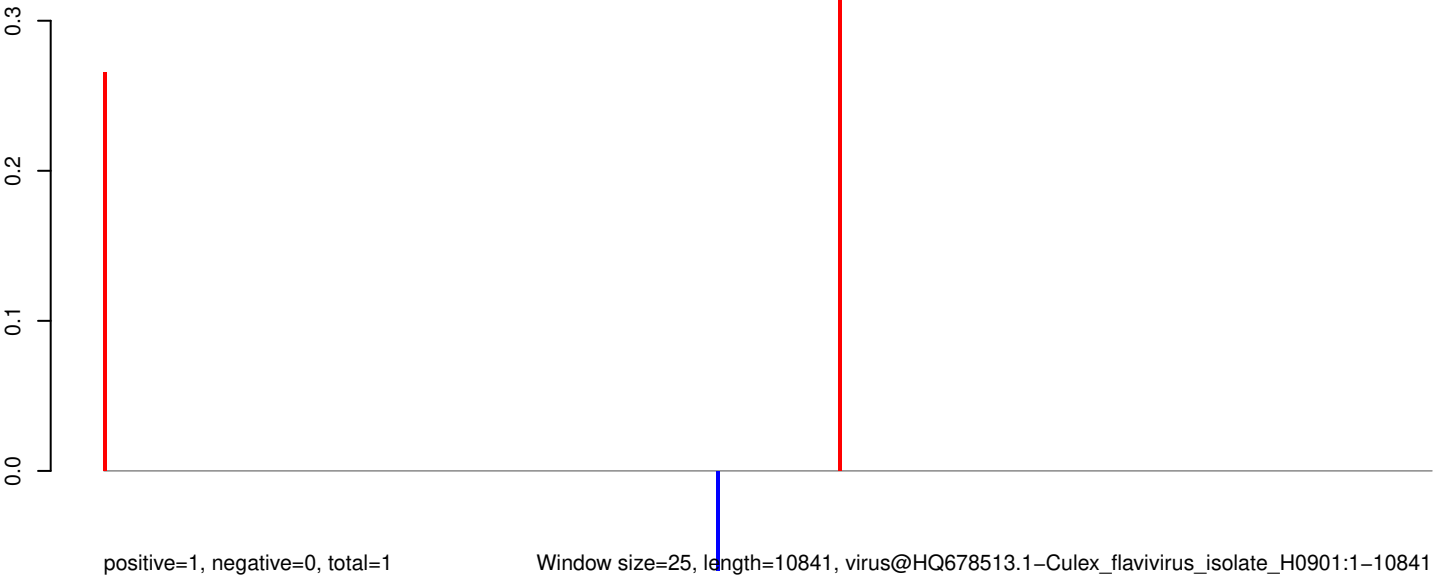
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

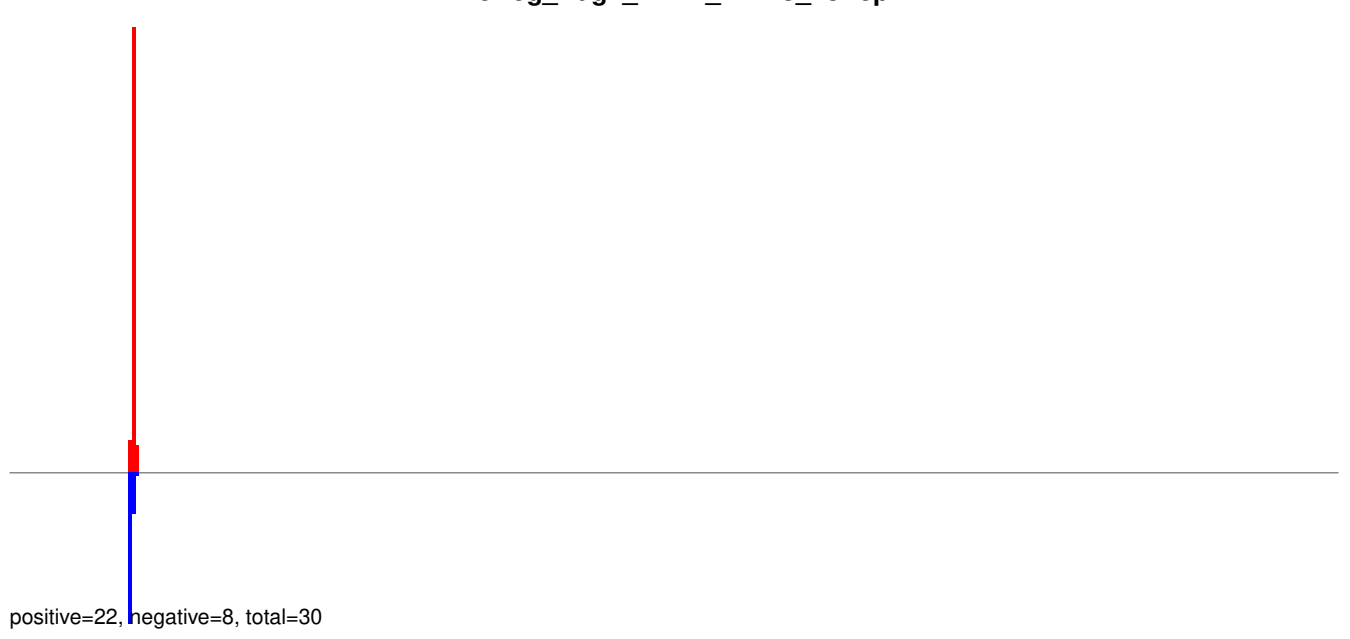


AeAeg_Aag2_BTV2_AK.rep

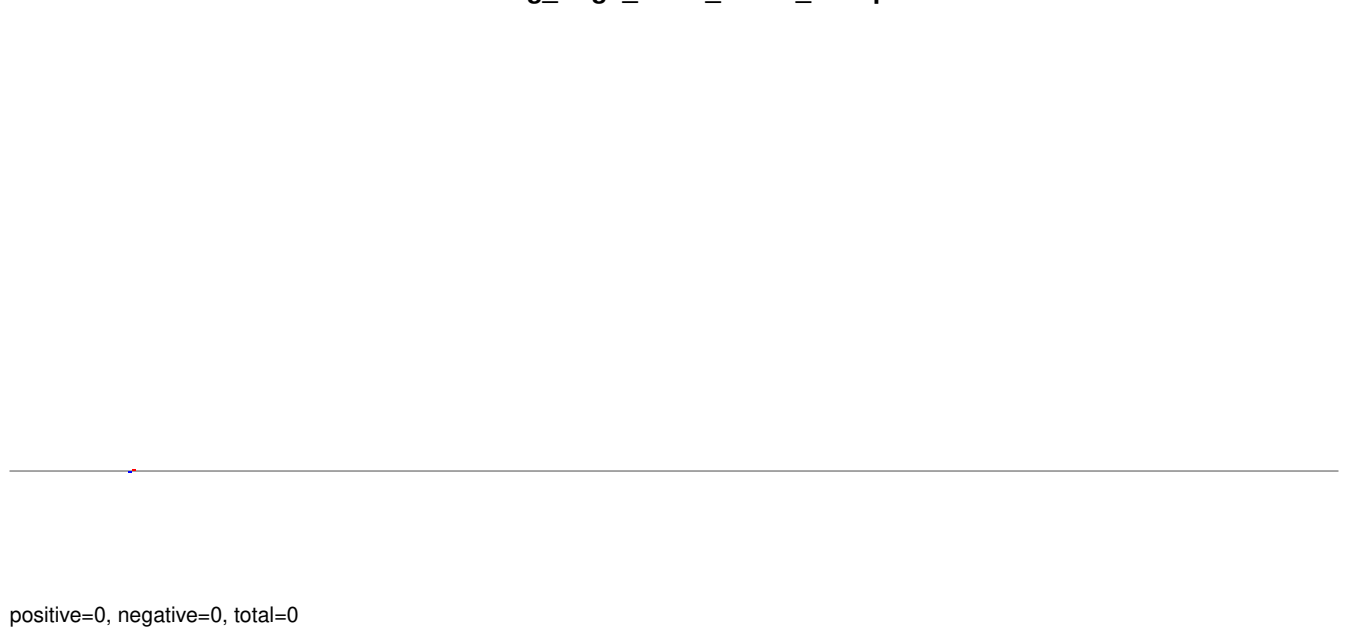


0 2000 4000 6000 8000 10000

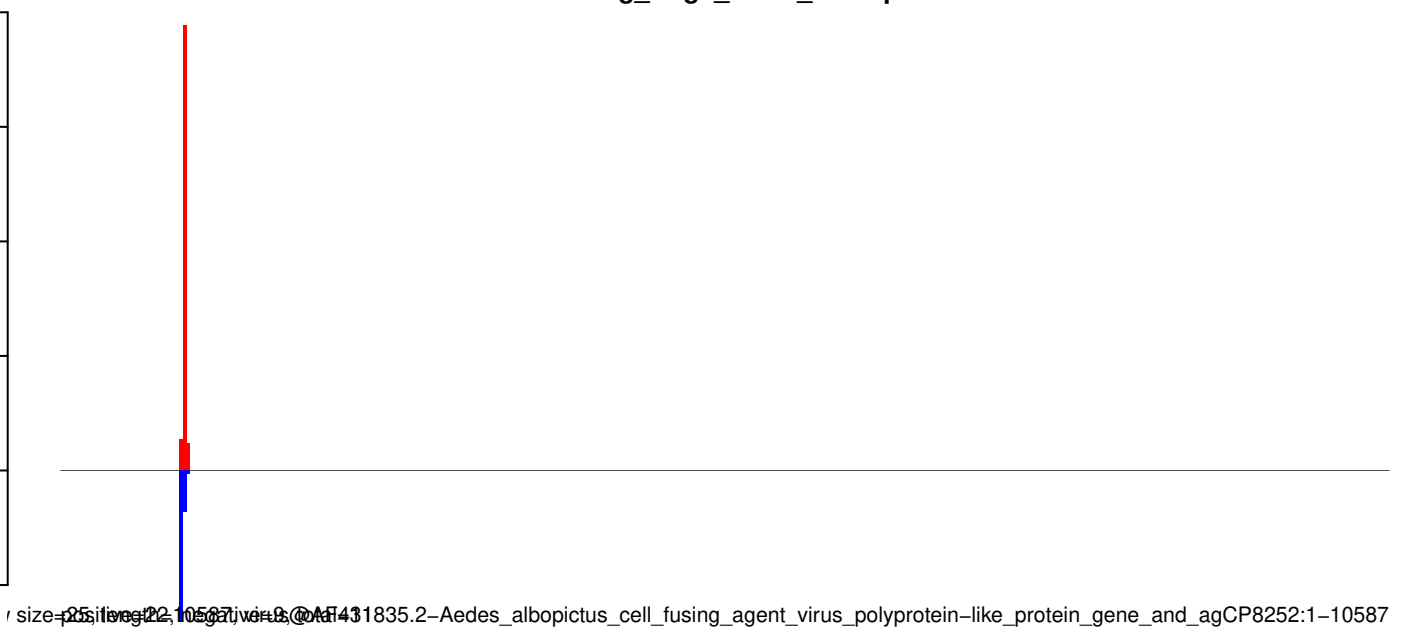
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



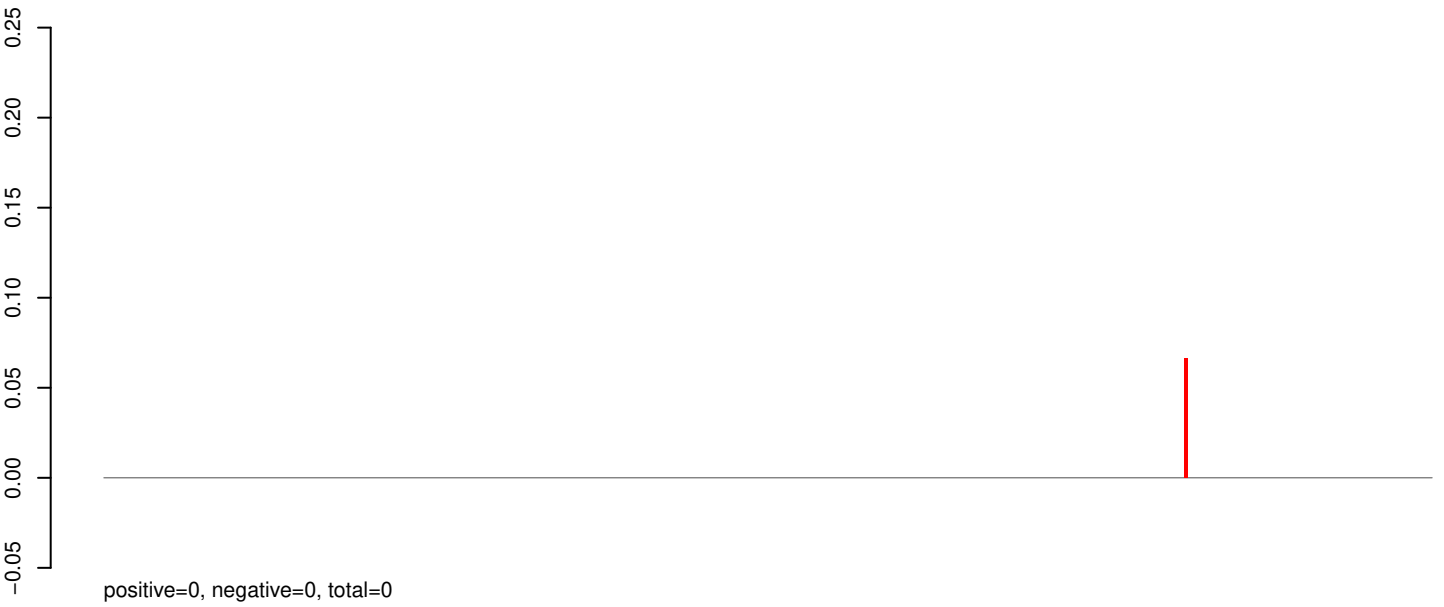
AeAeg_Aag2_BTV2_AK.rep



AeAeg_Aag2_BTV2_AK.18_23.rep



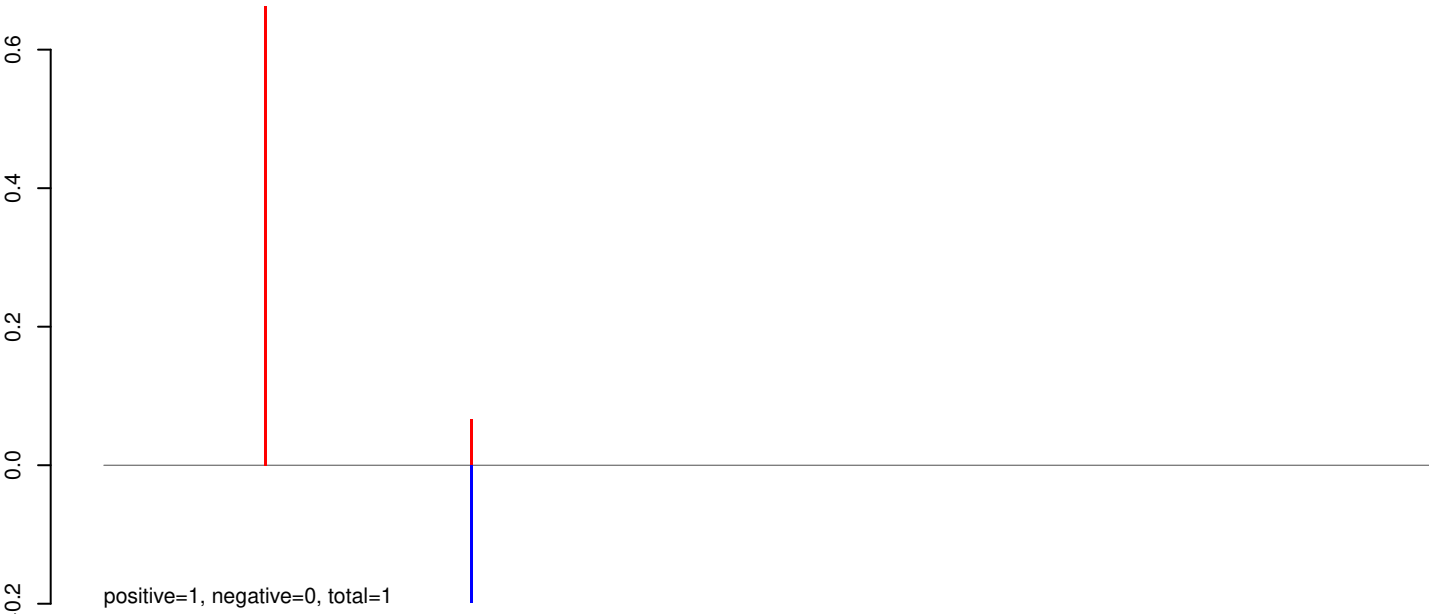
AeAeg_Aag2_BTV2_AK.24_35.rep



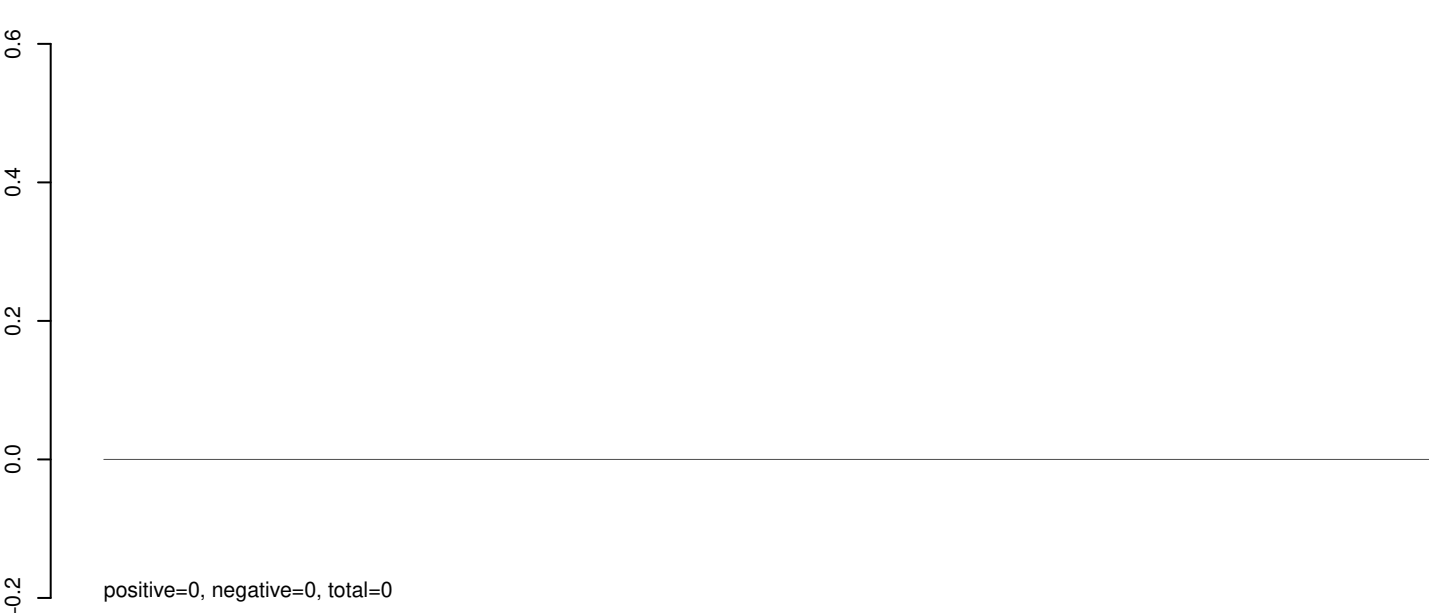
AeAeg_Aag2_BTV2_AK.rep



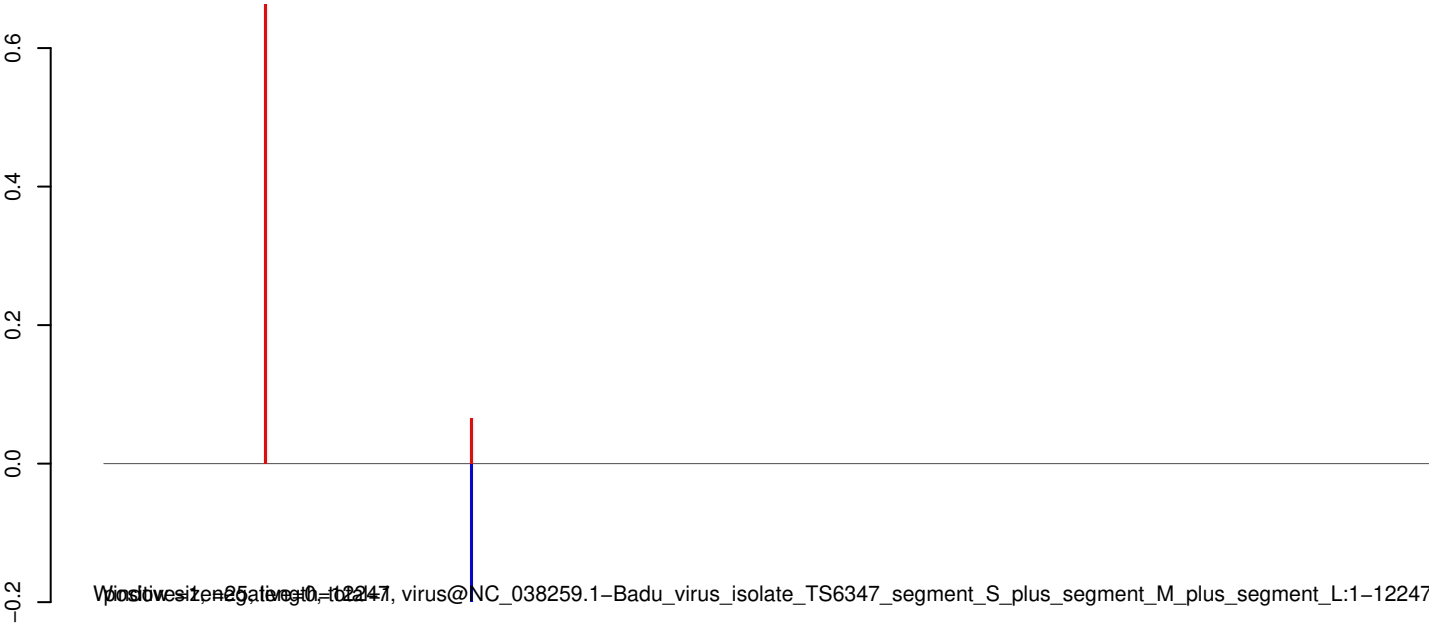
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

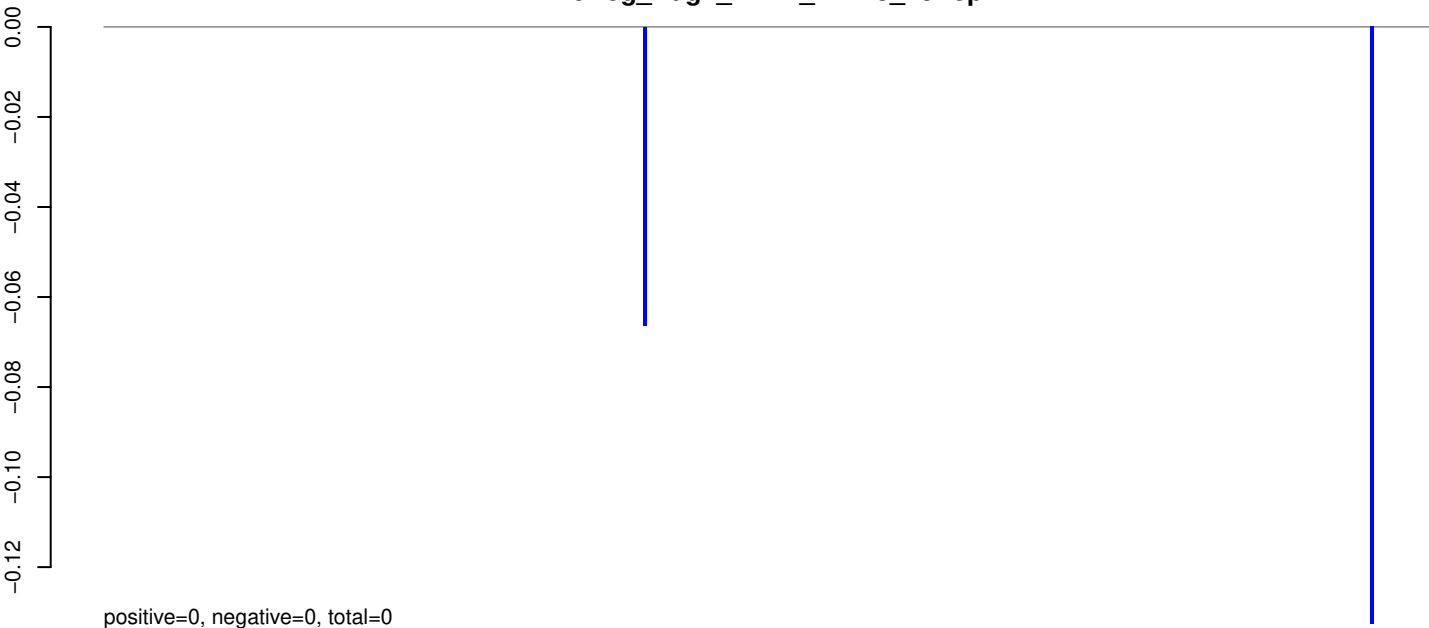


AeAeg_Aag2_BTV2_AK.rep

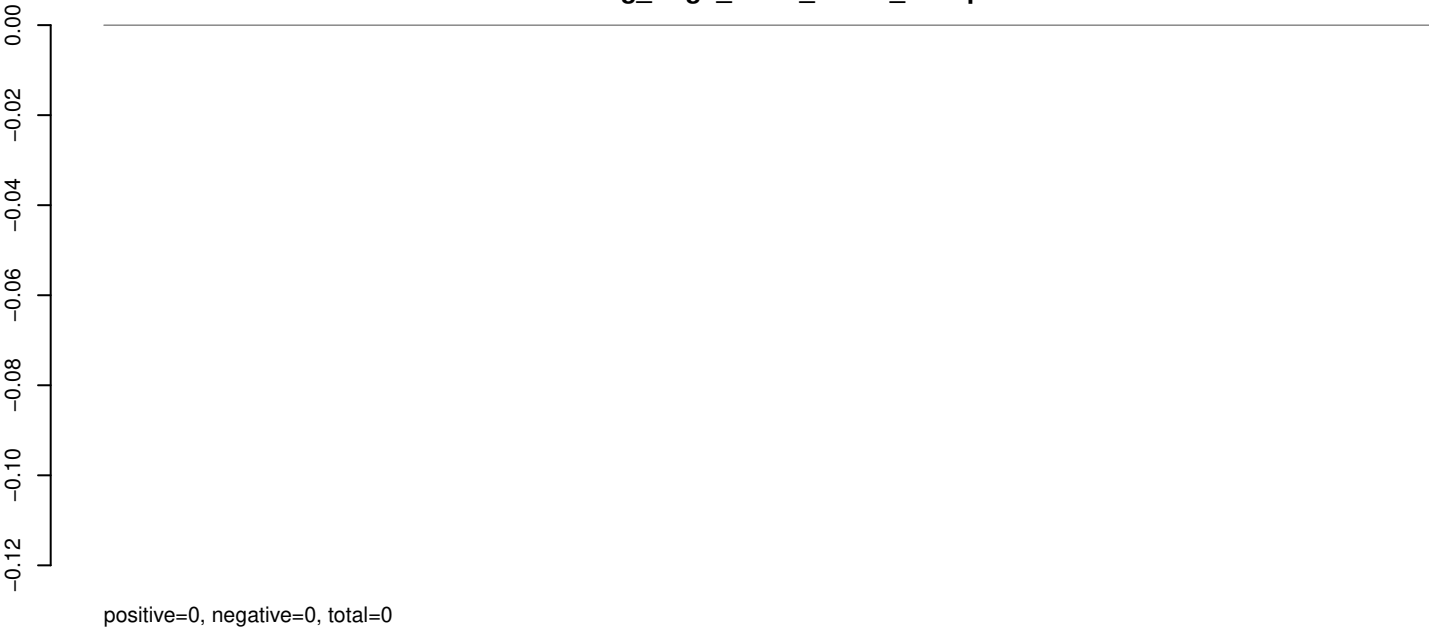


positives=1, negatives=1, total=2, virus@NC_038259.1-Badu_virus_isolate_TS6347_segment_S_plus_segment_M_plus_segment_L:1-12247

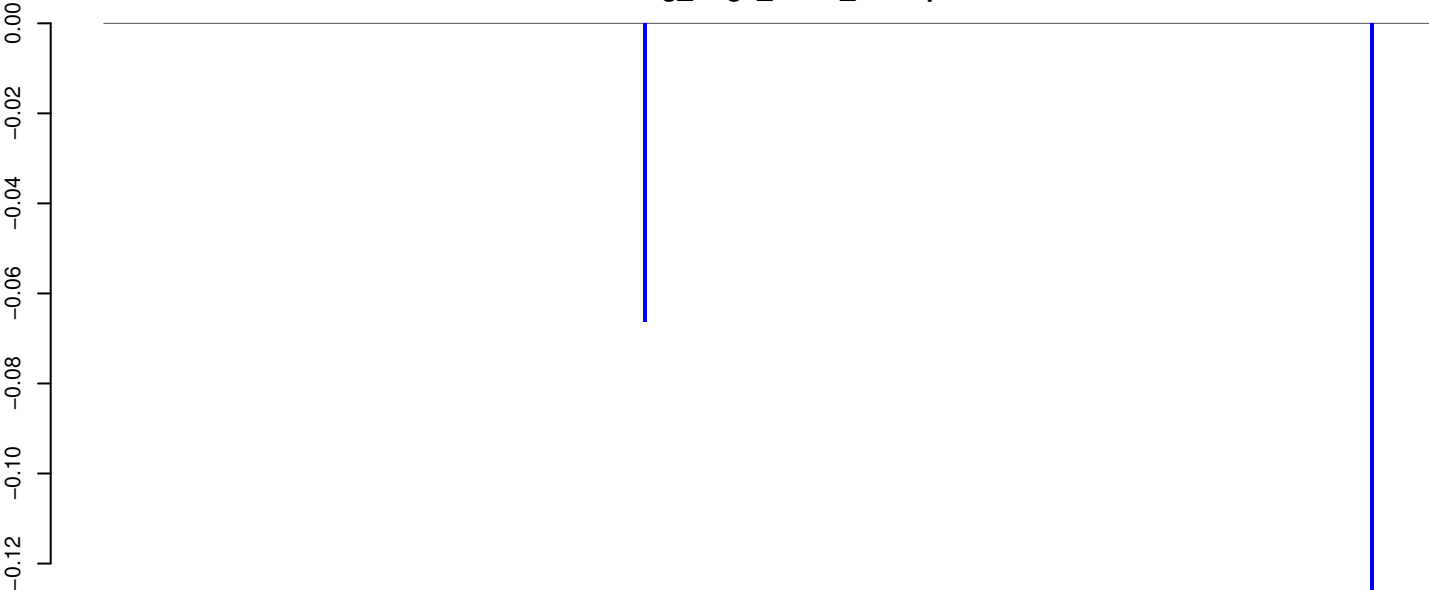
AeAeg_Aag2_BTV2_AK.18_23.rep



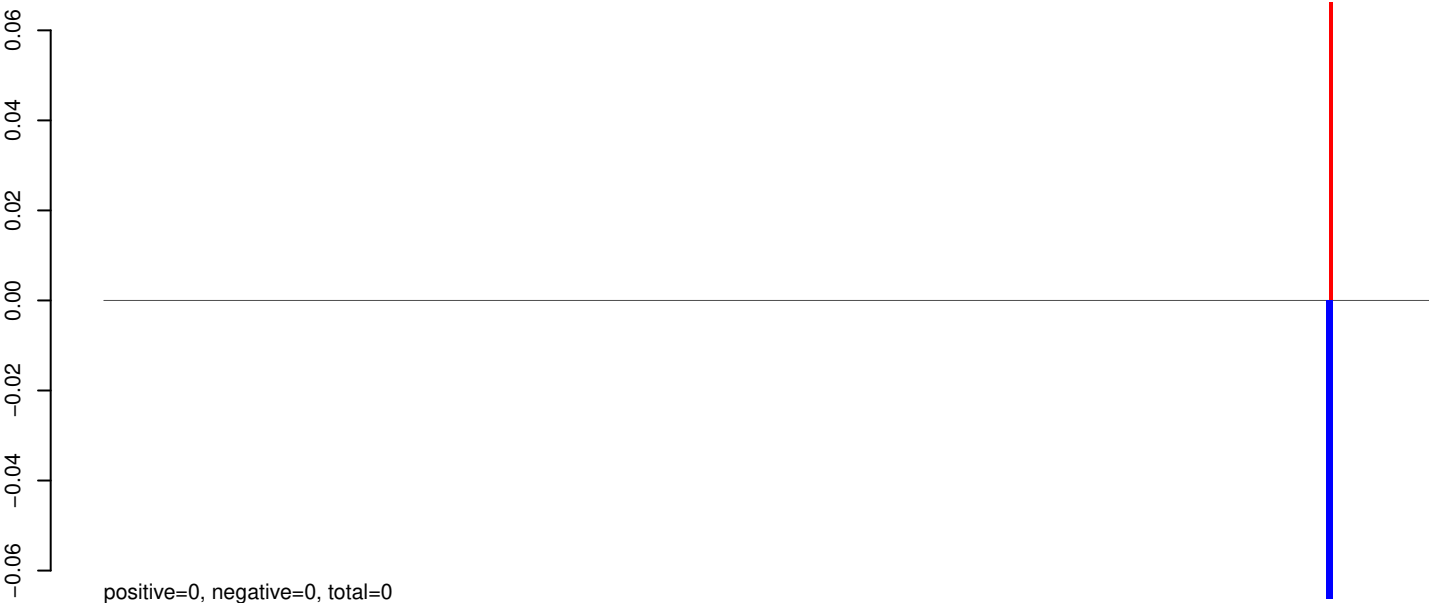
AeAeg_Aag2_BTV2_AK.24_35.rep



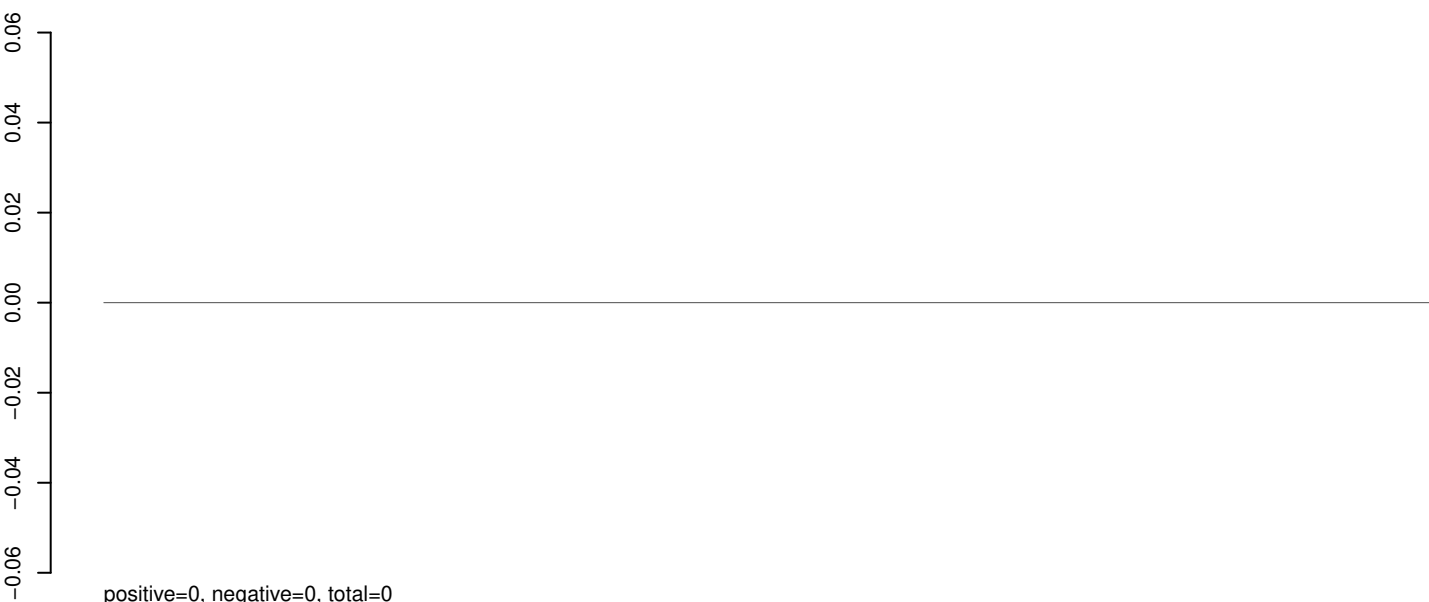
AeAeg_Aag2_BTV2_AK.rep



AeAeg_Aag2_BTV2_AK.18_23.rep



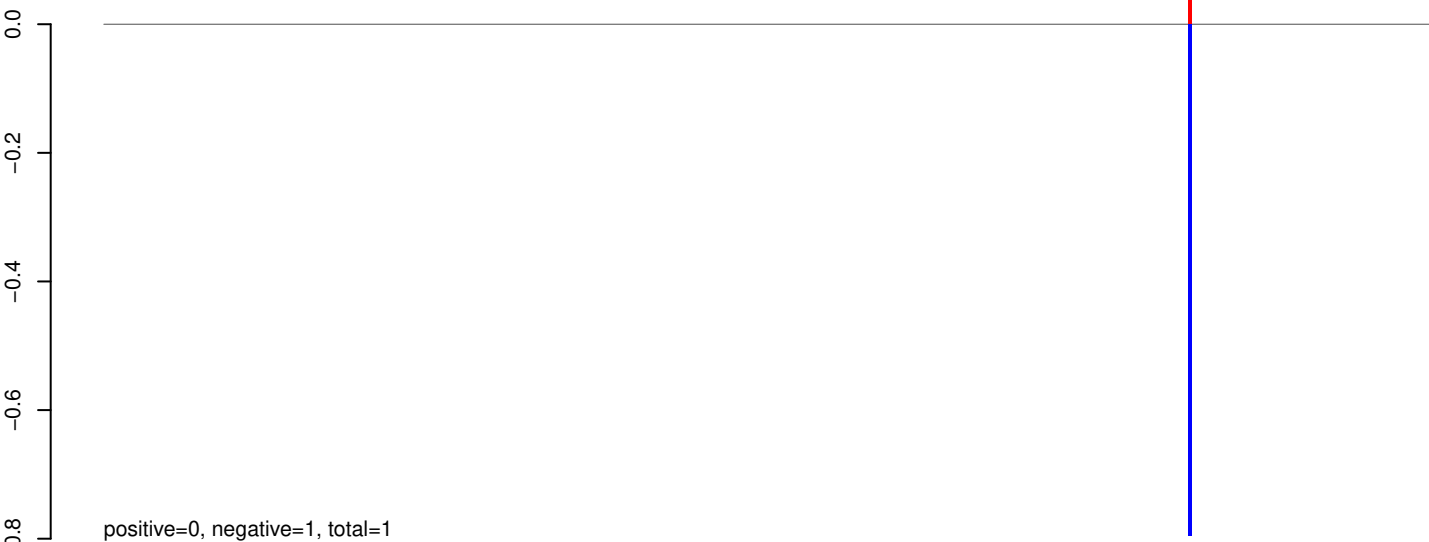
AeAeg_Aag2_BTV2_AK.24_35.rep



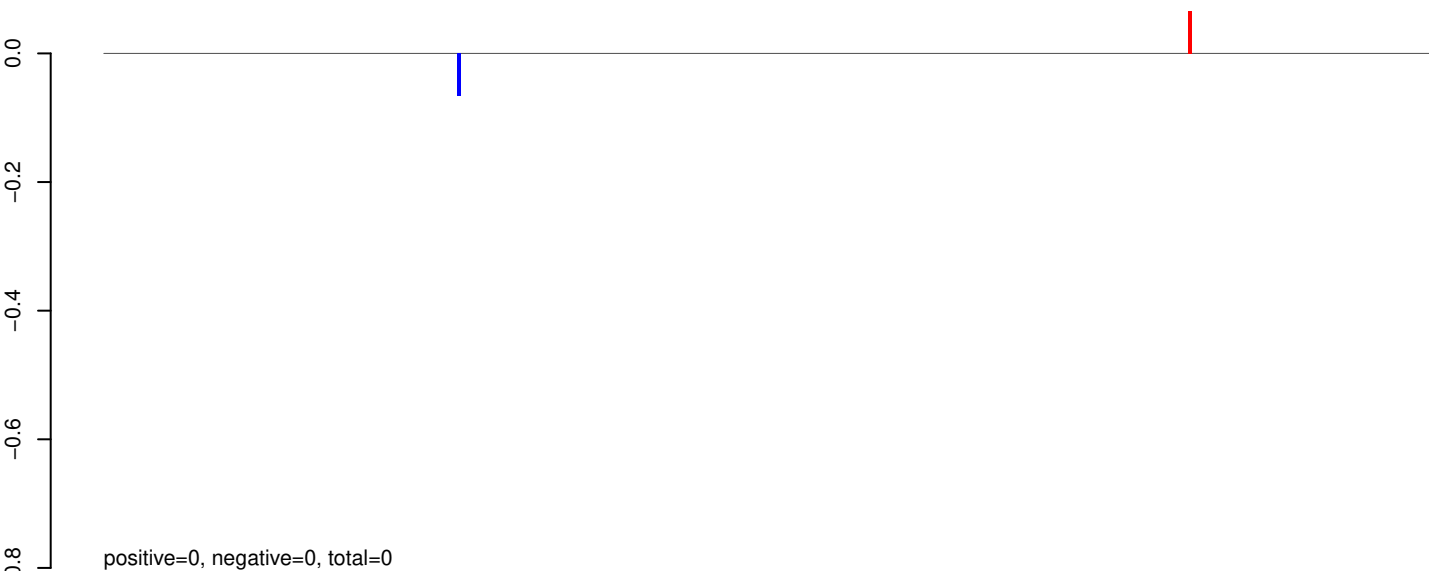
AeAeg_Aag2_BTV2_AK.rep



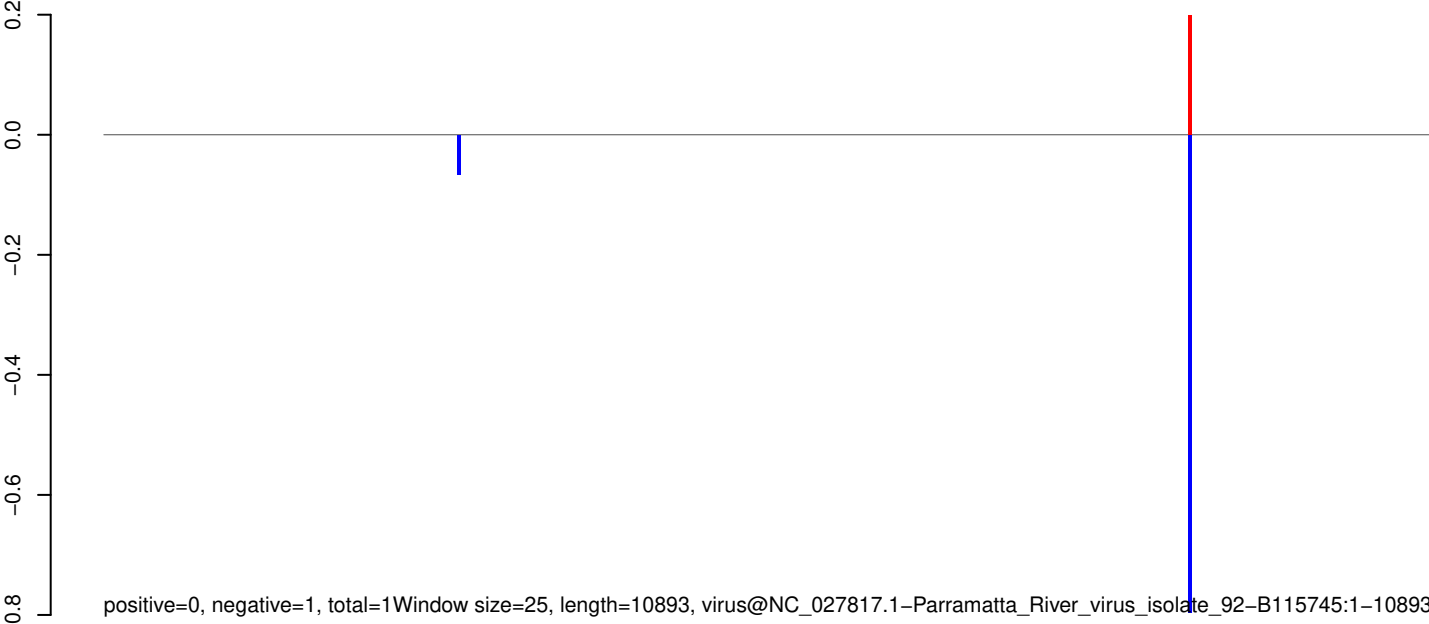
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



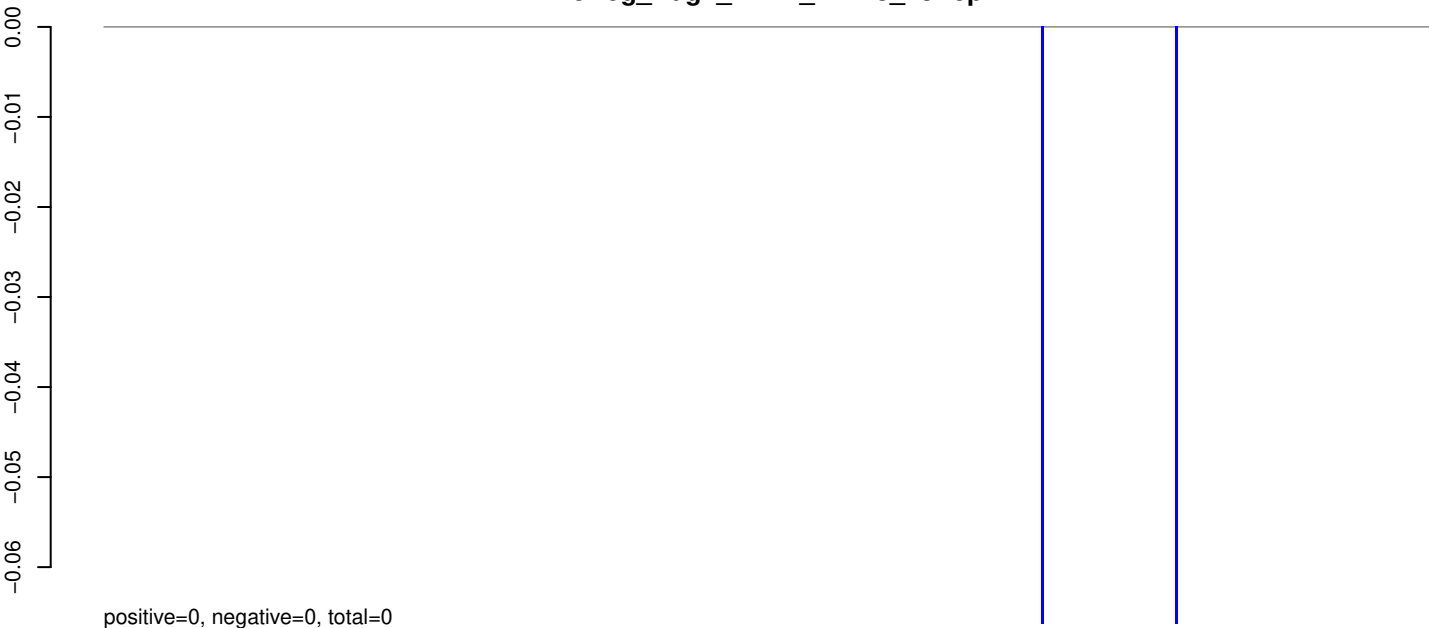
AeAeg_Aag2_BTV2_AK.rep



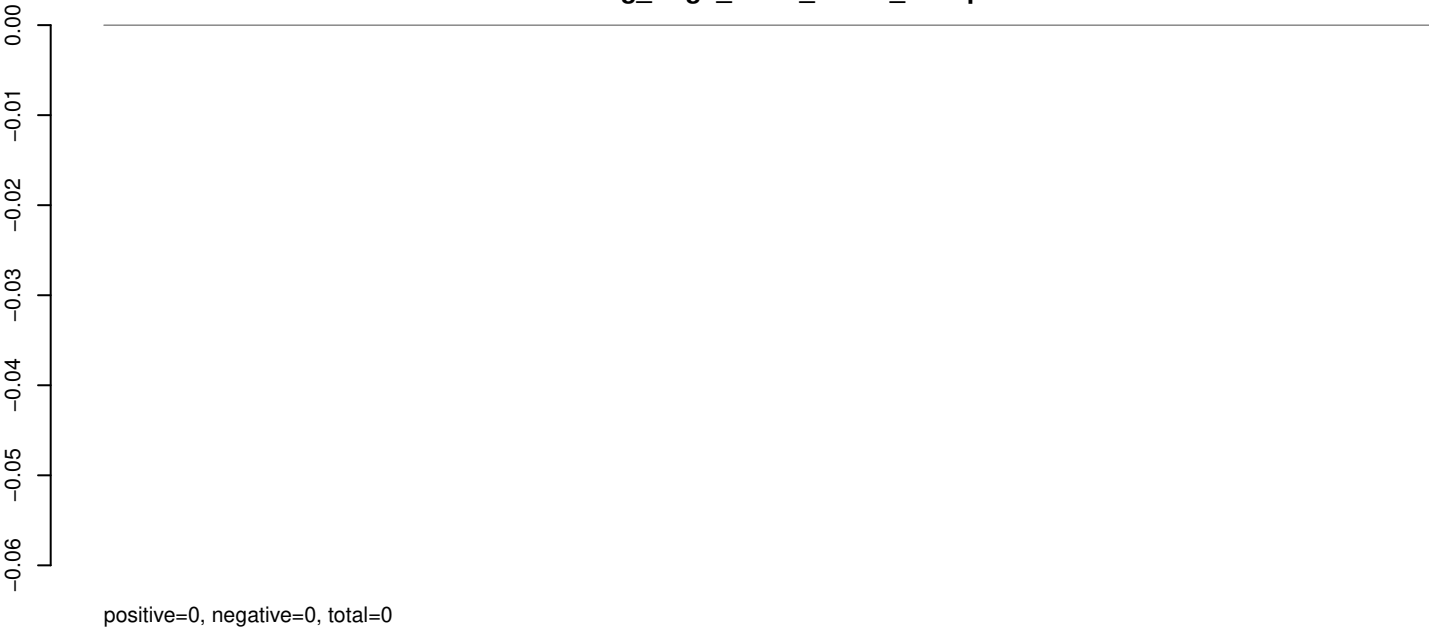
Window size=25, length=10893, virus@NC_027817.1-Parramatta_River_virus_isolate_92-B115745:1-10893

0 2000 4000 6000 8000 10000

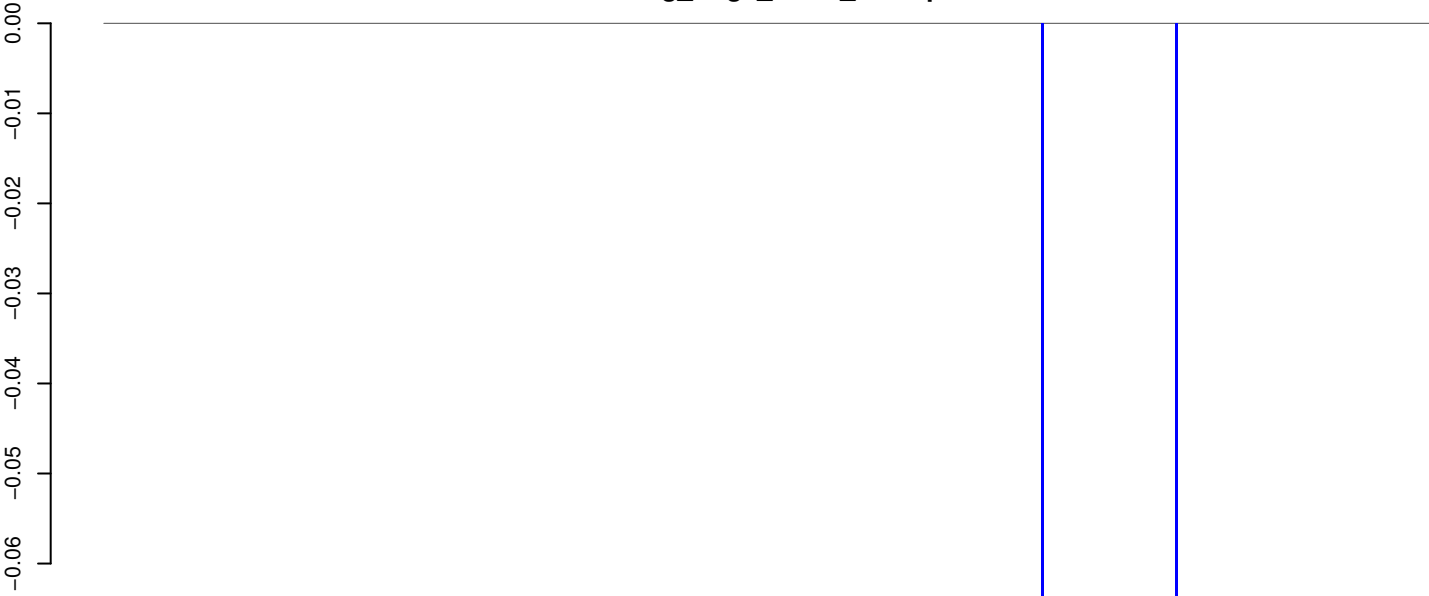
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



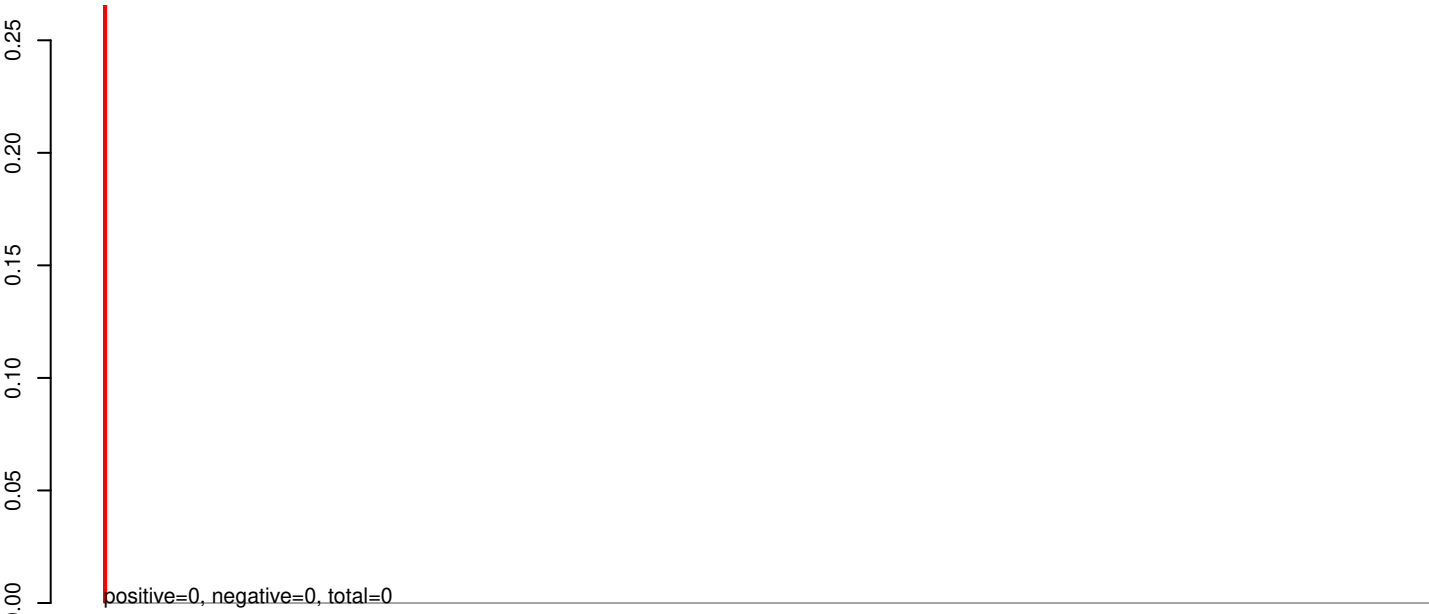
AeAeg_Aag2_BTV2_AK.rep



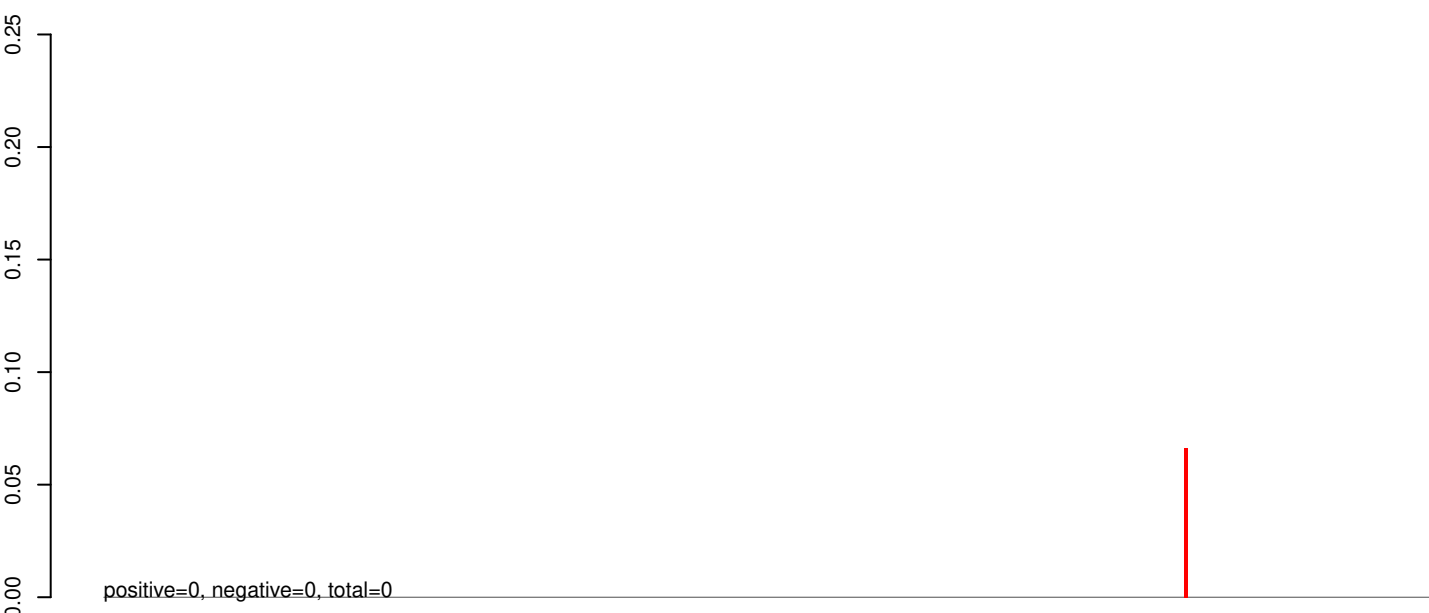
length=12390, coverage=0.161350 -Drosophila_melanogaster_sigma_virus_AP30_N_P_X_M_G_and_L_genes_genomic_RNA_isolate:1-12390

0 2000 4000 6000 8000 10000 12000

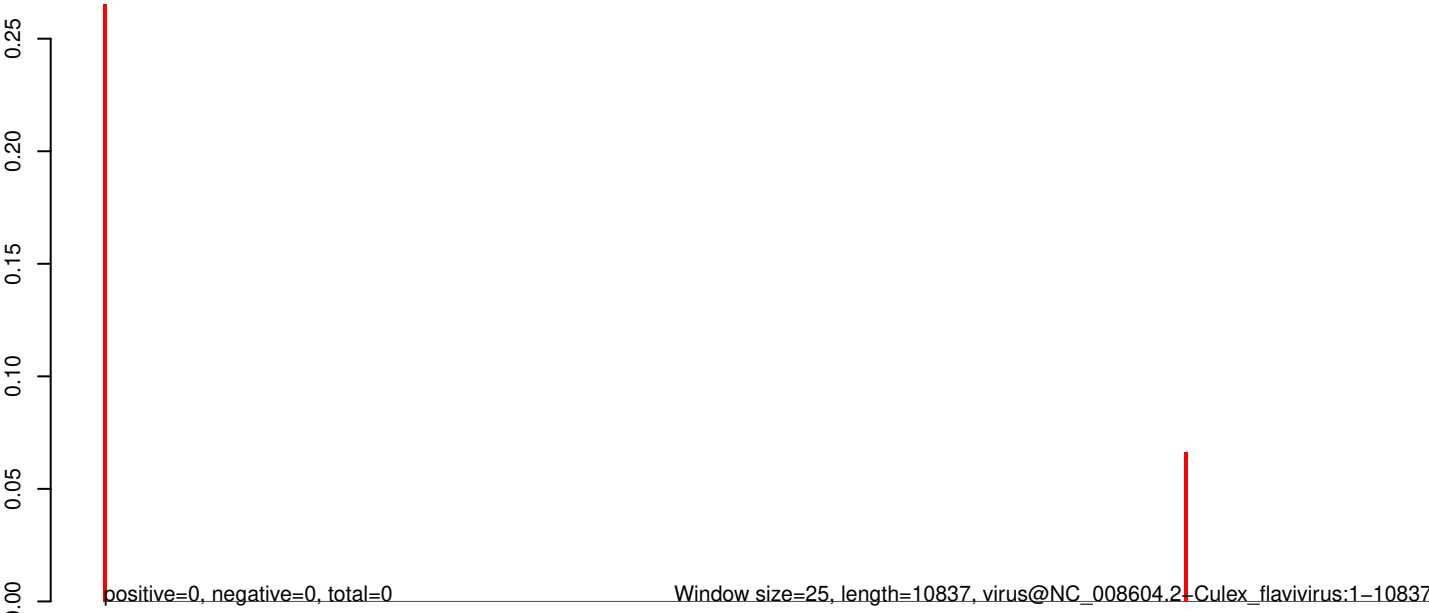
AeAeg_Aag2_BTV2_AK.18_23.rep



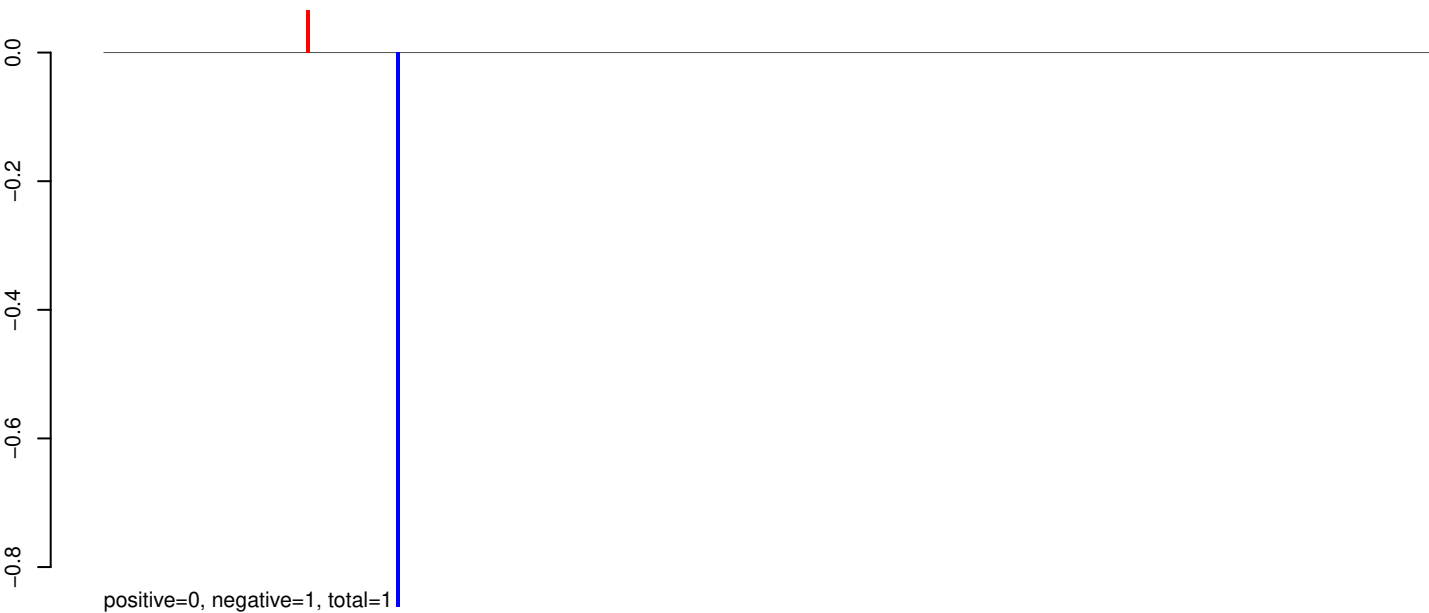
AeAeg_Aag2_BTV2_AK.24_35.rep



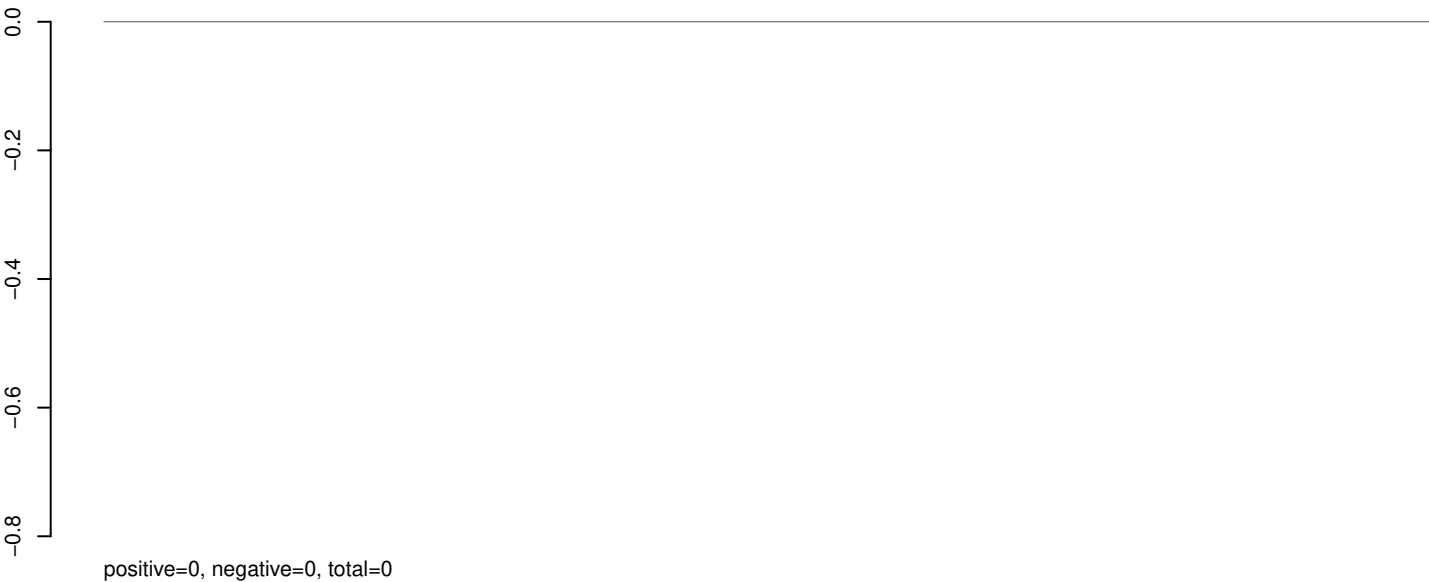
AeAeg_Aag2_BTV2_AK.rep



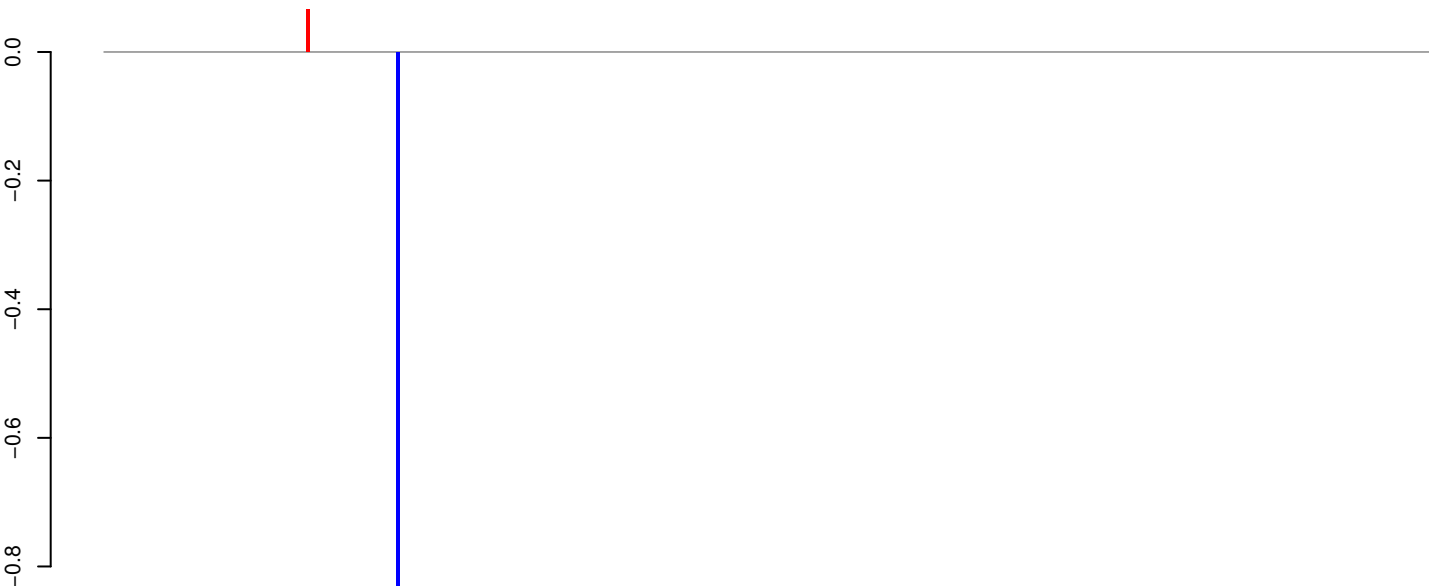
AeAeg_Aag2_BTV2_AK.18_23.rep



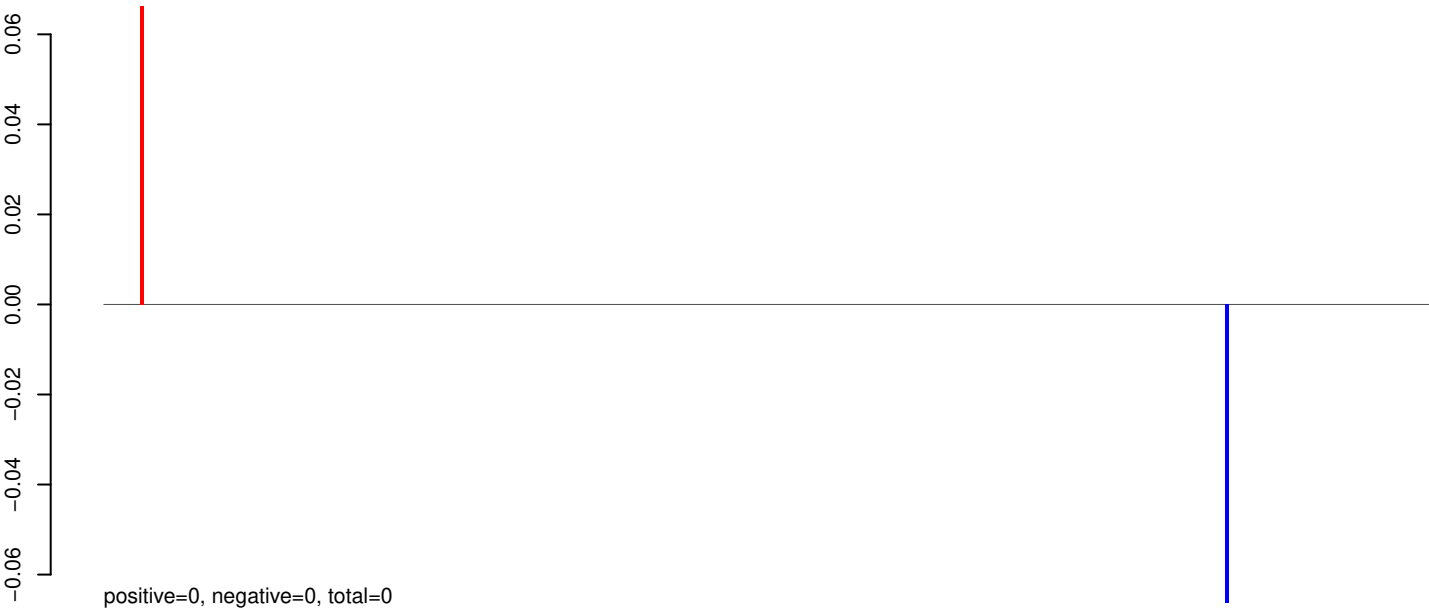
AeAeg_Aag2_BTV2_AK.24_35.rep



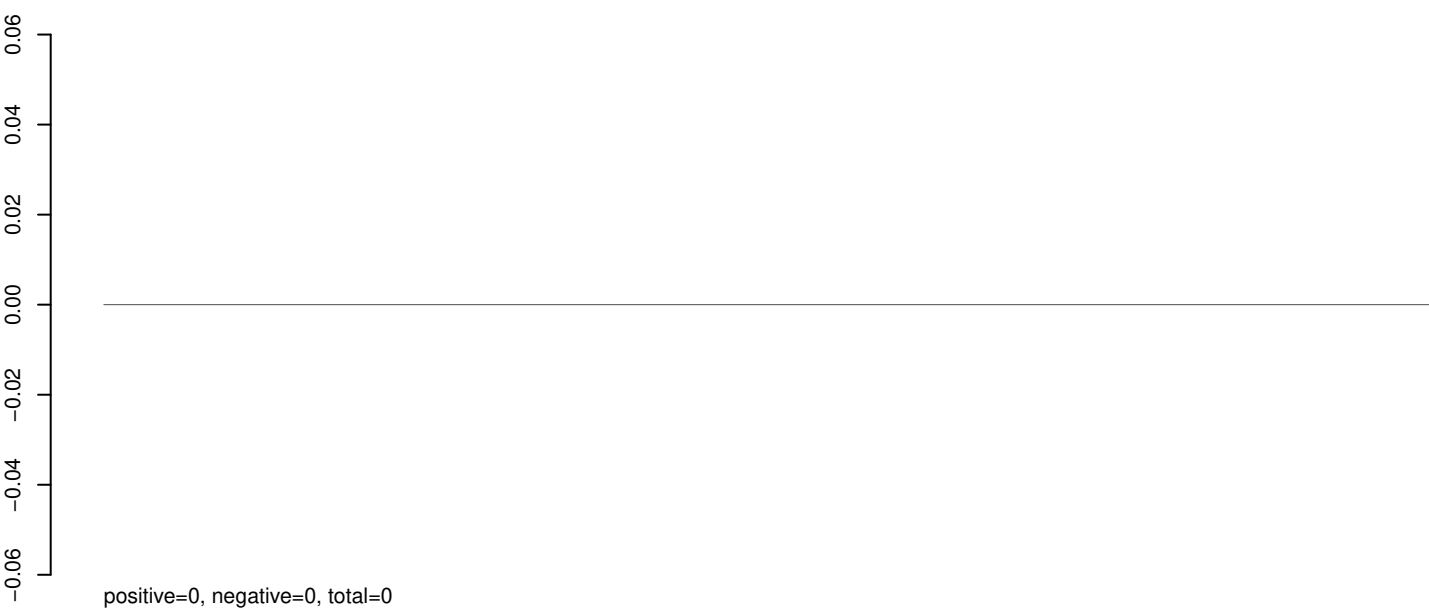
AeAeg_Aag2_BTV2_AK.rep



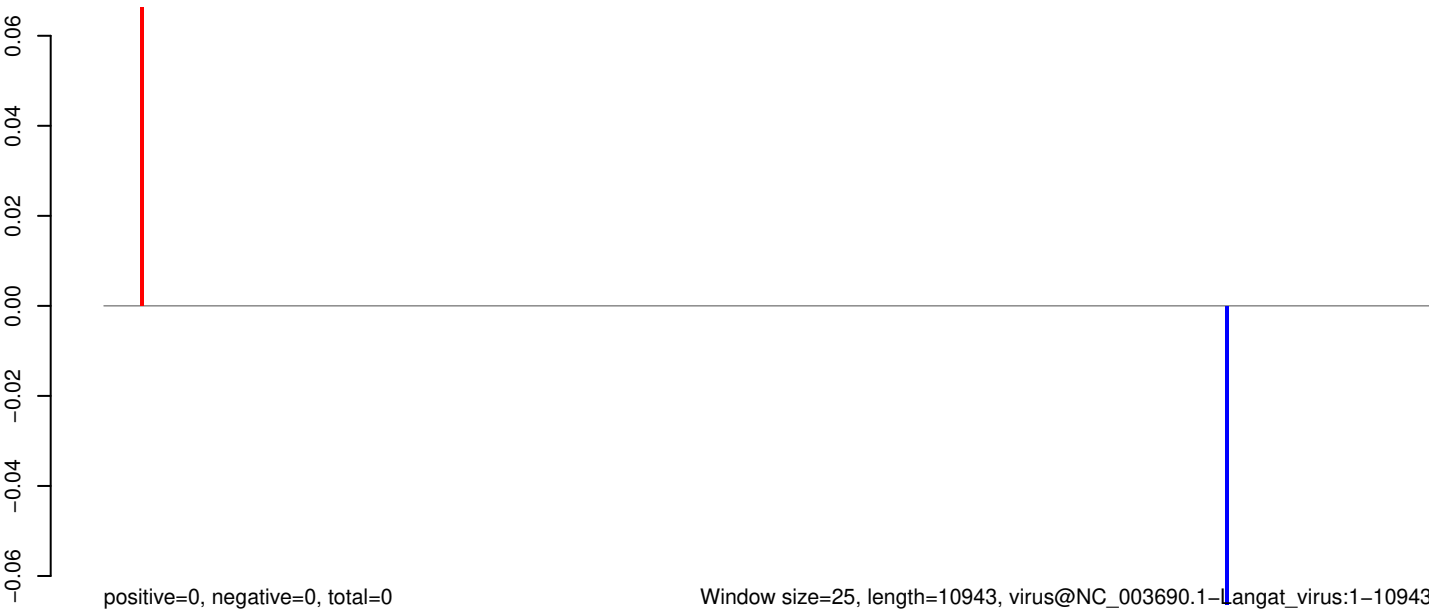
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



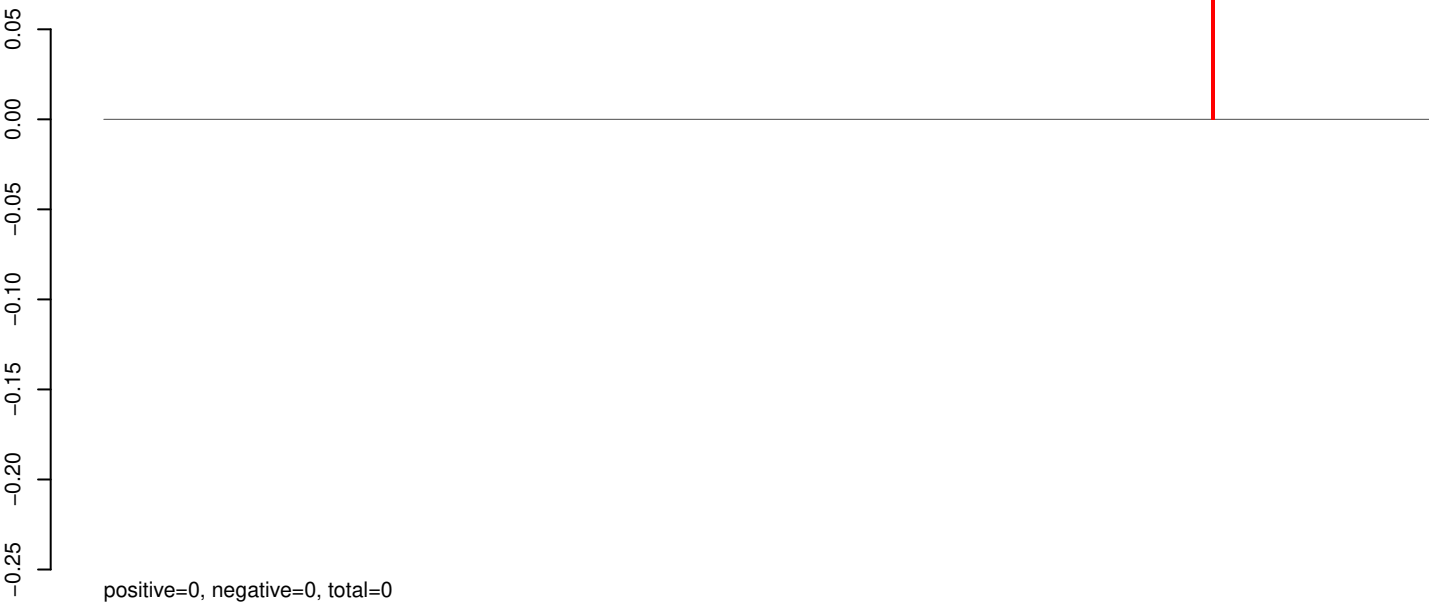
AeAeg_Aag2_BTV2_AK.rep



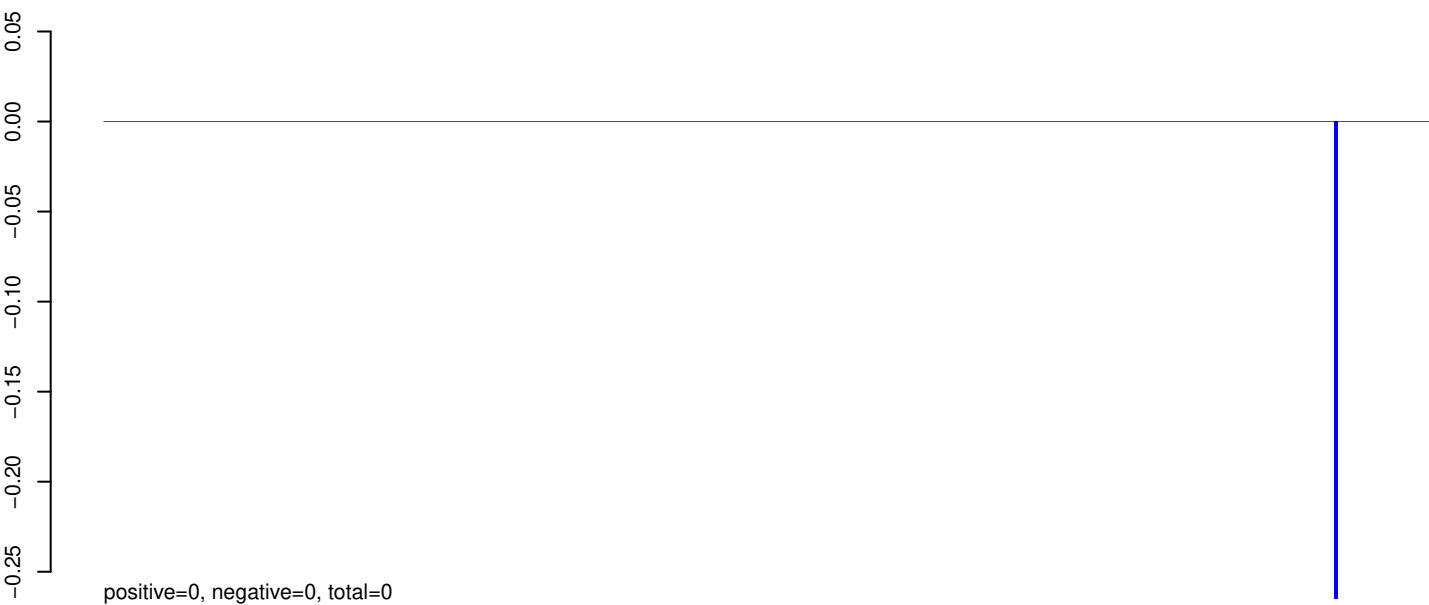
Window size=25, length=10943, virus@NC_003690.1-Langat_virus:1-10943

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

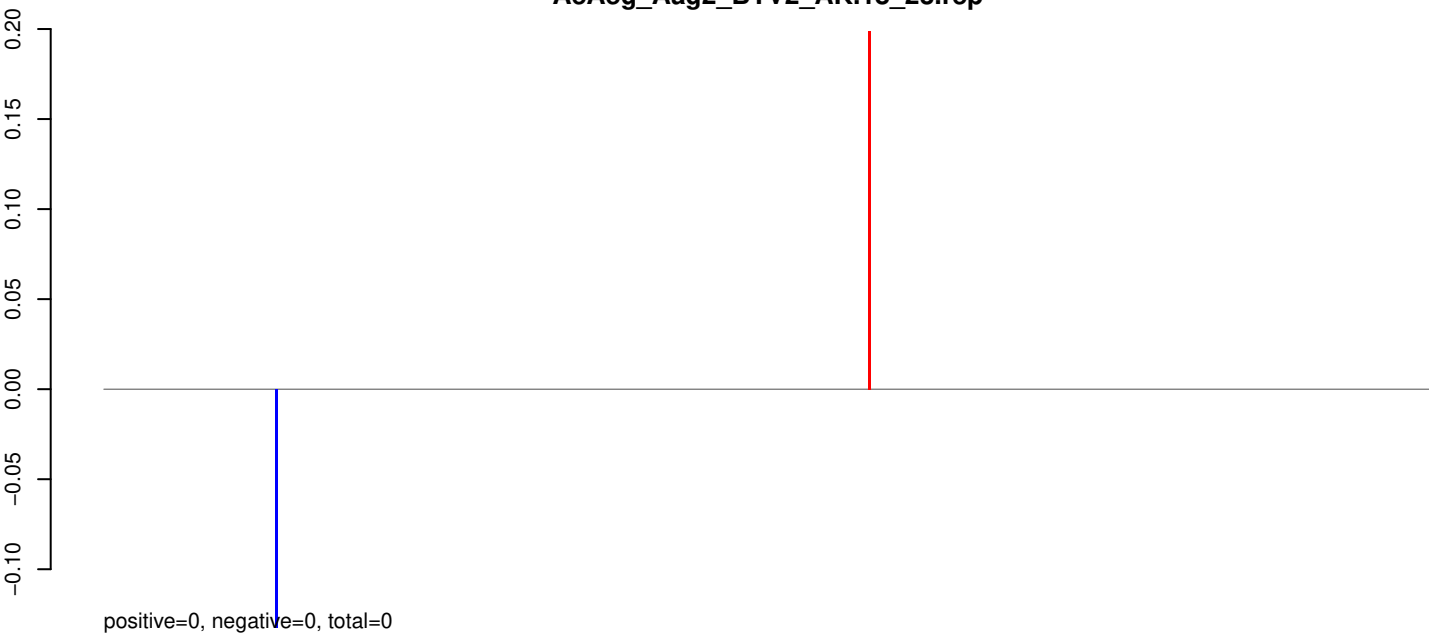


AeAeg_Aag2_BTV2_AK.rep

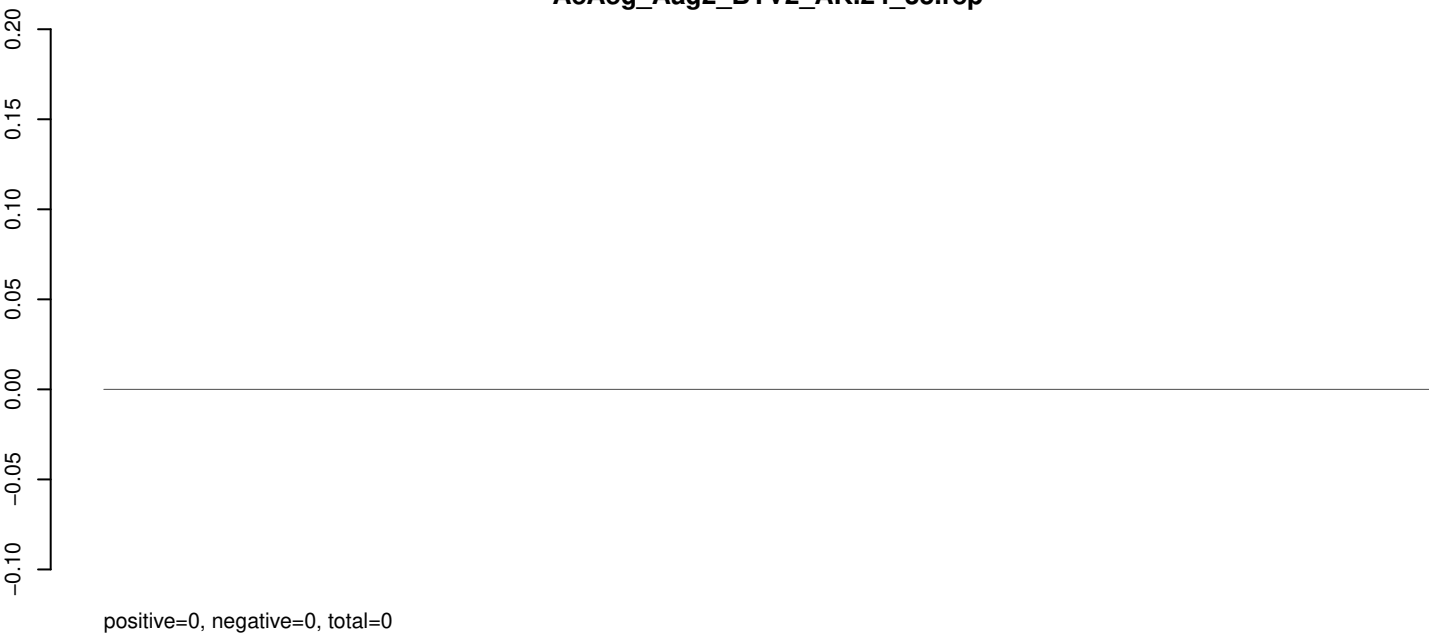


0 2000 4000 6000 8000 10000

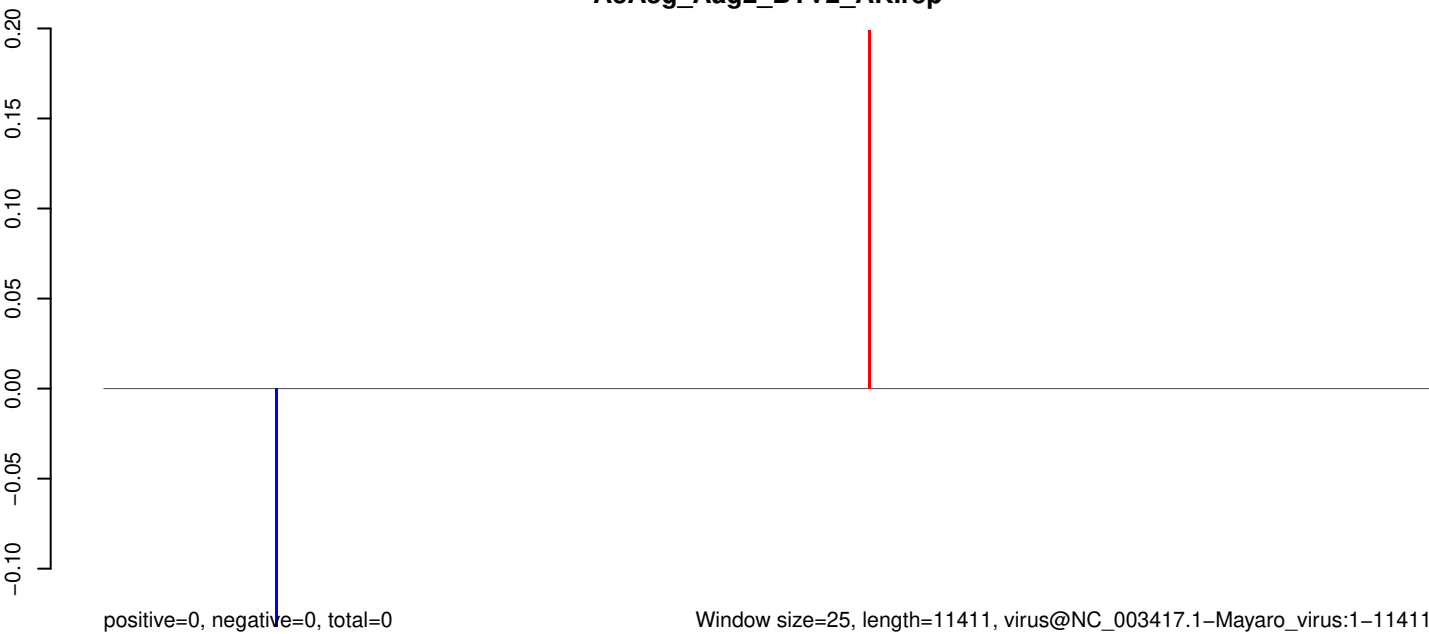
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



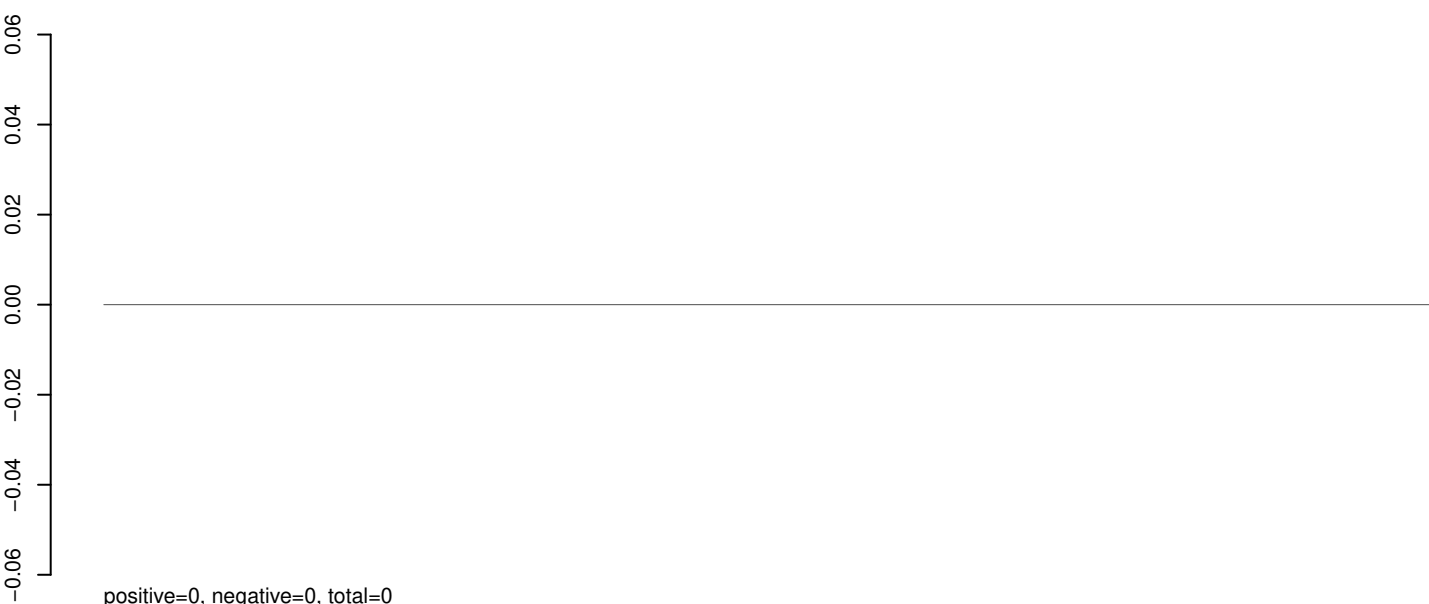
AeAeg_Aag2_BTV2_AK.rep



AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



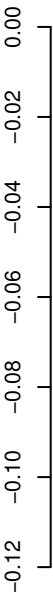
AeAeg_Aag2_BTV2_AK.rep



Window size=25, length=10707, virus@NC_001475.2-Dengue_virus_3:1-10707

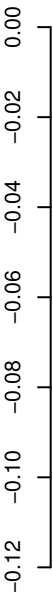
0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BTV2_AK.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BTV2_AK.rep

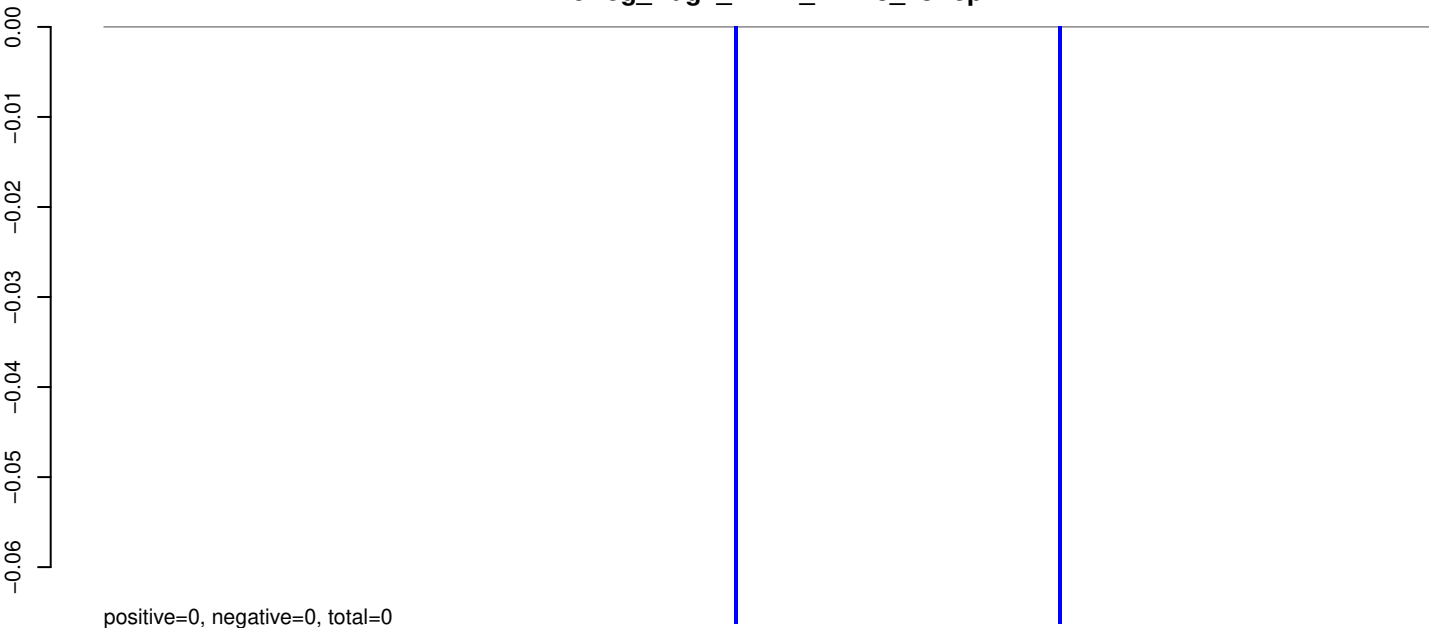


positive=0, negative=0, total=0

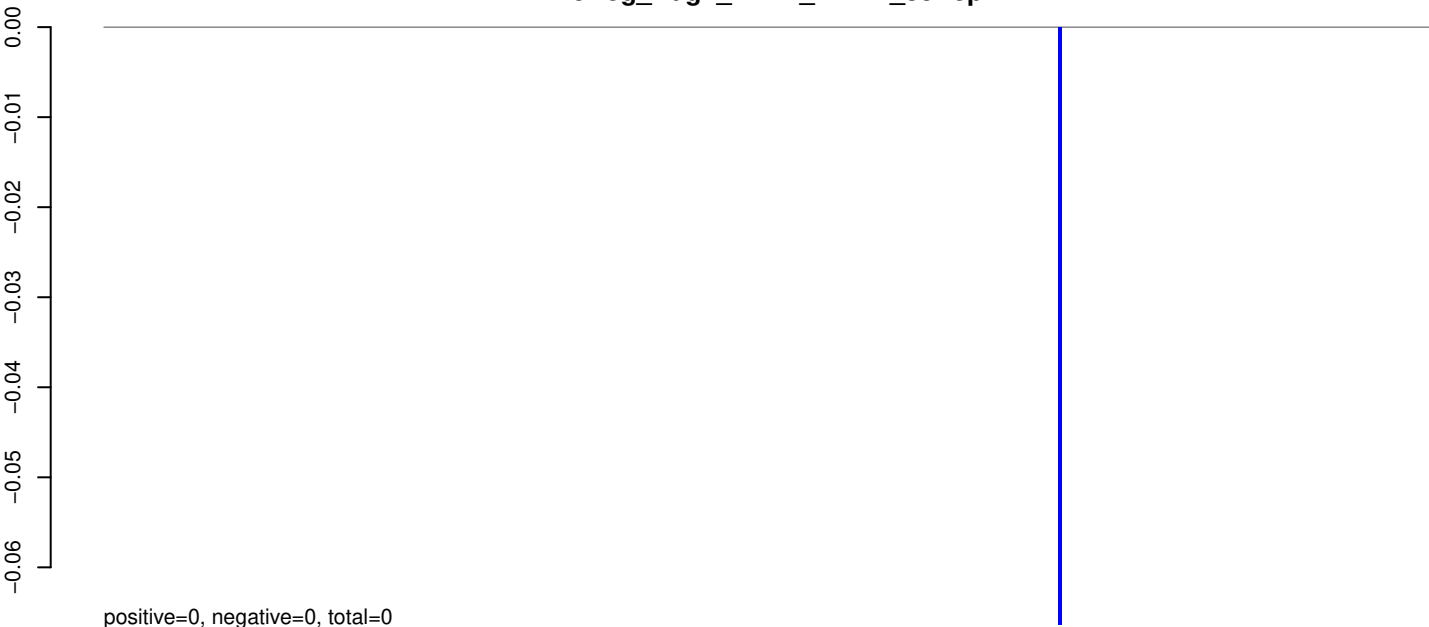
Window size=25, length=11014, virus@NC_000943.1-Murray_Valley_encephalitis_virus:1-11014



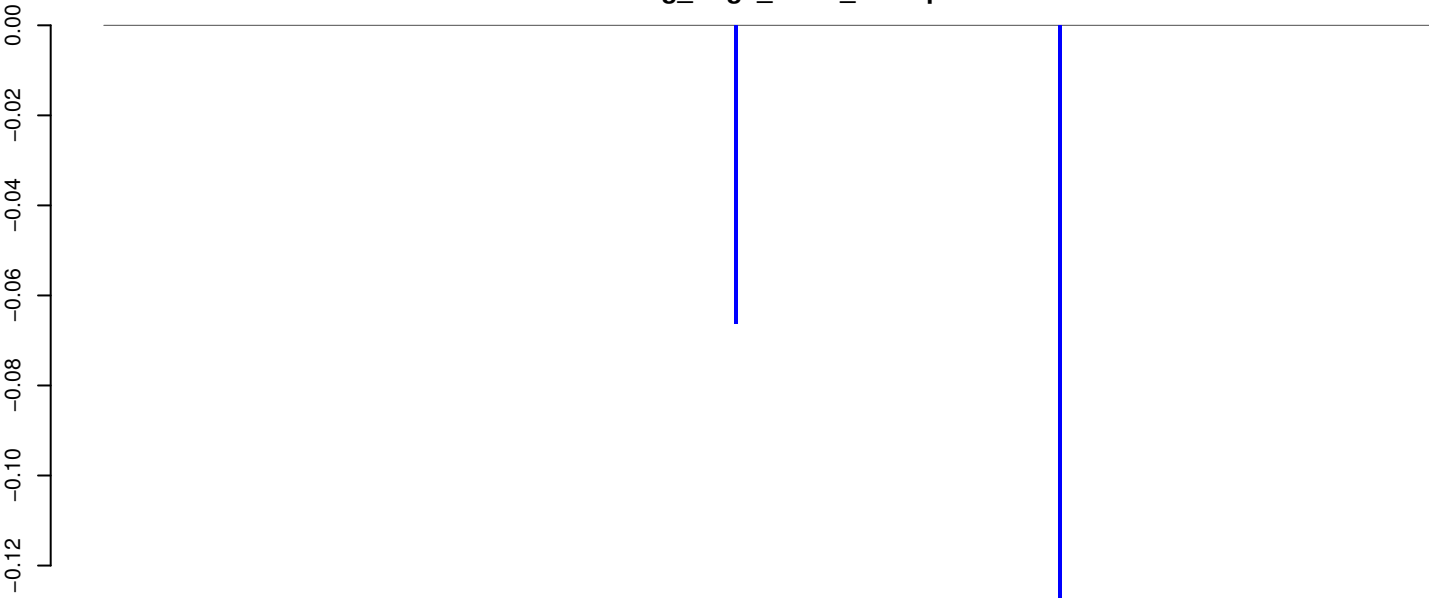
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



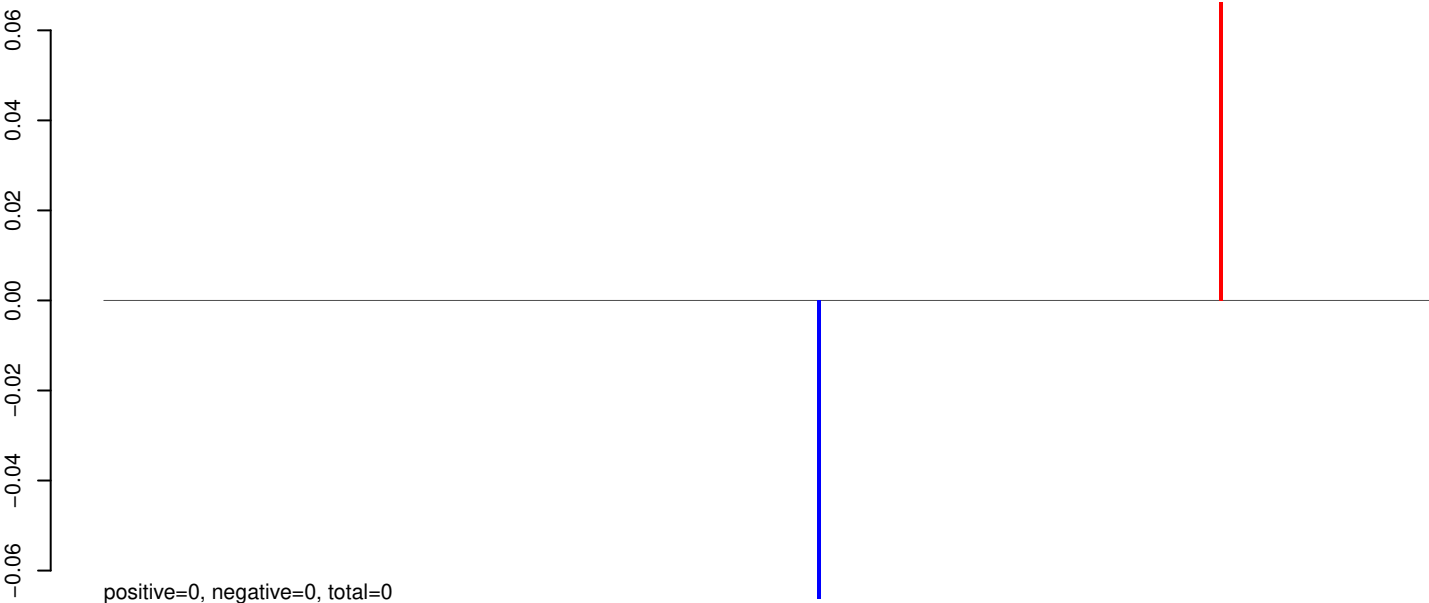
AeAeg_Aag2_BTV2_AK.rep



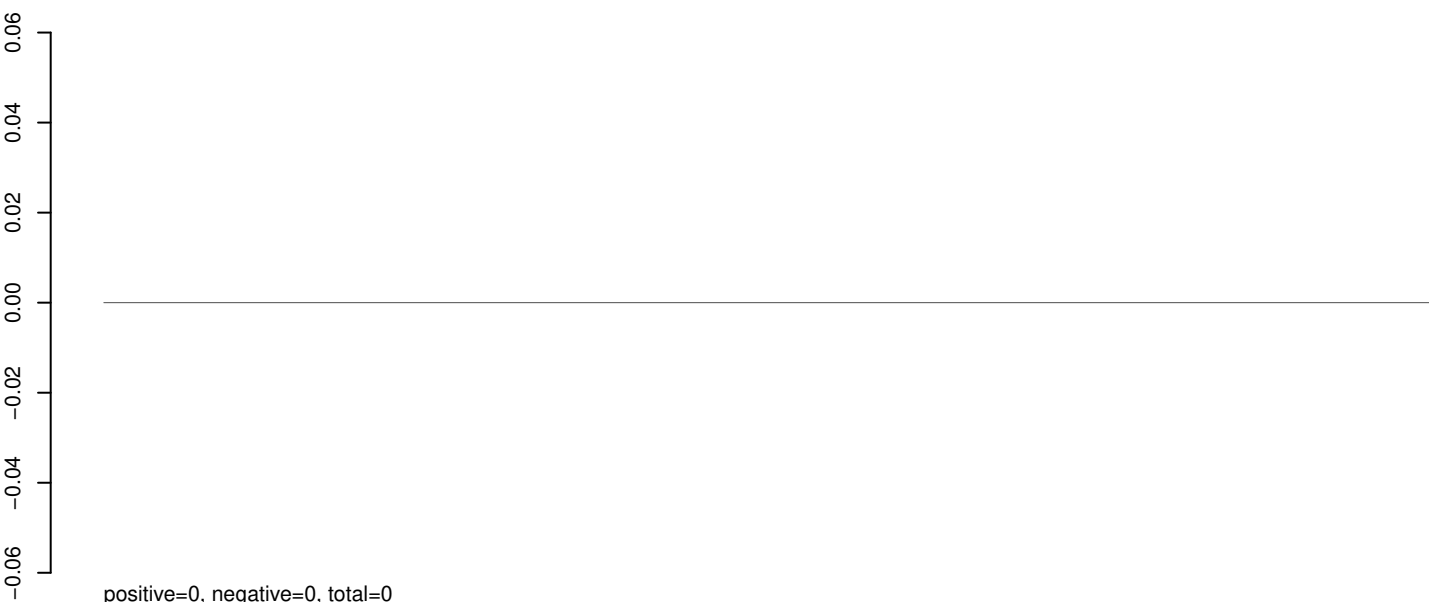
Window size=25, length=10749, virus@KX669682.1-Calbertado_virus_isolate_Cal_mos_2_polyprotein_gene:1-10749

0 2000 4000 6000 8000 10000

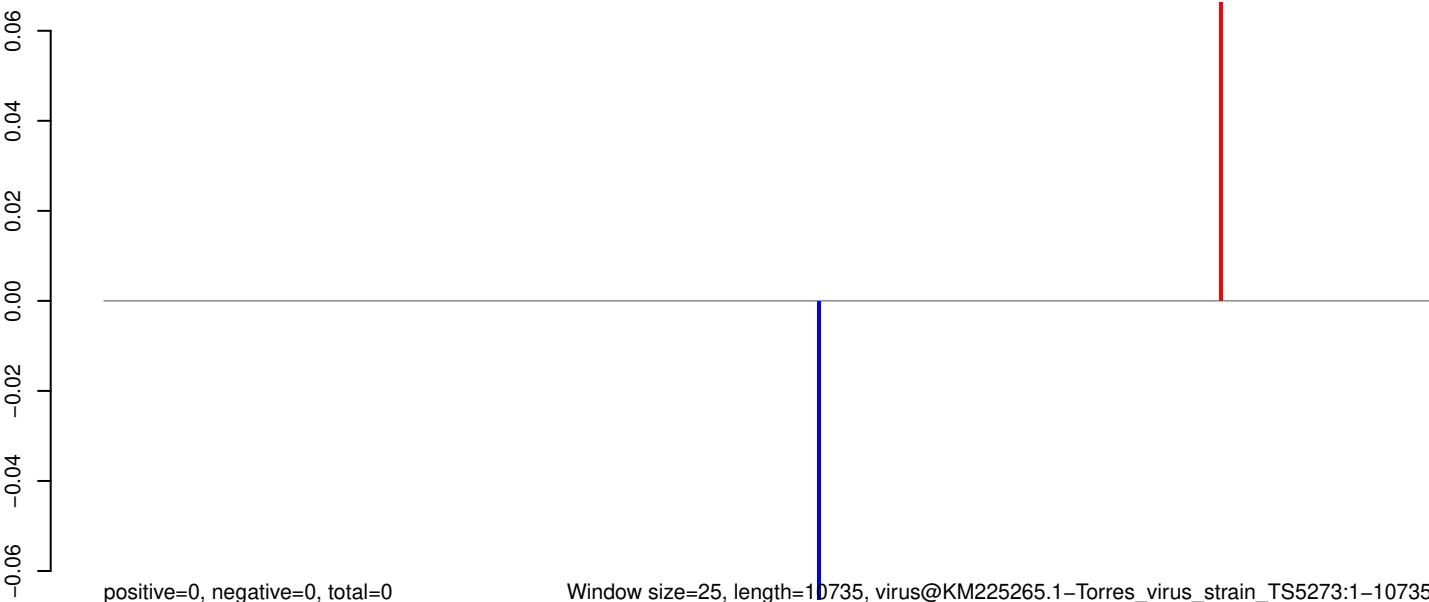
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

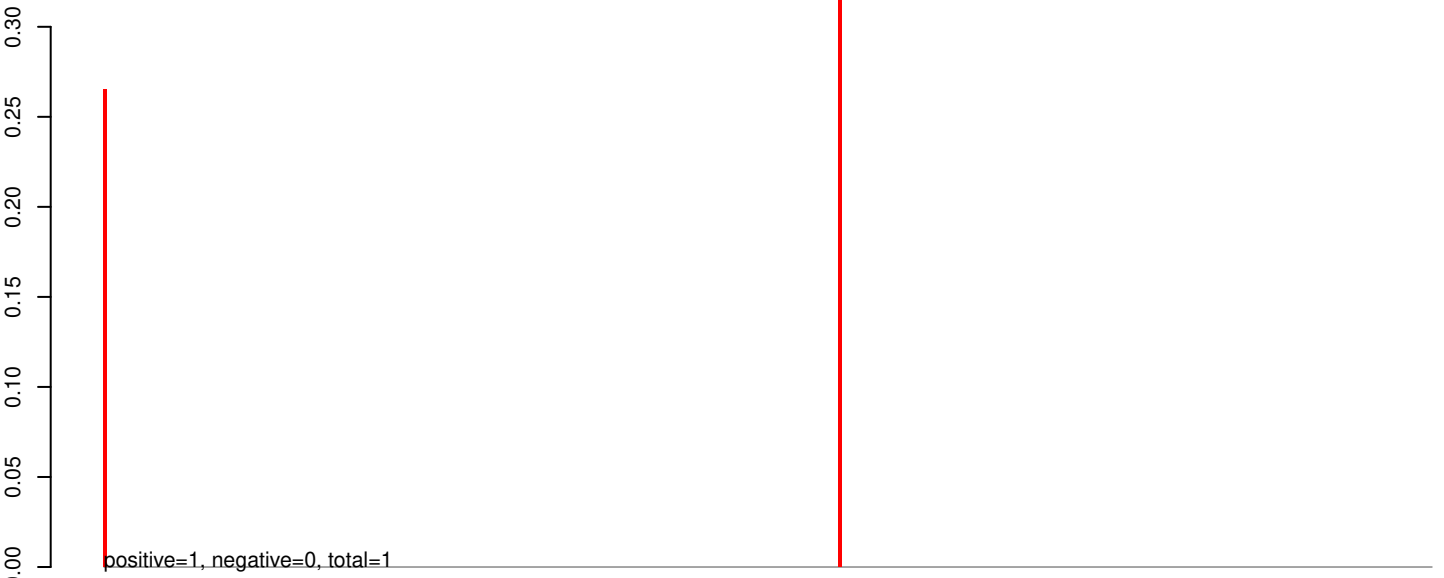


AeAeg_Aag2_BTV2_AK.rep

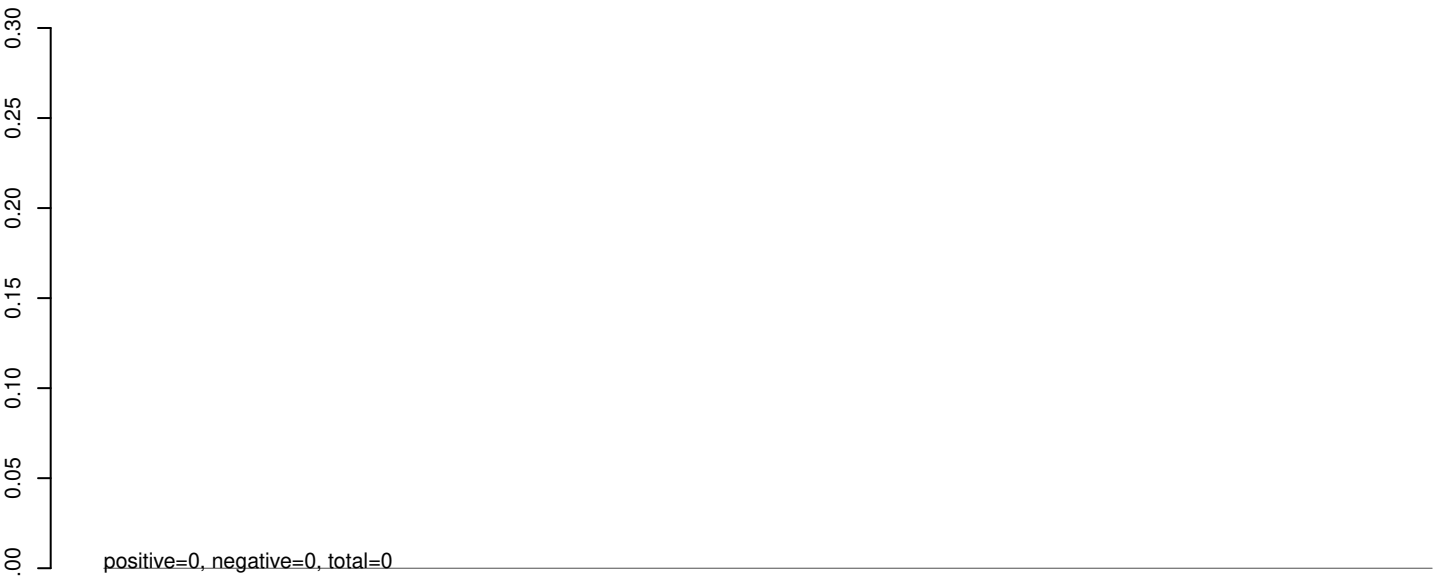


0 2000 4000 6000 8000 10000

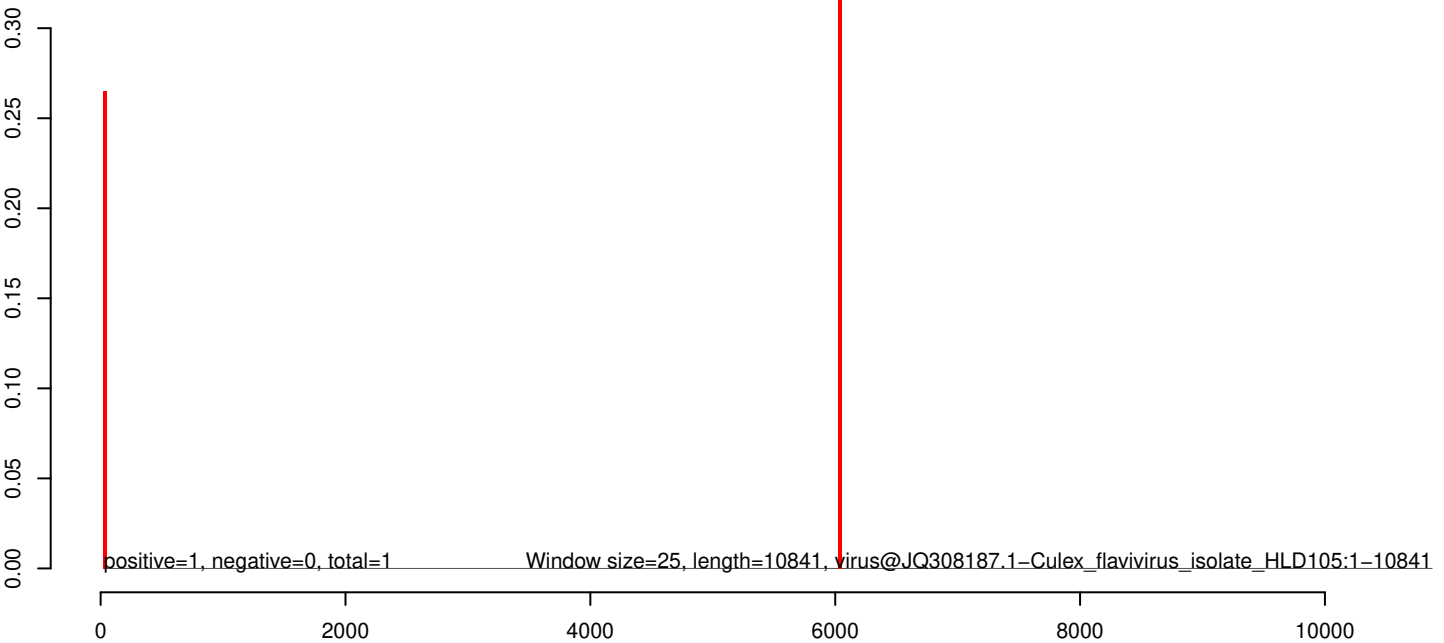
AeAeg_Aag2_BTV2_AK.18_23.rep



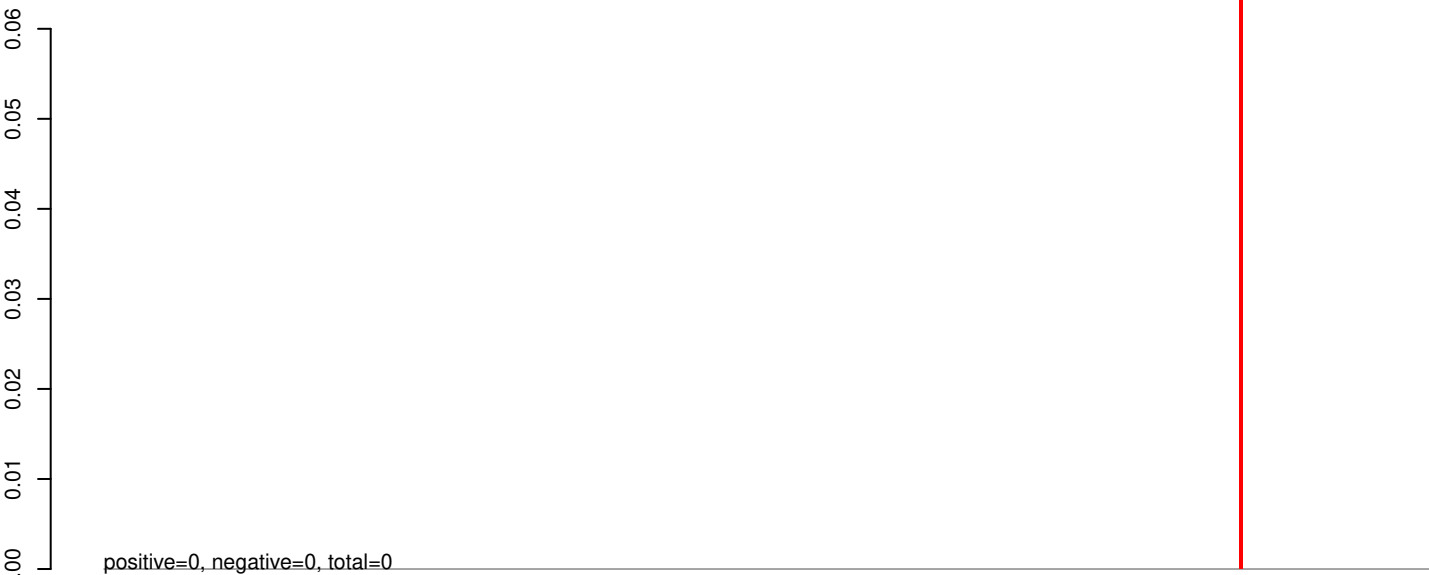
AeAeg_Aag2_BTV2_AK.24_35.rep



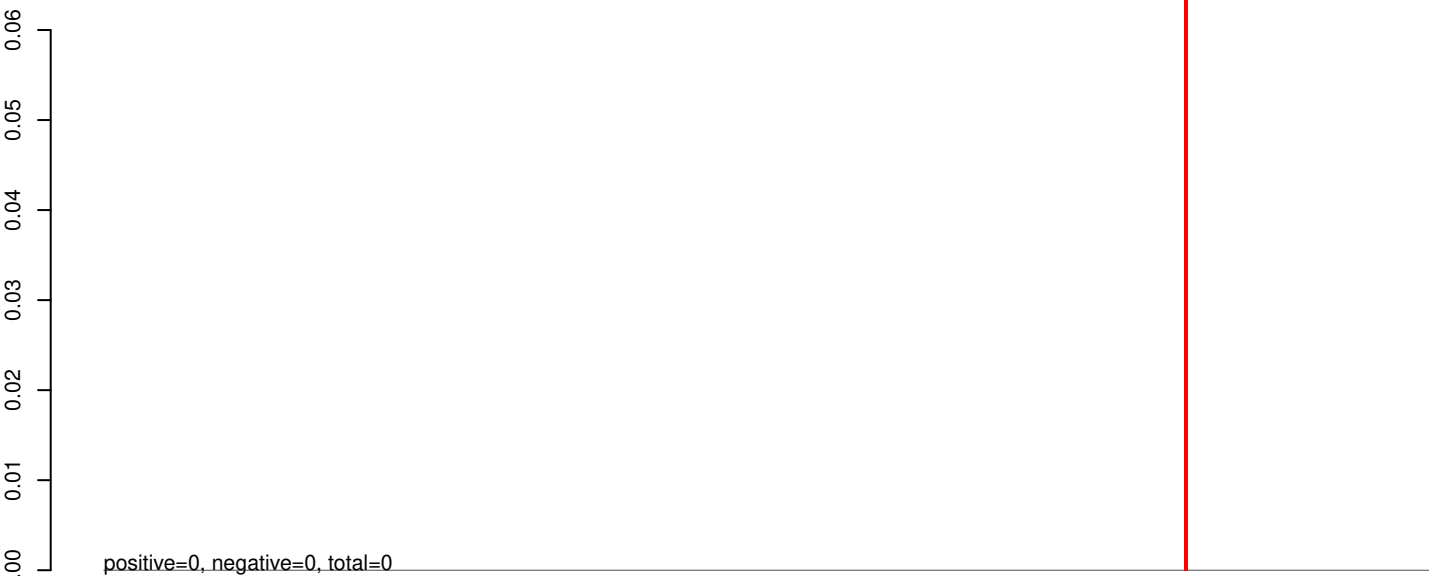
AeAeg_Aag2_BTV2_AK.rep



AeAeg_Aag2_BTV2_AK.18_23.rep



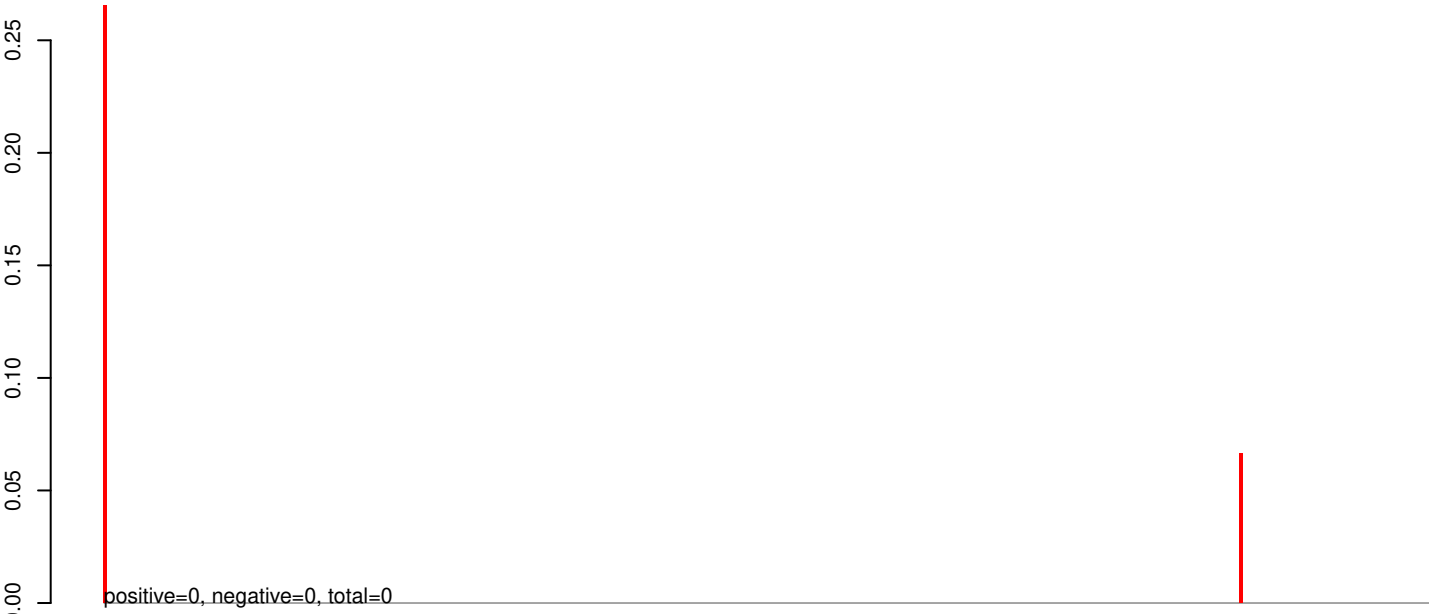
AeAeg_Aag2_BTV2_AK.24_35.rep



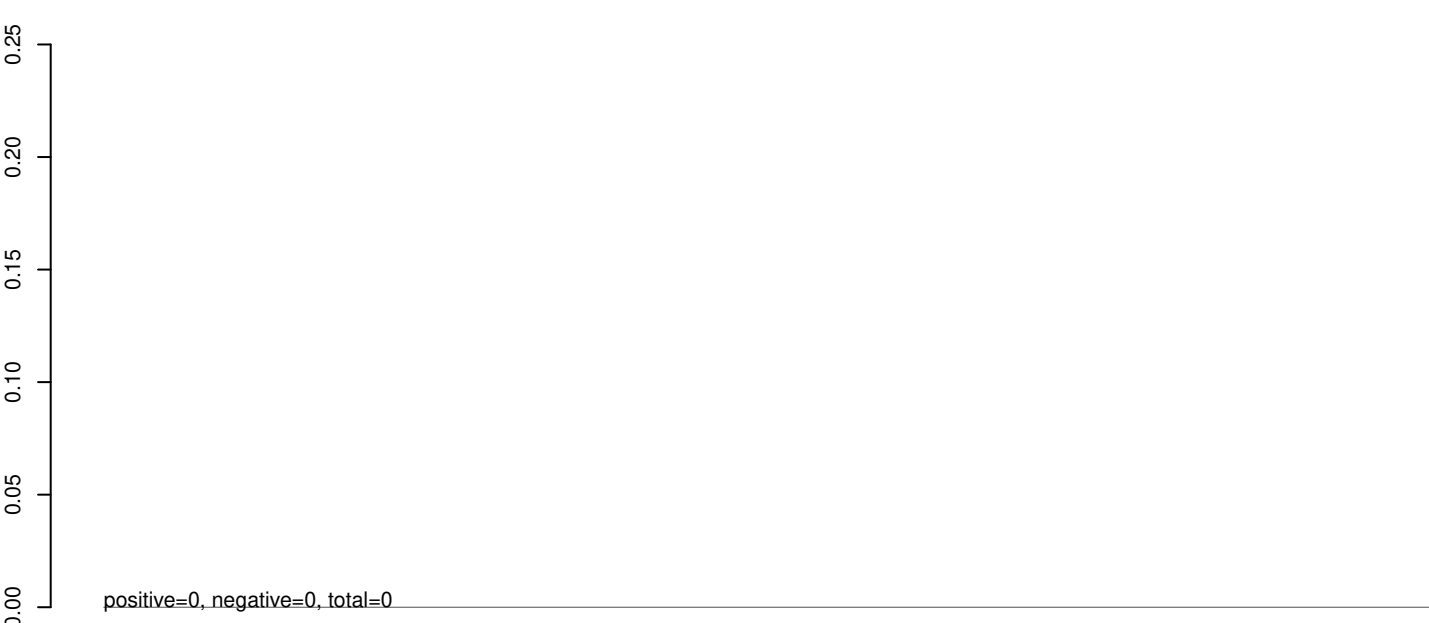
AeAeg_Aag2_BTV2_AK.rep



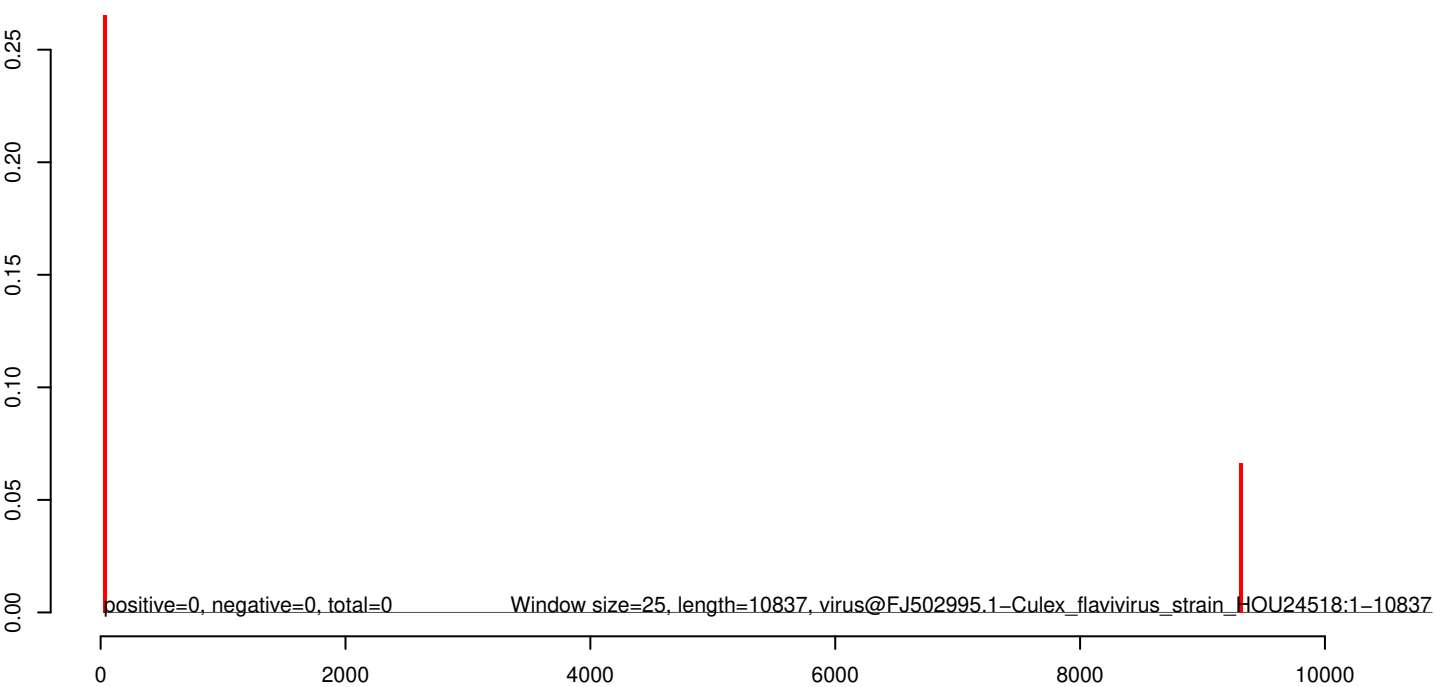
AeAeg_Aag2_BTV2_AK.18_23.rep



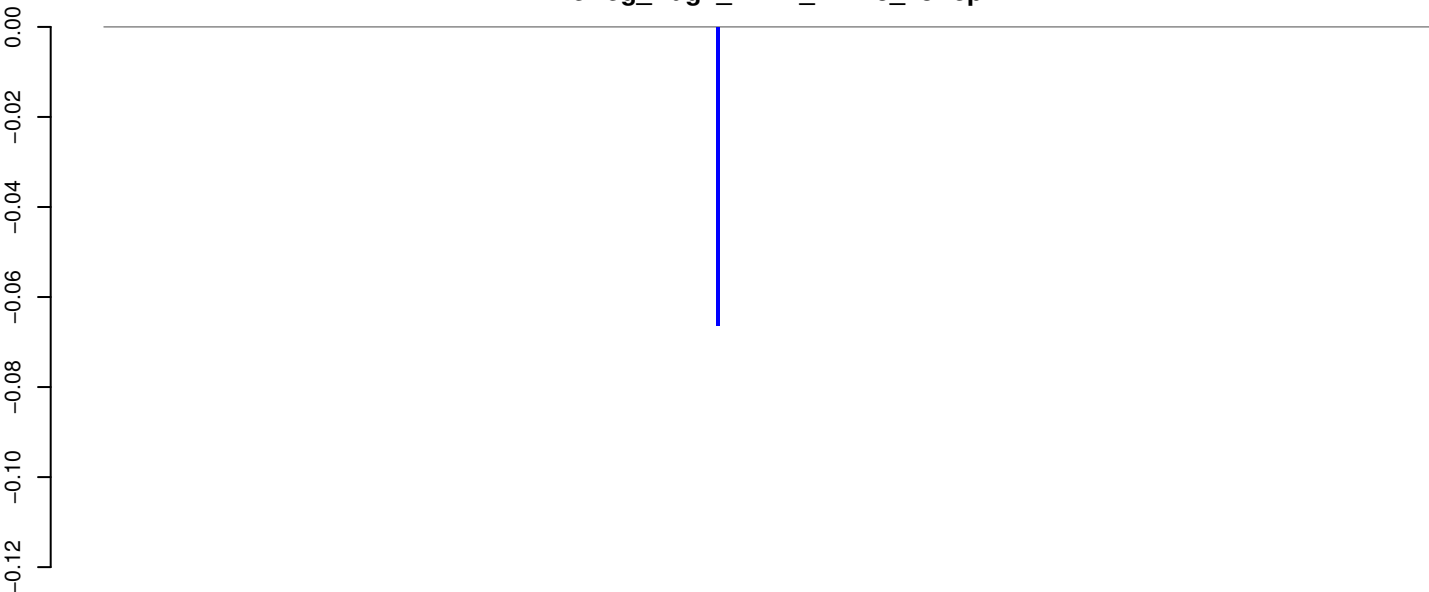
AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep

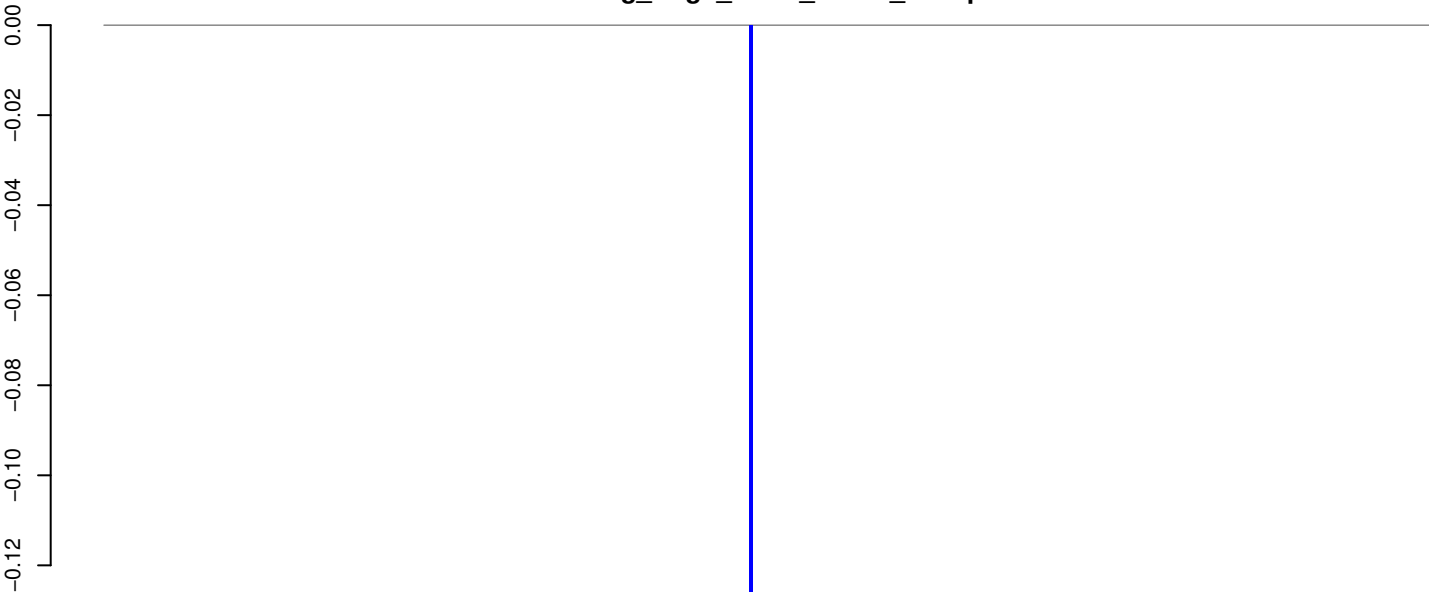


AeAeg_Aag2_BTV2_AK.18_23.rep



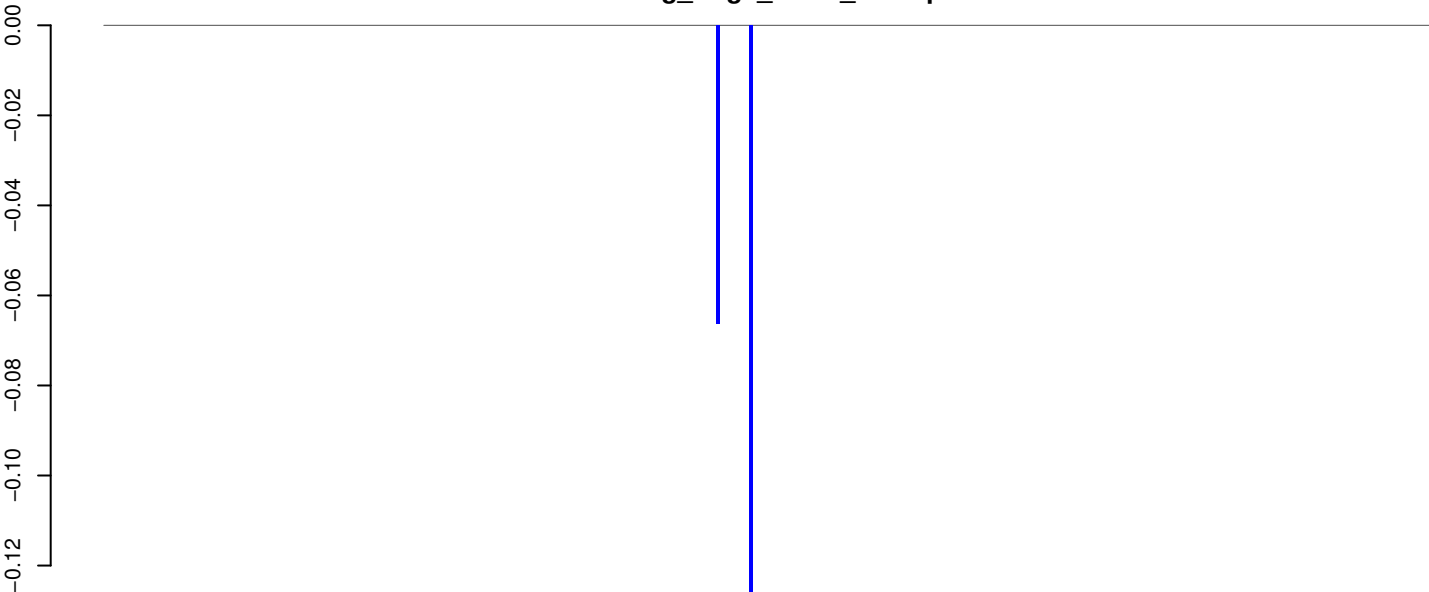
positive=0, negative=0, total=0

AeAeg_Aag2_BTV2_AK.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BTV2_AK.rep

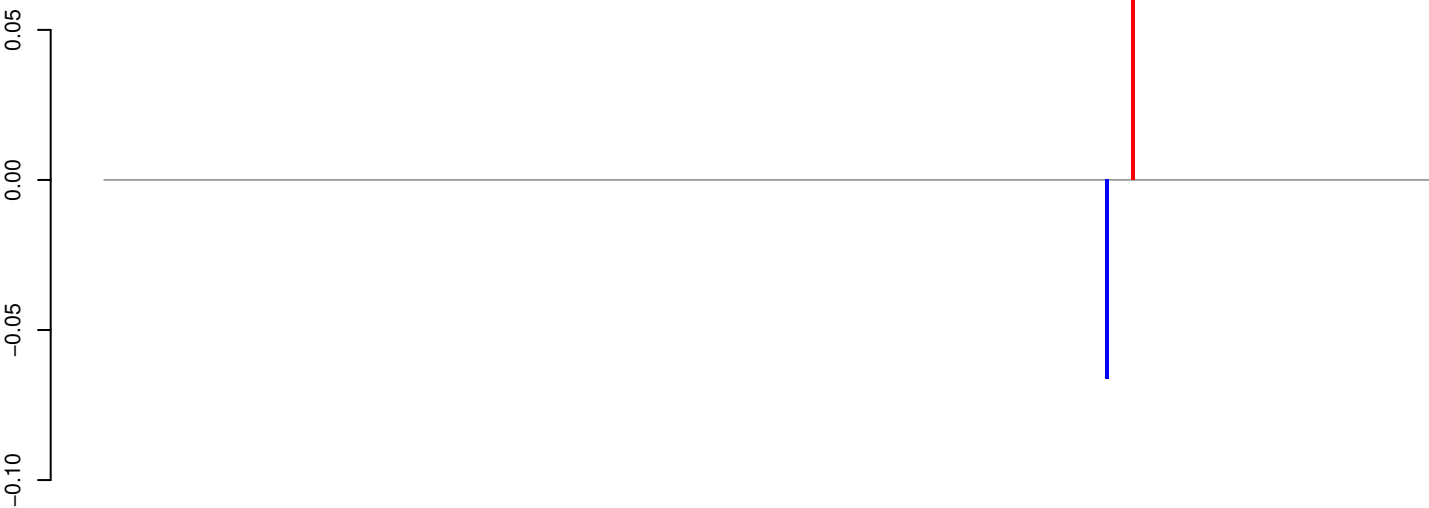


positive=0, negative=0, total=0

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

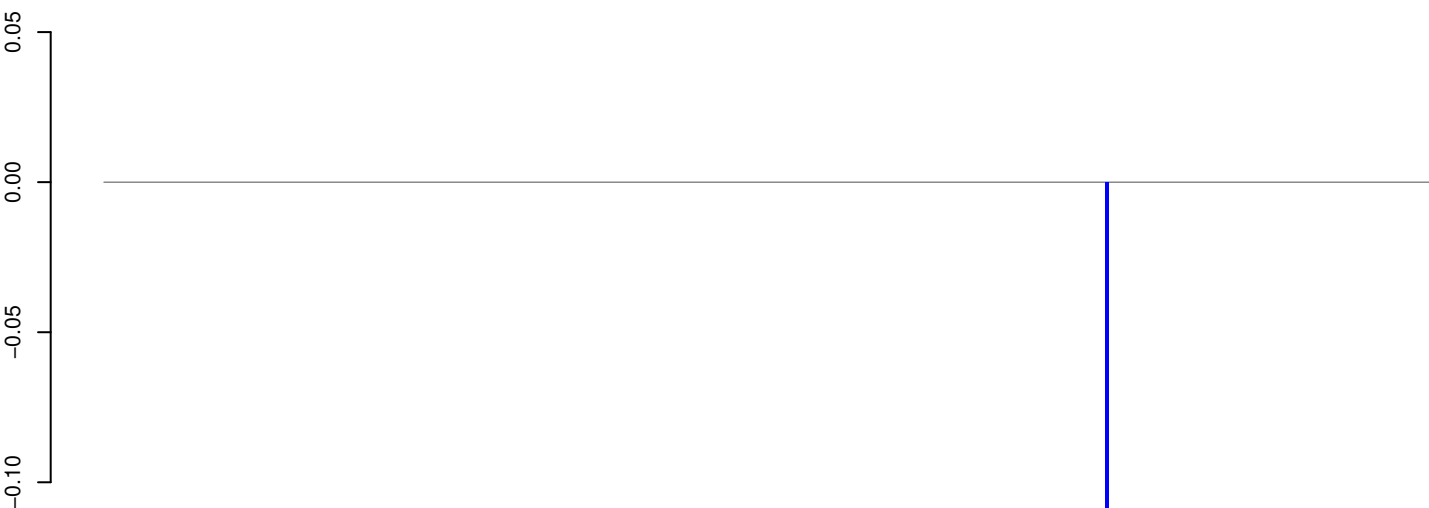


AeAeg_Aag2_BTV2_AK.18_23.rep



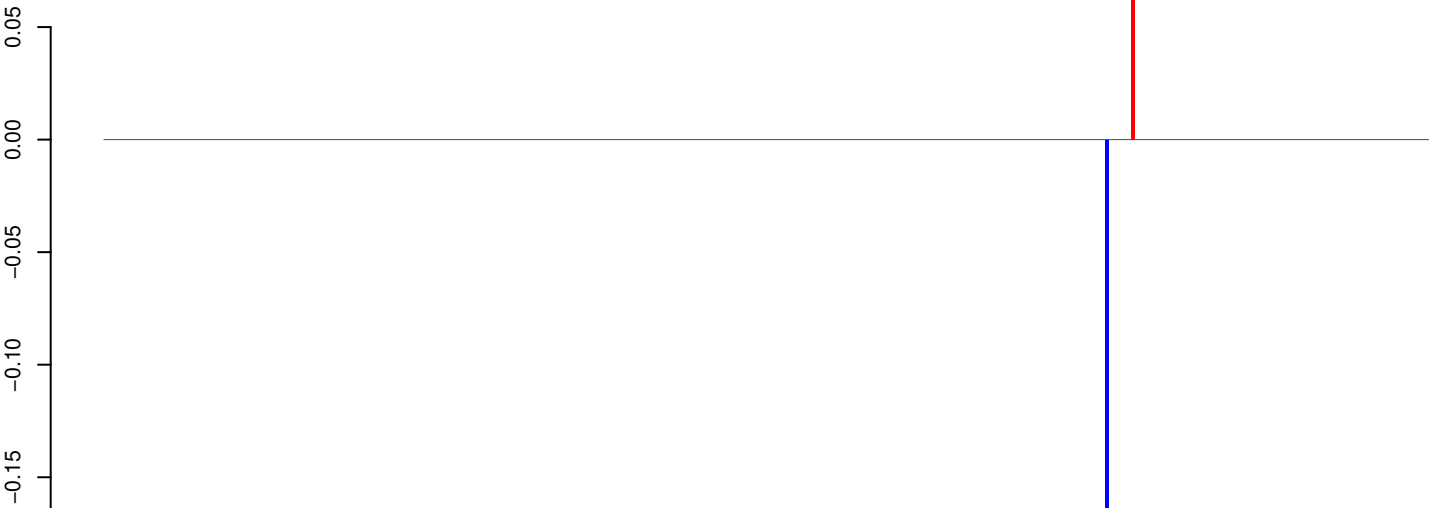
positive=0, negative=0, total=0

AeAeg_Aag2_BTV2_AK.24_35.rep



positive=0, negative=0, total=0

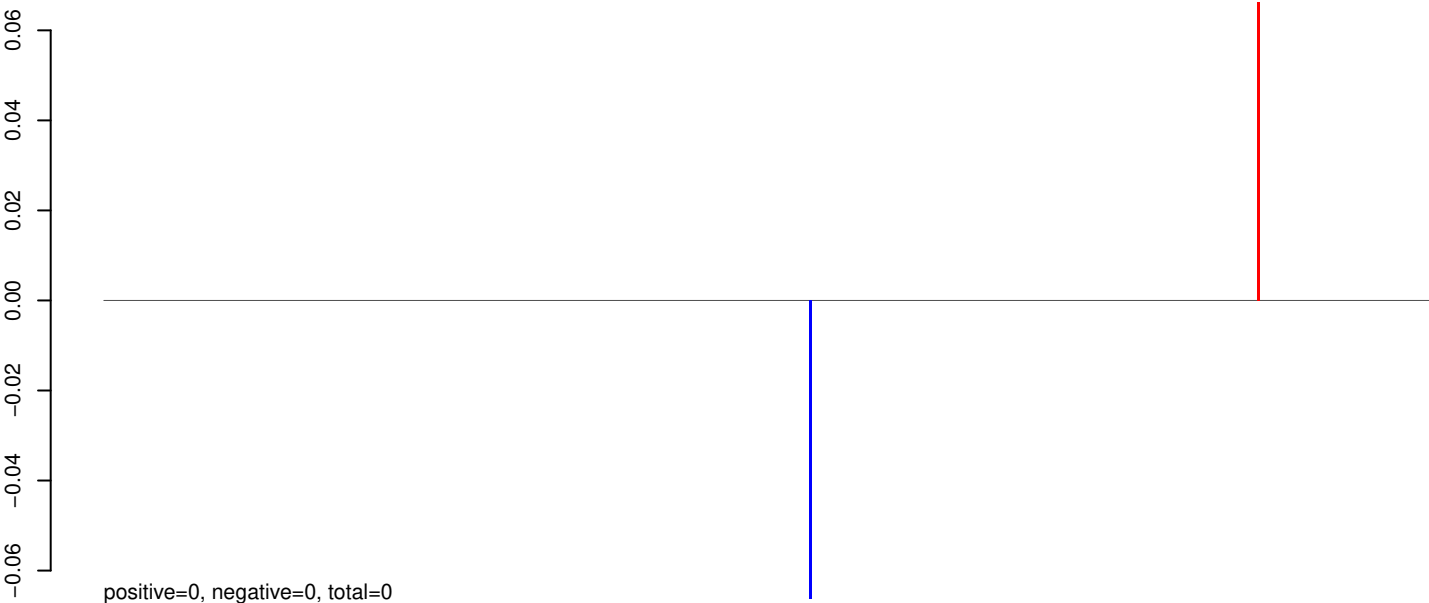
AeAeg_Aag2_BTV2_AK.rep



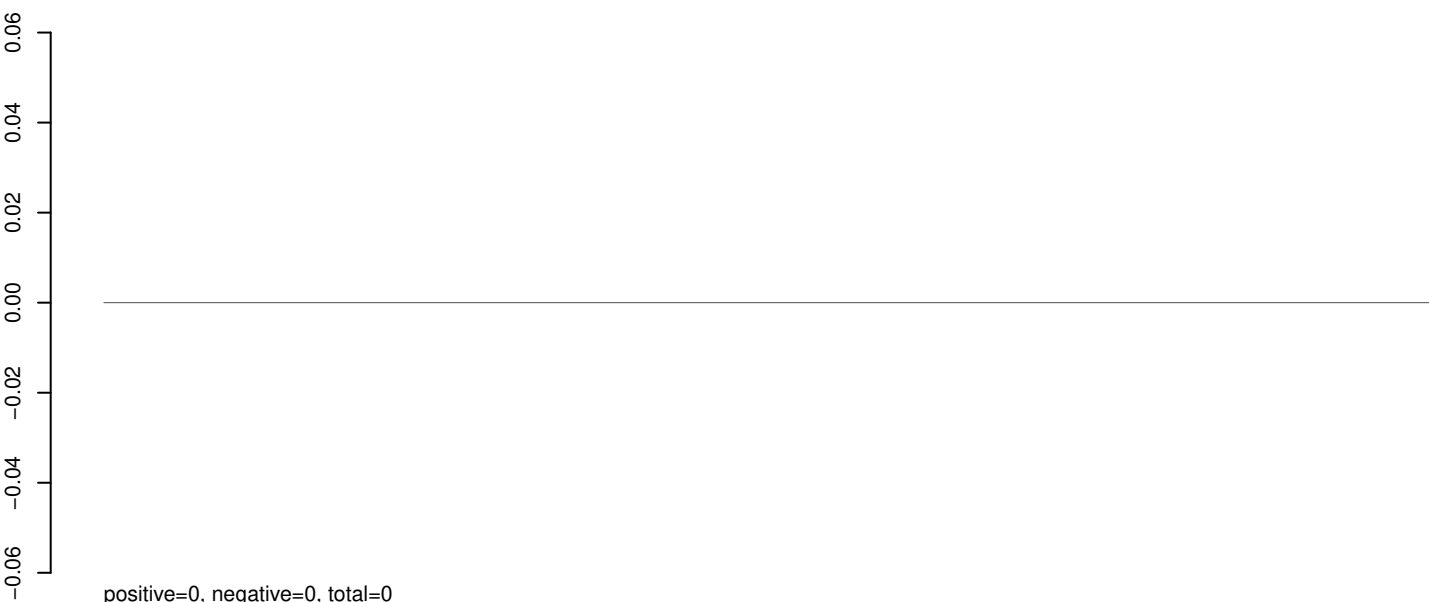
positive=0, negative=0, total=0
Window size=25, length=10254, virus@DQ235149.1-Royal_Farm_virus_from_Afghanistan_polyprotein_gene:1-10254

0 2000 4000 6000 8000 10000

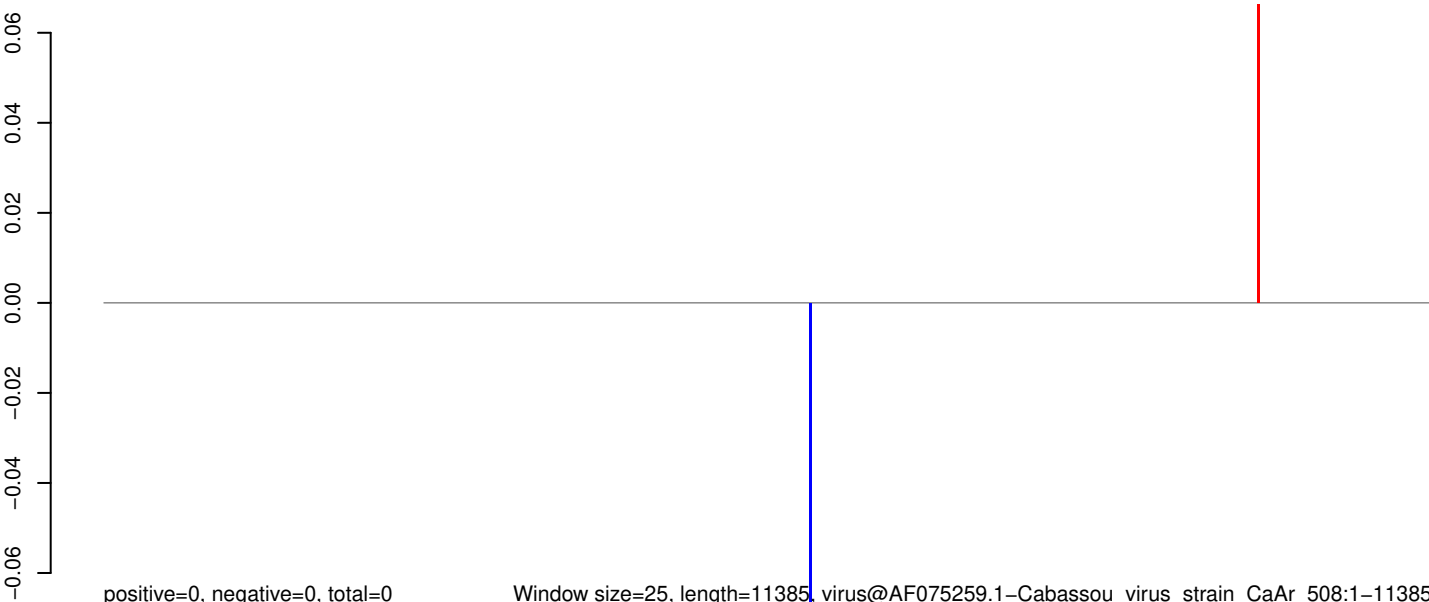
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

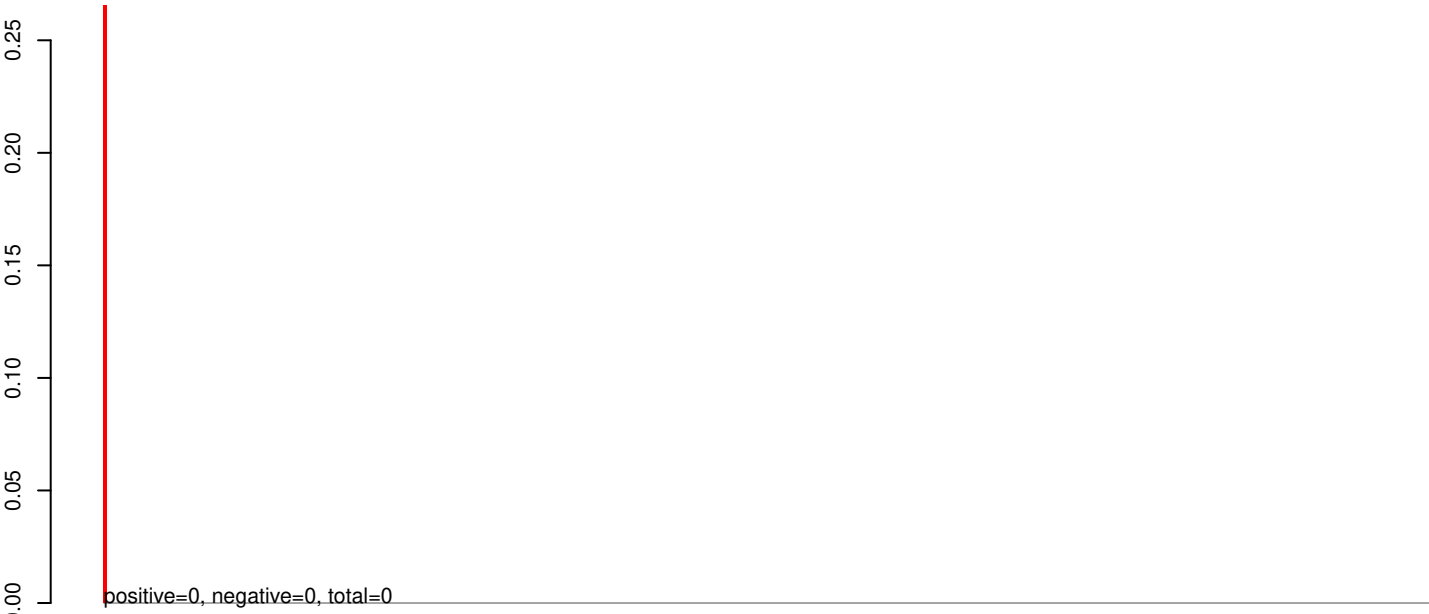


AeAeg_Aag2_BTV2_AK.rep

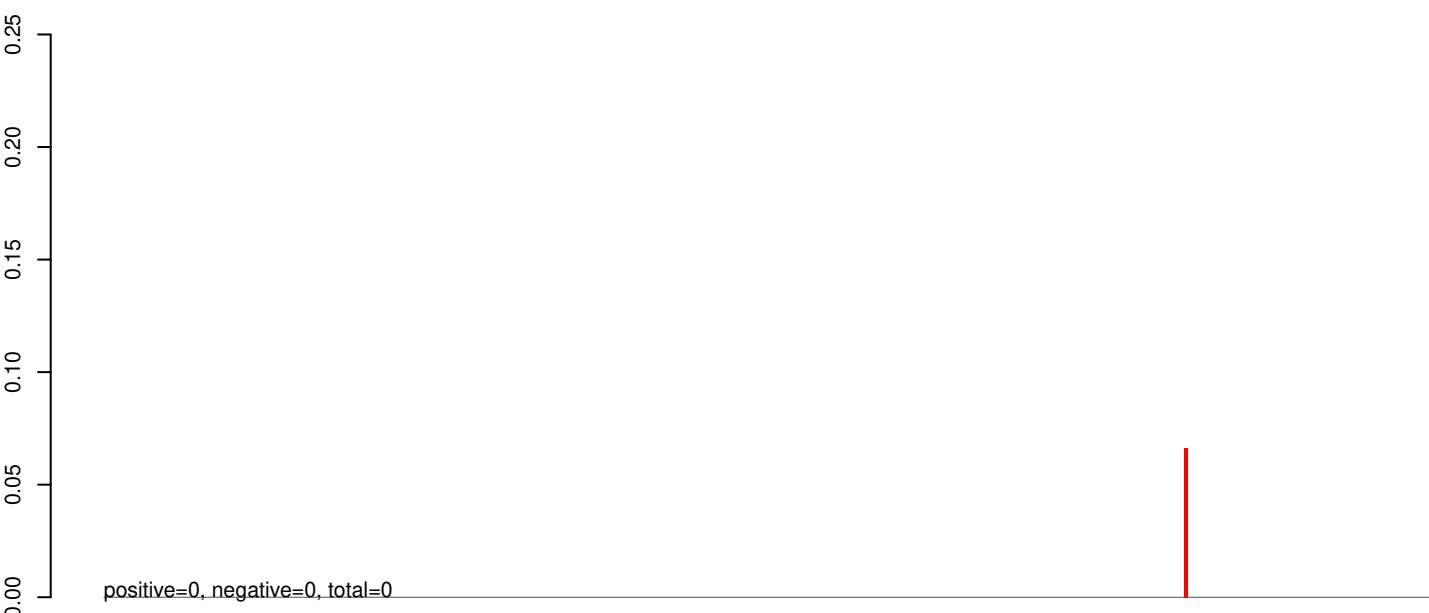


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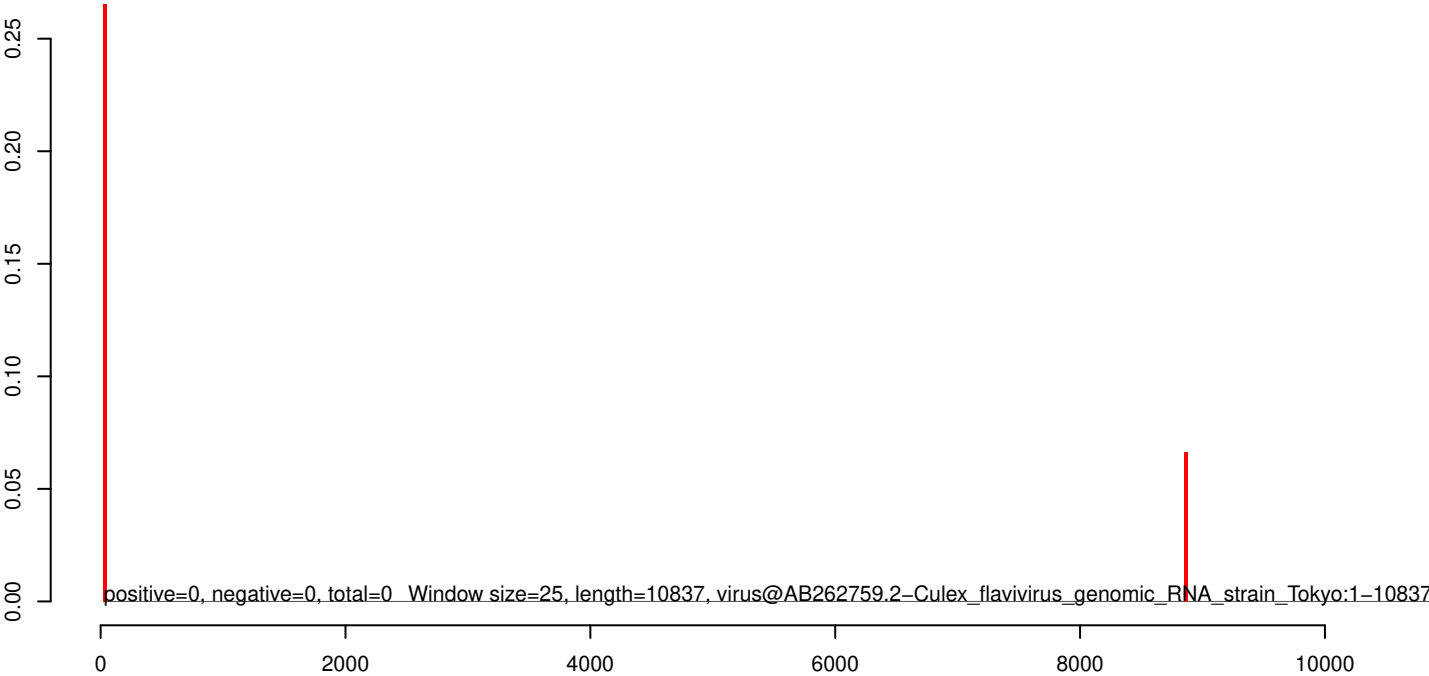
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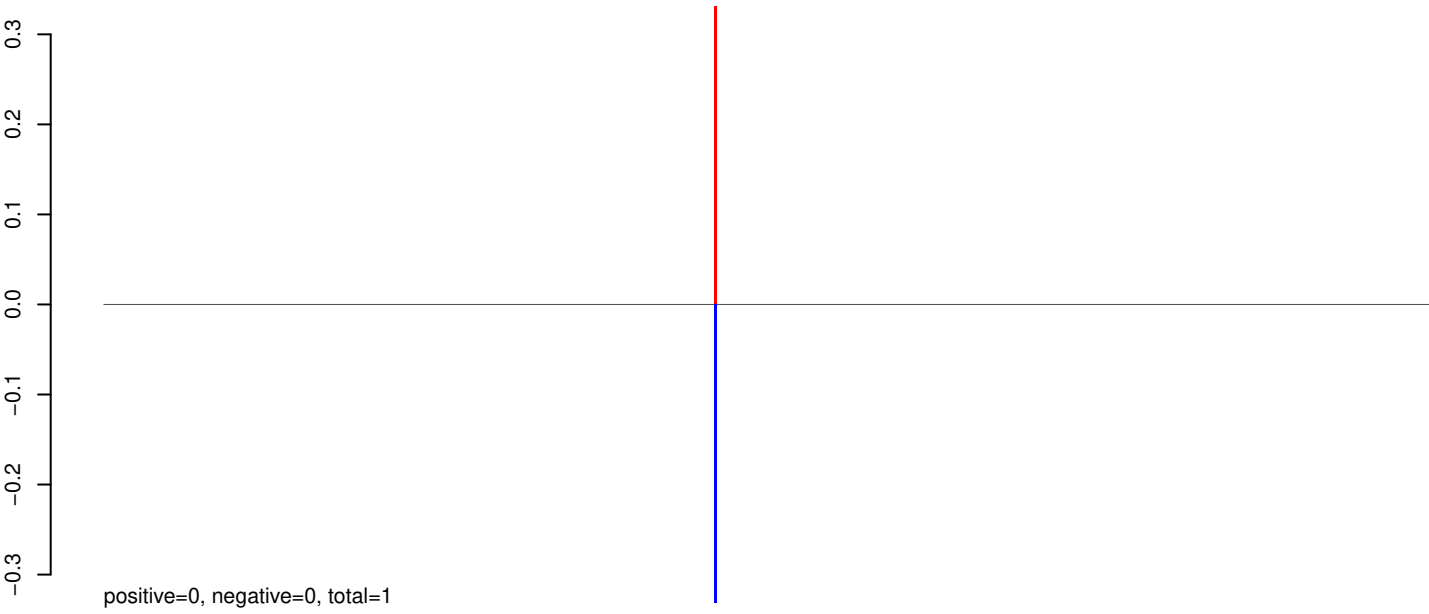
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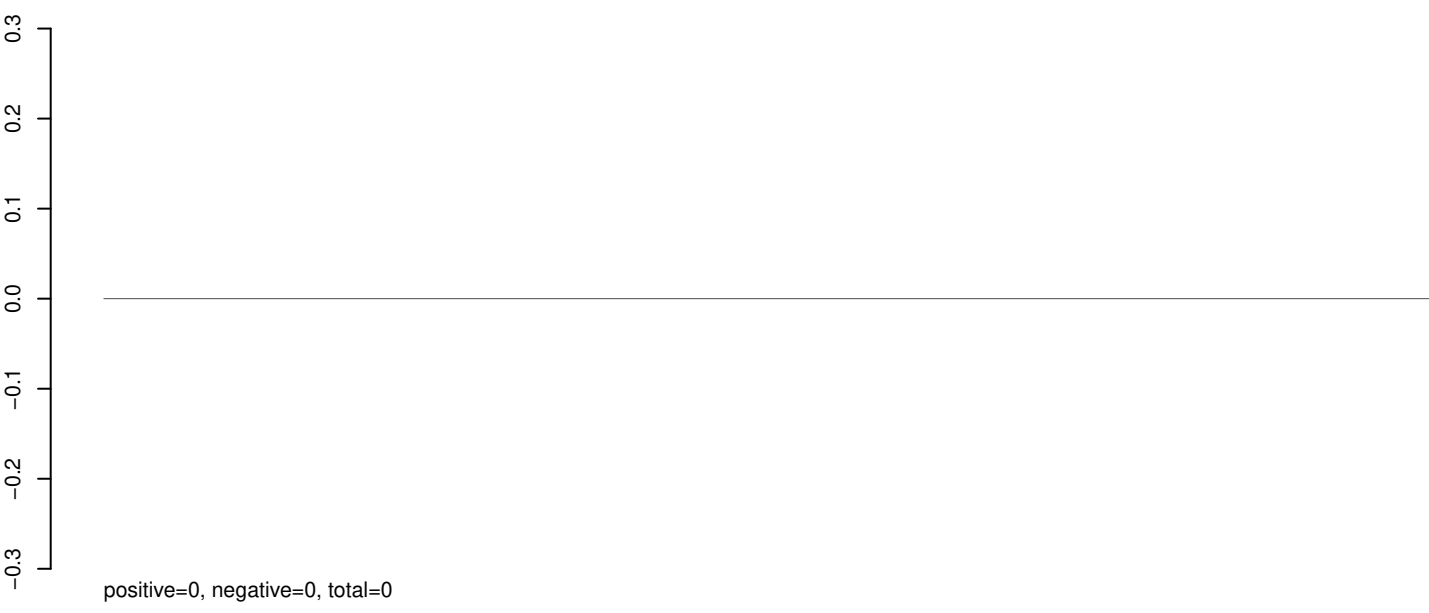
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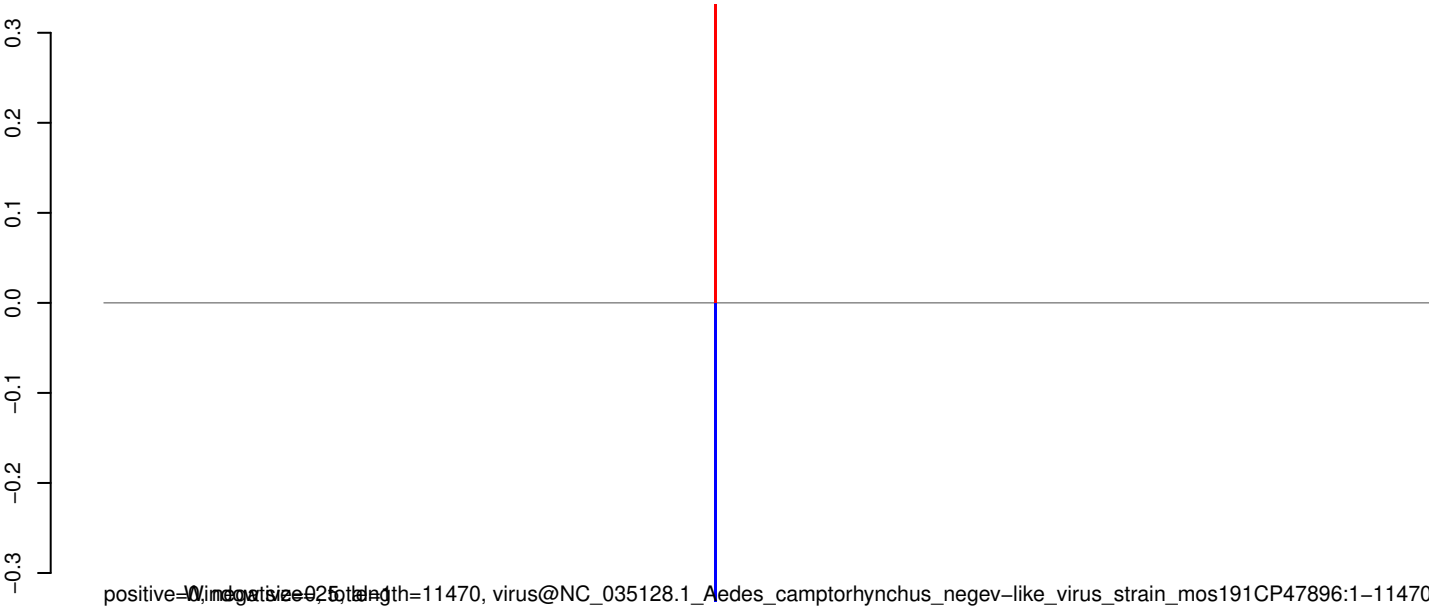
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AeAeg_Aag2_BTV2_AK.24_35.rep



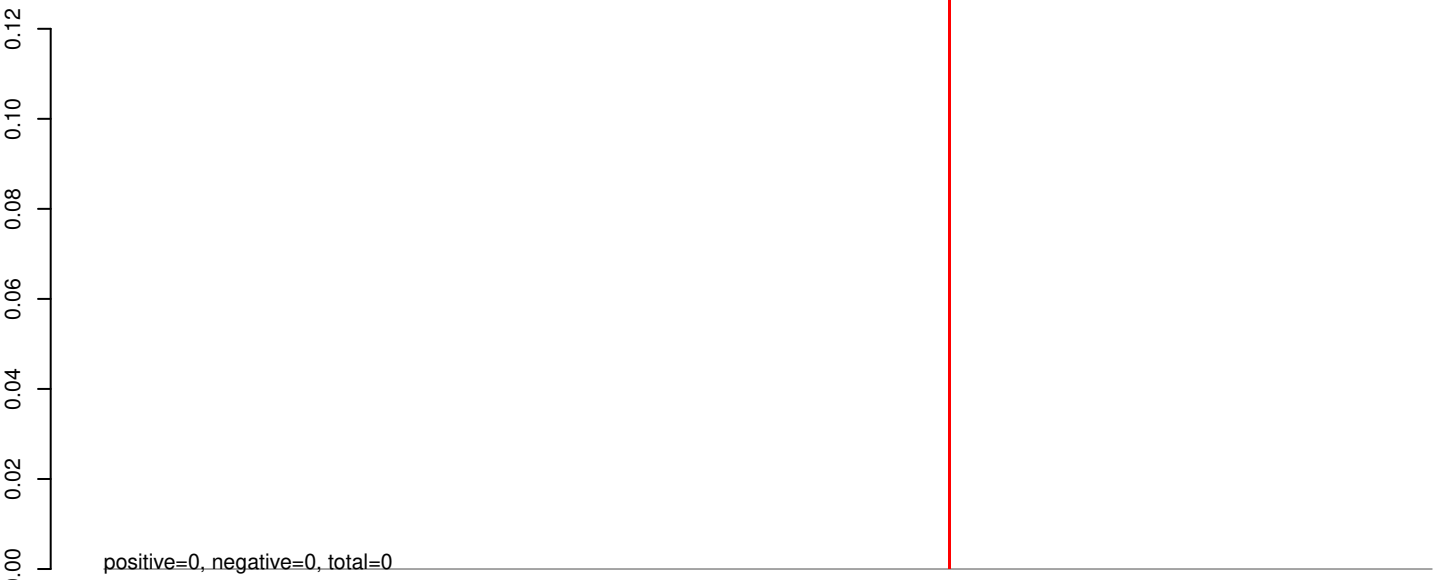
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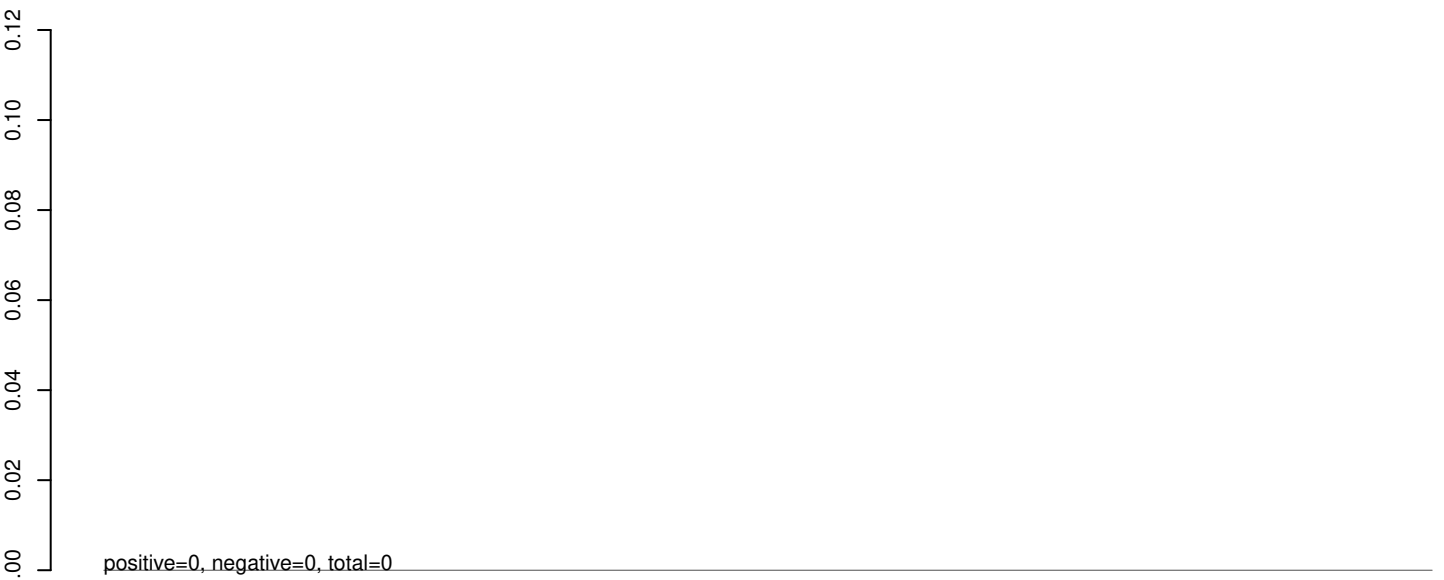
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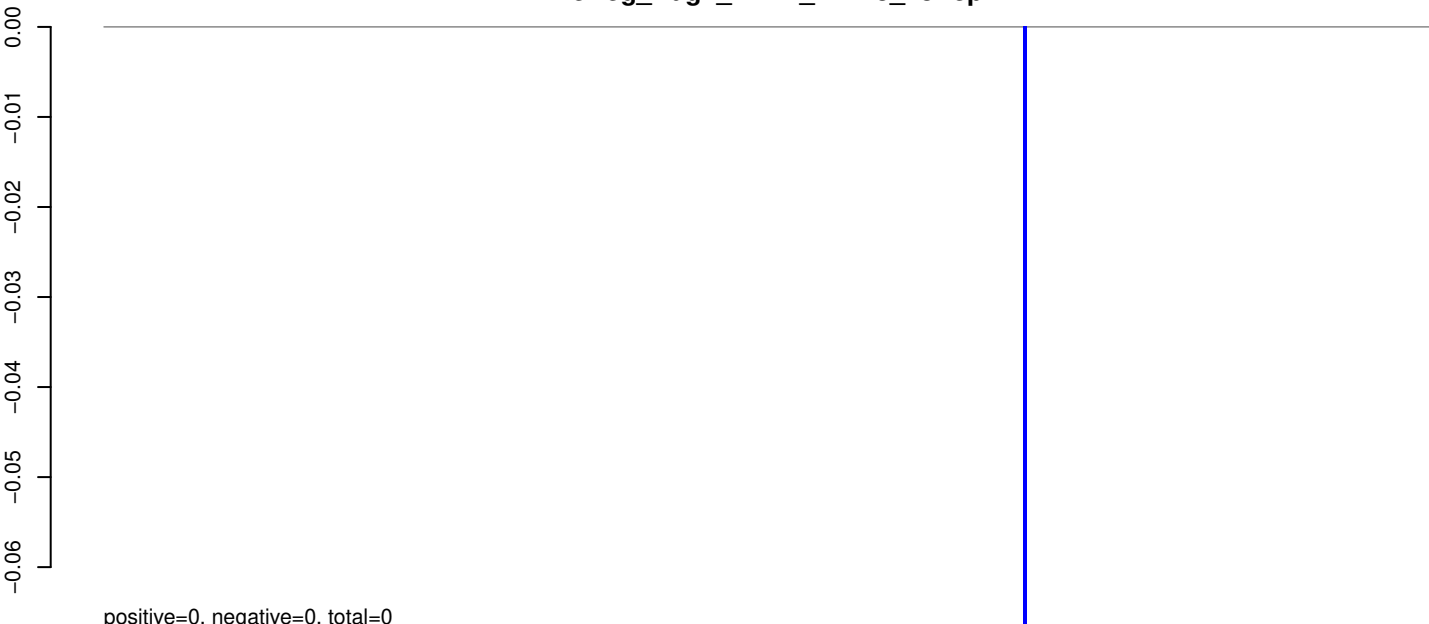
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AeAeg_Aag2_BTV2_AK.rep



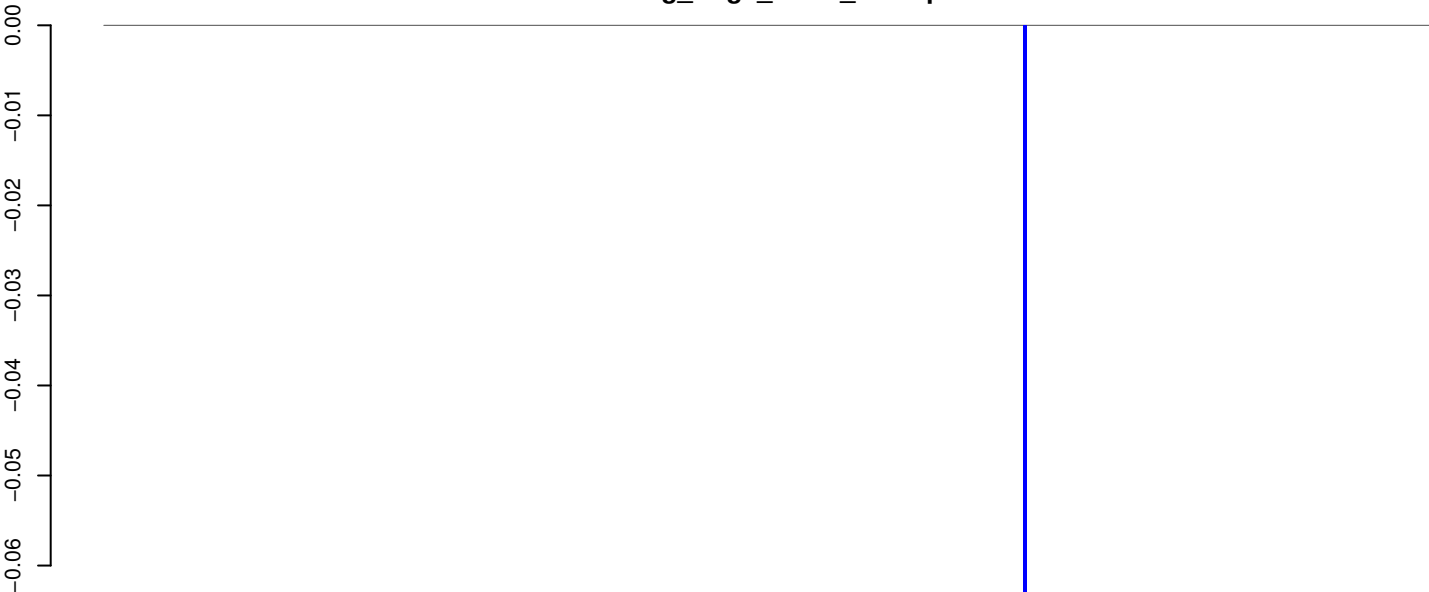
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AeAeg_Aag2_BTV2_AK.24_35.rep



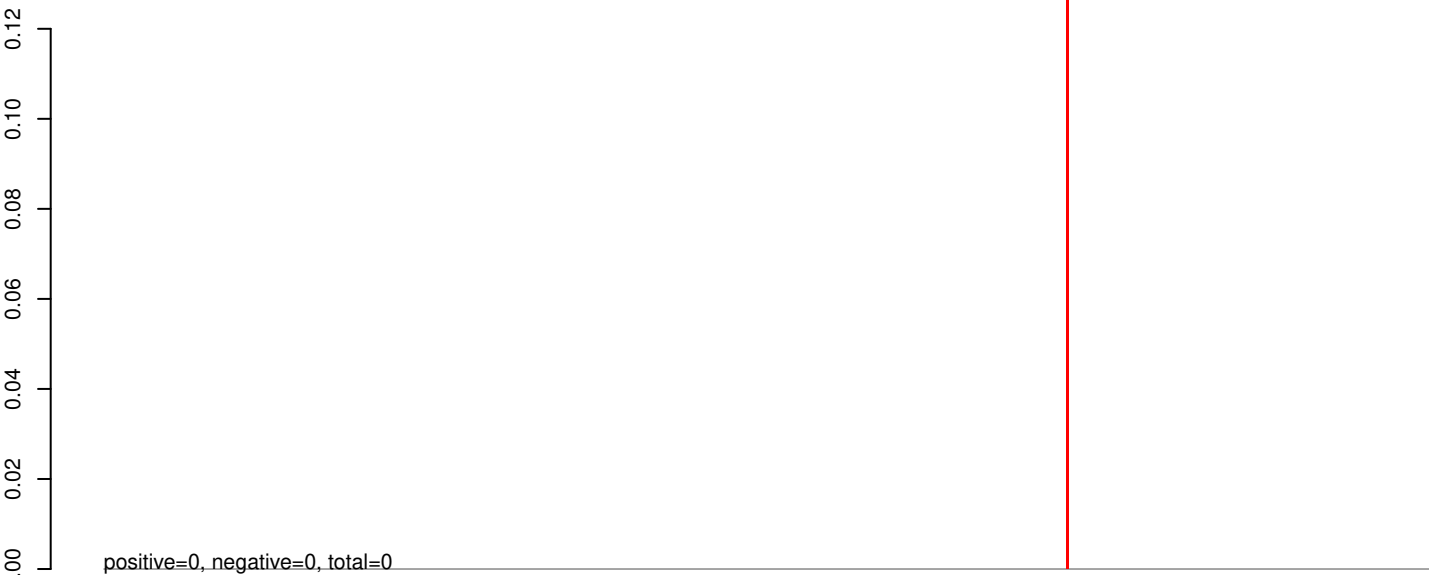
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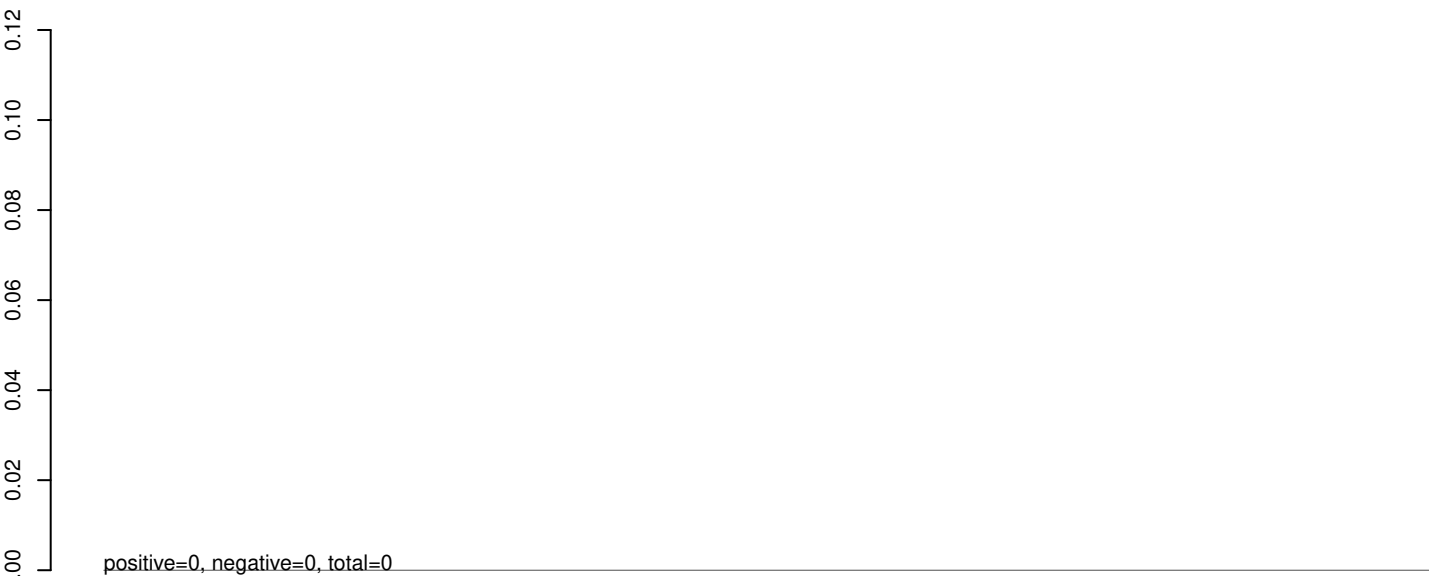
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0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



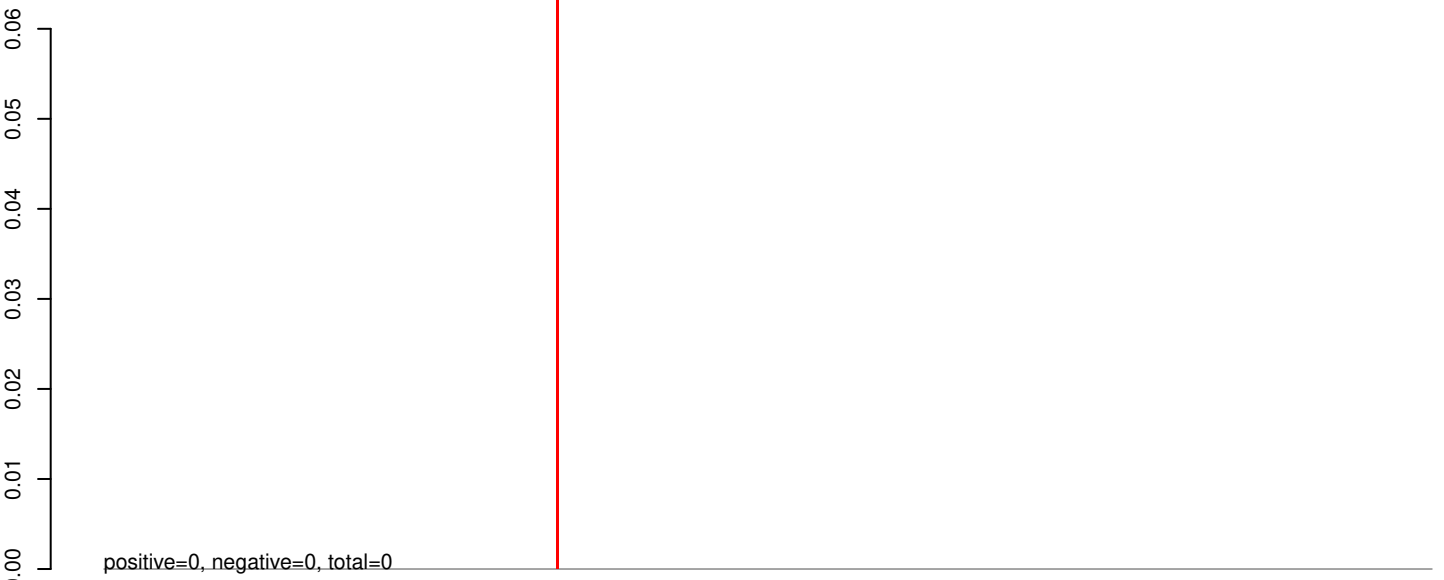
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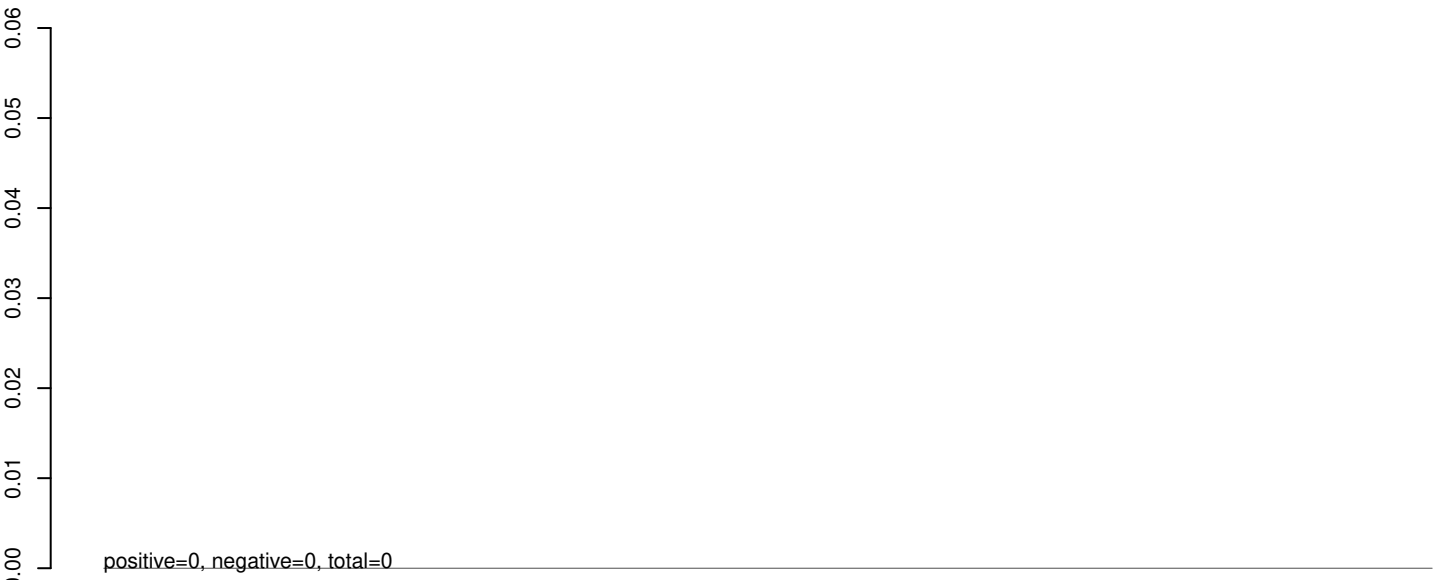
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AeAeg_Aag2_BTV2_AK.18_23.rep



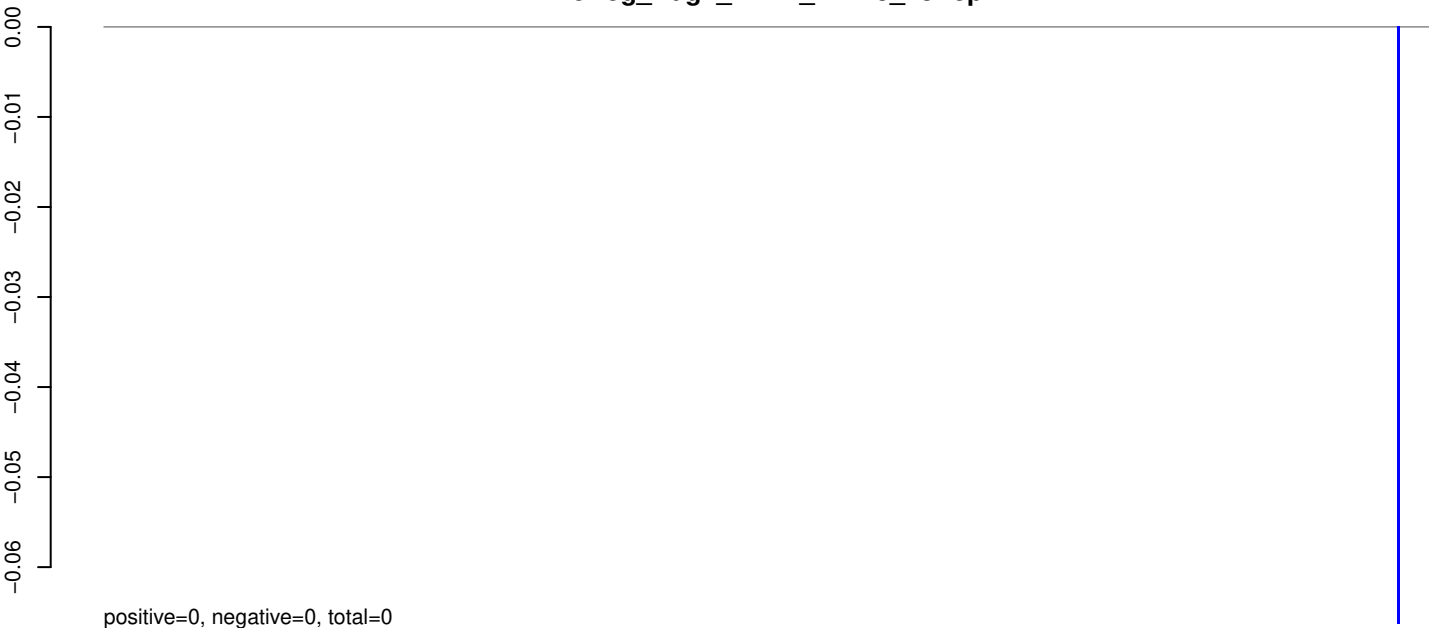
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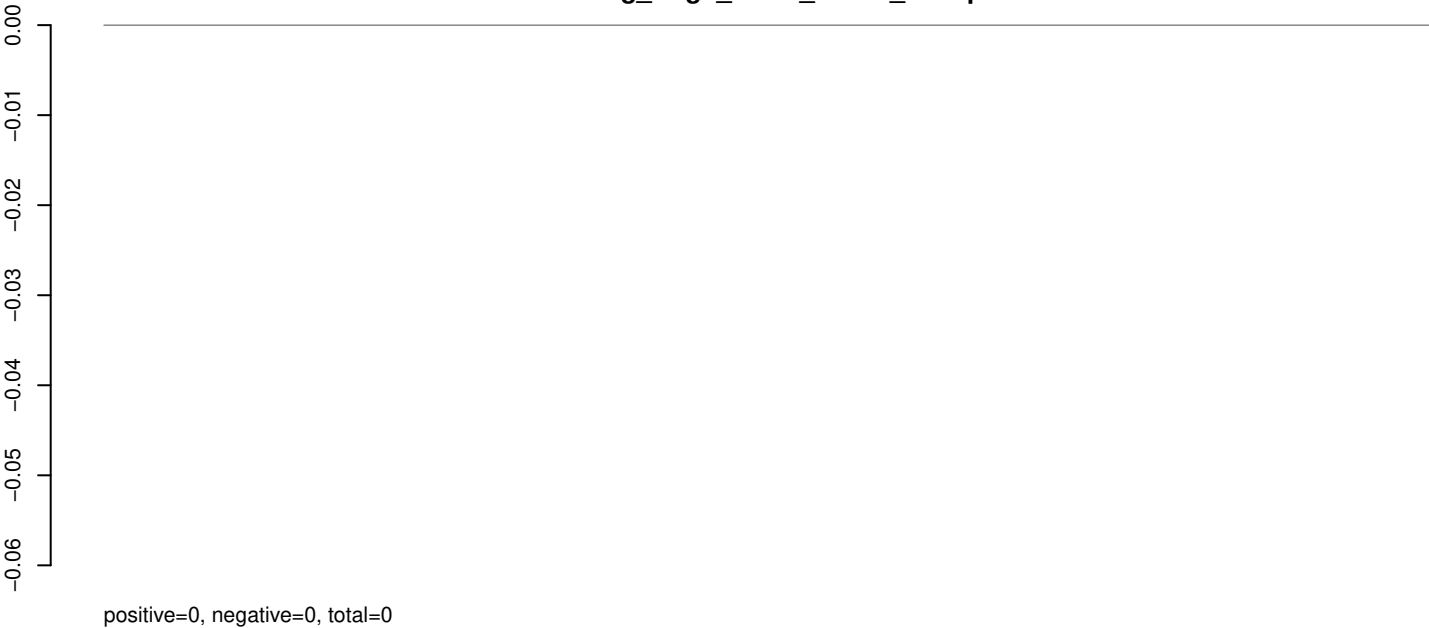
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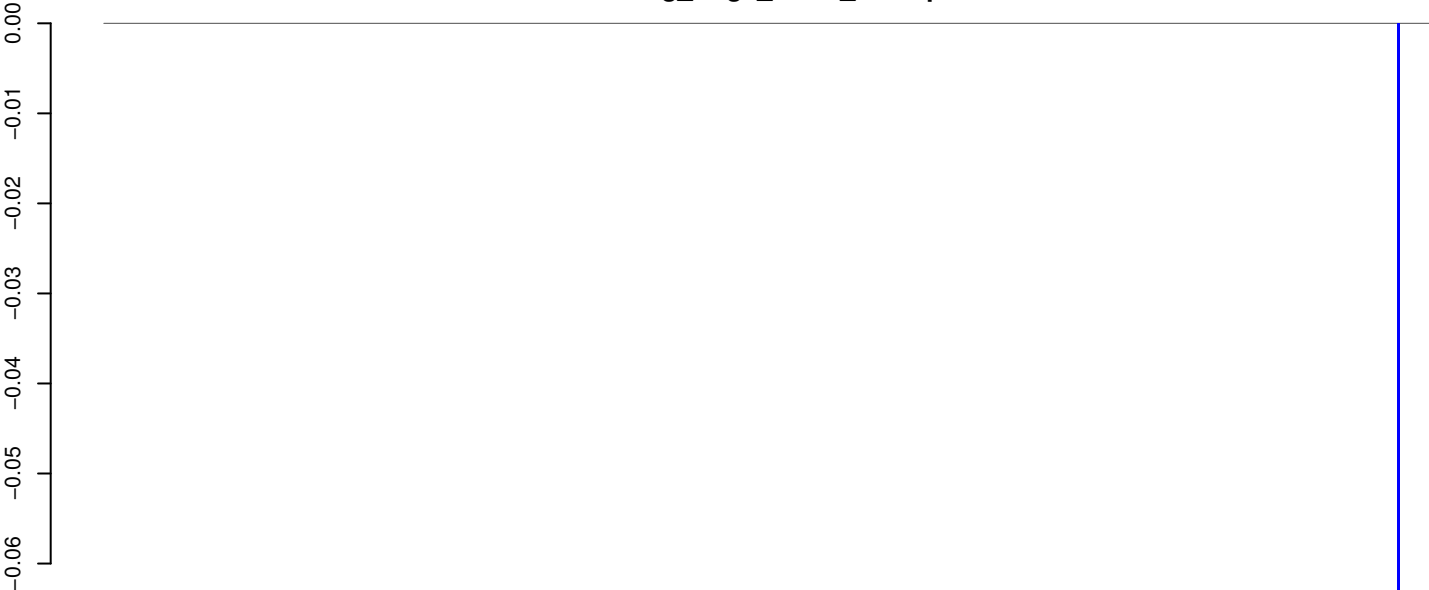
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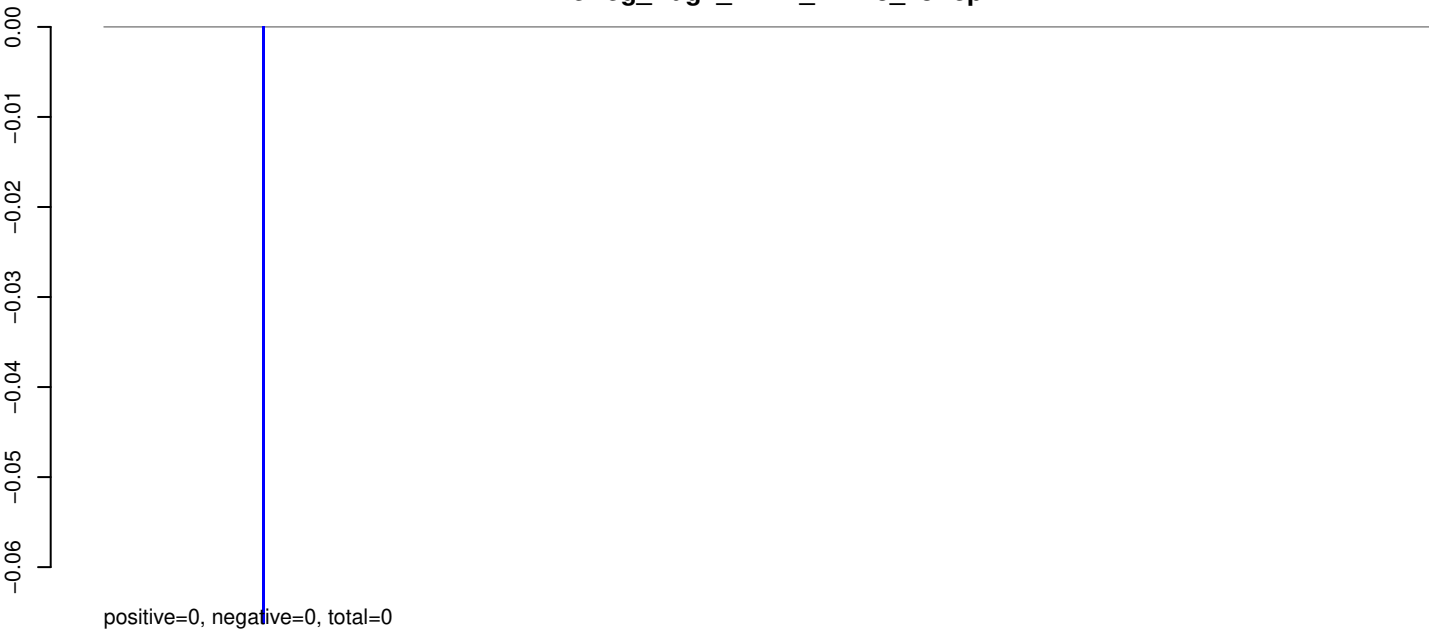
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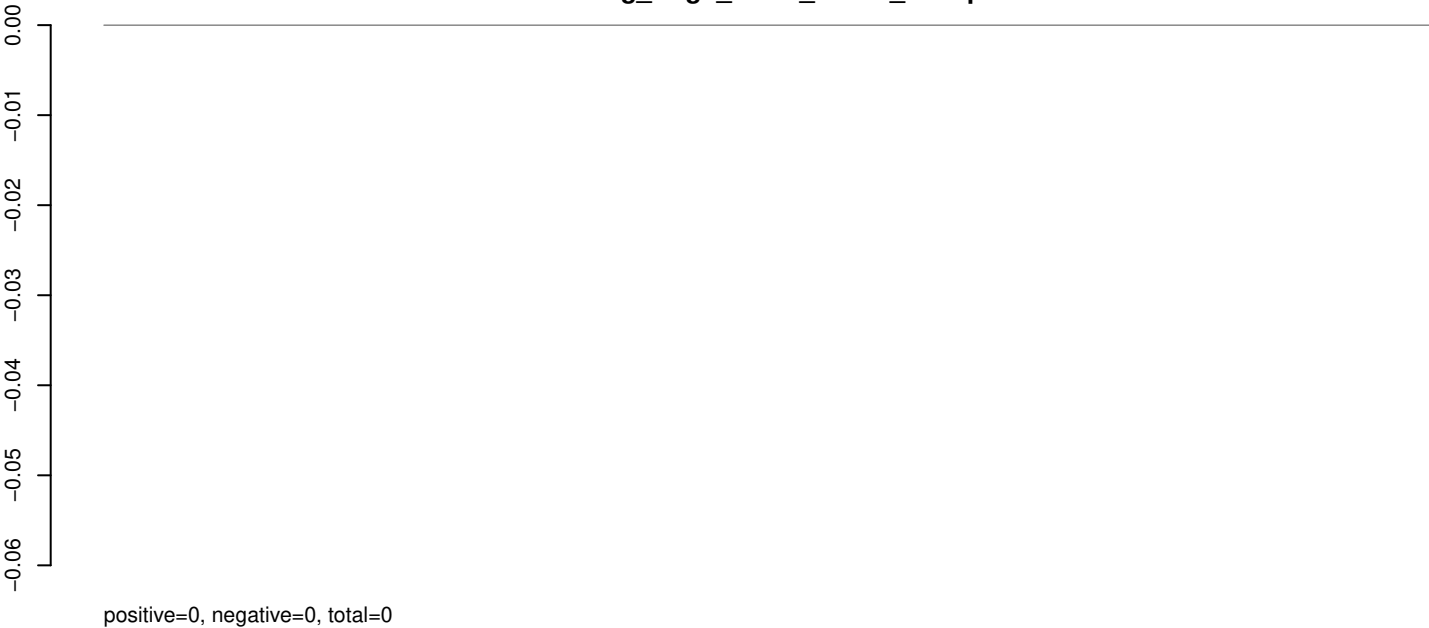
AeAeg_Aag2_BTV2_AK.rep



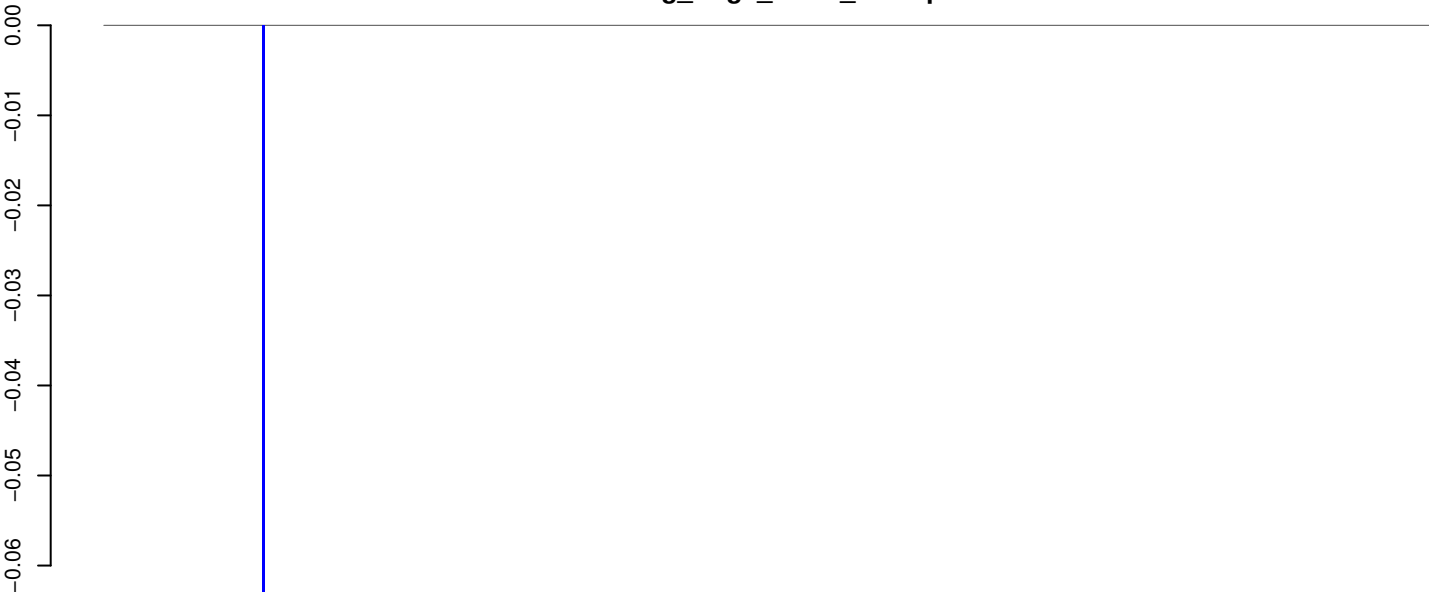
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



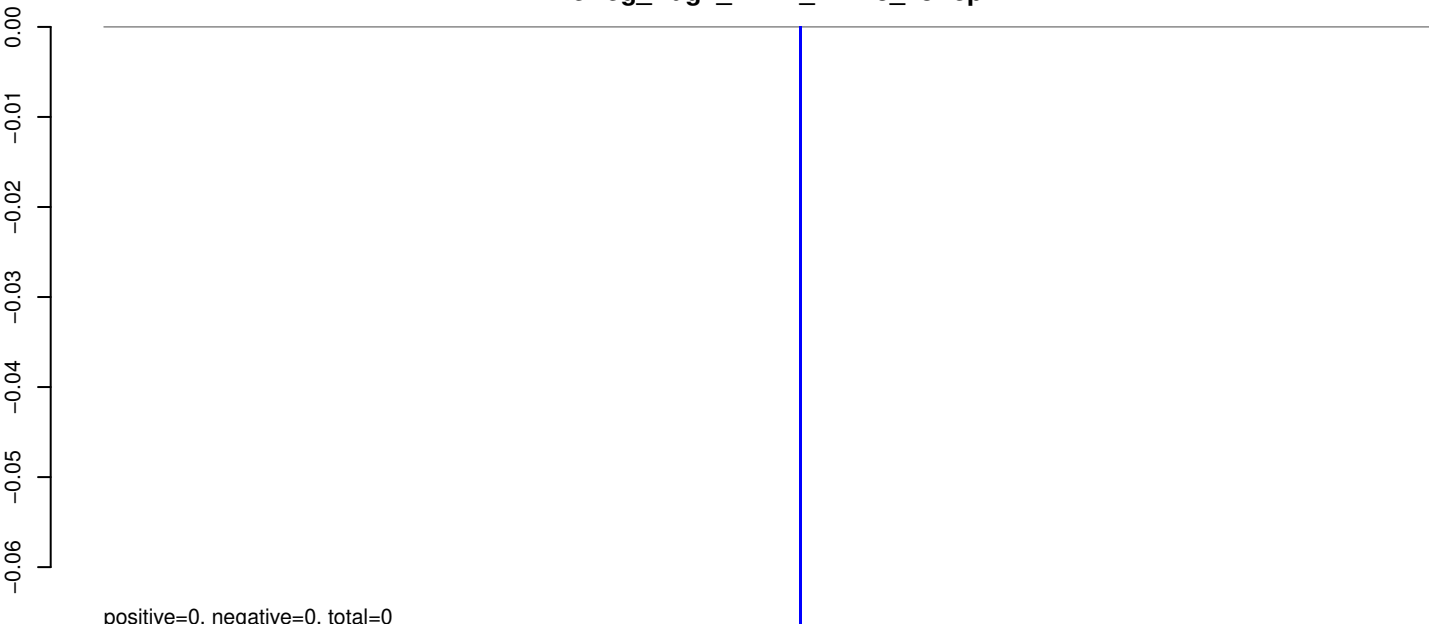
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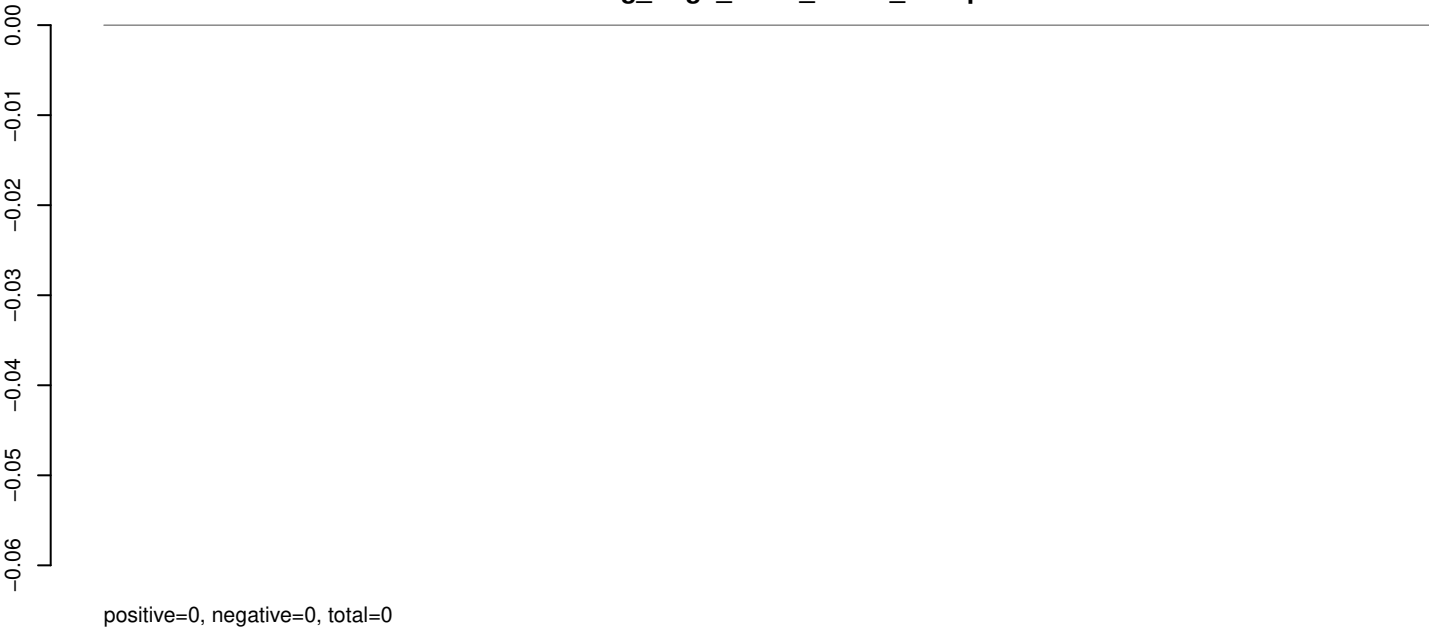
25, length=128, negais @ NC_013445.1-Tai_virus_strain_F47_CI_2004_segment_S_plus_segment_M_plus_segment_L_nucleocapsid_:1-11728

0 2000 4000 6000 8000 10000 12000

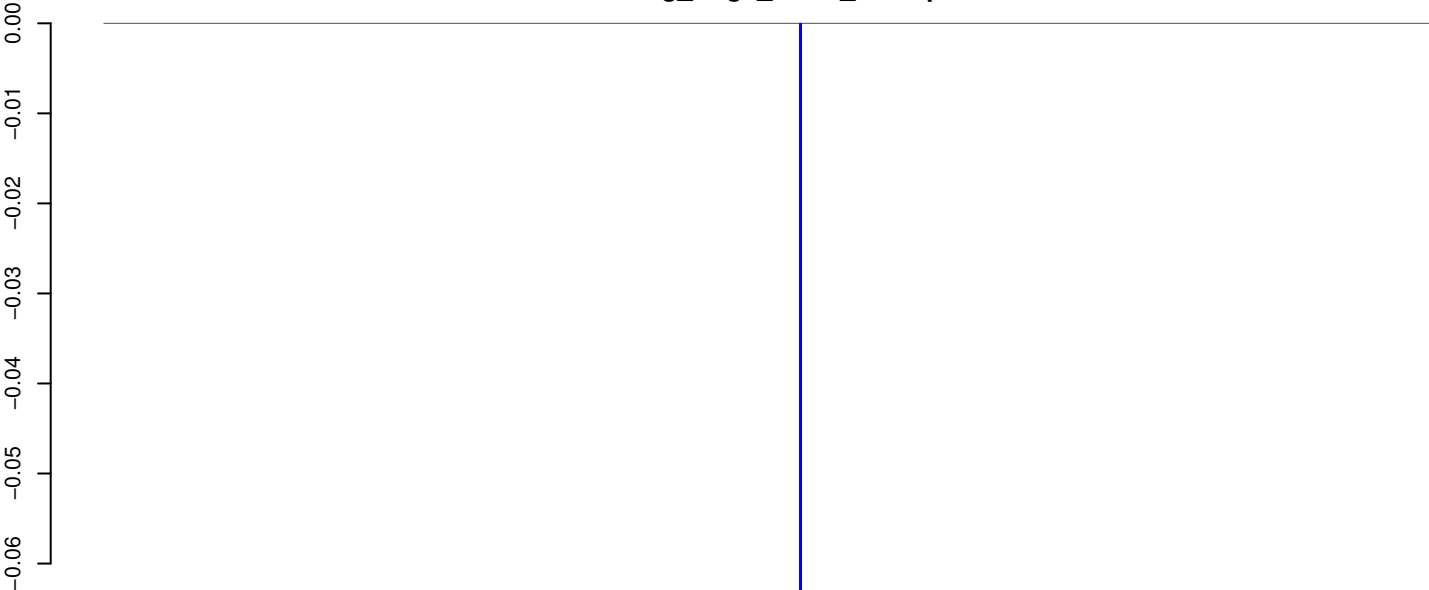
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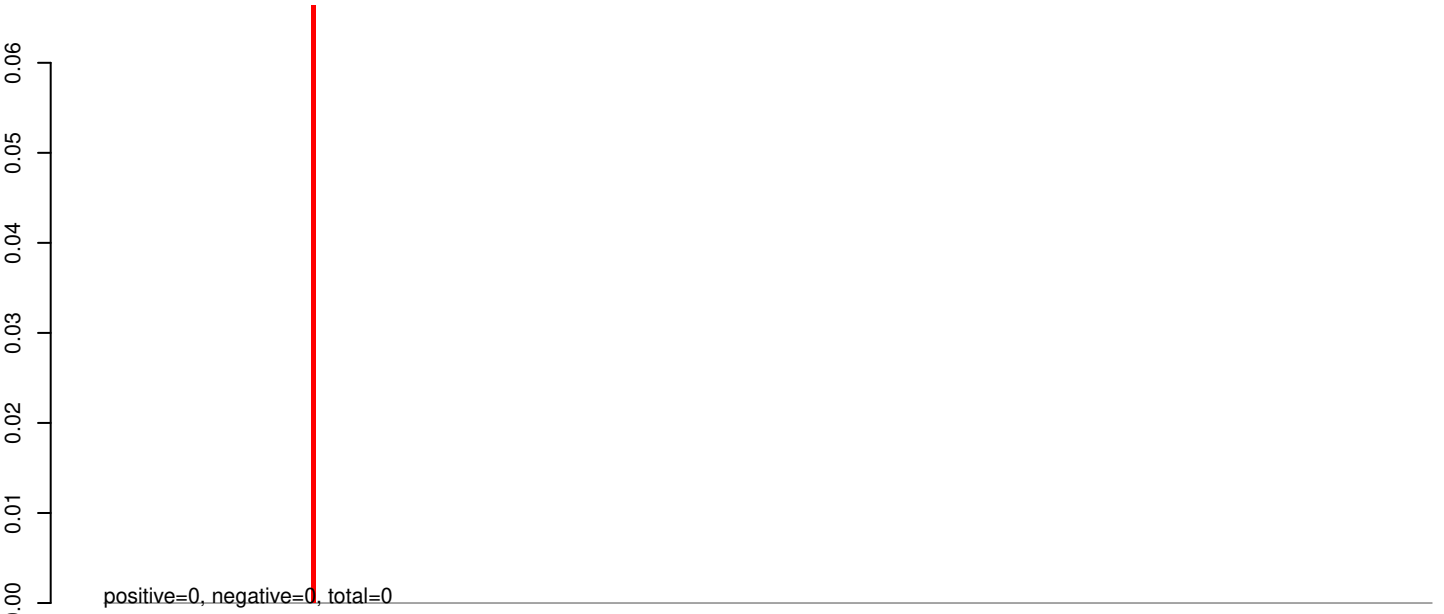
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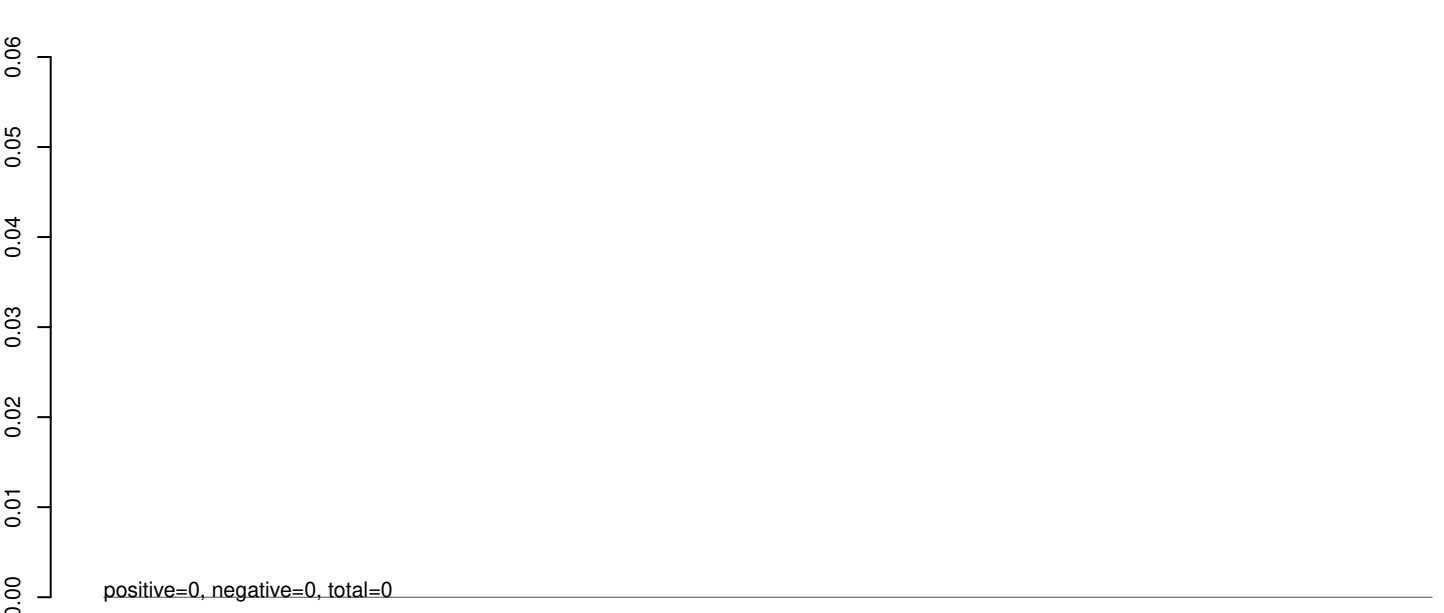
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AeAeg_Aag2_BTV2_AK.18_23.rep



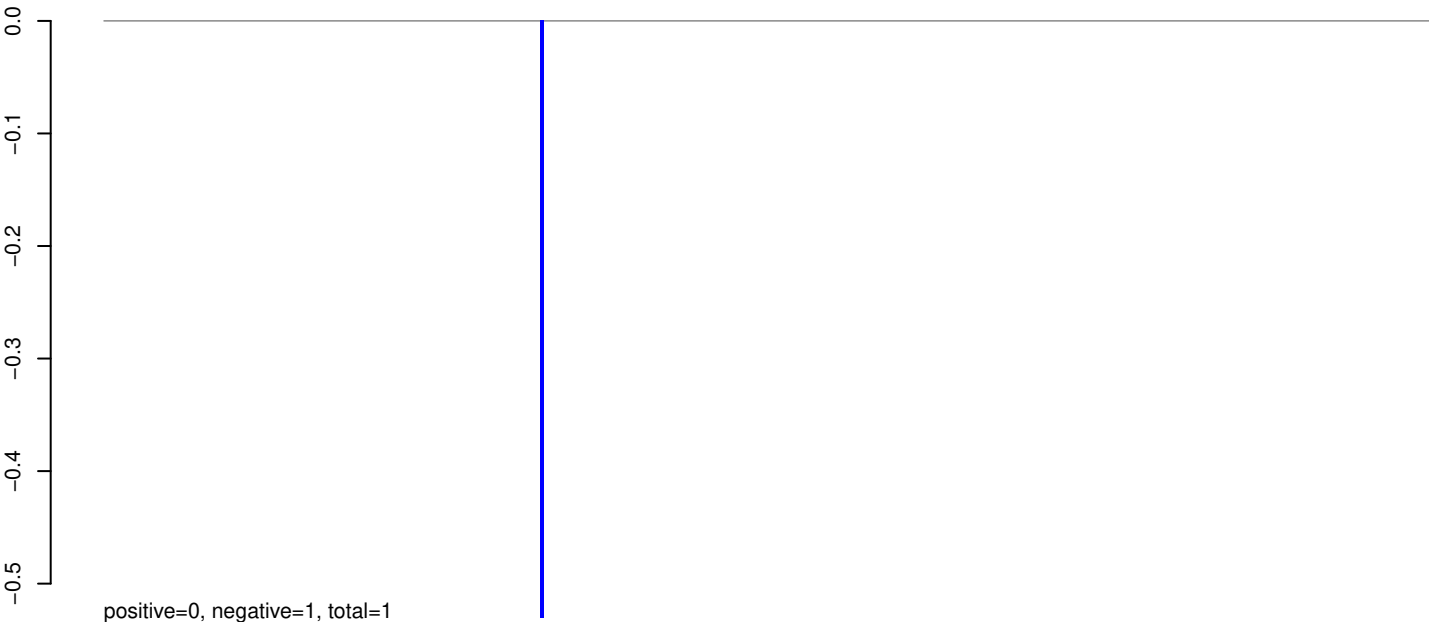
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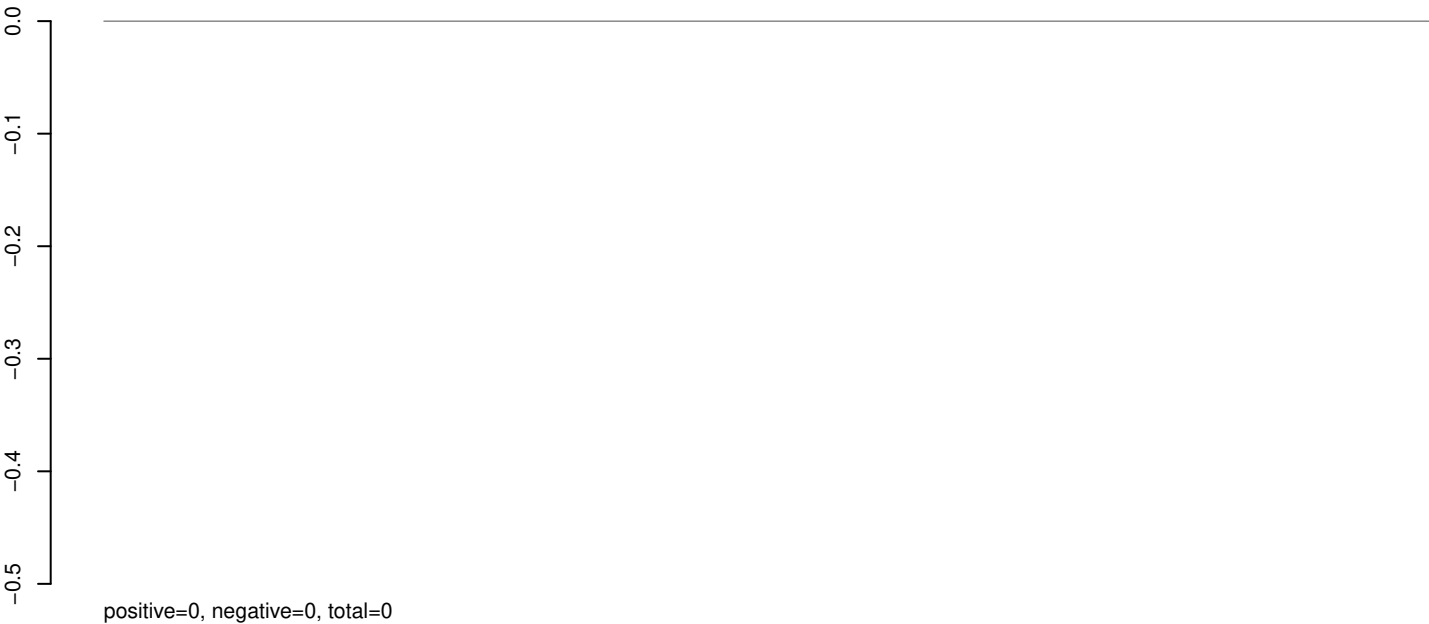
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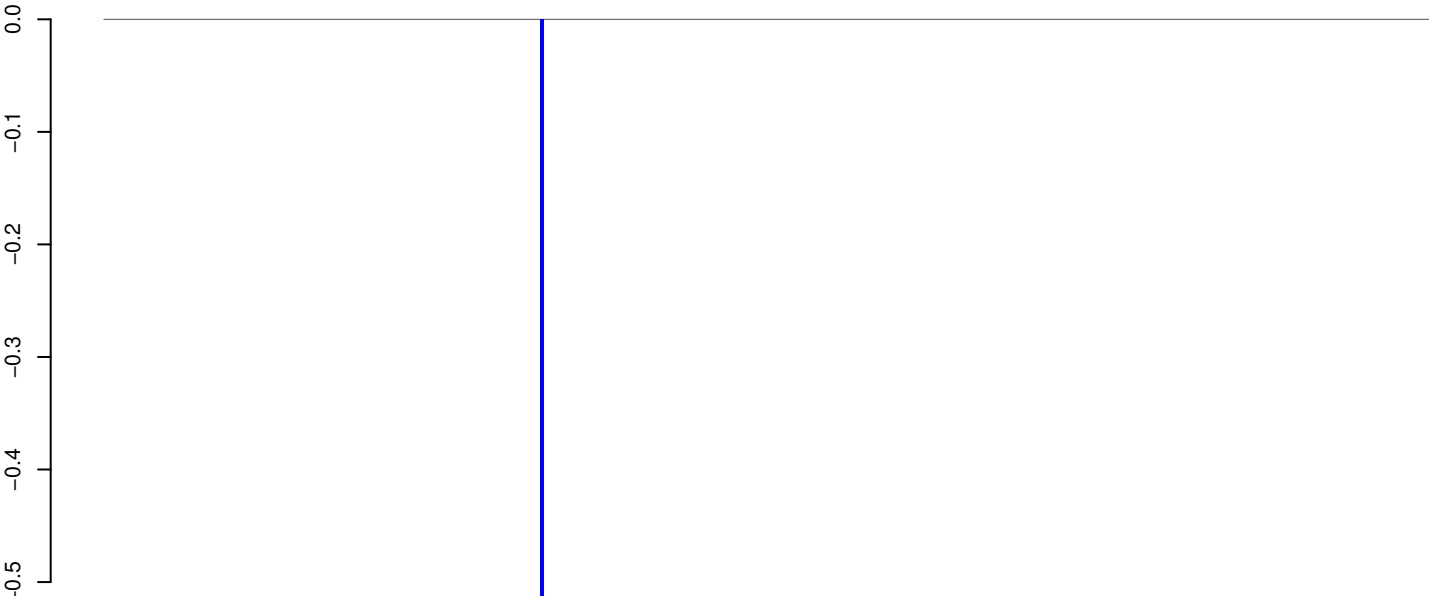
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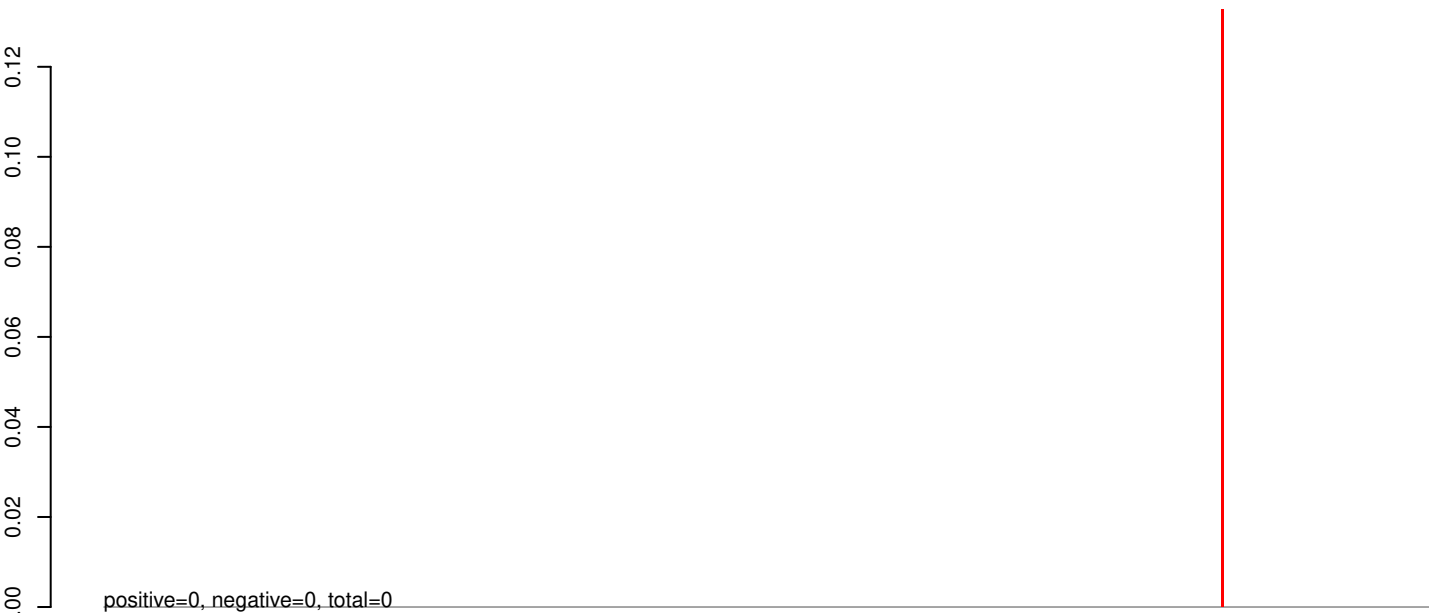
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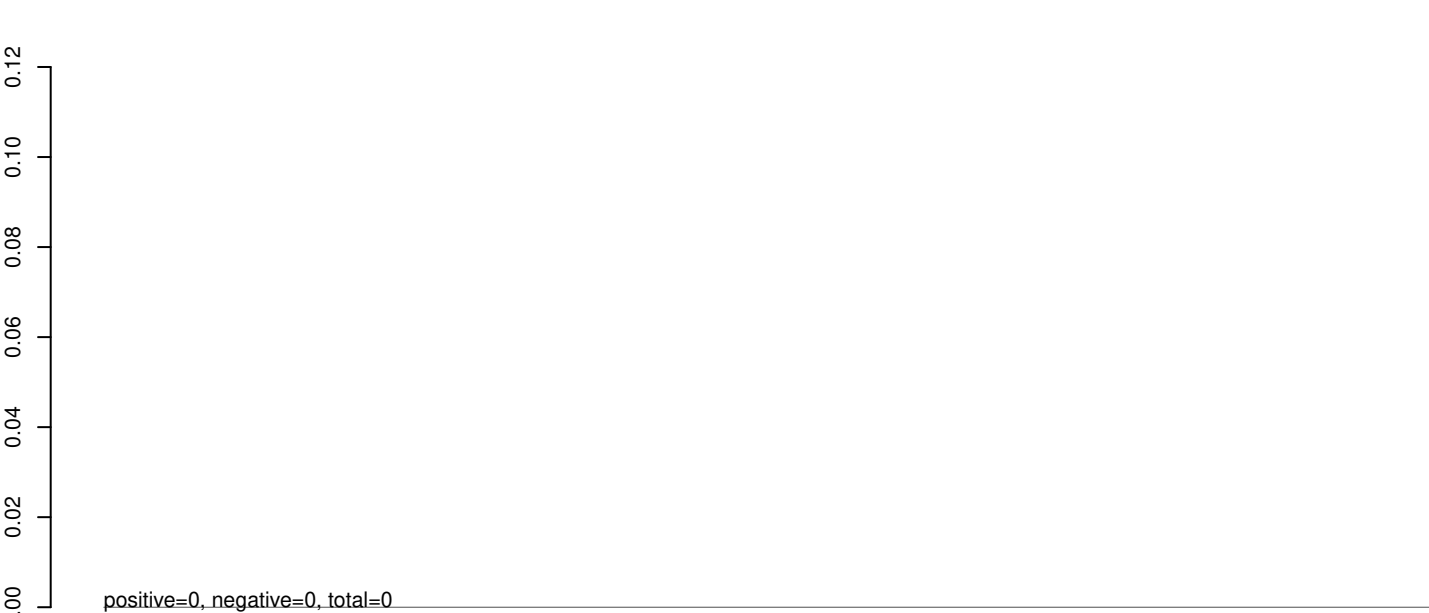
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0 2000 4000 6000 8000 10000

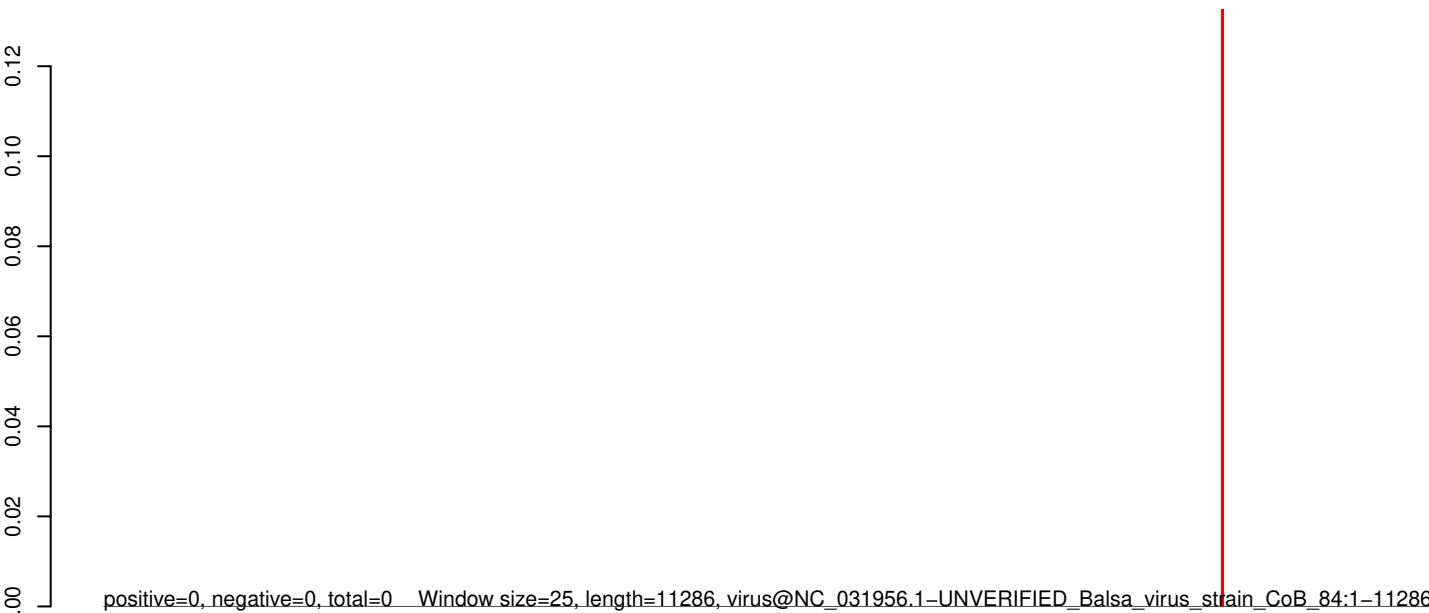
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AeAeg_Aag2_BTV2_AK.24_35.rep



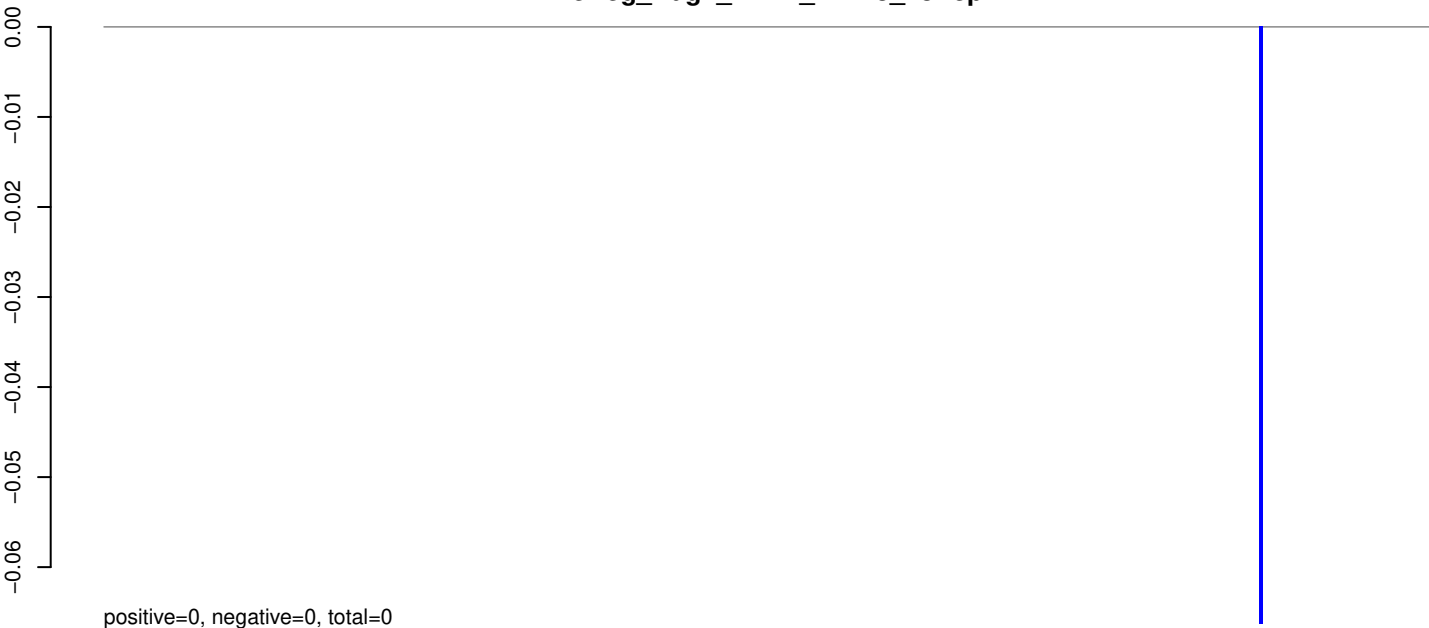
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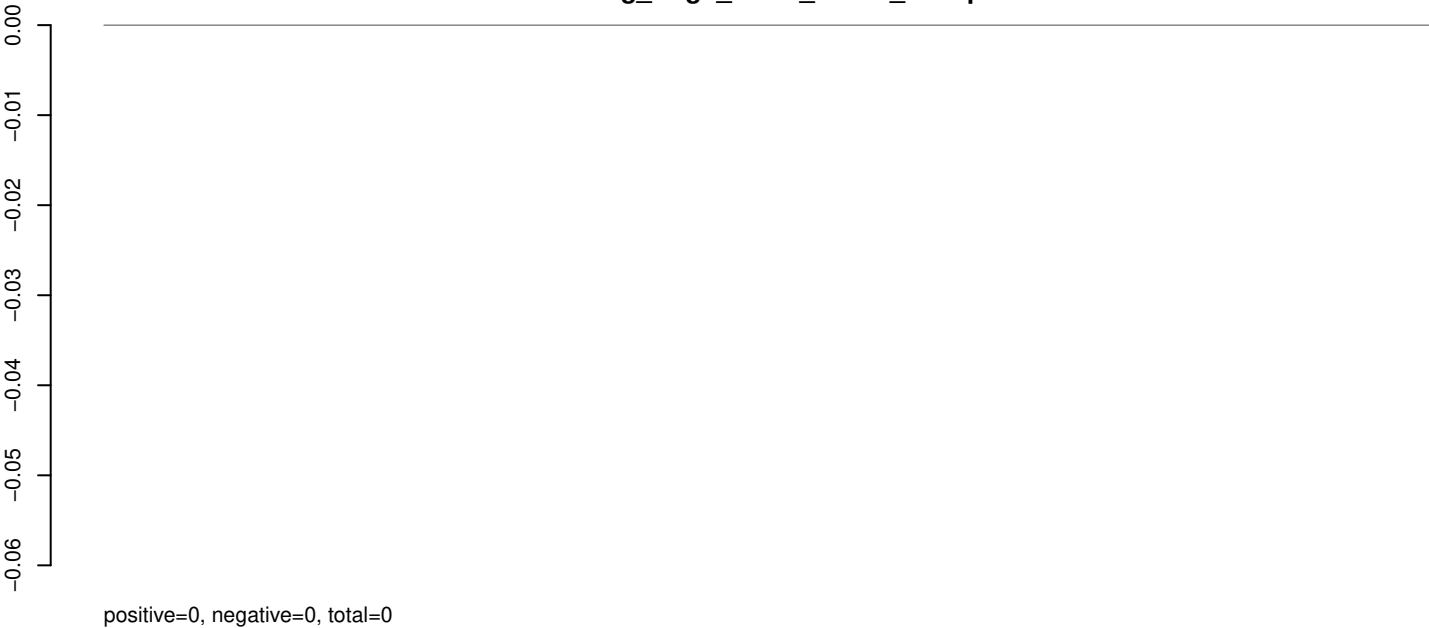
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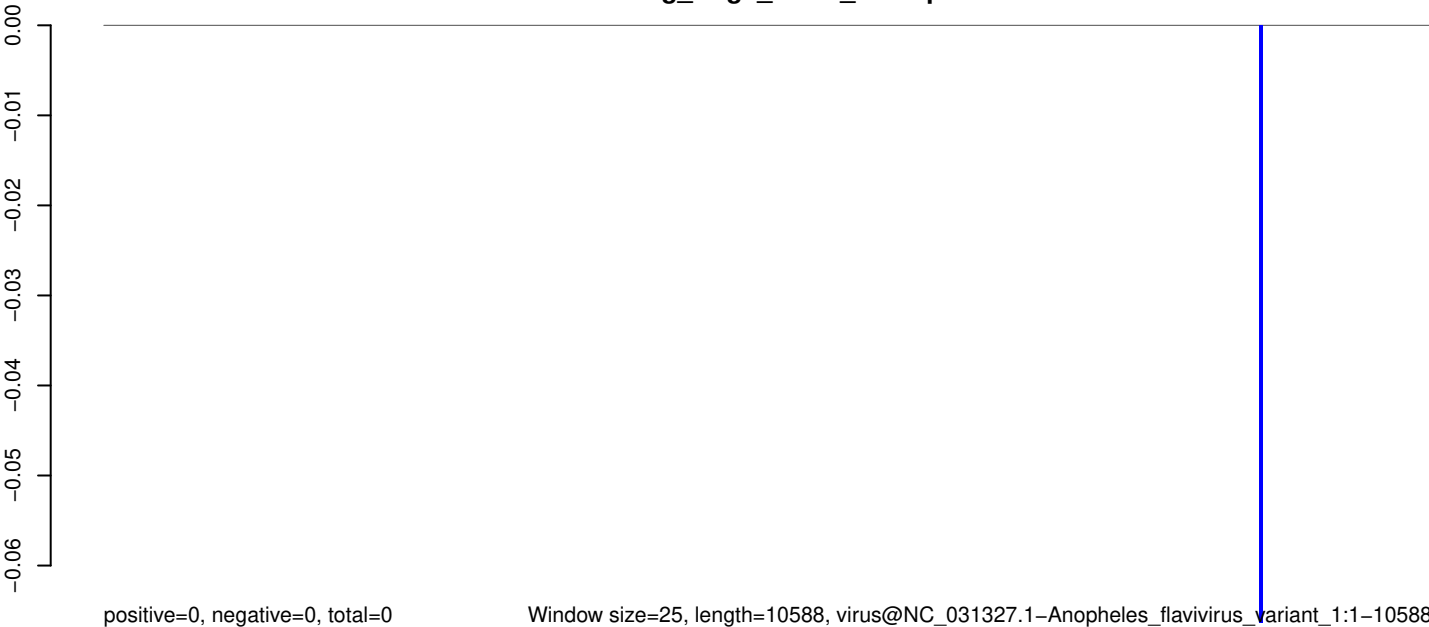
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



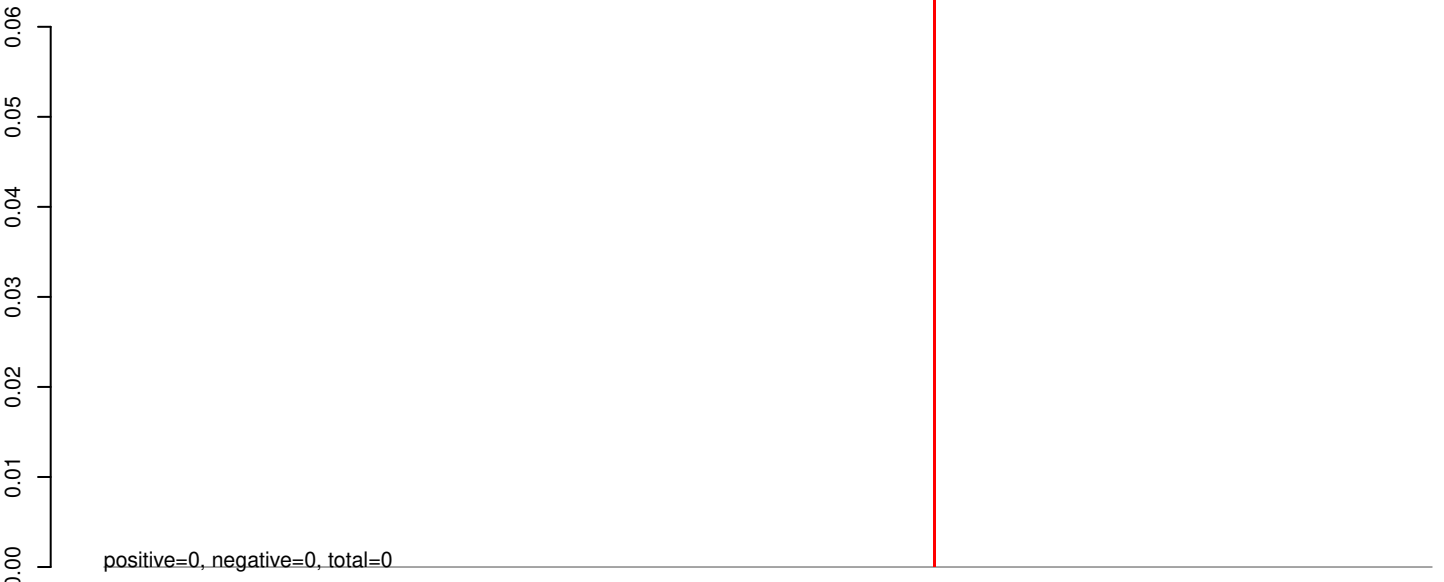
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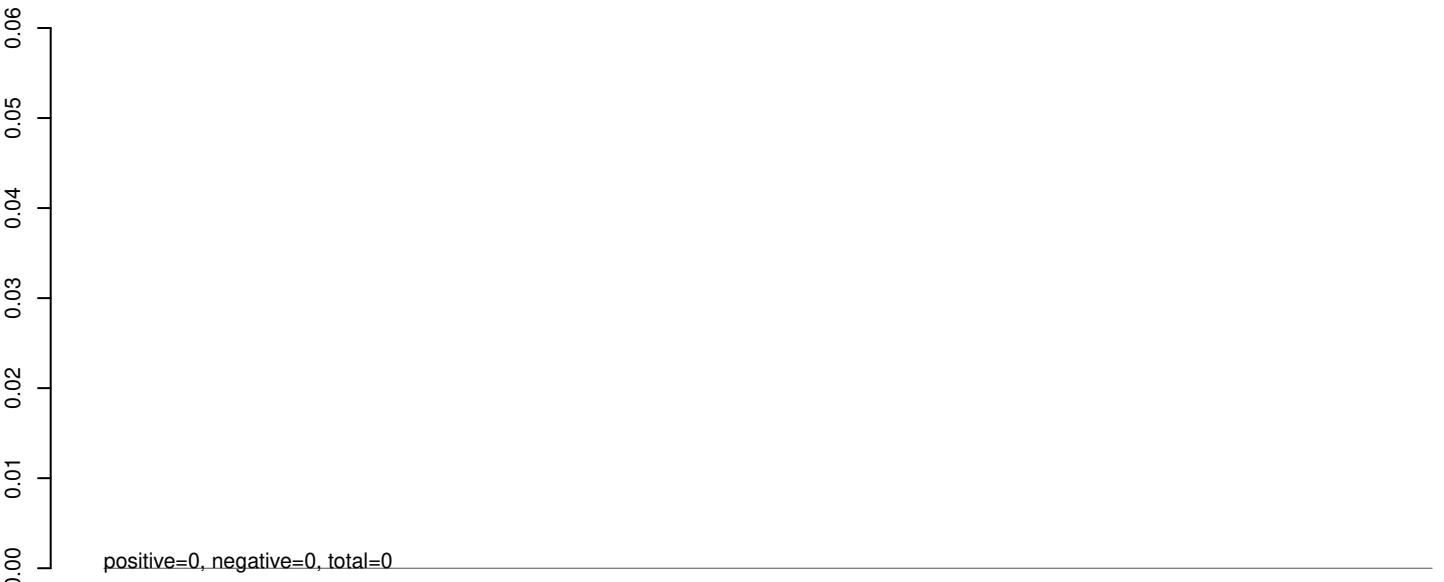
Window size=25, length=10588, virus@NC_031327.1-Anopheles_flavivirus_variant_1:1-10588

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

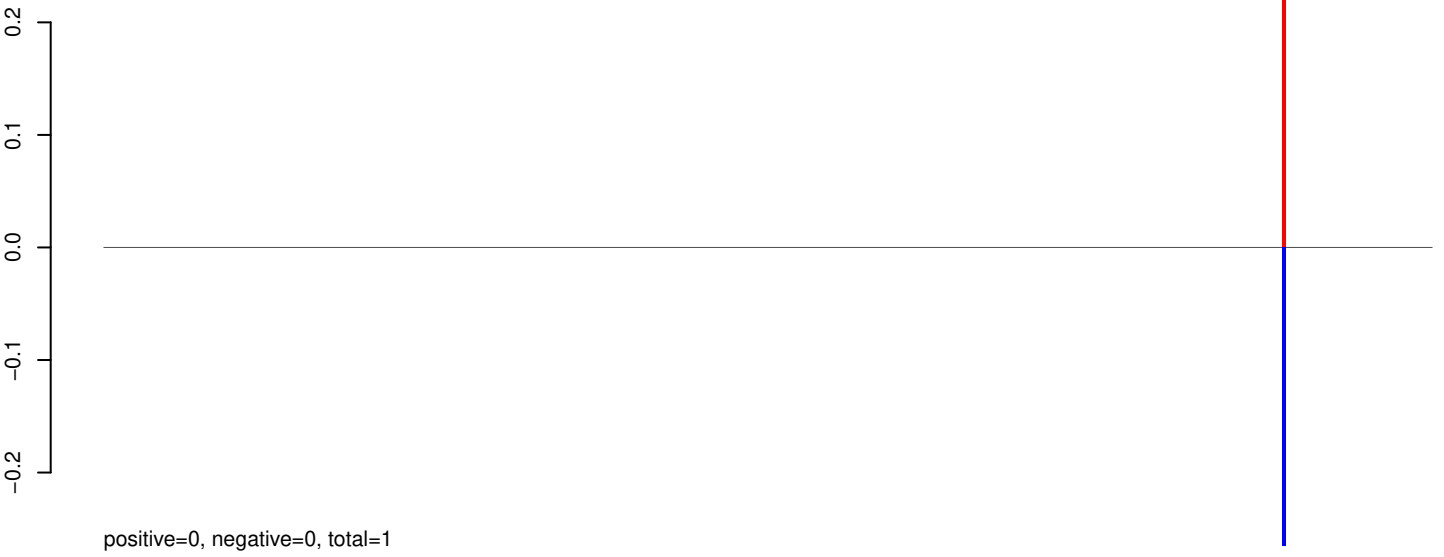


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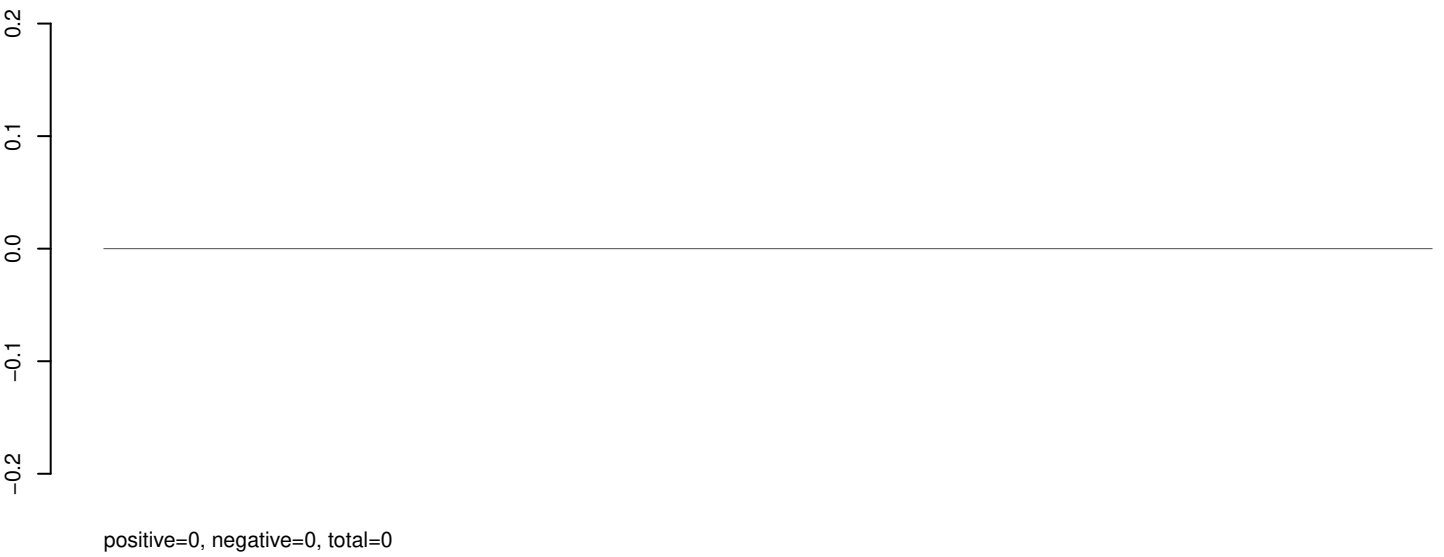


Xincheng_Mosquito_Virus_strain_XC1-6_ORF1_ORF1_ORF2_ORF2_glycoprotein_G_and_RNA-de:1-12774

AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



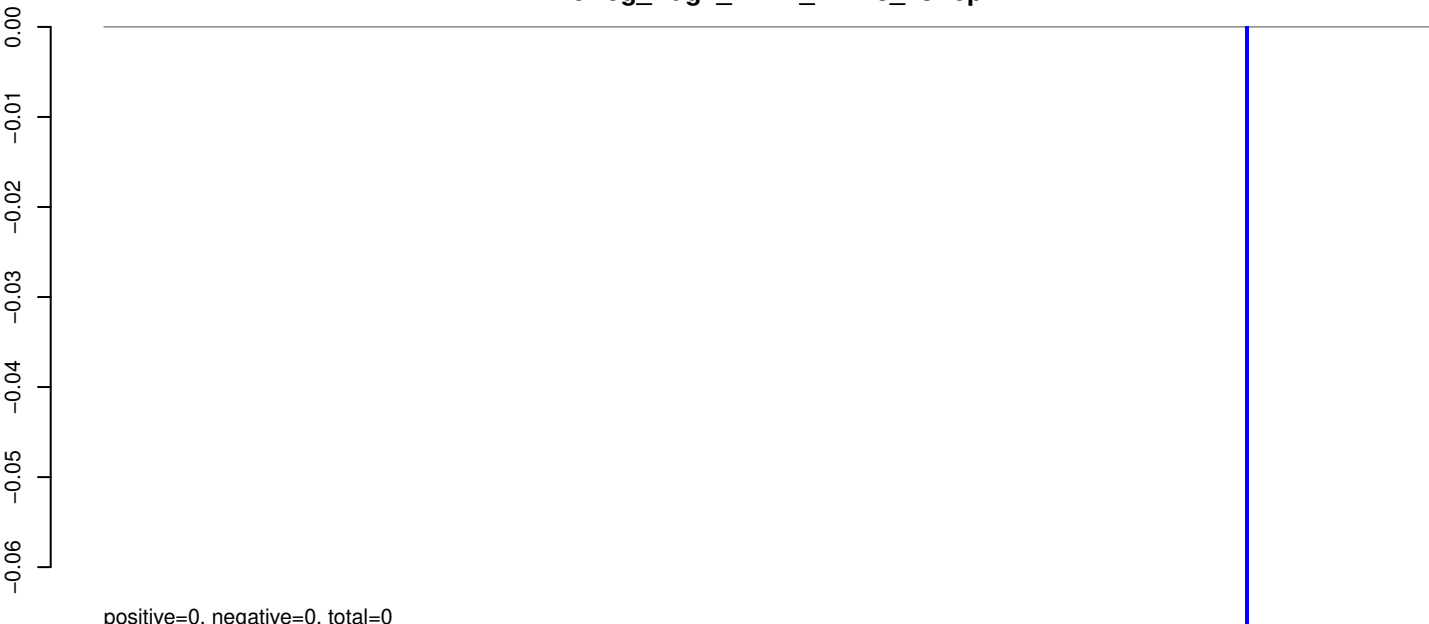
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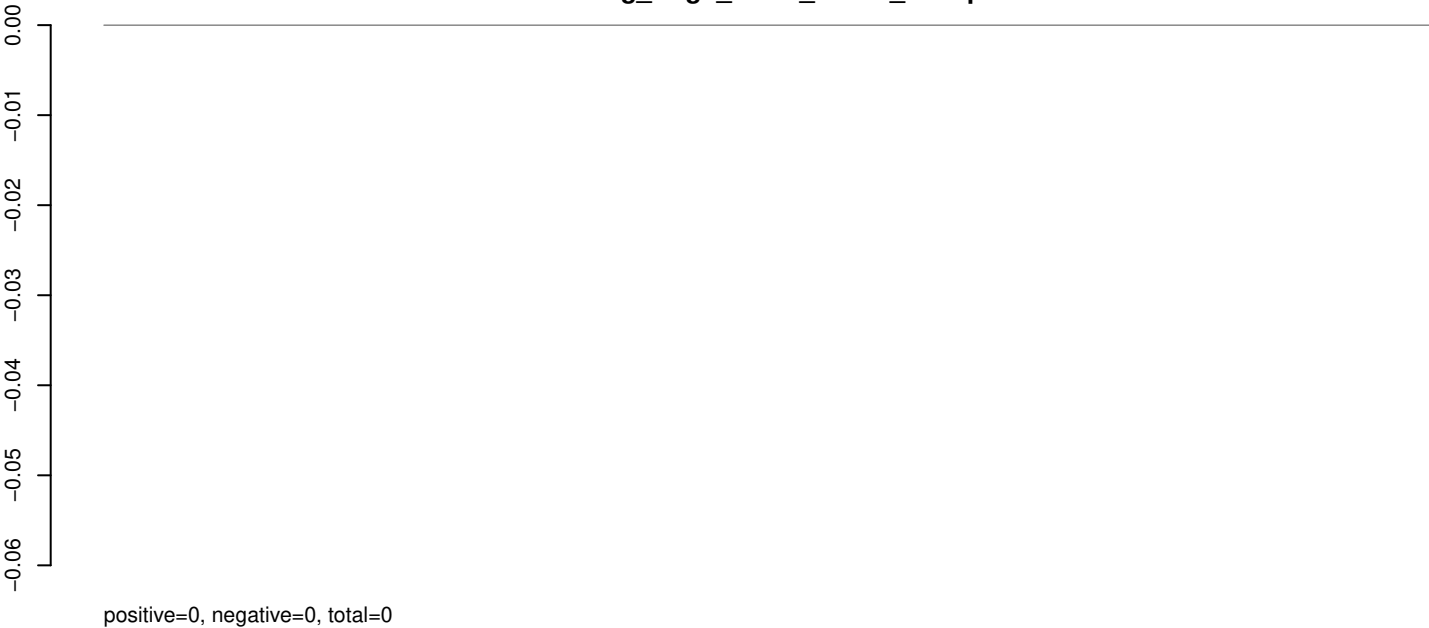
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0 2000 4000 6000 8000 10000

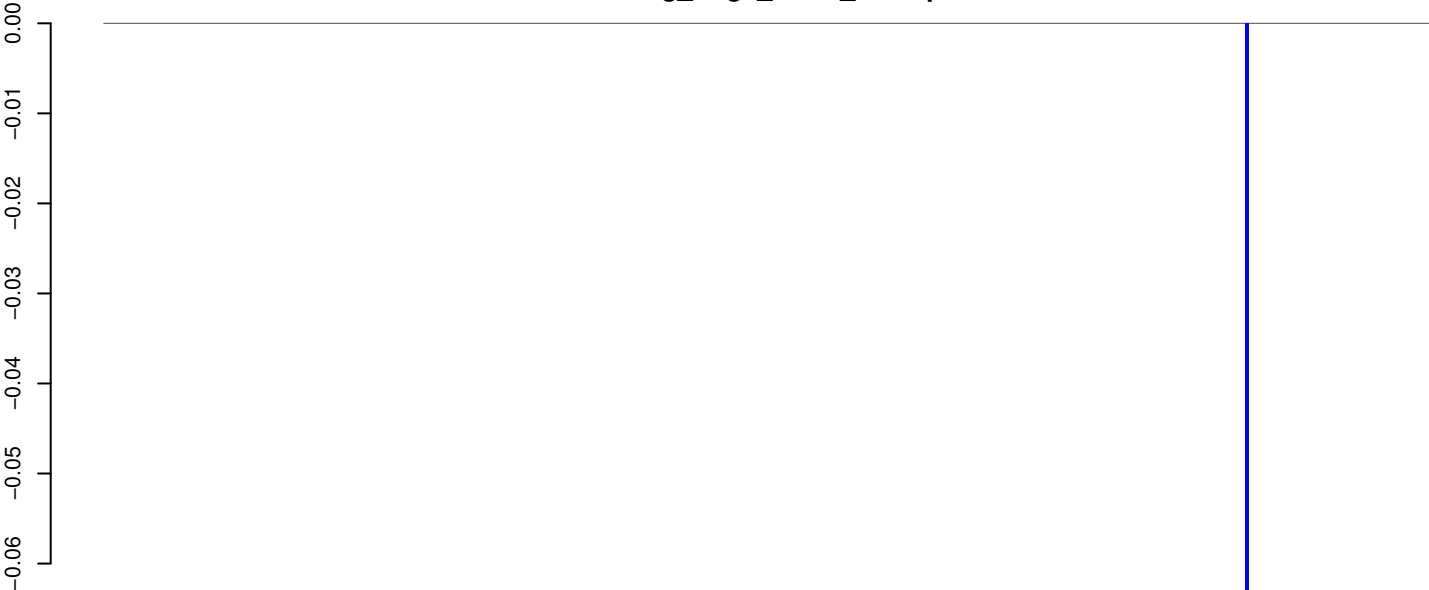
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



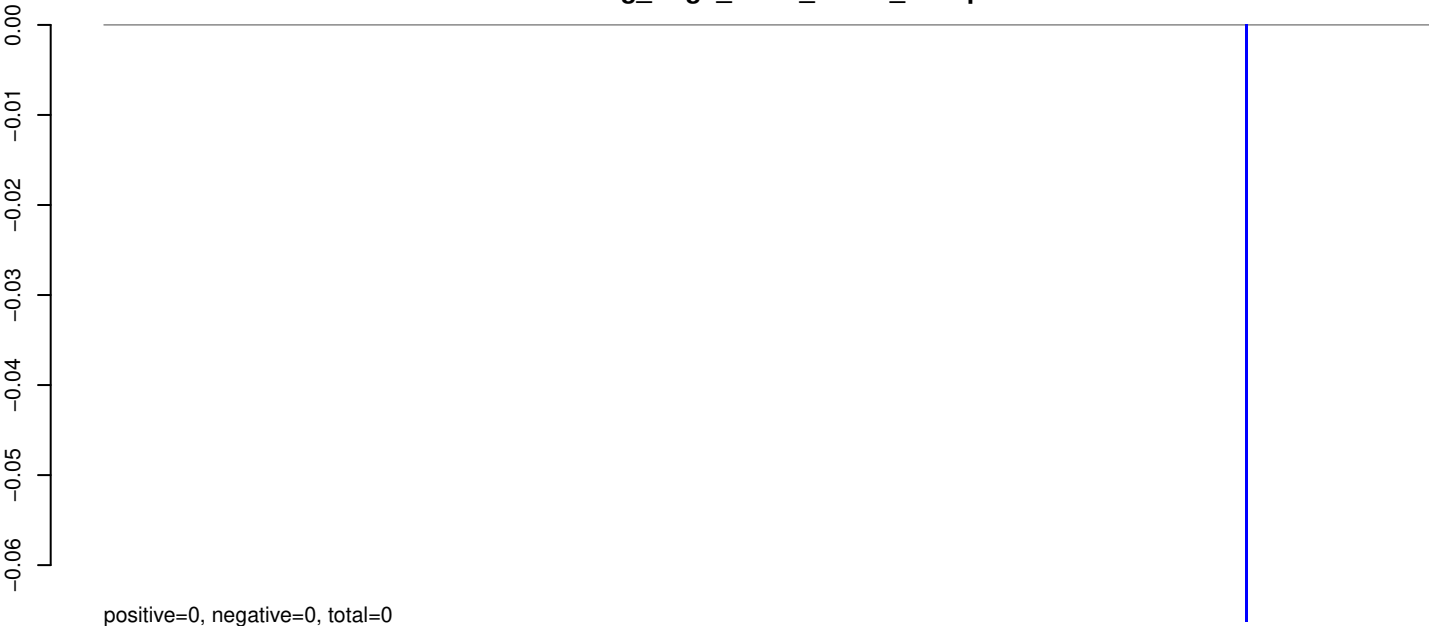
AeAeg_Aag2_BTV2_AK.rep



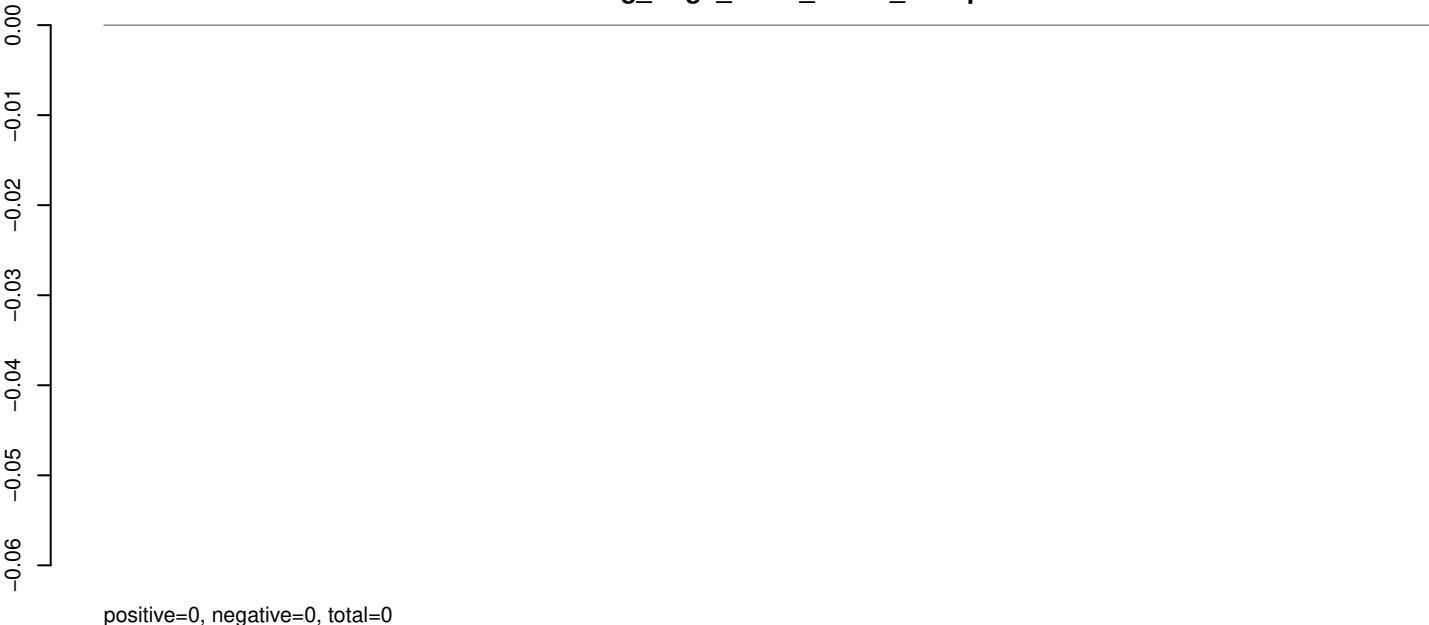
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0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



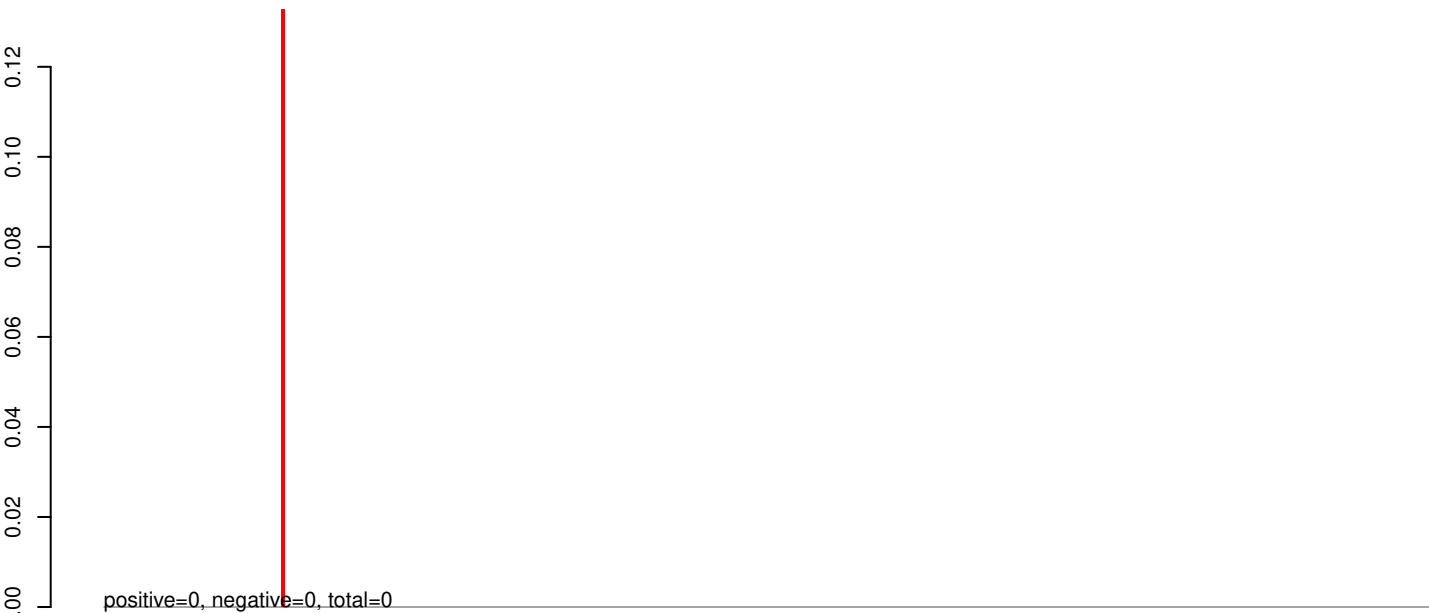
AeAeg_Aag2_BTV2_AK.rep



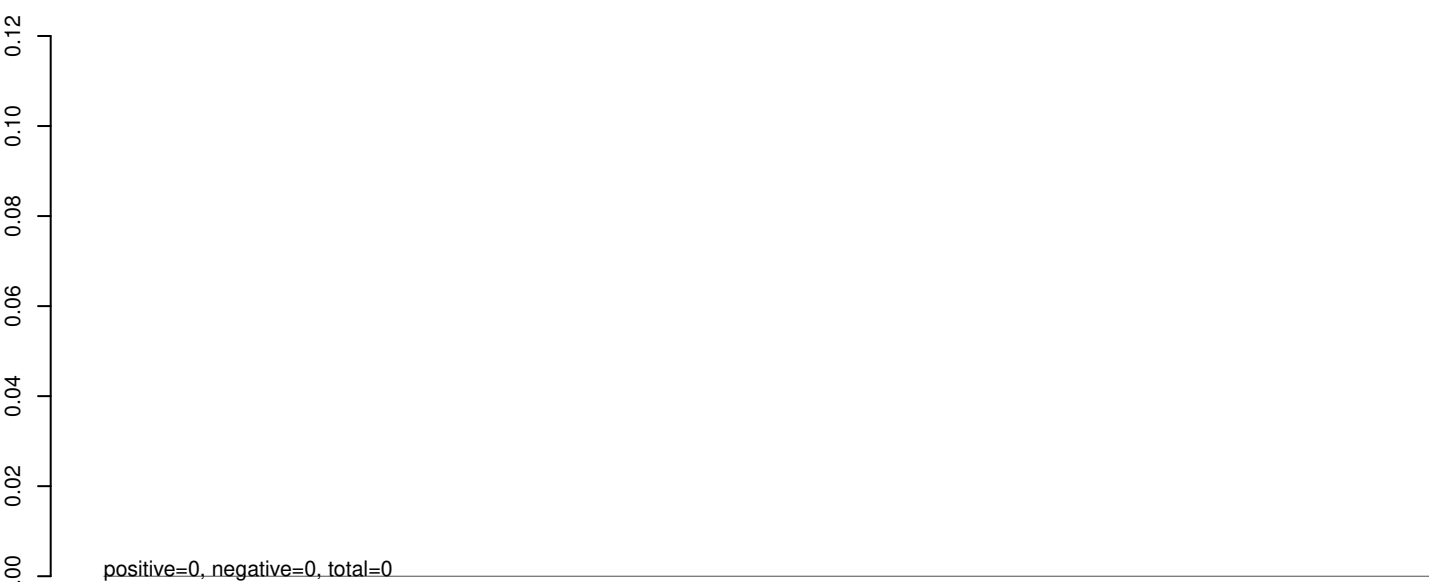
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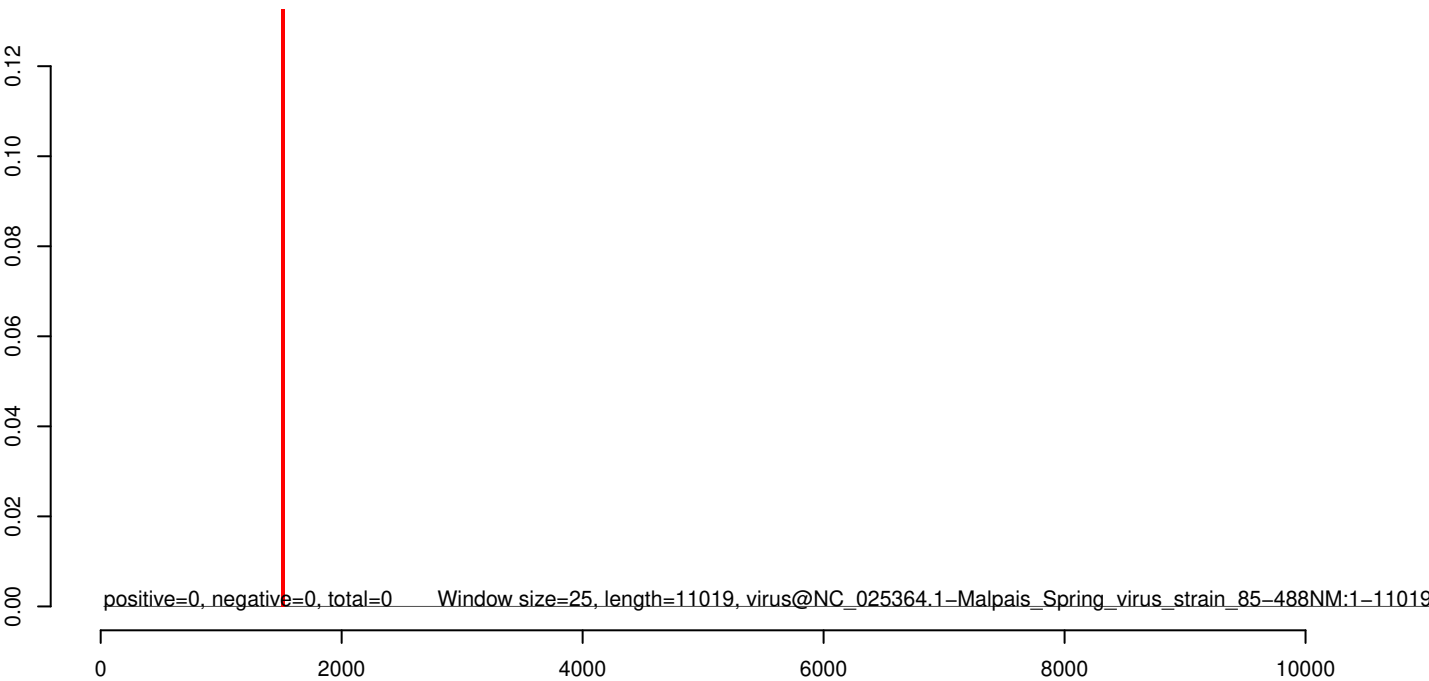
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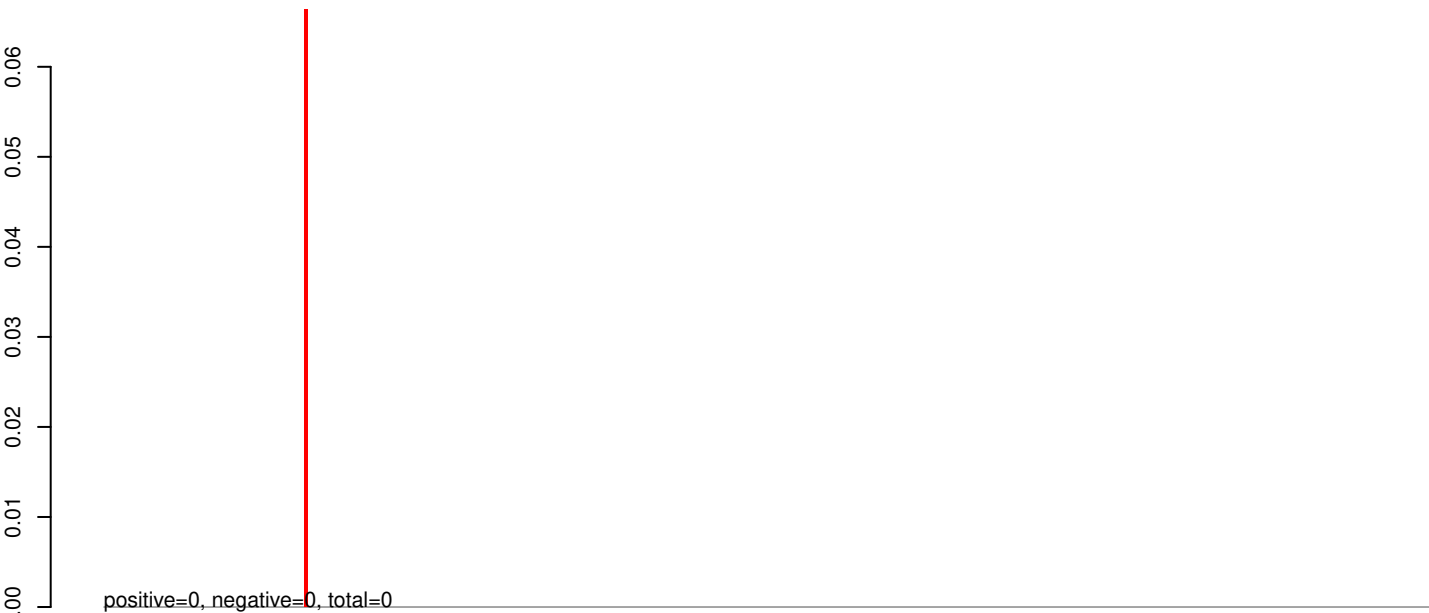
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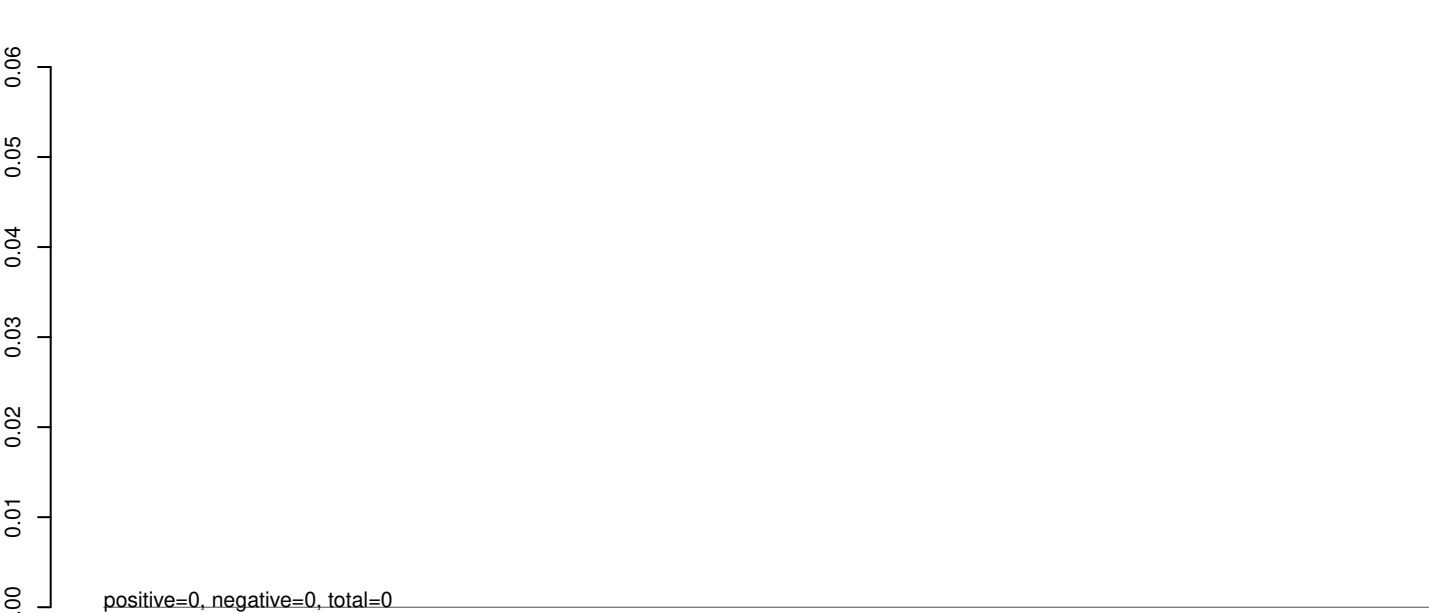
AeAeg_Aag2_BTV2_AK.rep



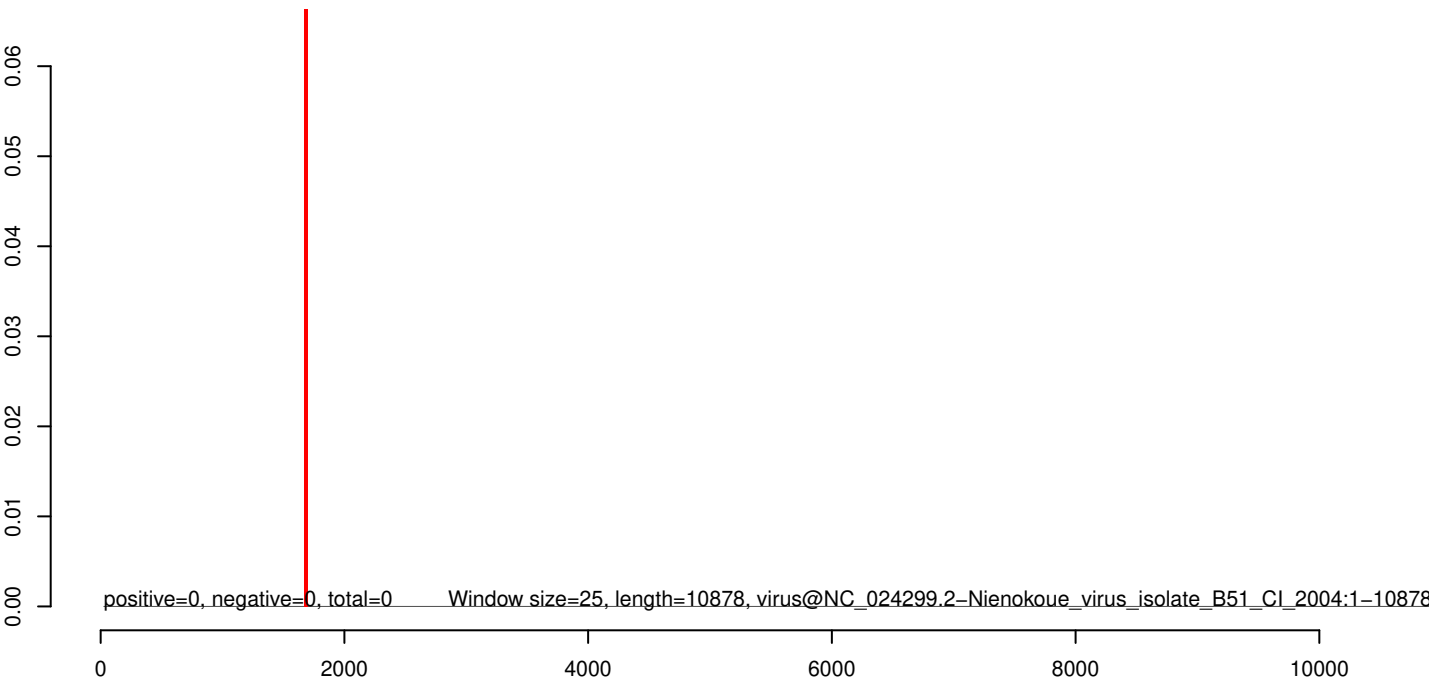
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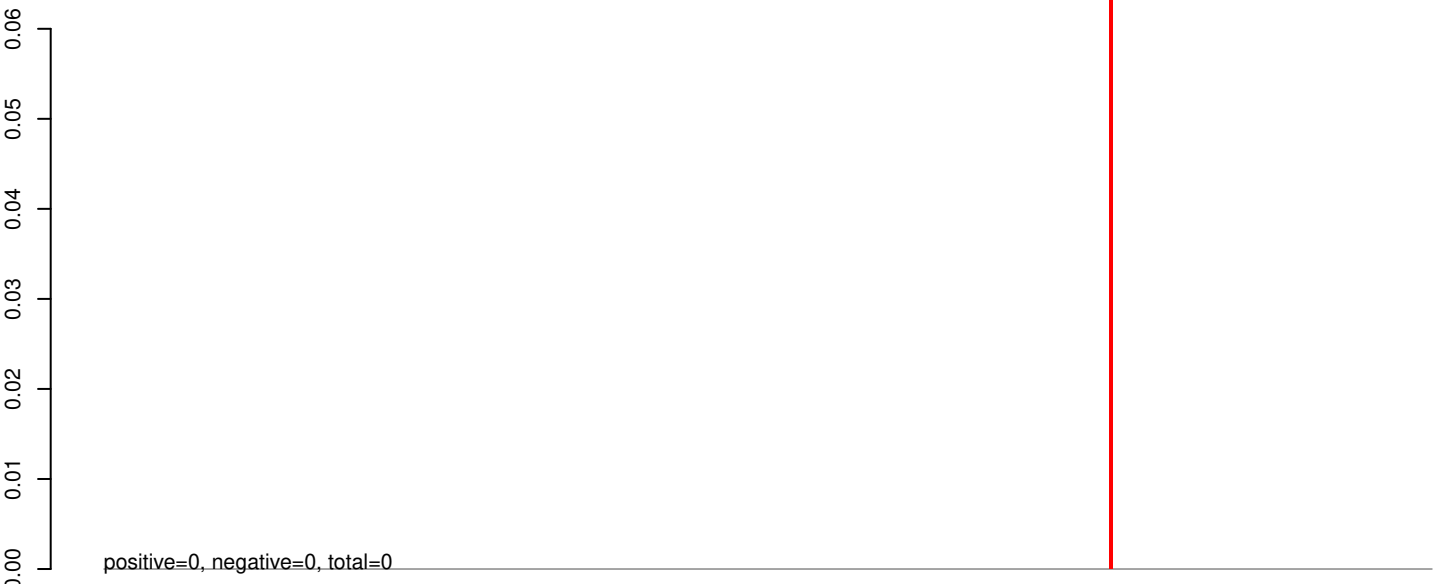
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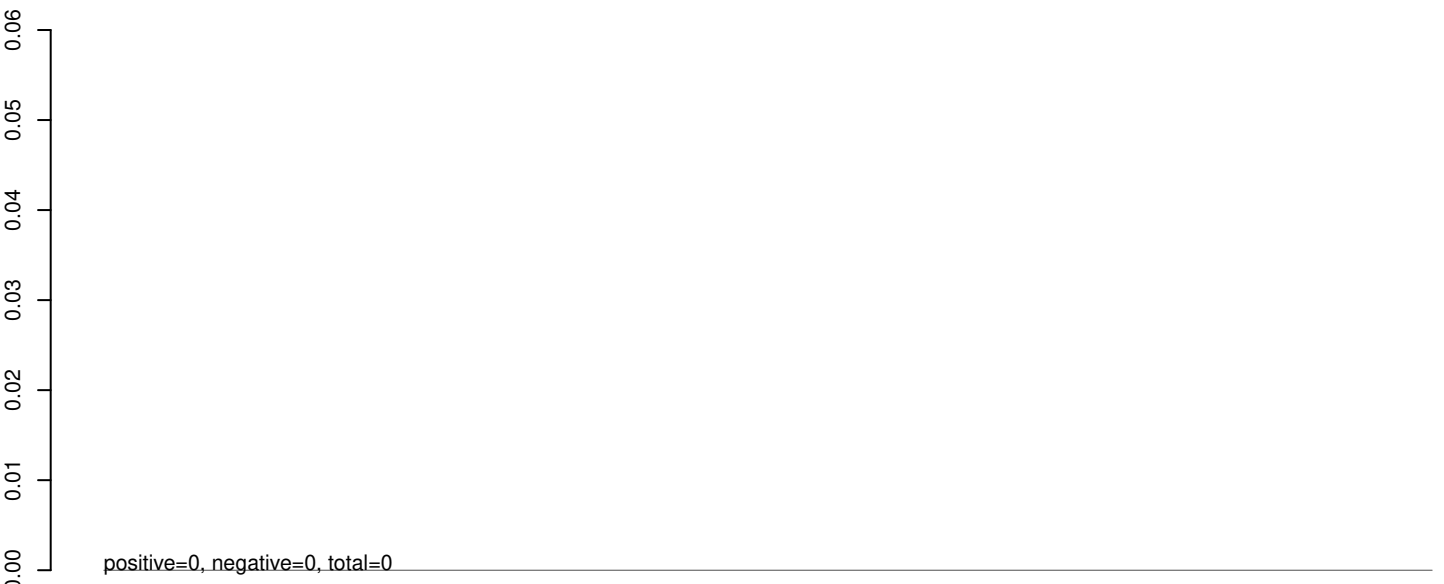
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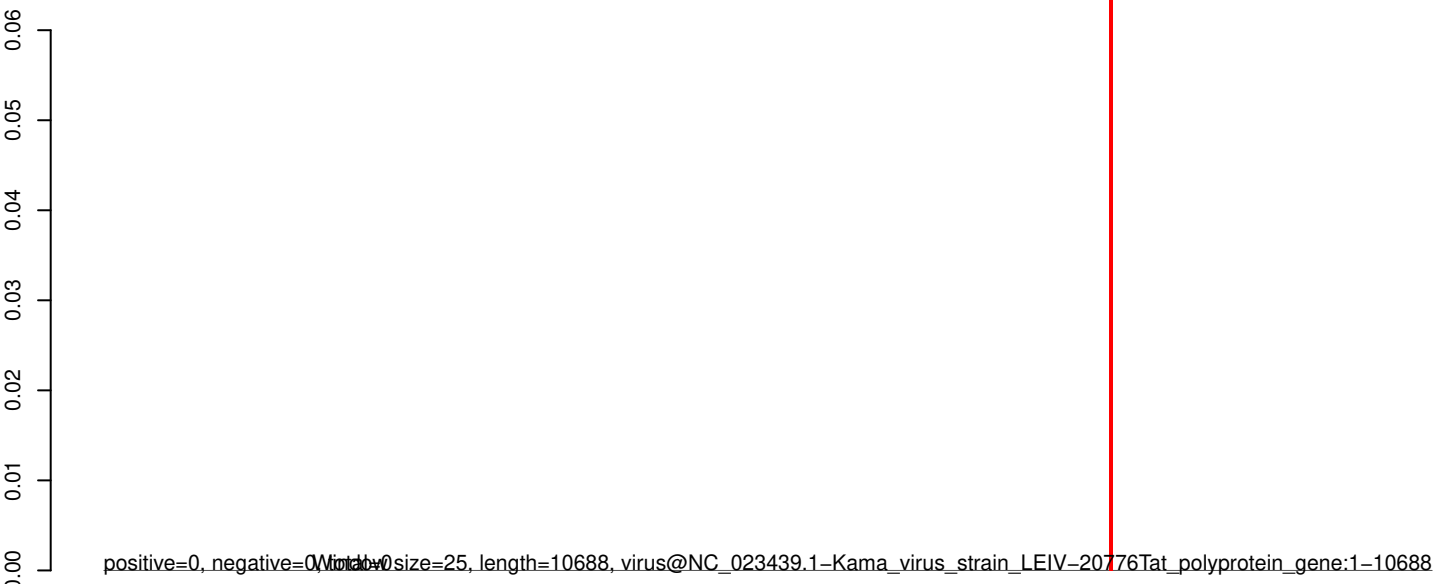
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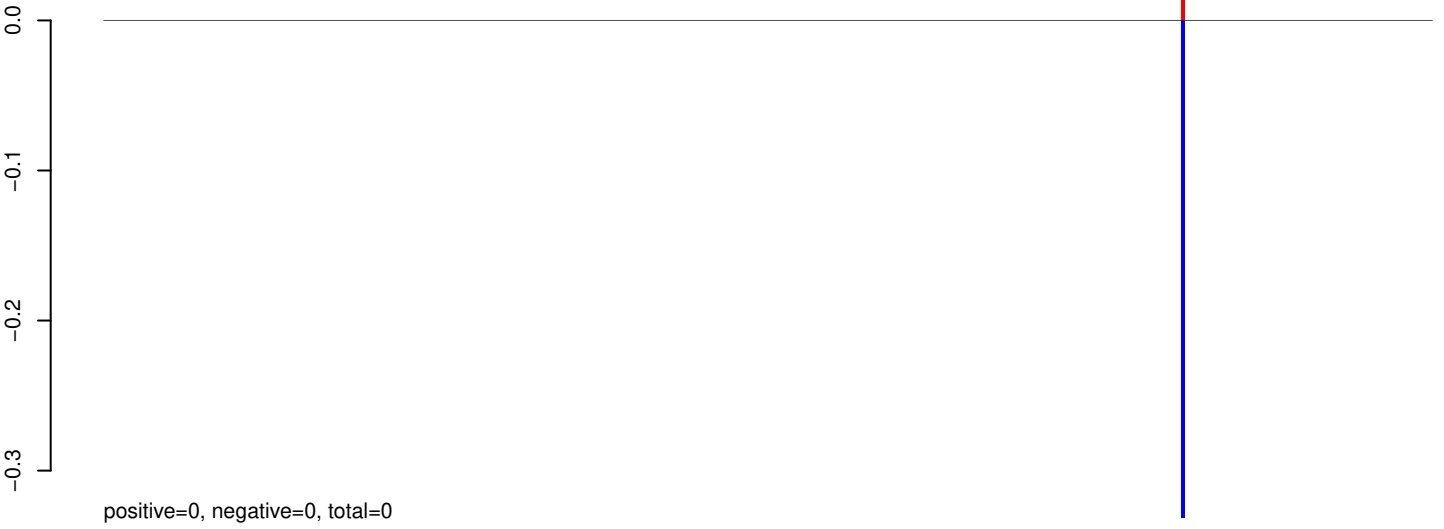
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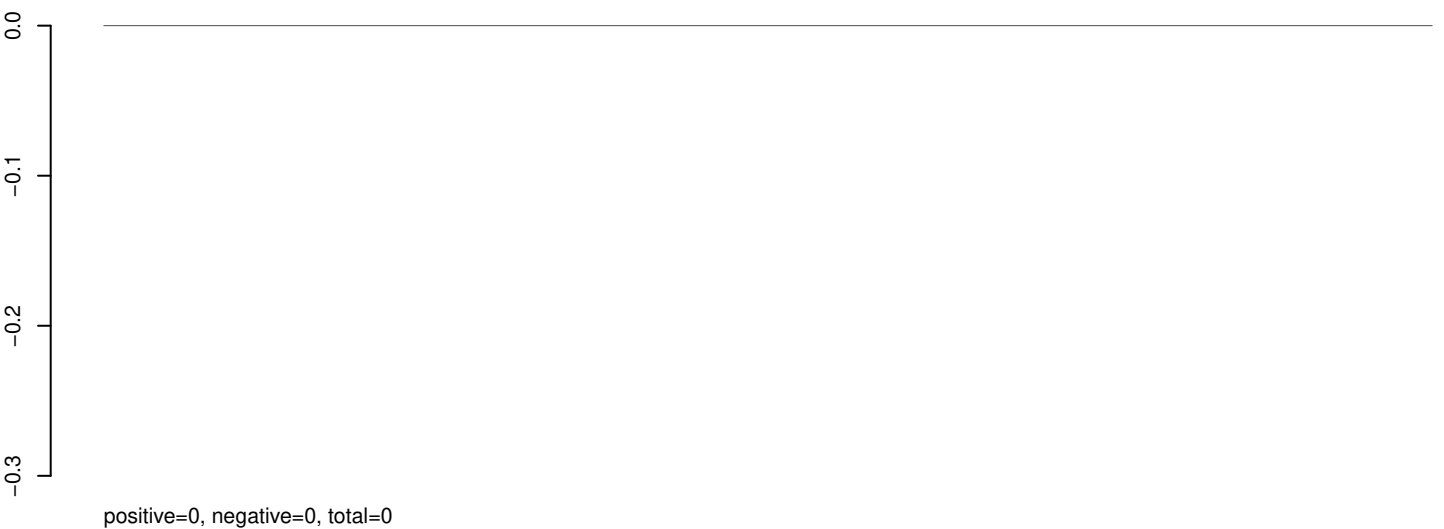
AeAeg_Aag2_BTV2_AK.rep



AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



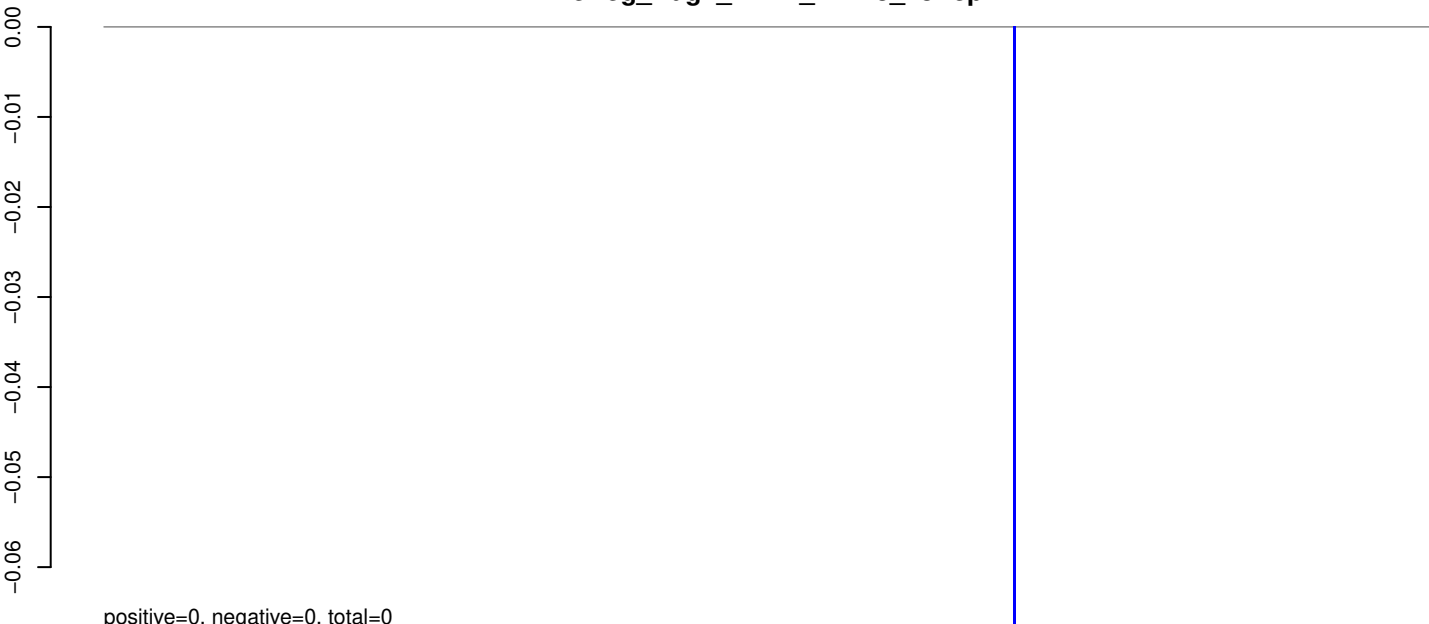
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Window size=25, length=10865, virus@NC_021069.1-Mosquito_flavivirus_isolate_LSFlaviV-A20-09:1-10865

0 2000 4000 6000 8000 10000

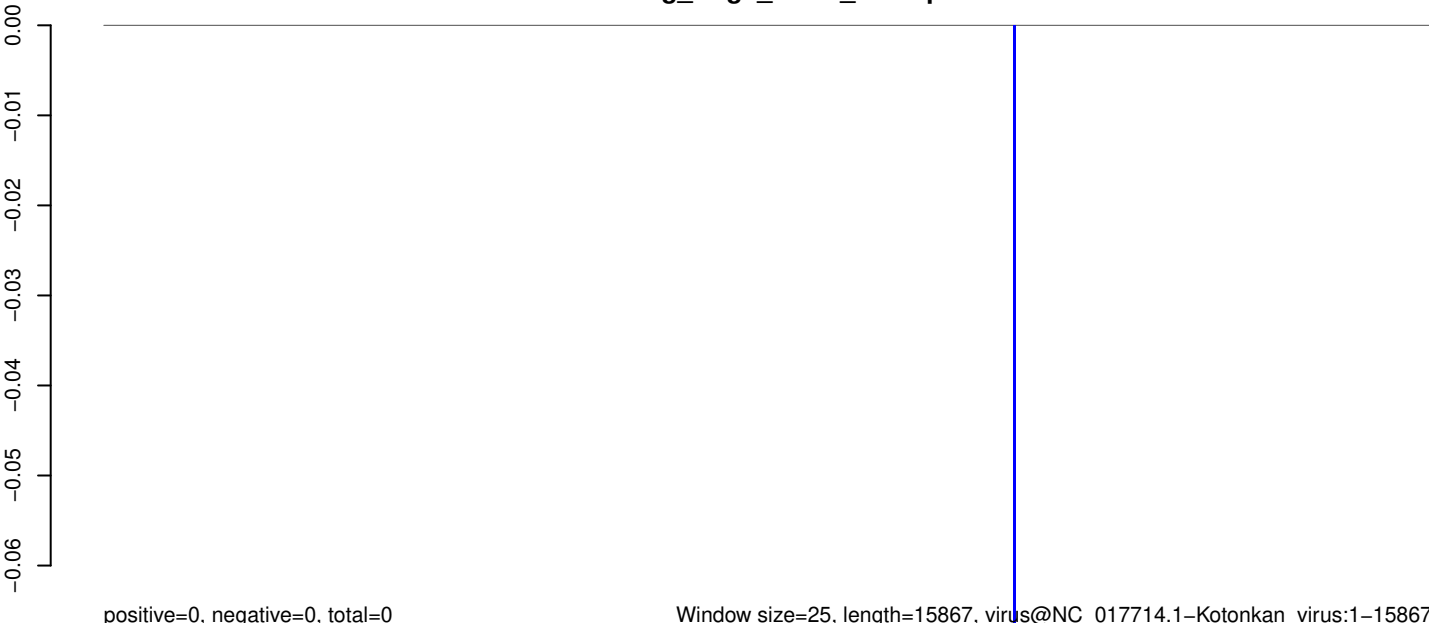
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AeAeg_Aag2_BTV2_AK.24_35.rep



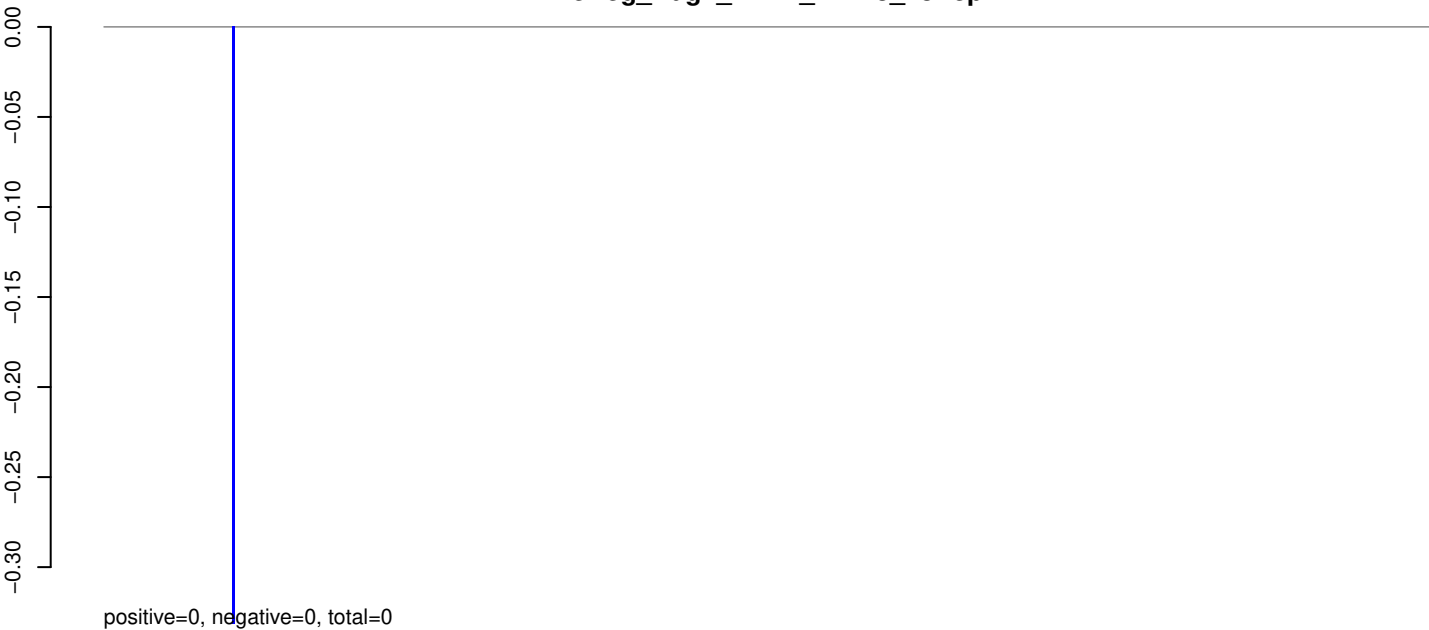
AeAeg_Aag2_BTV2_AK.rep



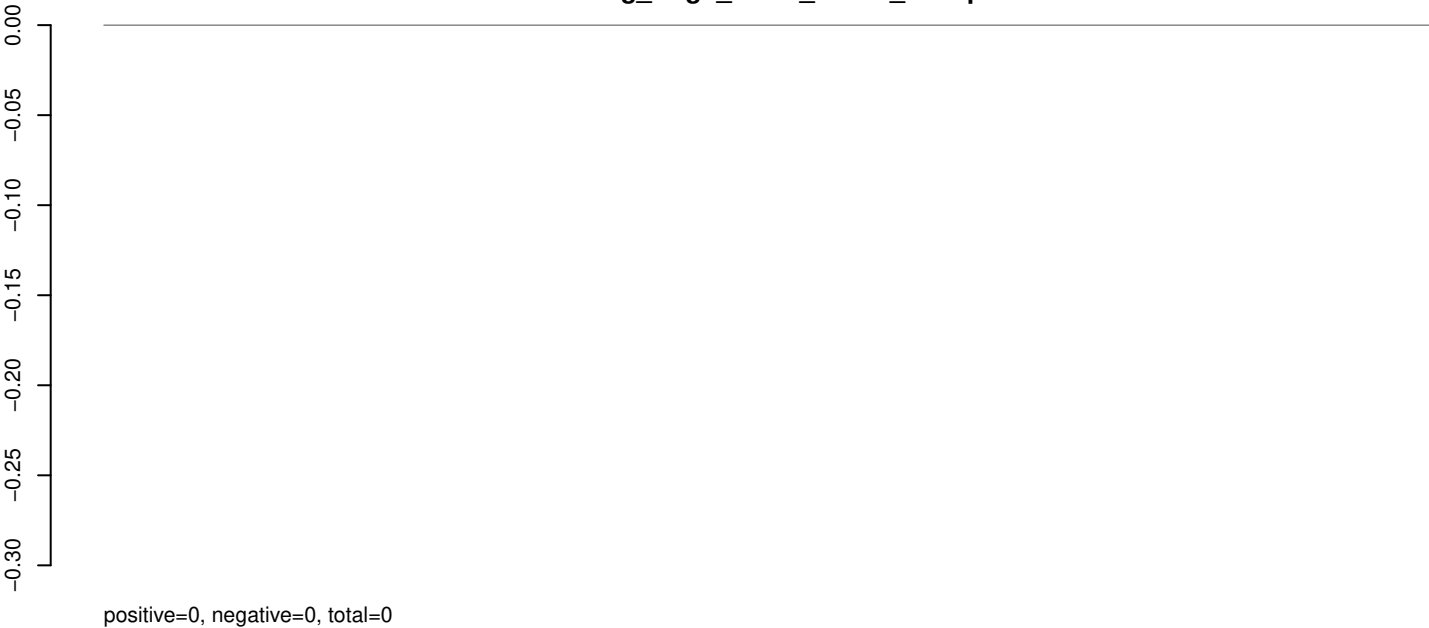
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0 5000 10000 15000

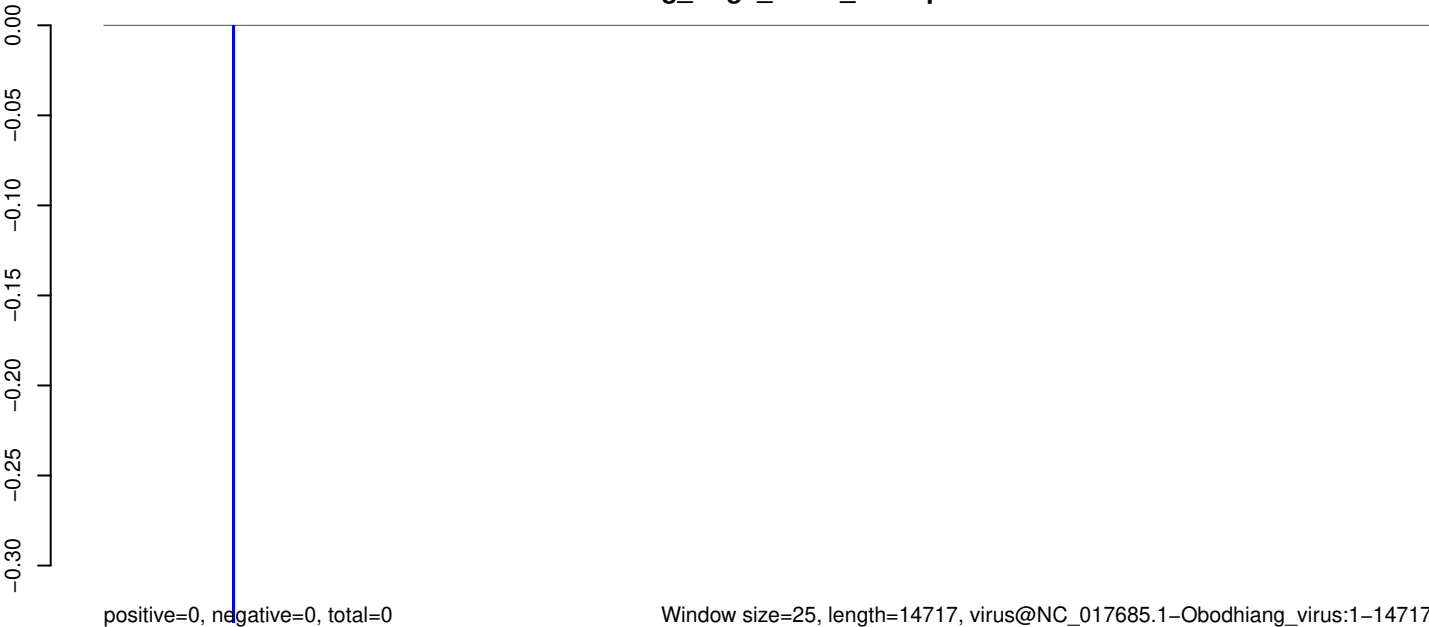
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



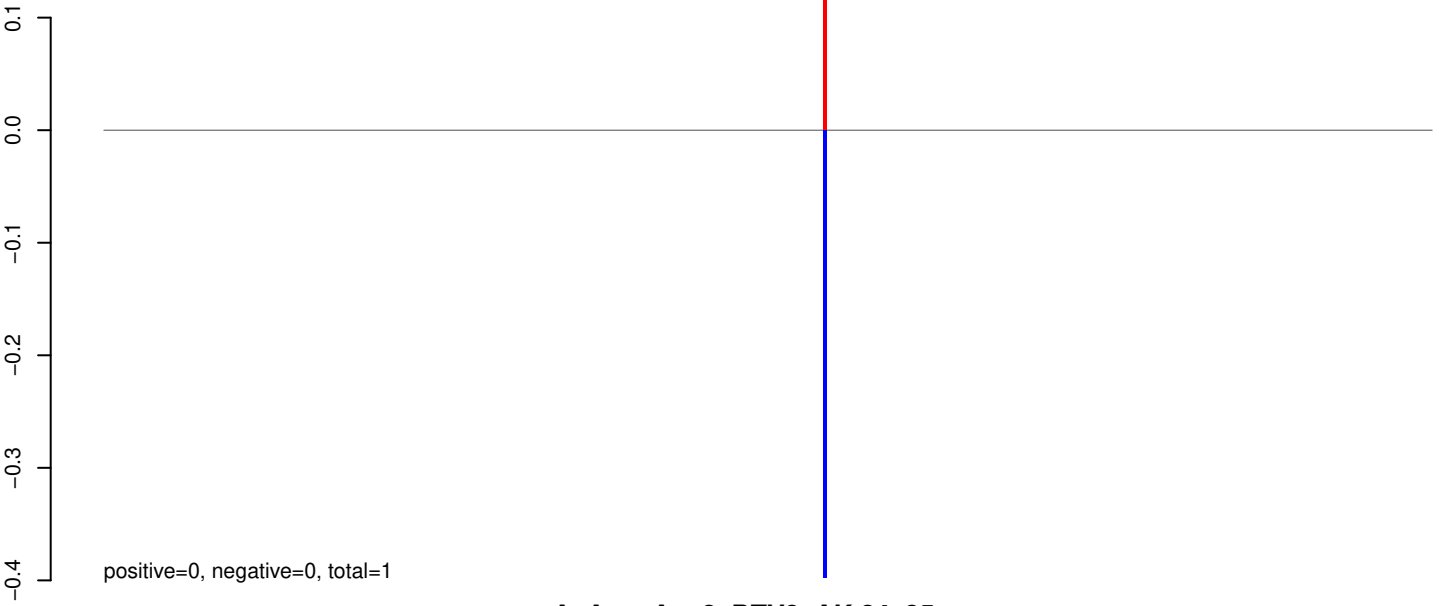
AeAeg_Aag2_BTV2_AK.rep



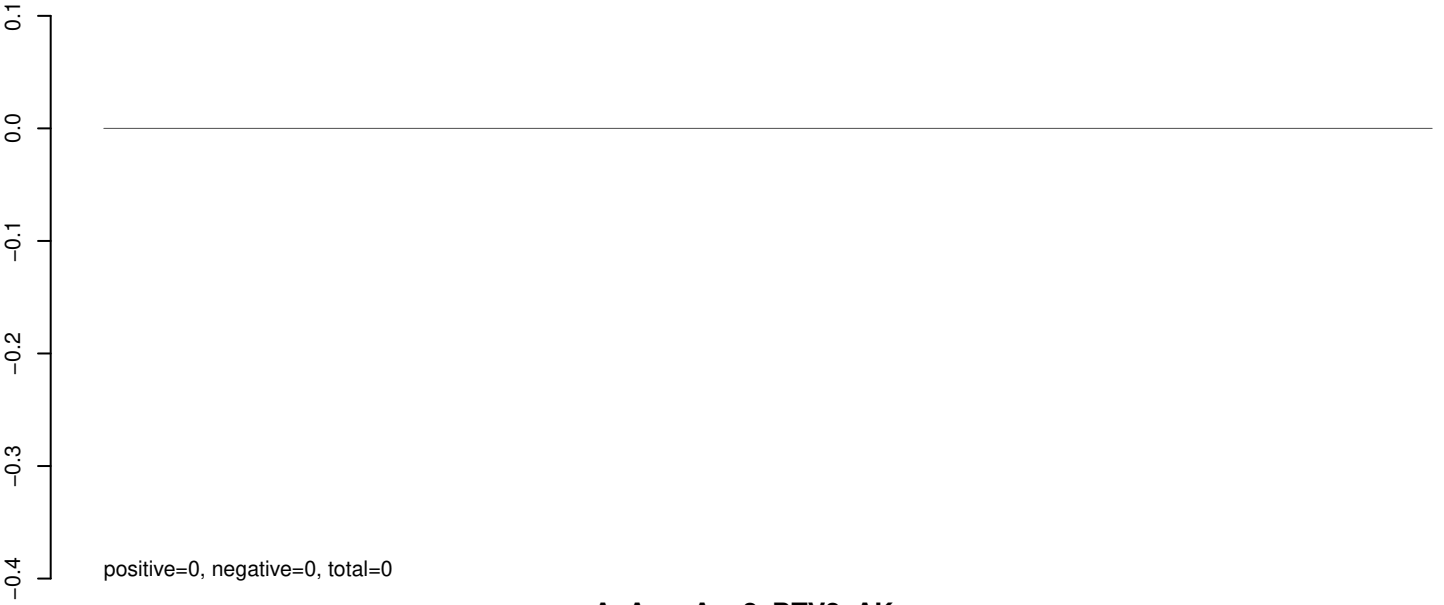
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0 5000 10000 15000

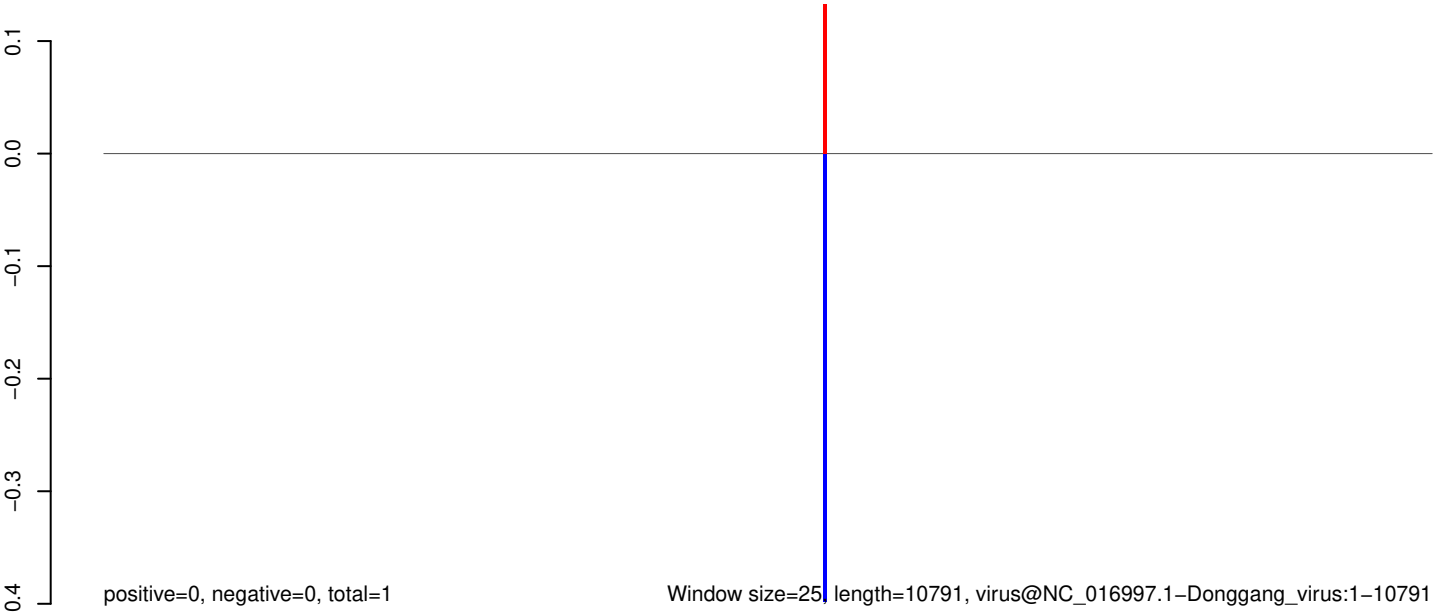
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

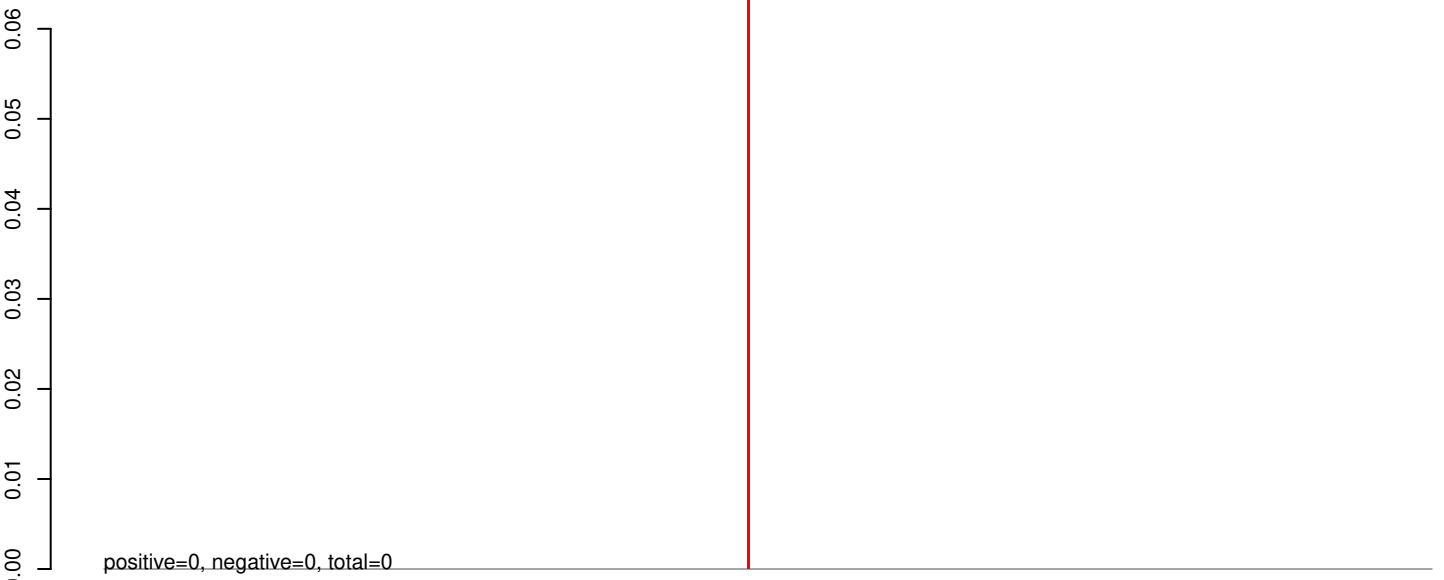


AeAeg_Aag2_BTV2_AK.rep

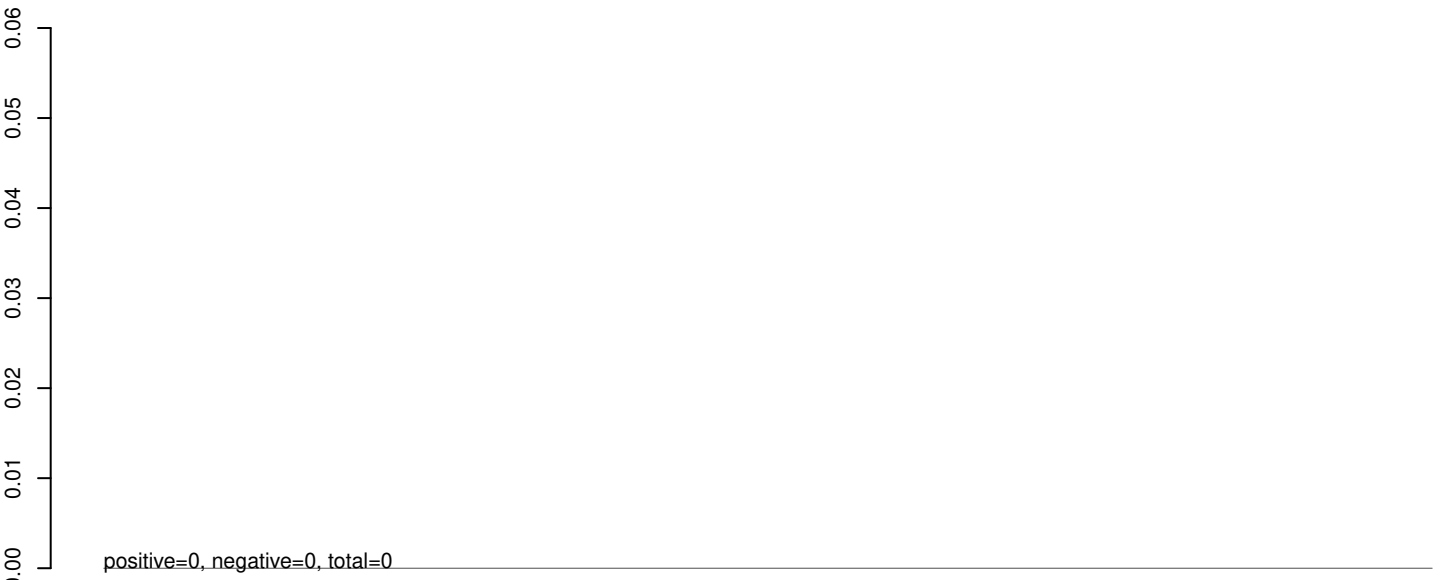


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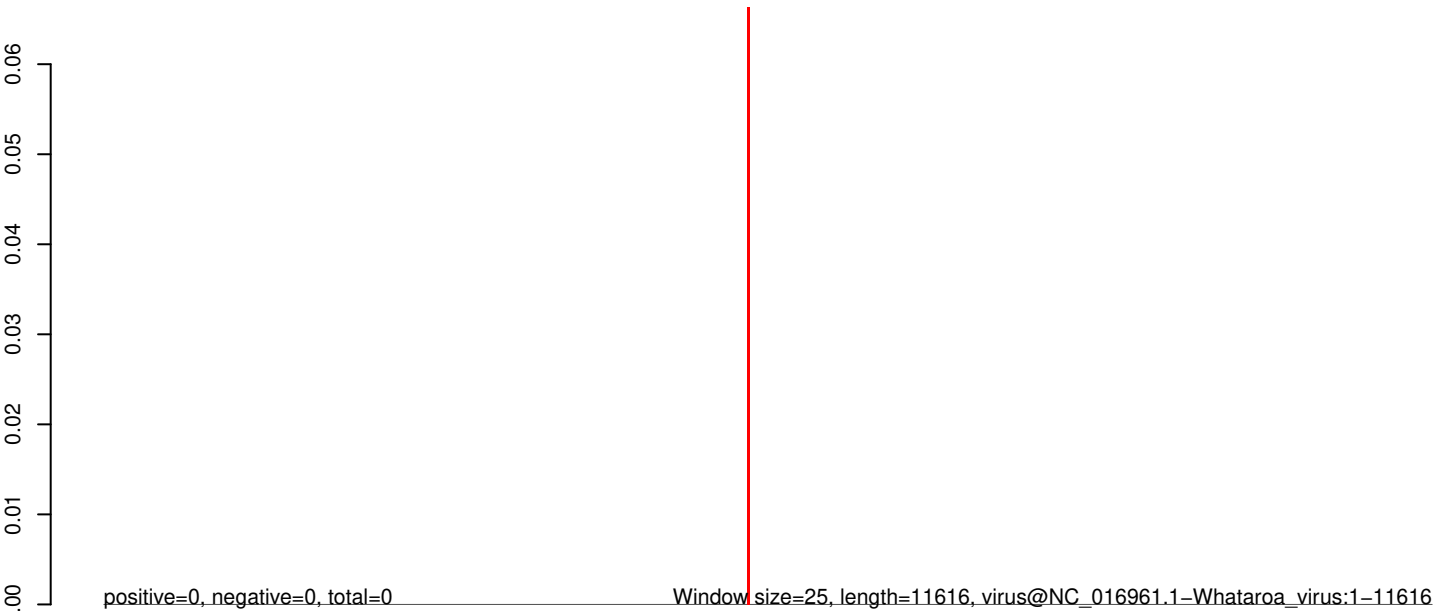
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AeAeg_Aag2_BTV2_AK.24_35.rep

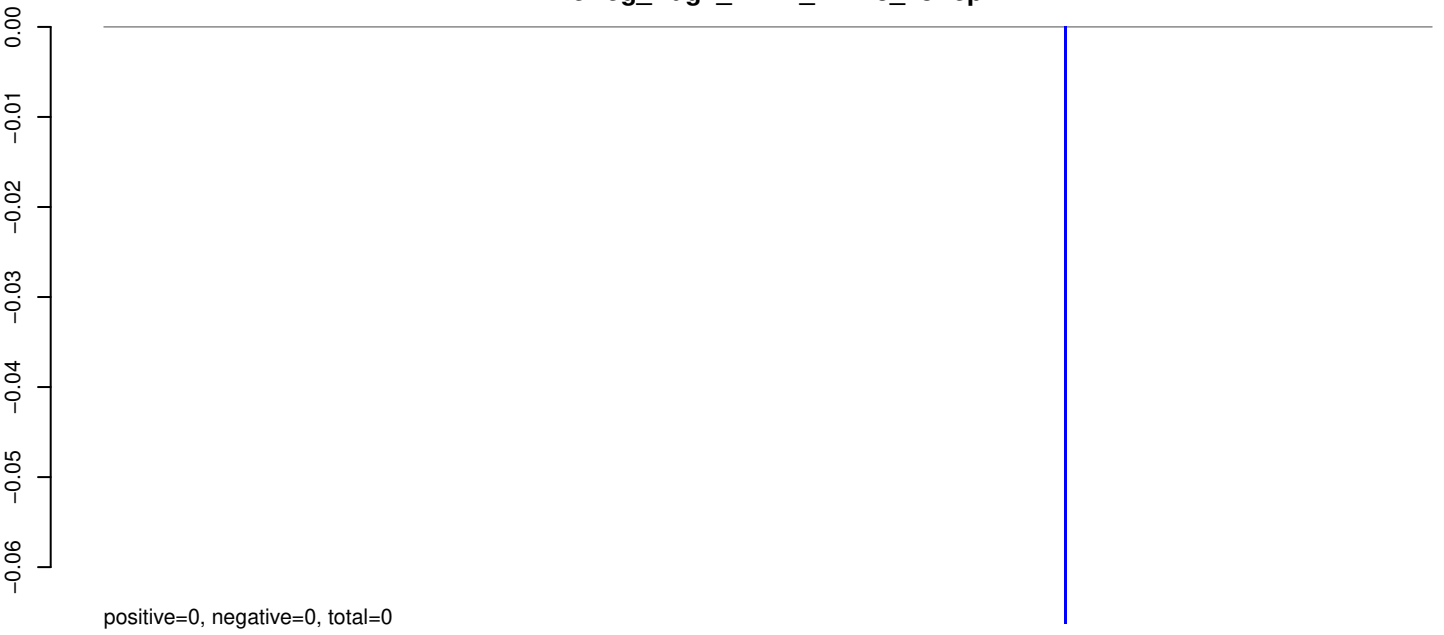


AeAeg_Aag2_BTV2_AK.rep

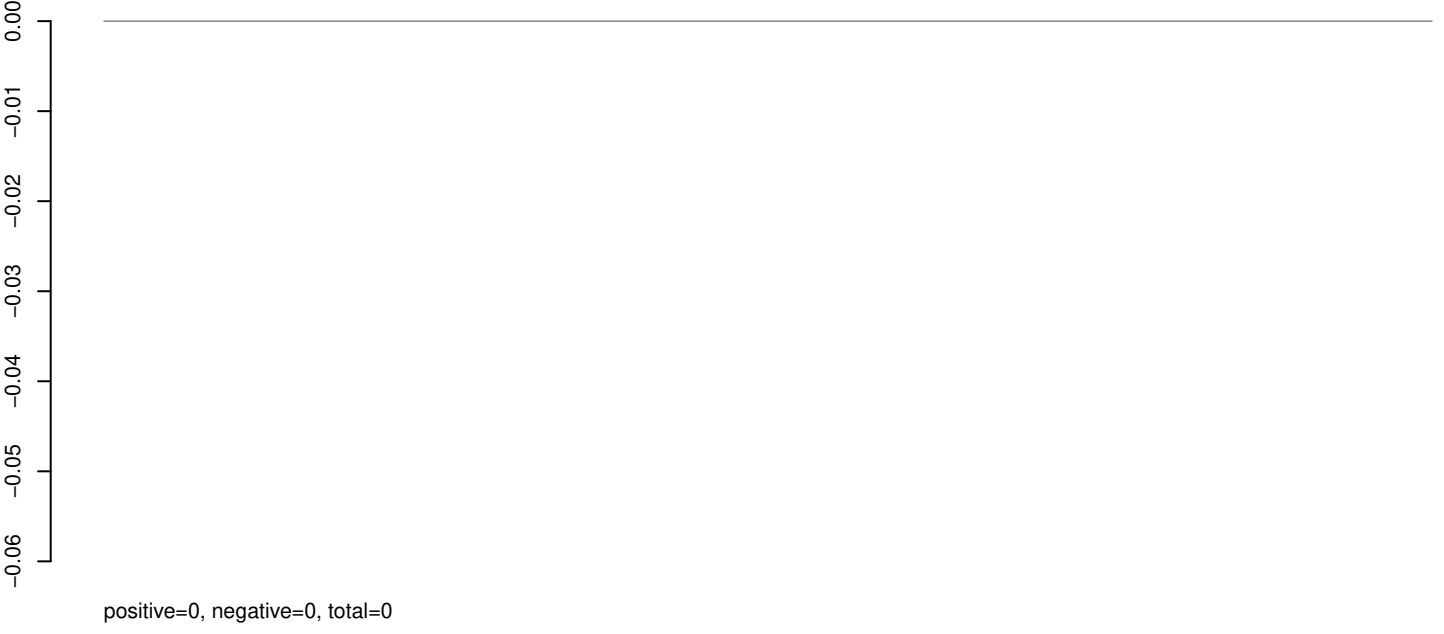


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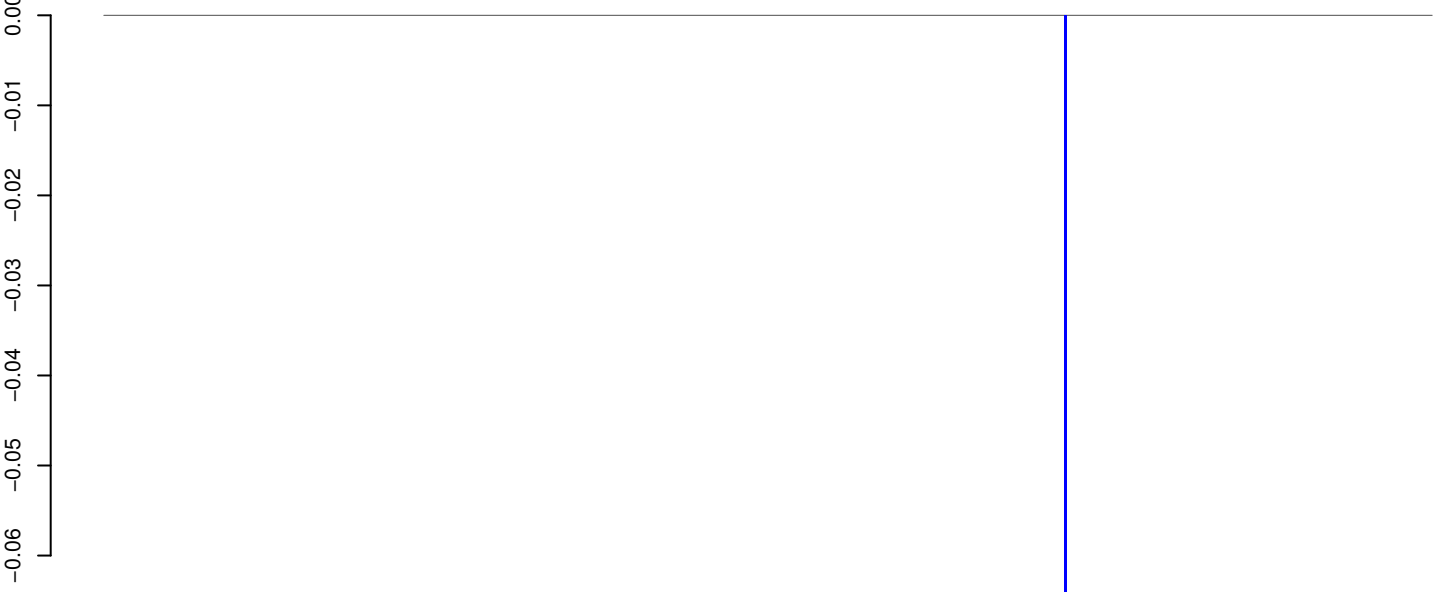
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



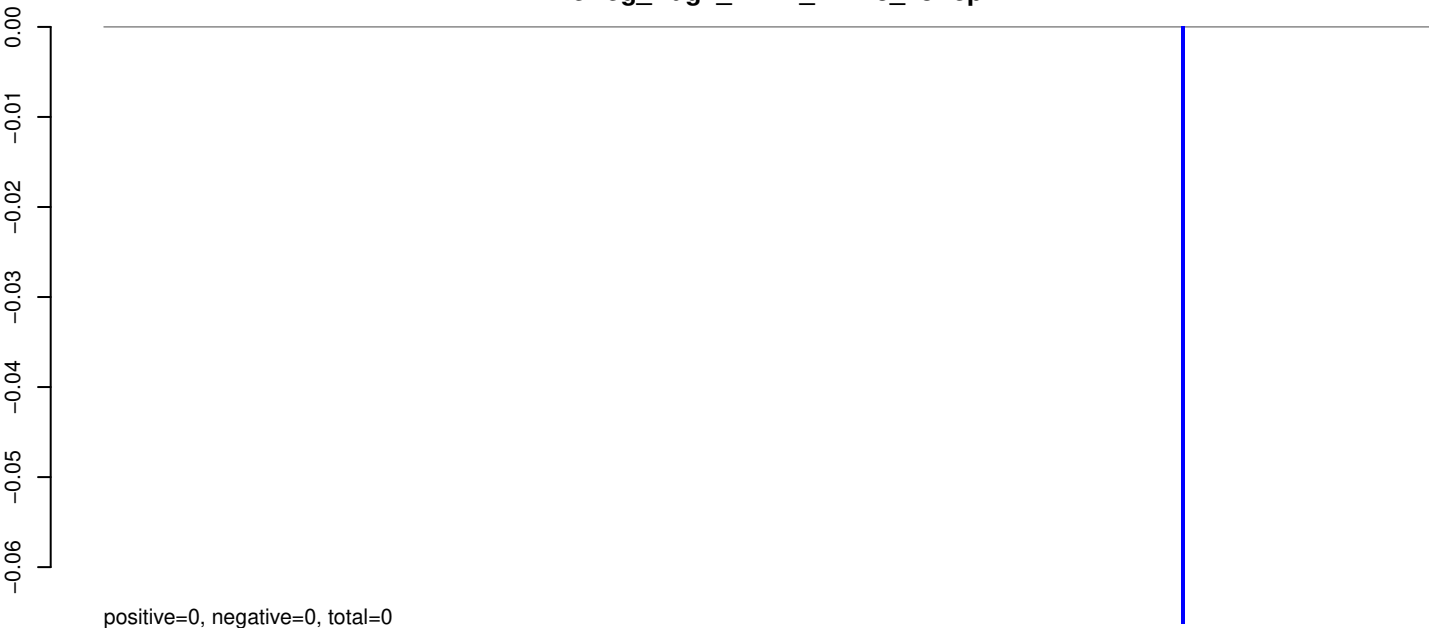
AeAeg_Aag2_BTV2_AK.rep



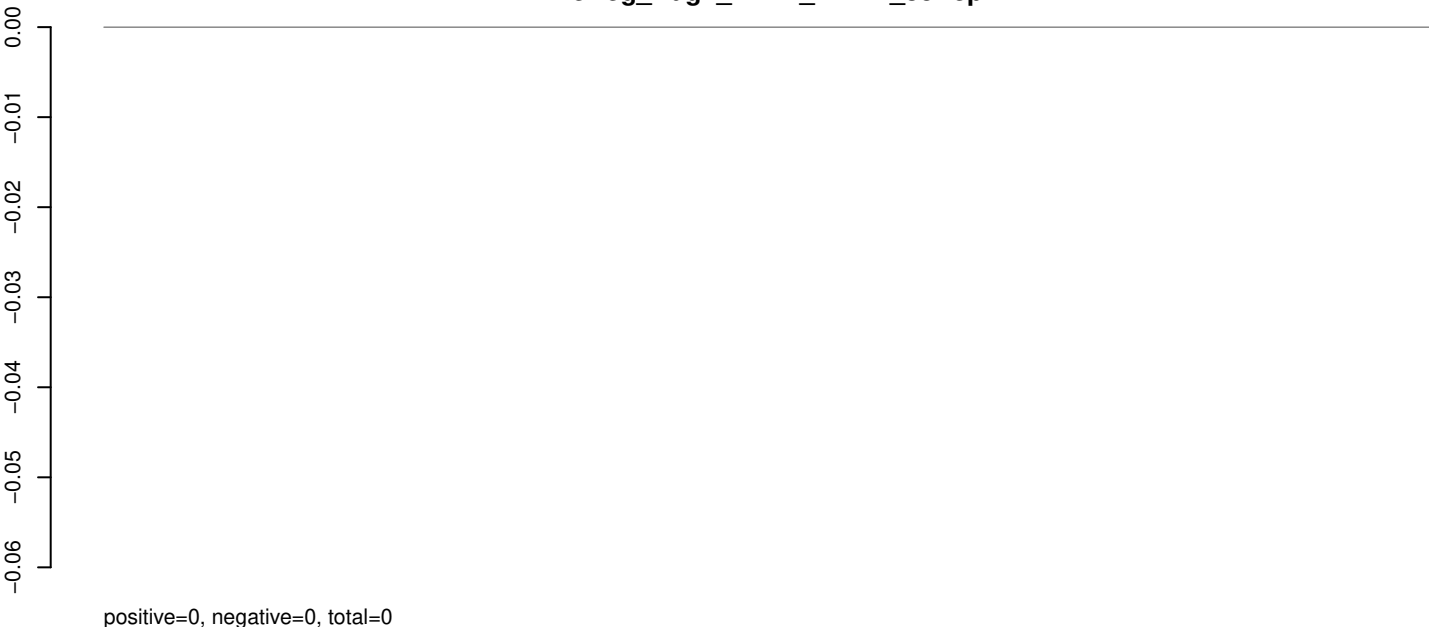
Window size: 25, step: 1 1979, virus@NC_014395.1-Rift_Valley_fever_virus_segment_S_plus_segment_M_plus_segment_L:1-11979

0 2000 4000 6000 8000 10000 12000

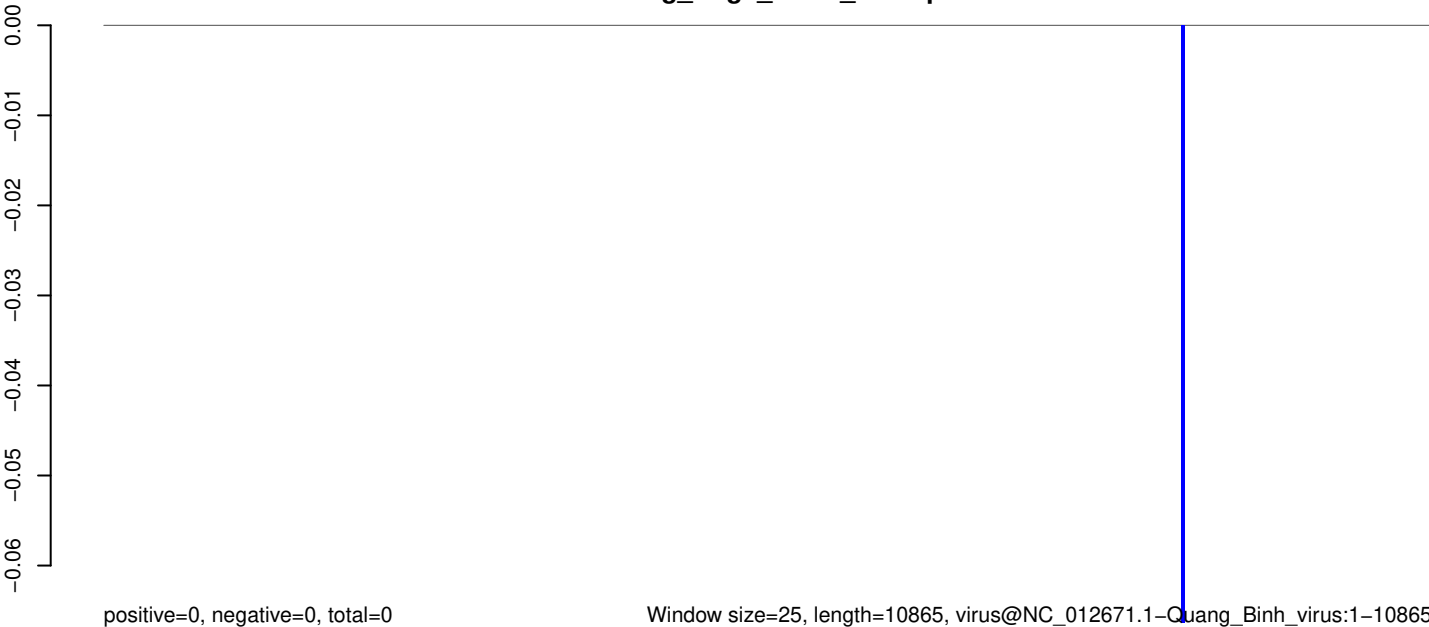
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



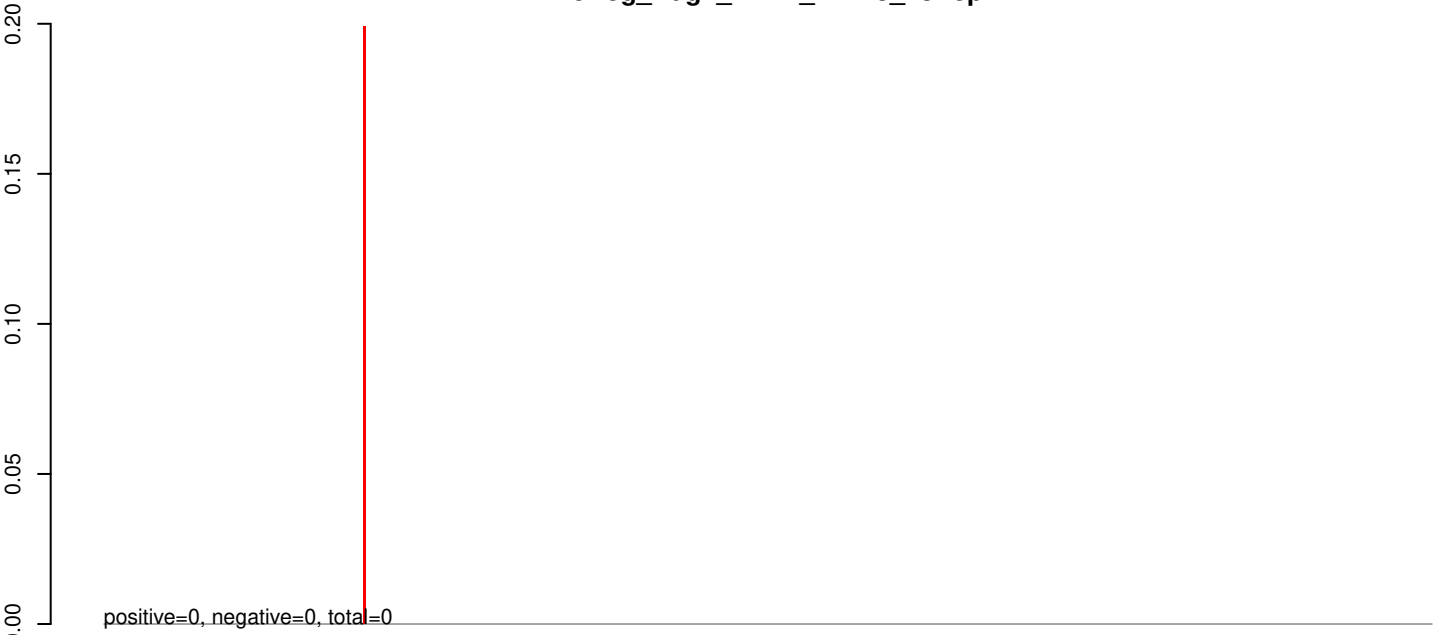
AeAeg_Aag2_BTV2_AK.rep



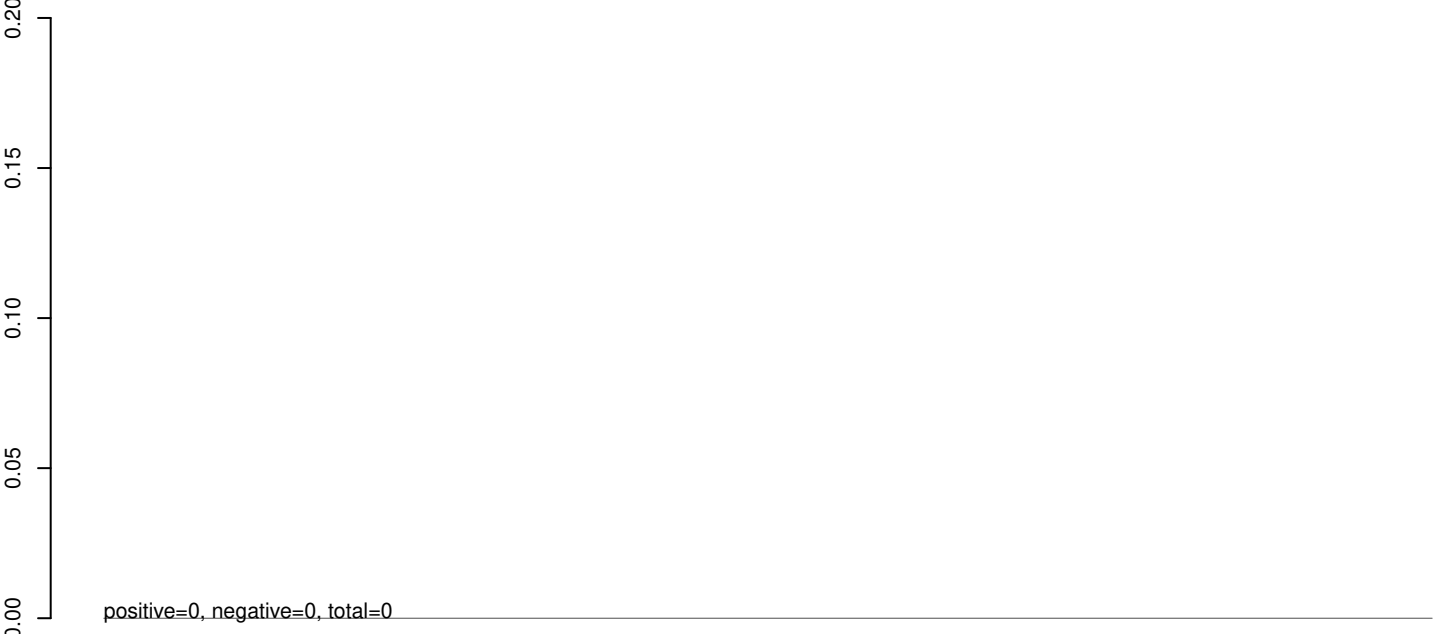
Window size=25, length=10865, virus@NC_012671.1-Quang_Binh_virus:1-10865

0 2000 4000 6000 8000 10000

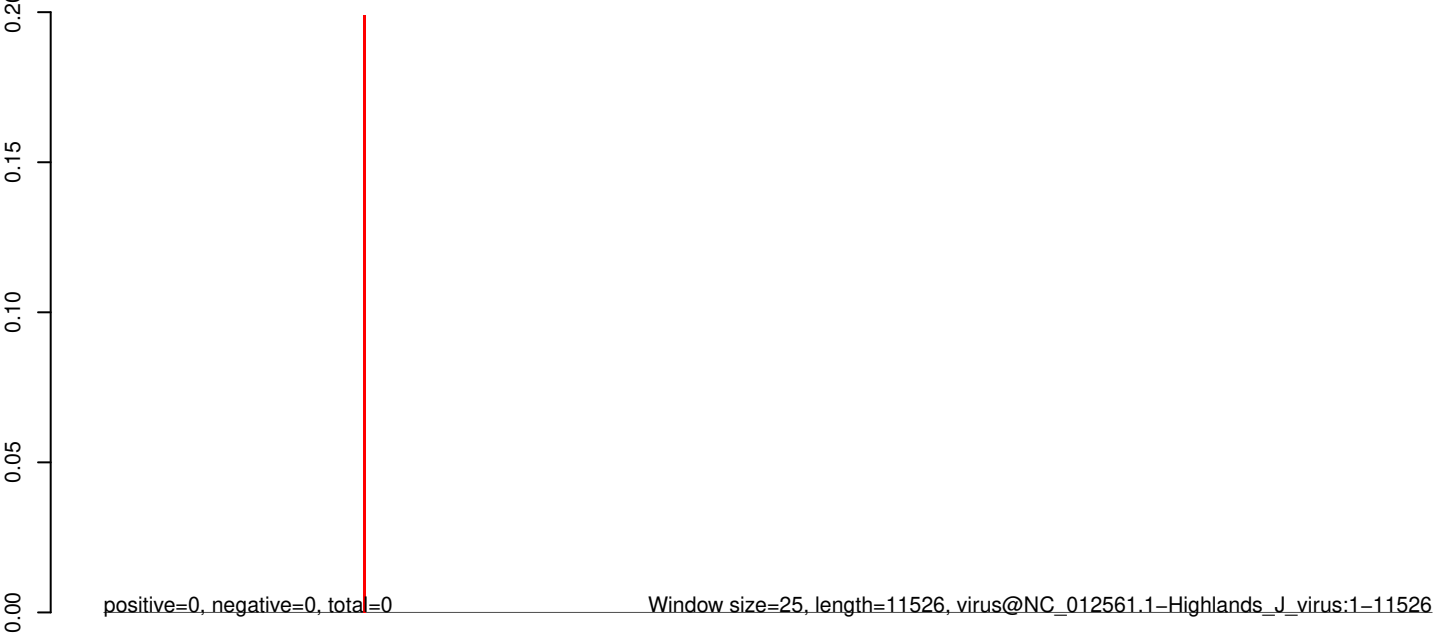
AeAeg_Aag2_BTV2_AK.18_23.rep



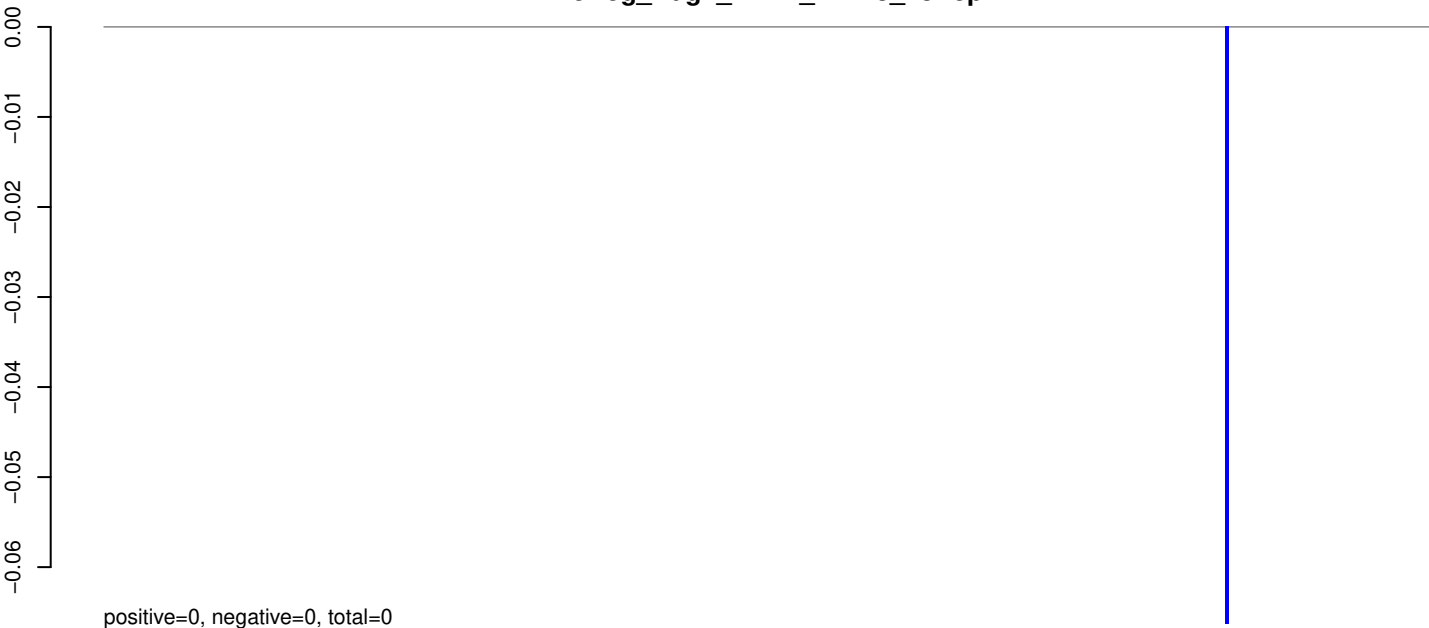
AeAeg_Aag2_BTV2_AK.24_35.rep



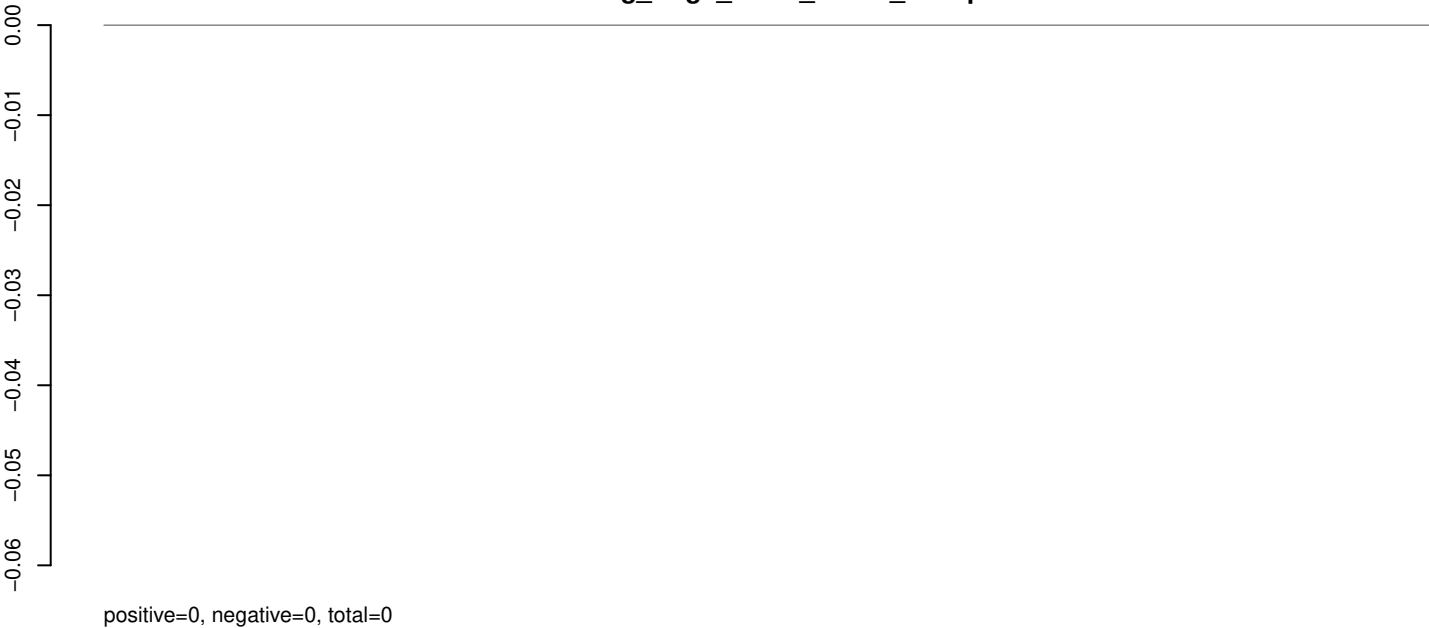
AeAeg_Aag2_BTV2_AK.rep



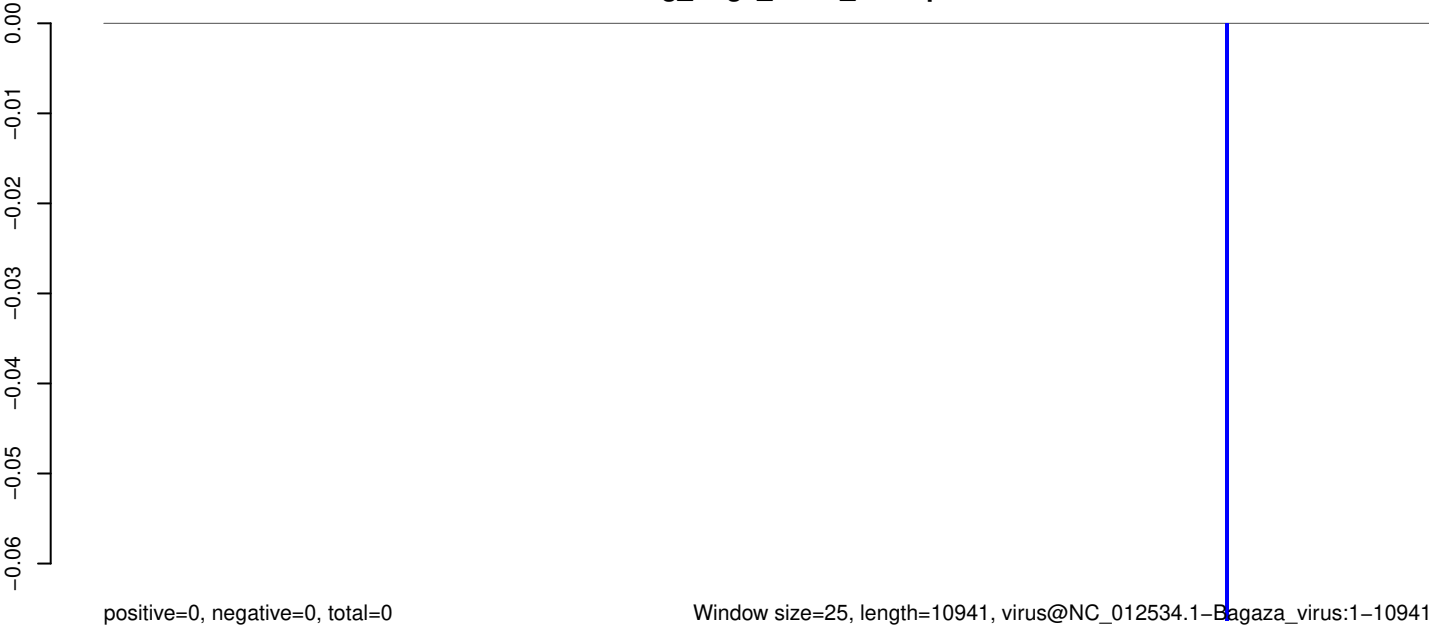
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



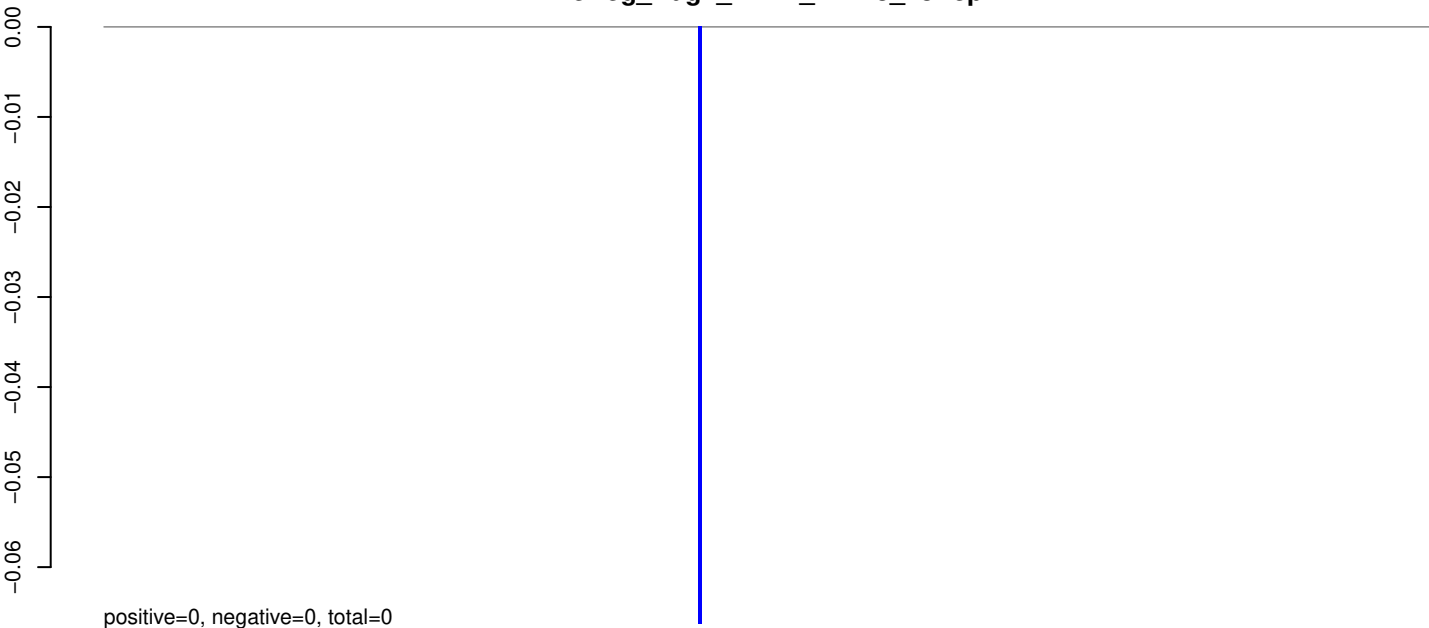
AeAeg_Aag2_BTV2_AK.rep



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

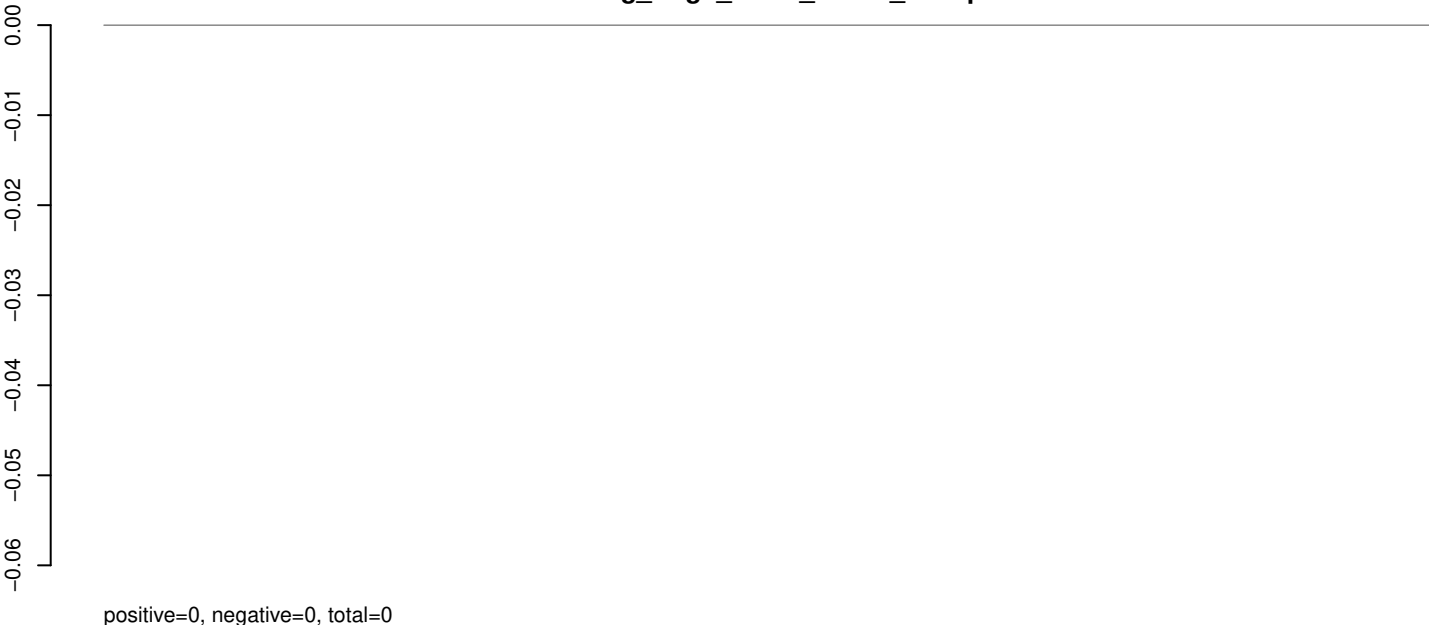
0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BTV2_AK.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BTV2_AK.rep

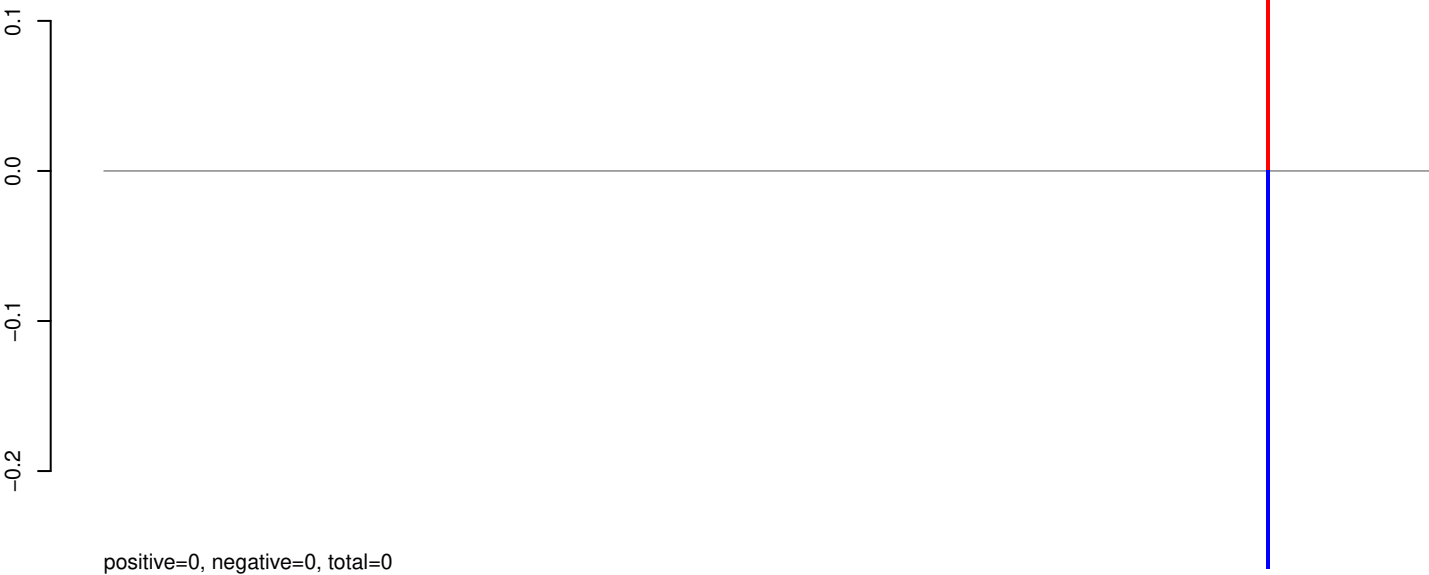


positive=0, negative=0, total=0

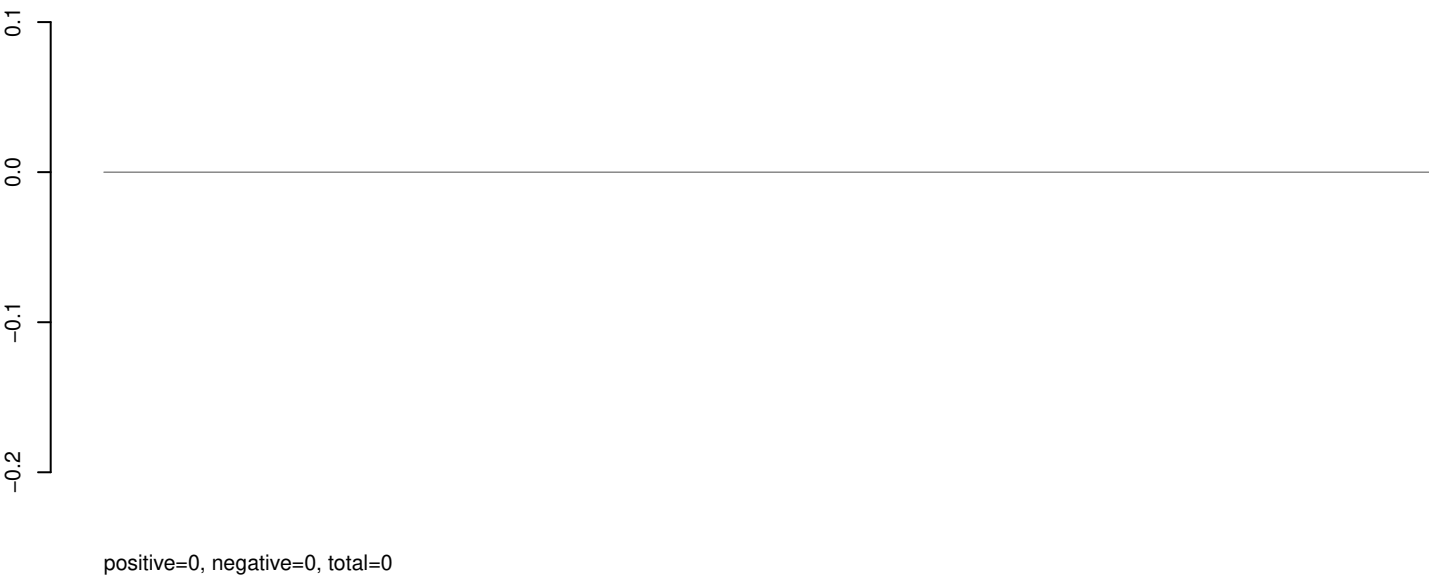
Window size=25, length=10723, virus@NC_012533.1-Kedougou_virus:1-10723

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



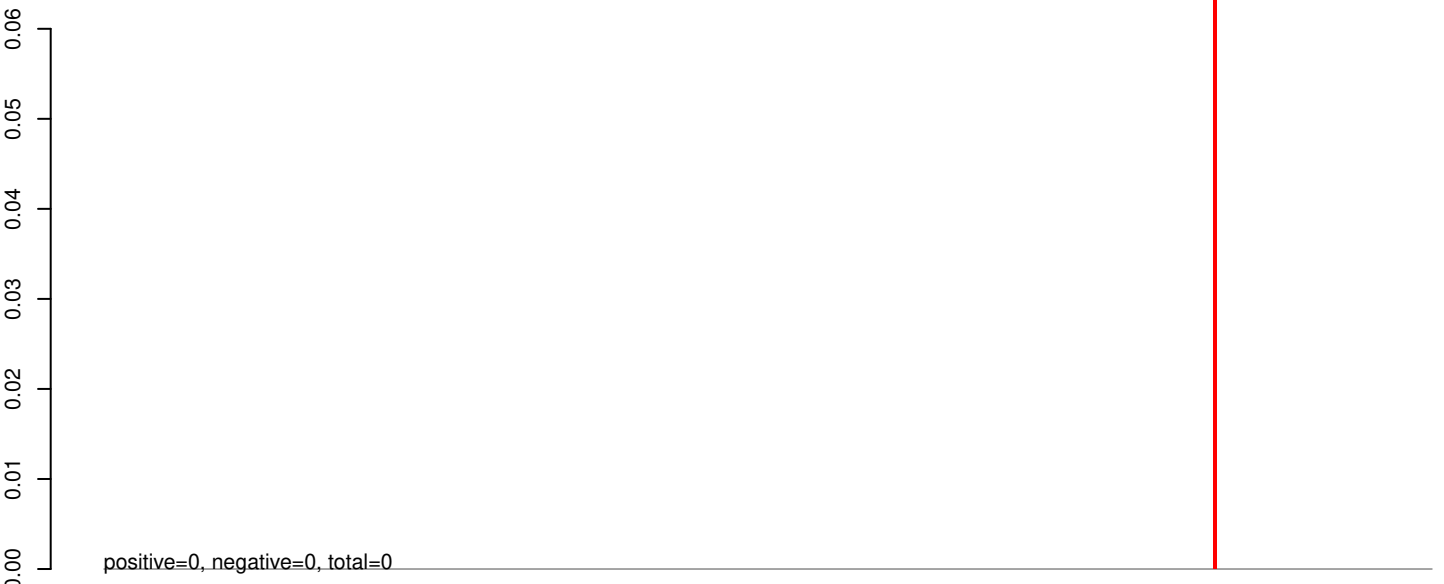
AeAeg_Aag2_BTV2_AK.rep



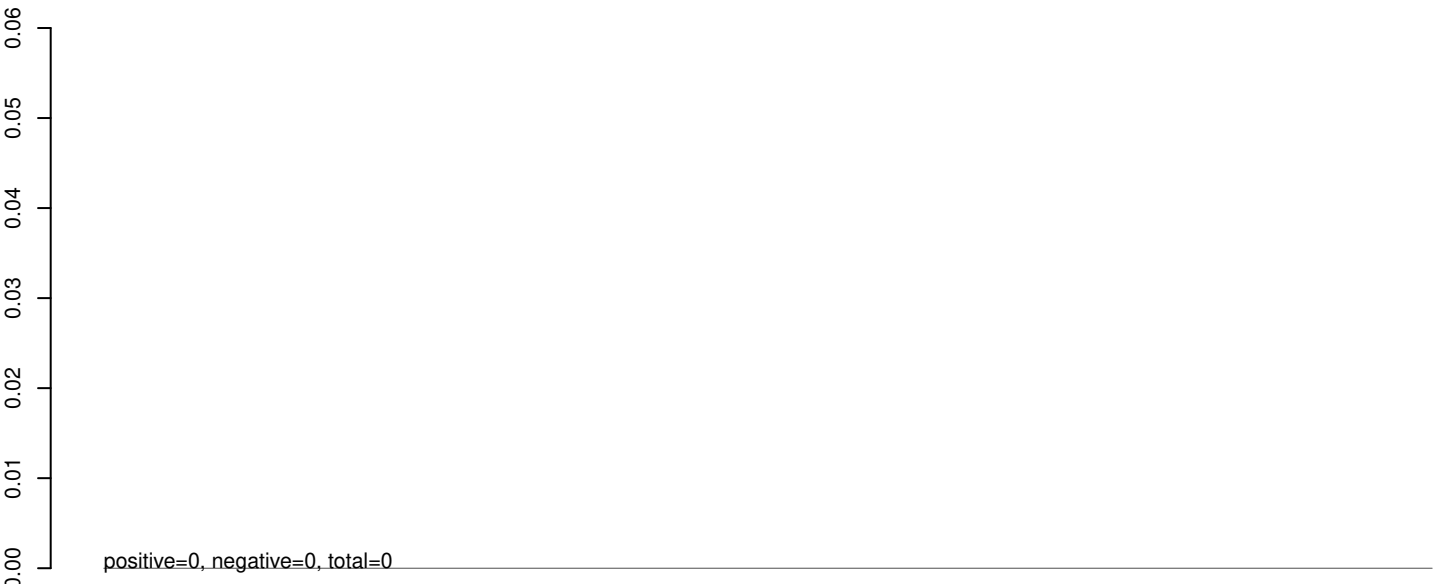
Window size=25, length=10815, virus@NC_009026.2-Bussuquara_virus:1-10815

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



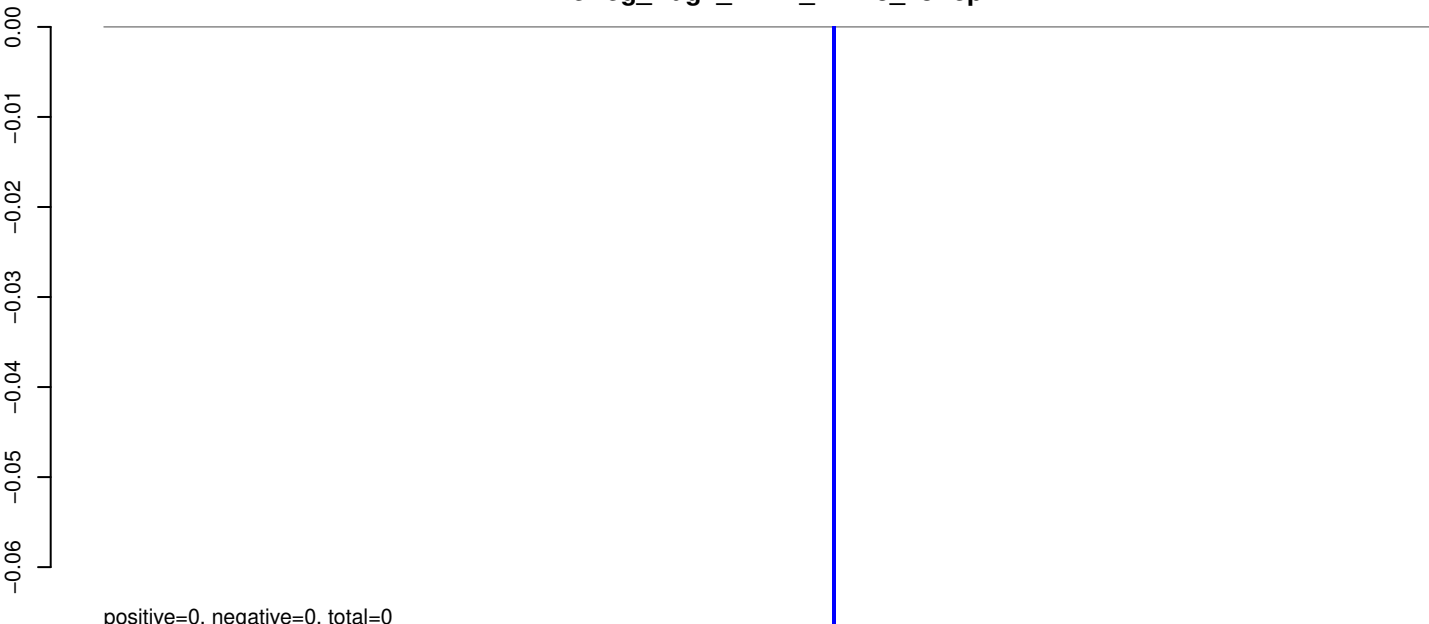
AeAeg_Aag2_BTV2_AK.24_35.rep



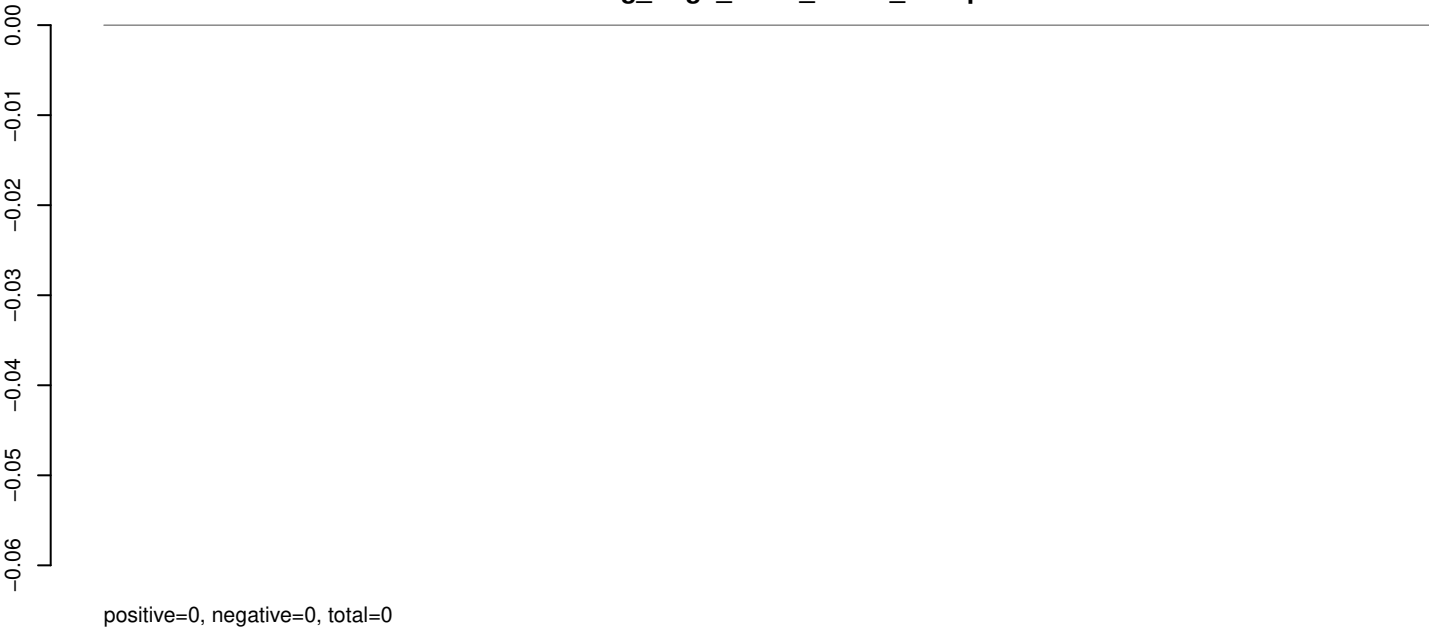
AeAeg_Aag2_BTV2_AK.rep



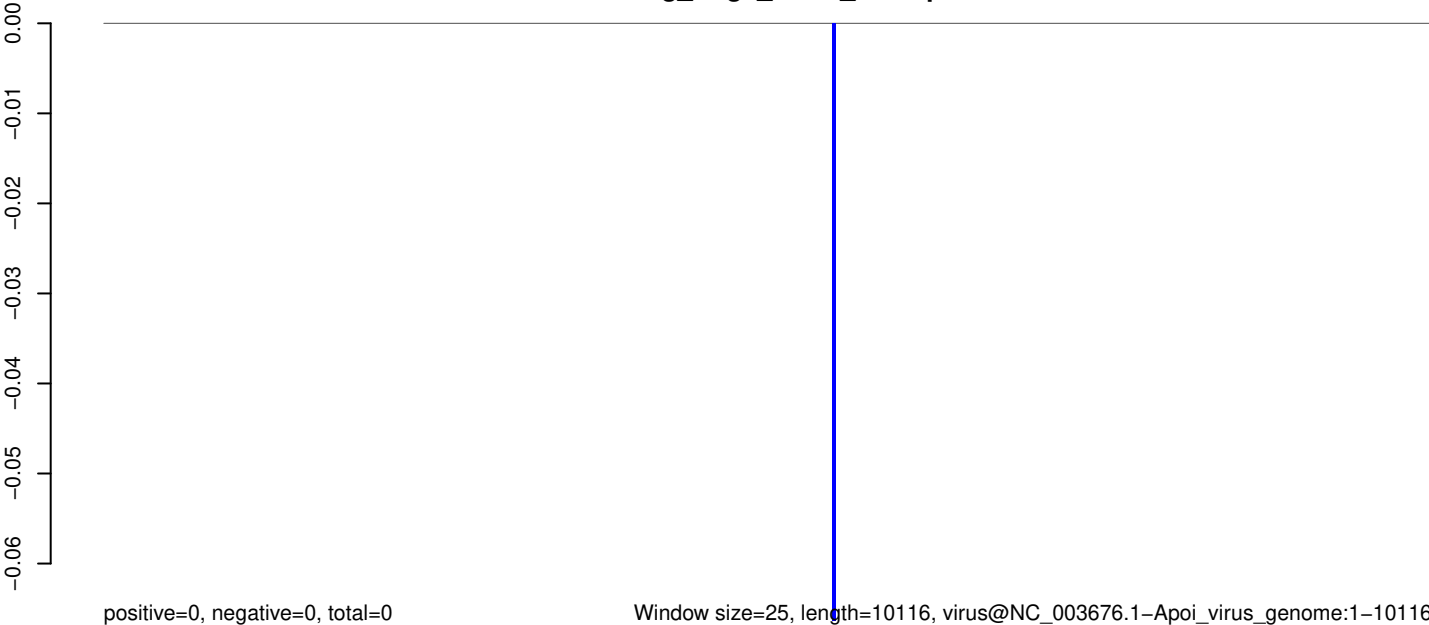
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



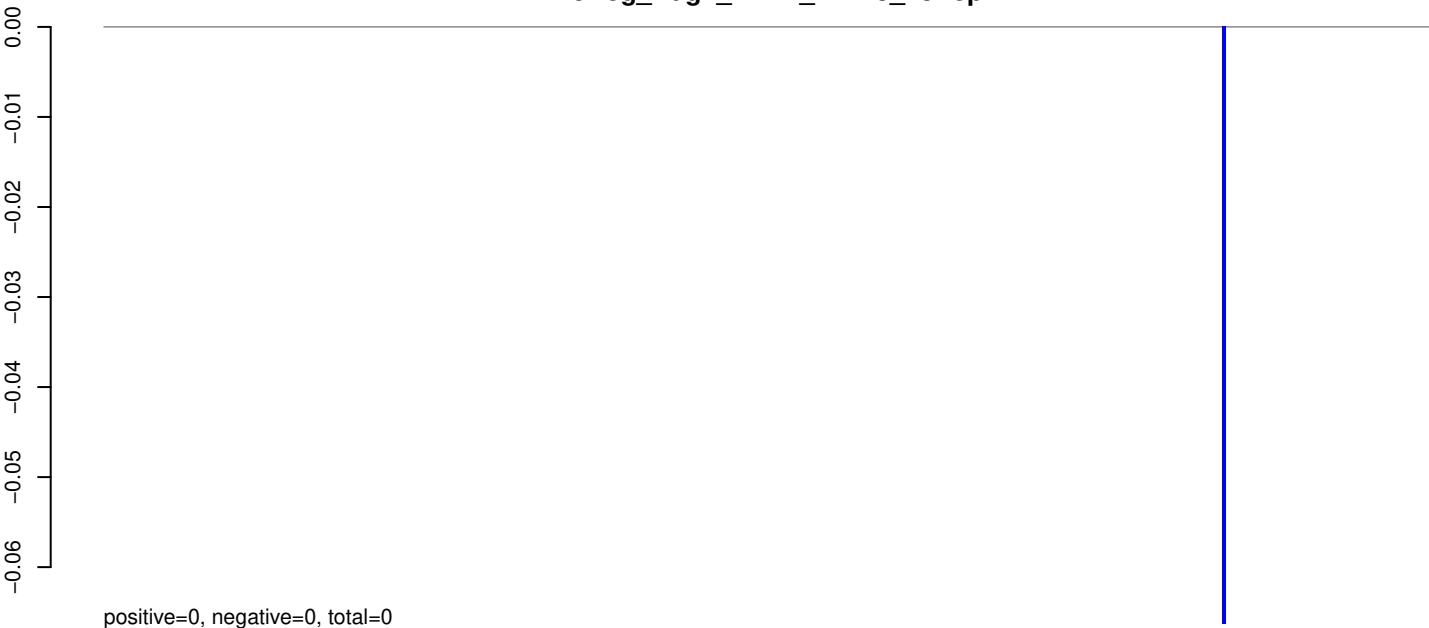
AeAeg_Aag2_BTV2_AK.rep



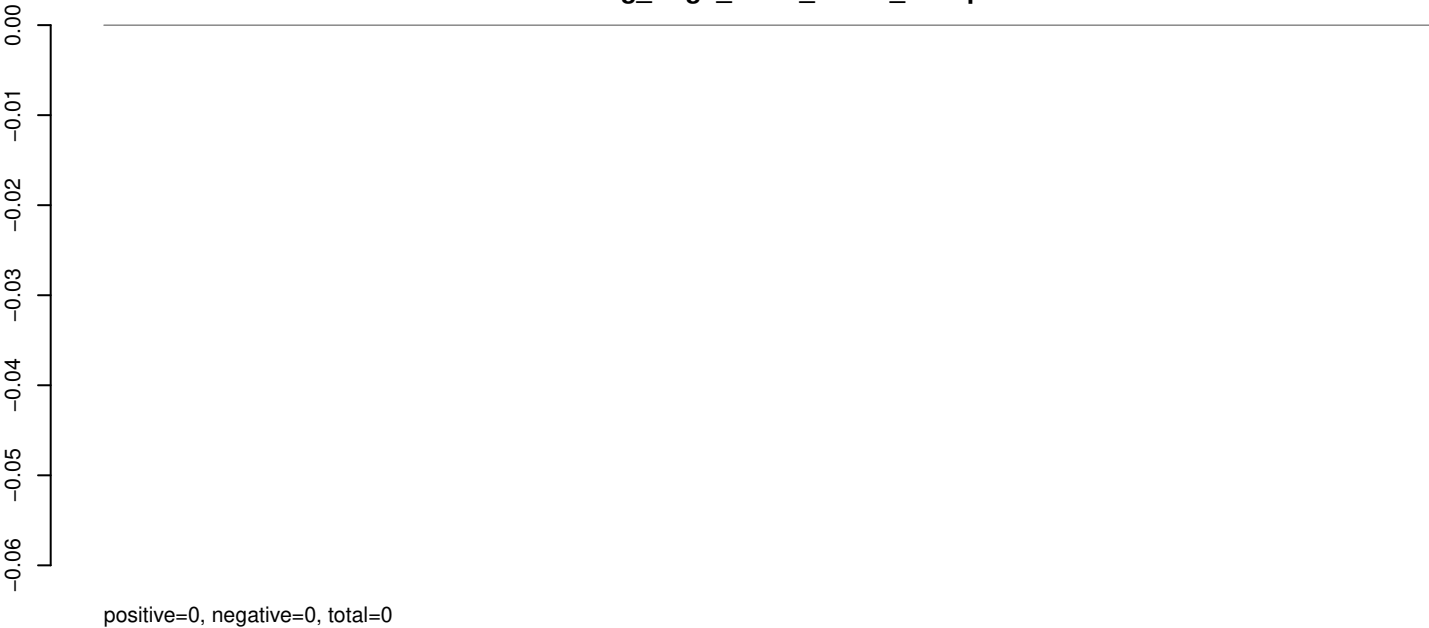
Window size=25, length=10116, virus@NC_003676.1-Apoi_virus_genome:1-10116

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



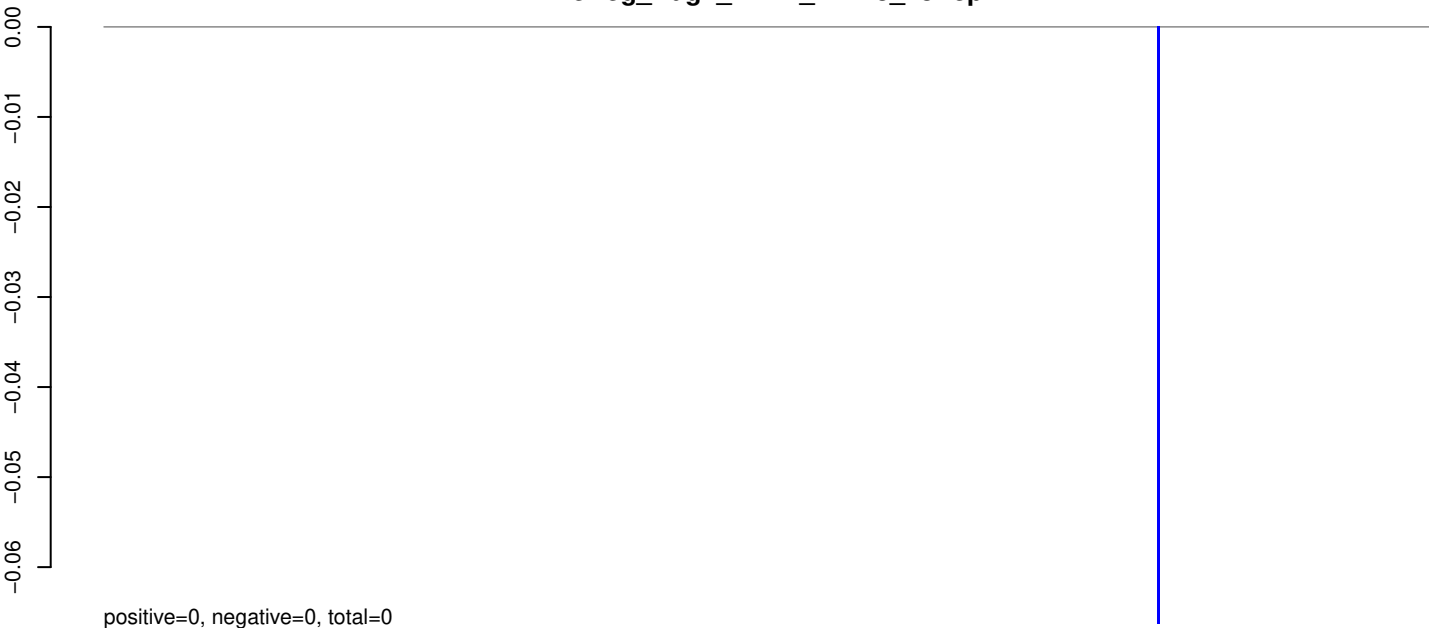
AeAeg_Aag2_BTV2_AK.rep



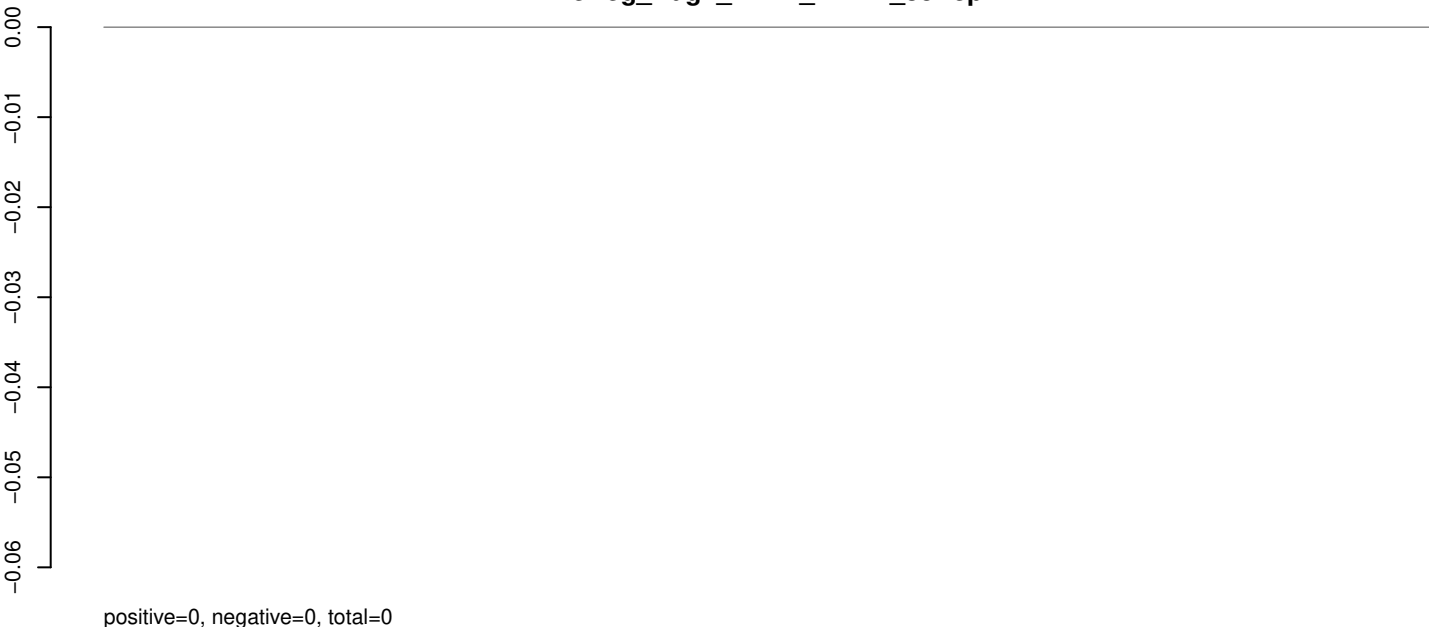
Window size=25, length=10140, virus@NC_003675.1-Rio_Bravo_virus_genome:1-10140



AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



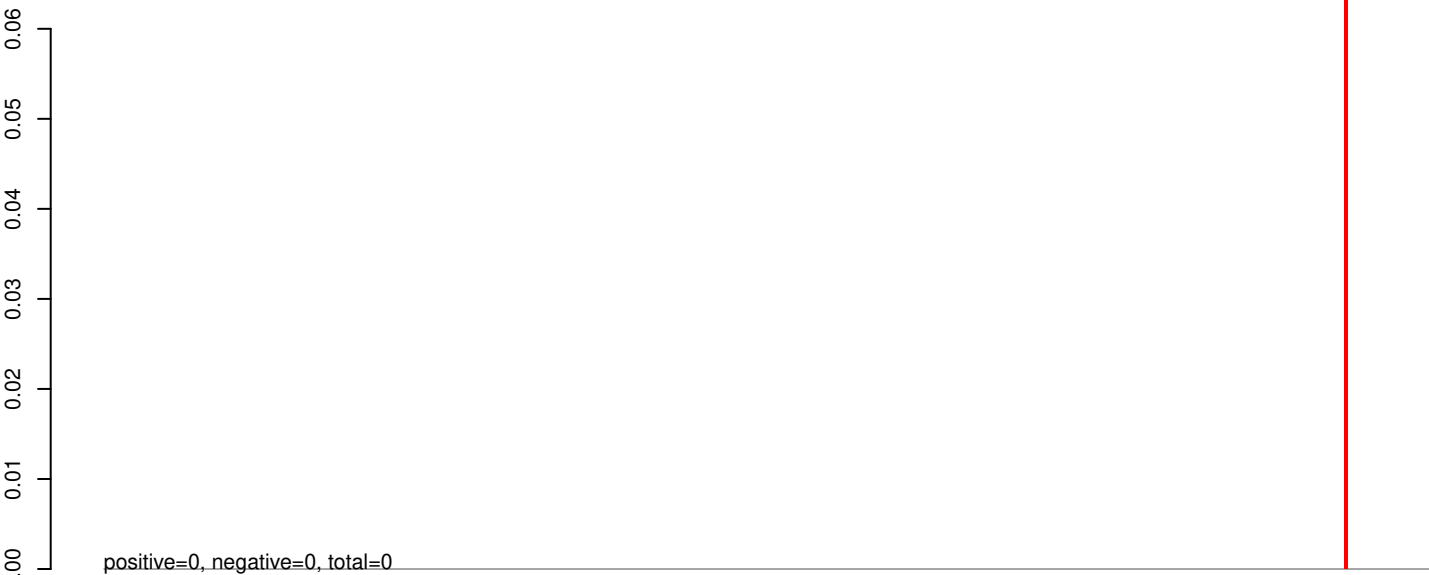
AeAeg_Aag2_BTV2_AK.rep



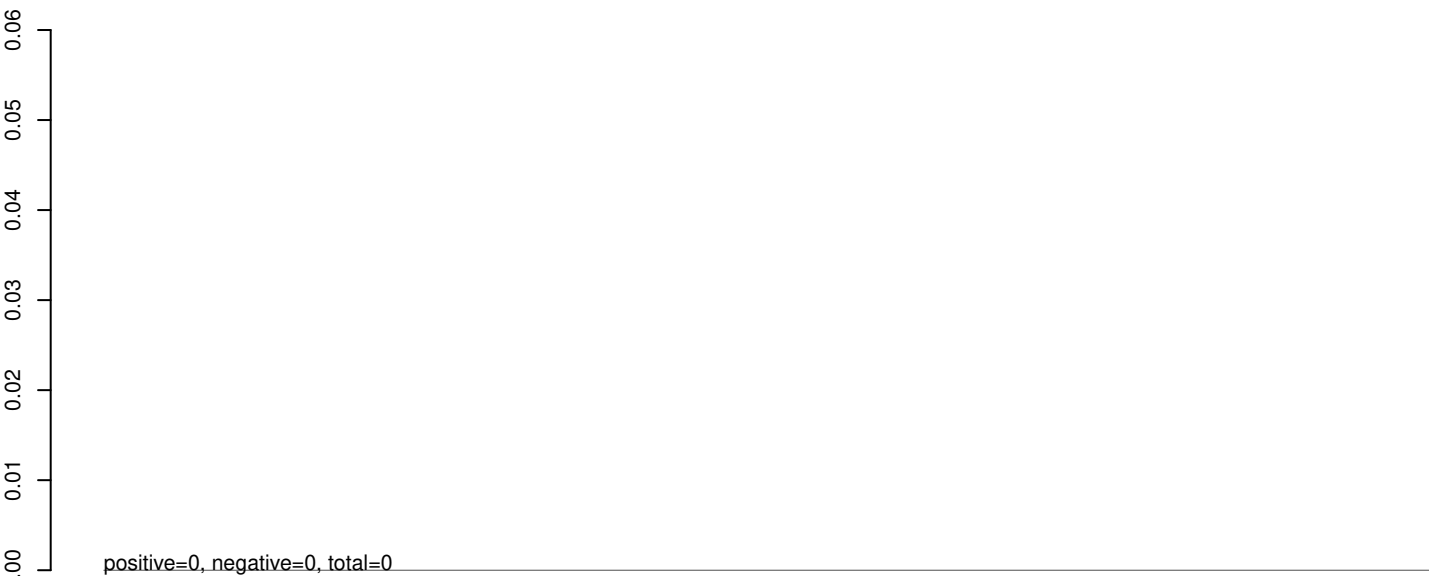
Window size=25, length=11442, virus@NC_003215.1-Semliki_forest_virus:1-11442

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

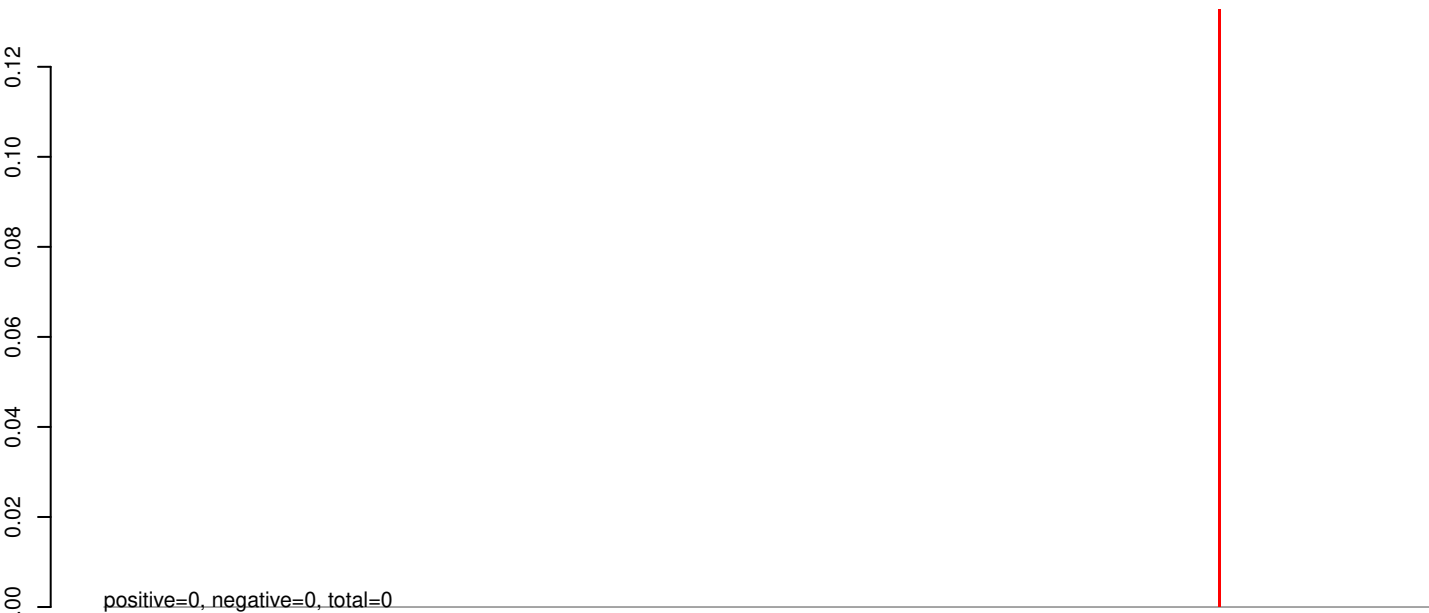


AeAeg_Aag2_BTV2_AK.rep

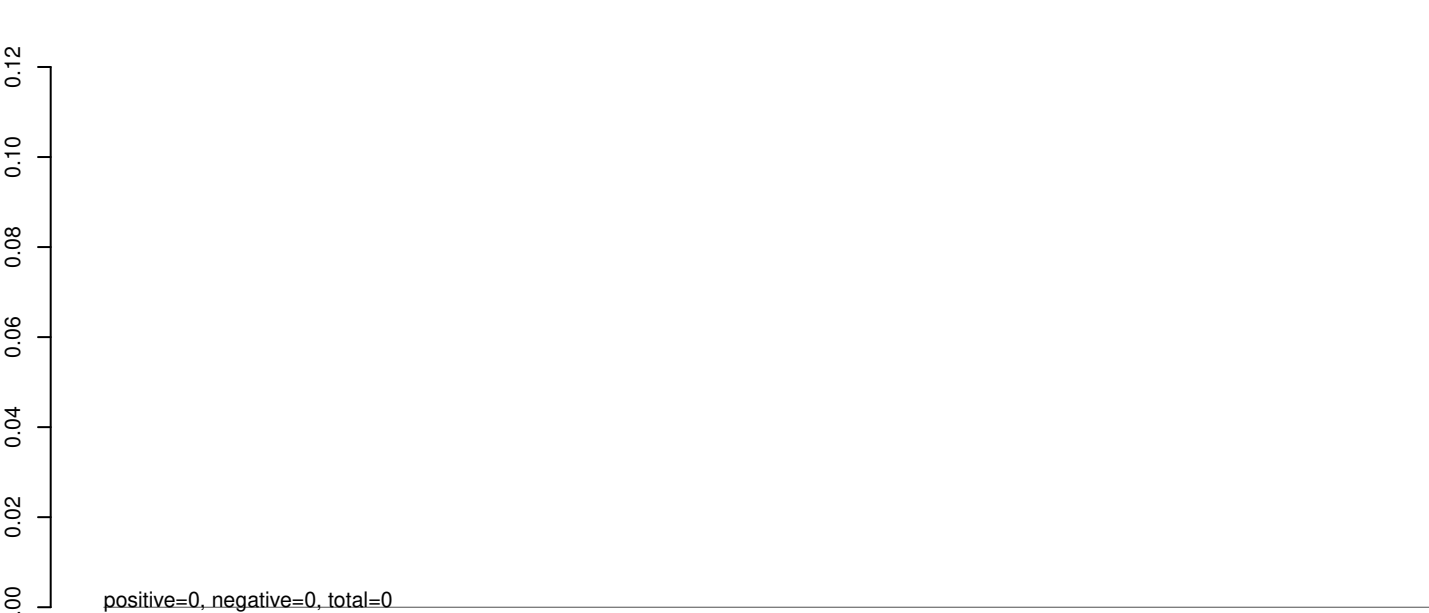


0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

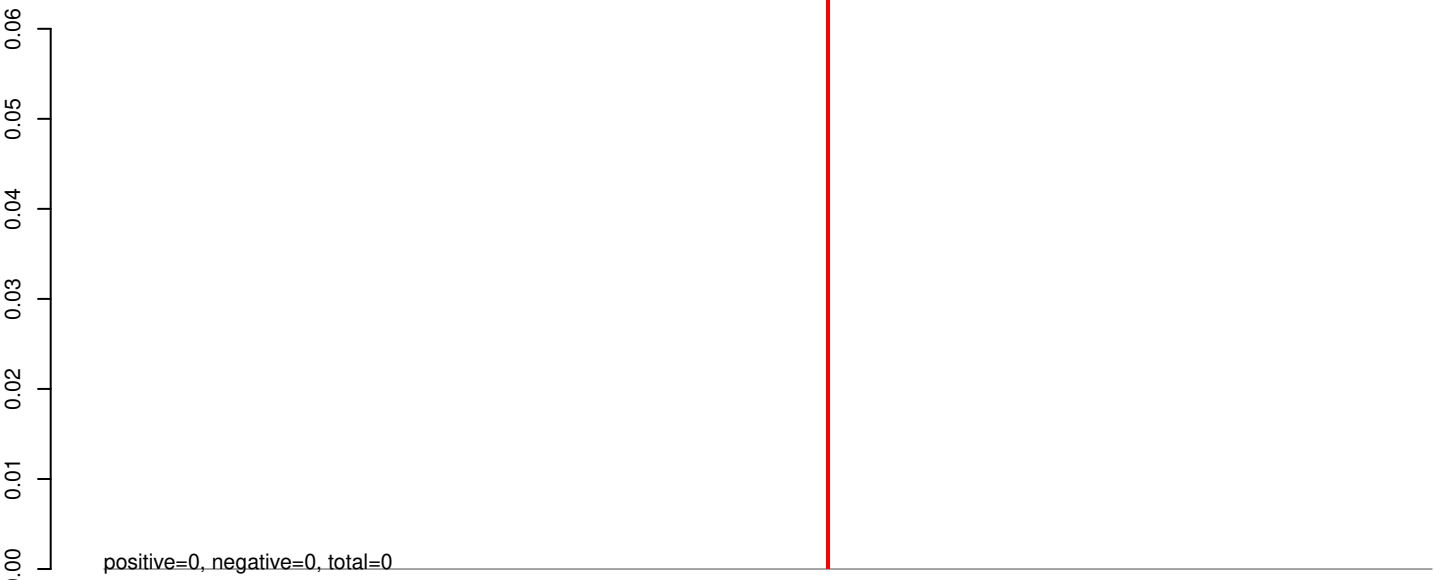


AeAeg_Aag2_BTV2_AK.rep

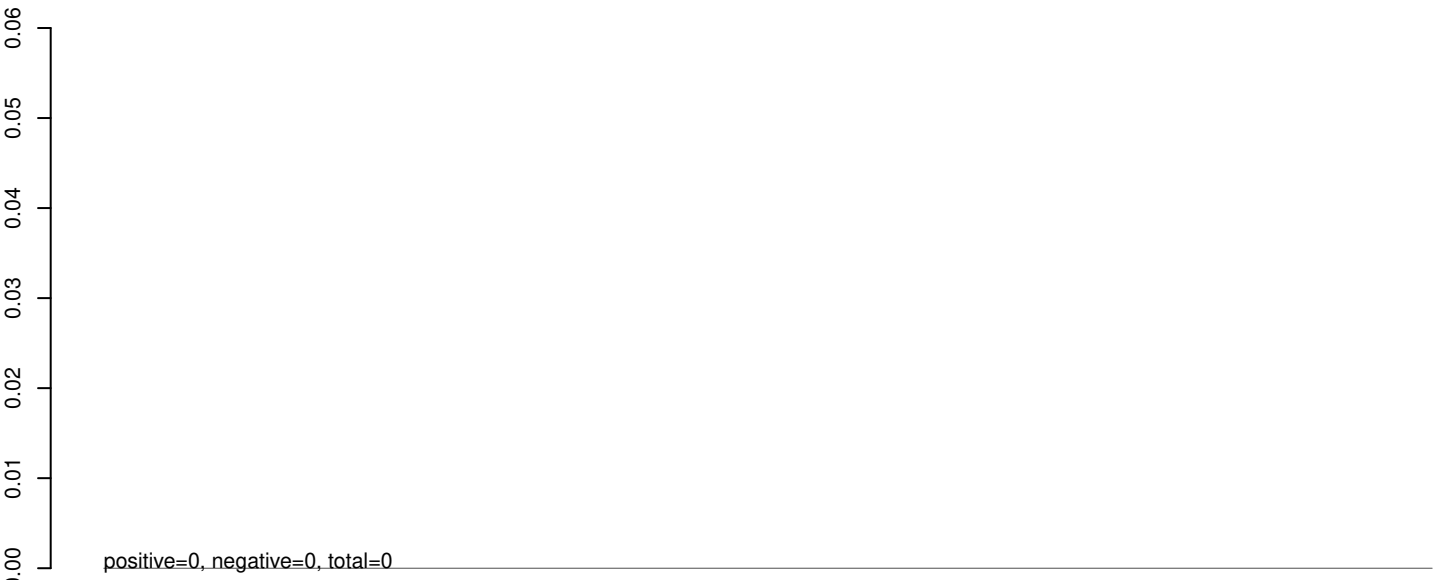


0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

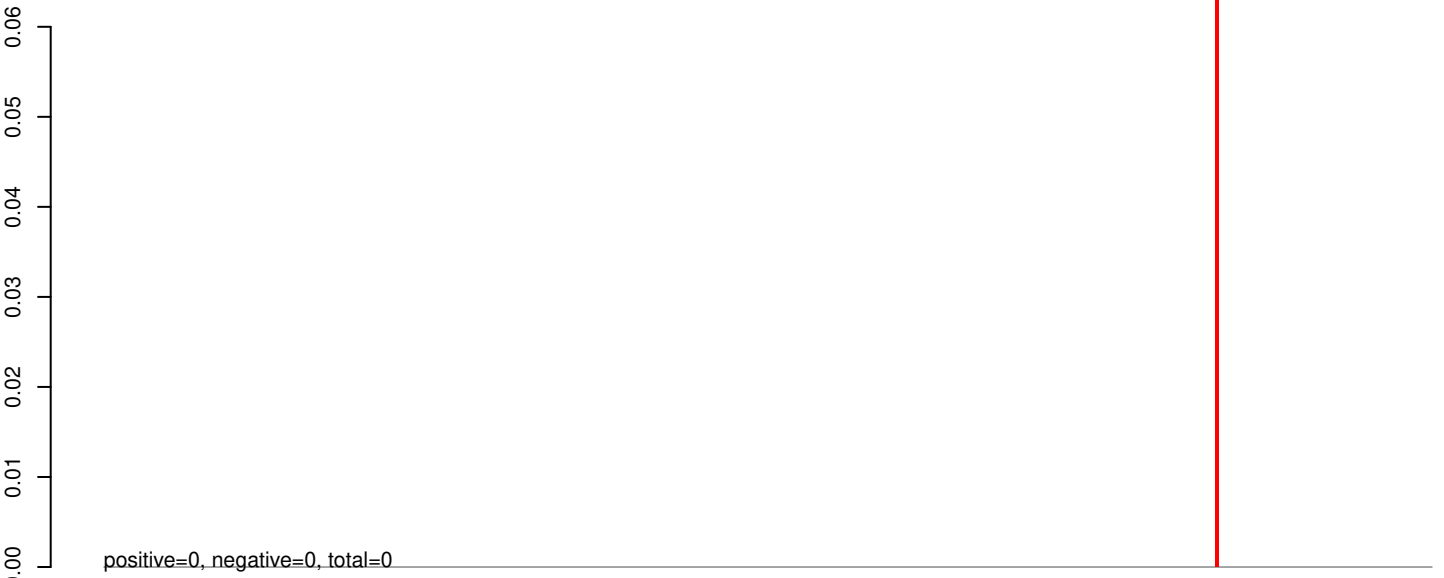


AeAeg_Aag2_BTV2_AK.rep

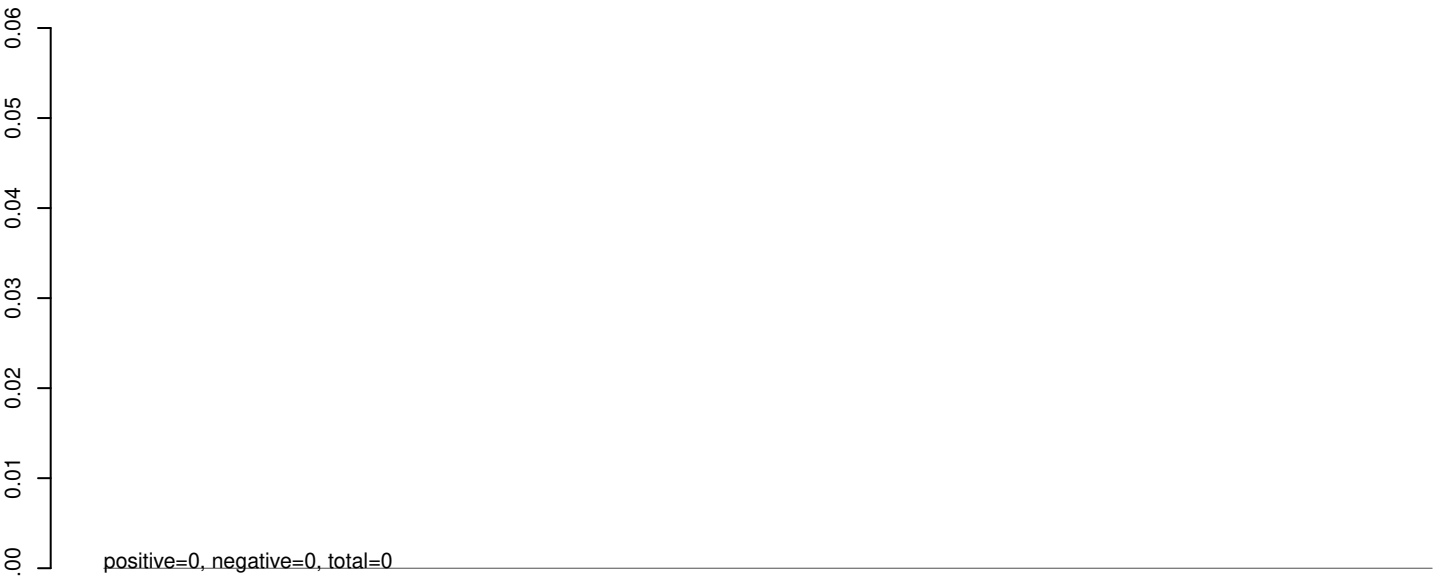


0 2000 4000 6000 8000 10000

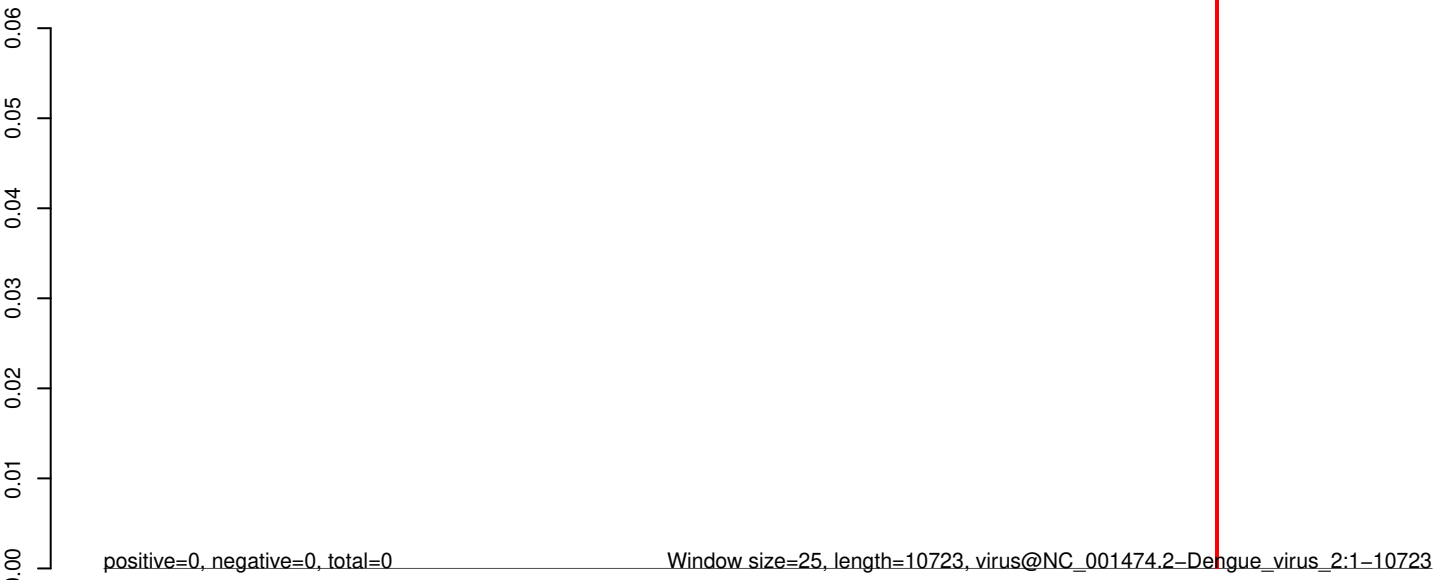
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

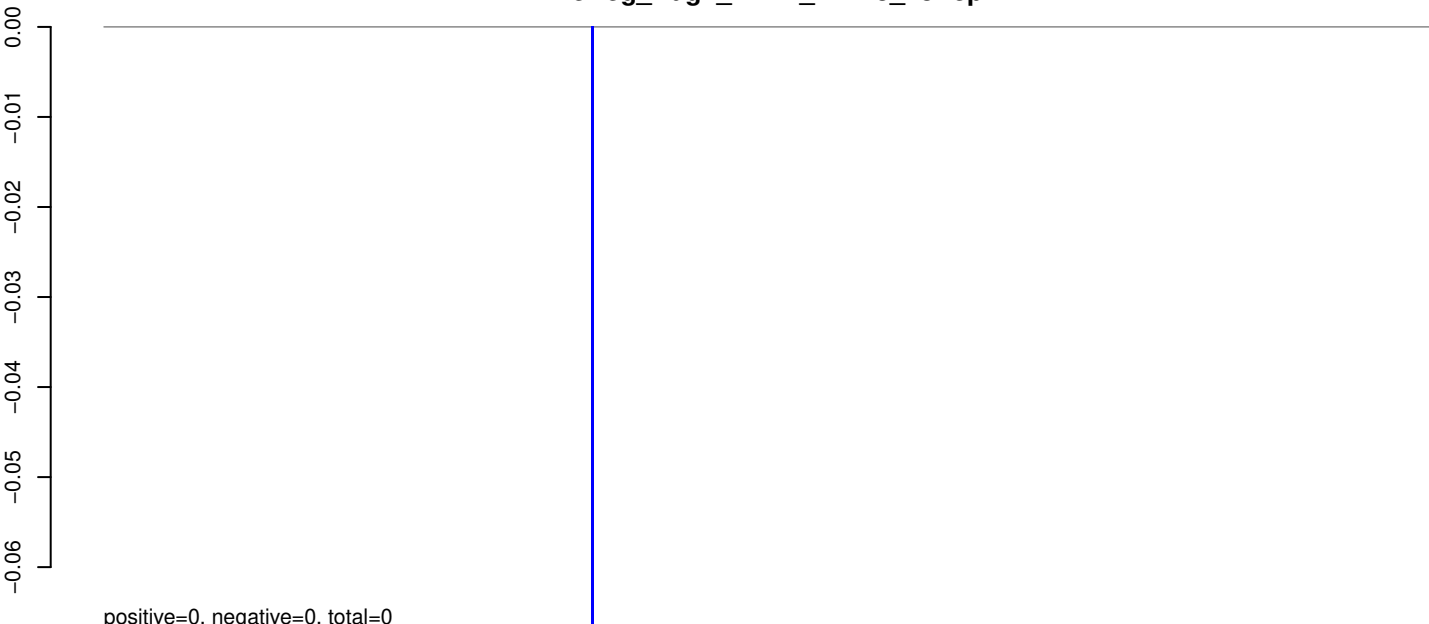


AeAeg_Aag2_BTV2_AK.rep

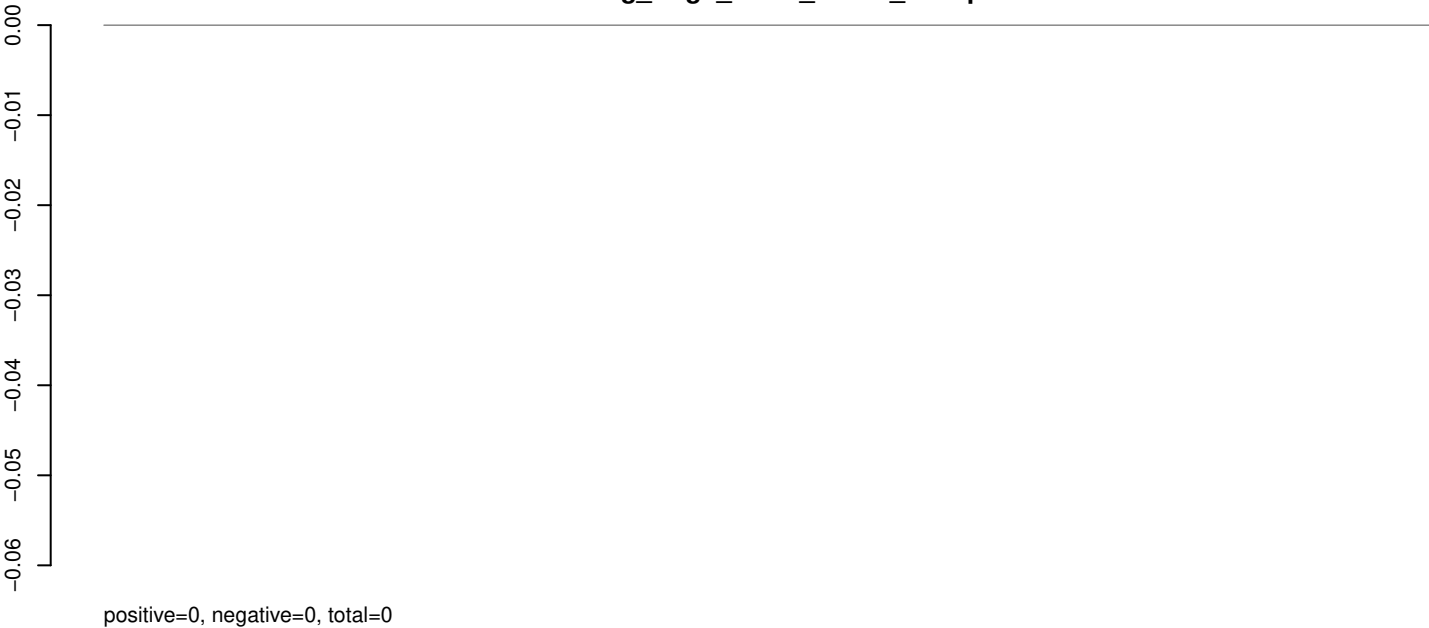


0 2000 4000 6000 8000 10000

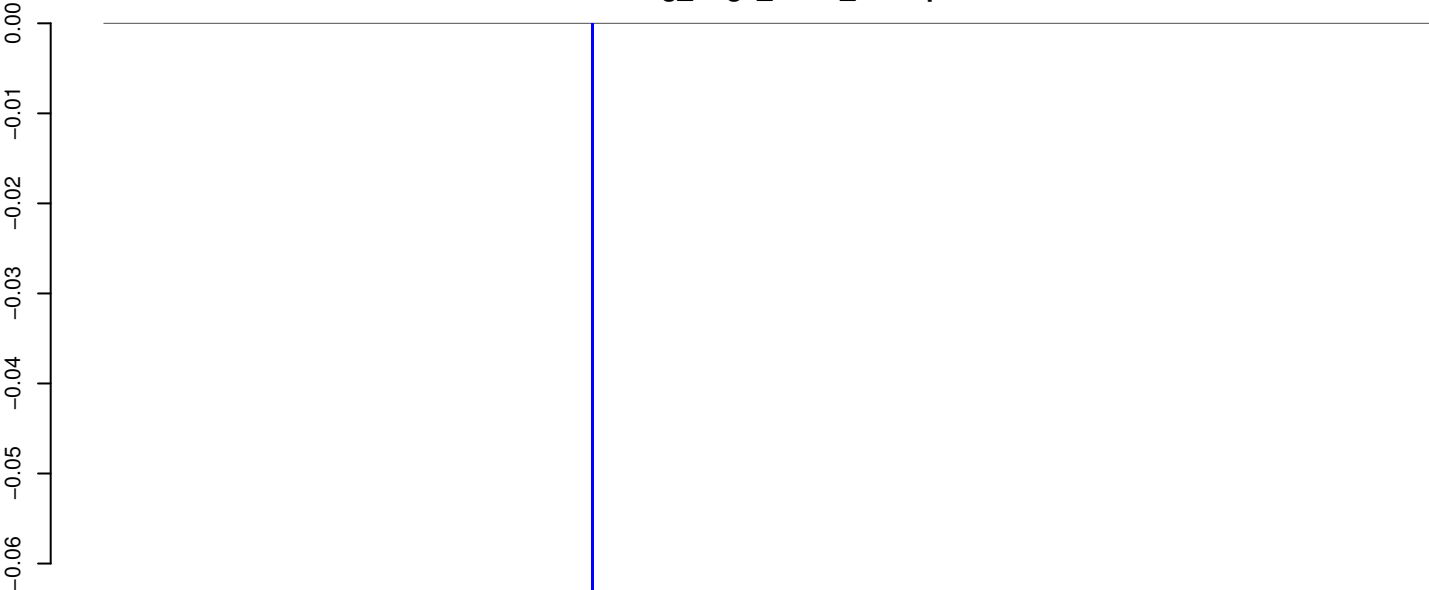
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



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

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep



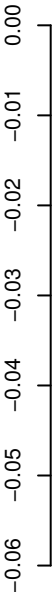
positive=0, negative=0, total=0

AeAeg_Aag2_BTV2_AK.24_35.rep



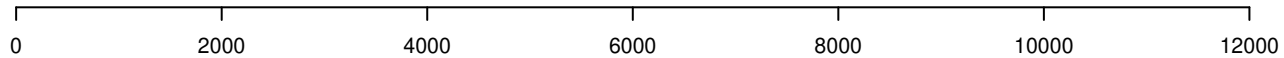
positive=0, negative=0, total=0

AeAeg_Aag2_BTV2_AK.rep

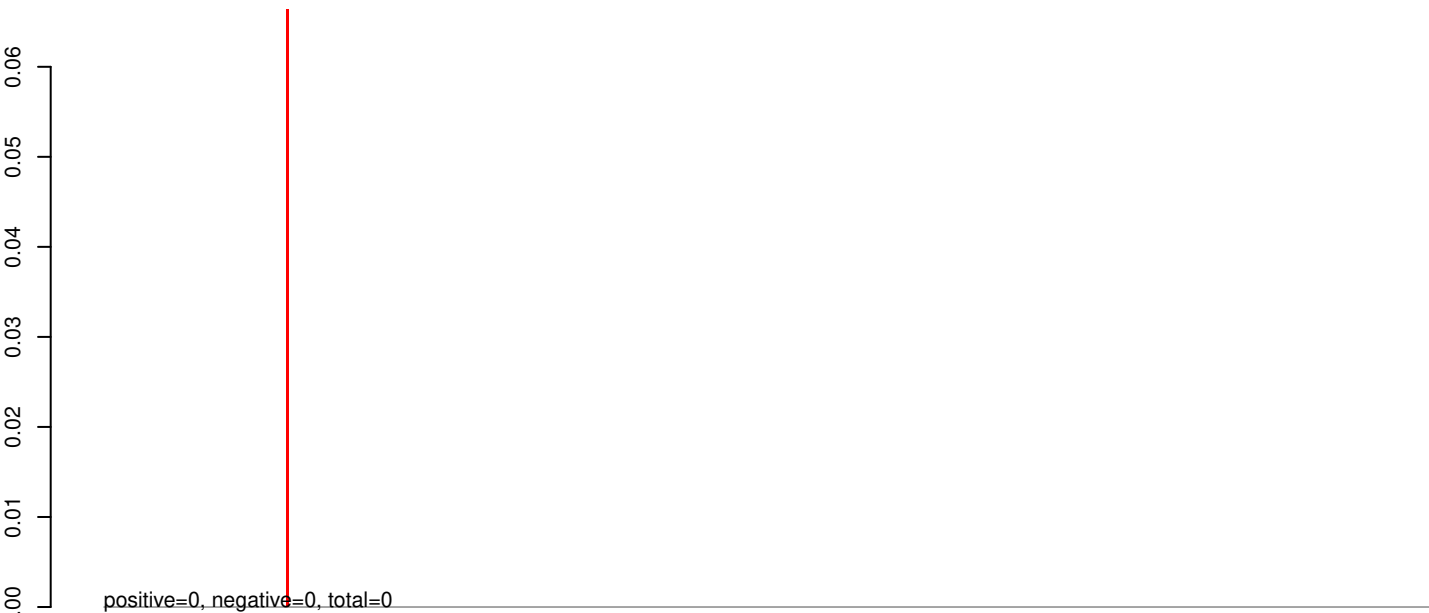


positive=0, negative=0, total=0

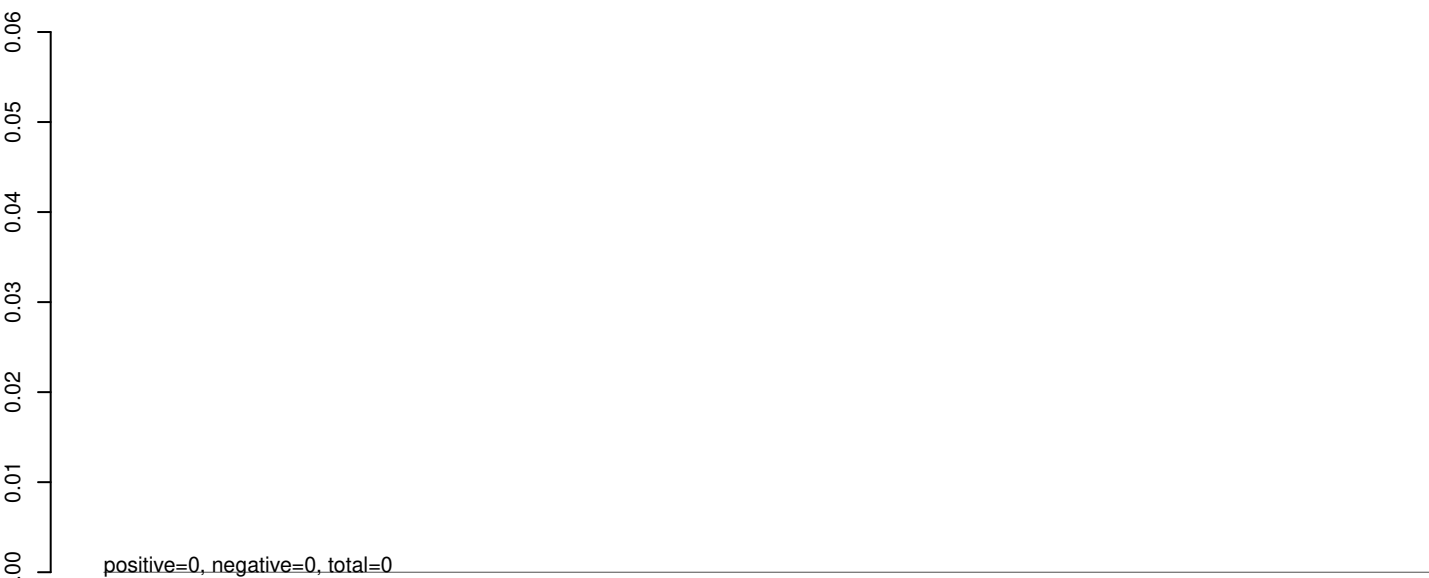
Window size=25, length=12916, virus@MH237595.1-Aedes_aegypti_anphevirus_strain_1a:1-12916



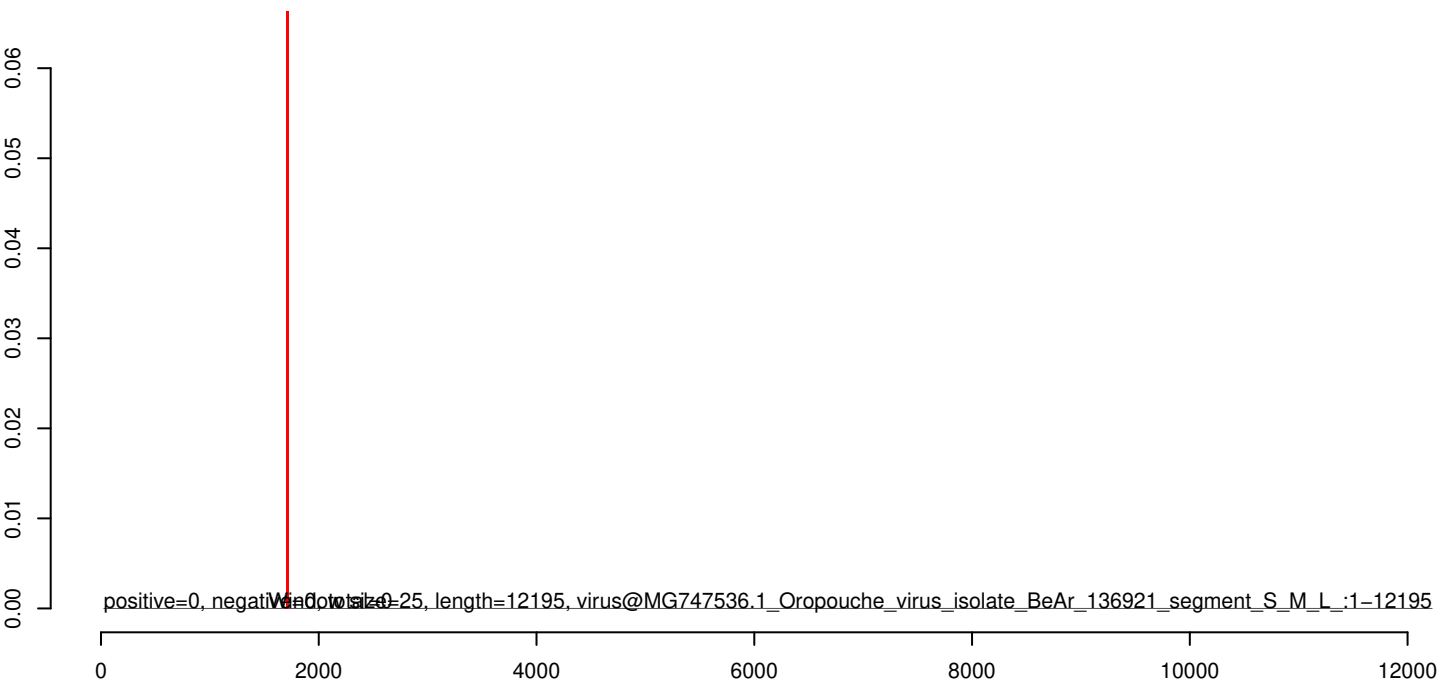
AeAeg_Aag2_BTV2_AK.18_23.rep



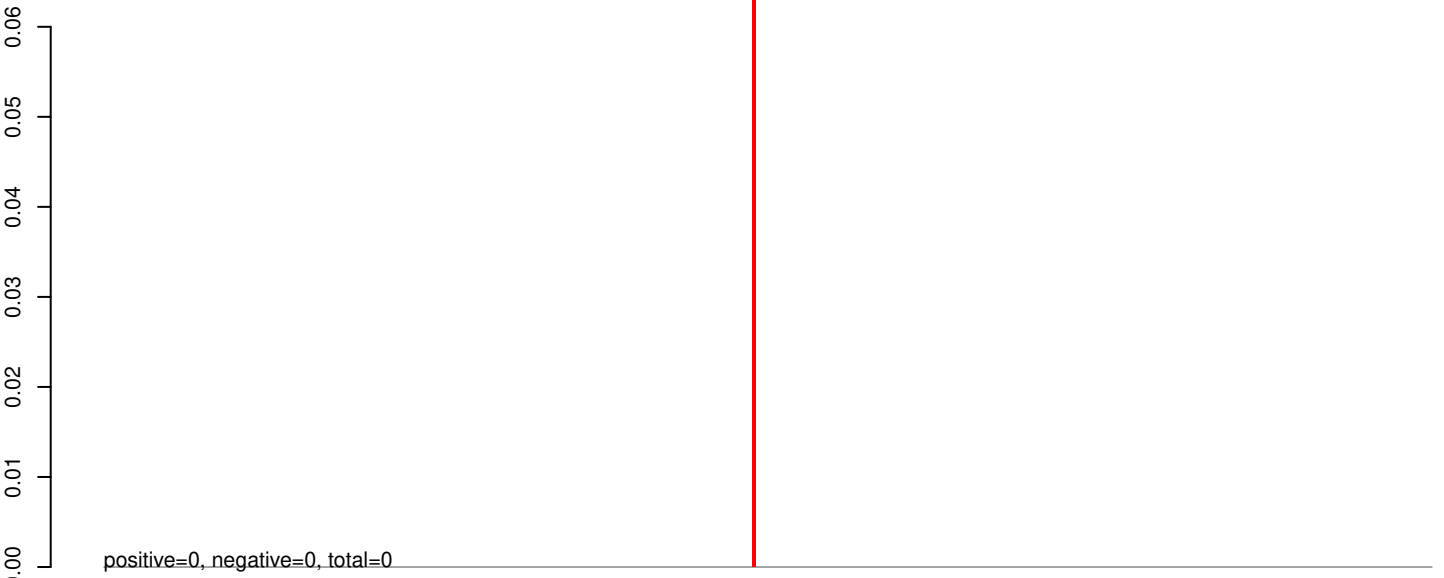
AeAeg_Aag2_BTV2_AK.24_35.rep



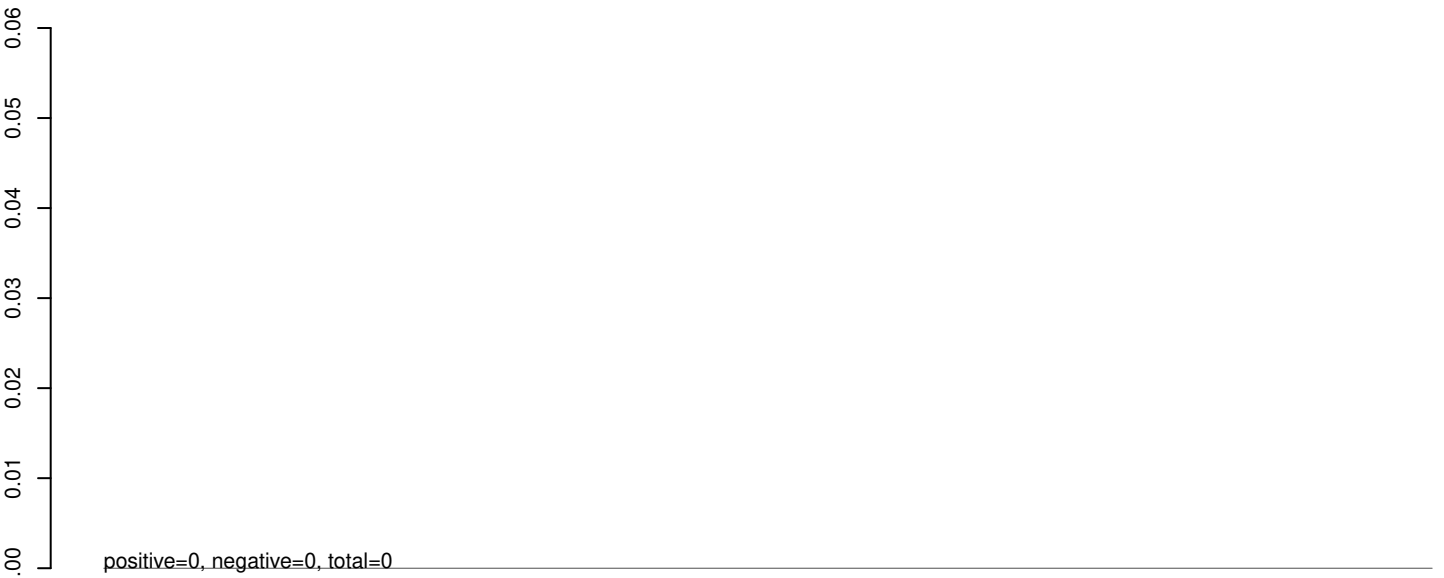
AeAeg_Aag2_BTV2_AK.rep



AeAeg_Aag2_BTV2_AK.18_23.rep



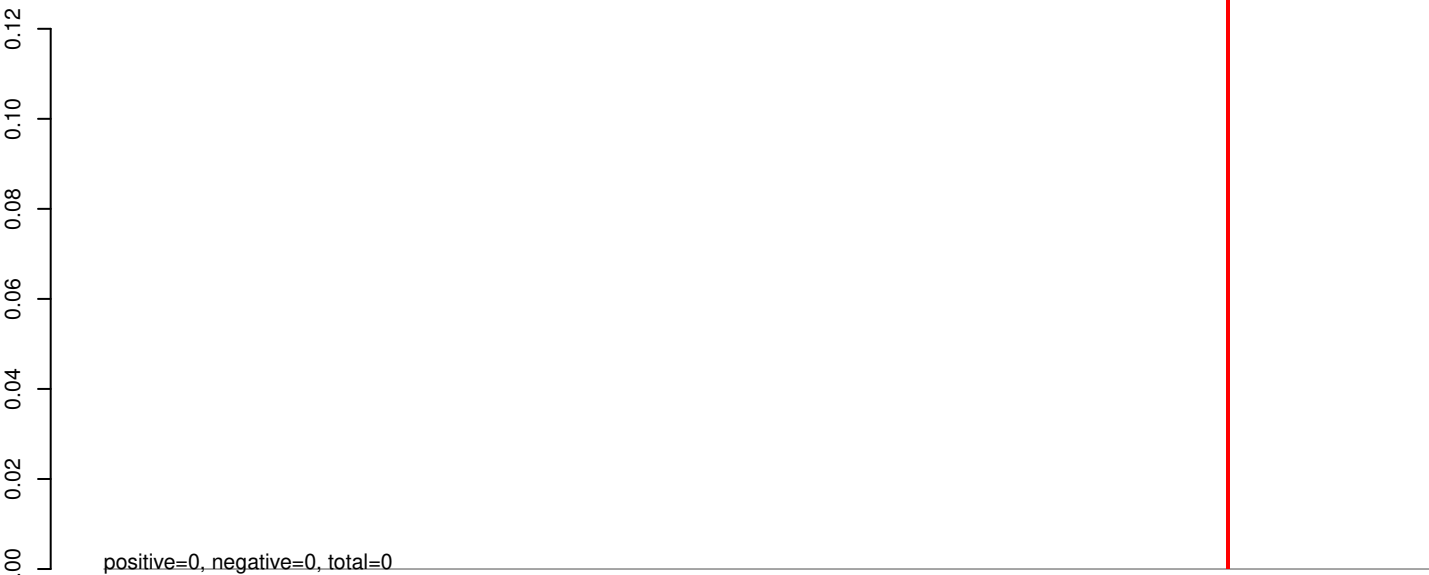
AeAeg_Aag2_BTV2_AK.24_35.rep



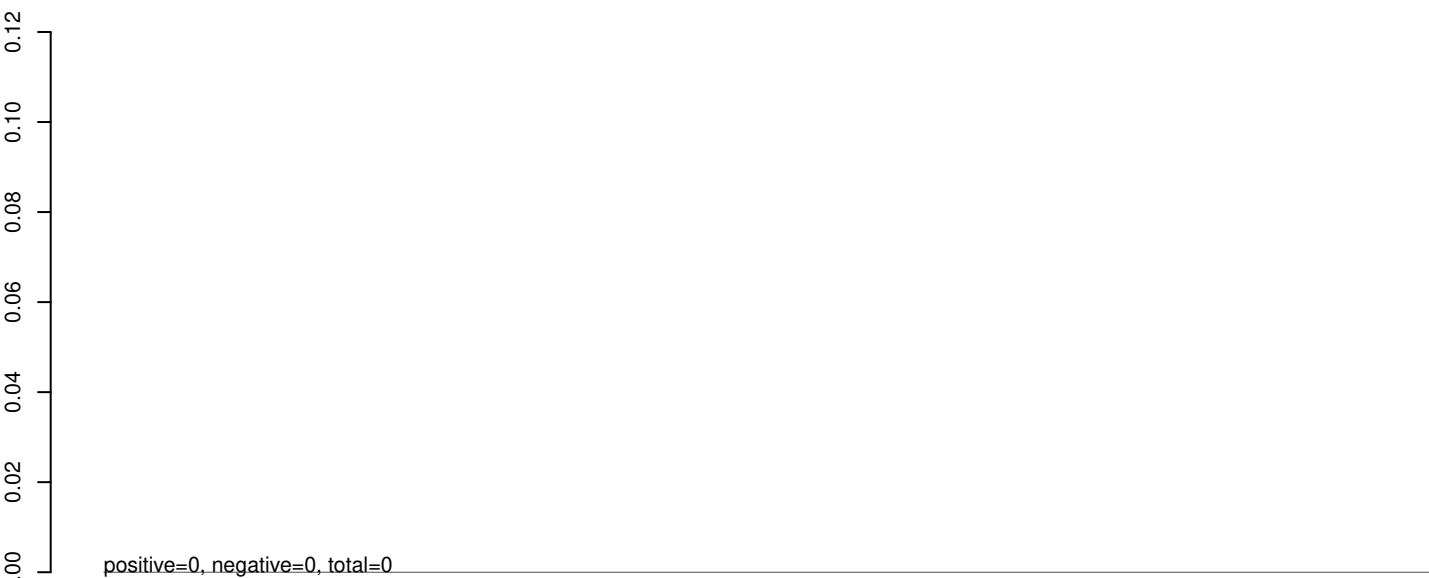
AeAeg_Aag2_BTV2_AK.rep



AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep

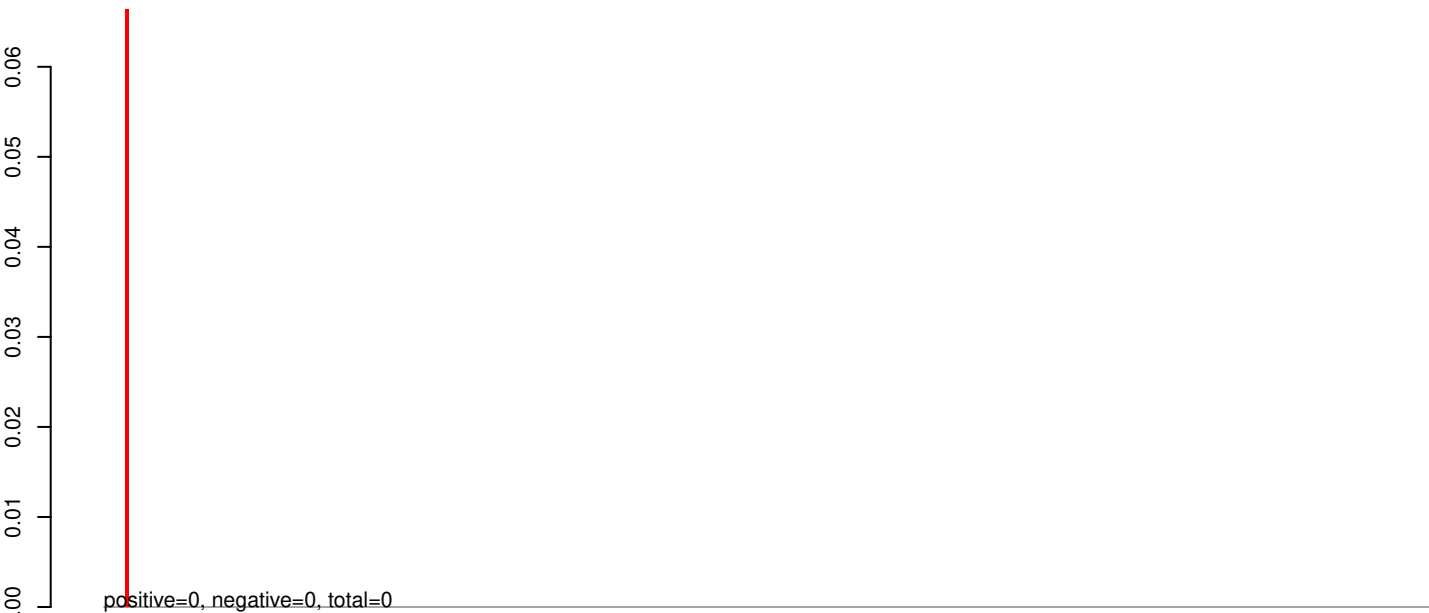


AeAeg_Aag2_BTV2_AK.rep

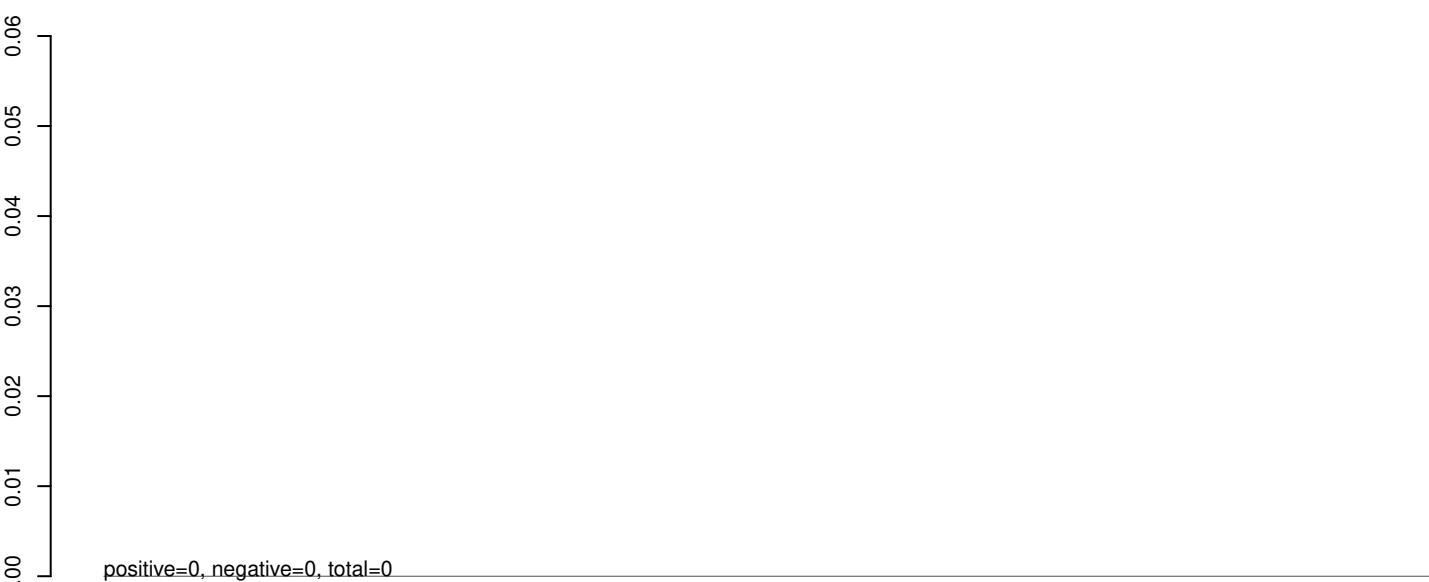


0 2000 4000 6000 8000 10000

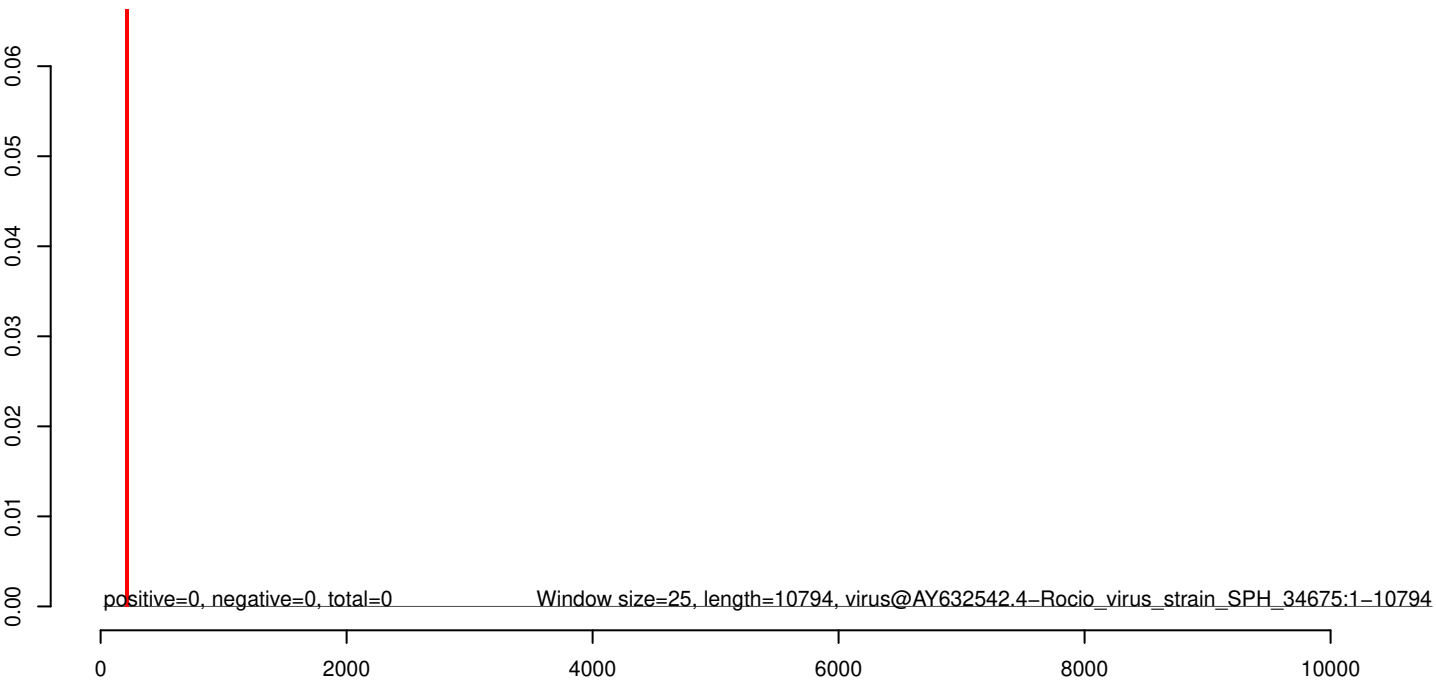
AeAeg_Aag2_BTV2_AK.18_23.rep



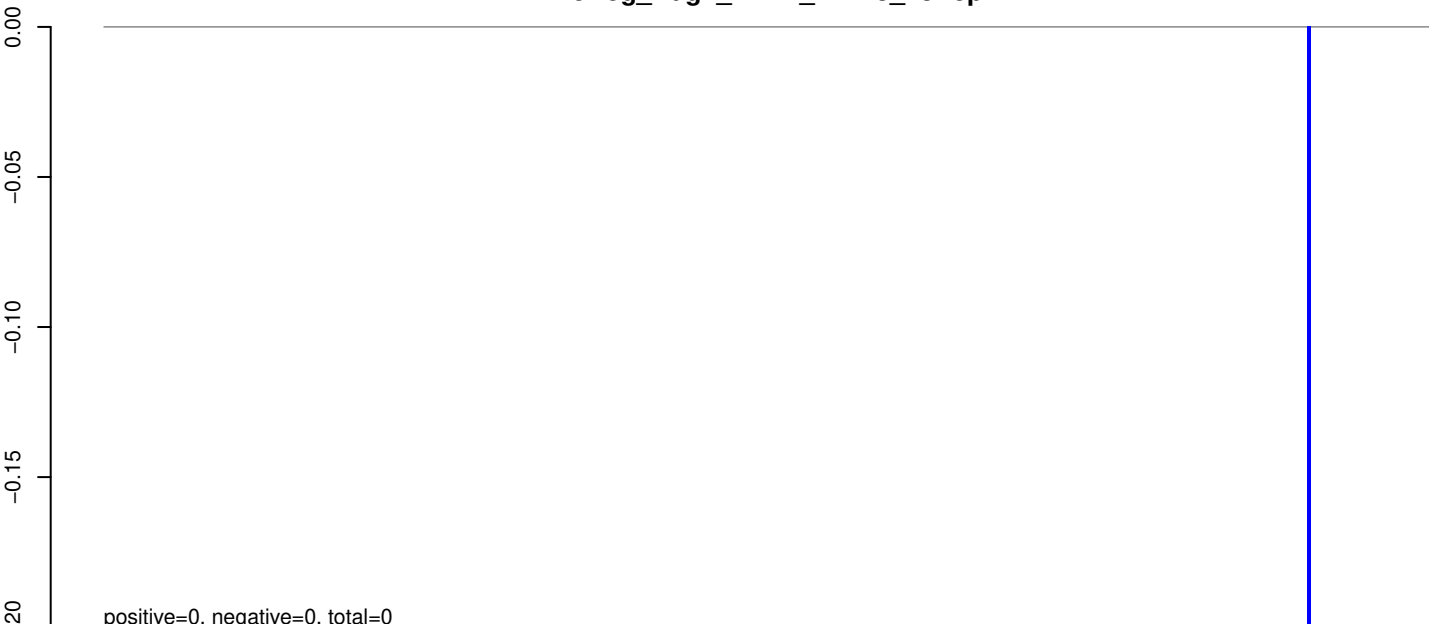
AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



Window size=25, length=10922, virus@AY632538.4-Iguape_virus_strain_SPAn_71686:1-10922

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV2_AK.18_23.rep

0.00
-0.01
-0.02
-0.03
-0.04
-0.05
-0.06

positive=0, negative=0, total=0

AeAeg_Aag2_BTV2_AK.24_35.rep

0.00
-0.01
-0.02
-0.03
-0.04
-0.05
-0.06

positive=0, negative=0, total=0

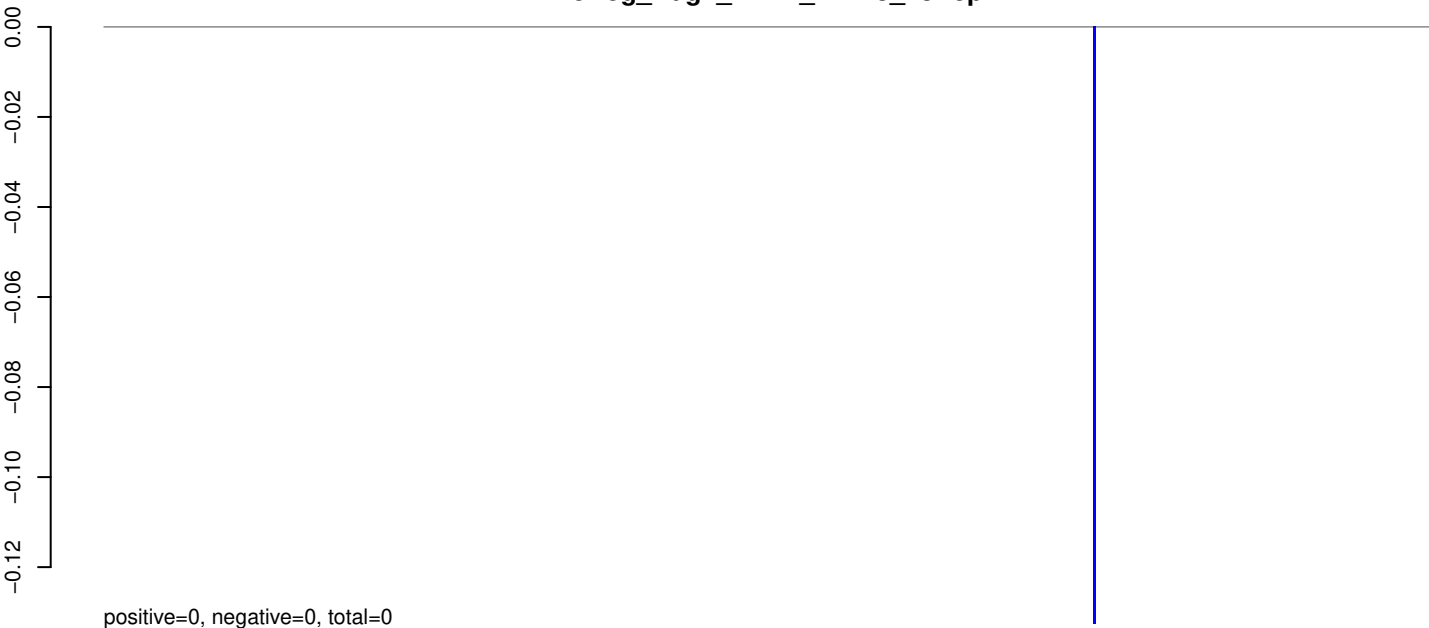
AeAeg_Aag2_BTV2_AK.rep

0.00
-0.01
-0.02
-0.03
-0.04
-0.05
-0.06

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

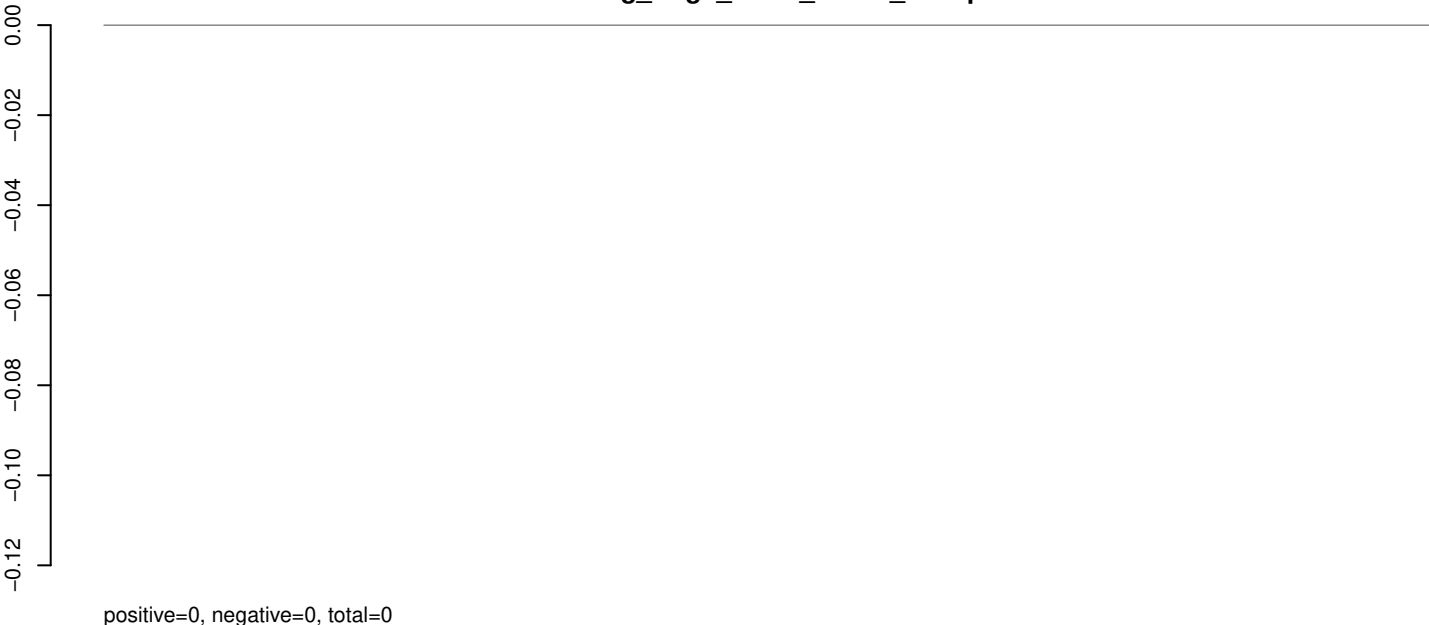
0 200 400 600 800 1000

AeAeg_Aag2_BTV2_AK.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BTV2_AK.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BTV2_AK.rep



positive=0, negative=0, total=0

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



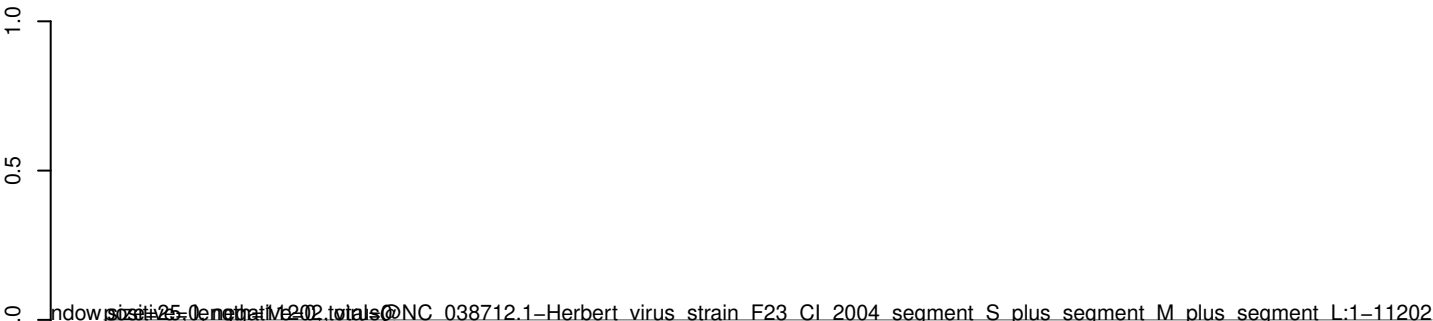
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



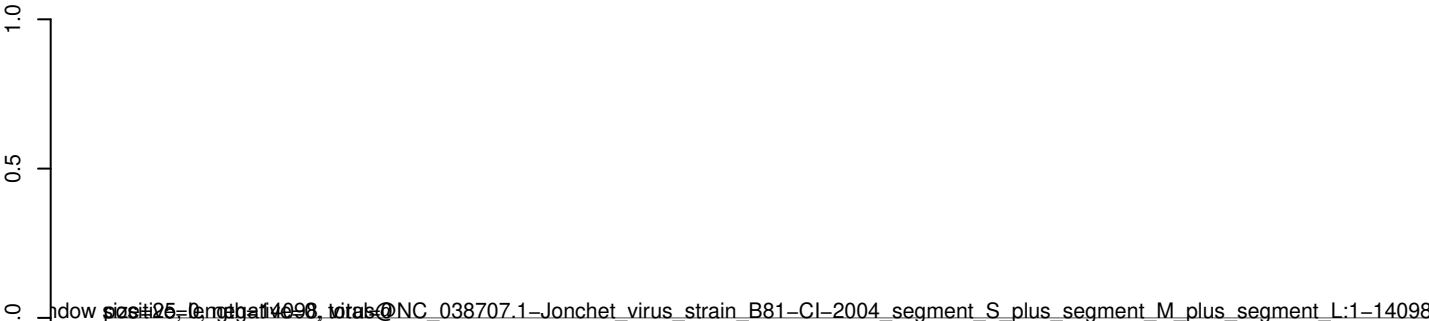
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



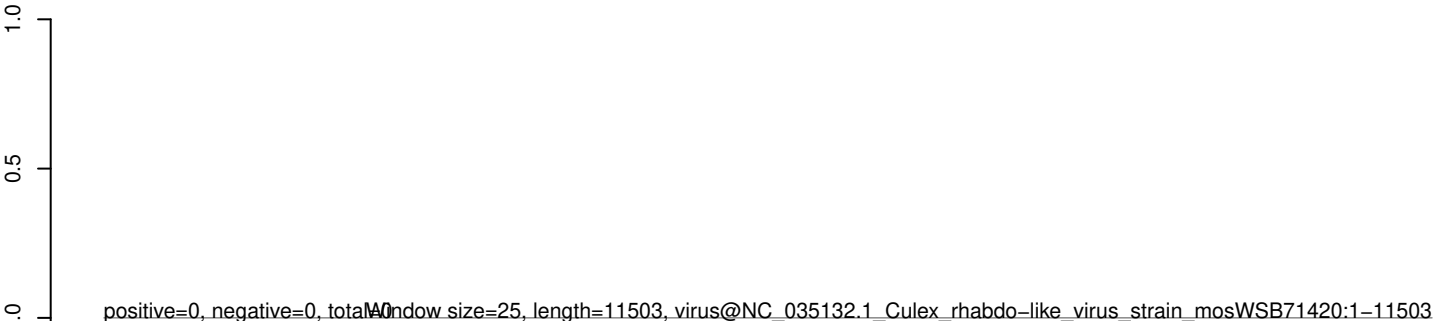
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



0 2000 4000 6000 8000 10000 12000

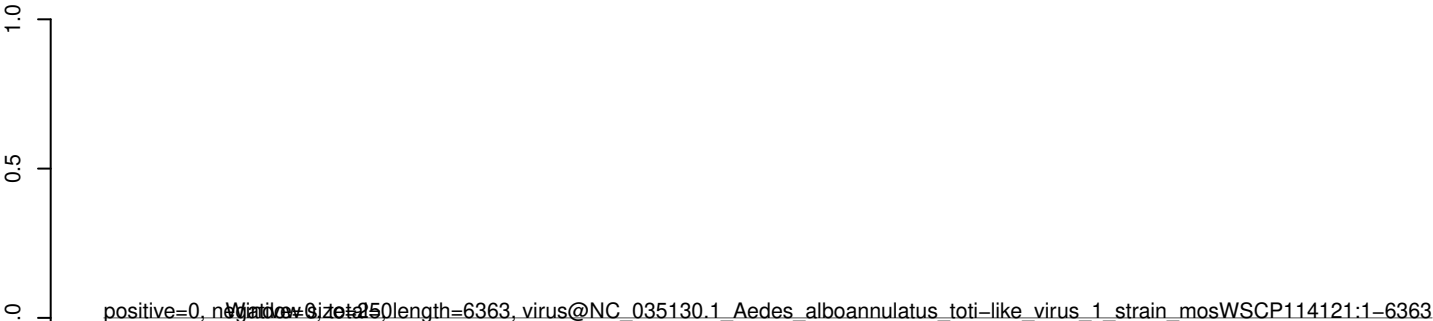
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



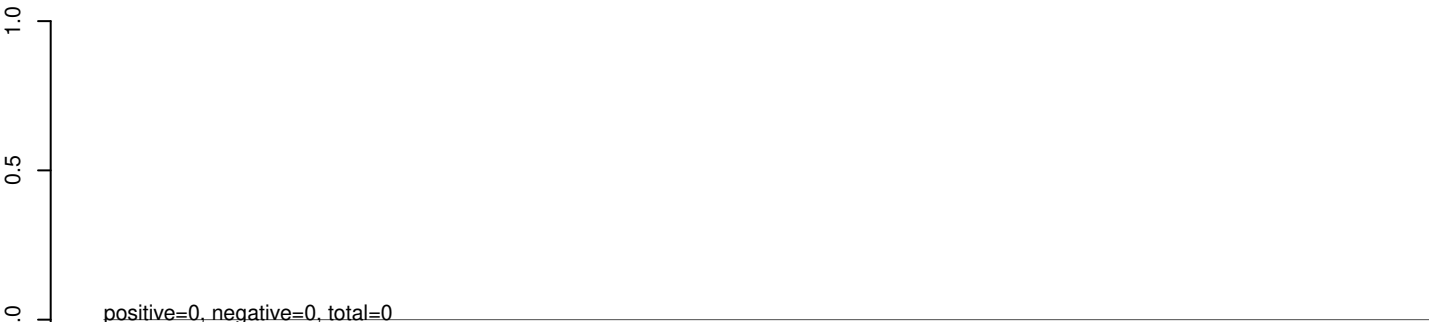
AeAeg_Aag2_BTV2_AK.rep



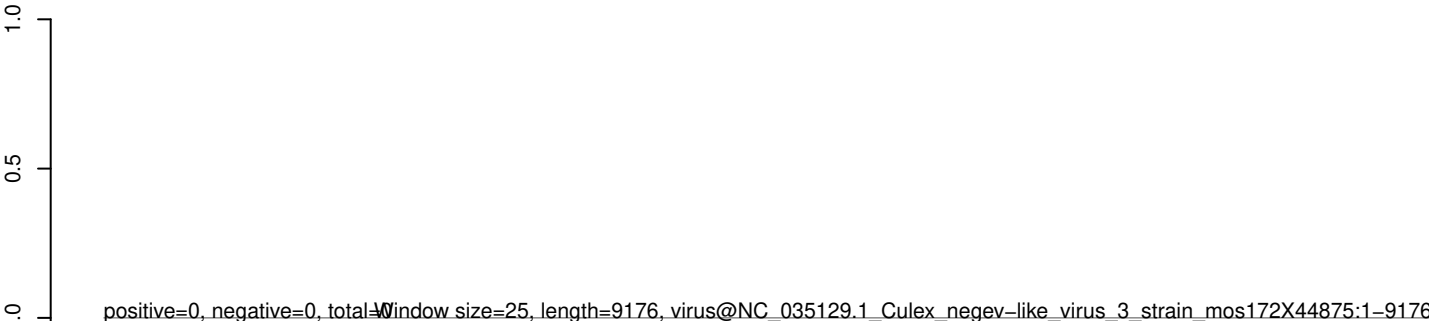
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



0 2000 4000 6000 8000

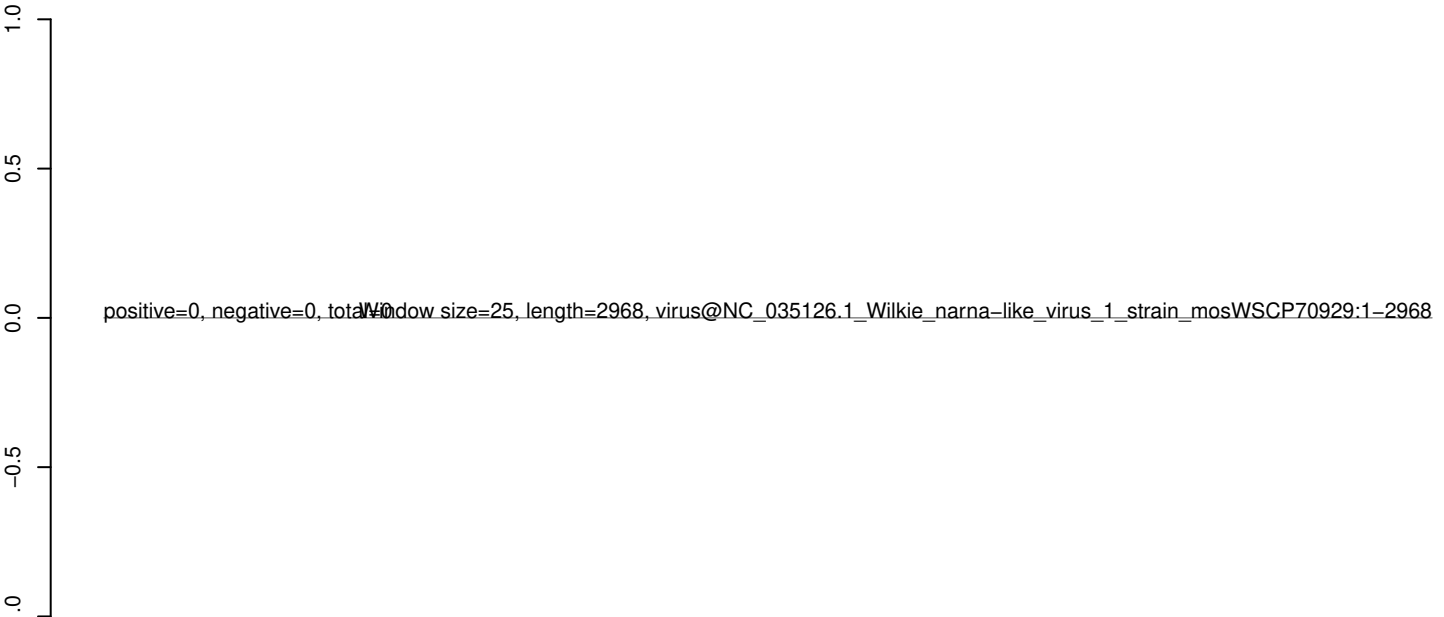
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



0 500 1000 1500 2000 2500 3000

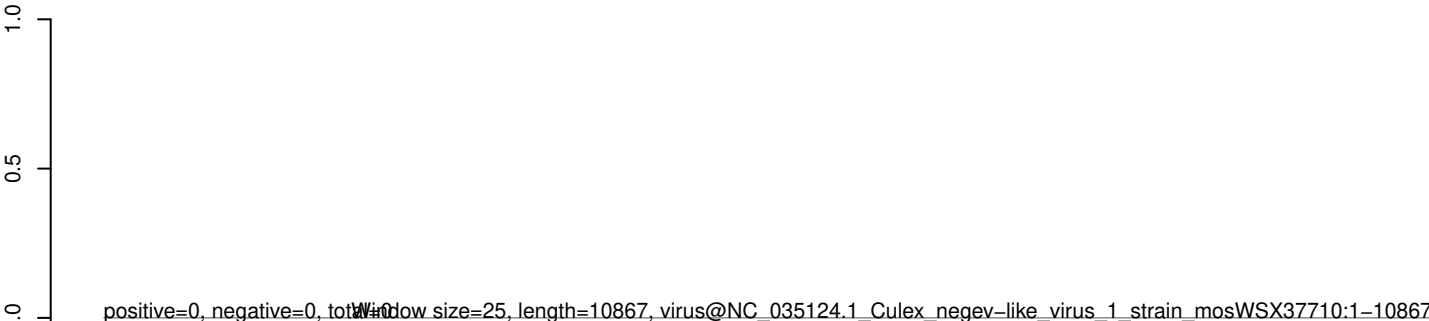
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



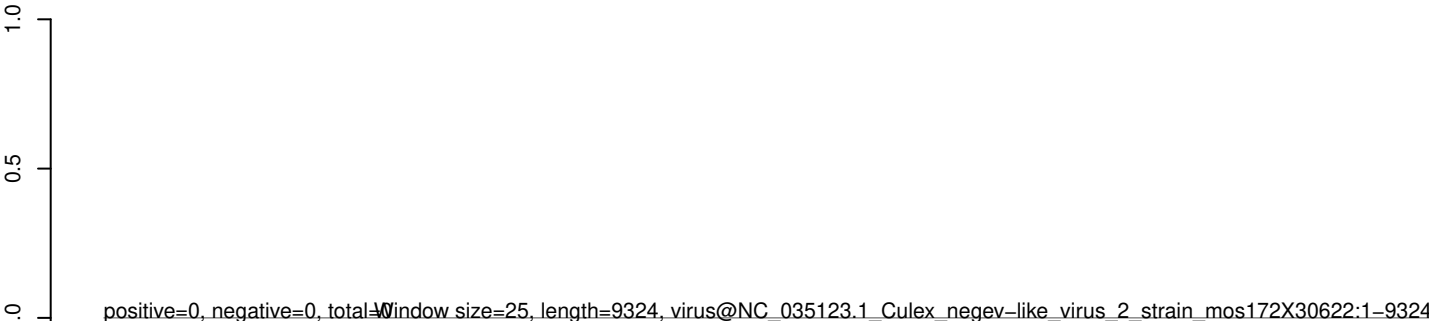
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep

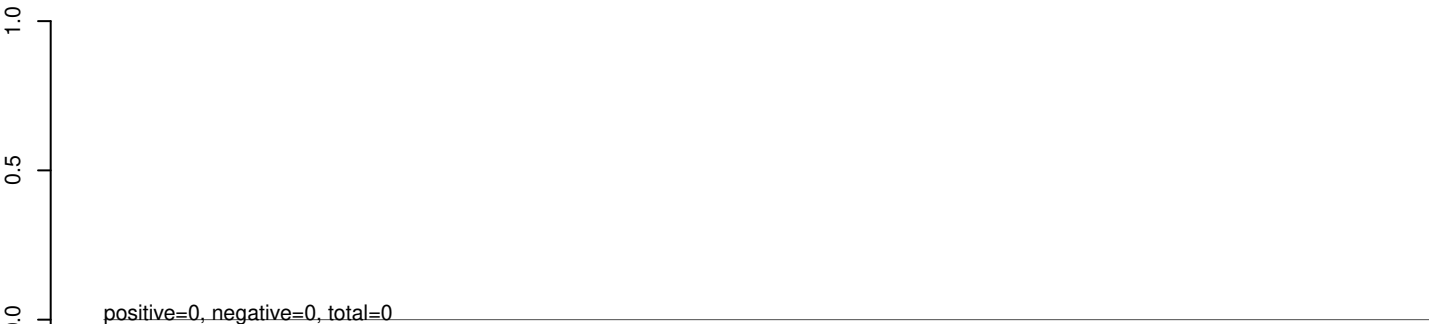


0 2000 4000 6000 8000

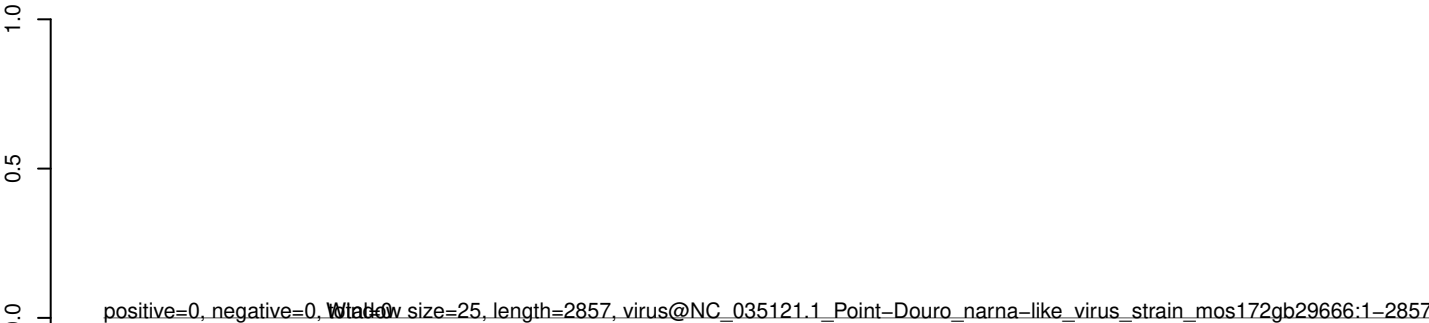
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



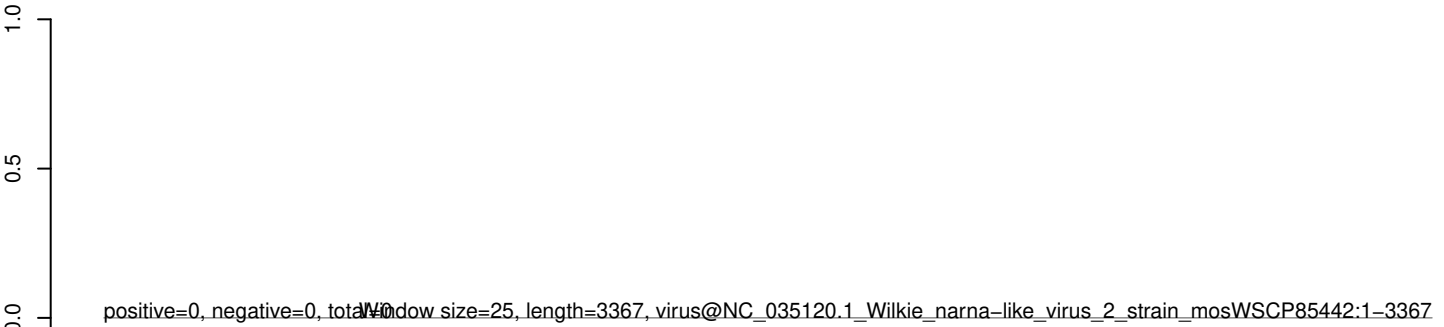
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



0 500 1000 1500 2000 2500 3000 3500

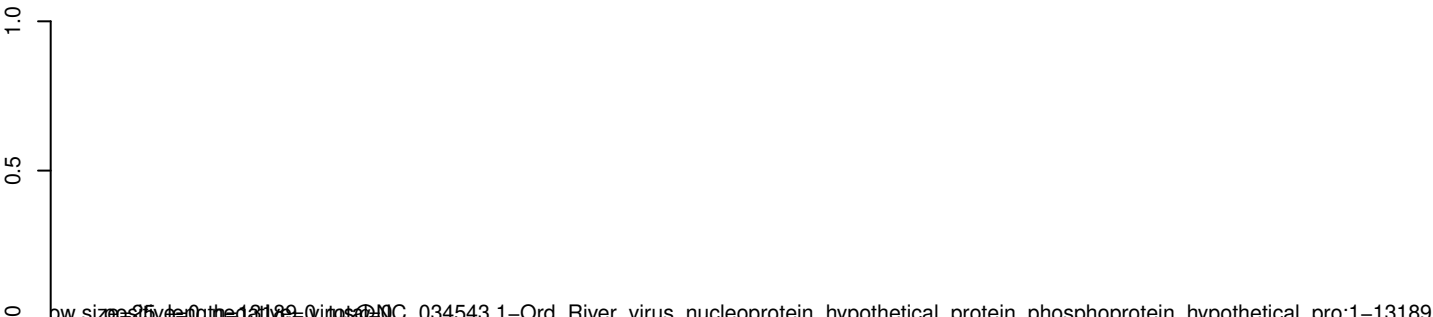
AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



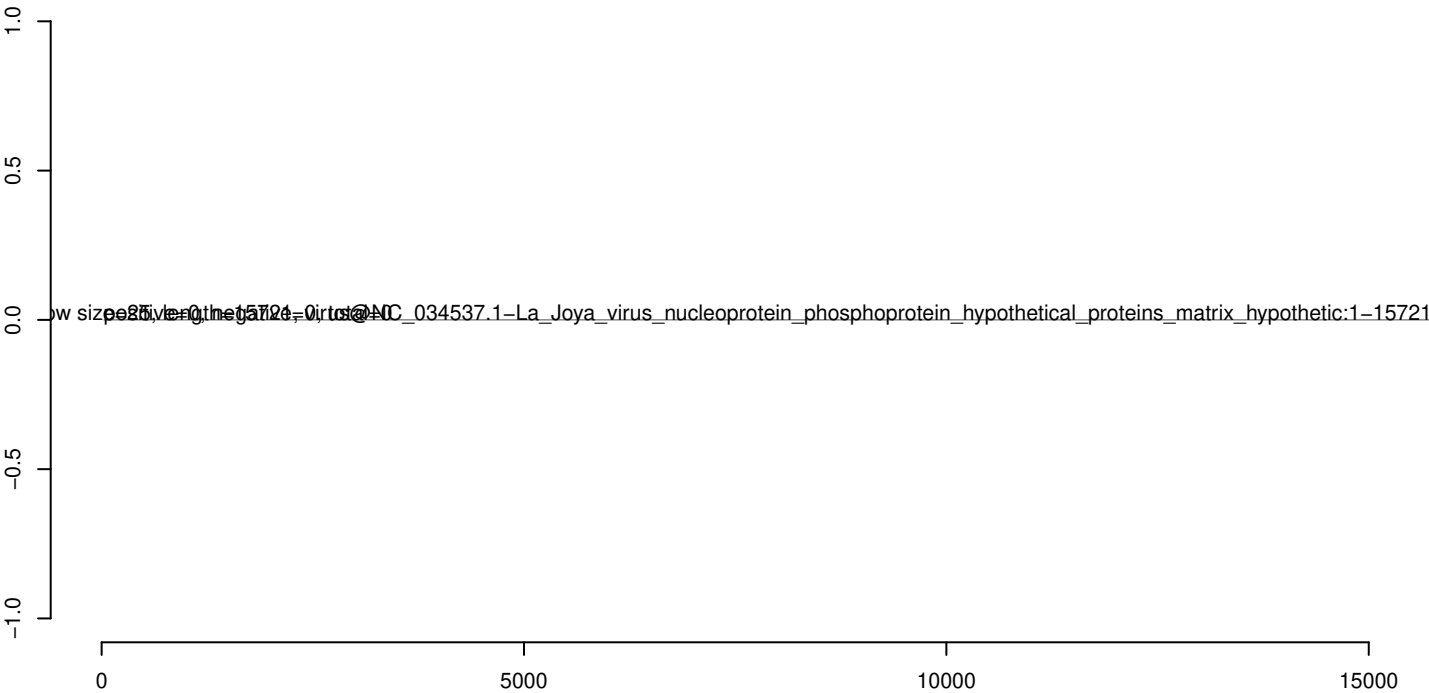
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AeAeg_Aag2_BTV2_AK.24_35.rep



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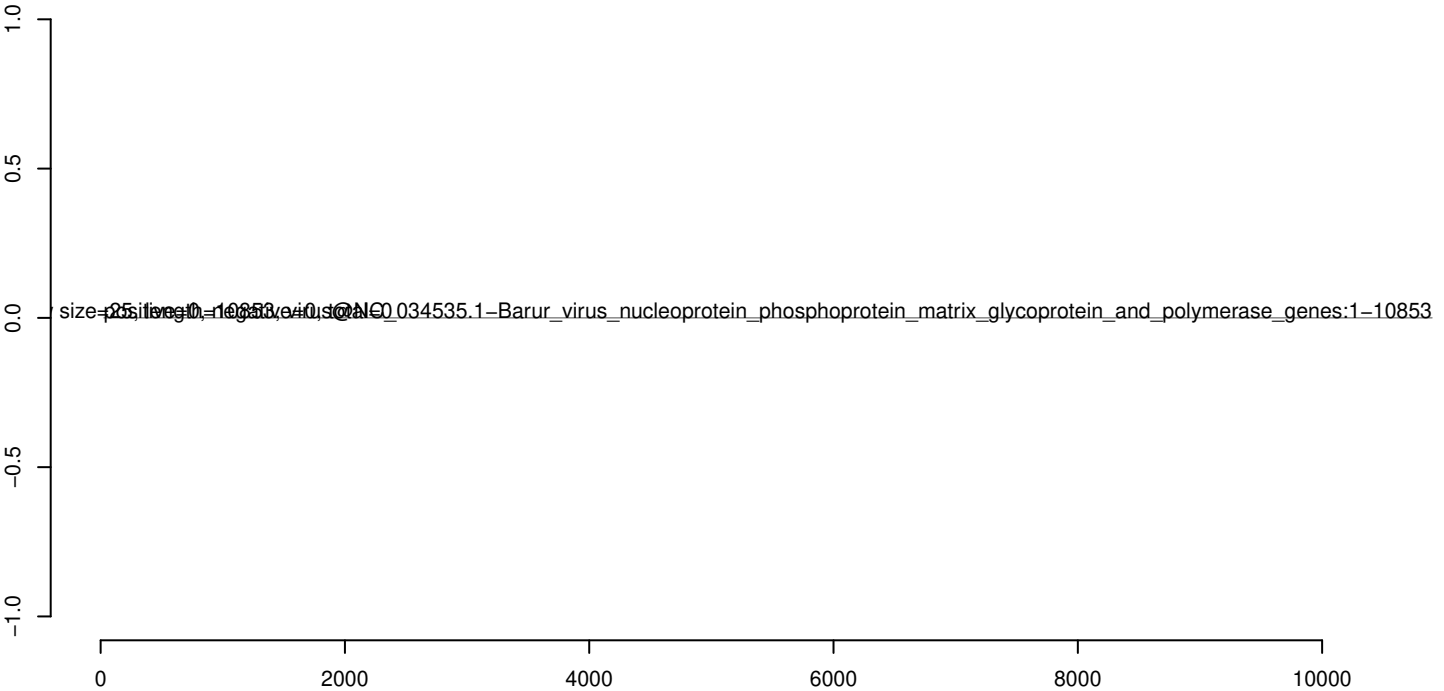
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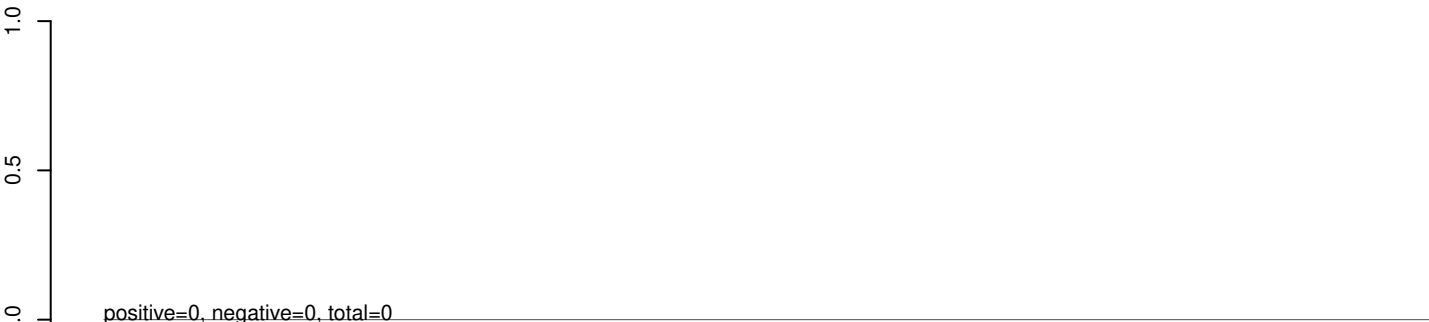
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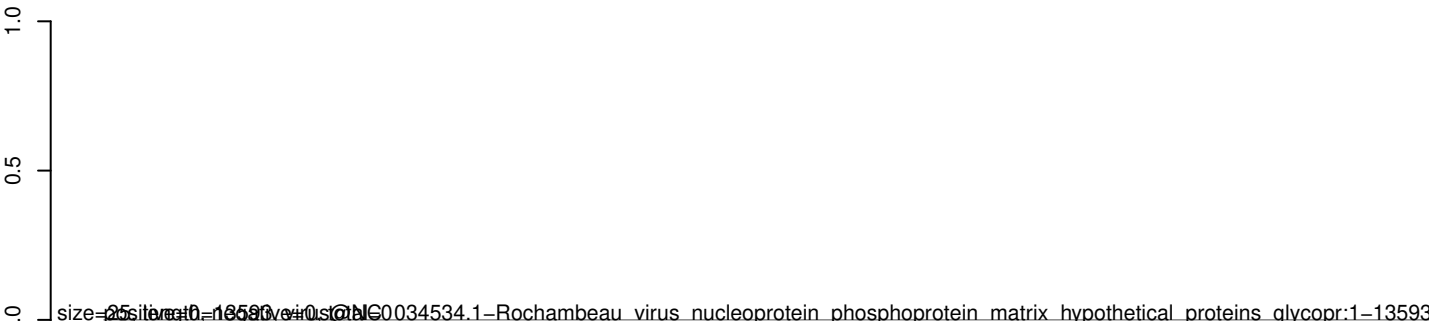
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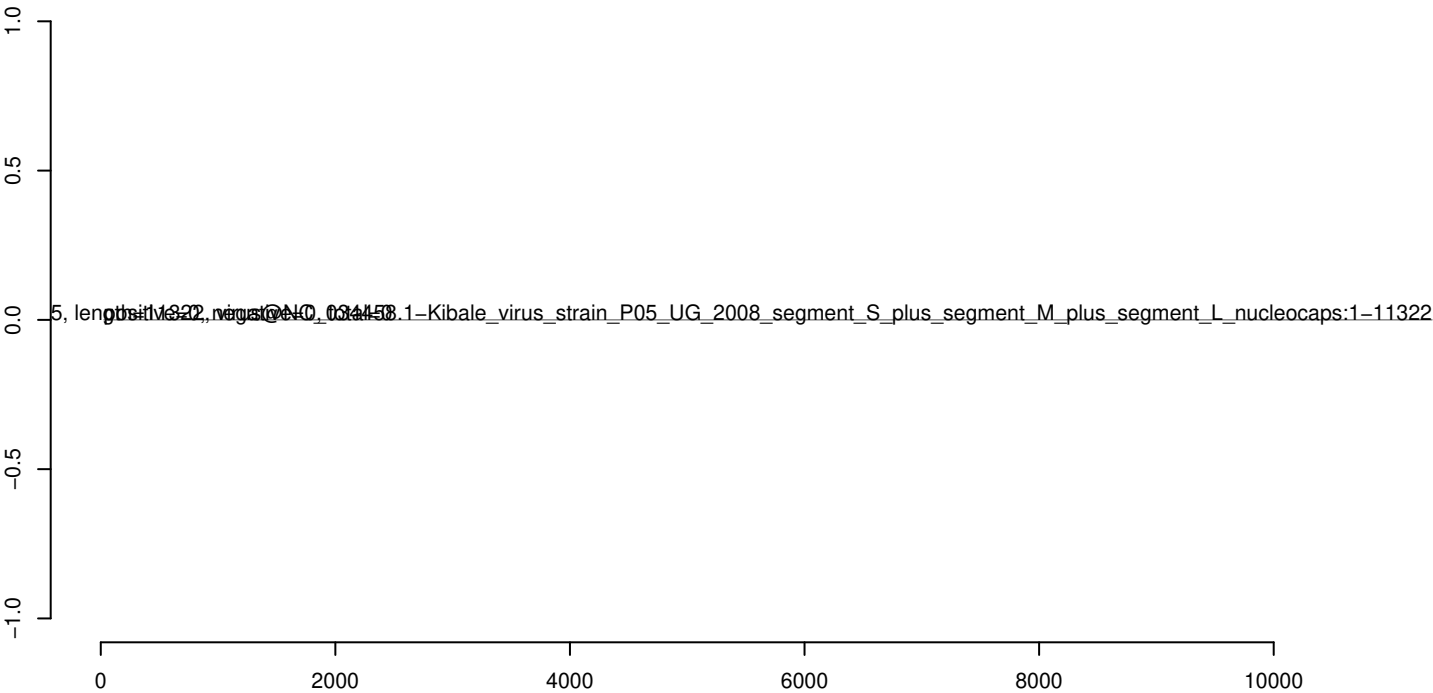
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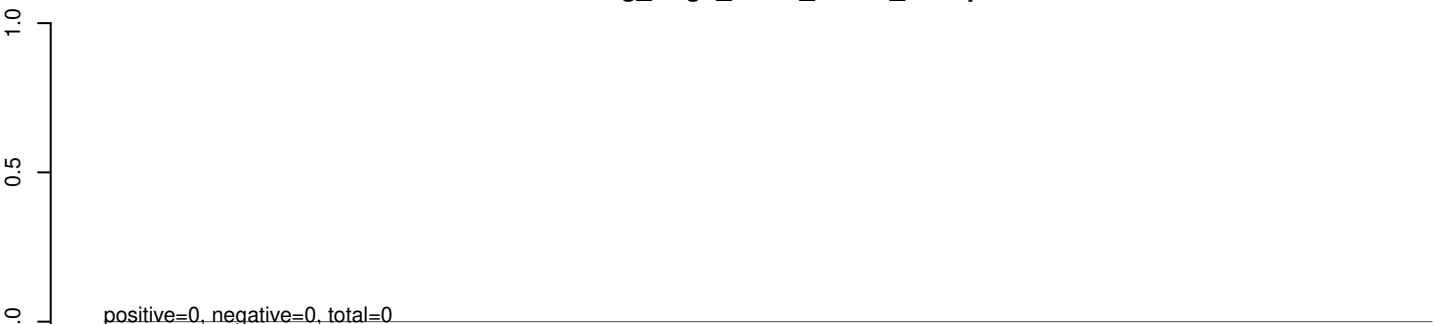
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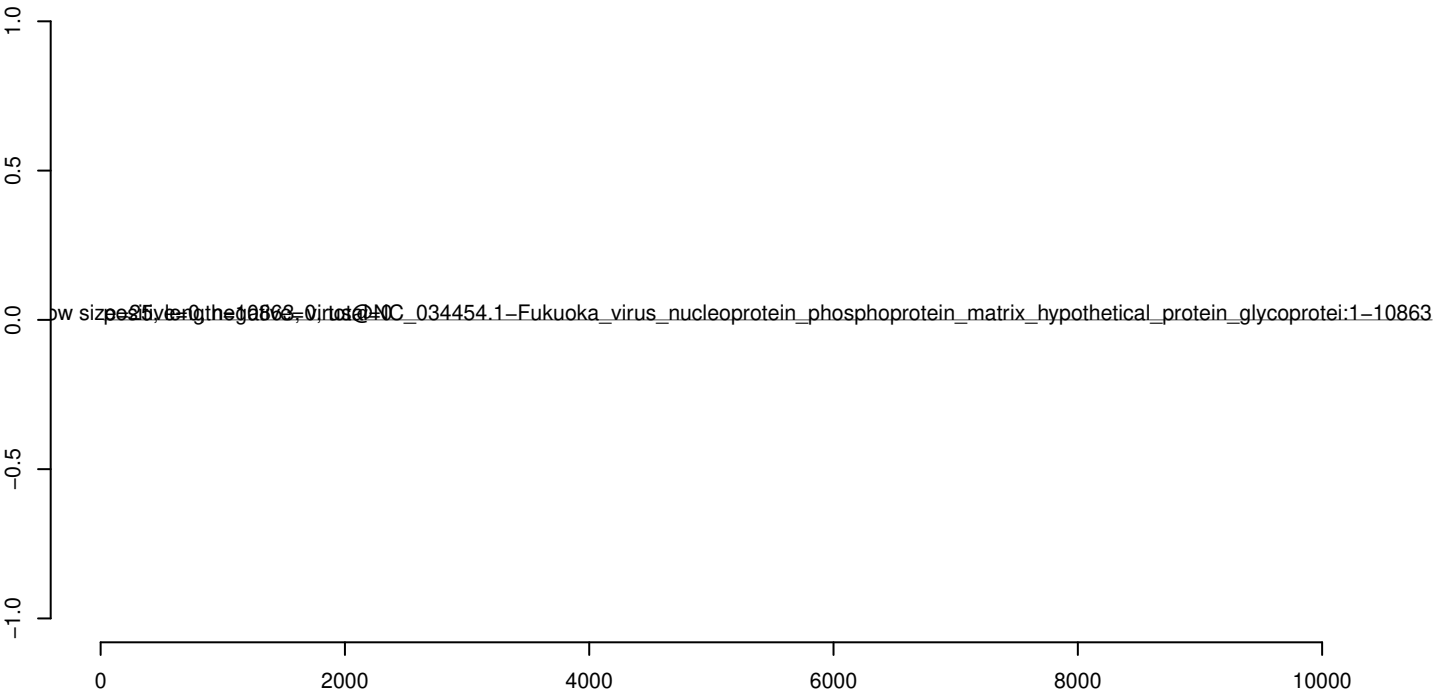
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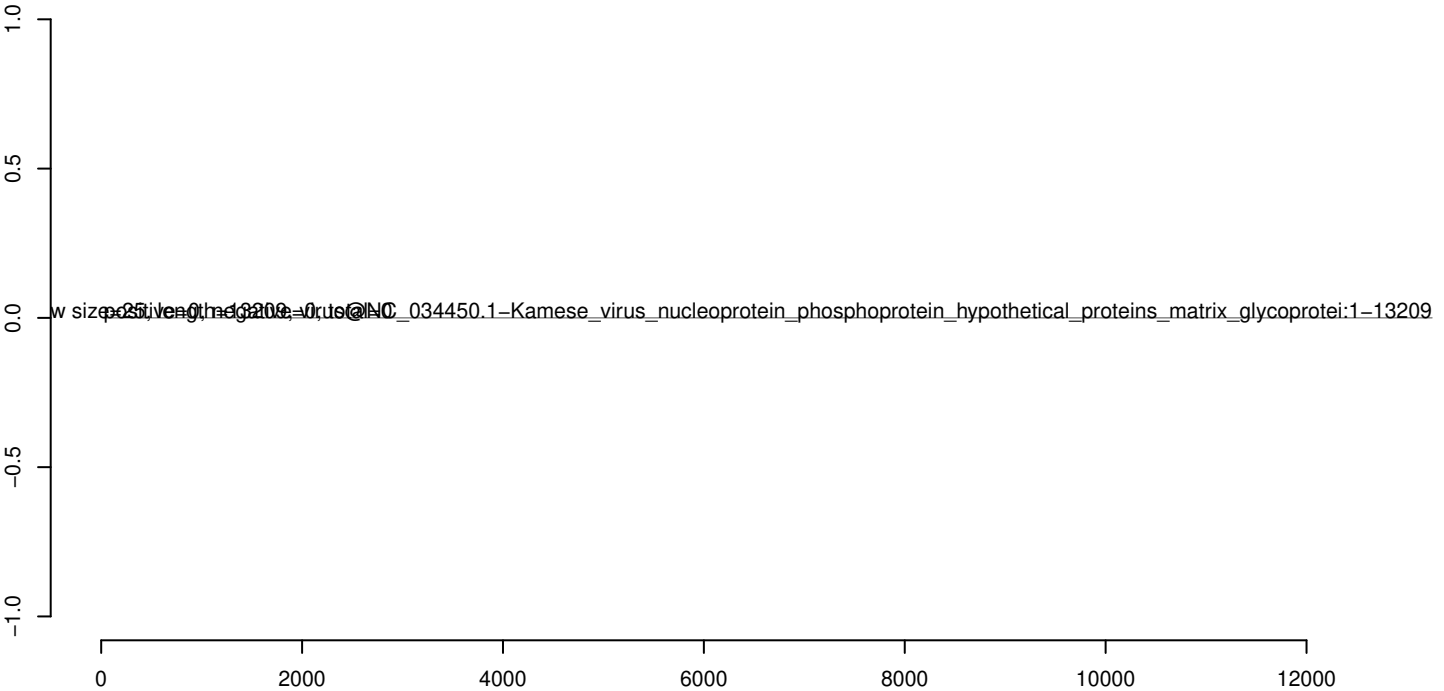
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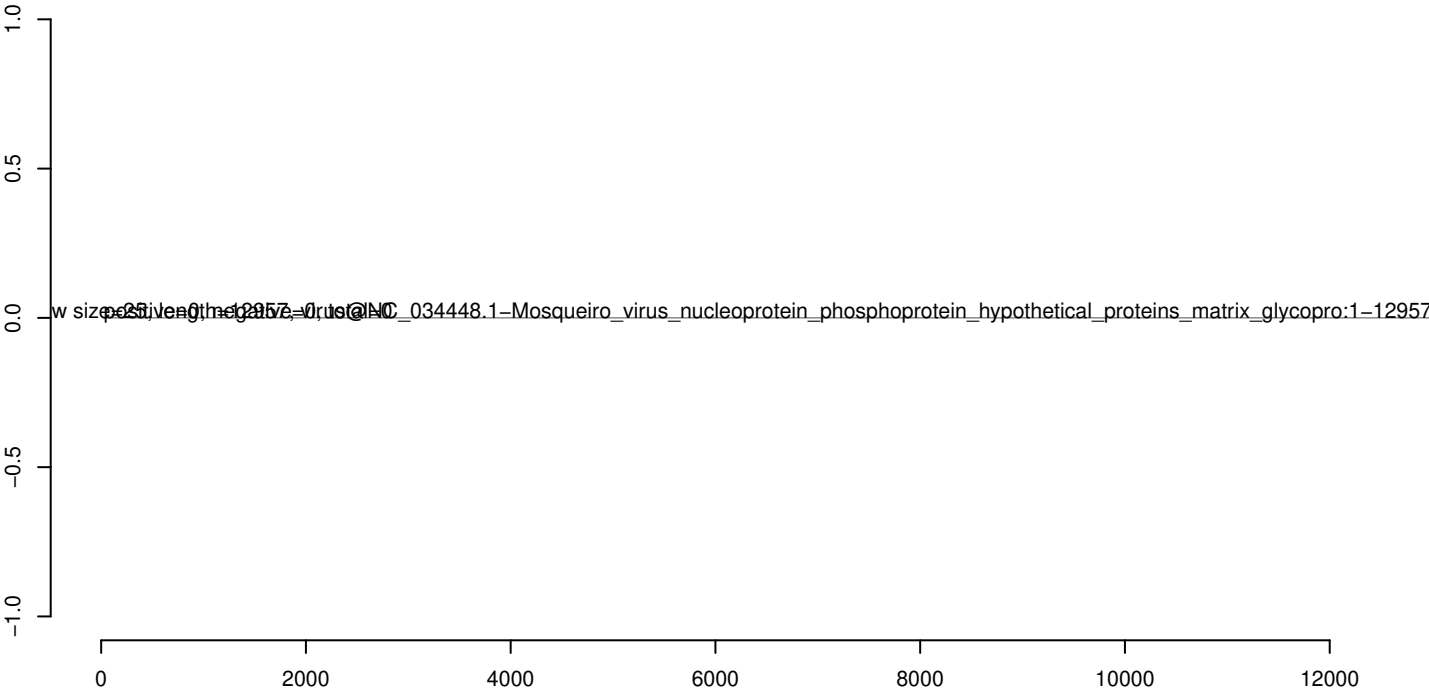
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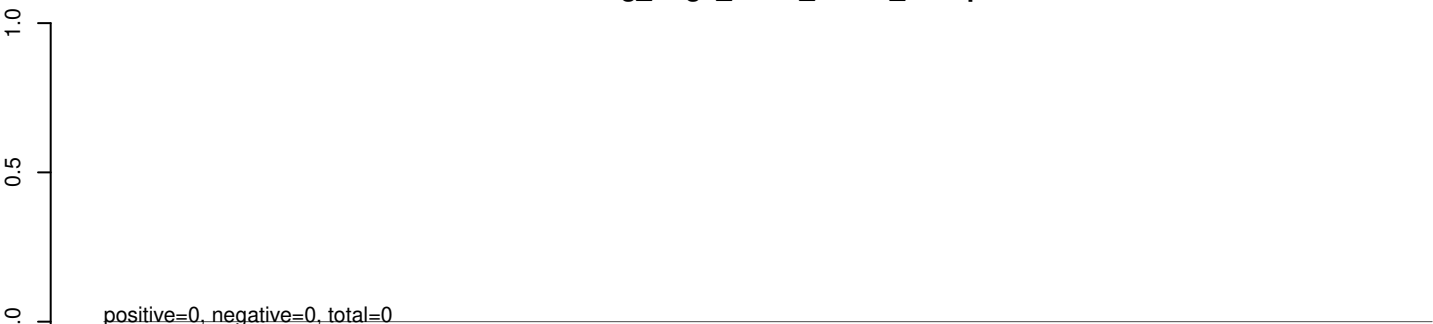
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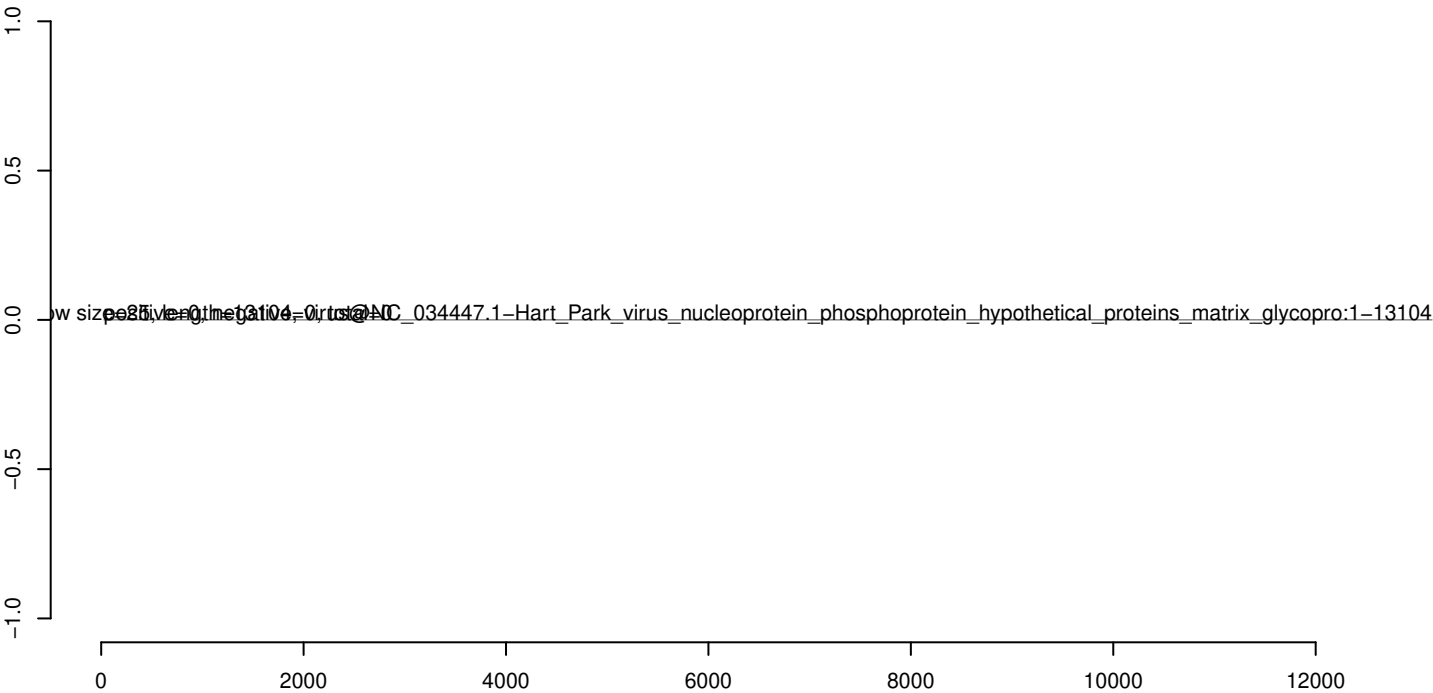
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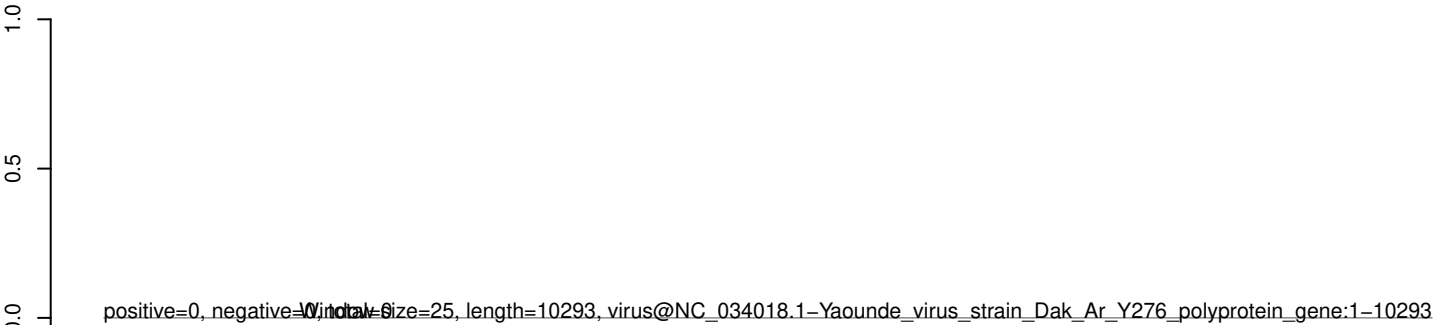
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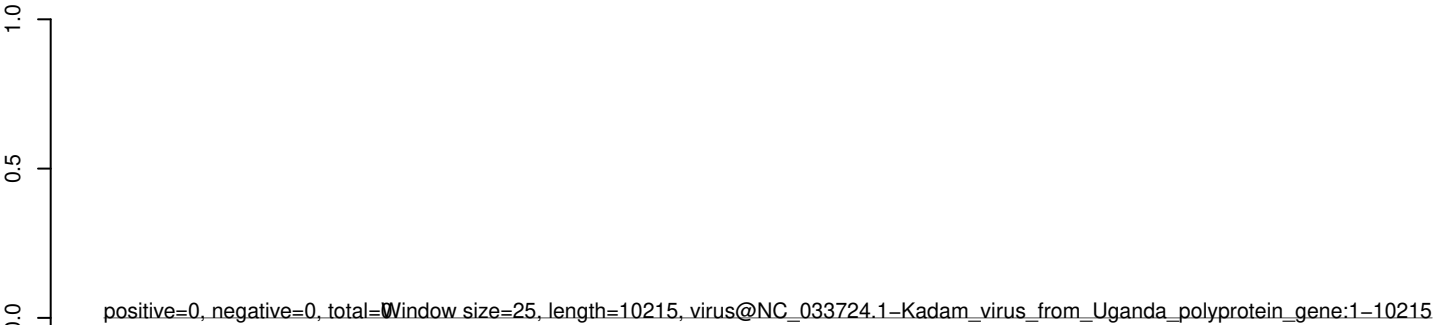
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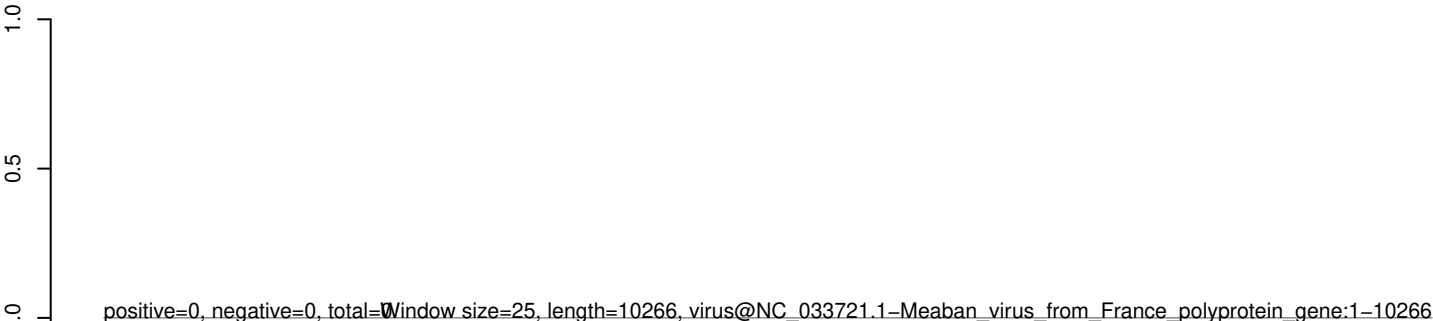
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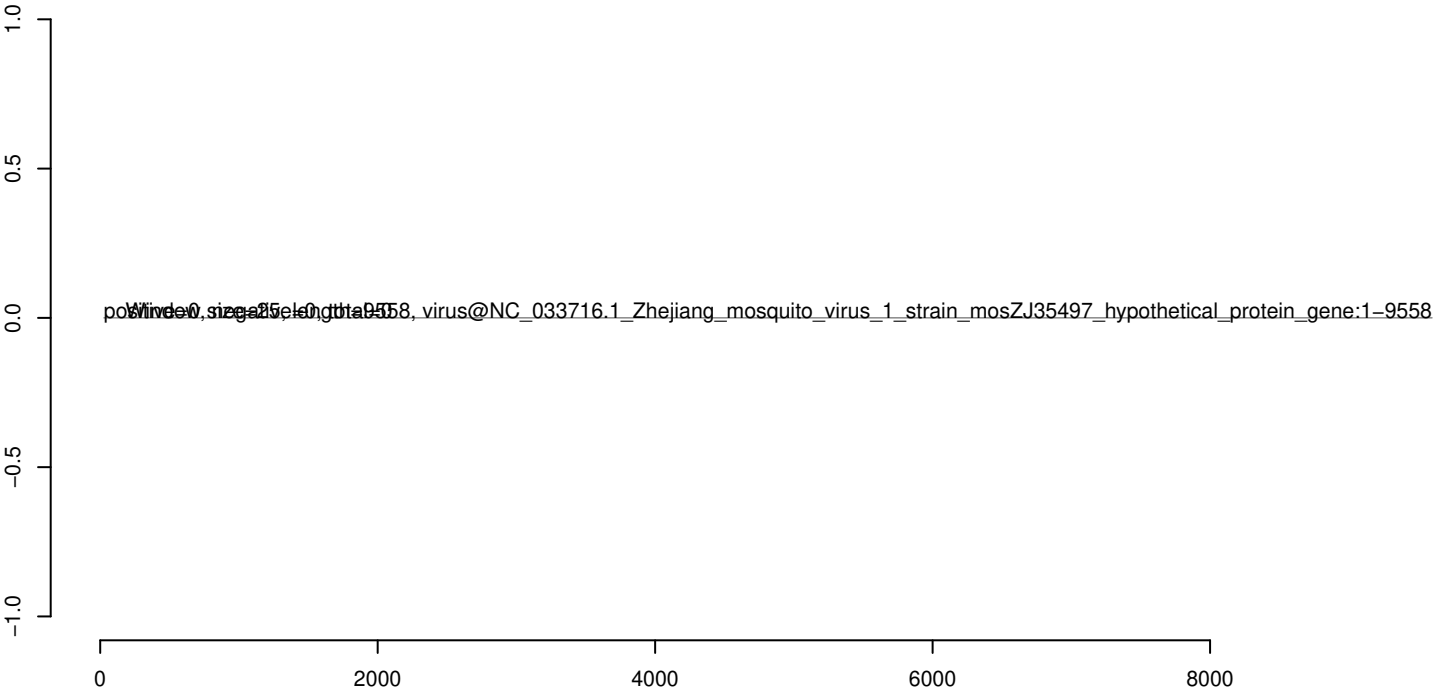
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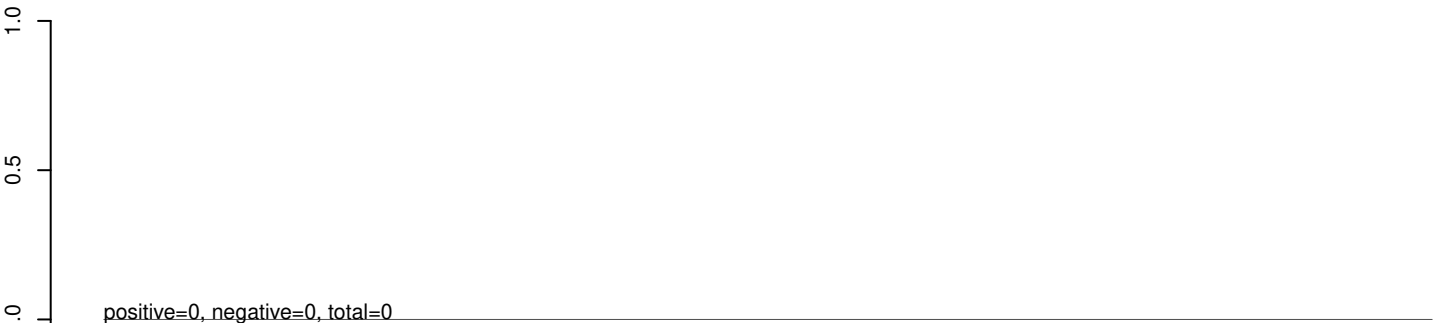
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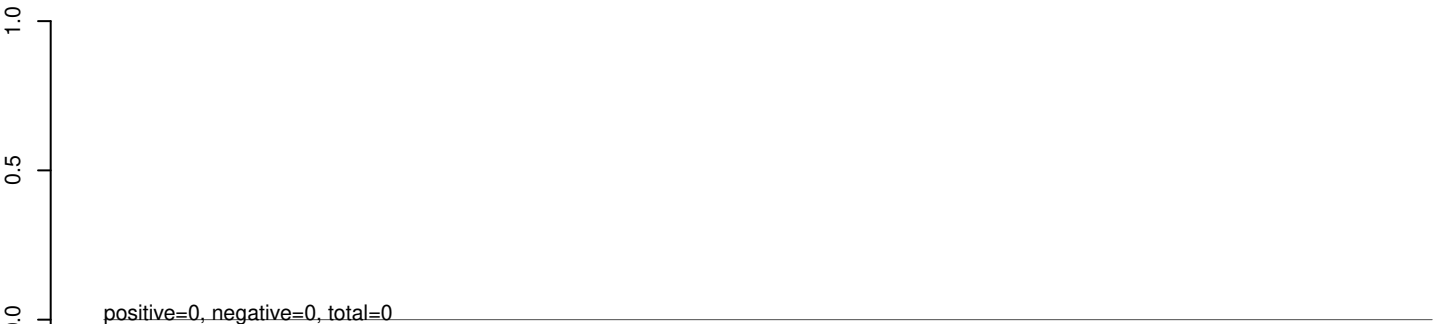
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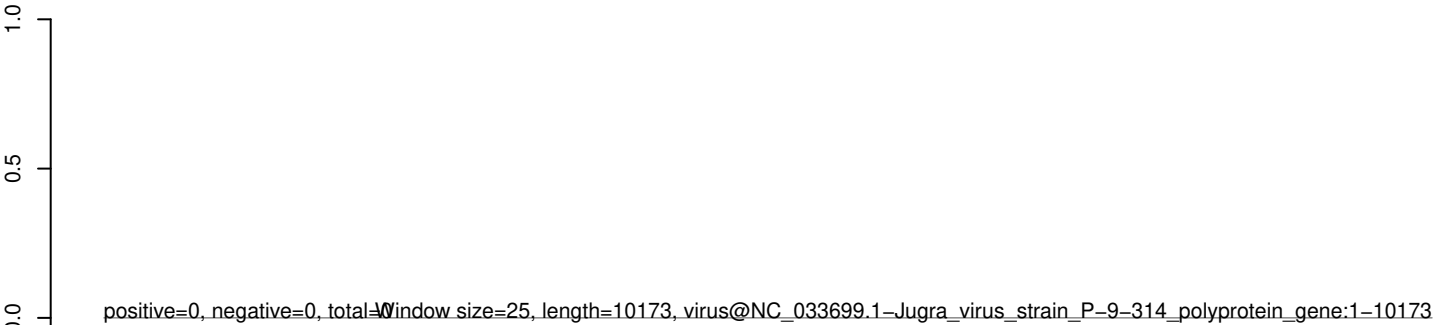
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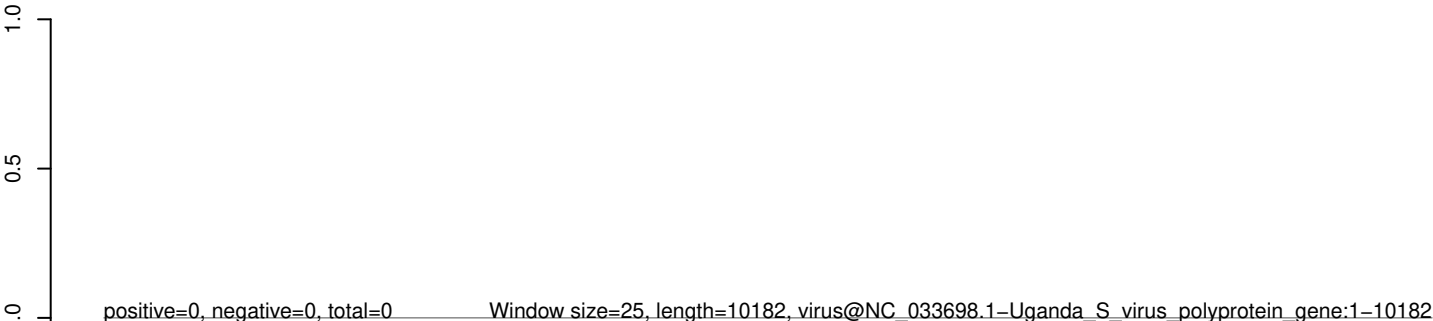
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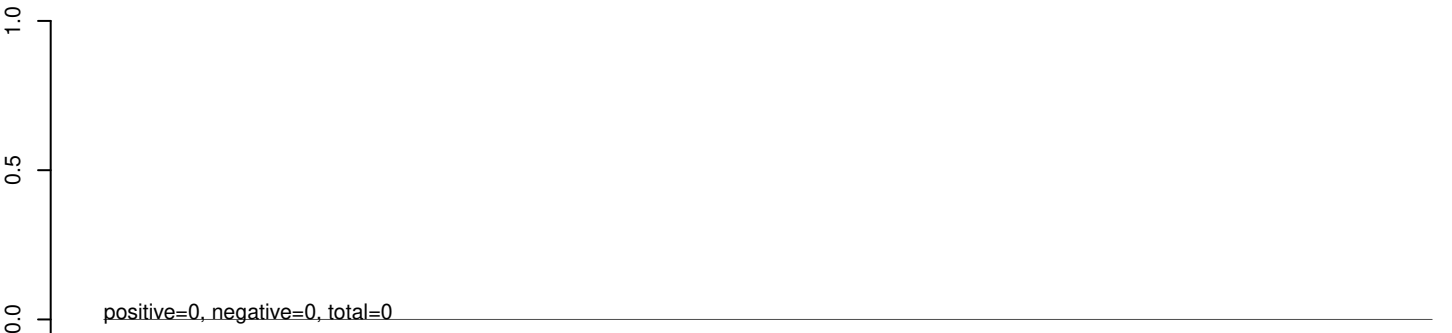
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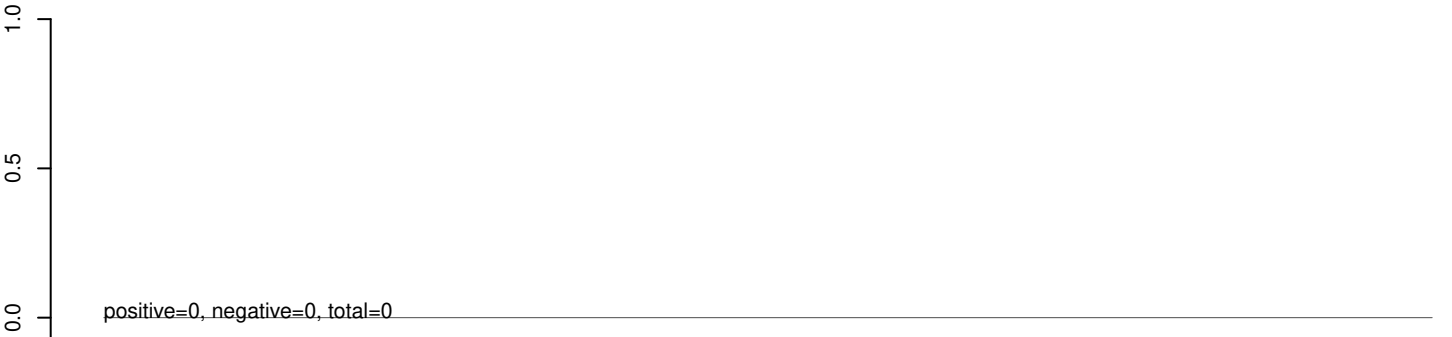
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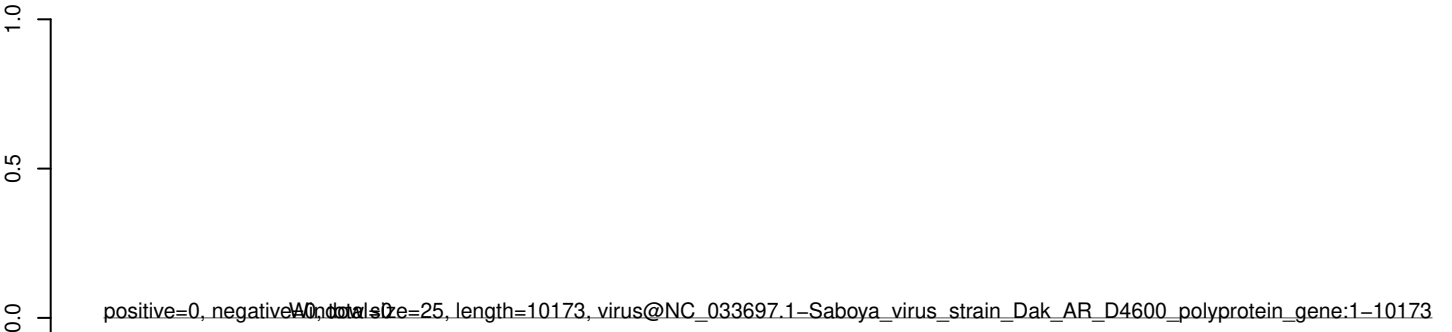
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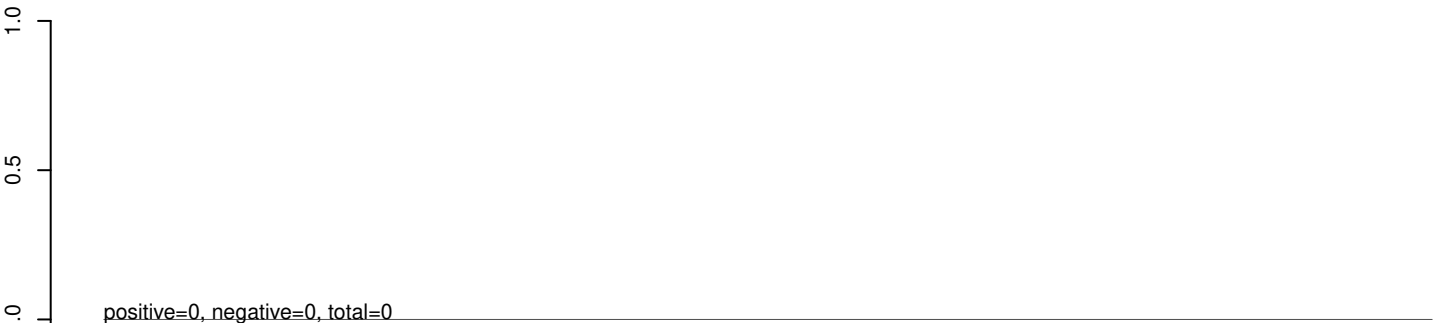
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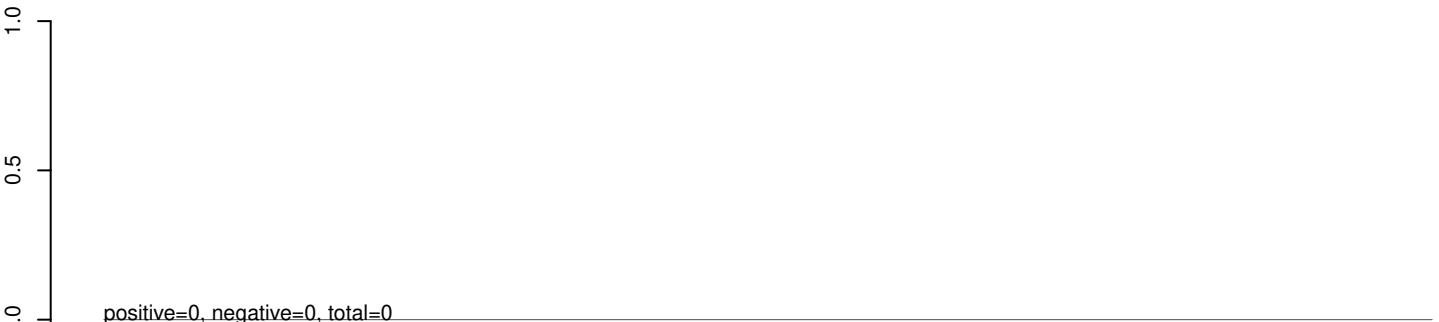
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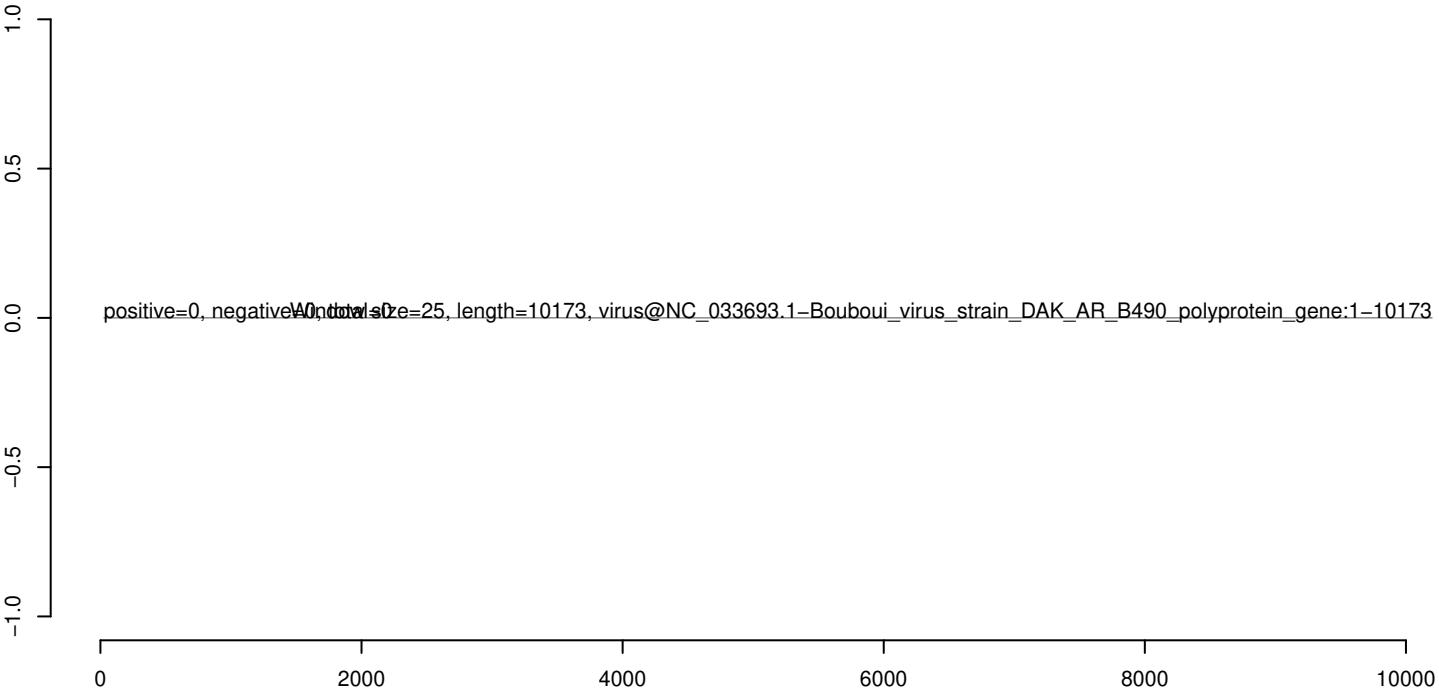
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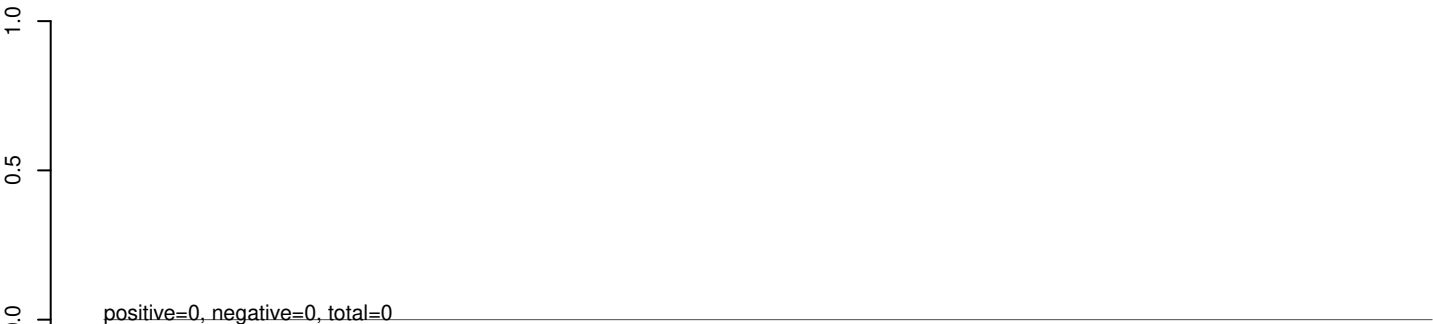
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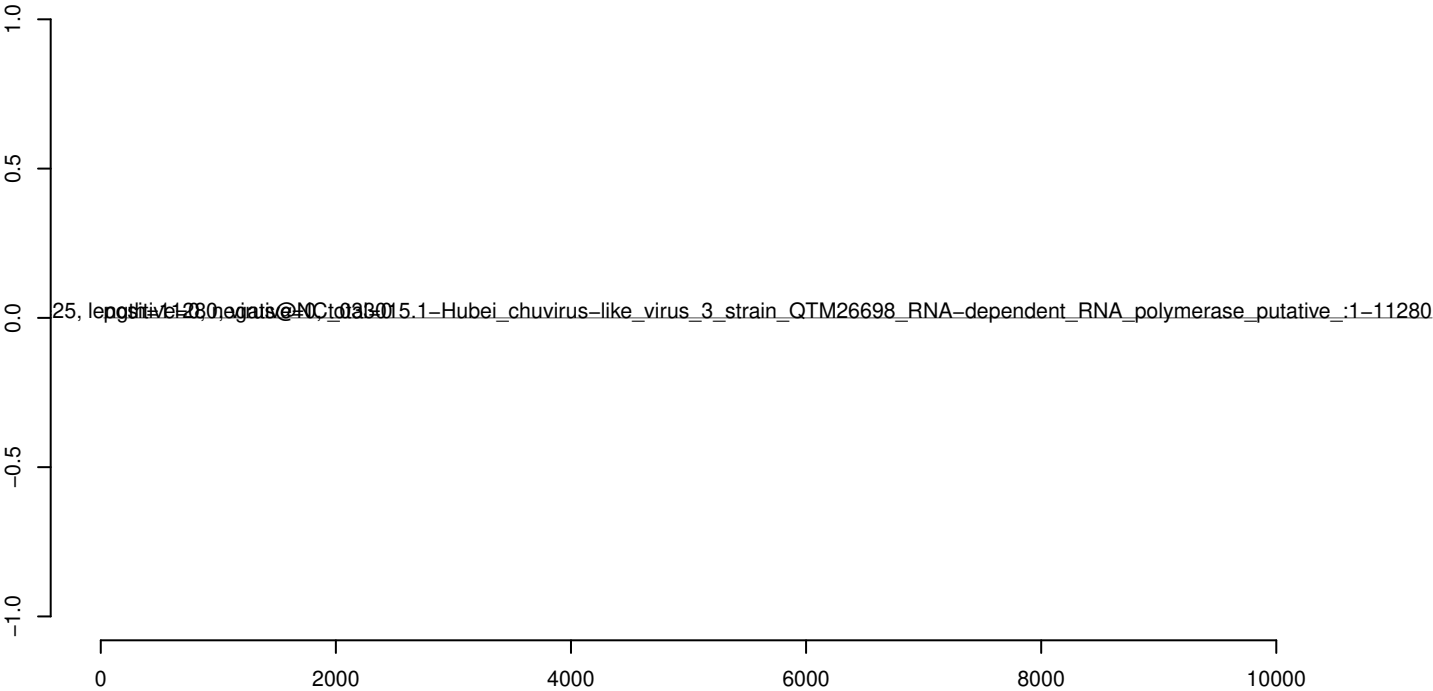
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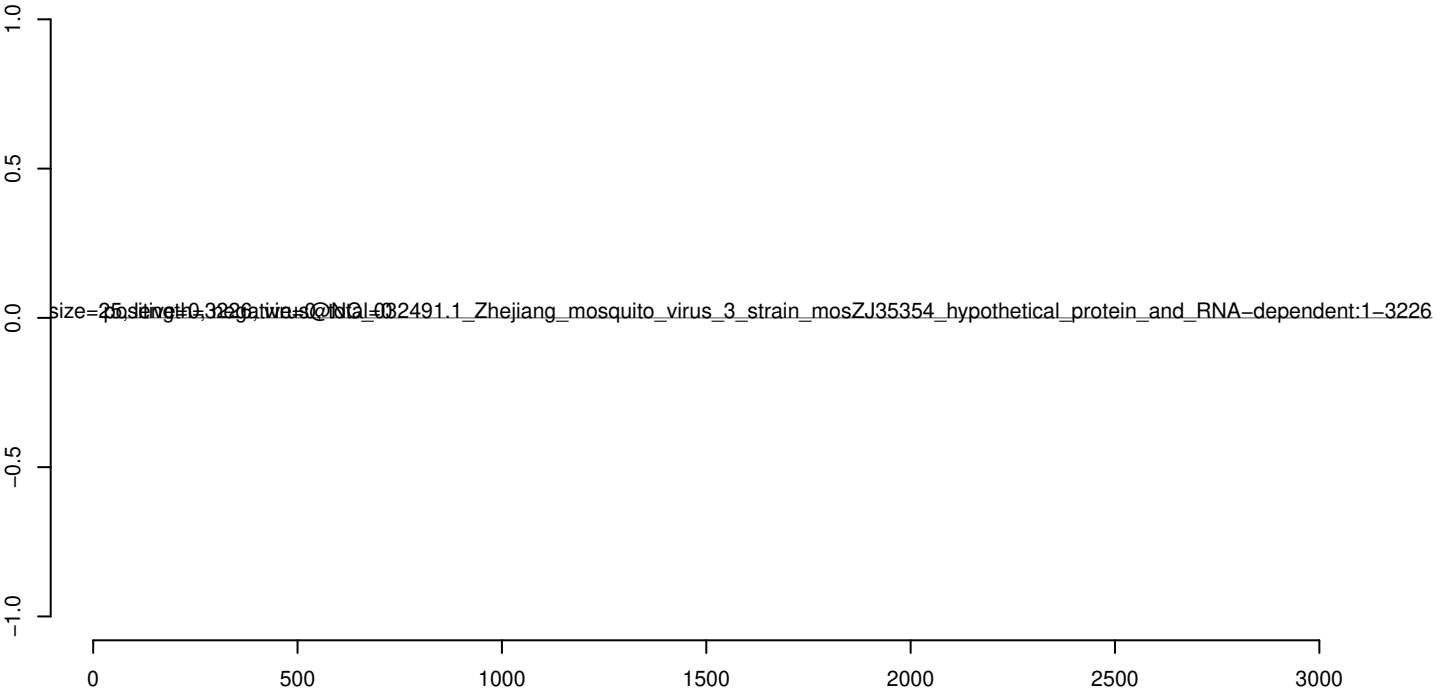
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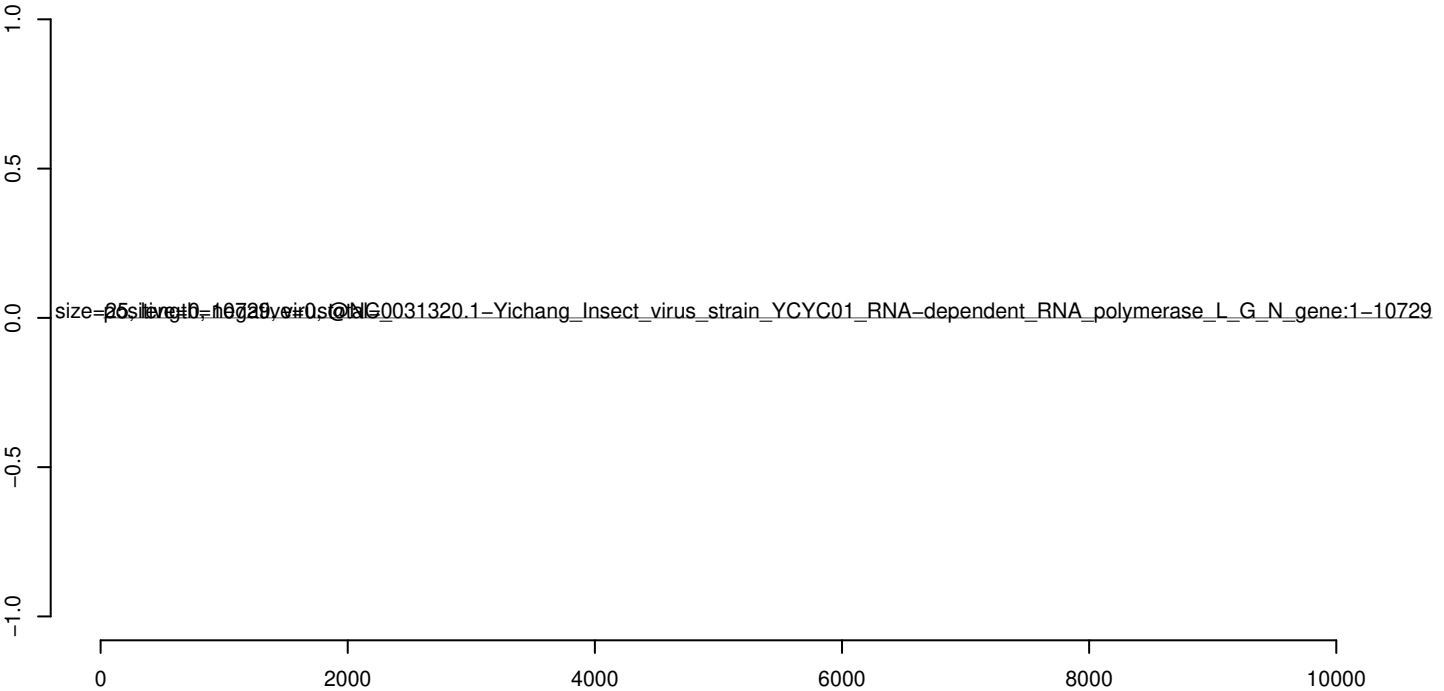
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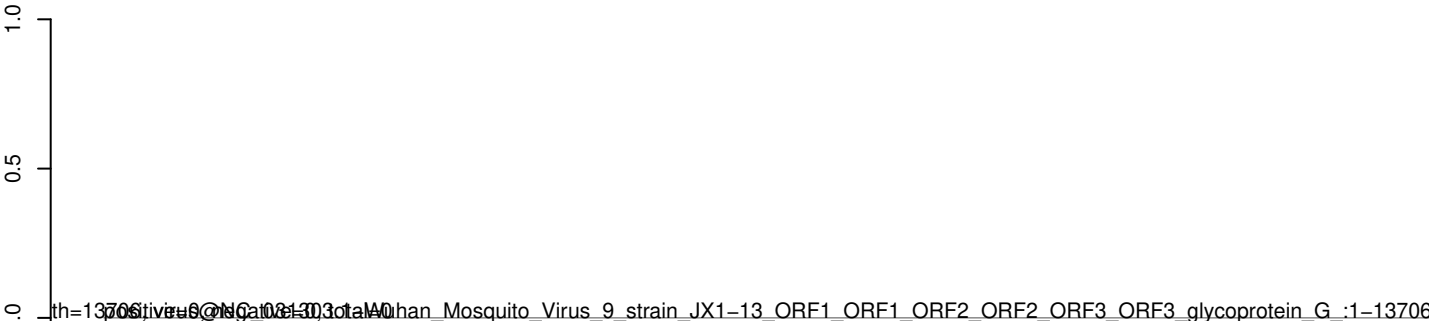
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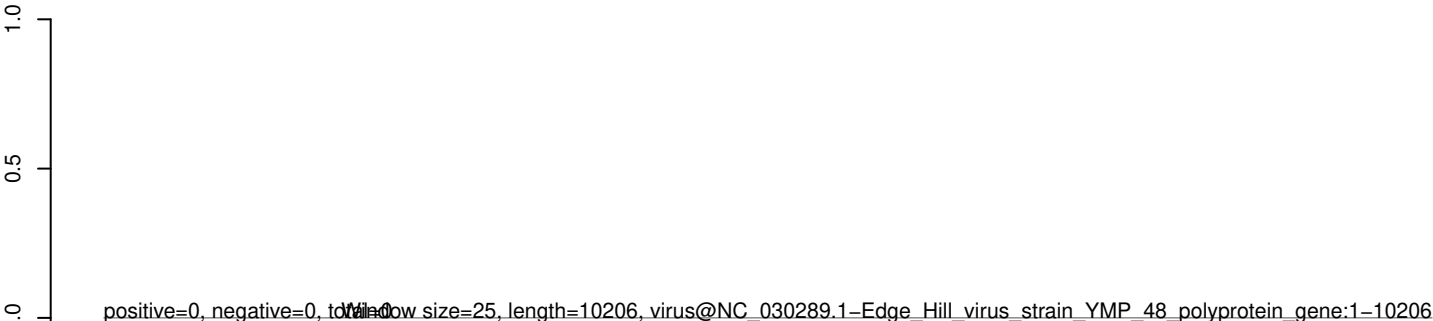
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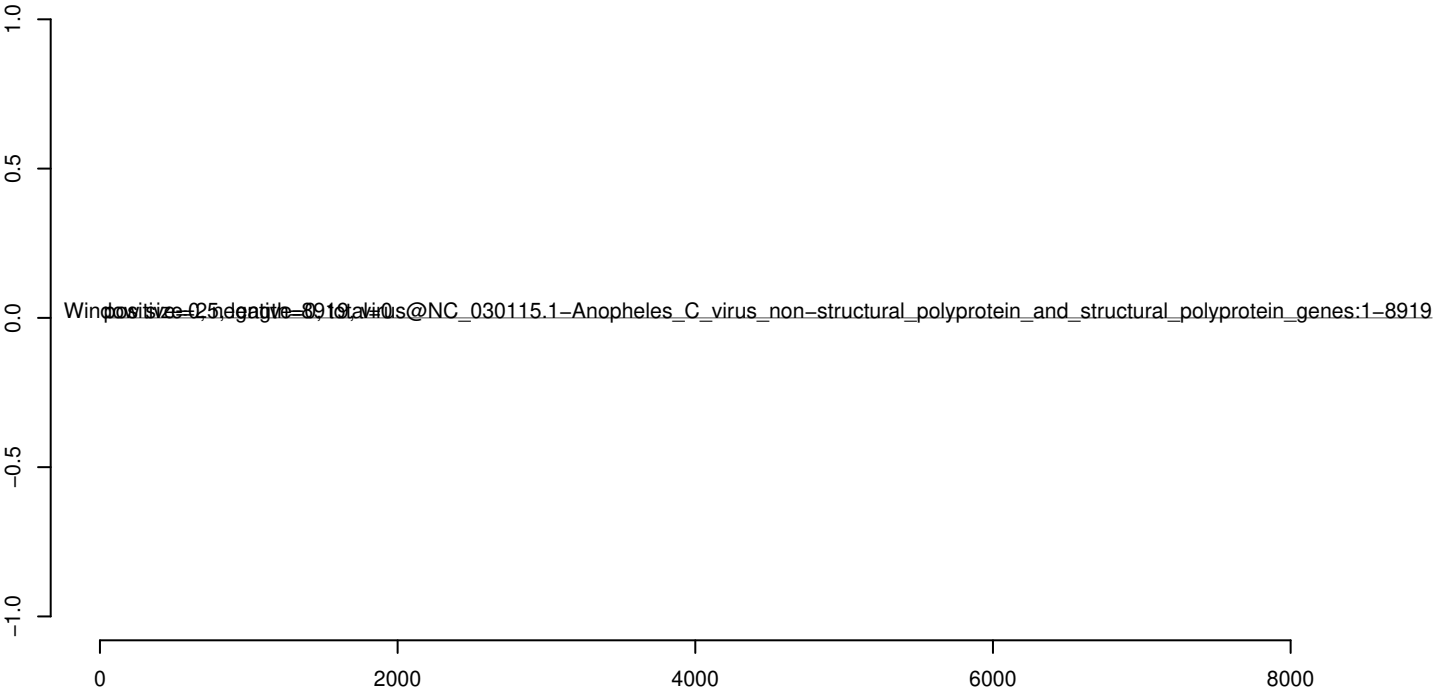
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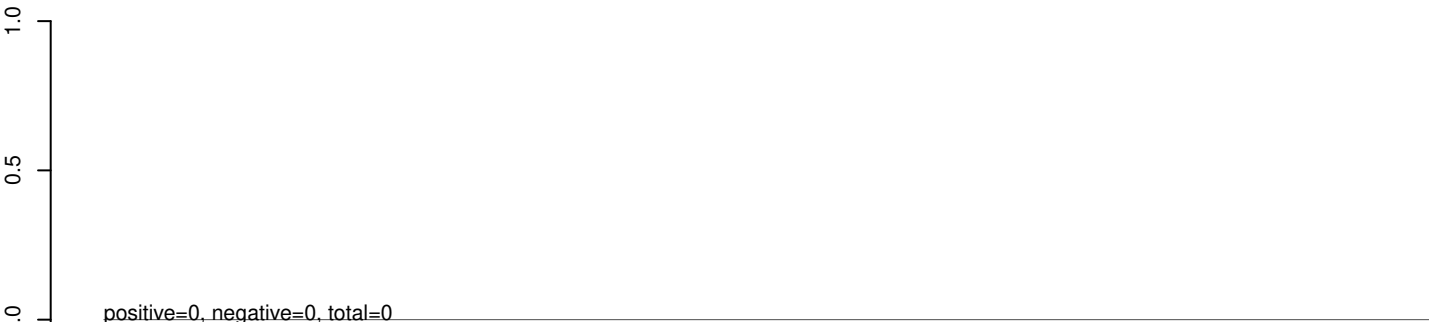
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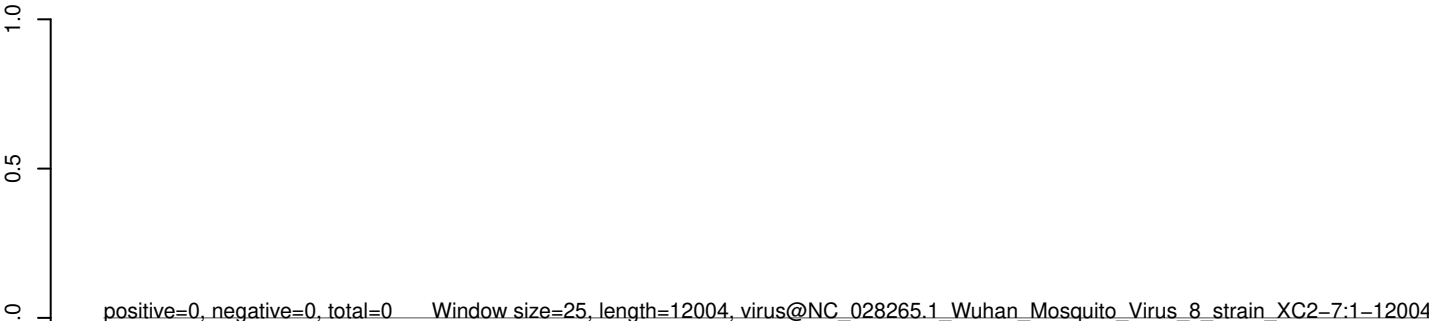
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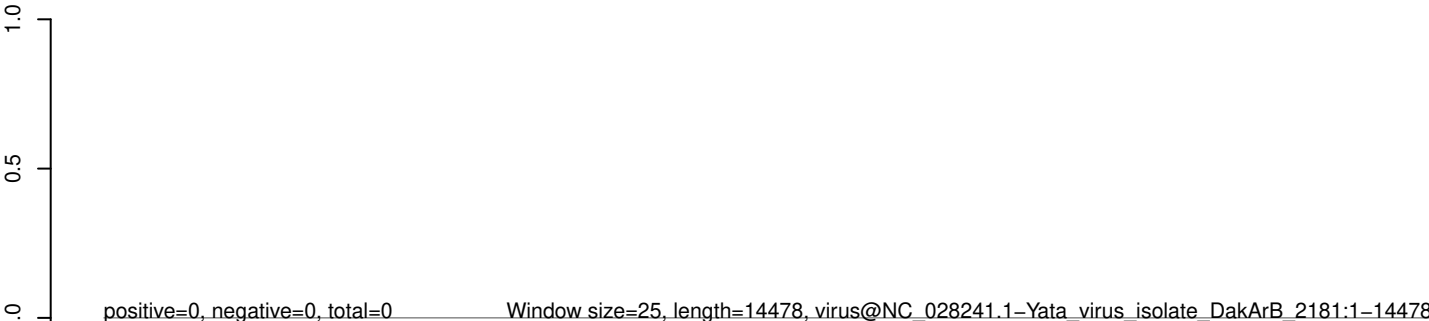
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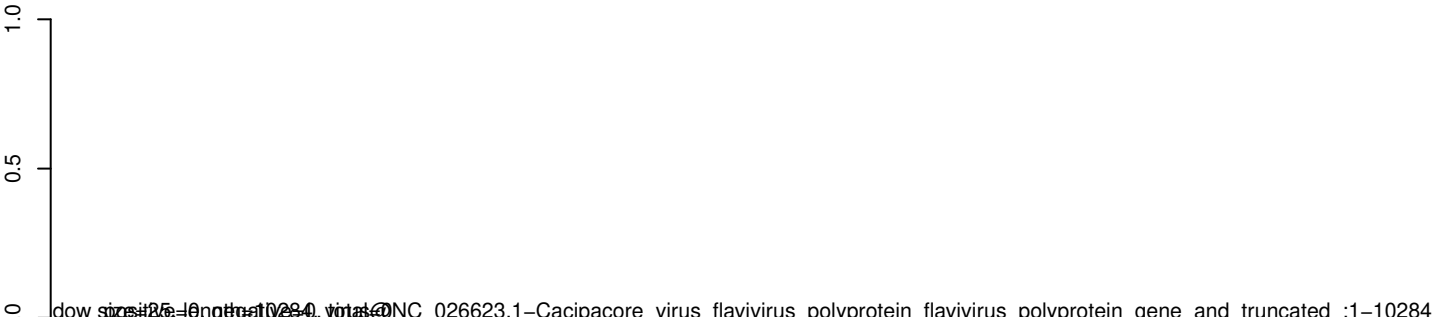
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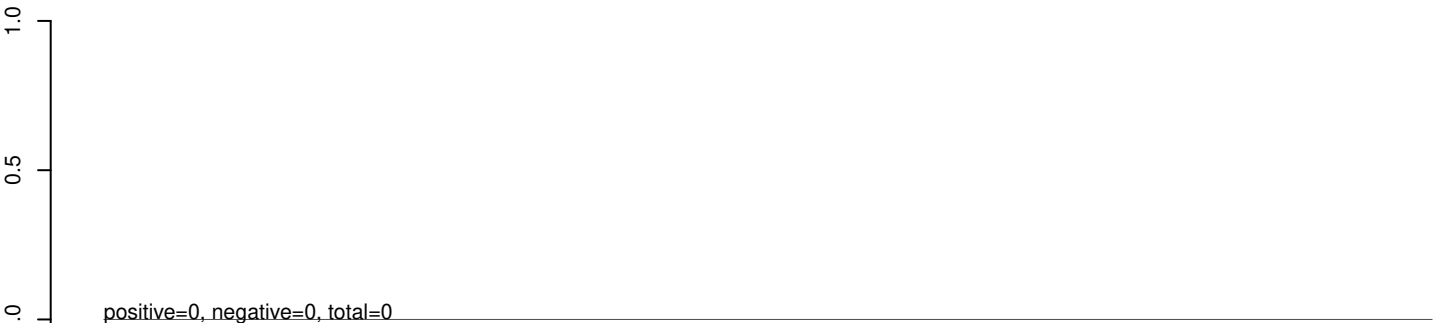


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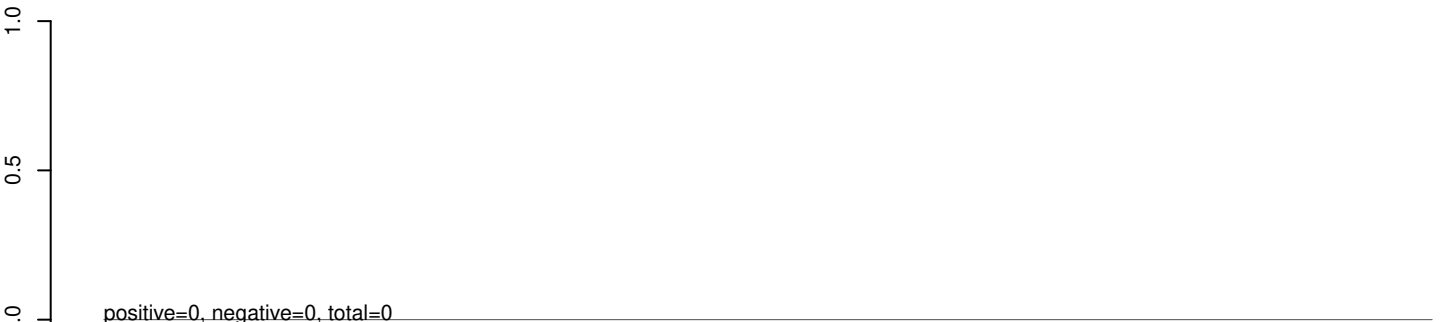


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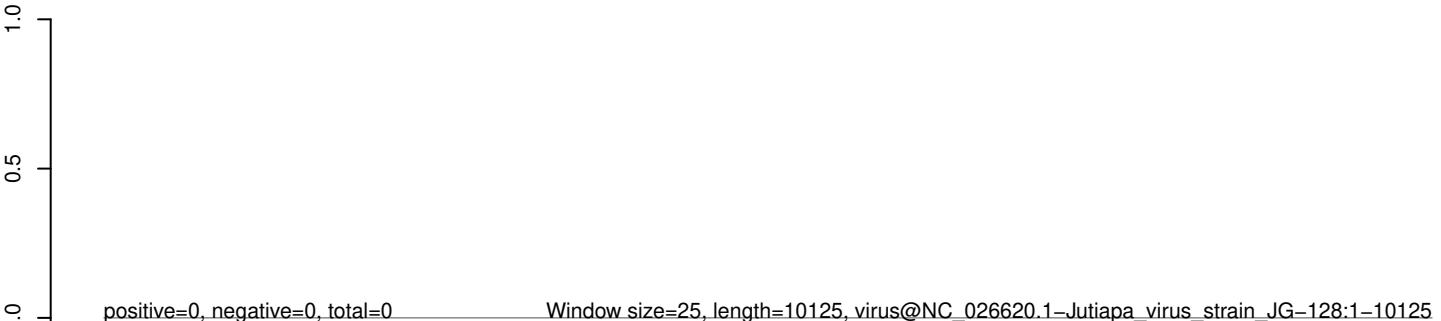
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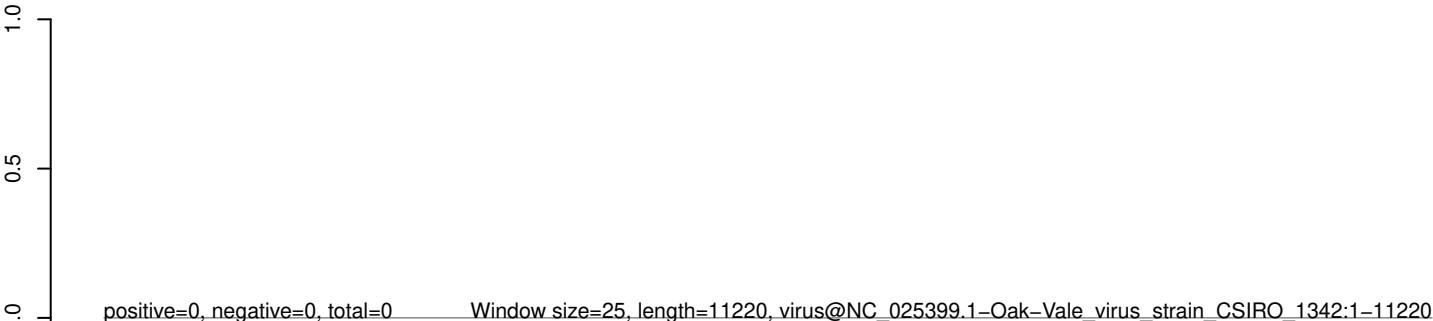
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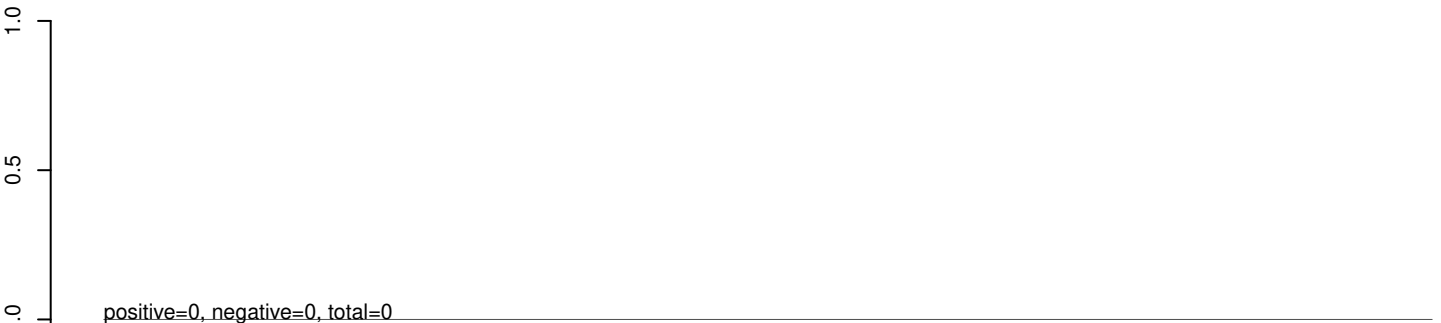


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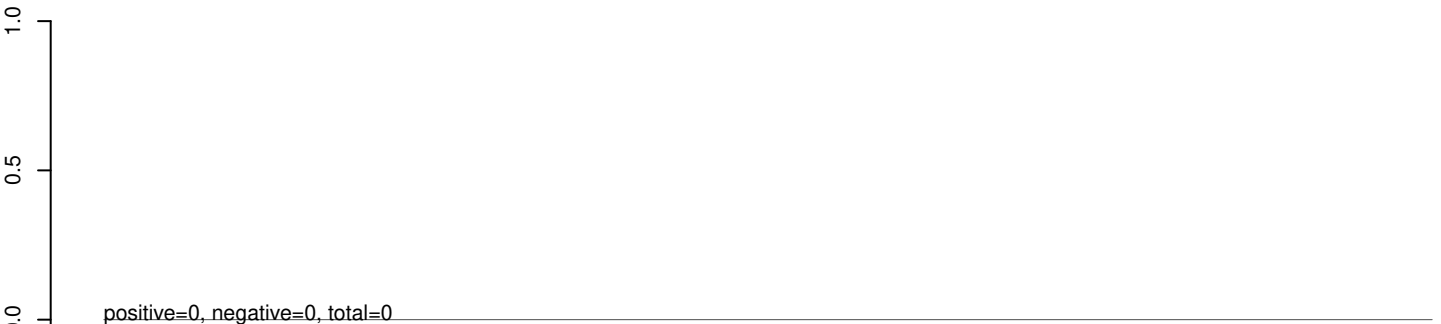


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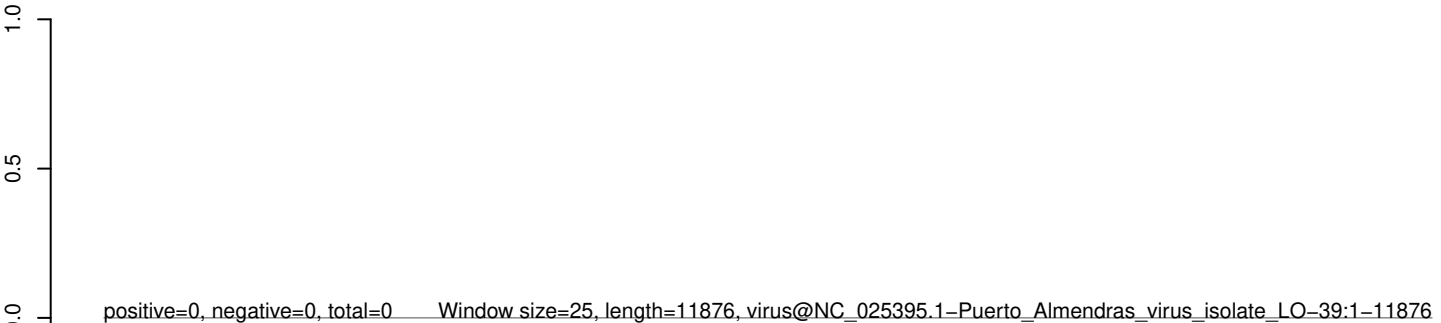
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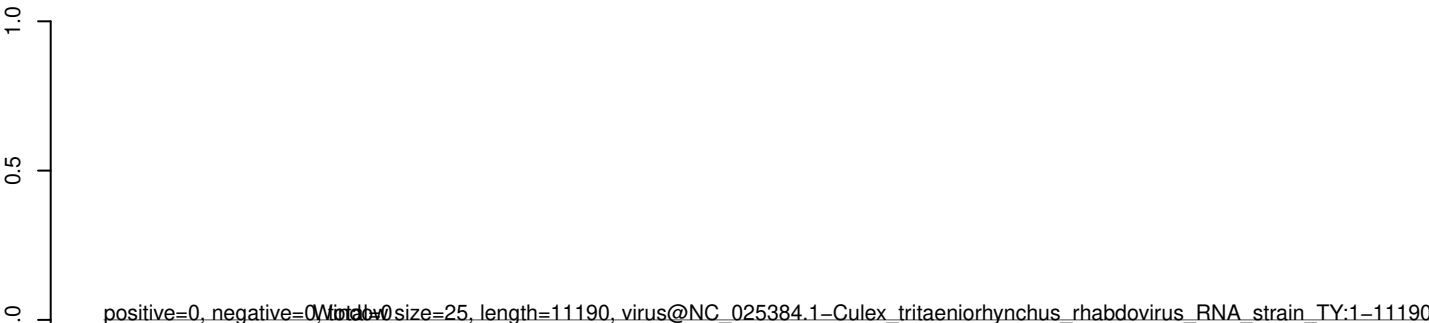
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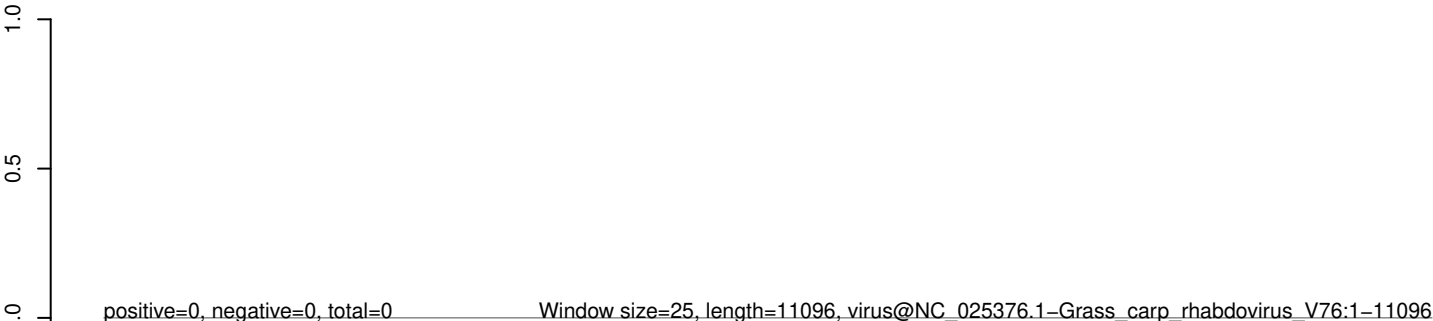
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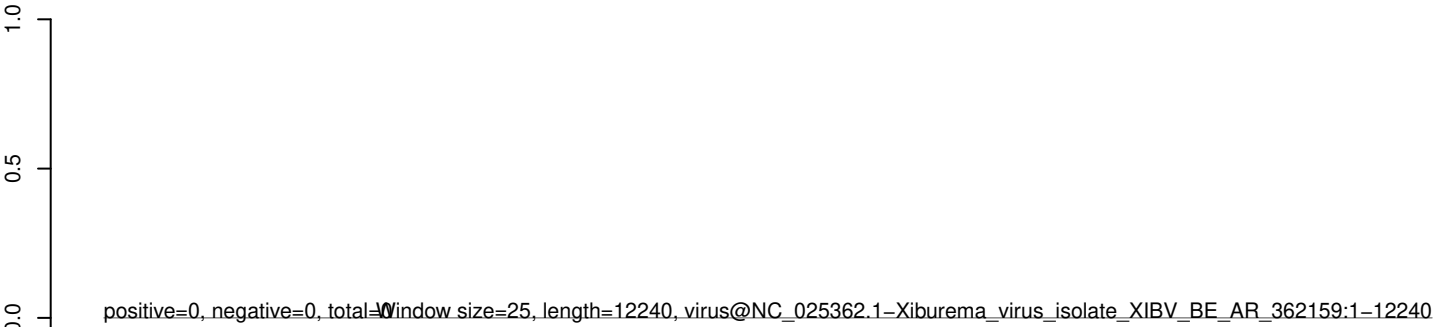
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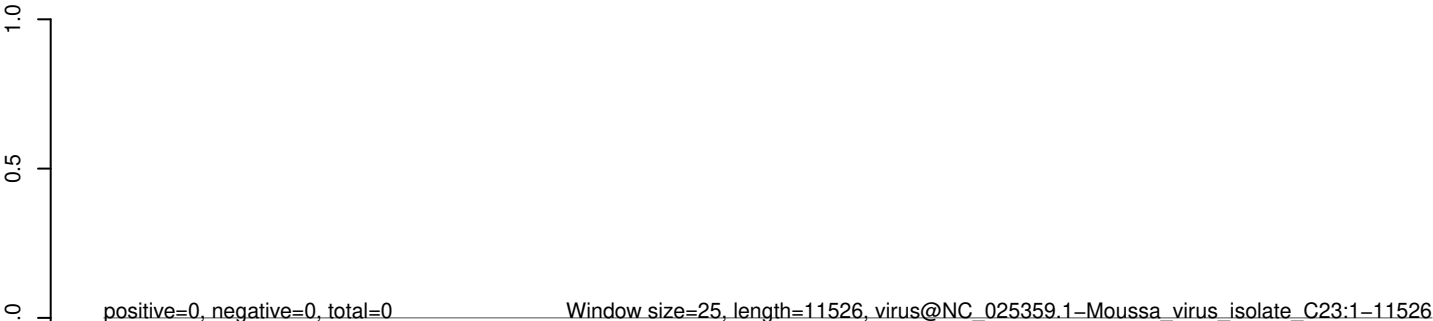
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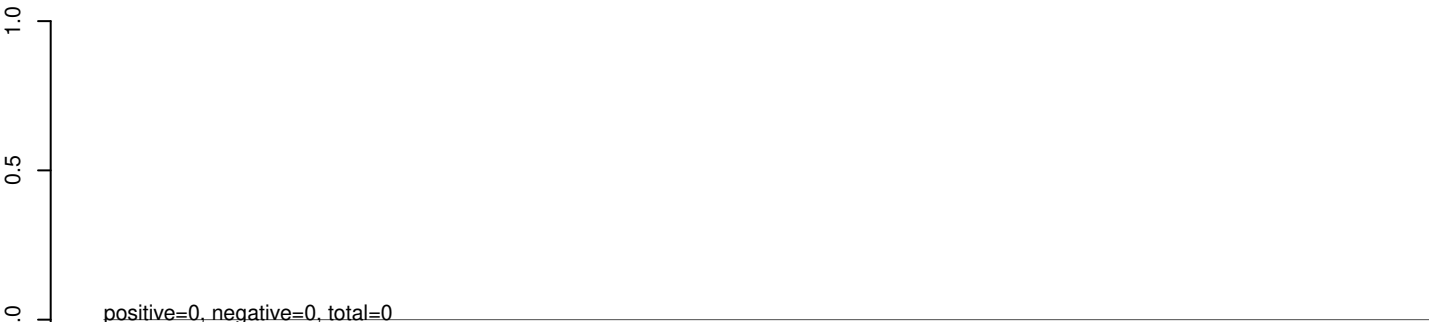


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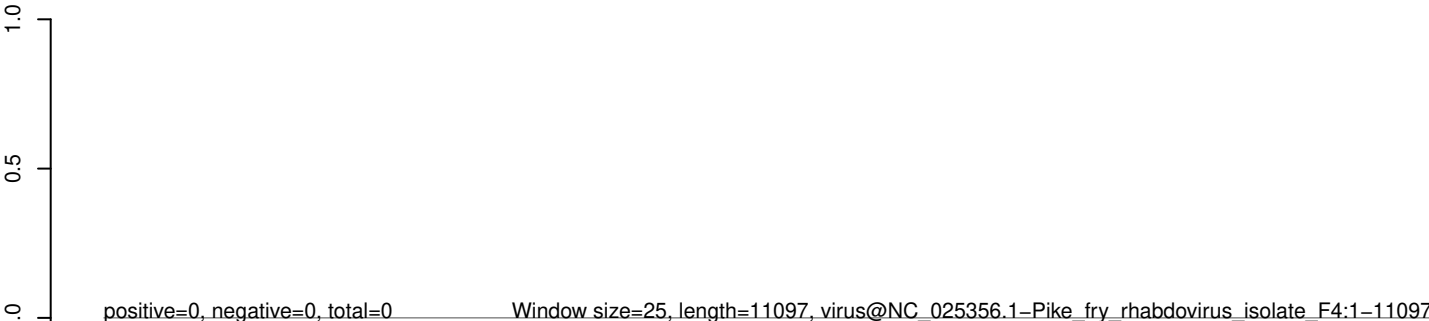
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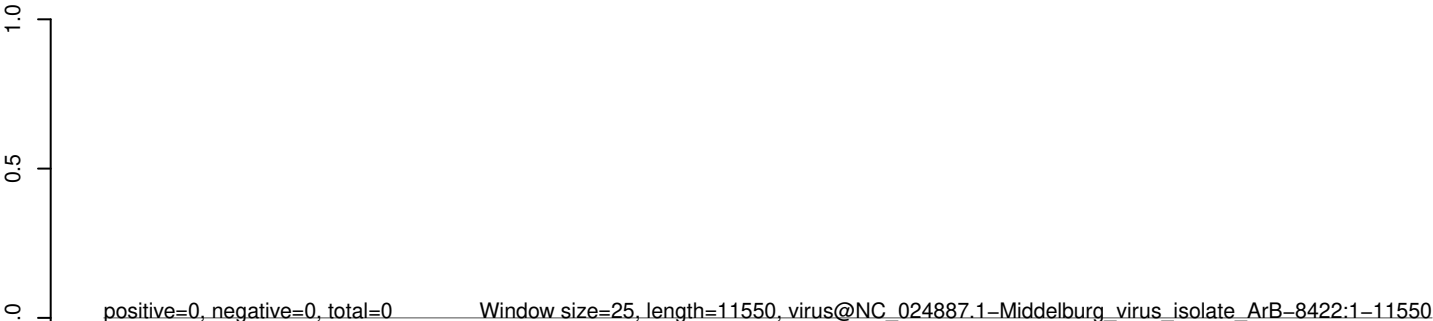
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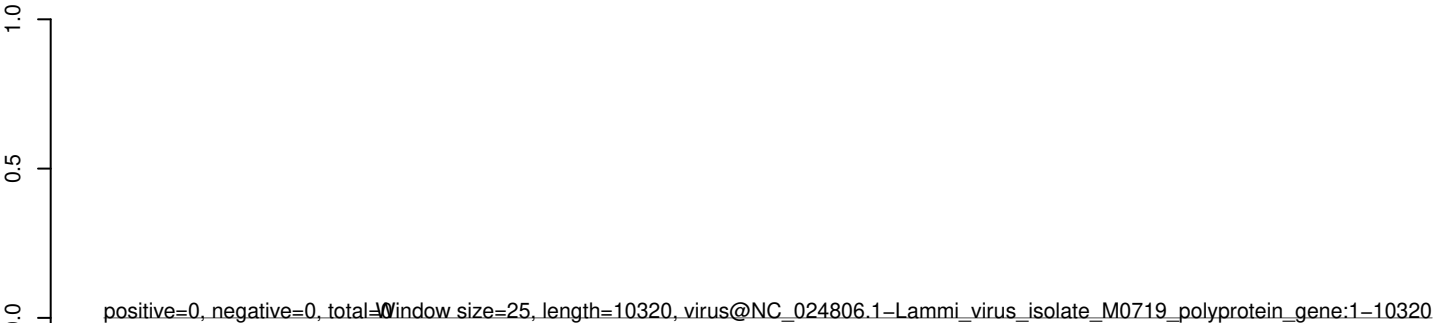
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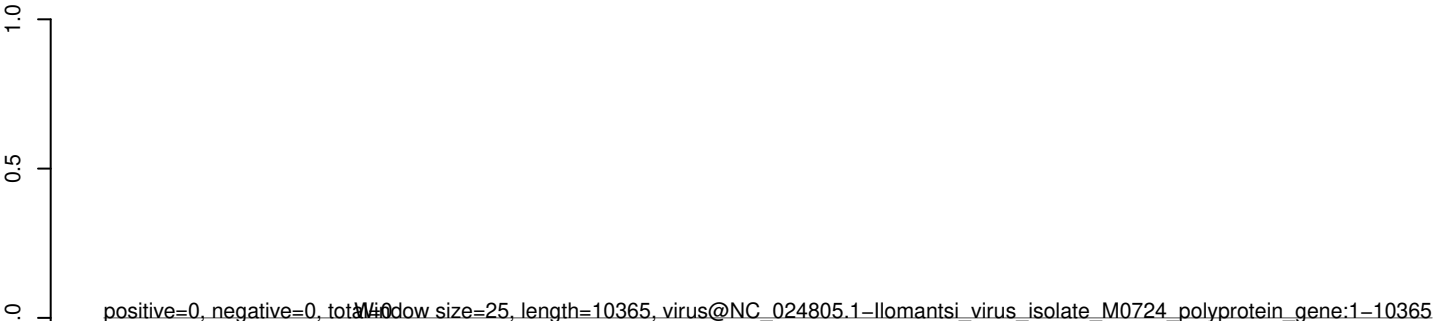
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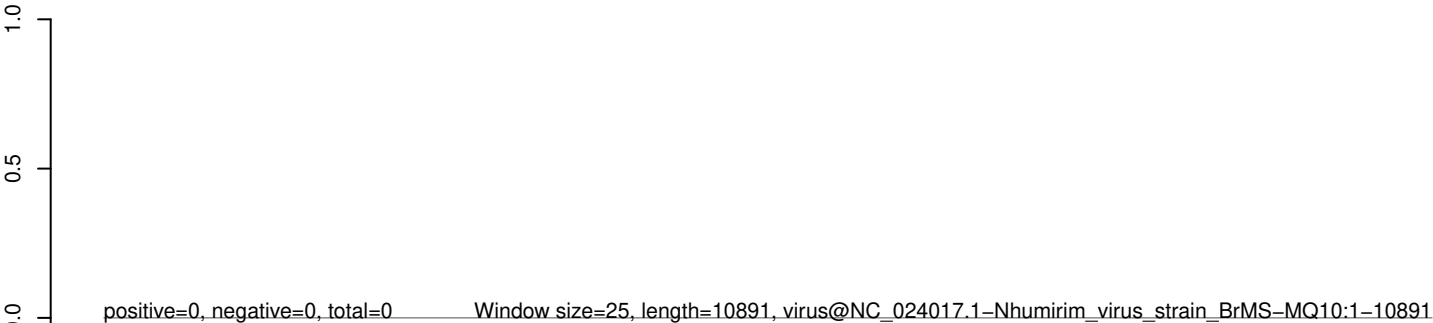
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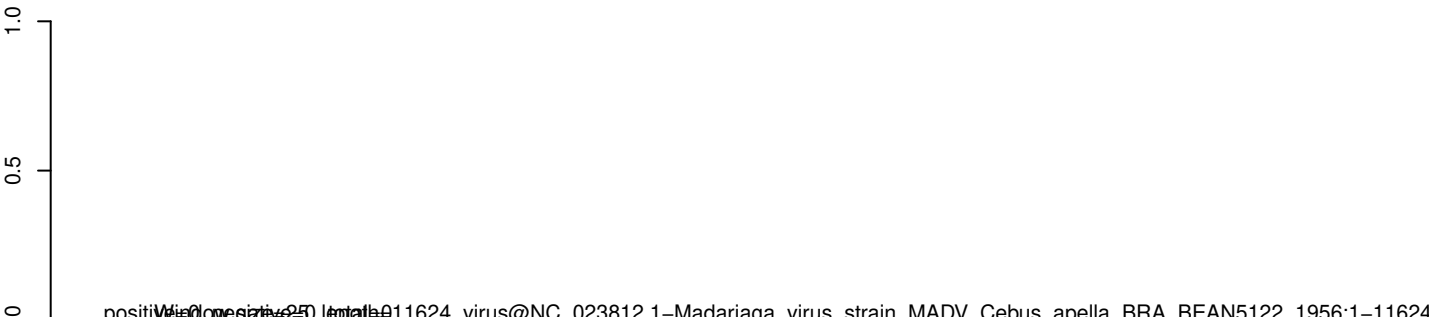
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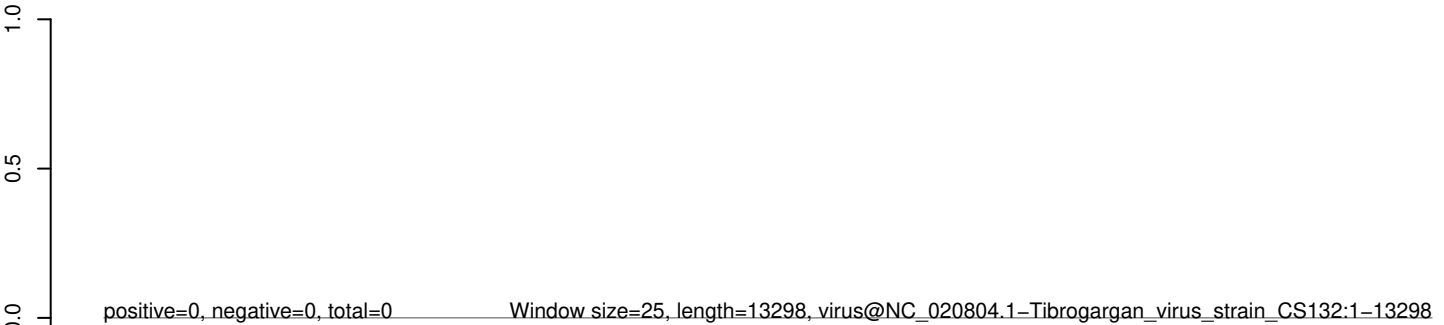
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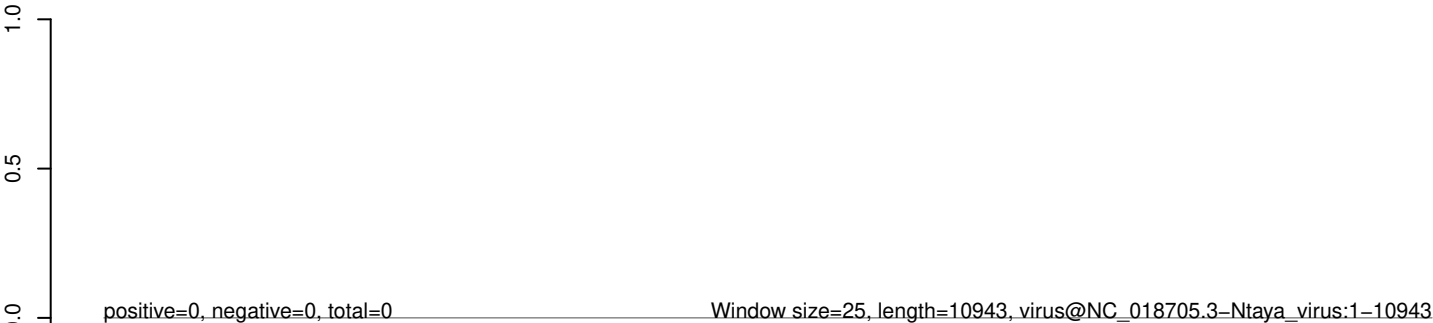
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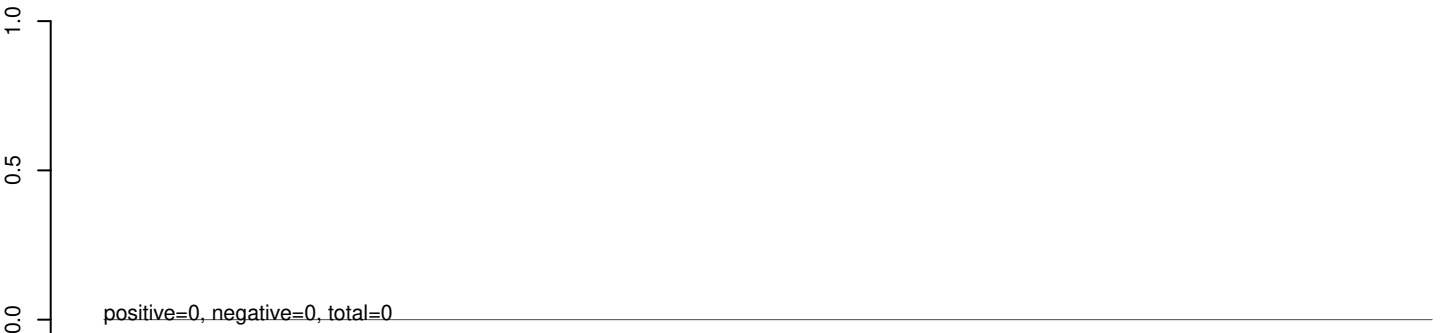
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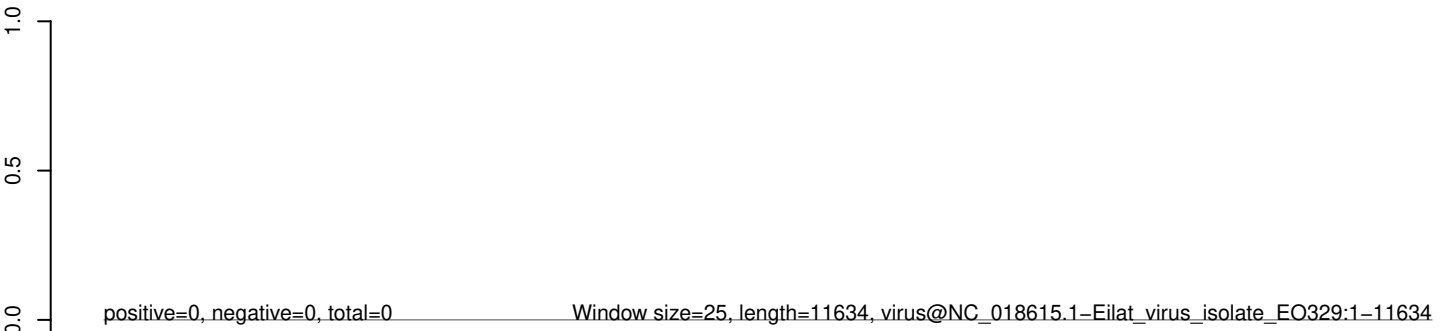
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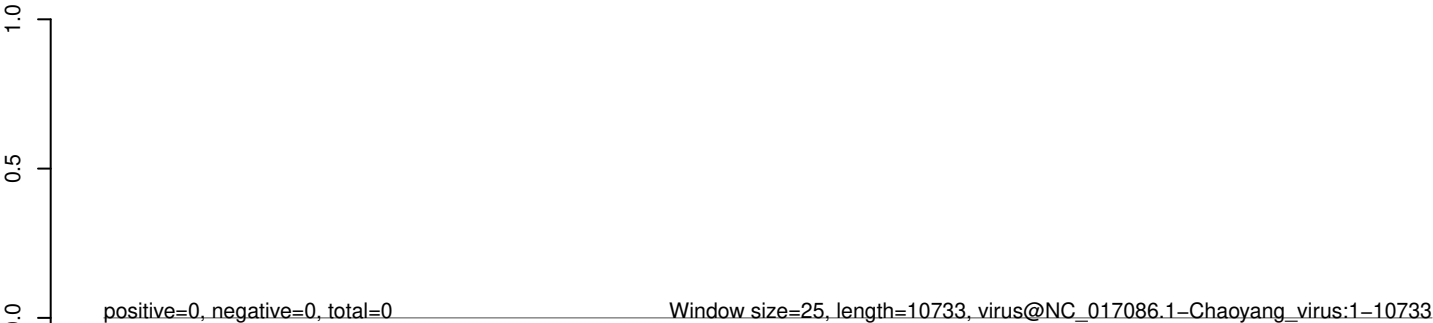
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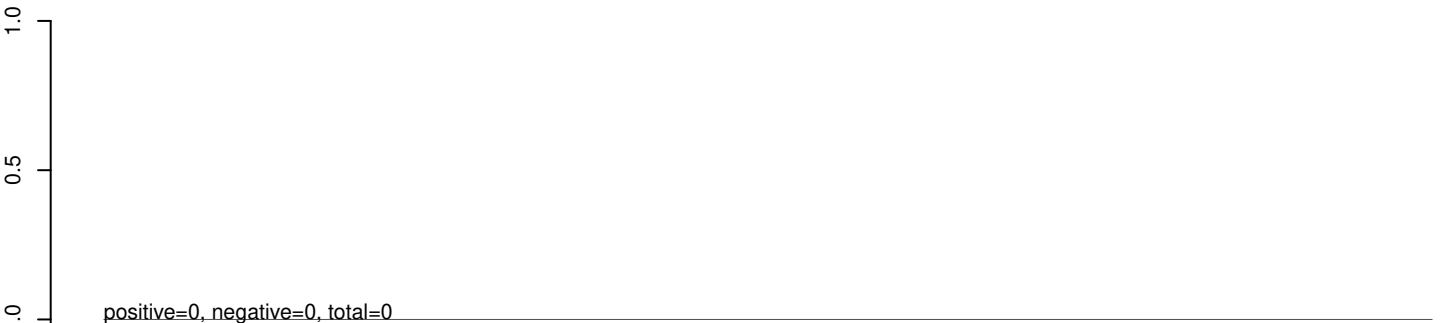


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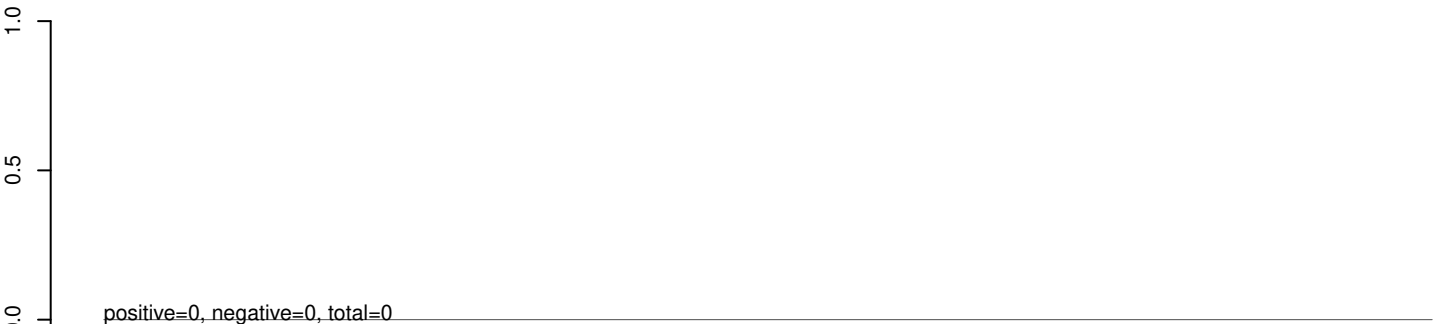


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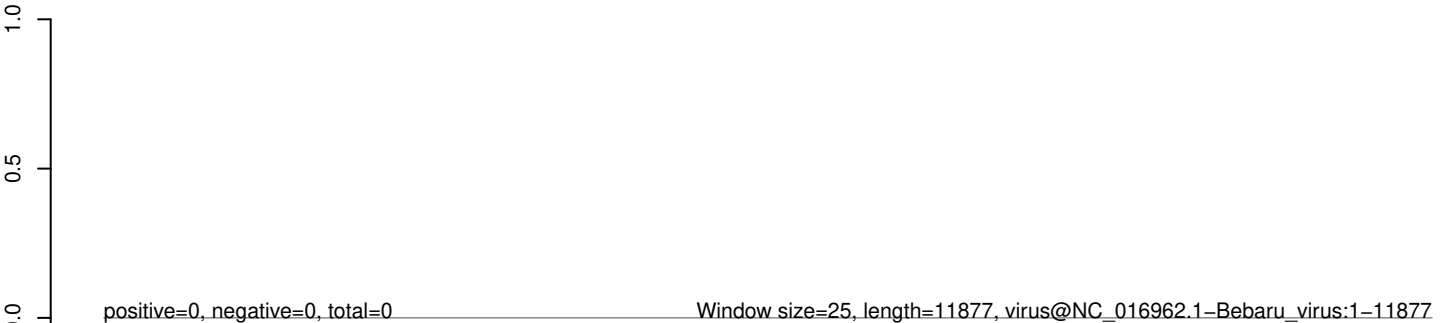
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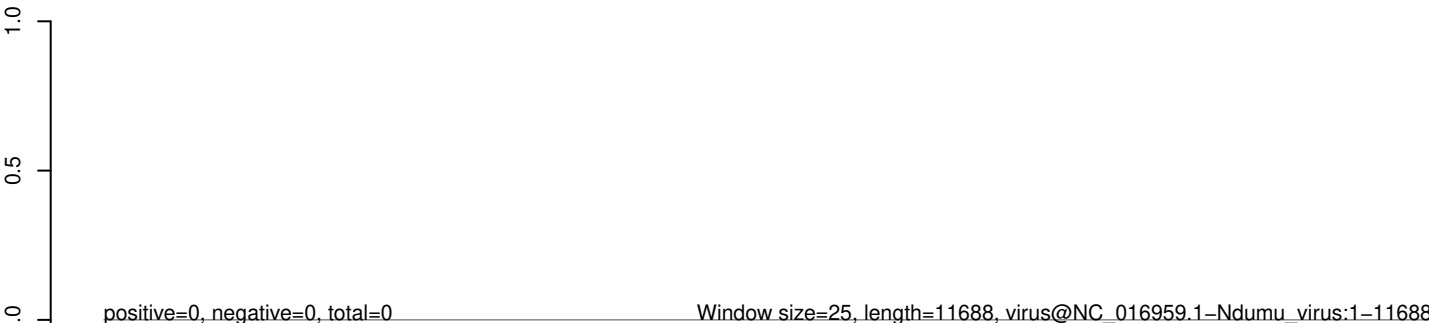
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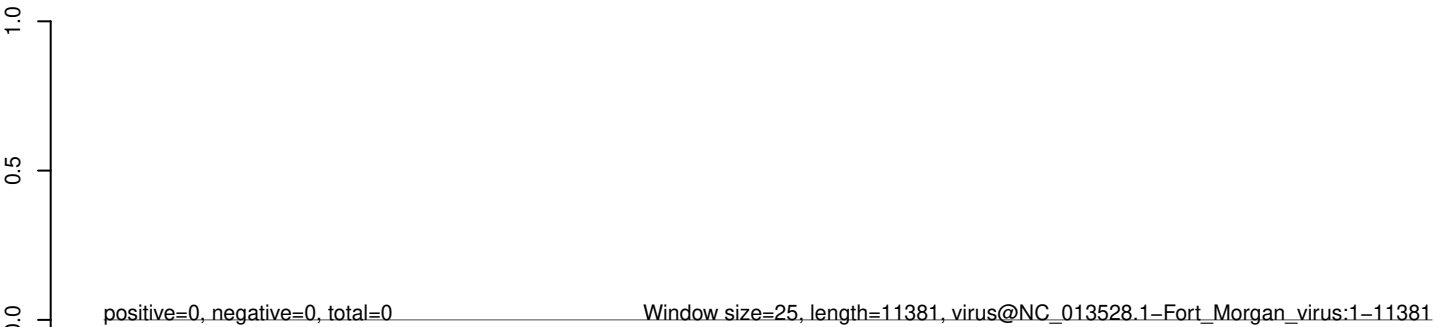
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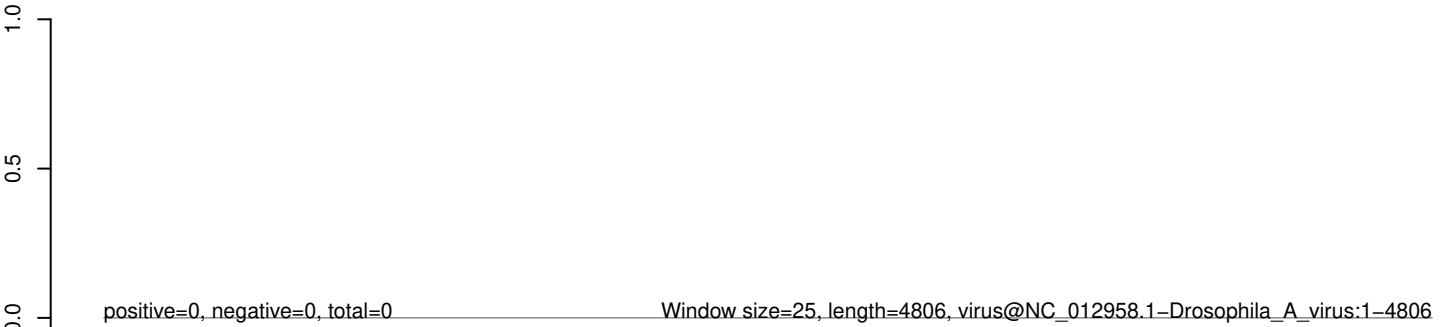
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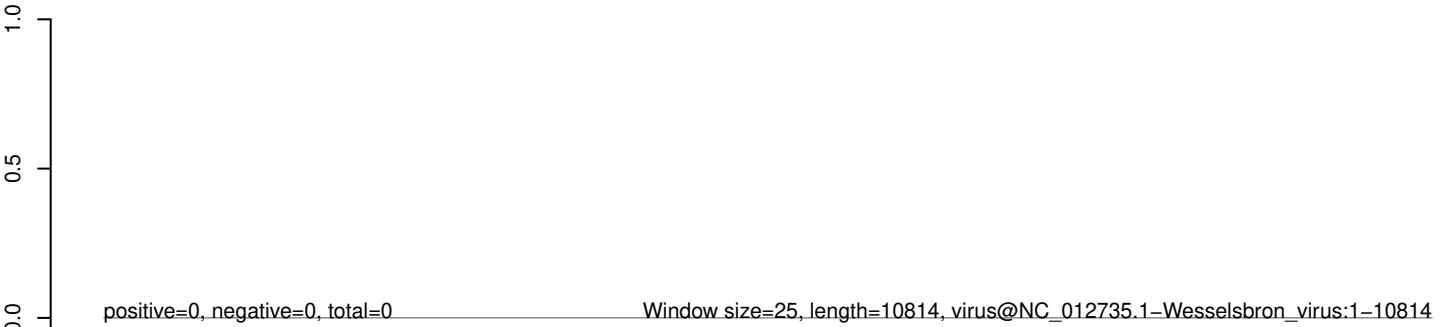
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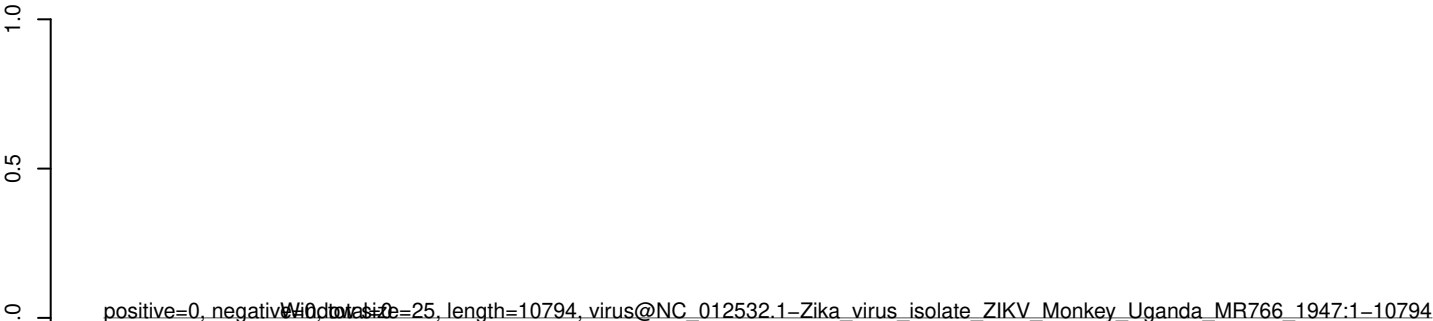
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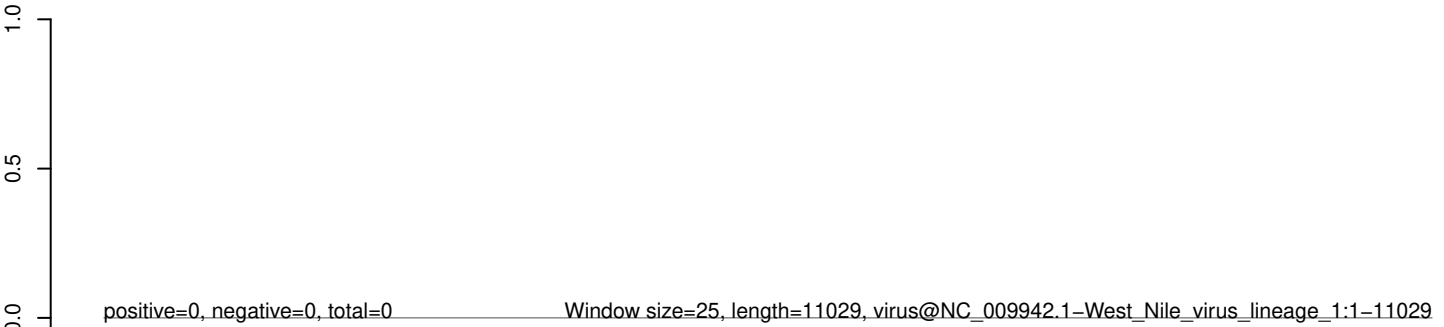
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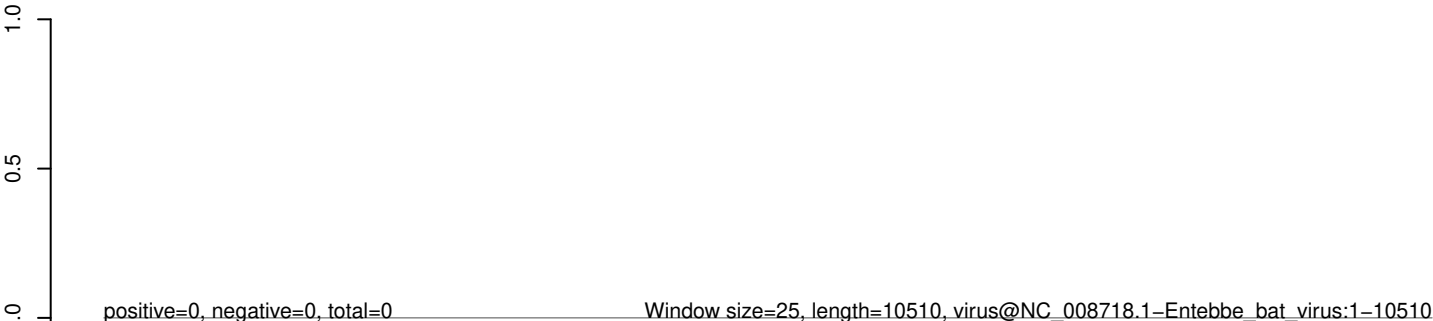
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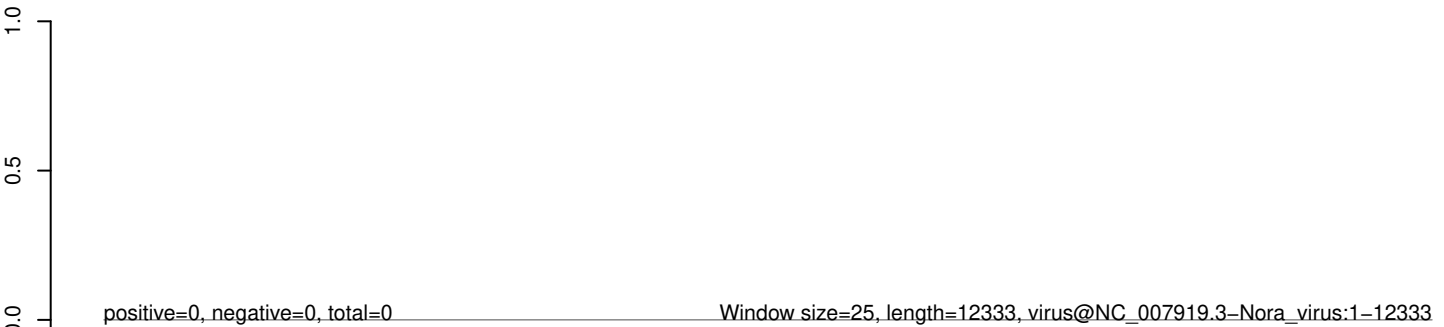
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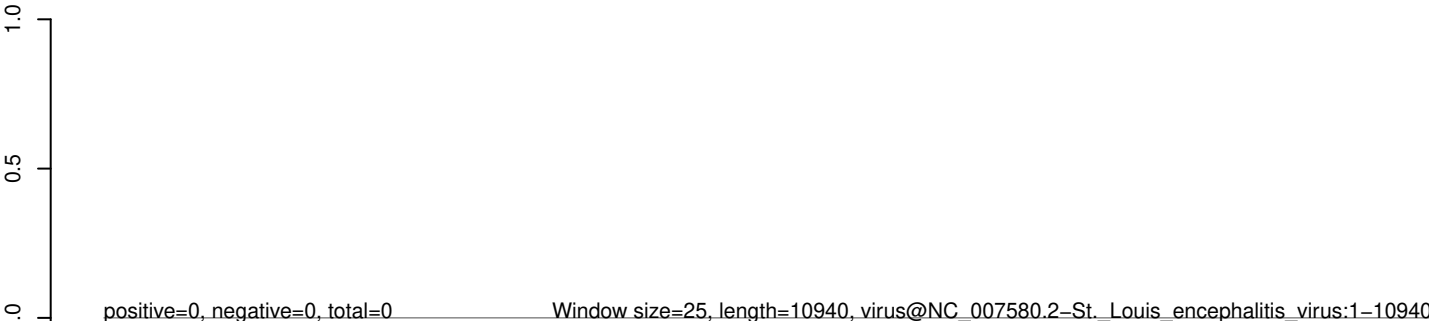
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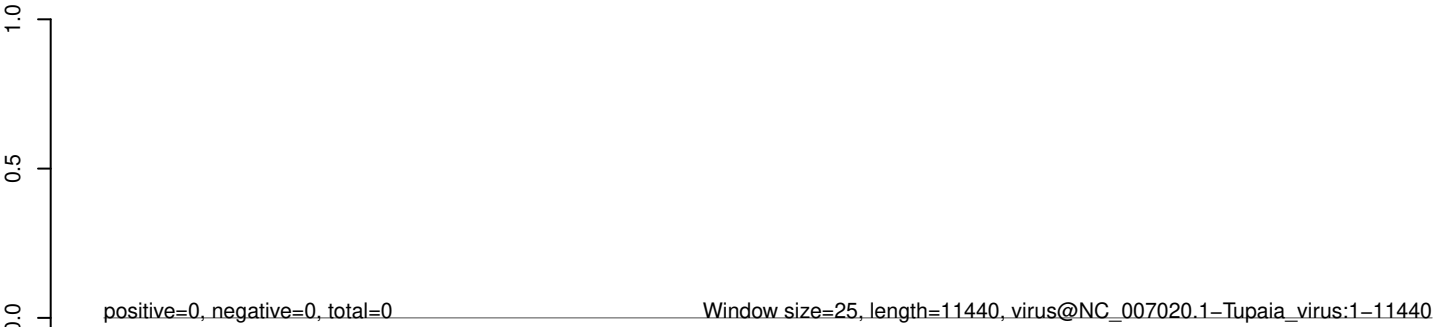
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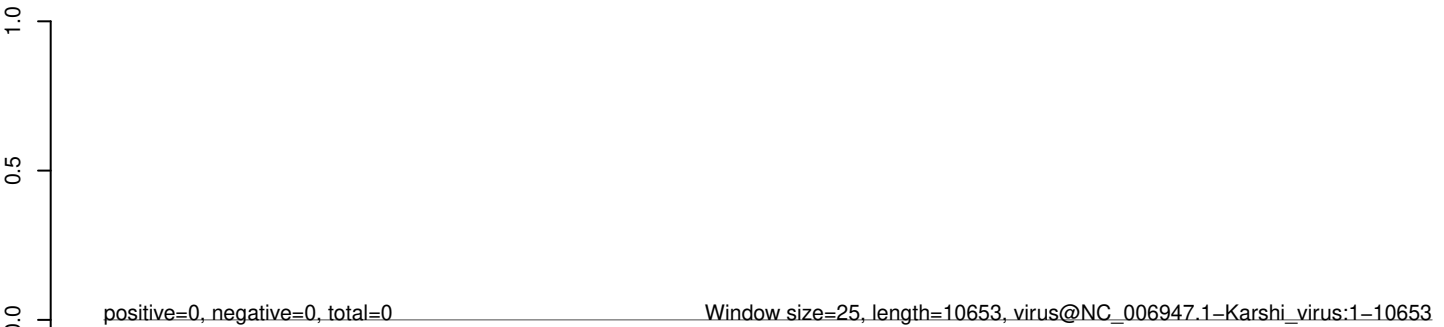
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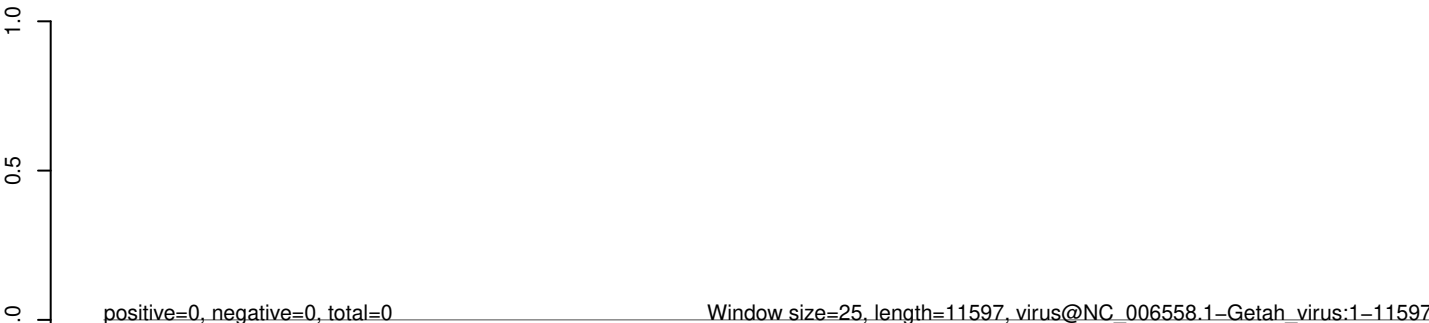
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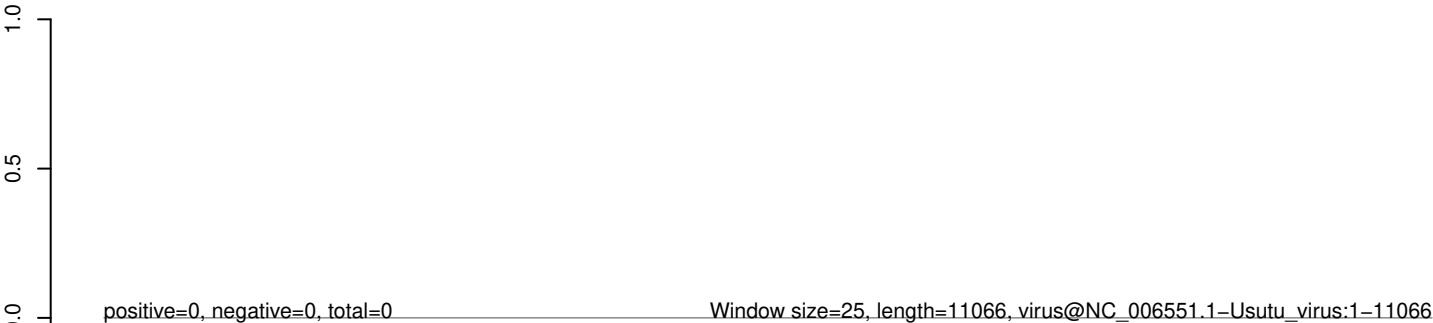
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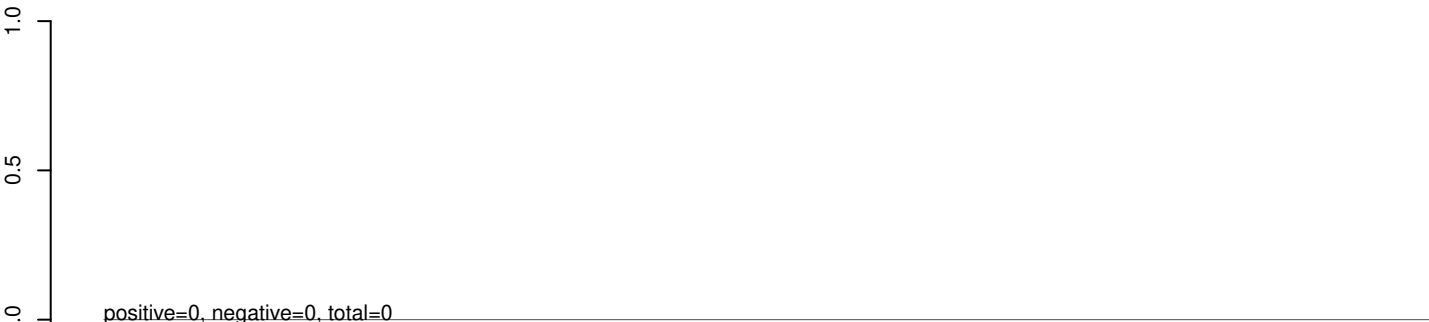


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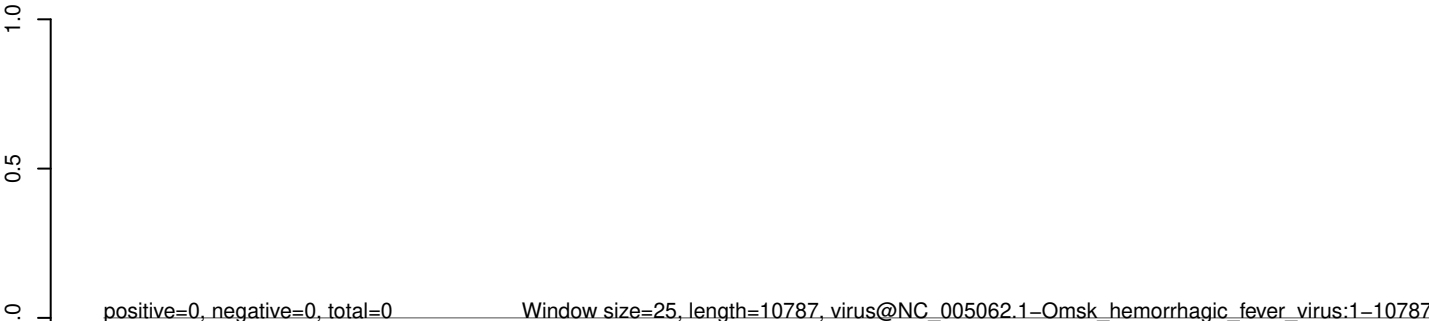
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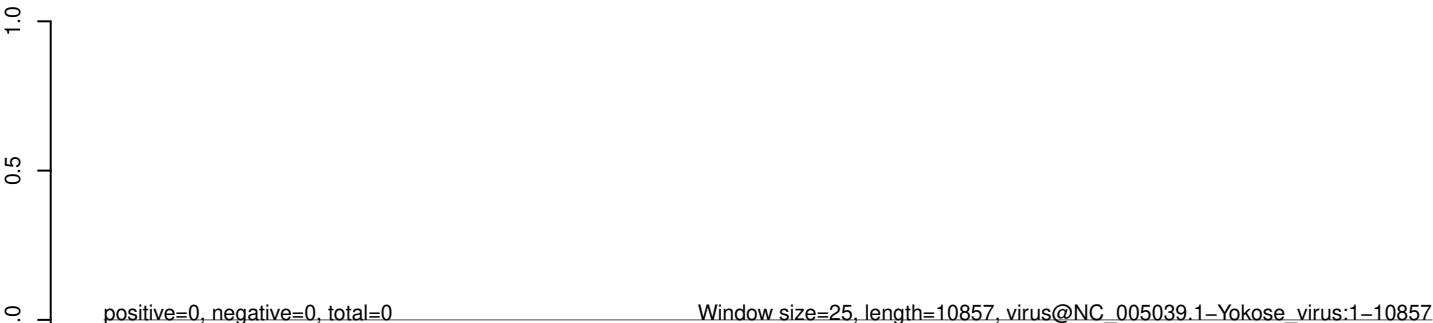
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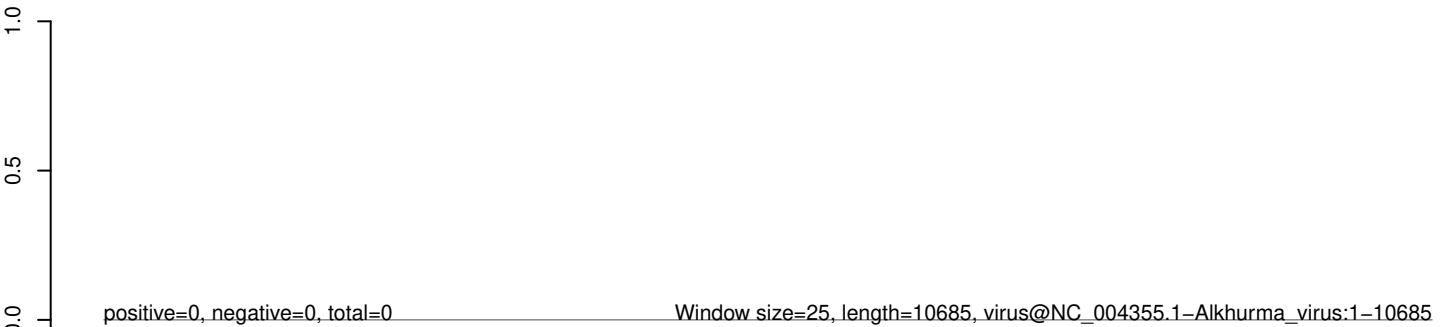
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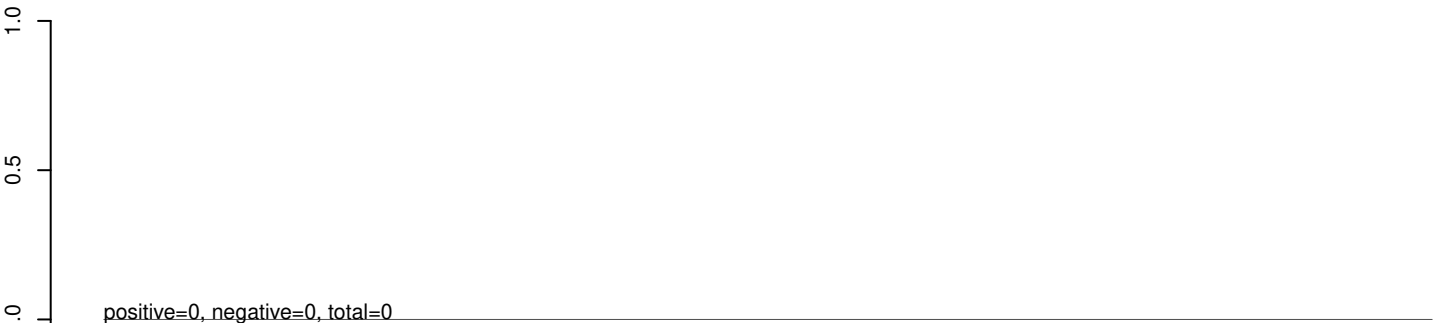


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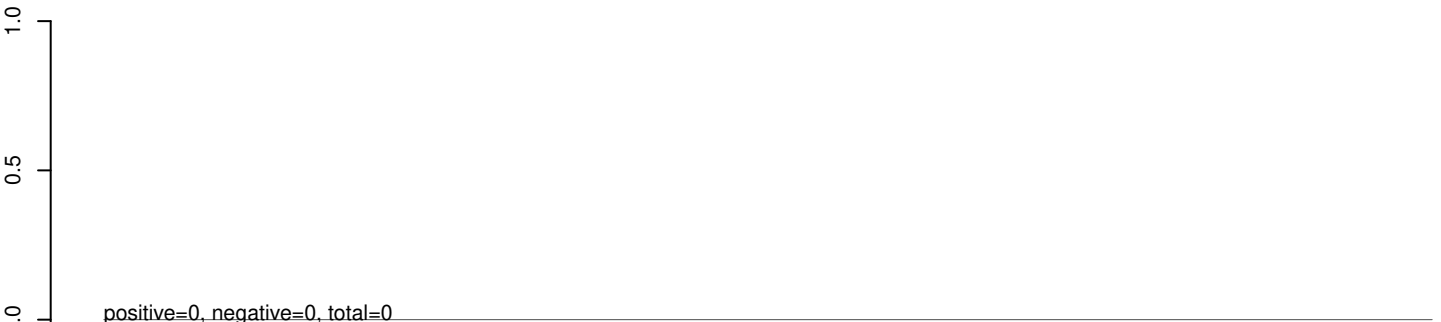


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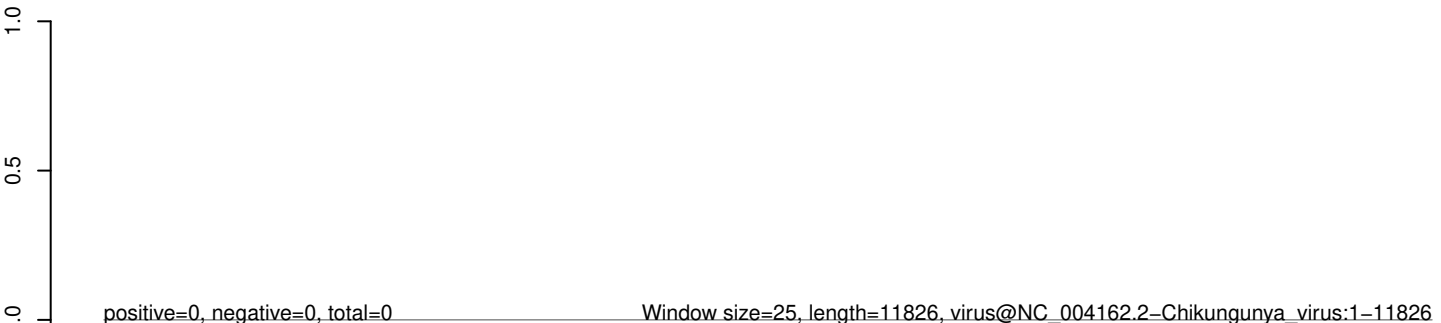
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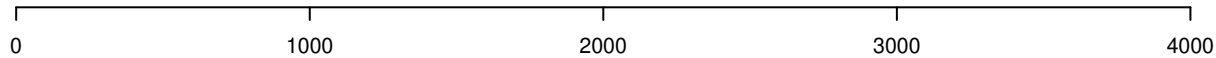
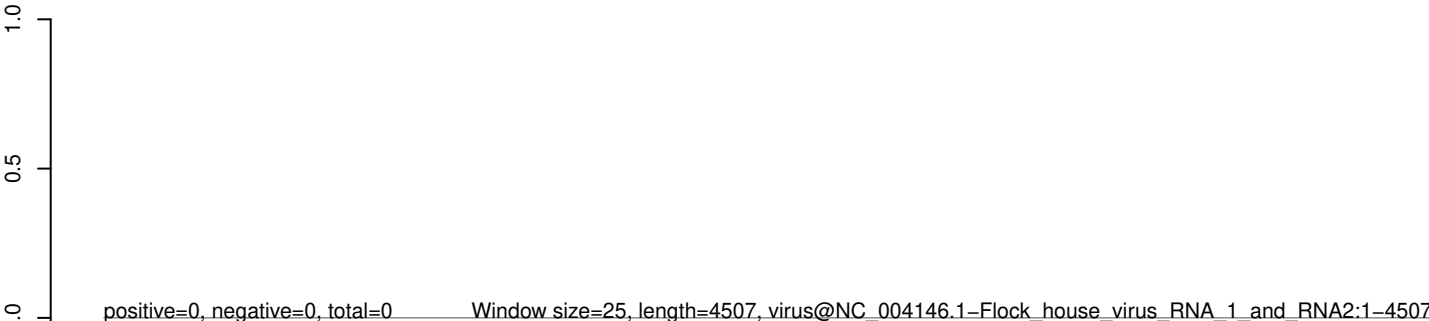
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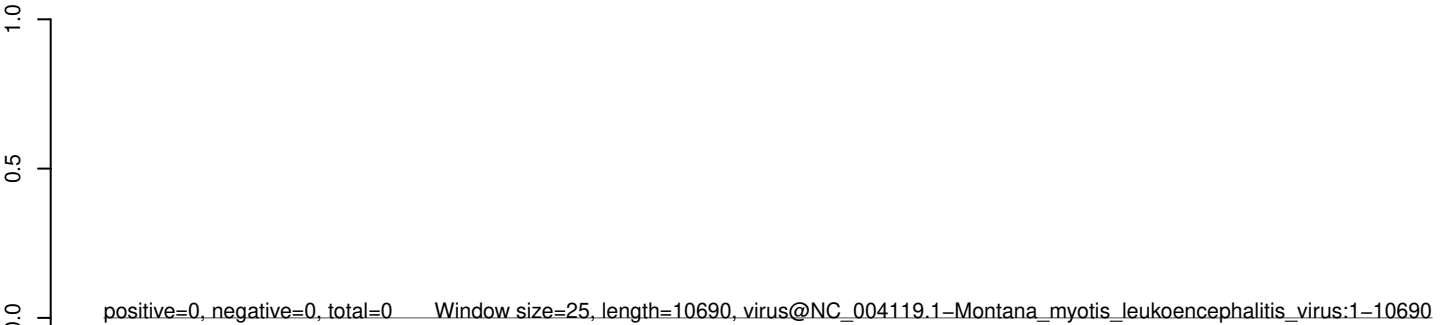
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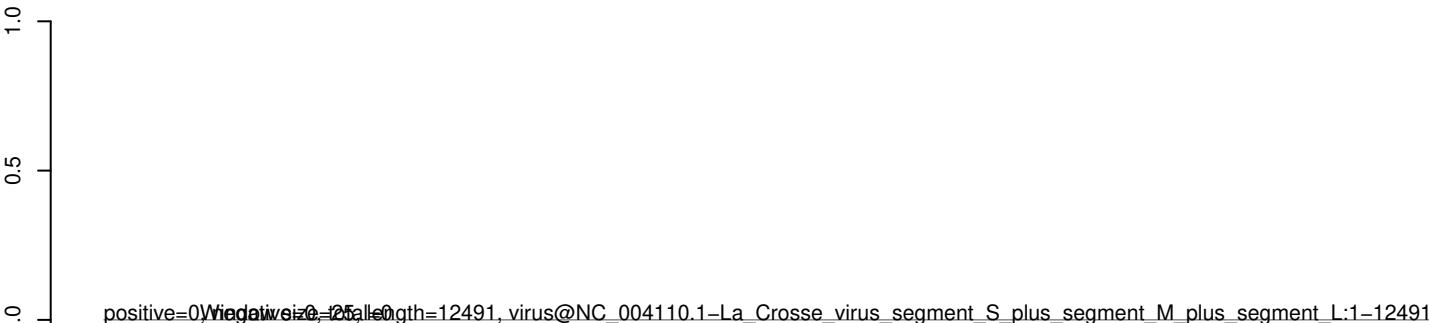
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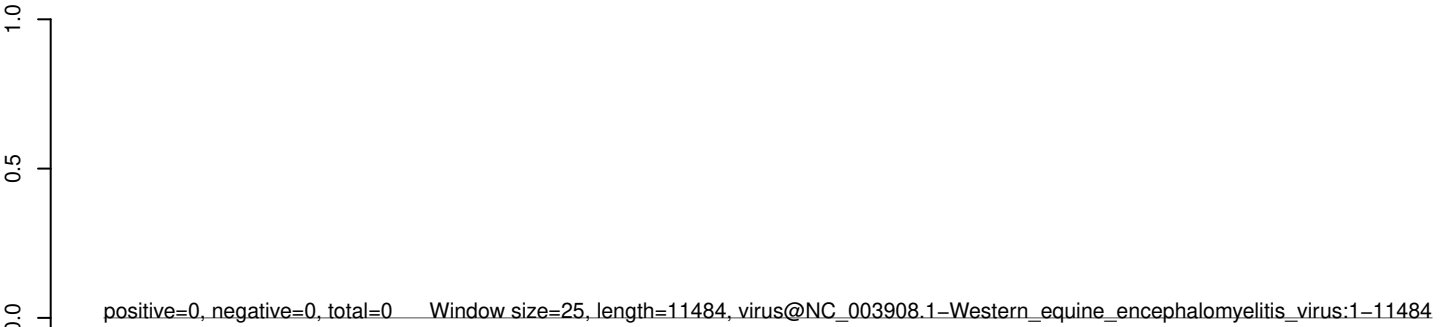
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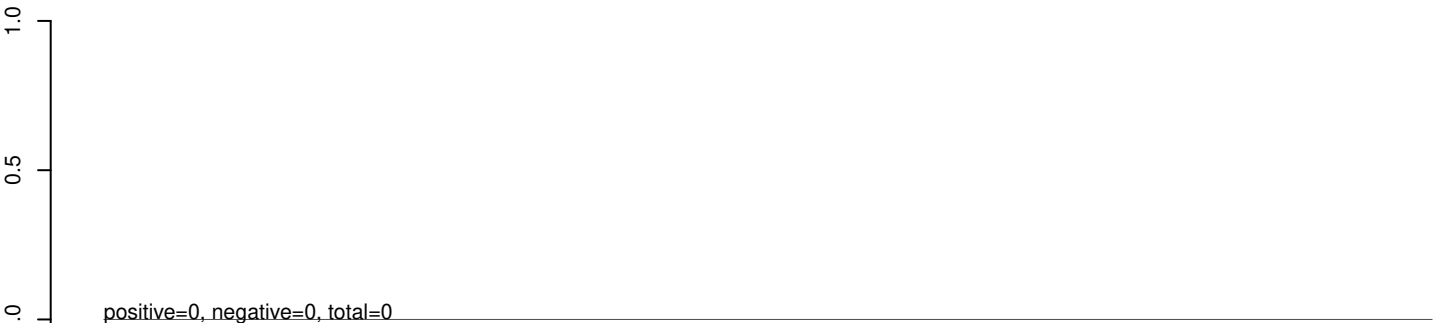


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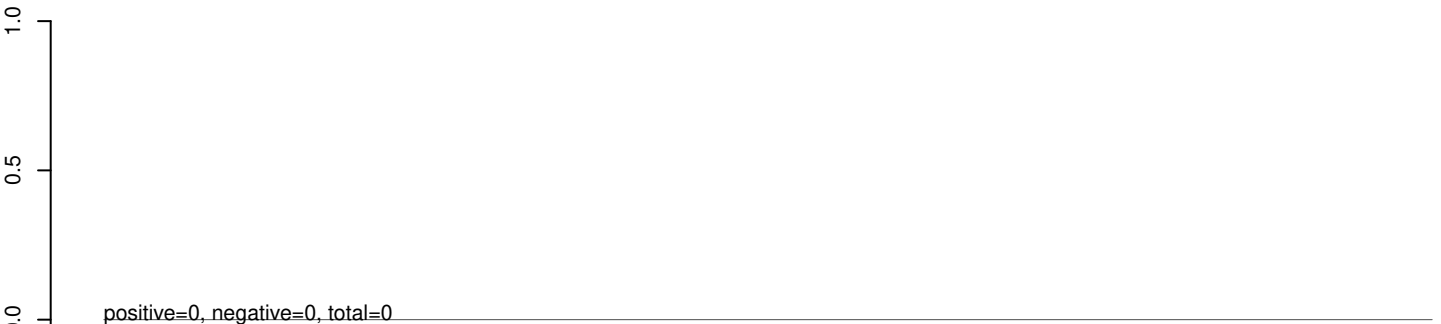


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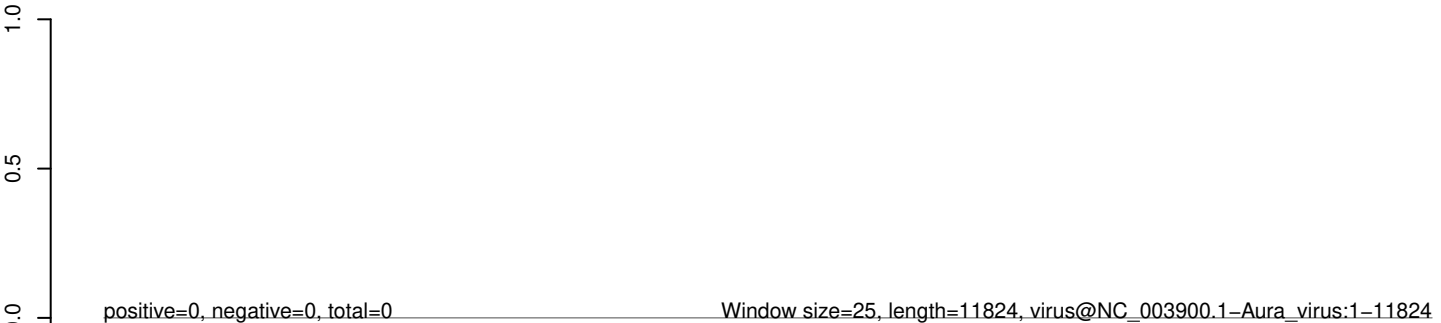
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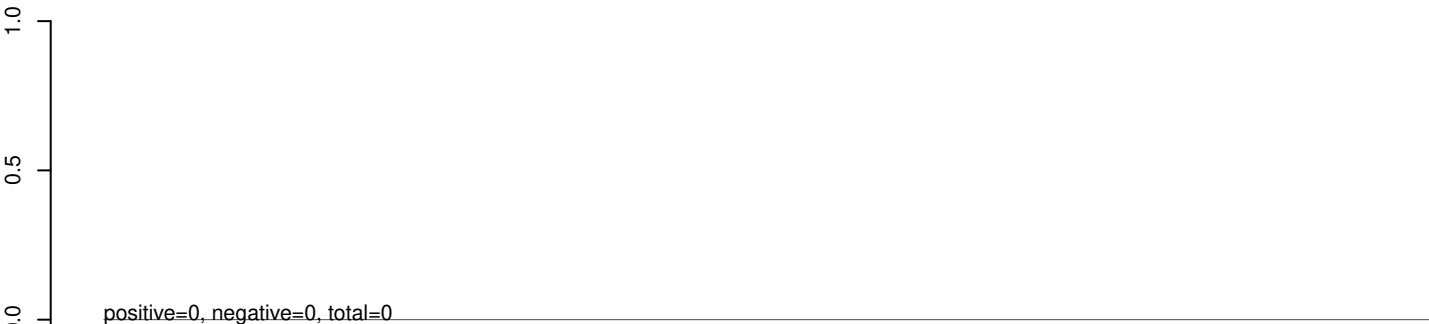


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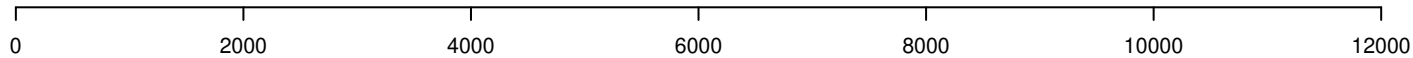
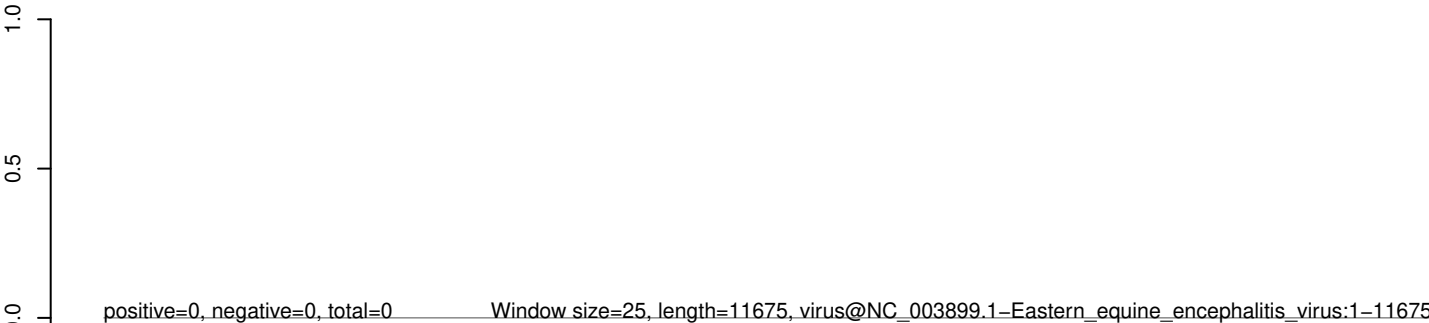
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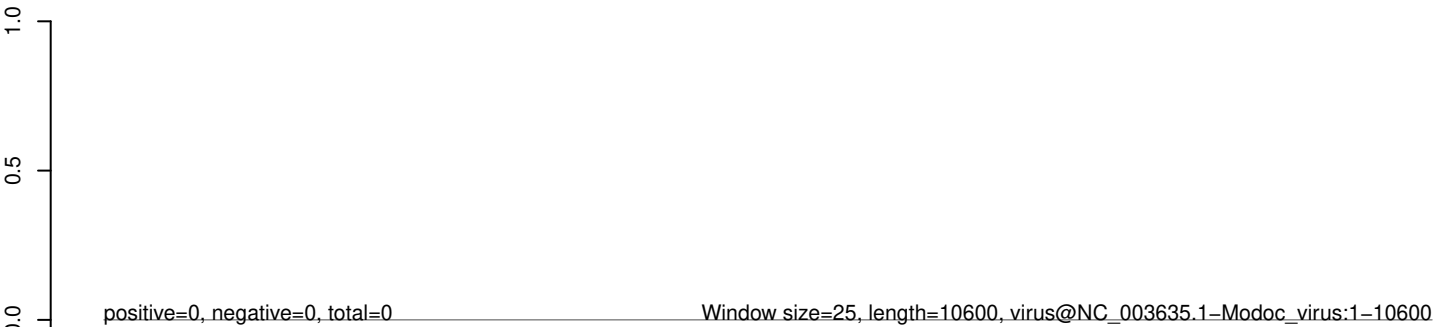
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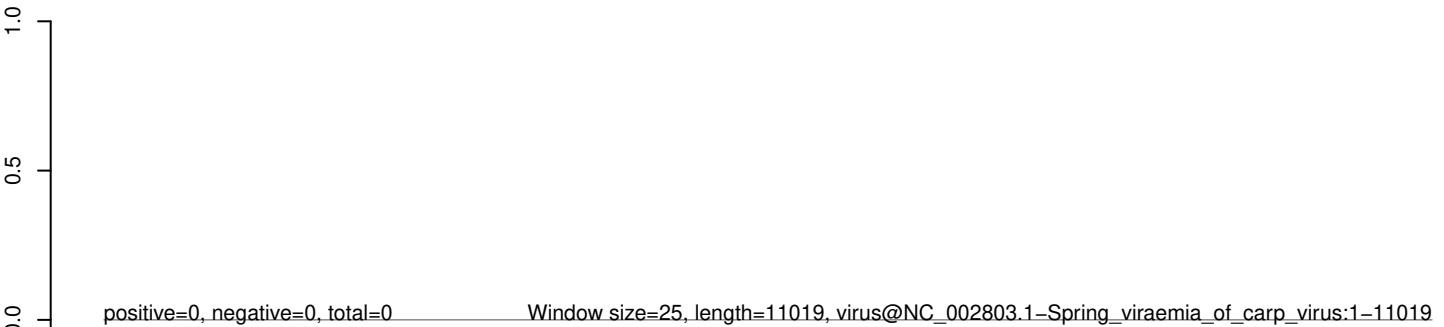
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AeAeg_Aag2_BTV2_AK.rep



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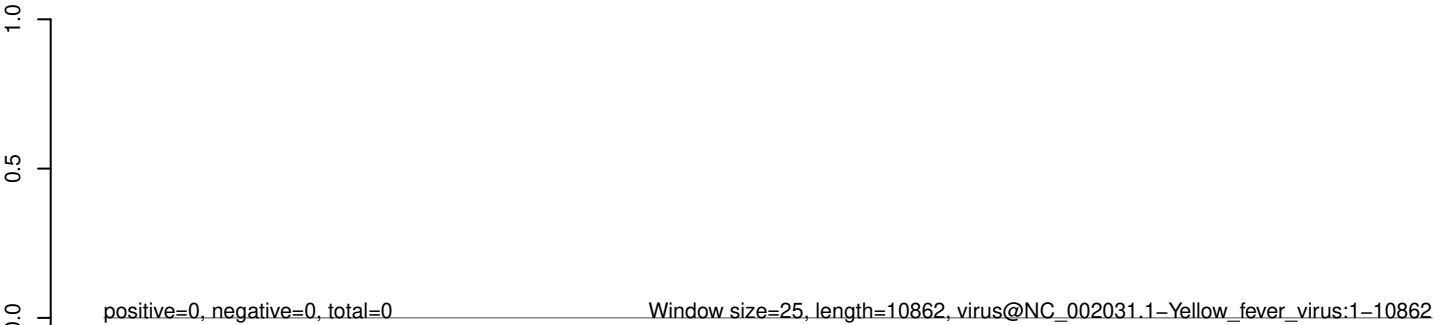
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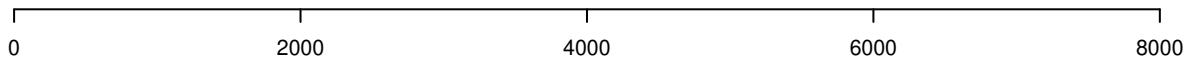
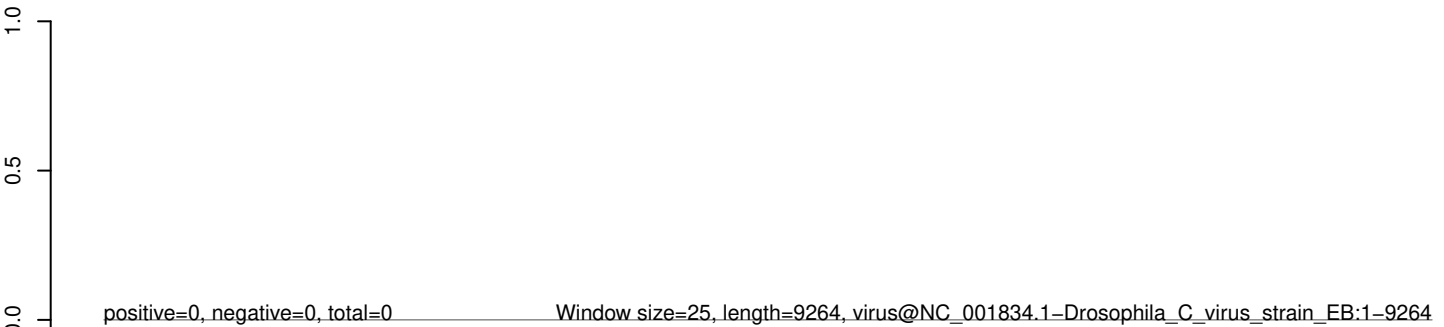
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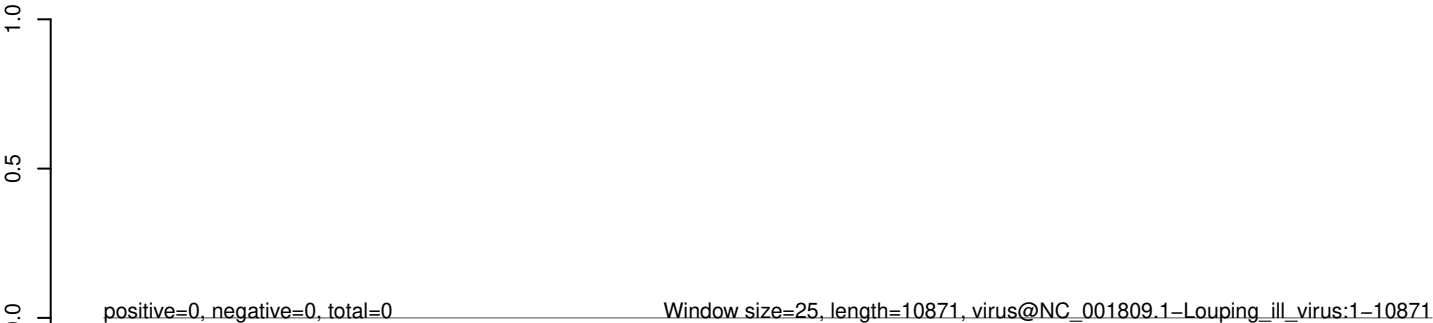
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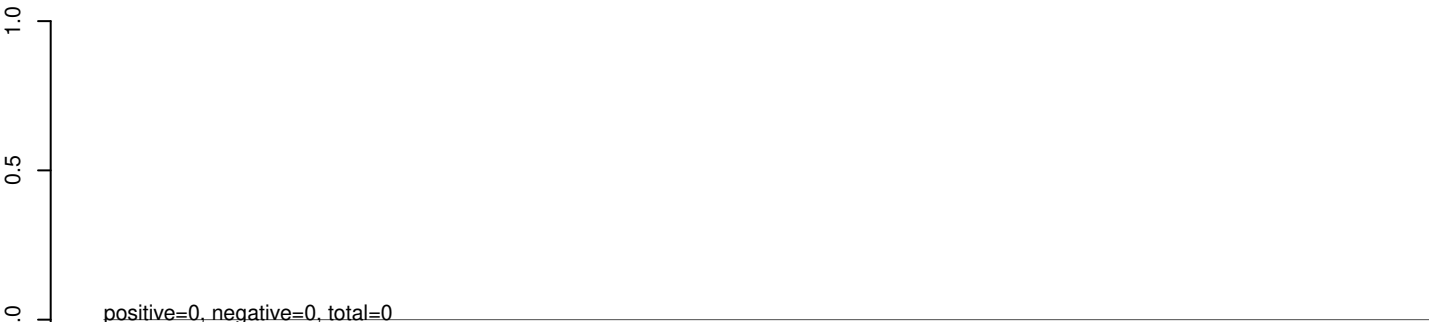


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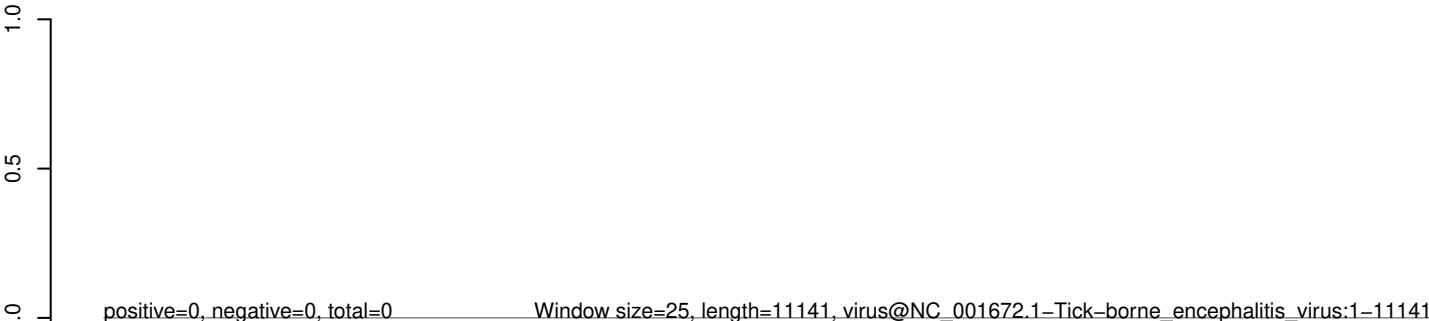
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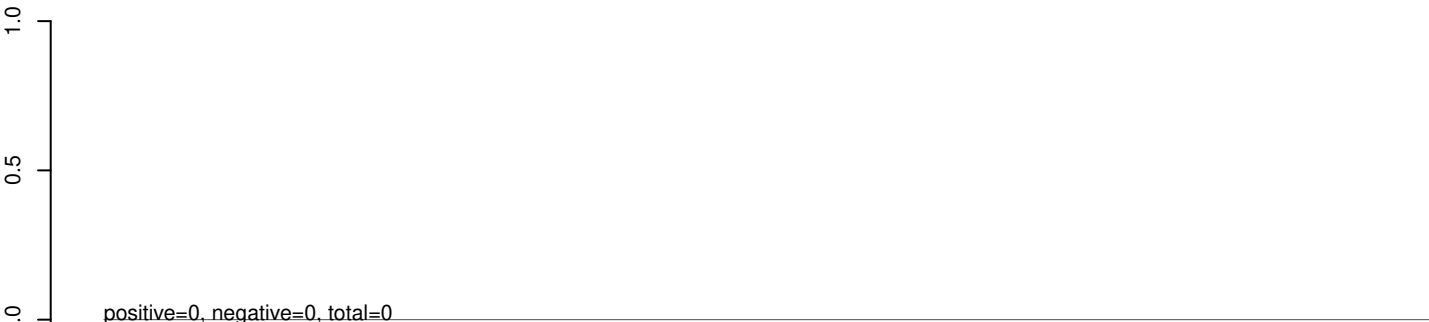


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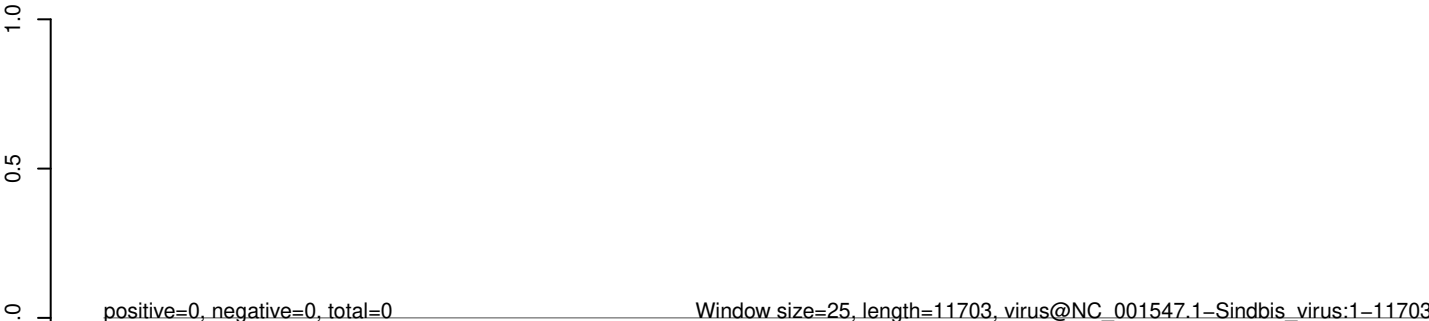
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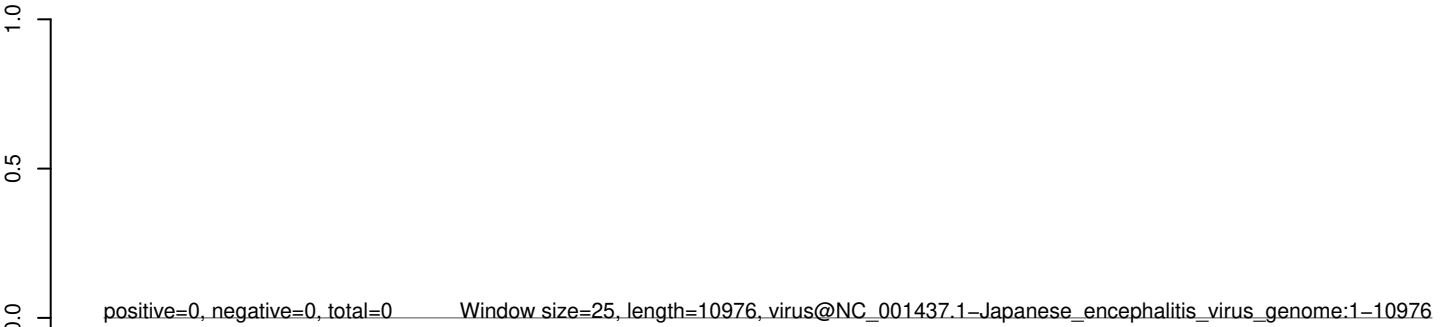
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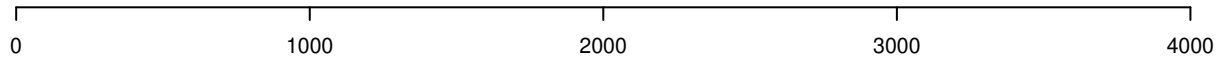
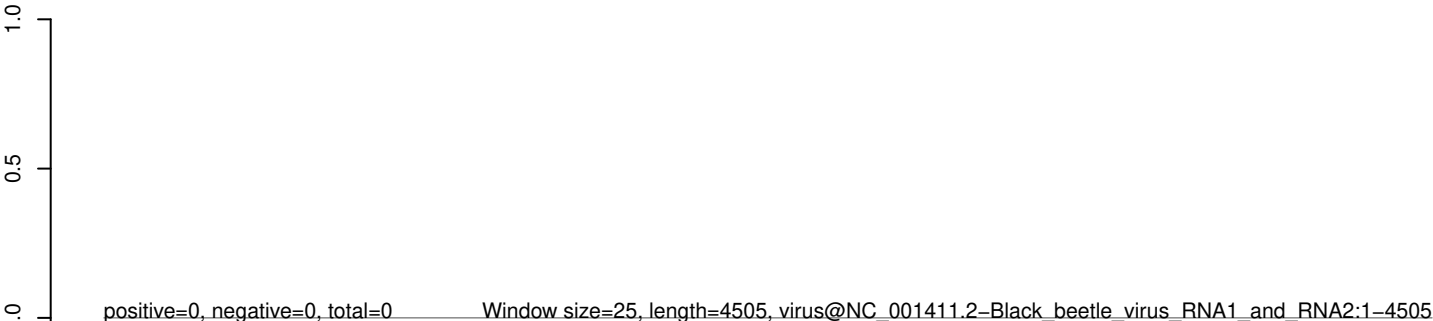
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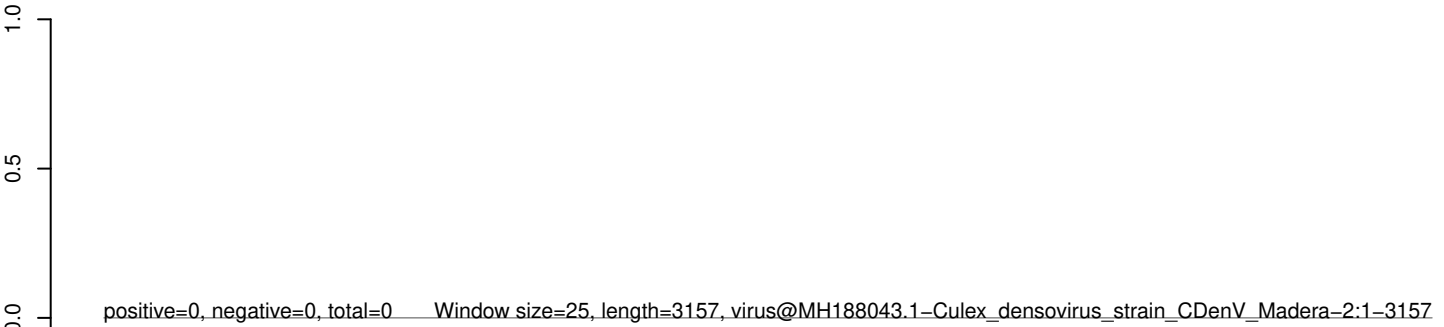
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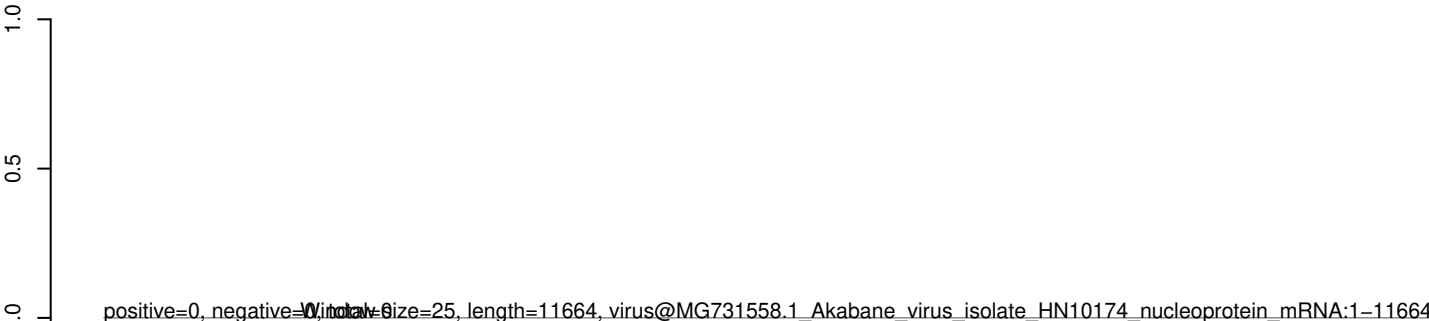
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AeAeg_Aag2_BTV2_AK.rep



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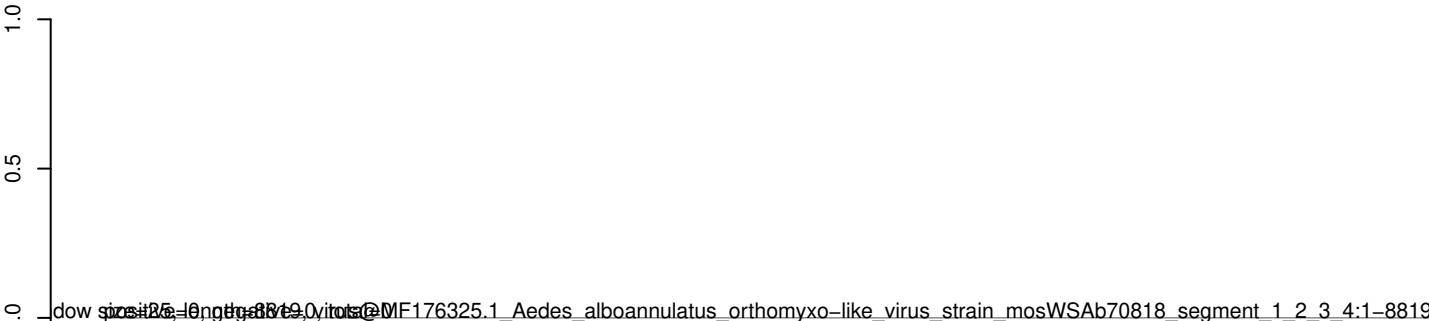
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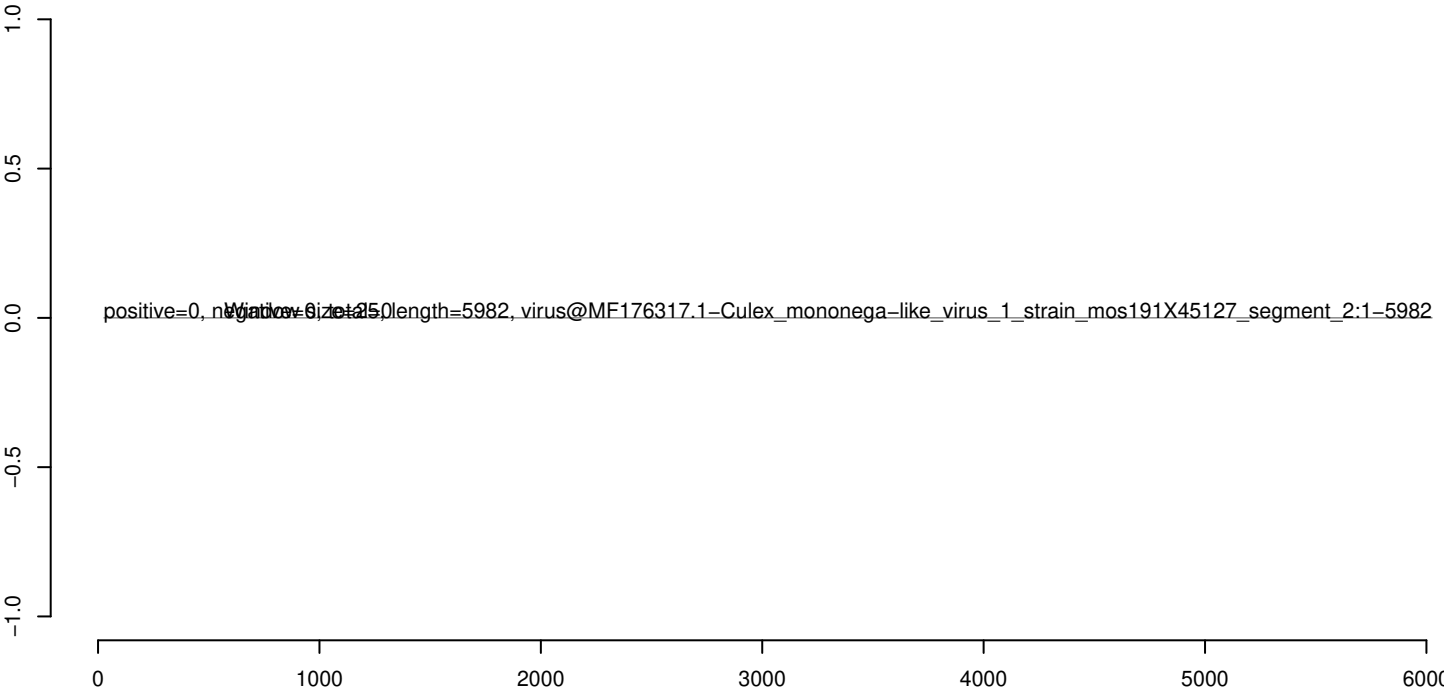
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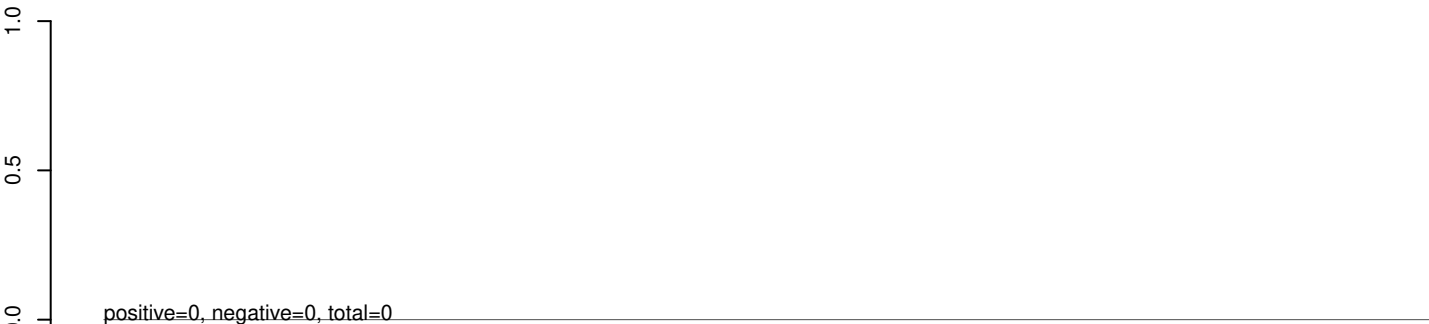
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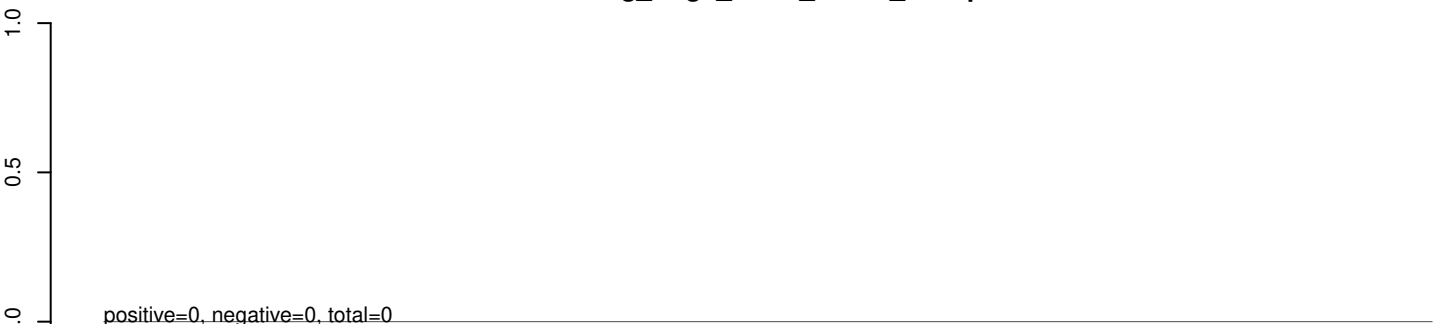
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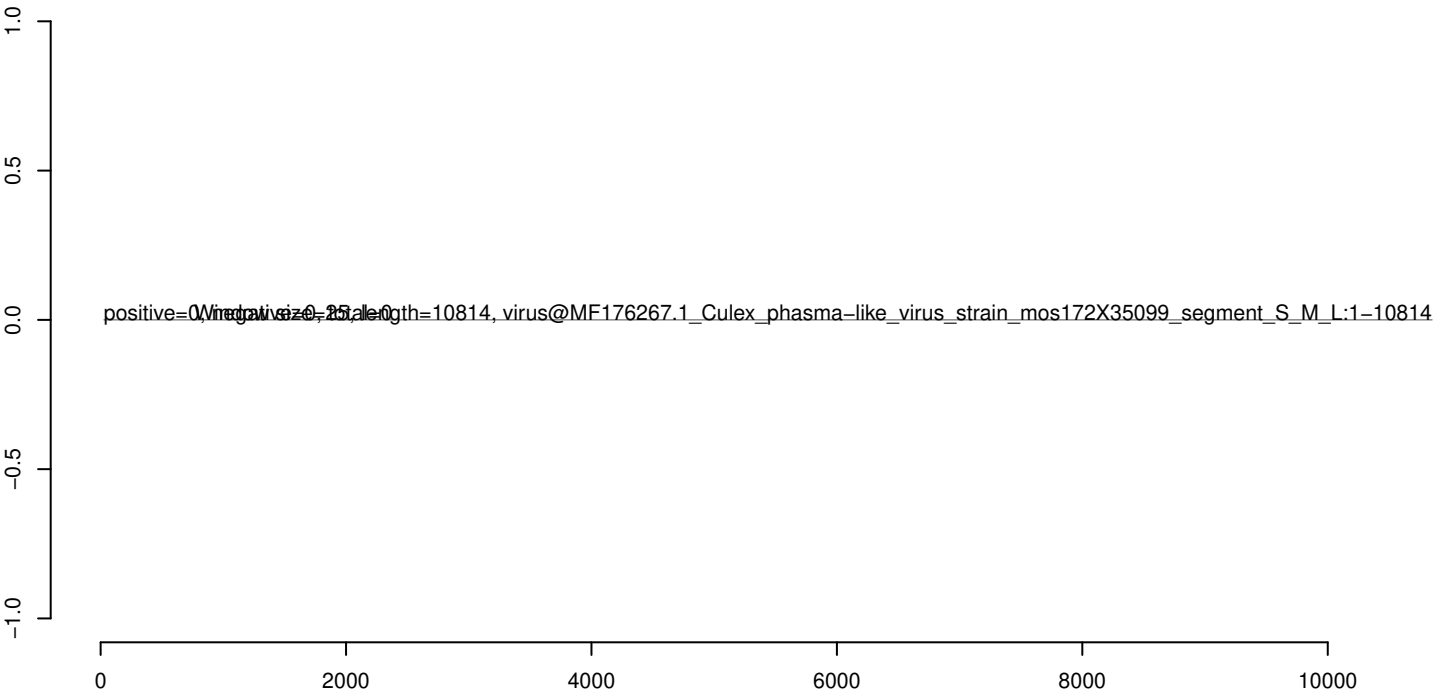
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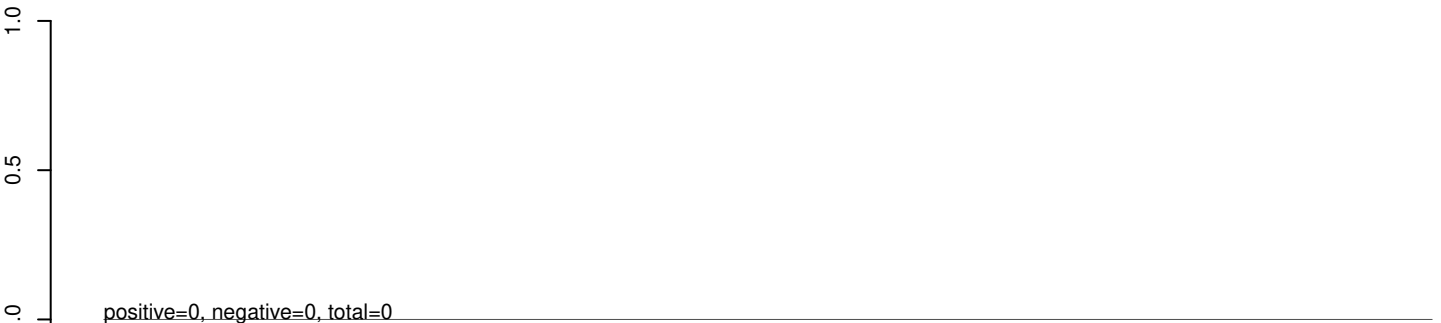
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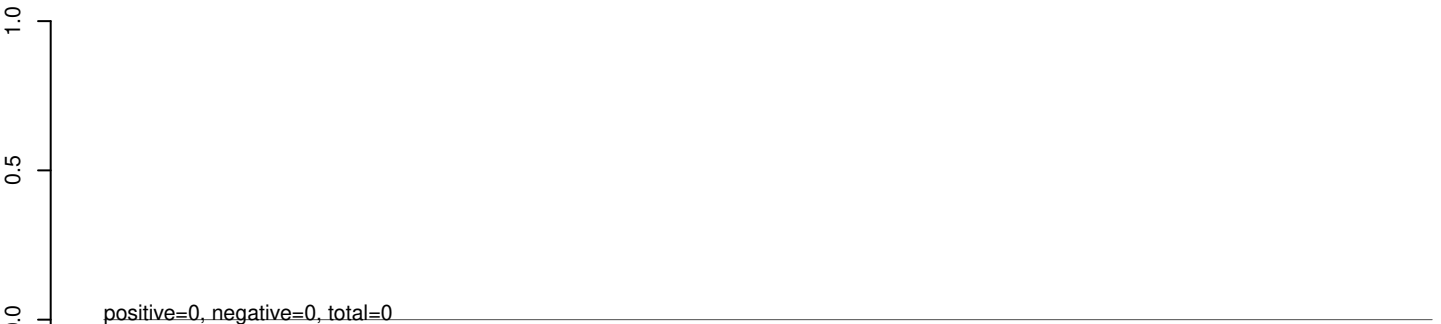
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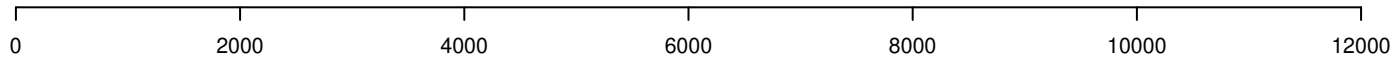
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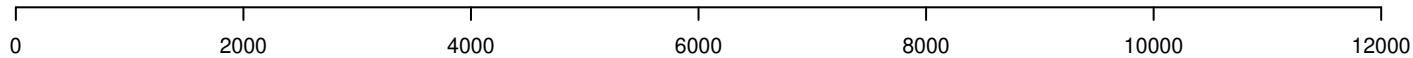
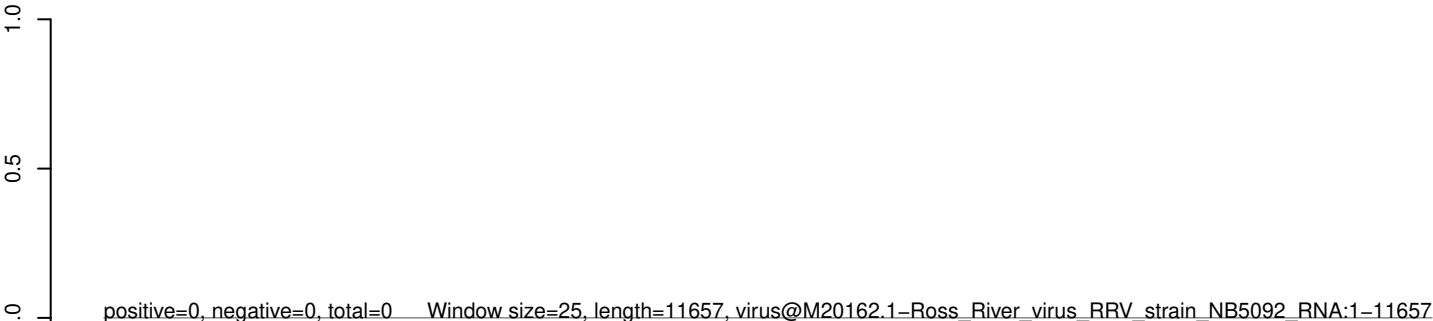
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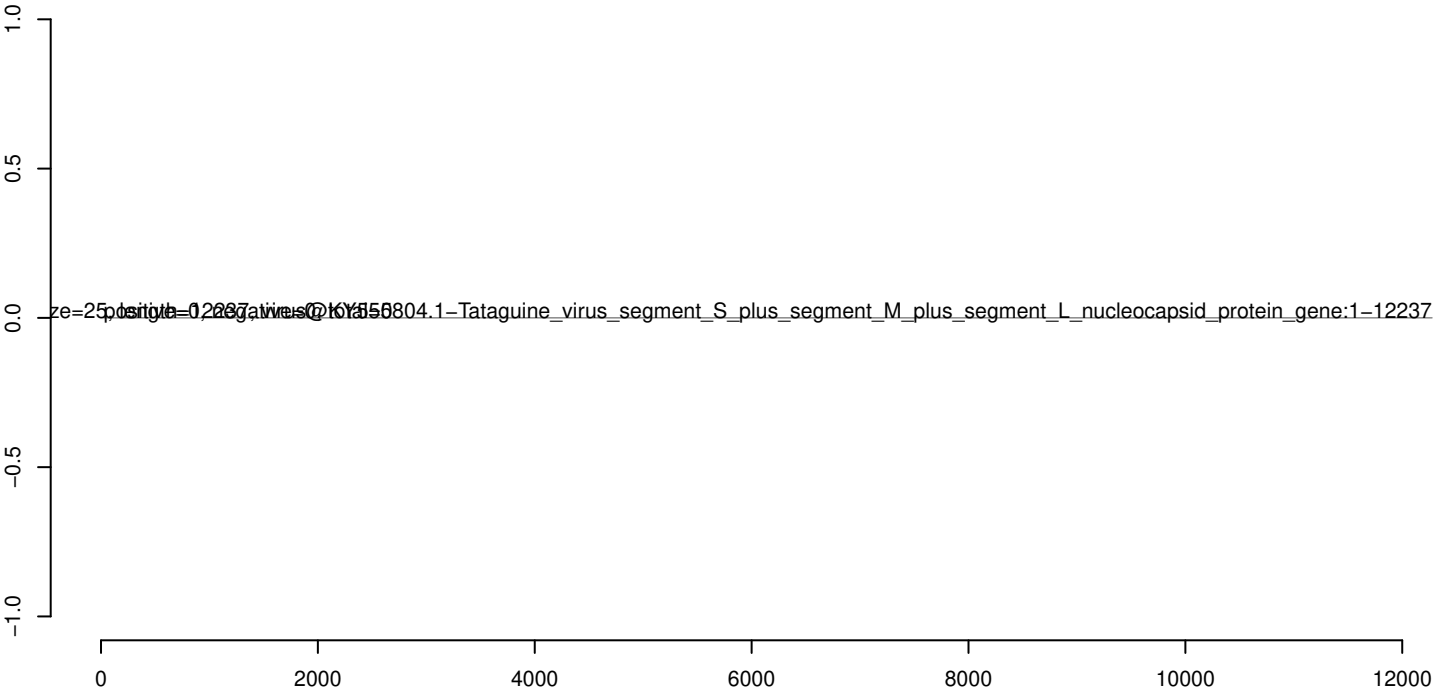
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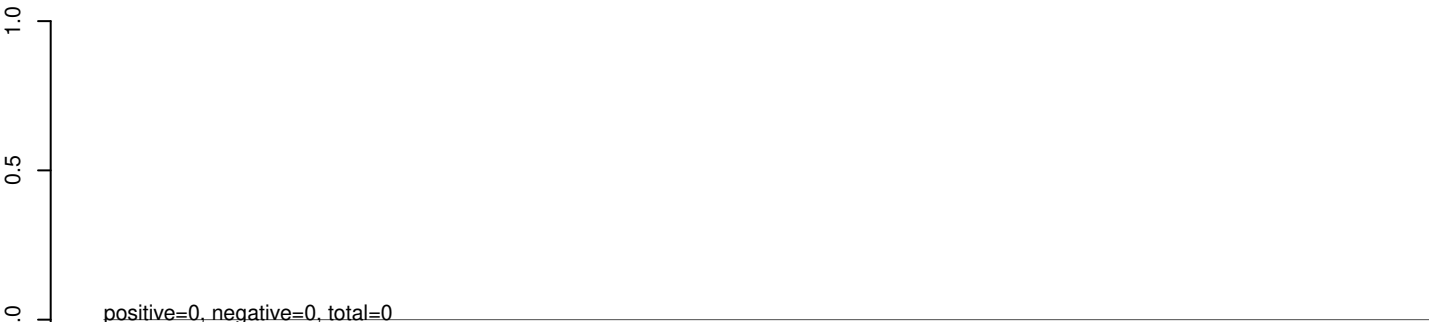
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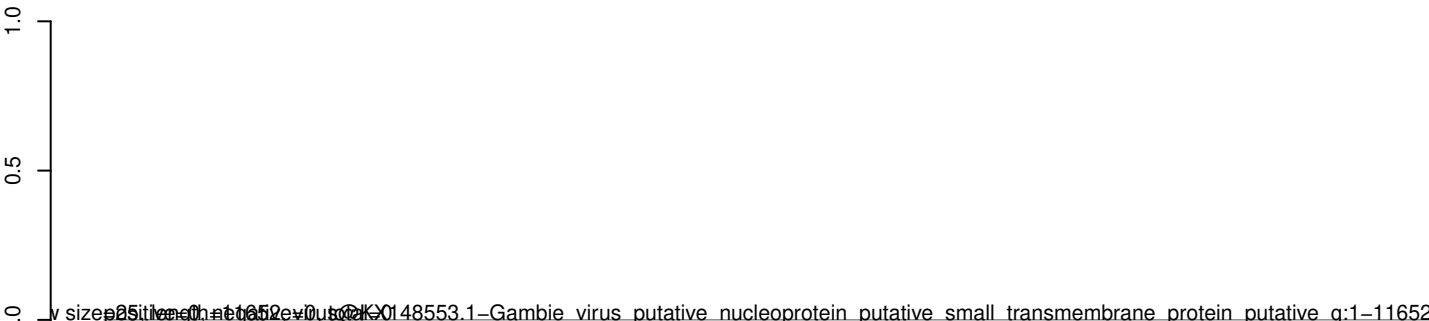
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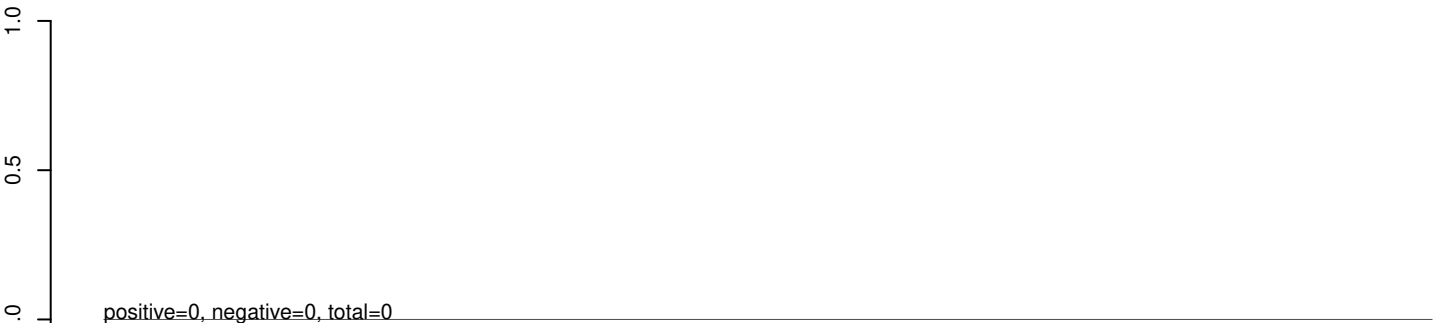


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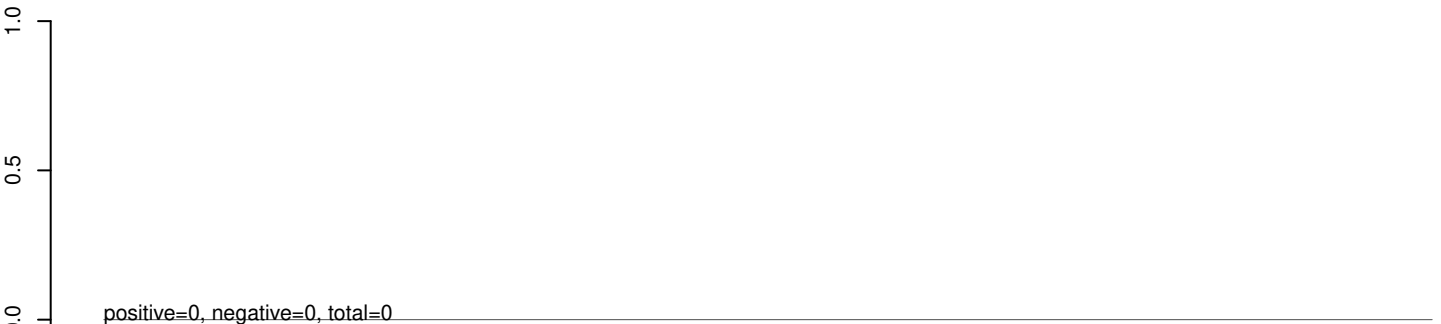


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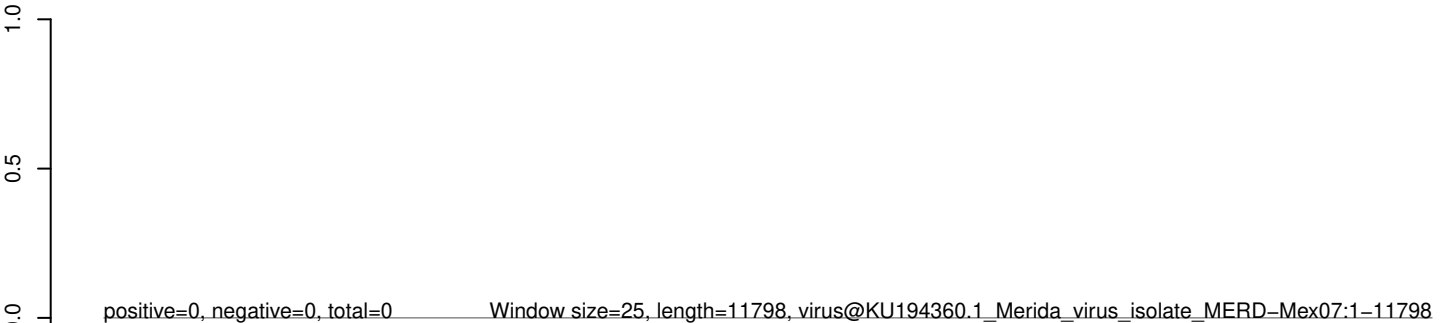
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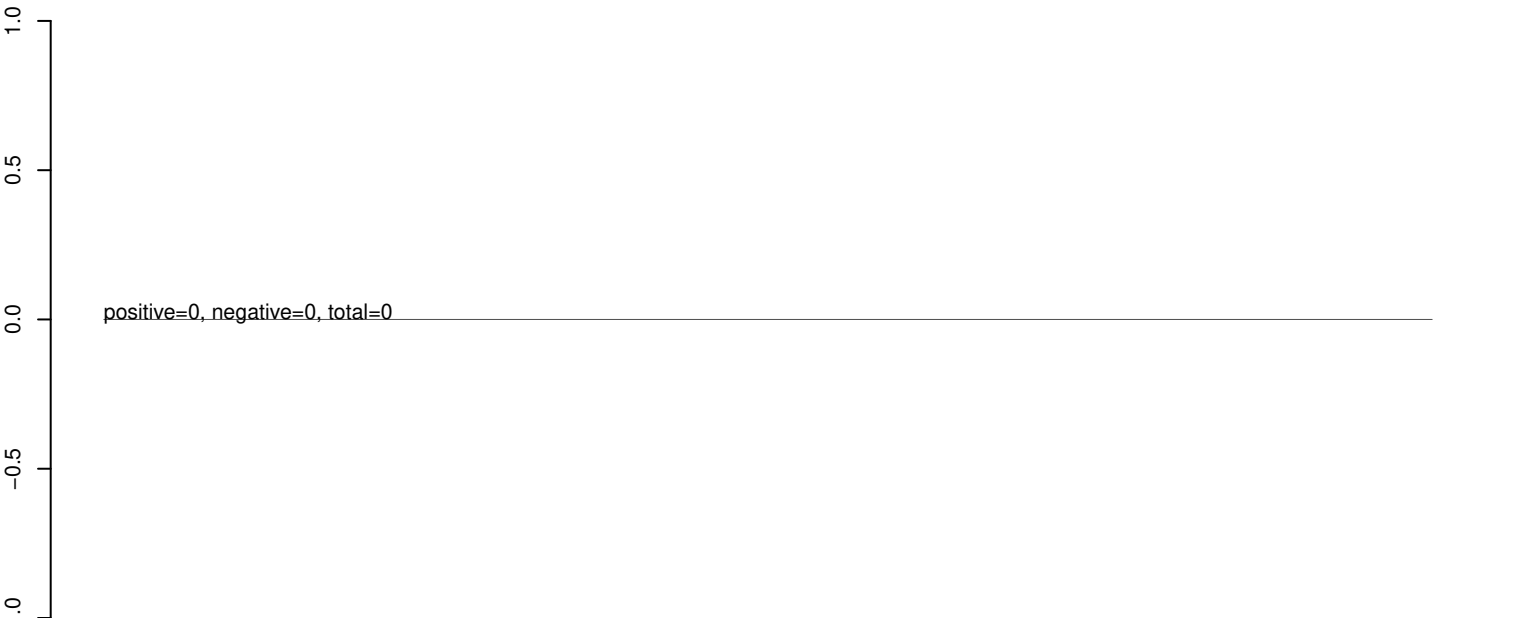


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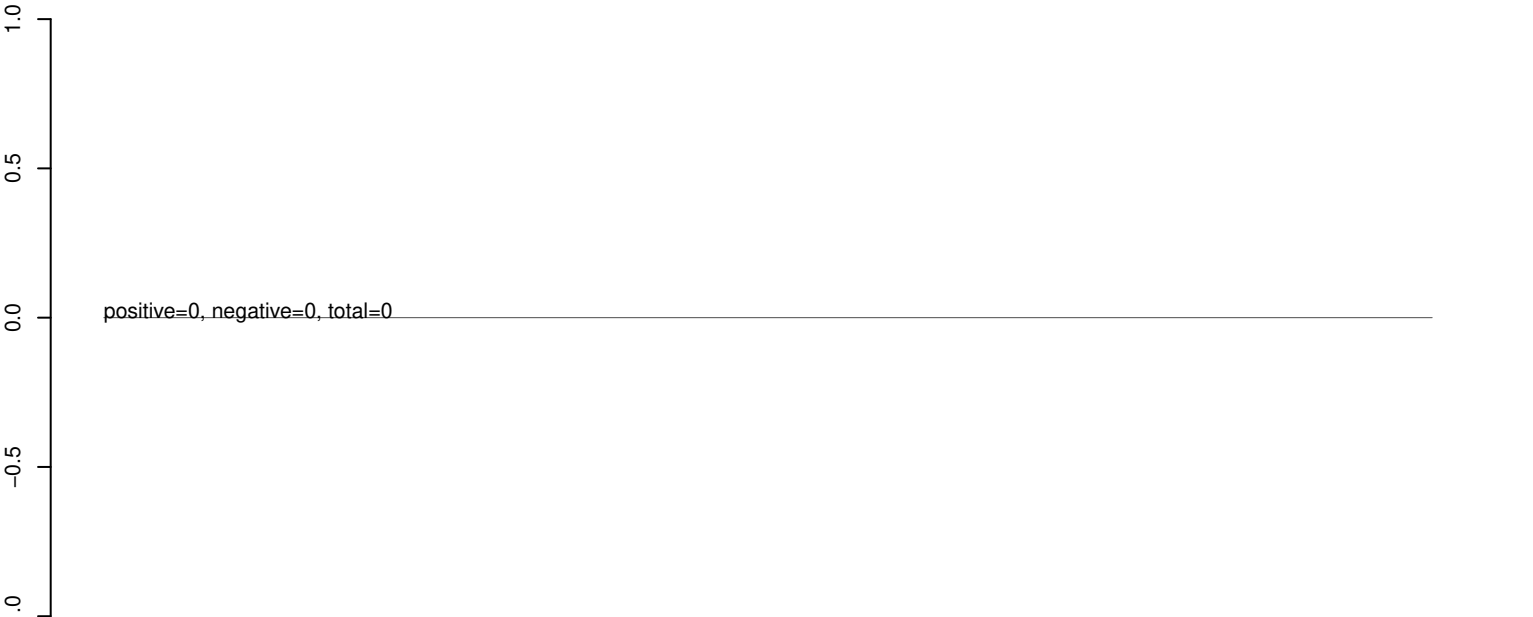


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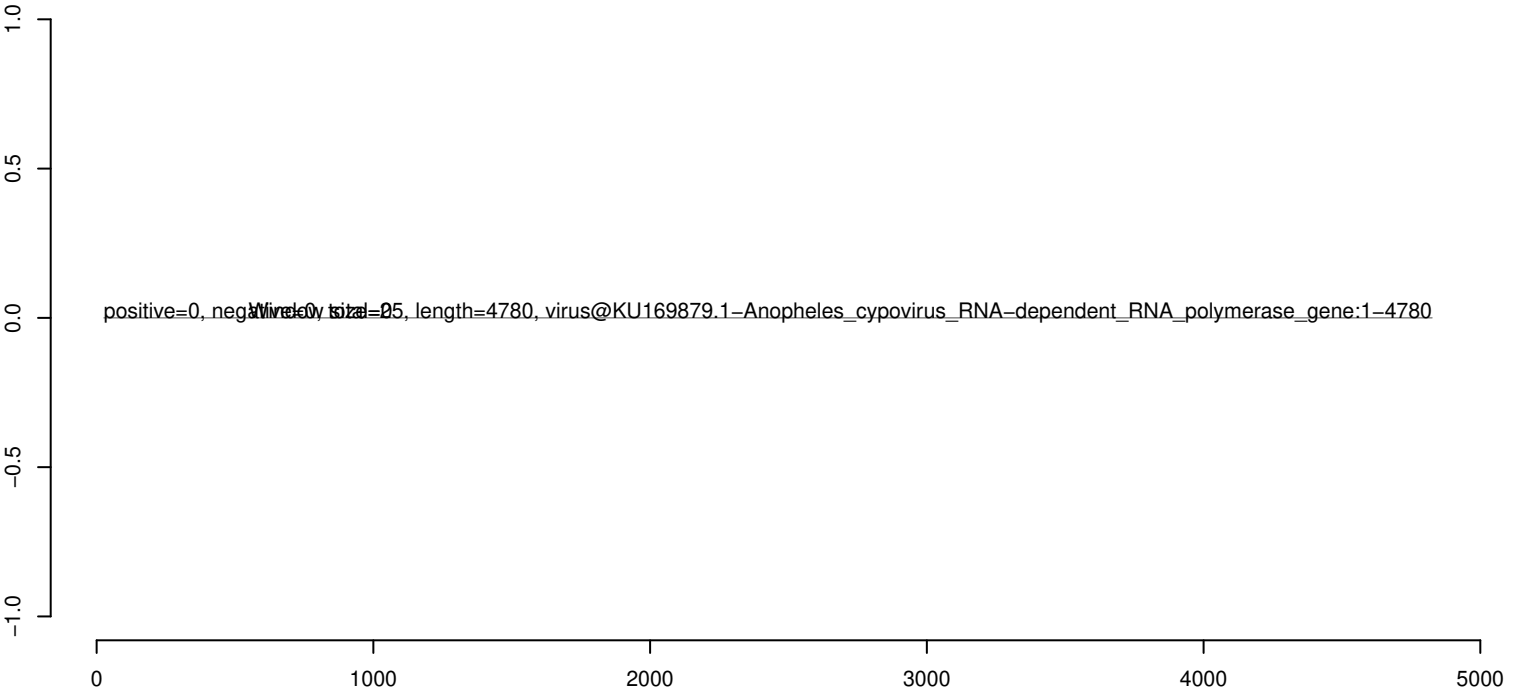
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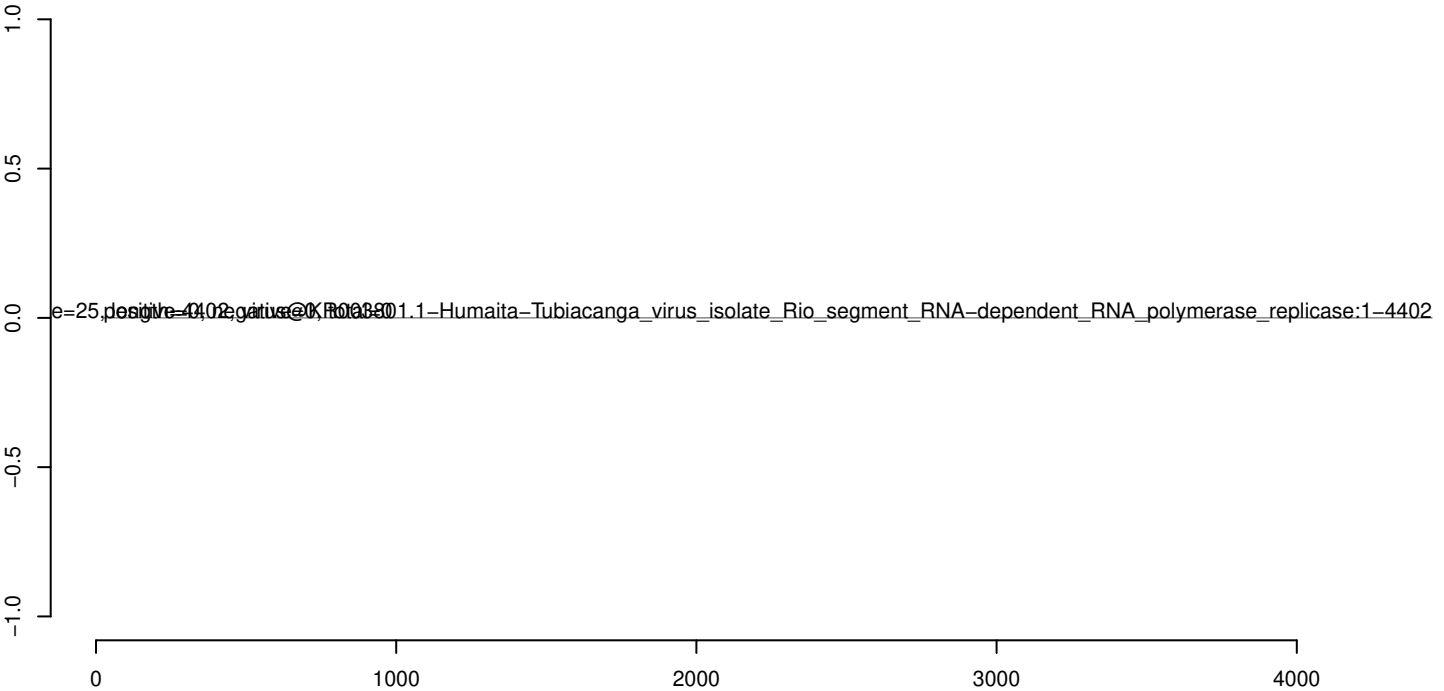
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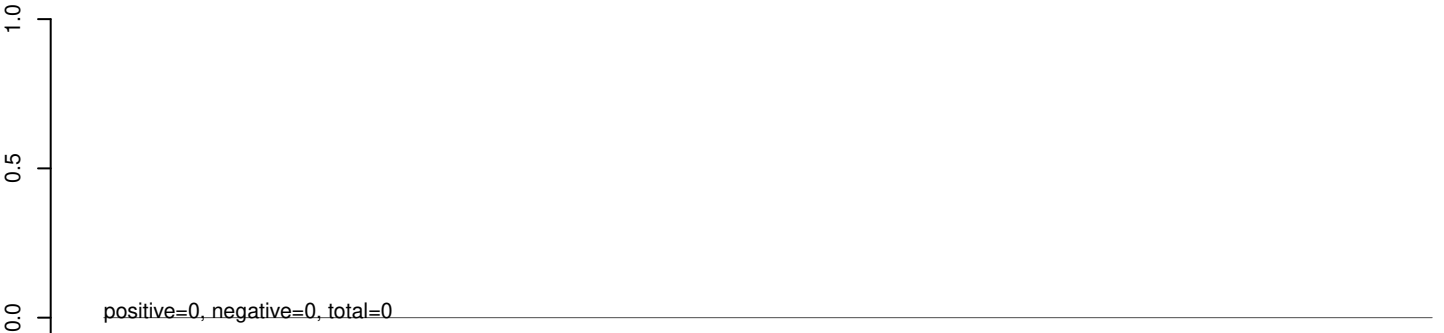
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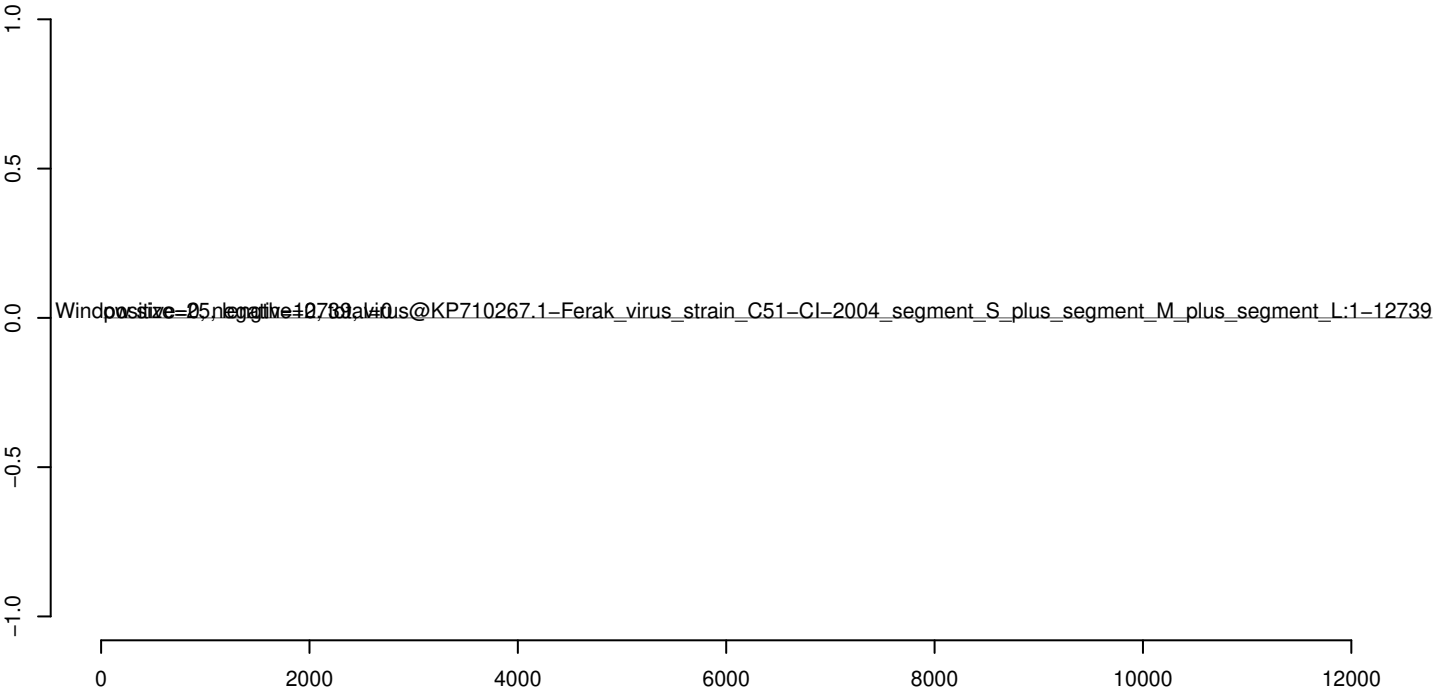
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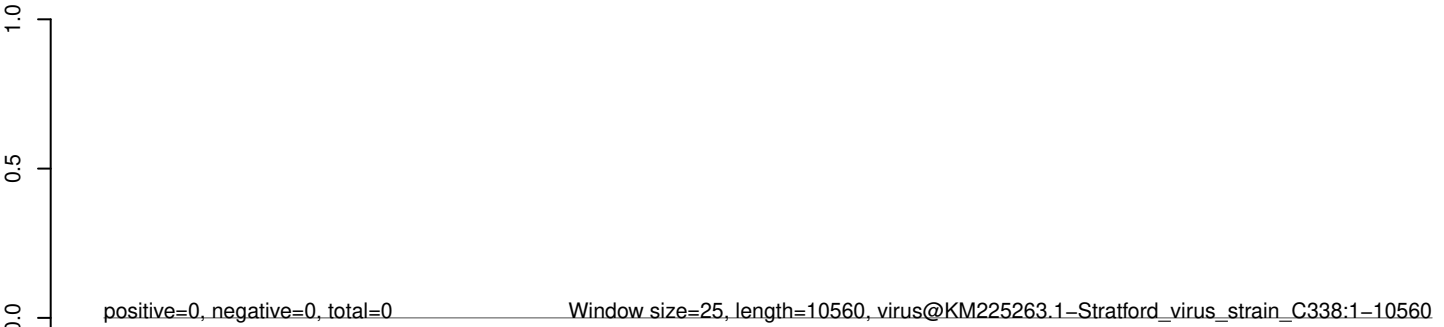
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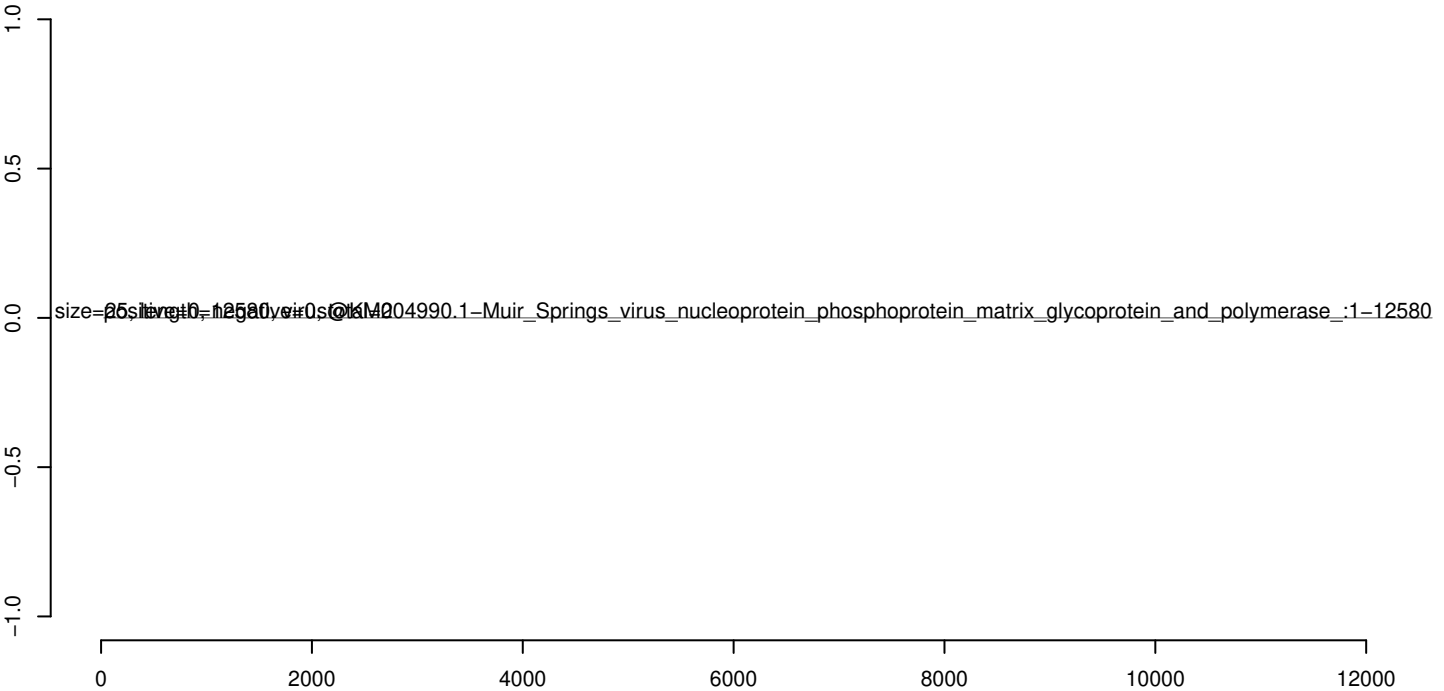
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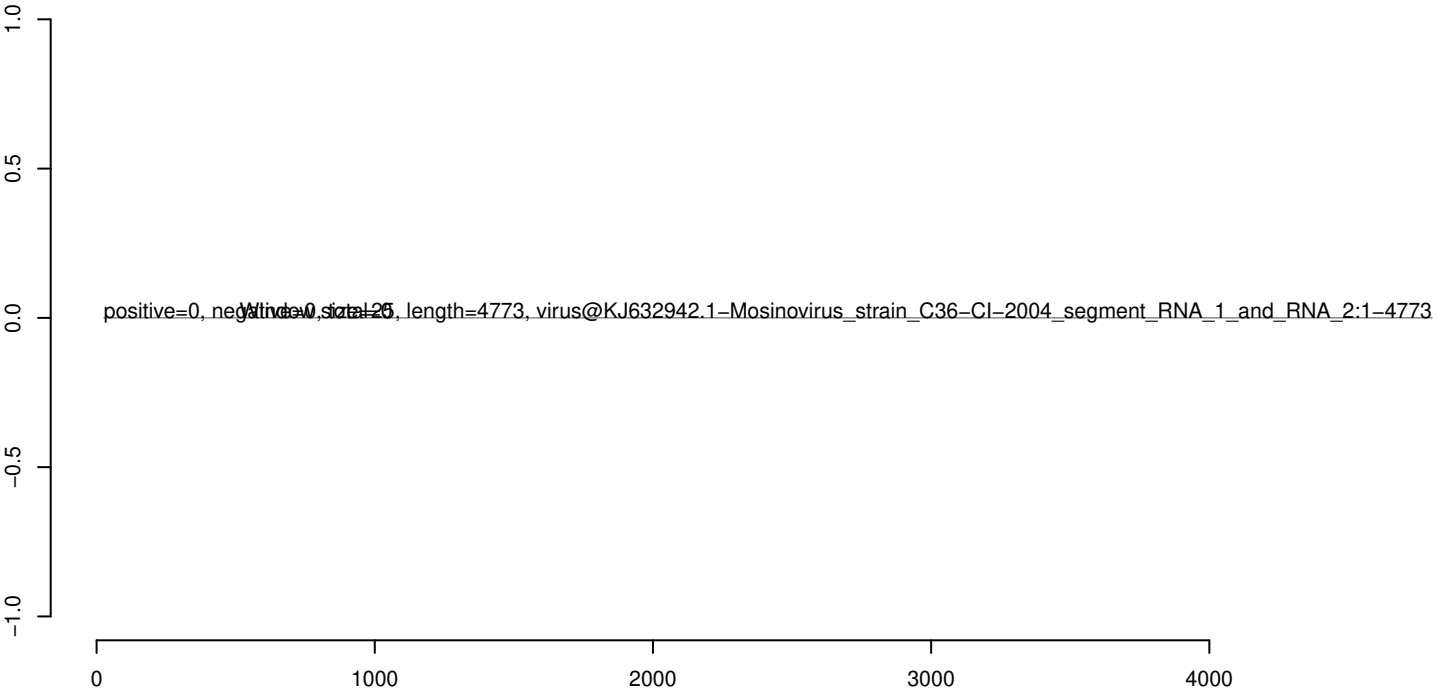
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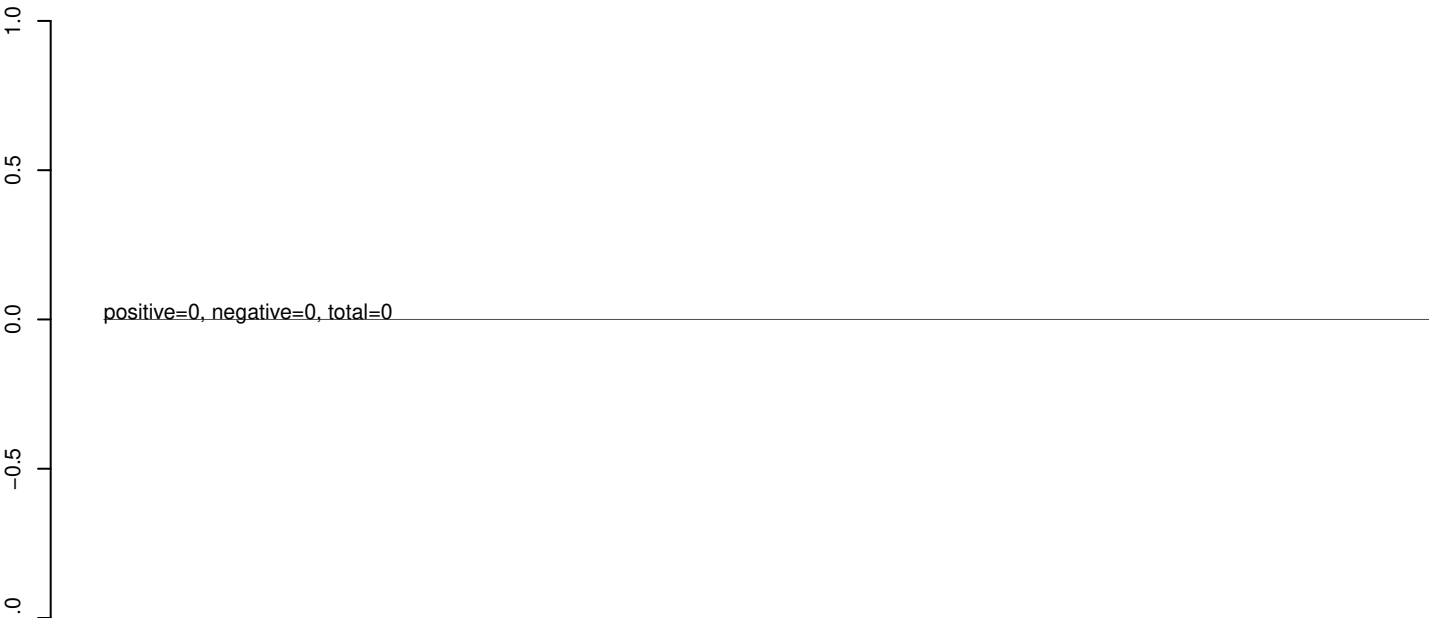
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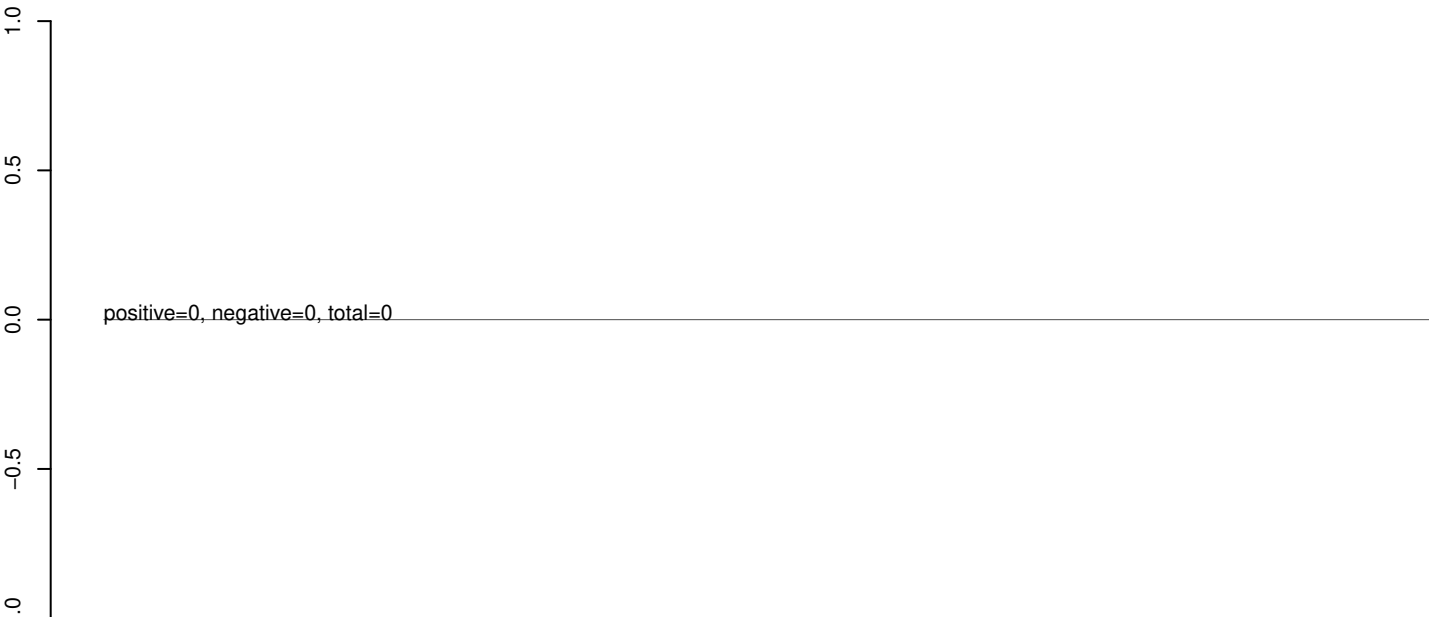
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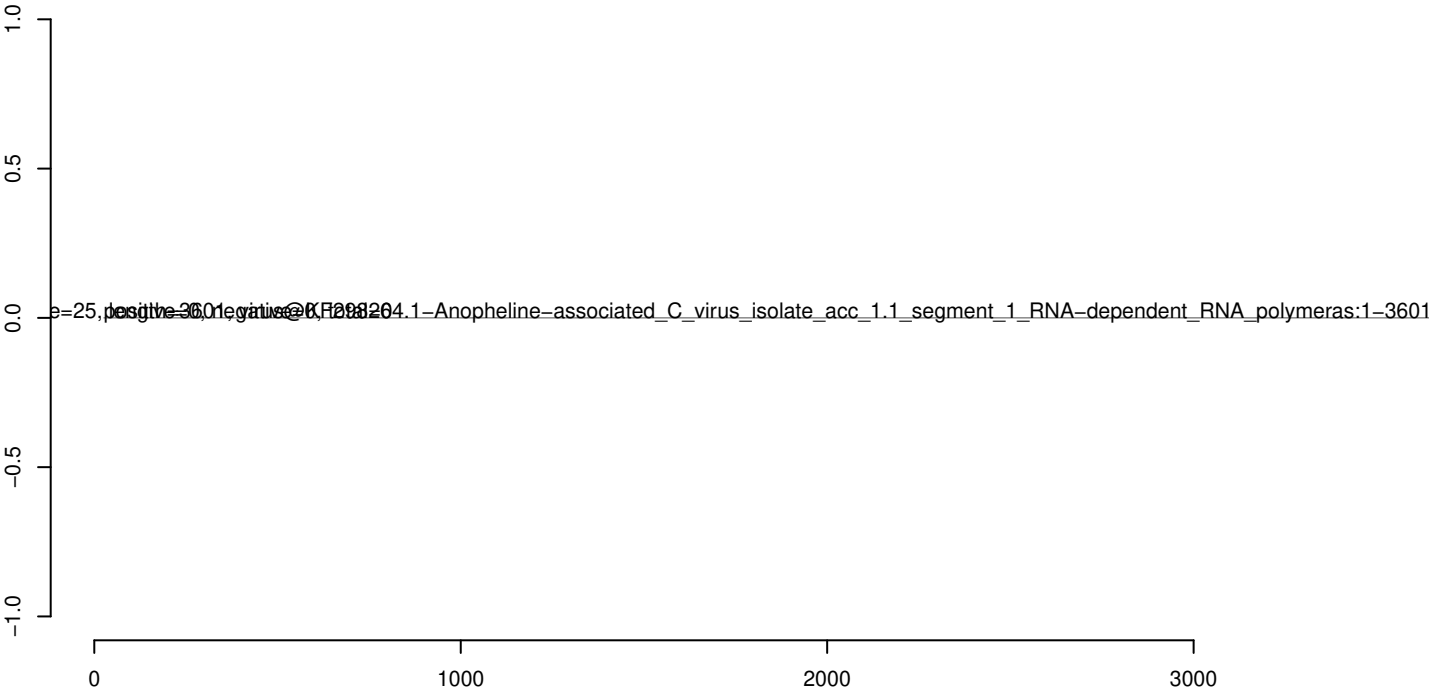
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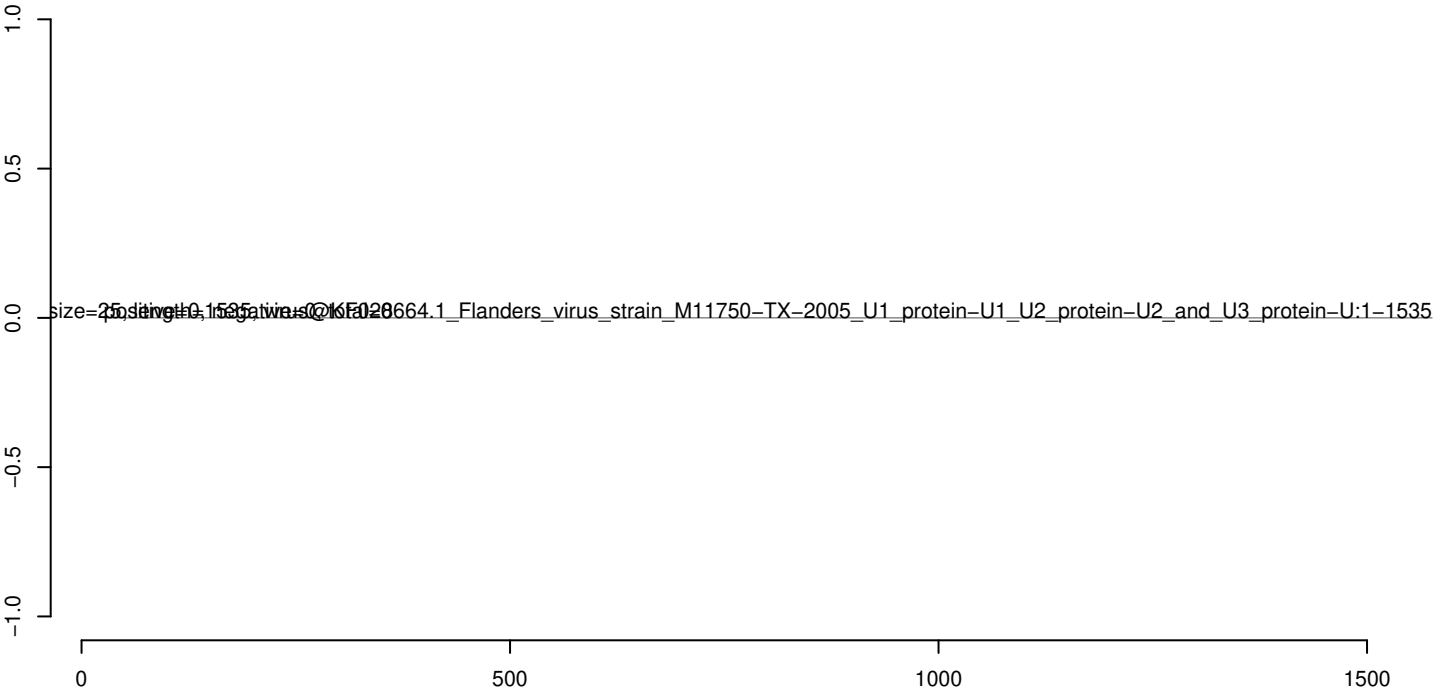
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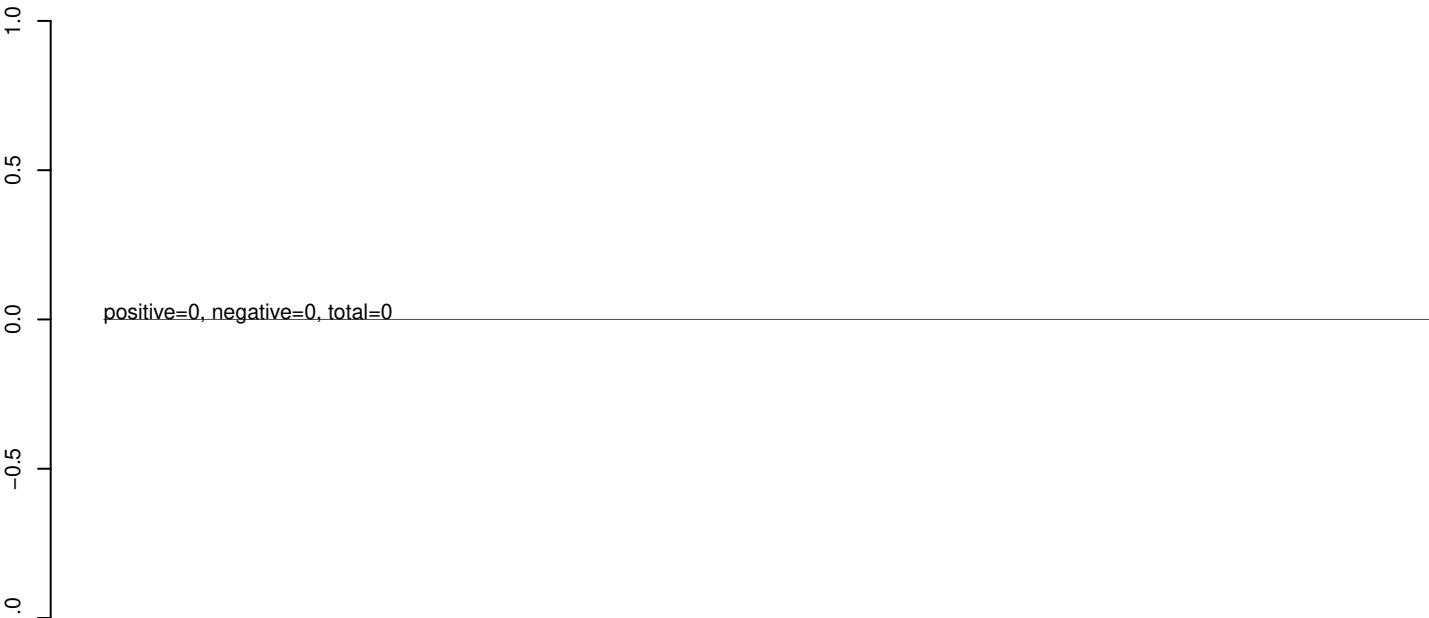
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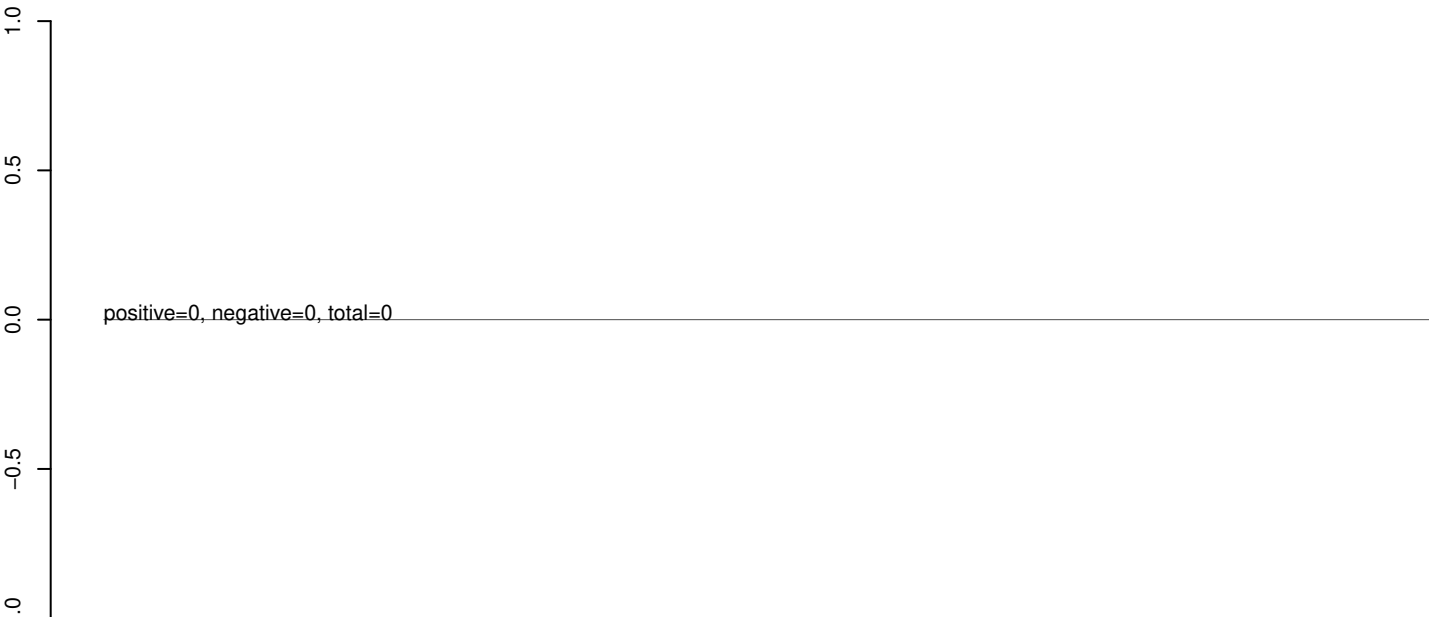
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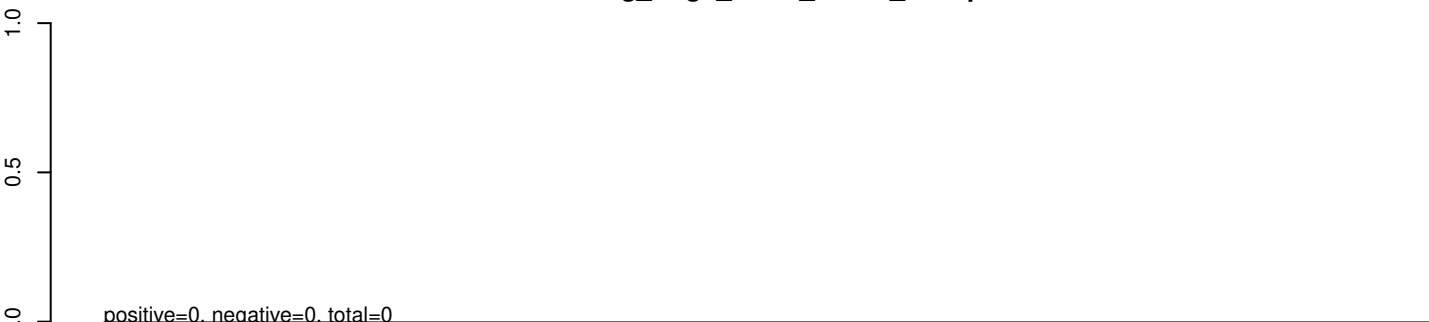
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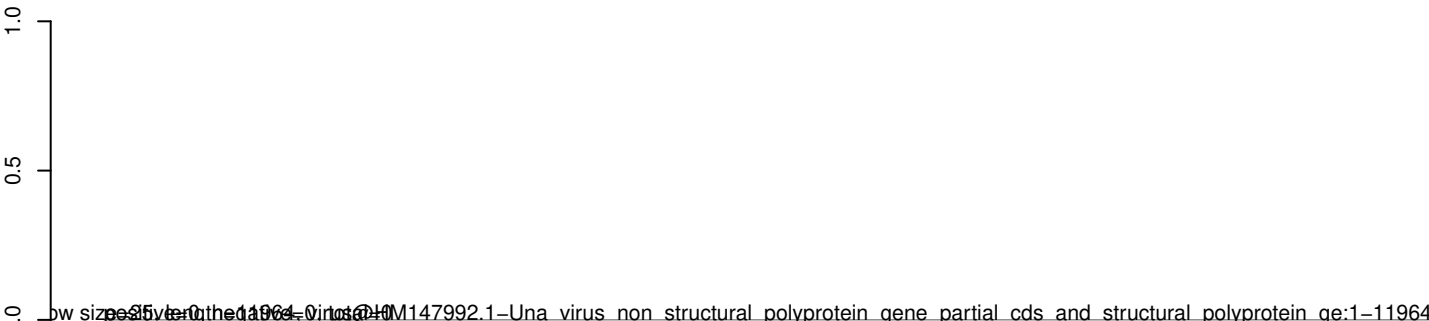
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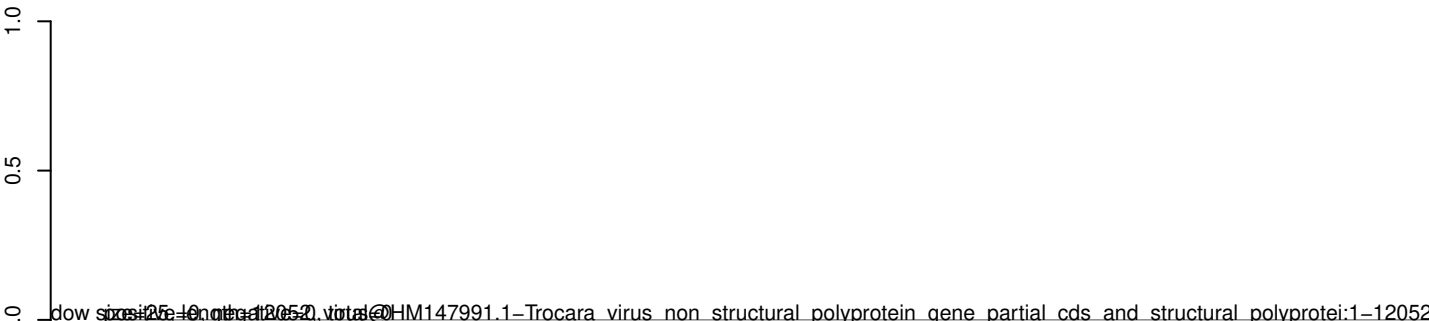
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0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



AeAeg_Aag2_BTV2_AK.18_23.rep



AeAeg_Aag2_BTV2_AK.24_35.rep



AeAeg_Aag2_BTV2_AK.rep



165.1-Drosophila_melanogaster_American_nodavirus_ANV_SW-2009a_segment_RNA1_and_RNA2:1-4523

AeAeg_Aag2_BTV2_AK.18_23.rep



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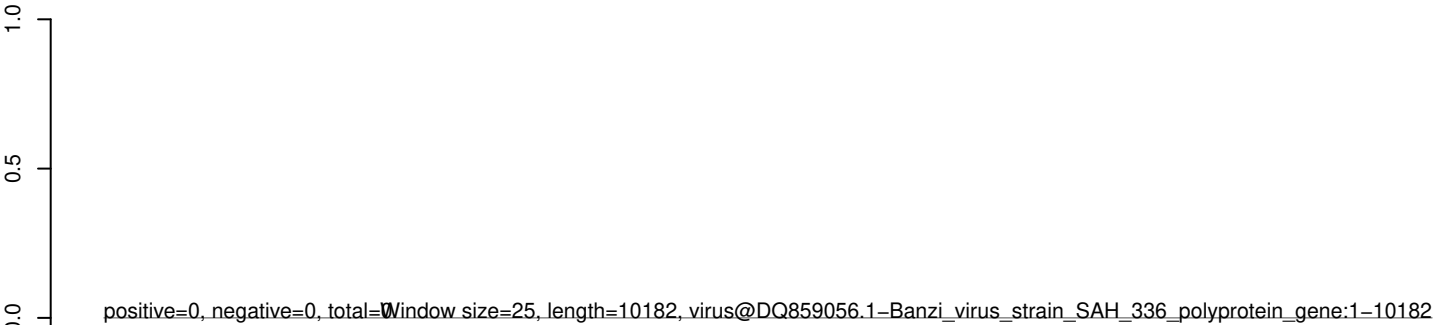
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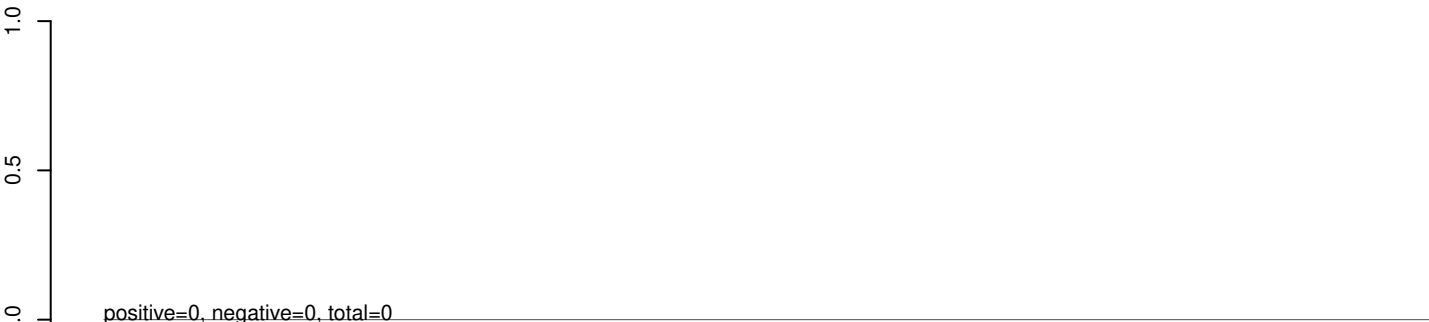
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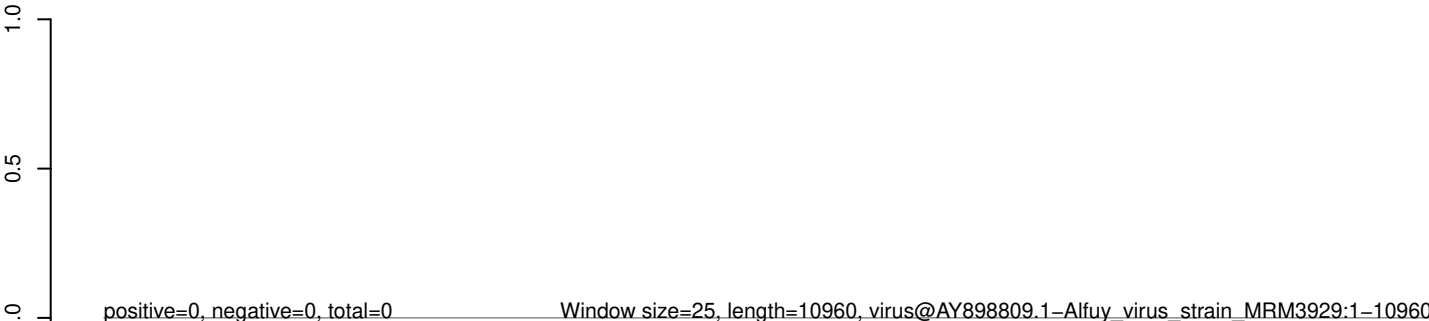
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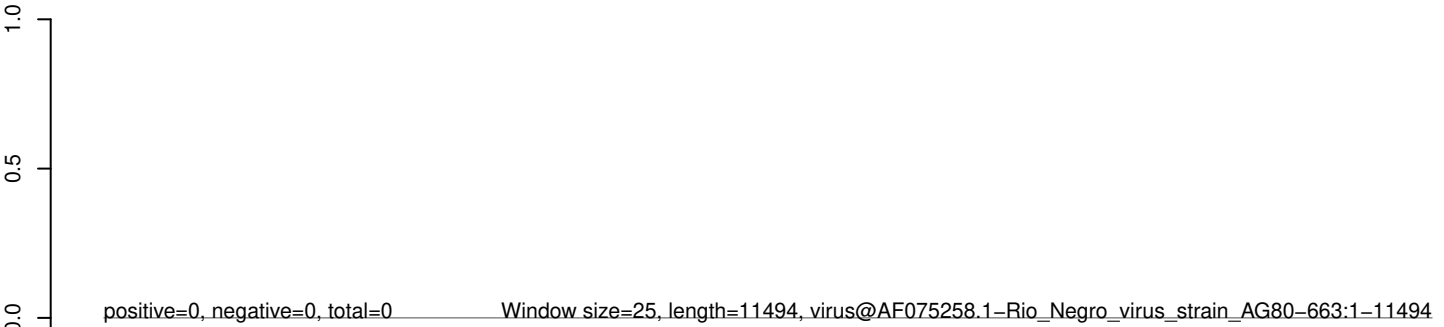
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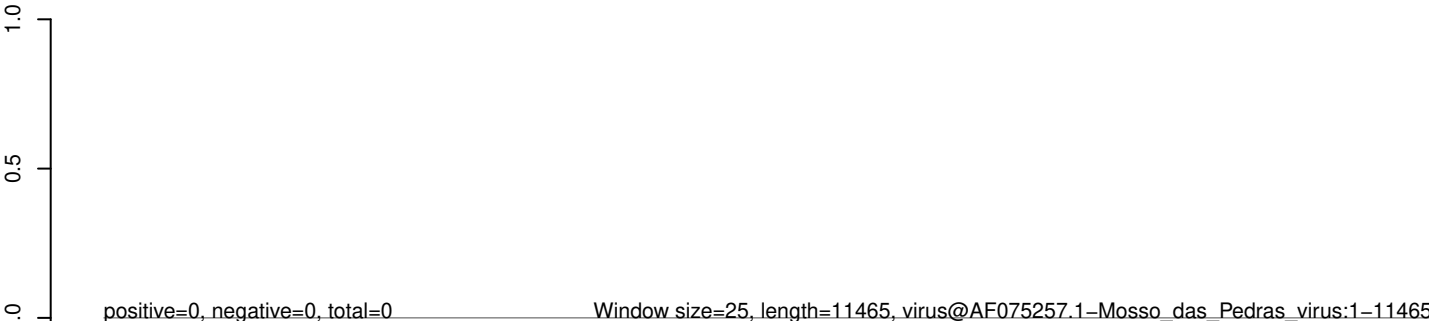
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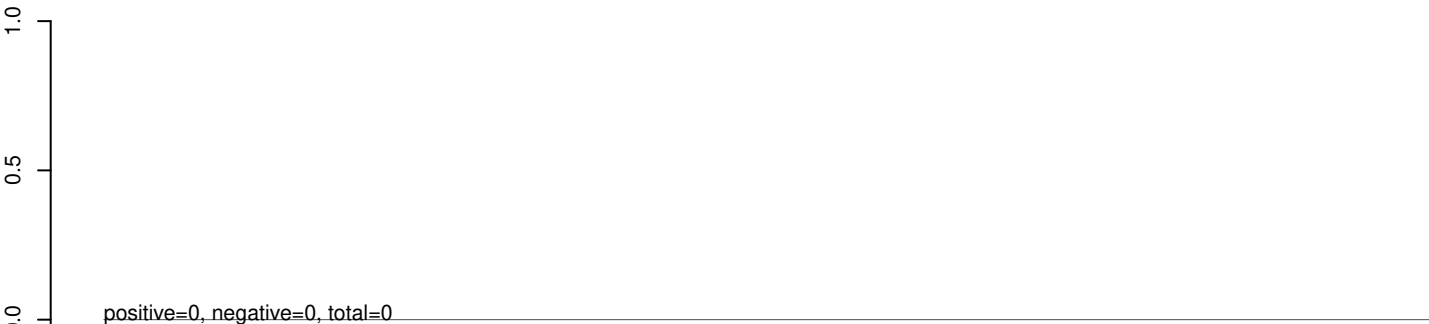


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