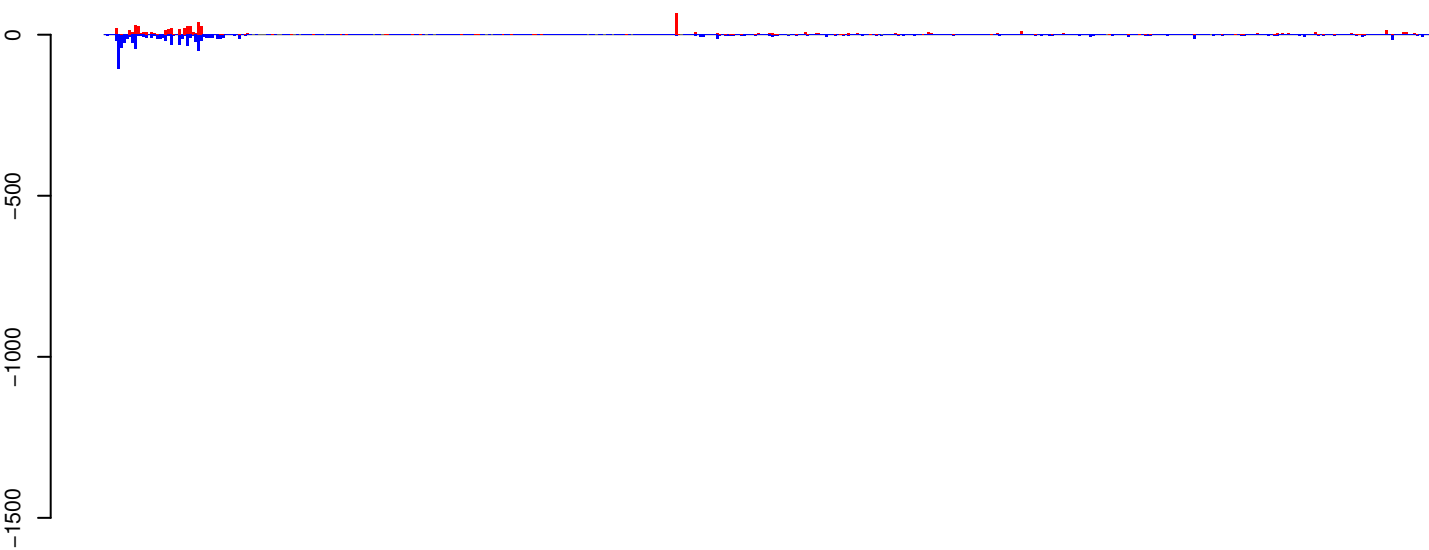
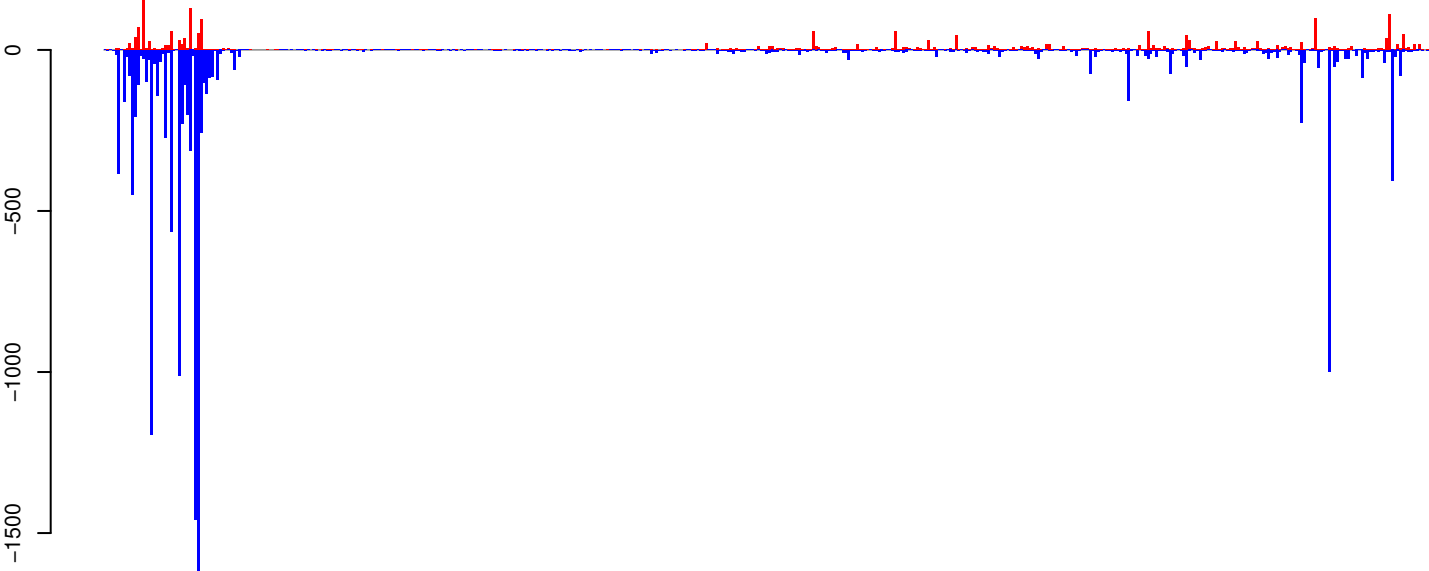


AeAeg_Aag2SINV_dsLuc.18_23.rep



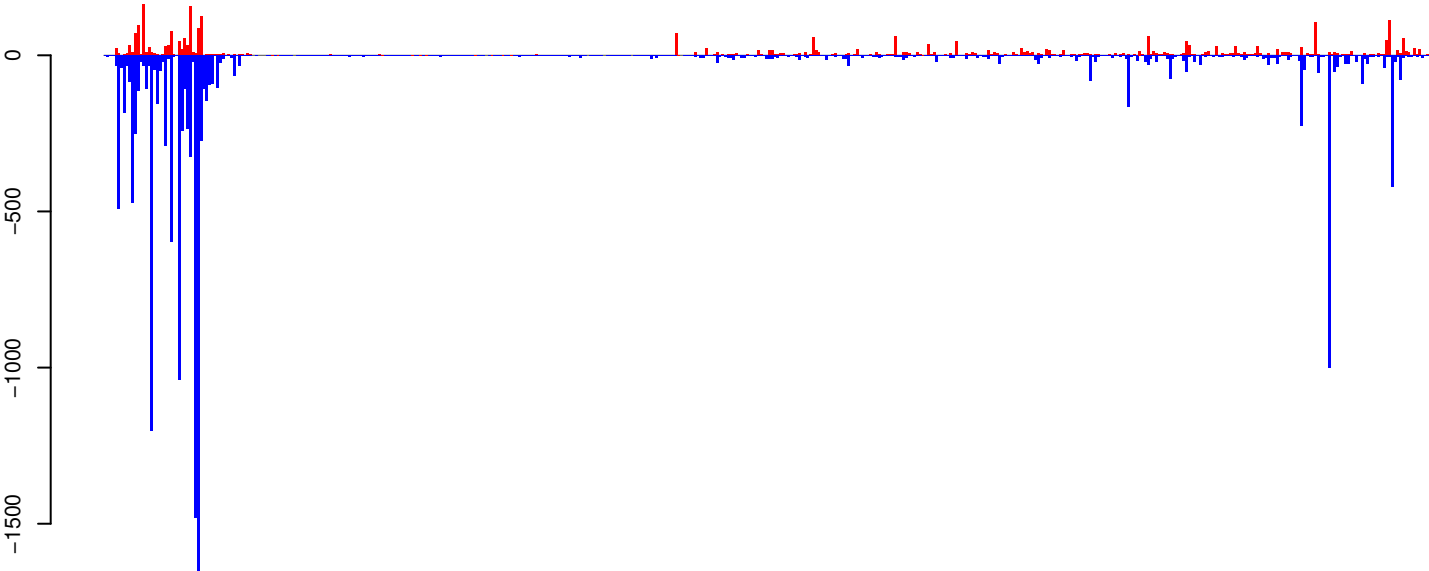
positive=836, negative=1092, total=1928

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=2417, negative=13293, total=15710

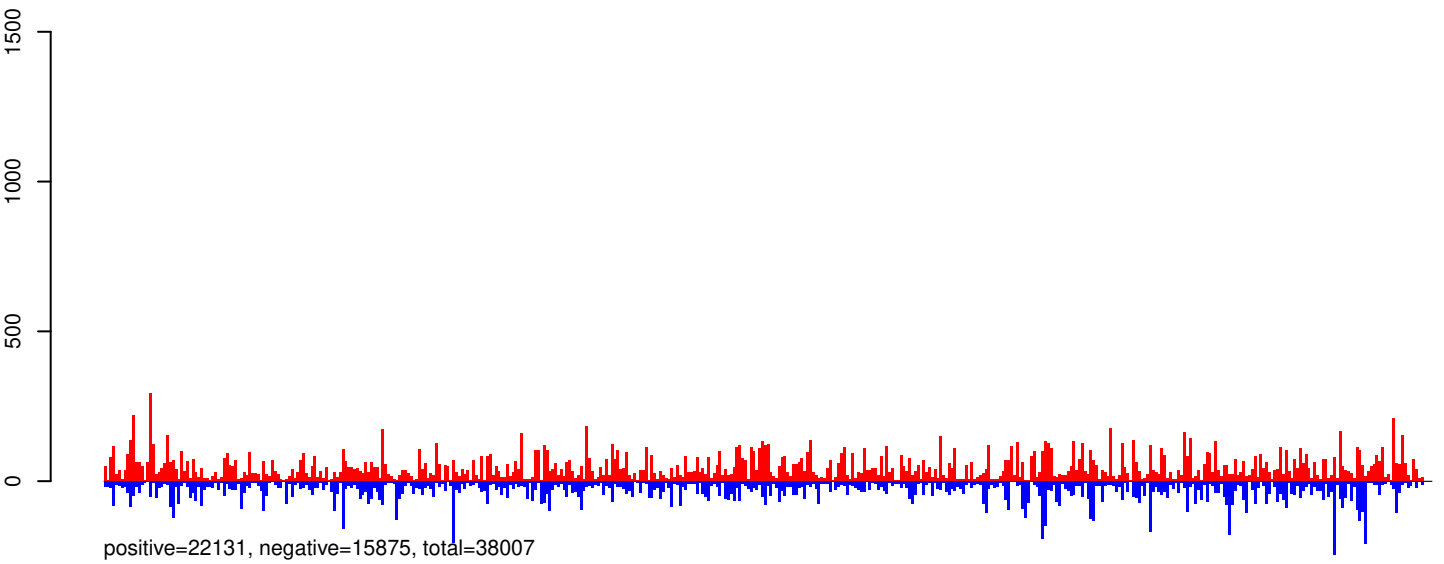
AeAeg_Aag2SINV_dsLuc.rep



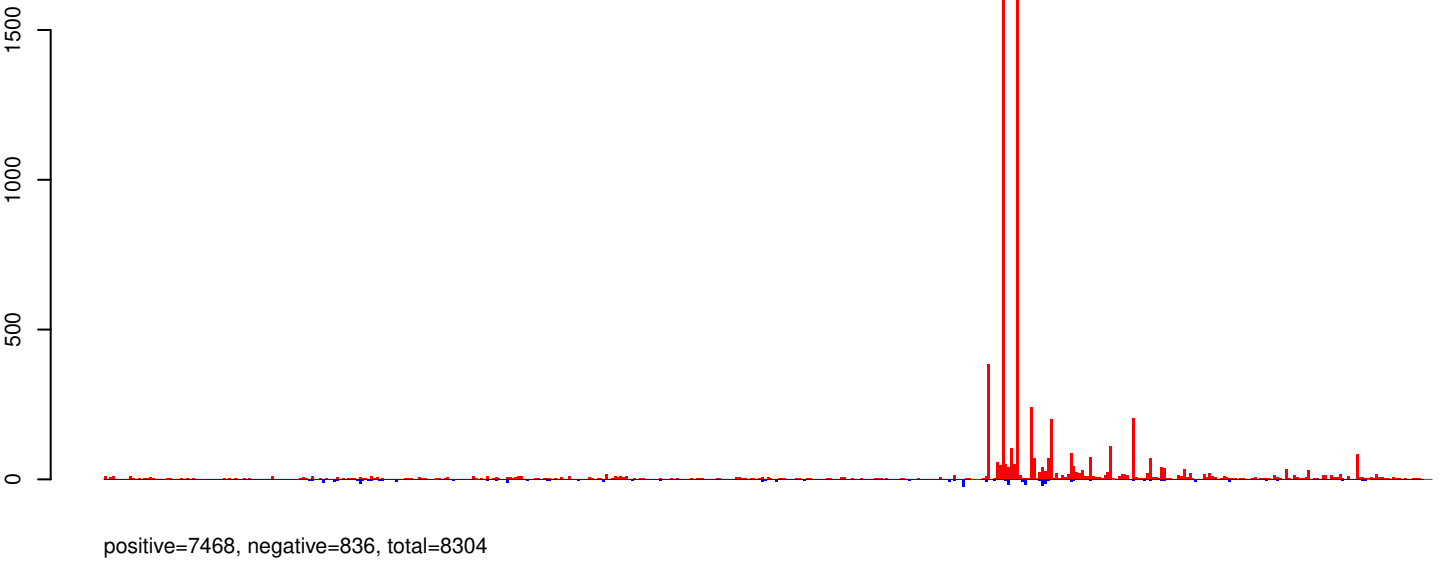
-25, positive=42893, negative=103825, total=146718 Charoen-like_virus_isolate_Rio_segment_S_plus_segment_M_plus_segment_L_Nucle:1-12099

0 2000 4000 6000 8000 10000 12000

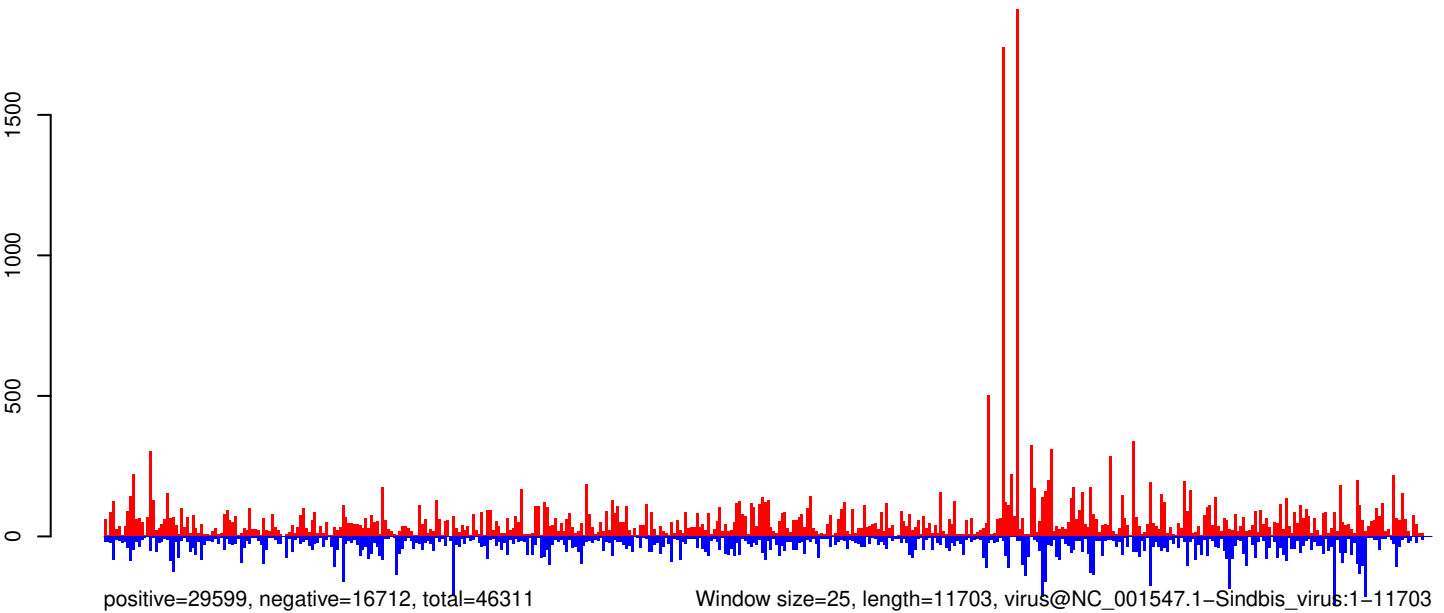
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

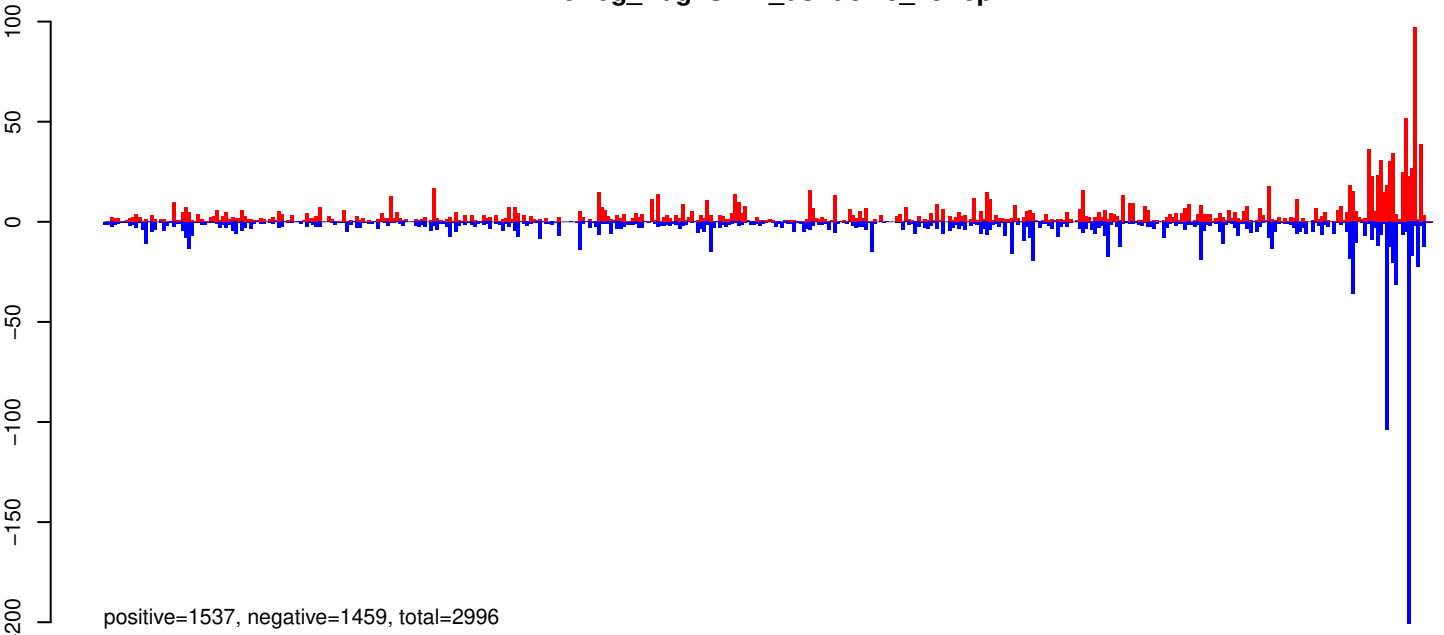


AeAeg_Aag2SINV_dsLuc.rep

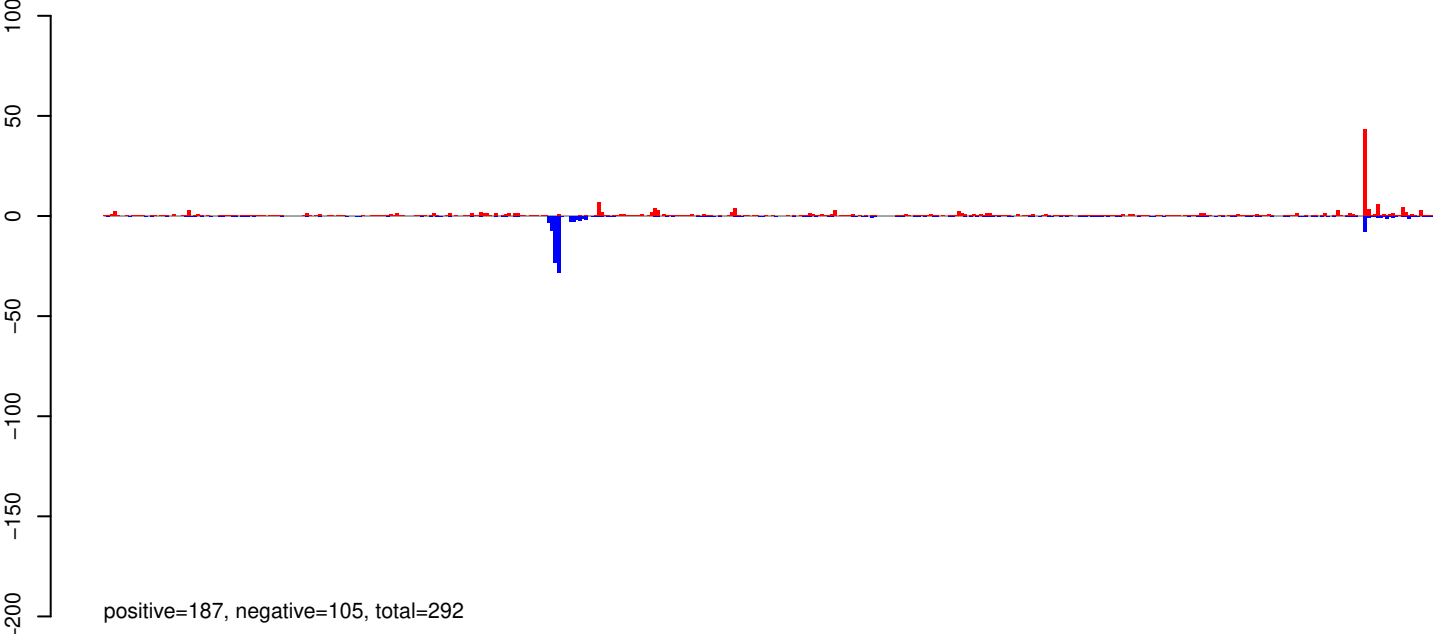


0 2000 4000 6000 8000 10000 12000

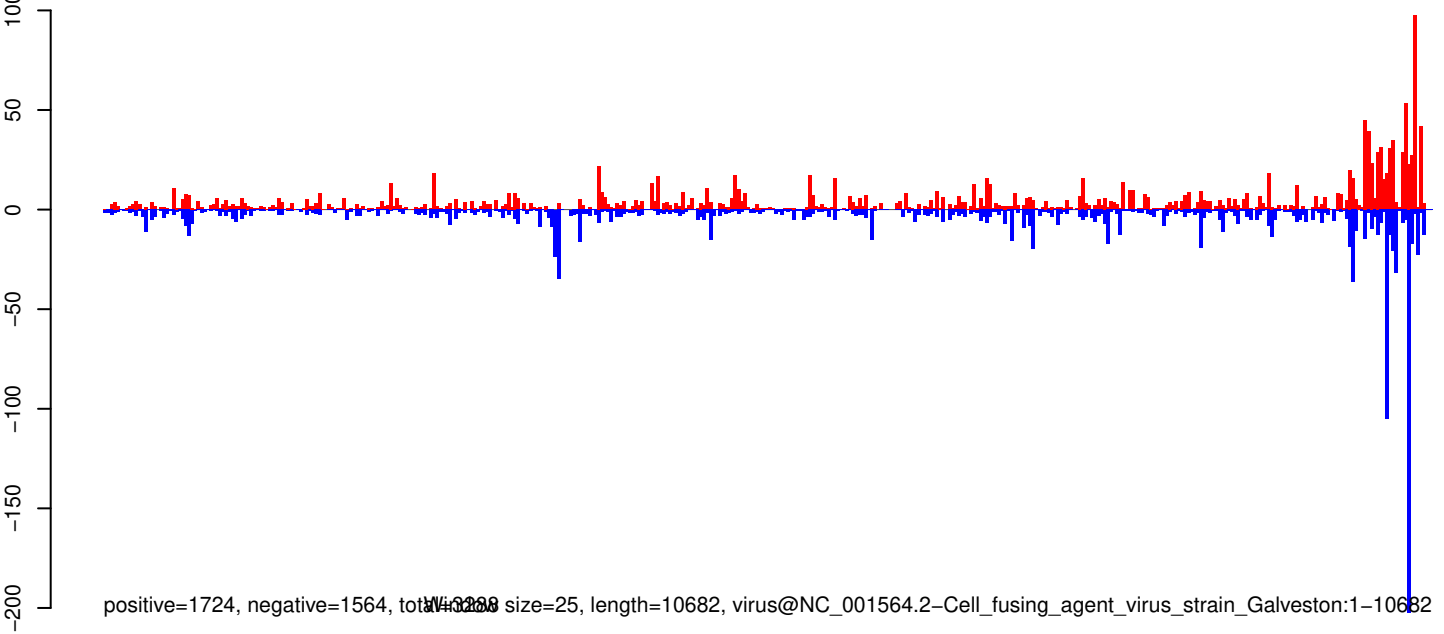
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

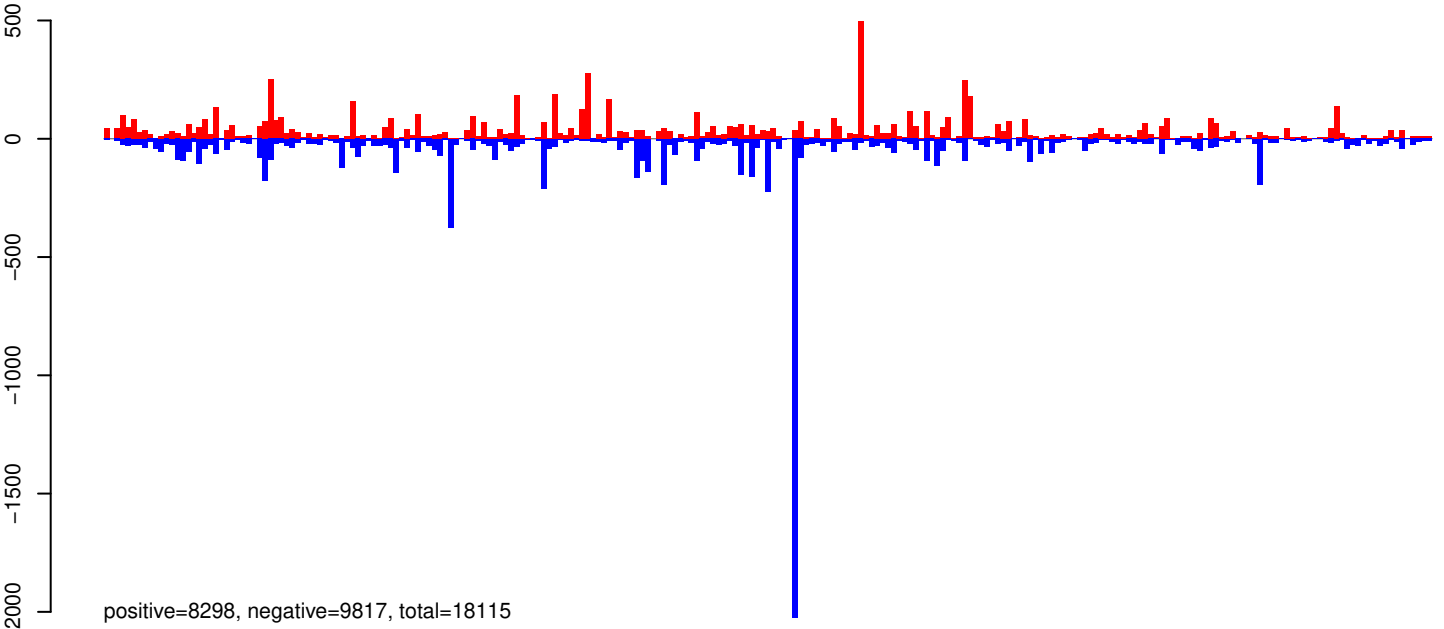


AeAeg_Aag2SINV_dsLuc.rep

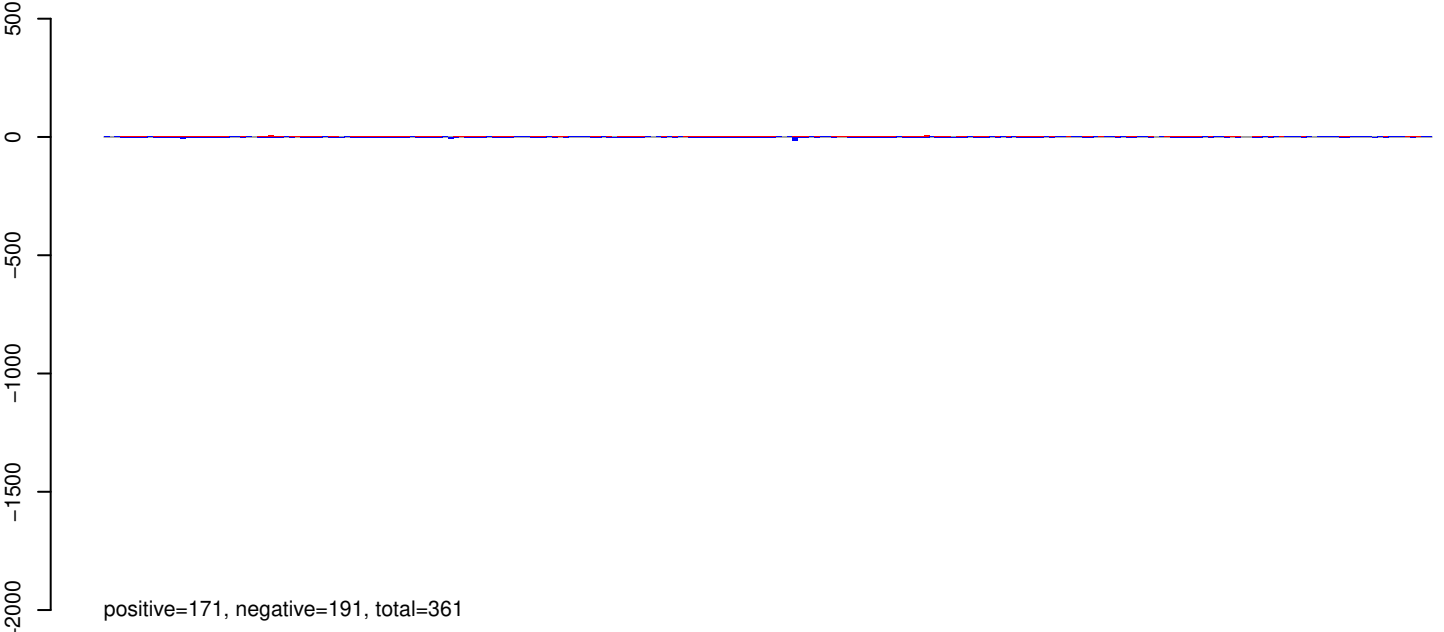


1143268 size=25, length=10682, virus@NC_001564.2-Cell_fusing_agent_virus_strain_Galveston:1-10682

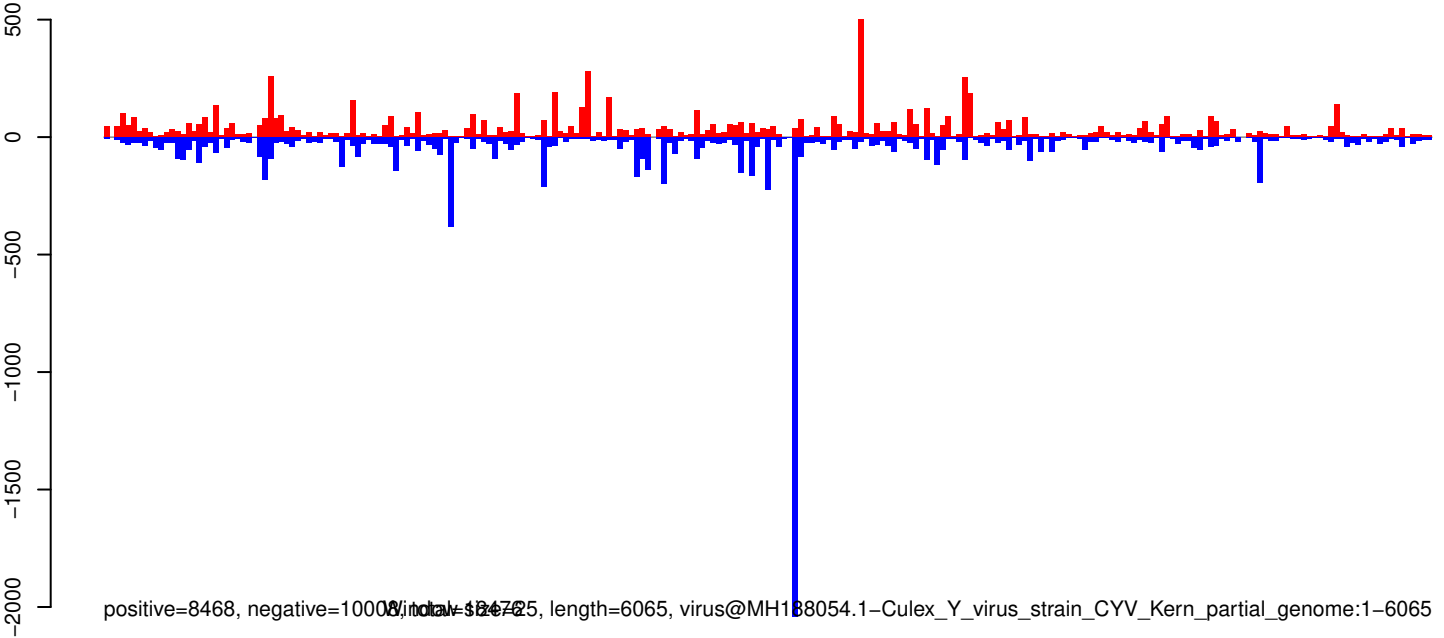
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

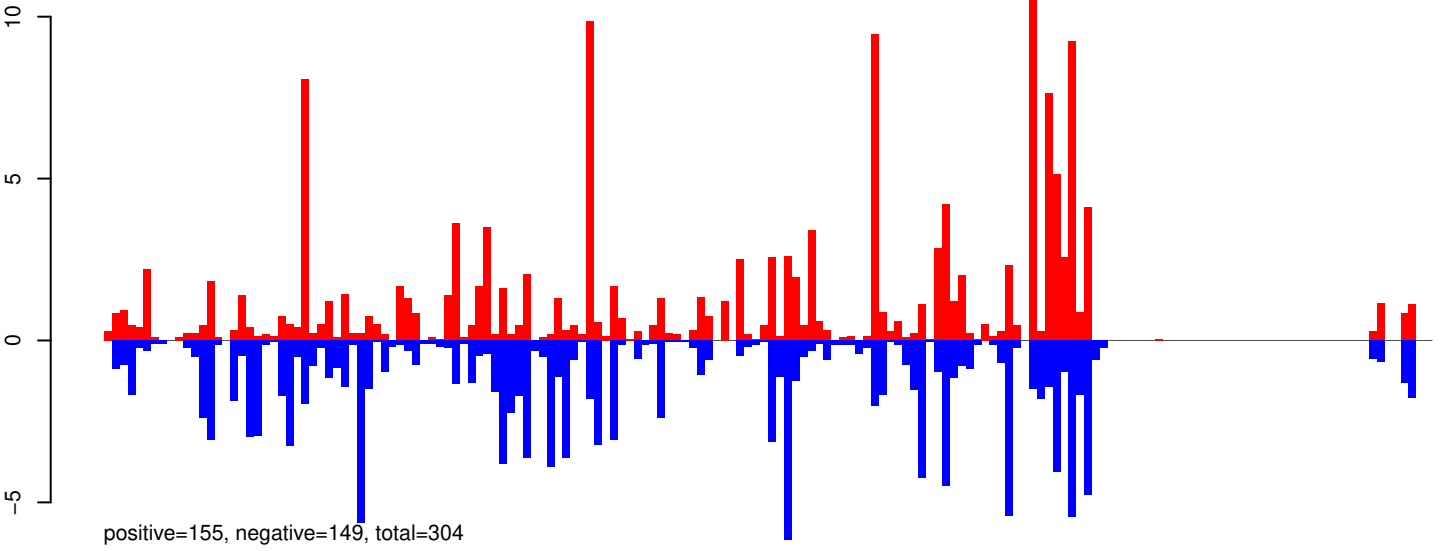


AeAeg_Aag2SINV_dsLuc.rep

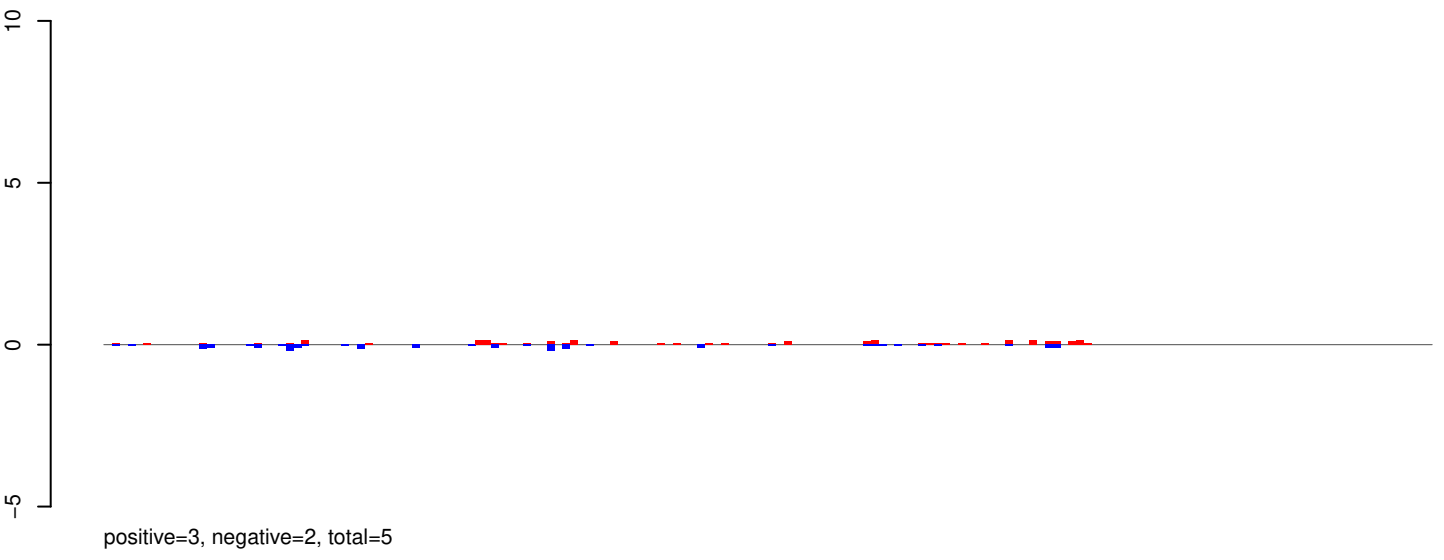


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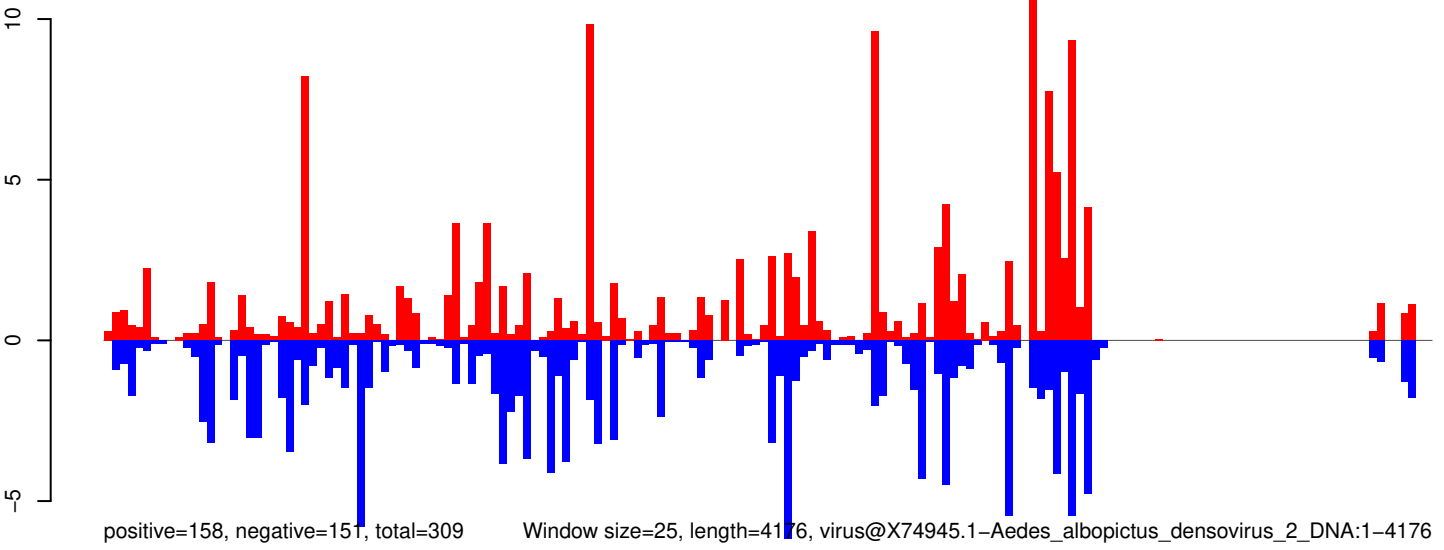
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



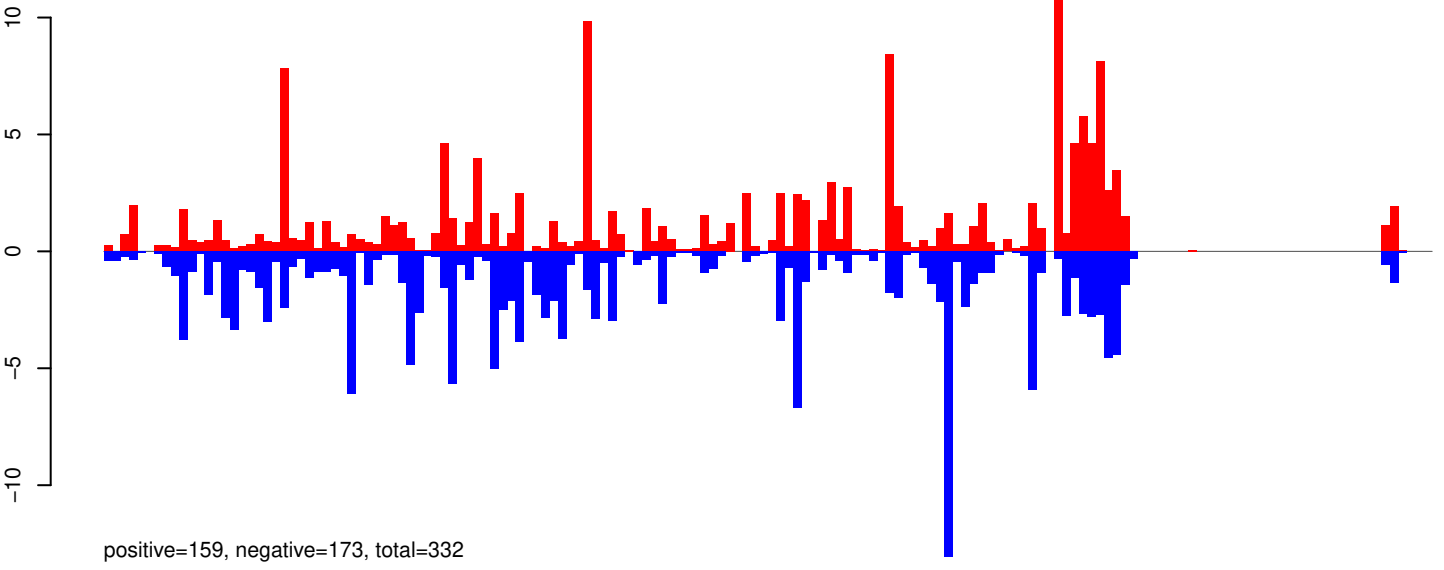
AeAeg_Aag2SINV_dsLuc.rep



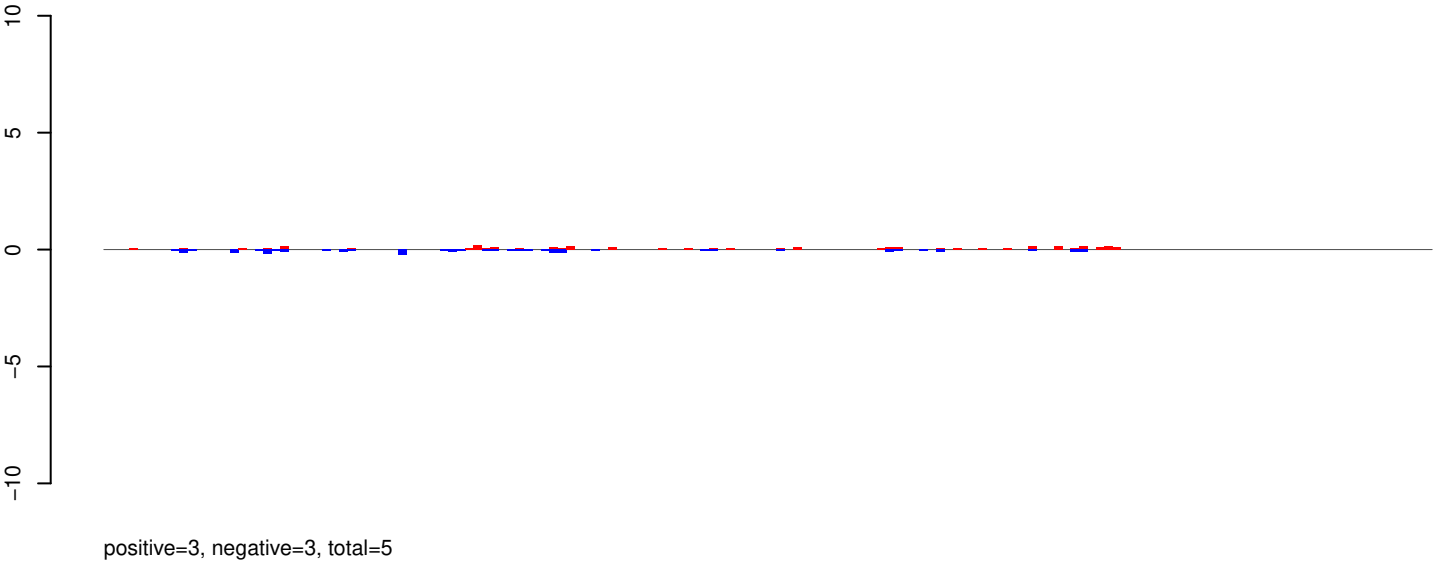
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0 1000 2000 3000 4000

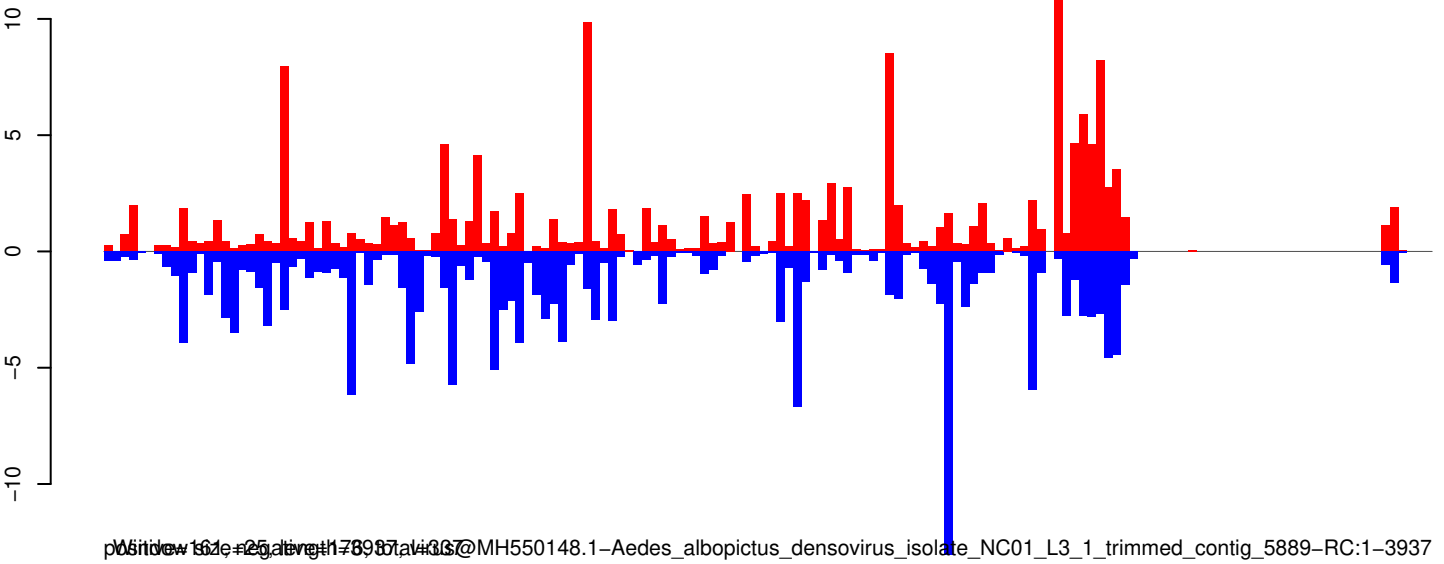
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep

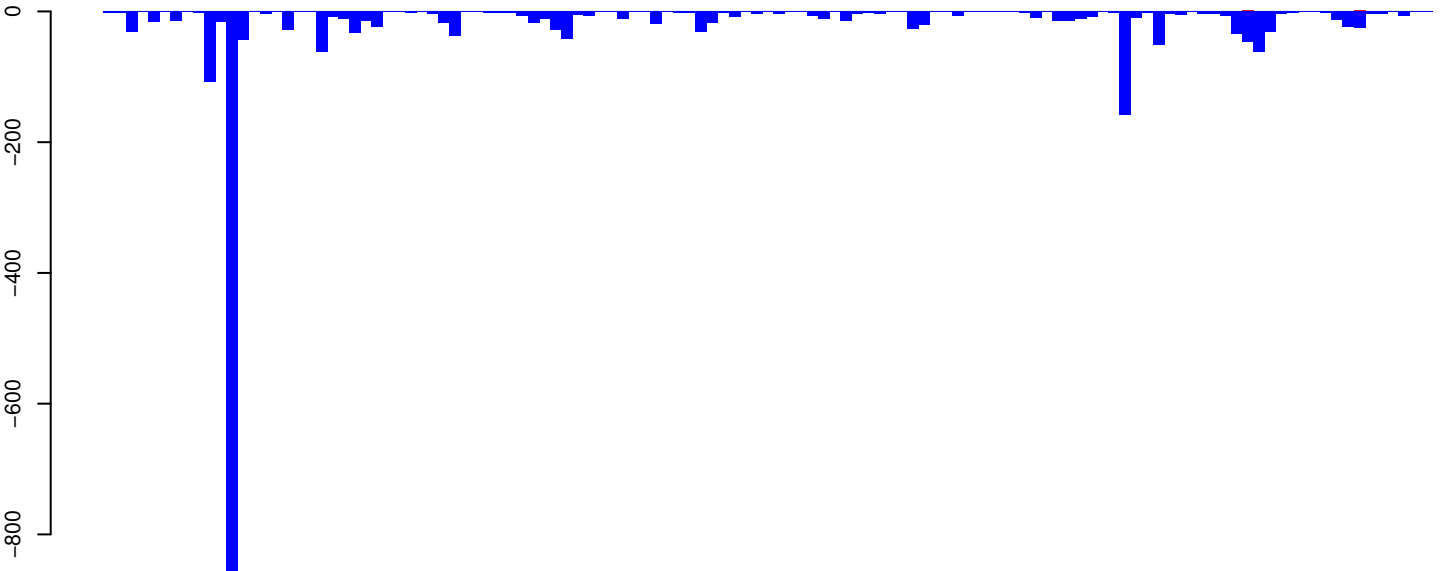


AeAeg_Aag2SINV_dsLuc.18_23.rep



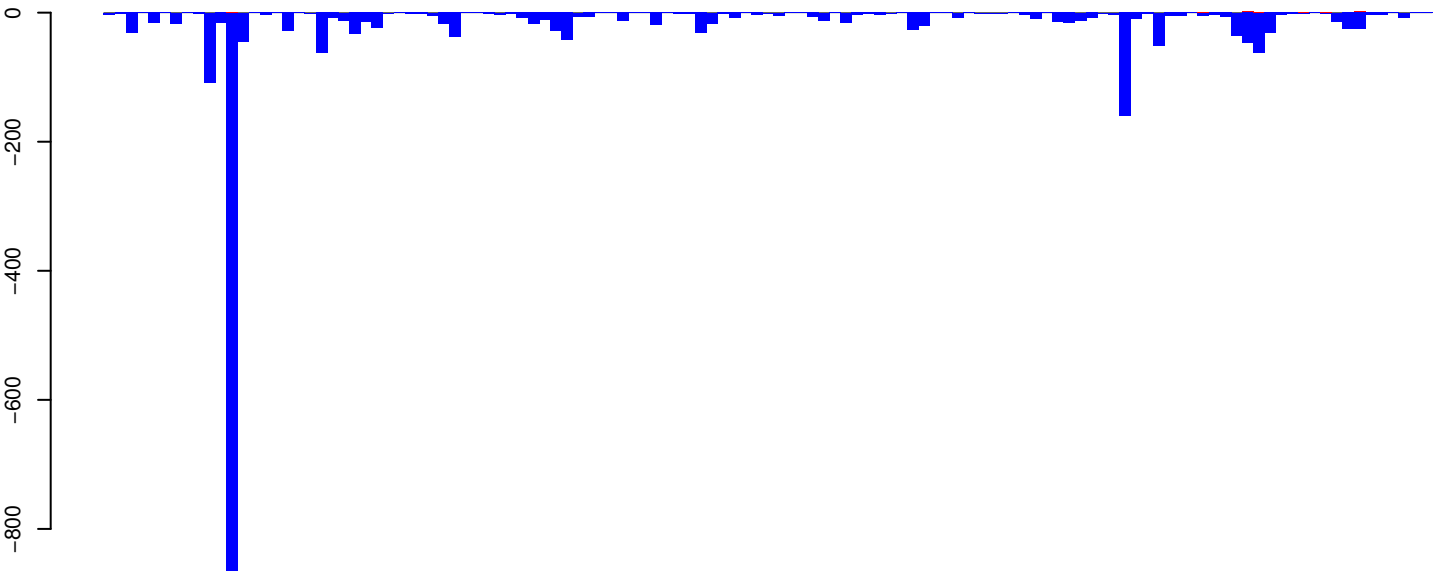
positive=1, negative=40, total=41

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=6, negative=2315, total=2320

AeAeg_Aag2SINV_dsLuc.rep

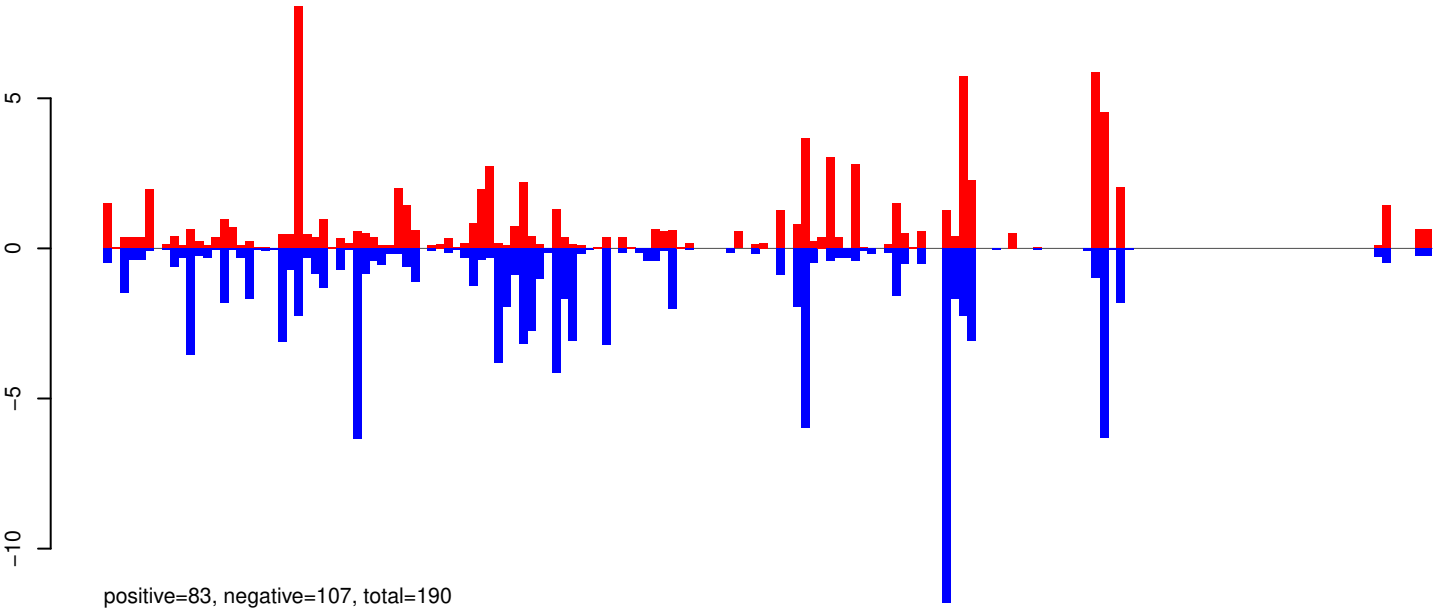


positive=7, negative=2354, total=2361

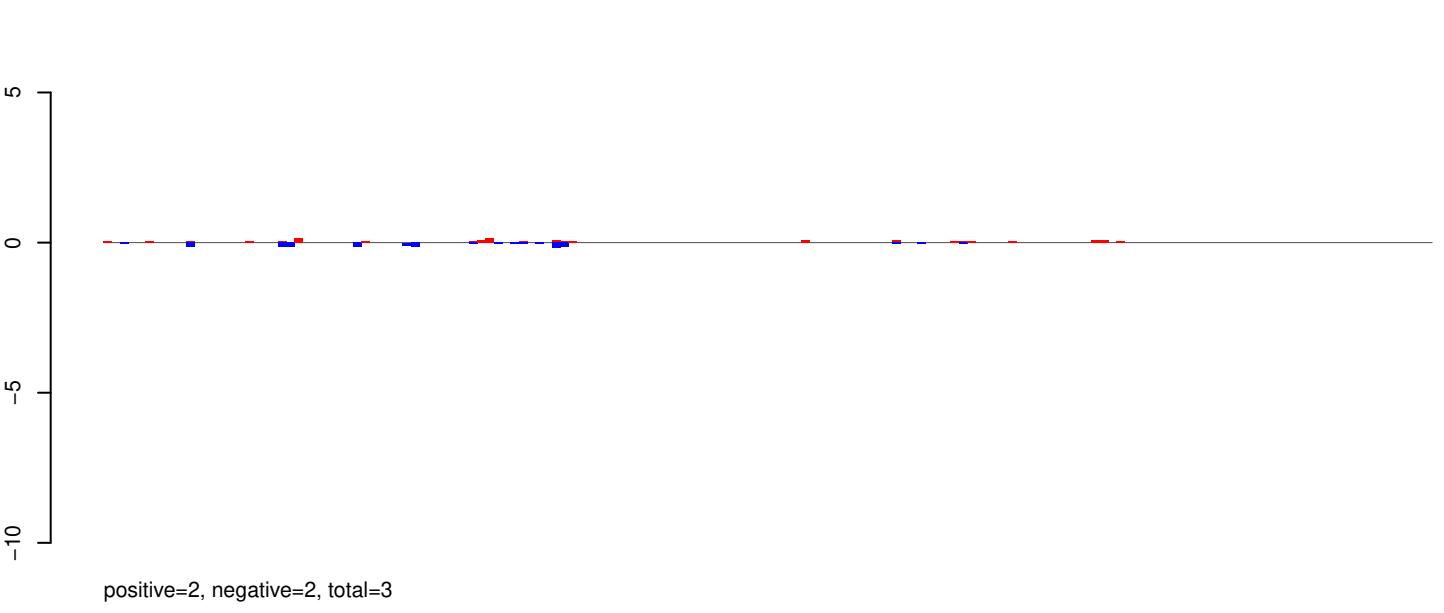
Window size=25, length=2965, virus@AY347953.2-Aedes_aegypti_genomic_sequence:1-2965

0 500 1000 1500 2000 2500 3000

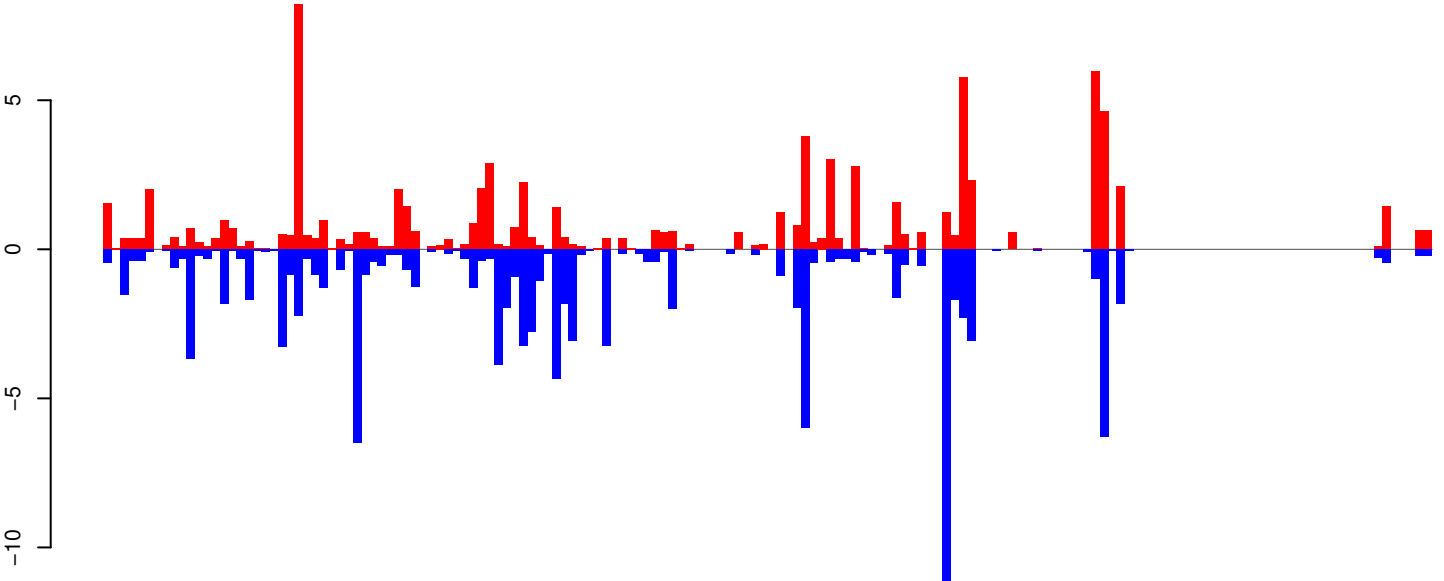
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



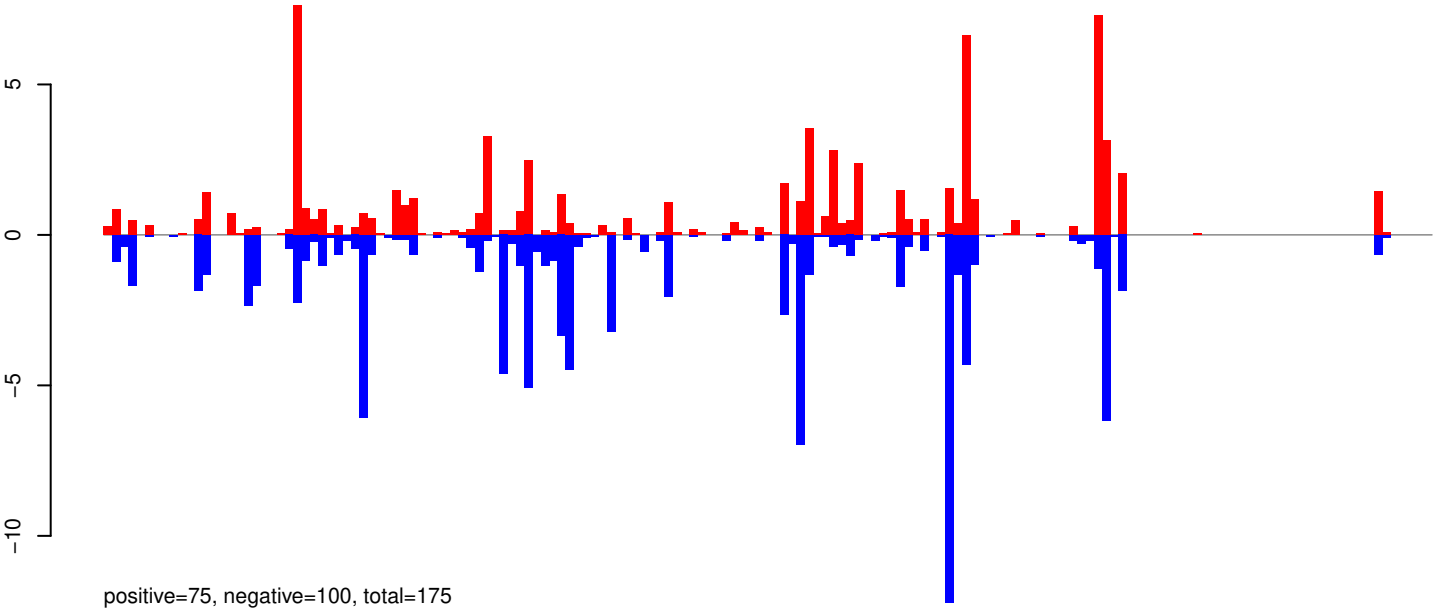
AeAeg_Aag2SINV_dsLuc.rep



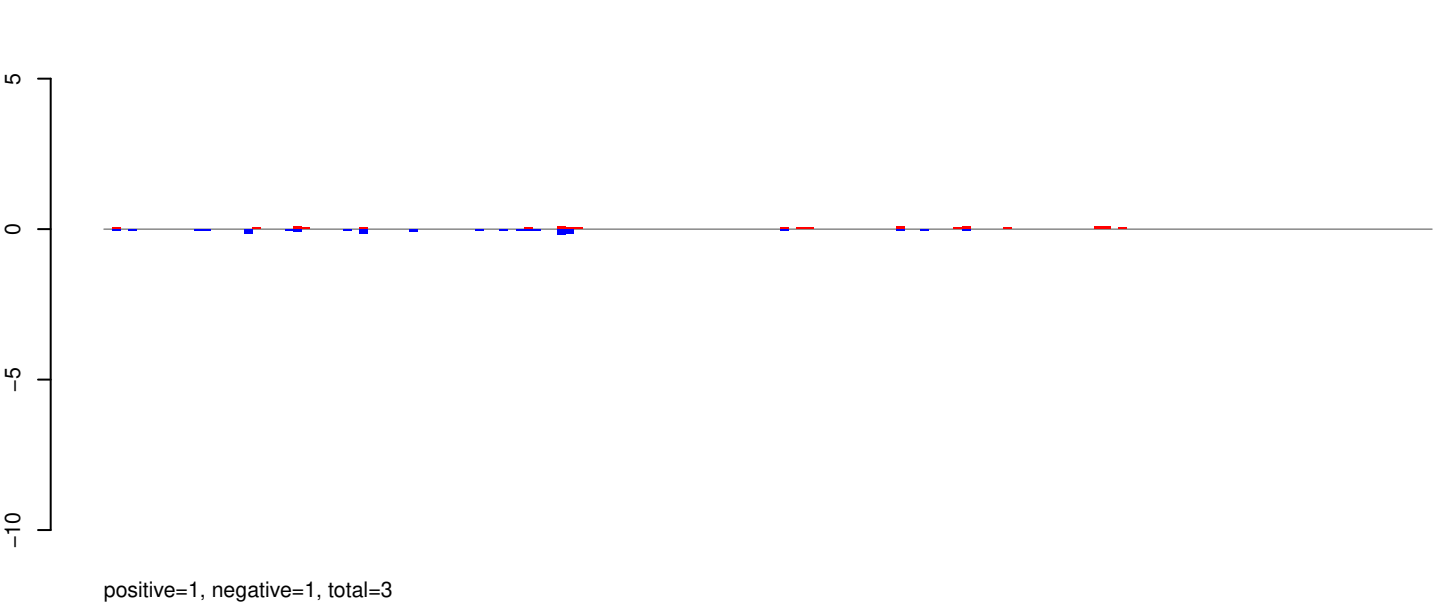
www.ebi.ac.uk/Ensembl/Genomes/100/AY010505.1-Haemagogus_equinus_densovirus_nonstructural_protein_1_nonstructural_protein_2_and_c:1-3986

0 1000 2000 3000 4000

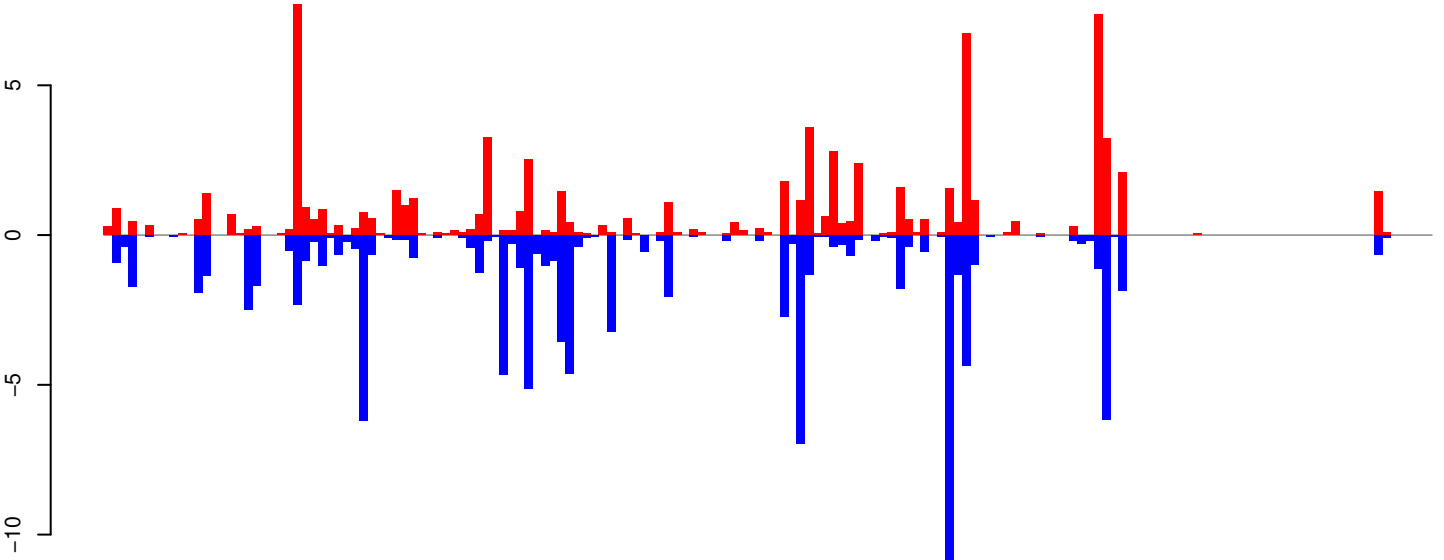
AeAeg_Aag2SINV_dsLuc.18_23.rep



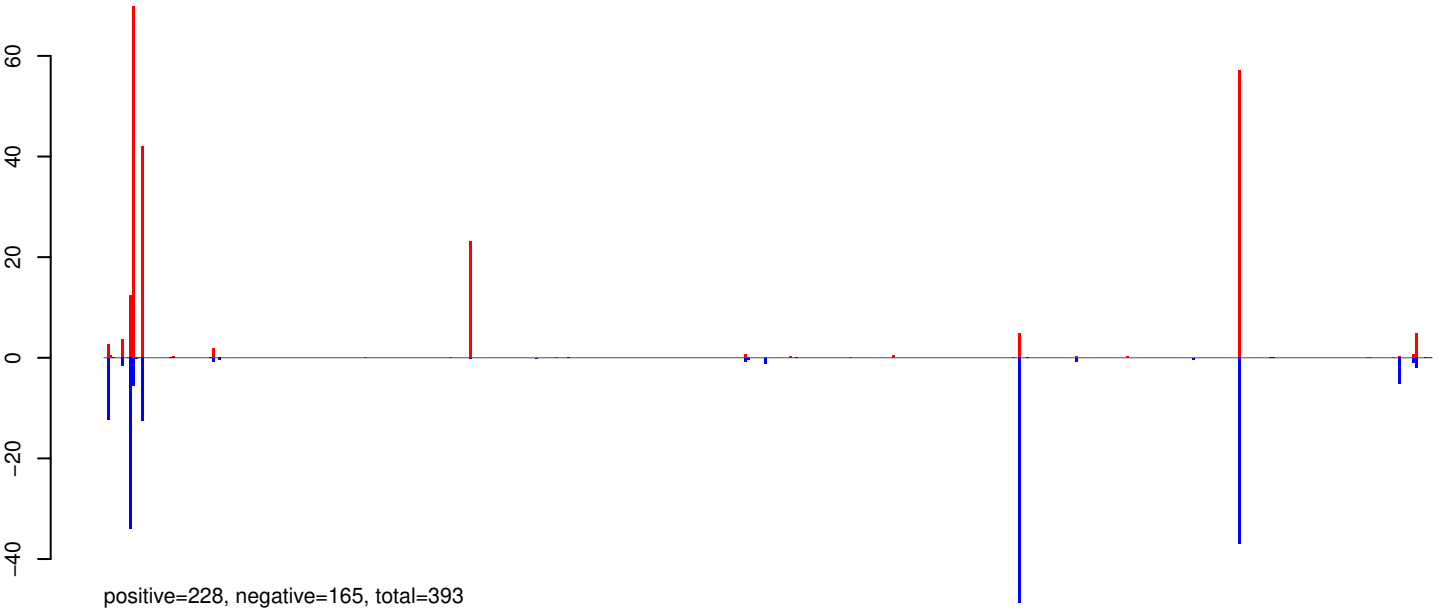
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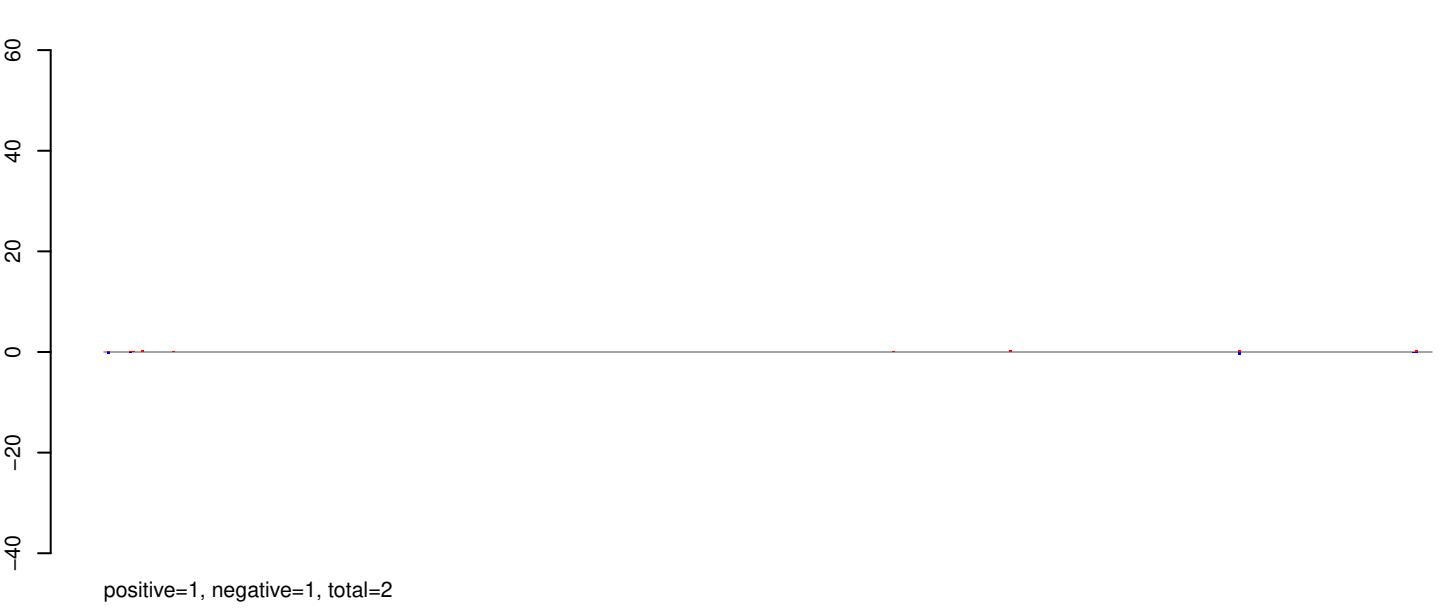
AeAeg_Aag2SINV_dsLuc.rep



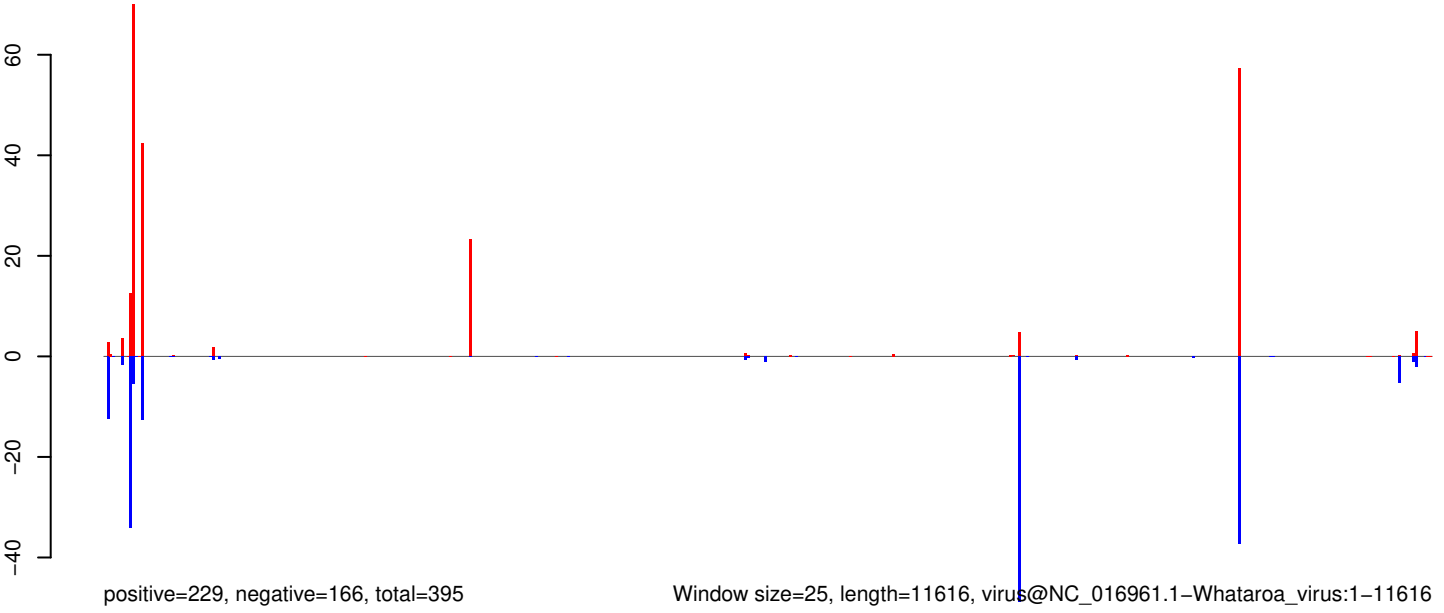
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



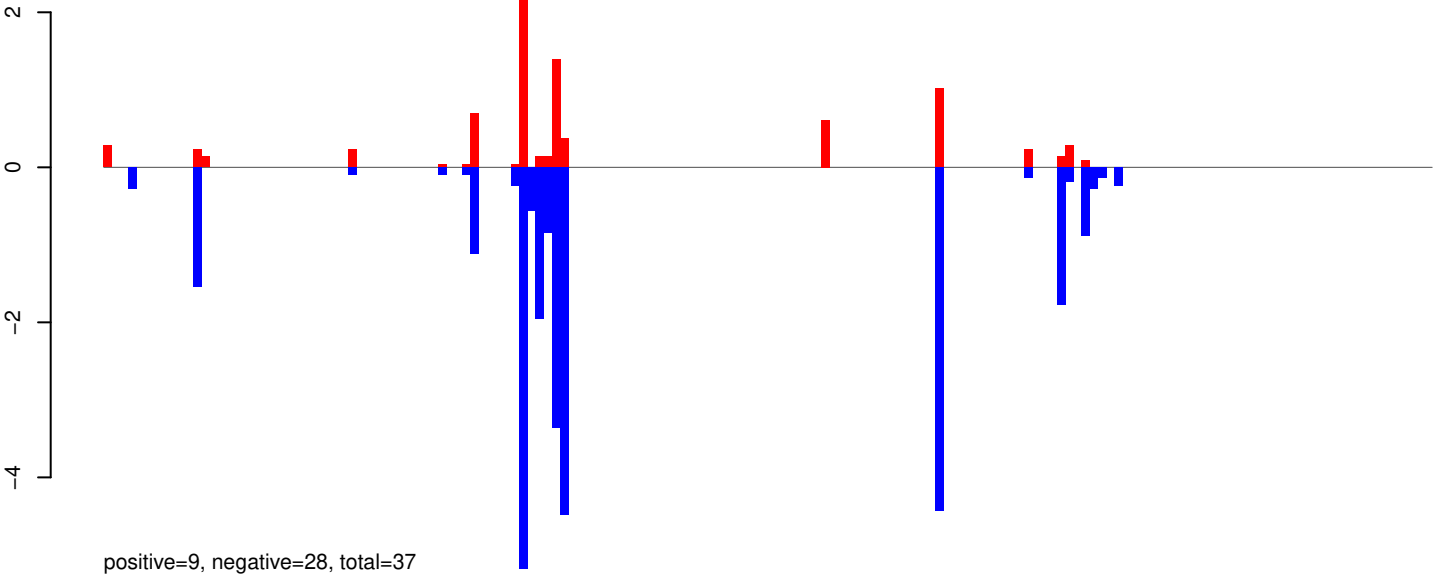
AeAeg_Aag2SINV_dsLuc.rep



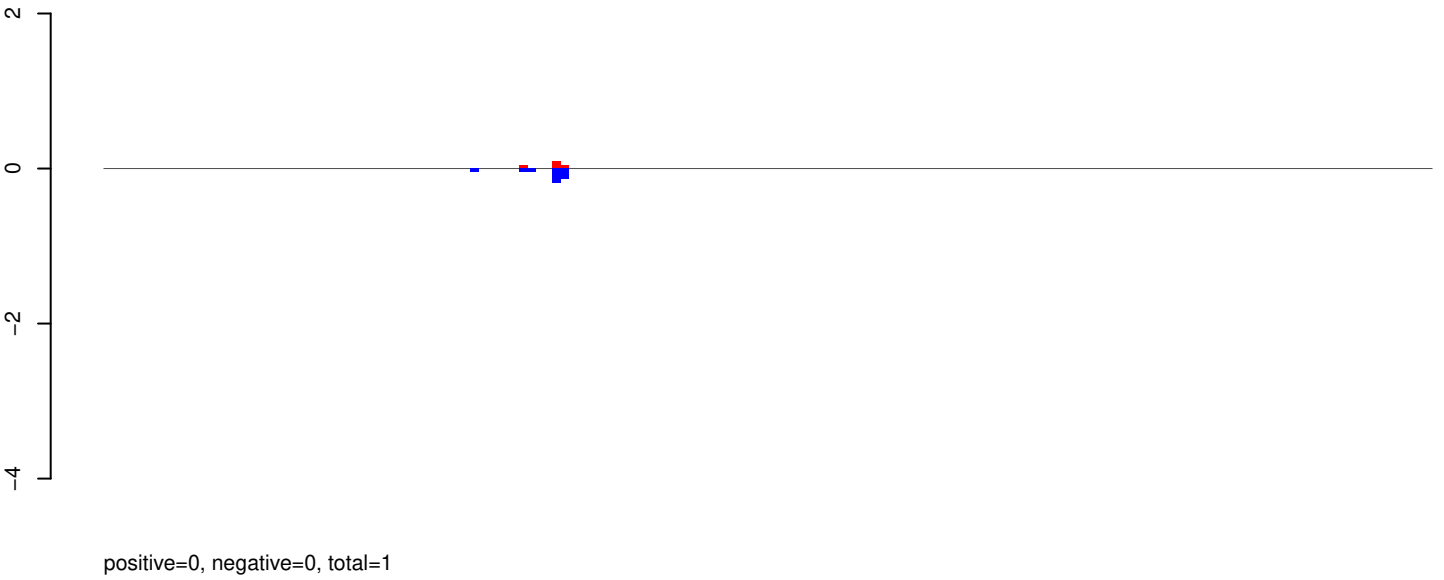
Window size=25, length=11616, virus@NC_016961.1-Whataroa_virus:1-11616

0 2000 4000 6000 8000 10000 12000

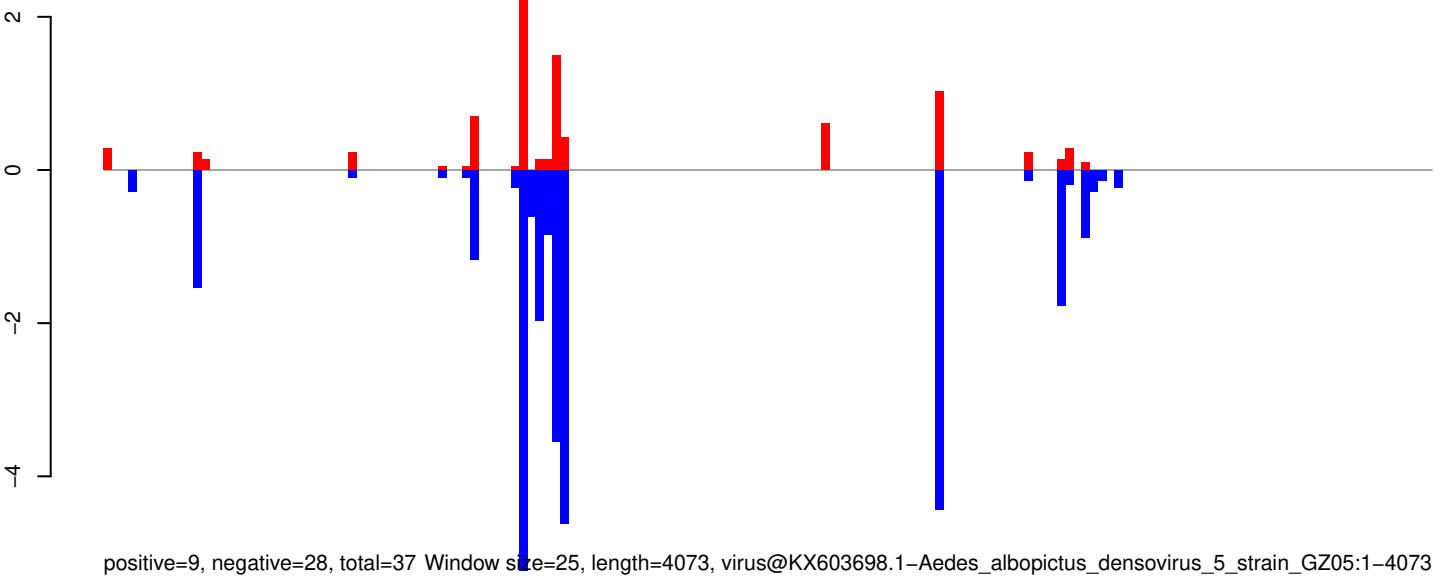
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

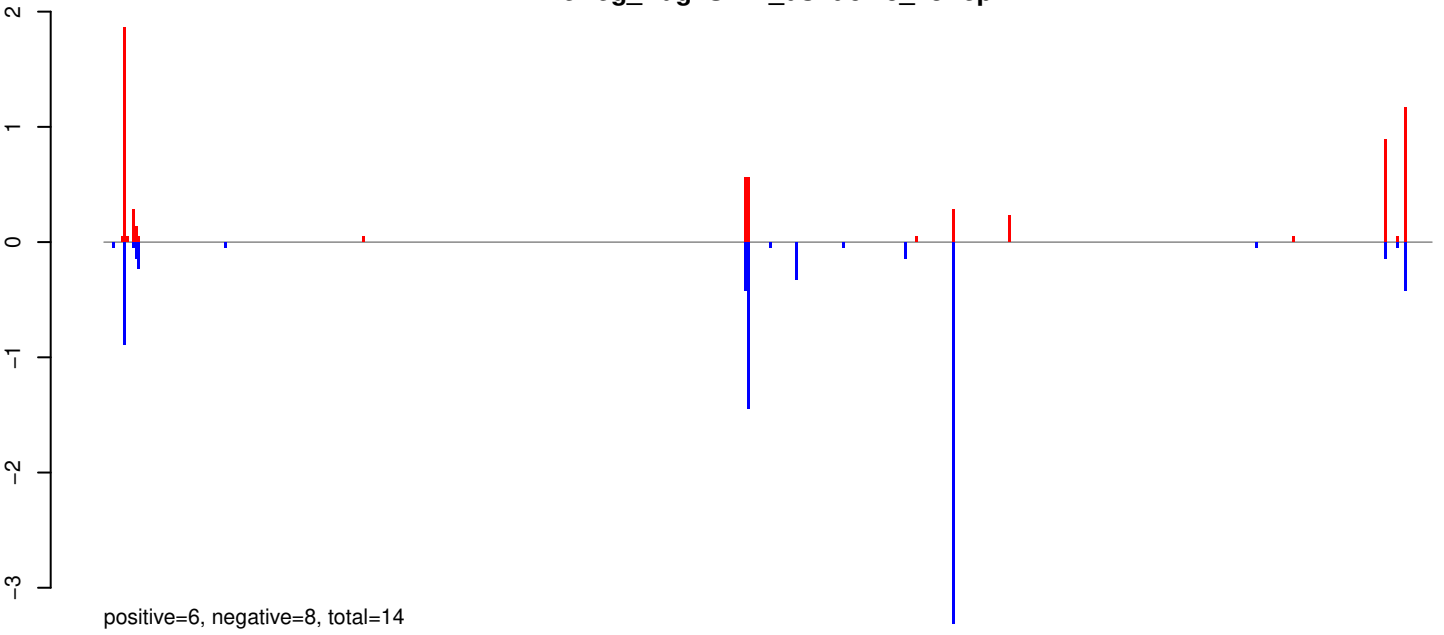


AeAeg_Aag2SINV_dsLuc.rep

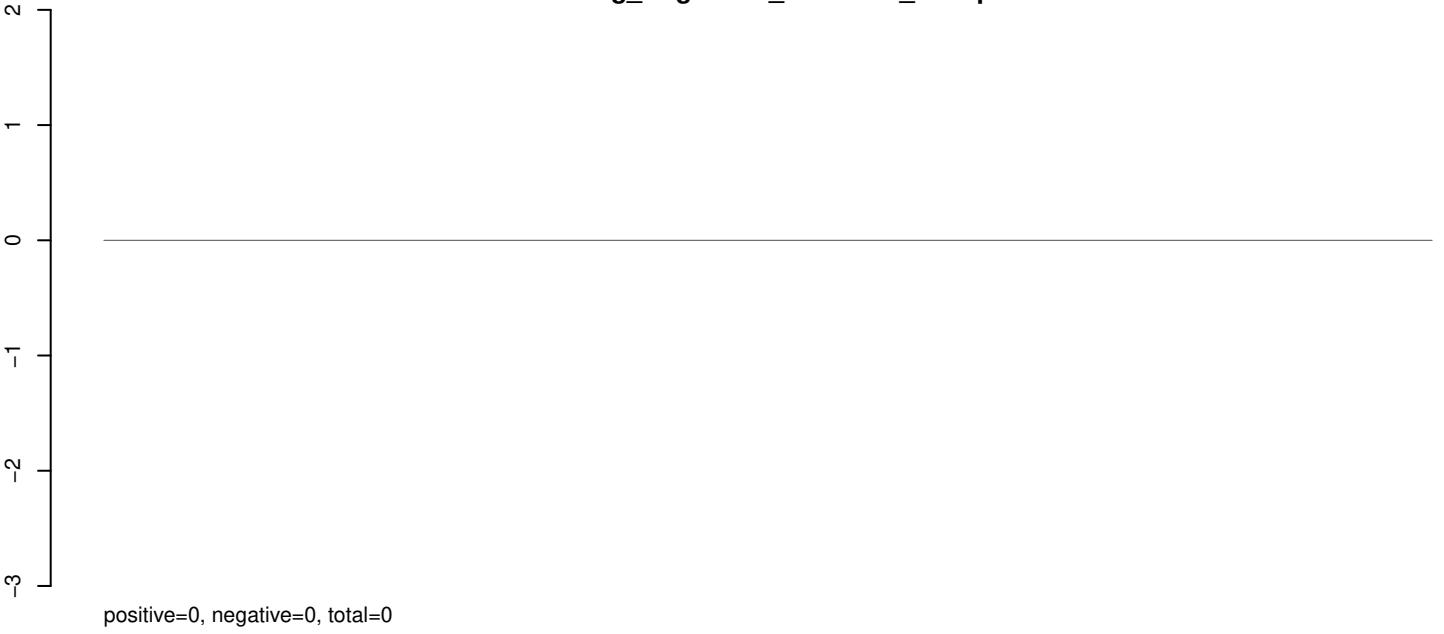


0 1000 2000 3000 4000

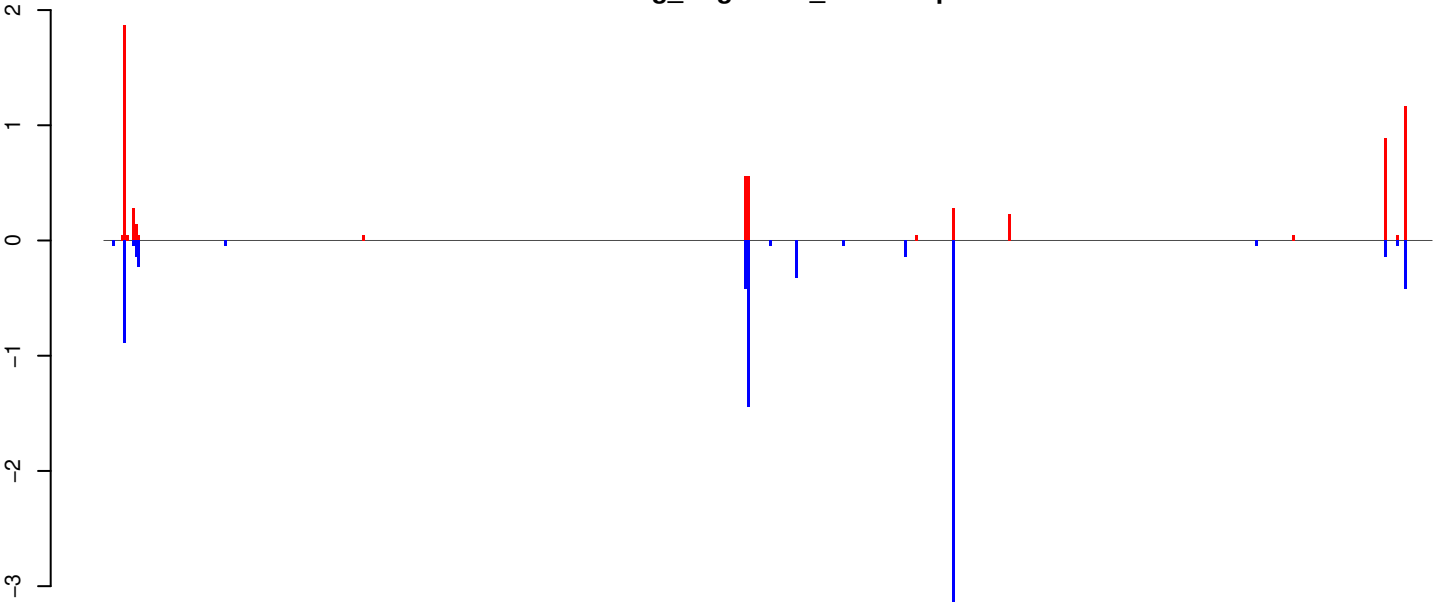
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



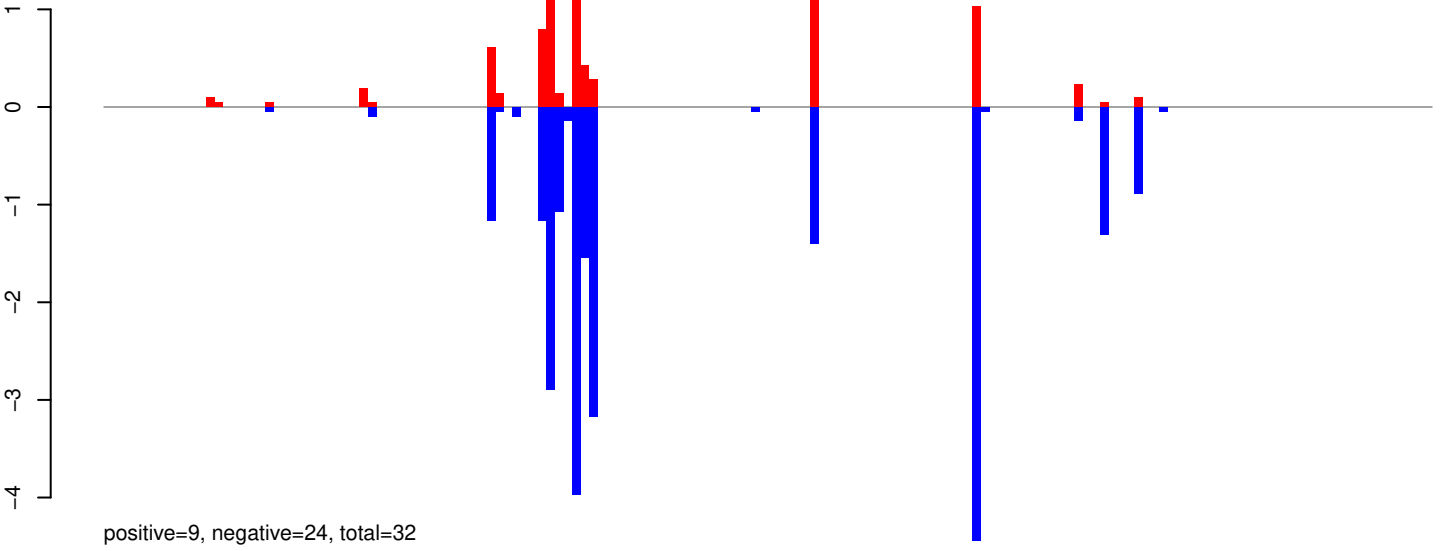
AeAeg_Aag2SINV_dsLuc.rep



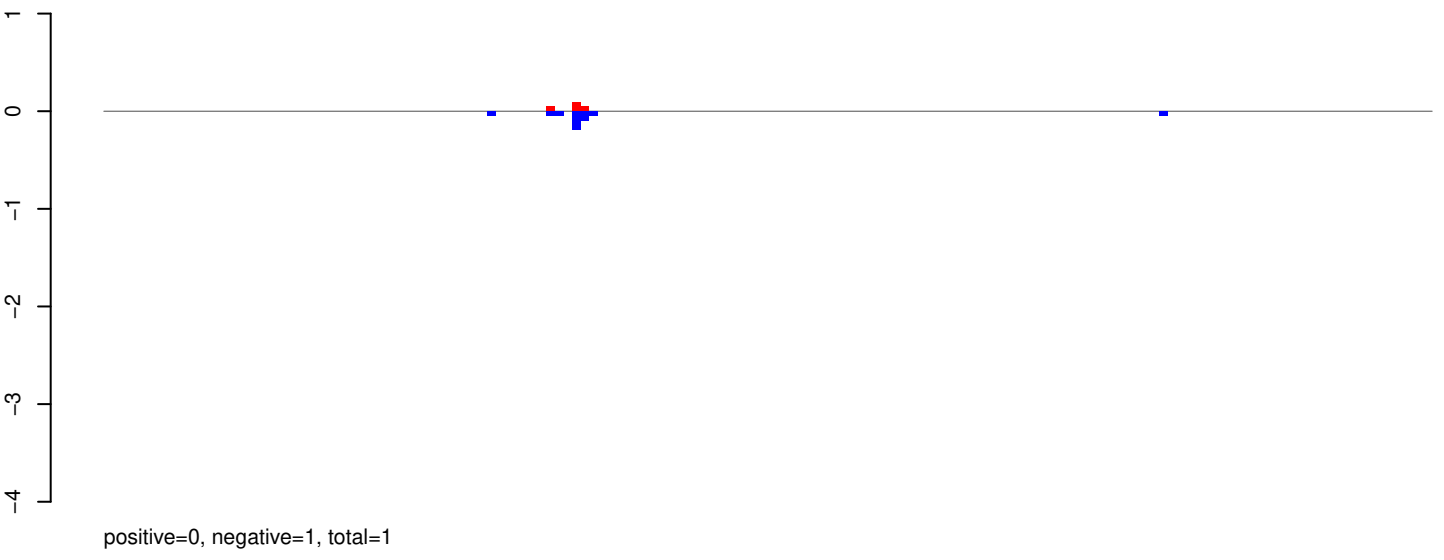
Window size=25, length=11824, virus@NC_003900.1-Aura_virus:1-11824

0 2000 4000 6000 8000 10000 12000

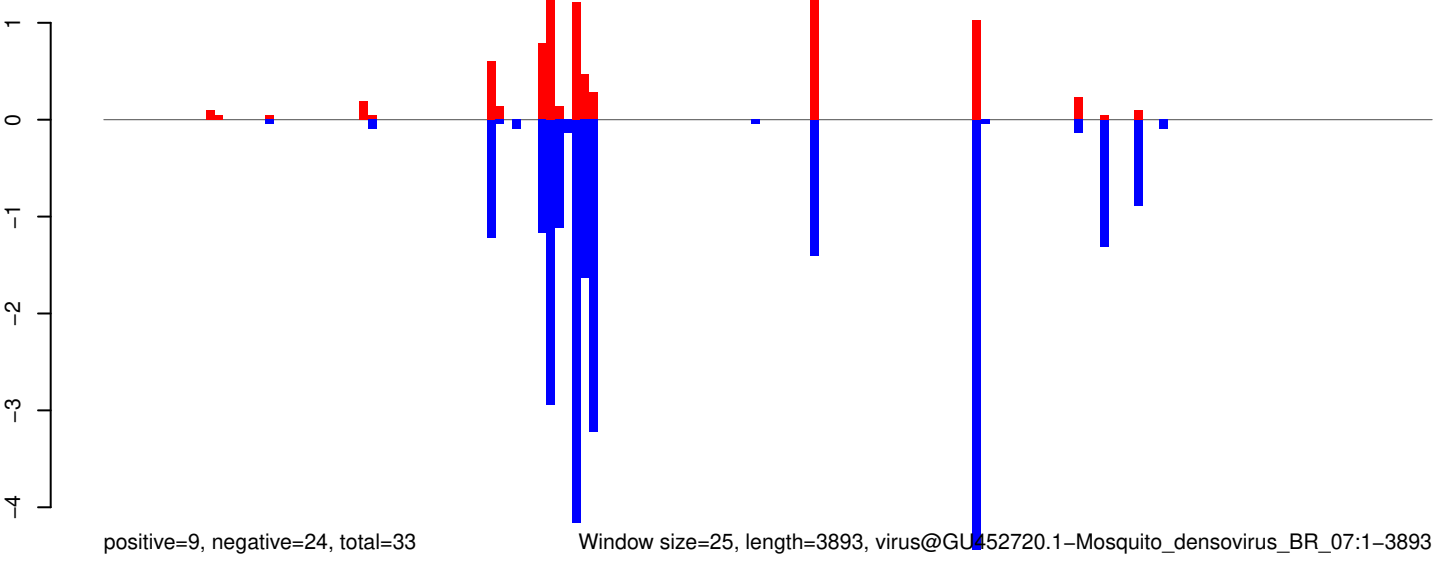
AeAeg_Aag2SINV_dsLuc.18_23.rep



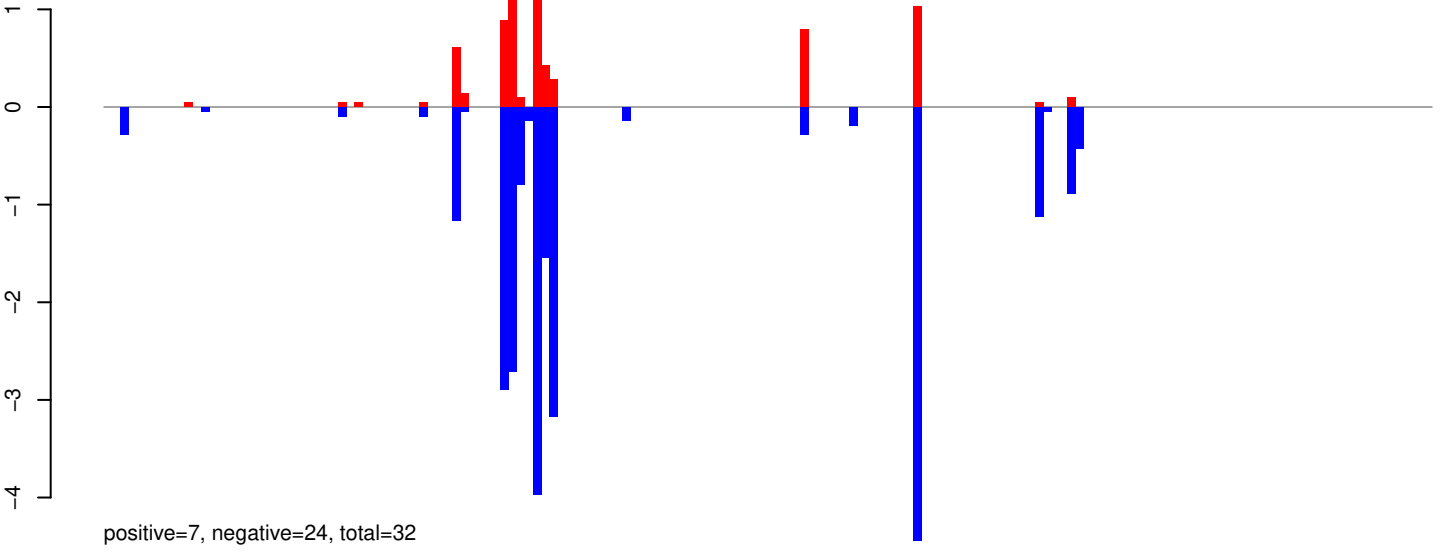
AeAeg_Aag2SINV_dsLuc.24_35.rep



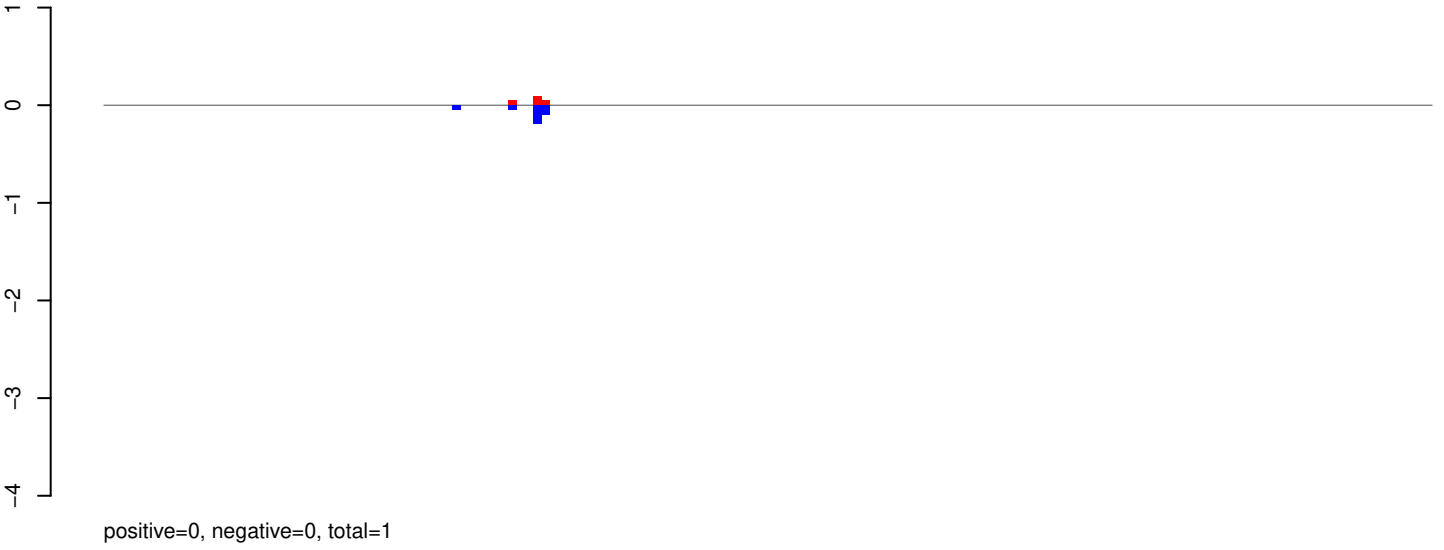
AeAeg_Aag2SINV_dsLuc.rep



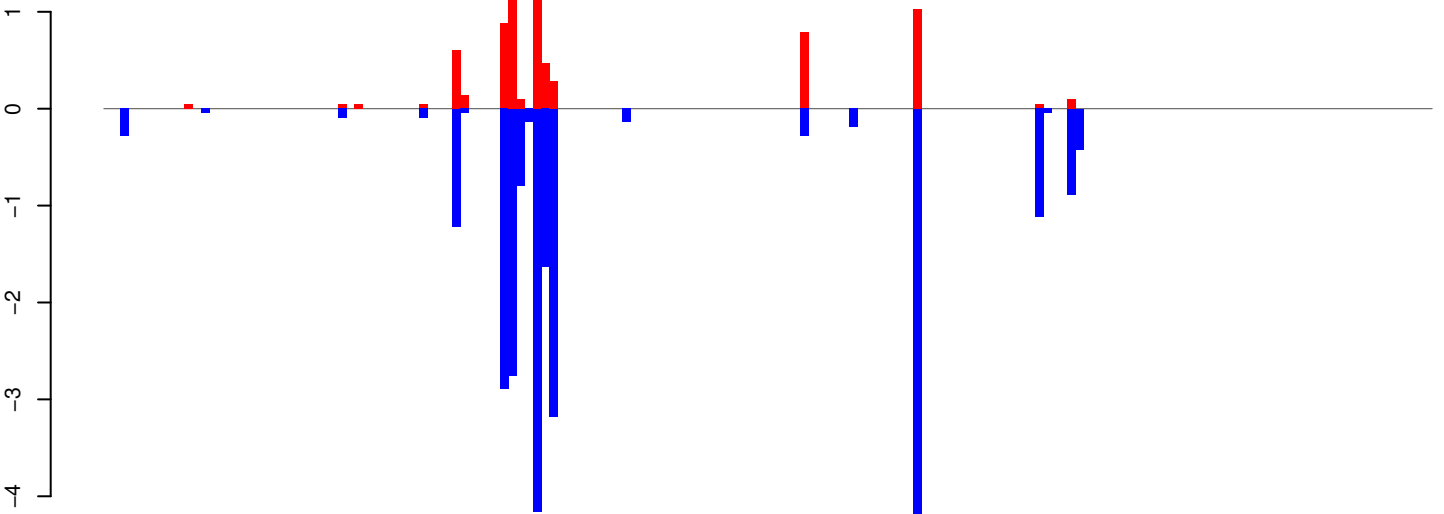
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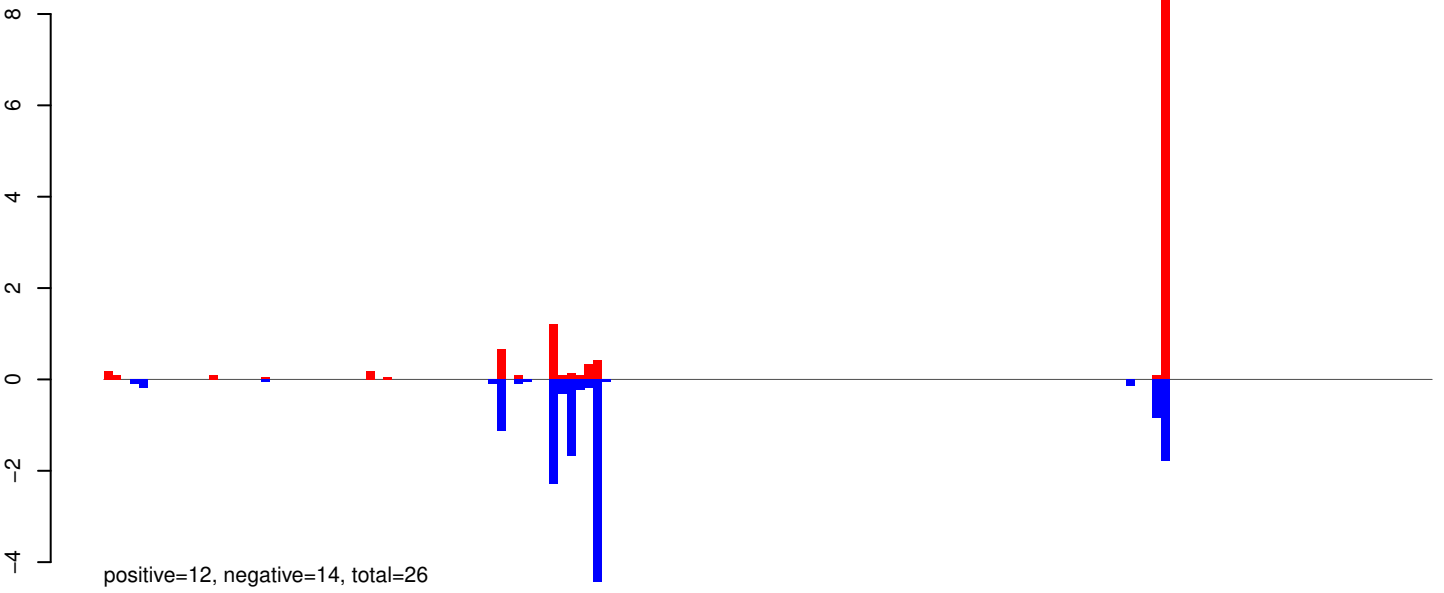
AeAeg_Aag2SINV_dsLuc.24_35.rep



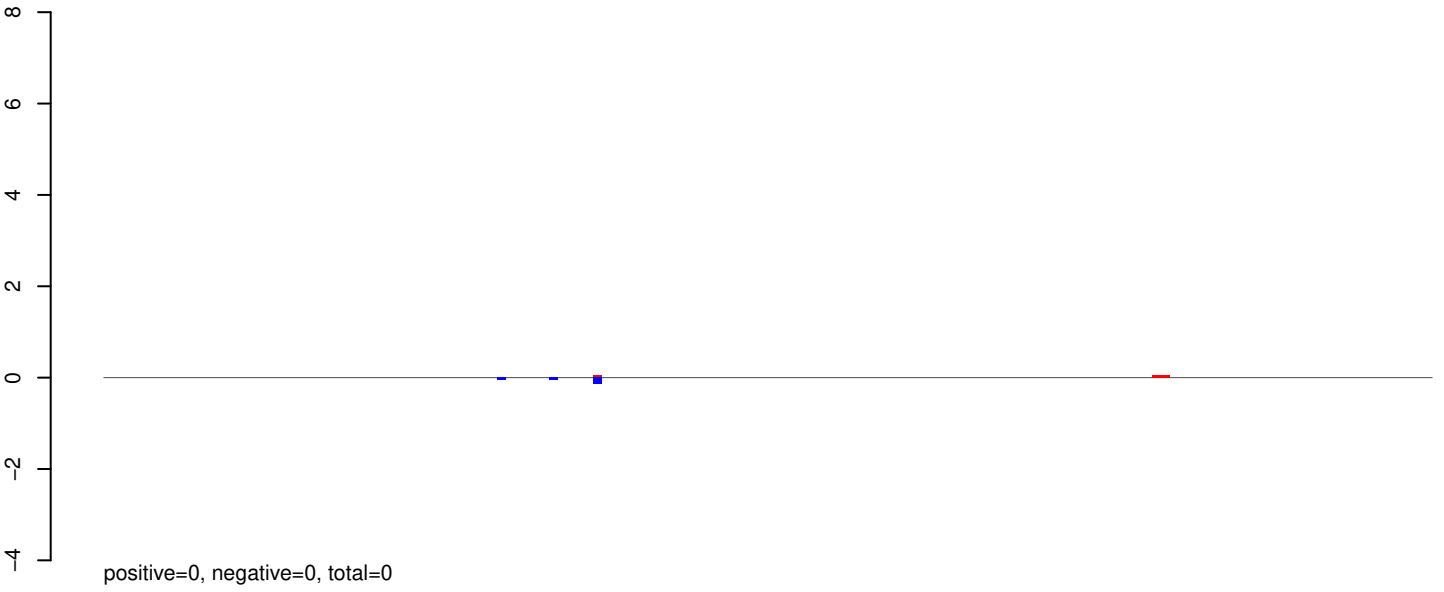
AeAeg_Aag2SINV_dsLuc.rep



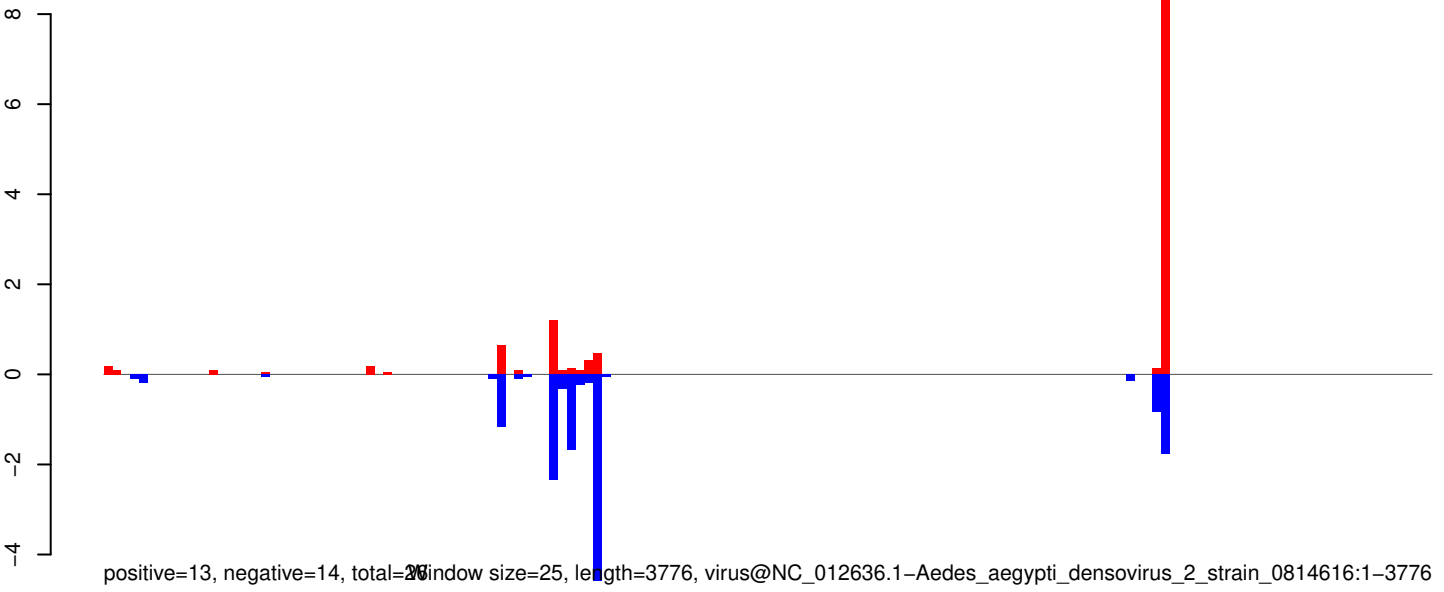
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

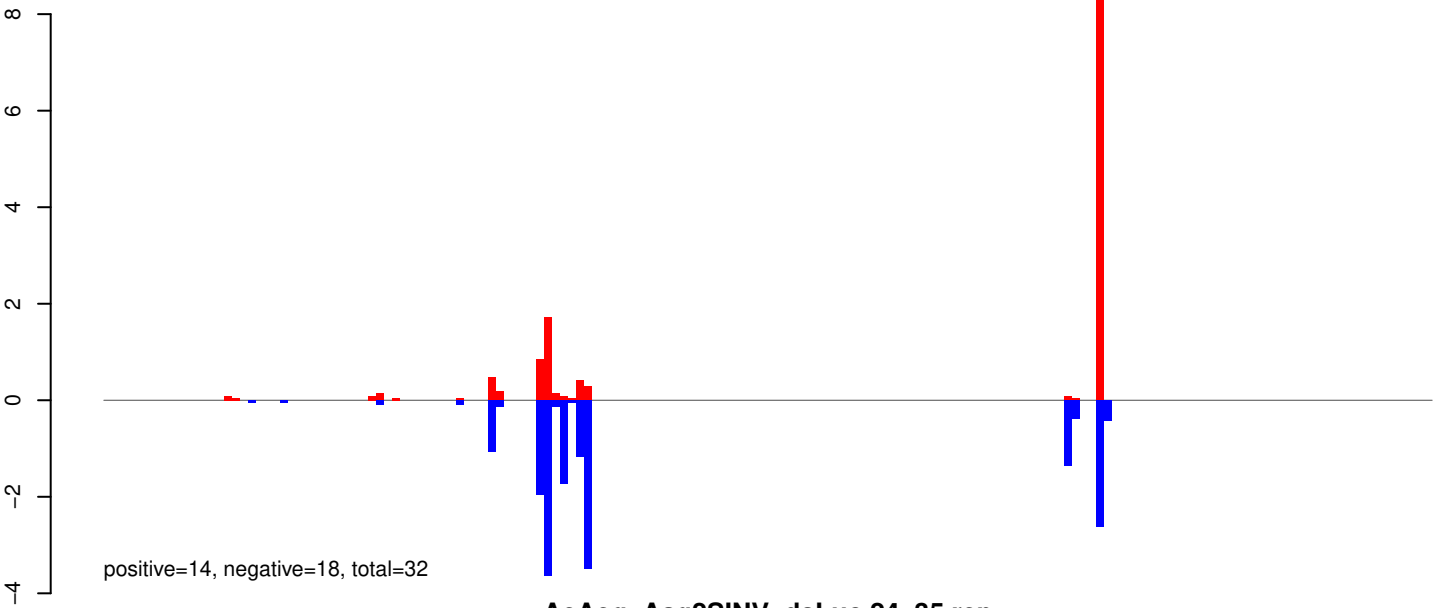


AeAeg_Aag2SINV_dsLuc.rep

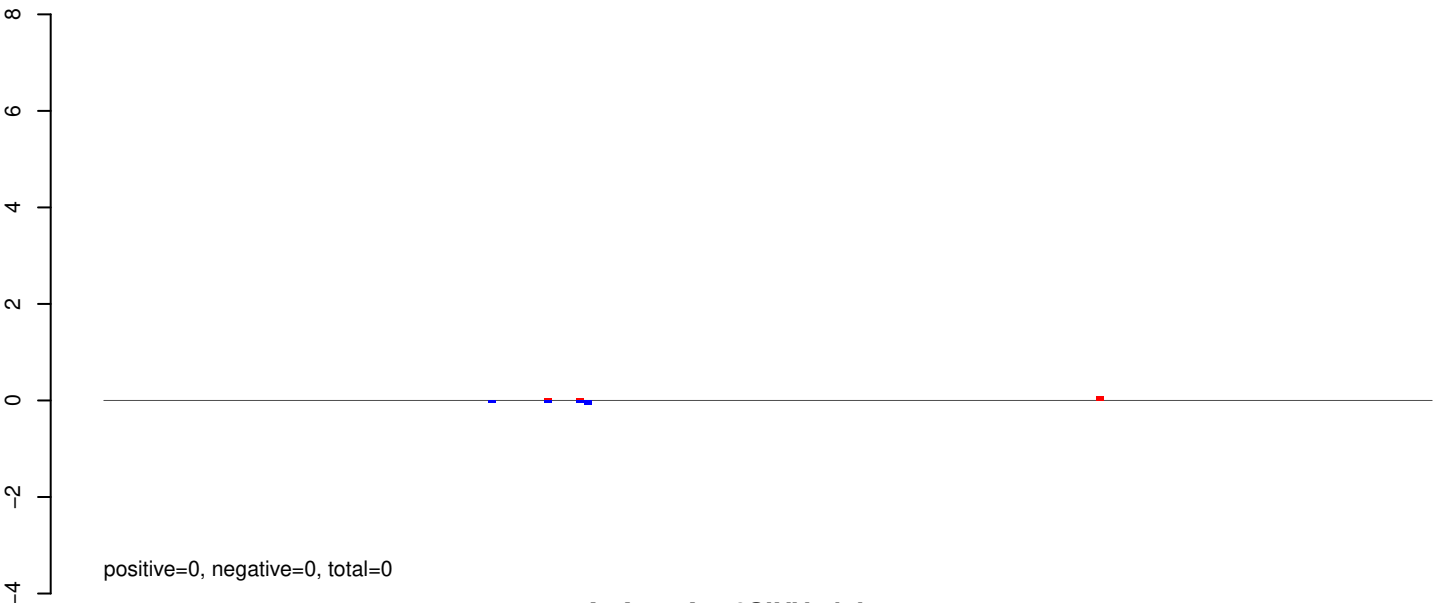


Window size=25, length=3776, virus@NC_012636.1-Aedes_aegypti_densovirus_2_strain_0814616:1-3776

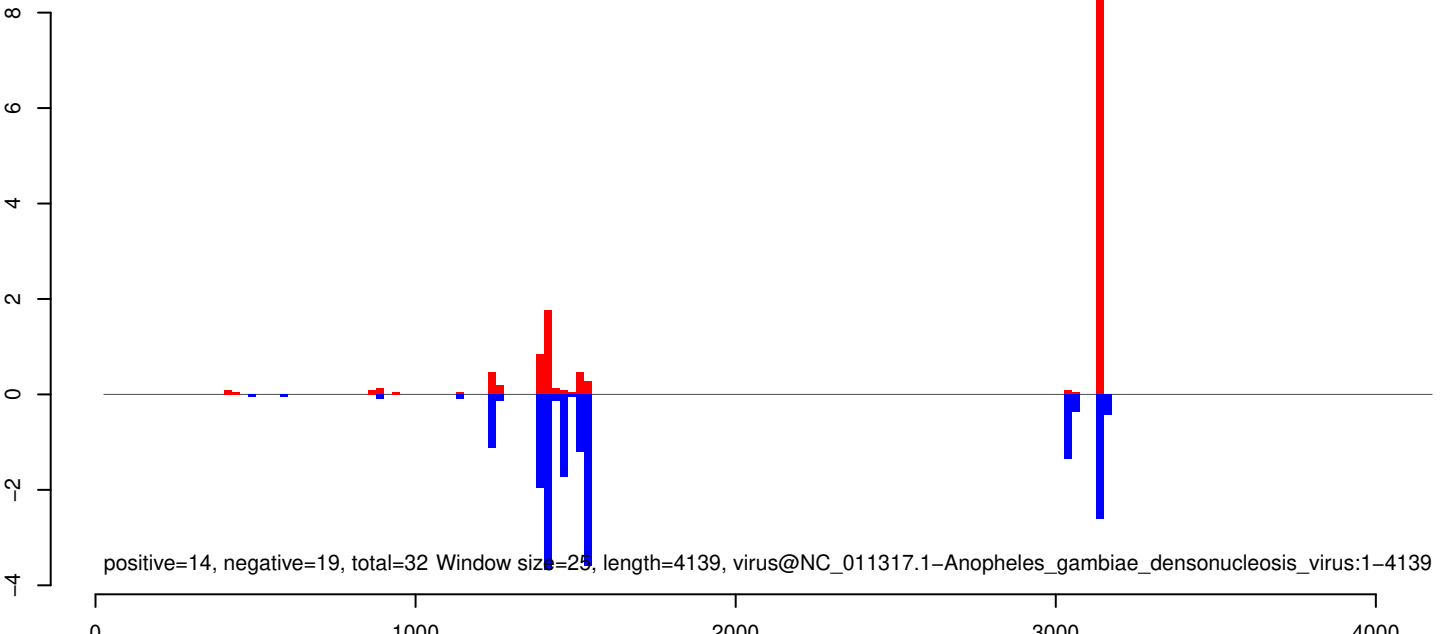
AeAeg_Aag2SINV_dsLuc.18_23.rep



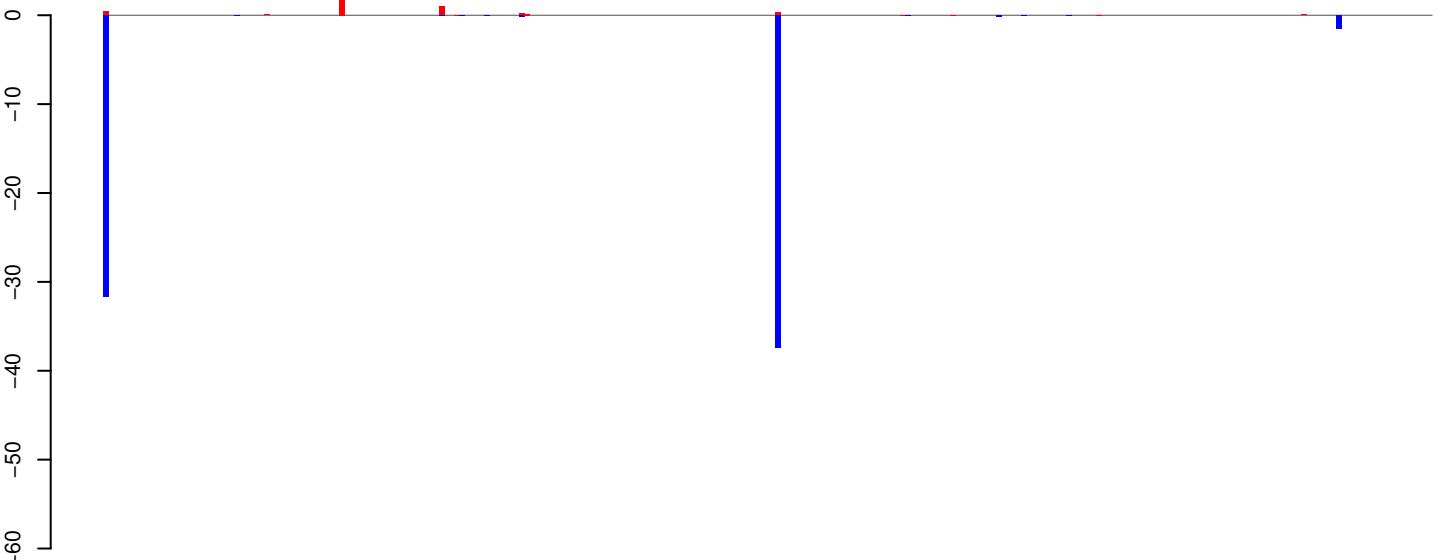
AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep

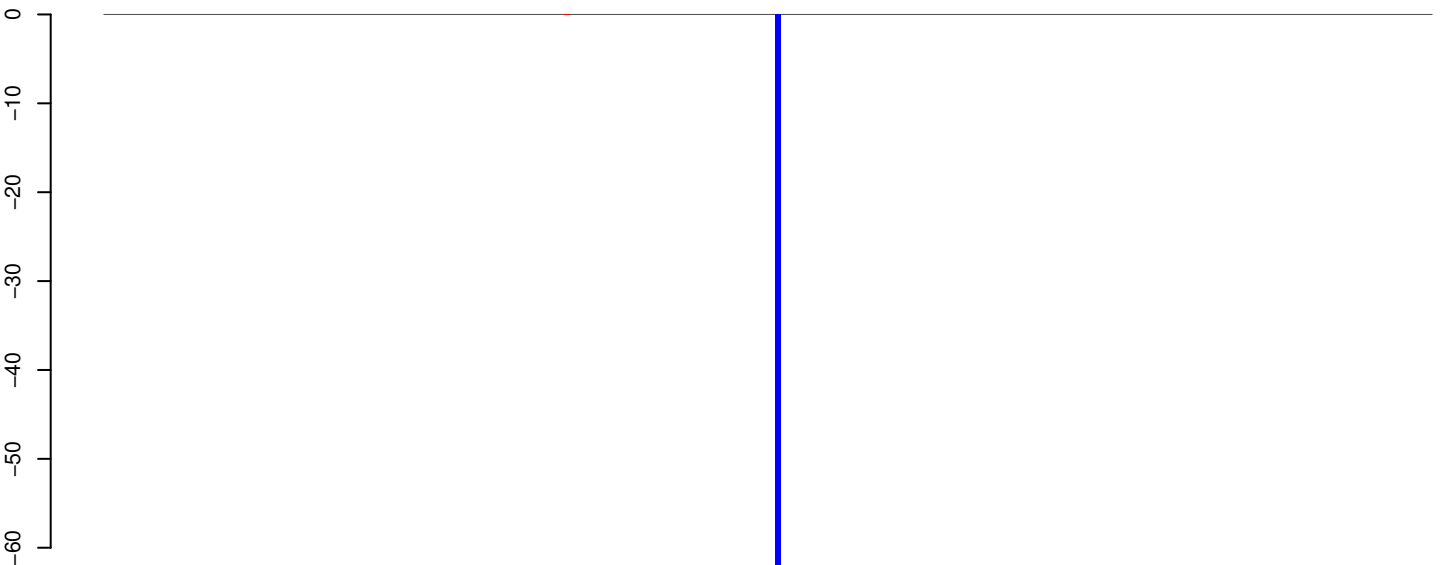


AeAeg_Aag2SINV_dsLuc.18_23.rep



positive=5, negative=71, total=76

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=0, negative=65, total=65

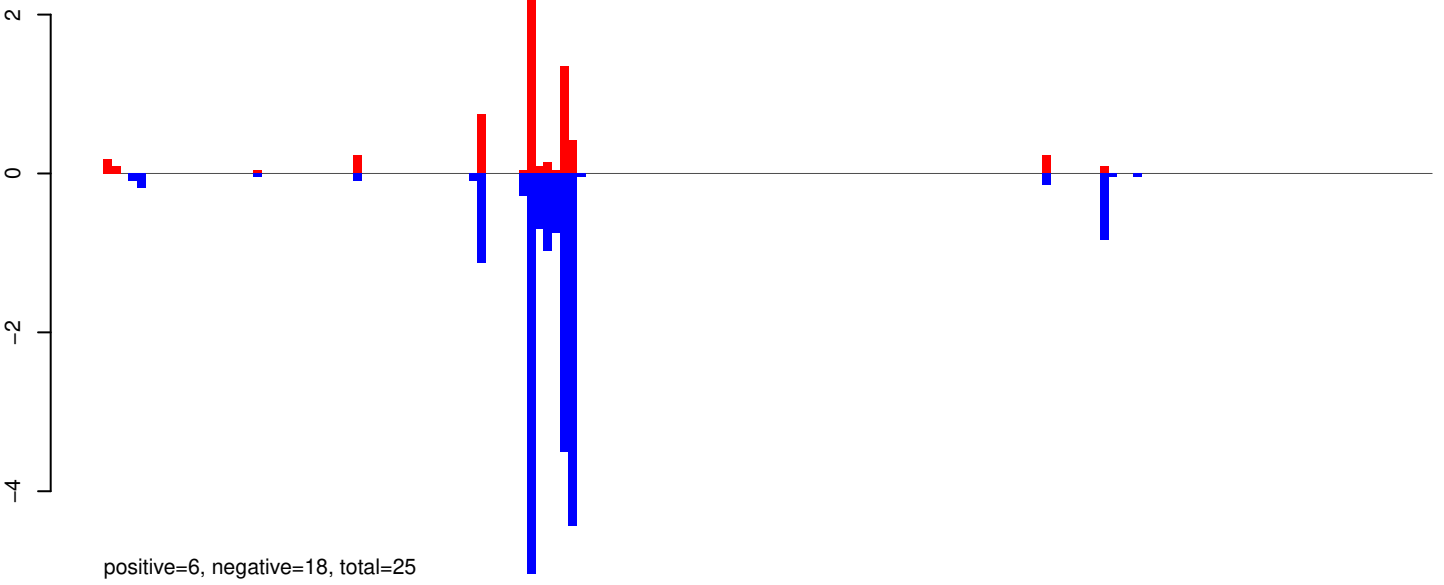
AeAeg_Aag2SINV_dsLuc.rep



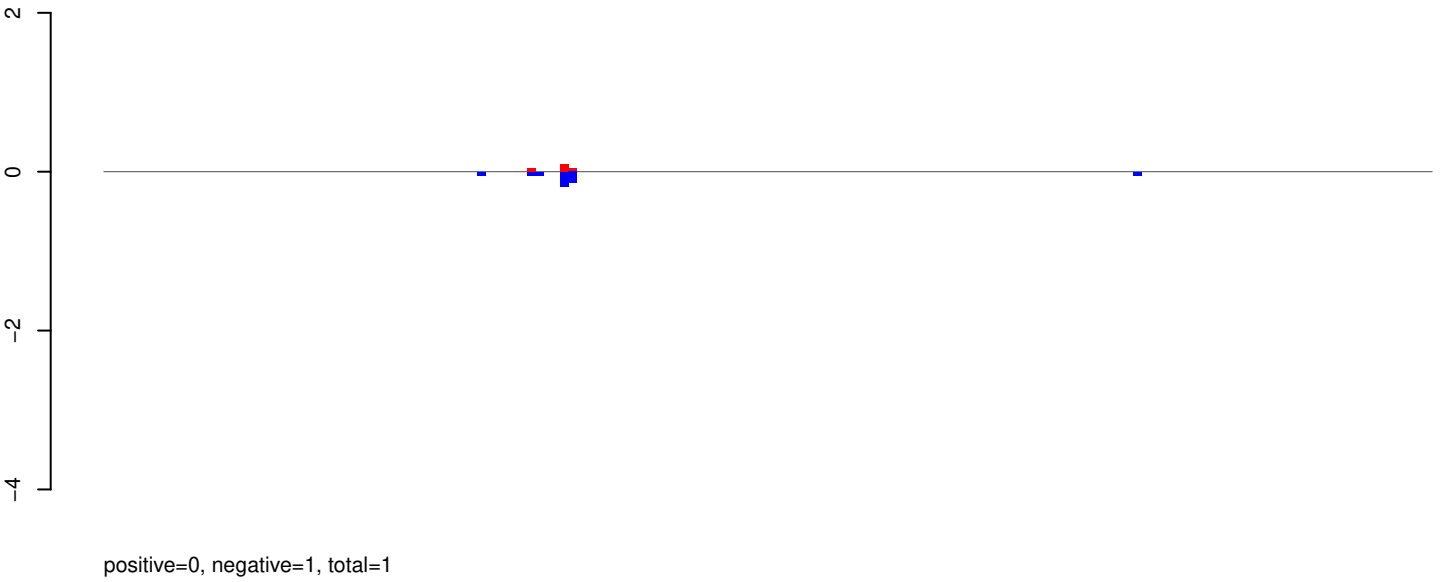
positive=5, negative=136, total=141, window size=25, length=6603, virus@NC_004177.1-Drosophila_x_virus_segment_A_and_segment_B:1-6603

0 1000 2000 3000 4000 5000 6000

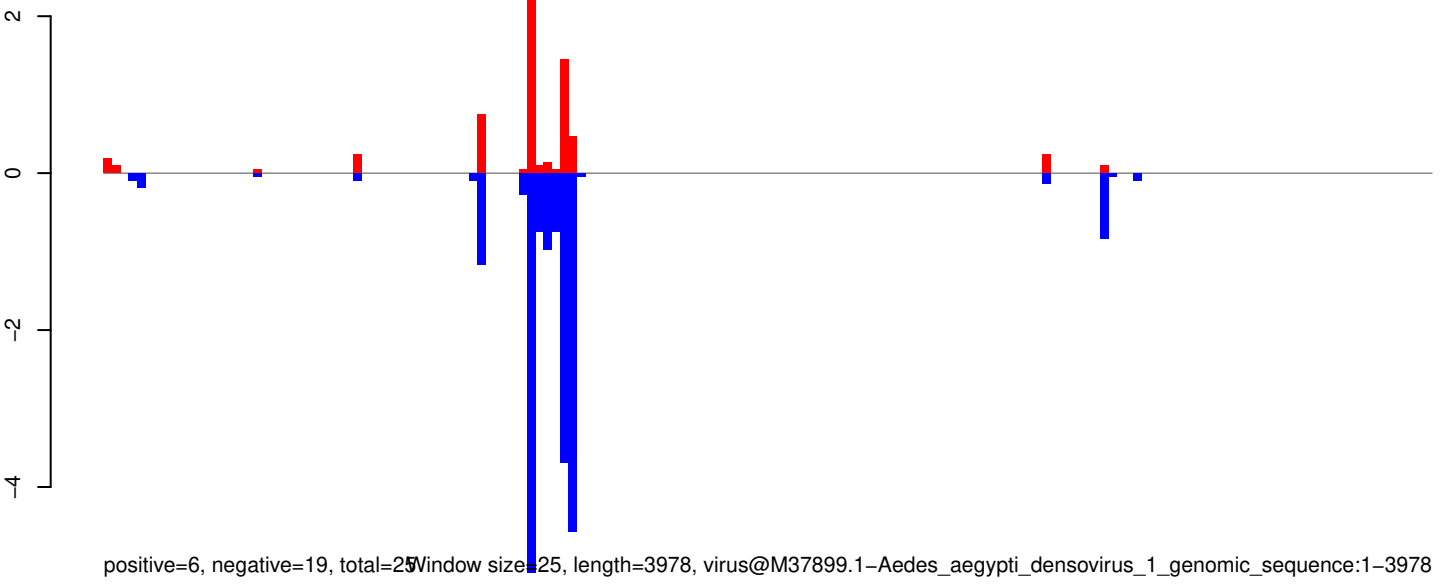
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

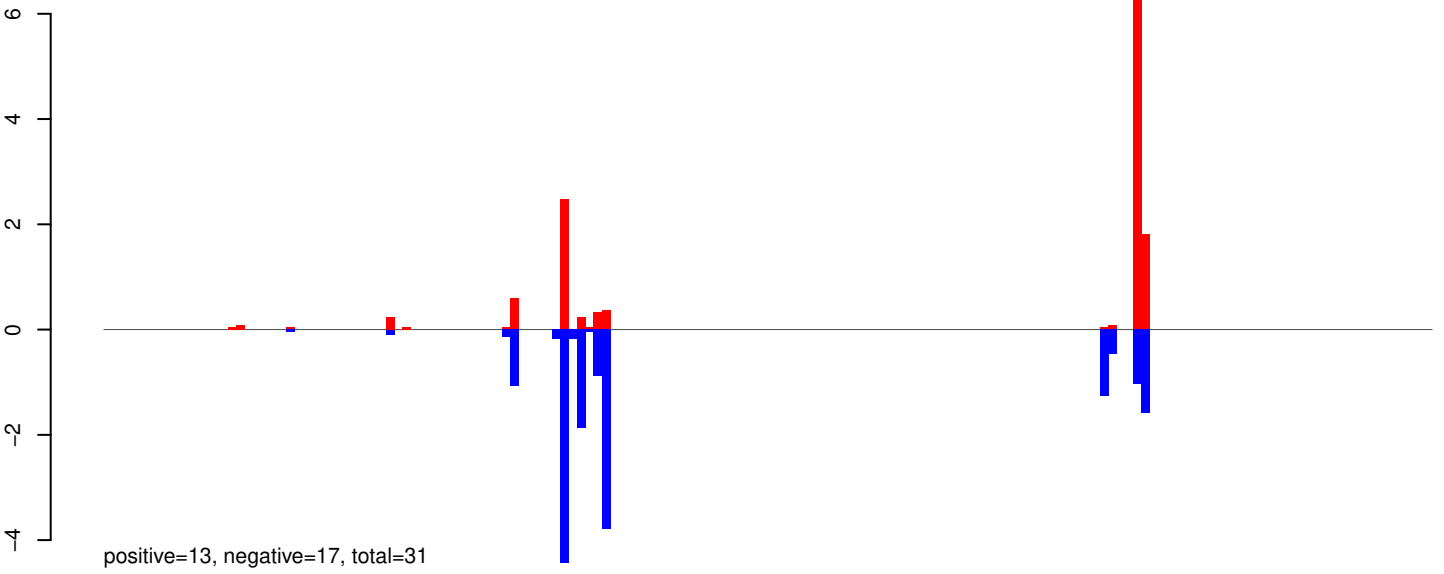


AeAeg_Aag2SINV_dsLuc.rep

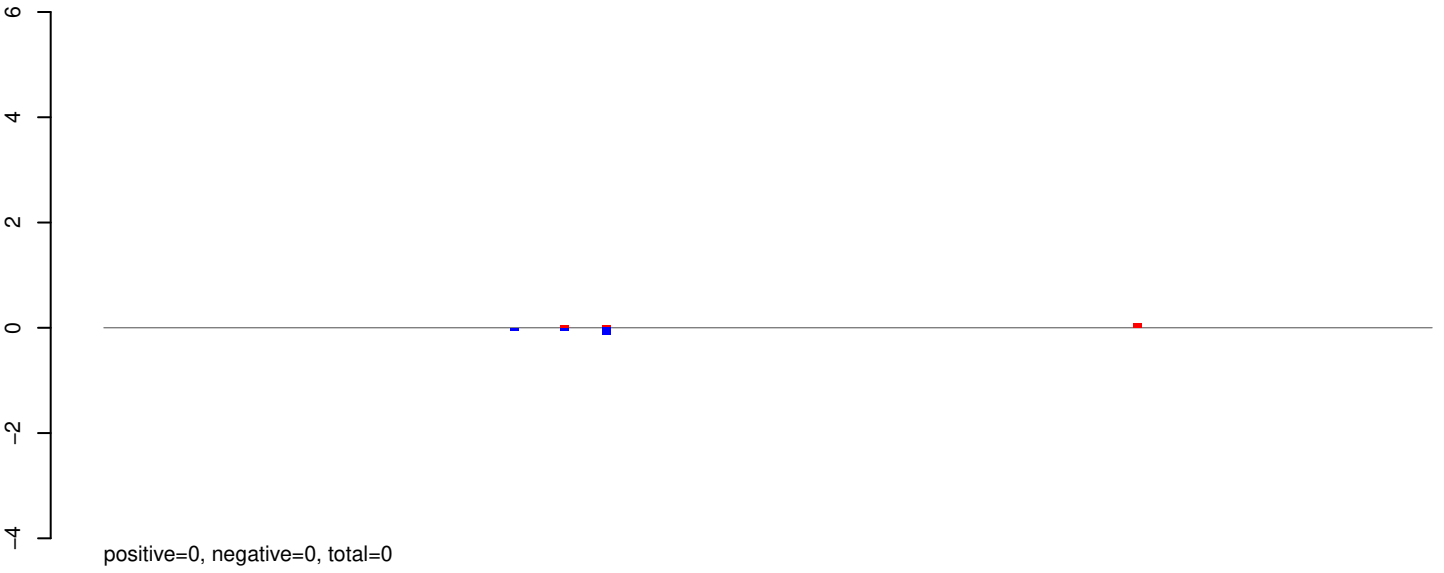


Window size=25, length=3978, virus@M37899.1-Aedes_aegypti_densovirus_1_genomic_sequence:1-3978

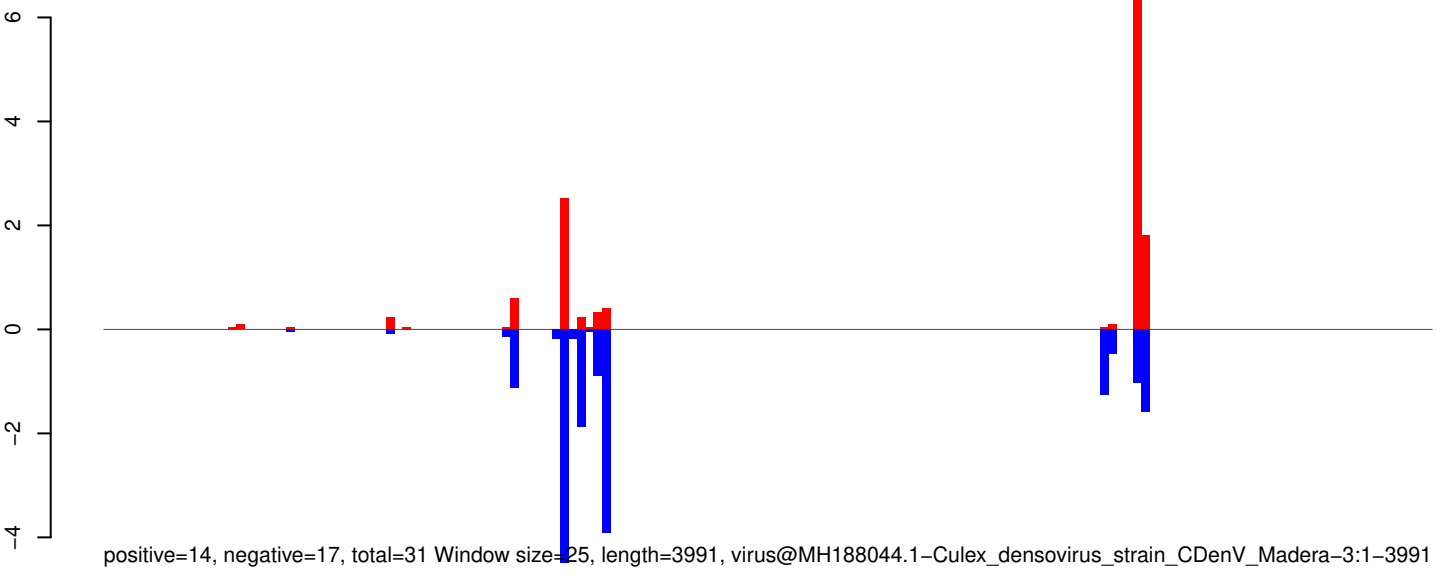
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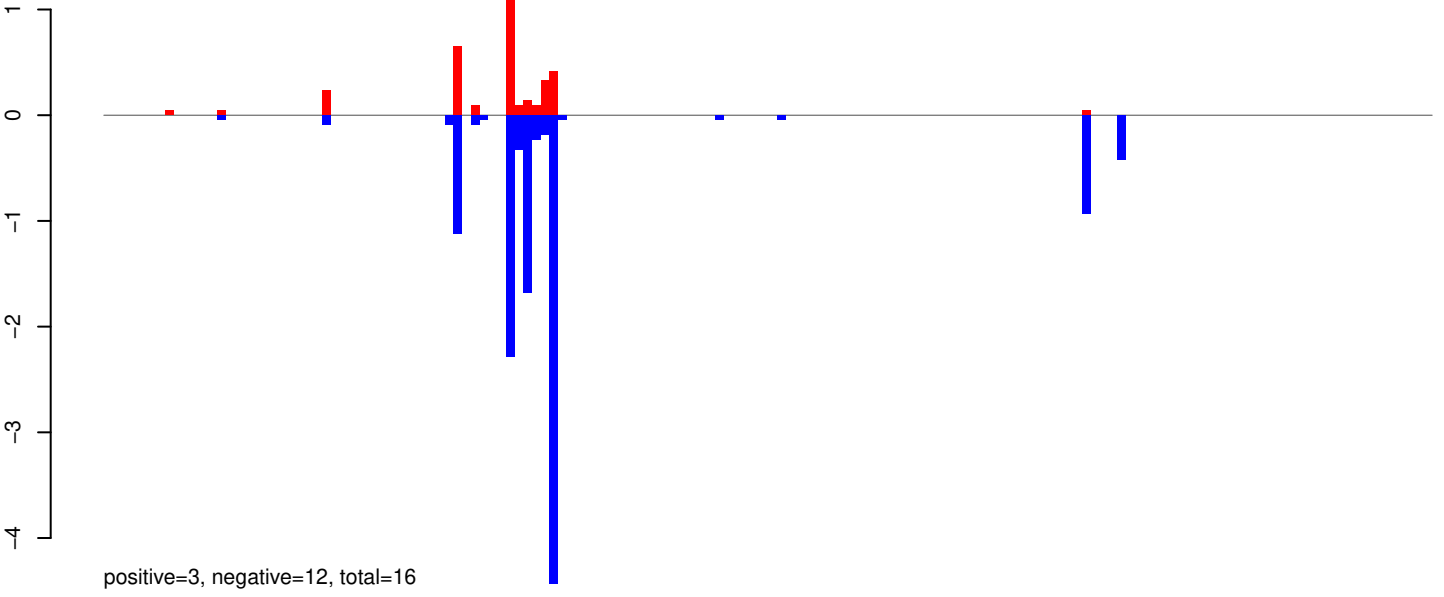
AeAeg_Aag2SINV_dsLuc.24_35.rep



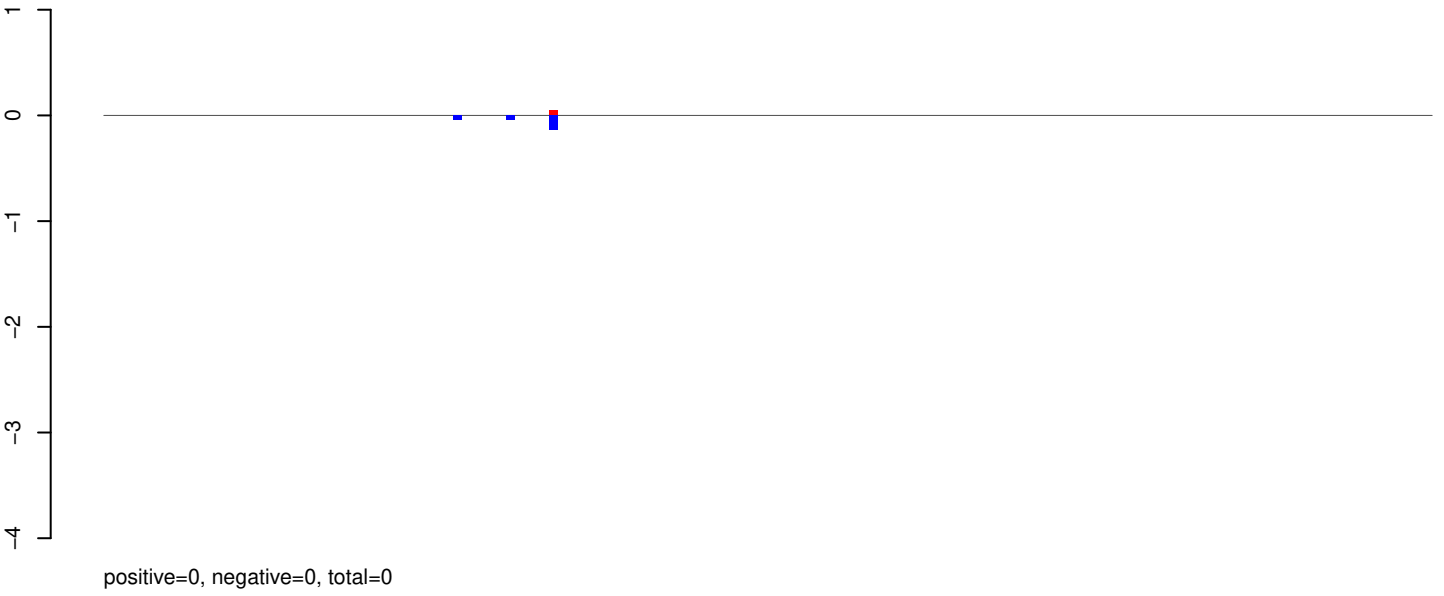
AeAeg_Aag2SINV_dsLuc.rep



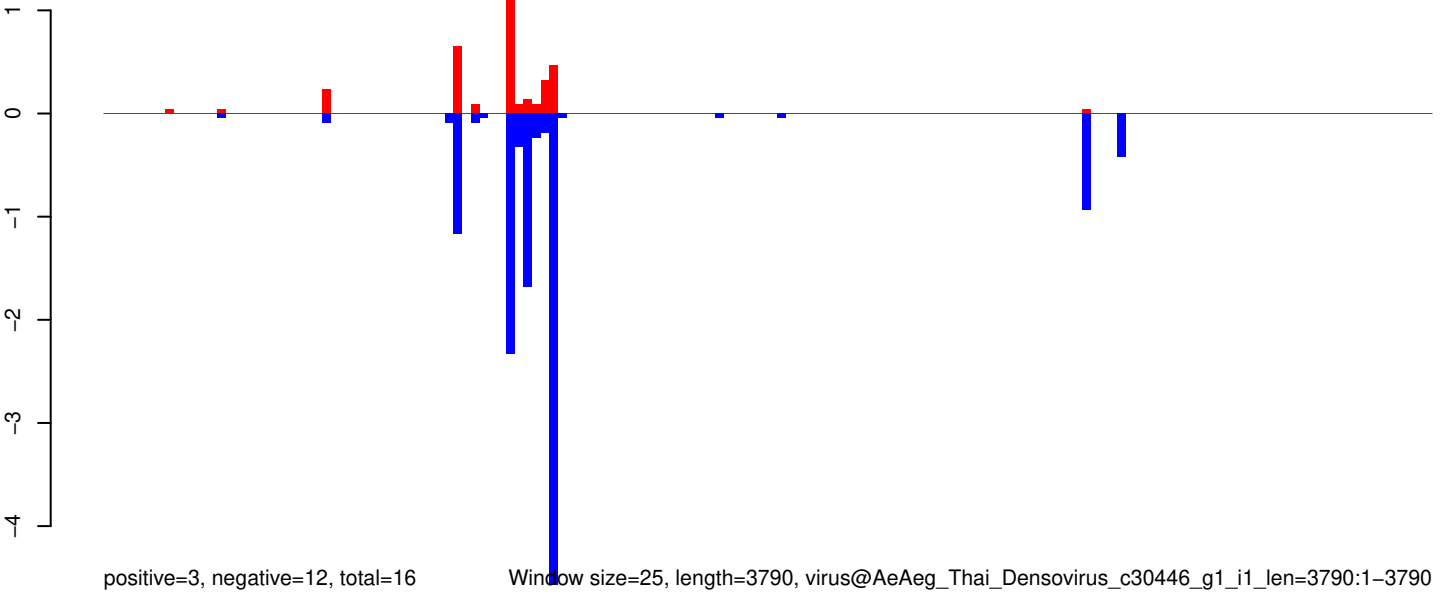
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



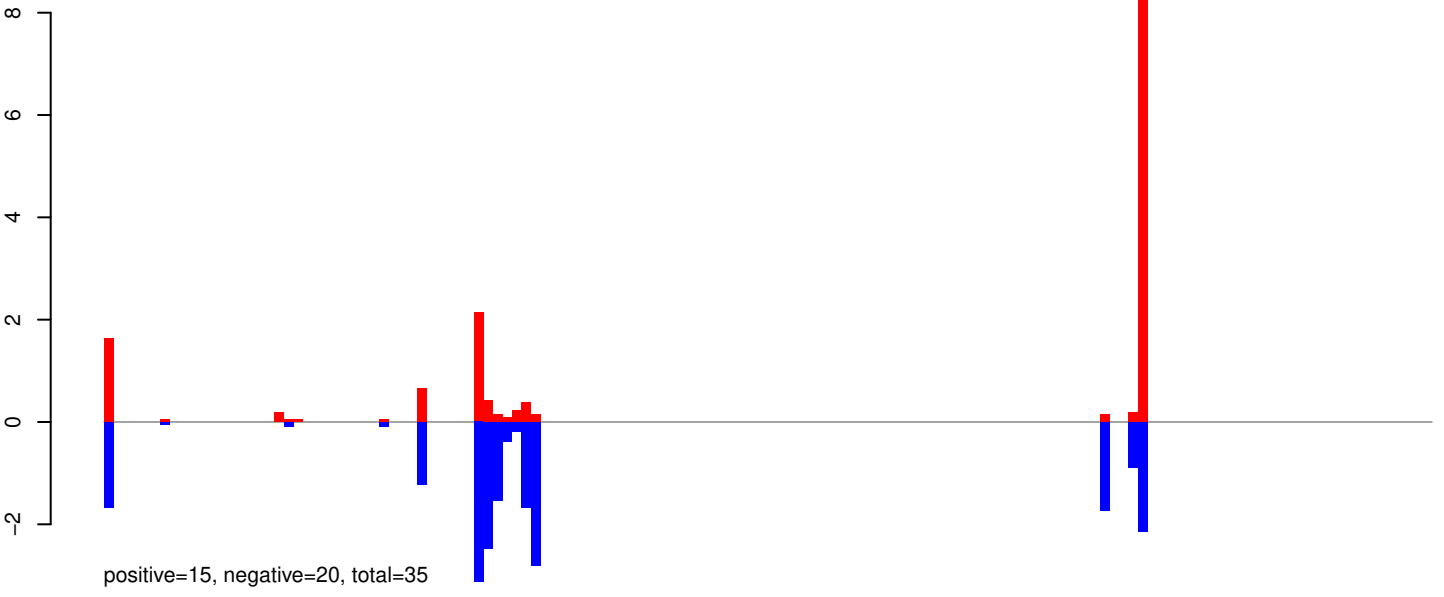
AeAeg_Aag2SINV_dsLuc.rep



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

0 1000 2000 3000

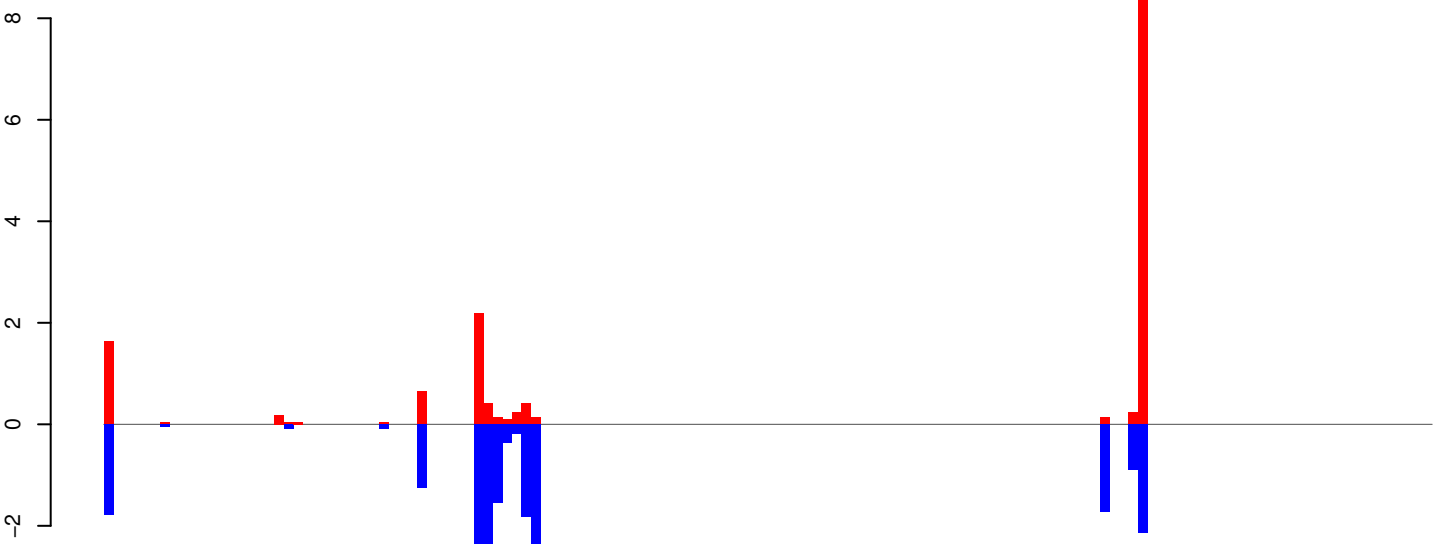
AeAeg_Aag2SINV_dsLuc.18_23.rep



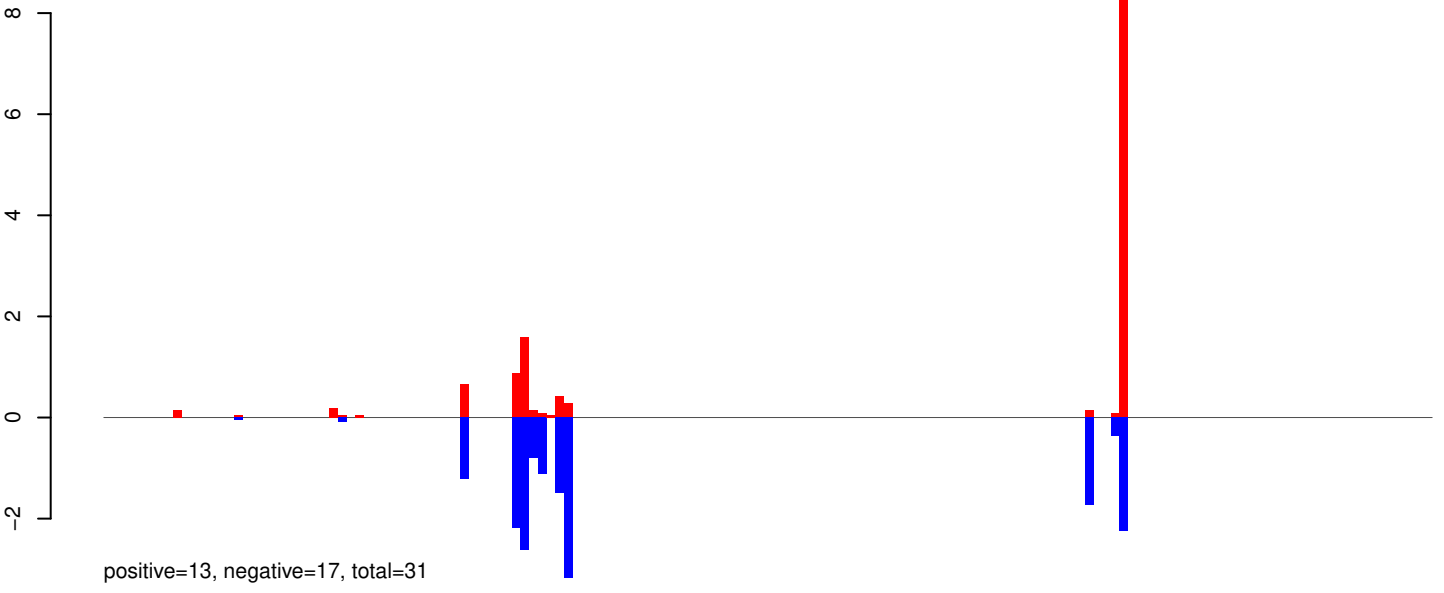
AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



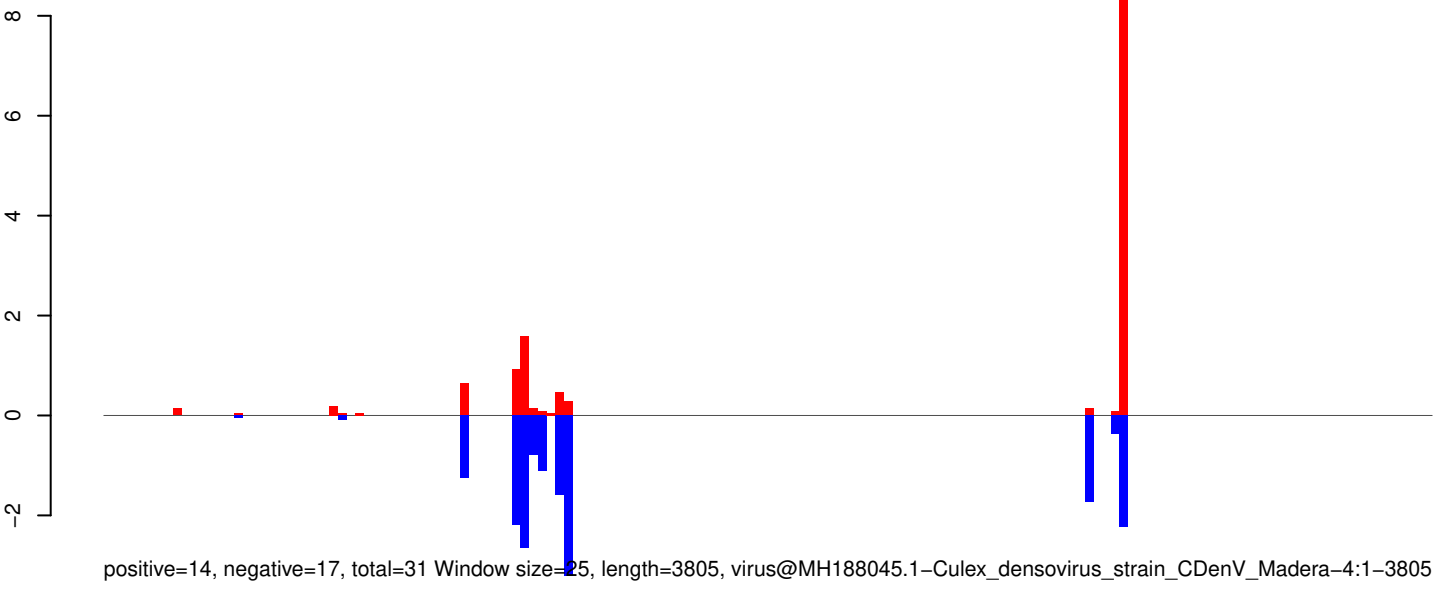
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep

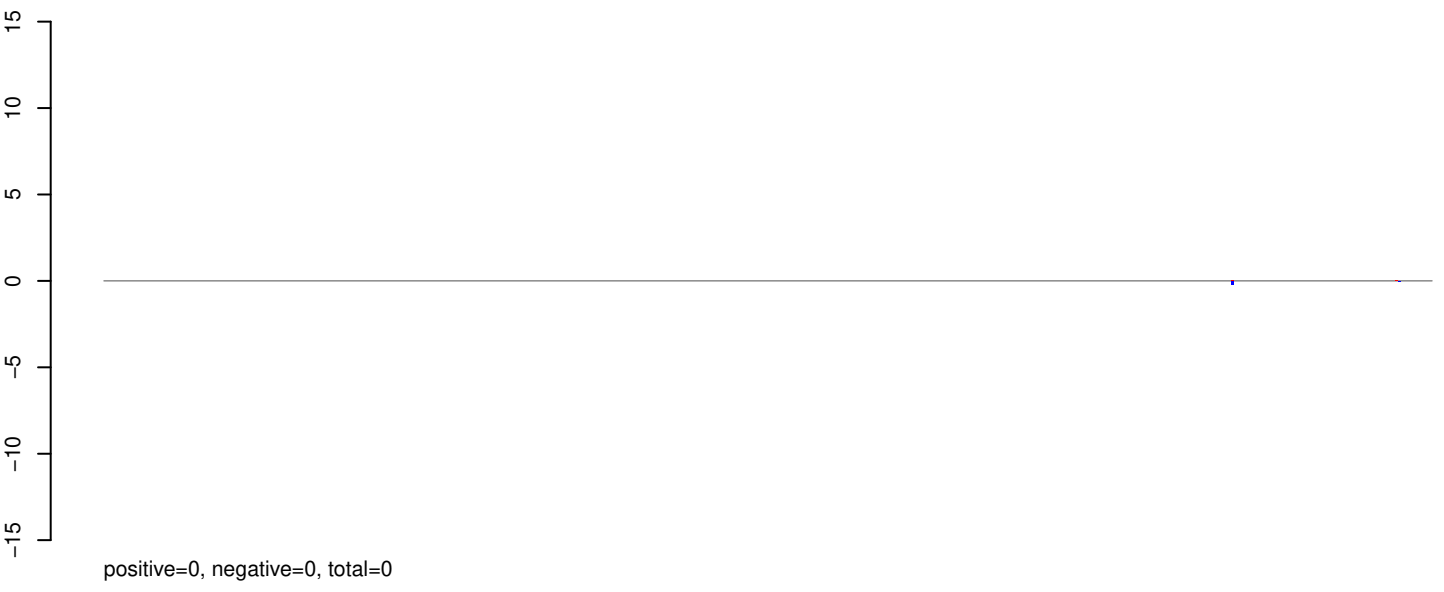


0 1000 2000 3000 4000

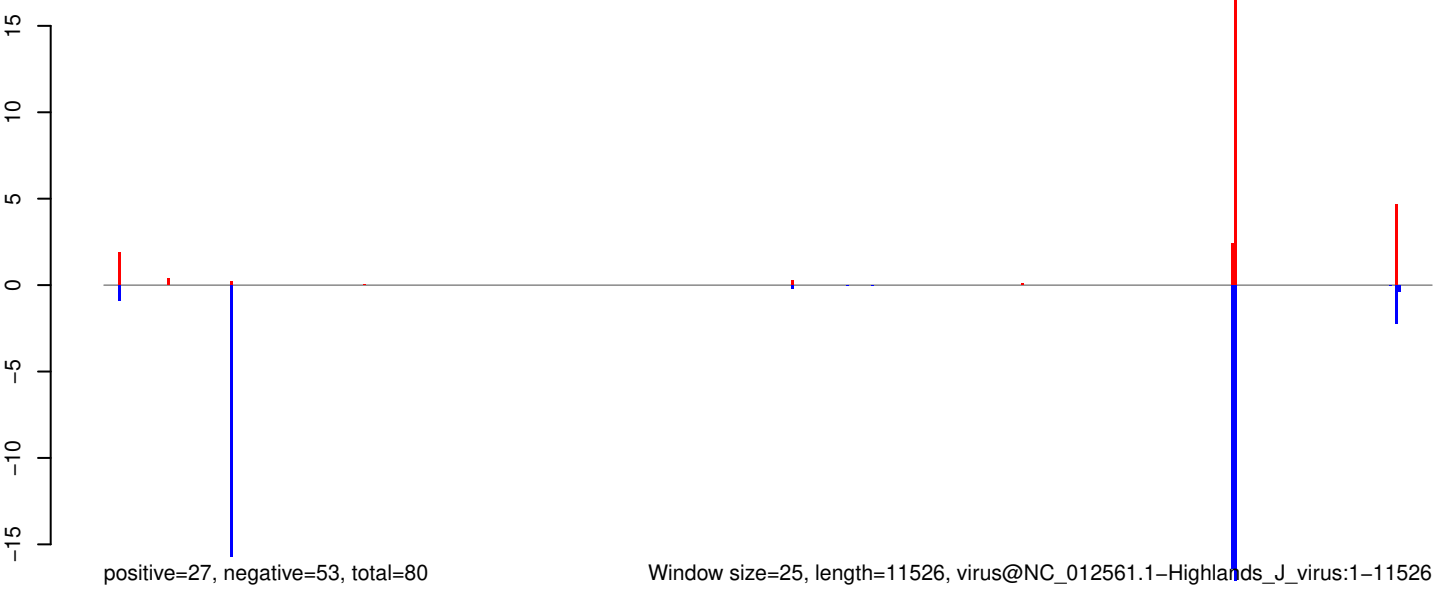
AeAeg_Aag2SINV_dsLuc.18_23.rep



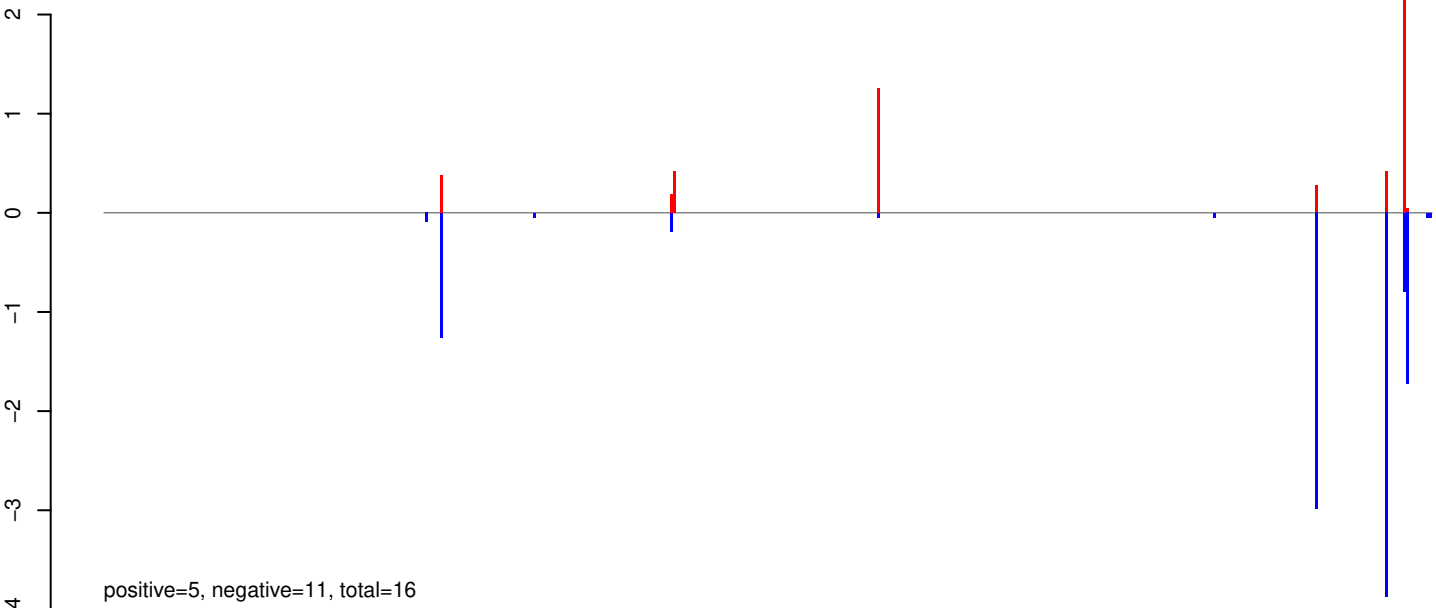
AeAeg_Aag2SINV_dsLuc.24_35.rep



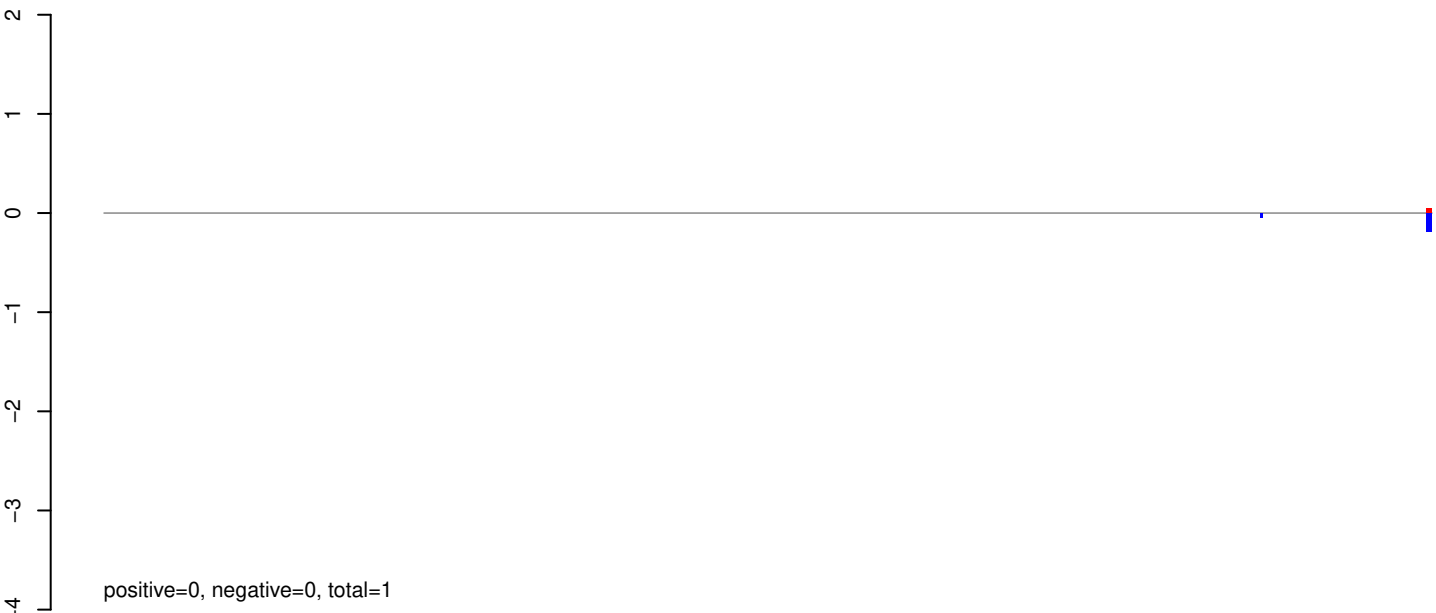
AeAeg_Aag2SINV_dsLuc.rep



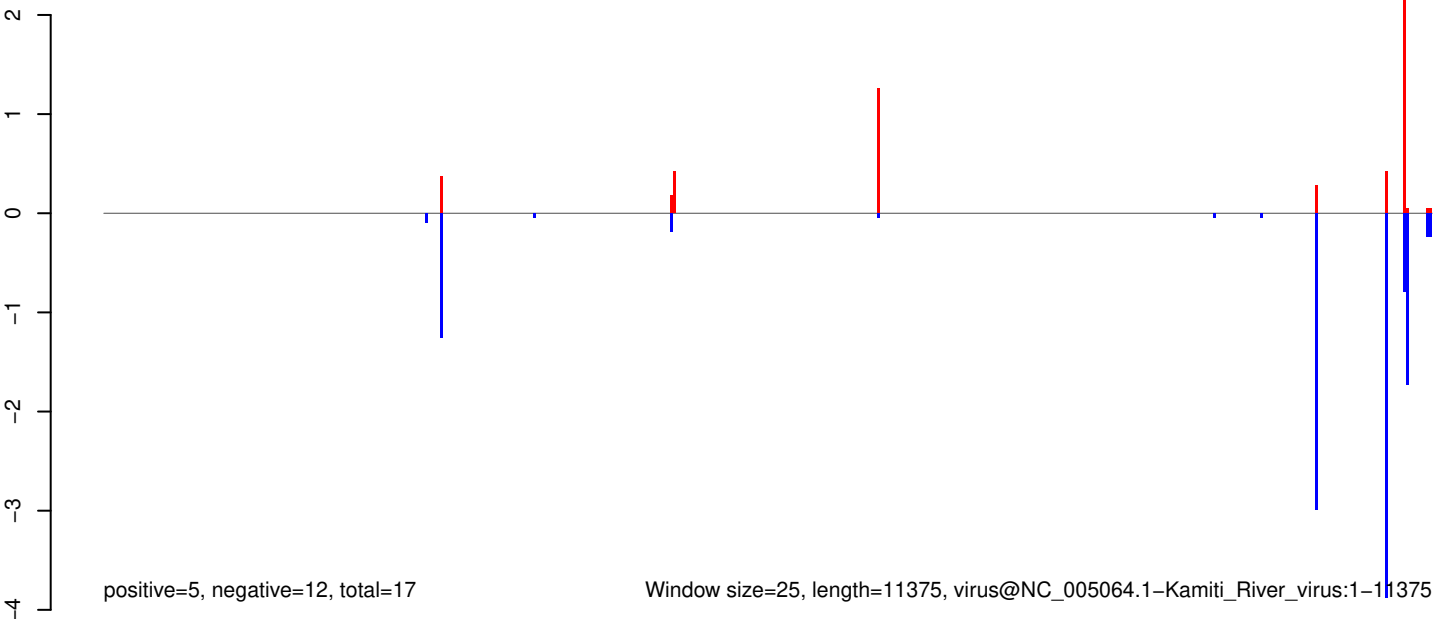
AeAeg_Aag2SINV_dsLuc.18_23.rep



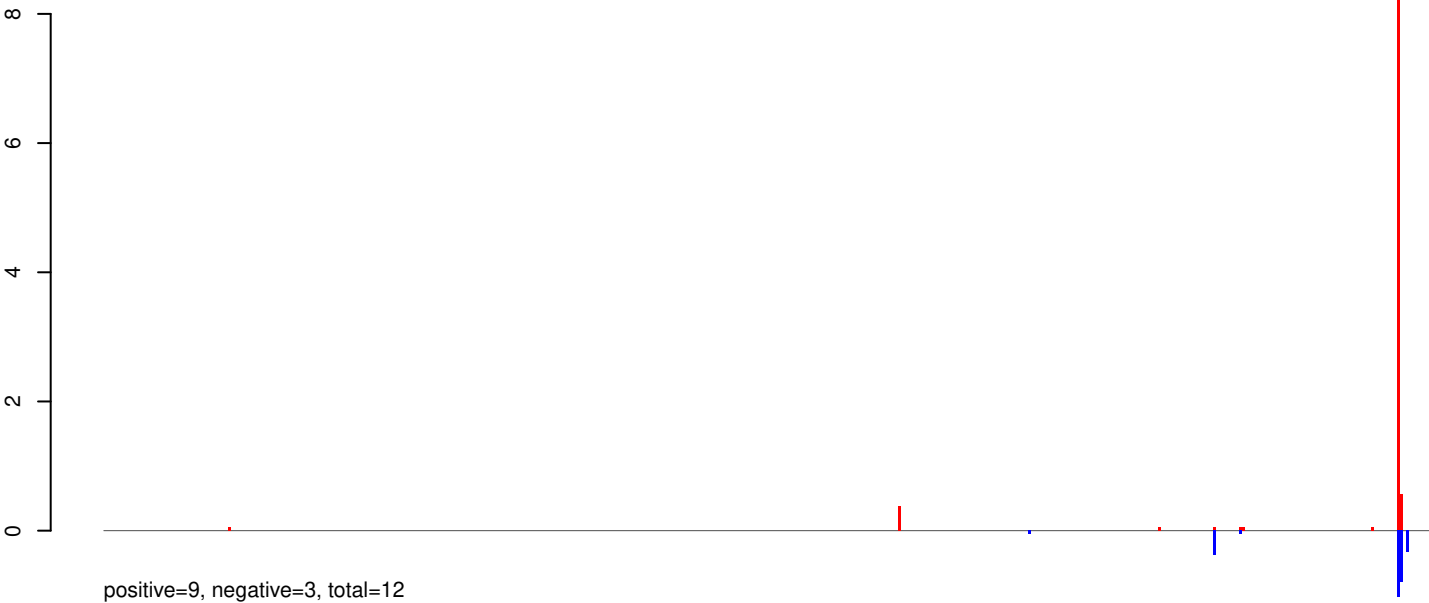
AeAeg_Aag2SINV_dsLuc.24_35.rep



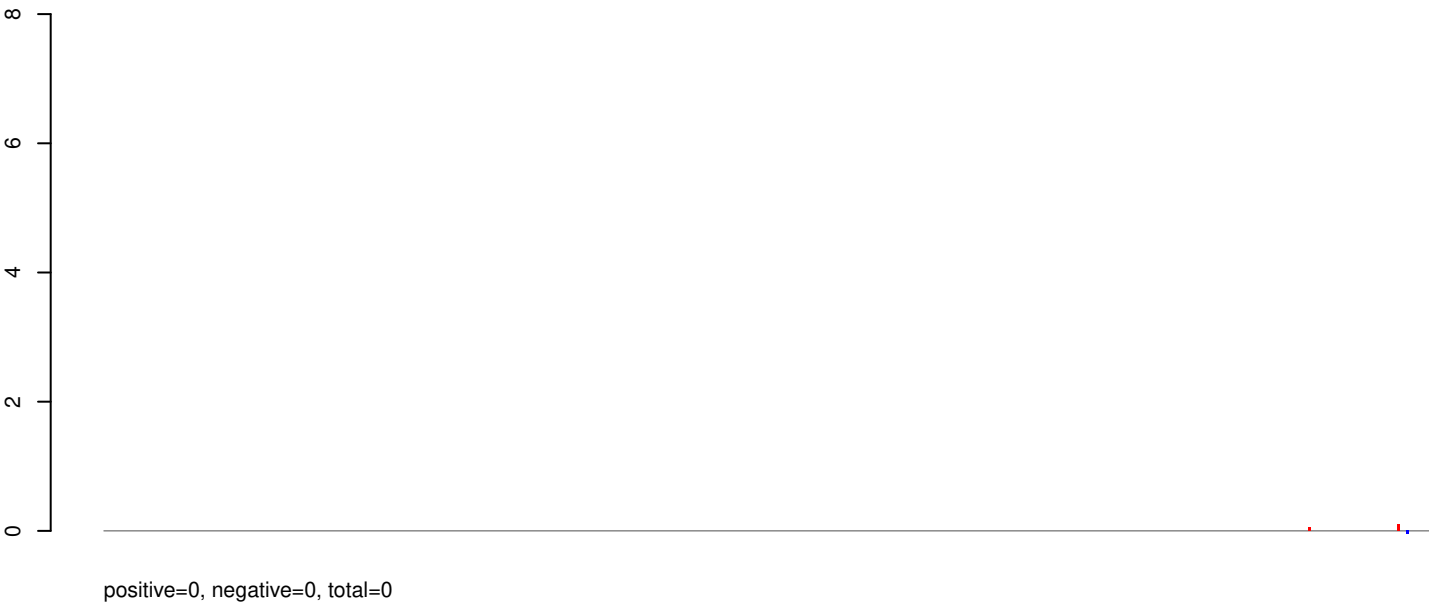
AeAeg_Aag2SINV_dsLuc.rep



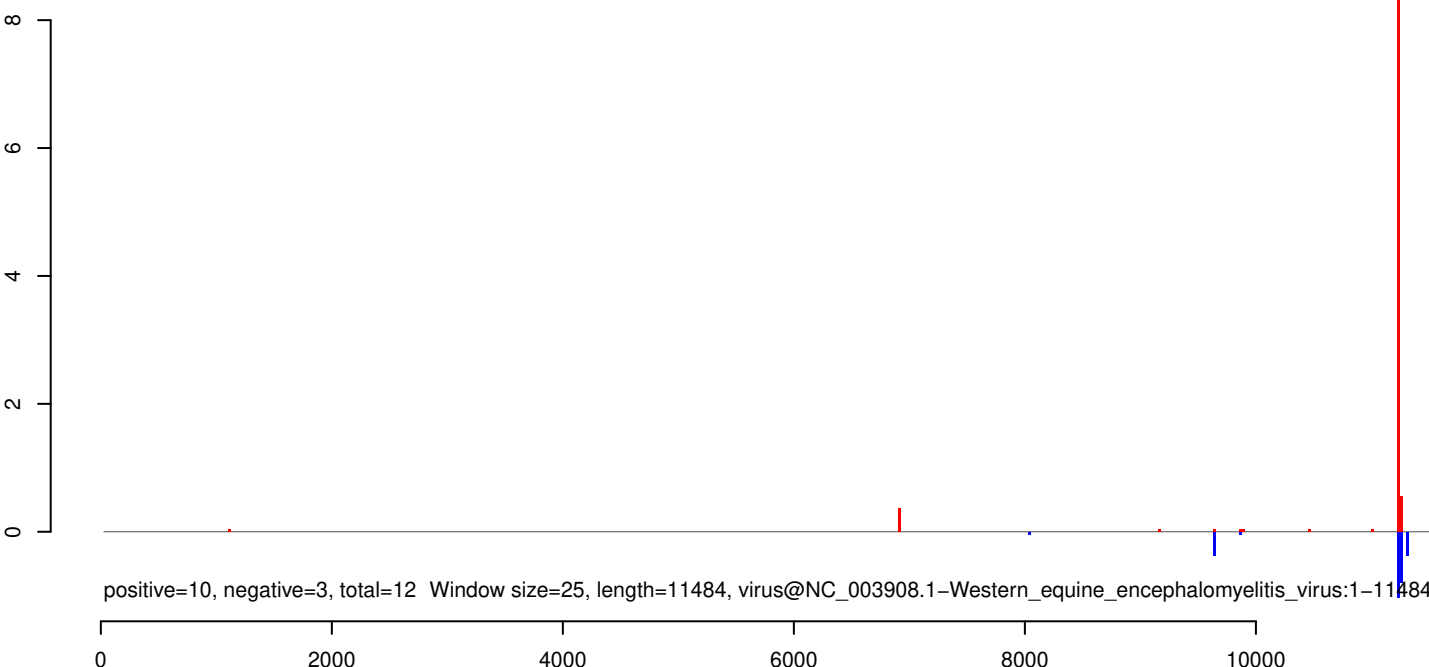
AeAeg_Aag2SINV_dsLuc.18_23.rep



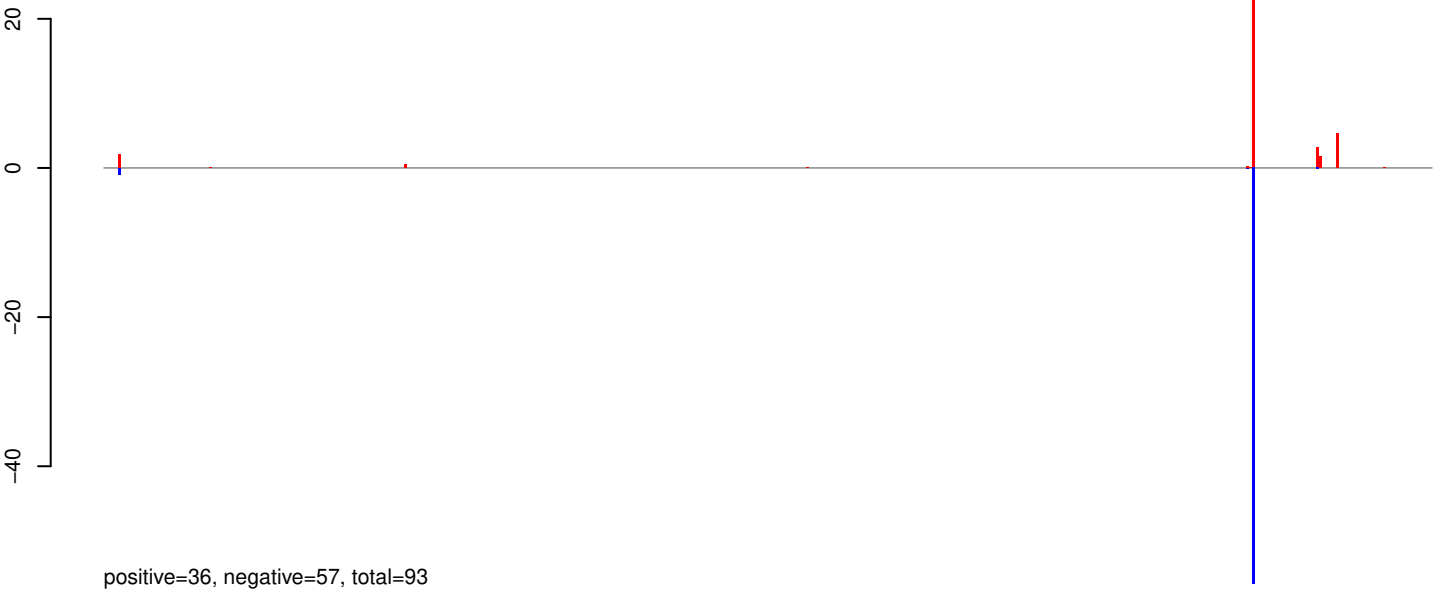
AeAeg_Aag2SINV_dsLuc.24_35.rep



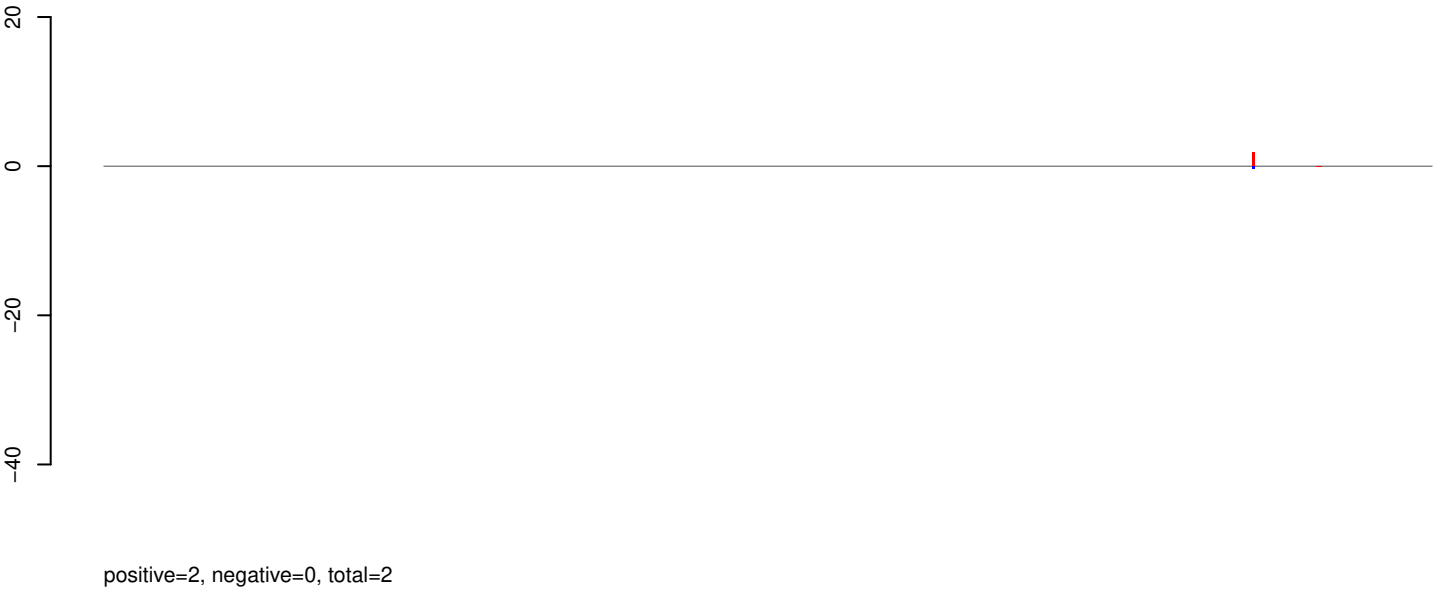
AeAeg_Aag2SINV_dsLuc.rep



AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

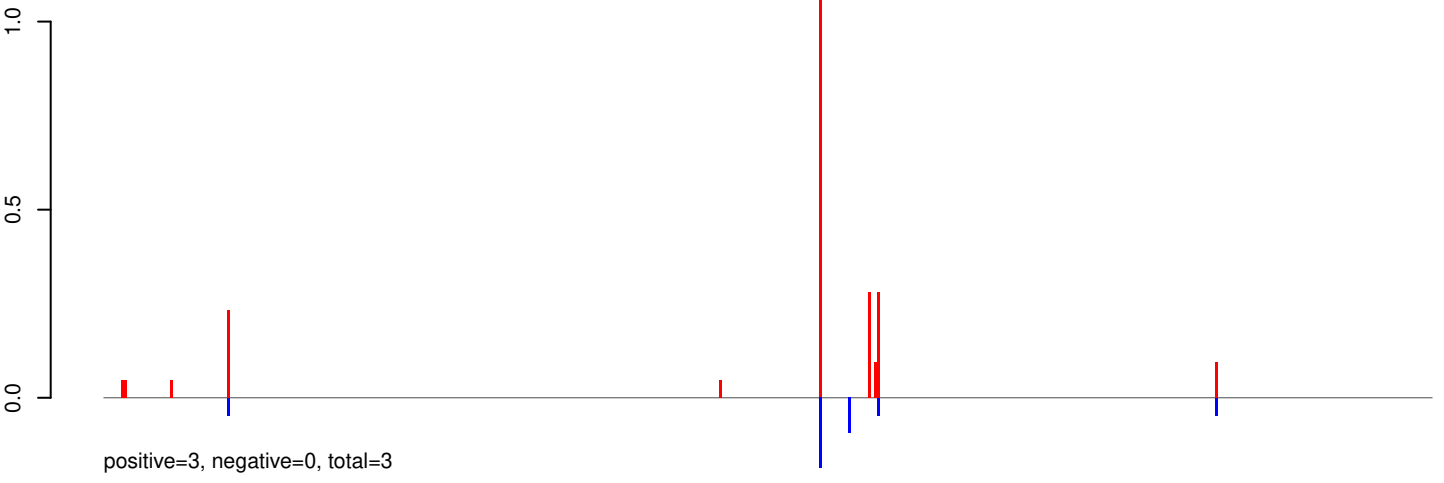


AeAeg_Aag2SINV_dsLuc.rep

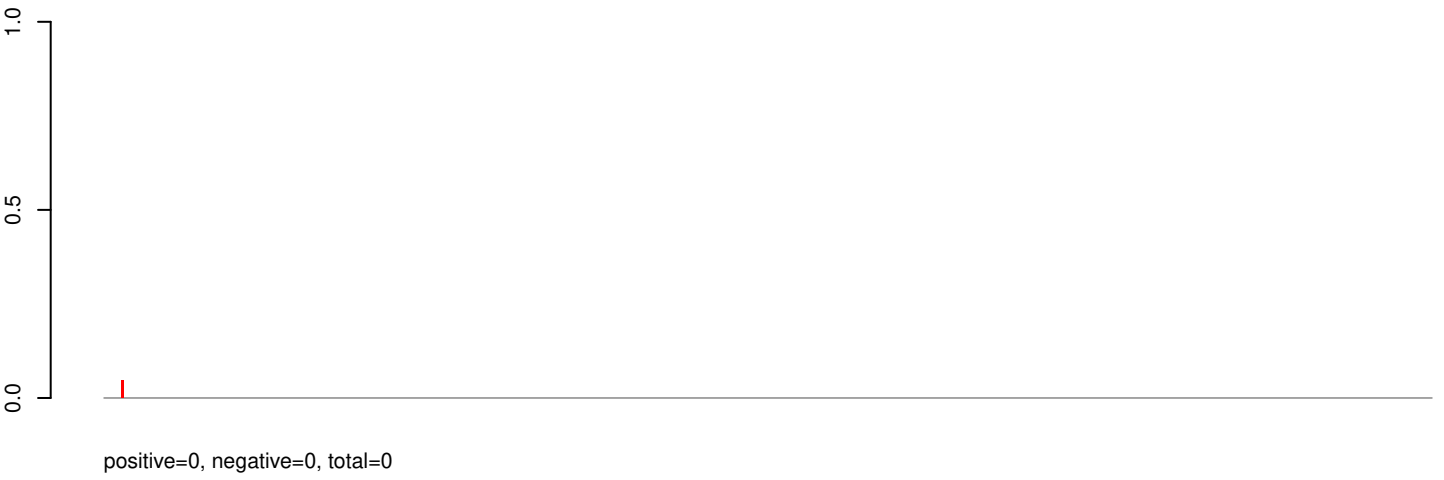


Window size=25, length=11381, virus@NC_013528.1-Fort_Morgan_virus:1-11381

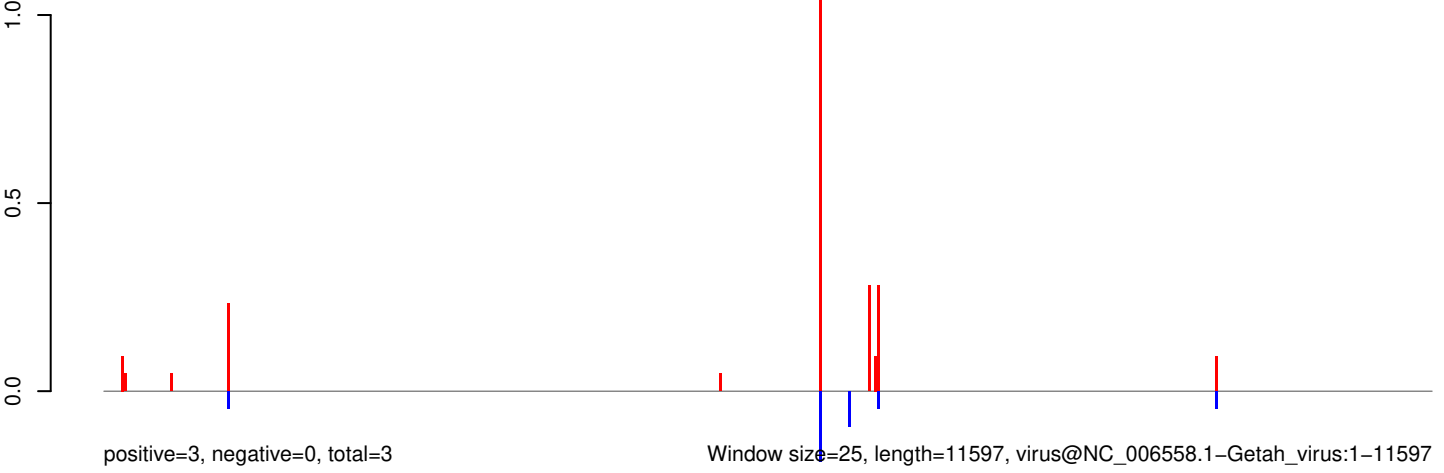
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



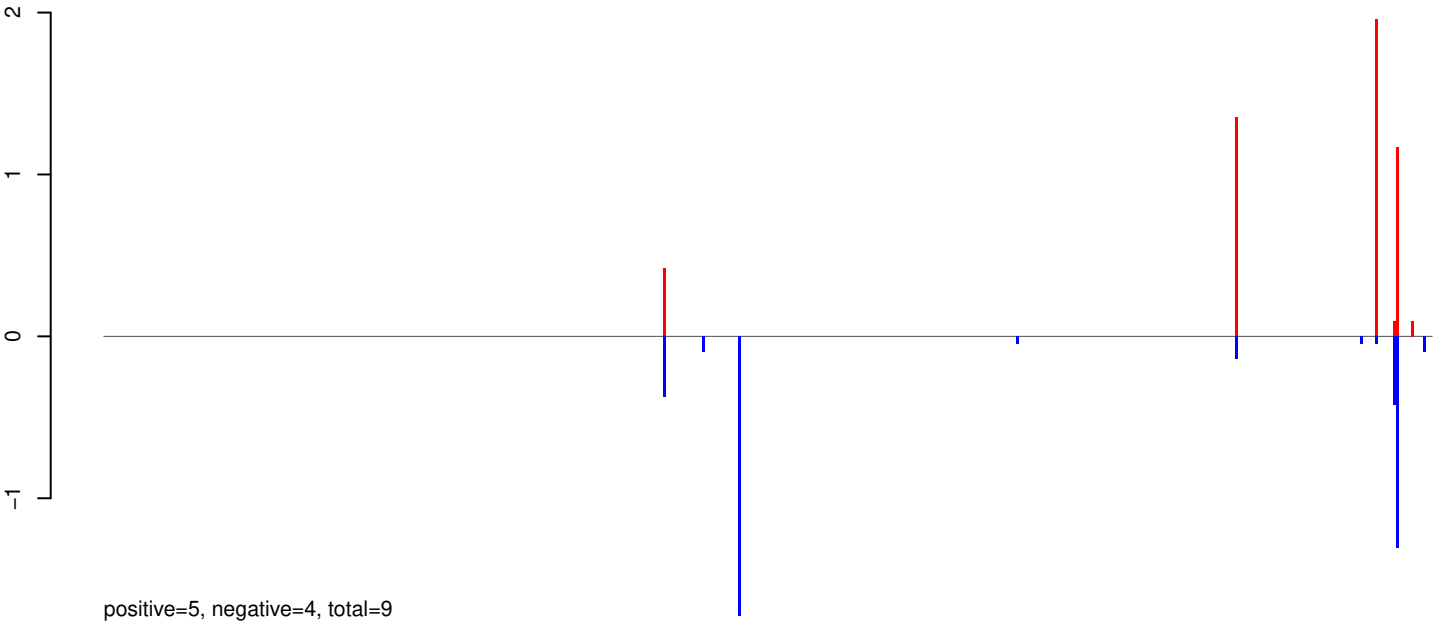
AeAeg_Aag2SINV_dsLuc.rep



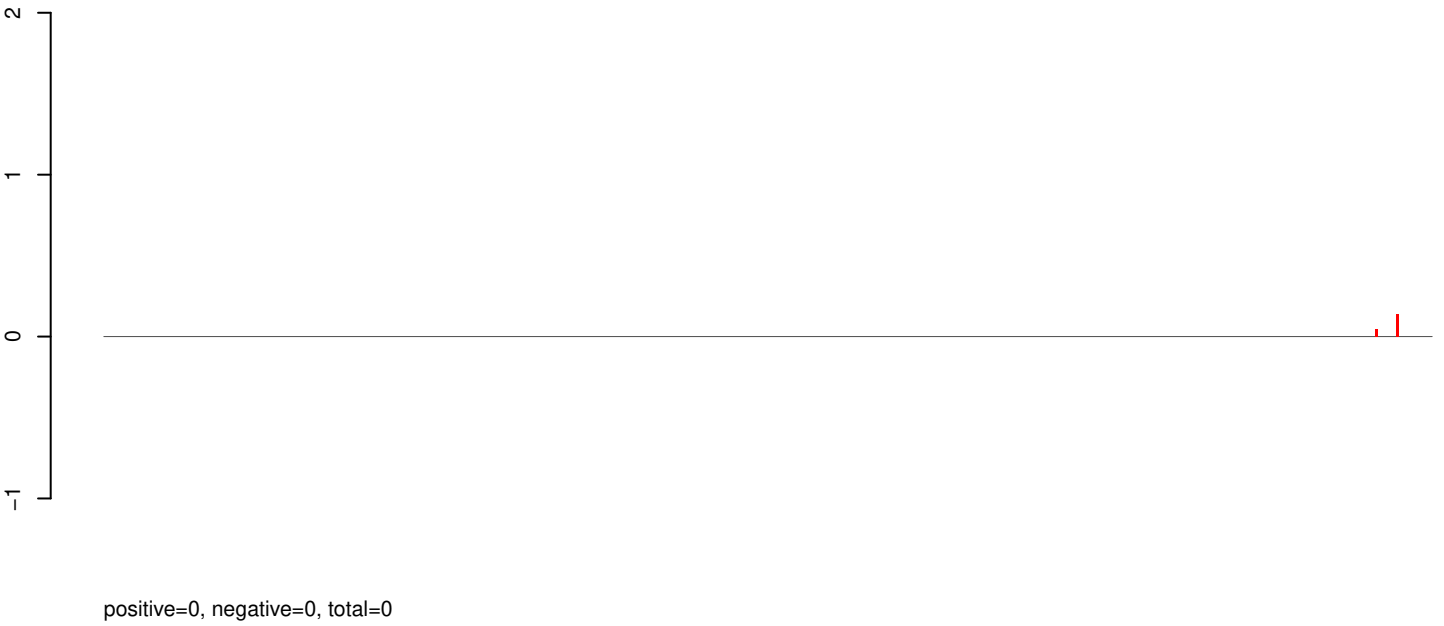
Window size=25, length=11597, virus@NC_006558.1-Getah_virus:1-11597

0 2000 4000 6000 8000 10000 12000

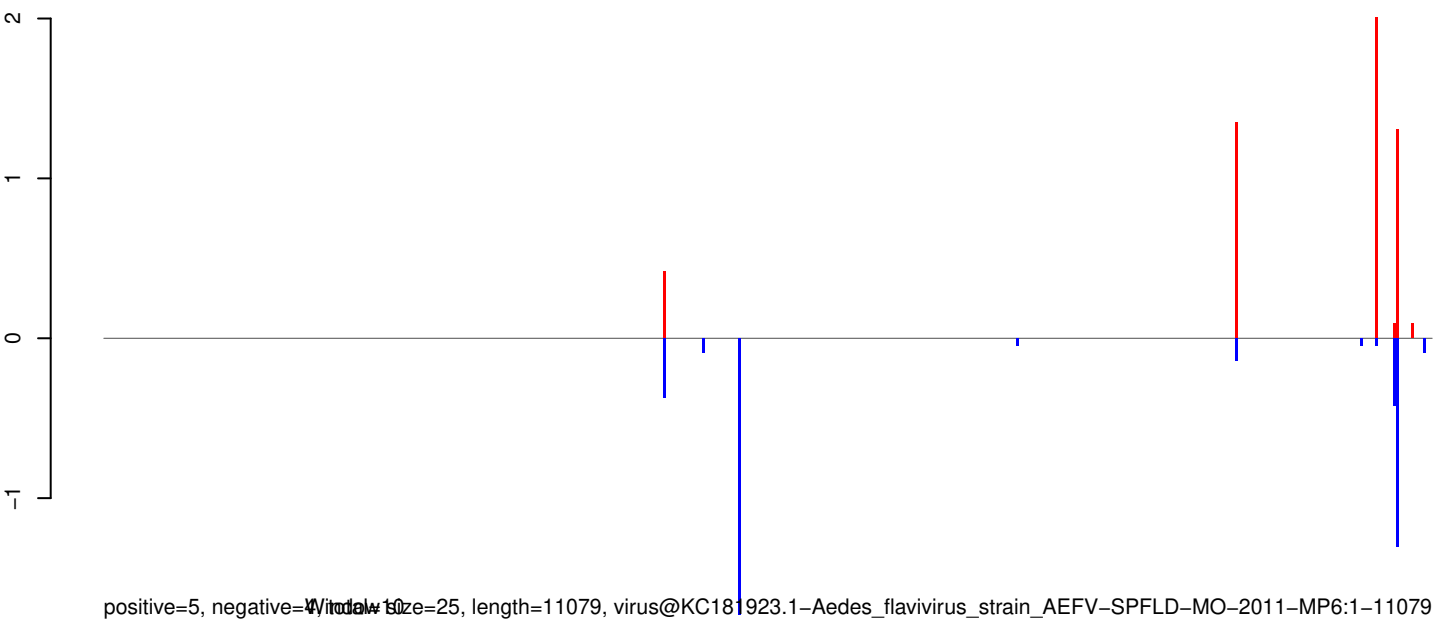
AeAeg_Aag2SINV_dsLuc.18_23.rep



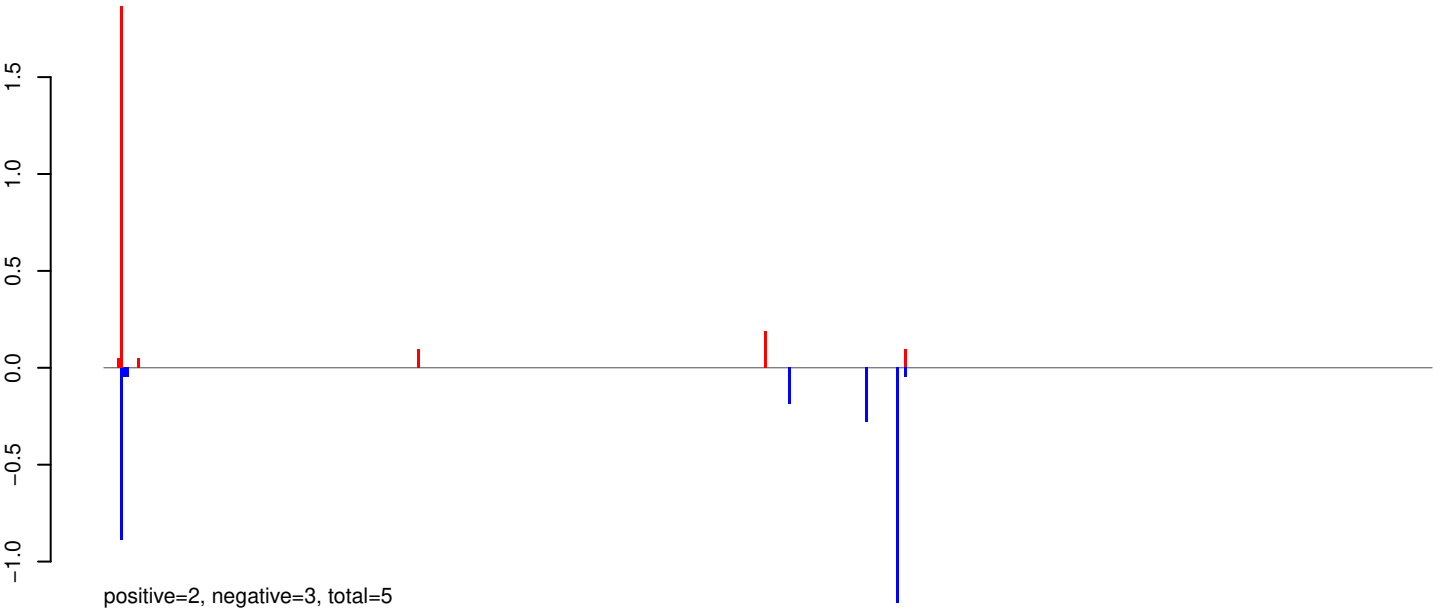
AeAeg_Aag2SINV_dsLuc.24_35.rep



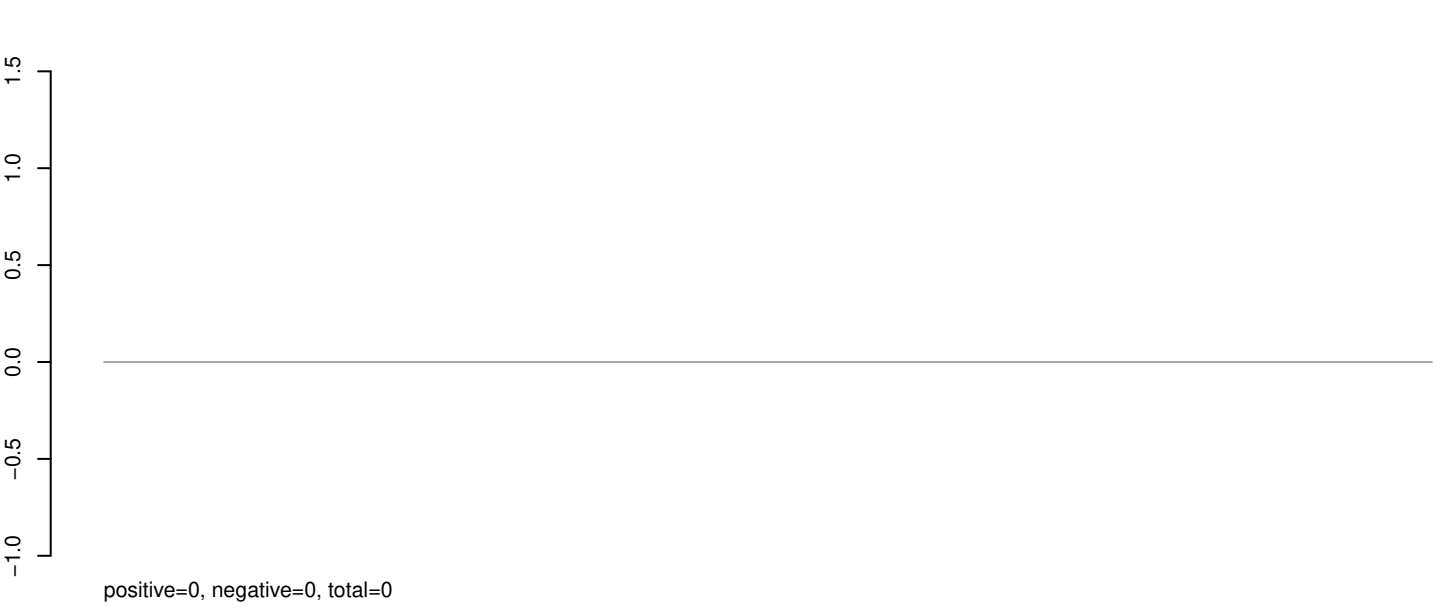
AeAeg_Aag2SINV_dsLuc.rep



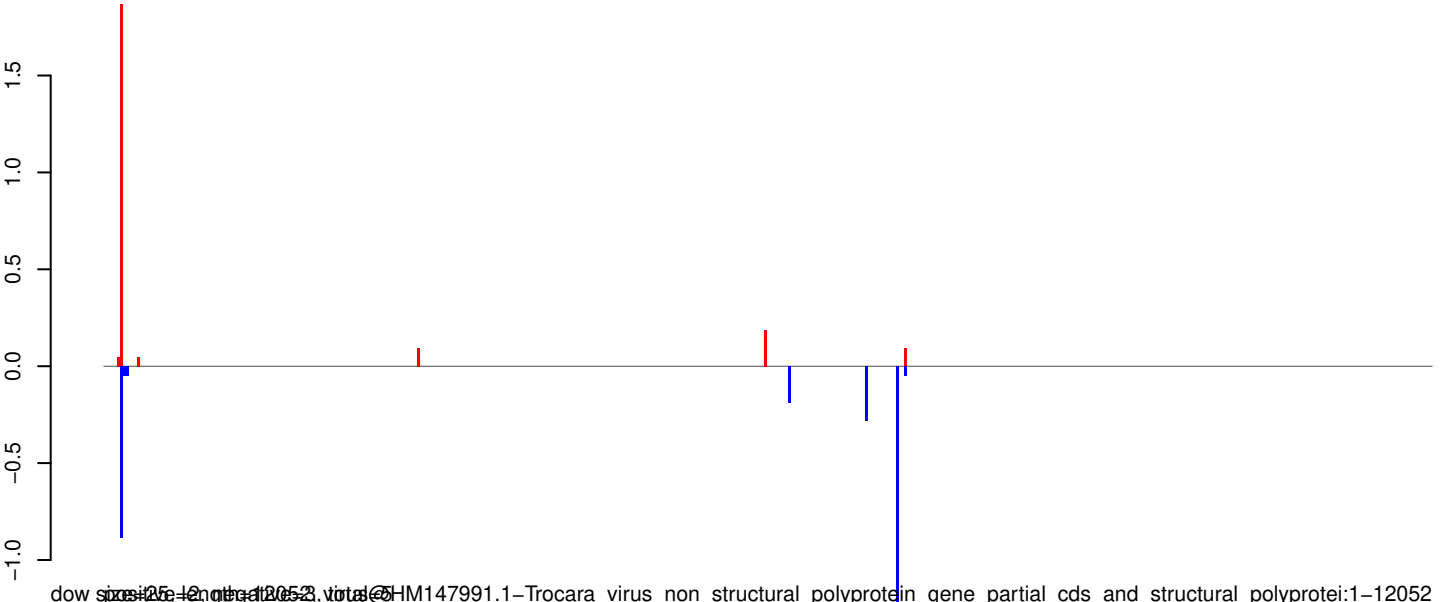
AeAeg_Aag2SINV_dsLuc.18_23.rep



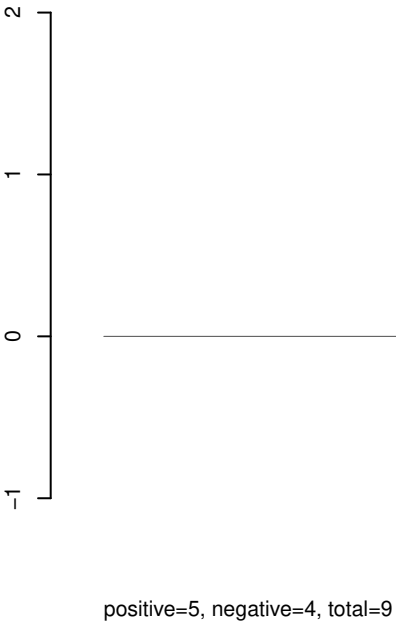
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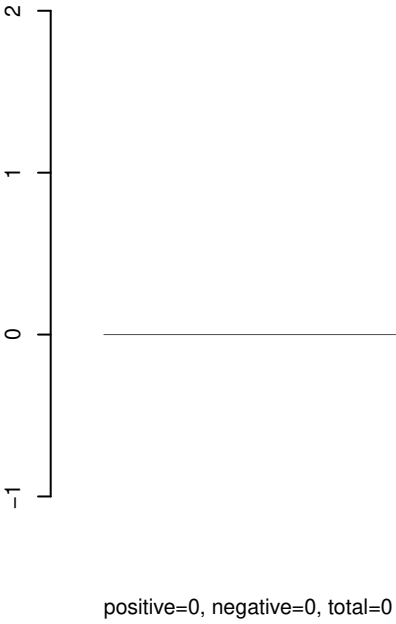
AeAeg_Aag2SINV_dsLuc.rep



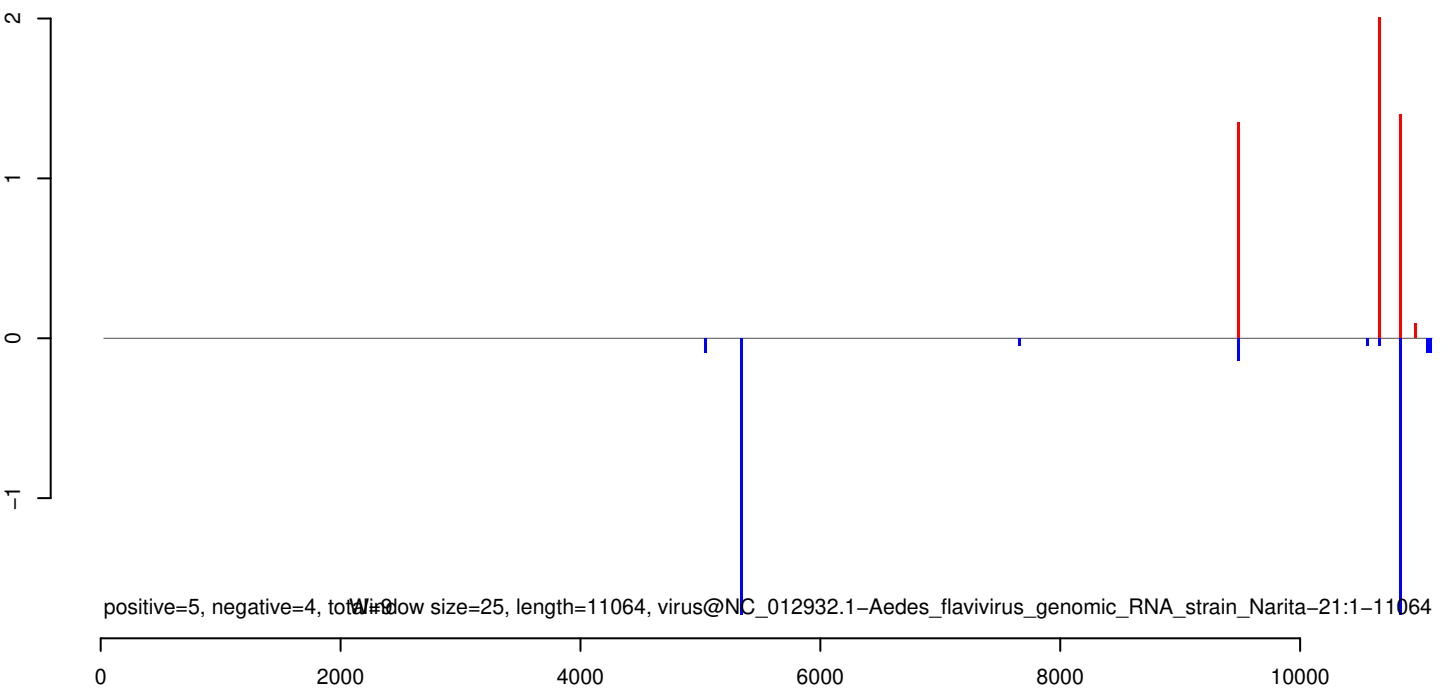
AeAeg_Aag2SINV_dsLuc.18_23.rep



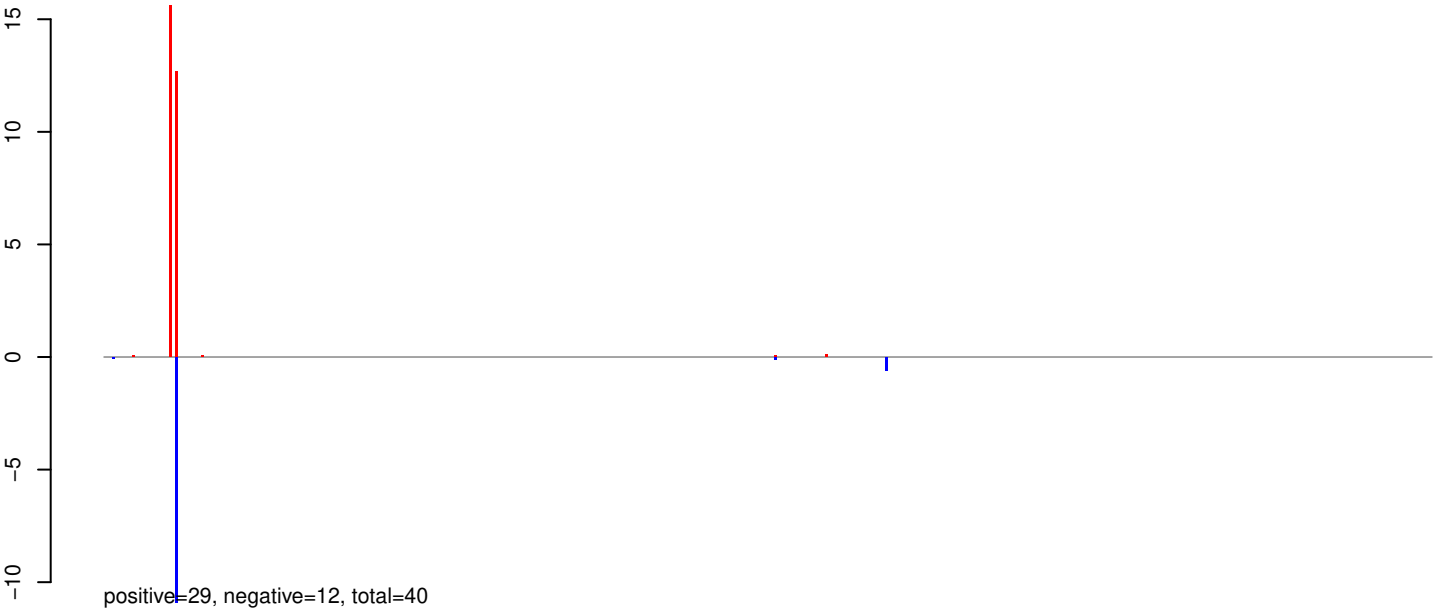
AeAeg_Aag2SINV_dsLuc.24_35.rep



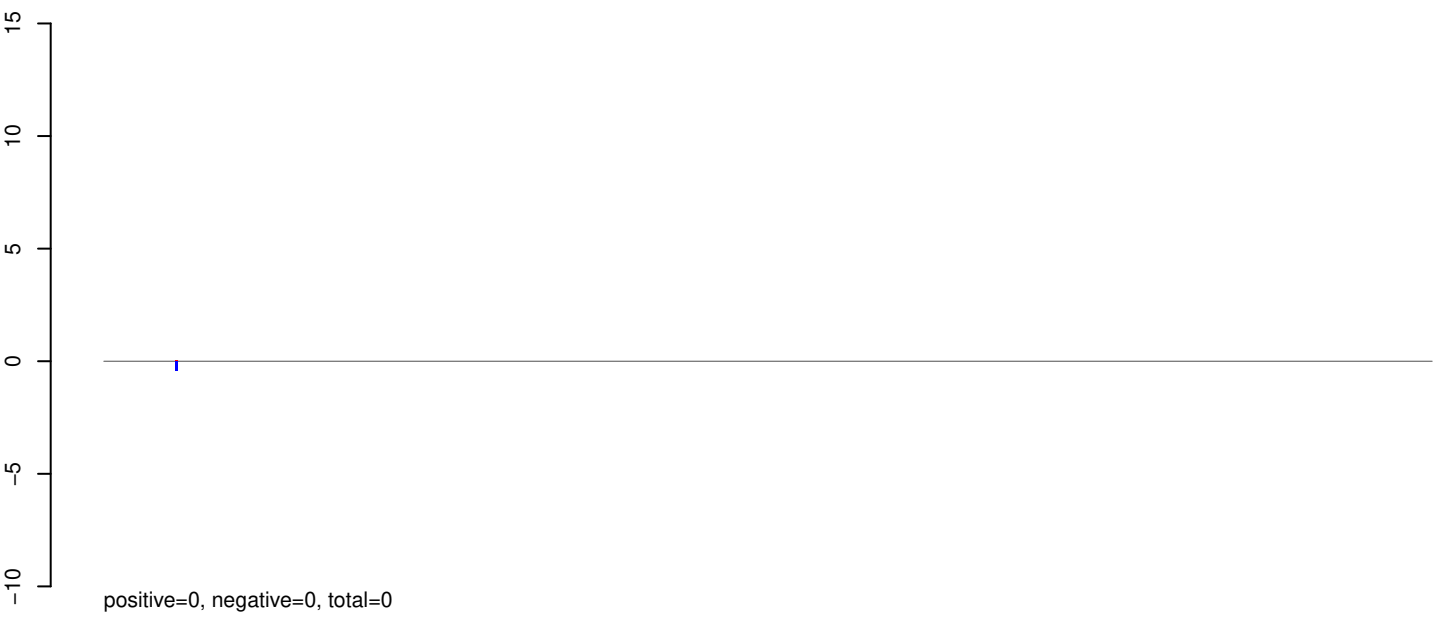
AeAeg_Aag2SINV_dsLuc.rep



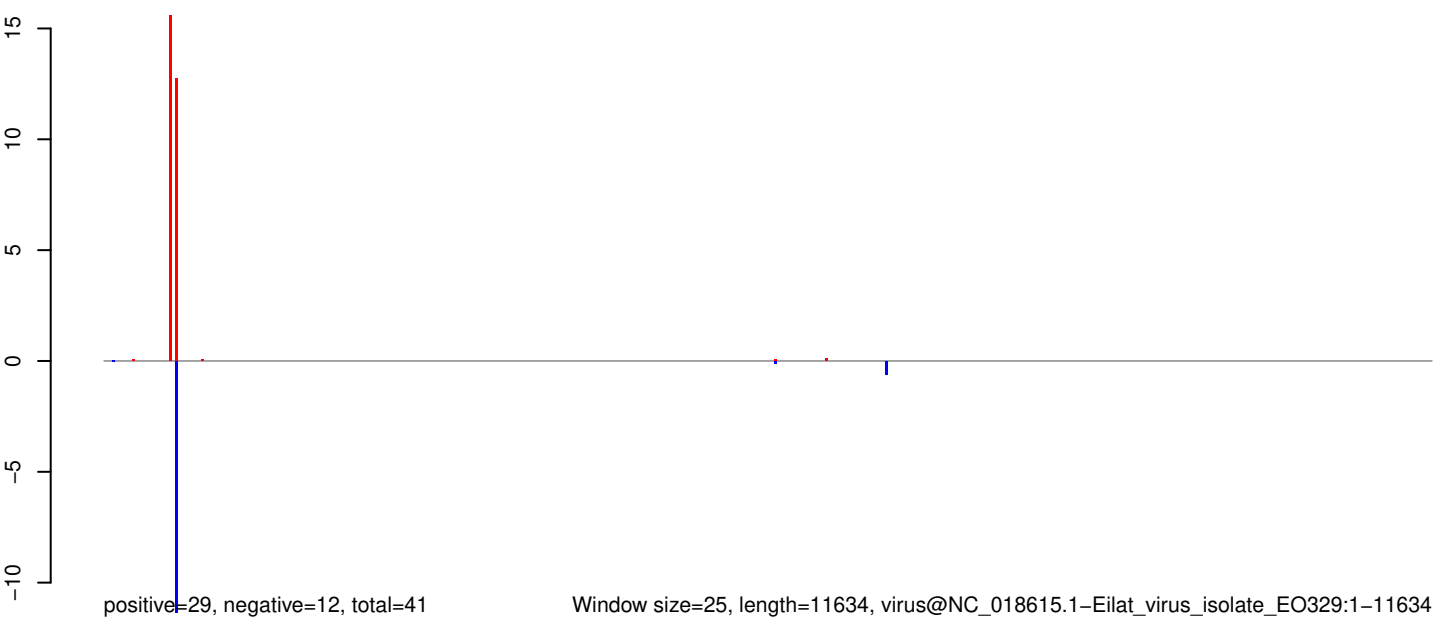
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



Window size=25, length=11634, virus@NC_018615.1-Eilat_virus_isolate_EO329:1-11634

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_dsLuc.18_23.rep

positive=10, negative=2, total=12

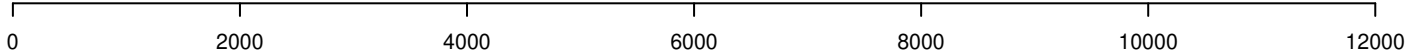
AeAeg_Aag2SINV_dsLuc.24_35.rep

positive=0, negative=0, total=0

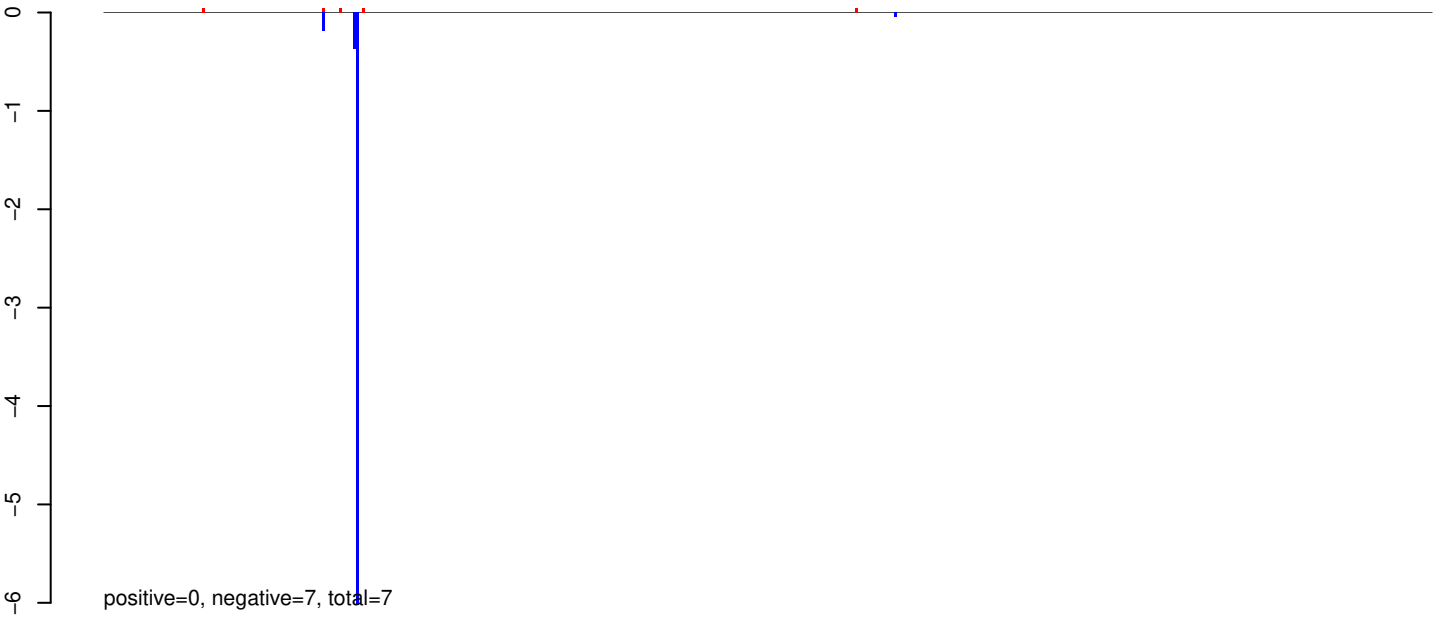
AeAeg_Aag2SINV_dsLuc.rep

positive=10, negative=2, total=12

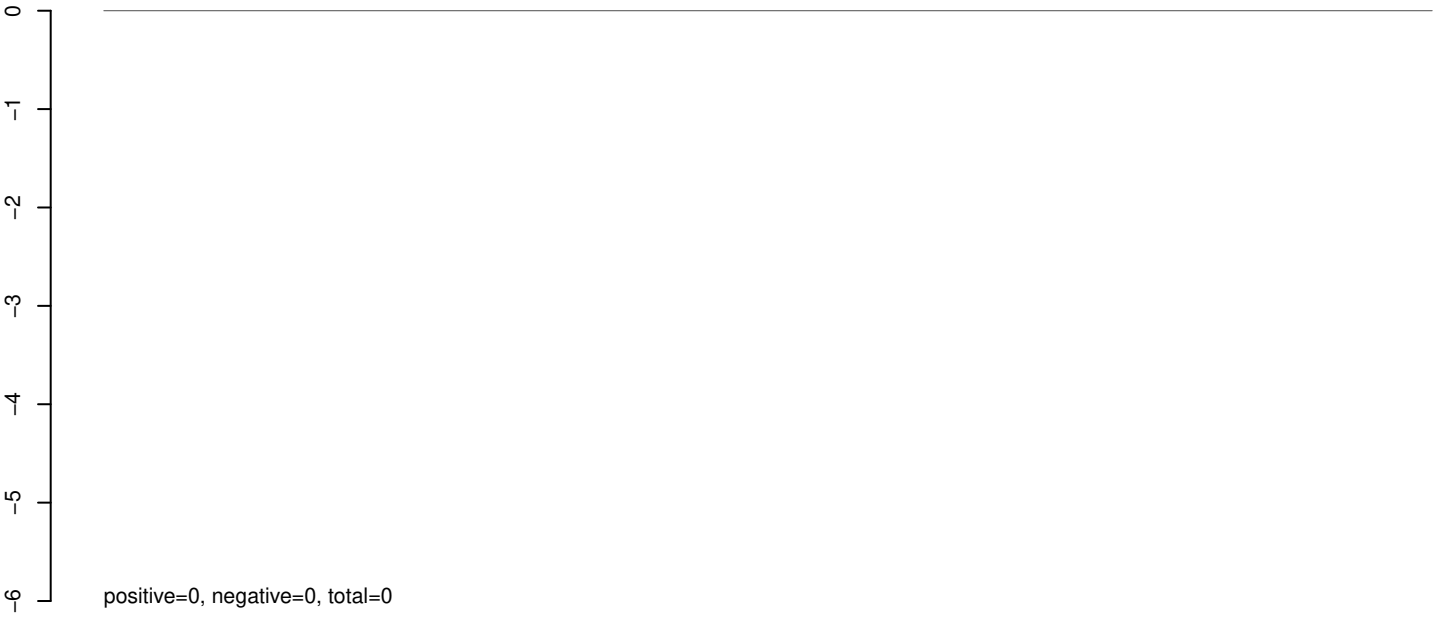
Window size=25, length=11688, virus@NC_016959.1-Ndumu_virus:1-11688



AeAeg_Aag2SINV_dsLuc.18_23.rep



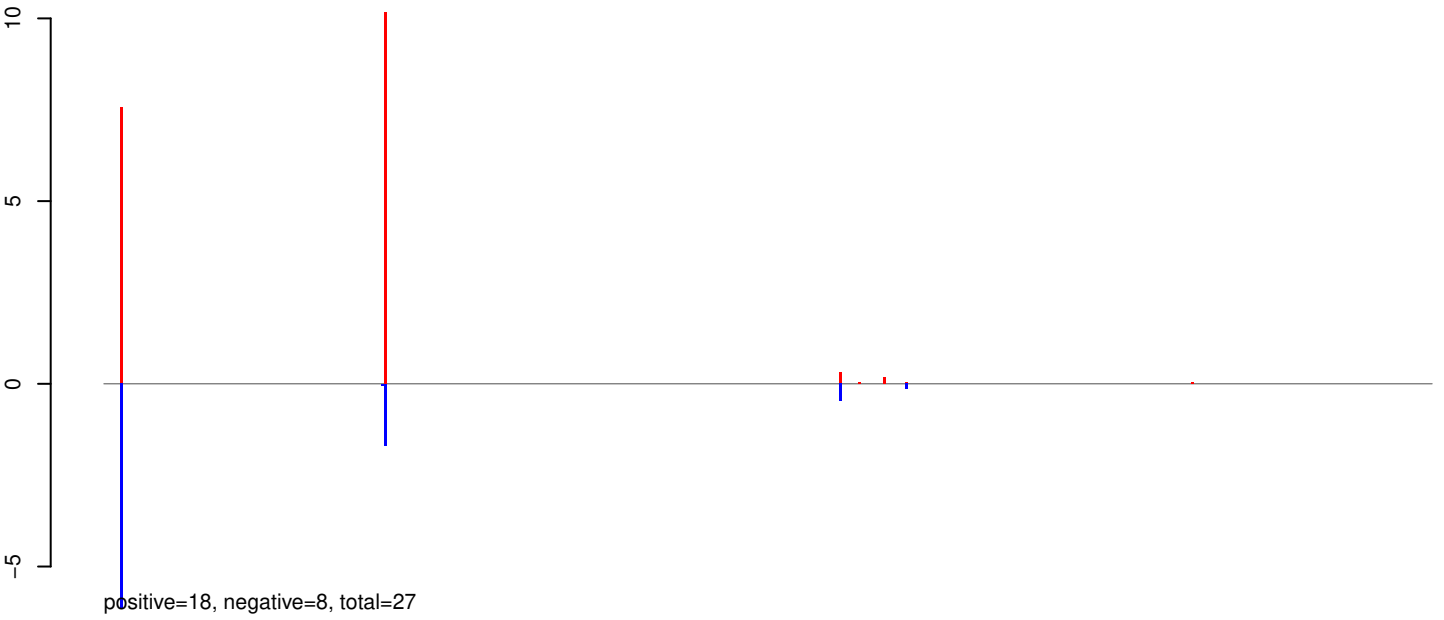
AeAeg_Aag2SINV_dsLuc.24_35.rep



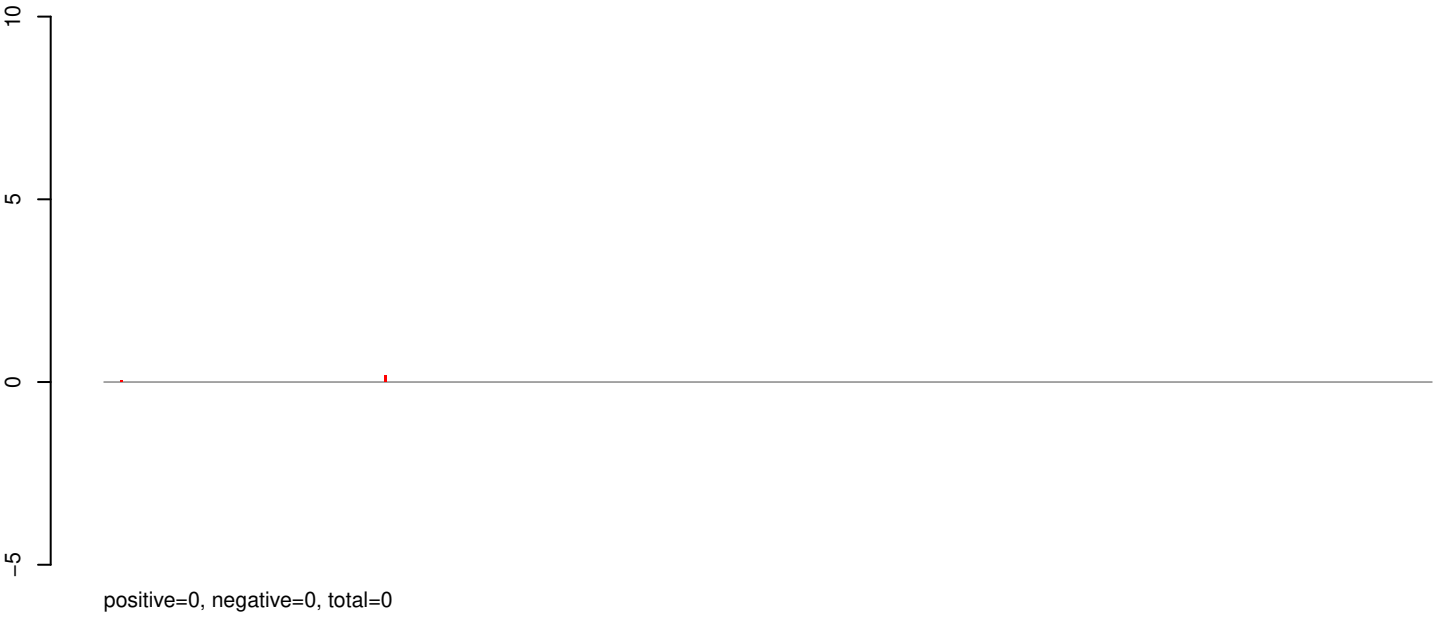
AeAeg_Aag2SINV_dsLuc.rep



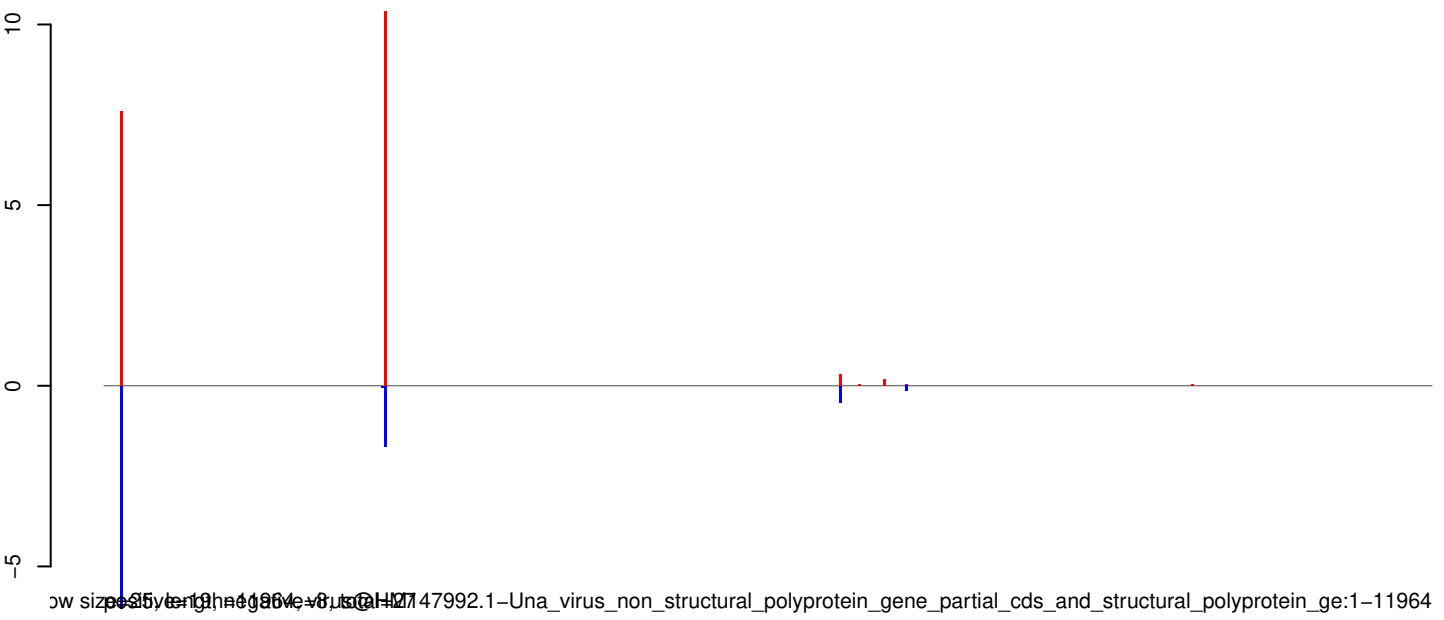
AeAeg_Aag2SINV_dsLuc.18_23.rep



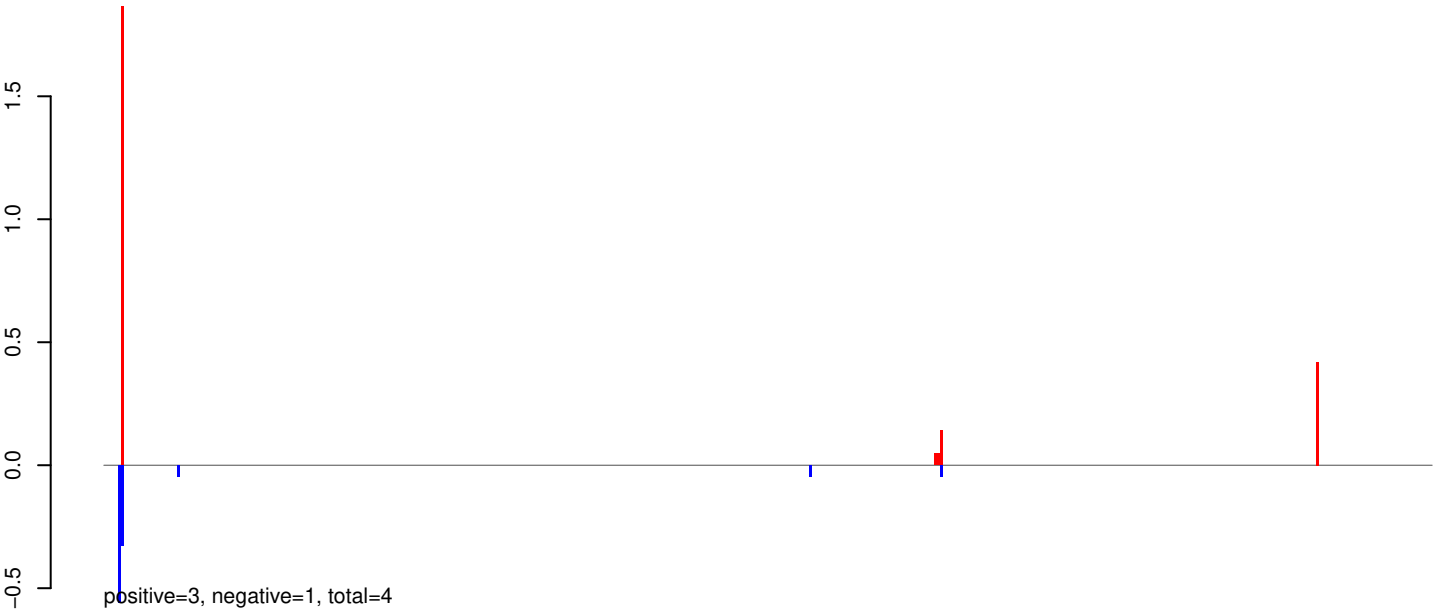
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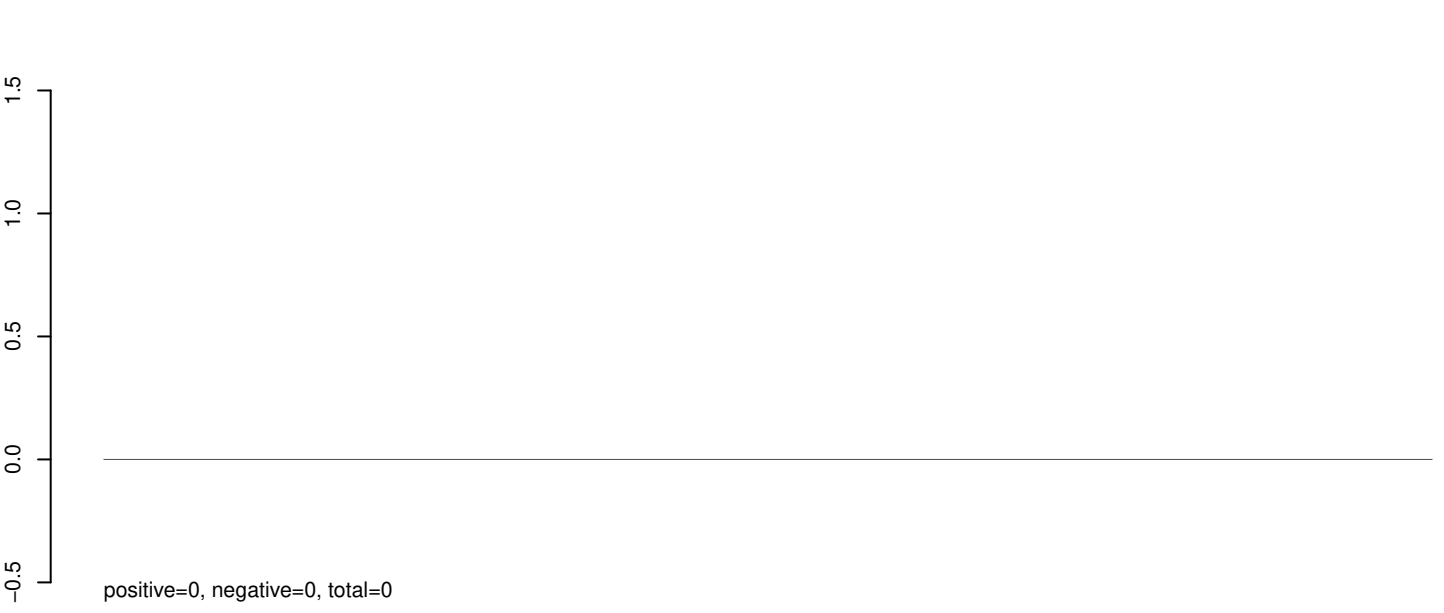
AeAeg_Aag2SINV_dsLuc.rep



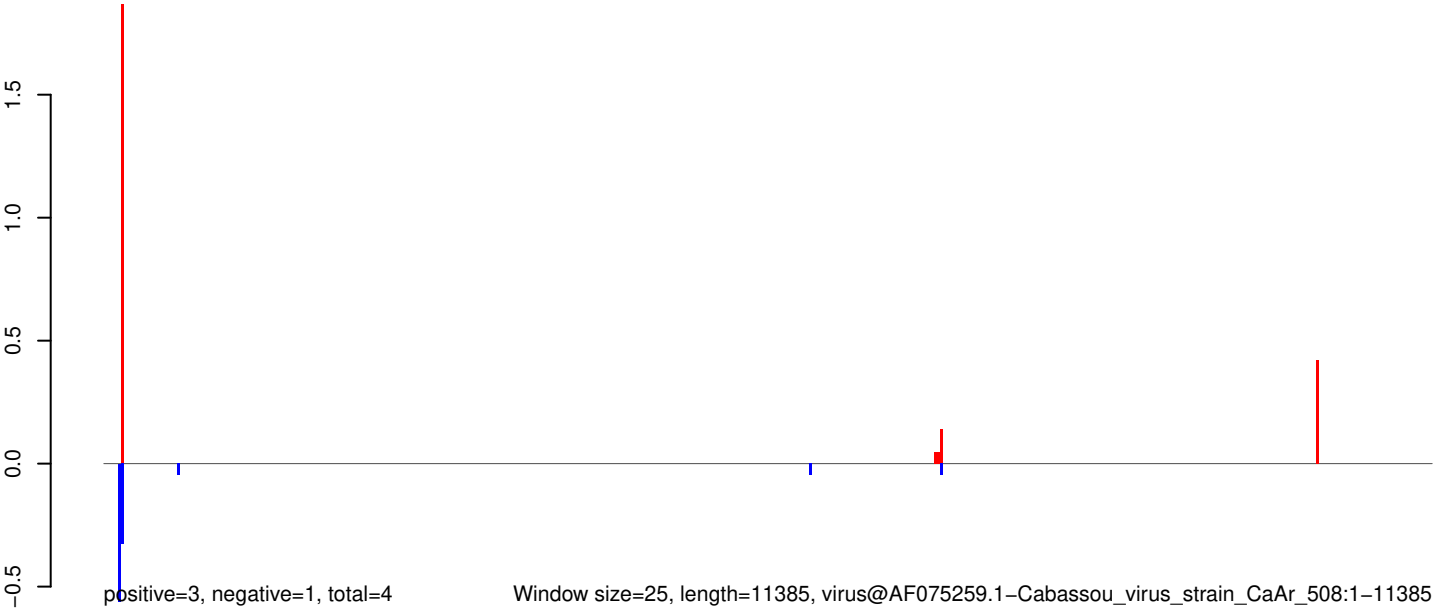
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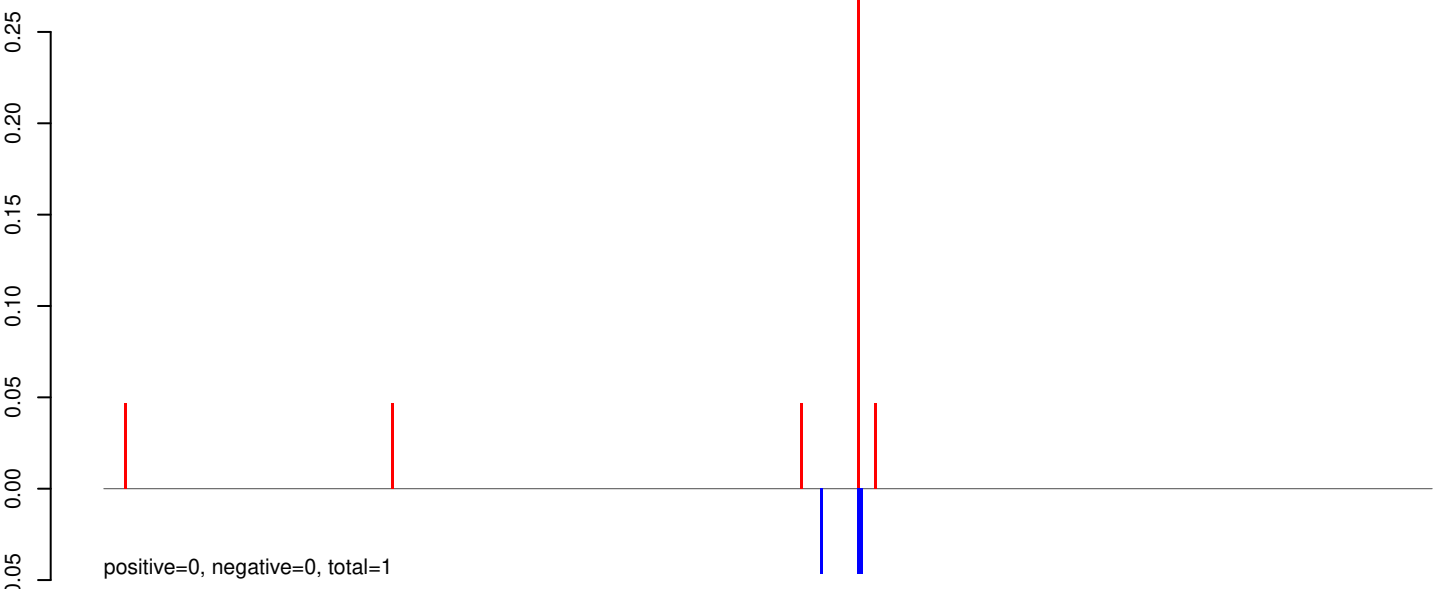
AeAeg_Aag2SINV_dsLuc.24_35.rep



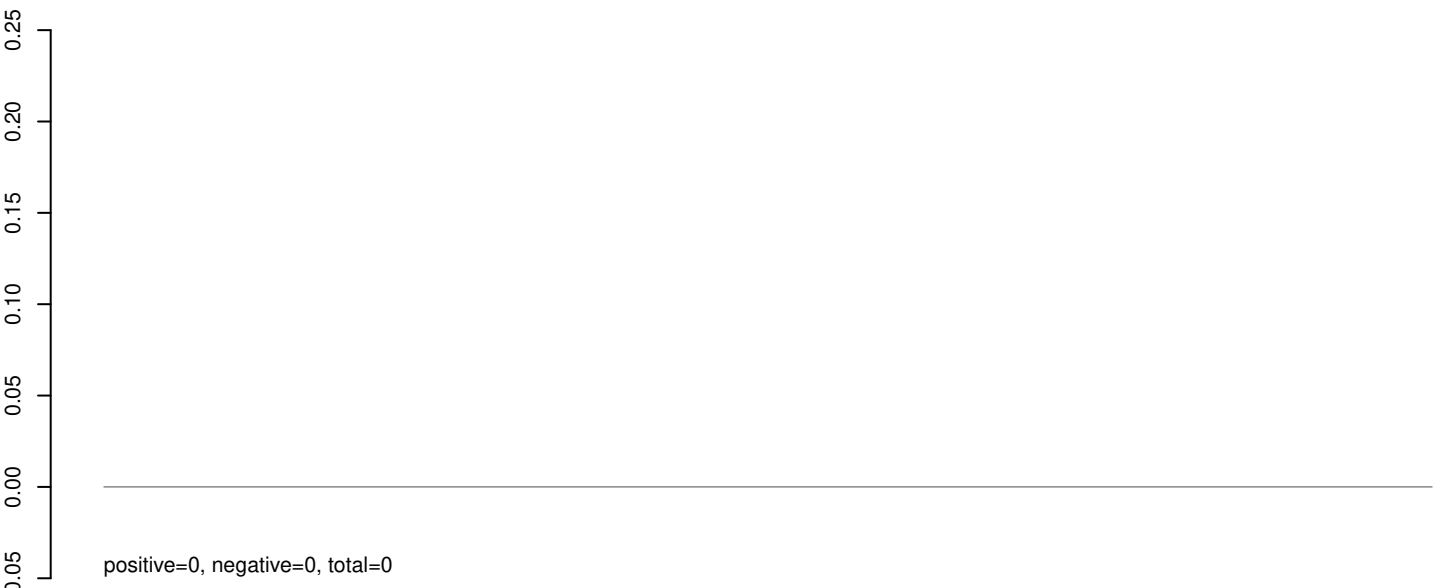
AeAeg_Aag2SINV_dsLuc.rep



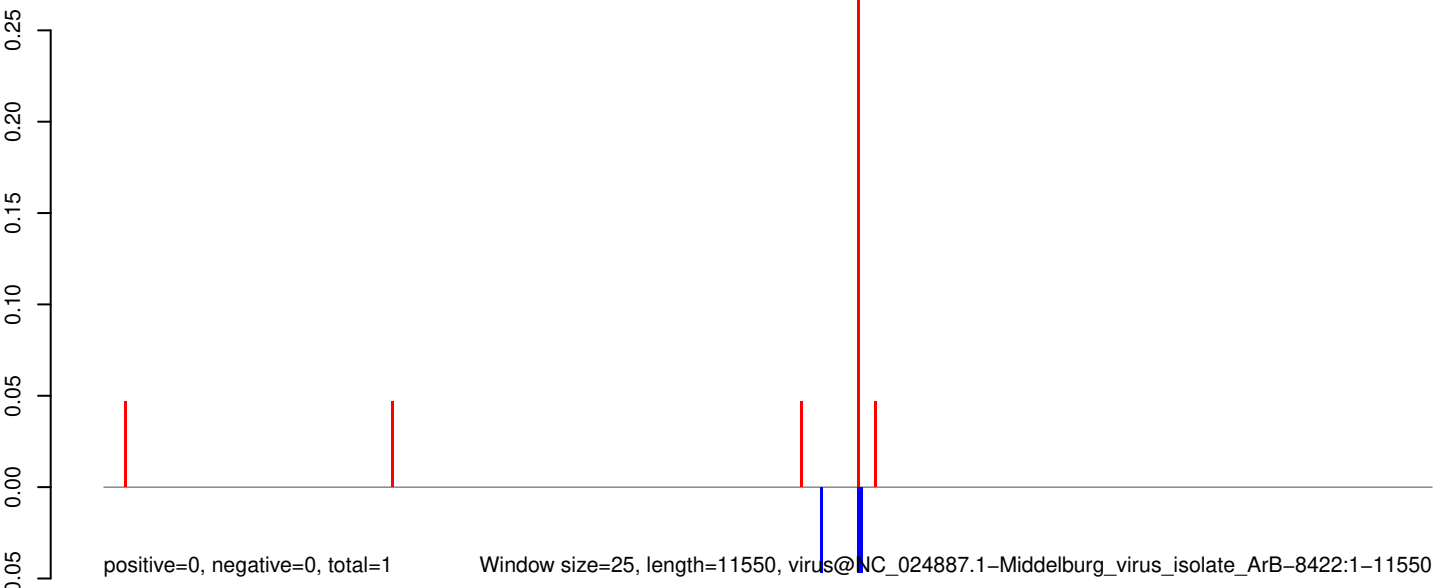
AeAeg_Aag2SINV_dsLuc.18_23.rep



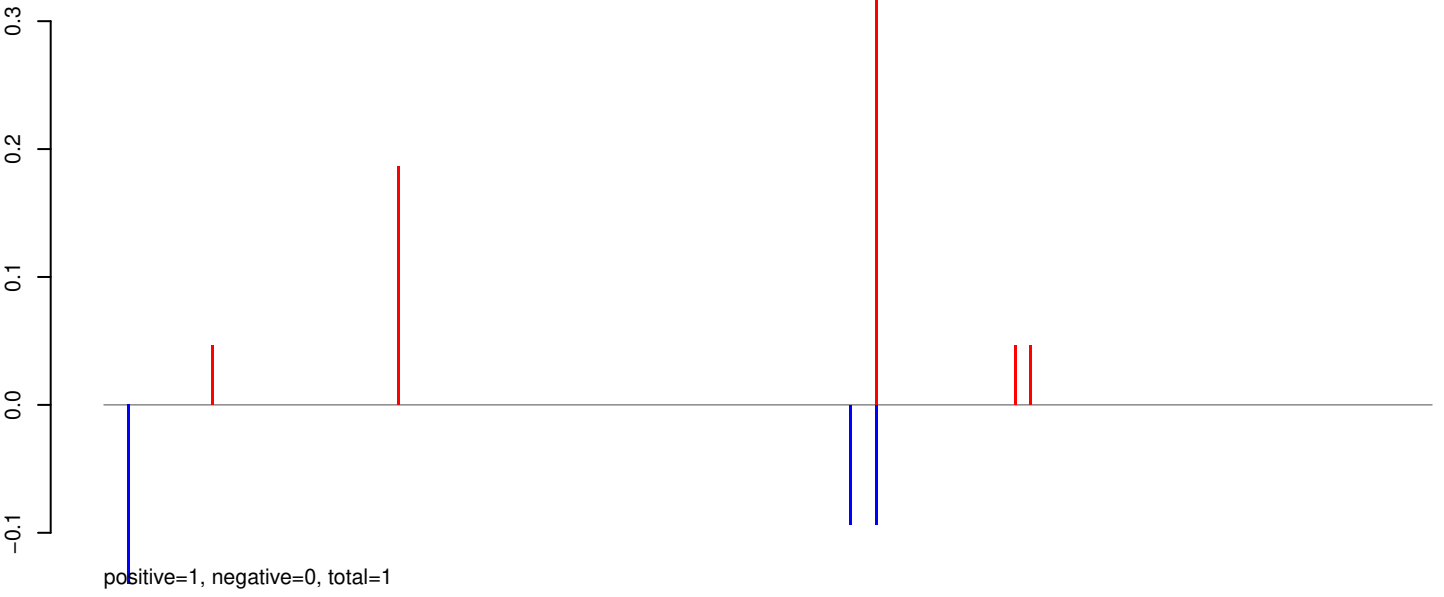
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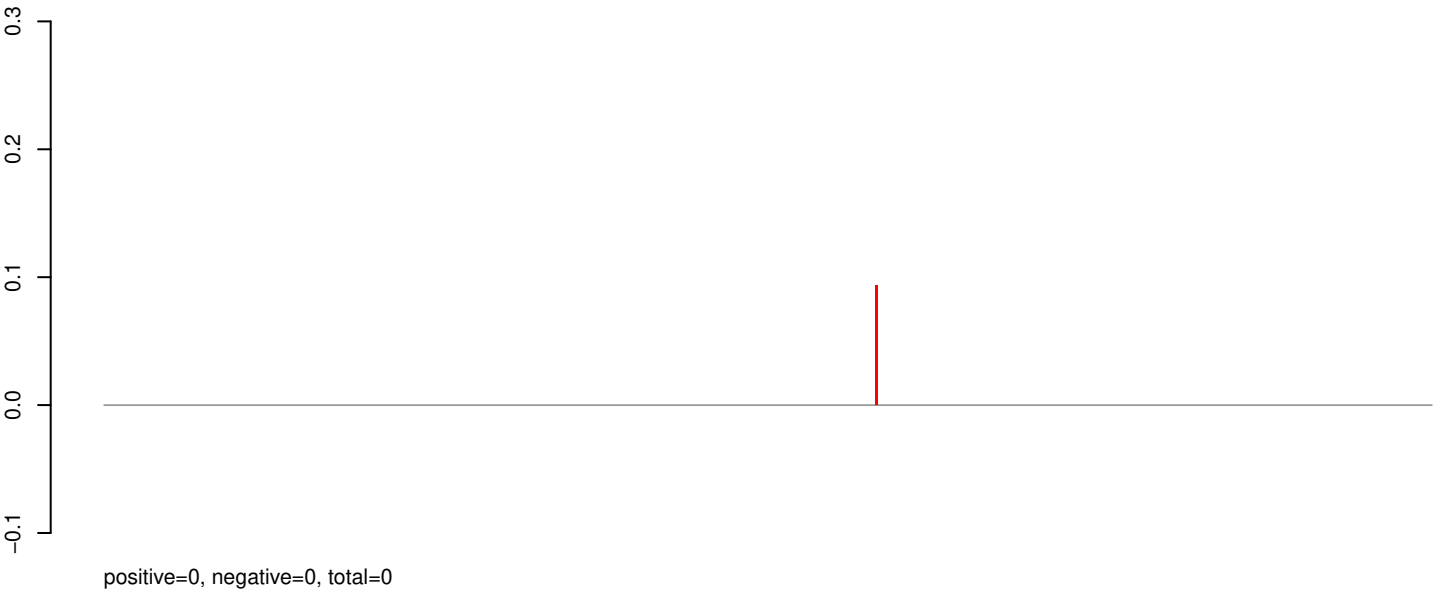
AeAeg_Aag2SINV_dsLuc.rep



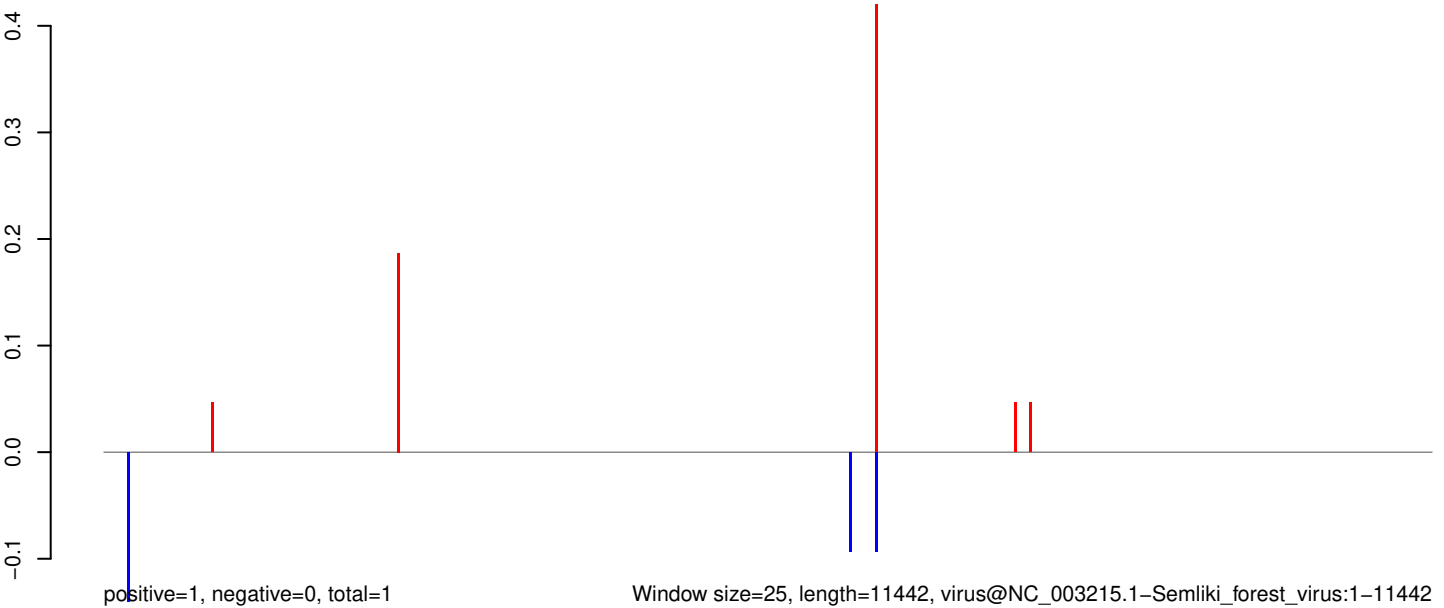
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AeAeg_Aag2SINV_dsLuc.24_35.rep



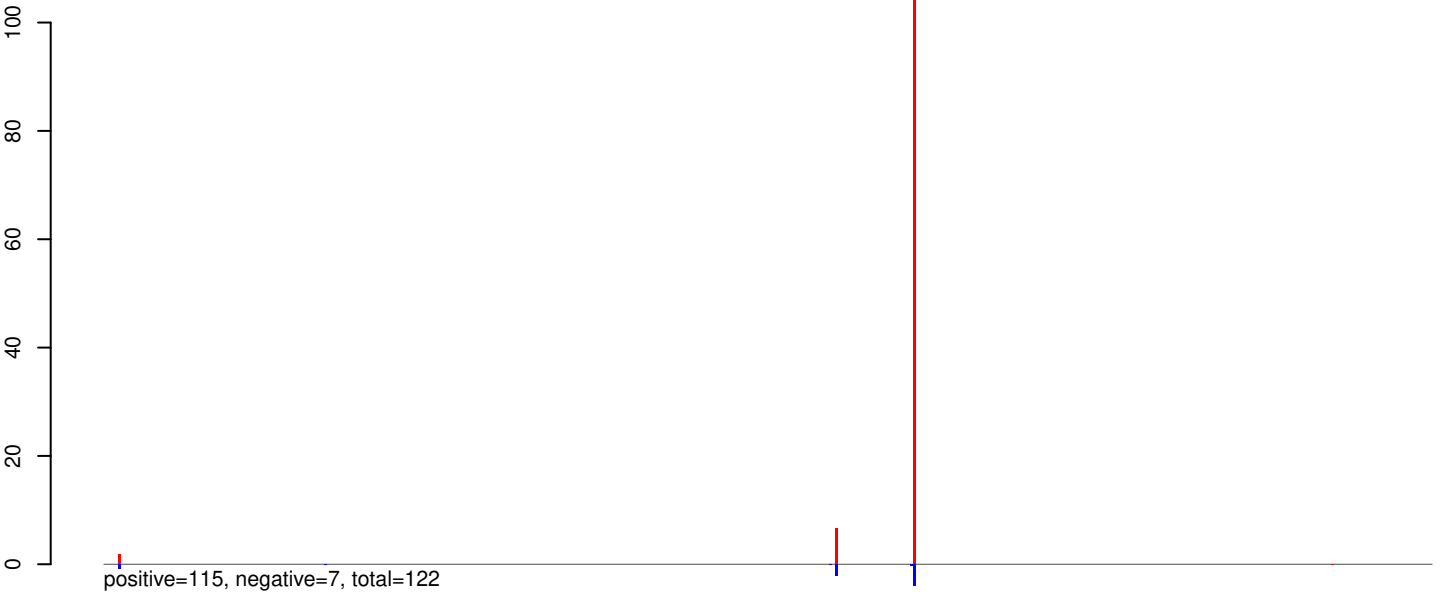
AeAeg_Aag2SINV_dsLuc.rep



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

0 2000 4000 6000 8000 10000

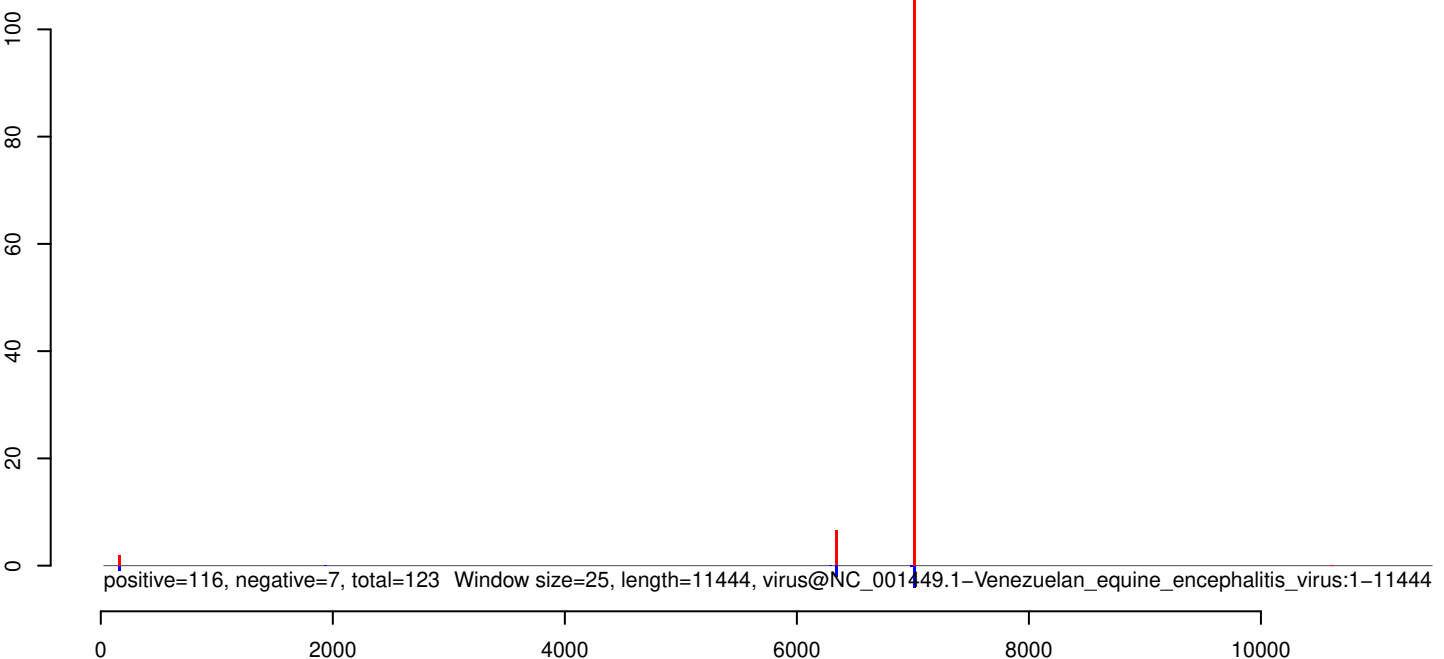
AeAeg_Aag2SINV_dsLuc.18_23.rep



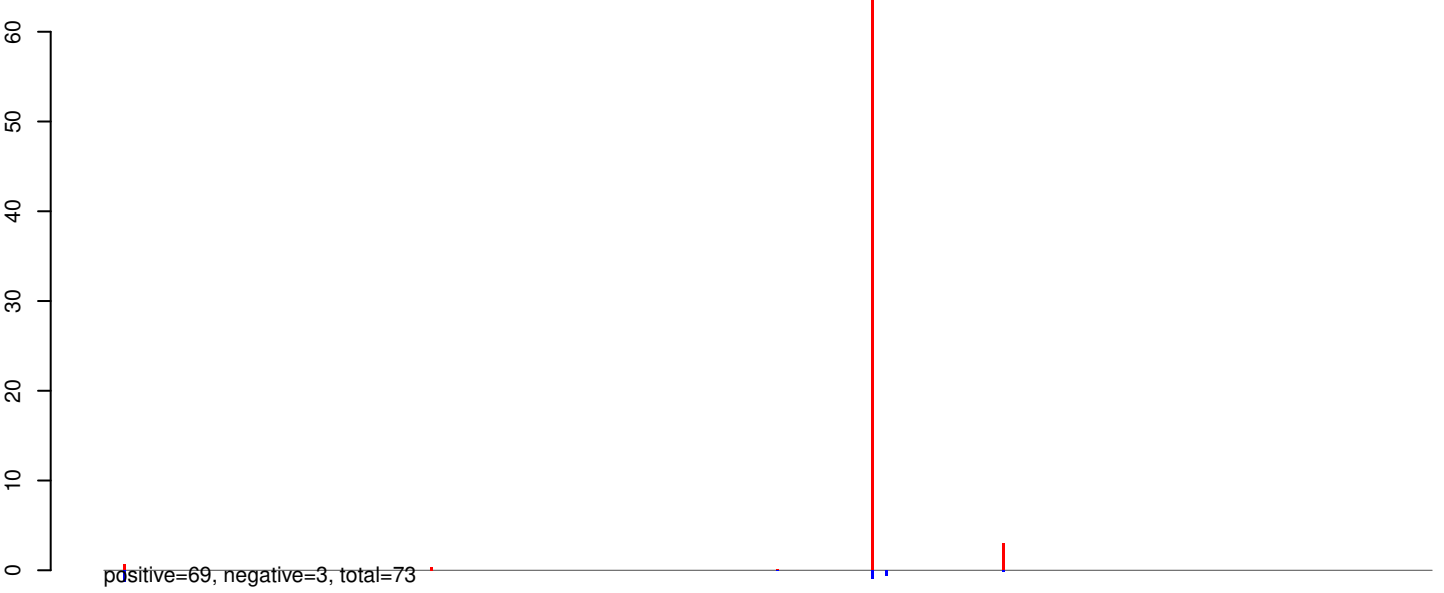
AeAeg_Aag2SINV_dsLuc.24_35.rep



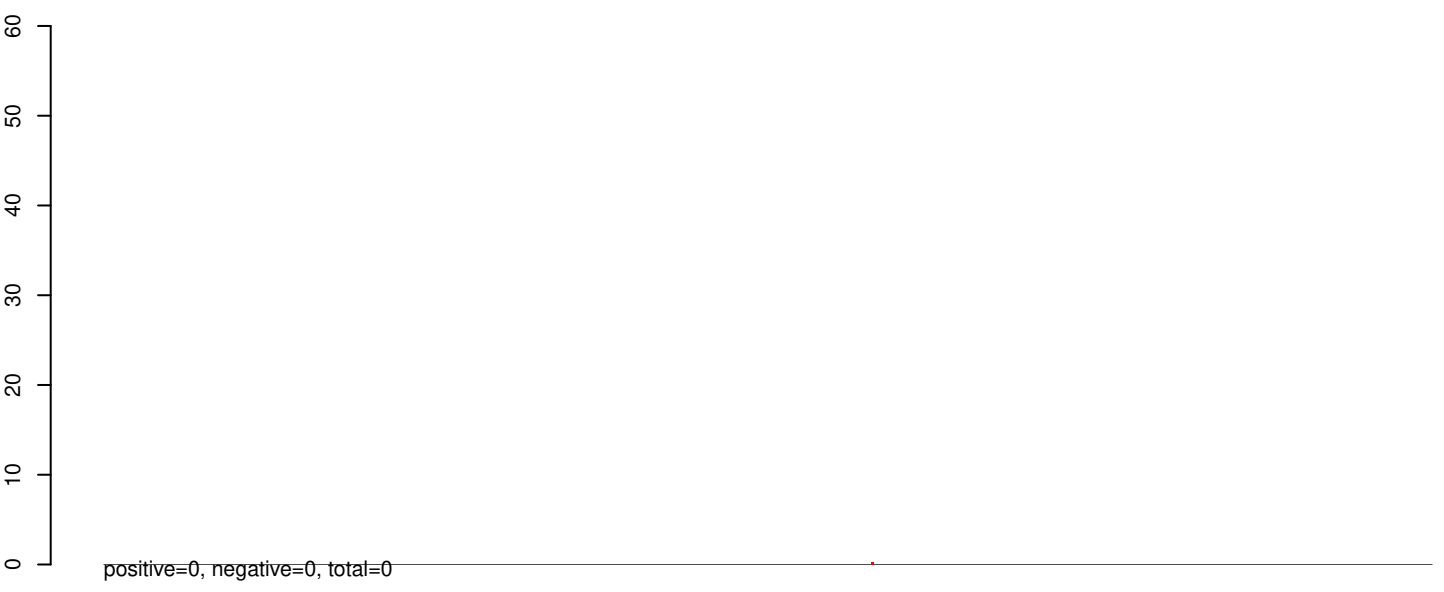
AeAeg_Aag2SINV_dsLuc.rep



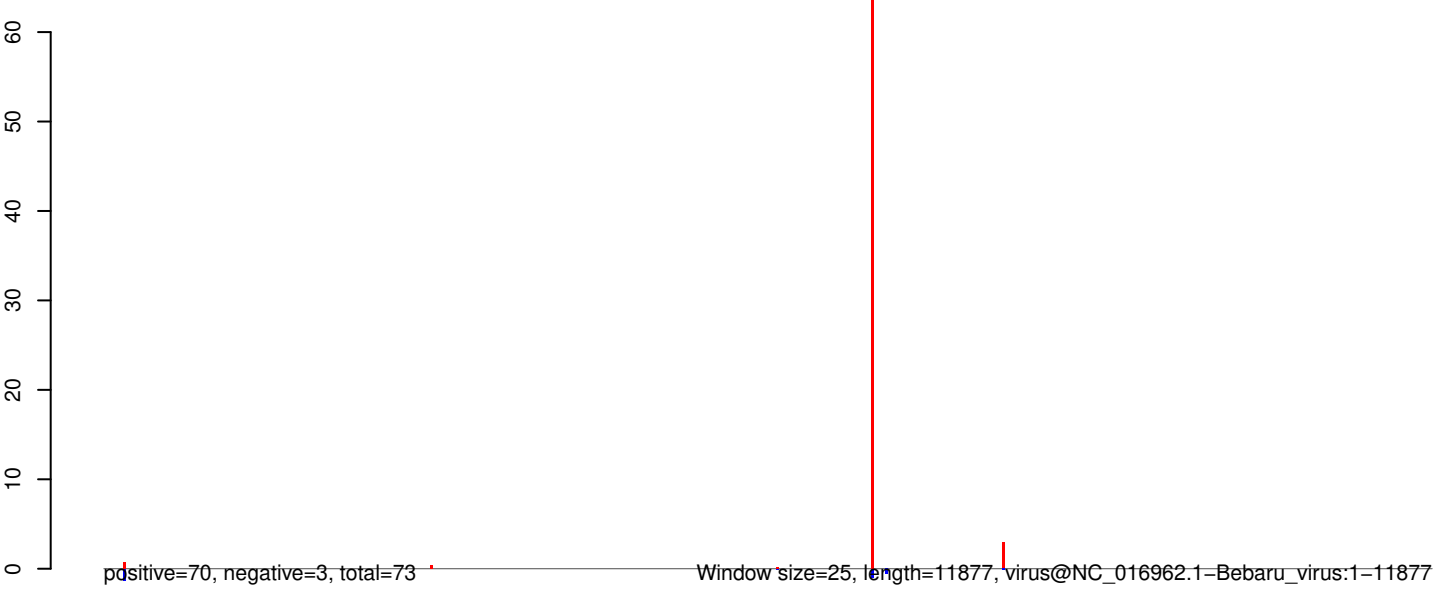
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

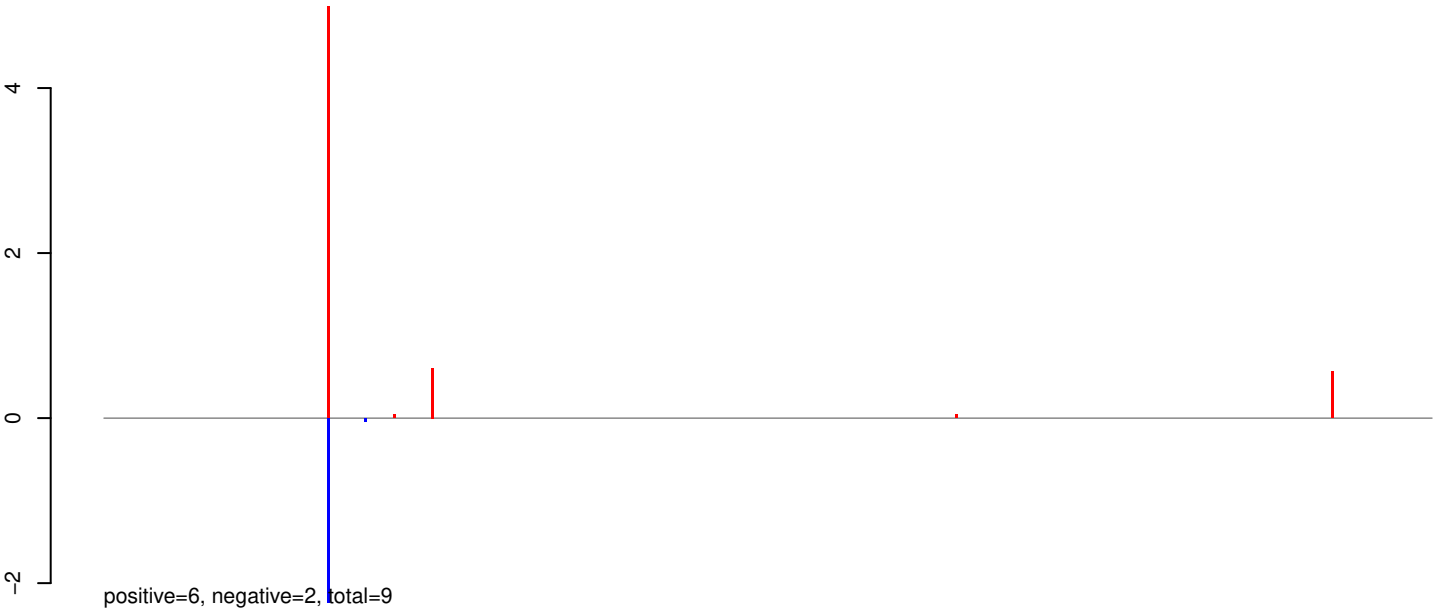


AeAeg_Aag2SINV_dsLuc.rep

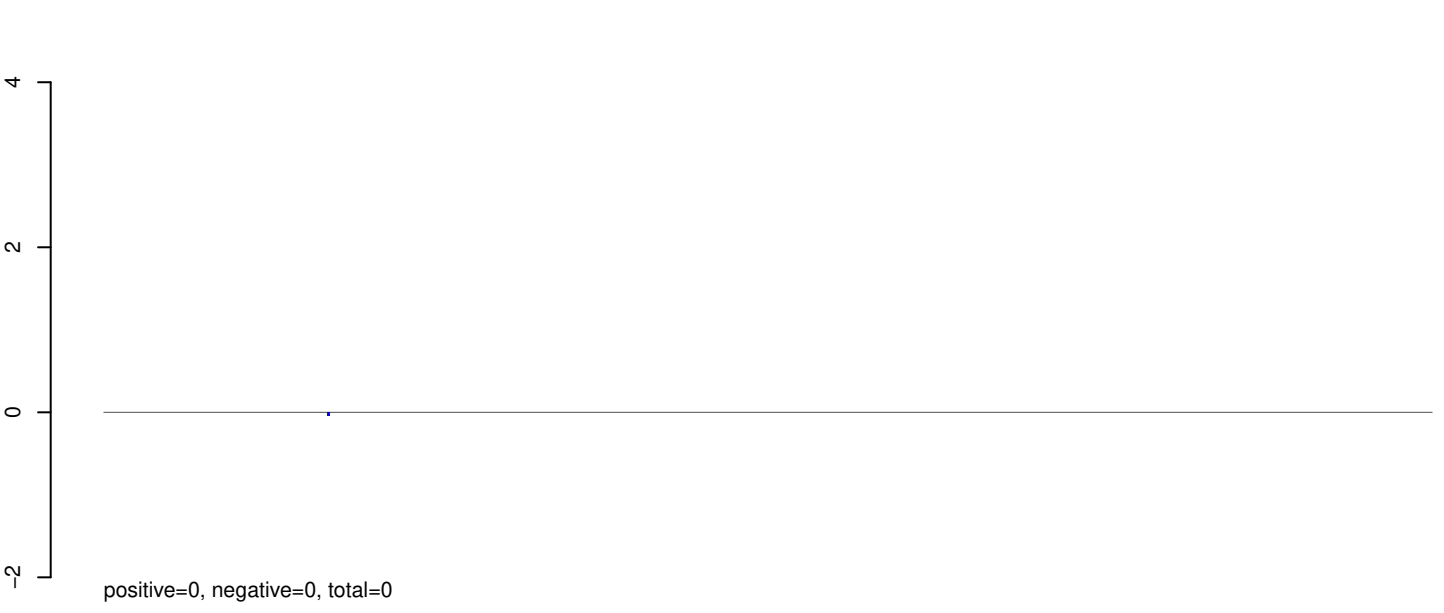


0 2000 4000 6000 8000 10000 12000

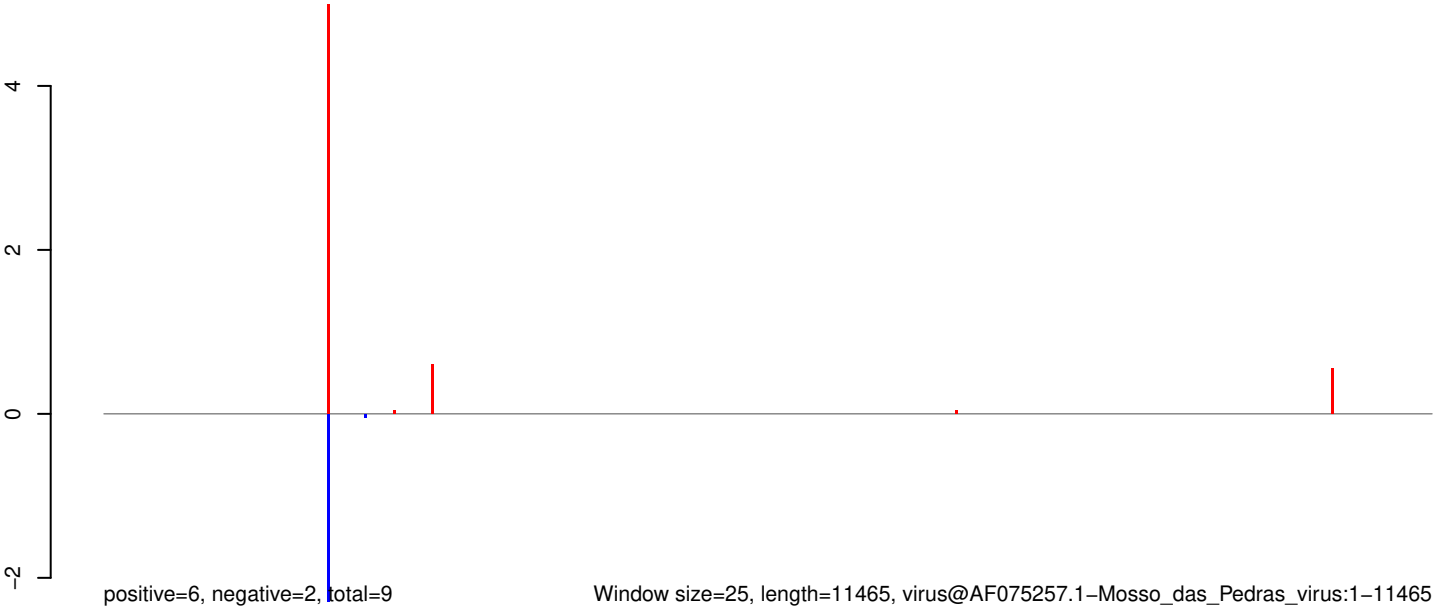
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



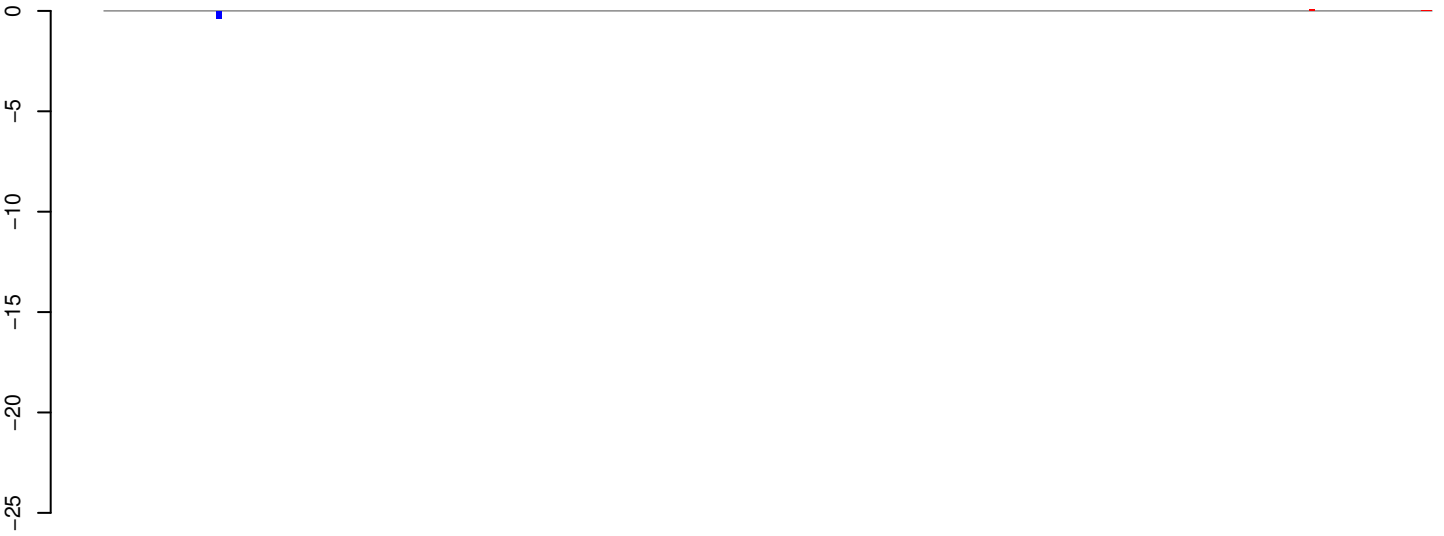
AeAeg_Aag2SINV_dsLuc.rep



Window size=25, length=11465, virus@AF075257.1-Mosso_das_Pedras_virus:1-11465

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_dsLuc.18_23.rep



positive=0, negative=0, total=1

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=2, negative=29, total=31

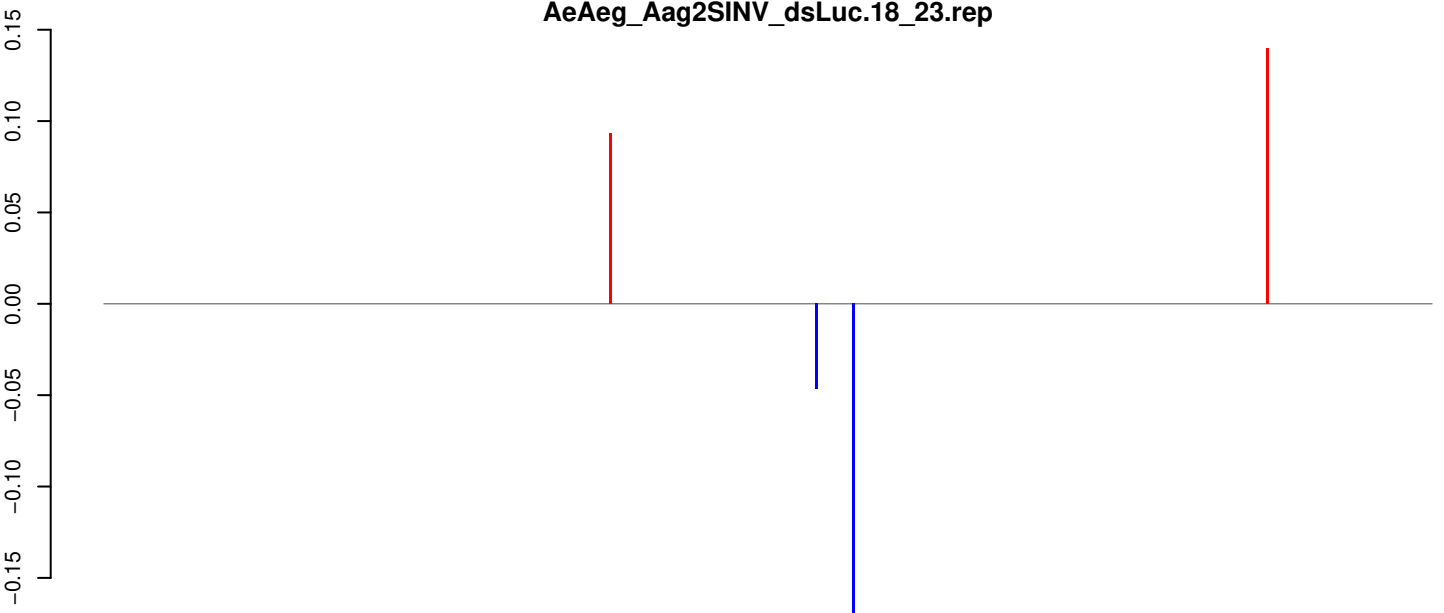
AeAeg_Aag2SINV_dsLuc.rep



positive=2, negative=29, window size=25, length=6203, virus@NC_035674.1-Australian_Anopheles_totivirus_isolate_AATV_150840:1-6203

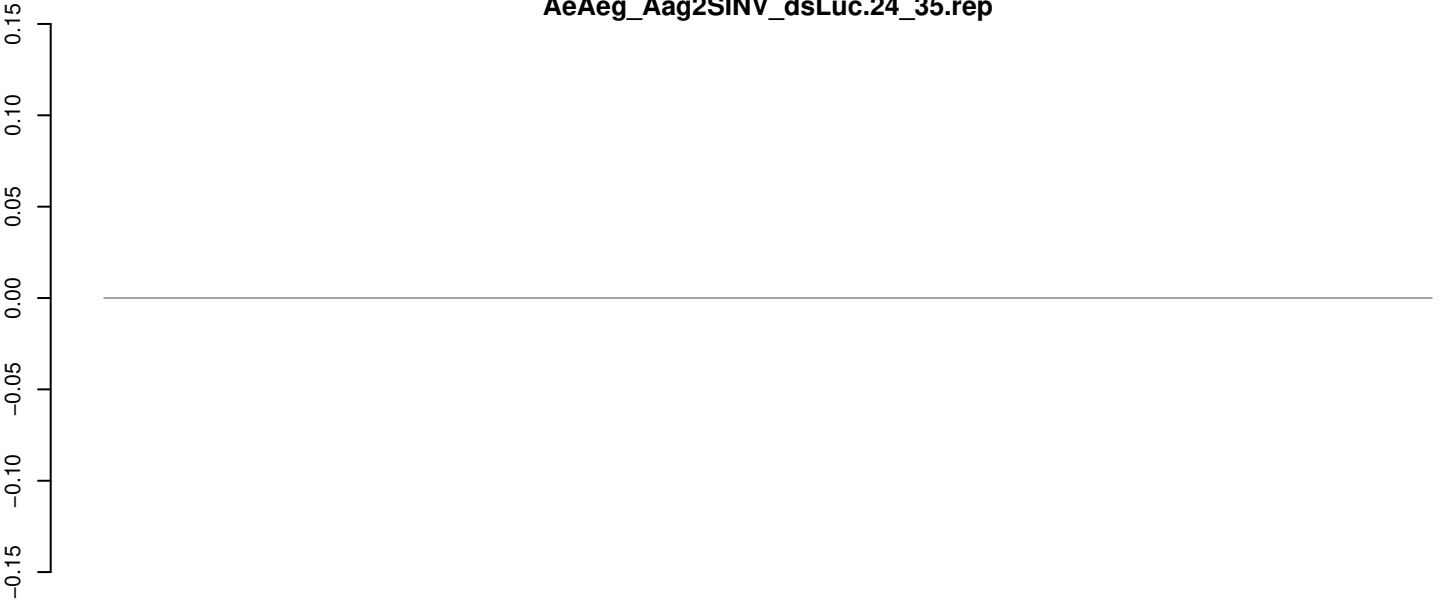
0 1000 2000 3000 4000 5000 6000

AeAeg_Aag2SINV_dsLuc.18_23.rep



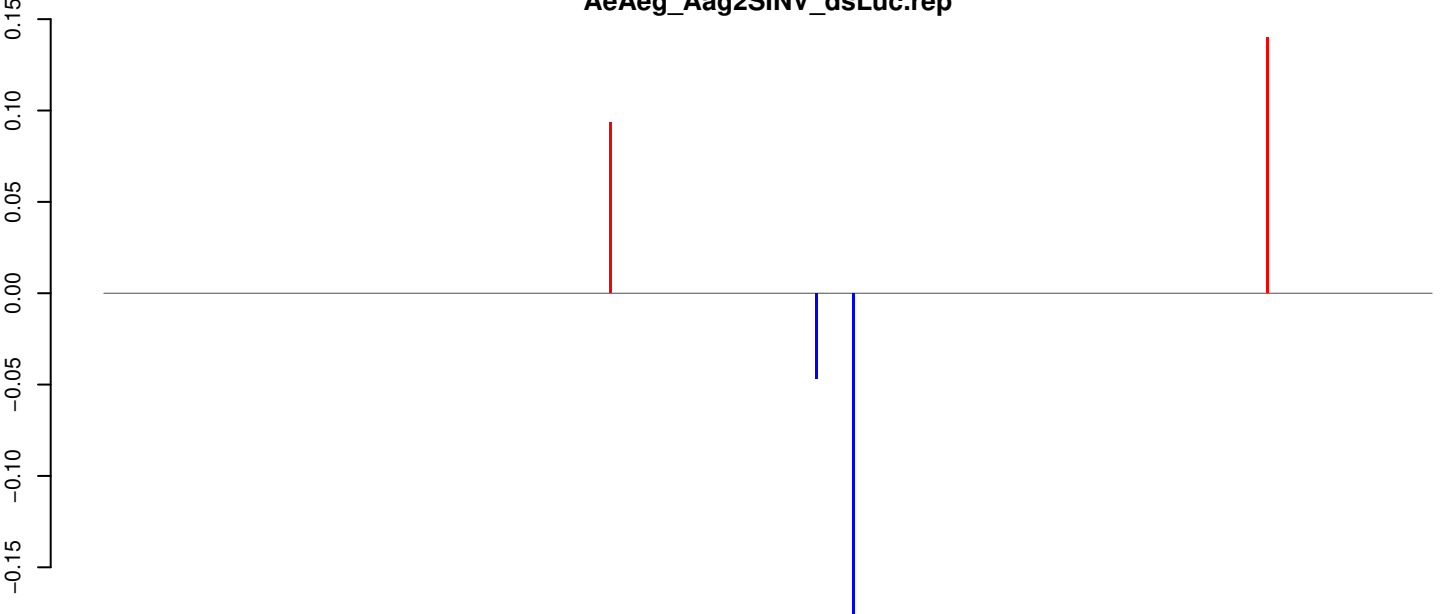
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=0, negative=0, total=0

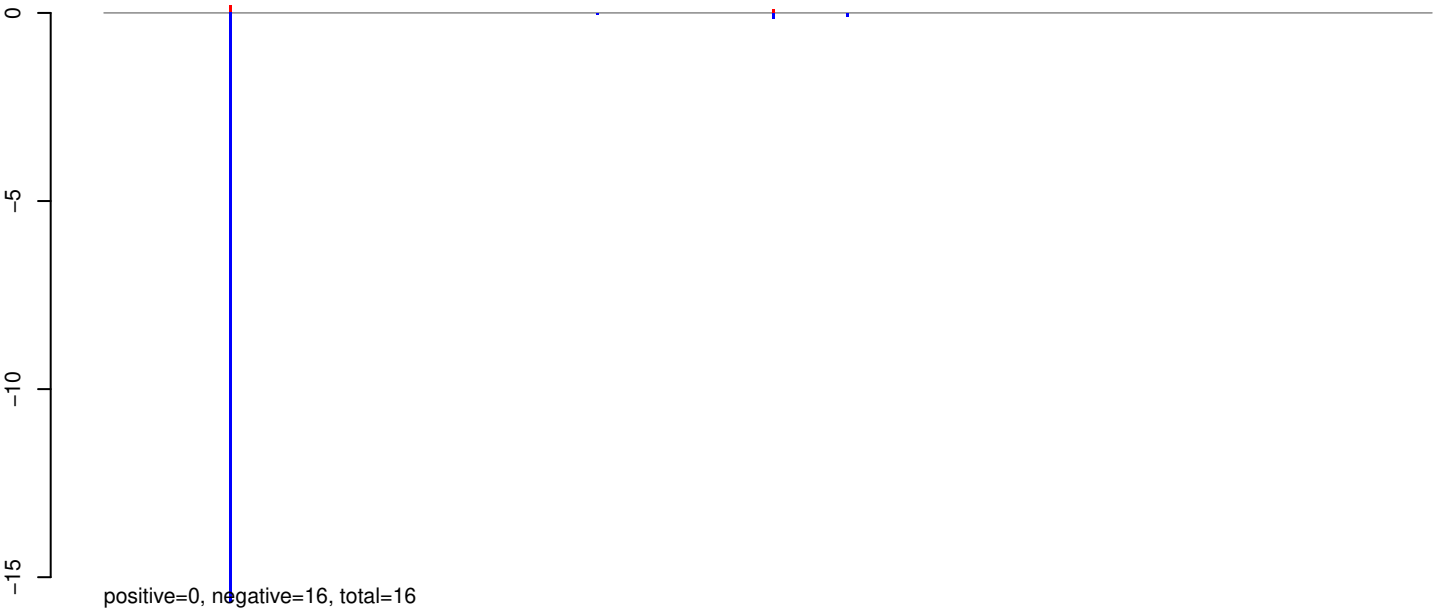
AeAeg_Aag2SINV_dsLuc.rep



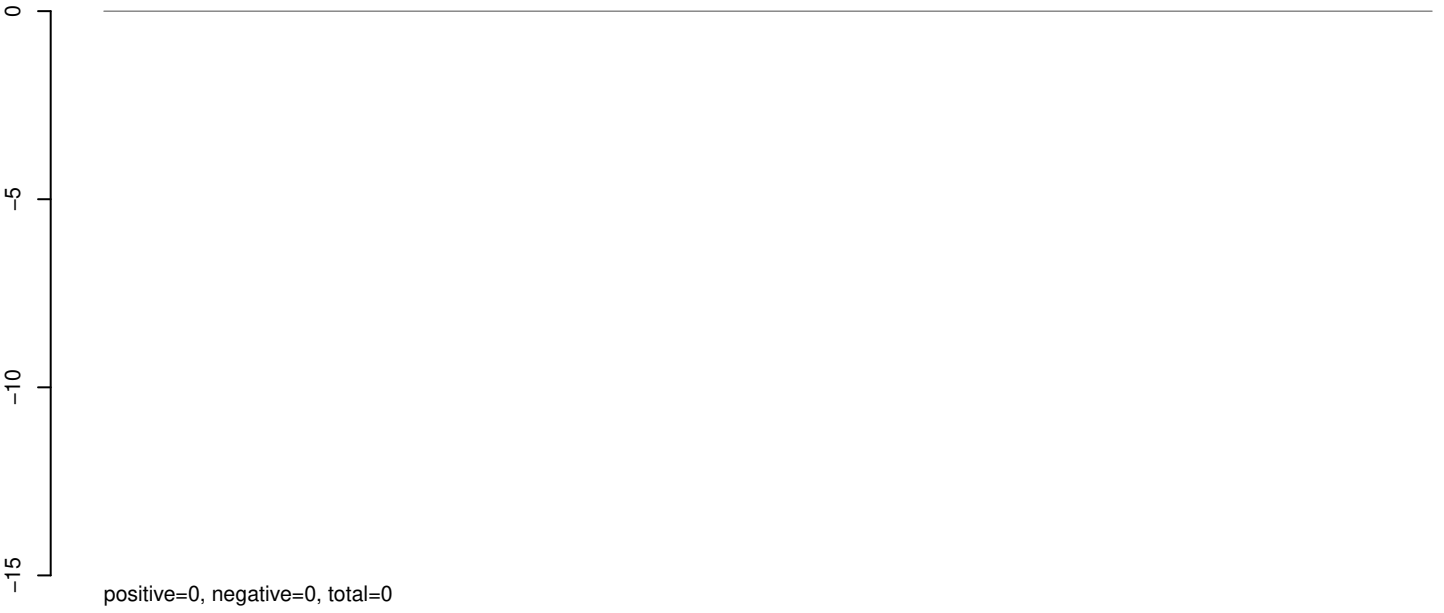
positive=0, negative=25, total=25, length=11624, virus@NC_023812.1-Madariaga_virus_strain_MADV_Cebus_apella_BRA_BEAN5122_1956:1-11624

0 2000 4000 6000 8000 10000 12000

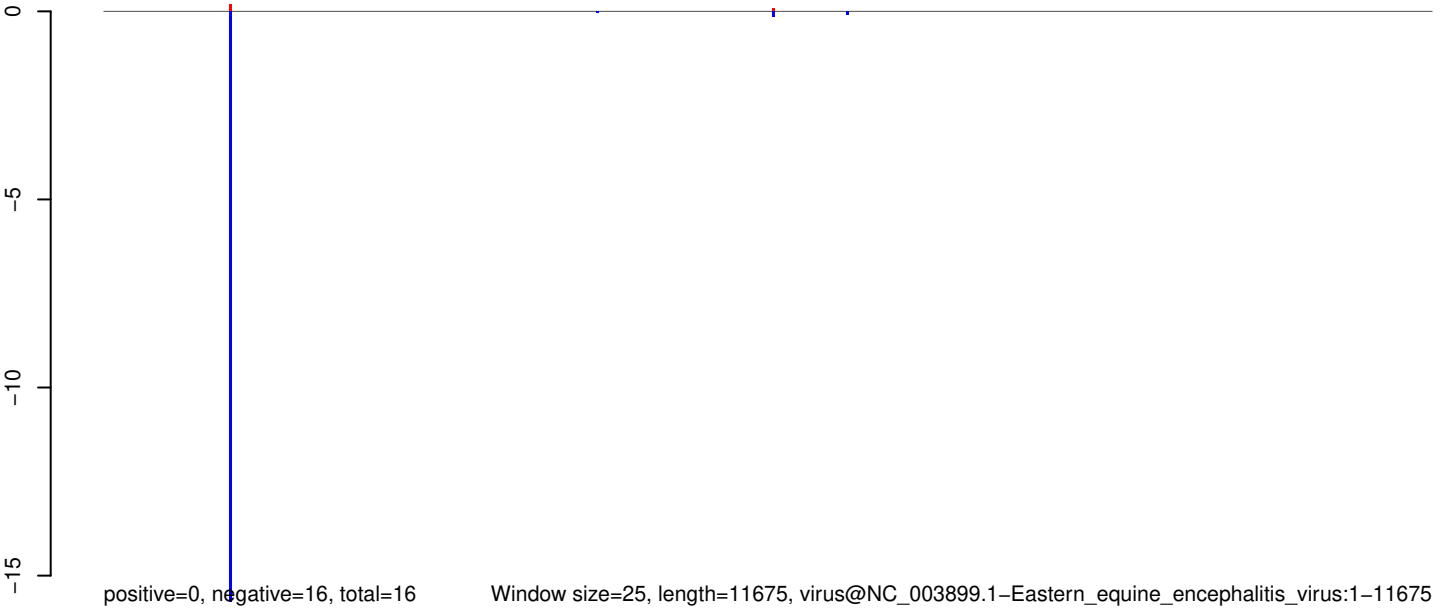
AeAeg_Aag2SINV_dsLuc.18_23.rep



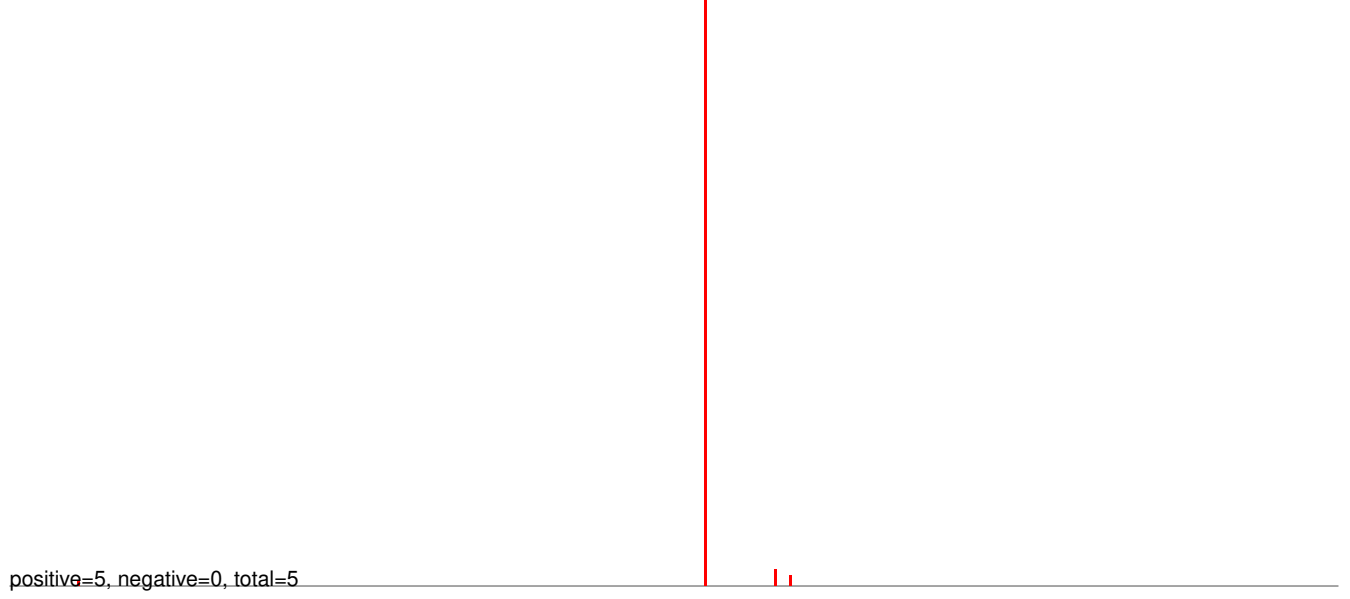
AeAeg_Aag2SINV_dsLuc.24_35.rep



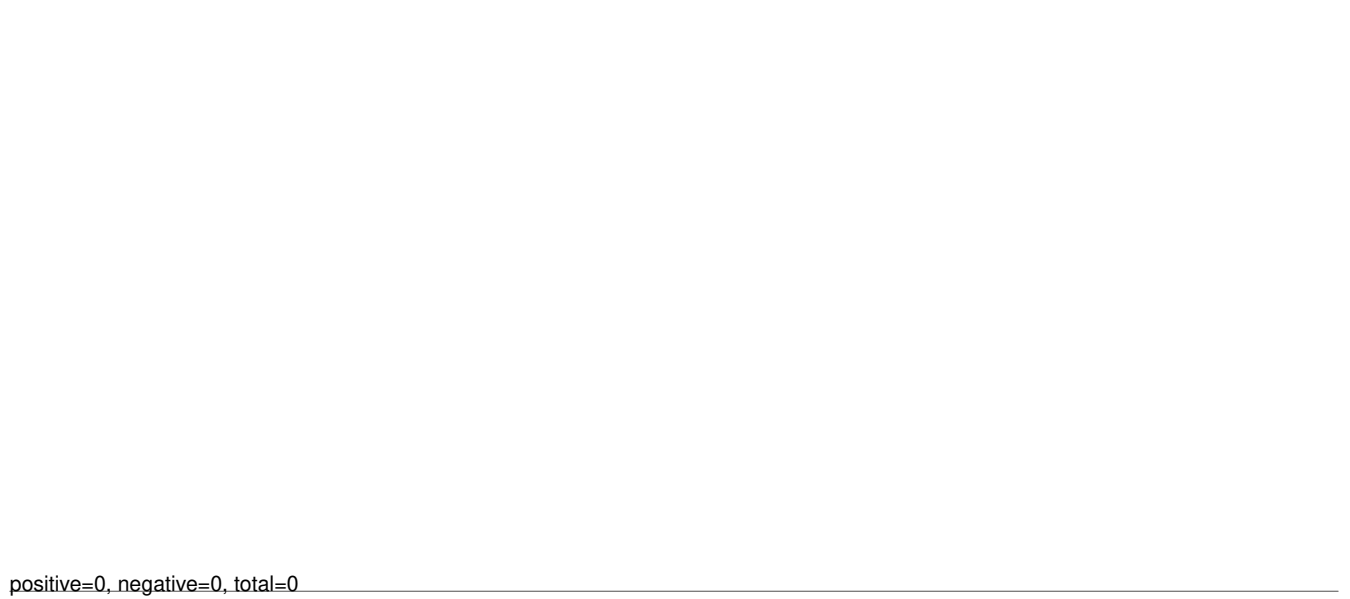
AeAeg_Aag2SINV_dsLuc.rep



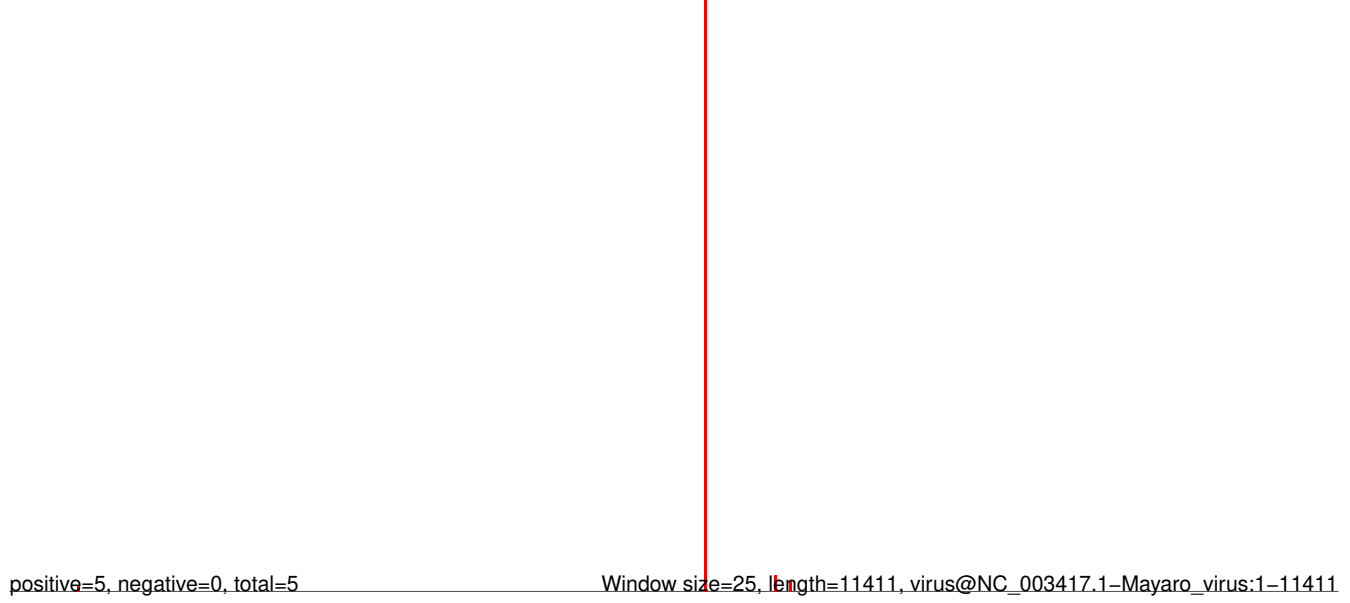
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

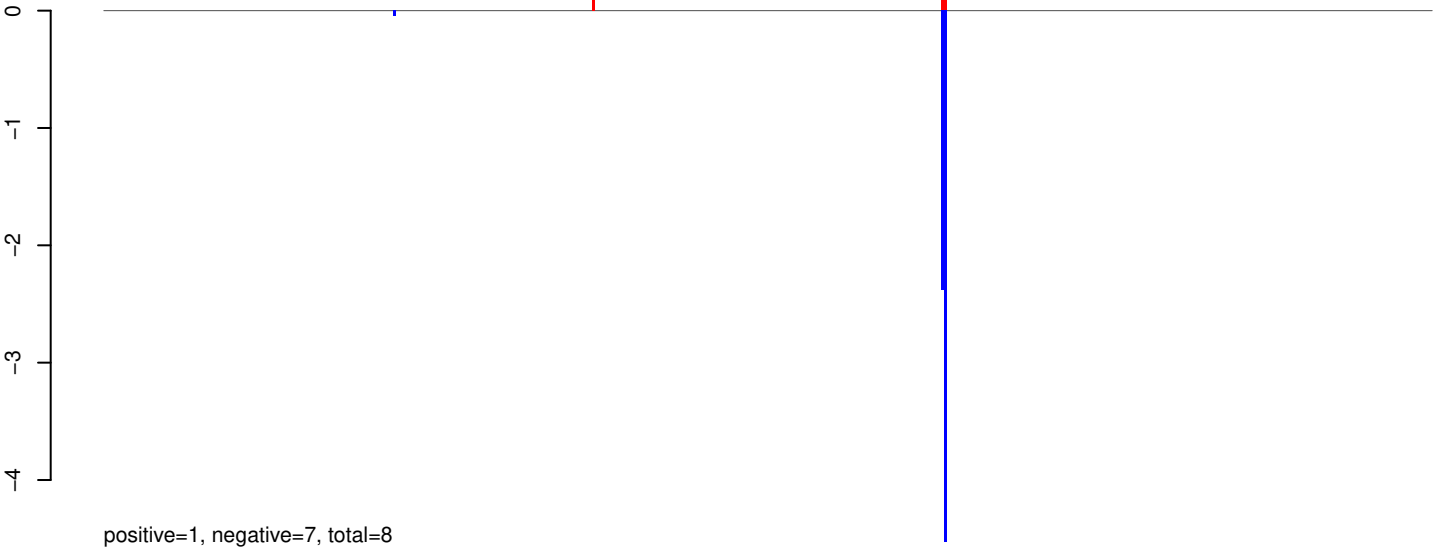


AeAeg_Aag2SINV_dsLuc.rep

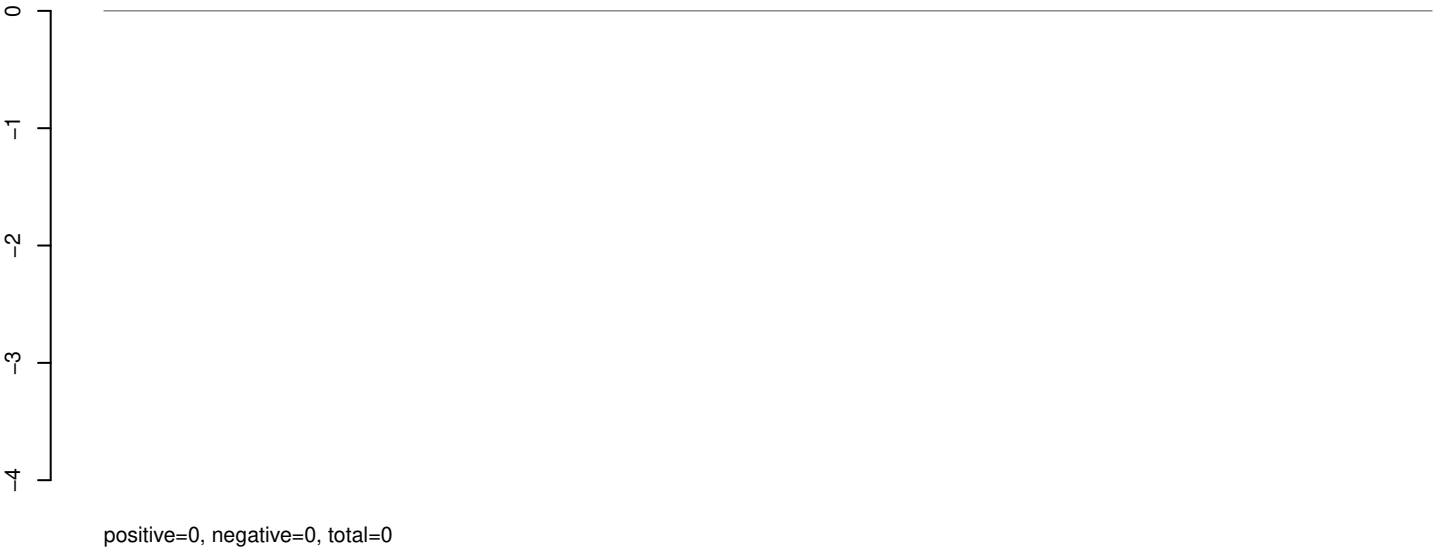


0 2000 4000 6000 8000 10000

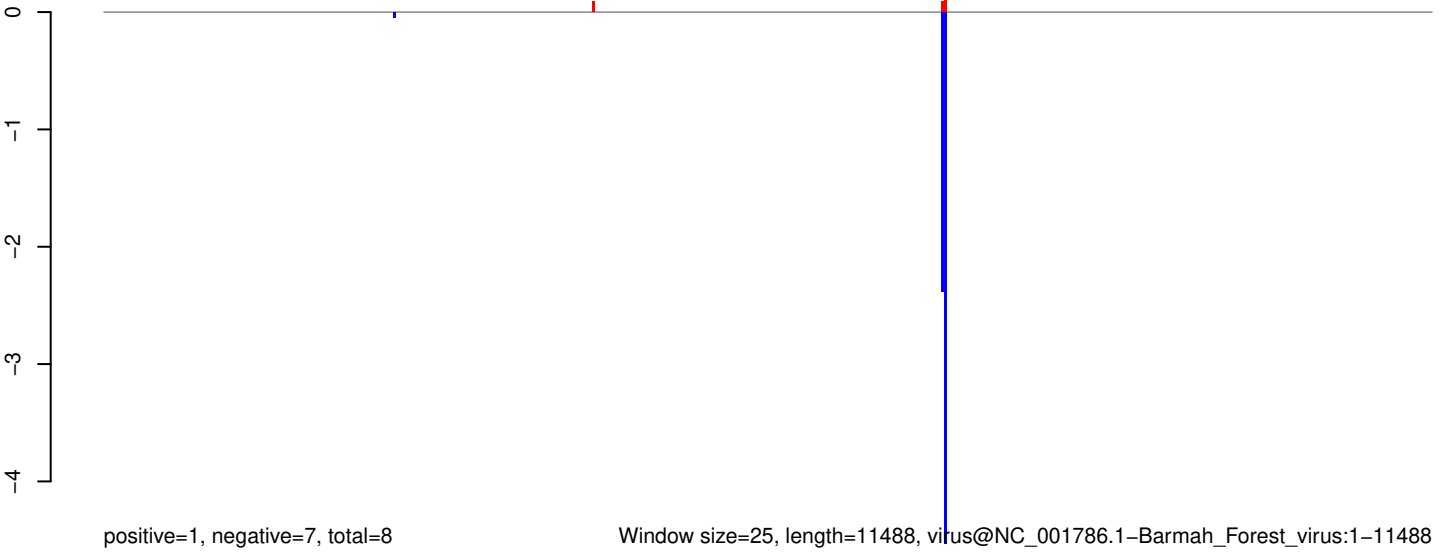
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



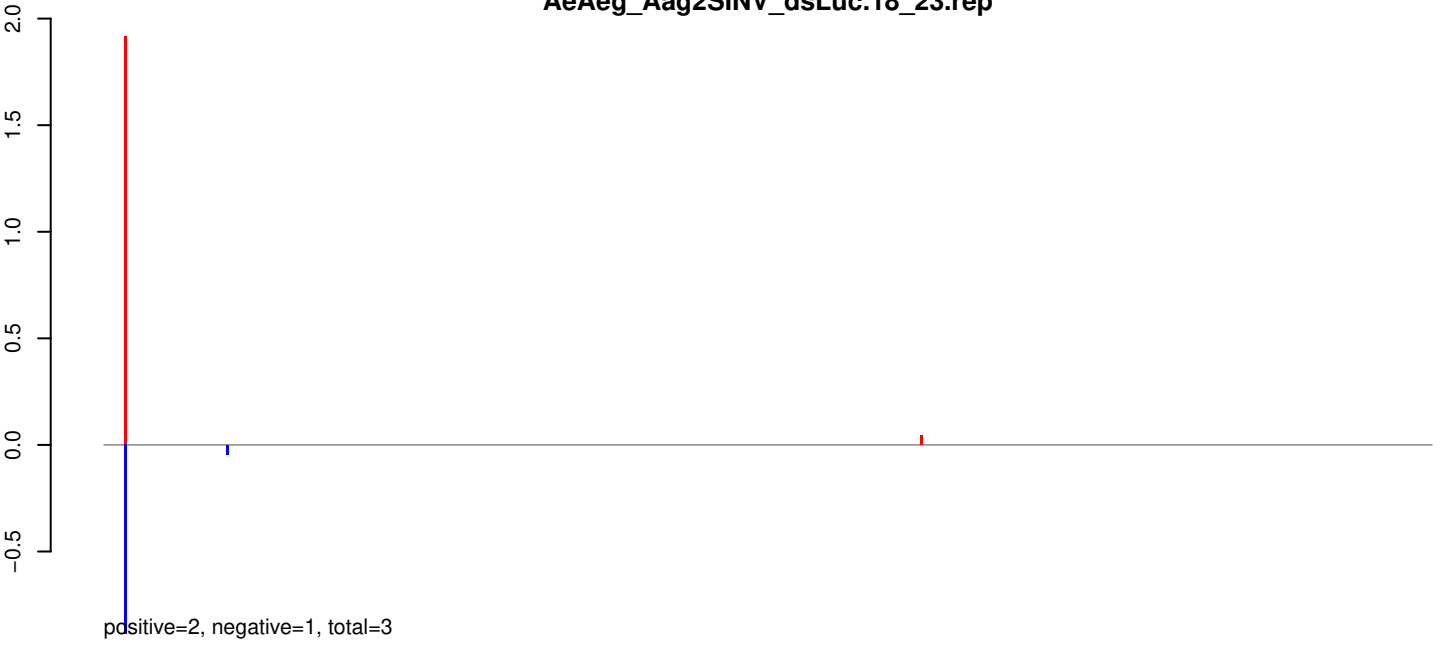
AeAeg_Aag2SINV_dsLuc.rep



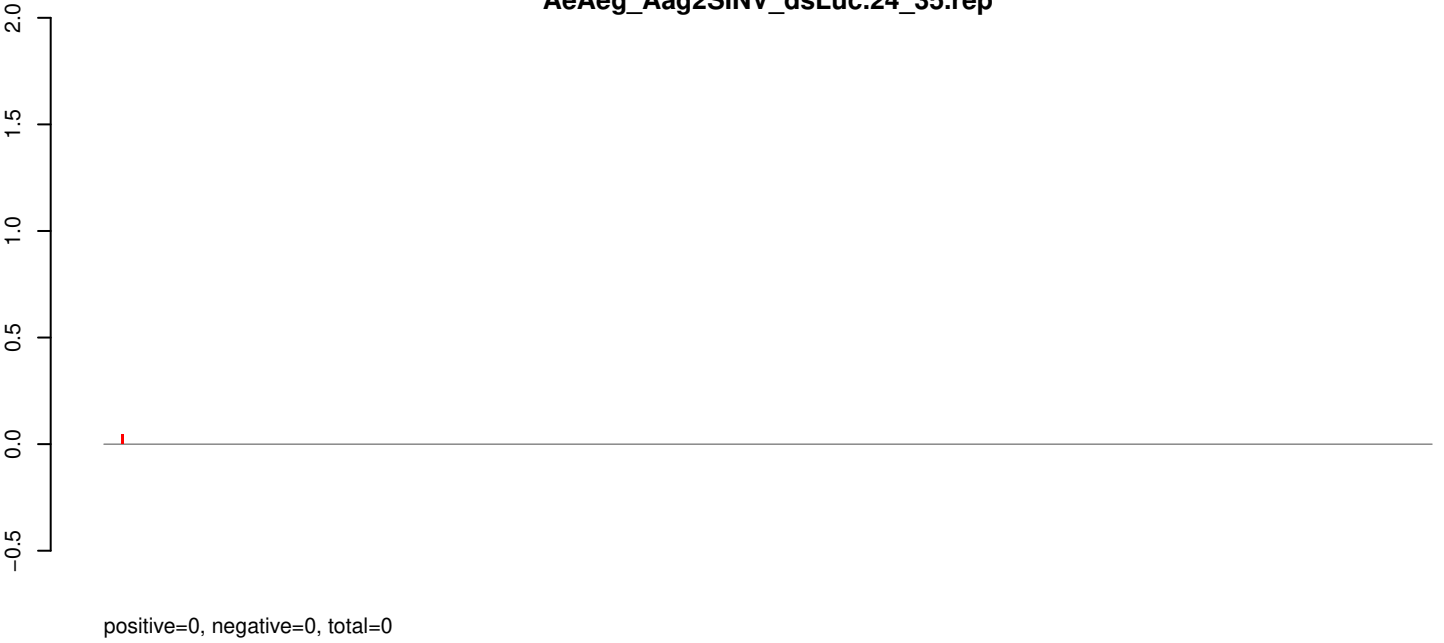
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0 2000 4000 6000 8000 10000

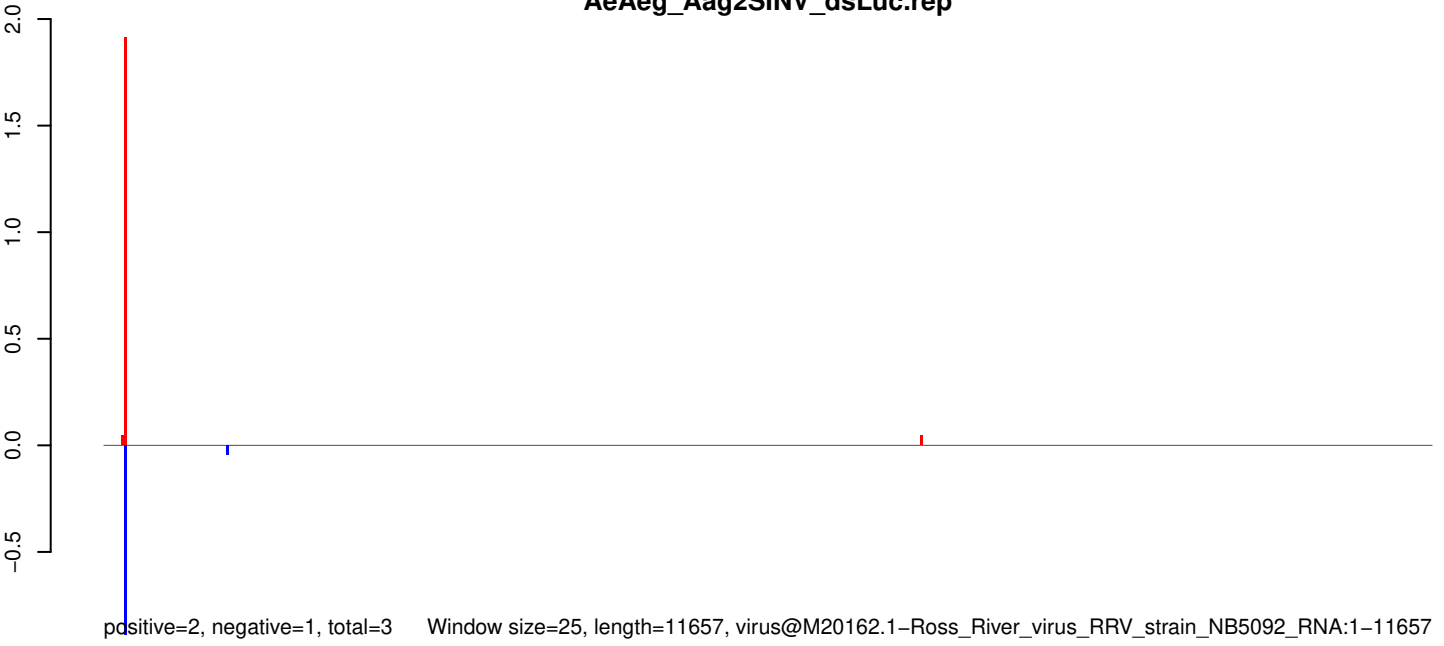
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



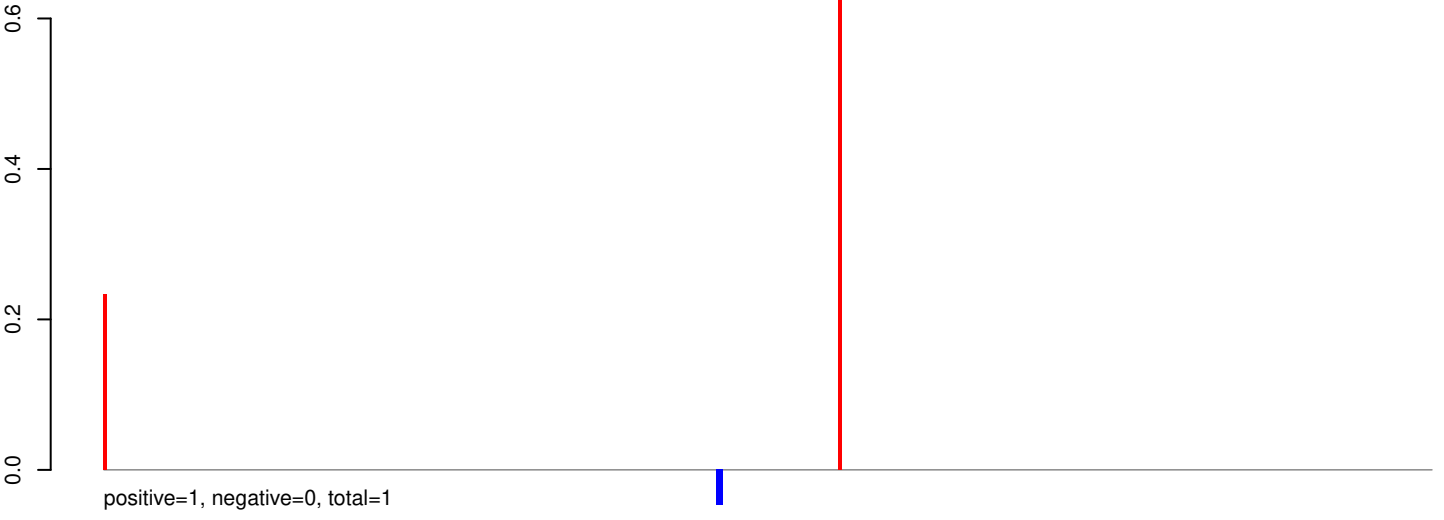
AeAeg_Aag2SINV_dsLuc.rep



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

0 2000 4000 6000 8000 10000 12000

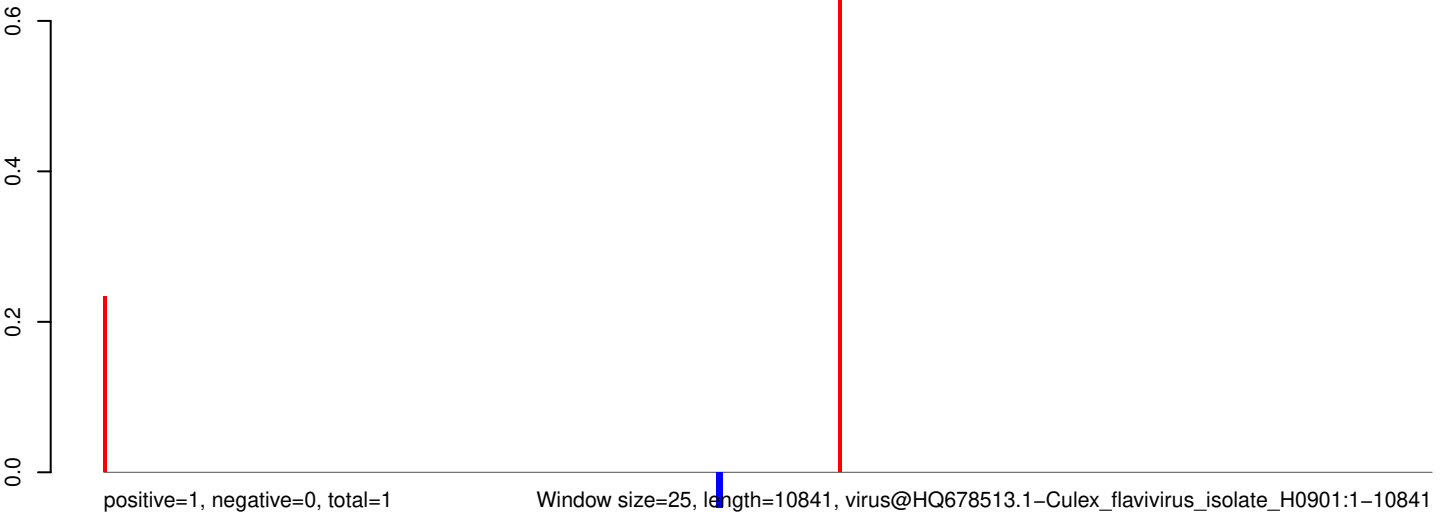
AeAeg_Aag2SINV_dsLuc.18_23.rep



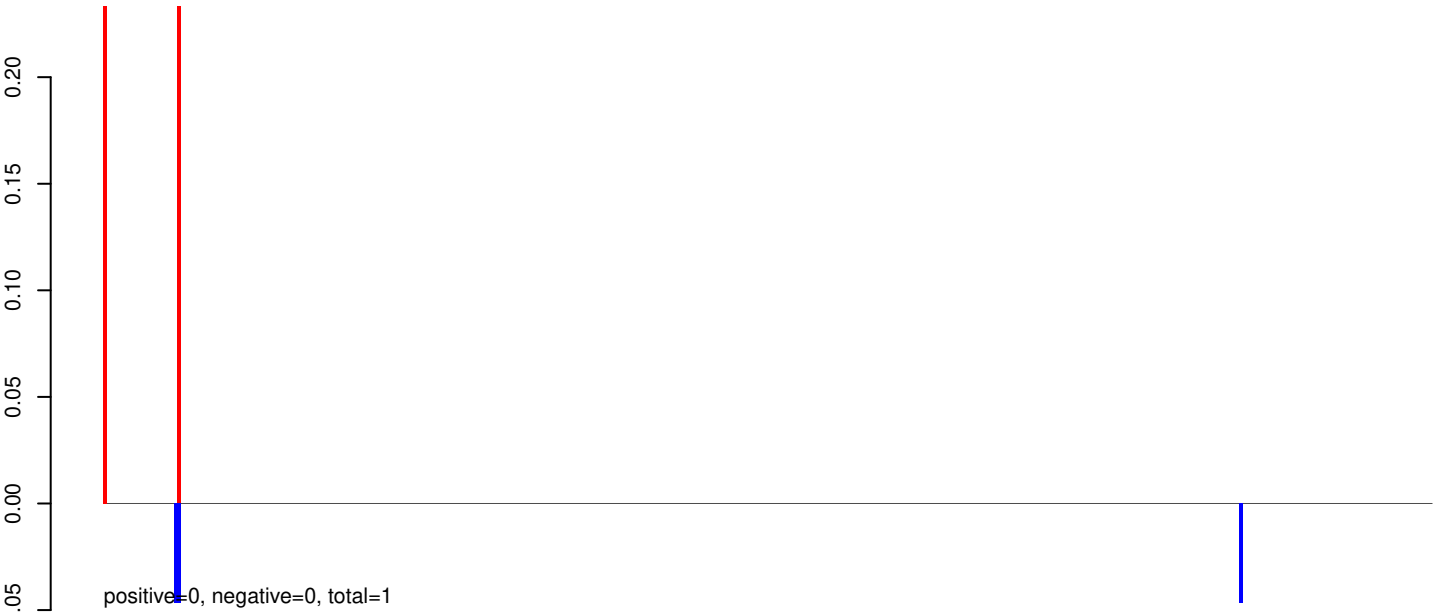
AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



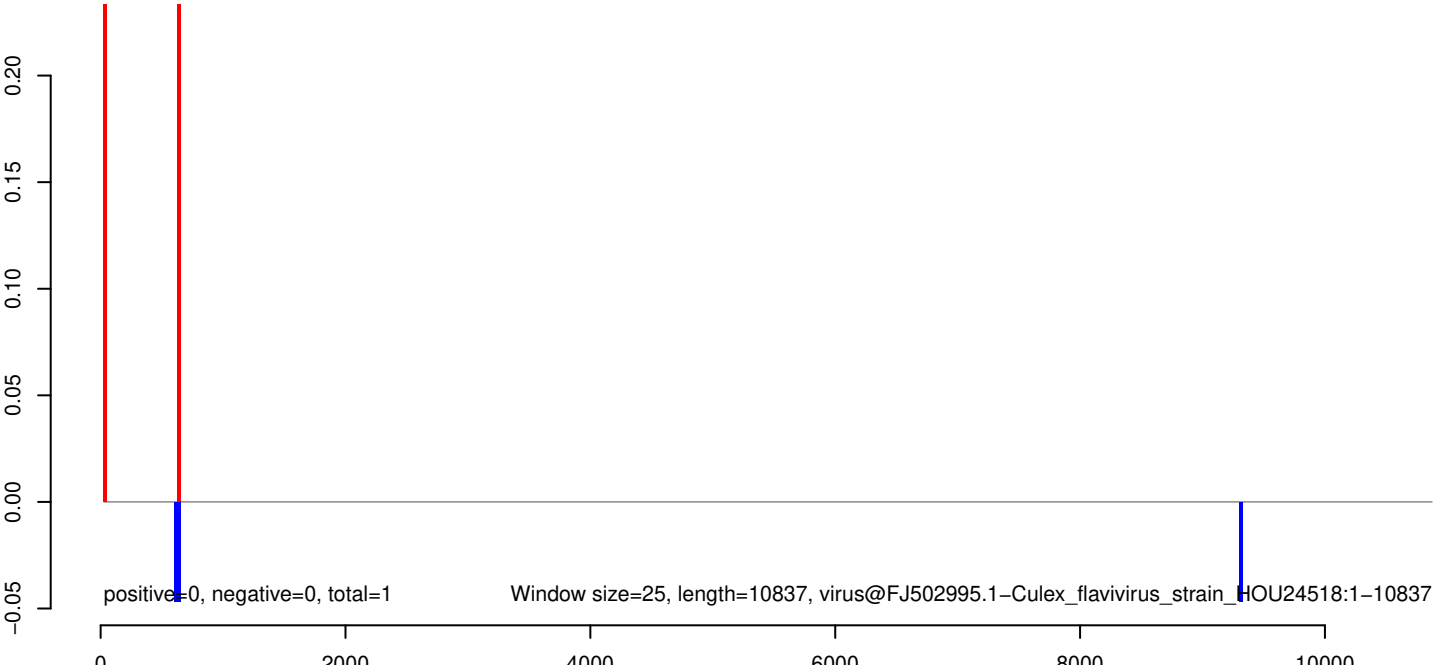
AeAeg_Aag2SINV_dsLuc.18_23.rep



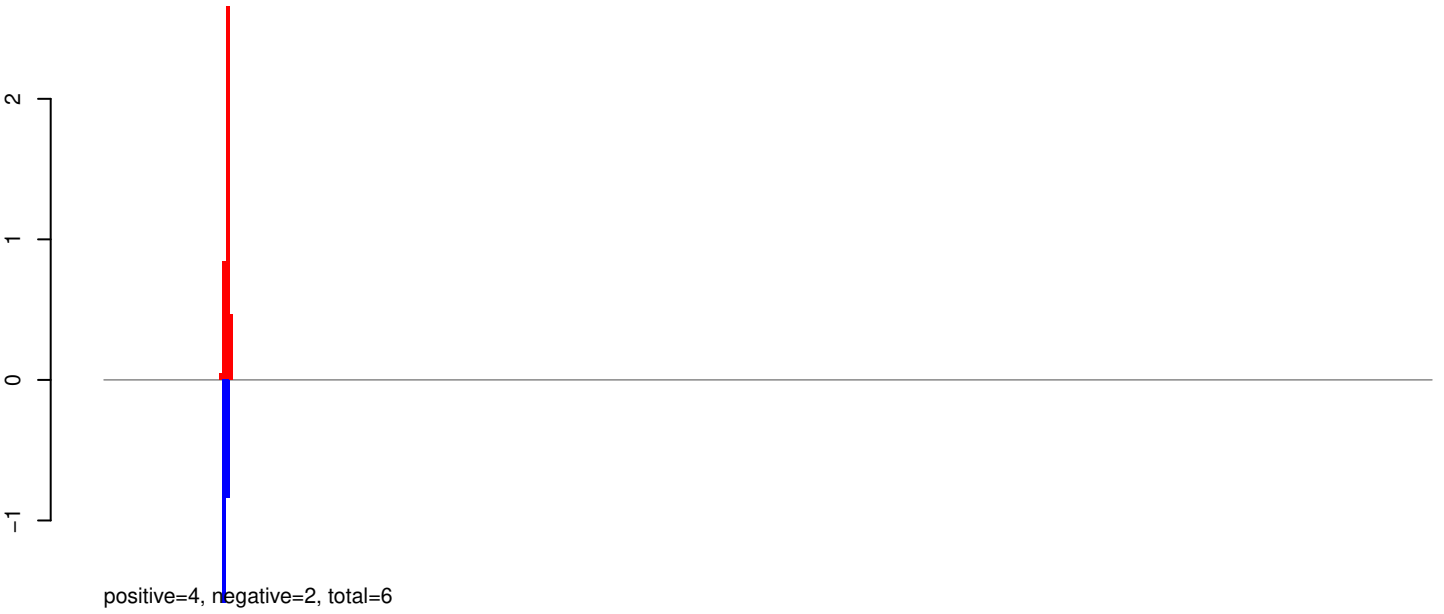
AeAeg_Aag2SINV_dsLuc.24_35.rep



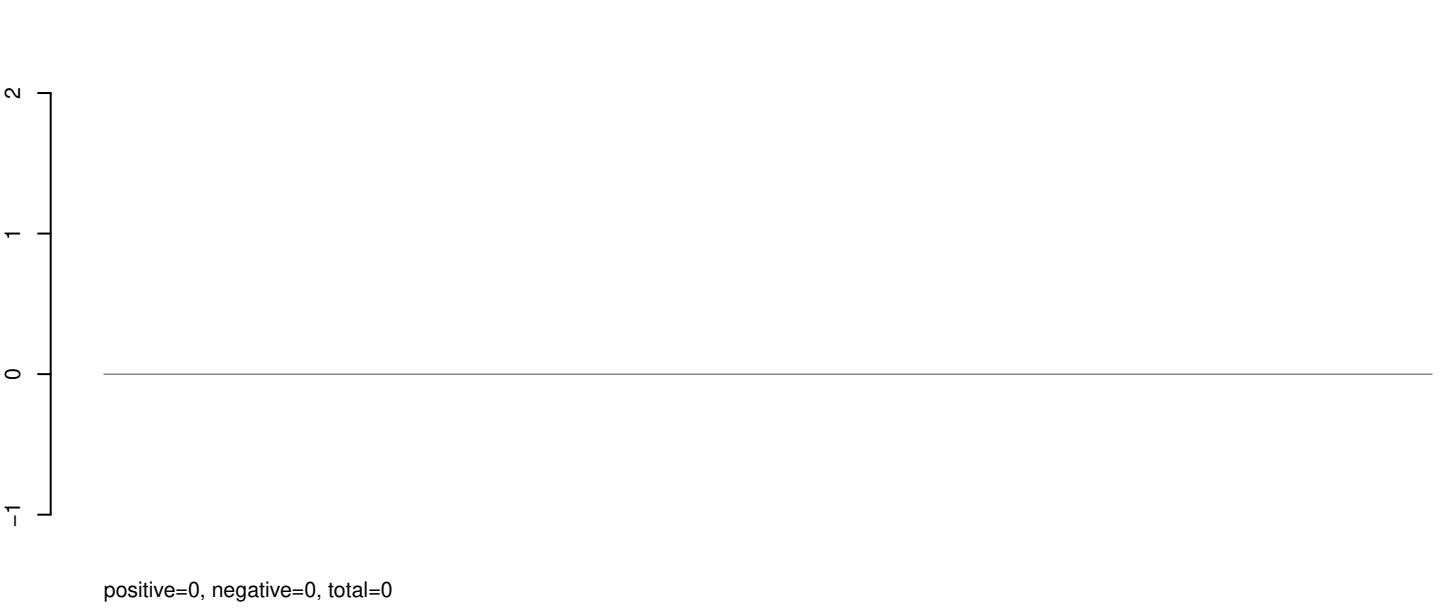
AeAeg_Aag2SINV_dsLuc.rep



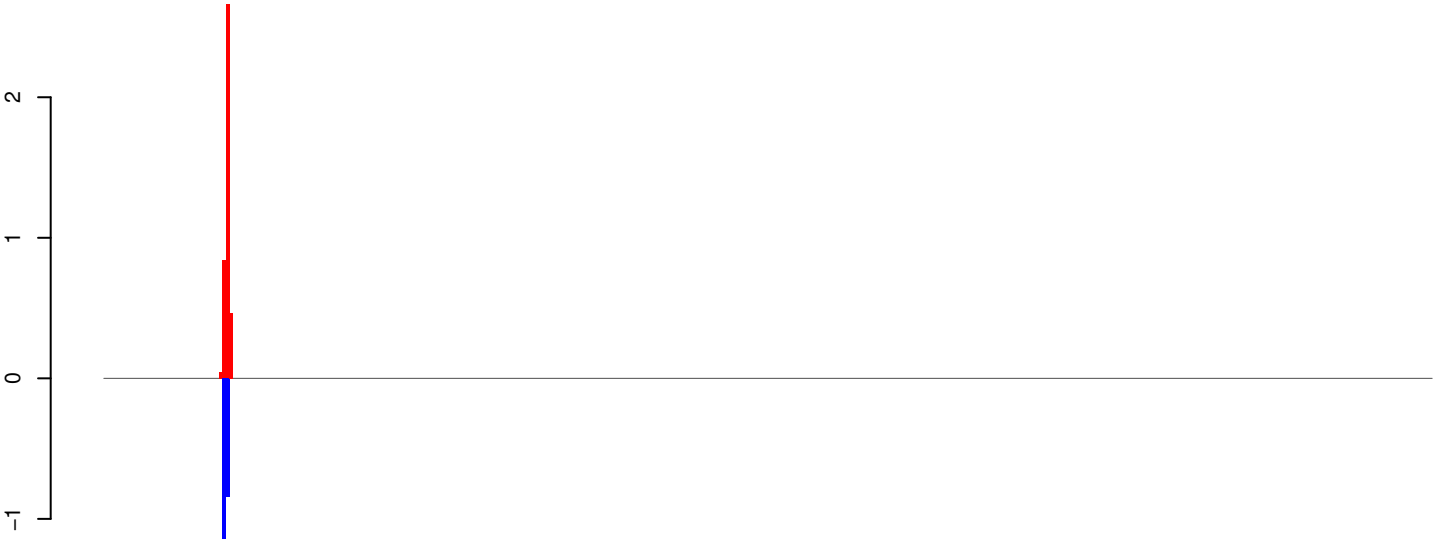
AeAeg_Aag2SINV_dsLuc.18_23.rep



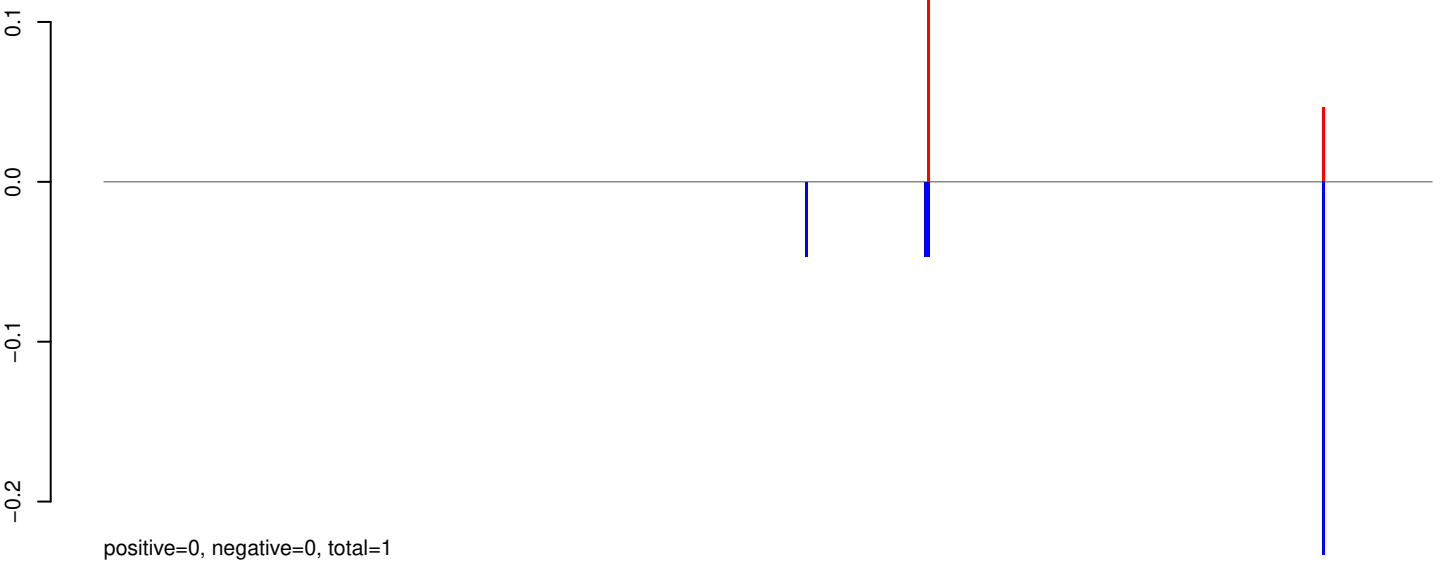
AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



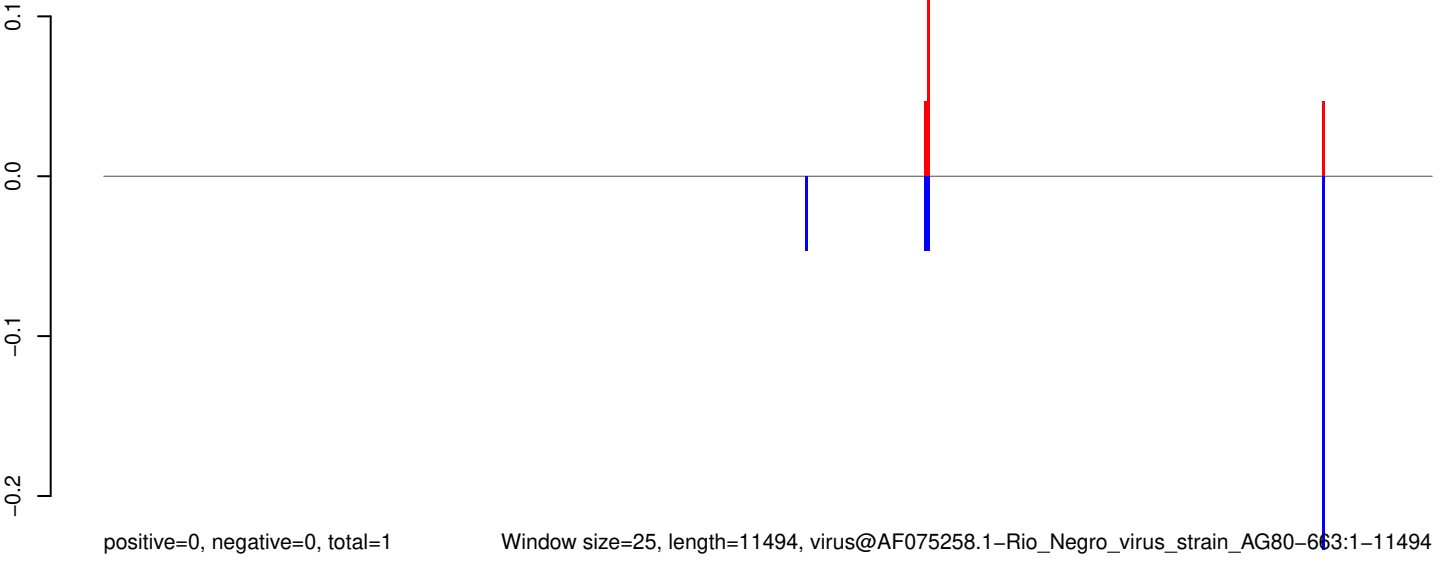
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



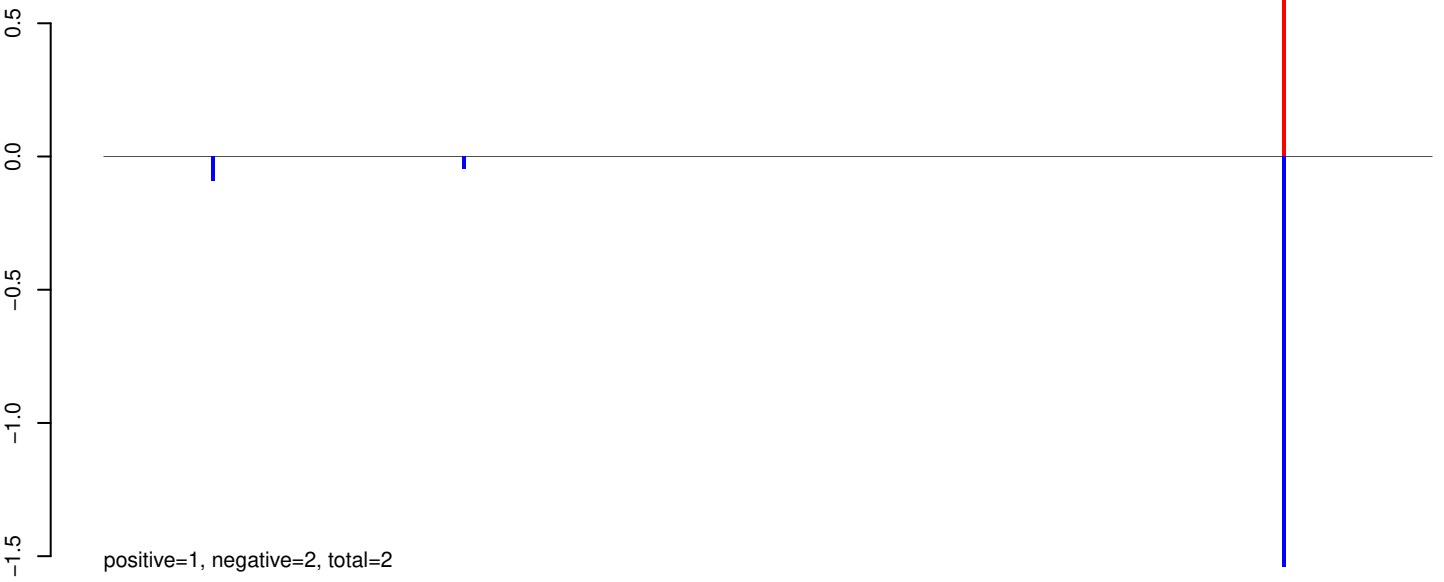
AeAeg_Aag2SINV_dsLuc.rep



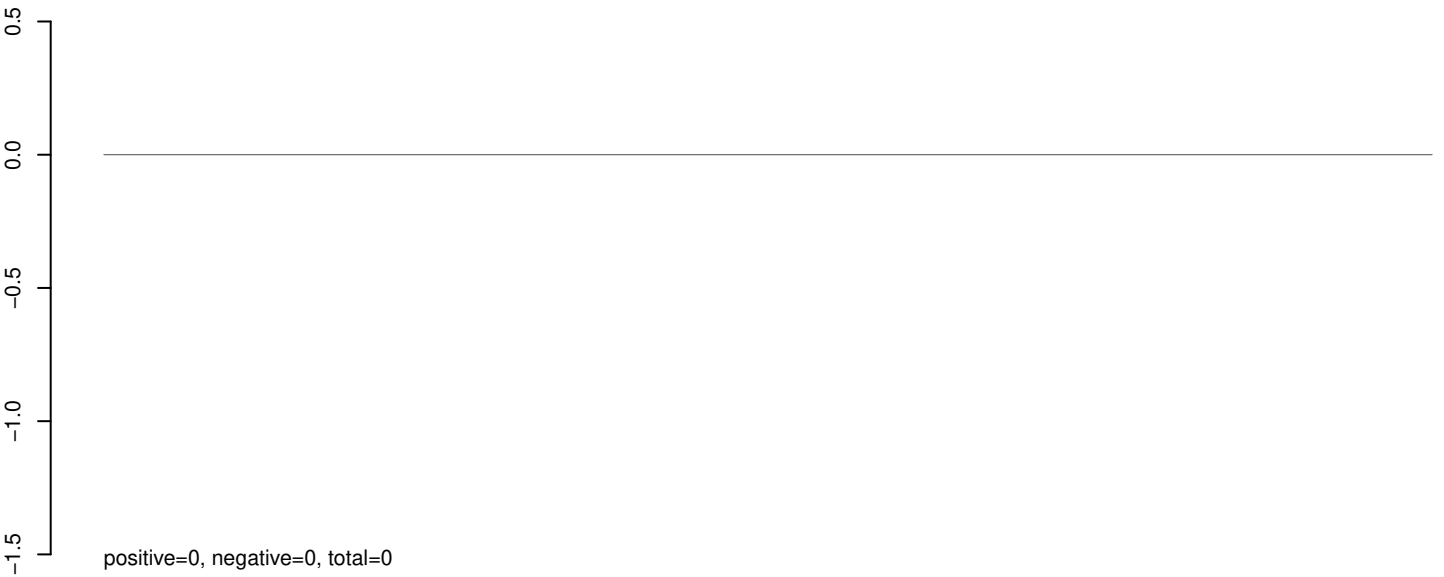
Window size=25, length=11494, virus@AF075258.1-Rio_Negro_virus_strain_AG80-663:1-11494

0 2000 4000 6000 8000 10000

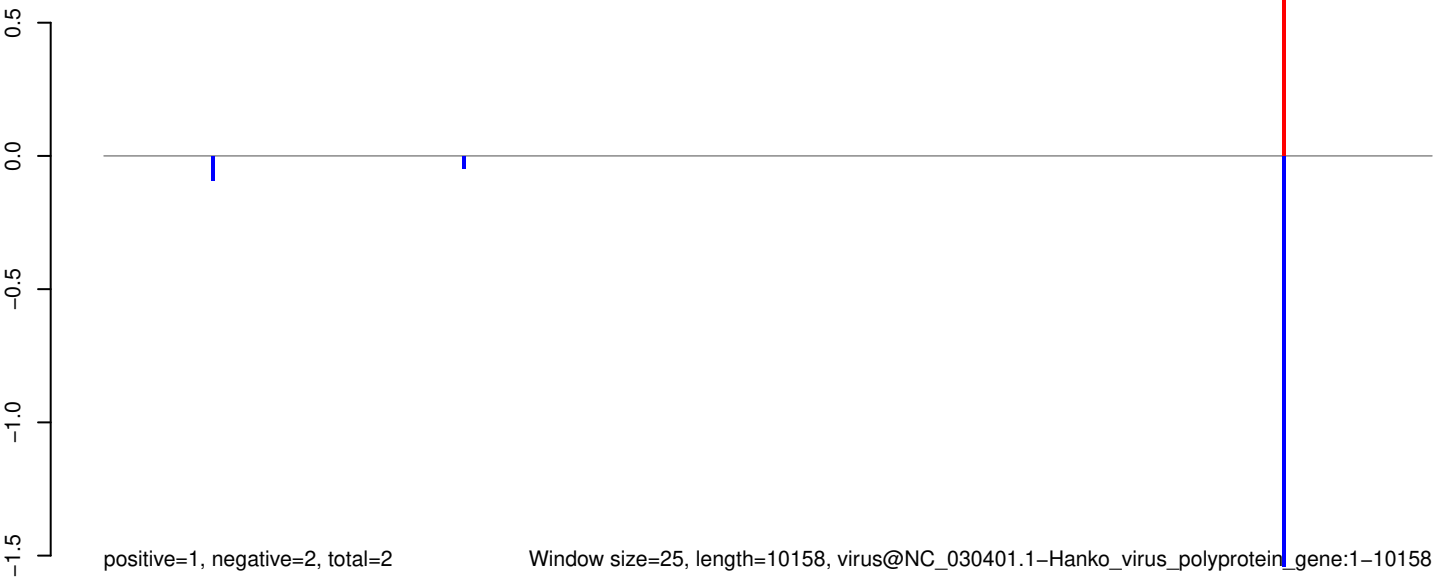
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



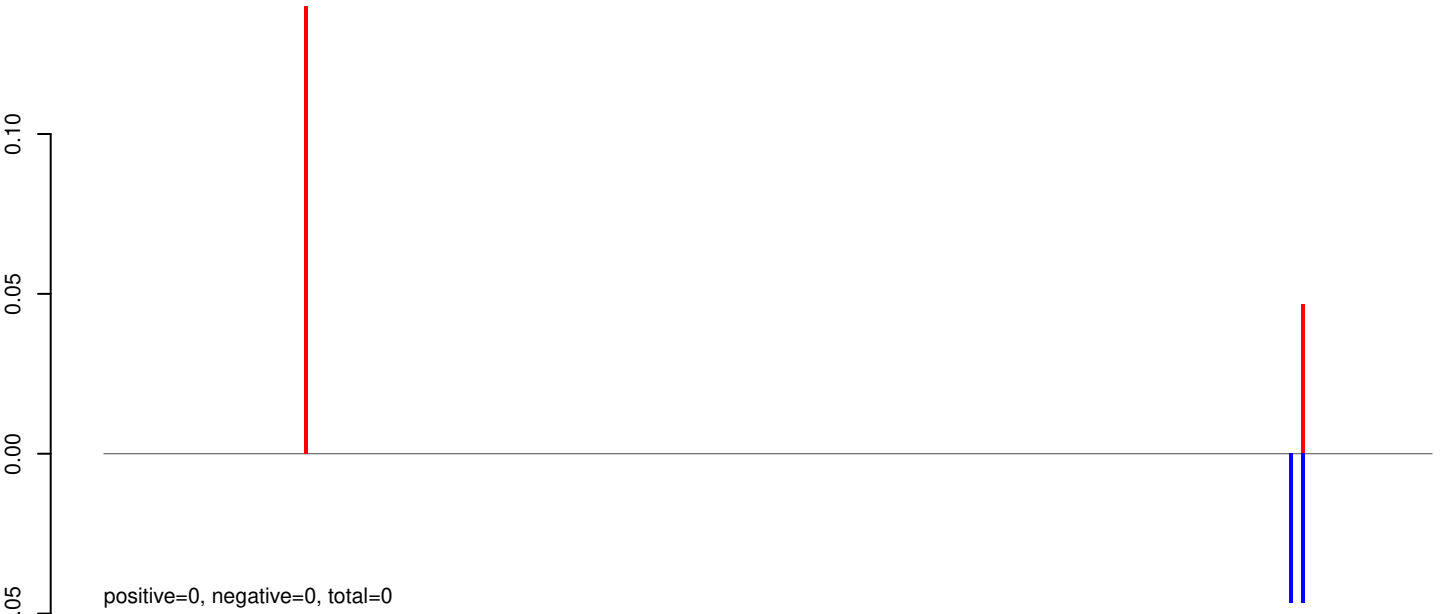
AeAeg_Aag2SINV_dsLuc.rep



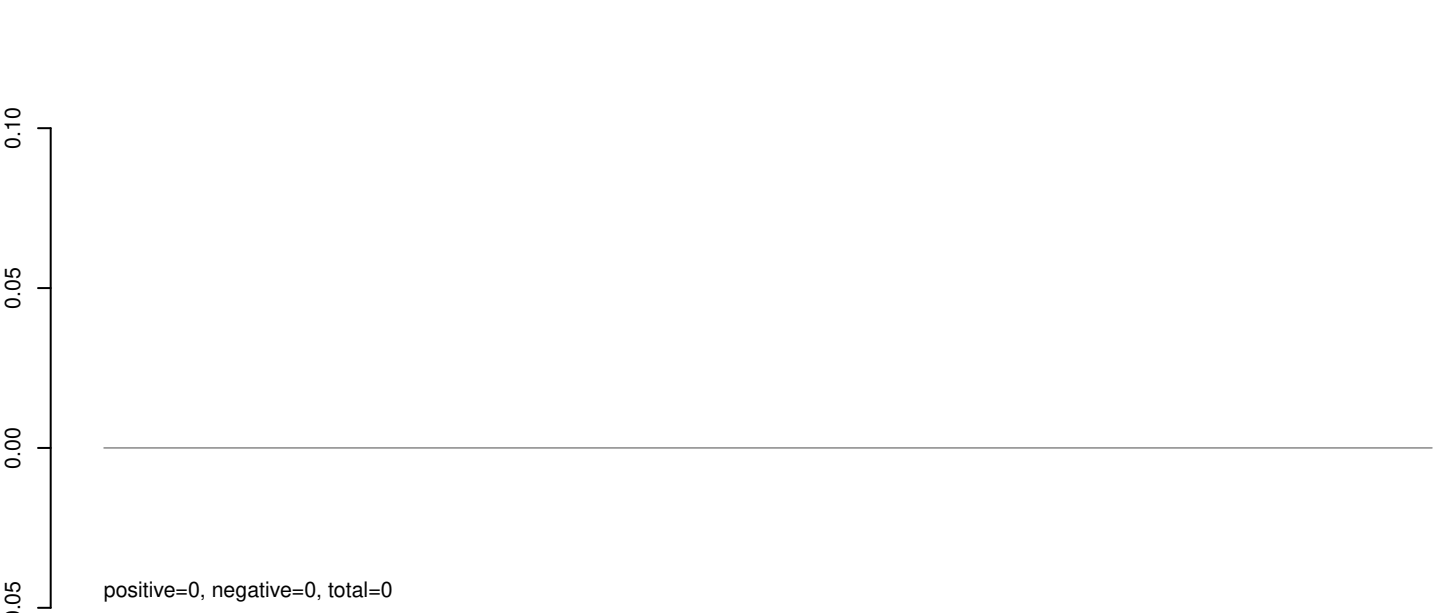
Window size=25, length=10158, virus@NC_030401.1-Hanko_virus_polyprotein_gene:1-10158

0 2000 4000 6000 8000 10000

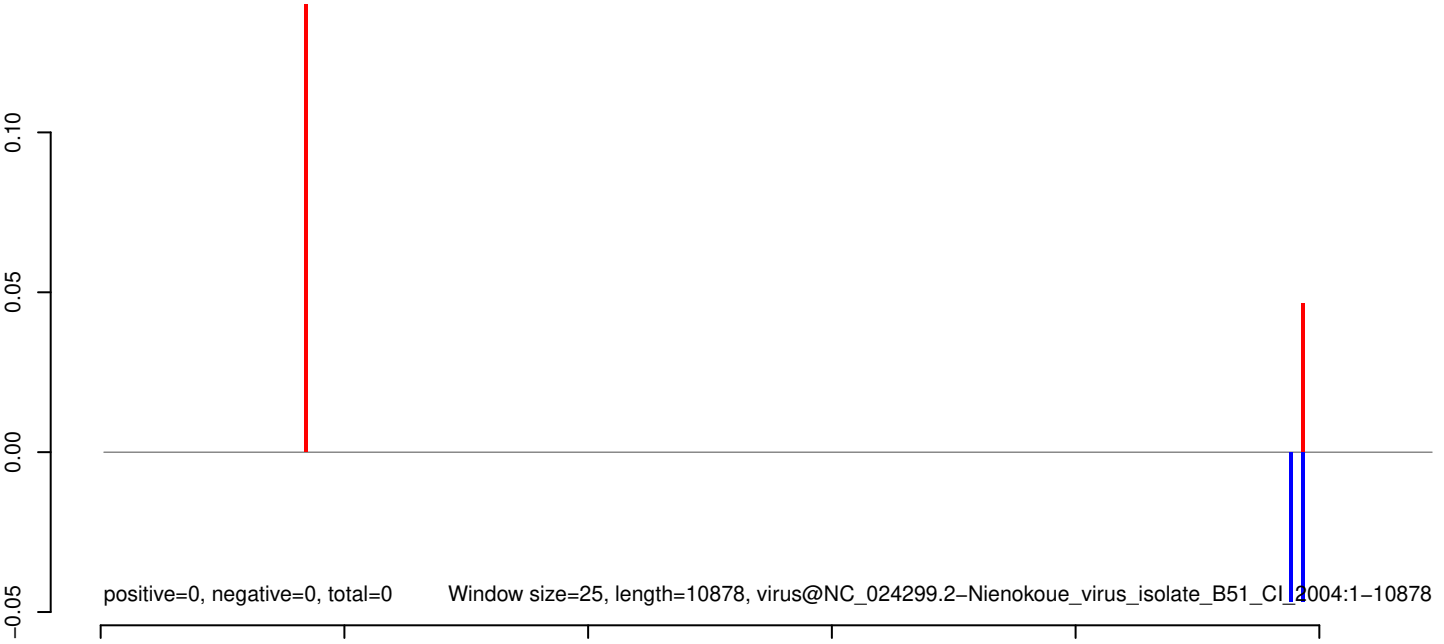
AeAeg_Aag2SINV_dsLuc.18_23.rep



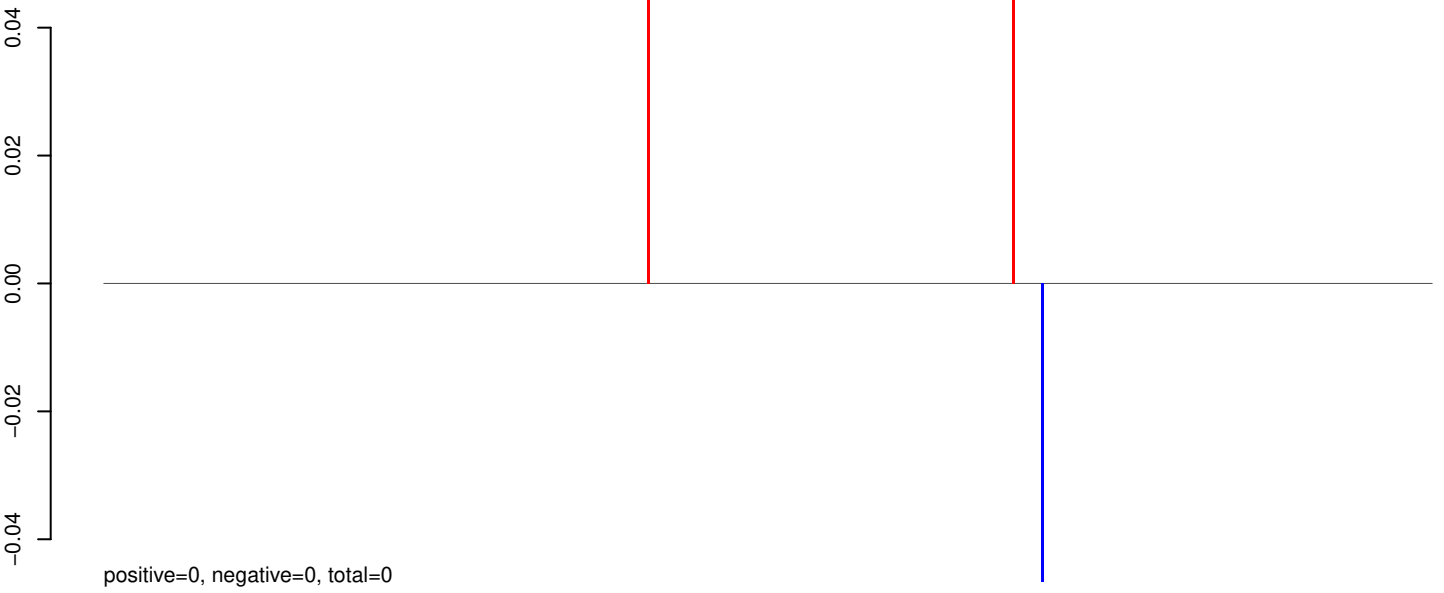
AeAeg_Aag2SINV_dsLuc.24_35.rep



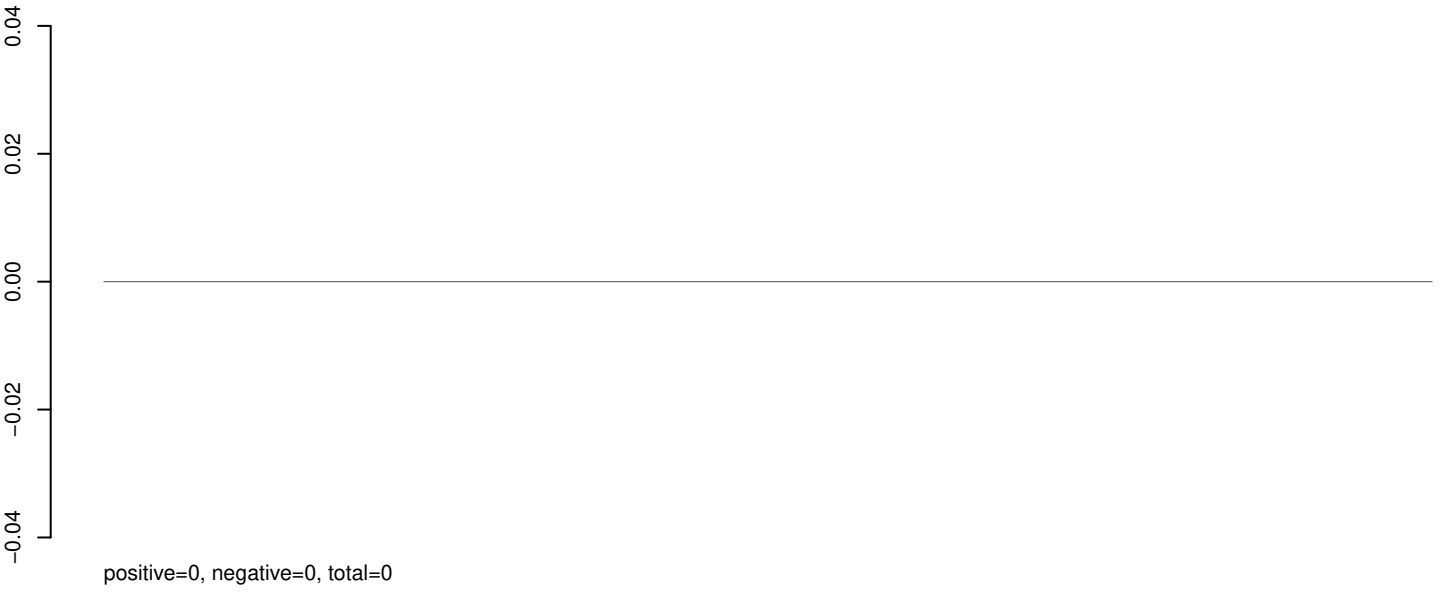
AeAeg_Aag2SINV_dsLuc.rep



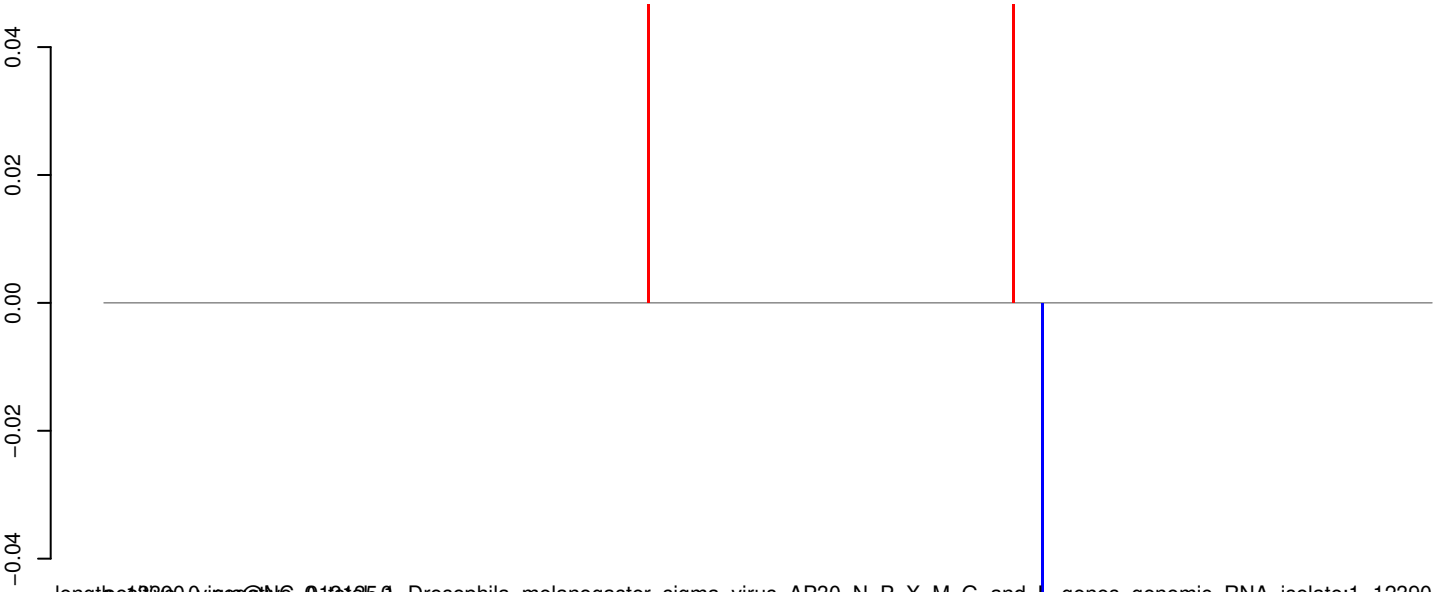
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



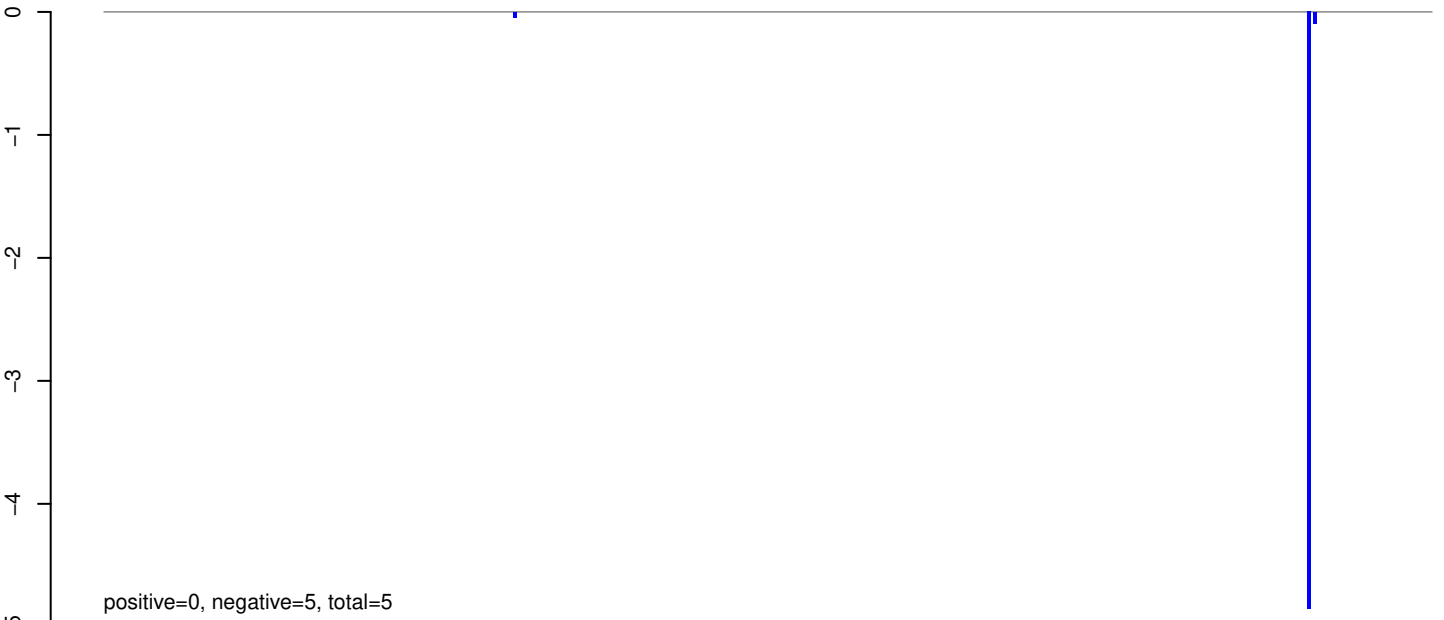
AeAeg_Aag2SINV_dsLuc.rep



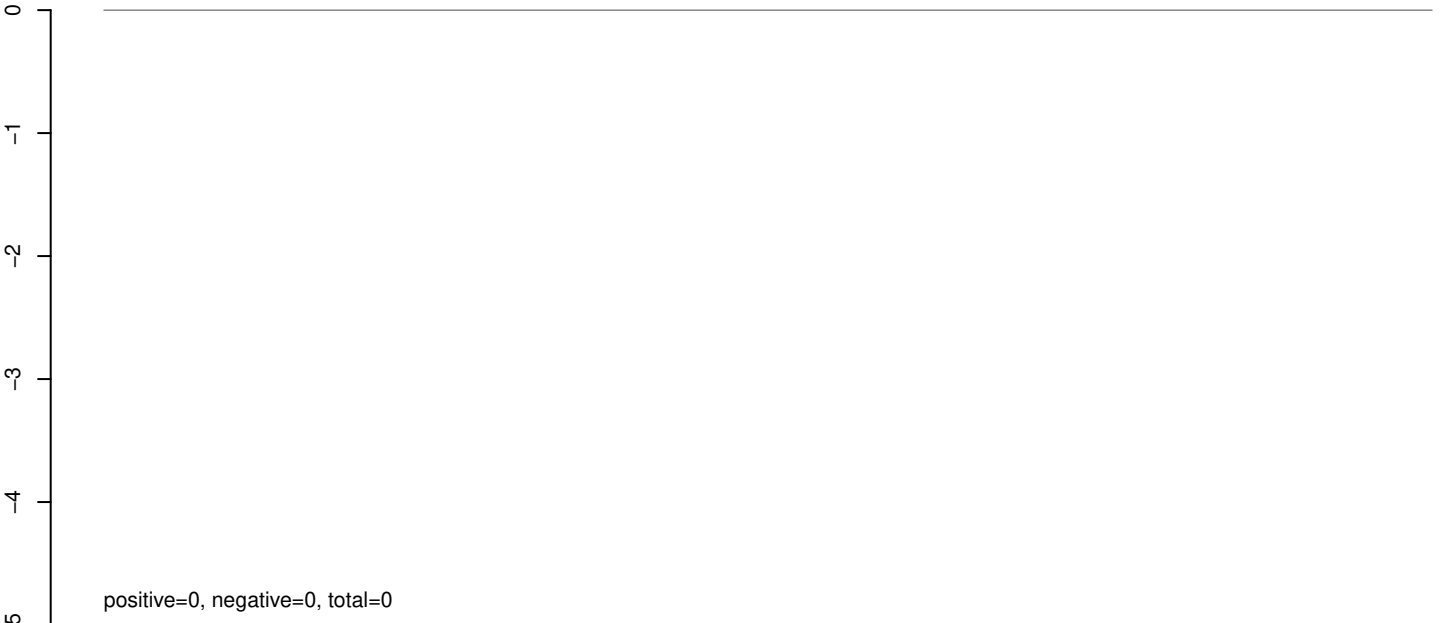
length=12390, virus=161350 -Drosophila_melanogaster_sigma_virus_AP30_N_P_X_M_G_and_L_genes_genomic_RNA_isolate:1-12390

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



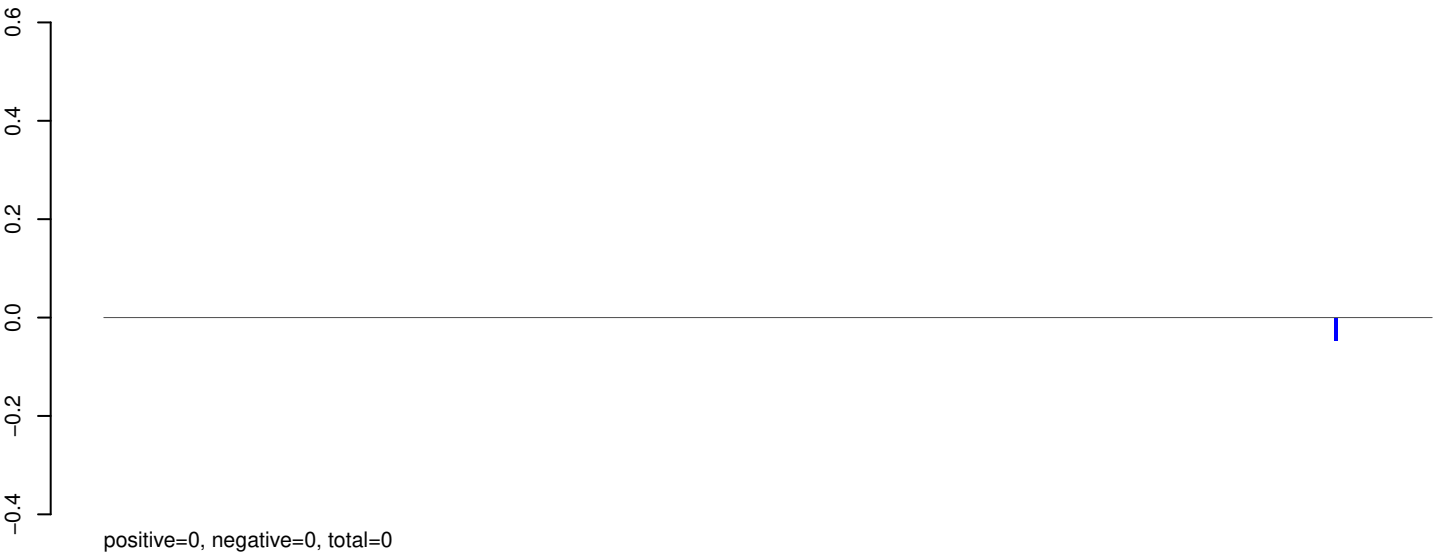
Window size=25, length=10874, virus@NC_009029.2-Kokobera_virus:1-10874

0 2000 4000 6000 8000 10000

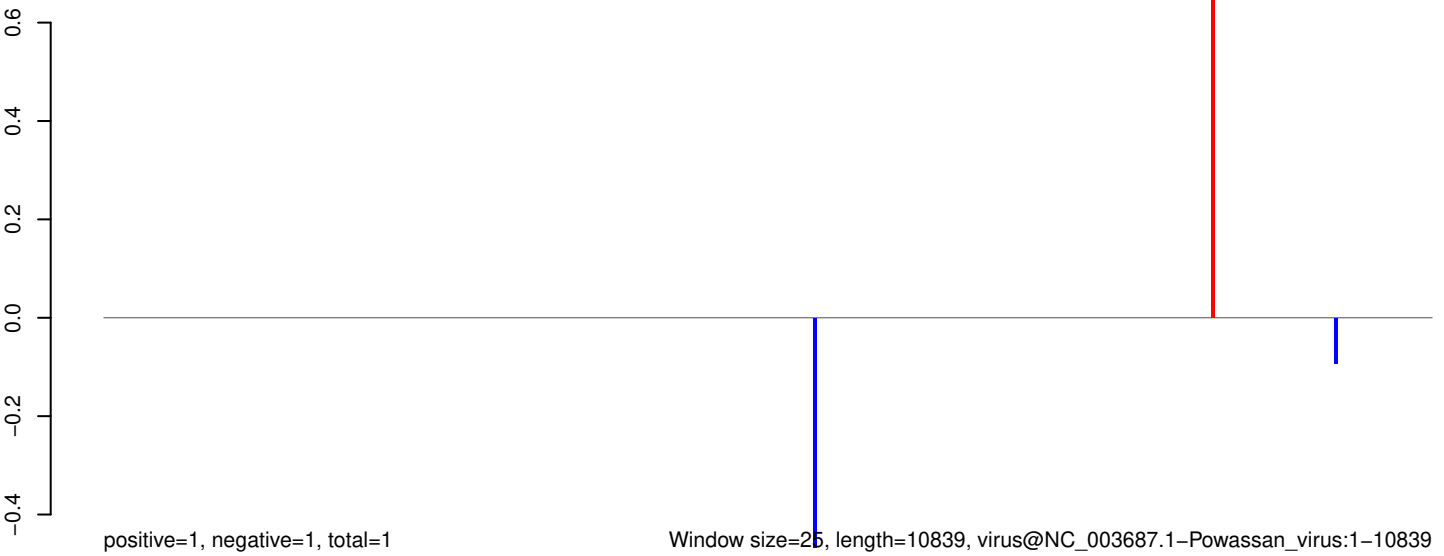
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



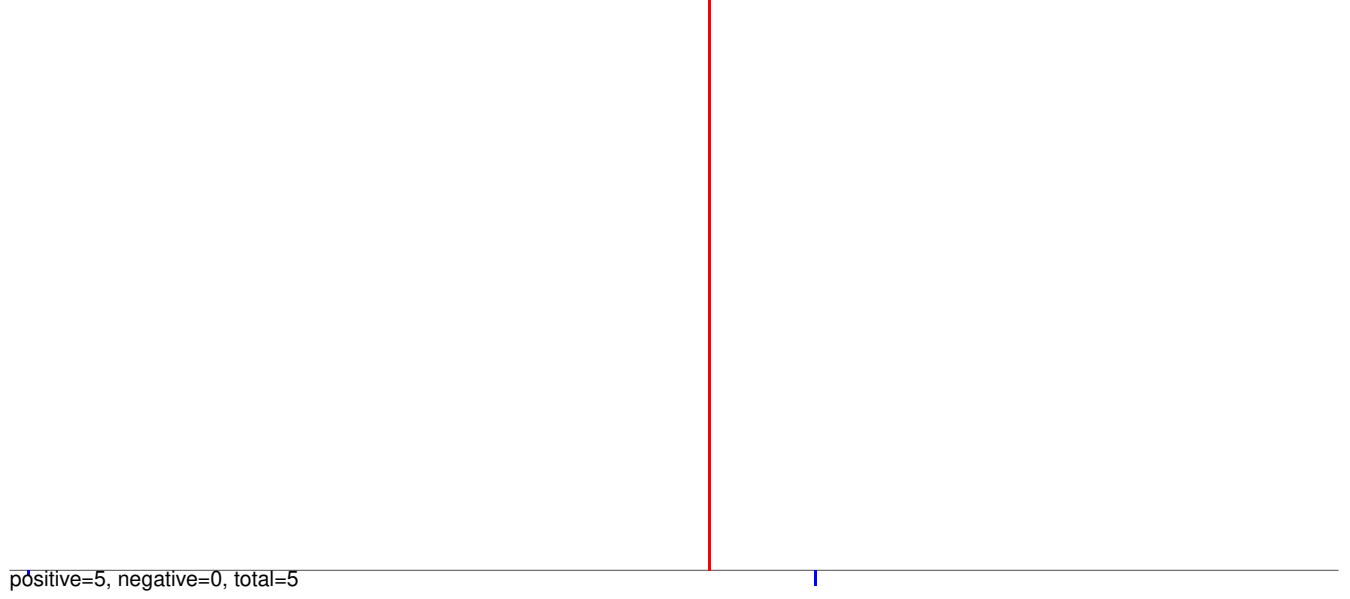
AeAeg_Aag2SINV_dsLuc.rep



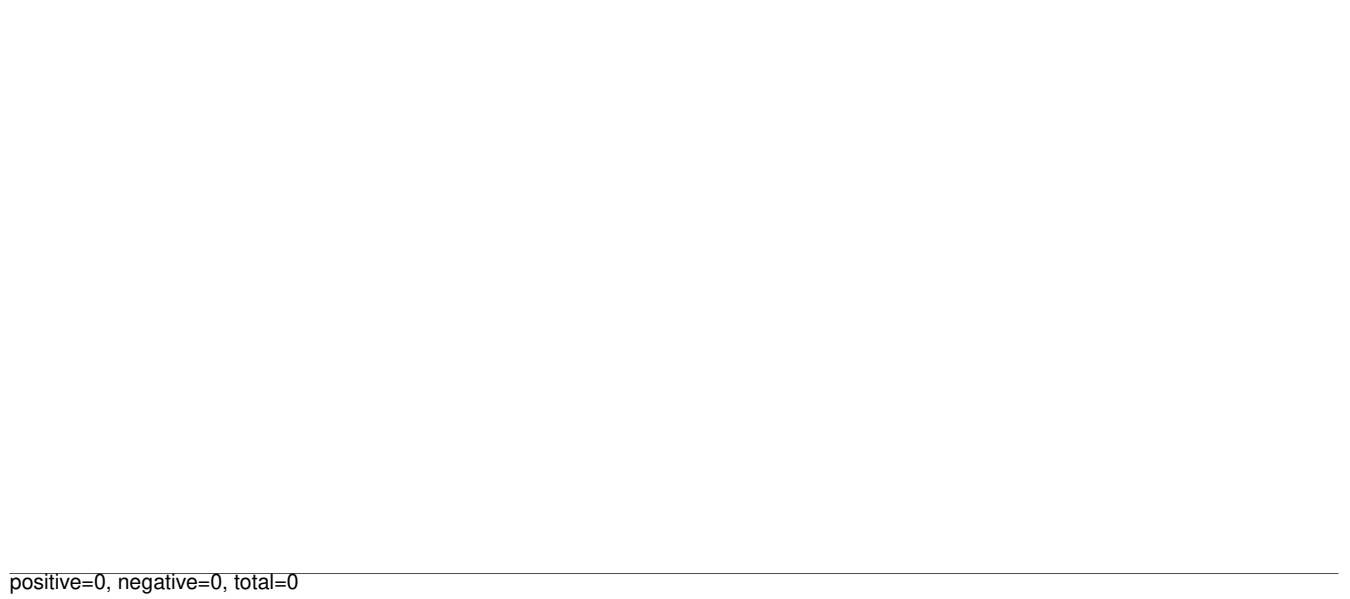
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0 2000 4000 6000 8000 10000

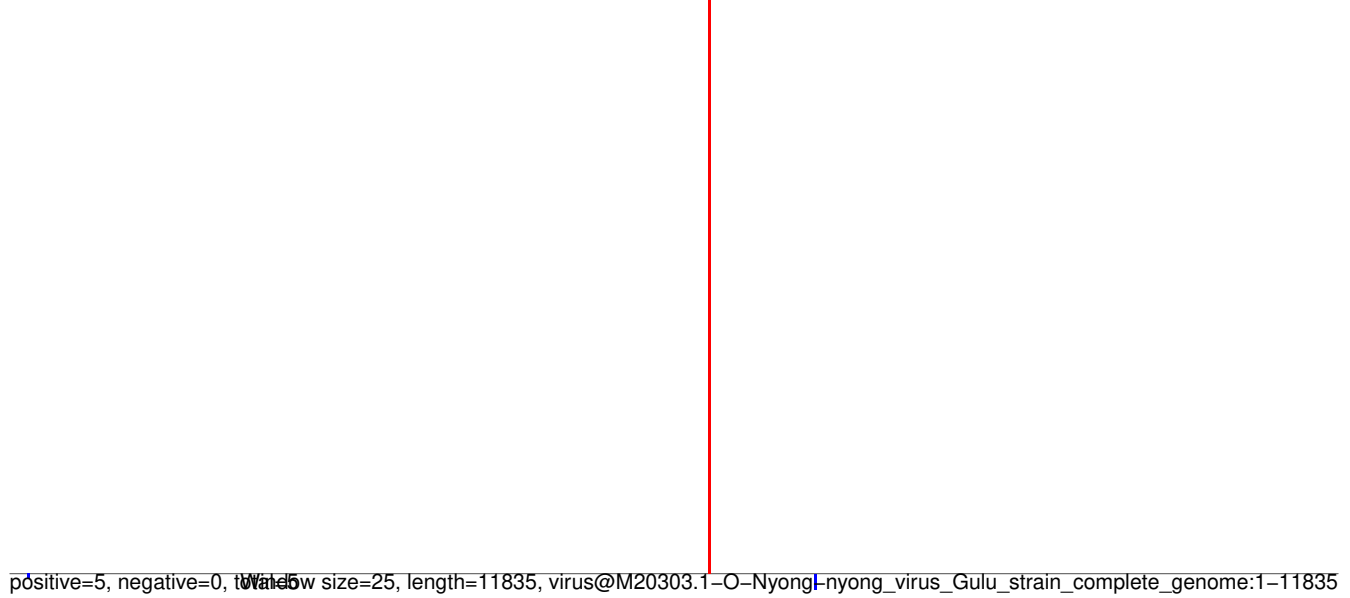
AeAeg_Aag2SINV_dsLuc.18_23.rep



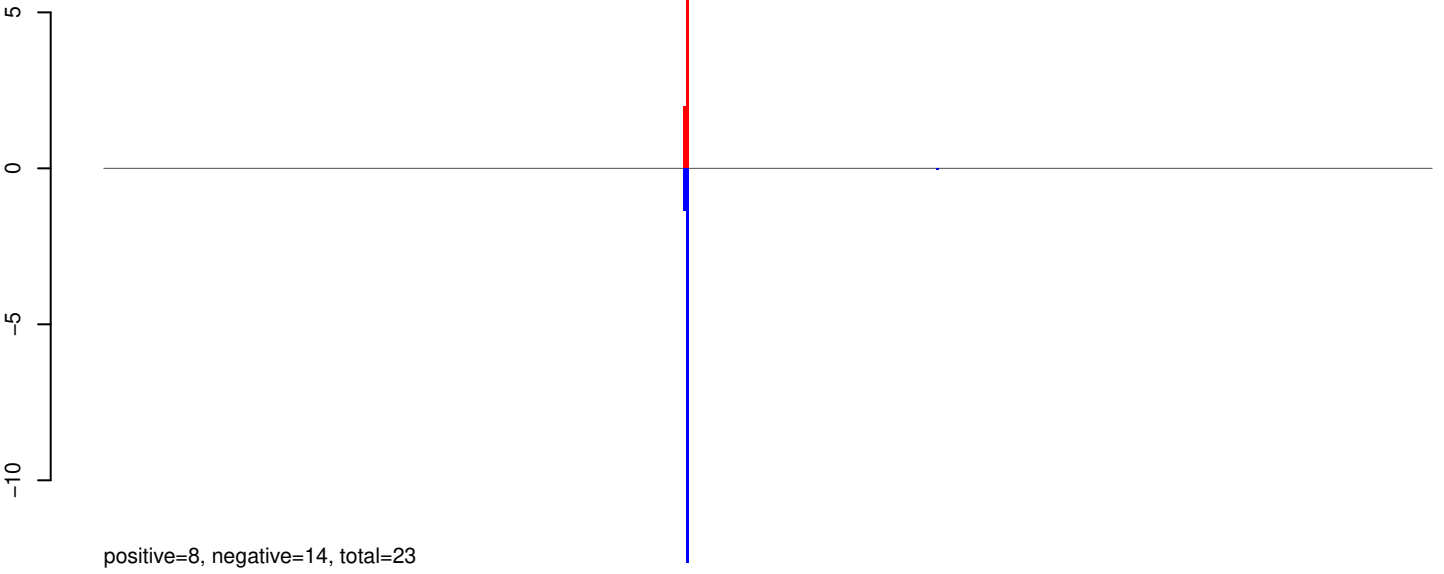
AeAeg_Aag2SINV_dsLuc.24_35.rep



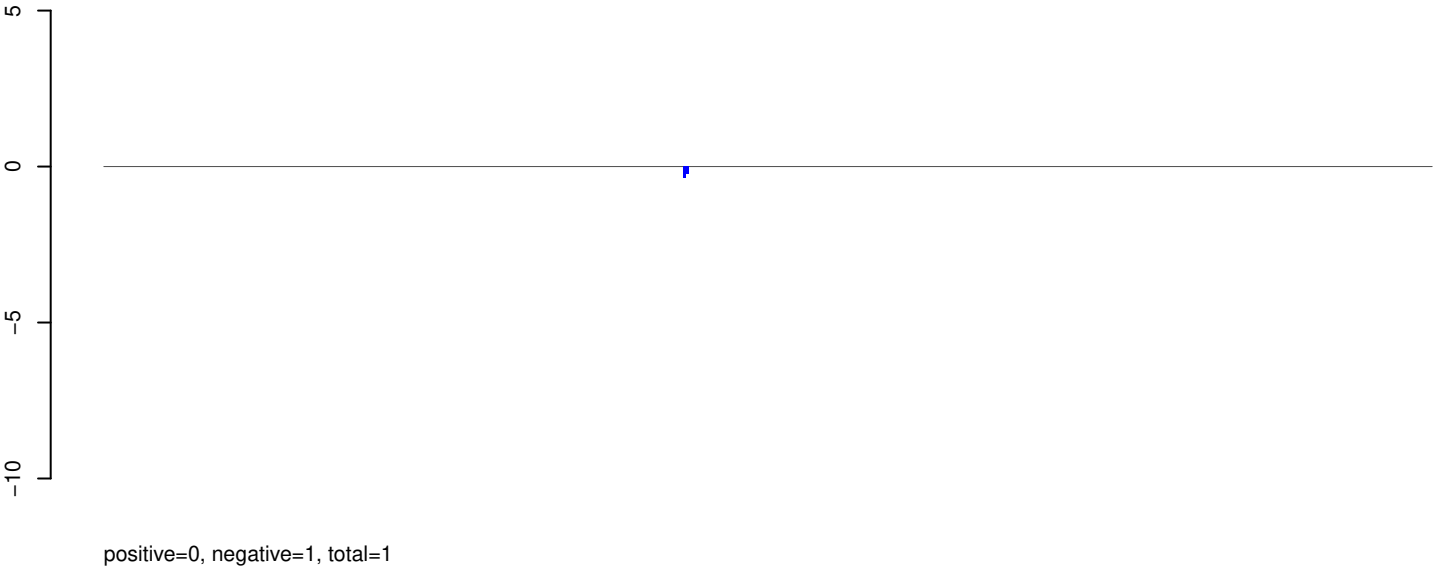
AeAeg_Aag2SINV_dsLuc.rep



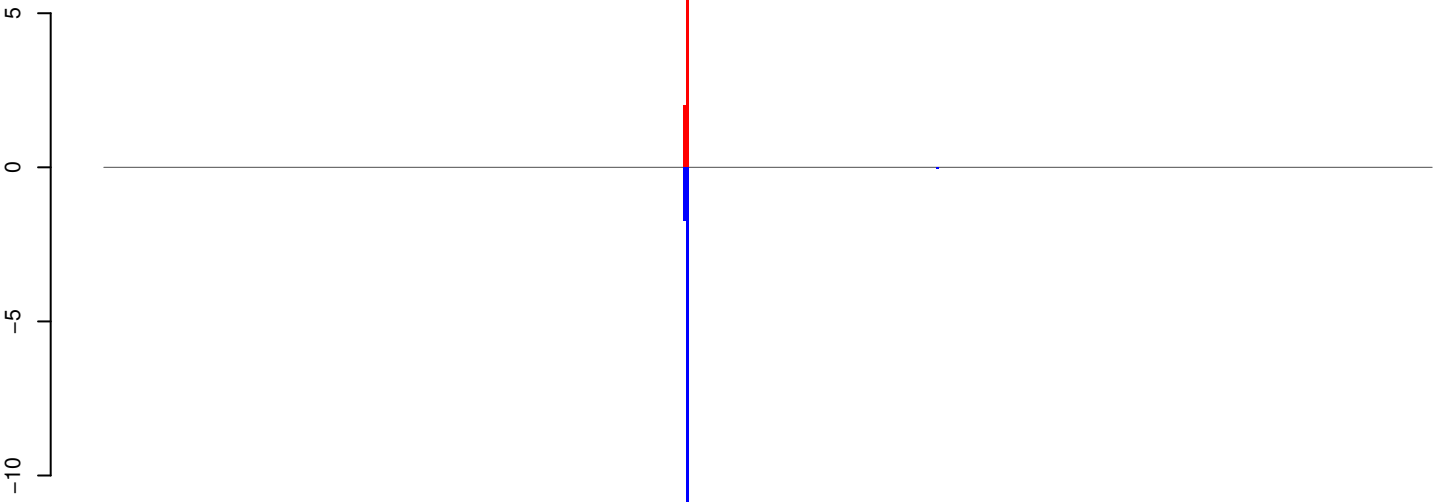
AeAeg_Aag2SINV_dsLuc.18_23.rep



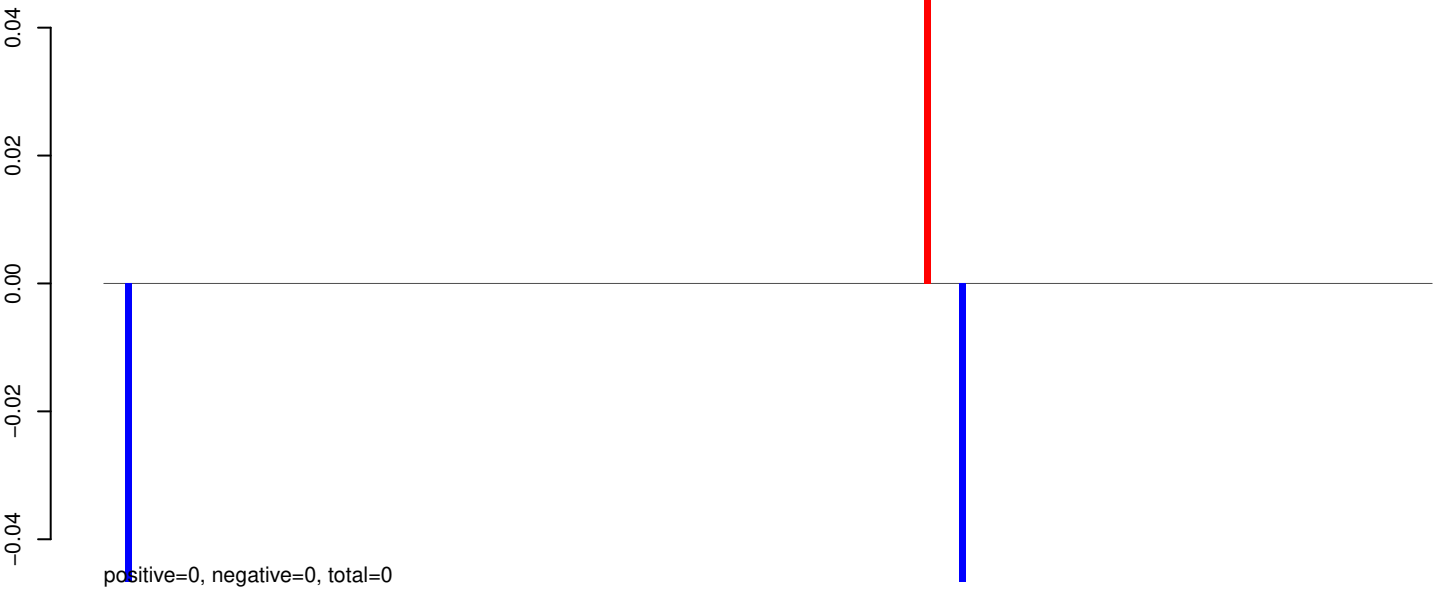
AeAeg_Aag2SINV_dsLuc.24_35.rep



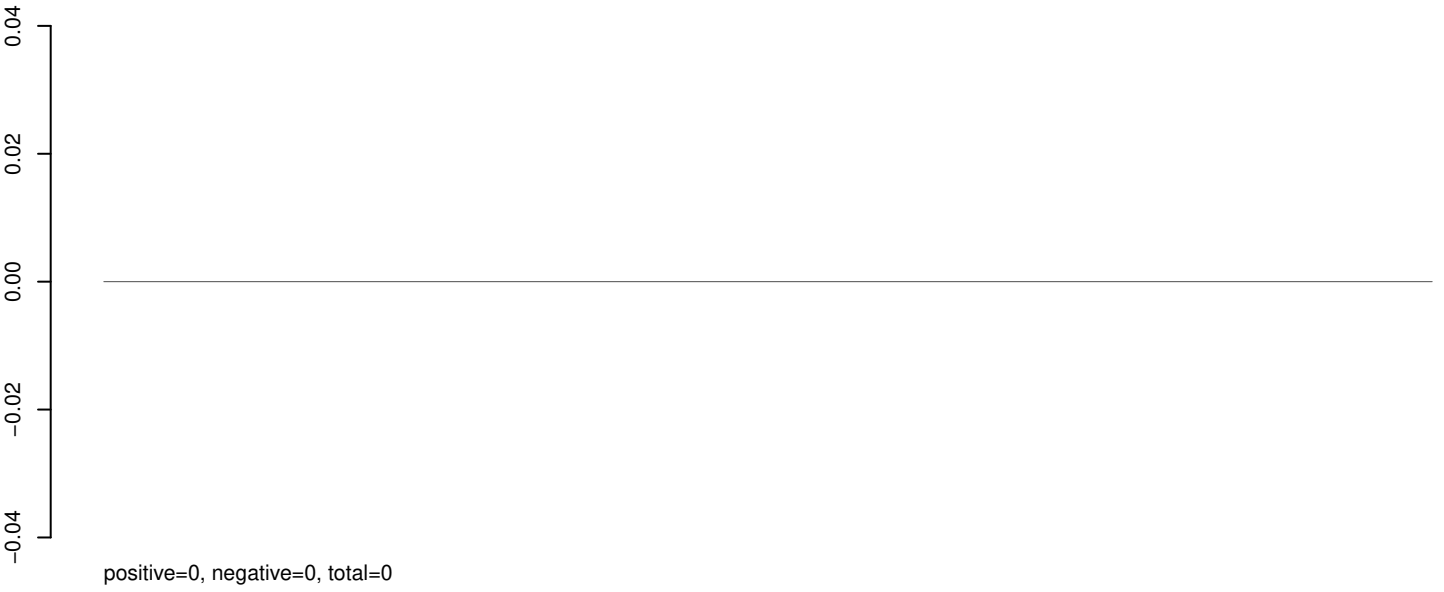
AeAeg_Aag2SINV_dsLuc.rep



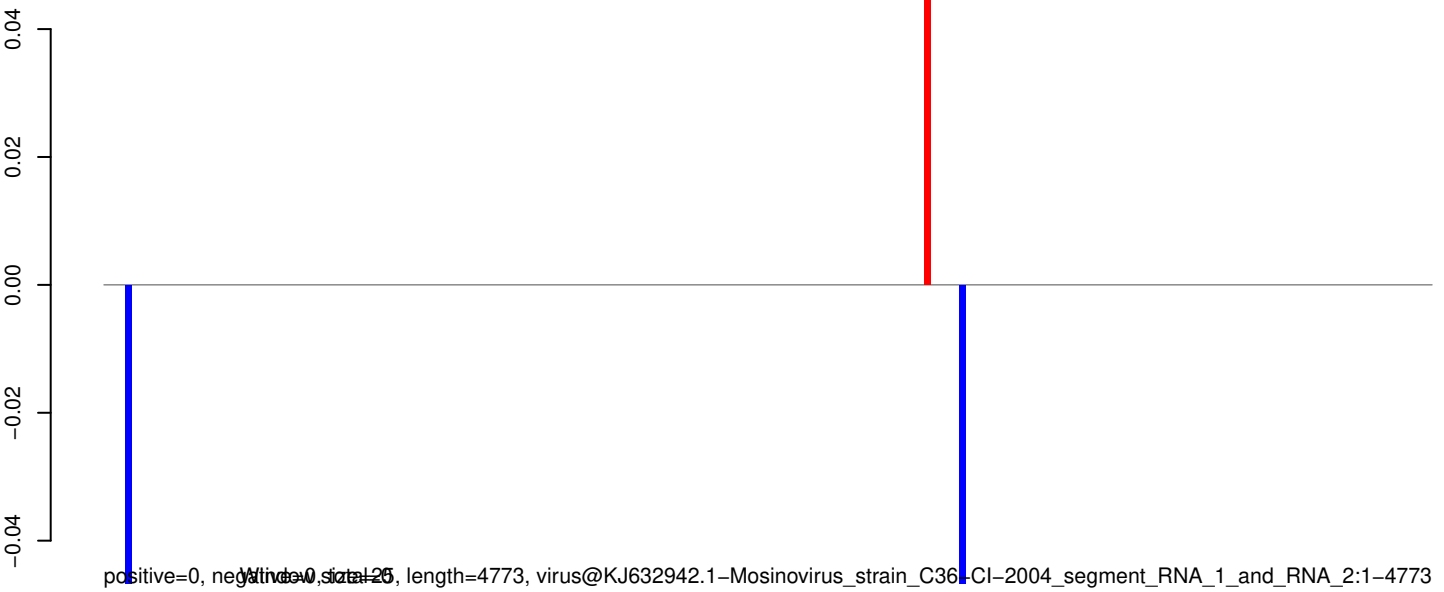
AeAeg_Aag2SINV_dsLuc.18_23.rep



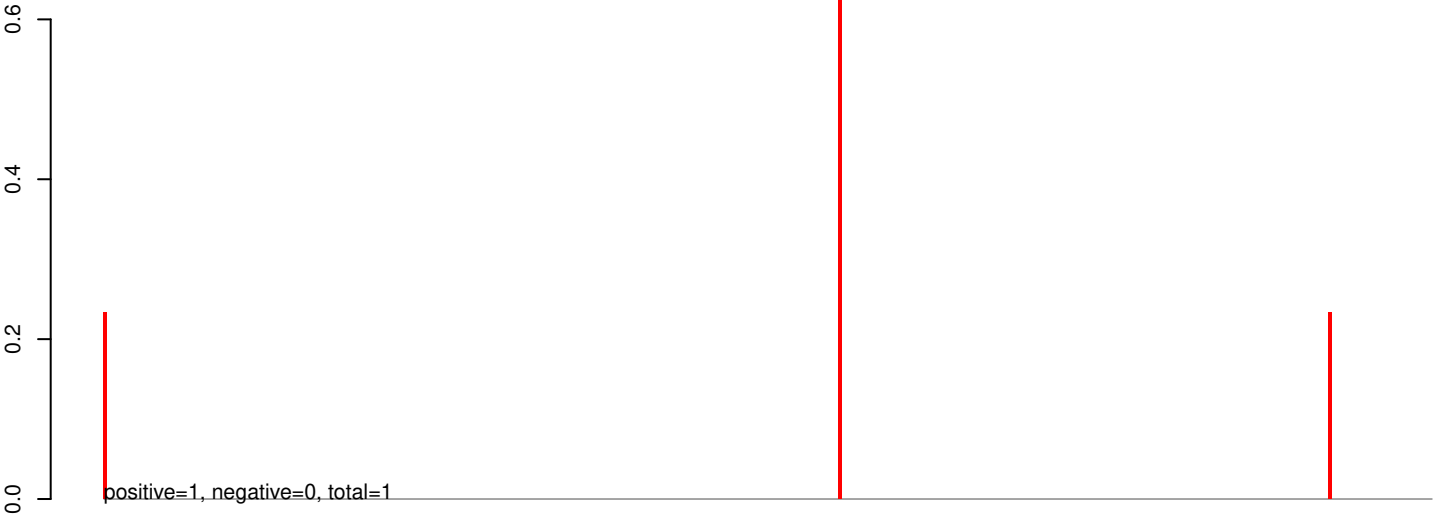
AeAeg_Aag2SINV_dsLuc.24_35.rep



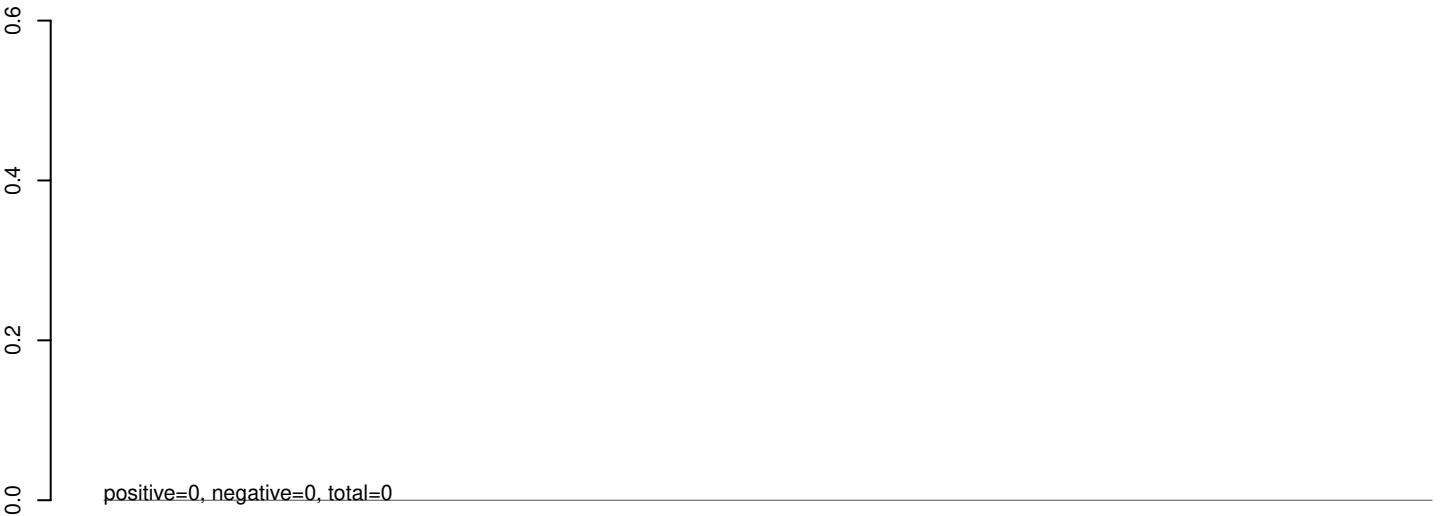
AeAeg_Aag2SINV_dsLuc.rep



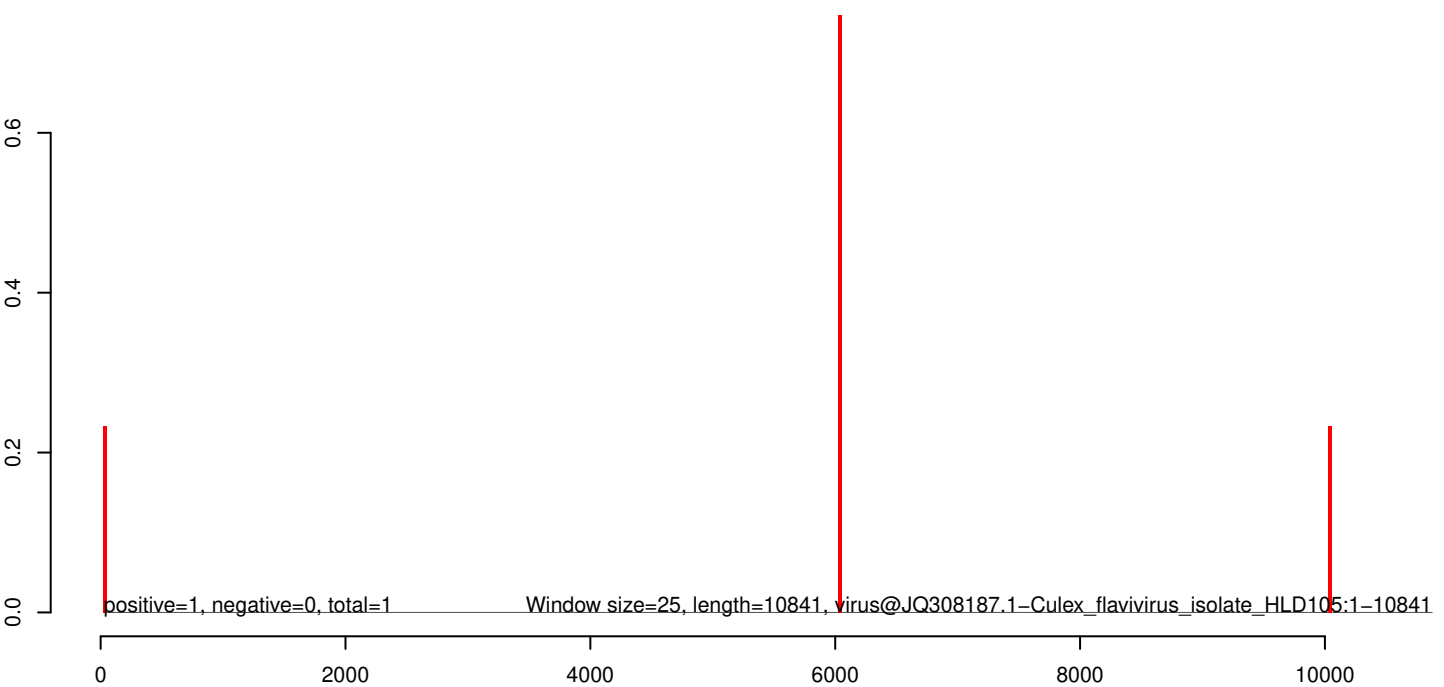
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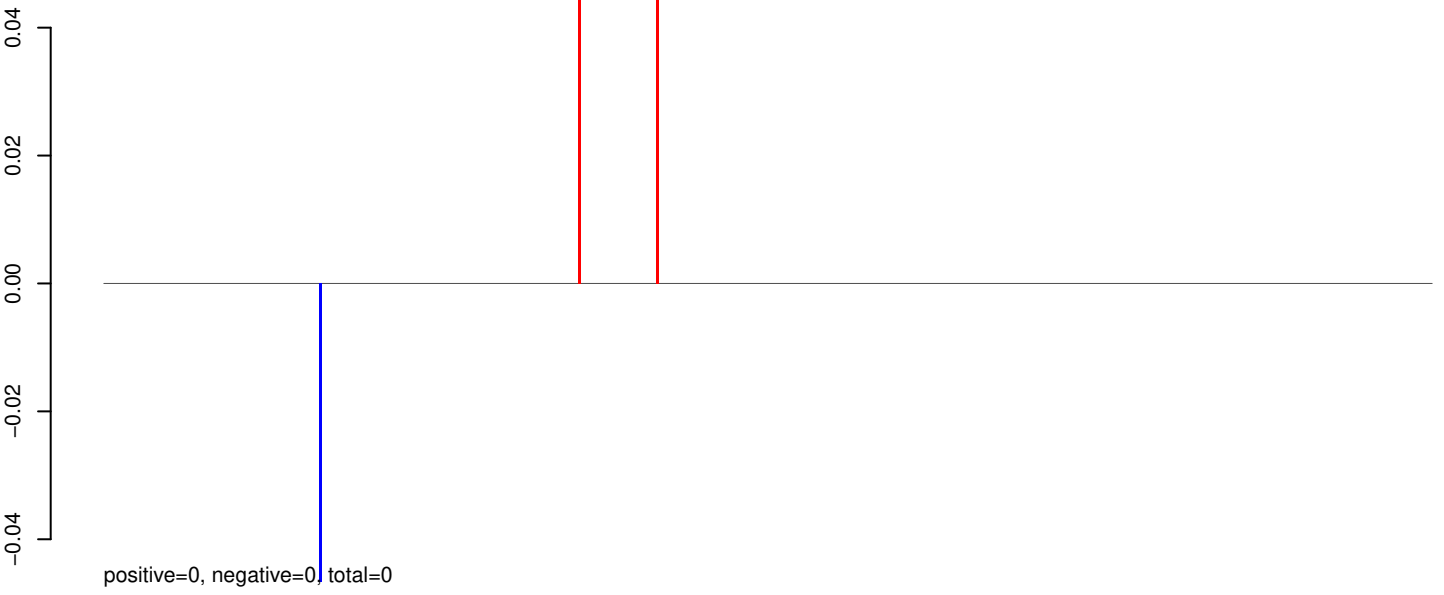
AeAeg_Aag2SINV_dsLuc.24_35.rep



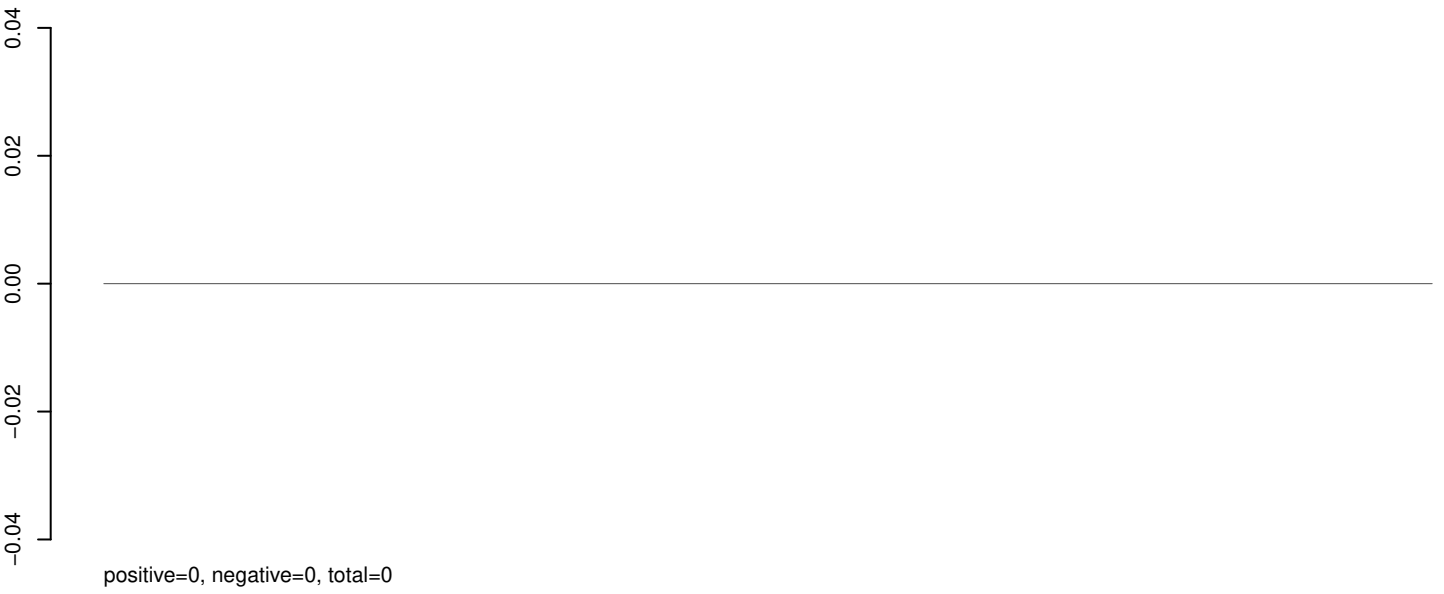
AeAeg_Aag2SINV_dsLuc.rep



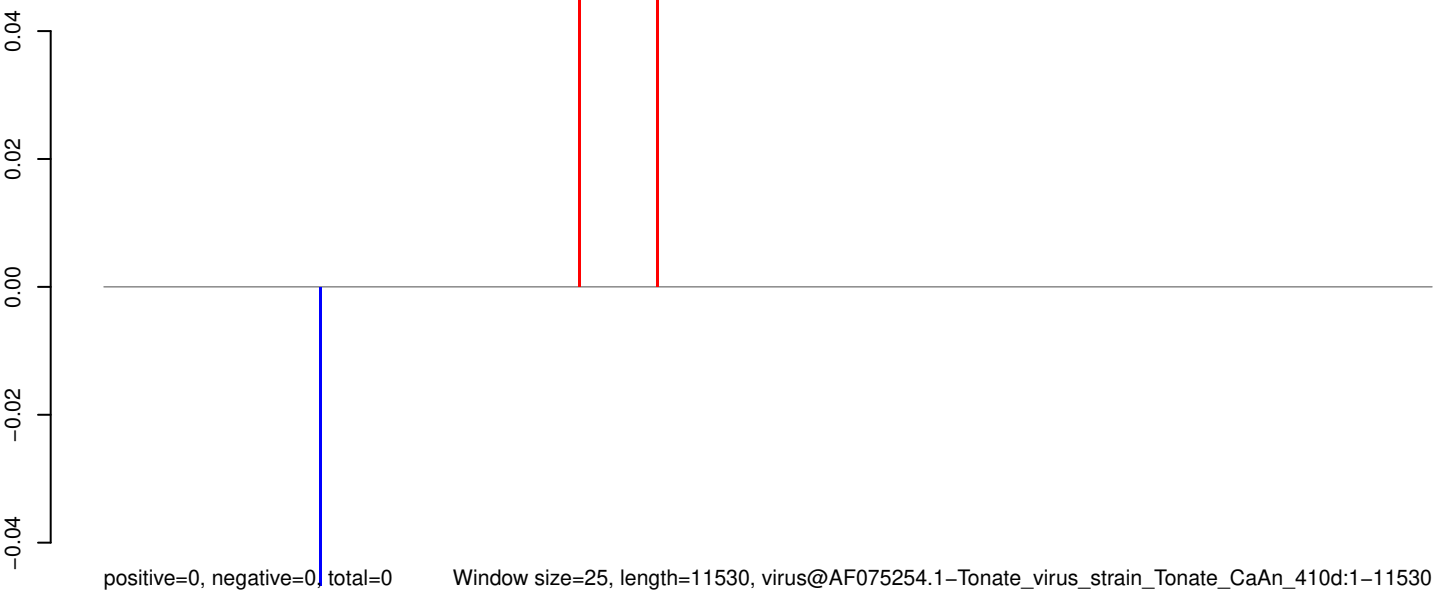
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



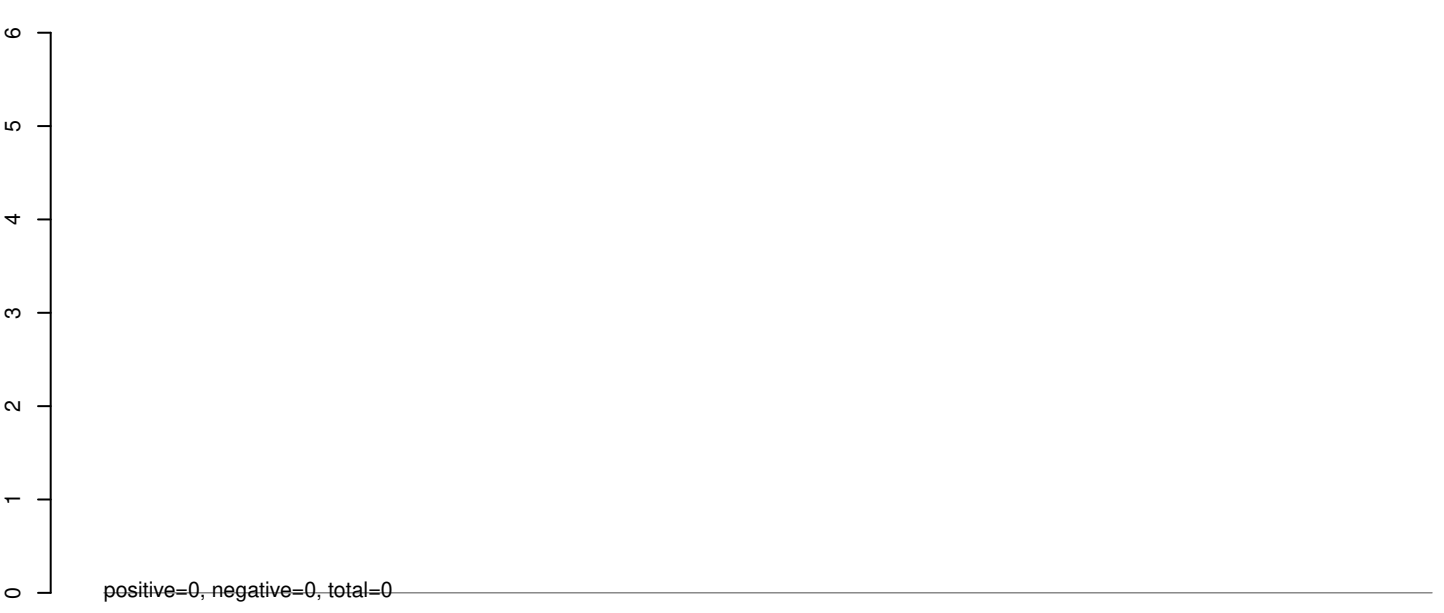
AeAeg_Aag2SINV_dsLuc.rep



AeAeg_Aag2SINV_dsLuc.18_23.rep



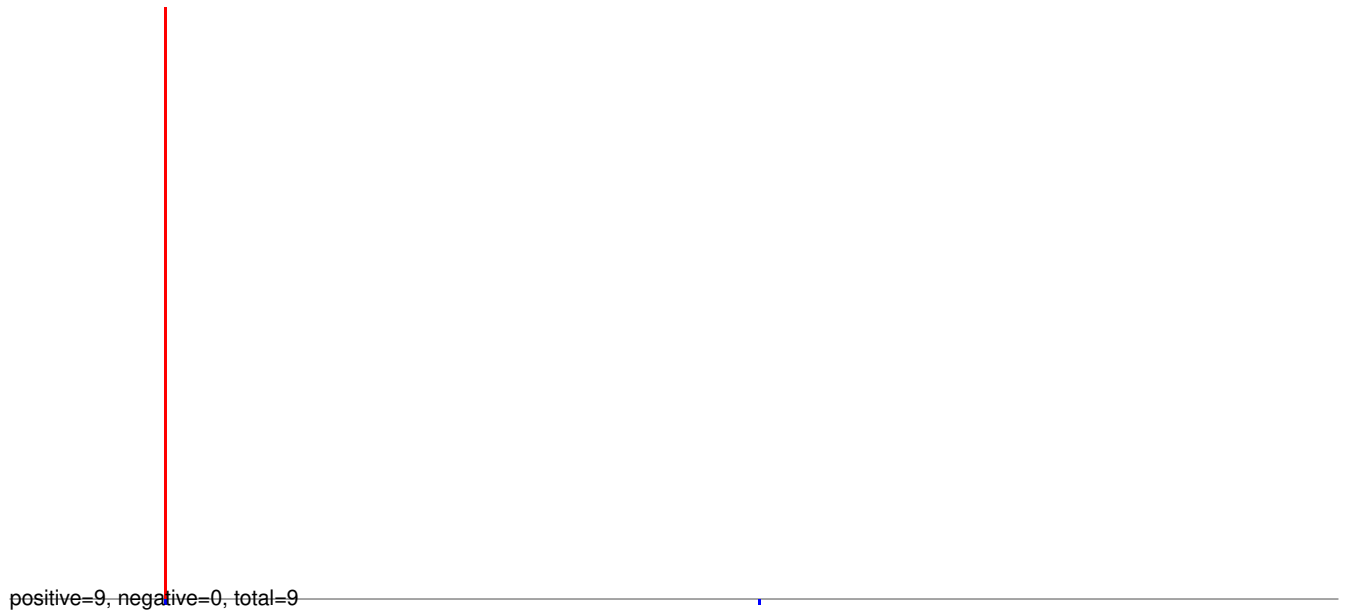
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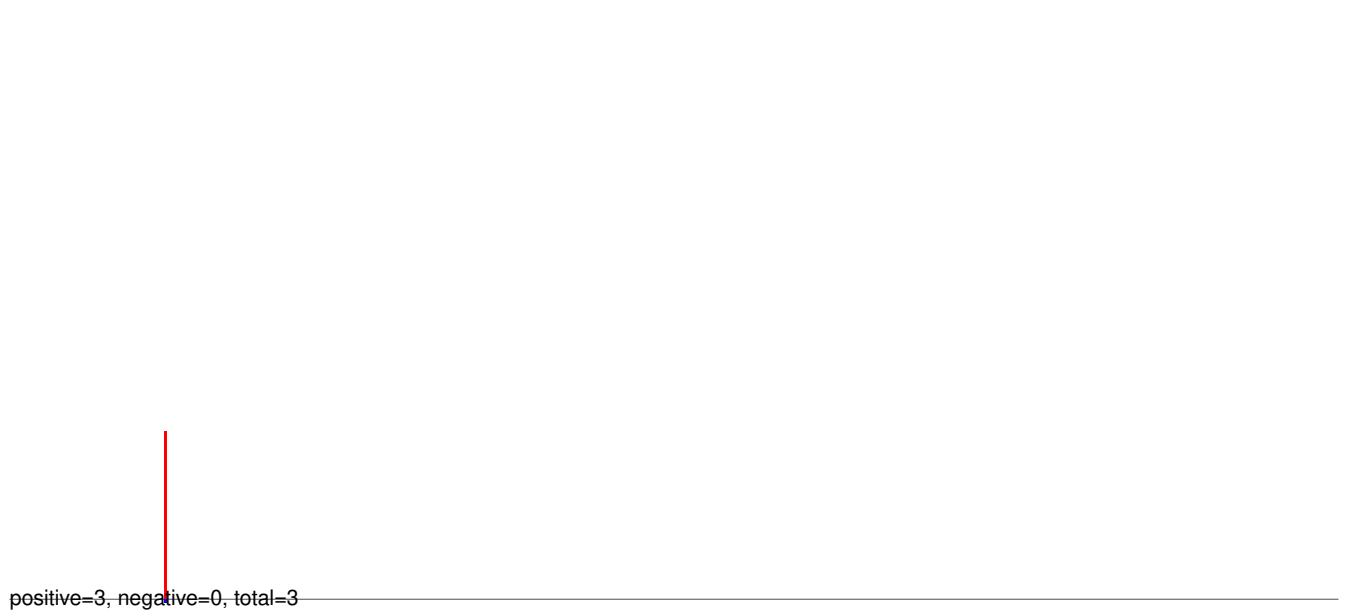
AeAeg_Aag2SINV_dsLuc.rep



AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



positive=12, negative=0, total=12, virus@NC_031319.1_Wutai_Mosquito_Virus_strain_QN3-5_nucleopasid_protein_N_G_L:1-12072

0 2000 4000 6000 8000 10000 12000

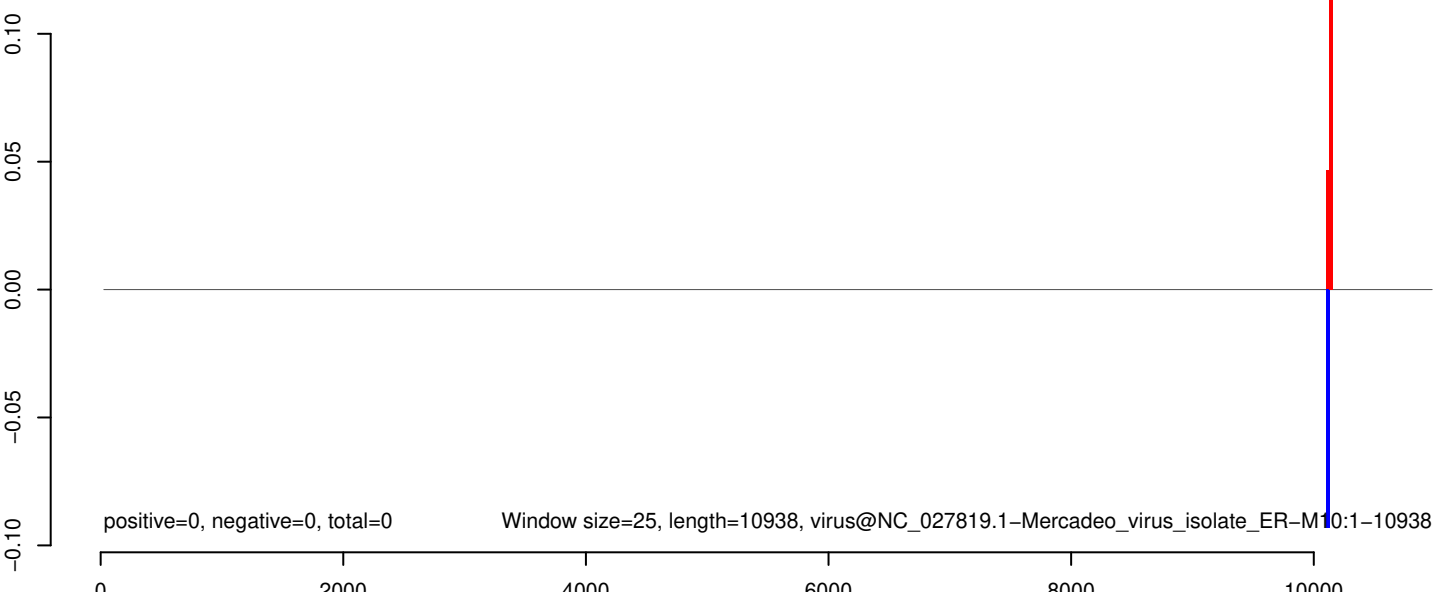
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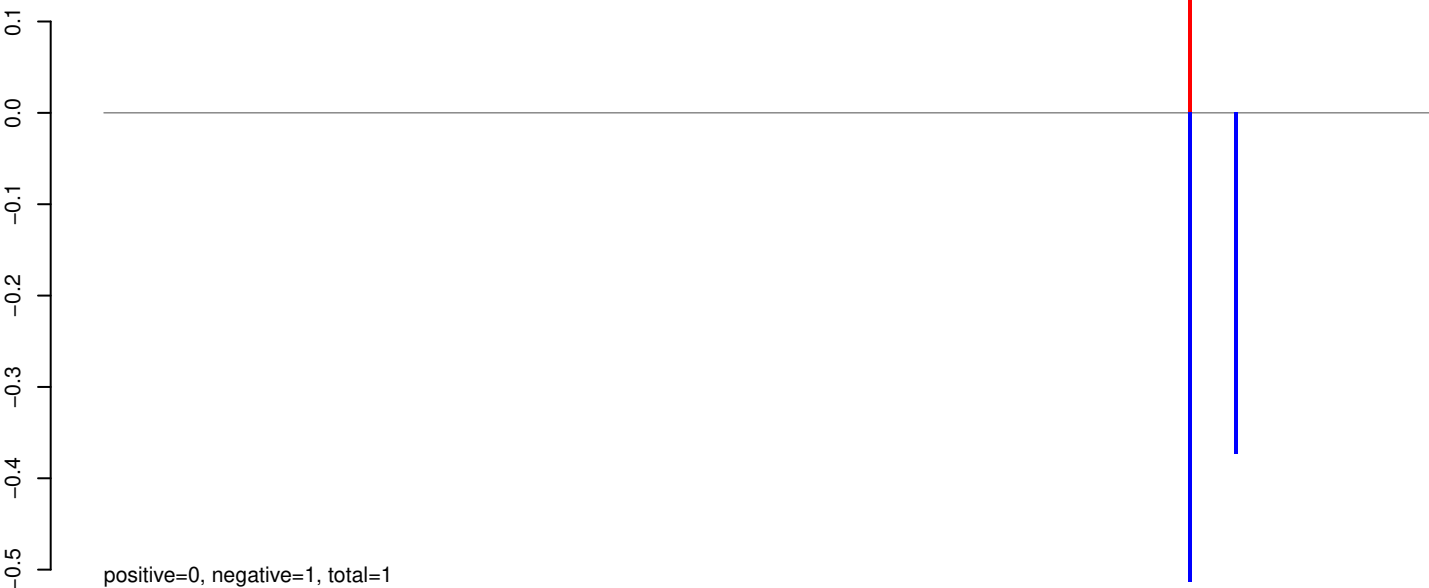
AeAeg_Aag2SINV_dsLuc.24_35.rep



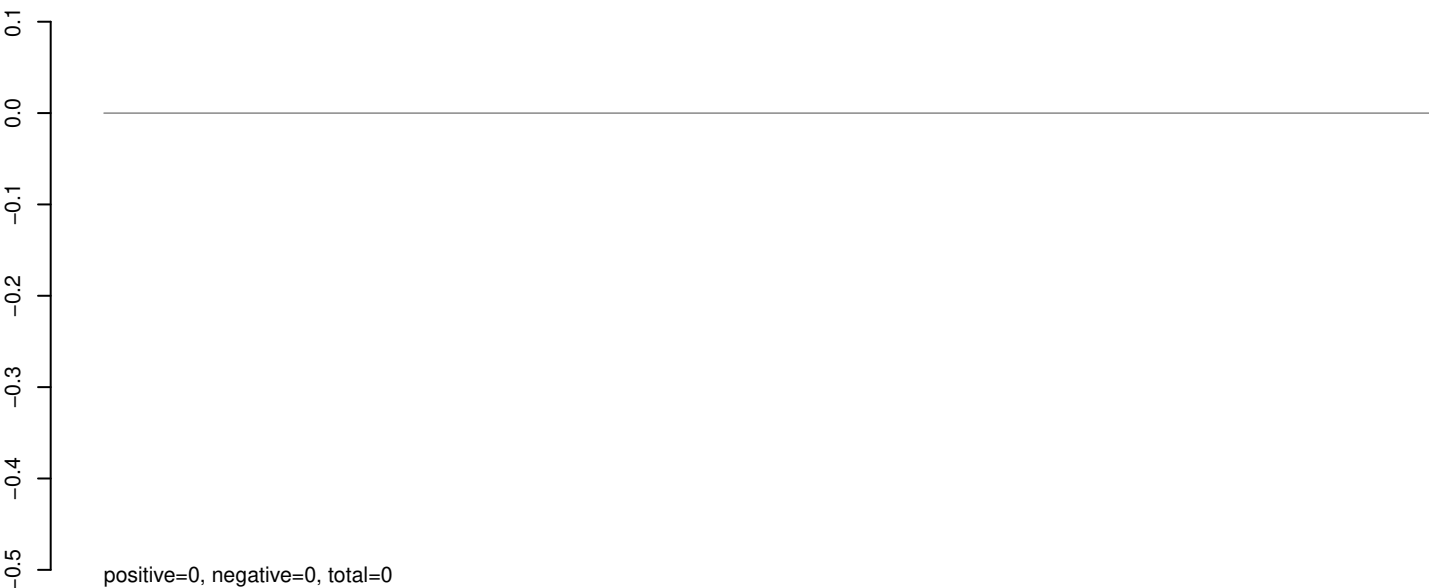
AeAeg_Aag2SINV_dsLuc.rep



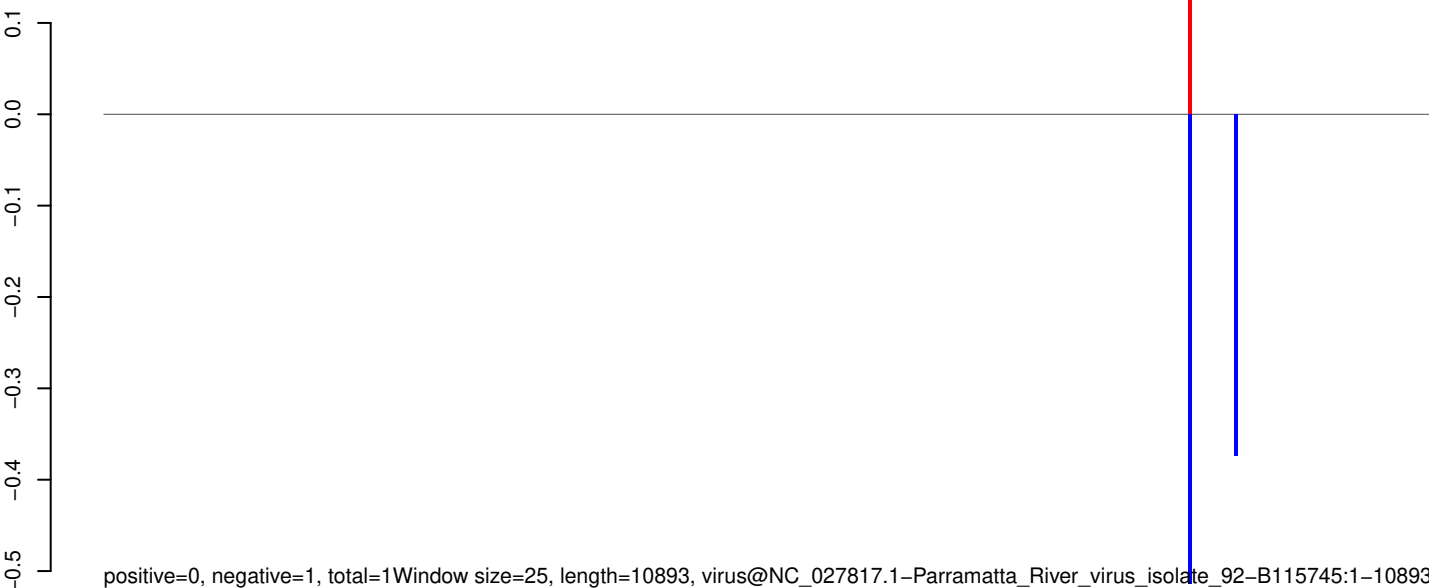
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AeAeg_Aag2SINV_dsLuc.24_35.rep

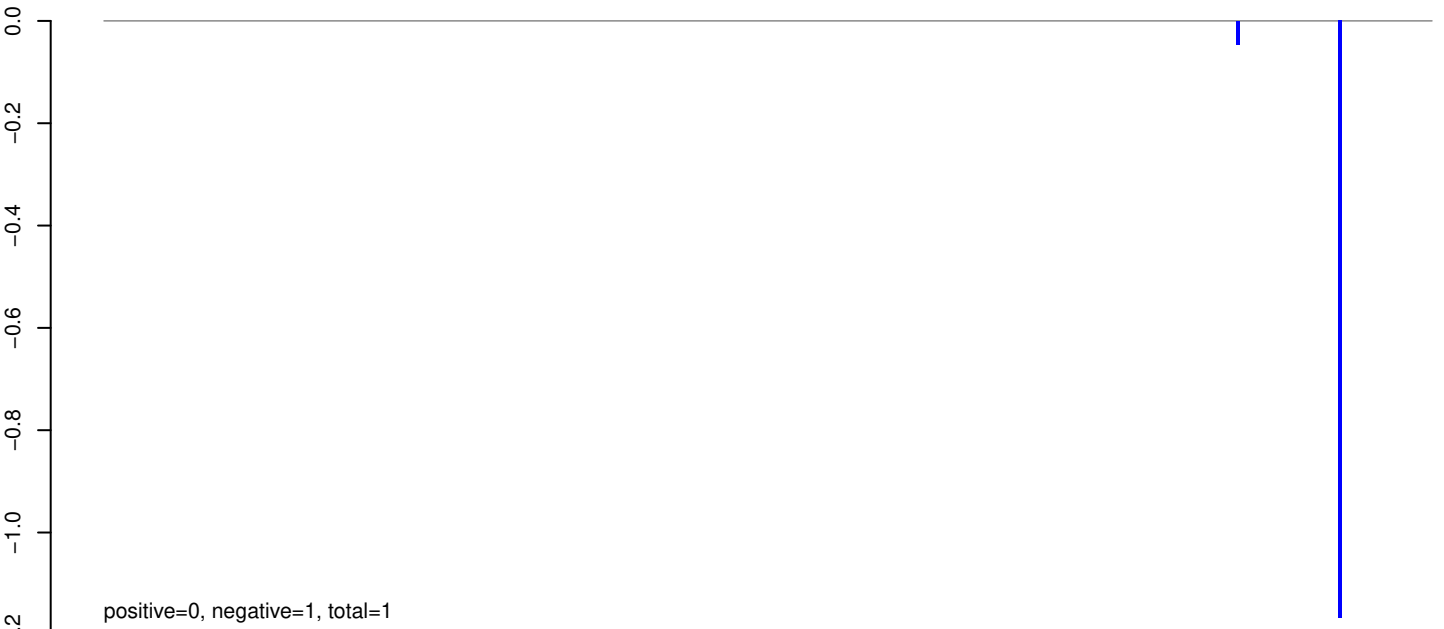


AeAeg_Aag2SINV_dsLuc.rep

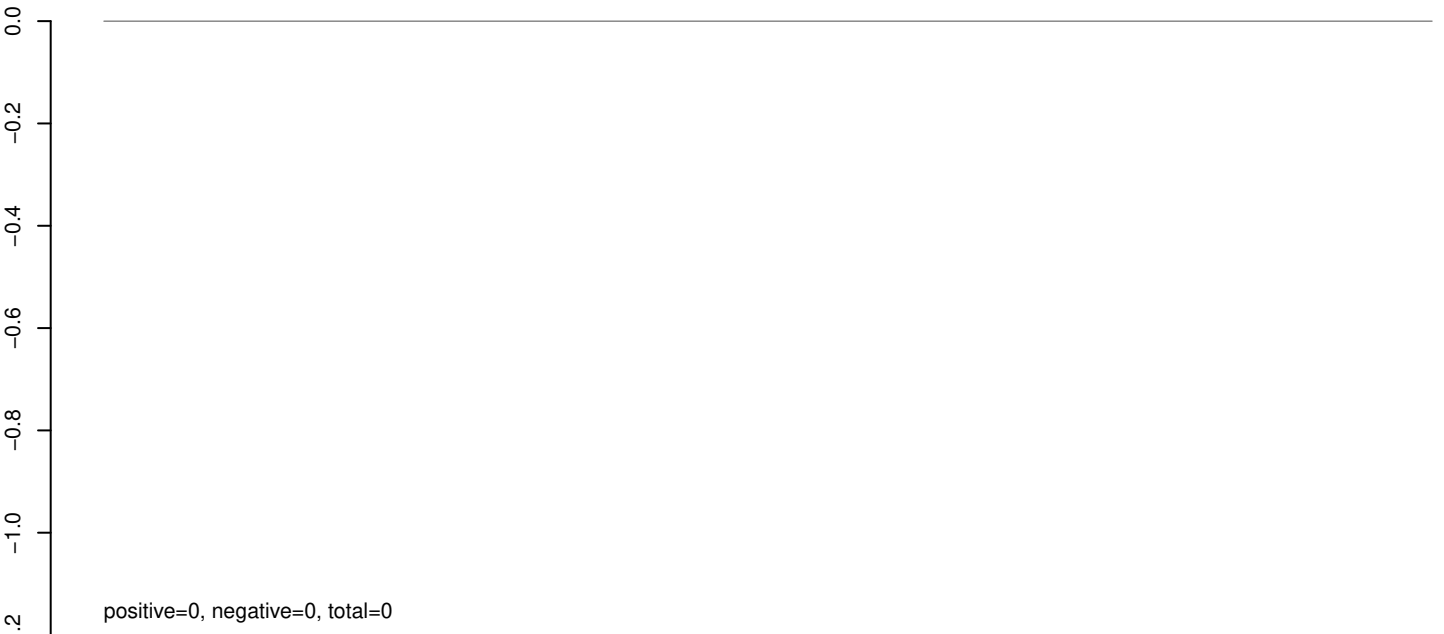


0 2000 4000 6000 8000 10000

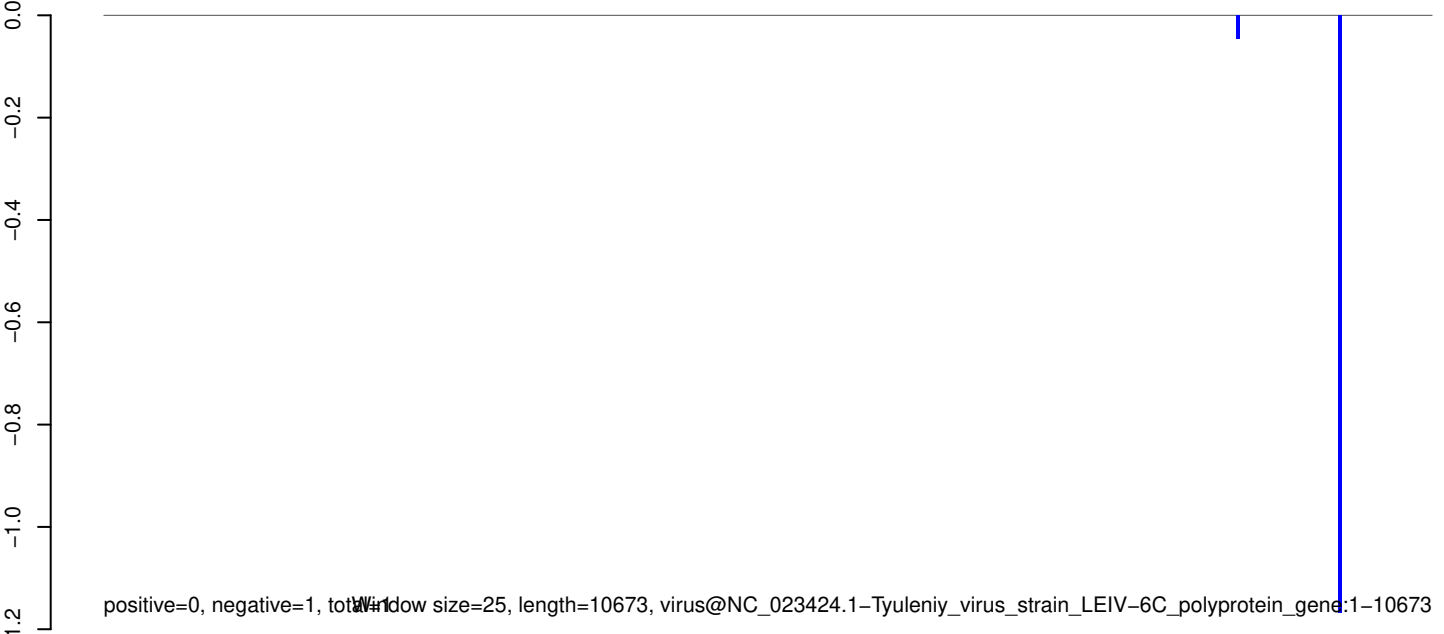
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



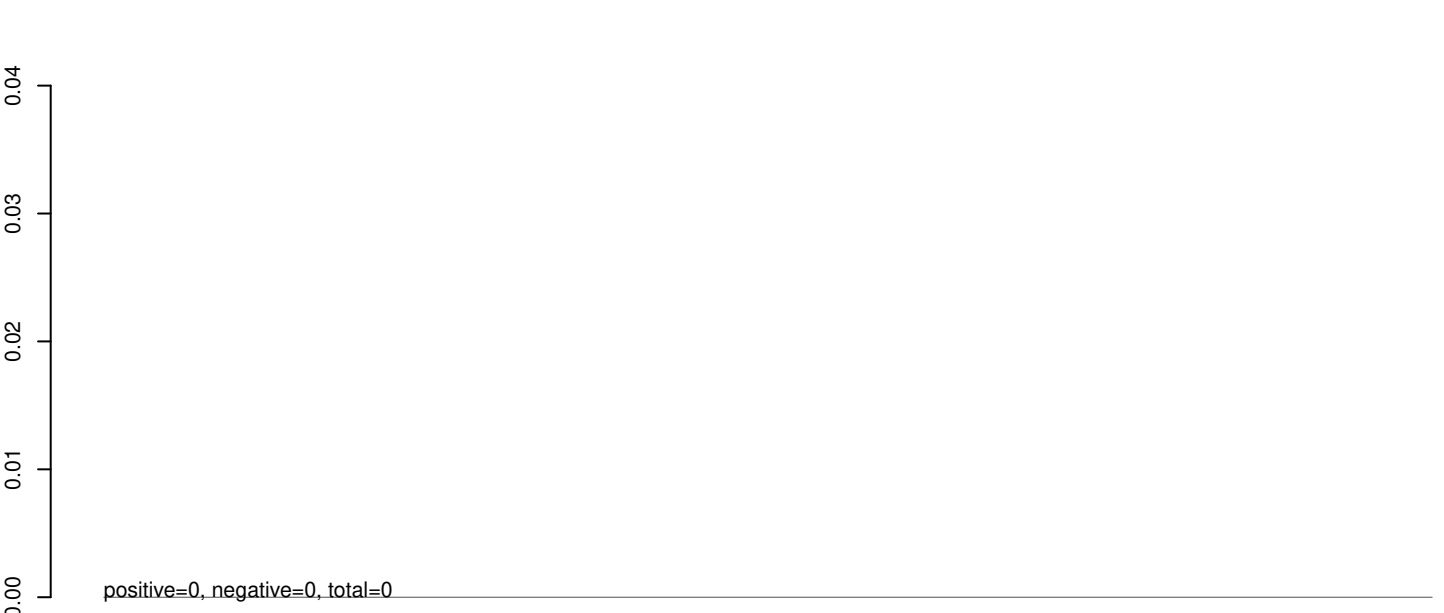
Window size=25, length=10673, virus@NC_023424.1-Tyuleniy_virus_strain_LEIV-6C_polyprotein_gene:1-10673

0 2000 4000 6000 8000 10000

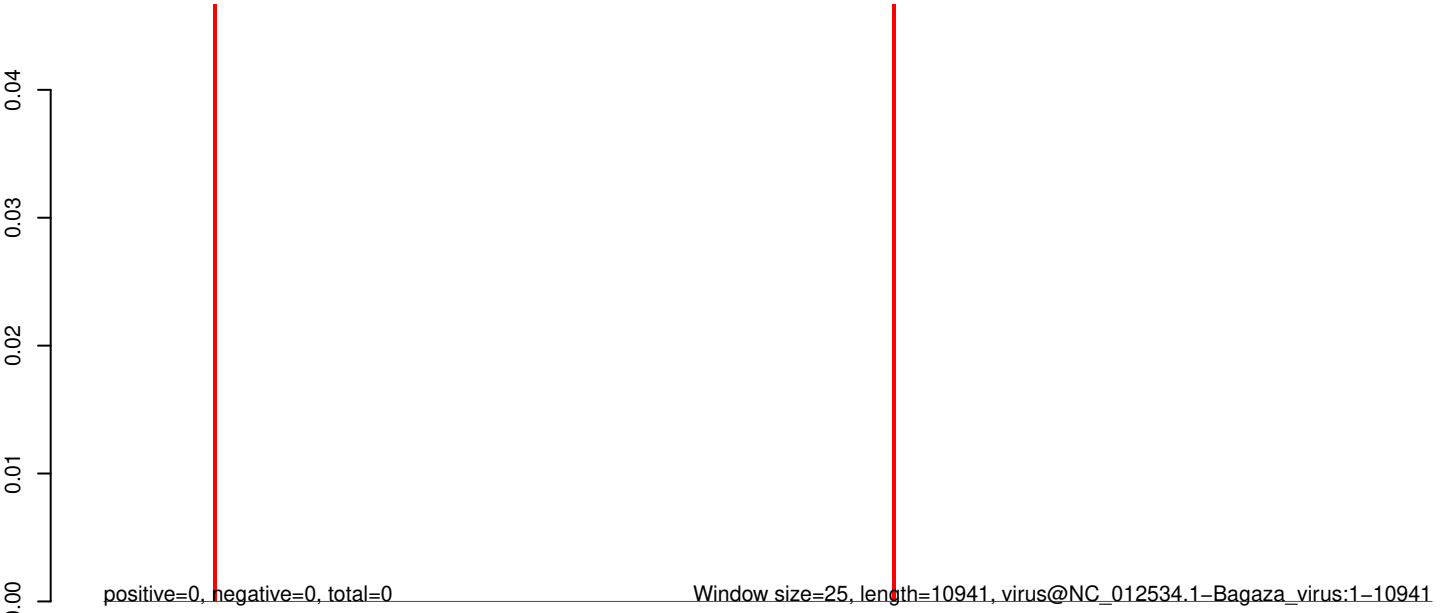
AeAeg_Aag2SINV_dsLuc.18_23.rep



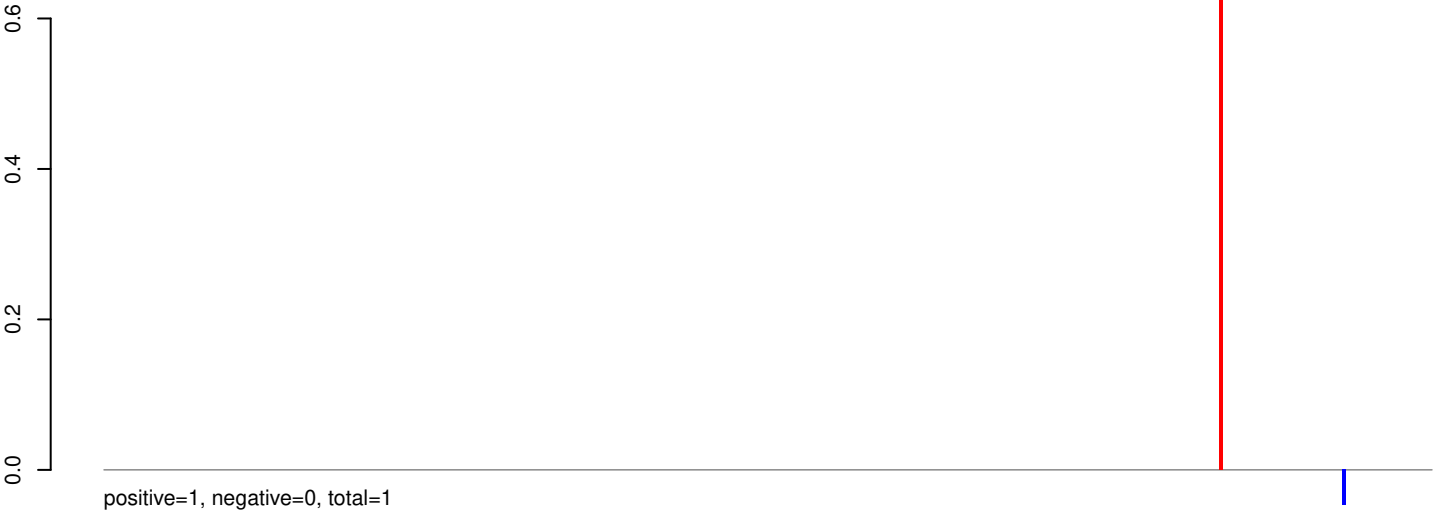
AeAeg_Aag2SINV_dsLuc.24_35.rep



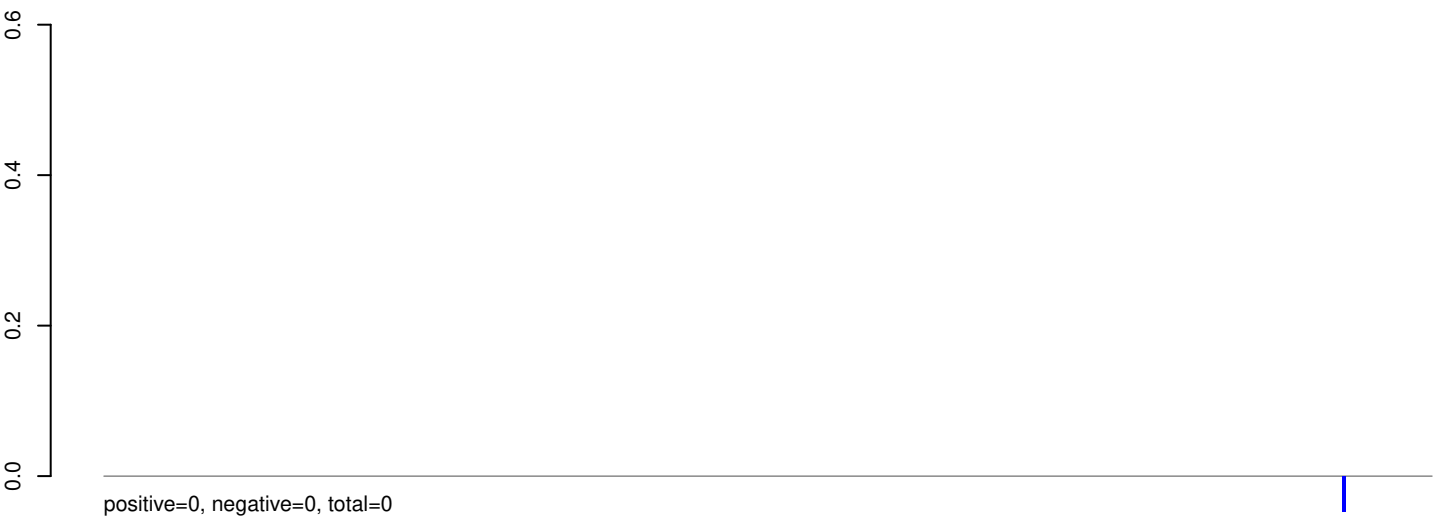
AeAeg_Aag2SINV_dsLuc.rep



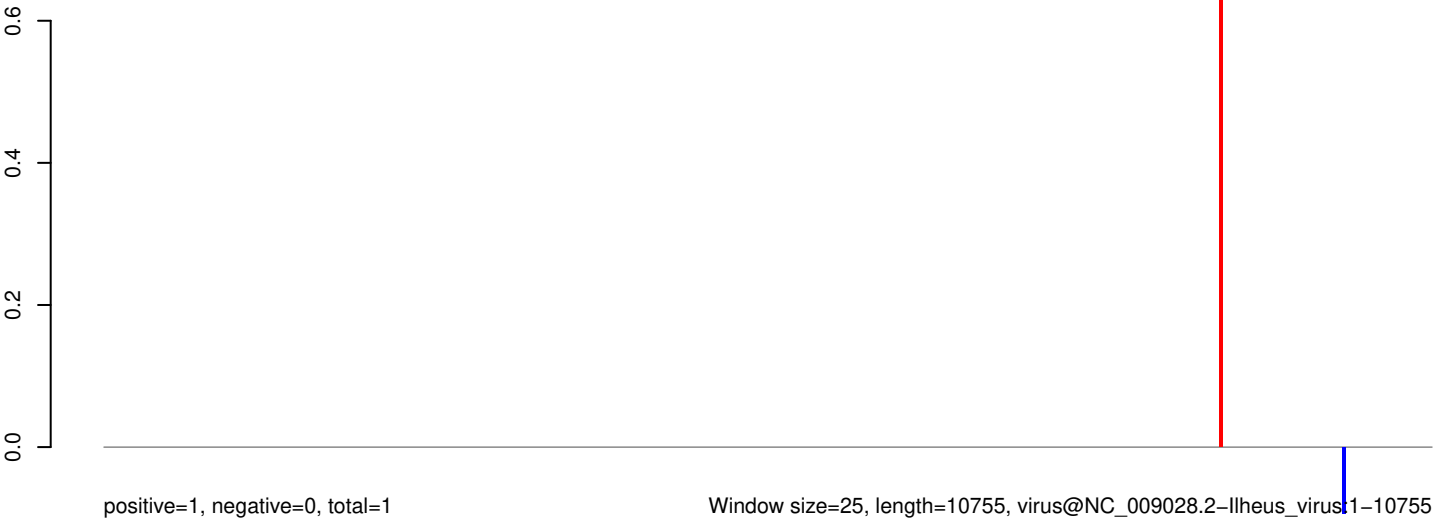
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep

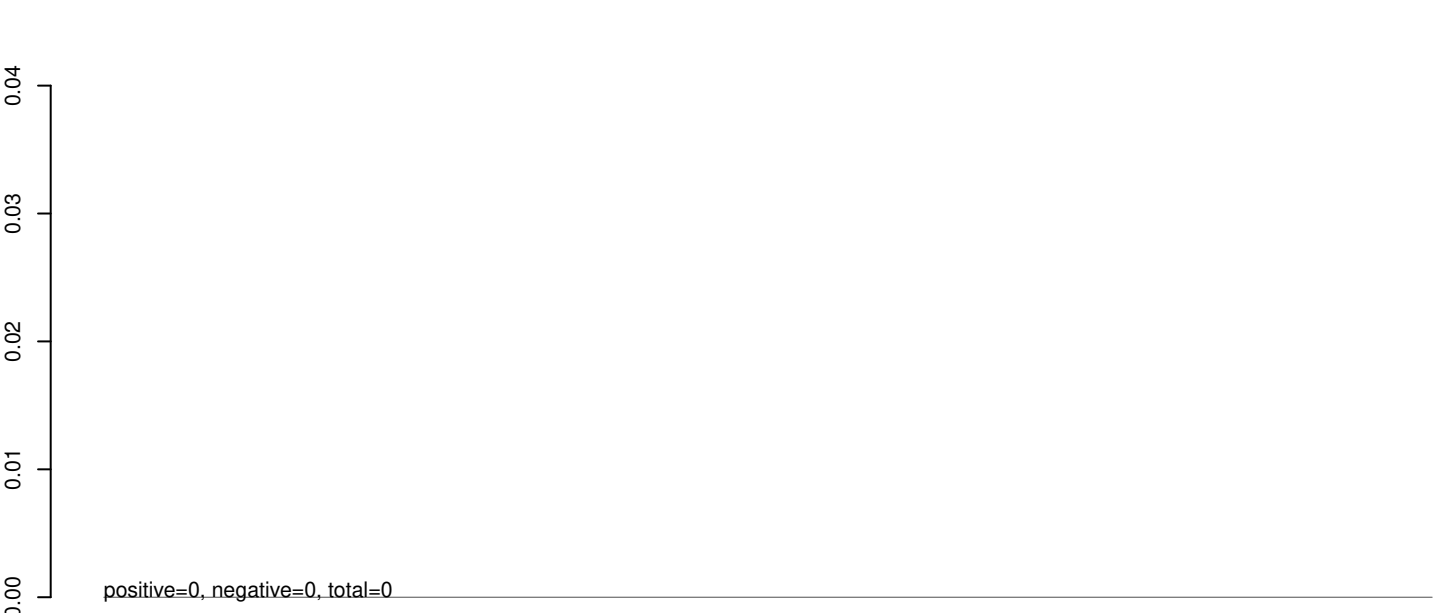


0 2000 4000 6000 8000 10000

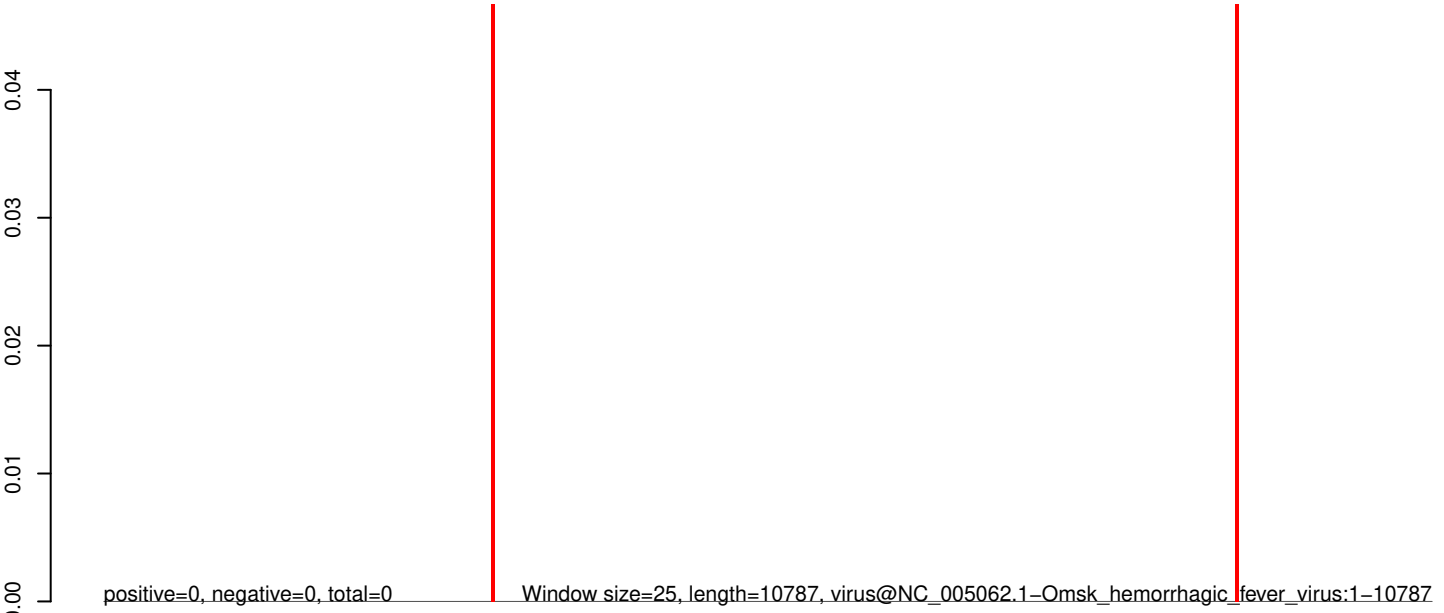
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

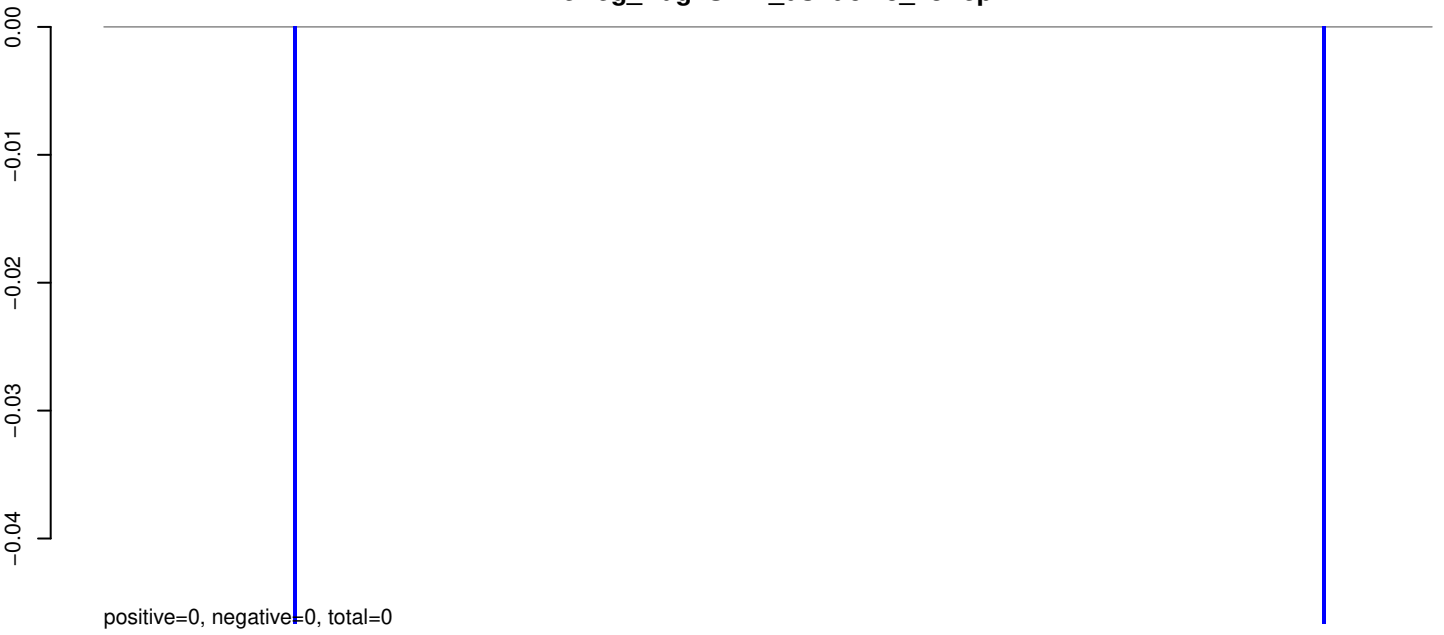


AeAeg_Aag2SINV_dsLuc.rep

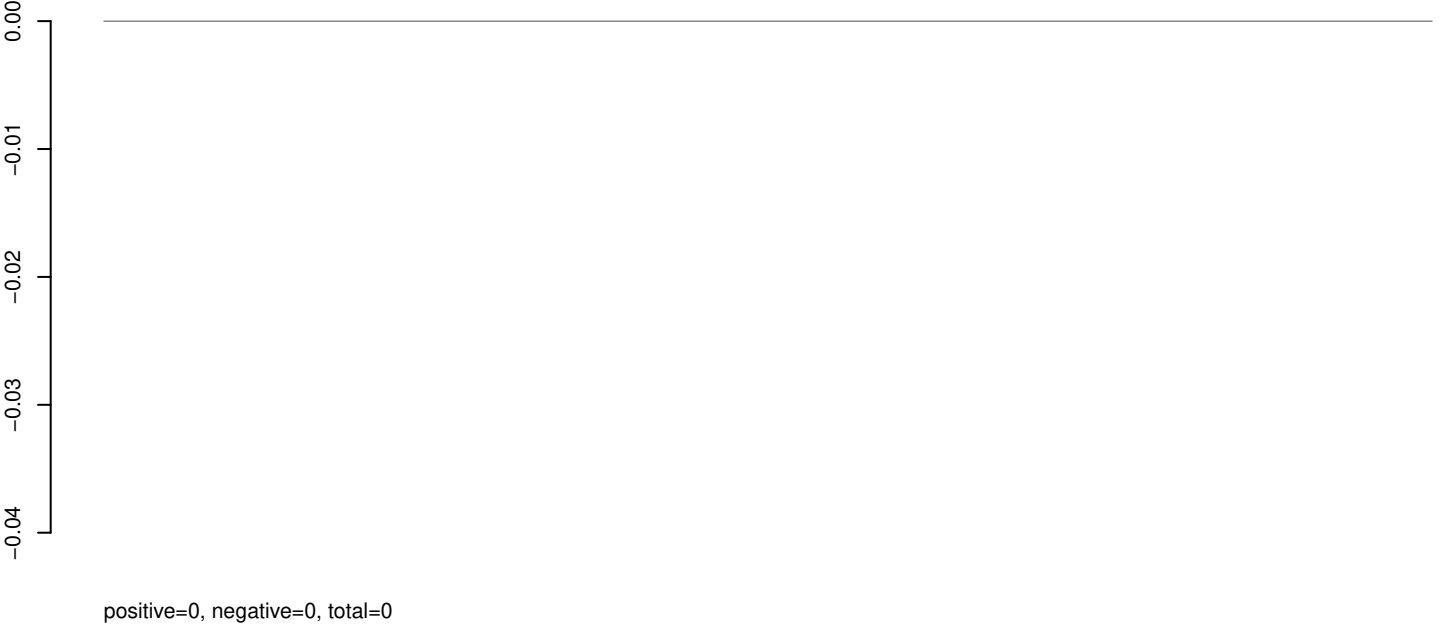


0 2000 4000 6000 8000 10000

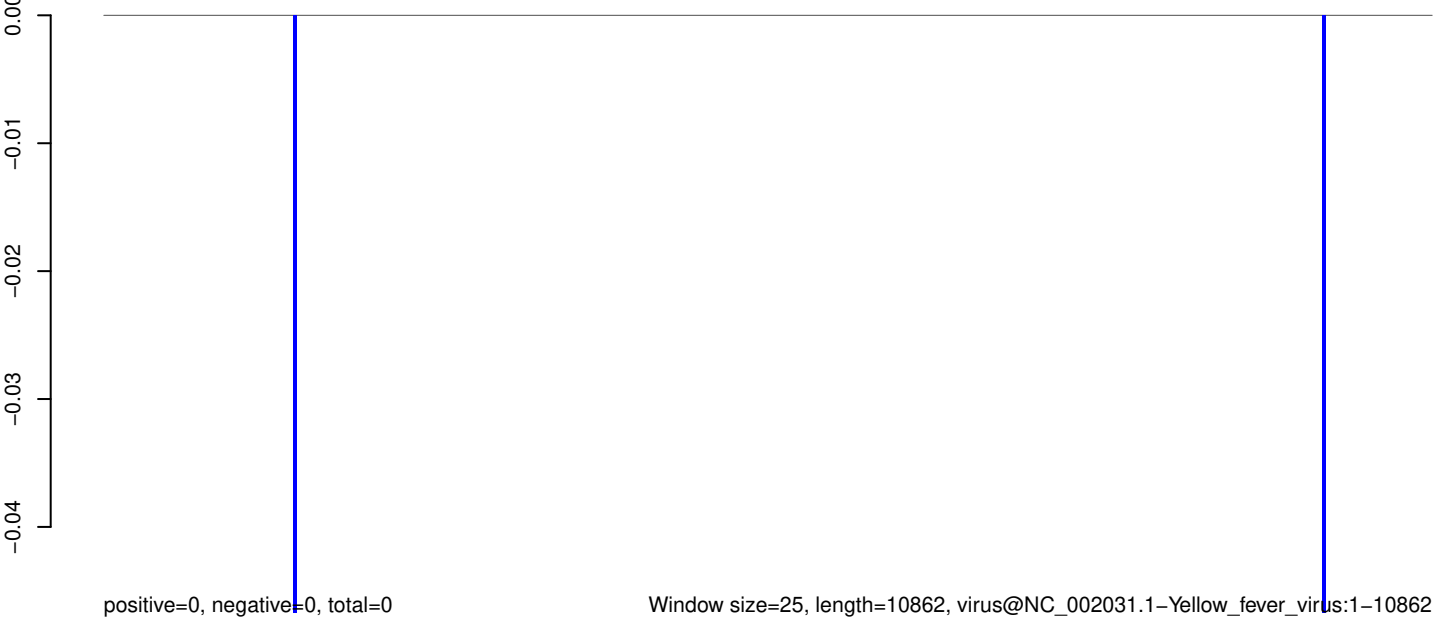
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

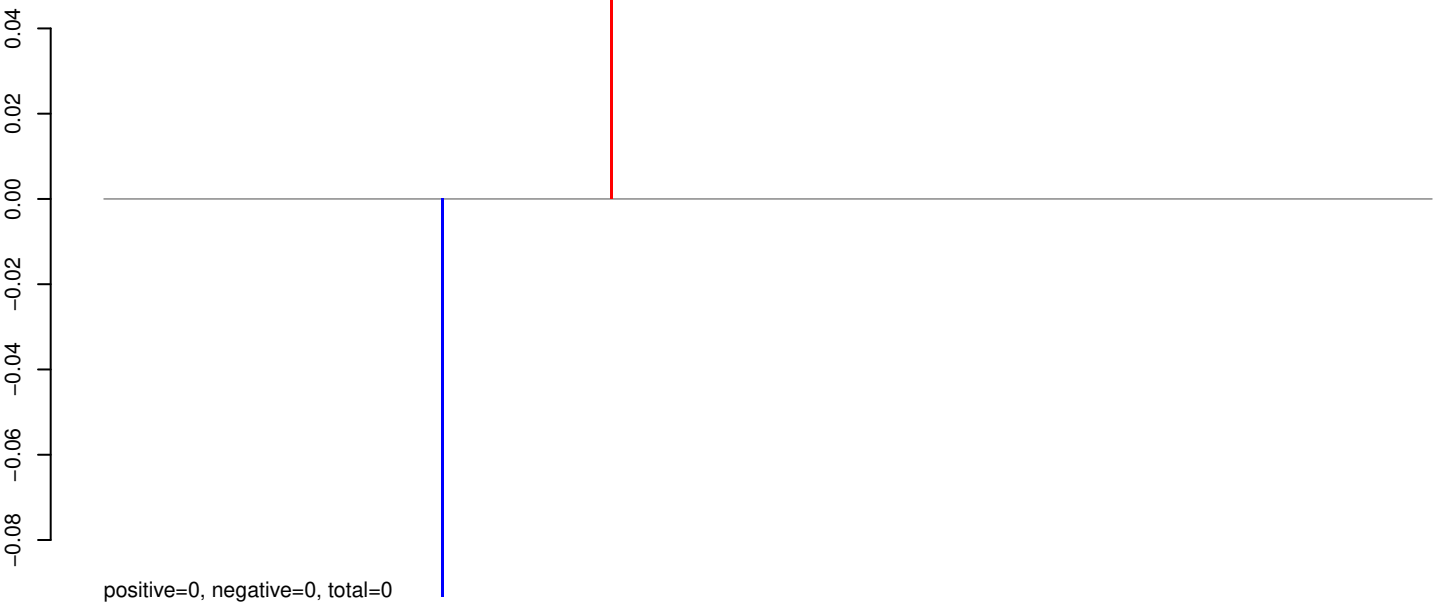


AeAeg_Aag2SINV_dsLuc.rep

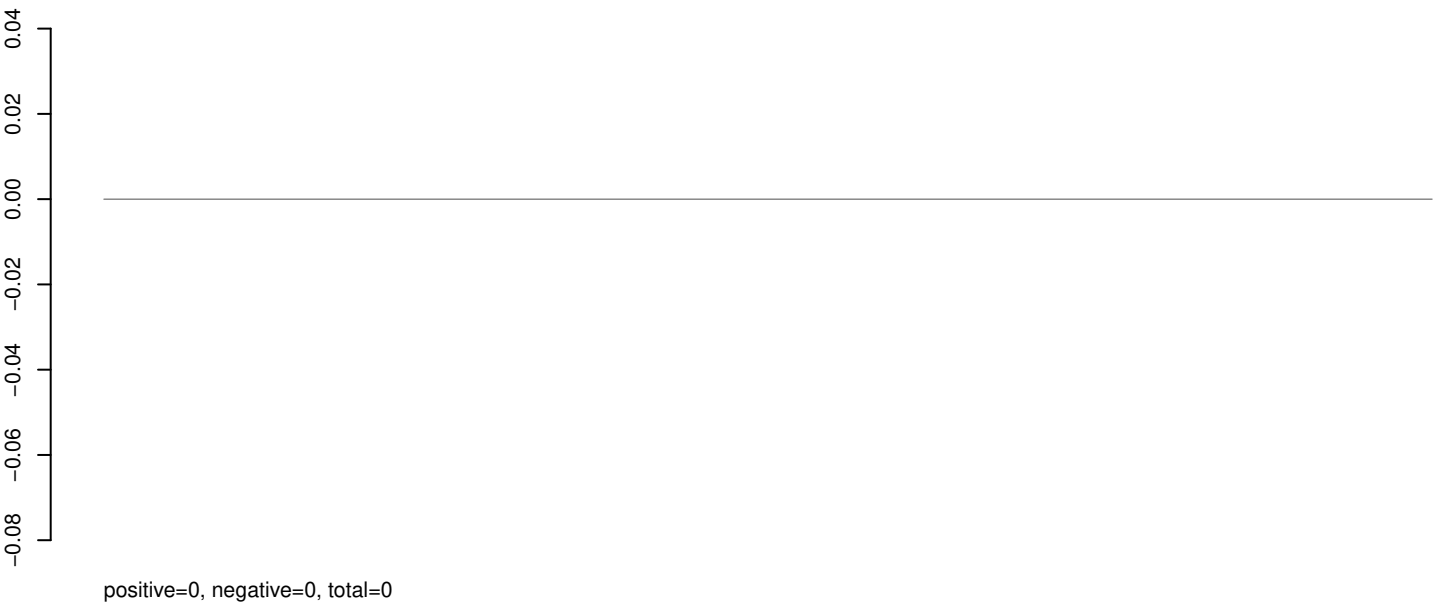


0 2000 4000 6000 8000 10000

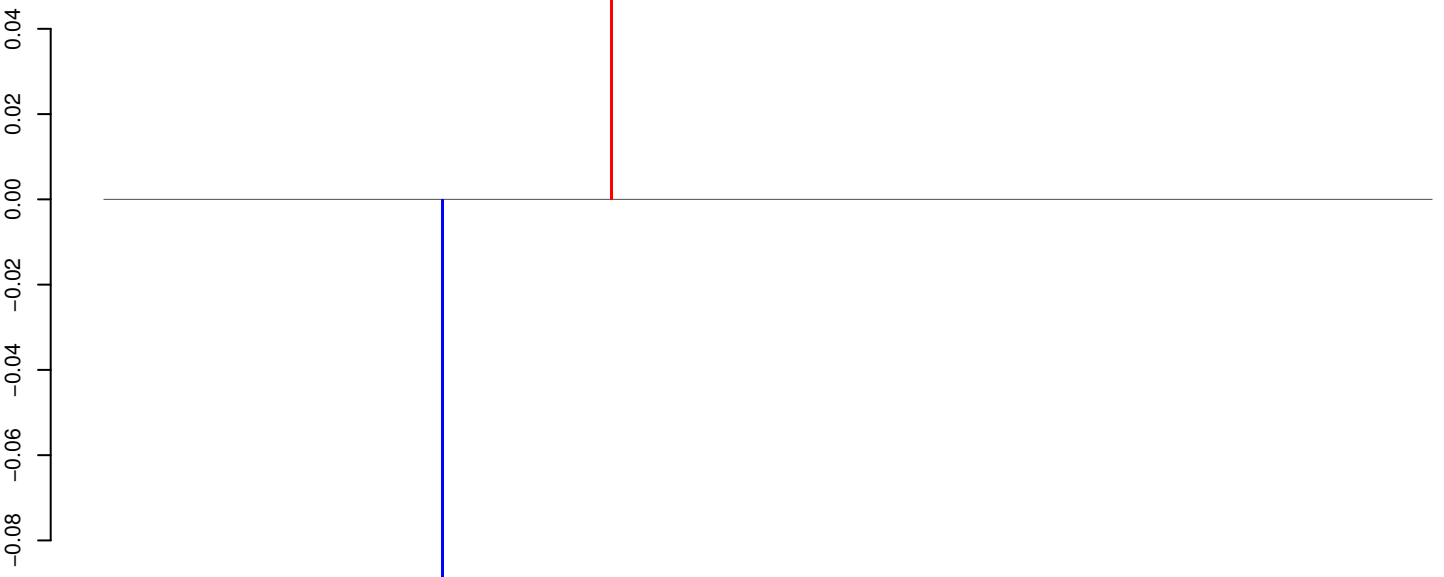
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



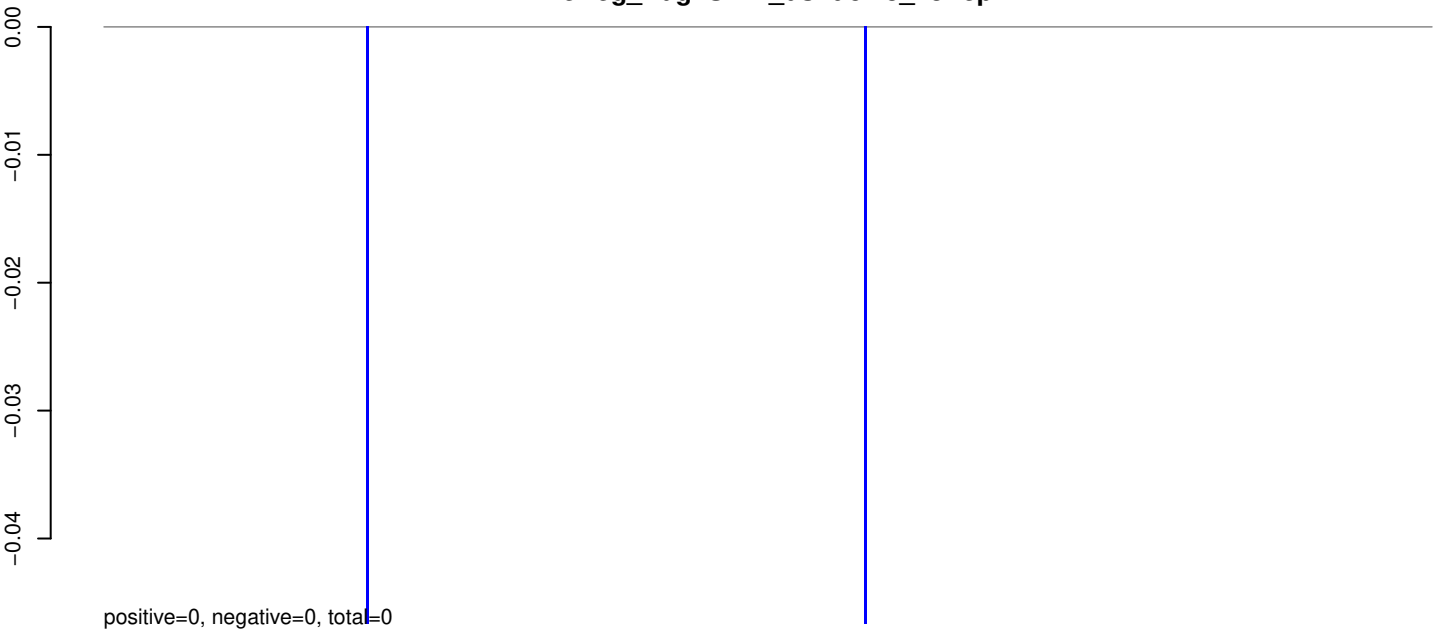
AeAeg_Aag2SINV_dsLuc.rep



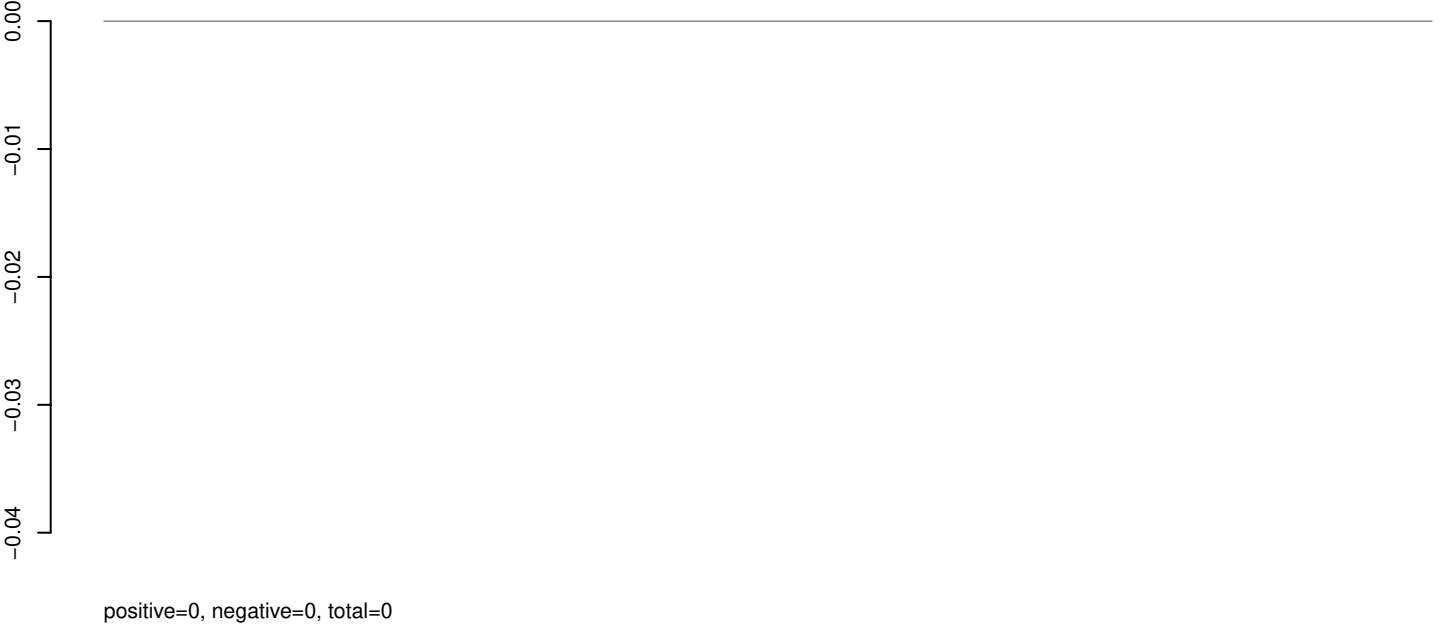
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0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

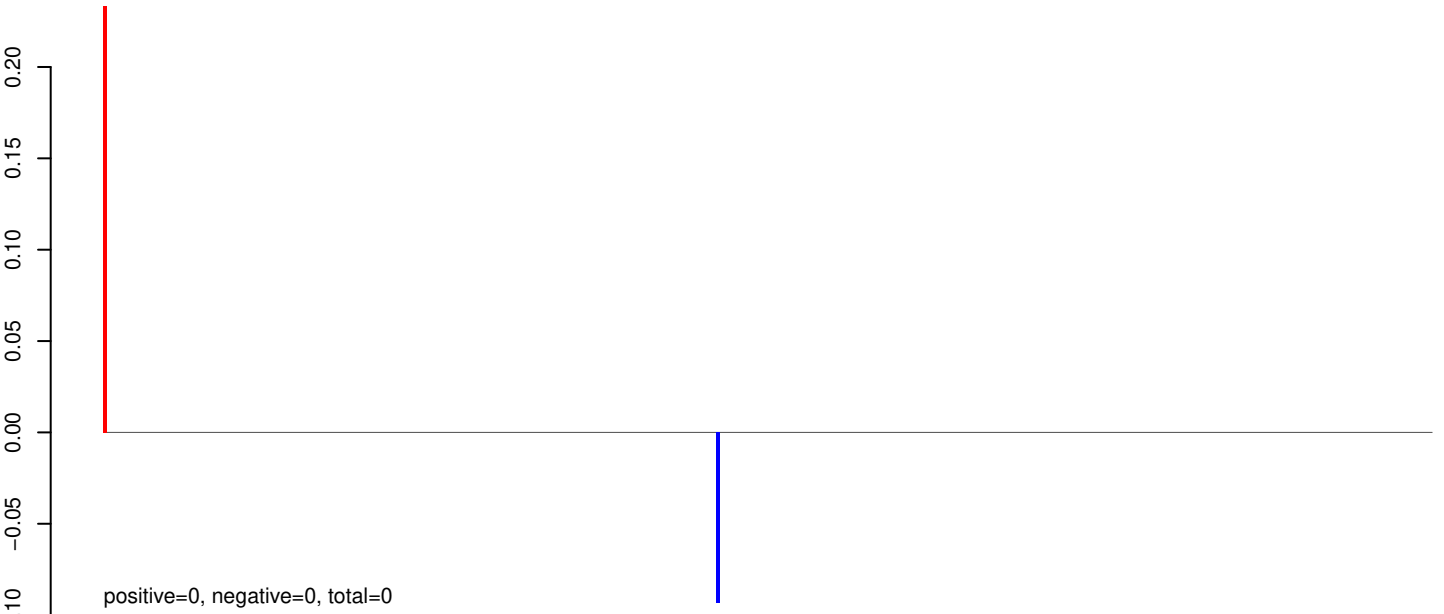


AeAeg_Aag2SINV_dsLuc.rep

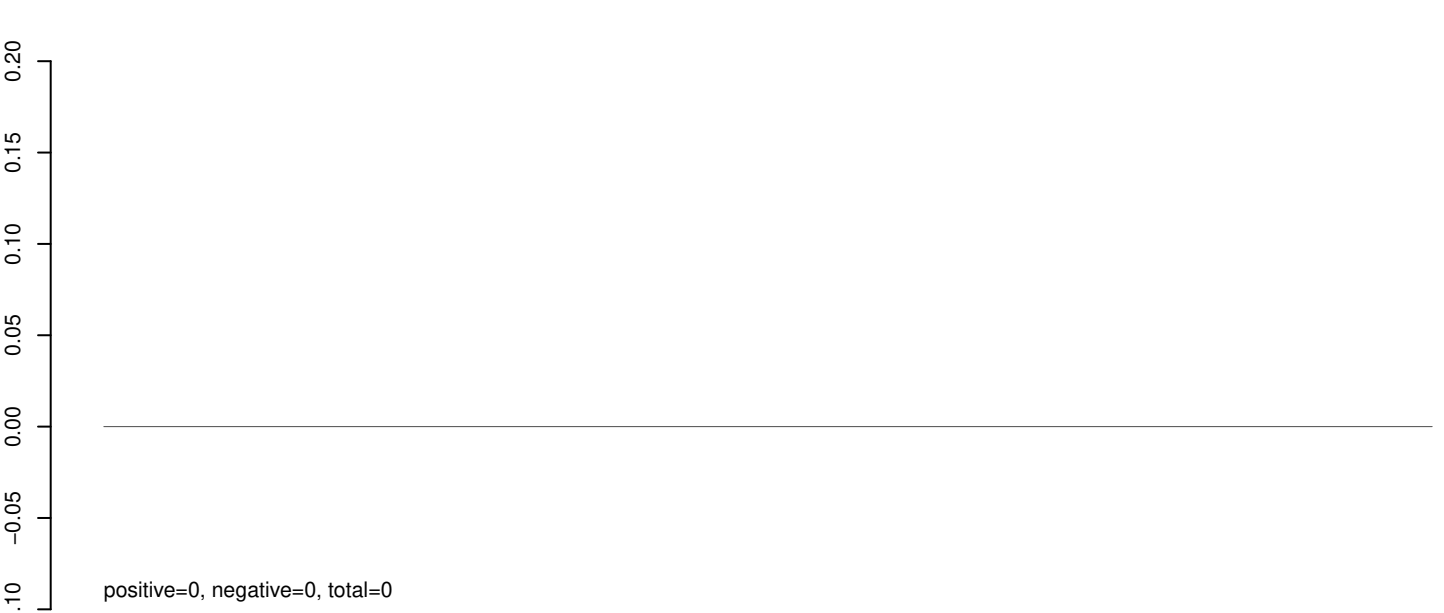


0 2000 4000 6000 8000 10000

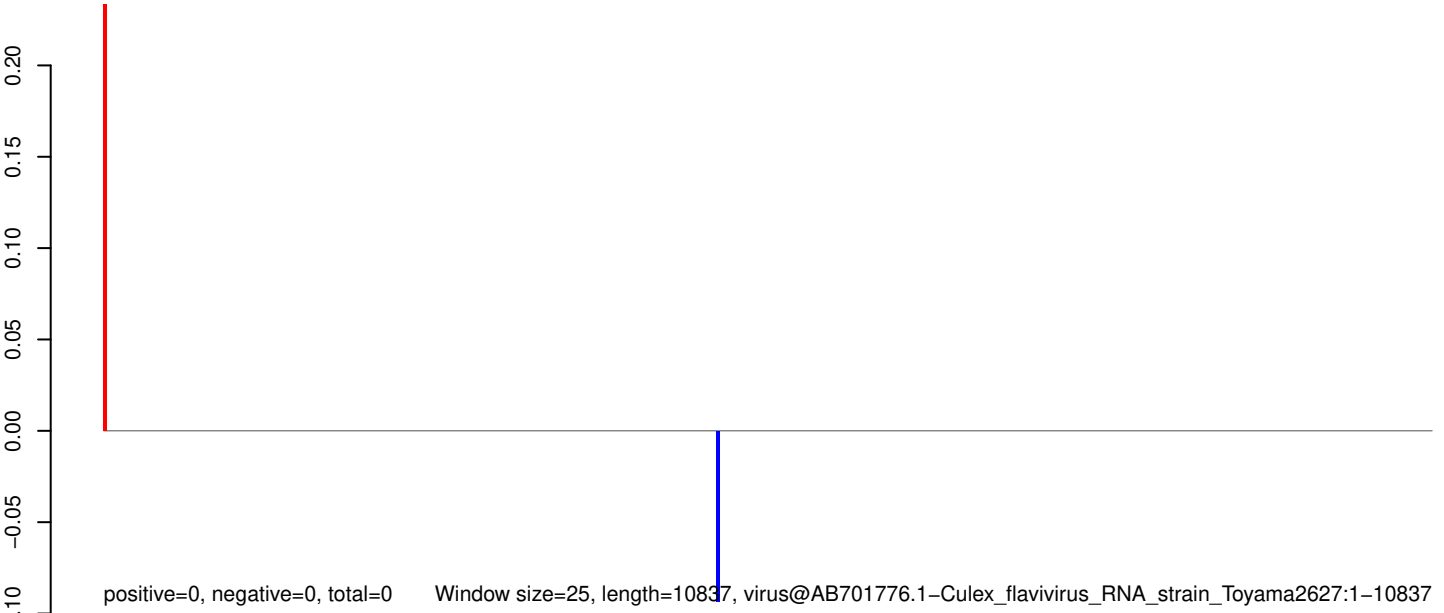
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



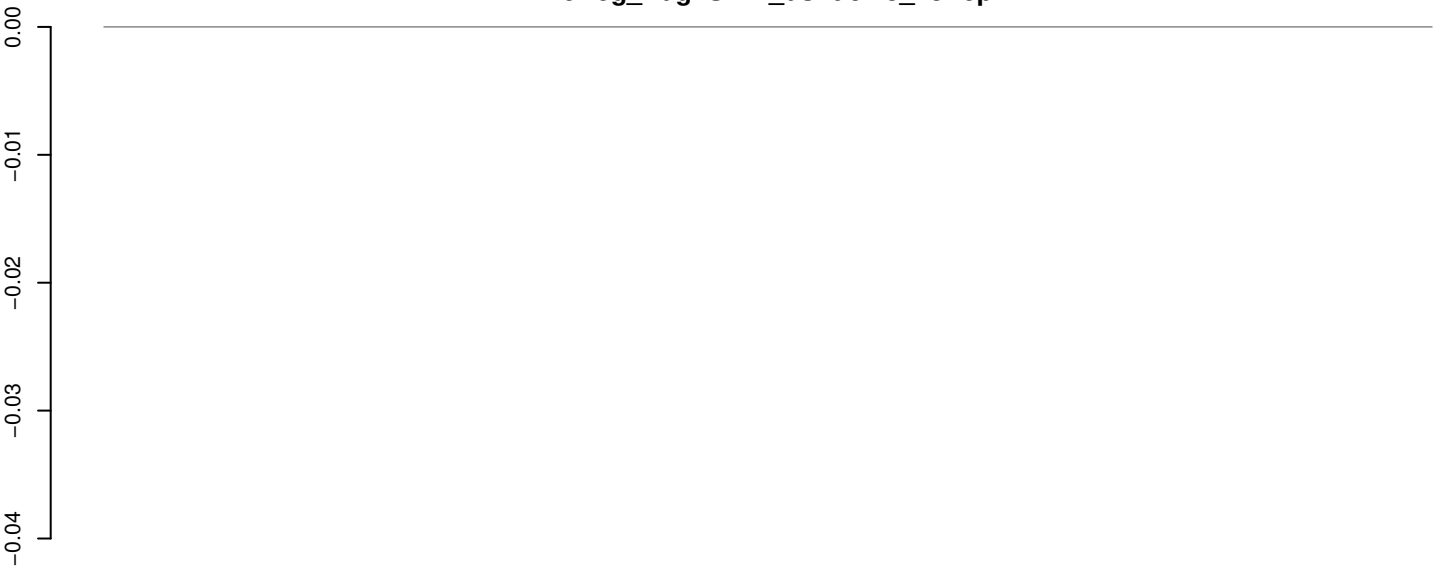
AeAeg_Aag2SINV_dsLuc.rep



Window size=25, length=10837, virus@AB701776.1-Culex_flavivirus_RNA_strain_Toyama2627:1-10837

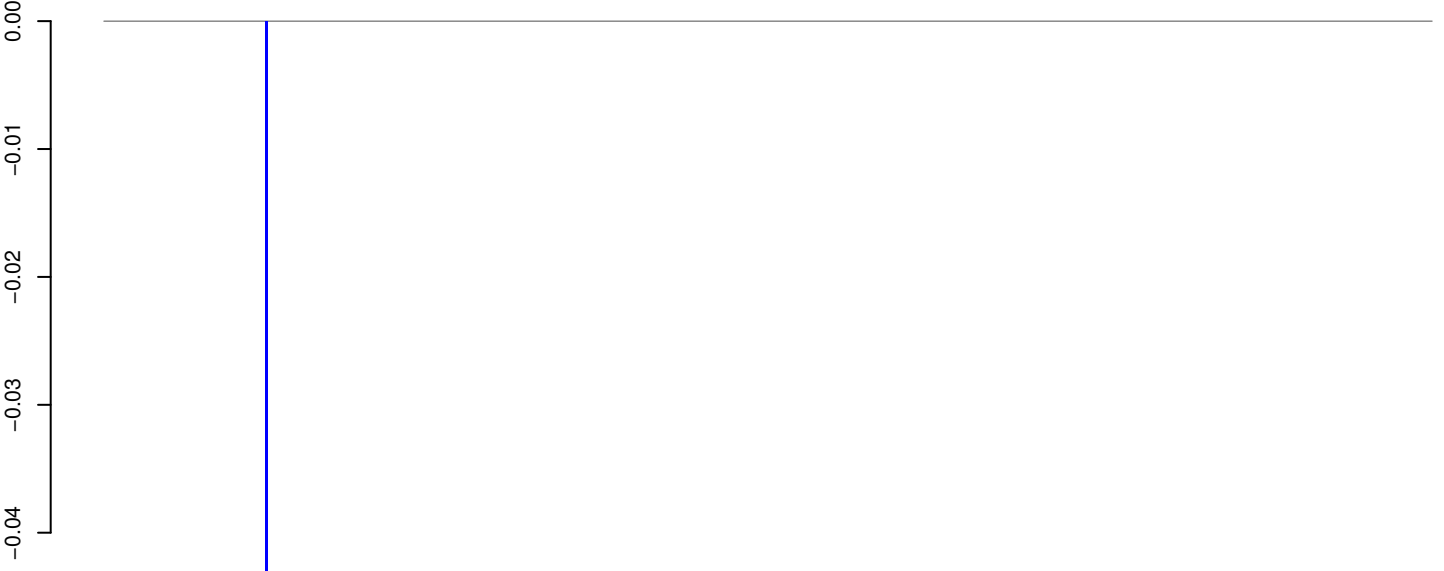
0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_dsLuc.18_23.rep



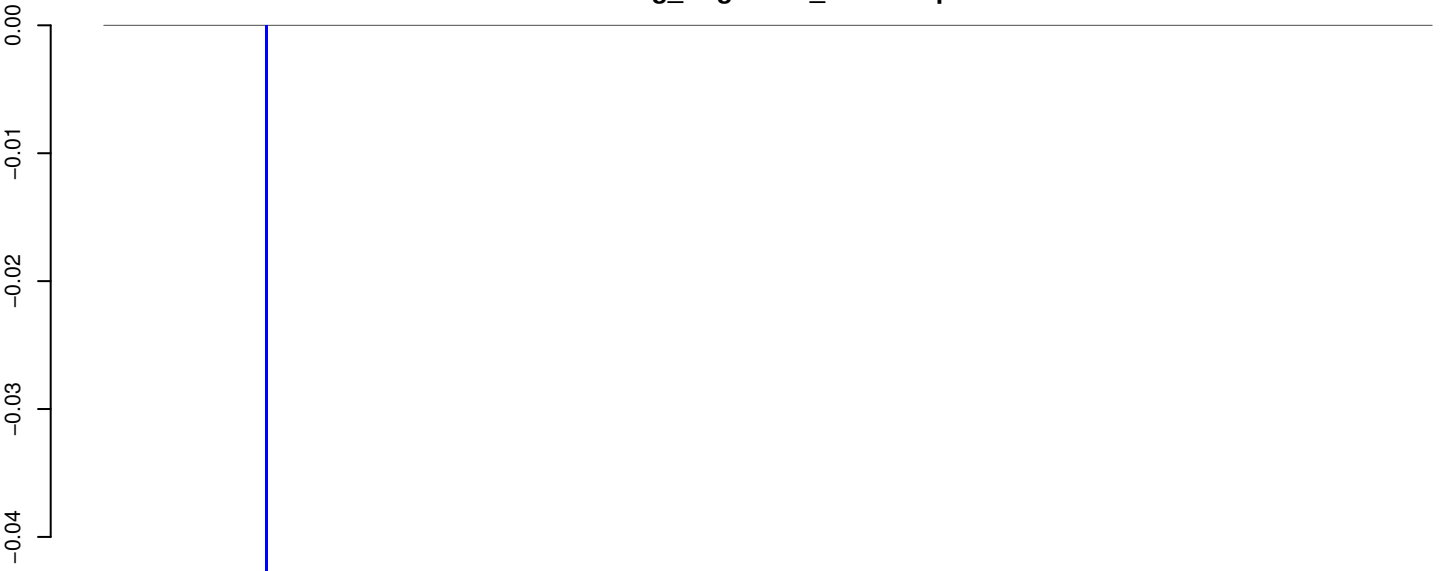
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep

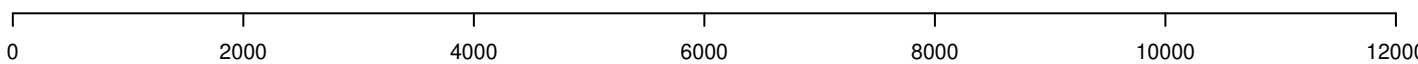


positive=0, negative=0, total=0

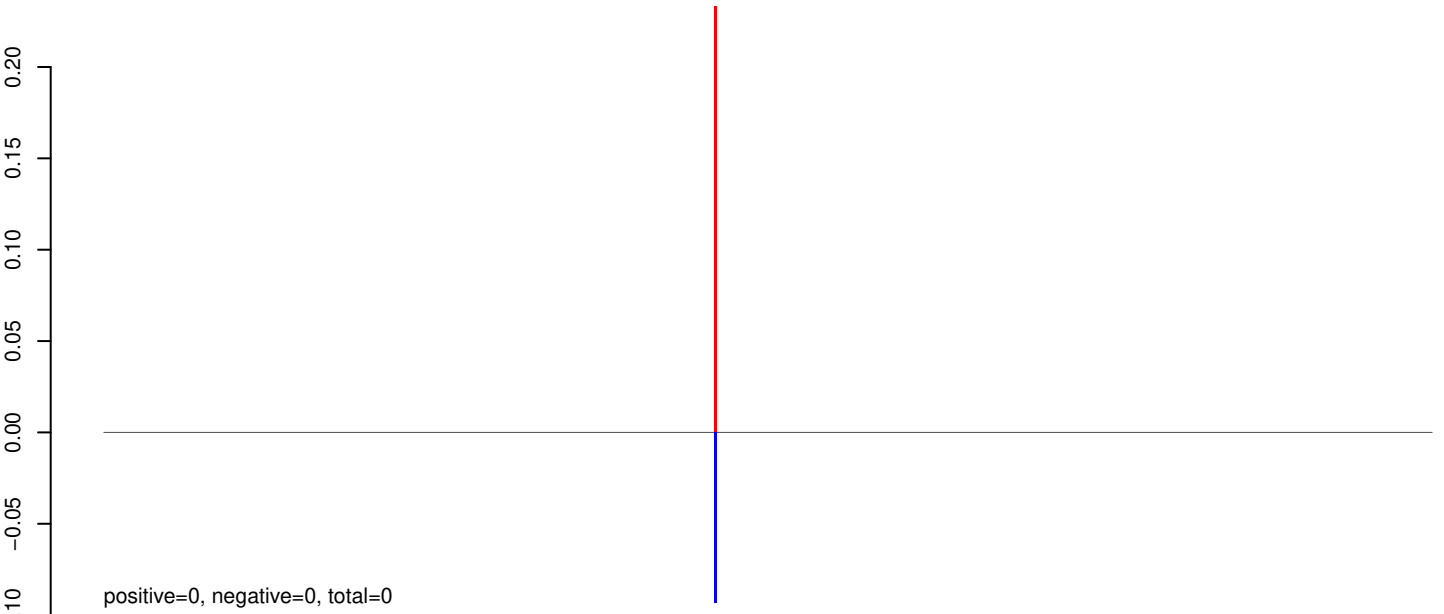
AeAeg_Aag2SINV_dsLuc.rep



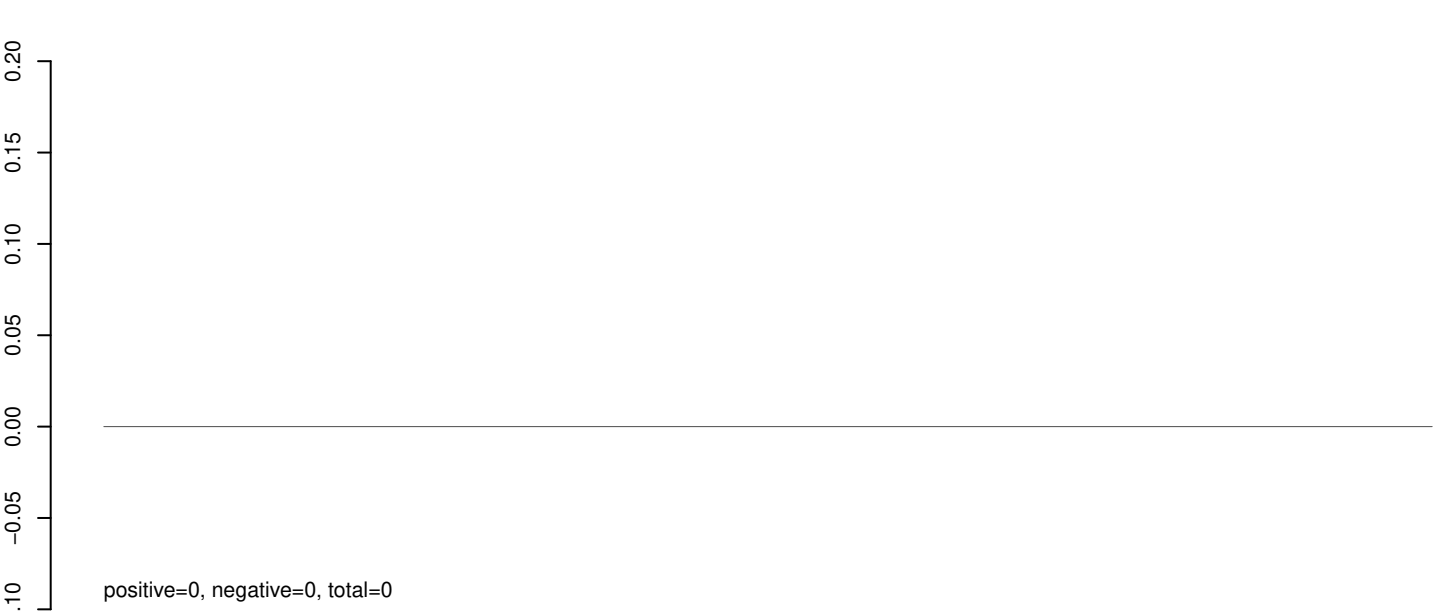
positive=0, negative=0, total=0 window size=25, length=11503, virus@NC_035132.1_Culex_rhabdo-like_virus_strain_mosWSB71420:1-11503



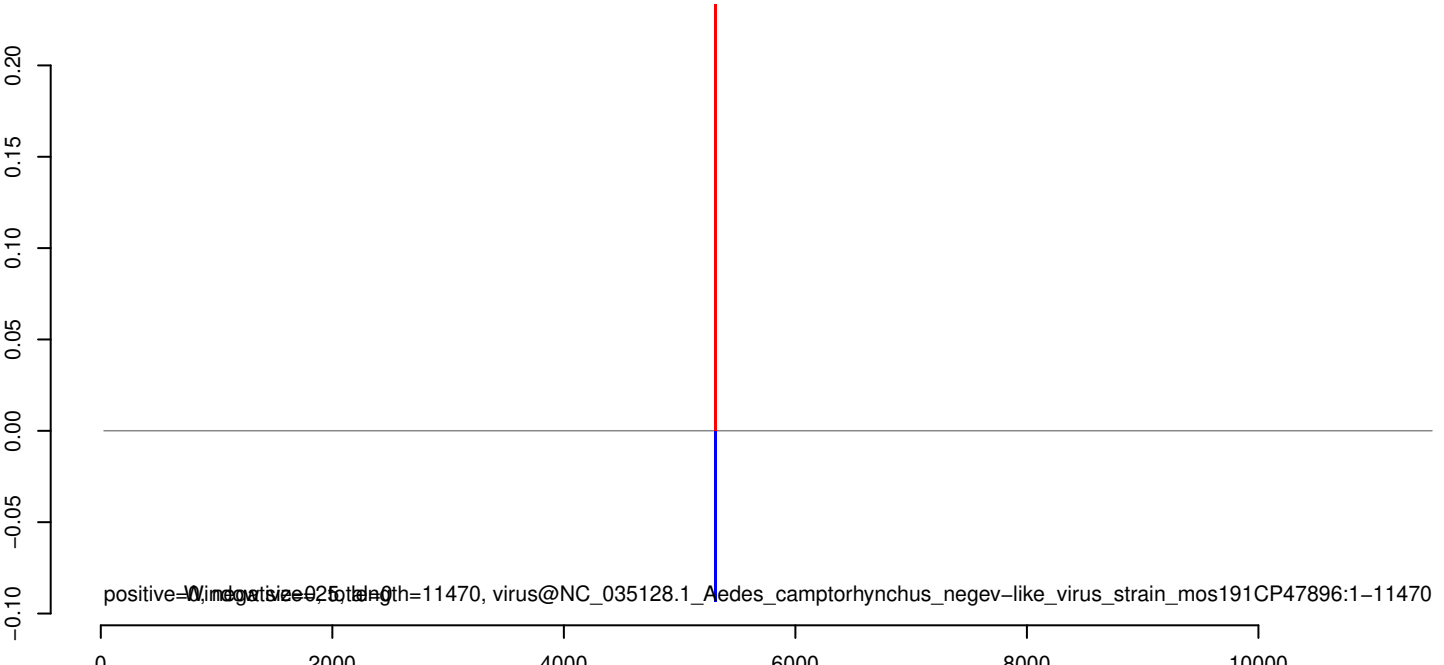
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



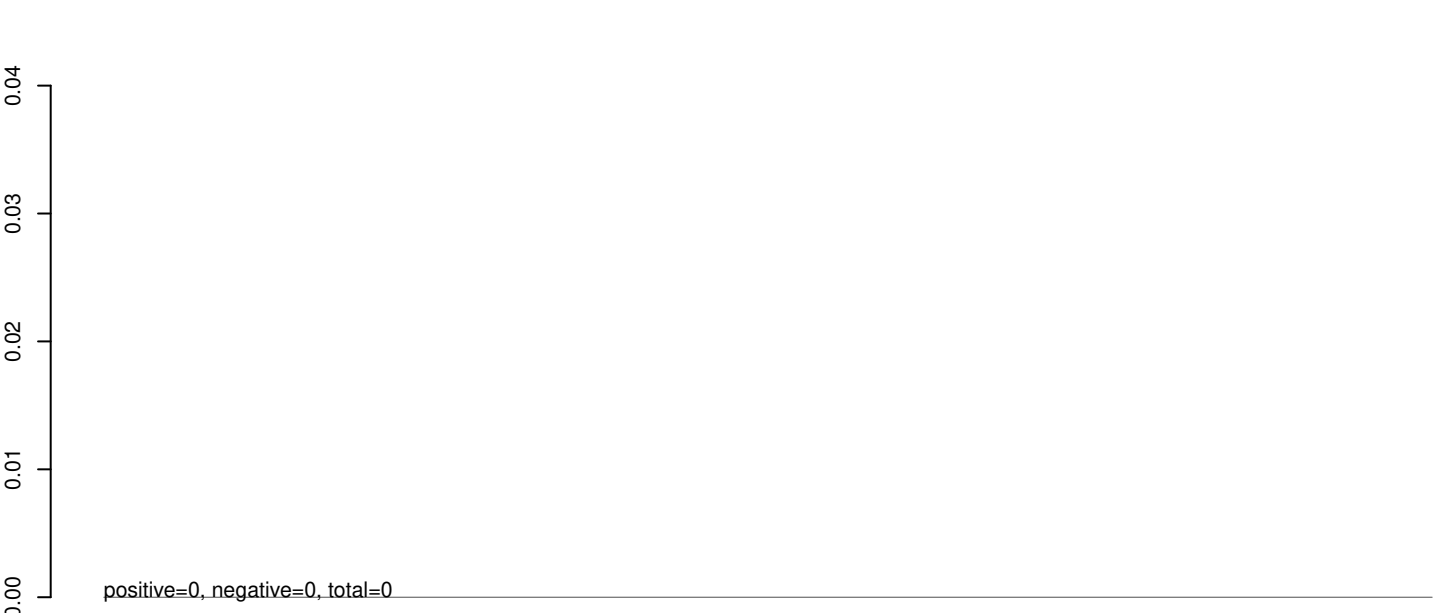
AeAeg_Aag2SINV_dsLuc.rep



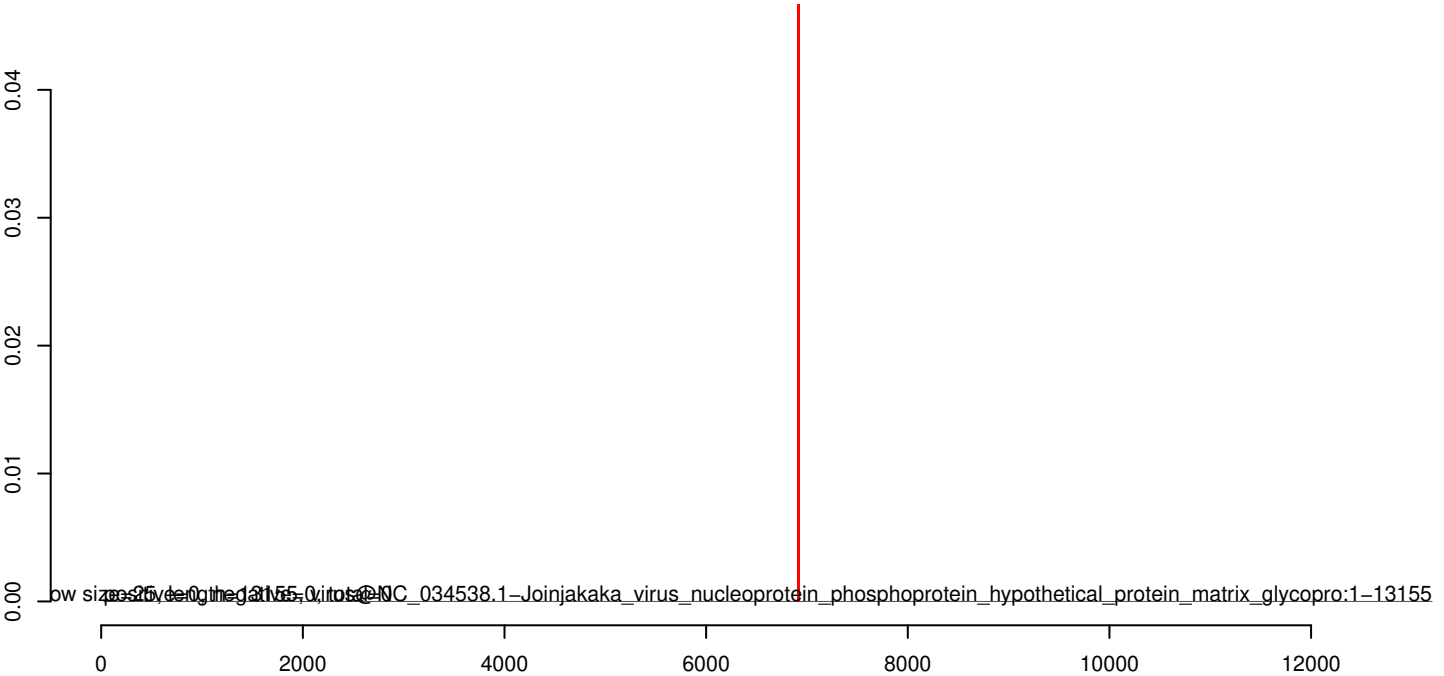
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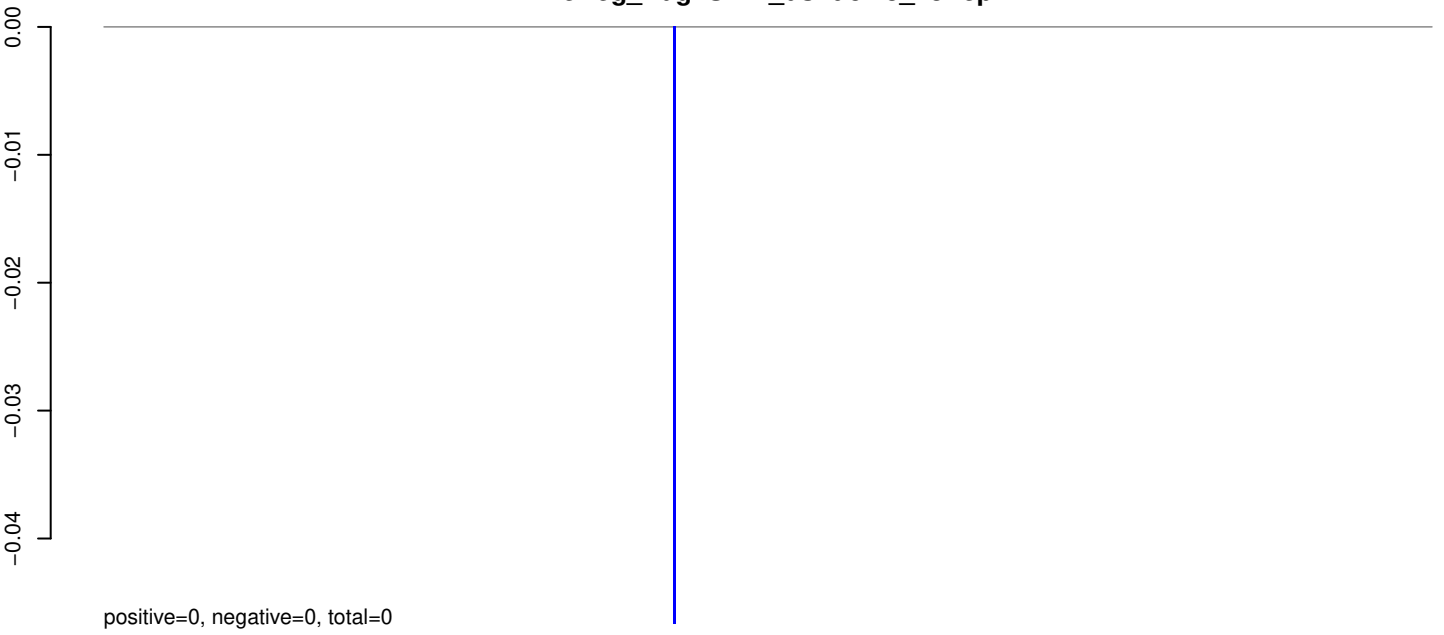
AeAeg_Aag2SINV_dsLuc.24_35.rep



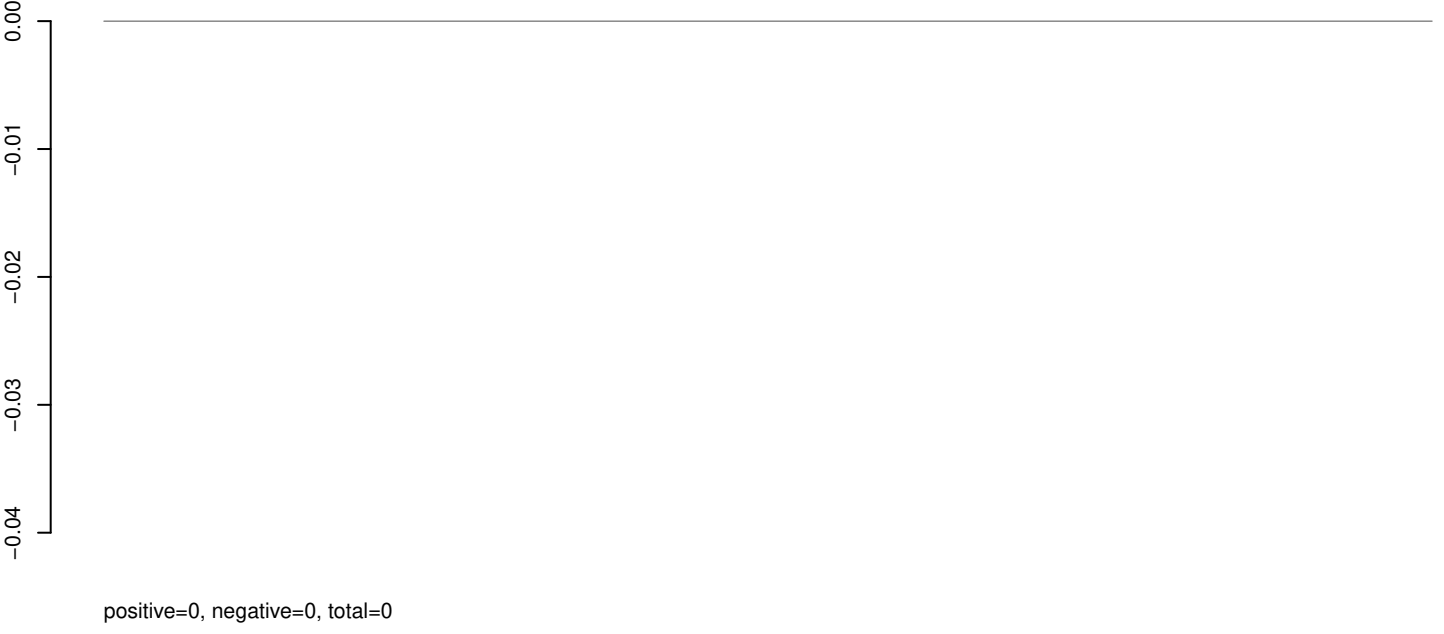
AeAeg_Aag2SINV_dsLuc.rep



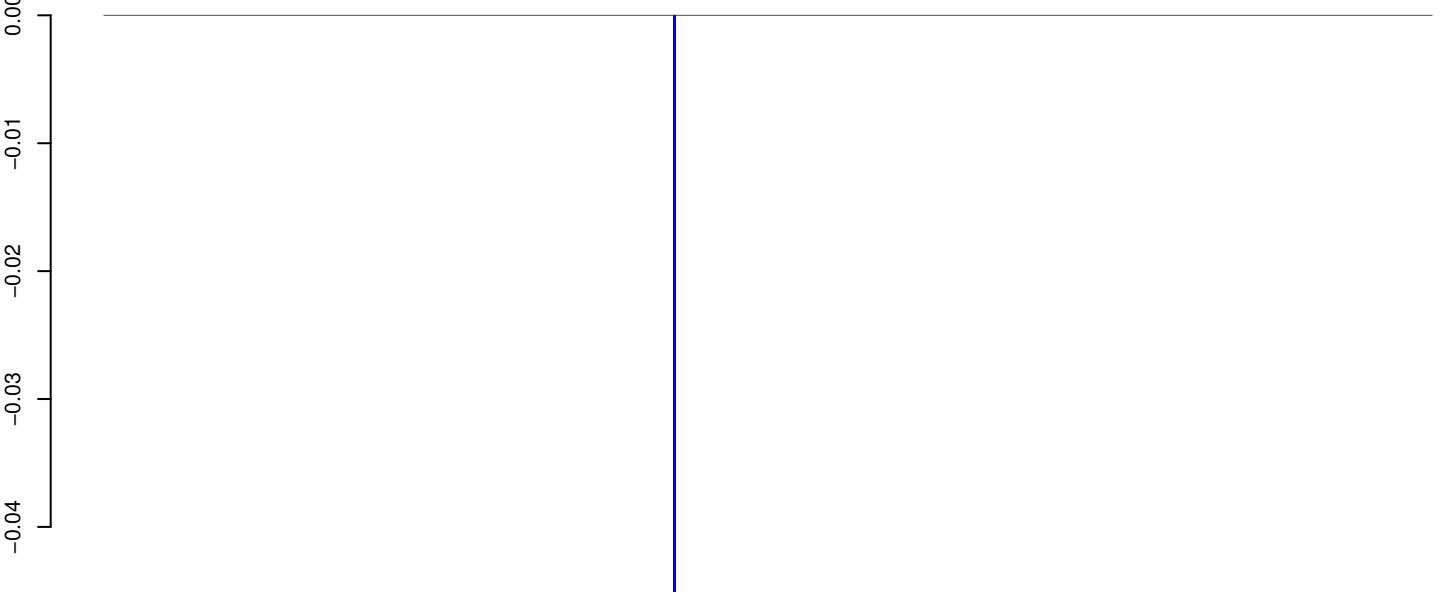
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



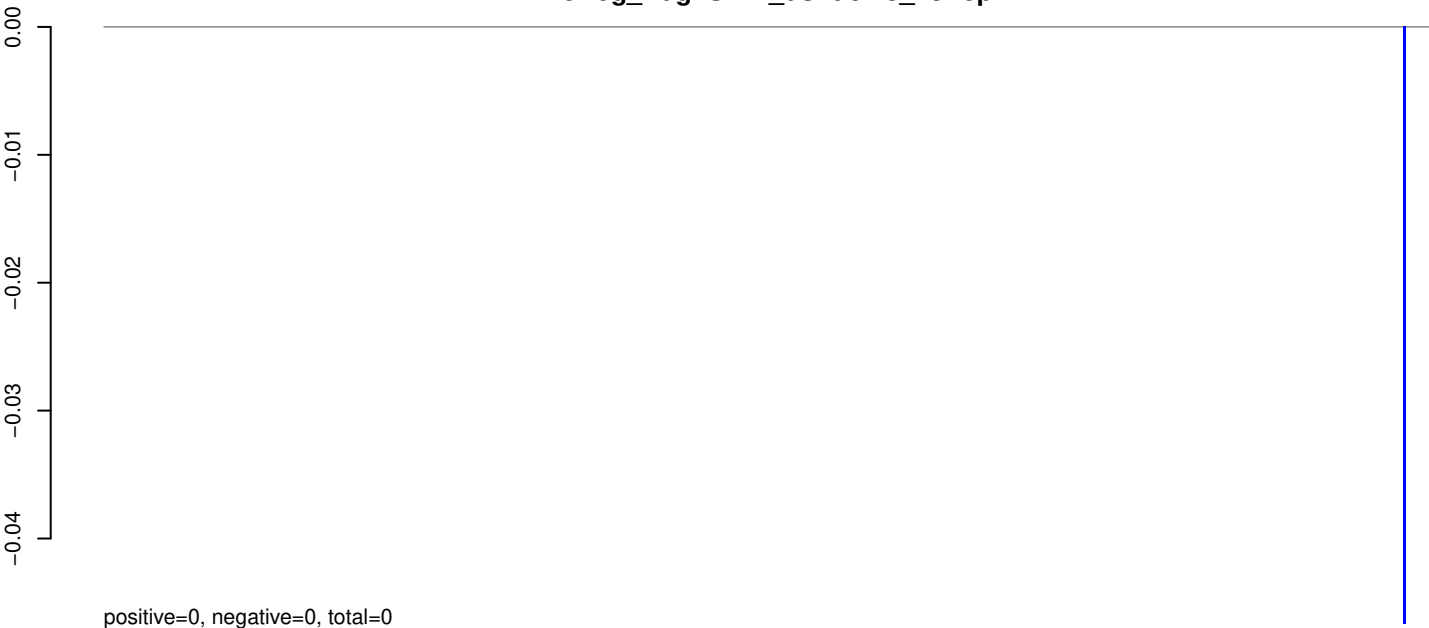
AeAeg_Aag2SINV_dsLuc.rep



00034534.1-Rochambeau_virus_nucleoprotein_phosphoprotein_matrix_hypothetical_proteins_glycopr:1-13593

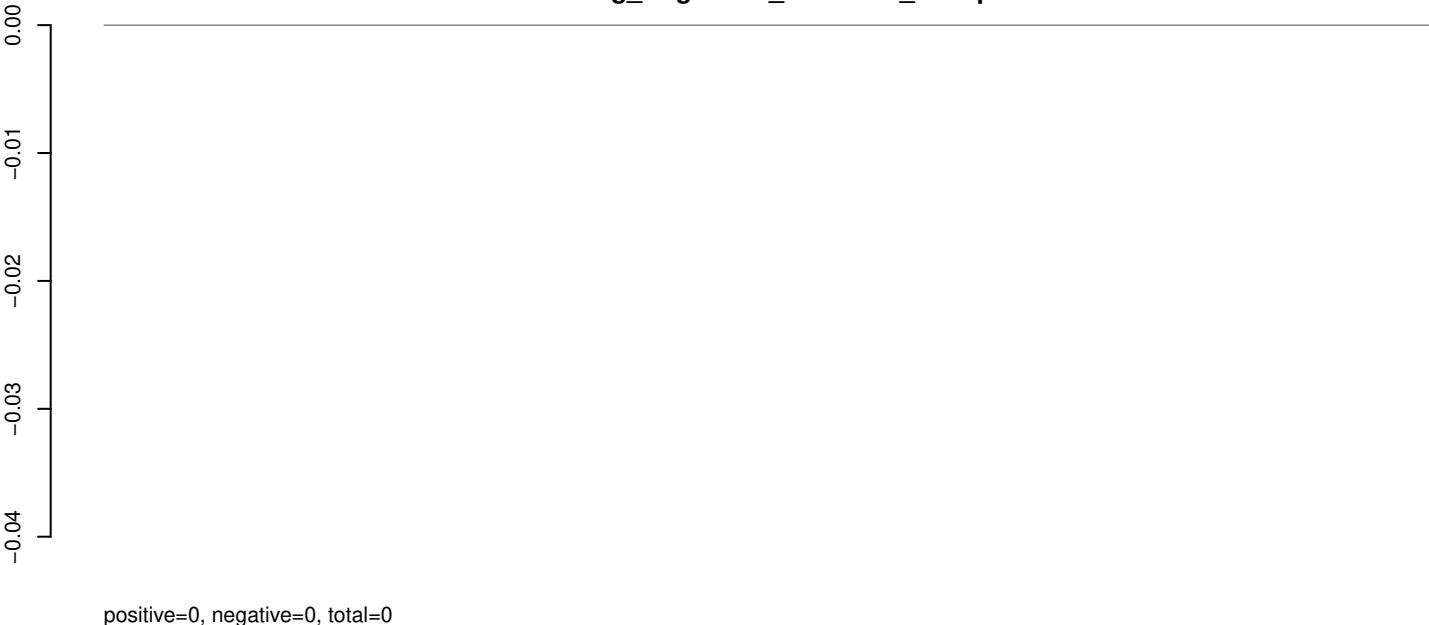
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AeAeg_Aag2SINV_dsLuc.18_23.rep



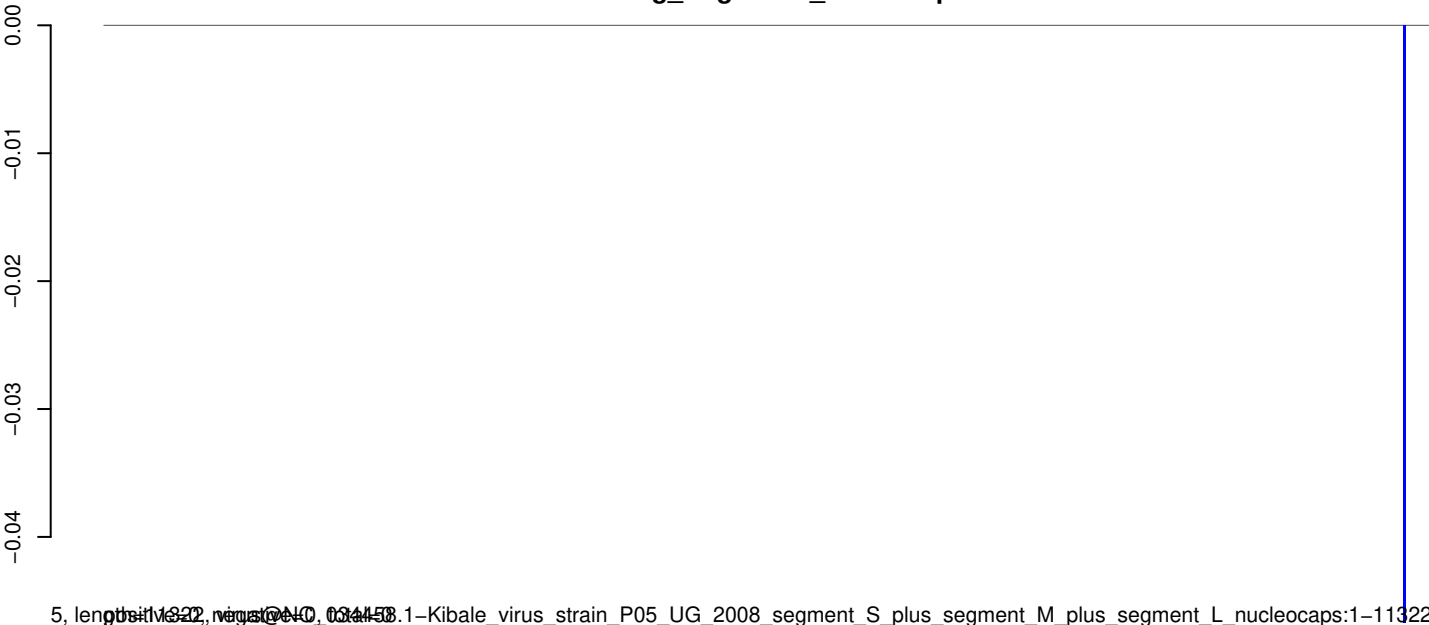
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=0, negative=0, total=0

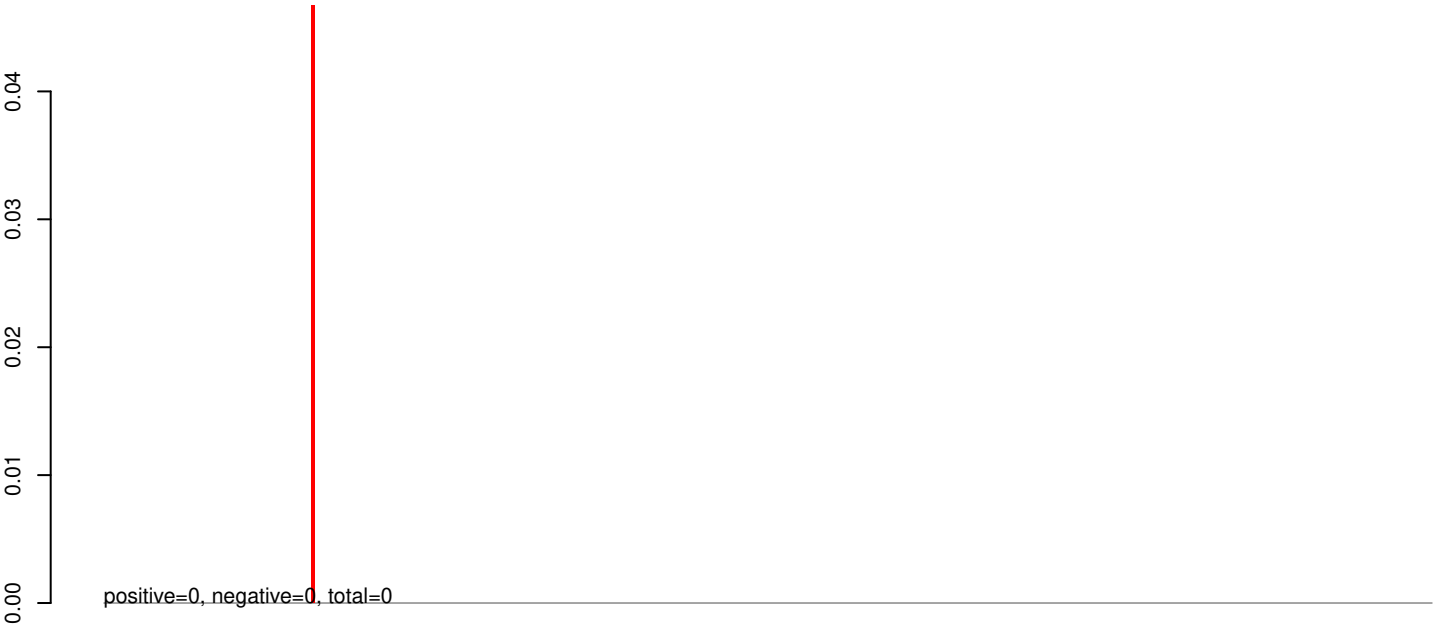
AeAeg_Aag2SINV_dsLuc.rep



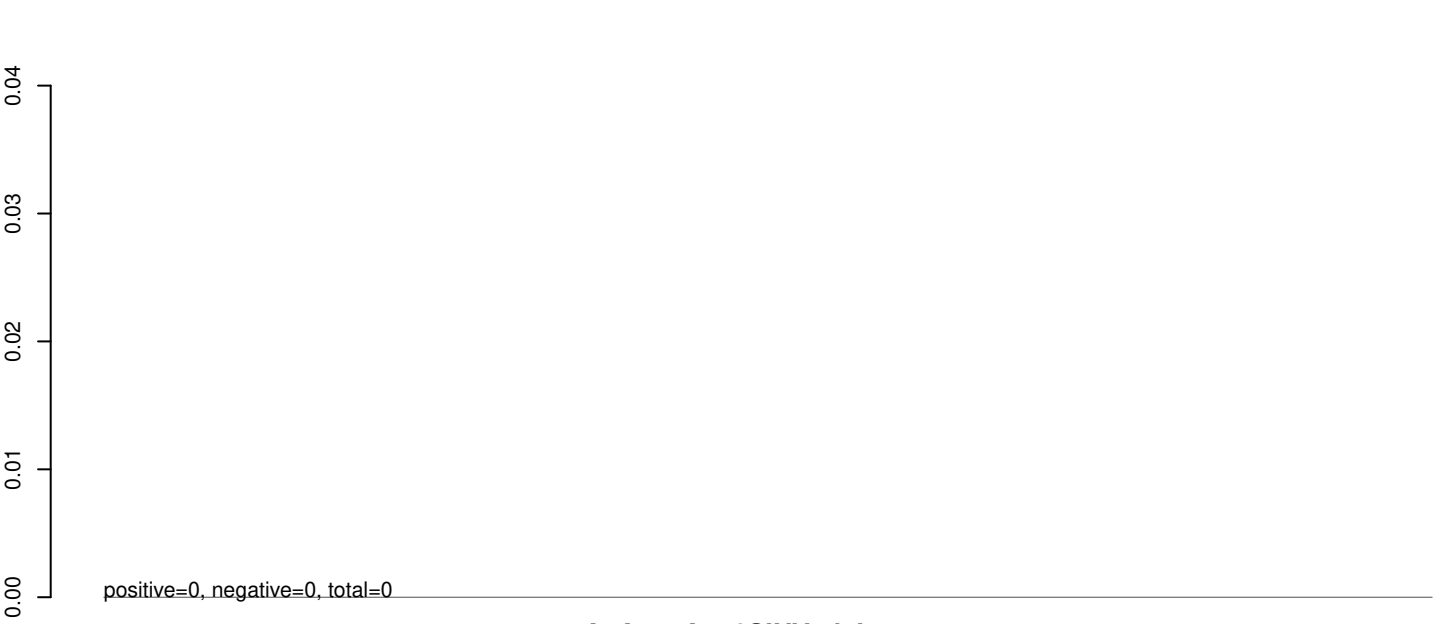
5, length=11322, negative=0, total=0 1-Kibale_virus_strain_P05_UG_2008_segment_S_plus_segment_M_plus_segment_L_nucleocaps:1-11322

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_dsLuc.18_23.rep



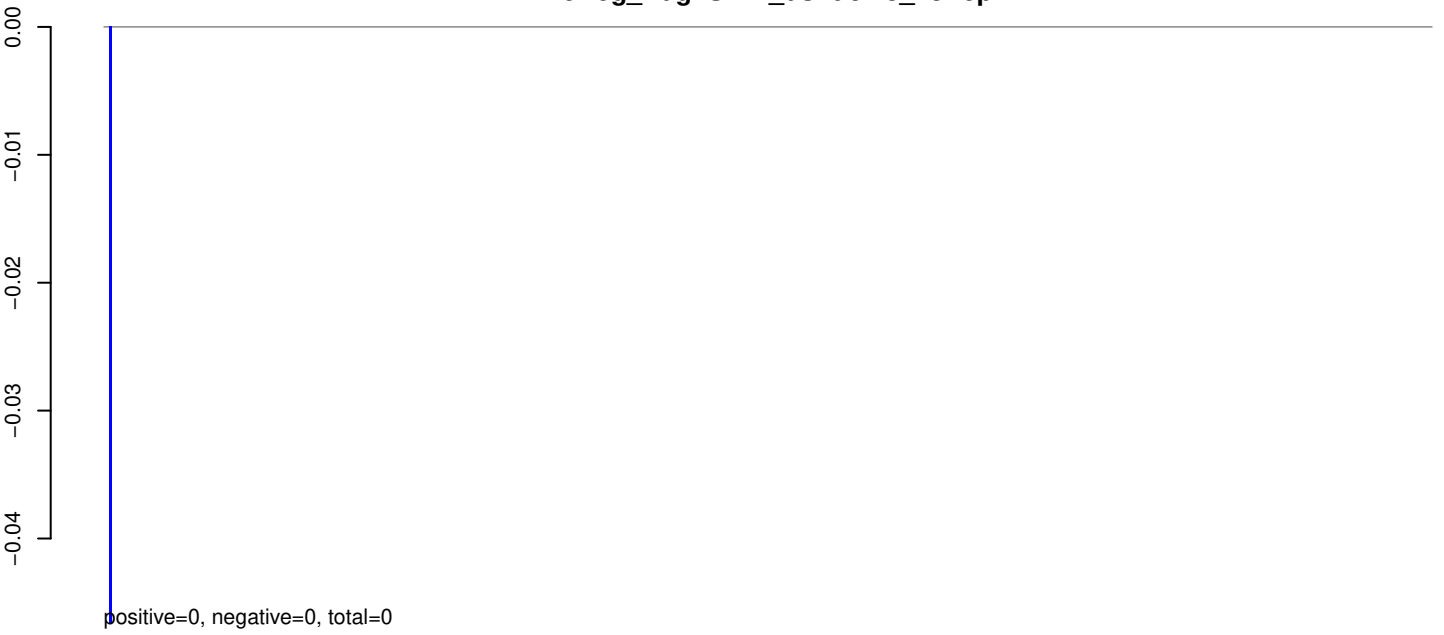
AeAeg_Aag2SINV_dsLuc.24_35.rep



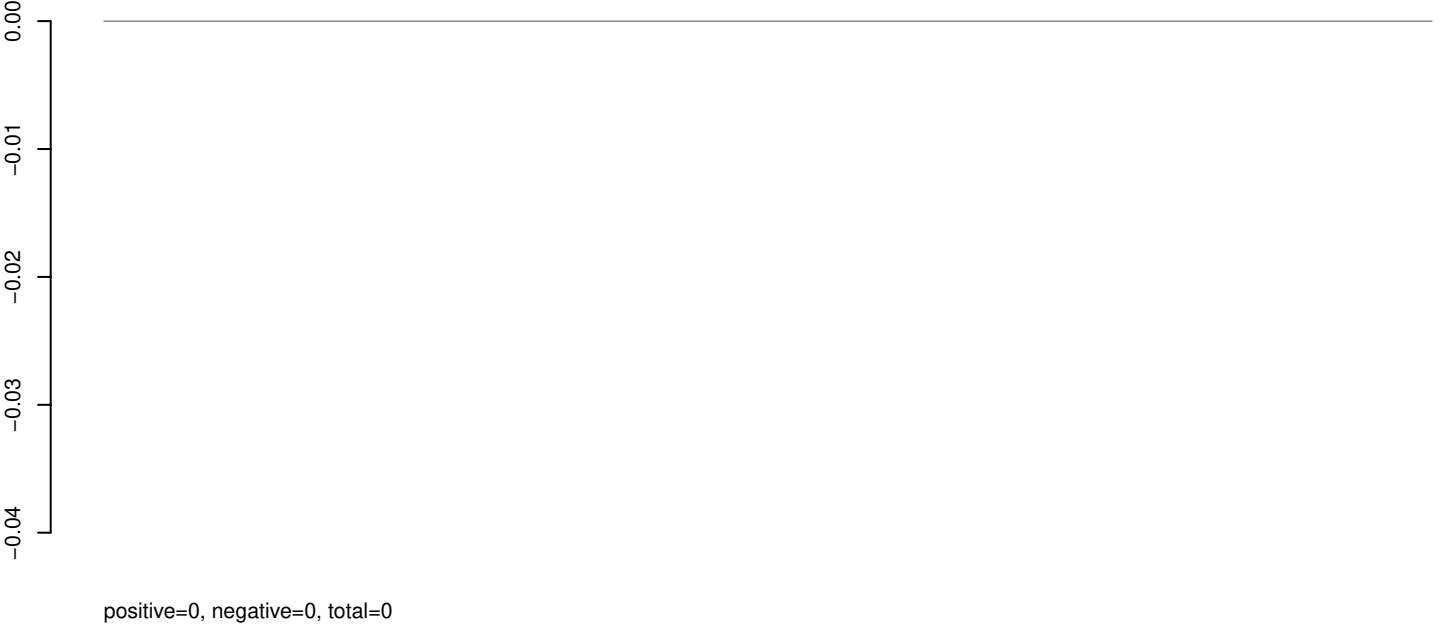
AeAeg_Aag2SINV_dsLuc.rep



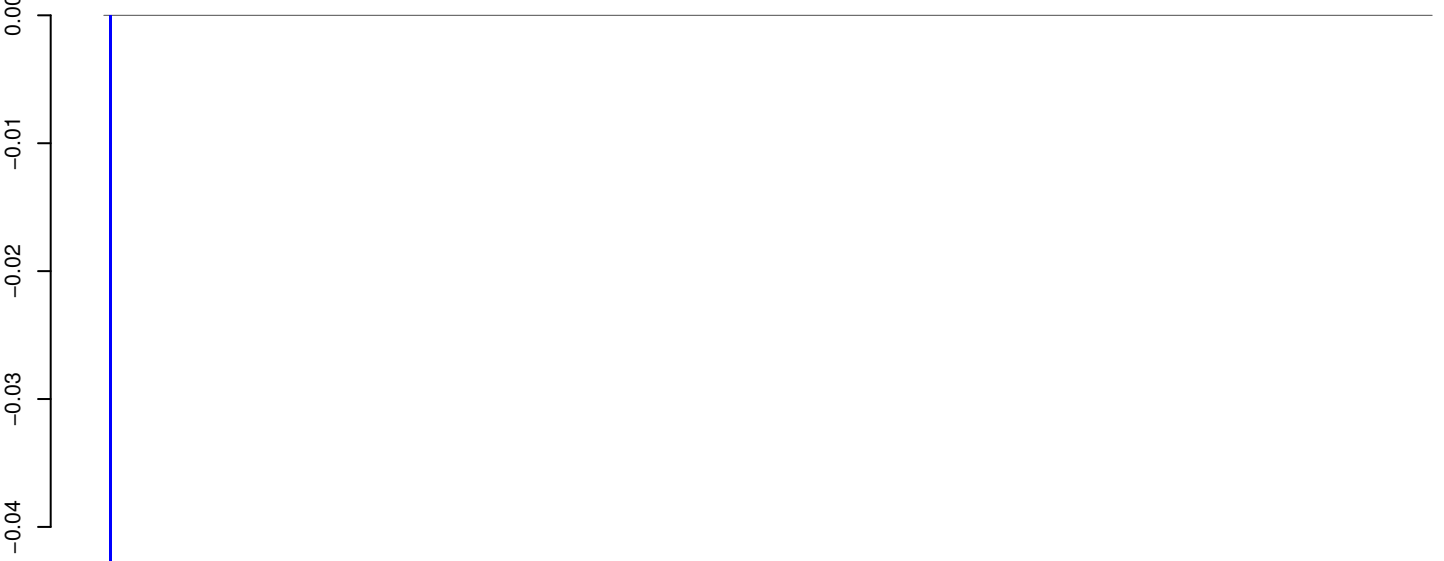
AeAeg_Aag2SINV_dsLuc.18_23.rep



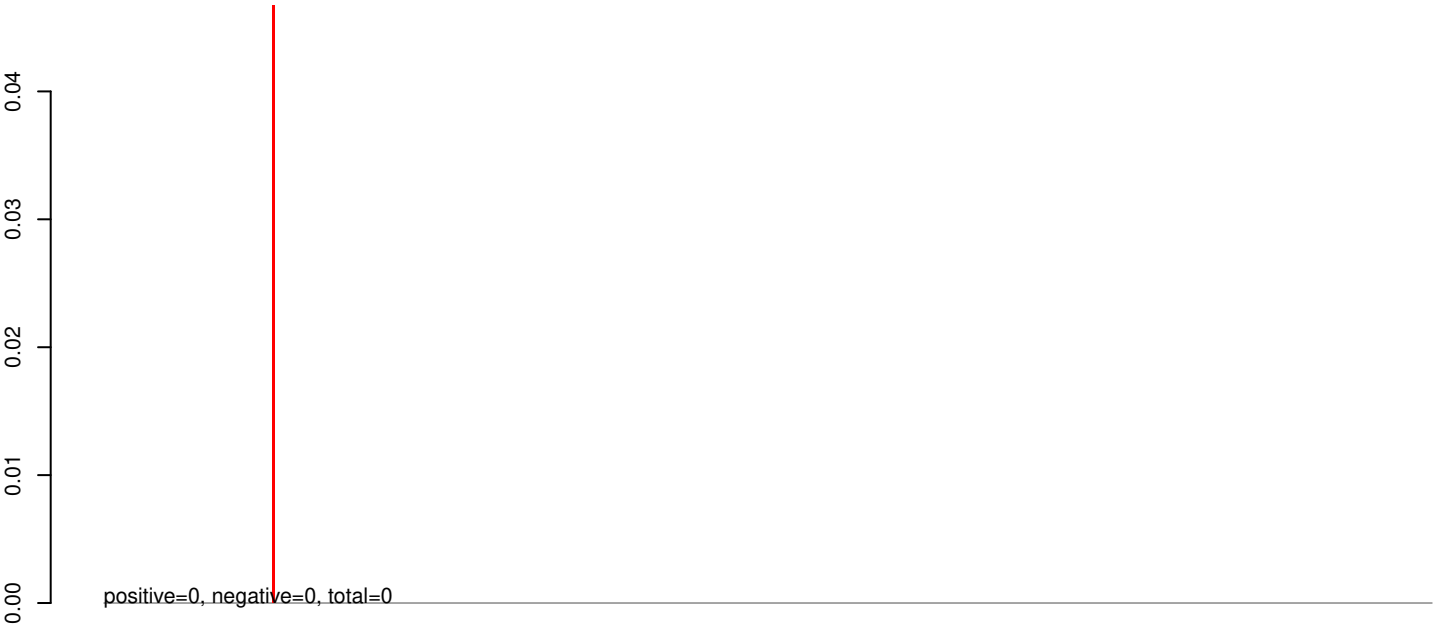
AeAeg_Aag2SINV_dsLuc.24_35.rep



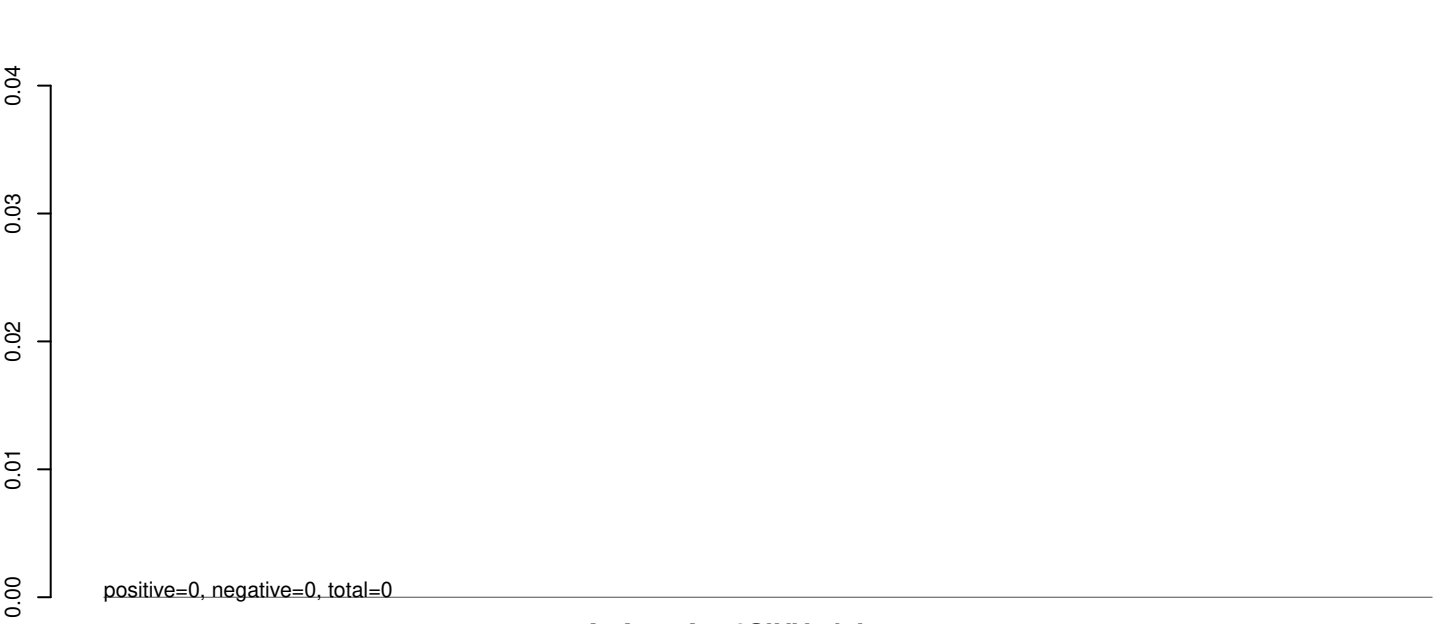
AeAeg_Aag2SINV_dsLuc.rep



AeAeg_Aag2SINV_dsLuc.18_23.rep



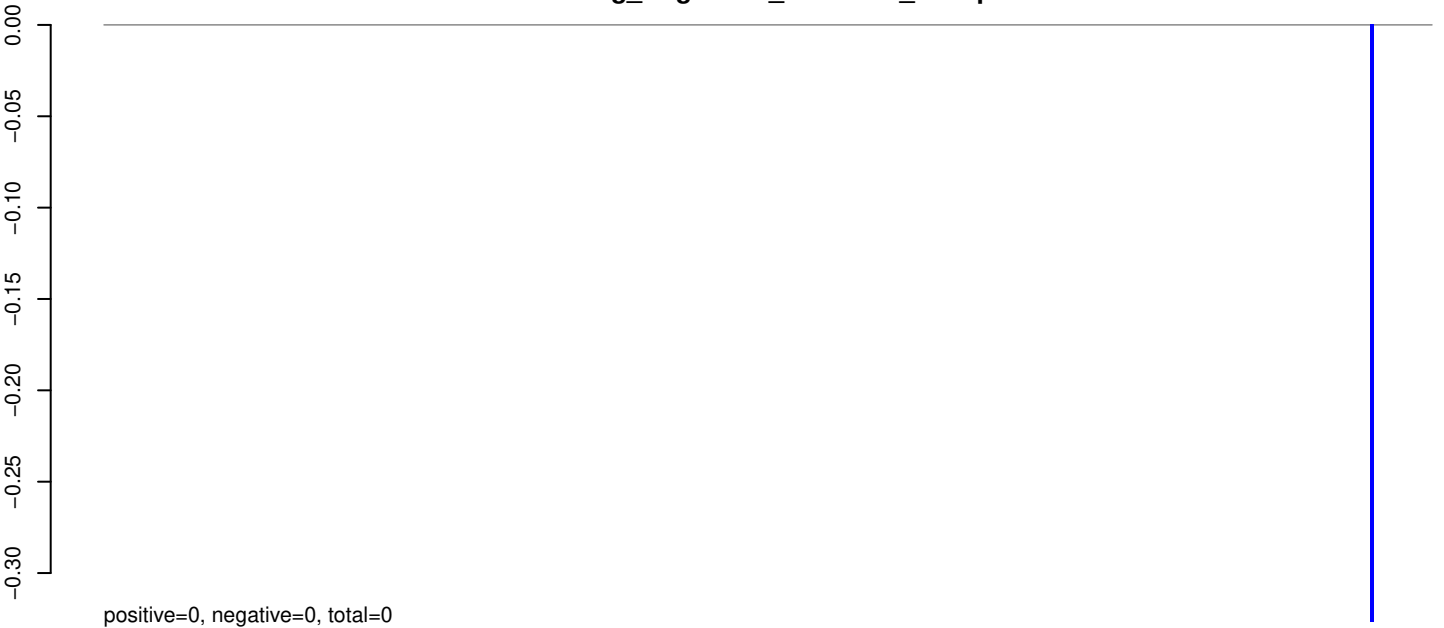
AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep

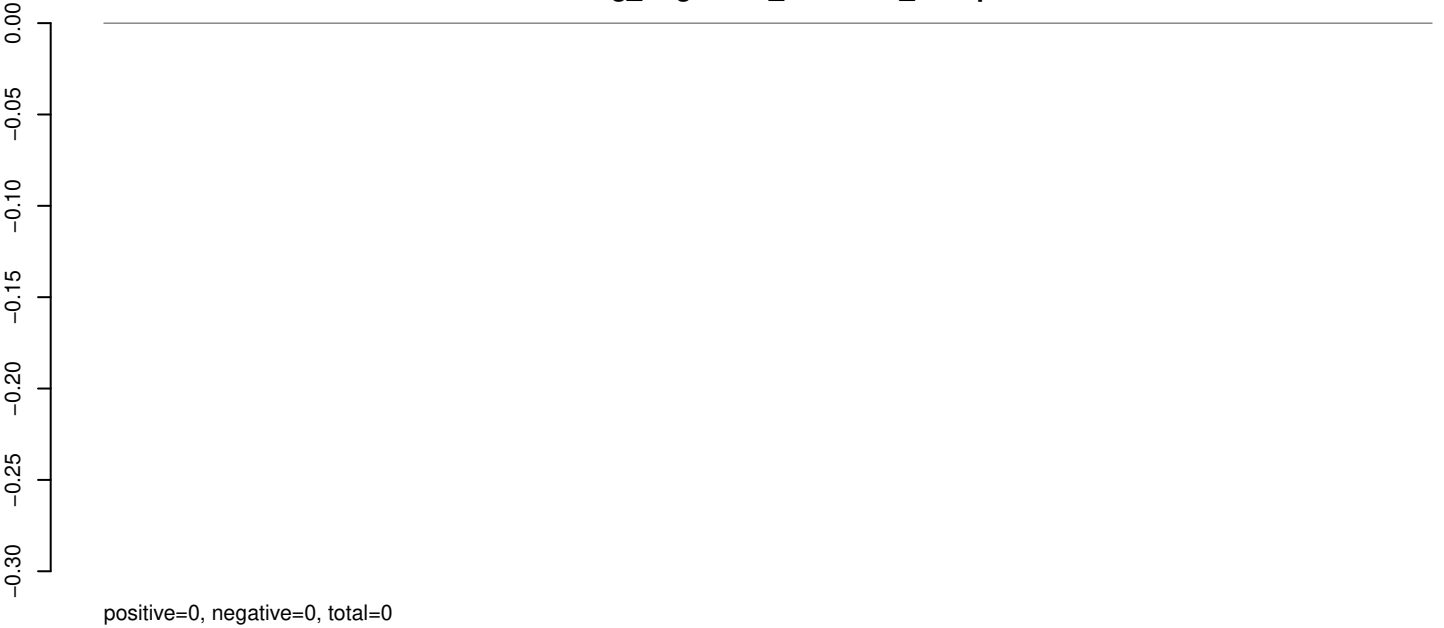


AeAeg_Aag2SINV_dsLuc.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=0, negative=0, total=0

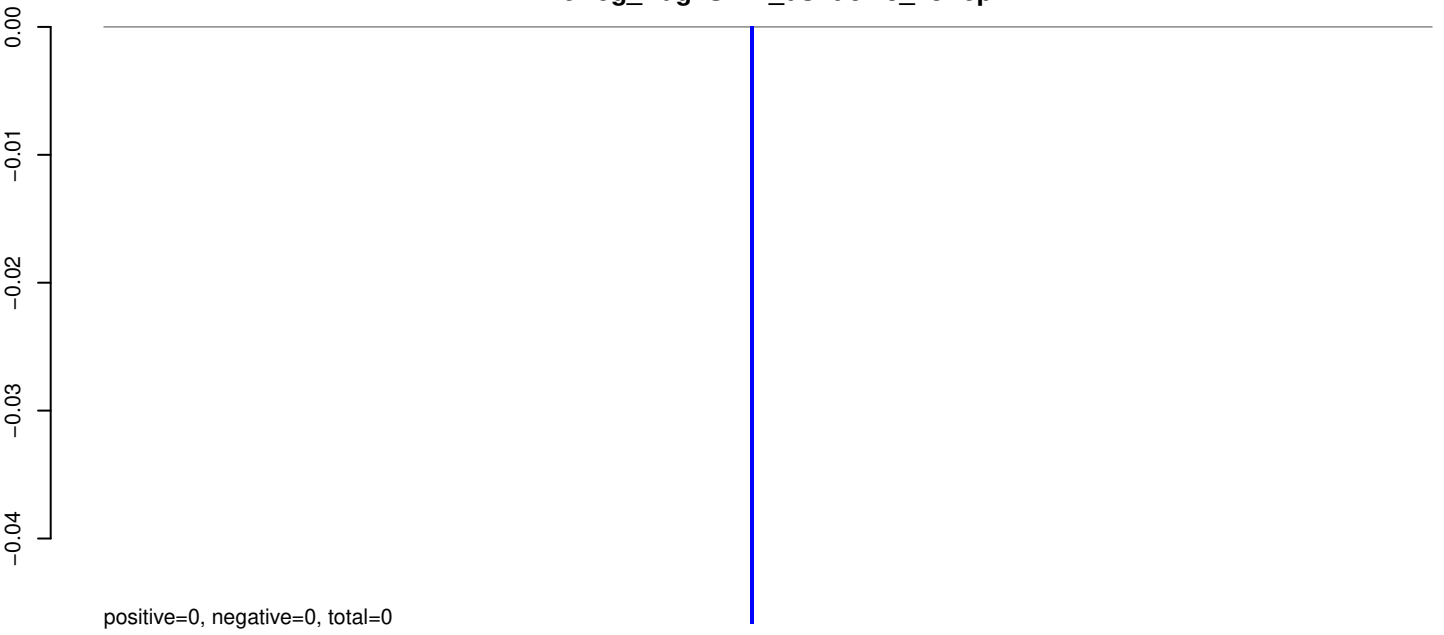
AeAeg_Aag2SINV_dsLuc.rep



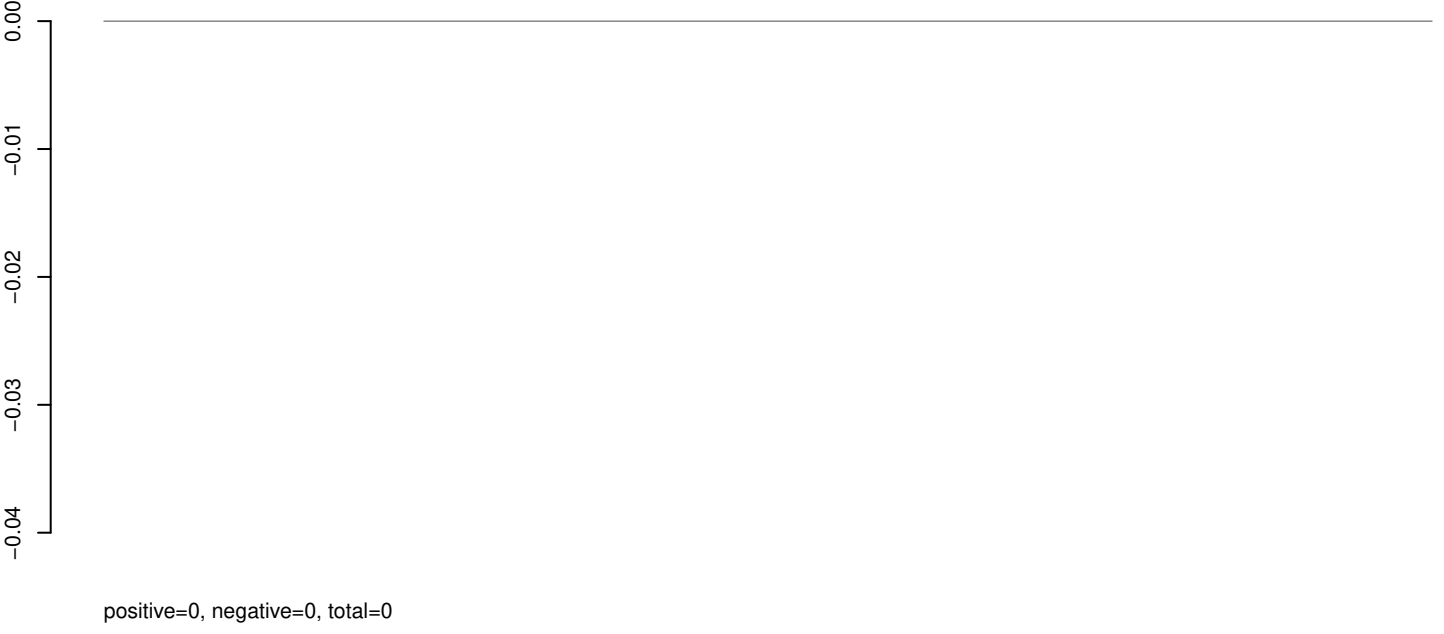
positive=0, negative=0, total=0, window-size=25, length=10251, virus@NC_033723.1-Gadgets_Gully_virus_from_Australia_polyprotein_gene:1-10251

0 2000 4000 6000 8000 10000

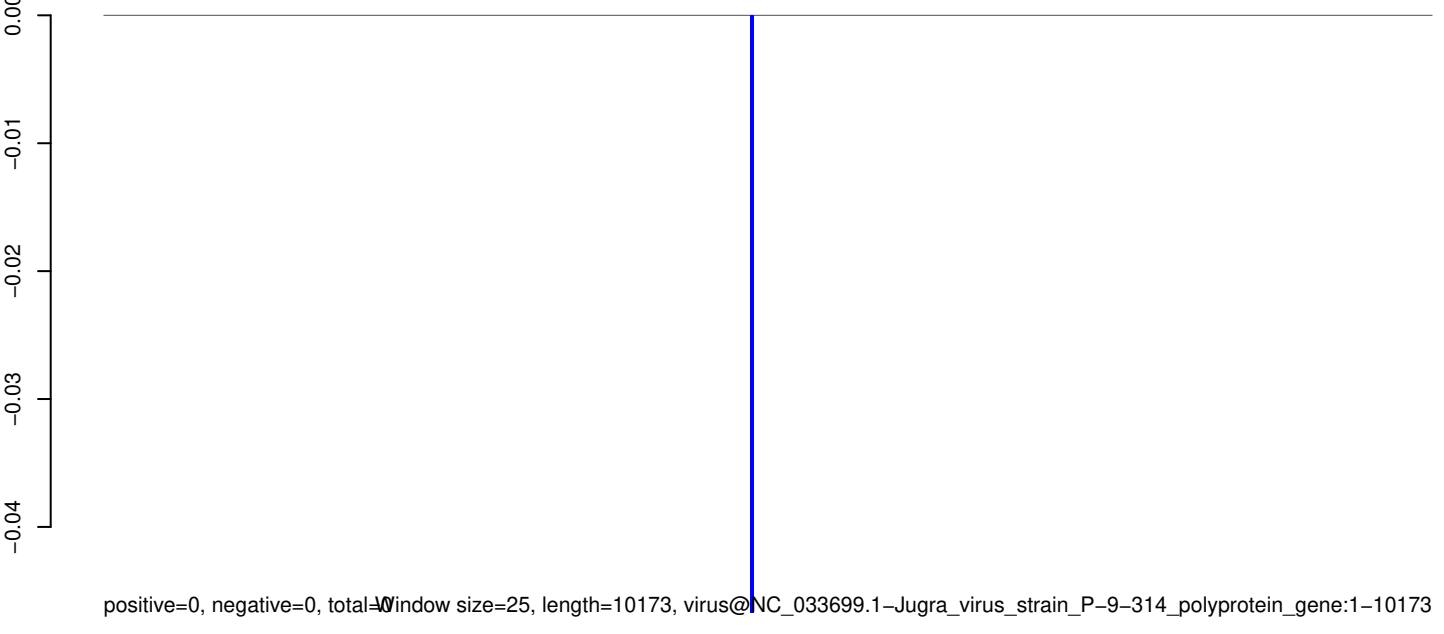
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



AeAeg_Aag2SINV_dsLuc.18_23.rep

0.00
-0.01
-0.02
-0.03
-0.04

positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep

0.00
-0.01
-0.02
-0.03
-0.04

positive=0, negative=0, total=0

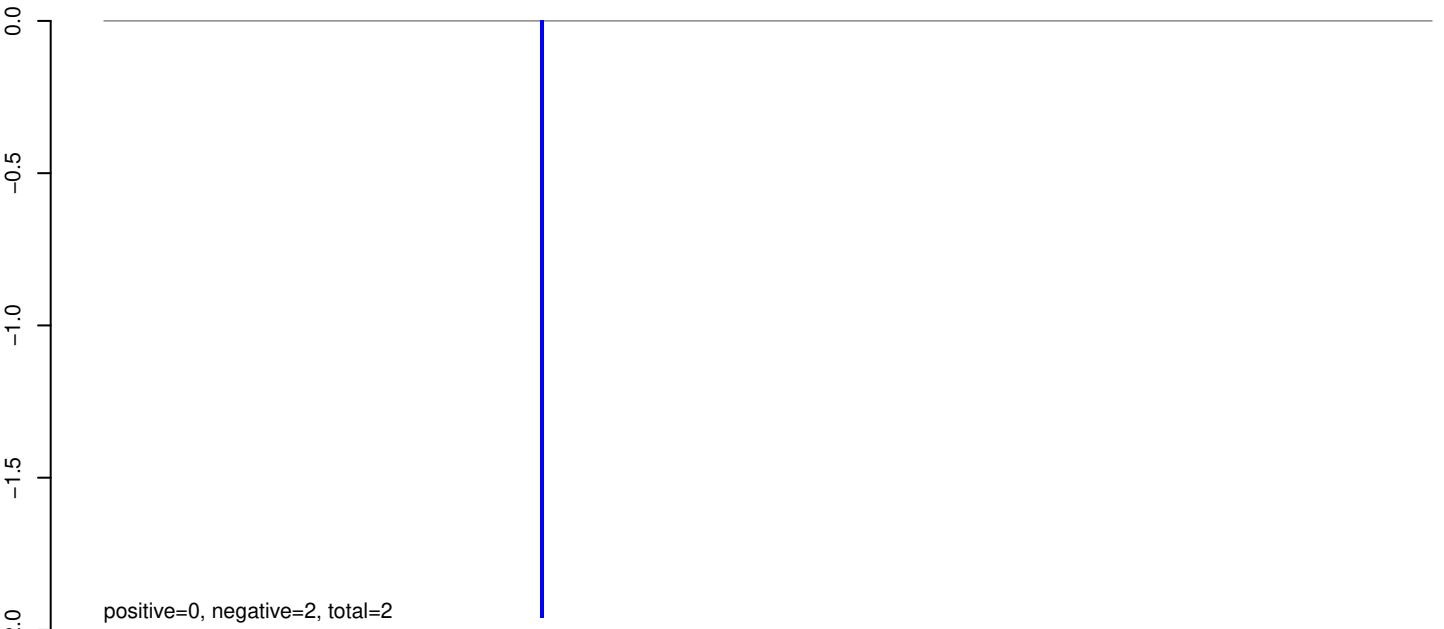
AeAeg_Aag2SINV_dsLuc.rep

0.00
-0.01
-0.02
-0.03
-0.04

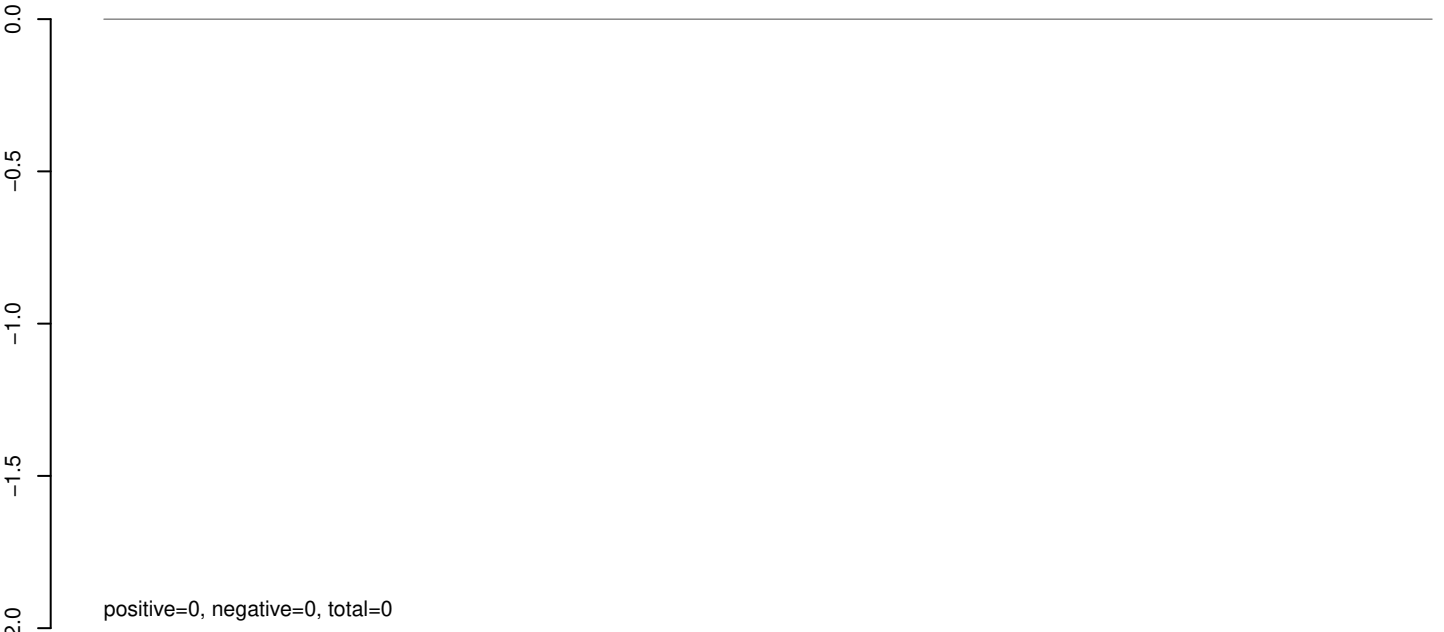
positive=0, negative=0, total=25, length=10173, virus@NC_033693.1-Bouboui_virus_strain_DAK_AR_B490_polyprotein_gene:1-10173

0 2000 4000 6000 8000 10000

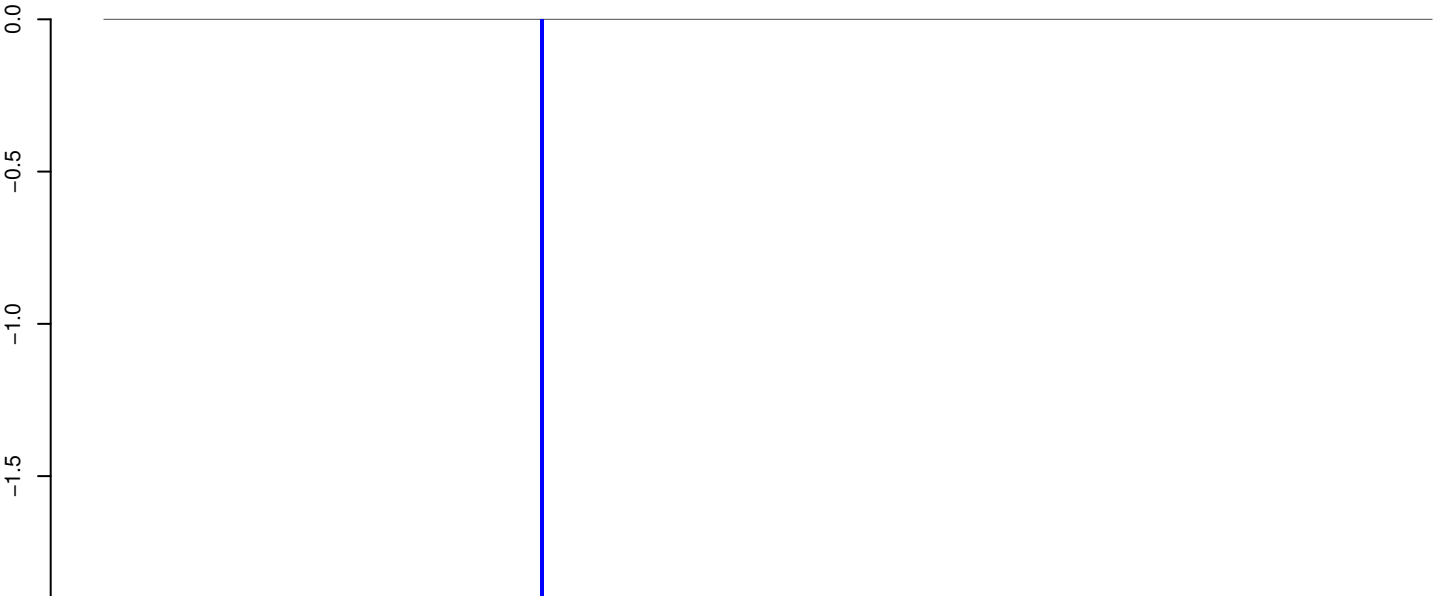
AeAeg_Aag2SINV_dsLuc.18_23.rep



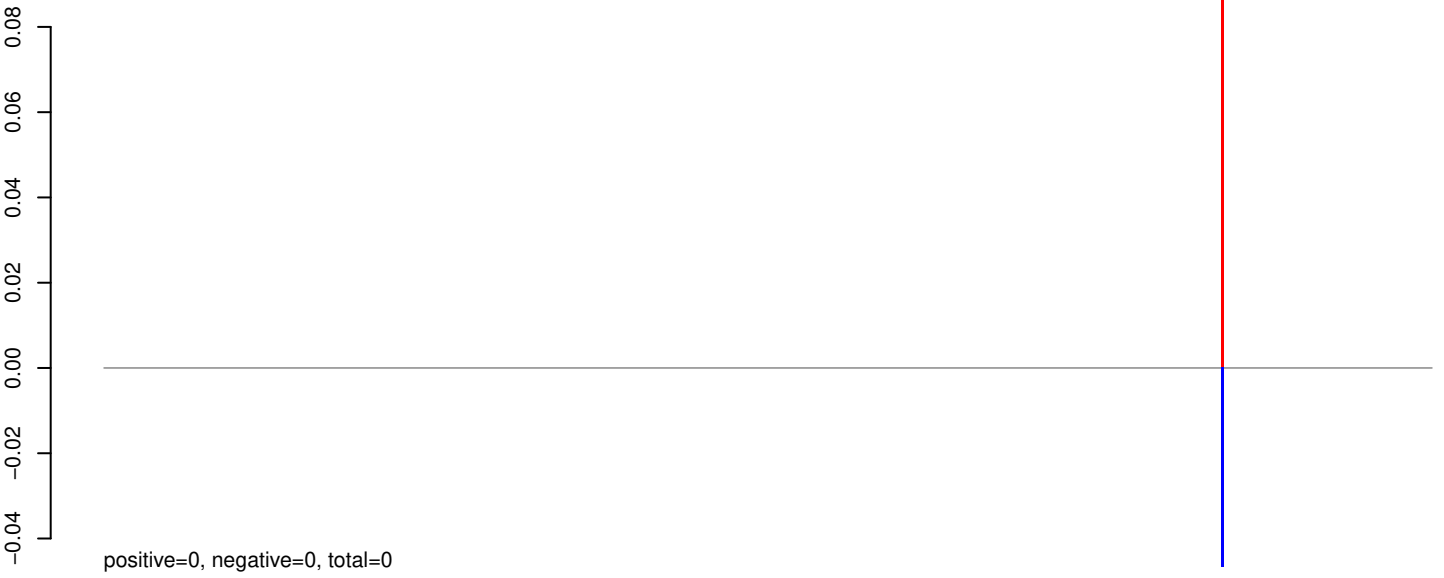
AeAeg_Aag2SINV_dsLuc.24_35.rep



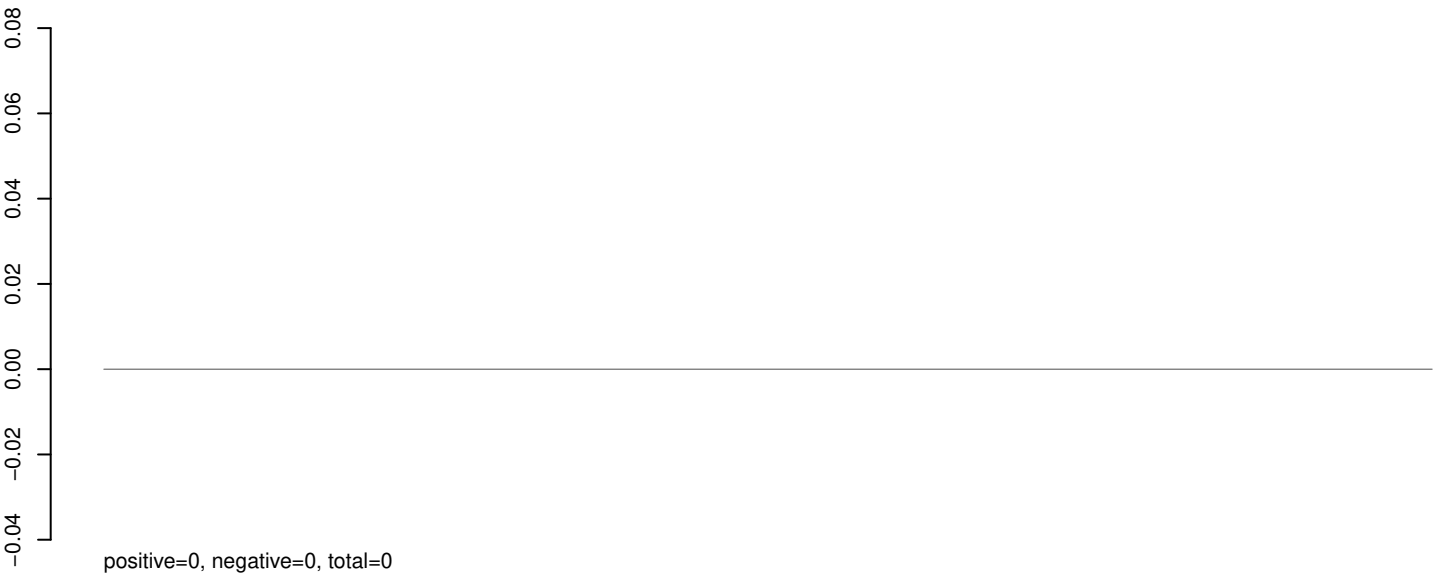
AeAeg_Aag2SINV_dsLuc.rep



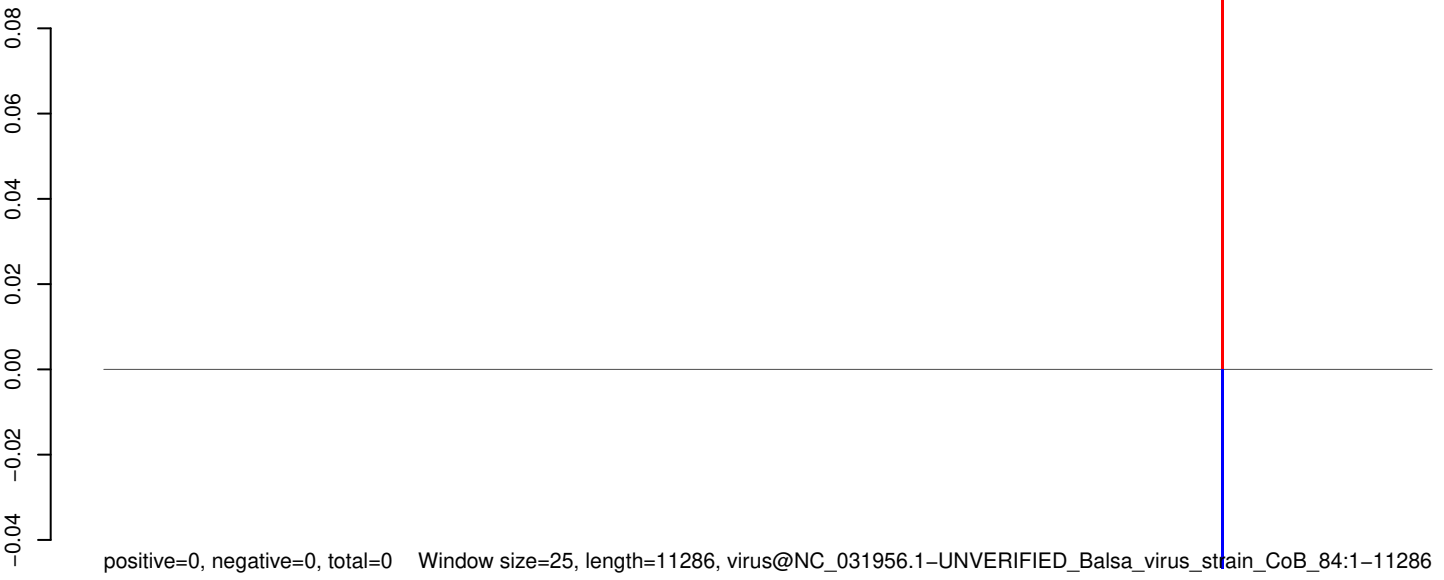
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



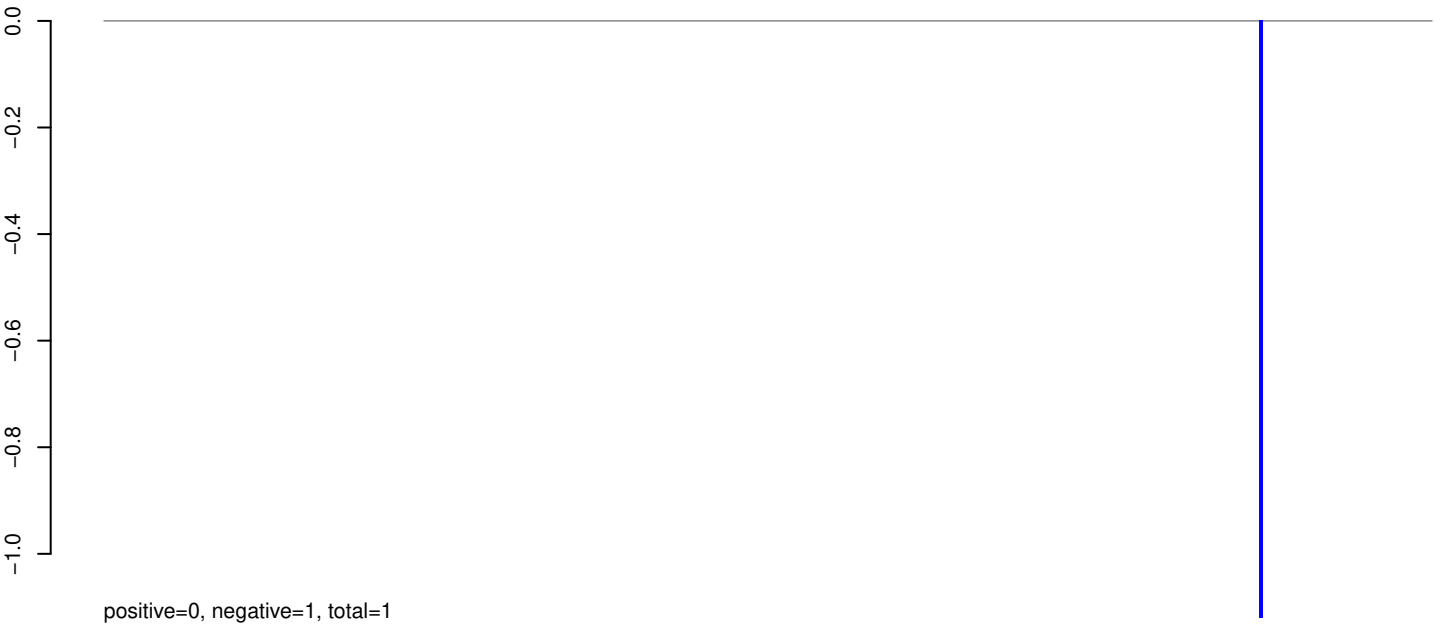
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Window size=25, length=11286, virus@NC_031956.1-UNVERIFIED_Balsa_virus_strain_CoB_84:1-11286

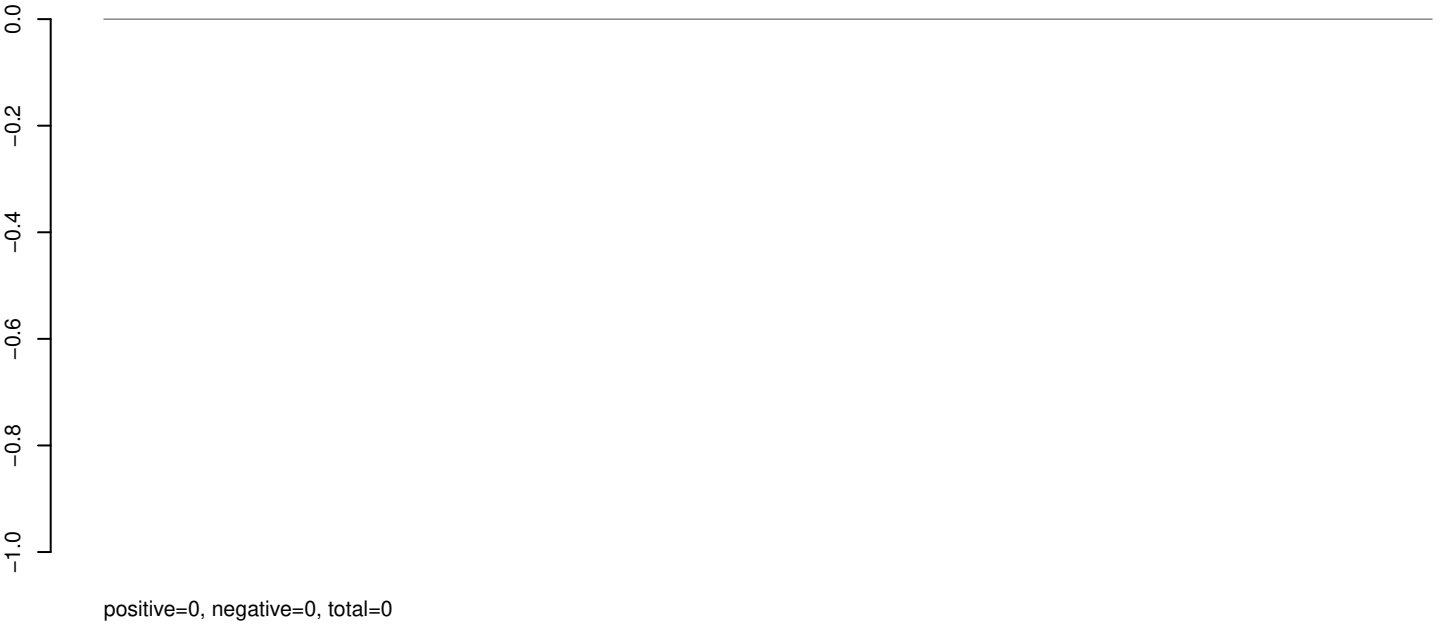
0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_dsLuc.18_23.rep



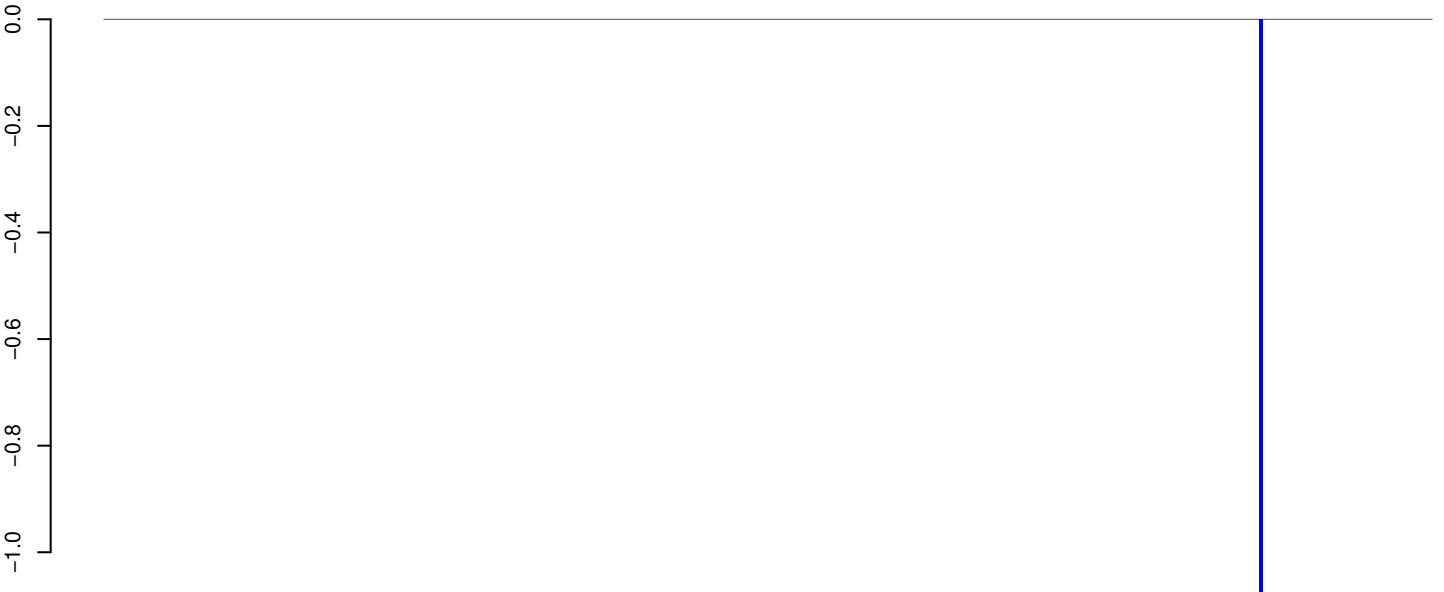
positive=0, negative=1, total=1

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.rep

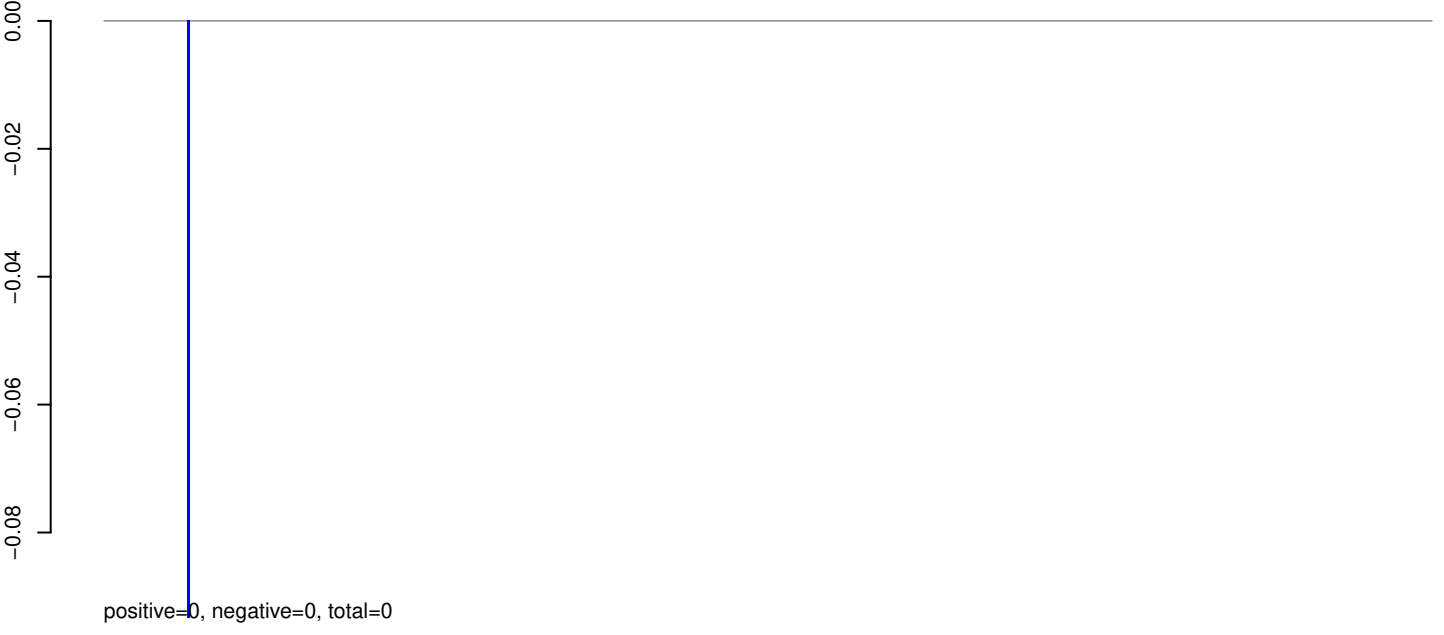


positive=0, negative=1, total=1

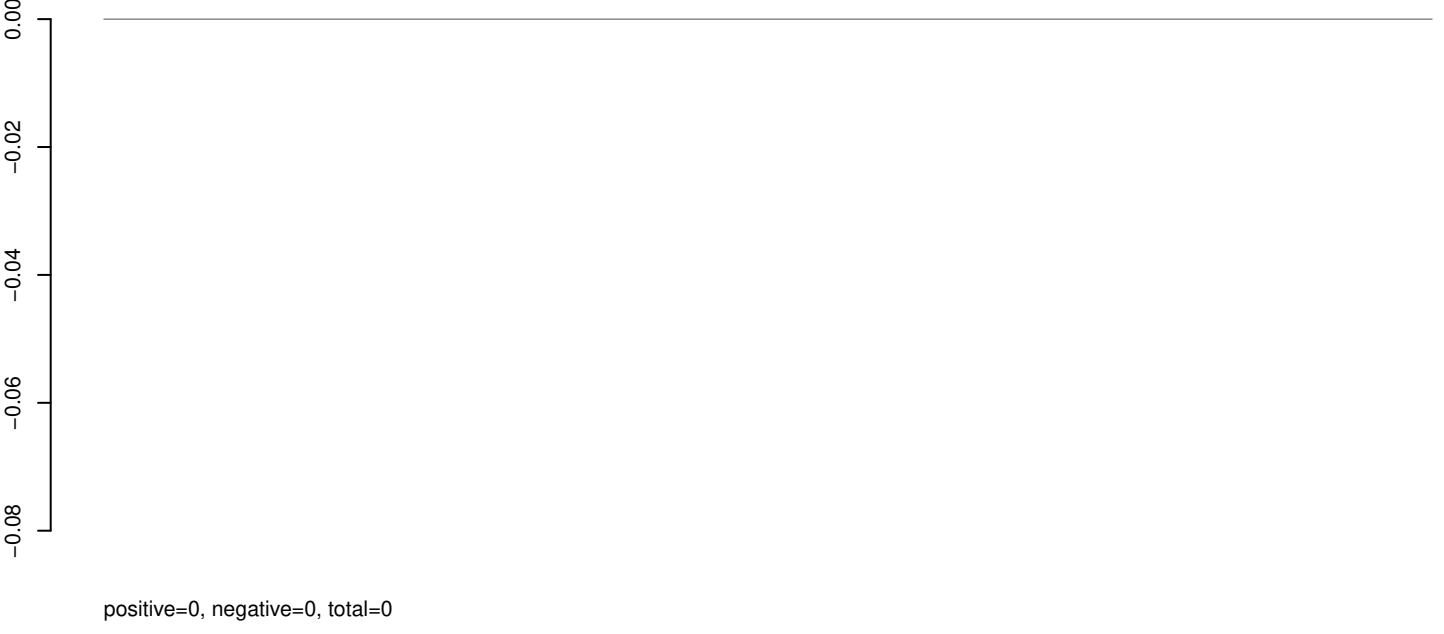
Window size=25, length=10588, virus@NC_031327.1-Anopheles_flavivirus_variant_1:1-10588

0 2000 4000 6000 8000 10000

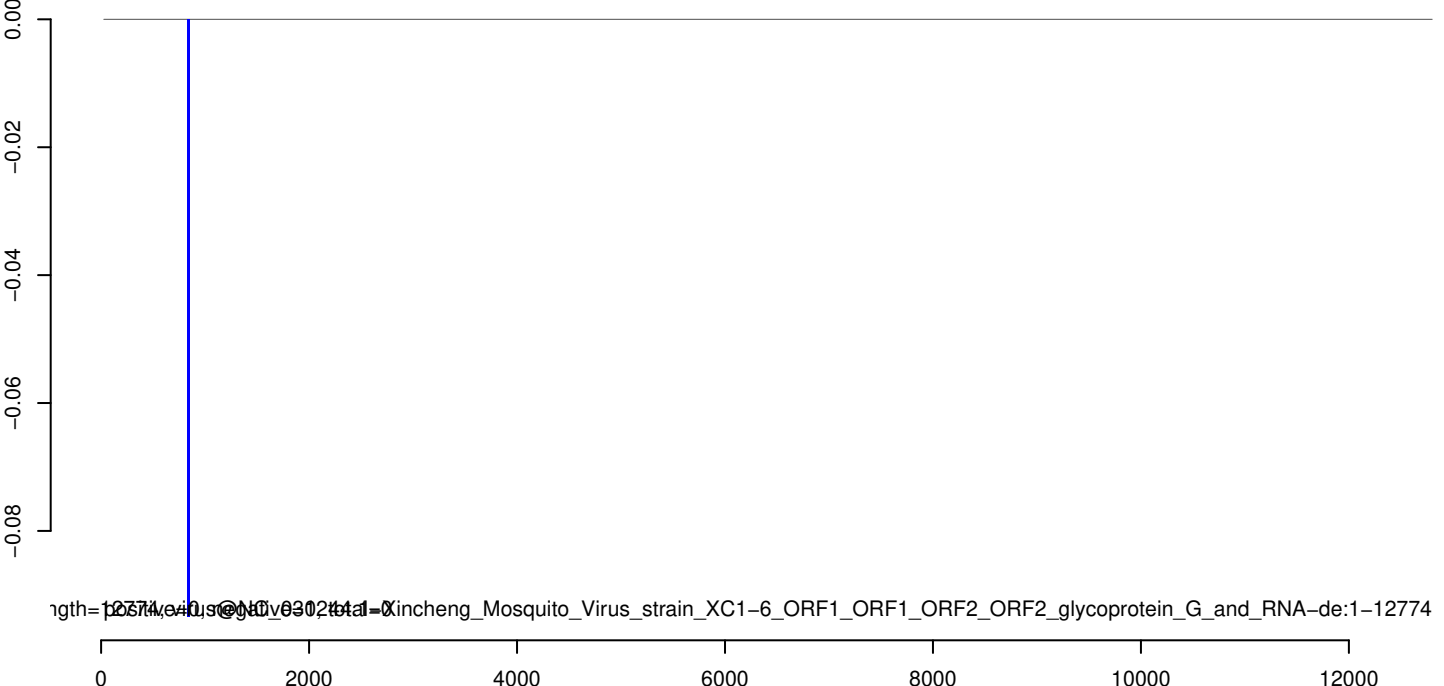
AeAeg_Aag2SINV_dsLuc.18_23.rep



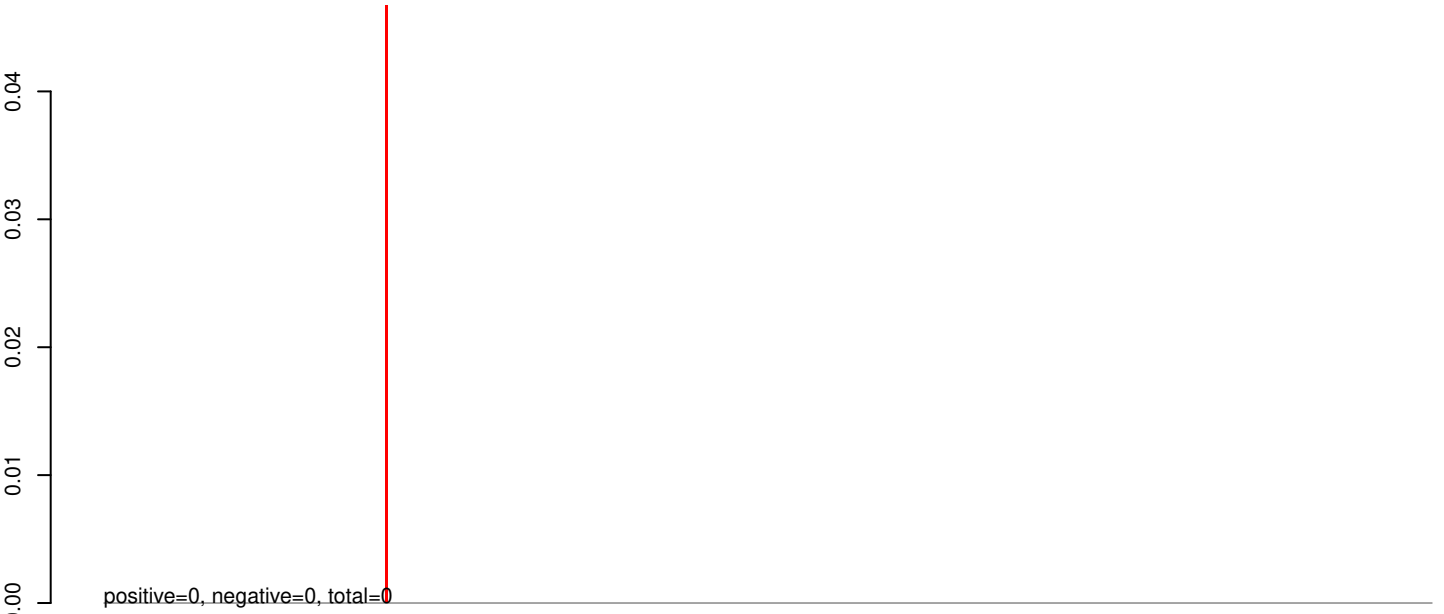
AeAeg_Aag2SINV_dsLuc.24_35.rep



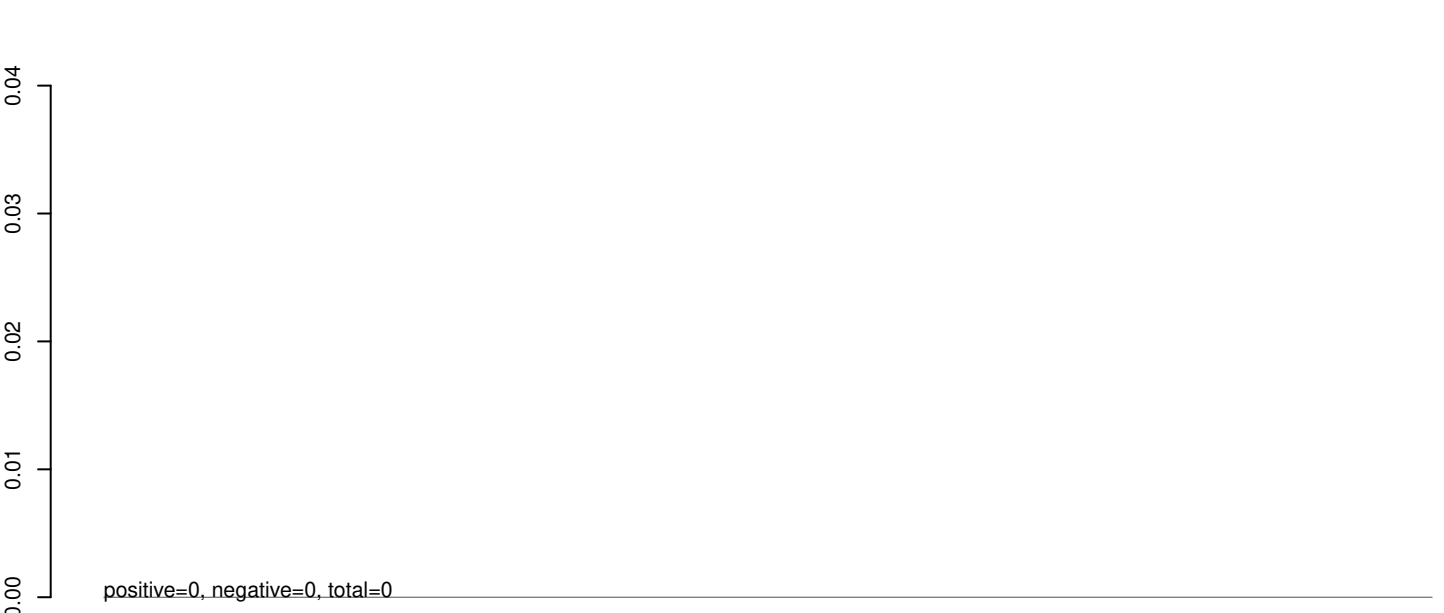
AeAeg_Aag2SINV_dsLuc.rep



AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



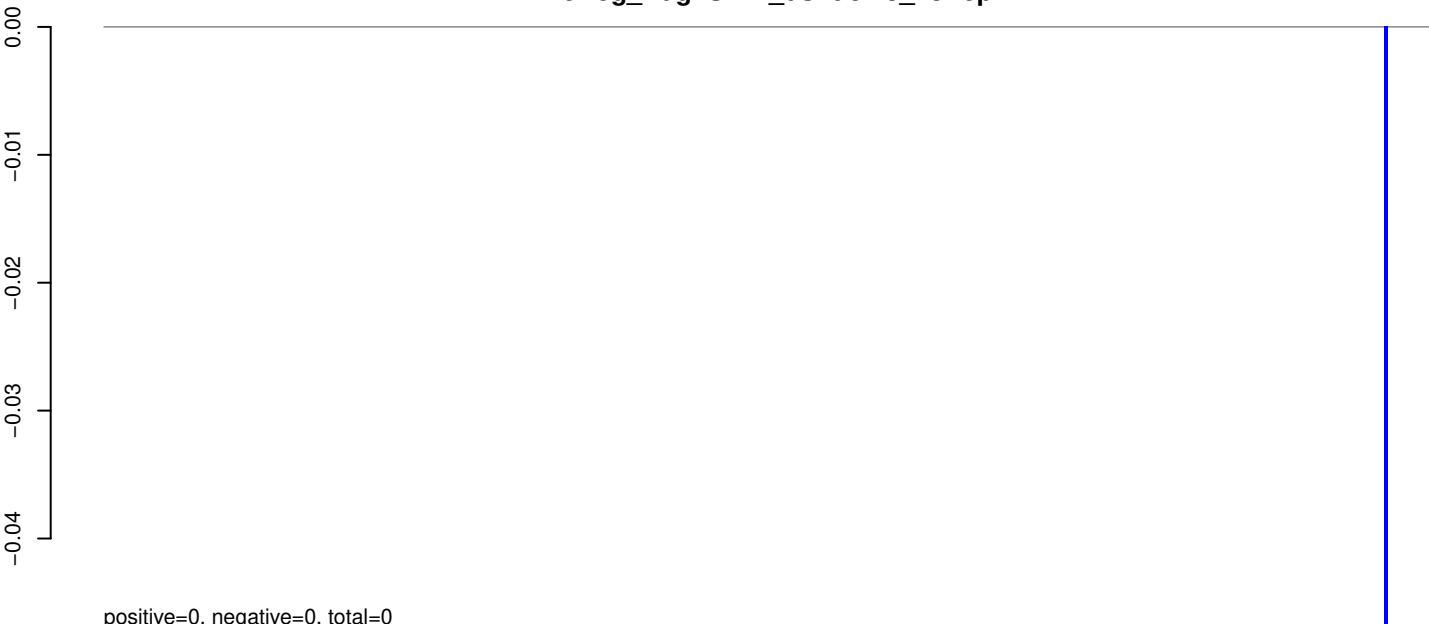
AeAeg_Aag2SINV_dsLuc.rep



Window size=25, length=15444, virus@NC_025400.1-Malaka virus isolate_SudAr_1169-64:1-15444

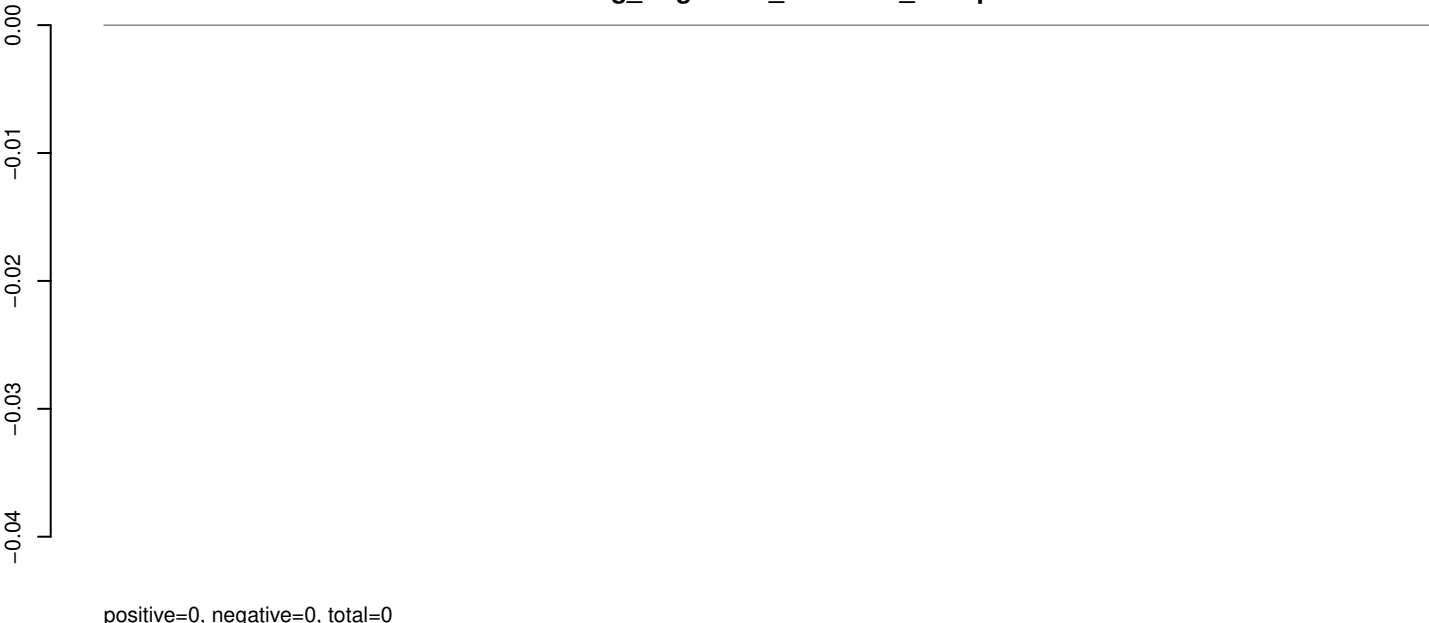
0 5000 10000 15000

AeAeg_Aag2SINV_dsLuc.18_23.rep



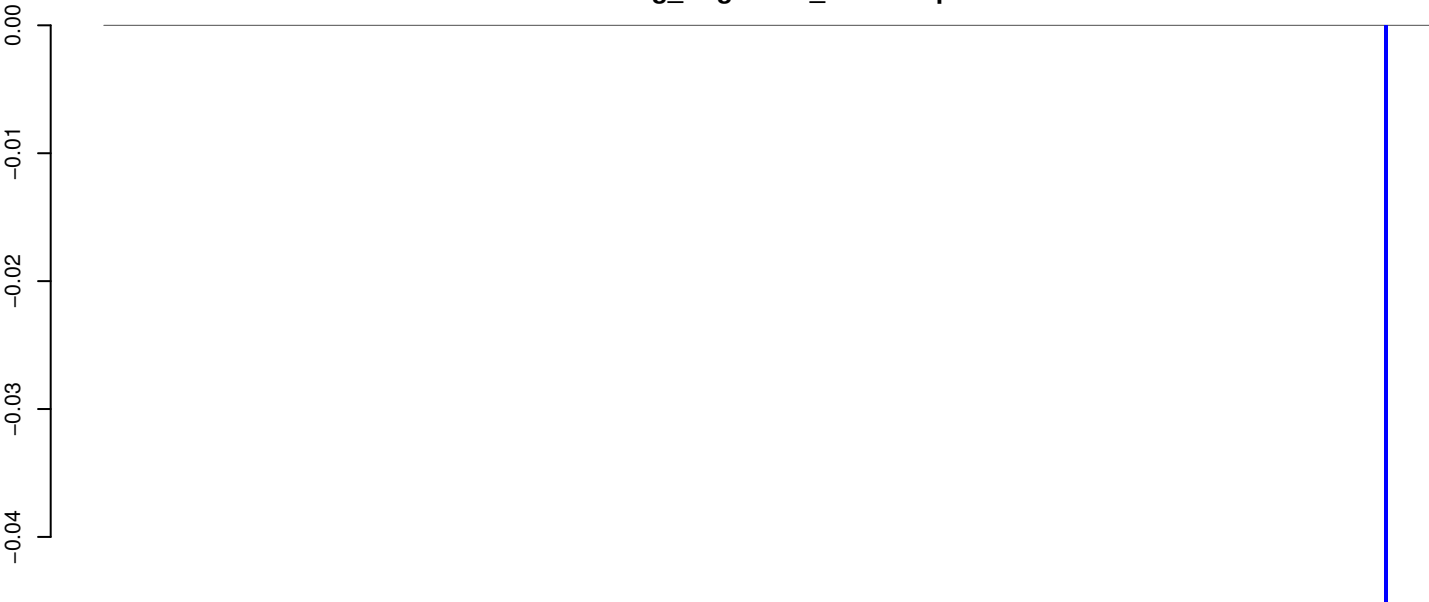
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=0, negative=0, total=0

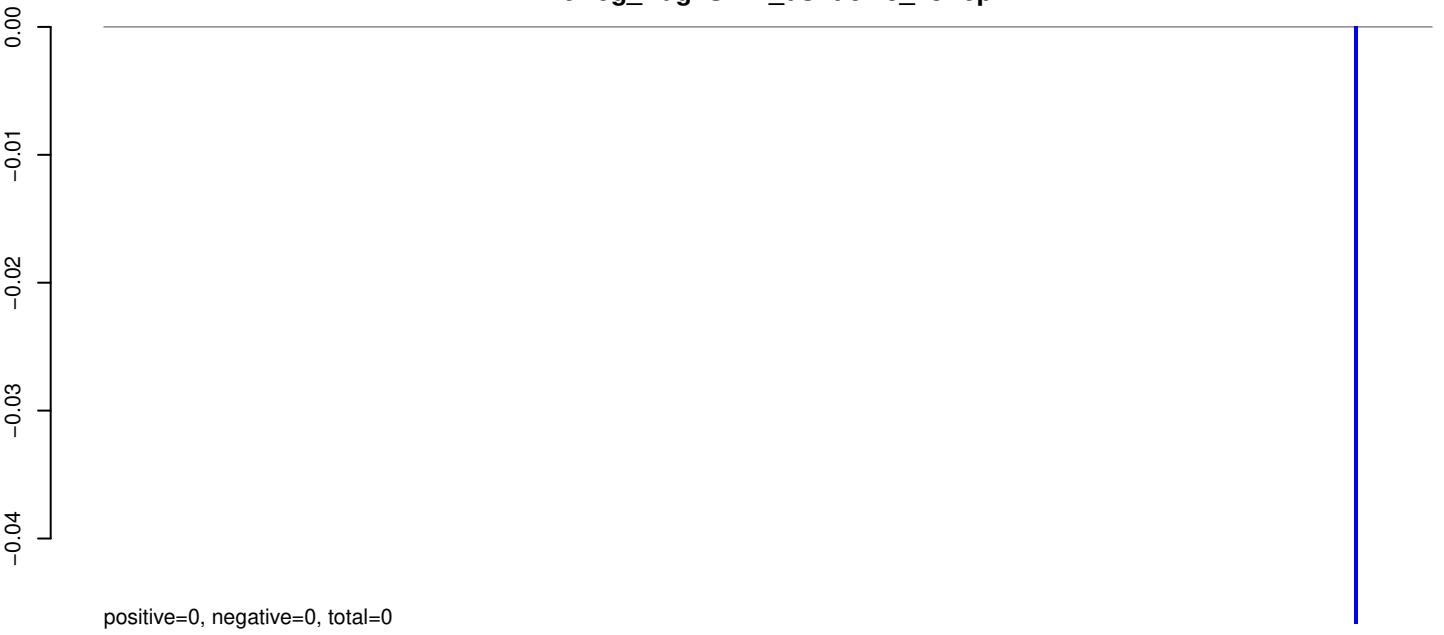
AeAeg_Aag2SINV_dsLuc.rep



positive=0, negative=0, total=0 window size=25, length=10320, virus@NC_024806.1-Lammi_virus_isolate_M0719_polyprotein_gene:1-10320

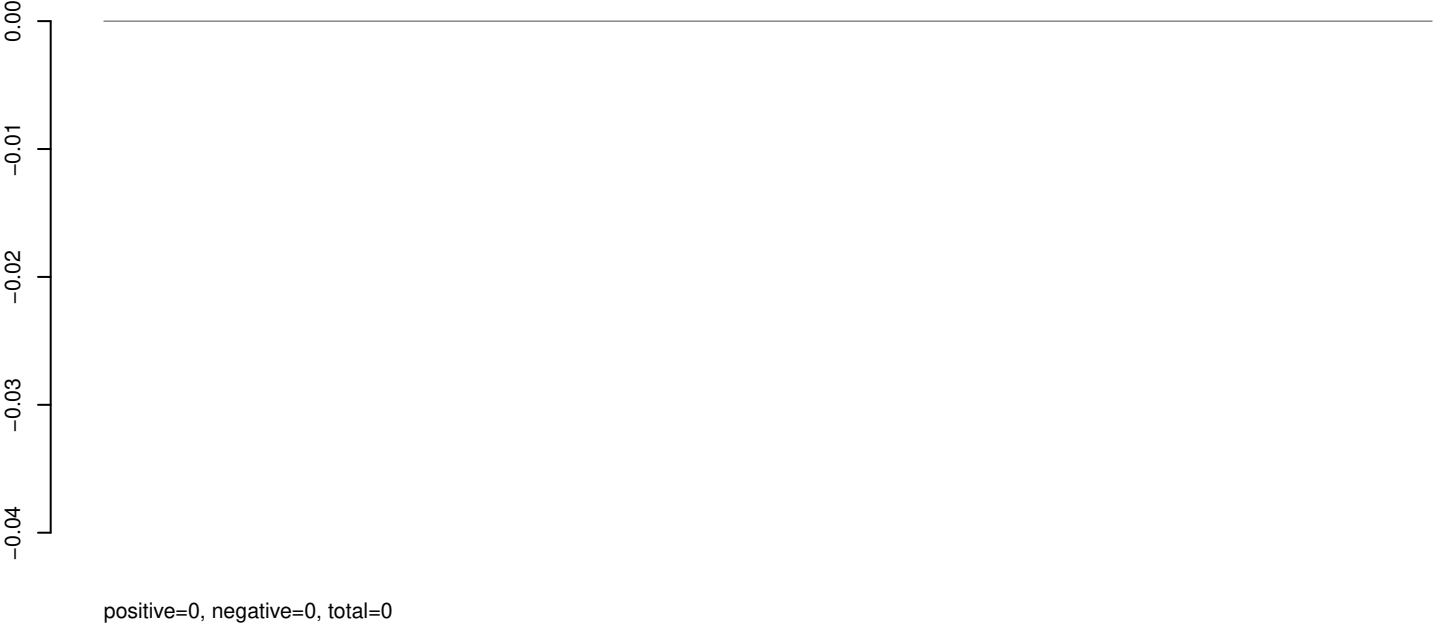


AeAeg_Aag2SINV_dsLuc.18_23.rep



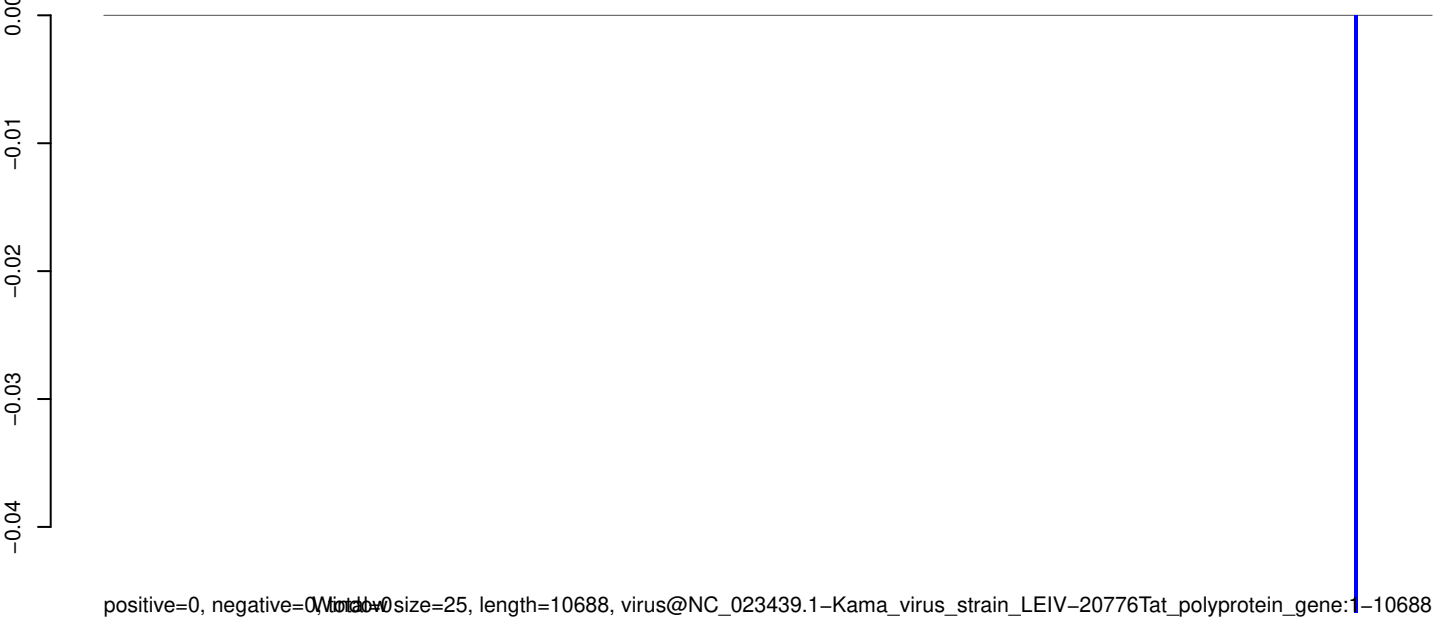
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=0, negative=0, total=0

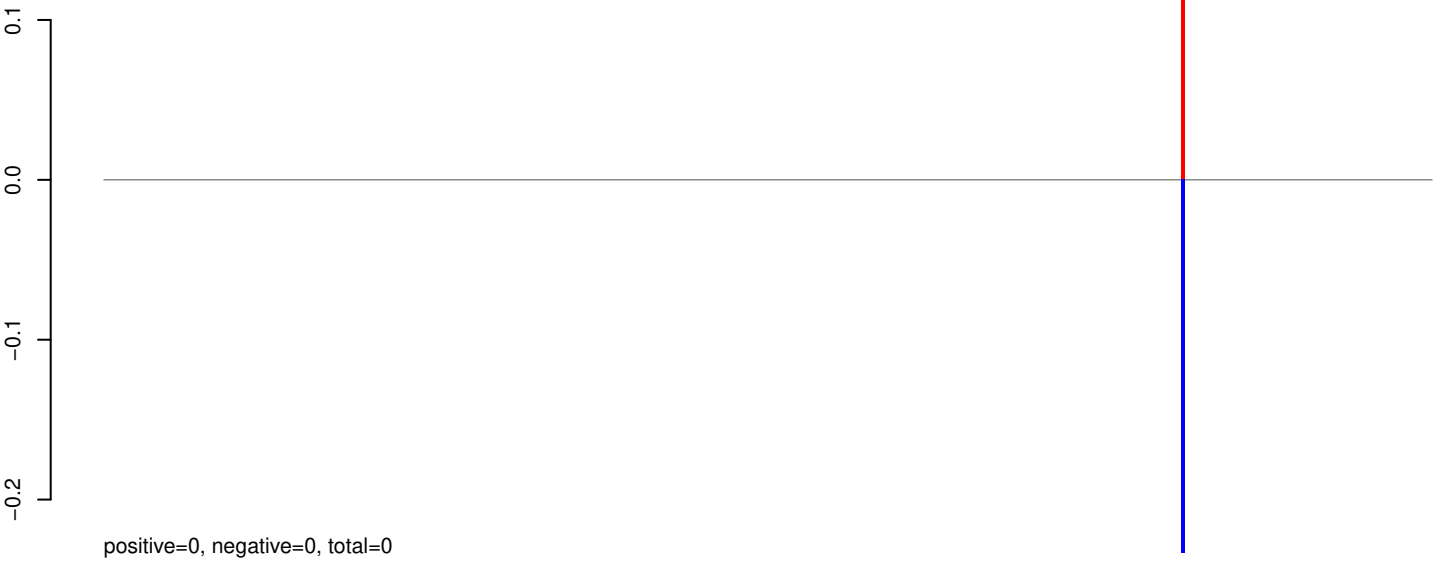
AeAeg_Aag2SINV_dsLuc.rep



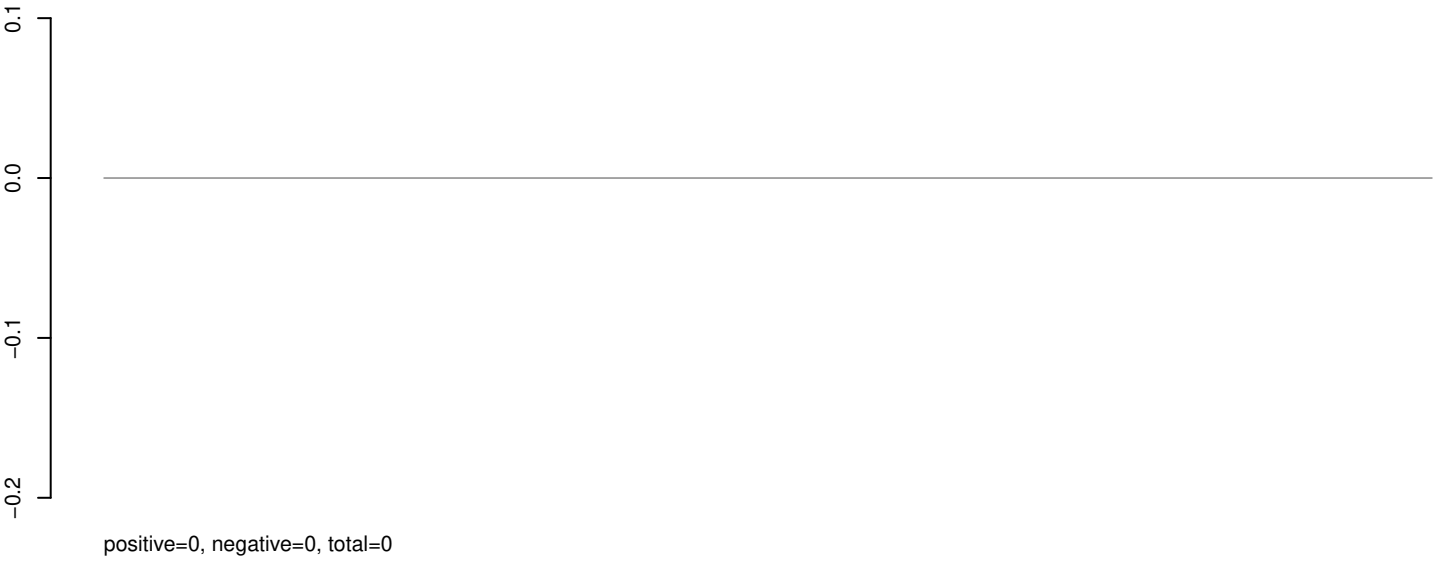
positive=0, negative=0, total=0, window size=25, length=10688, virus@NC_023439.1-Kama_virus_strain_LEIV-20776Tat_polyprotein_gene:1-10688

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



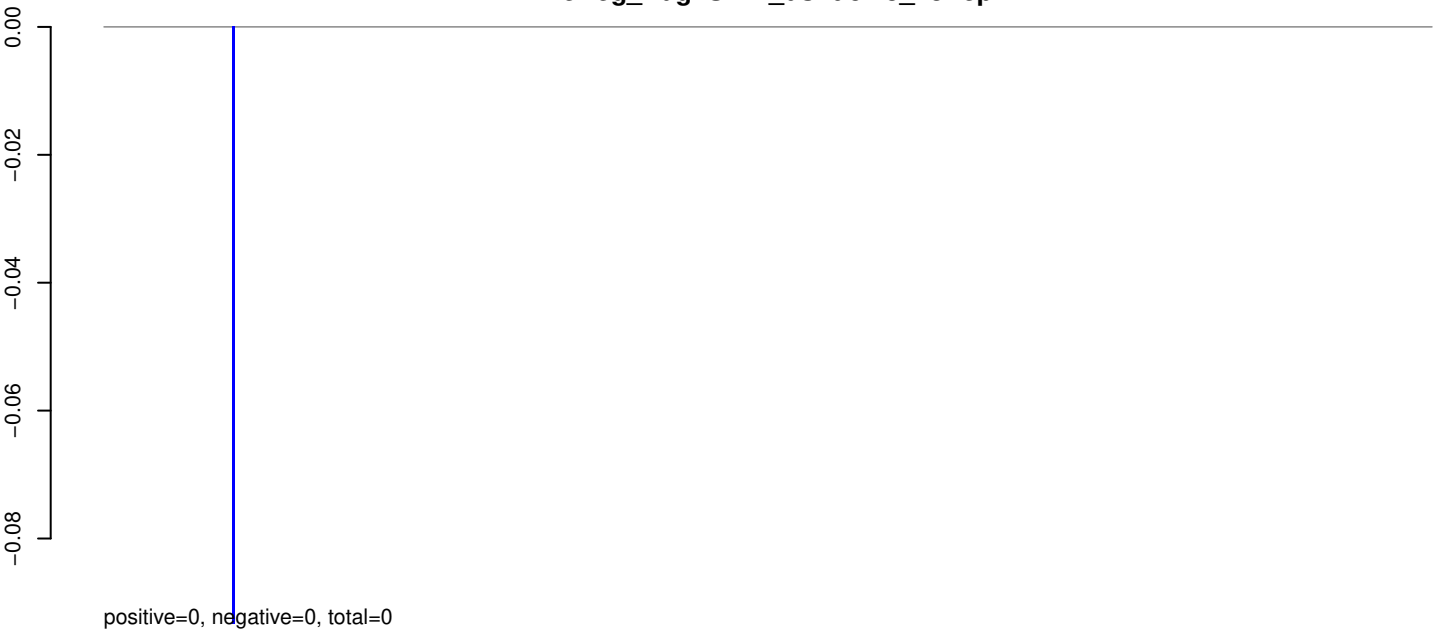
AeAeg_Aag2SINV_dsLuc.rep



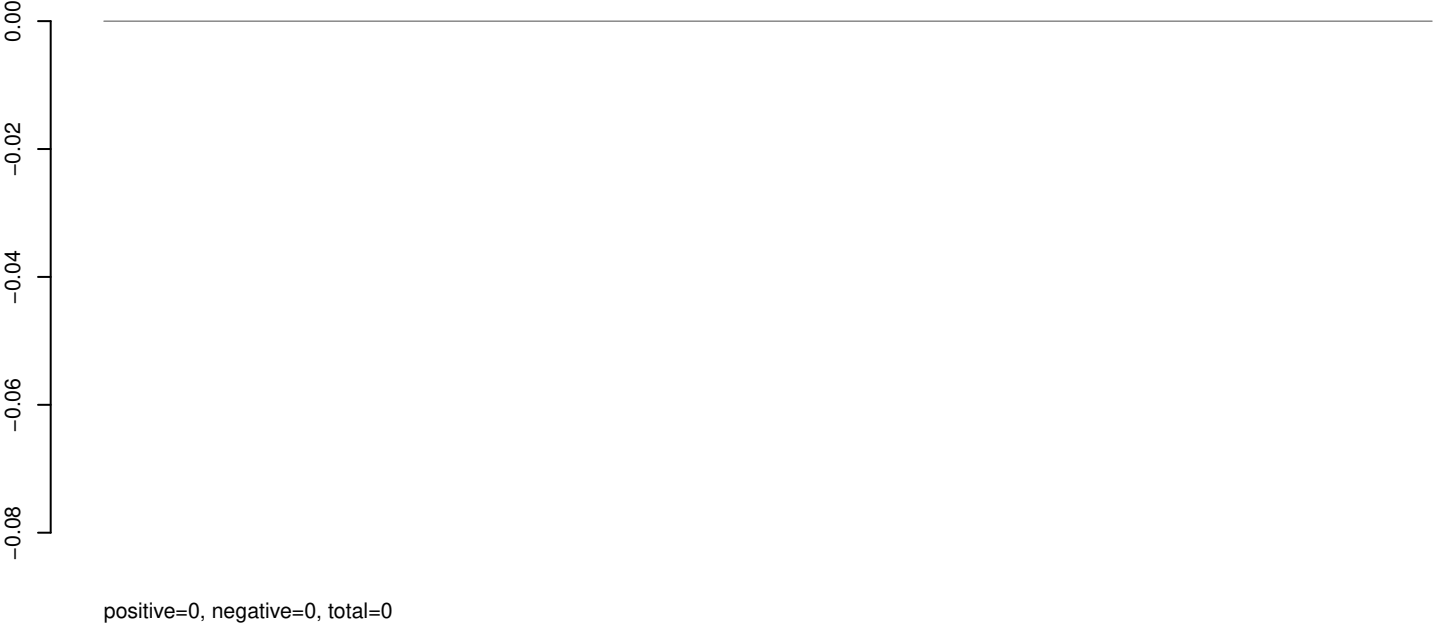
Window size=25, length=10865, virus@NC_021069.1-Mosquito_flavivirus_isolate_LSFlaviV-A20-09:1-10865

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

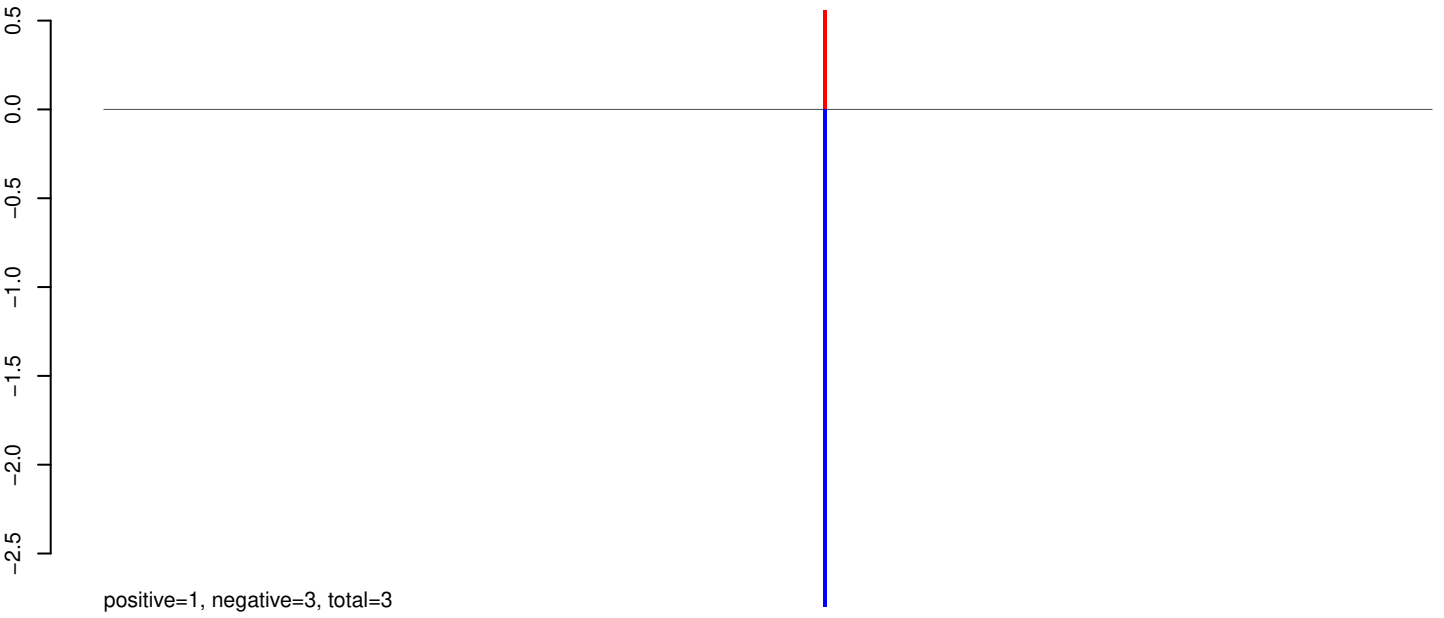


AeAeg_Aag2SINV_dsLuc.rep

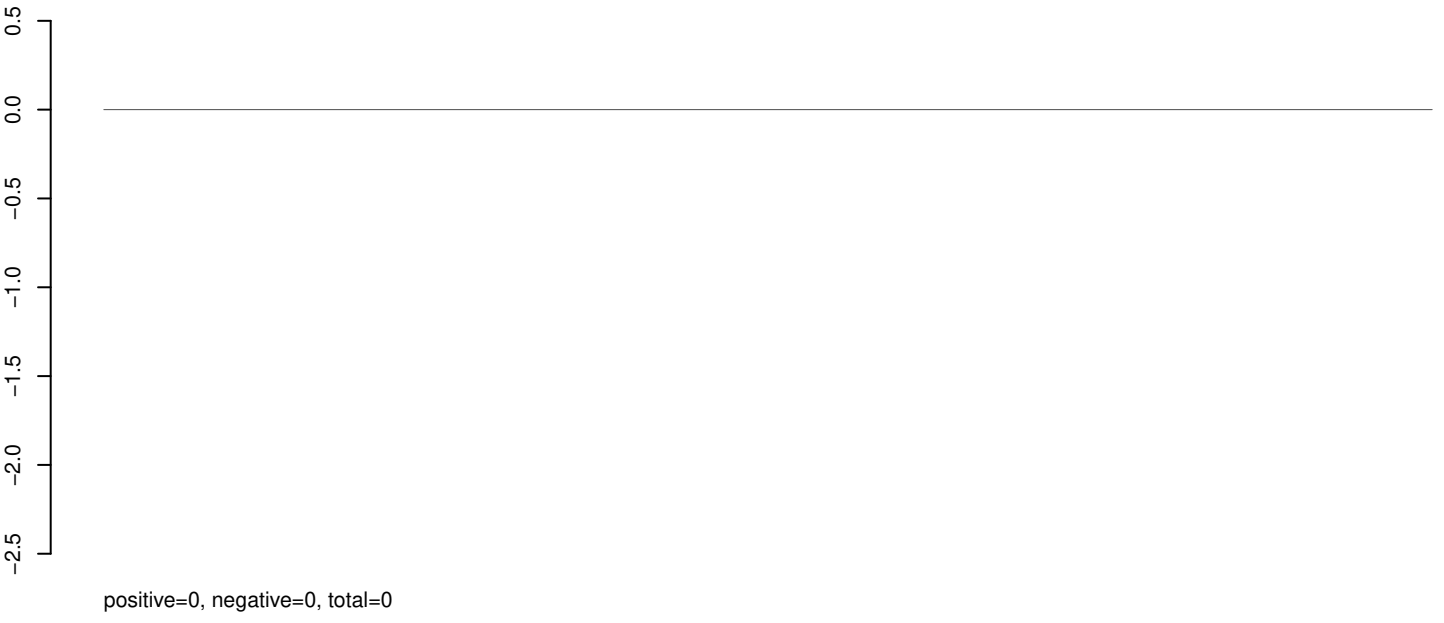


Window size=25, length=14717, virus@NC_017685.1-Obodhiang_virus:1-14717

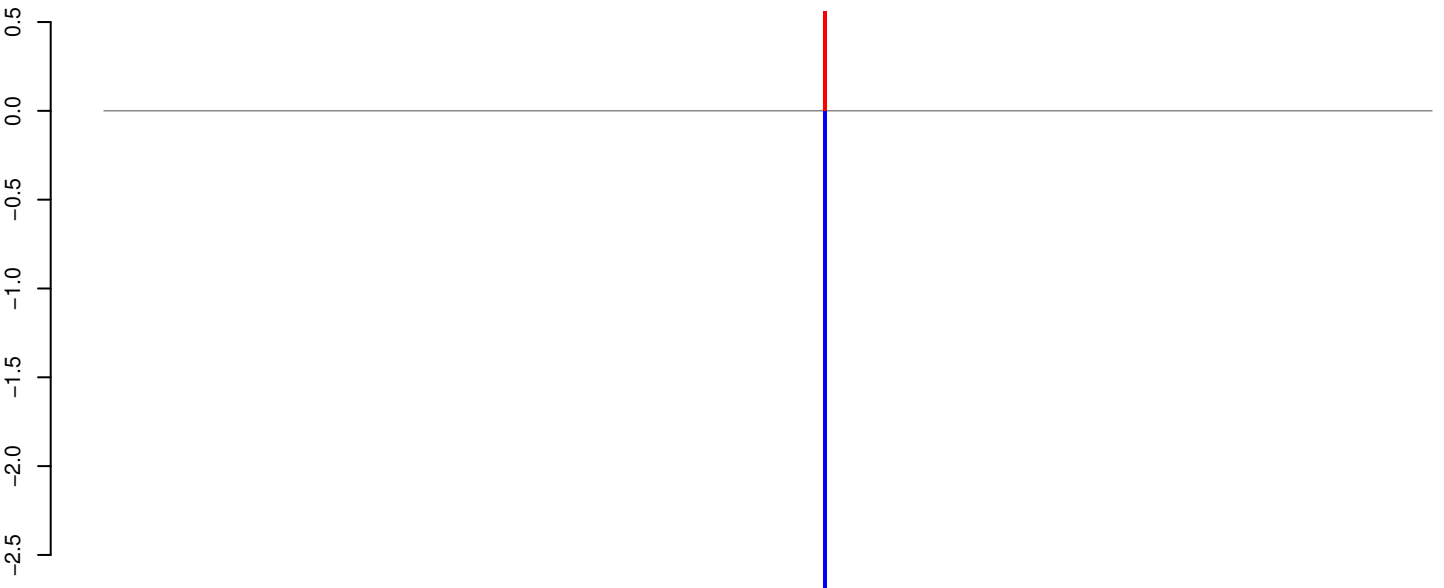
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



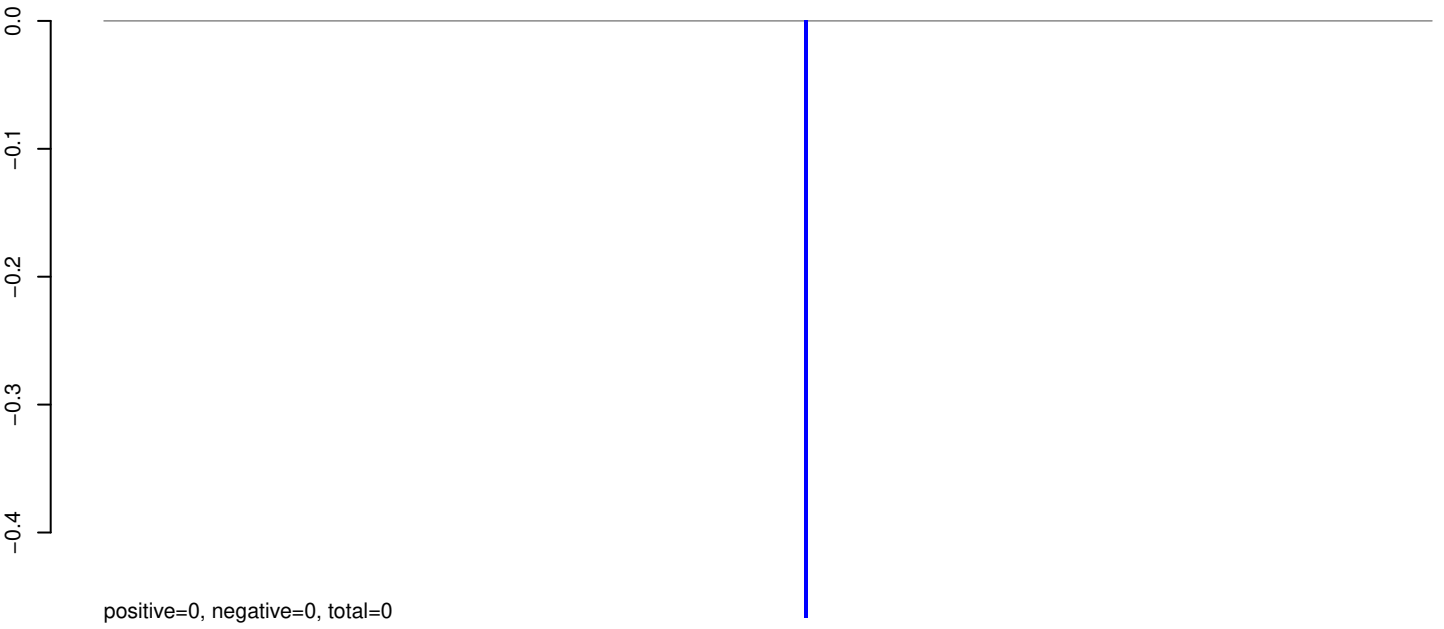
AeAeg_Aag2SINV_dsLuc.rep



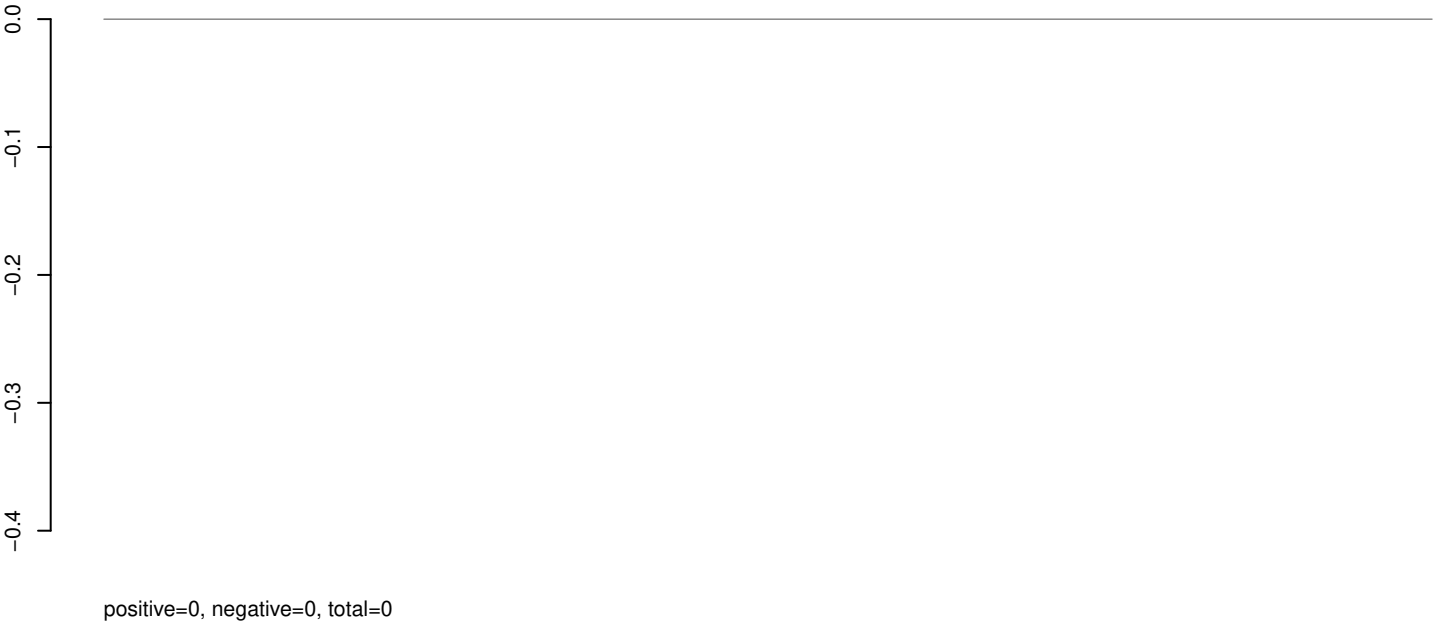
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0 2000 4000 6000 8000 10000

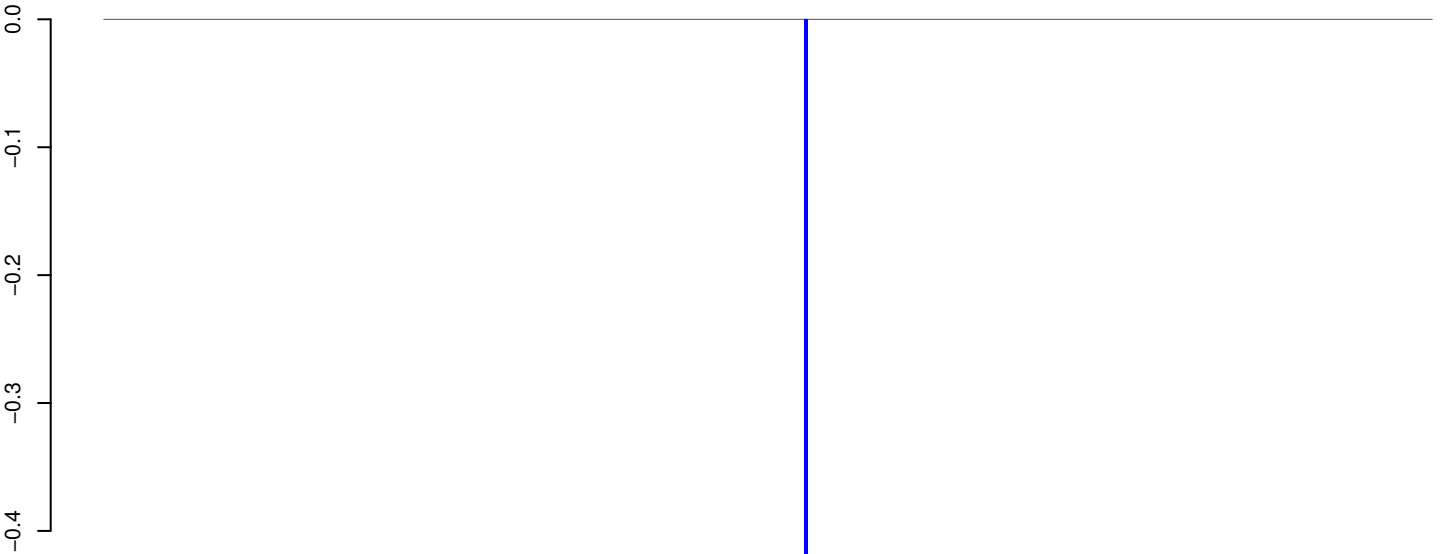
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



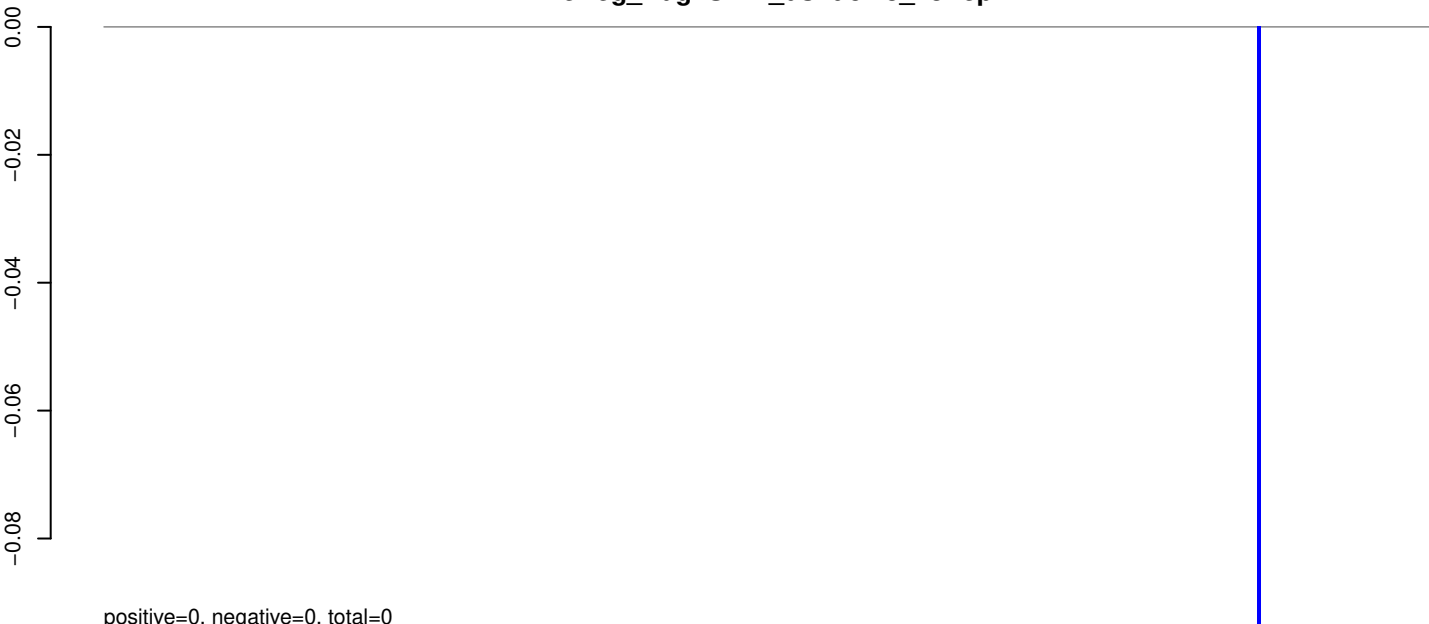
AeAeg_Aag2SINV_dsLuc.rep



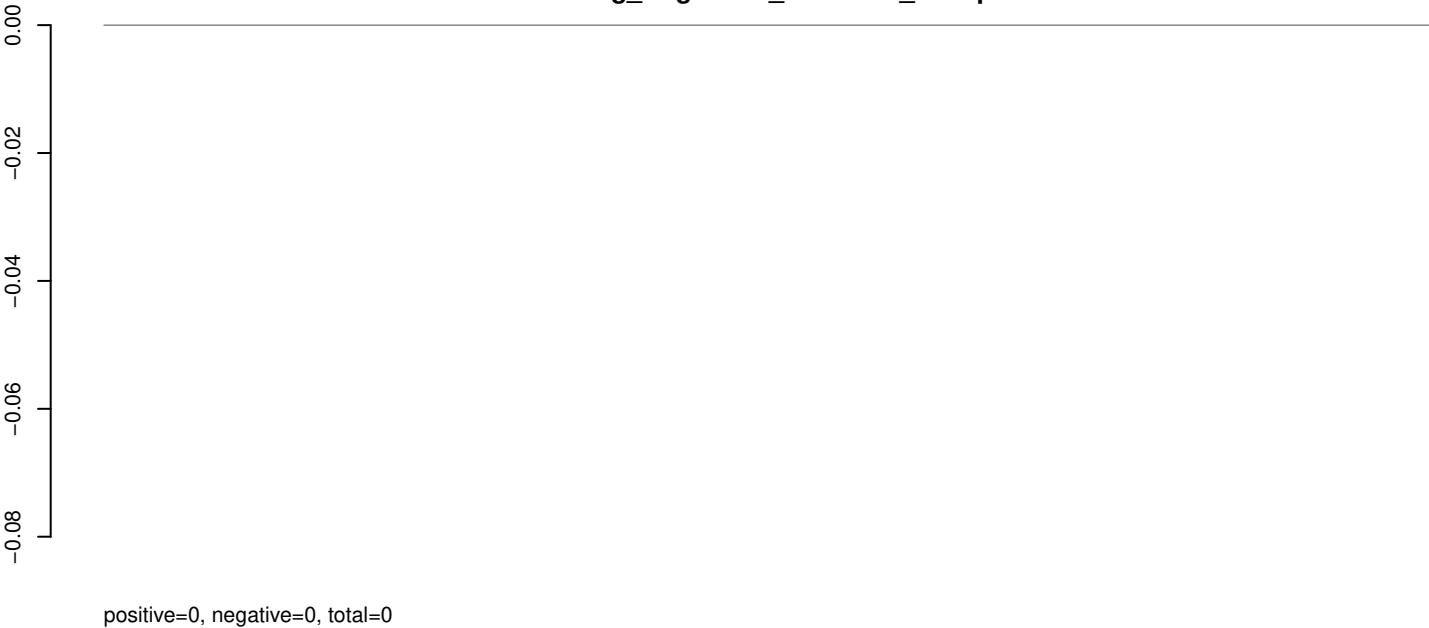
Windows size=25, length=10990, virus@NC_015843.2-Tembusu_virus_flavivirus_polyprotein_flavivirus_polyprotein_gene_gene:1-10990

0 2000 4000 6000 8000 10000

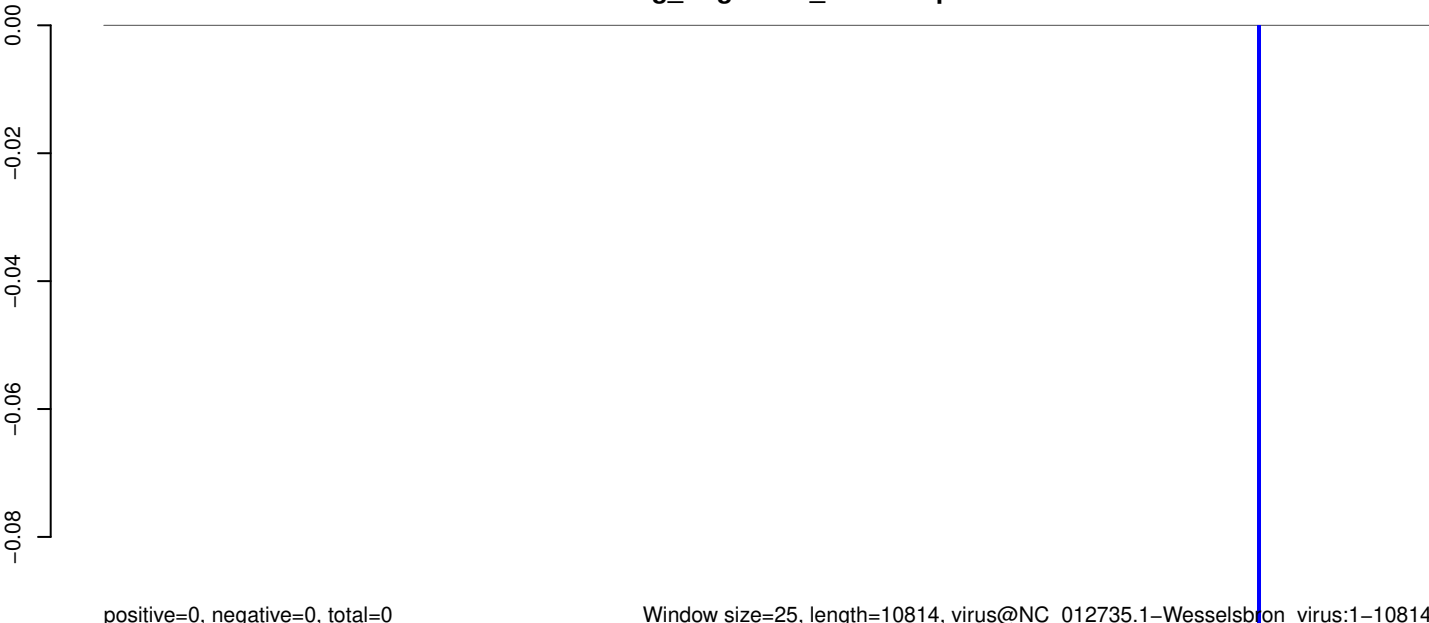
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



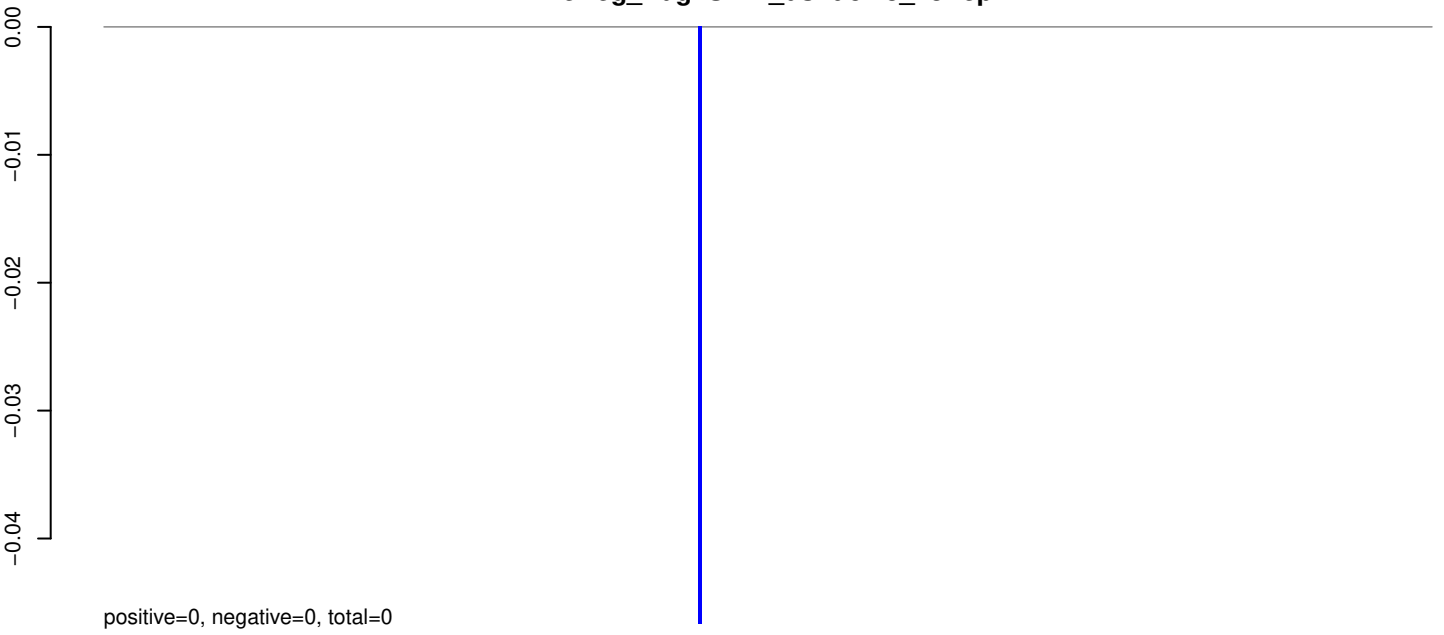
AeAeg_Aag2SINV_dsLuc.rep



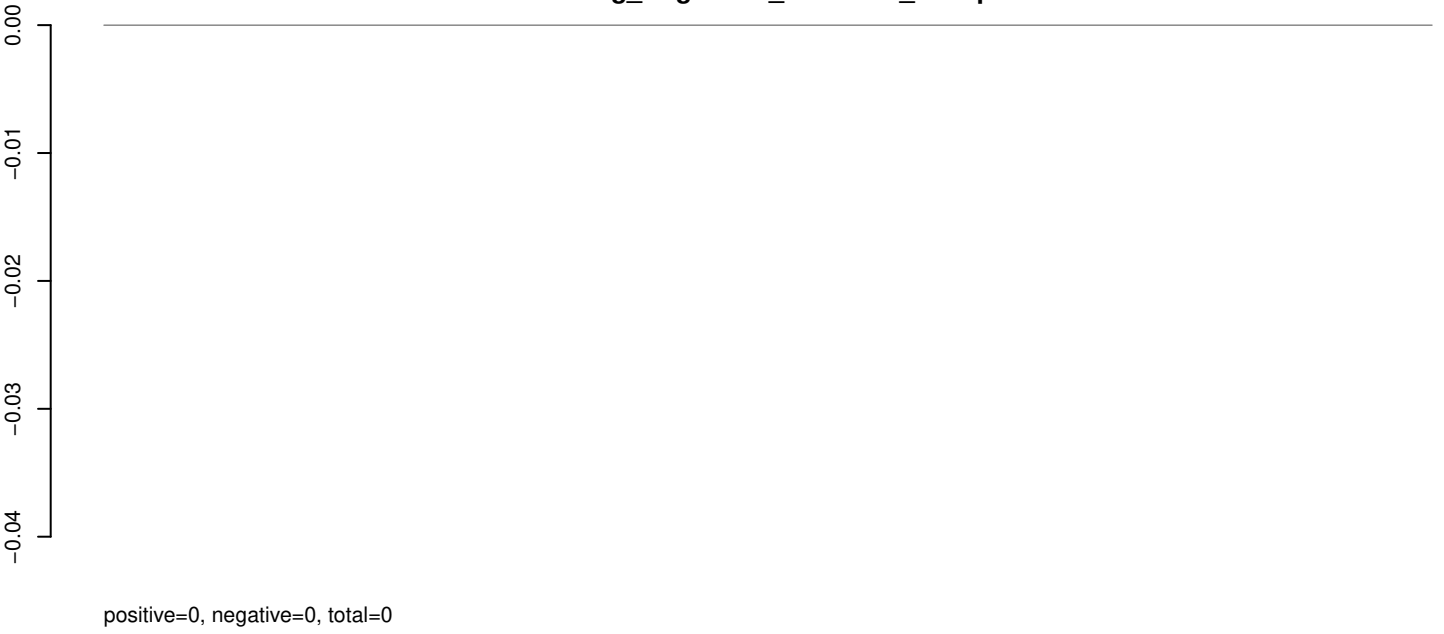
Window size=25, length=10814, virus@NC_012735.1-Wesselsbron_virus:1-10814

0 2000 4000 6000 8000 10000

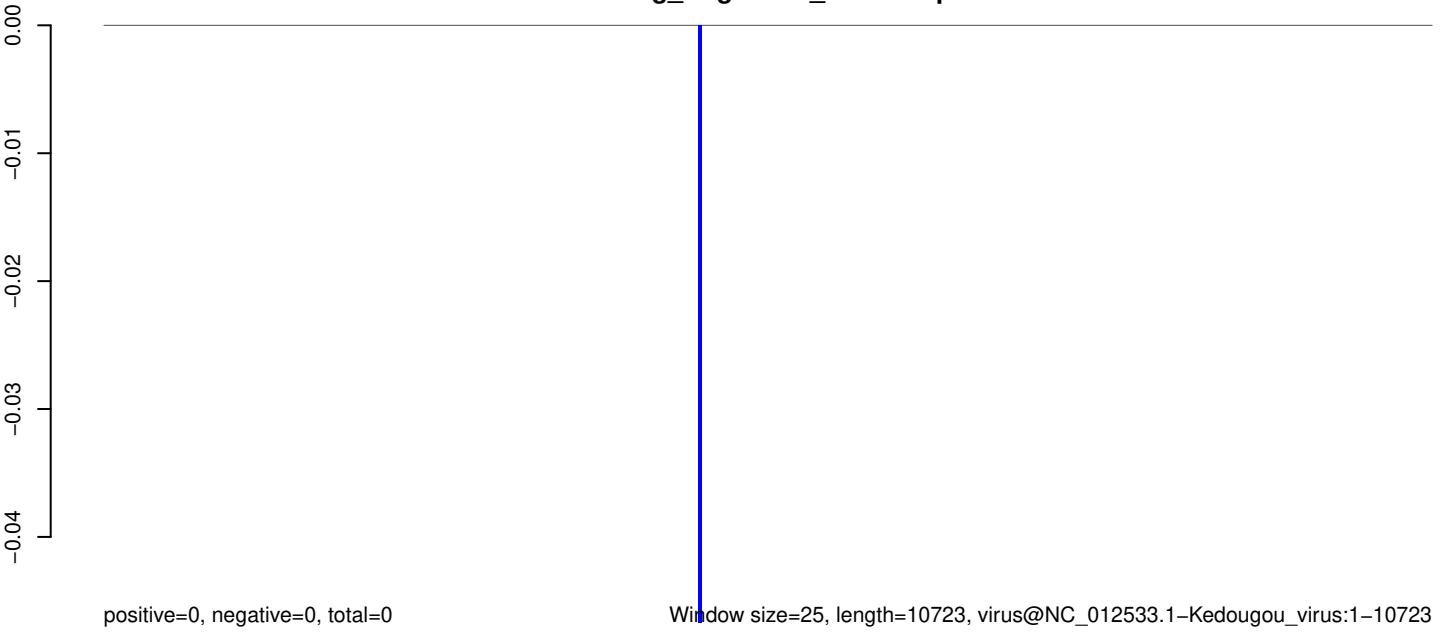
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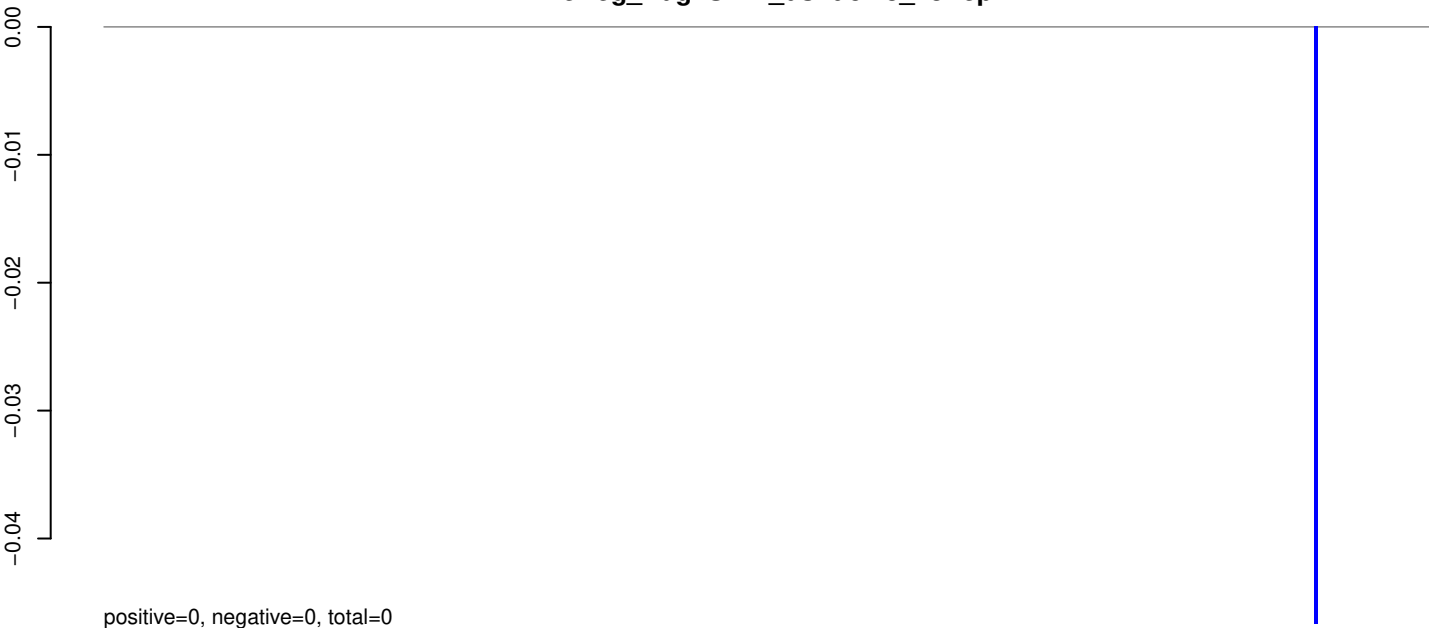
AeAeg_Aag2SINV_dsLuc.24_35.rep



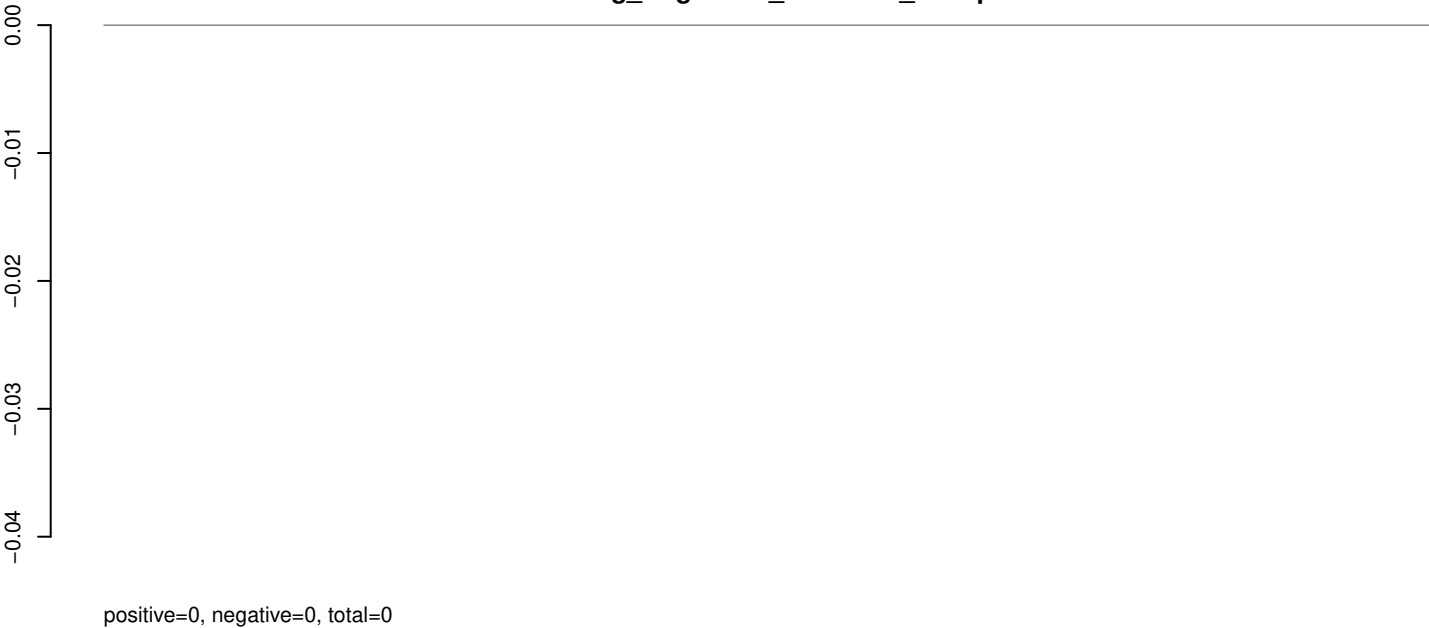
AeAeg_Aag2SINV_dsLuc.rep



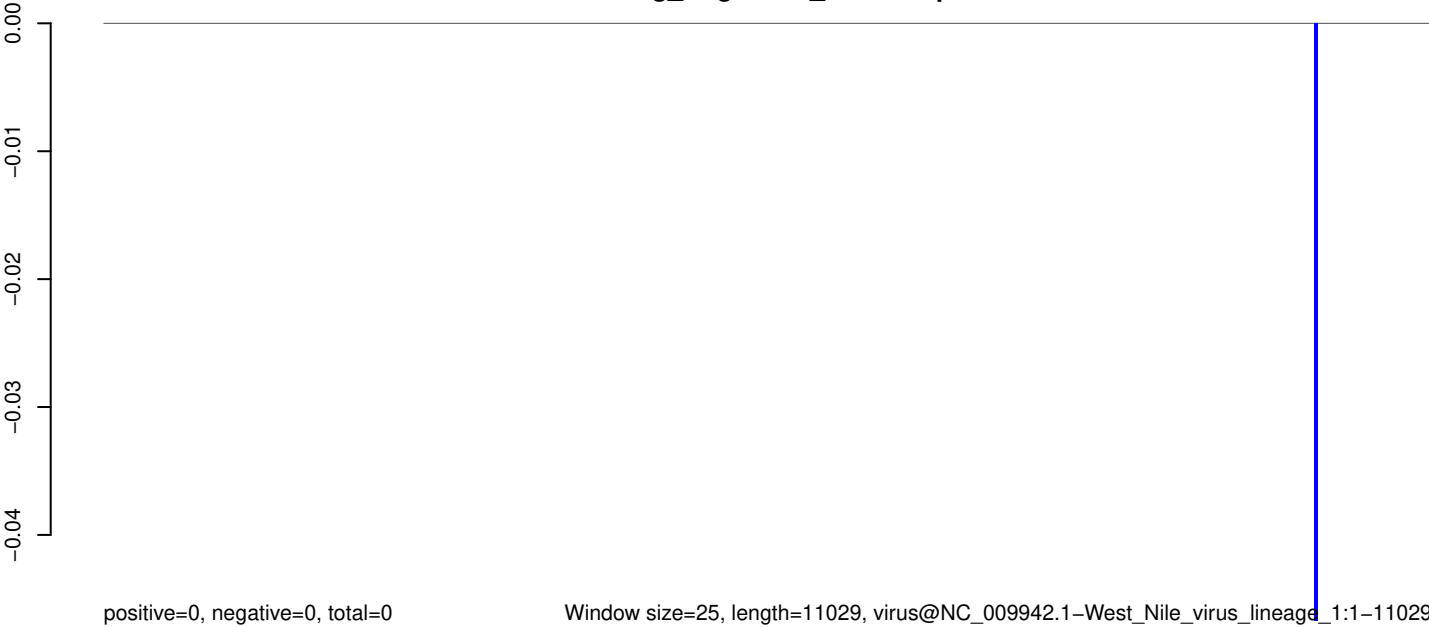
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AeAeg_Aag2SINV_dsLuc.24_35.rep



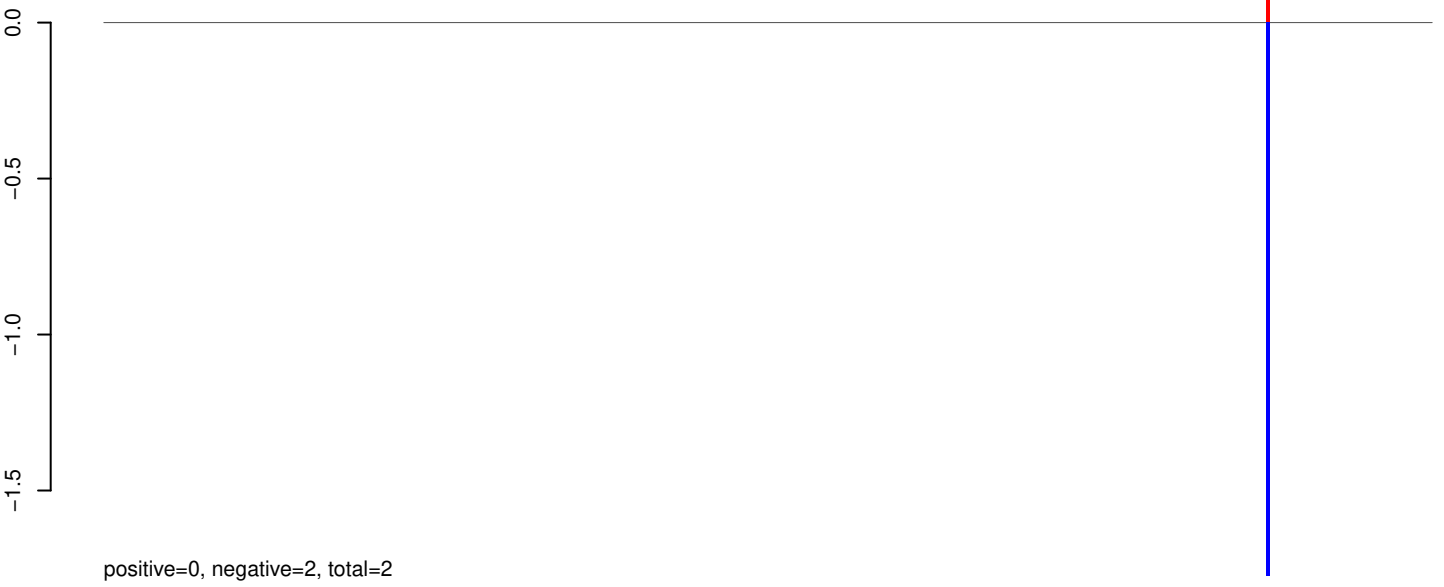
AeAeg_Aag2SINV_dsLuc.rep



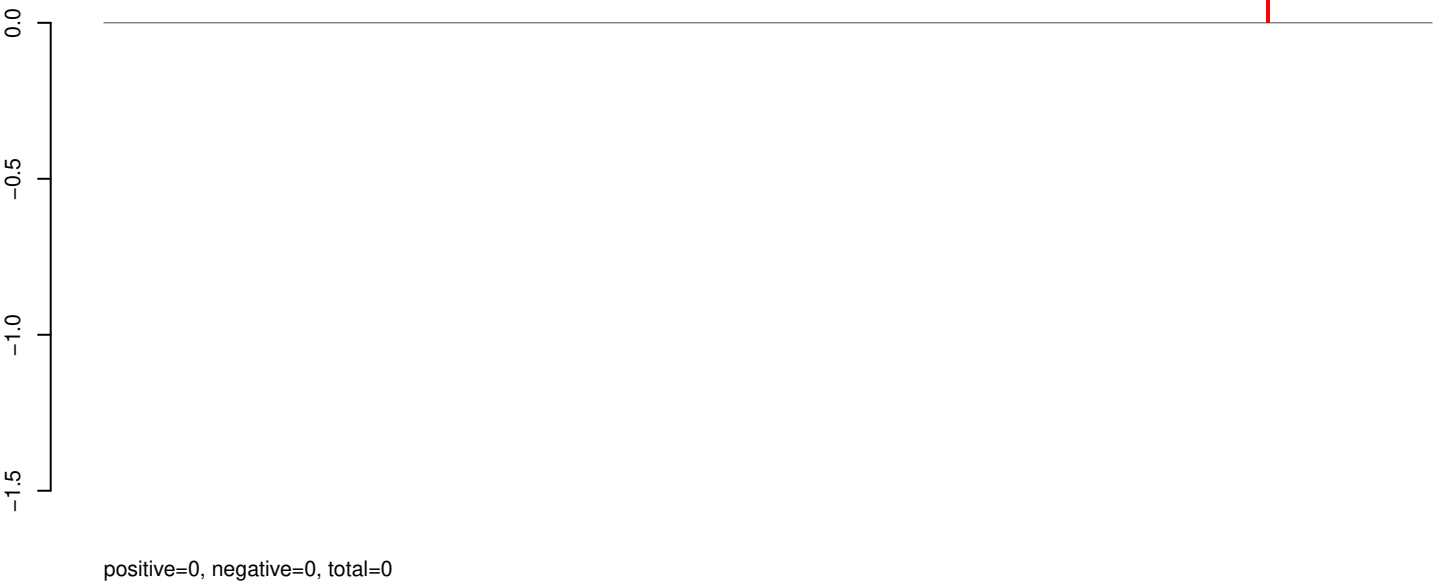
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0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



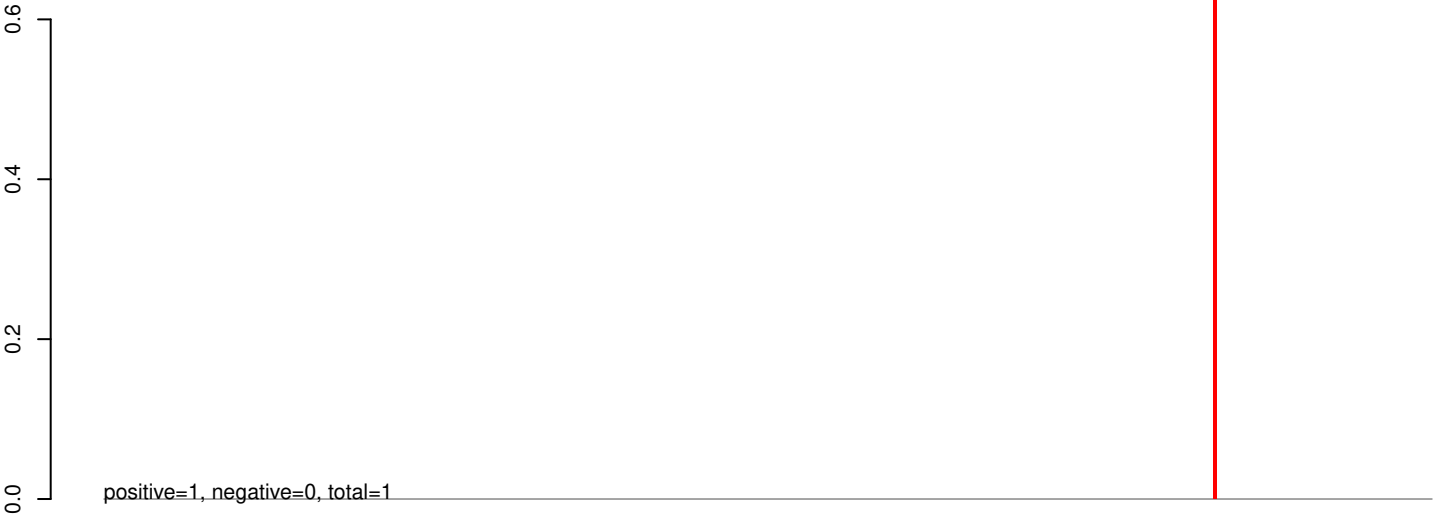
AeAeg_Aag2SINV_dsLuc.rep



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

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_dsLuc.18_23.rep



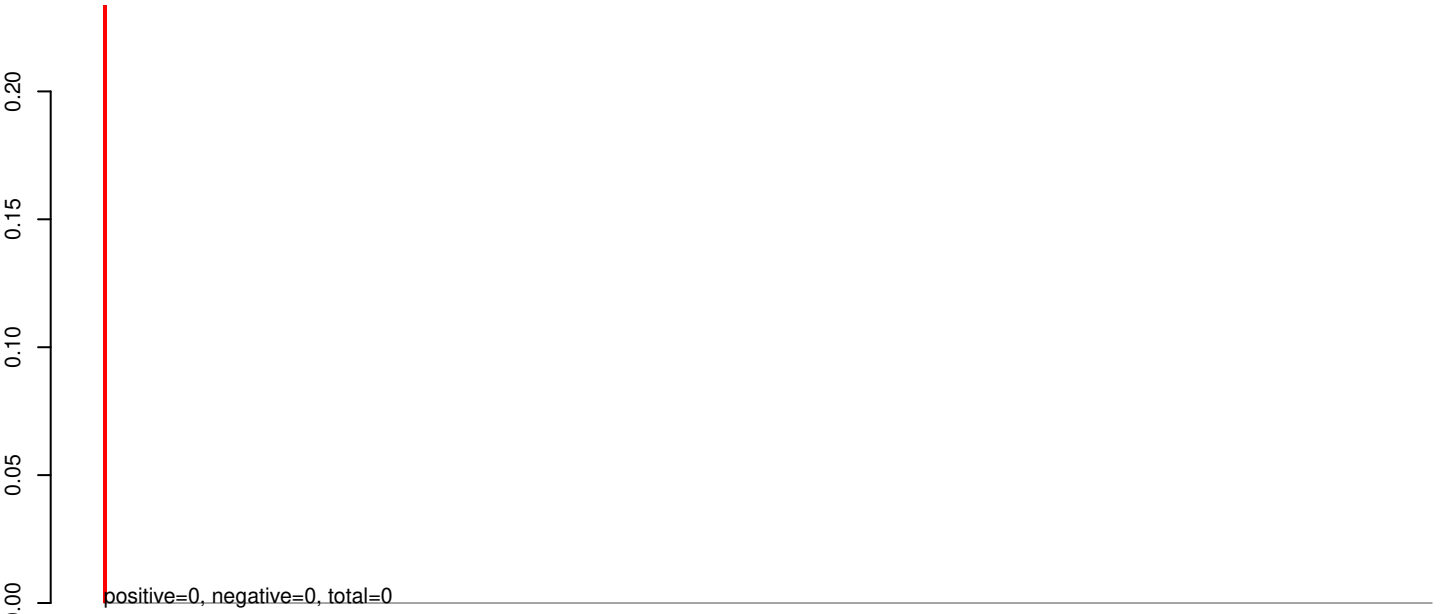
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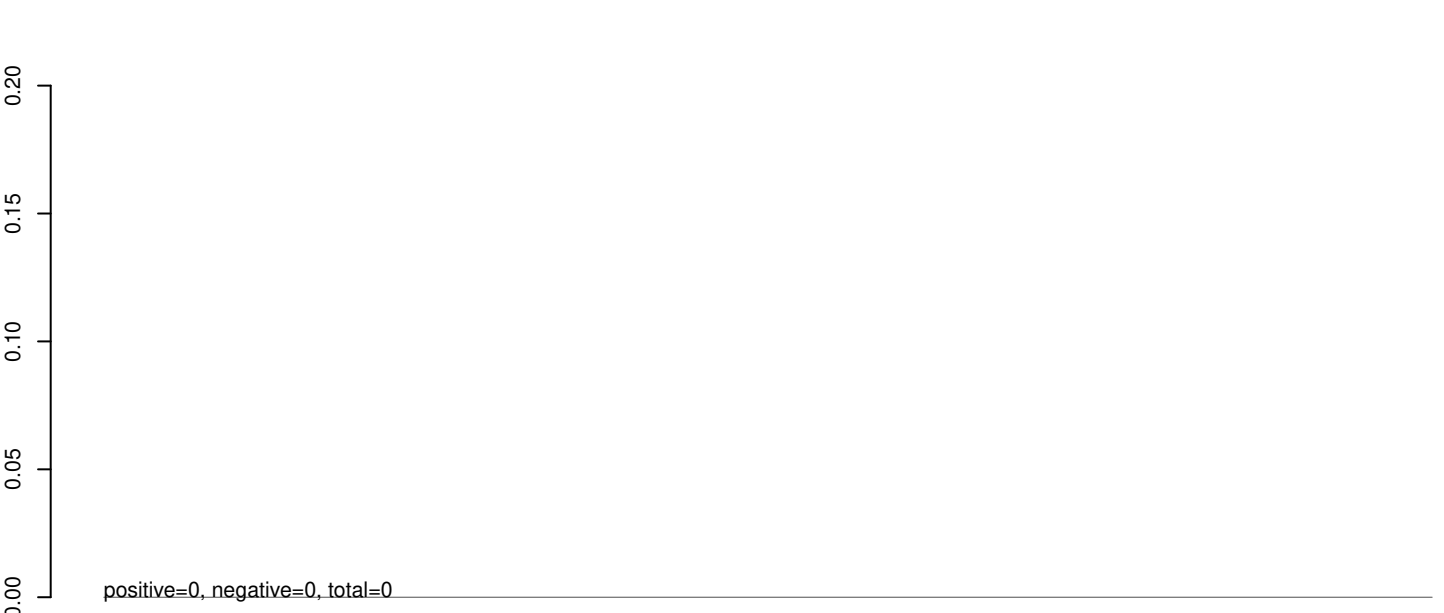
AeAeg_Aag2SINV_dsLuc.rep



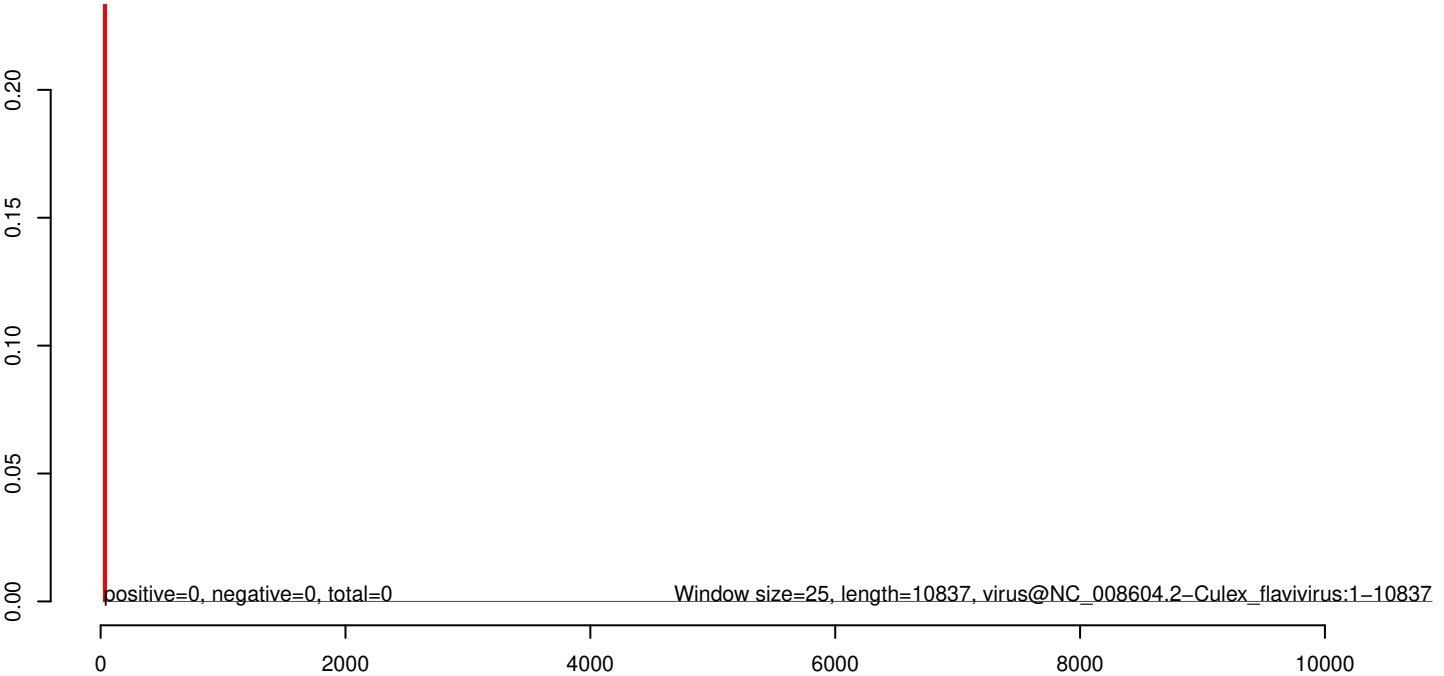
AeAeg_Aag2SINV_dsLuc.18_23.rep



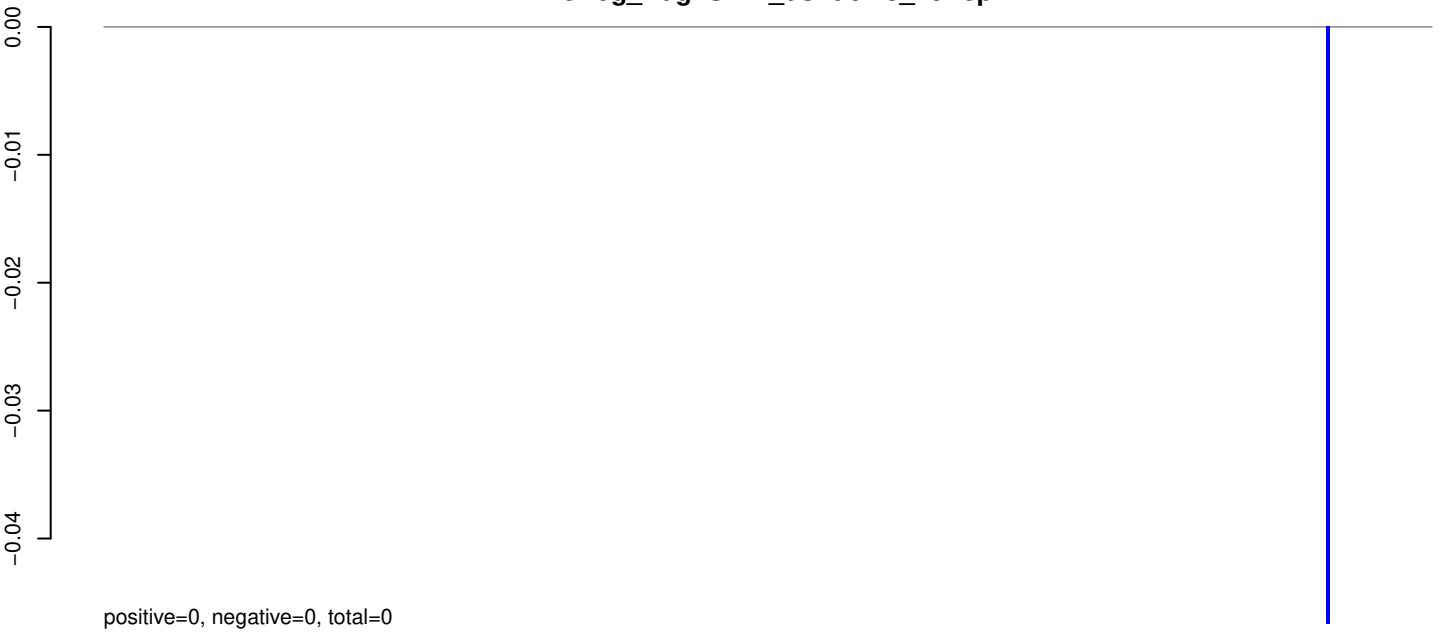
AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep

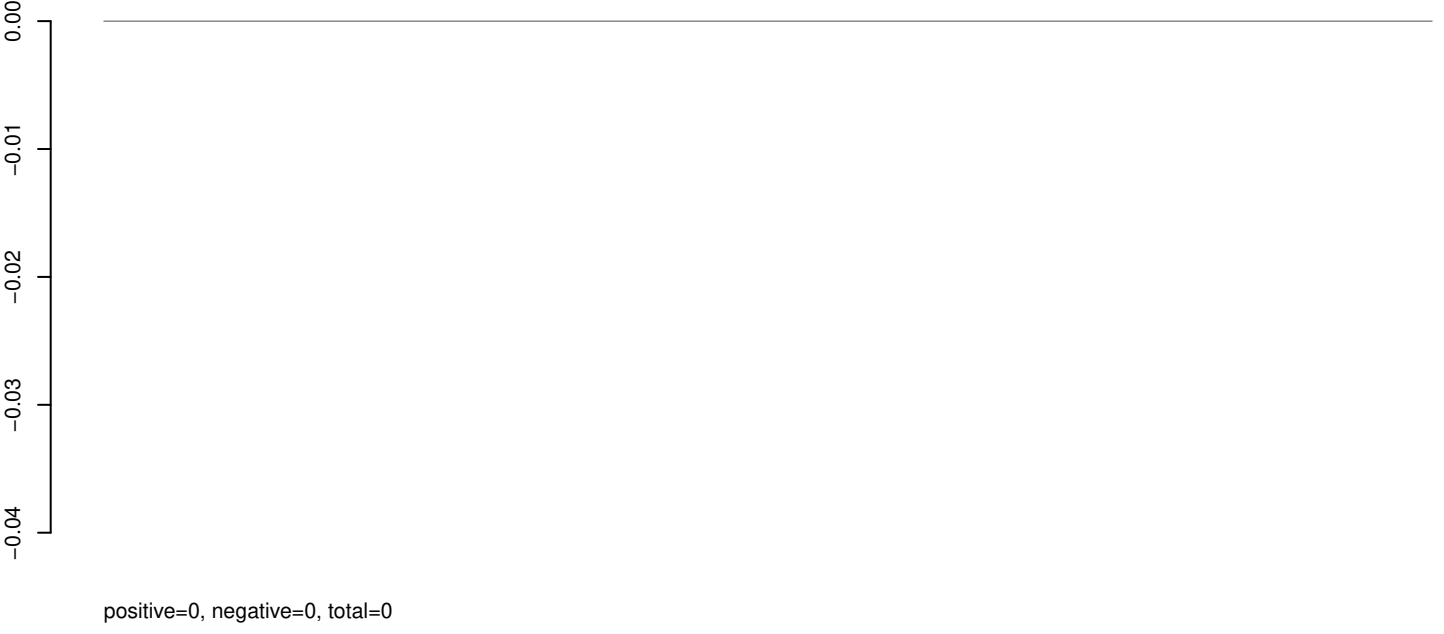


AeAeg_Aag2SINV_dsLuc.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep



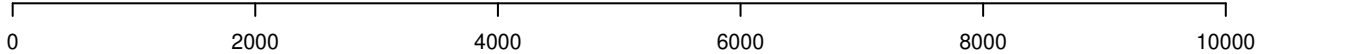
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.rep



positive=0, negative=0, total=0

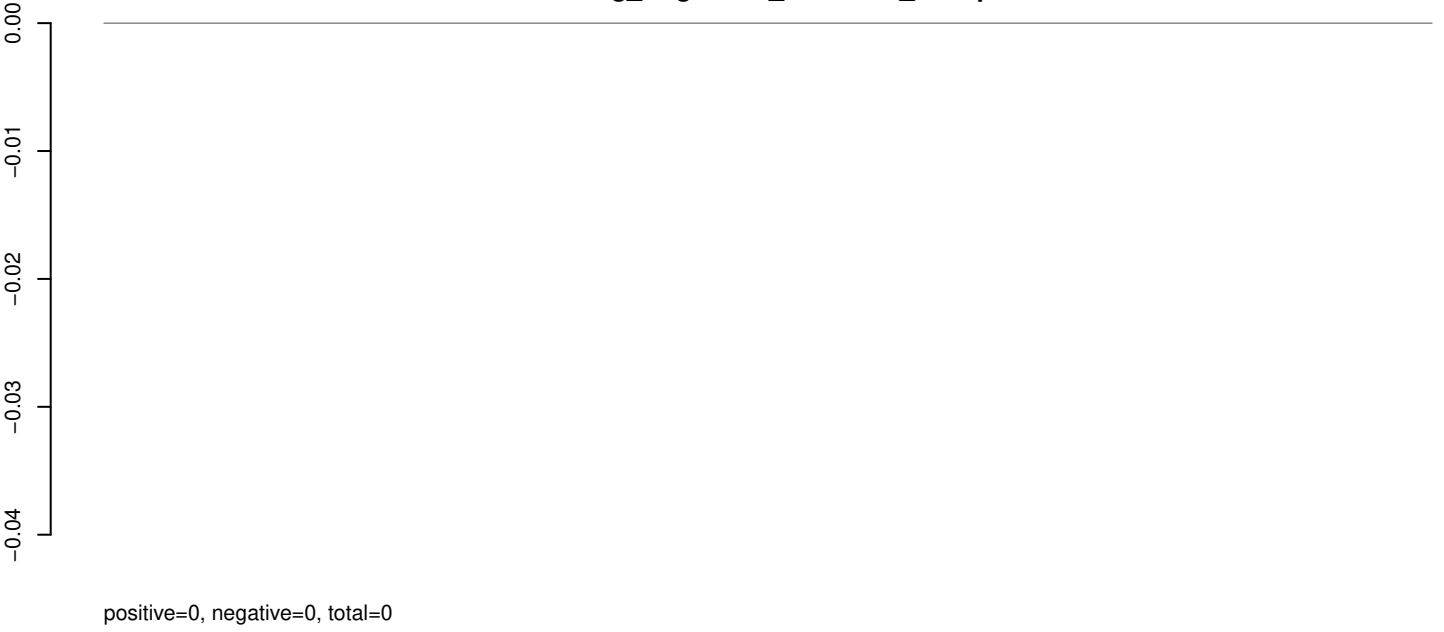
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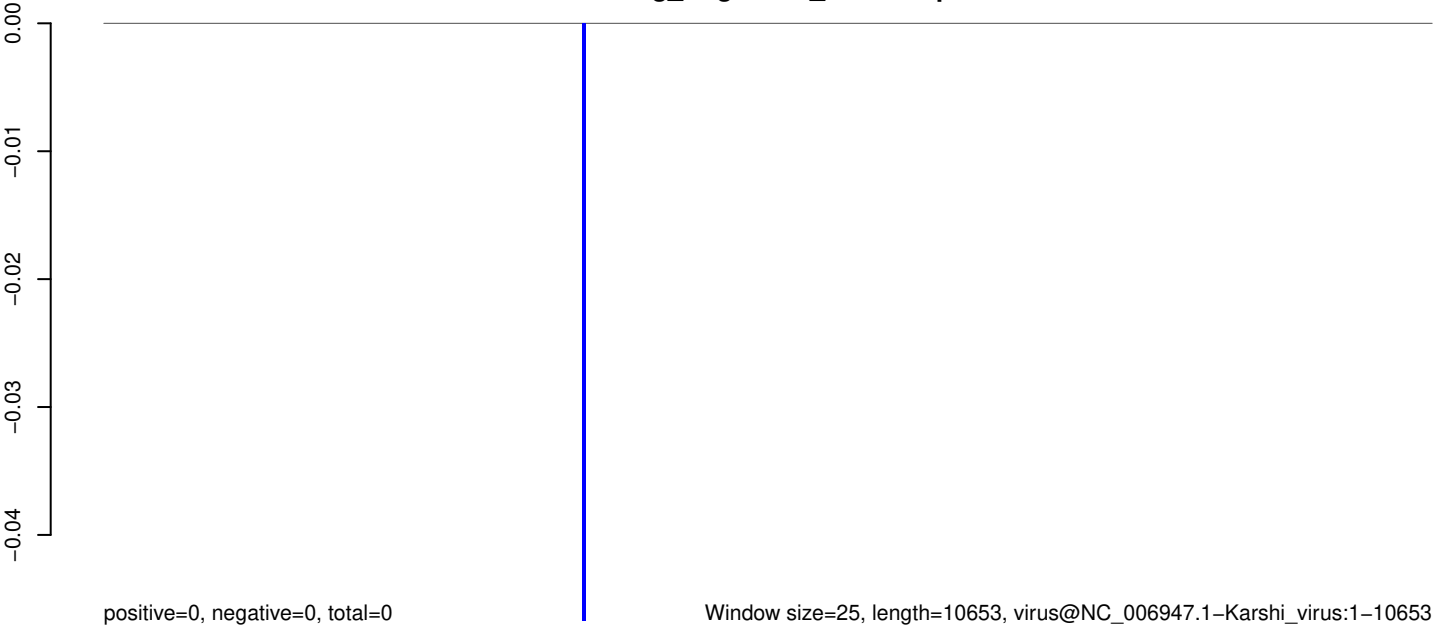
AeAeg_Aag2SINV_dsLuc.18_23.rep



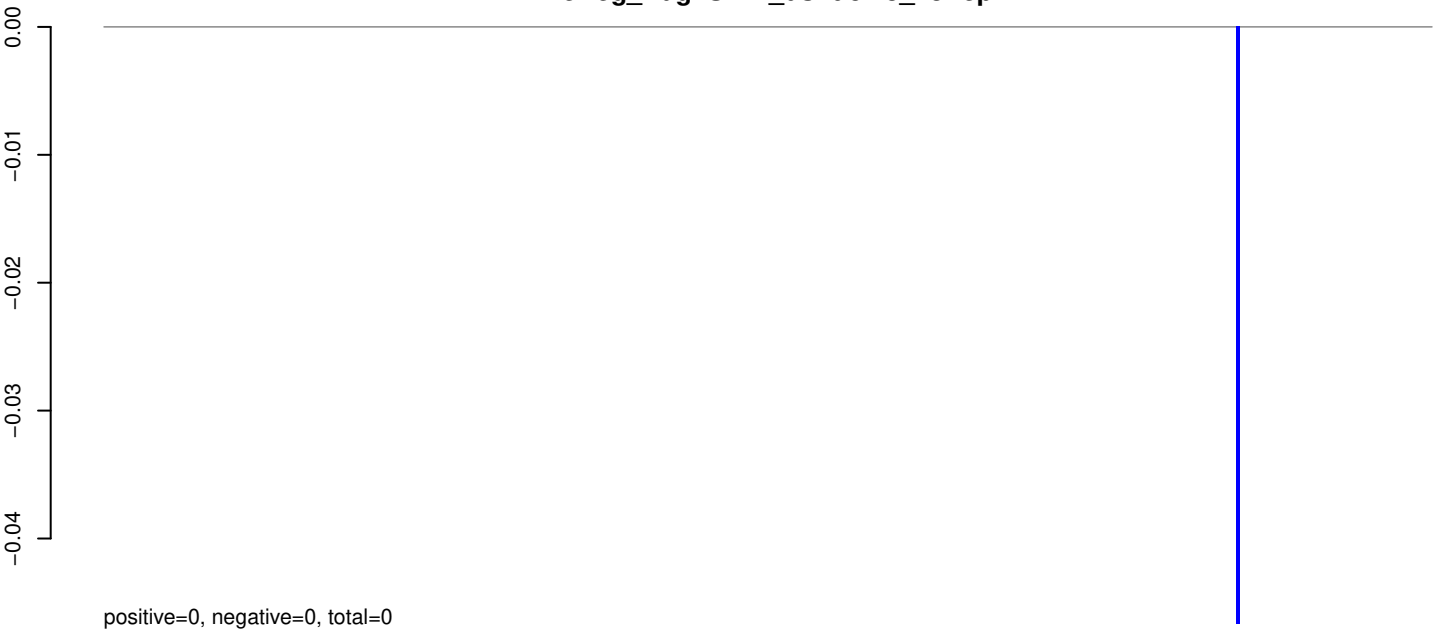
AeAeg_Aag2SINV_dsLuc.24_35.rep



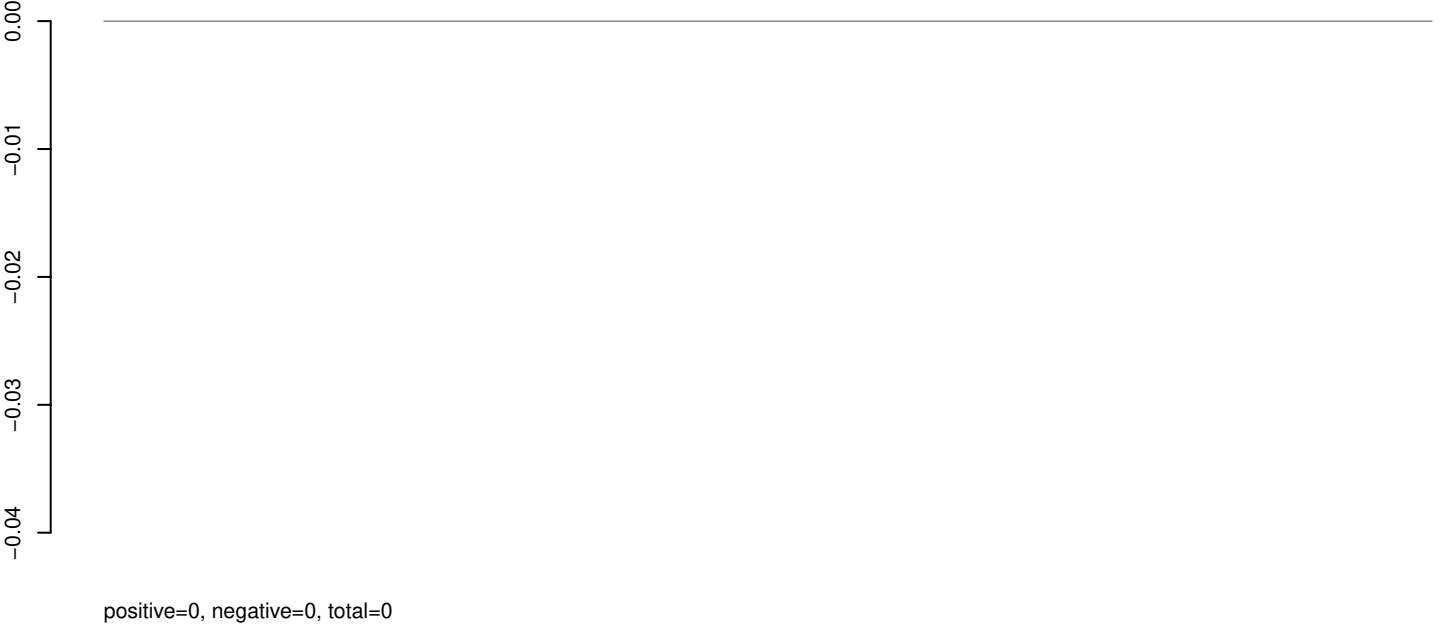
AeAeg_Aag2SINV_dsLuc.rep



AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



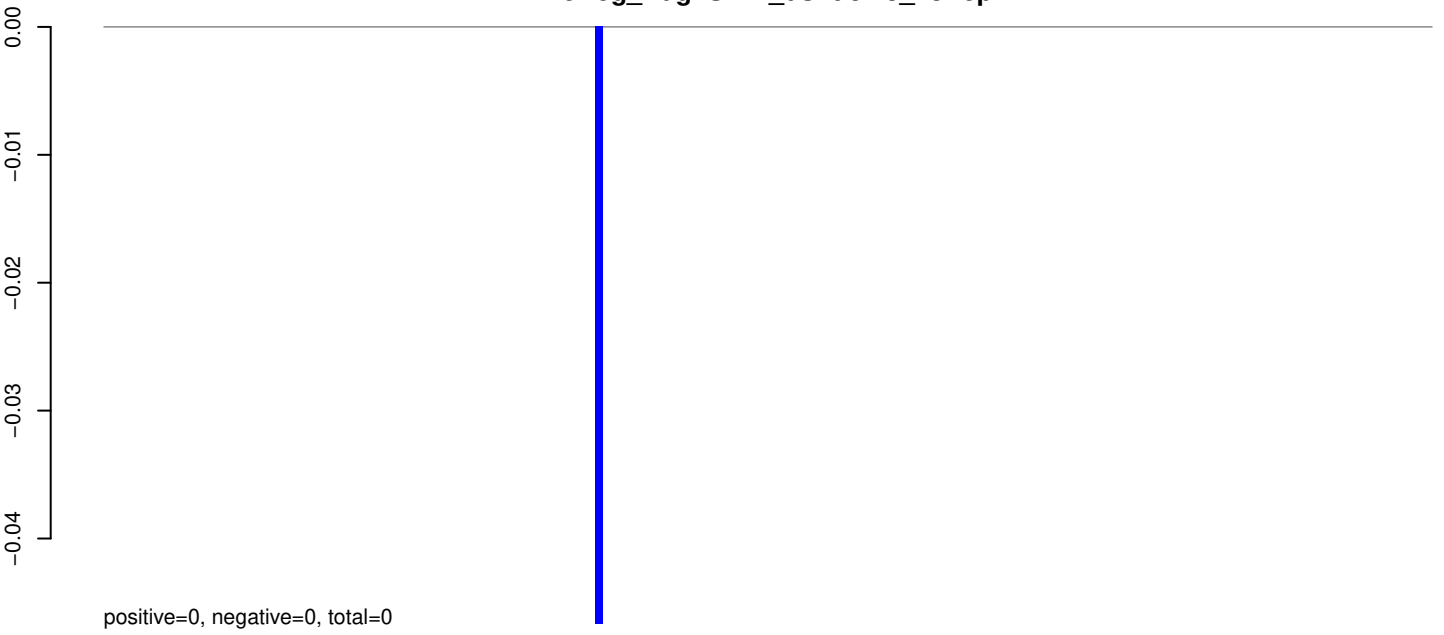
AeAeg_Aag2SINV_dsLuc.rep



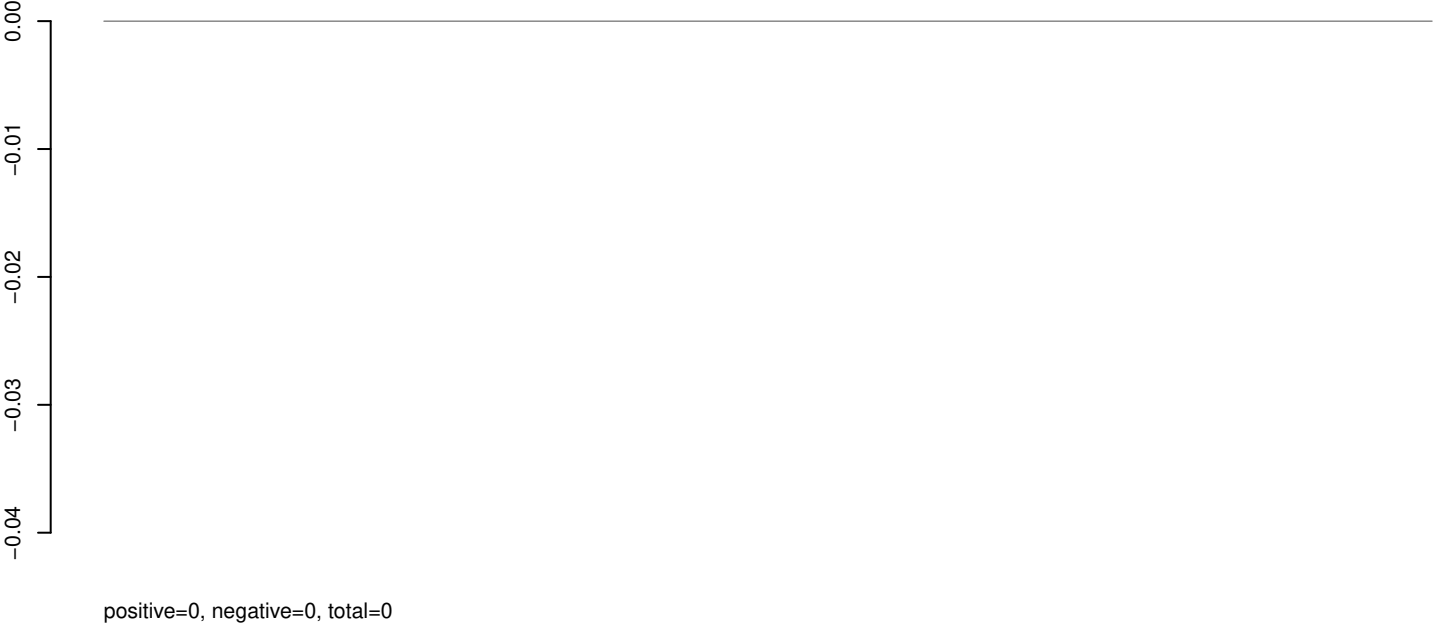
Window size=25, length=10685, virus@NC_004355.1-Alkhurma_virus:1-10685

0 2000 4000 6000 8000 10000

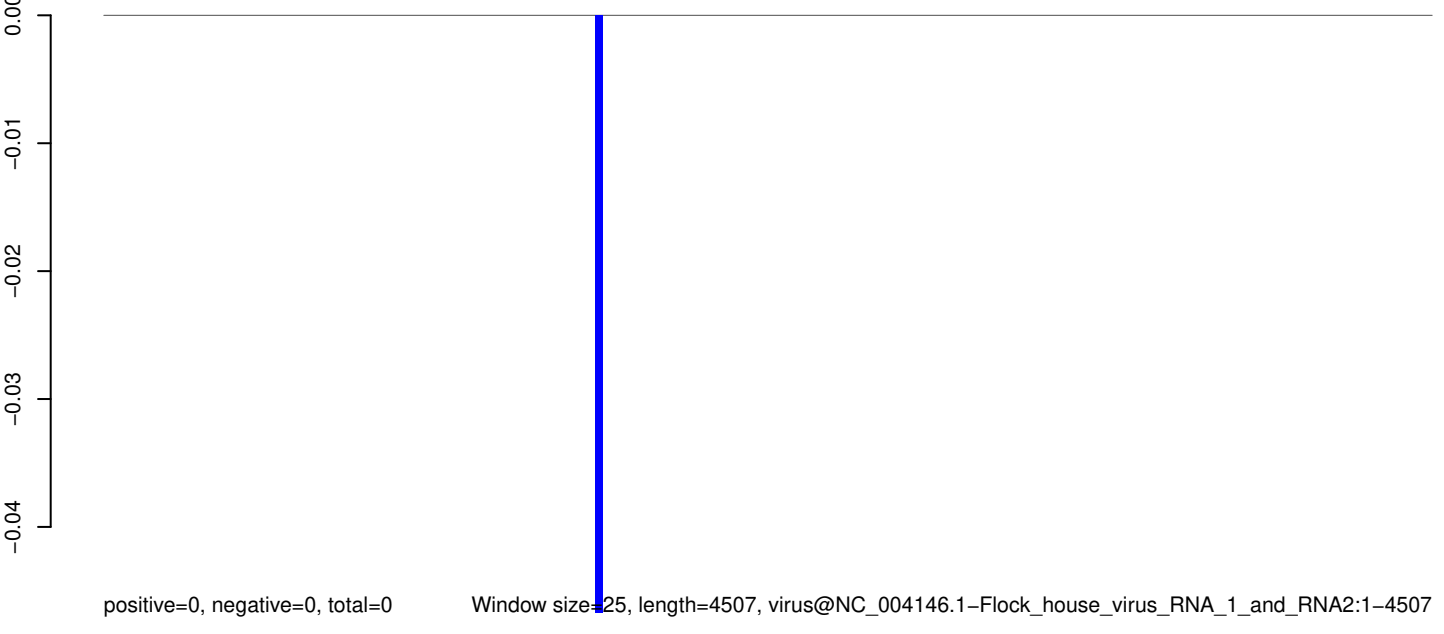
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

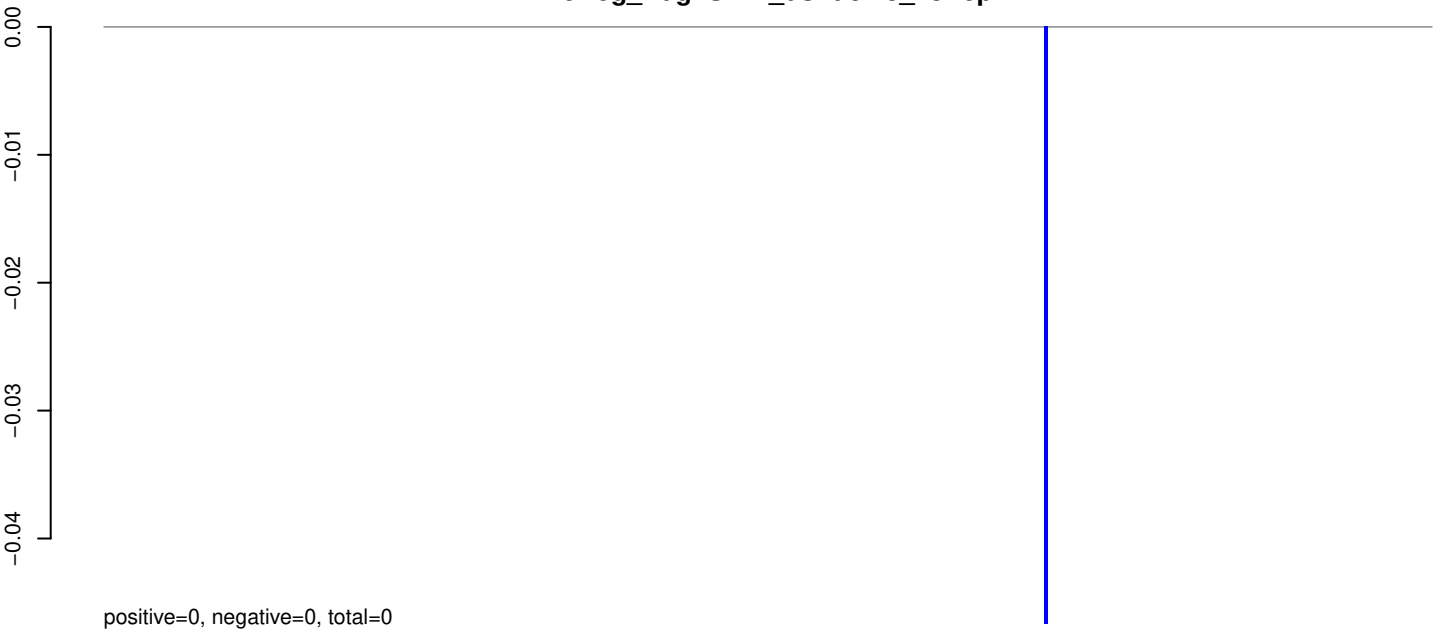


AeAeg_Aag2SINV_dsLuc.rep



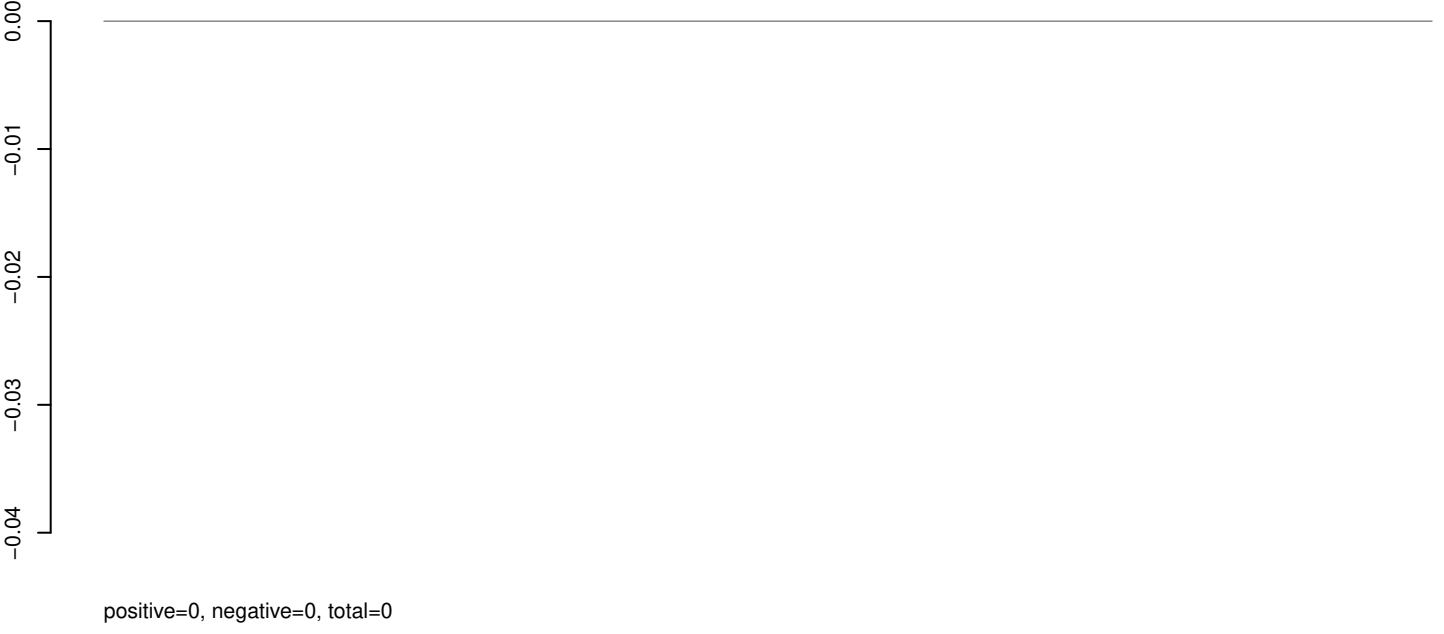
0 1000 2000 3000 4000

AeAeg_Aag2SINV_dsLuc.18_23.rep



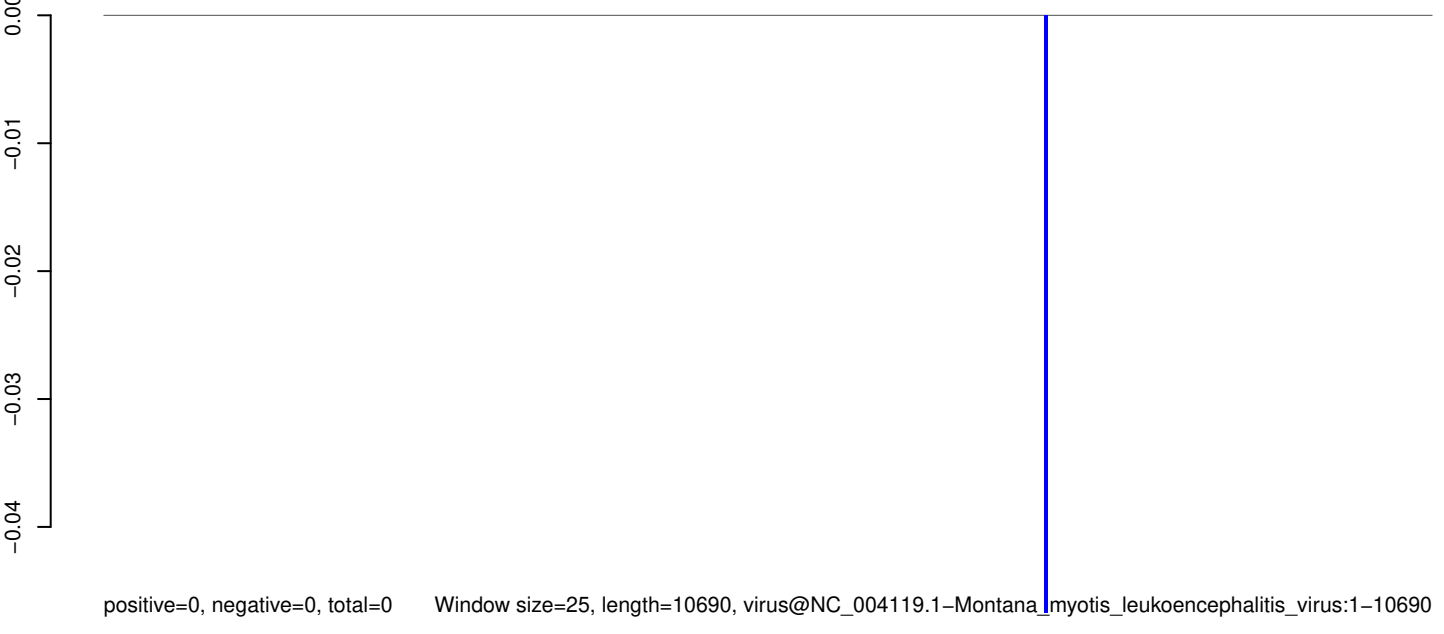
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.rep

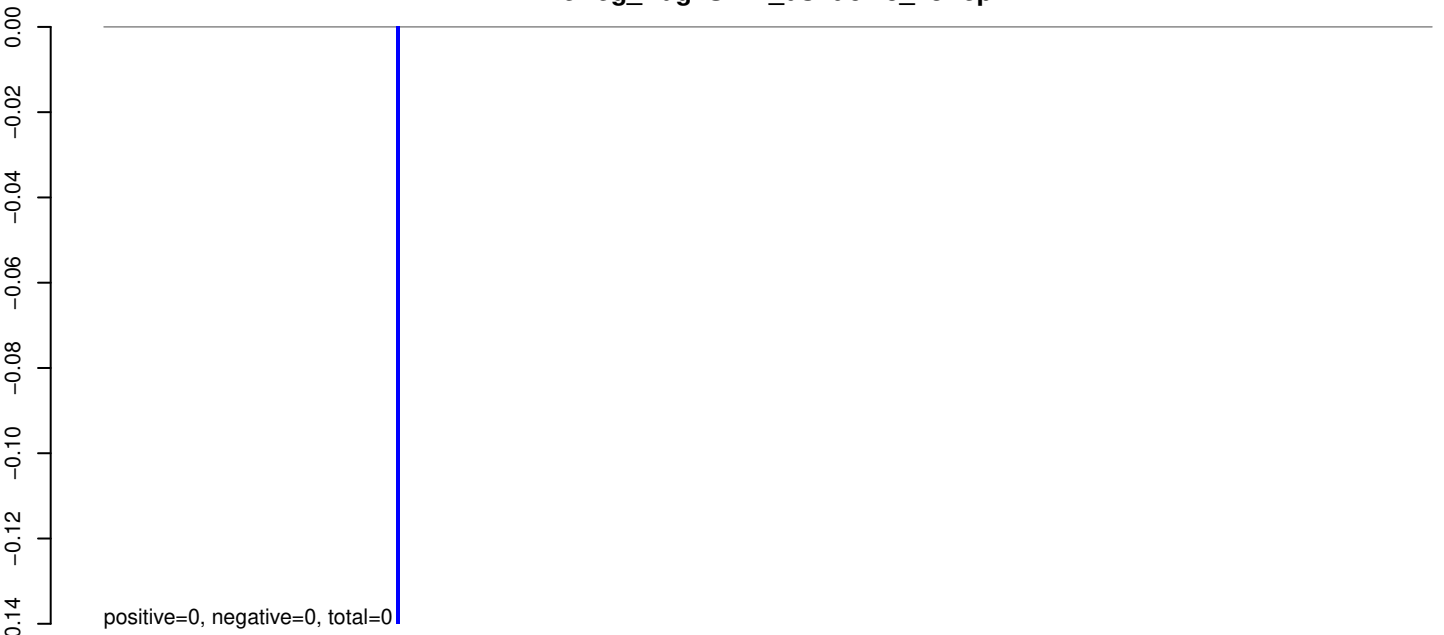


positive=0, negative=0, total=0

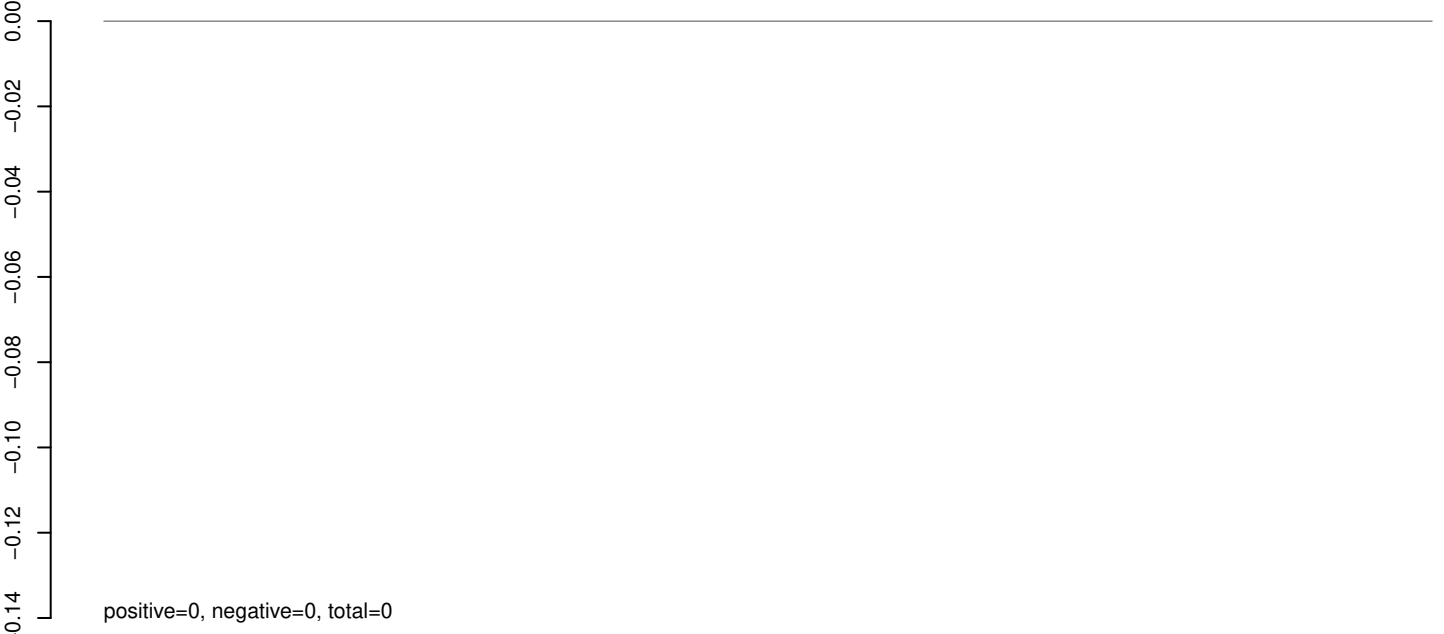
Window size=25, length=10690, virus@NC_004119.1-Montana_myotis_leukoencephalitis_virus:1-10690

0 2000 4000 6000 8000 10000

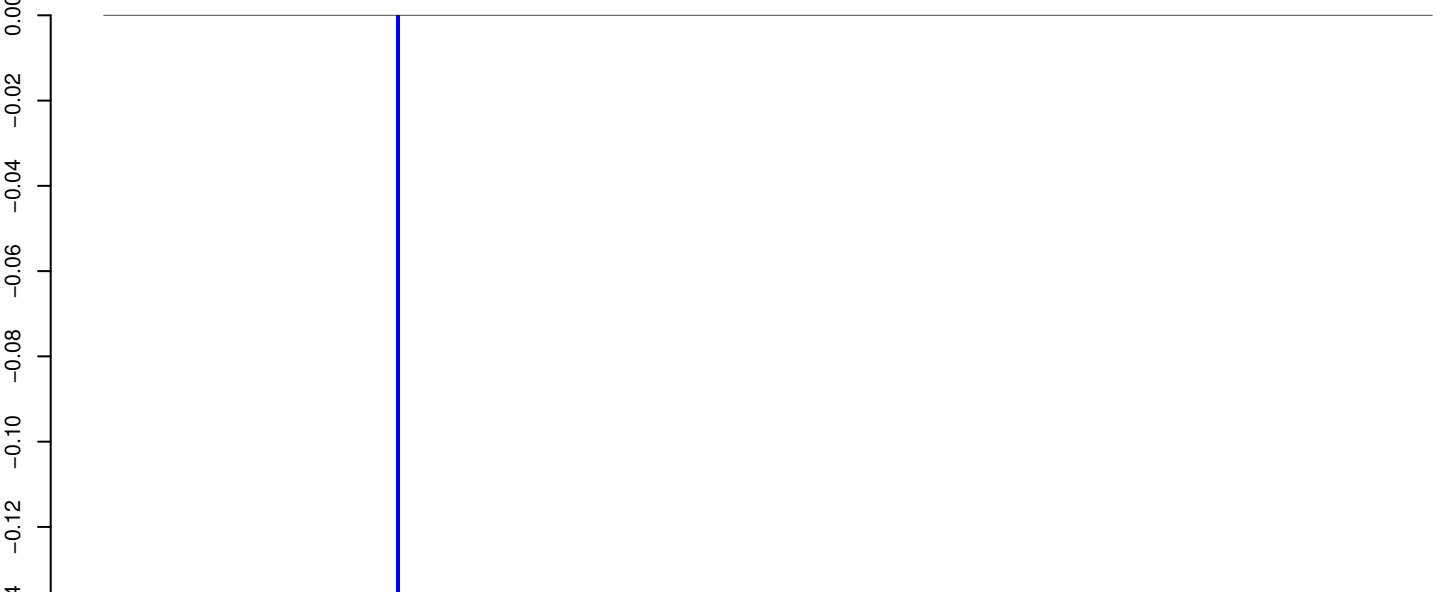
AeAeg_Aag2SINV_dsLuc.18_23.rep



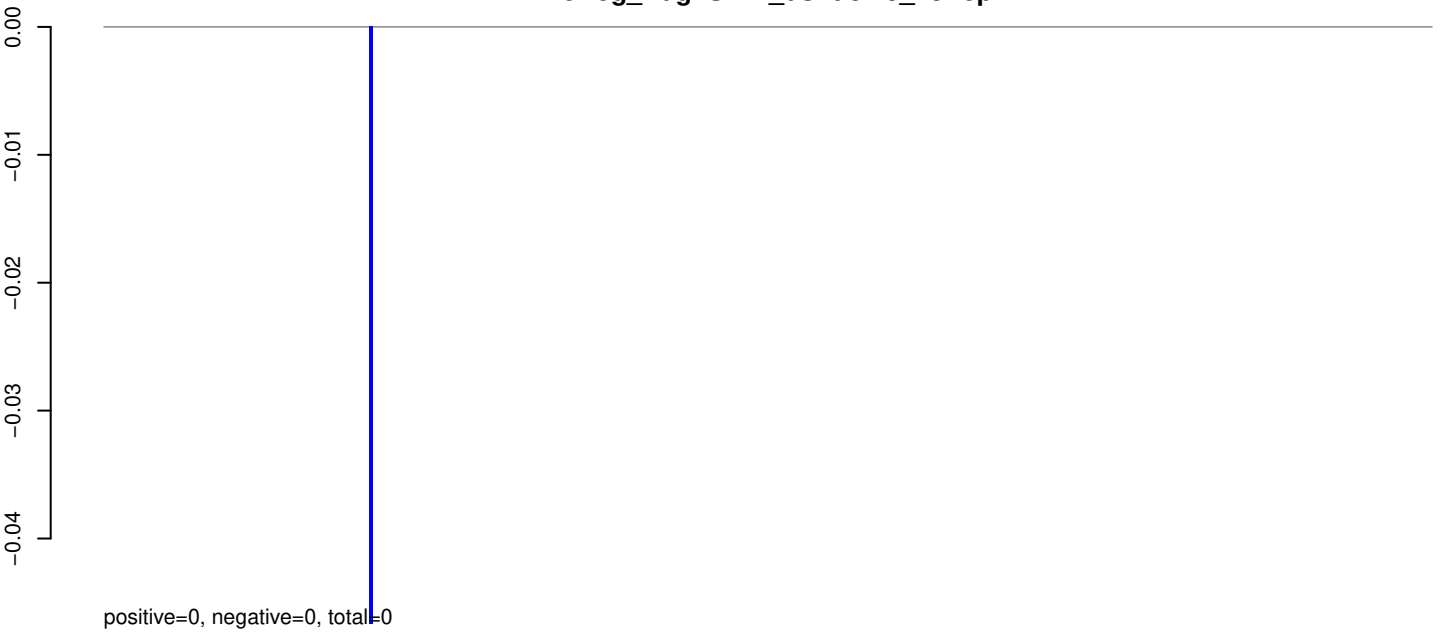
AeAeg_Aag2SINV_dsLuc.24_35.rep



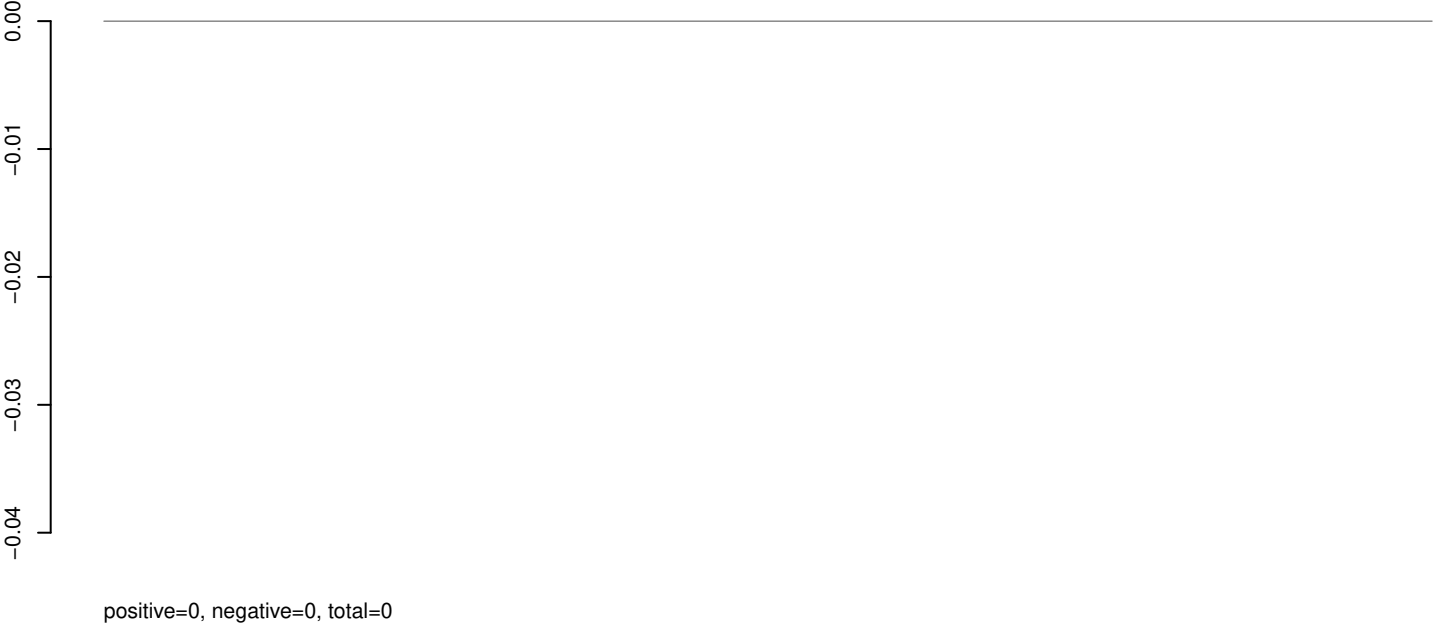
AeAeg_Aag2SINV_dsLuc.rep



AeAeg_Aag2SINV_dsLuc.18_23.rep



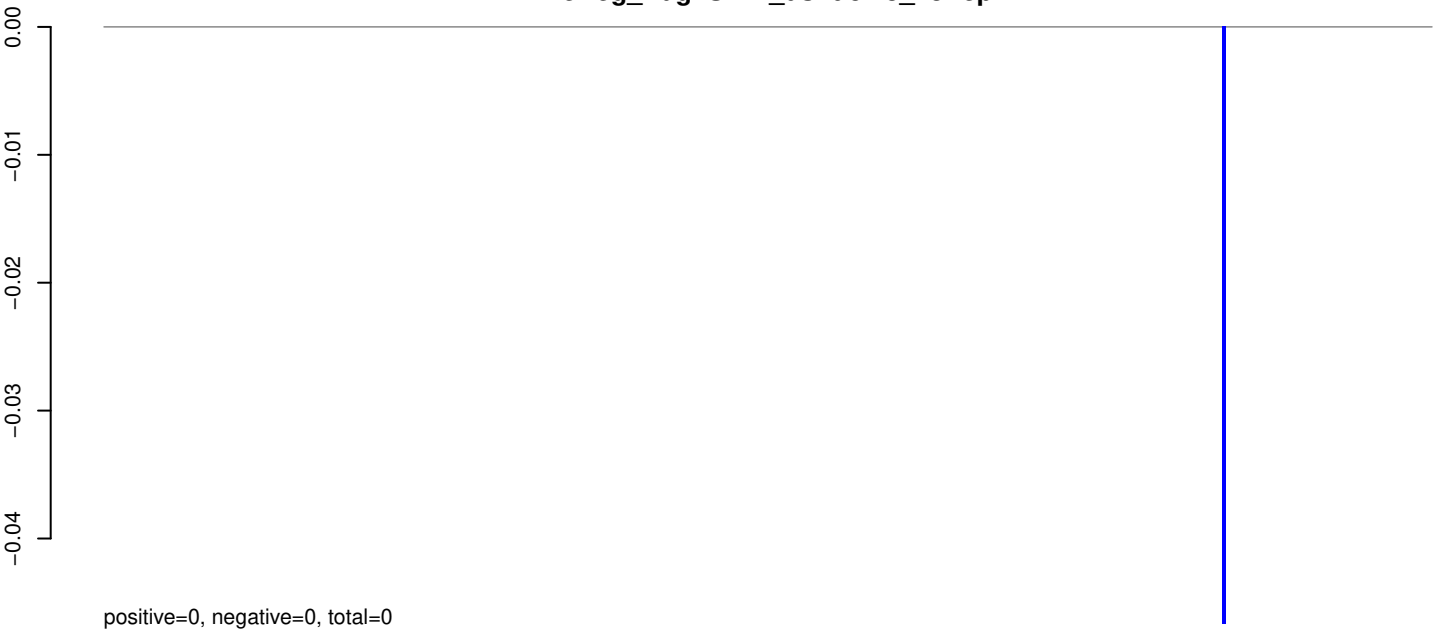
AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep

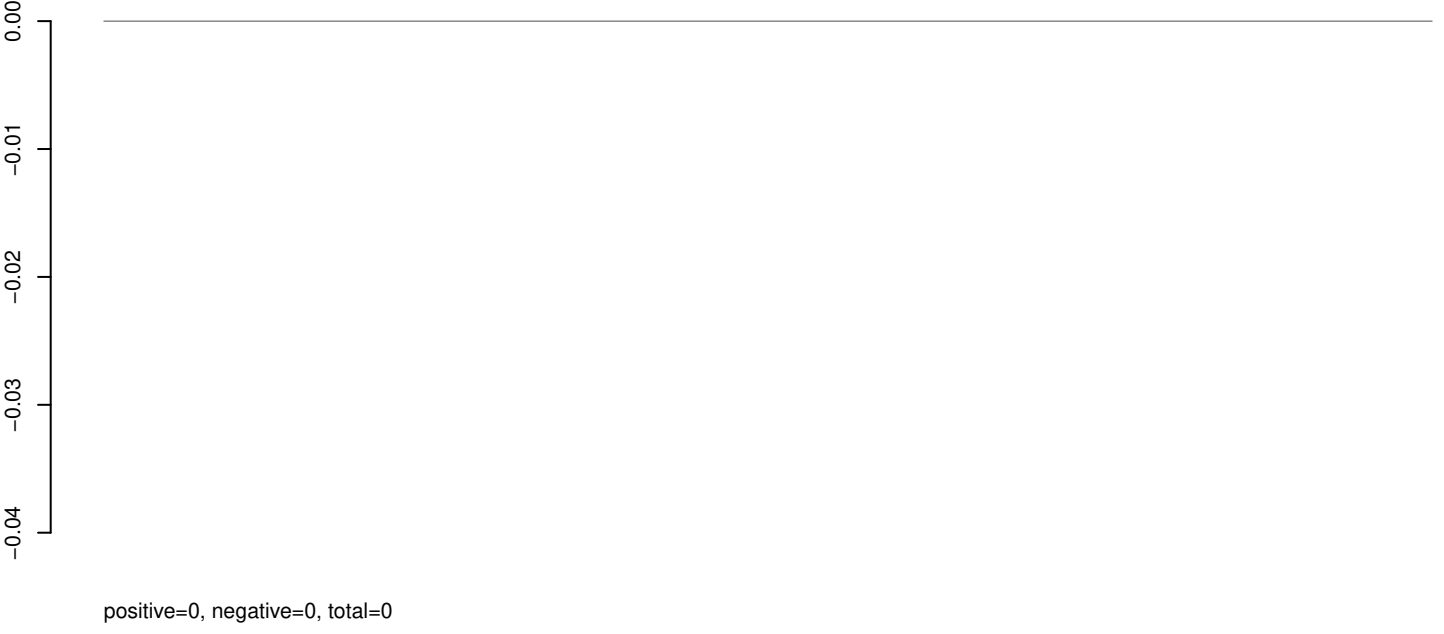


AeAeg_Aag2SINV_dsLuc.18_23.rep



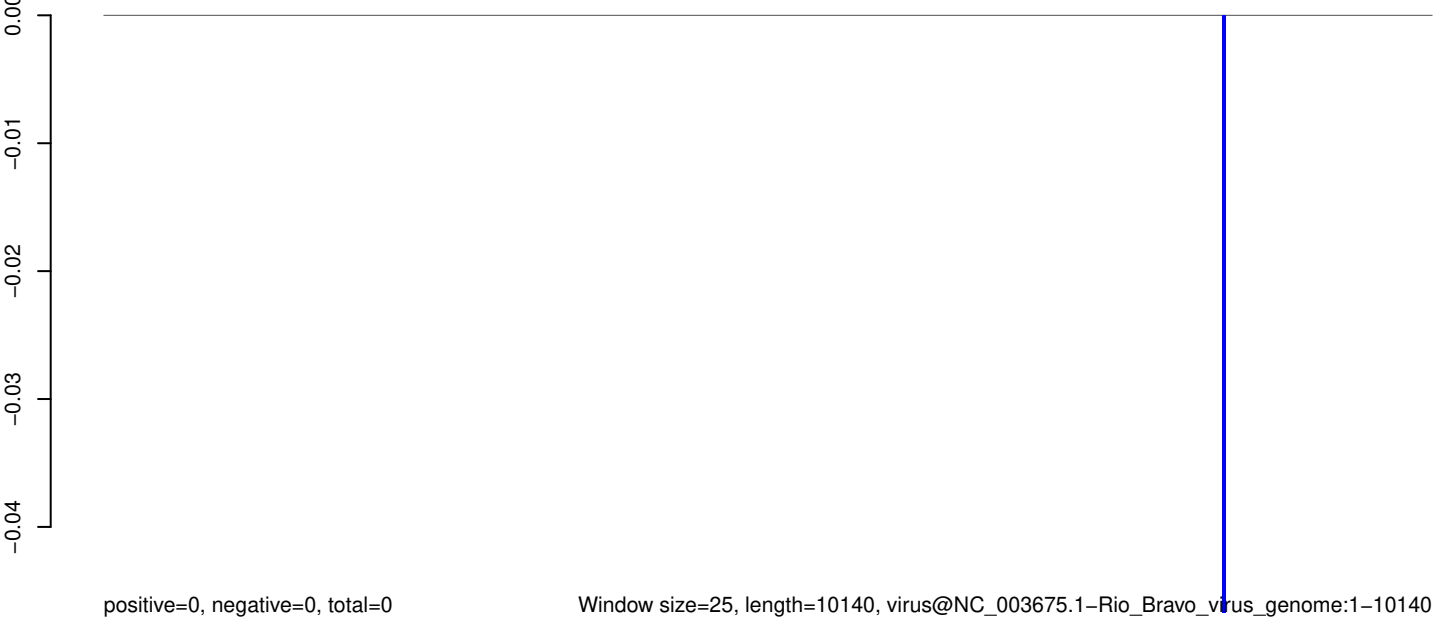
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep



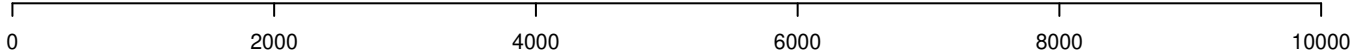
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.rep

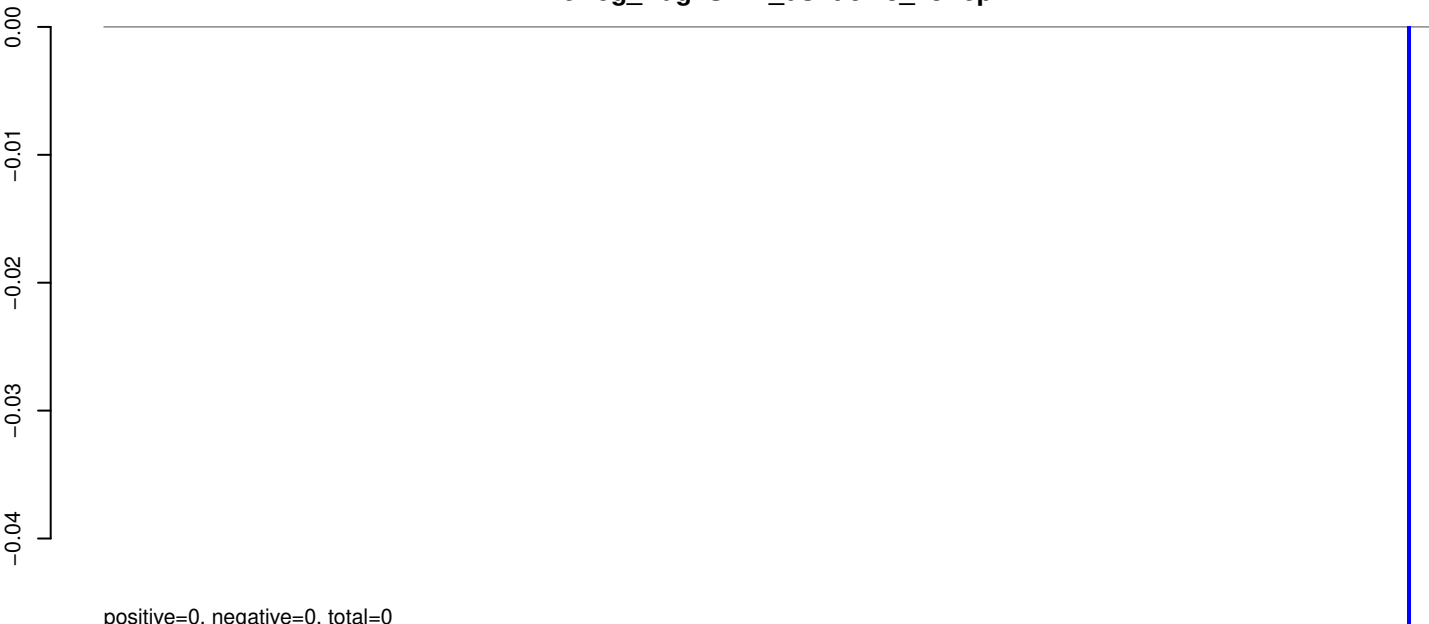


positive=0, negative=0, total=0

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

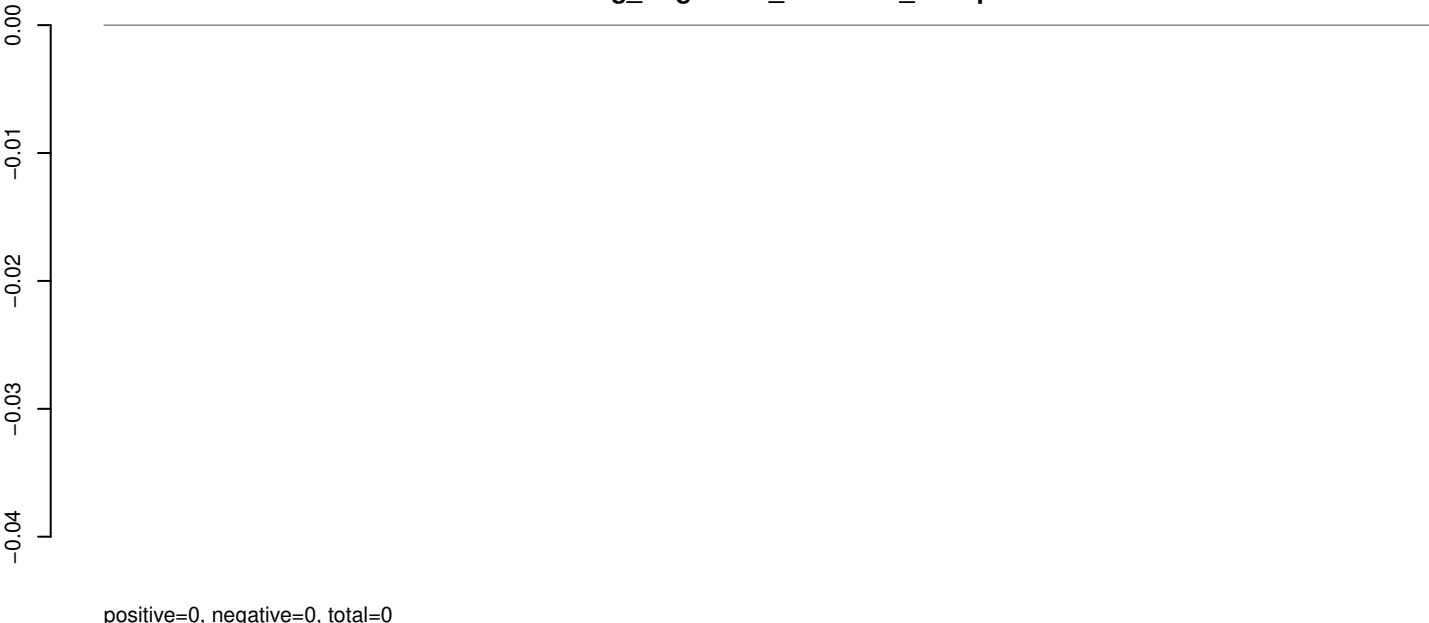


AeAeg_Aag2SINV_dsLuc.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.rep

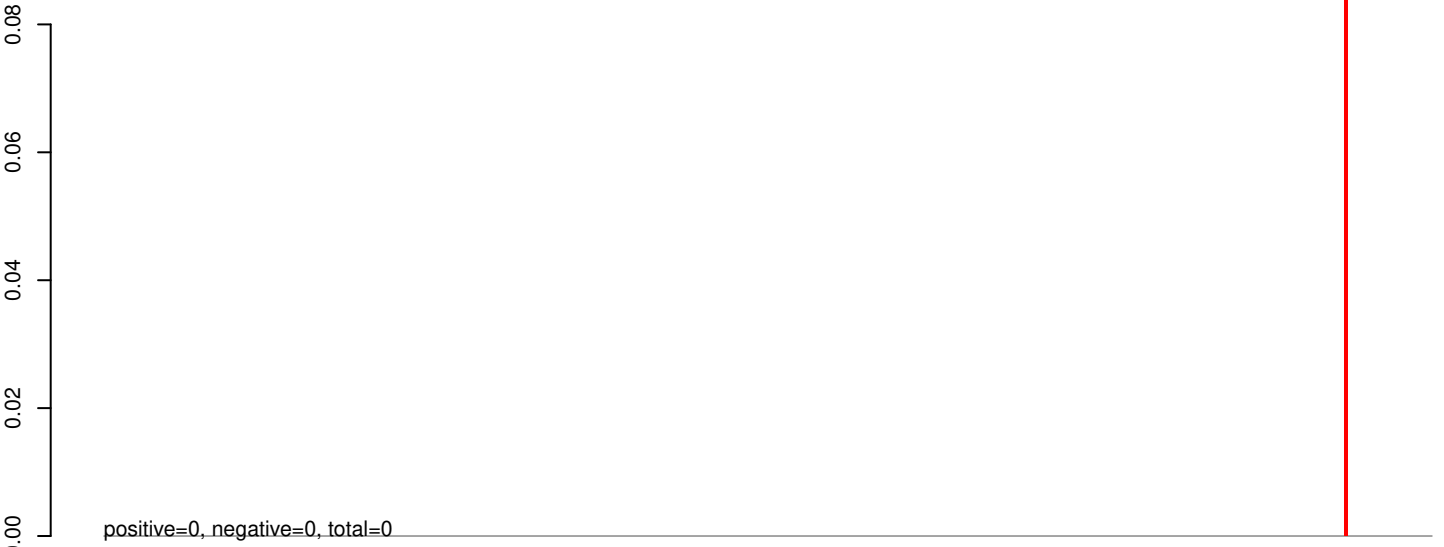


positive=0, negative=0, total=0

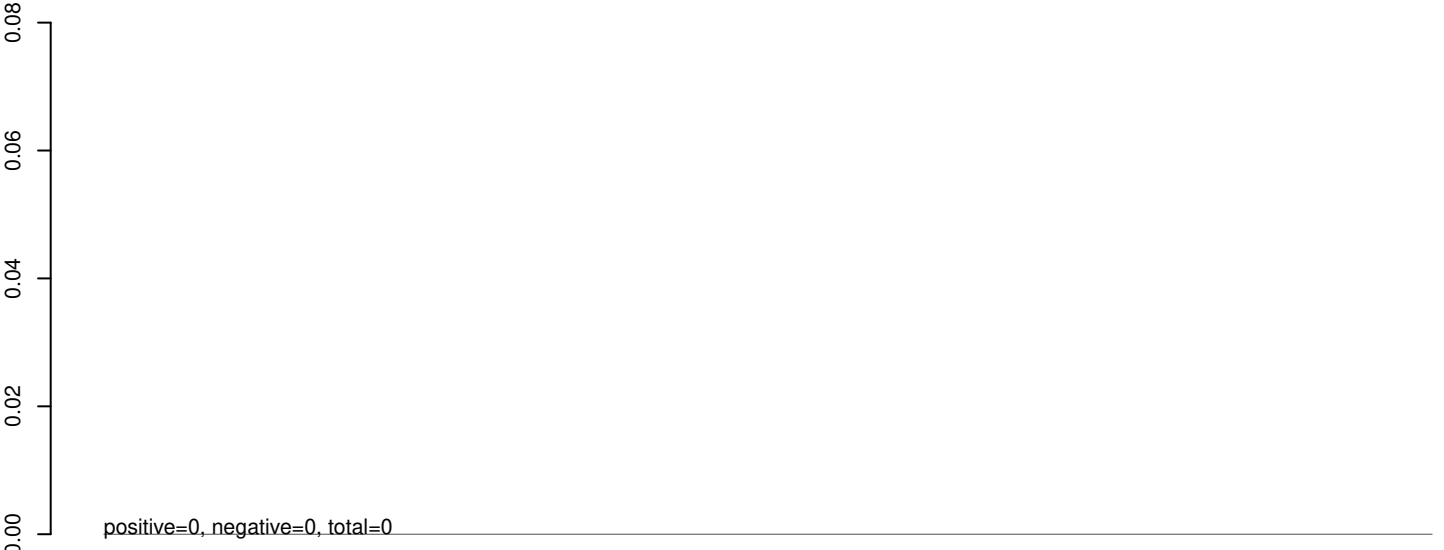
Window size=25, length=10600, virus@NC_003635.1-Modoc_virus:1-10600

0 2000 4000 6000 8000 10000

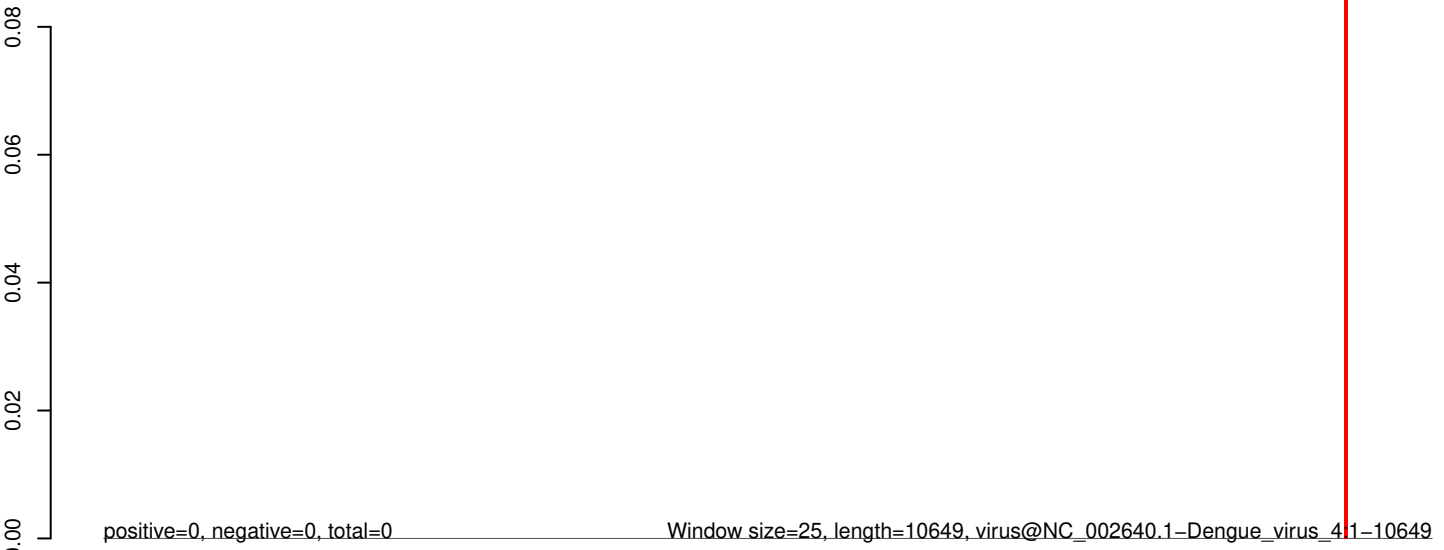
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

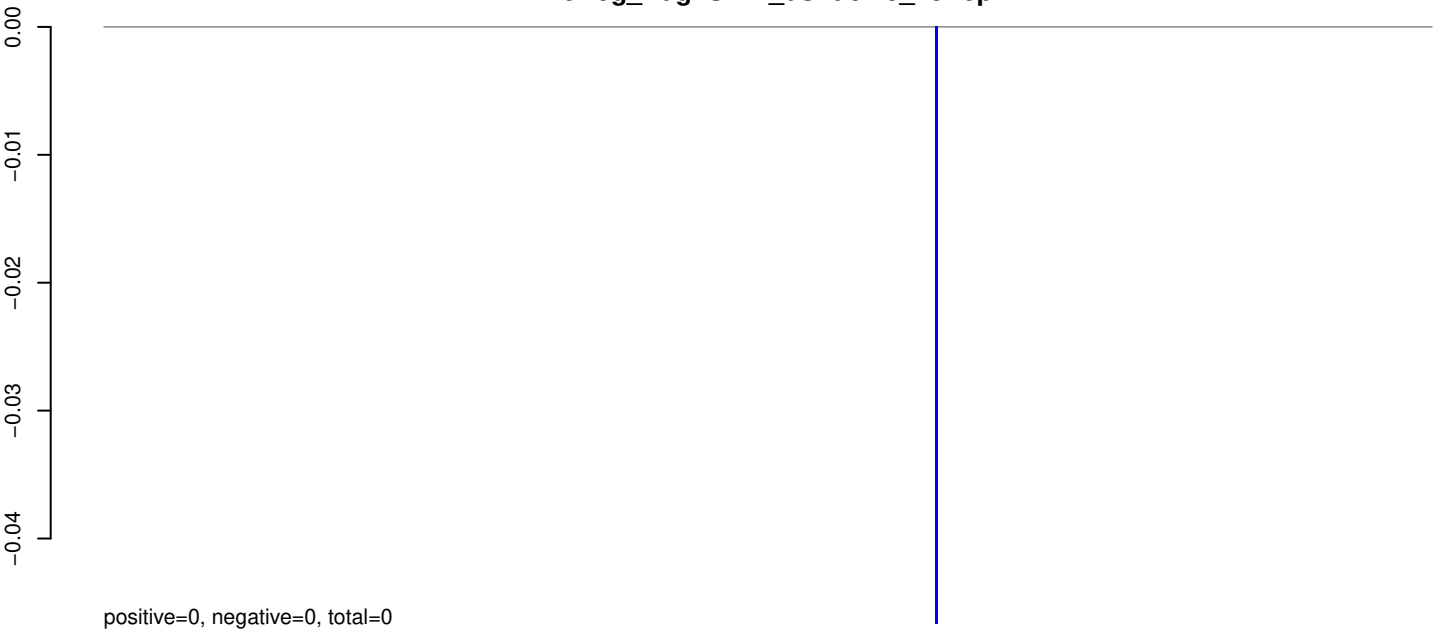


AeAeg_Aag2SINV_dsLuc.rep

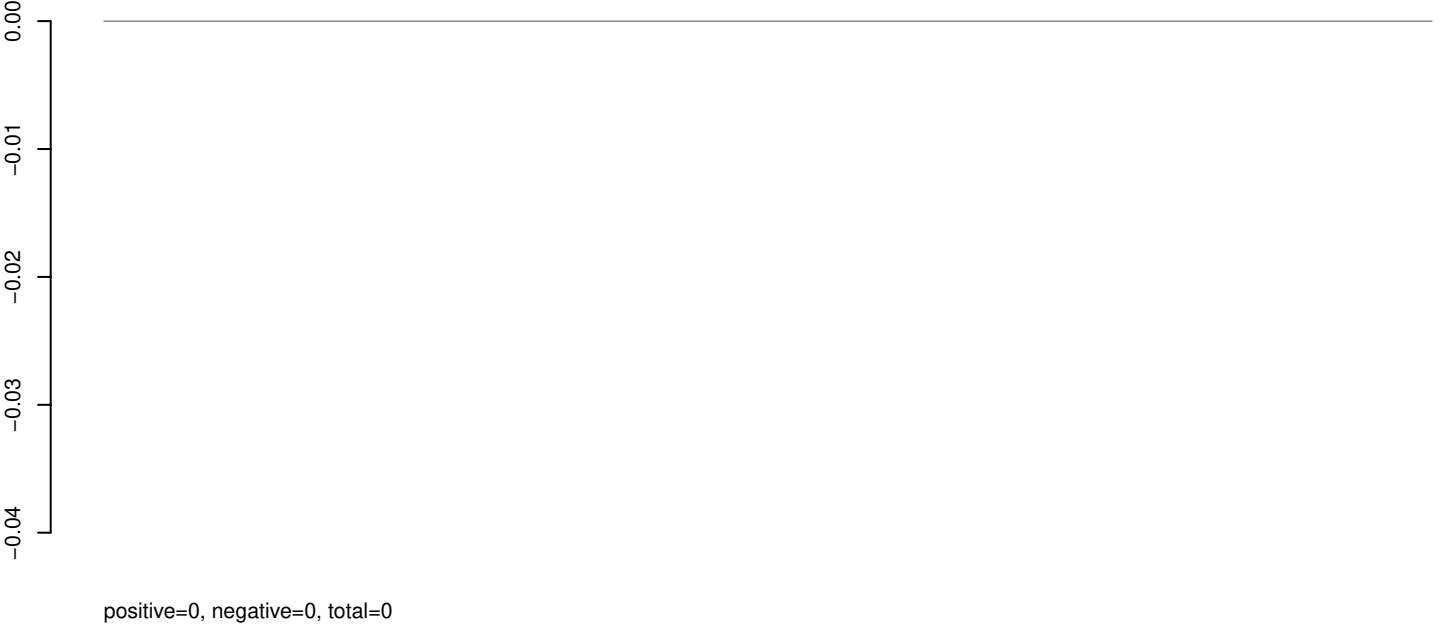


0 2000 4000 6000 8000 10000

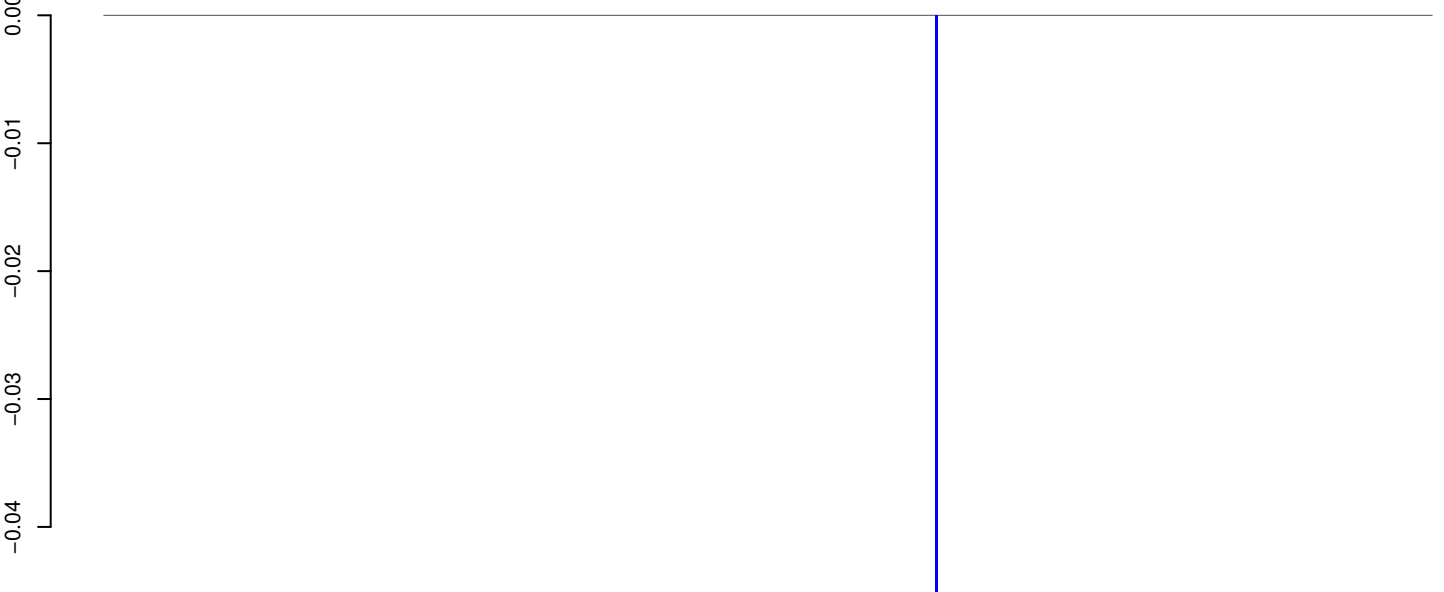
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



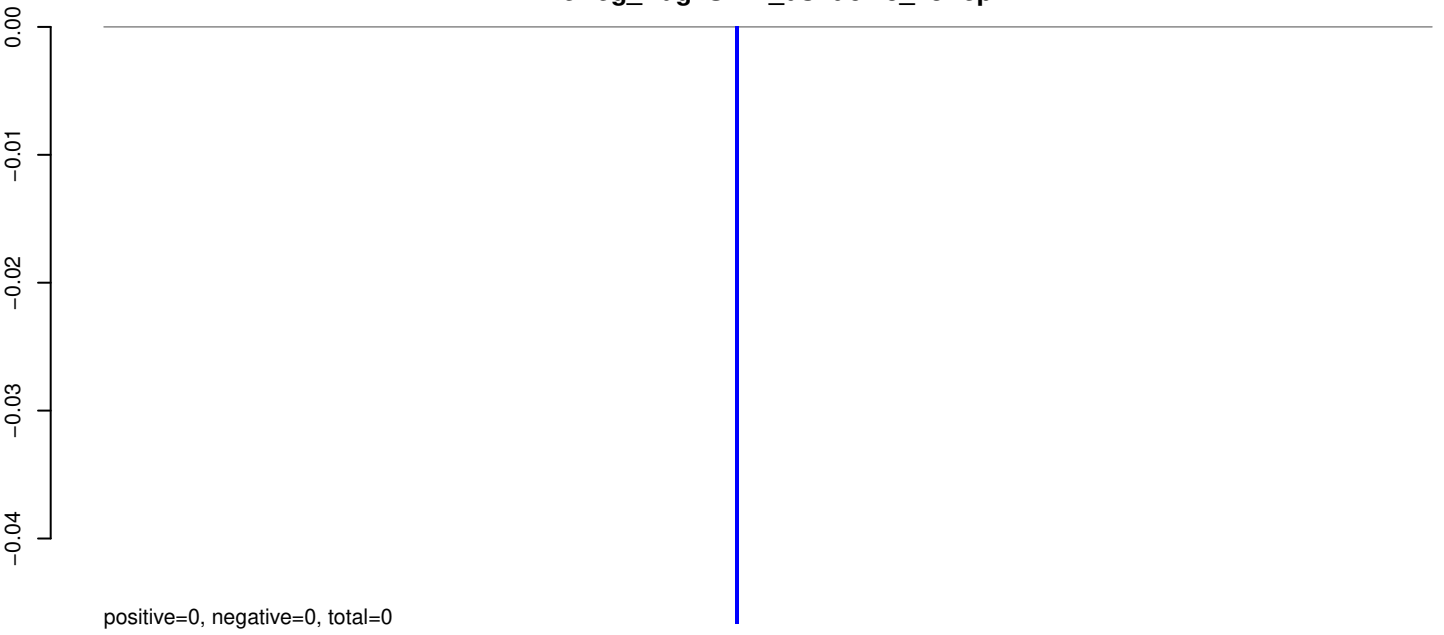
AeAeg_Aag2SINV_dsLuc.rep



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

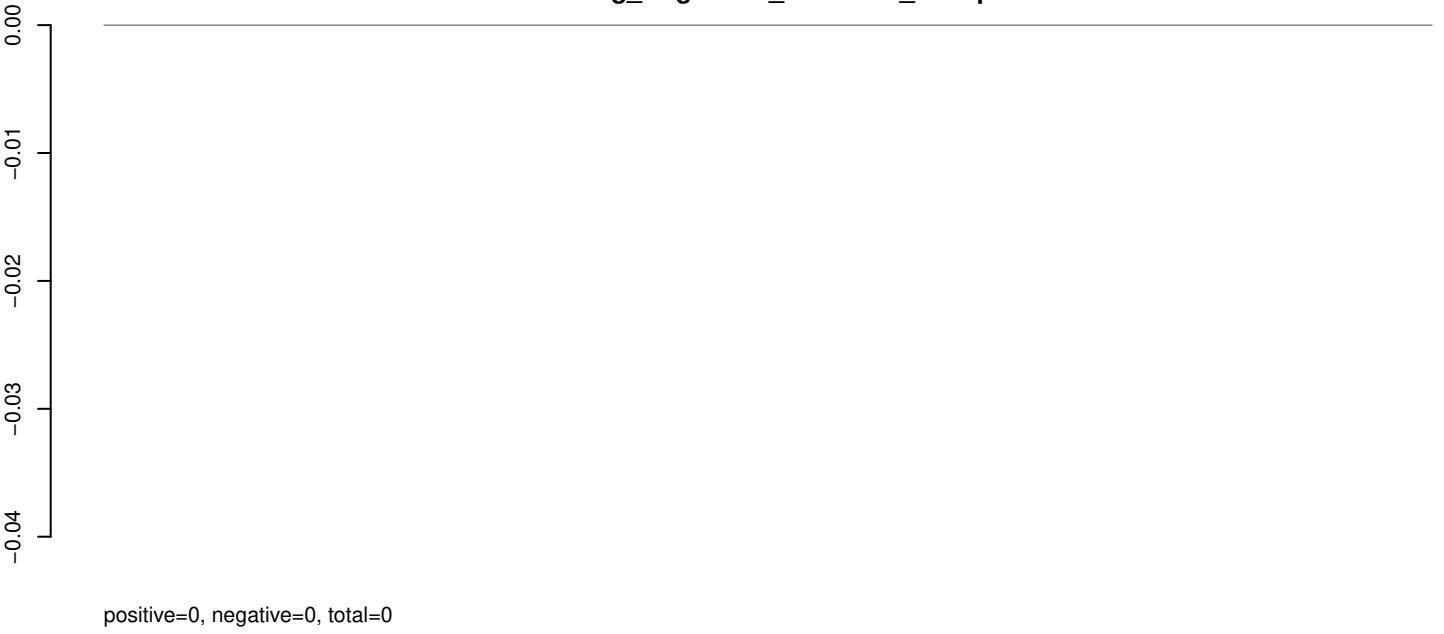
0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_dsLuc.18_23.rep



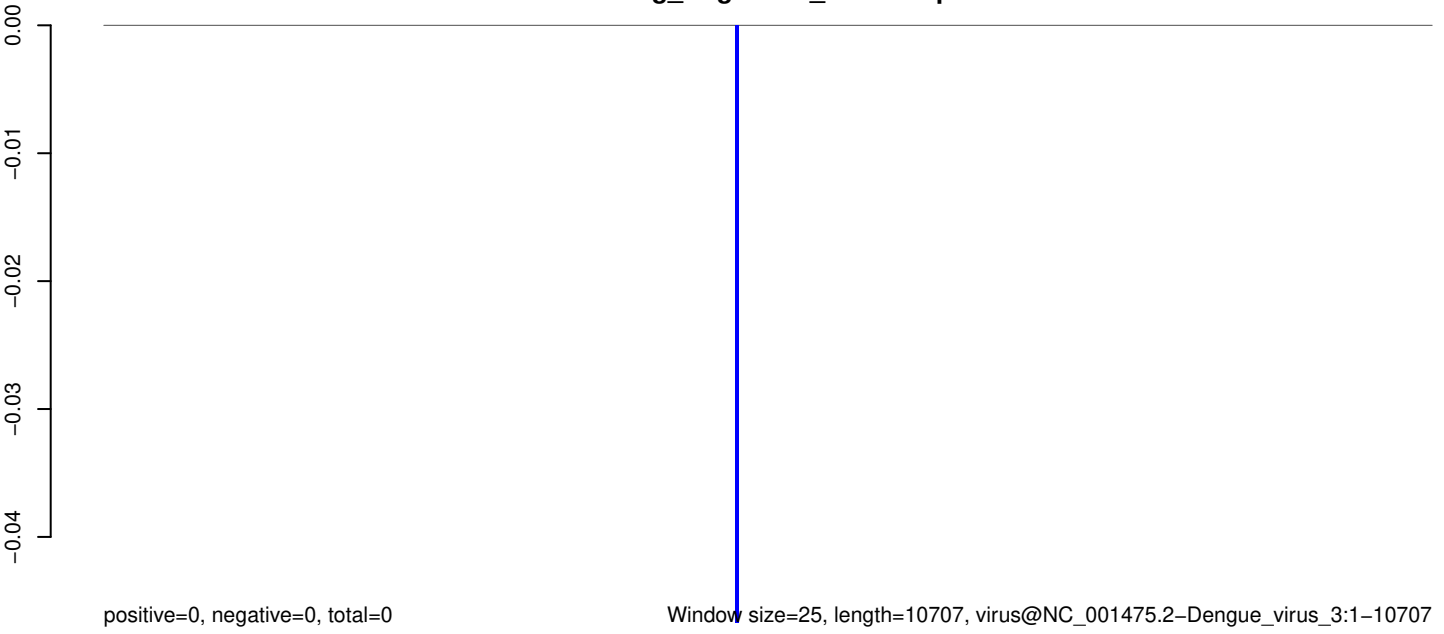
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.rep

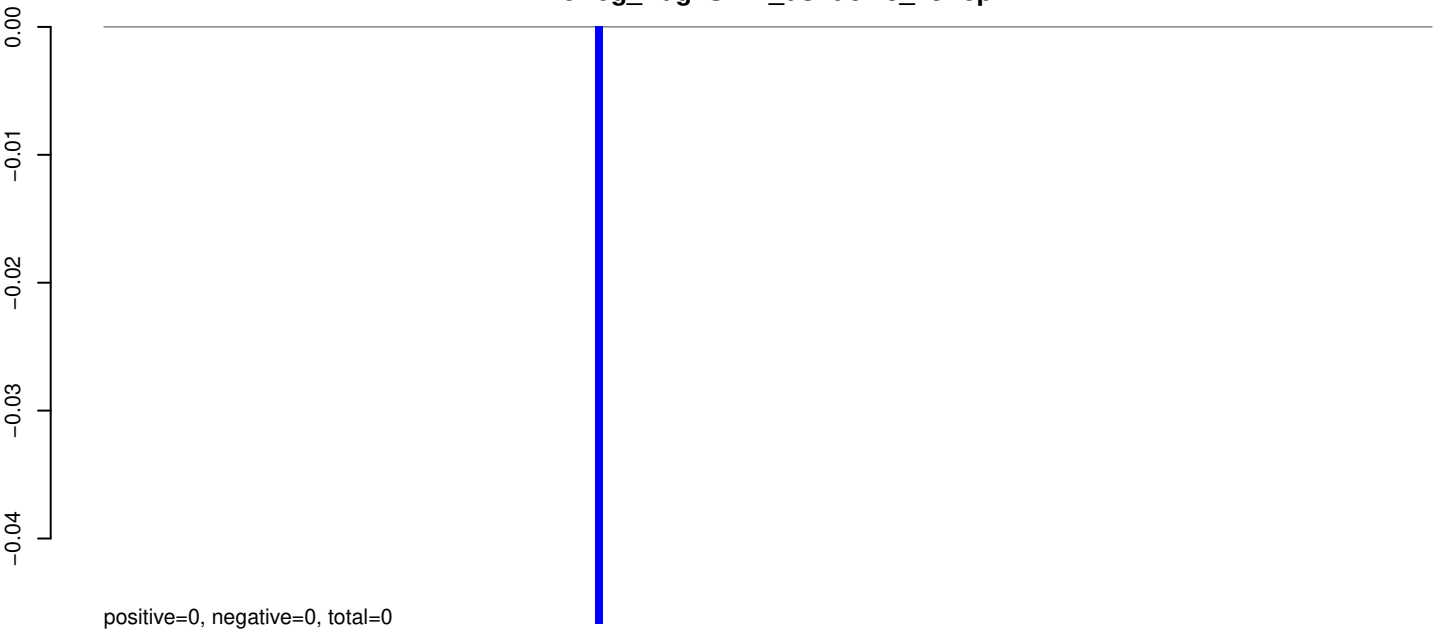


positive=0, negative=0, total=0

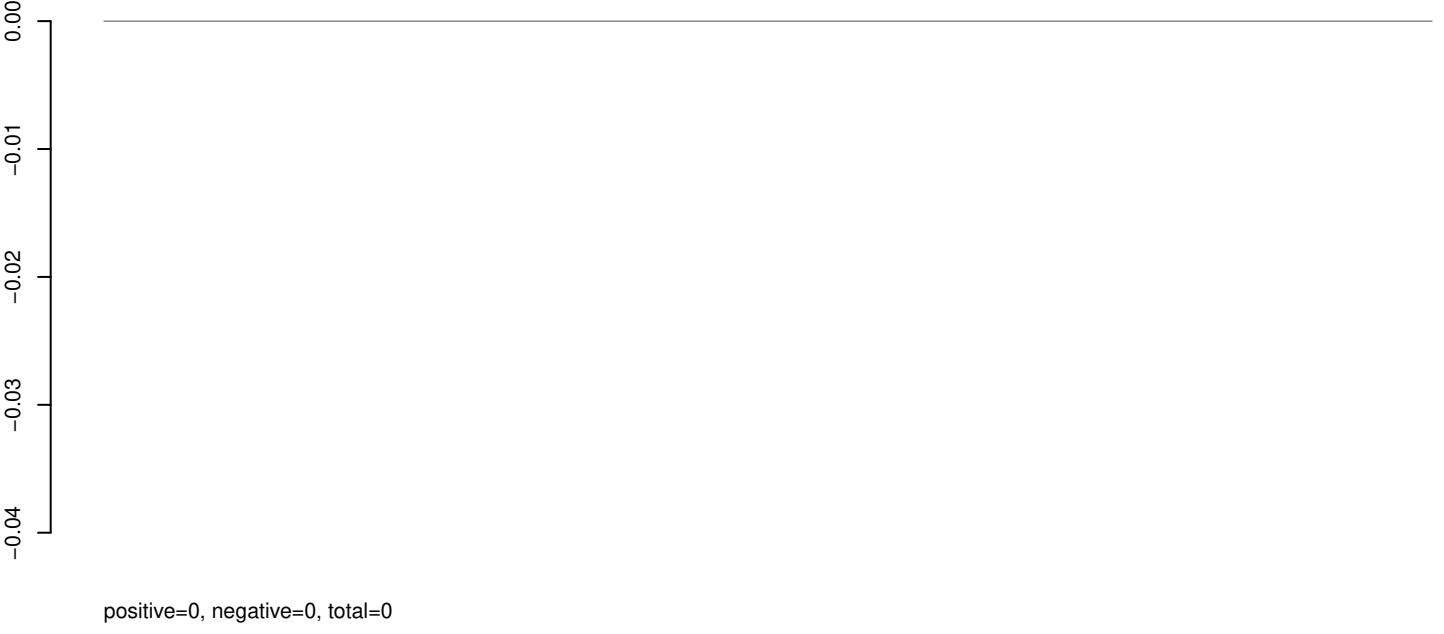
Window size=25, length=10707, virus@NC_001475.2-Dengue_virus_3:1-10707

0 2000 4000 6000 8000 10000

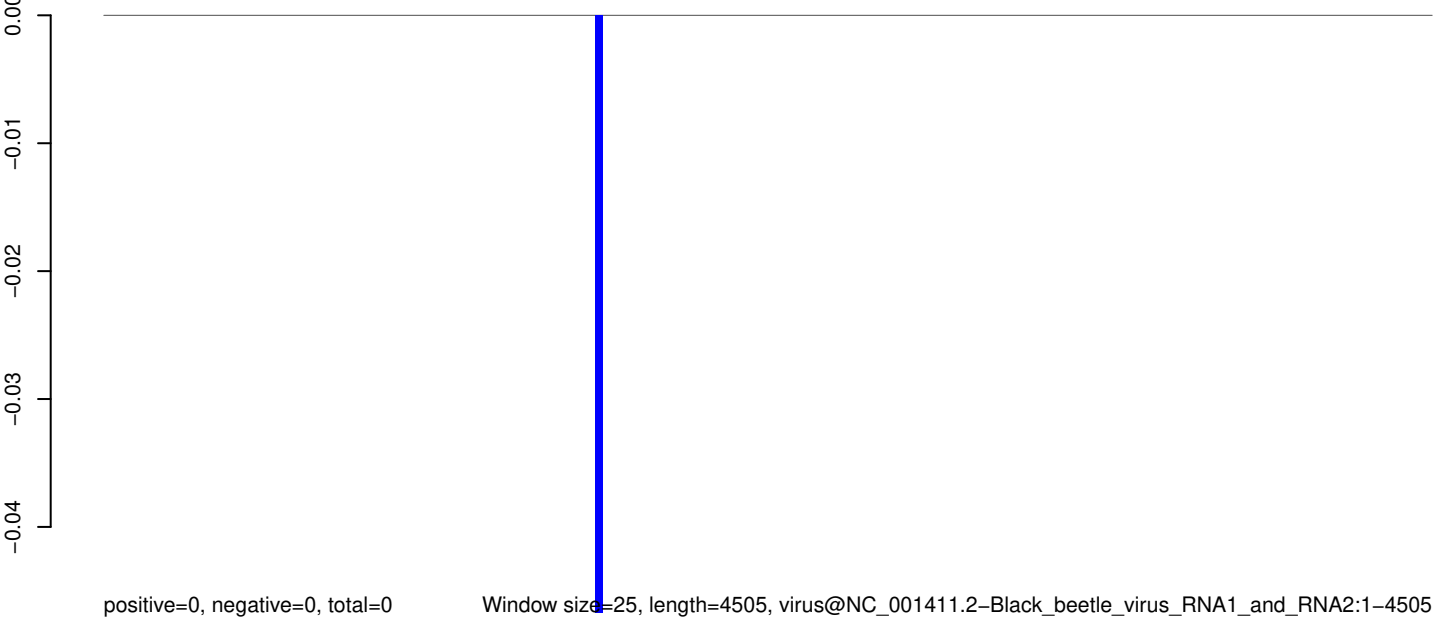
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



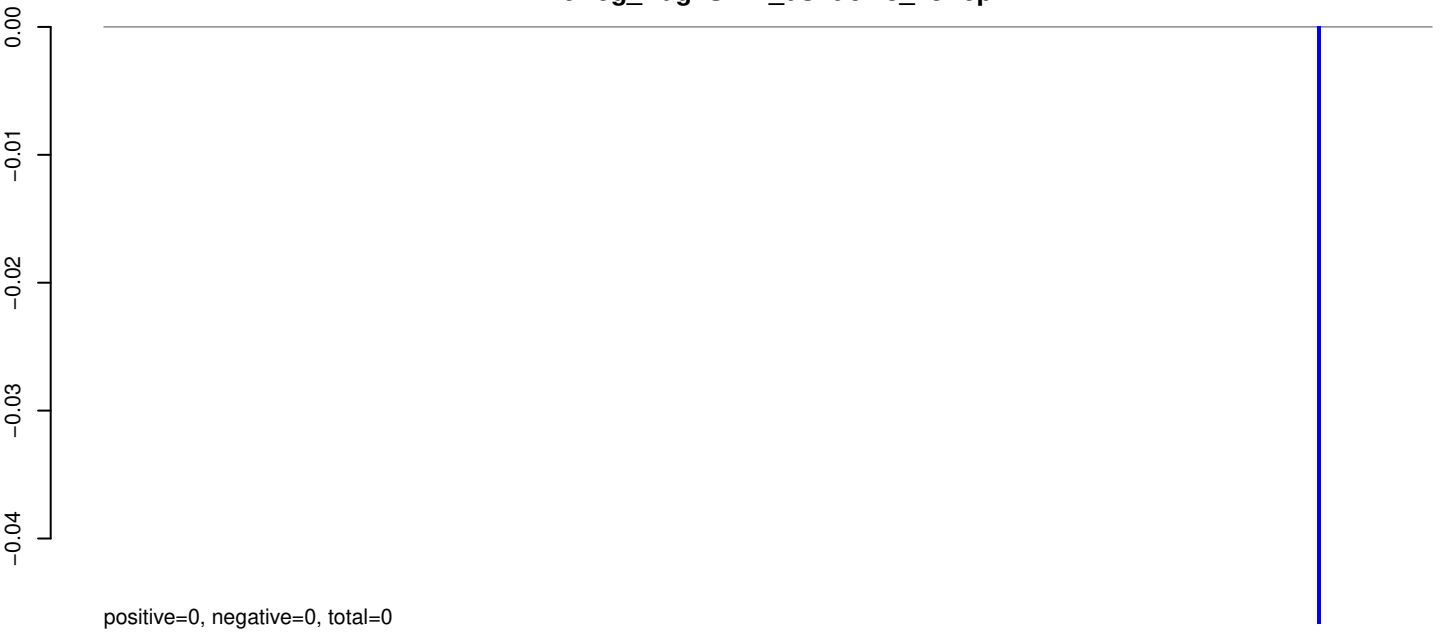
AeAeg_Aag2SINV_dsLuc.rep



positive=0, negative=0, total=0 Window size=25, length=4505, virus@NC_001411.2-Black_beetle_virus_RNA1_and_RNA2:1-4505

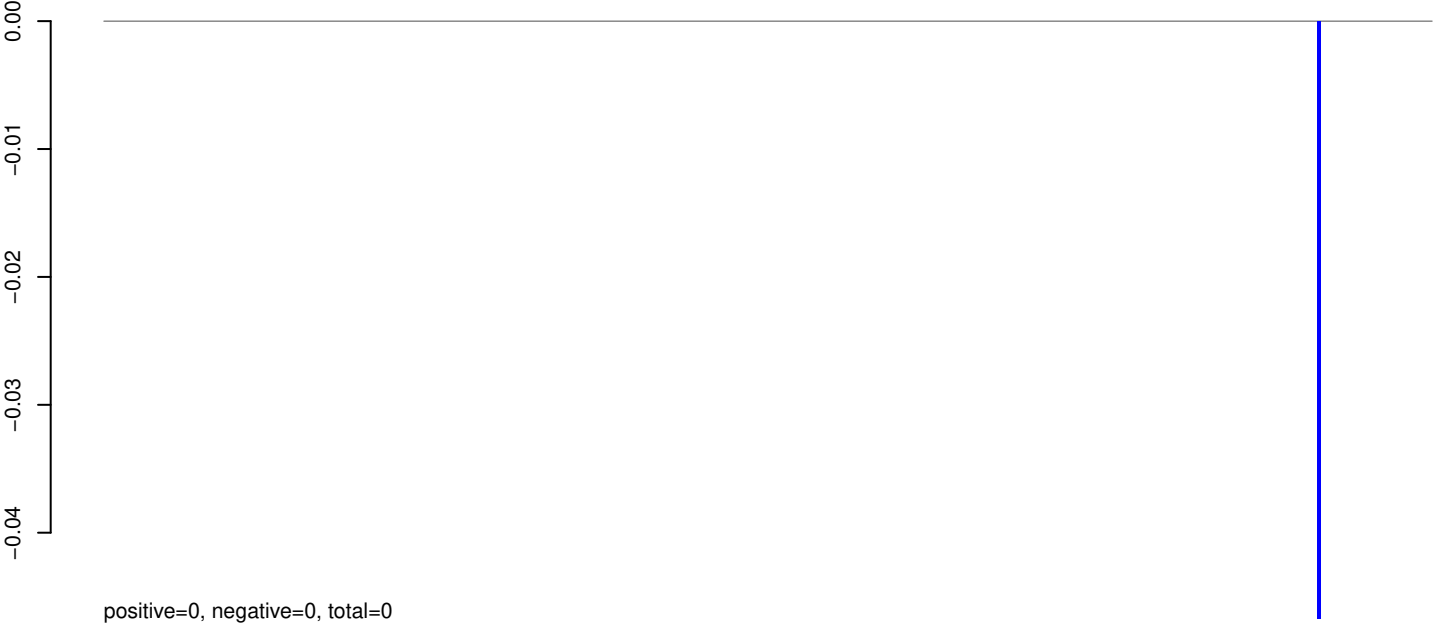
0 1000 2000 3000 4000

AeAeg_Aag2SINV_dsLuc.18_23.rep



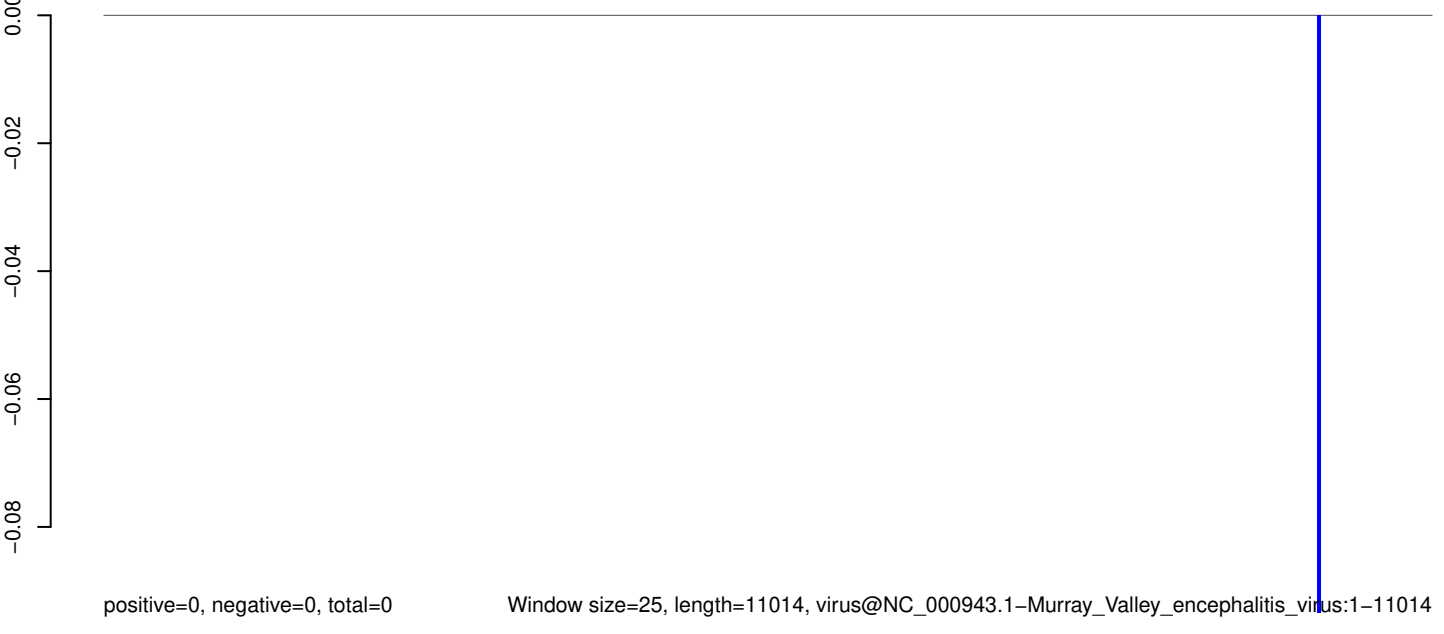
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep



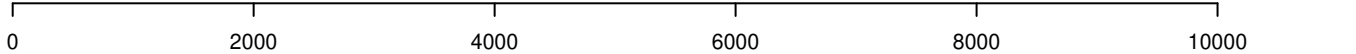
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.rep



positive=0, negative=0, total=0

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



AeAeg_Aag2SINV_dsLuc.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08

positive=0, negative=0, total=0

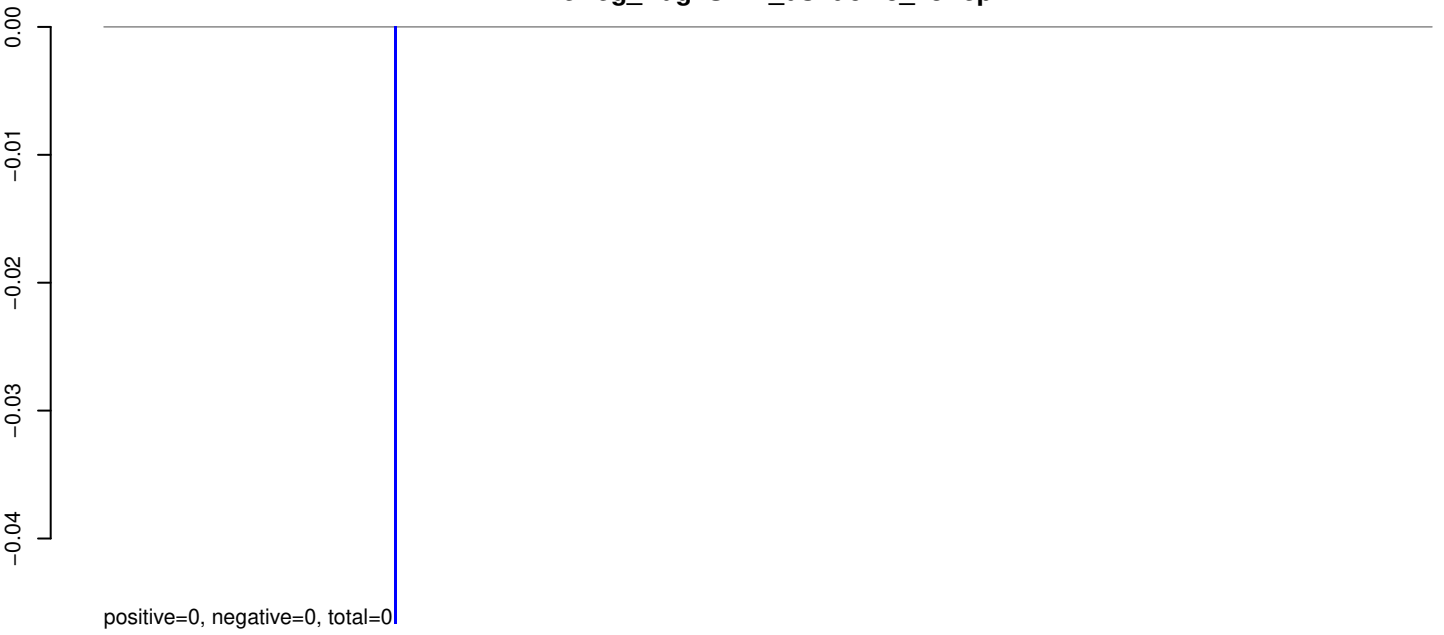
AeAeg_Aag2SINV_dsLuc.rep

0.00
-0.02
-0.04
-0.06
-0.08

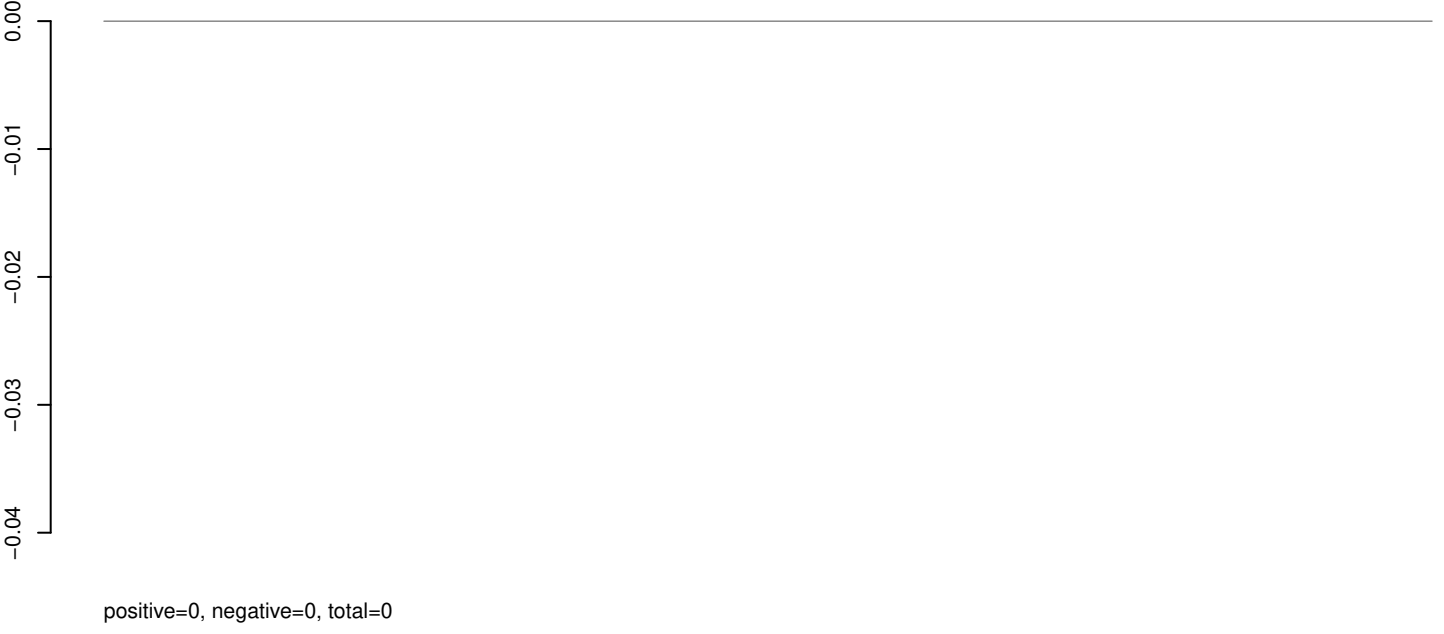
positive=0, negative=0, total=0
Window size=25, length=10749, virus@KX669682.1-Calbertado_virus_isolate_Cal_mos_2_polyprotein_gene:1-10749

0 2000 4000 6000 8000 10000

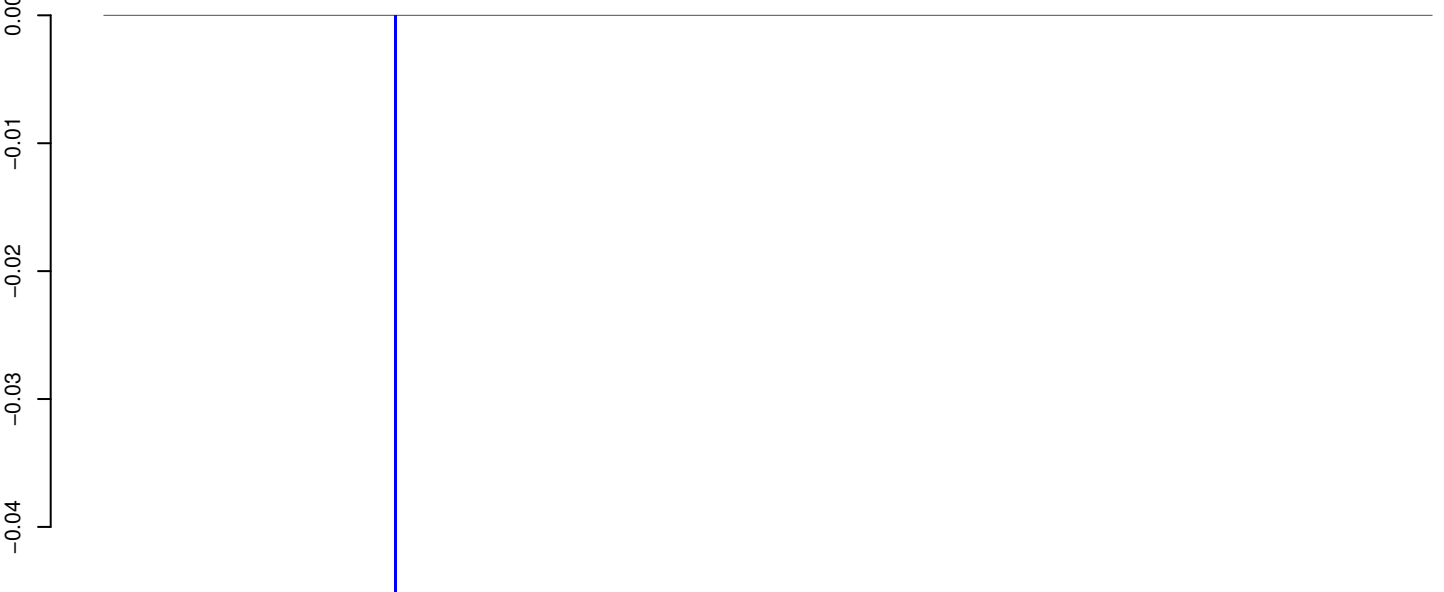
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



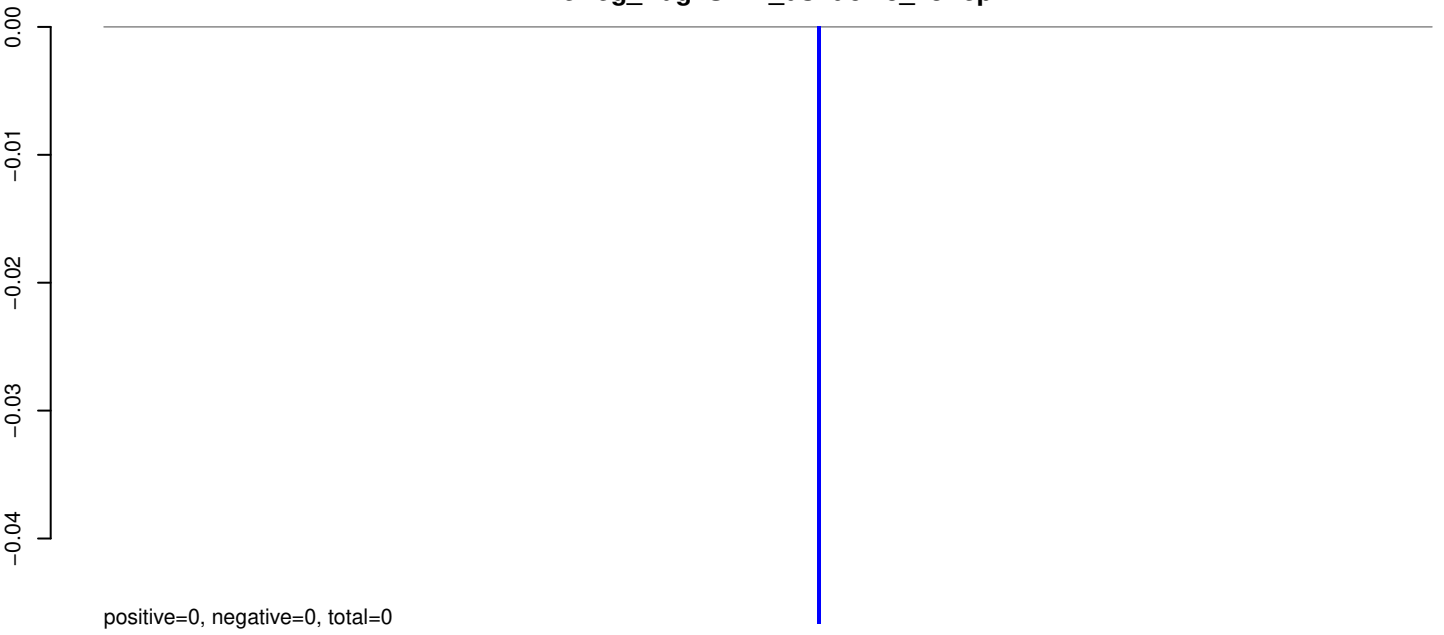
AeAeg_Aag2SINV_dsLuc.rep



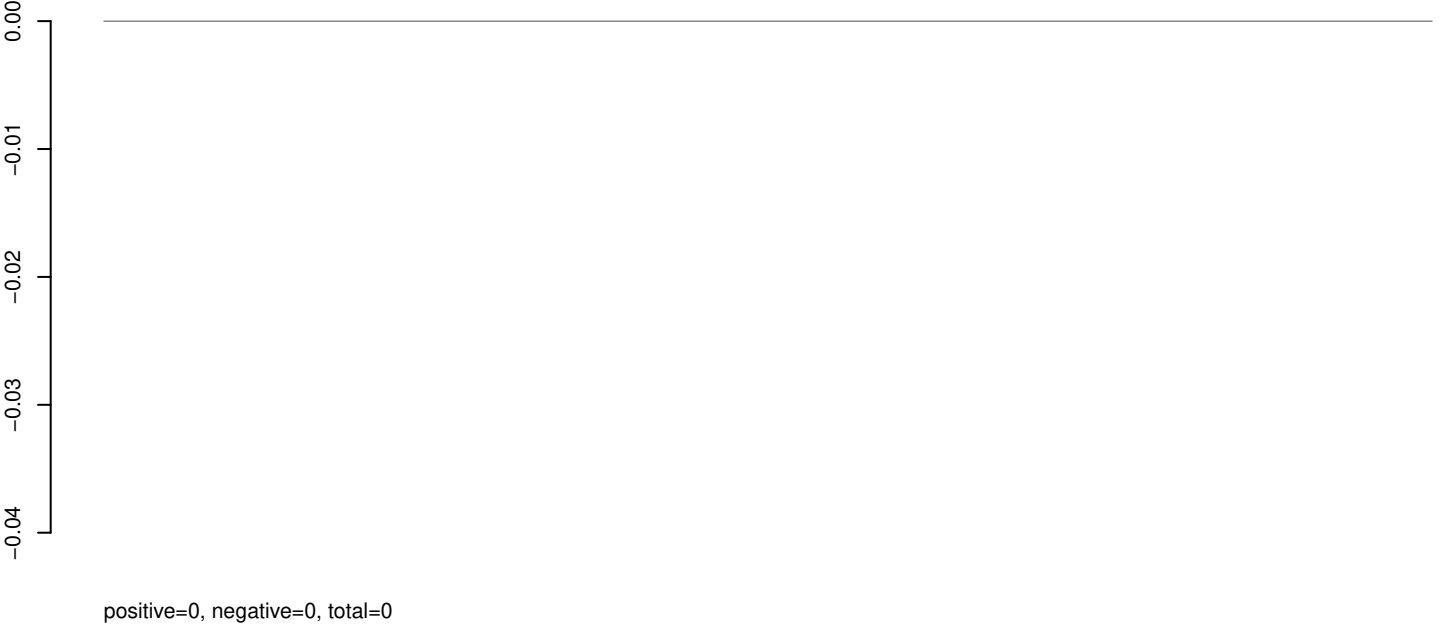
v size=2515, length=1652, total=0, GC=48.553.1-Gambie_virus_putative_nucleoprotein_putative_small_transmembrane_protein_putative_g:1-11652

0 2000 4000 6000 8000 10000 12000

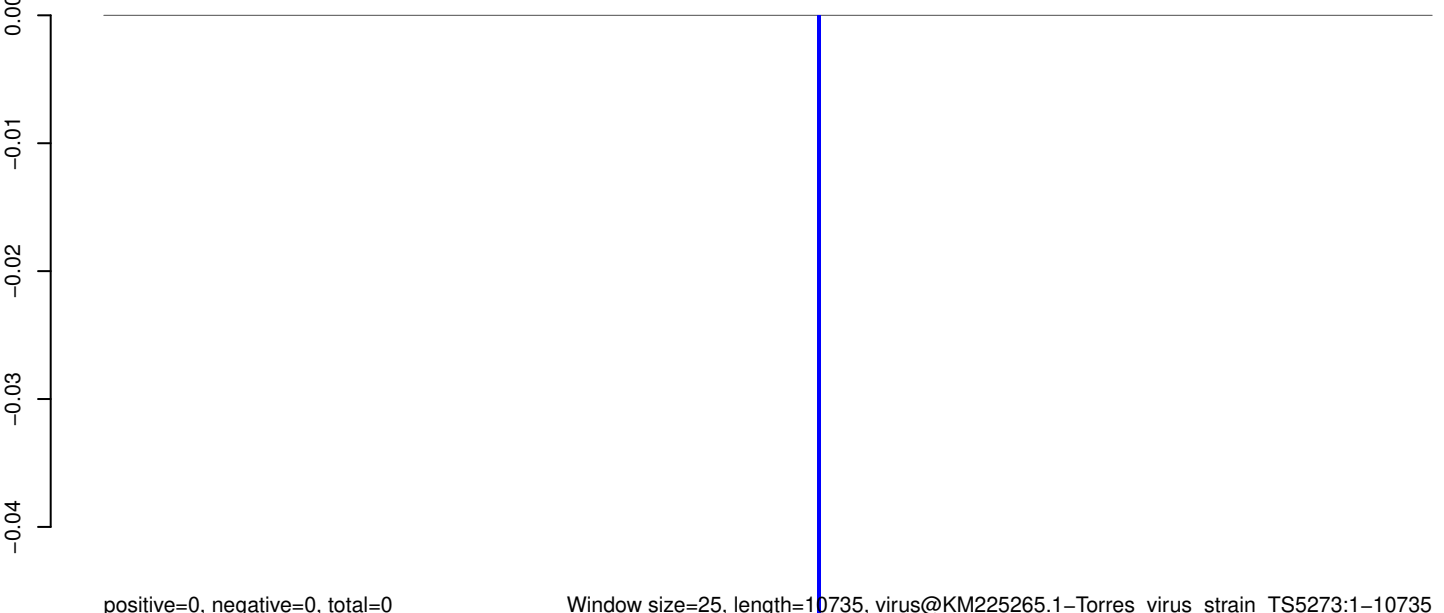
AeAeg_Aag2SINV_dsLuc.18_23.rep



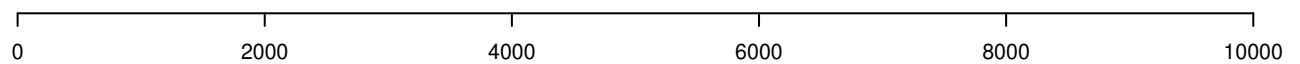
AeAeg_Aag2SINV_dsLuc.24_35.rep



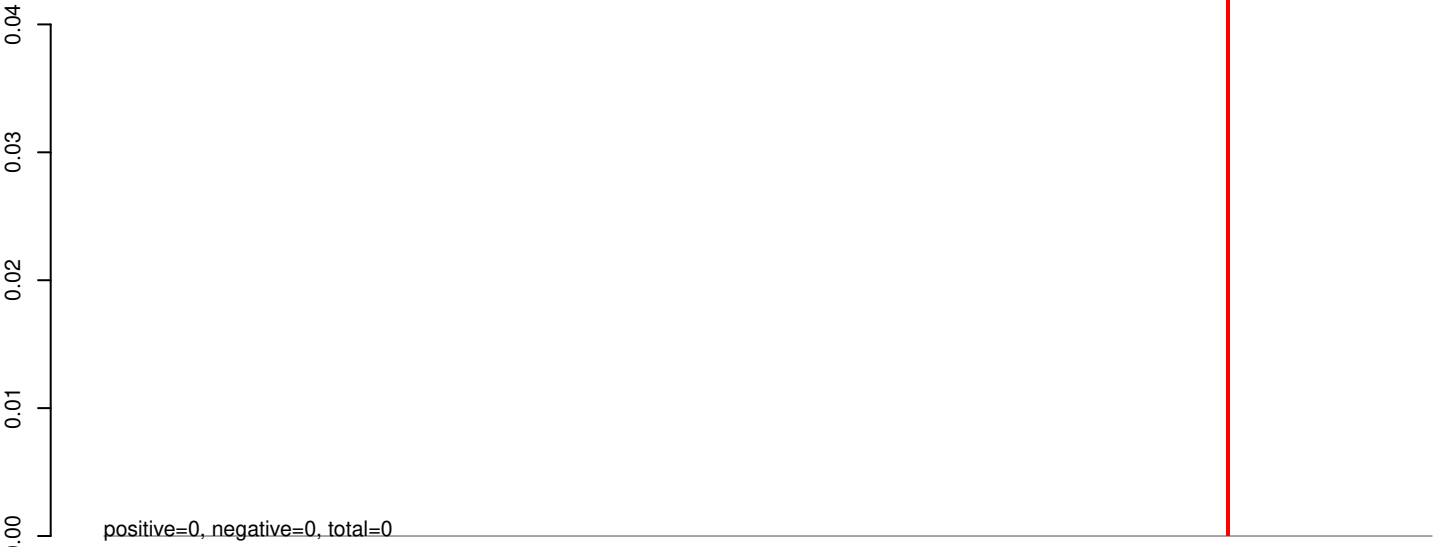
AeAeg_Aag2SINV_dsLuc.rep



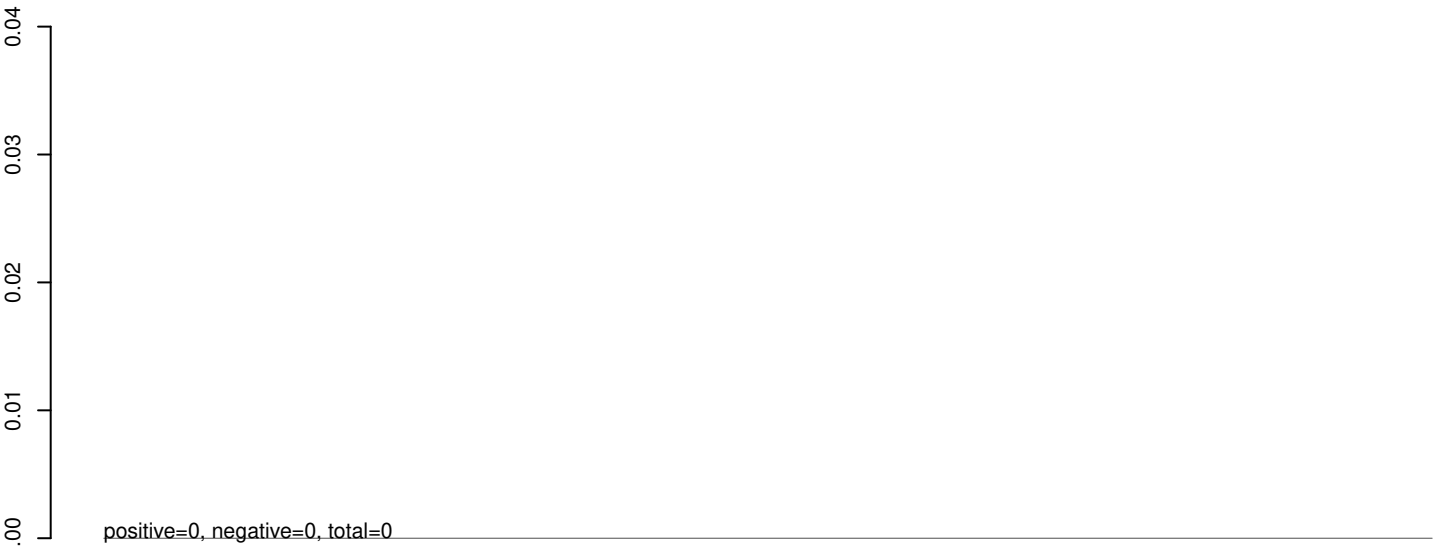
Window size=25, length=10735, virus@KM225265.1-Torres_virus_strain_TS5273:1-10735



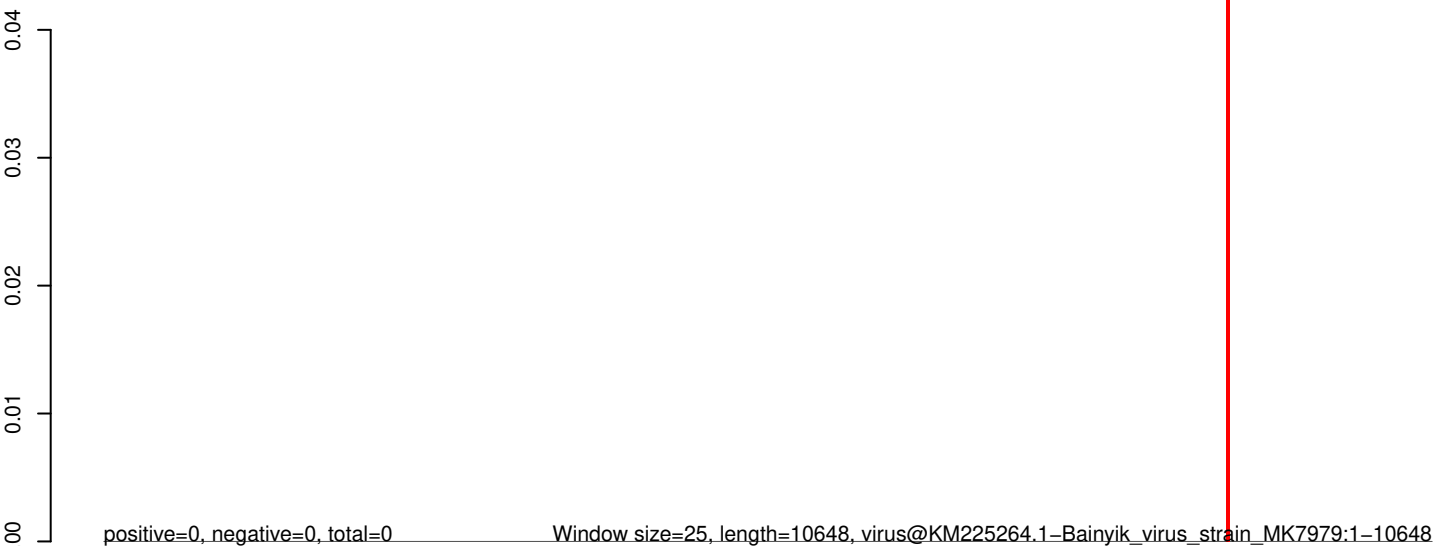
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

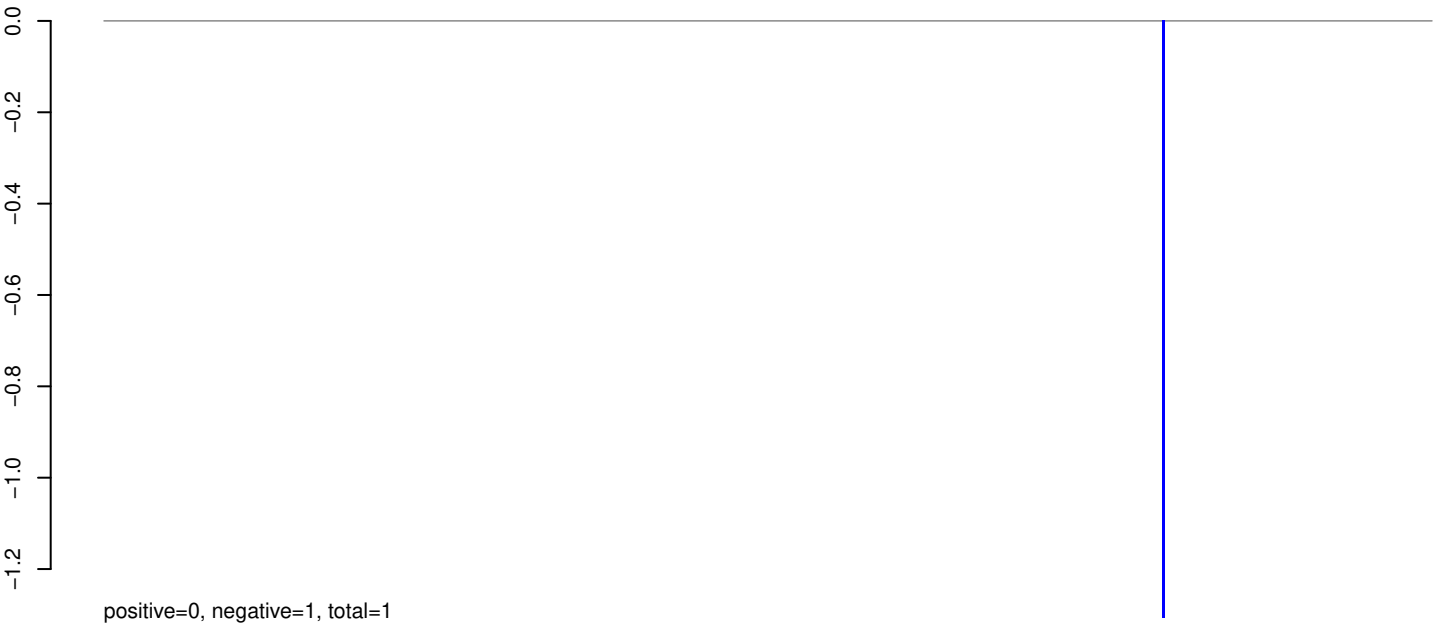


AeAeg_Aag2SINV_dsLuc.rep



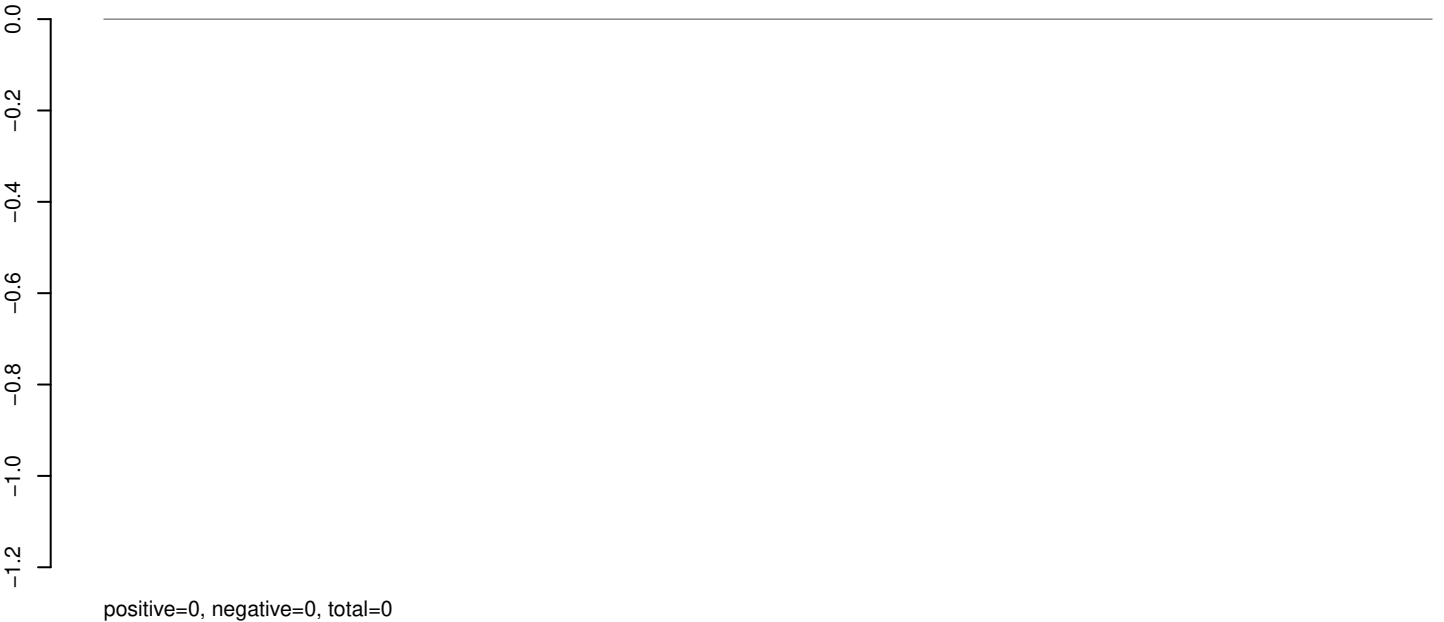
0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_dsLuc.18_23.rep



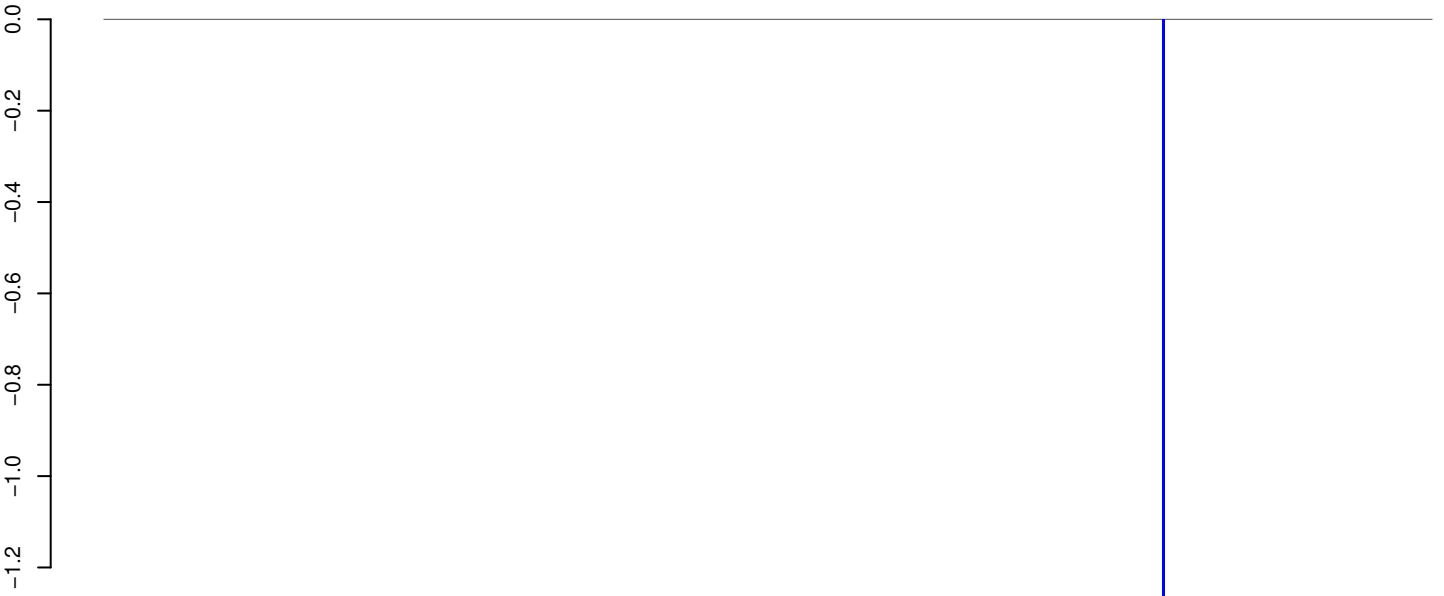
positive=0, negative=1, total=1

AeAeg_Aag2SINV_dsLuc.24_35.rep

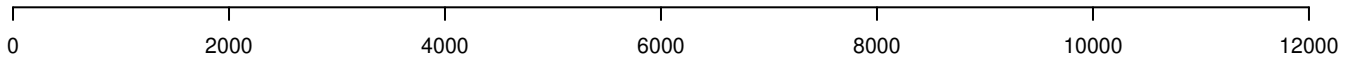


positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.rep



Window size=25, range=1, 2007, Virus@KC608151.1-Ilesha_virus_strain_ILESHA_8e_segment_S_plus_segment_M_plus_segment_L:1-12297



AeAeg_Aag2SINV_dsLuc.18_23.rep

0.04
0.03
0.02
0.01
0.00

positive=0, negative=0, total=0



AeAeg_Aag2SINV_dsLuc.24_35.rep

0.04
0.03
0.02
0.01
0.00

positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.rep

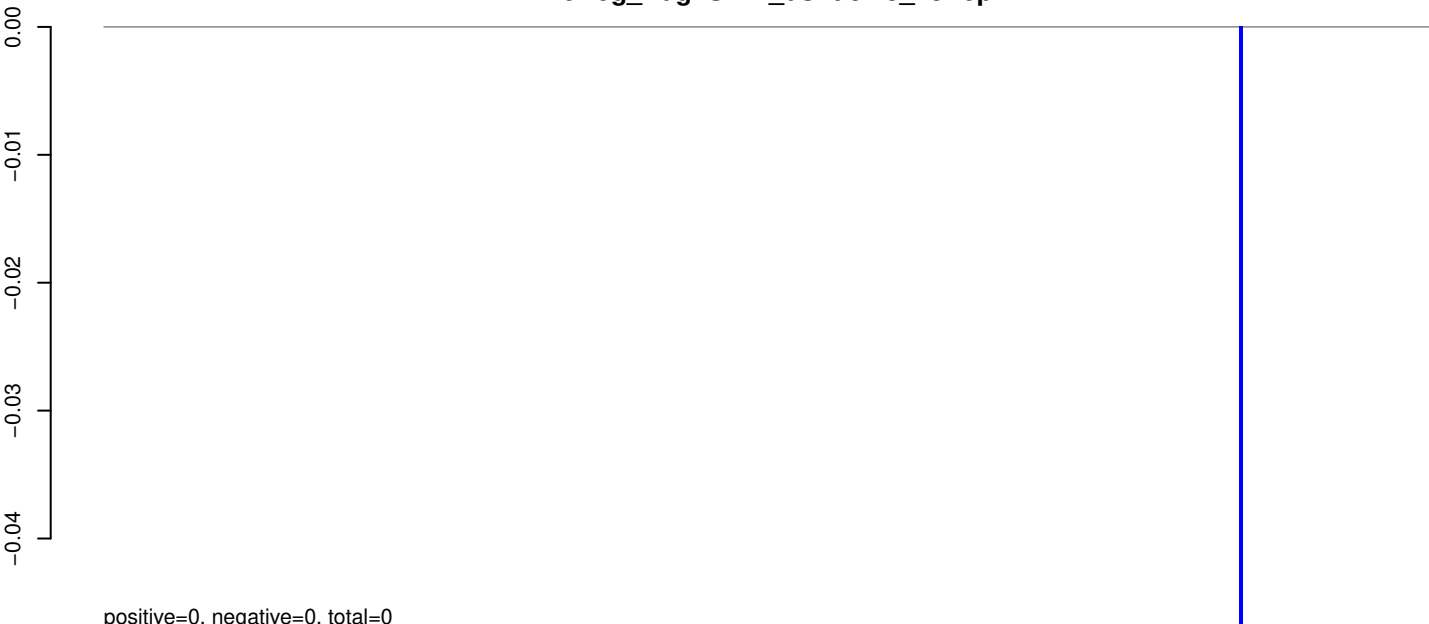
0.04
0.03
0.02
0.01
0.00

size=25 positive=4523 negative=0 total=4523 65.1-Drosophila_melanogaster_American_nodavirus_ANV_SW-2009a_segment_RNA1_and_RNA2:1-4523

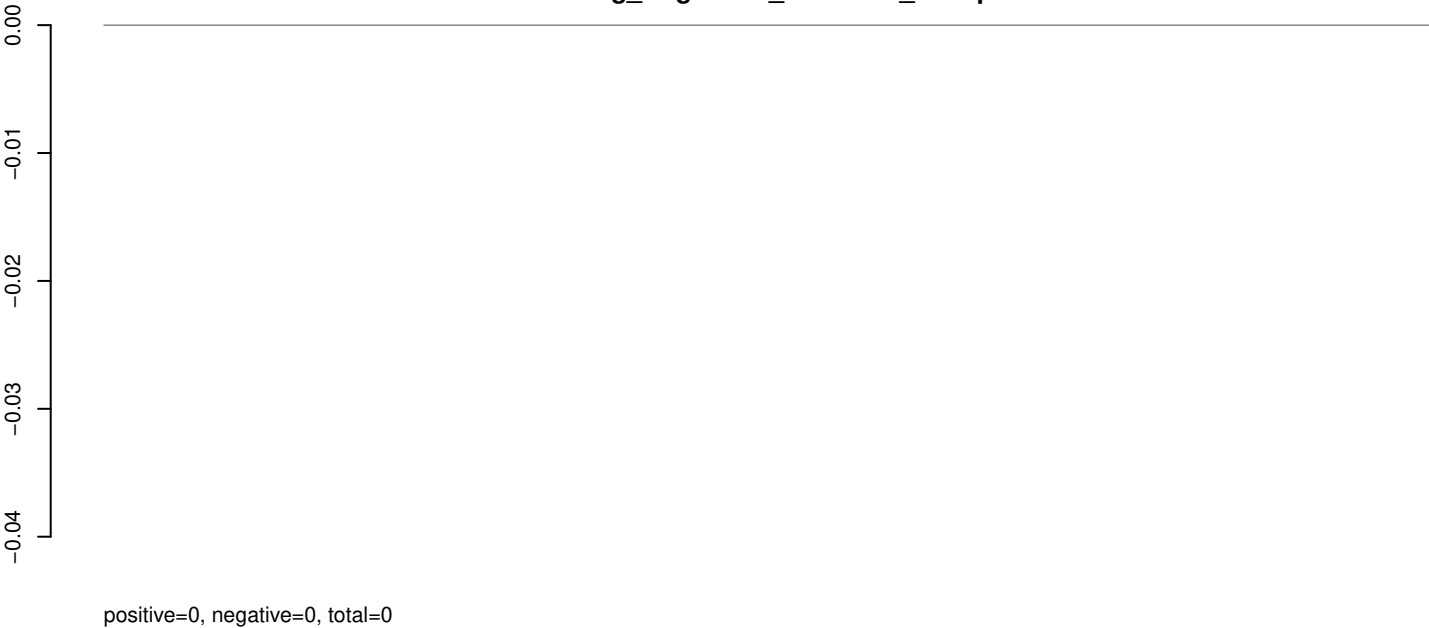


0 1000 2000 3000 4000

AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



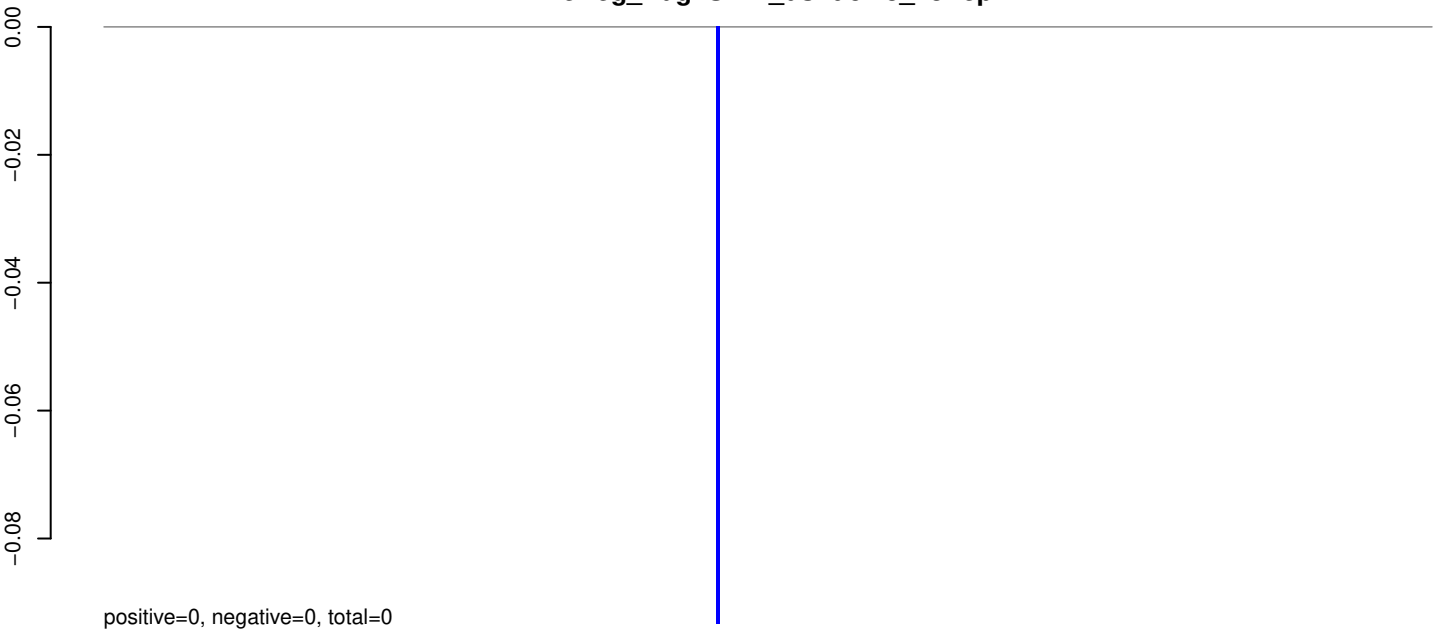
AeAeg_Aag2SINV_dsLuc.rep



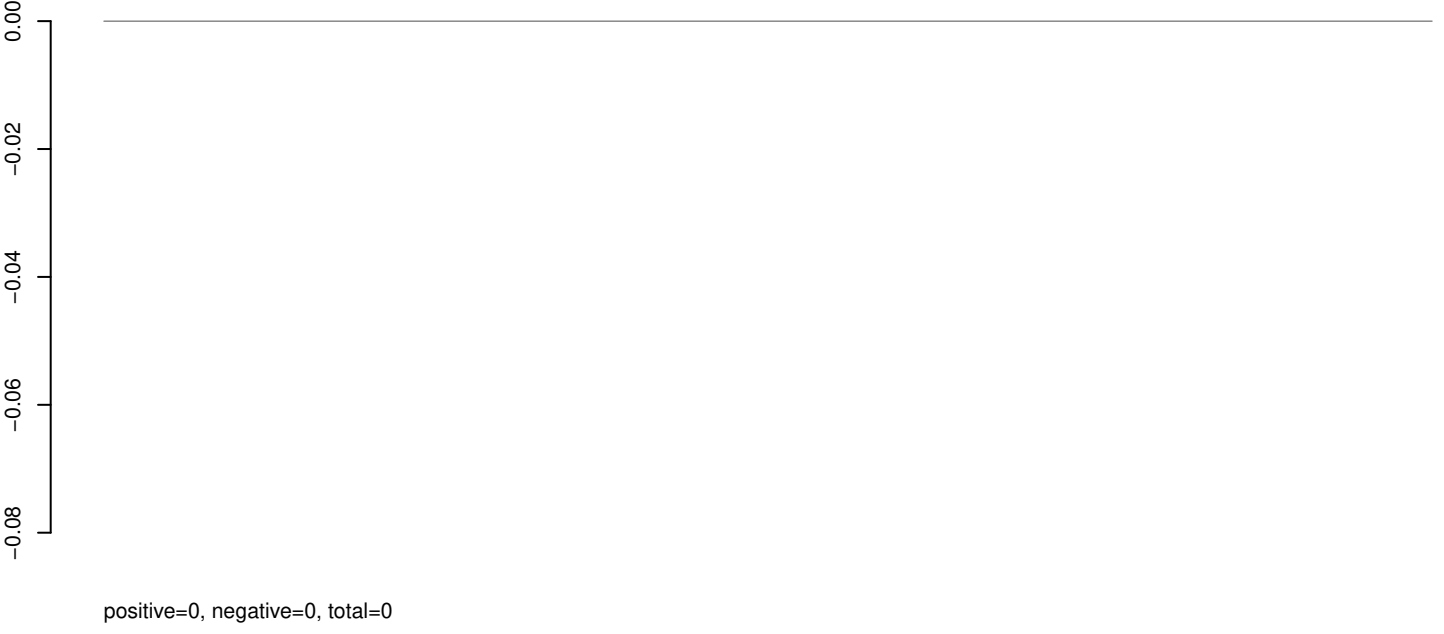
Window size=25, length=10837, virus@FJ663034.1-Culex_flavivirus_isolate_lowa07:1-10837

0 2000 4000 6000 8000 10000

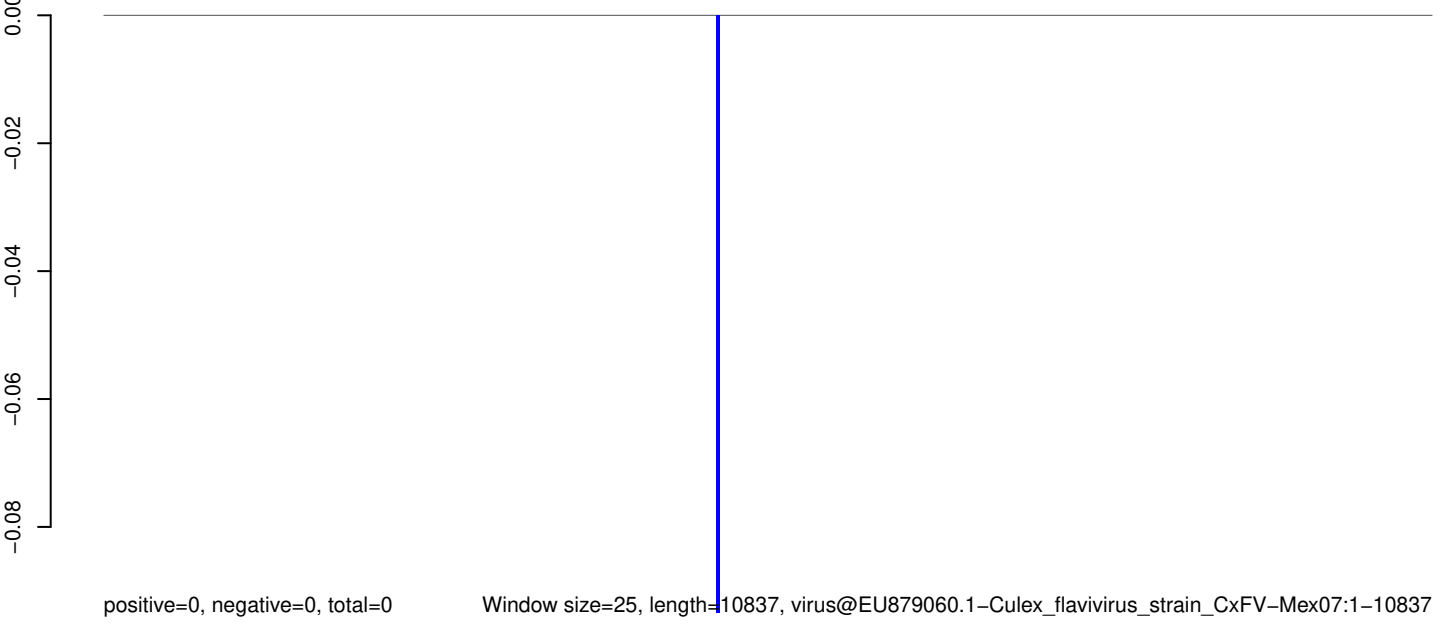
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



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

0 2000 4000 6000 8000 10000

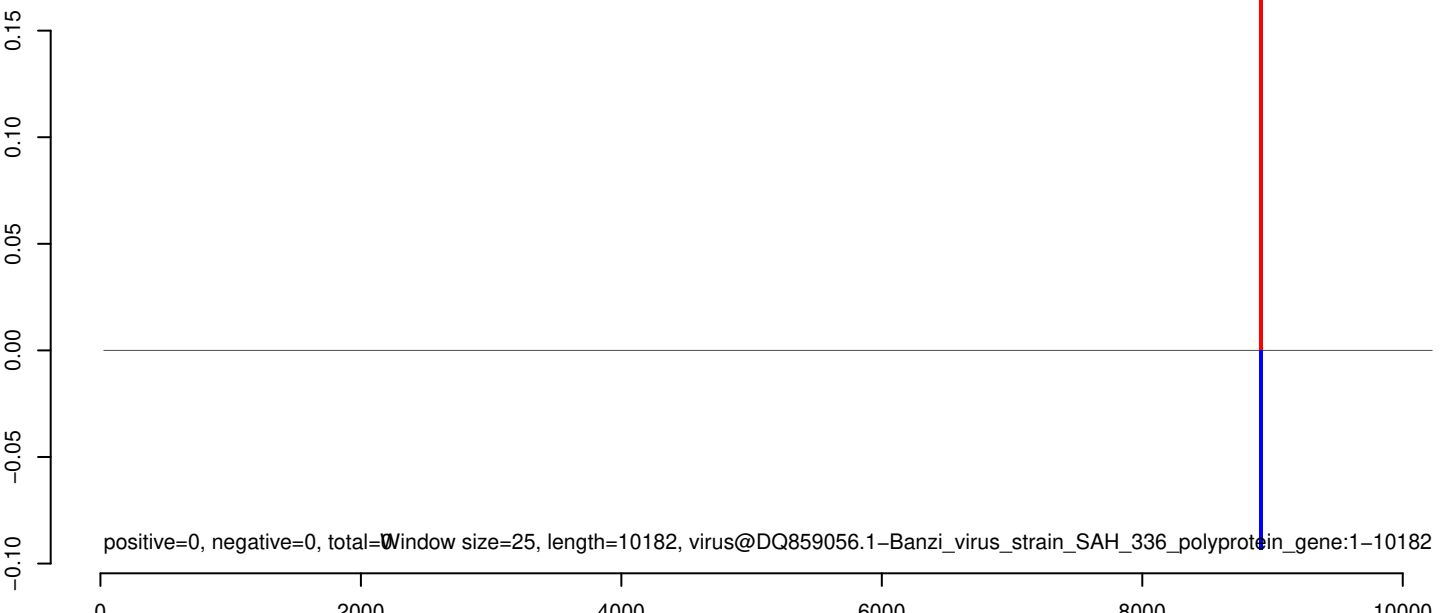
AeAeg_Aag2SINV_dsLuc.18_23.rep



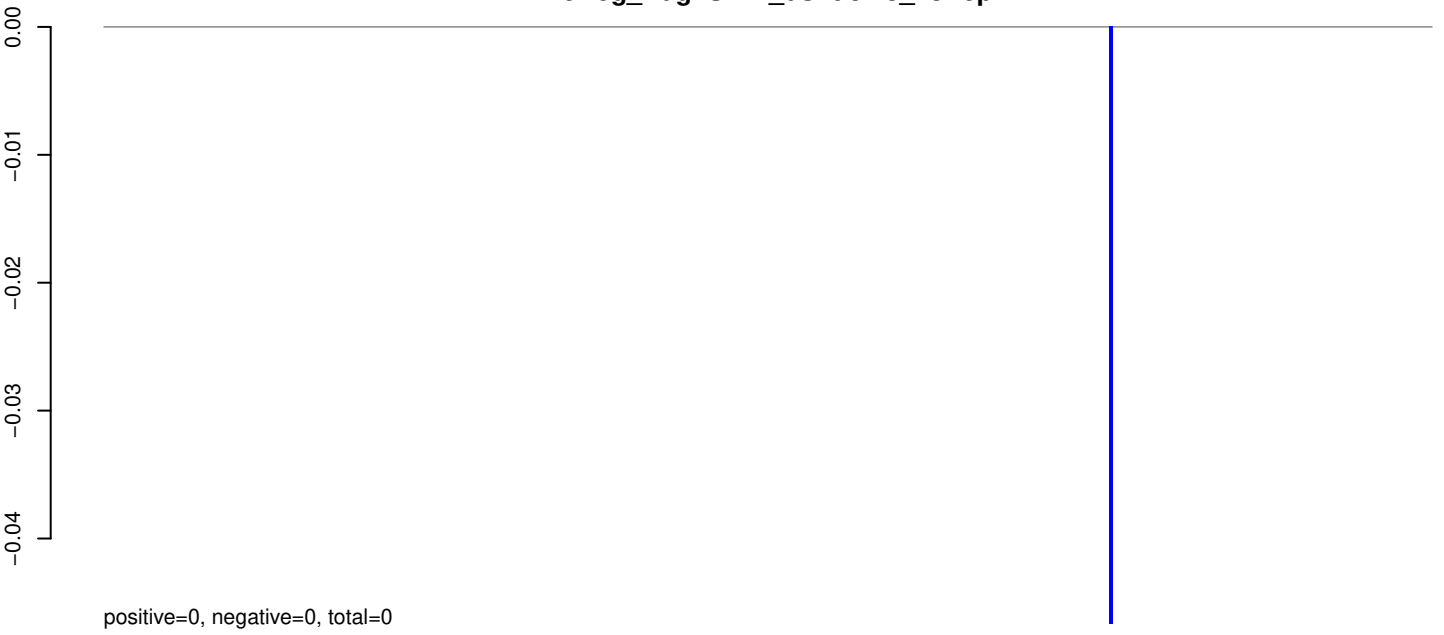
AeAeg_Aag2SINV_dsLuc.24_35.rep



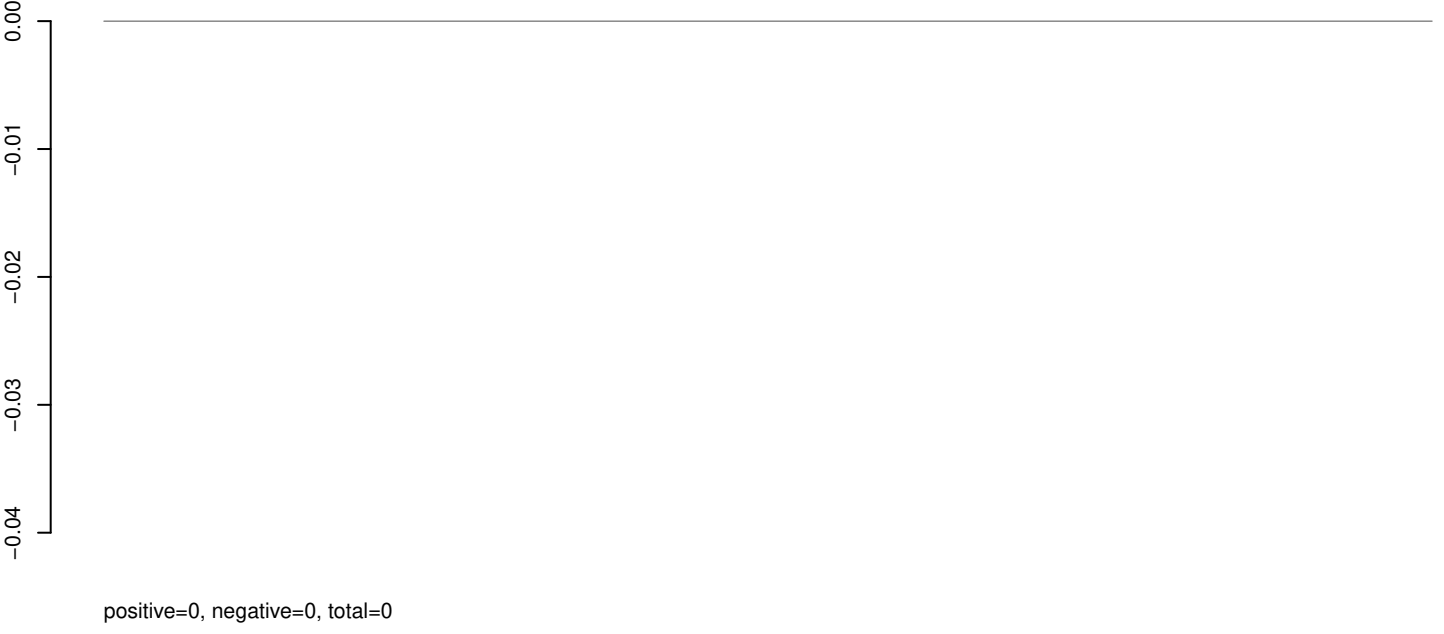
AeAeg_Aag2SINV_dsLuc.rep



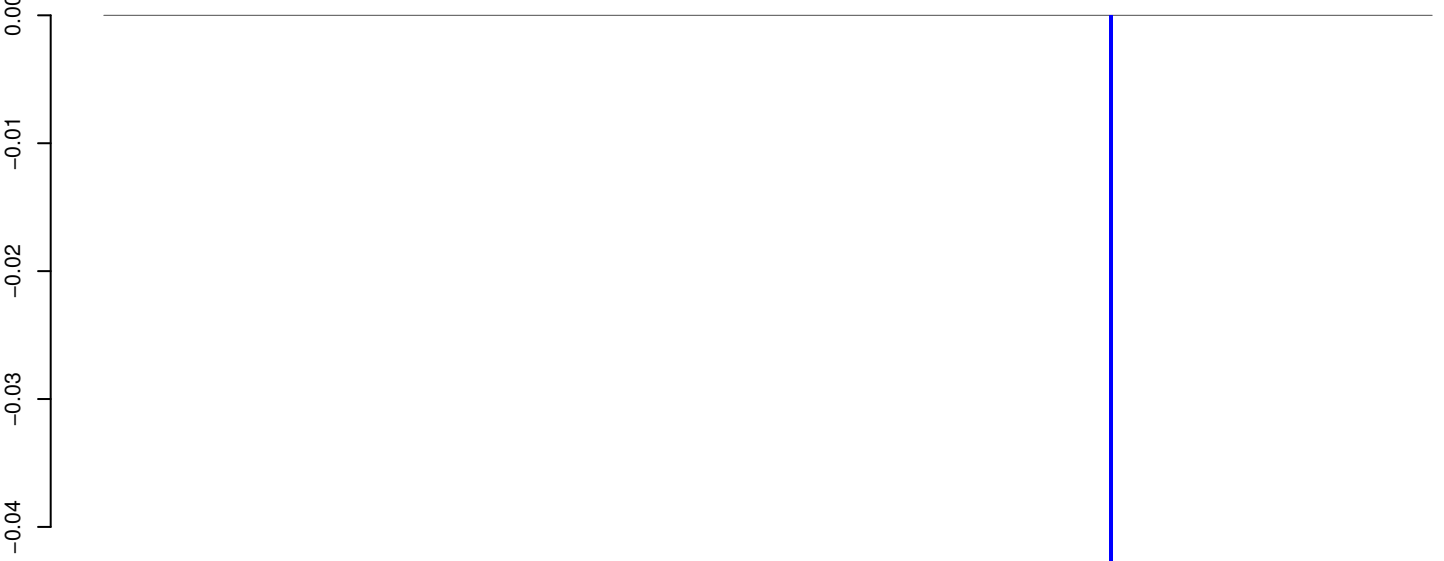
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



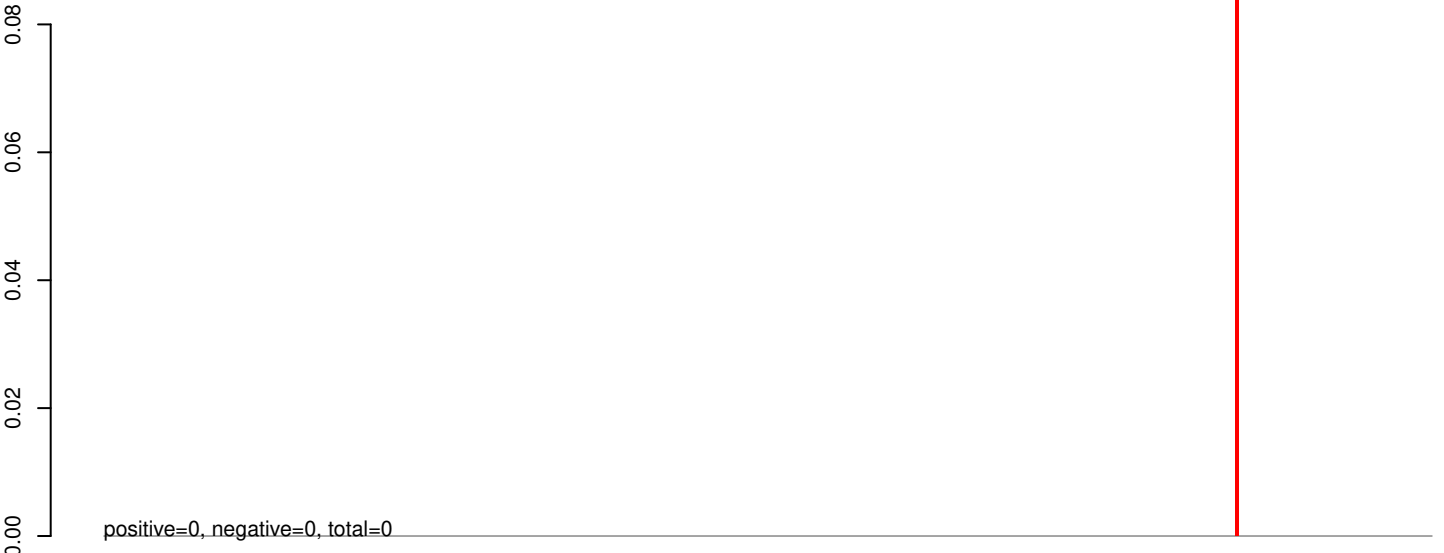
AeAeg_Aag2SINV_dsLuc.rep



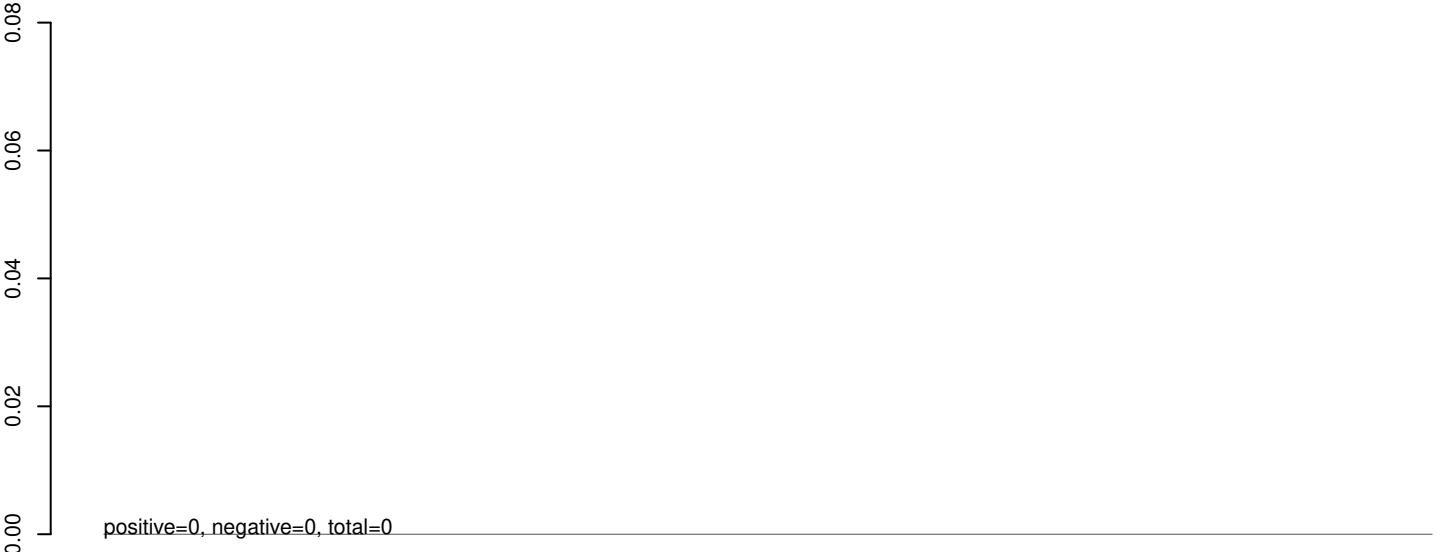
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0 2000 4000 6000 8000 10000

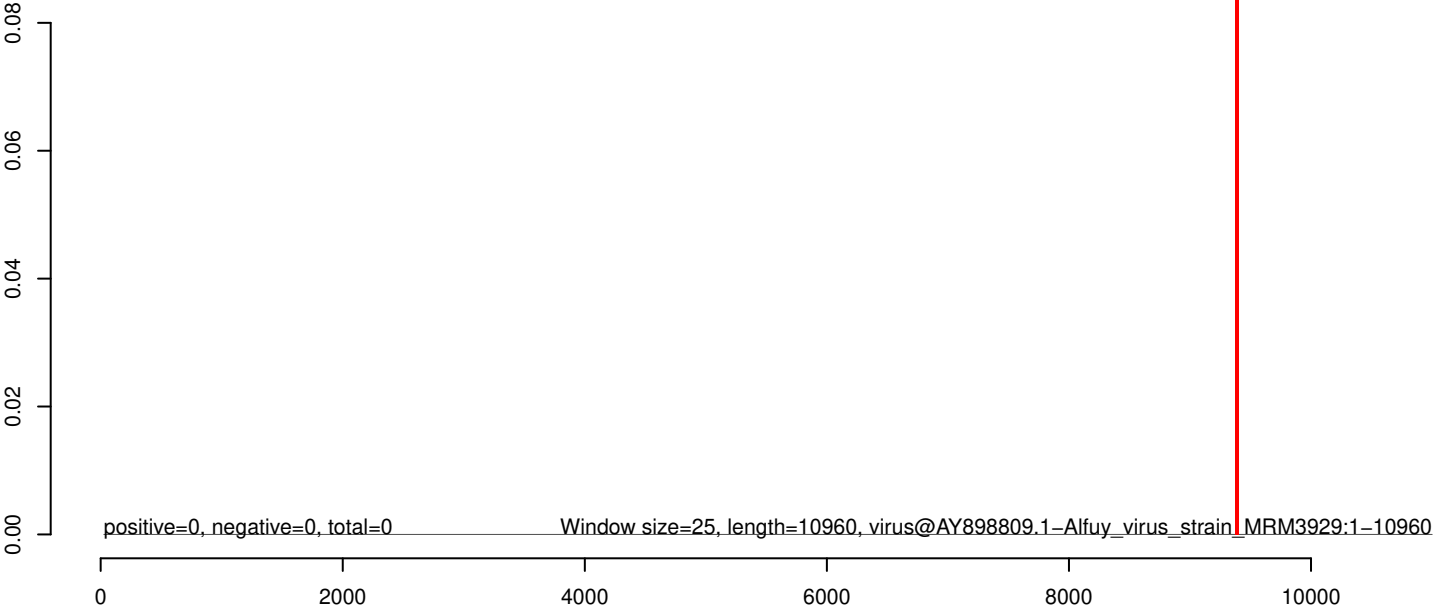
AeAeg_Aag2SINV_dsLuc.18_23.rep



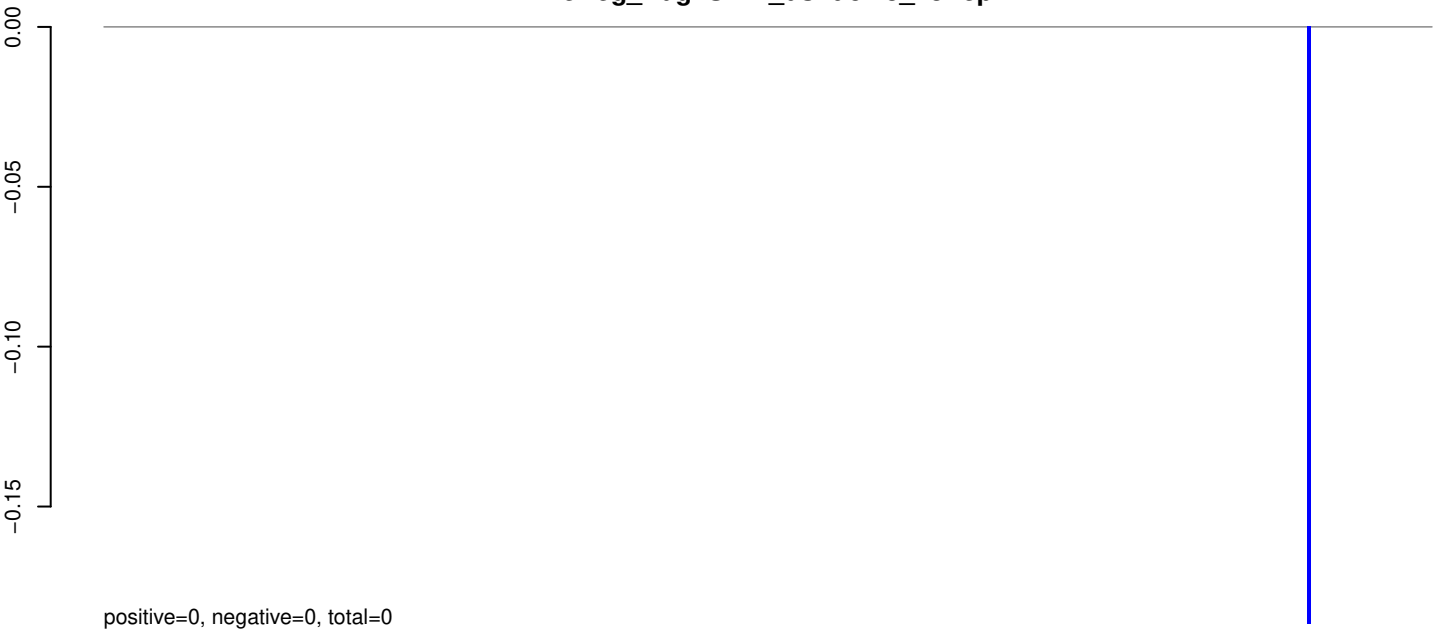
AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep

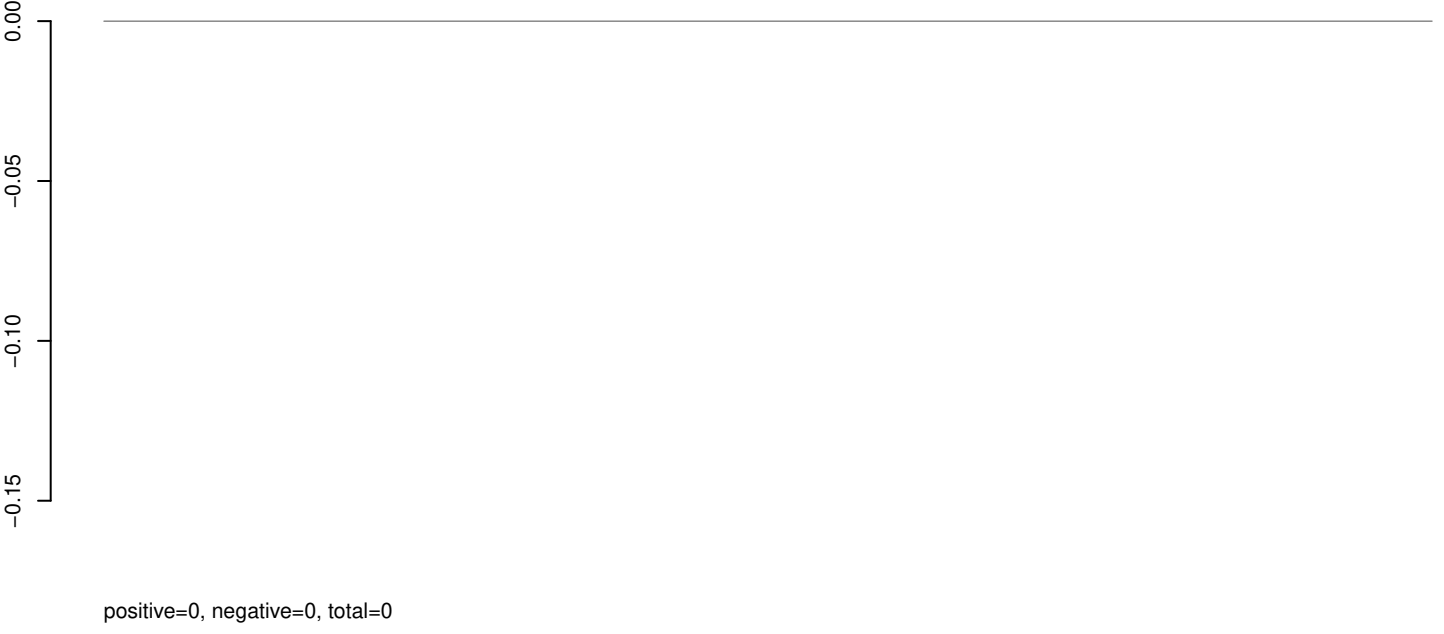


AeAeg_Aag2SINV_dsLuc.18_23.rep



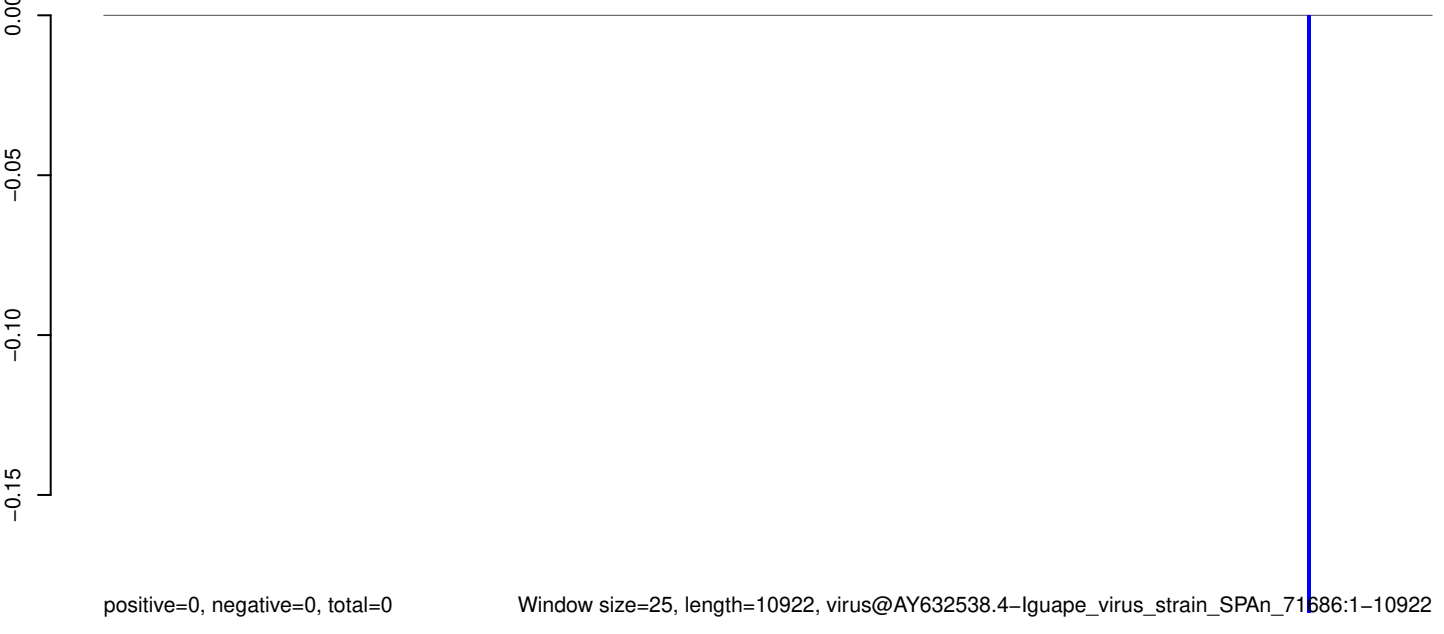
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.rep

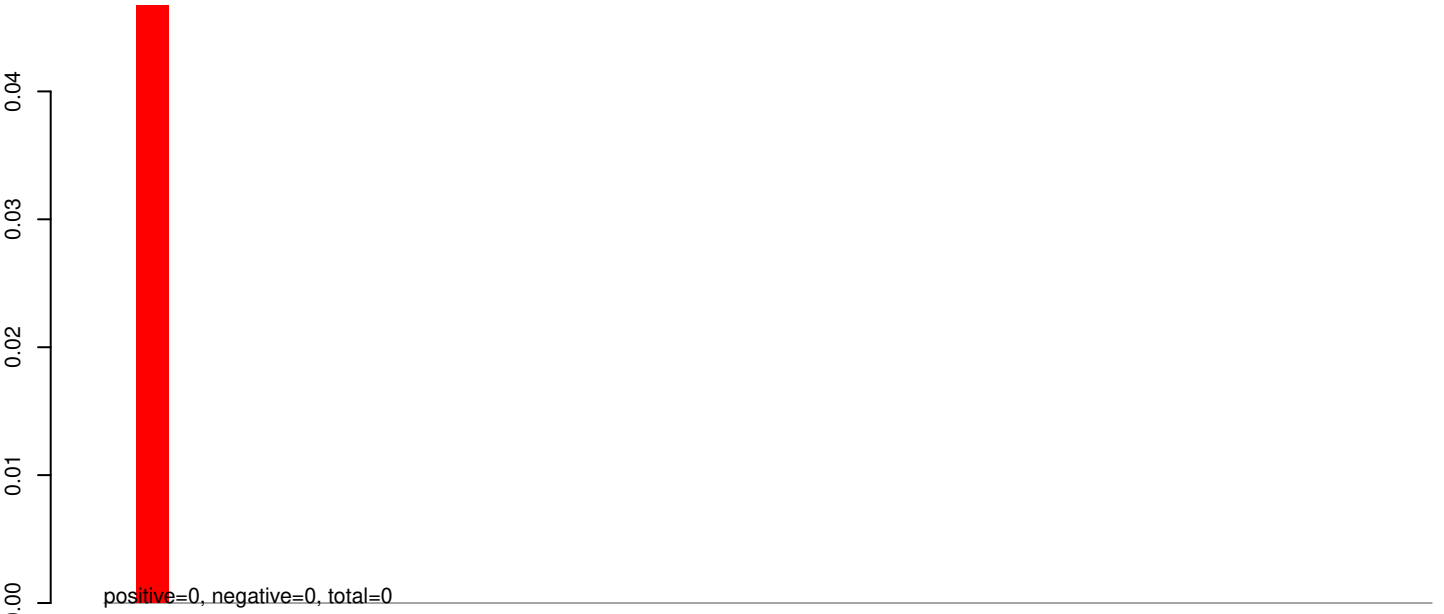


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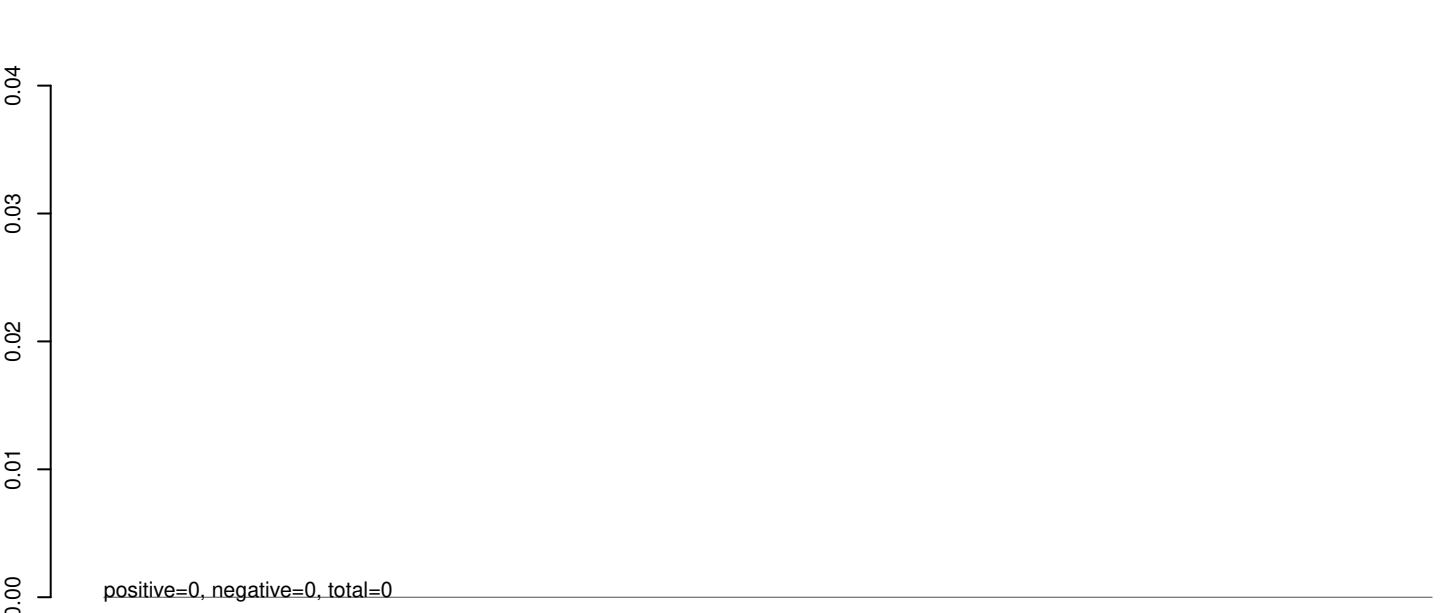
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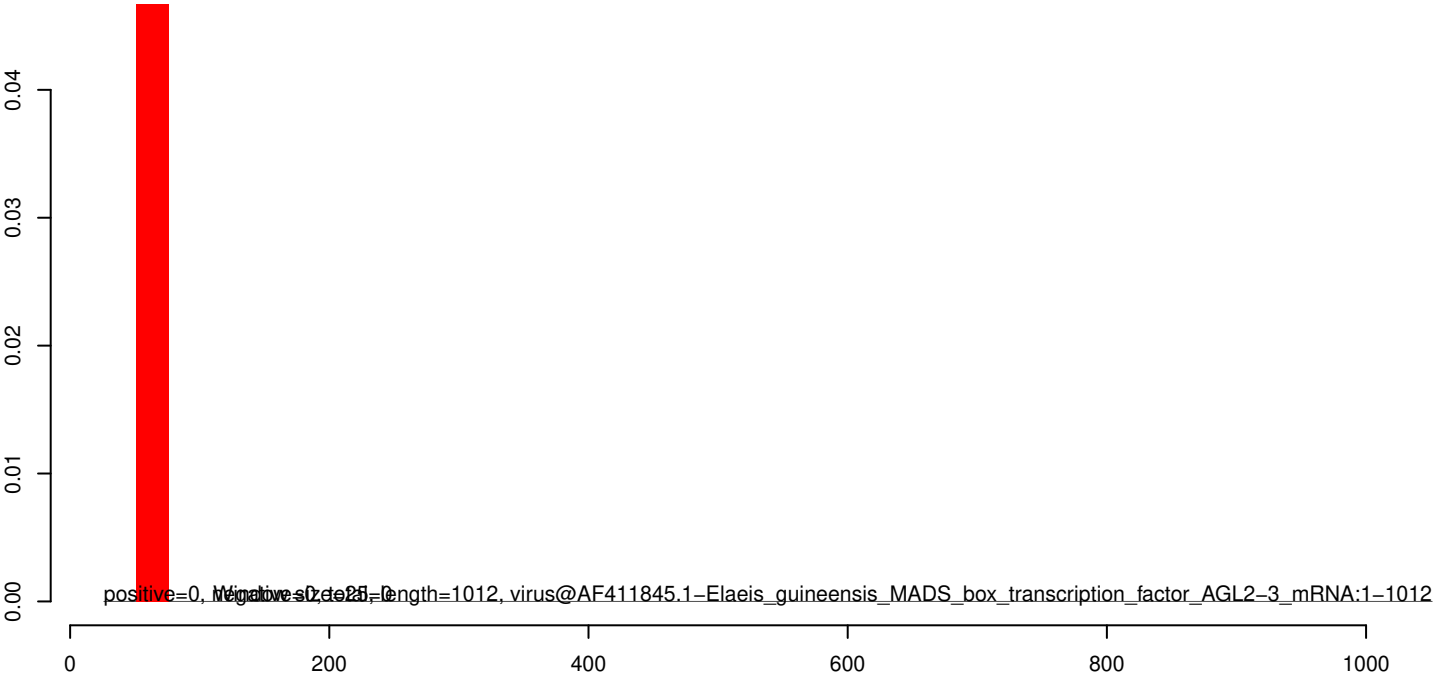
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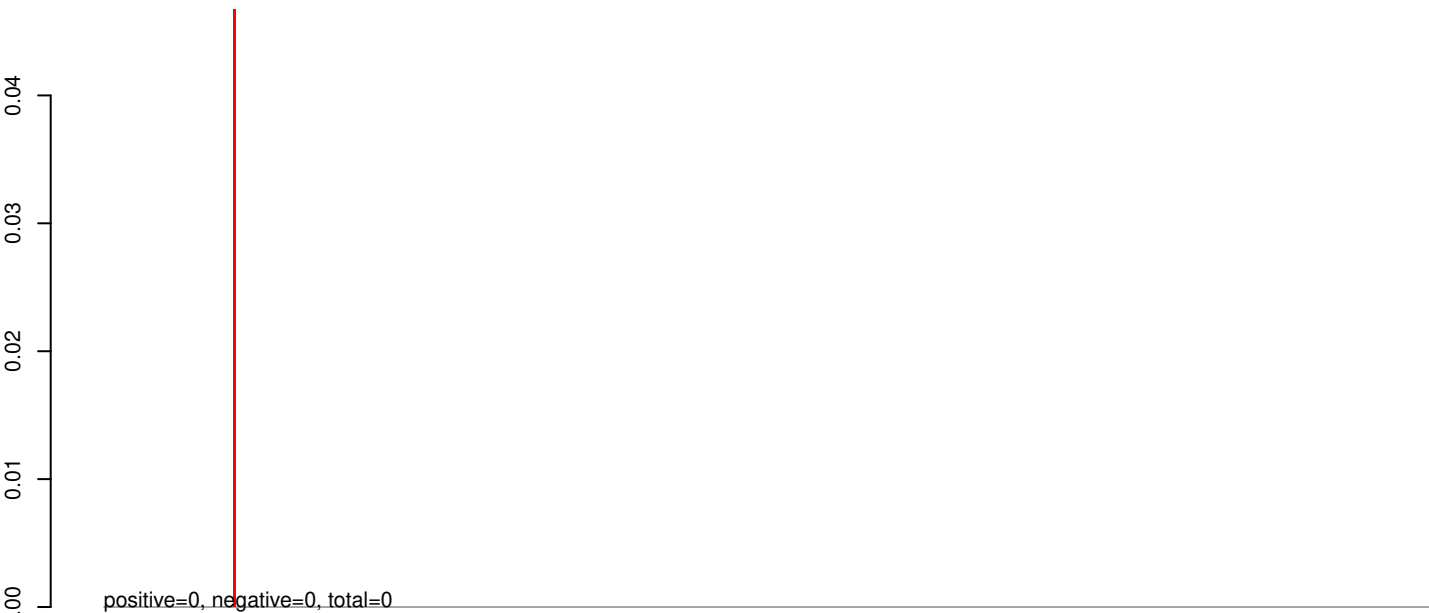
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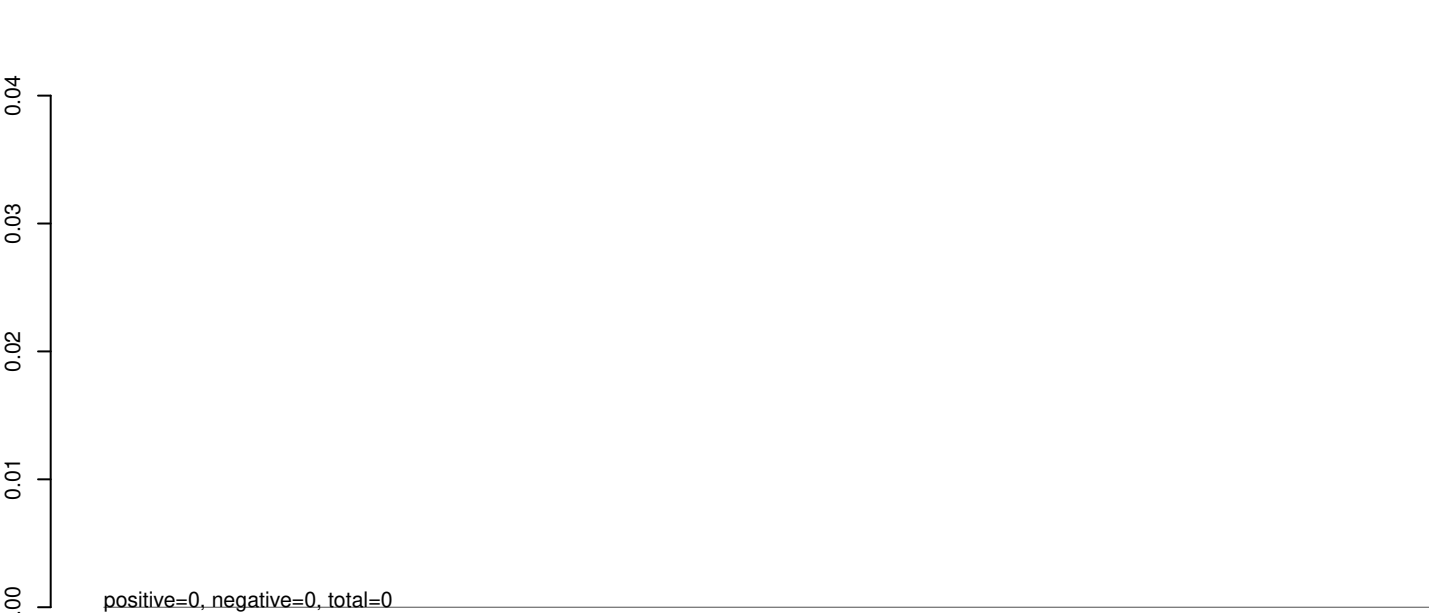
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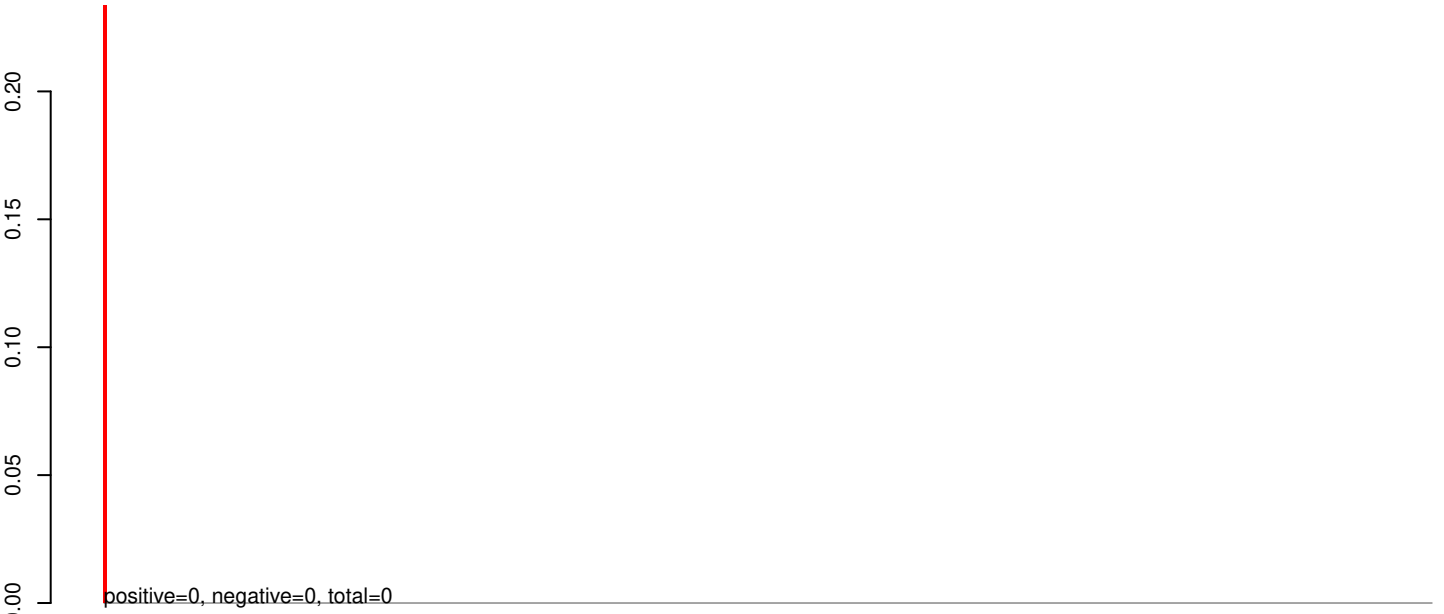
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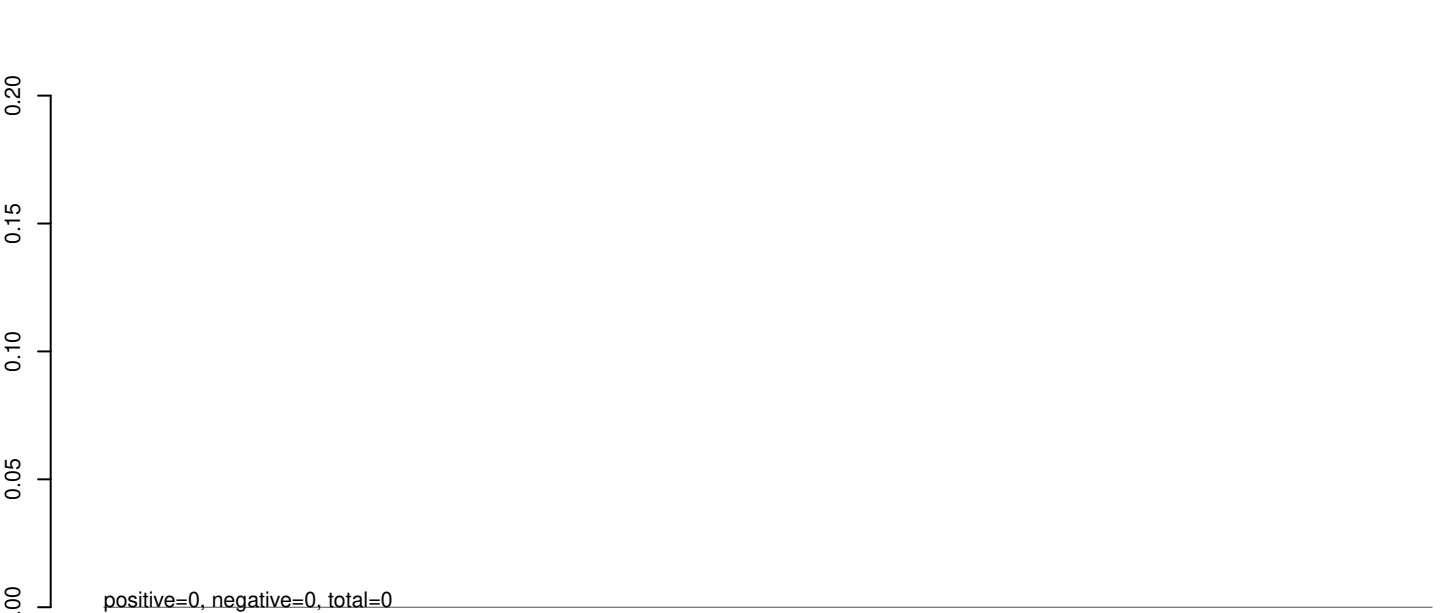
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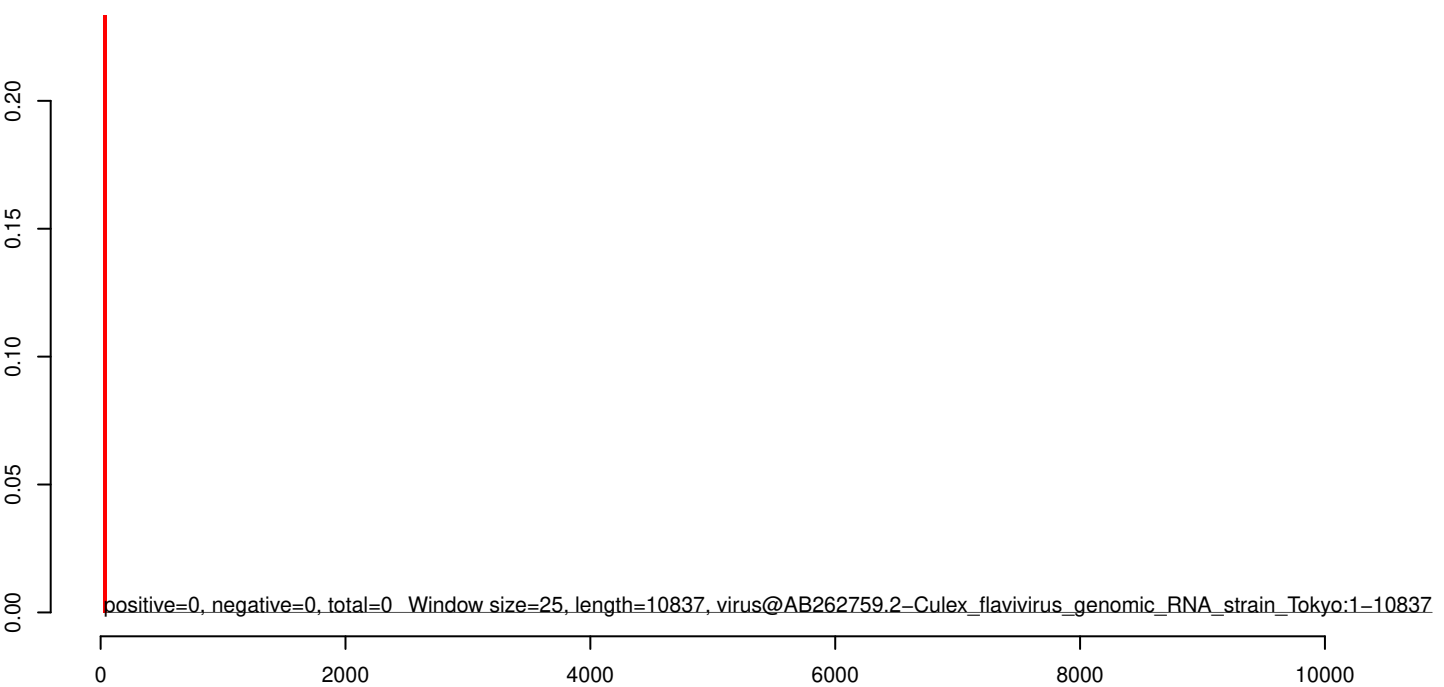
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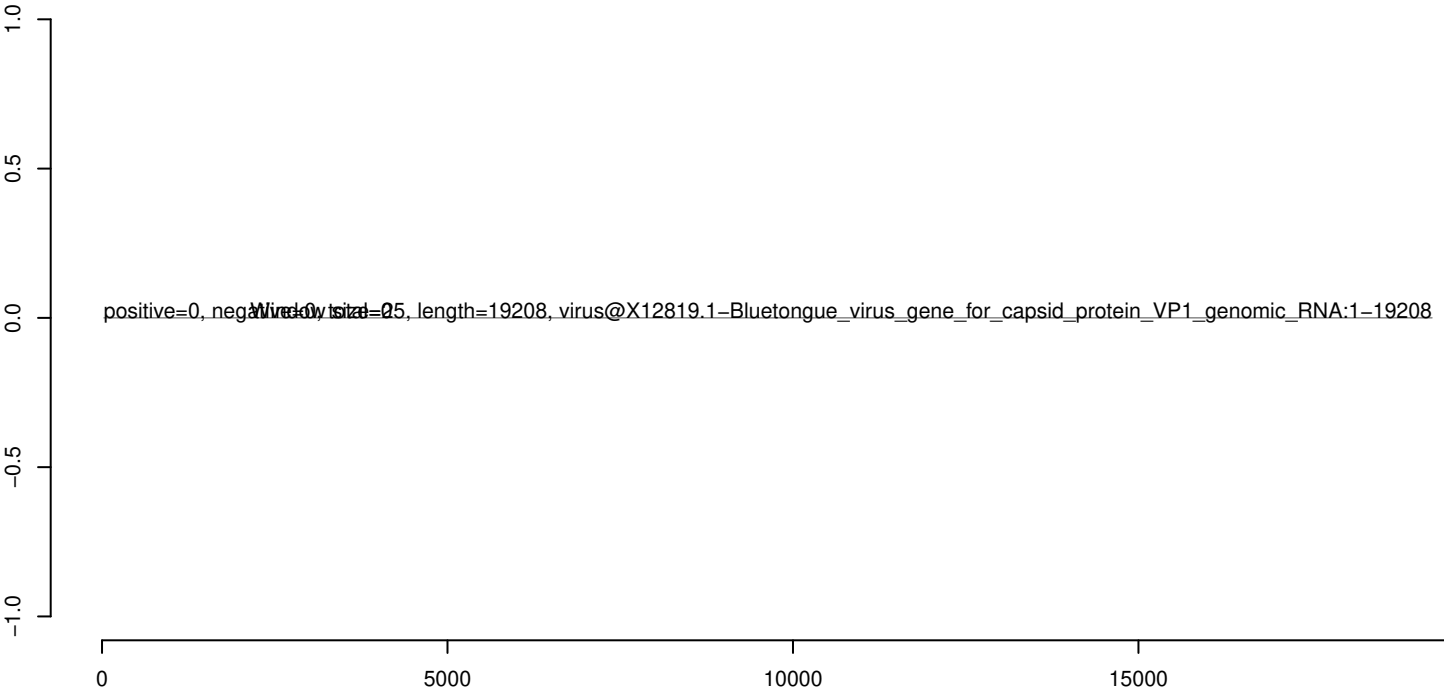
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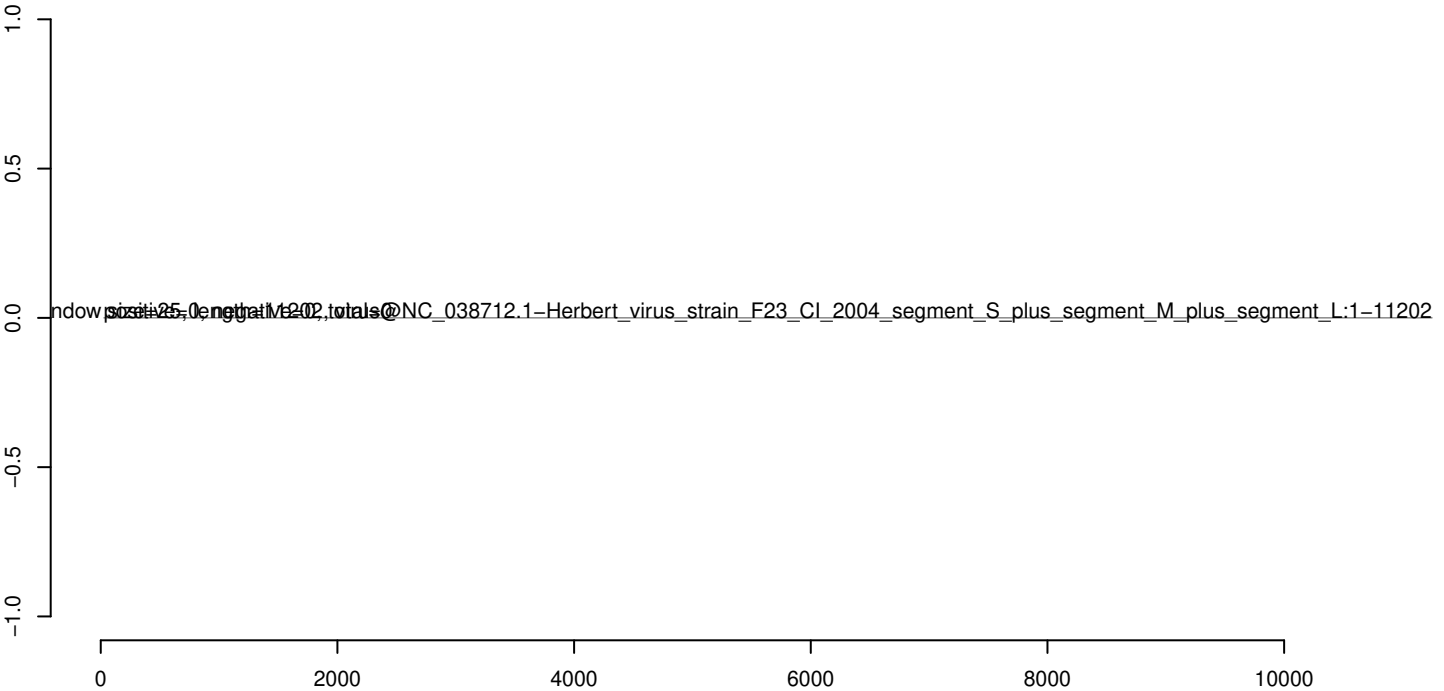
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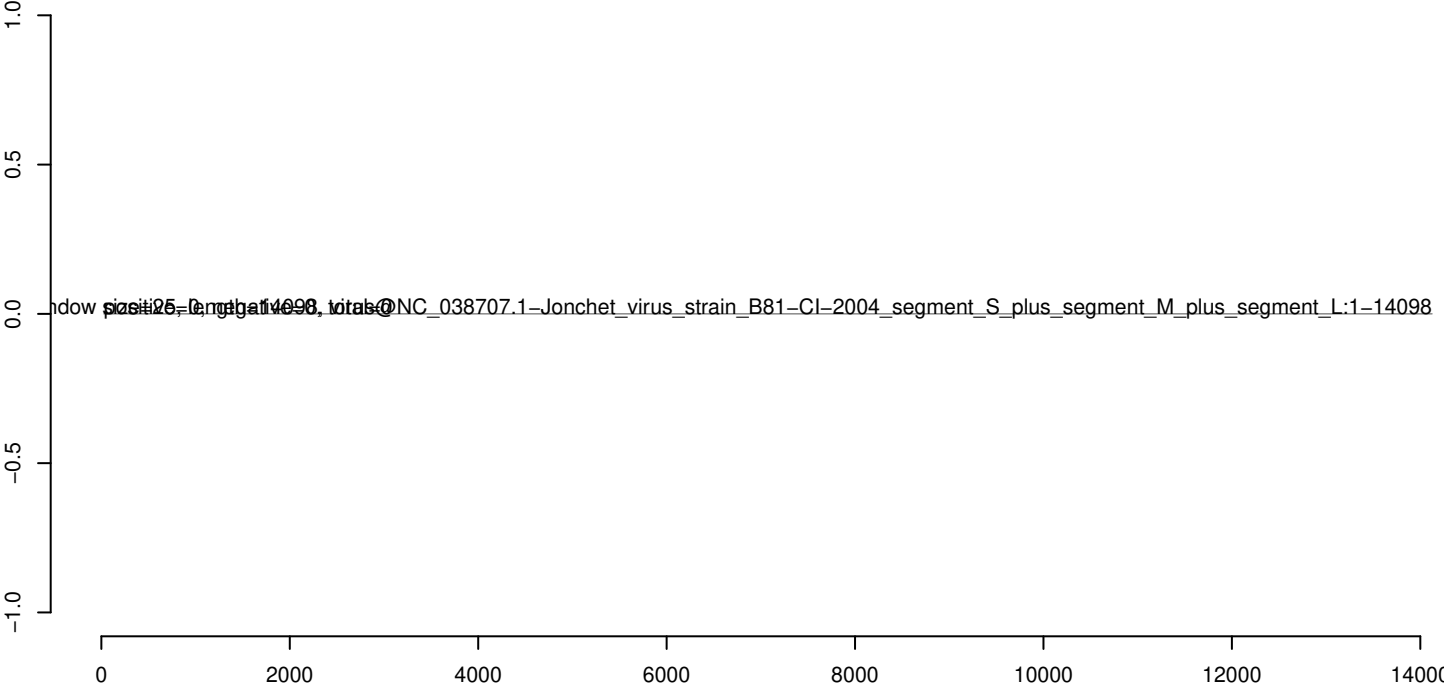
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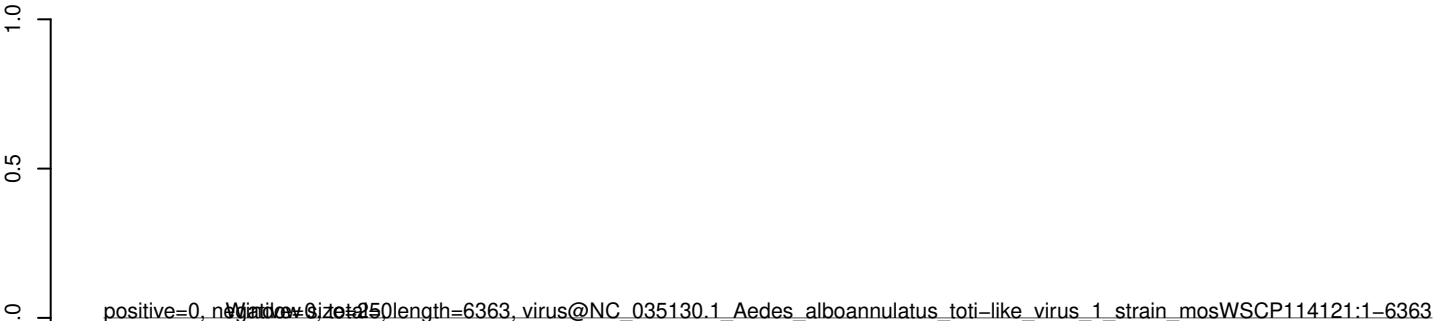
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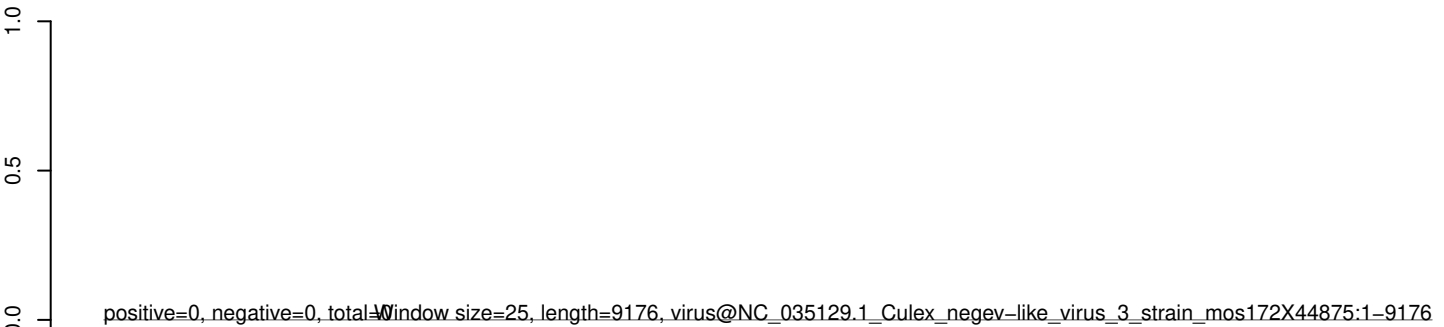
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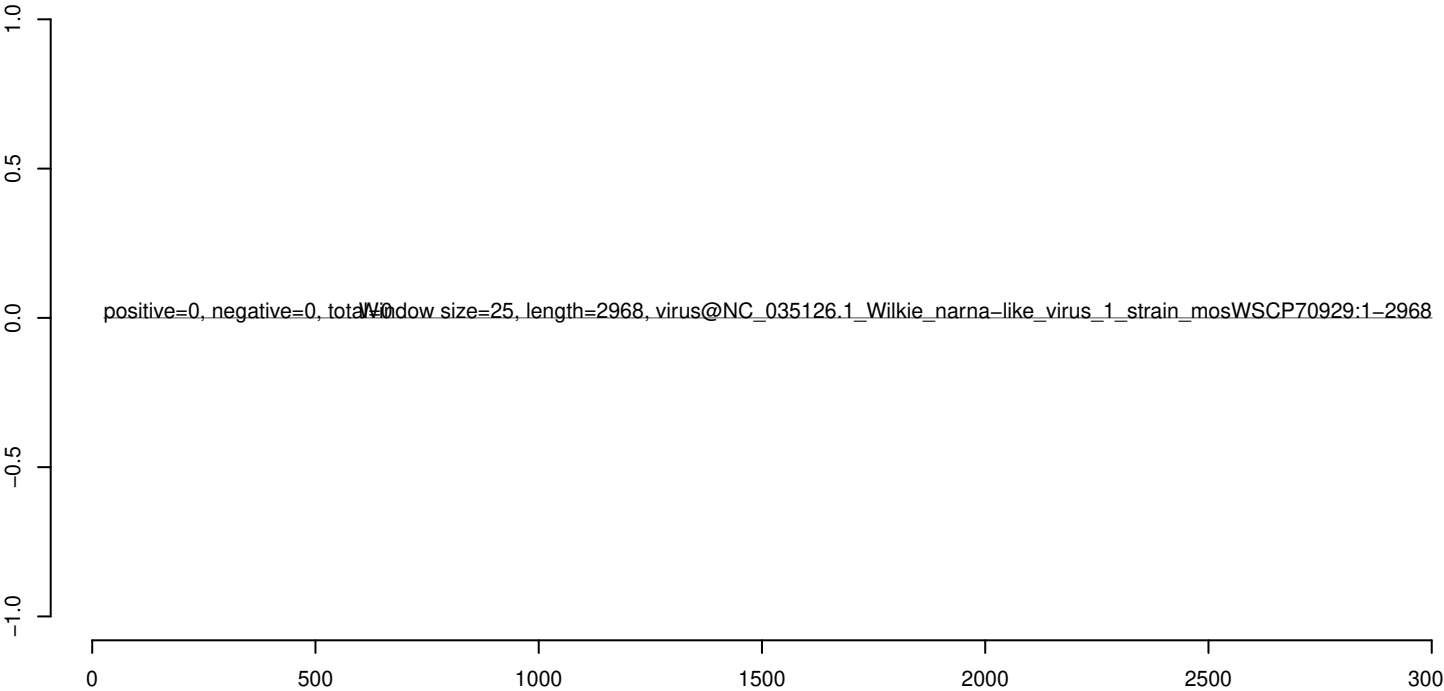
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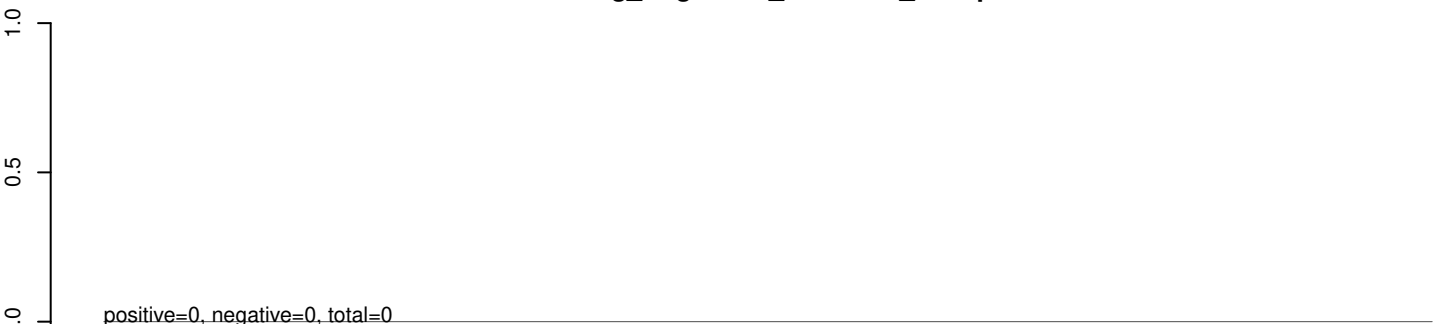
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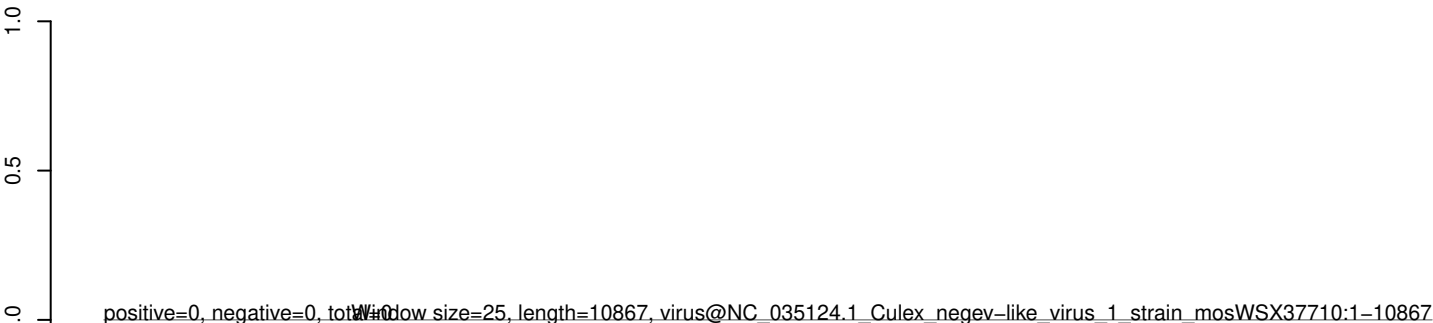
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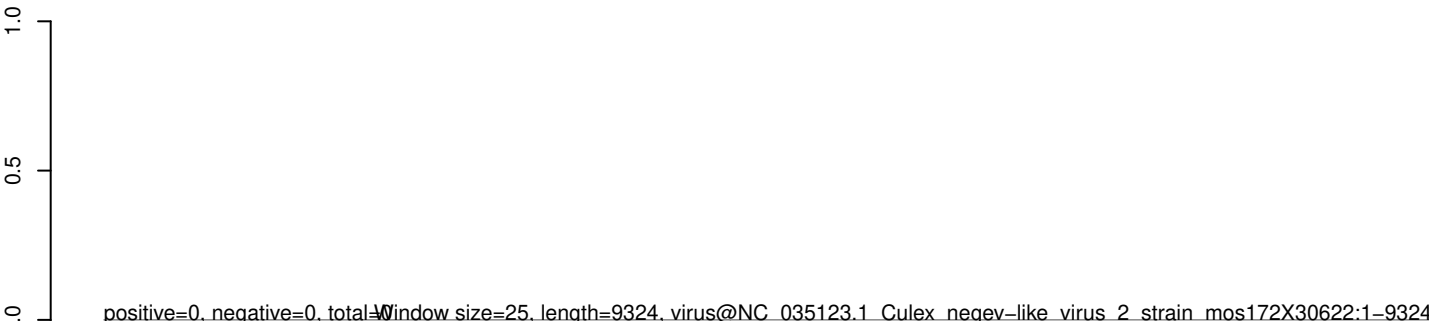
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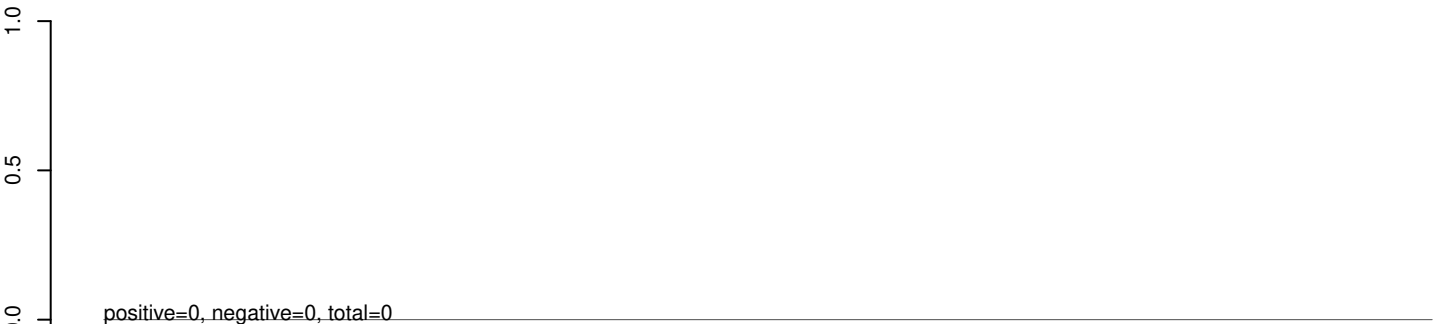


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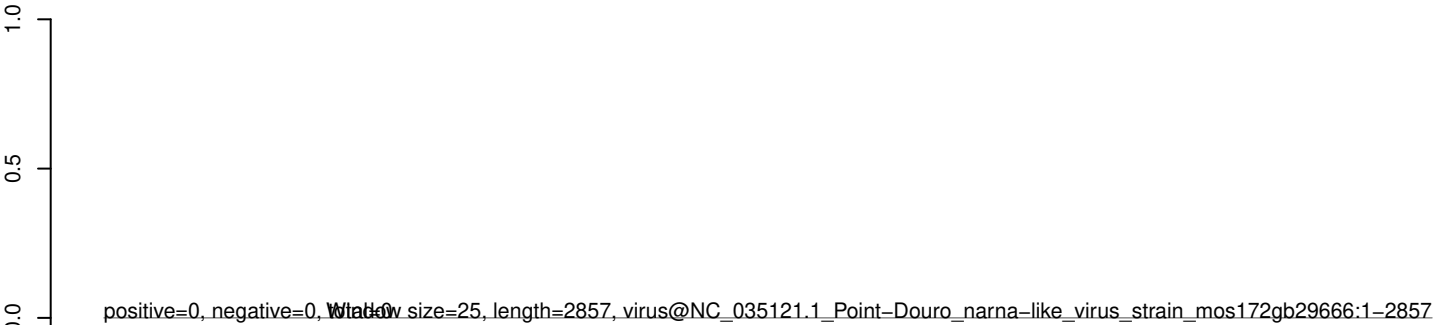
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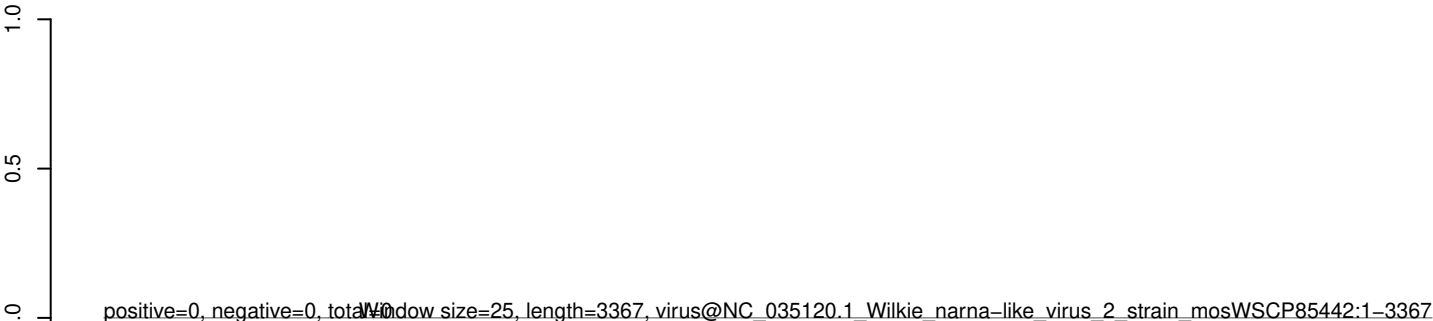
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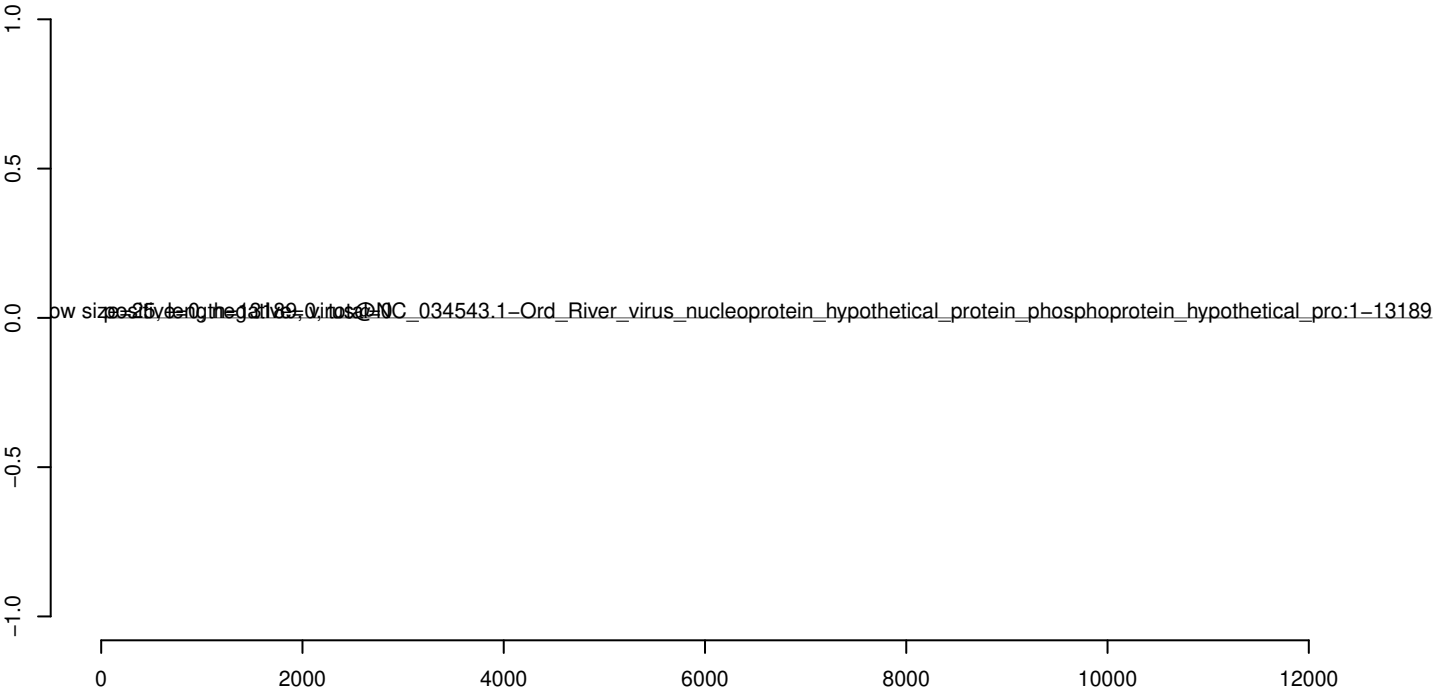
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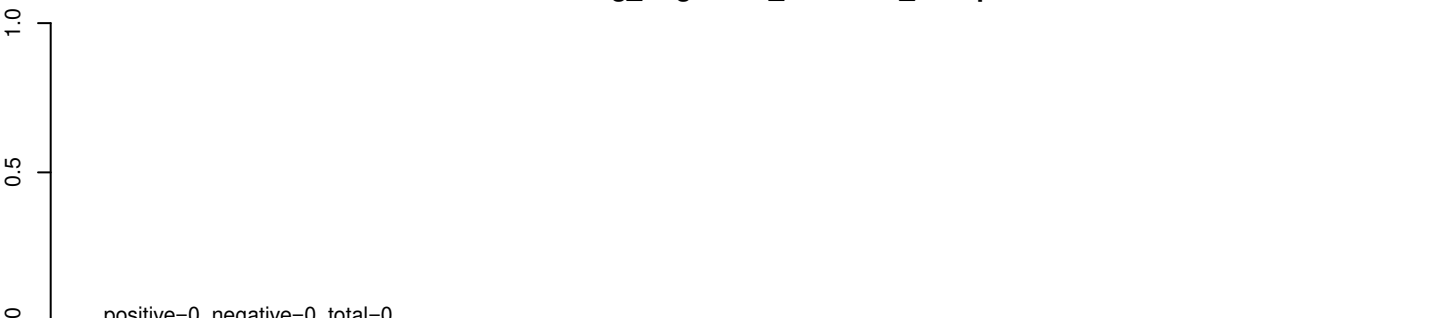
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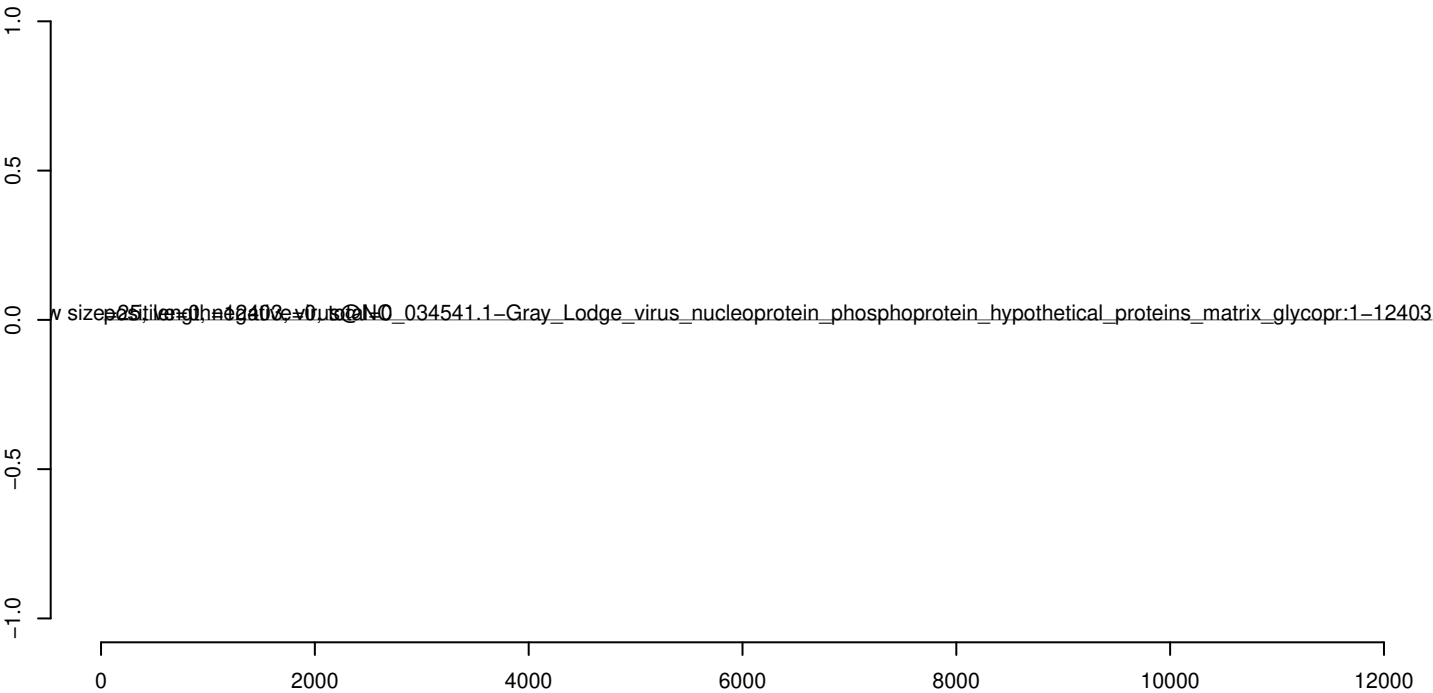
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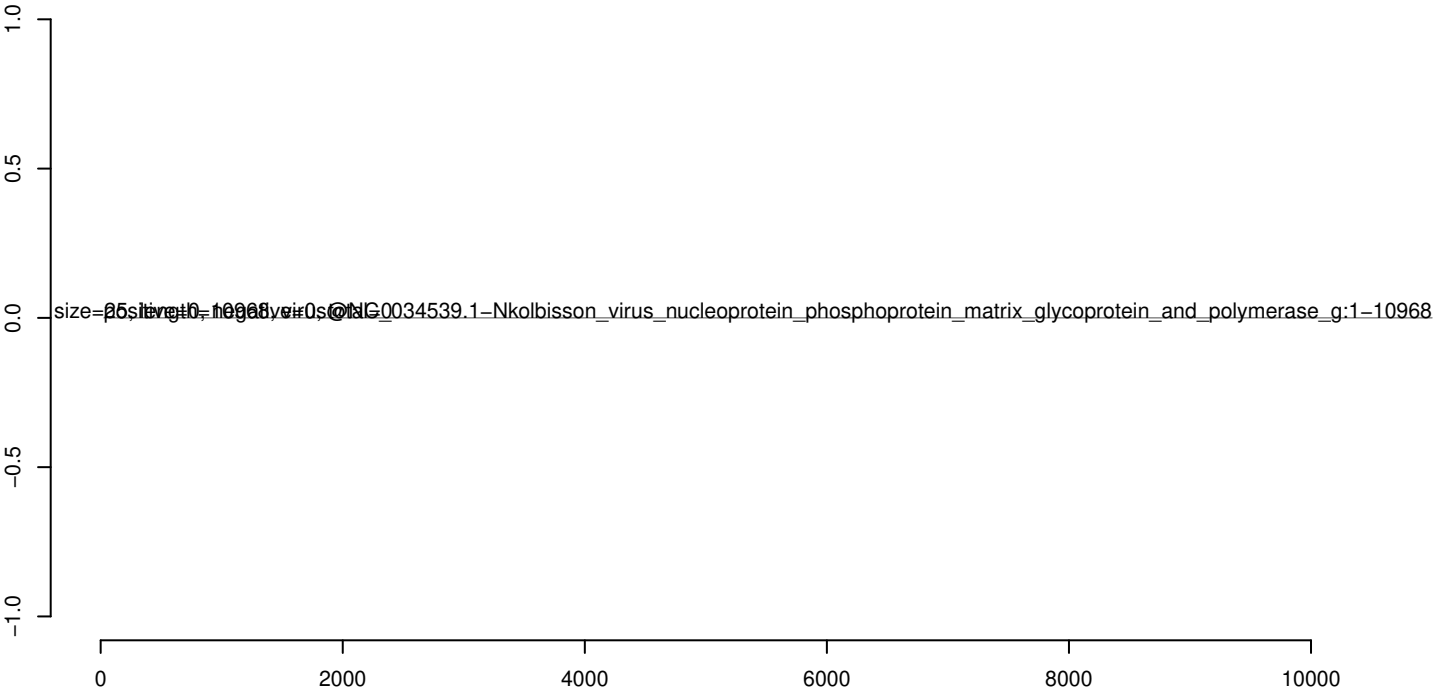
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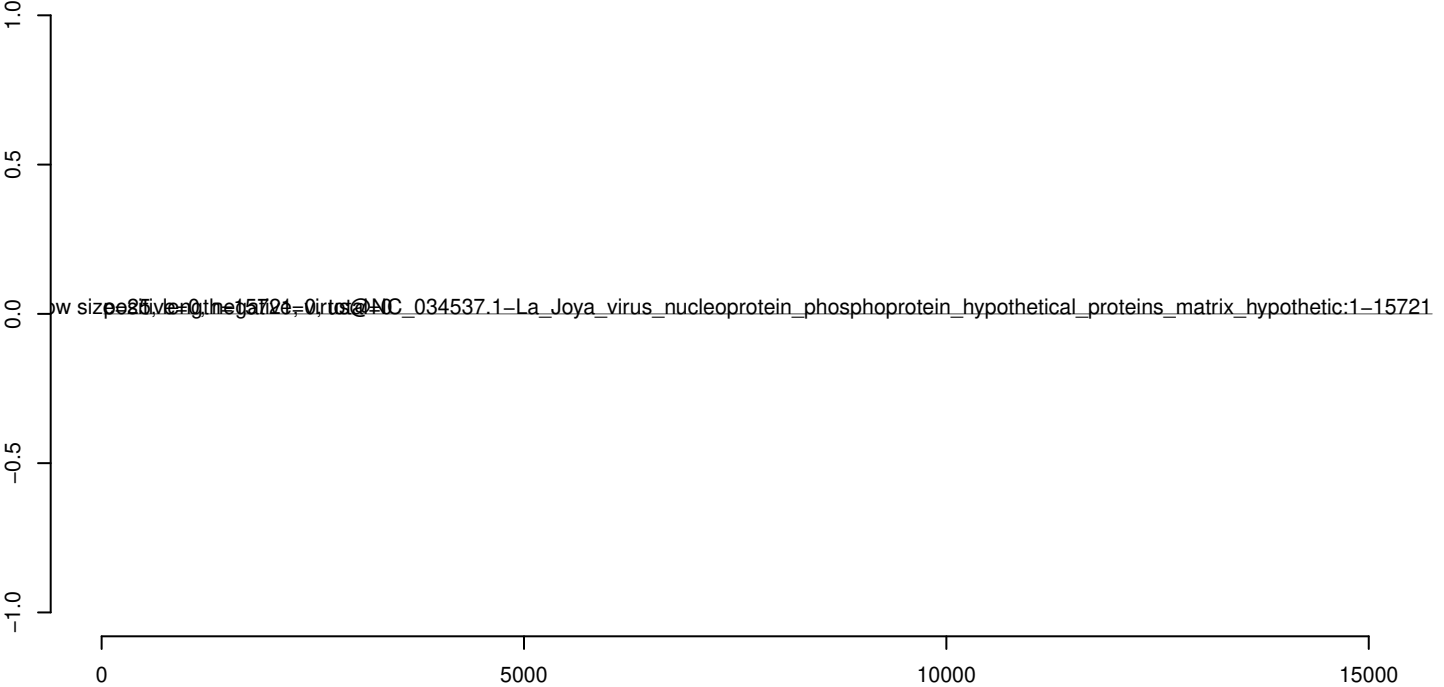
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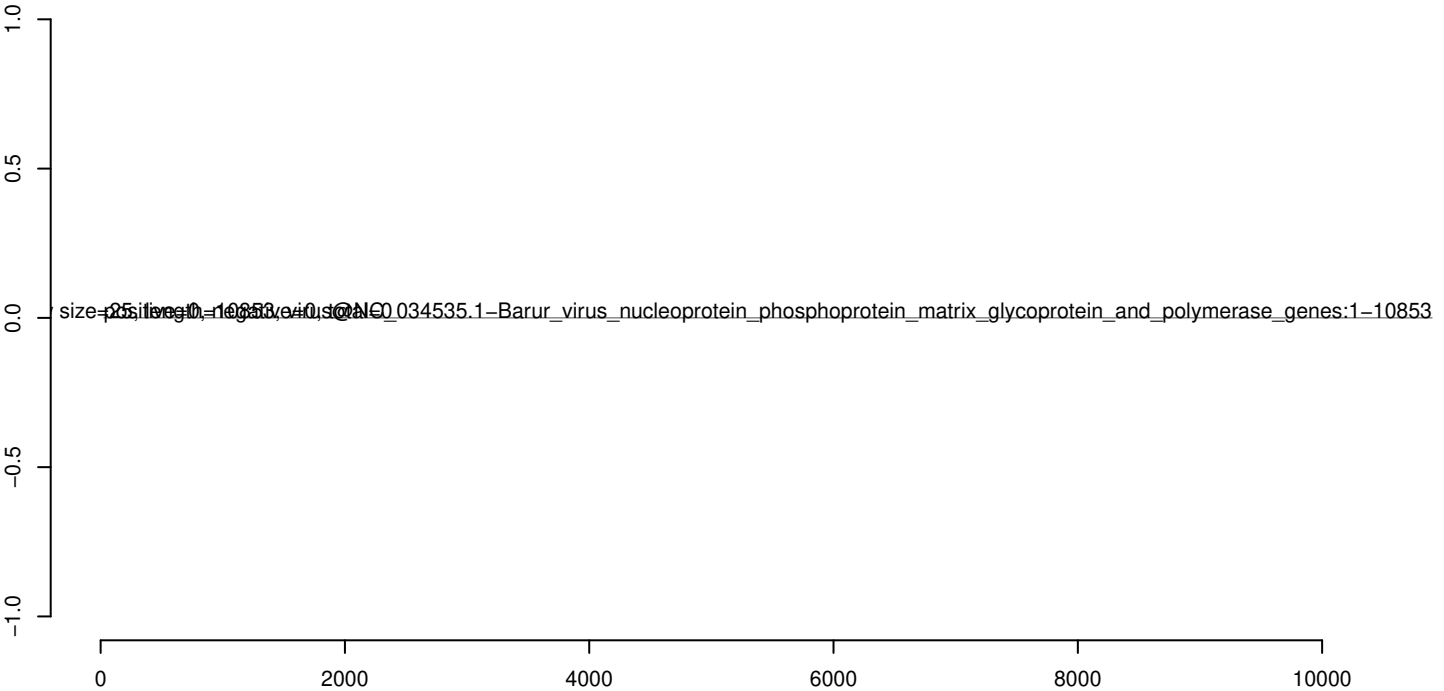
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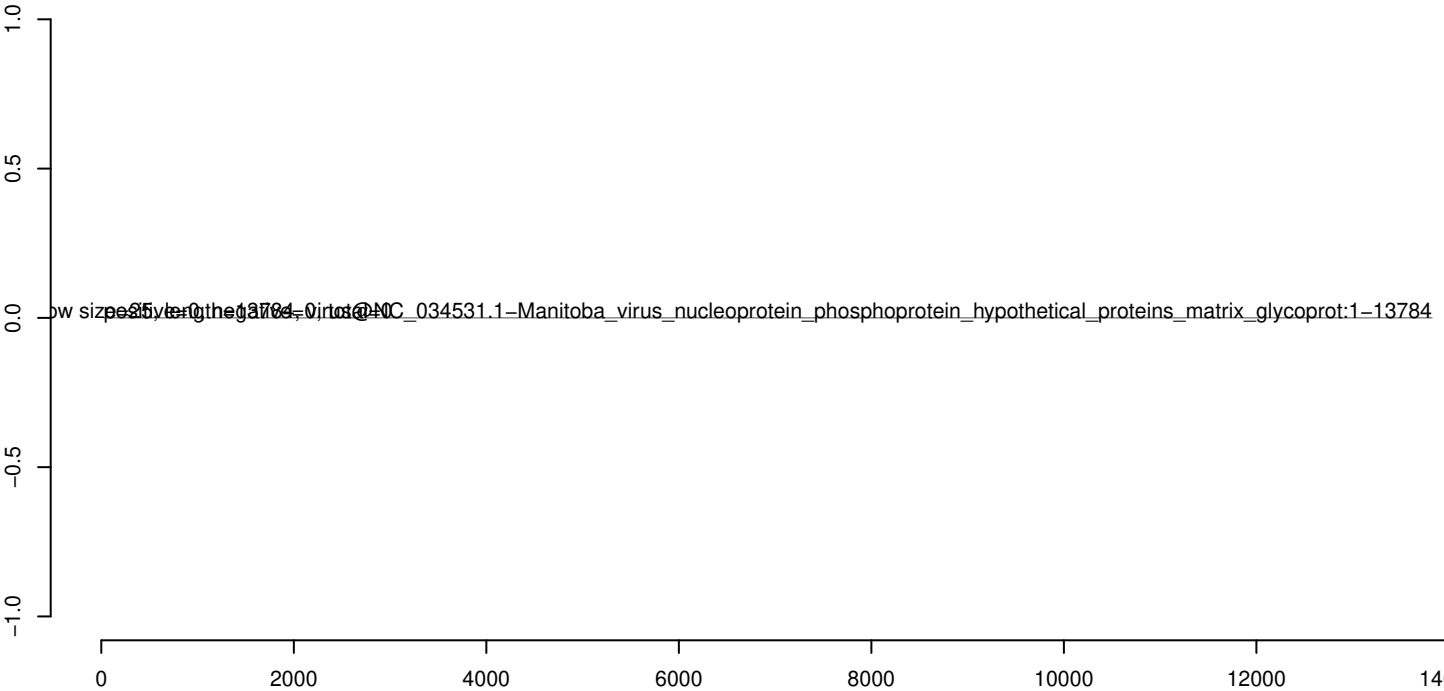
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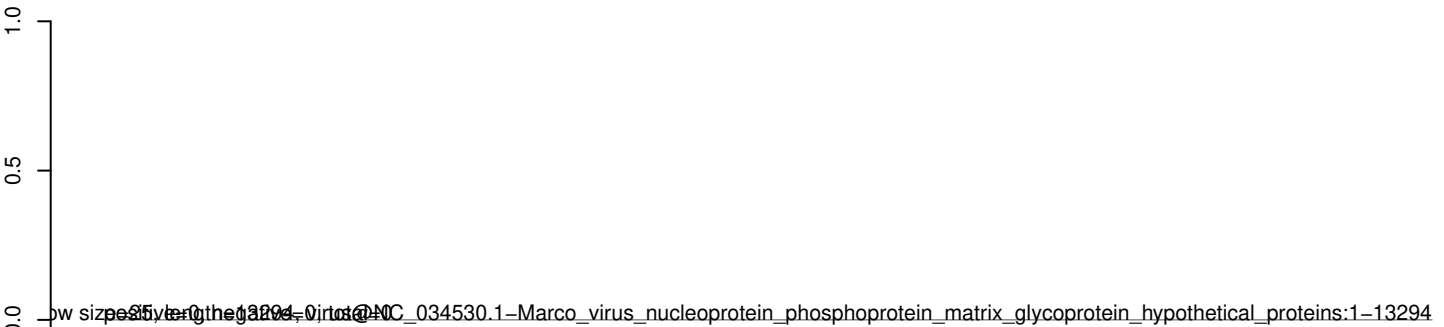
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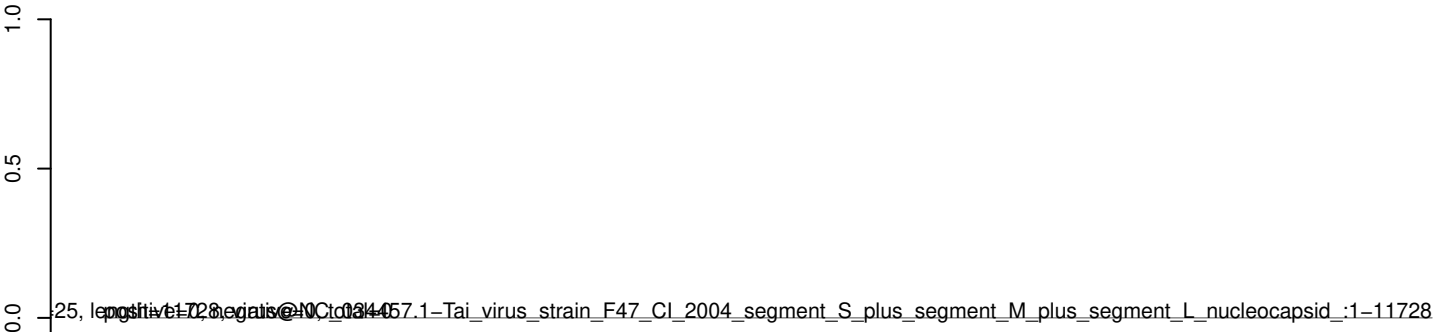
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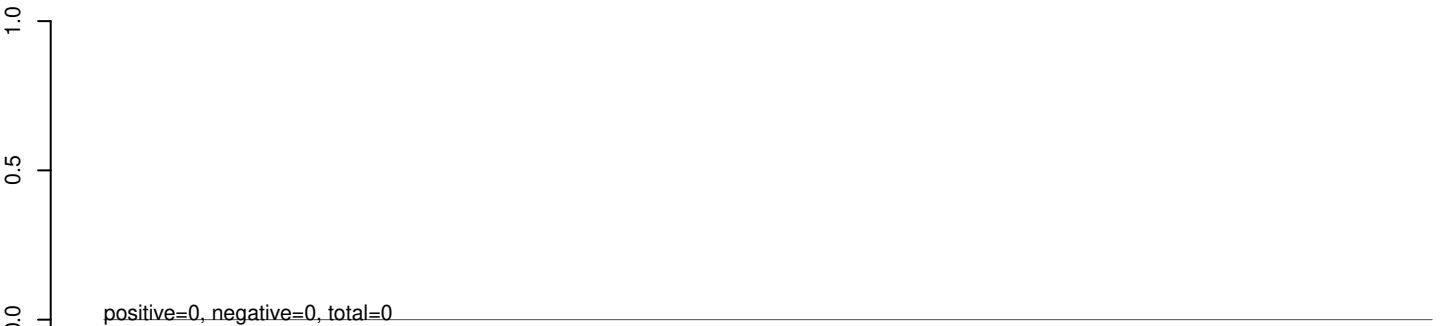
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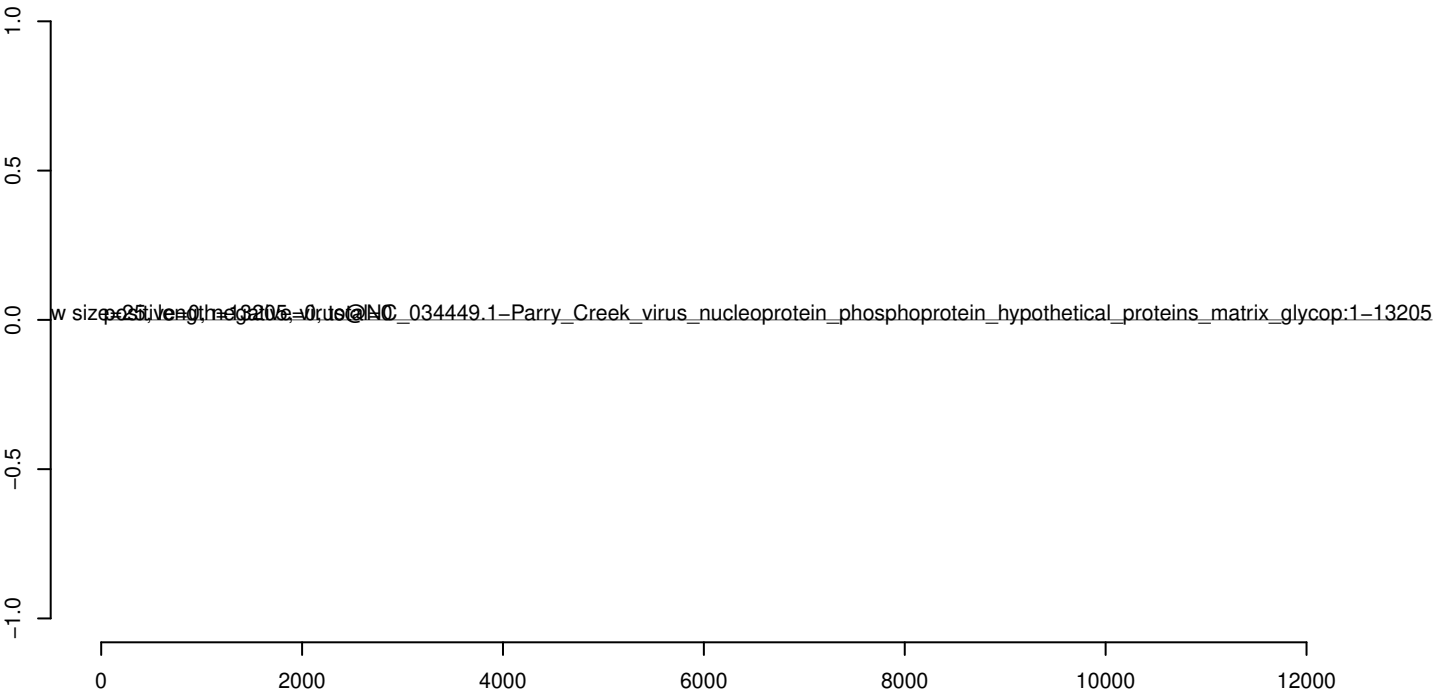
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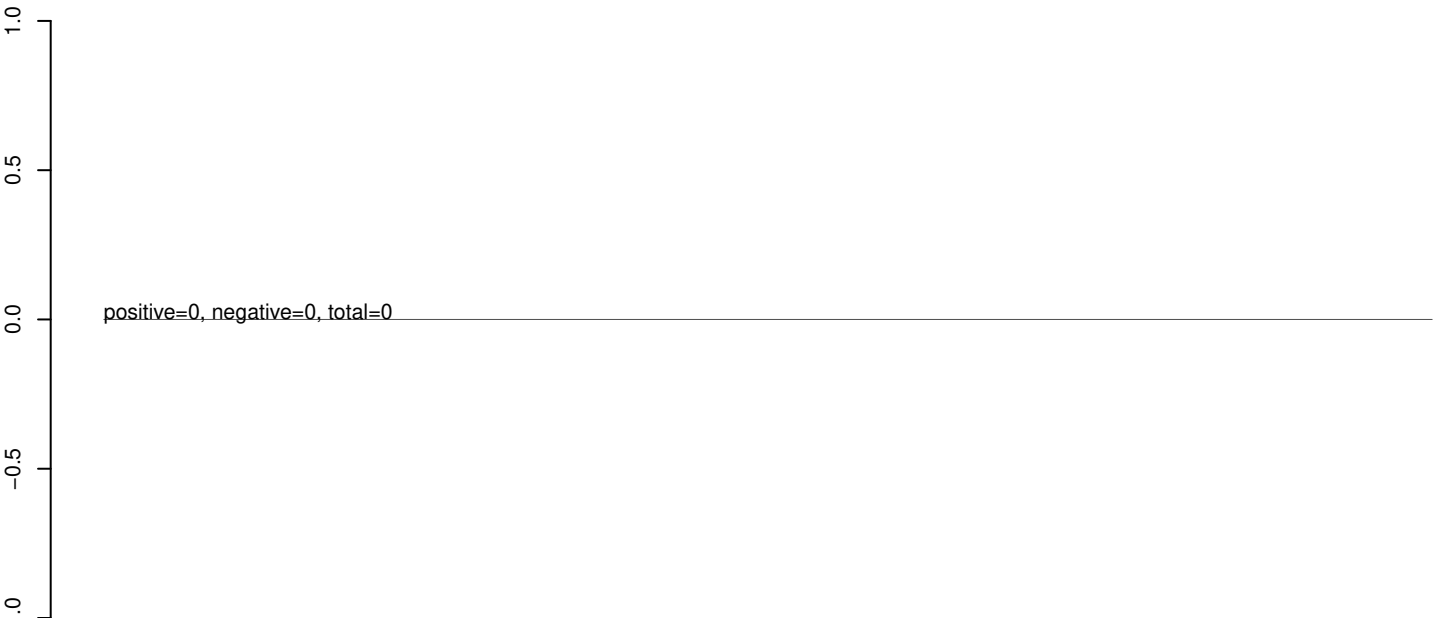
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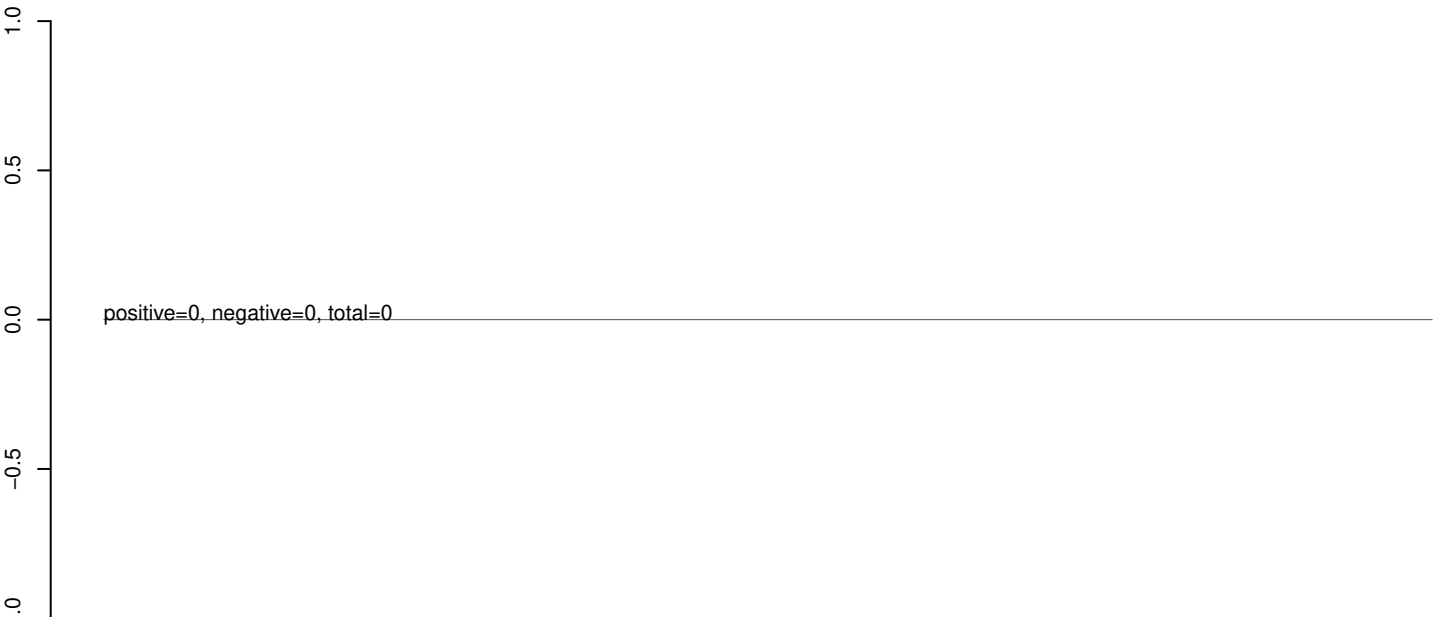
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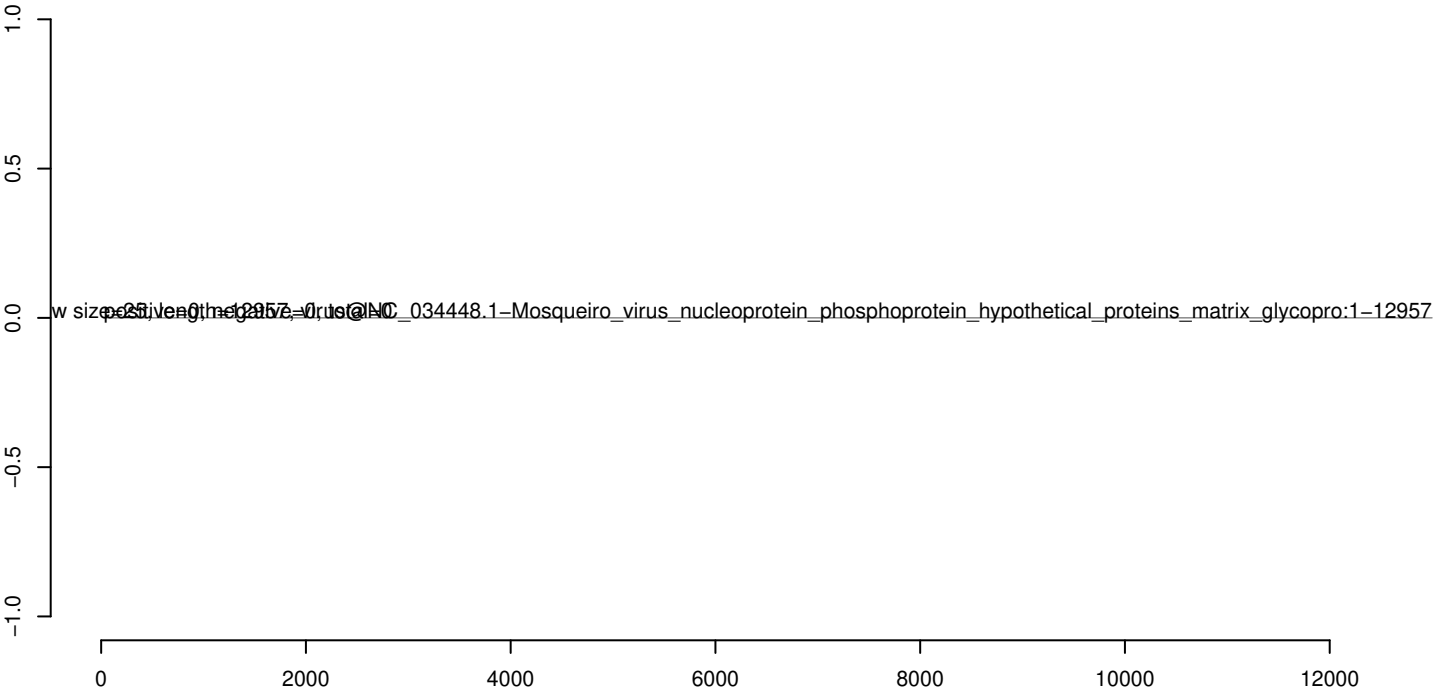
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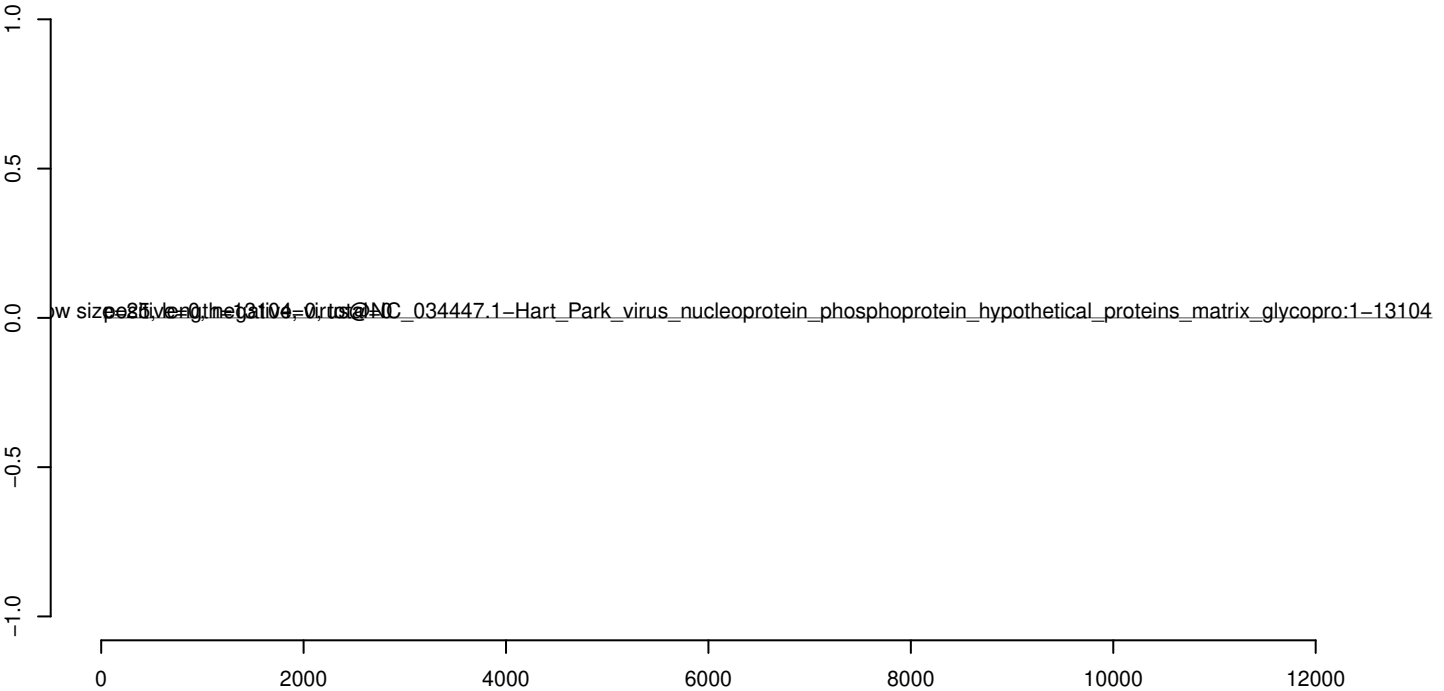
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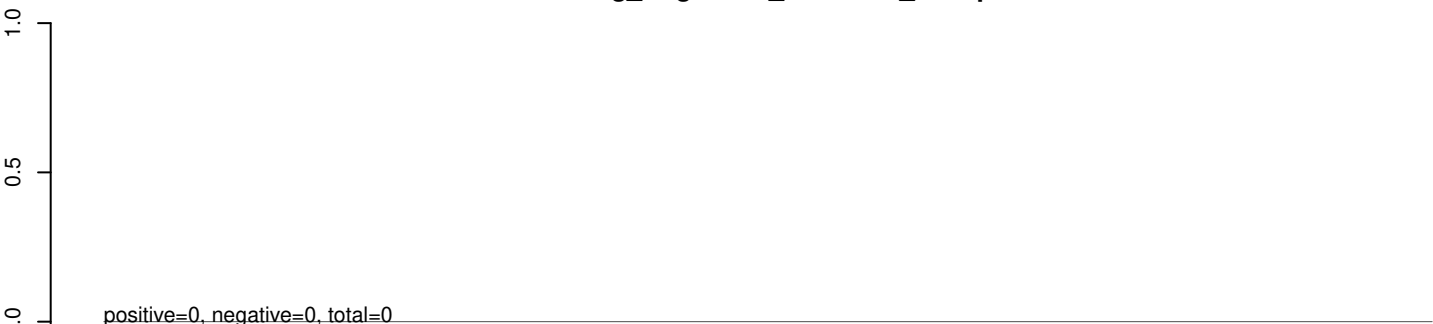
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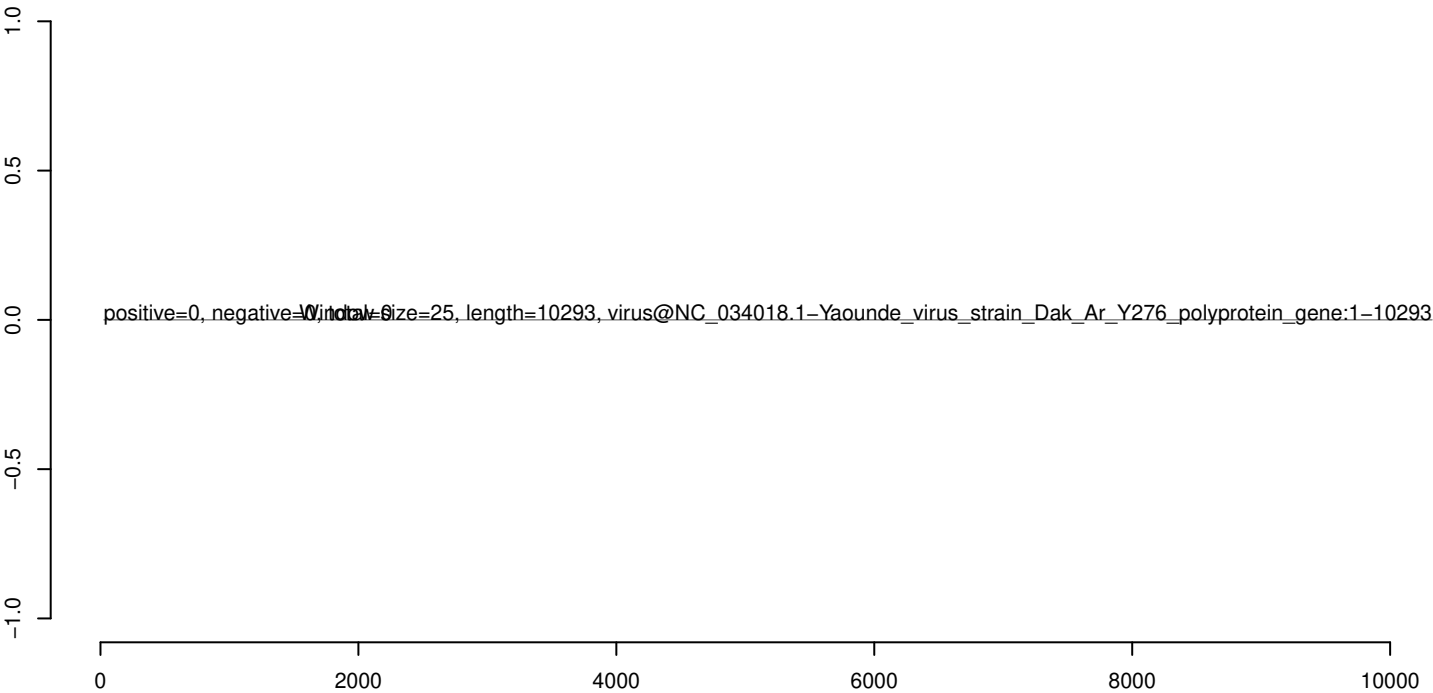
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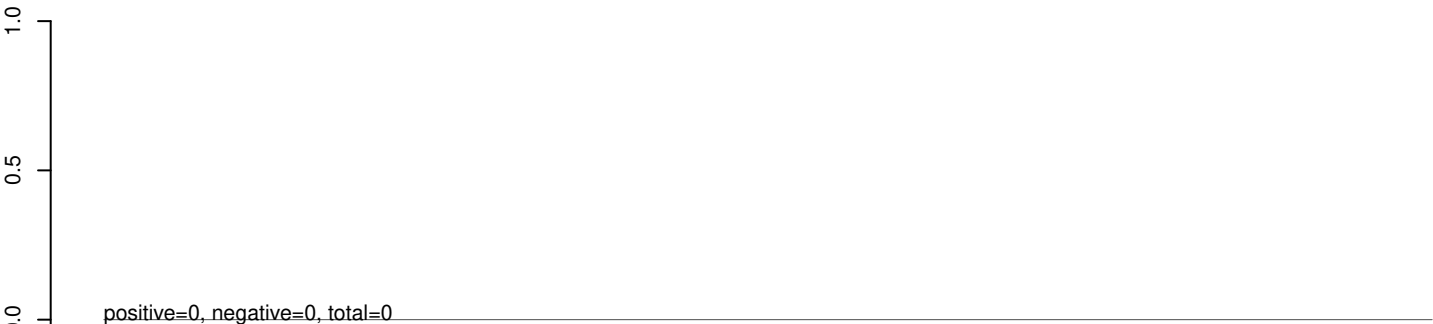
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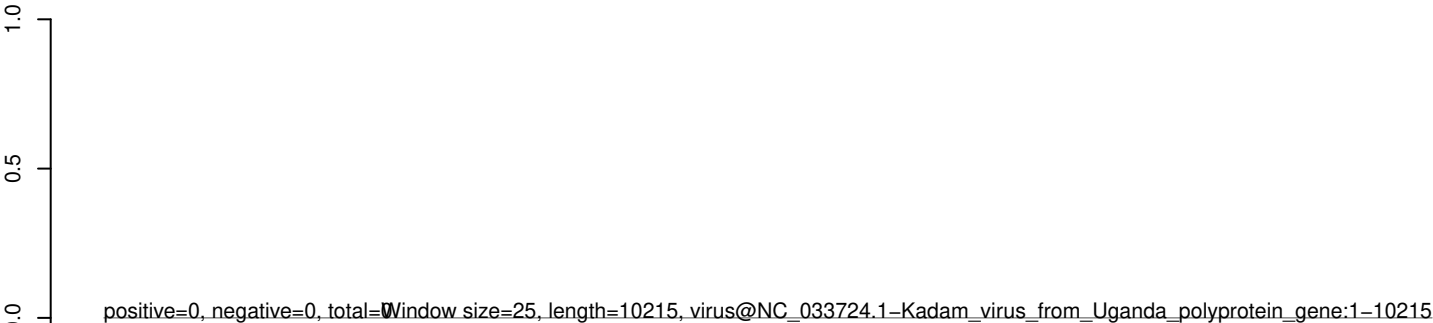
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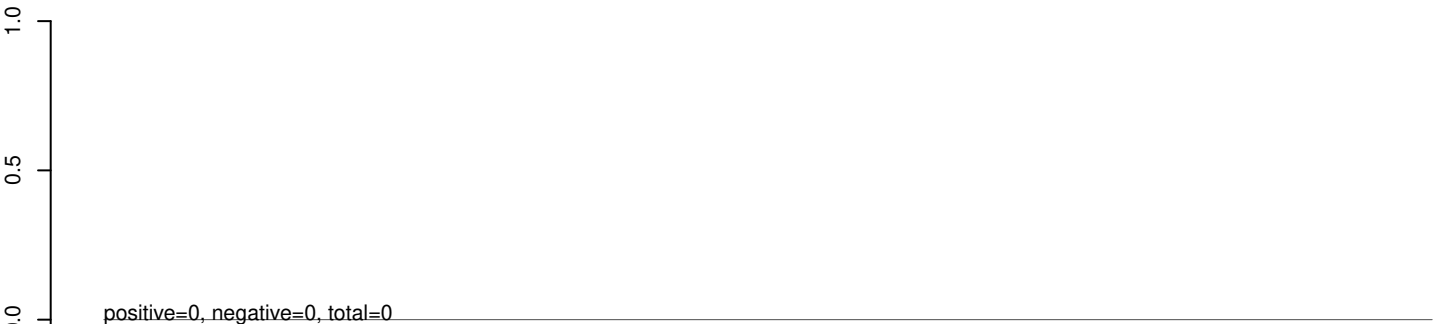
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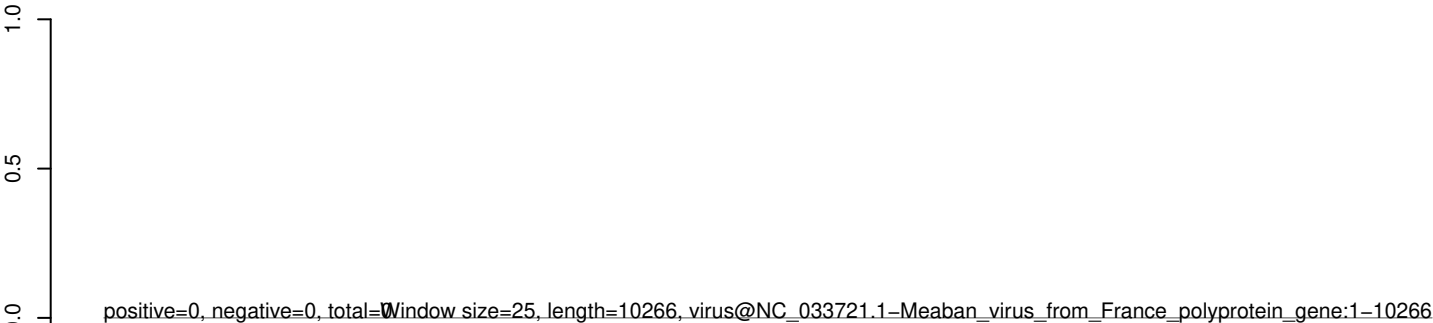
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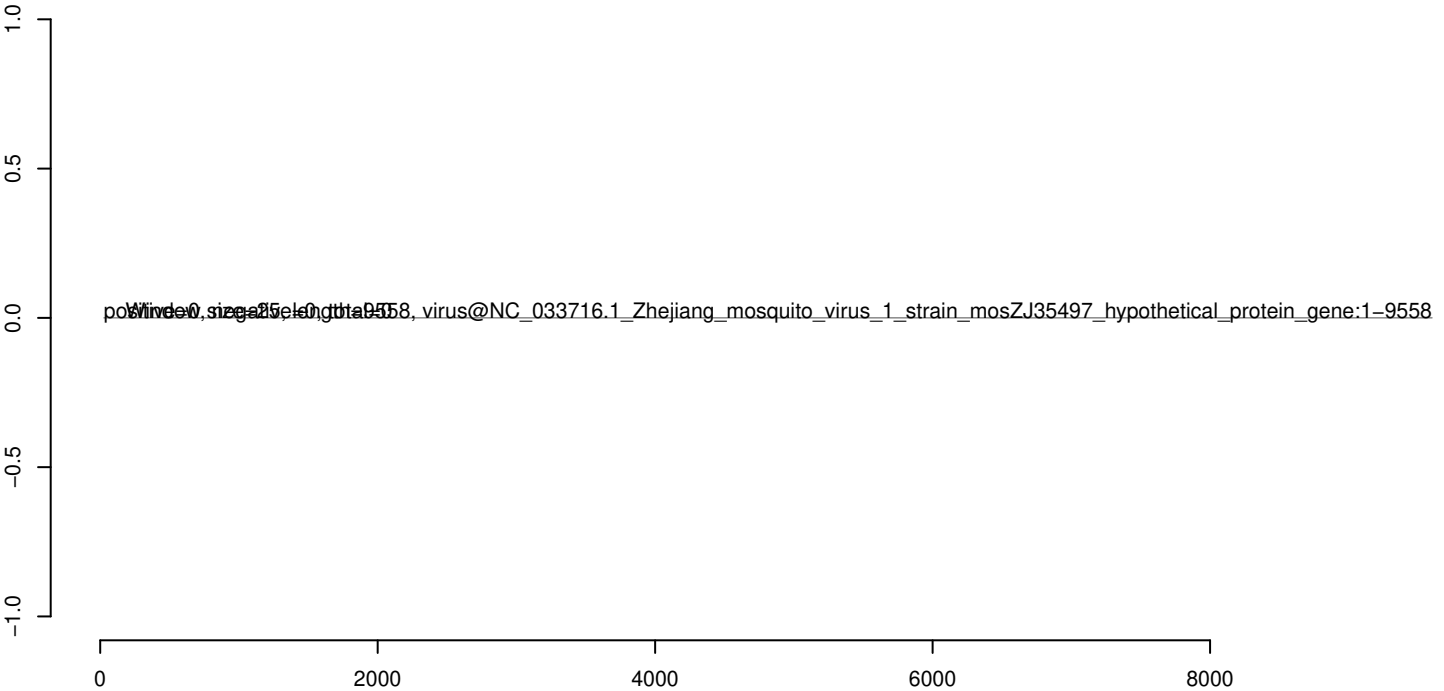
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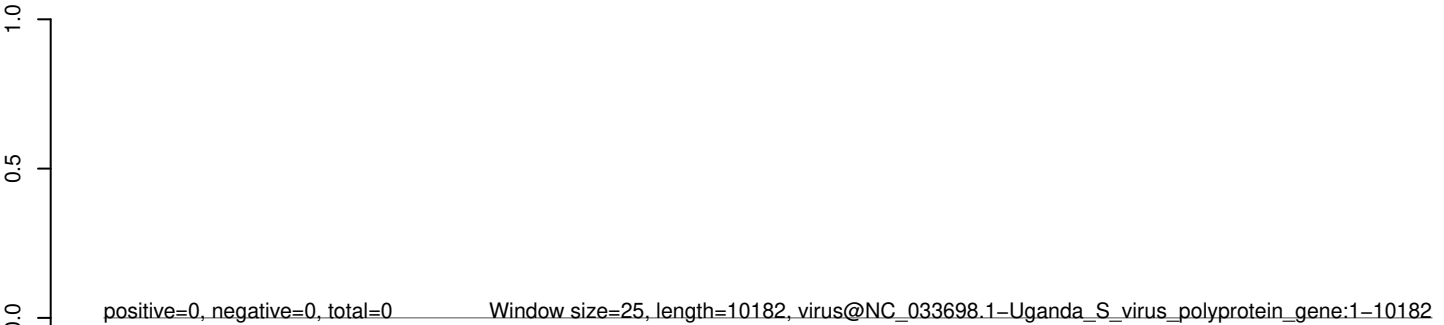
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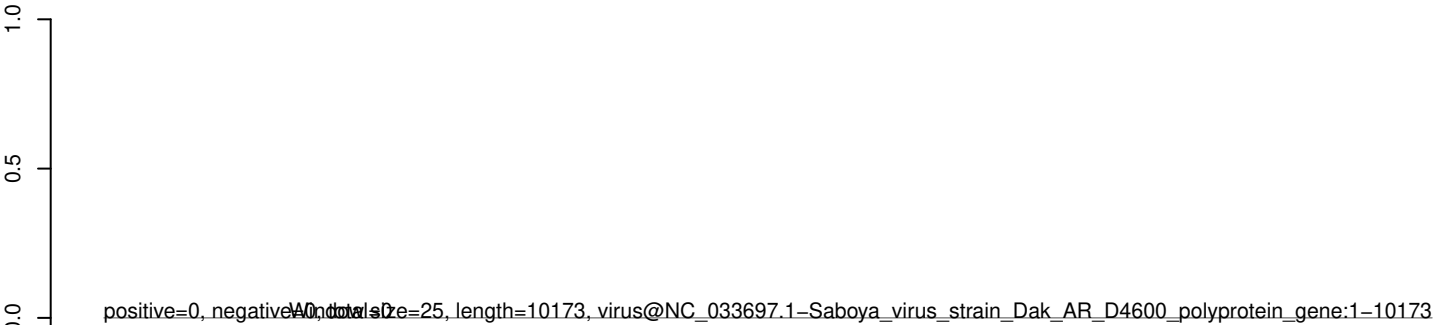
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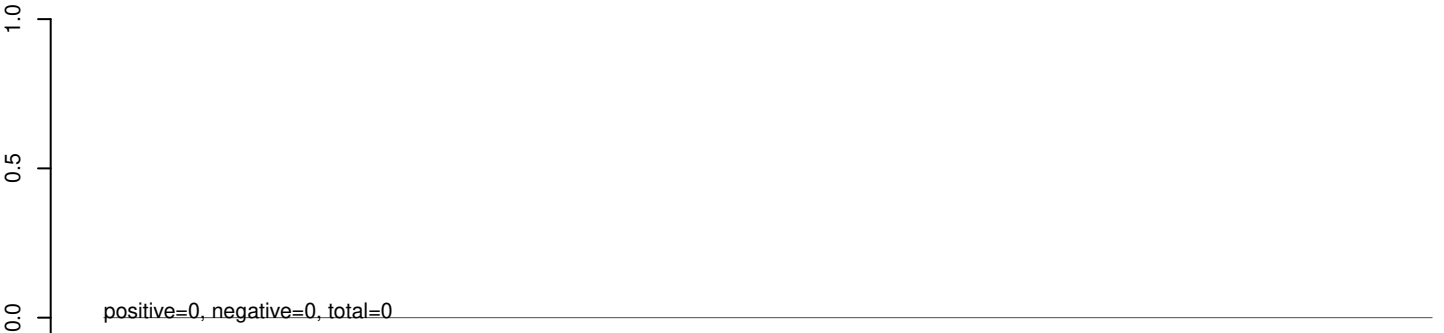
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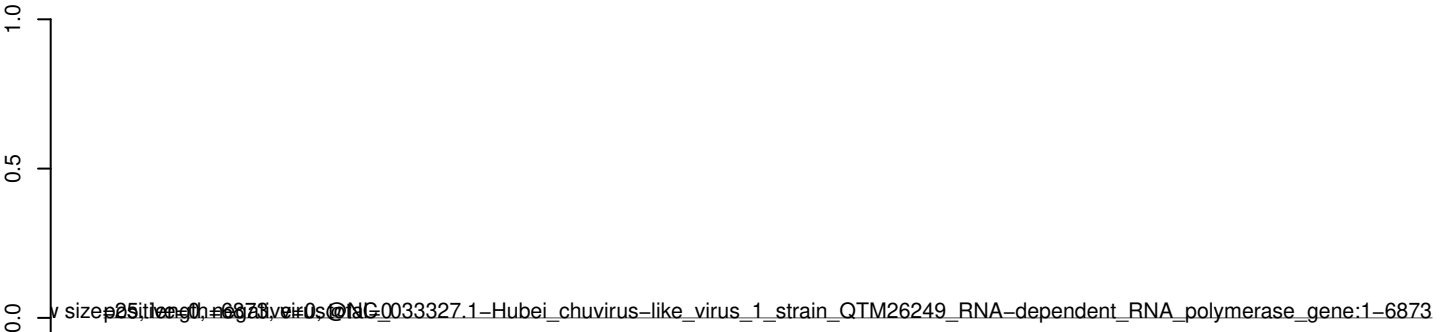
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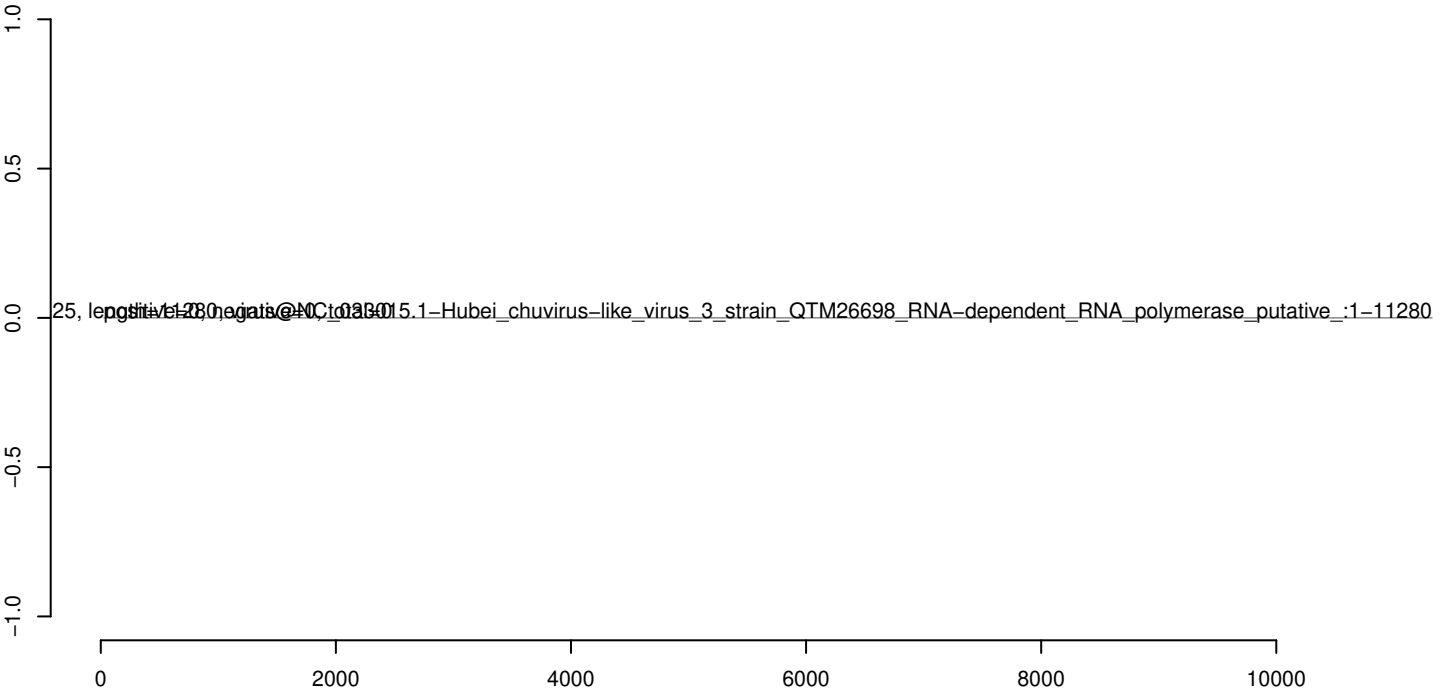
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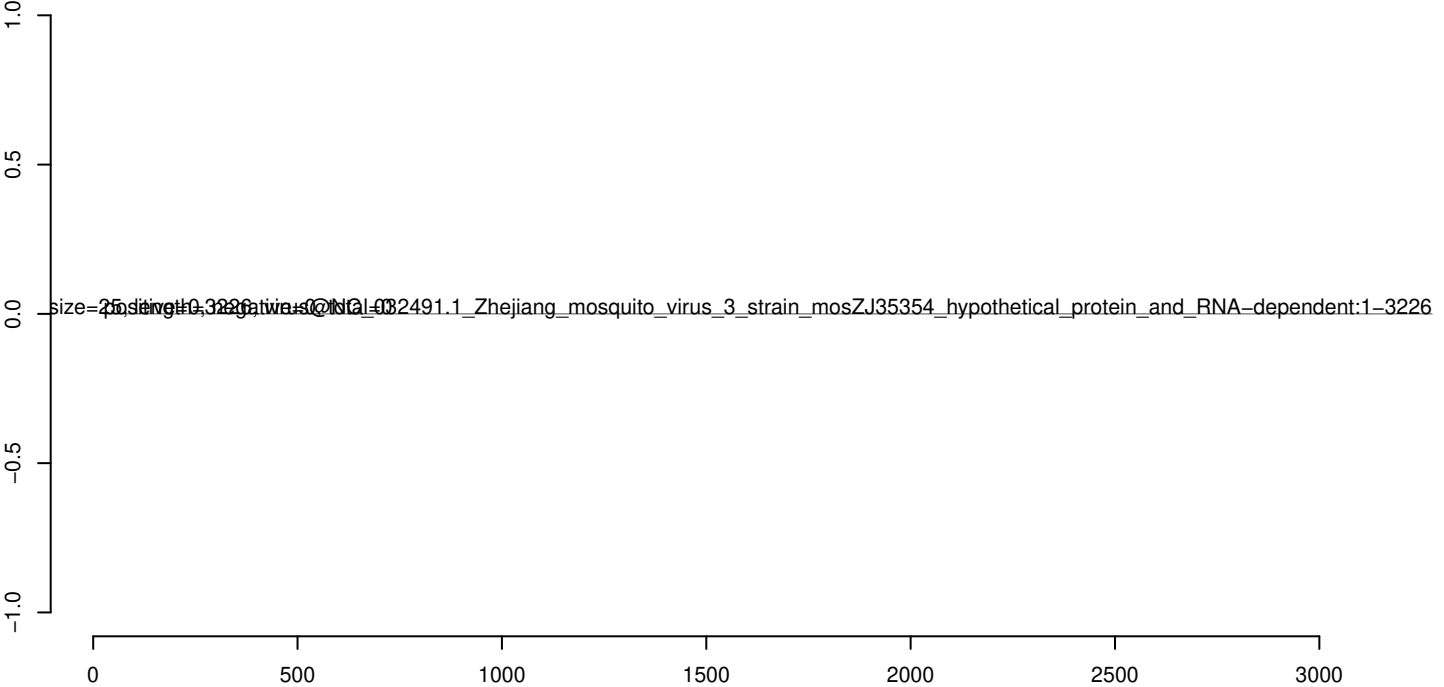
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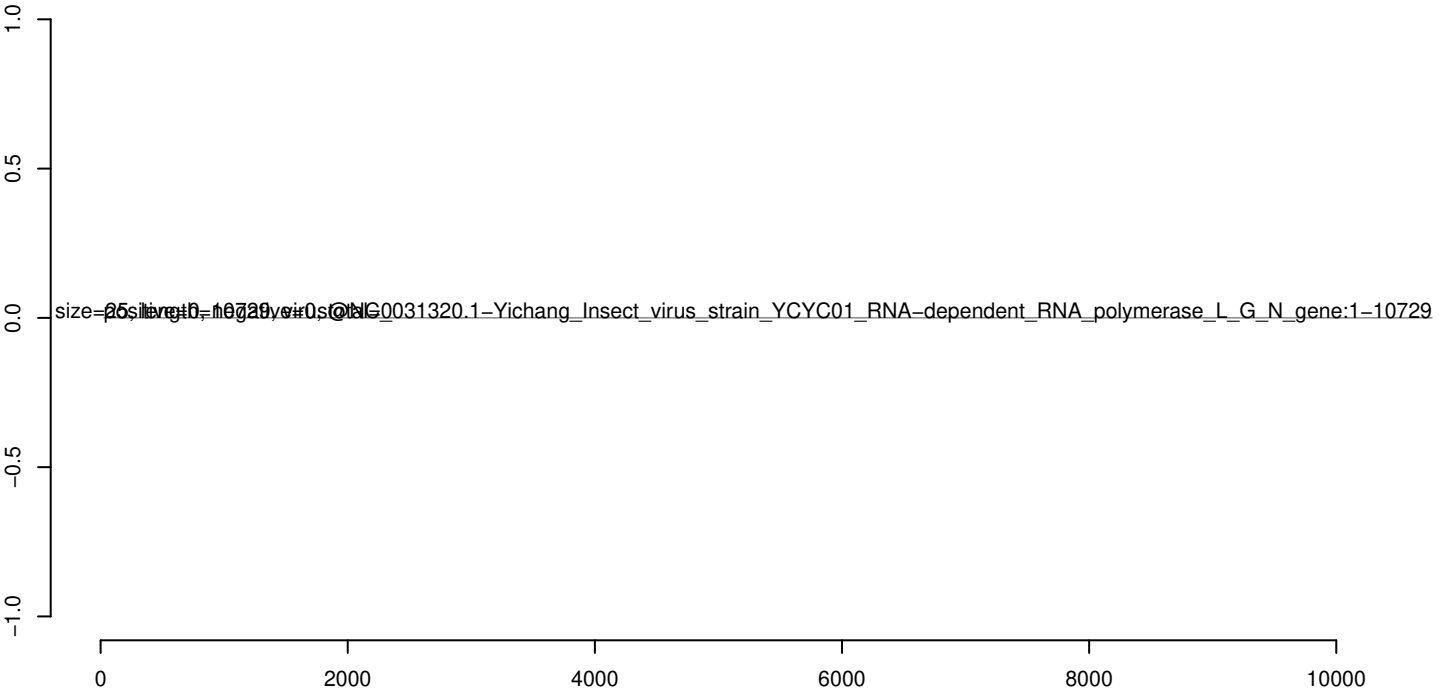
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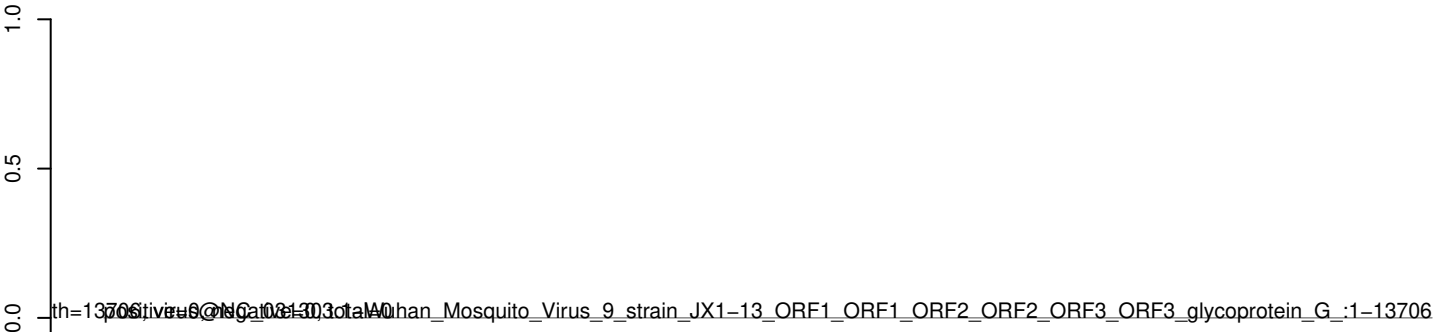
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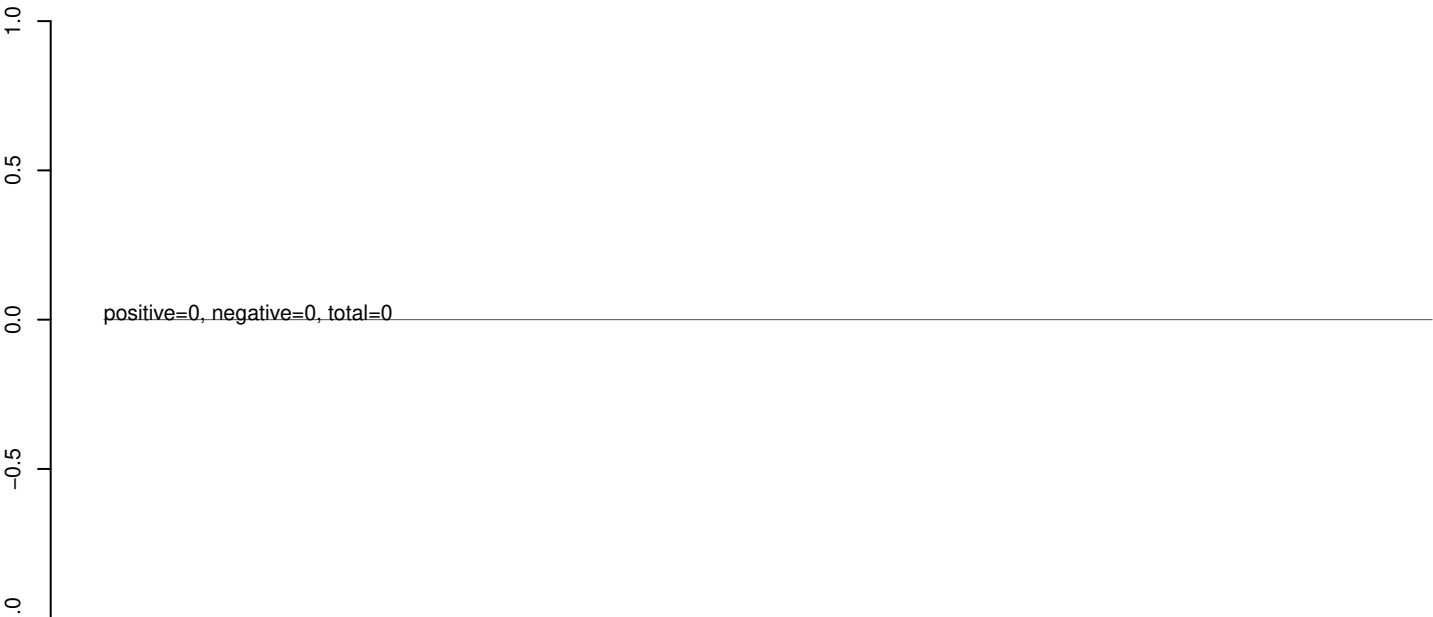


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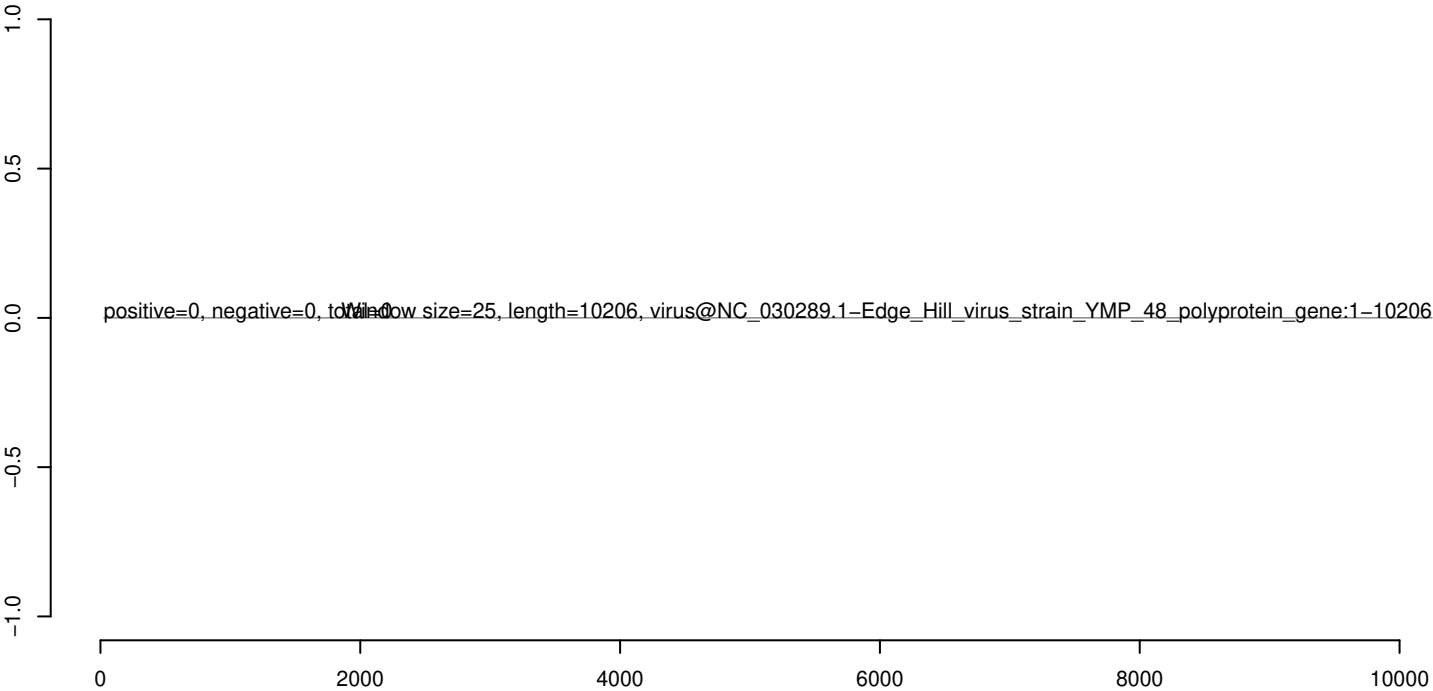
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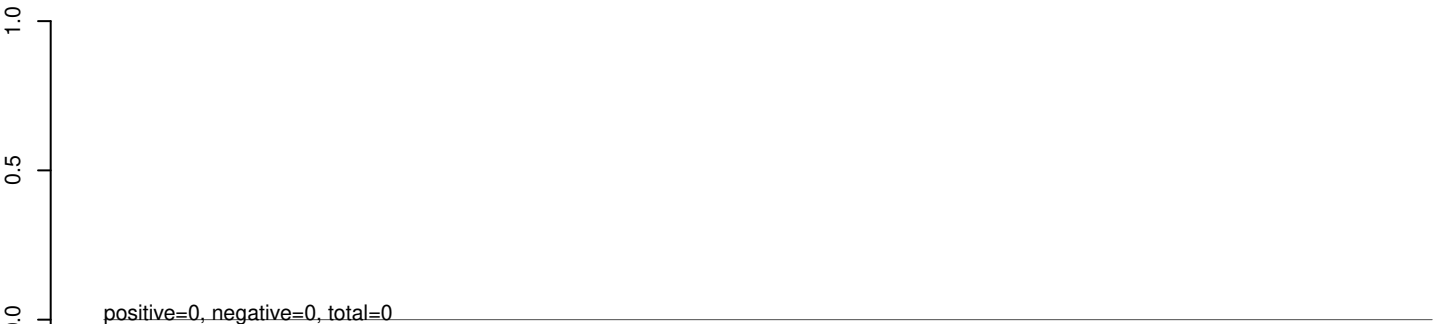
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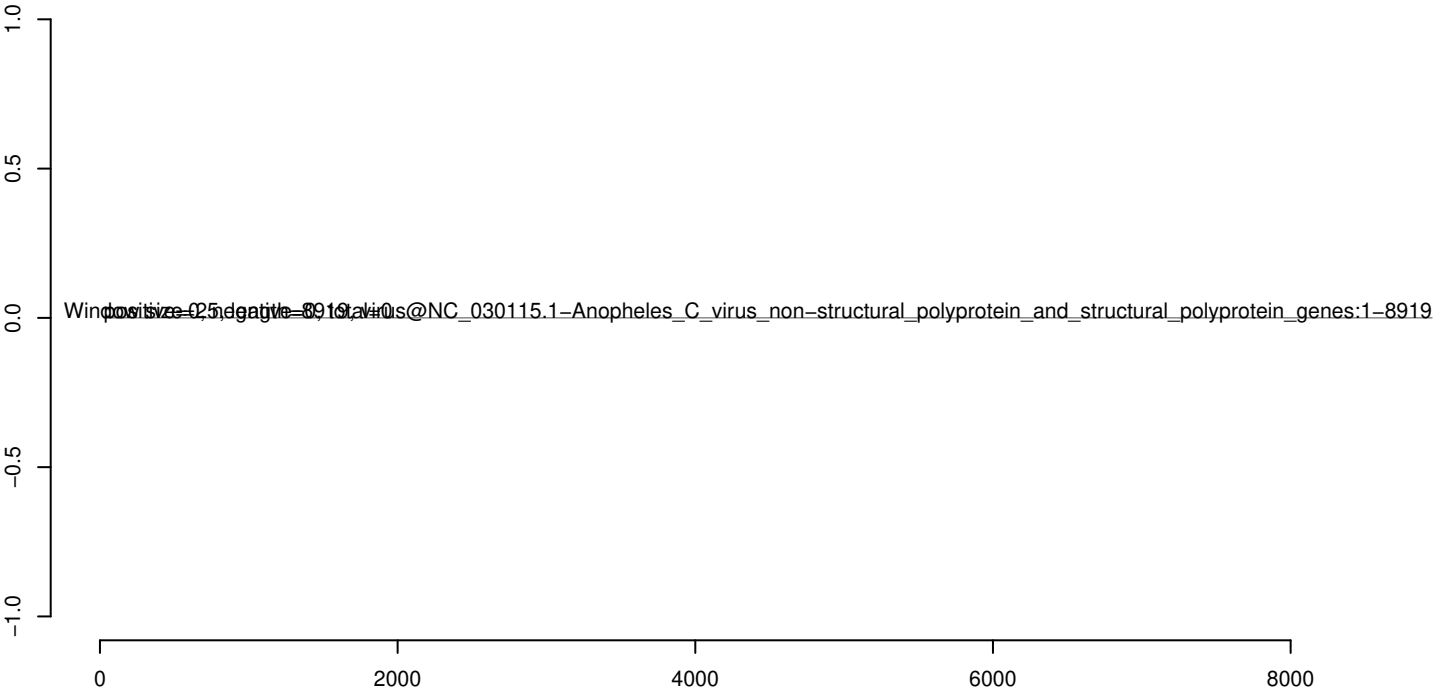
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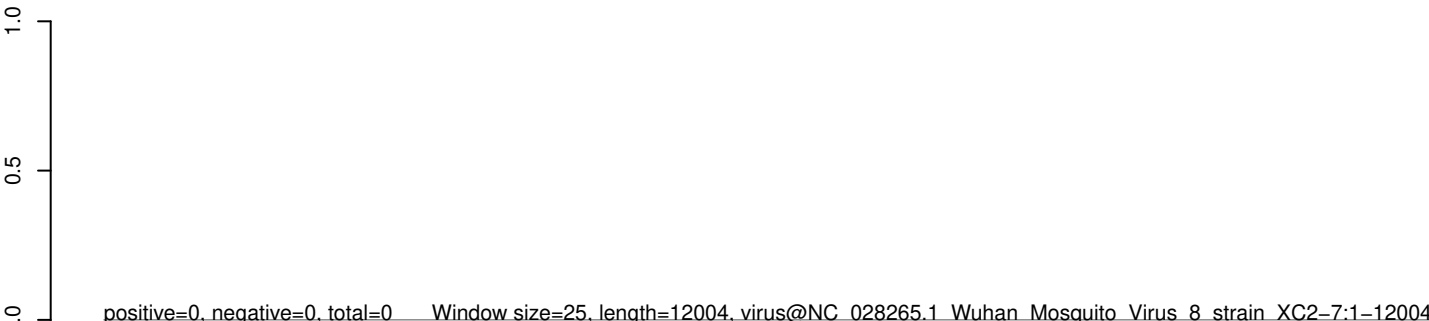
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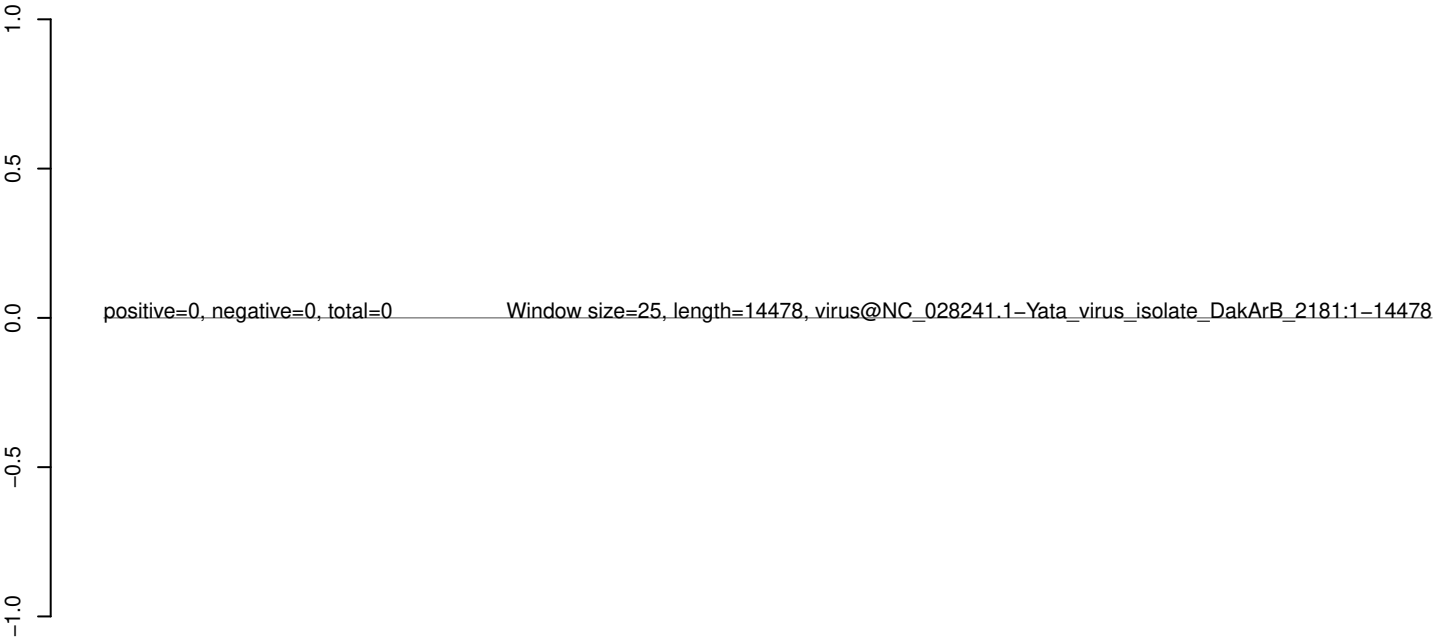
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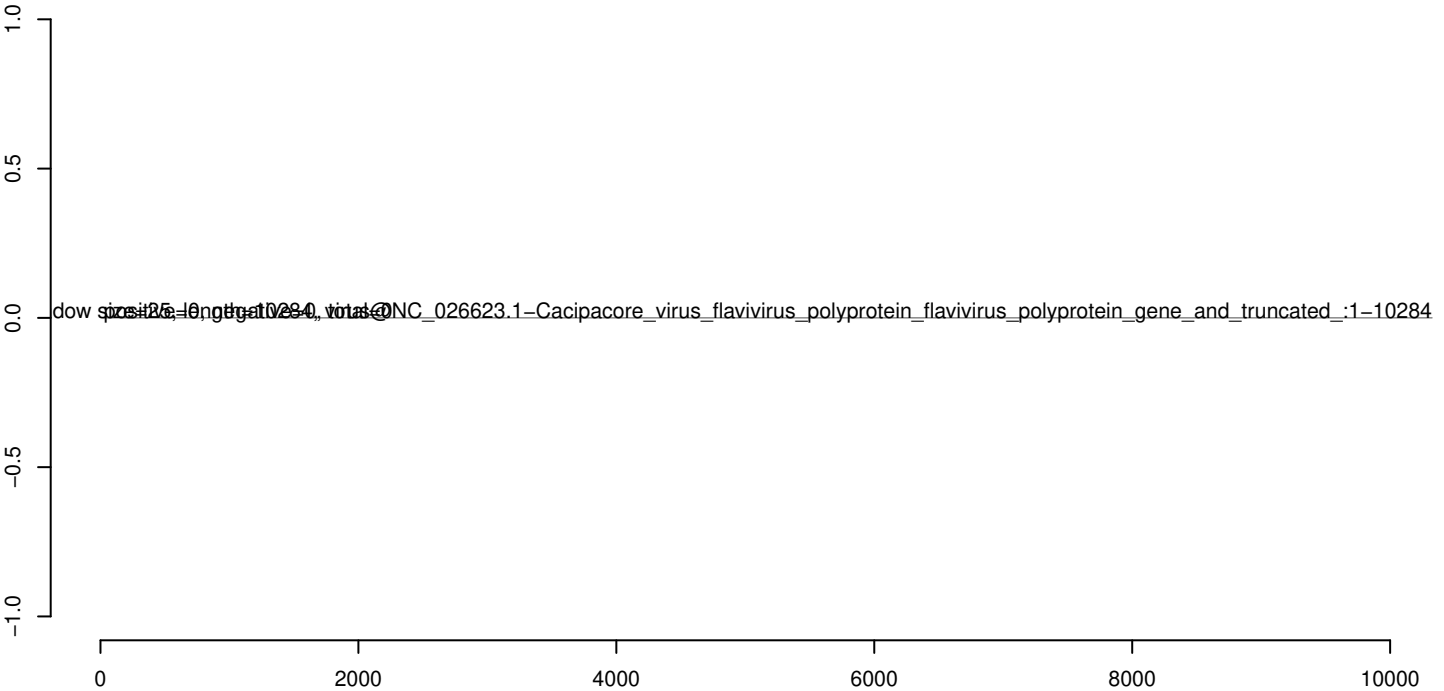
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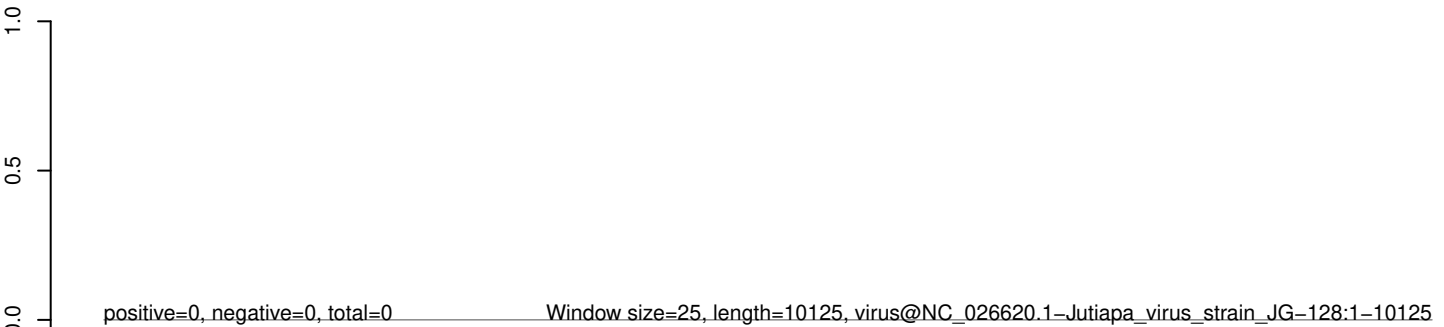
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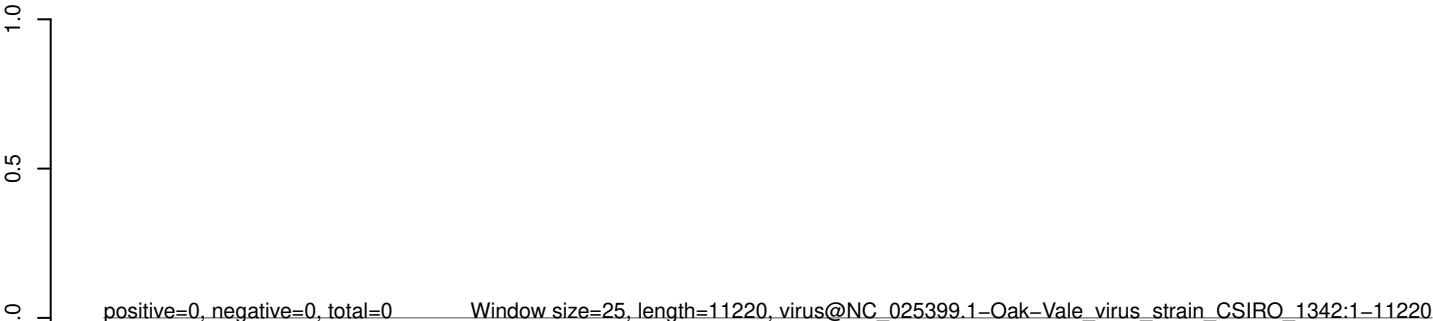
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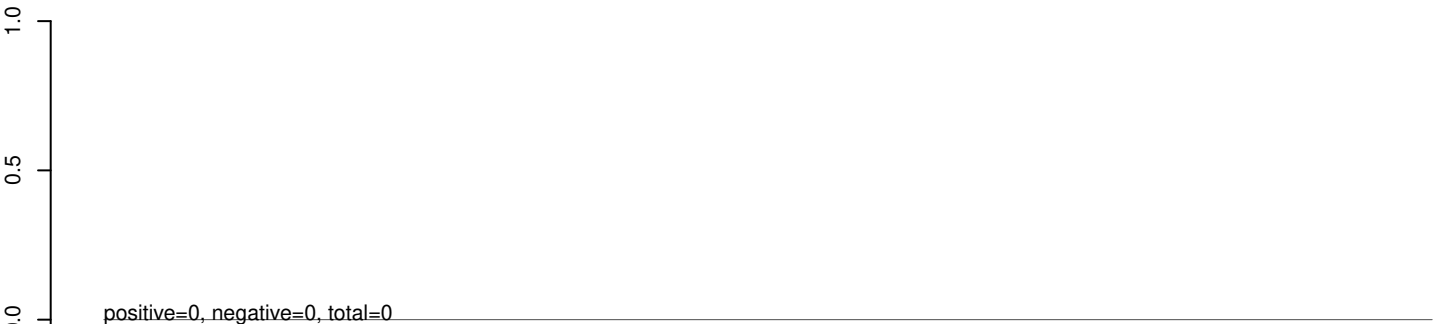


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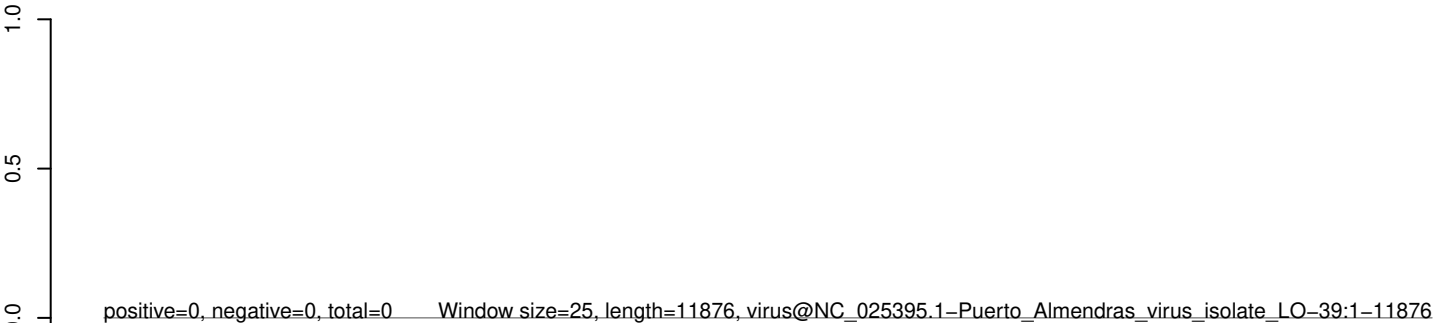
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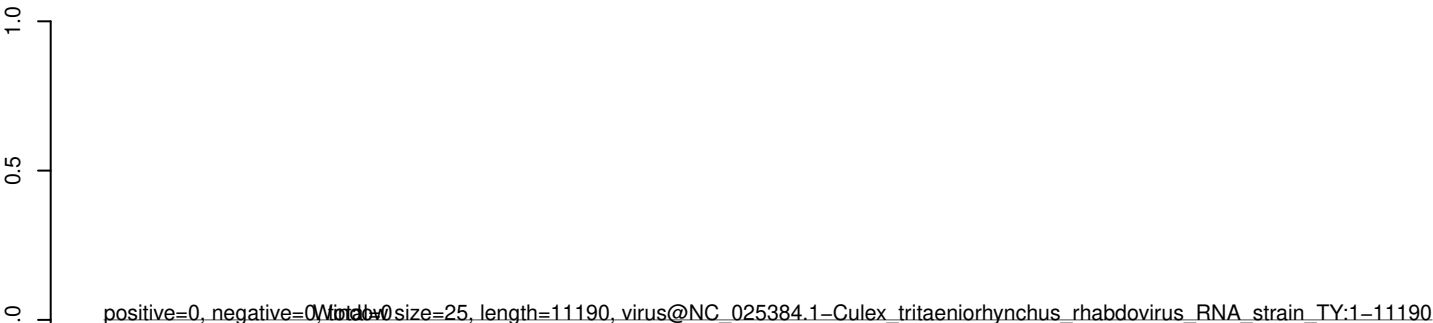
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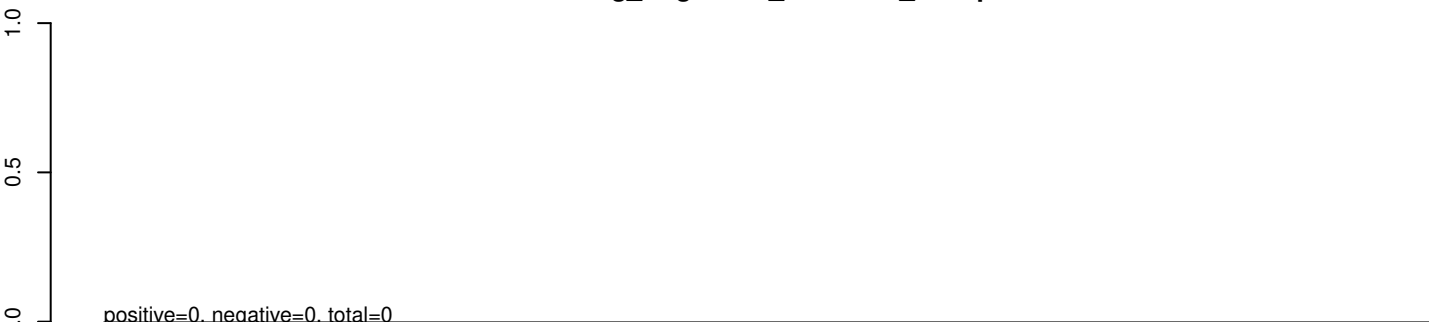


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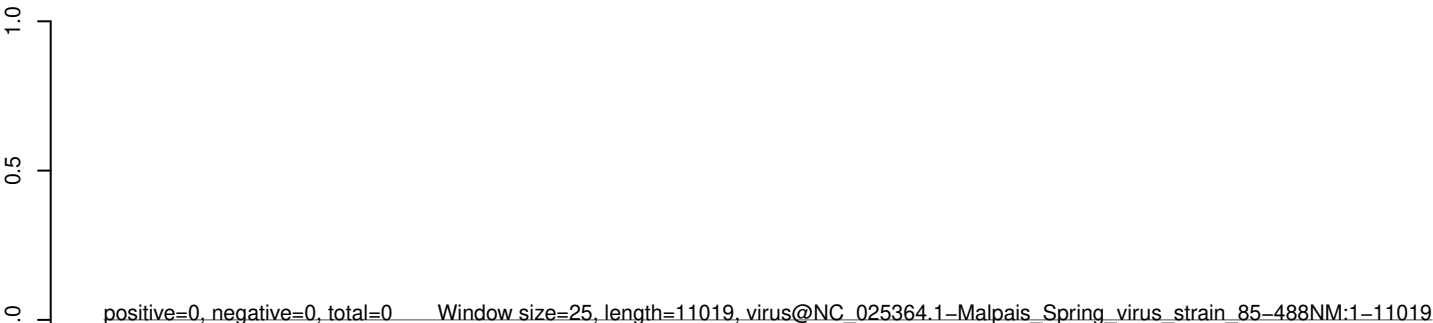
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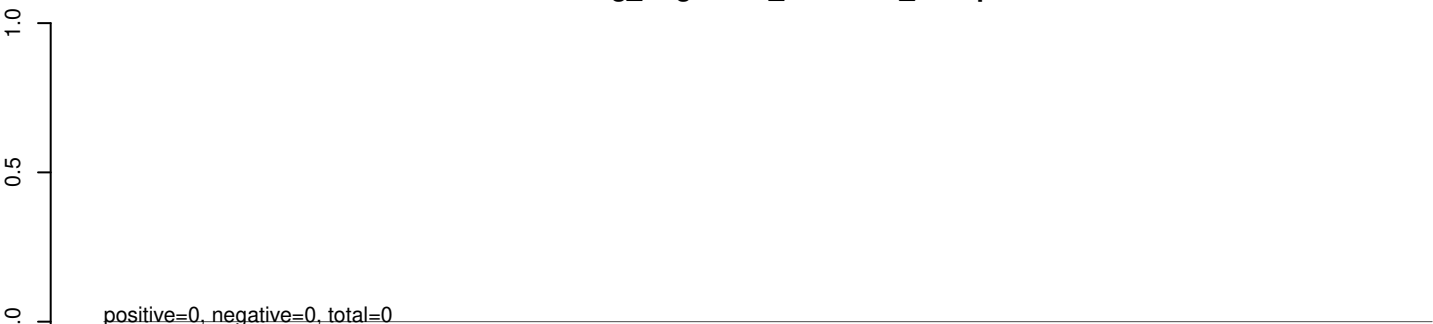


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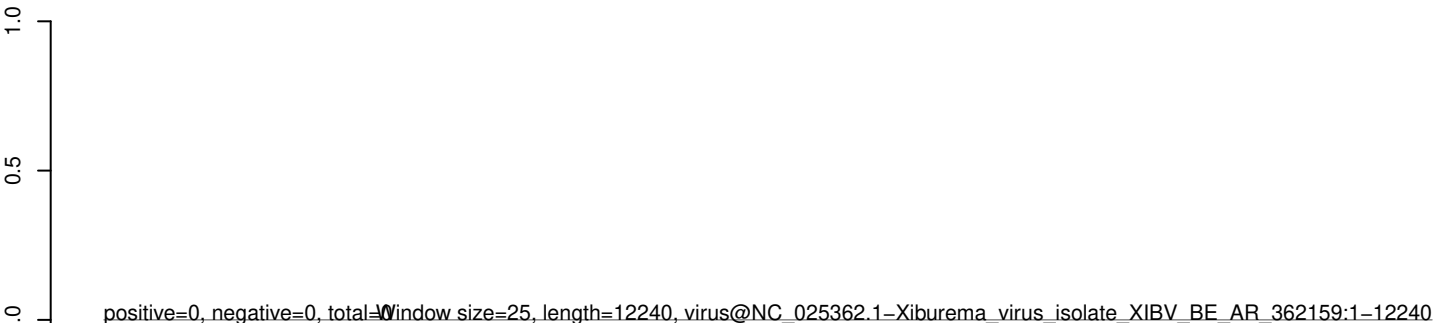
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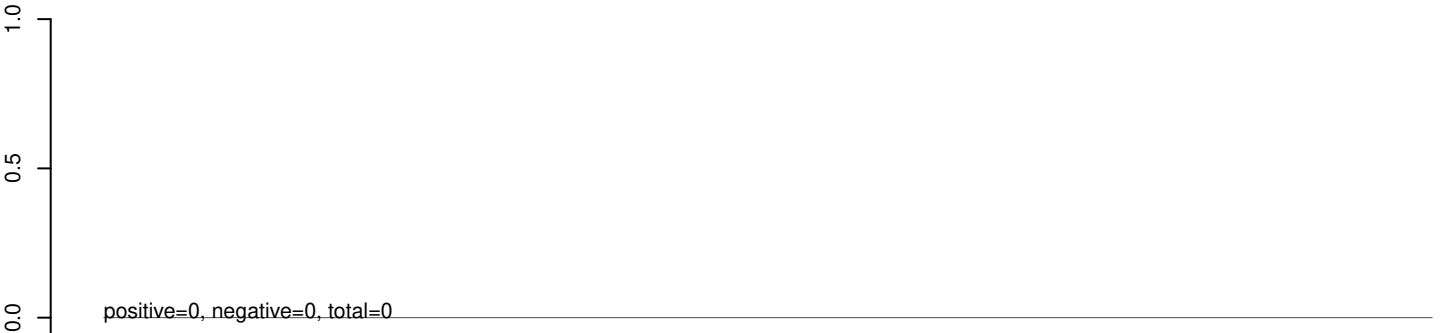
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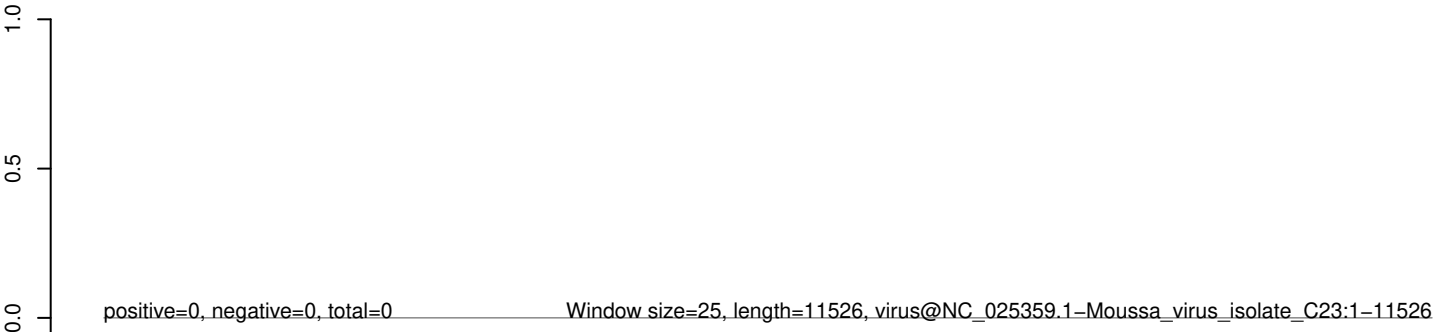
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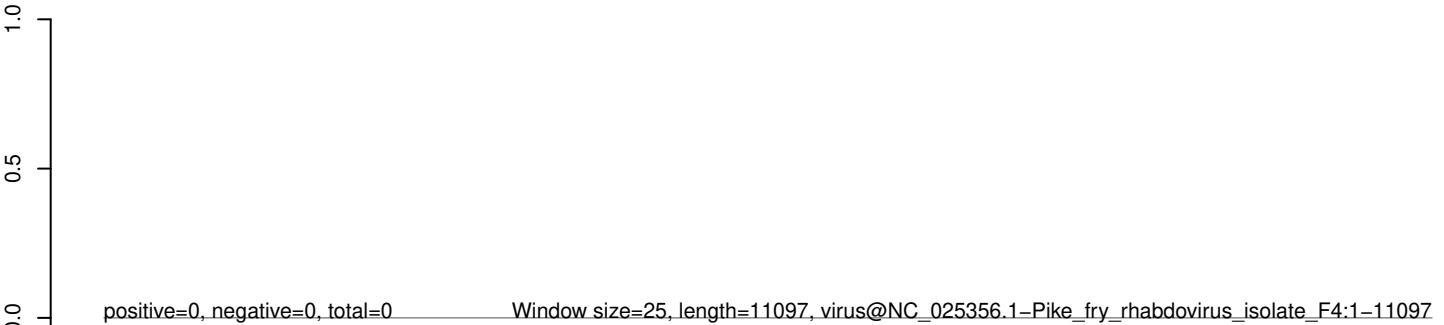
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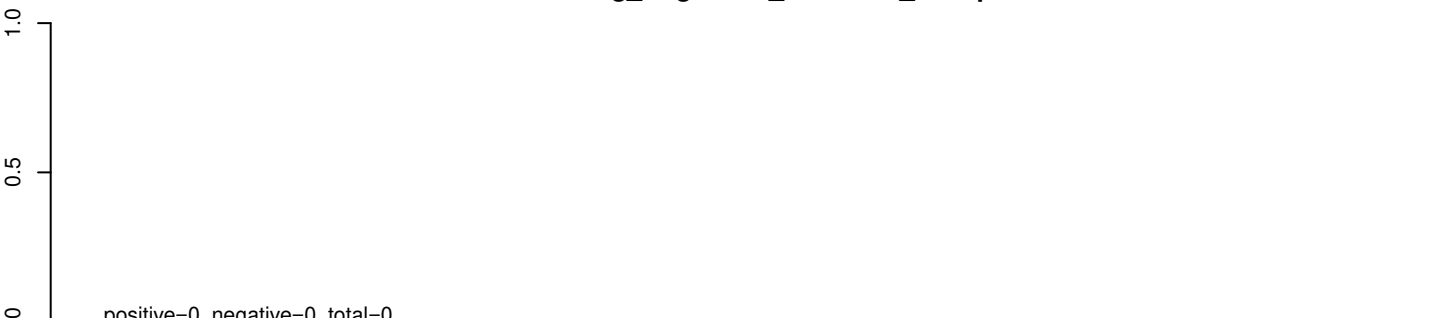


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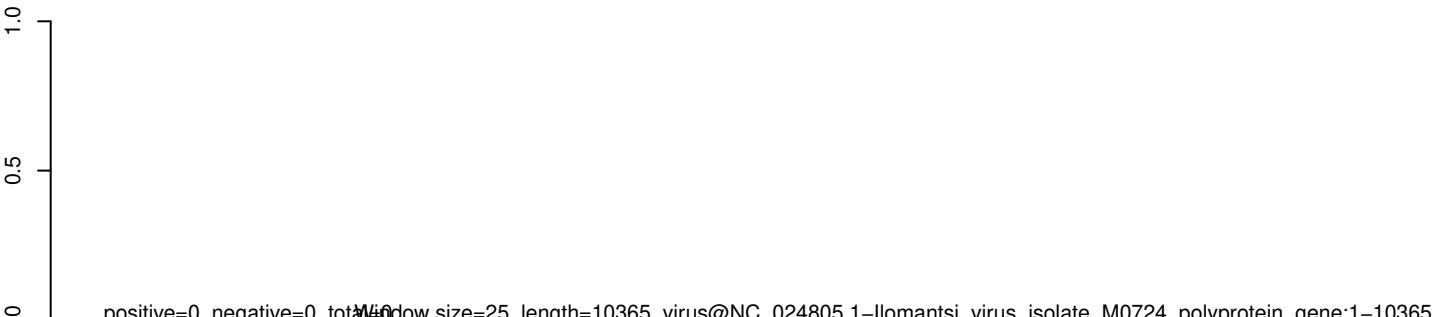
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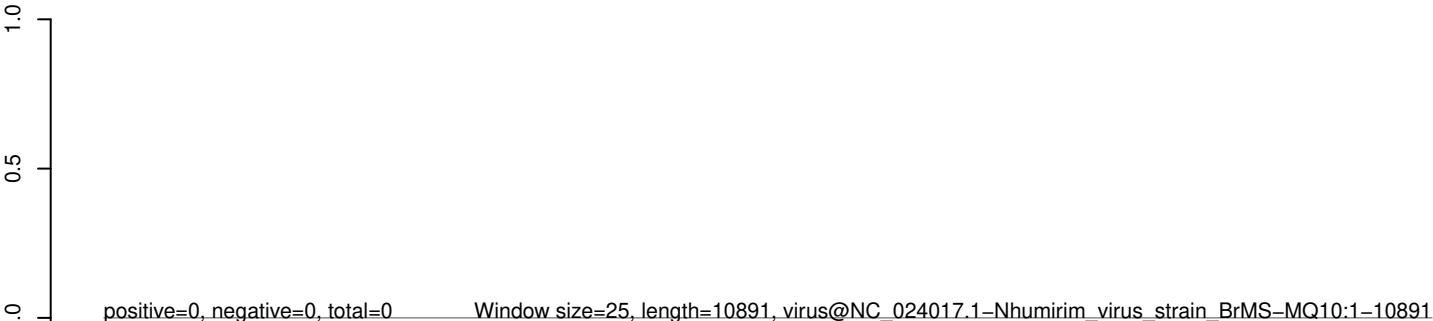
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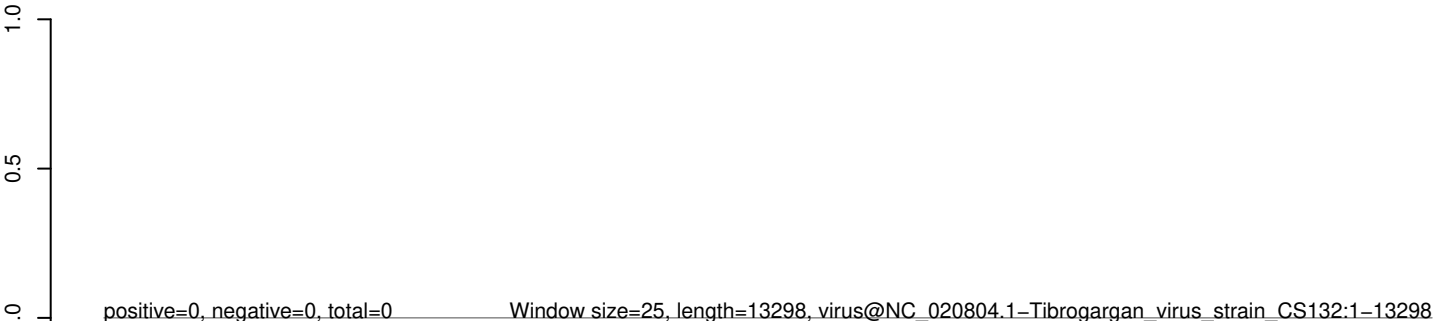
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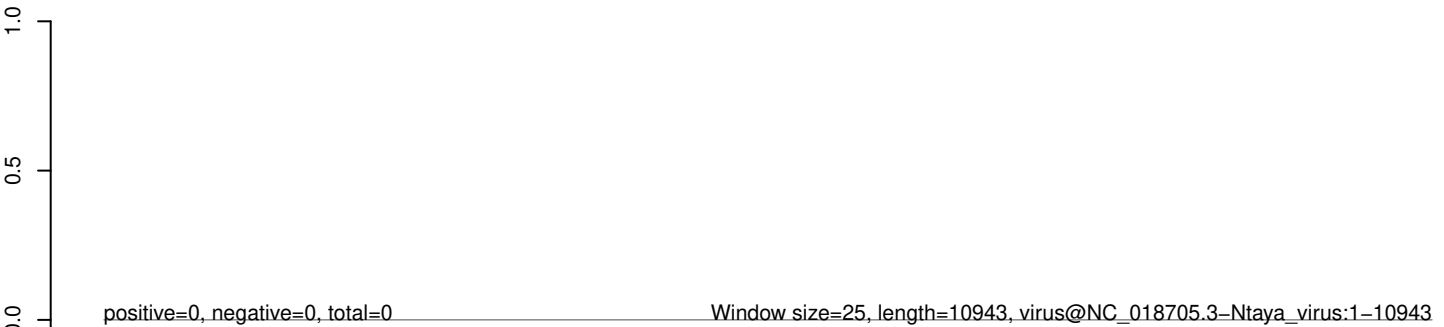
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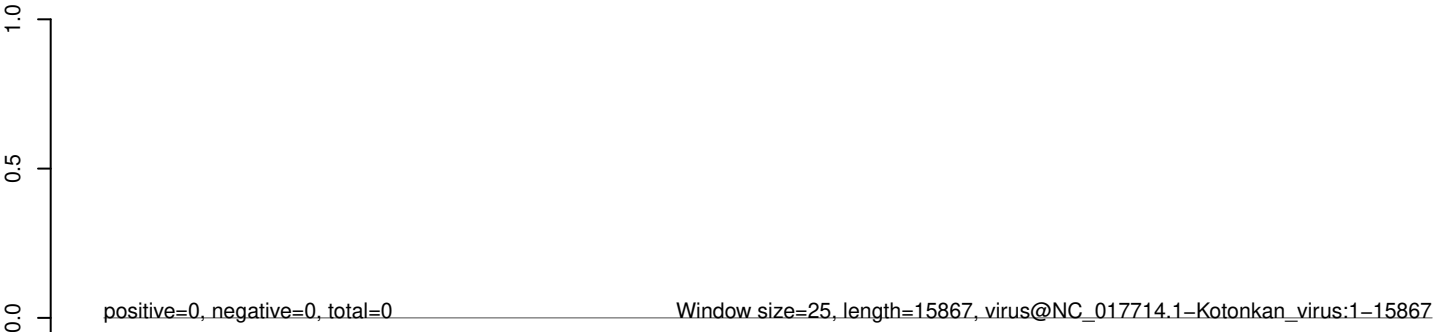
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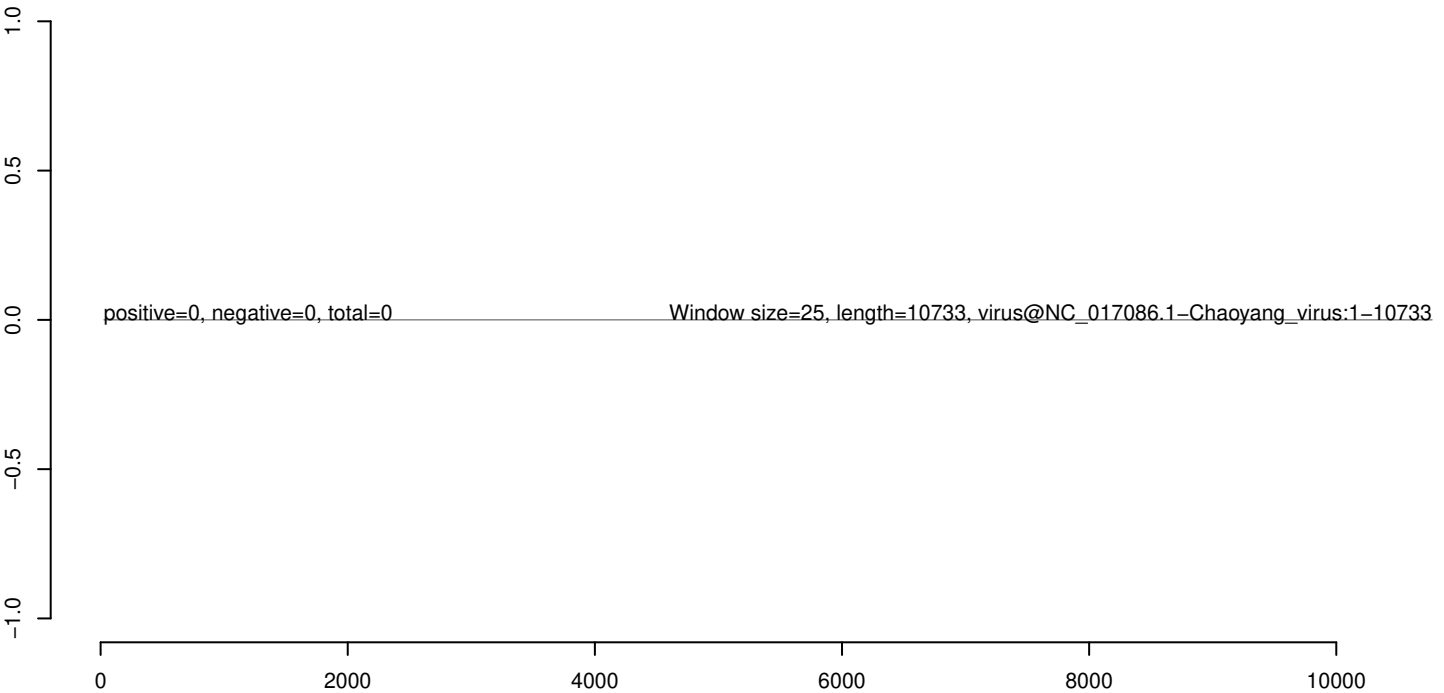
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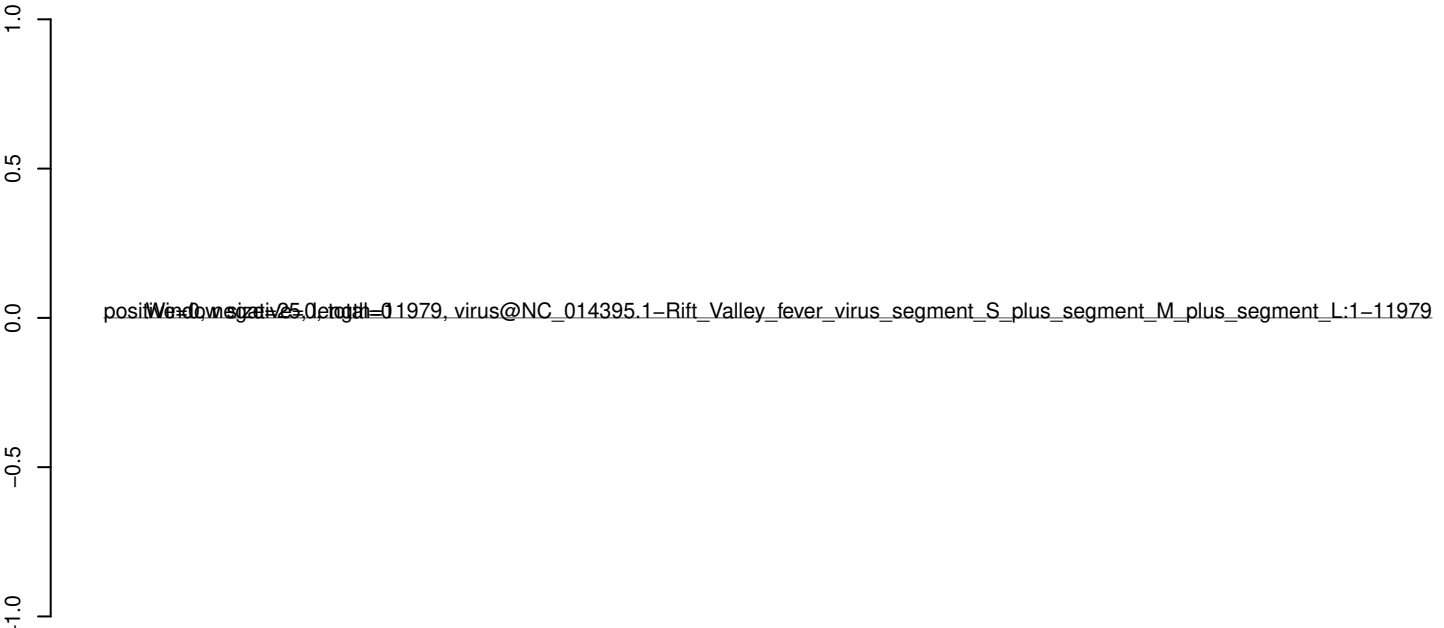
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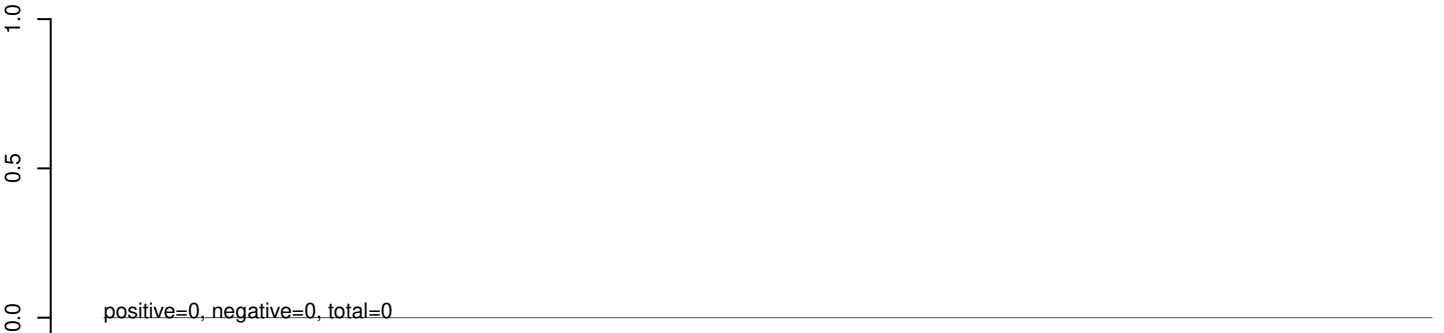
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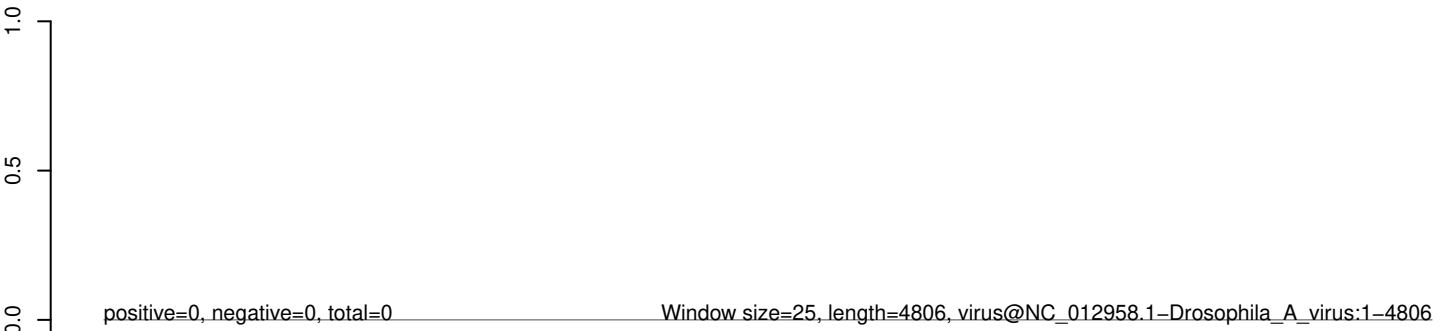
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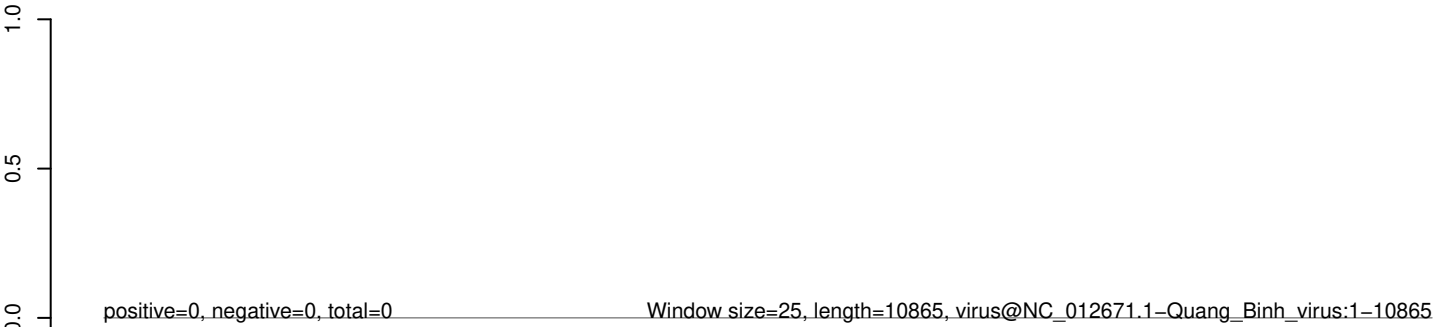
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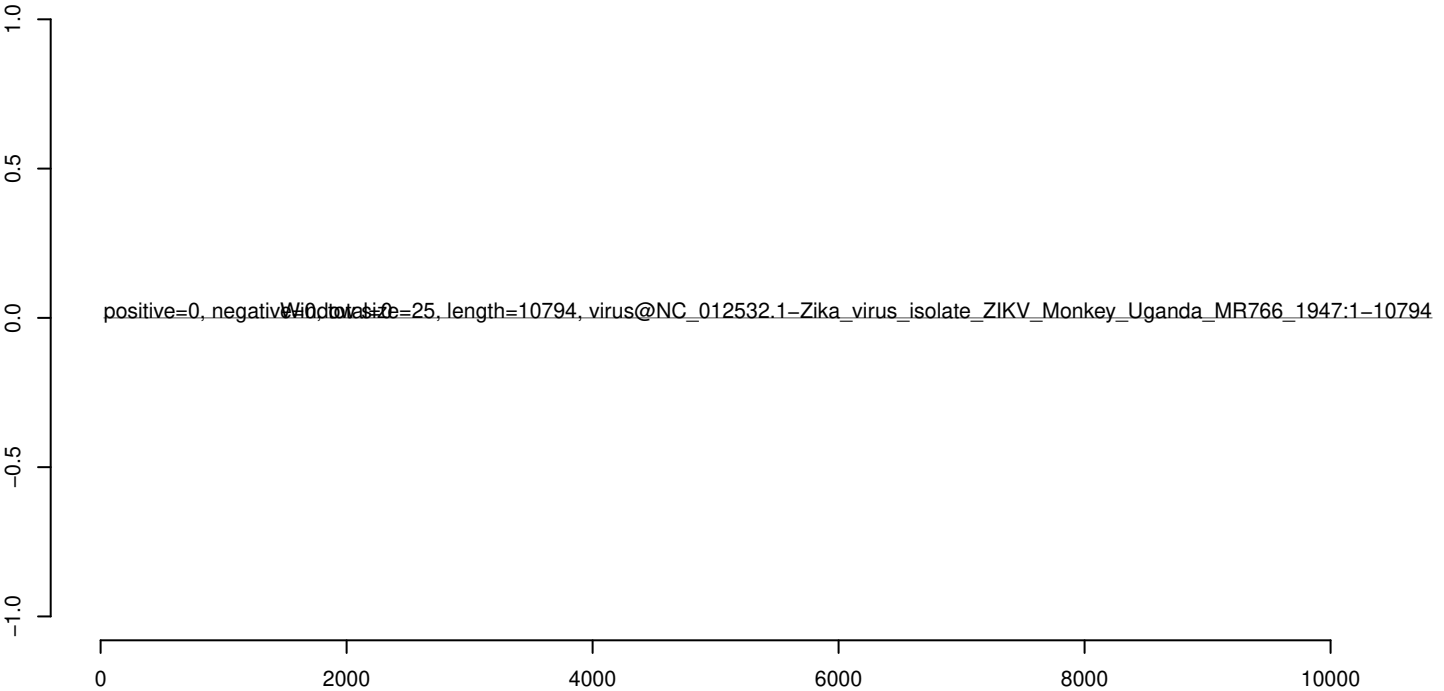
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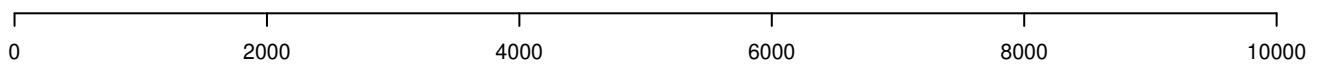
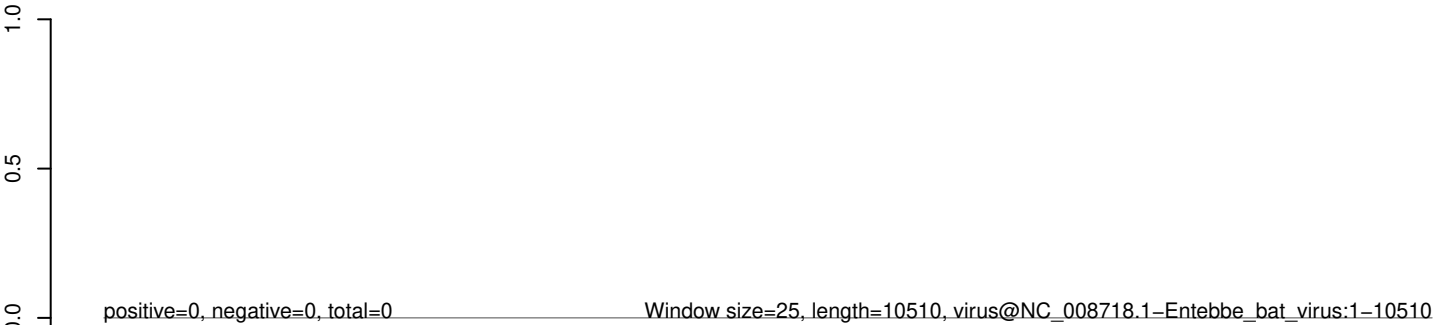
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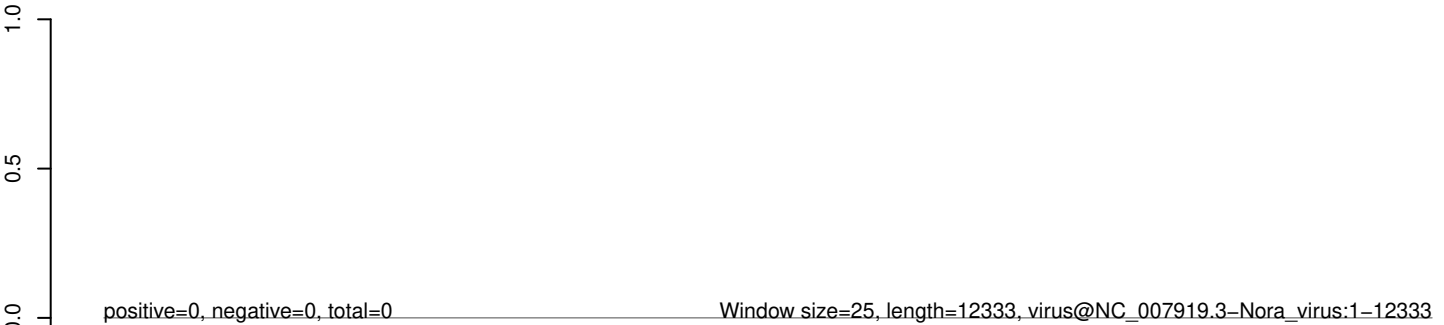
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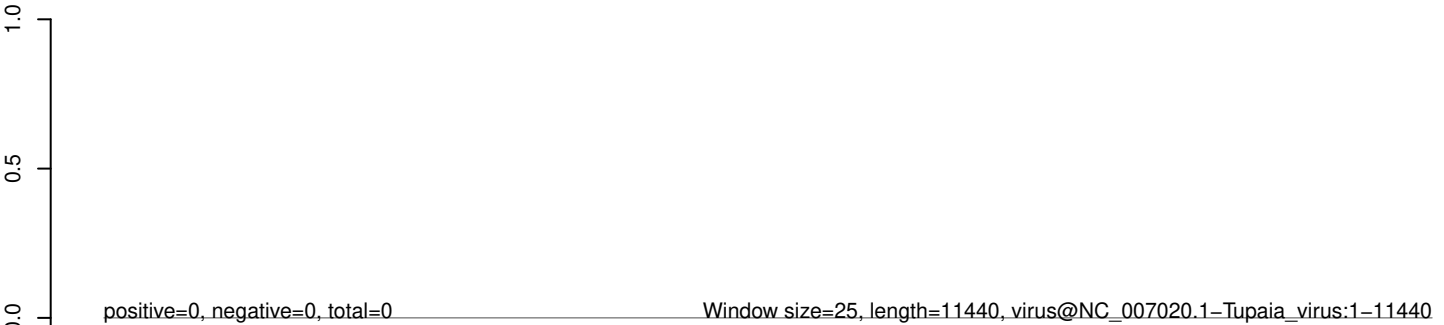
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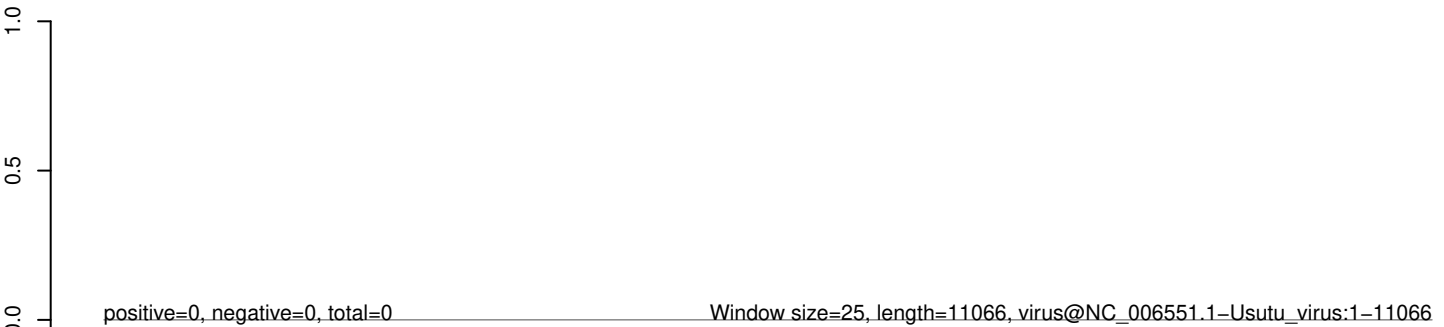
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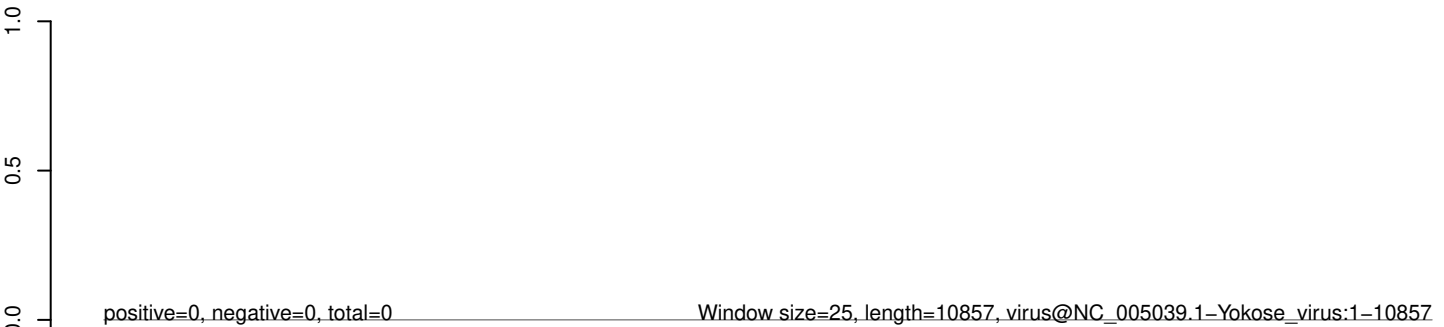
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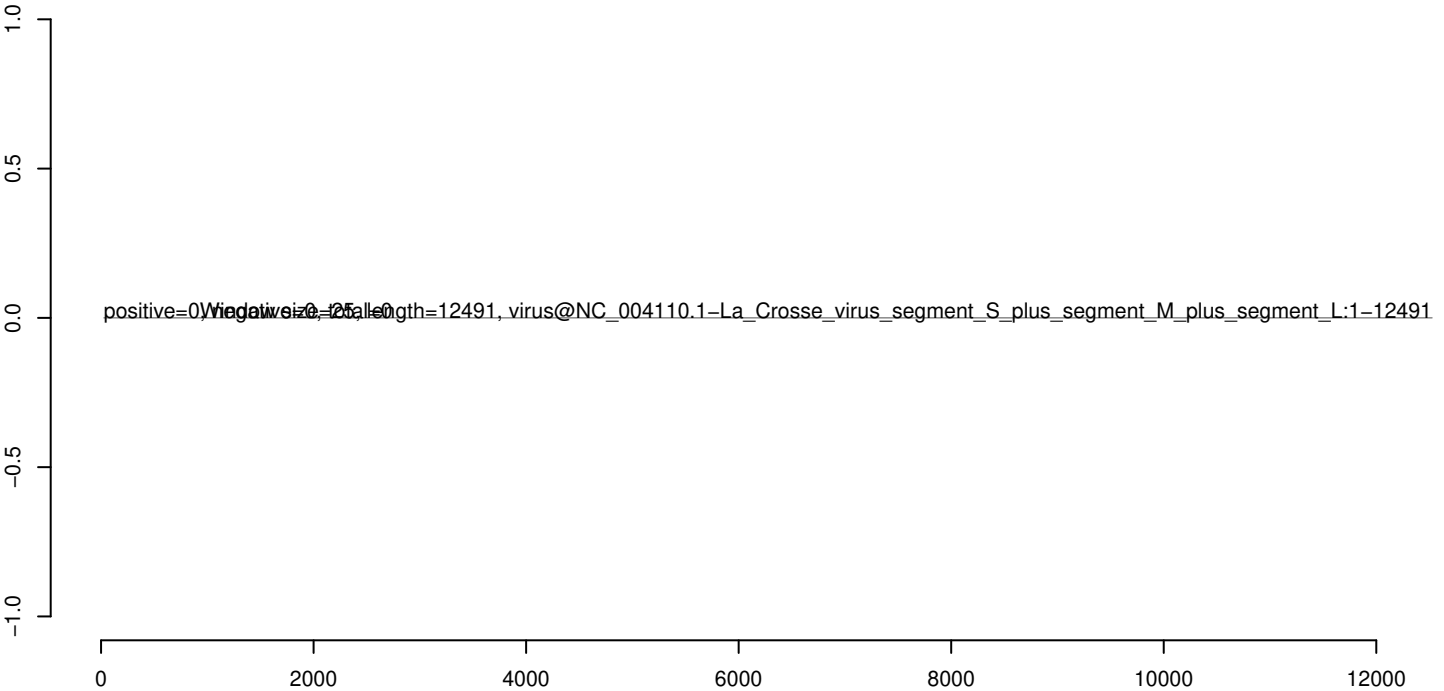
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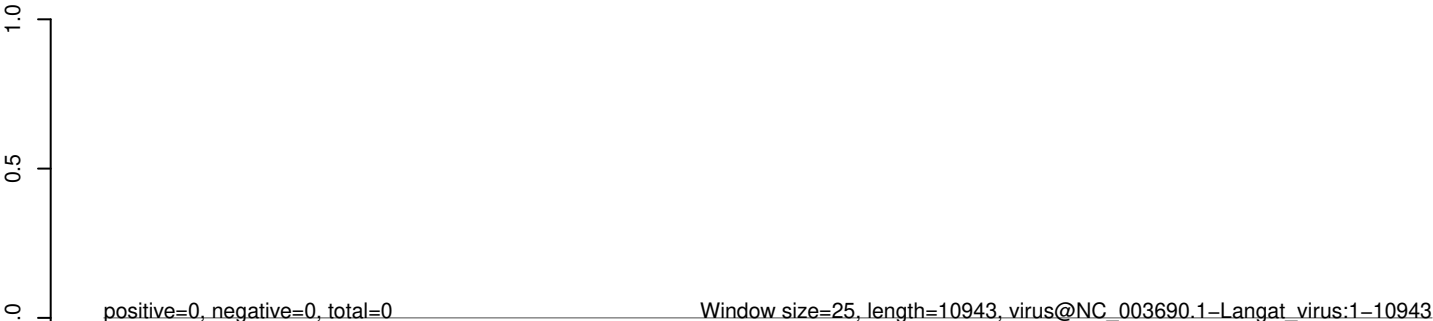
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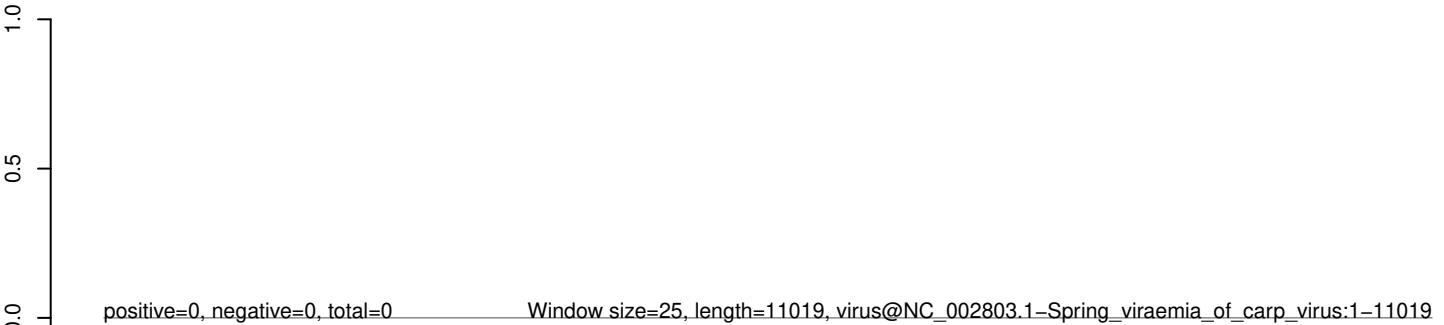
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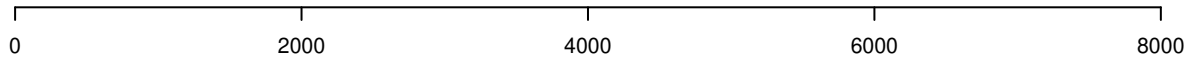
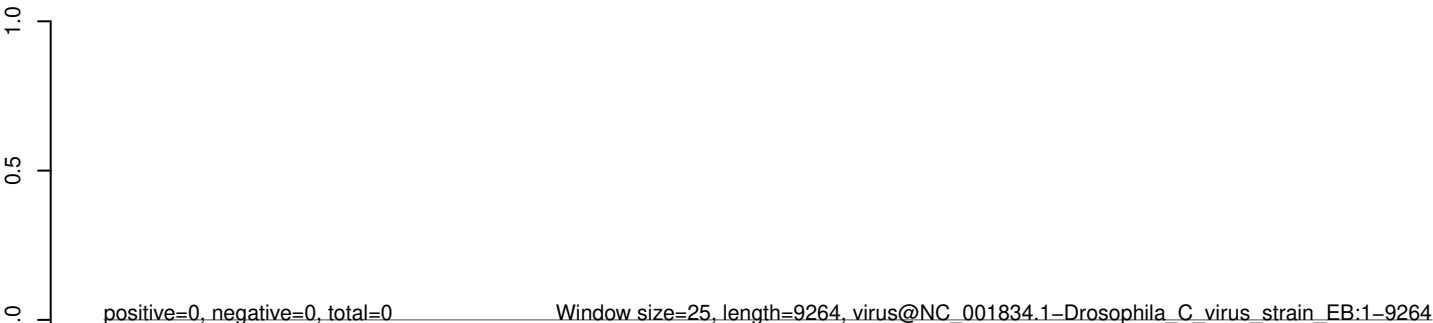
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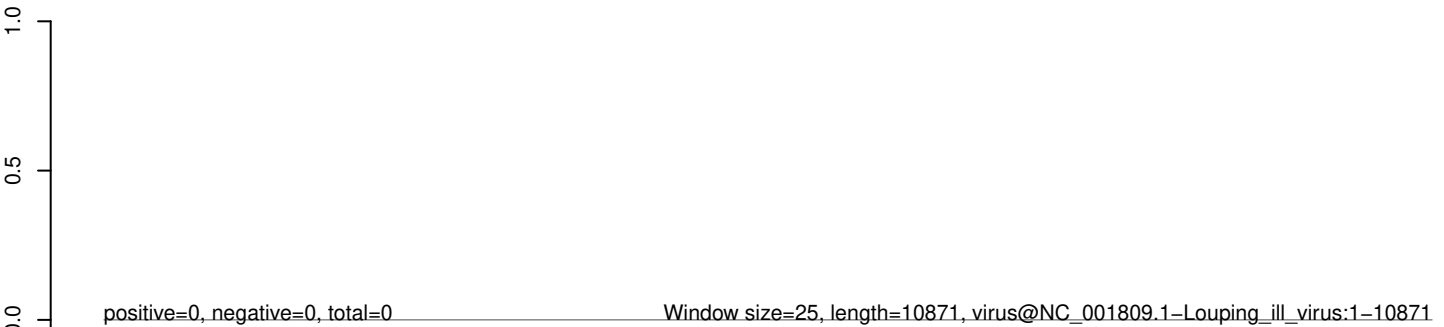
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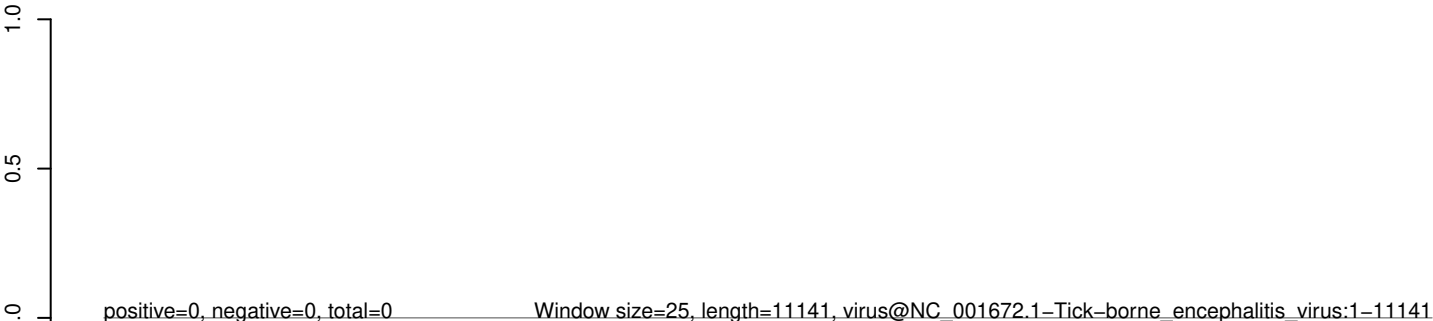
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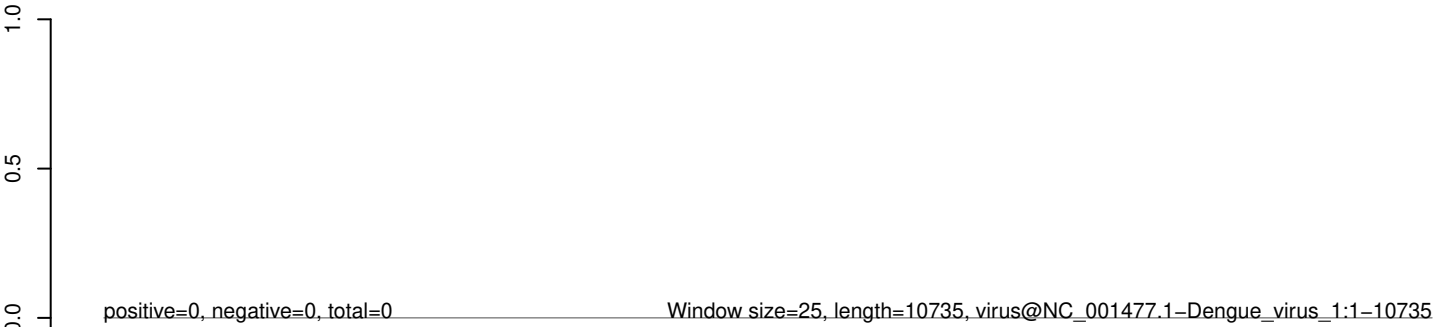
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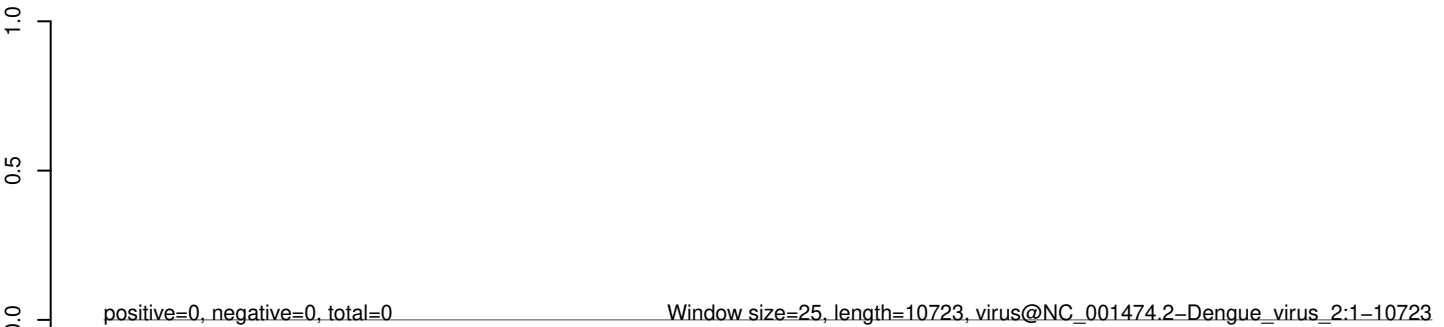
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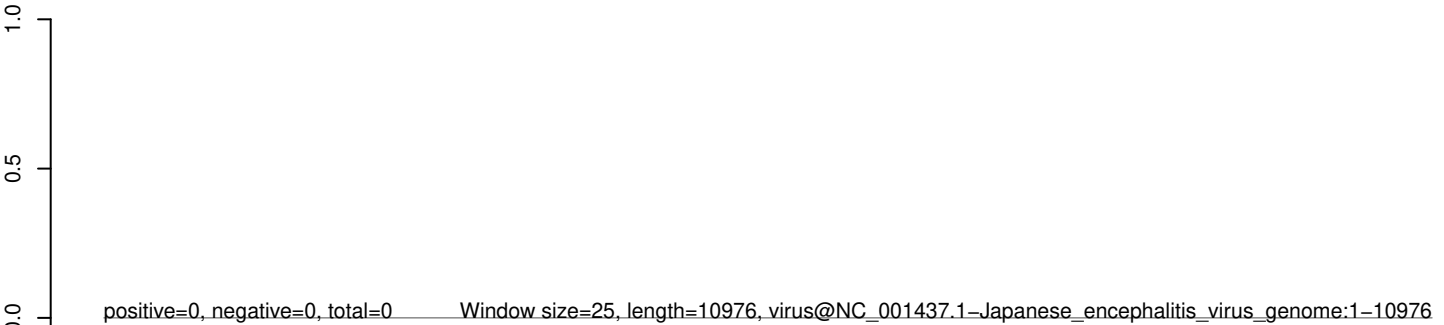
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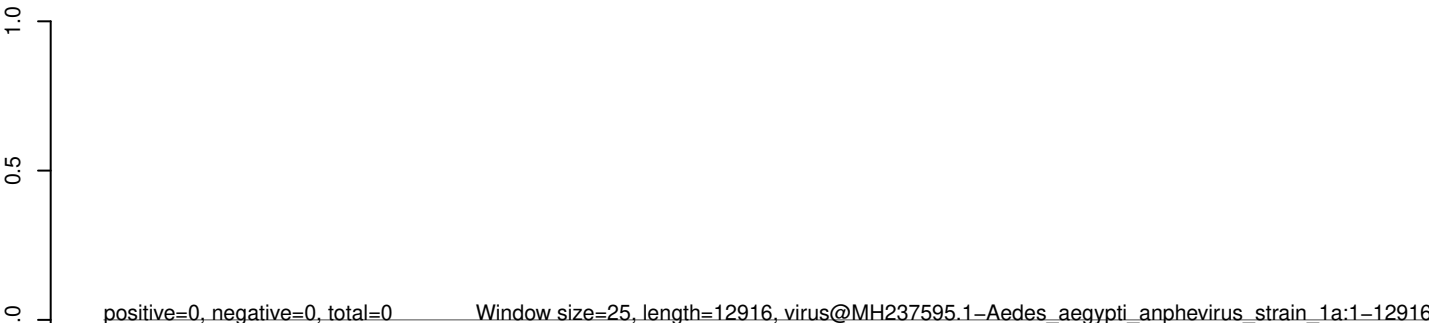
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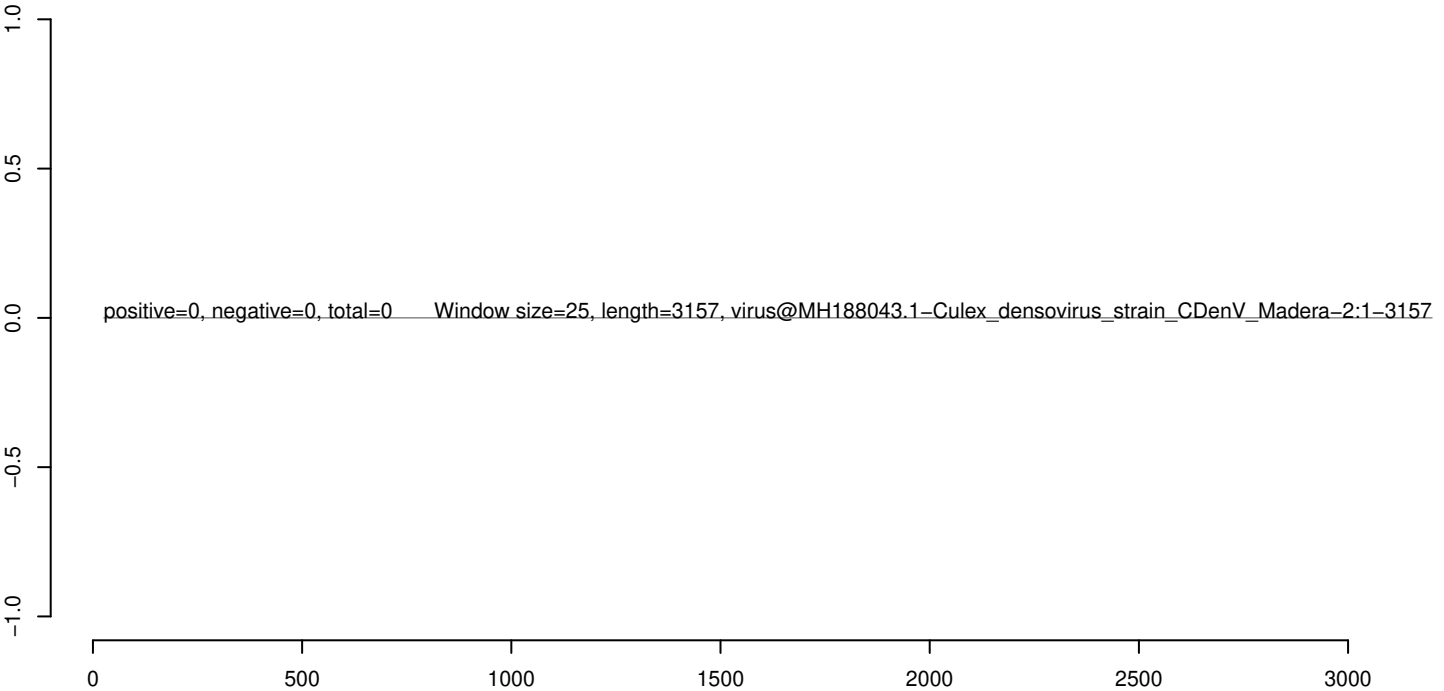
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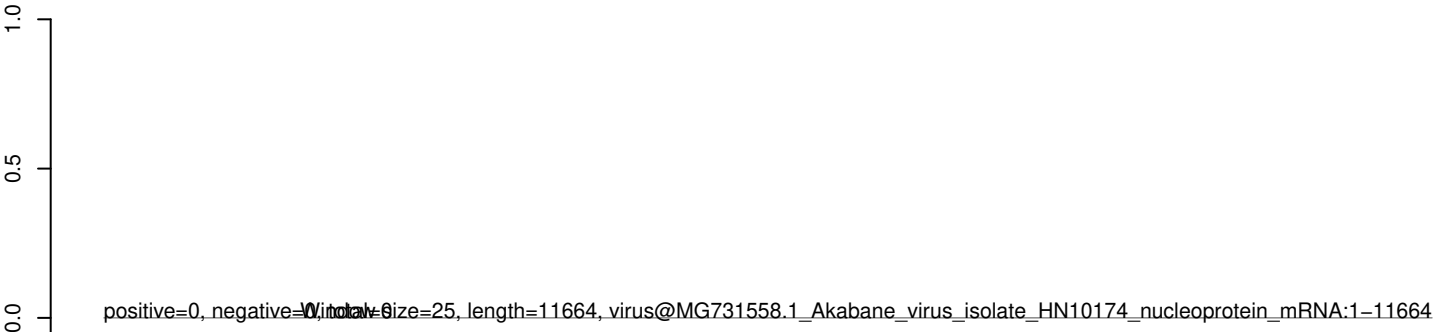
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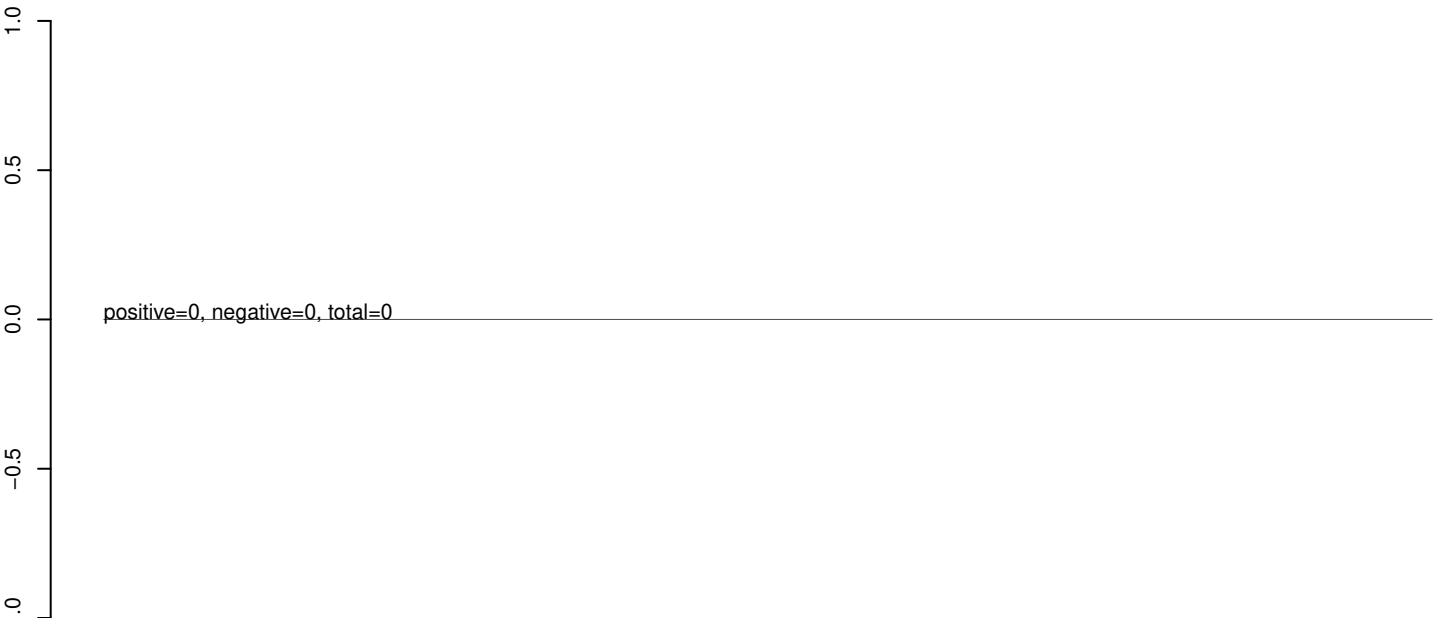
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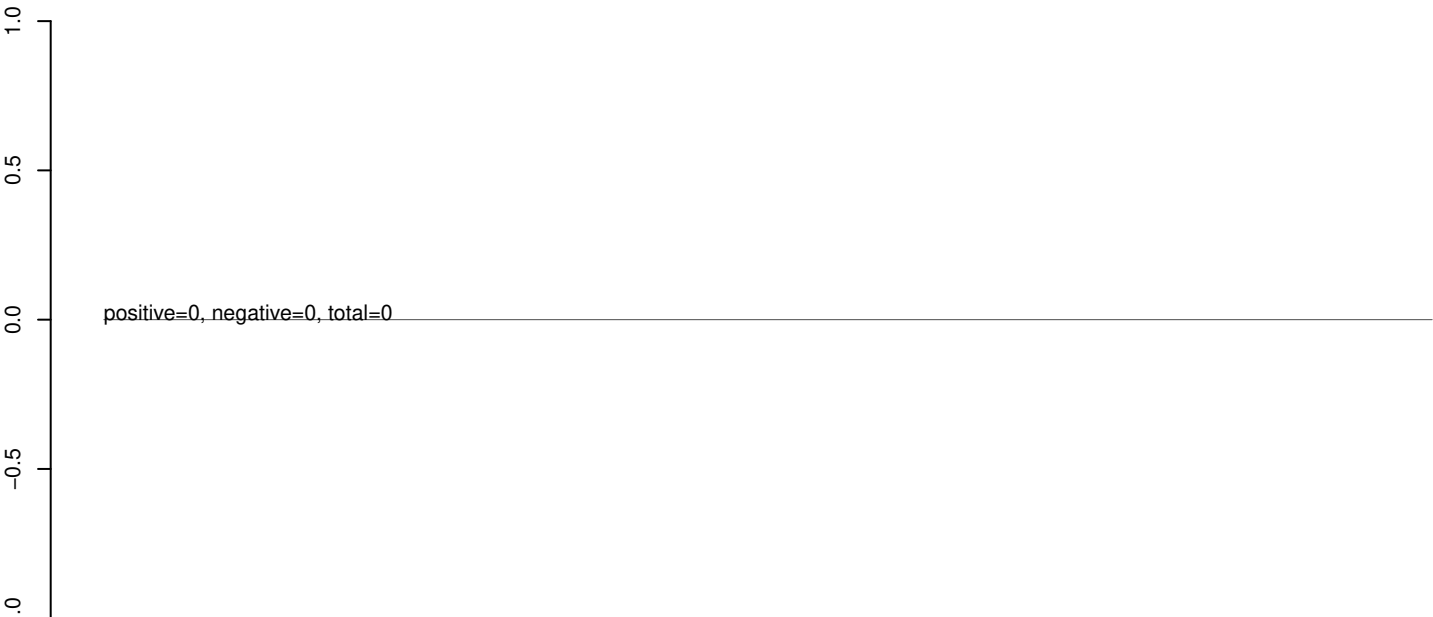
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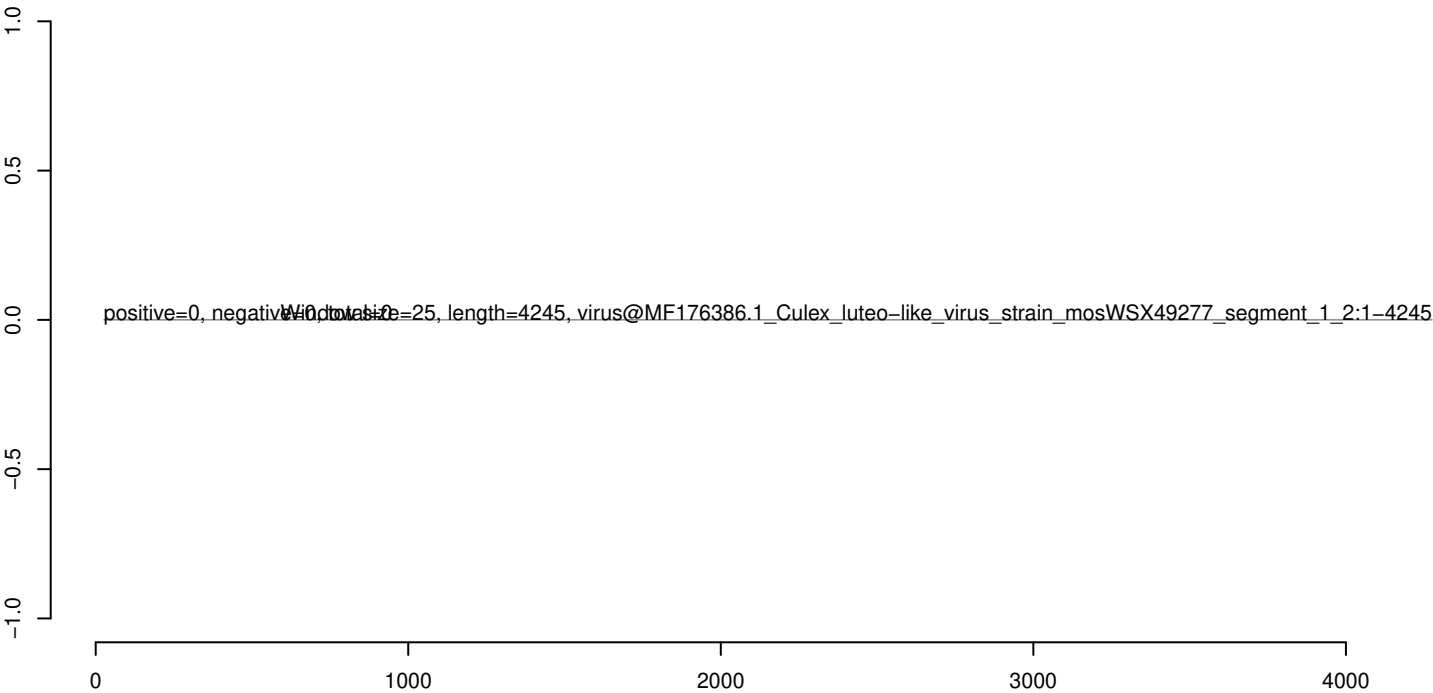
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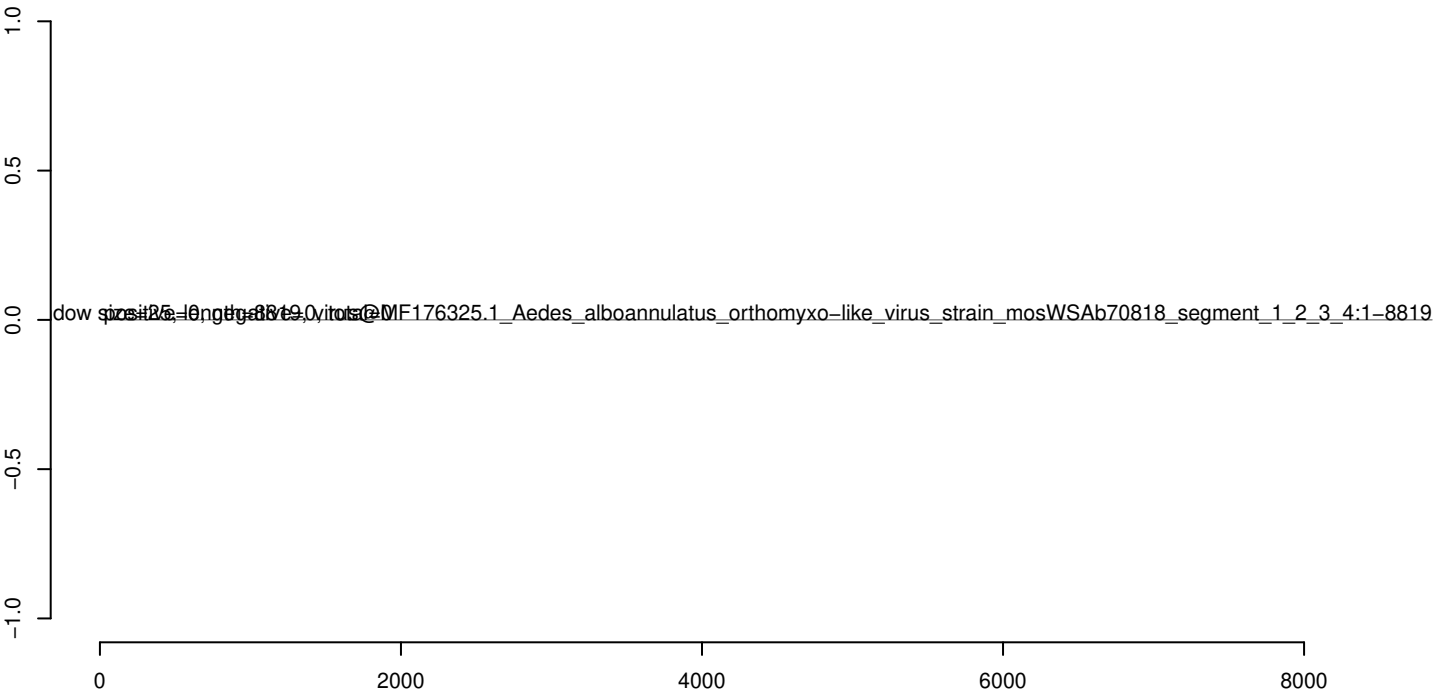
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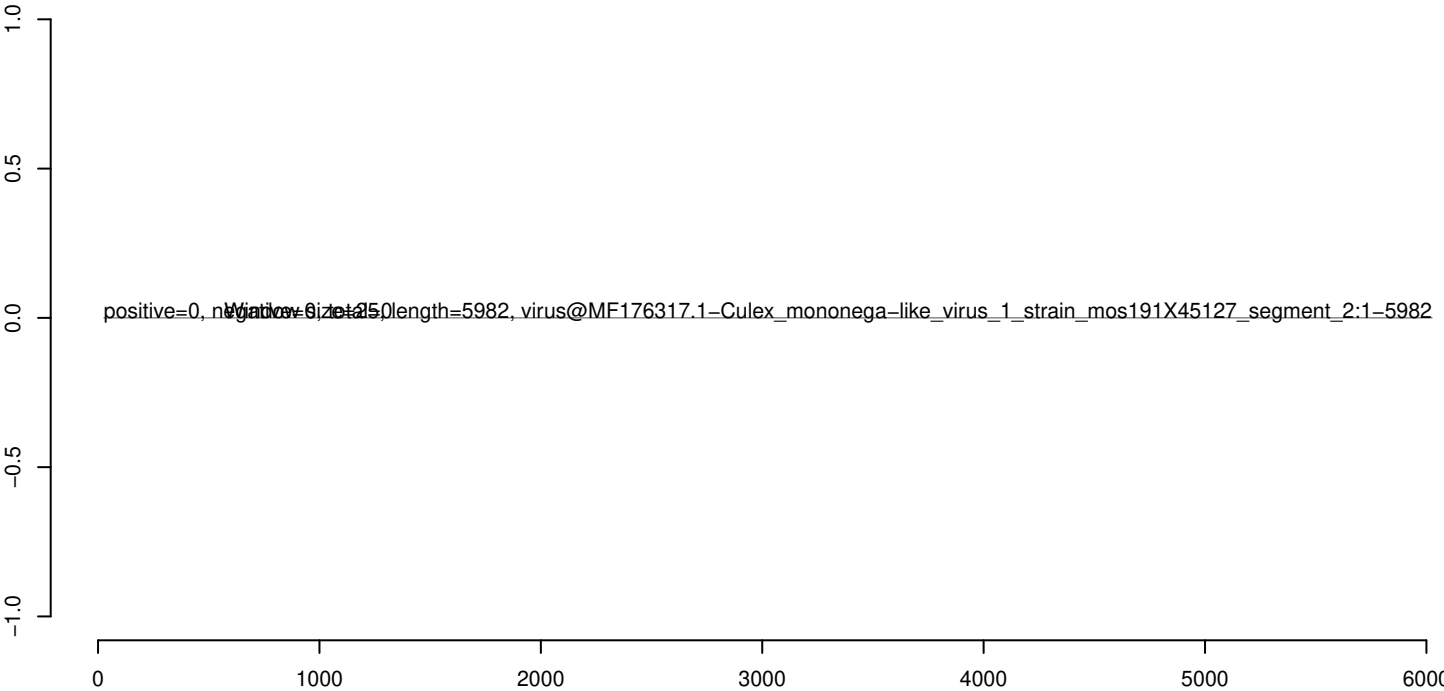
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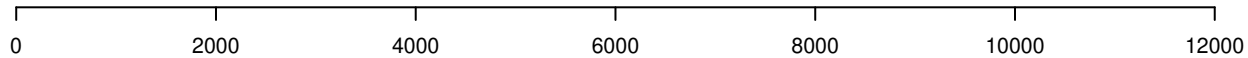
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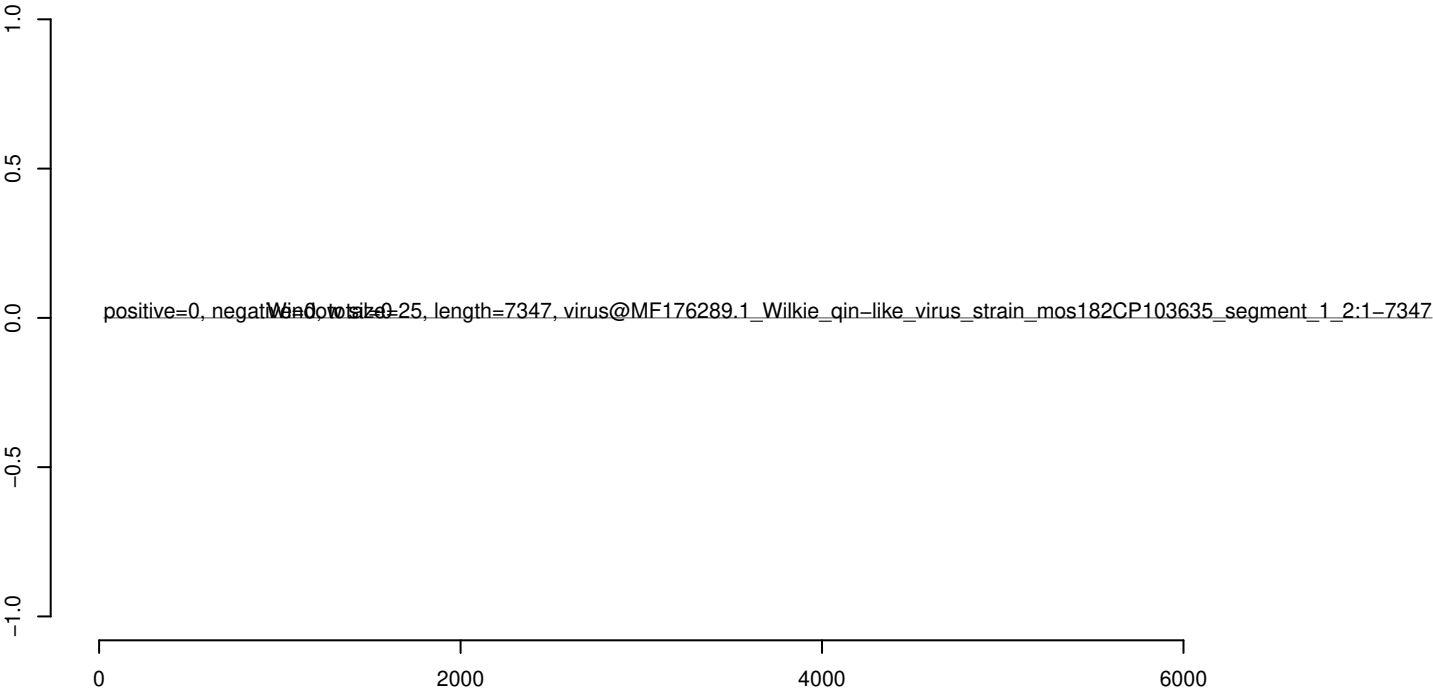
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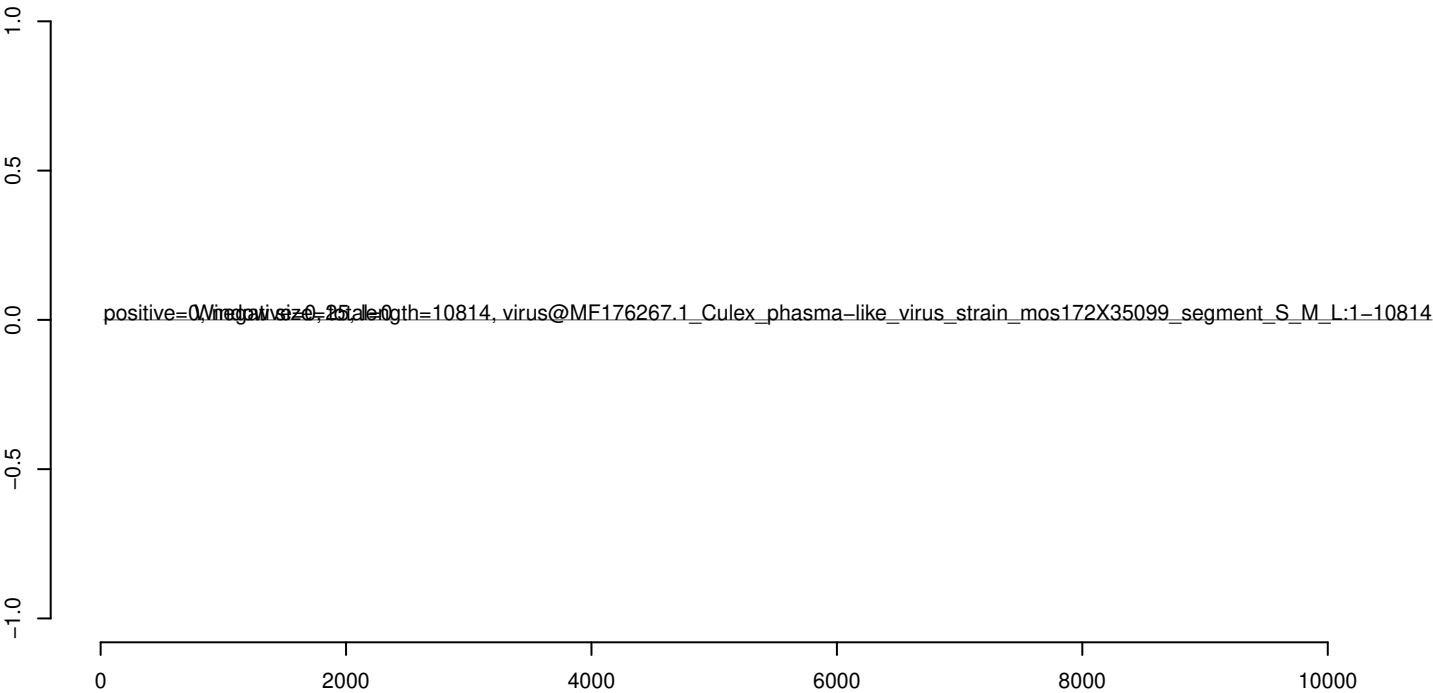
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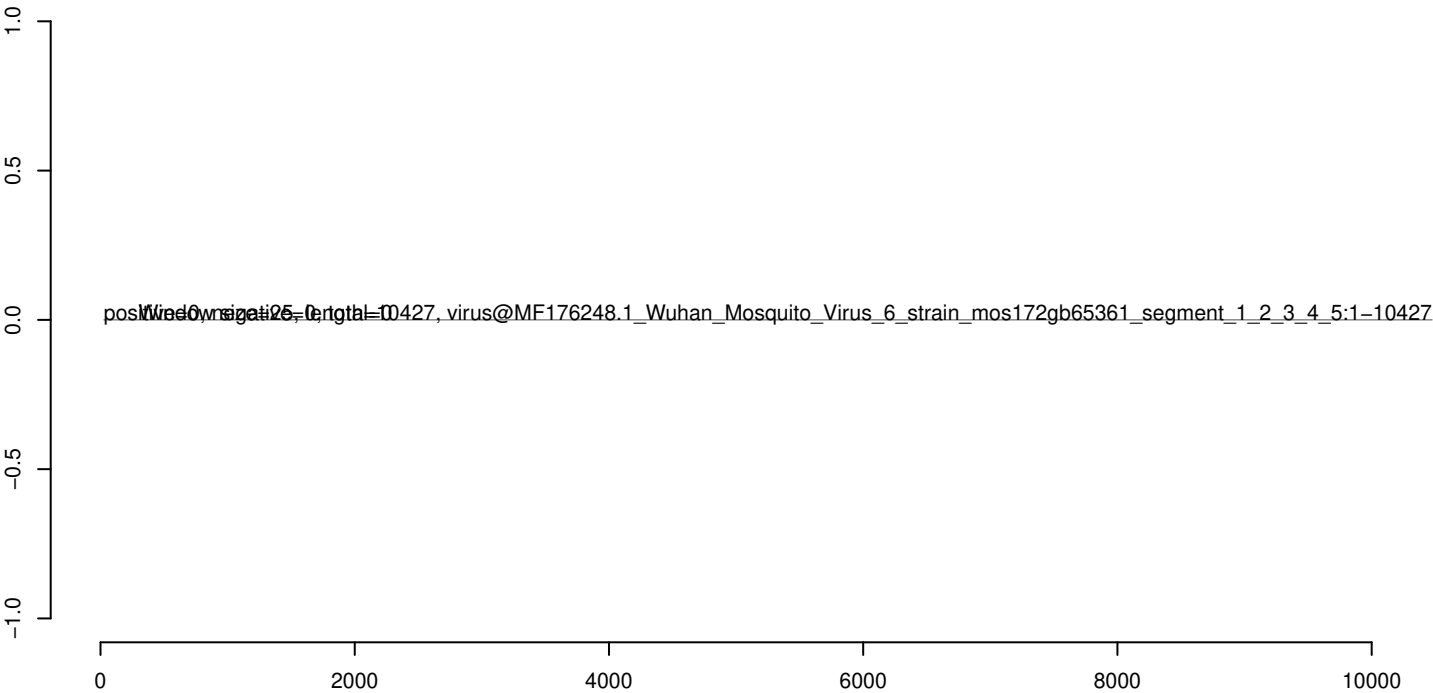
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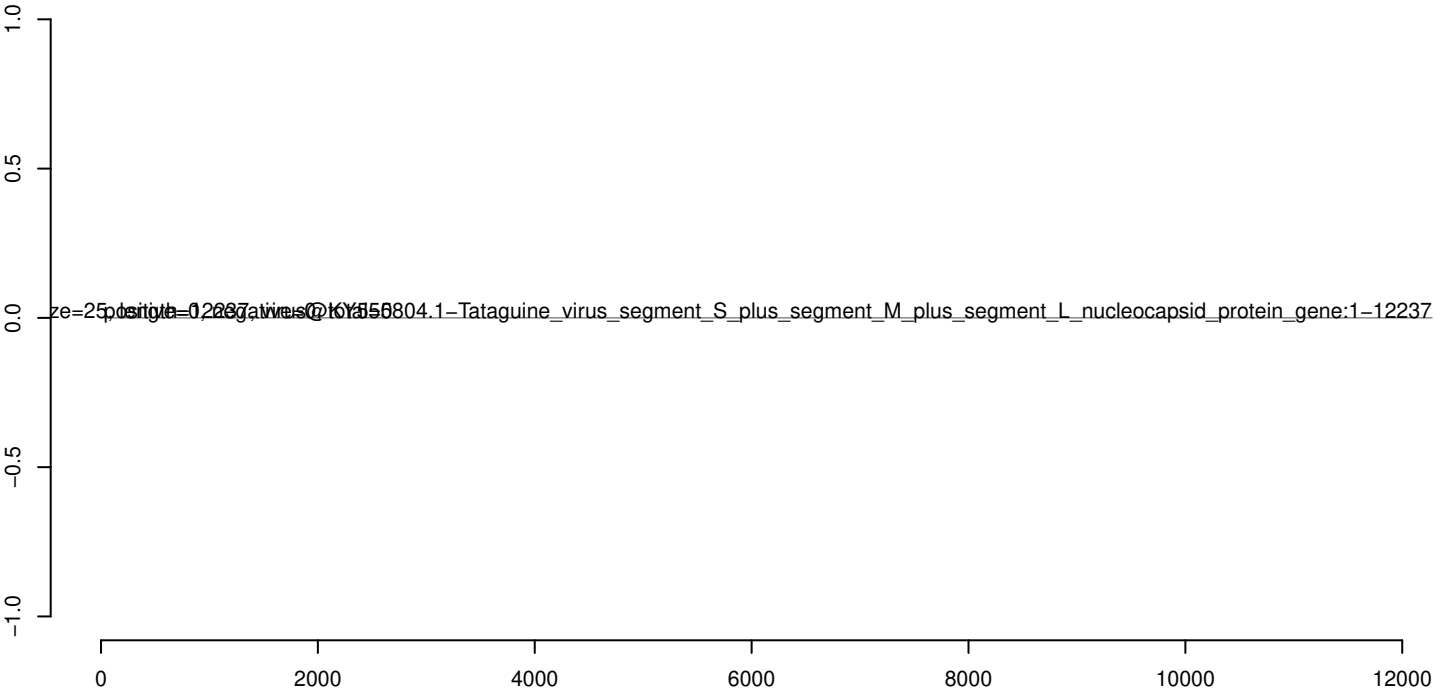
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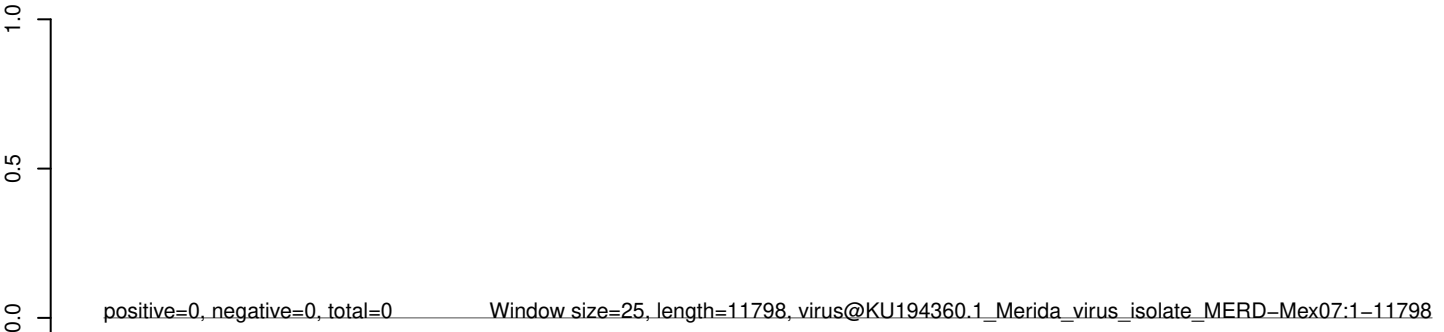
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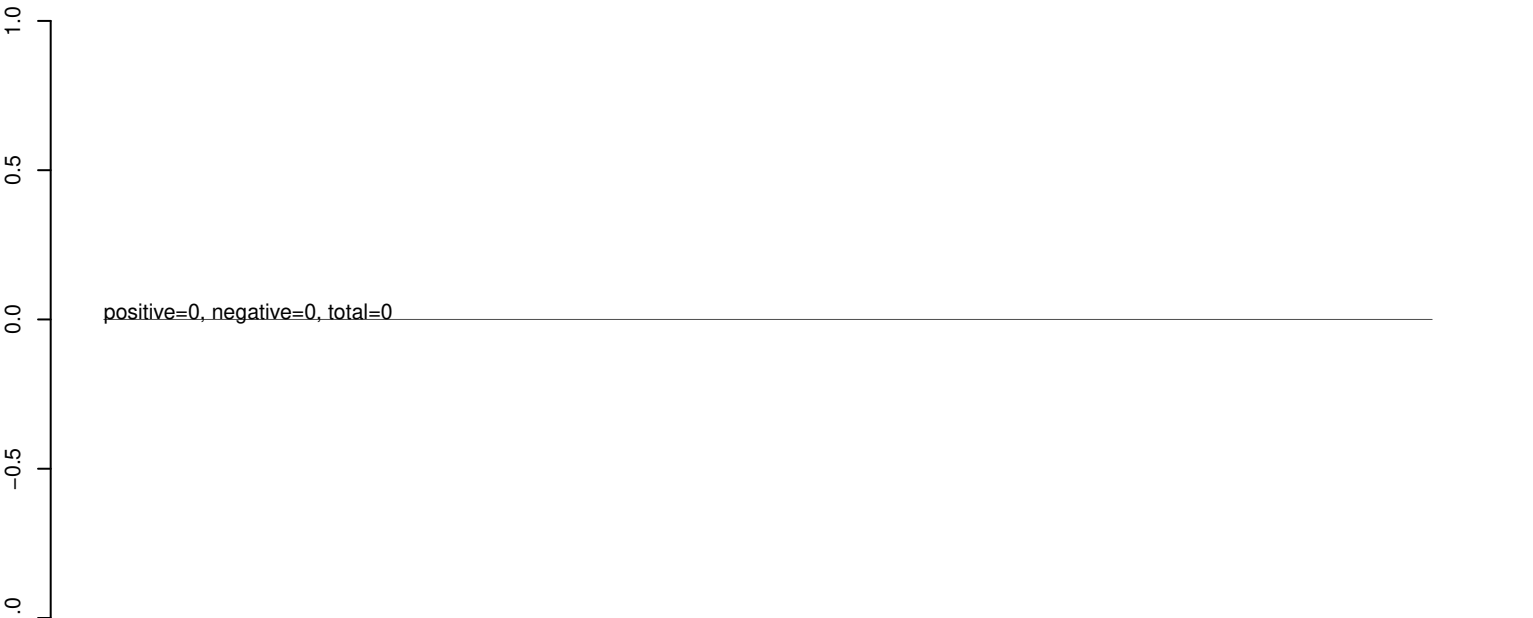


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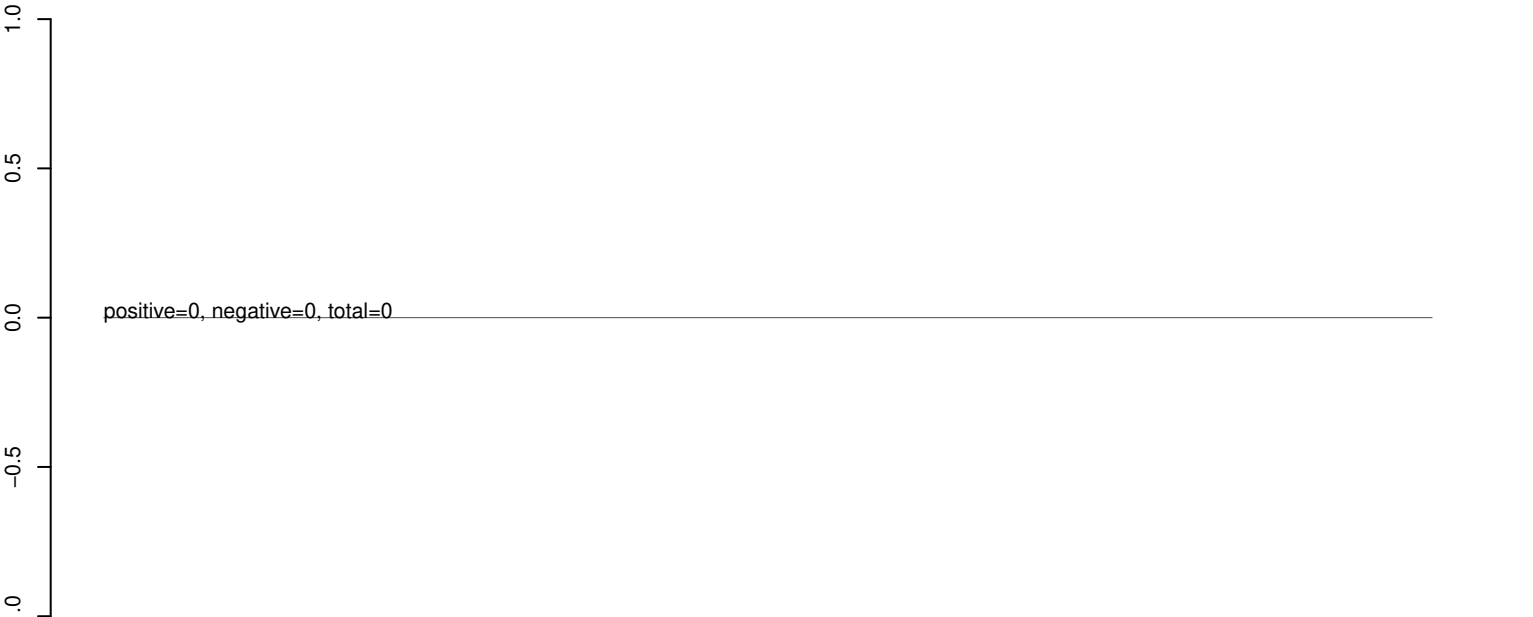


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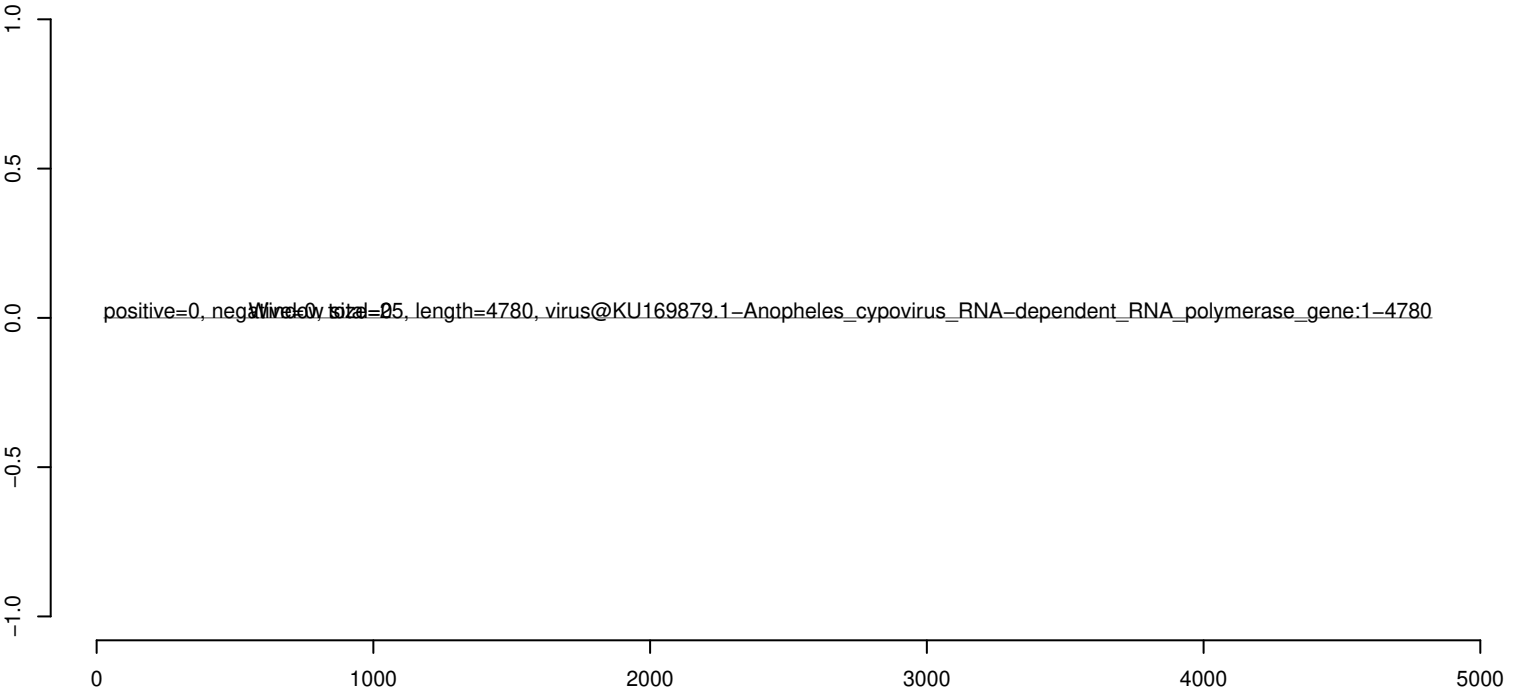
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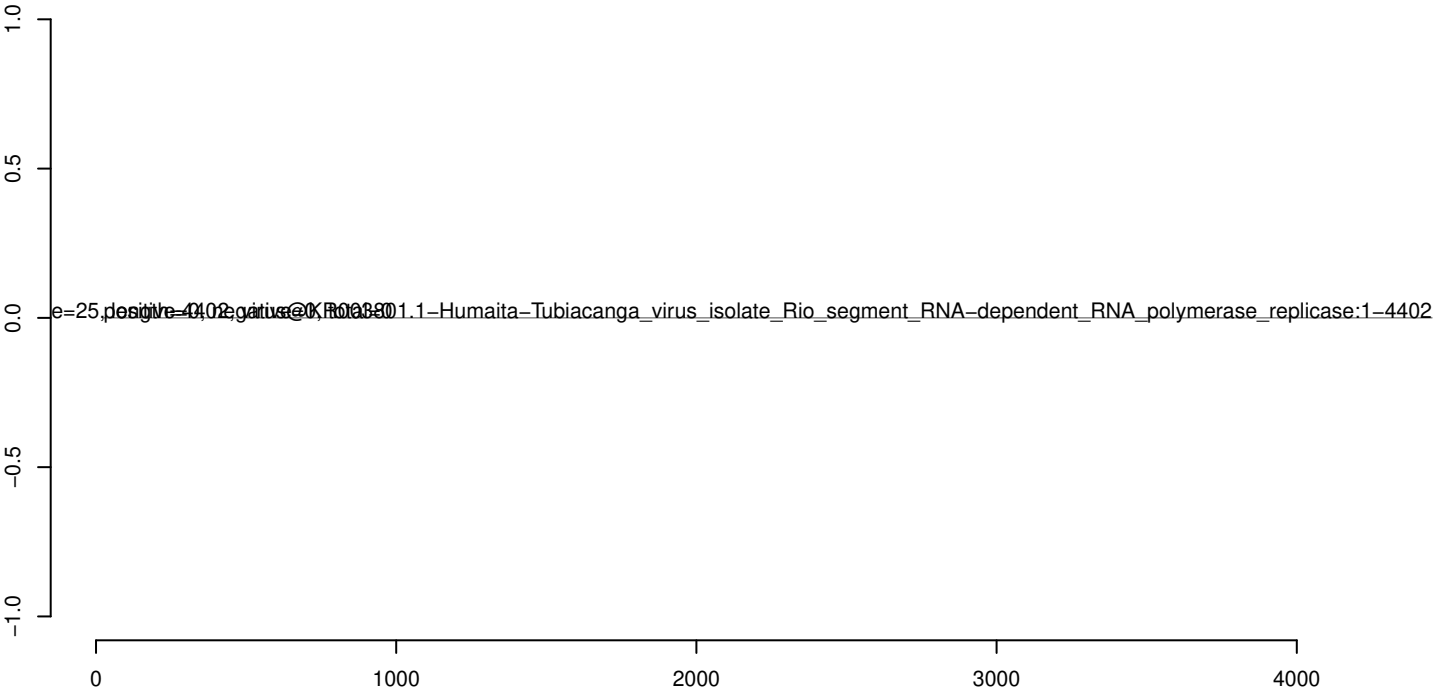
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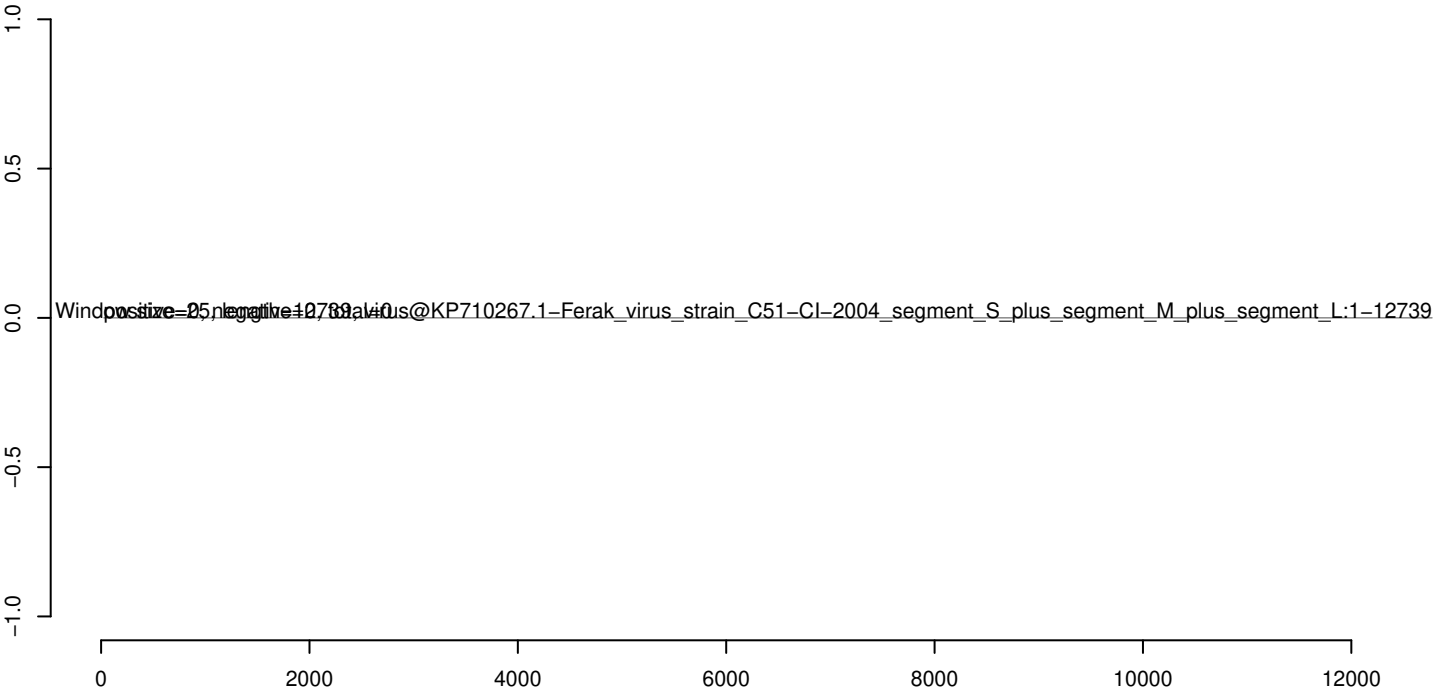
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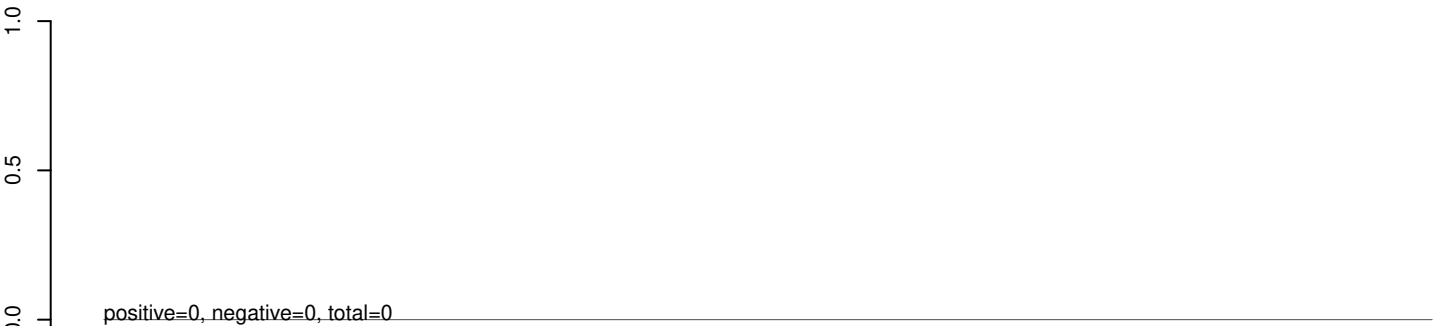
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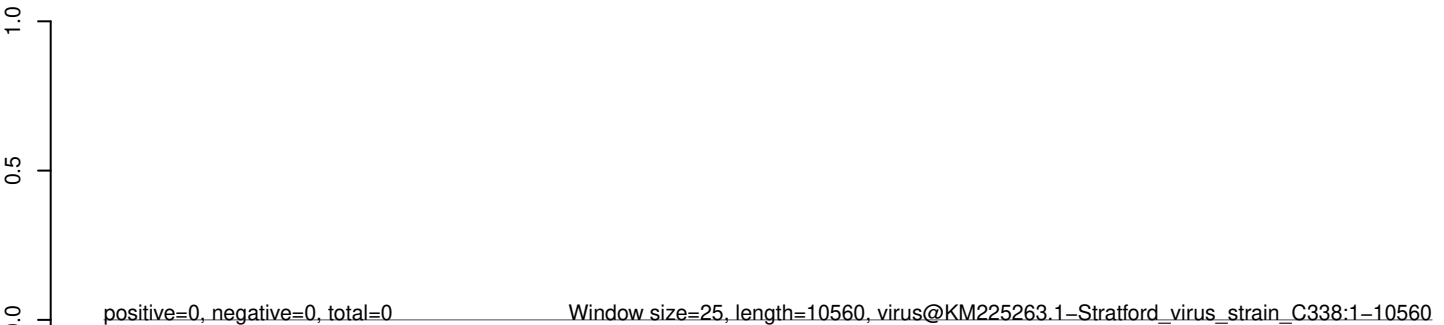
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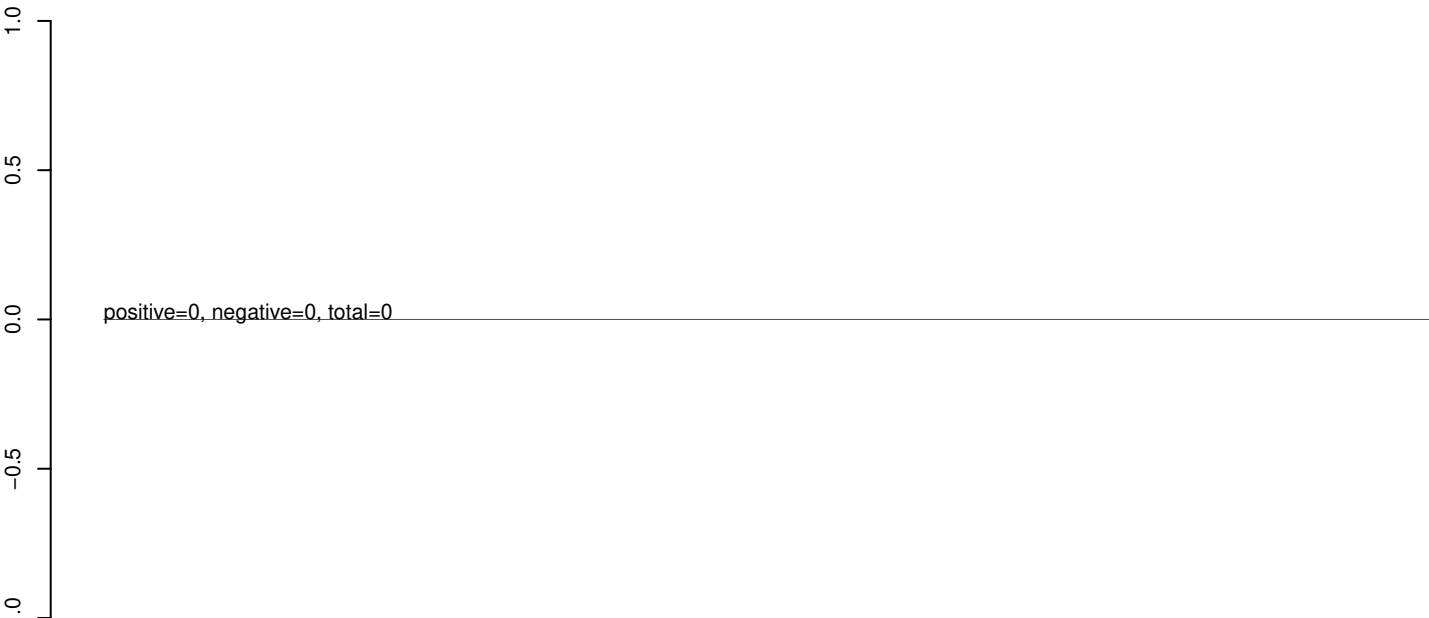


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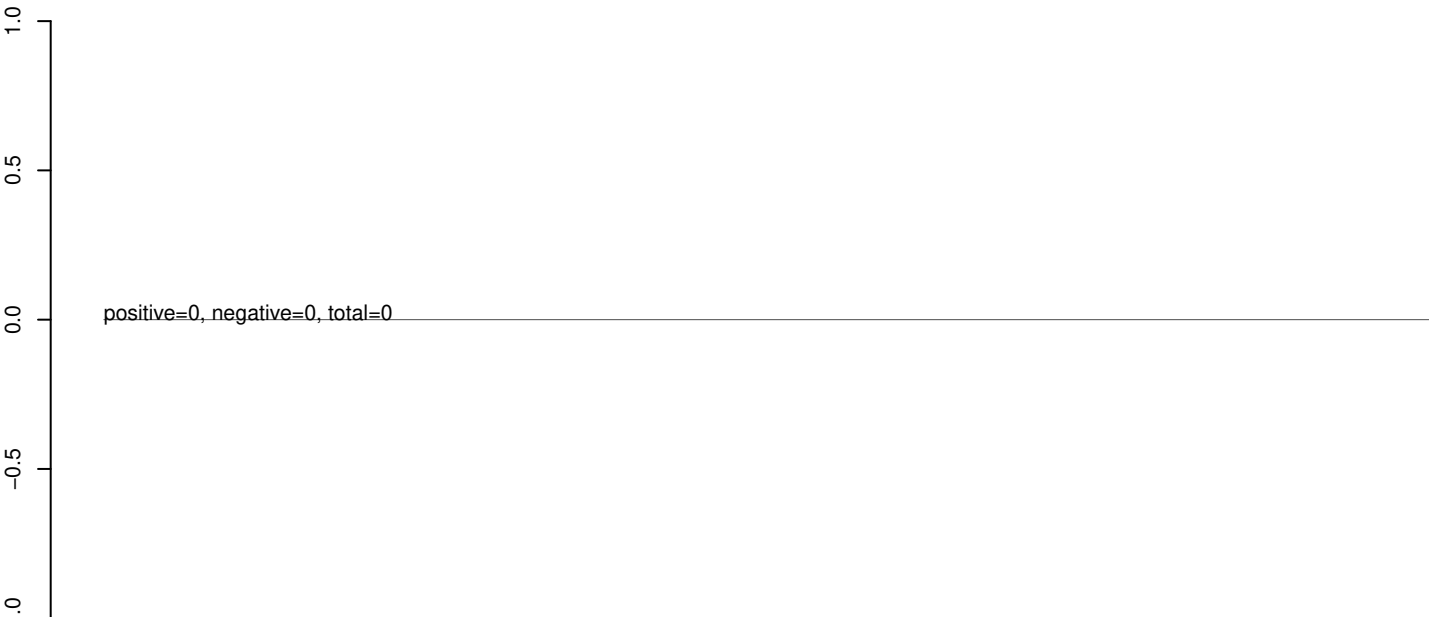


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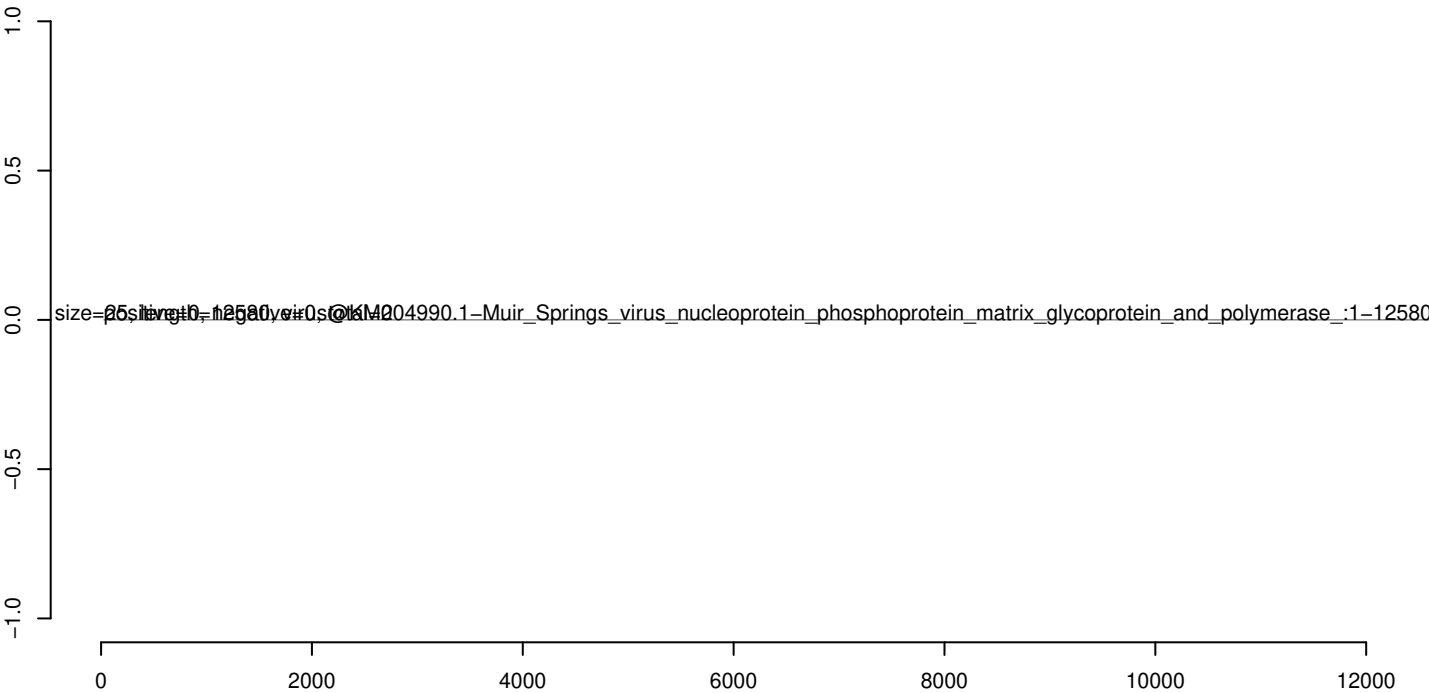
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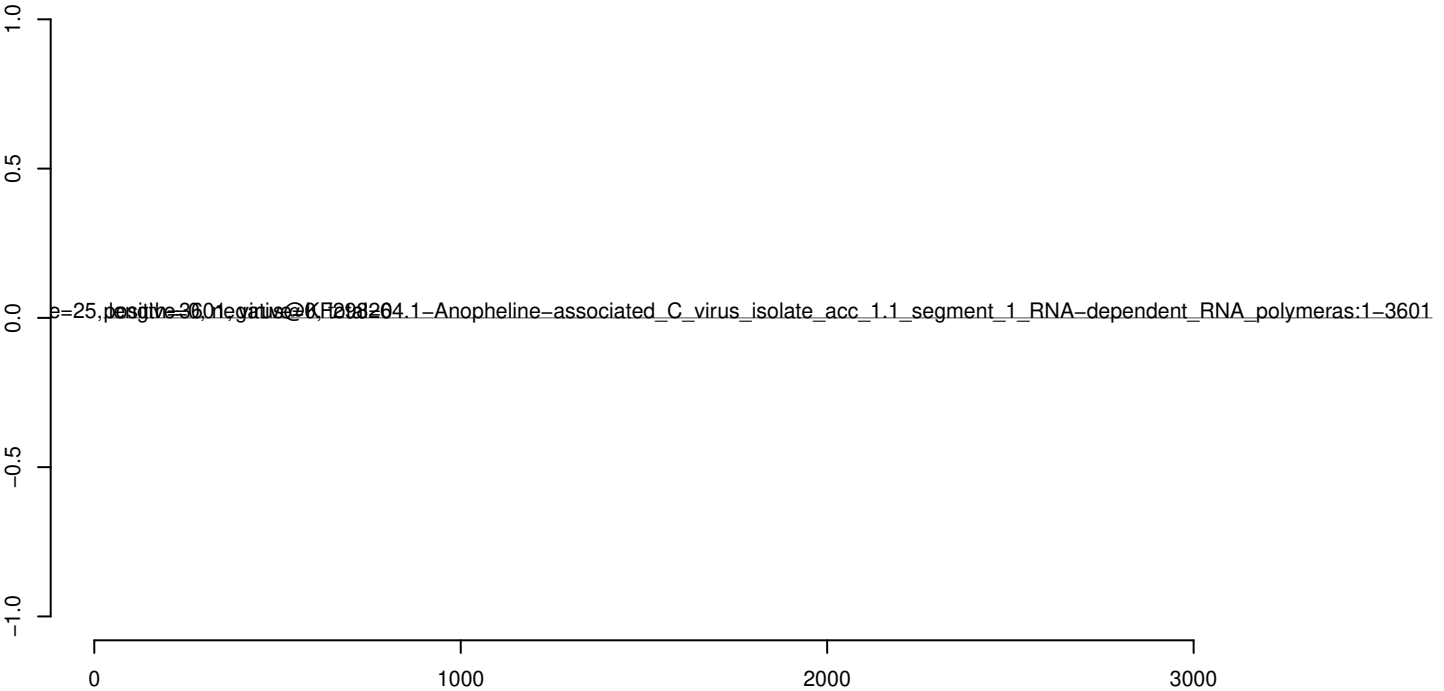
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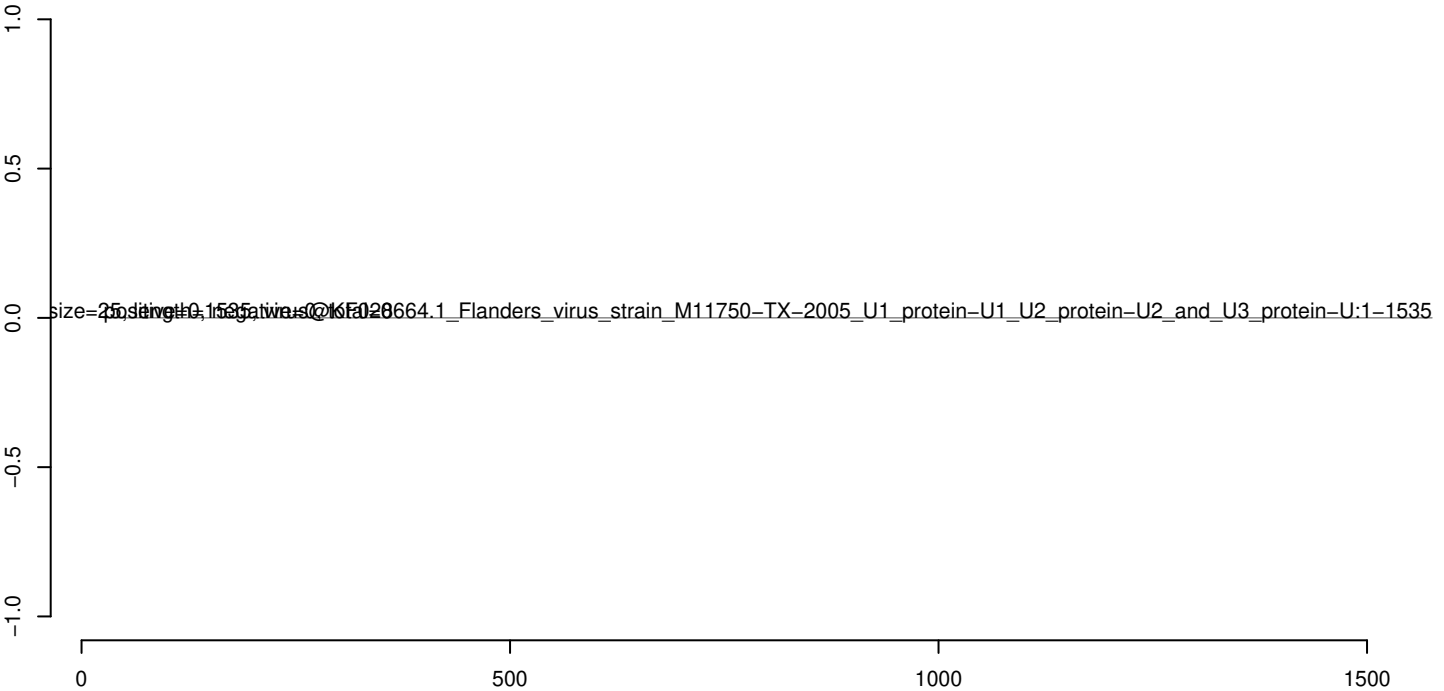
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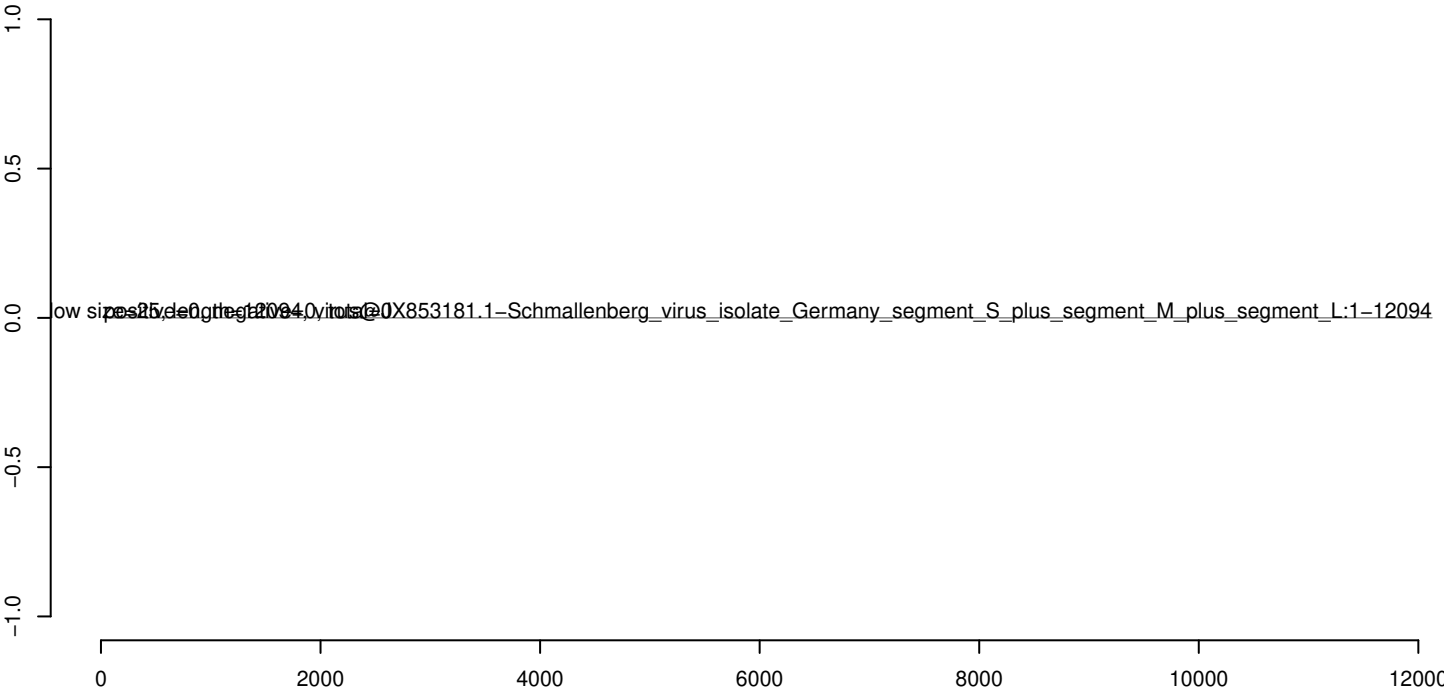
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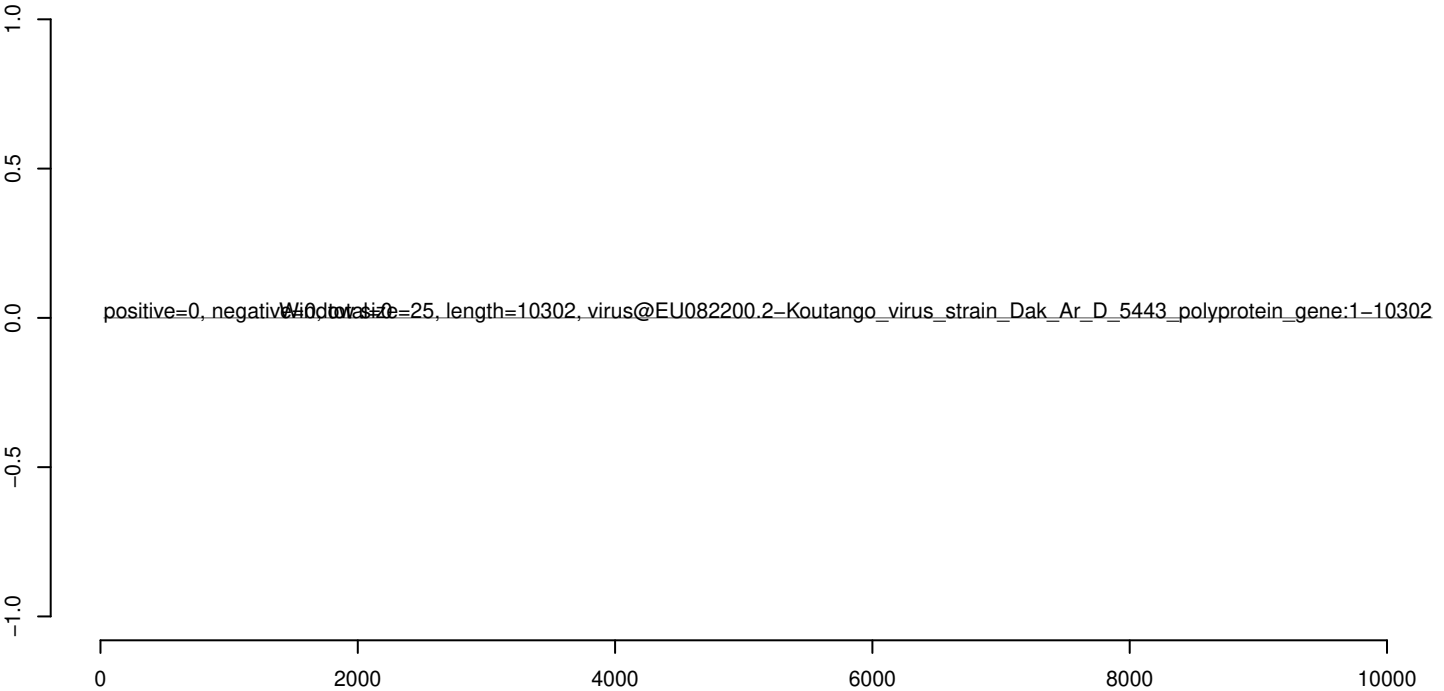
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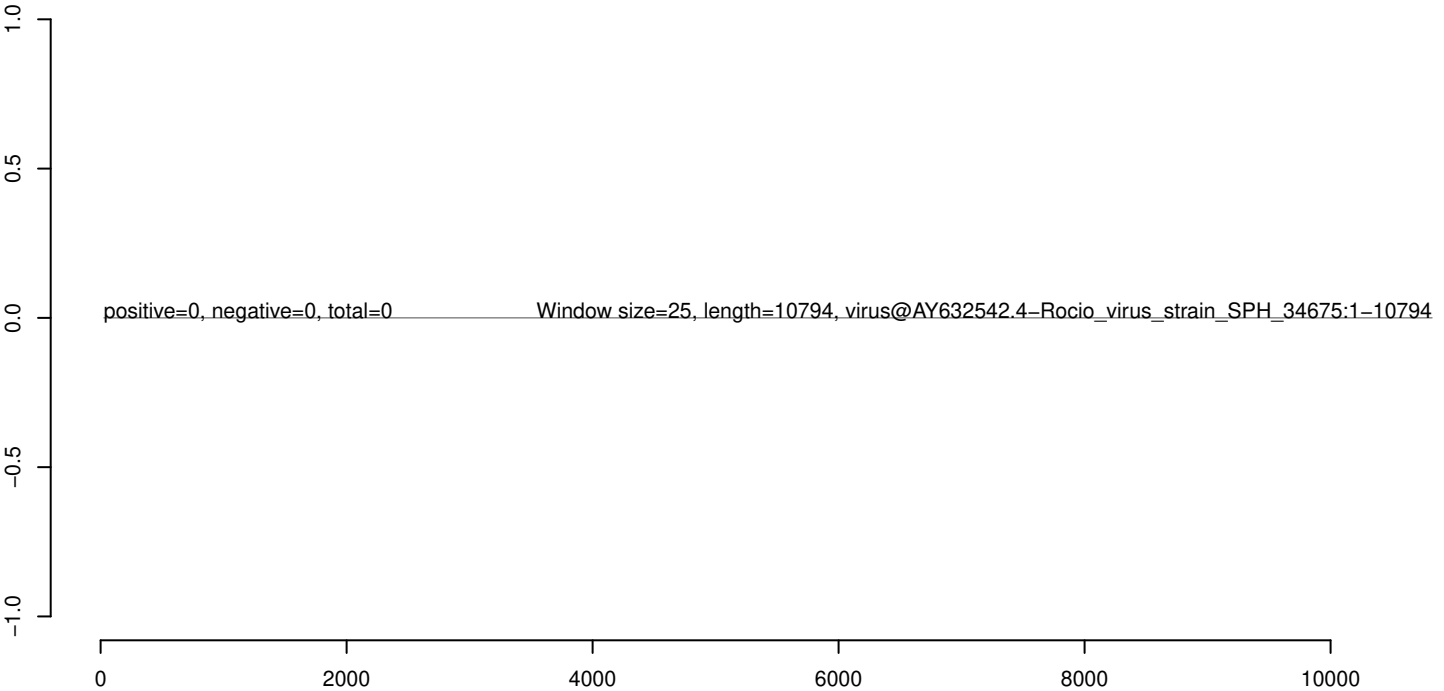
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