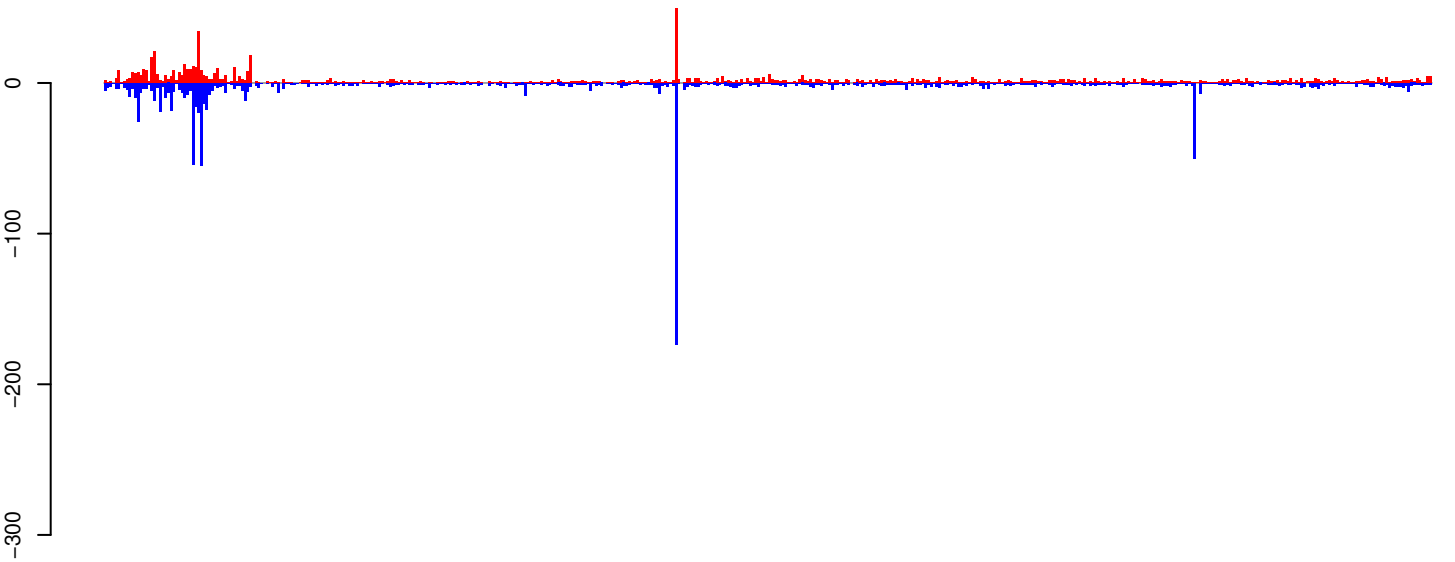
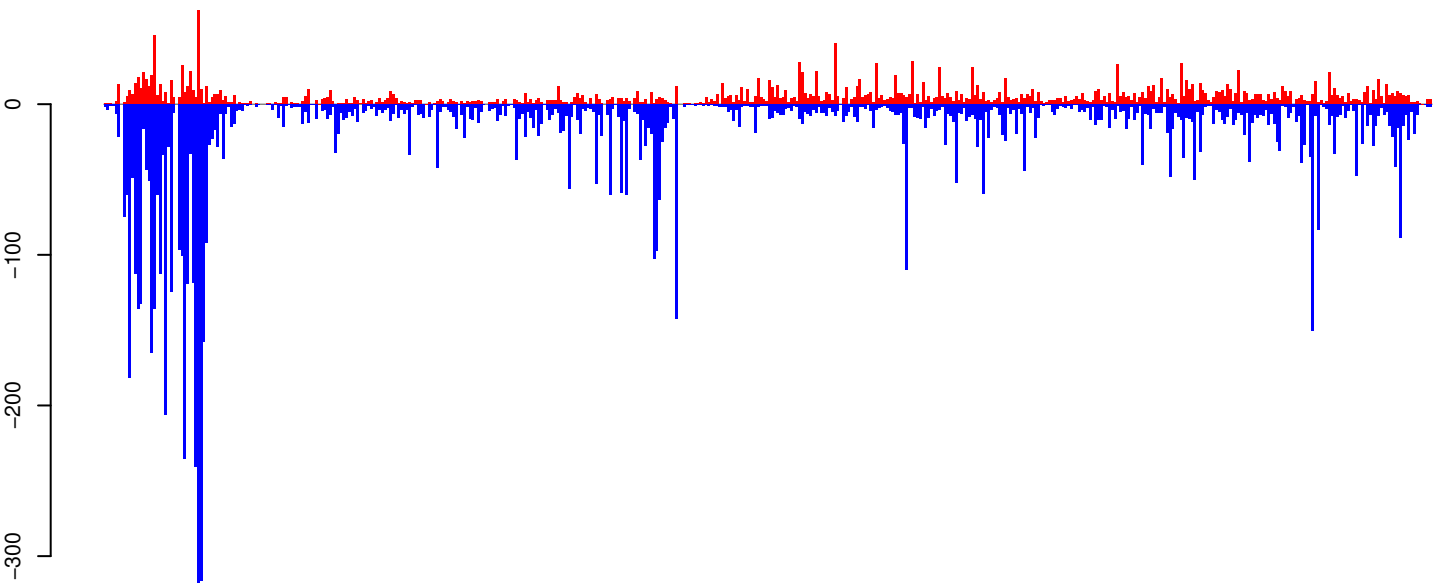


AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



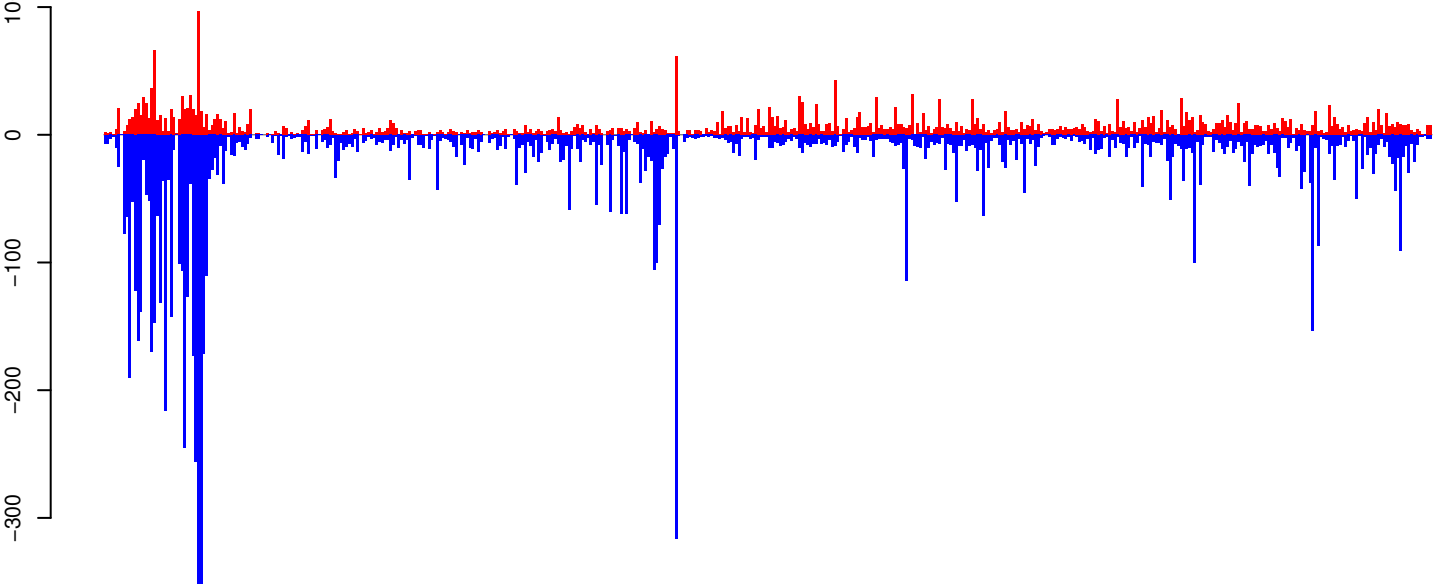
positive=826, negative=1155, total=1981

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



positive=2450, negative=8304, total=10753

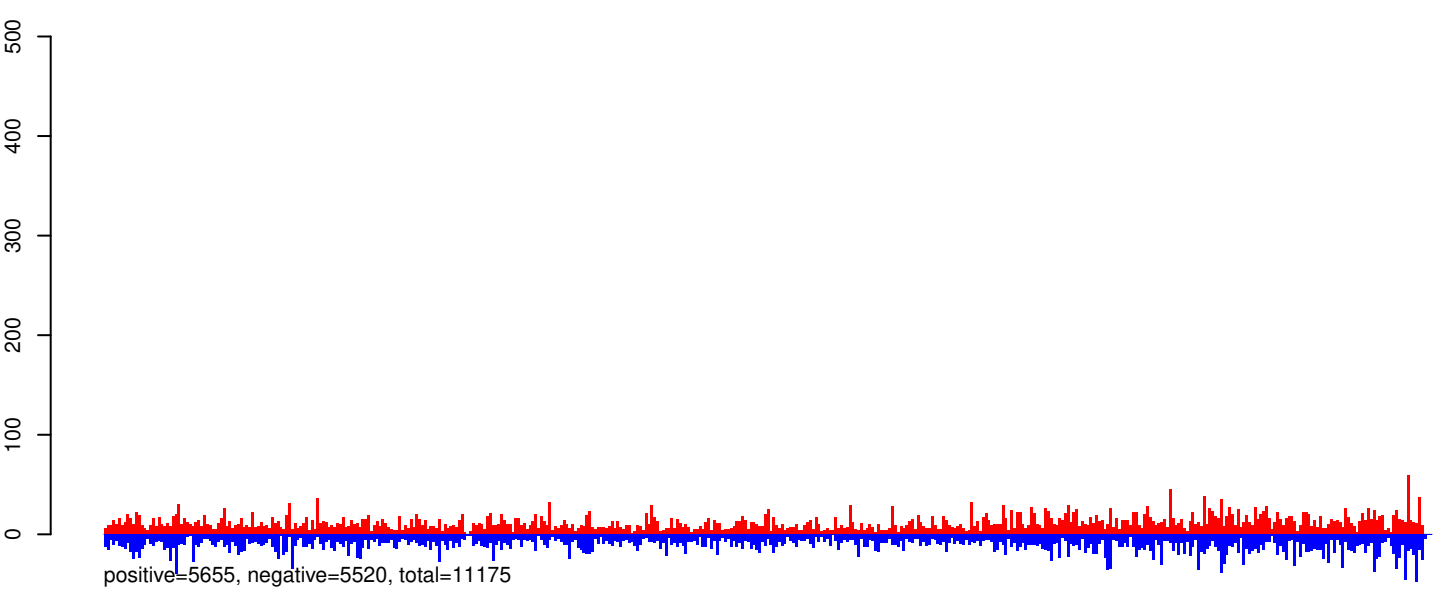
AeAeg_Aag2_mock_dsFluc_RA.rep



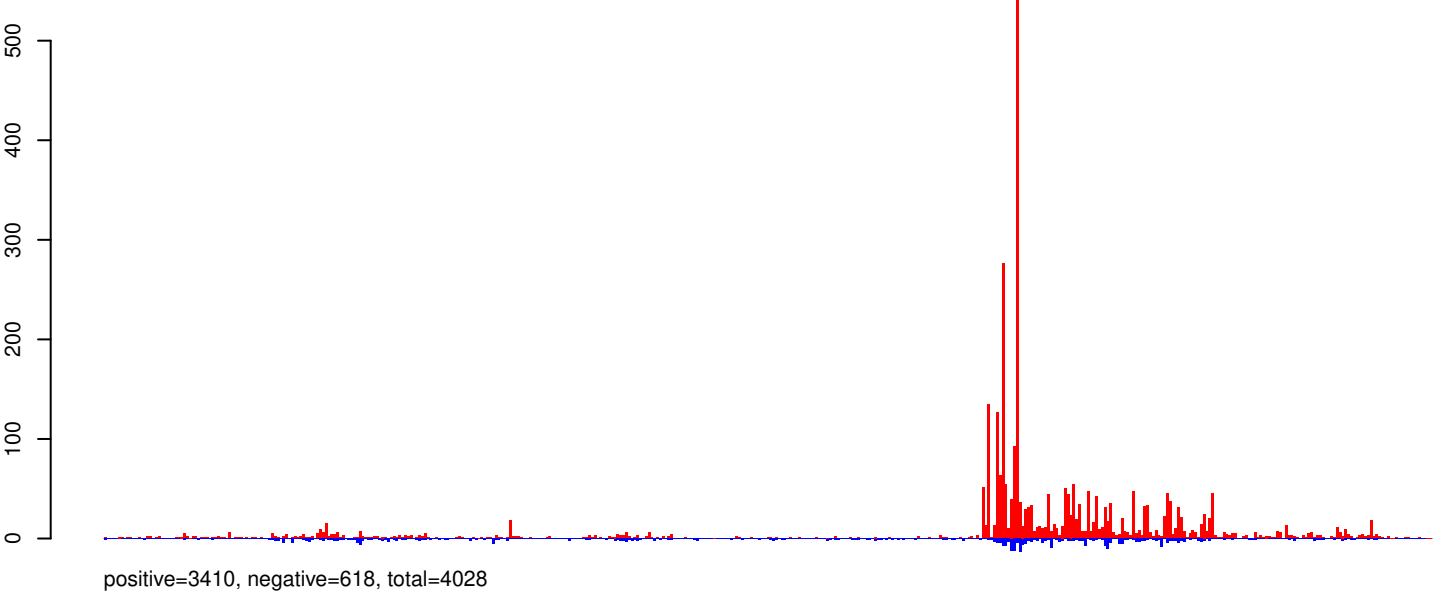
=25, length=12099, vireg@MC-9052, total=10754 Charoen-like_virus_isolate_Rio_segment_S_plus_segment_M_plus_segment_L_Nucle:1-12099

0 2000 4000 6000 8000 10000 12000

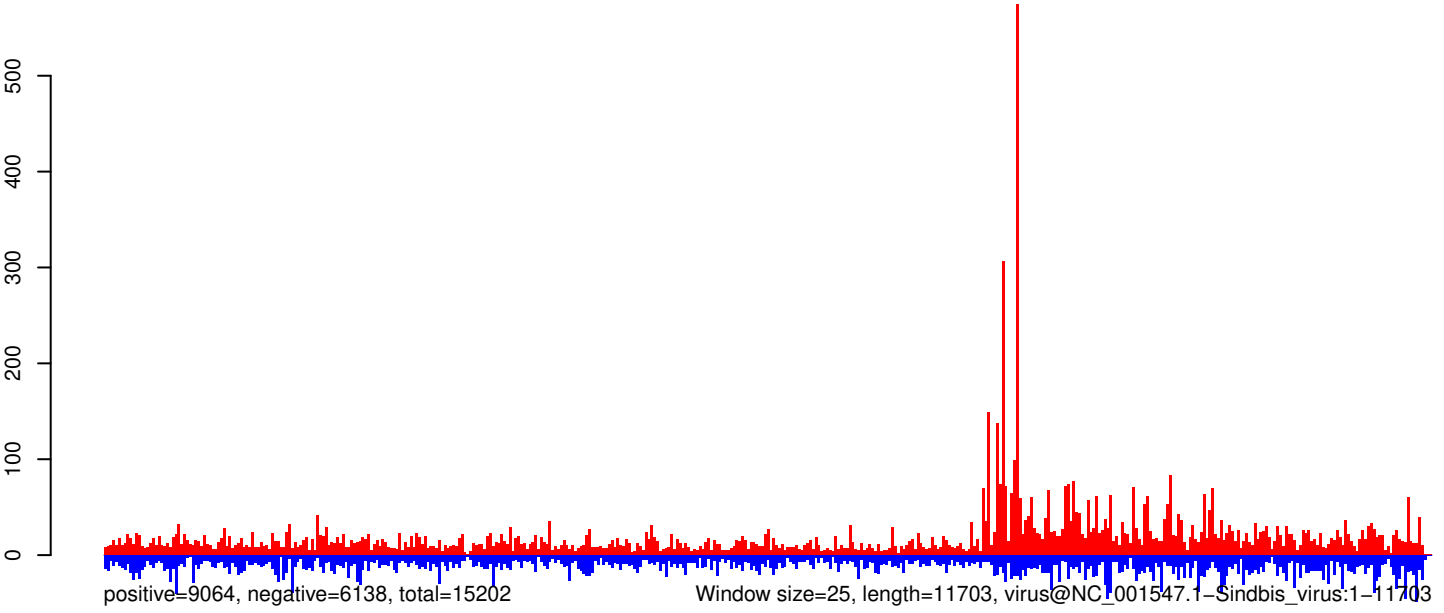
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



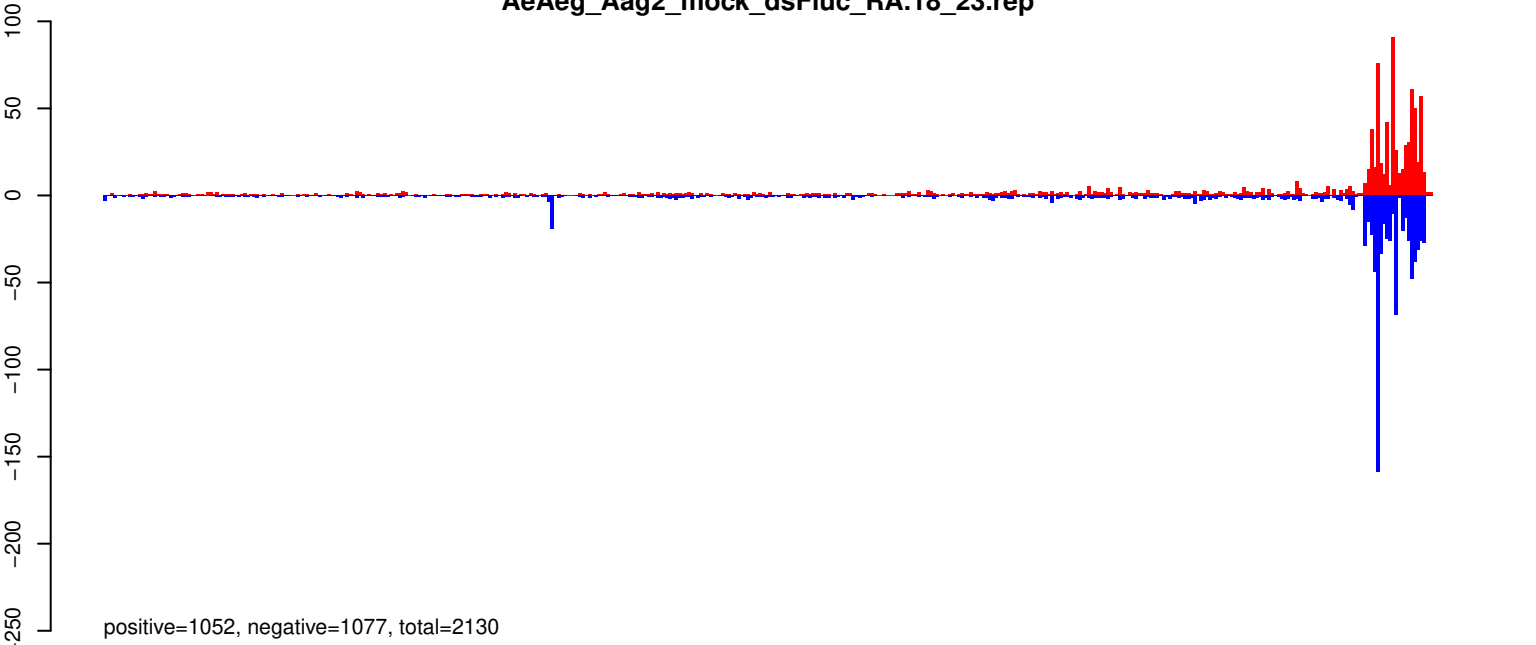
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



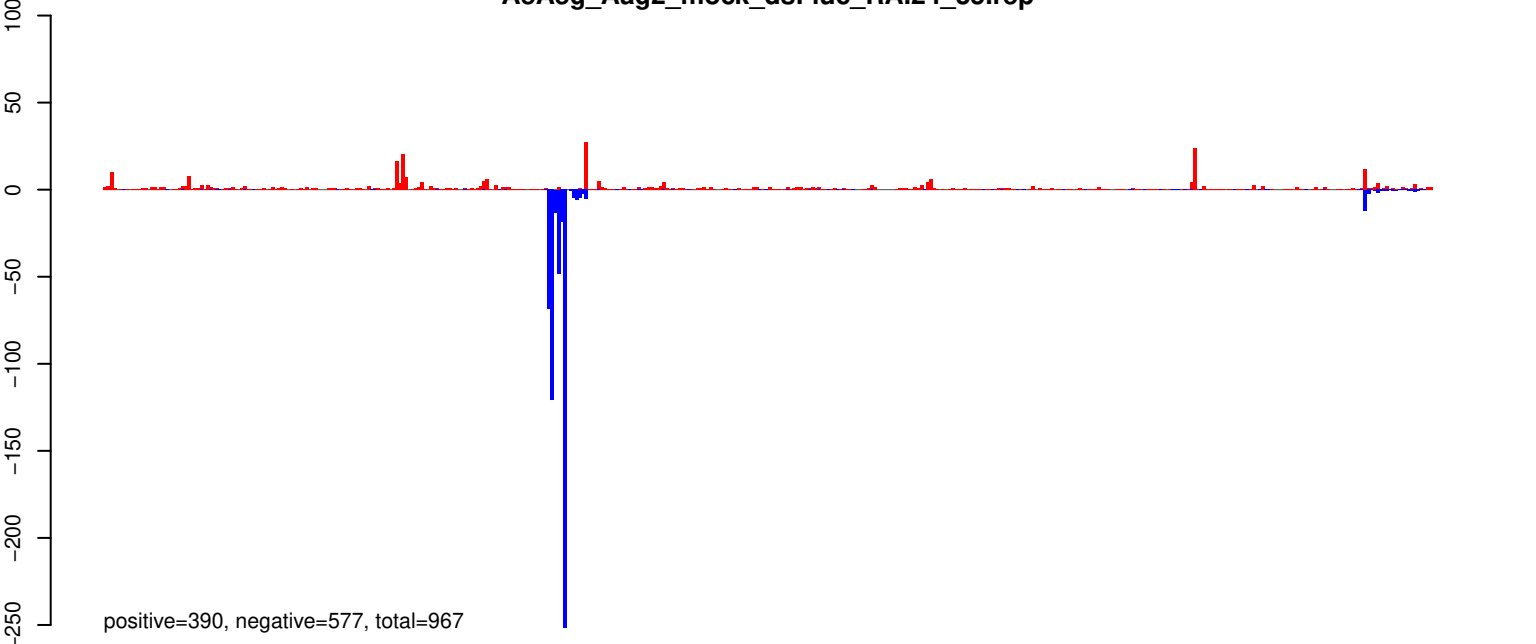
AeAeg_Aag2_mock_dsFluc_RA.rep



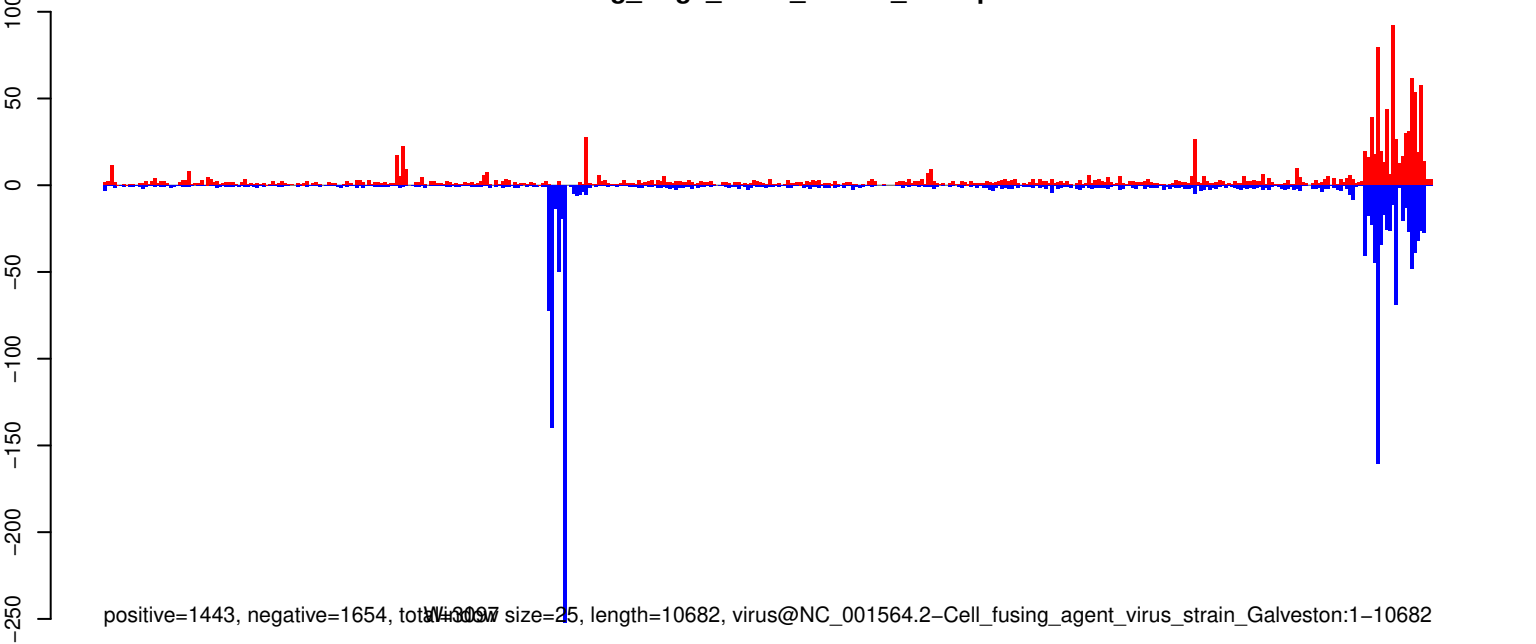
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

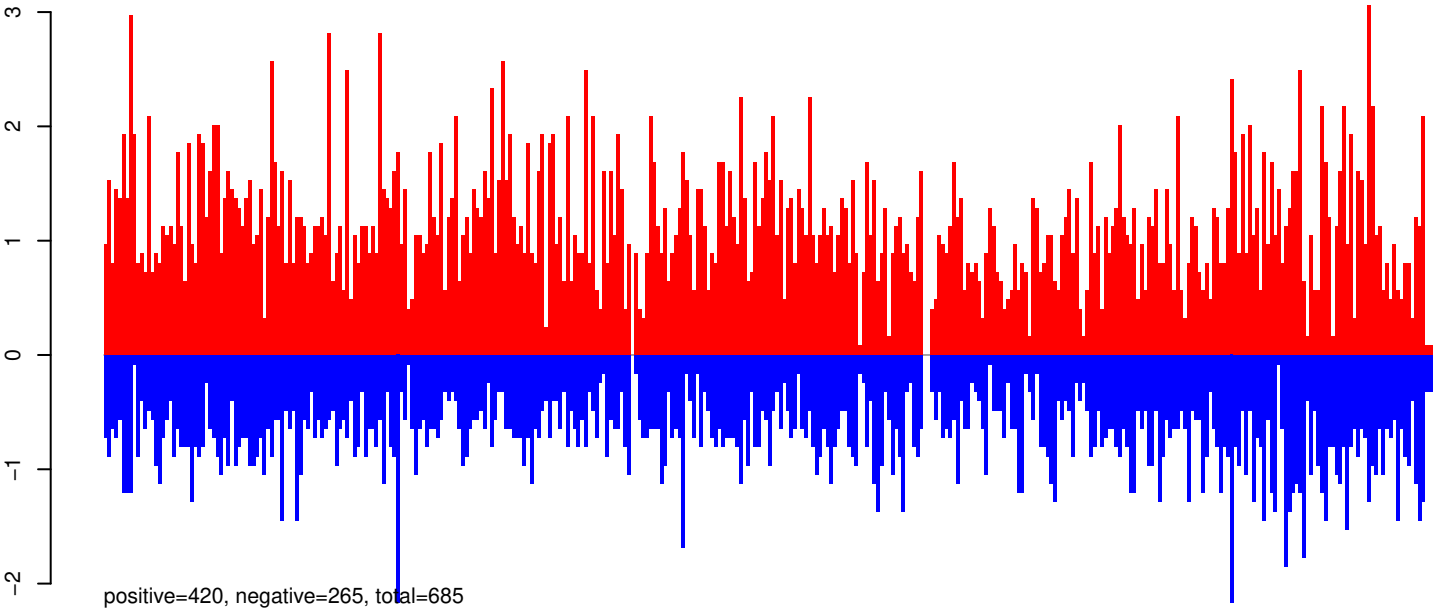


AeAeg_Aag2_mock_dsFluc_RA.rep

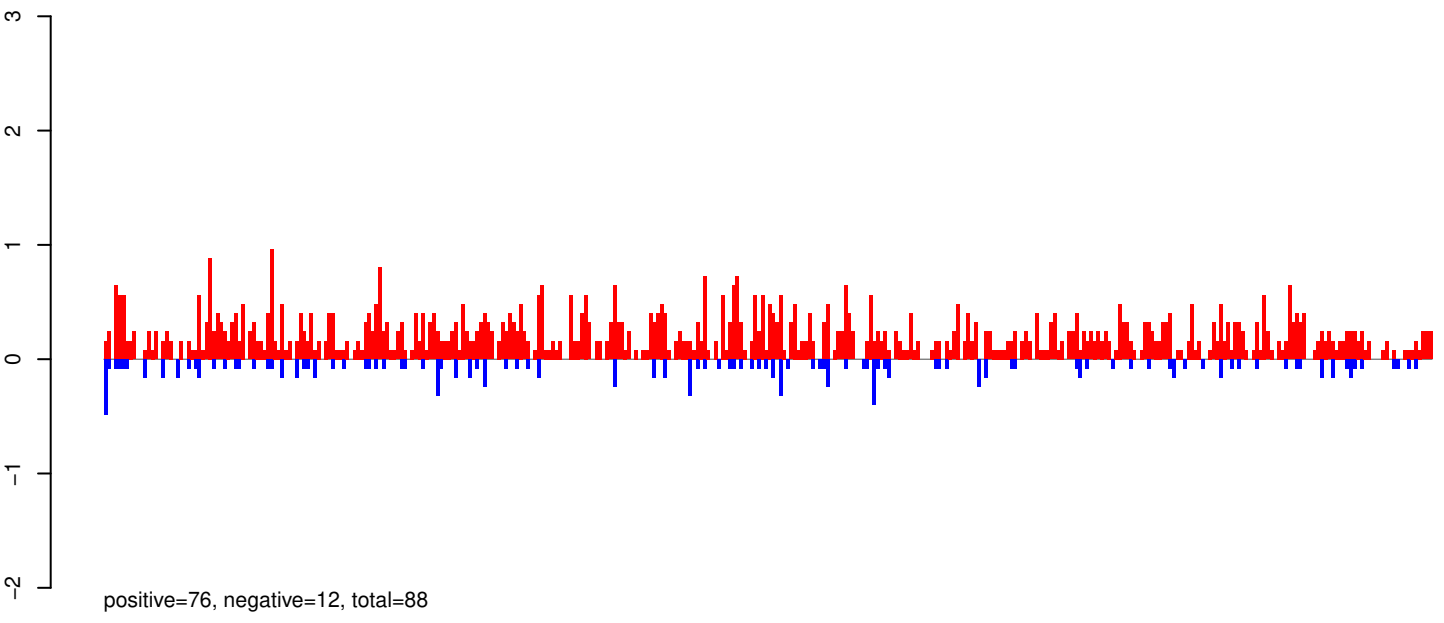


0 2000 4000 6000 8000 10000

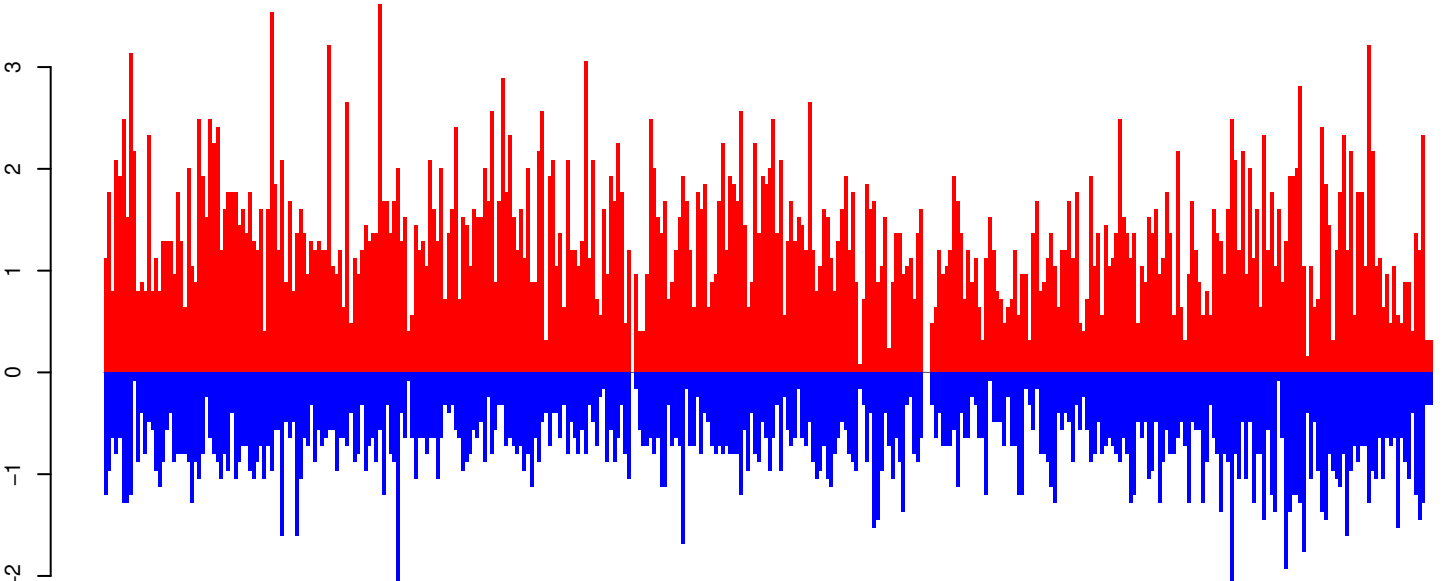
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



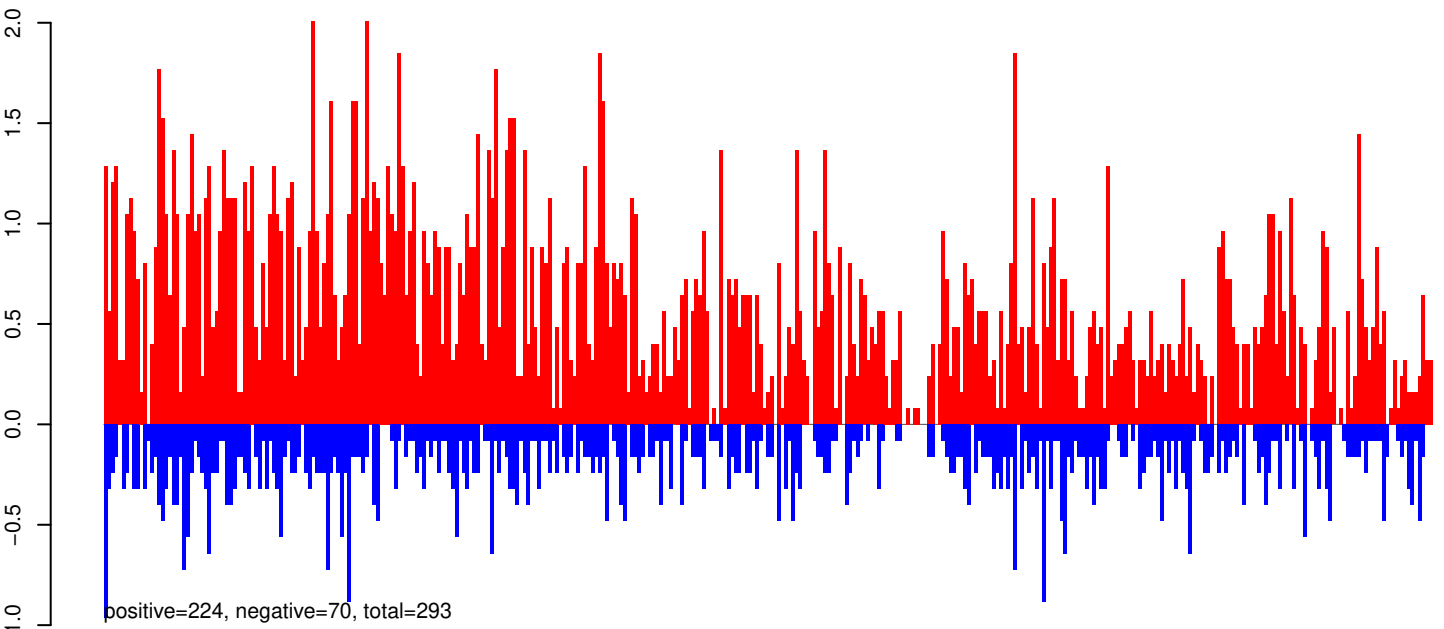
AeAeg_Aag2_mock_dsFluc_RA.rep



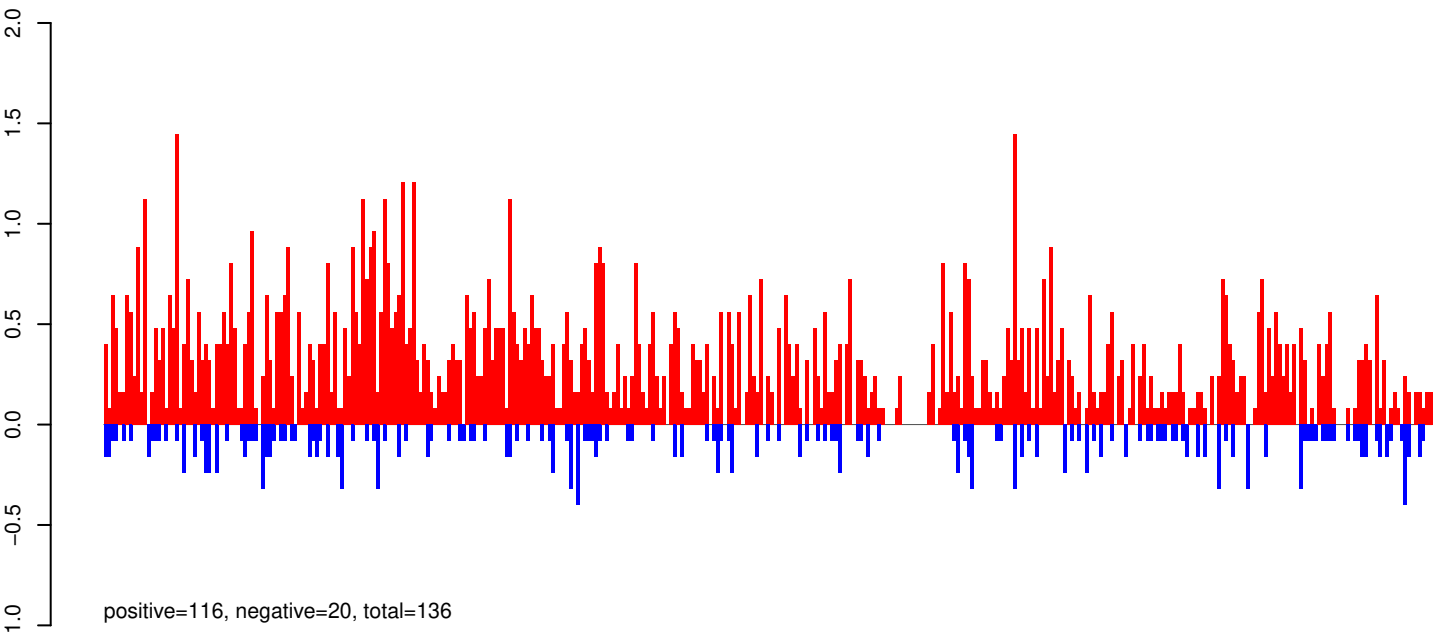
Window size=25, length=1852, use=0.7003924.1-Cricket_paralysis_virus_nonstructural_polyprotein_and_structural_polyprotein_genes:1-9185

0 2000 4000 6000 8000

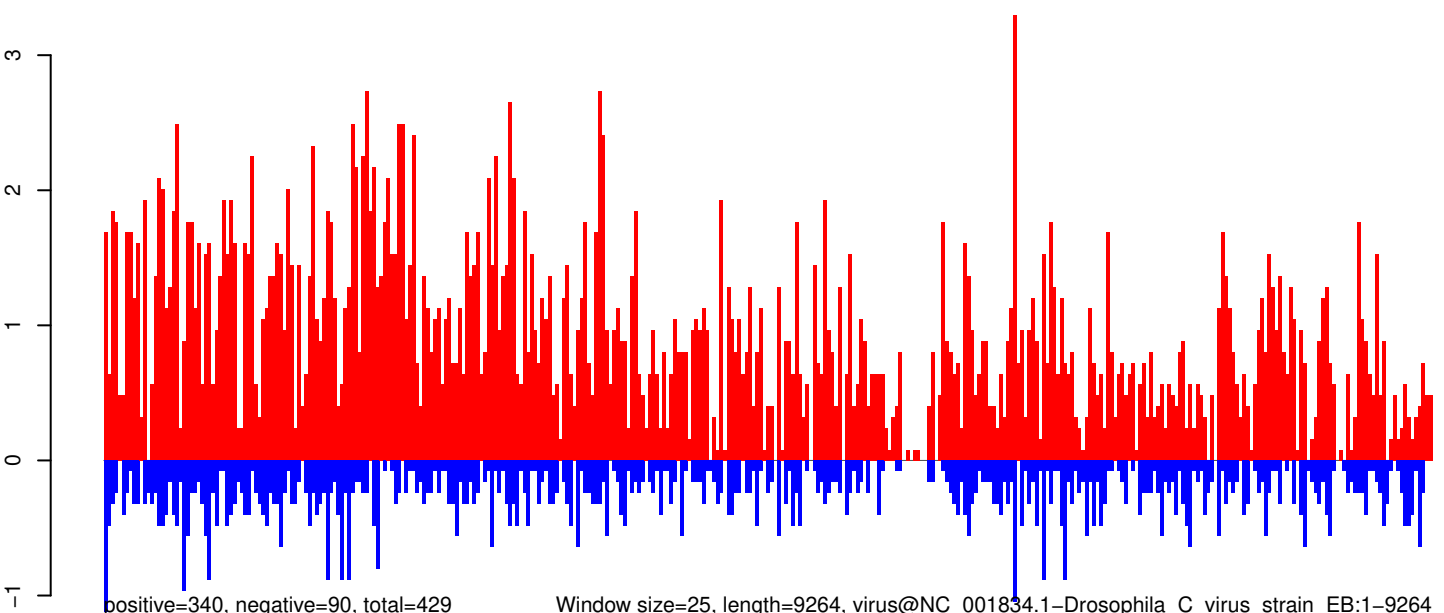
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



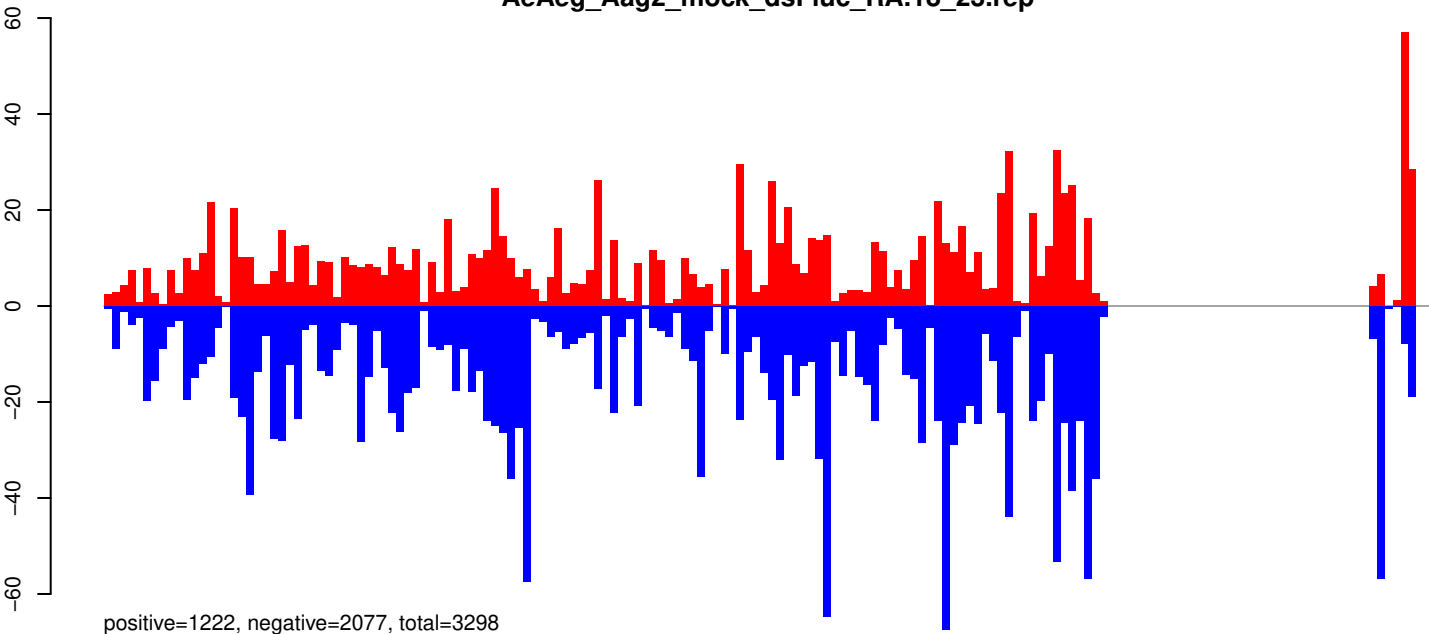
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



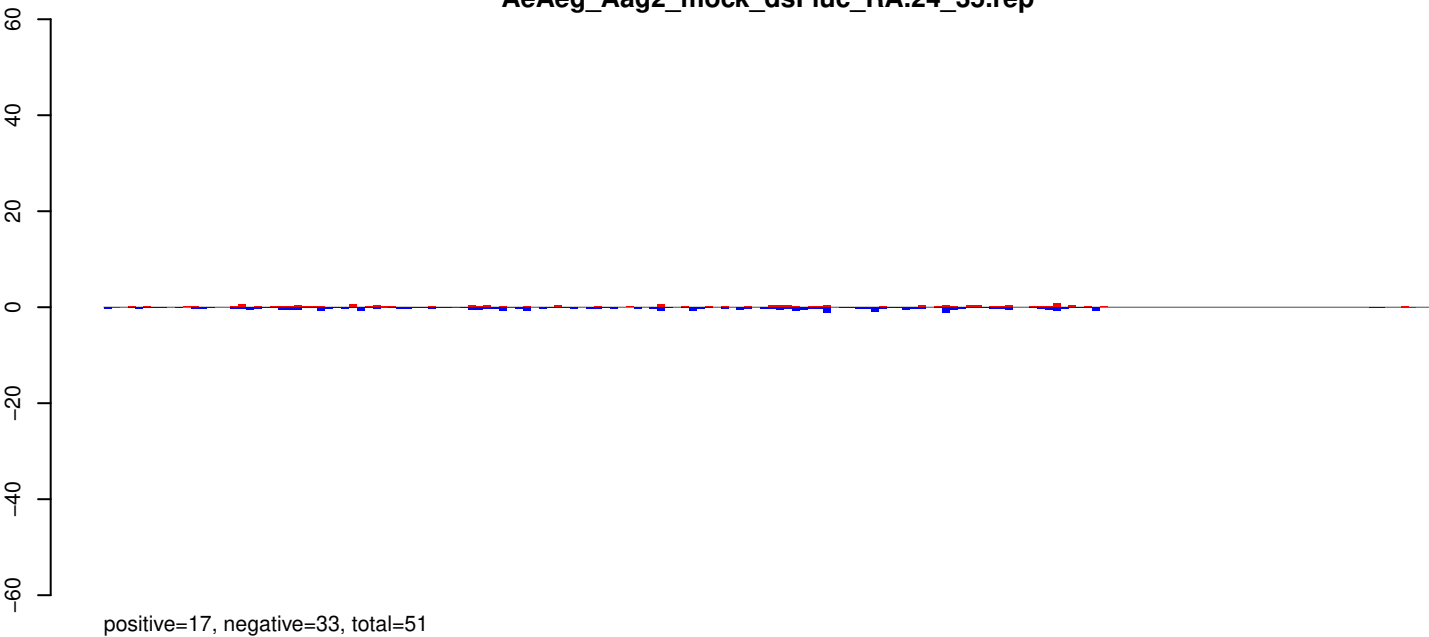
AeAeg_Aag2_mock_dsFluc_RA.rep



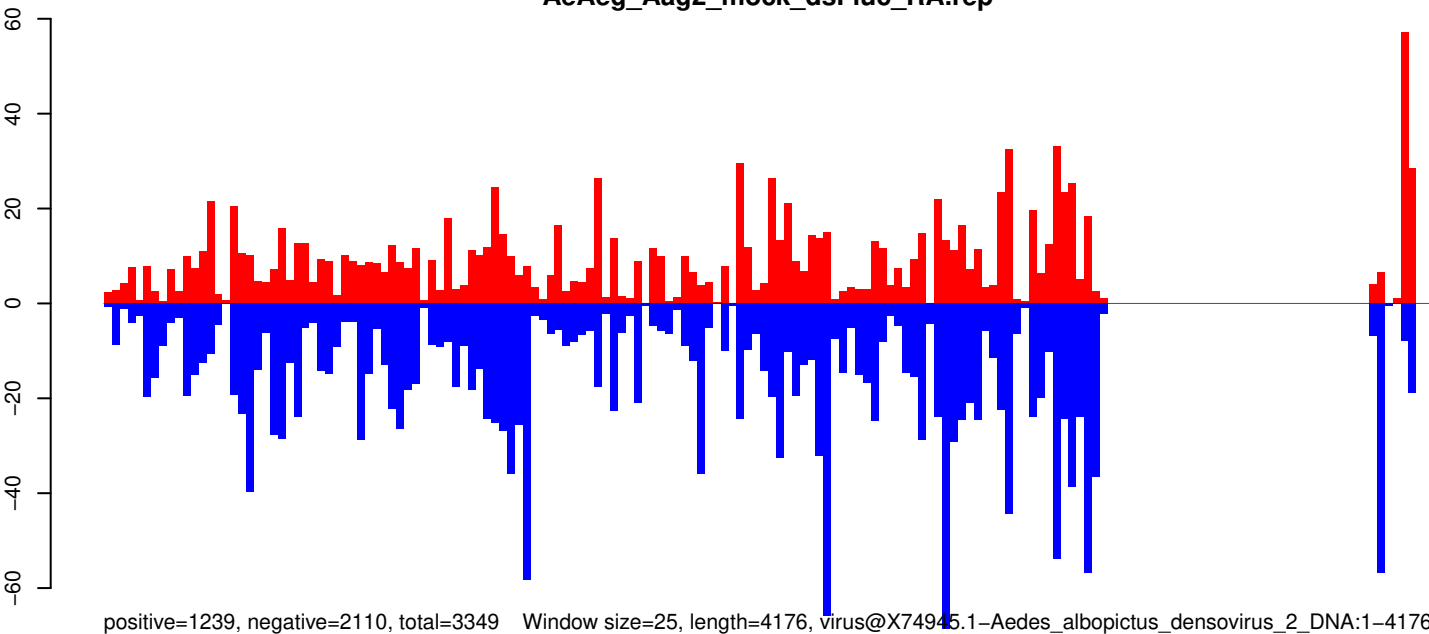
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

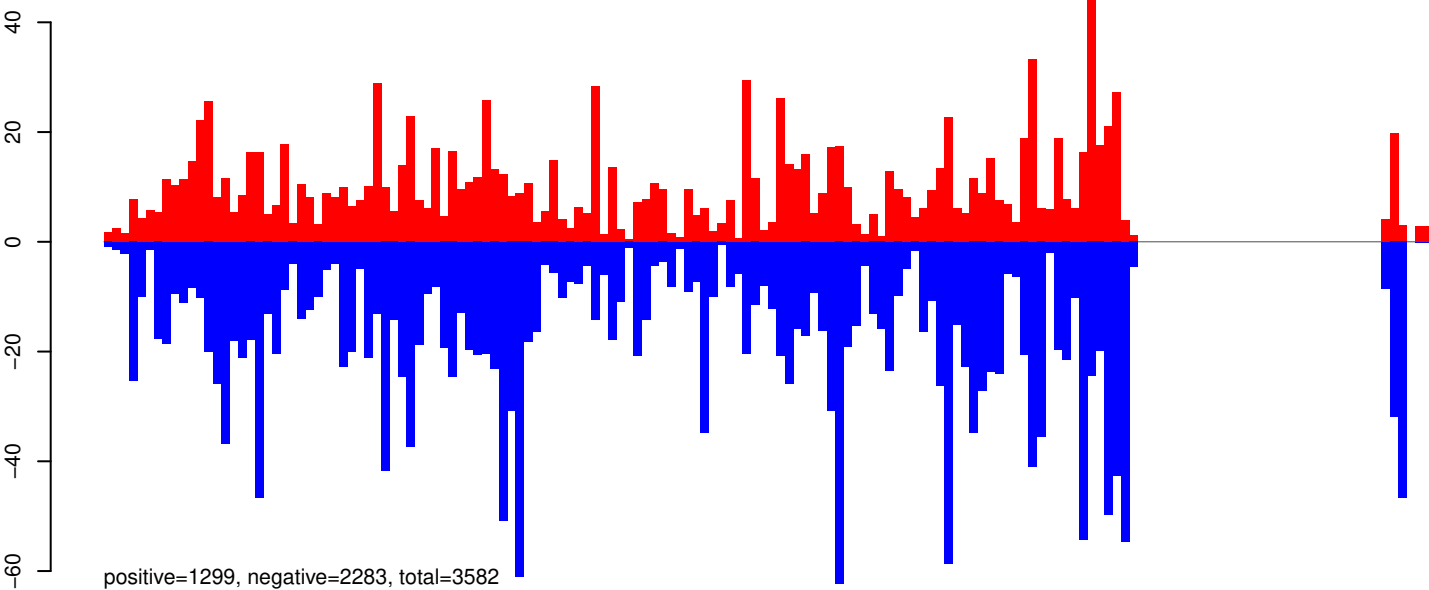


AeAeg_Aag2_mock_dsFluc_RA.rep

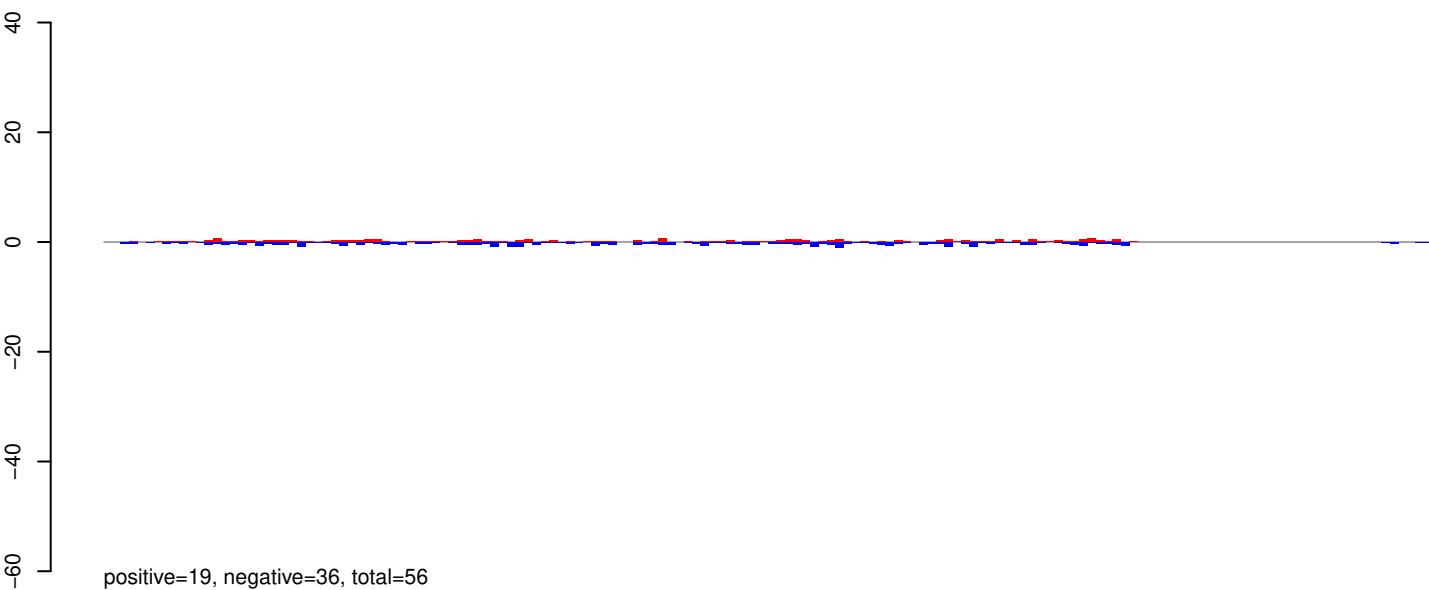


0 1000 2000 3000 4000

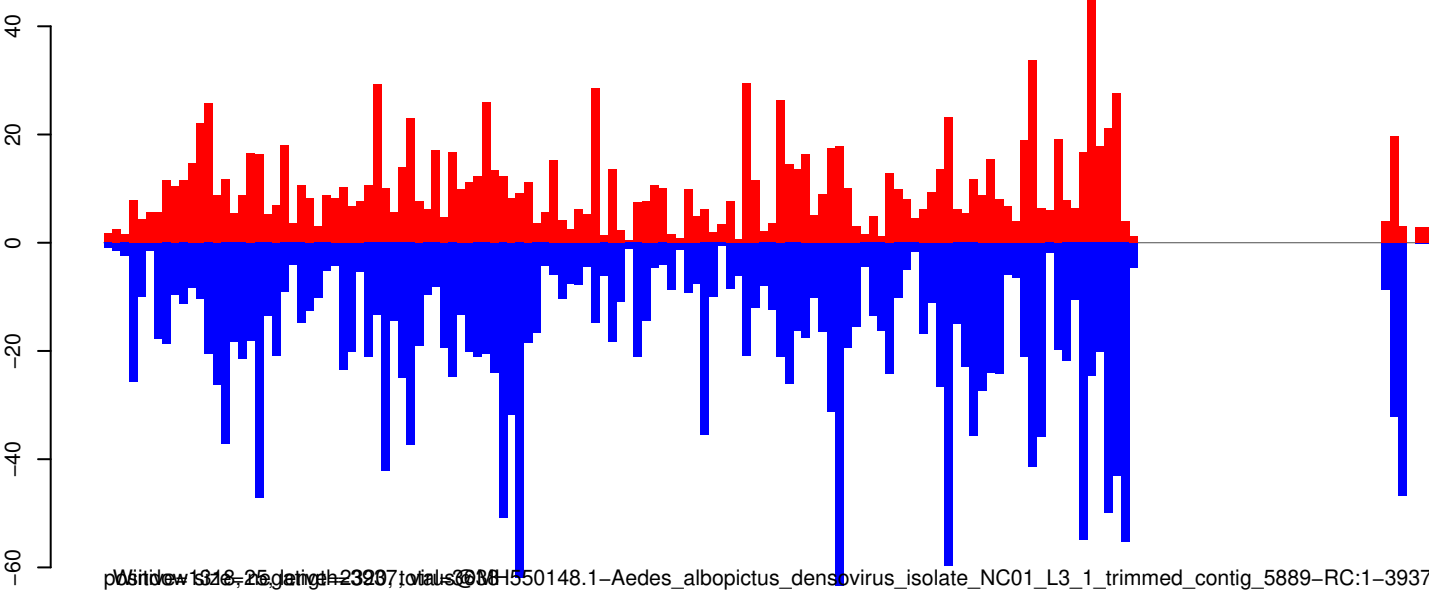
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



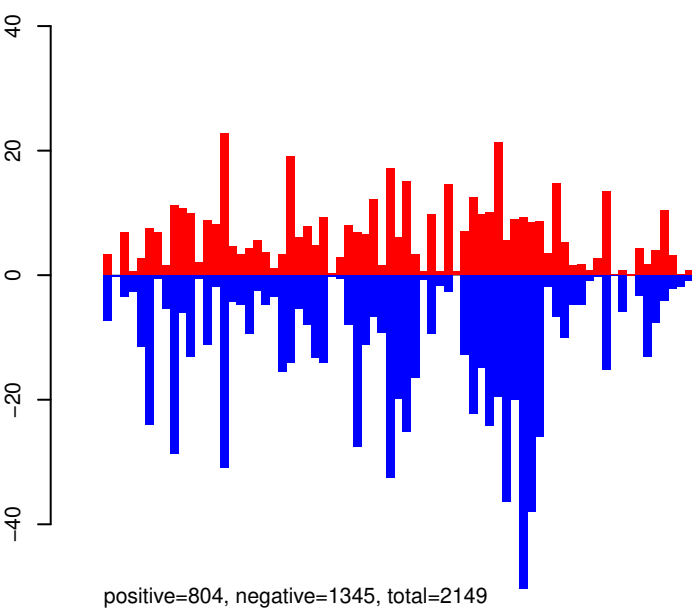
AeAeg_Aag2_mock_dsFluc_RA.rep



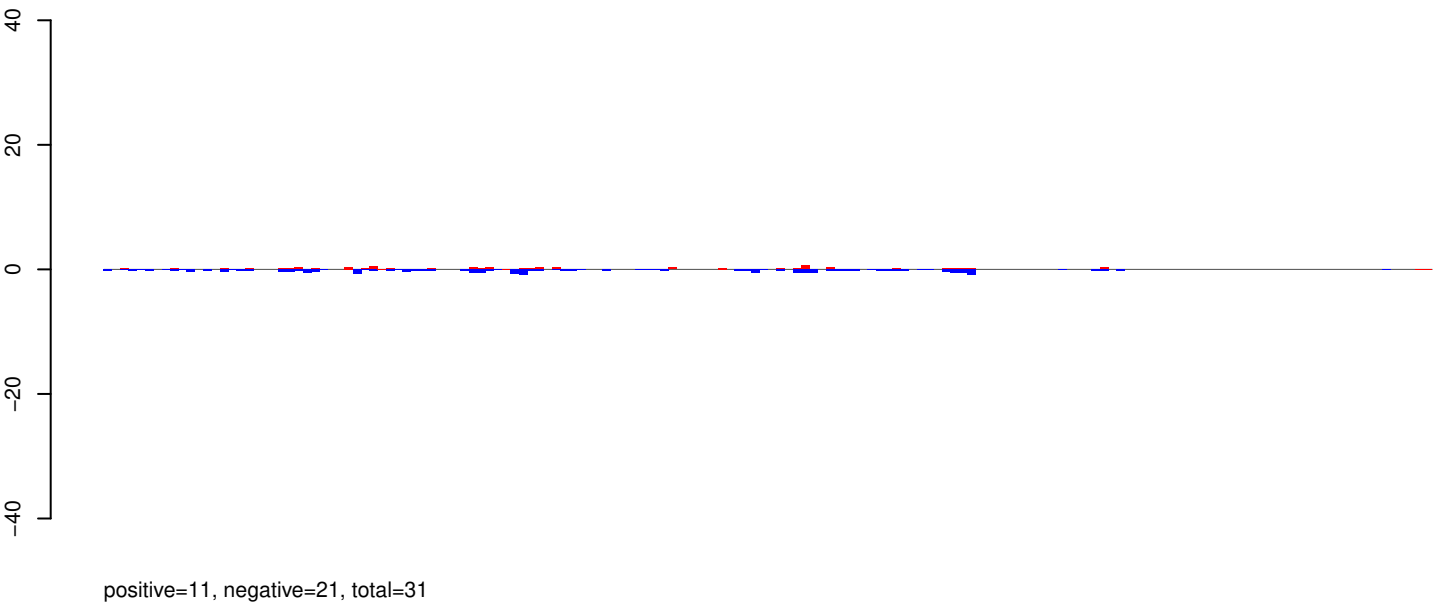
Window Size=25, length=3937, total=3645
H50148.1-Aedes albopictus_densivirus_isolate_NC01_L3_1_trimmed_contig_5889-RC:1-3937

0 1000 2000 3000 4000

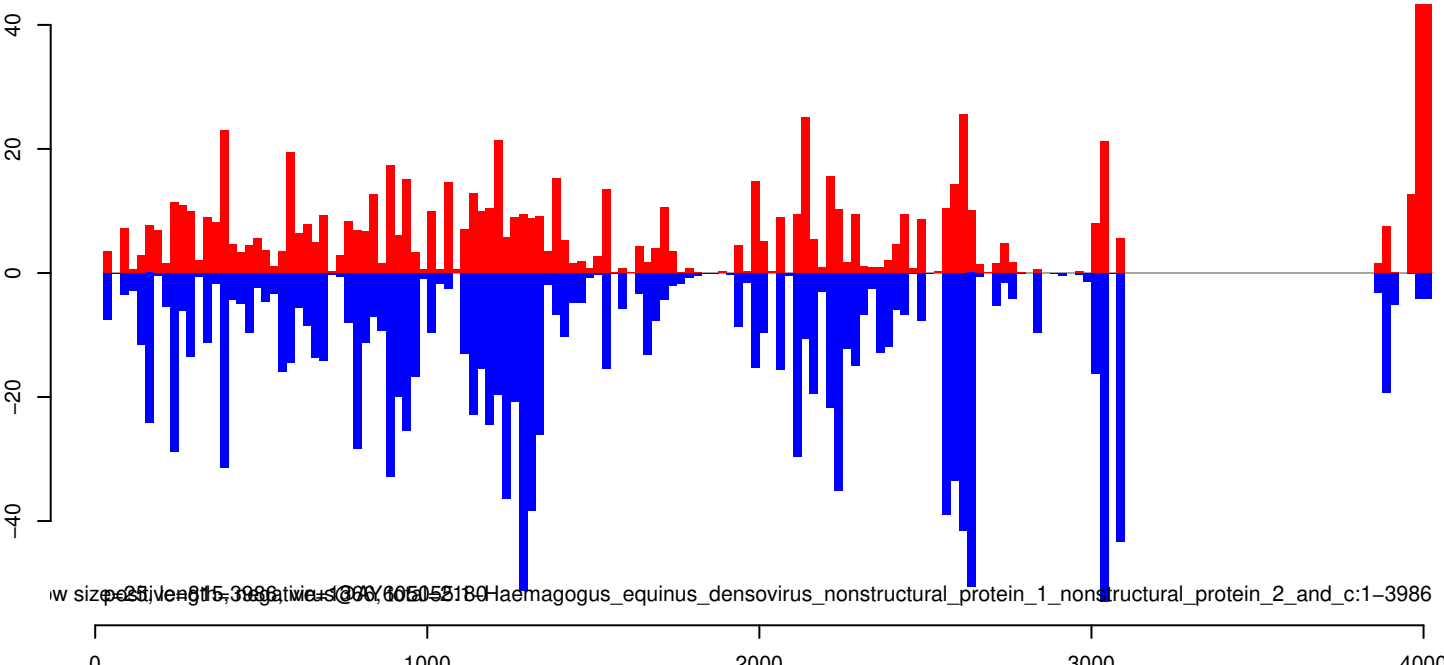
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



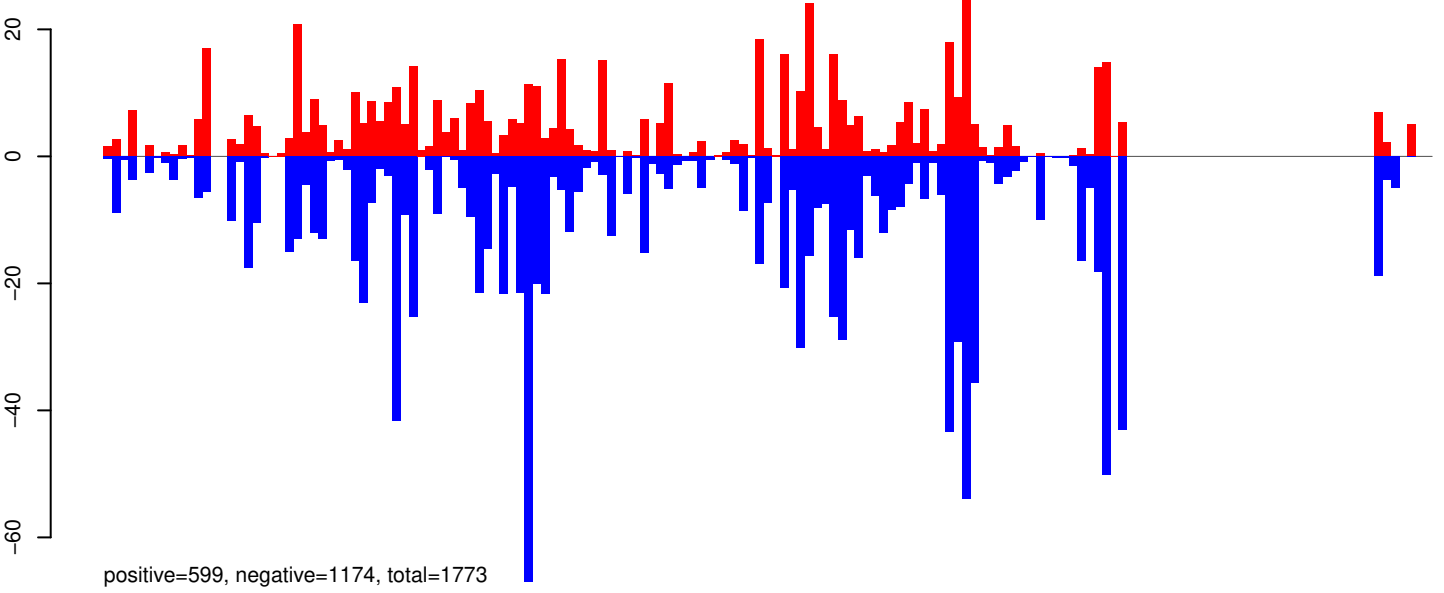
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



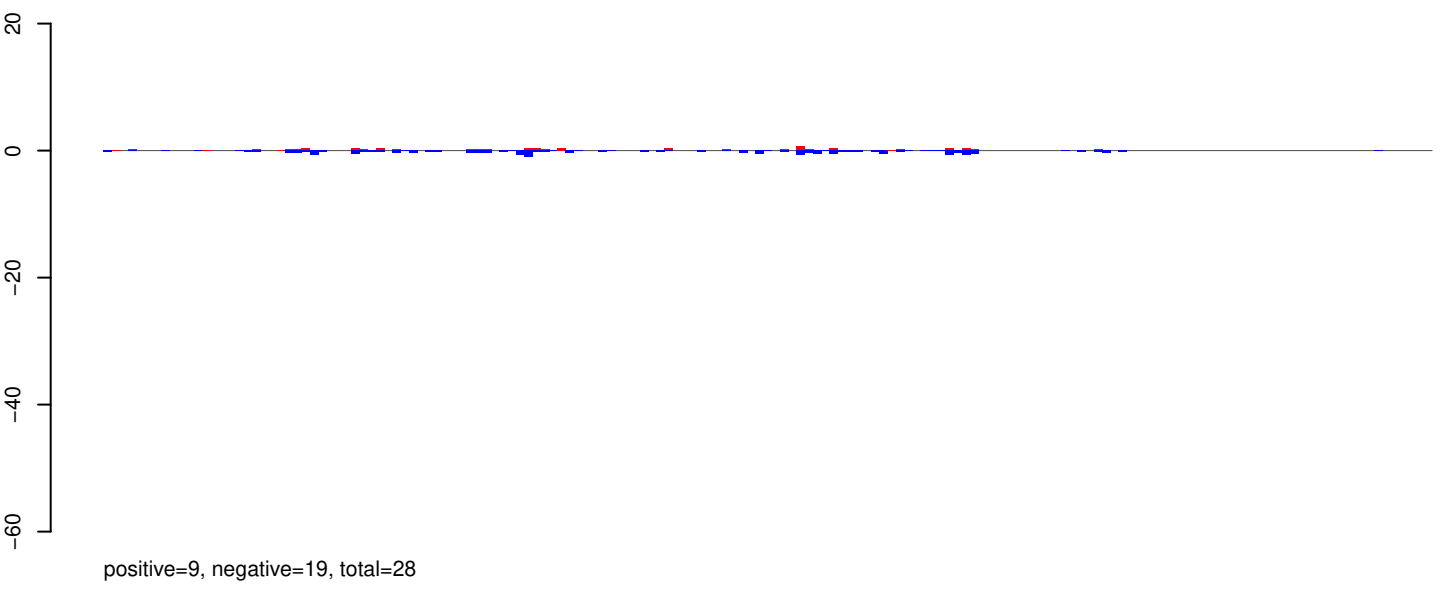
AeAeg_Aag2_mock_dsFluc_RA.rep



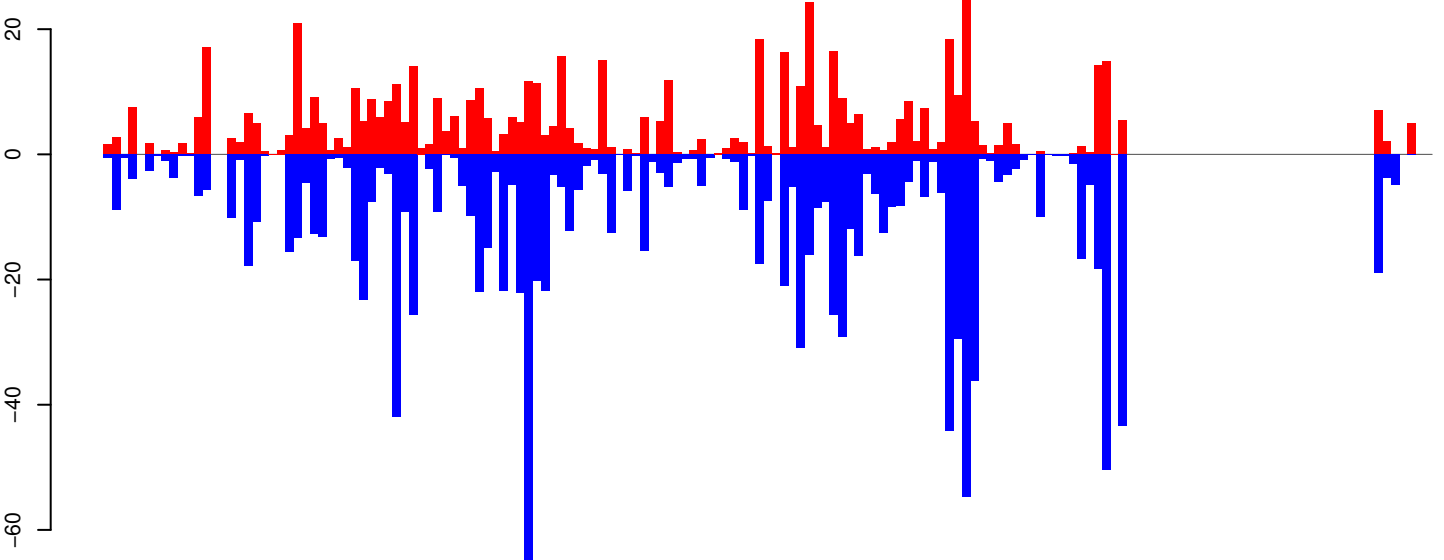
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



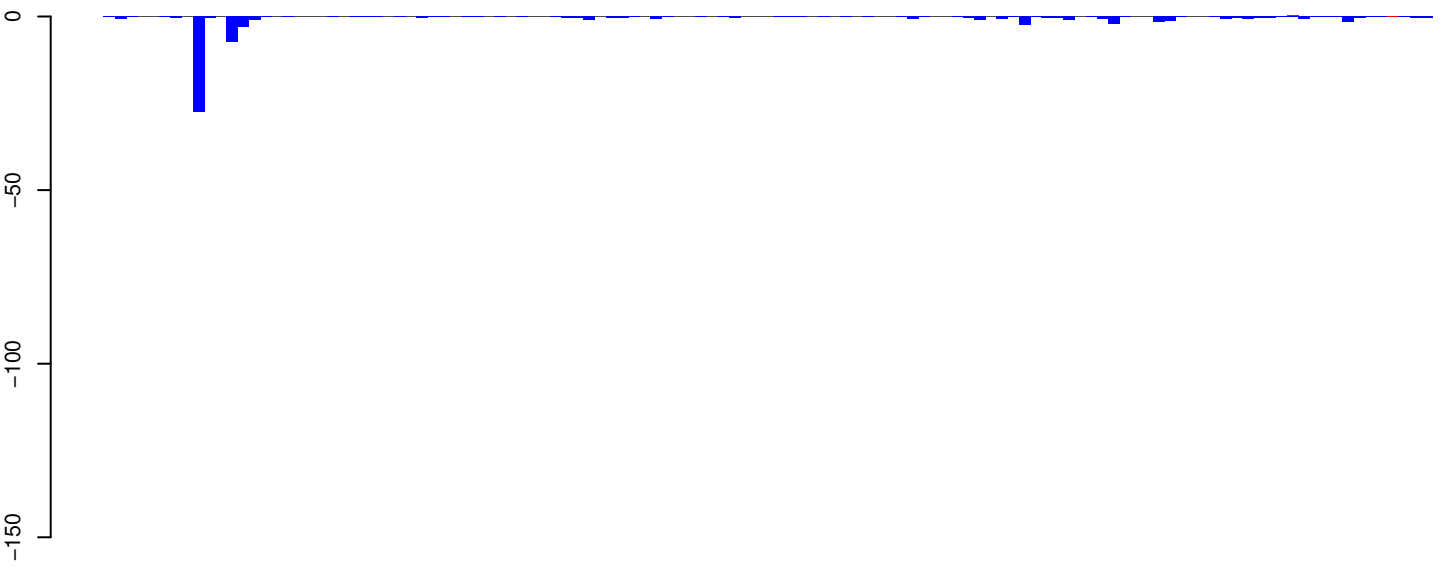
AeAeg_Aag2_mock_dsFluc_RA.rep



downloaded from <https://www.ncbi.nlm.nih.gov/nuccore/192419871> - Aedes albopictus_C6_36_cell_densovirus_non-structural_protein_1_non-structural_prot:1-4006

0 1000 2000 3000 4000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



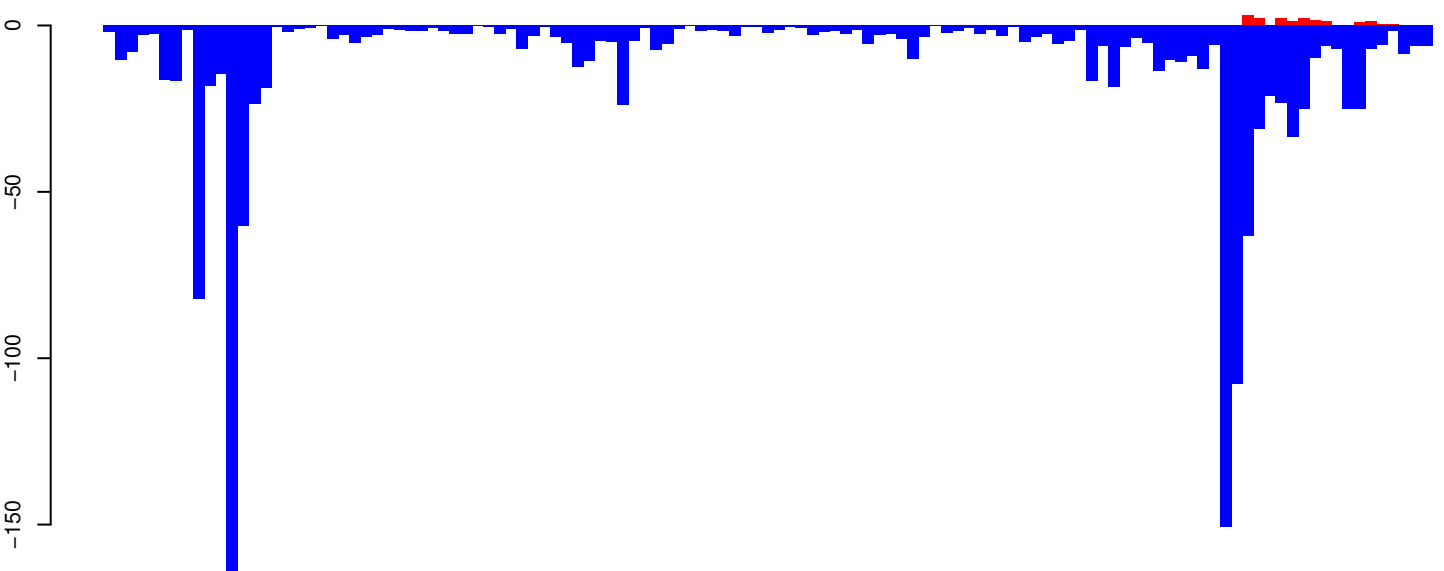
positive=2, negative=70, total=72

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



positive=17, negative=1291, total=1308

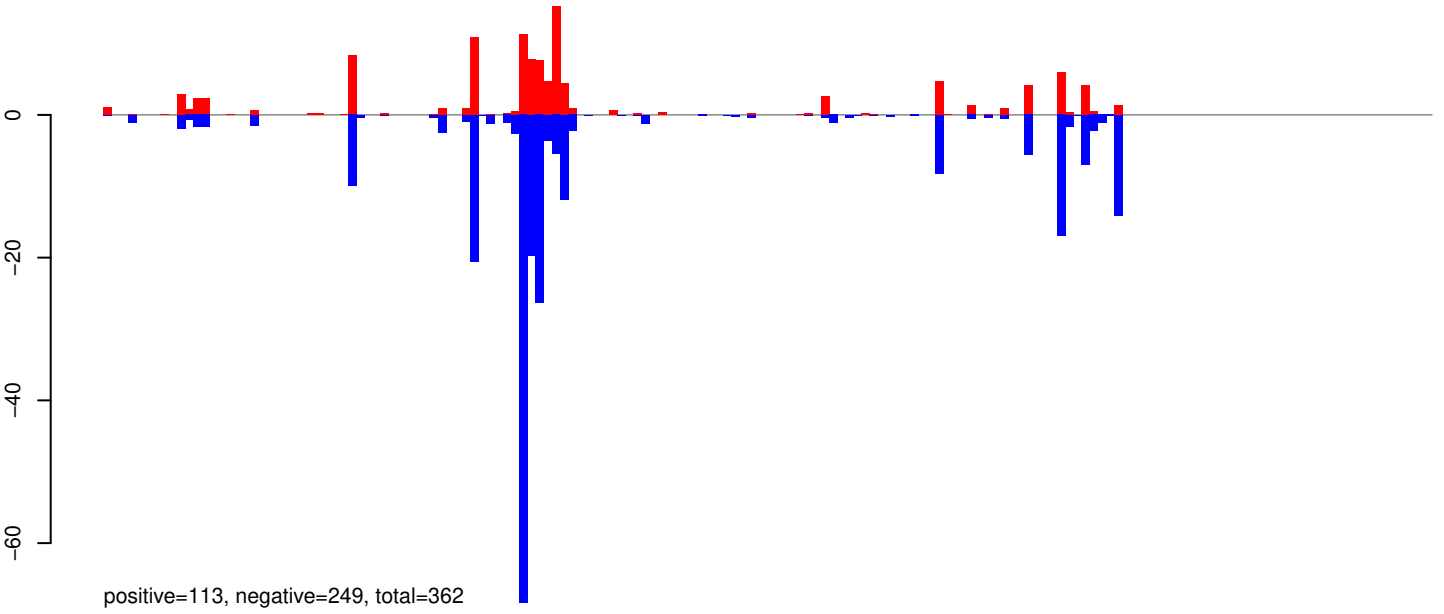
AeAeg_Aag2_mock_dsFluc_RA.rep



positive=19, negative=1361, total=1380 Window size=25, length=2965, virus@AY347953.2-Aedes_aegypti_genomic_sequence:1-2965

0 500 1000 1500 2000 2500 3000

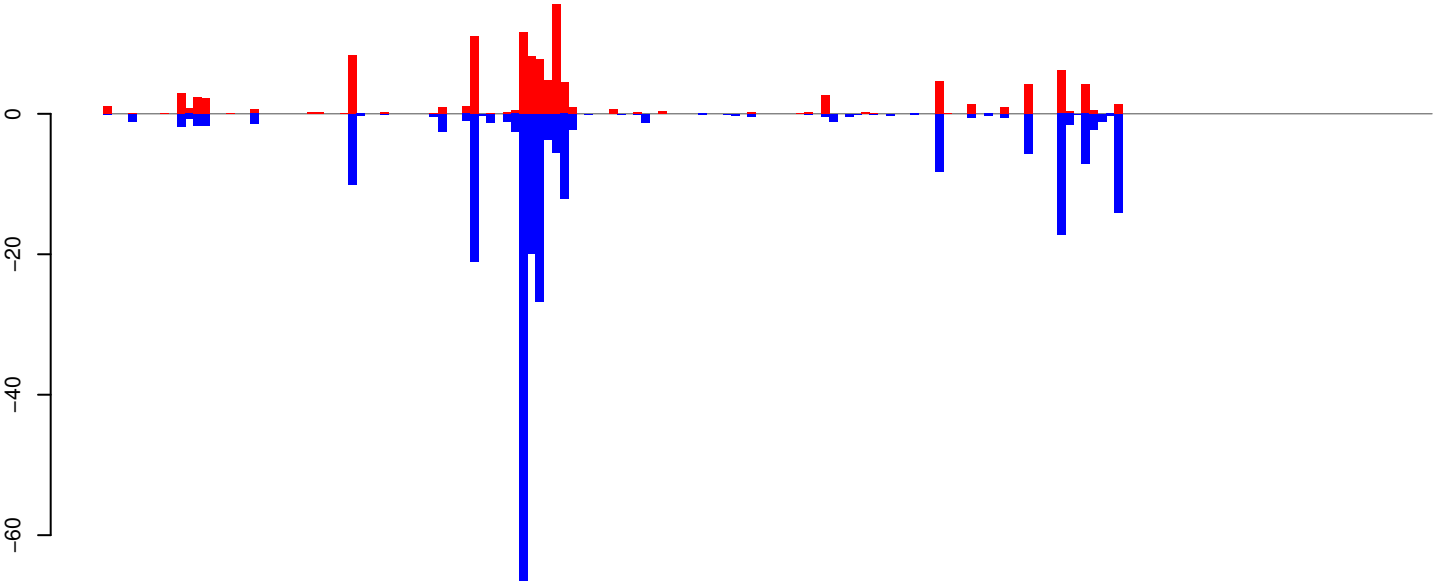
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



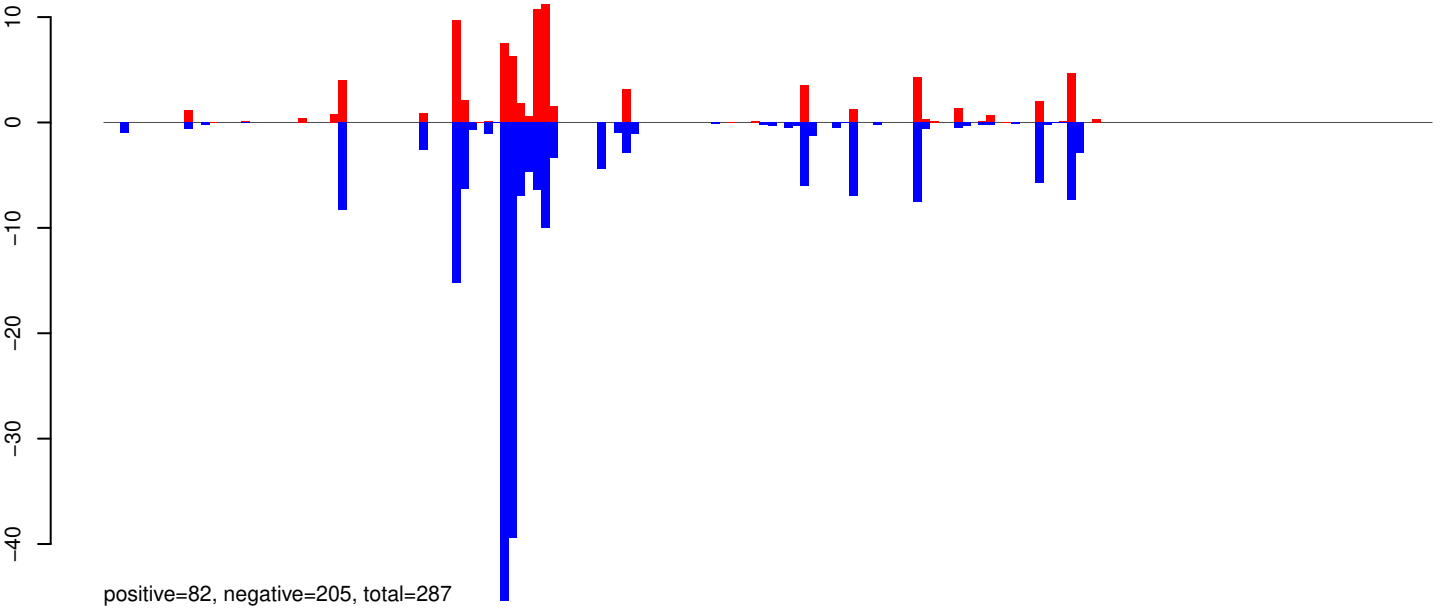
AeAeg_Aag2_mock_dsFluc_RA.rep



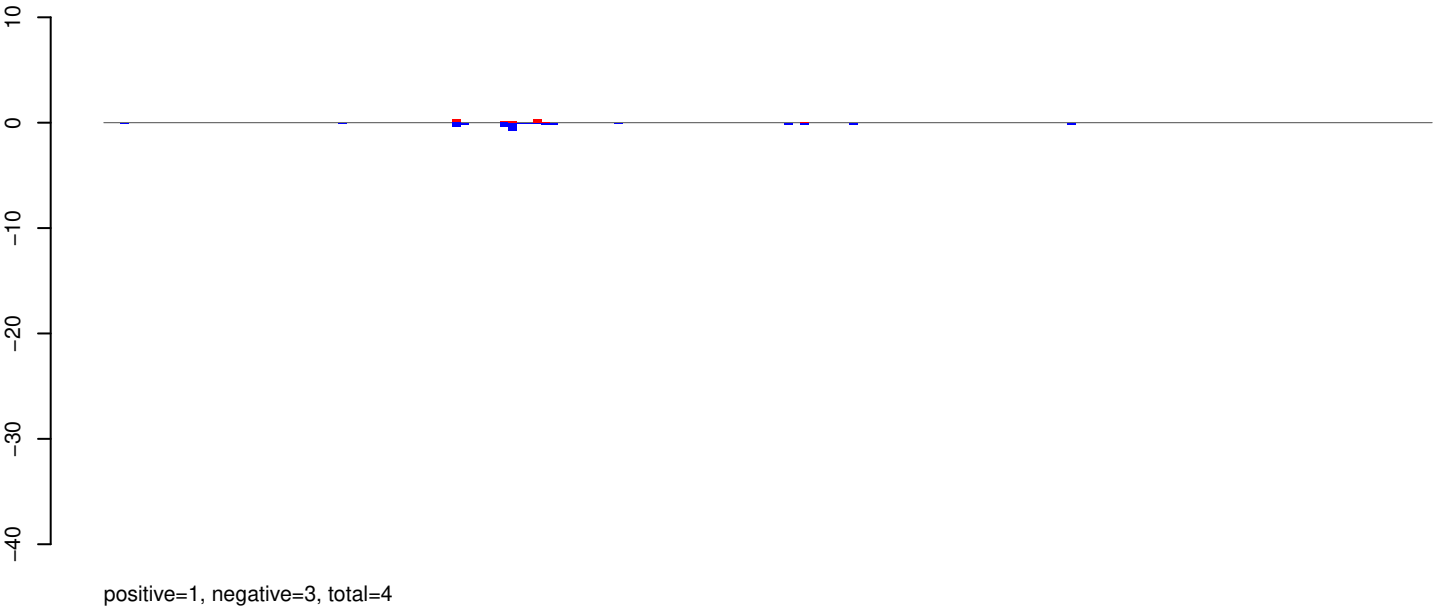
Window size=25, length=4073, virus@KX603698.1-Aedes albopictus_densovirus_5_strain_GZ05:1-4073

0 1000 2000 3000 4000

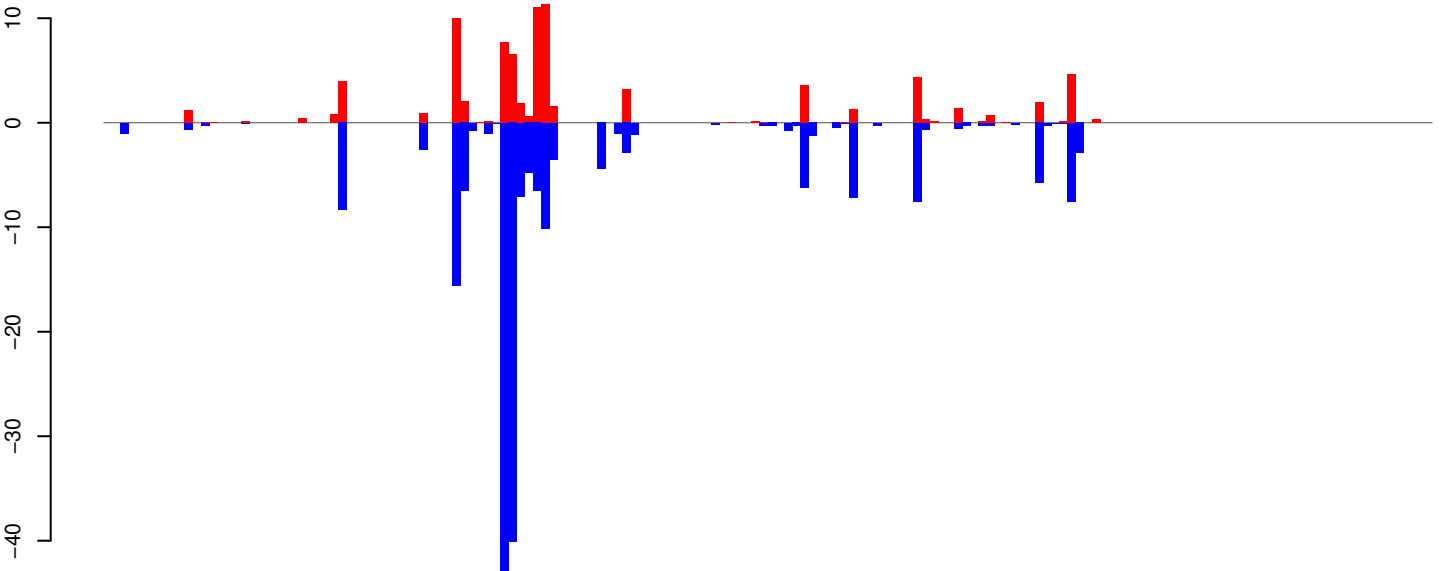
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



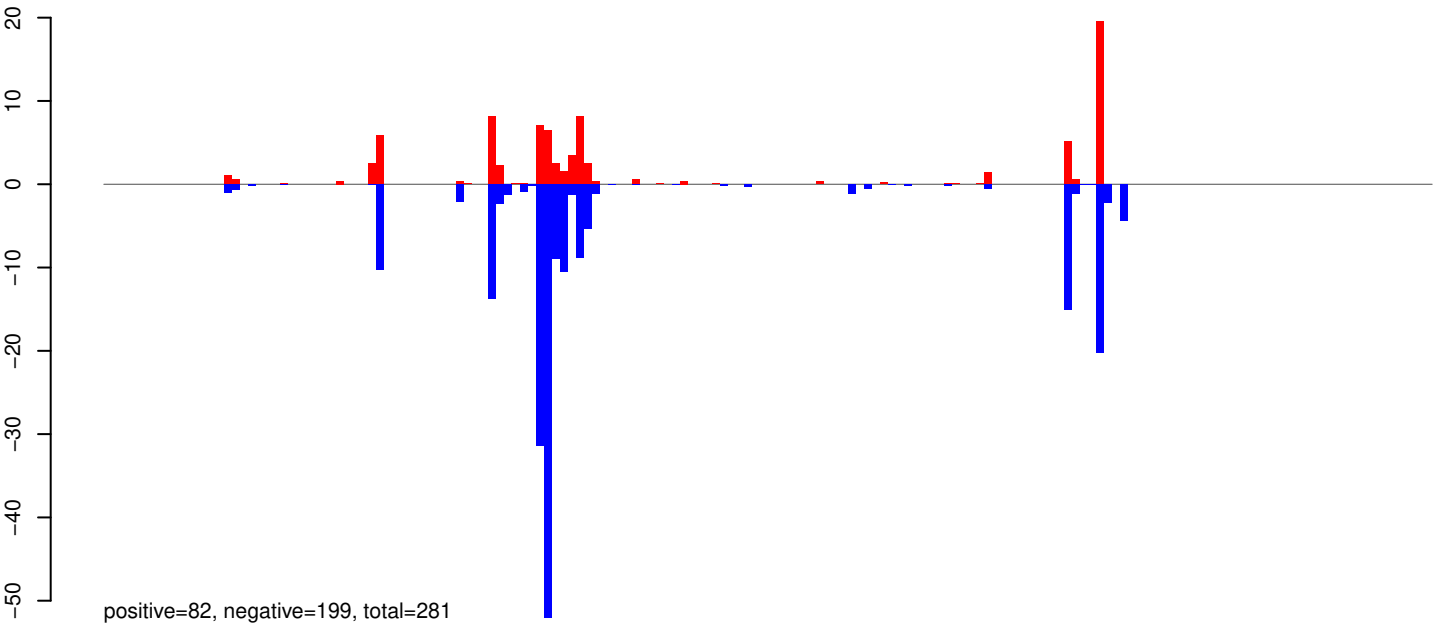
AeAeg_Aag2_mock_dsFluc_RA.rep



bow size=35, length=4094, coverage=268, AY092351.2-Aedes albopictus_C6_36_cell_densovirus_non-structural_protein_1_NS-1_non-structural:1-4094

0 1000 2000 3000 4000

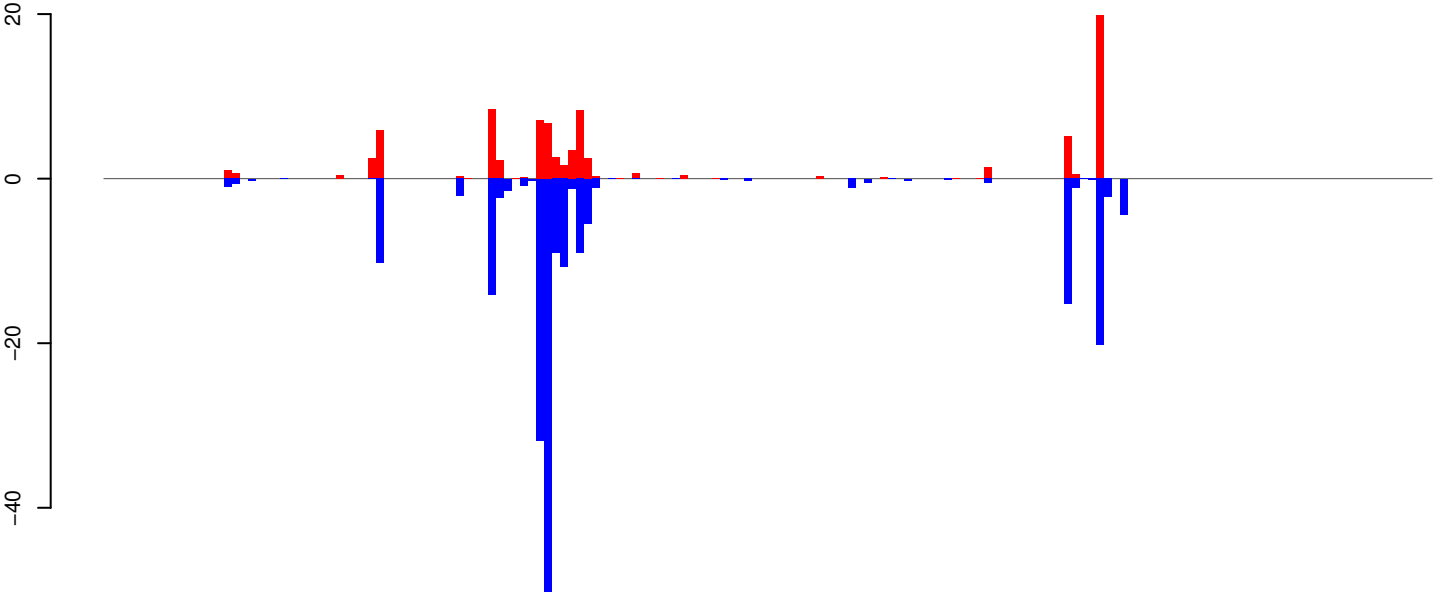
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



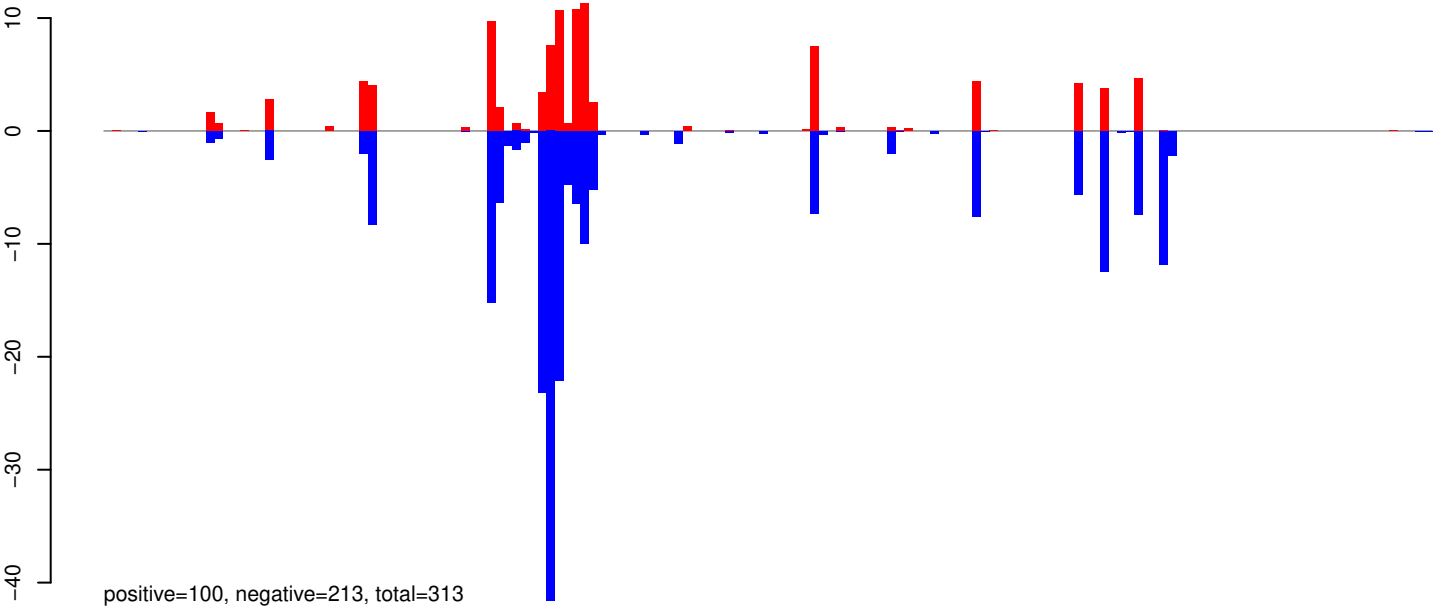
AeAeg_Aag2_mock_dsFluc_RA.rep



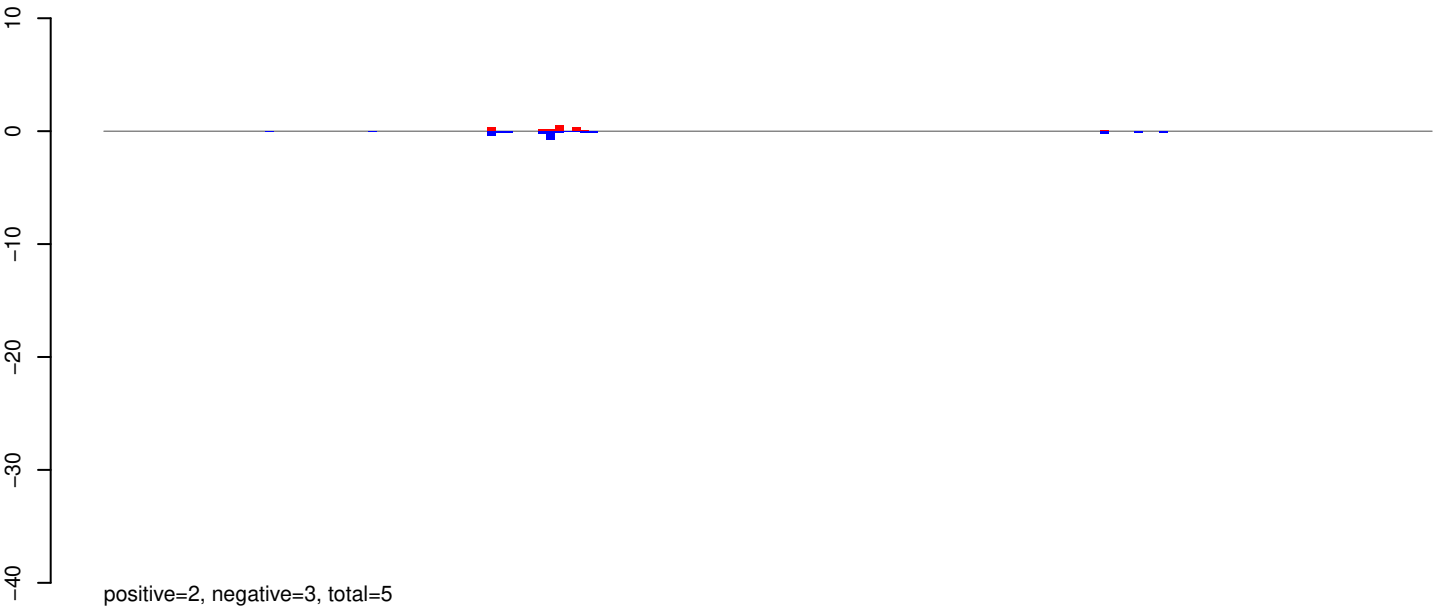
Window size=25, length=4139, virus@NC_011317.1-Anopheles_gambiae_densonucleosis_virus:1-4139

0 1000 2000 3000 4000

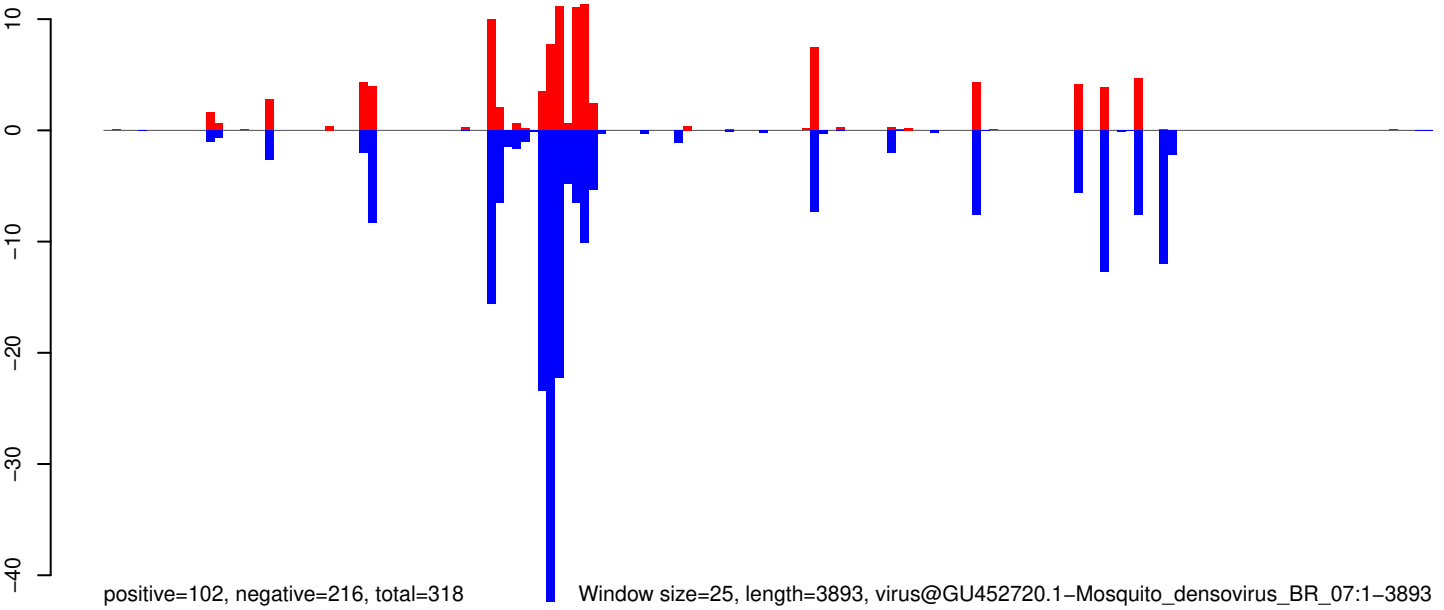
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



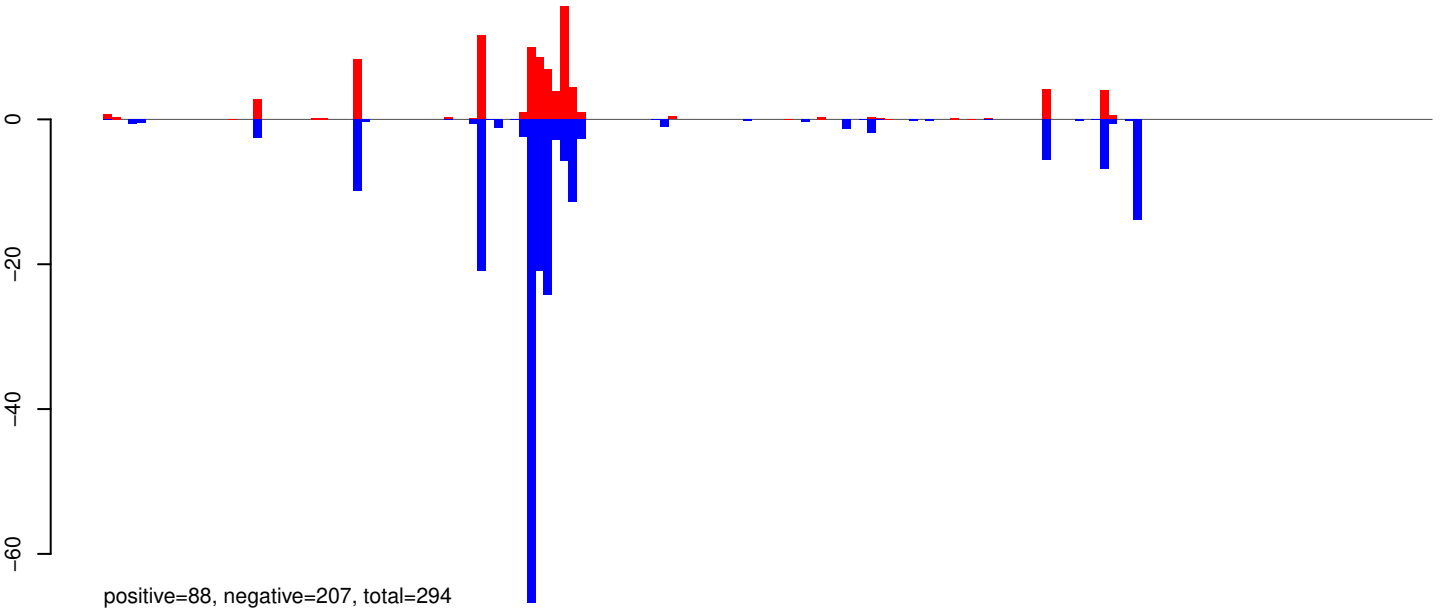
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



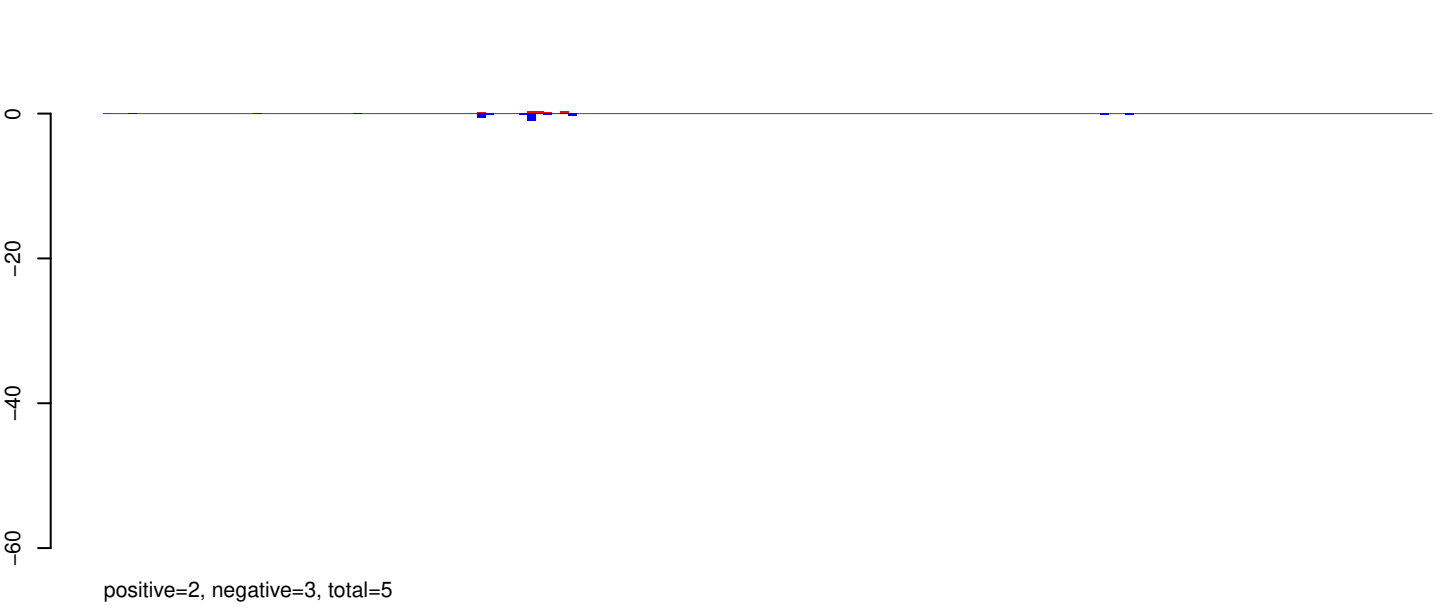
AeAeg_Aag2_mock_dsFluc_RA.rep



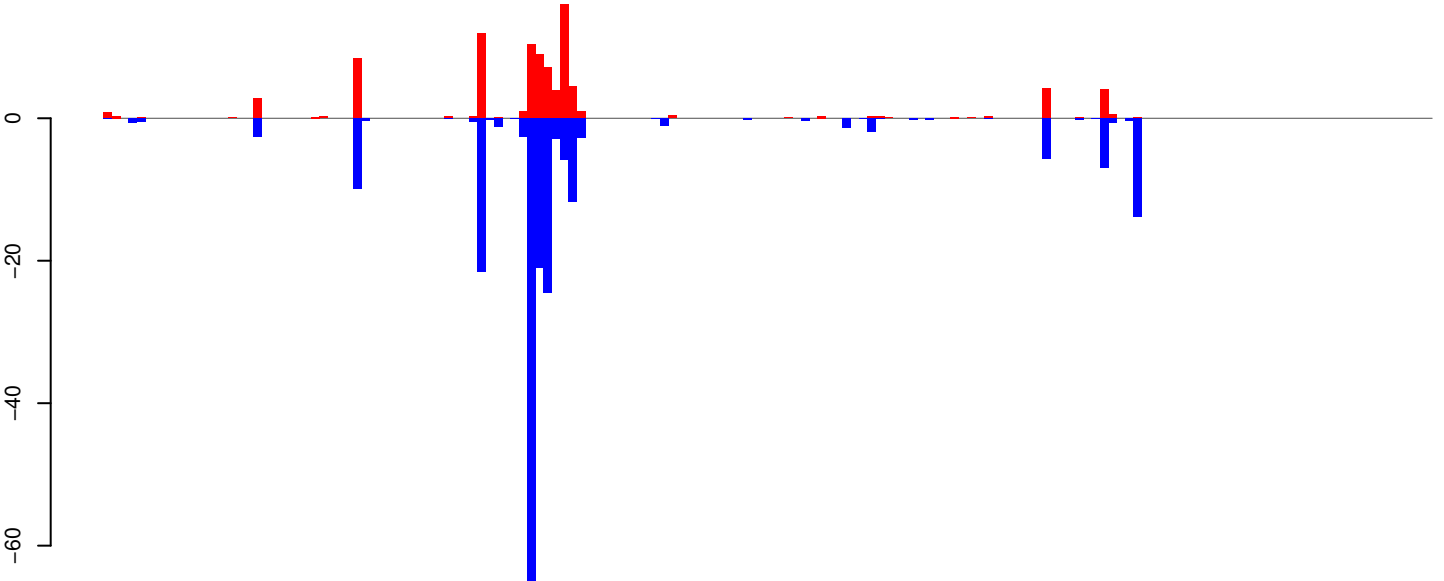
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



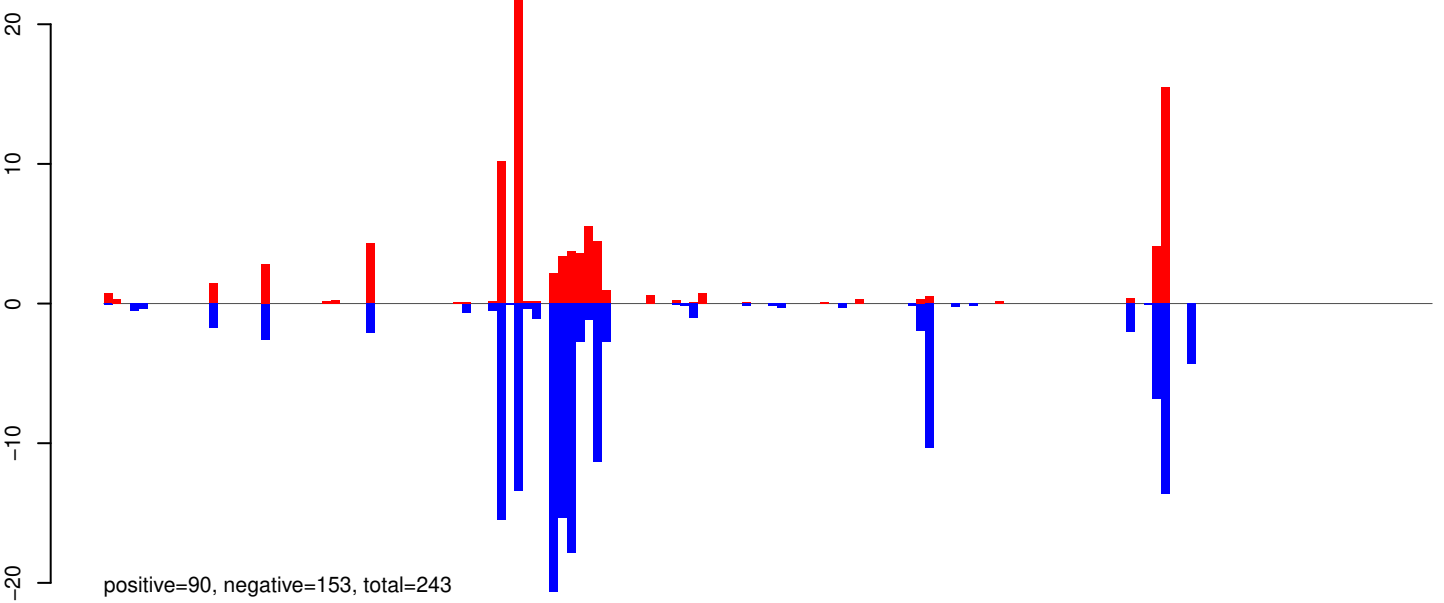
AeAeg_Aag2_mock_dsFluc_RA.rep



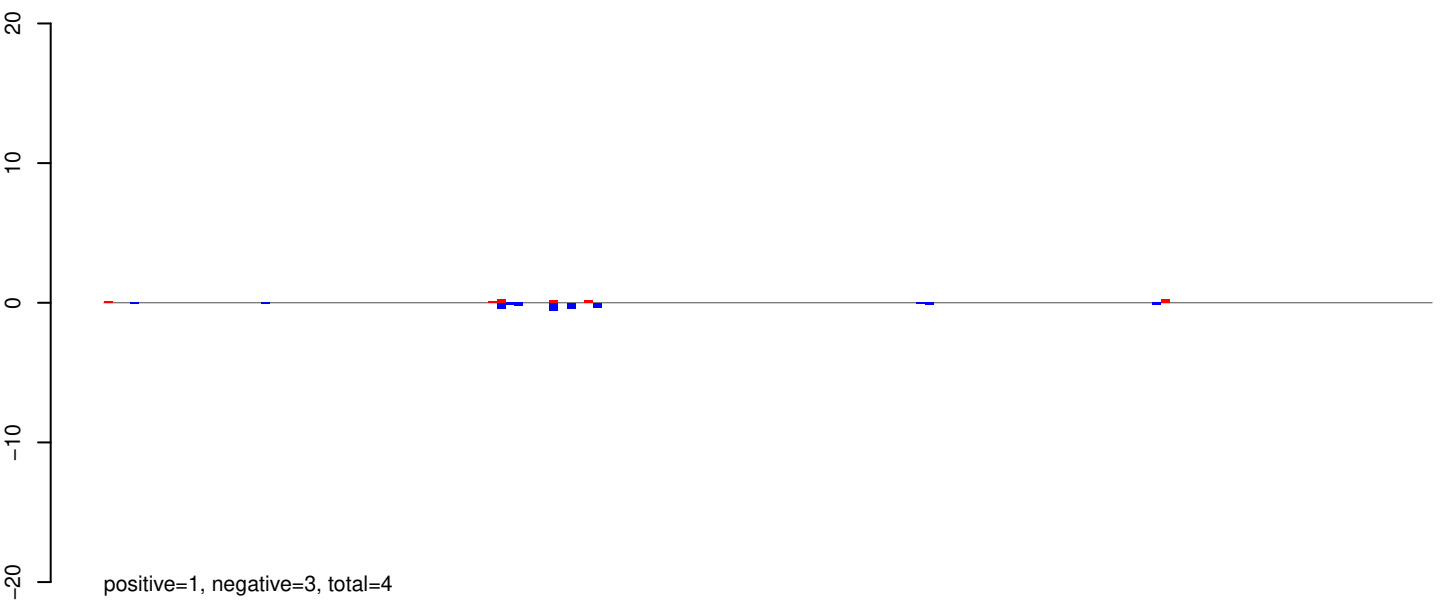
low size=25, length=3978, virus@M37899.1-Aedes_aegypti_densovirus_1_genomic_sequence:1-3978

0 1000 2000 3000 4000

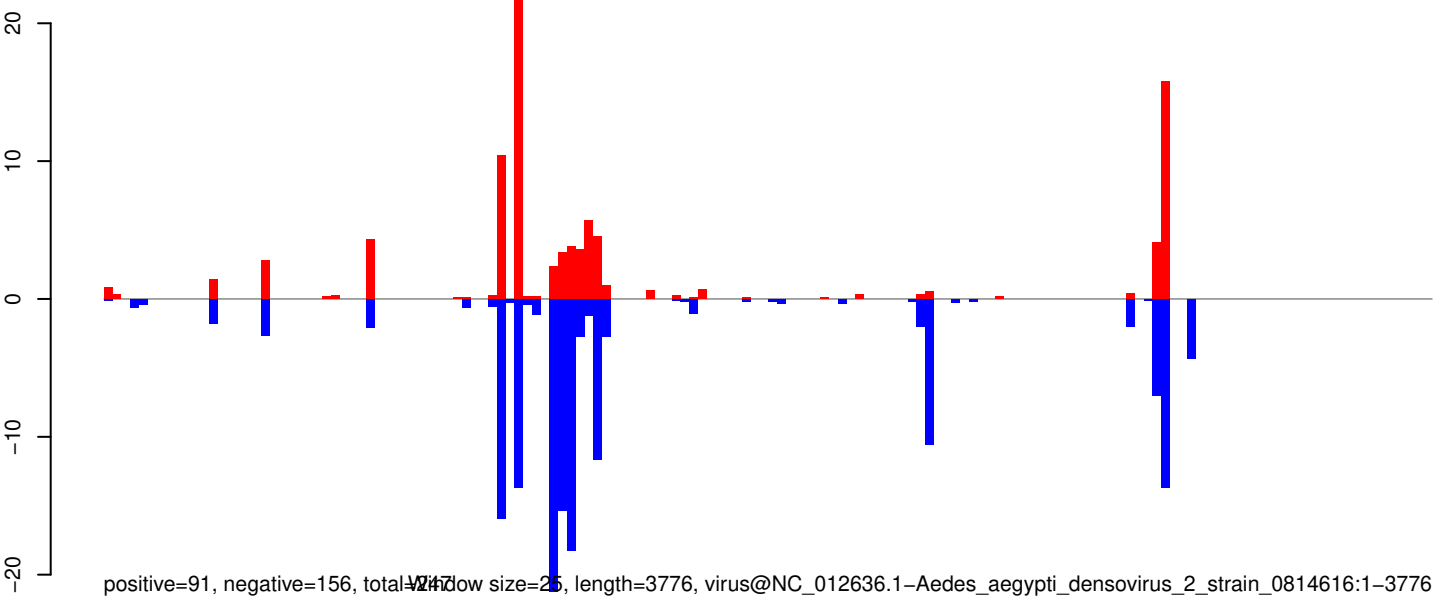
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

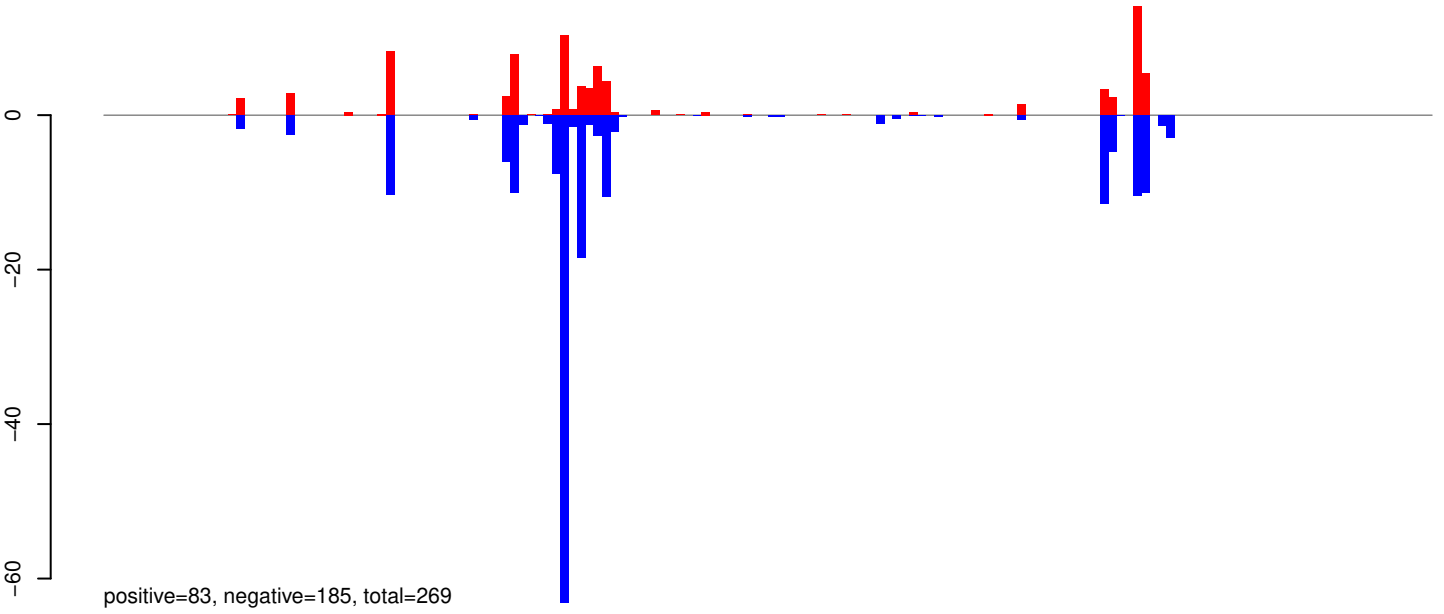


AeAeg_Aag2_mock_dsFluc_RA.rep



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

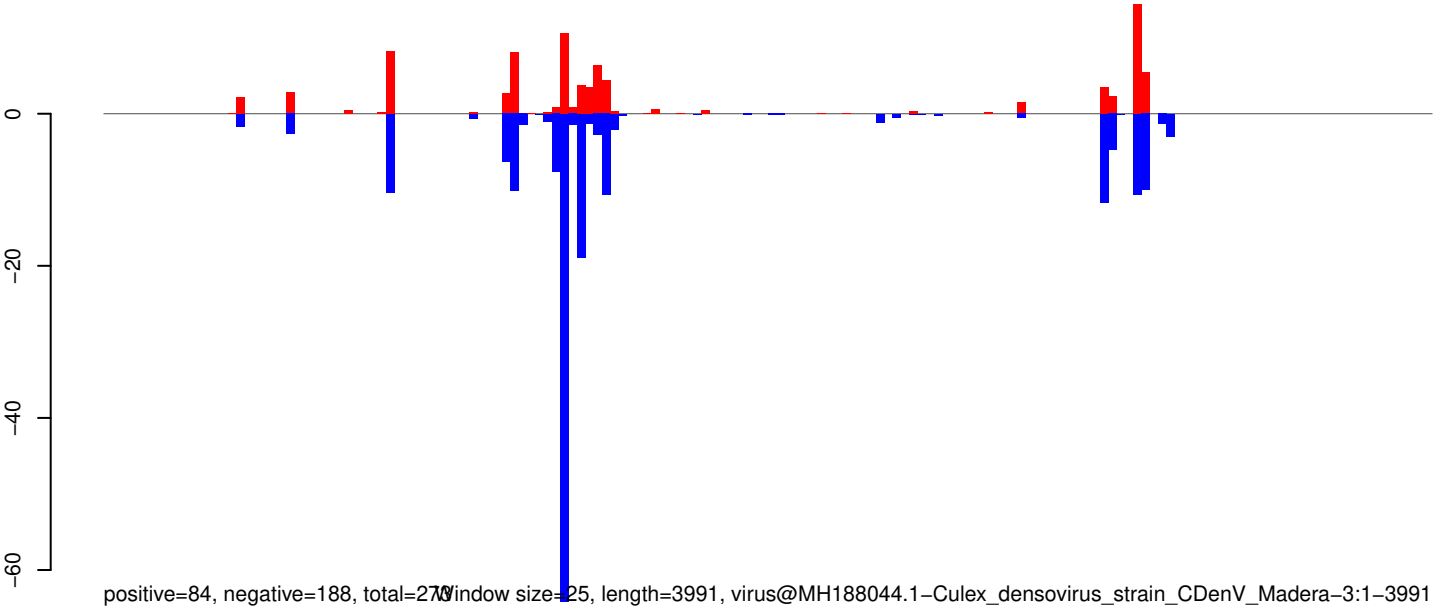
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



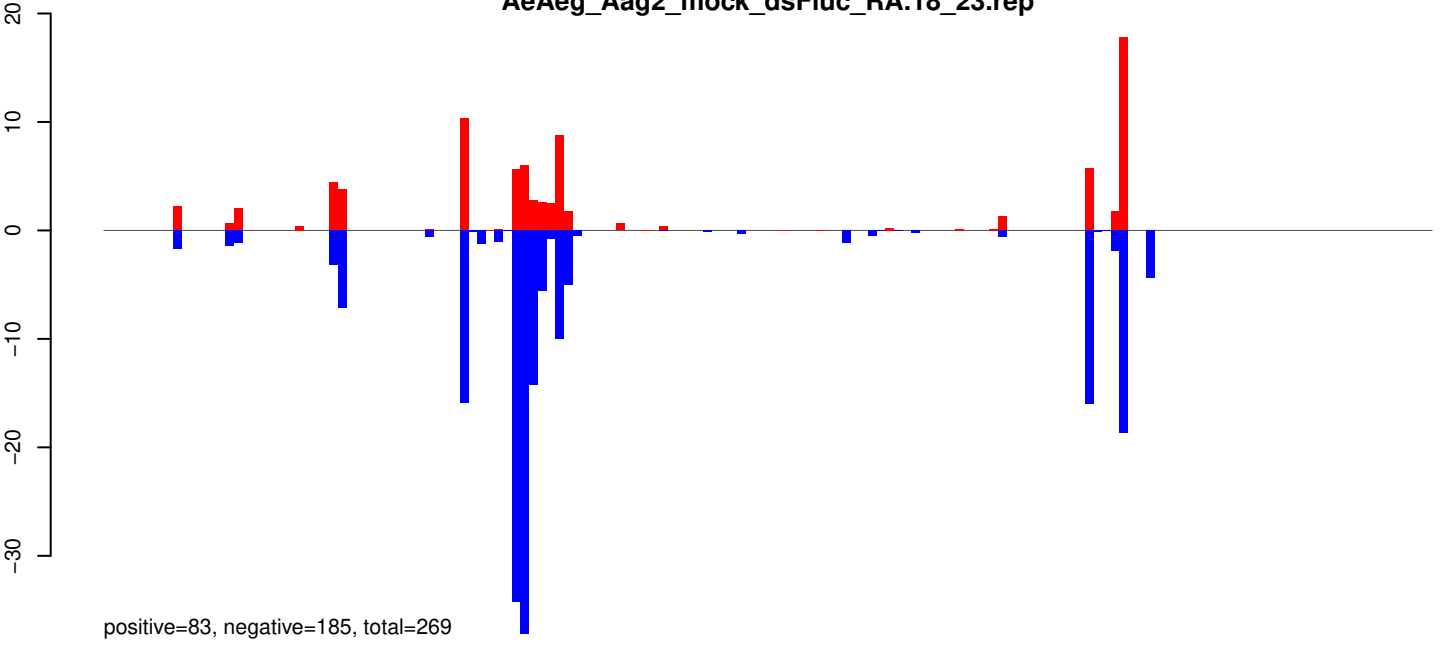
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep

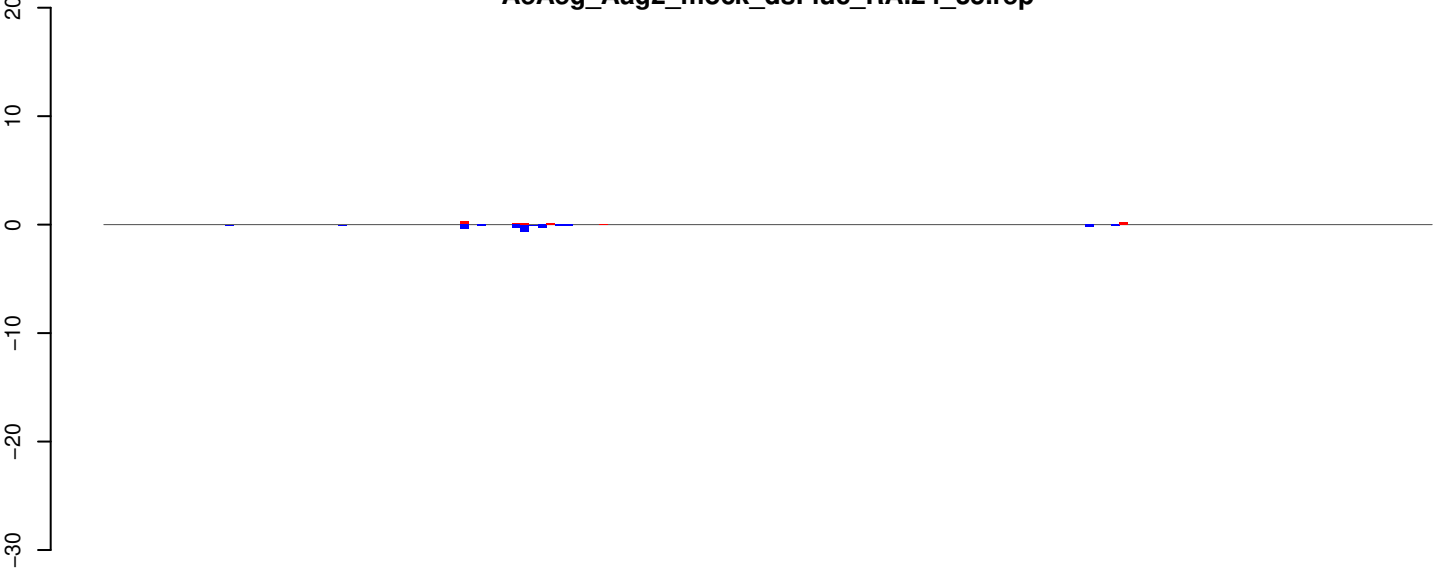


AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



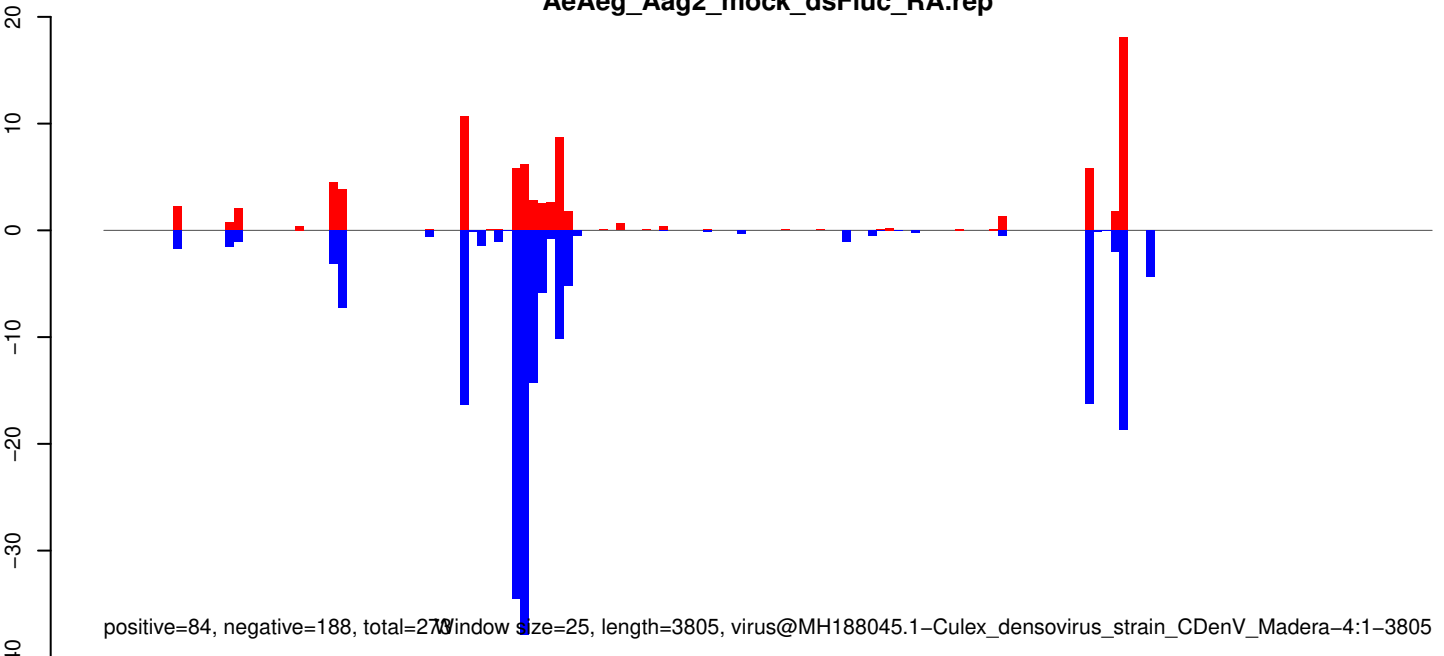
positive=83, negative=185, total=269

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



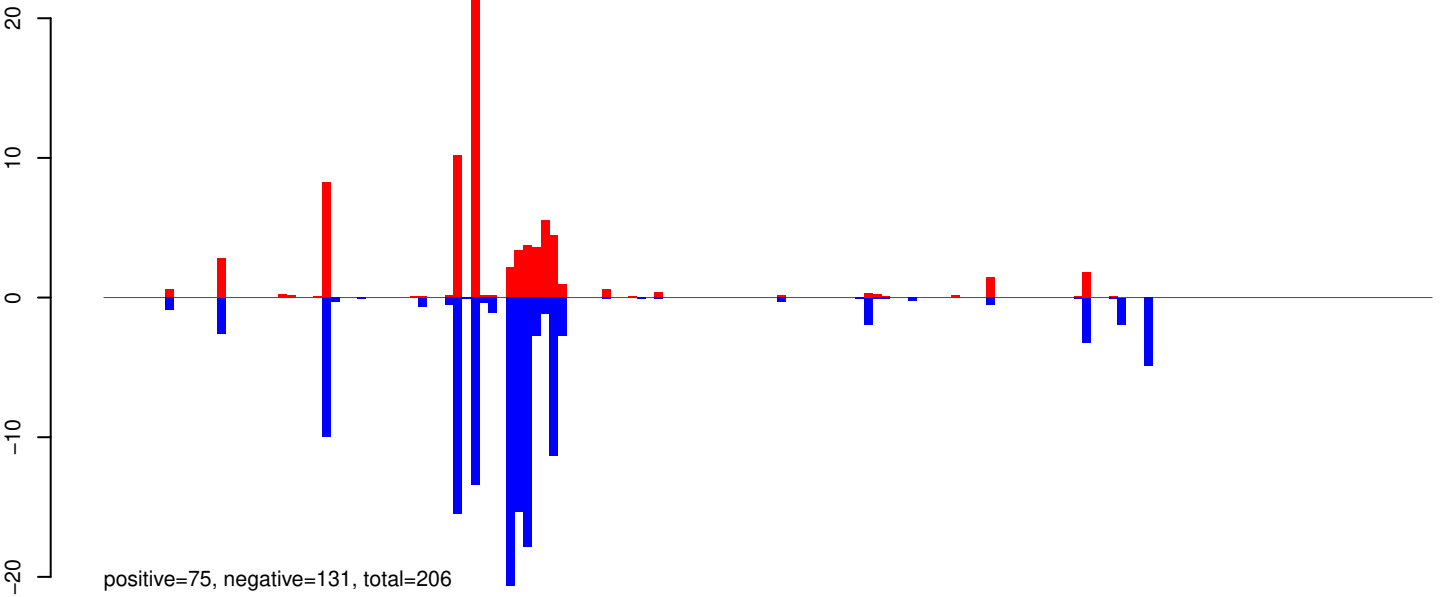
positive=1, negative=3, total=4

AeAeg_Aag2_mock_dsFluc_RA.rep

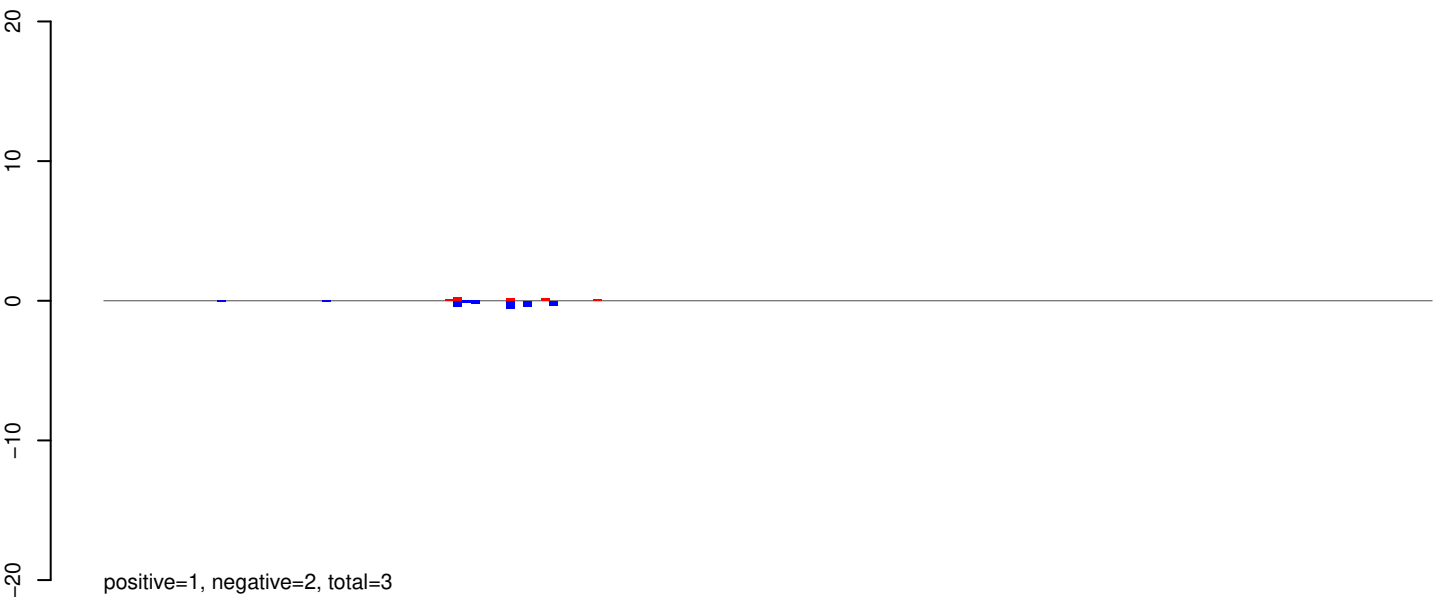


positive=84, negative=188, total=272, window size=25, length=3805, virus@MH188045.1-Culex_densovirus_strain_CDenV_Madera-4:1-3805

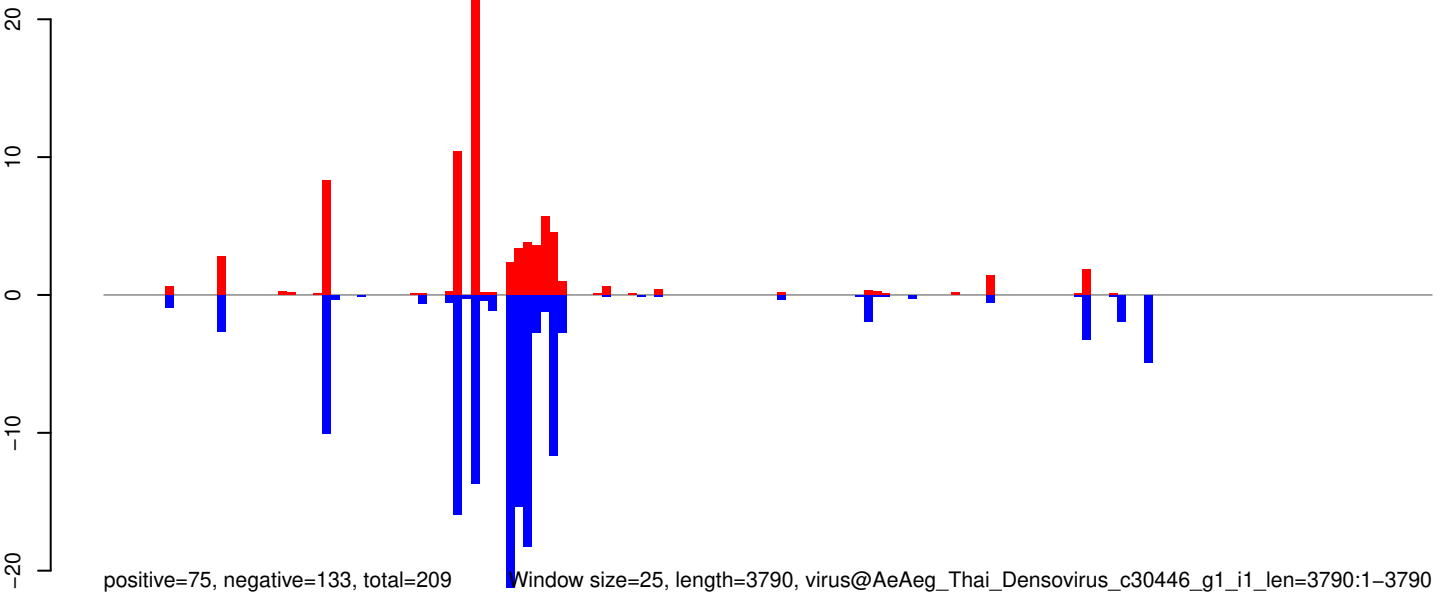
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



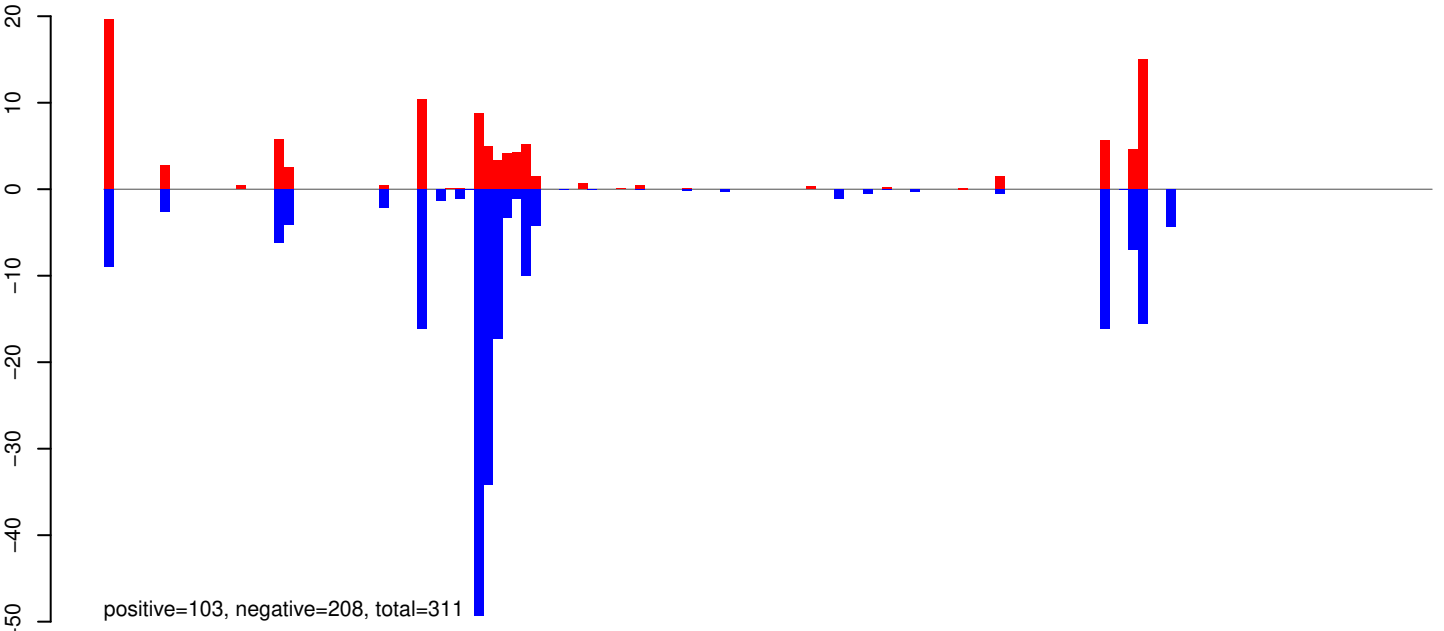
AeAeg_Aag2_mock_dsFluc_RA.rep



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

0 1000 2000 3000

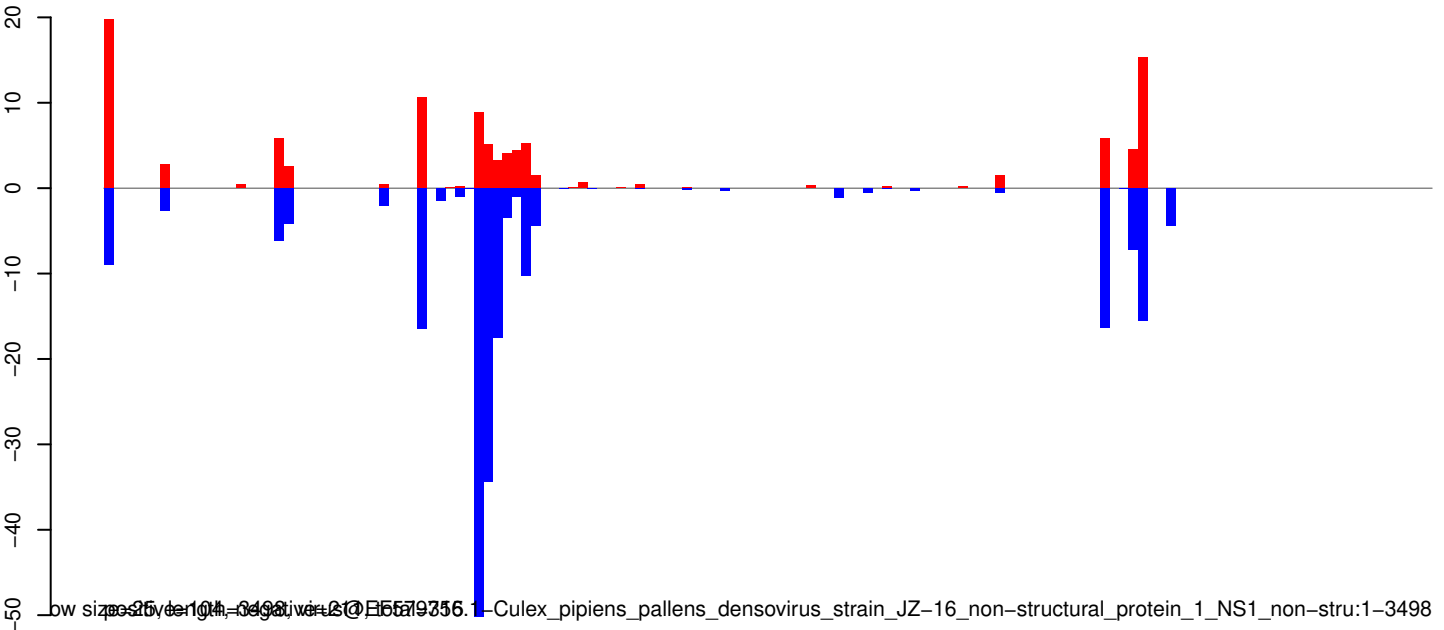
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



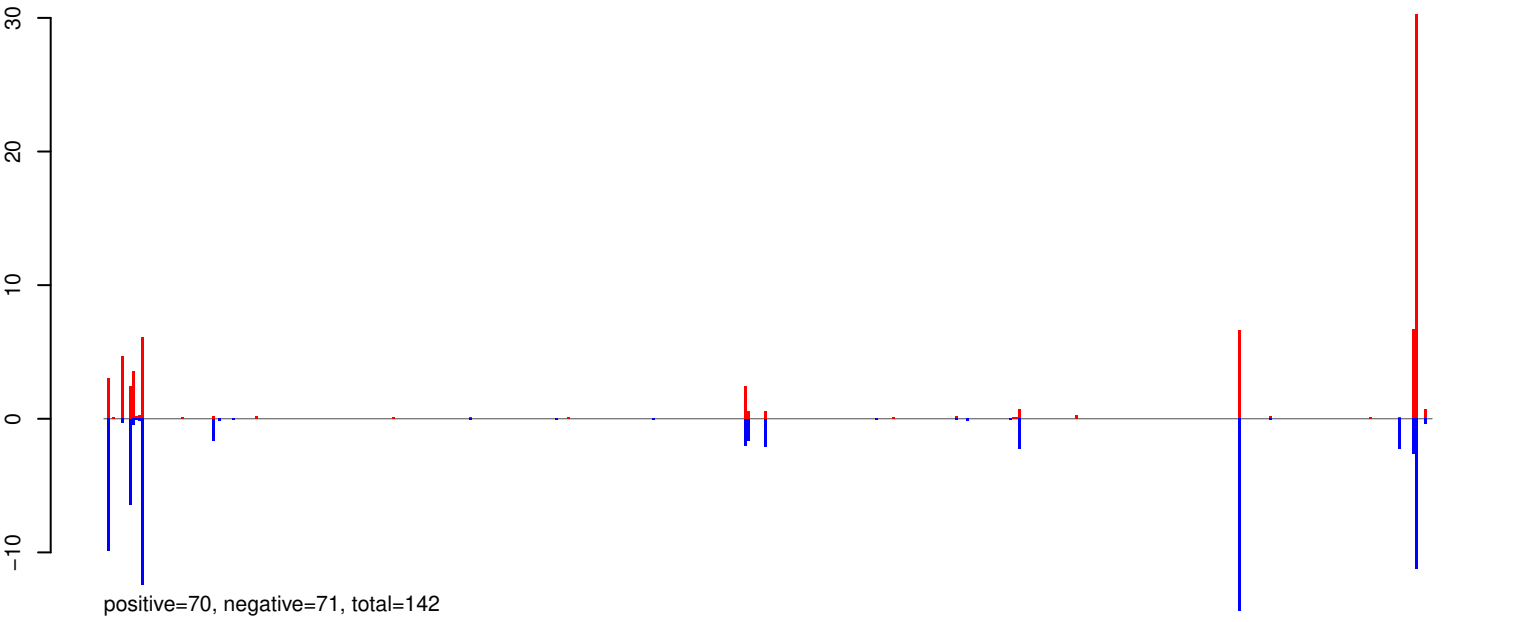
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep



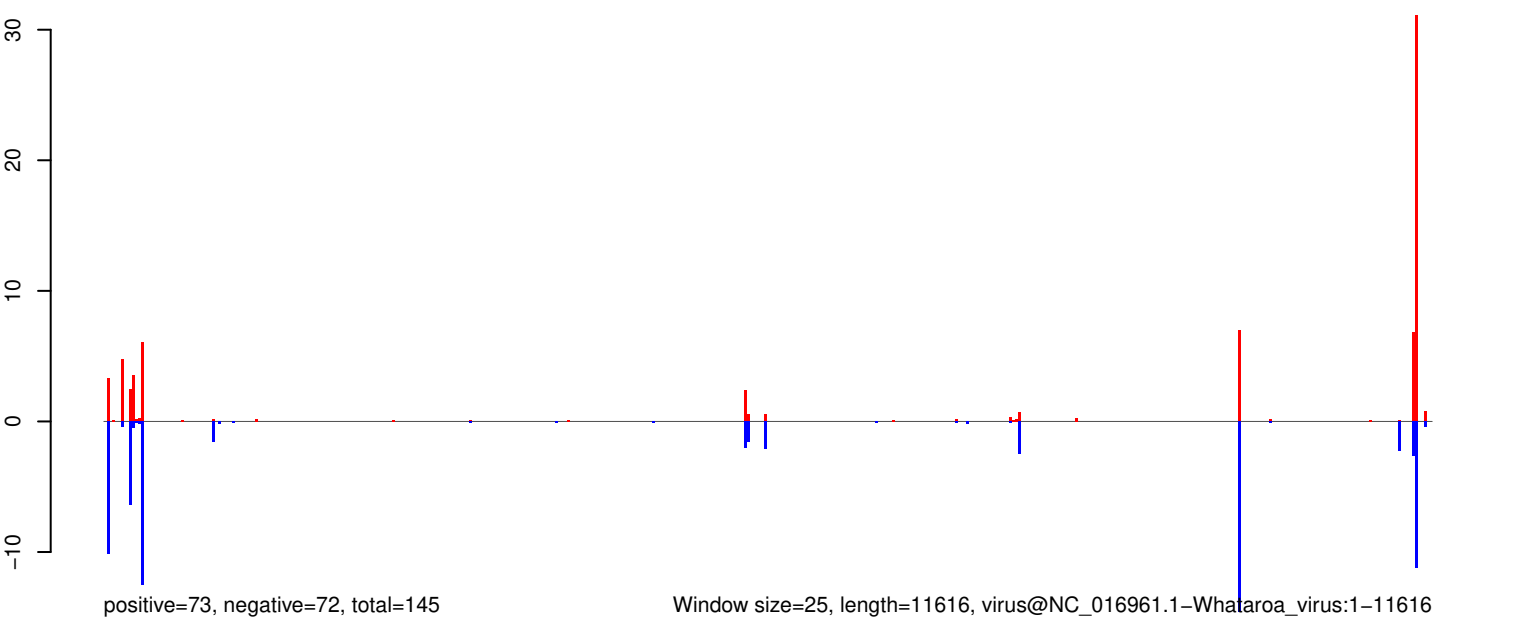
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



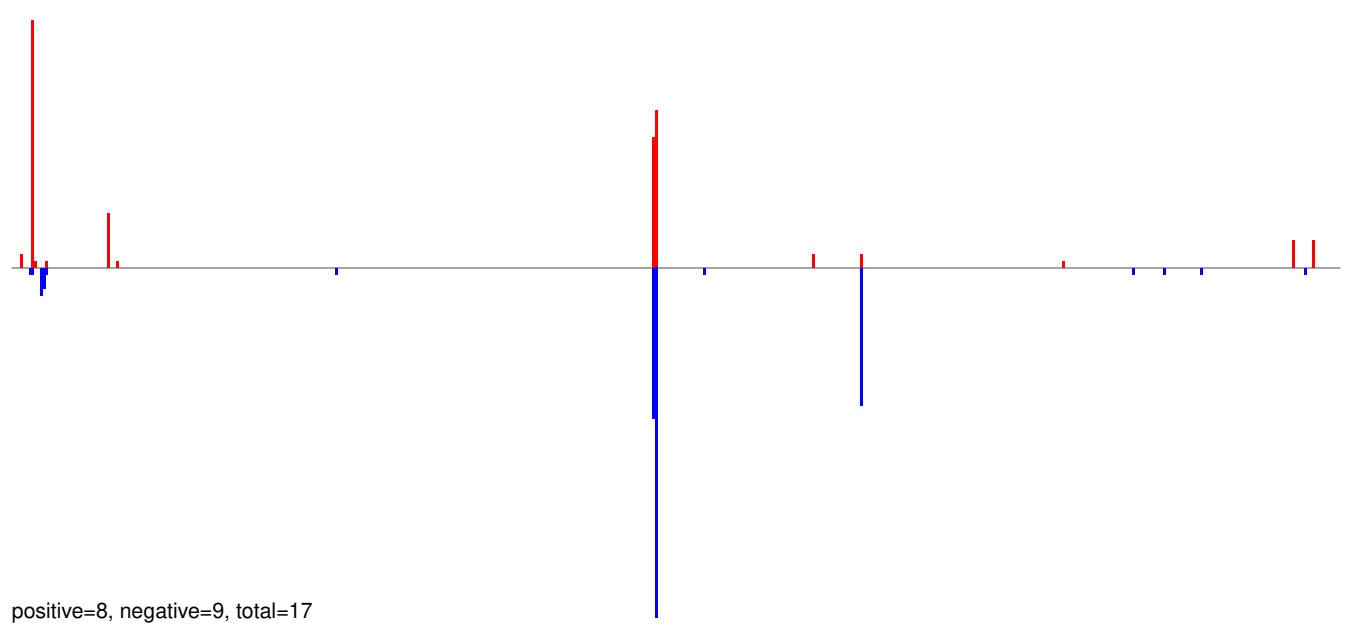
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



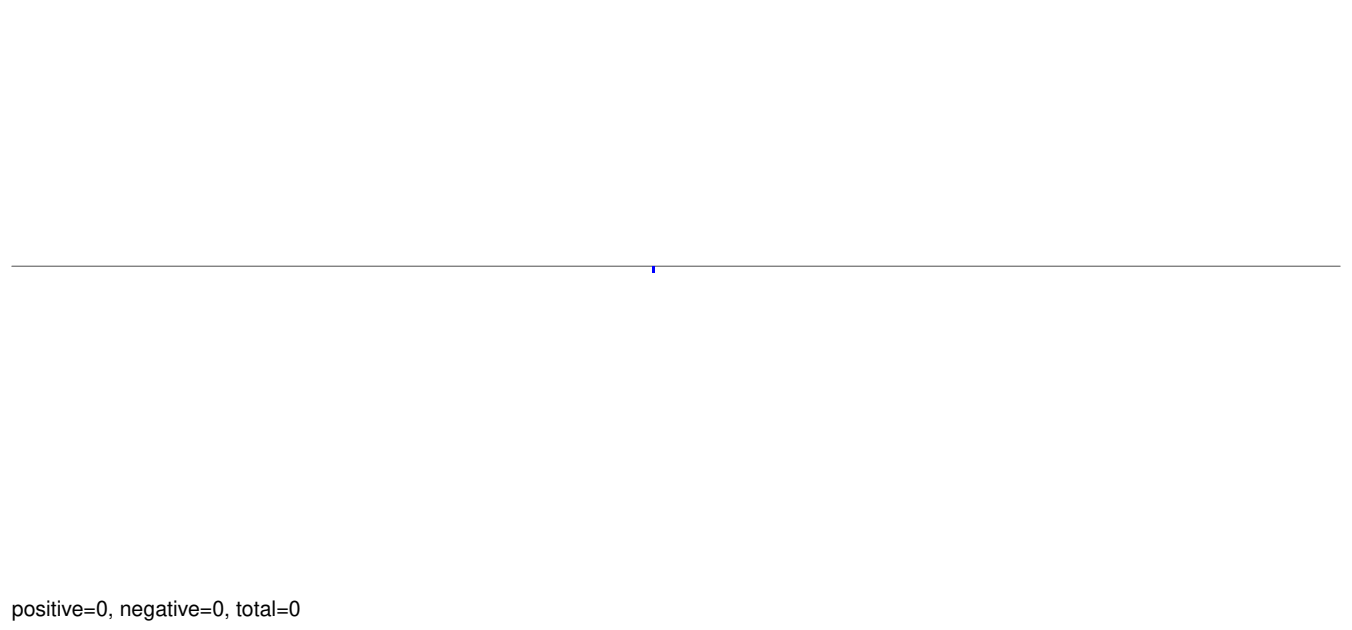
AeAeg_Aag2_mock_dsFluc_RA.rep



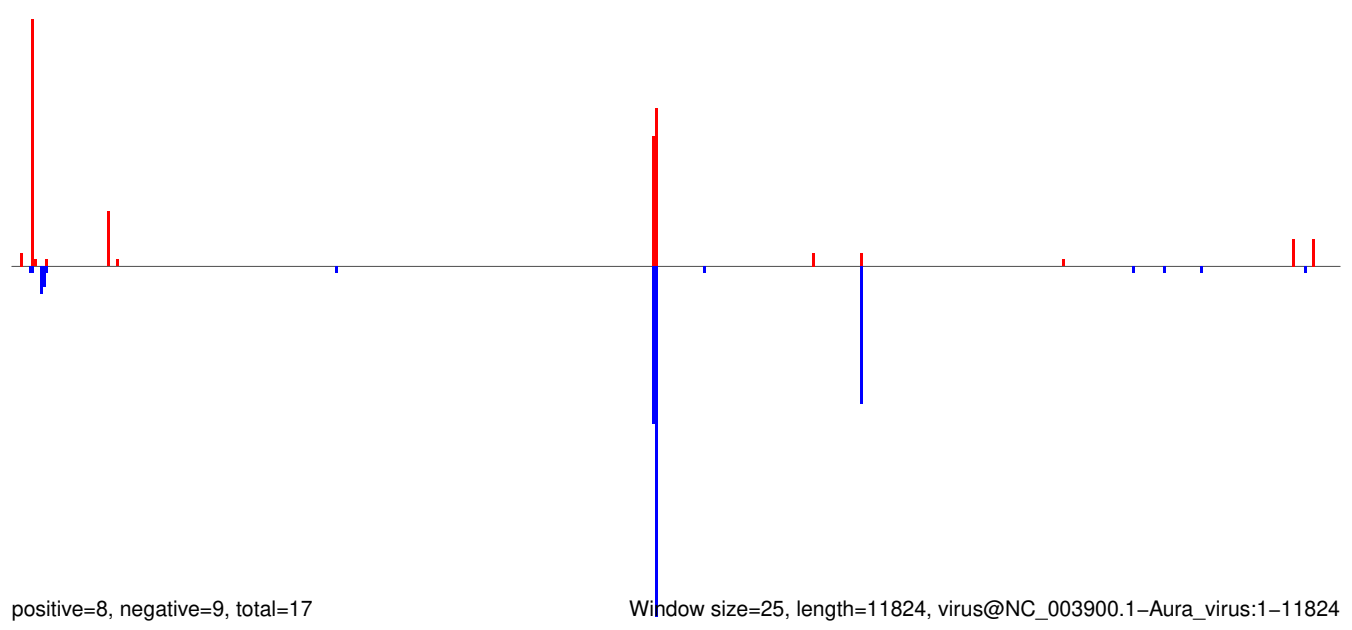
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep



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

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep

positive=12, negative=9, total=21

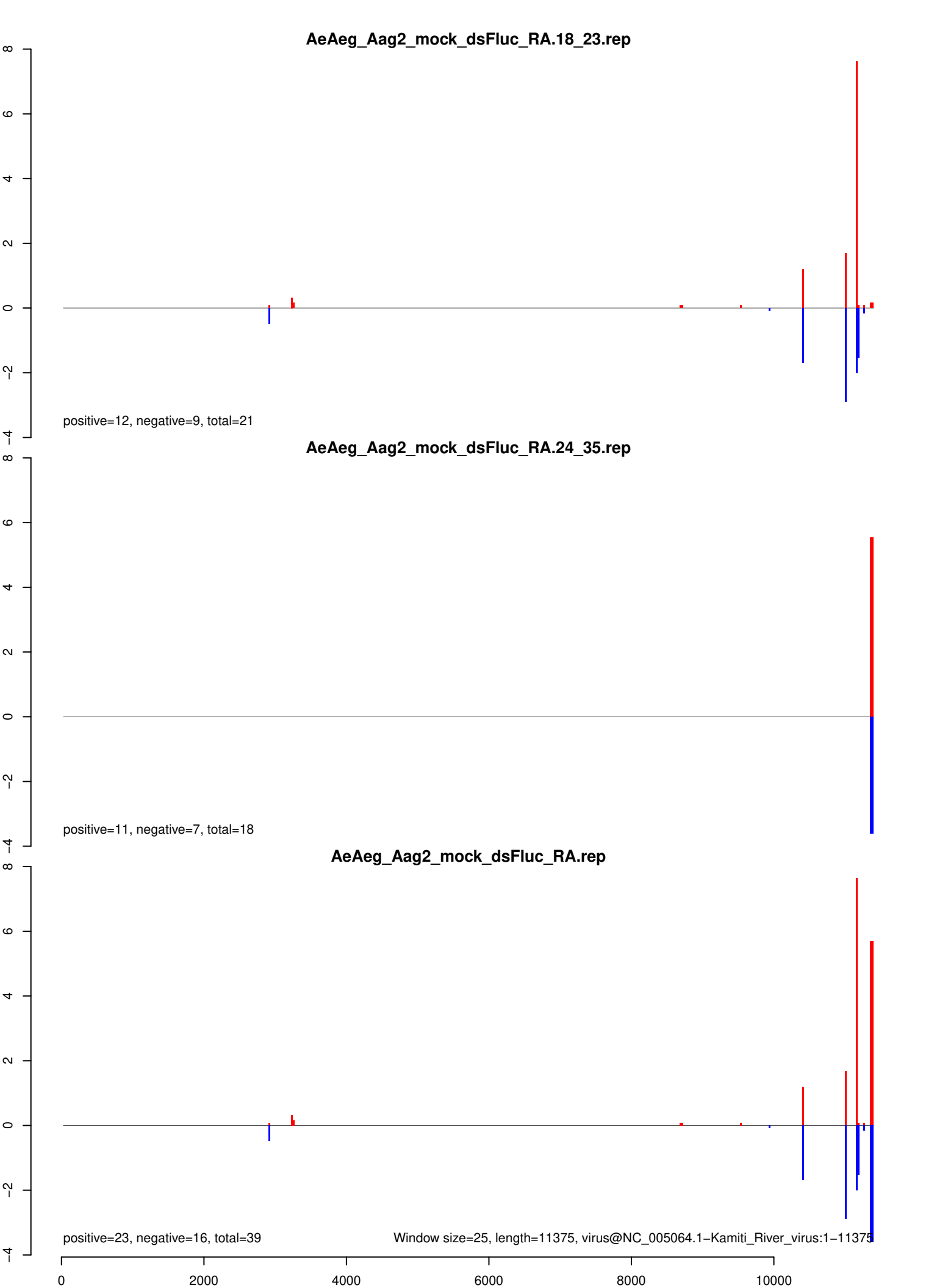
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

positive=11, negative=7, total=18

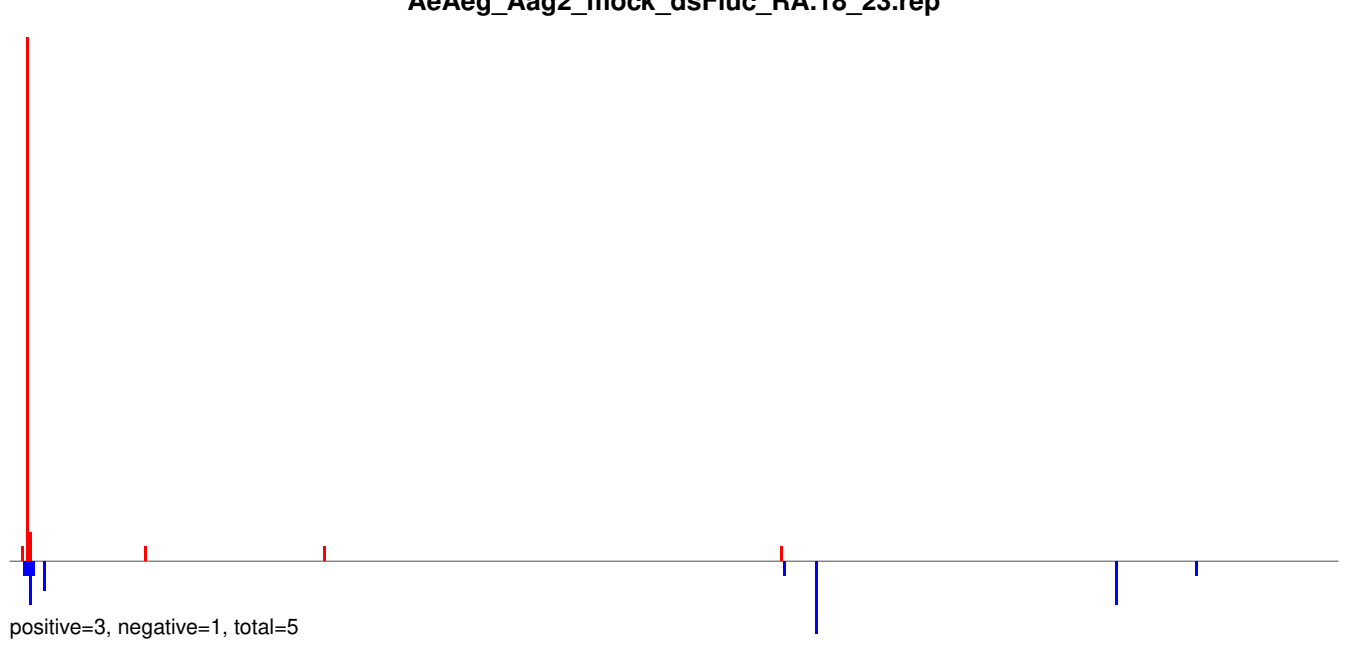
AeAeg_Aag2_mock_dsFluc_RA.rep

positive=23, negative=16, total=39

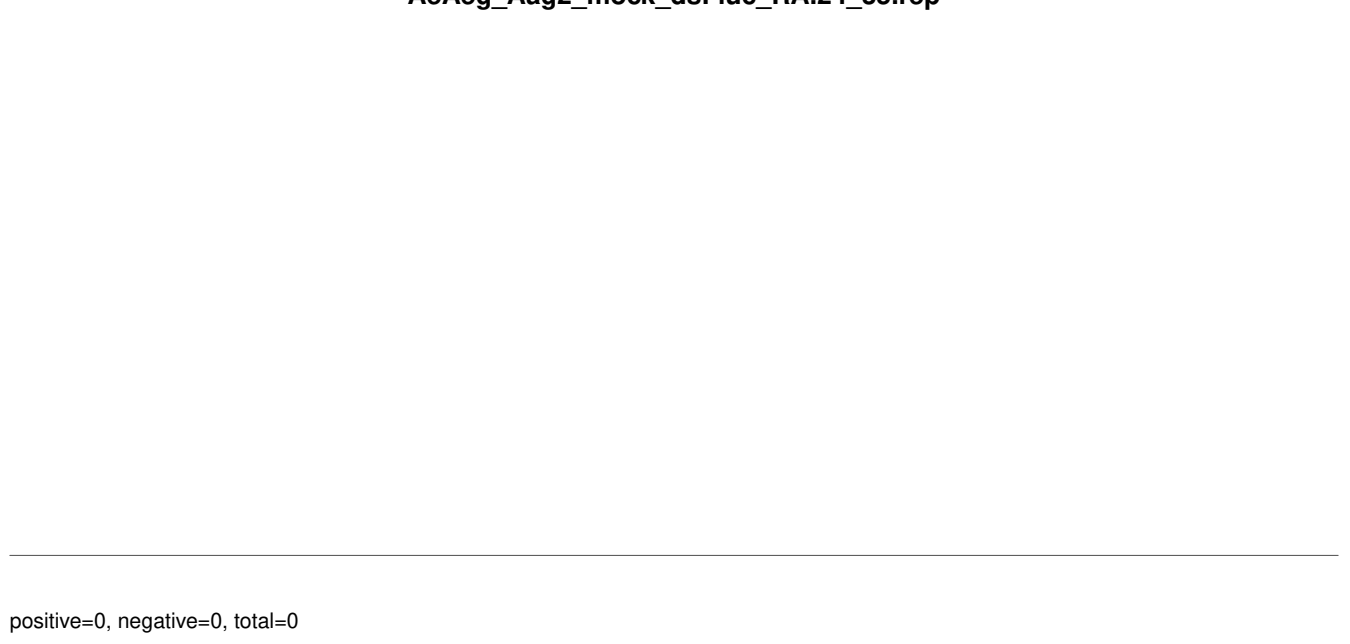
Window size=25, length=11375, virus@NC_005064.1-Kamiti_River_virus:1-11375



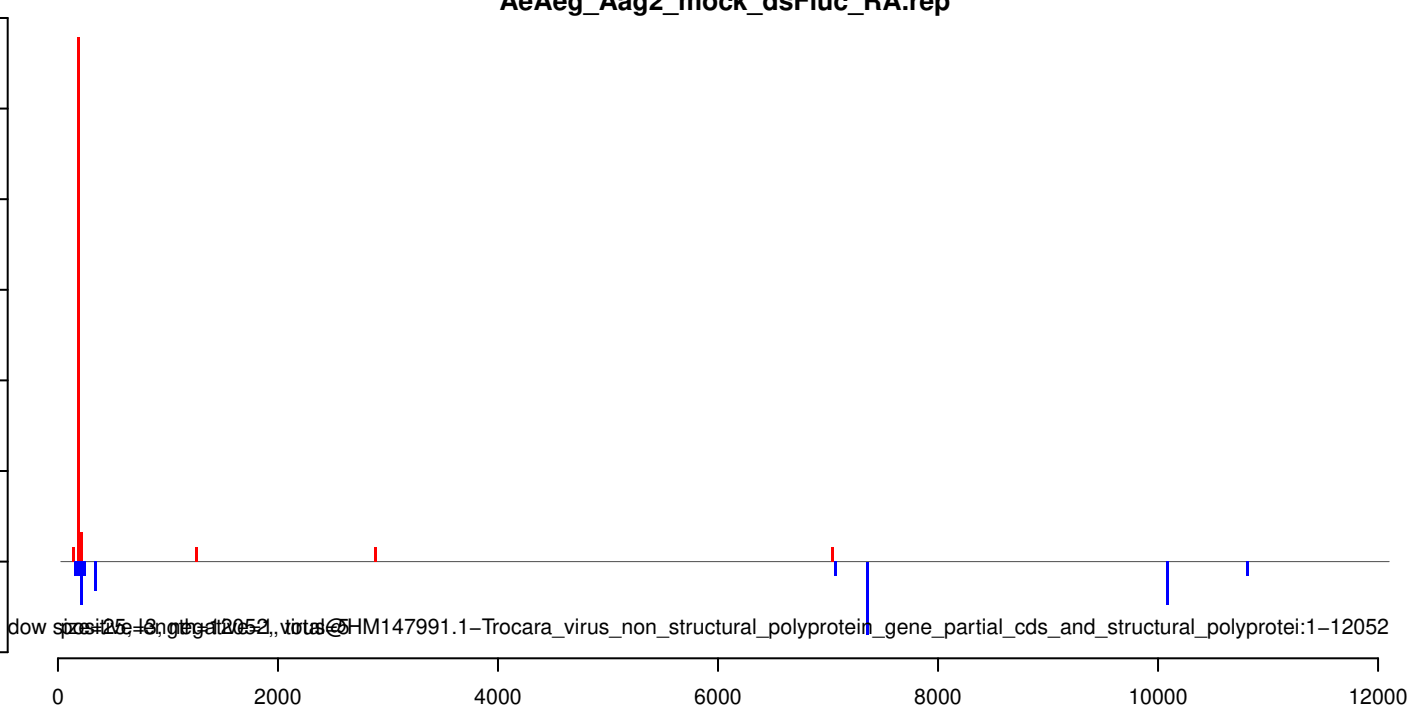
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

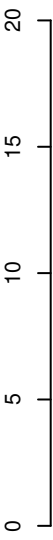


AeAeg_Aag2_mock_dsFluc_RA.rep



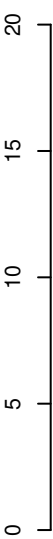
downloaded from SRA on 08/12/2015. Virus: HM147991.1-Trocaravirus non structural polyprotein gene partial cds and structural polyprotein:1-12052

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



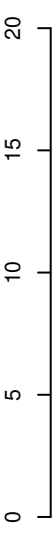
positive=27, negative=6, total=33

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

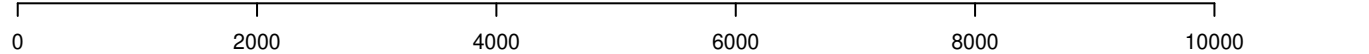


positive=2, negative=0, total=2

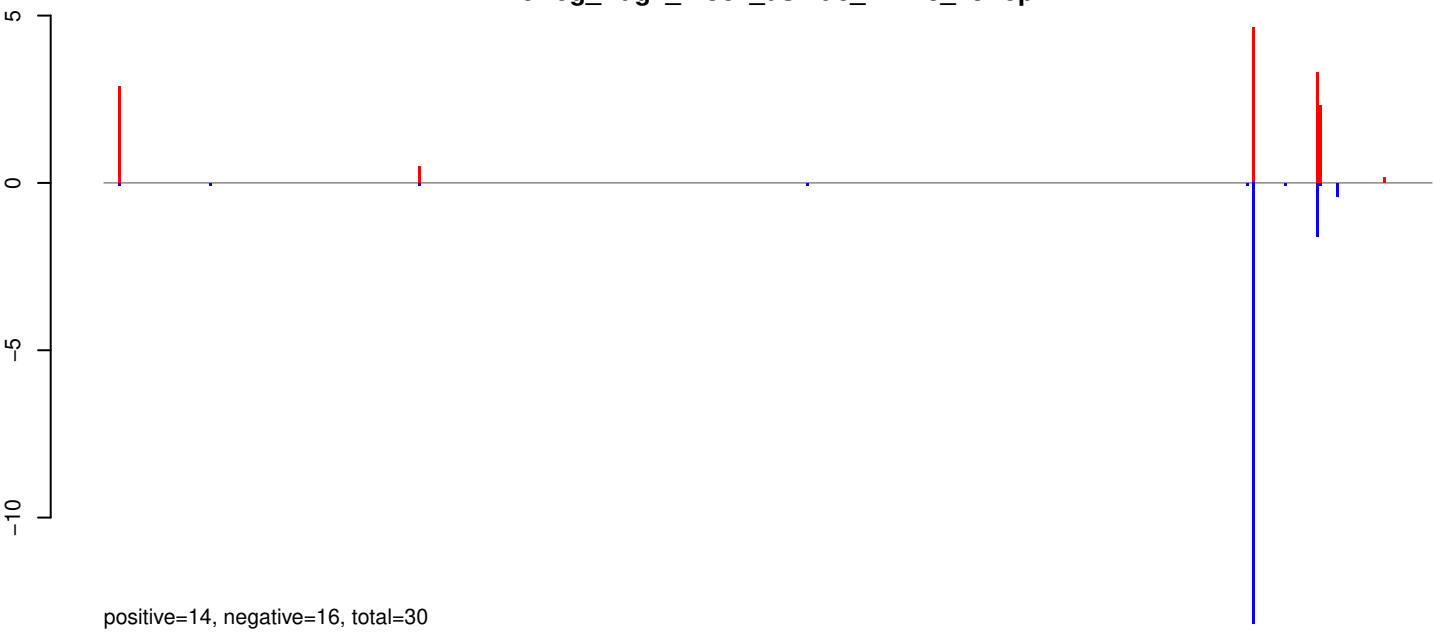
AeAeg_Aag2_mock_dsFluc_RA.rep



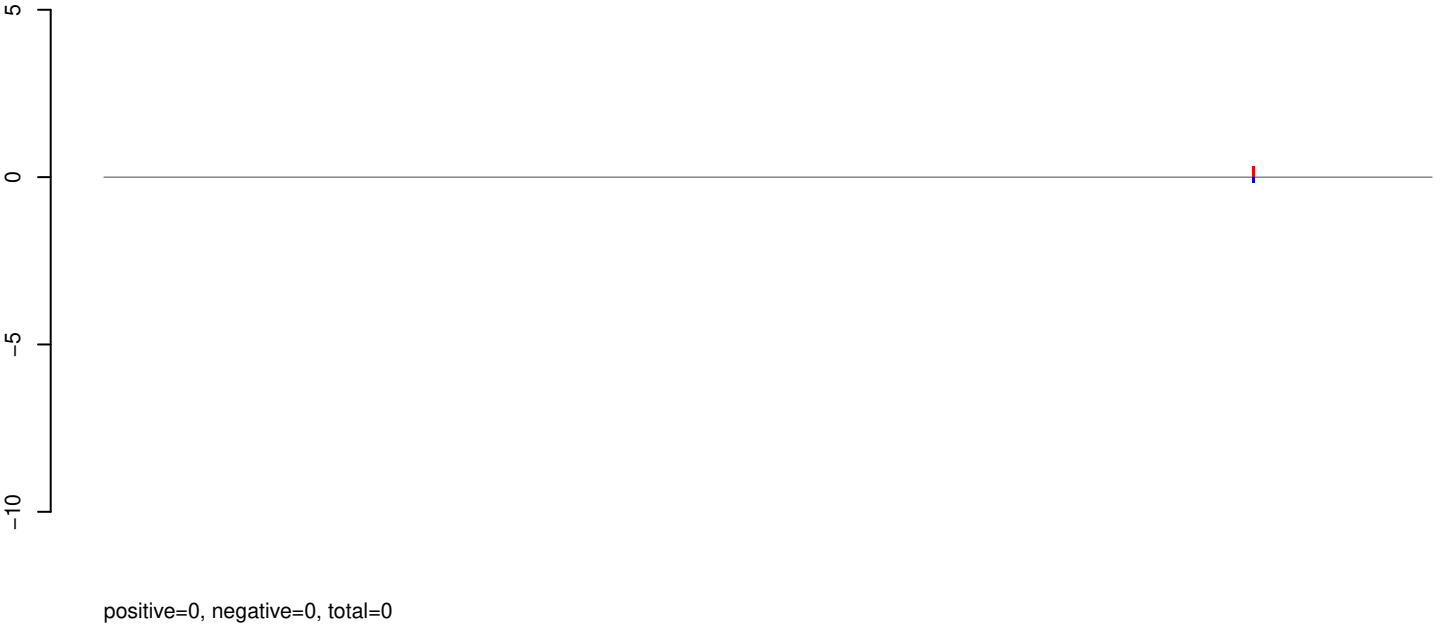
positive=29, negative=6, total=35, length=11079, virus@KC181923.1-Aedes_flavivirus_strain_AEFV-SPFLD-MO-2011-MP6:1-11079



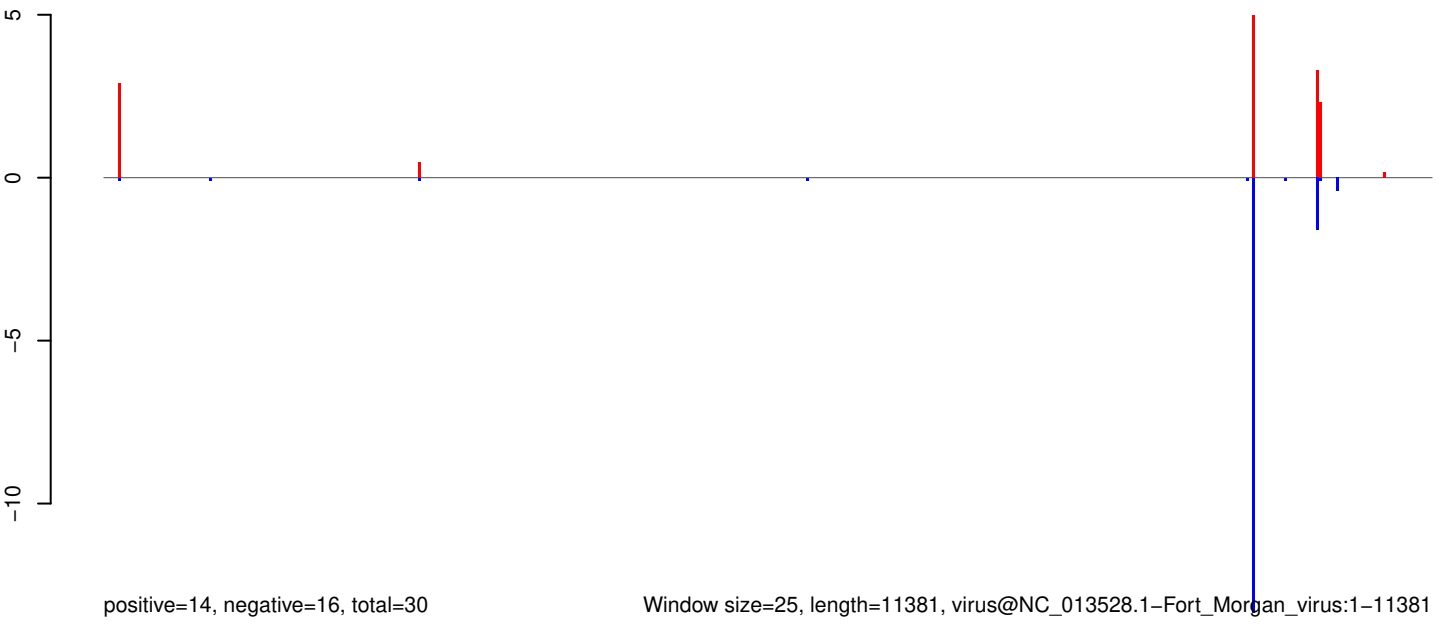
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep



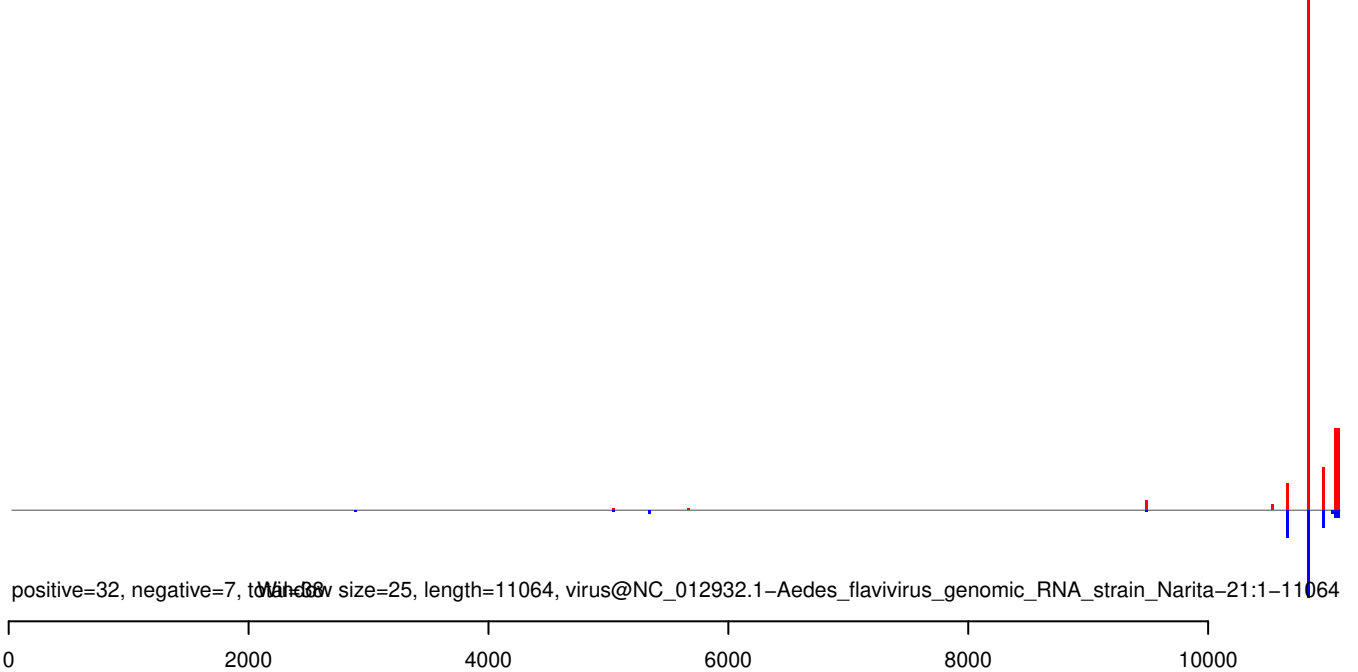
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



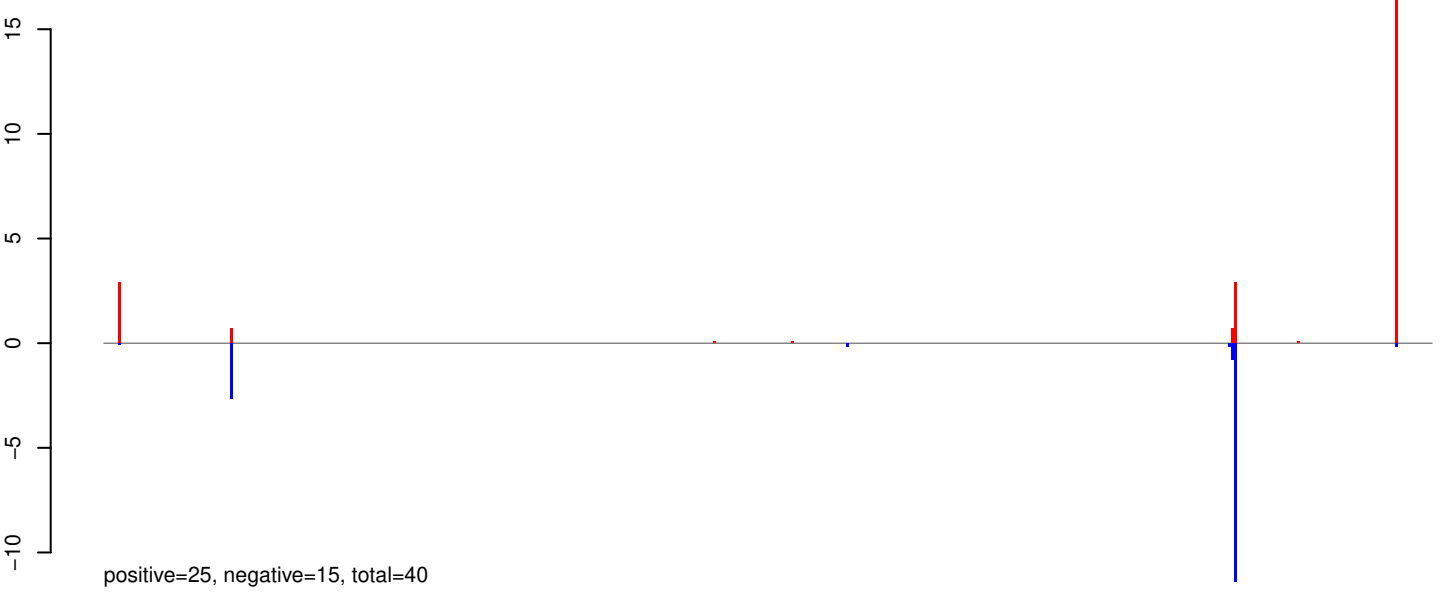
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



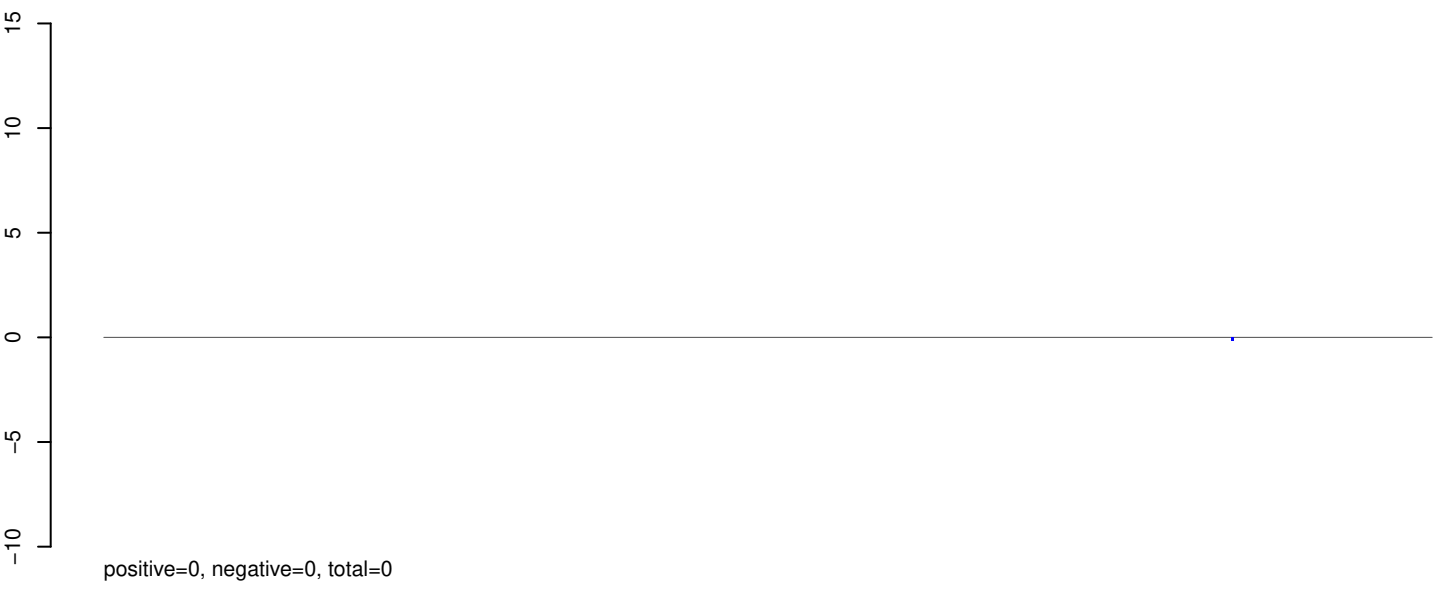
AeAeg_Aag2_mock_dsFluc_RA.rep



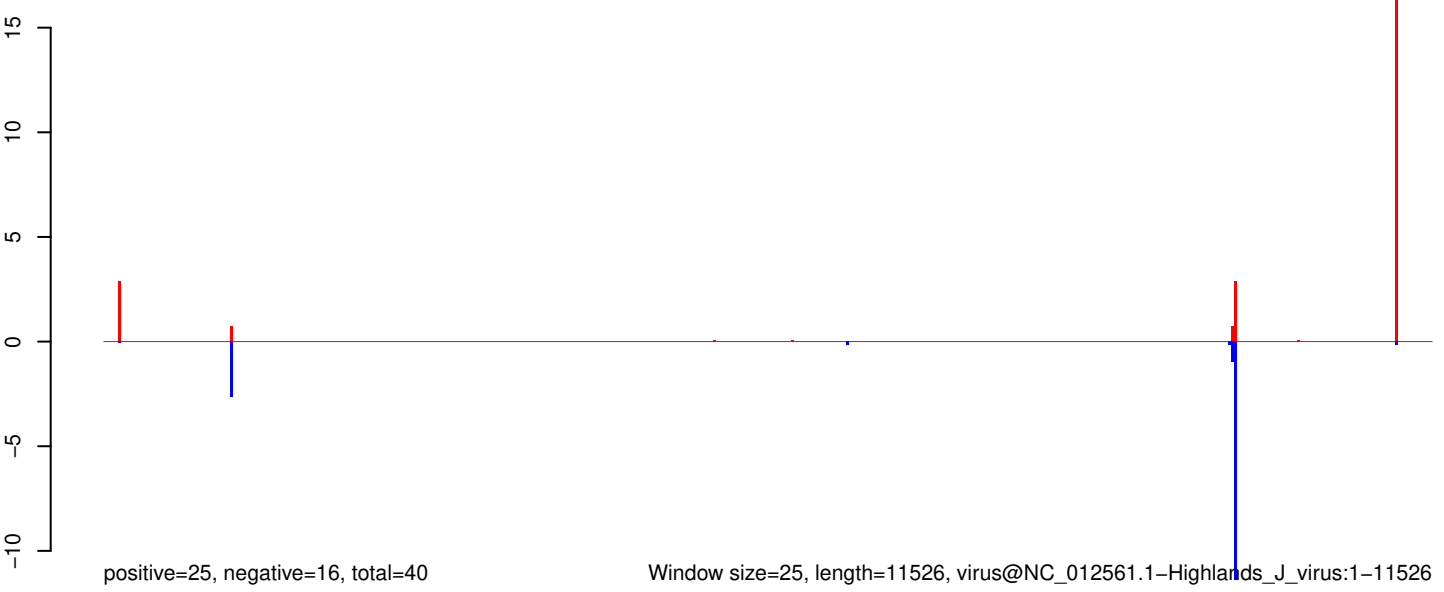
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

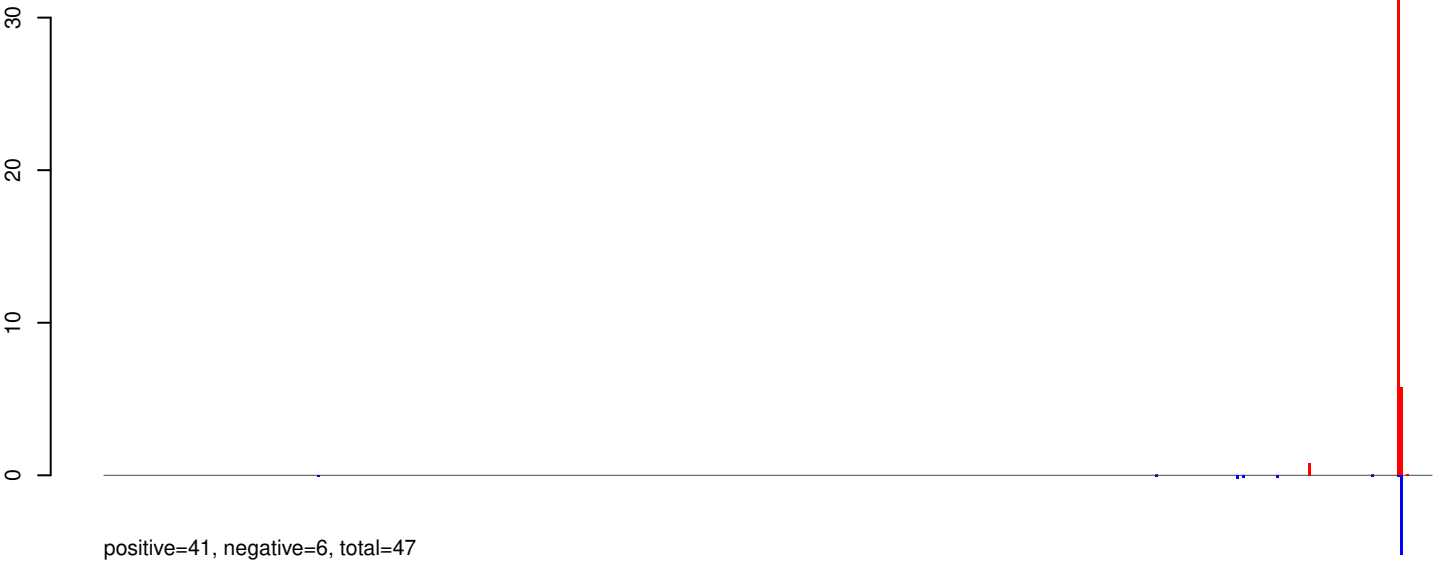


AeAeg_Aag2_mock_dsFluc_RA.rep

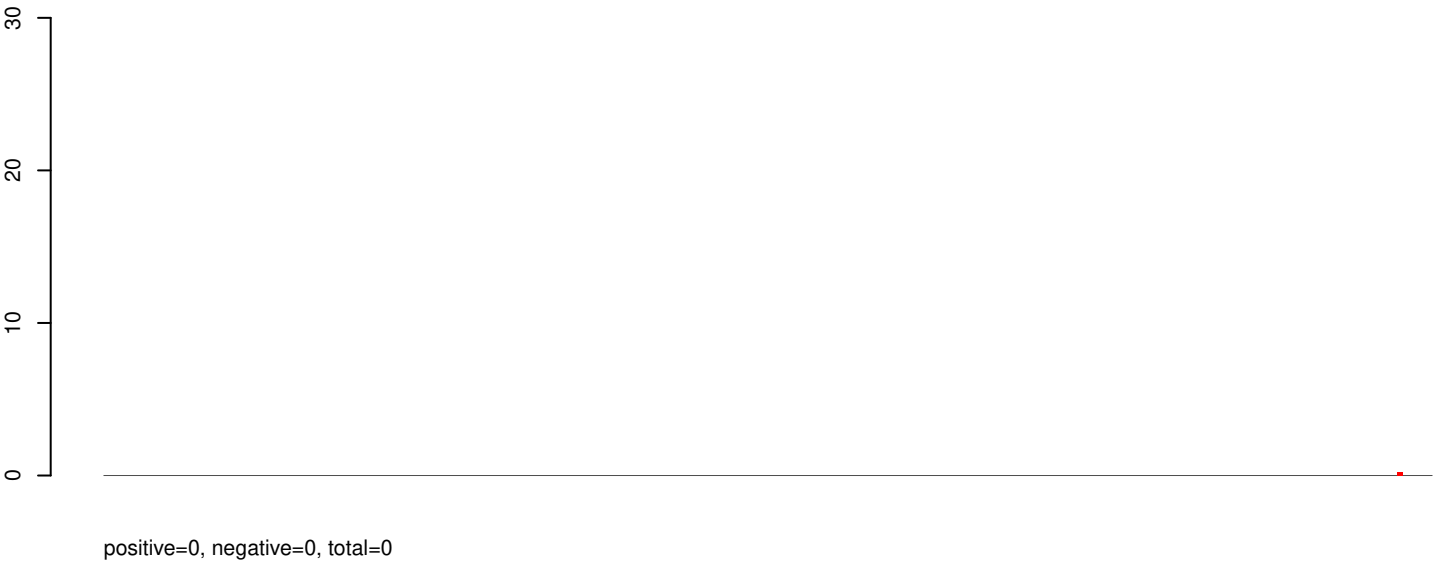


Window size=25, length=11526, virus@NC_012561.1-Highlands_J_virus:1-11526

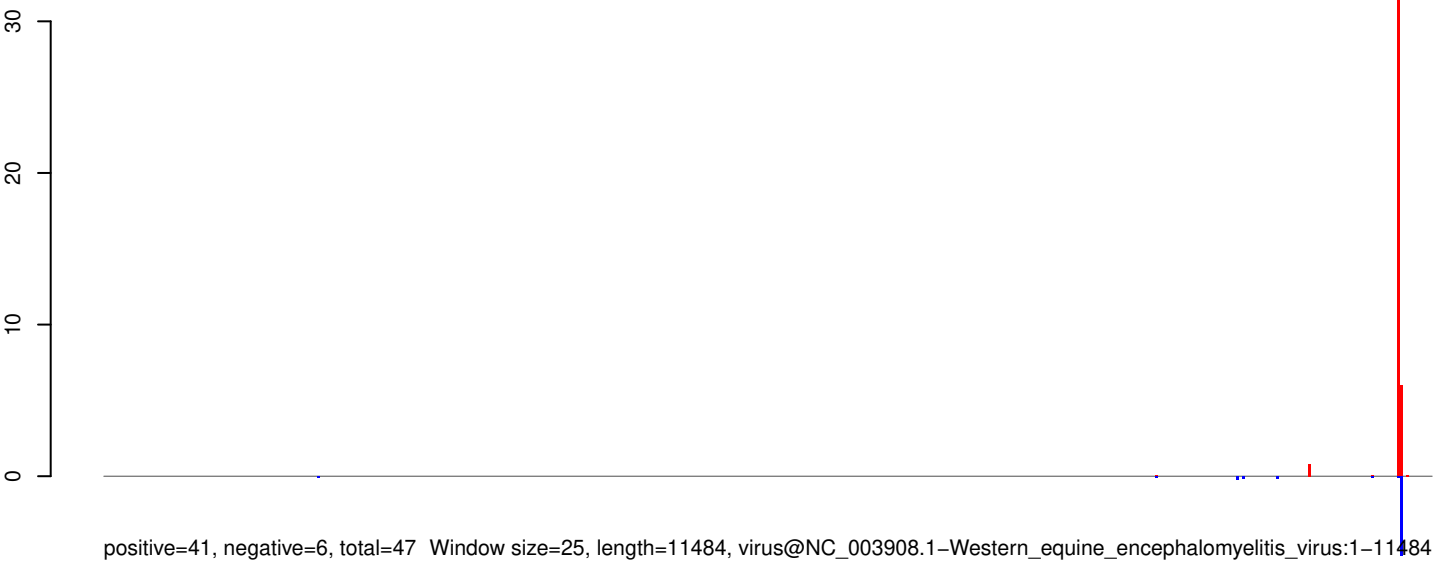
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



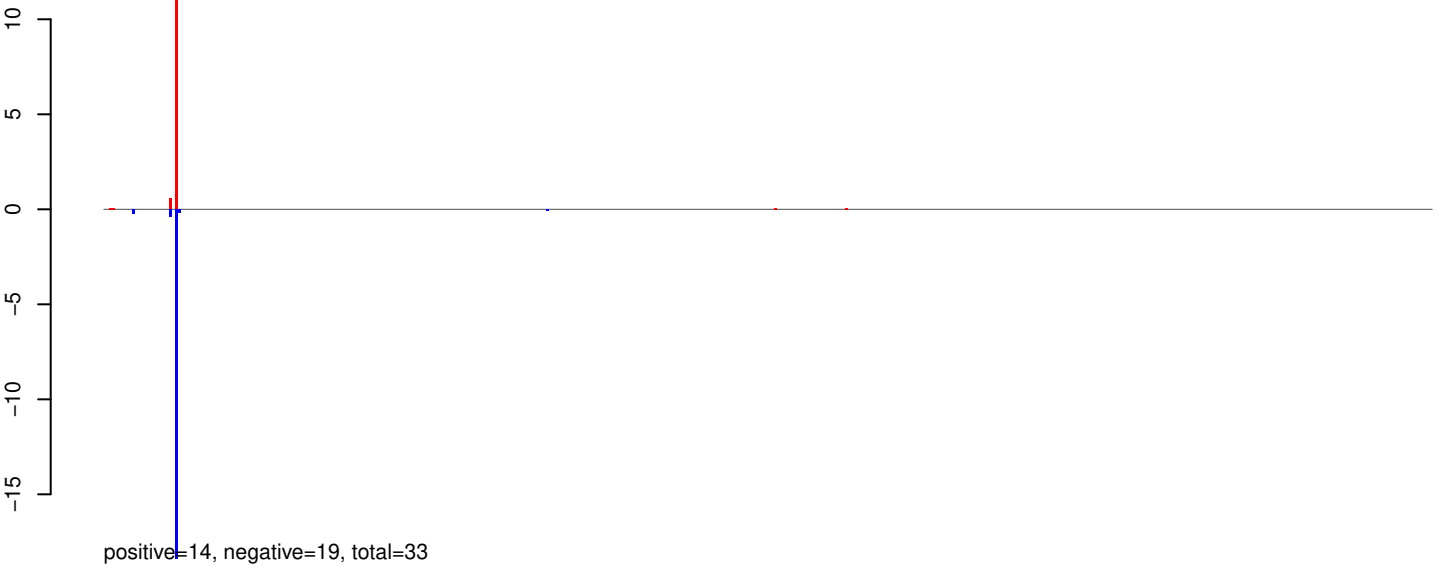
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



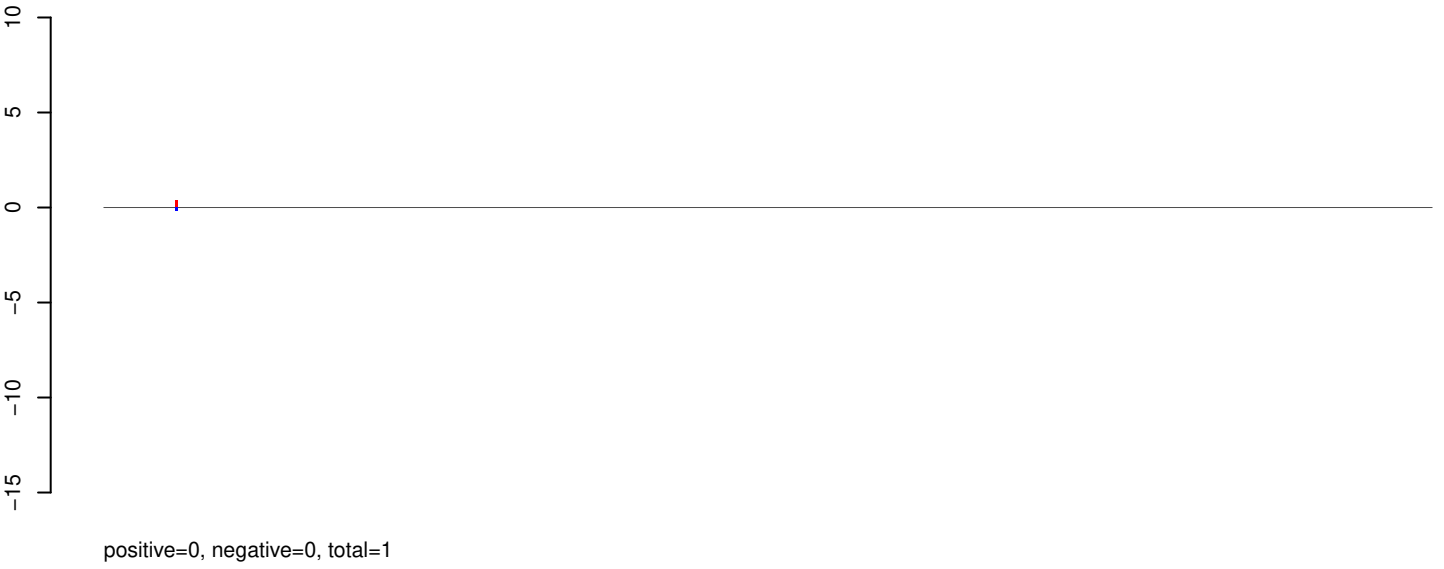
AeAeg_Aag2_mock_dsFluc_RA.rep



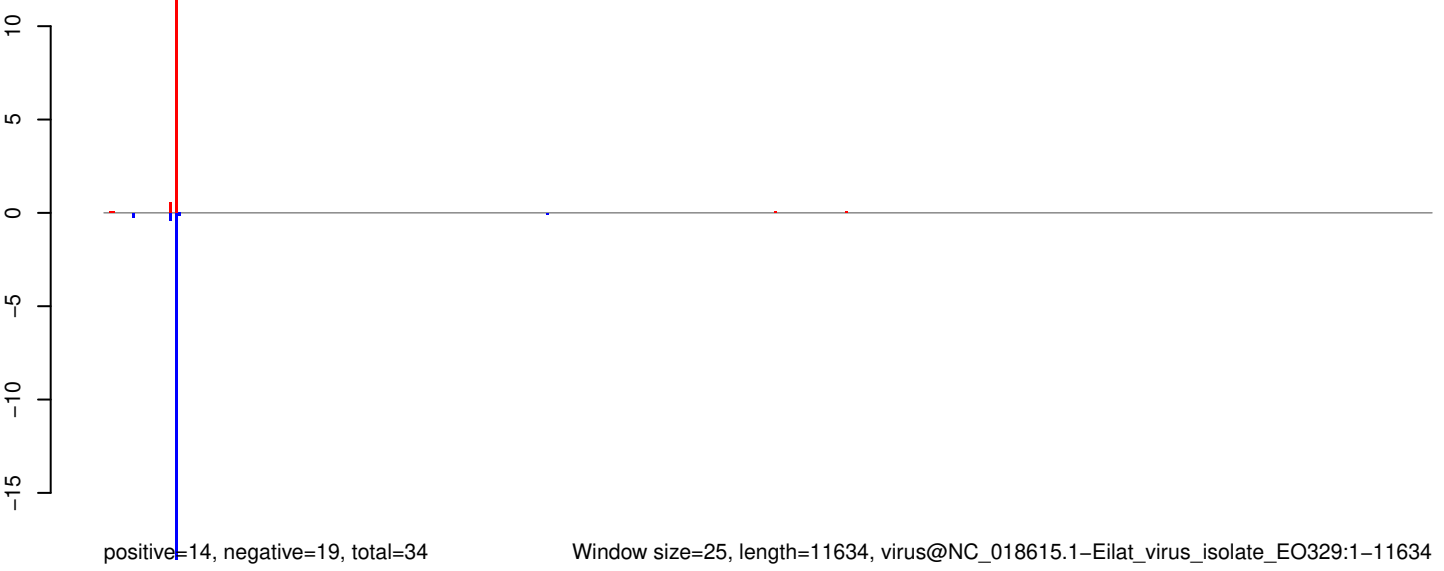
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



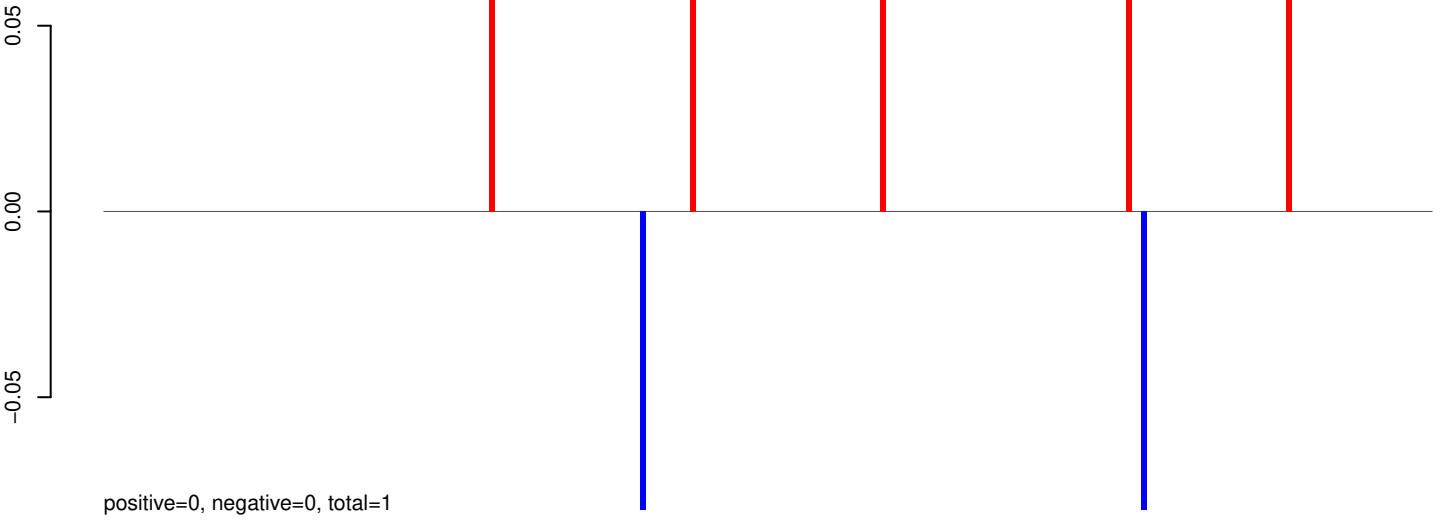
AeAeg_Aag2_mock_dsFluc_RA.rep



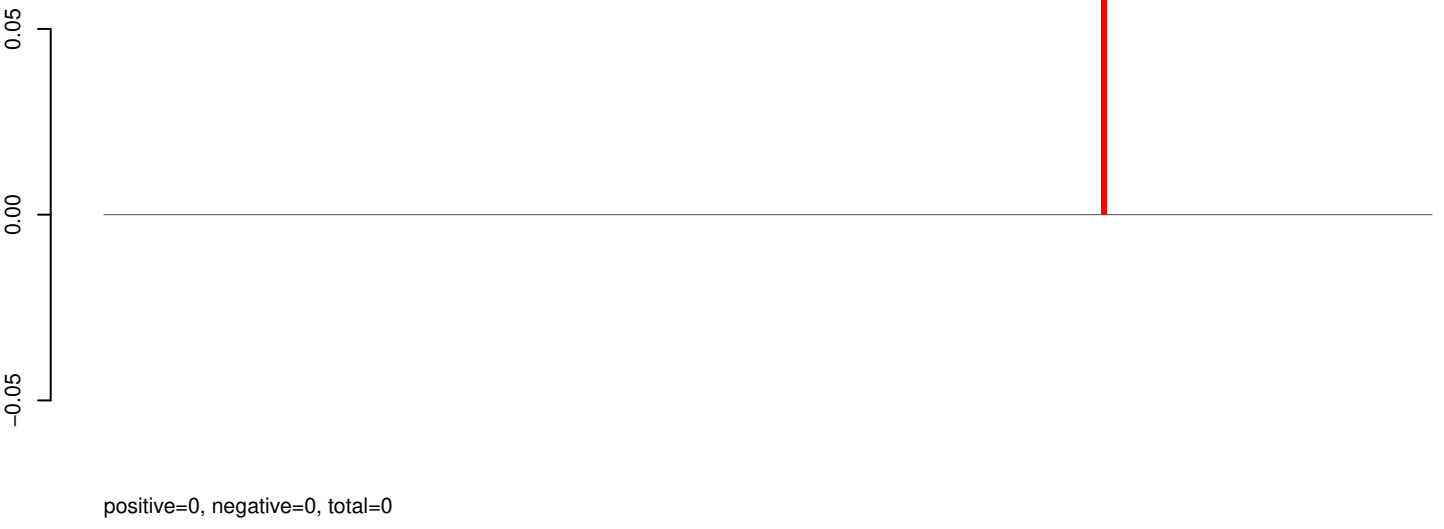
Window size=25, length=11634, virus@NC_018615.1-Eilat_virus_isolate_EO329:1-11634

0 2000 4000 6000 8000 10000 12000

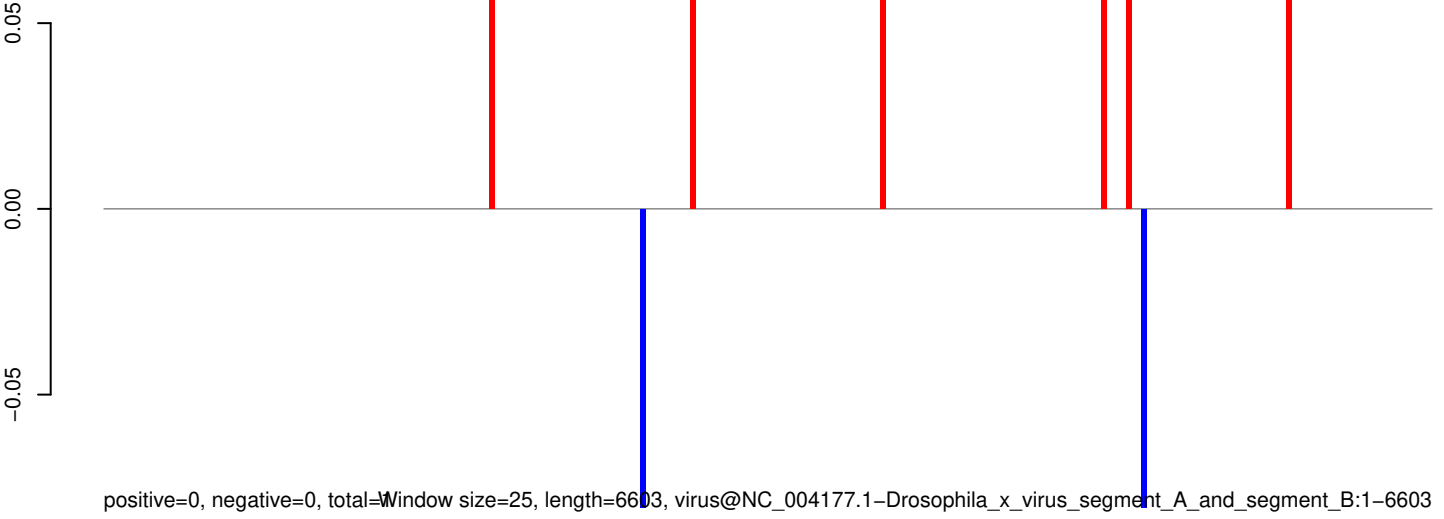
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



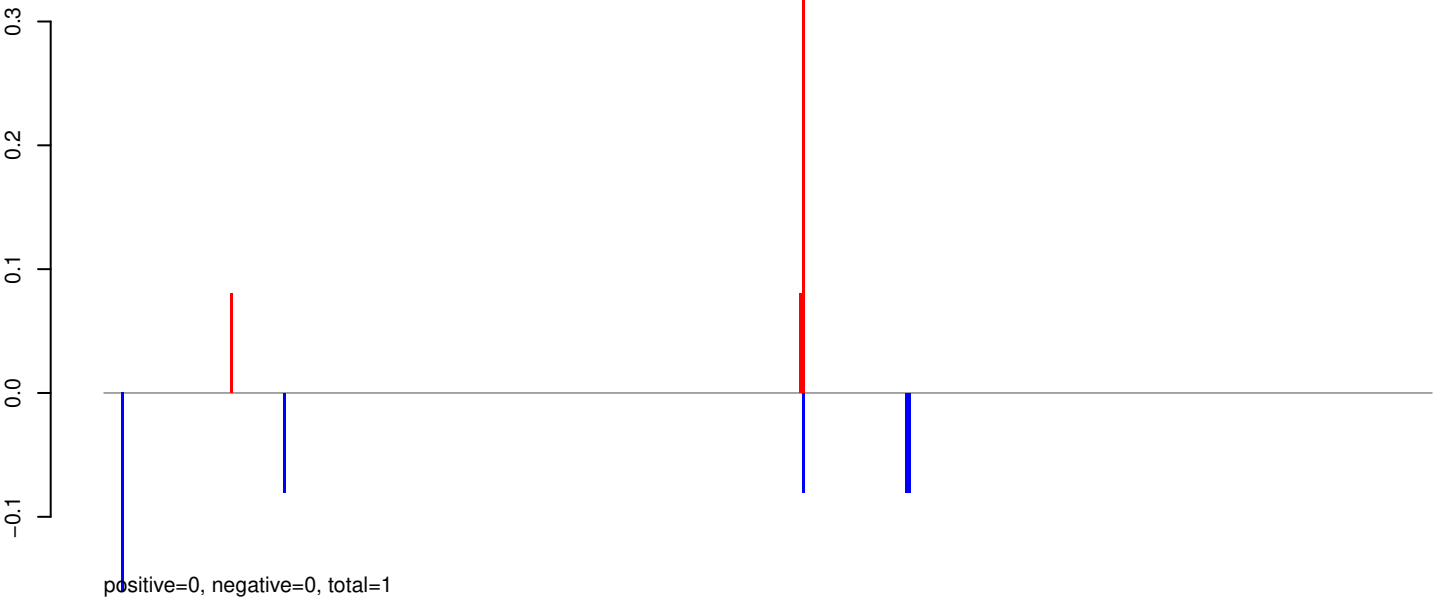
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



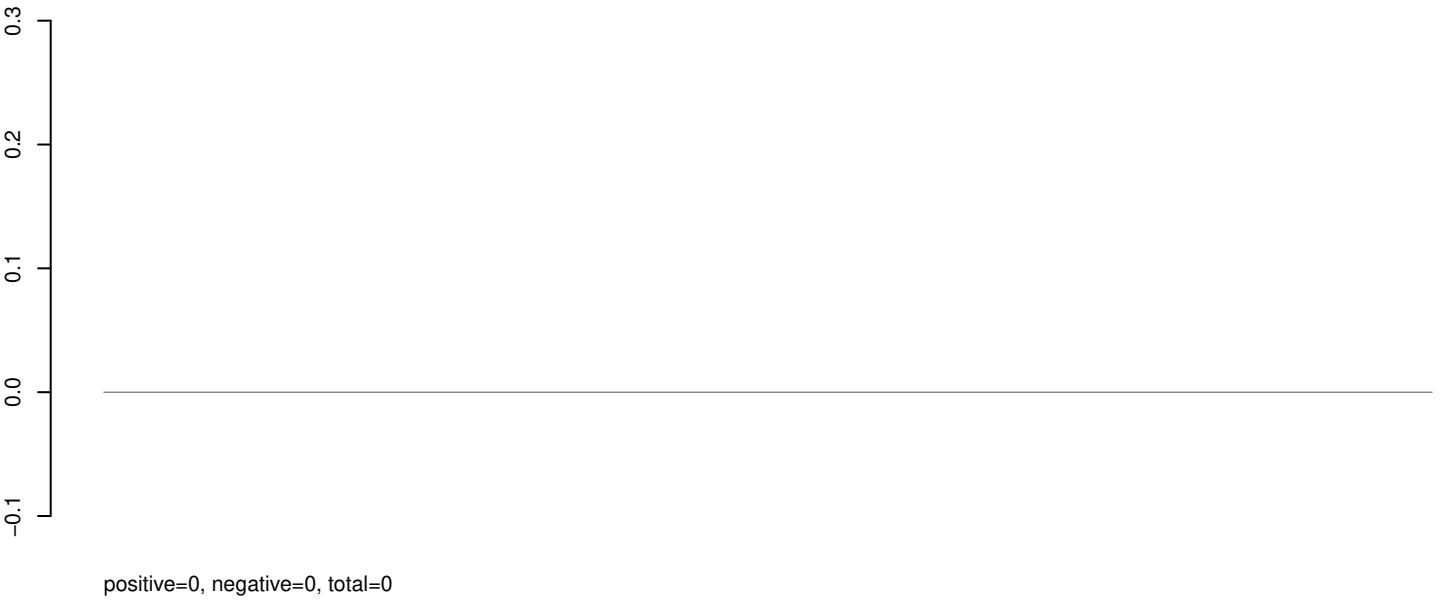
AeAeg_Aag2_mock_dsFluc_RA.rep



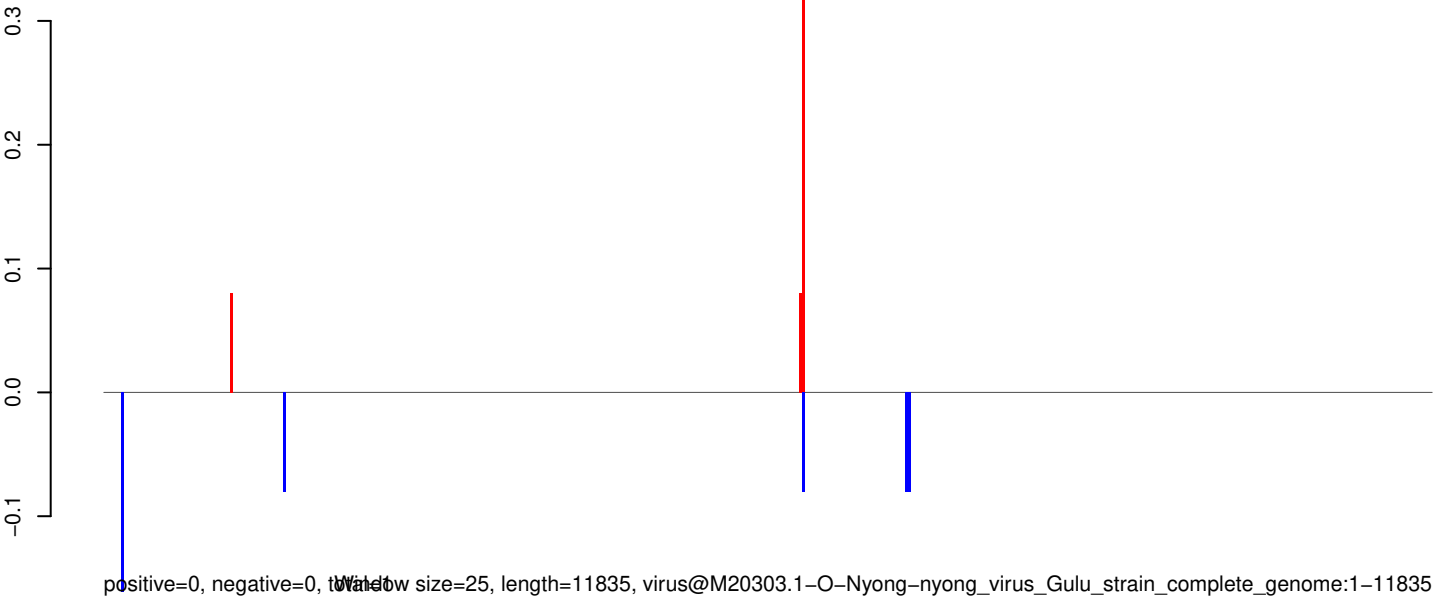
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



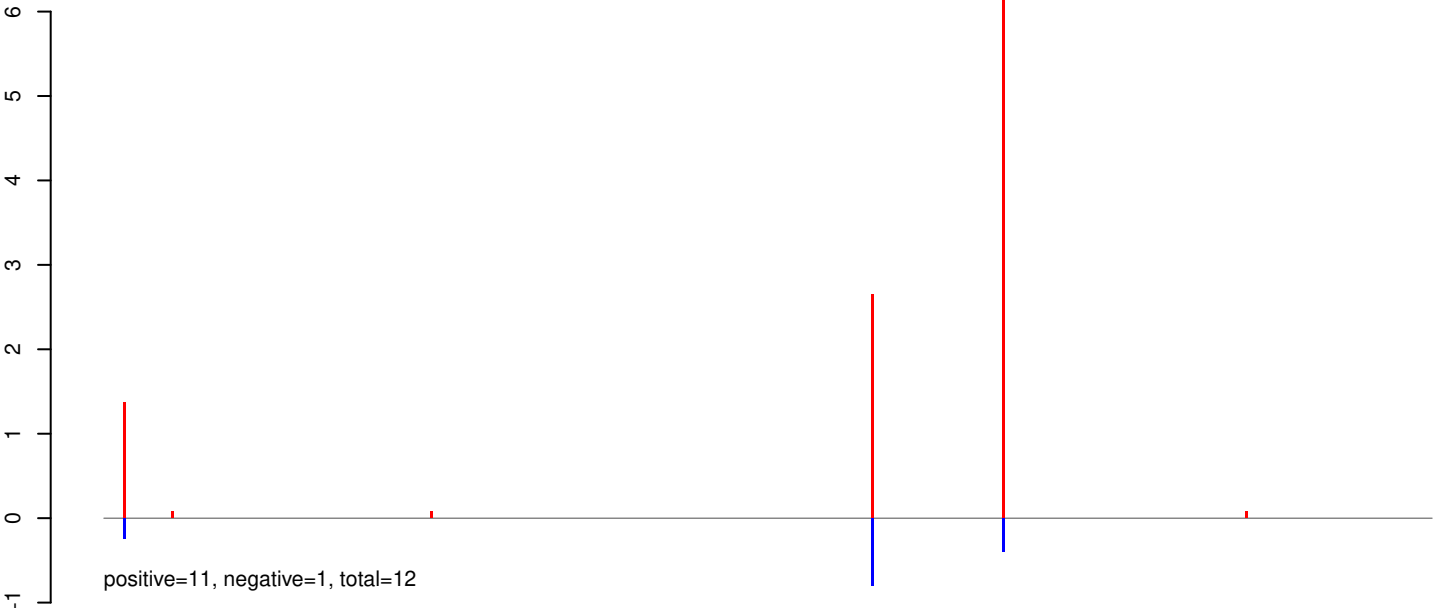
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



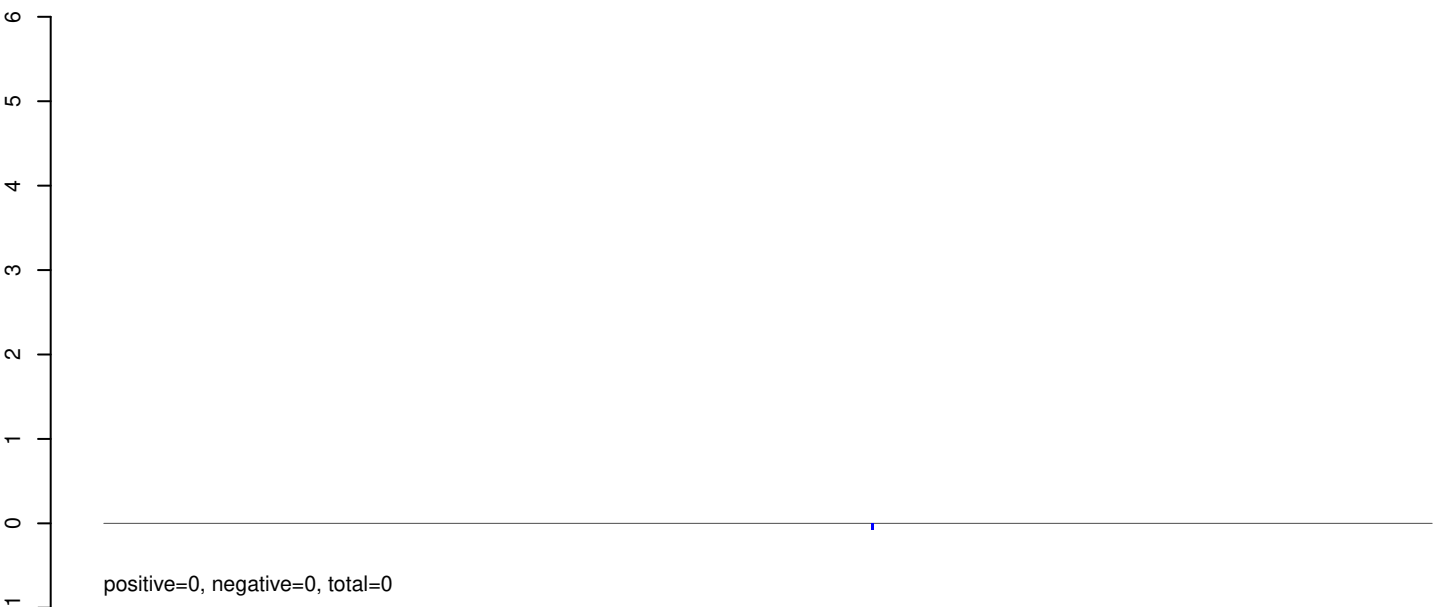
AeAeg_Aag2_mock_dsFluc_RA.rep



AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



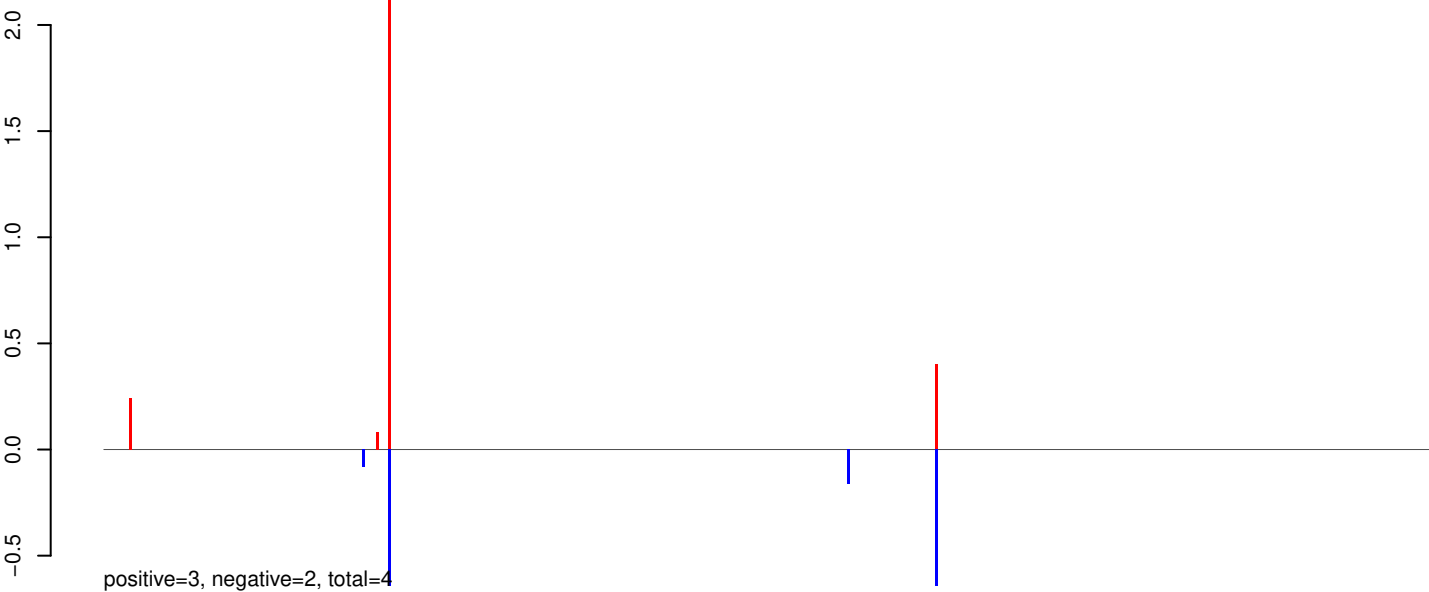
AeAeg_Aag2_mock_dsFluc_RA.rep



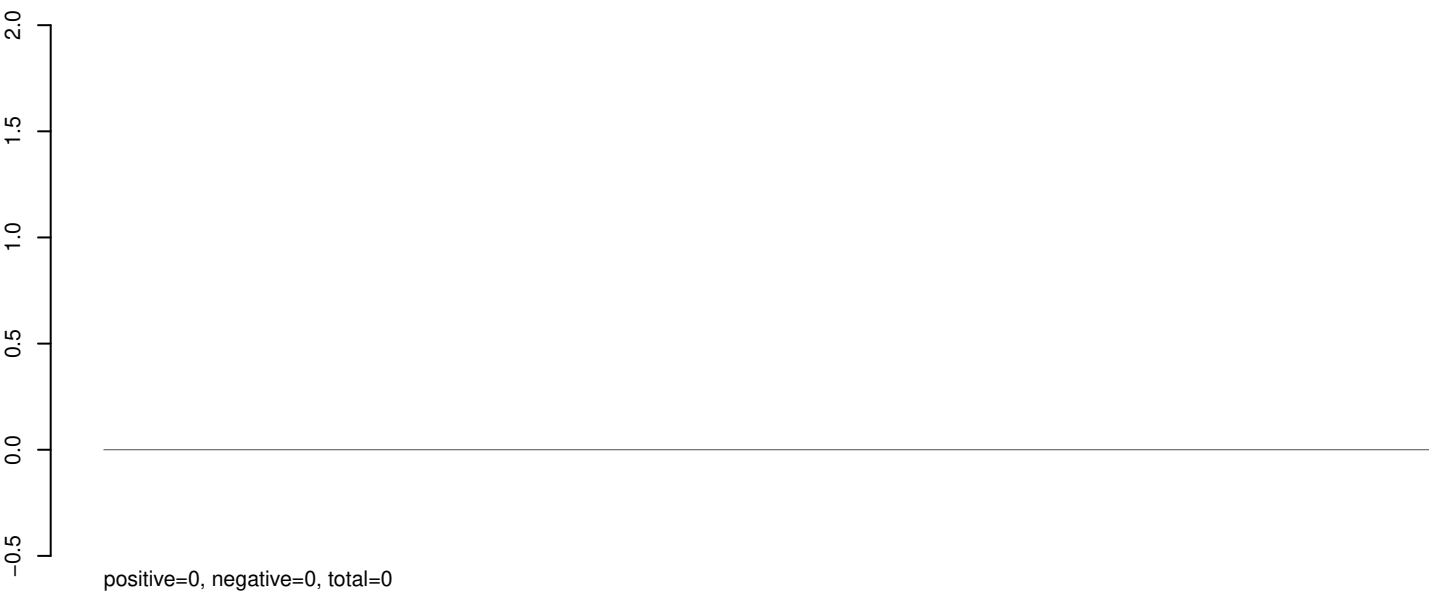
Window size=25, length=11877, virus@NC_016962.1-Bebaru_virus:1-11877

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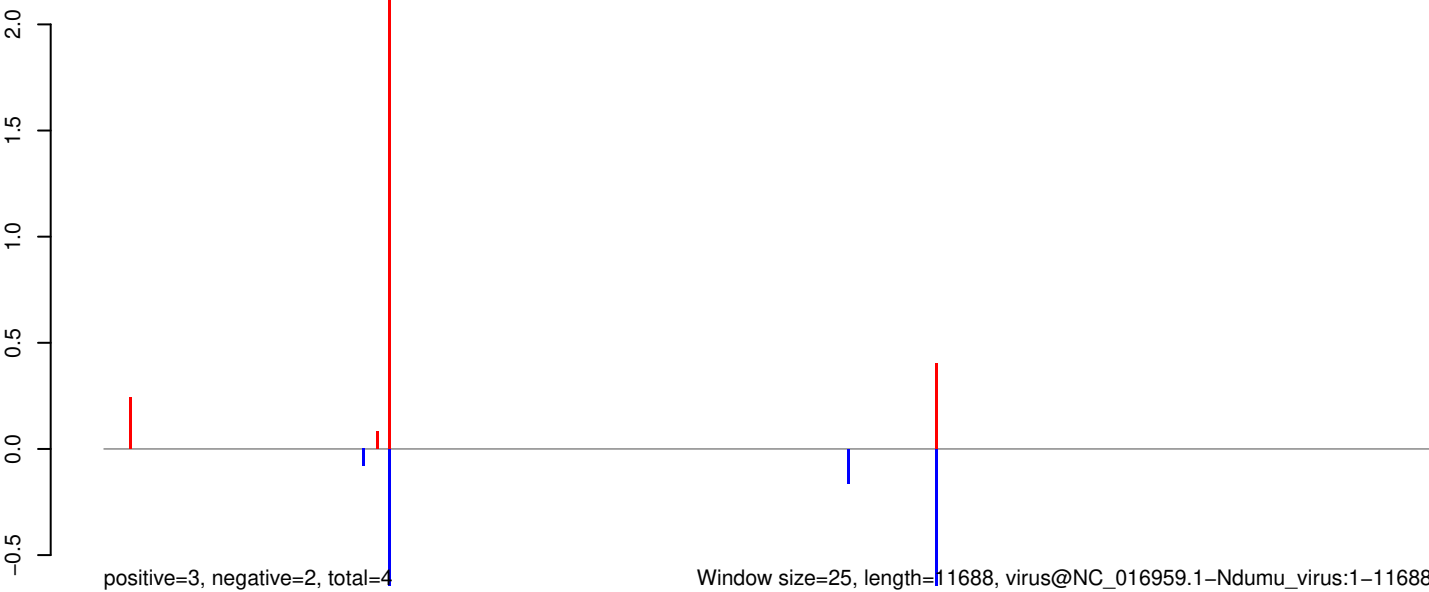
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



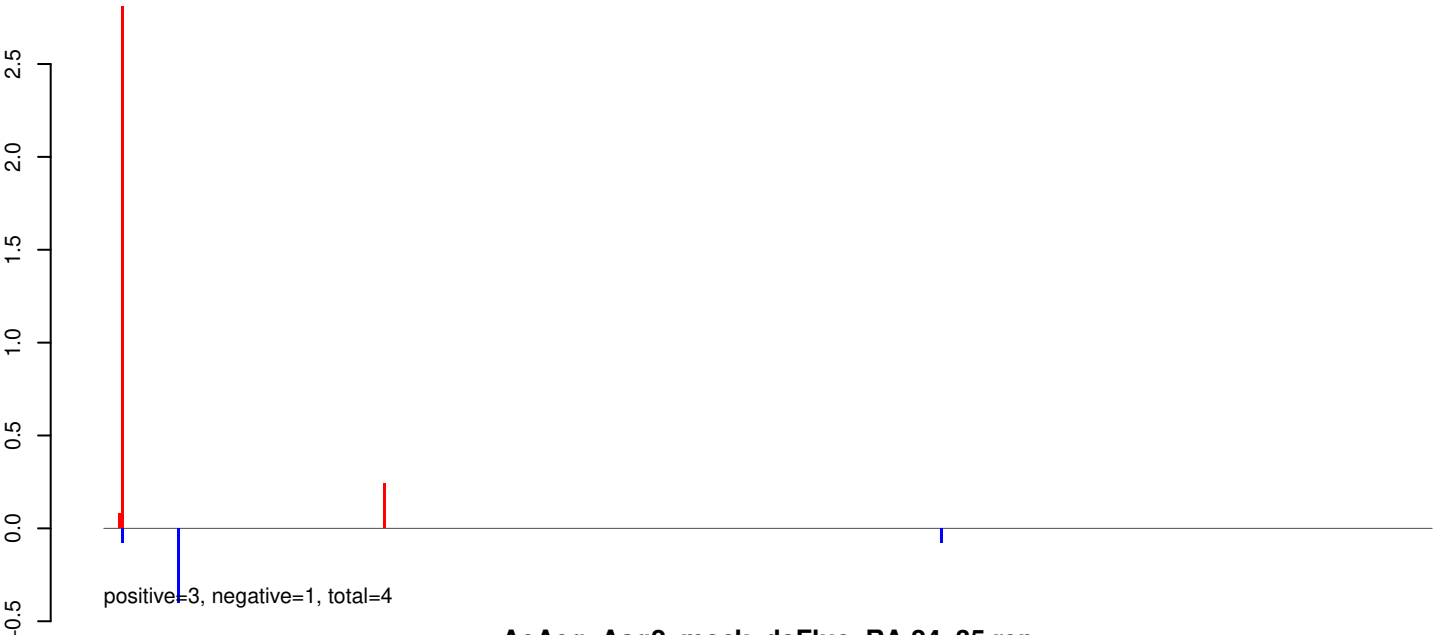
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



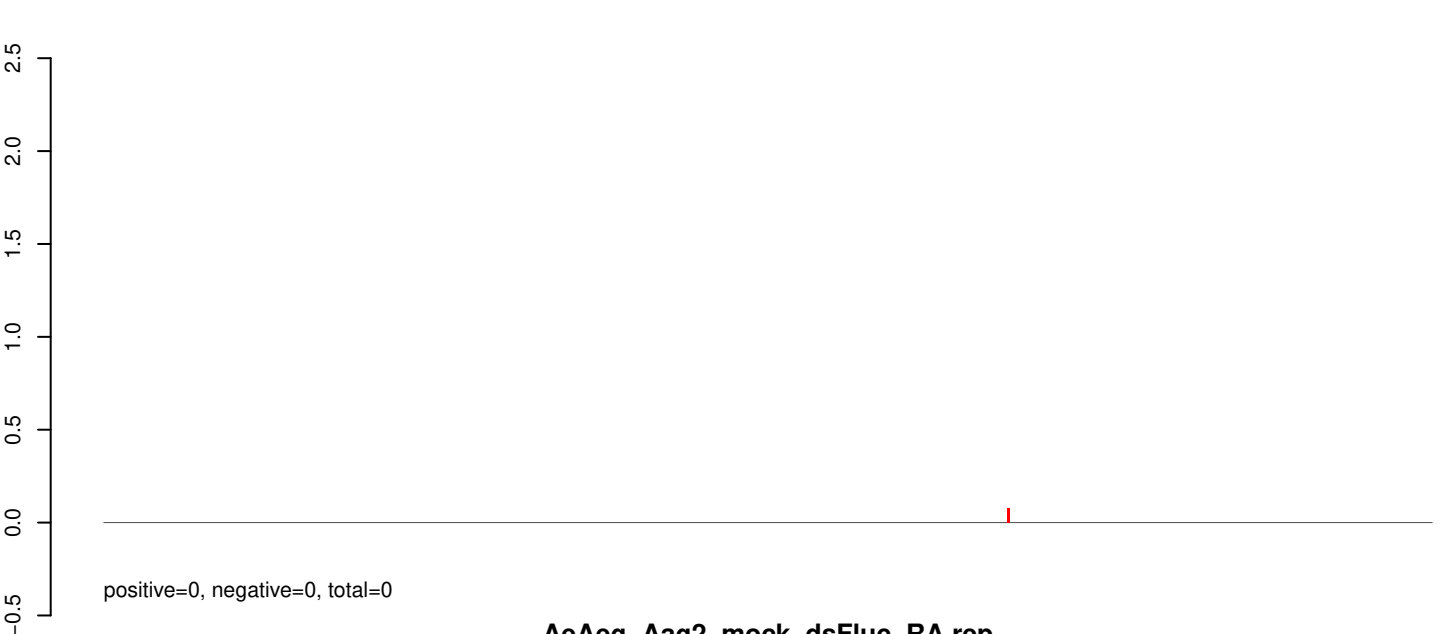
AeAeg_Aag2_mock_dsFluc_RA.rep



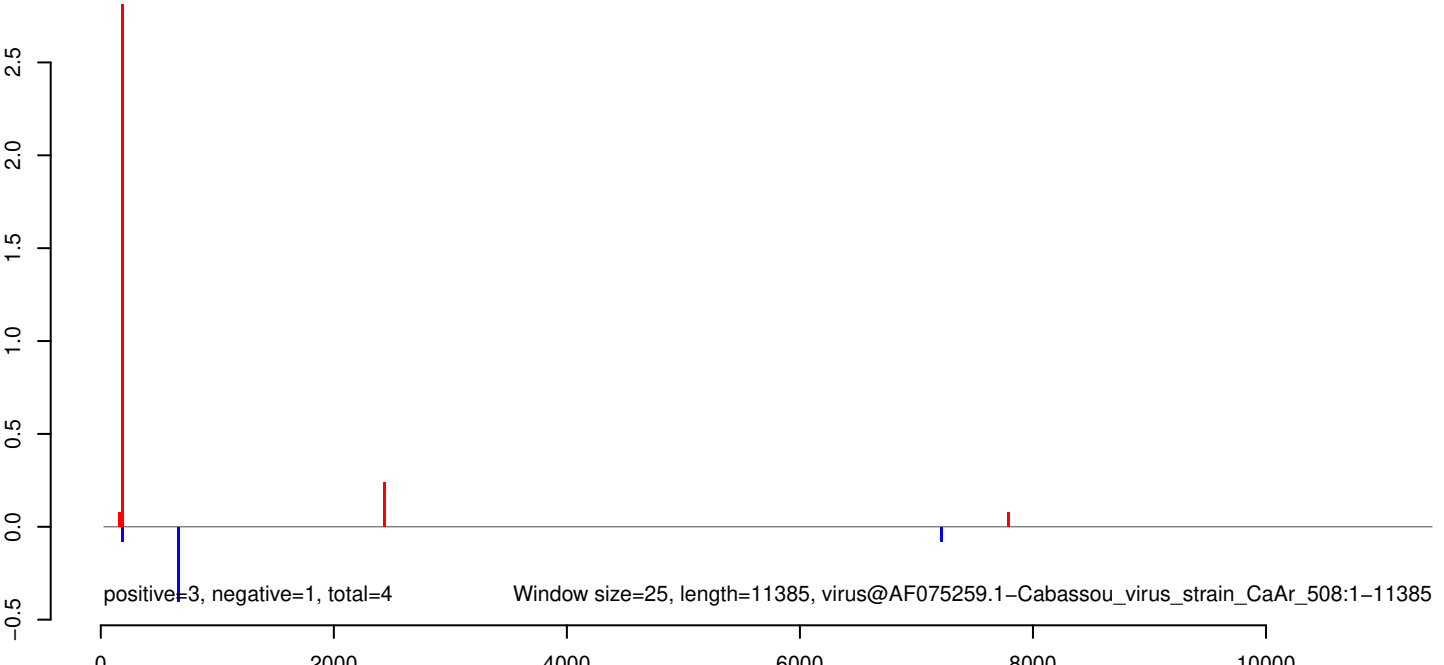
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep



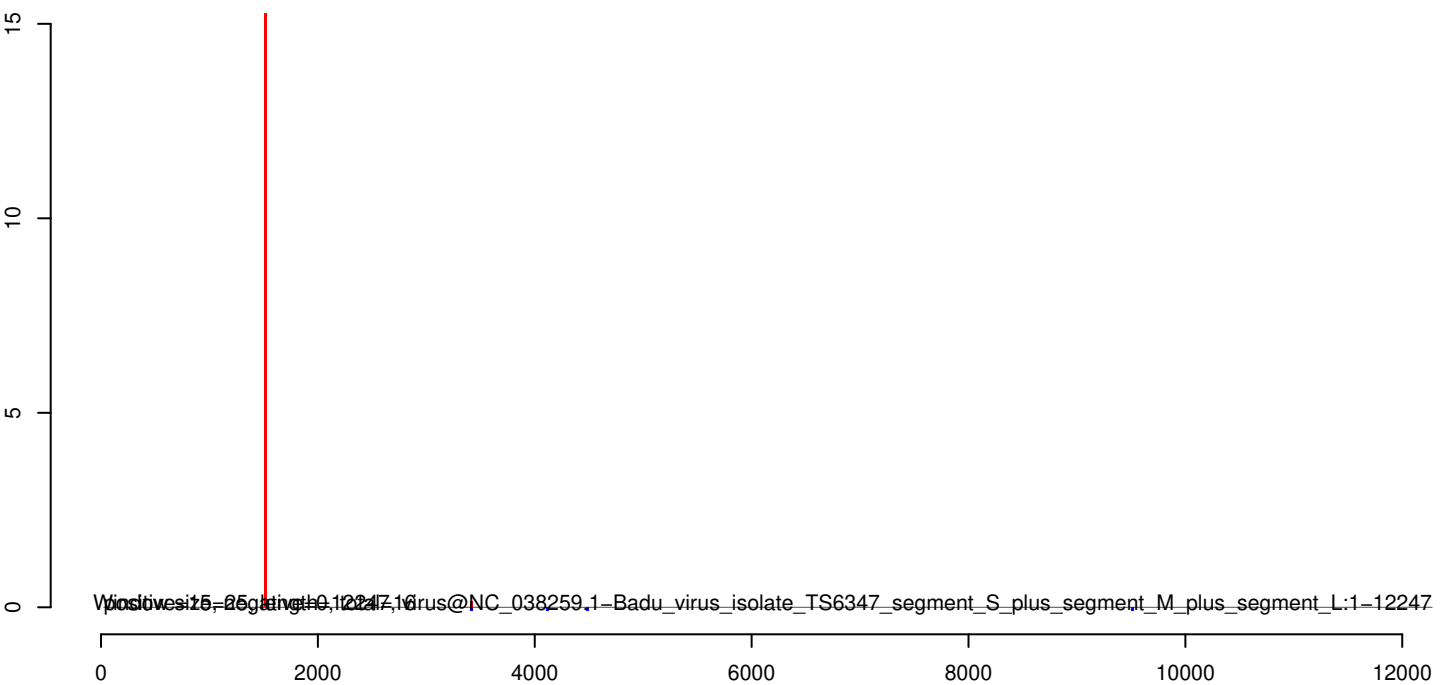
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



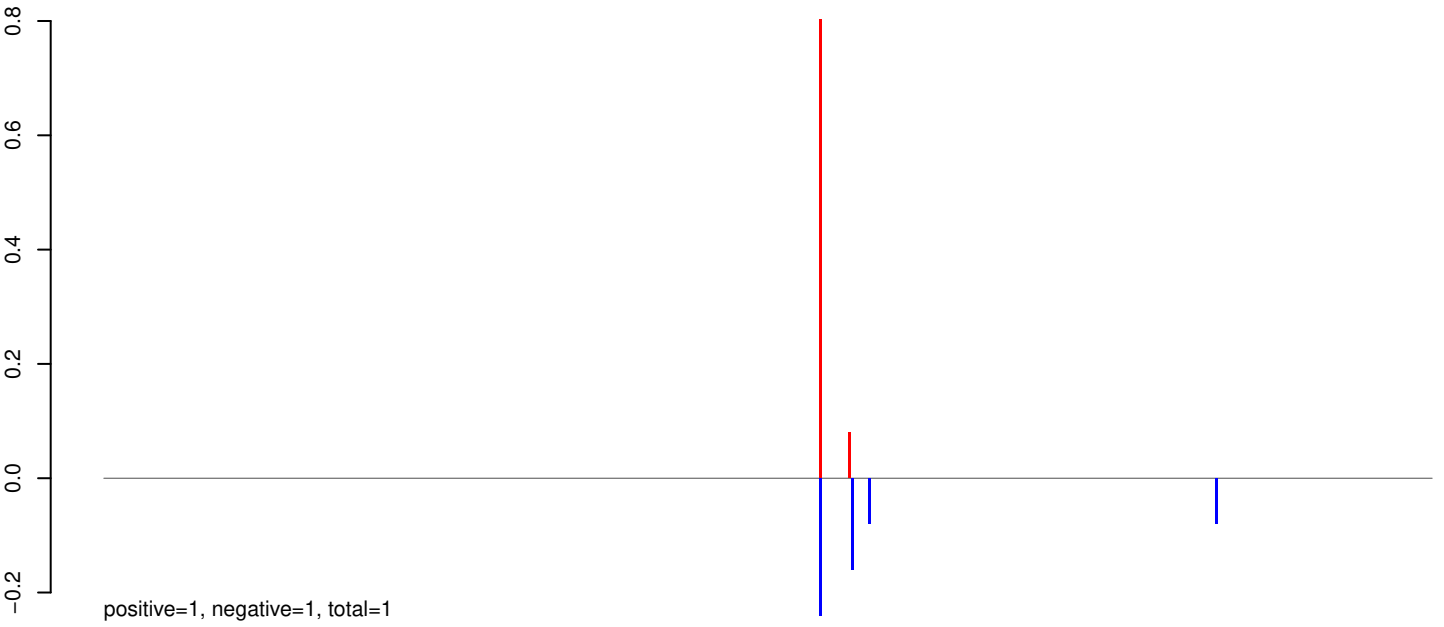
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



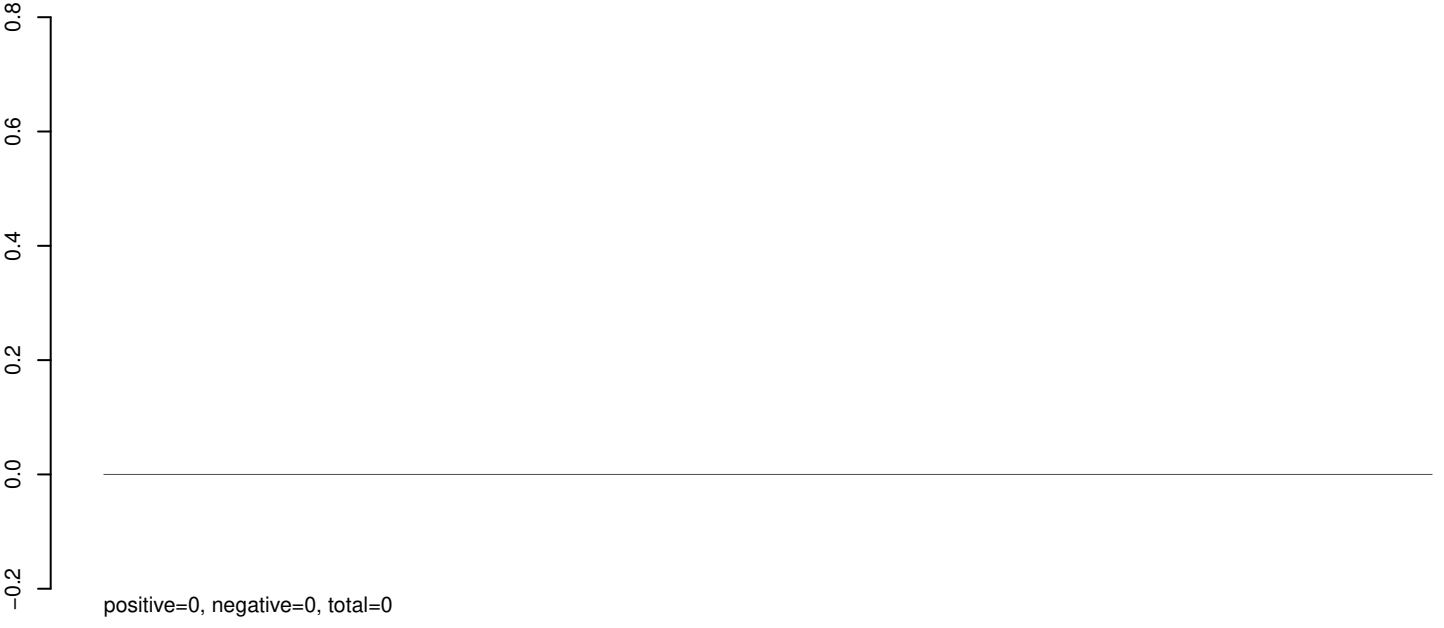
AeAeg_Aag2_mock_dsFluc_RA.rep



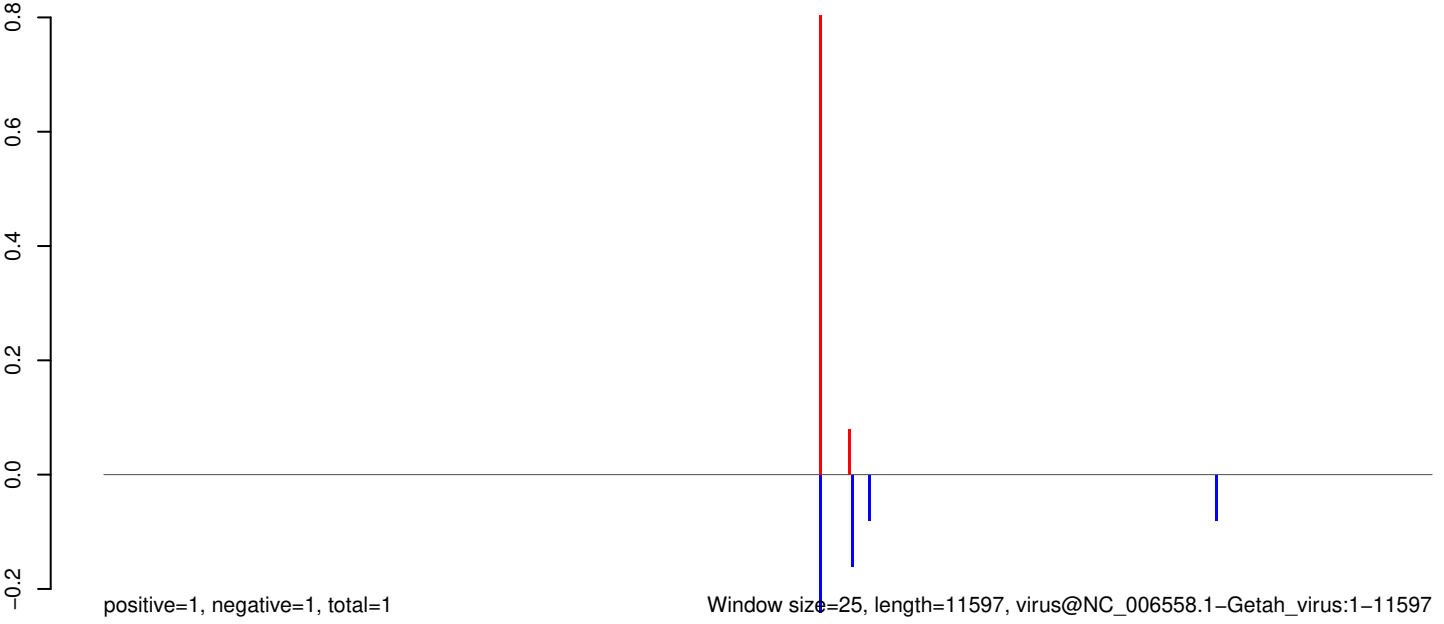
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

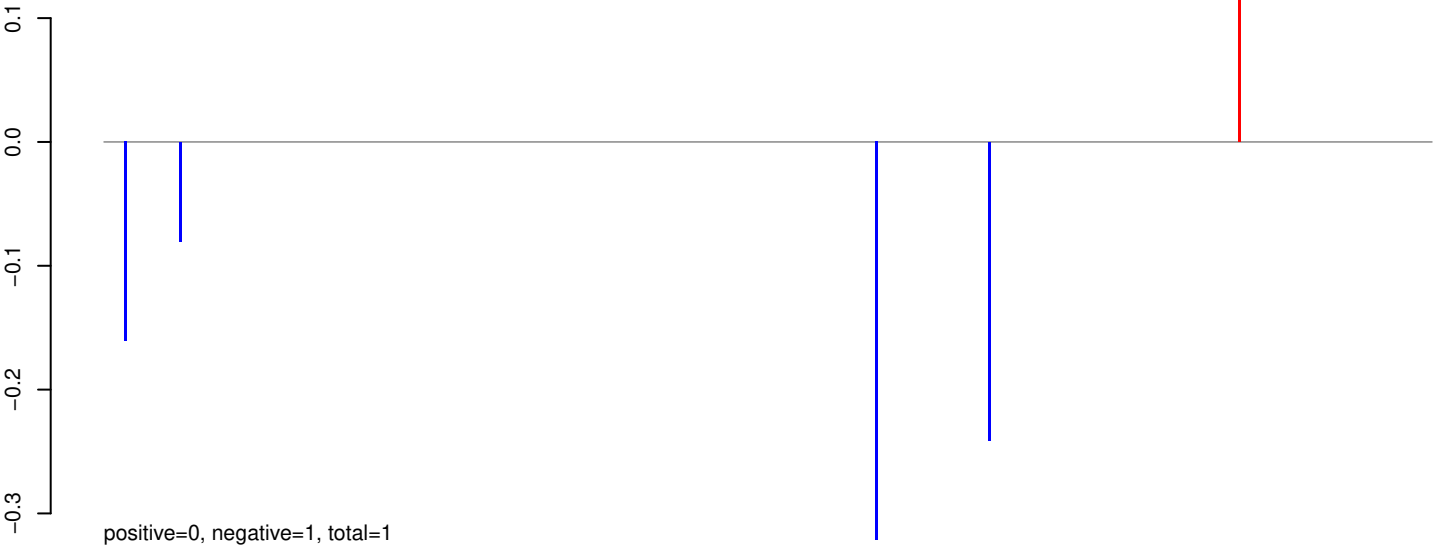


AeAeg_Aag2_mock_dsFluc_RA.rep

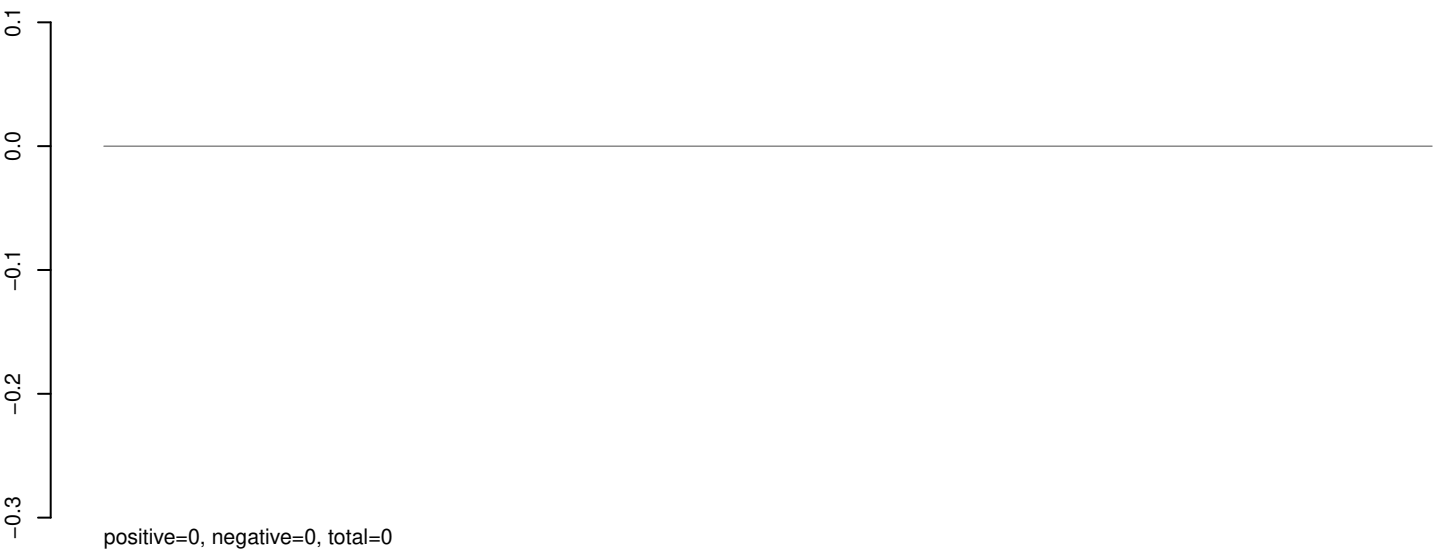


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AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

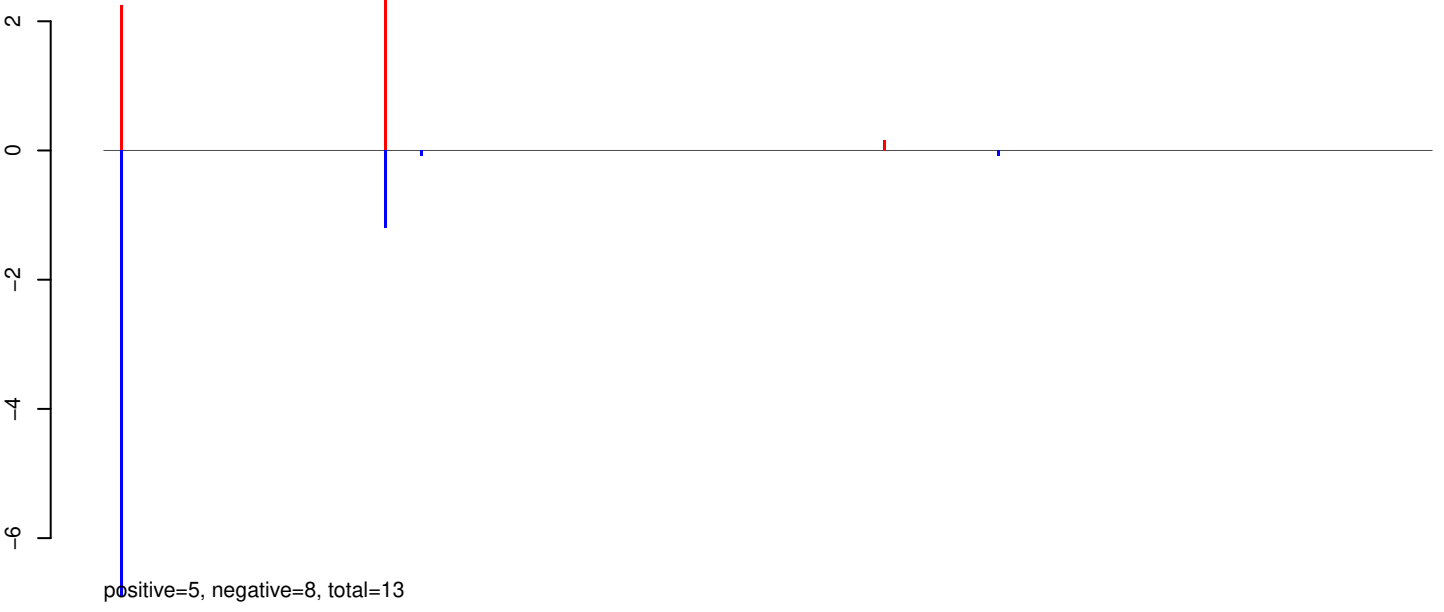


AeAeg_Aag2_mock_dsFluc_RA.rep

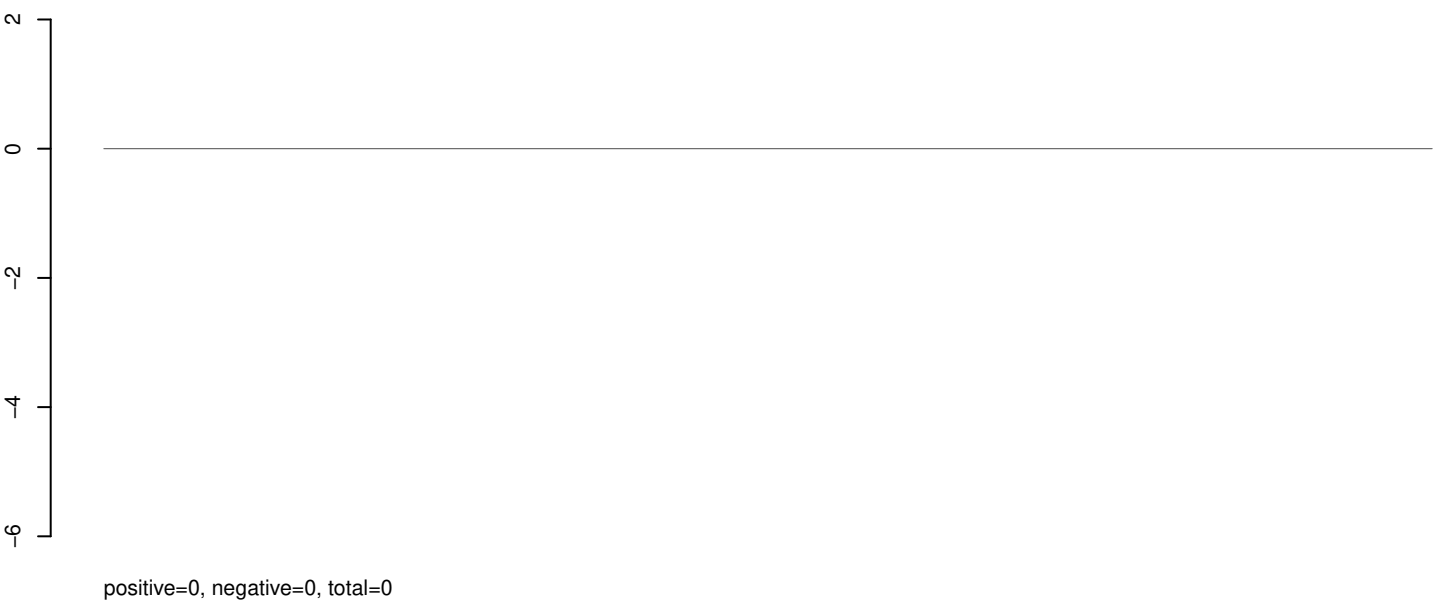


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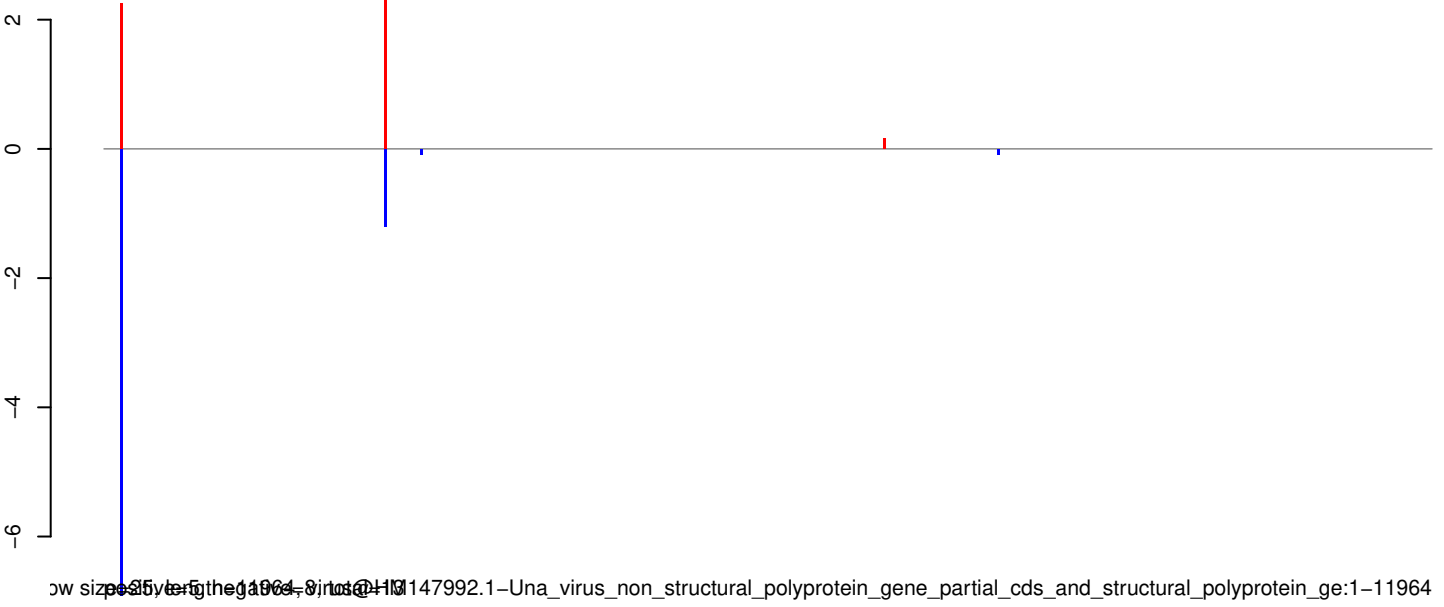
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep



AeAeg_Aag2_mock_dsFluc_RA.18_23.rep

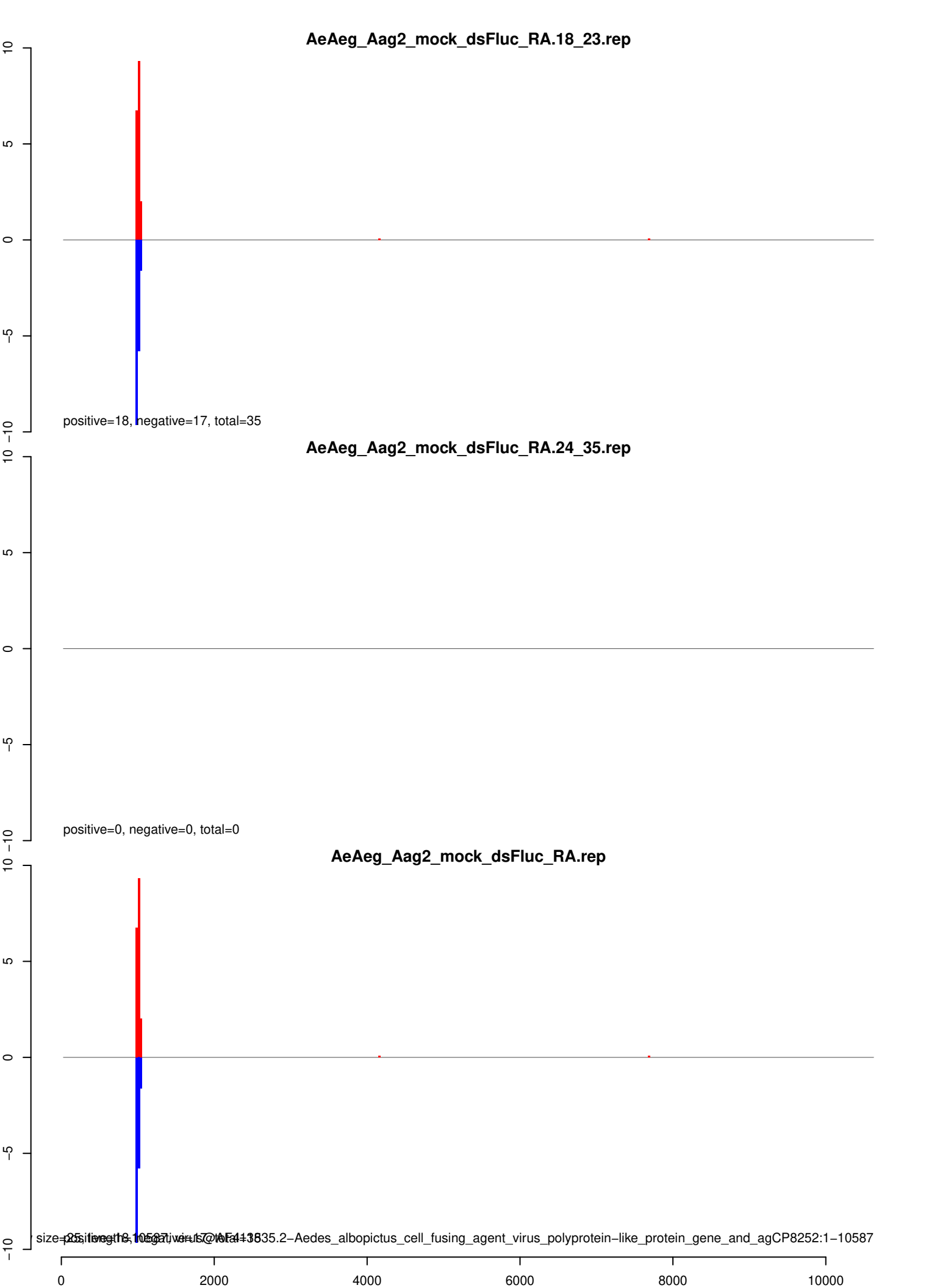
positive=18, negative=17, total=35

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

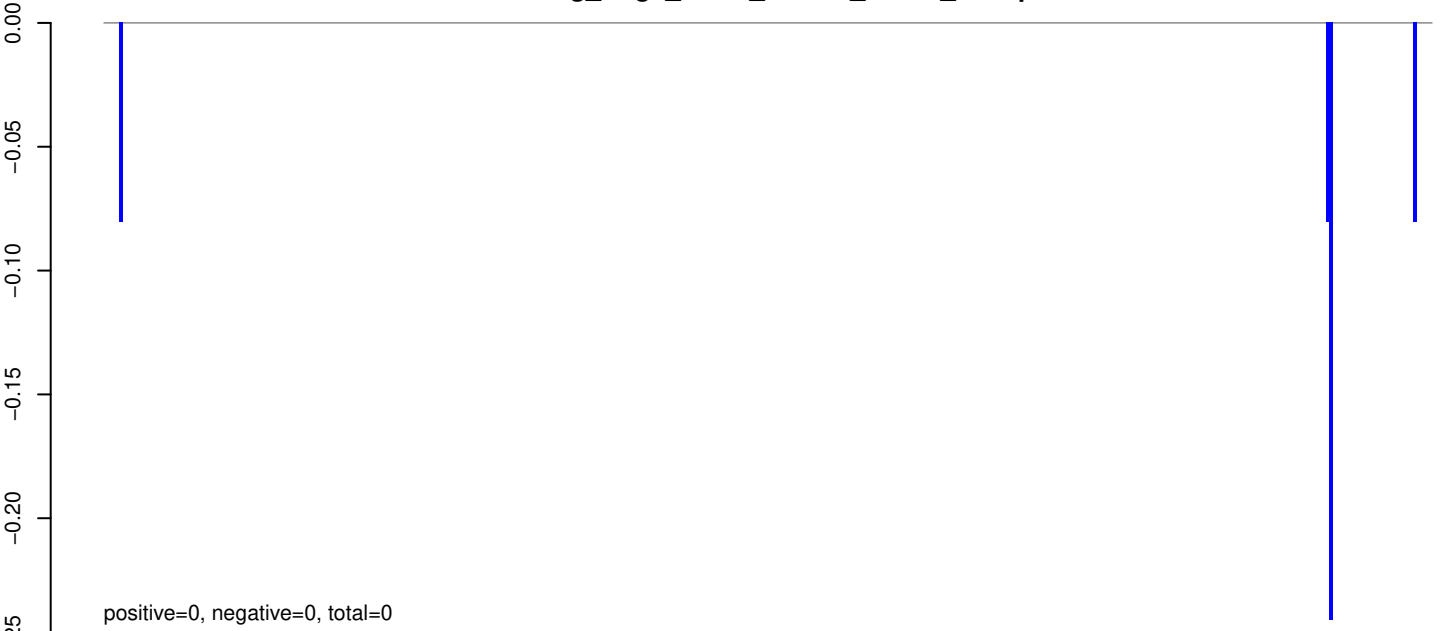
positive=0, negative=0, total=0

AeAeg_Aag2_mock_dsFluc_RA.rep

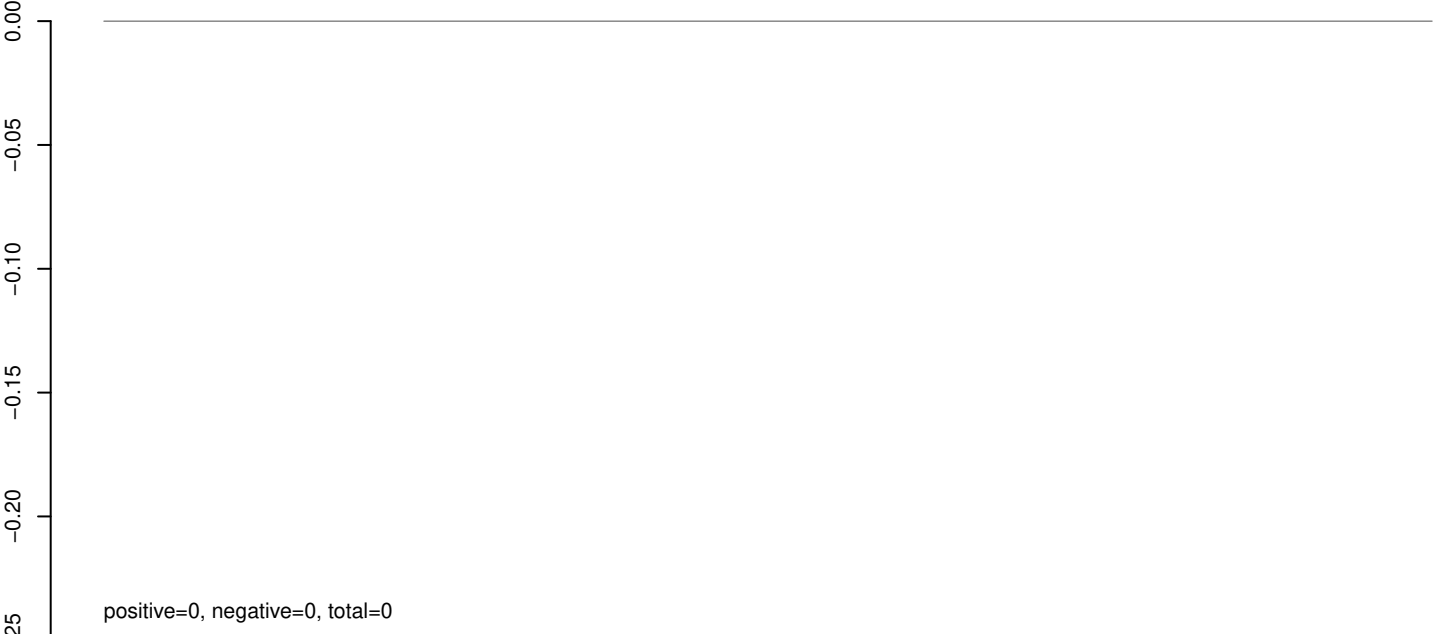
size=25, length=10587, version=16, Fa413835.2-Aedes_albopictus_cell_fusing_agent_virus_polyprotein-like_protein_gene_and_agCP8252:1-10587



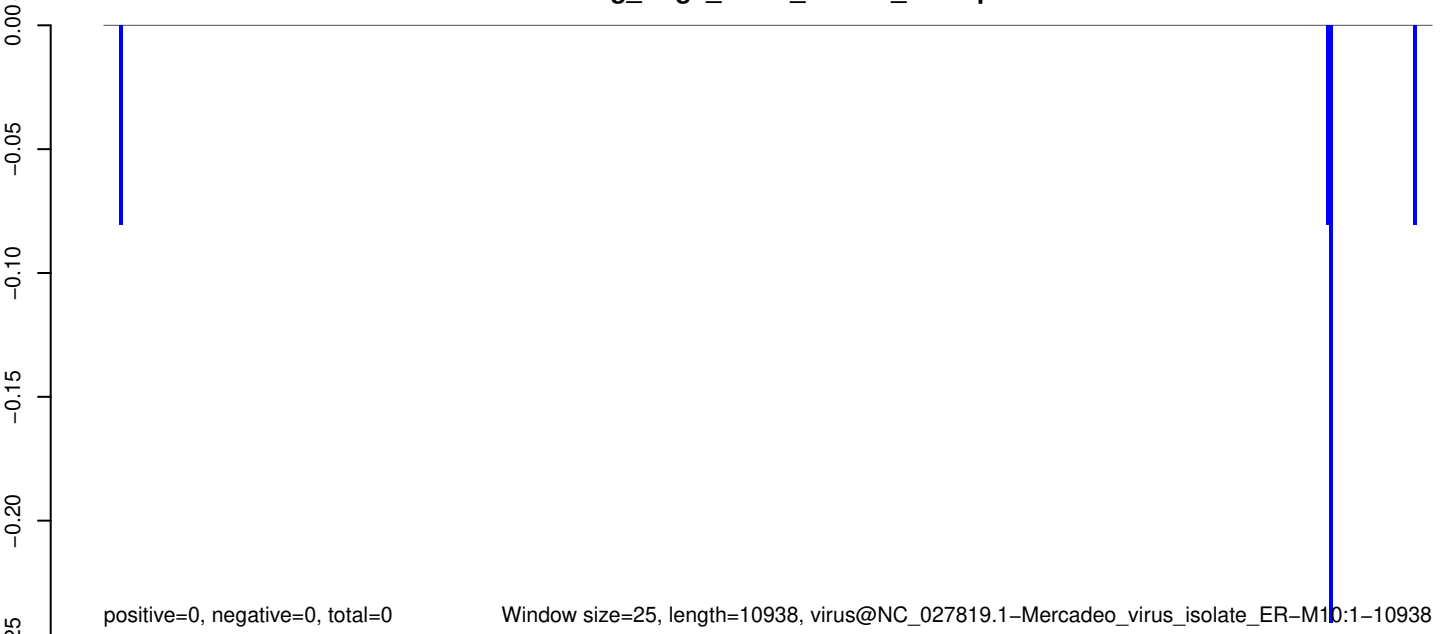
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

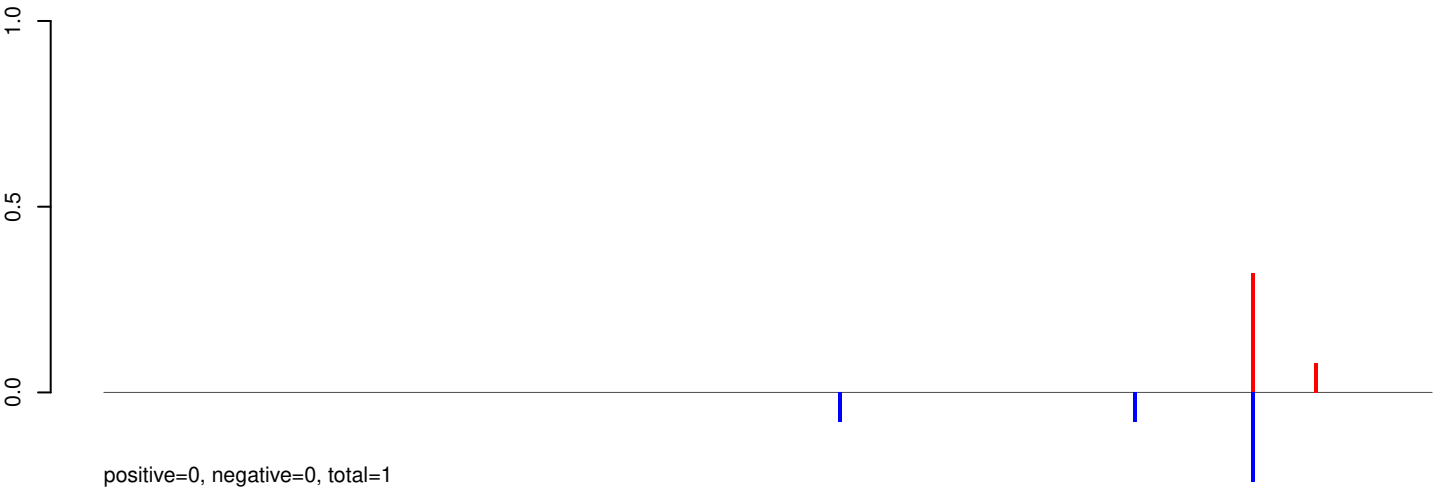


AeAeg_Aag2_mock_dsFluc_RA.rep

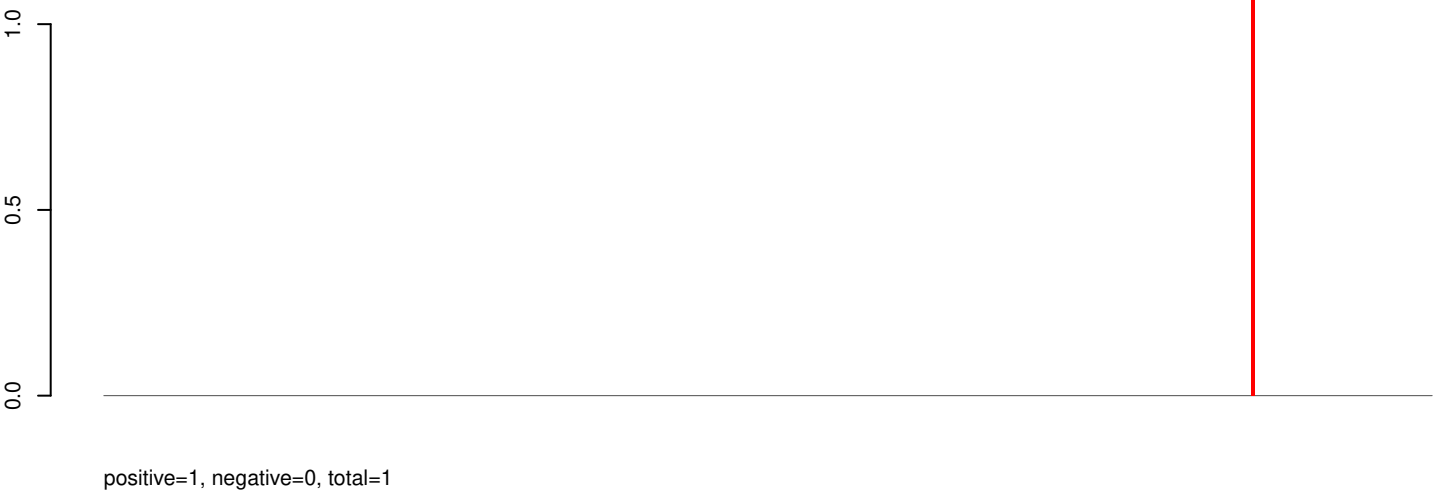


Window size=25, length=10938, virus@NC_027819.1-Mercadeo_virus_isolate_ER-M10:1-10938

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



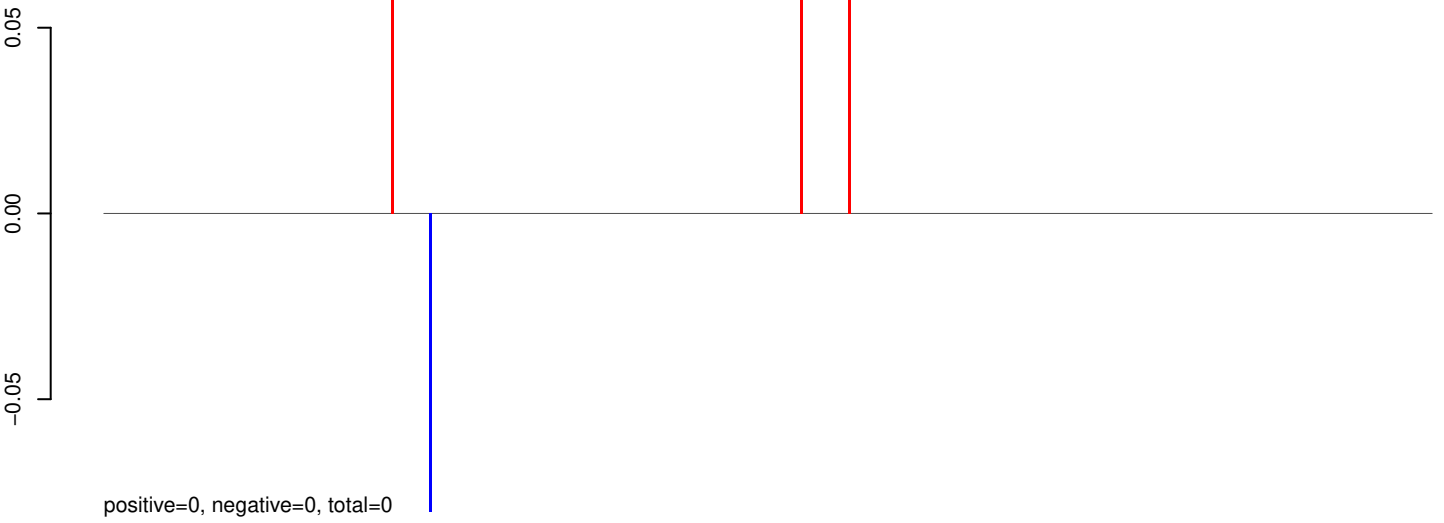
AeAeg_Aag2_mock_dsFluc_RA.rep



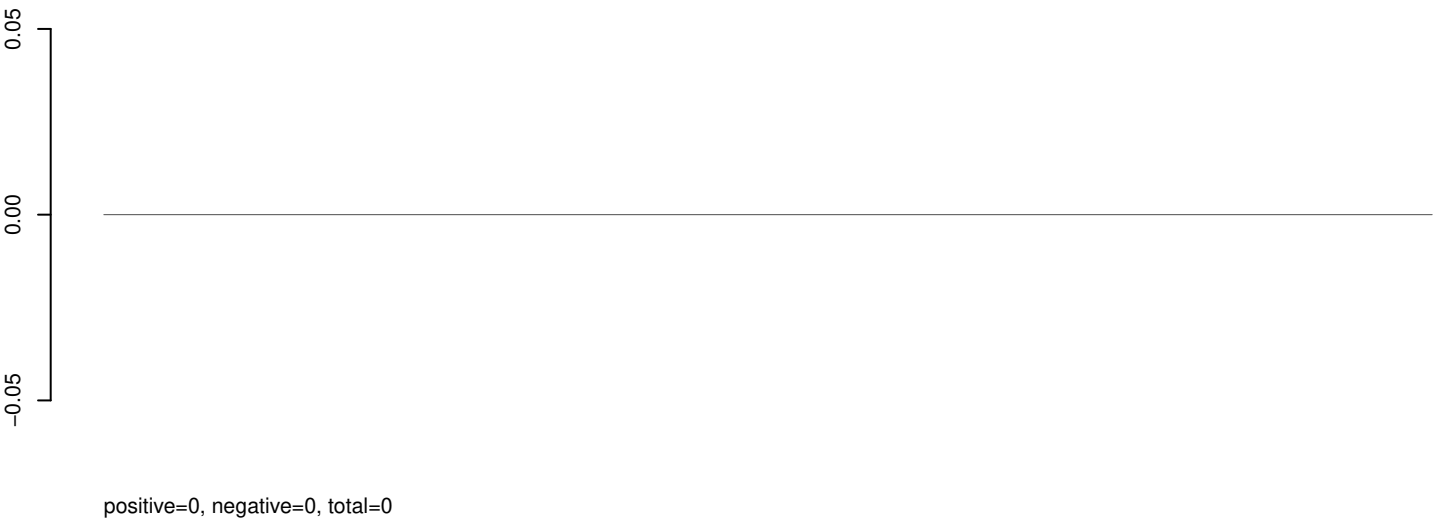
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0 2000 4000 6000 8000 10000

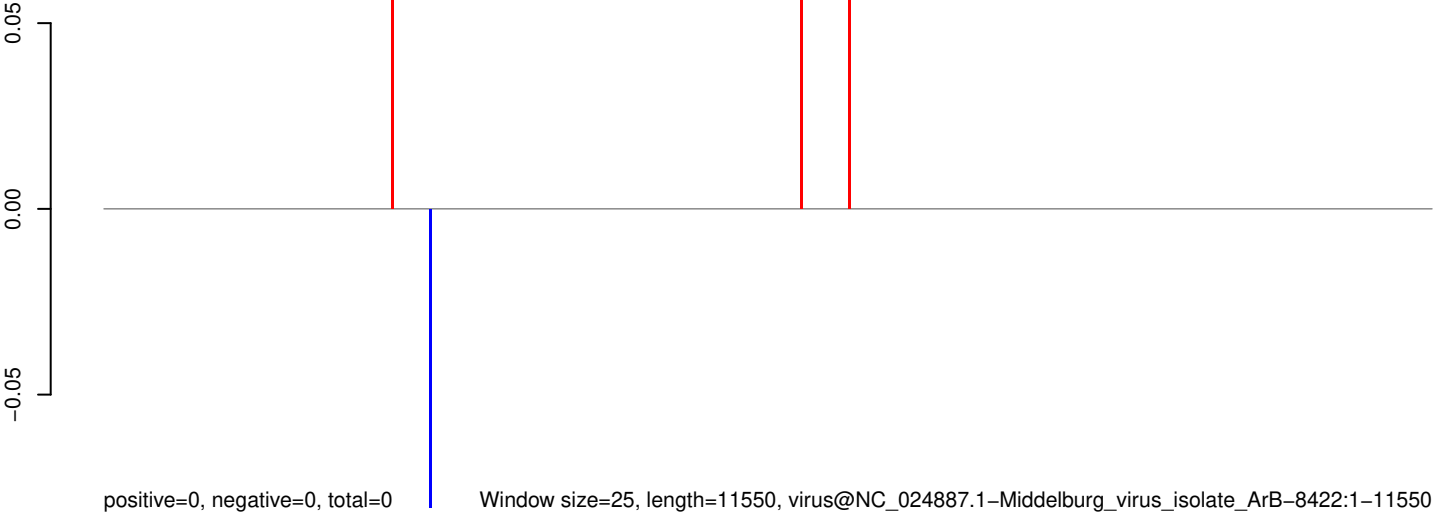
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



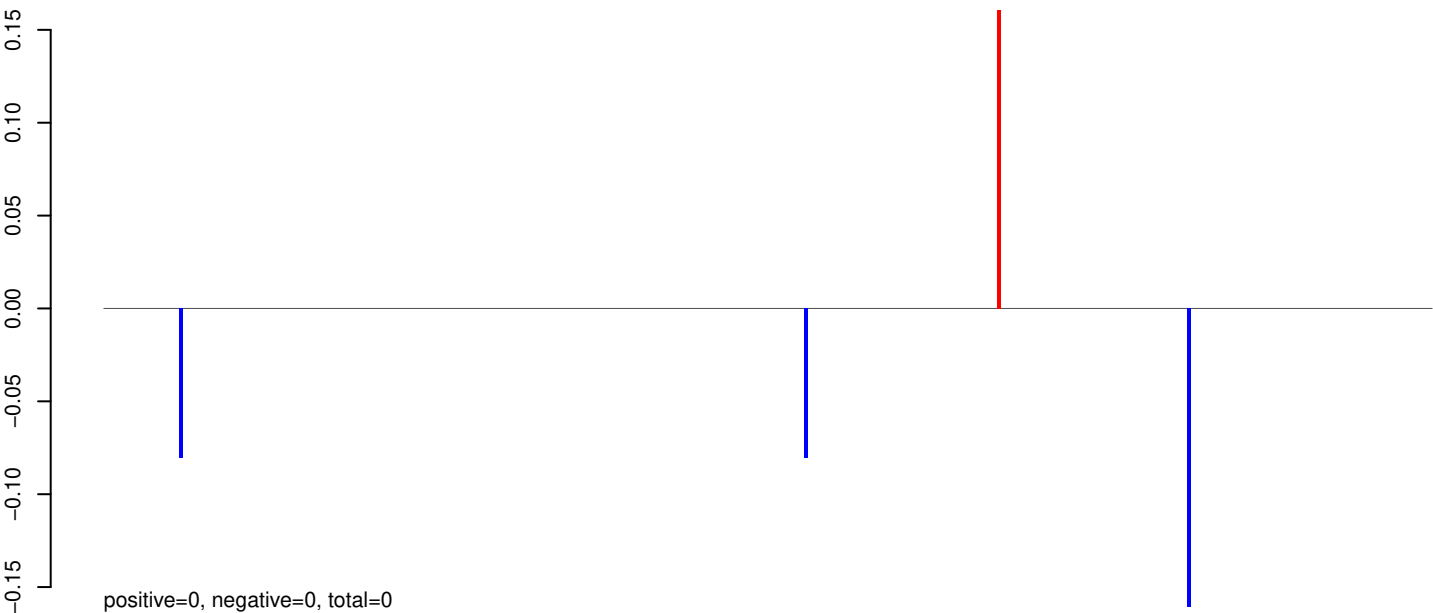
AeAeg_Aag2_mock_dsFluc_RA.rep



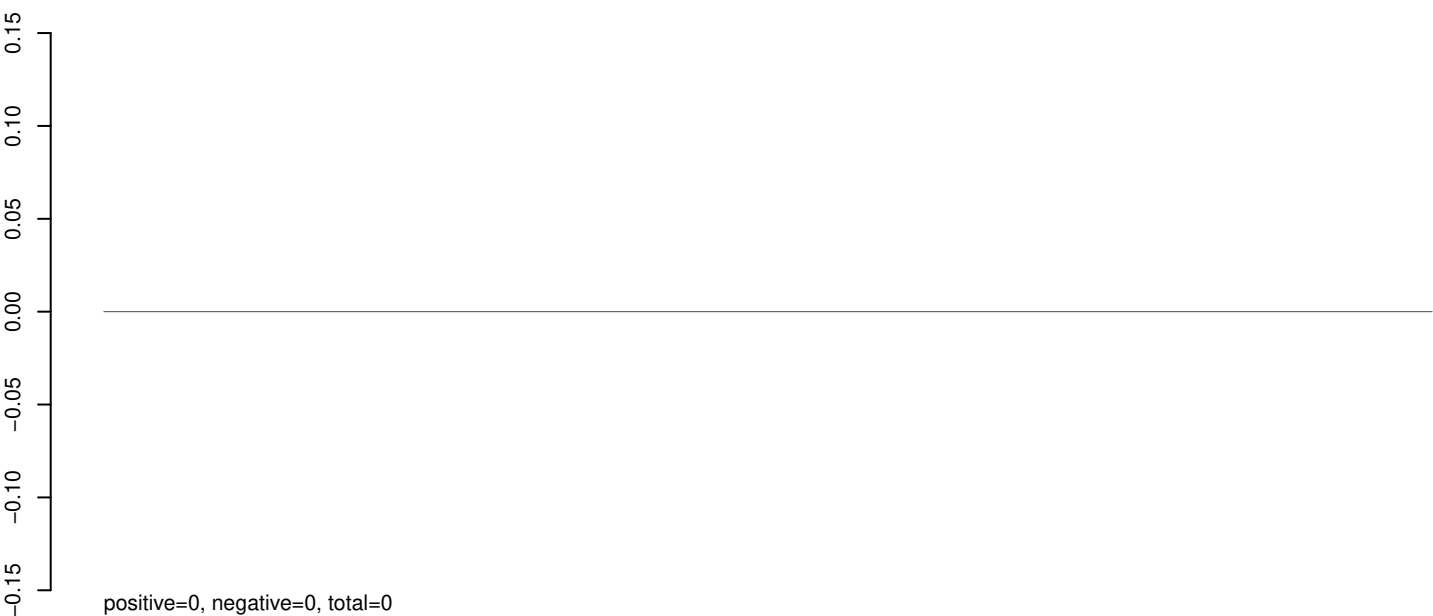
Window size=25, length=11550, virus@NC_024887.1-Middelburg_virus_isolate_ArB-8422:1-11550

0 2000 4000 6000 8000 10000 12000

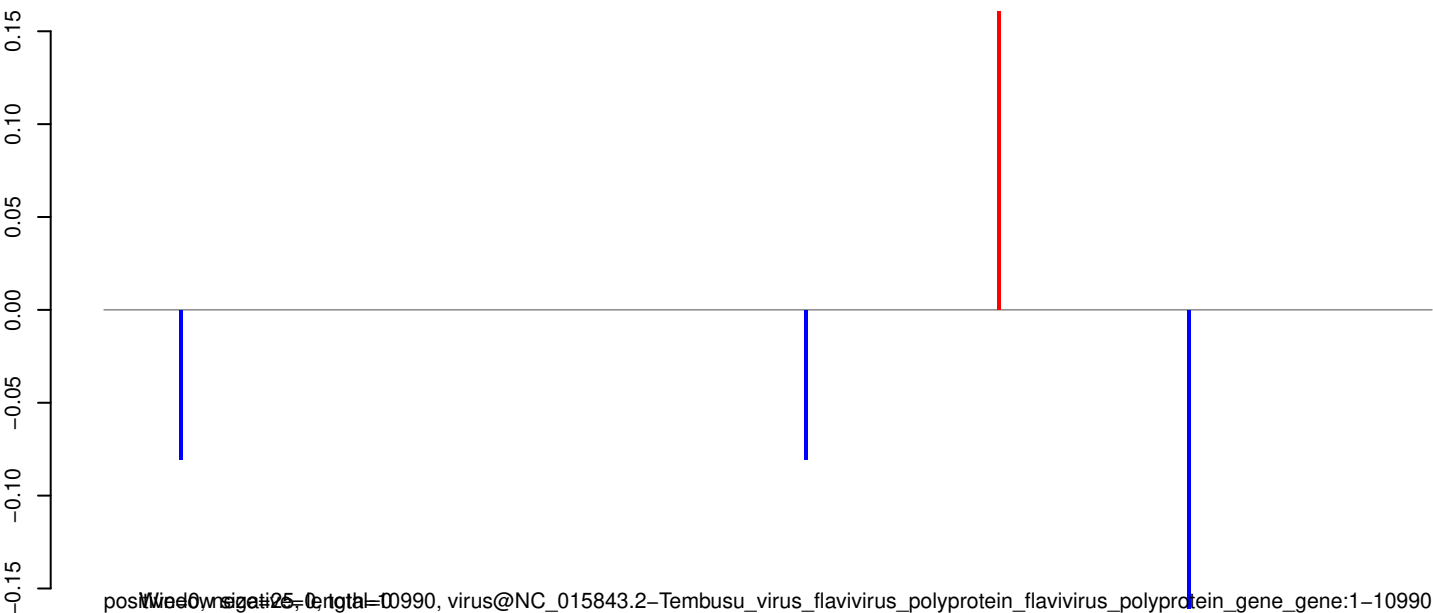
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

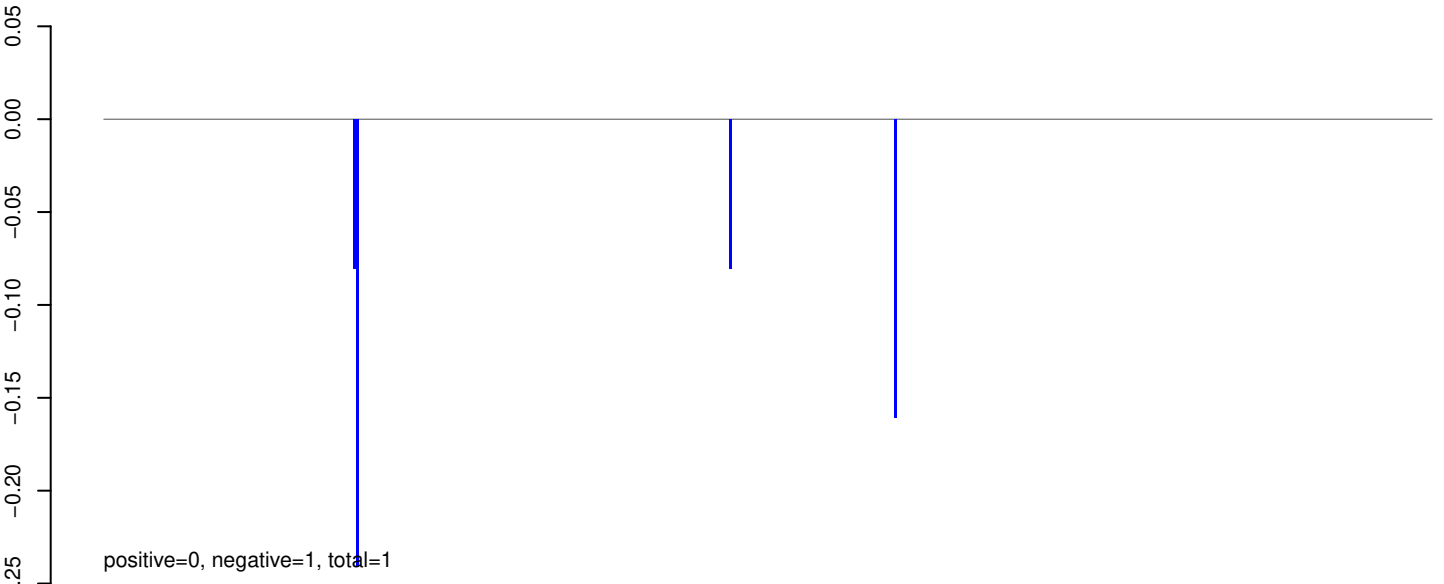


AeAeg_Aag2_mock_dsFluc_RA.rep

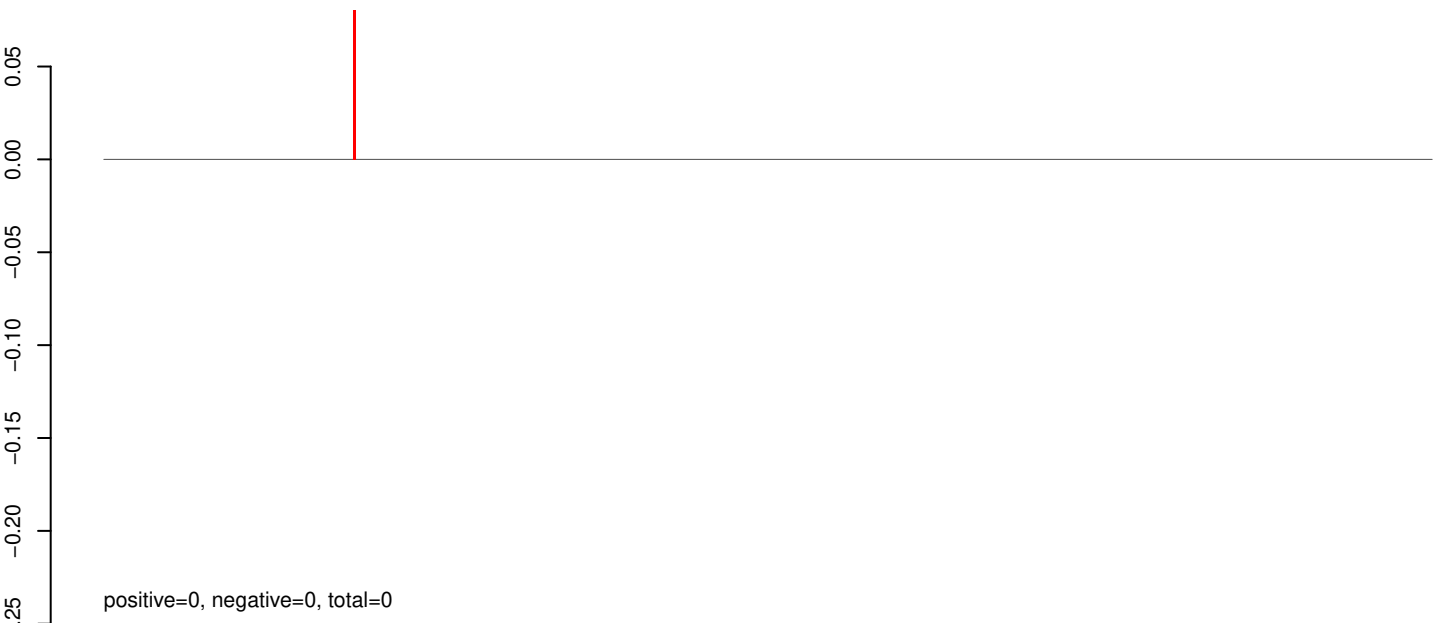


Window size=25, length=10990, virus@NC_015843.2-Tembusu_virus_flavivirus_polyprotein_flavivirus_polyprotein_gene_gene:1-10990

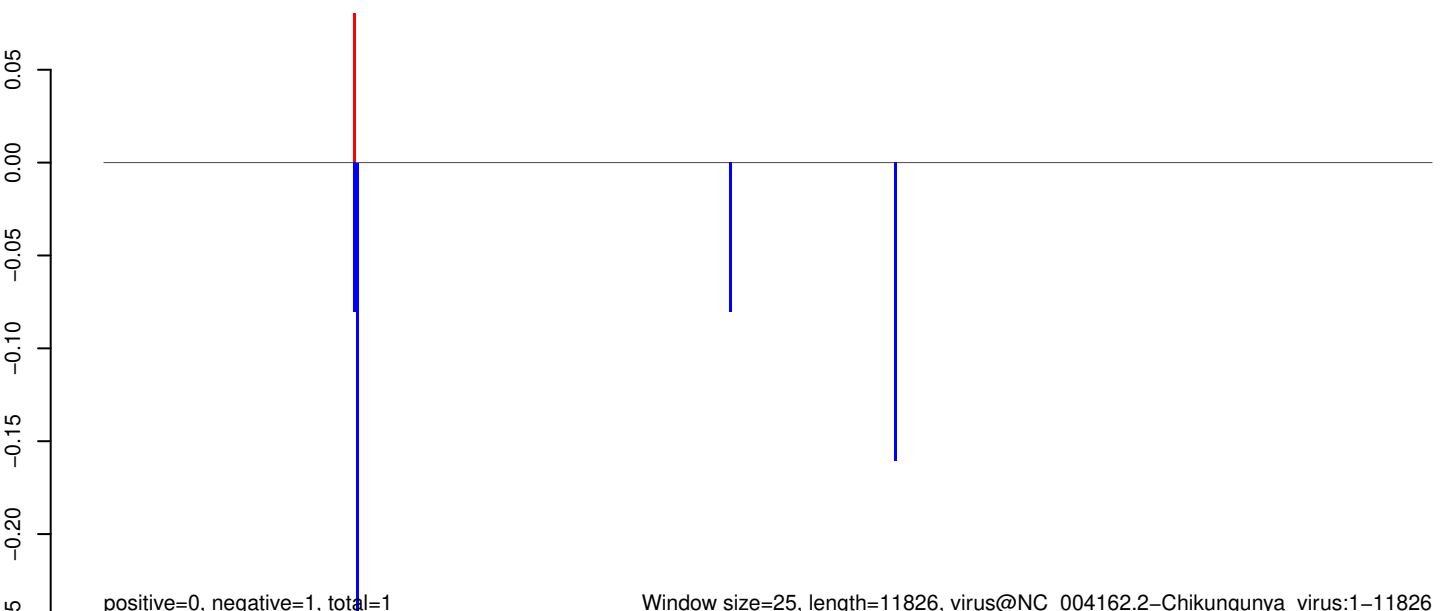
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



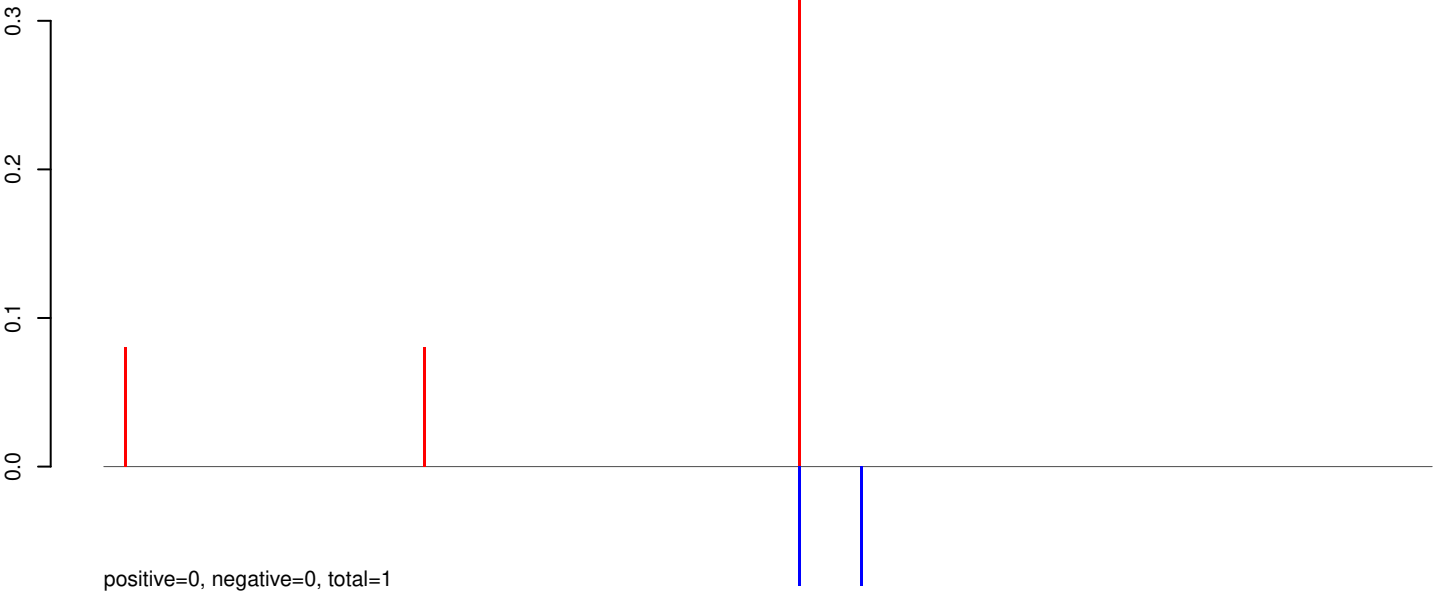
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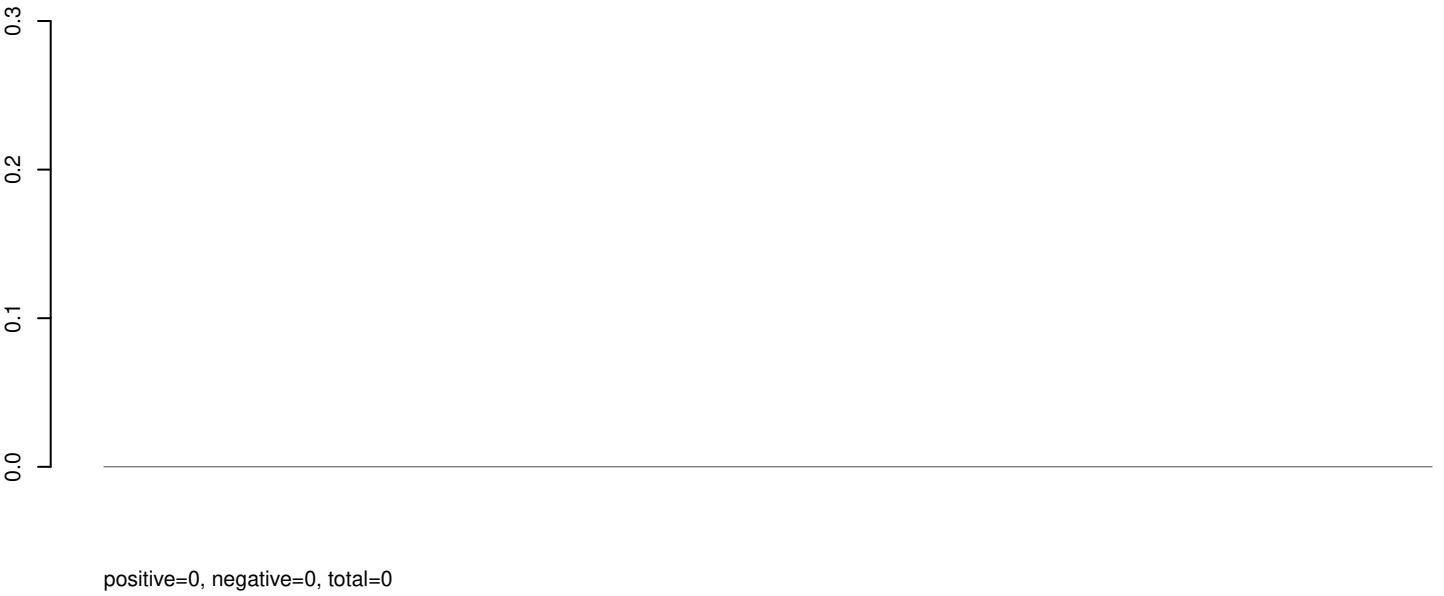
AeAeg_Aag2_mock_dsFluc_RA.rep



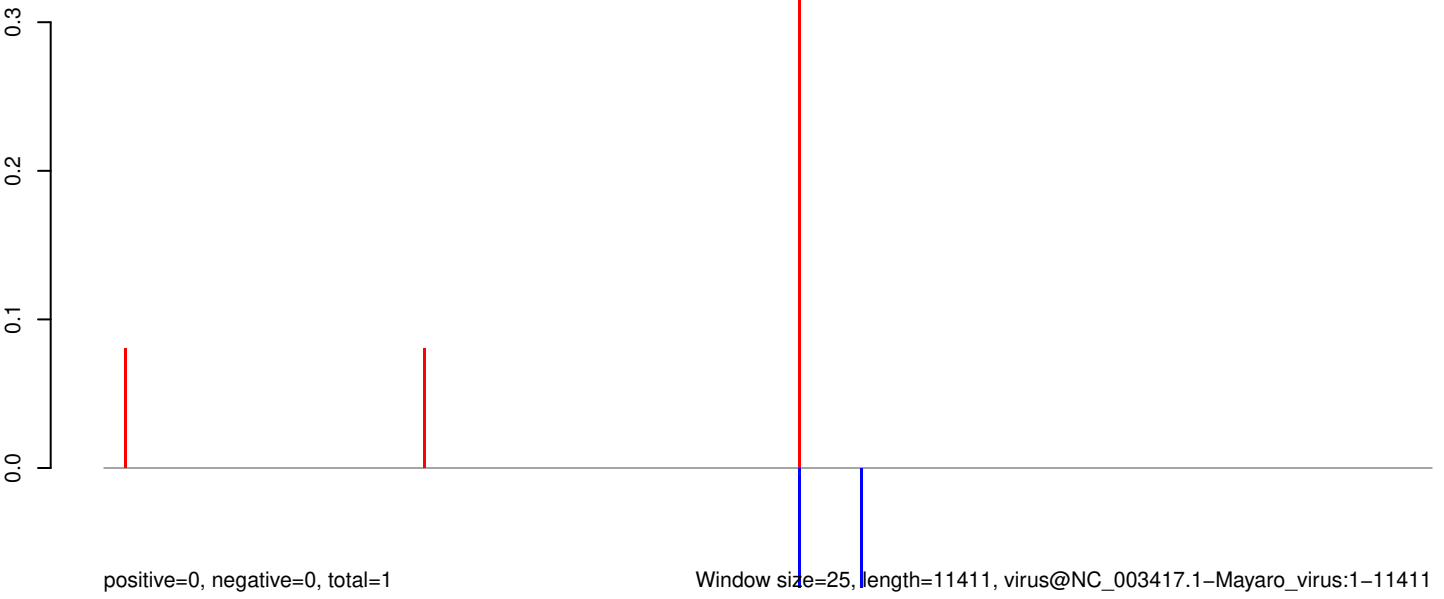
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

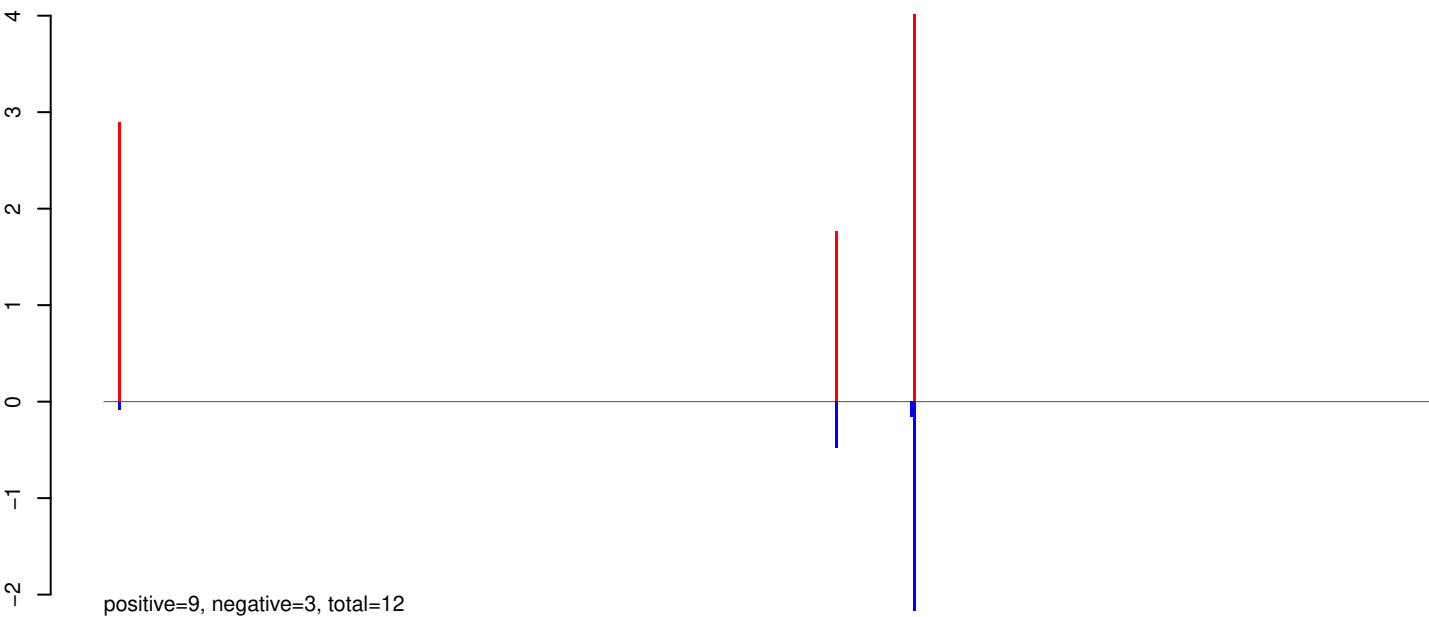


AeAeg_Aag2_mock_dsFluc_RA.rep

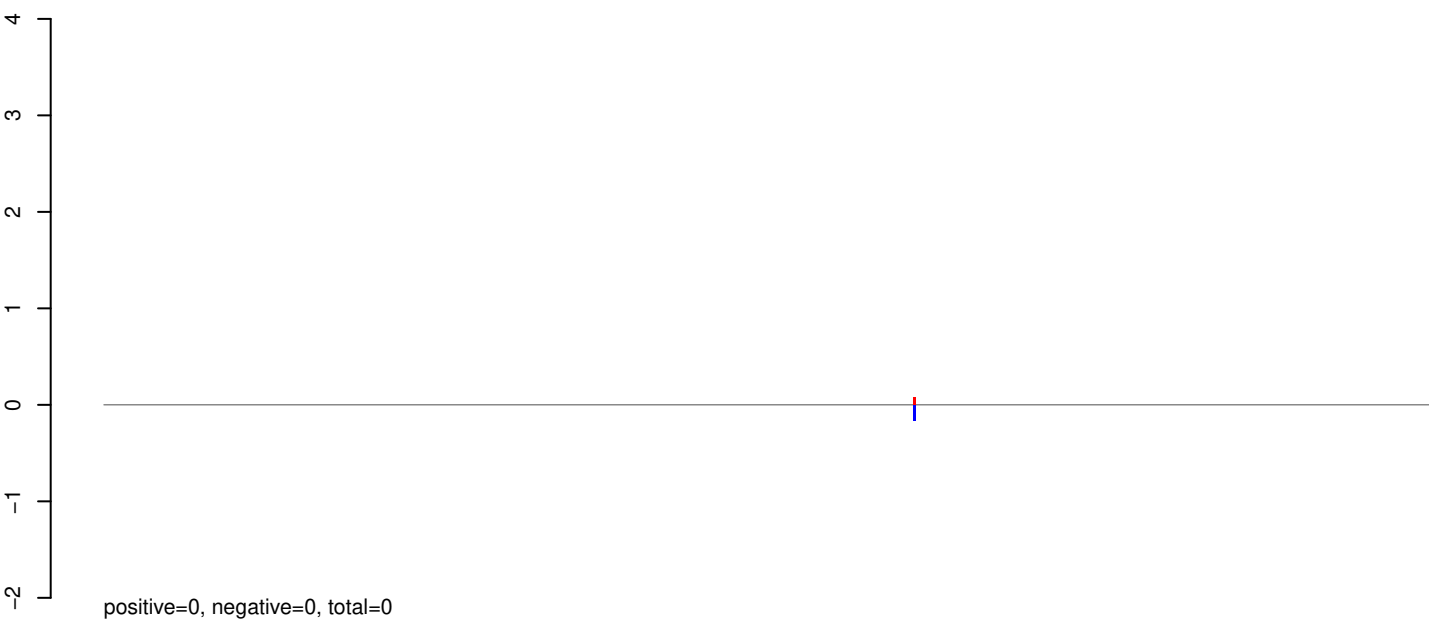


0 2000 4000 6000 8000 10000

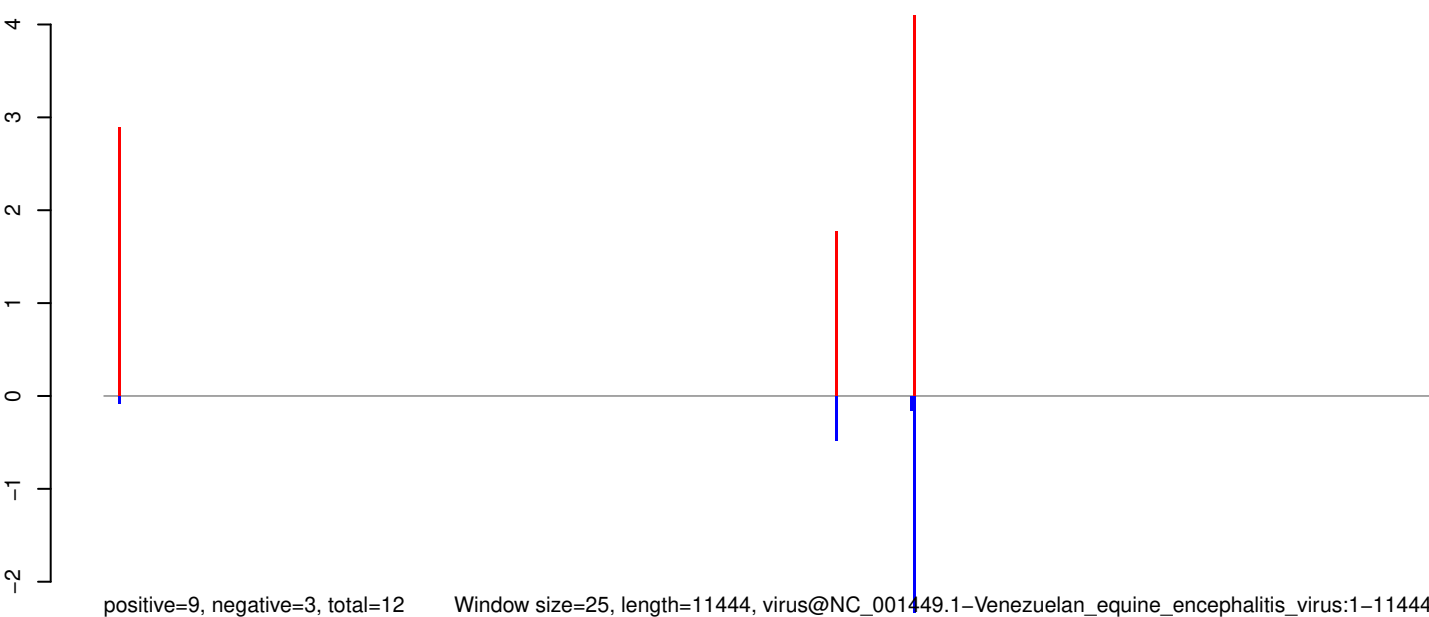
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



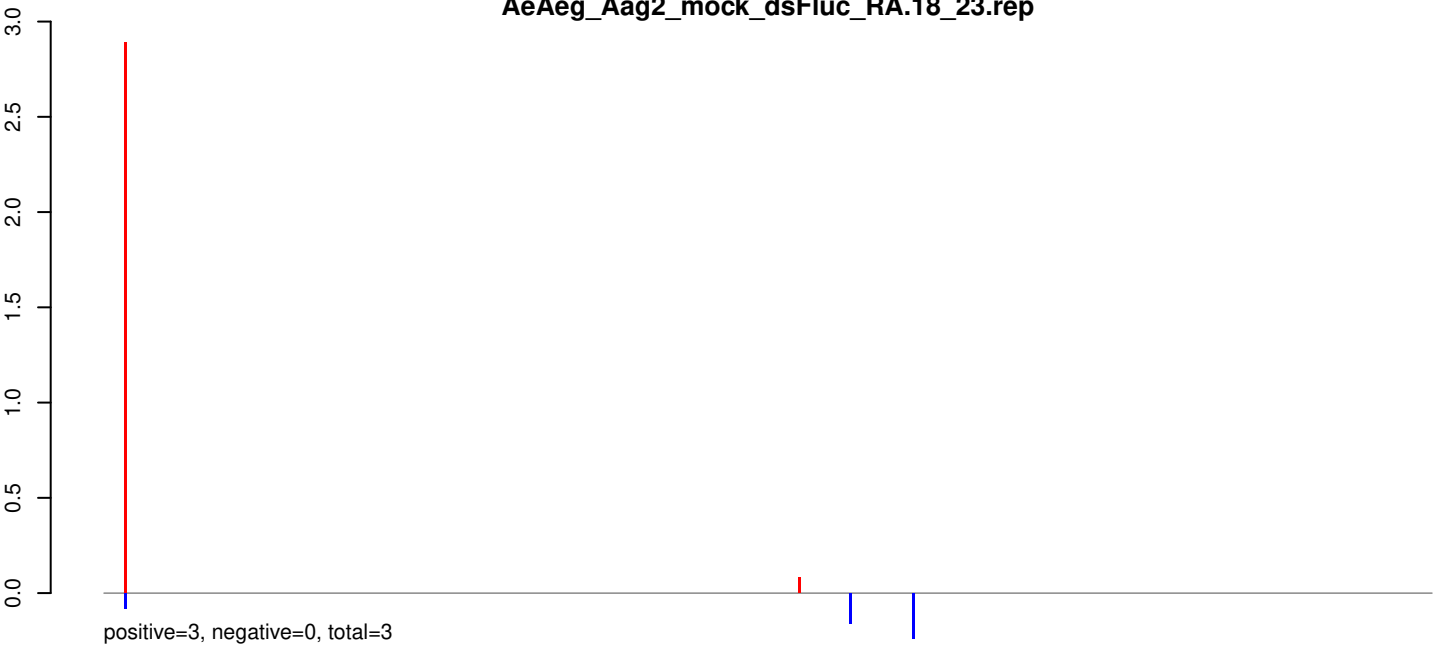
AeAeg_Aag2_mock_dsFluc_RA.rep



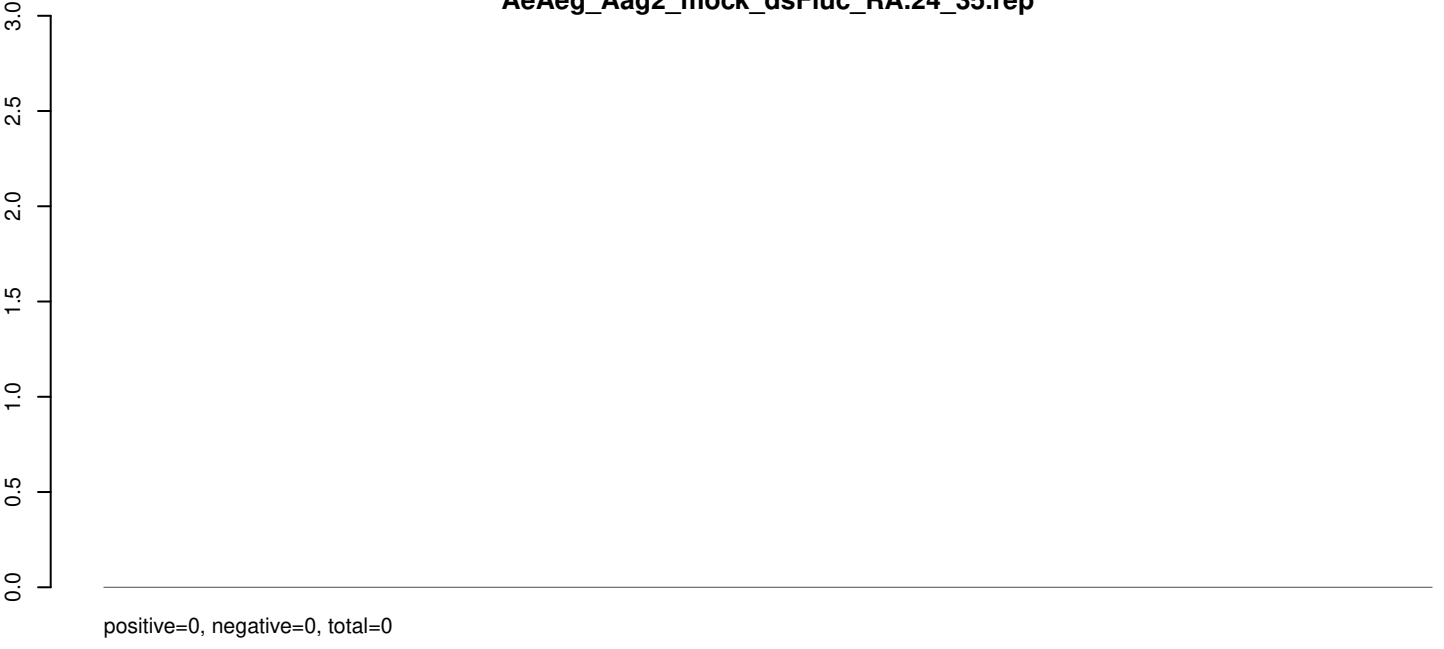
Window size=25, length=11444, virus@NC_001449.1-Venezuelan_equine_encephalitis_virus:1-11444

0 2000 4000 6000 8000 10000

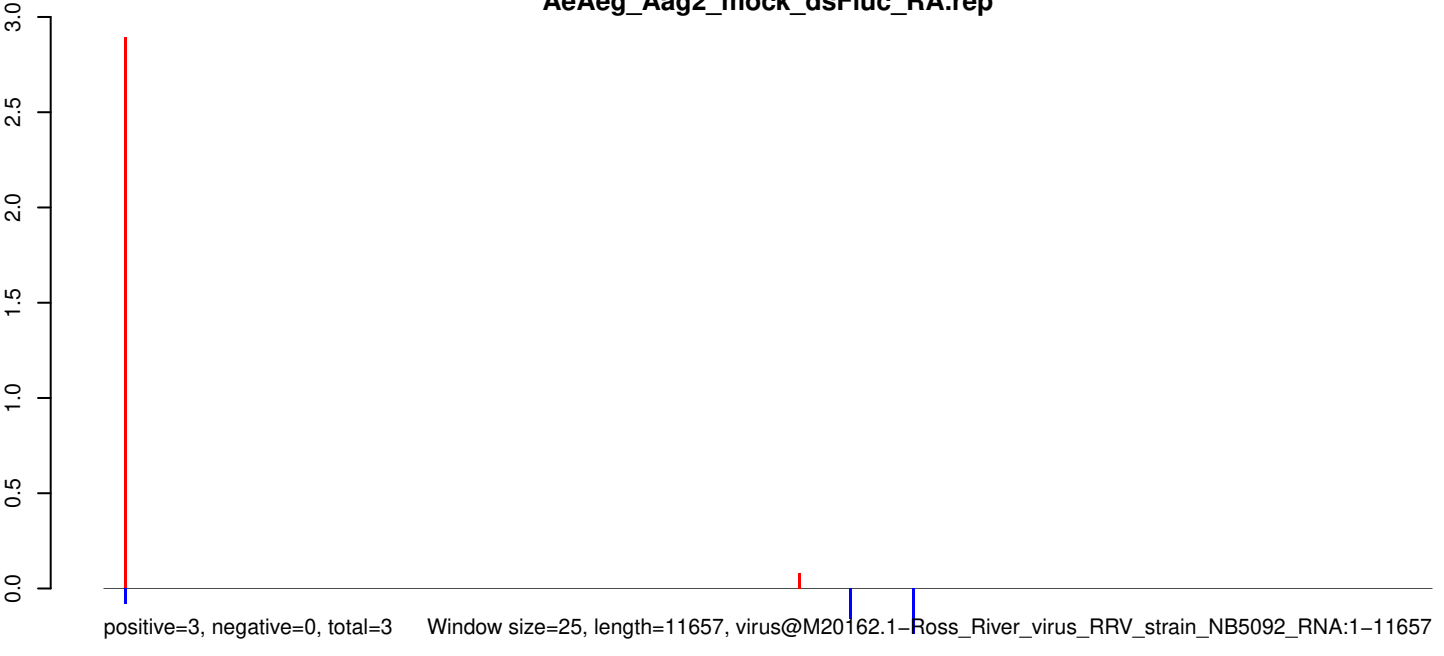
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



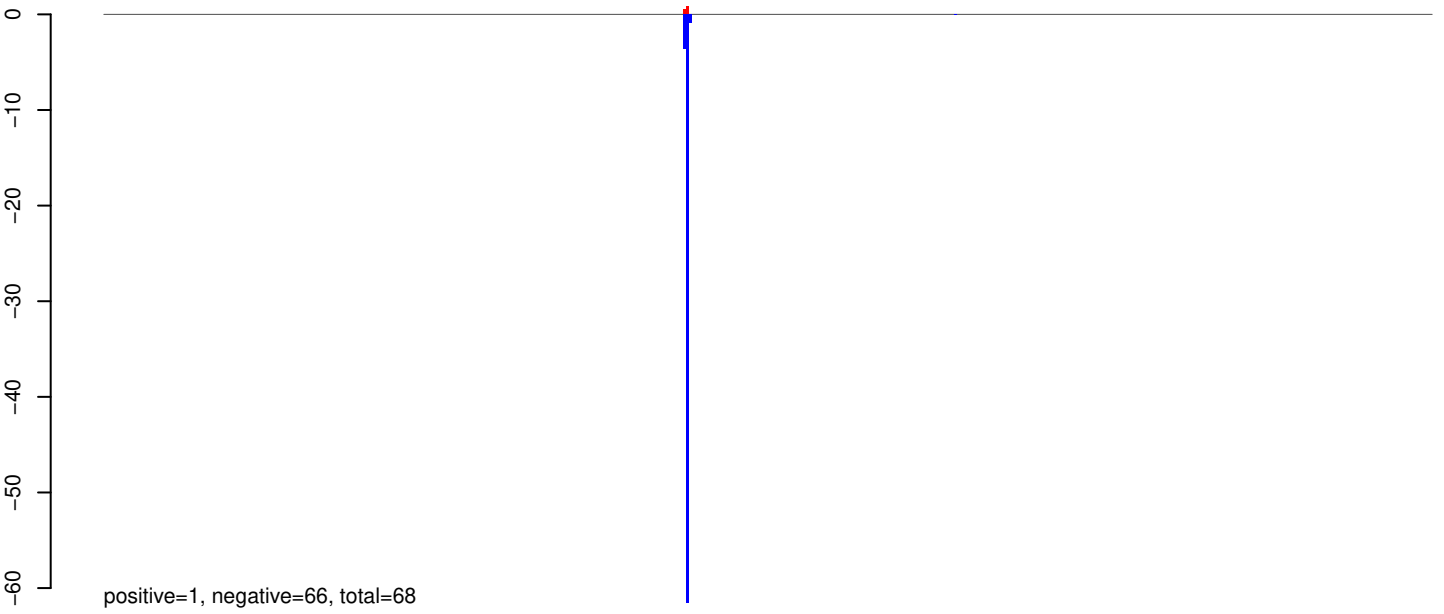
AeAeg_Aag2_mock_dsFluc_RA.rep



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

0 2000 4000 6000 8000 10000 12000

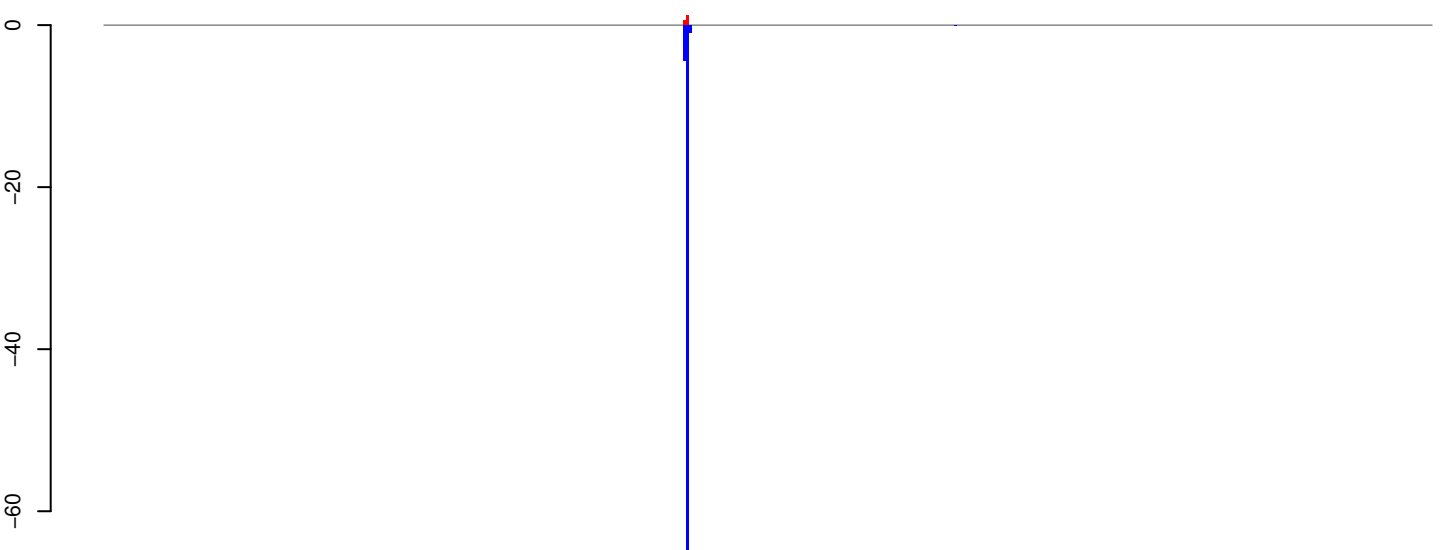
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



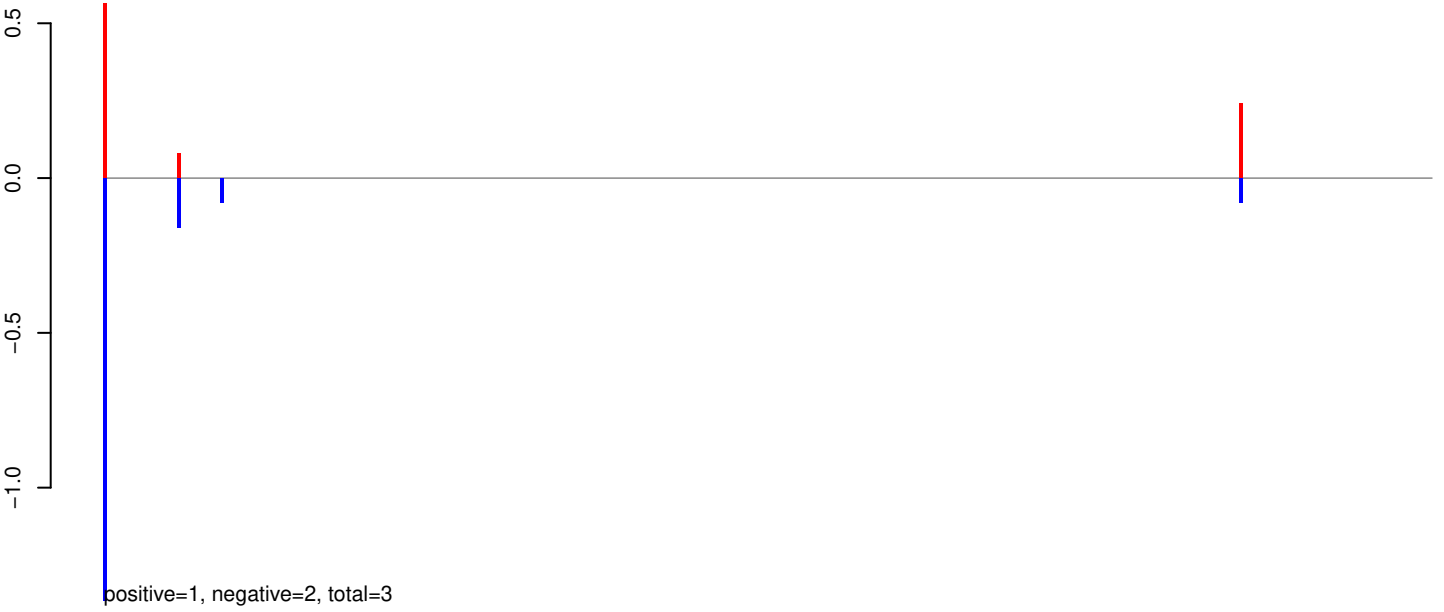
AeAeg_Aag2_mock_dsFluc_RA.rep



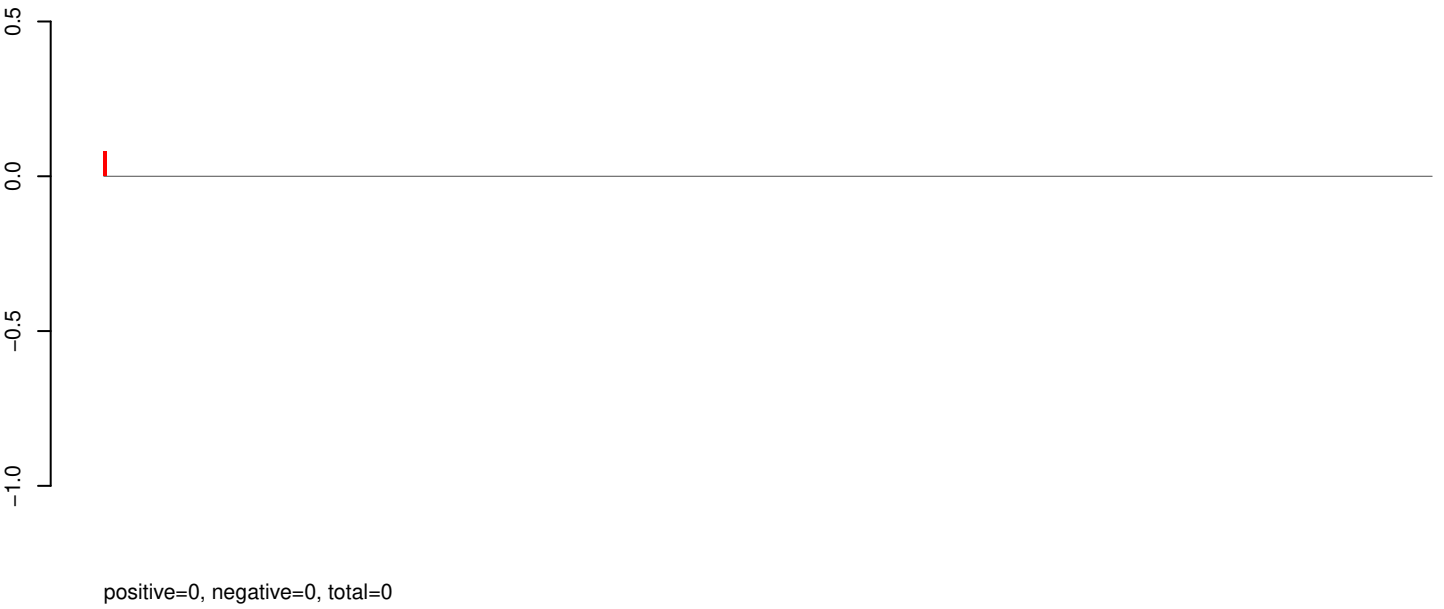
Wendy, size=25, total=68, virus@KJ716848.1-Ngari_virus_strain_Adrar_segment_S_plus_segment_M_plus_segment_L:1-12369

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



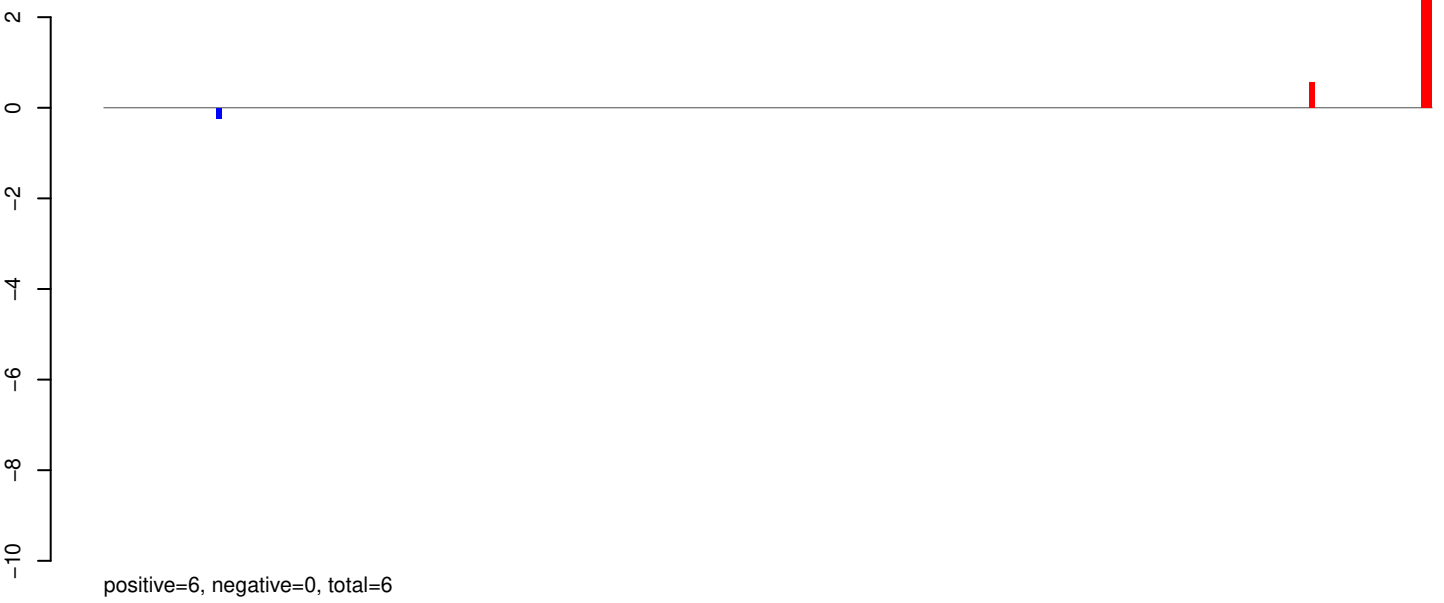
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



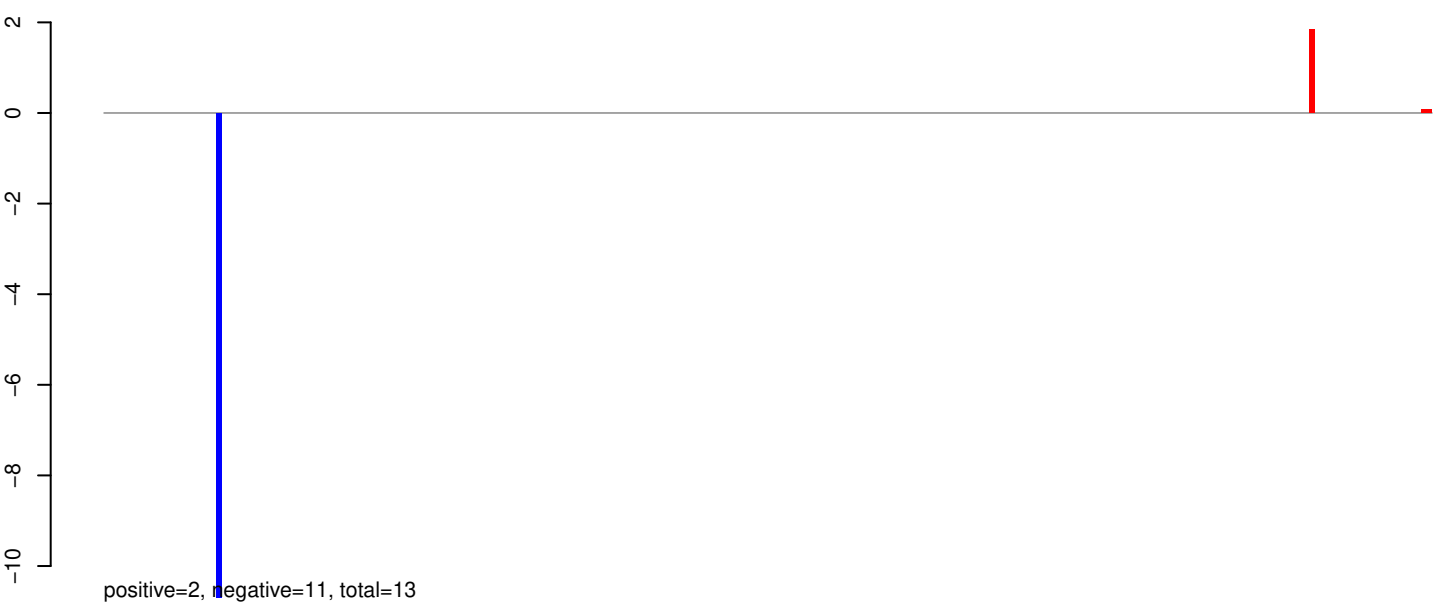
AeAeg_Aag2_mock_dsFluc_RA.rep



AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



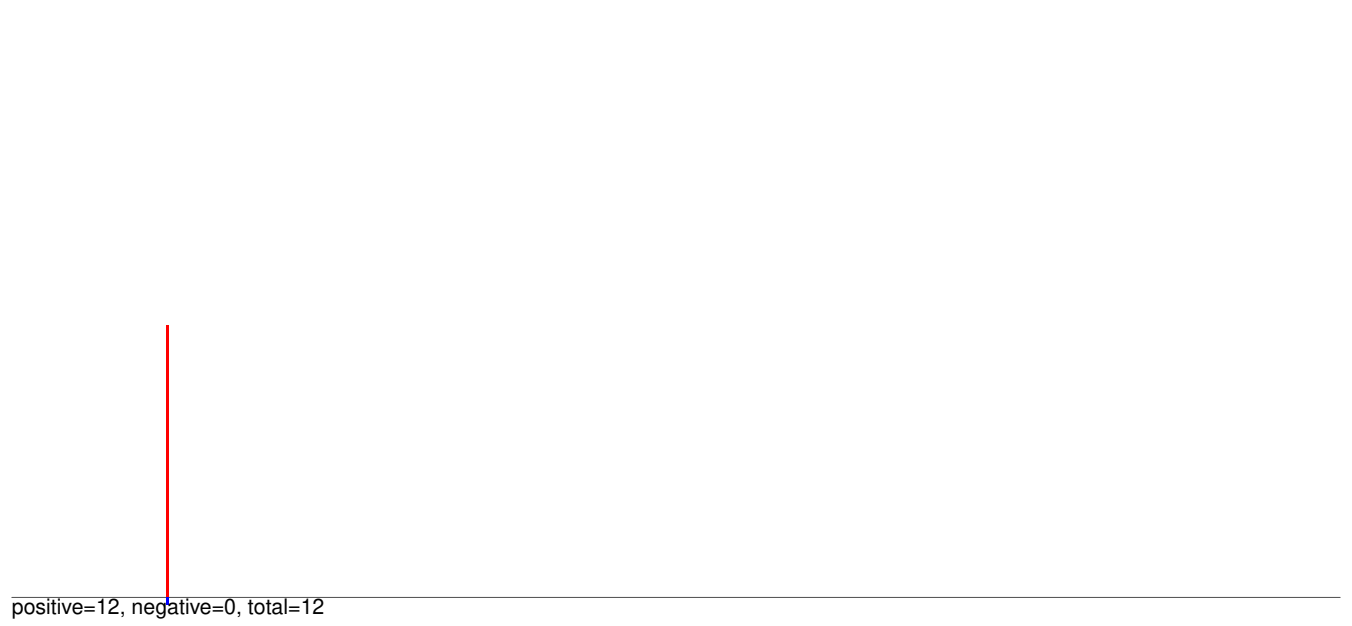
AeAeg_Aag2_mock_dsFluc_RA.rep



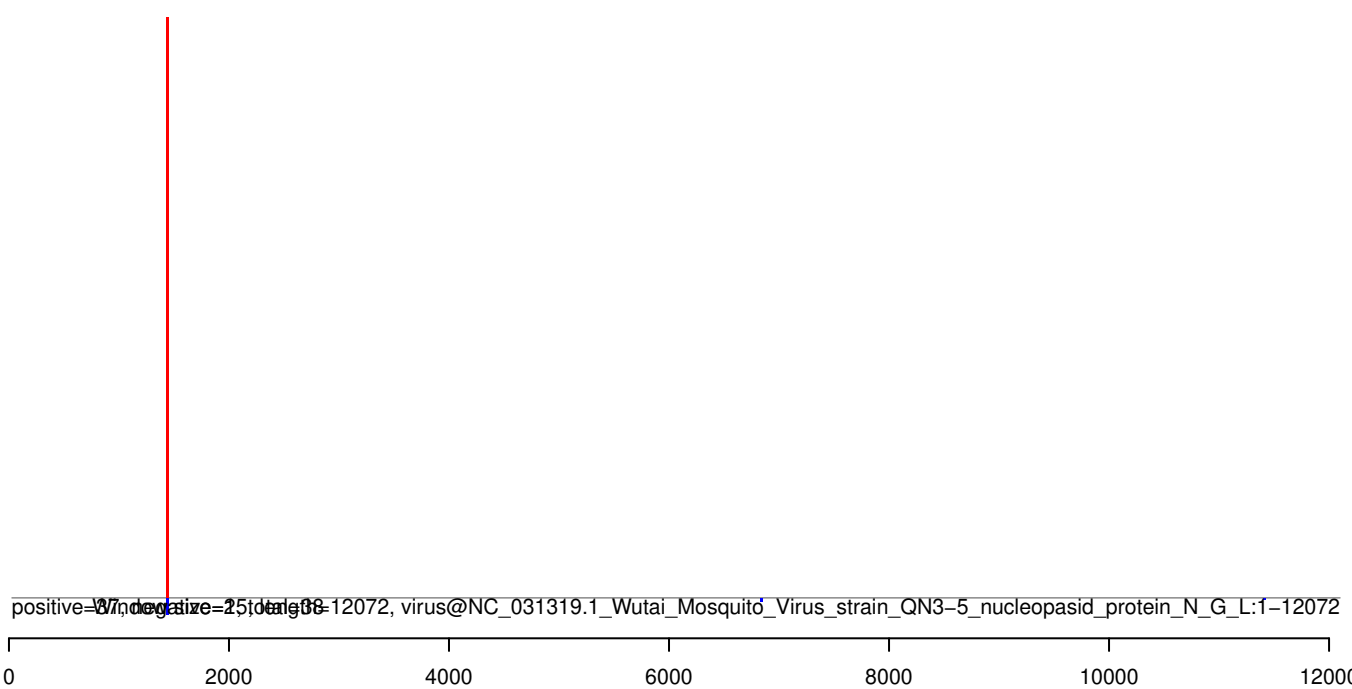
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

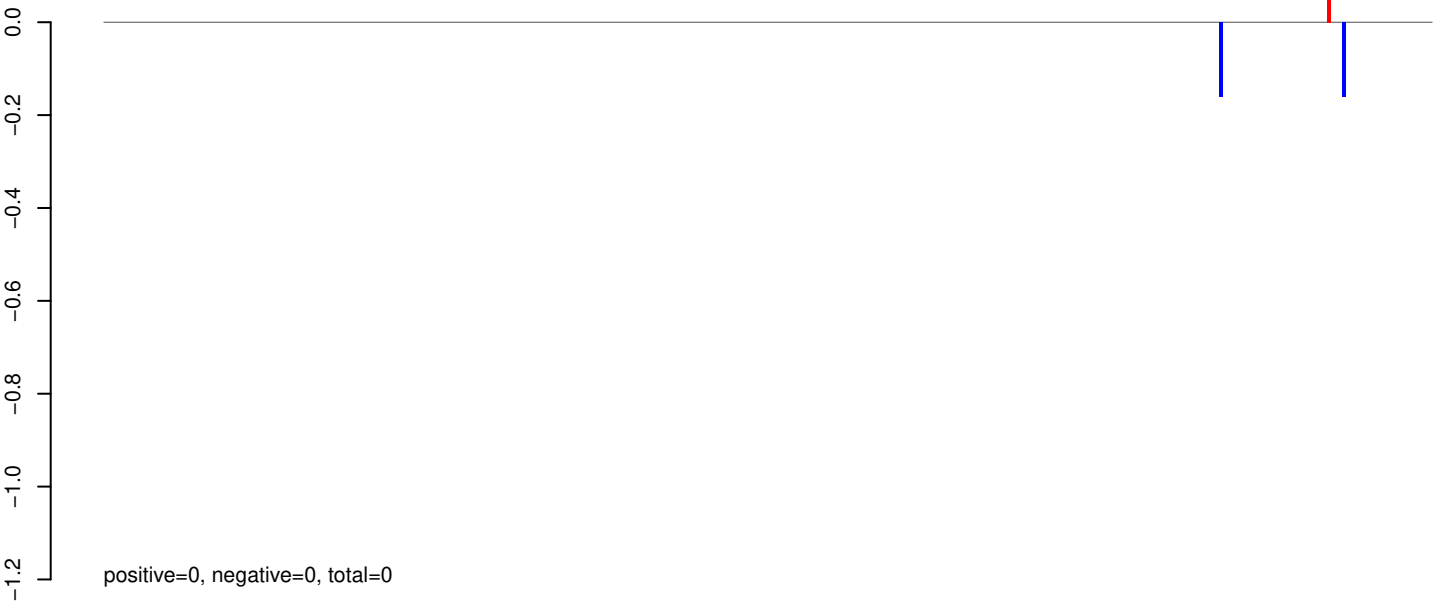


AeAeg_Aag2_mock_dsFluc_RA.rep

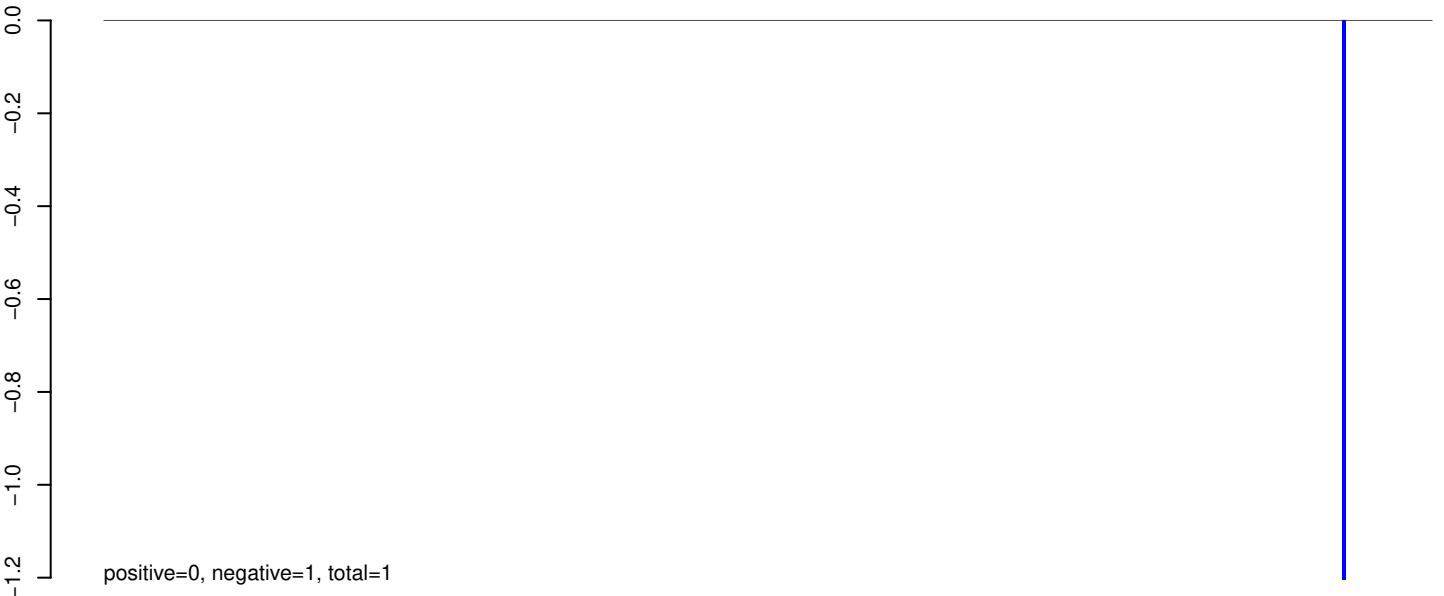


1-12072, virus@NC_031319.1_Wutai_Mosquito_Virus_strain_QN3-5_nucleopasid_protein_N_G_L:1-12072

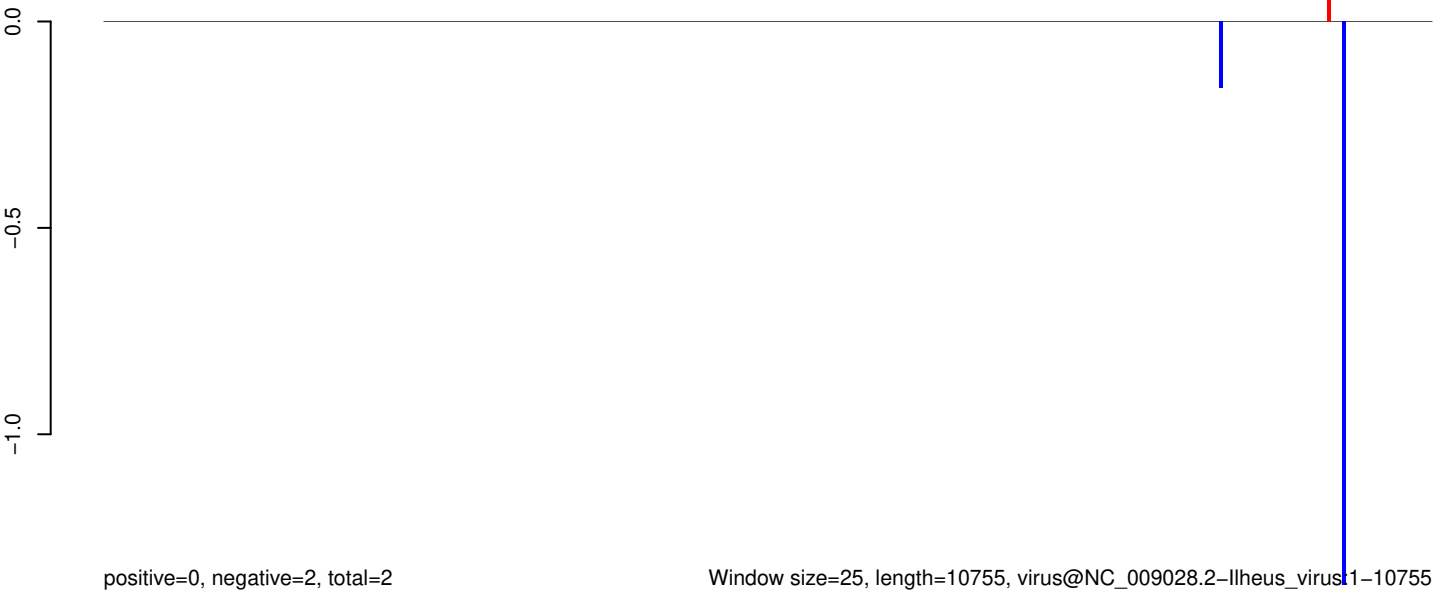
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



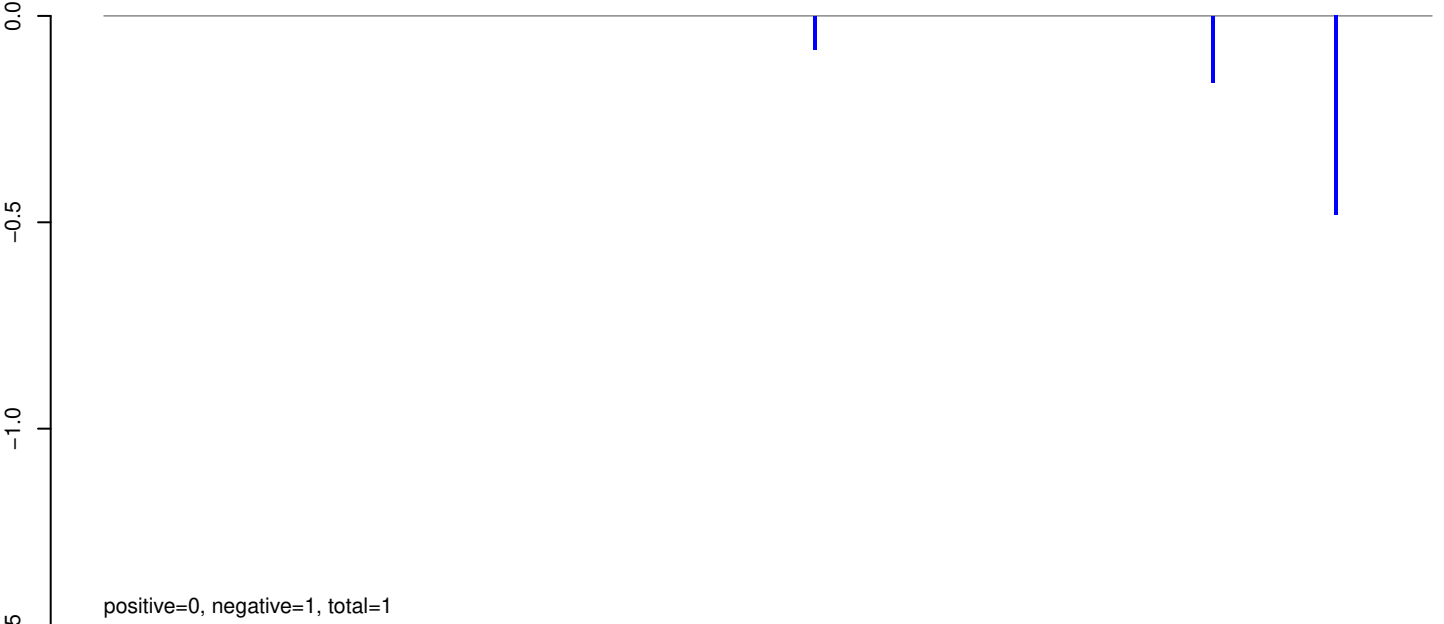
AeAeg_Aag2_mock_dsFluc_RA.rep



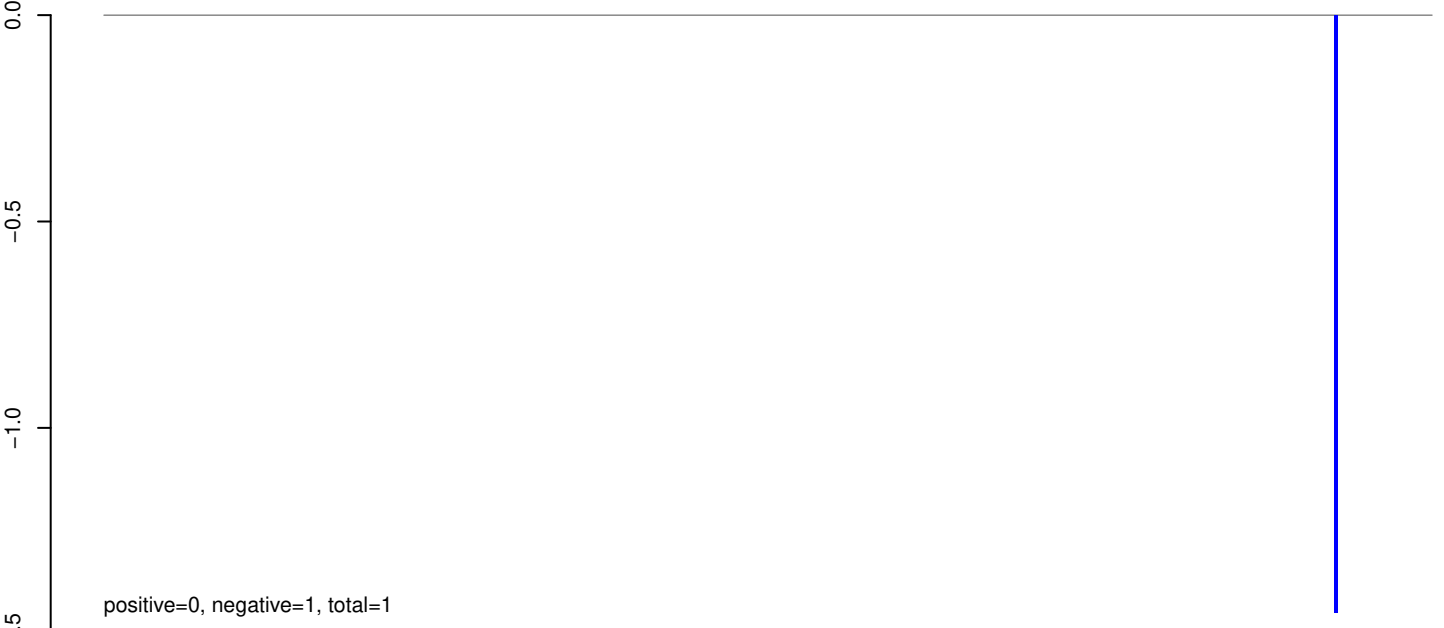
Window size=25, length=10755, virus@NC_009028.2-Ilheus_virus1-10755

0 2000 4000 6000 8000 10000

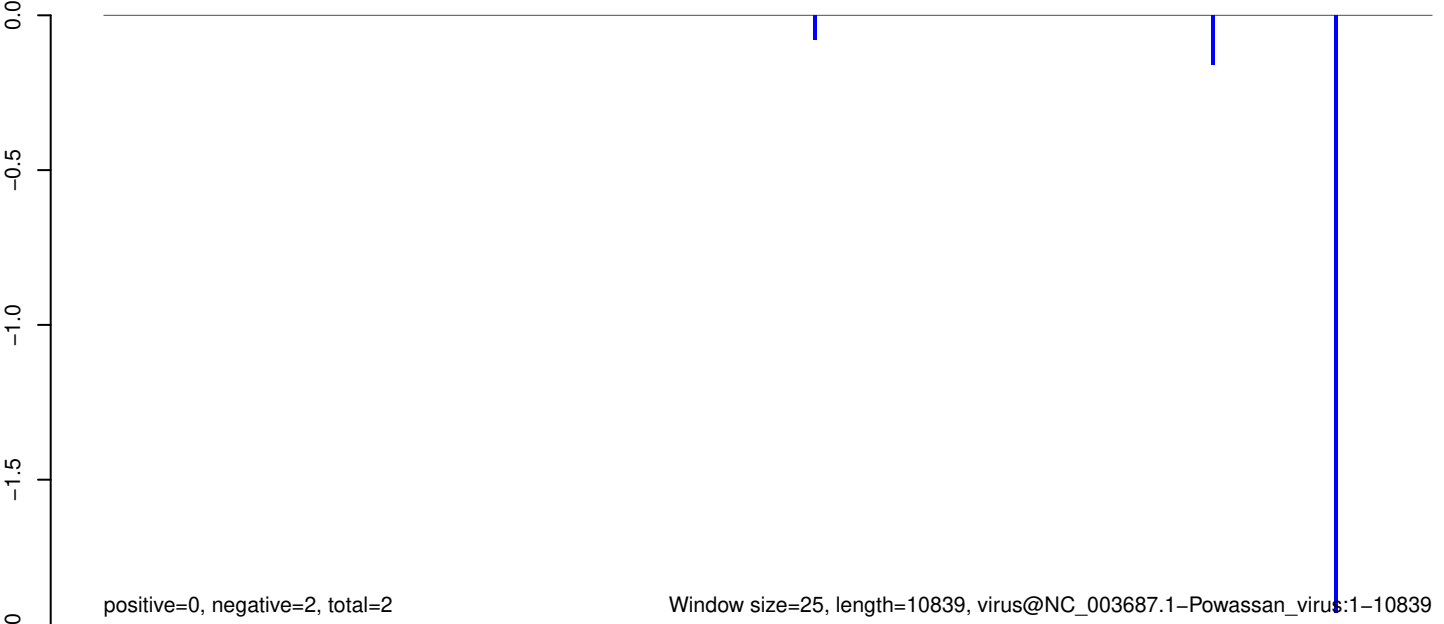
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



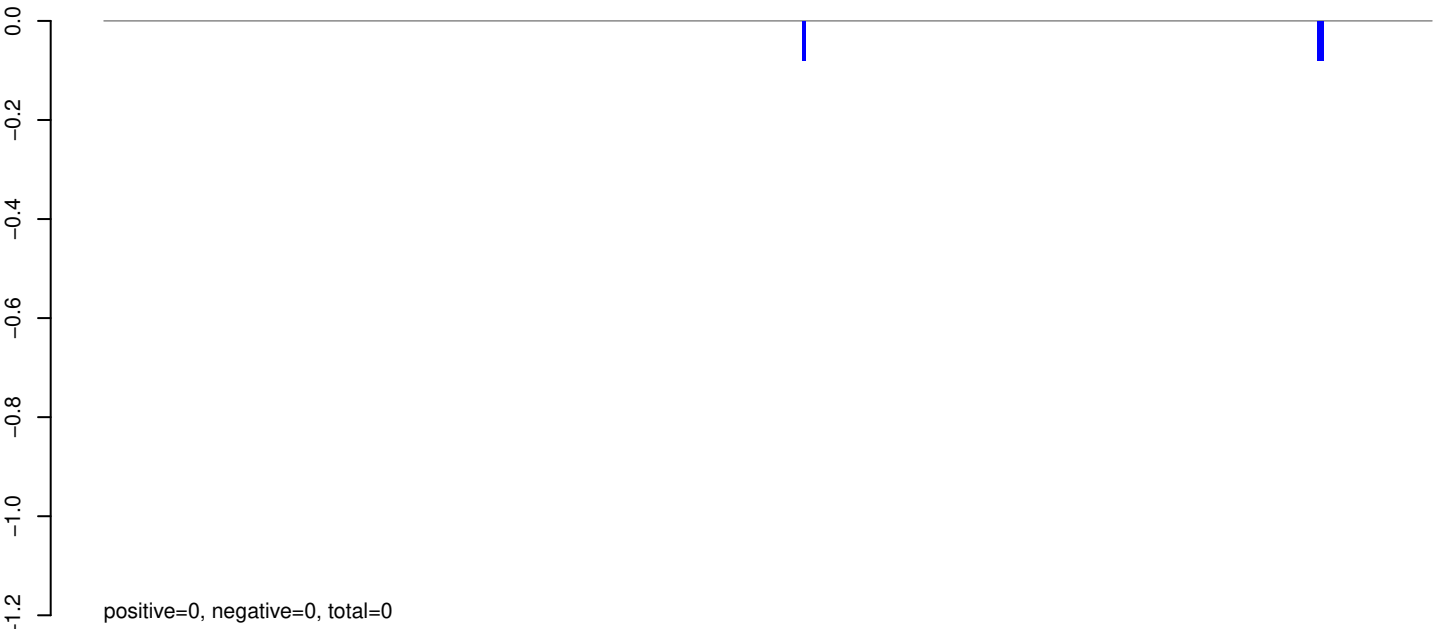
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



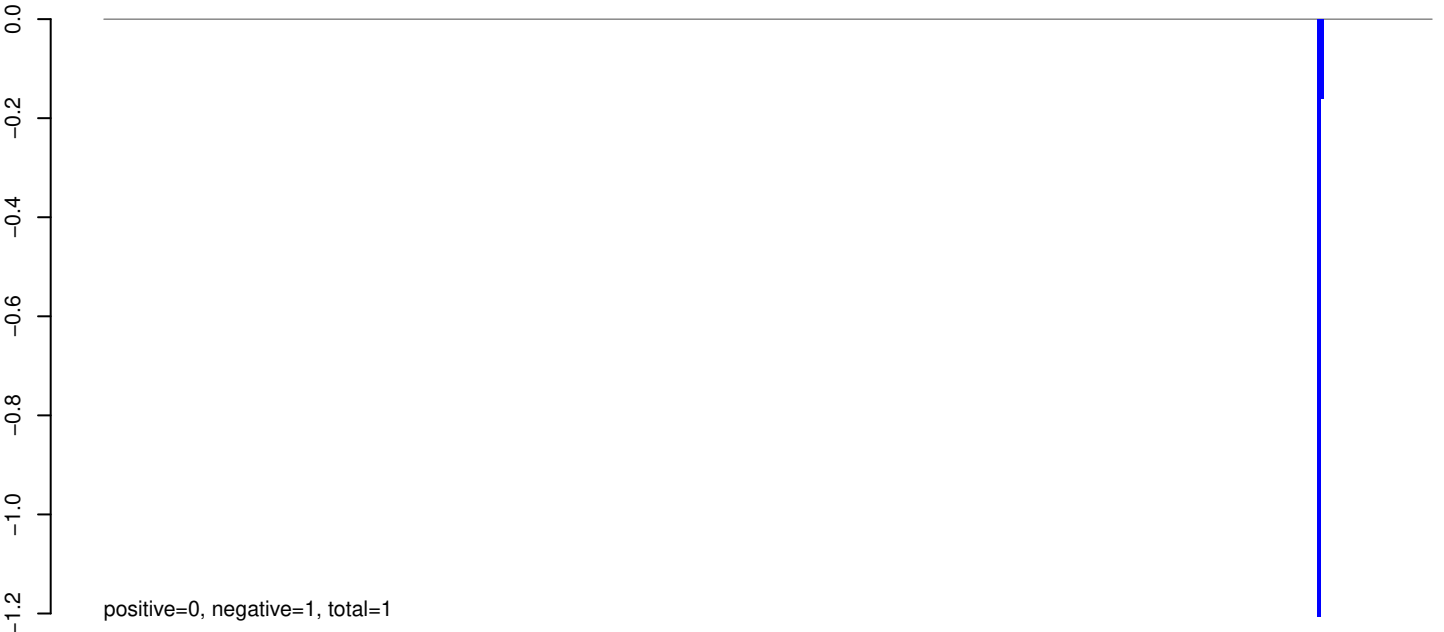
AeAeg_Aag2_mock_dsFluc_RA.rep



AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



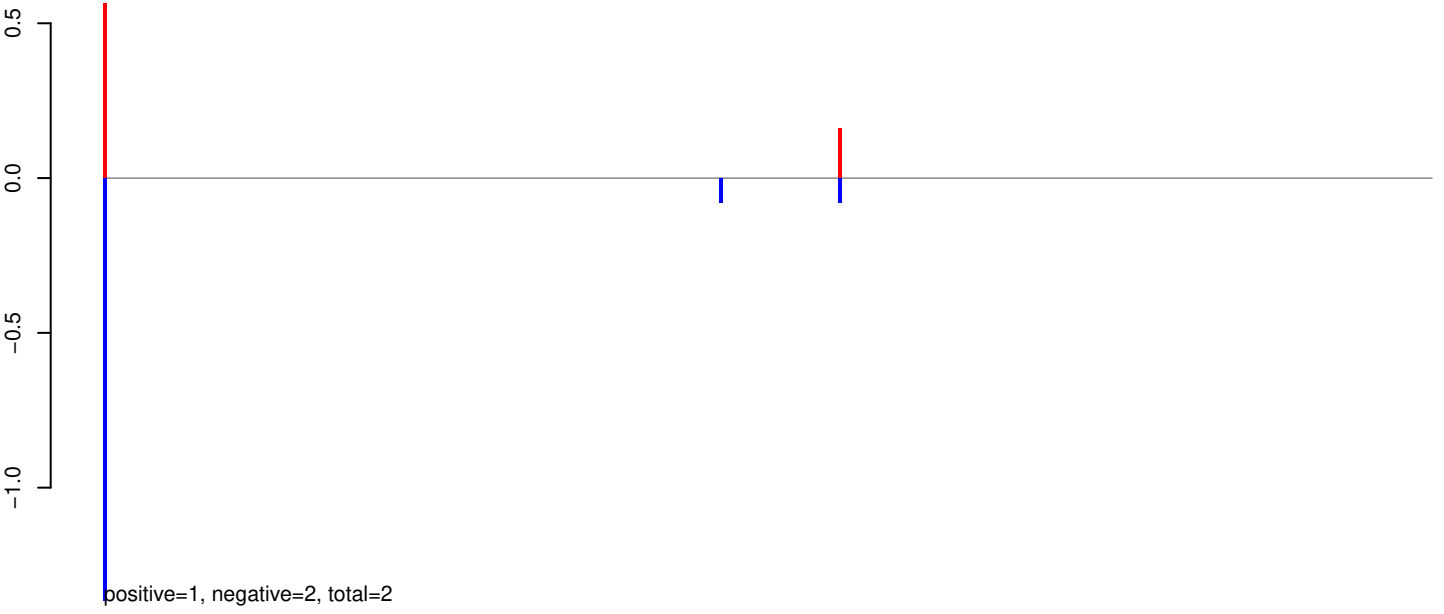
AeAeg_Aag2_mock_dsFluc_RA.rep



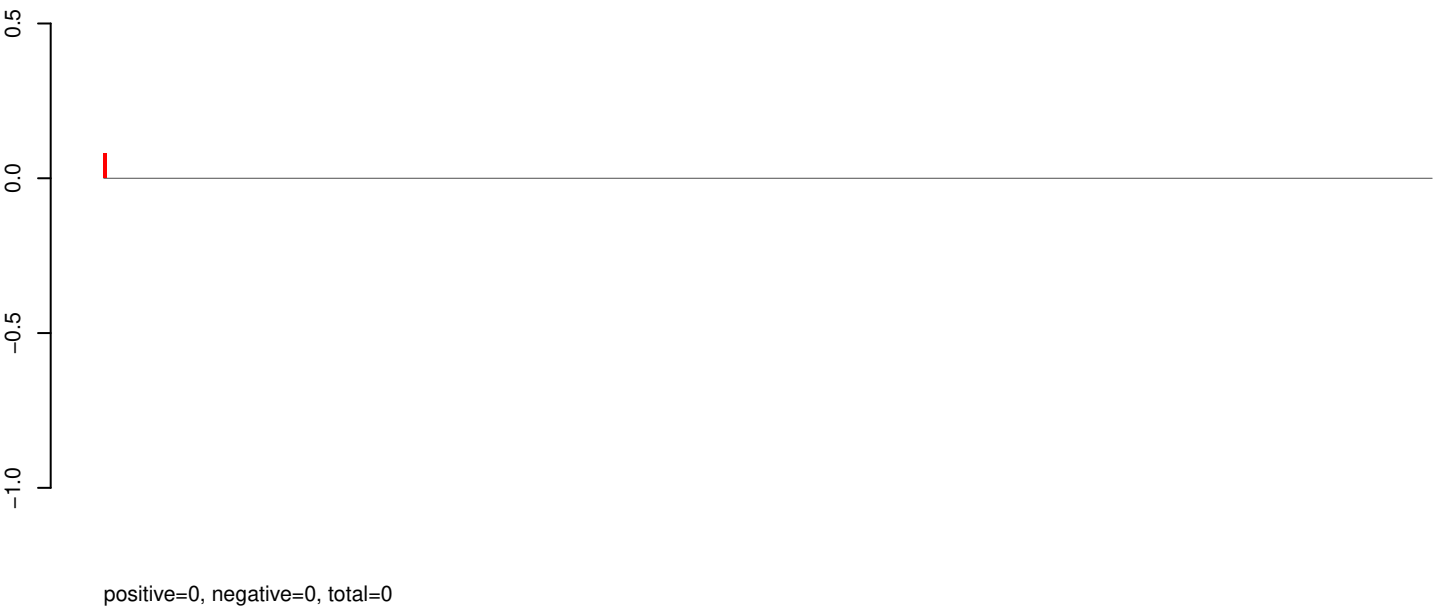
Window size=25, length=11014, virus@NC_000943.1-Murray_Valley_encephalitis_virus:1-11014

0 2000 4000 6000 8000 10000

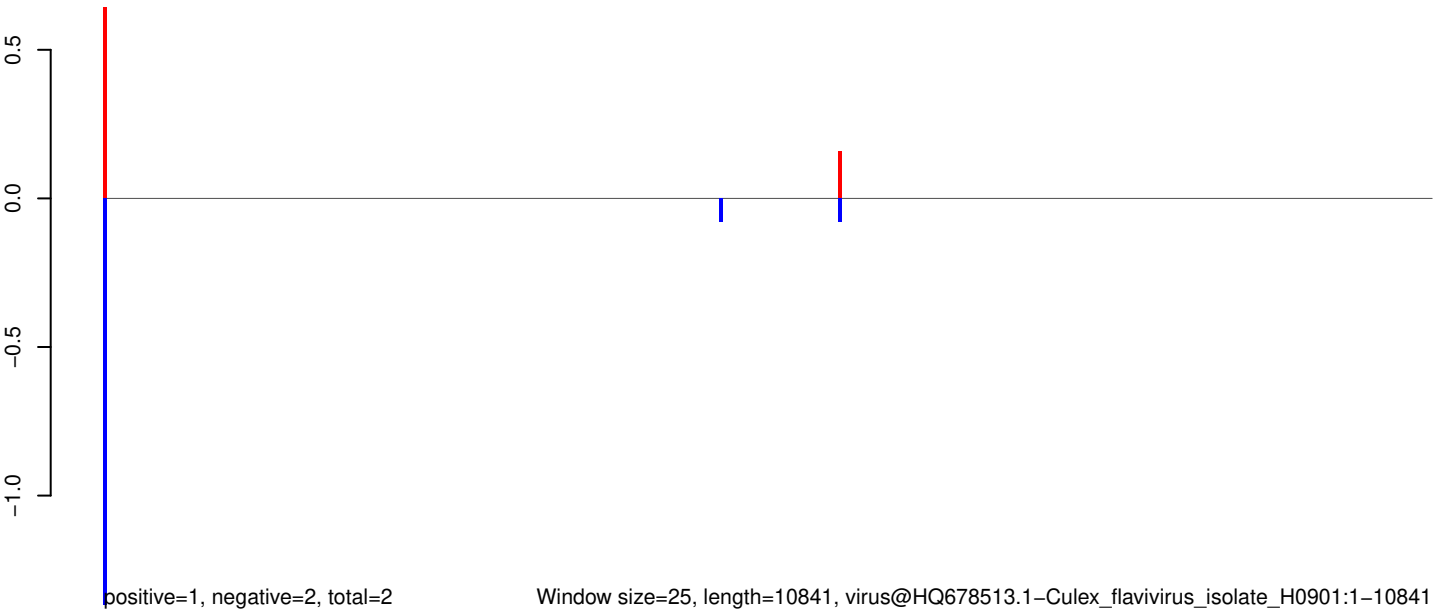
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

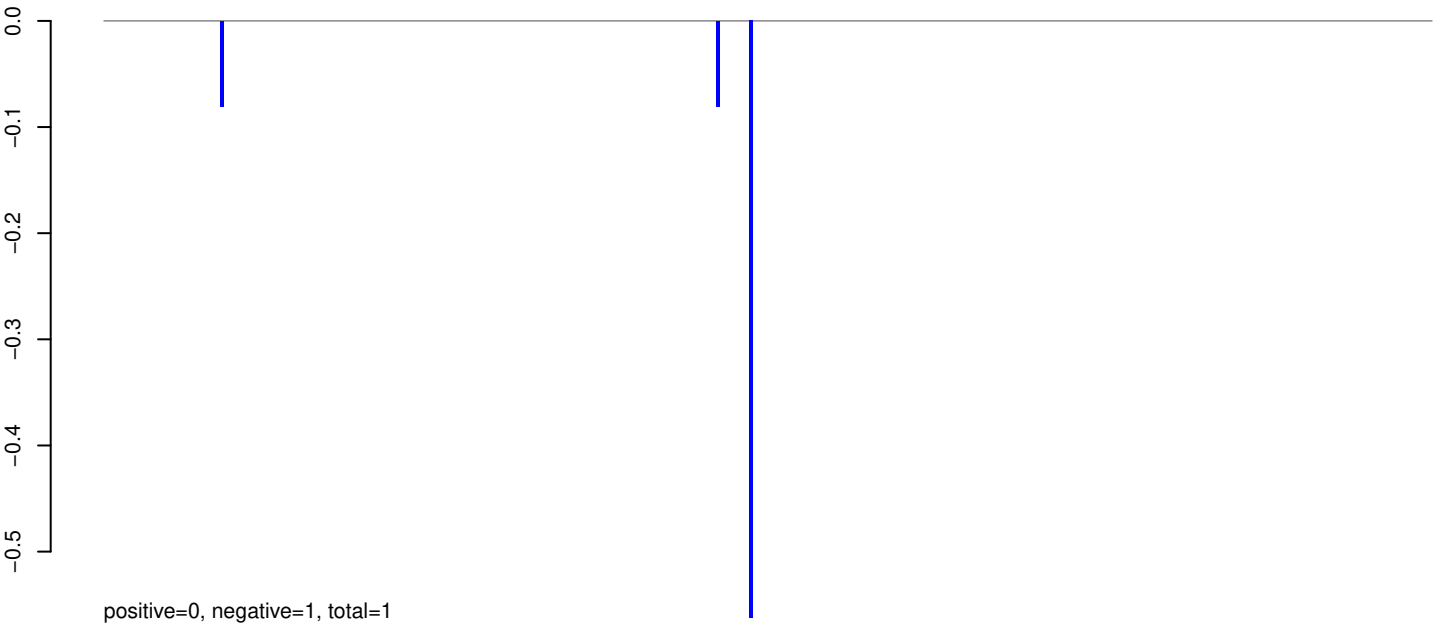


AeAeg_Aag2_mock_dsFluc_RA.rep

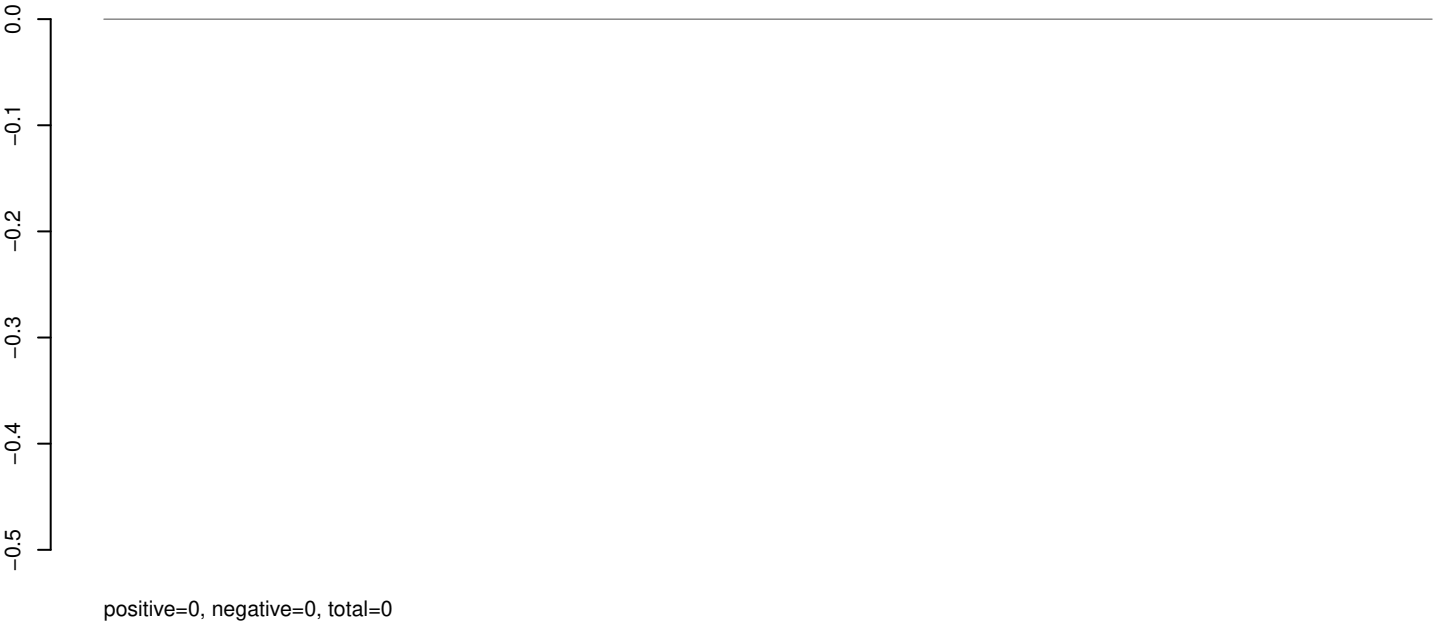


Window size=25, length=10841, virus@HQ678513.1-Culex_flavivirus_isolate_H0901:1-10841

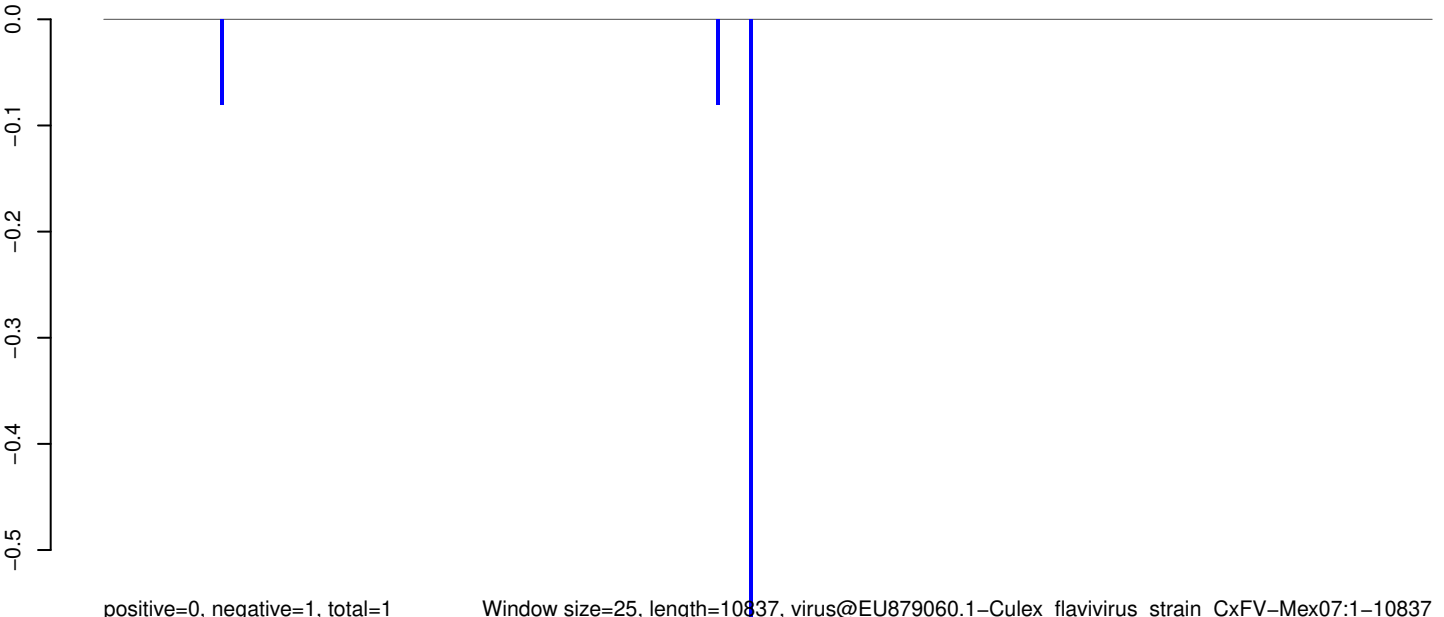
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



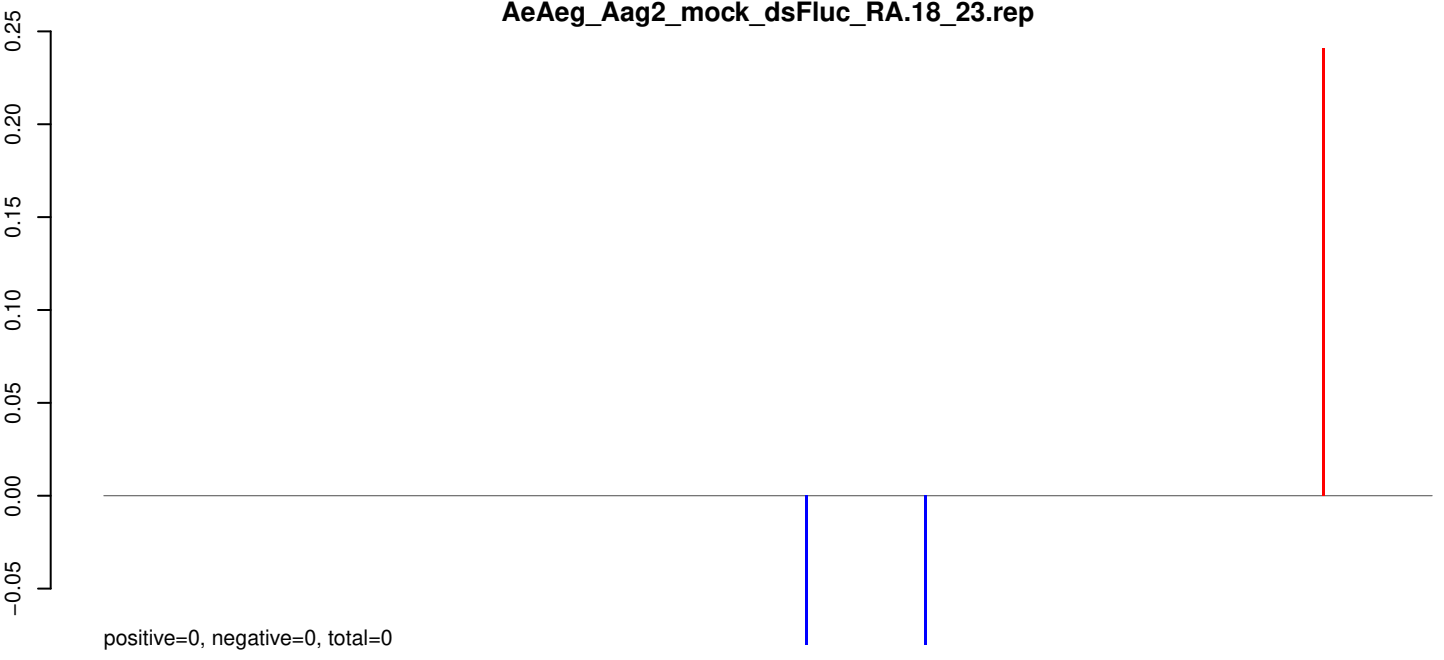
AeAeg_Aag2_mock_dsFluc_RA.rep



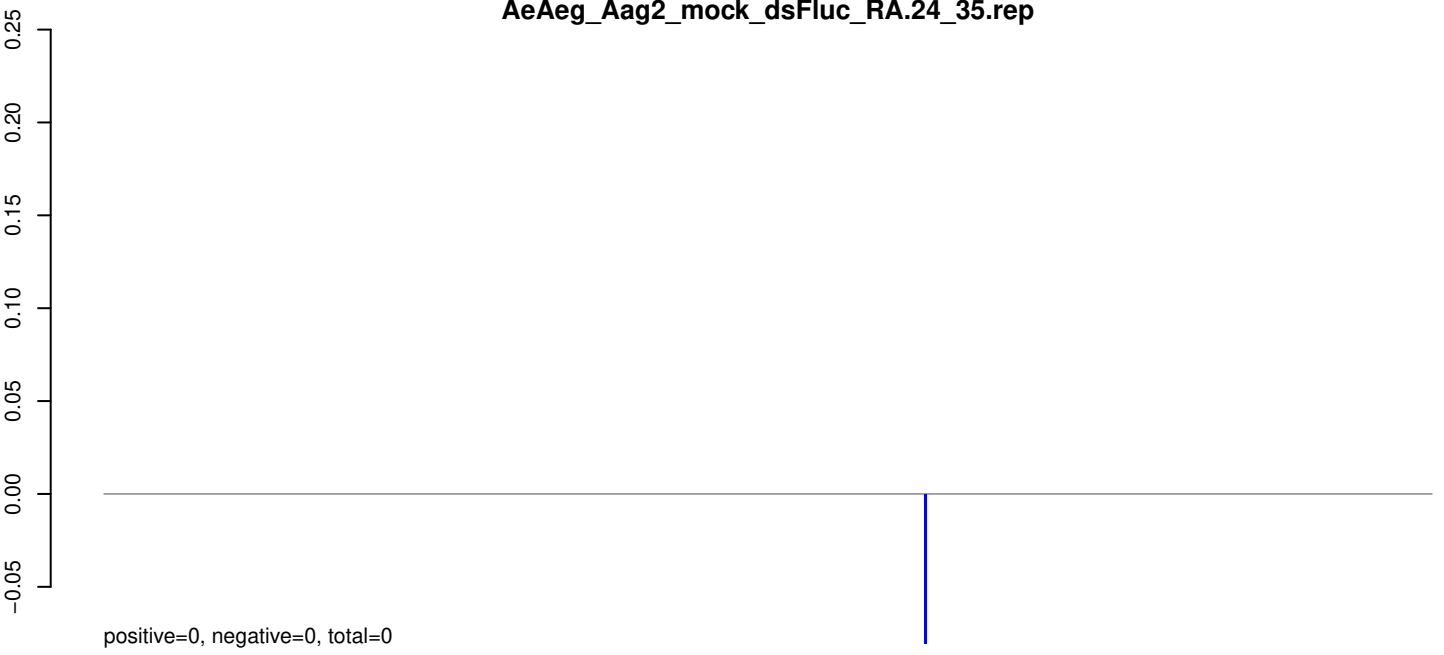
Window size=25, length=10837, virus@EU879060.1-Culex_flavivirus_strain_CxFV-Mex07:1-10837

0 2000 4000 6000 8000 10000

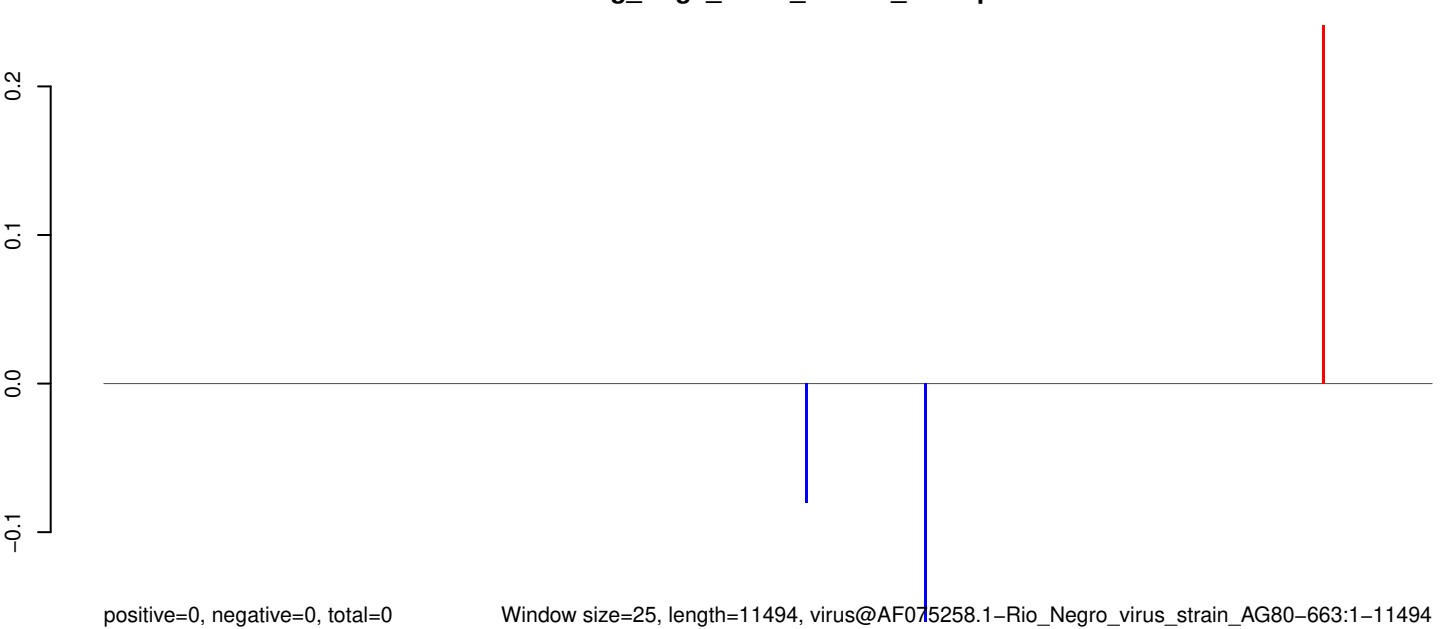
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



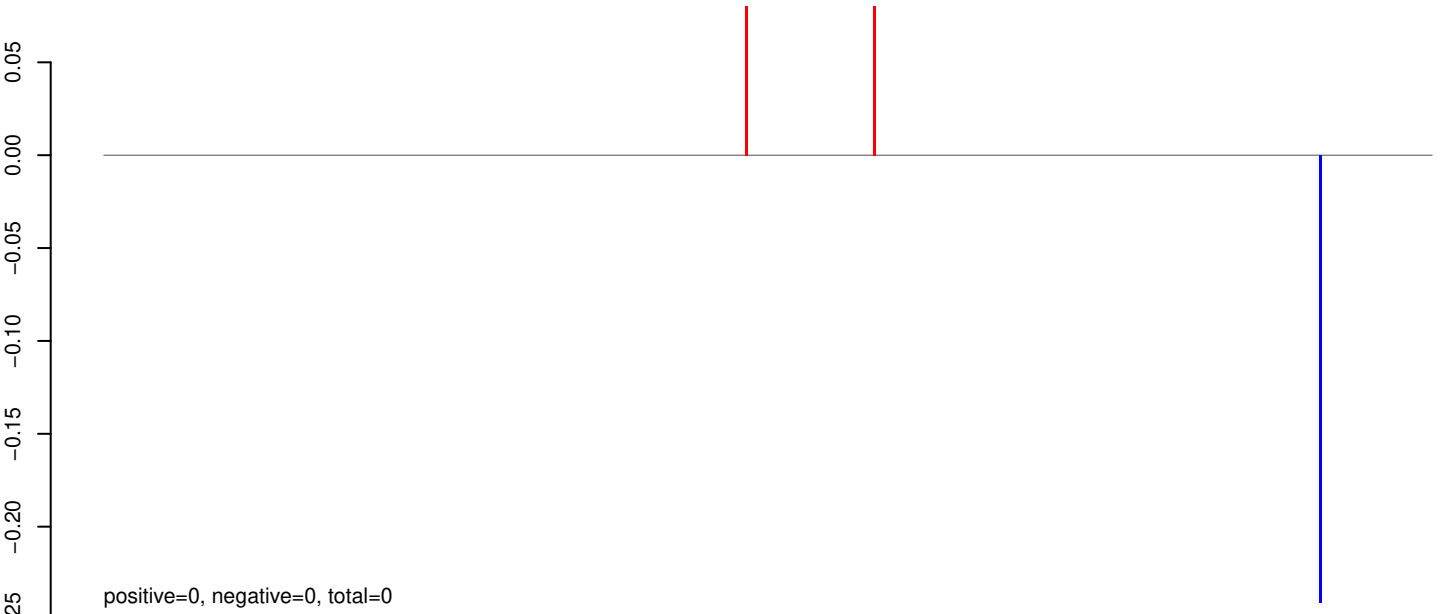
AeAeg_Aag2_mock_dsFluc_RA.rep



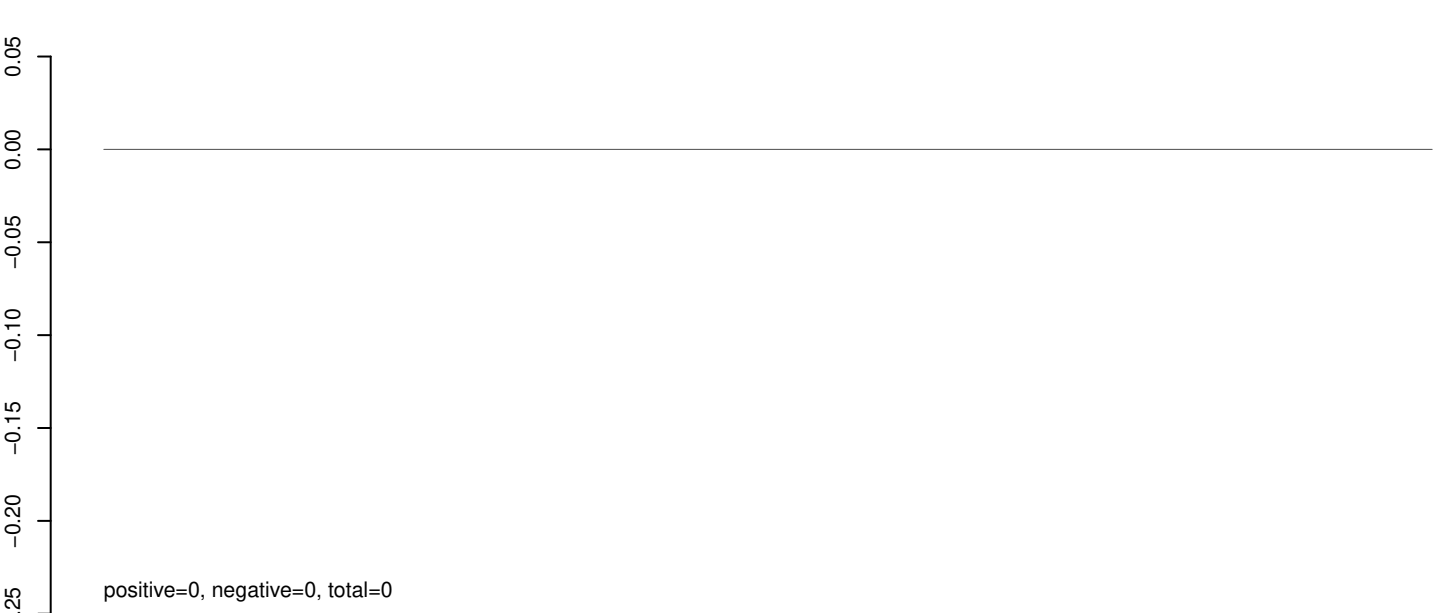
Window size=25, length=11494, virus@AF075258.1-Rio_Negro_virus_strain_AG80-663:1-11494

0 2000 4000 6000 8000 10000

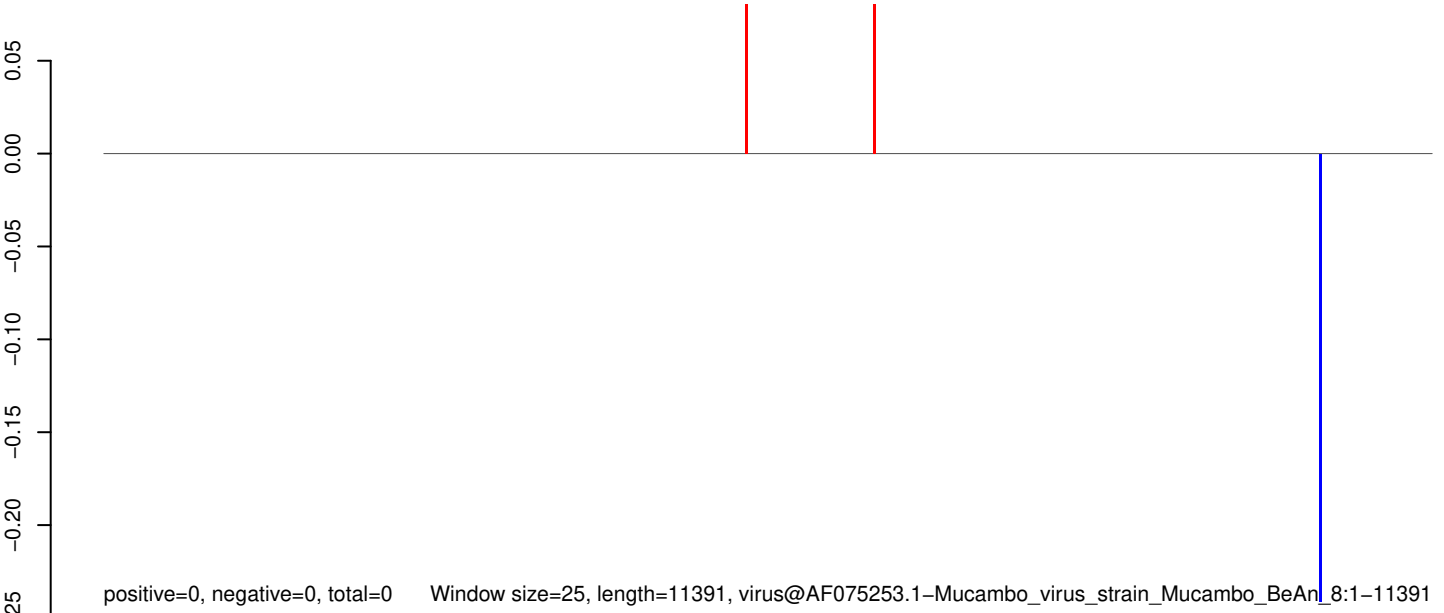
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



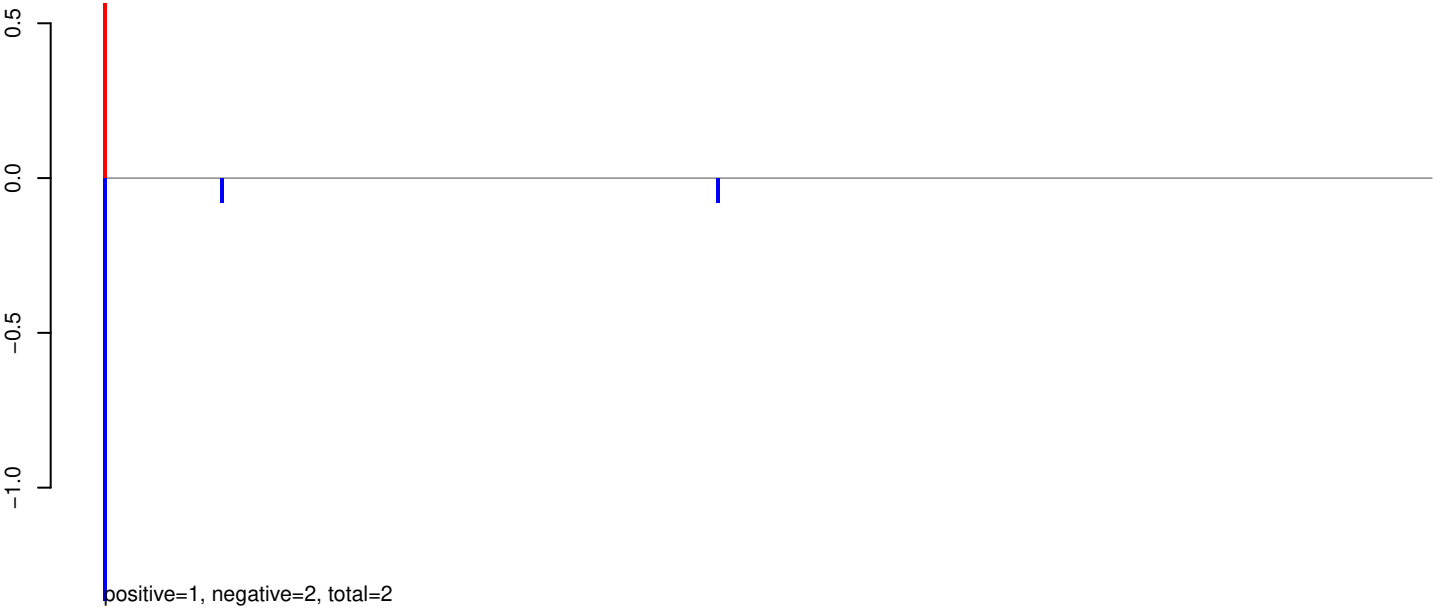
AeAeg_Aag2_mock_dsFluc_RA.rep



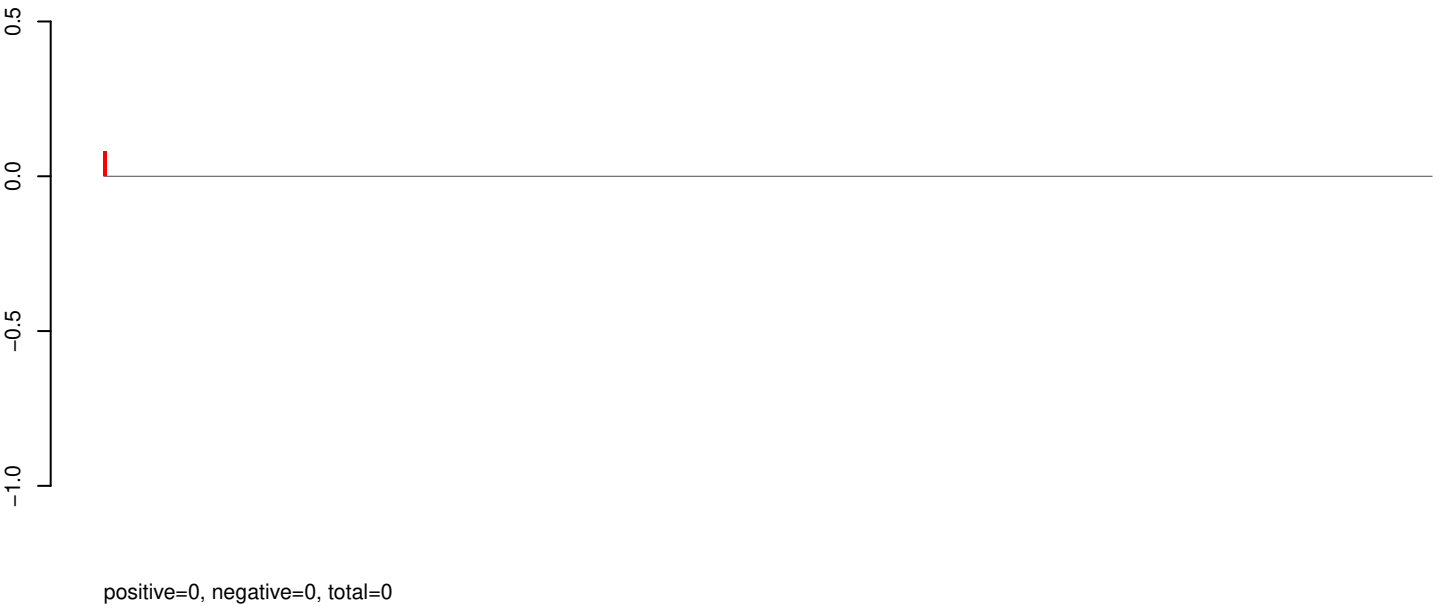
Window size=25, length=11391, virus@AF075253.1-Mucambo_virus_strain_Mucambo_BeAn_8:1-11391

0 2000 4000 6000 8000 10000

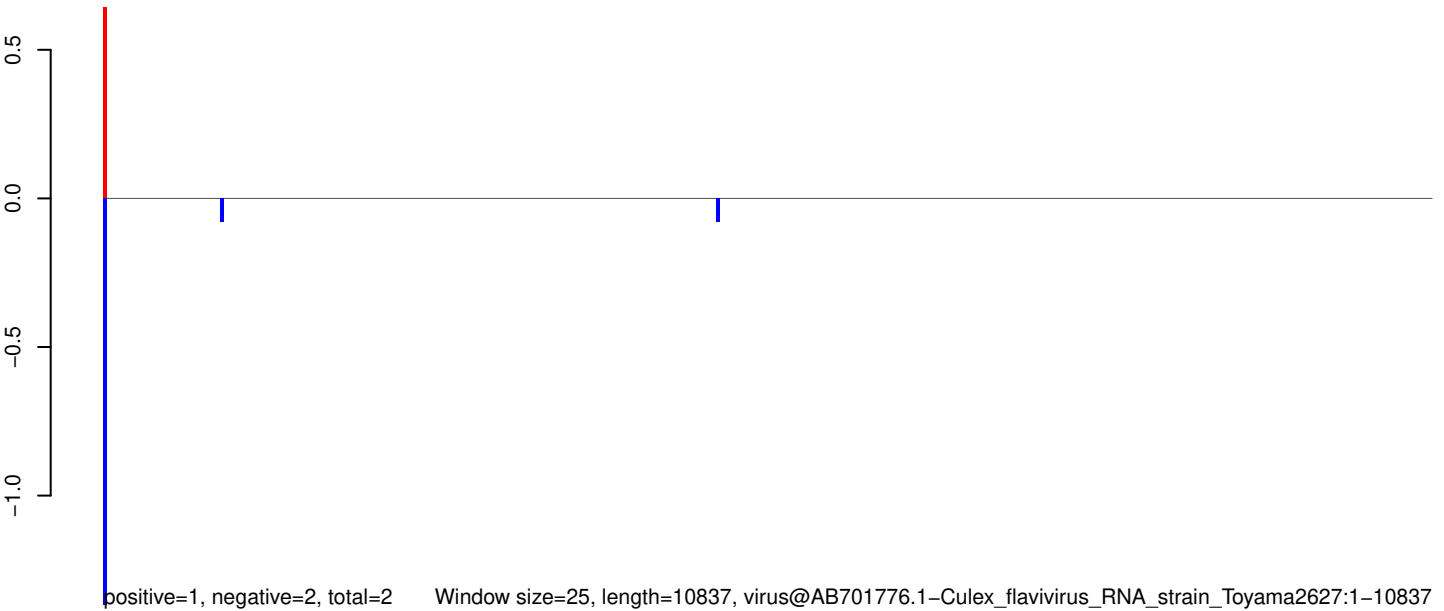
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep



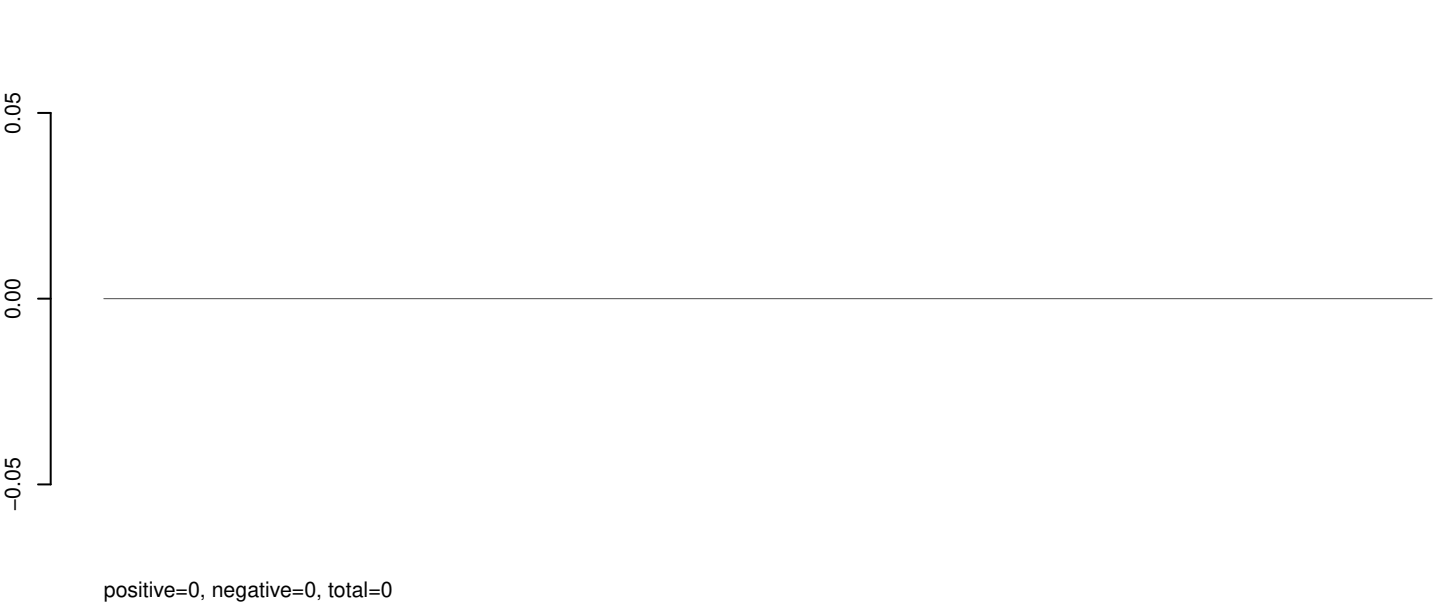
Window size=25, length=10837, virus@AB701776.1-Culex_flavivirus_RNA_strain_Toyama2627:1-10837

0 2000 4000 6000 8000 10000

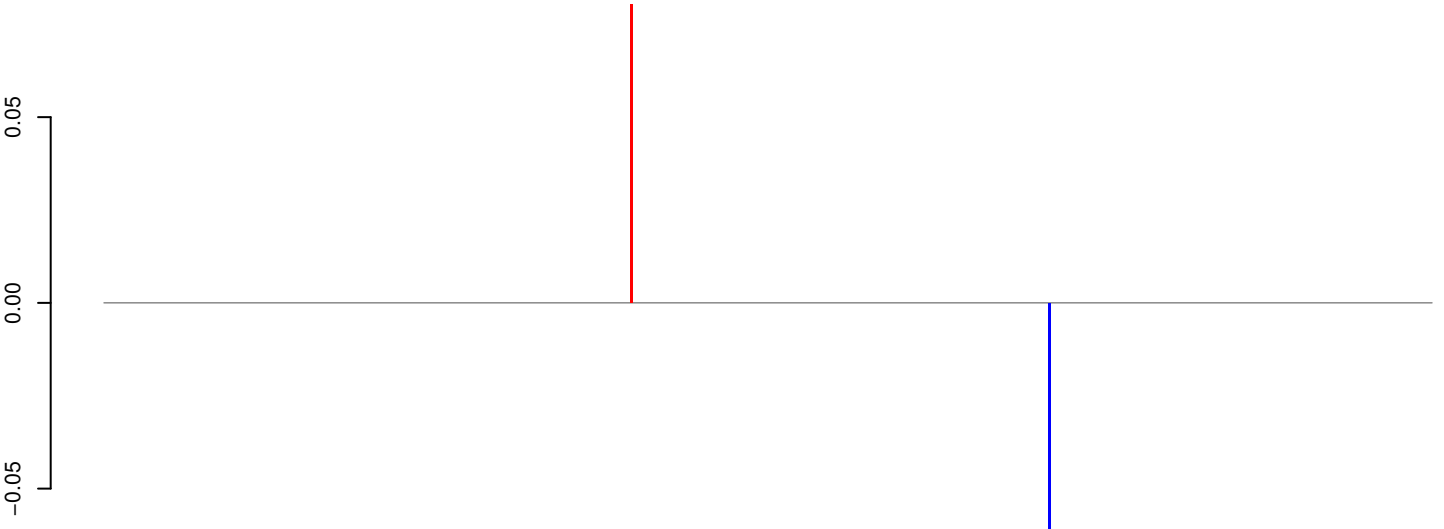
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



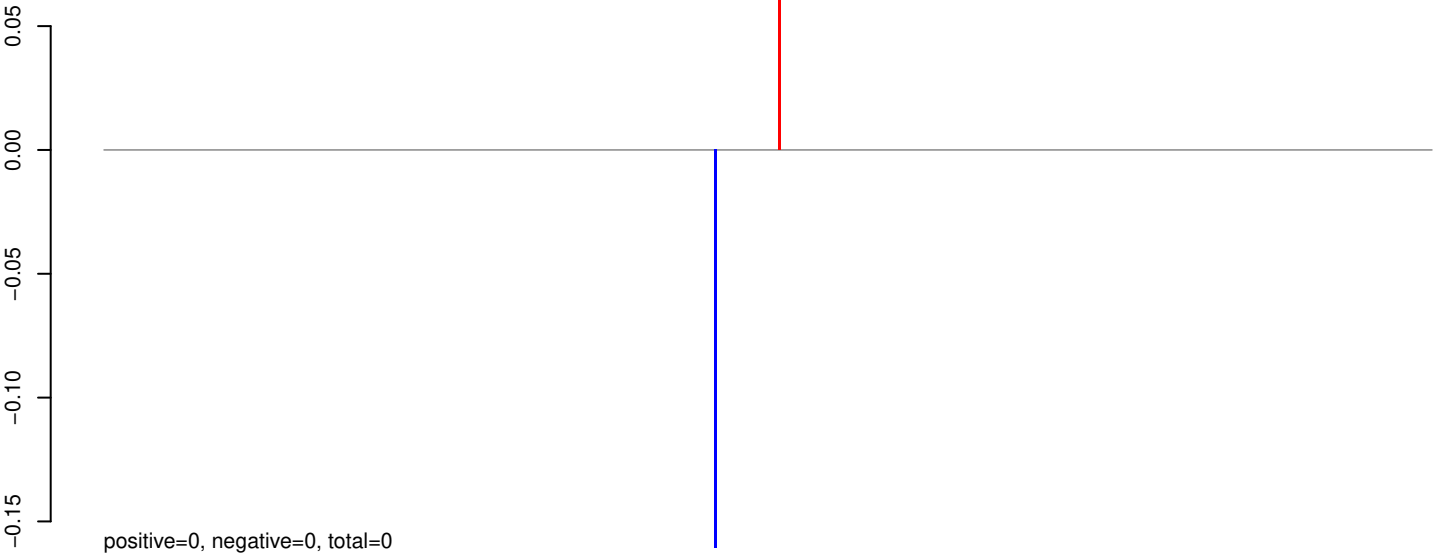
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



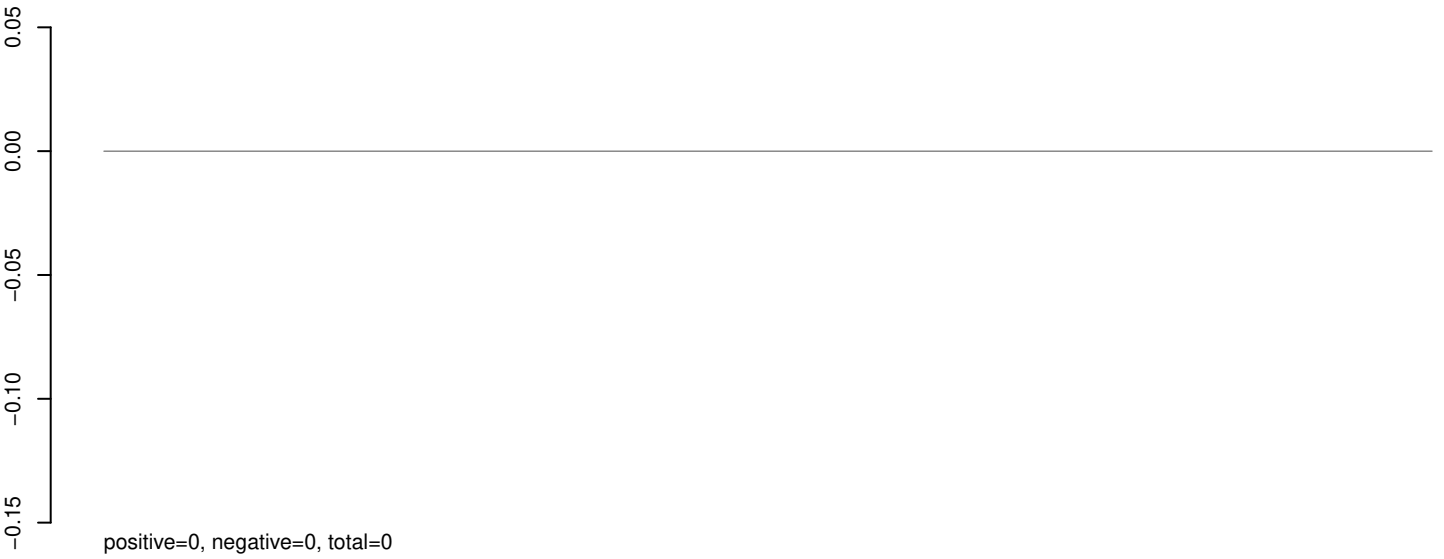
AeAeg_Aag2_mock_dsFluc_RA.rep



AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

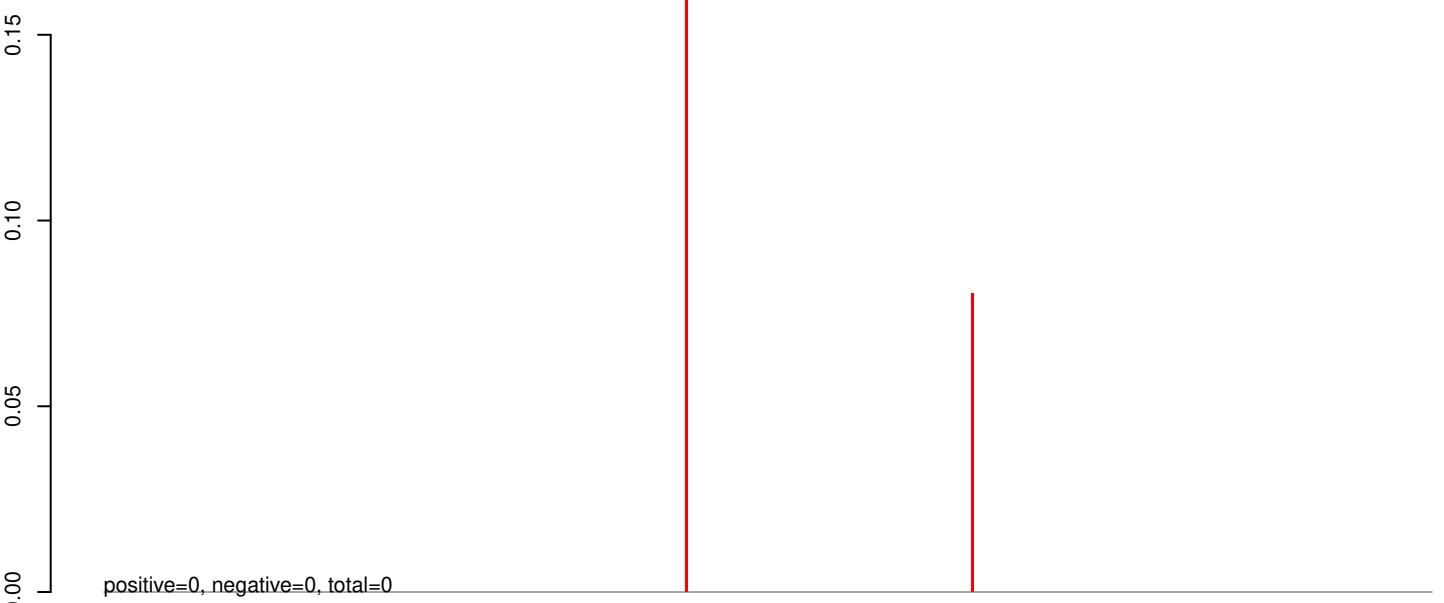


AeAeg_Aag2_mock_dsFluc_RA.rep

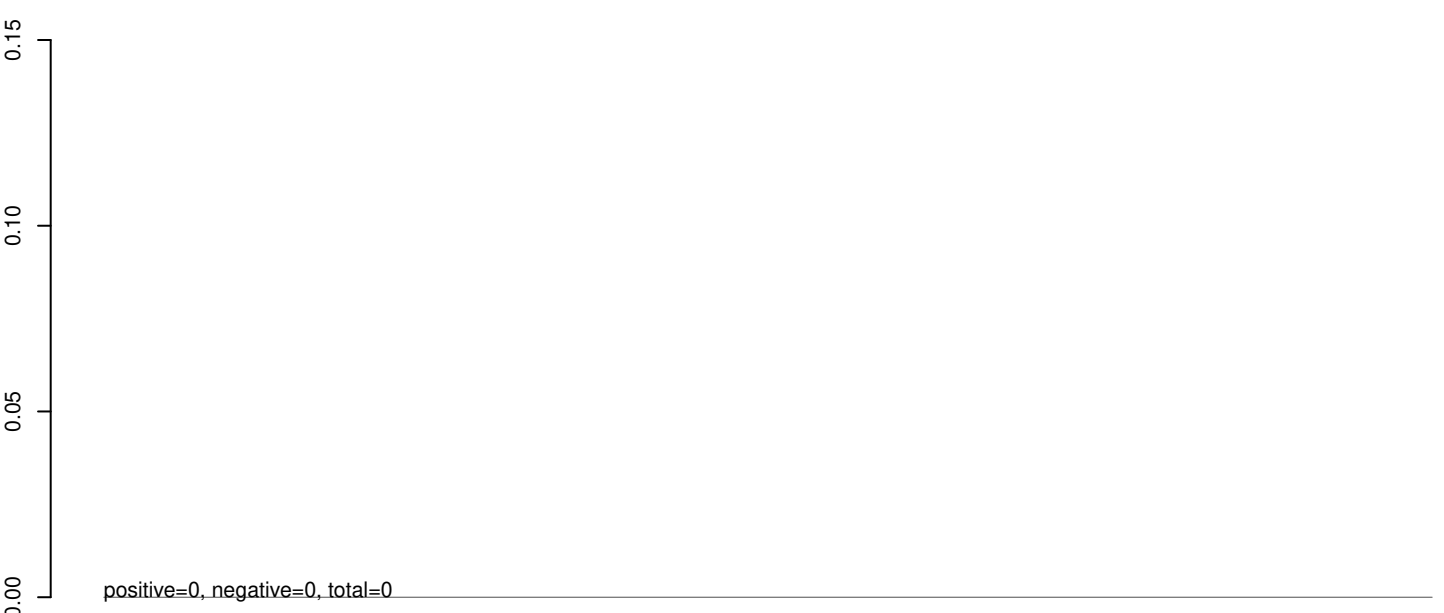


0 2000 4000 6000 8000 10000
positive=0, indel size=25, length=11470, virus@NC_035128.1_Aedes_camptorhynchus_negev-like_virus_strain_mos191CP47896:1-11470

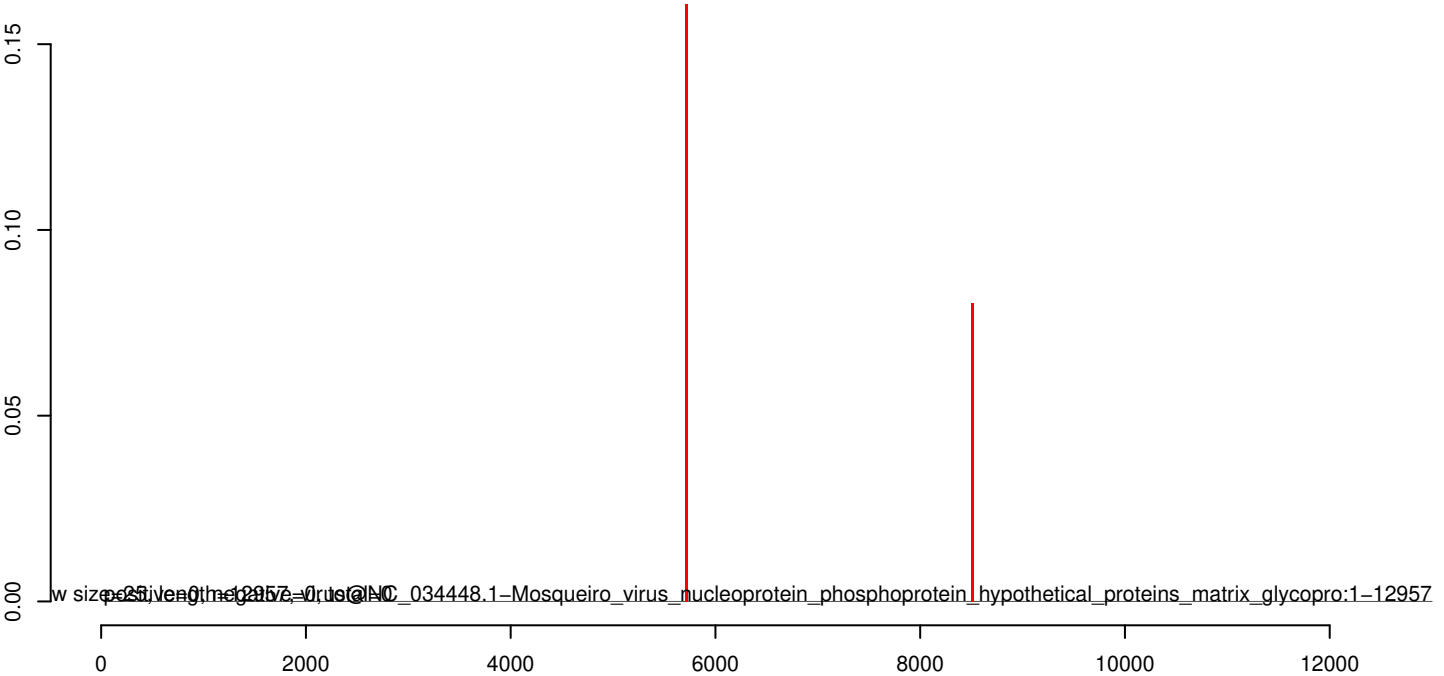
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



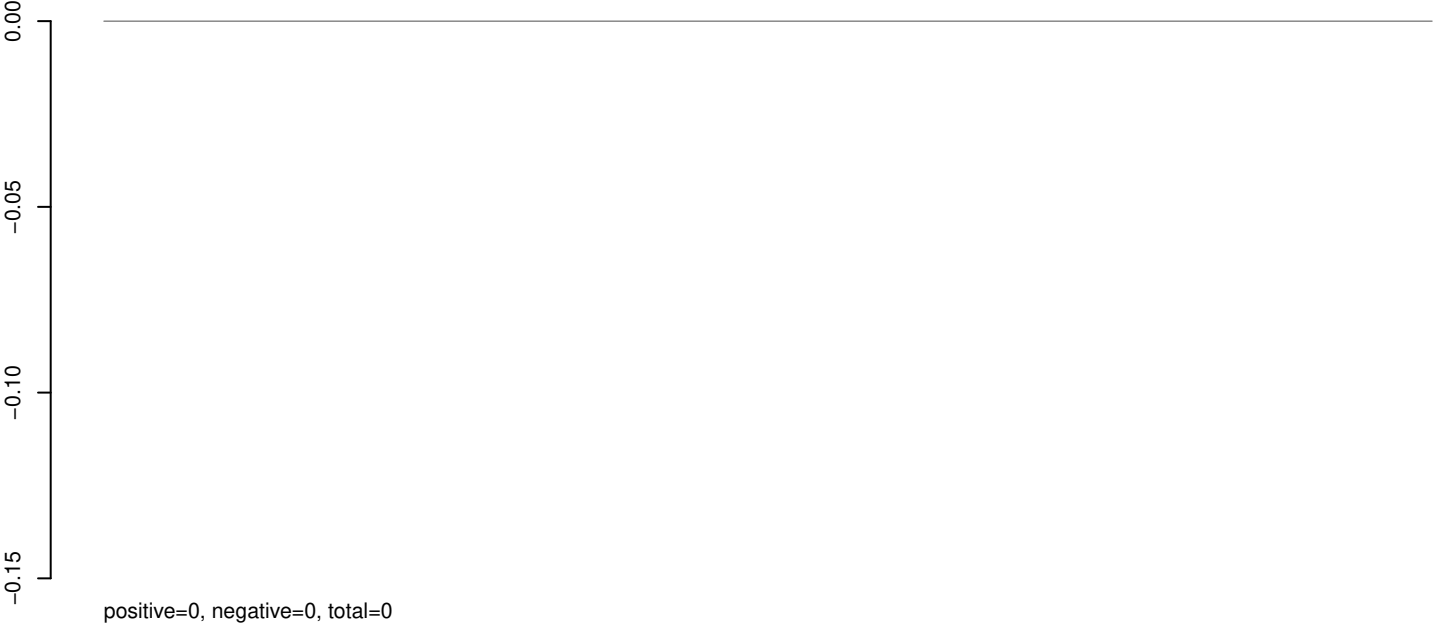
AeAeg_Aag2_mock_dsFluc_RA.rep



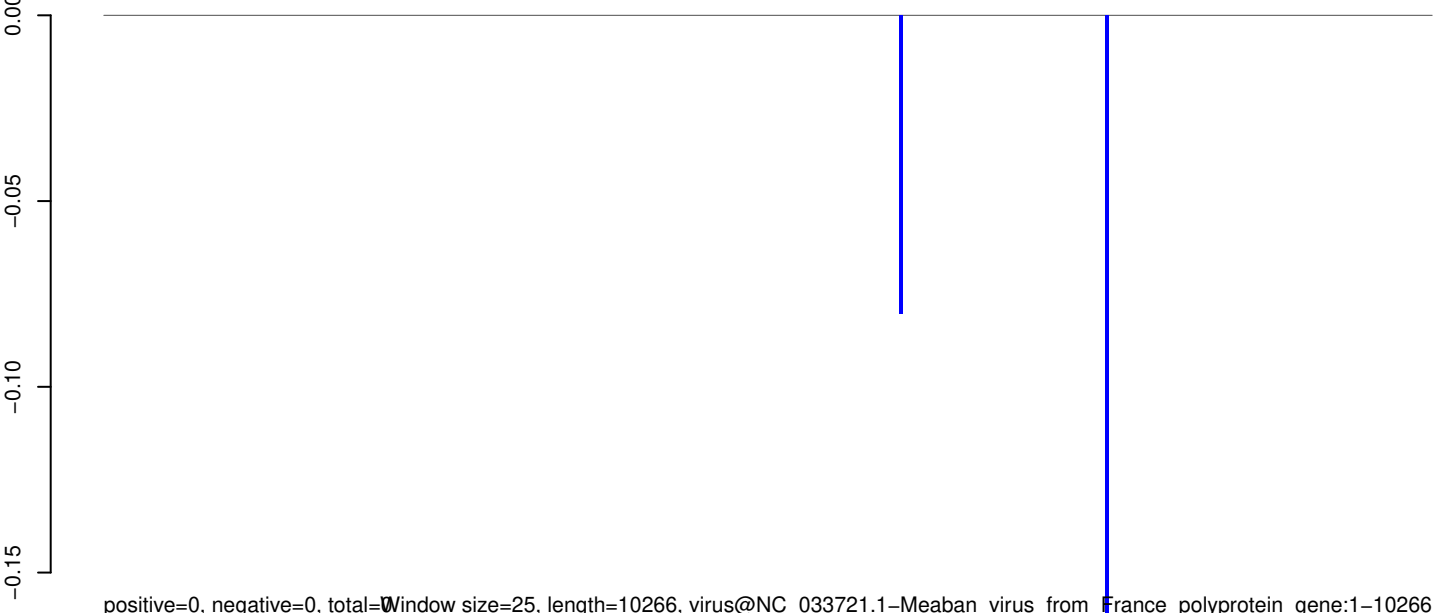
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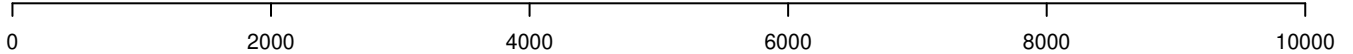
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



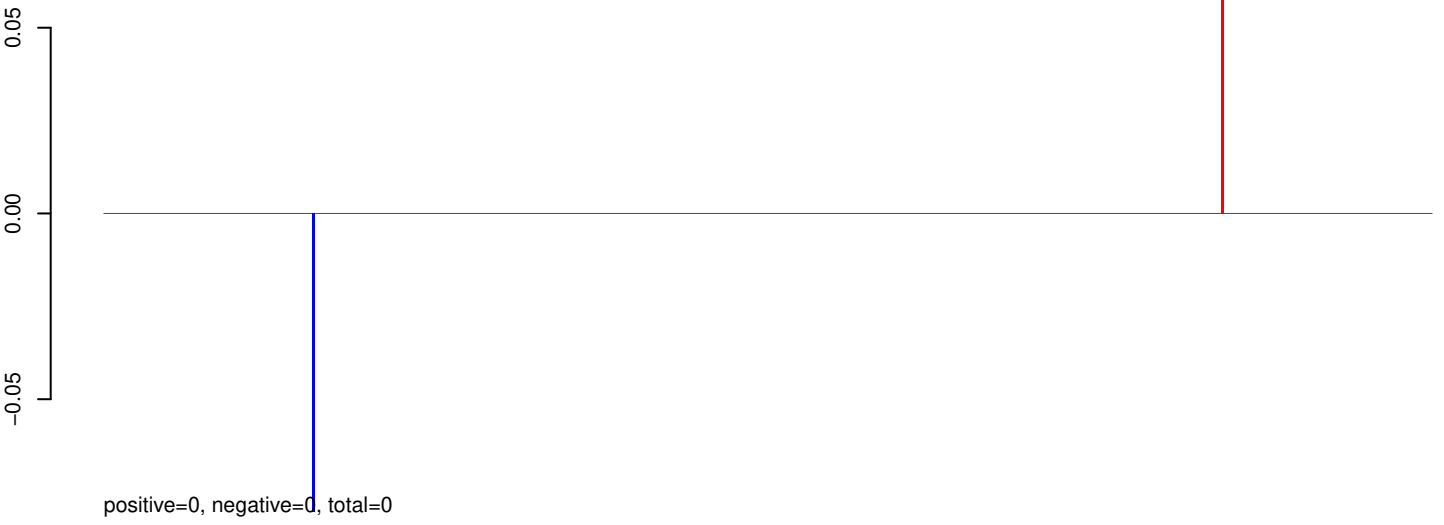
AeAeg_Aag2_mock_dsFluc_RA.rep



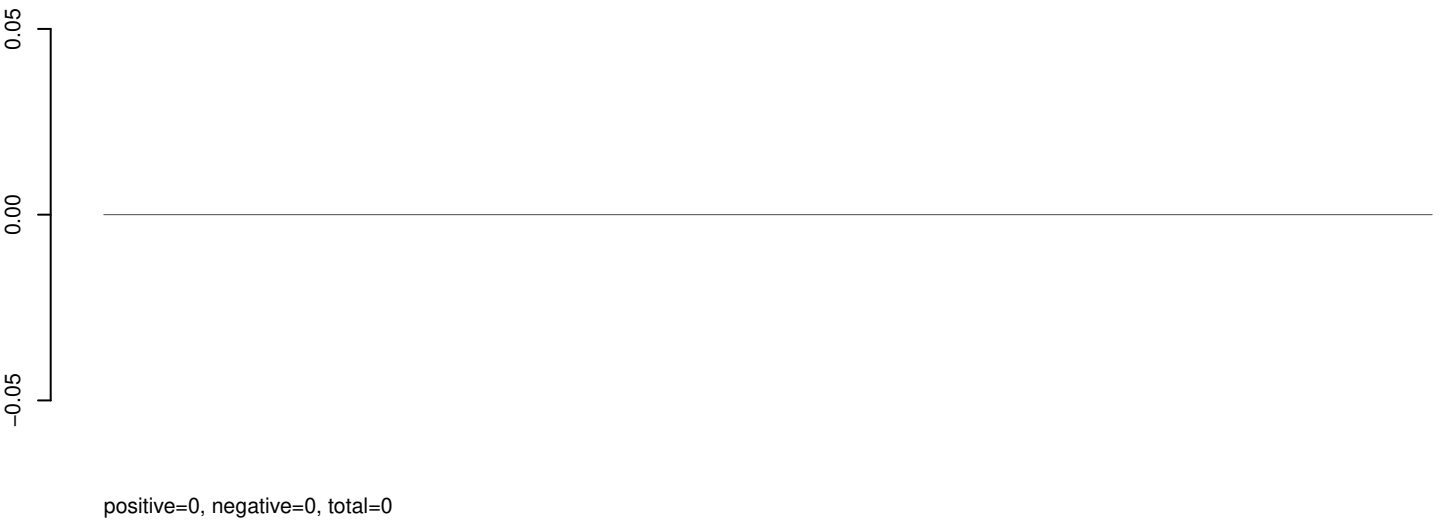
Window size=25, length=10266, virus@NC_033721.1-Meaban_virus_from_France_polyprotein_gene:1-10266



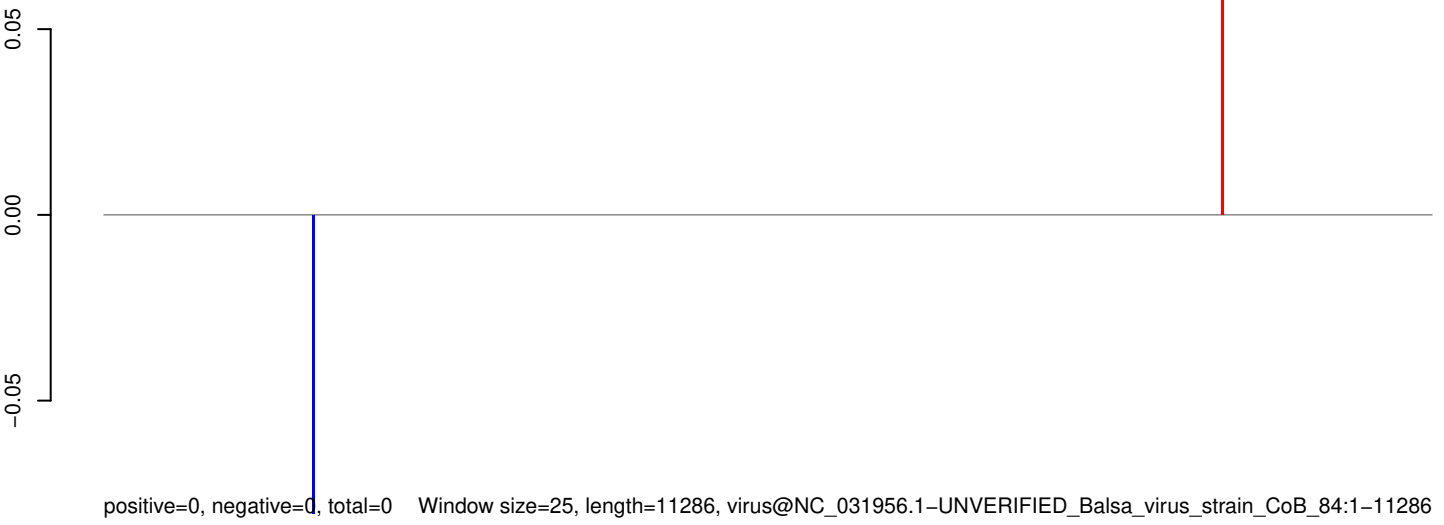
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



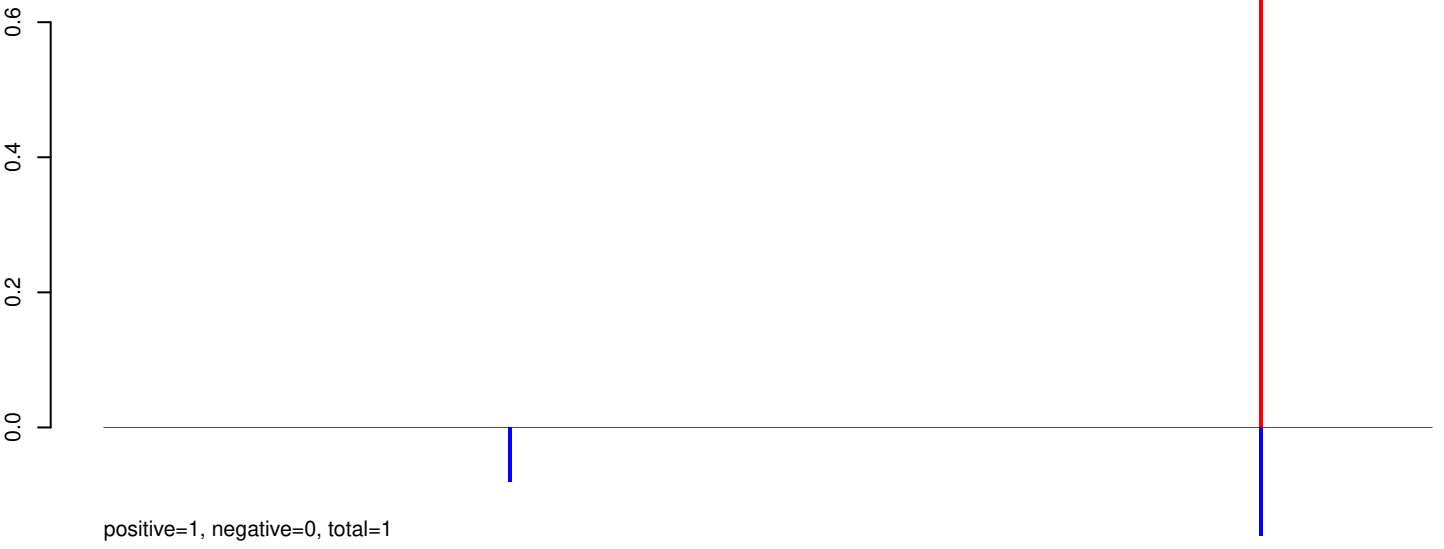
AeAeg_Aag2_mock_dsFluc_RA.rep



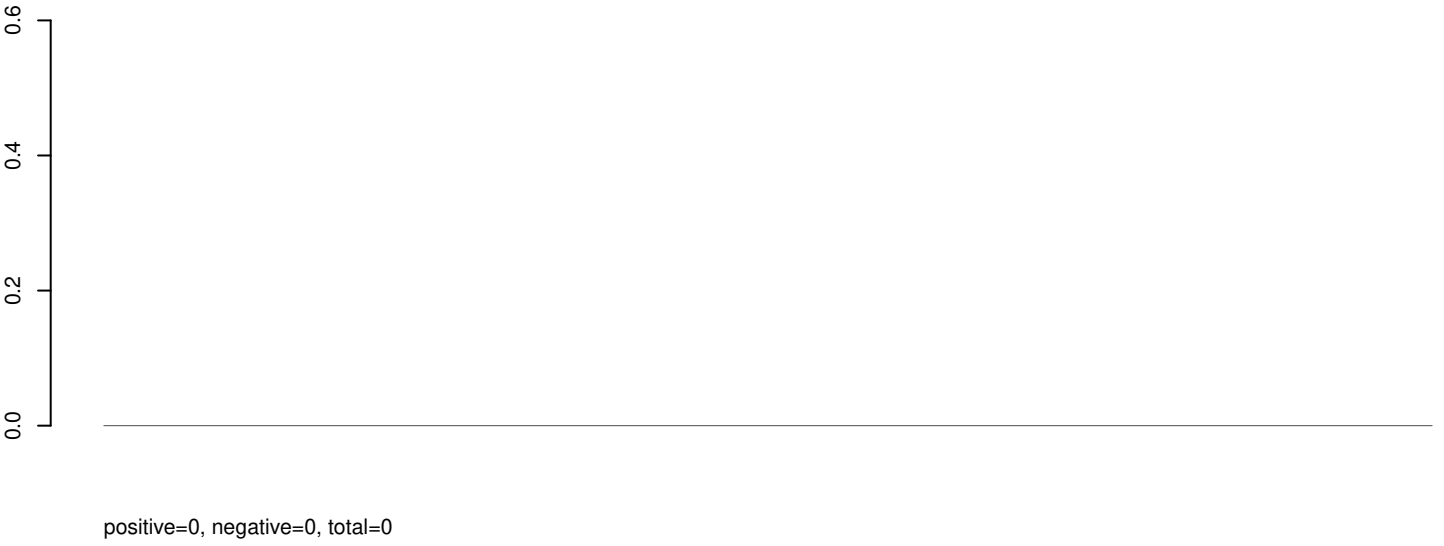
Window size=25, length=11286, virus@NC_031956.1-UNVERIFIED_Balsa_virus_strain_CoB_84:1-11286

0 2000 4000 6000 8000 10000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

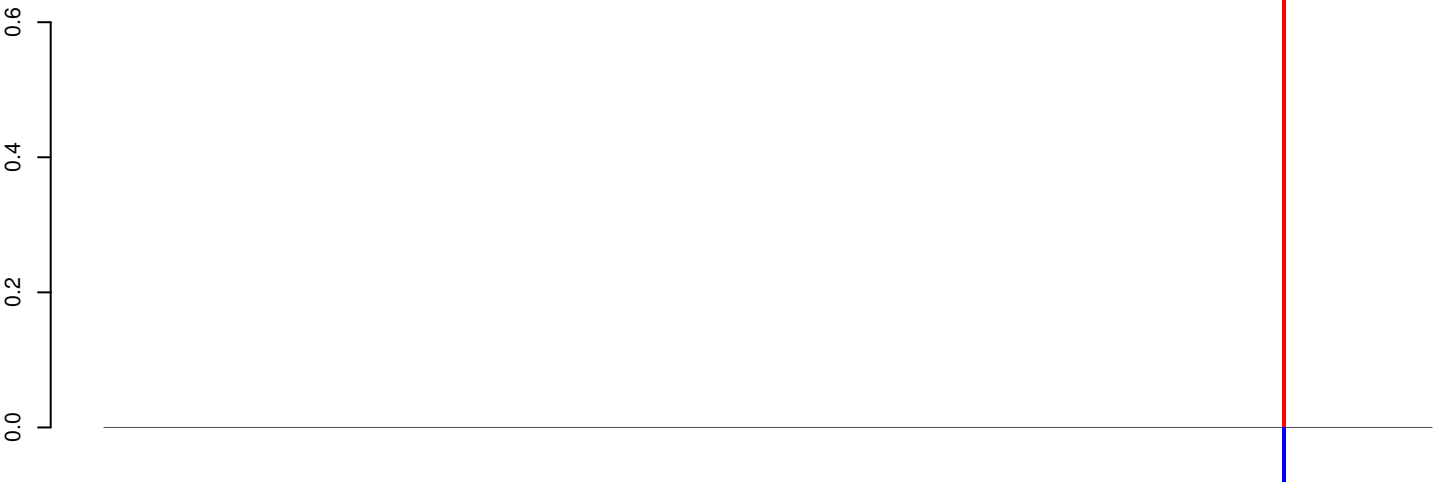


AeAeg_Aag2_mock_dsFluc_RA.rep



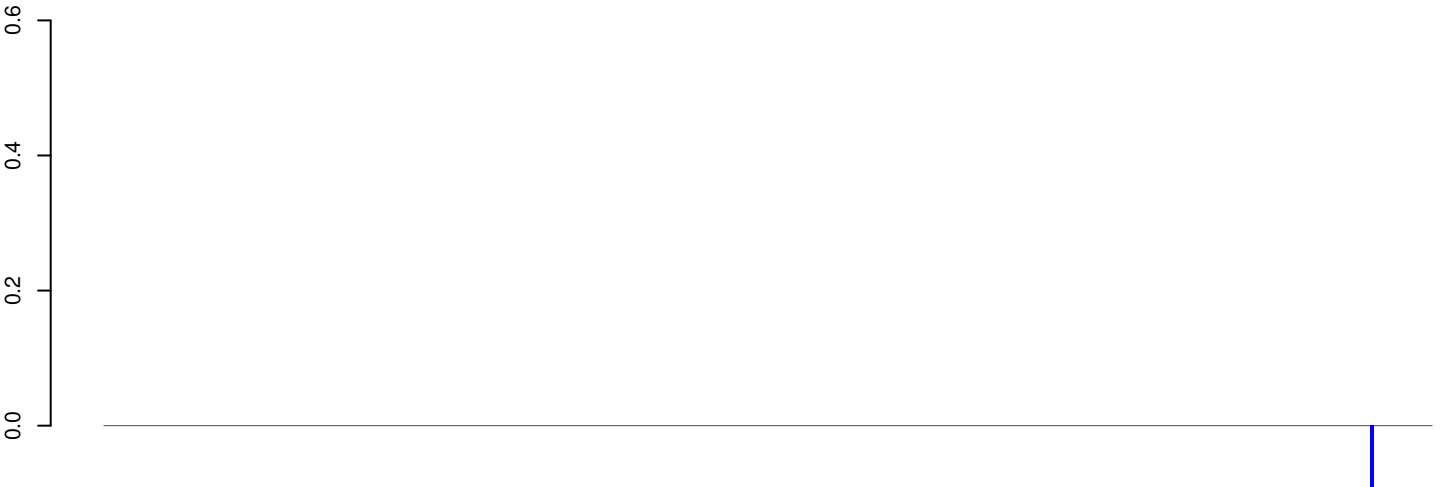
0 2000 4000 6000 8000 10000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



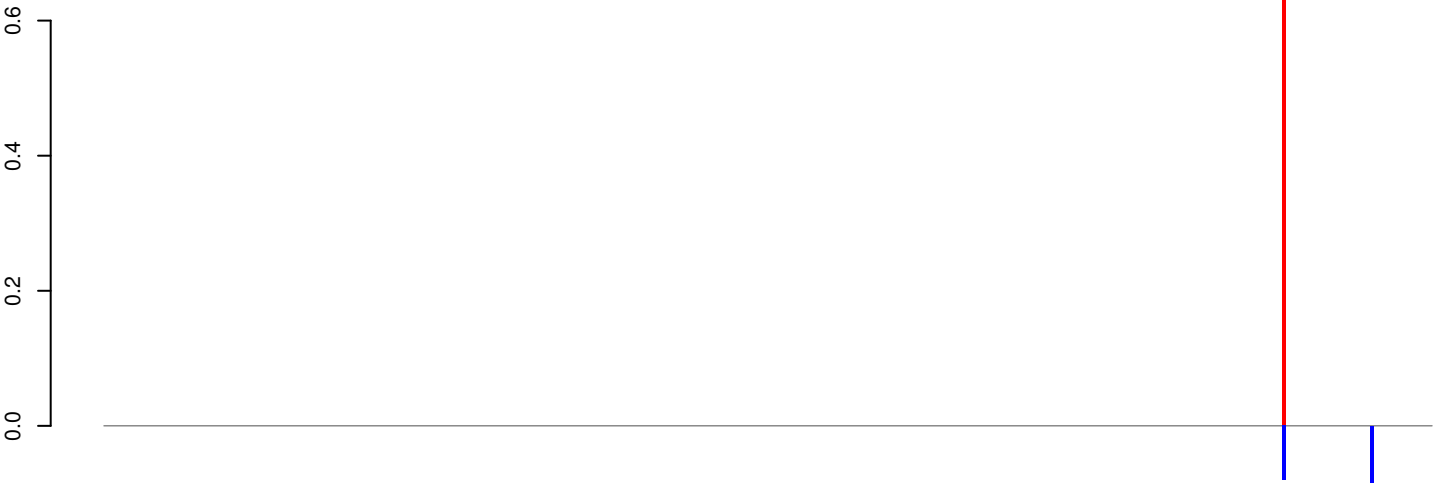
positive=1, negative=0, total=1

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



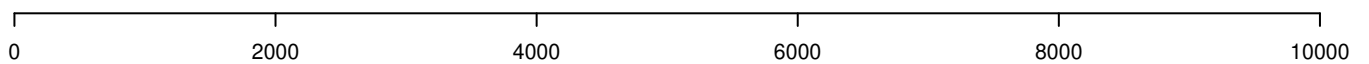
positive=0, negative=0, total=0

AeAeg_Aag2_mock_dsFluc_RA.rep

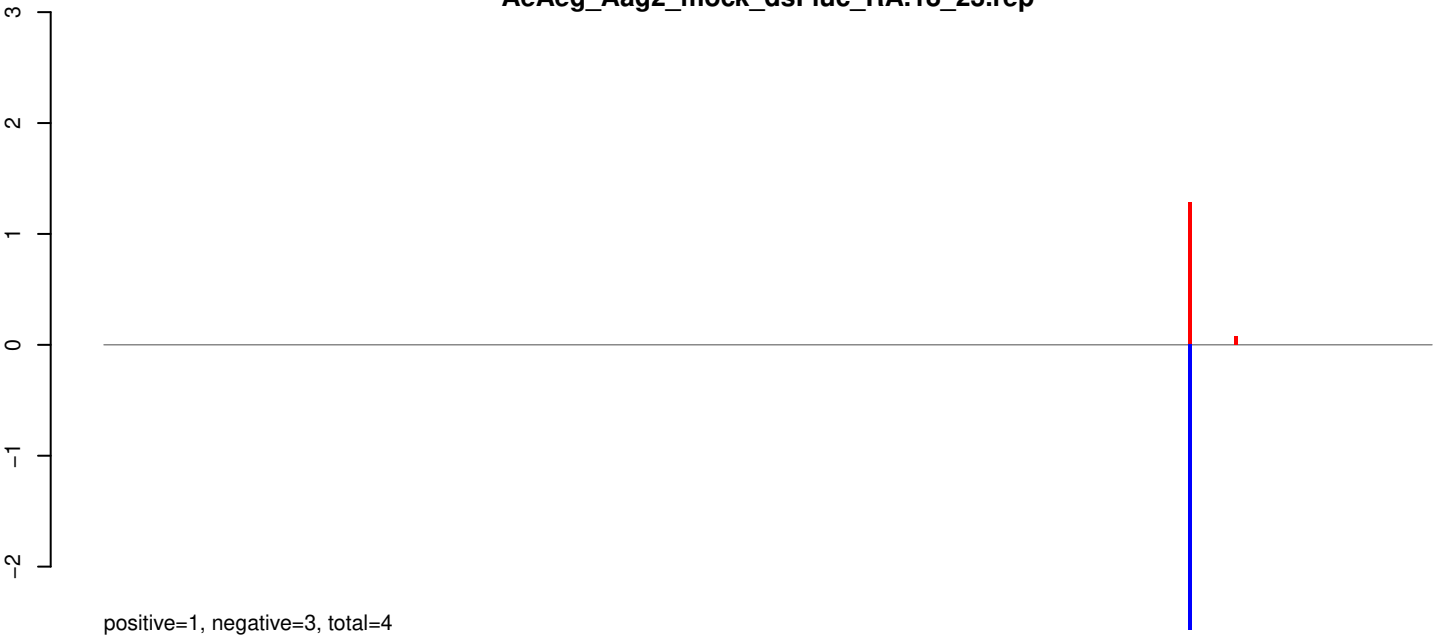


positive=1, negative=0, total=1

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

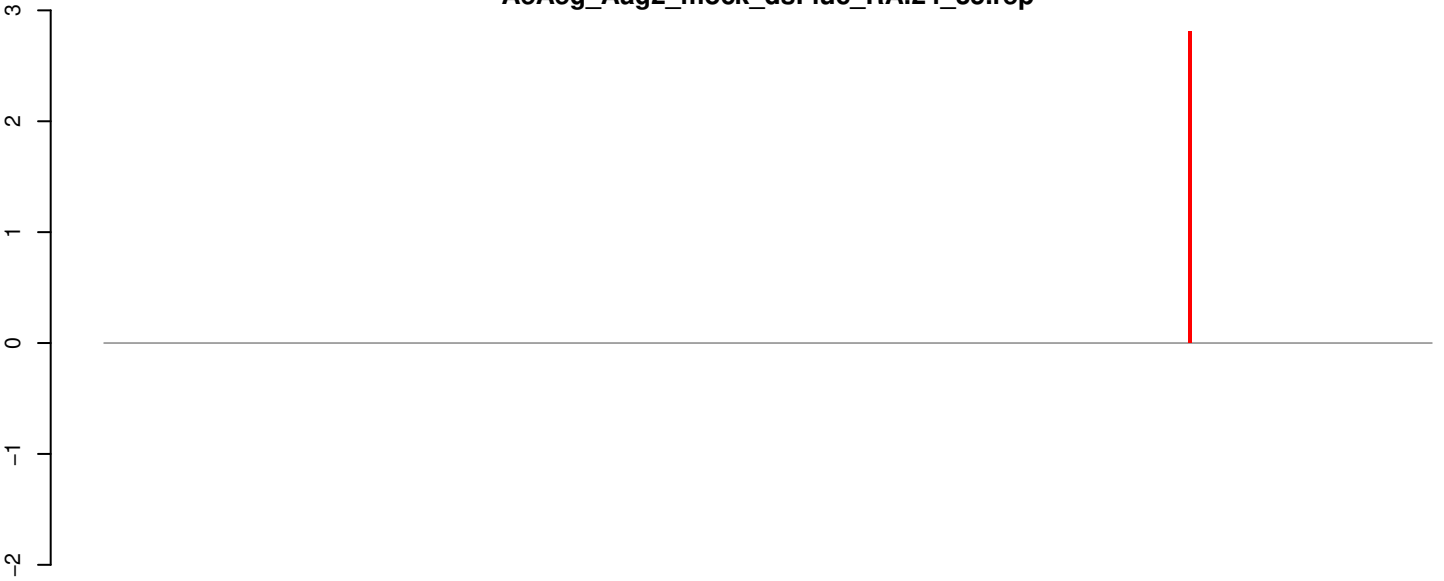


AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



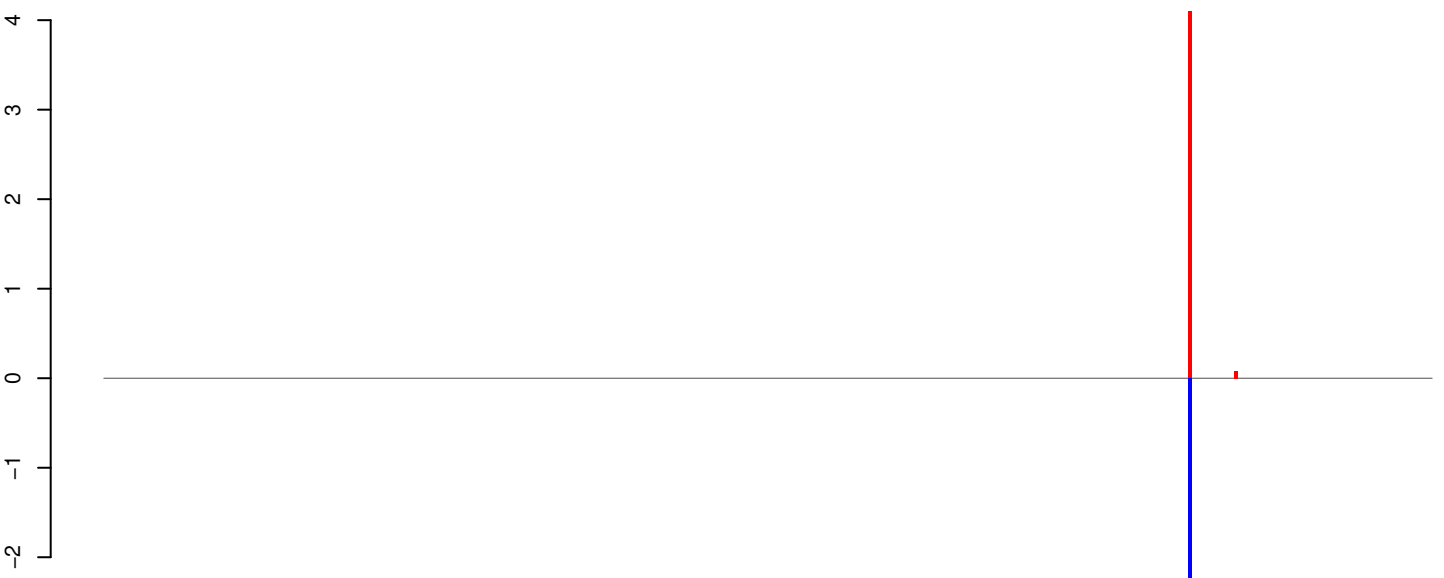
positive=1, negative=3, total=4

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



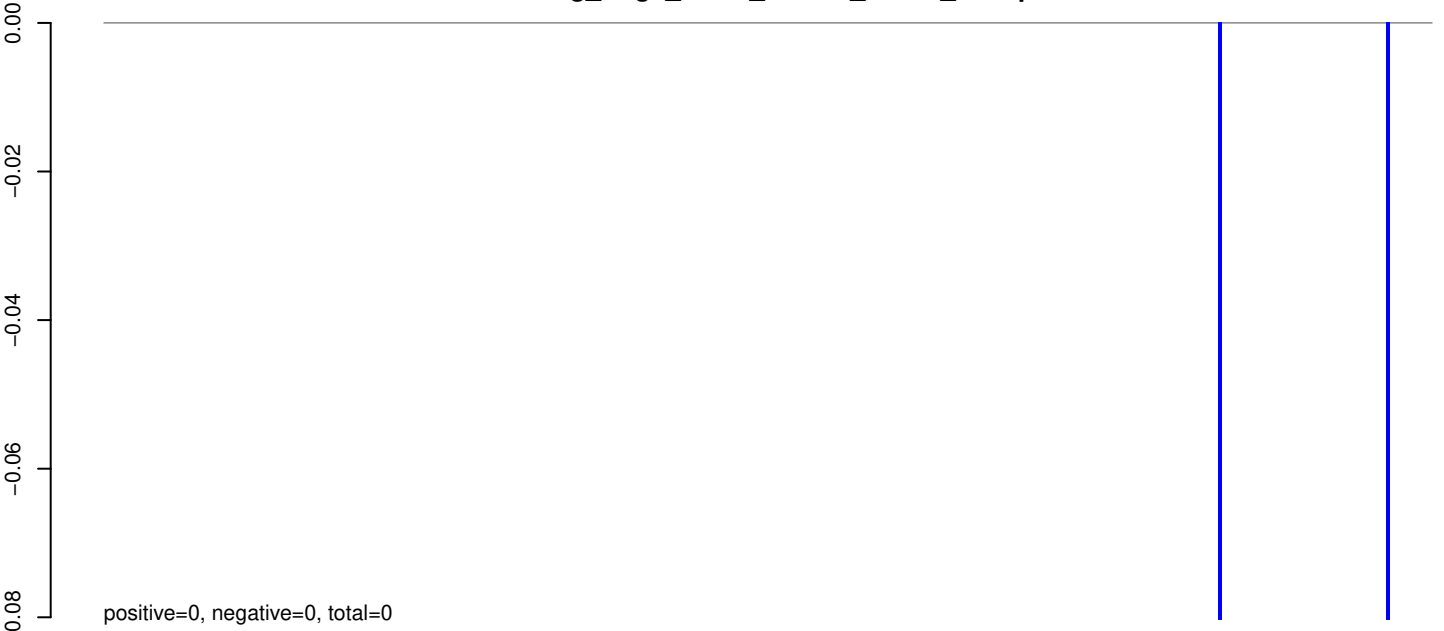
positive=3, negative=0, total=3

AeAeg_Aag2_mock_dsFluc_RA.rep

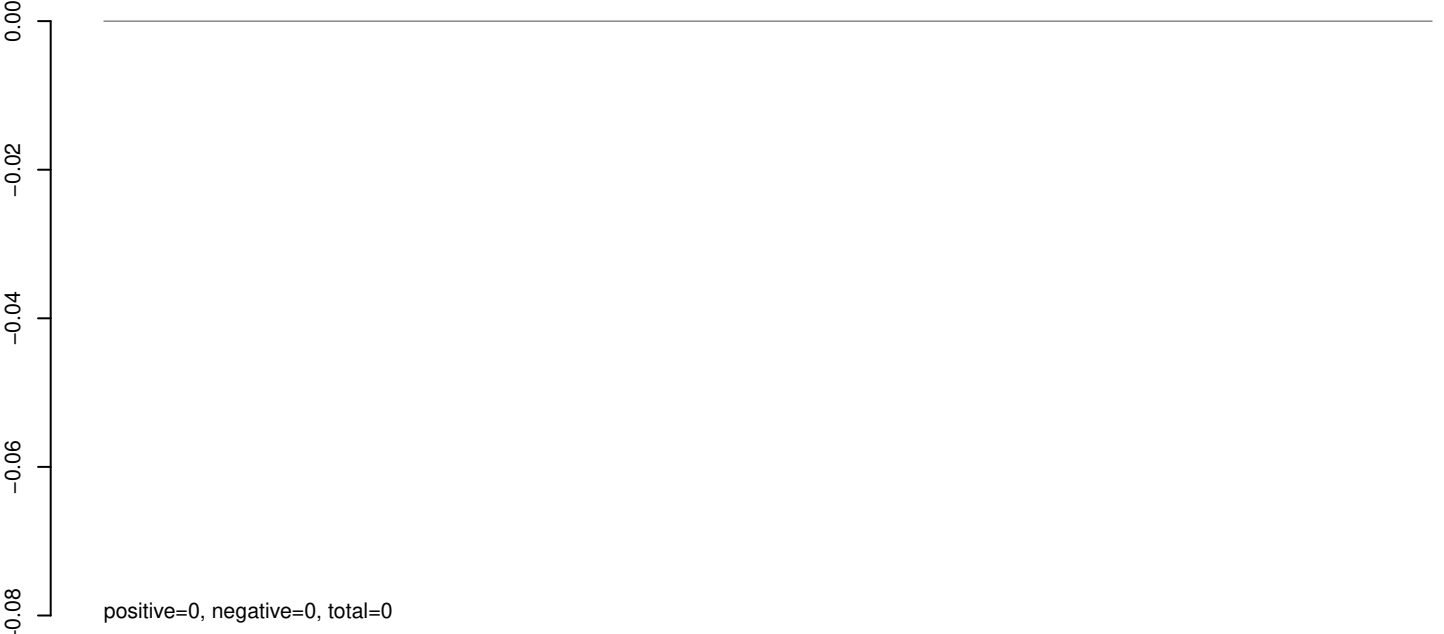


positive=4, negative=3, total=7 Window size=25, length=10893, virus@NC_027817.1-Parramatta_River_virus_isolate_92-B115745:1-10893

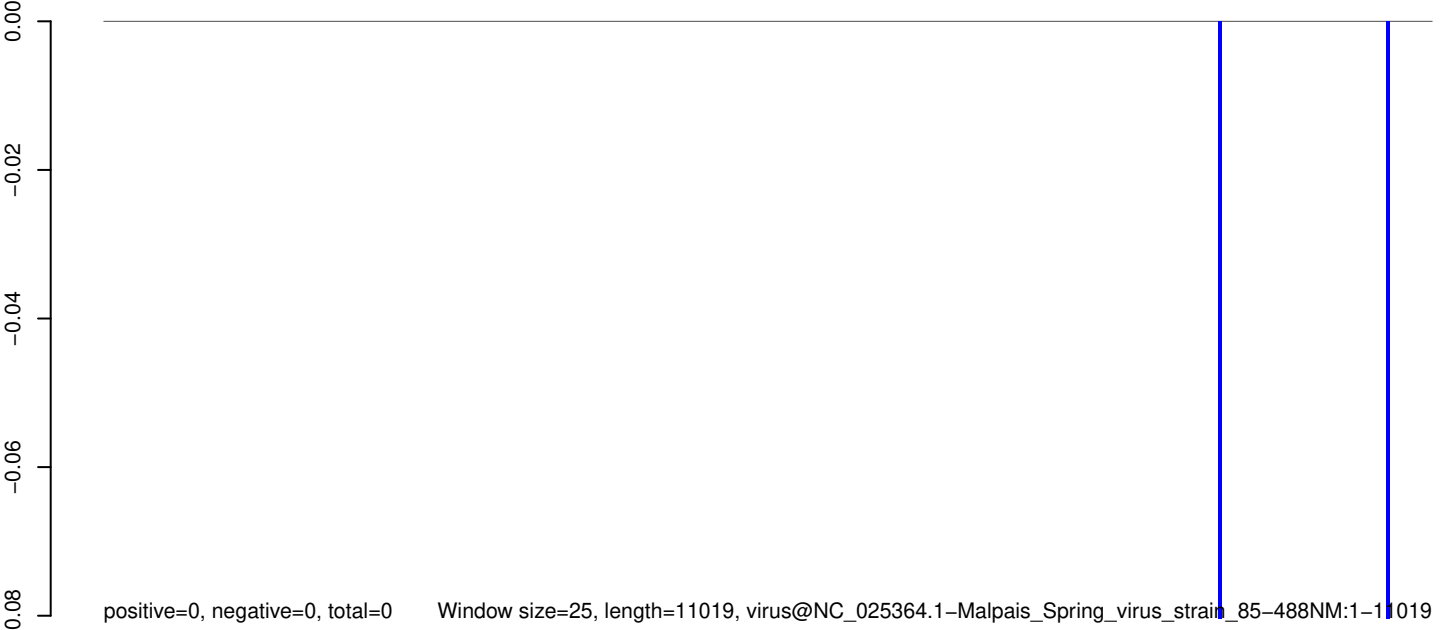
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



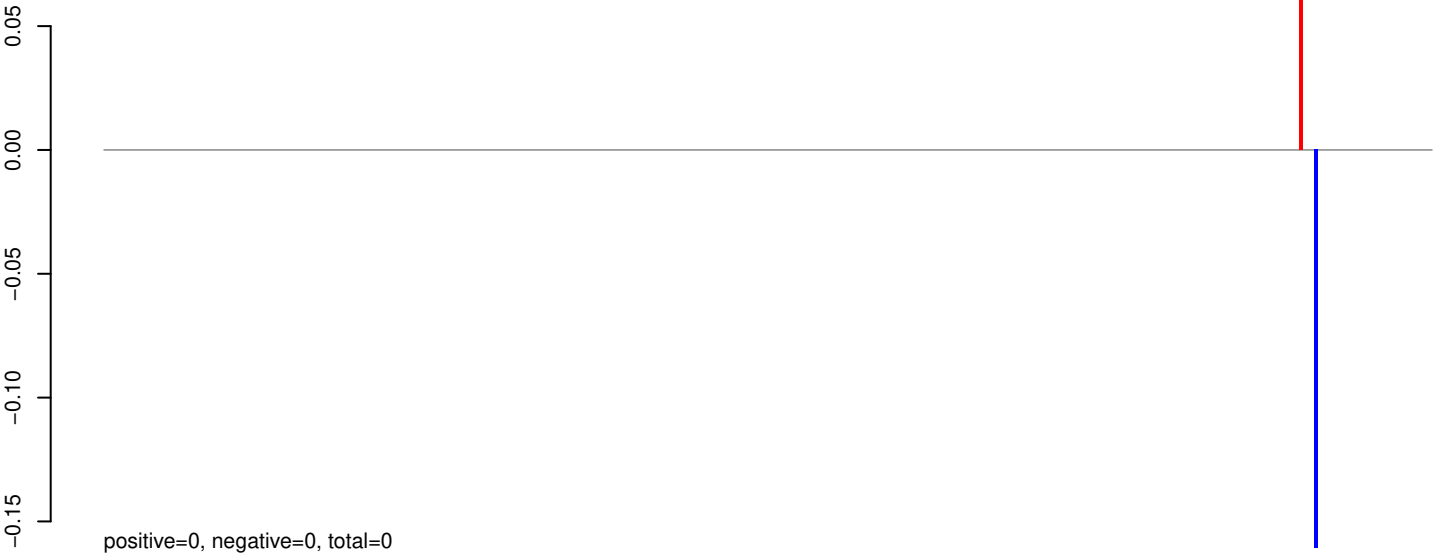
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



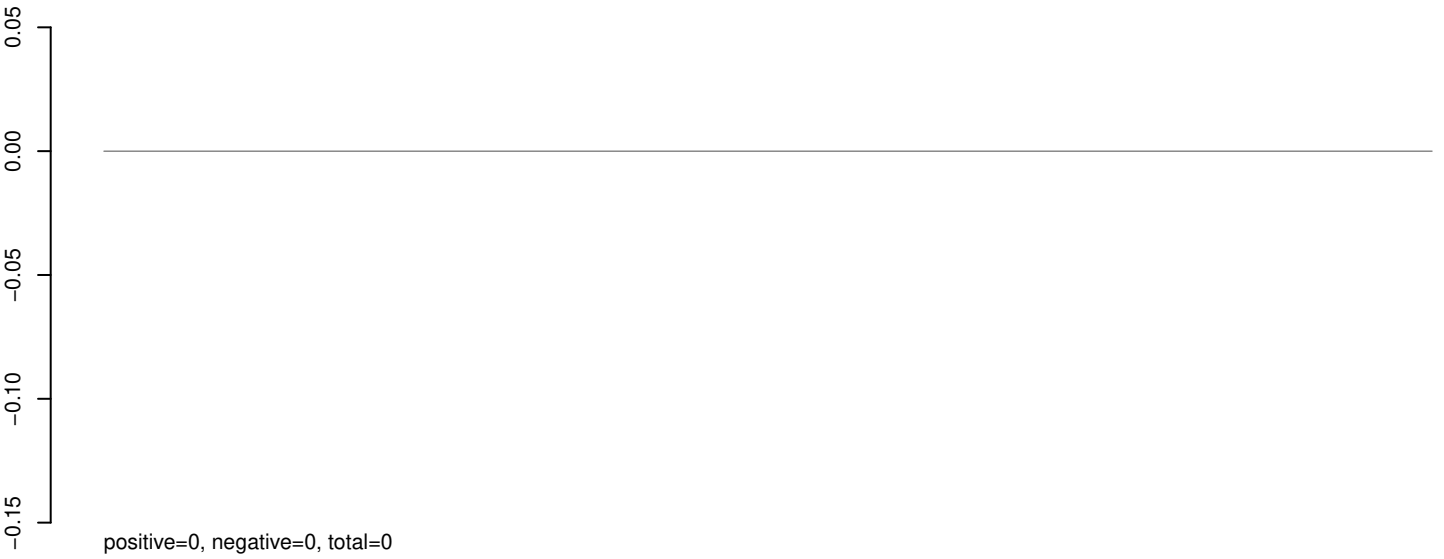
AeAeg_Aag2_mock_dsFluc_RA.rep



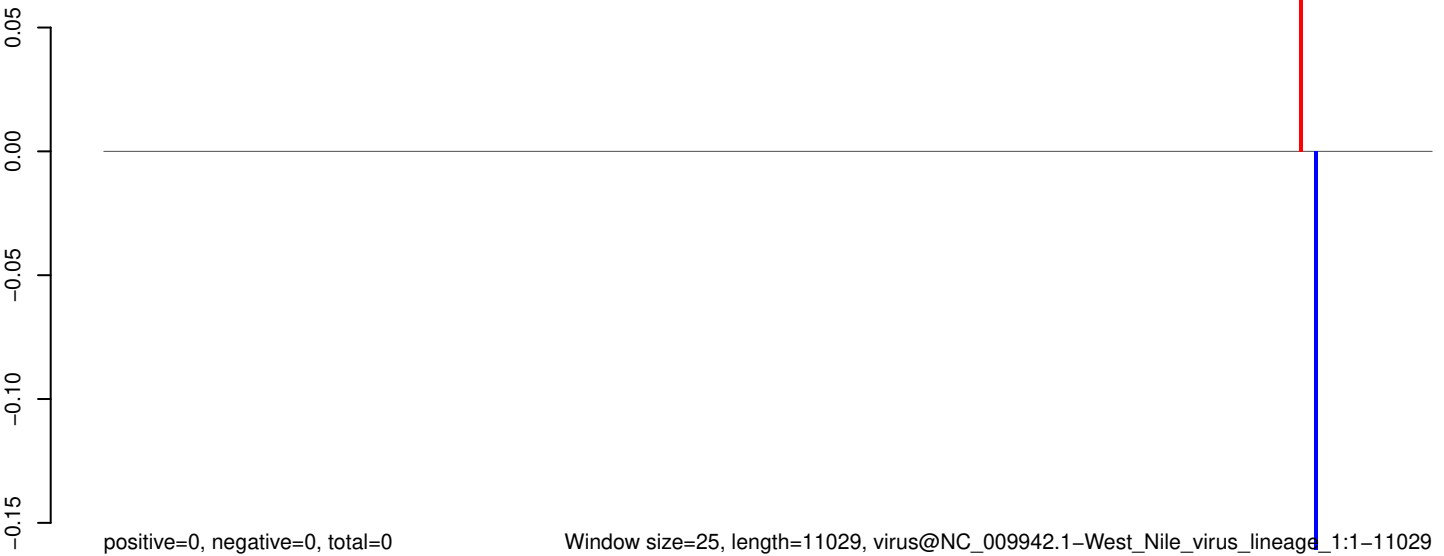
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep

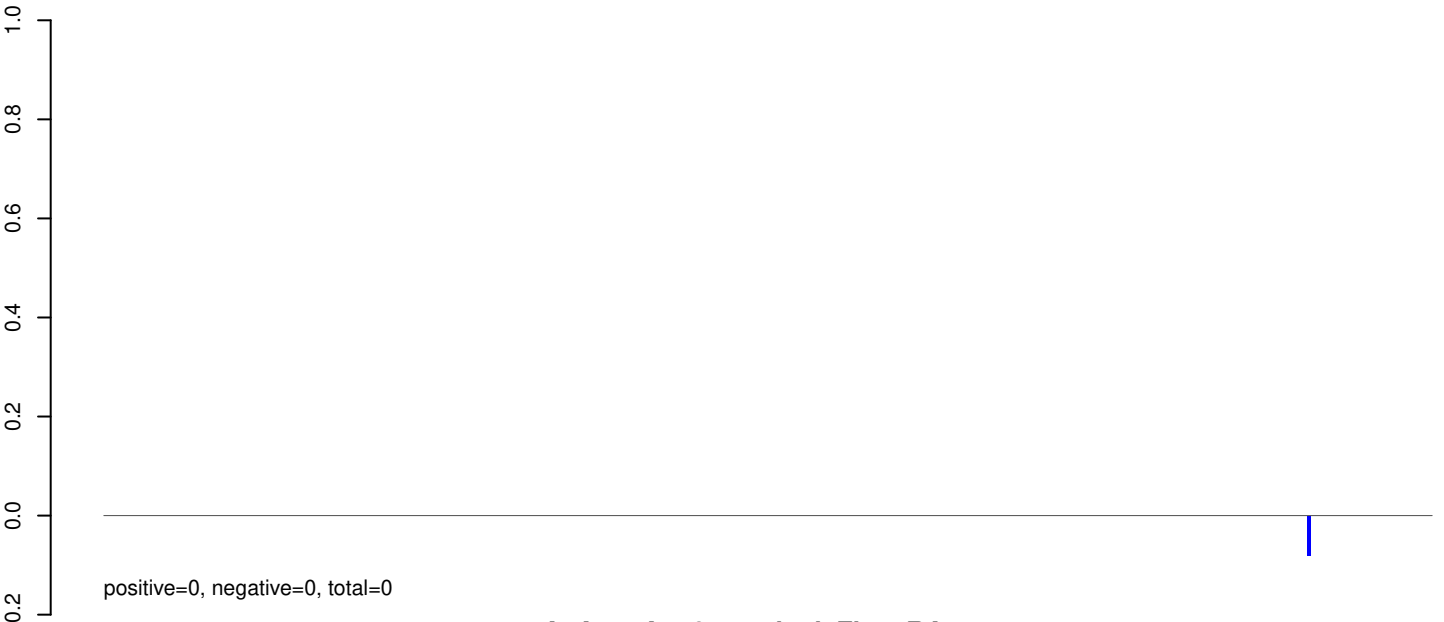


0 2000 4000 6000 8000 10000

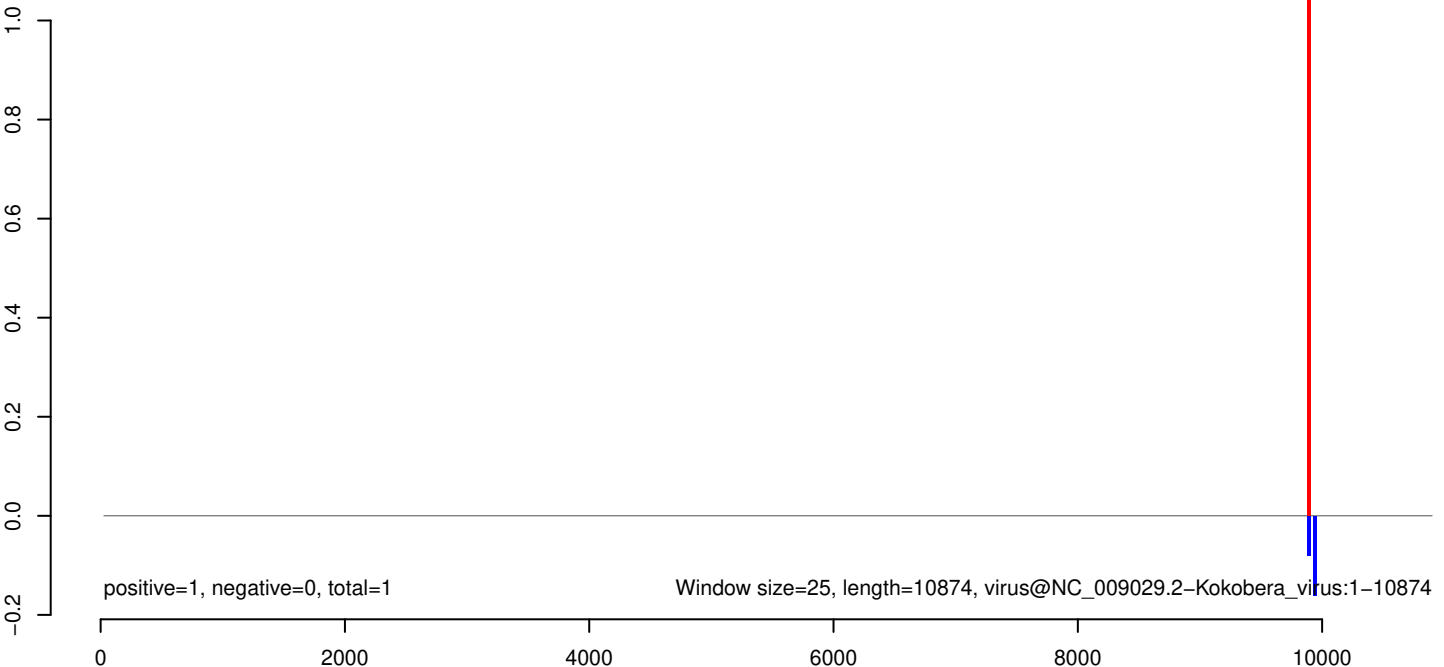
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



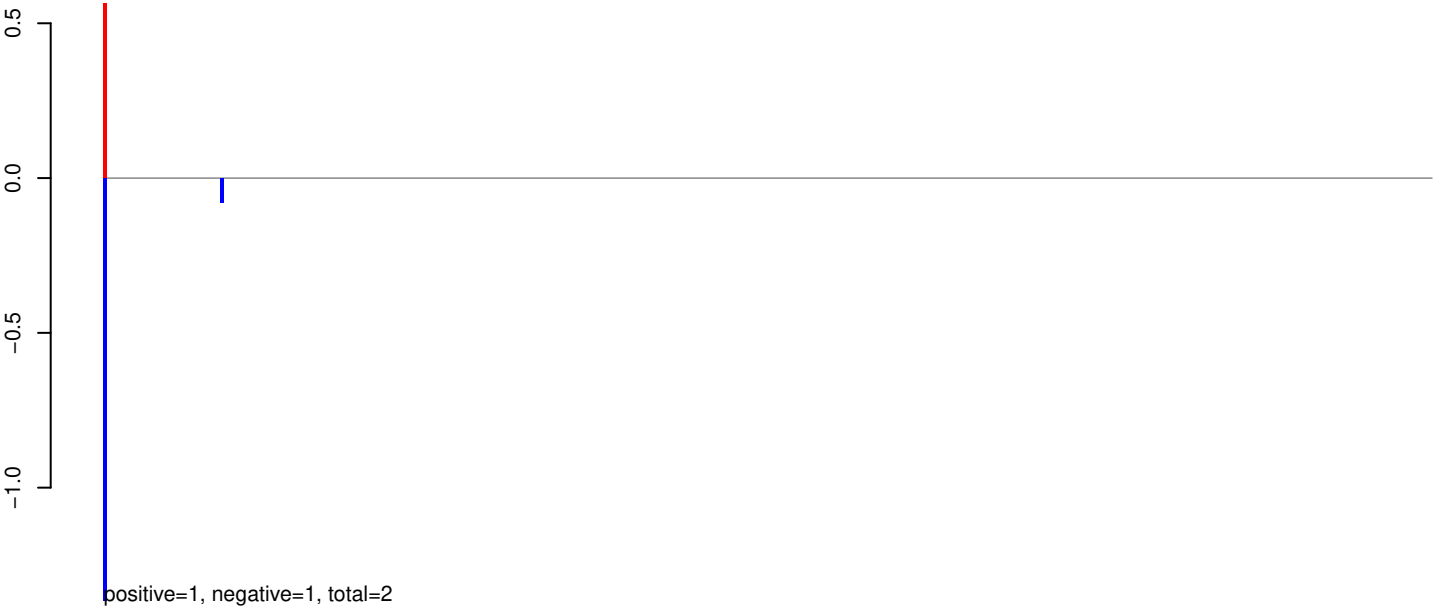
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



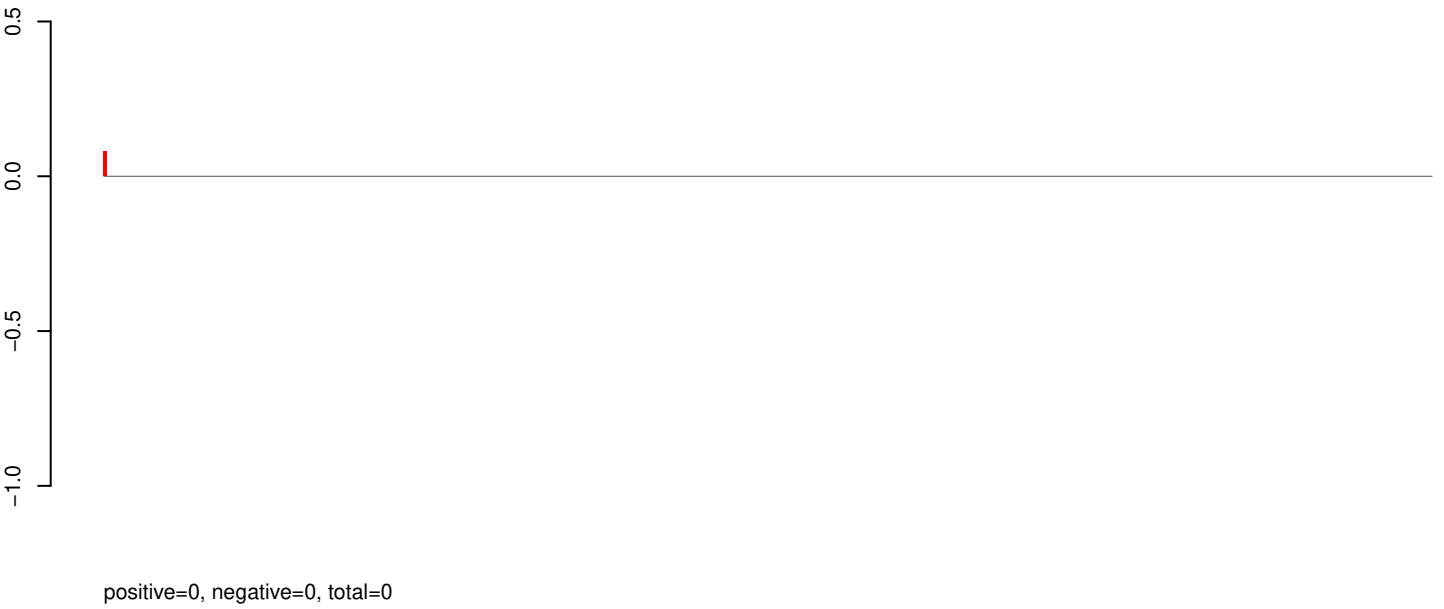
AeAeg_Aag2_mock_dsFluc_RA.rep



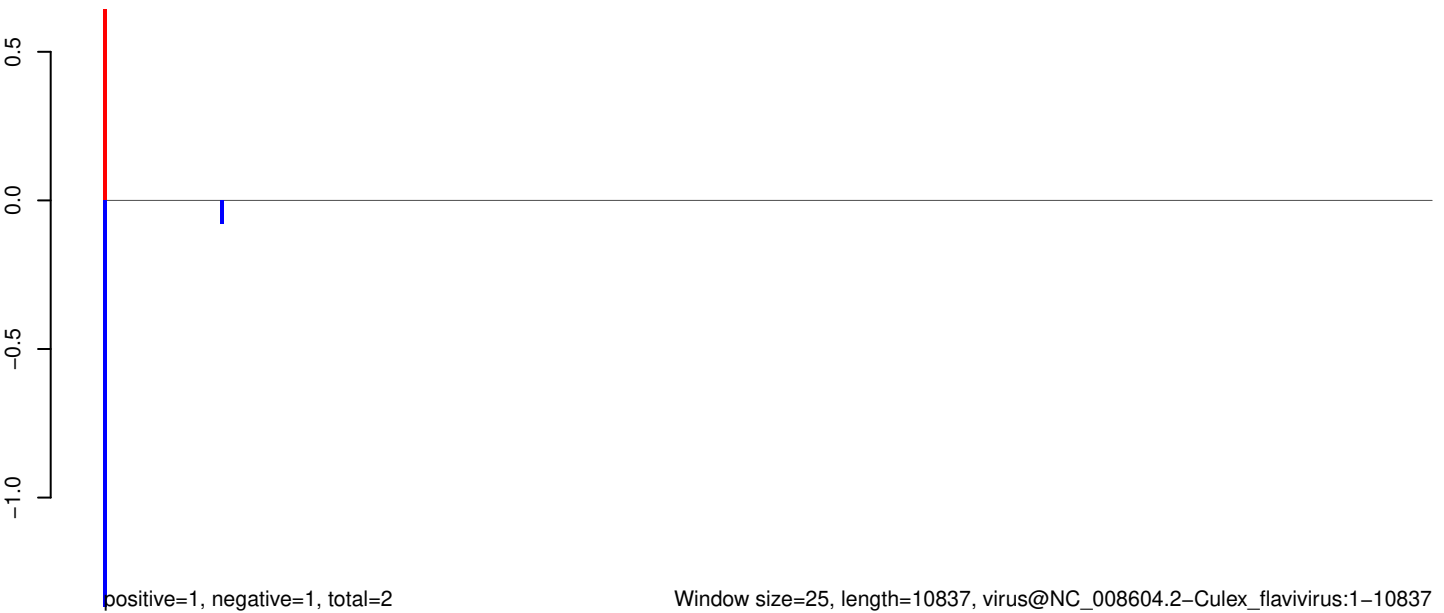
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



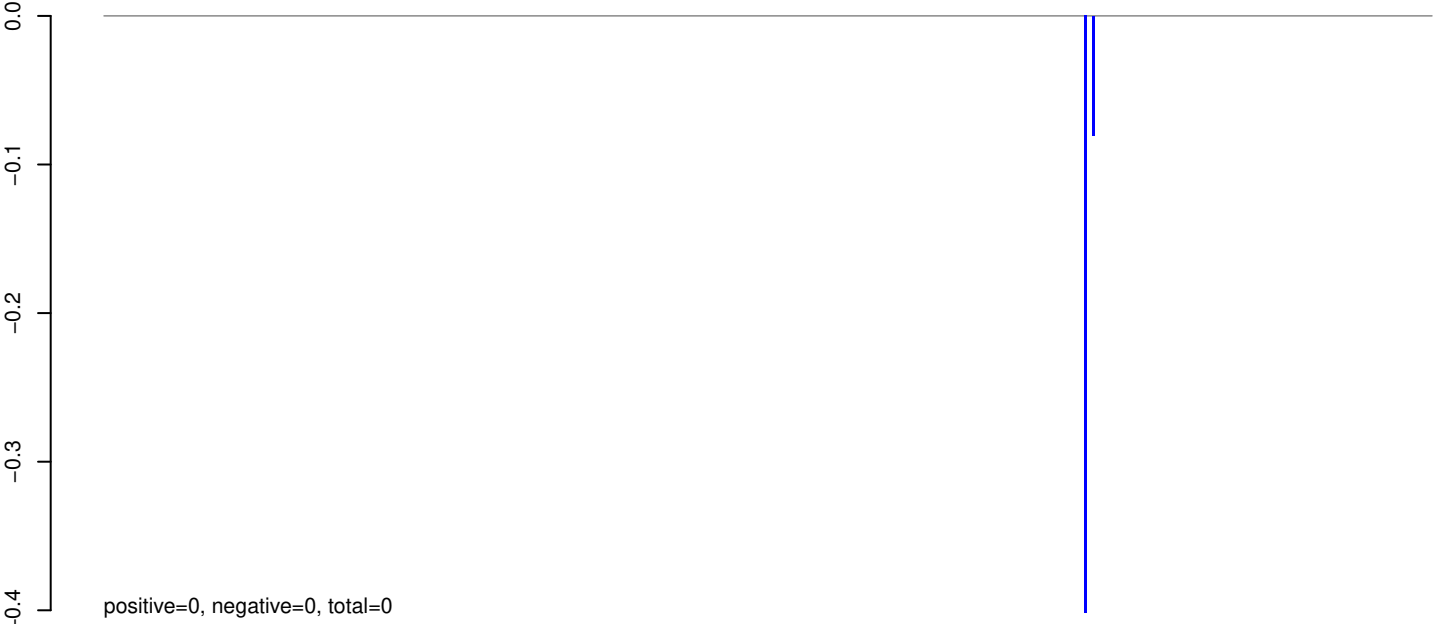
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



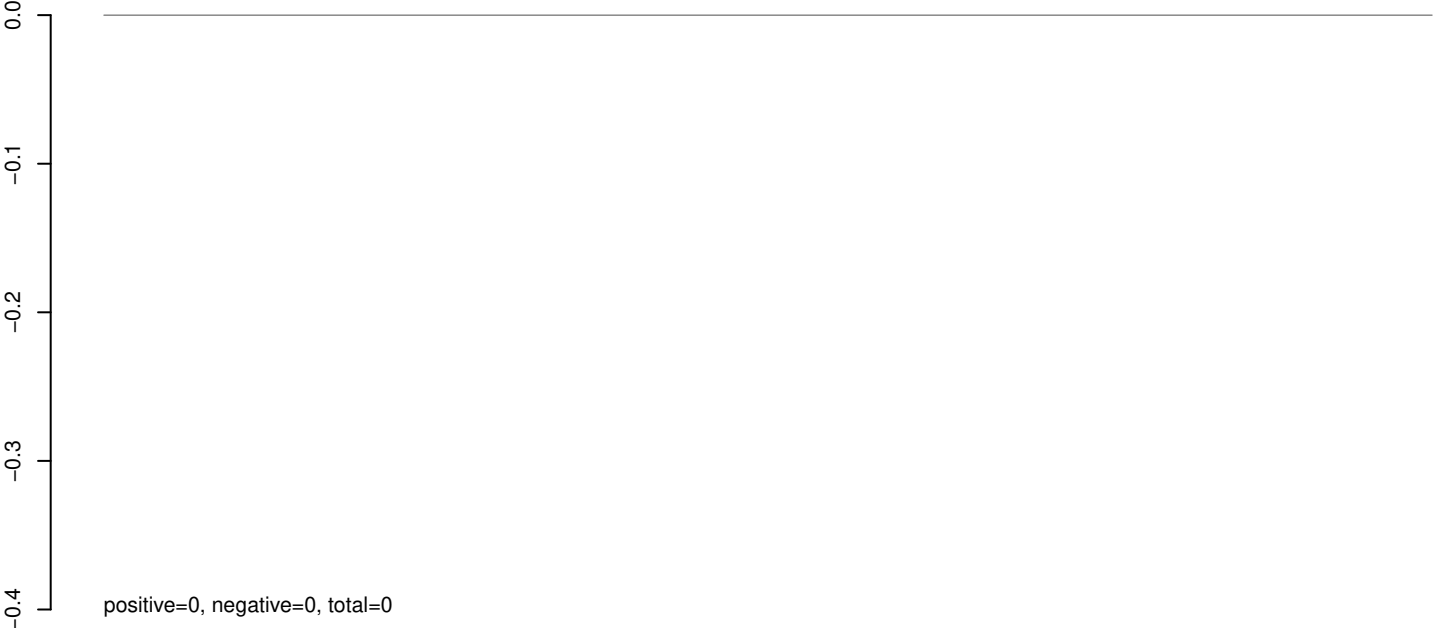
AeAeg_Aag2_mock_dsFluc_RA.rep



AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep

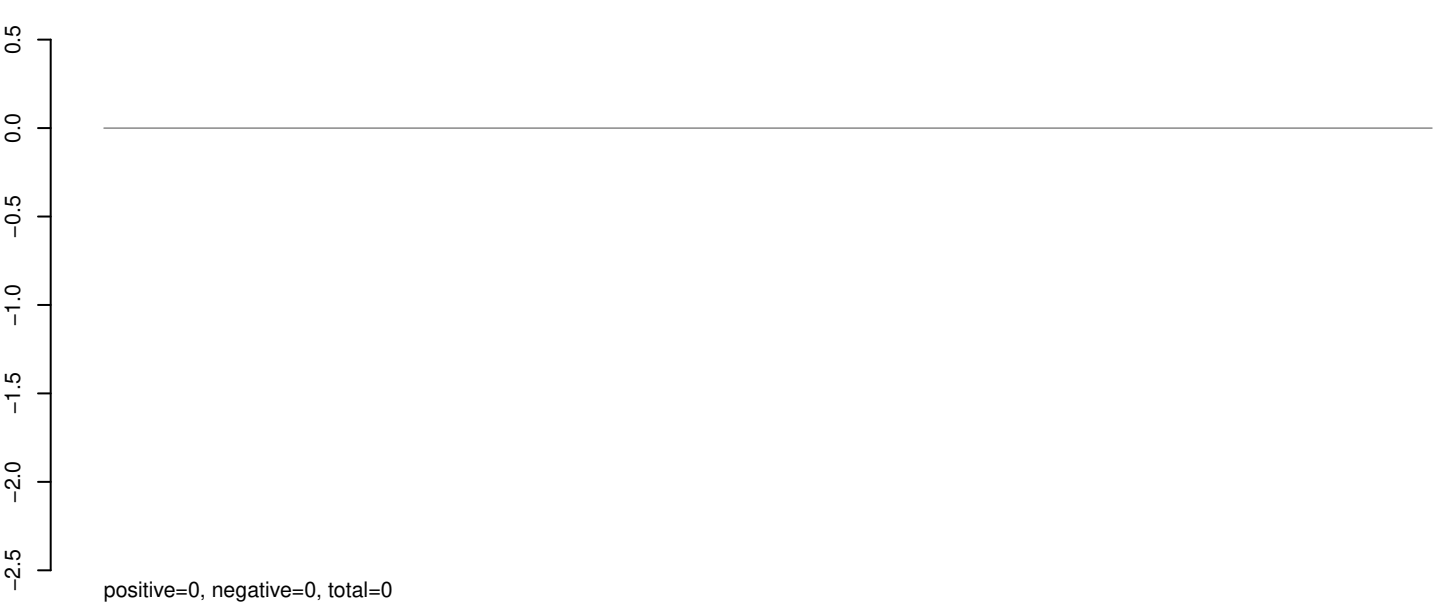


0 2000 4000 6000 8000 10000 12000

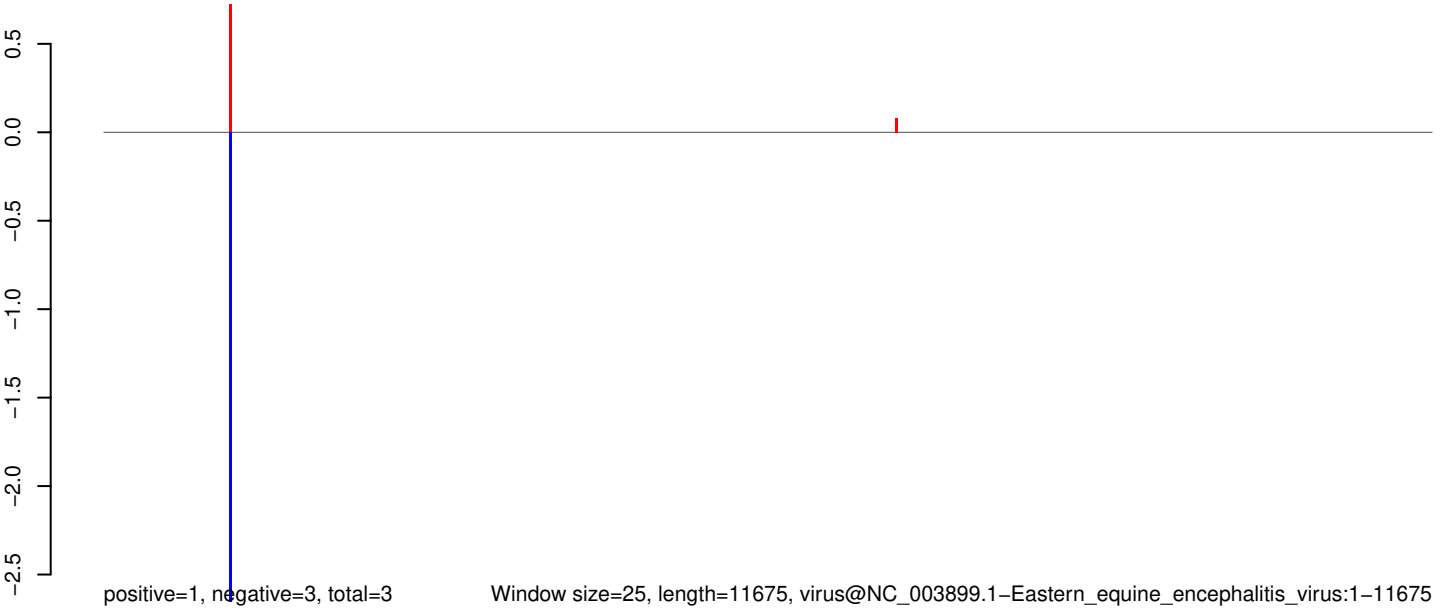
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



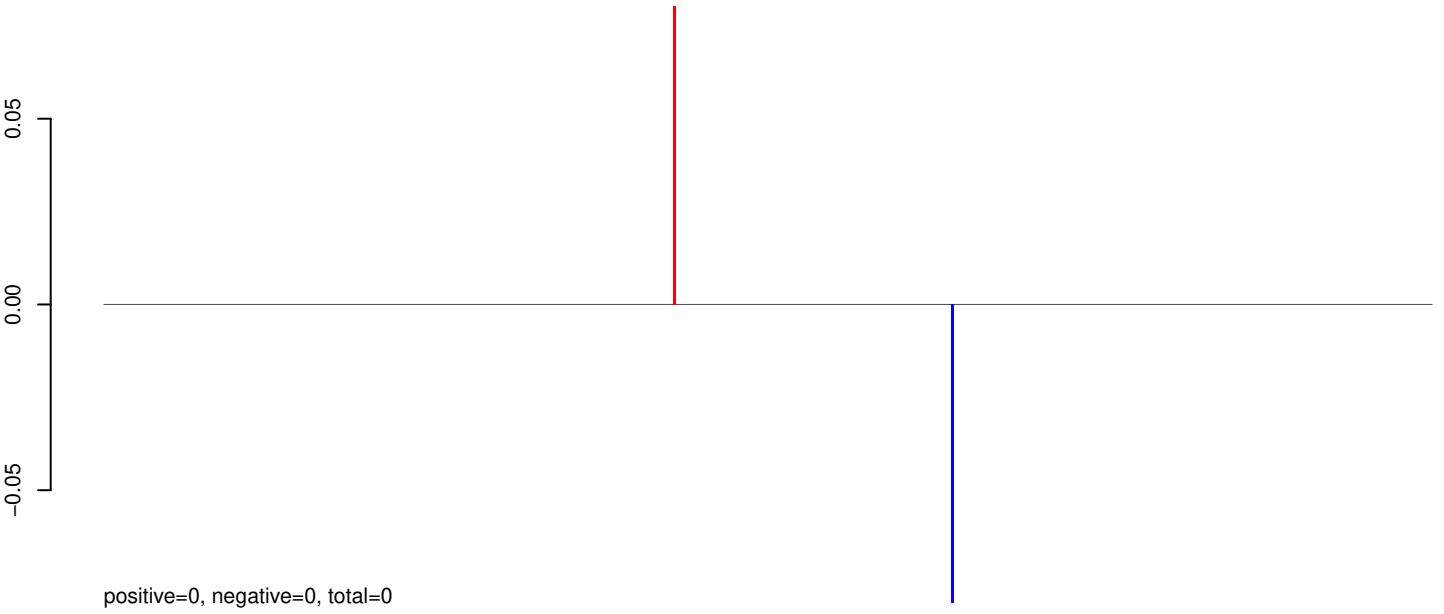
AeAeg_Aag2_mock_dsFluc_RA.rep



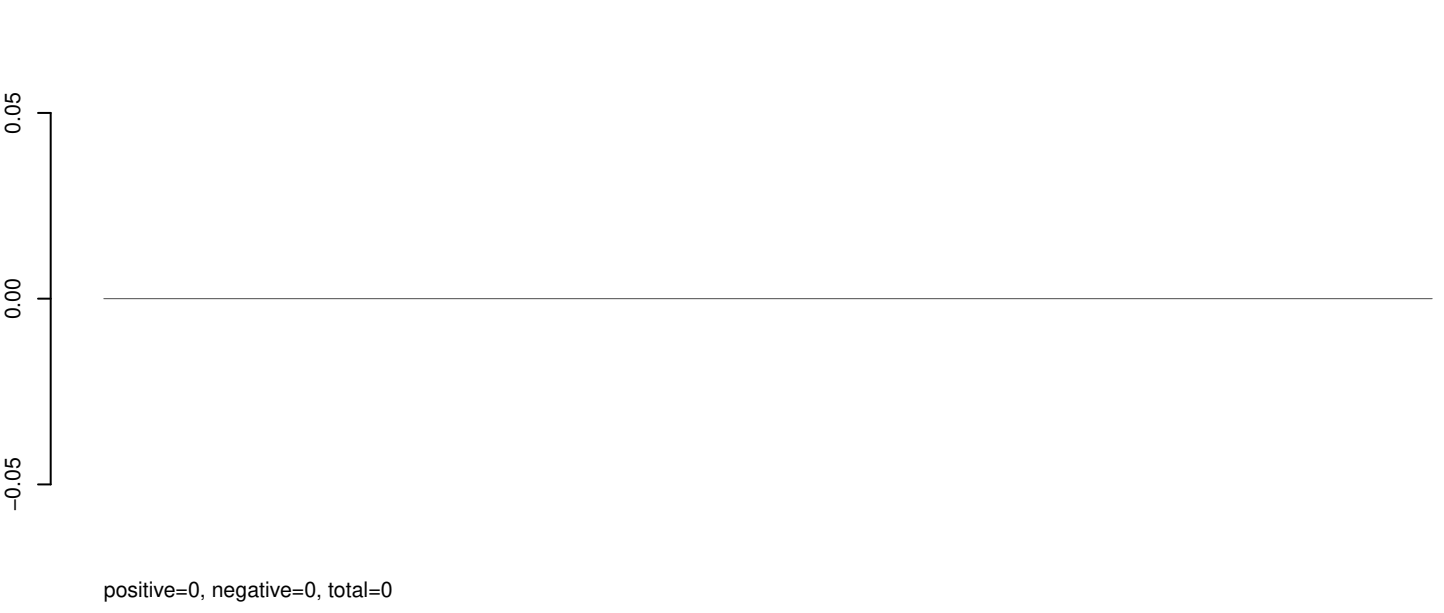
Window size=25, length=11675, virus@NC_003899.1-Eastern_equine_encephalitis_virus:1-11675

0 2000 4000 6000 8000 10000 12000

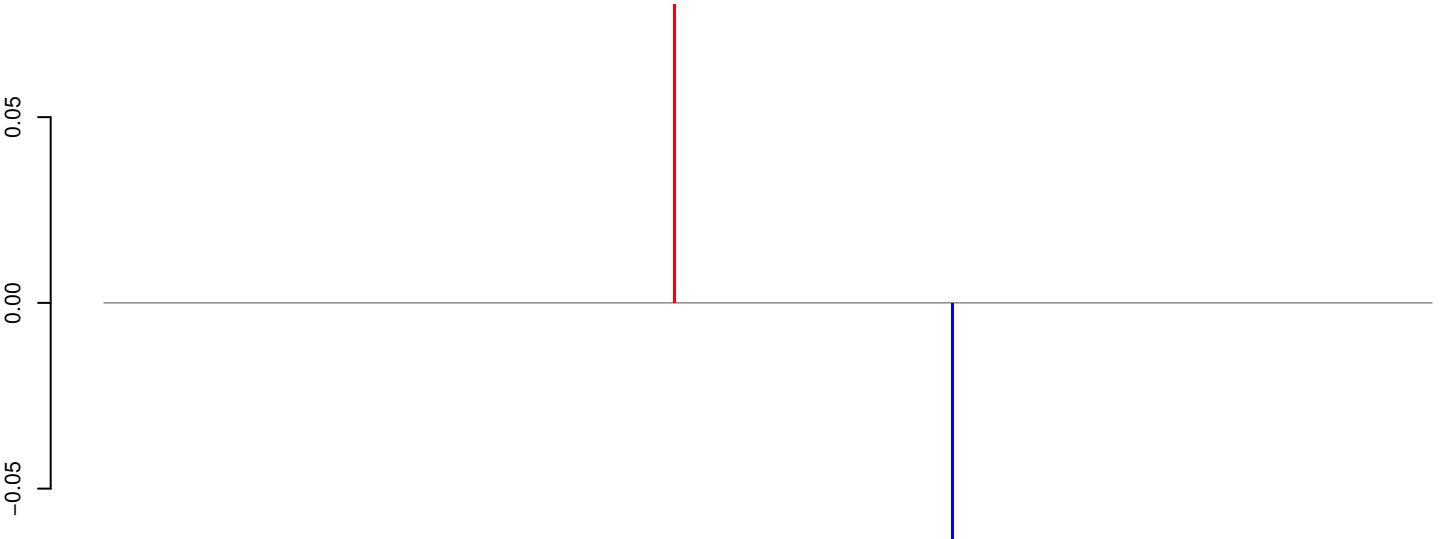
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



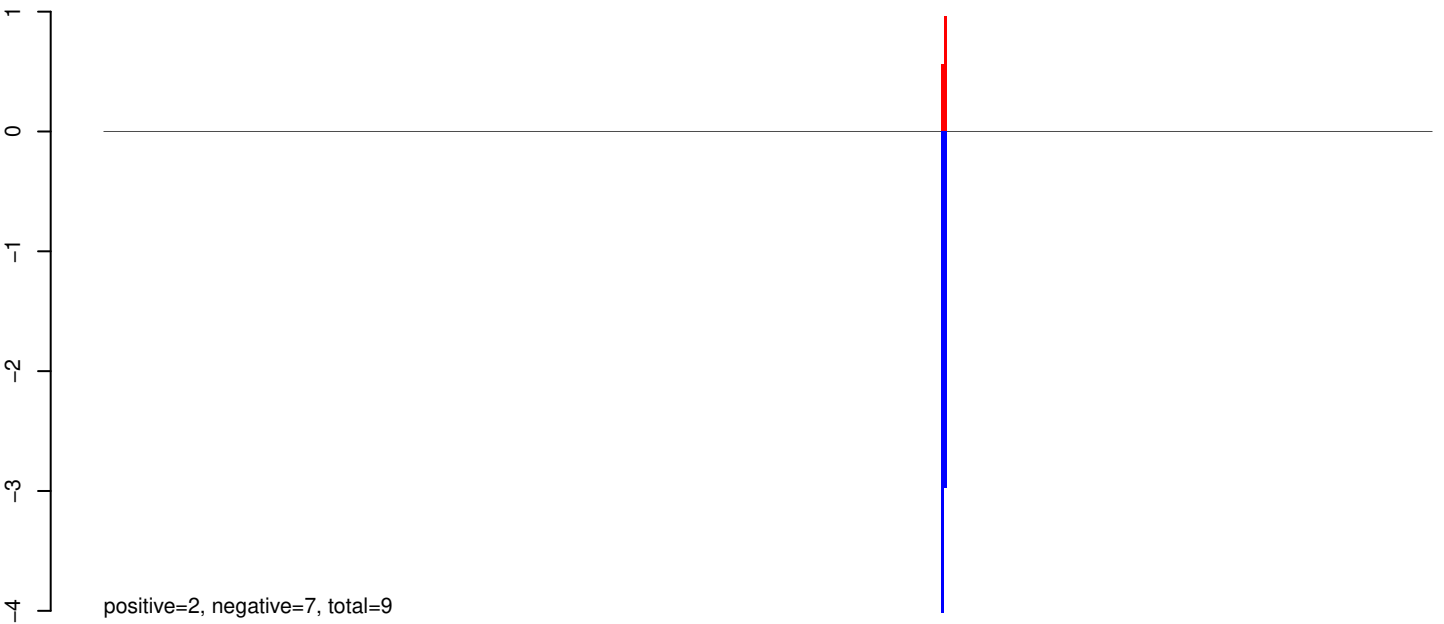
AeAeg_Aag2_mock_dsFluc_RA.rep



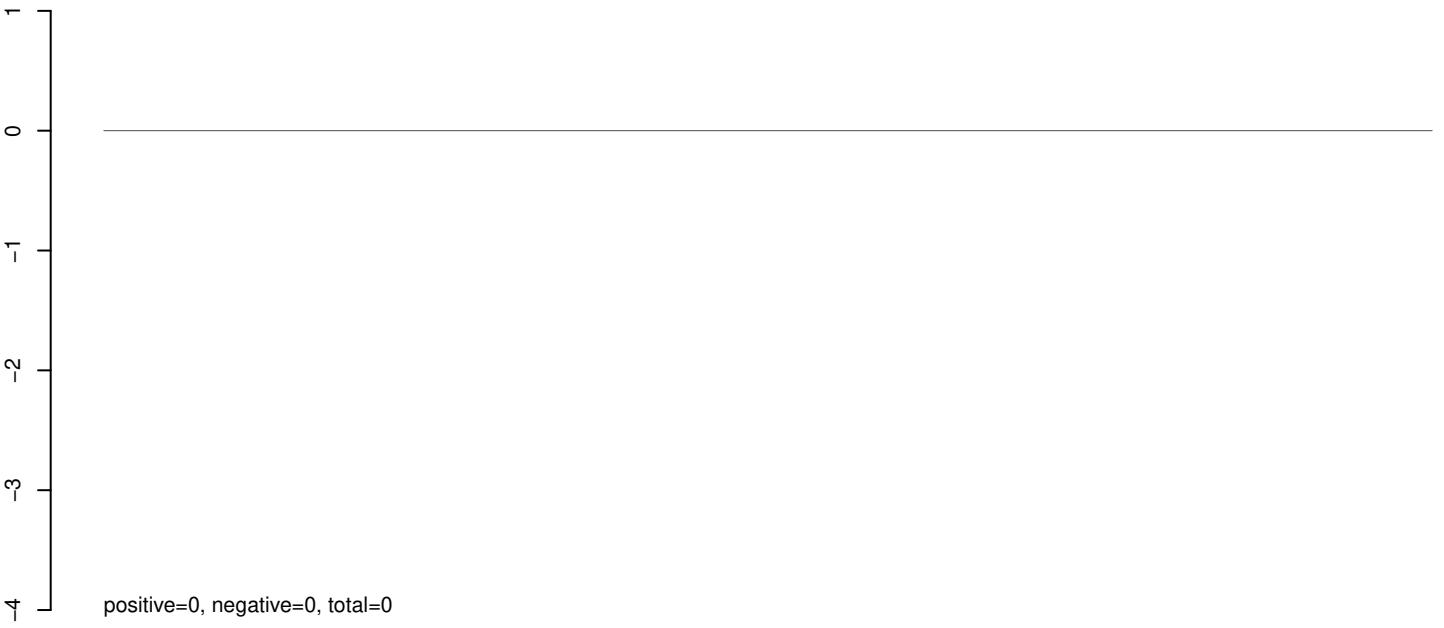
Window size=25, total=12294, virus@NC_001927.1-Bunyamwera_virus_segment_S_plus_segment_M_plus_segment_L:1-12294

0 2000 4000 6000 8000 10000 12000

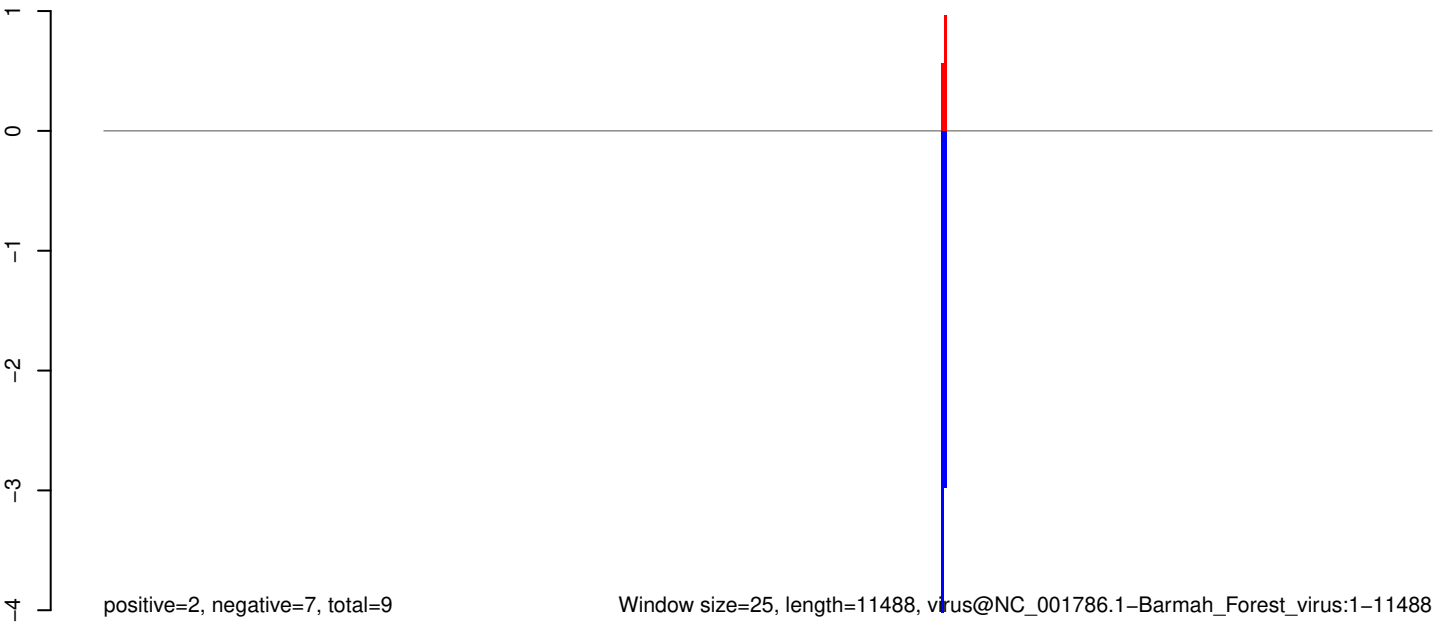
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



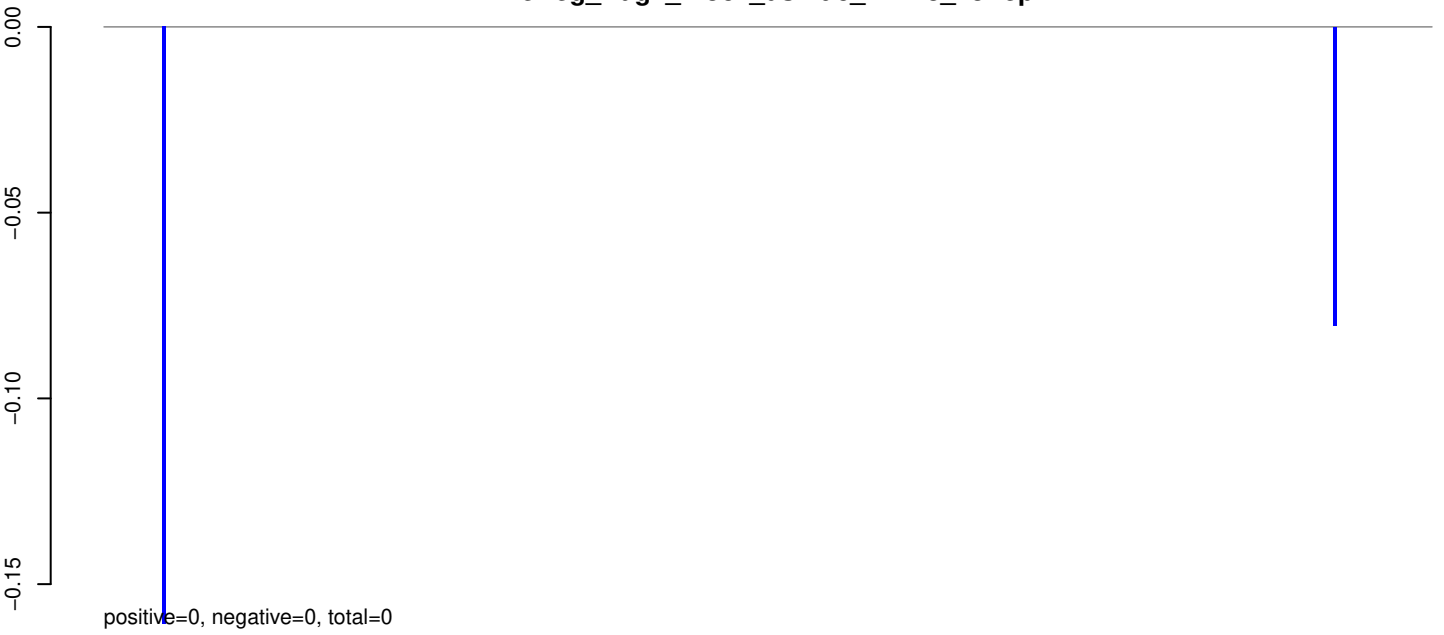
AeAeg_Aag2_mock_dsFluc_RA.rep



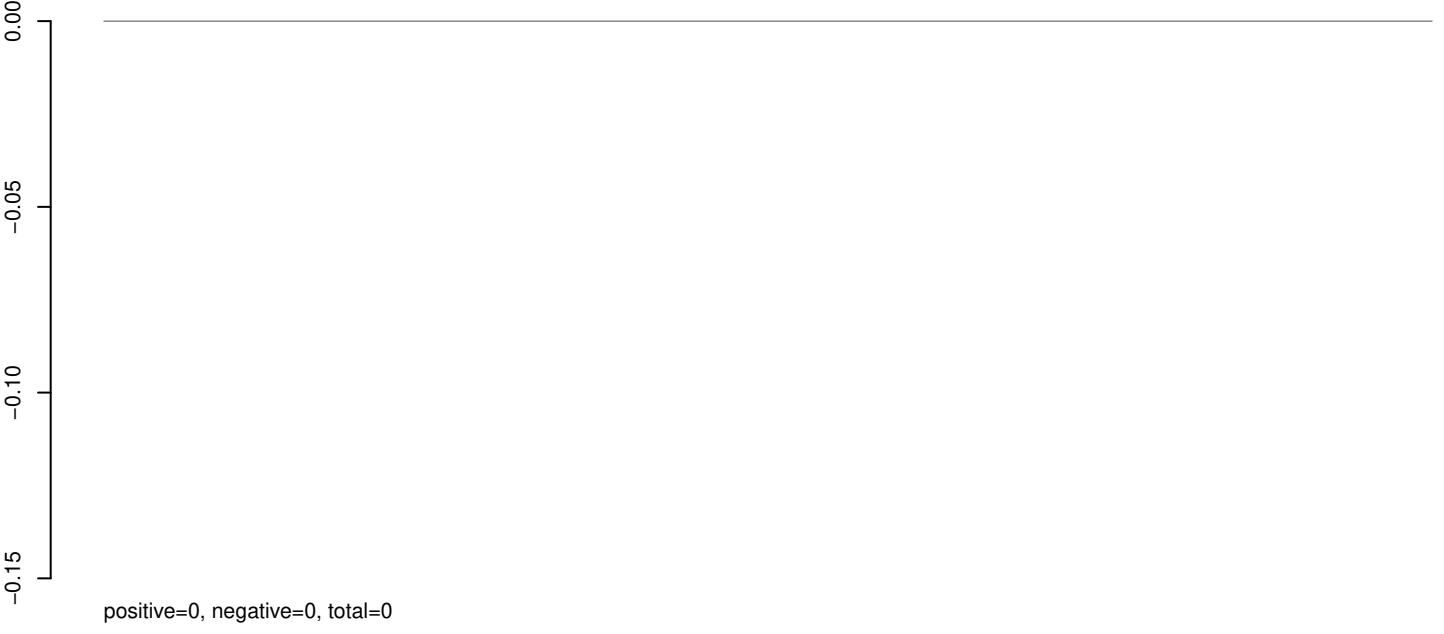
Window size=25, length=11488, virus@NC_001786.1-Barmah_Forest_virus:1-11488

0 2000 4000 6000 8000 10000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep



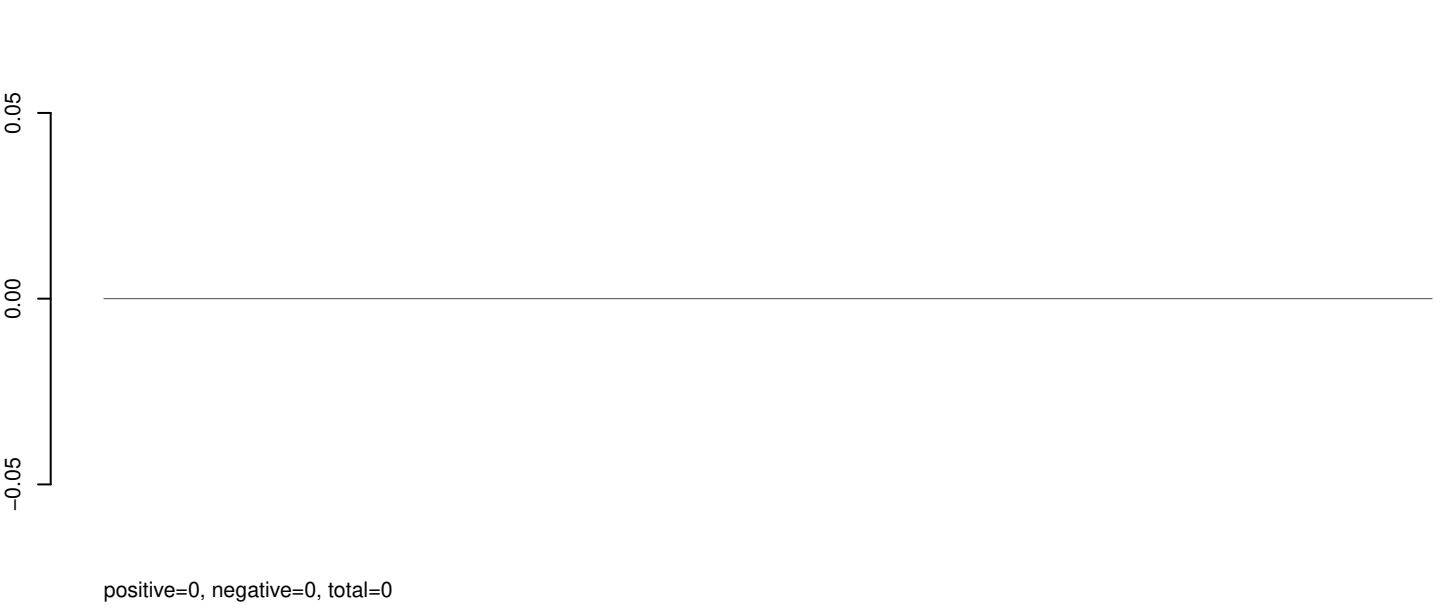
Window size=25, length=10735, virus@NC_001477.1-Dengue_virus_1:1-10735

0 2000 4000 6000 8000 10000

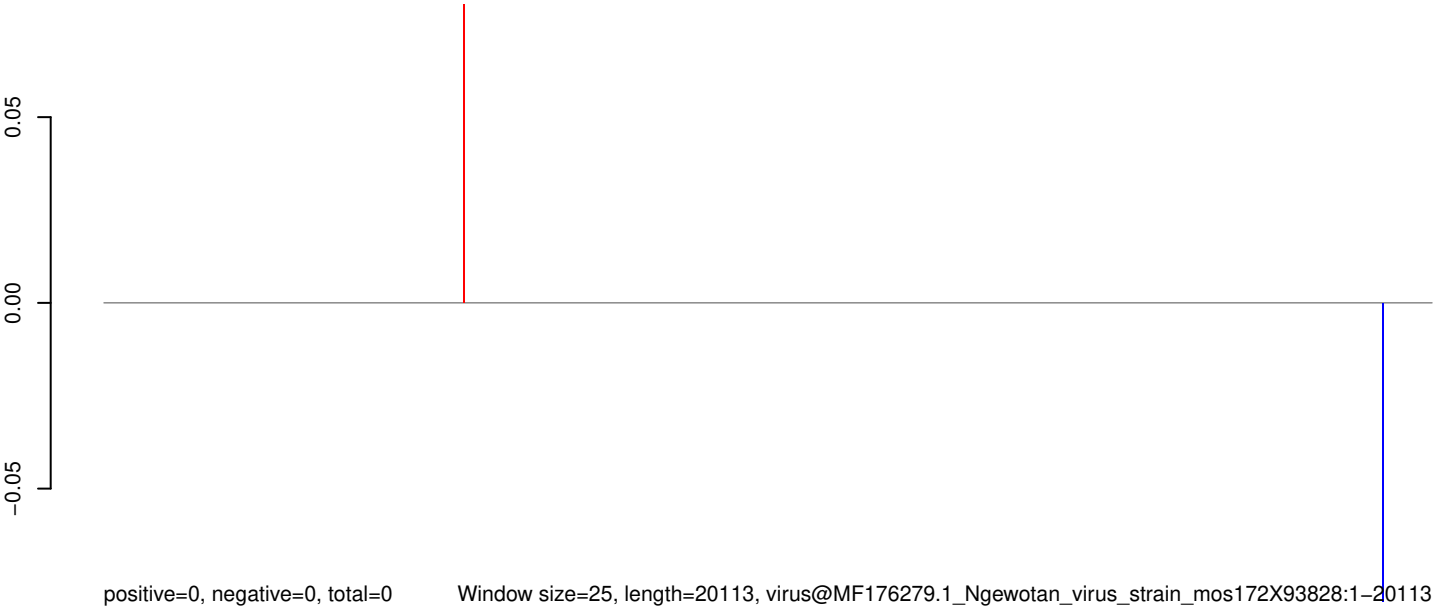
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

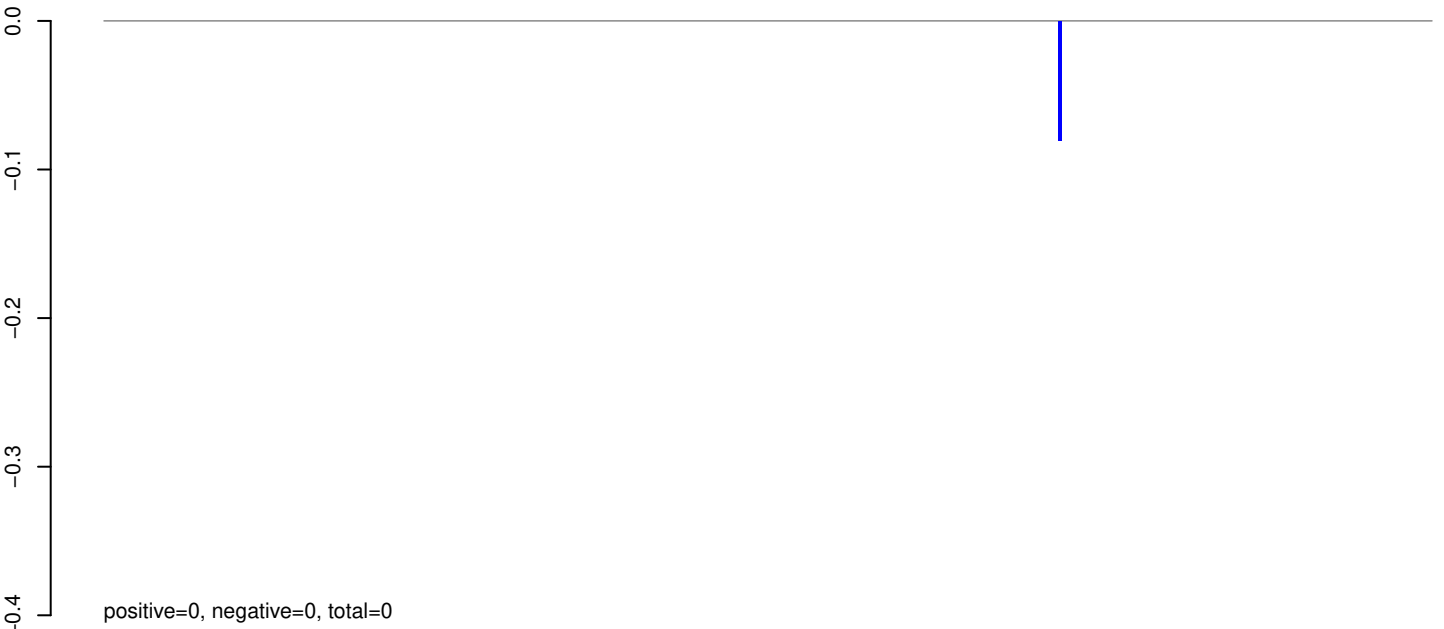


AeAeg_Aag2_mock_dsFluc_RA.rep

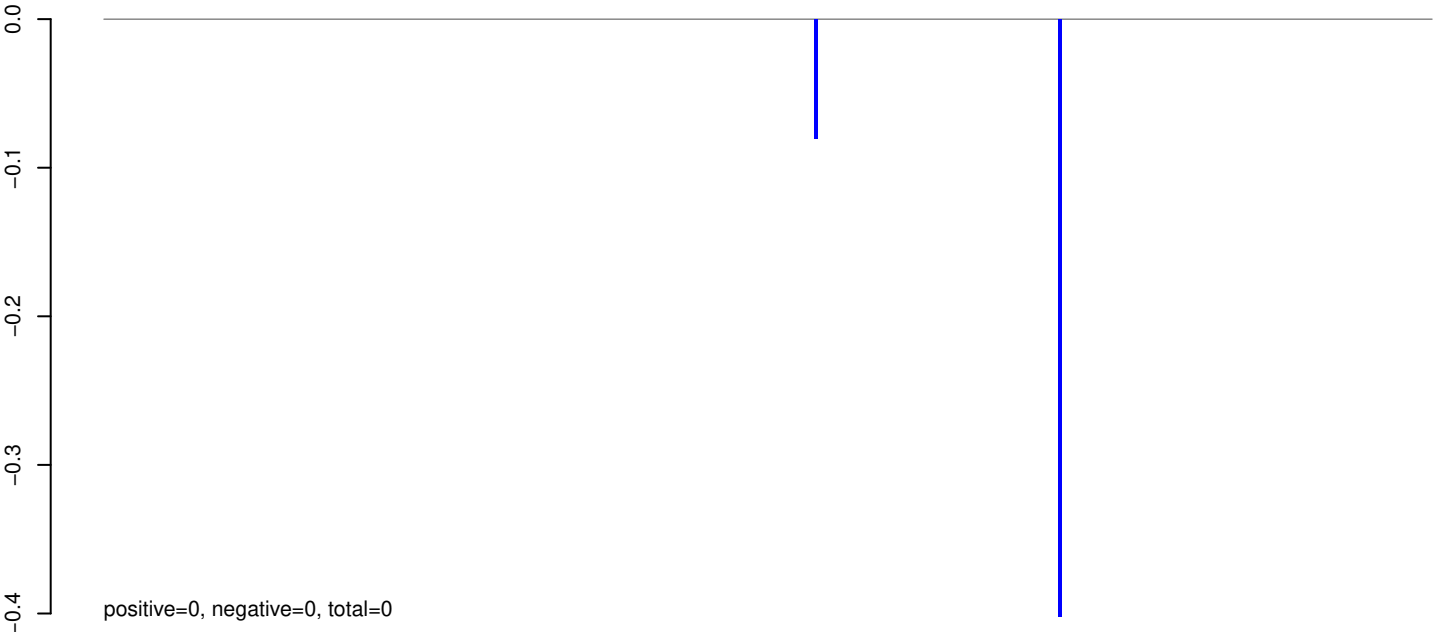


0 5000 10000 15000 20000

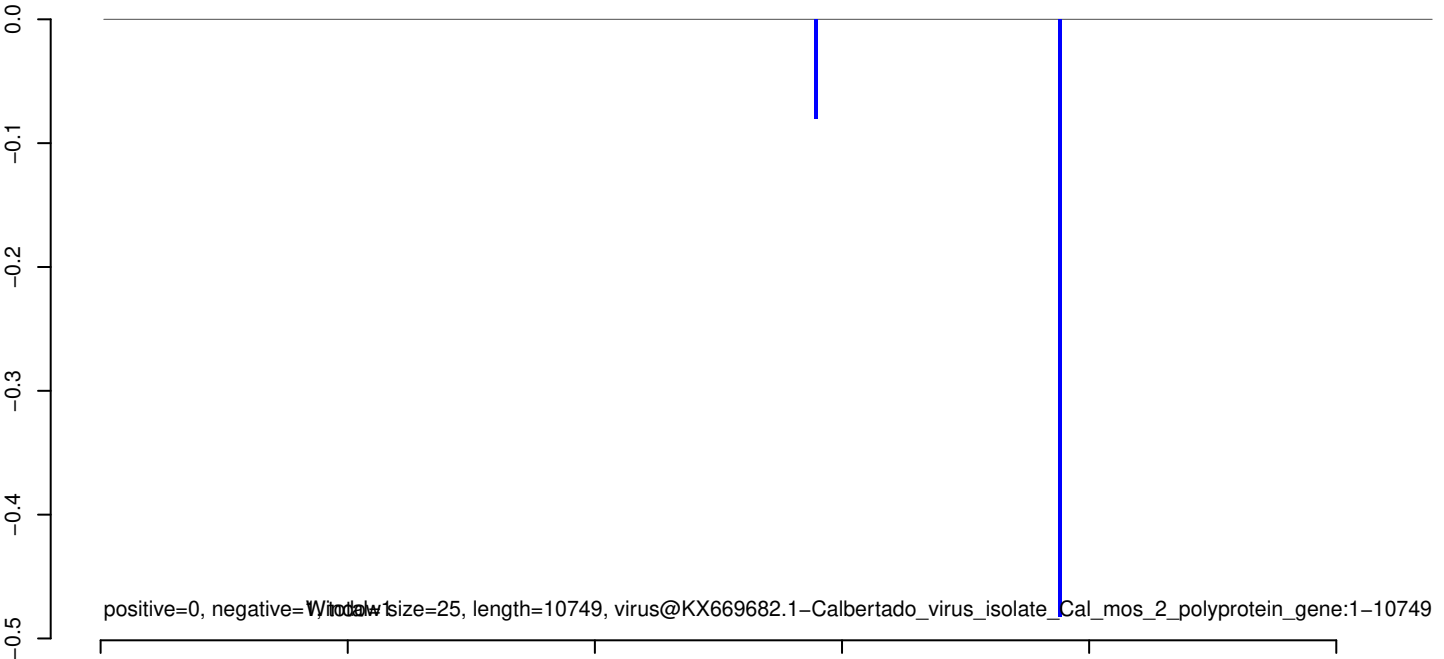
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

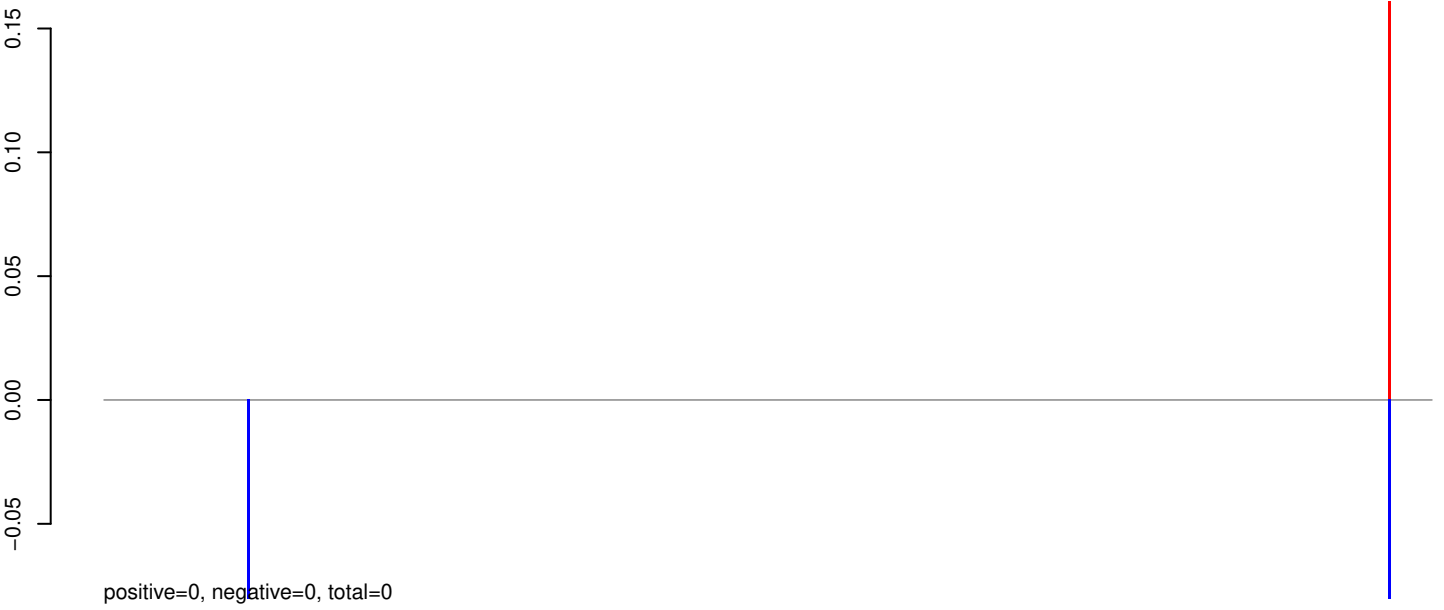


AeAeg_Aag2_mock_dsFluc_RA.rep

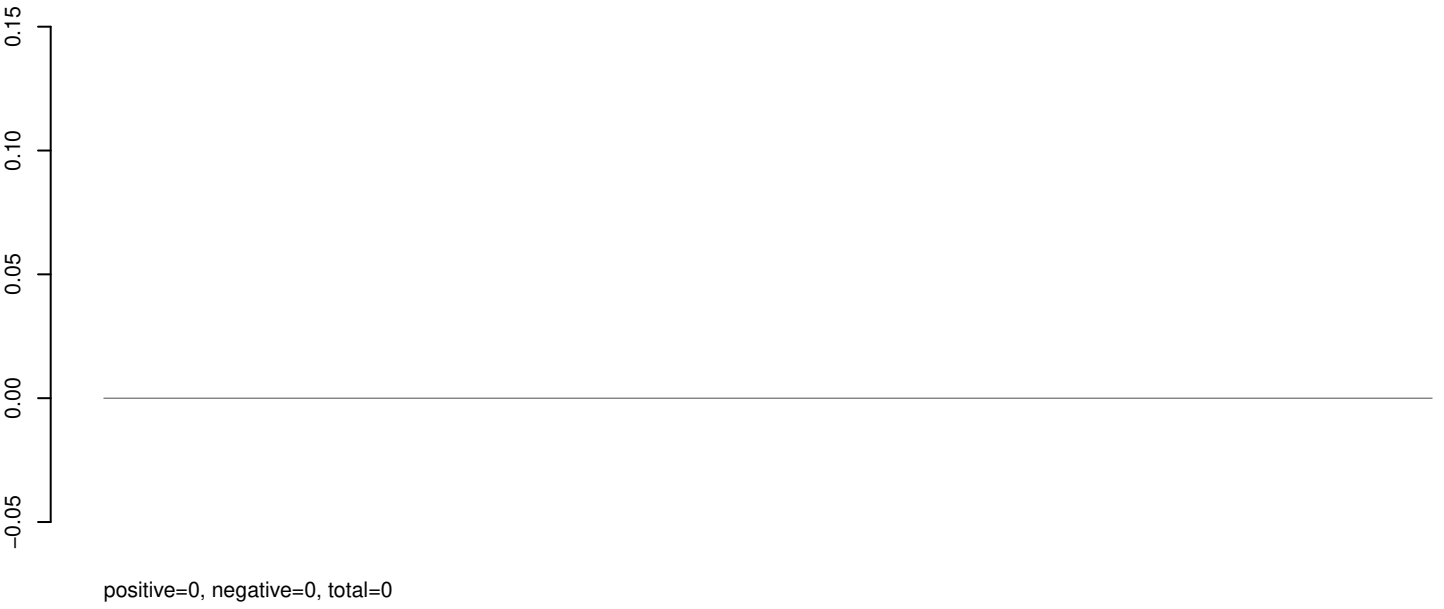


Window size=25, length=10749, virus@KX669682.1-Calbertado_virus_isolate_Cal_mos_2_polyprotein_gene:1-10749

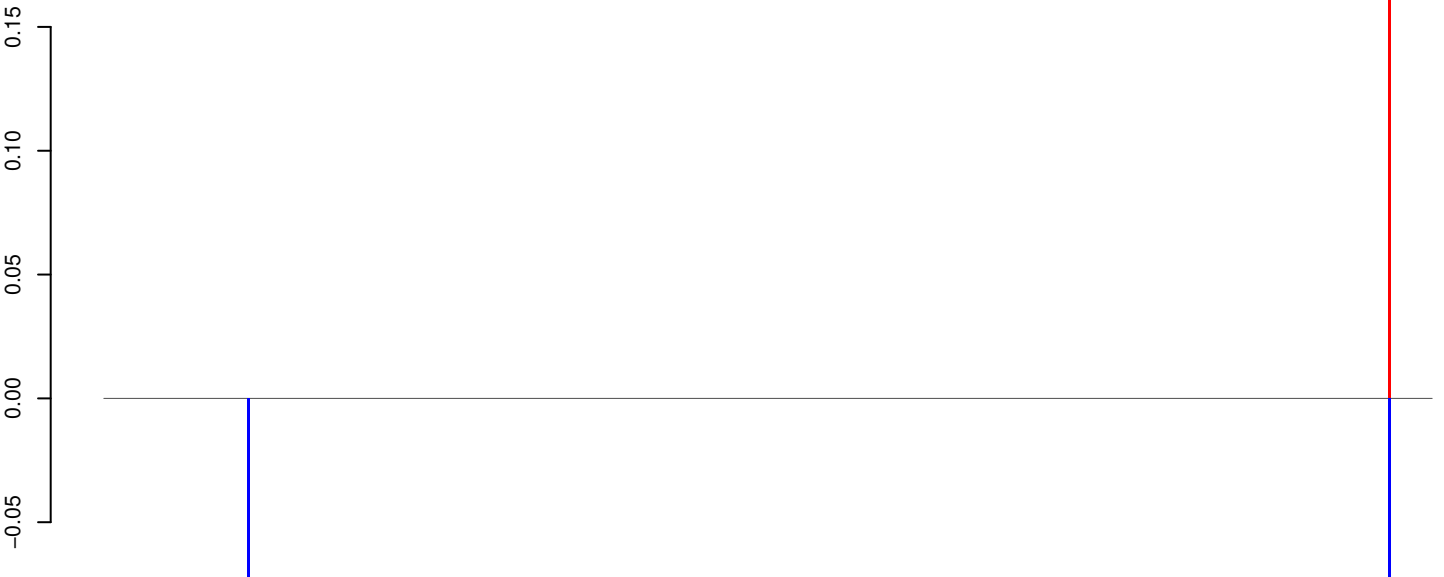
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



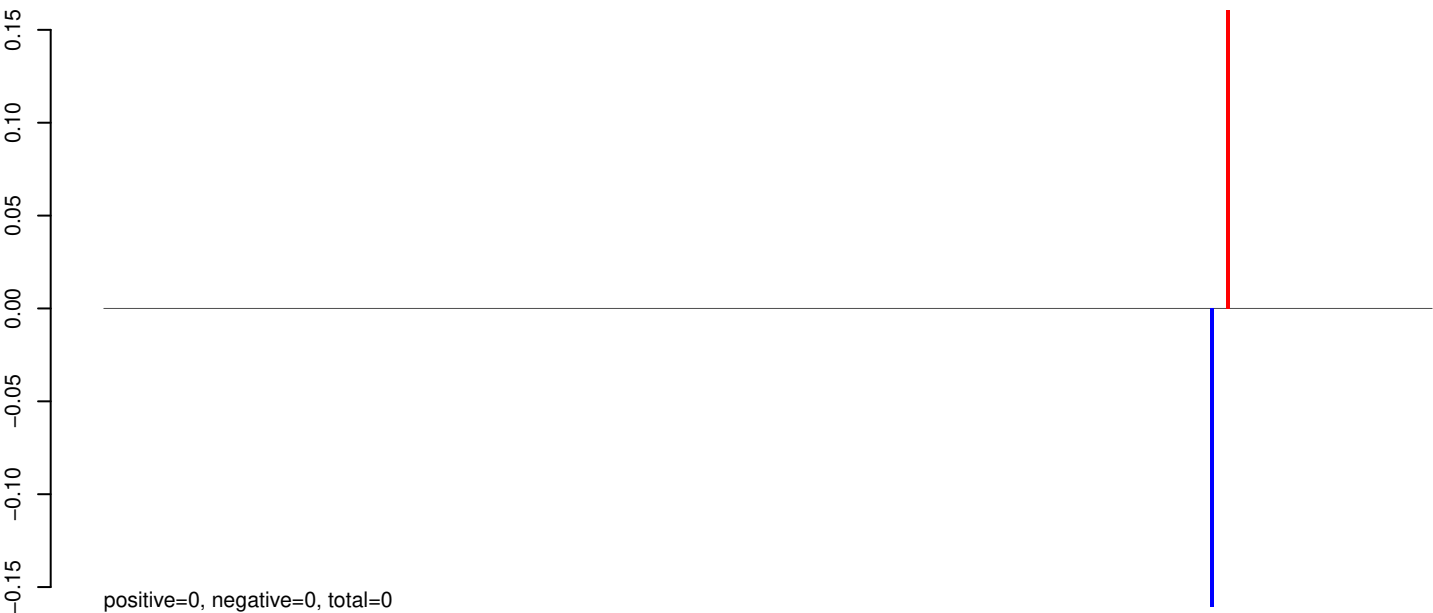
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



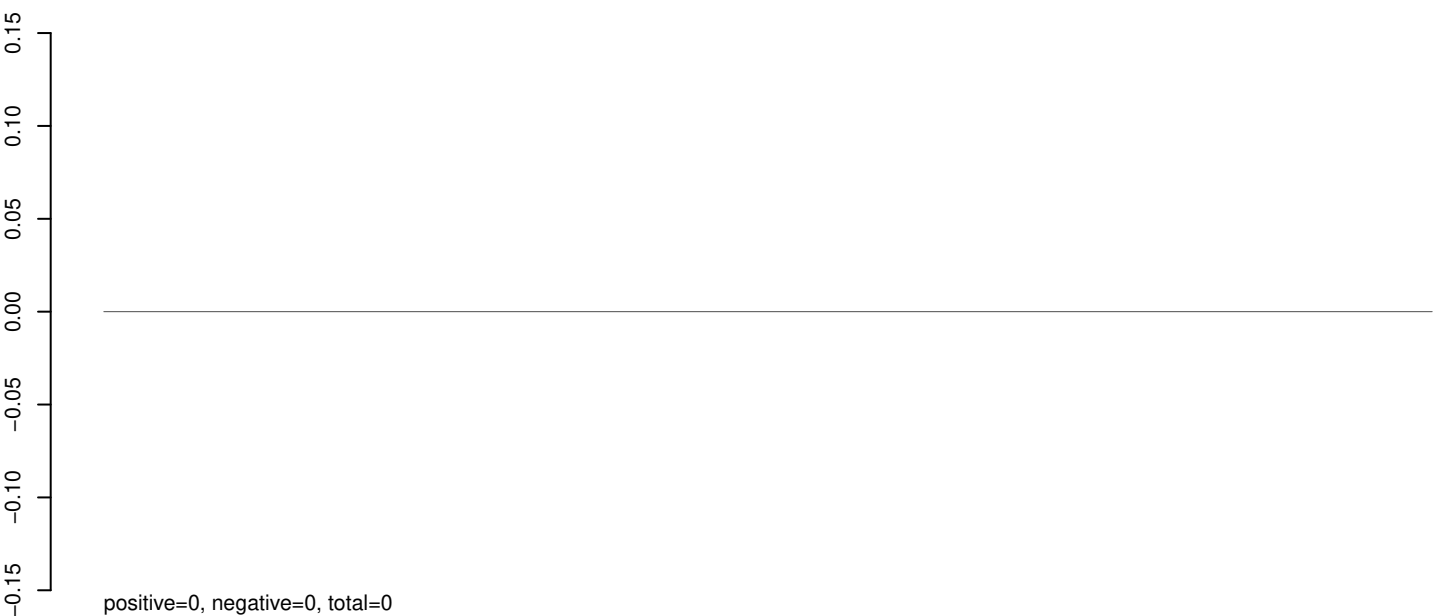
AeAeg_Aag2_mock_dsFluc_RA.rep



AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



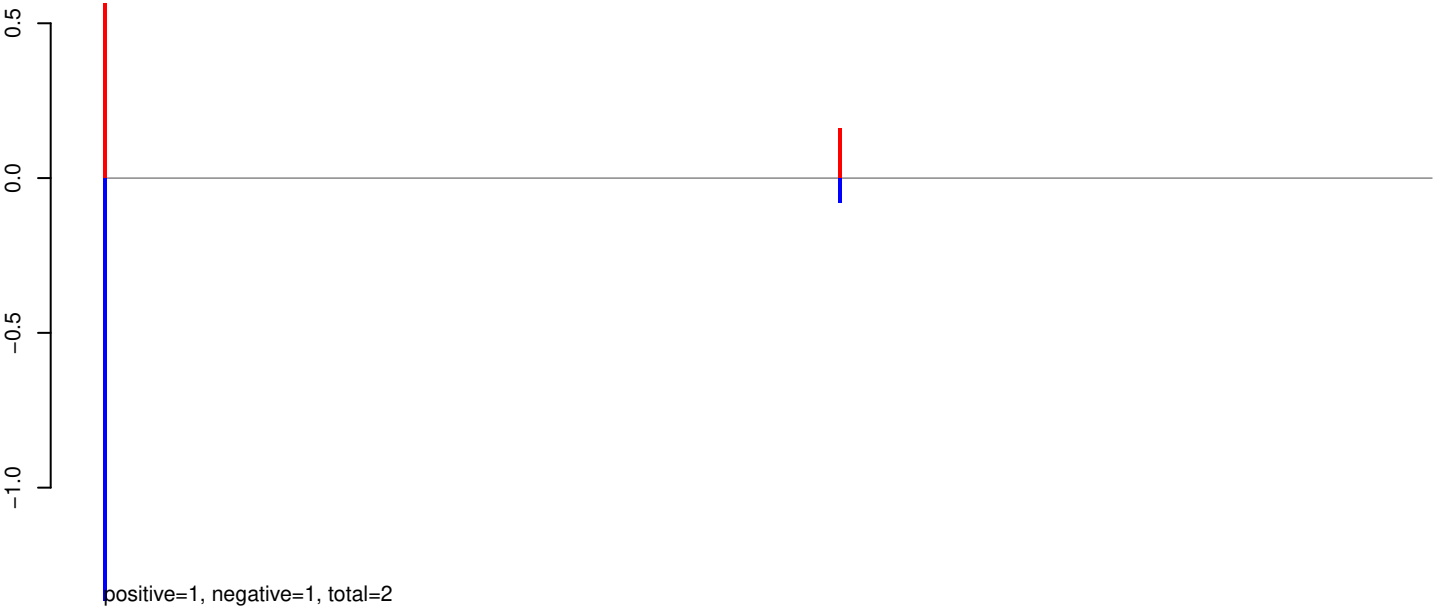
AeAeg_Aag2_mock_dsFluc_RA.rep



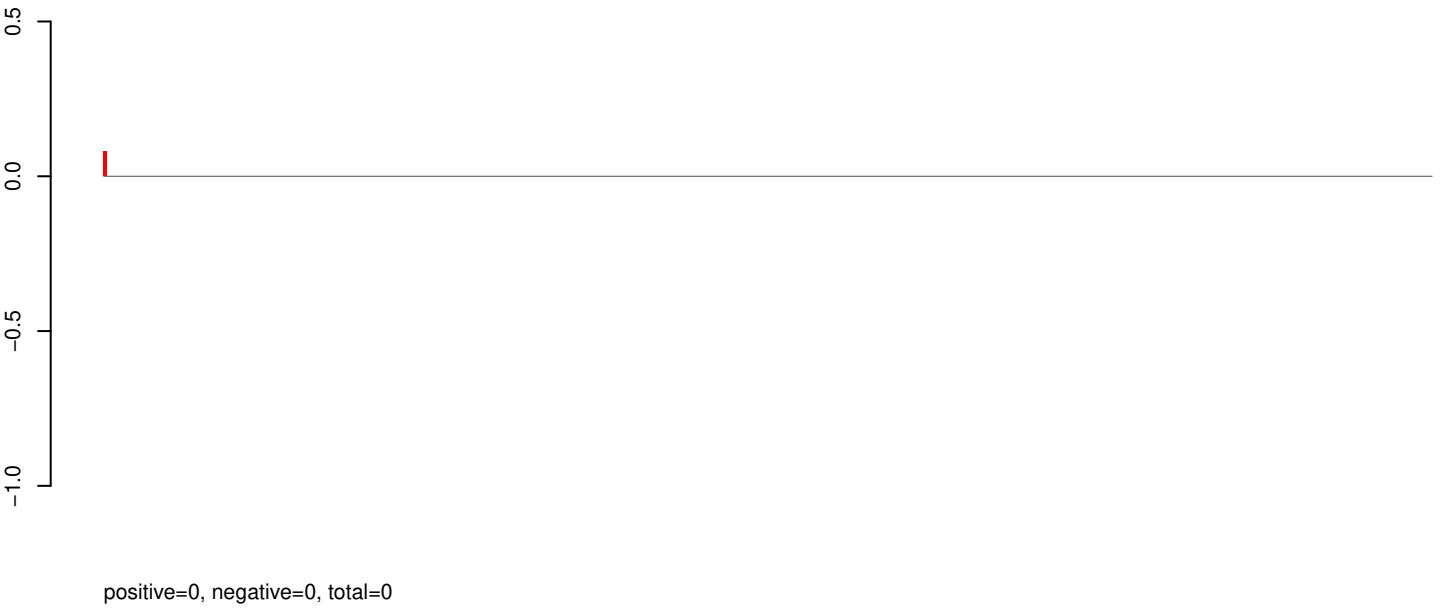
Window size=25, length=10648, virus@KM225264.1-Bainyik_virus_strain_MK7979:1-10648

0 2000 4000 6000 8000 10000

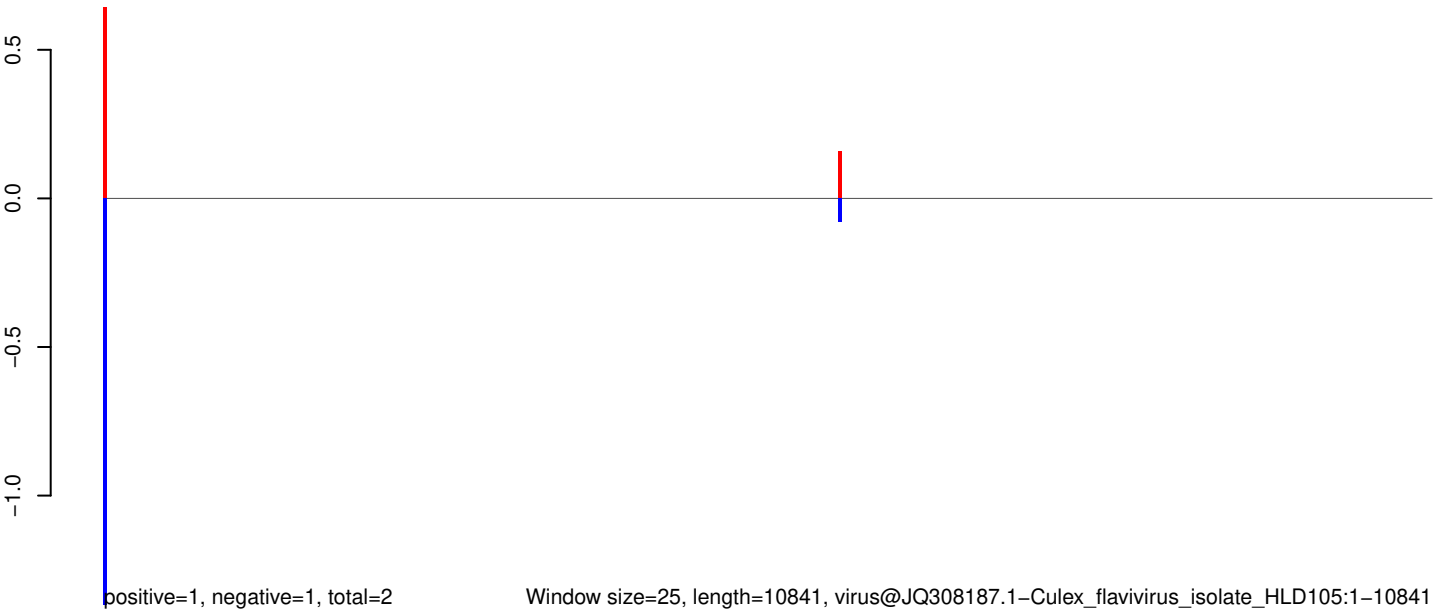
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

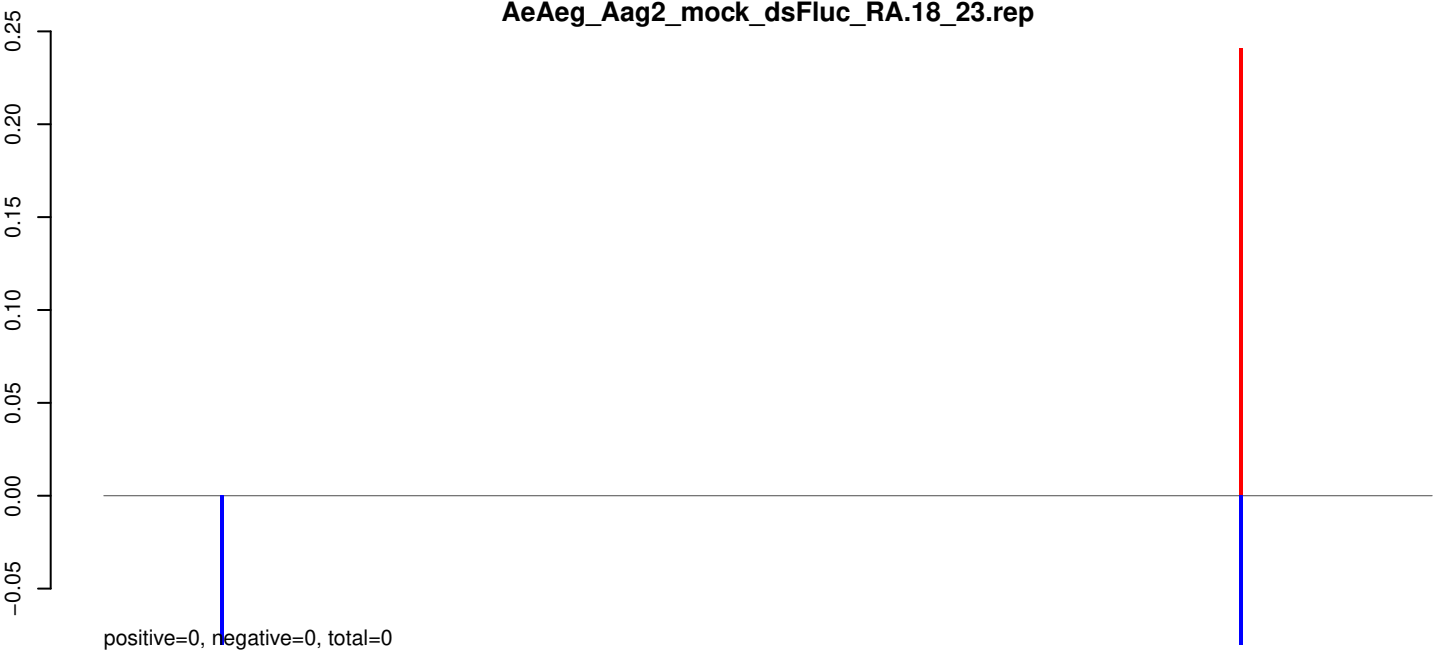


AeAeg_Aag2_mock_dsFluc_RA.rep



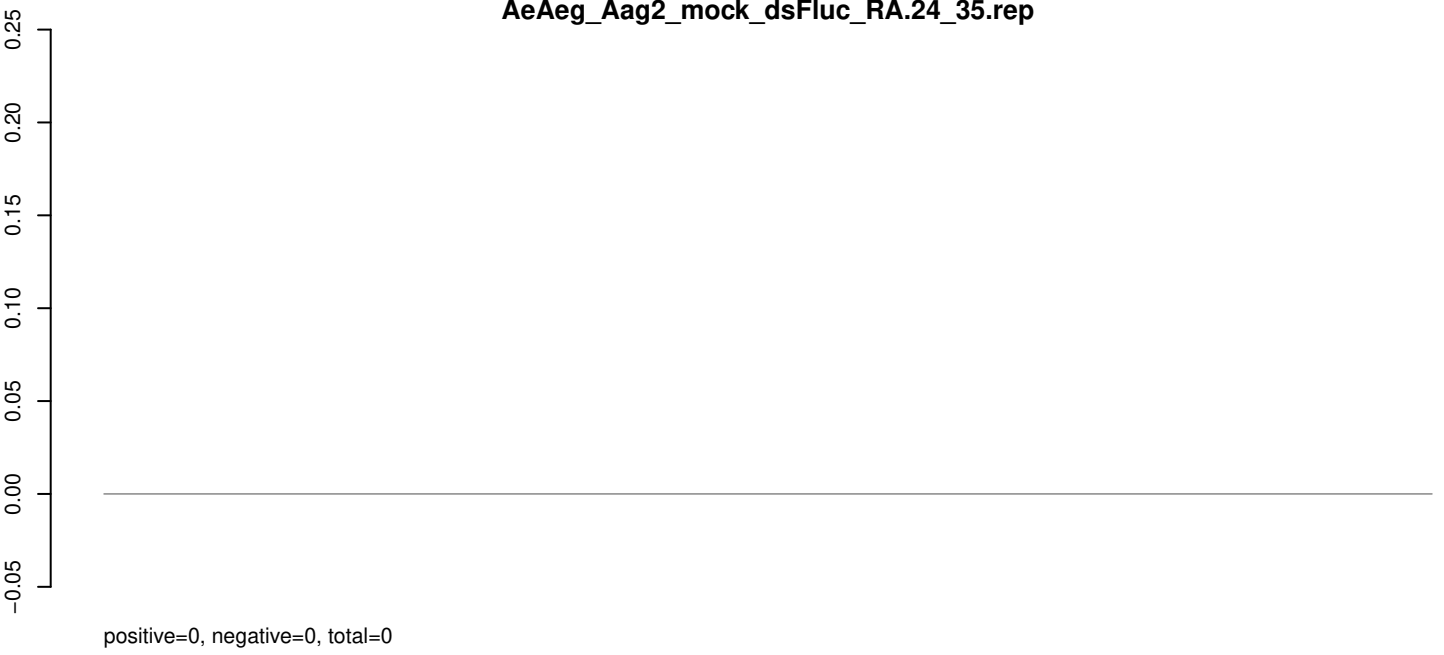
Window size=25, length=10841, virus@JQ308187.1-Culex_flavivirus_isolate_HLD105:1-10841

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_mock_dsFluc_RA.rep

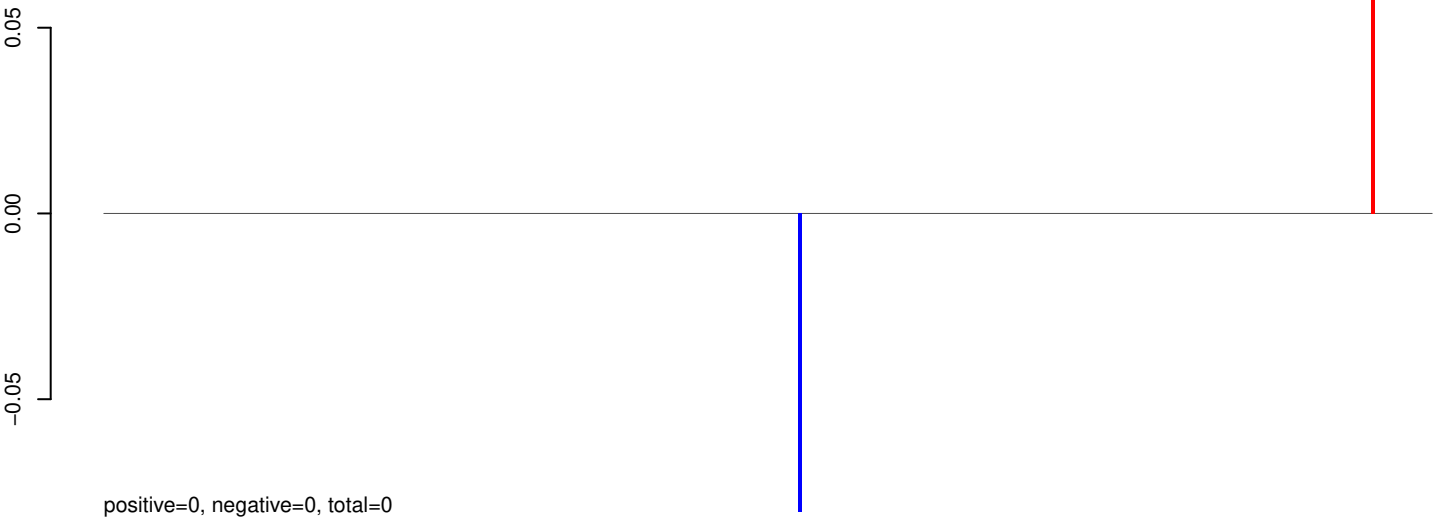


positive=0, negative=0, total=0

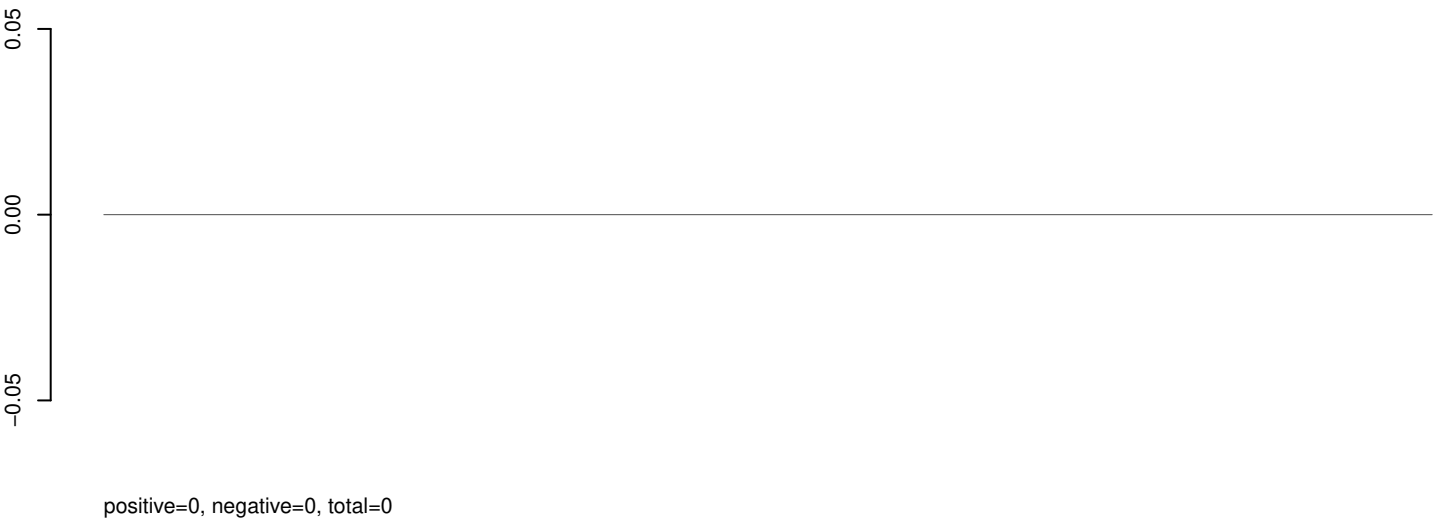
Window size=25, length=10837, virus@FJ663034.1-Culex_flavivirus_isolate_lowa07:1-10837

0 2000 4000 6000 8000 10000

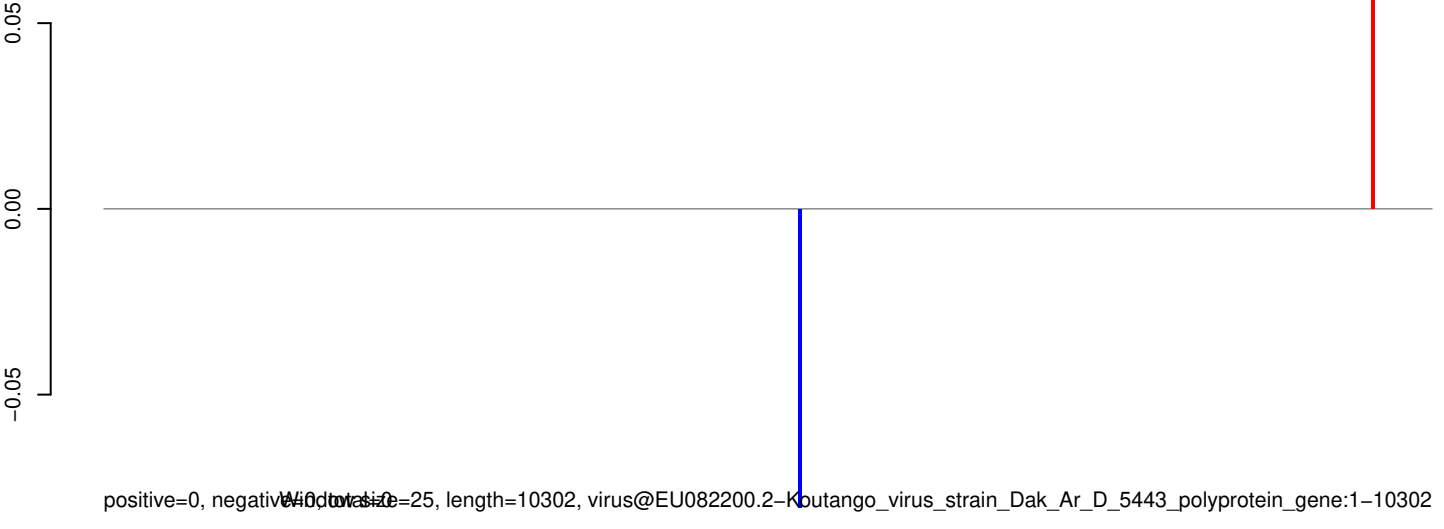
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

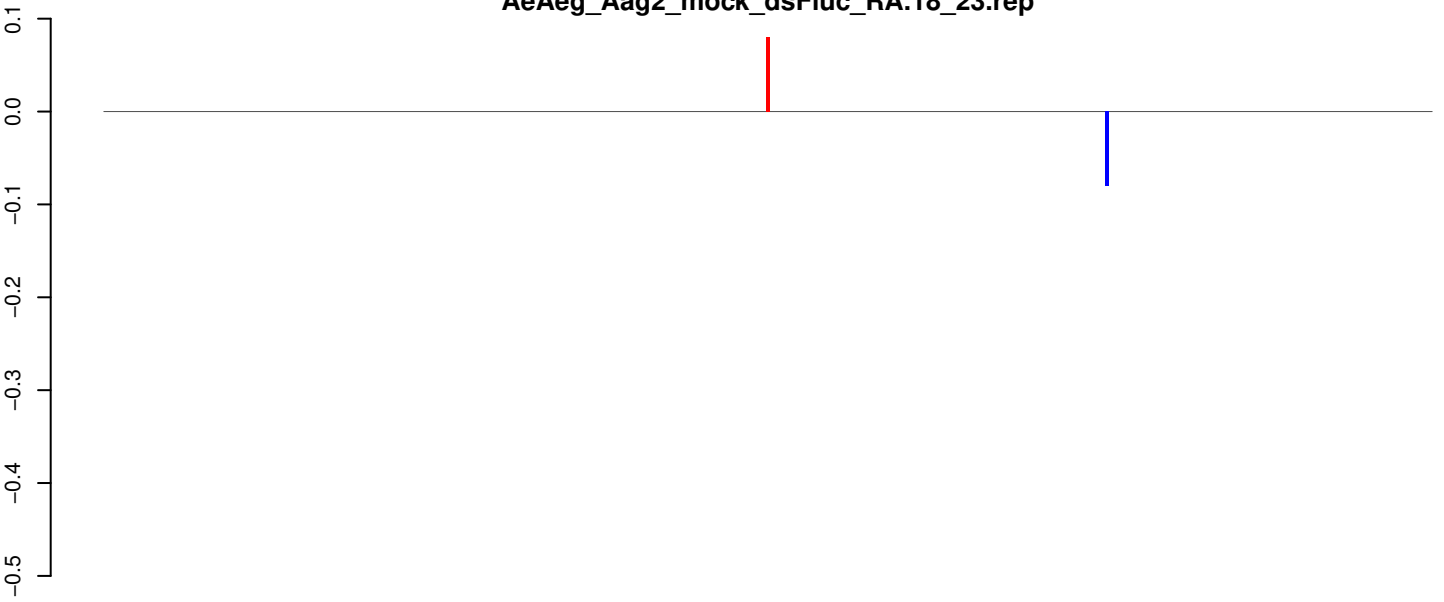


AeAeg_Aag2_mock_dsFluc_RA.rep



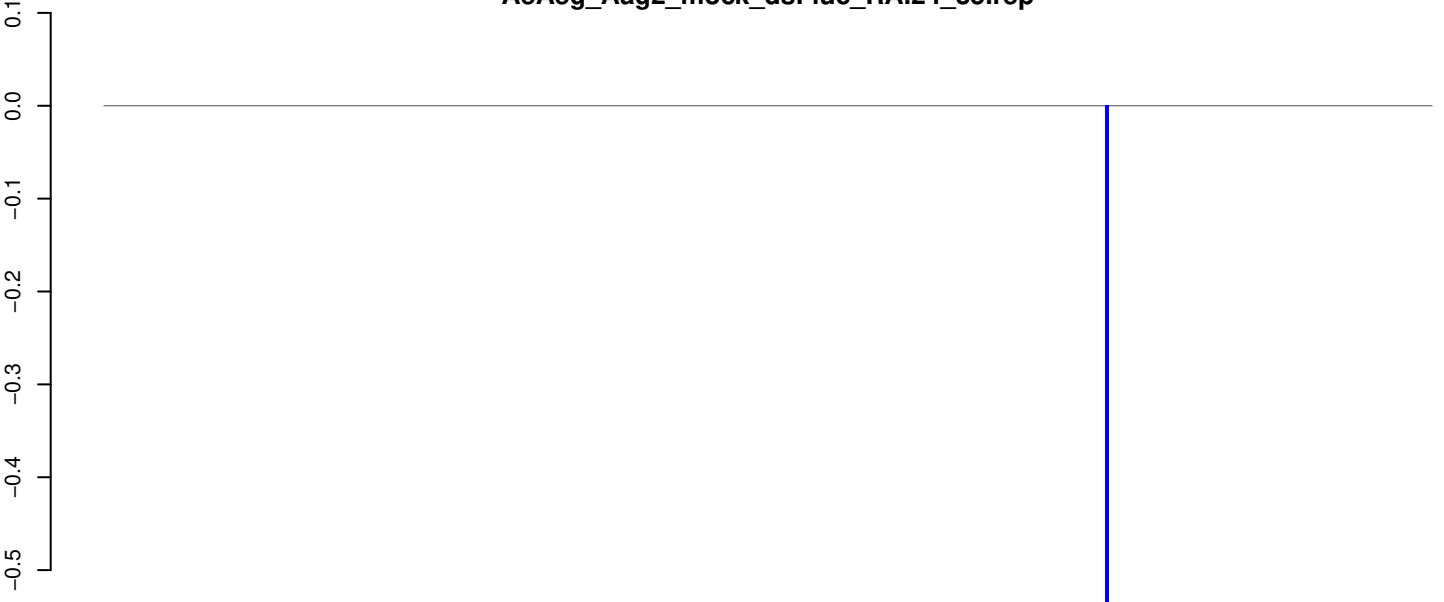
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AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



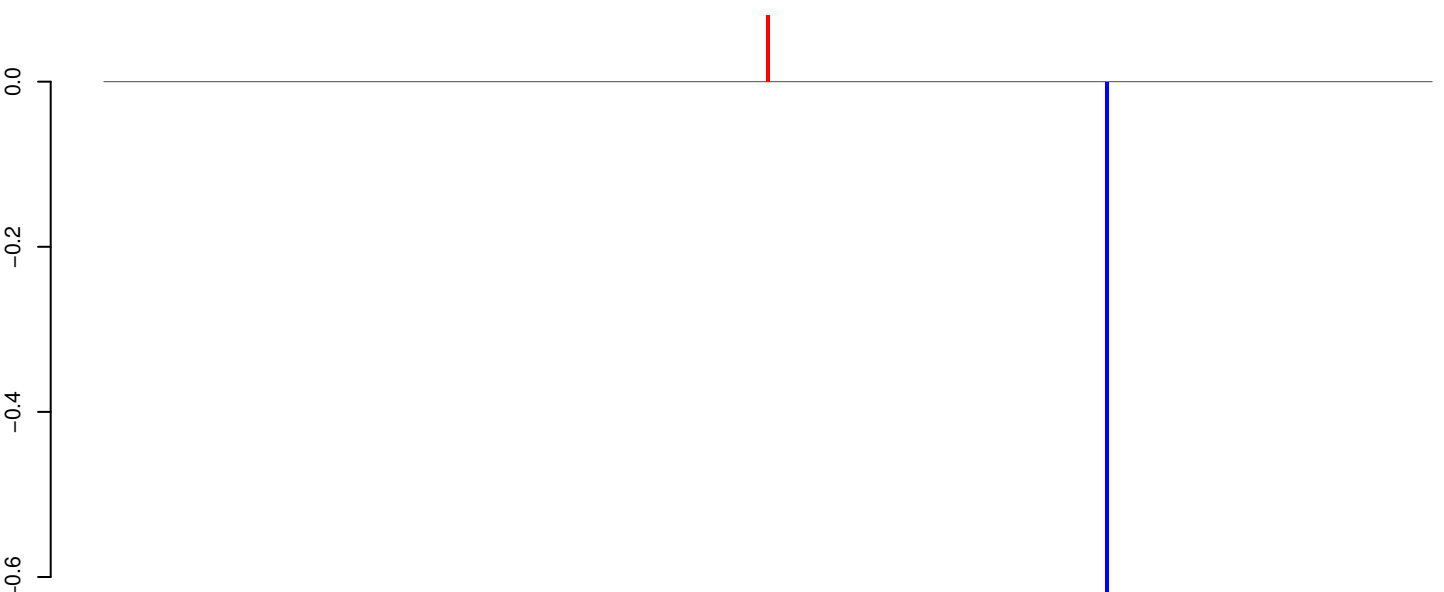
positive=0, negative=0, total=0

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

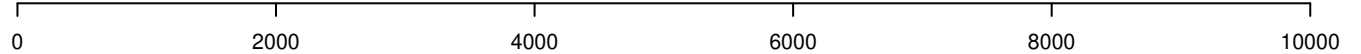


positive=0, negative=1, total=1

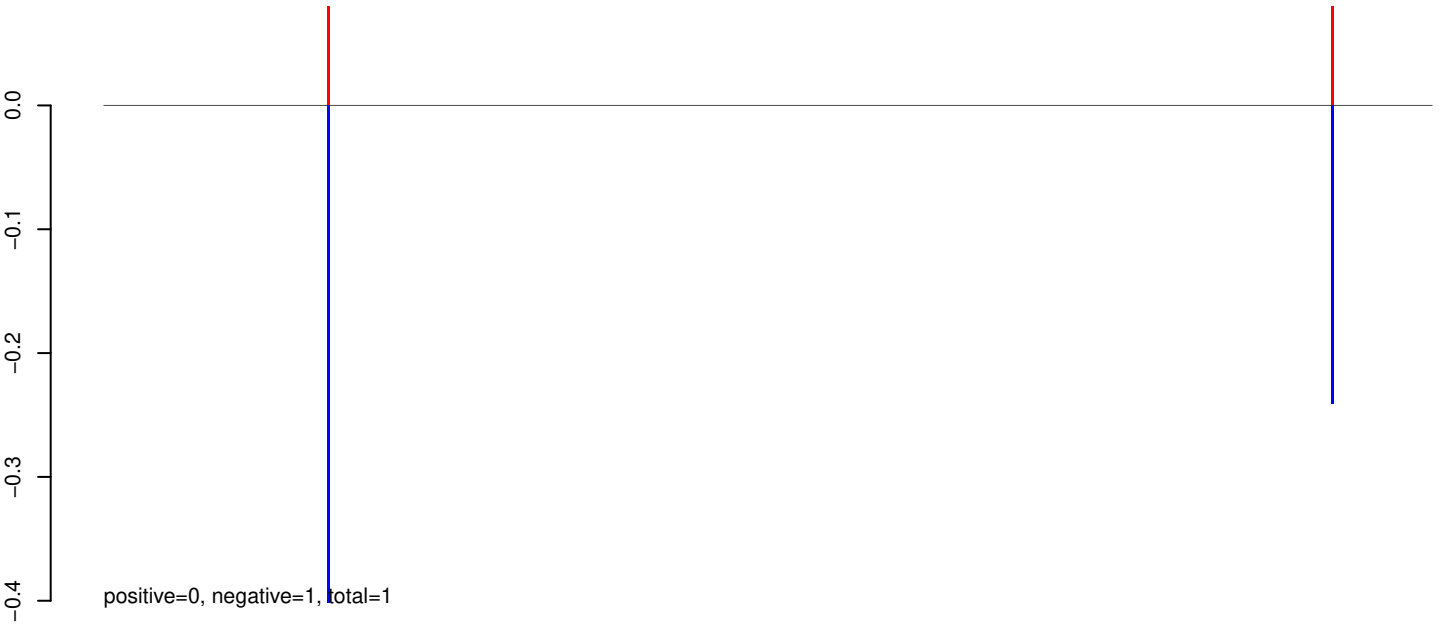
AeAeg_Aag2_mock_dsFluc_RA.rep



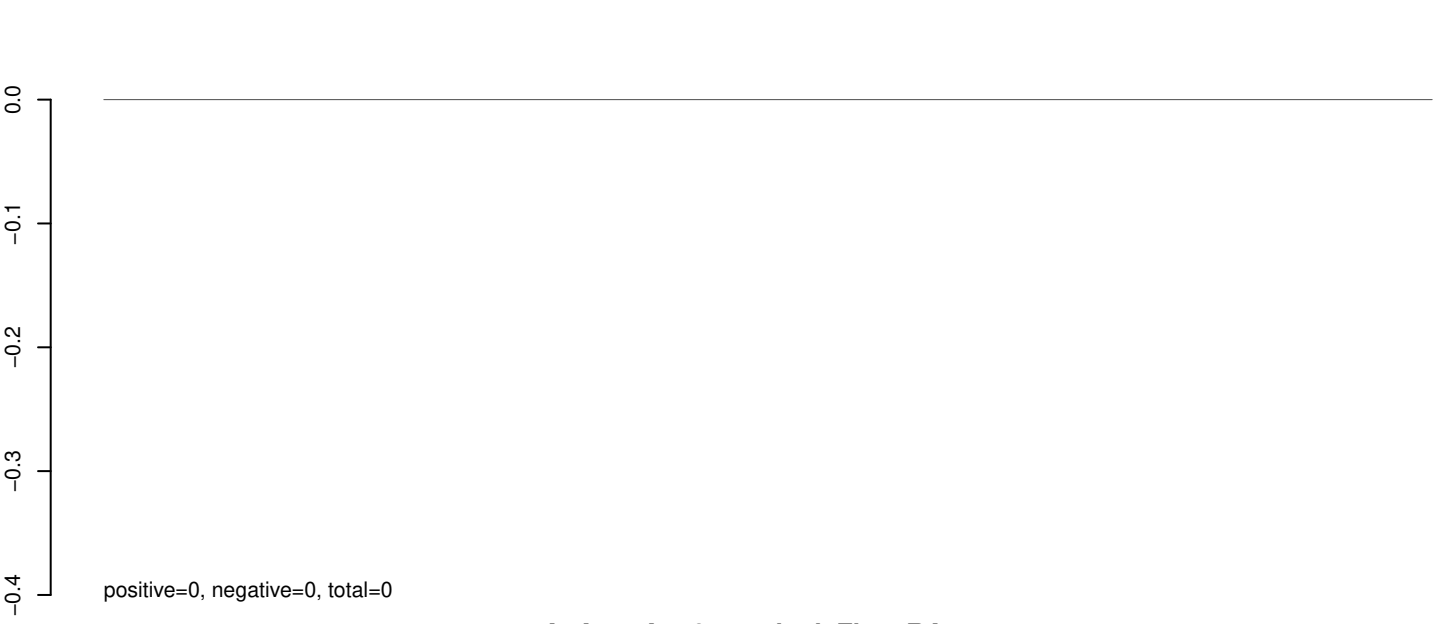
positive=0, negative=1, total=1
Window size=25, length=10254, virus@DQ235149.1-Royal_Farm_virus_from_Afghanistan_polyprotein_gene:1-10254



AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



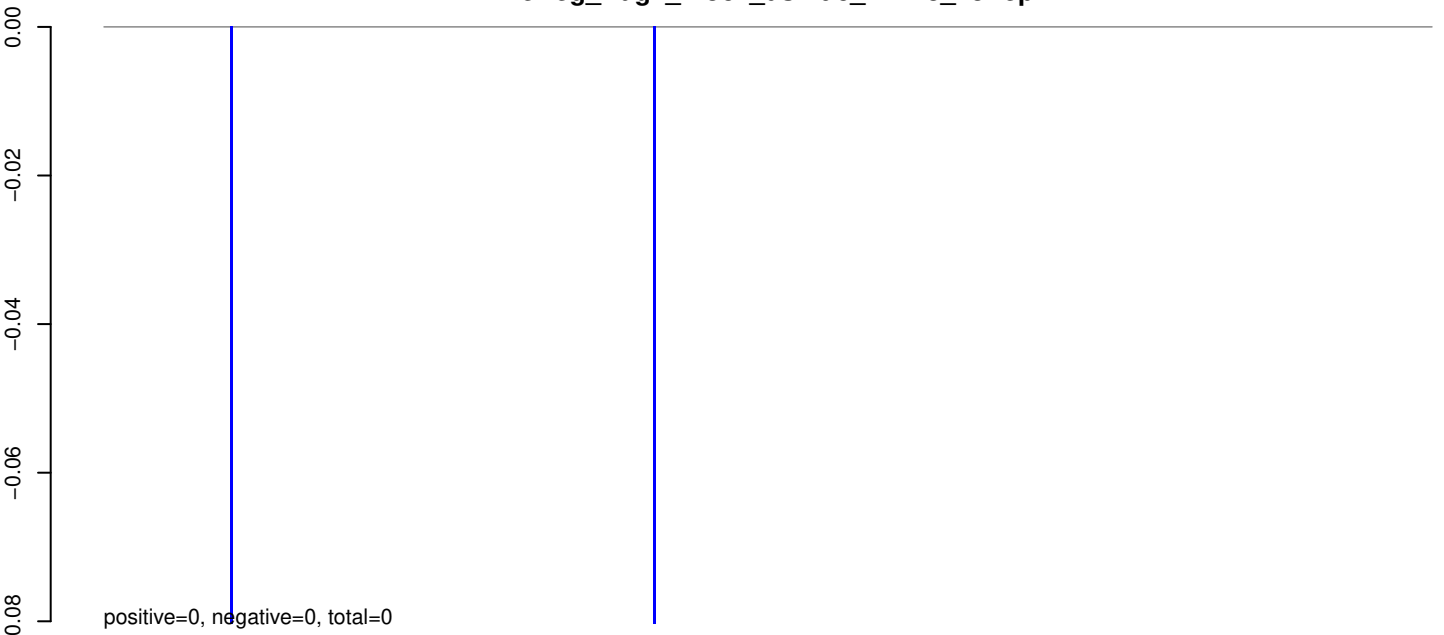
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



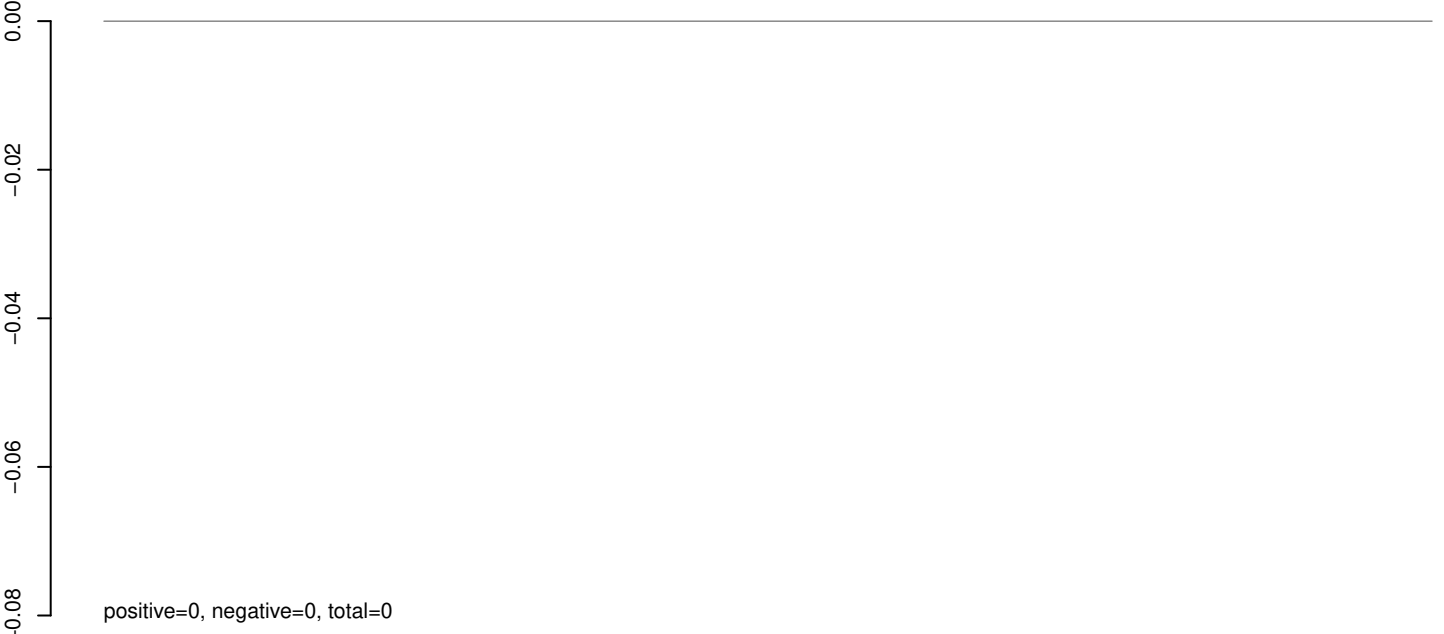
AeAeg_Aag2_mock_dsFluc_RA.rep



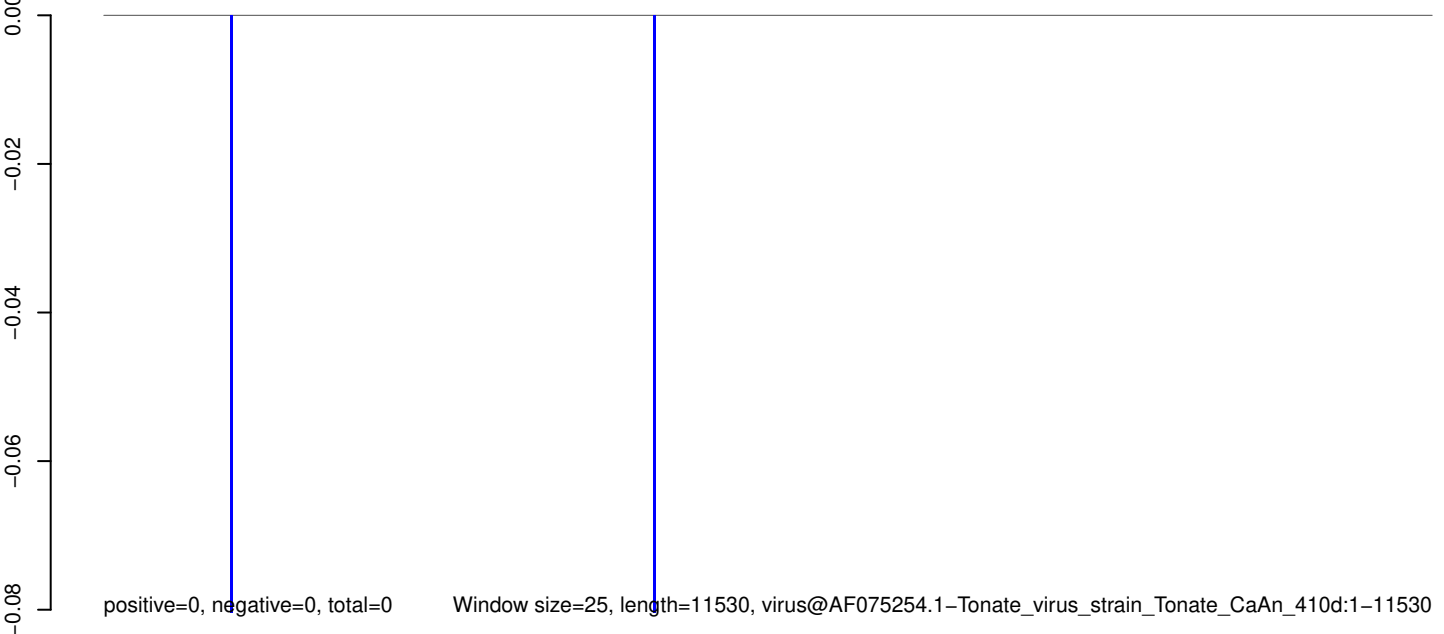
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

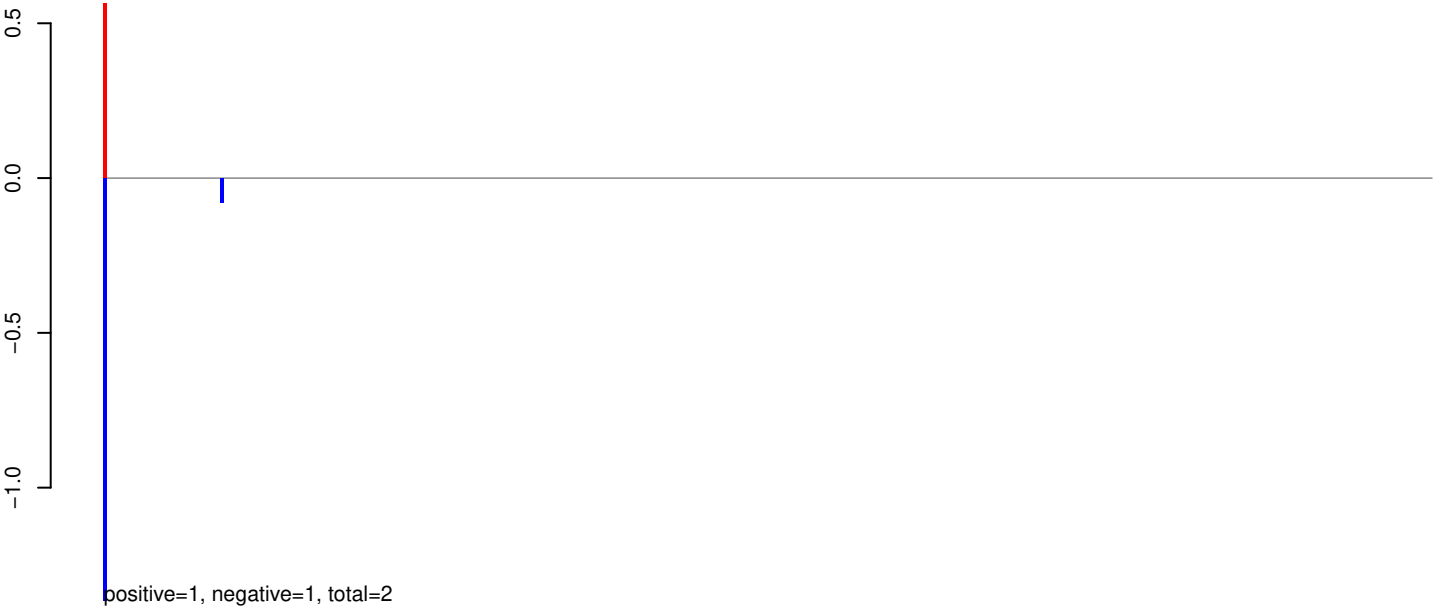


AeAeg_Aag2_mock_dsFluc_RA.rep

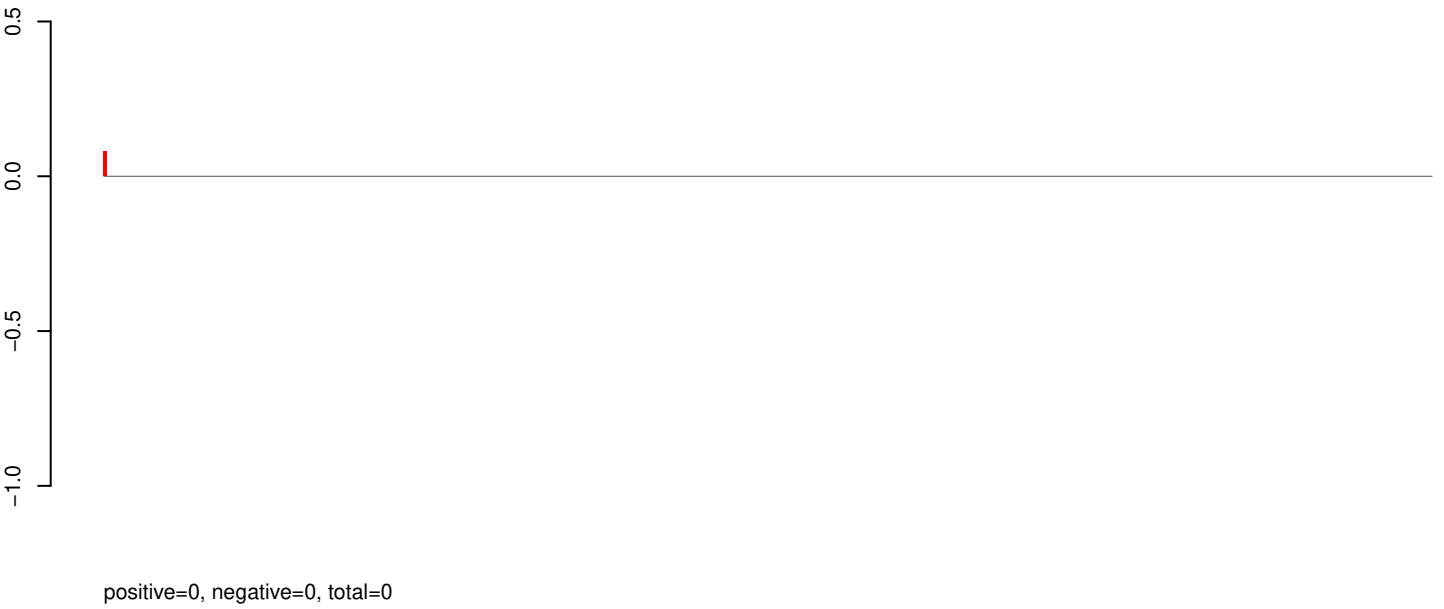


0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



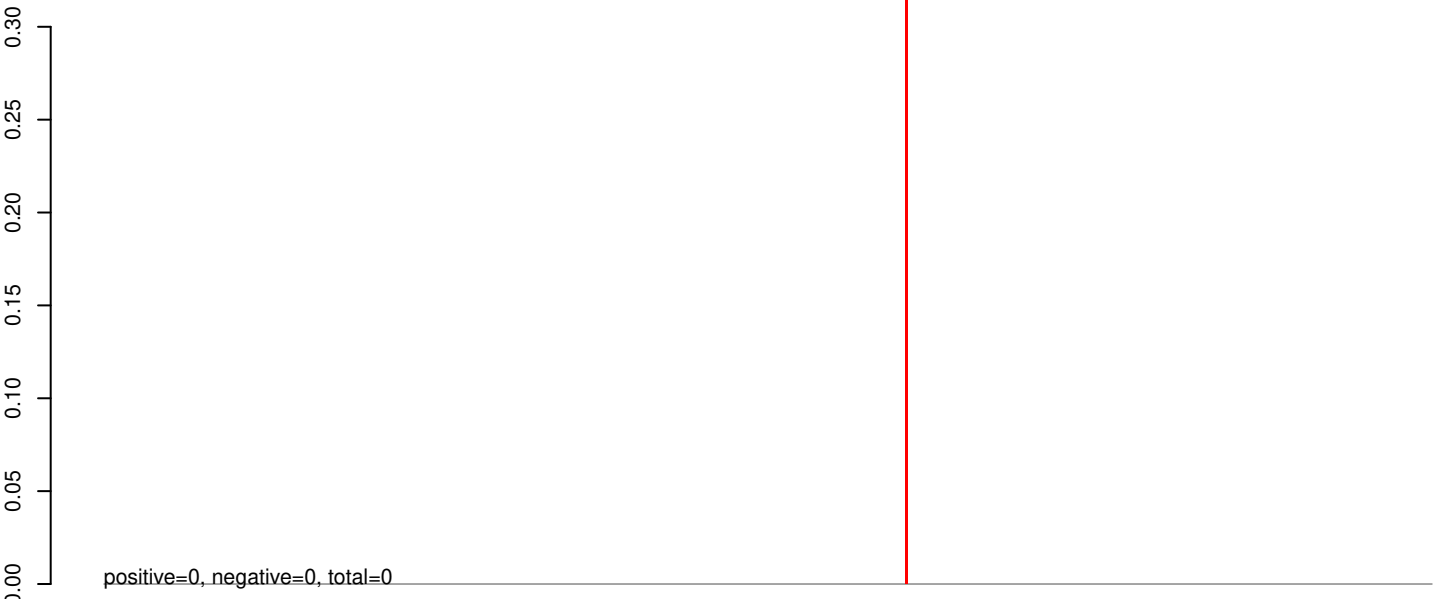
AeAeg_Aag2_mock_dsFluc_RA.rep



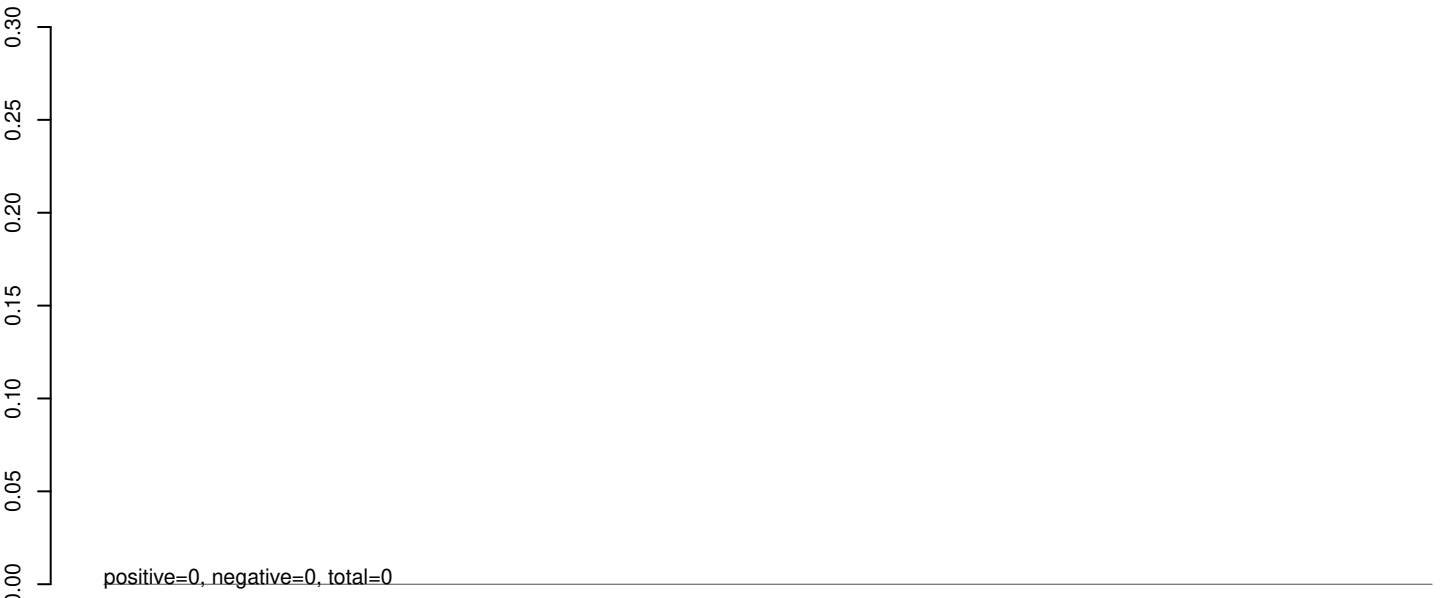
Window size=25, length=10837, virus@AB262759.2-Culex_flavivirus_genomic_RNA_strain_Tokyo:1-10837

0 2000 4000 6000 8000 10000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



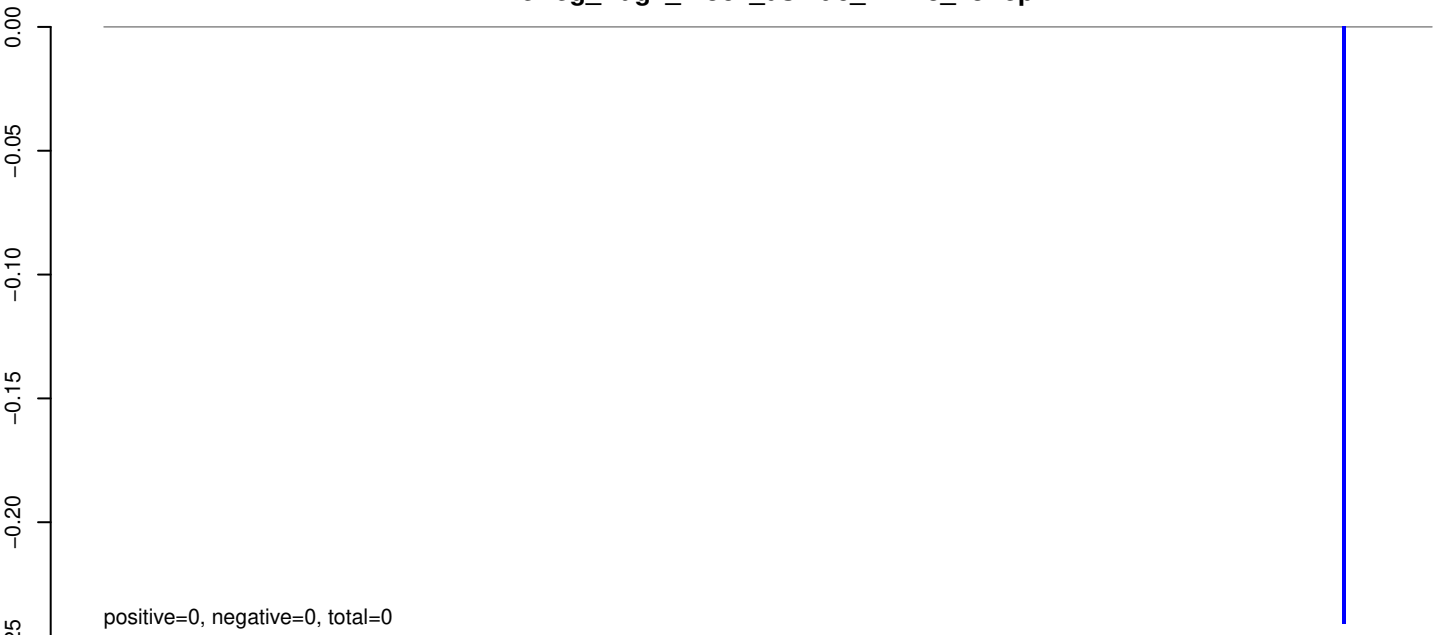
AeAeg_Aag2_mock_dsFluc_RA.rep



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

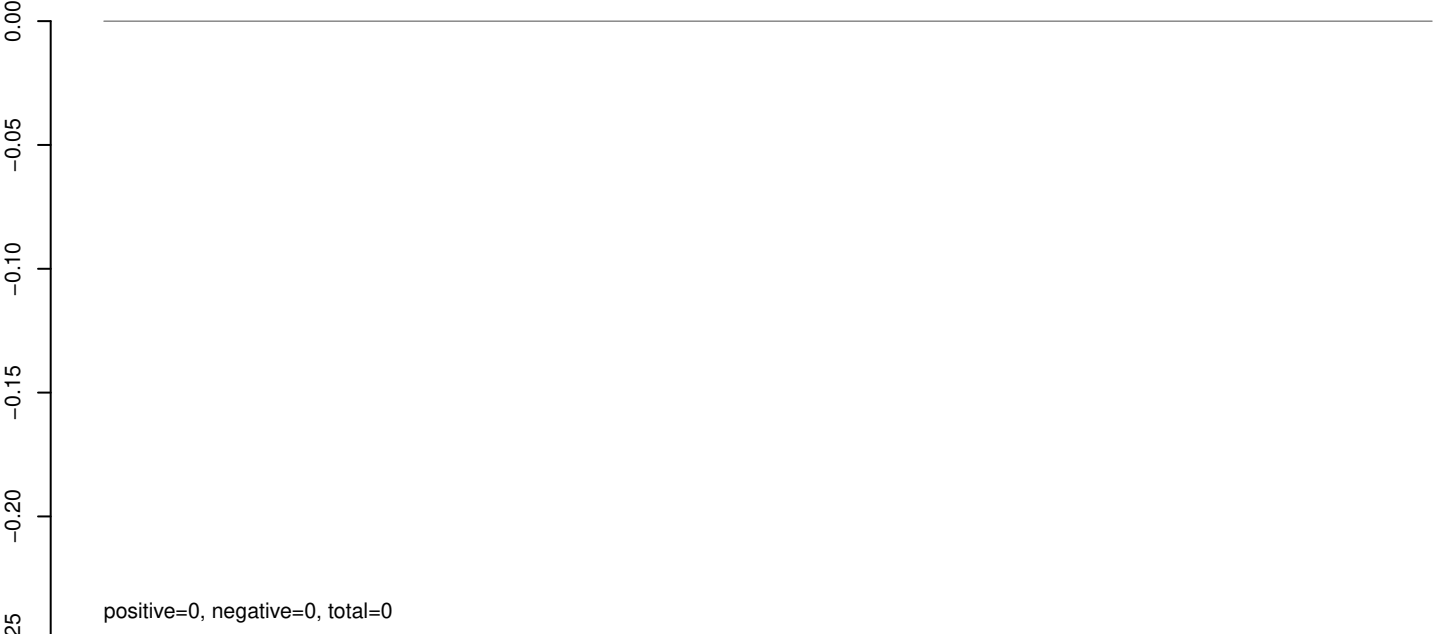
0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



positive=0, negative=0, total=0

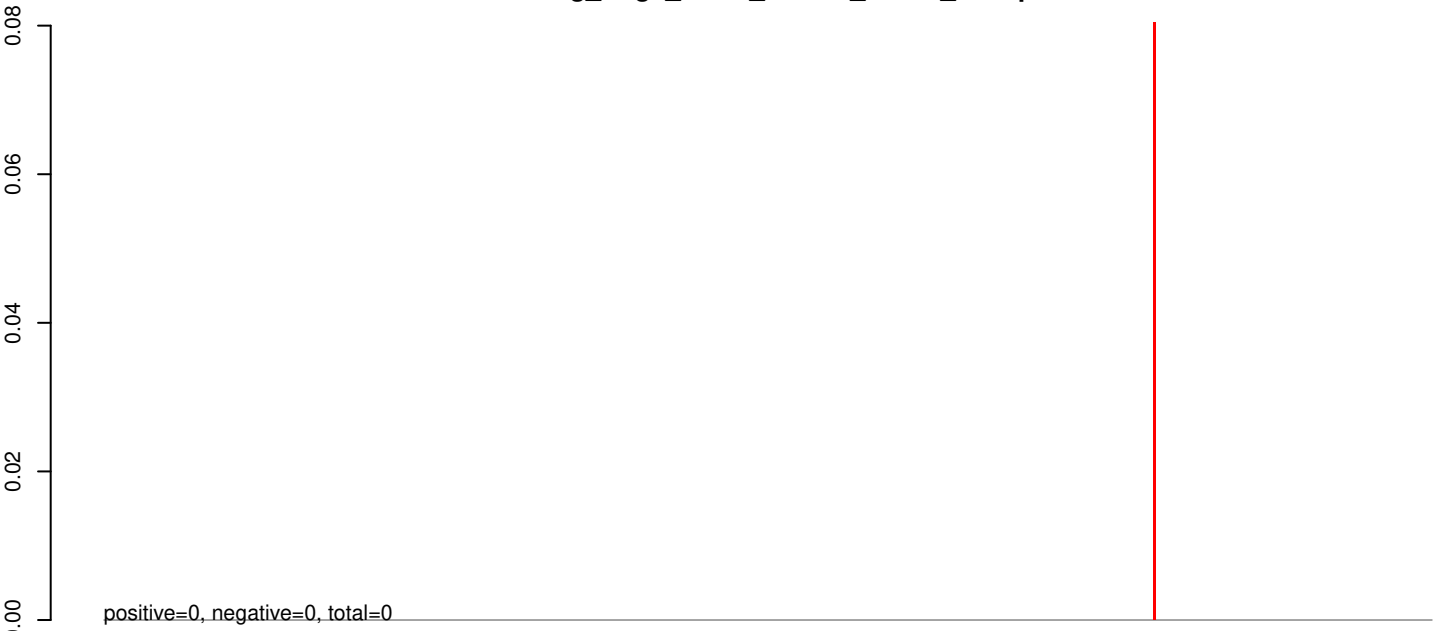
AeAeg_Aag2_mock_dsFluc_RA.rep



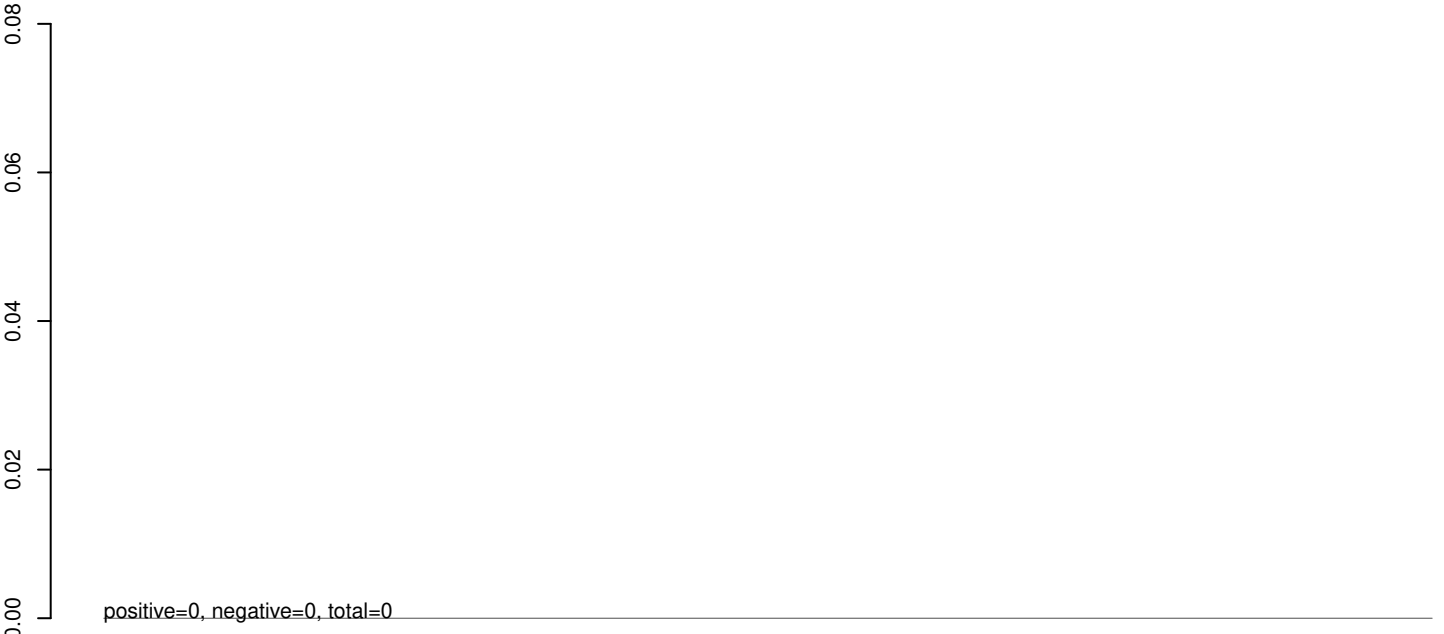
positive=0, negative=0, total=0 window size=25, length=9176, virus@NC_035129.1_Culex_negev-like_virus_3_strain_mos172X44875:1-9176

0 2000 4000 6000 8000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

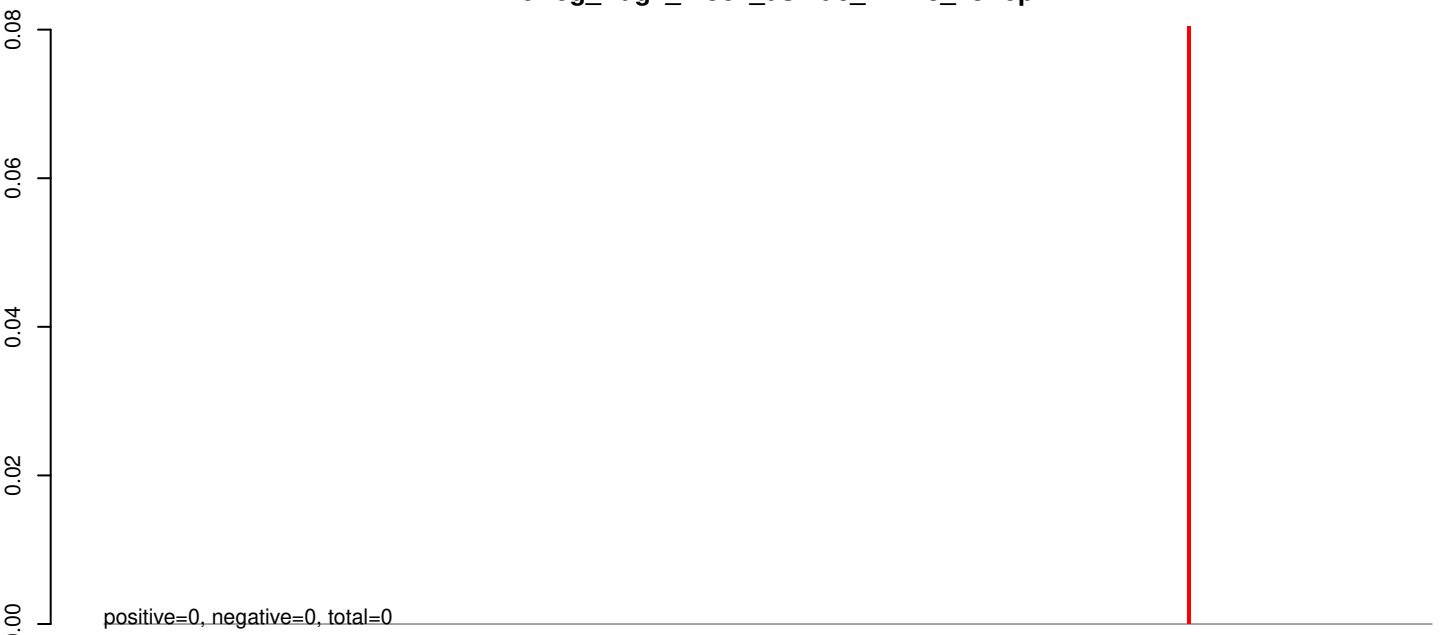


AeAeg_Aag2_mock_dsFluc_RA.rep

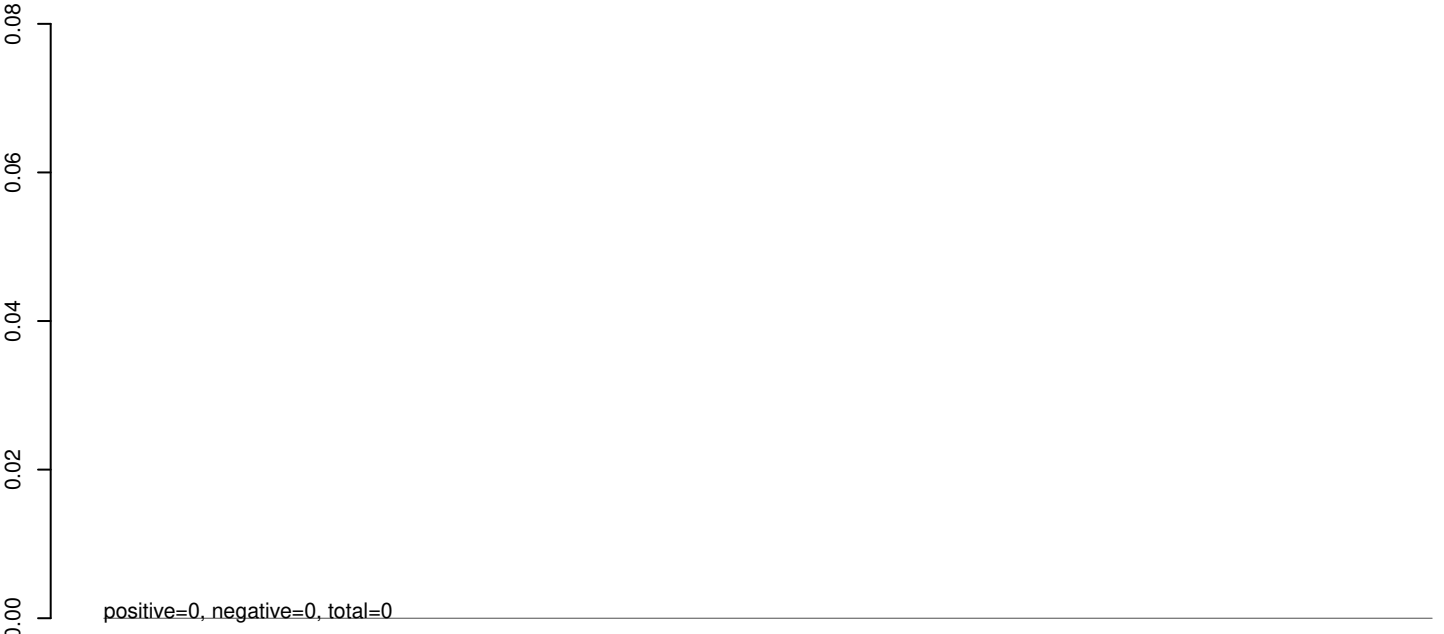


0 2000 4000 6000 8000 10000 12000

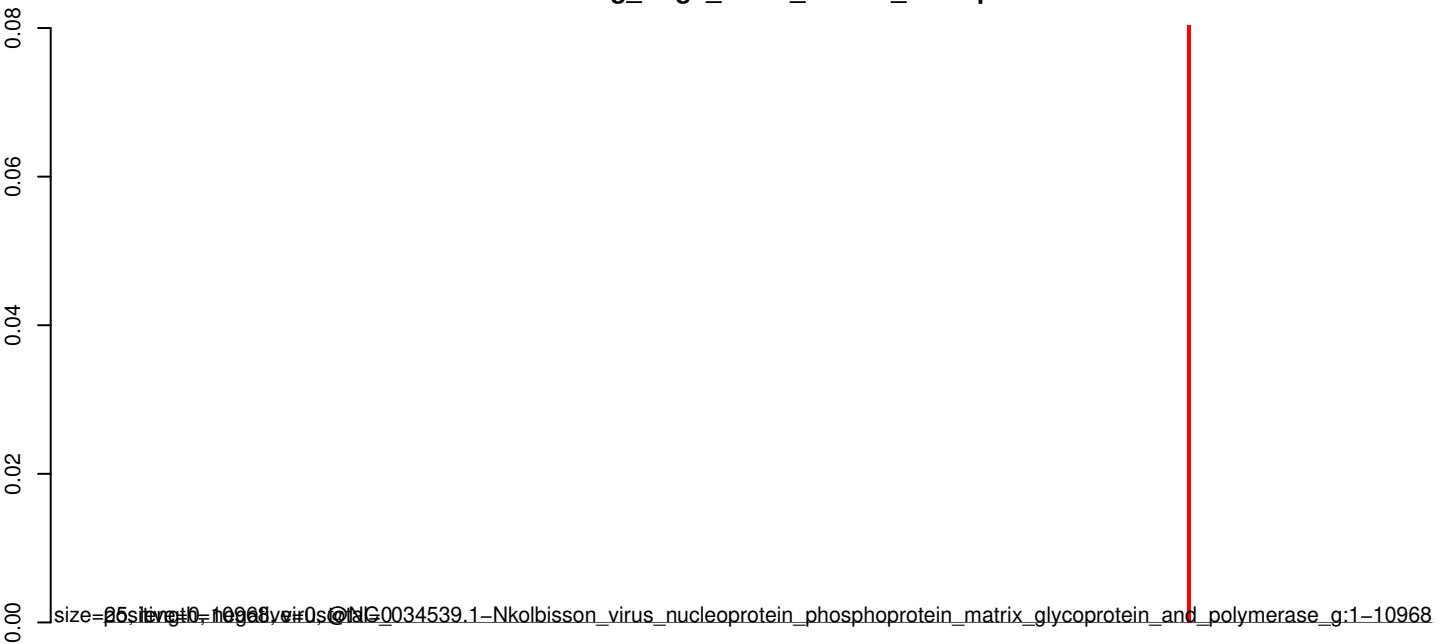
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

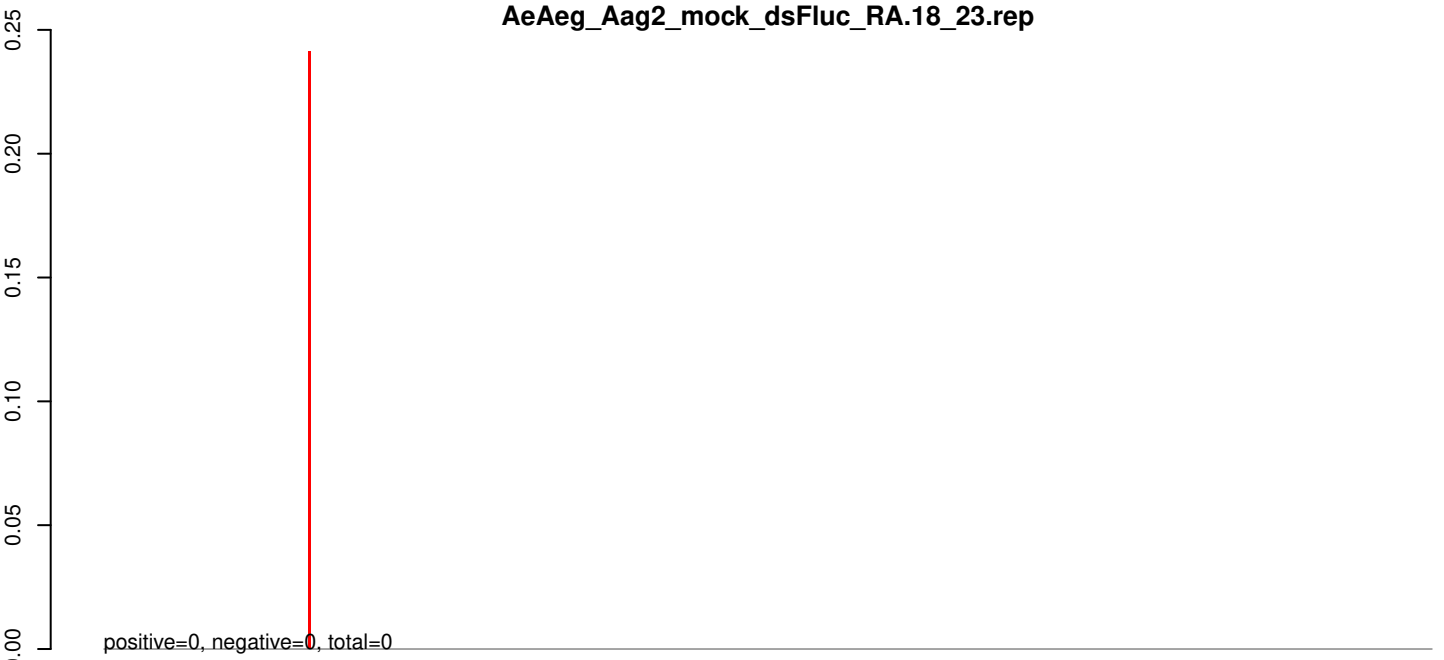


AeAeg_Aag2_mock_dsFluc_RA.rep

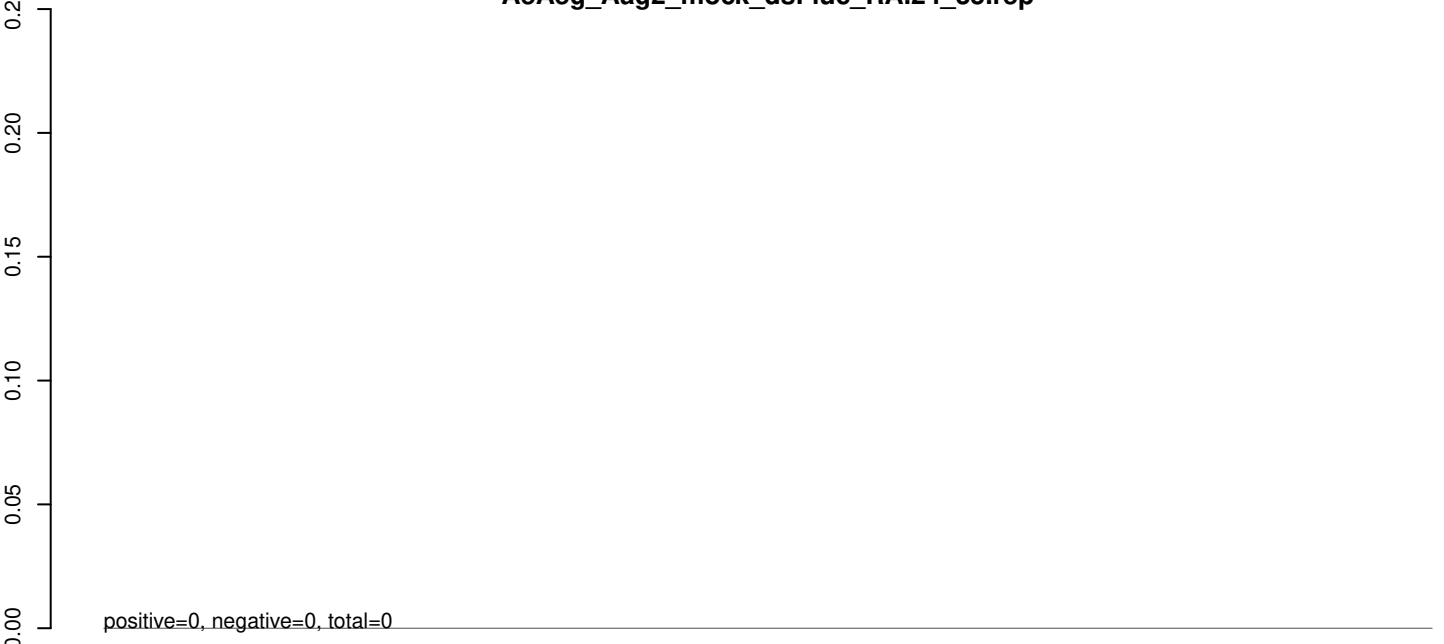


0 2000 4000 6000 8000 10000

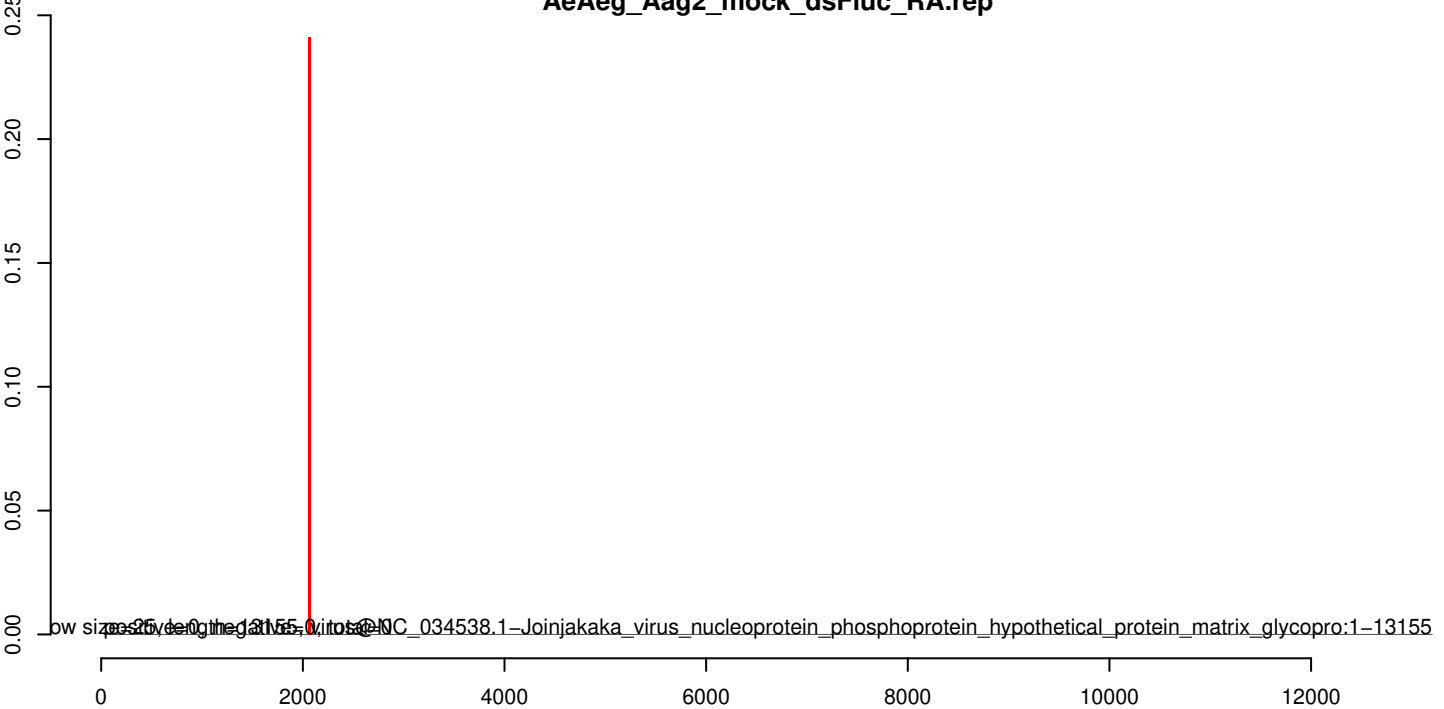
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



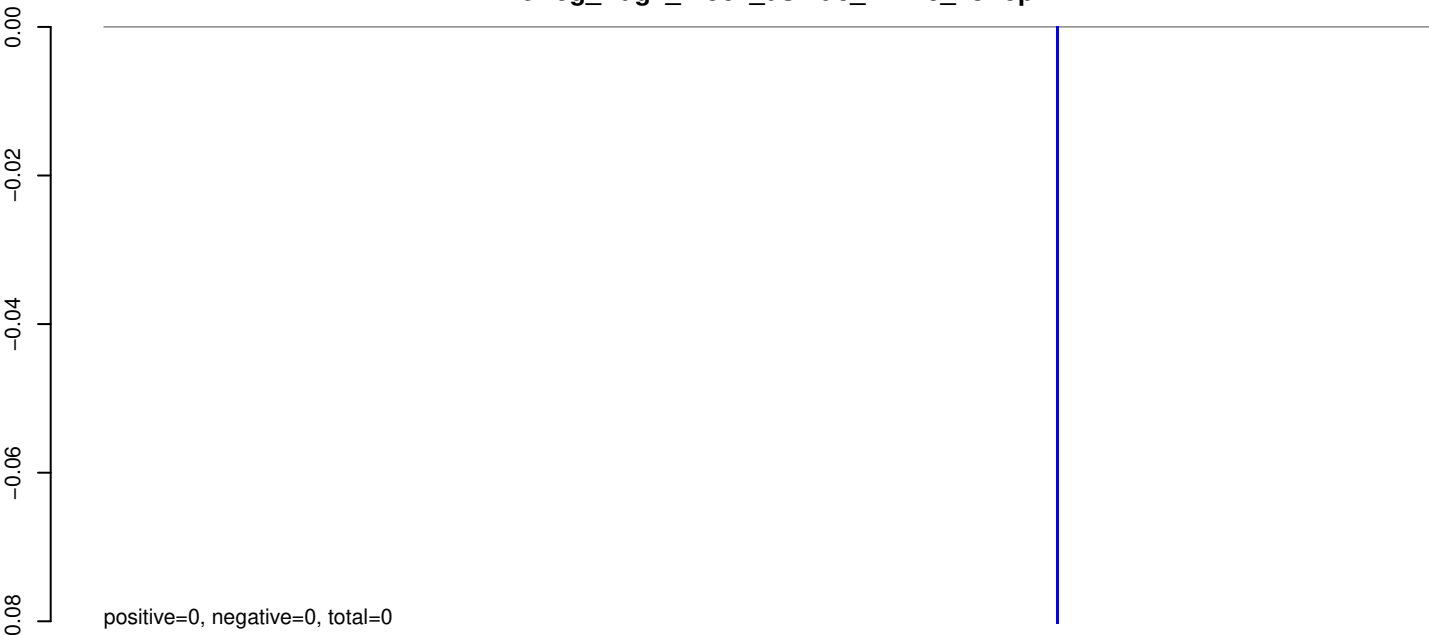
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



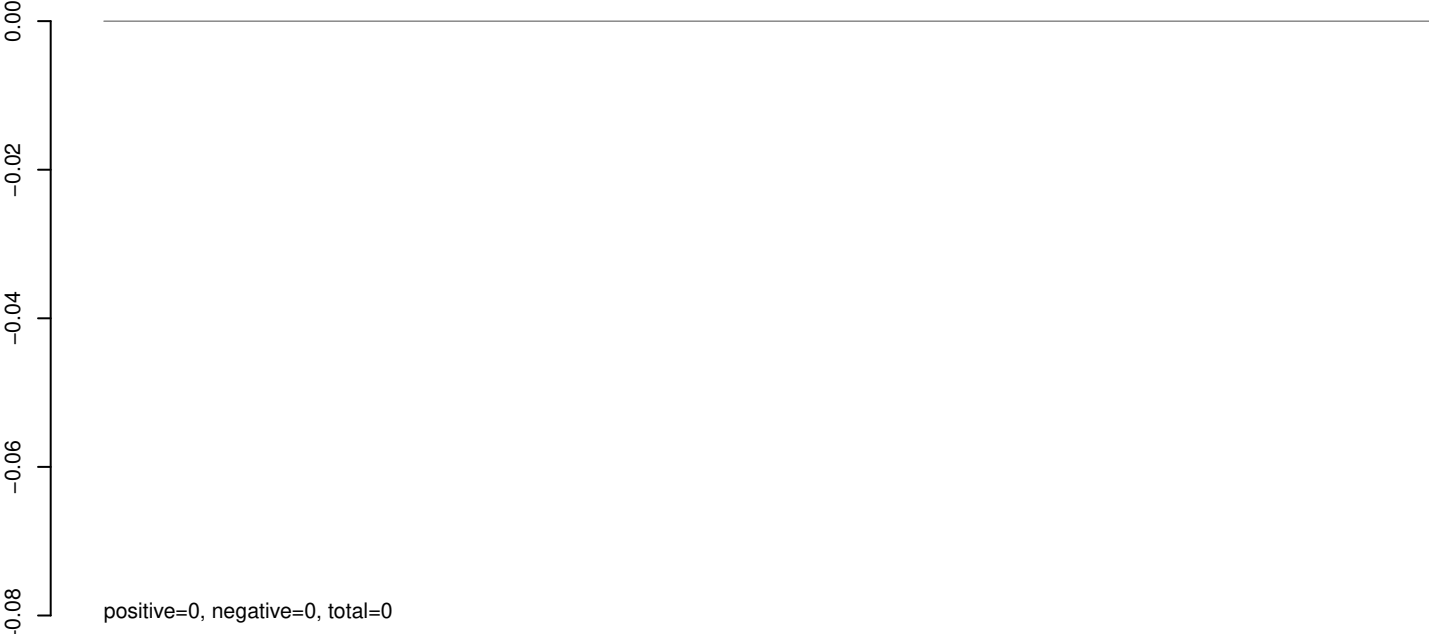
AeAeg_Aag2_mock_dsFluc_RA.rep



AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



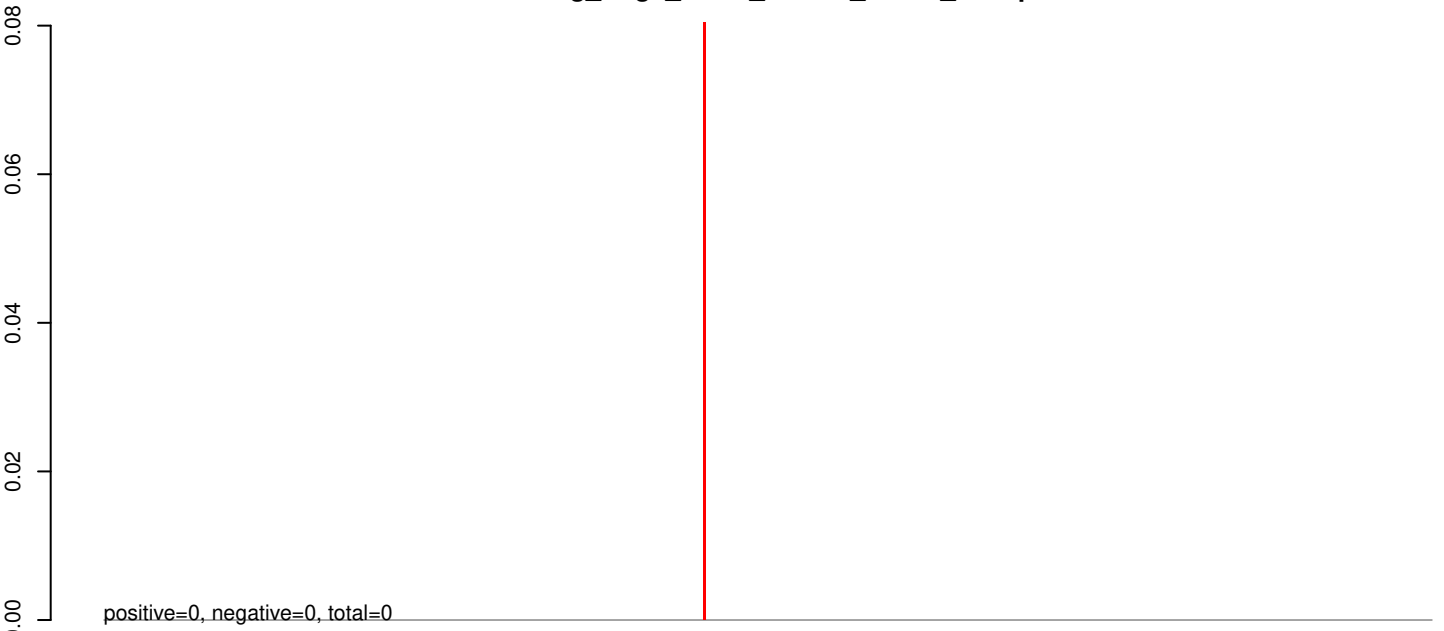
AeAeg_Aag2_mock_dsFluc_RA.rep



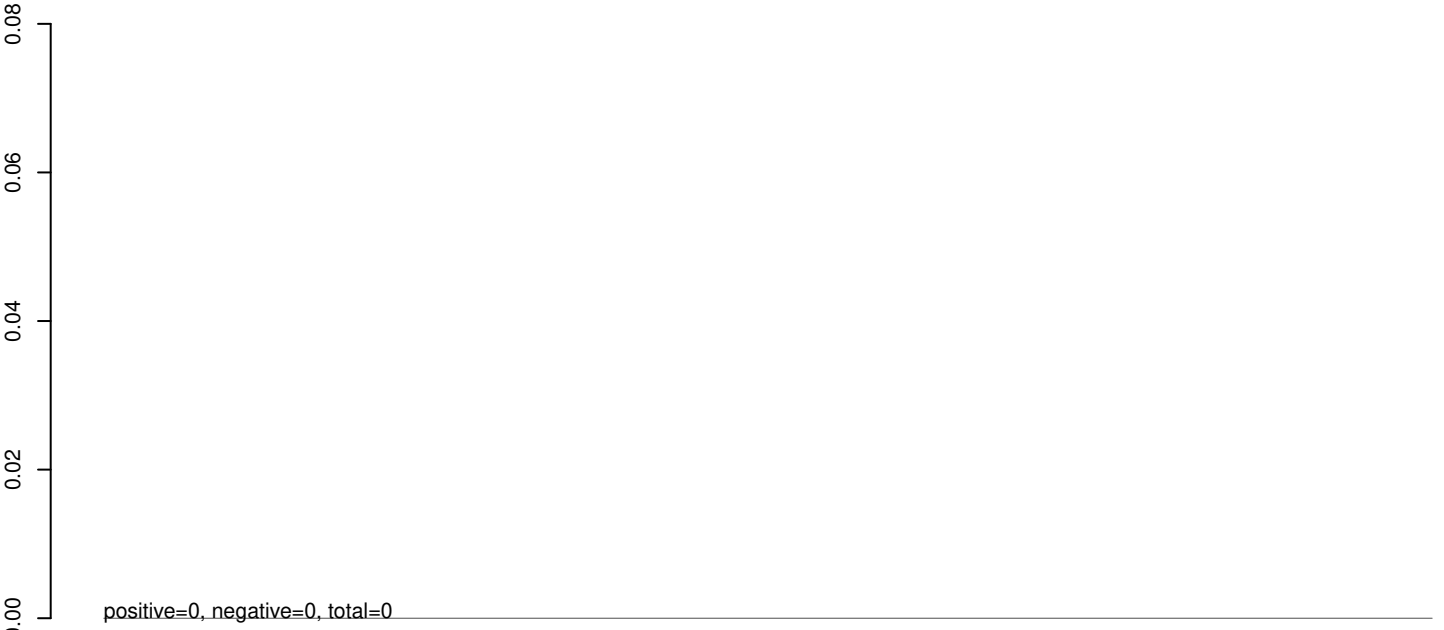
LOC0034534.1-Rochambeau_virus_nucleoprotein_phosphoprotein_matrix_hypothetical_proteins_glycopr:1-13593

0 2000 4000 6000 8000 10000 12000 14000

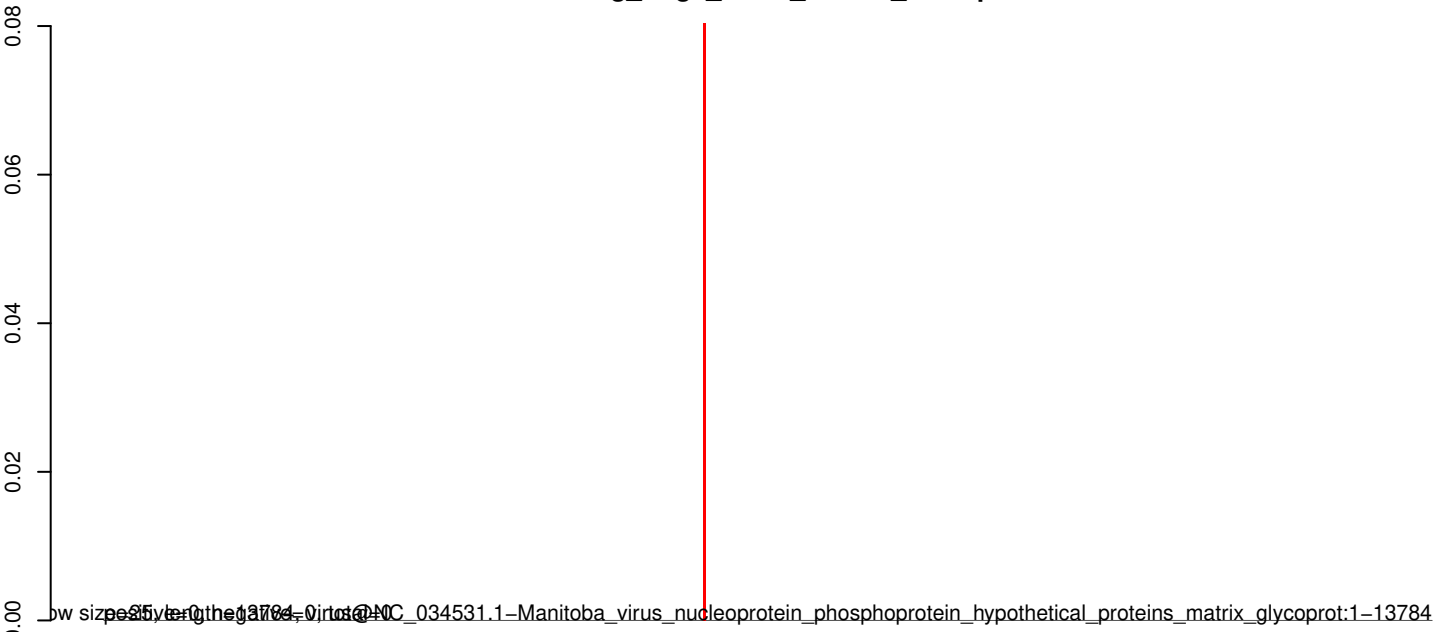
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep

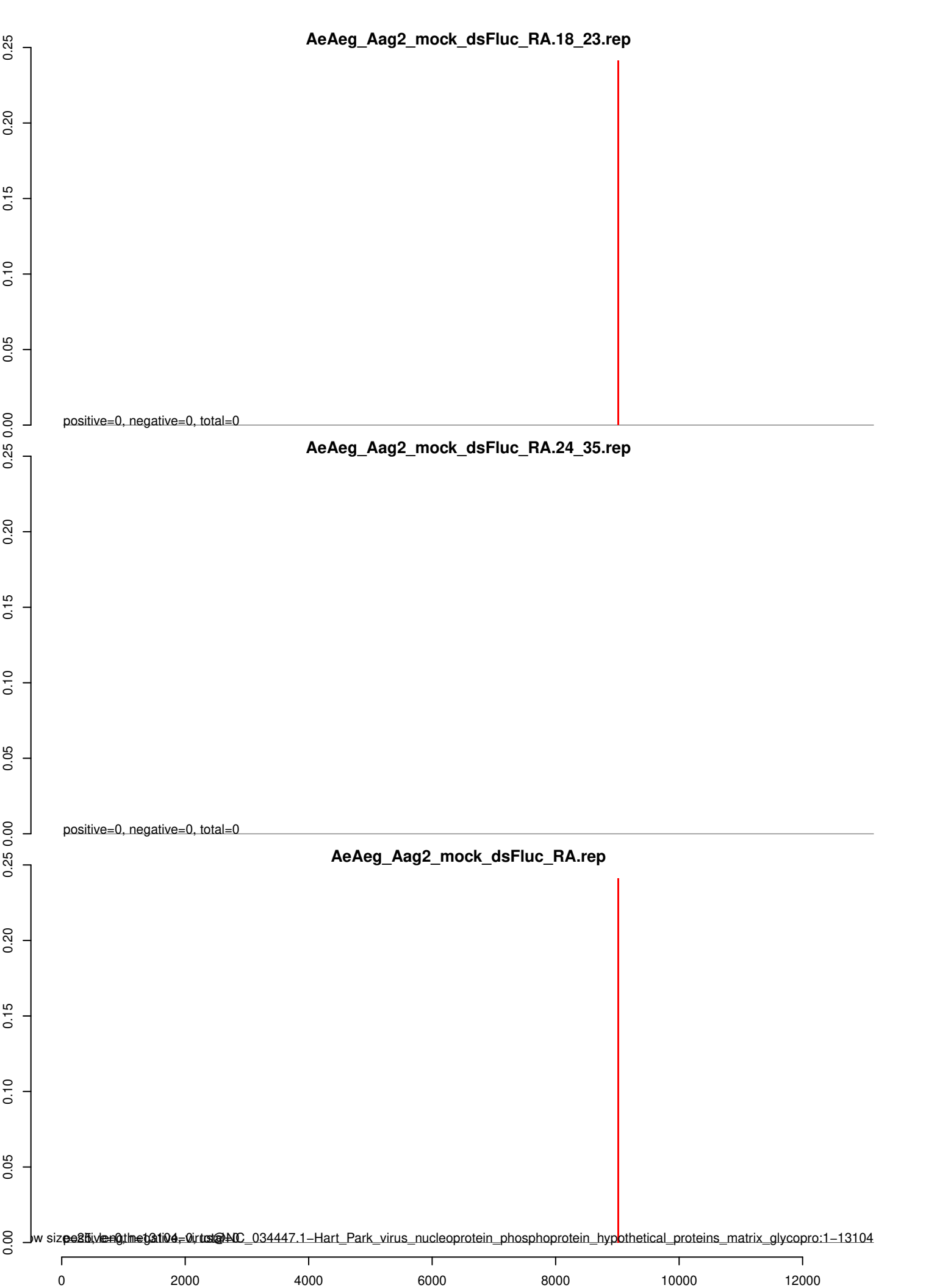


AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

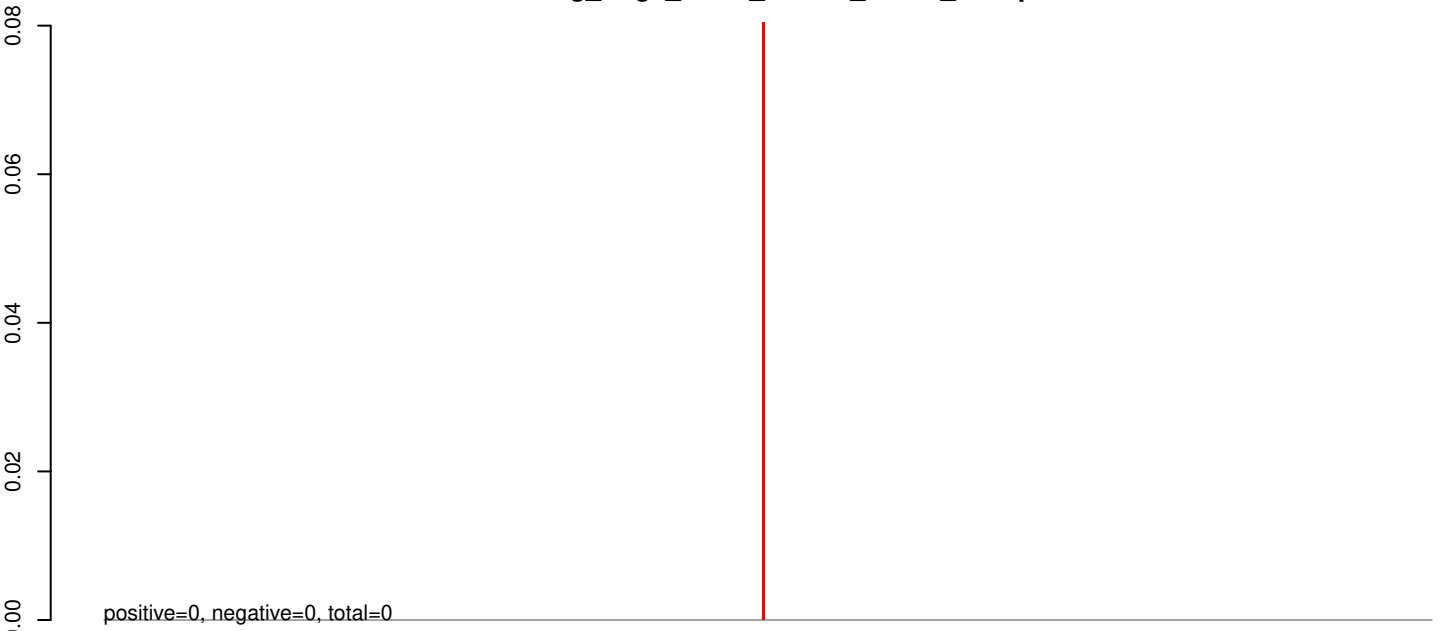


AeAeg_Aag2_mock_dsFluc_RA.rep

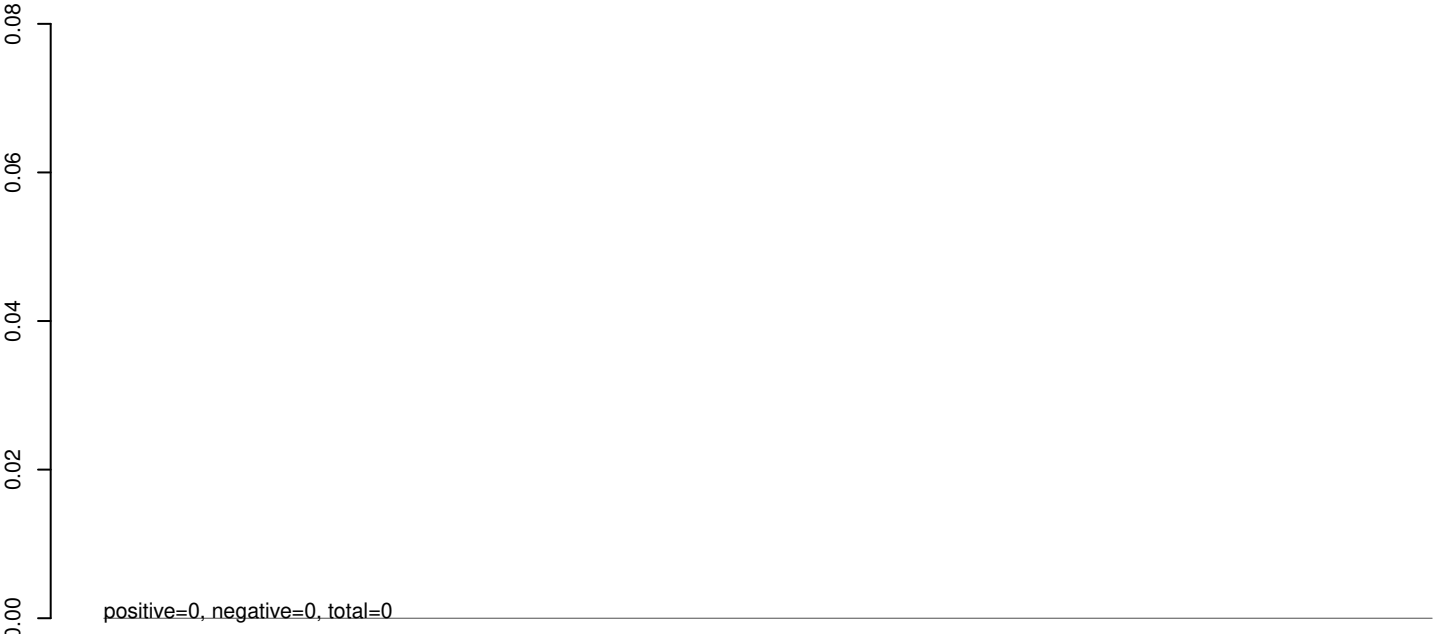




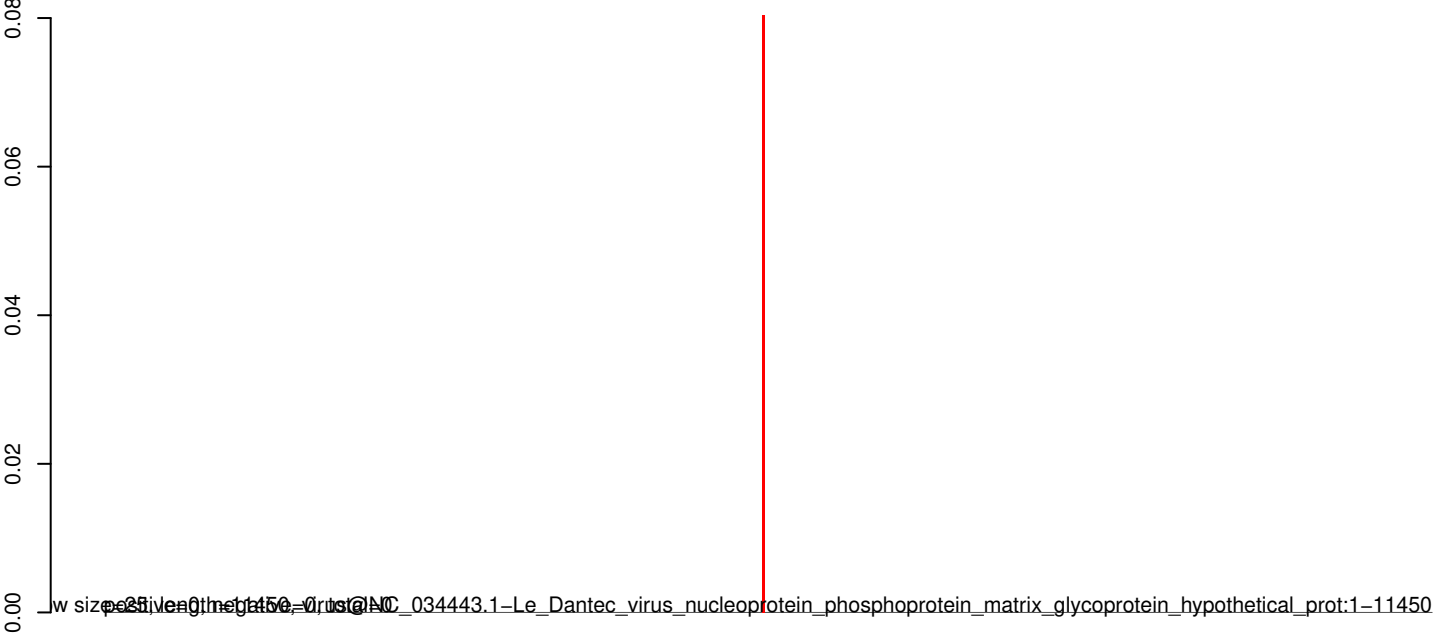
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

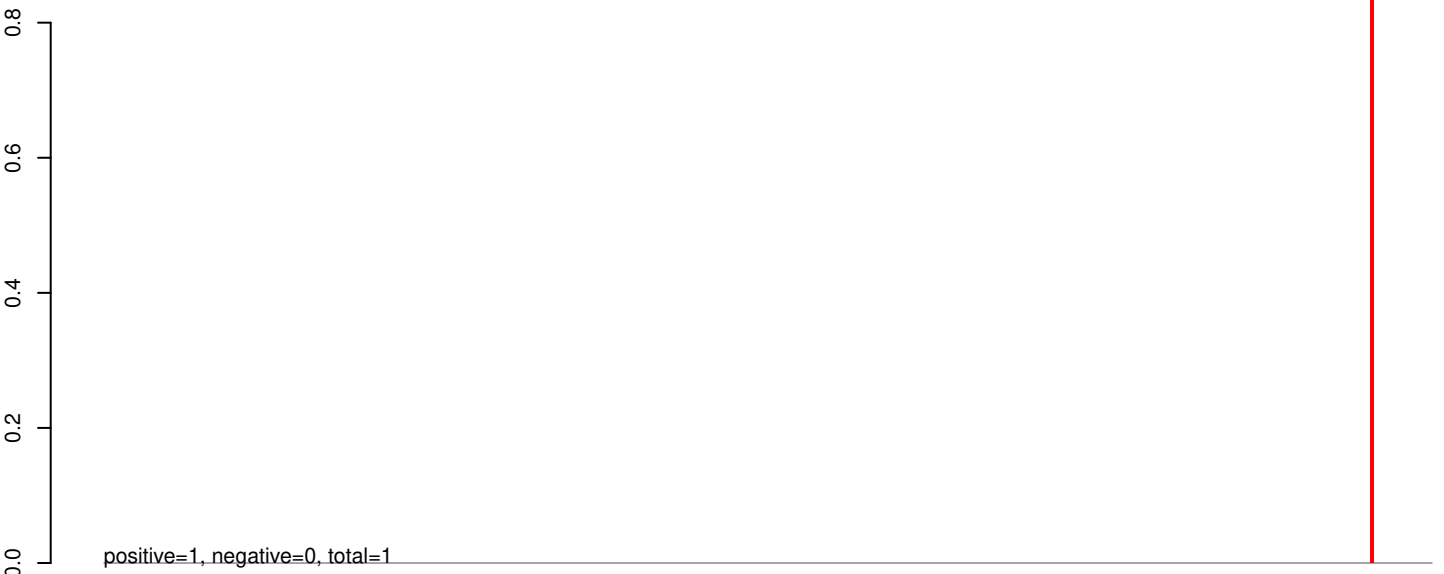


AeAeg_Aag2_mock_dsFluc_RA.rep

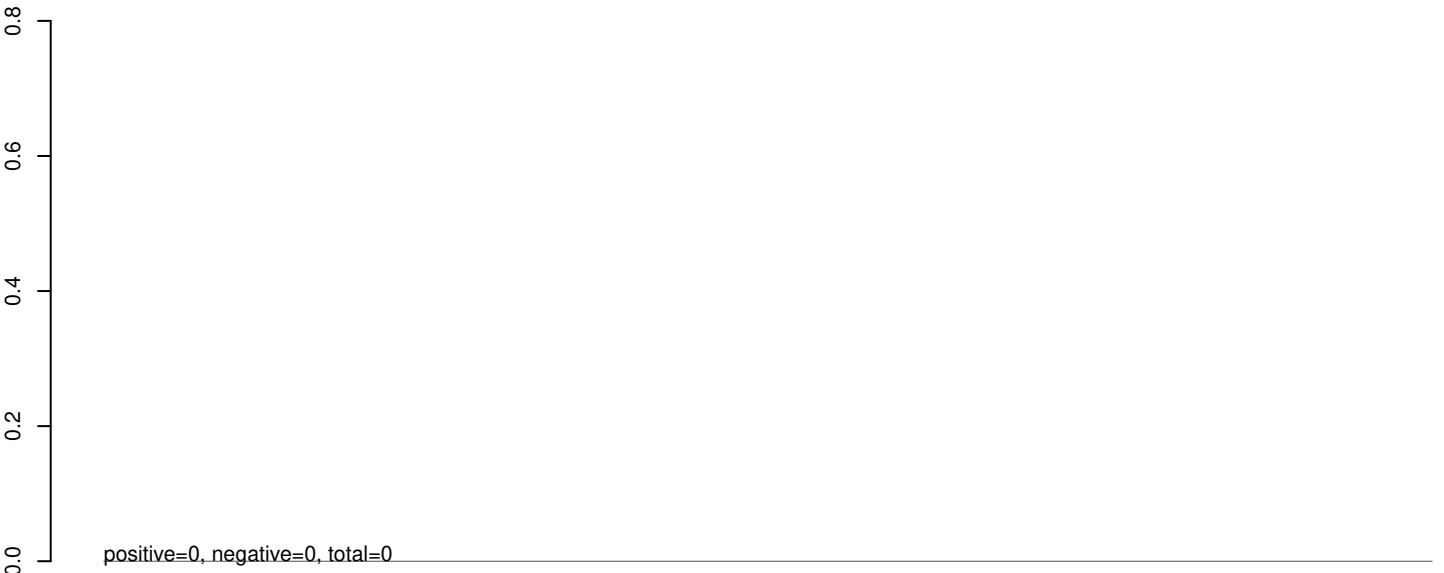


NC_034443.1-Le_Dantec_virus_nucleoprotein_phosphoprotein_matrix_glycoprotein_hypothetical_prot:1-11450

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

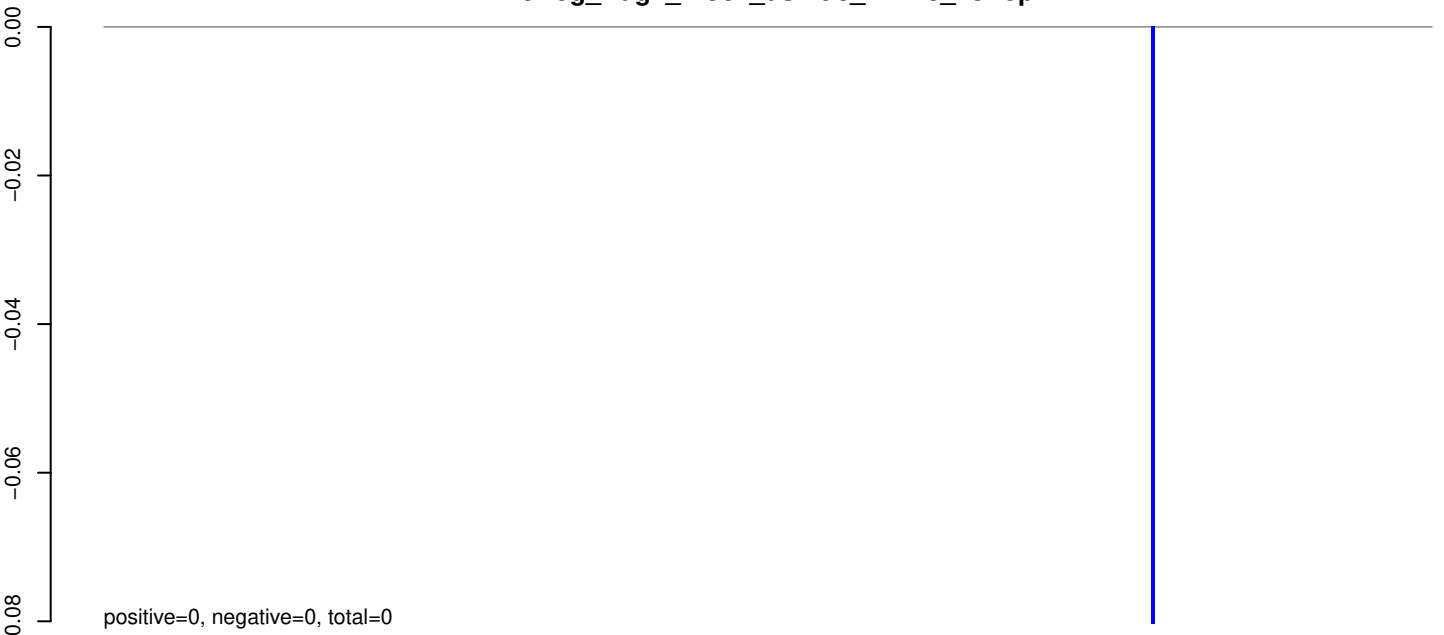


AeAeg_Aag2_mock_dsFluc_RA.rep

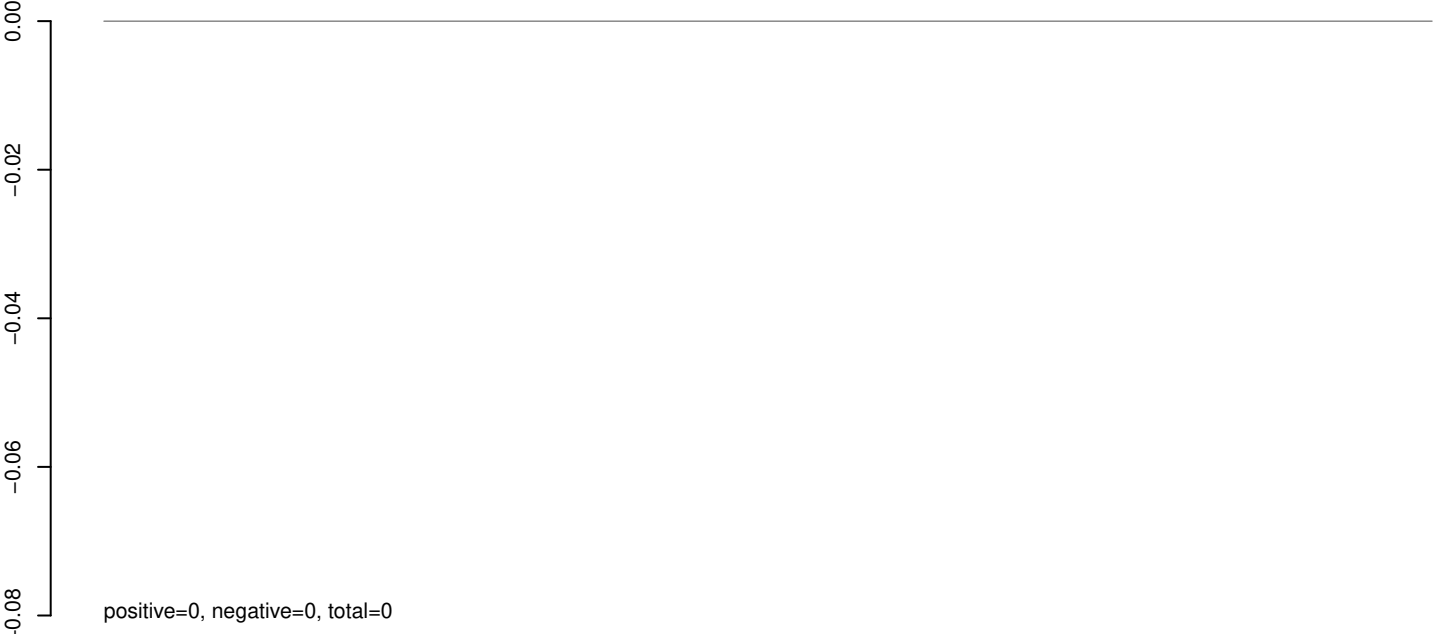


0 2000 4000 6000 8000 10000
window size=25, length=10251, virus@NC_033723.1-Gadgets_Gully_virus_from_Australia_polyprotein_gene:1-10251

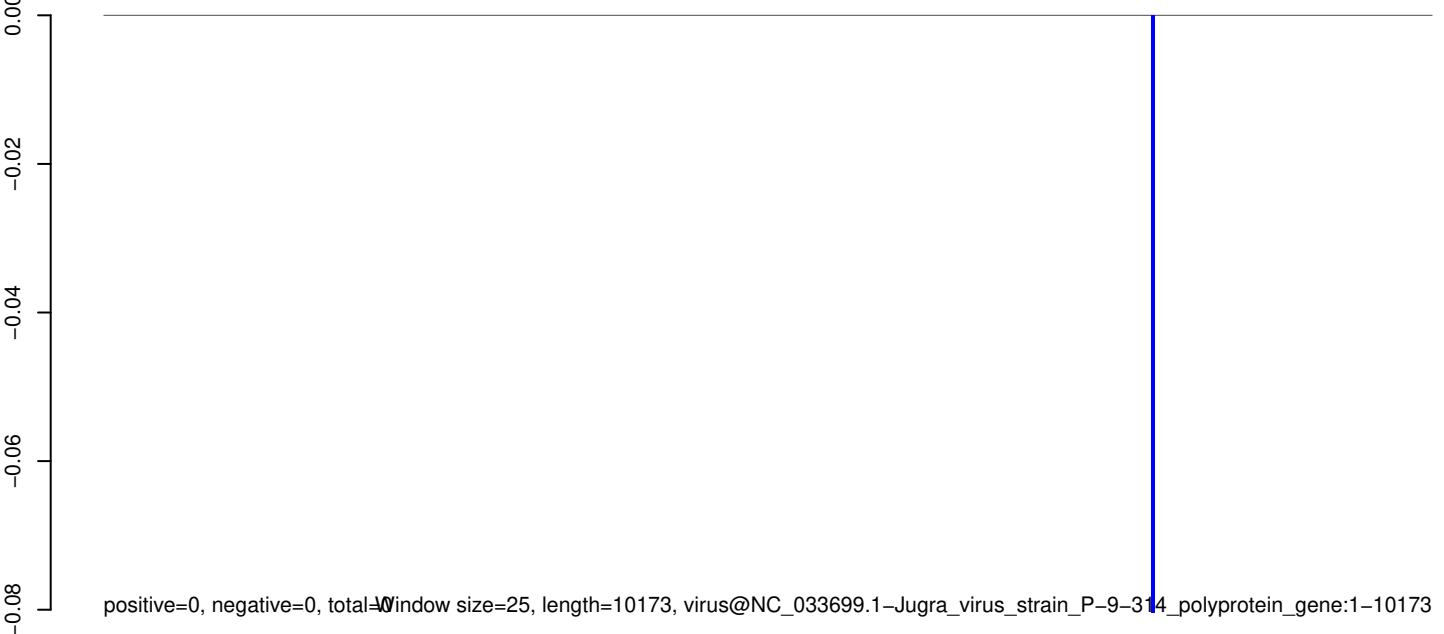
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

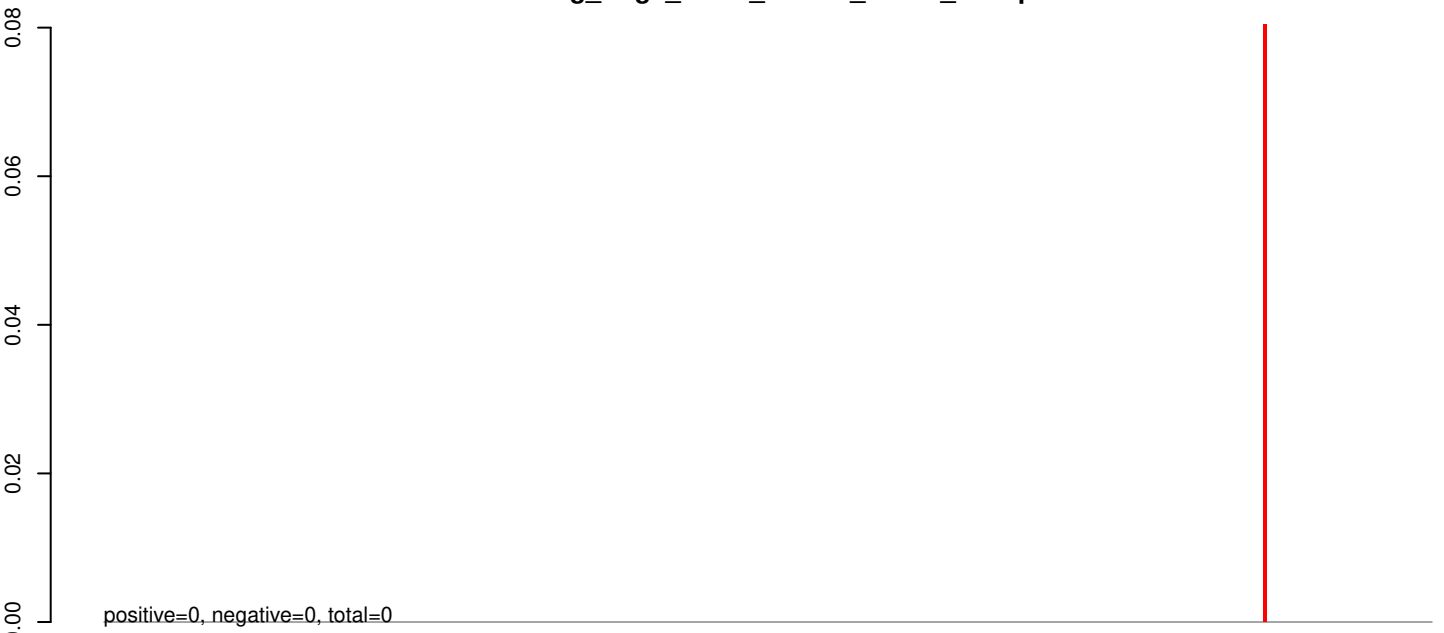


AeAeg_Aag2_mock_dsFluc_RA.rep

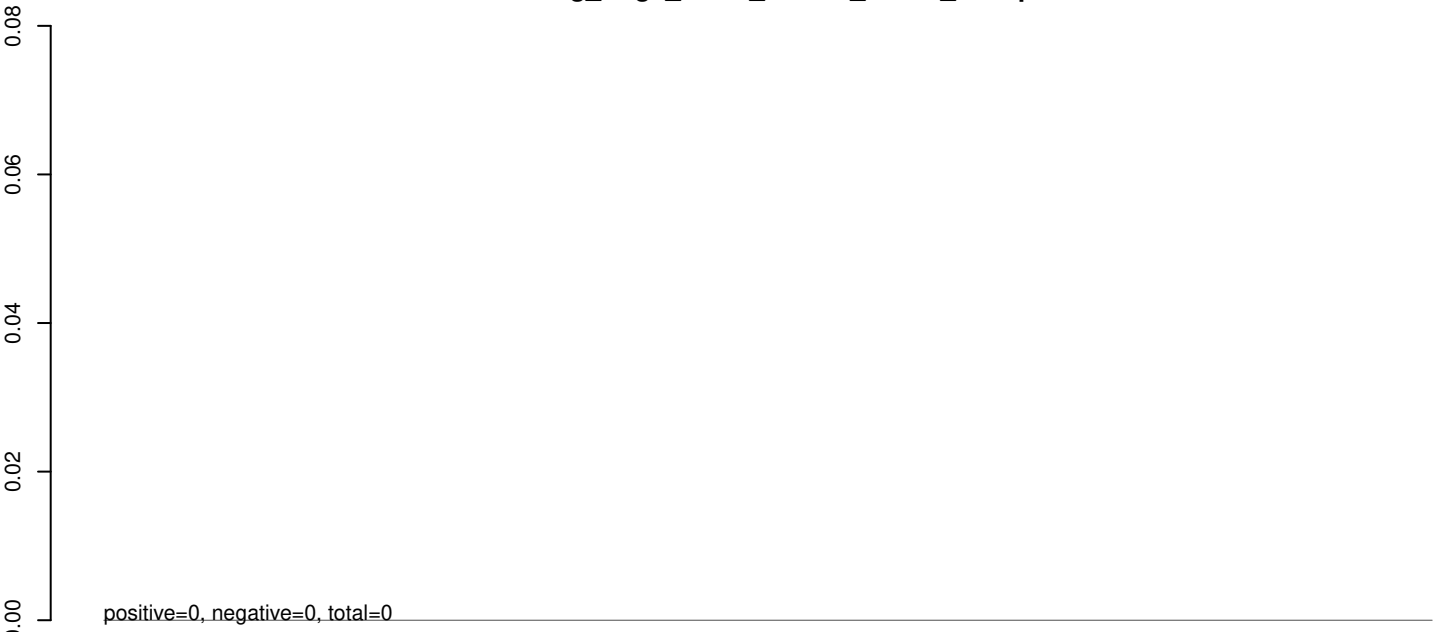


Window size=25, length=10173, virus@NC_033699.1-Jugra_virus_strain_P-9-314_polyprotein_gene:1-10173

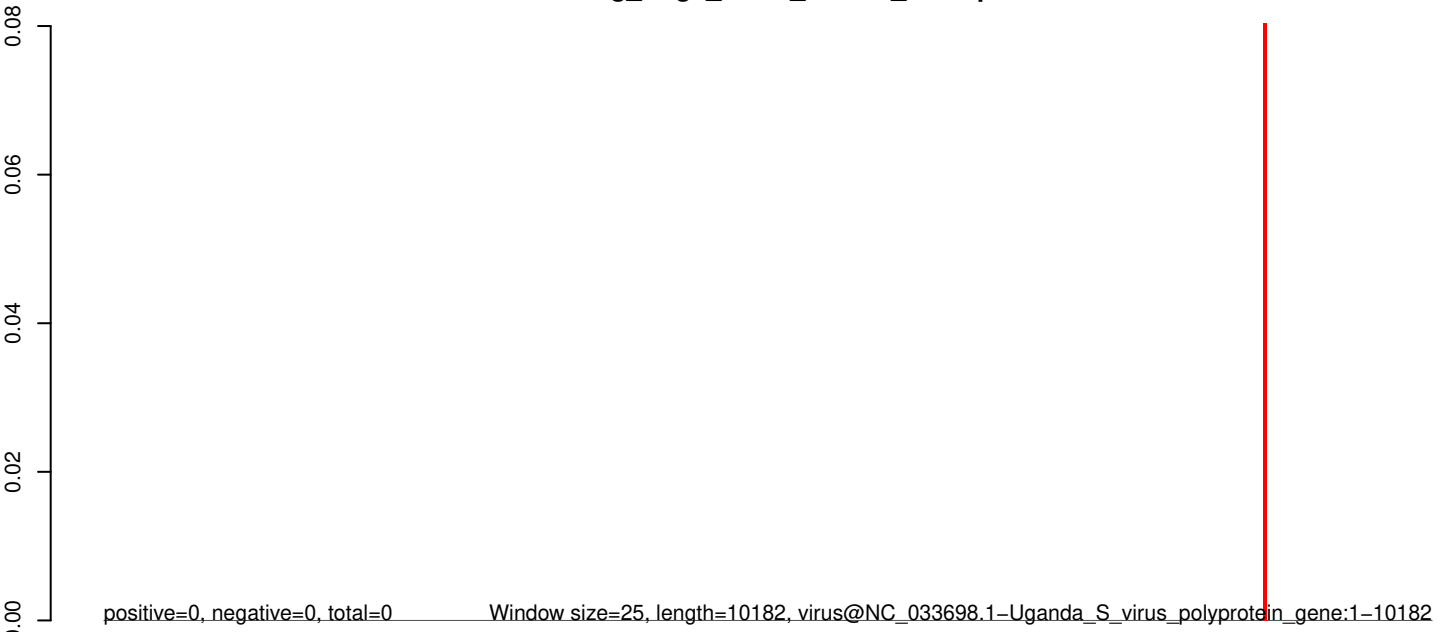
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

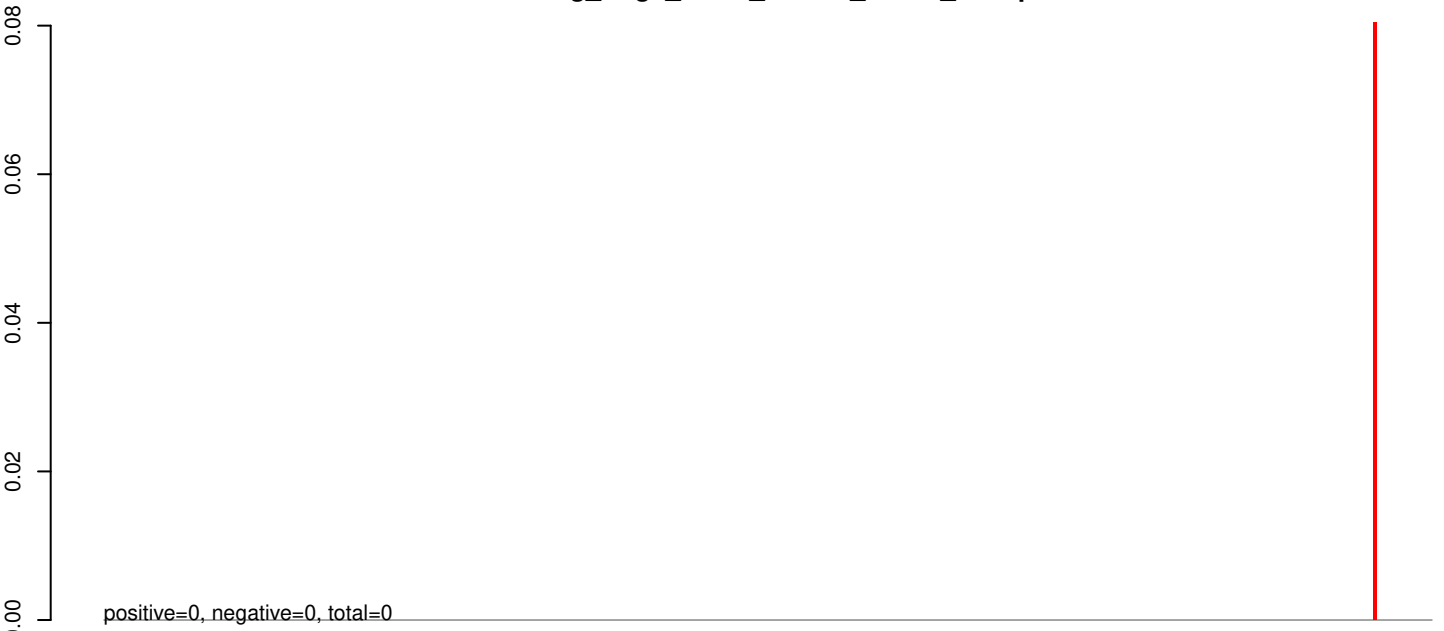


AeAeg_Aag2_mock_dsFluc_RA.rep

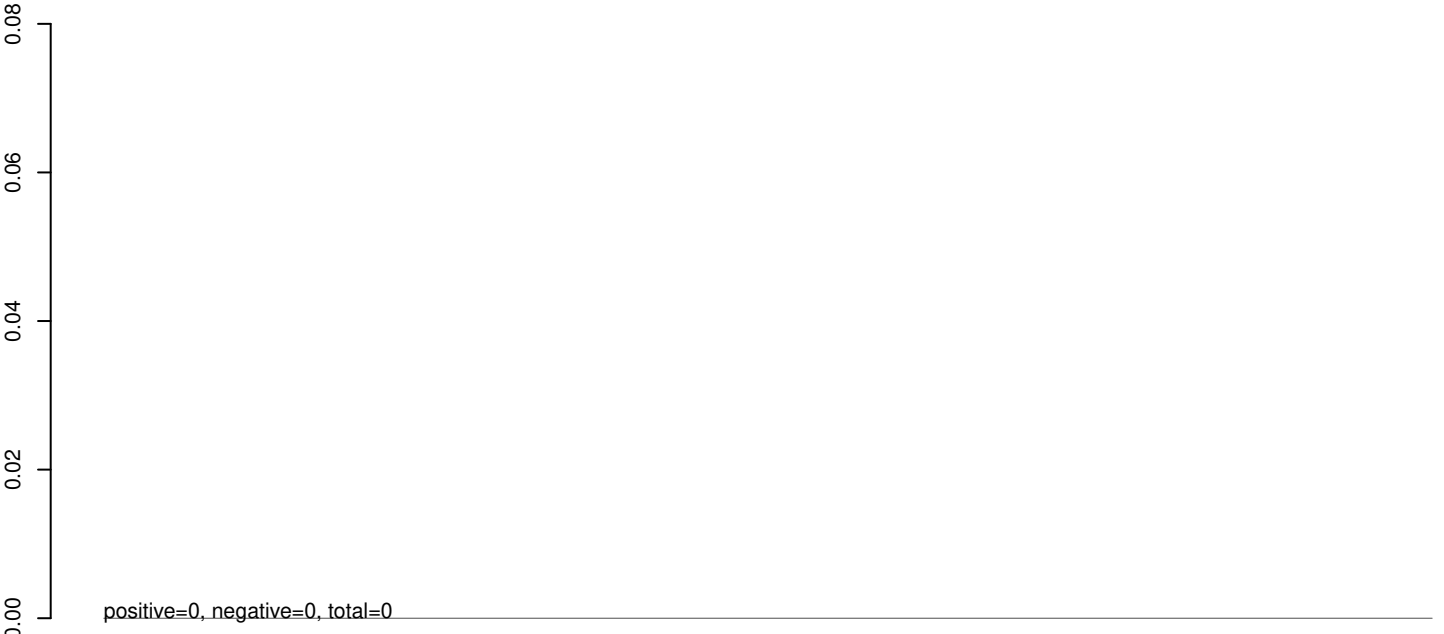


0 2000 4000 6000 8000 10000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



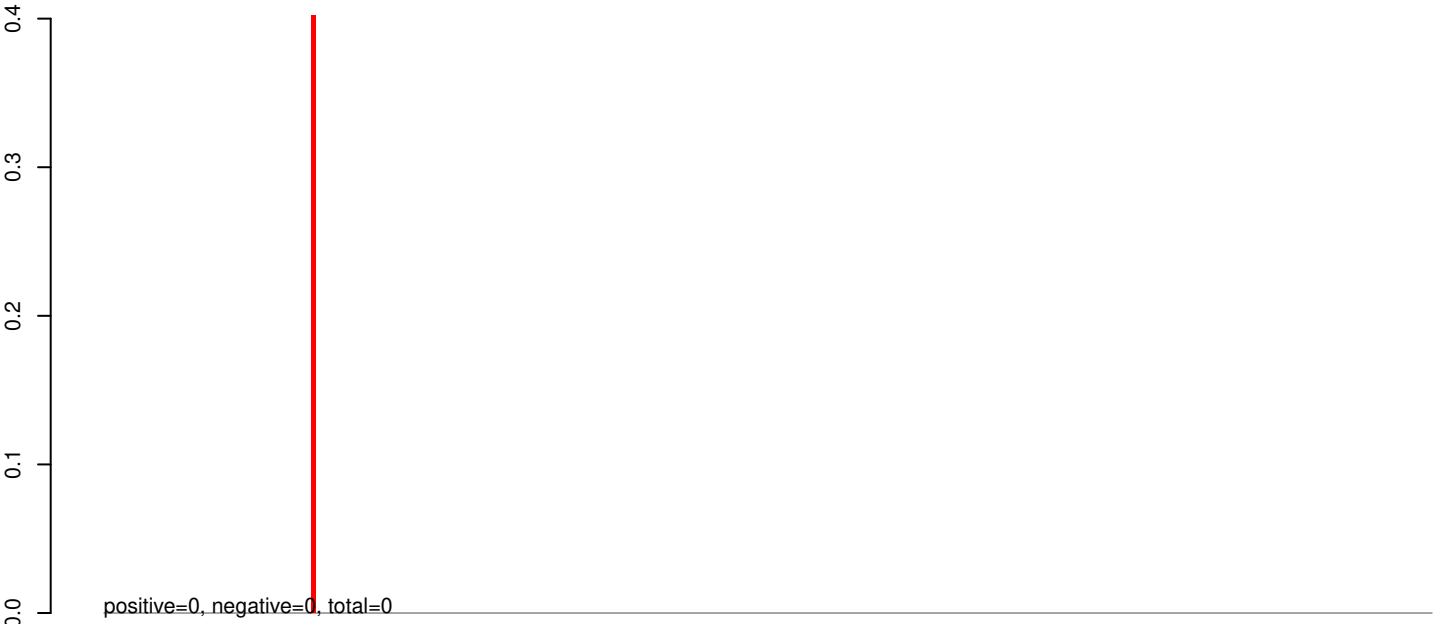
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



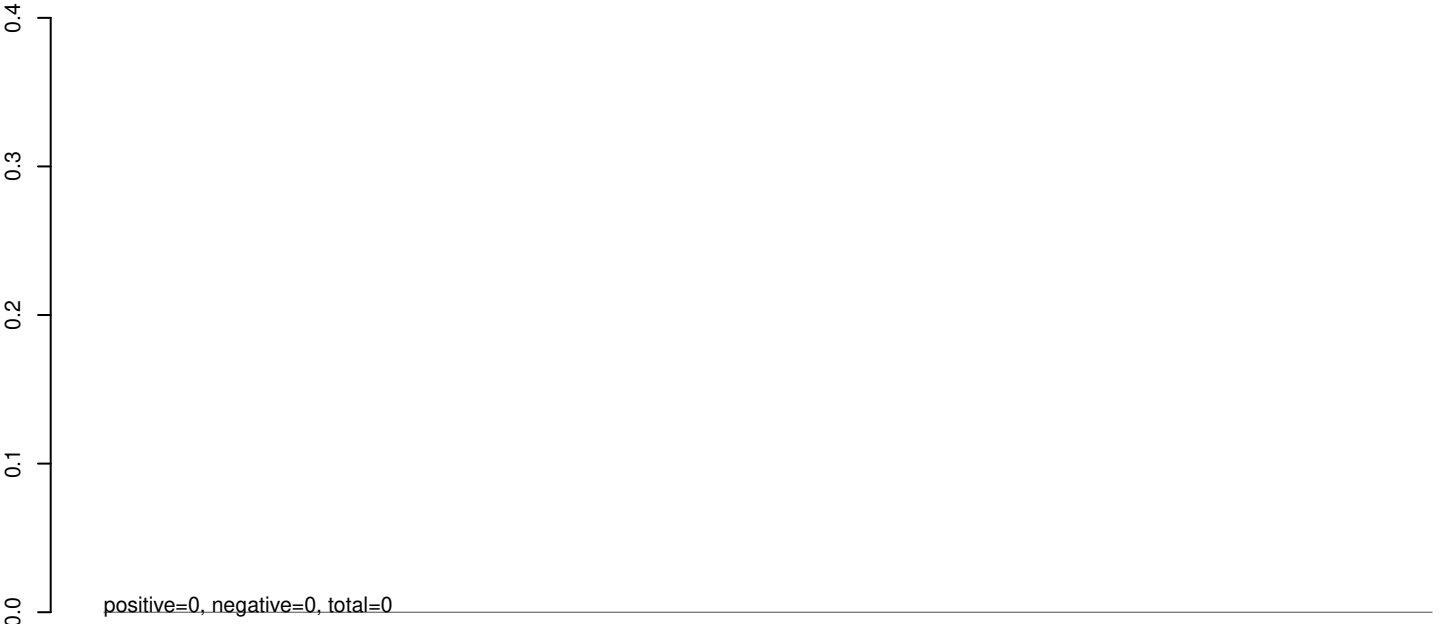
AeAeg_Aag2_mock_dsFluc_RA.rep



AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



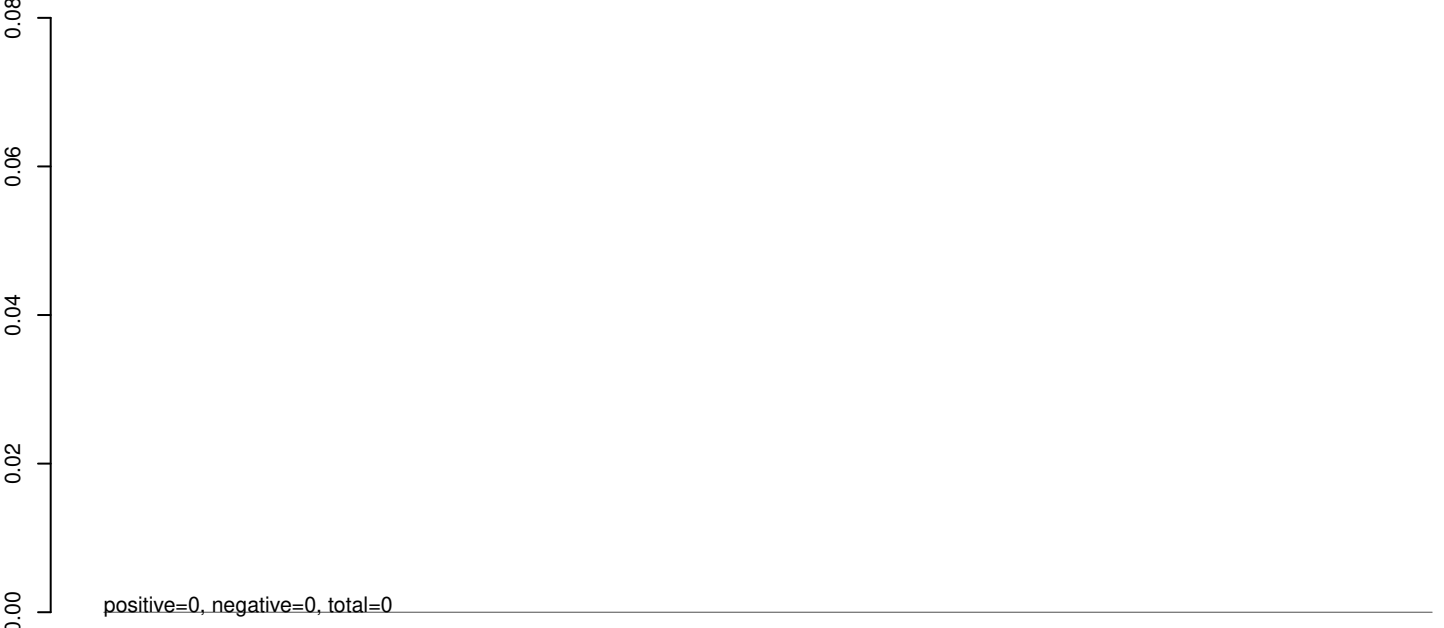
AeAeg_Aag2_mock_dsFluc_RA.rep



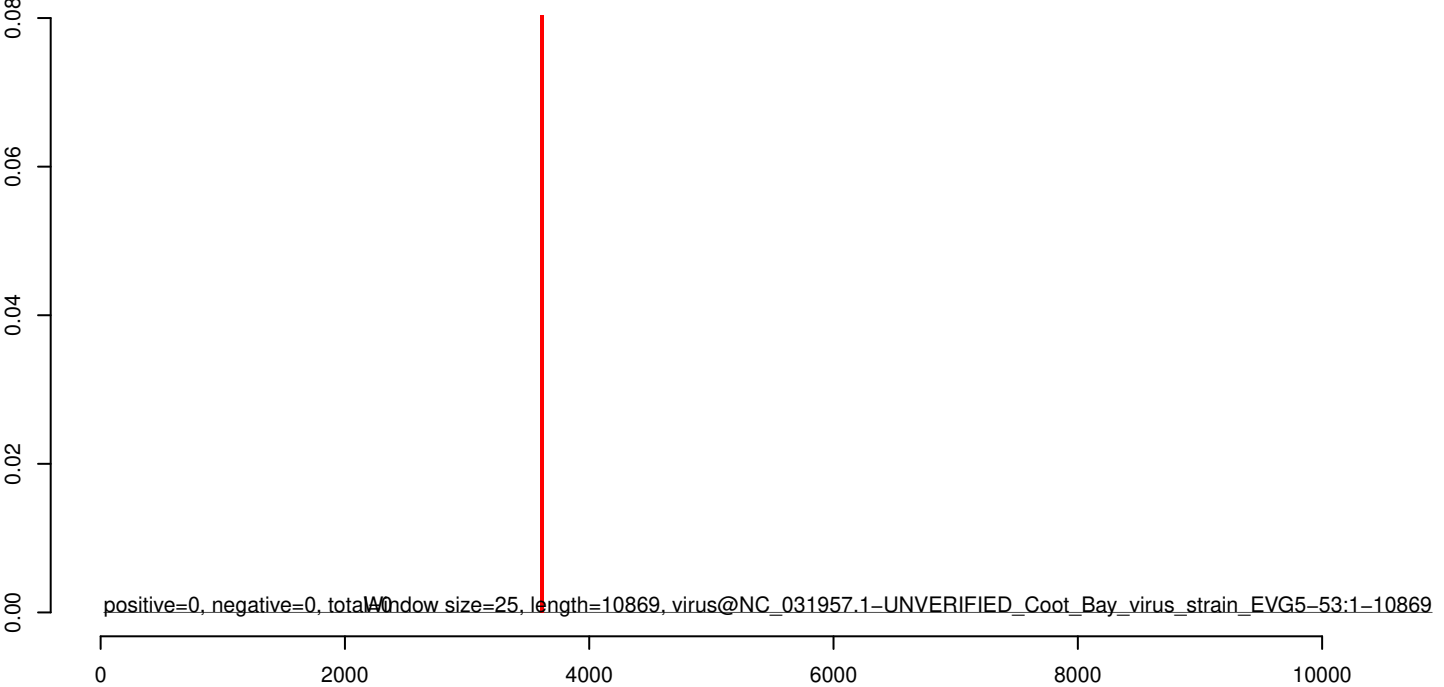
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

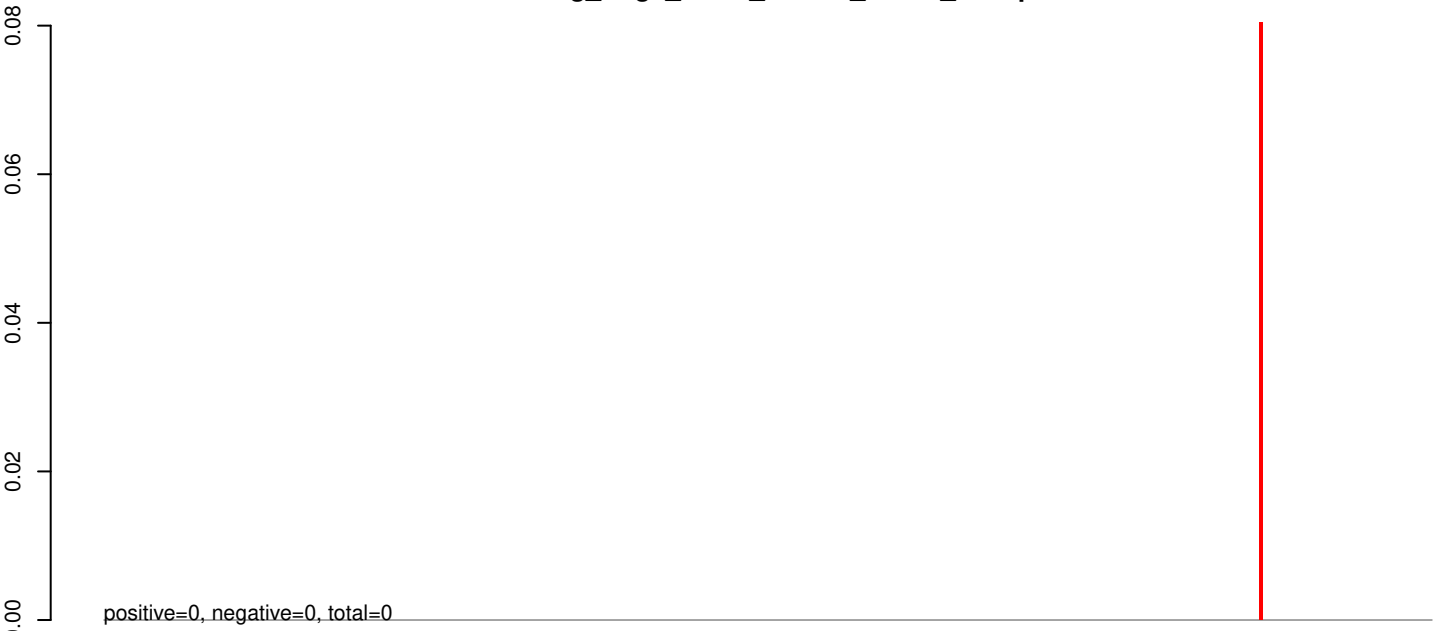


AeAeg_Aag2_mock_dsFluc_RA.rep

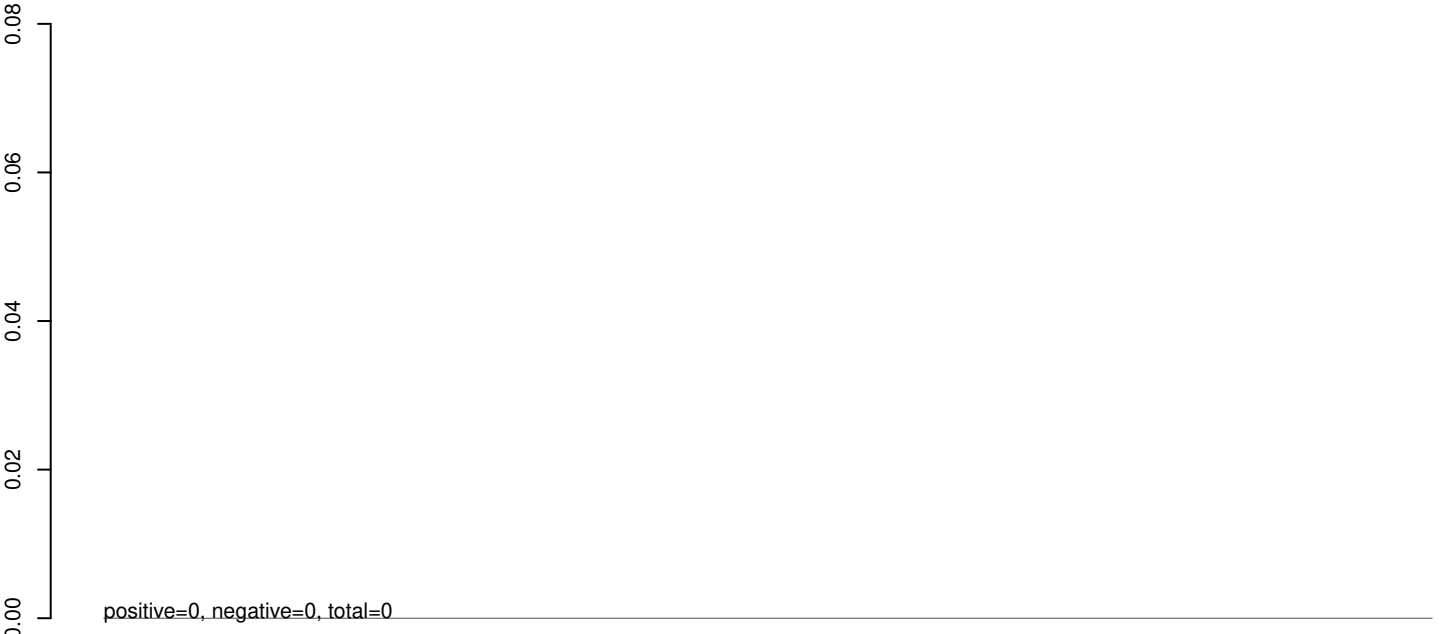


Window size=25, length=10869, virus@NC_031957.1-UNVERIFIED_Coot_Bay_virus_strain_EVG5-53:1-10869

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



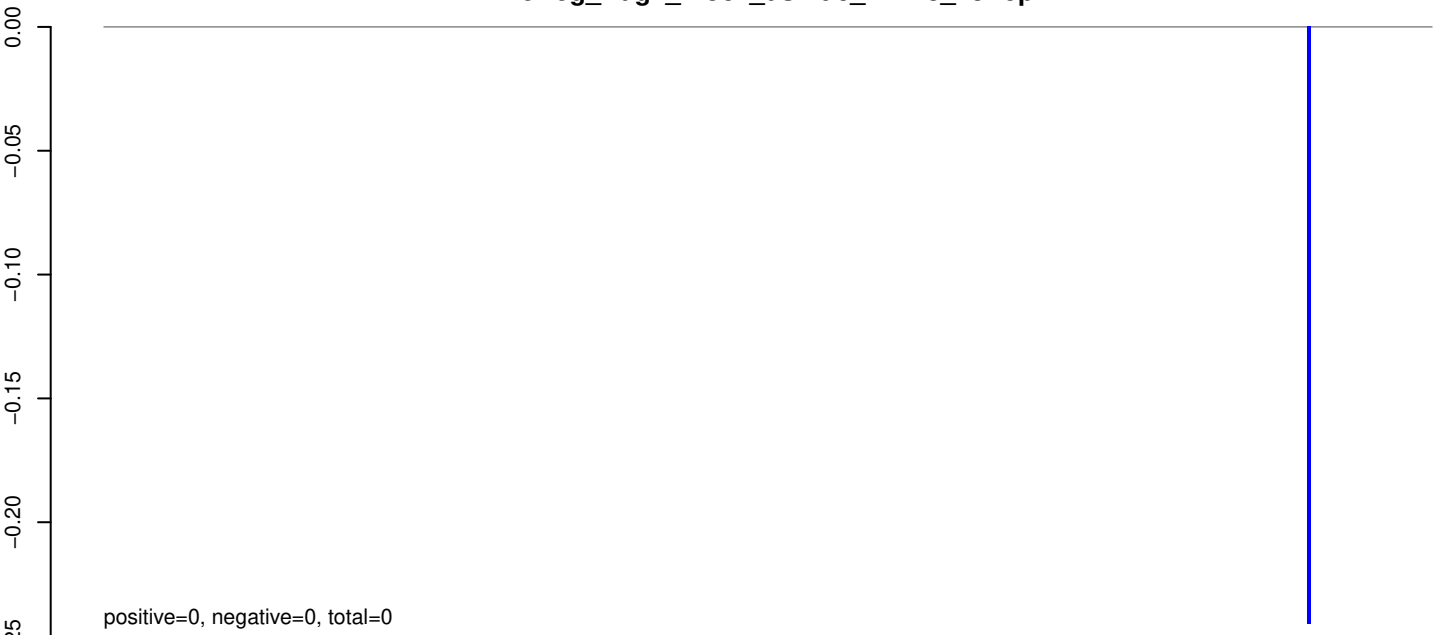
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep

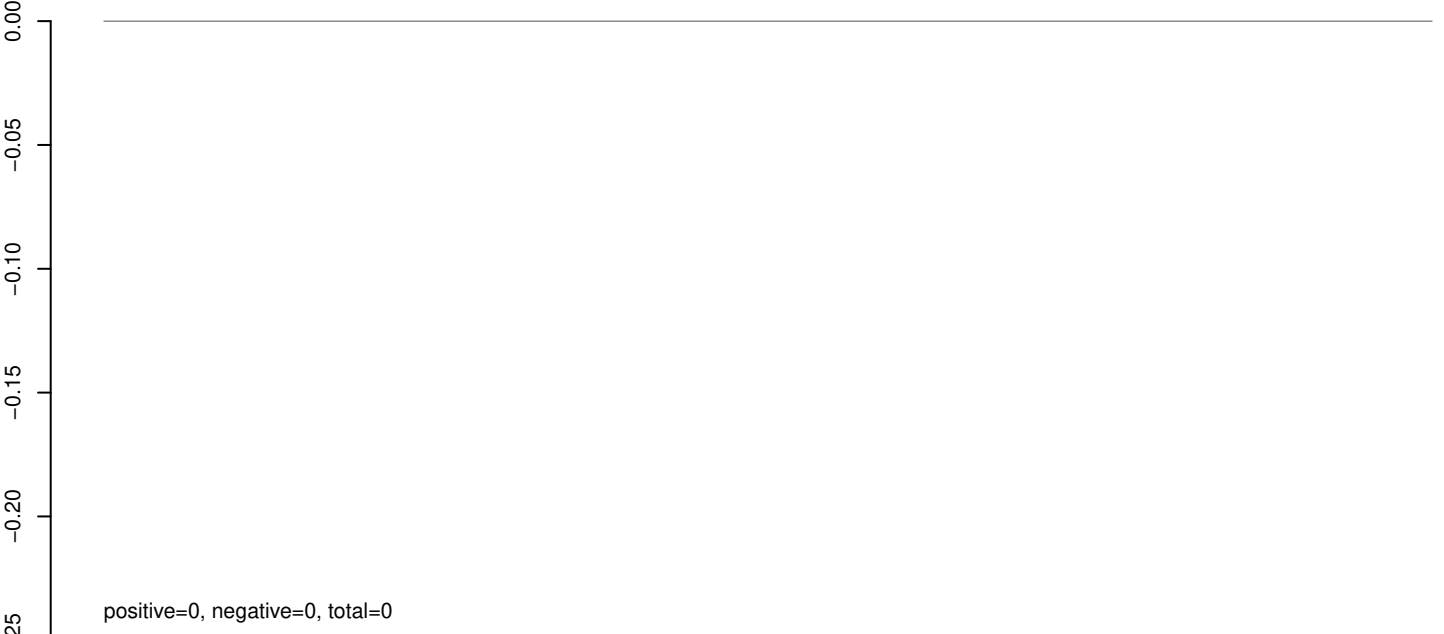


AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



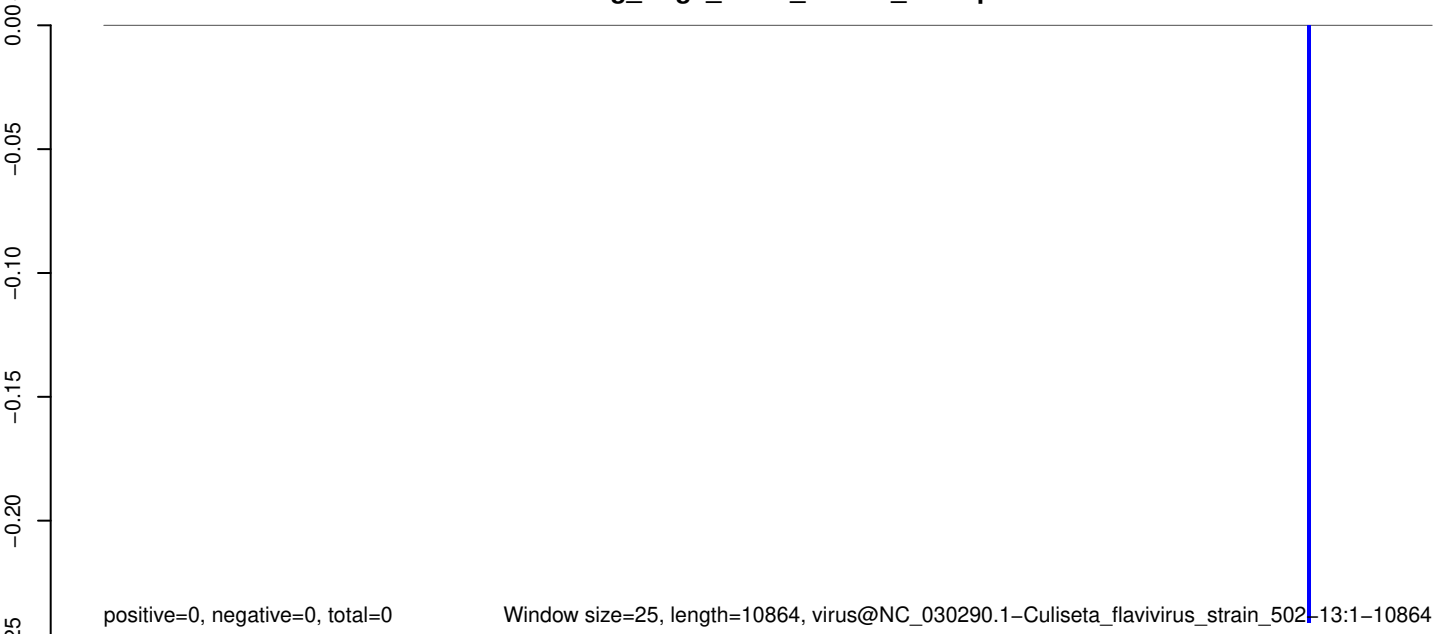
positive=0, negative=0, total=0

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



positive=0, negative=0, total=0

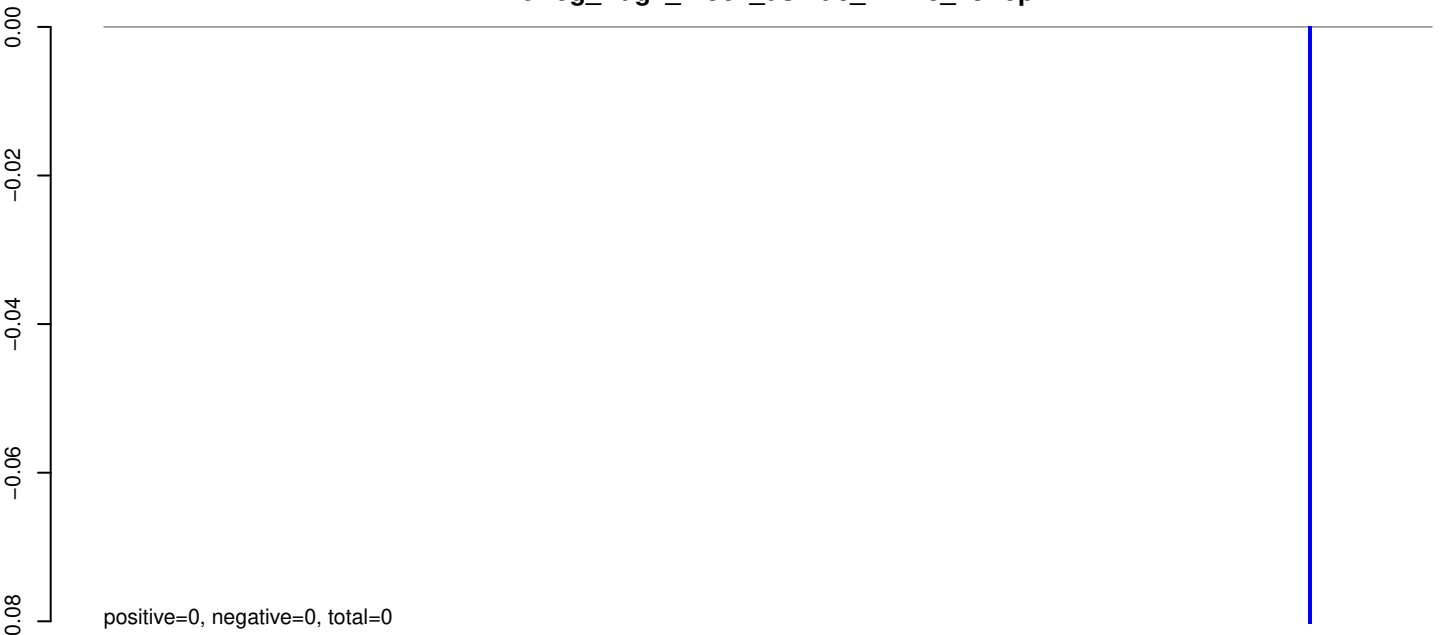
AeAeg_Aag2_mock_dsFluc_RA.rep



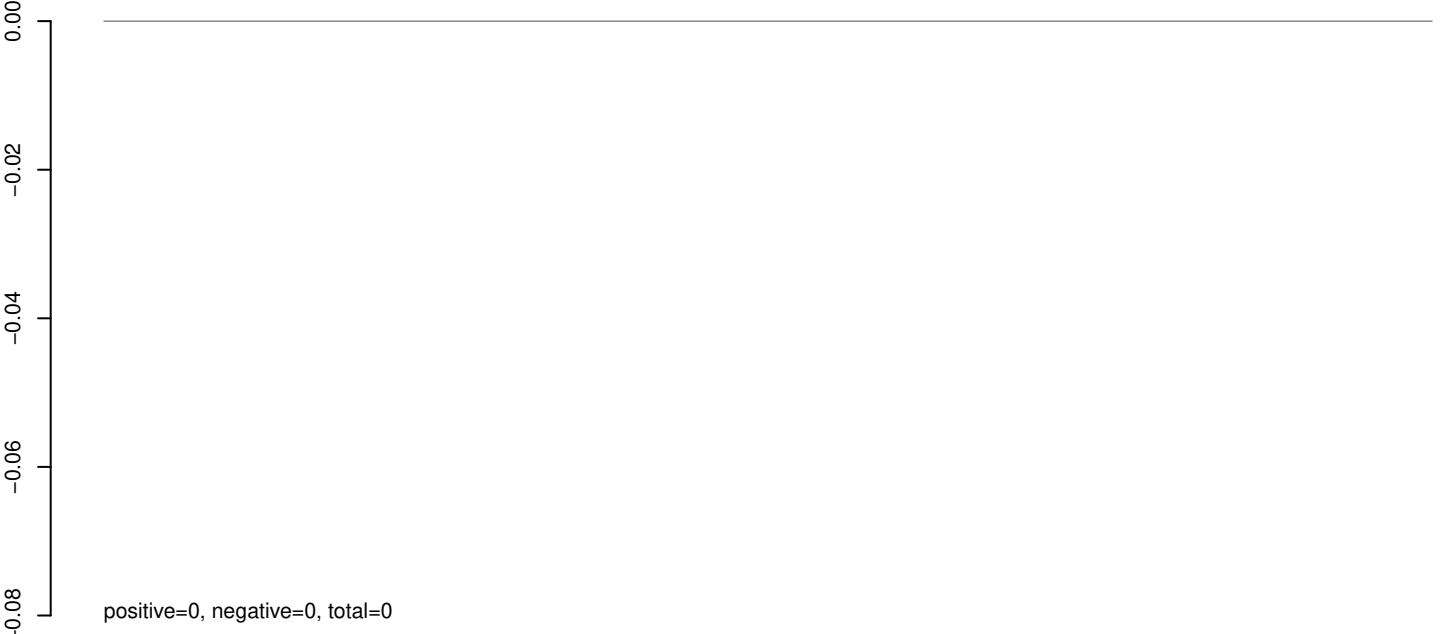
positive=0, negative=0, total=0

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

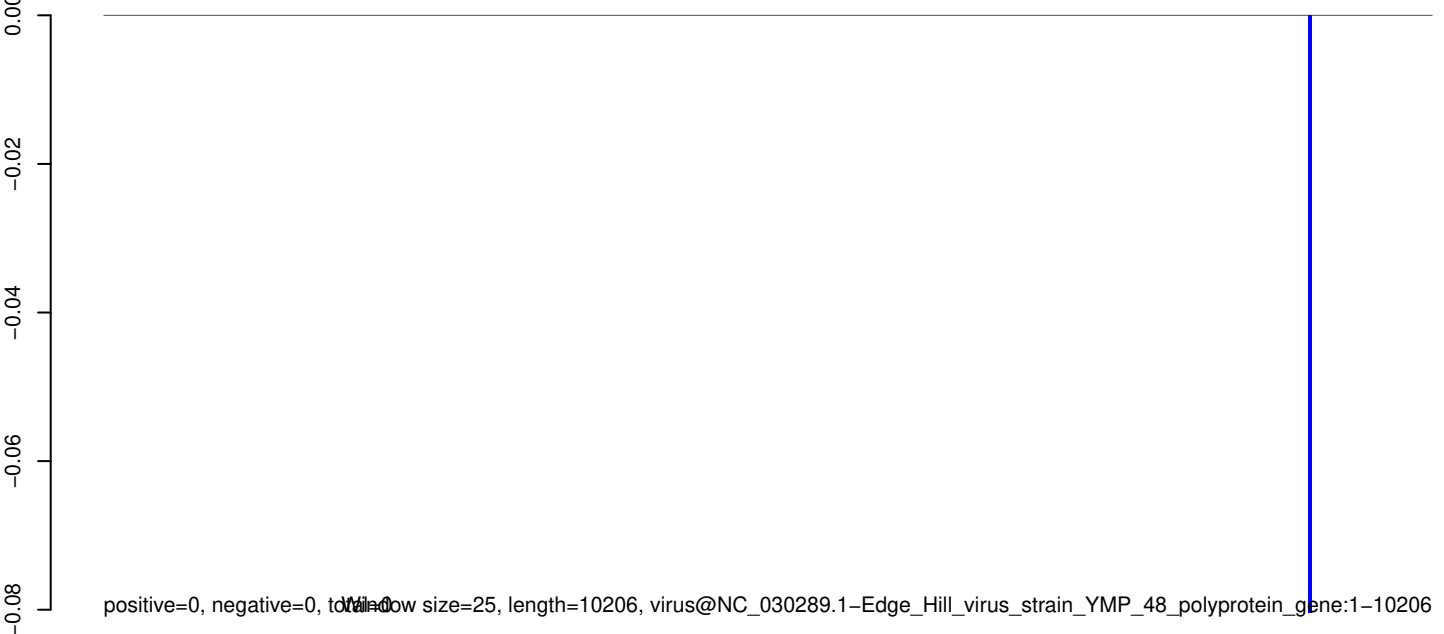
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



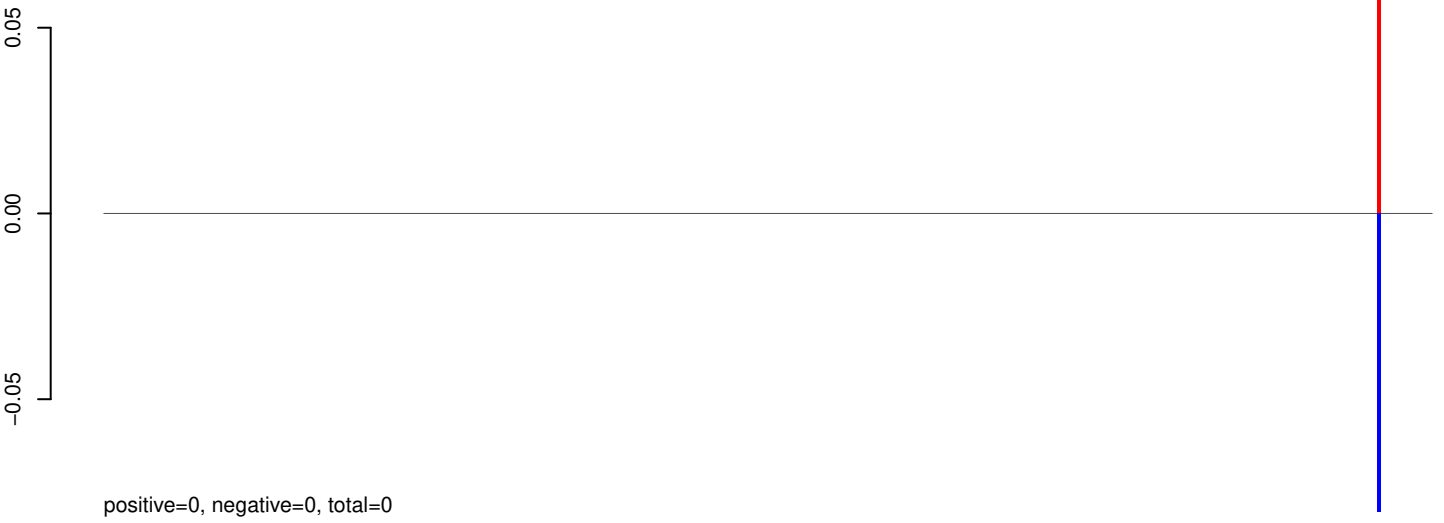
AeAeg_Aag2_mock_dsFluc_RA.rep



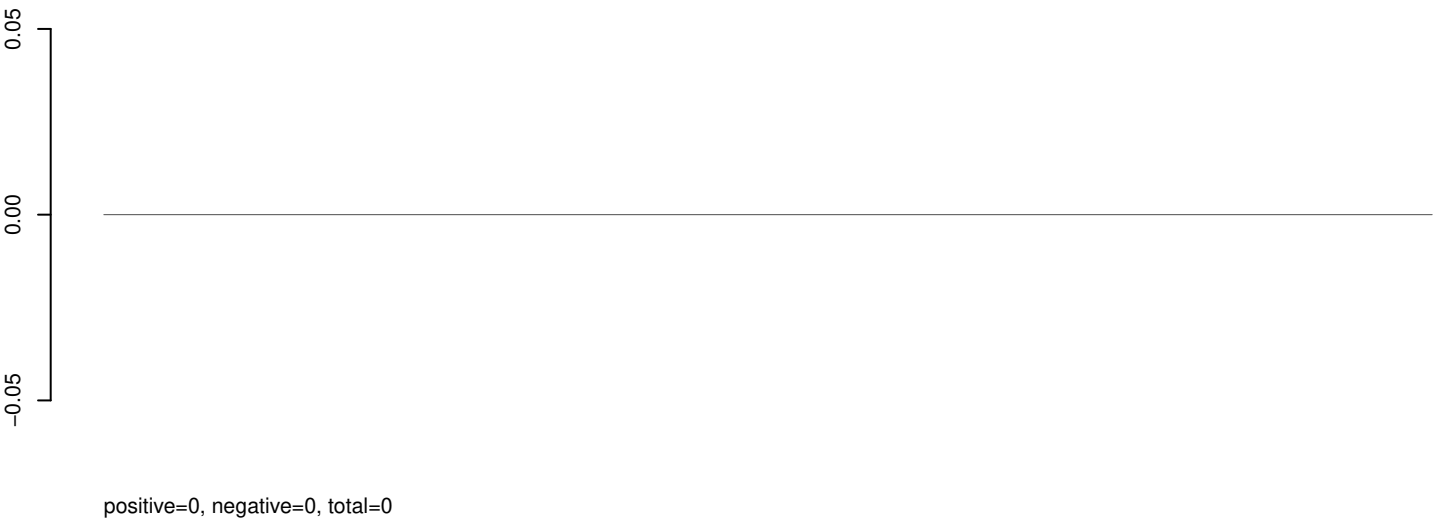
Window size=25, length=10206, virus@NC_030289.1-Edge_Hill_virus_strain_YMP_48_polyprotein_gene:1-10206

0 2000 4000 6000 8000 10000

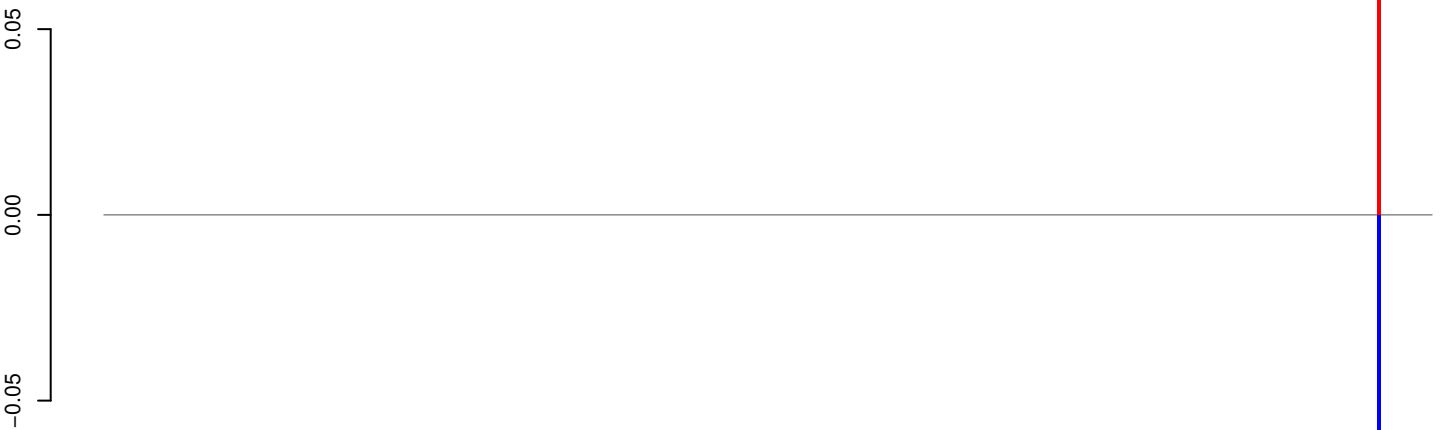
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



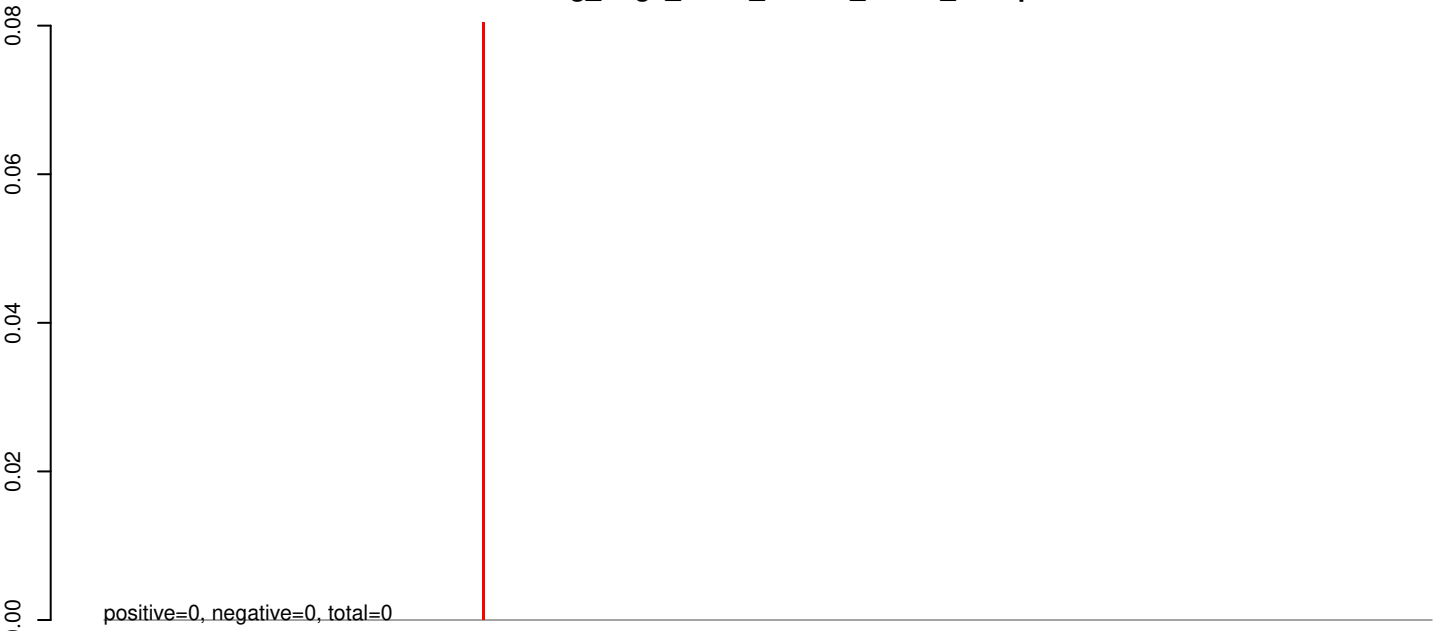
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



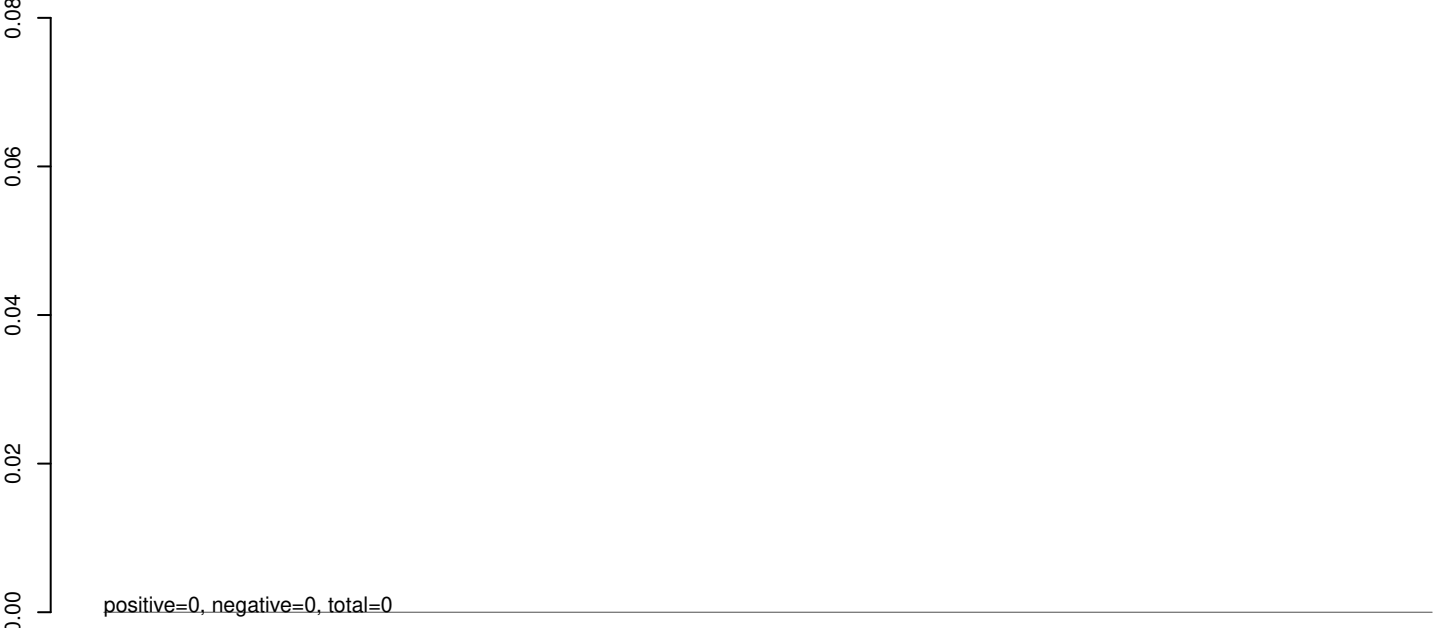
AeAeg_Aag2_mock_dsFluc_RA.rep



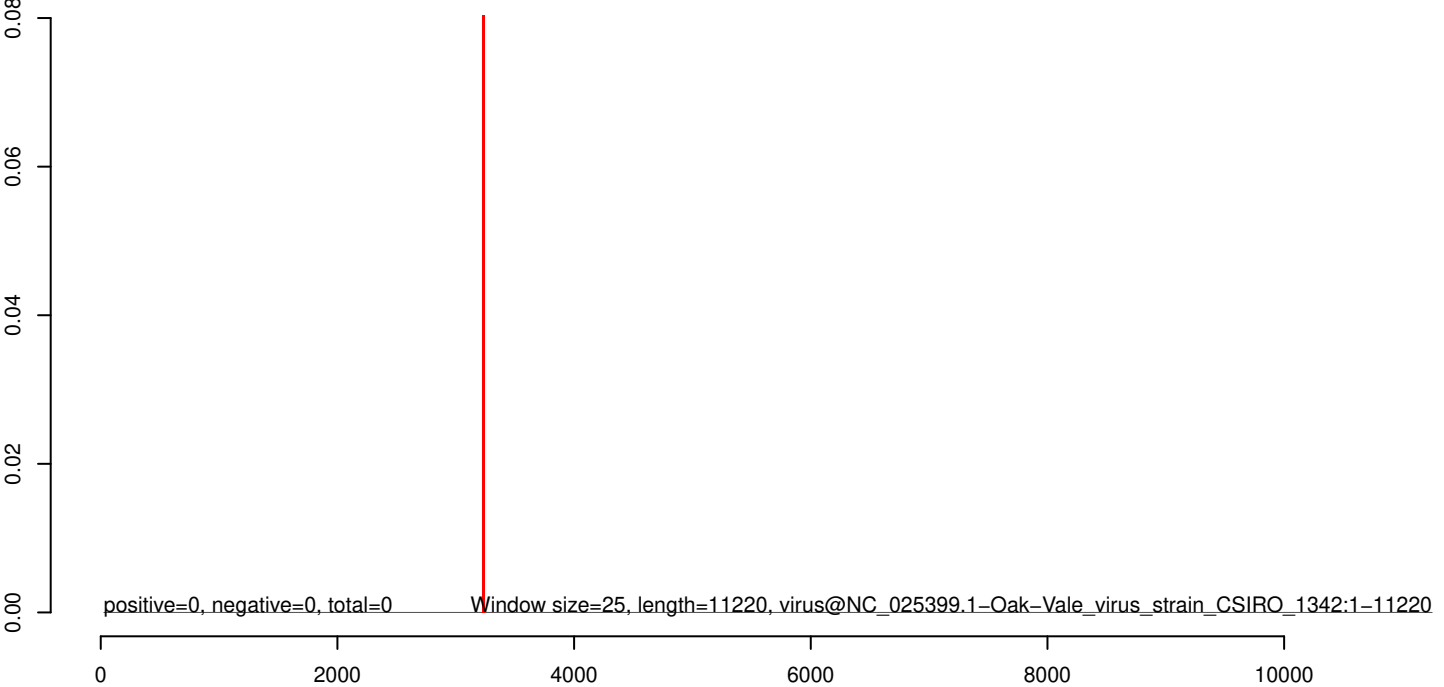
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



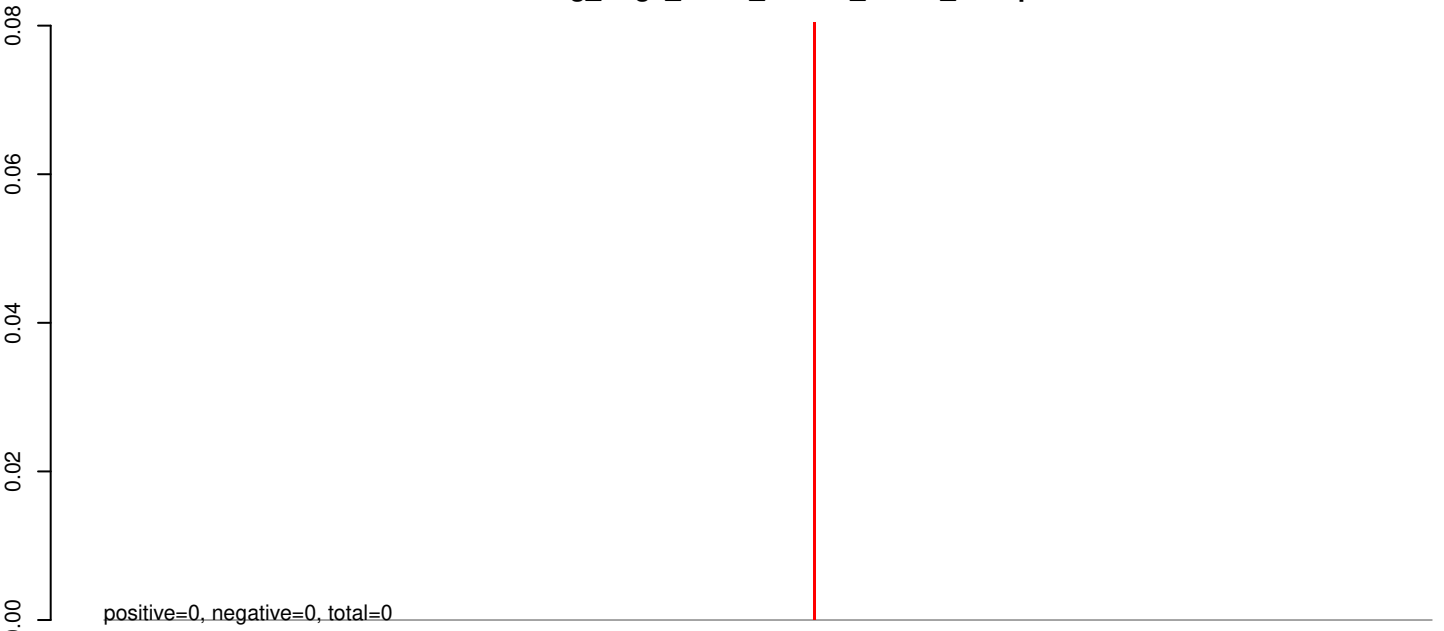
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



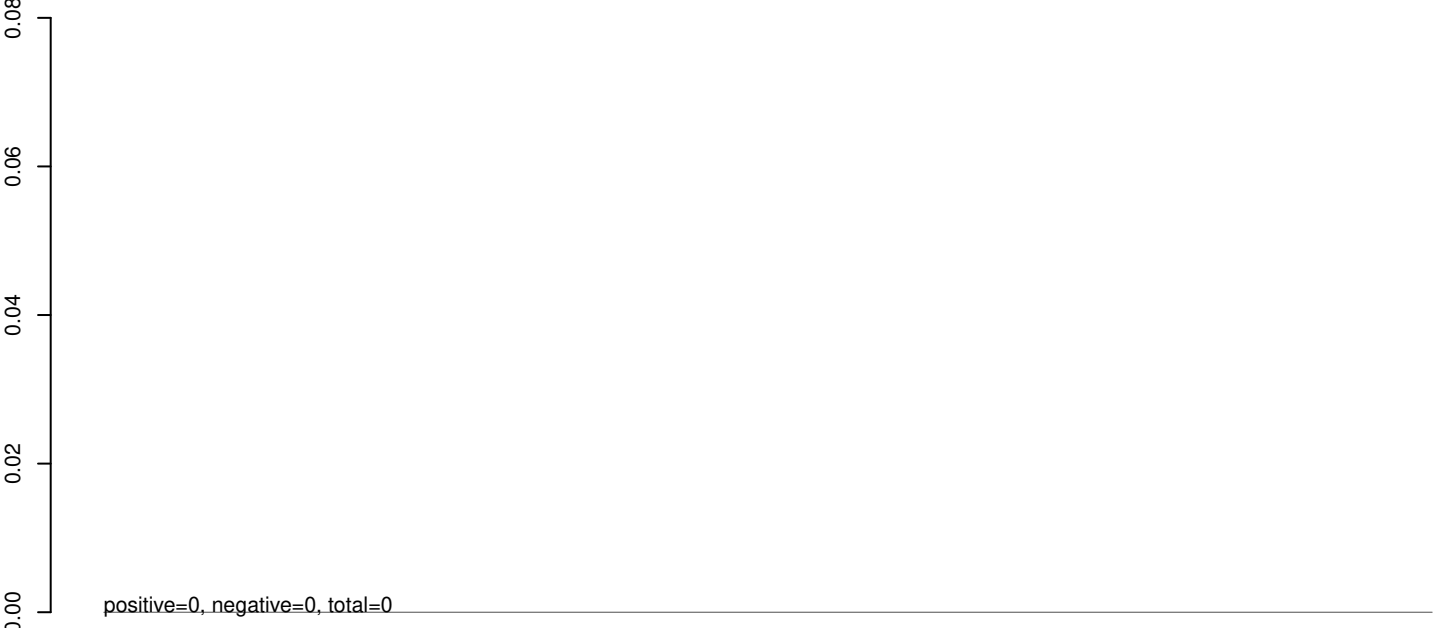
AeAeg_Aag2_mock_dsFluc_RA.rep



AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



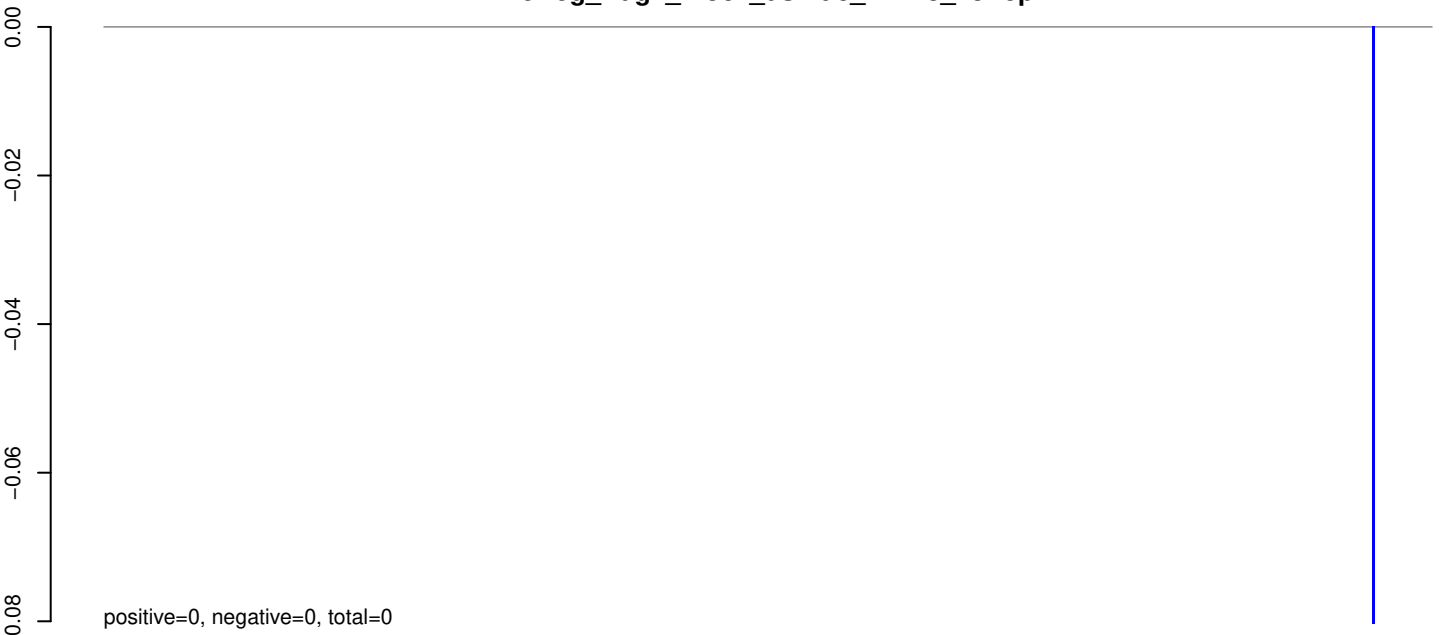
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



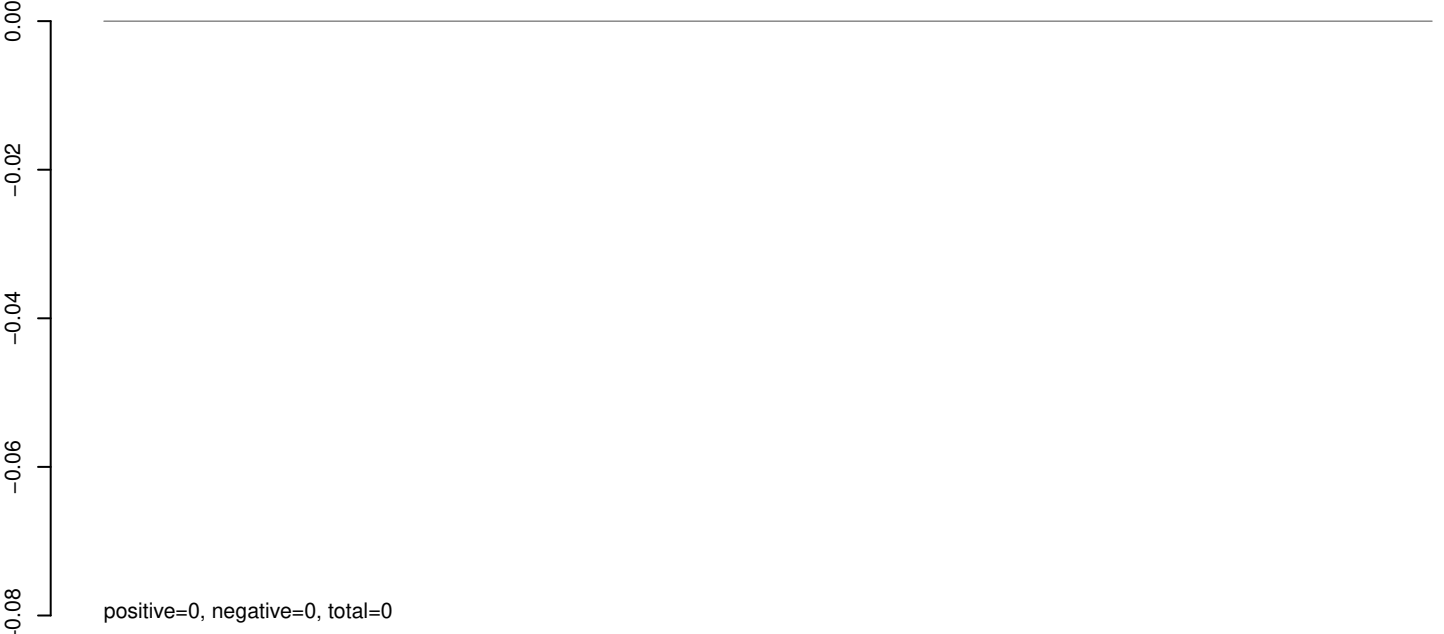
AeAeg_Aag2_mock_dsFluc_RA.rep



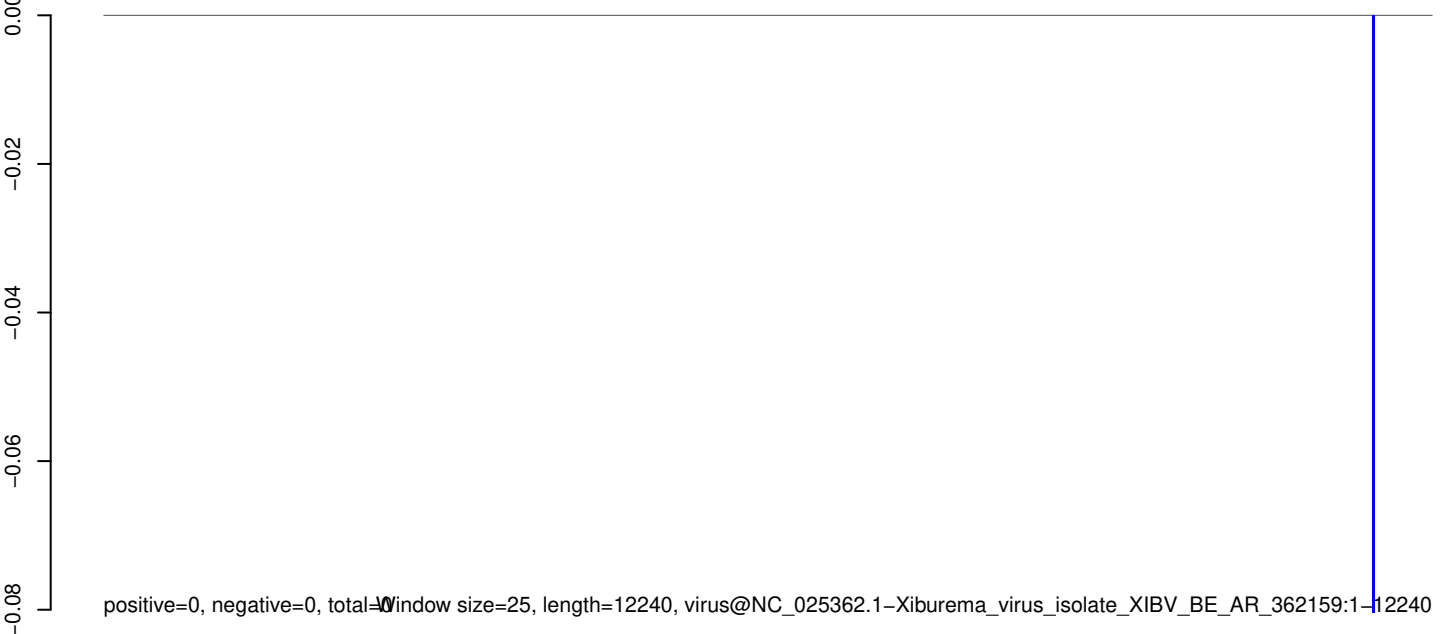
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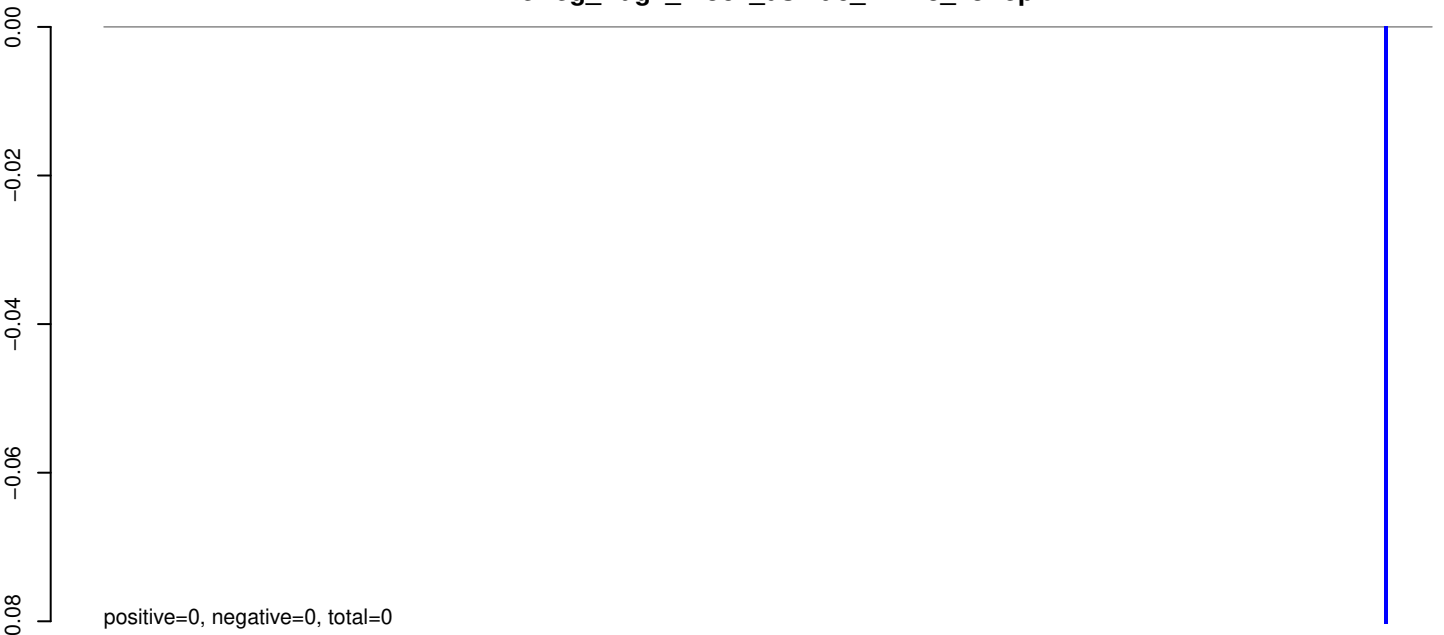
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep

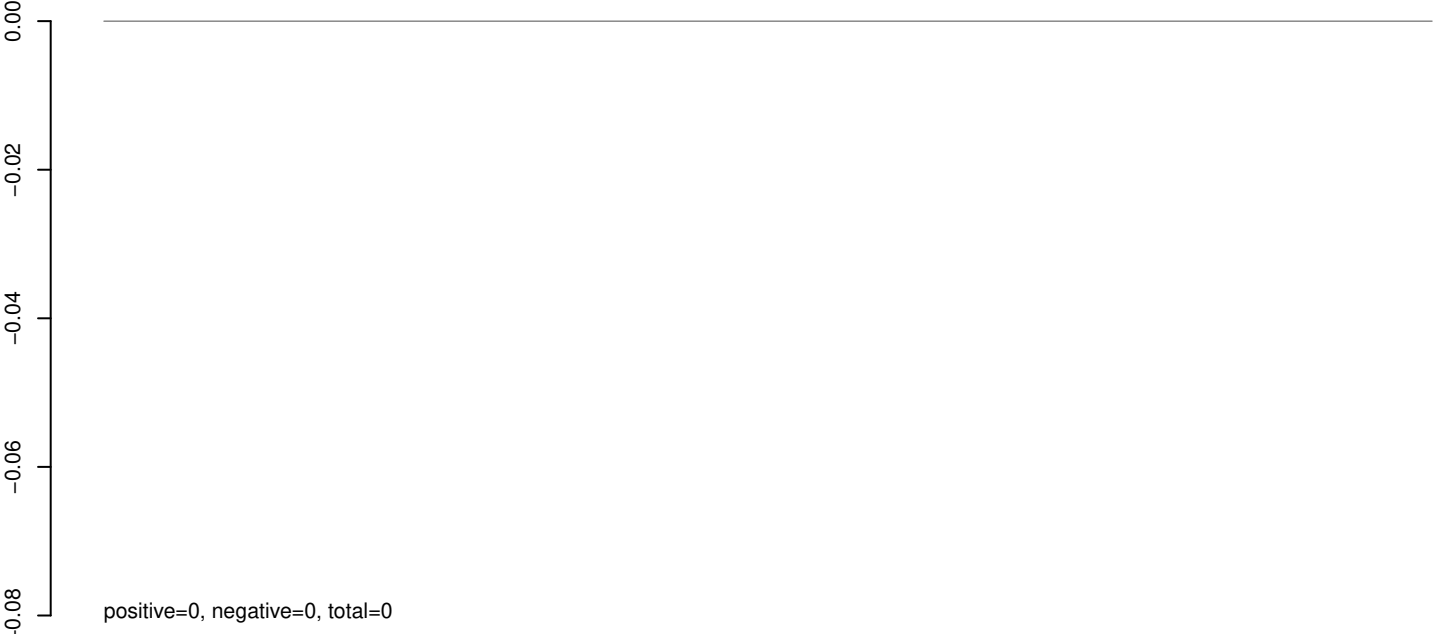


AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



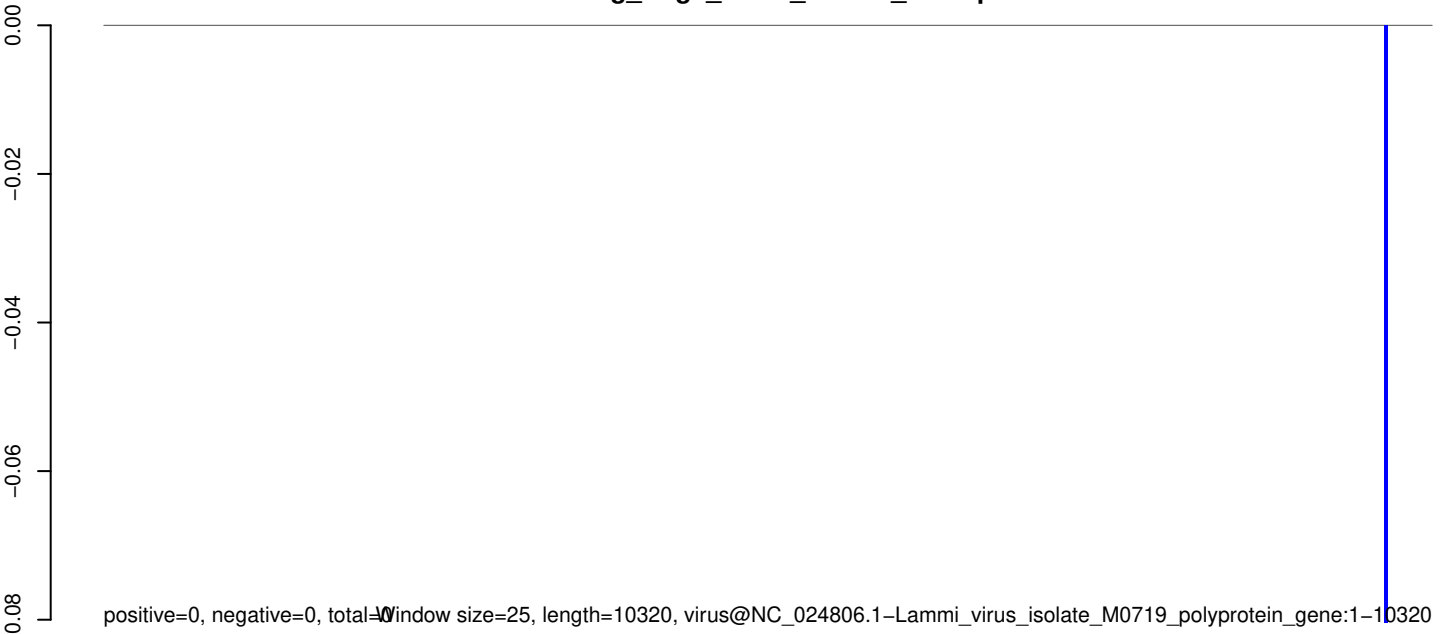
positive=0, negative=0, total=0

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



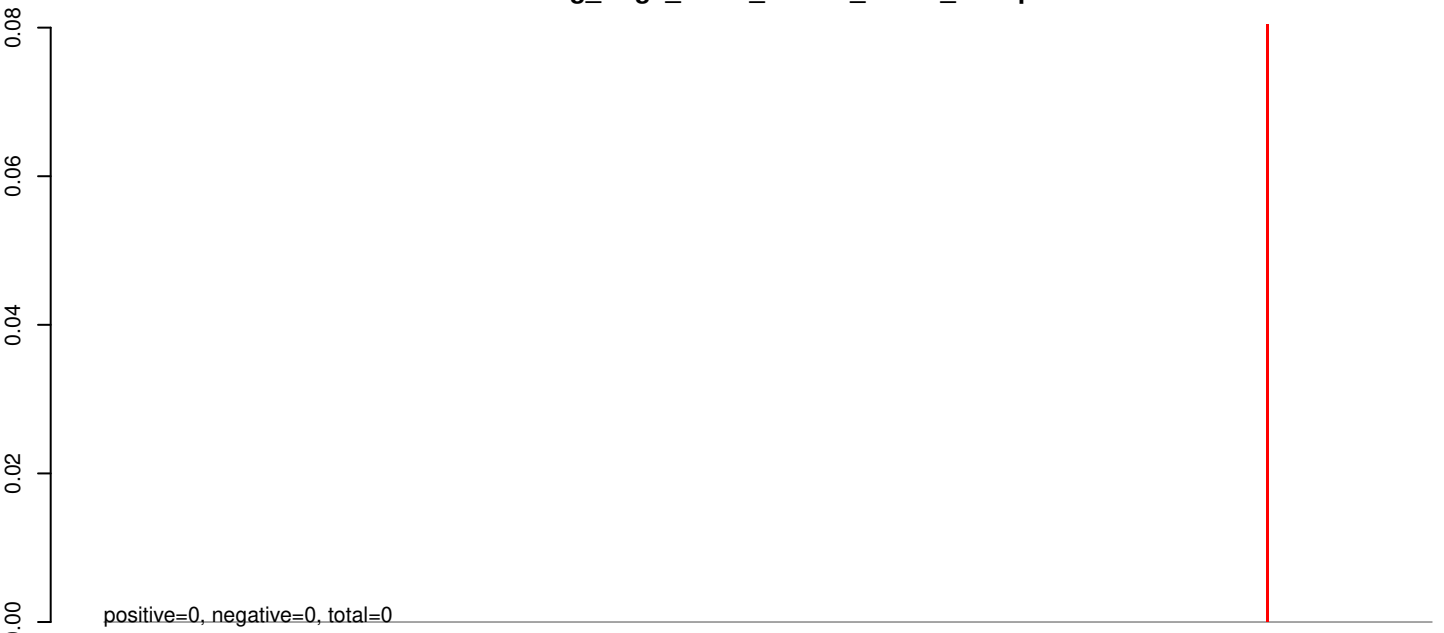
positive=0, negative=0, total=0

AeAeg_Aag2_mock_dsFluc_RA.rep

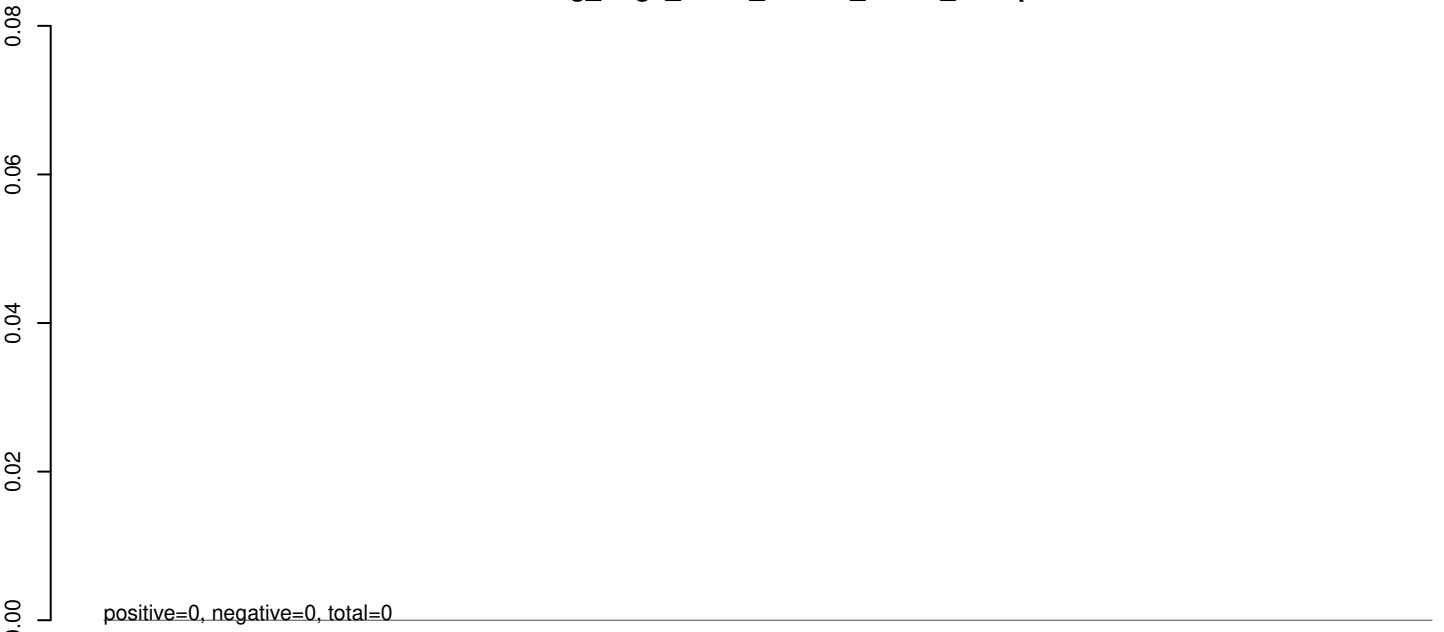


positive=0, negative=0, total=0 window size=25, length=10320, virus@NC_024806.1-Lammi_virus_isolate_M0719_polyprotein_gene:1-10320

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



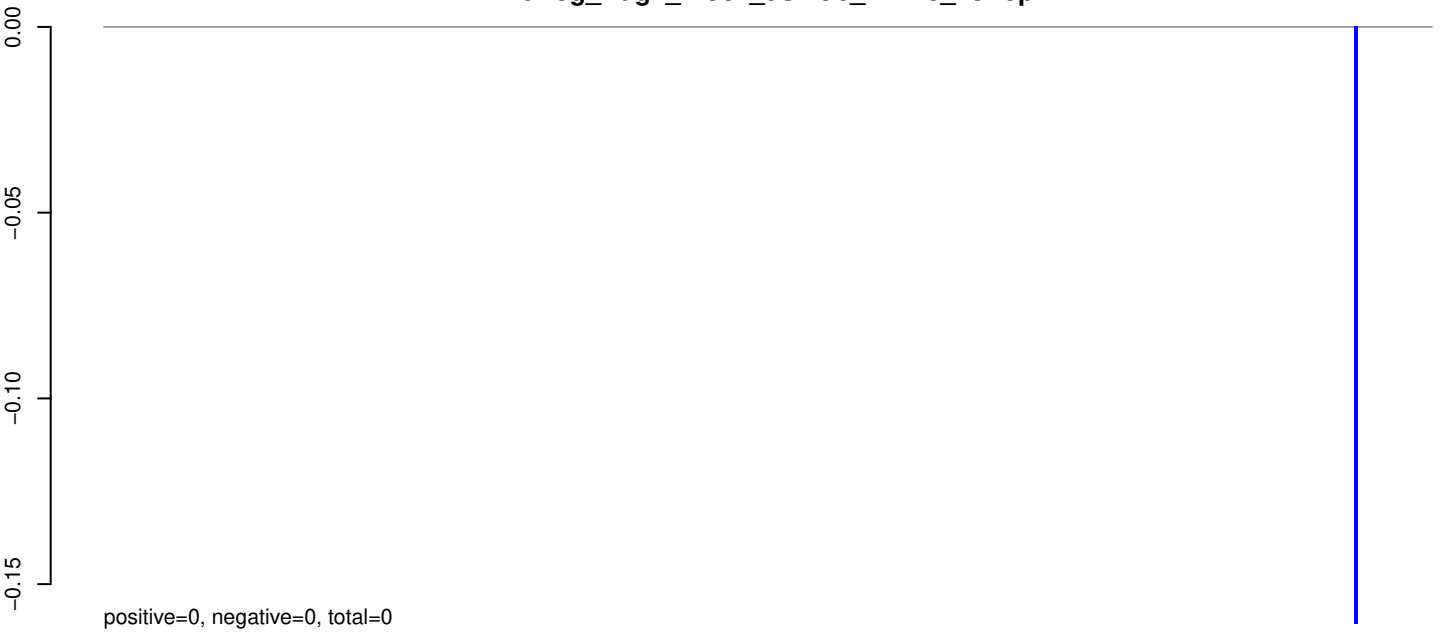
AeAeg_Aag2_mock_dsFluc_RA.rep



11624, virus@NC_023812.1-Madariaga_virus_strain_MADV_Cebus_apella_BRA_BEAN5122_1956:1-11624

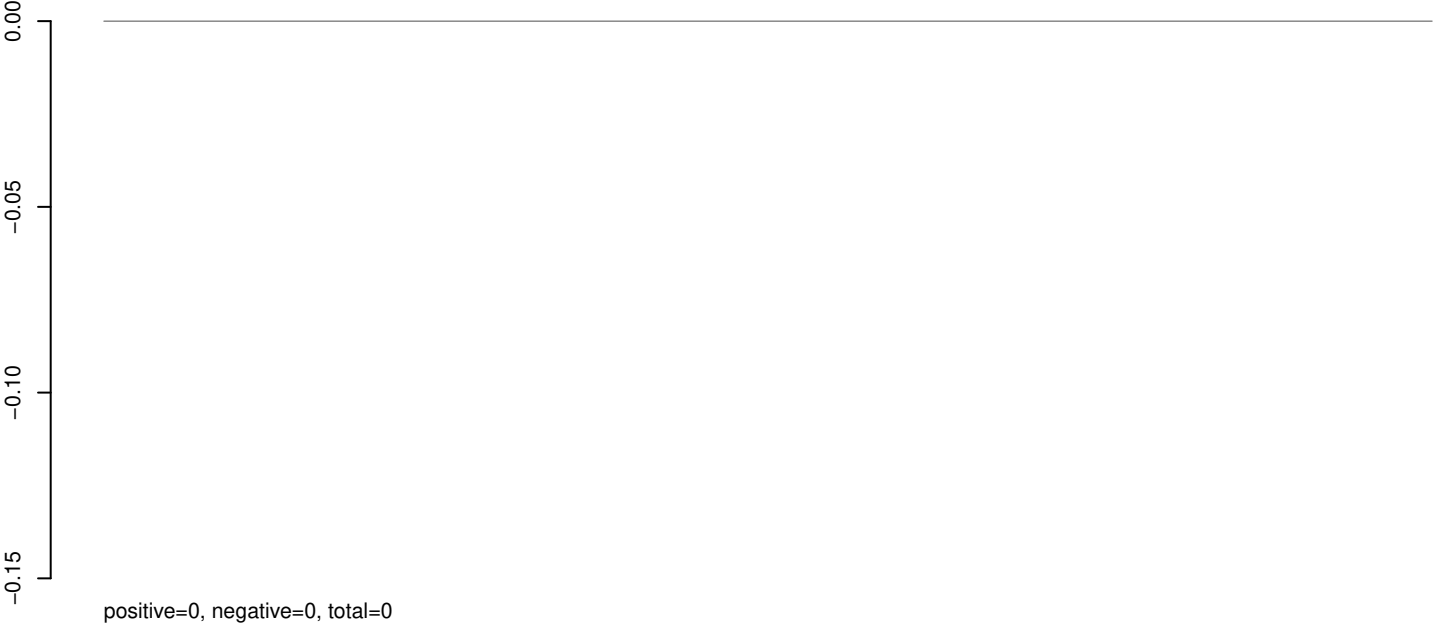
0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



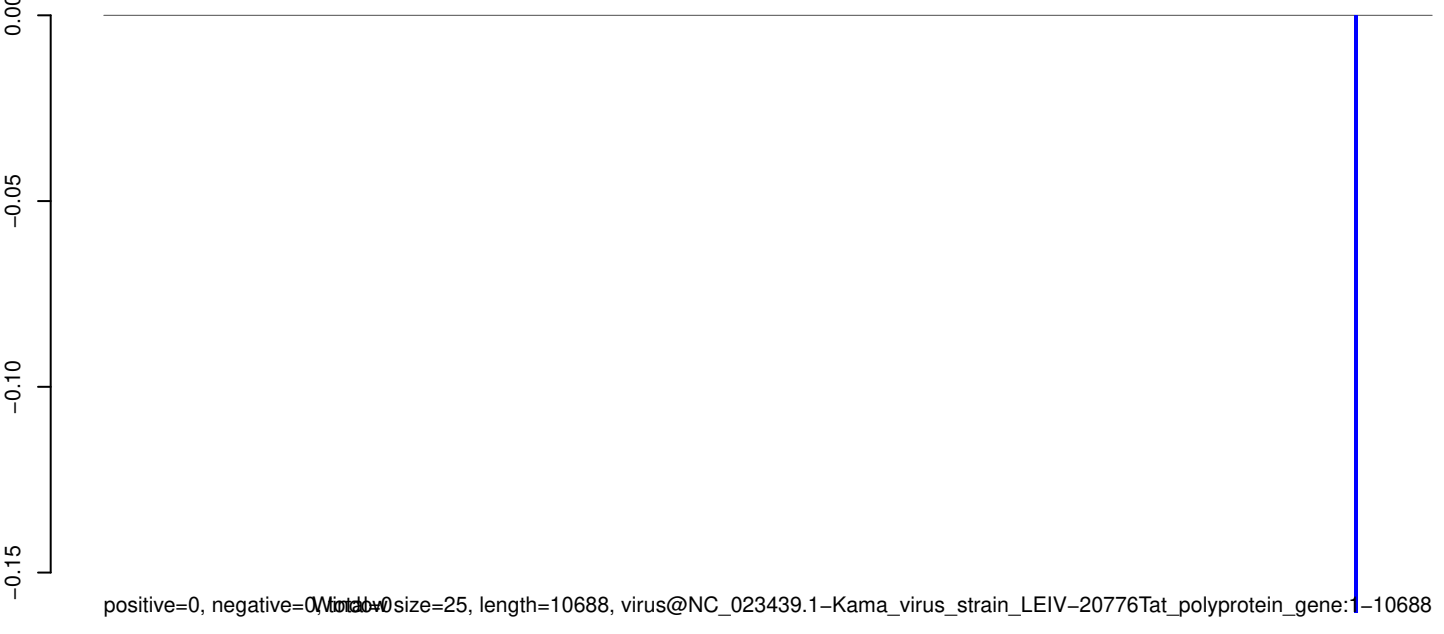
positive=0, negative=0, total=0

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



positive=0, negative=0, total=0

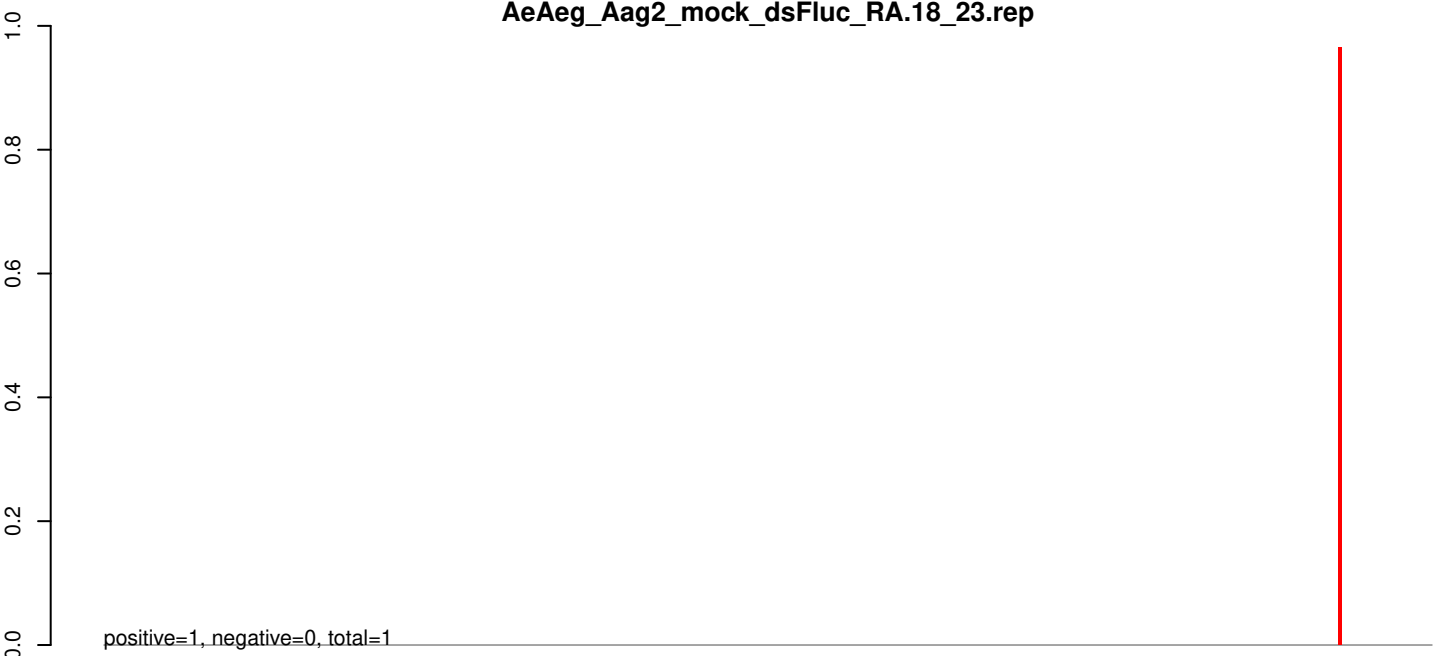
AeAeg_Aag2_mock_dsFluc_RA.rep



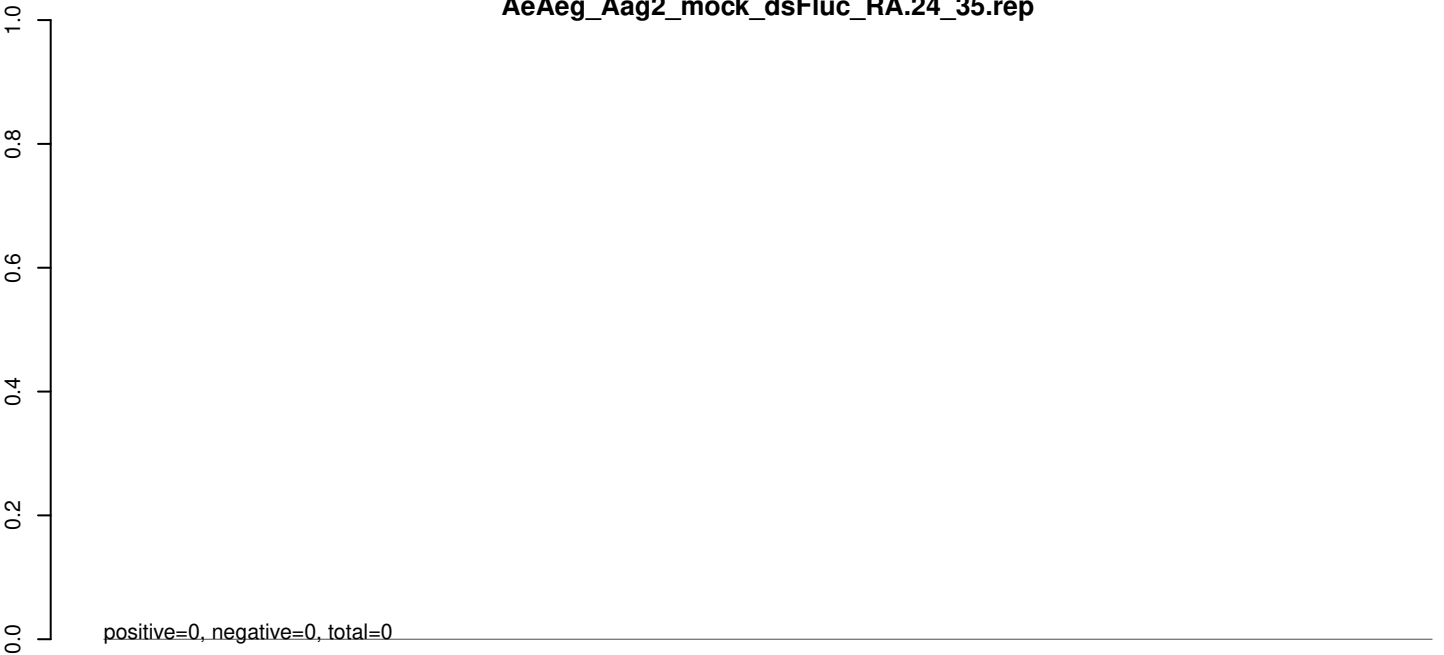
positive=0, negative=0, total=0, width=25, length=10688, virus@NC_023439.1-Kama_virus_strain_LEIV-20776Tat_polyprotein_gene:1-10688

0 2000 4000 6000 8000 10000

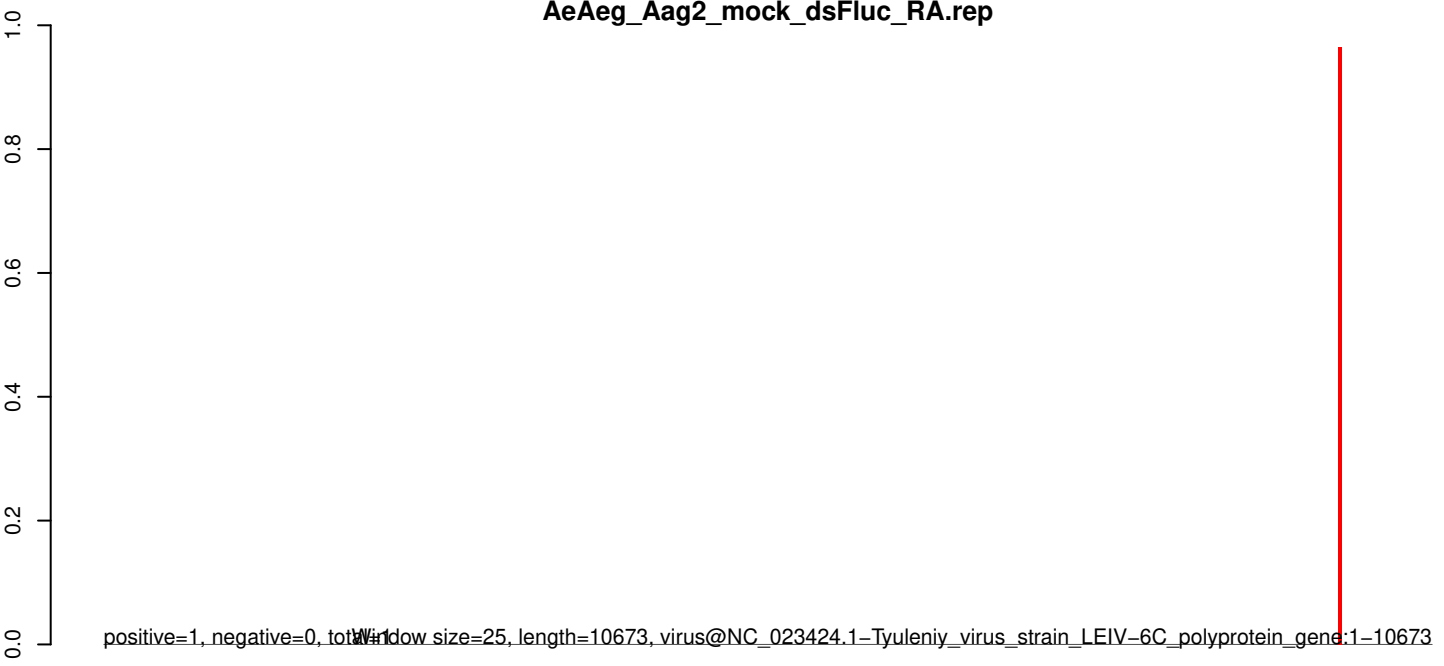
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



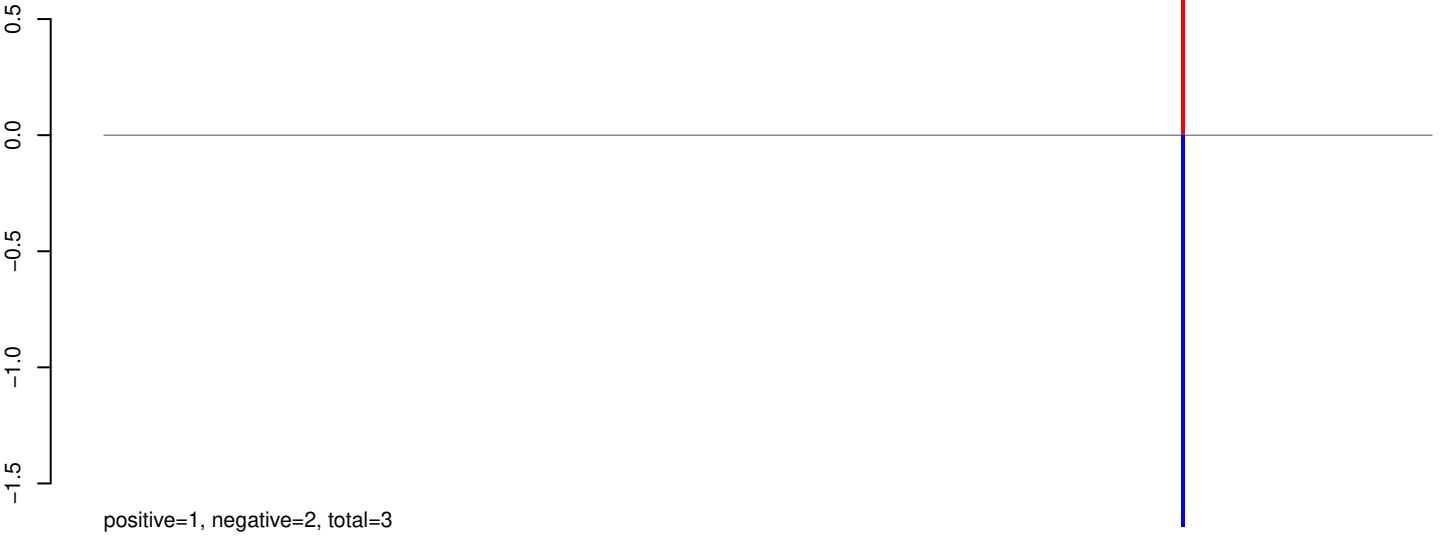
AeAeg_Aag2_mock_dsFluc_RA.rep



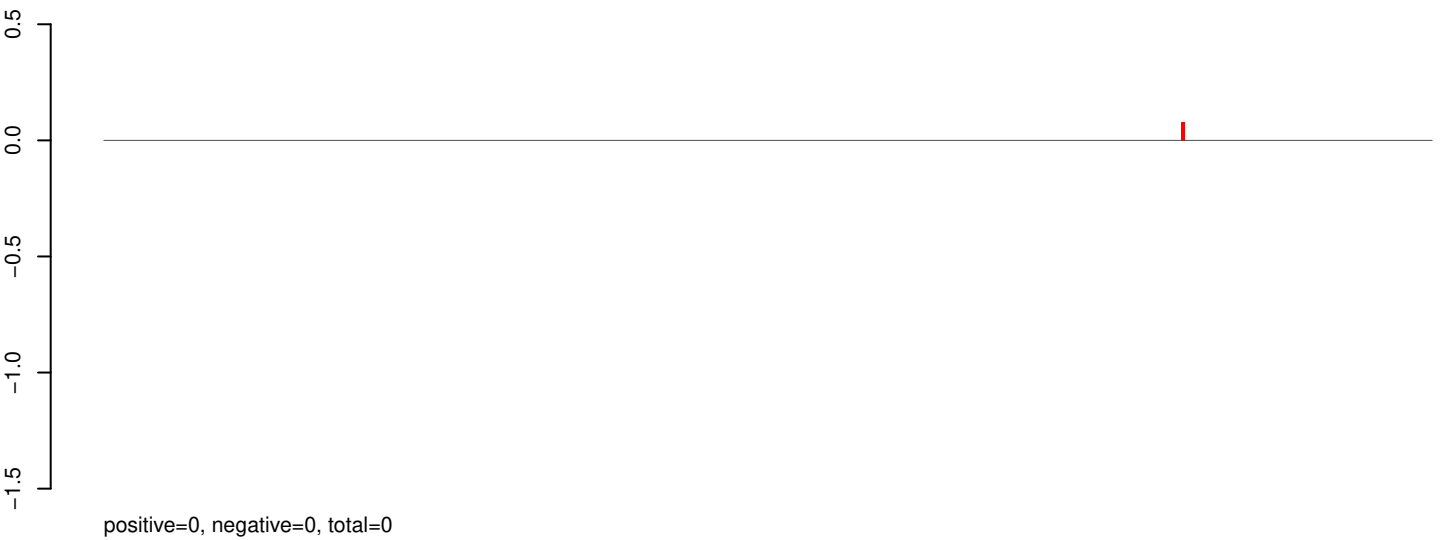
Window size=25, length=10673, virus@NC_023424.1-Tyuleniy_virus_strain_LEIV-6C_polyprotein_gene:1-10673

0 2000 4000 6000 8000 10000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



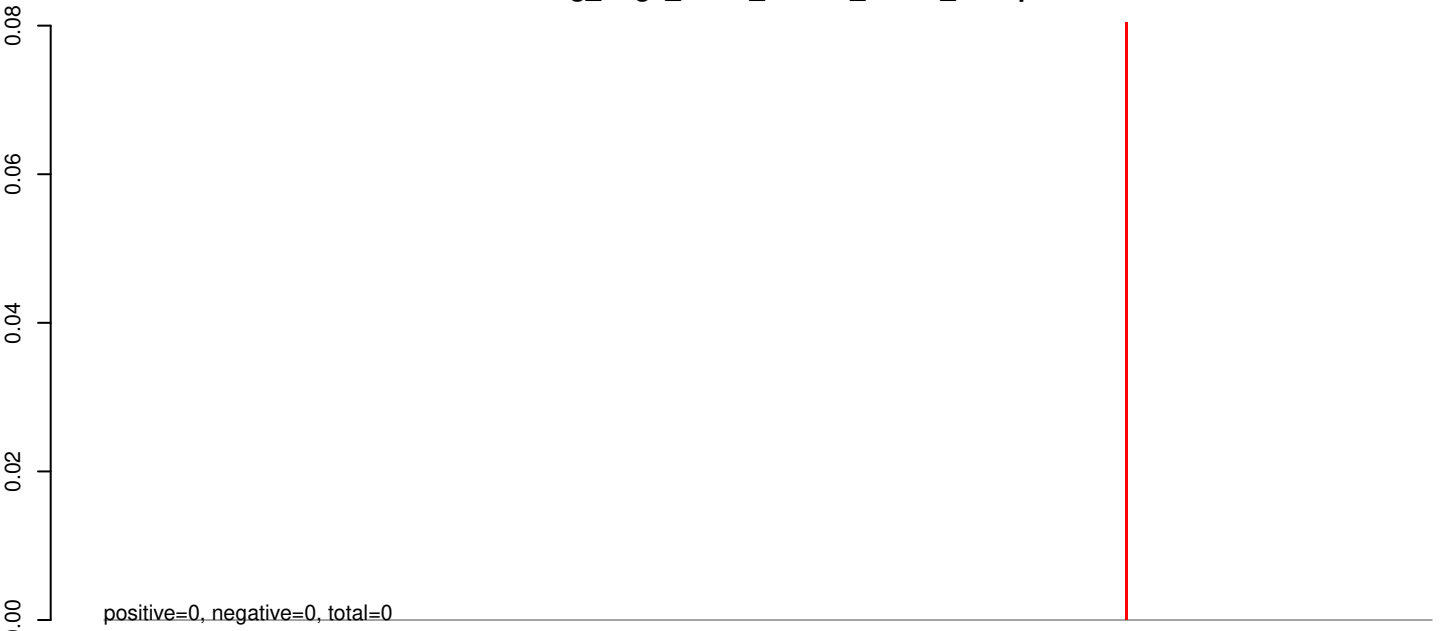
AeAeg_Aag2_mock_dsFluc_RA.rep



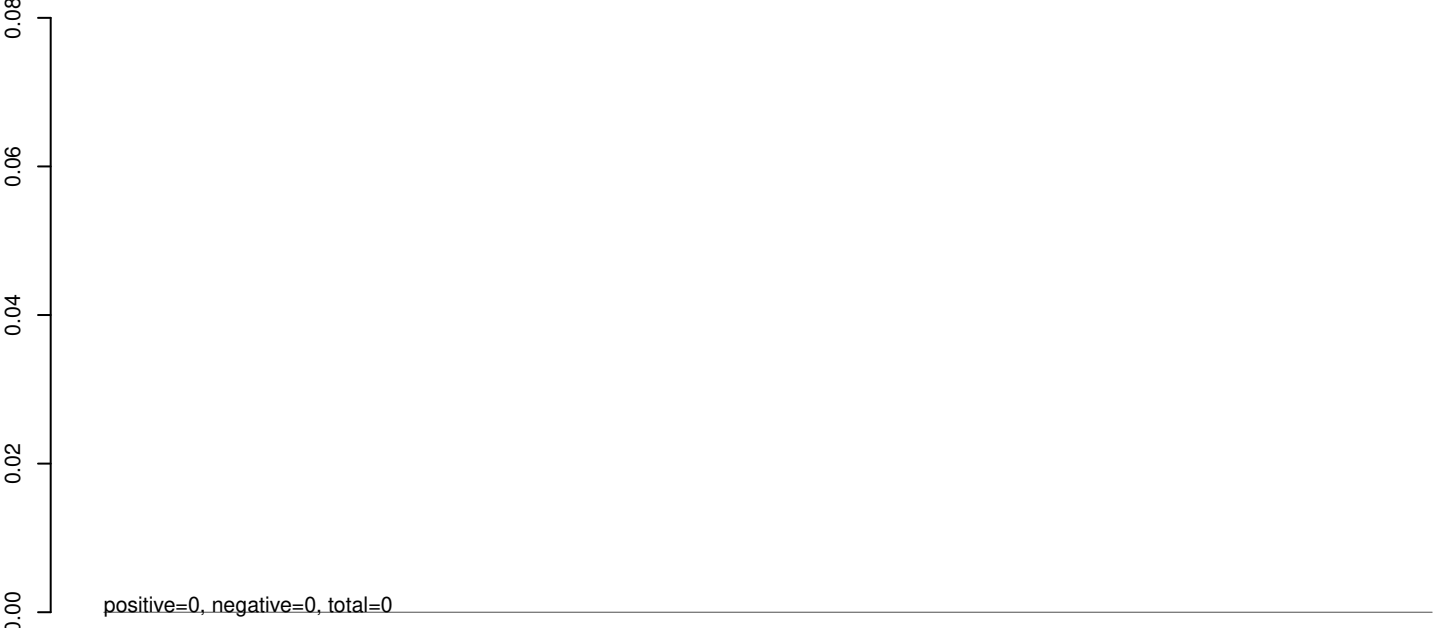
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0 2000 4000 6000 8000 10000

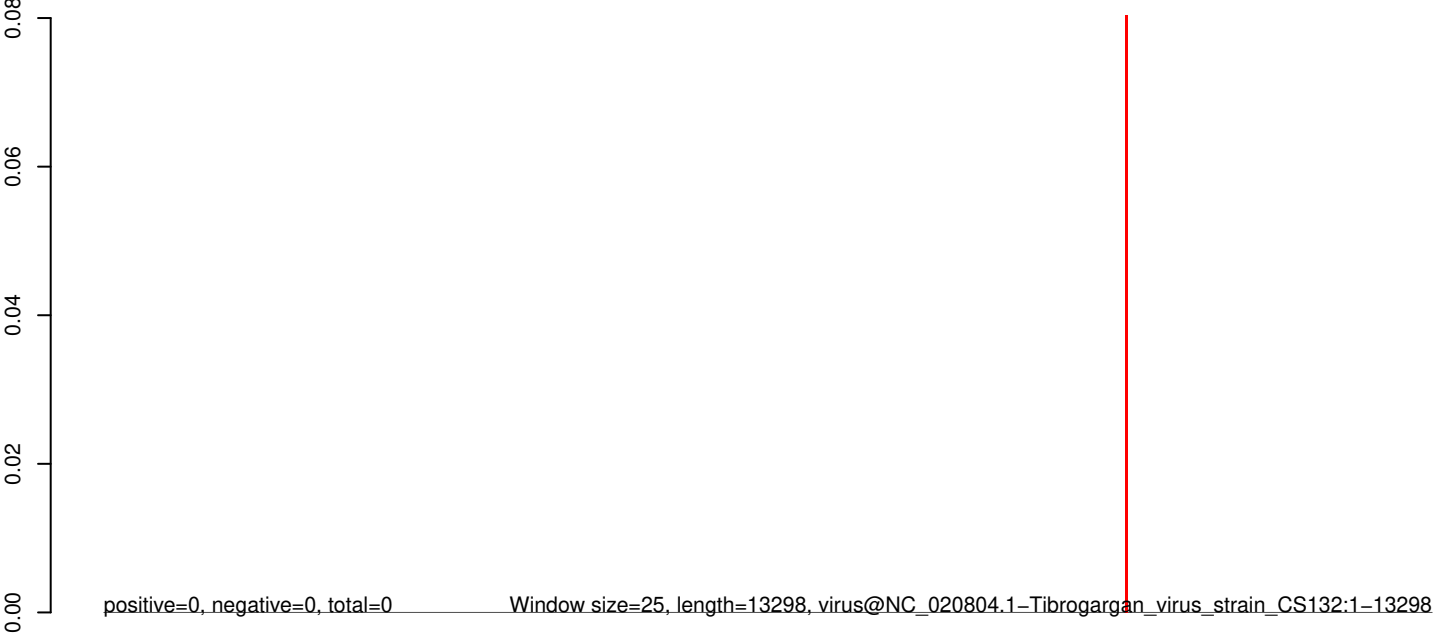
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

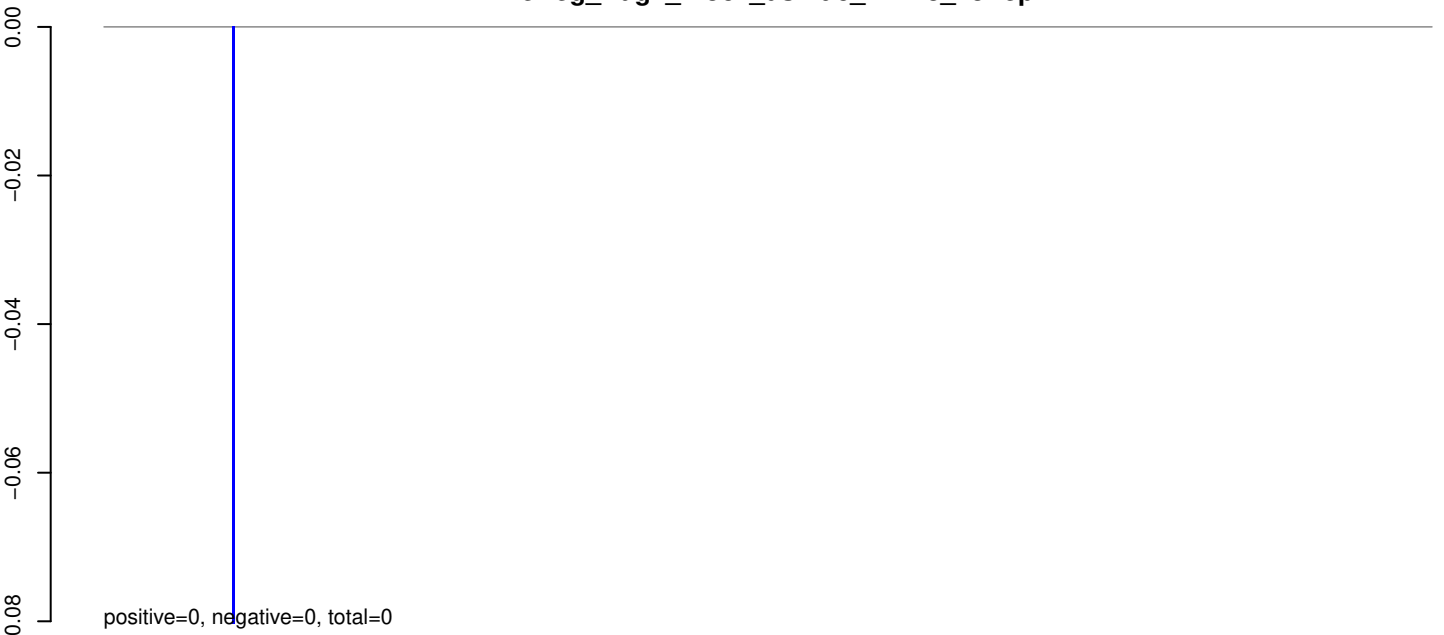


AeAeg_Aag2_mock_dsFluc_RA.rep

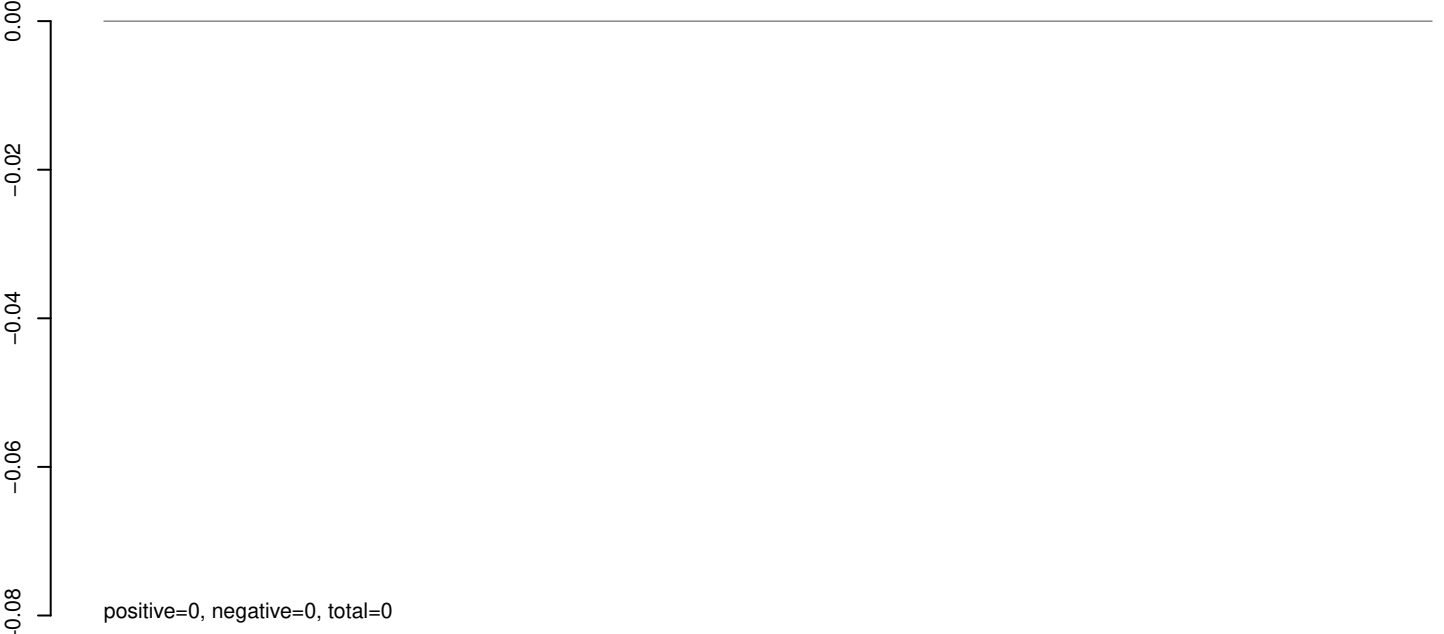


0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



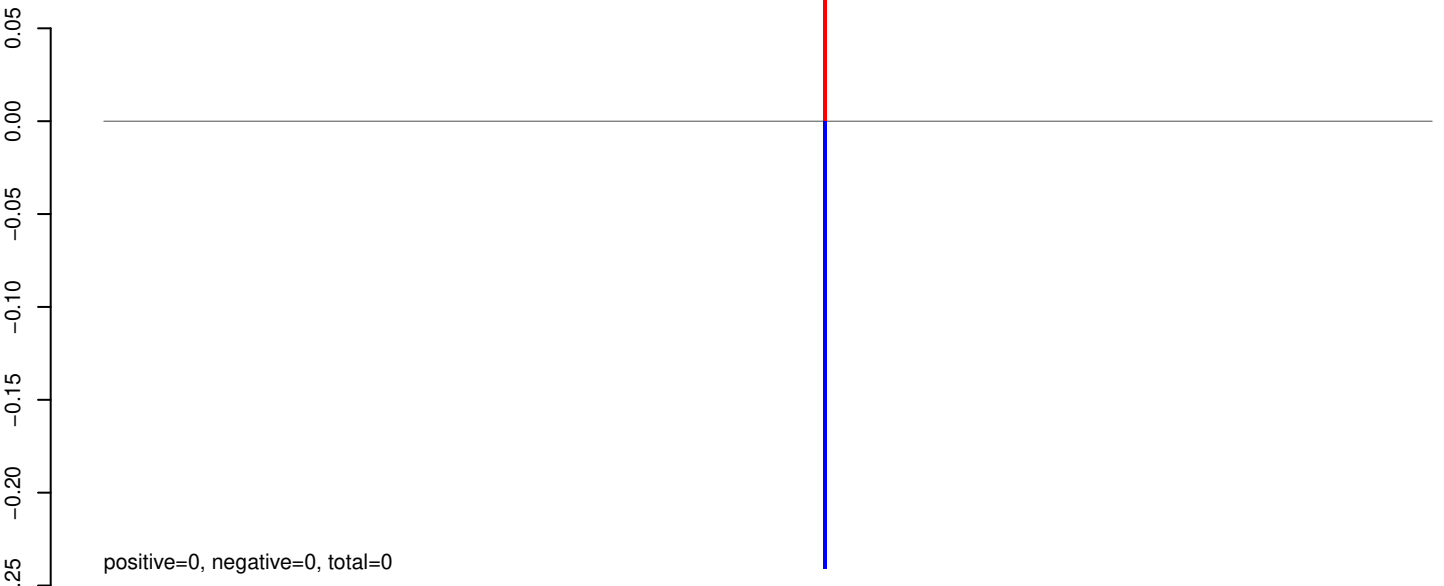
AeAeg_Aag2_mock_dsFluc_RA.rep



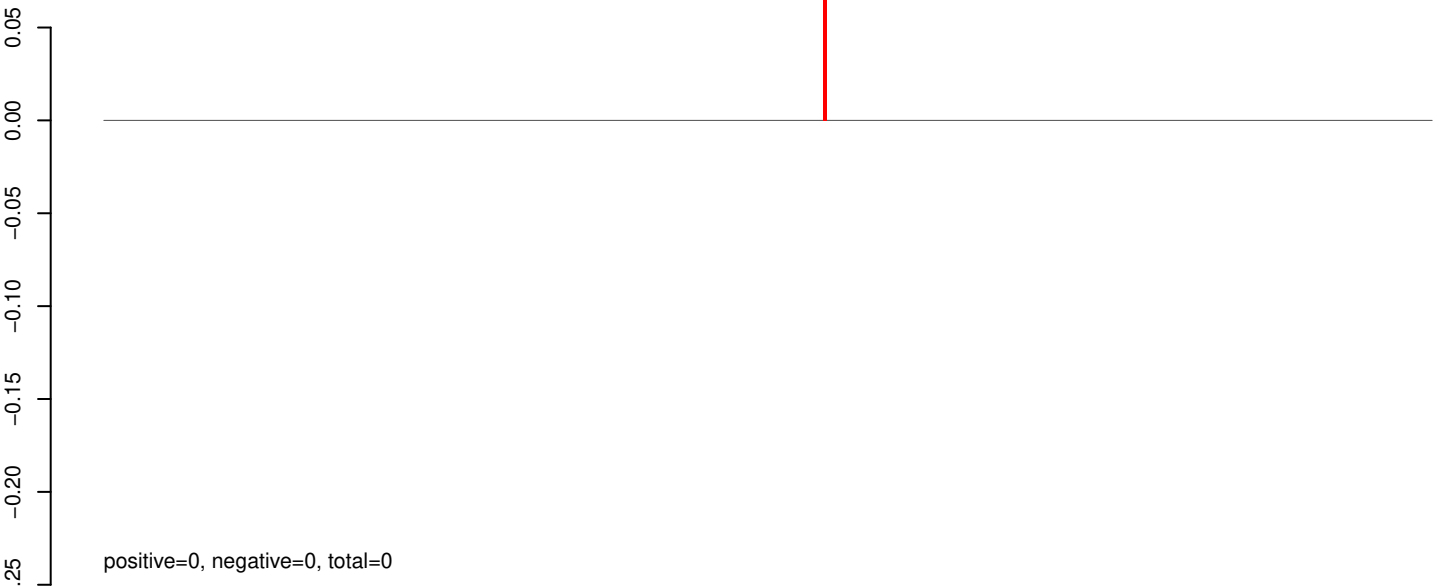
Window size=25, length=14717, virus@NC_017685.1-Obodhiang_virus:1-14717

0 5000 10000 15000

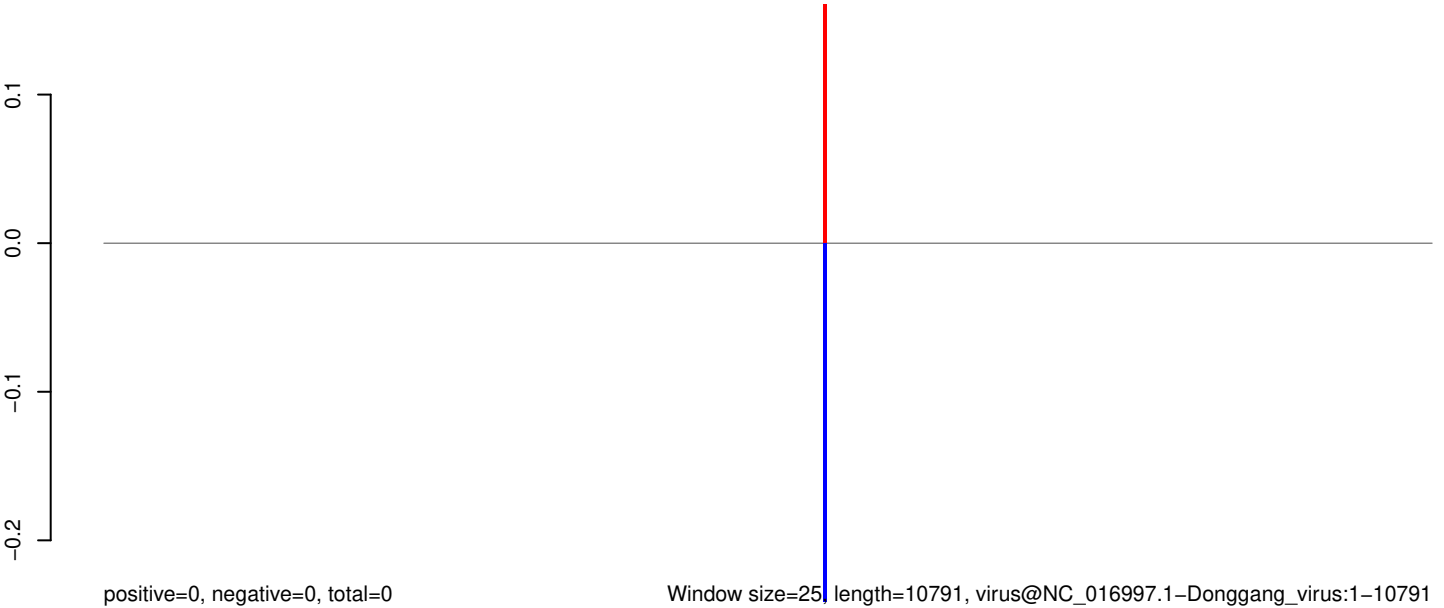
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



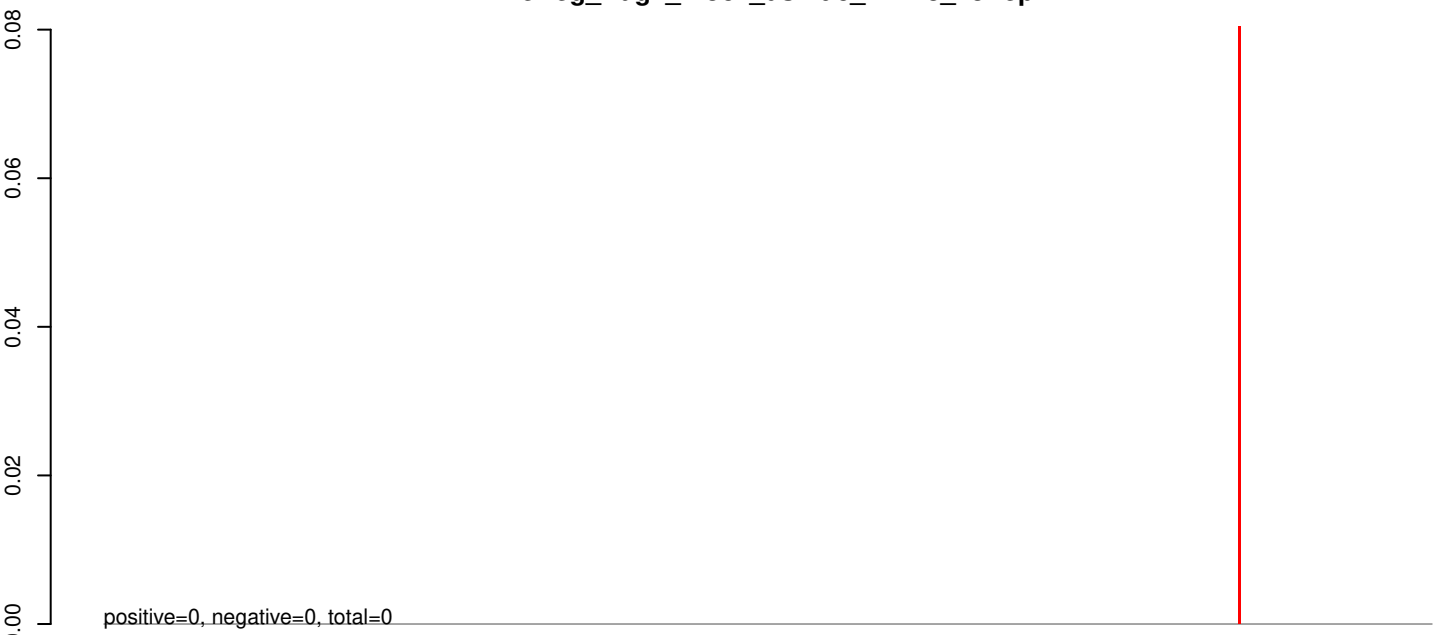
AeAeg_Aag2_mock_dsFluc_RA.rep



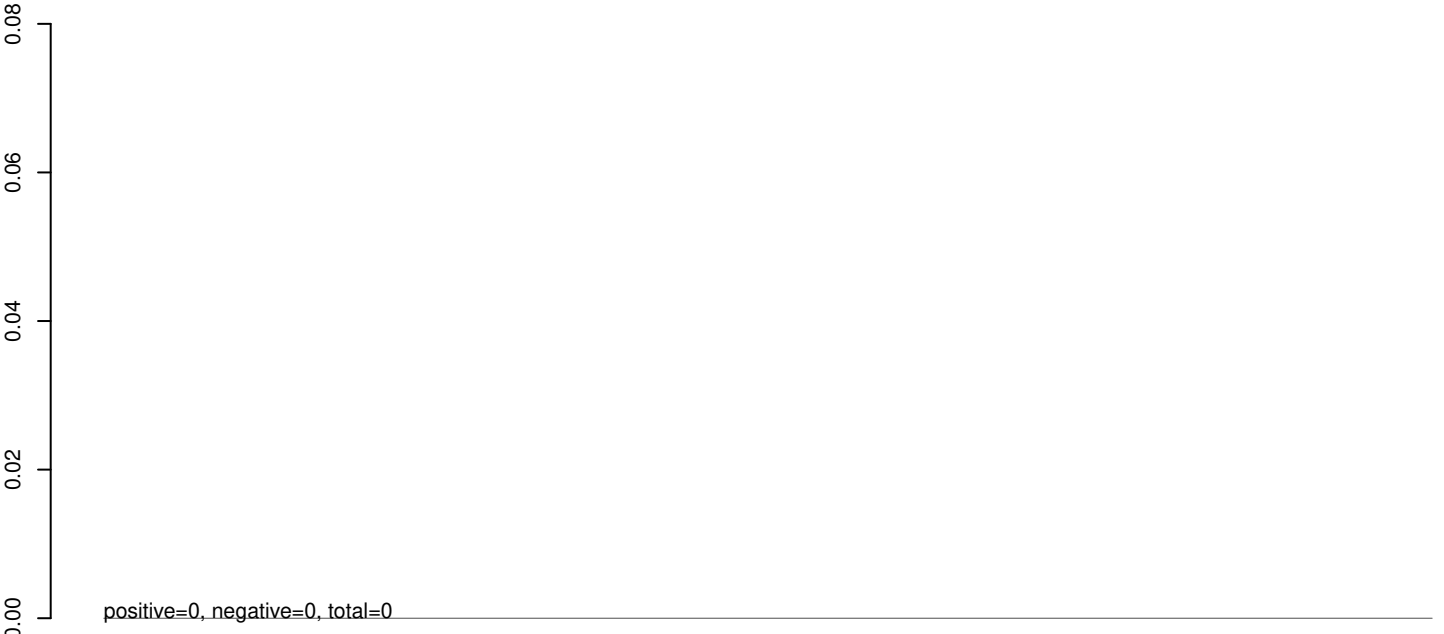
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0 2000 4000 6000 8000 10000

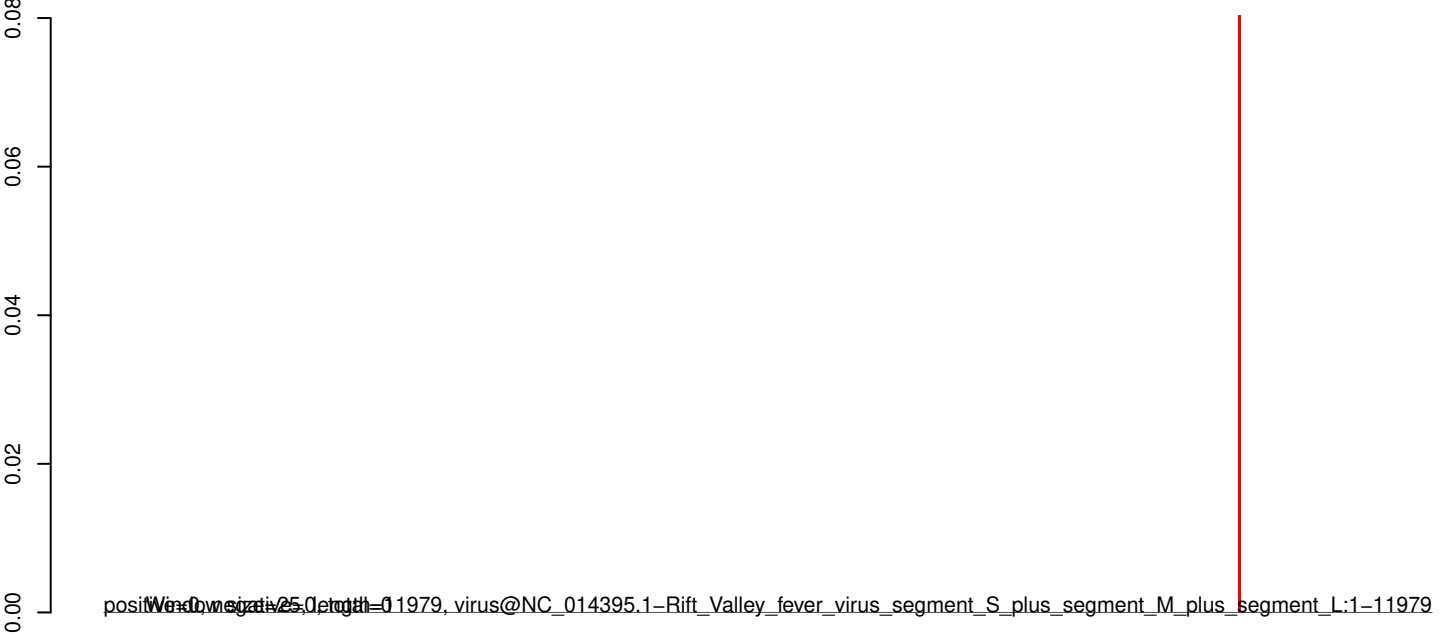
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

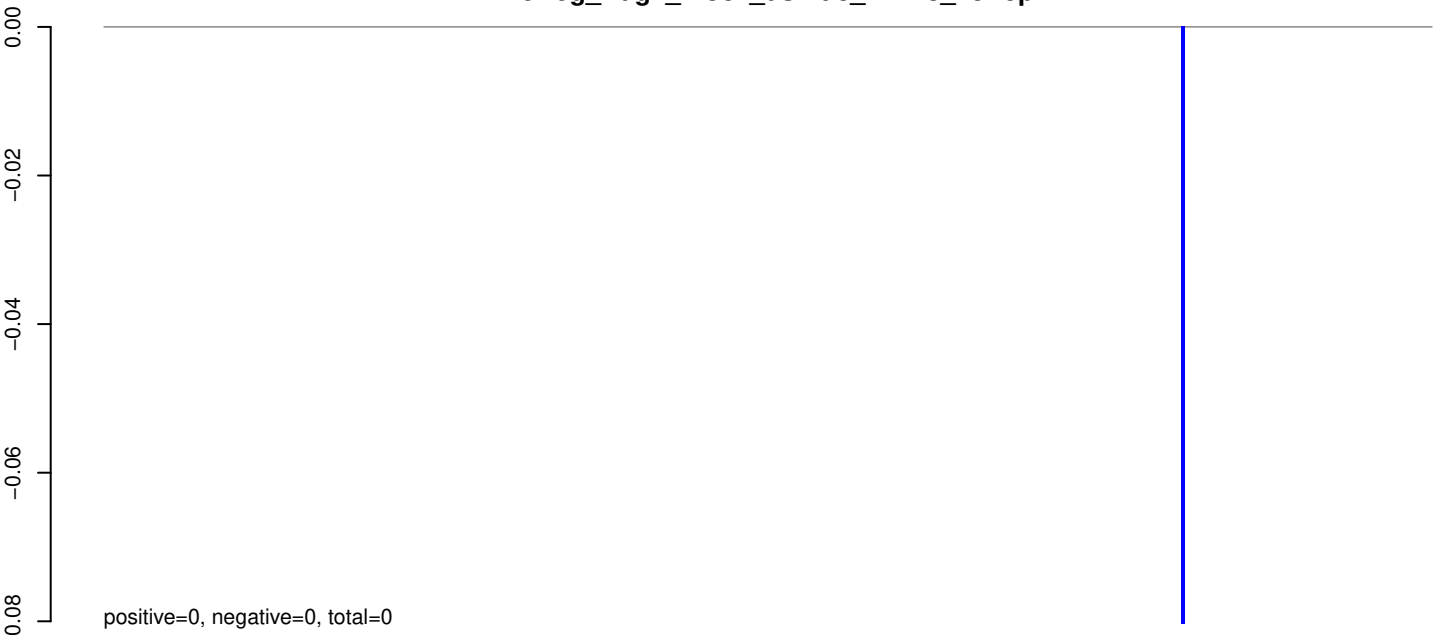


AeAeg_Aag2_mock_dsFluc_RA.rep

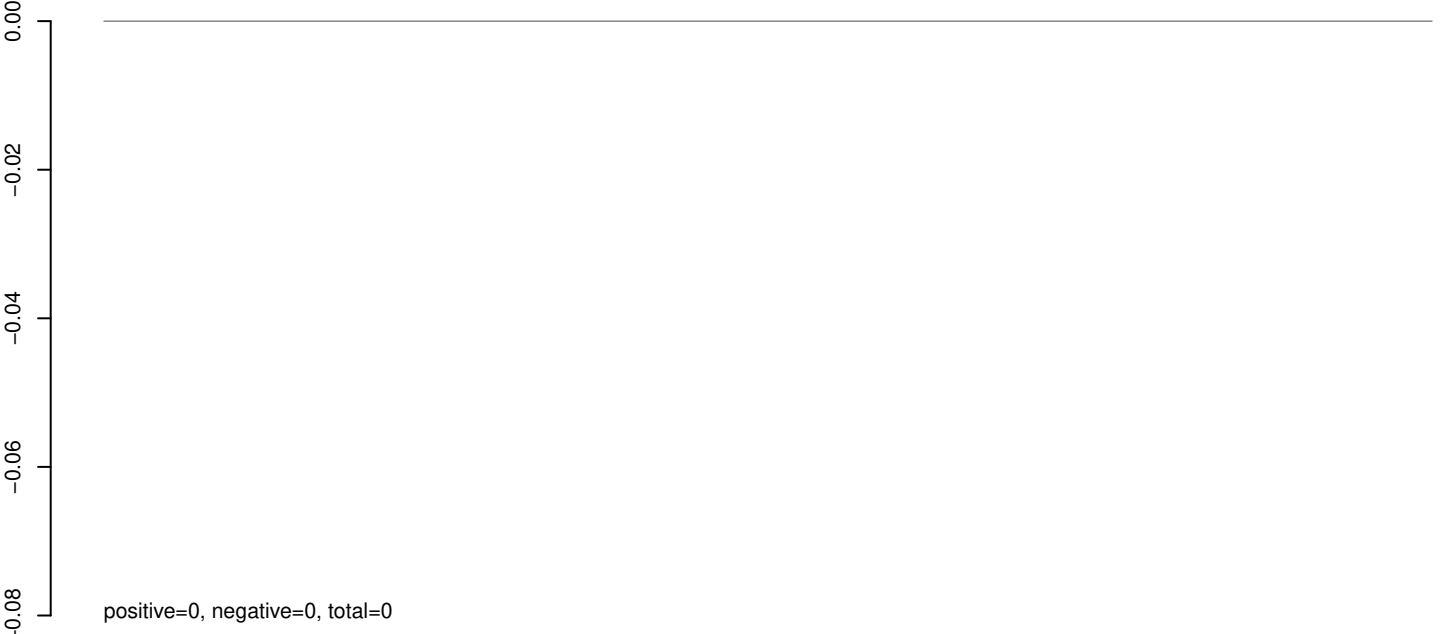


11979, virus@NC_014395.1-Rift_Valley_fever_virus_segment_S_plus_segment_M_plus_segment_L:1-11979

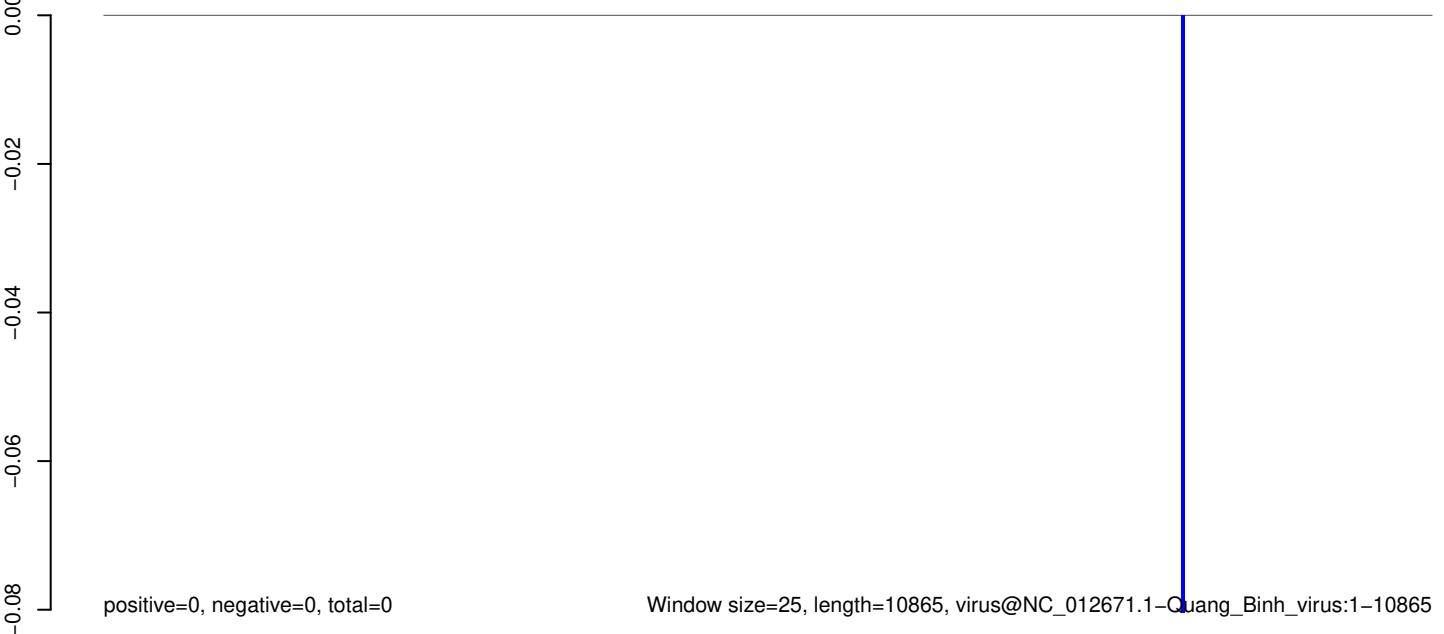
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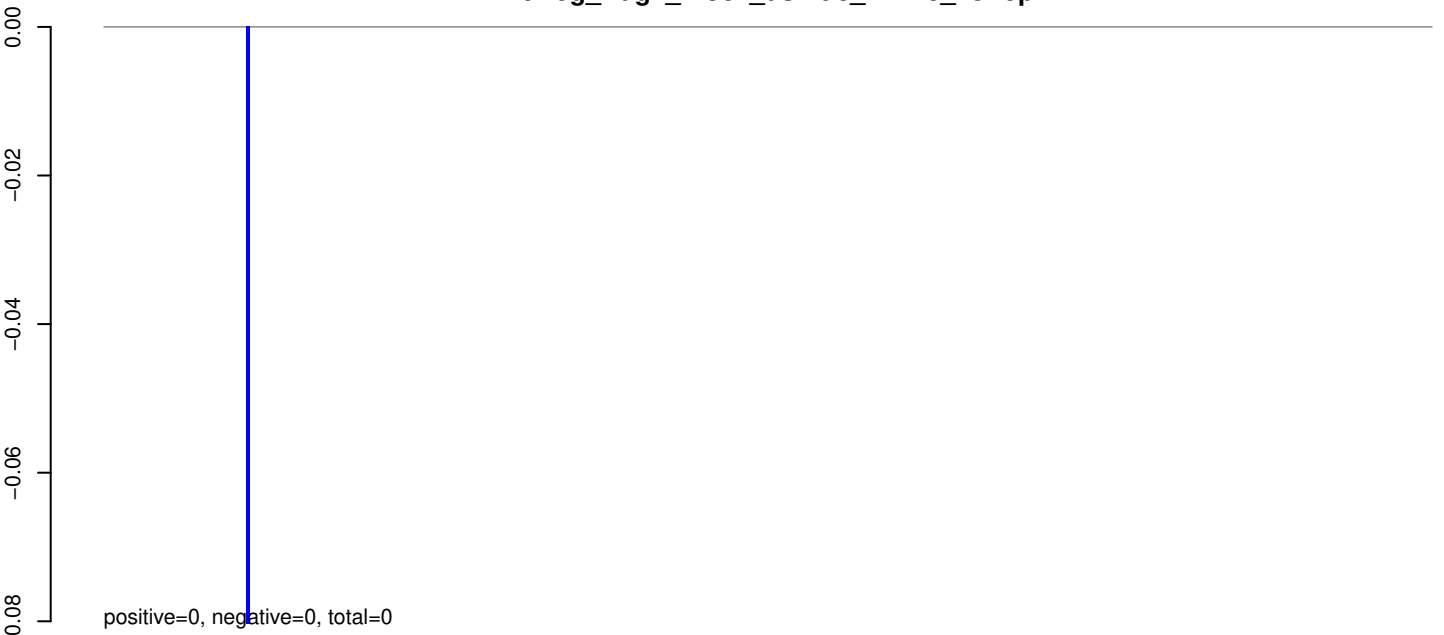
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



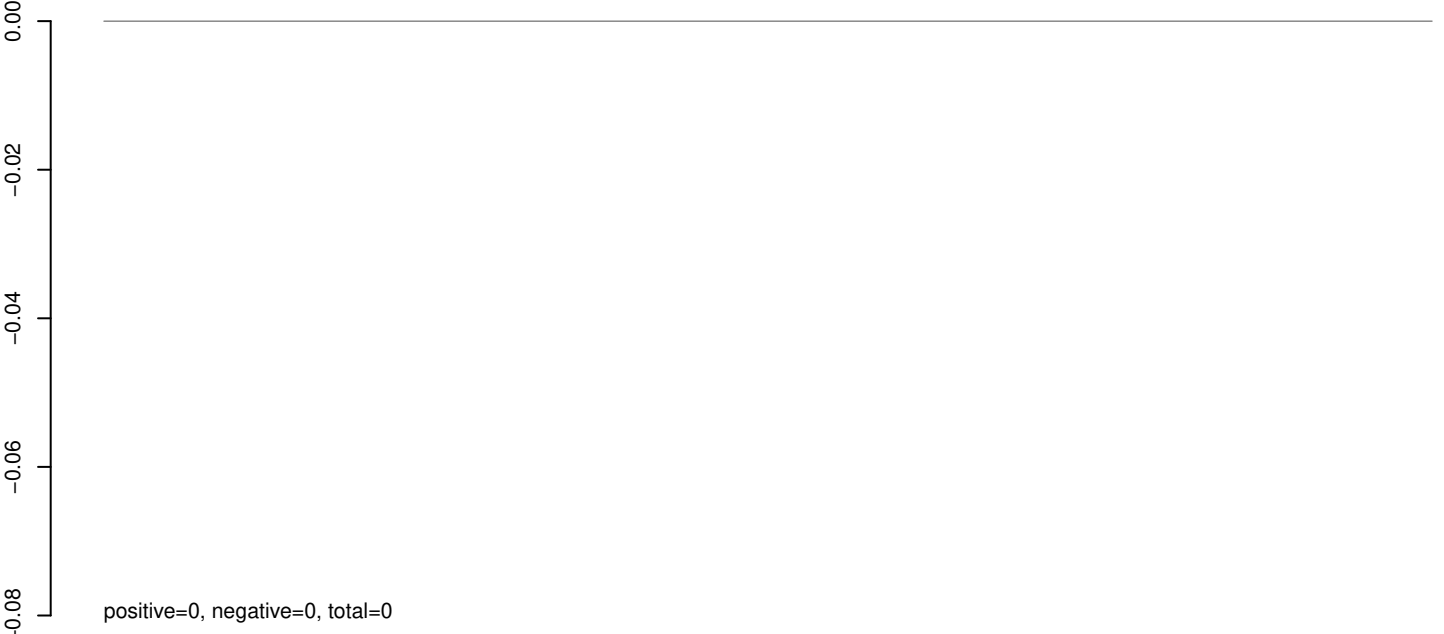
AeAeg_Aag2_mock_dsFluc_RA.rep



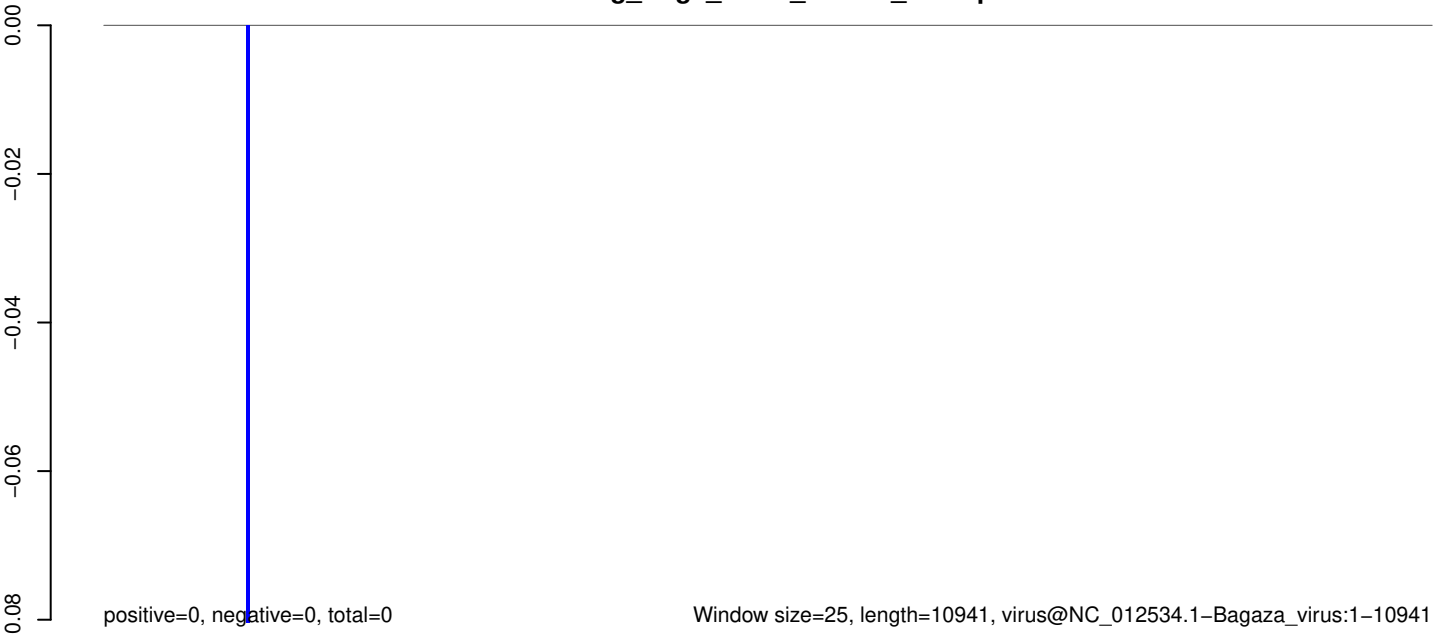
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



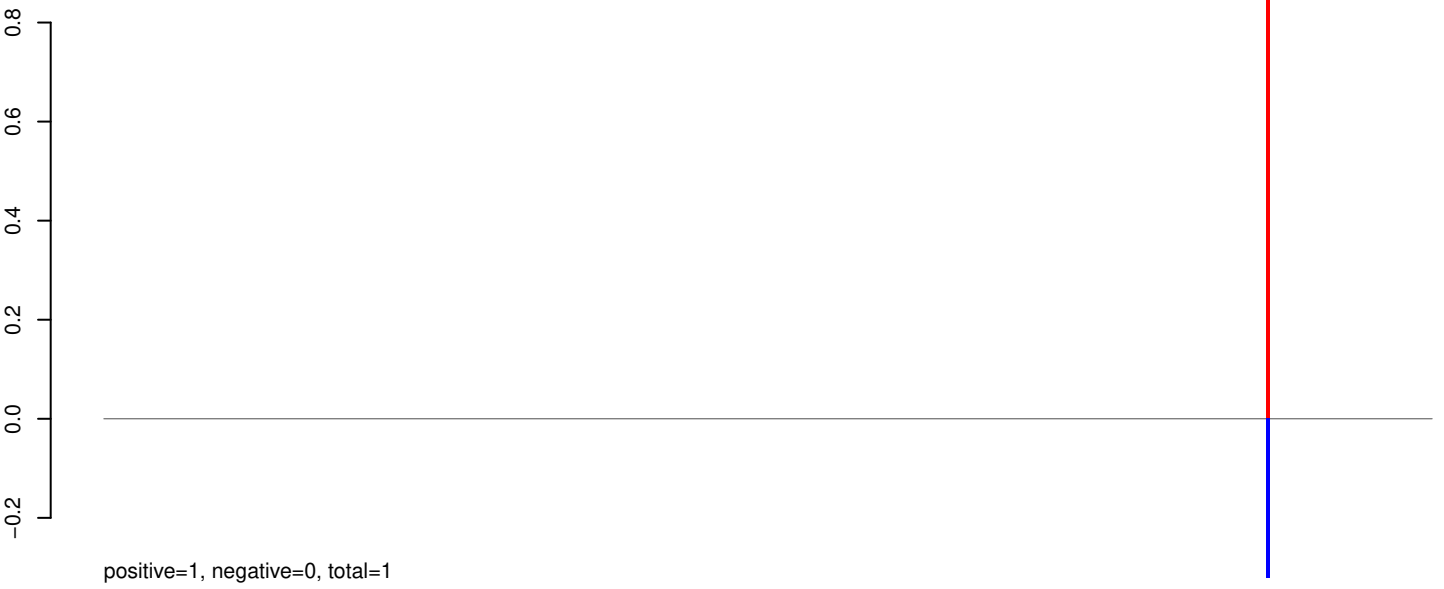
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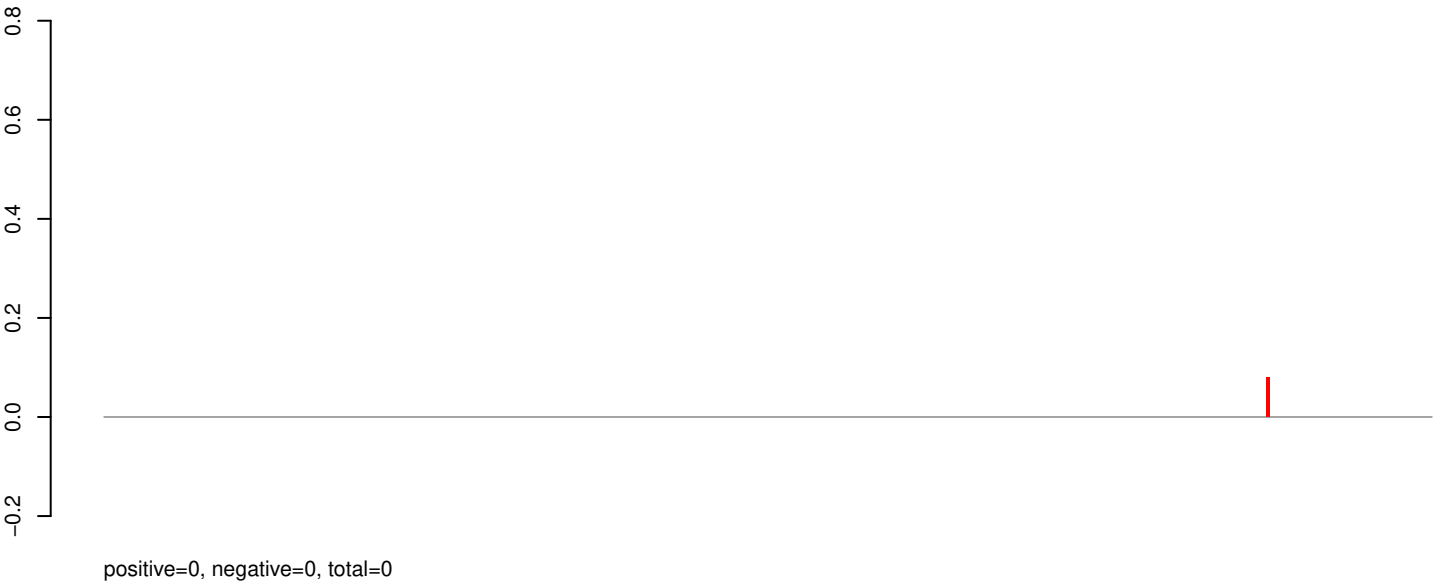
AeAeg_Aag2_mock_dsFluc_RA.rep



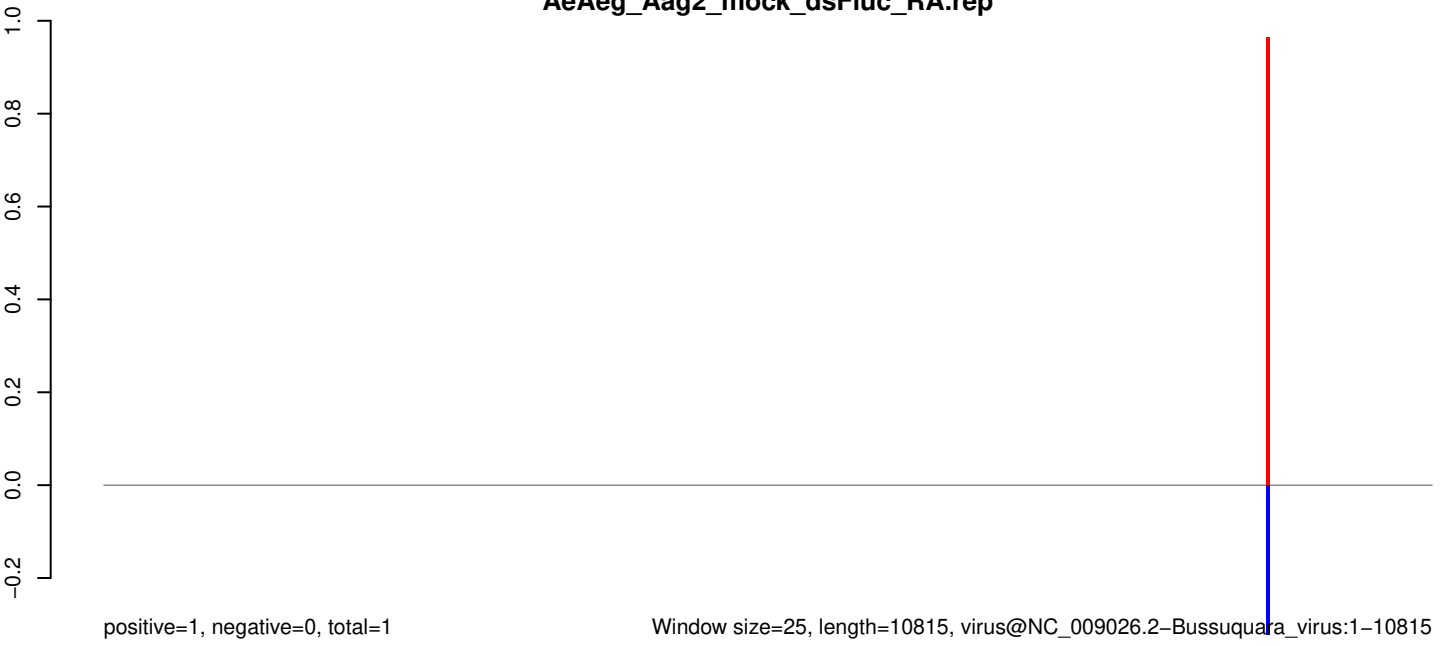
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

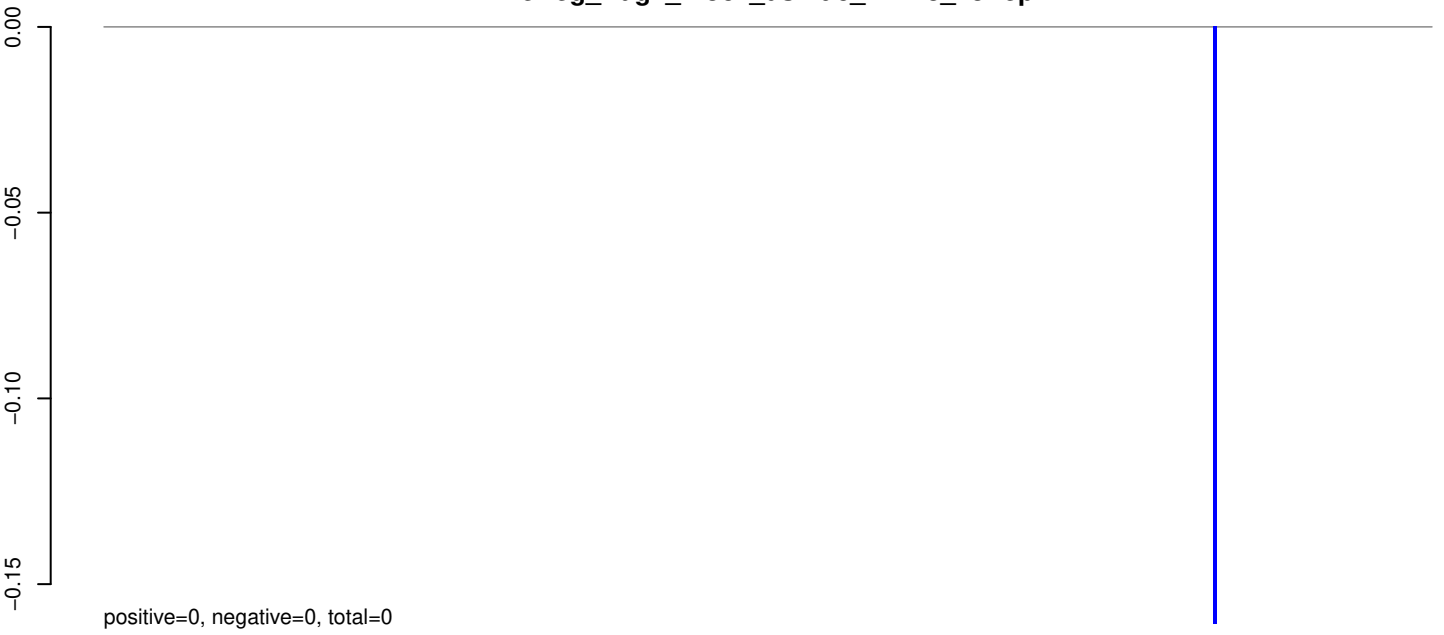


AeAeg_Aag2_mock_dsFluc_RA.rep

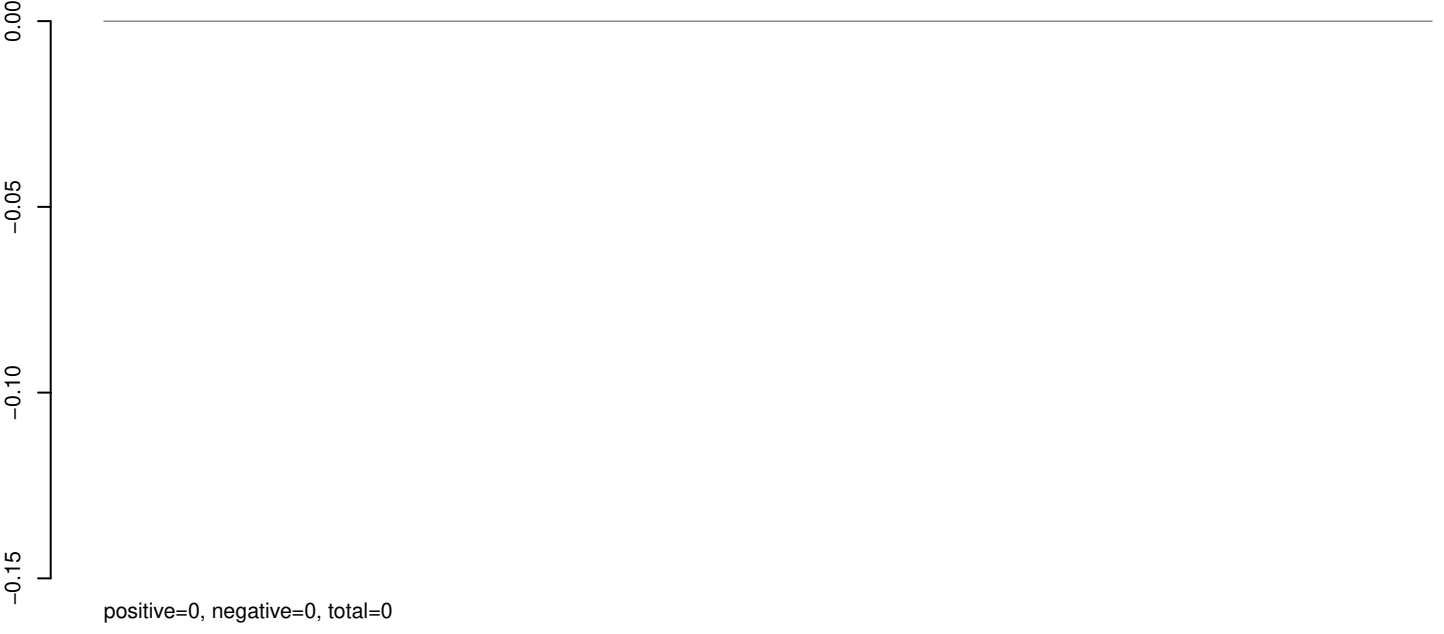


0 2000 4000 6000 8000 10000

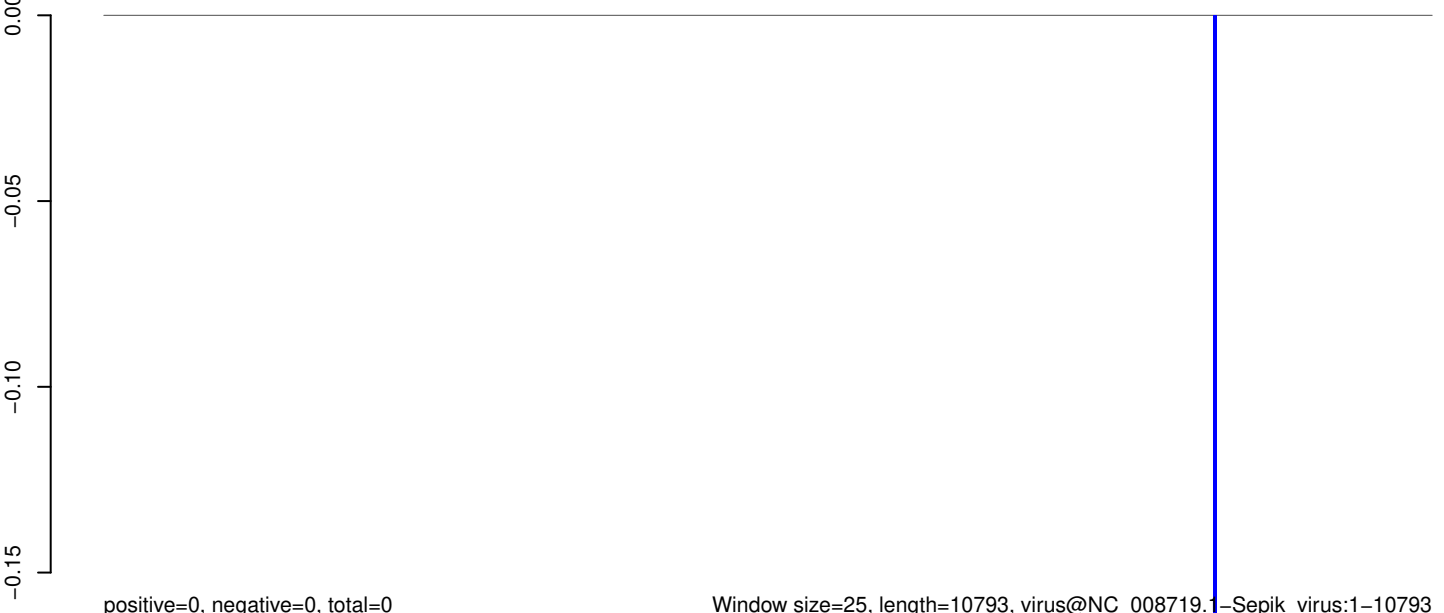
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



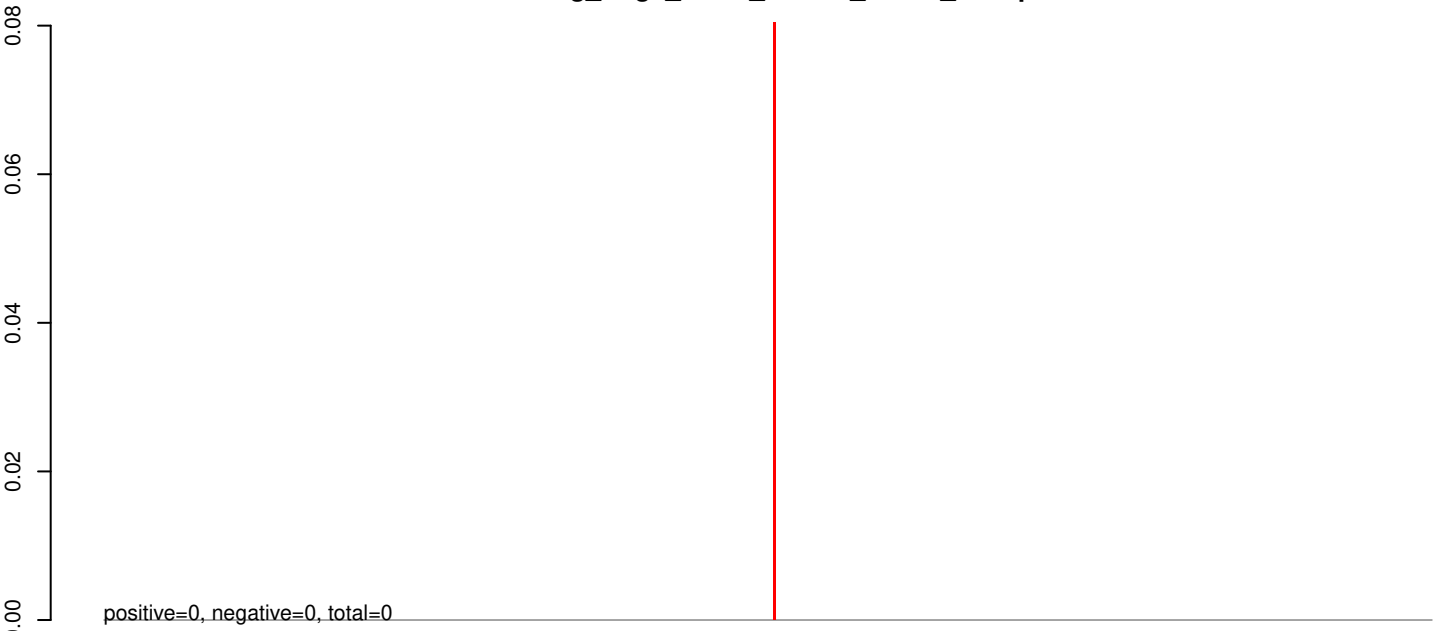
AeAeg_Aag2_mock_dsFluc_RA.rep



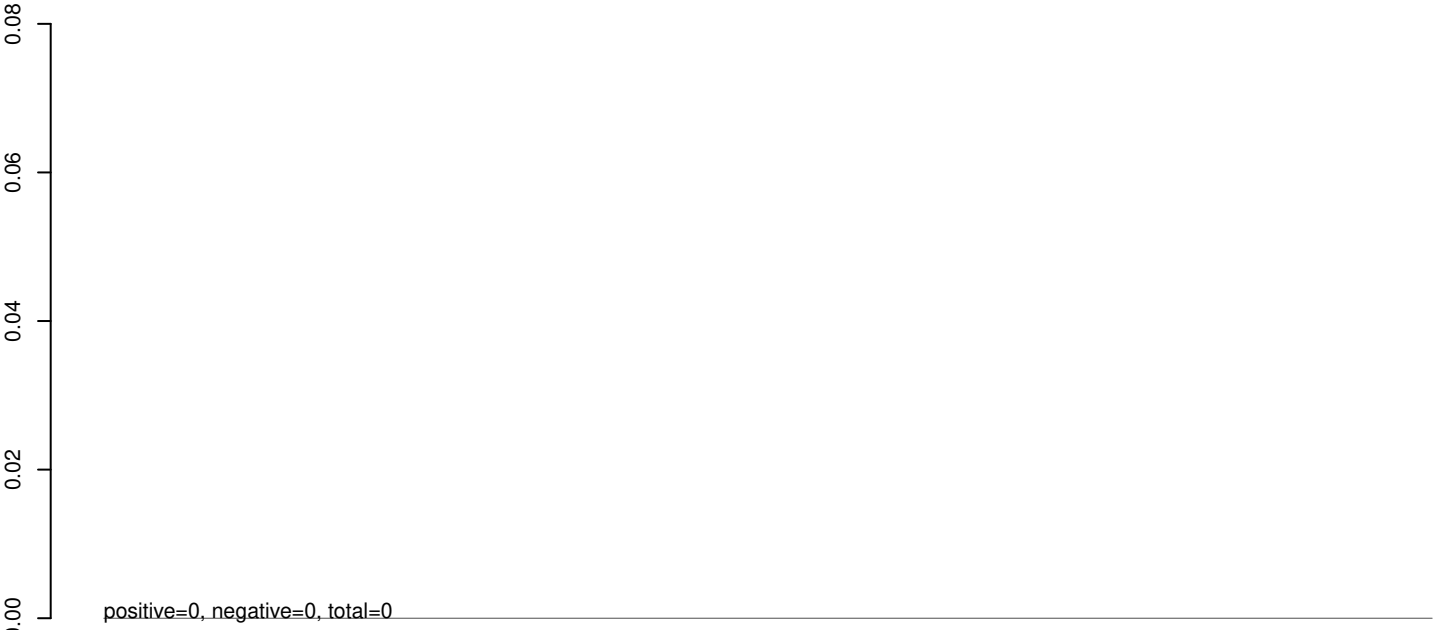
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0 2000 4000 6000 8000 10000

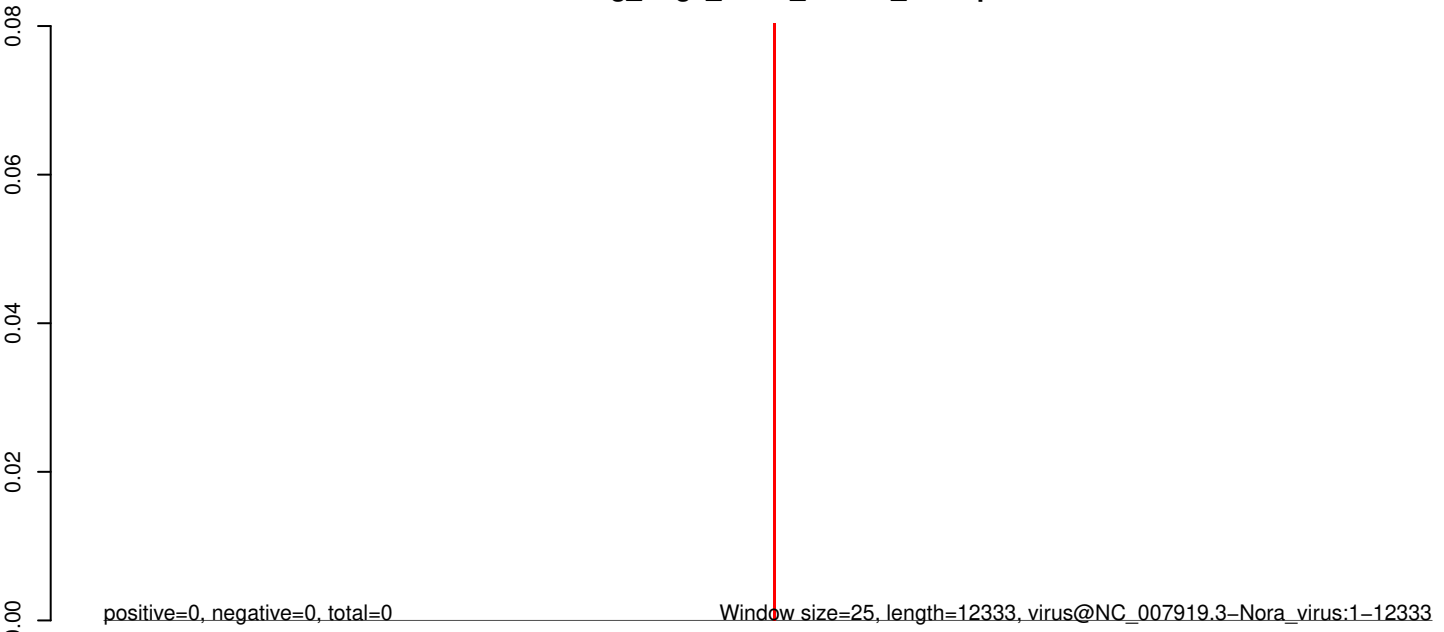
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



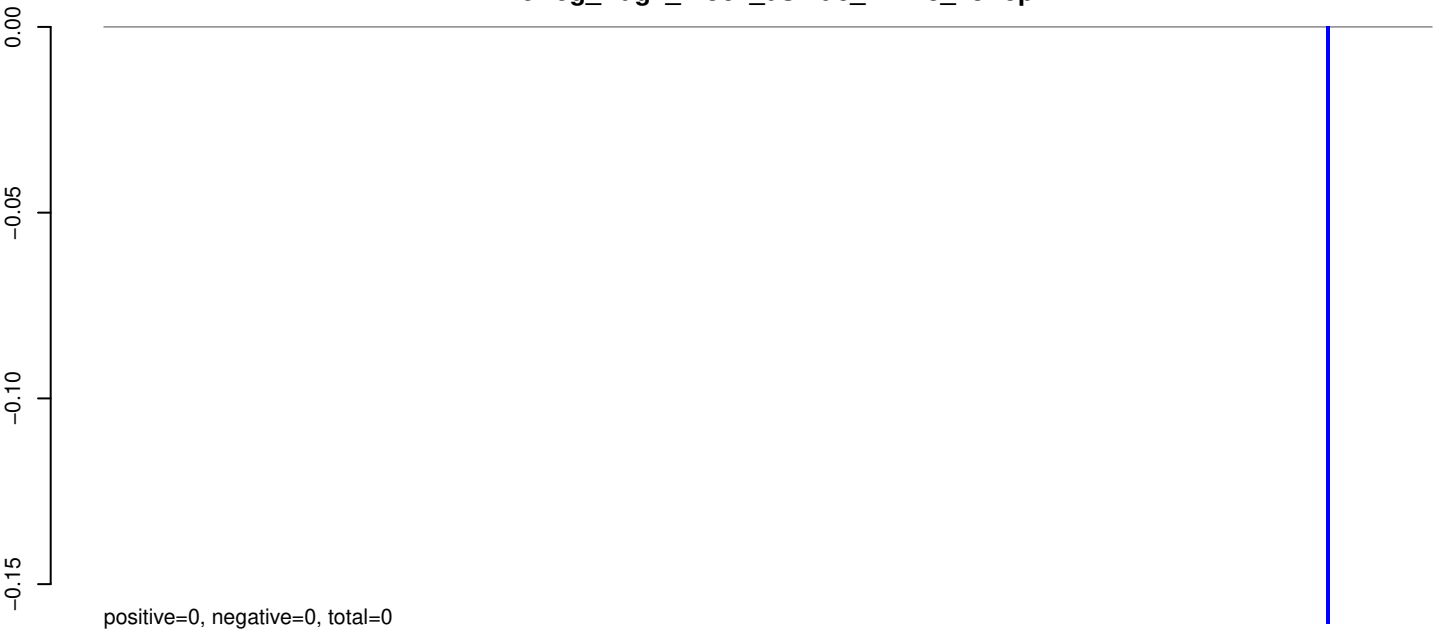
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



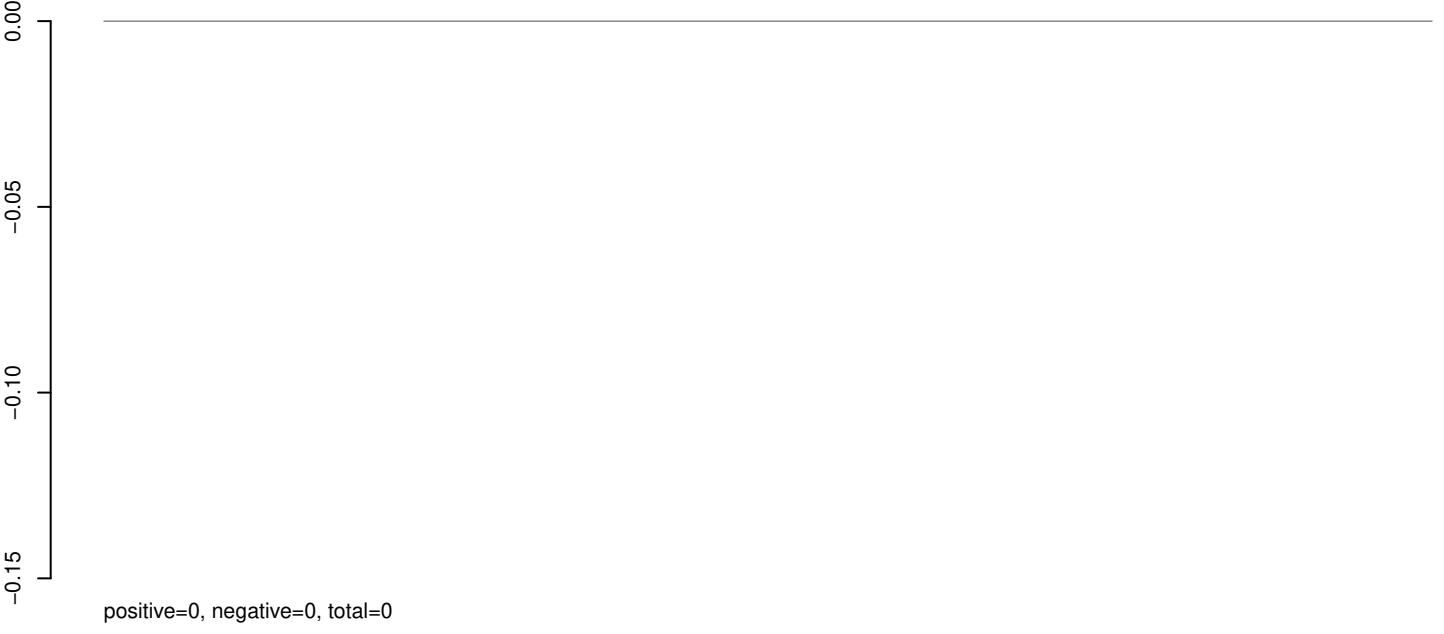
AeAeg_Aag2_mock_dsFluc_RA.rep



AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



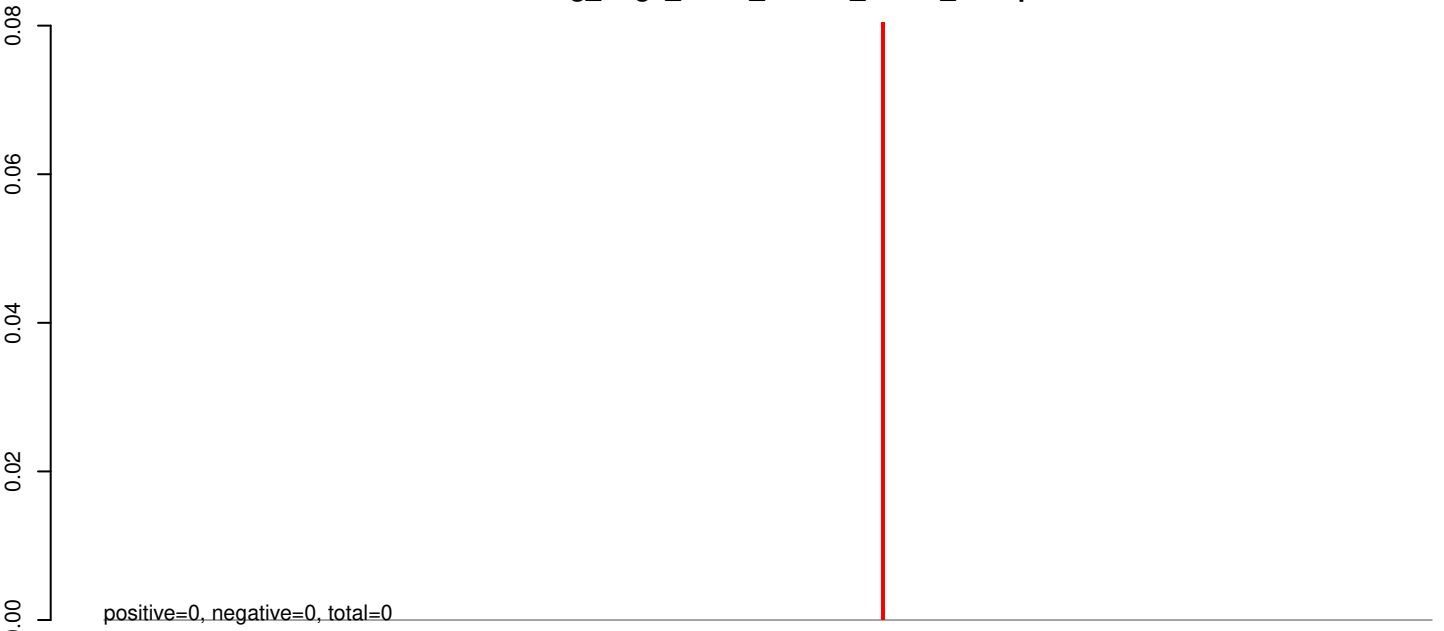
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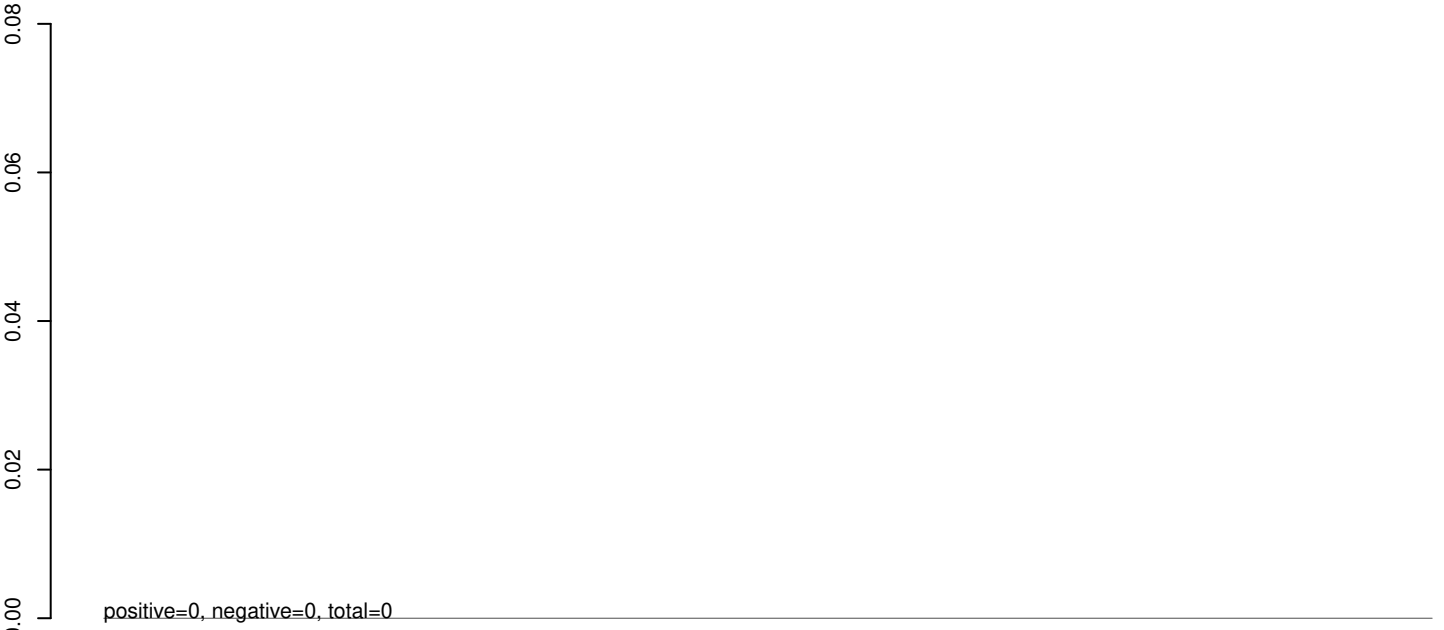
AeAeg_Aag2_mock_dsFluc_RA.rep



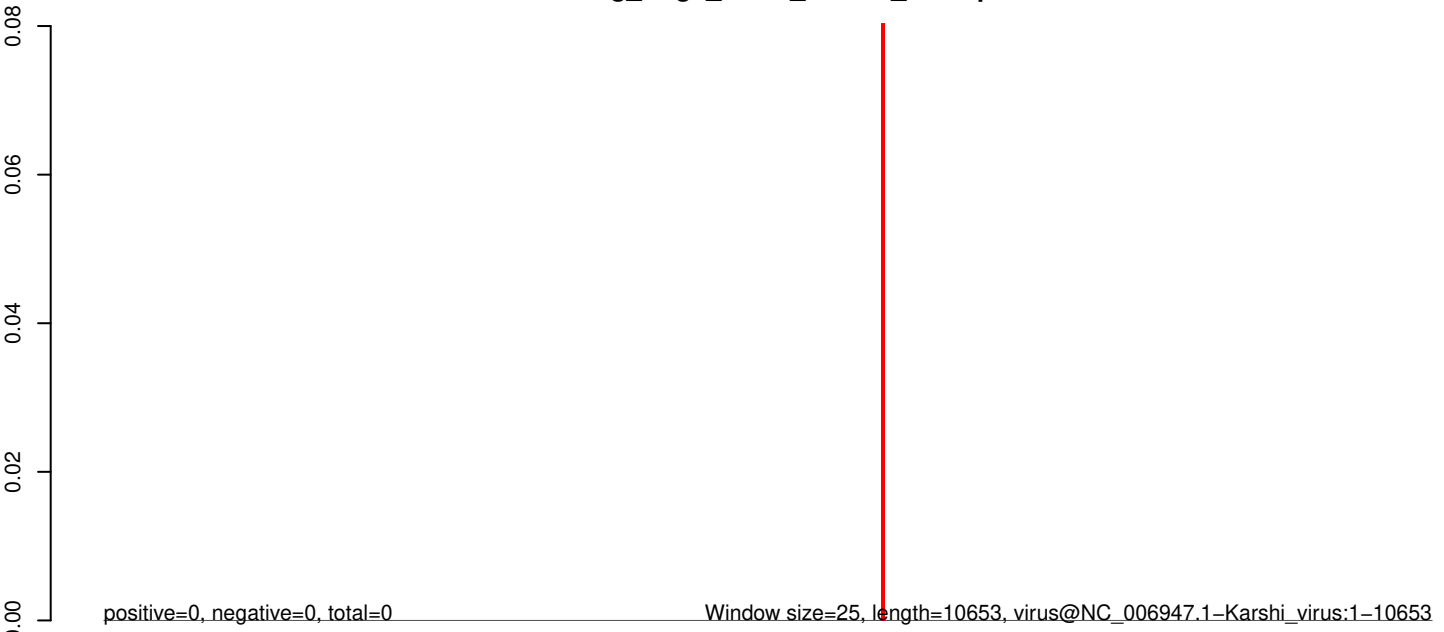
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



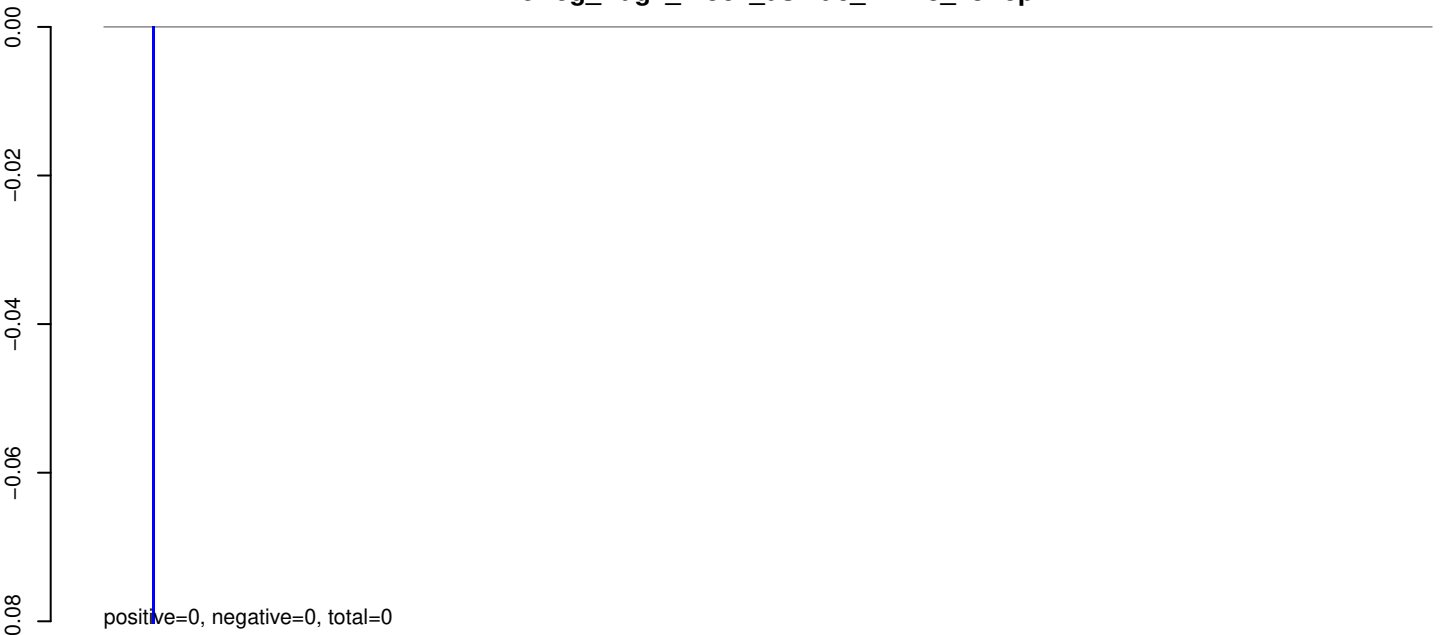
AeAeg_Aag2_mock_dsFluc_RA.rep



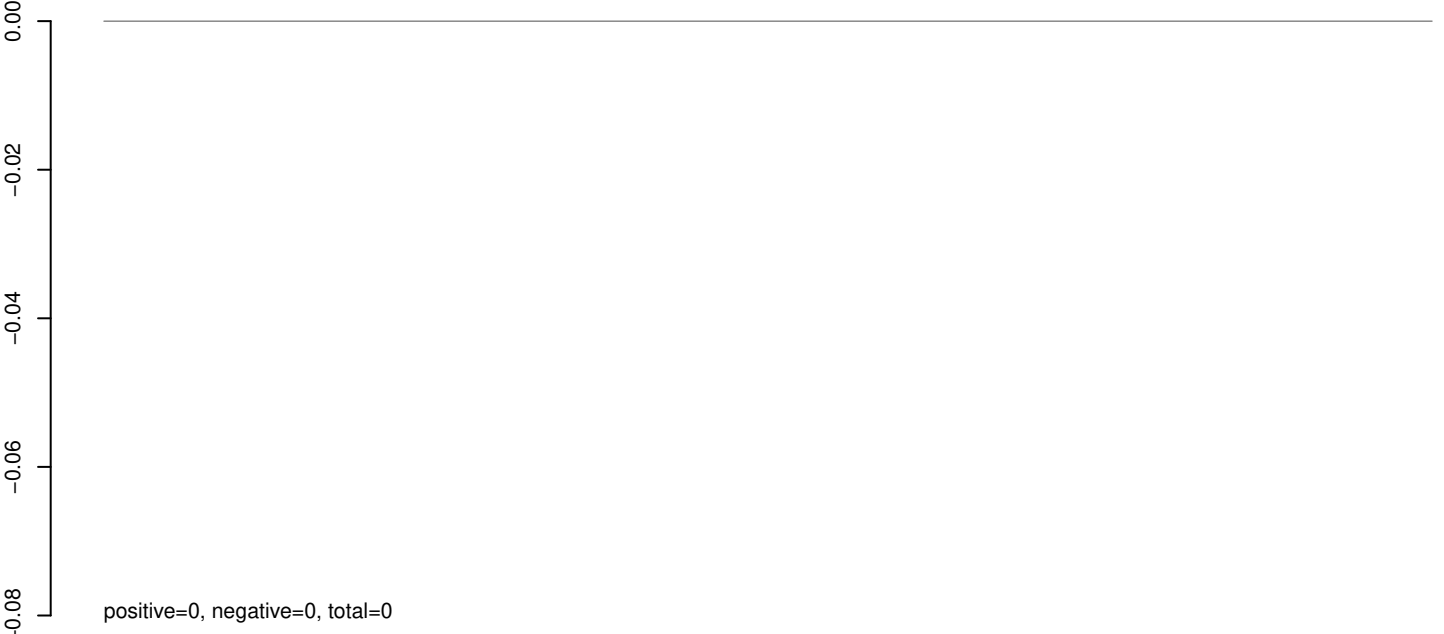
Window size=25, length=10653, virus@NC_006947.1-Karshi_virus:1-10653

0 2000 4000 6000 8000 10000

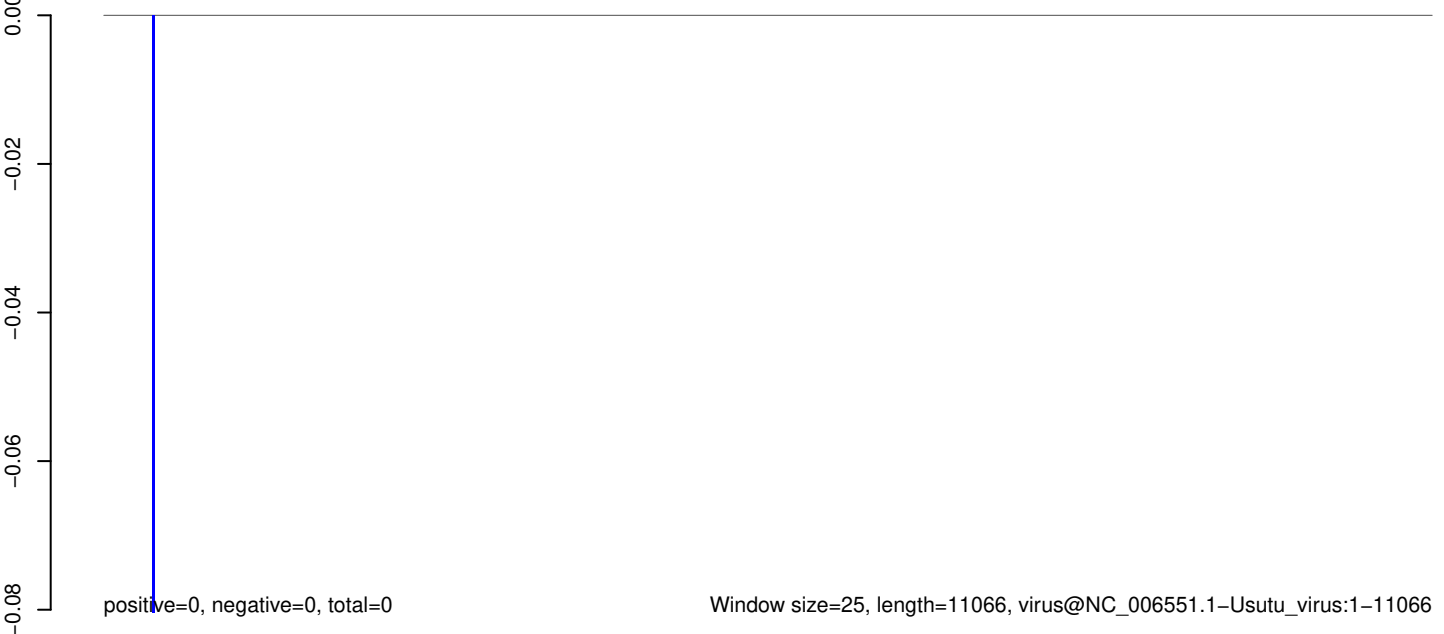
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



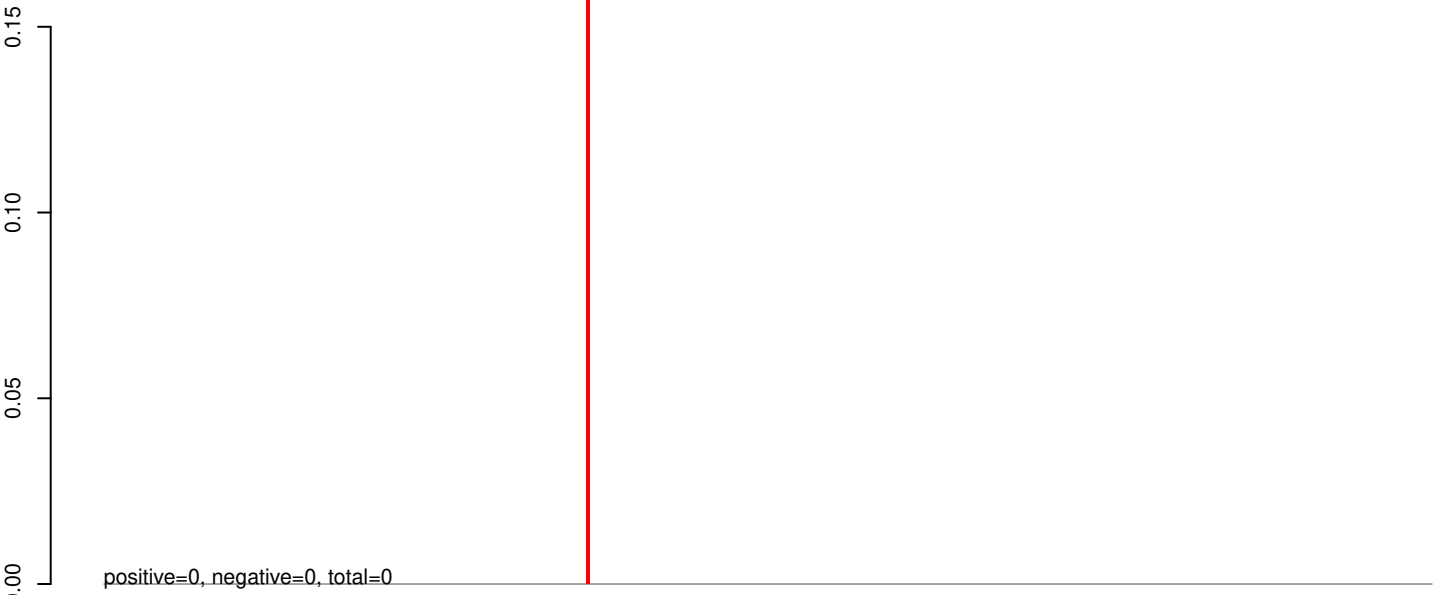
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



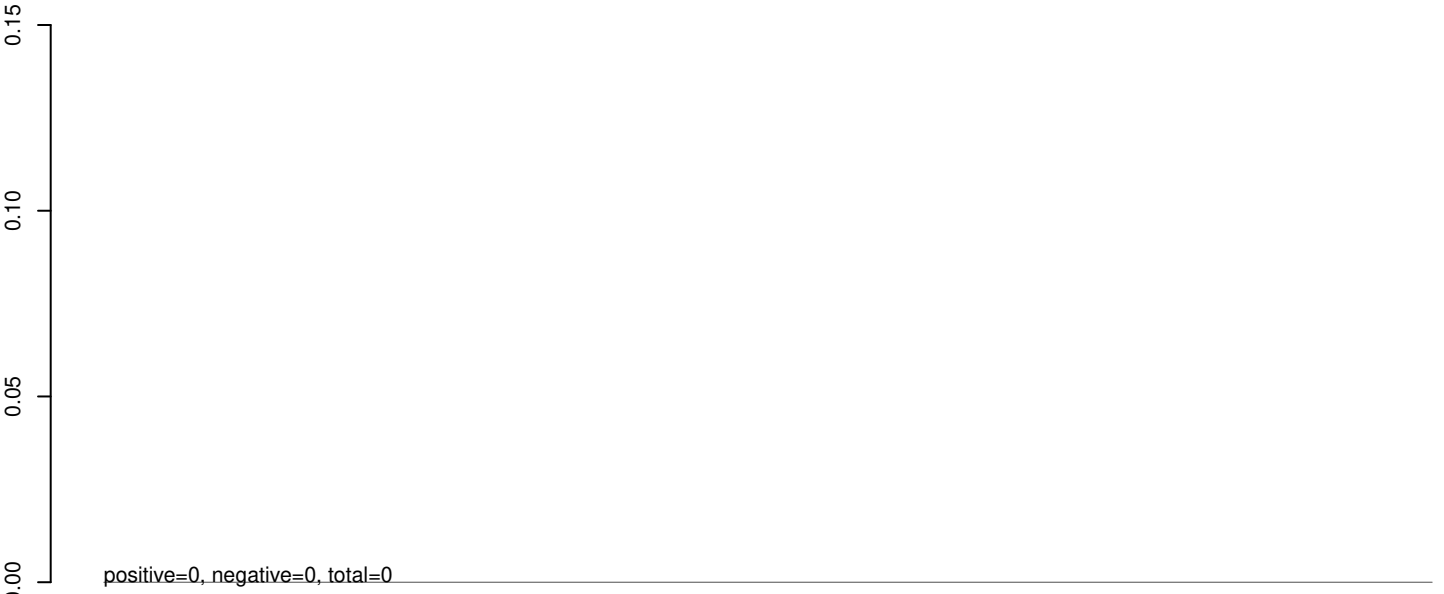
AeAeg_Aag2_mock_dsFluc_RA.rep



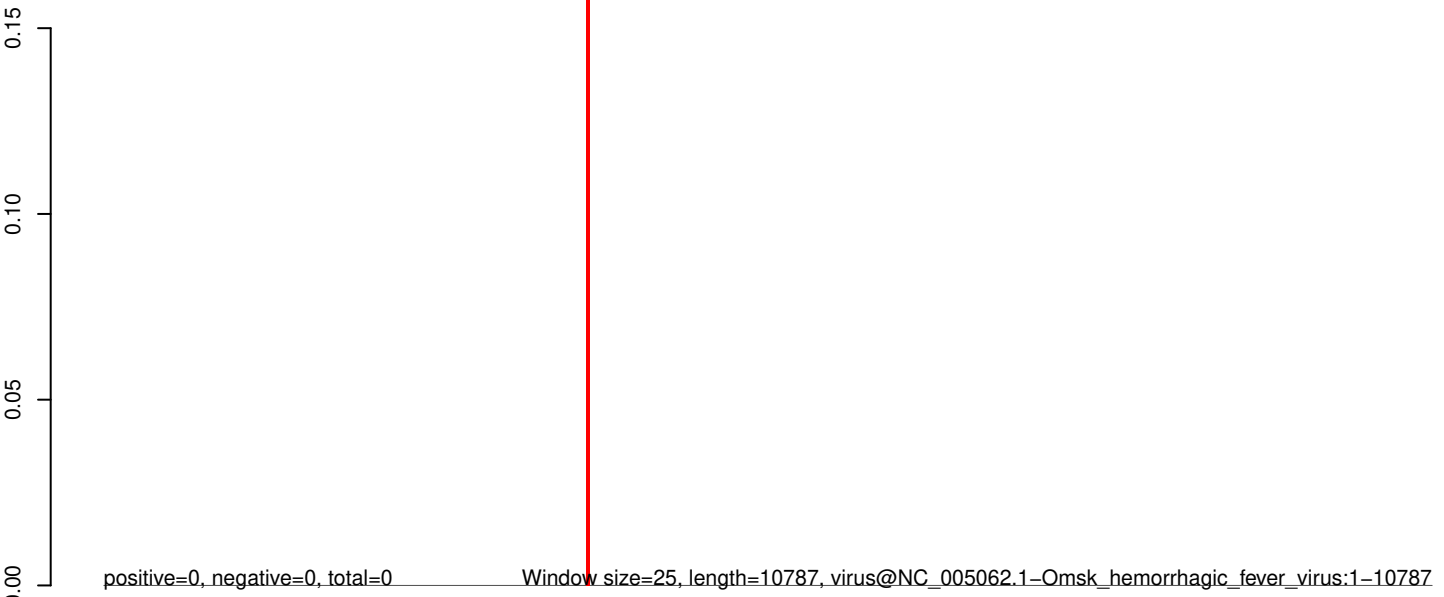
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



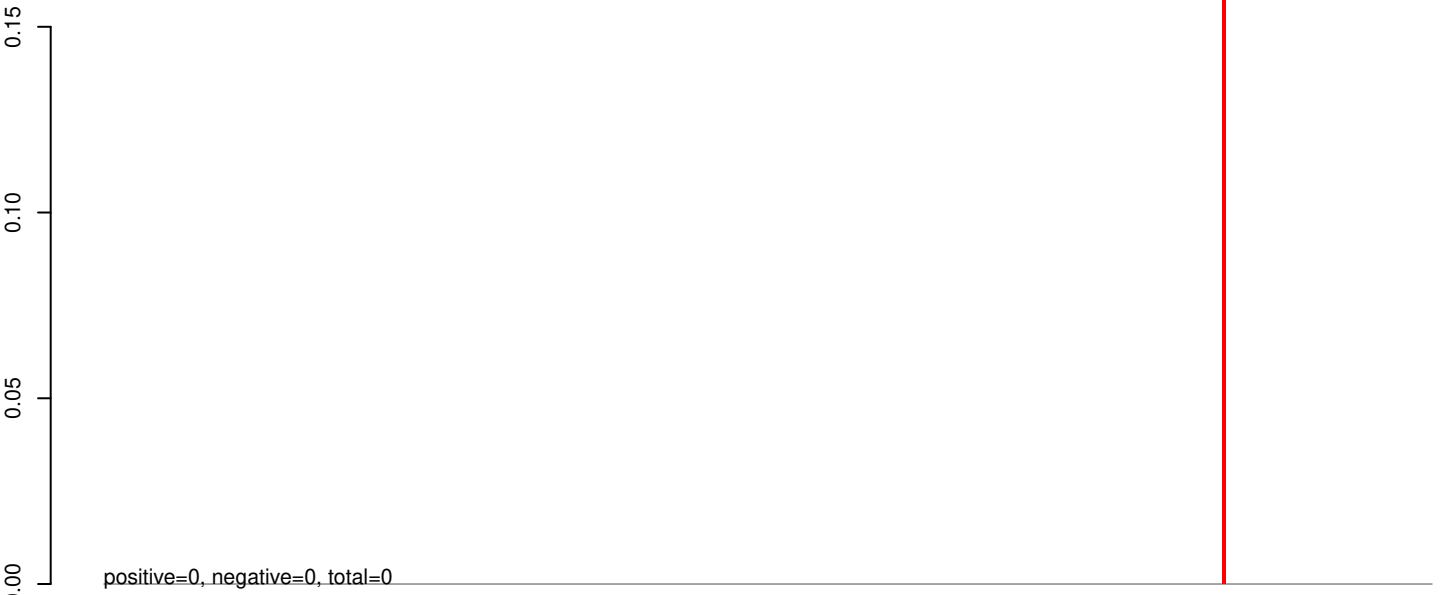
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



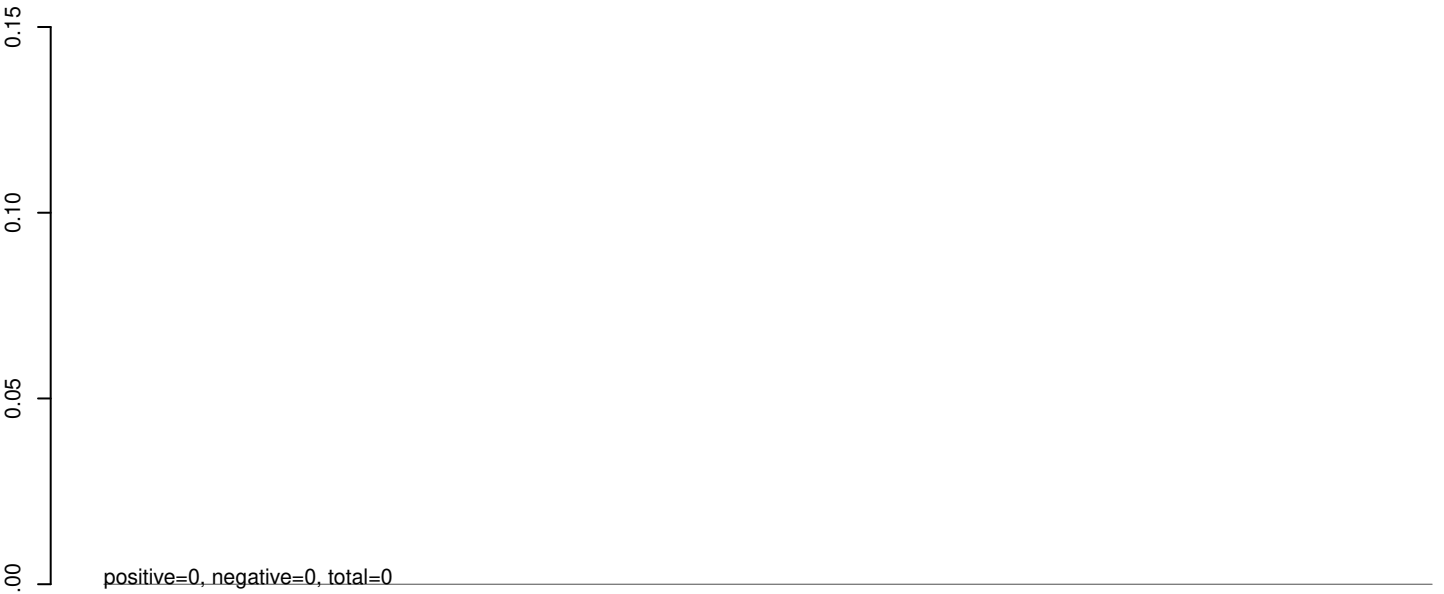
AeAeg_Aag2_mock_dsFluc_RA.rep



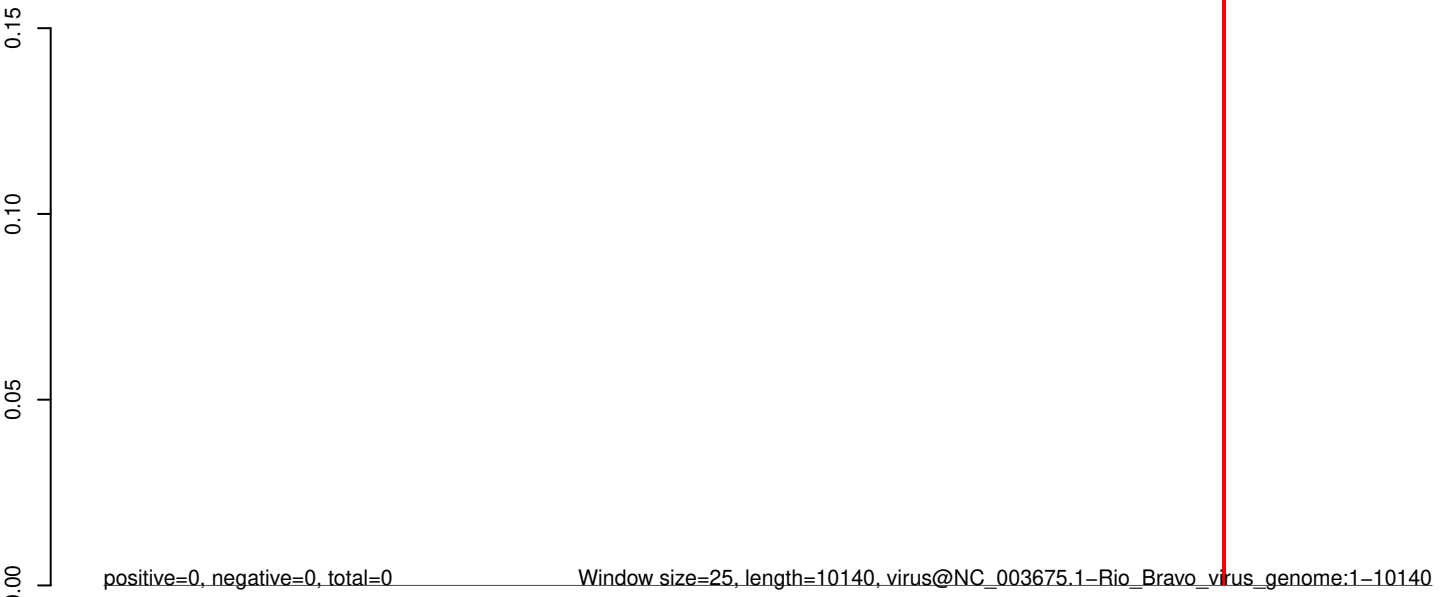
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

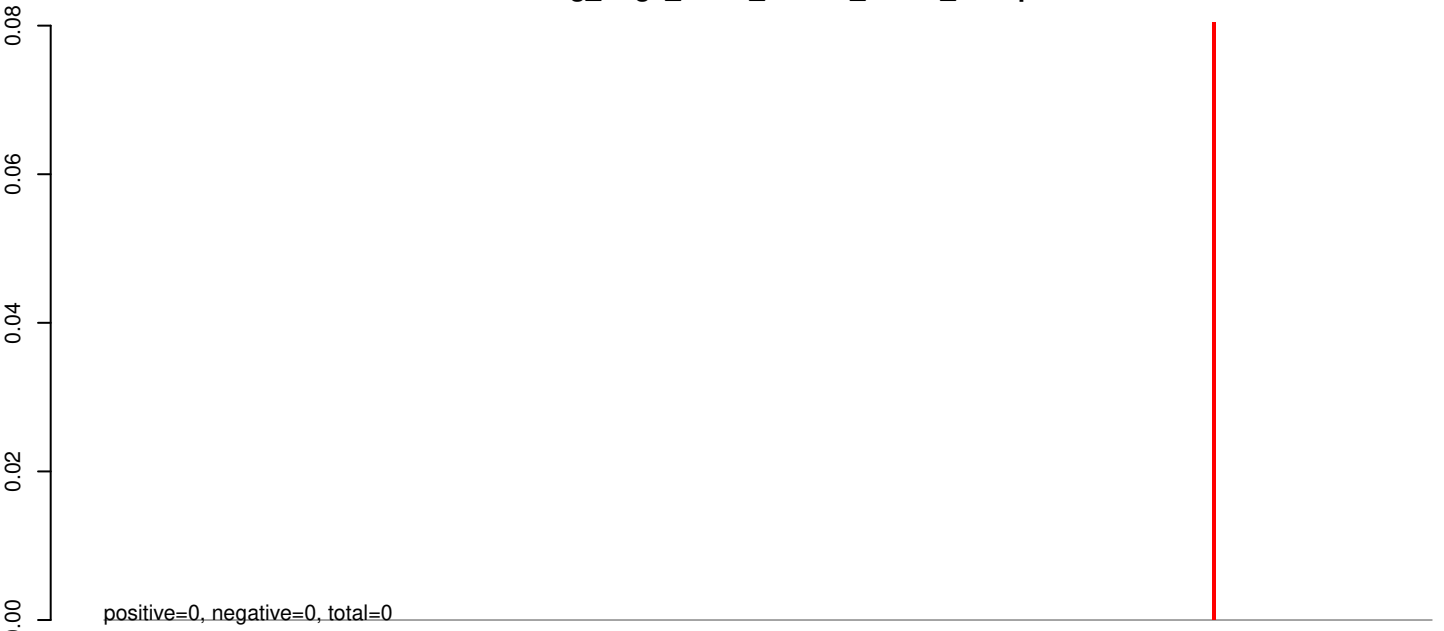


AeAeg_Aag2_mock_dsFluc_RA.rep

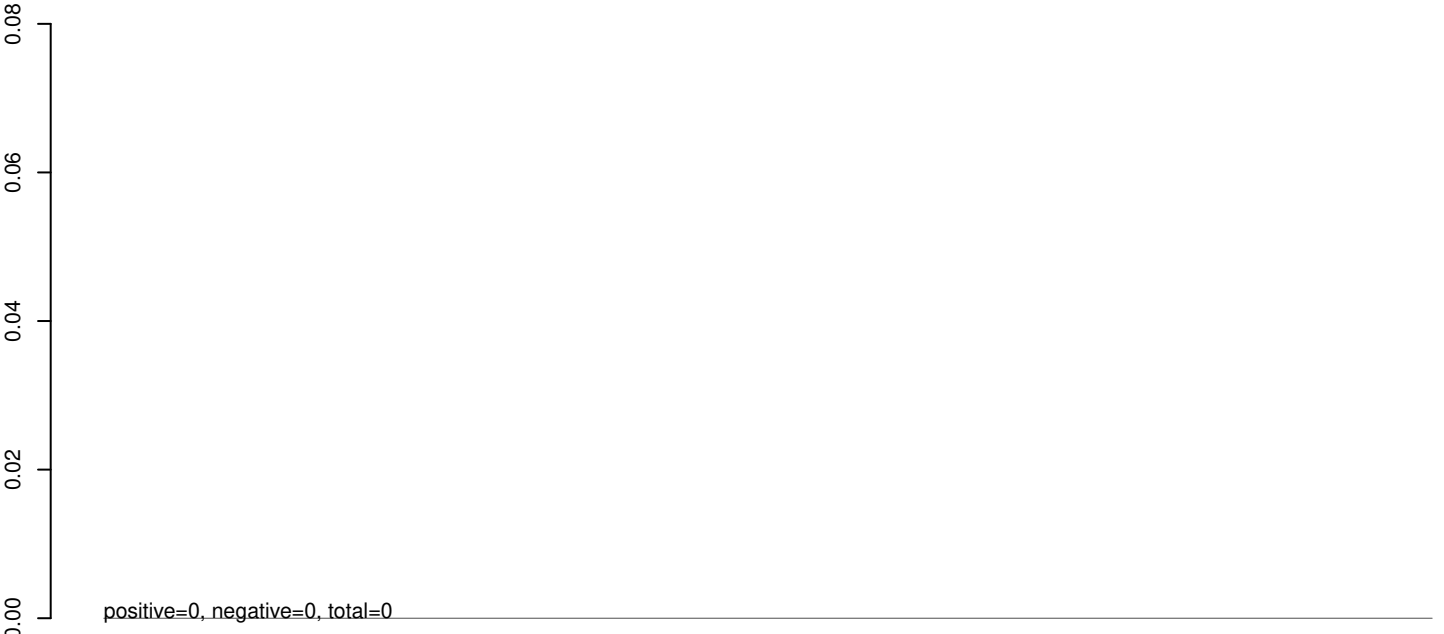


0 2000 4000 6000 8000 10000

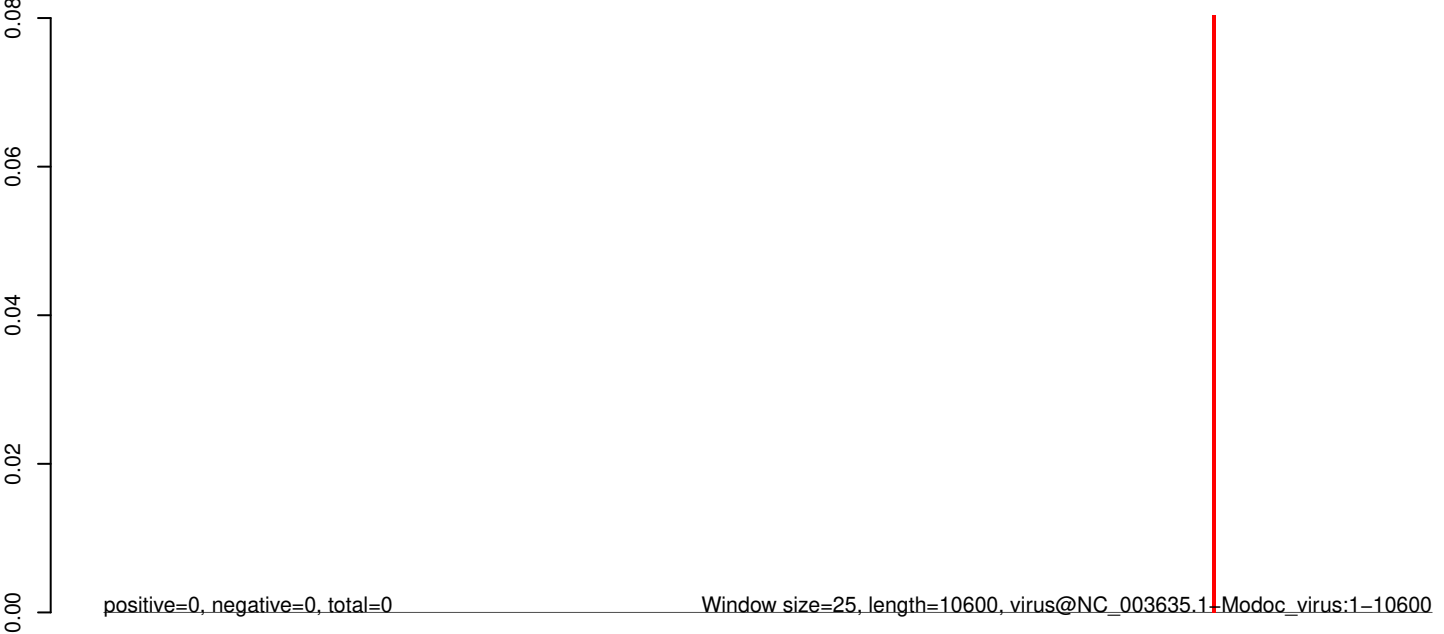
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

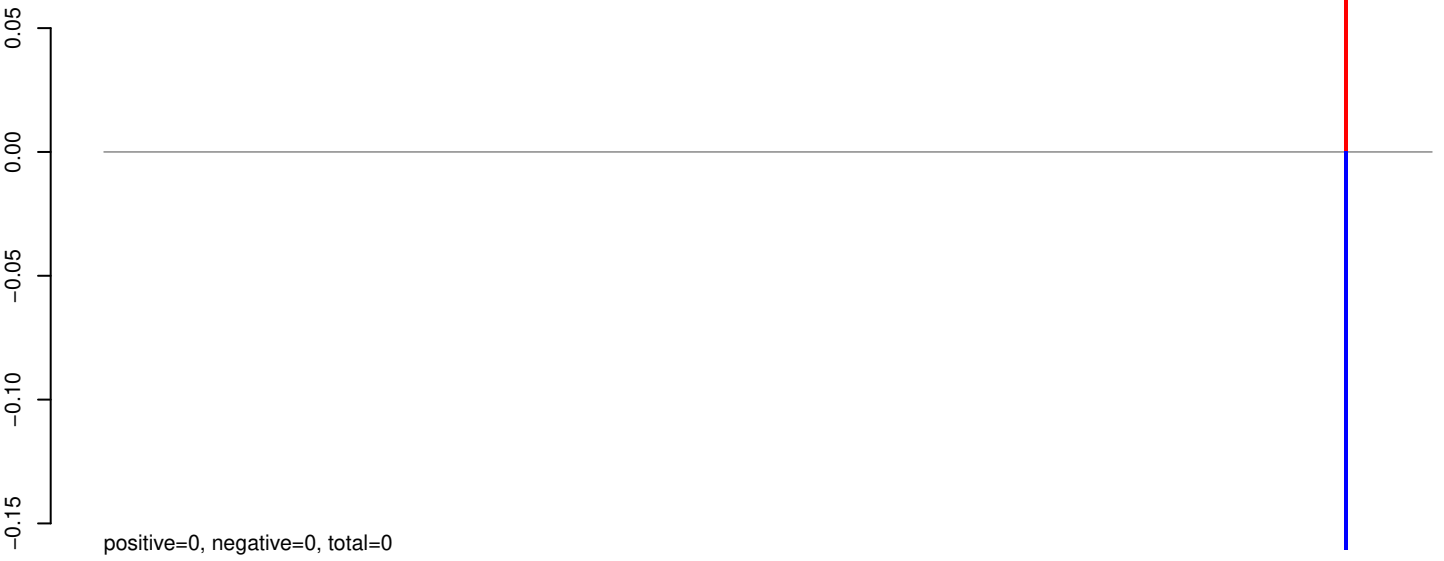


AeAeg_Aag2_mock_dsFluc_RA.rep

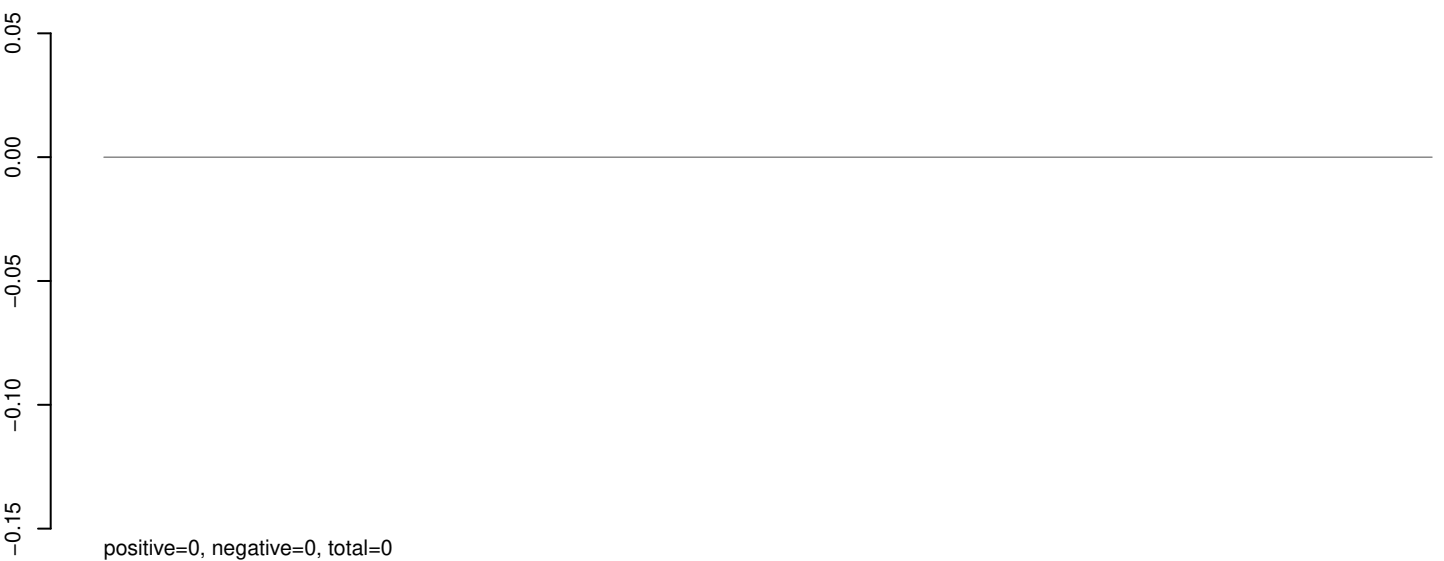


0 2000 4000 6000 8000 10000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



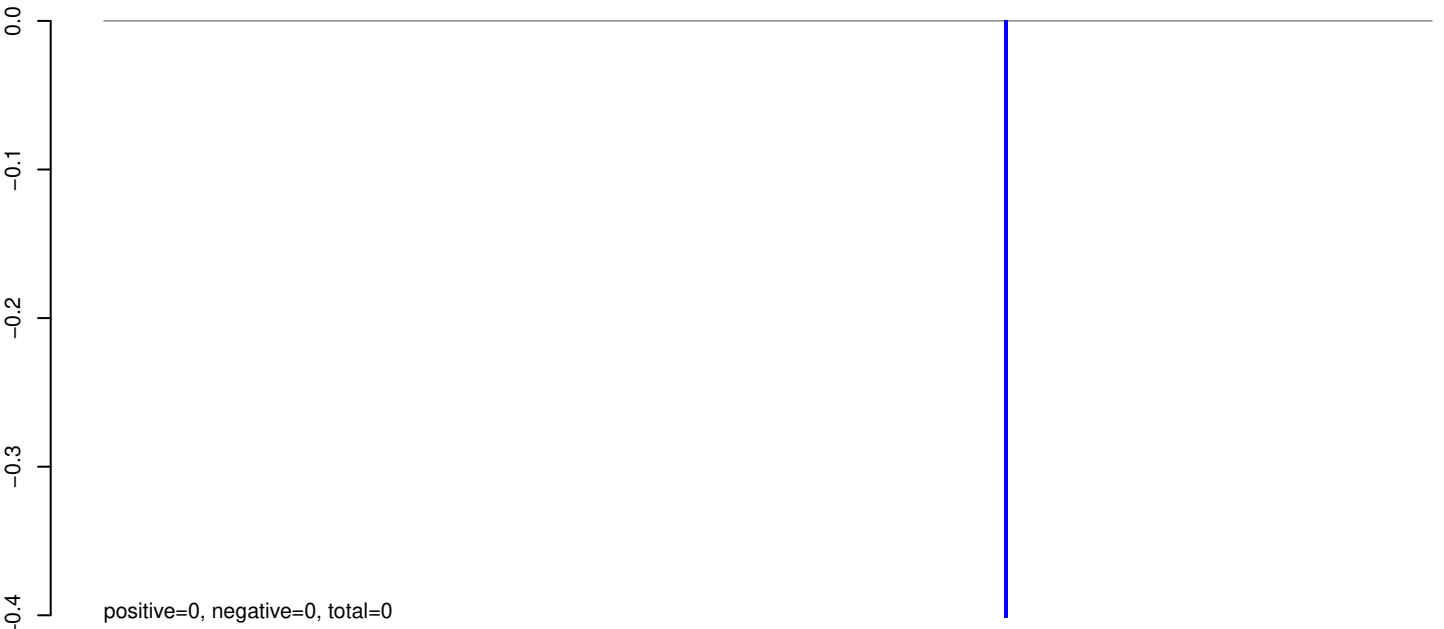
AeAeg_Aag2_mock_dsFluc_RA.rep



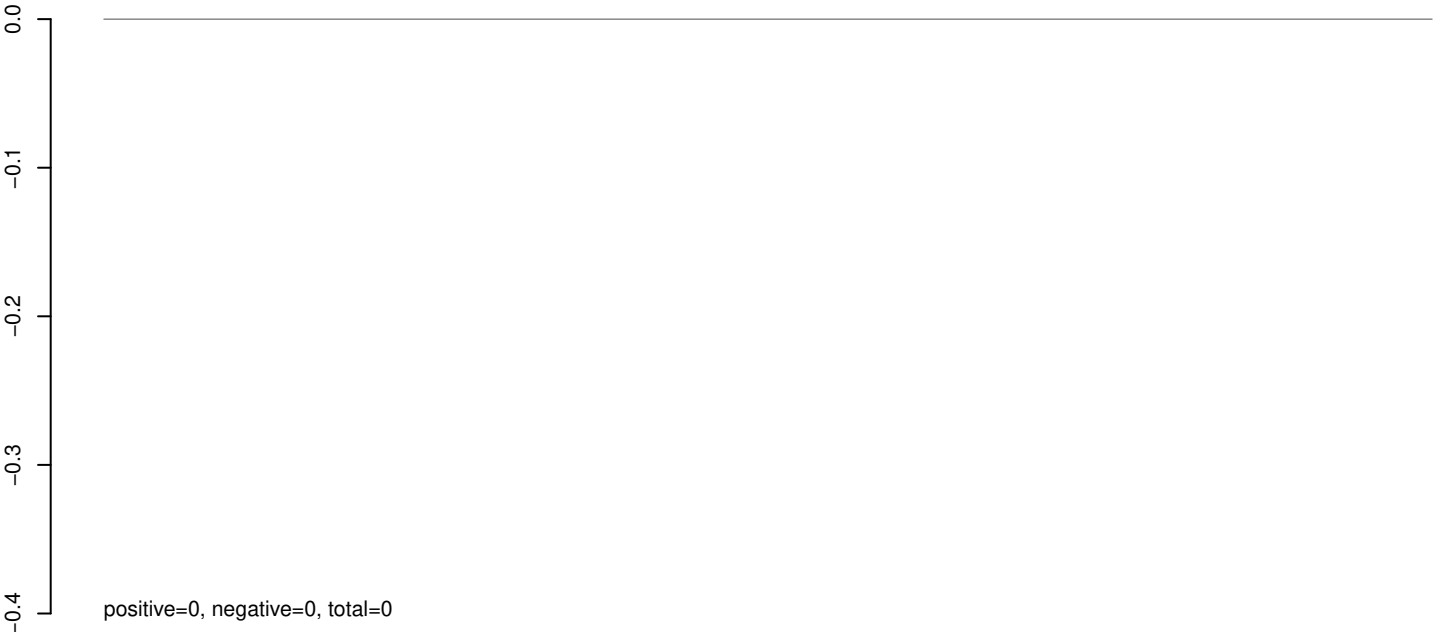
Window size=25, length=10649, virus@NC_002640.1-Dengue_virus_411-10649

0 2000 4000 6000 8000 10000

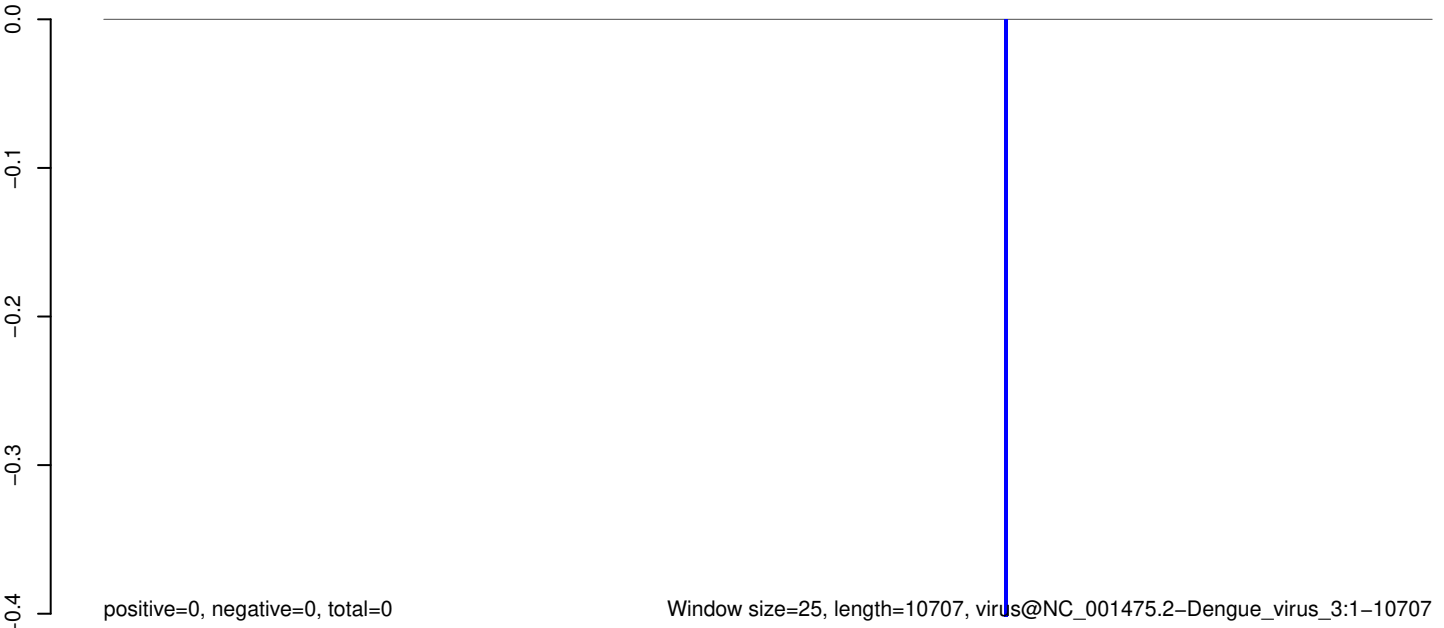
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



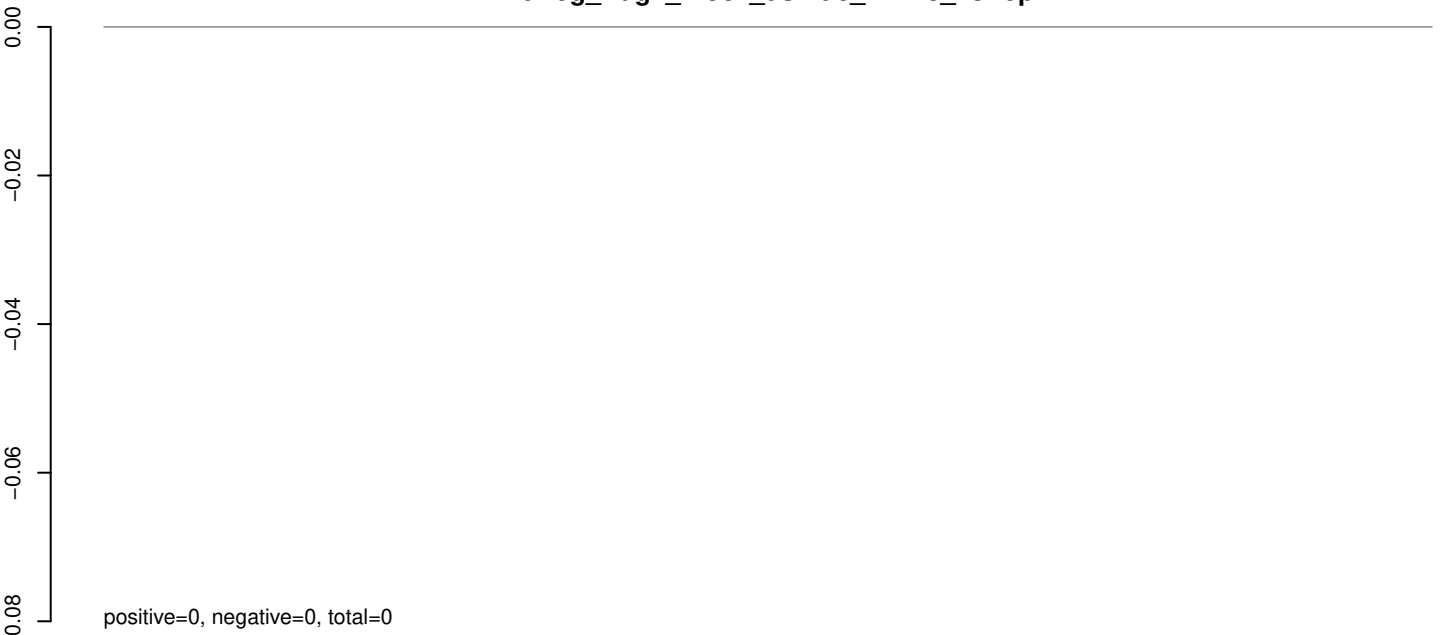
AeAeg_Aag2_mock_dsFluc_RA.rep



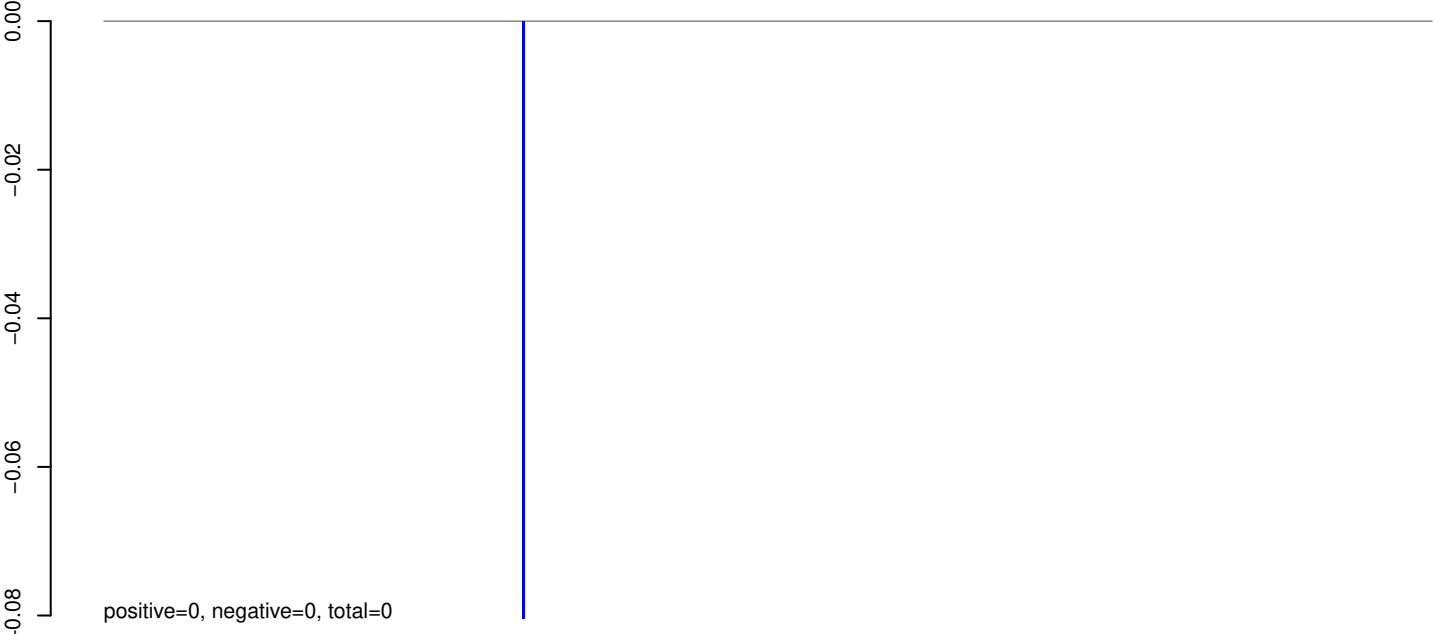
Window size=25, length=10707, virus@NC_001475.2-Dengue_virus_3:1-10707

0 2000 4000 6000 8000 10000

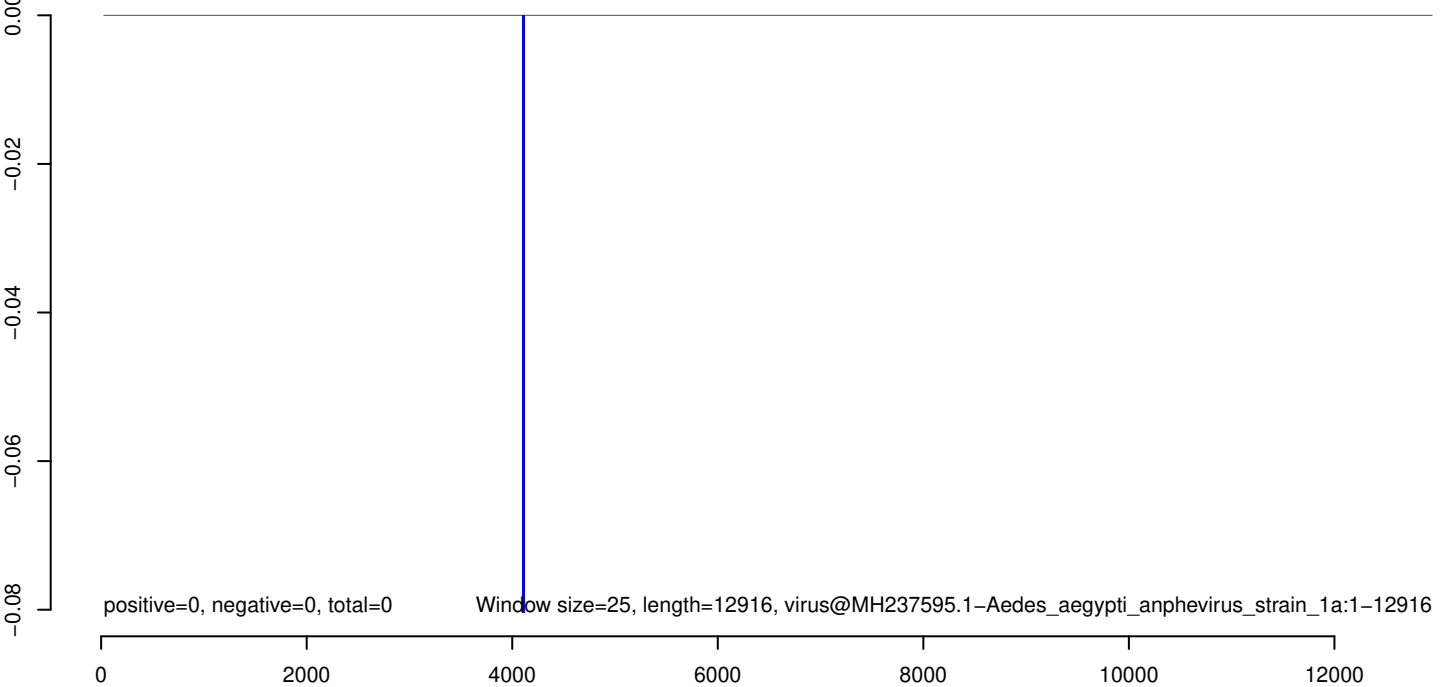
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



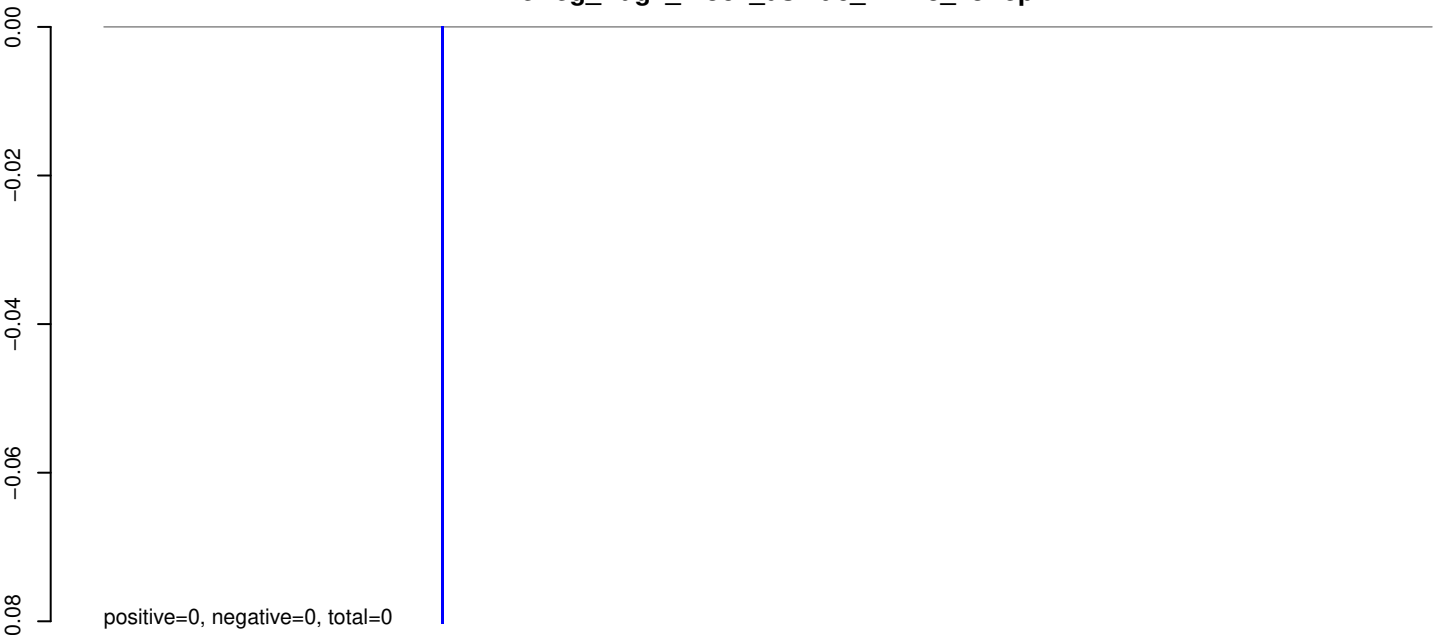
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



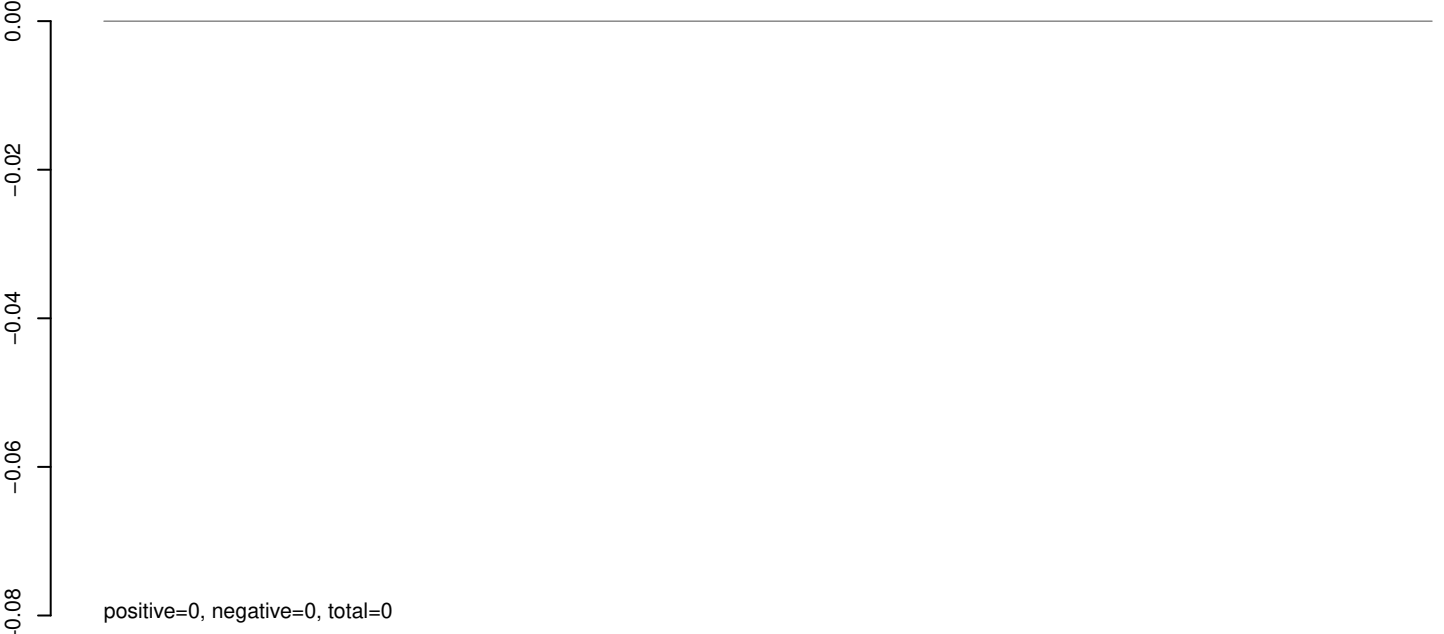
AeAeg_Aag2_mock_dsFluc_RA.rep



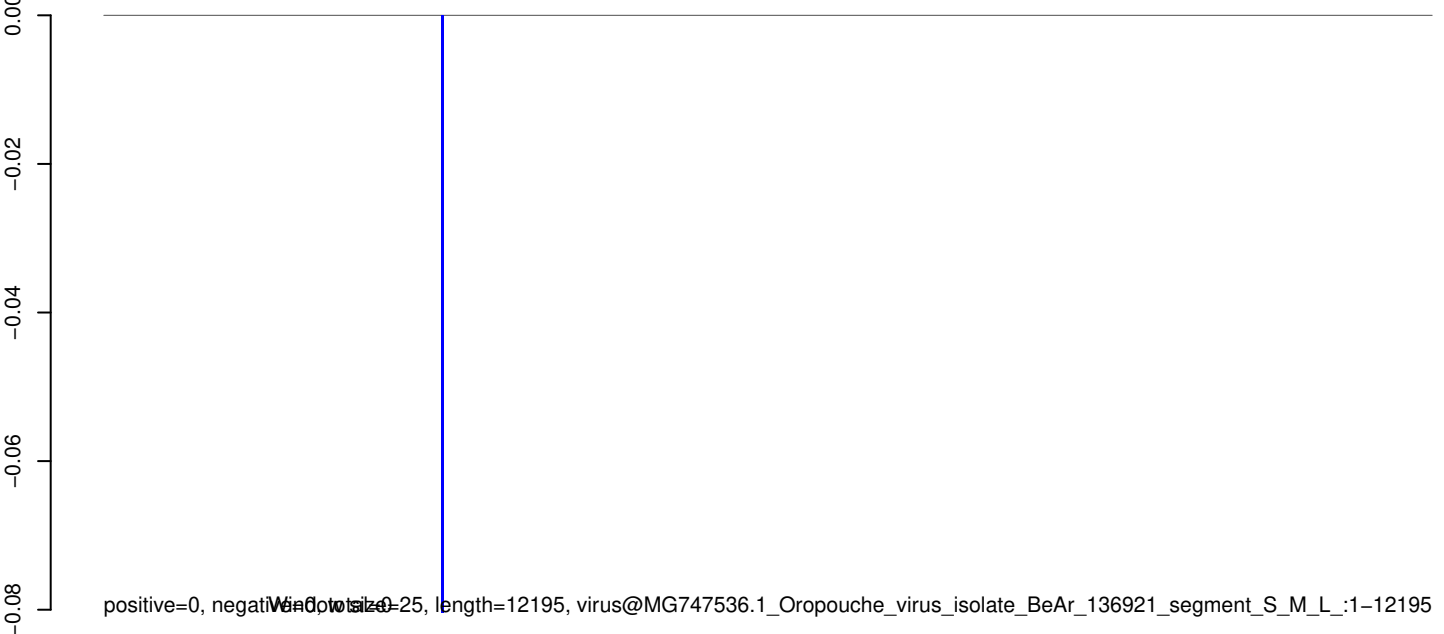
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep



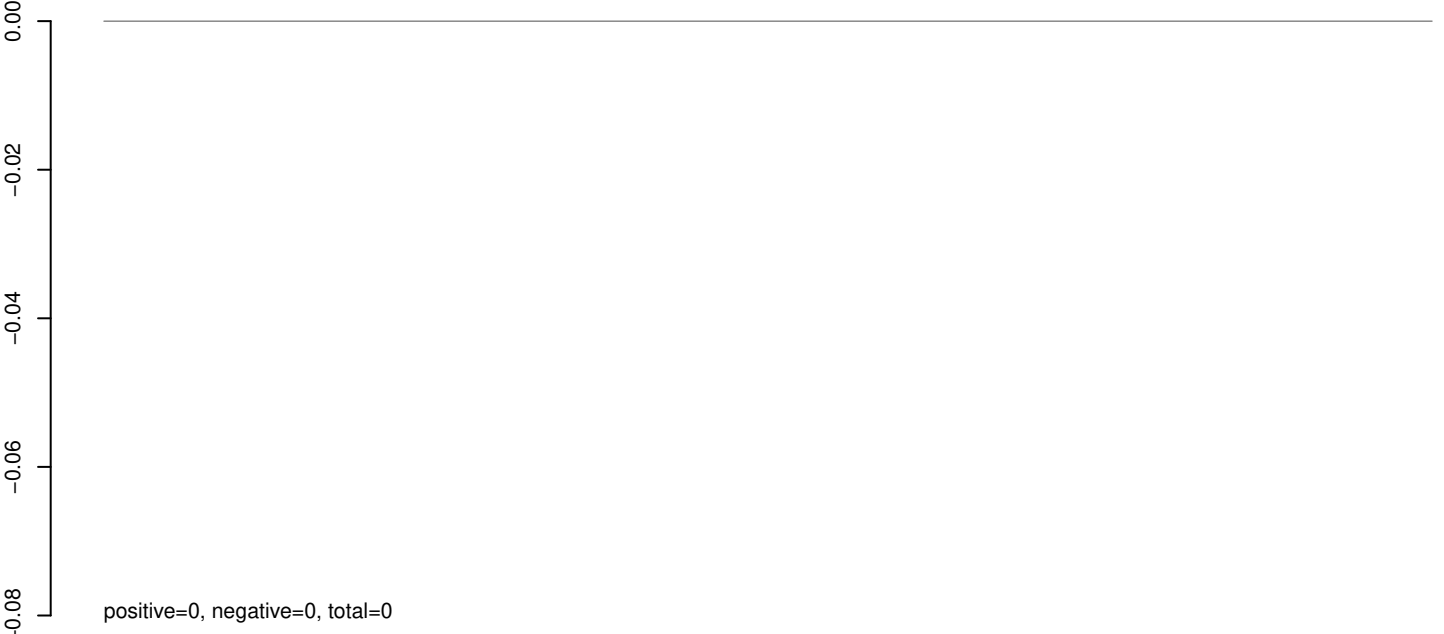
Window size=25, length=12195, virus@MG747536.1_Oropouche_virus_isolate_BeAr_136921_segment_S_M_L_:1-12195

0 2000 4000 6000 8000 10000 12000

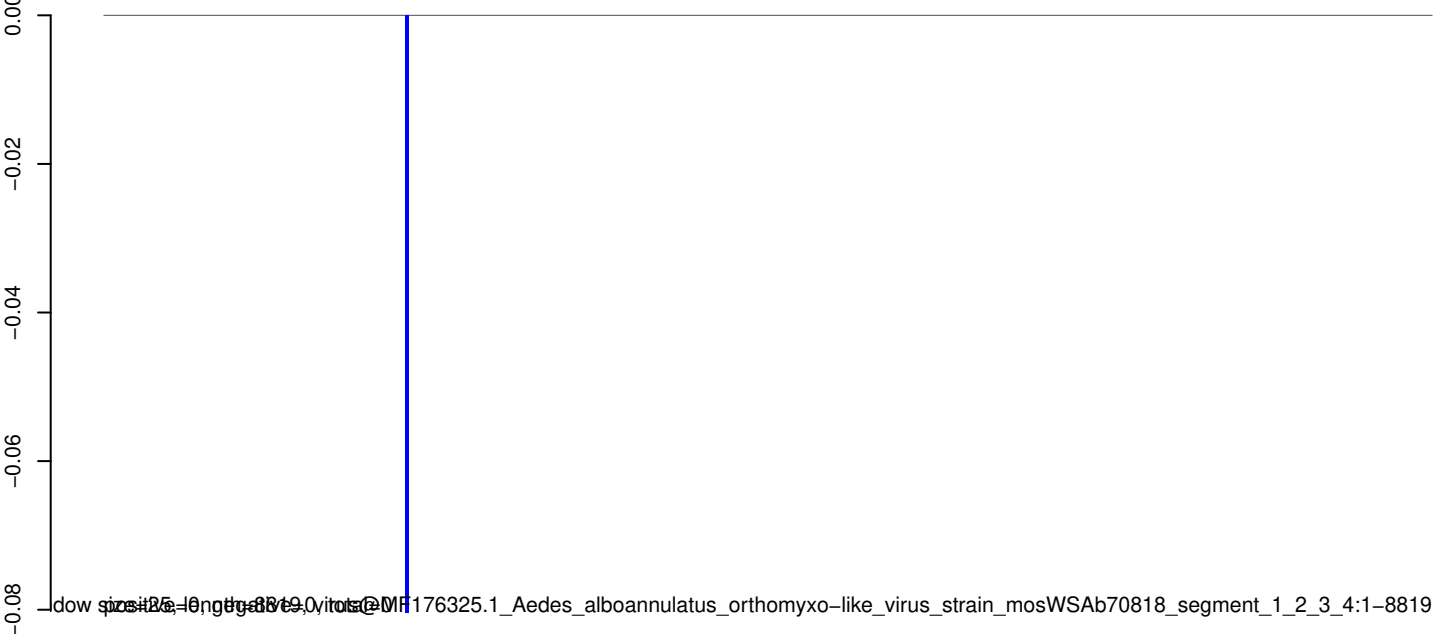
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



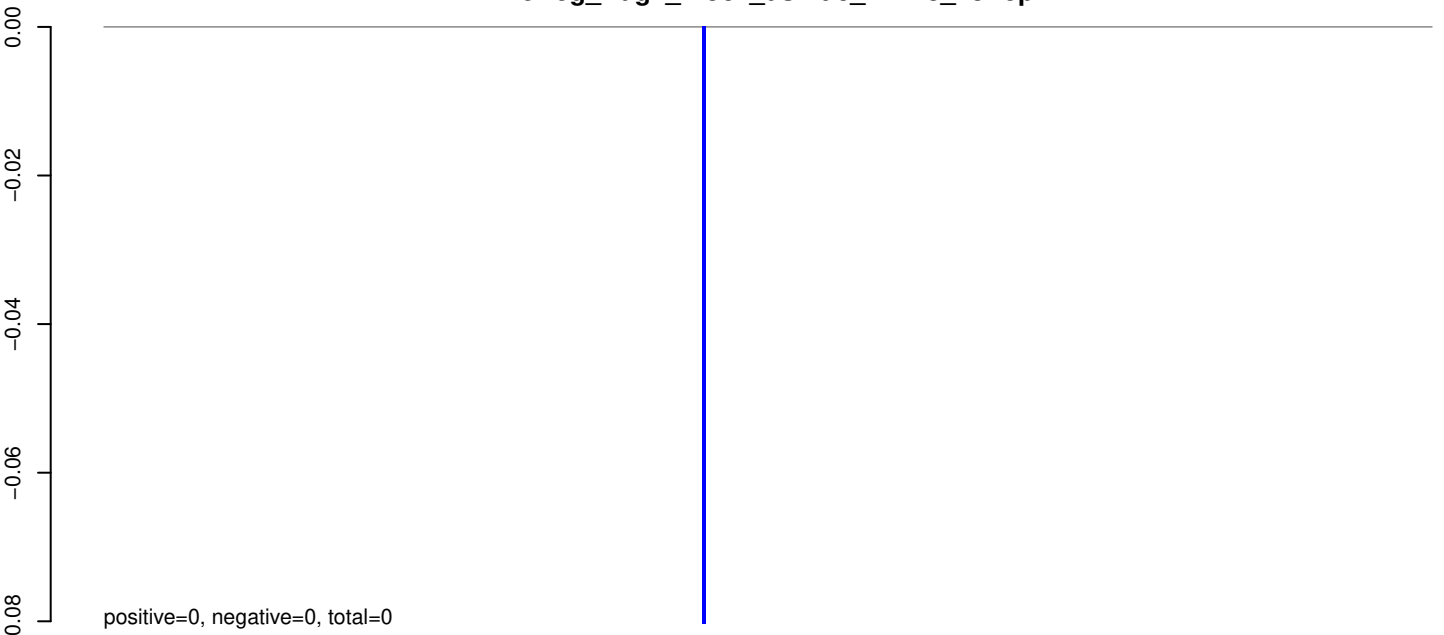
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



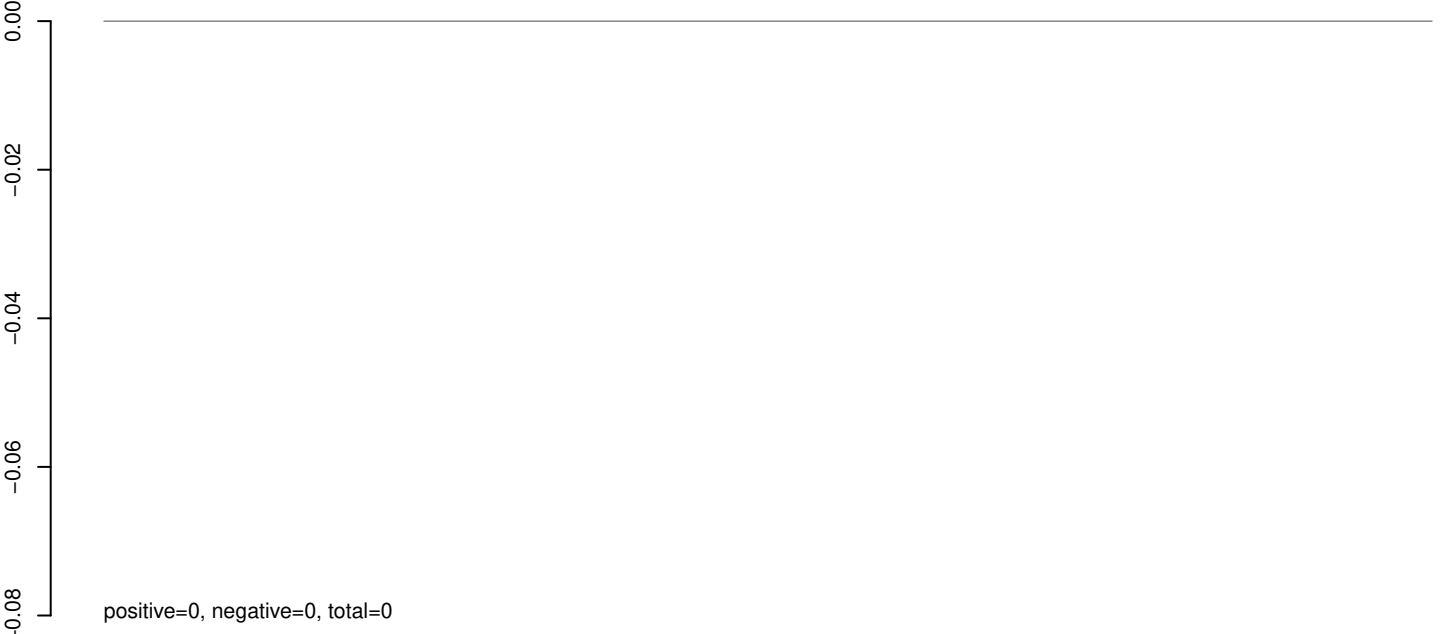
AeAeg_Aag2_mock_dsFluc_RA.rep



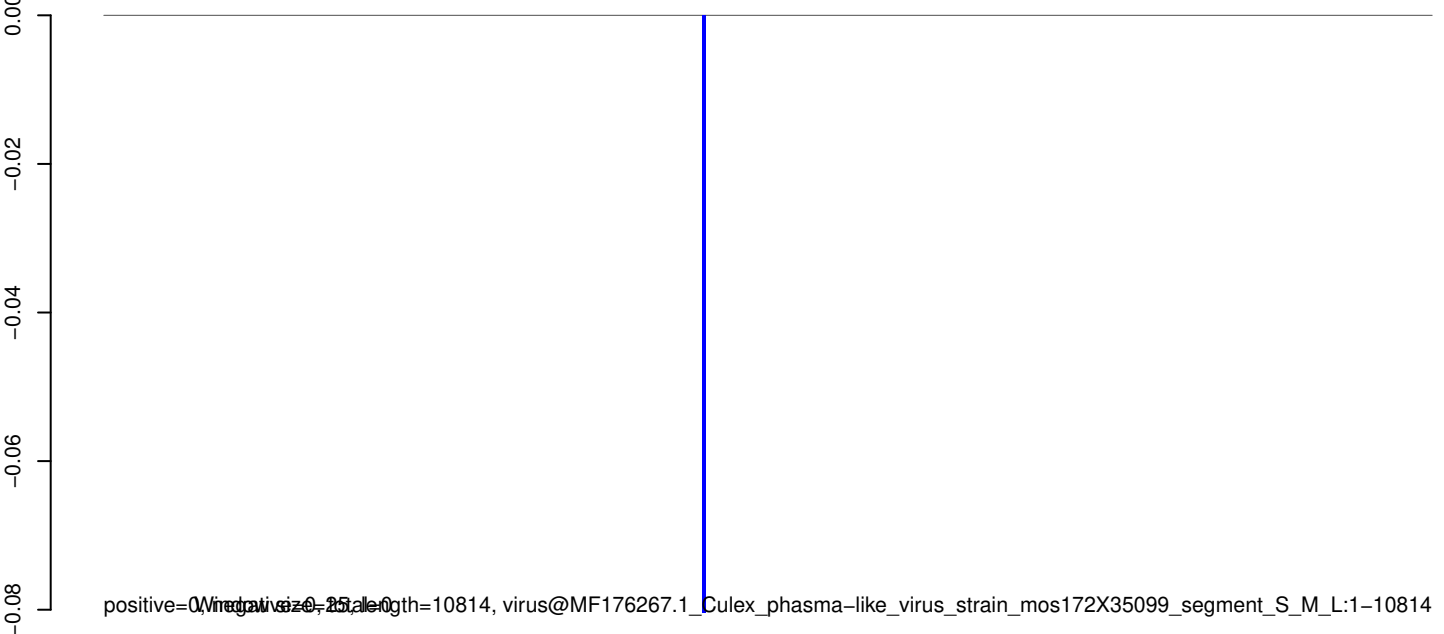
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep



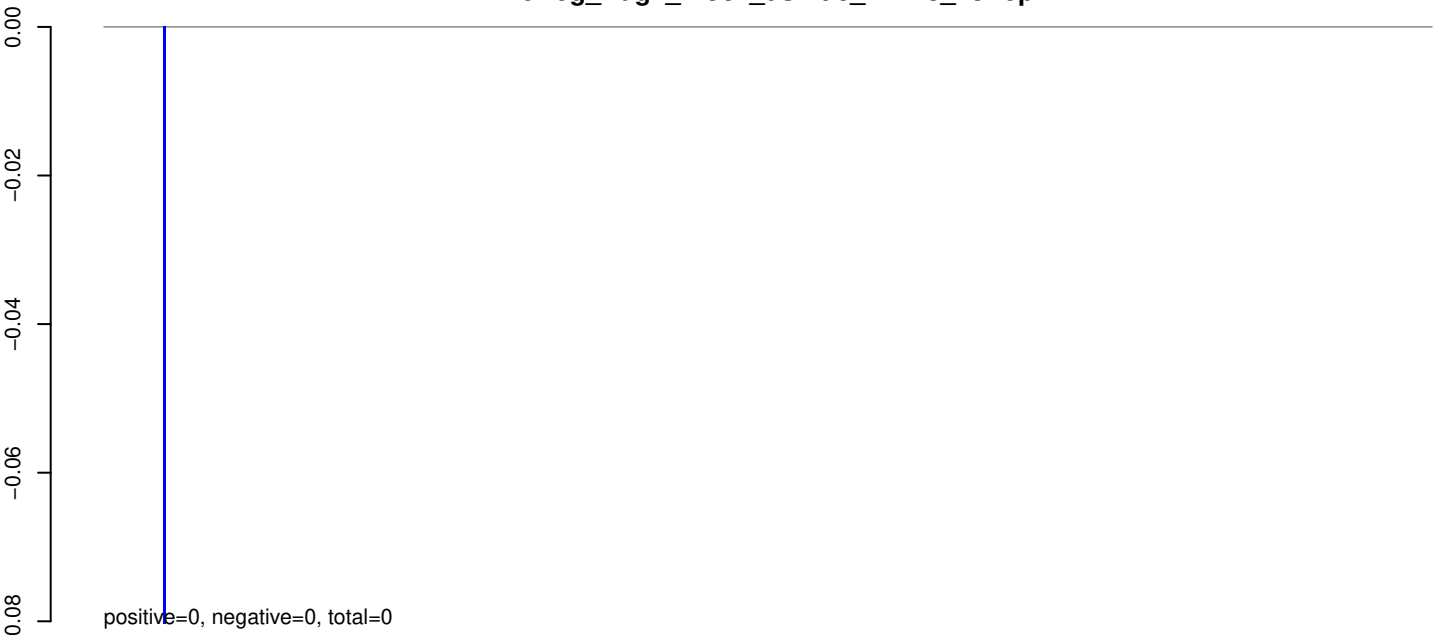
0 2000 4000 6000 8000 10000

0.00 -0.02 -0.04 -0.06 -0.08

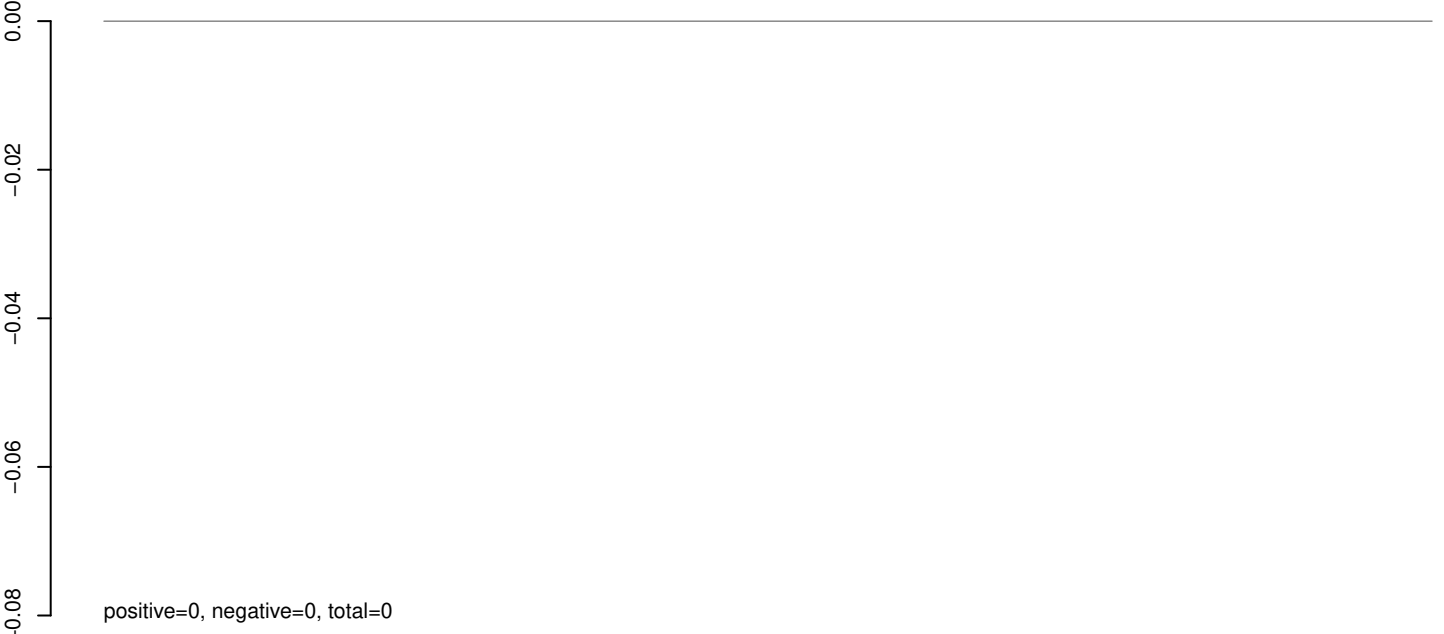
positive=0, negative=0, total=0

10814, virus@MF176267.1_Culex_phasma-like_virus_strain_mos172X35099_segment_S_M_L:1-10814

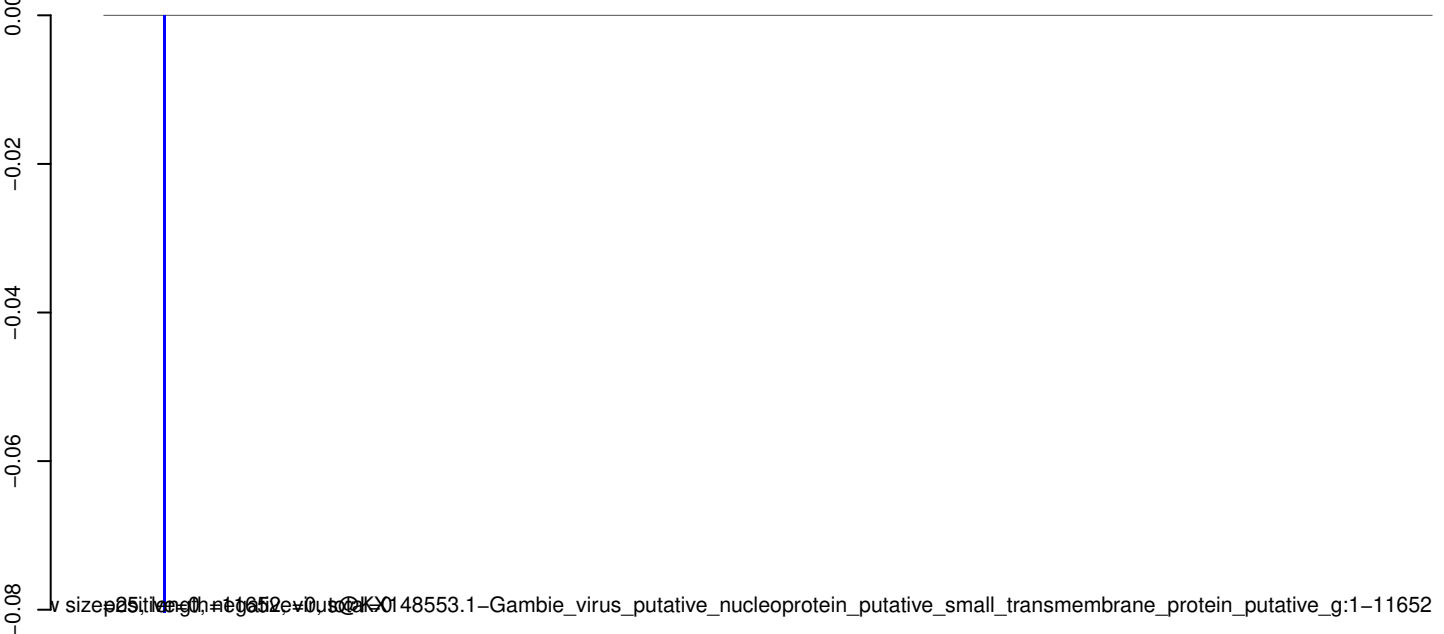
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



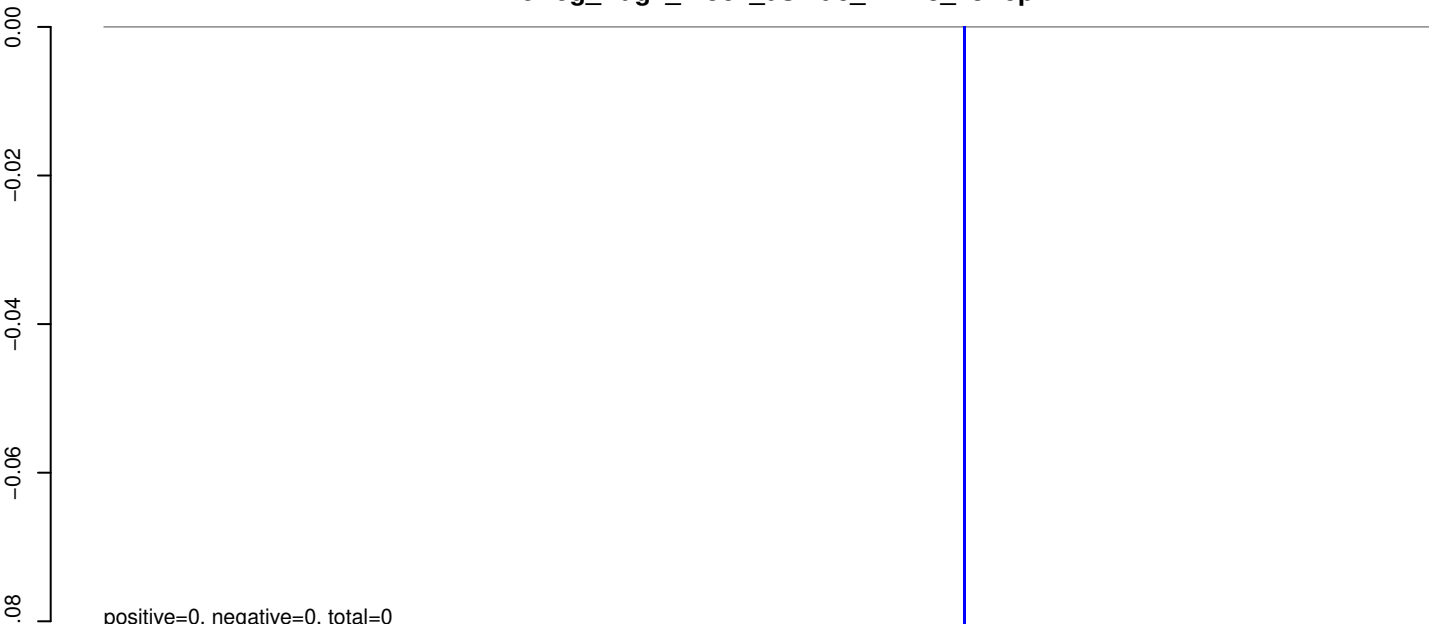
AeAeg_Aag2_mock_dsFluc_RA.rep



v size=25, time=0, n=1052, n_pos=0, n_neg=0, n_tot=1052, K=0.148553.1-Gambie_virus_putative_nucleoprotein_putative_small_transmembrane_protein_putative_g:1-11652

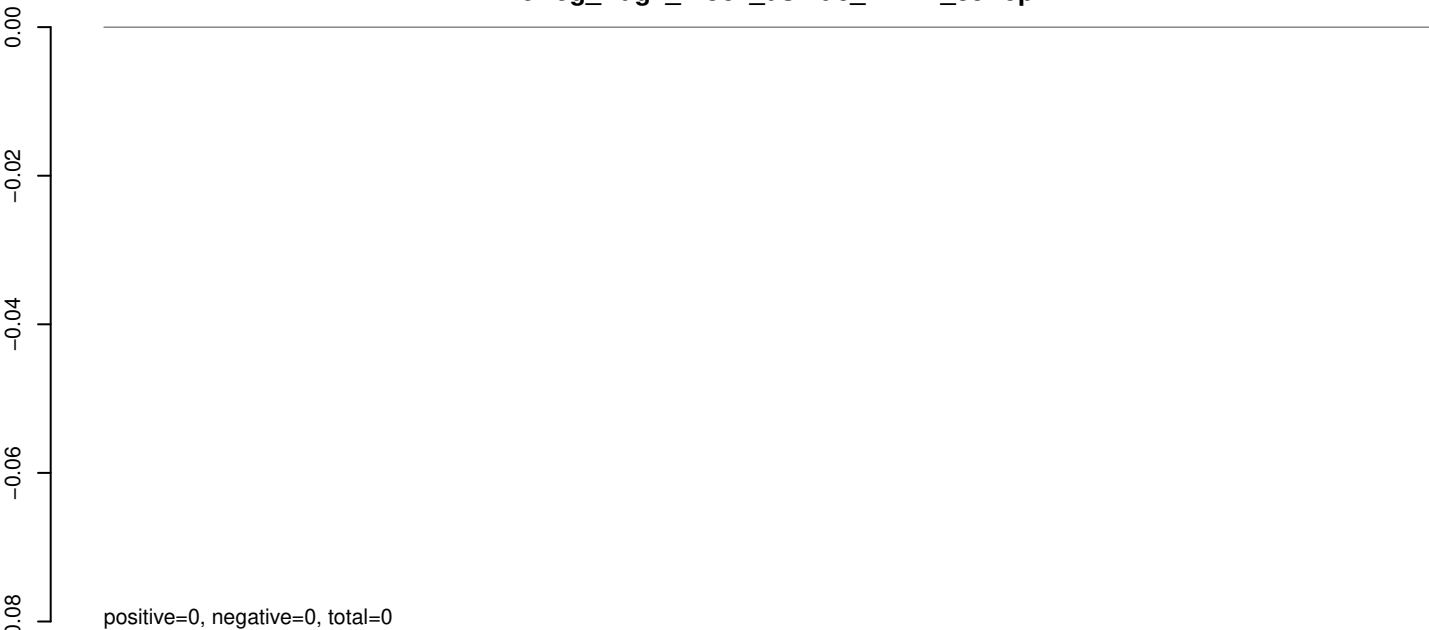
0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



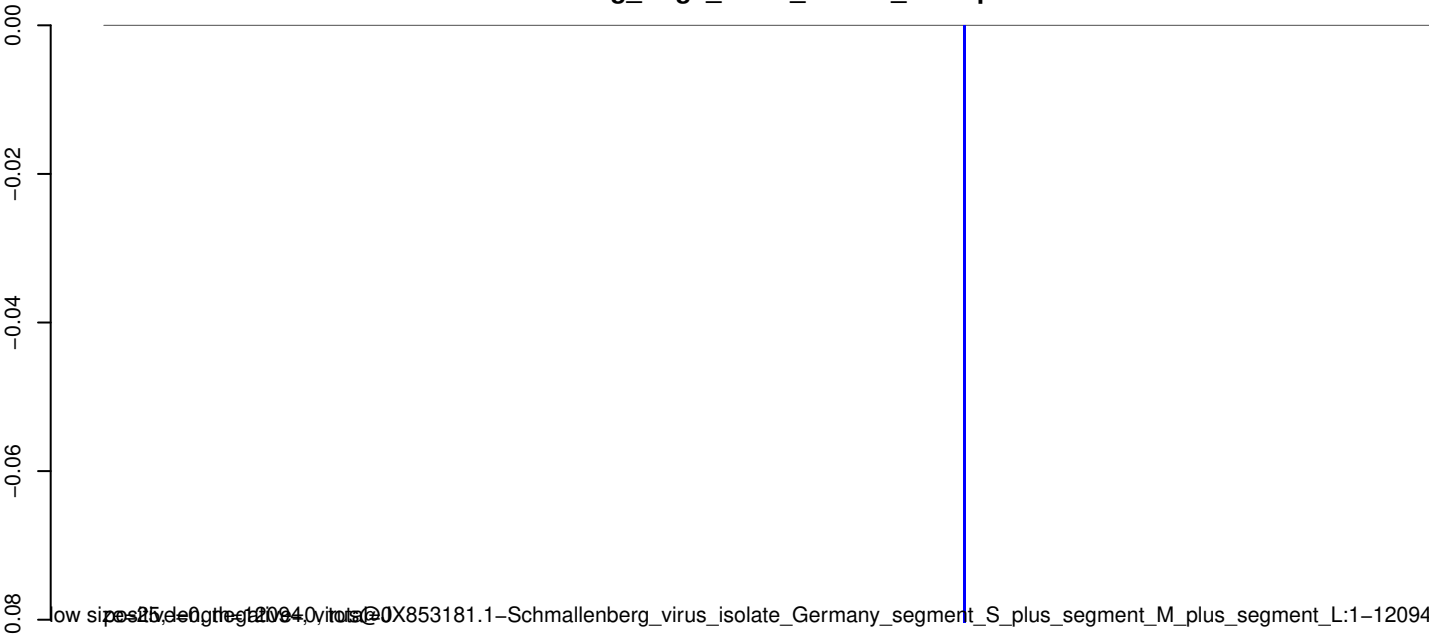
positive=0, negative=0, total=0

AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



positive=0, negative=0, total=0

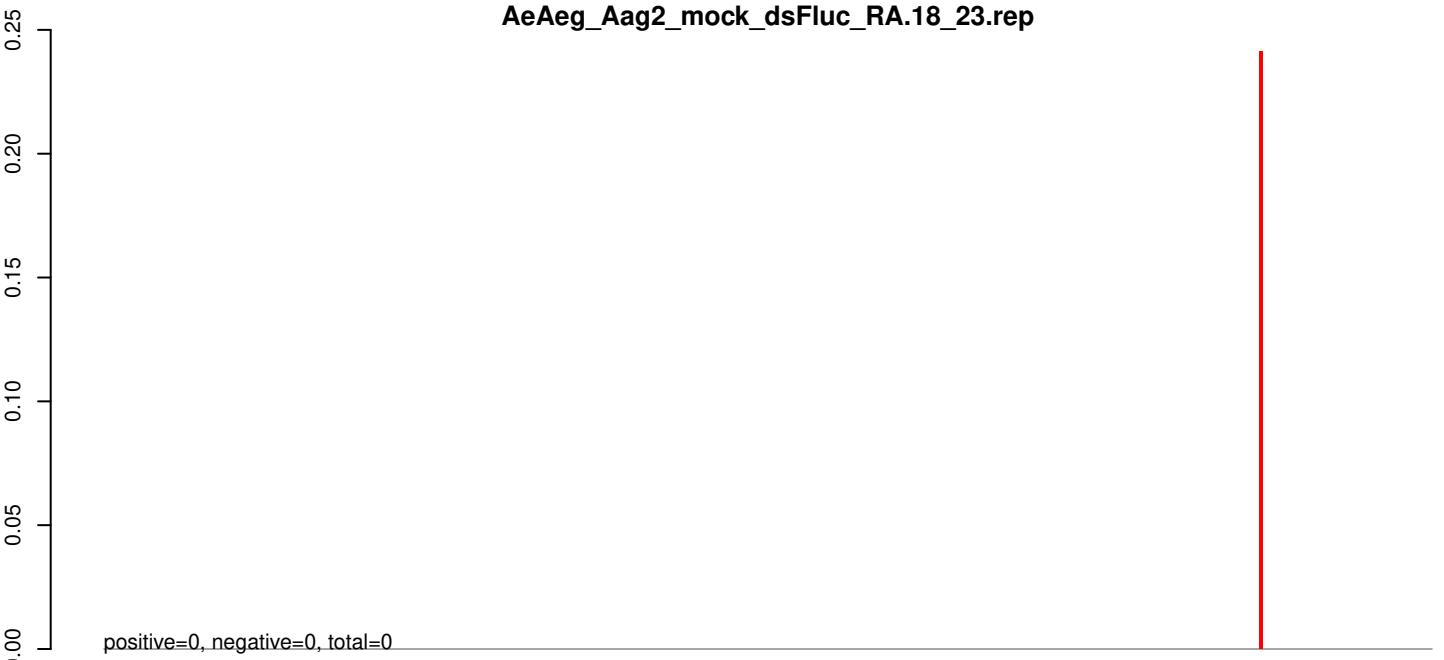
AeAeg_Aag2_mock_dsFluc_RA.rep



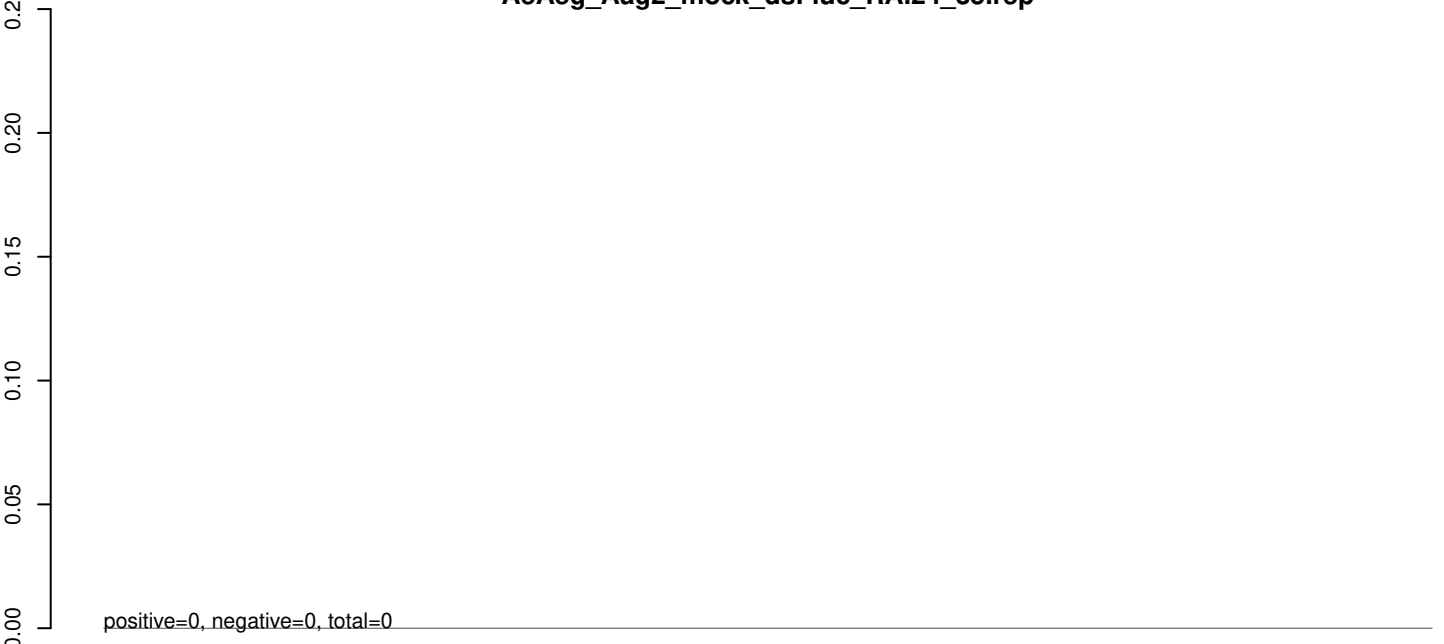
low size def of the 2004 virus @ X853181.1-Schmallenberg_virus_isolate_Germany_segment_S_plus_segment_M_plus_segment_L:1-12094

0 2000 4000 6000 8000 10000 12000

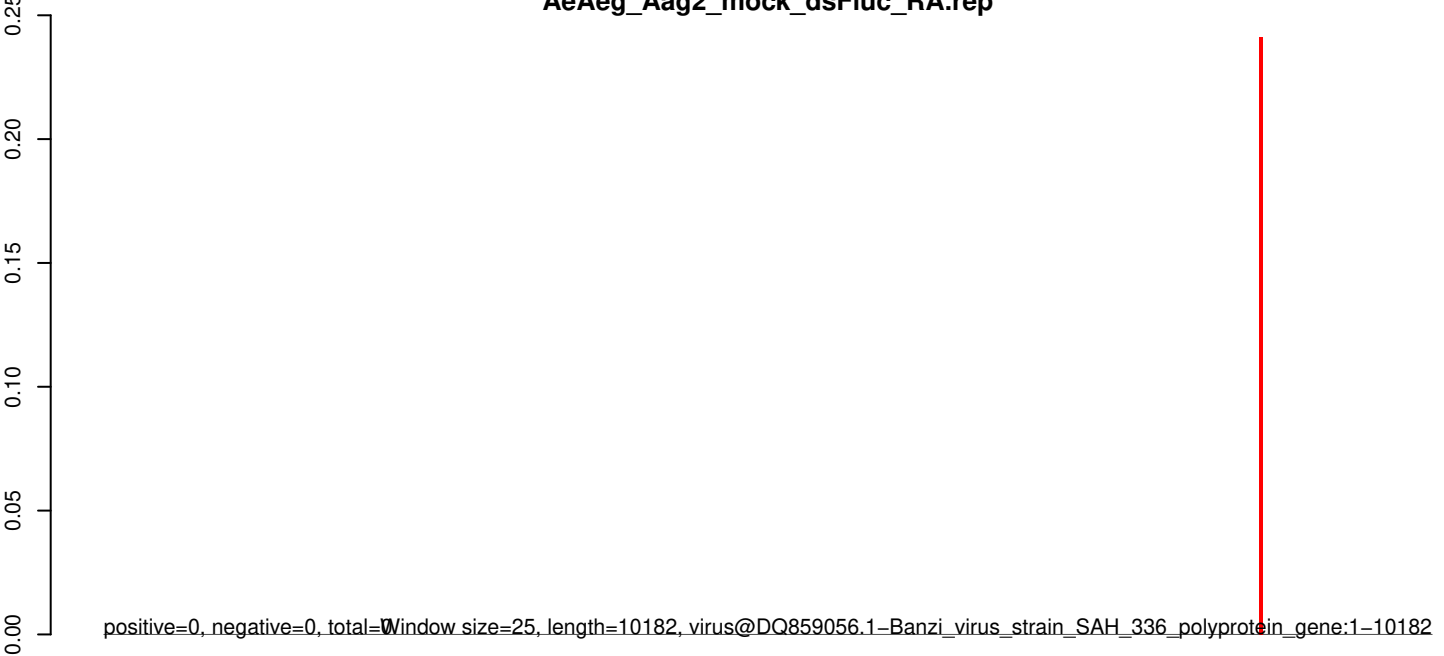
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

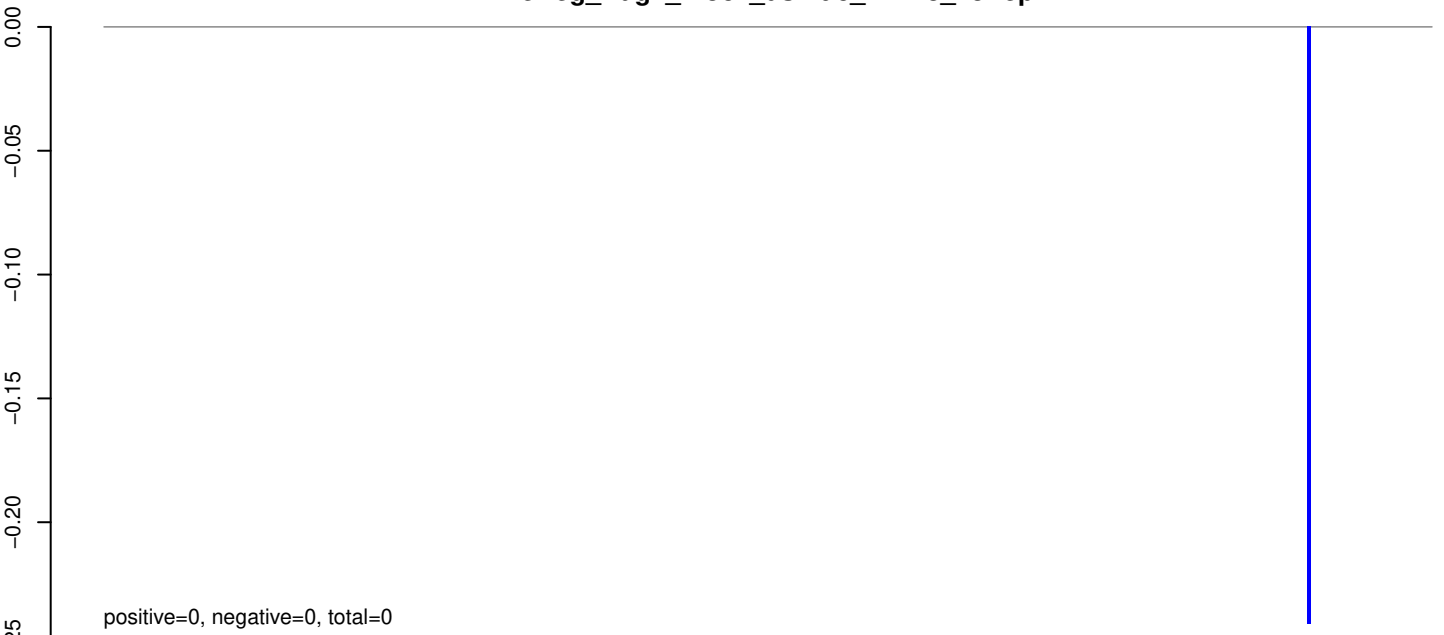


AeAeg_Aag2_mock_dsFluc_RA.rep

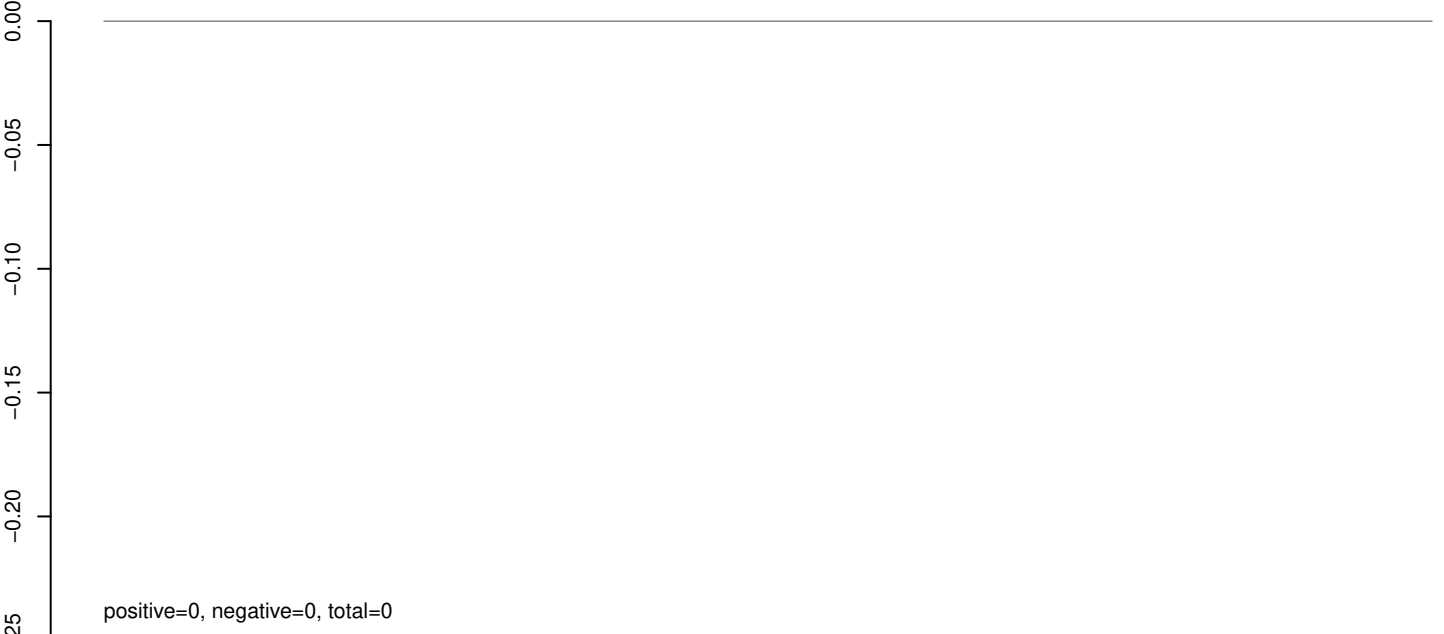


0 2000 4000 6000 8000 10000

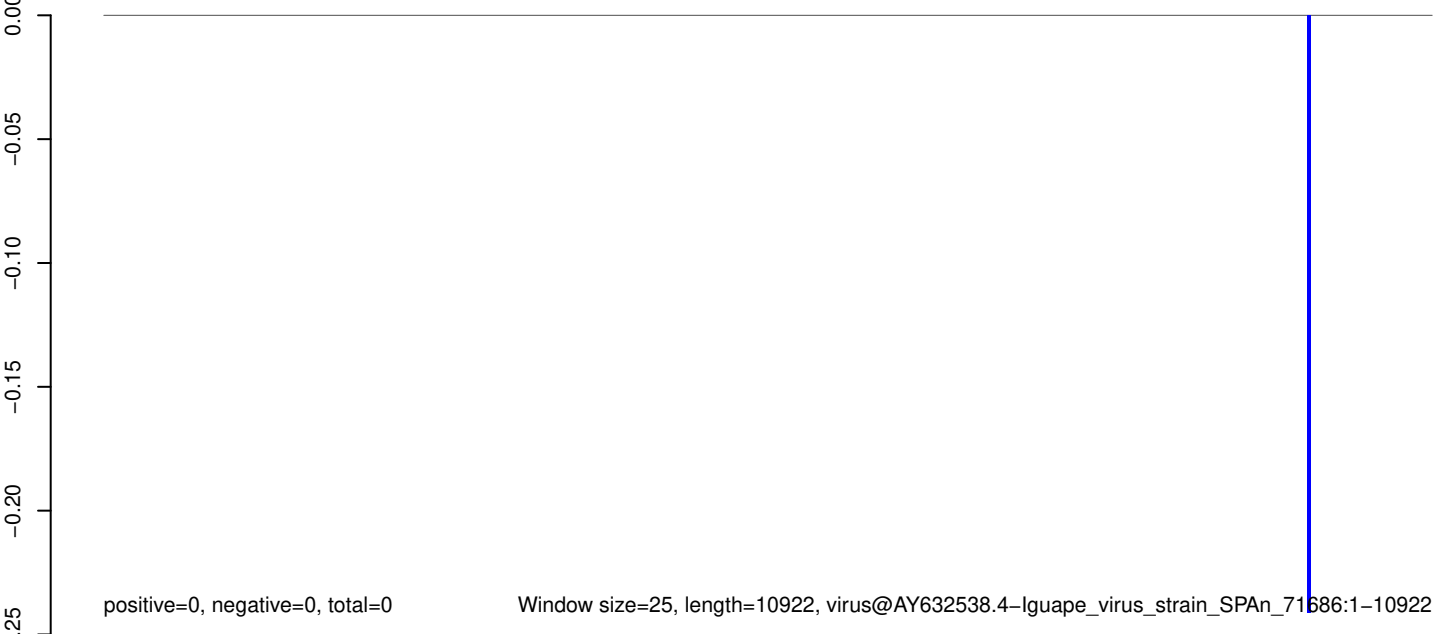
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

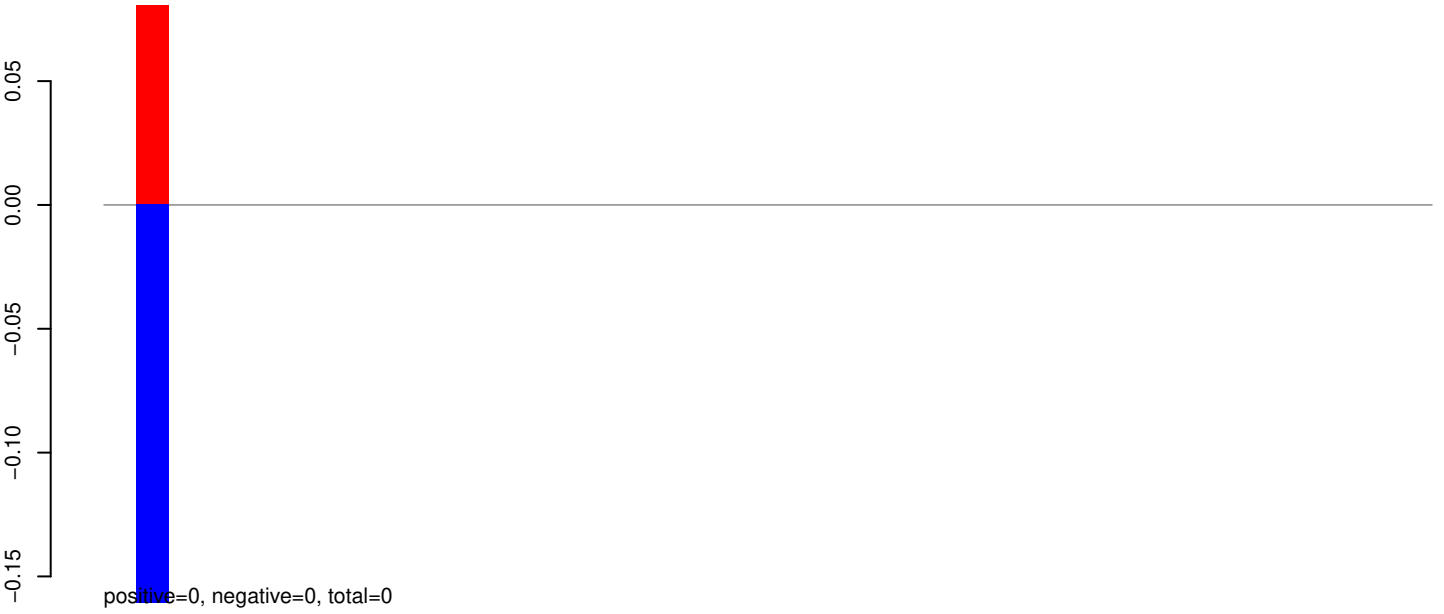


AeAeg_Aag2_mock_dsFluc_RA.rep

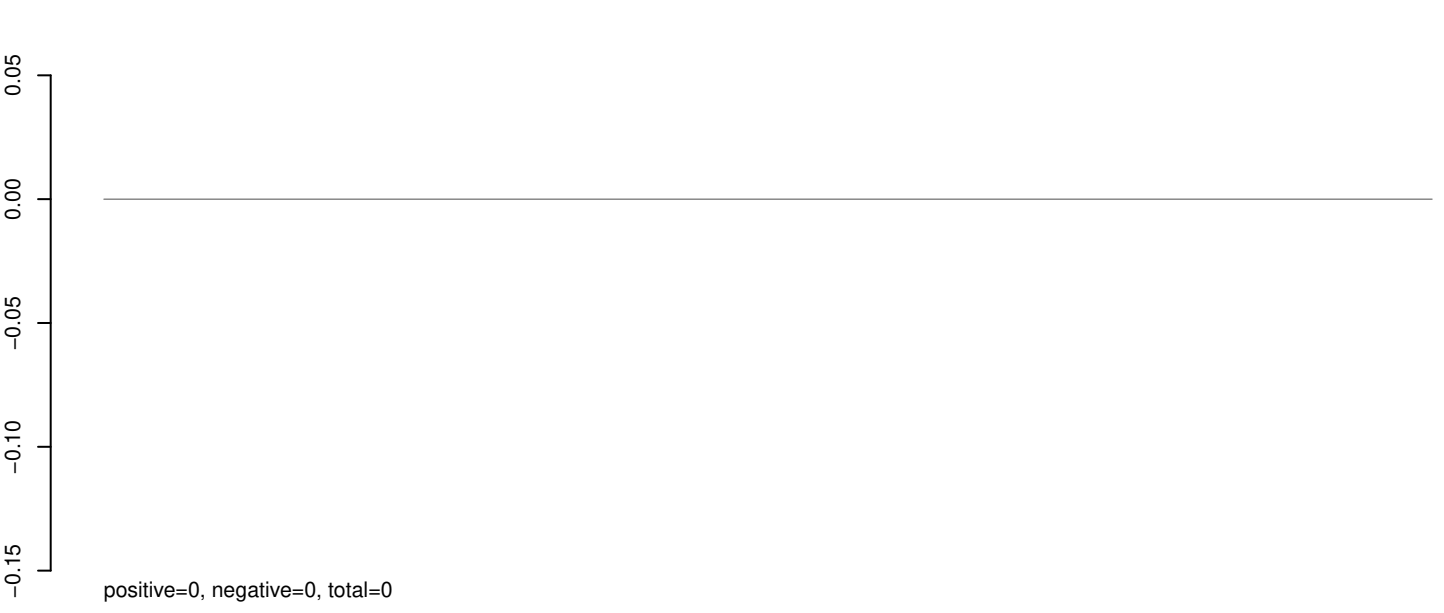


Window size=25, length=10922, virus@AY632538.4-Iguape_virus_strain_SPAn_71686:1-10922

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

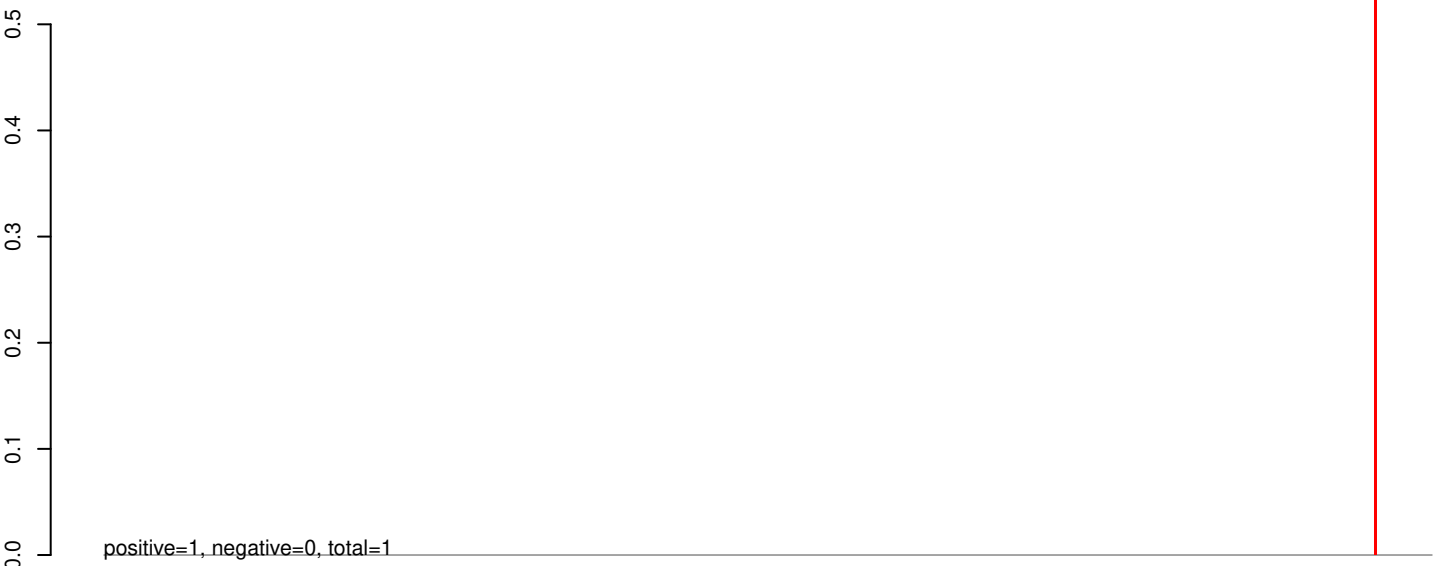


AeAeg_Aag2_mock_dsFluc_RA.rep

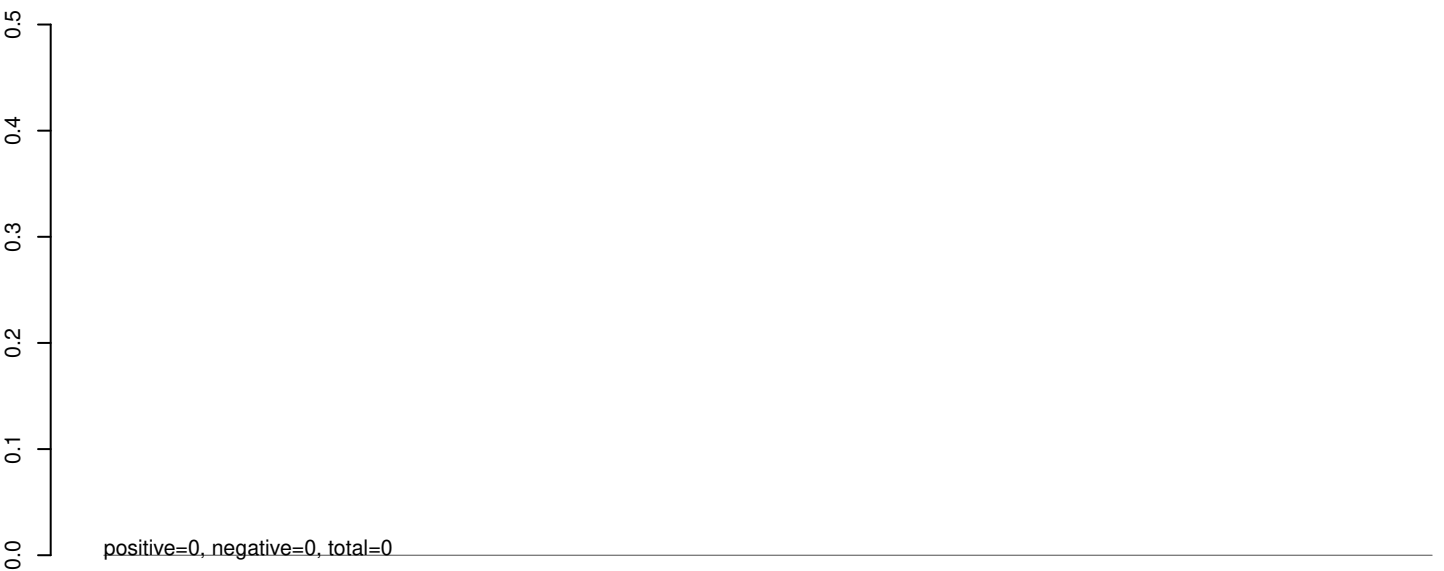


0 200 400 600 800 1000

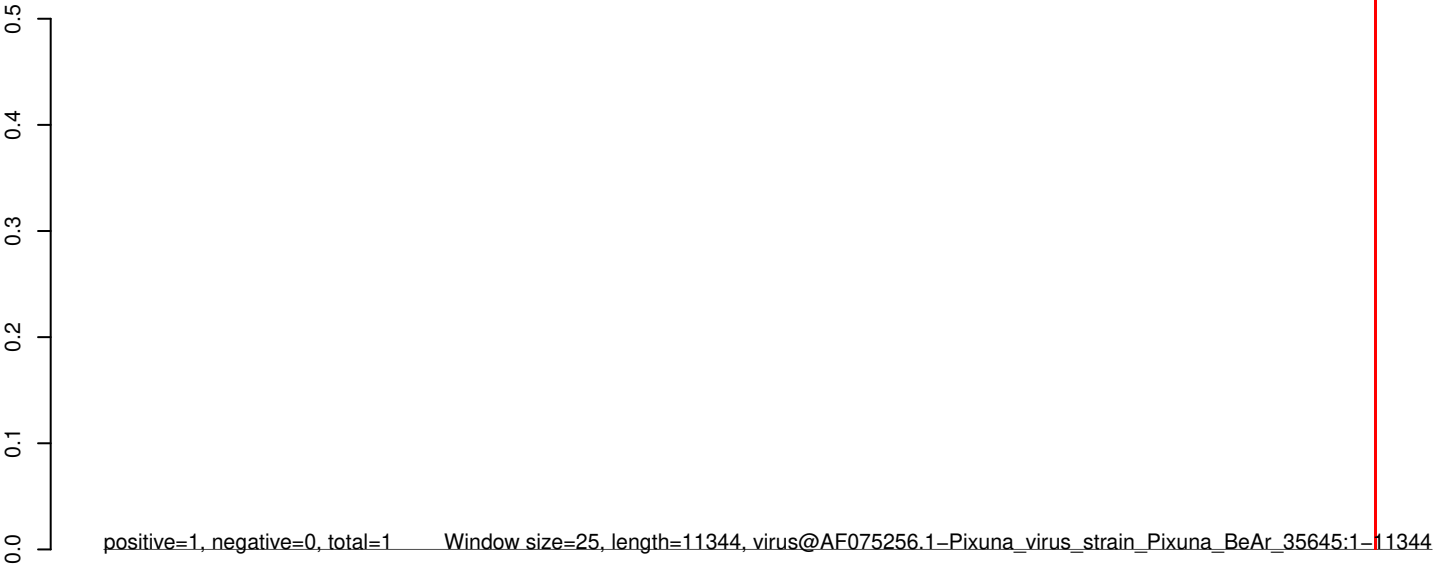
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



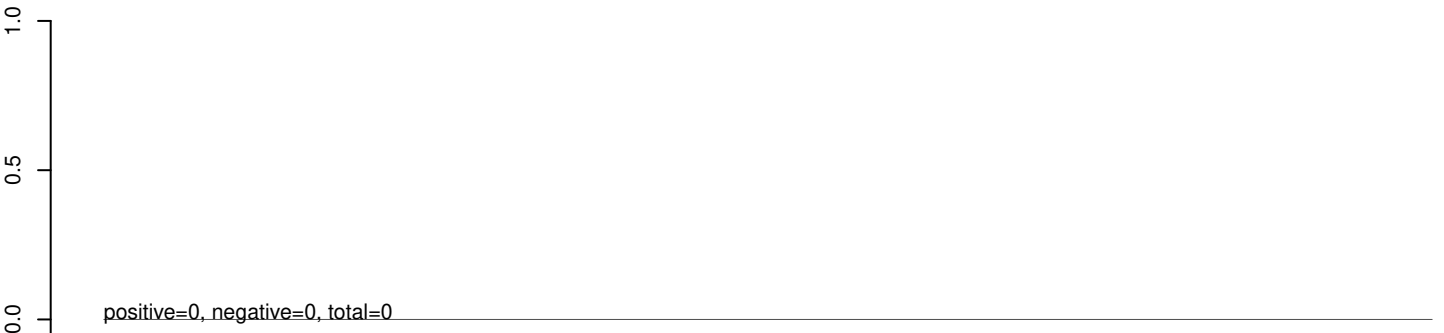
AeAeg_Aag2_mock_dsFluc_RA.rep



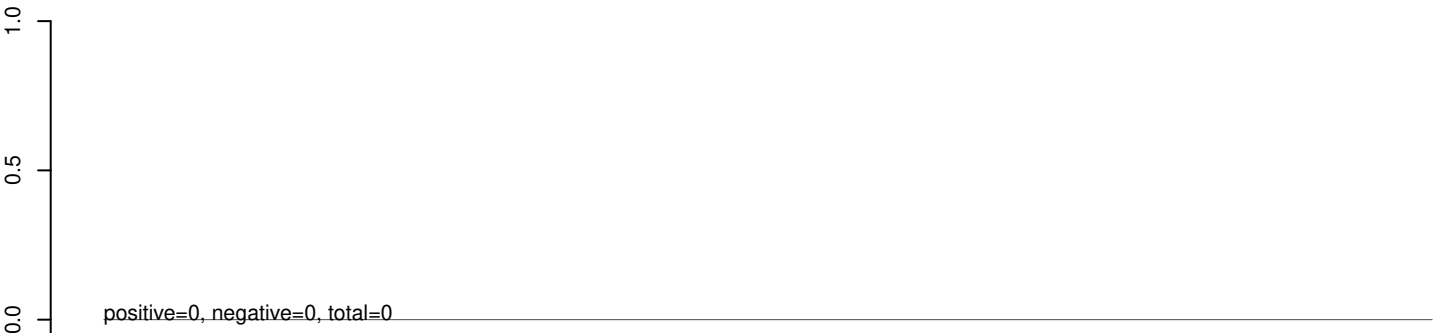
Window size=25, length=11344, virus@AF075256.1-Pixuna_virus_strain_Pixuna_BeAr_35645:1-11344

0 2000 4000 6000 8000 10000

AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep

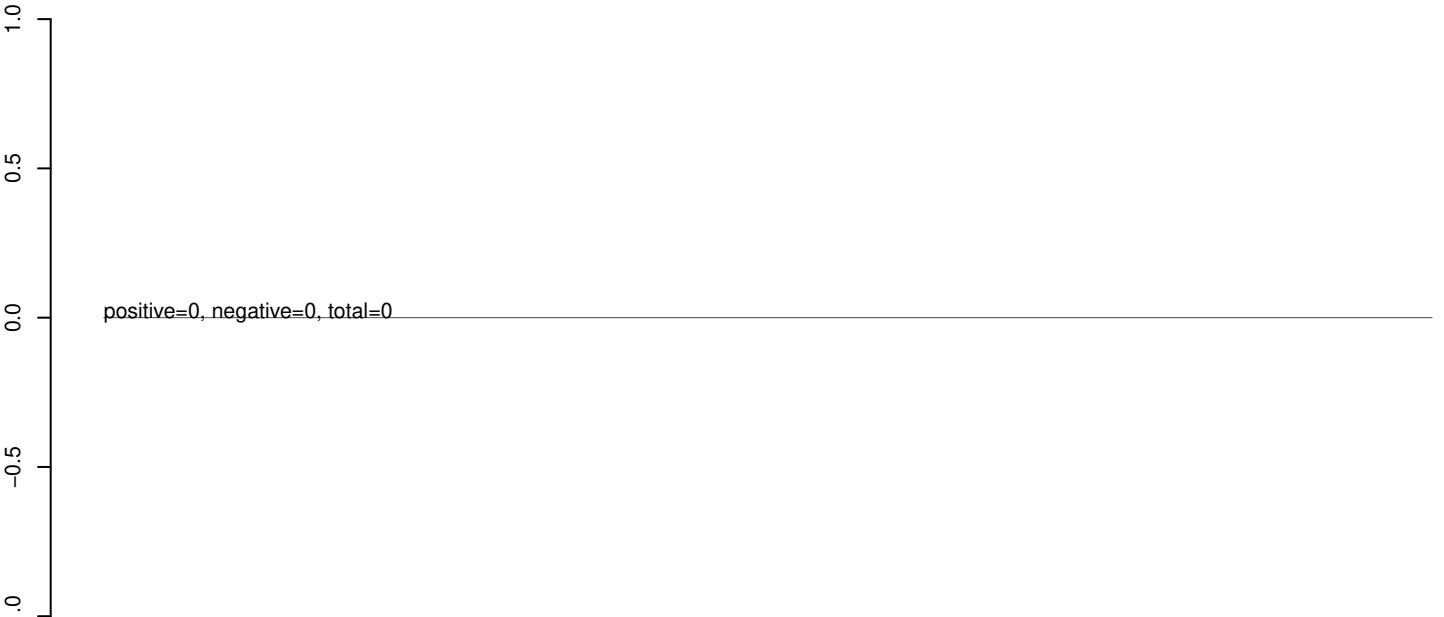


0 5000 10000 15000 20000

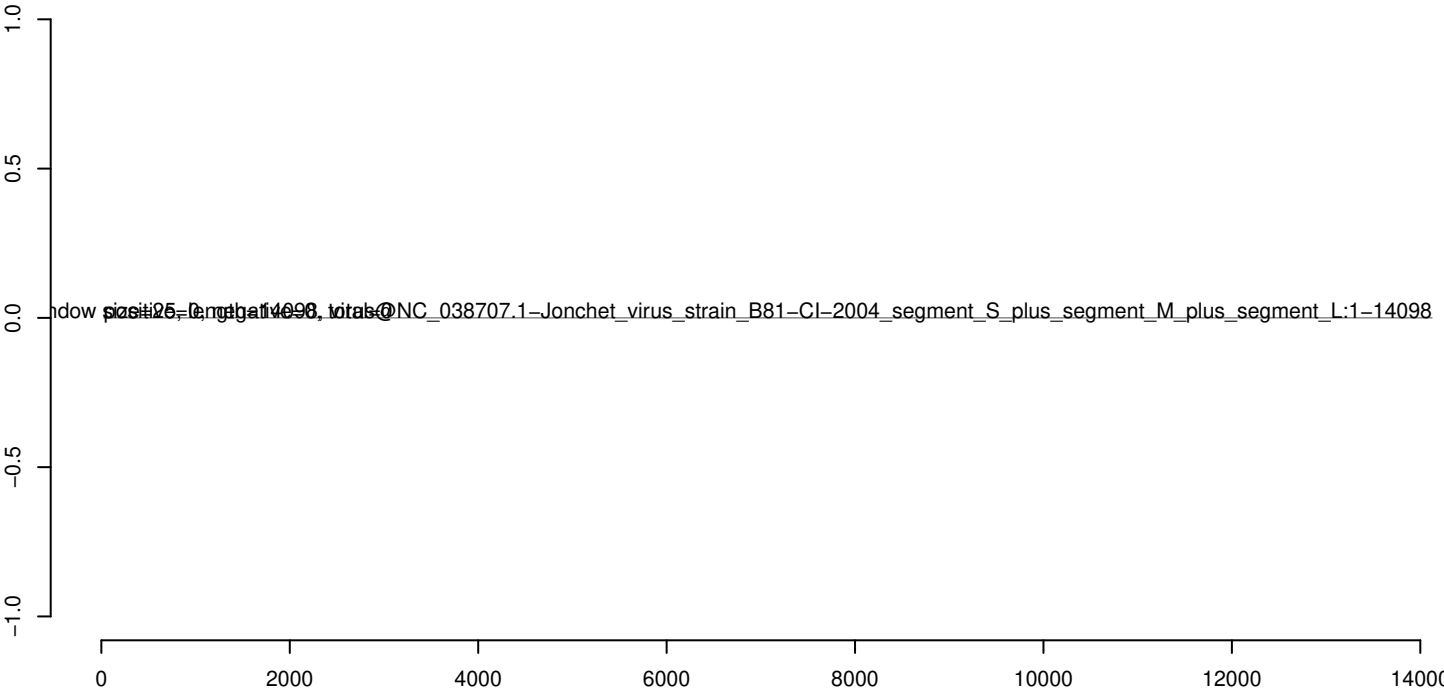
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



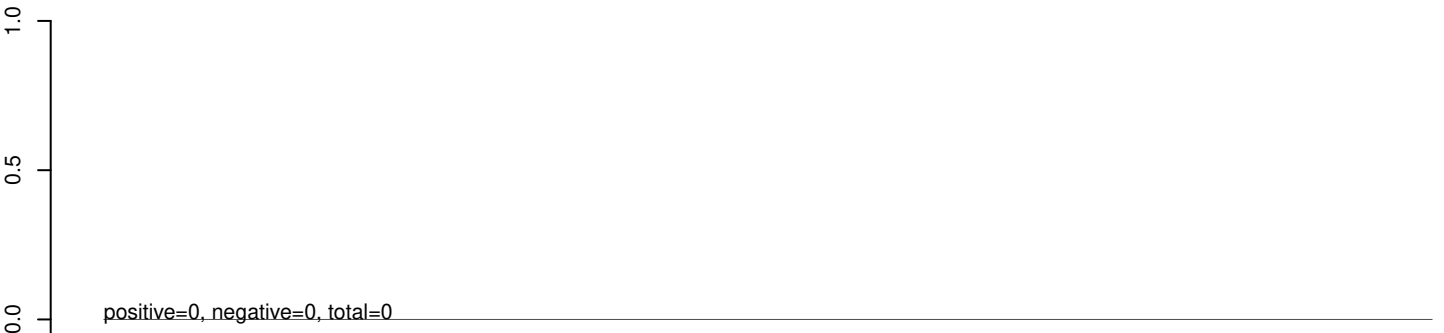
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



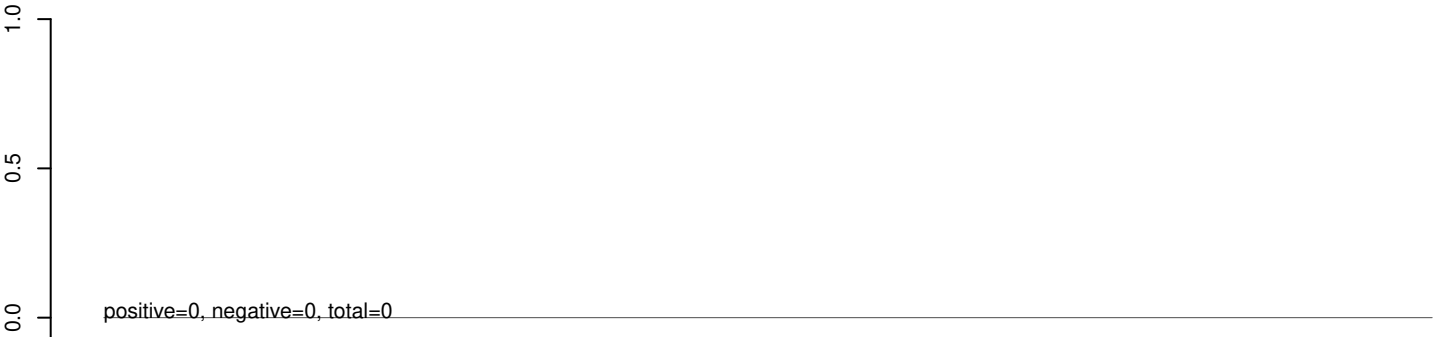
AeAeg_Aag2_mock_dsFluc_RA.rep



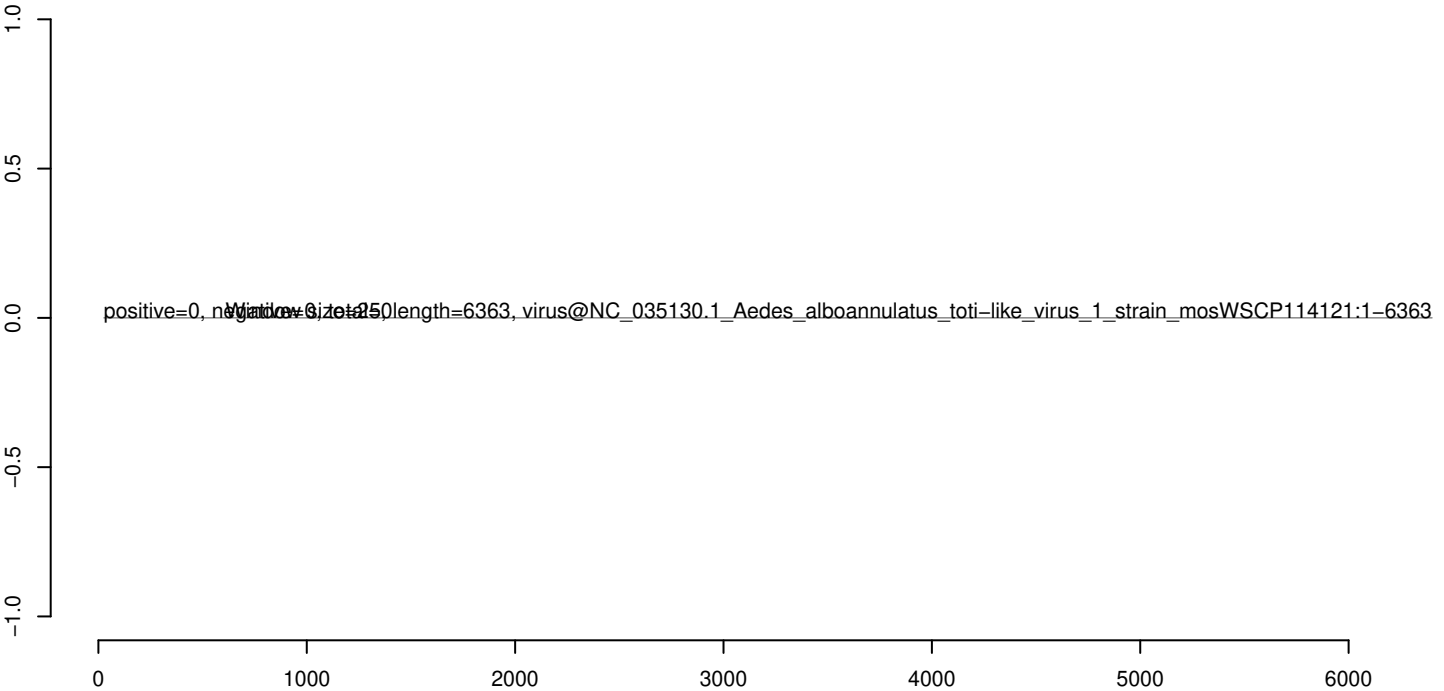
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



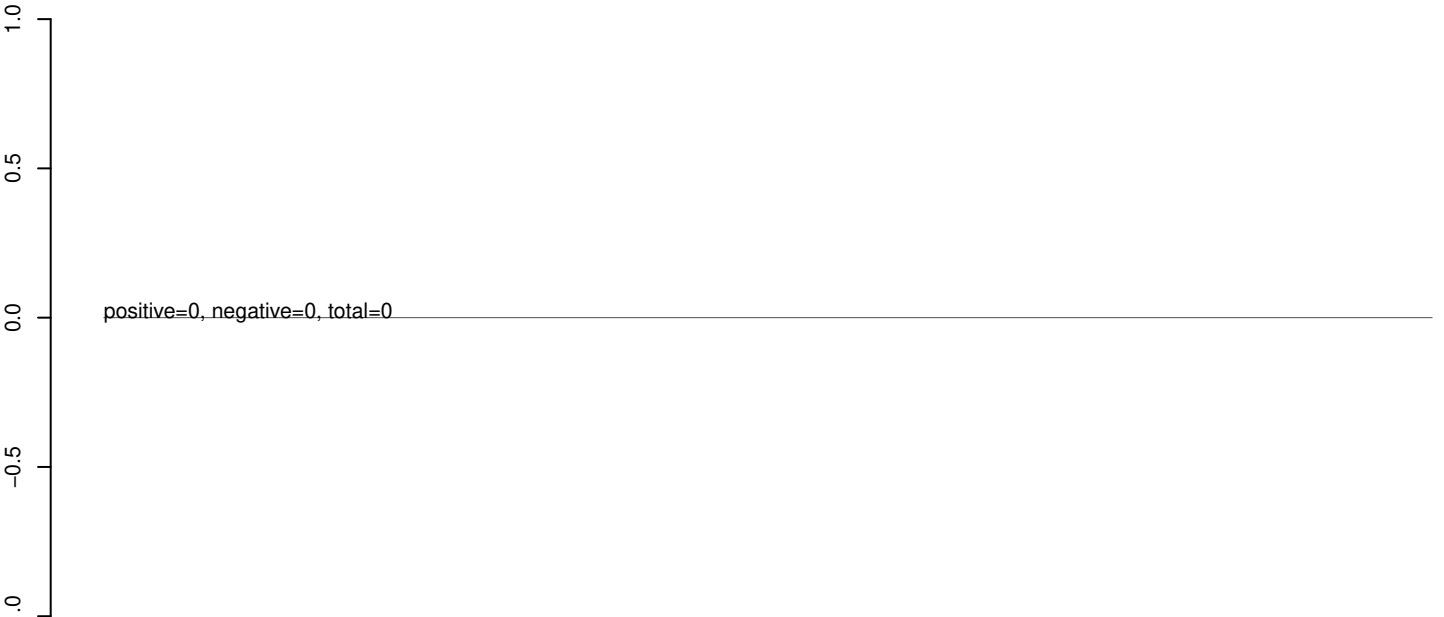
AeAeg_Aag2_mock_dsFluc_RA.rep



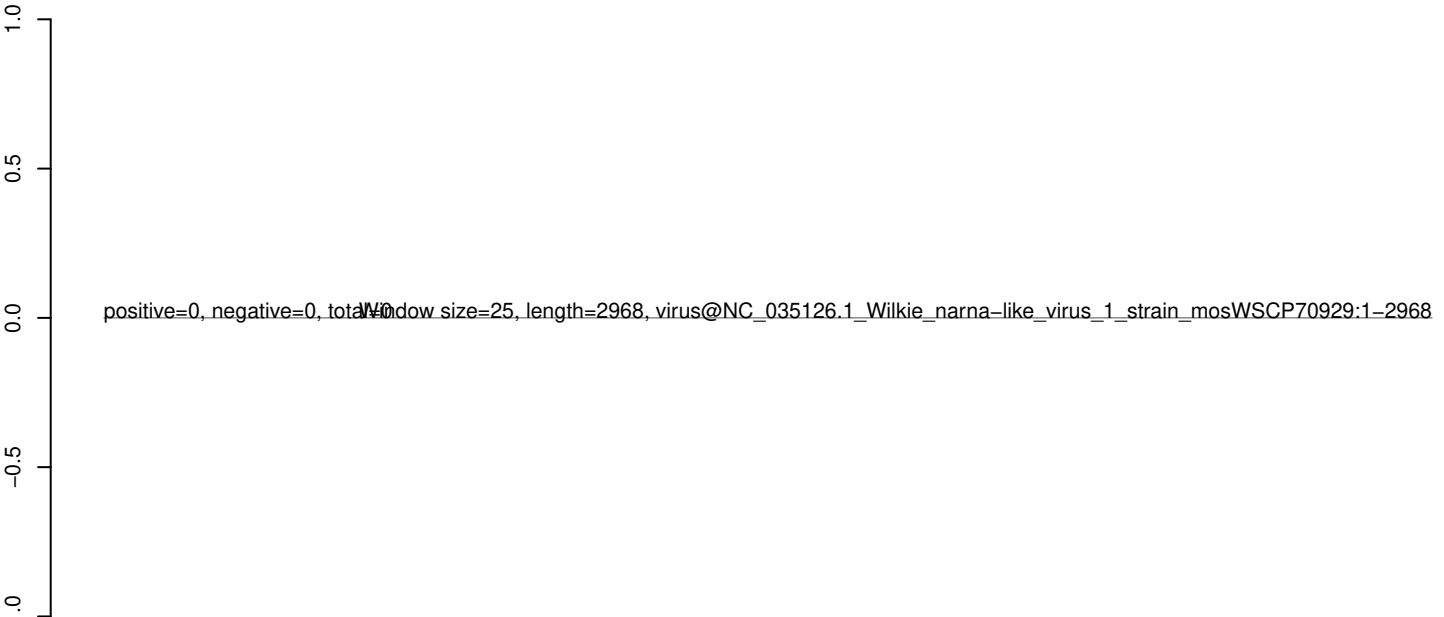
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep

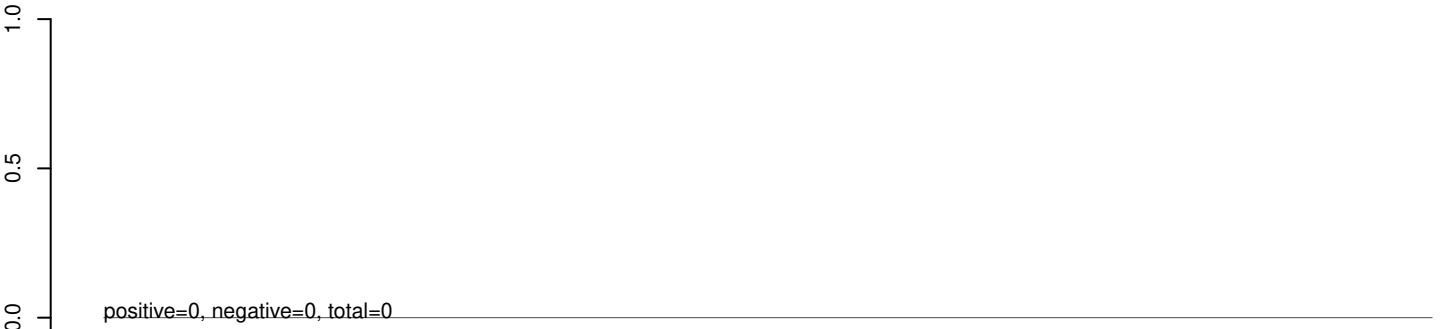


0 500 1000 1500 2000 2500 3000

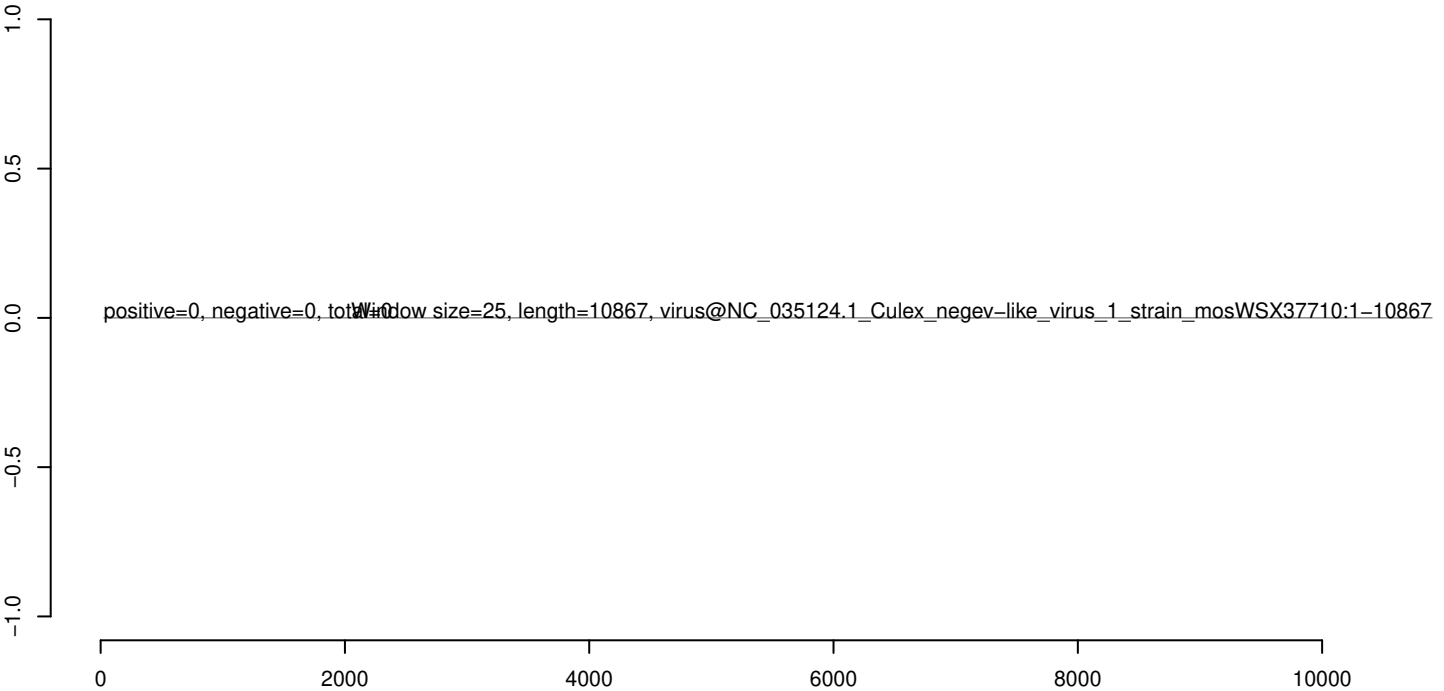
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep



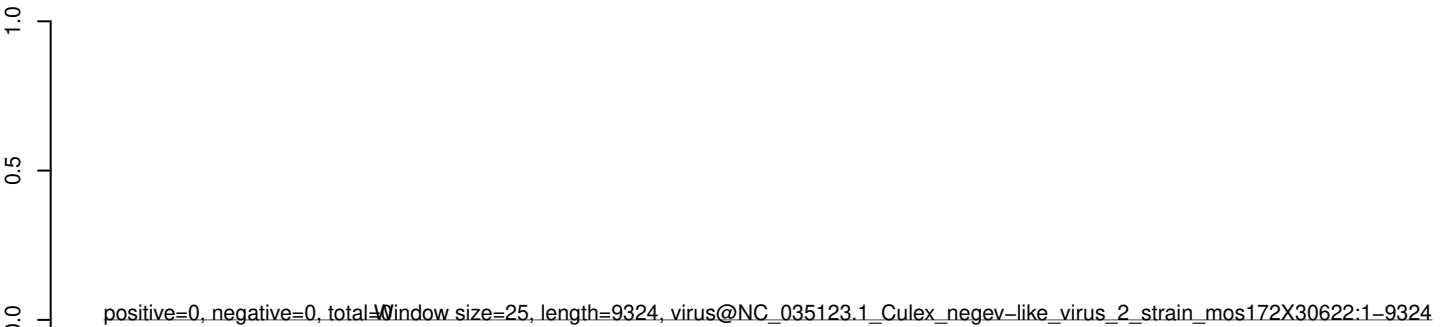
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



AeAeg_Aag2_mock_dsFluc_RA.rep

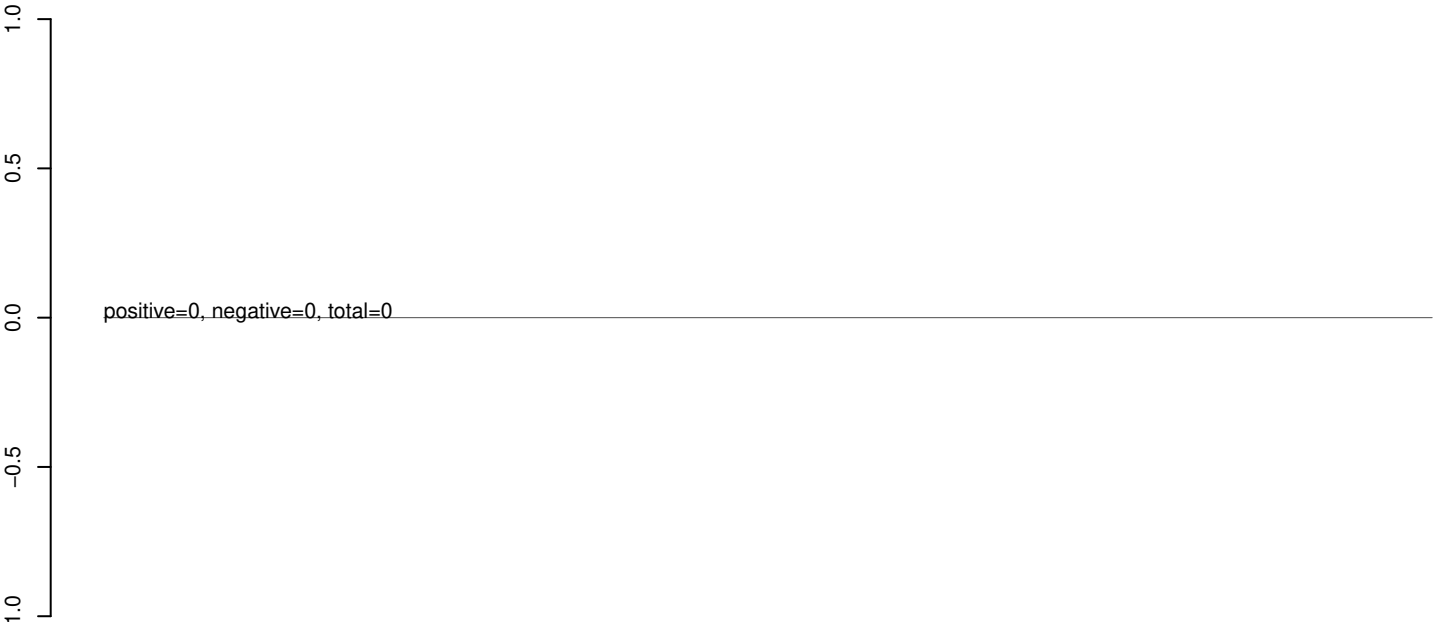


0 2000 4000 6000 8000

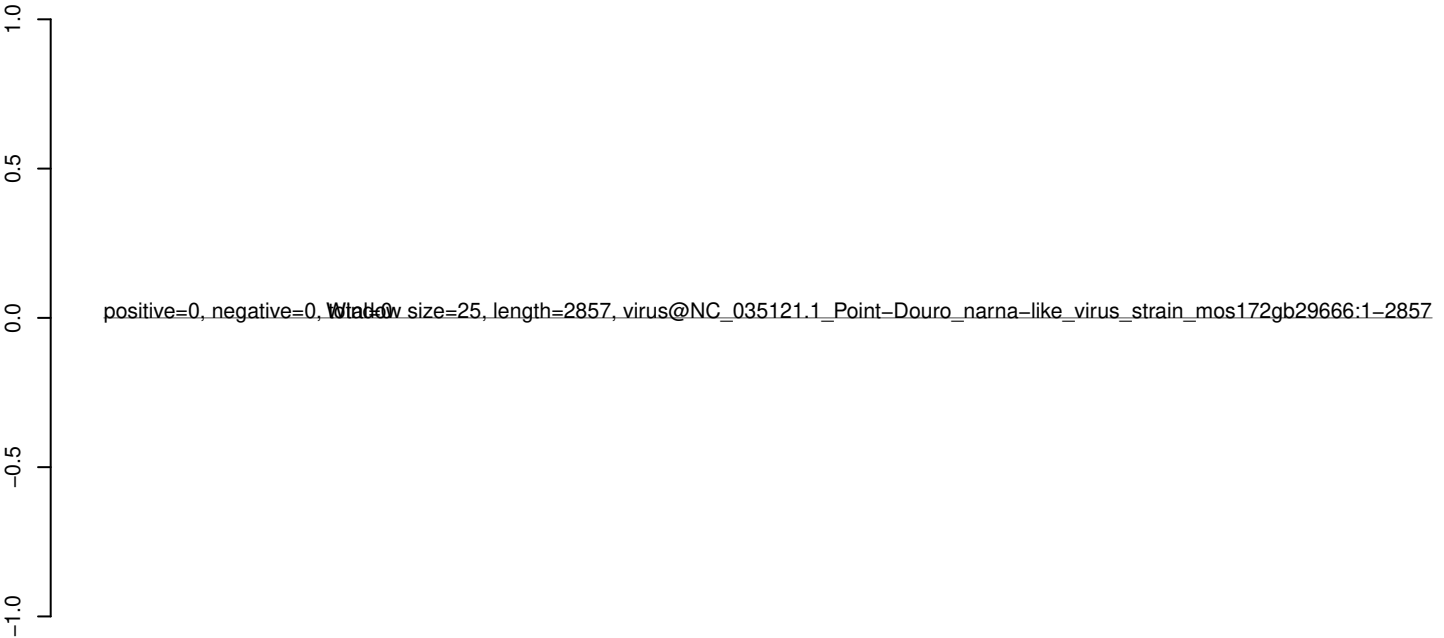
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



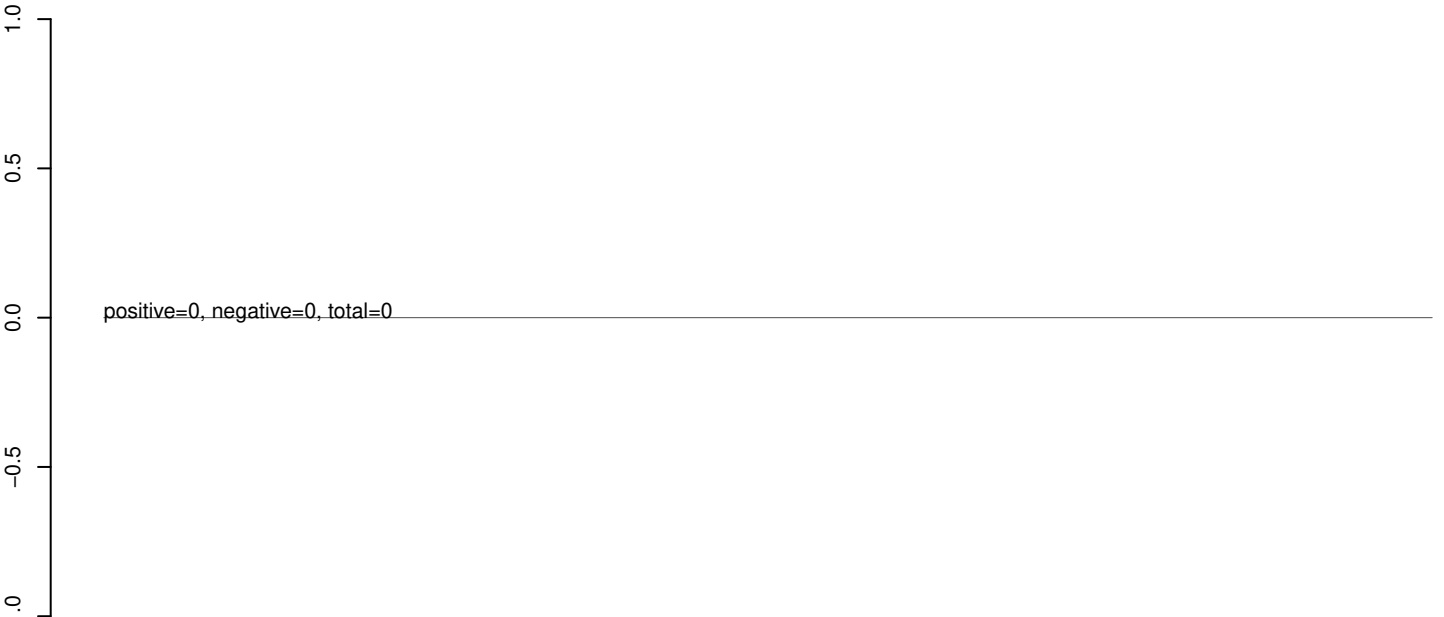
AeAeg_Aag2_mock_dsFluc_RA.rep



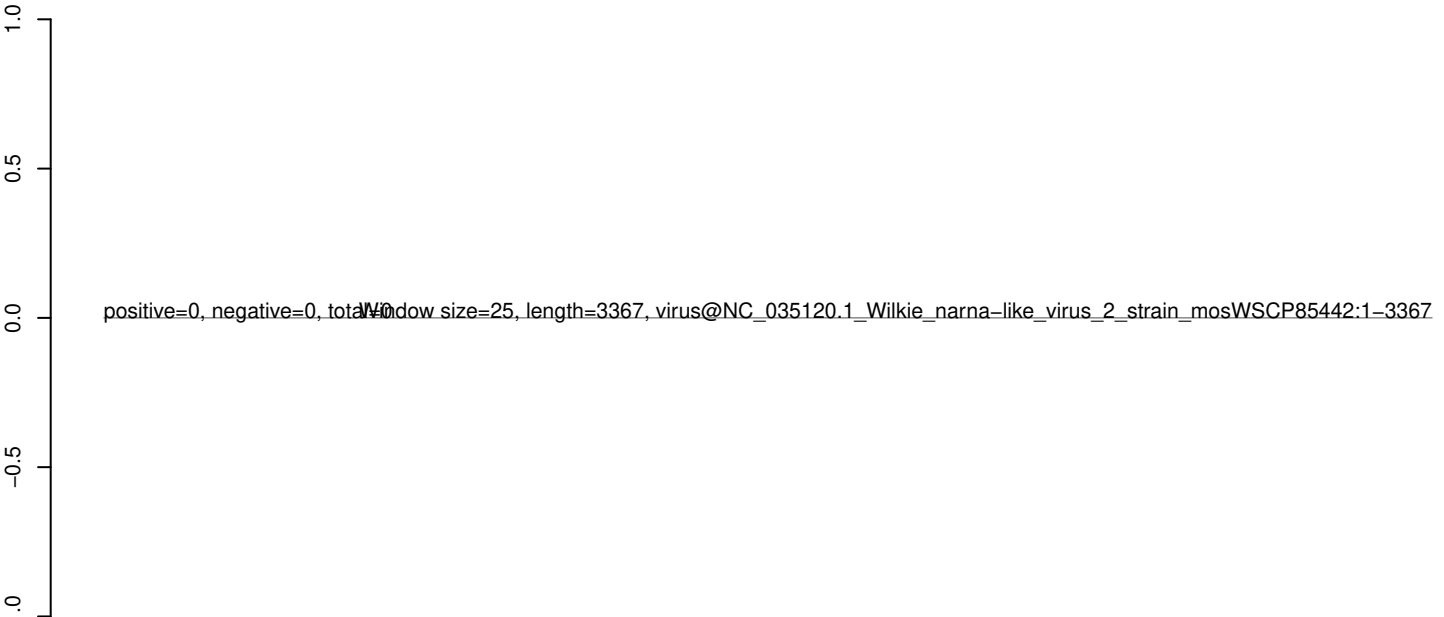
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



AeAeg_Aag2_mock_dsFluc_RA.24_35.rep

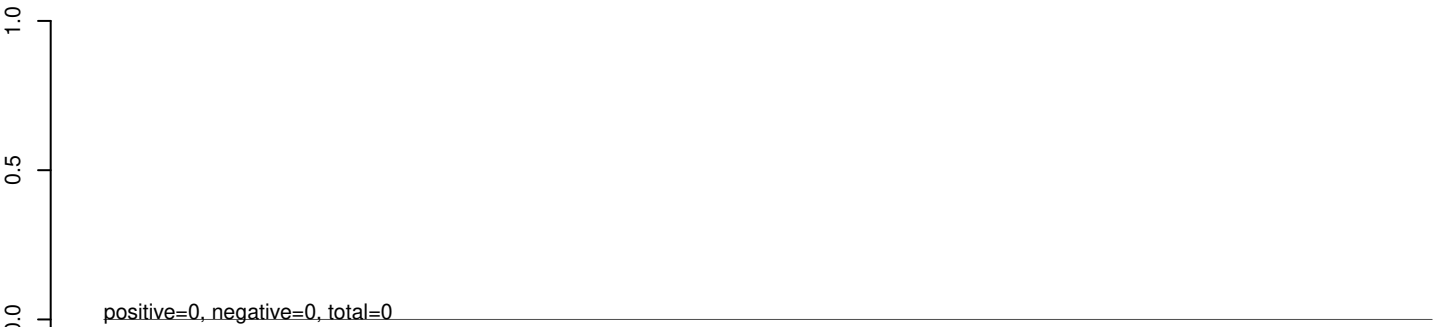


AeAeg_Aag2_mock_dsFluc_RA.rep

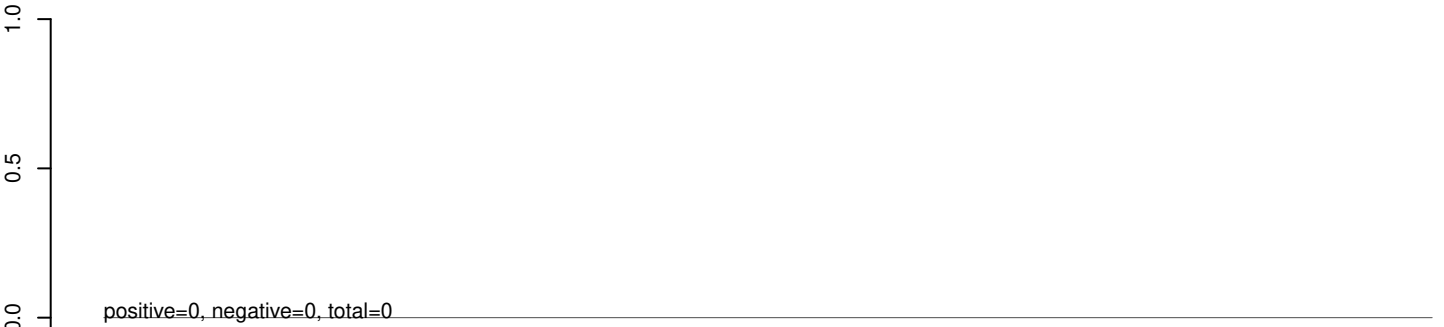


0 500 1000 1500 2000 2500 3000 3500

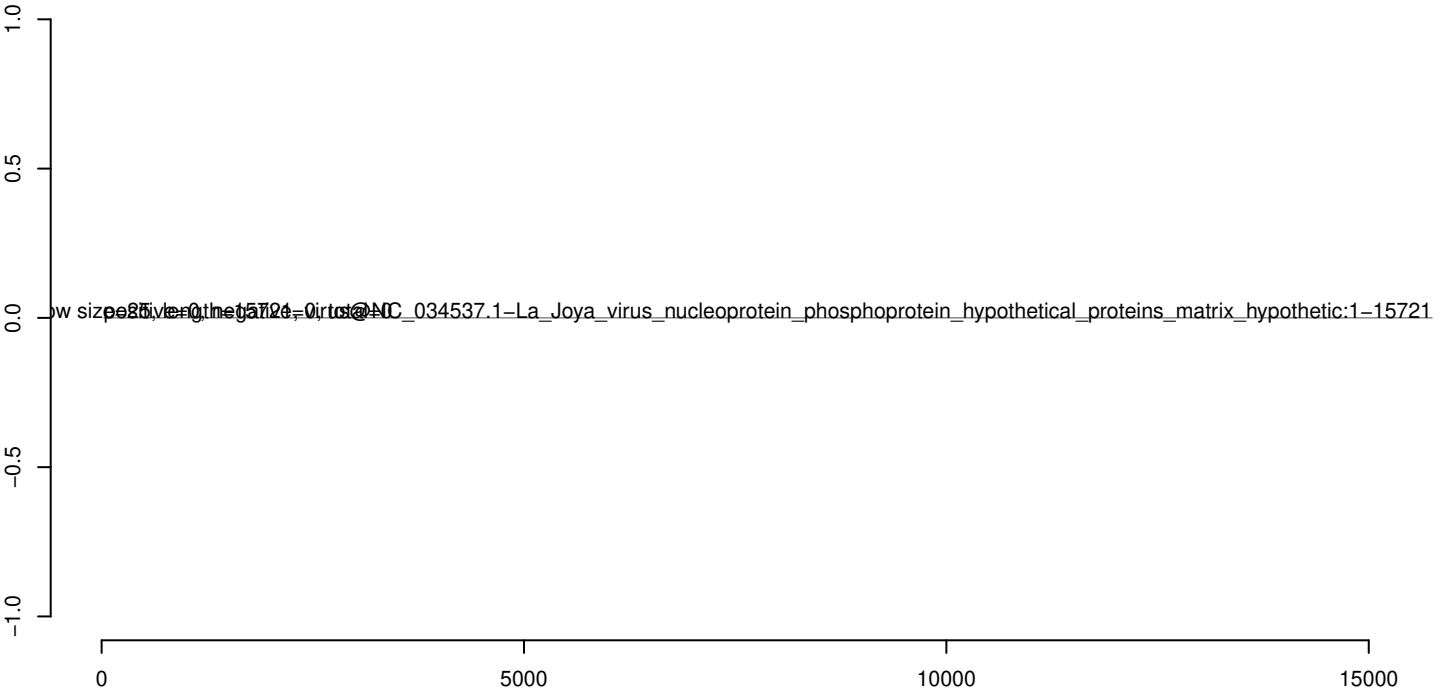
AeAeg_Aag2_mock_dsFluc_RA.18_23.rep



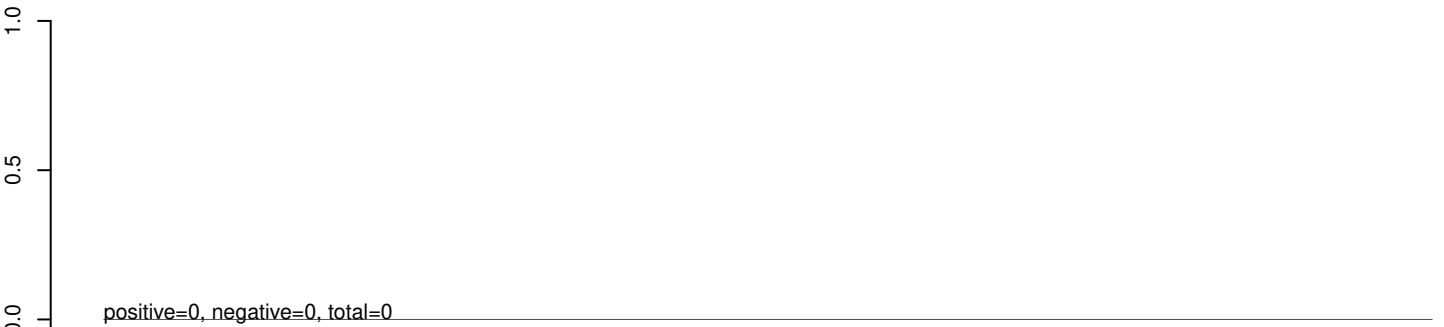
AeAeg_Aag2_mock_dsFluc_RA.24_35.rep



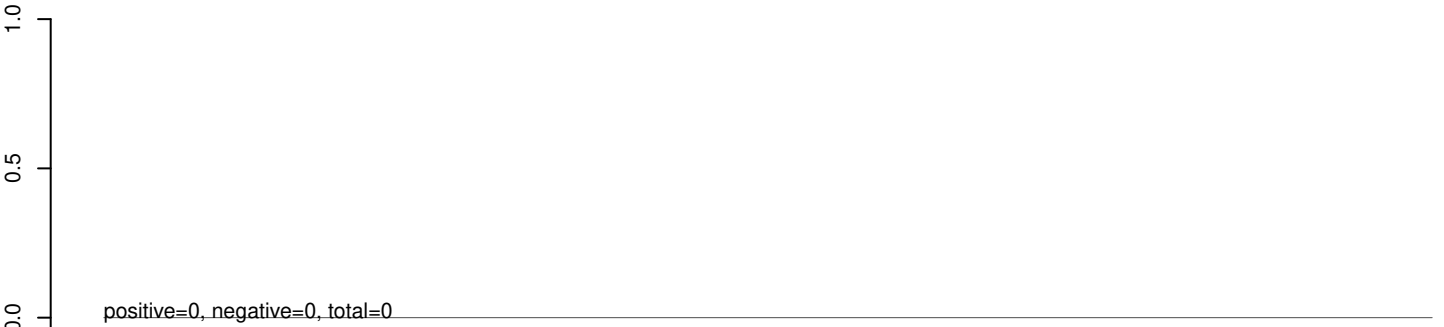
AeAeg_Aag2_mock_dsFluc_RA.rep



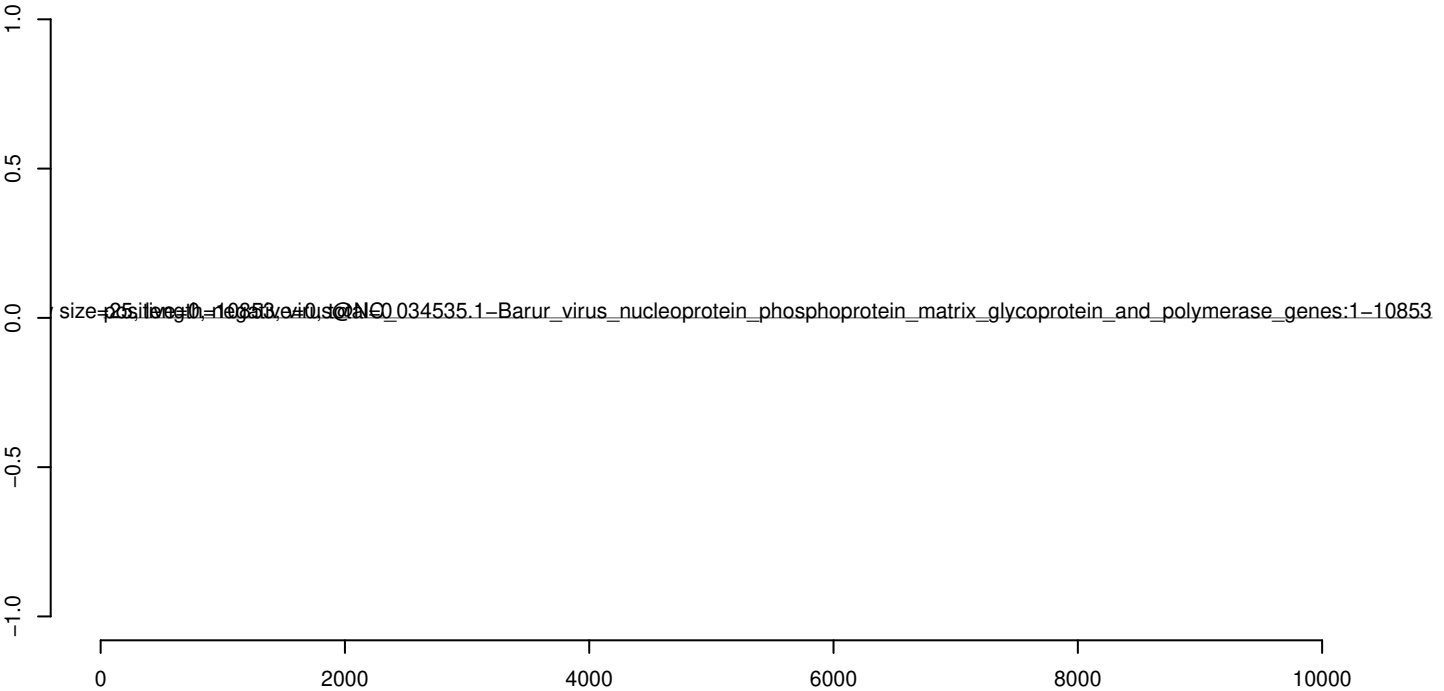
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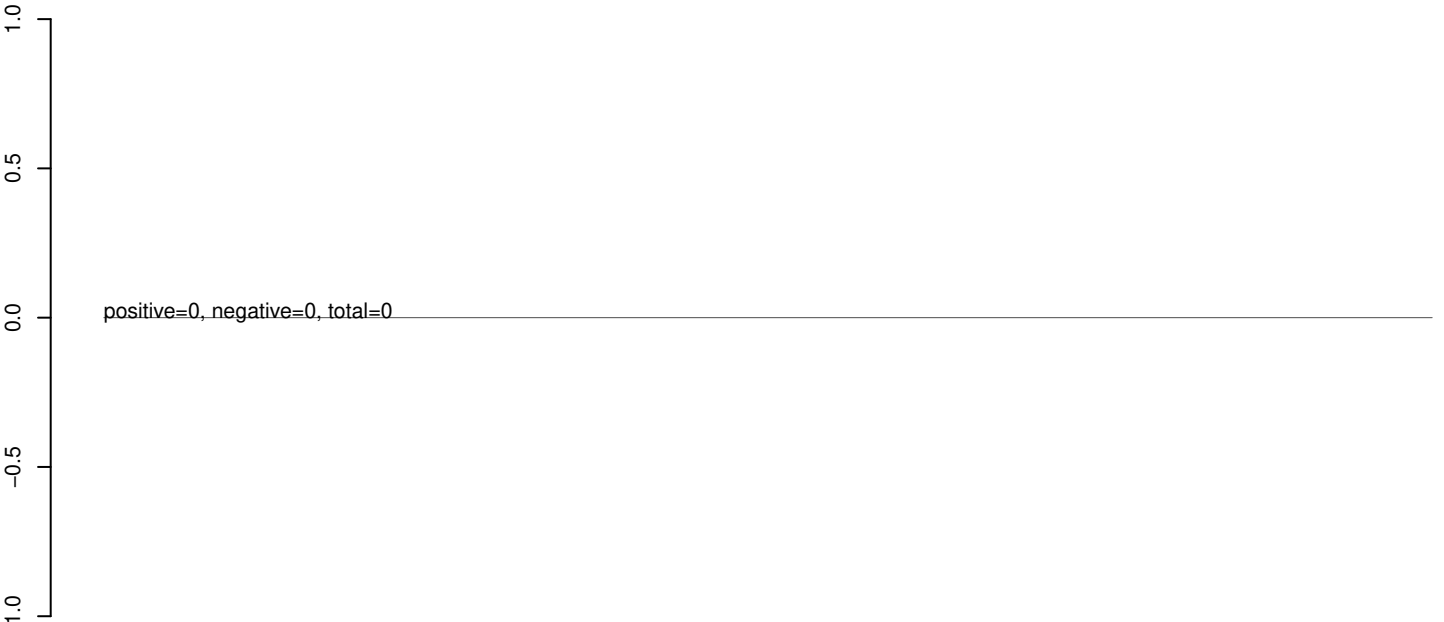
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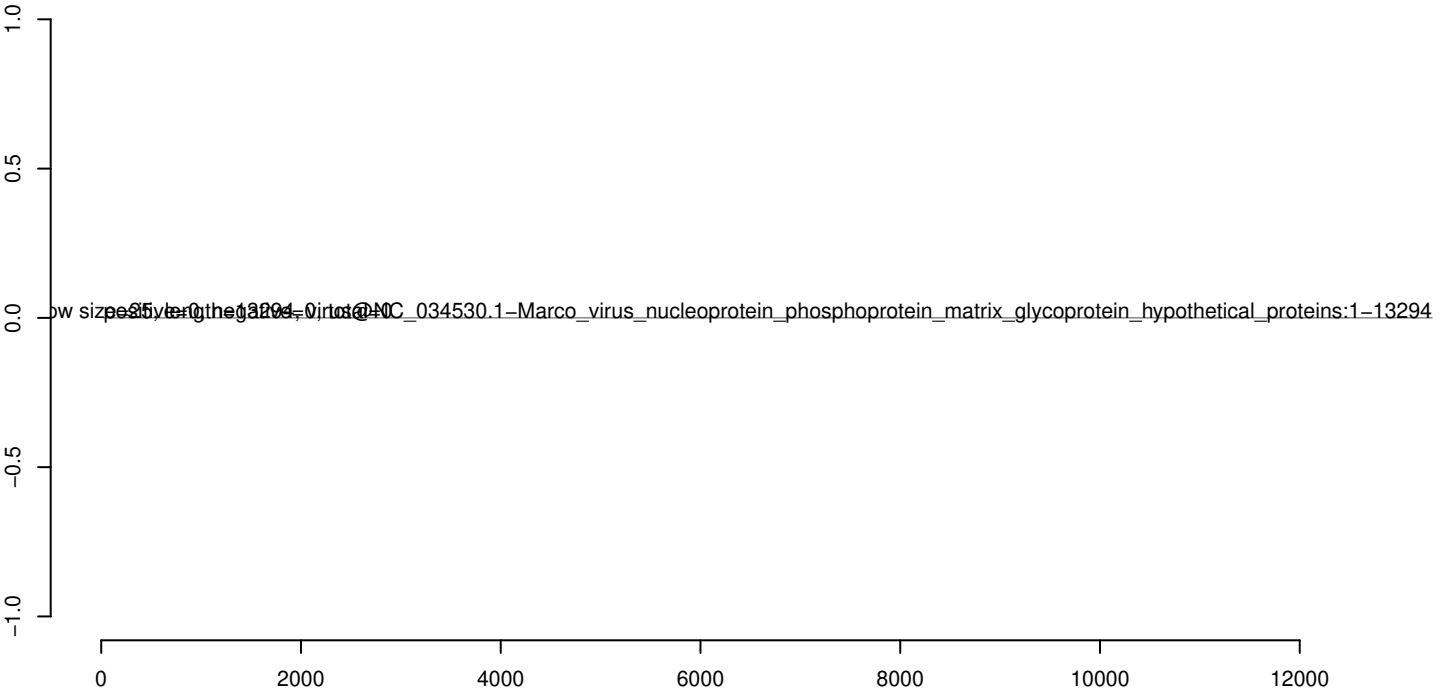
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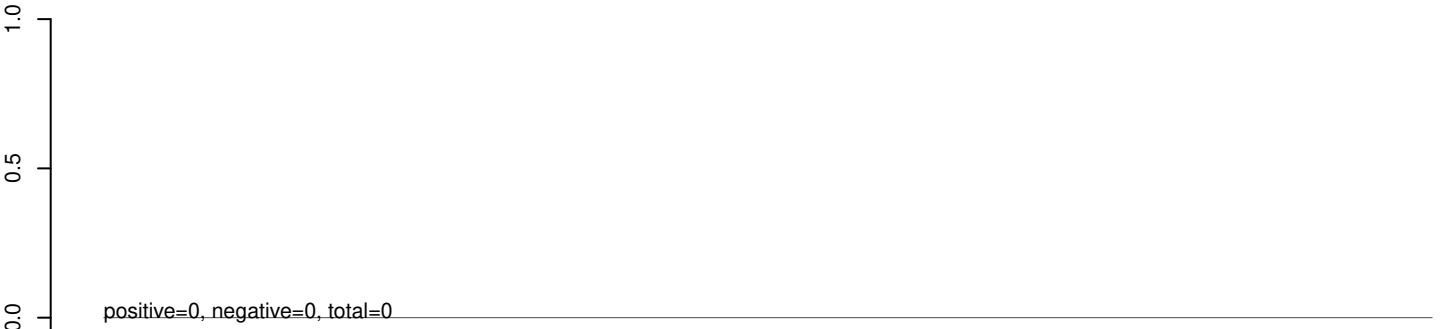
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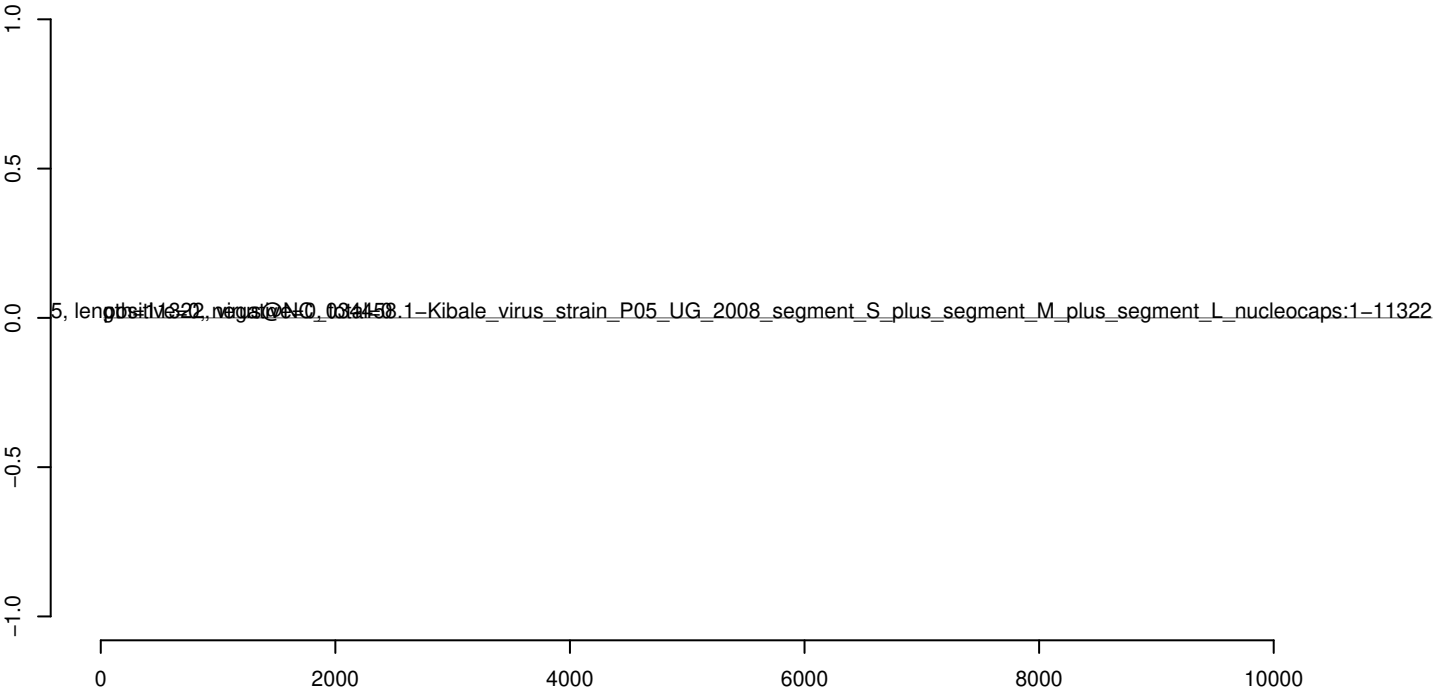
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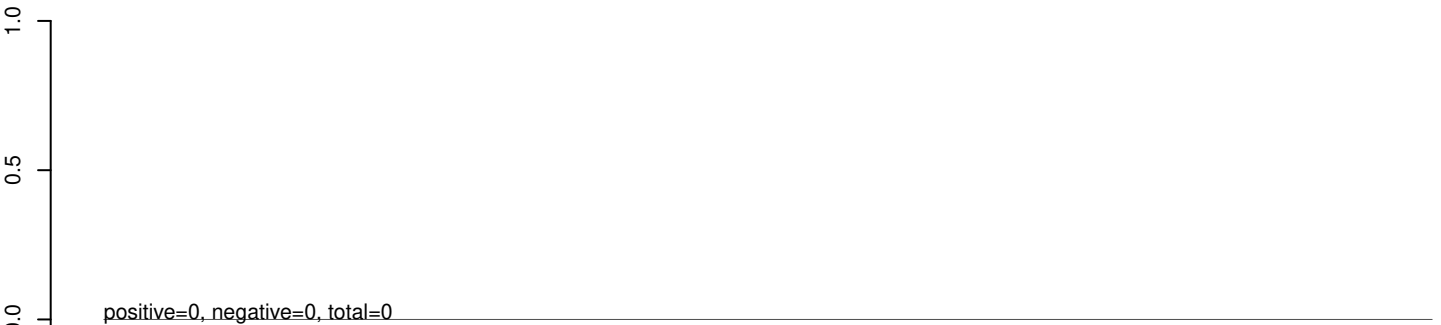
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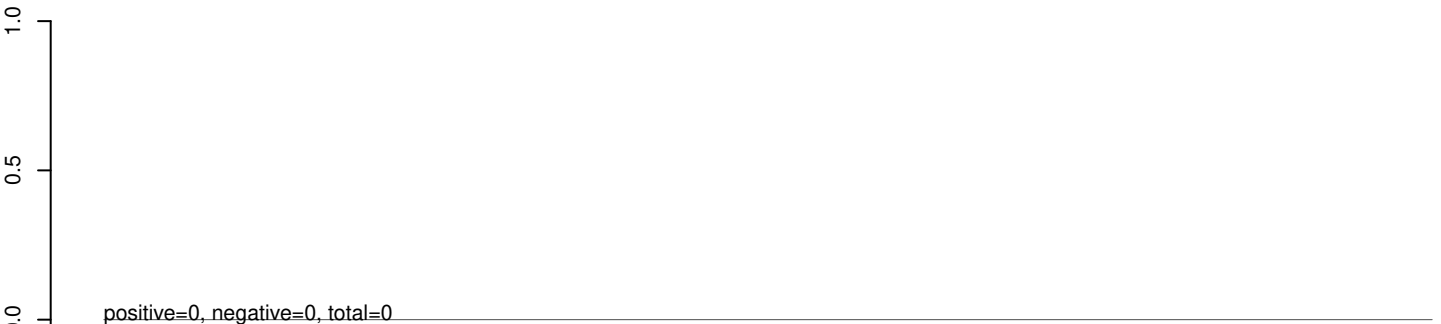
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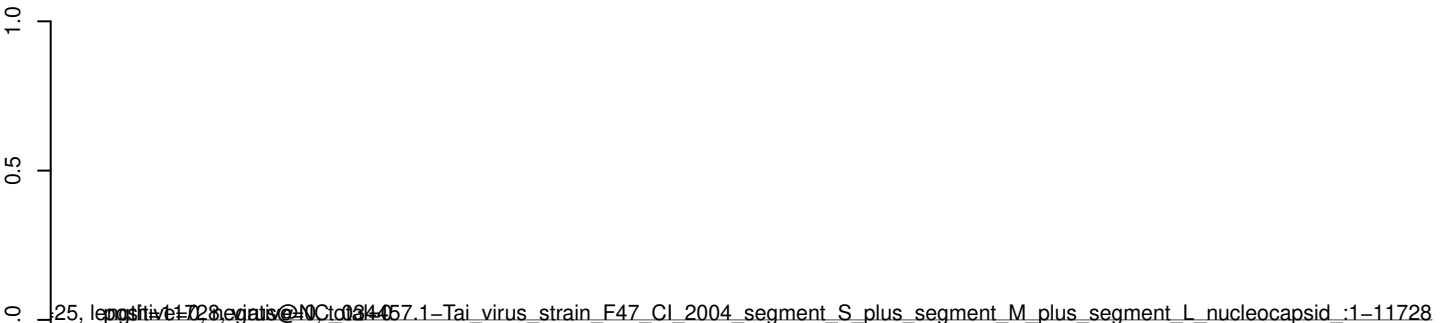
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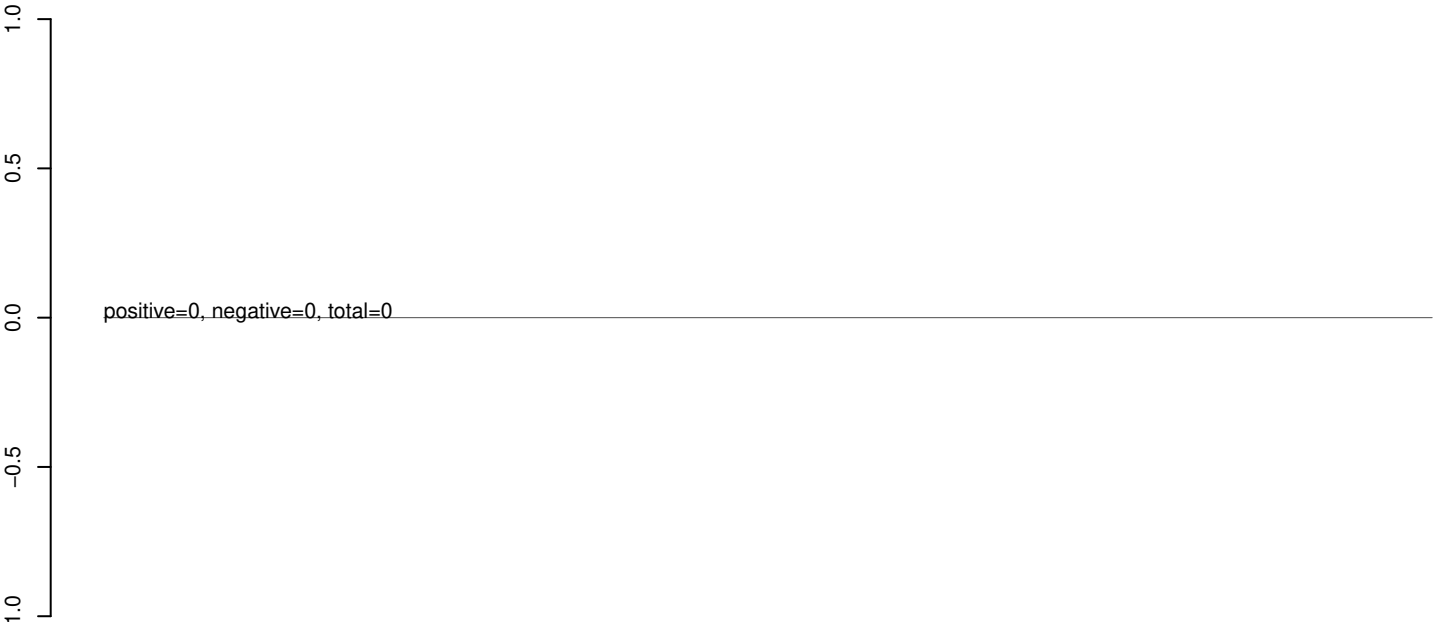
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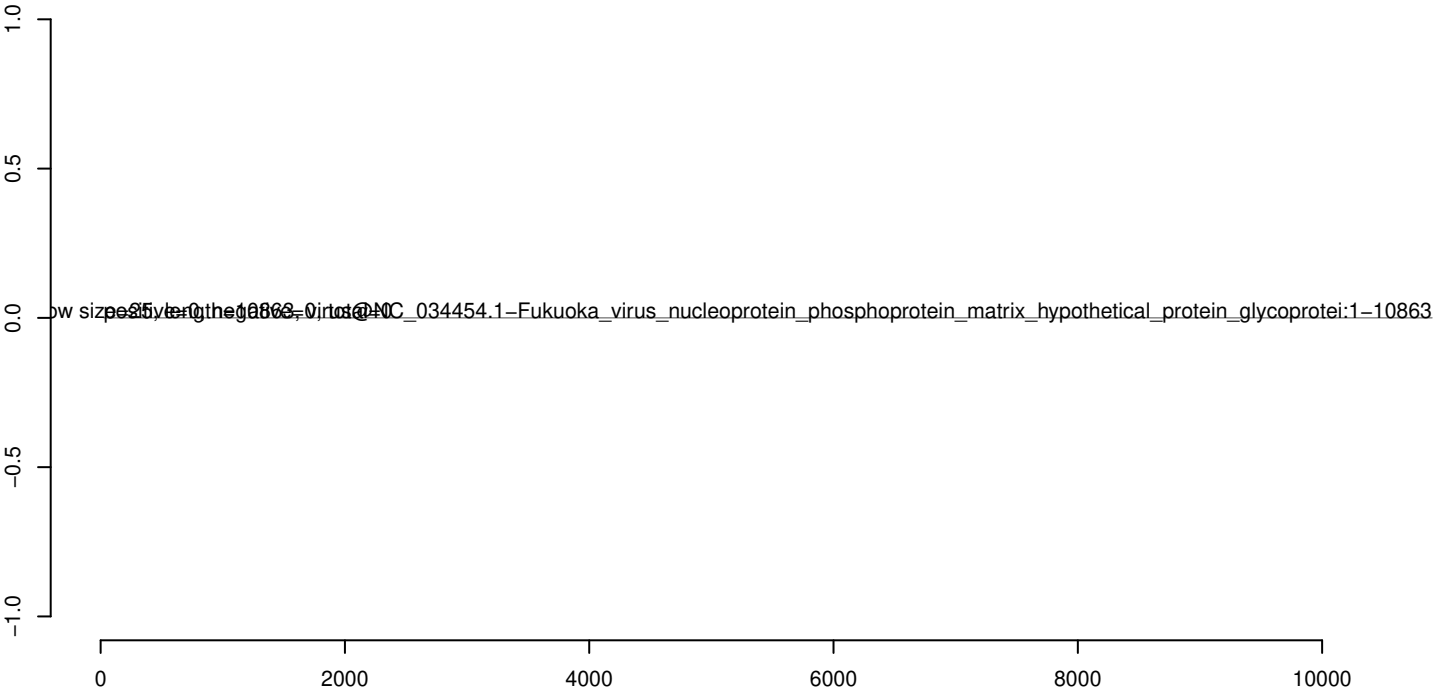
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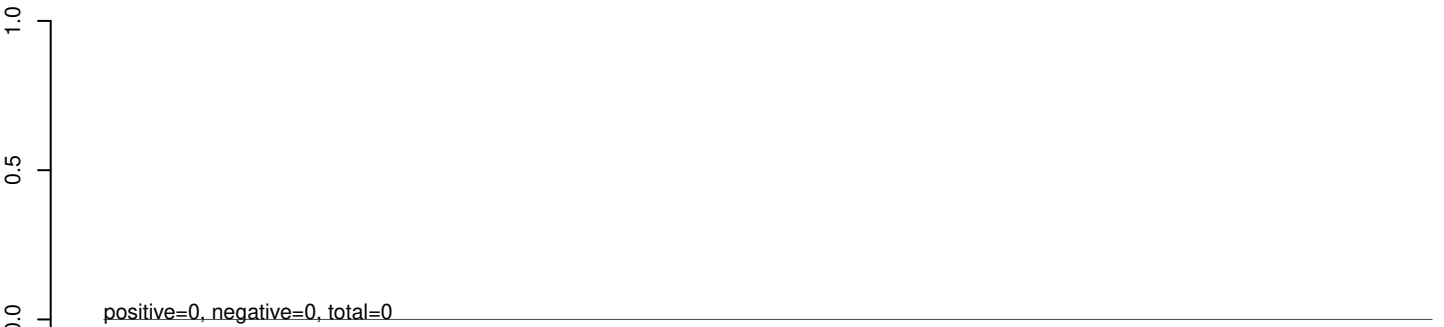
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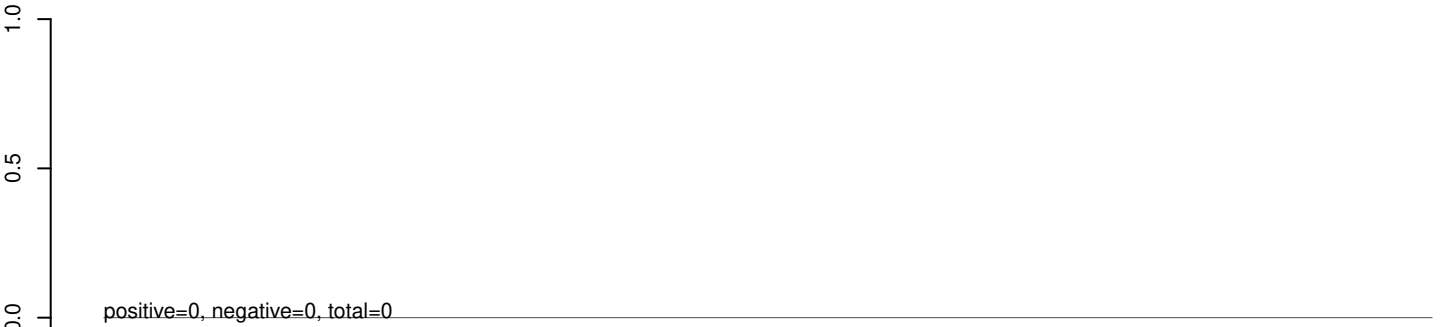
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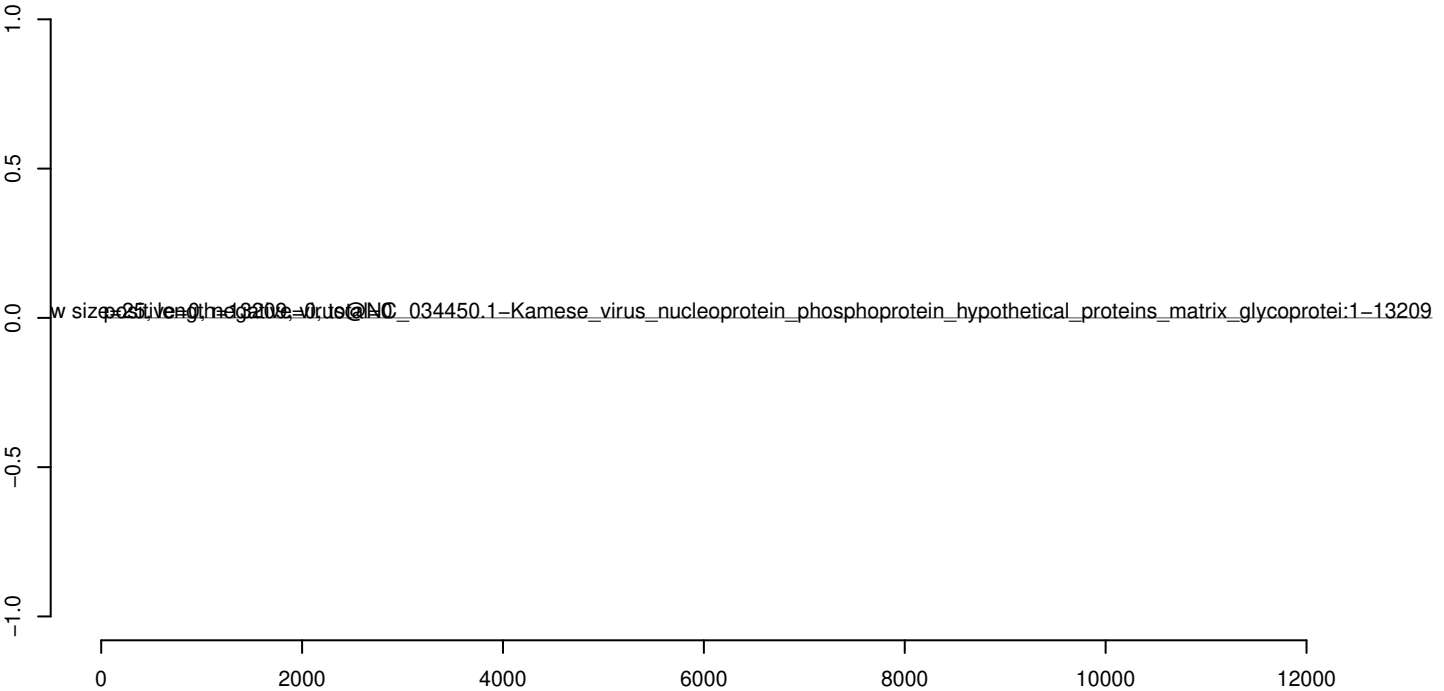
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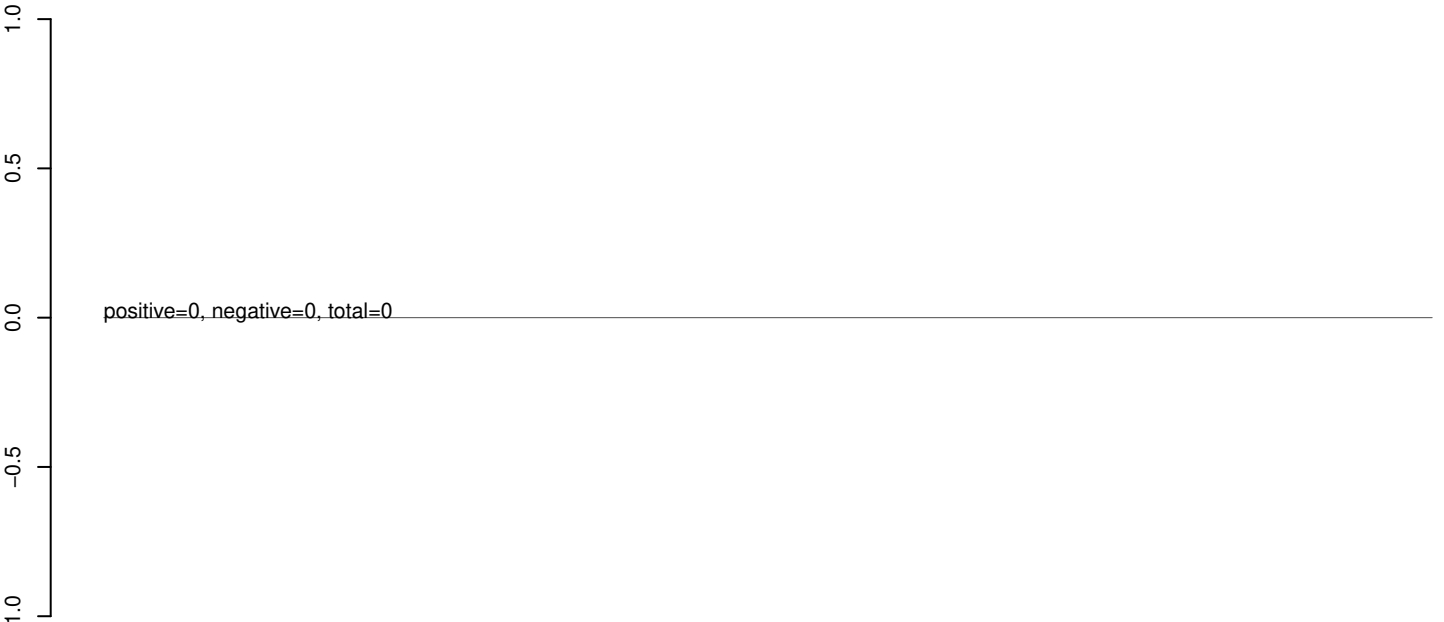
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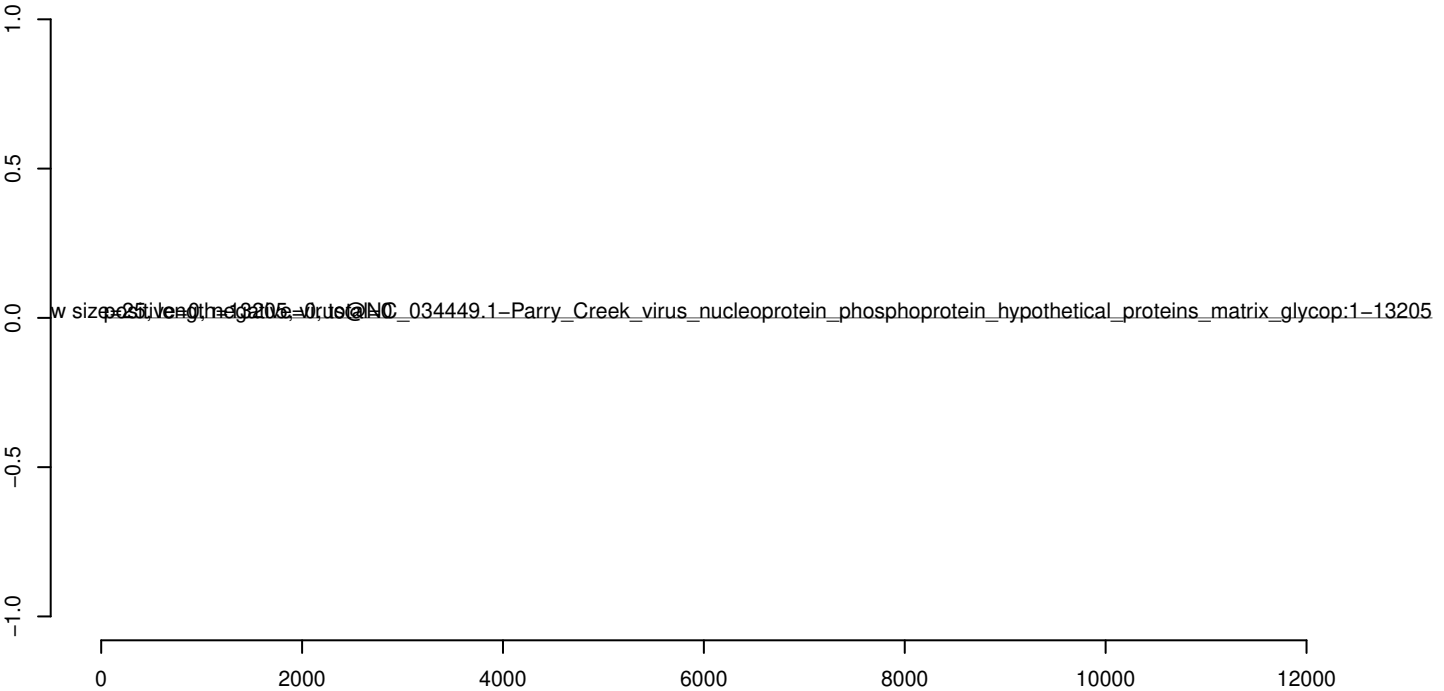
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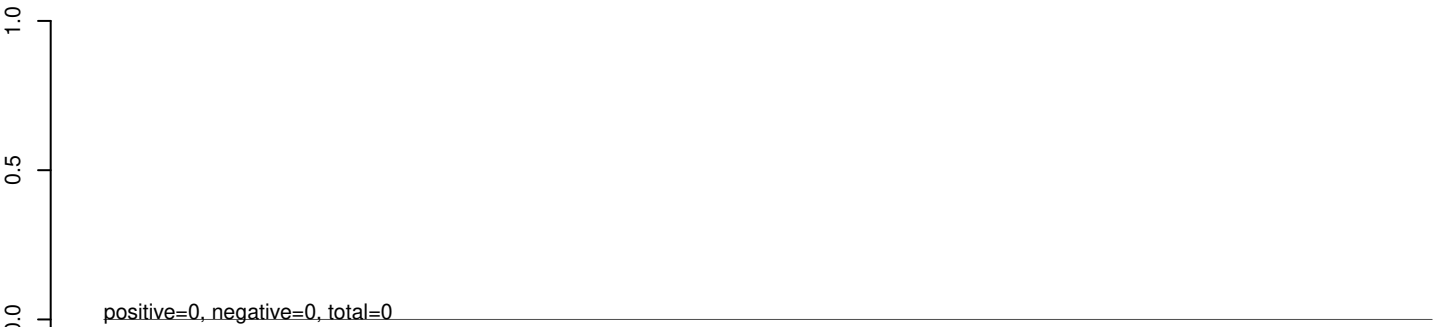
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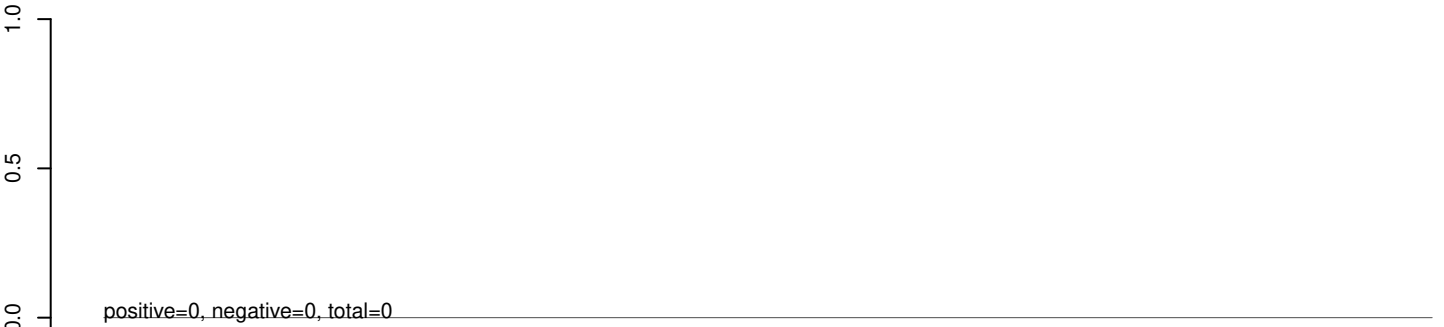
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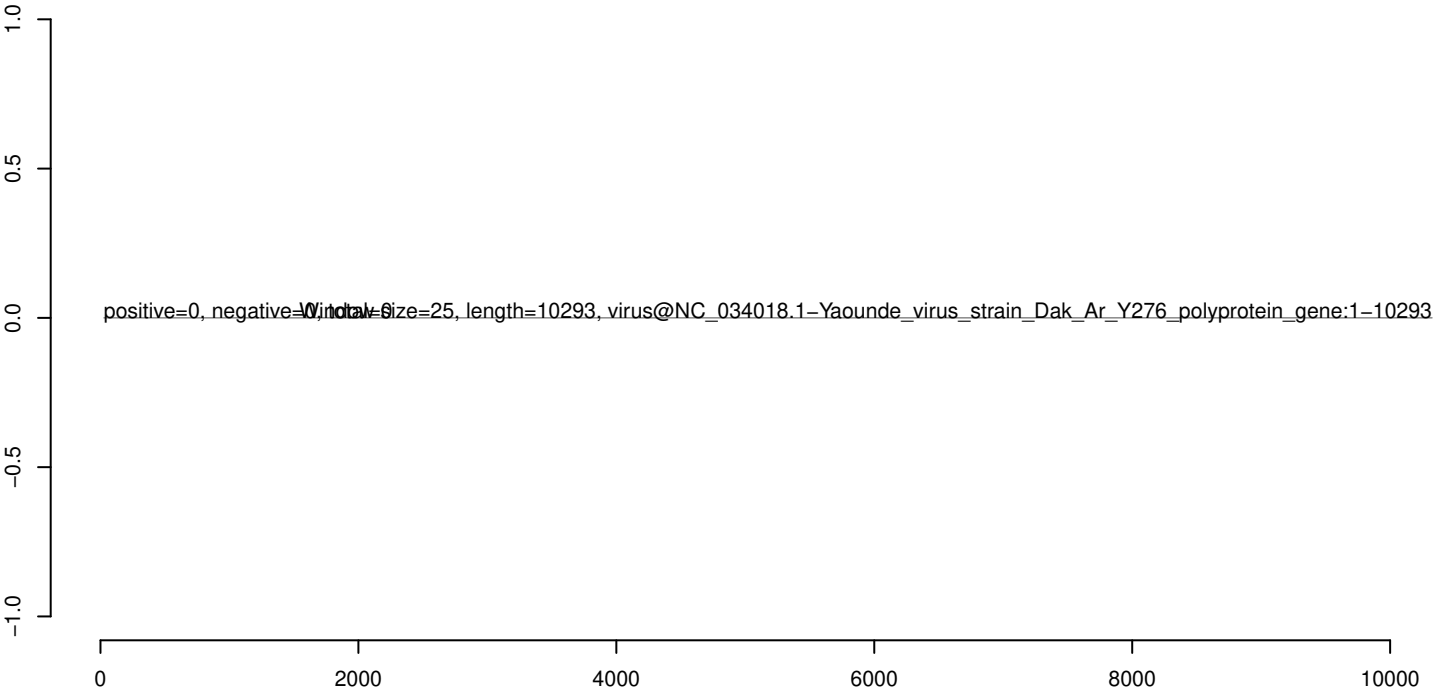
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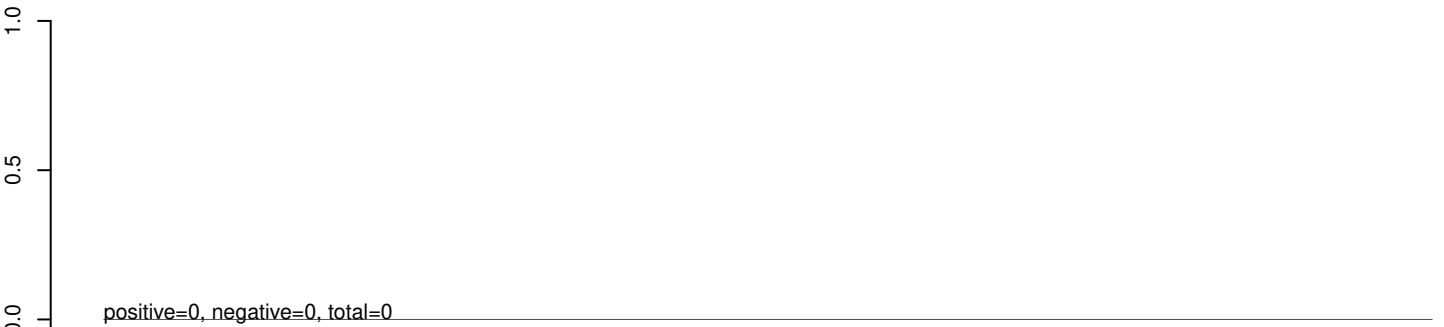
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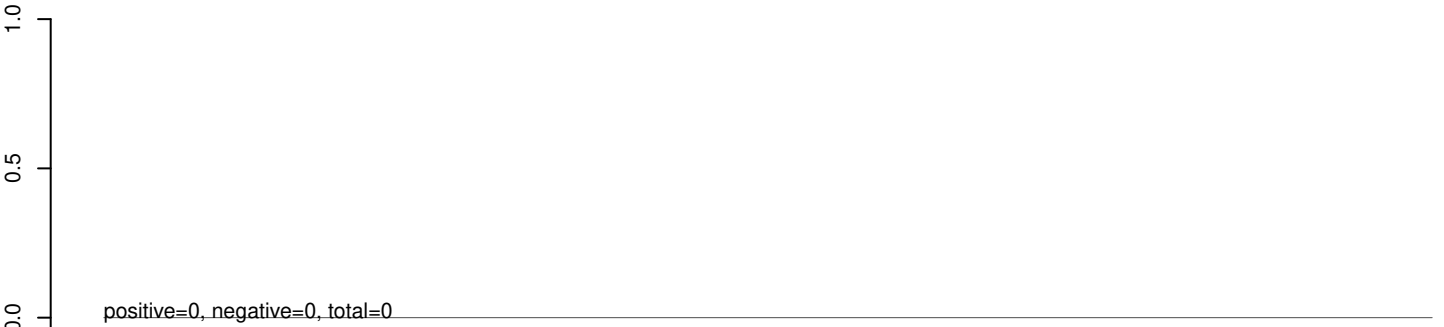
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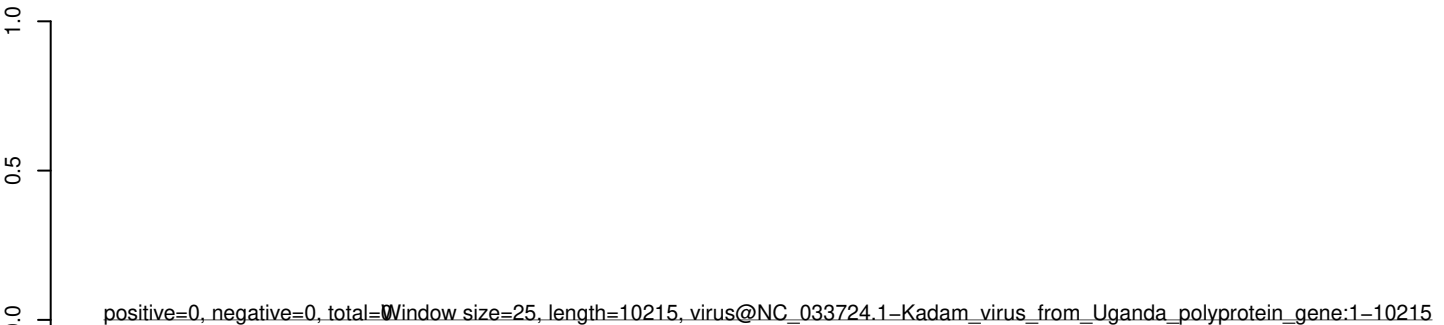
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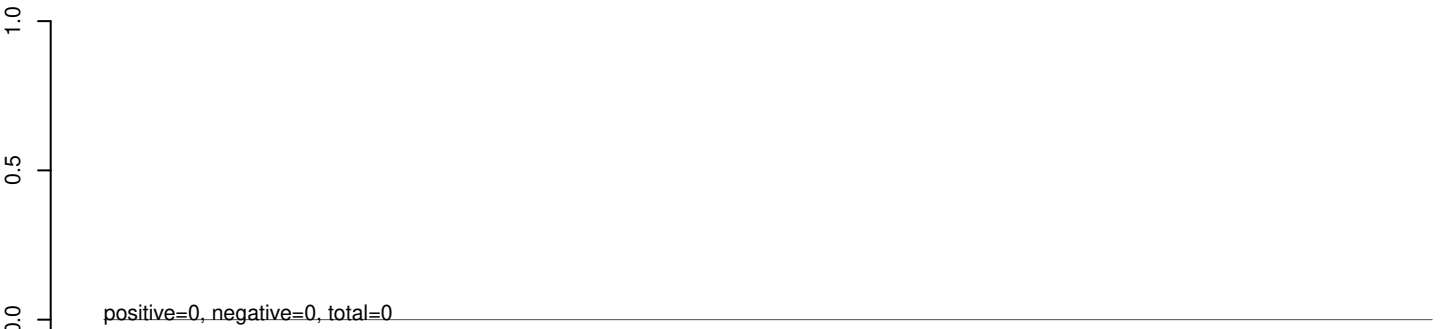


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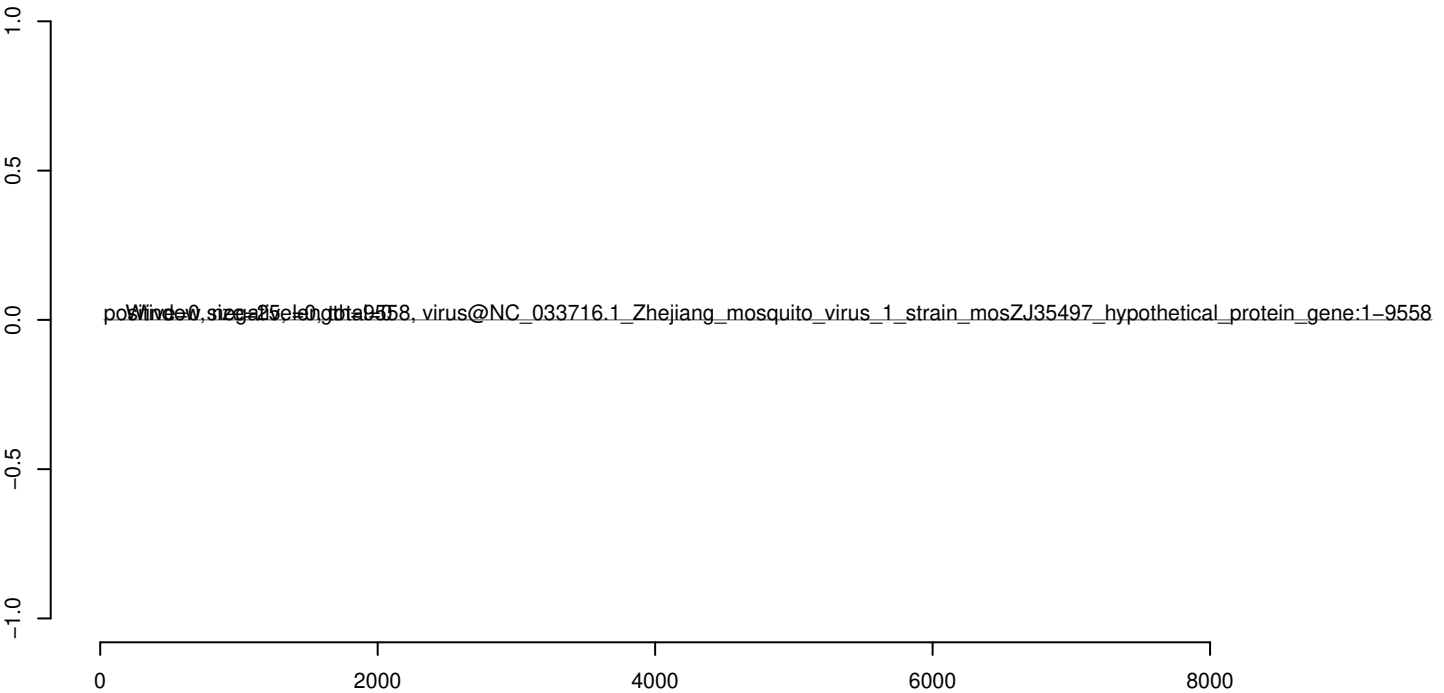
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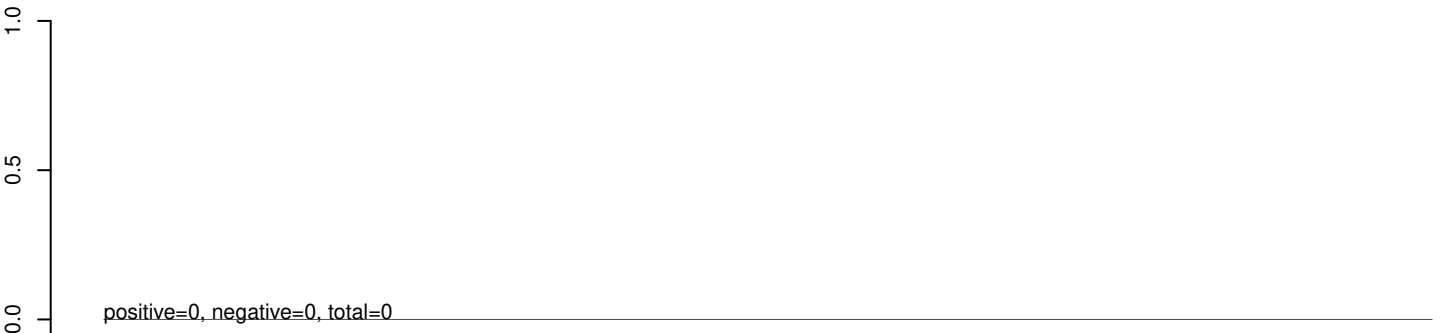
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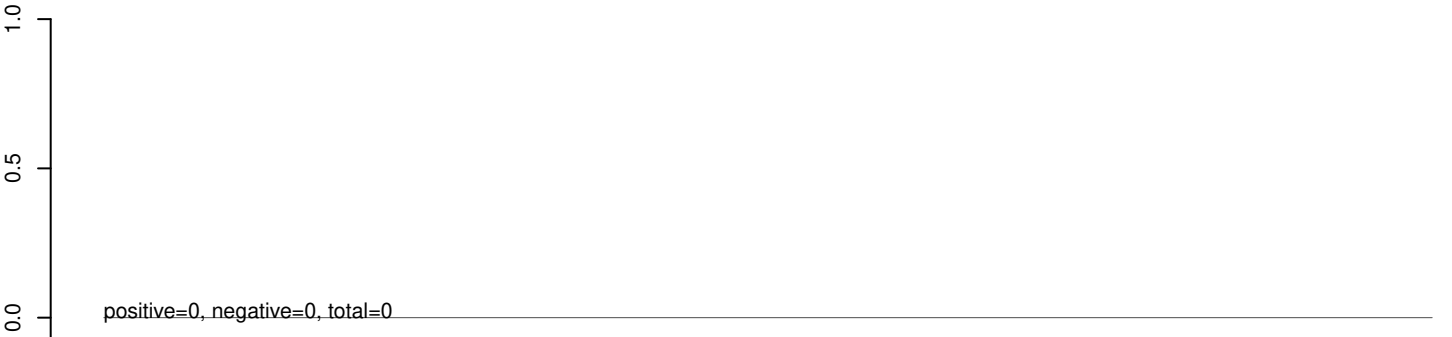
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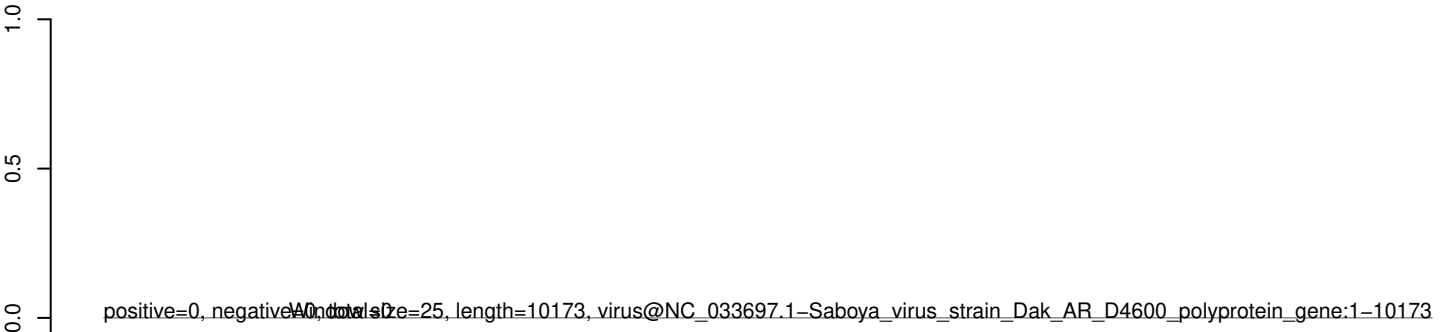
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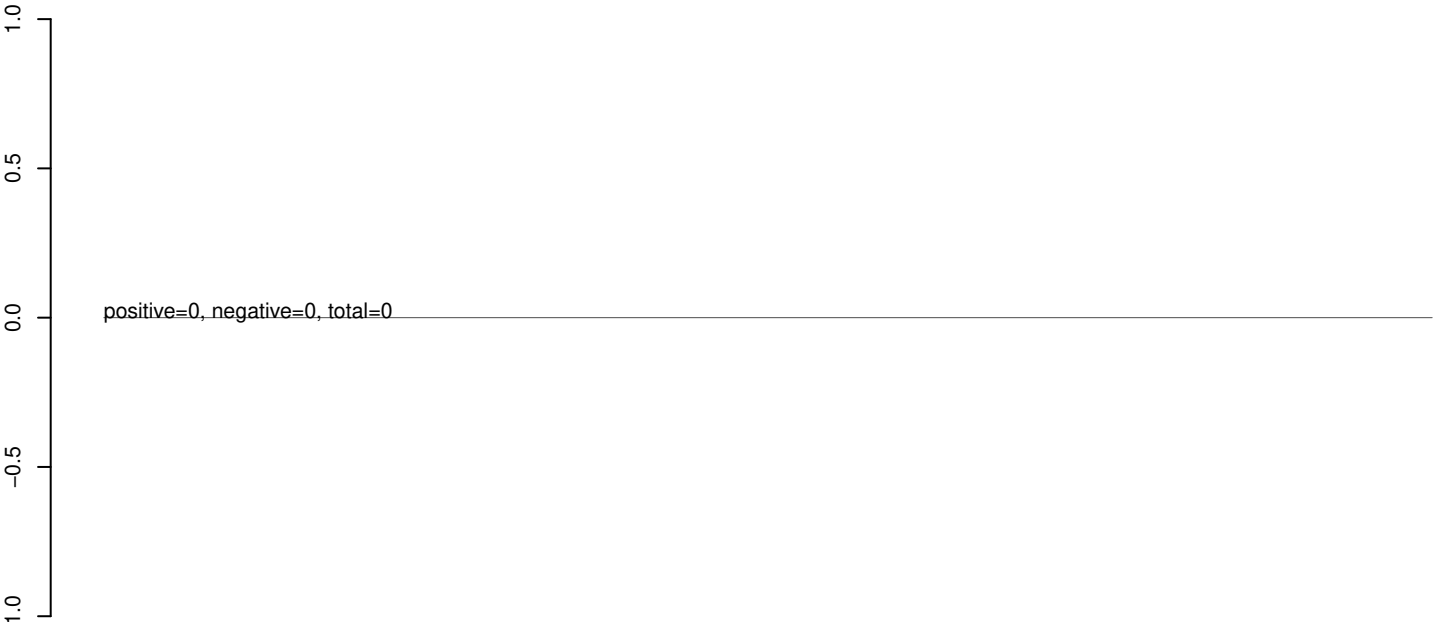


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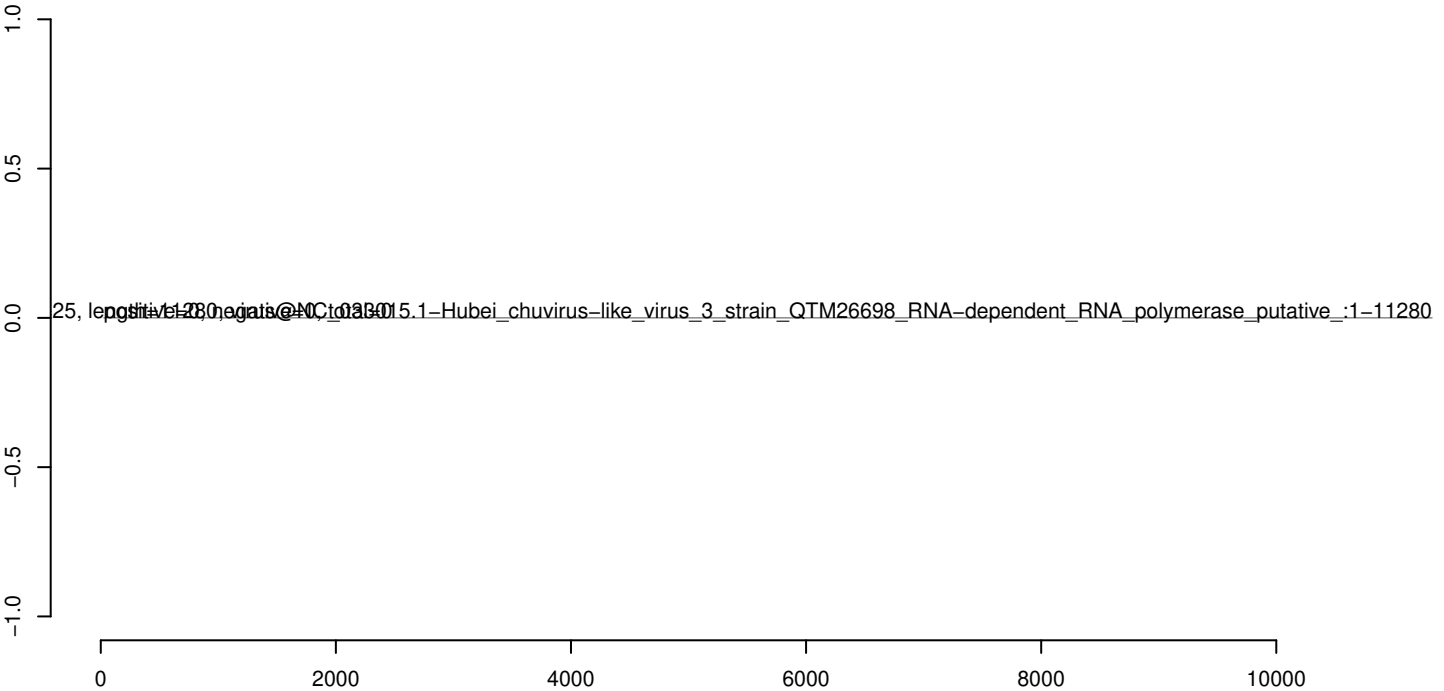
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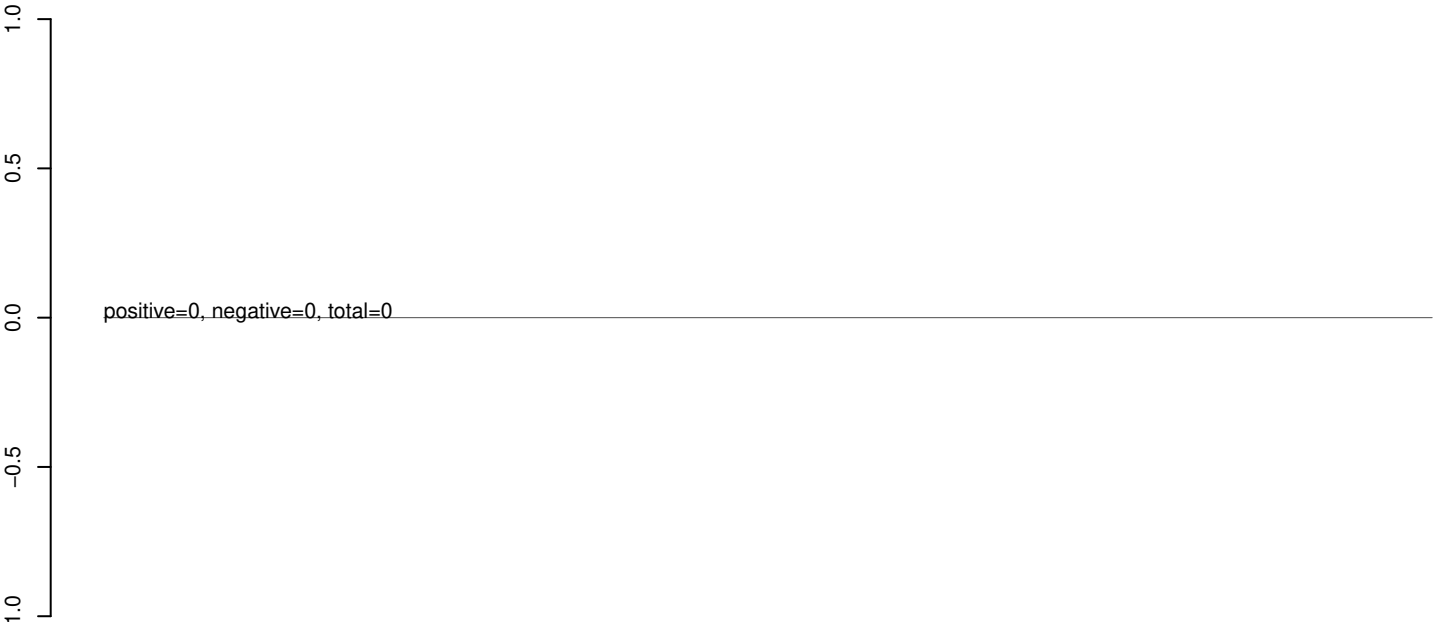
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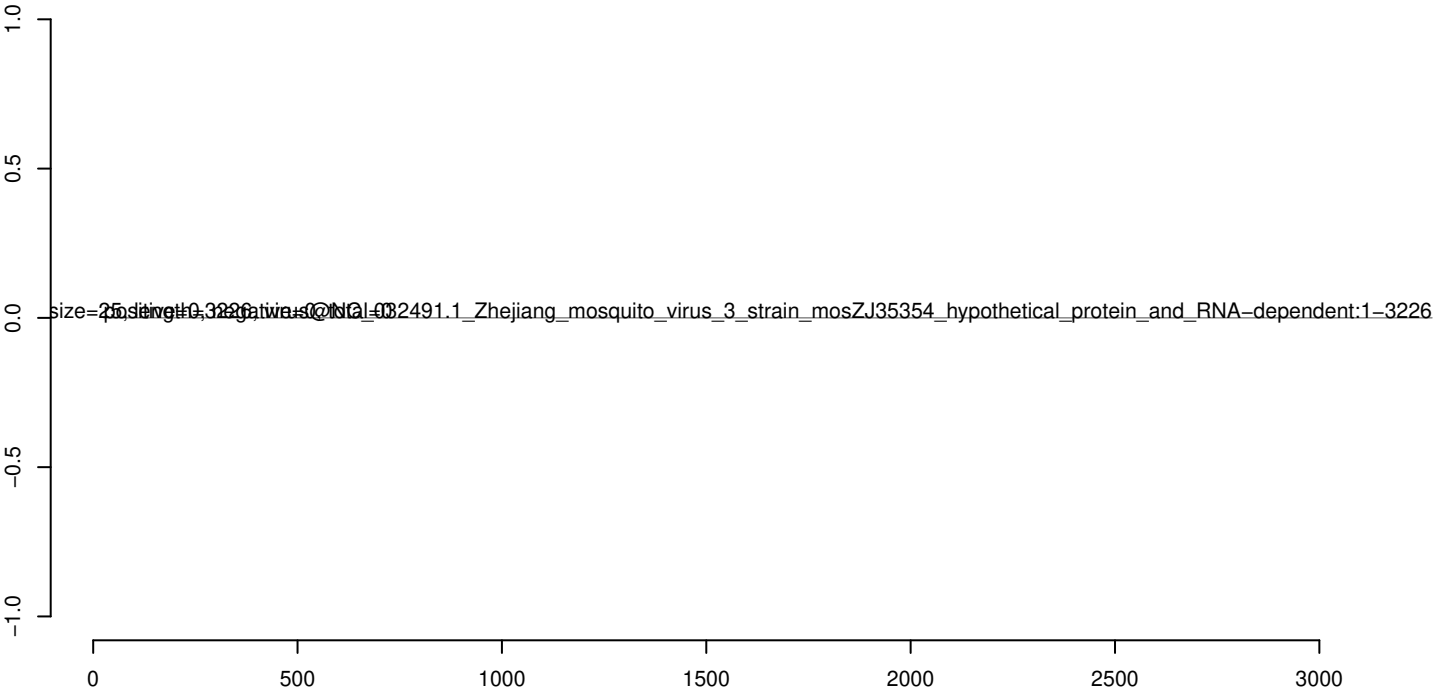
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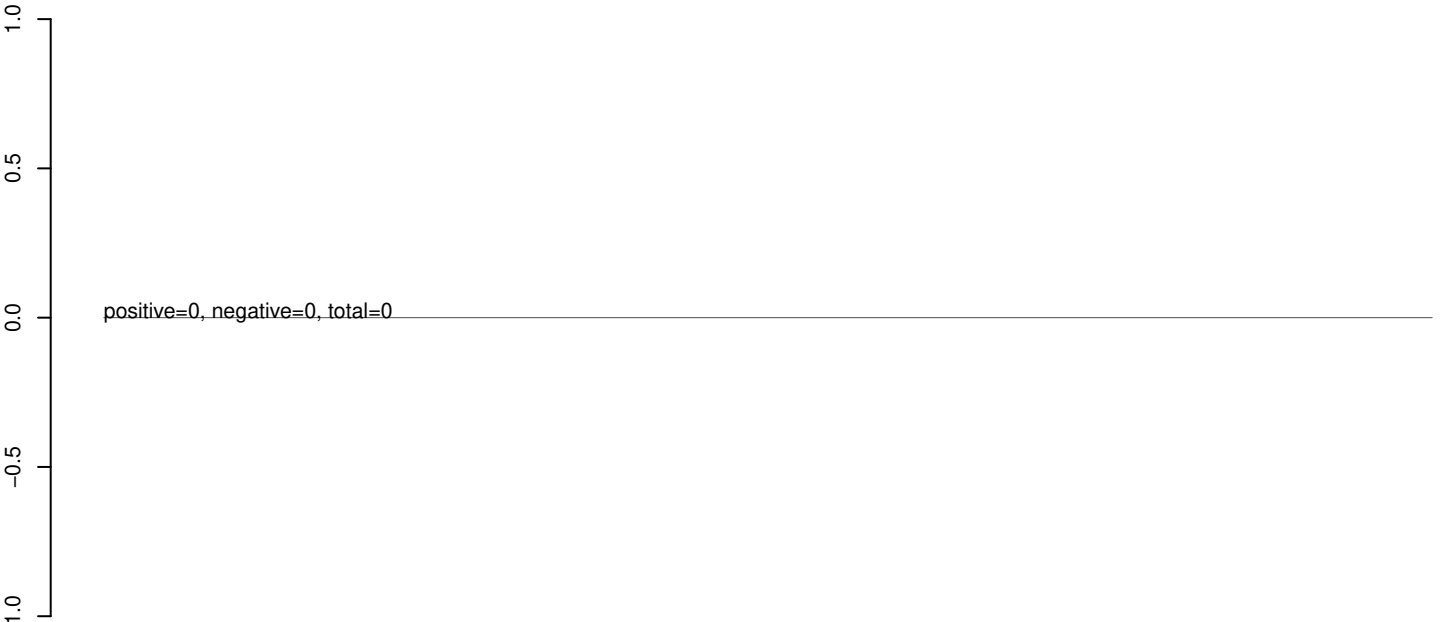
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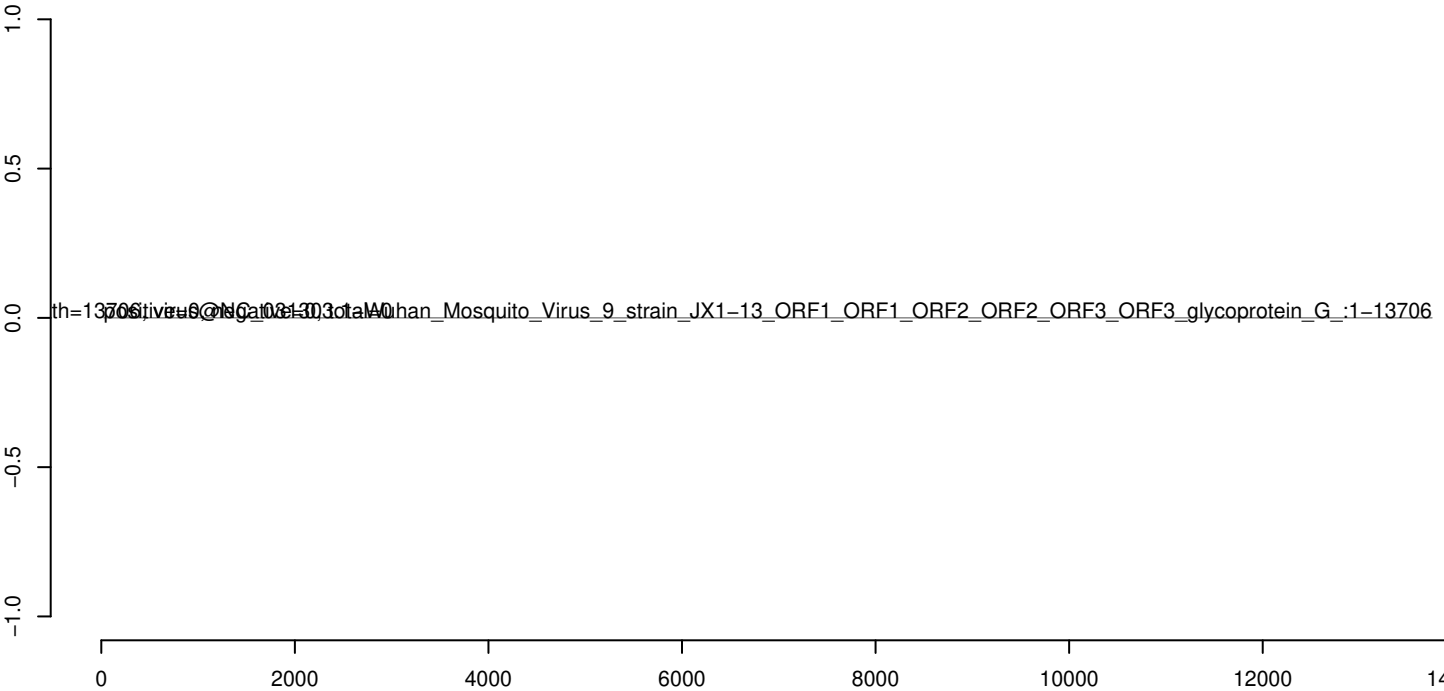
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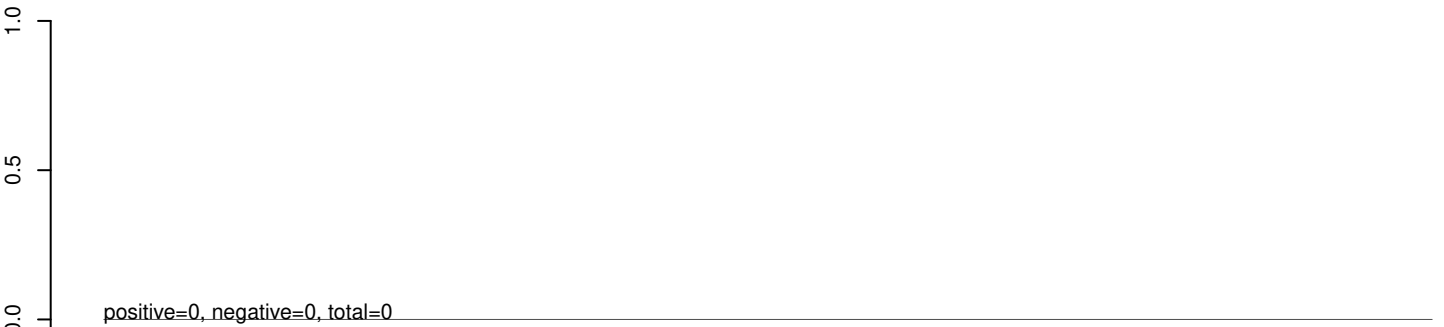
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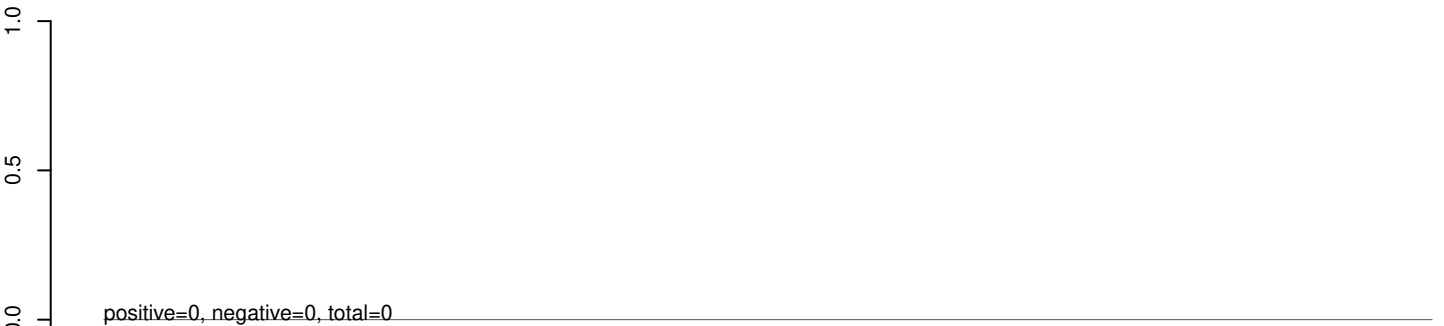
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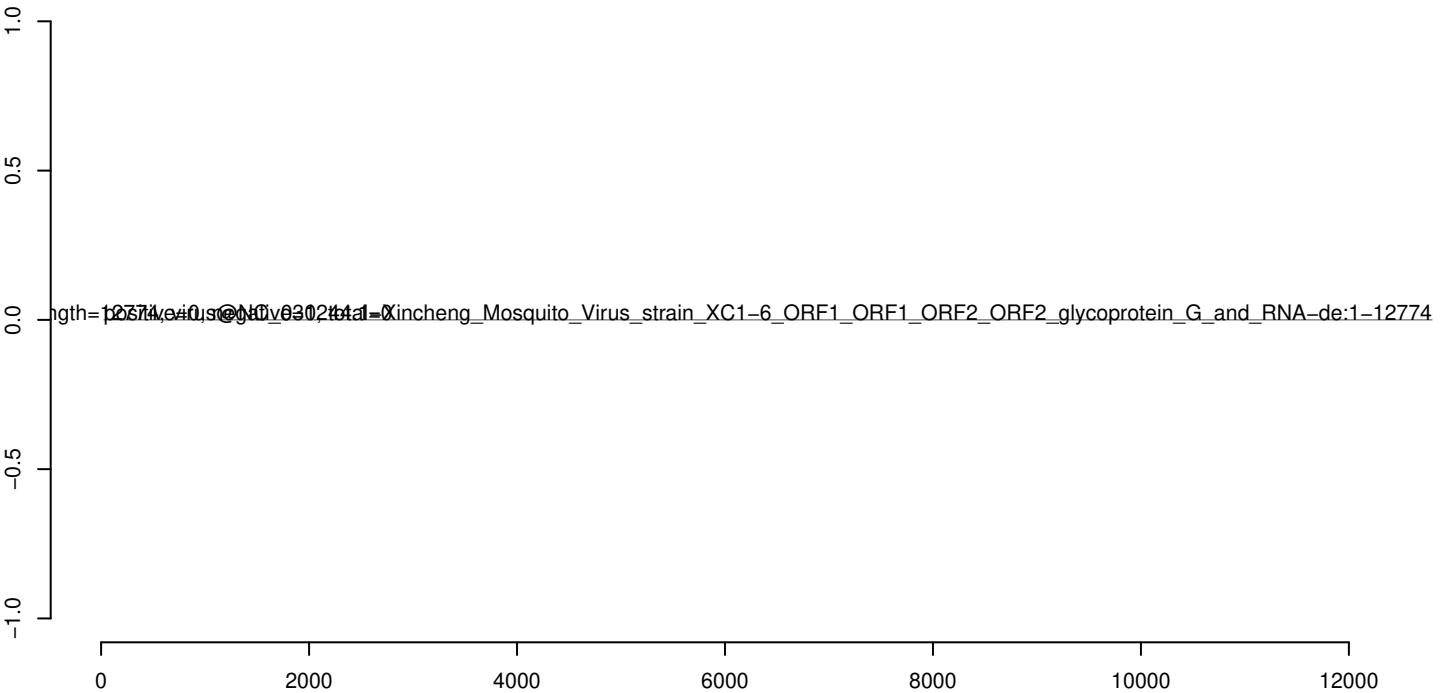
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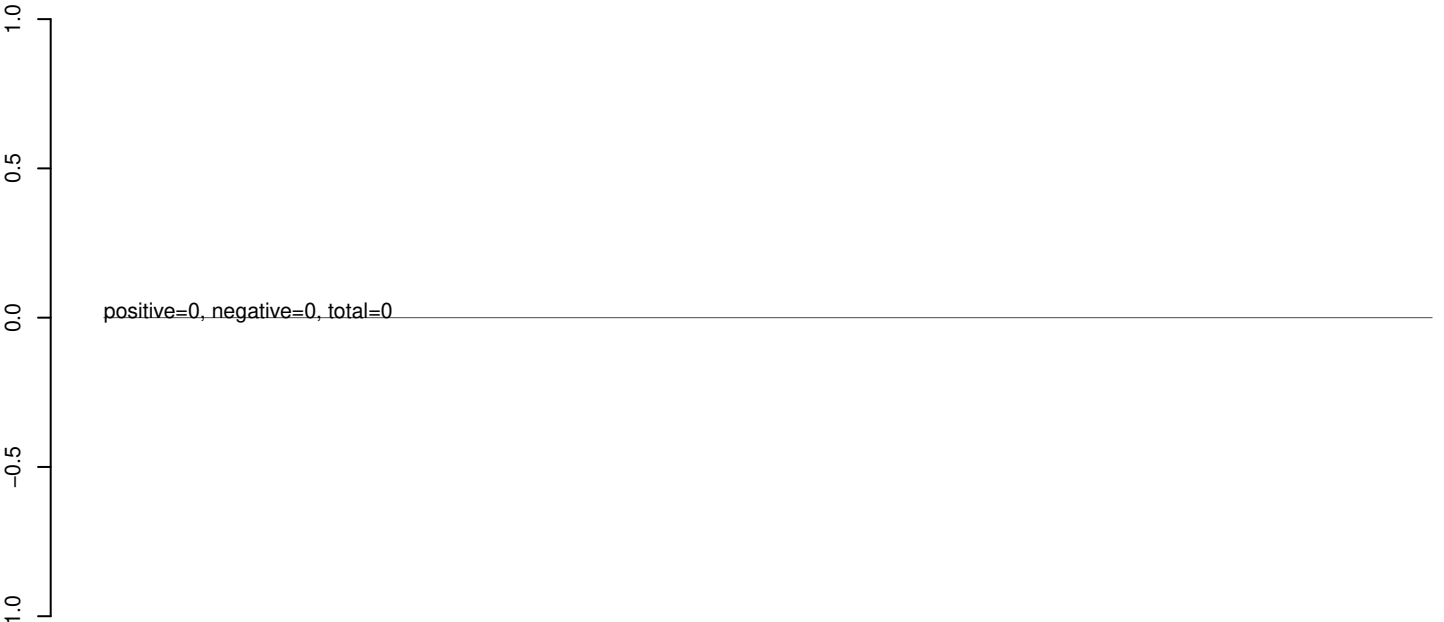
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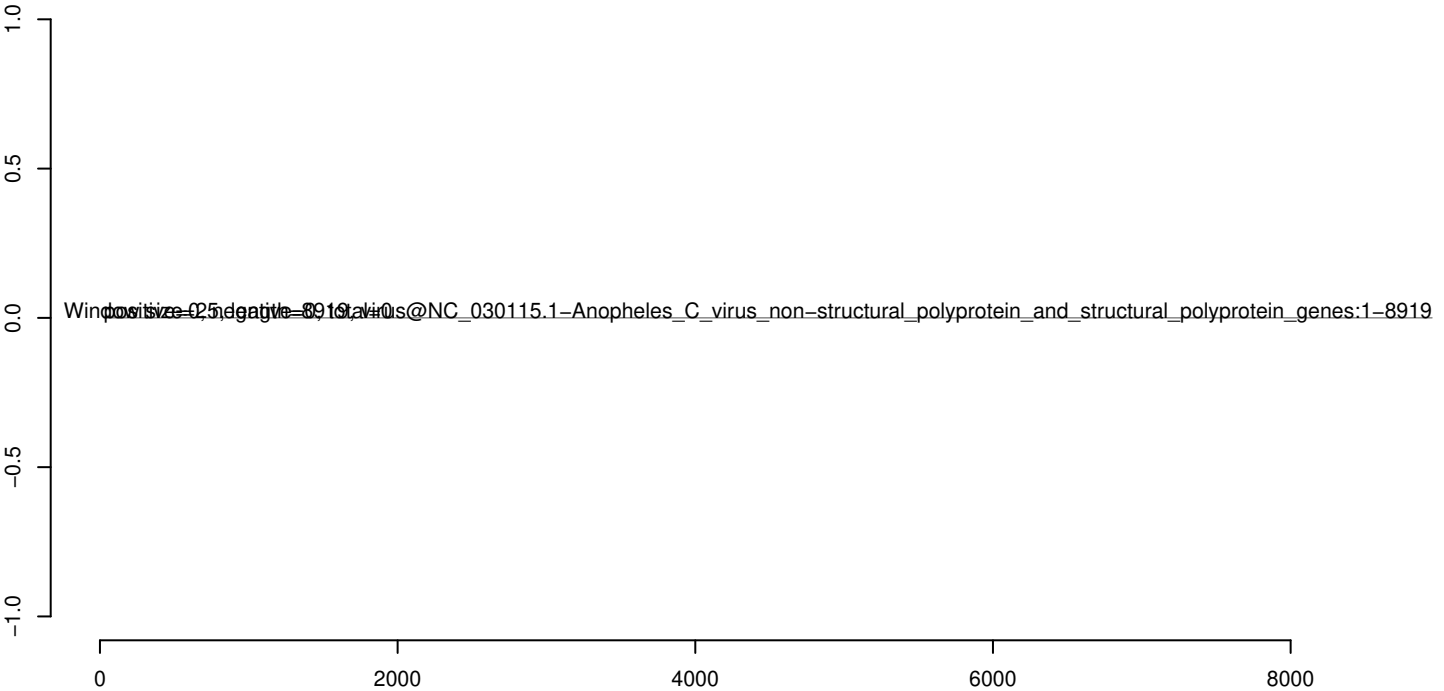
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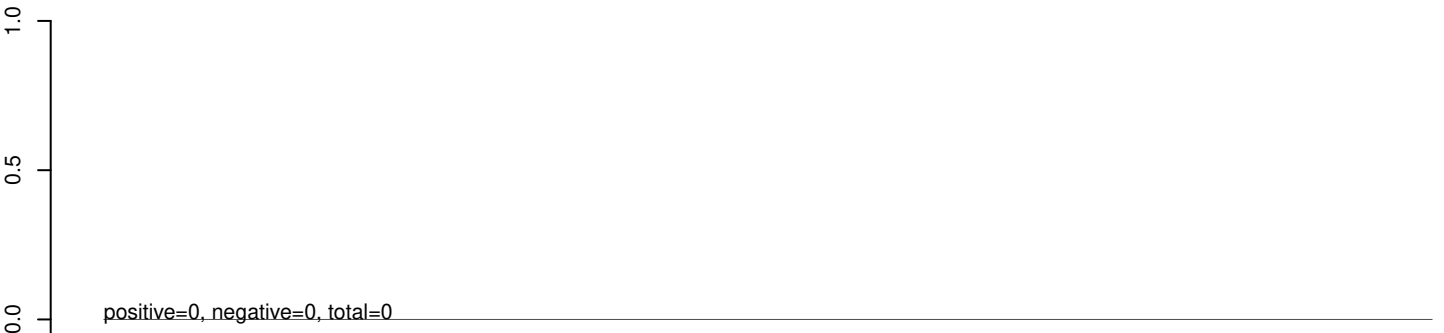
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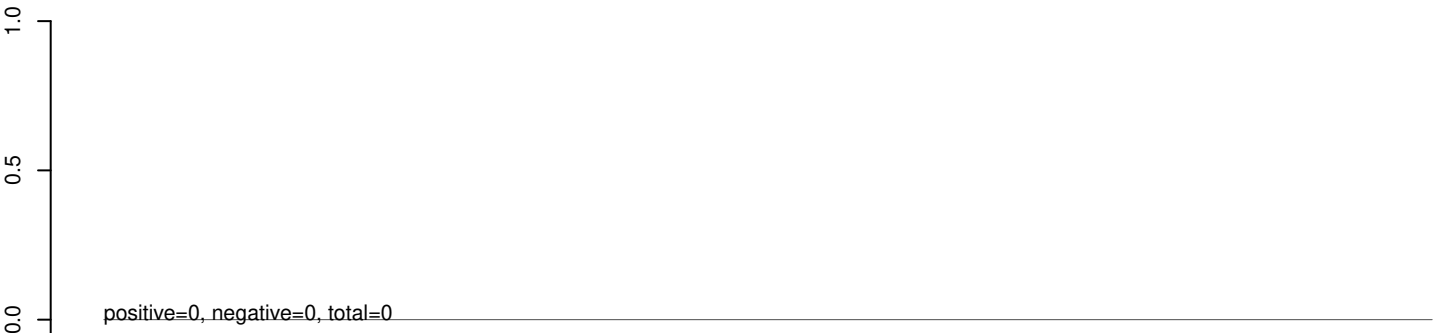
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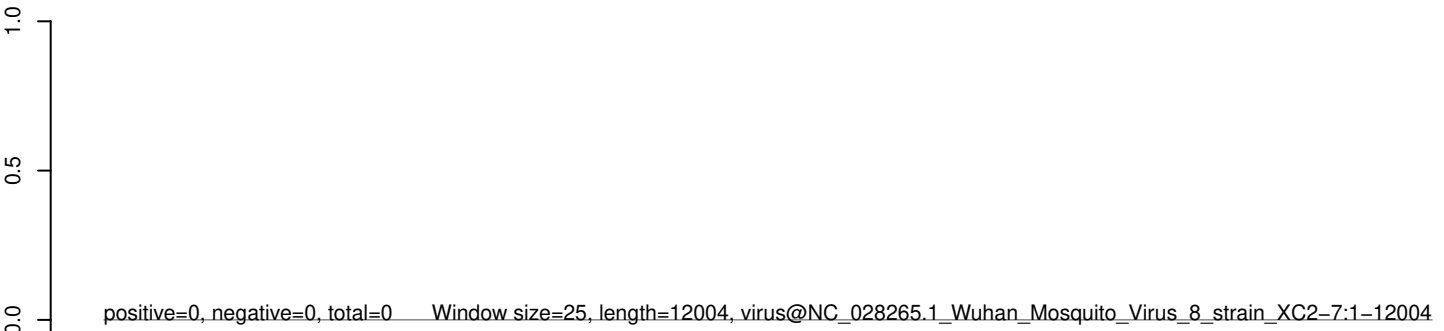
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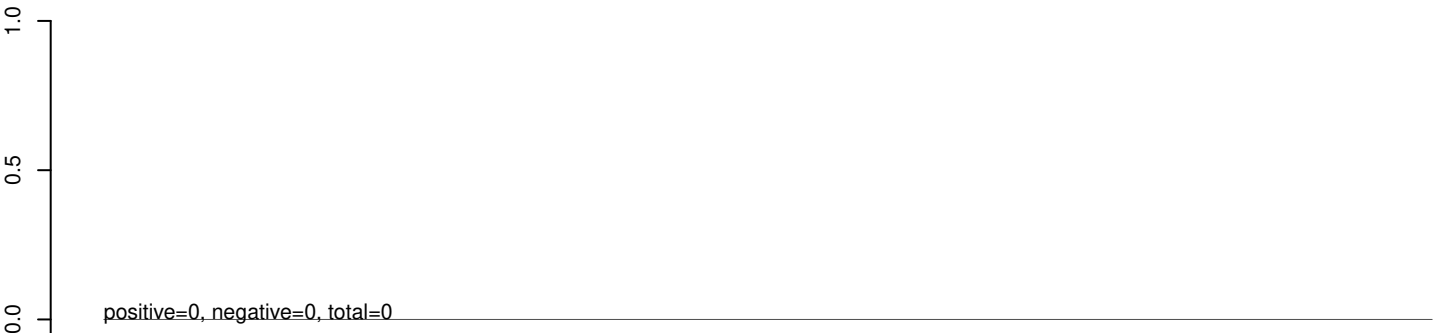


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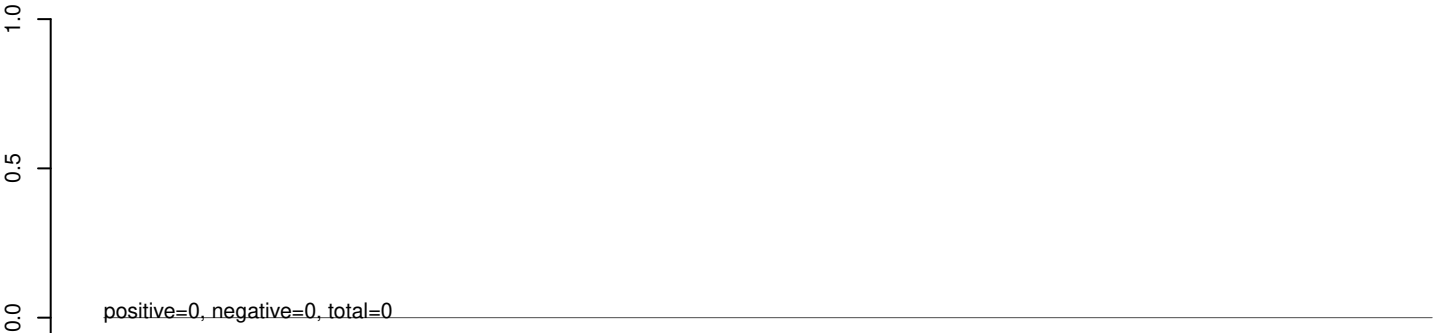


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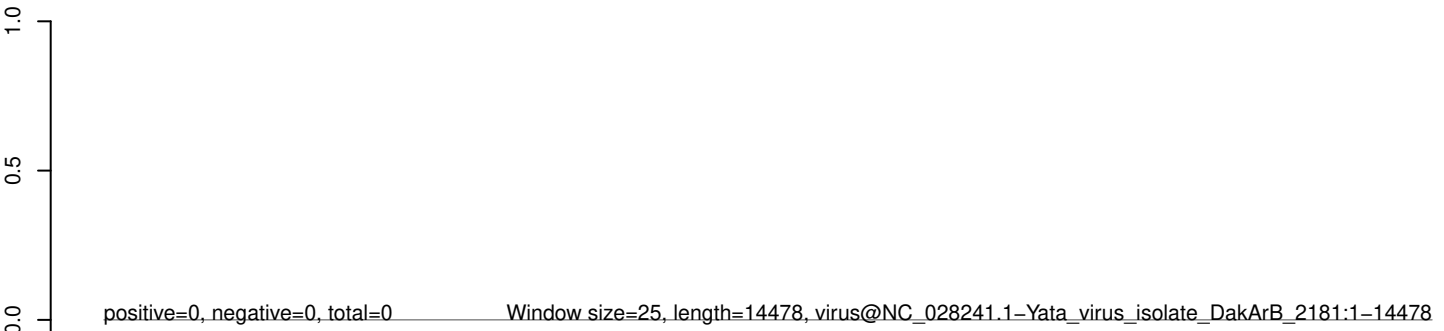
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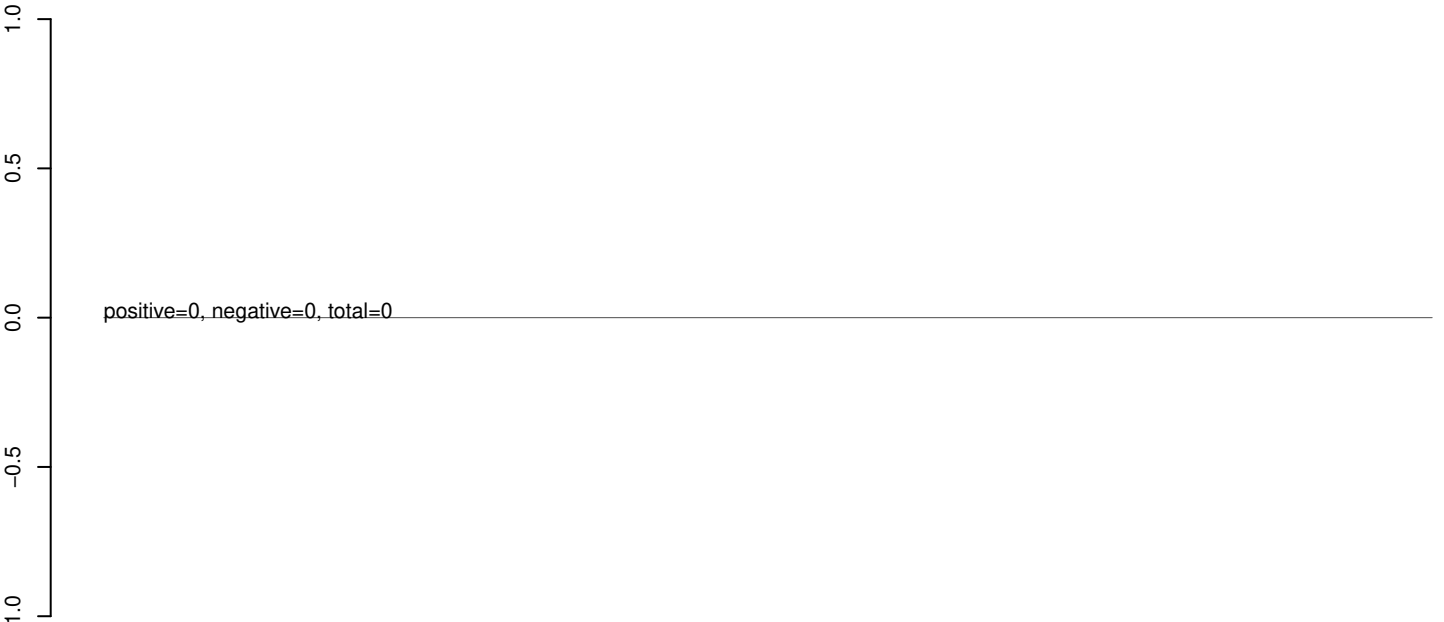
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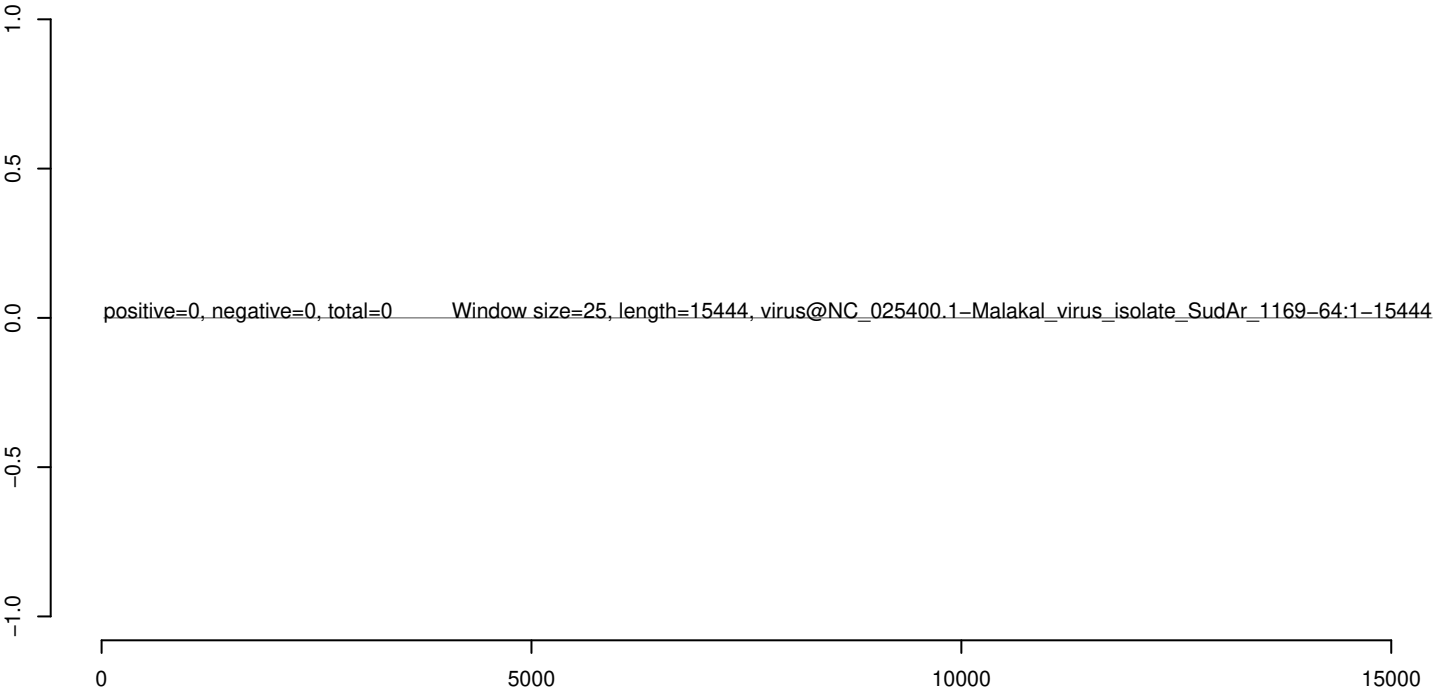
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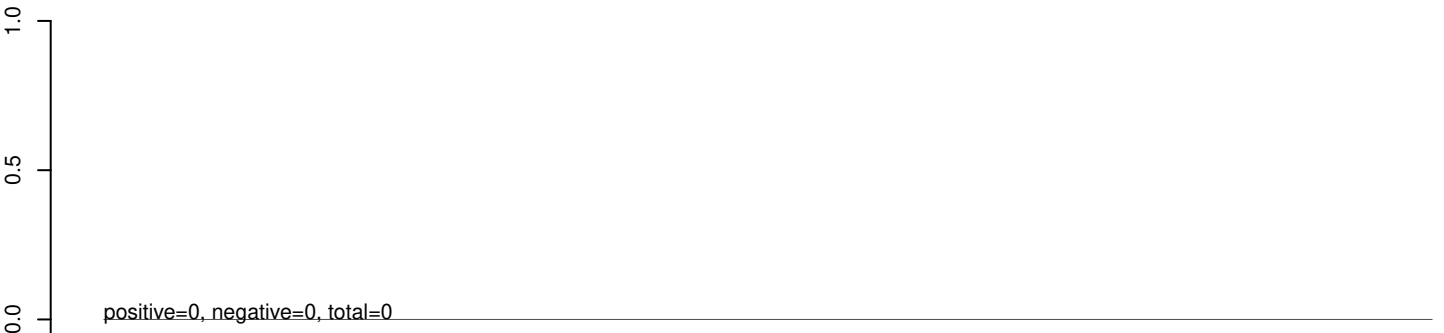
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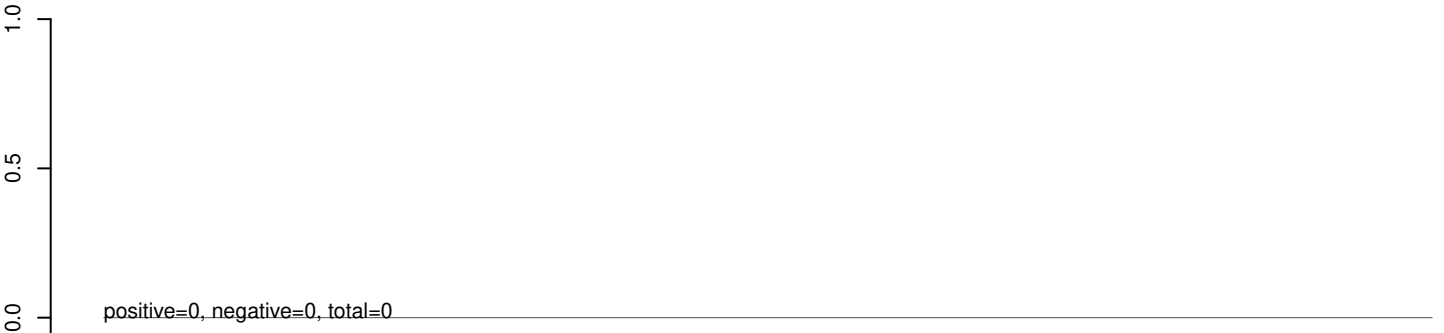
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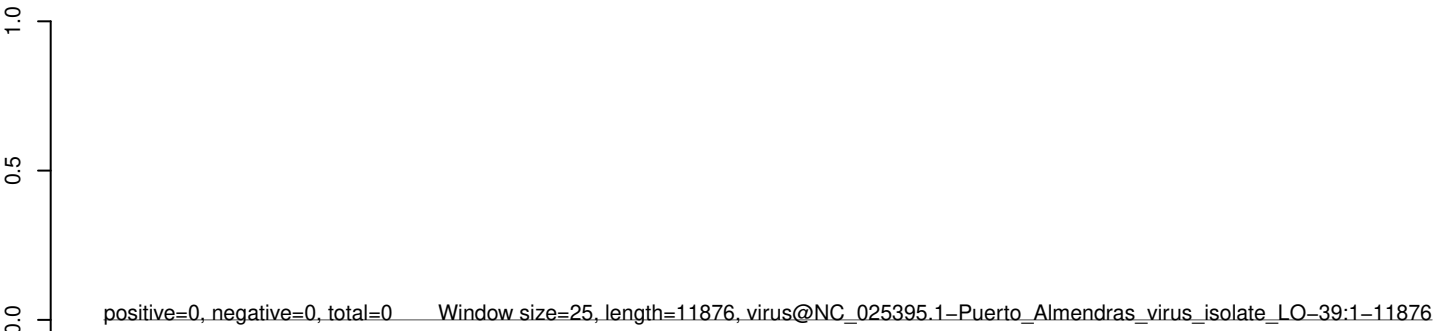
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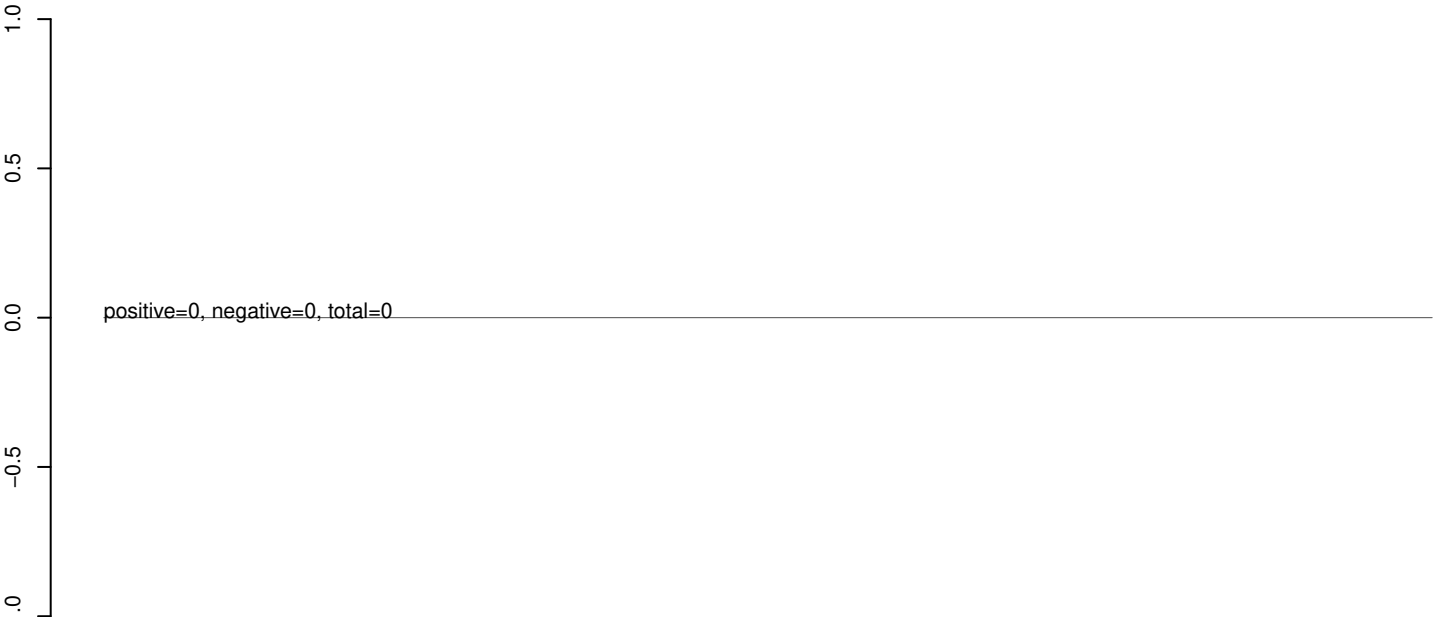
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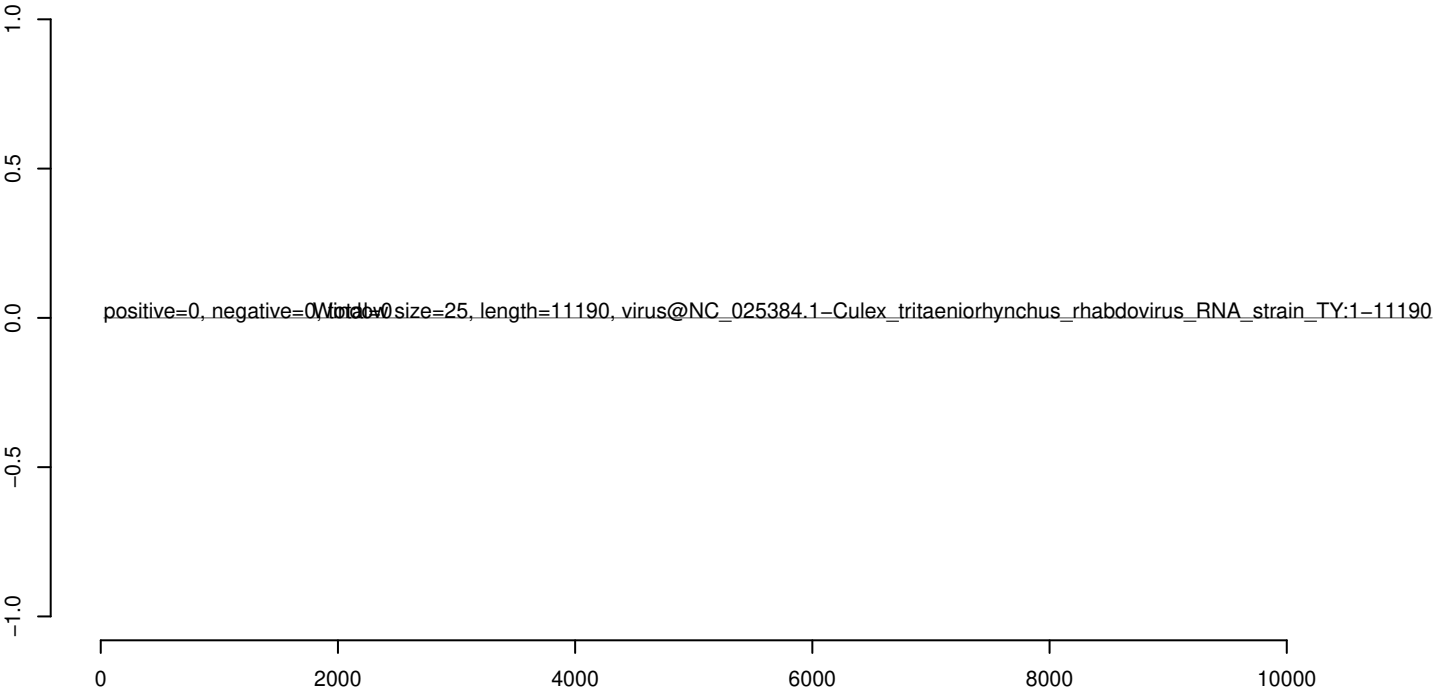
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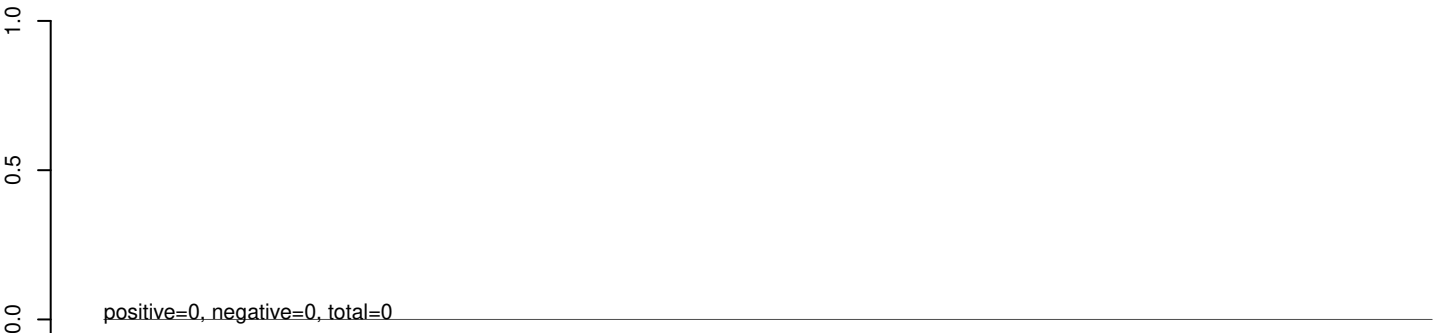
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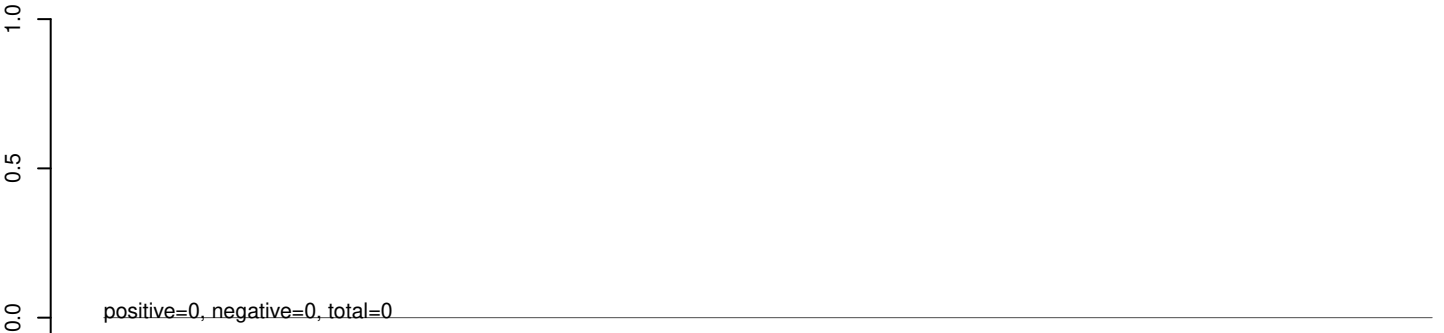
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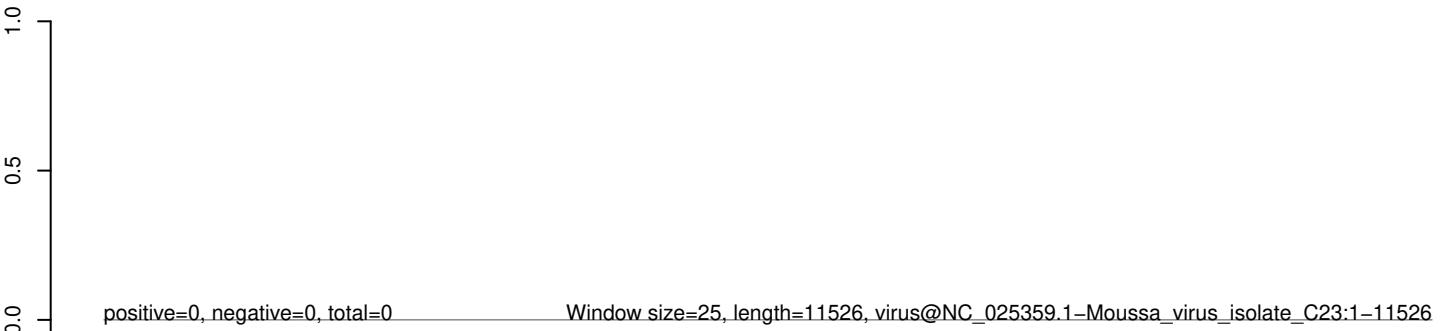
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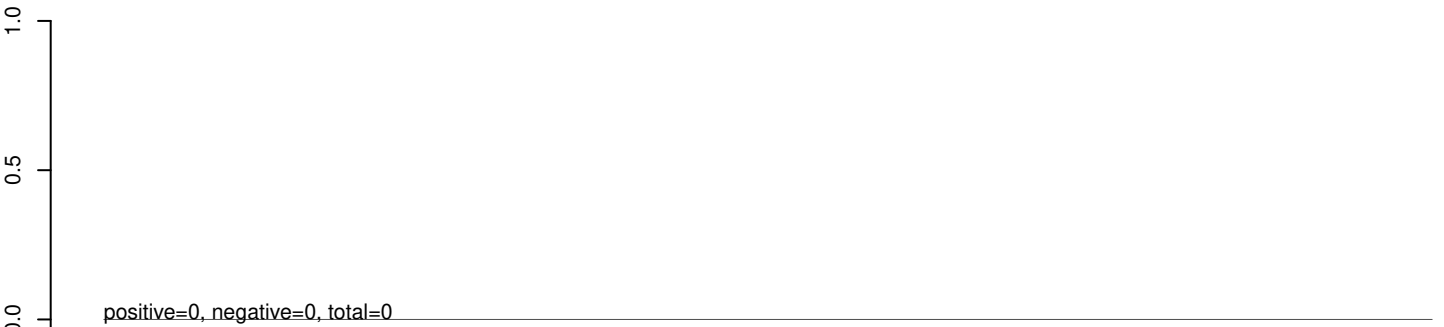
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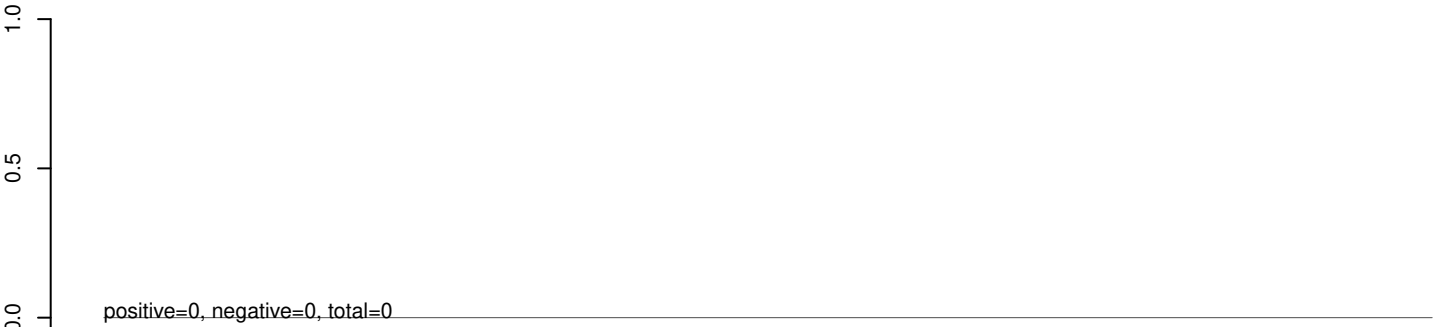
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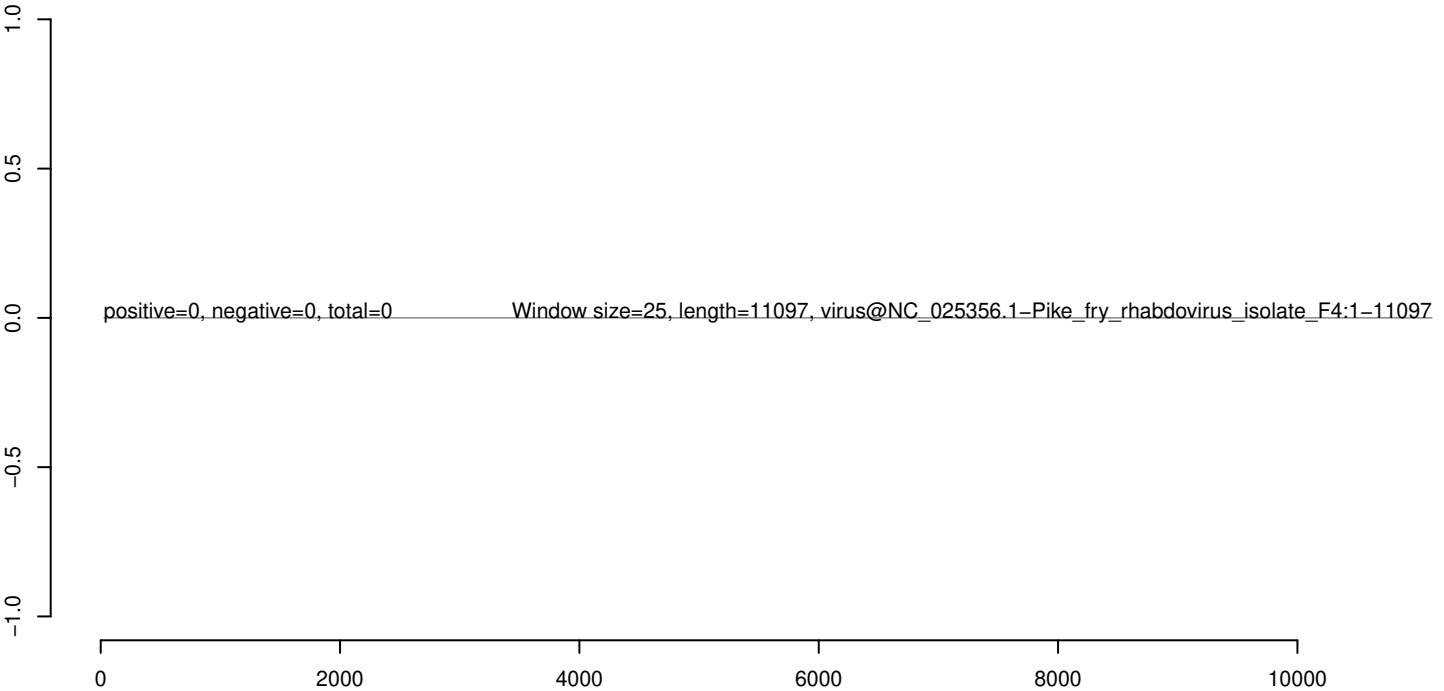
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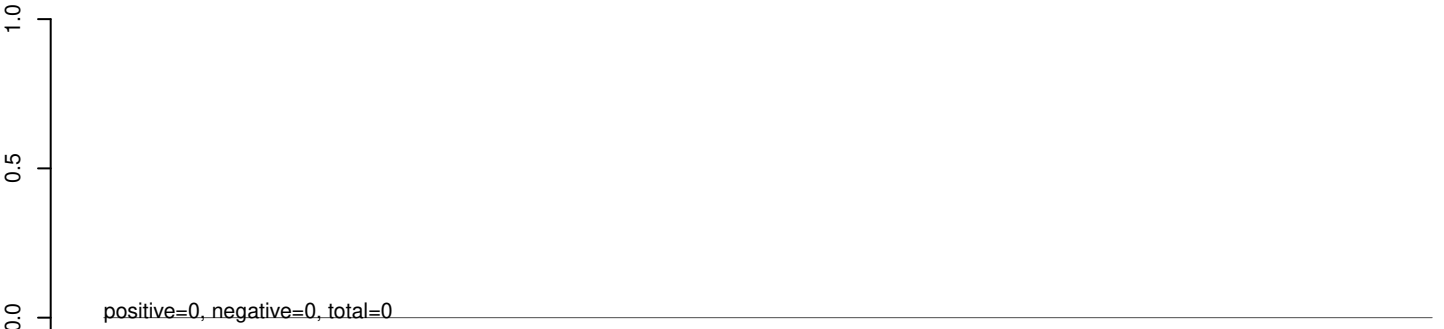
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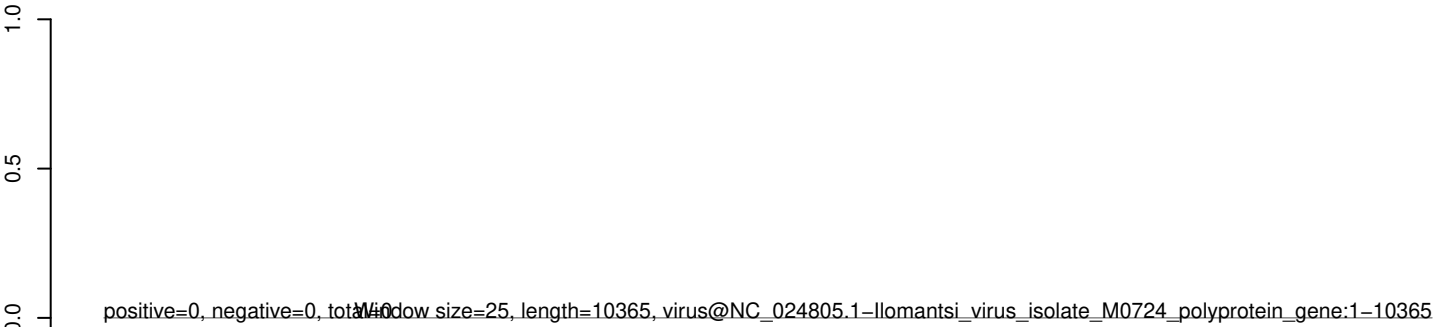
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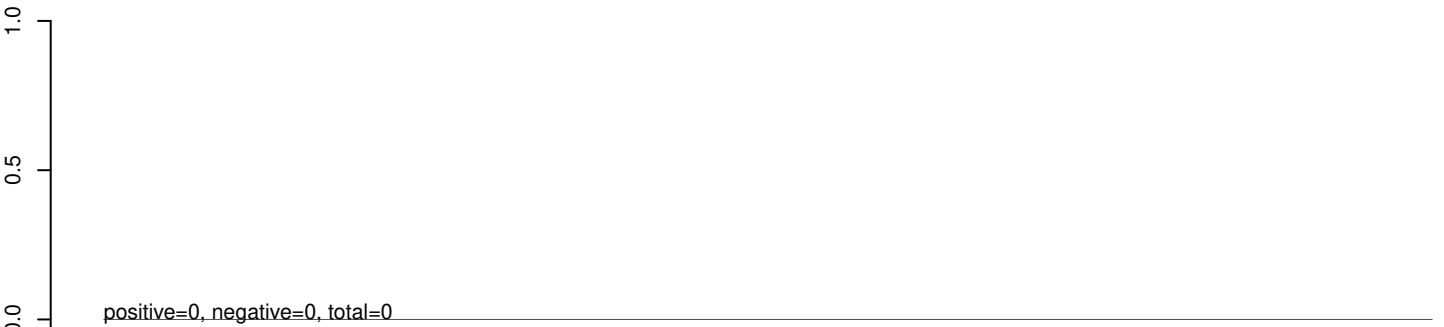


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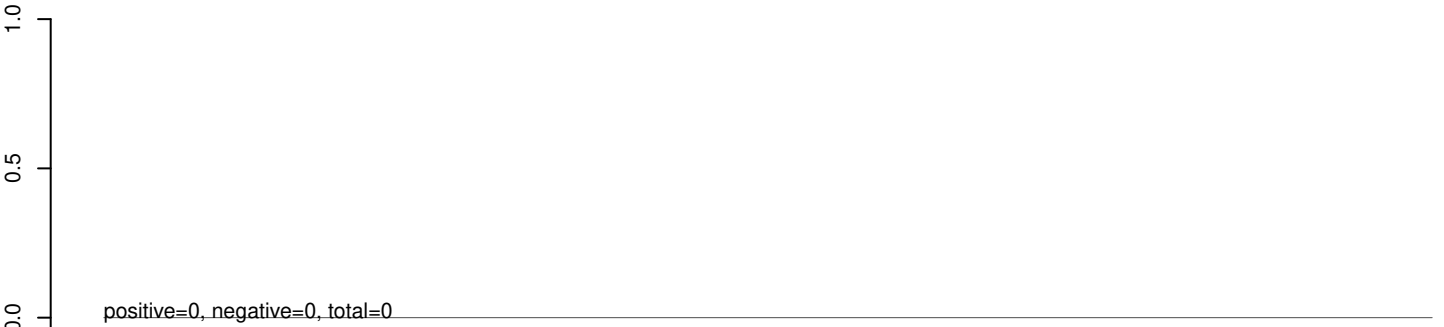


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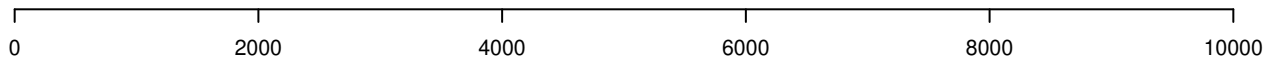
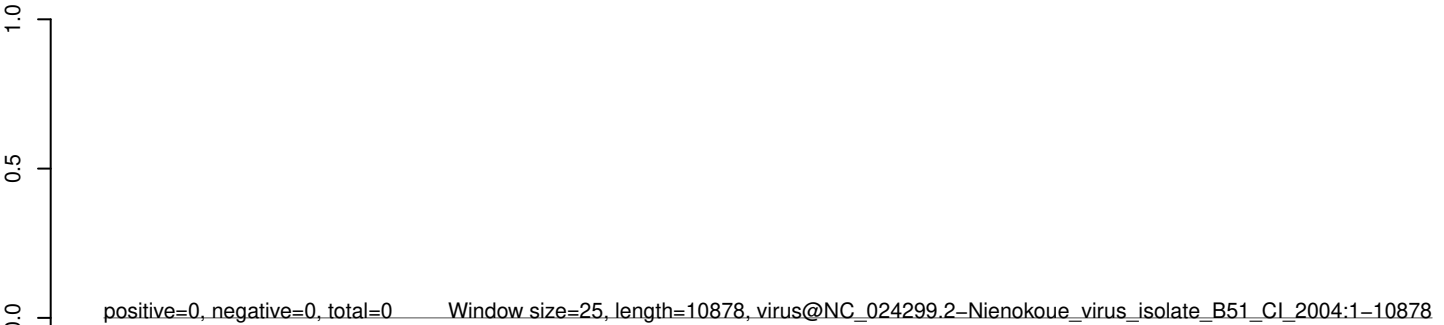
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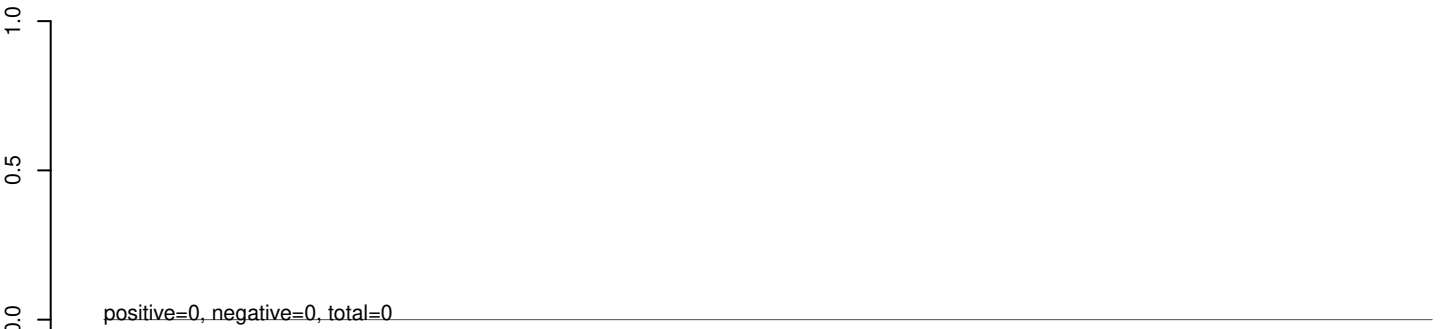
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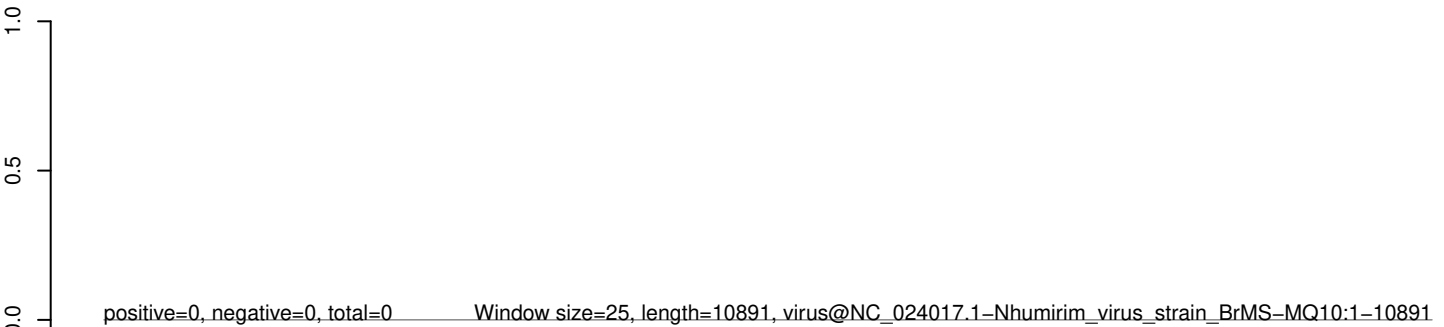
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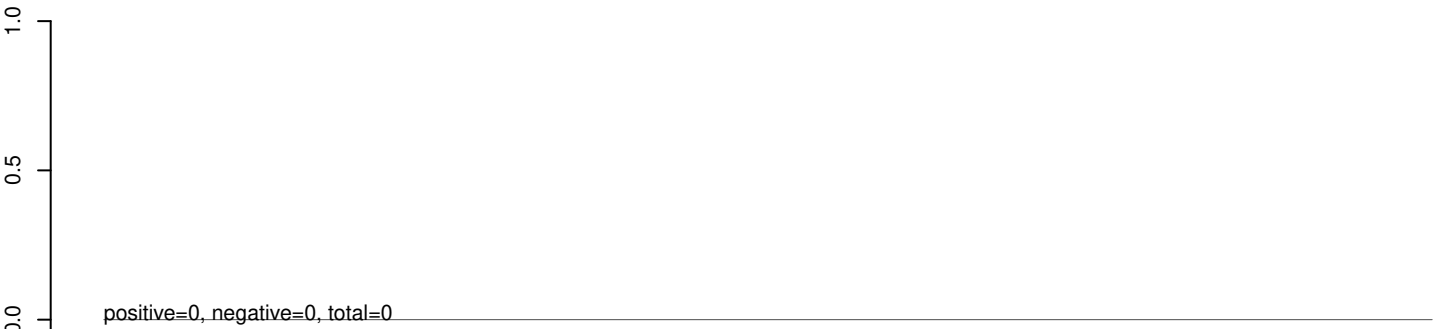


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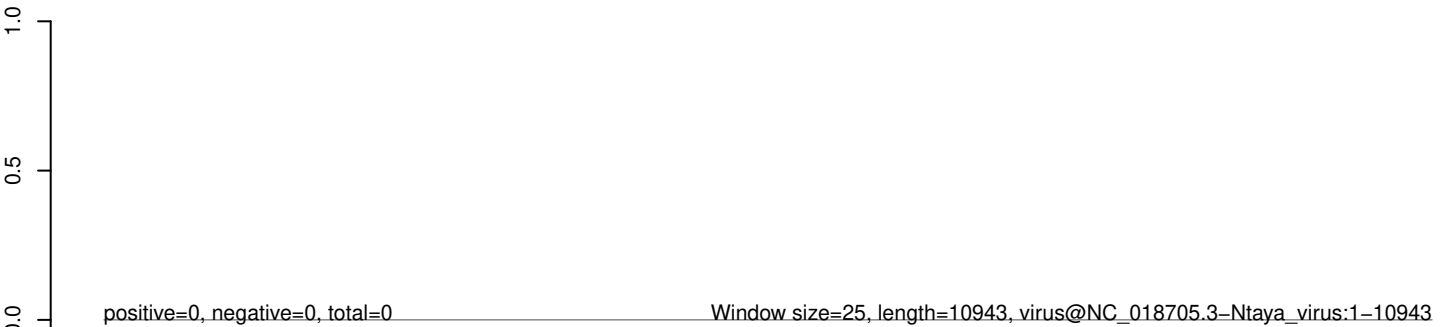
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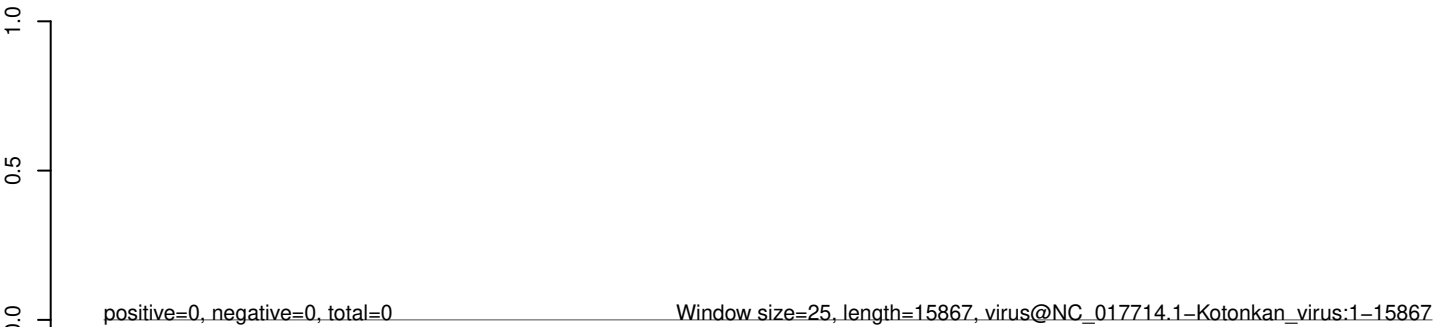
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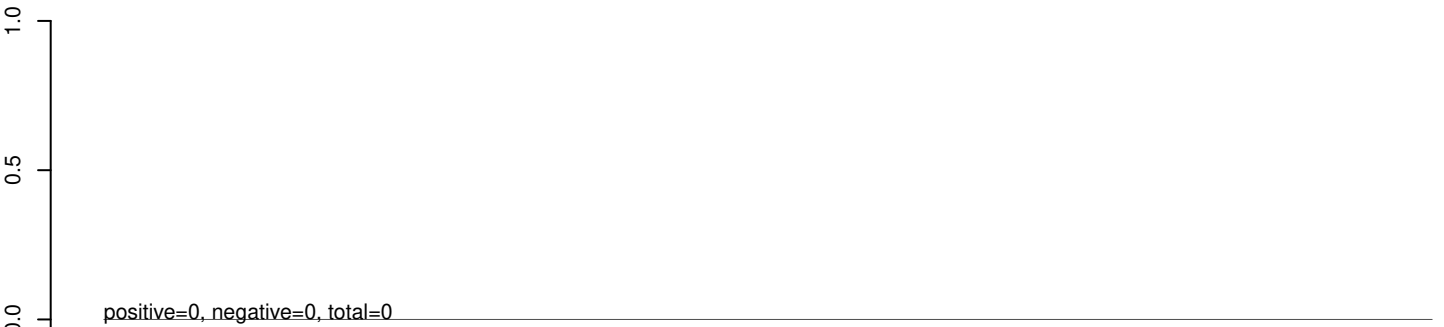
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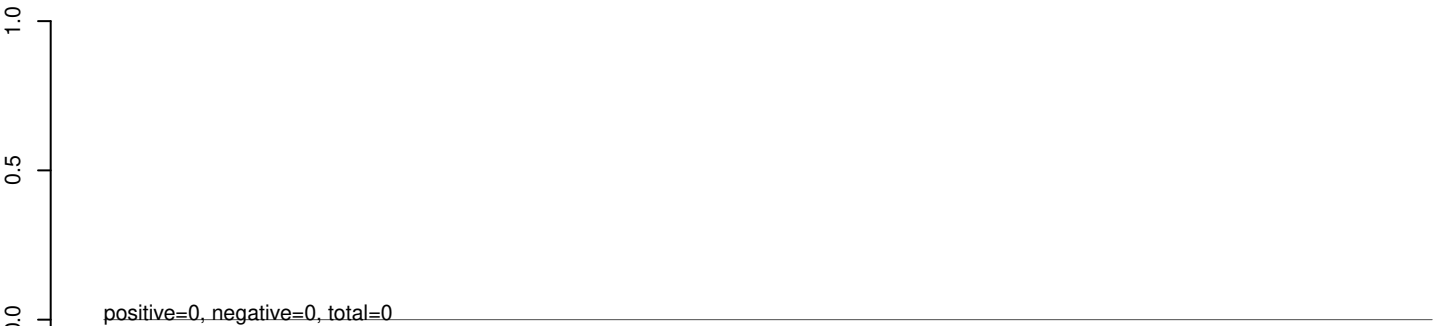
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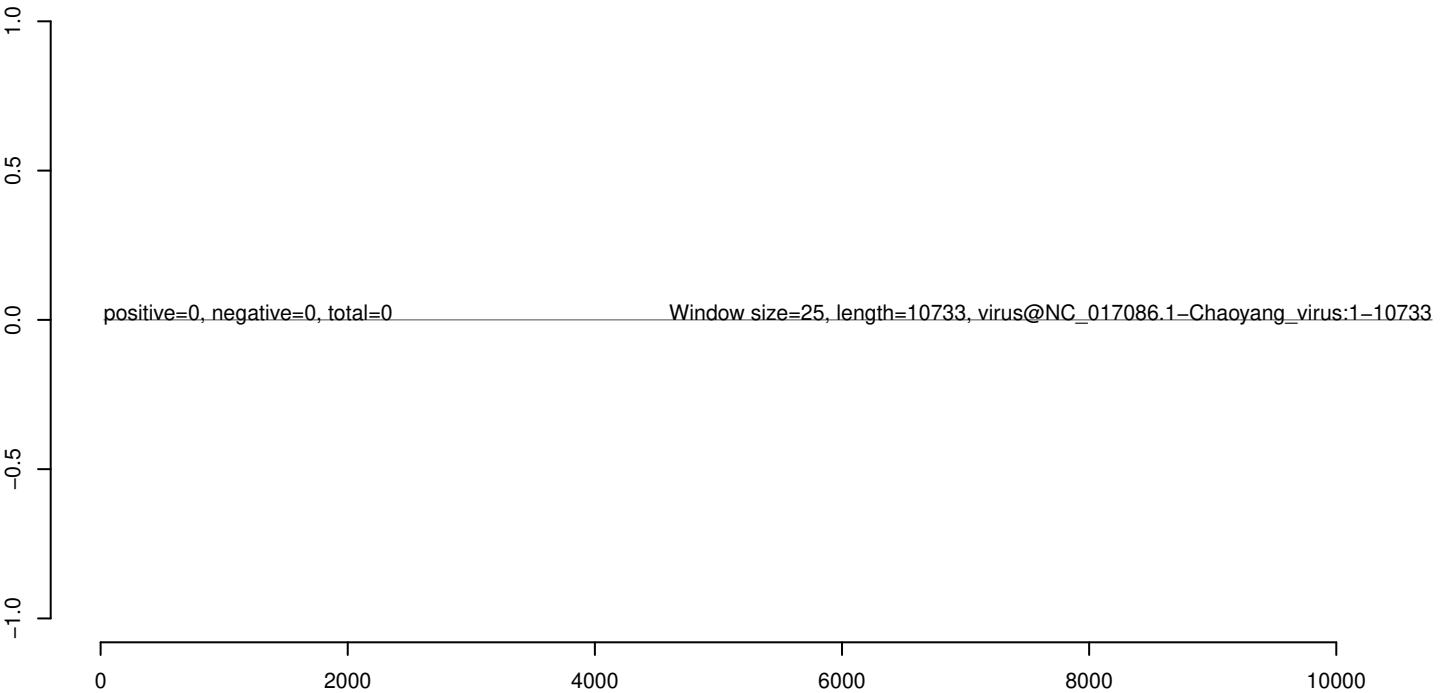
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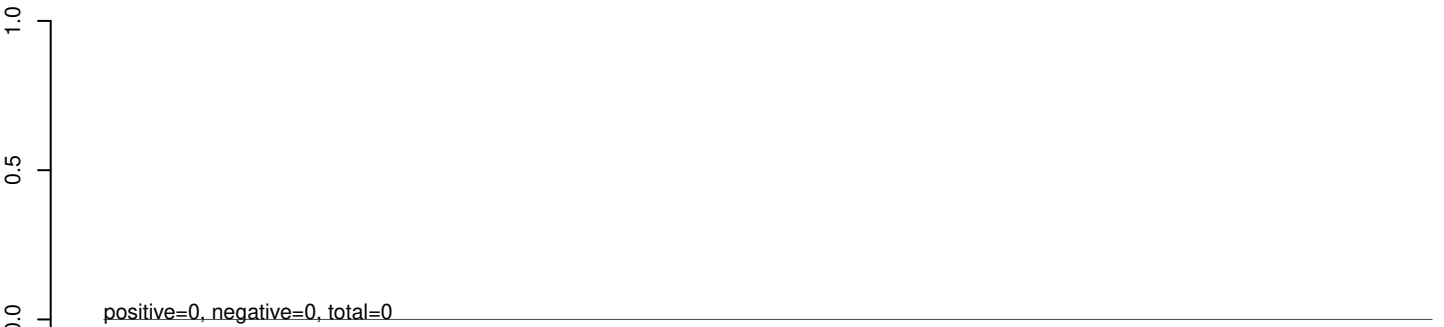
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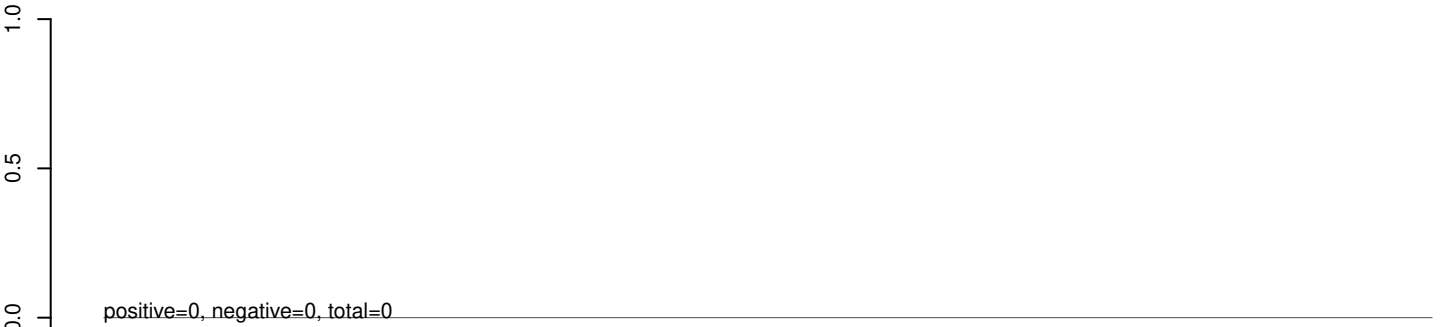
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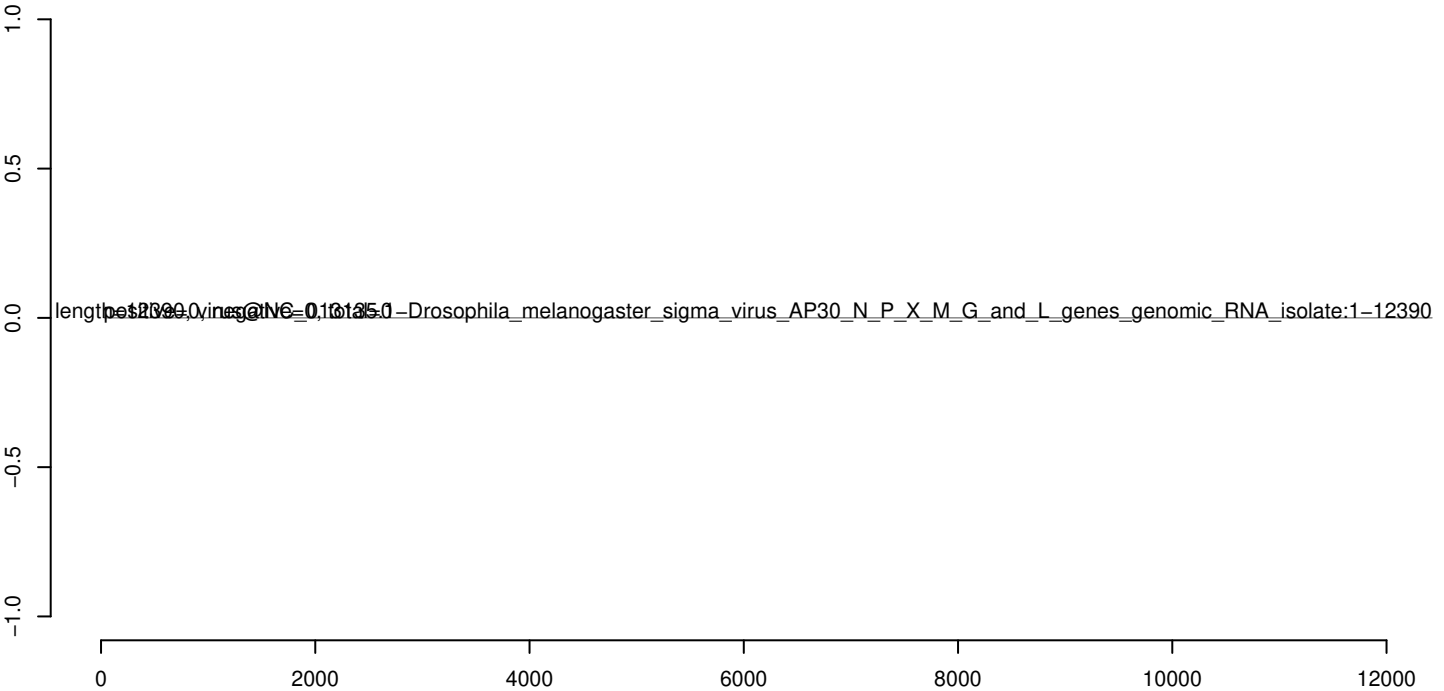
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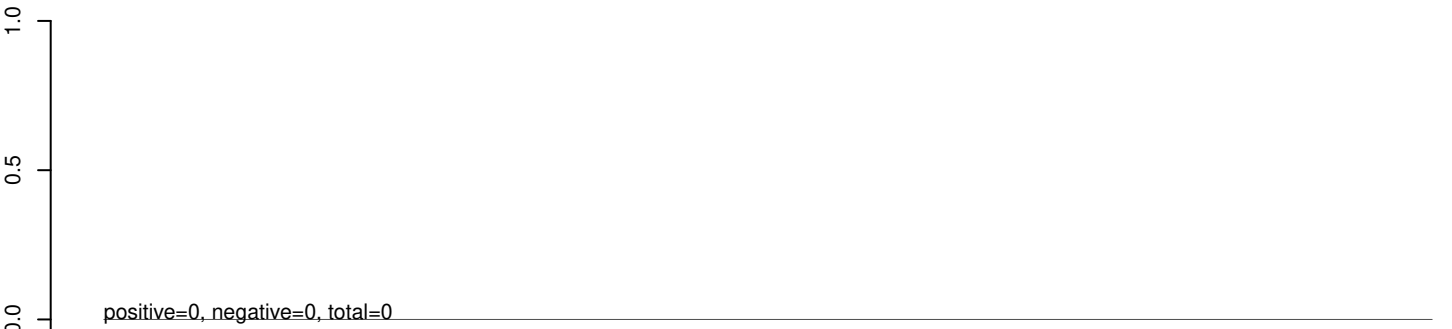
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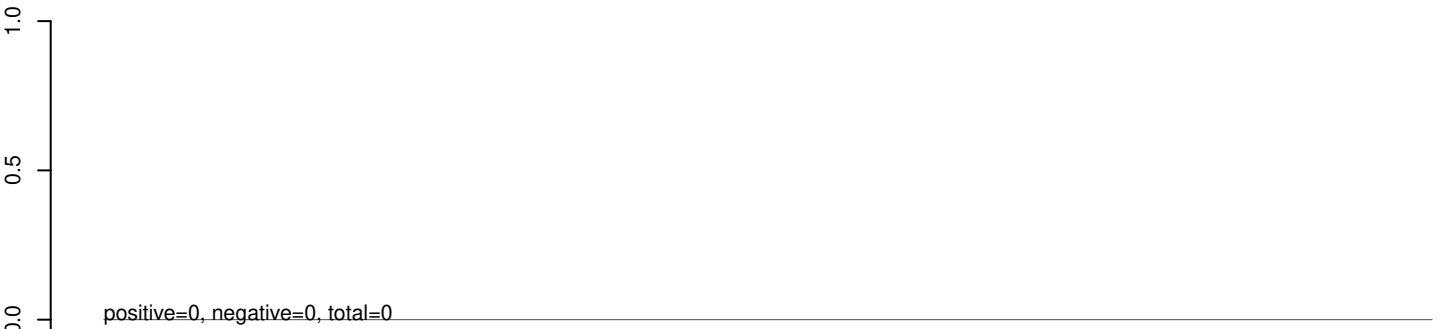
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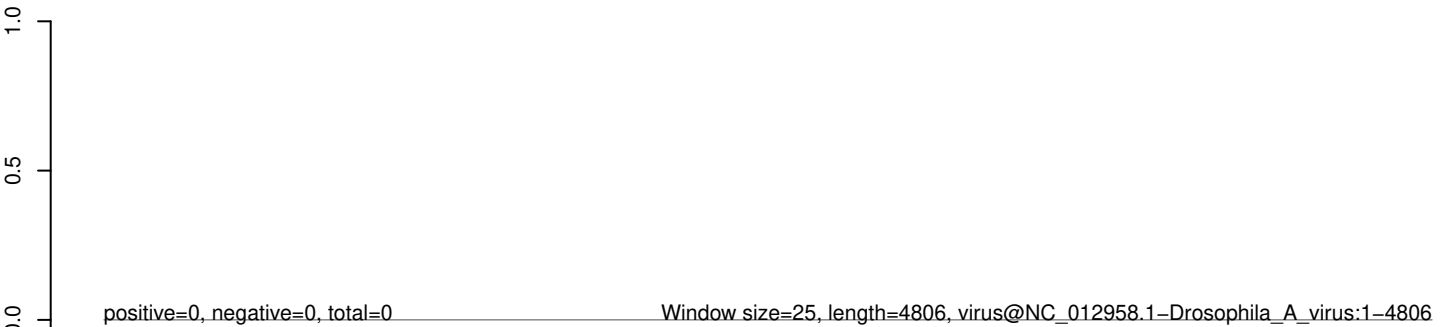
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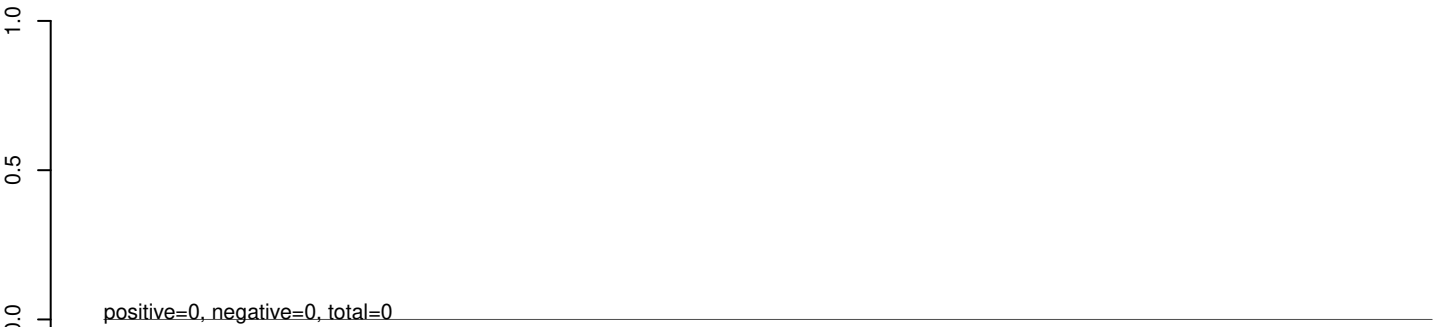
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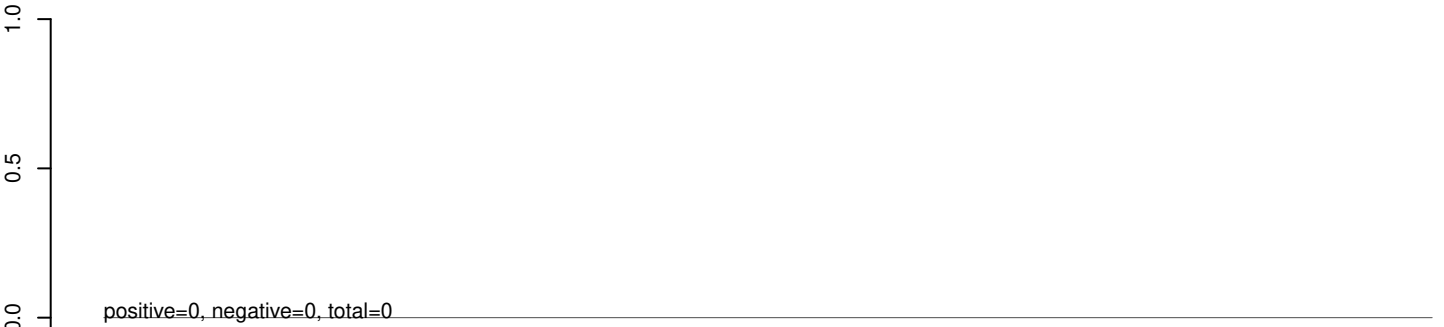
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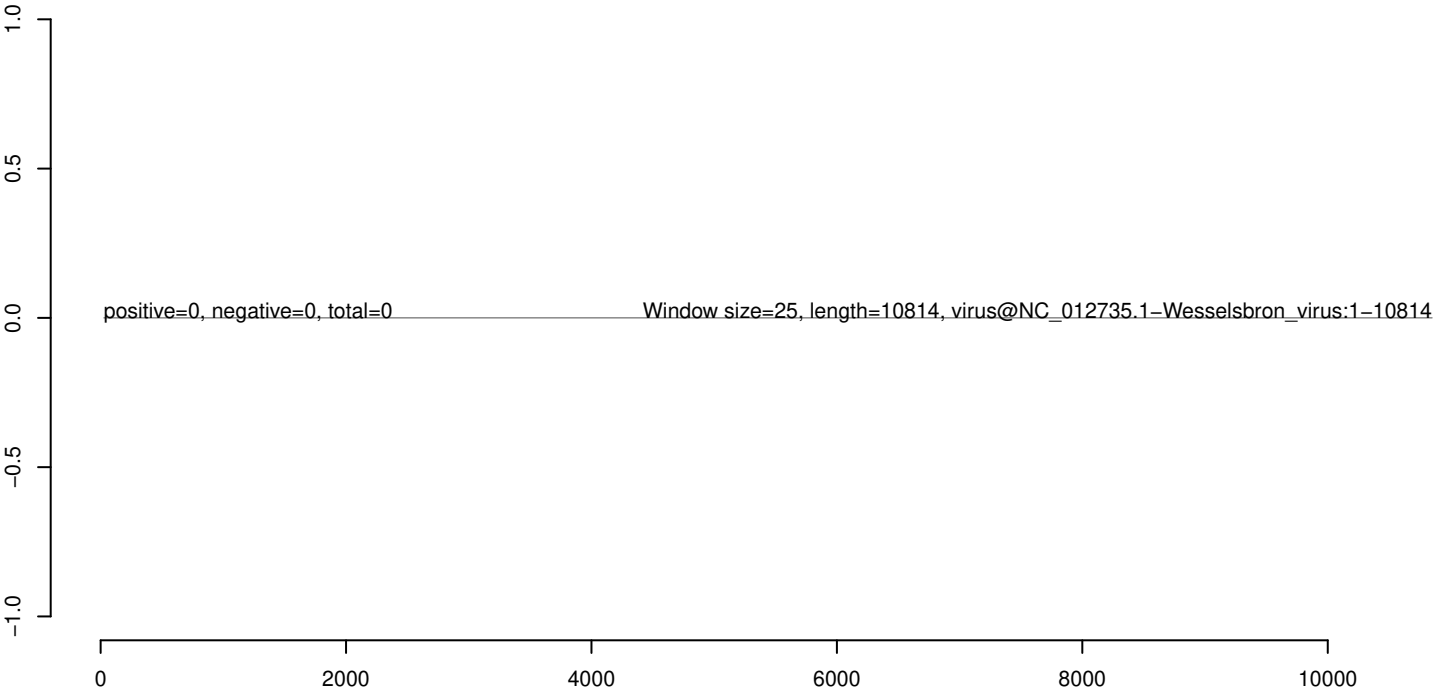
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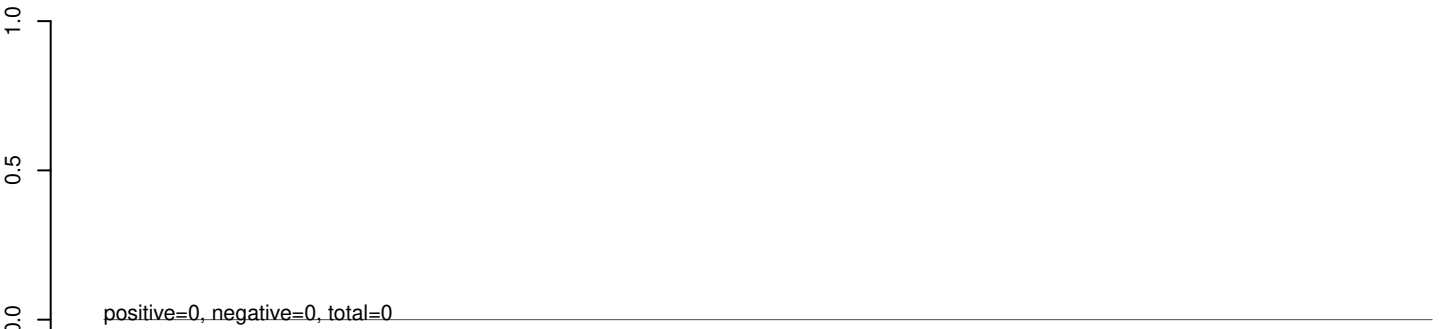
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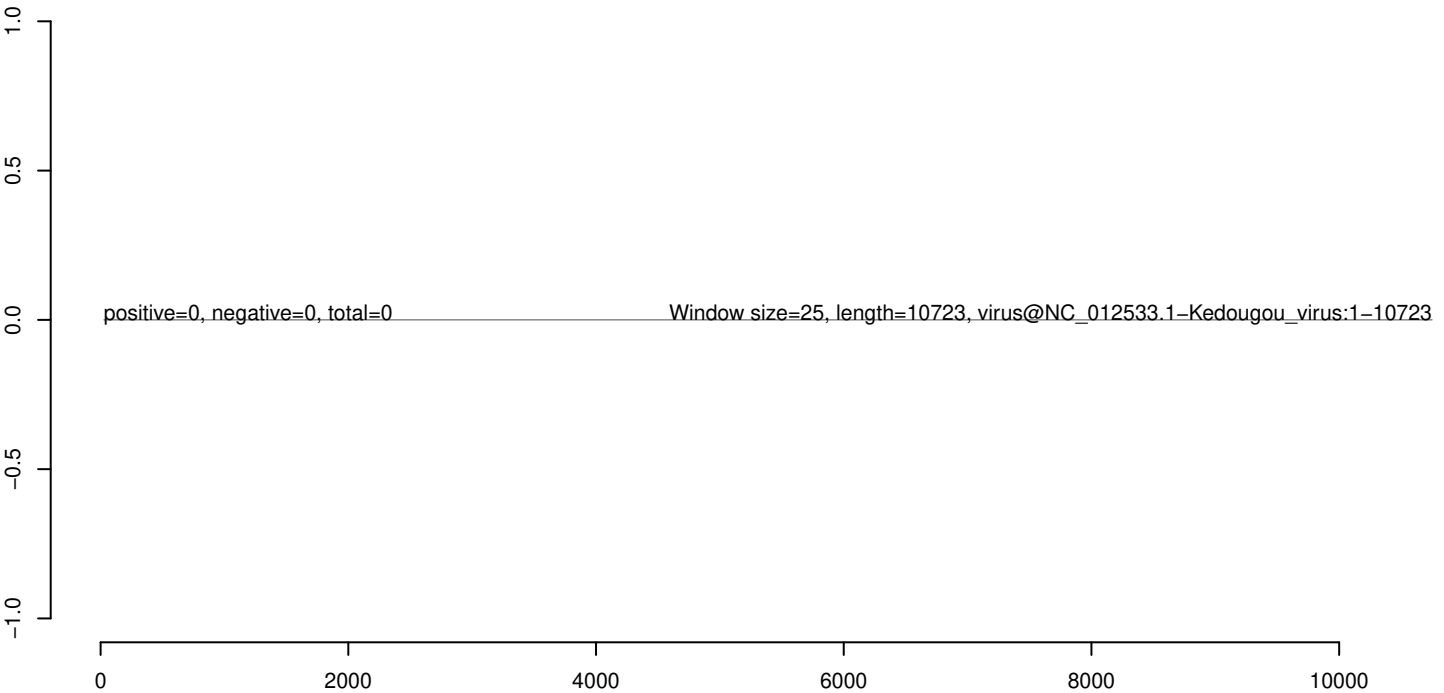
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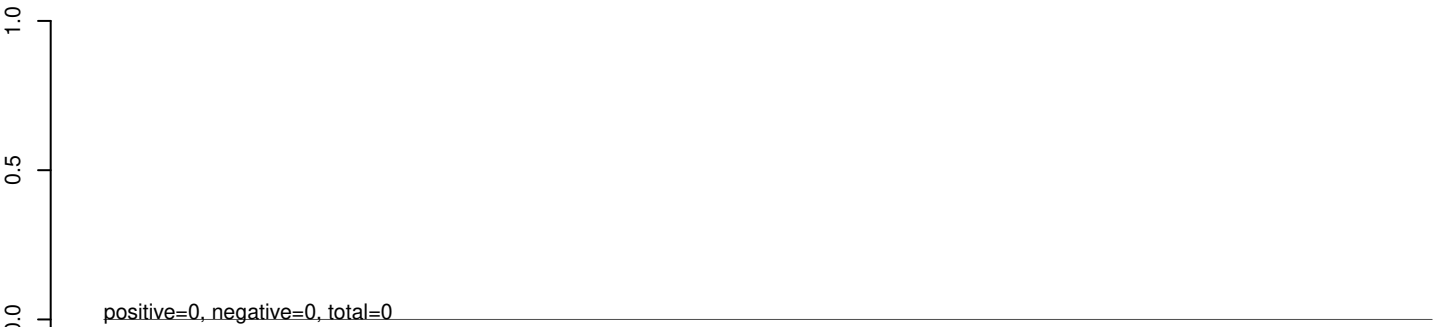
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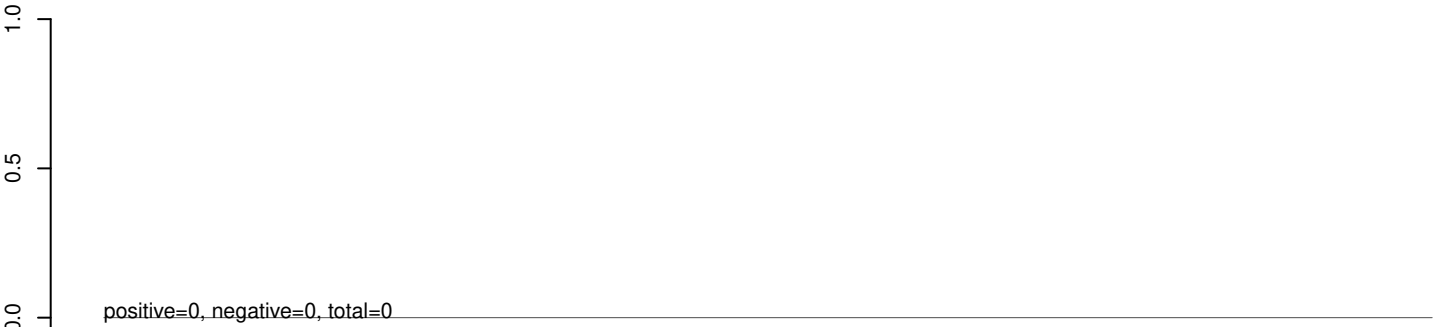
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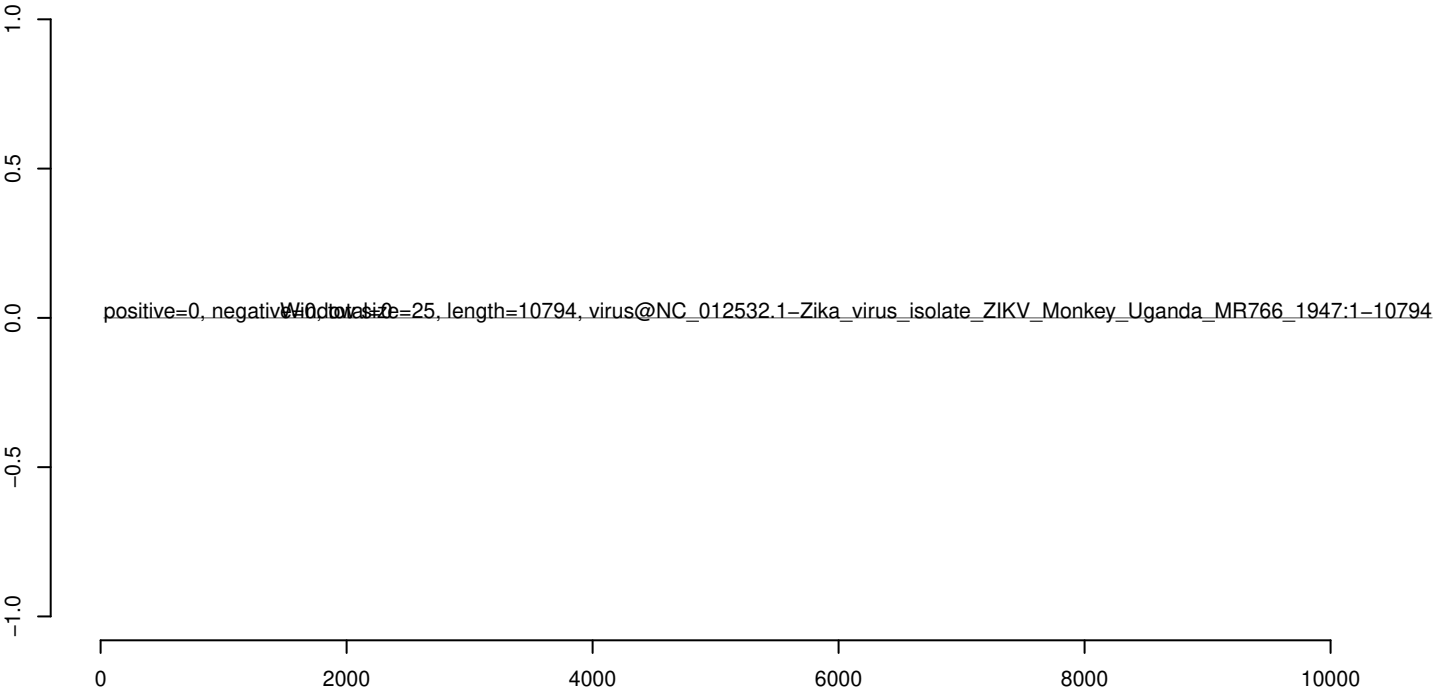
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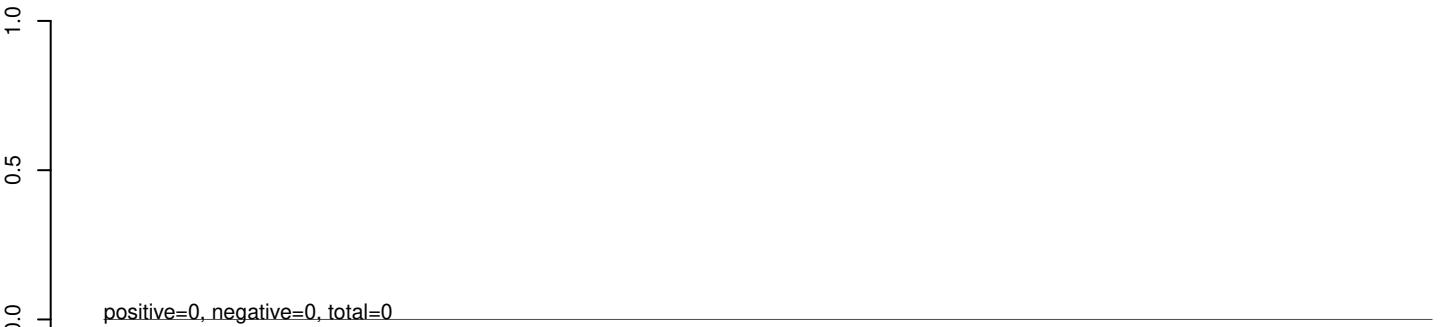
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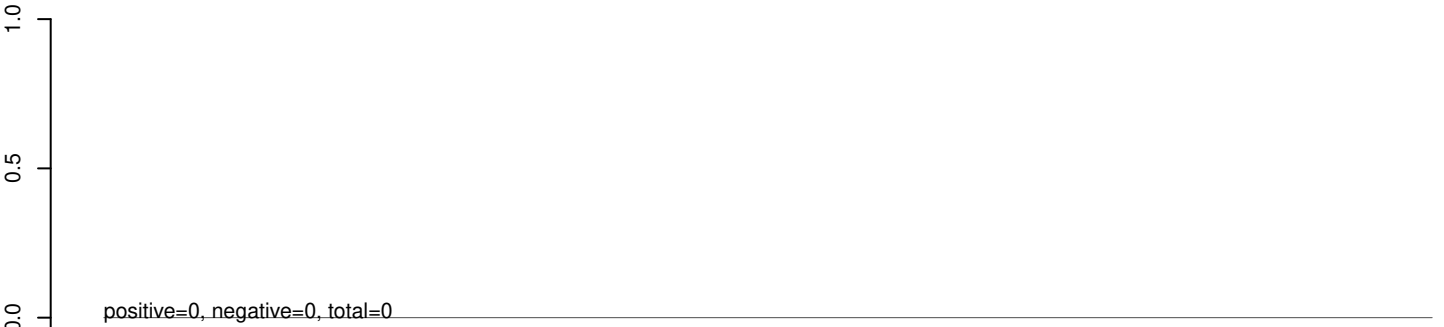
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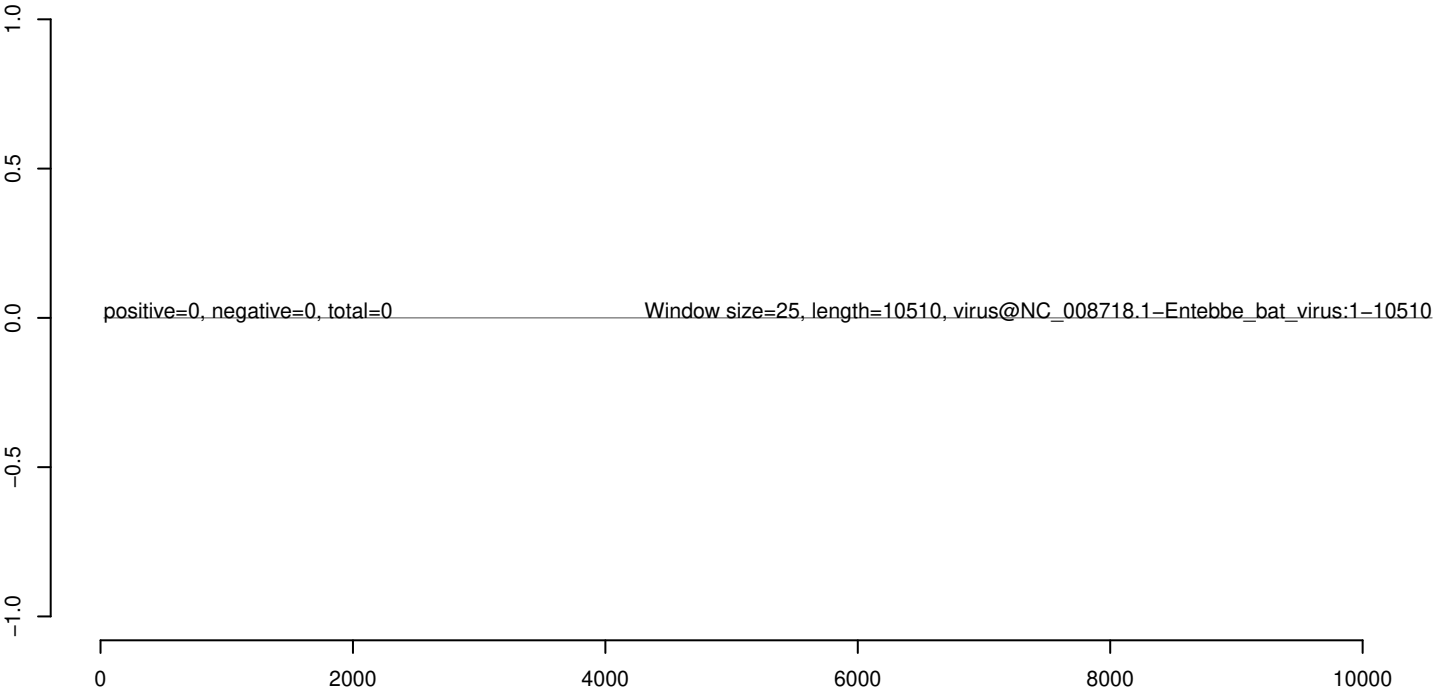
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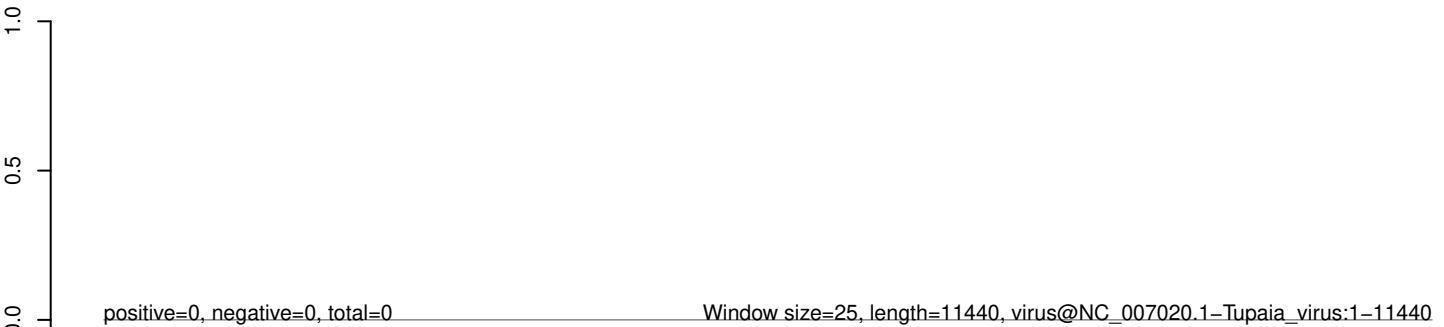
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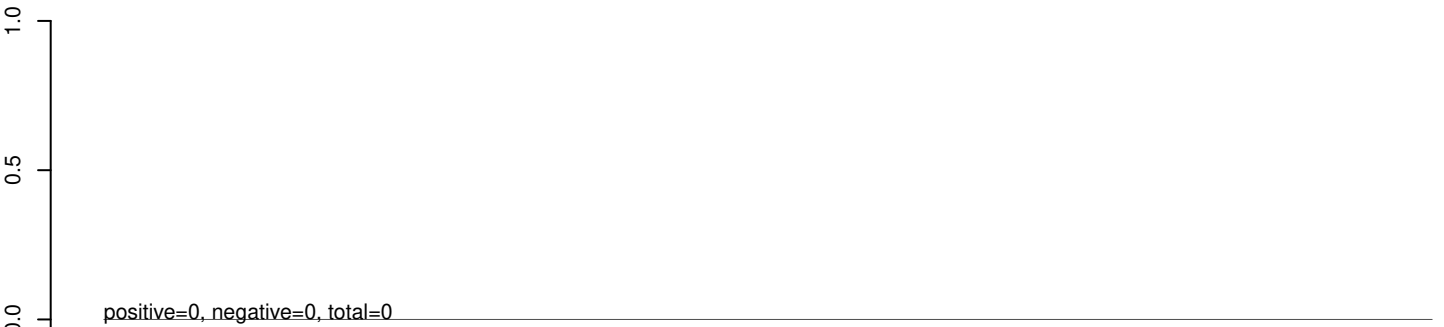


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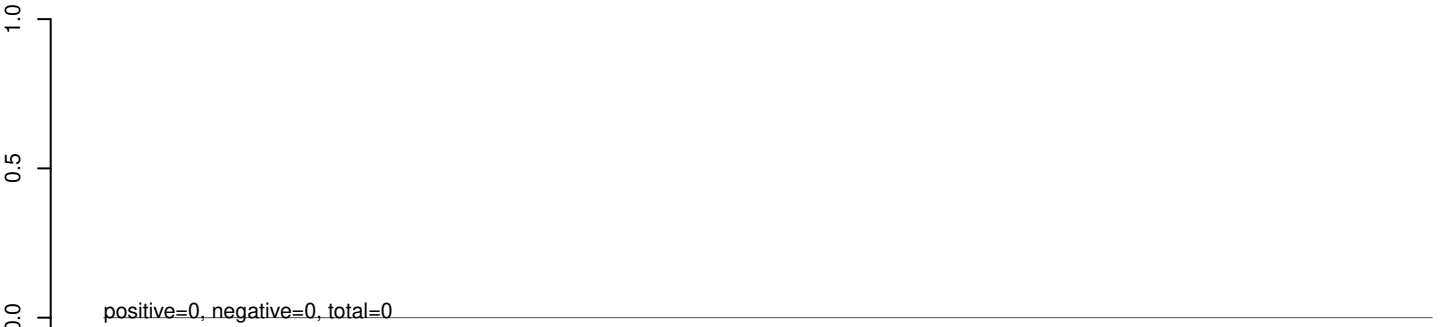


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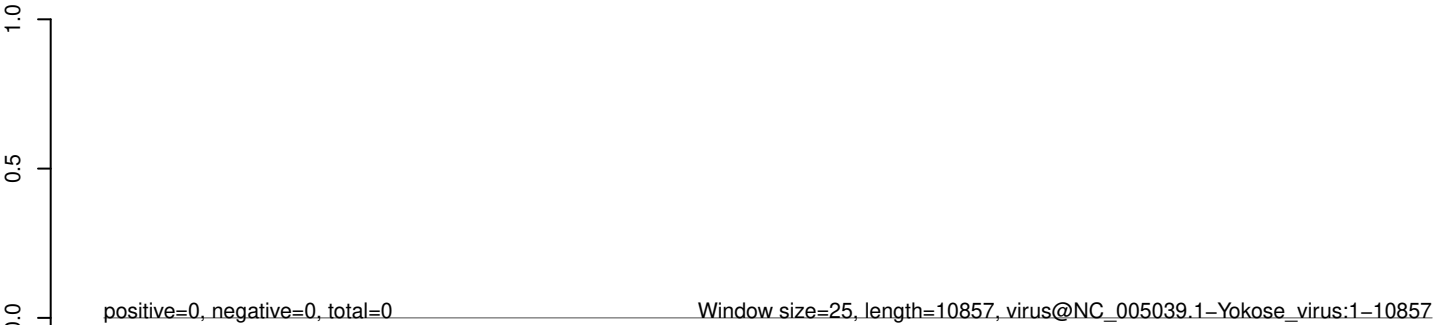
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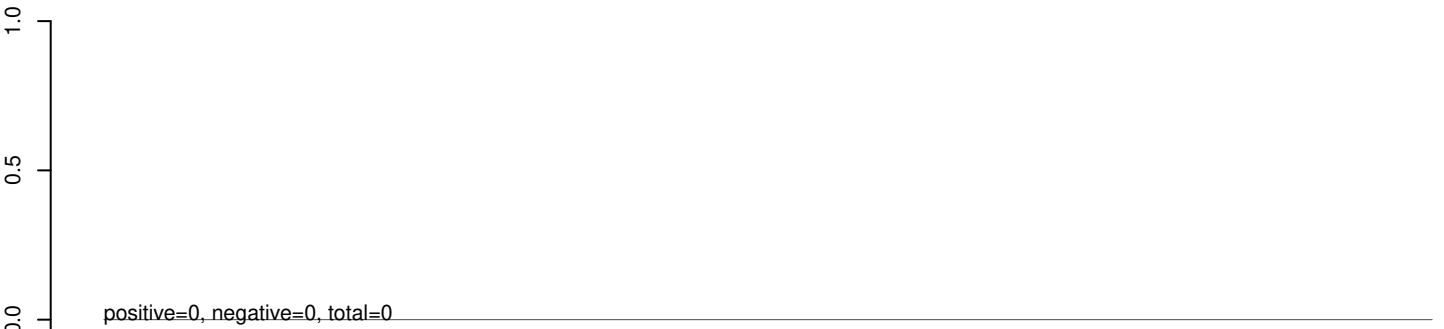


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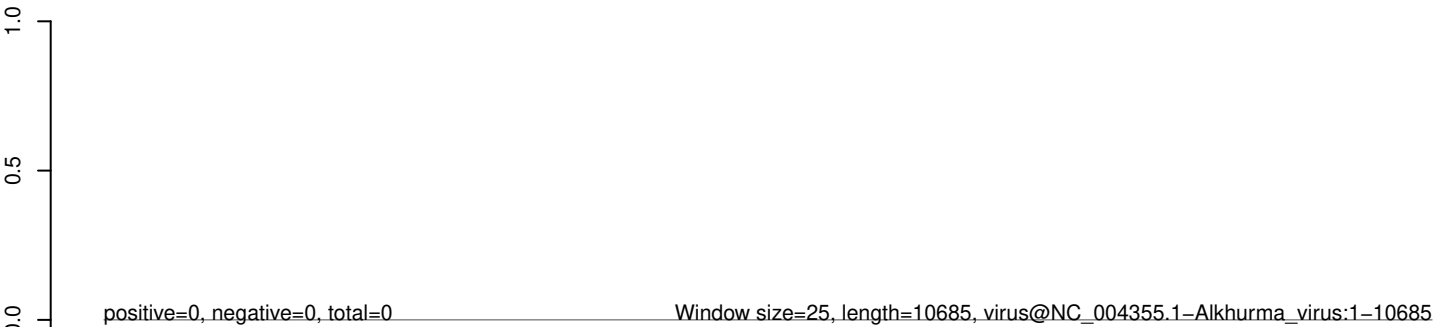
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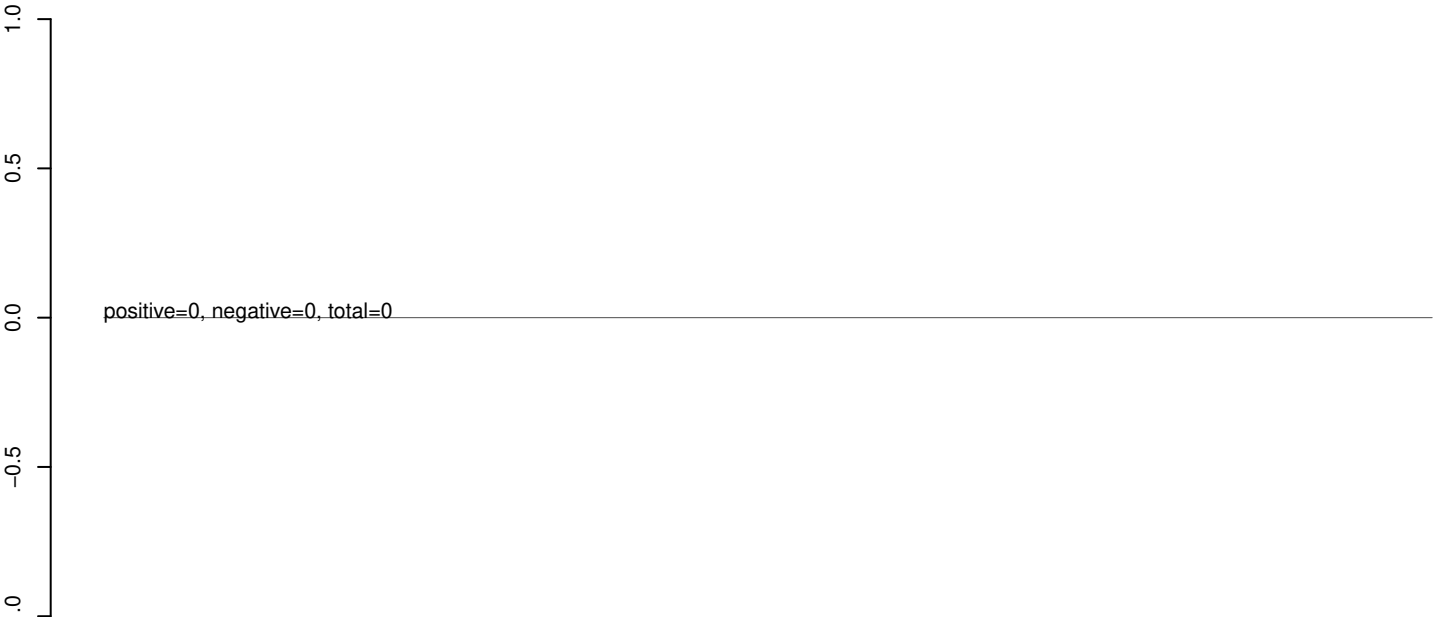


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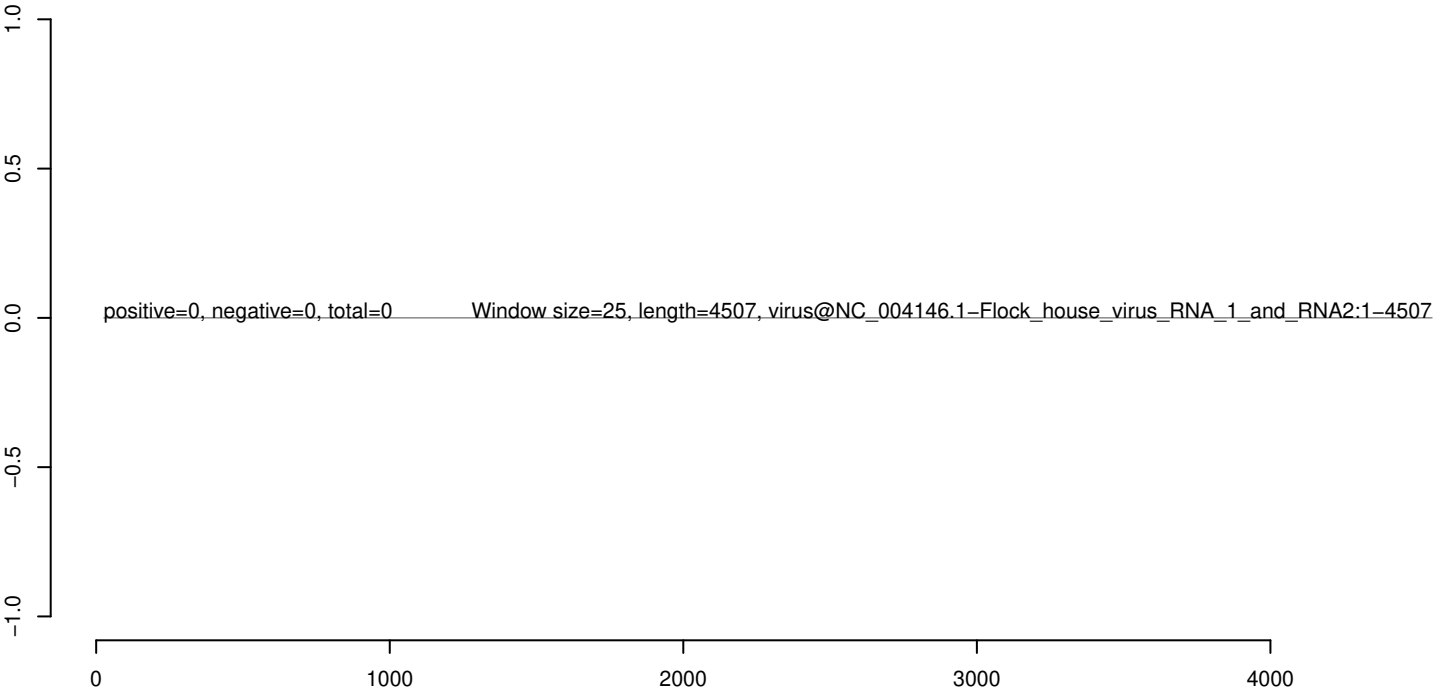
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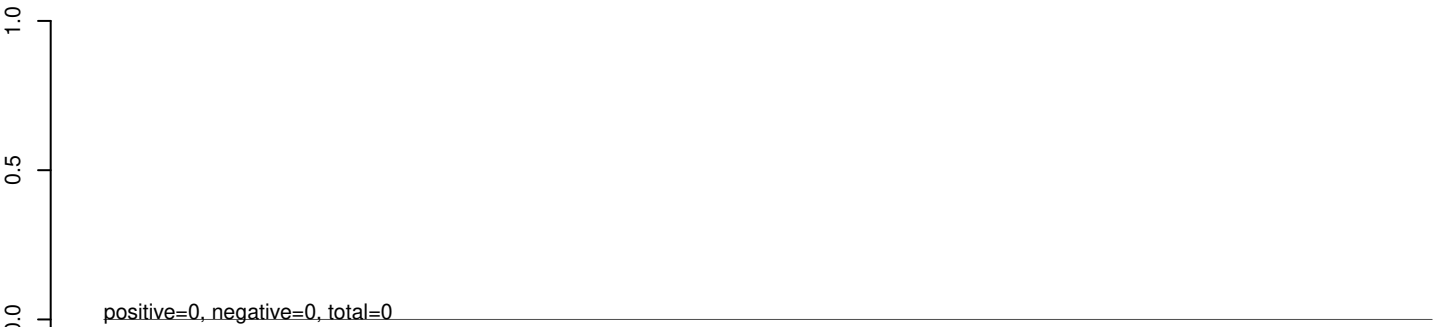
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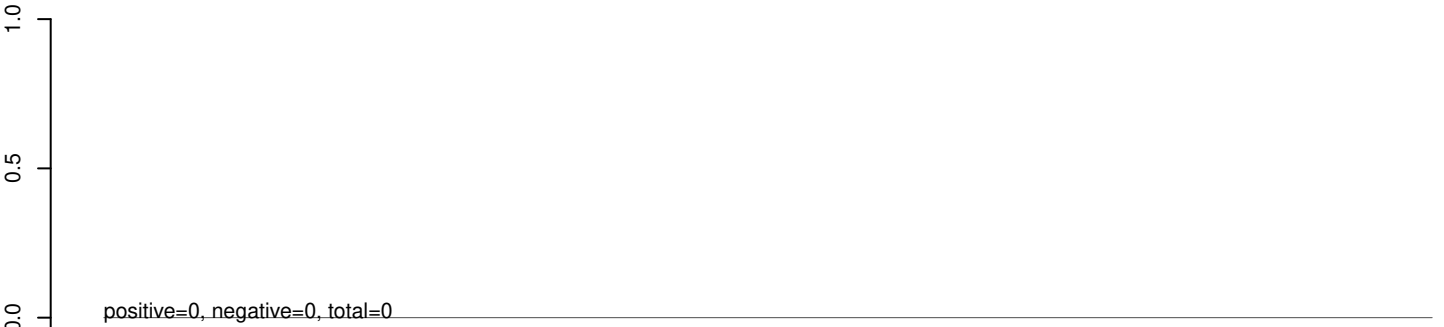
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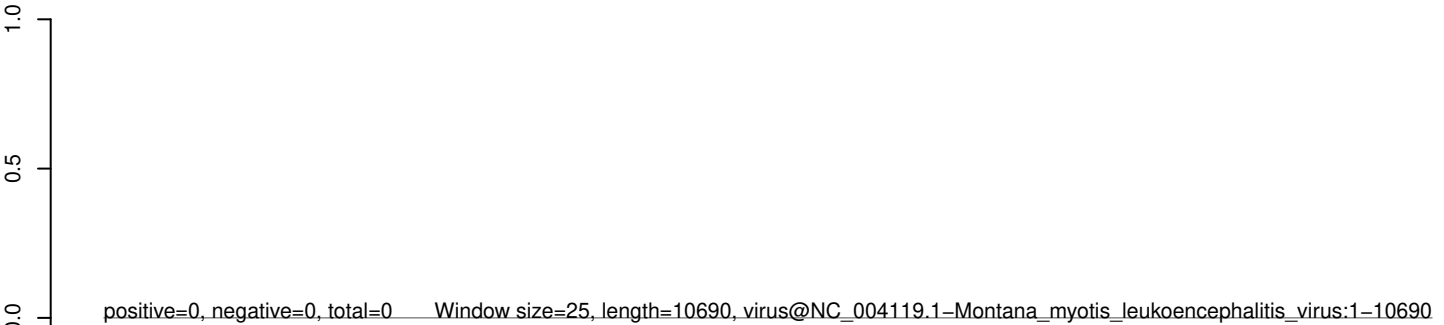
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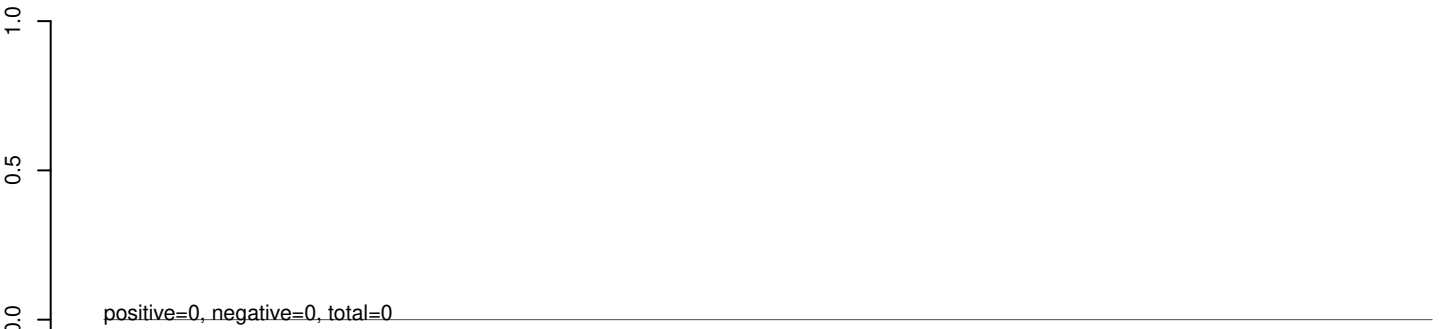


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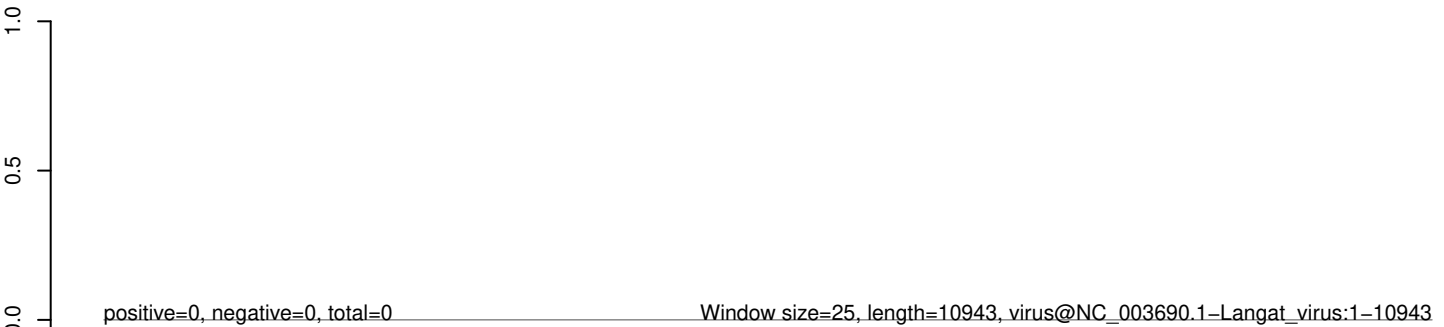
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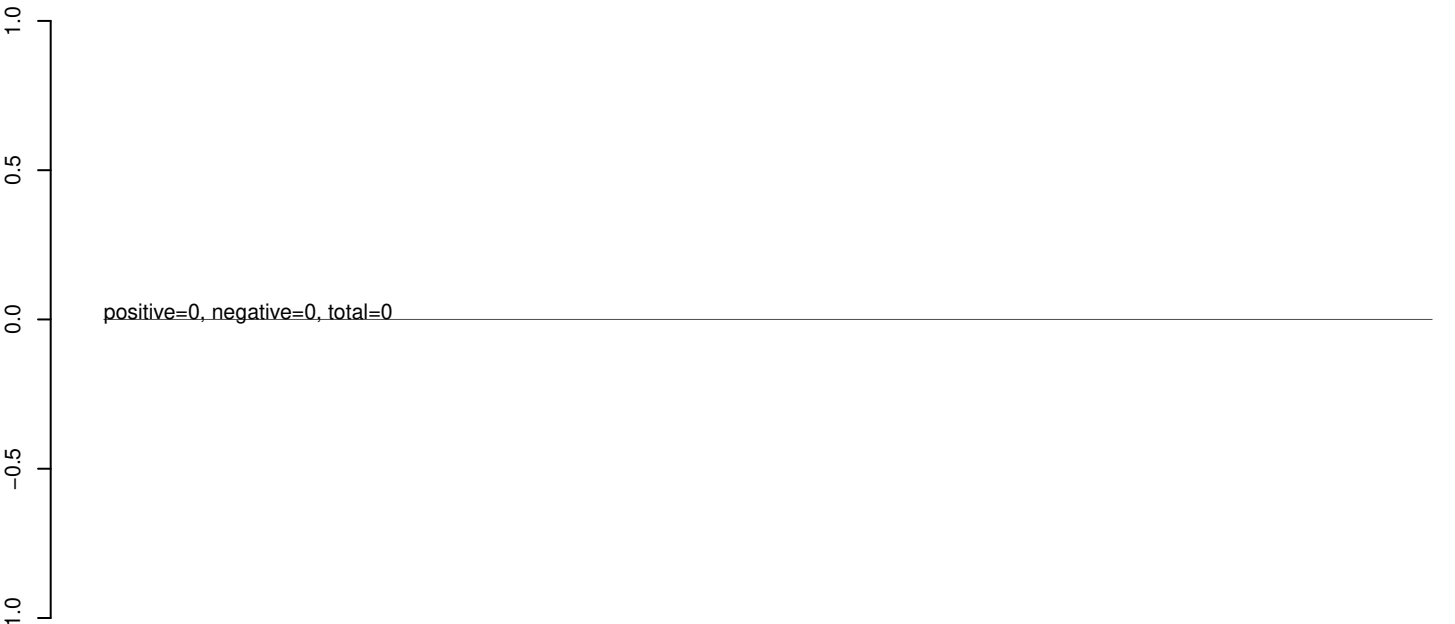


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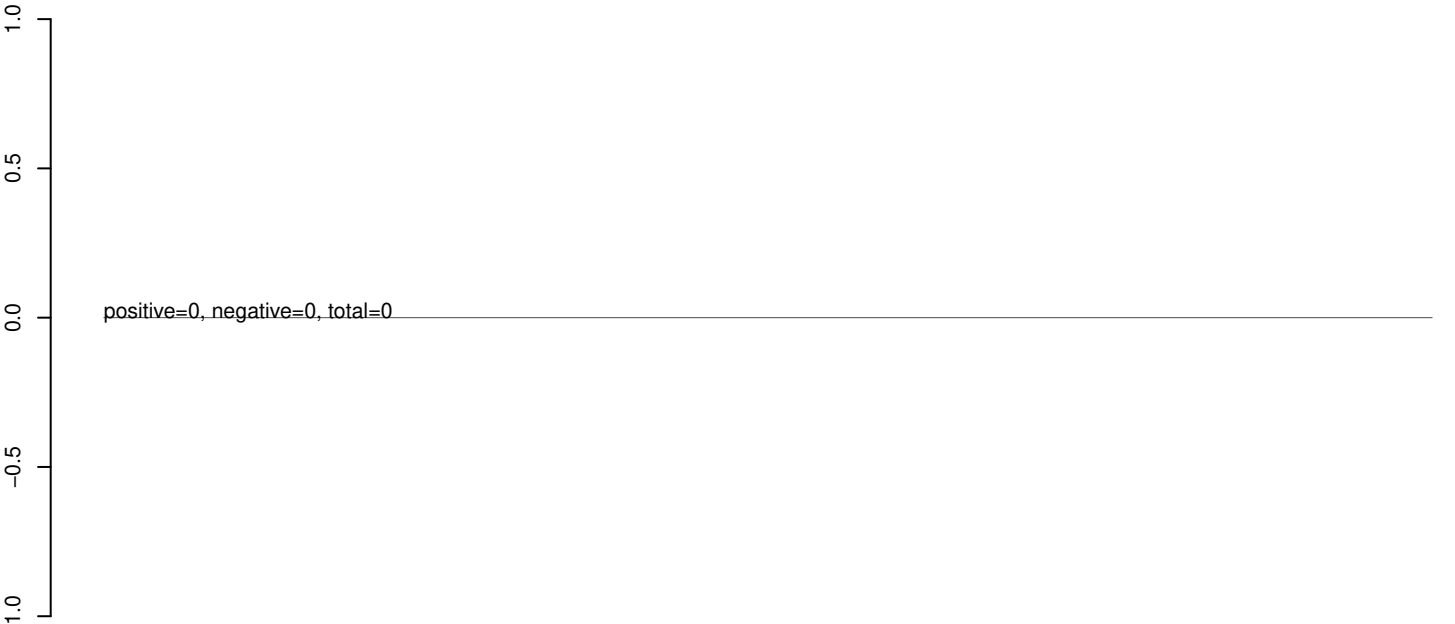


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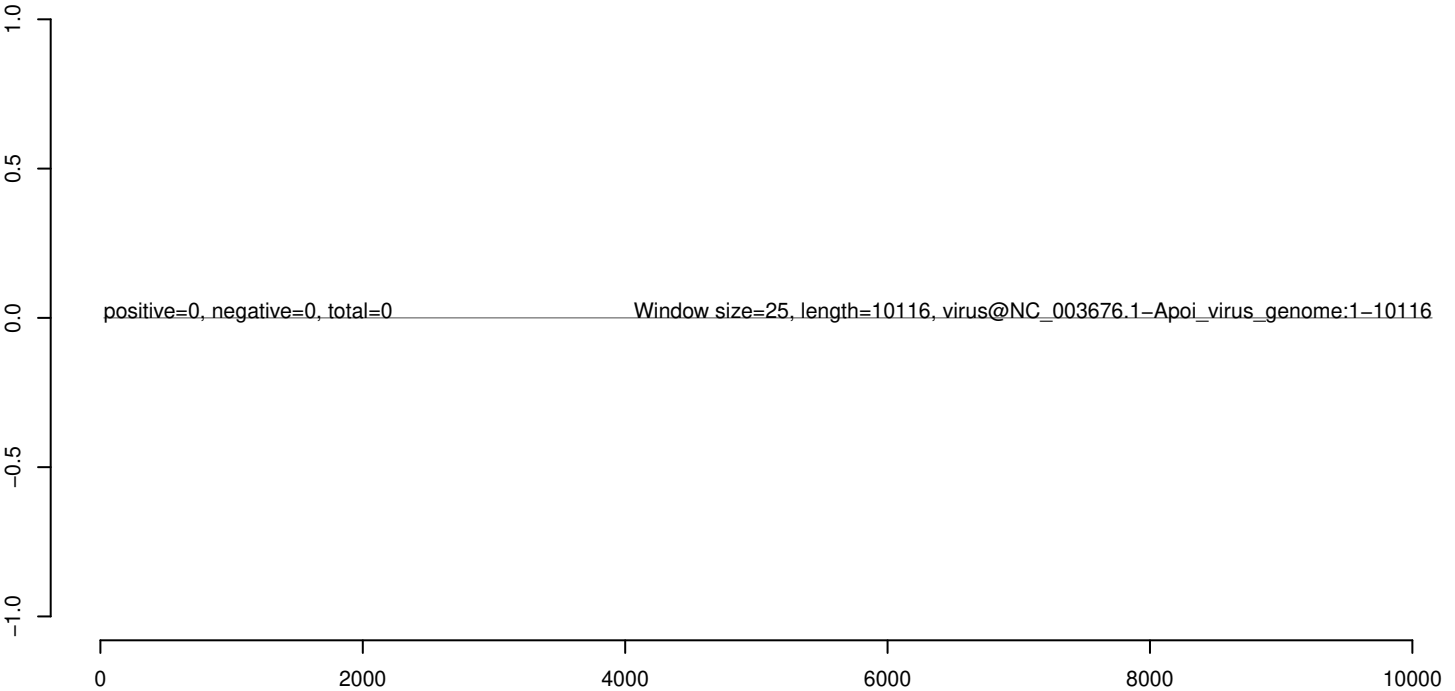
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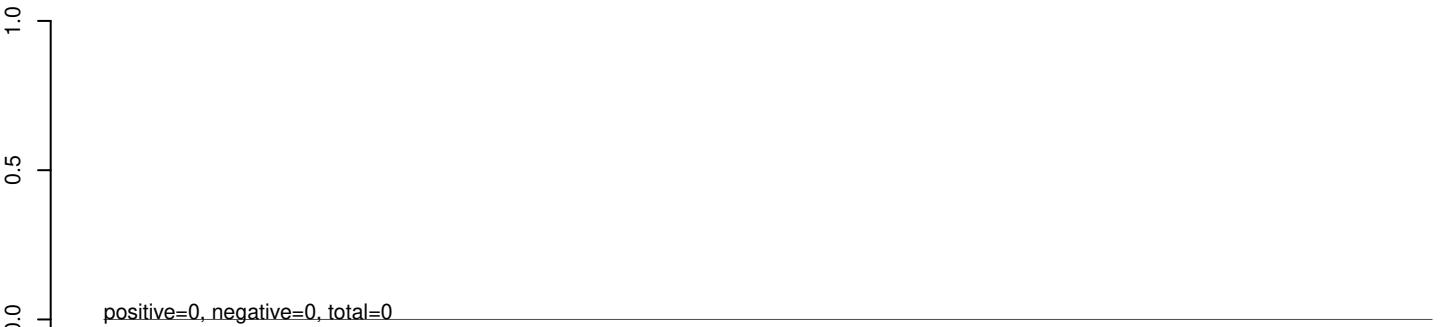
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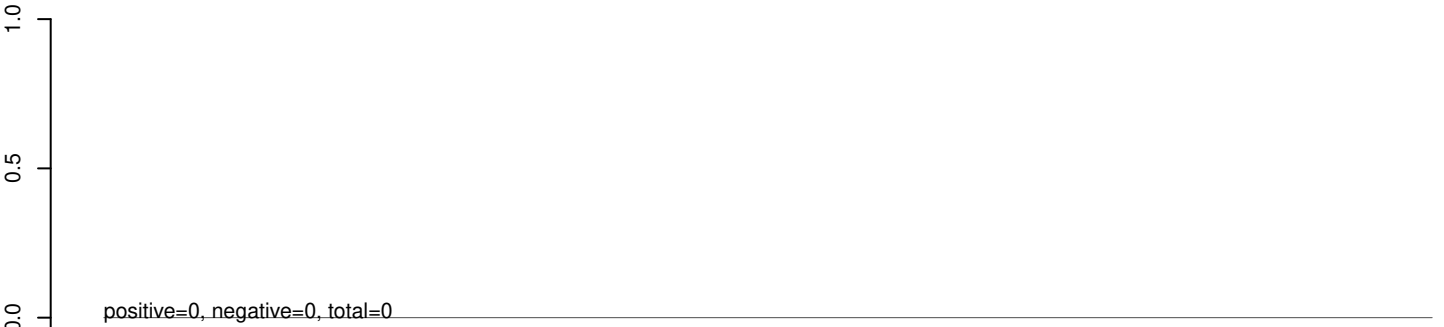
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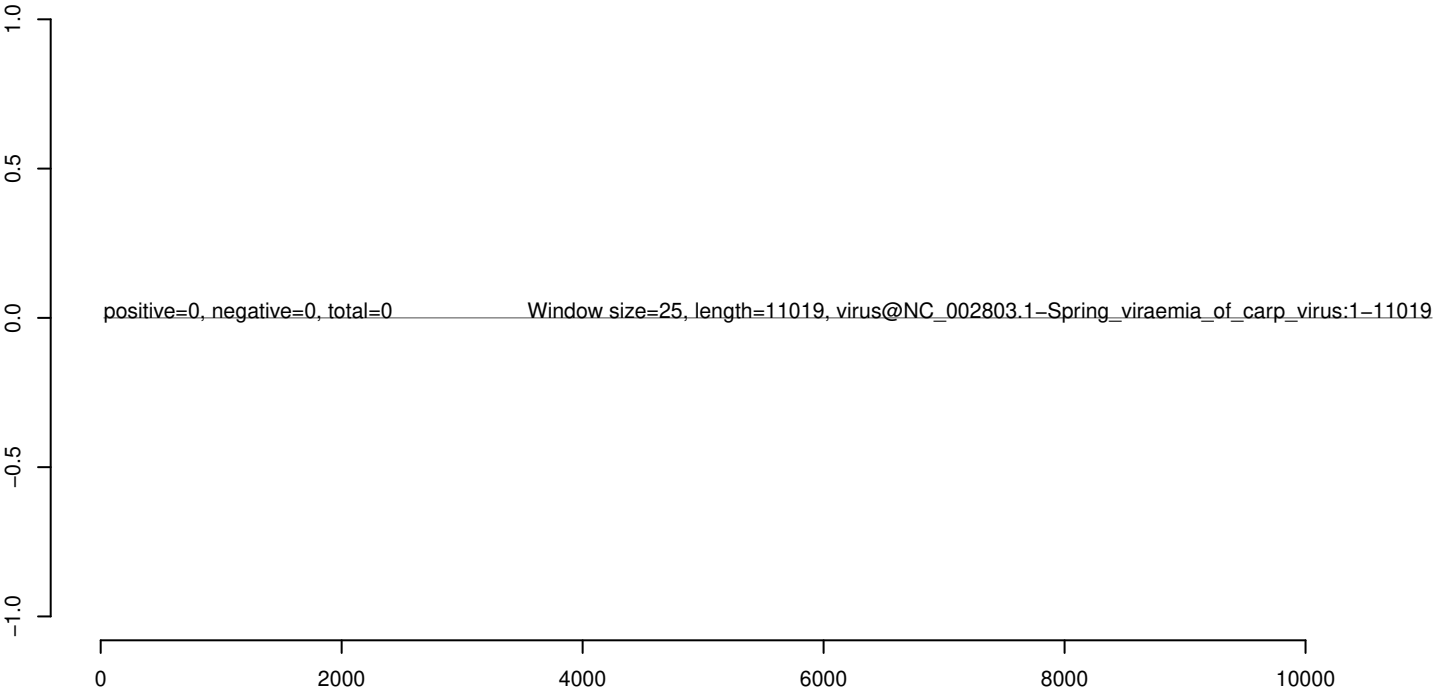
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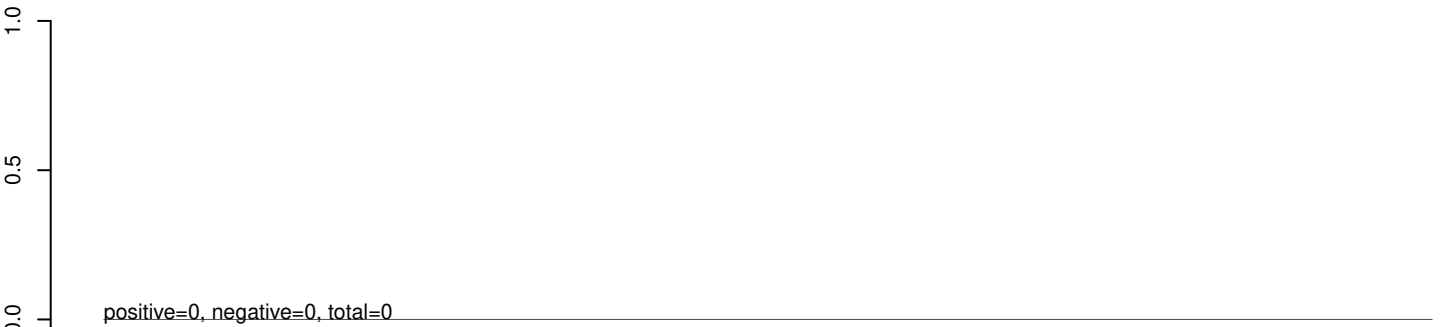
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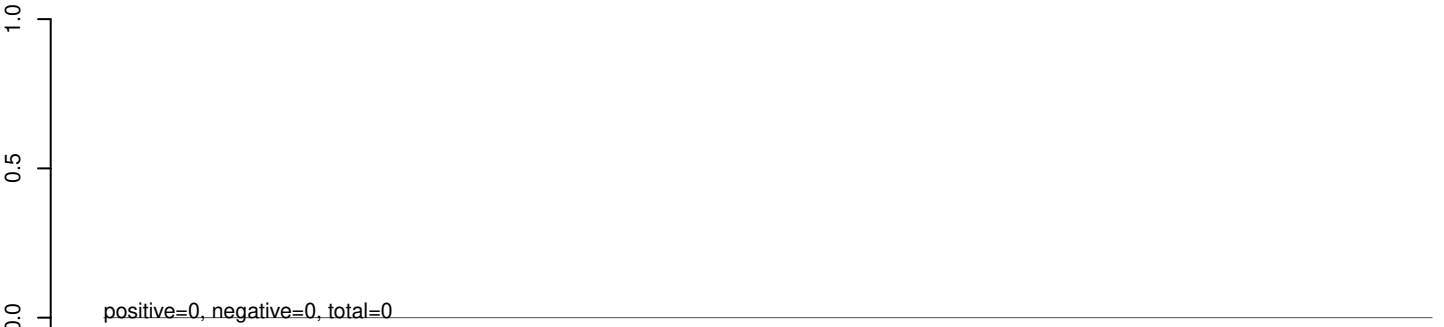
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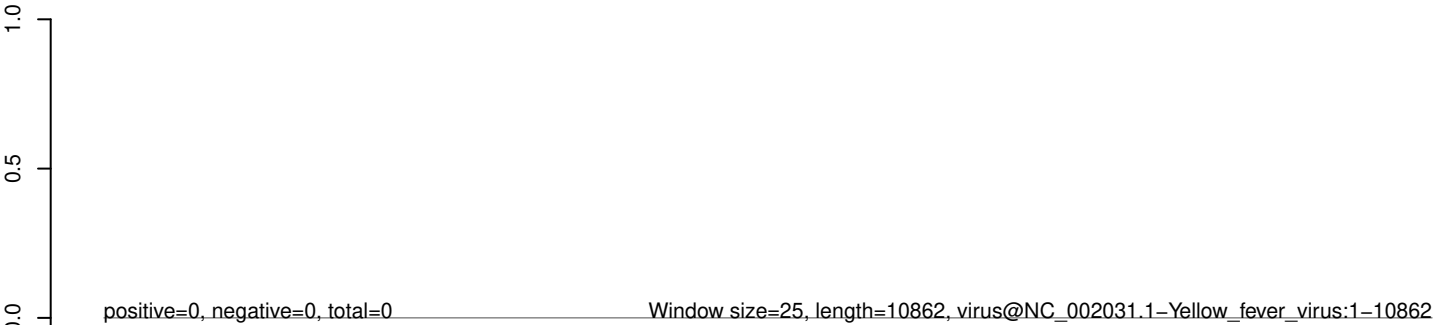
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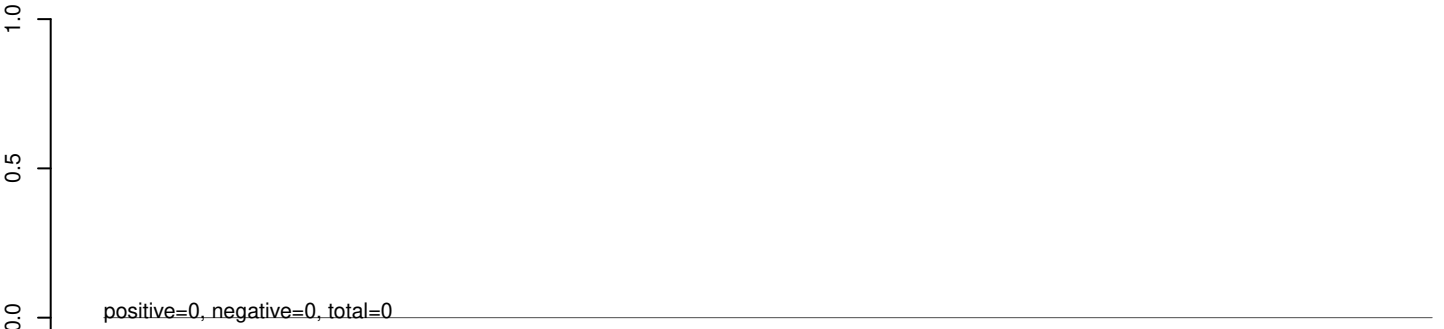


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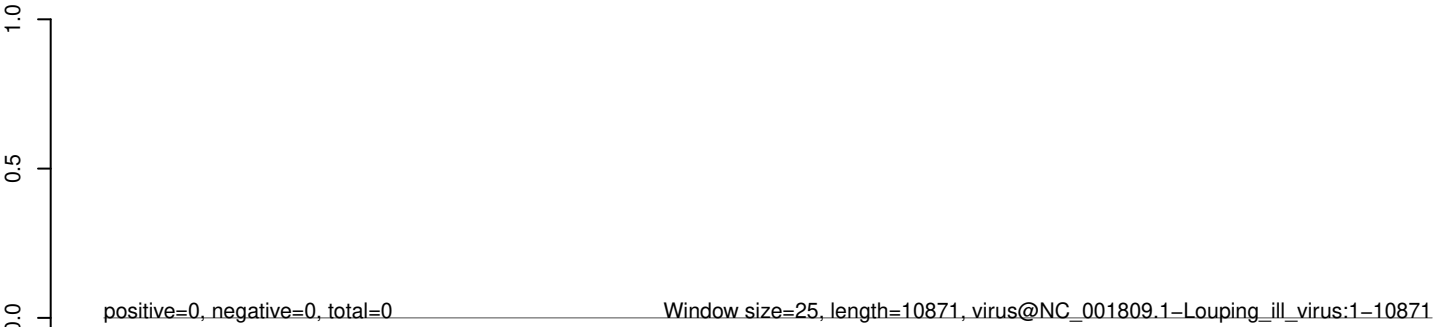
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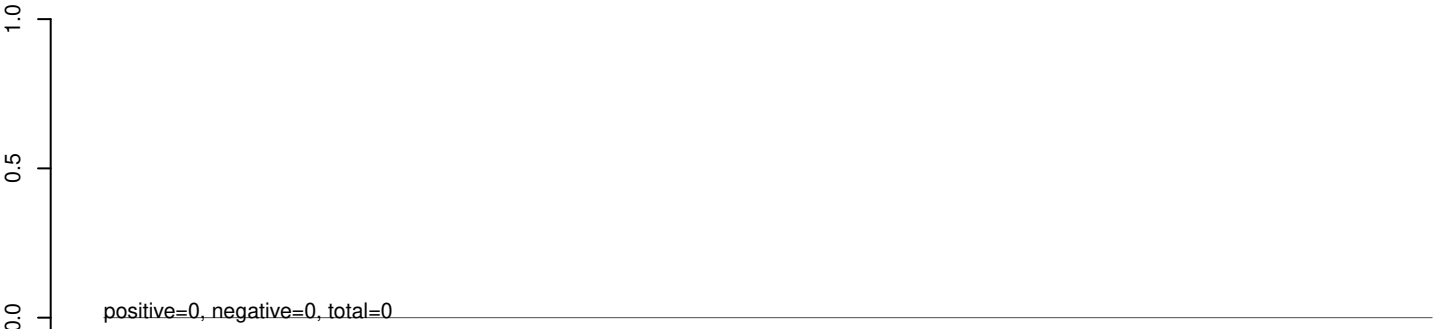


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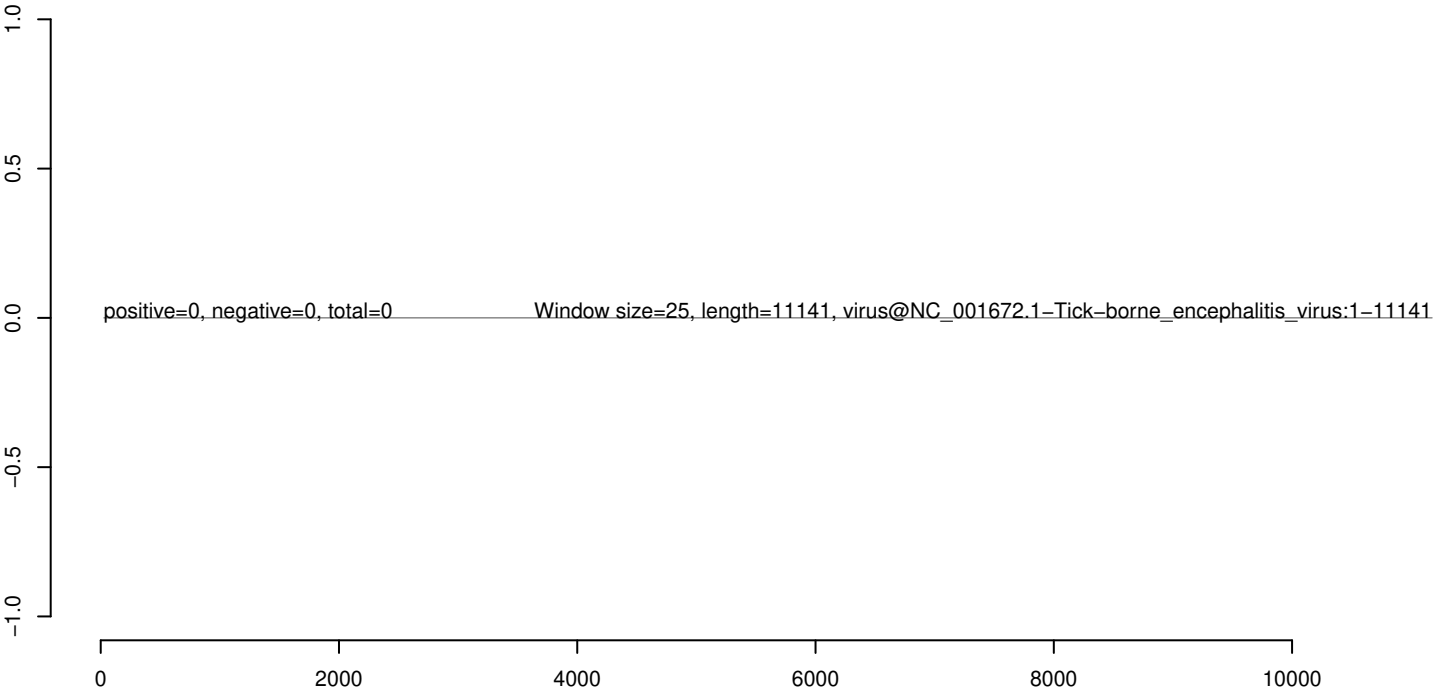
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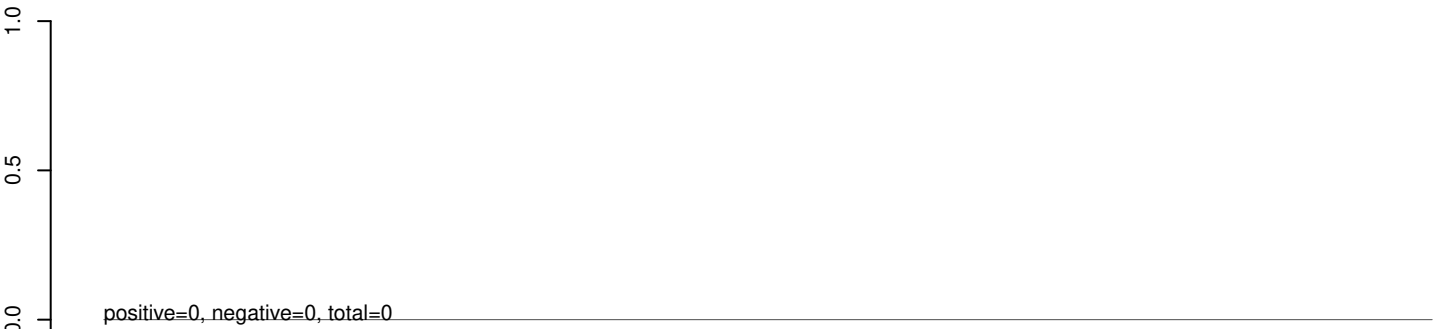
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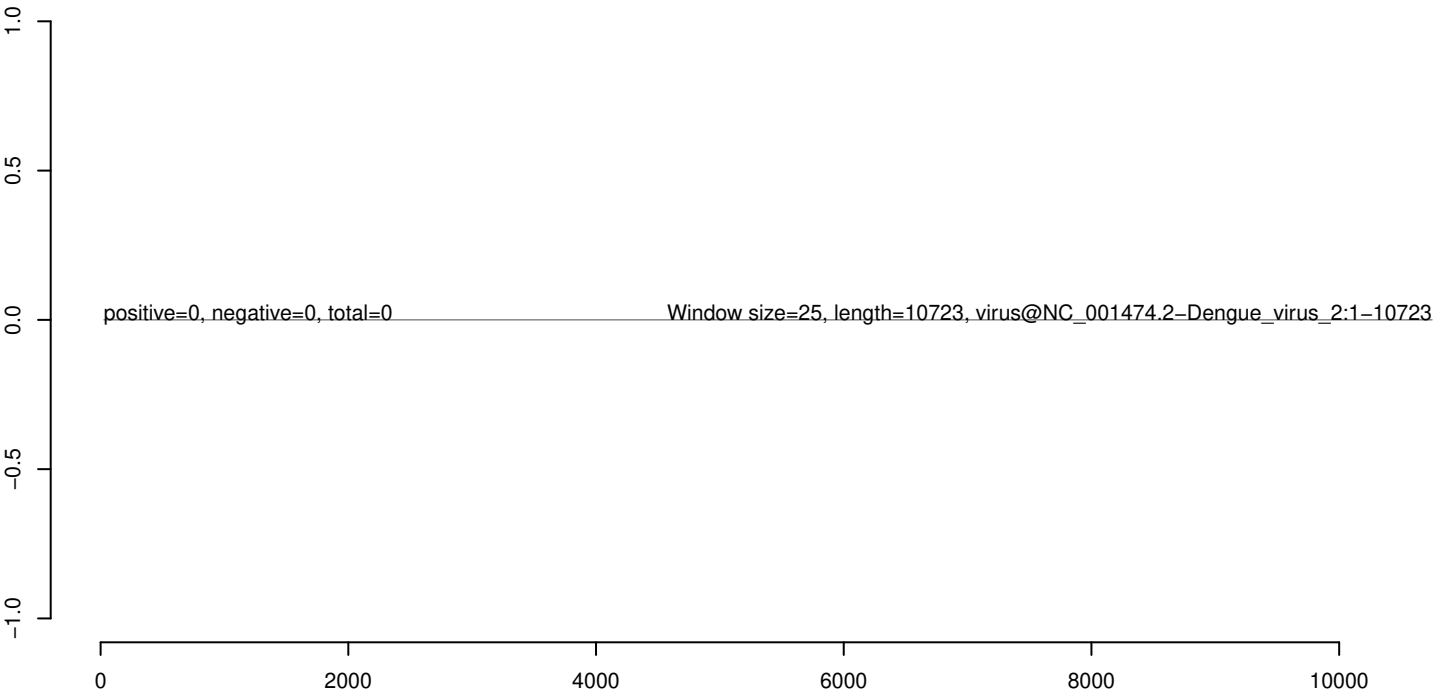
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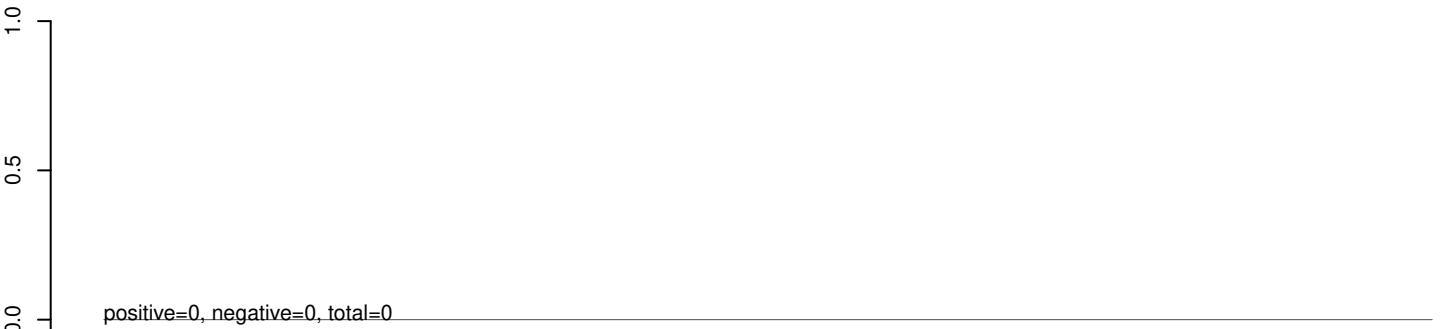
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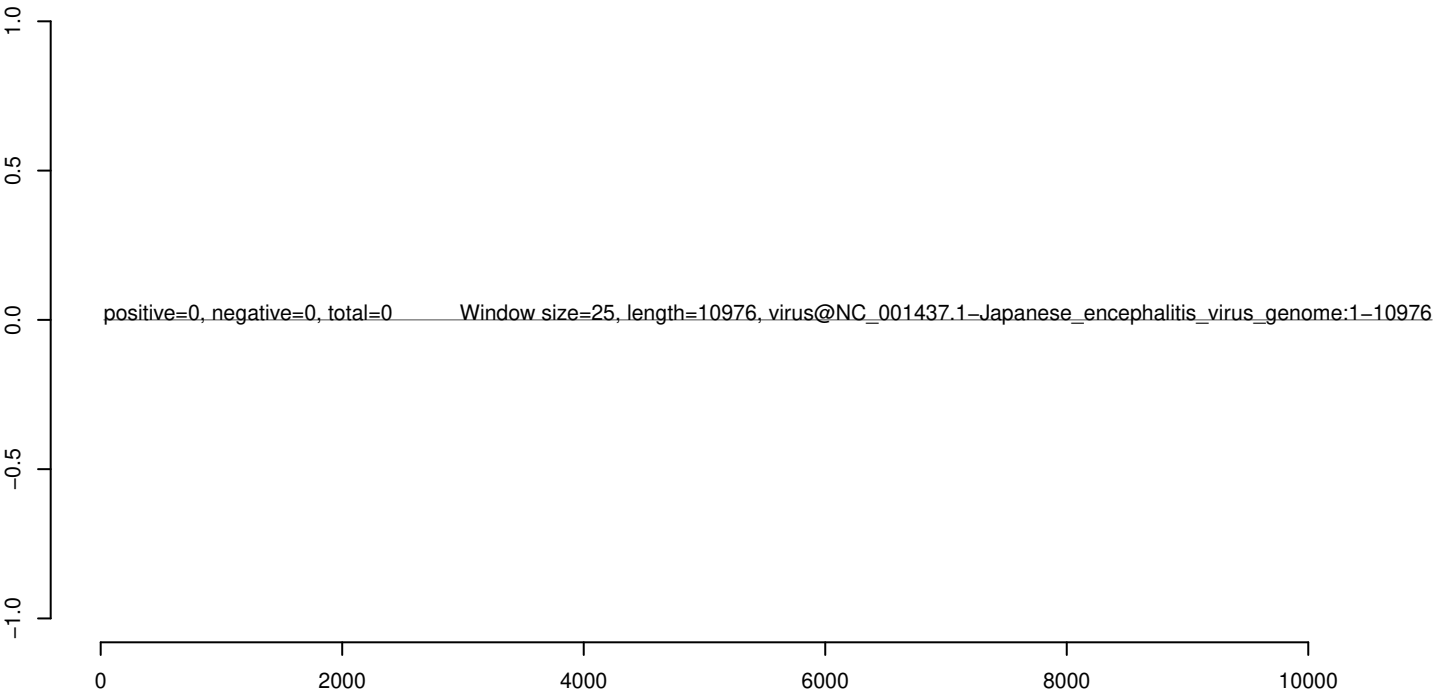
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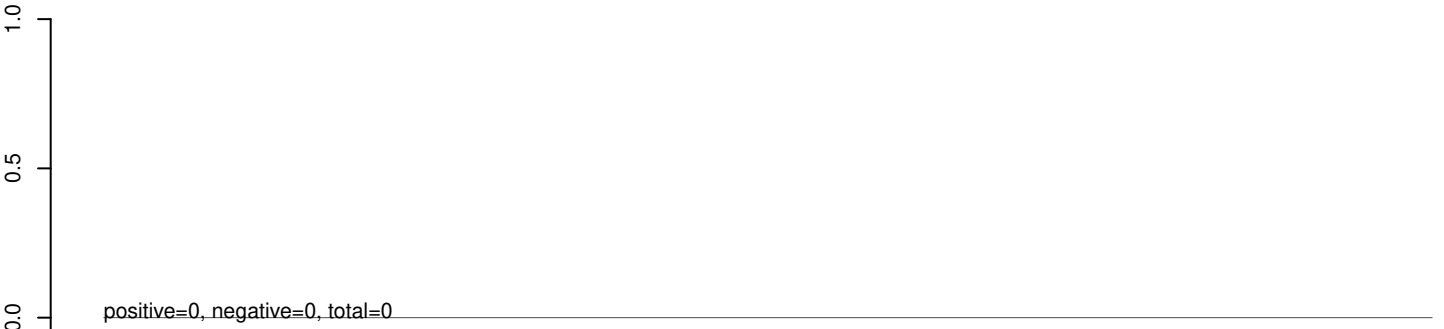
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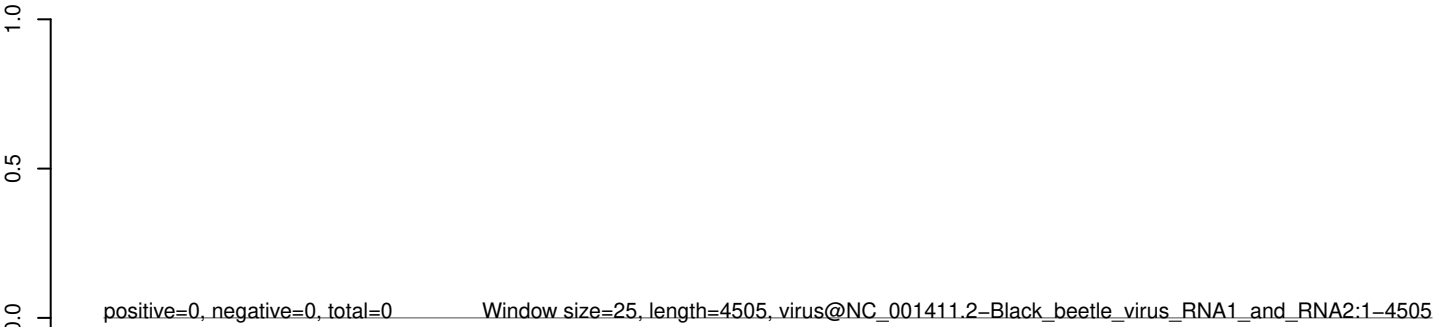
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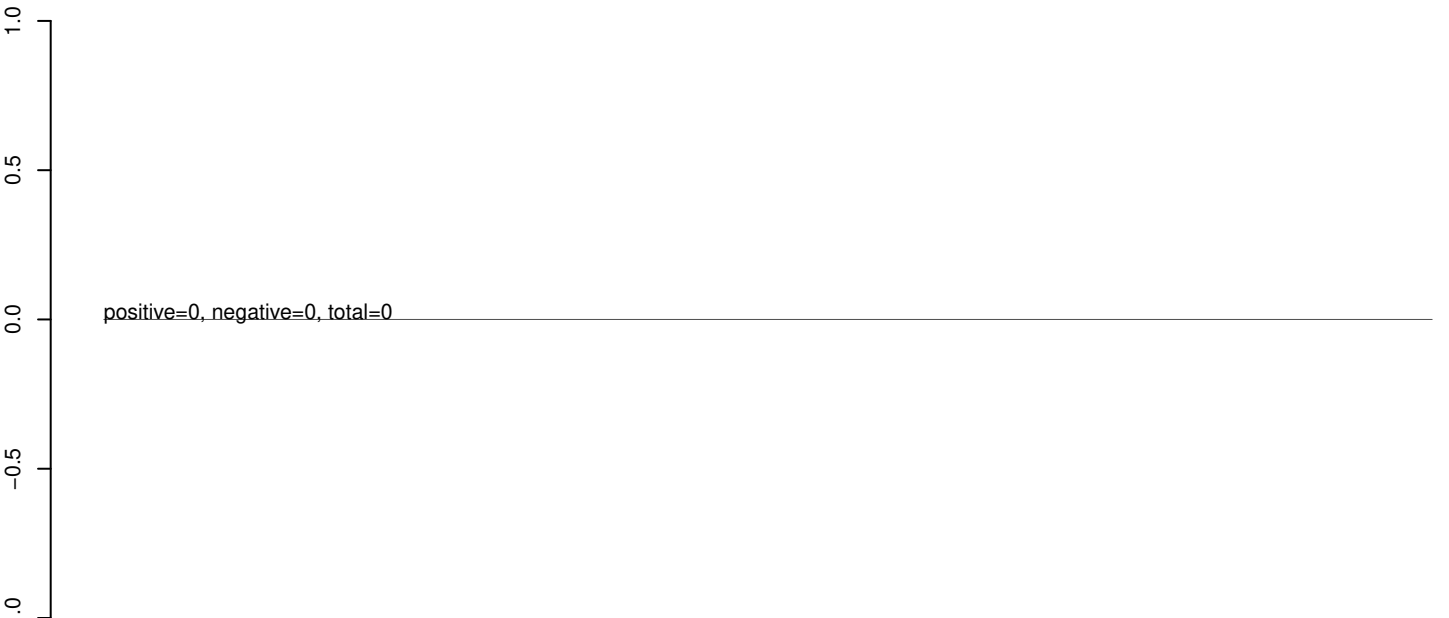


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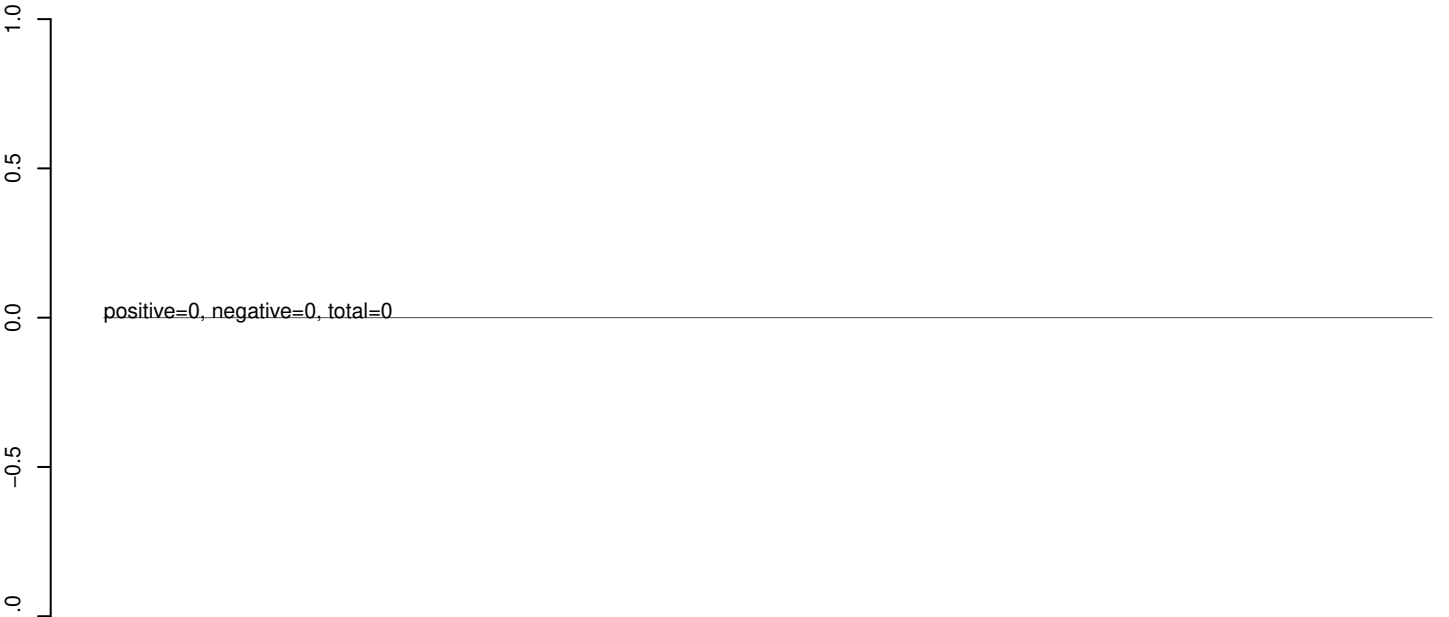


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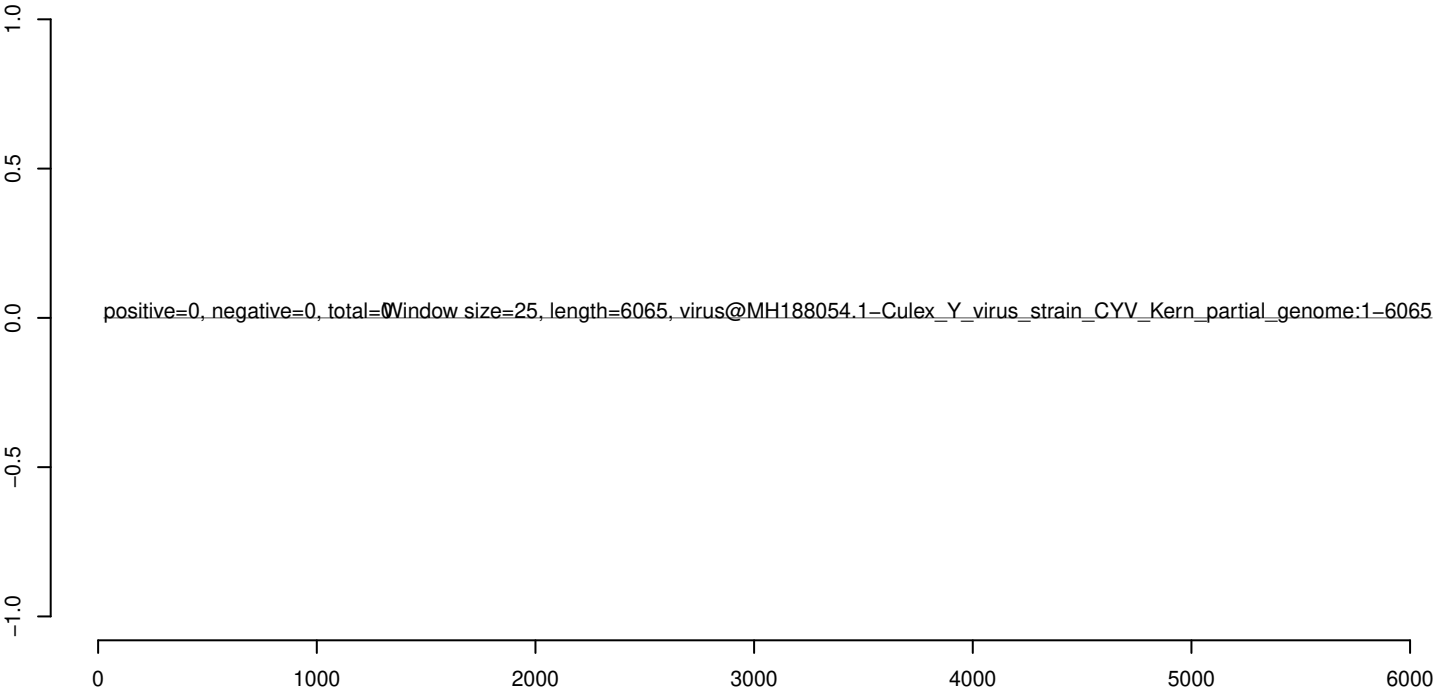
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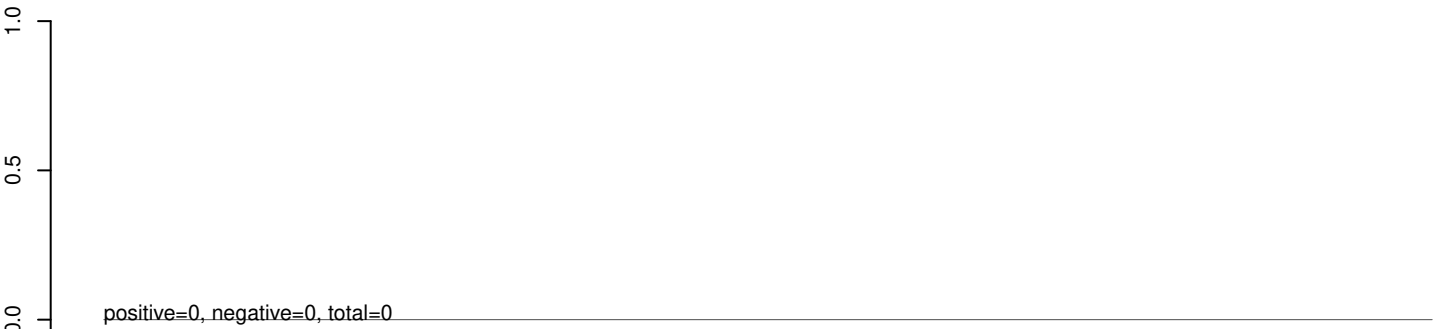
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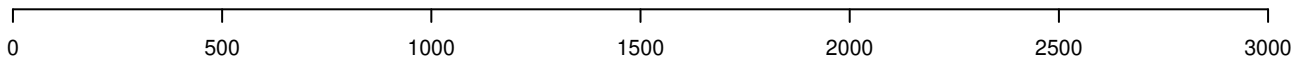
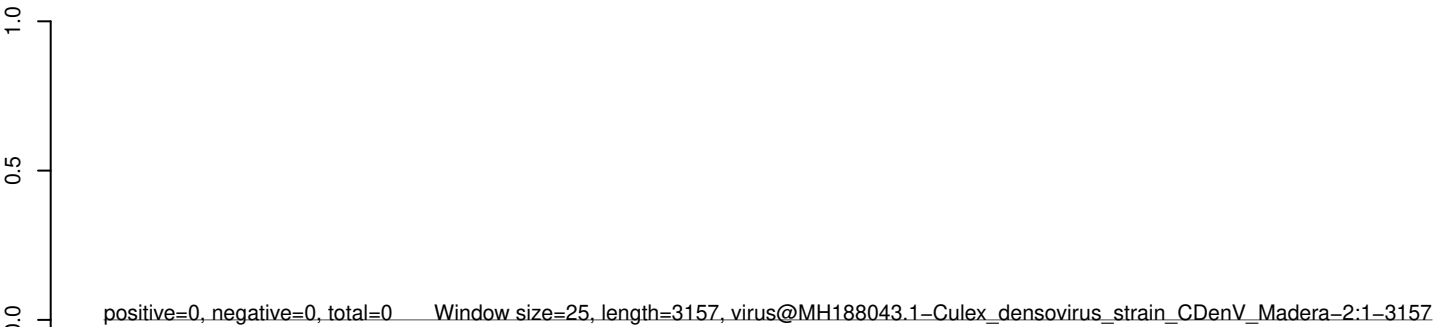
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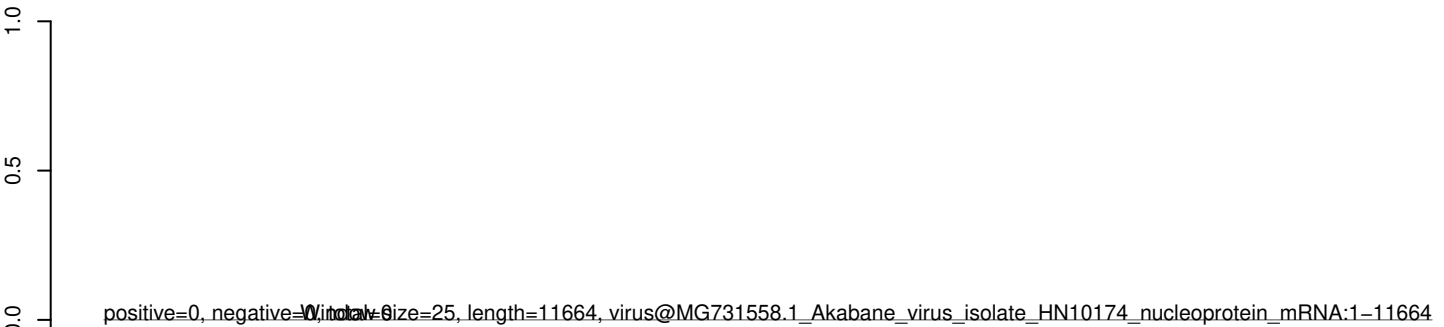
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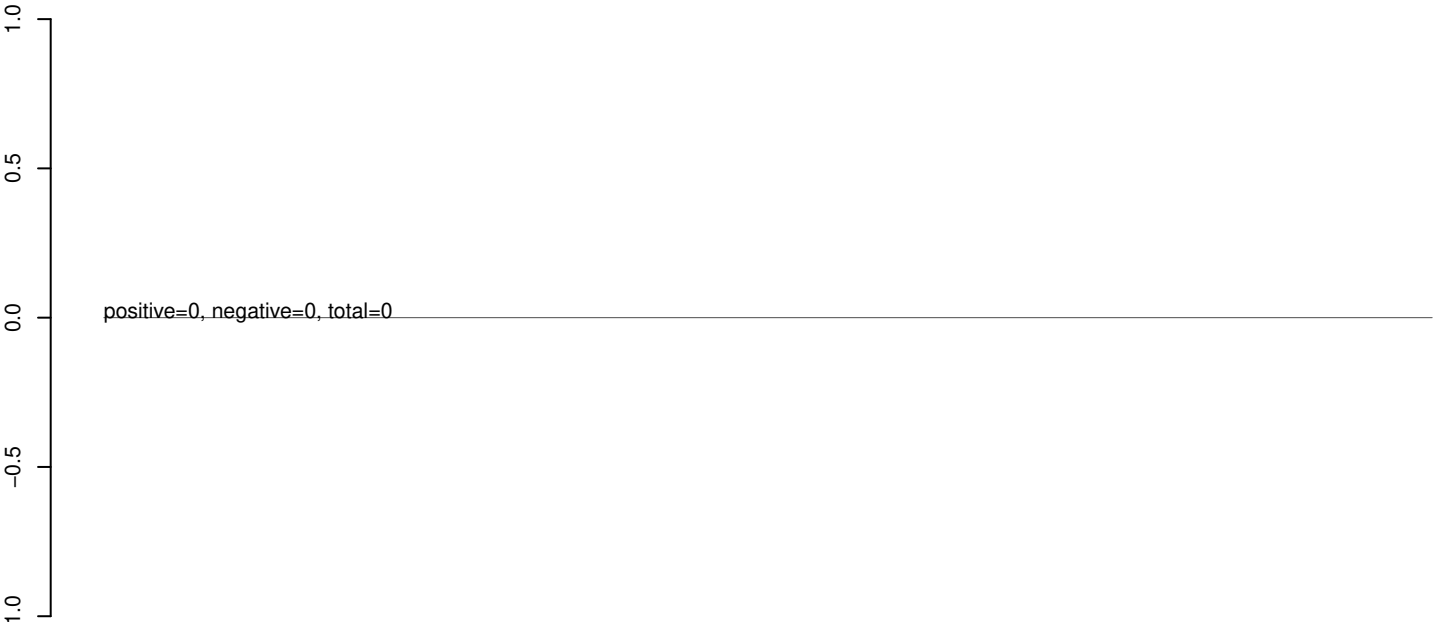


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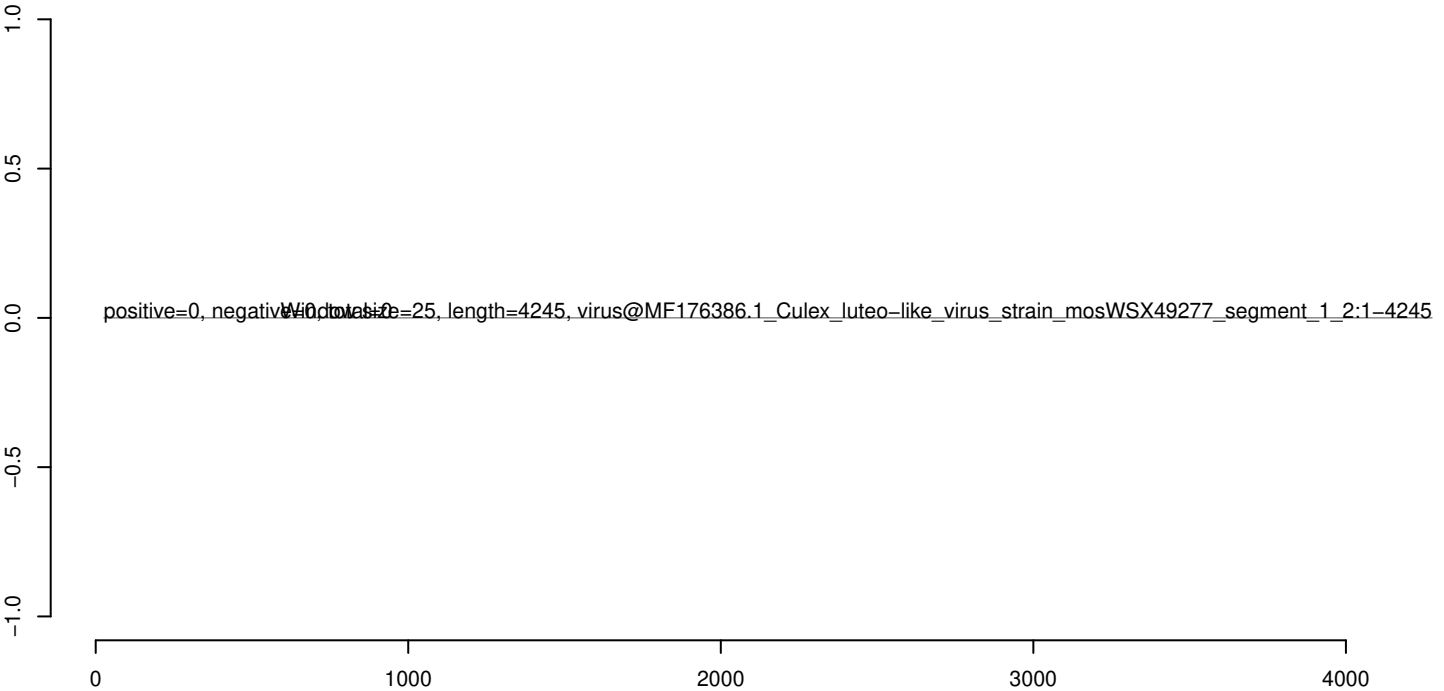
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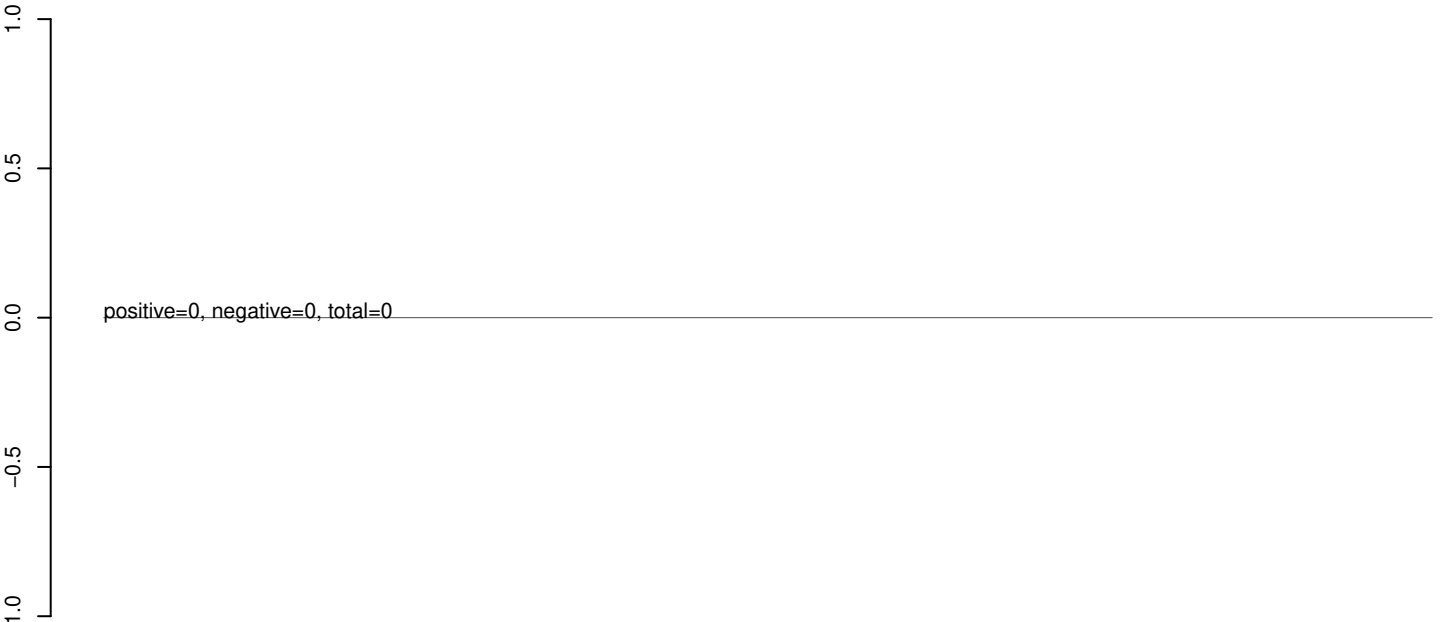
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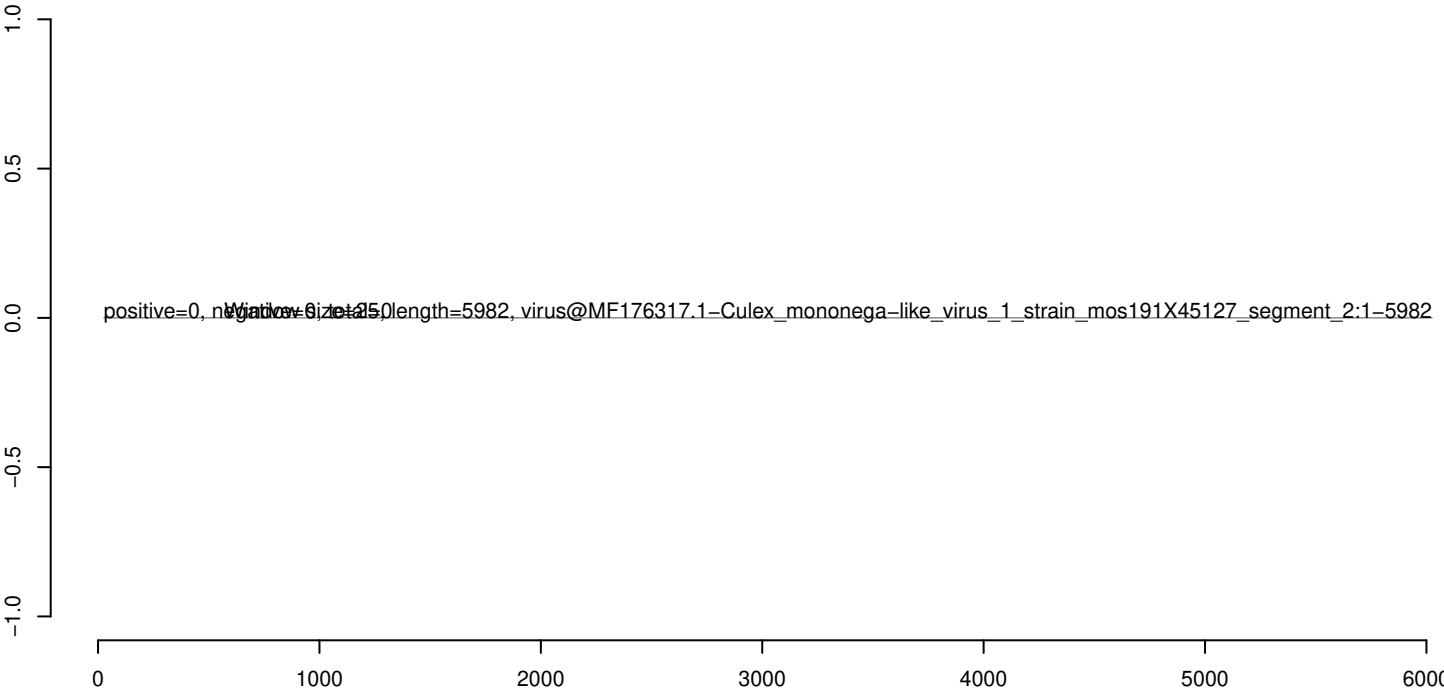
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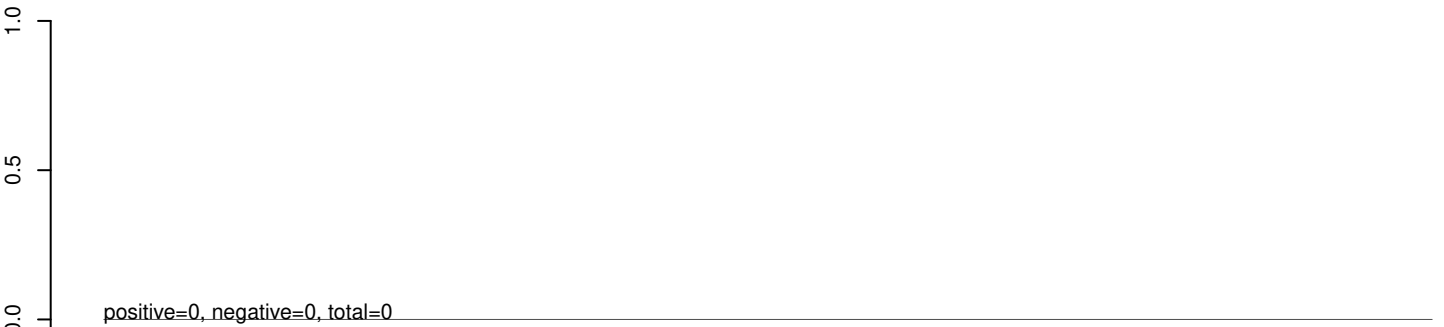
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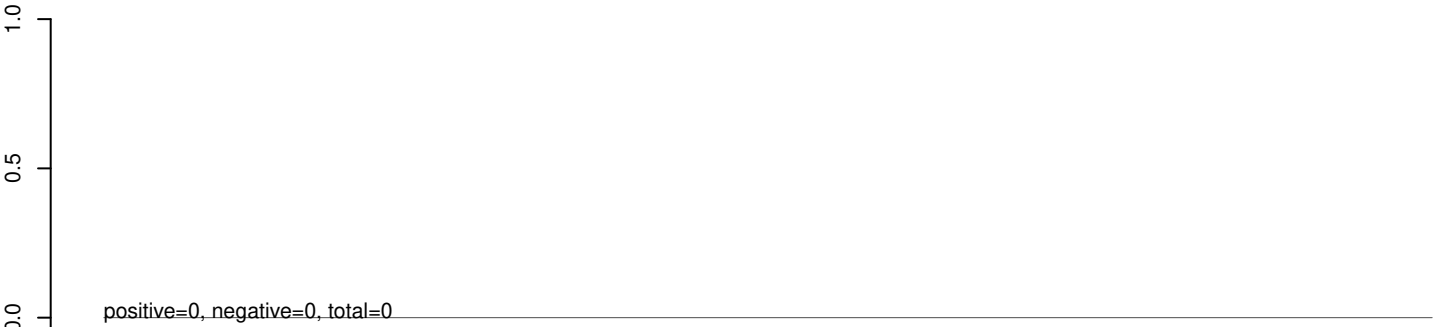
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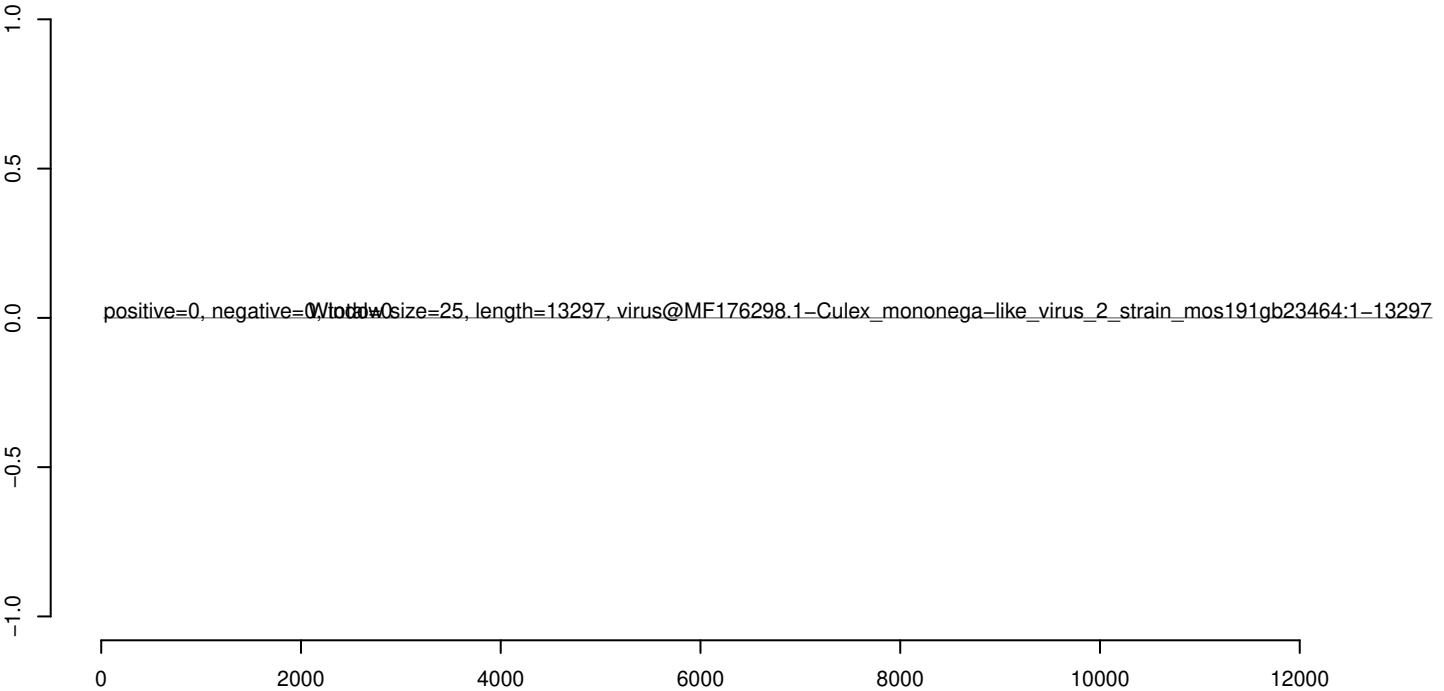
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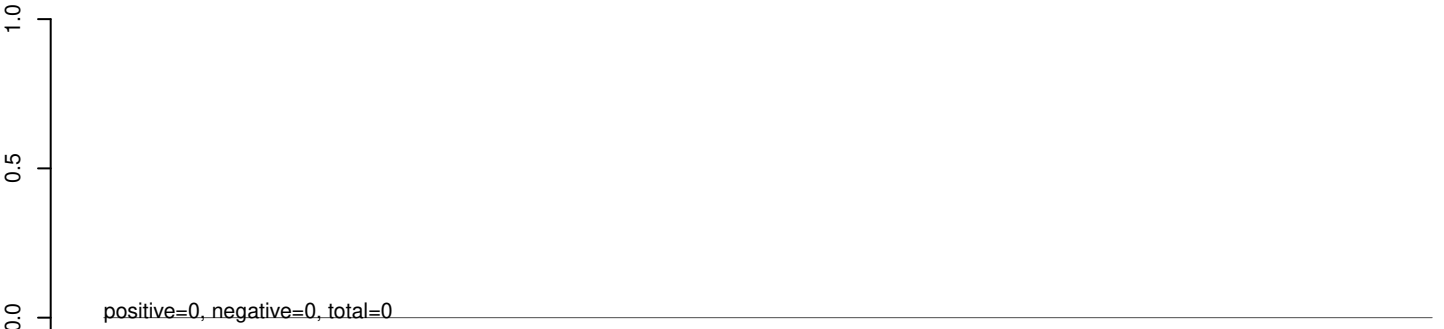
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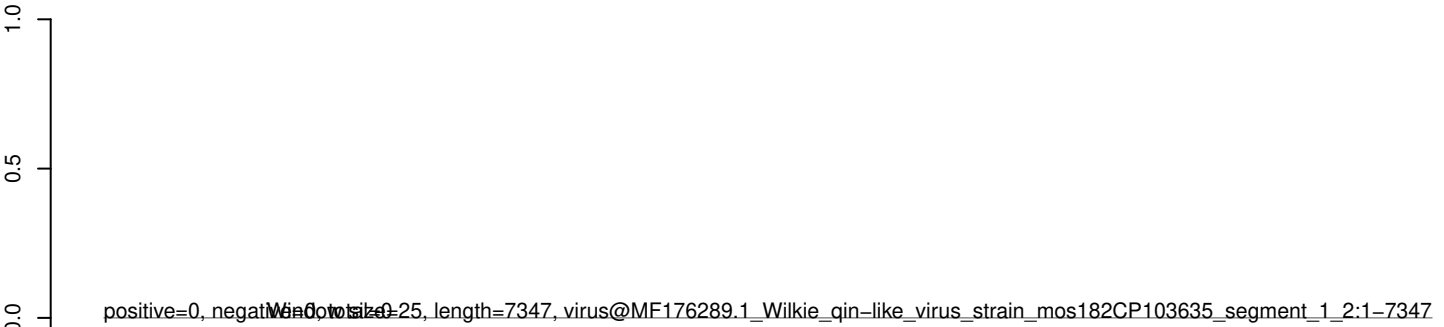
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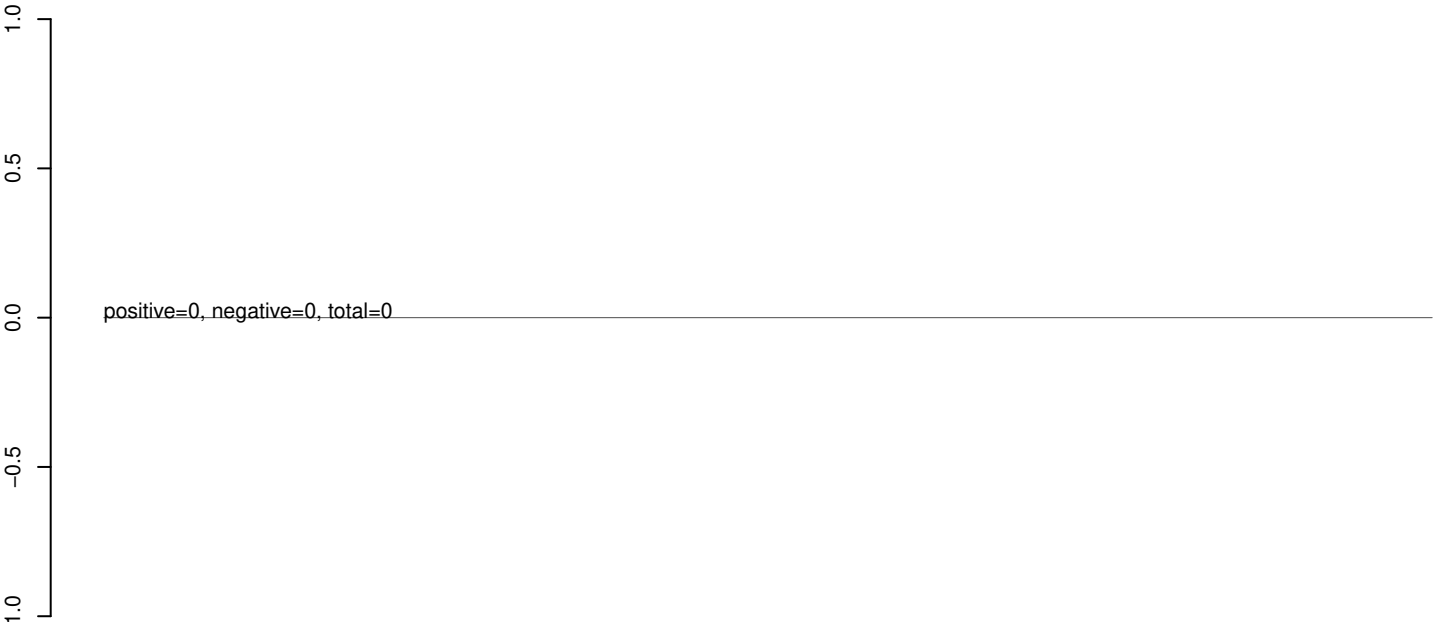


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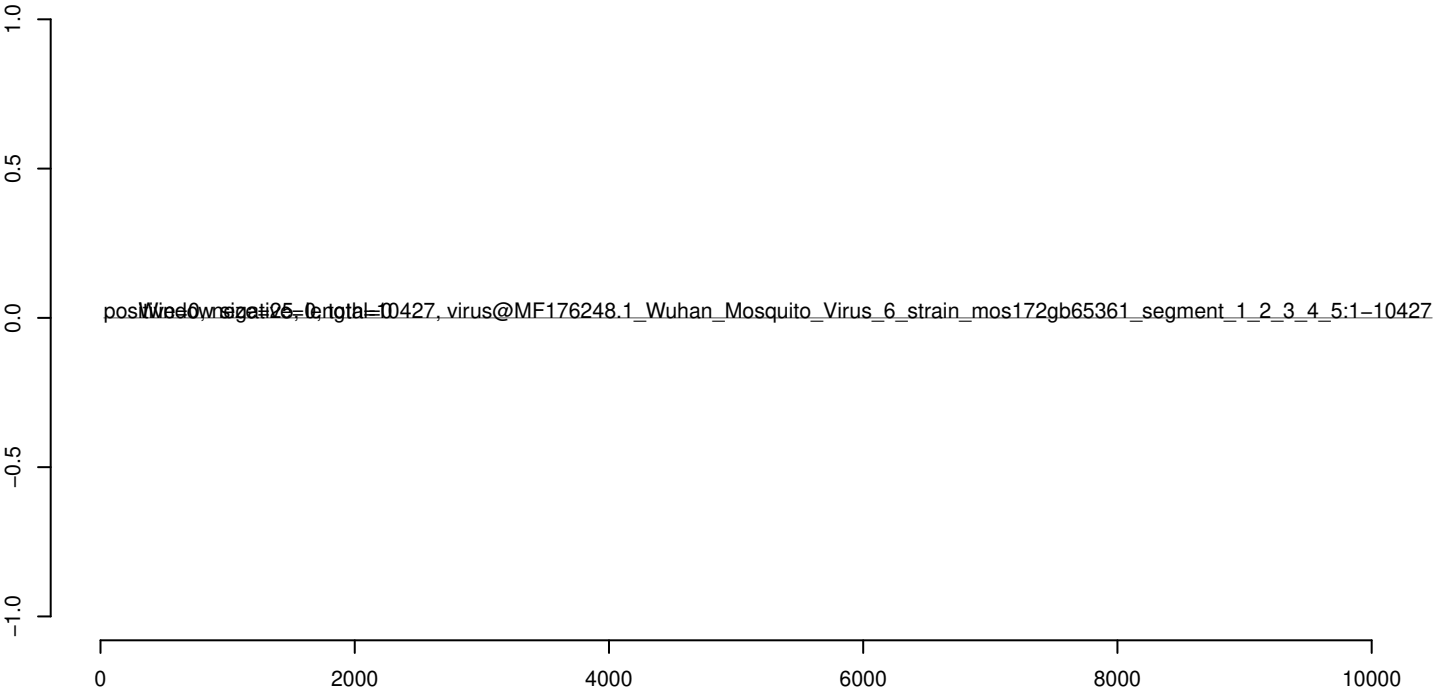
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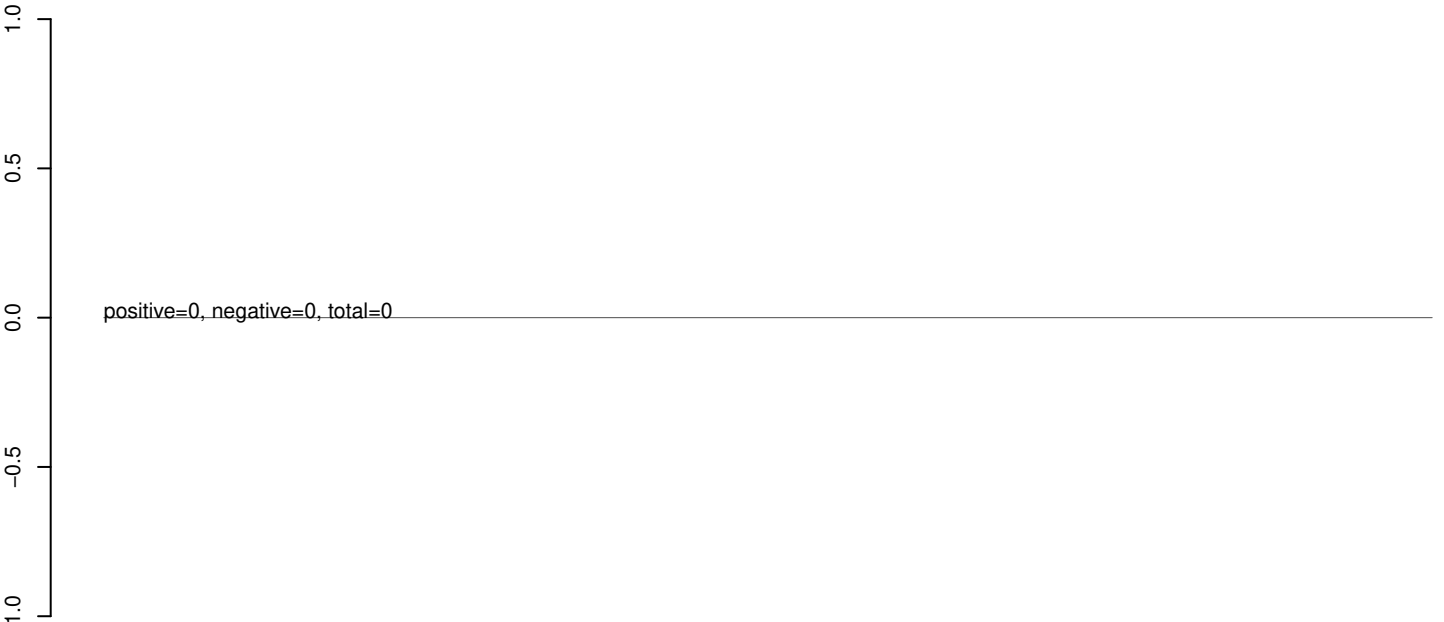
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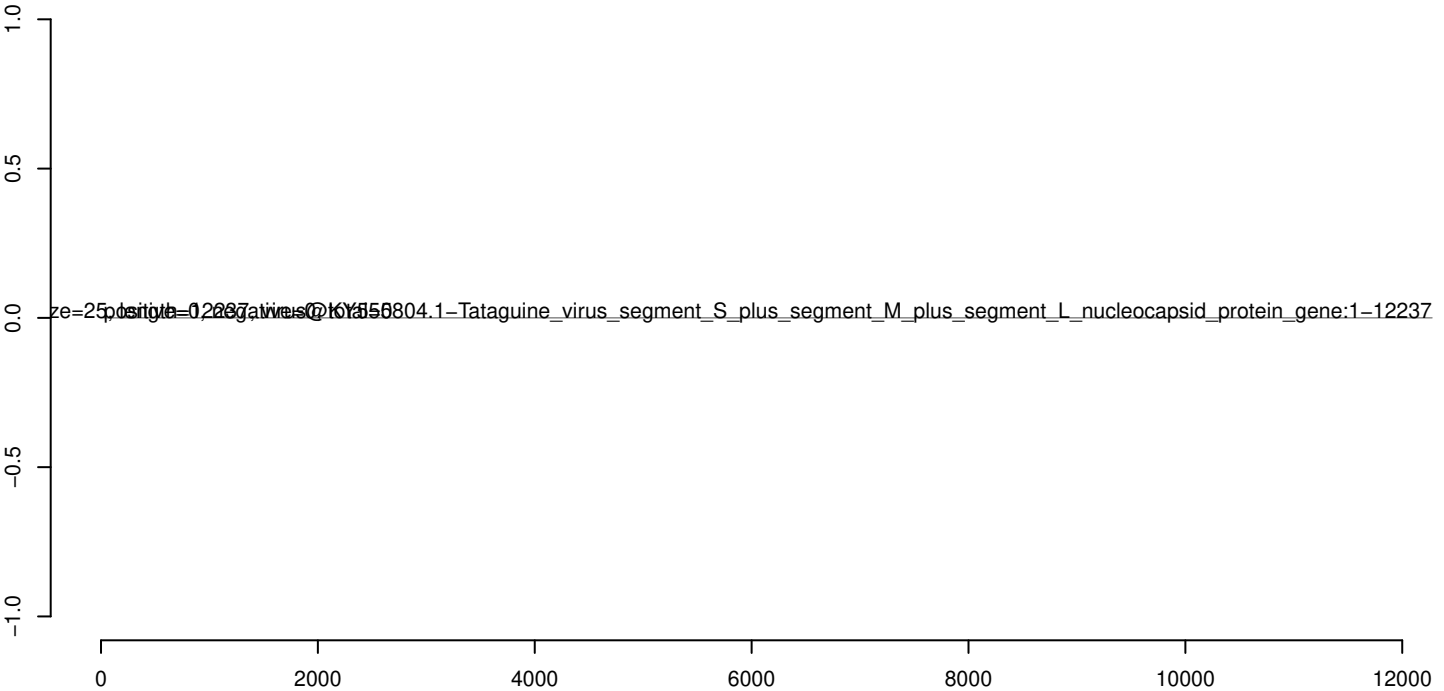
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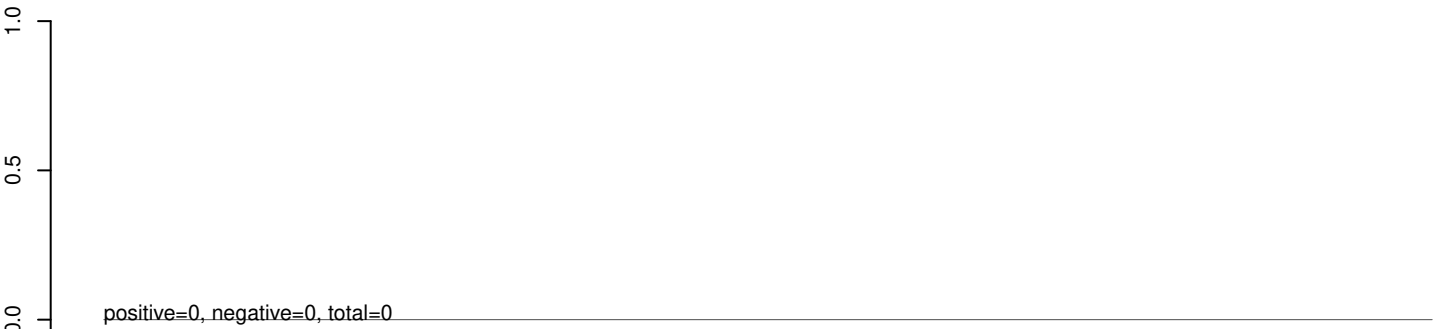
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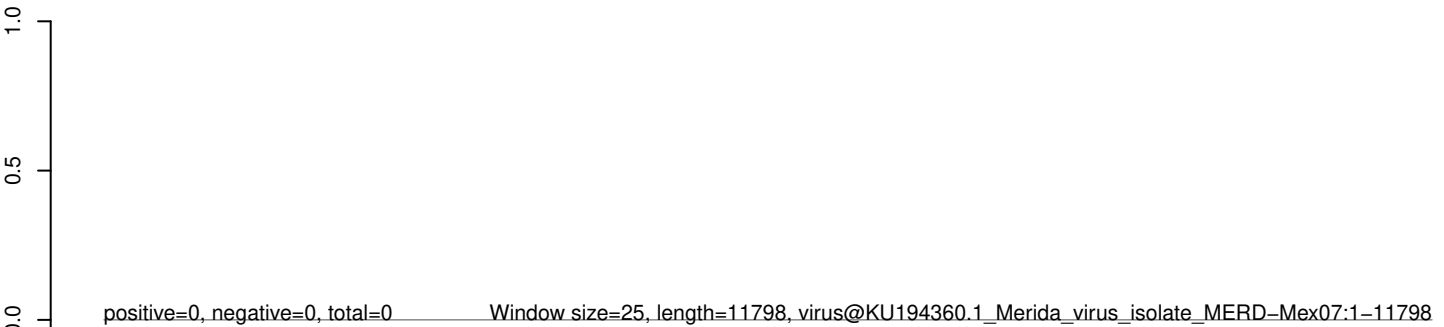
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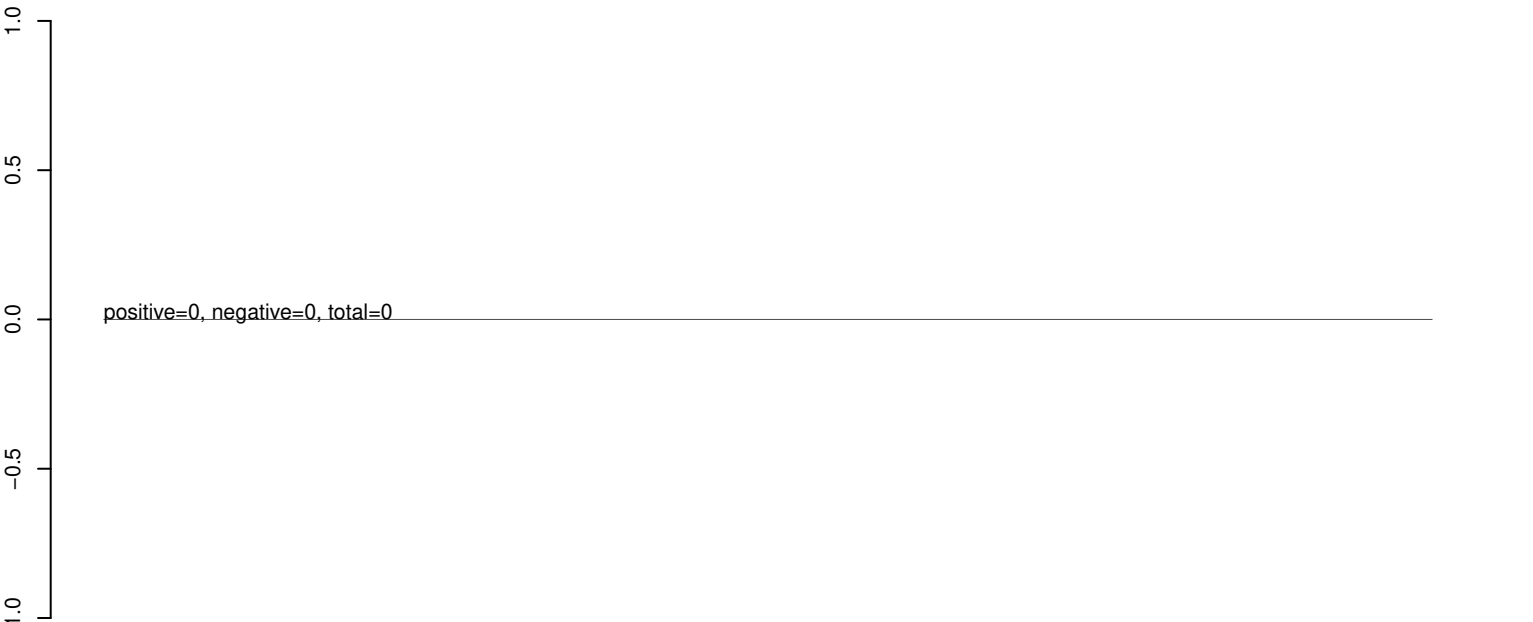


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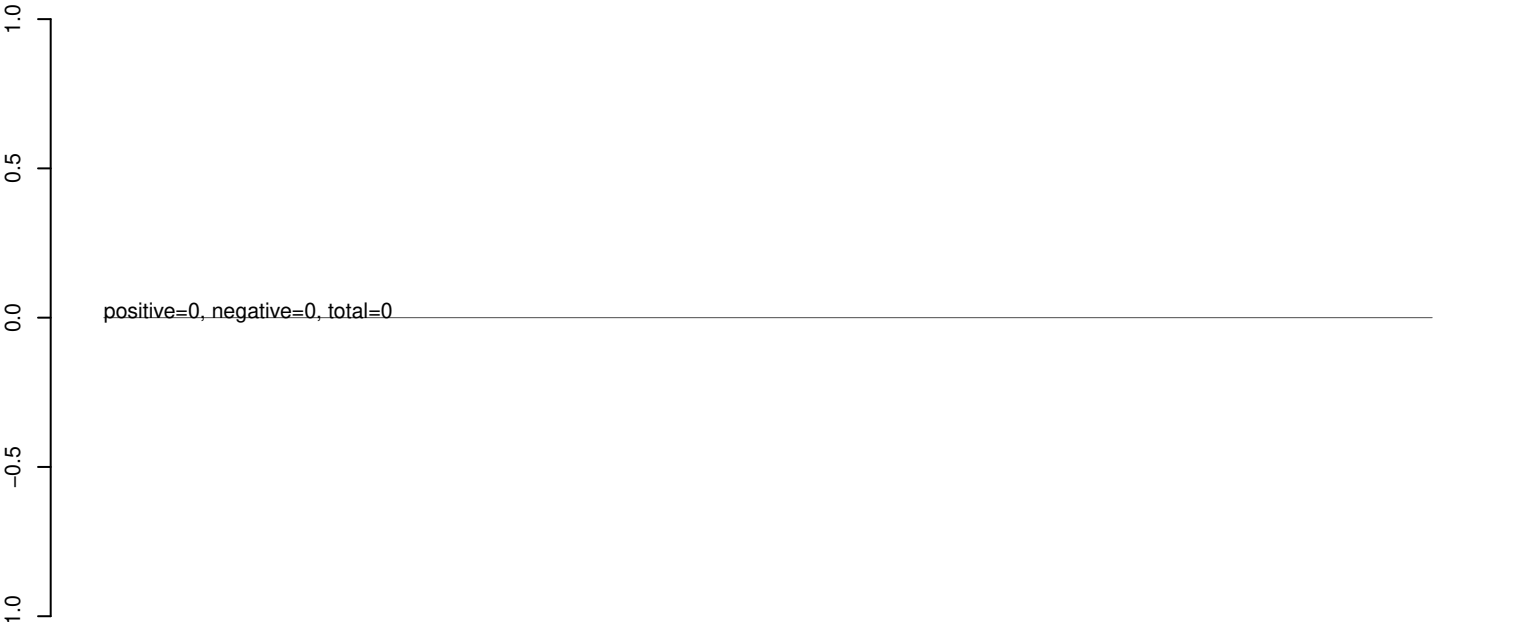


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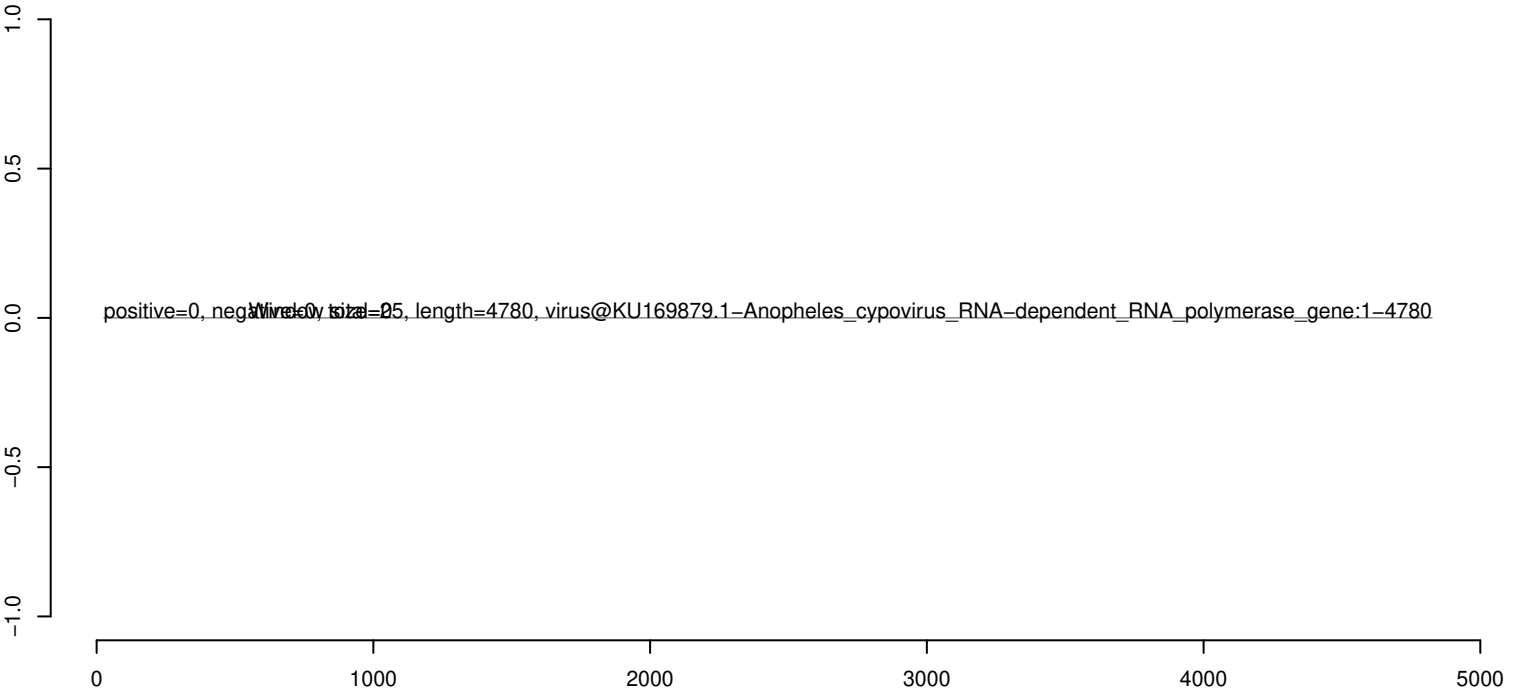
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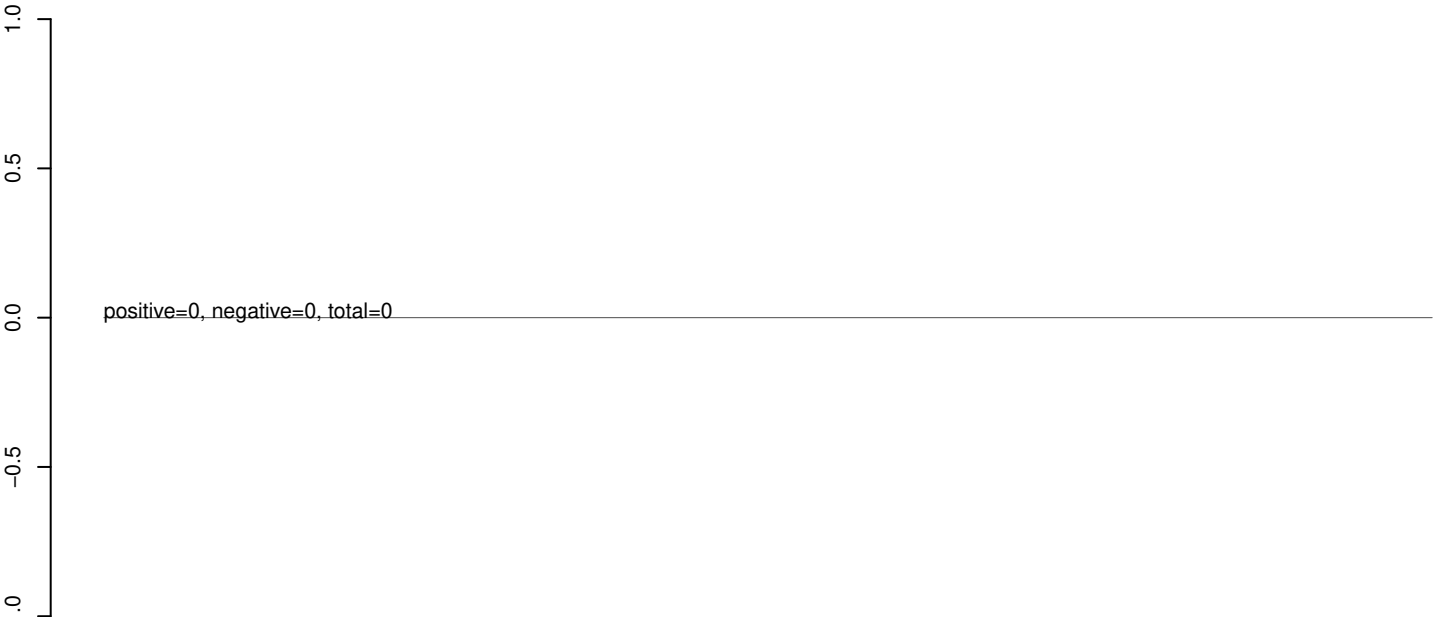
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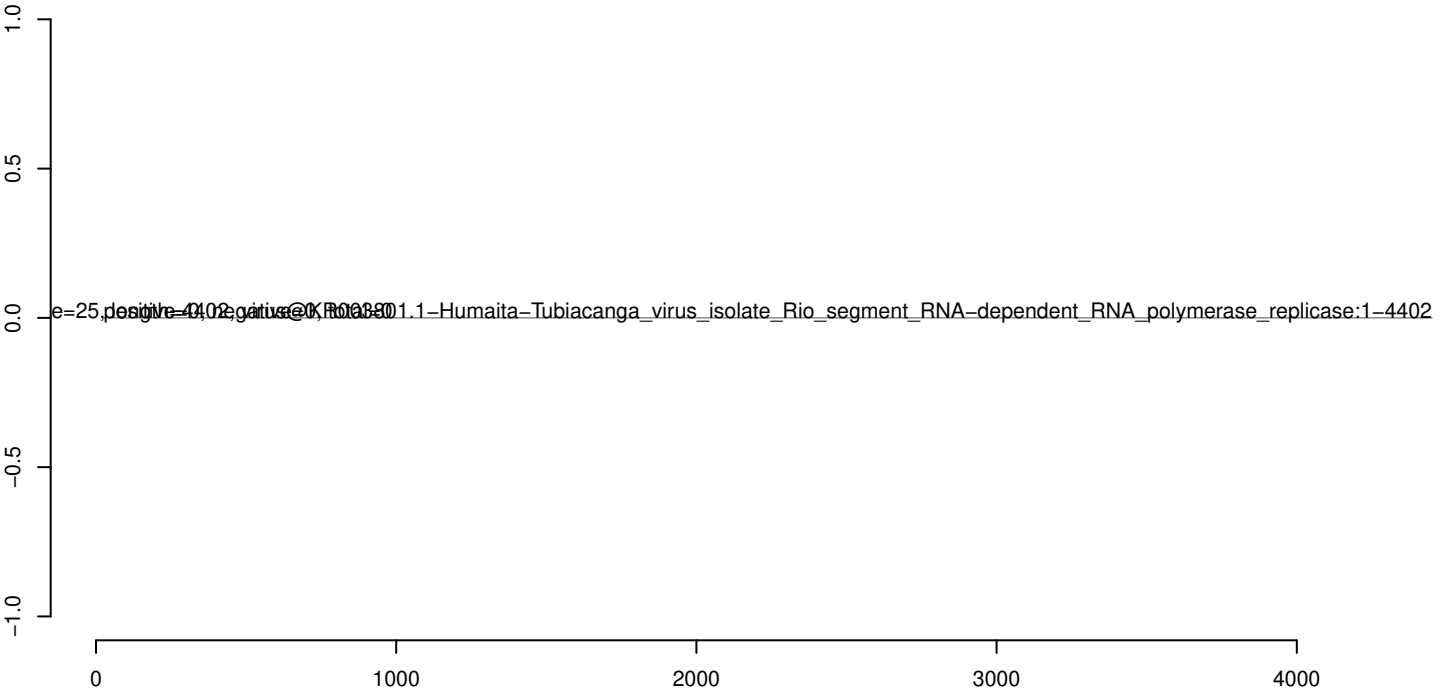
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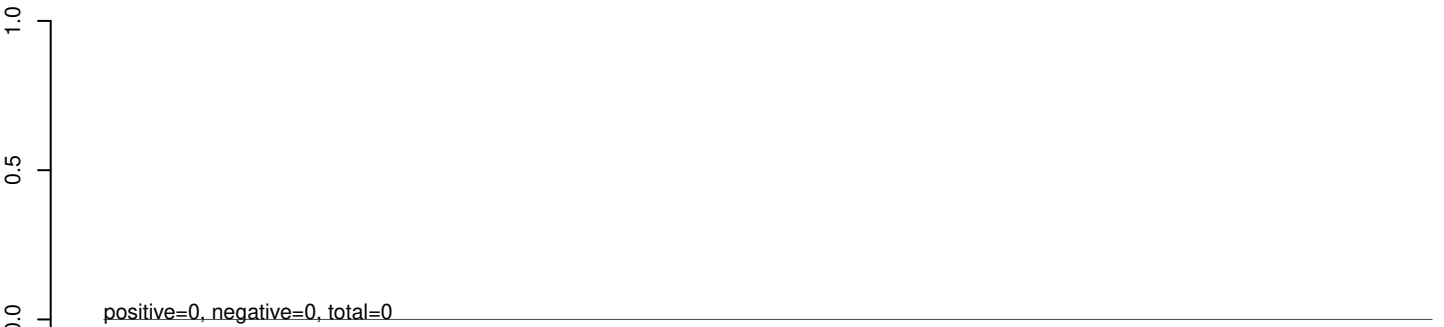
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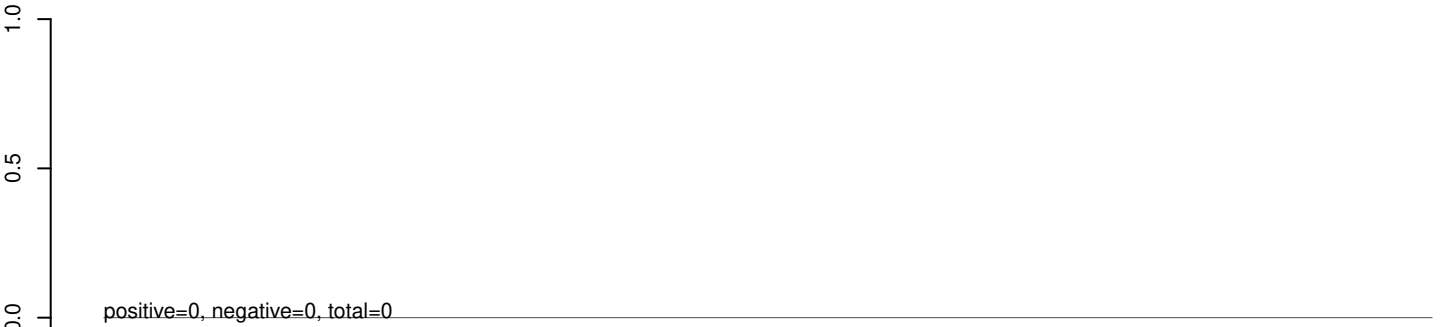
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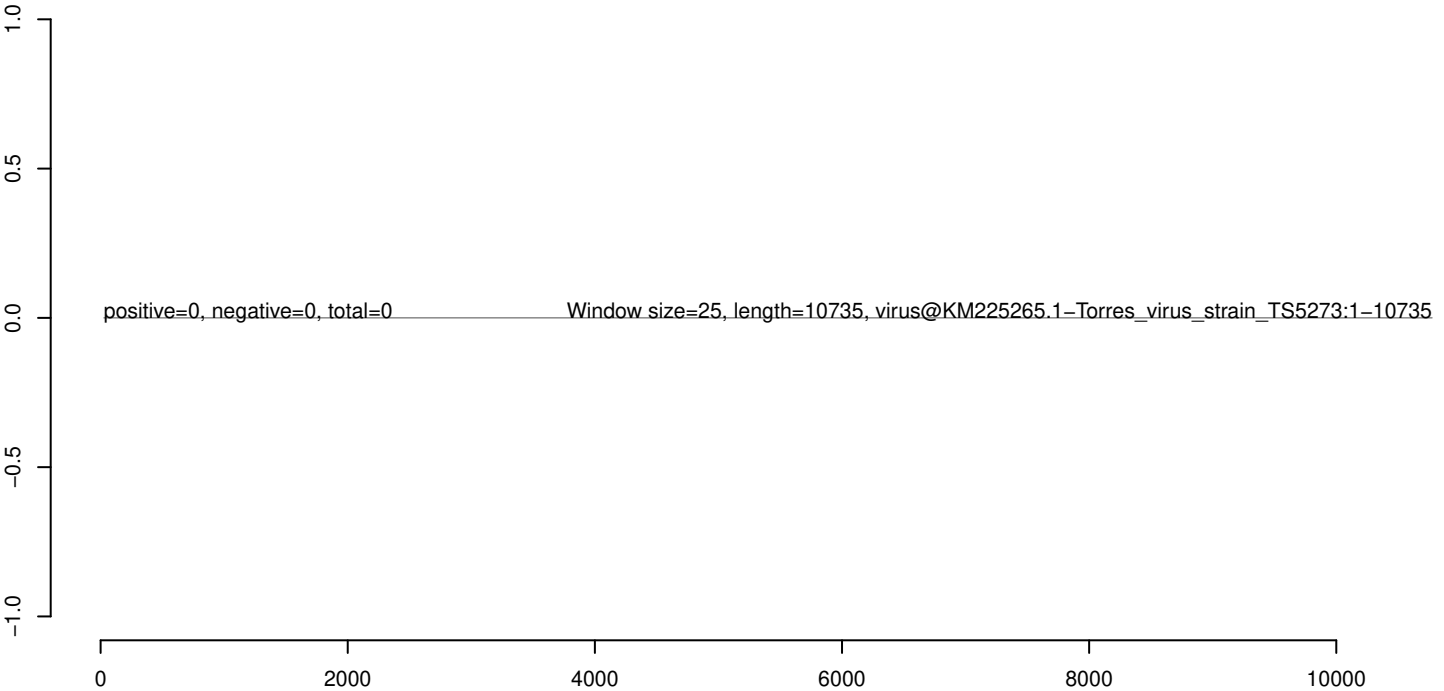
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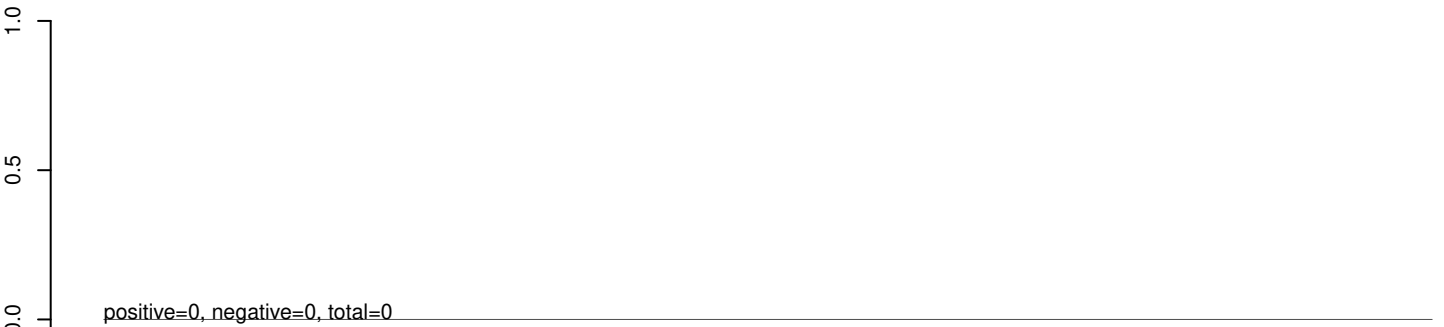
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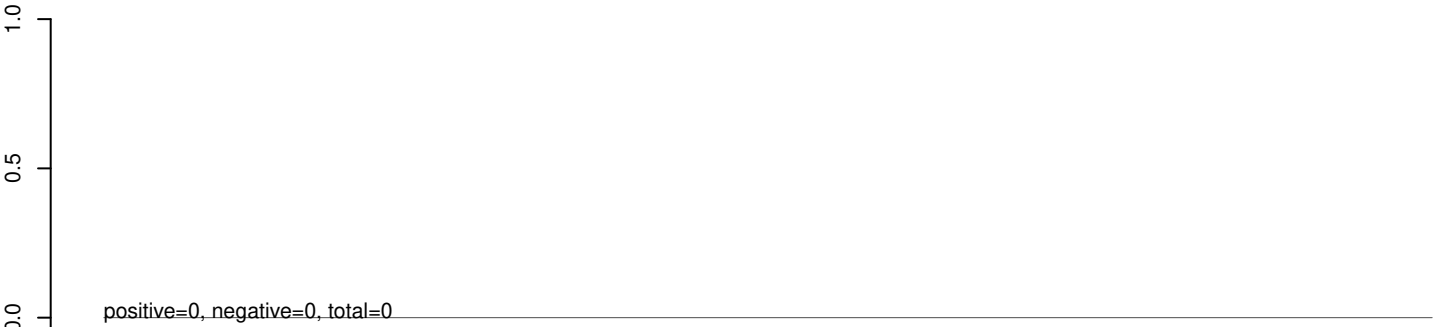
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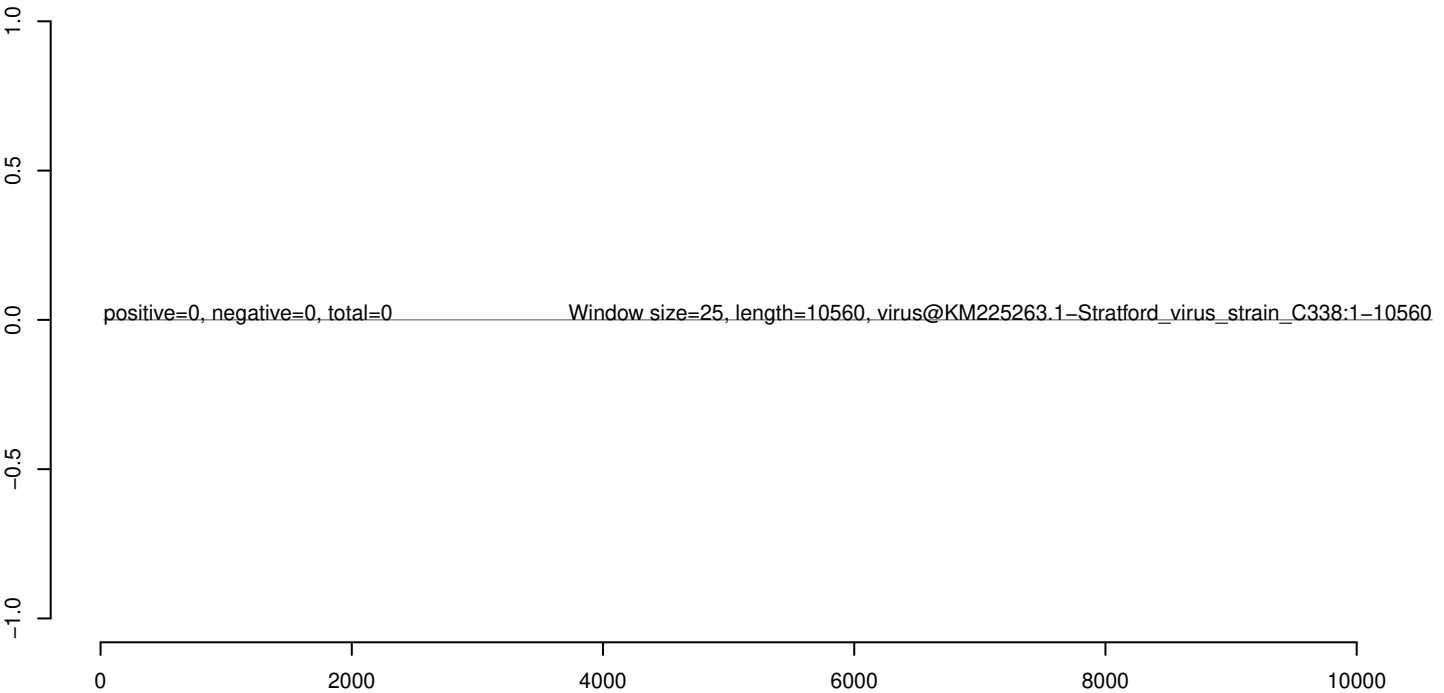
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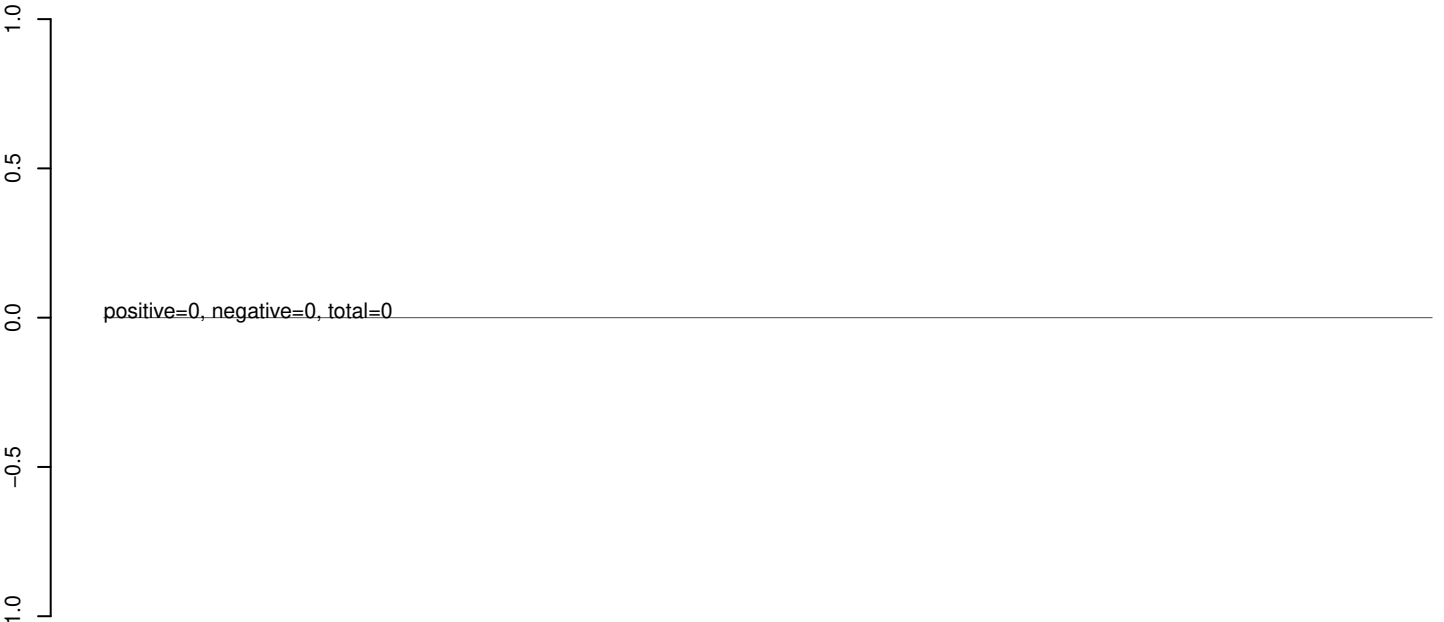
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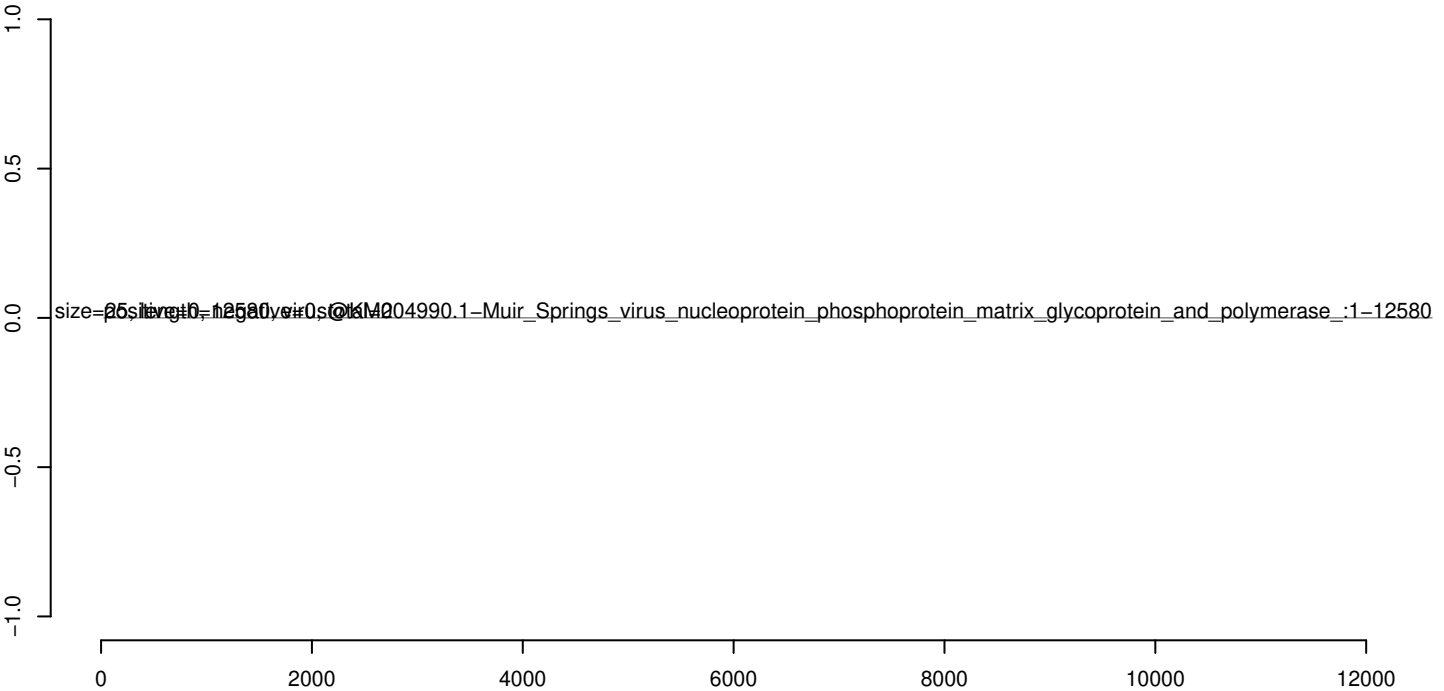
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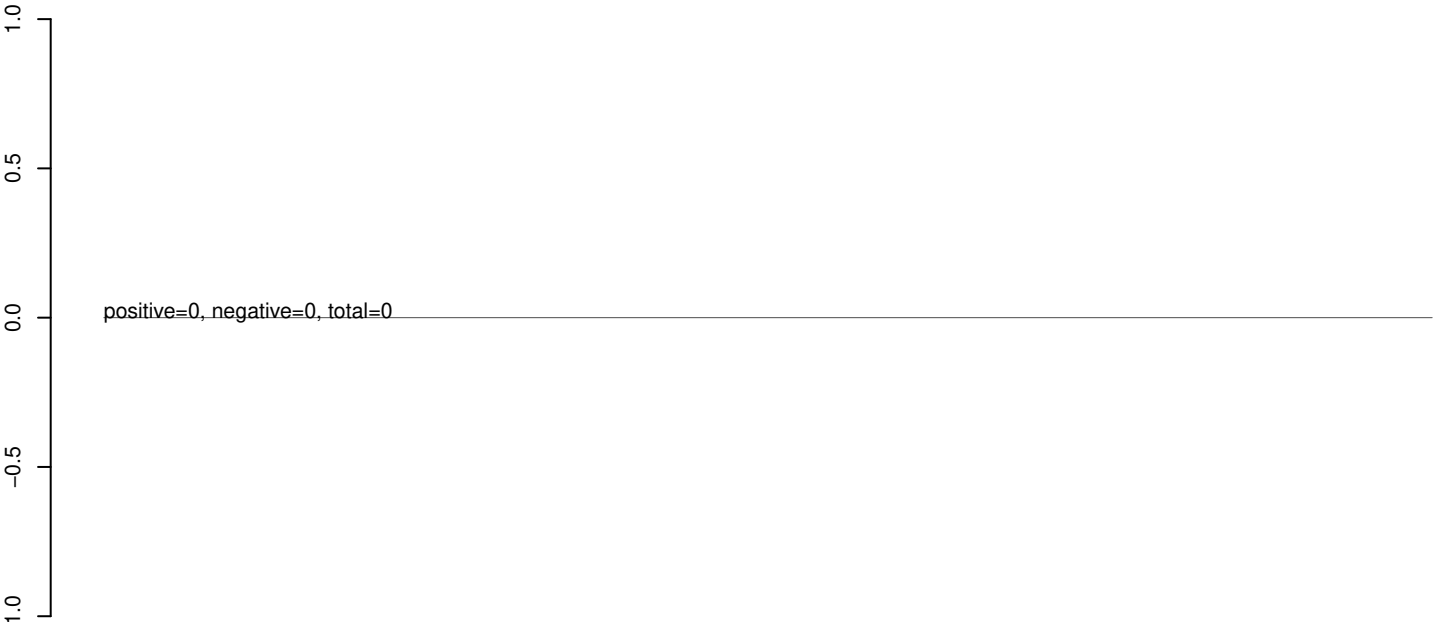
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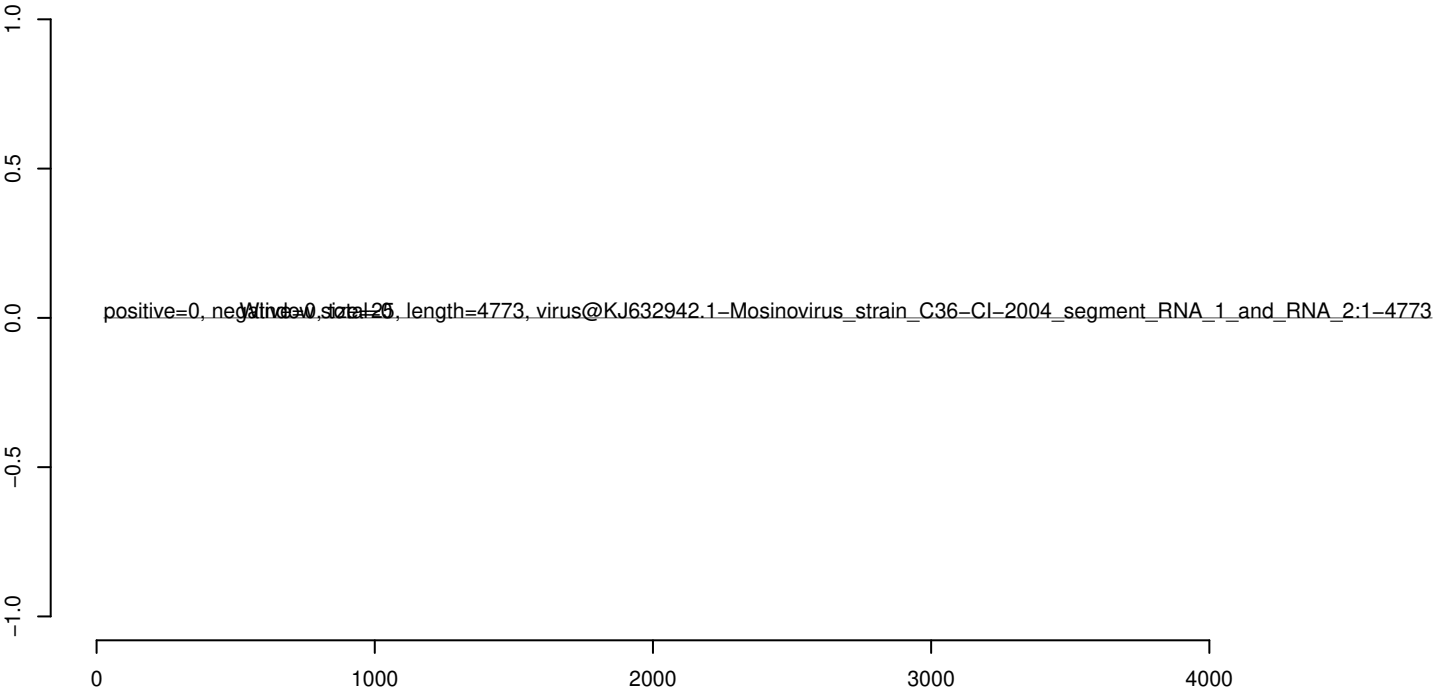
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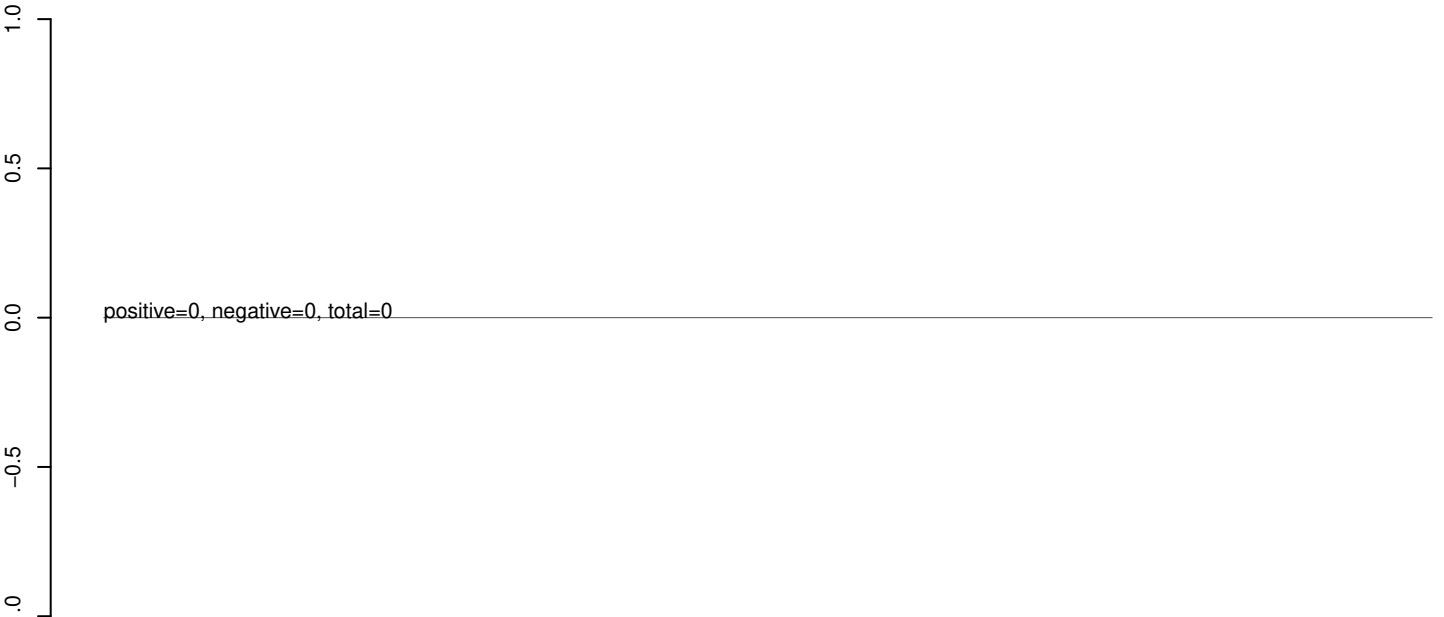
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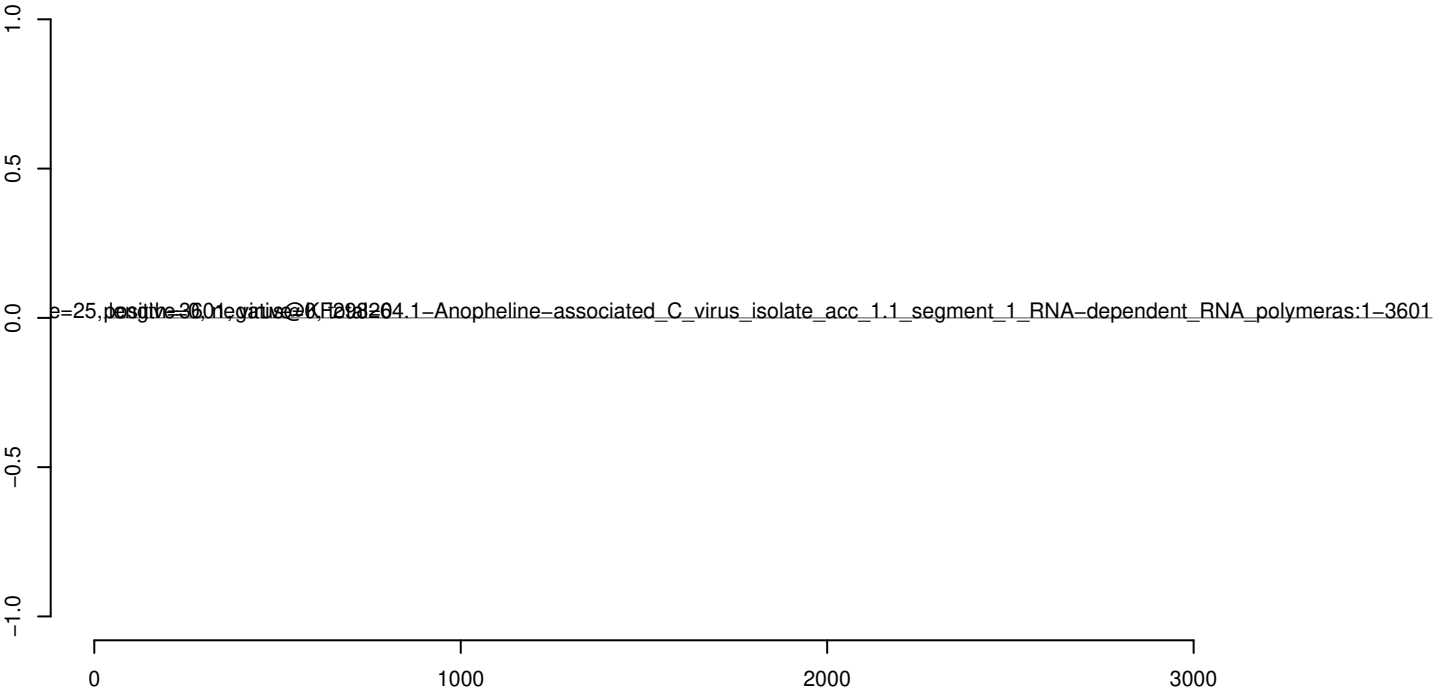
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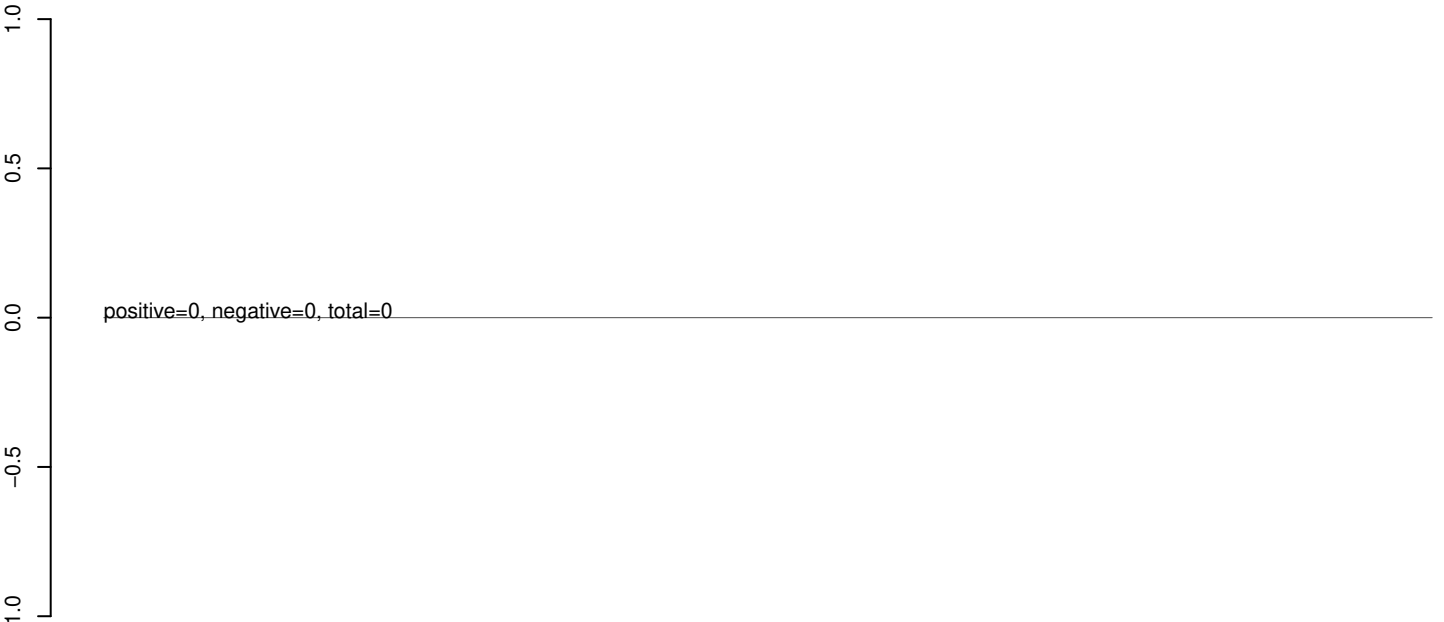
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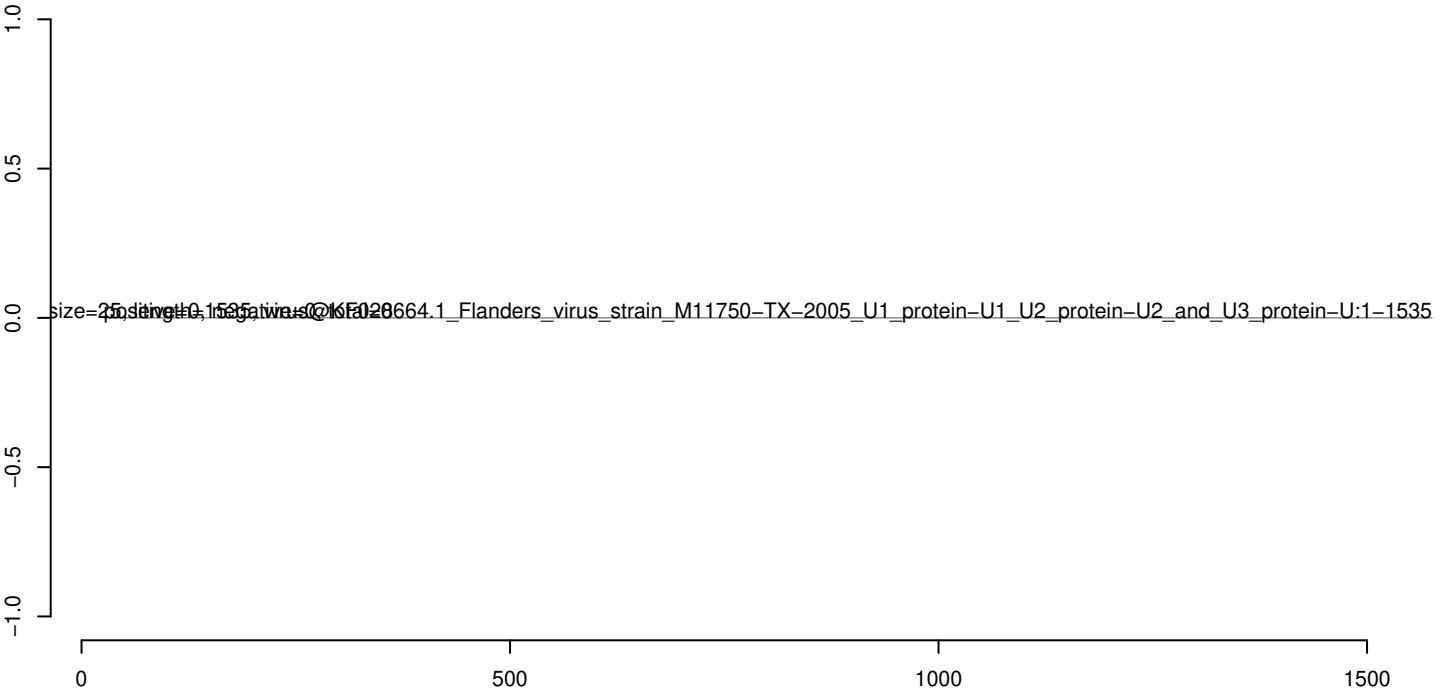
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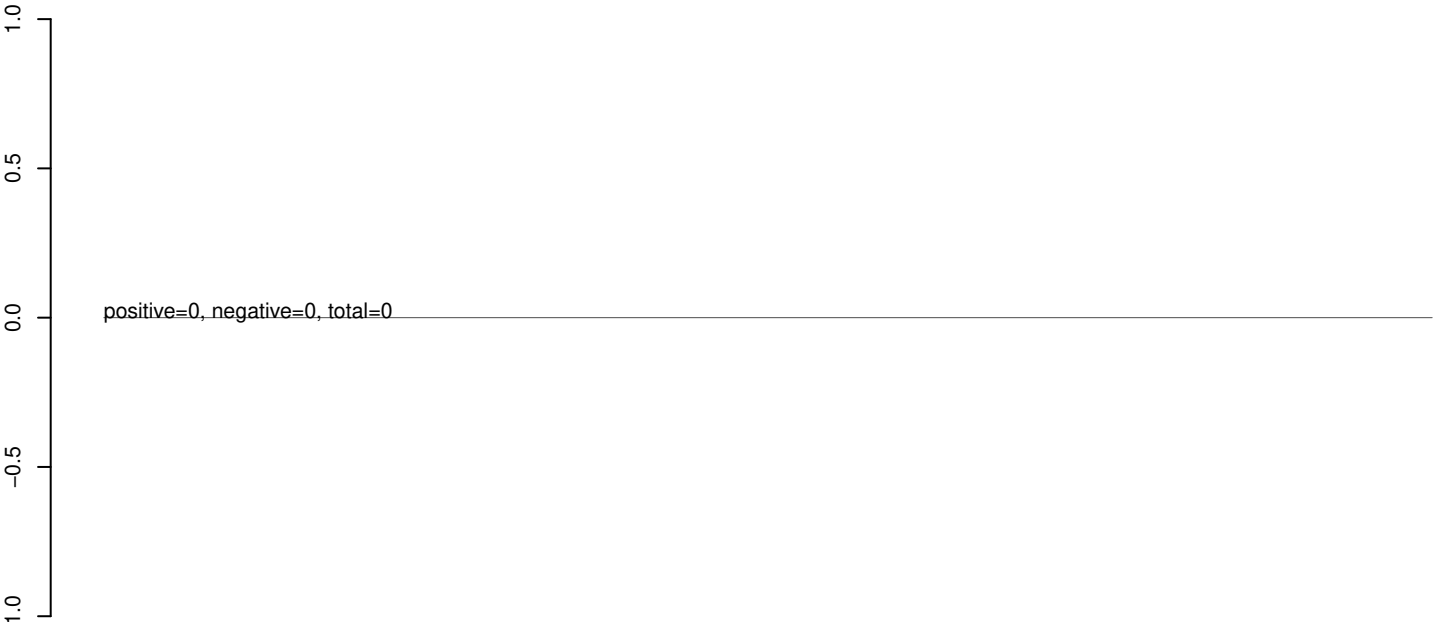
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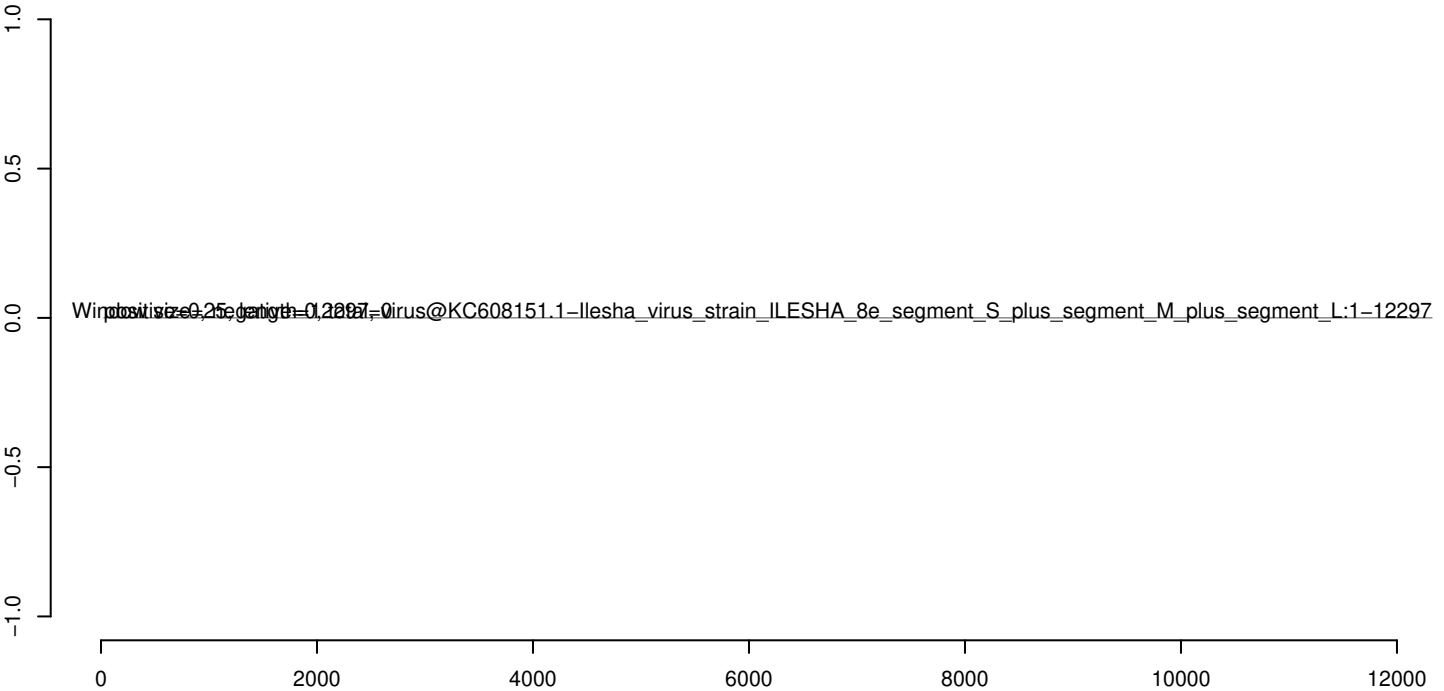
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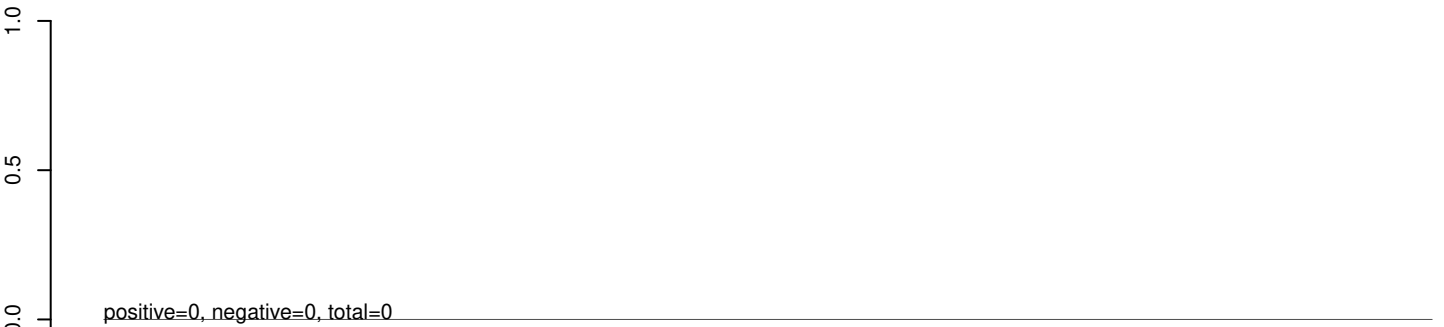
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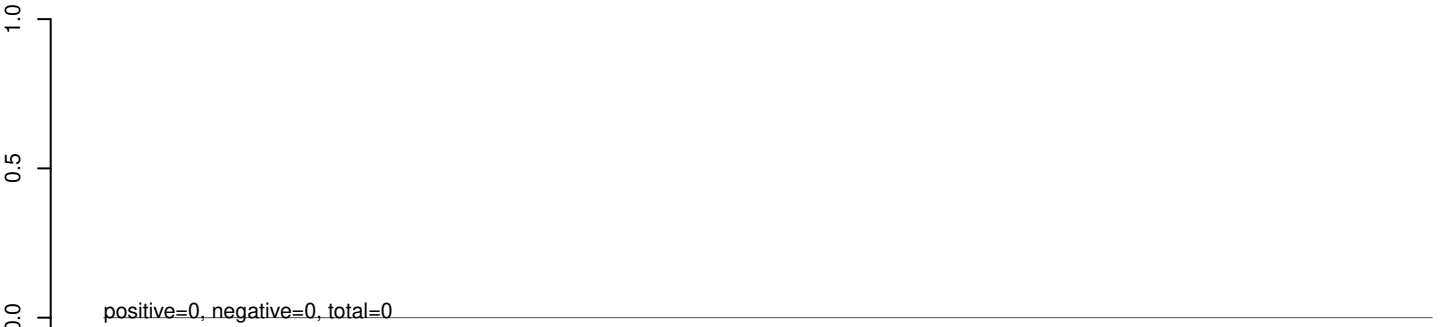
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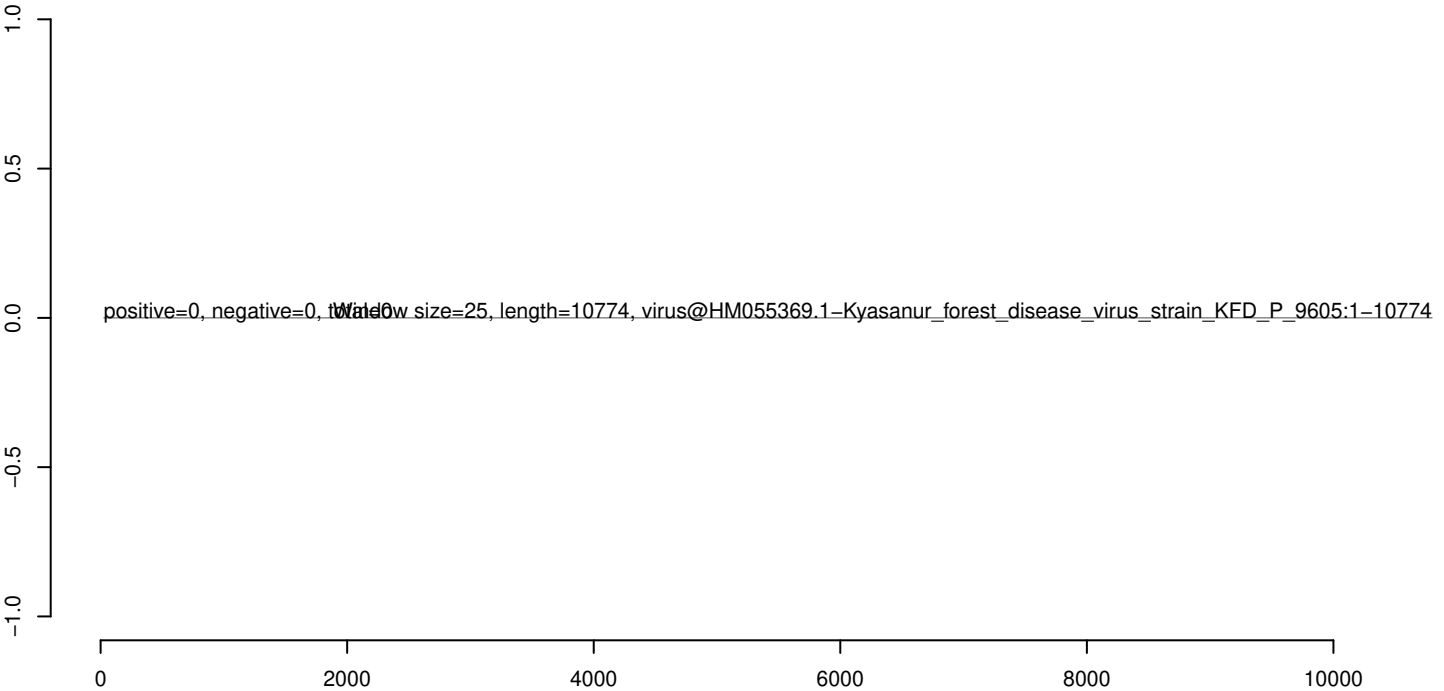
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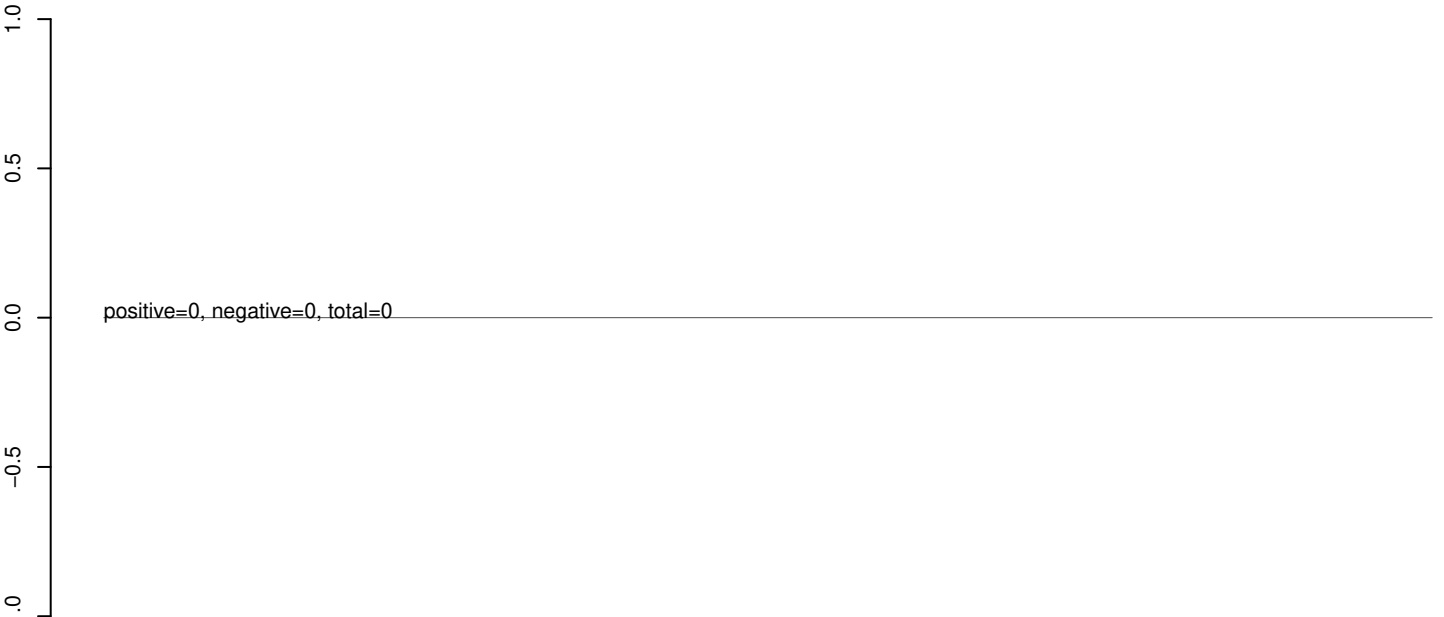
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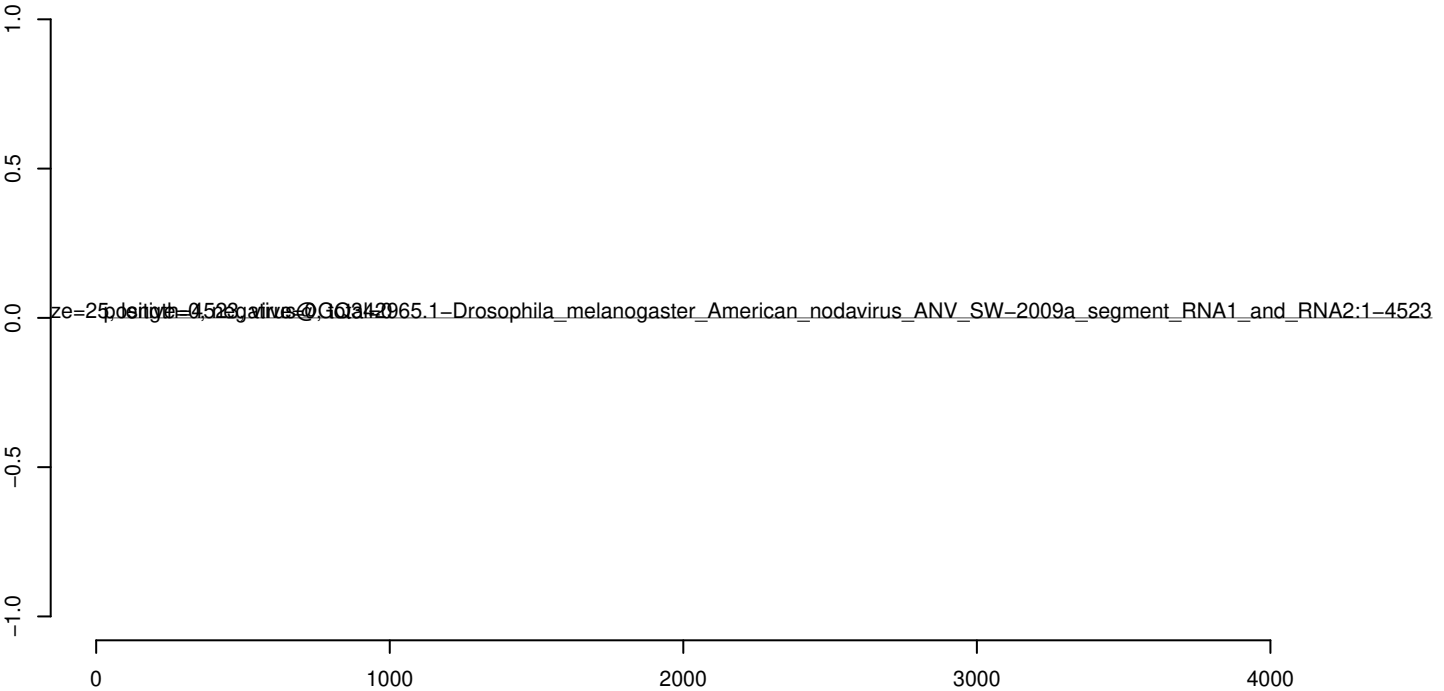
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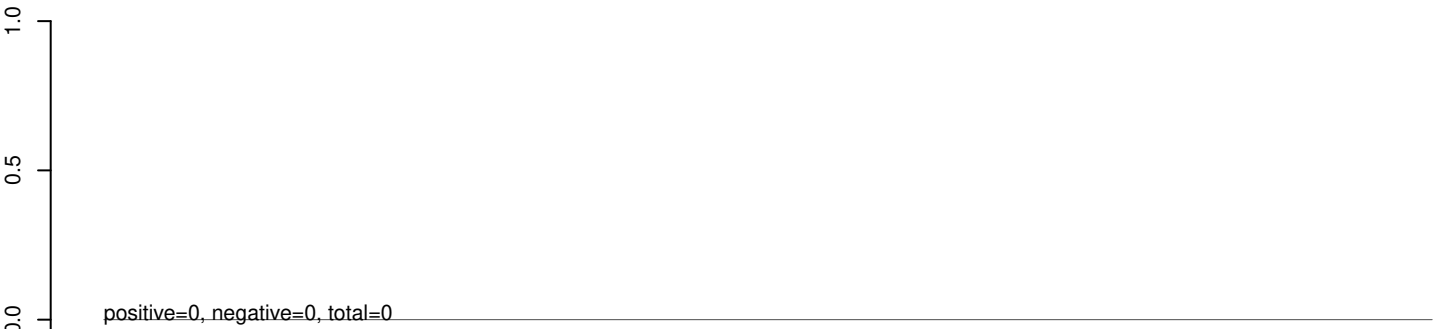
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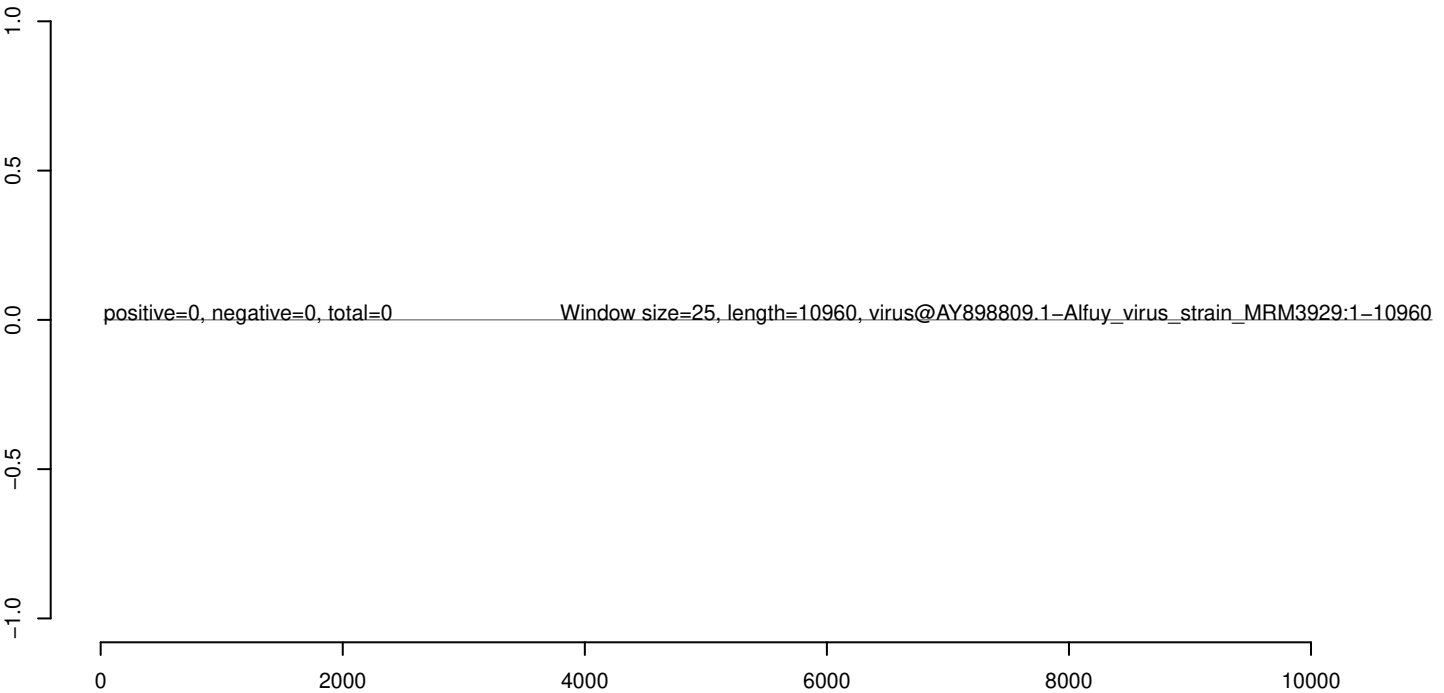
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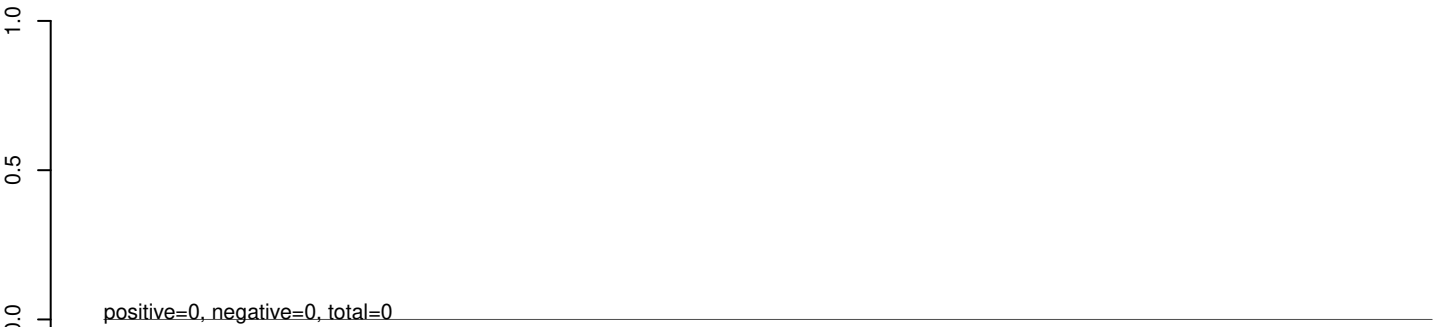
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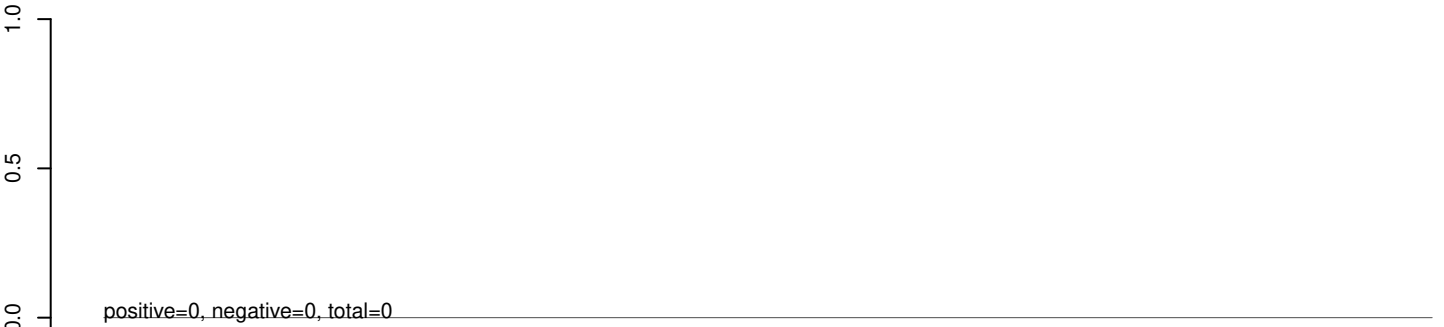
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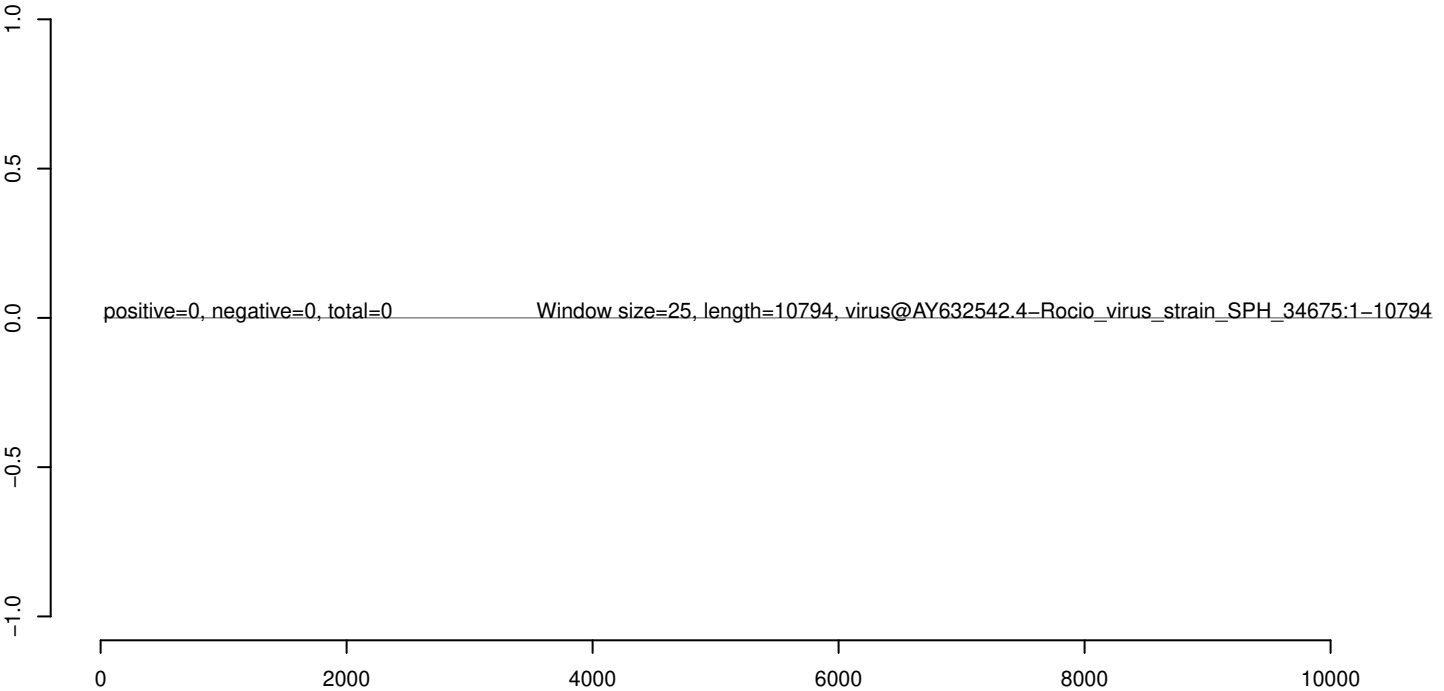
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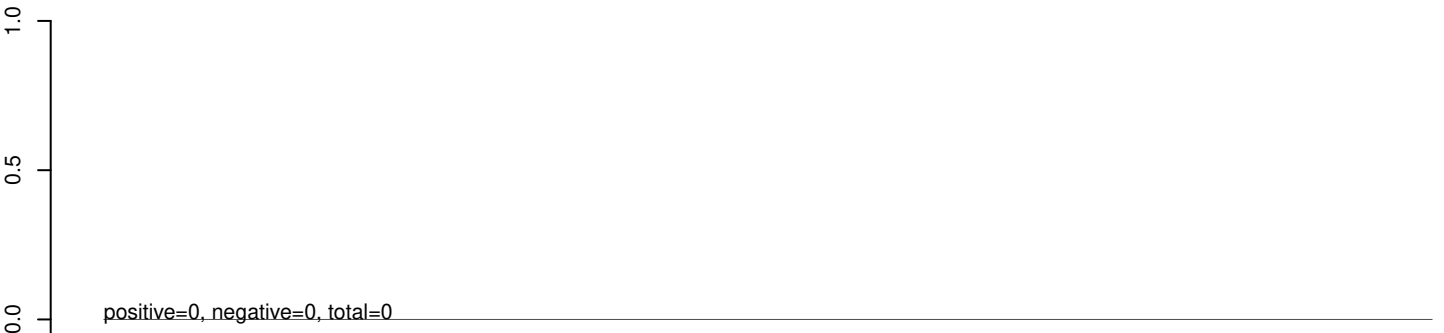
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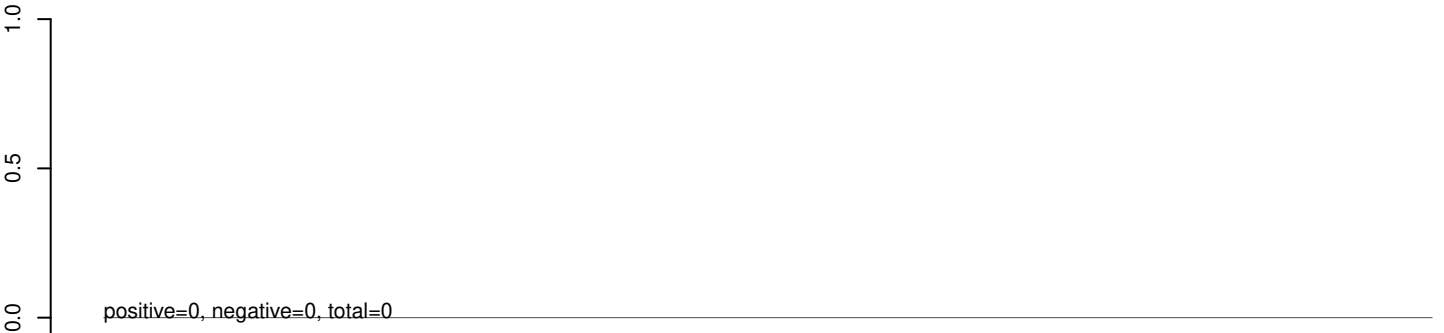
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AeAeg_Aag2_mock_dsFluc_RA.rep

