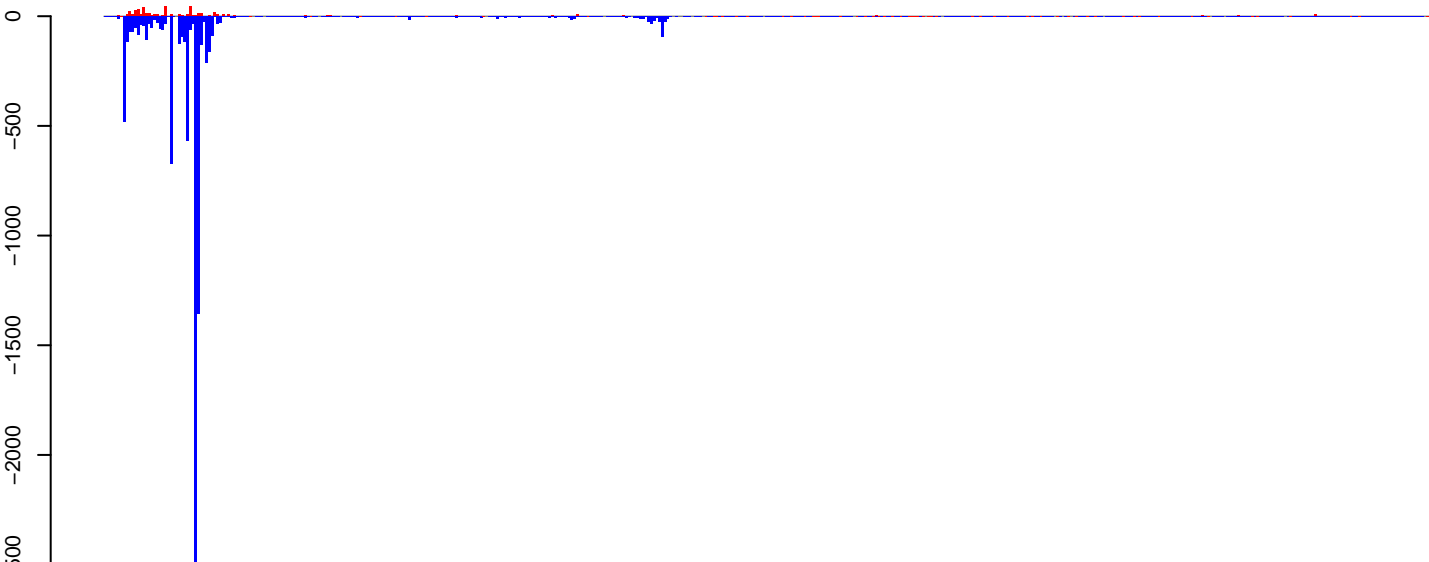


AeAeg_Aag2_cells.18_23.rep



positive=418, negative=2513, total=2930

AeAeg_Aag2_cells.24_35.rep



positive=590, negative=8378, total=8968

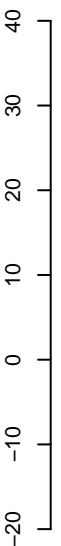
AeAeg_Aag2_cells.rep



25, length=12099, vireo@MC=108263, total=11898 Charoen-like_virus_isolate_Rio_segment_S_plus_segment_M_plus_segment_L_Nucle:1-12099

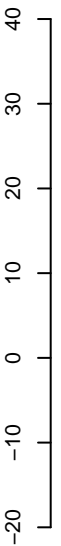
0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_cells.18_23.rep



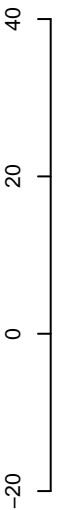
positive=451, negative=302, total=753

AeAeg_Aag2_cells.24_35.rep



positive=104, negative=70, total=173

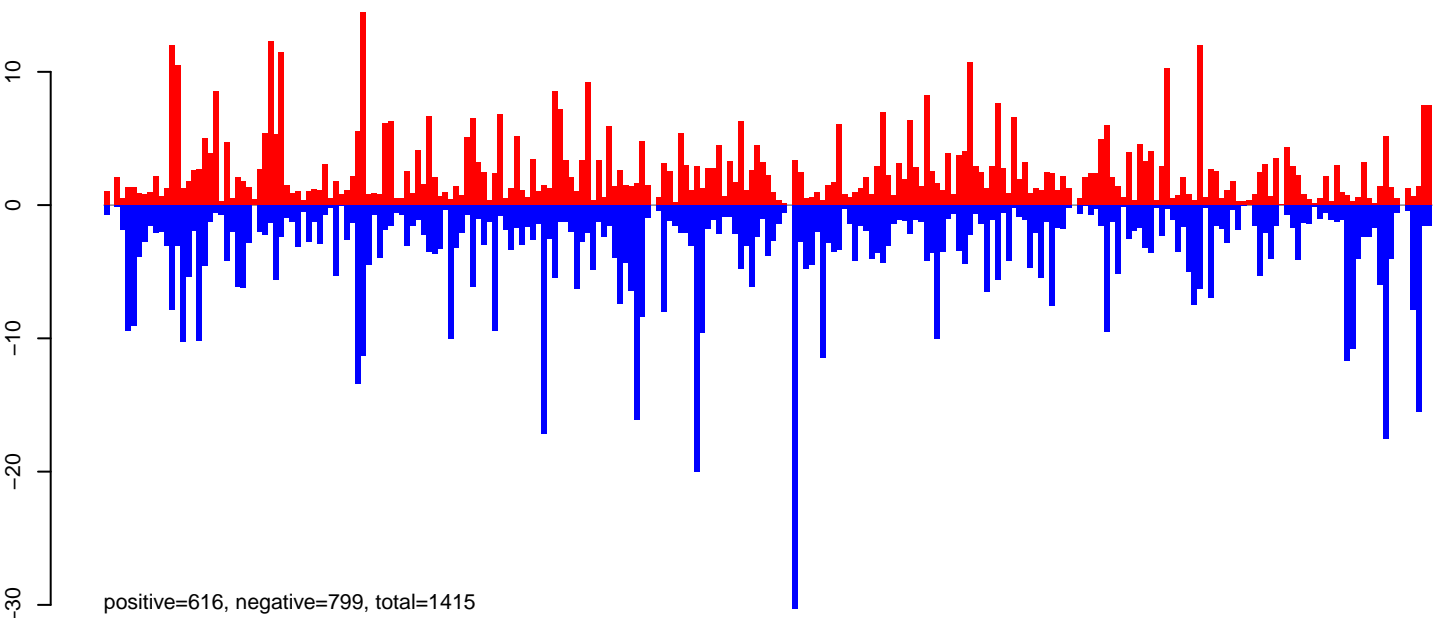
AeAeg_Aag2_cells.rep



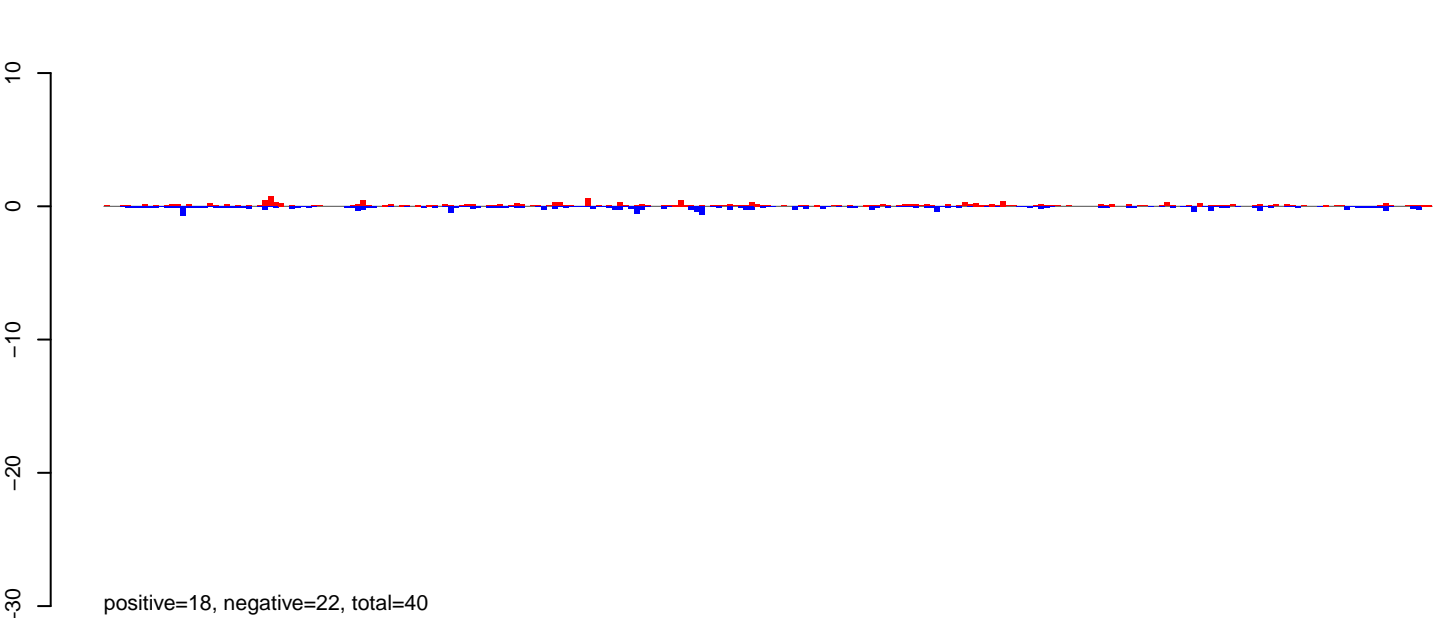
positive=555, negative=372, total=927, window size=25, length=10682, virus@NC_001564.2-Cell_fusing_agent_virus_strain_Galveston:1-10682



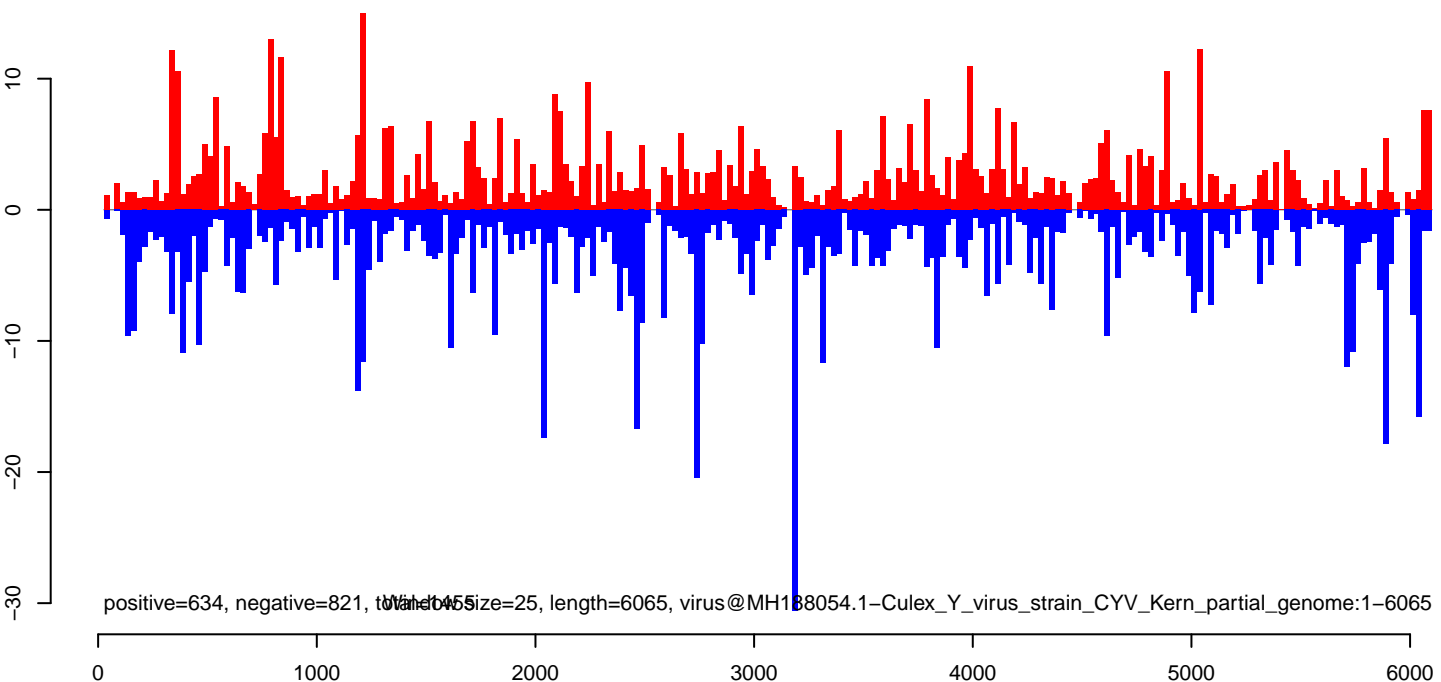
AeAeg_Aag2_cells.18_23.rep



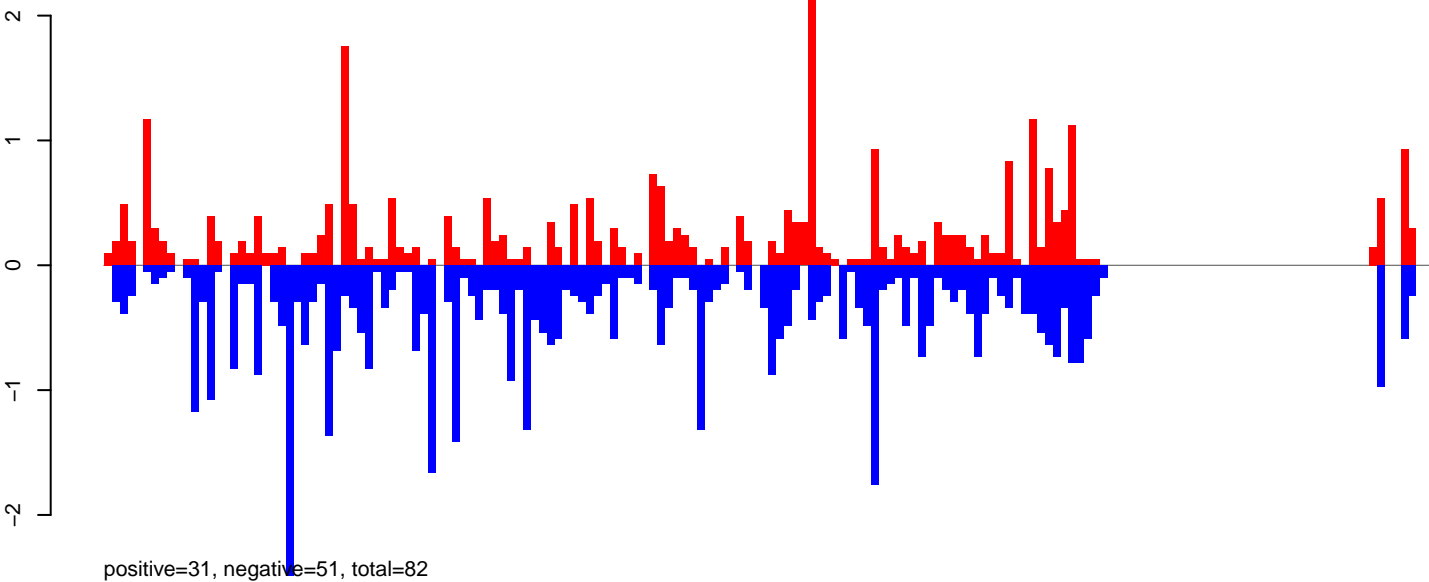
AeAeg_Aag2_cells.24_35.rep



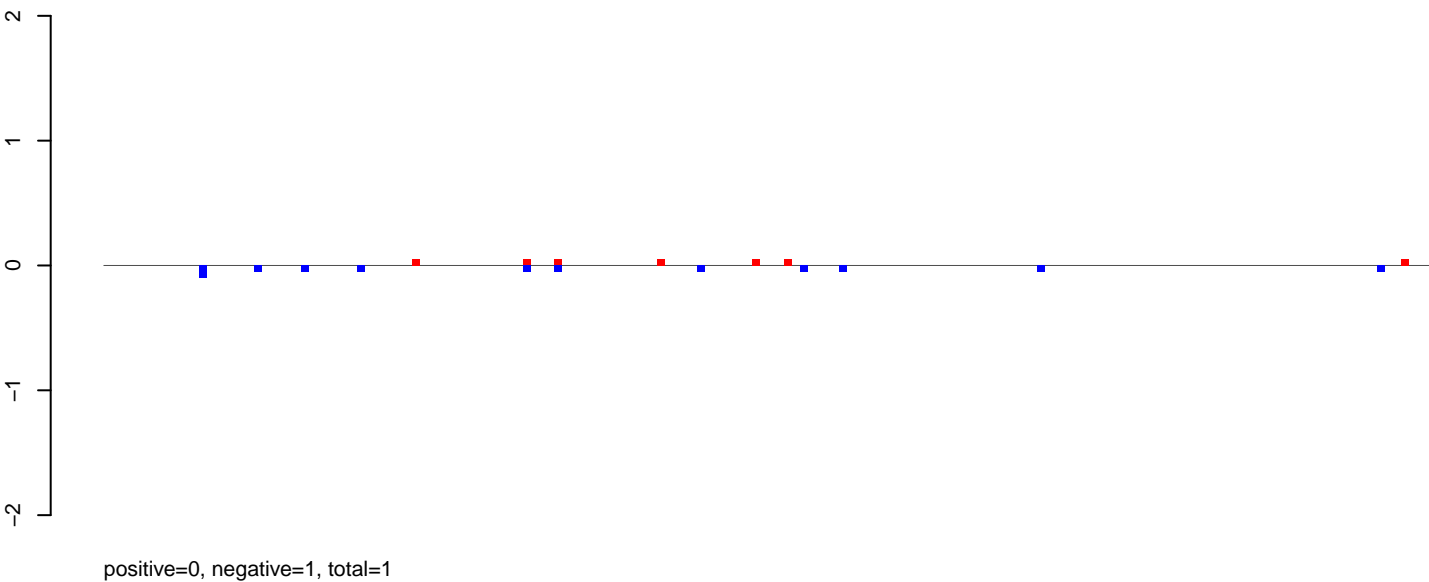
AeAeg_Aag2_cells.rep



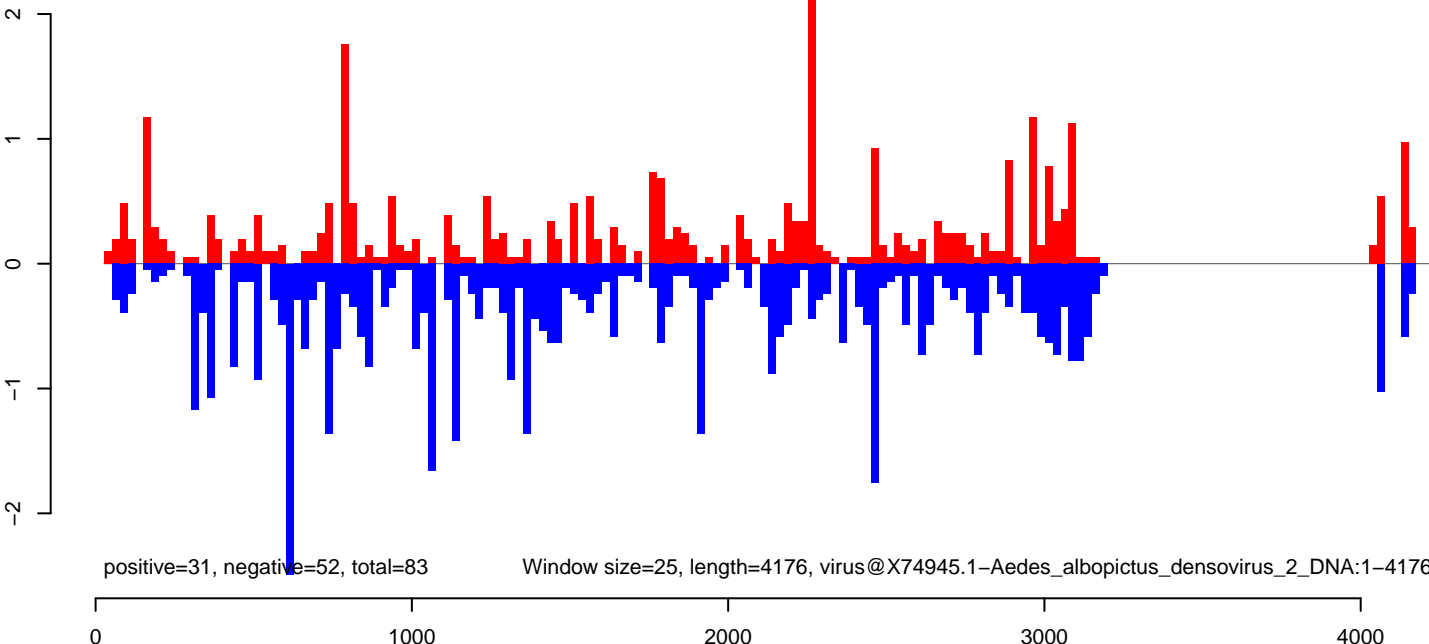
AeAeg_Aag2_cells.18_23.rep



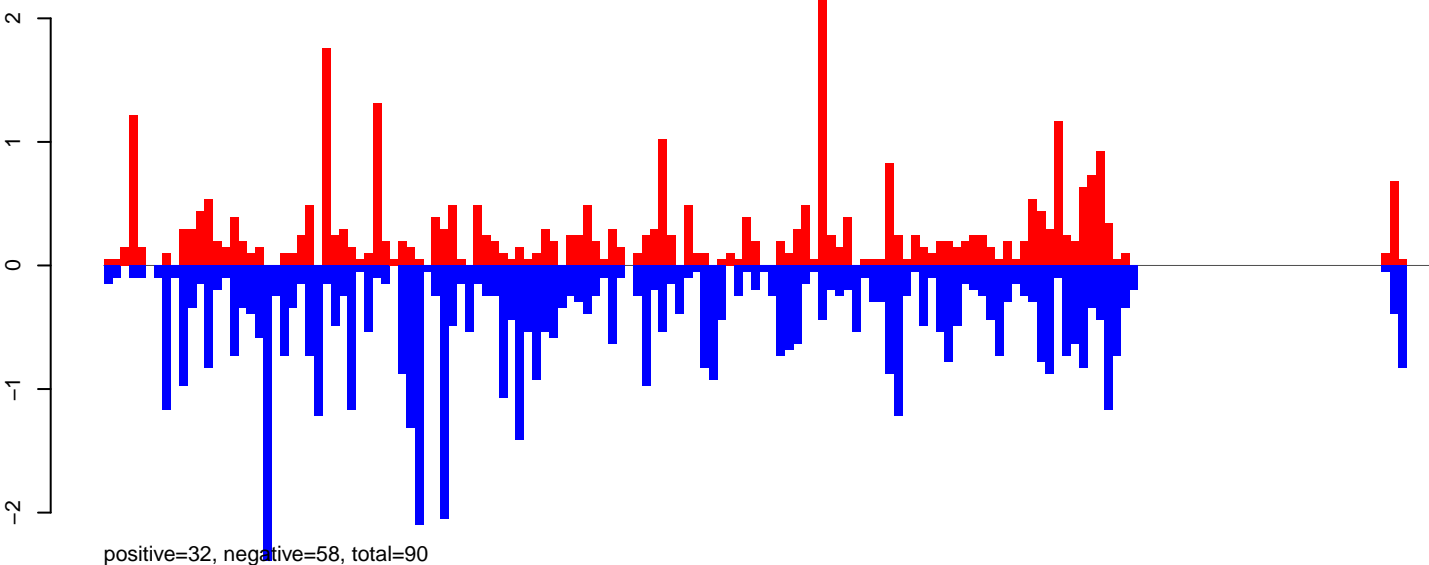
AeAeg_Aag2_cells.24_35.rep



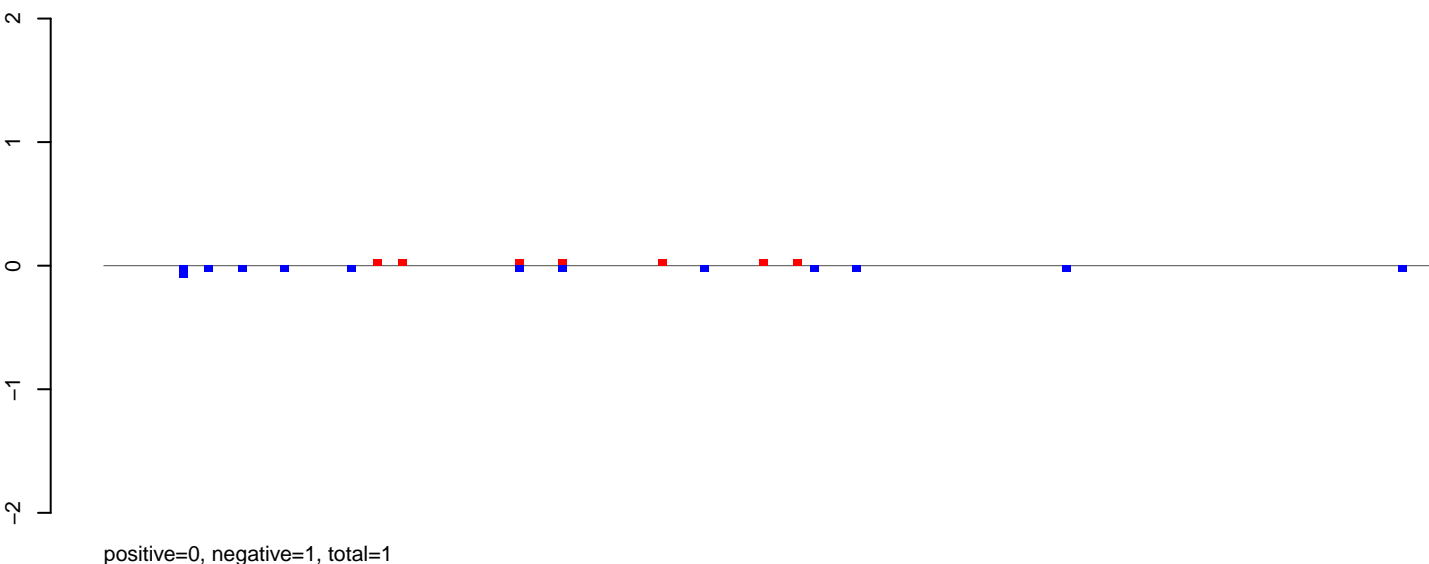
AeAeg_Aag2_cells.rep



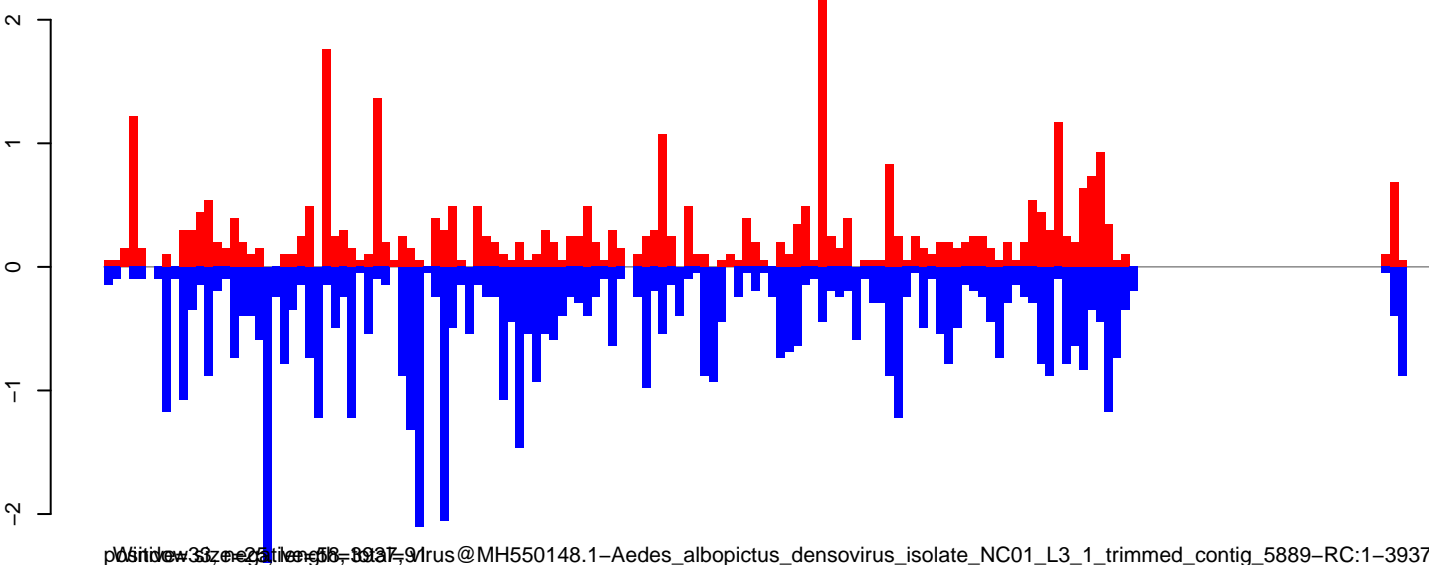
AeAeg_Aag2_cells.18_23.rep



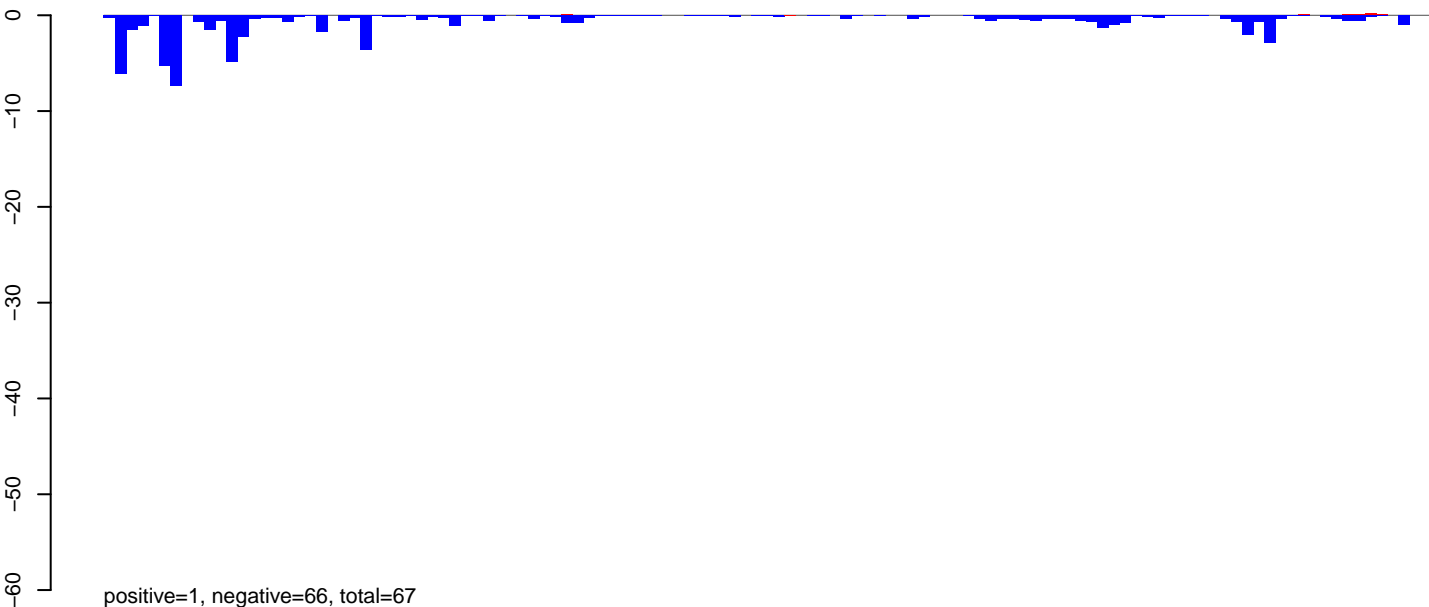
AeAeg_Aag2_cells.24_35.rep



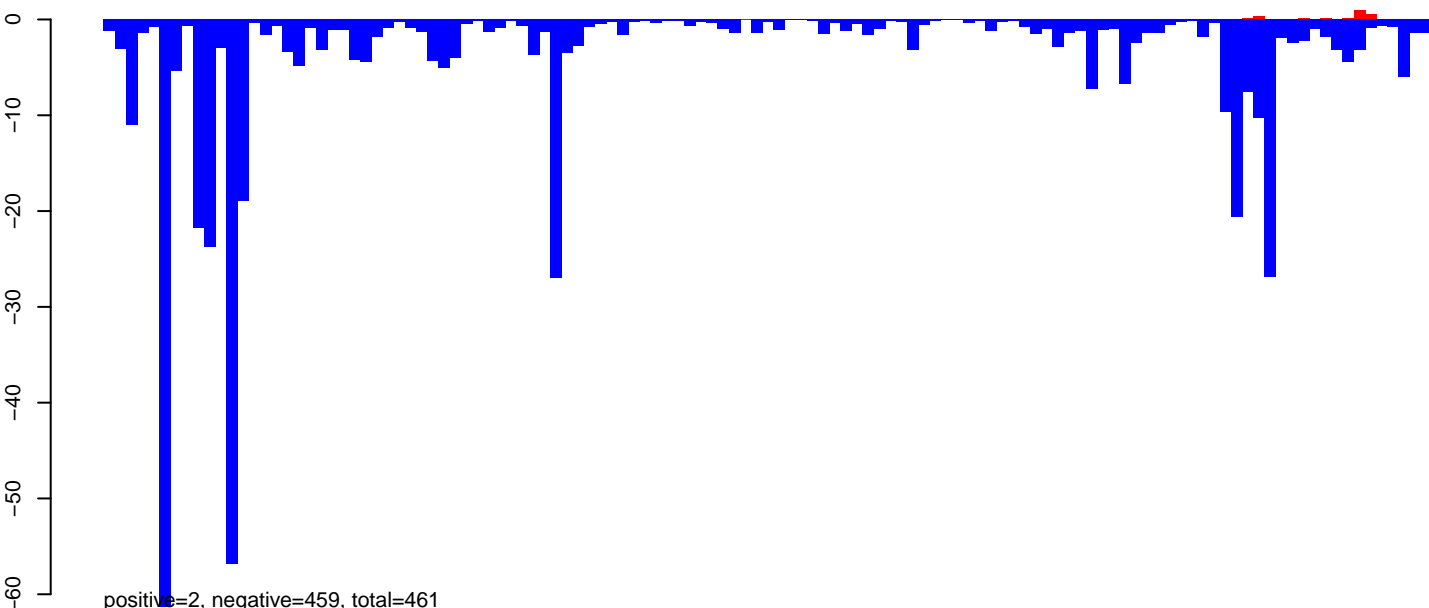
AeAeg_Aag2_cells.rep



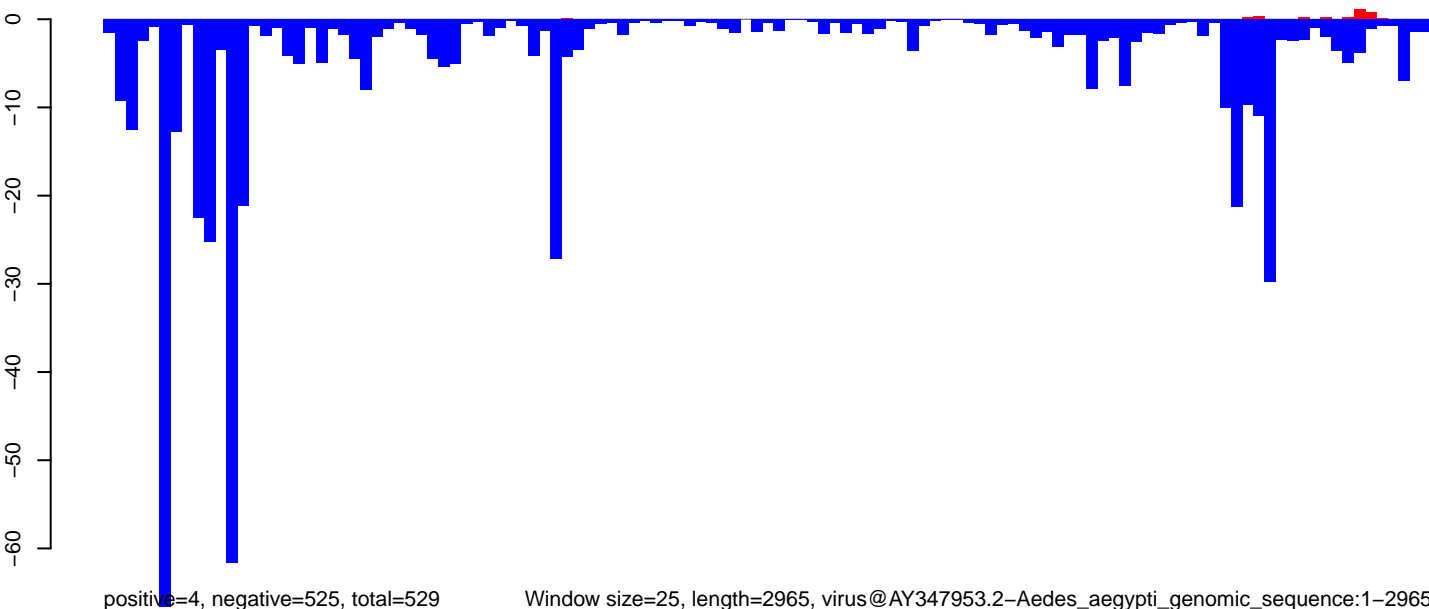
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



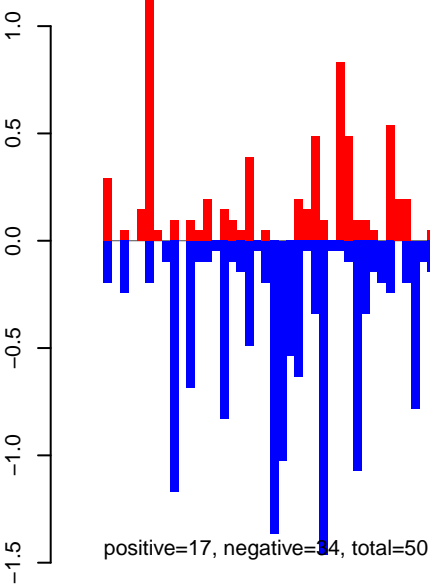
AeAeg_Aag2_cells.rep



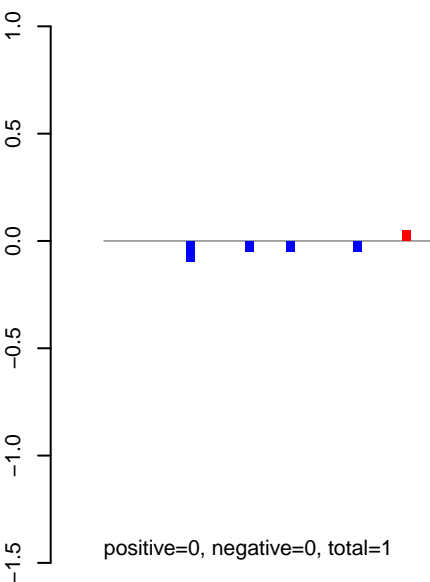
Window size=25, length=2965, virus@AY347953.2-Aedes_aegypti_genomic_sequence:1-2965

0 500 1000 1500 2000 2500 3000

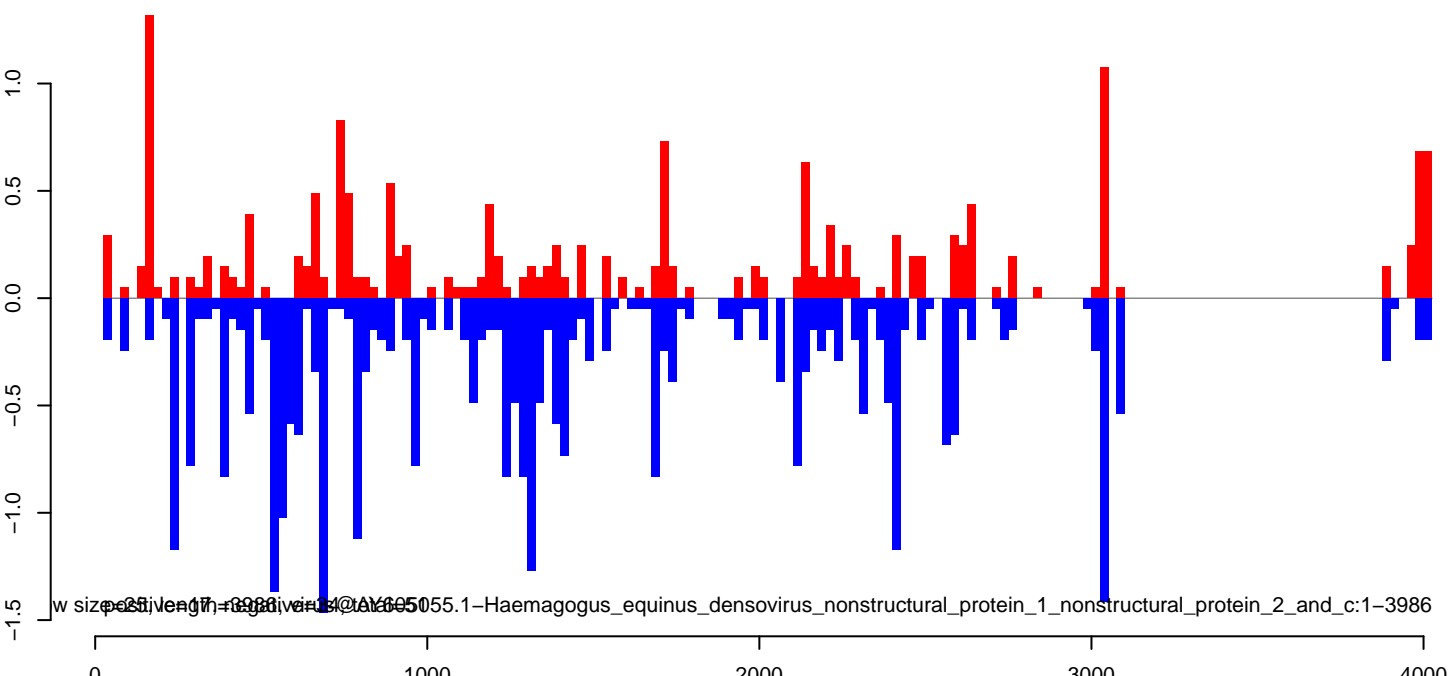
AeAeg_Aag2_cells.18_23.rep



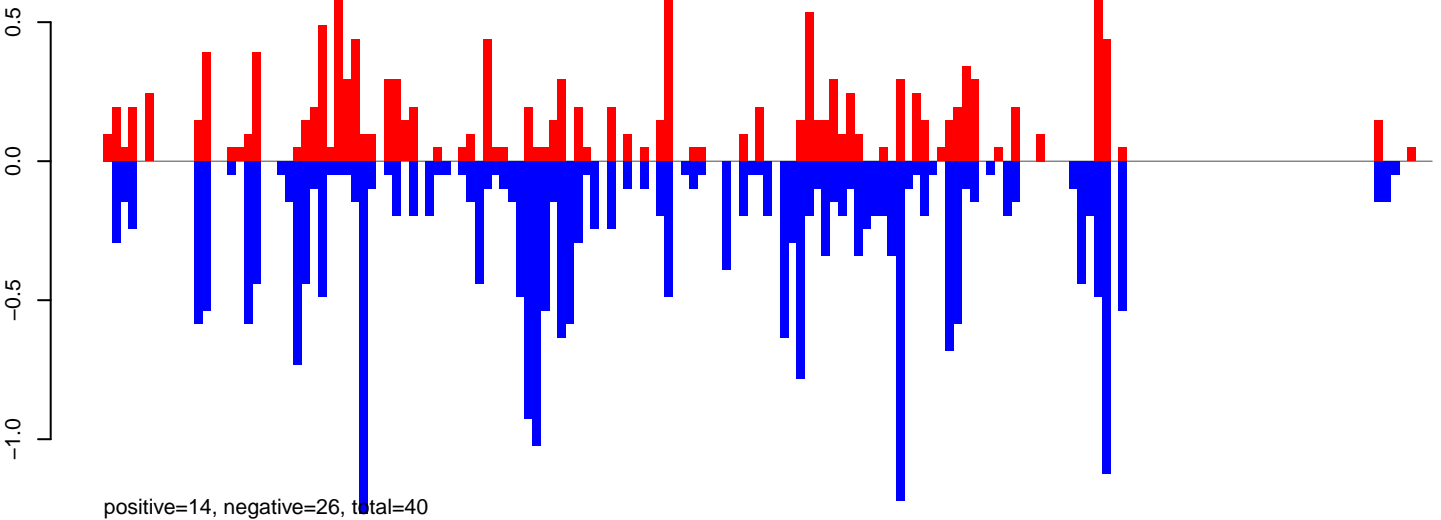
AeAeg_Aag2_cells.24_35.rep



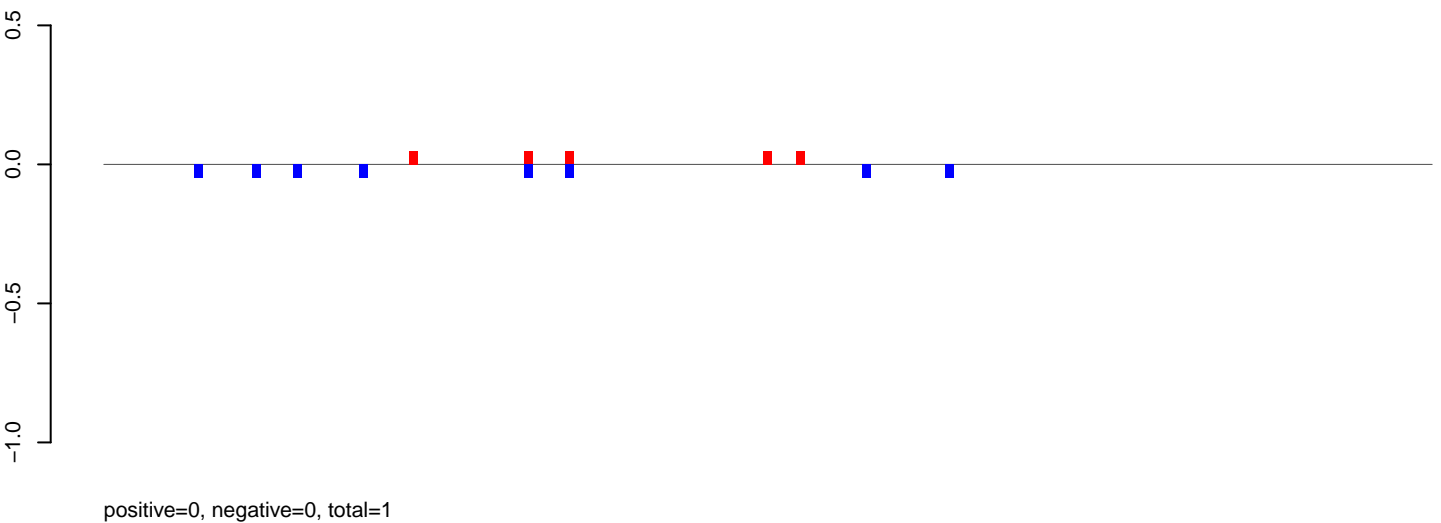
AeAeg_Aag2_cells.rep



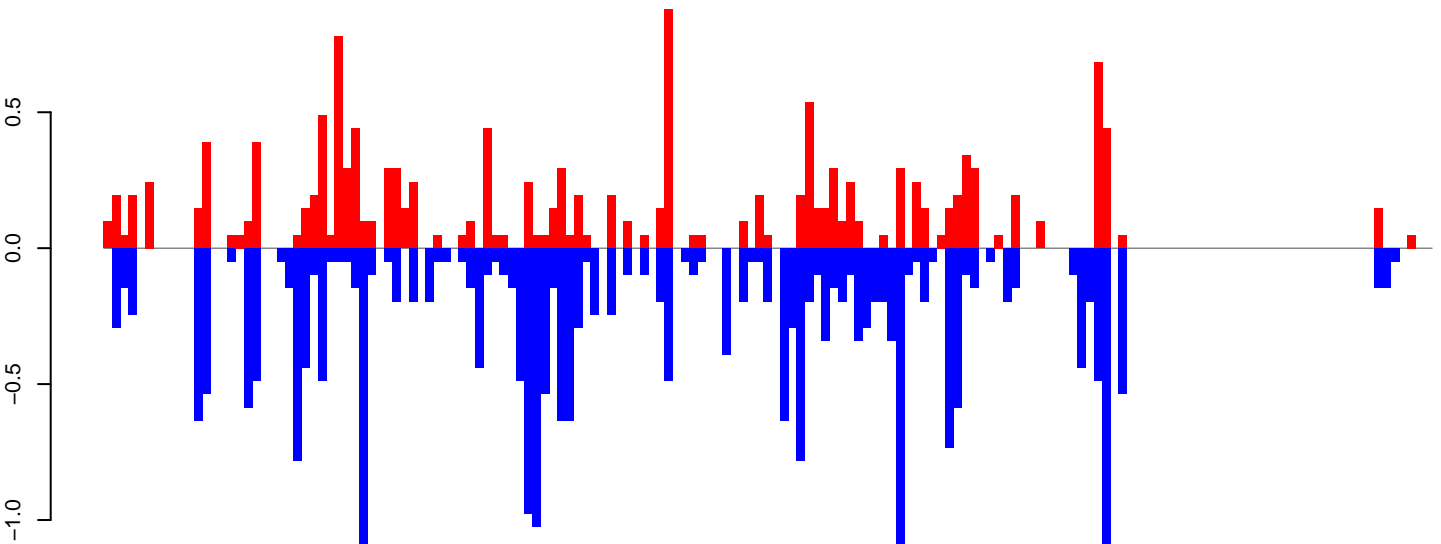
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



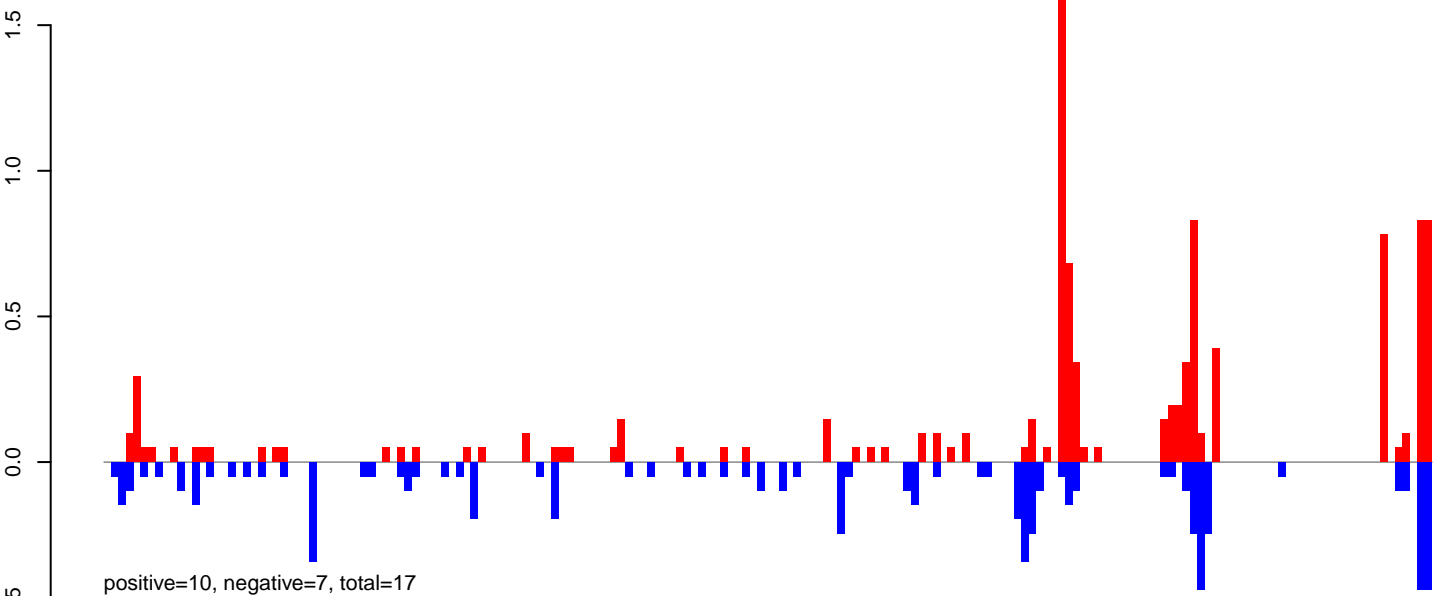
AeAeg_Aag2_cells.rep



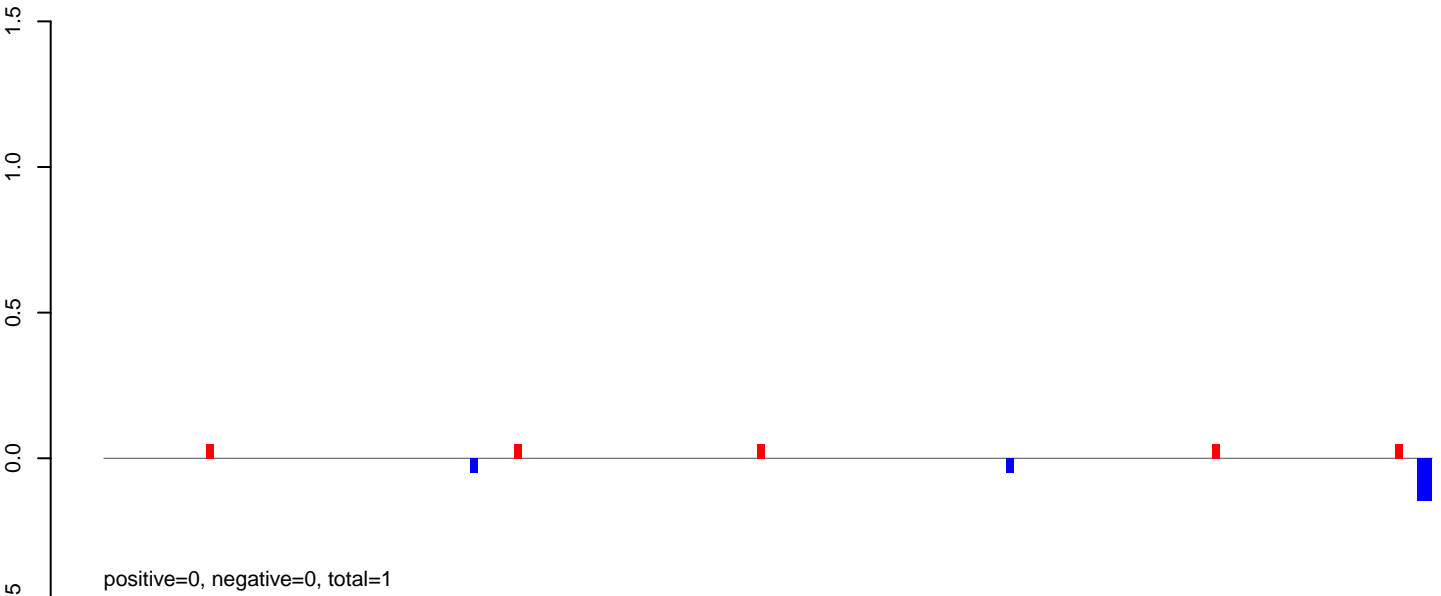
10877.1-Aedes albopictus_C6_36_cell_densovirus_non-structural_protein_1_non-structural_prot:1-4006

0 1000 2000 3000 4000

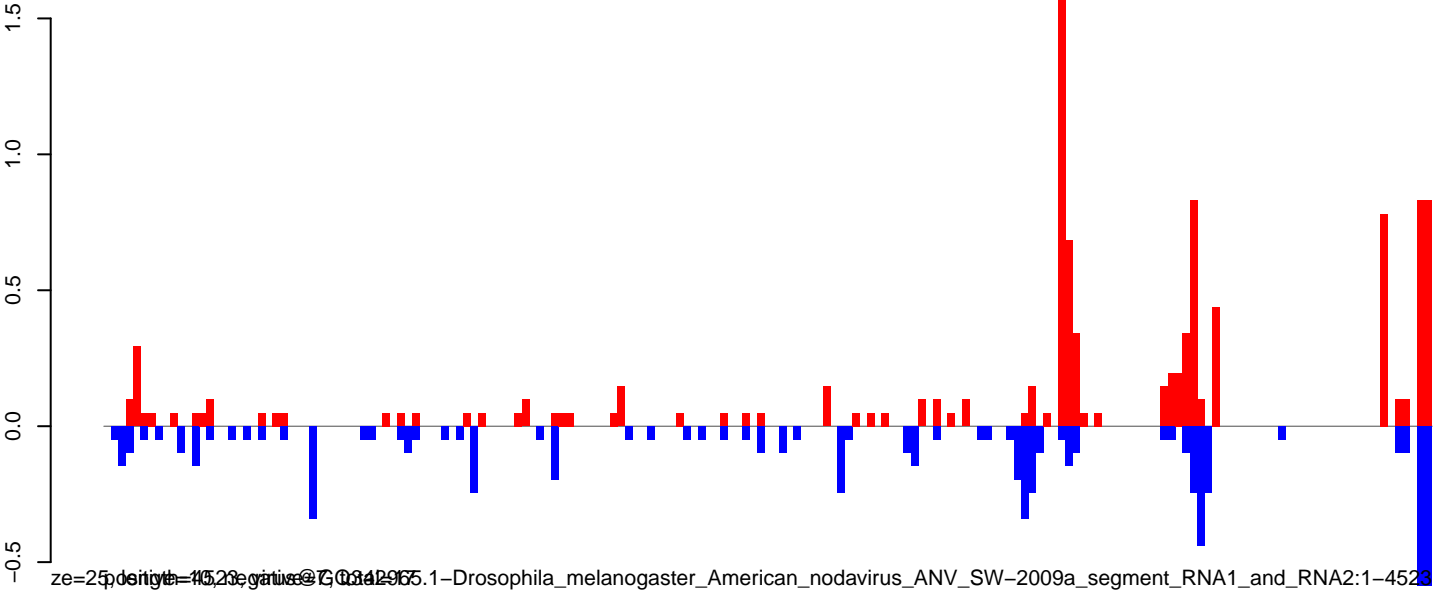
AeAeg_Aag2_cells.18_23.rep



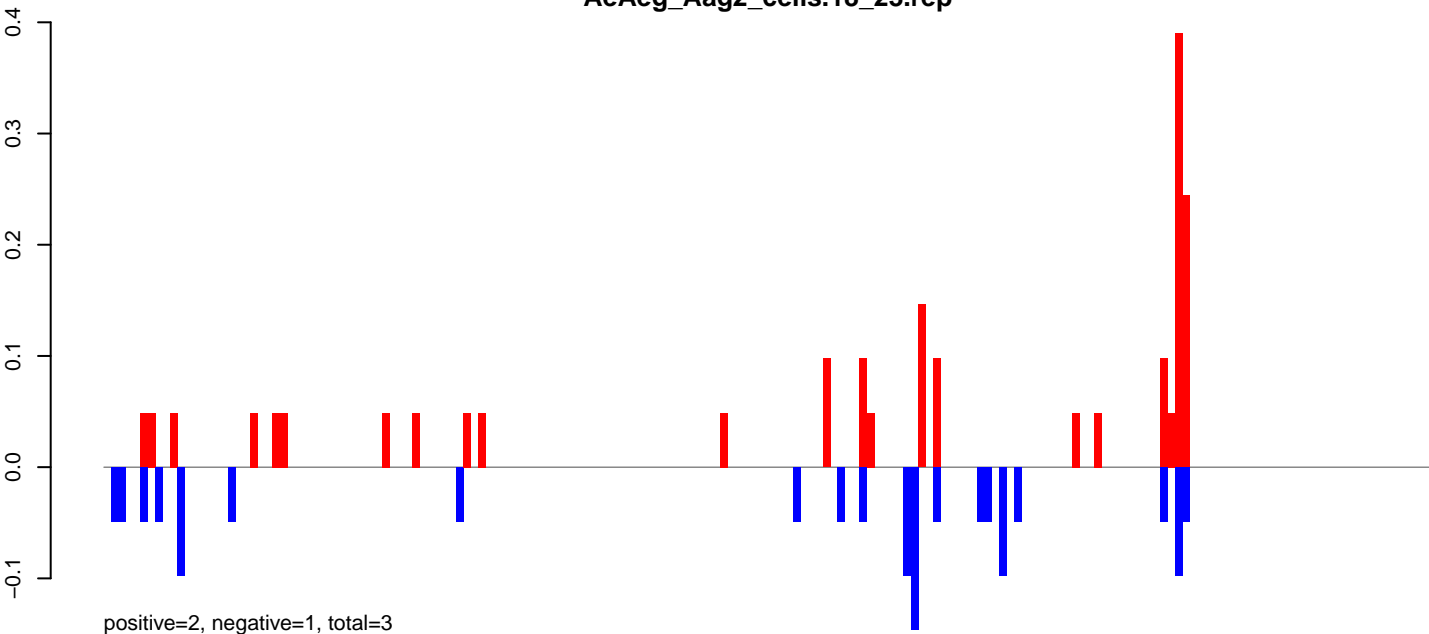
AeAeg_Aag2_cells.24_35.rep



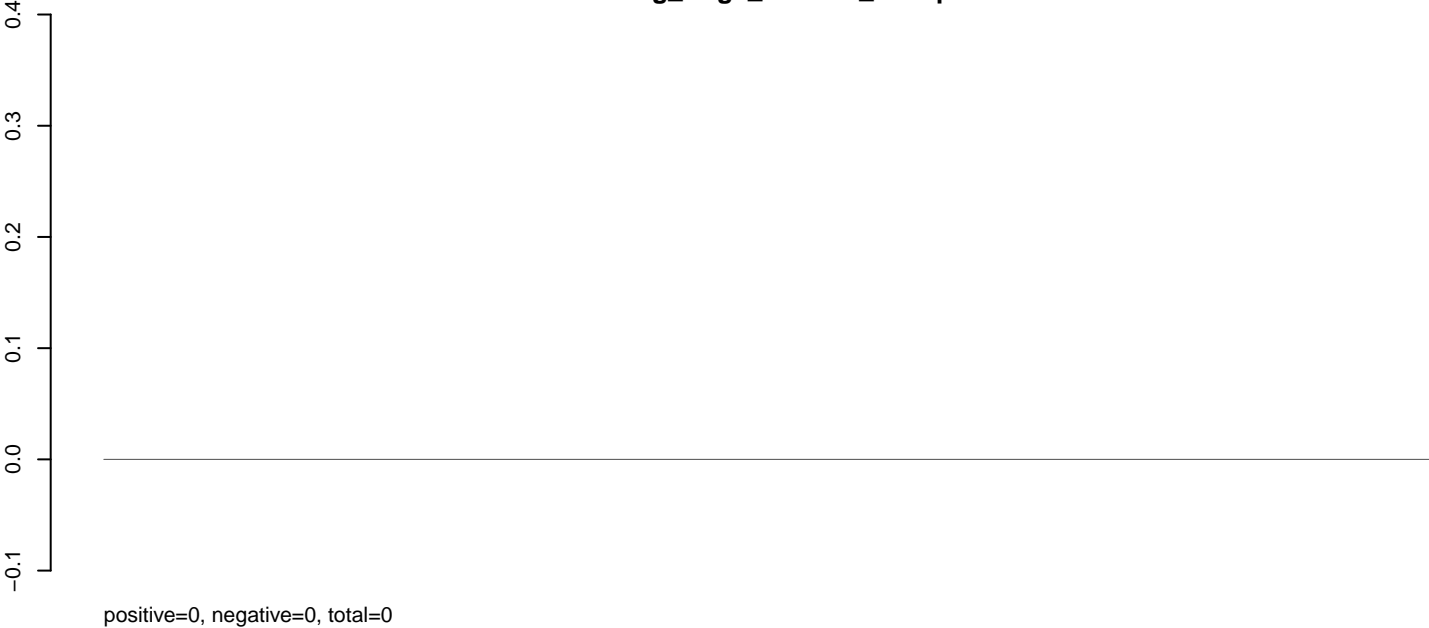
AeAeg_Aag2_cells.rep



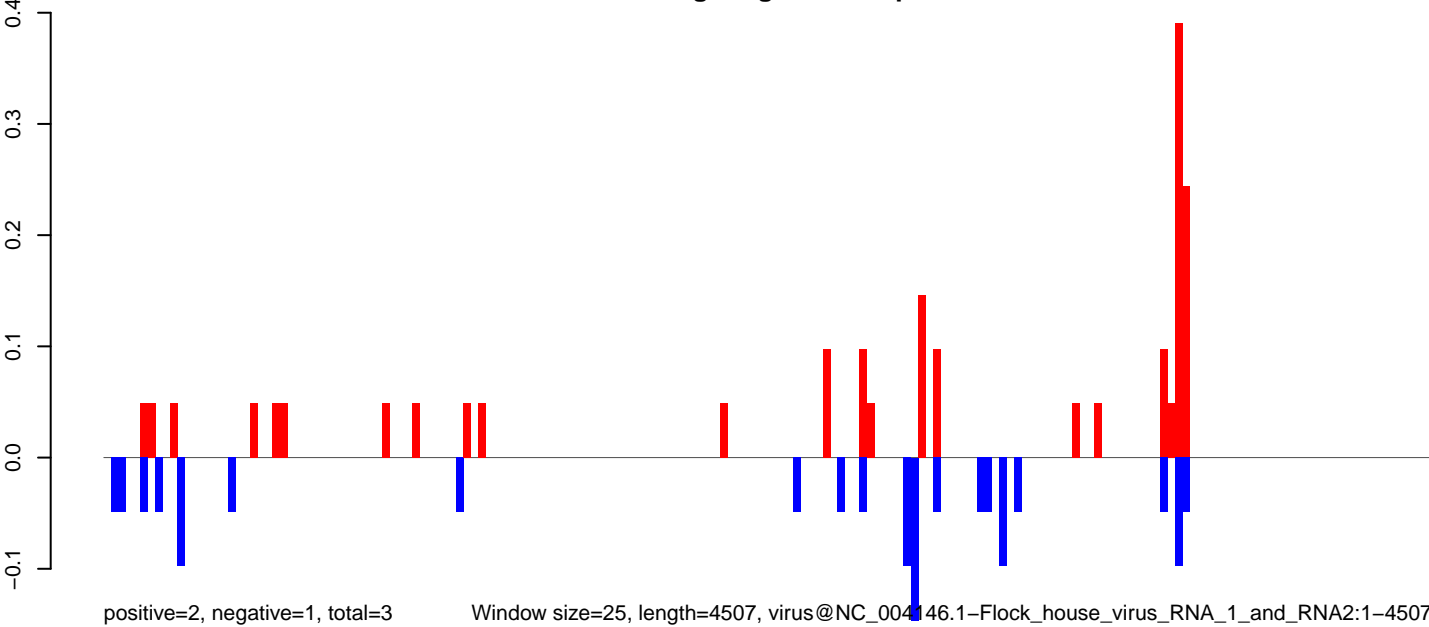
AeAeg_Aag2_cells.18_23.rep



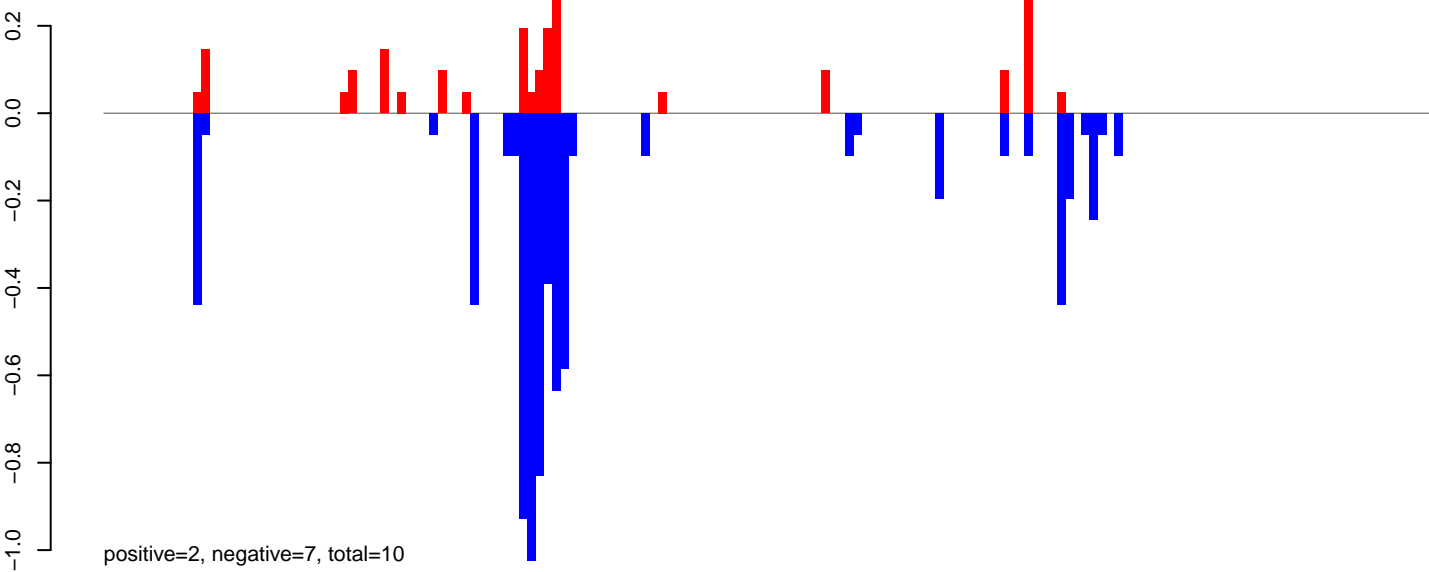
AeAeg_Aag2_cells.24_35.rep



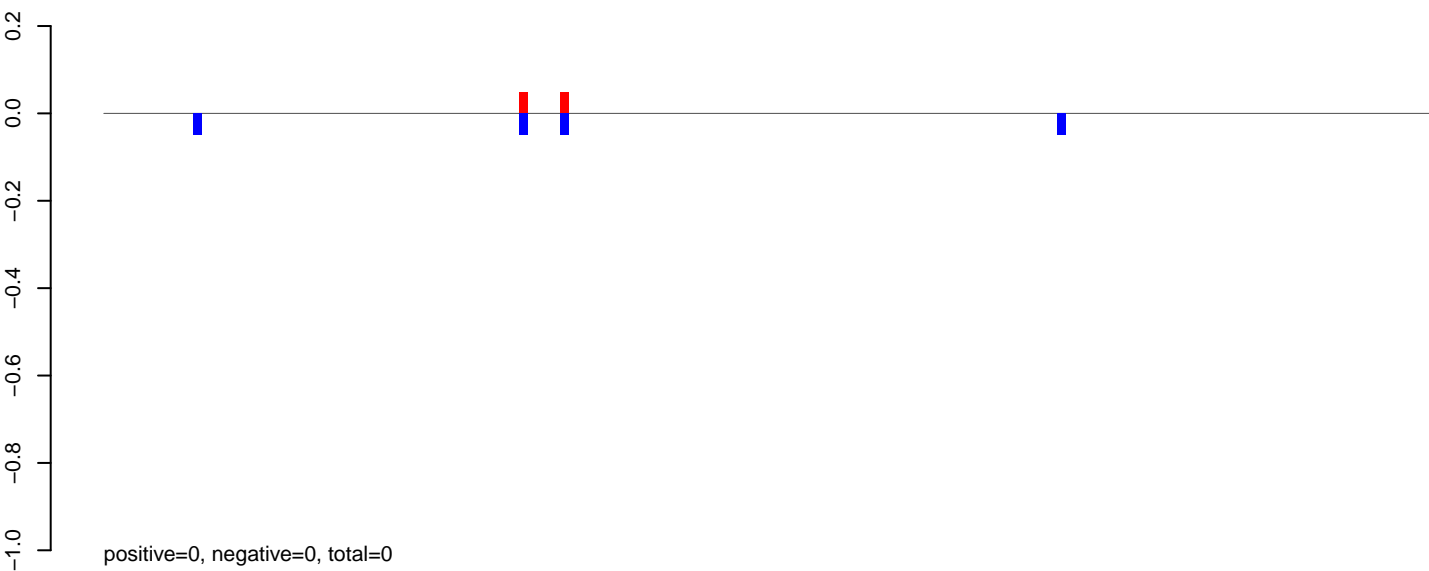
AeAeg_Aag2_cells.rep



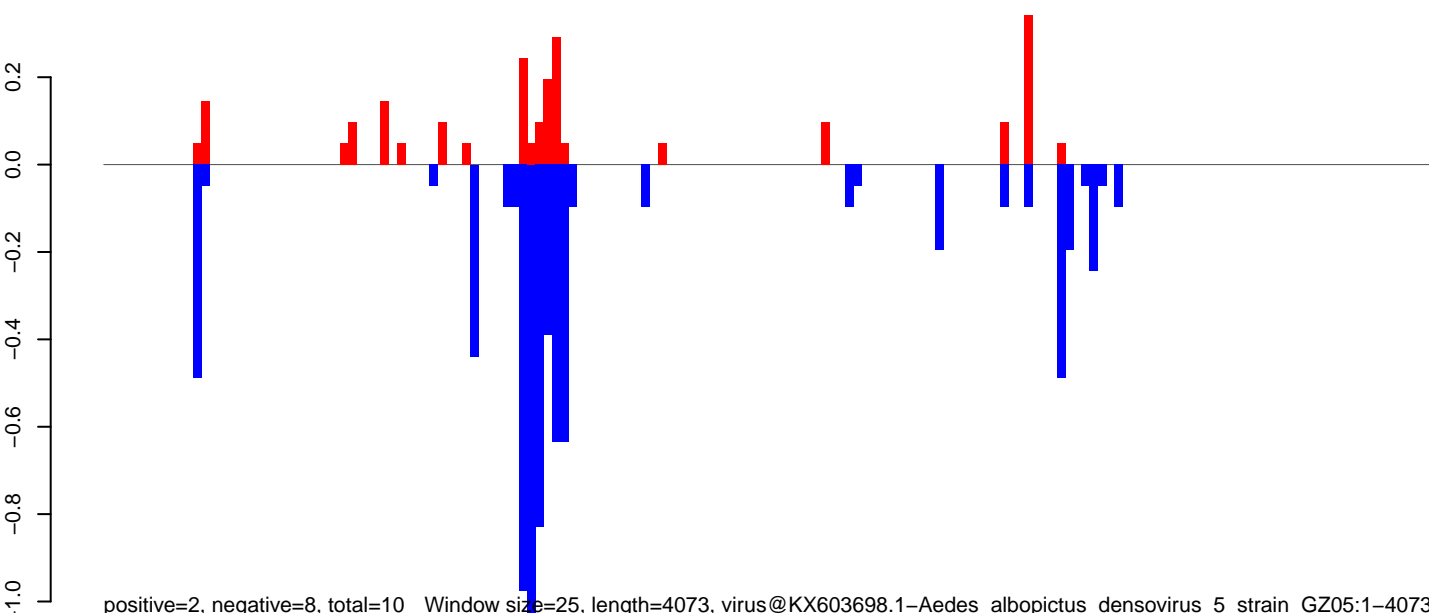
AeAeg_Aag2_cells.18_23.rep



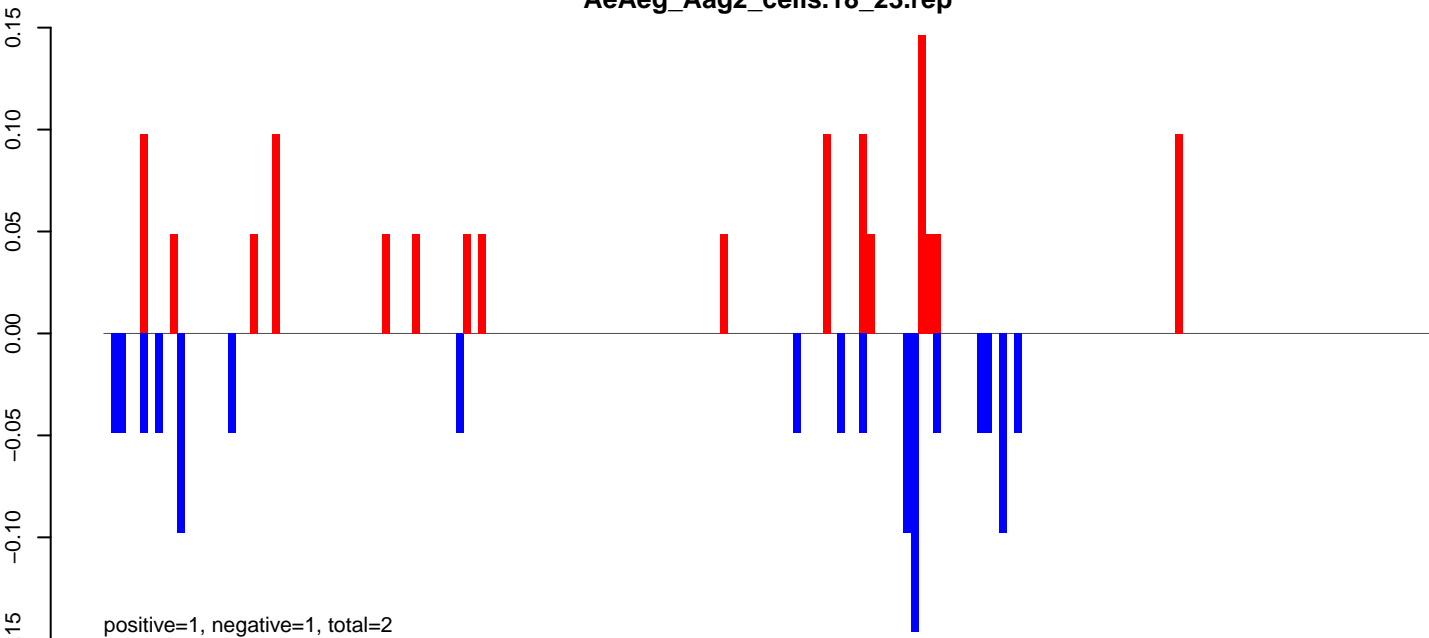
AeAeg_Aag2_cells.24_35.rep



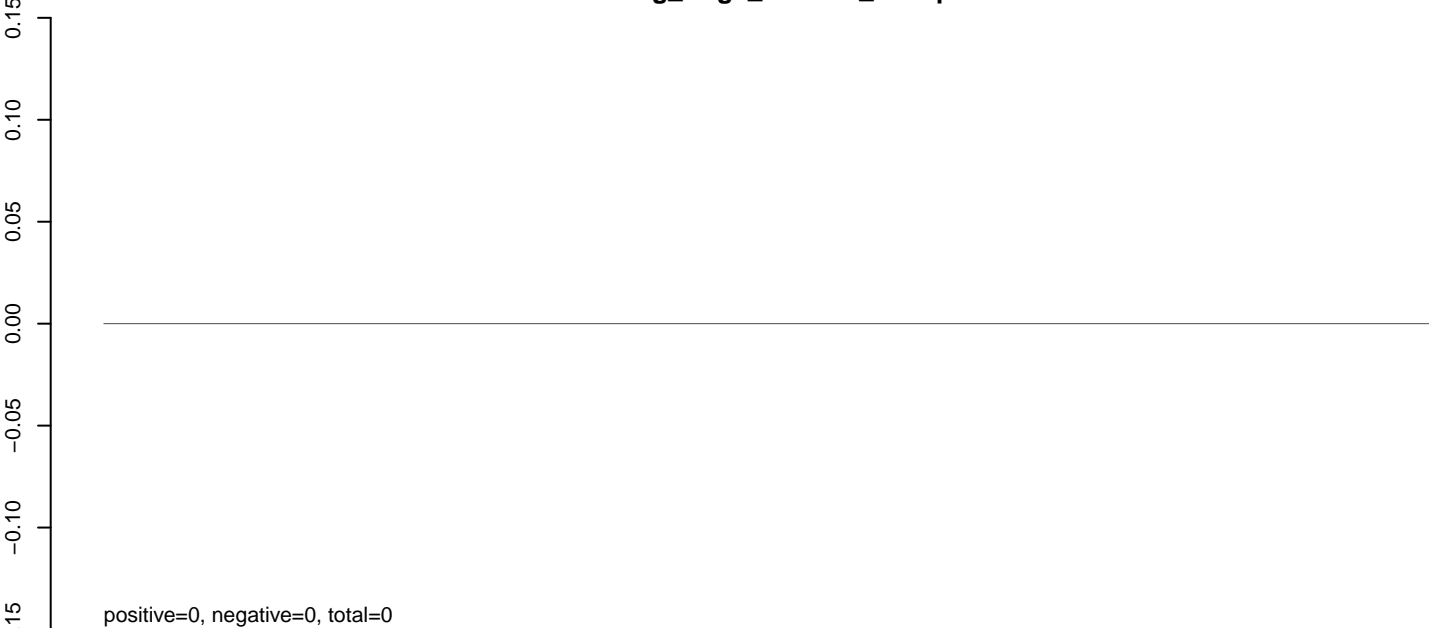
AeAeg_Aag2_cells.rep



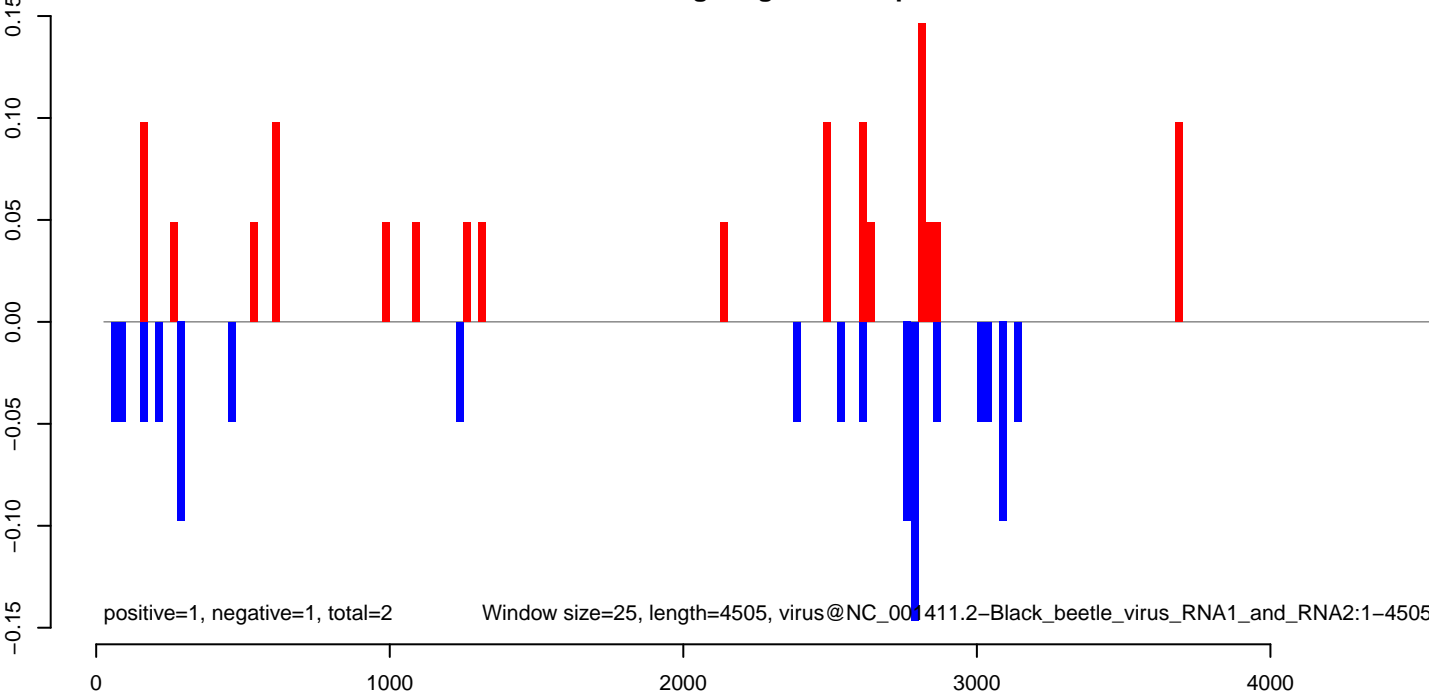
AeAeg_Aag2_cells.18_23.rep



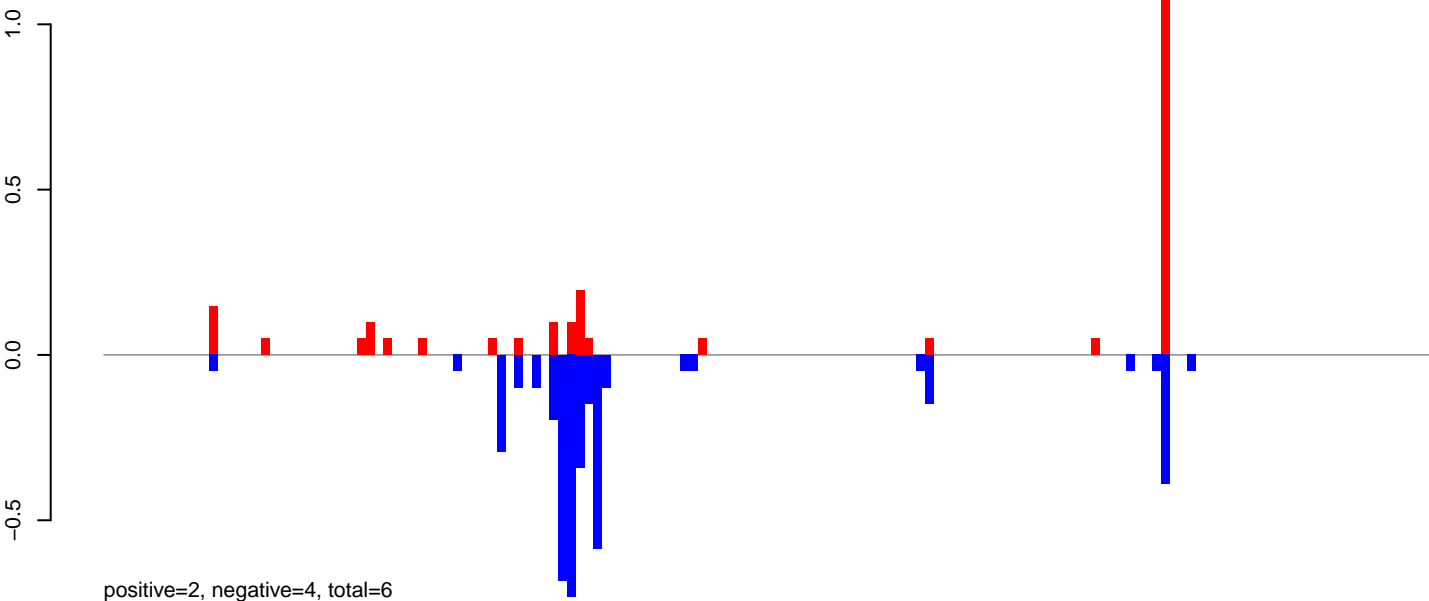
AeAeg_Aag2_cells.24_35.rep



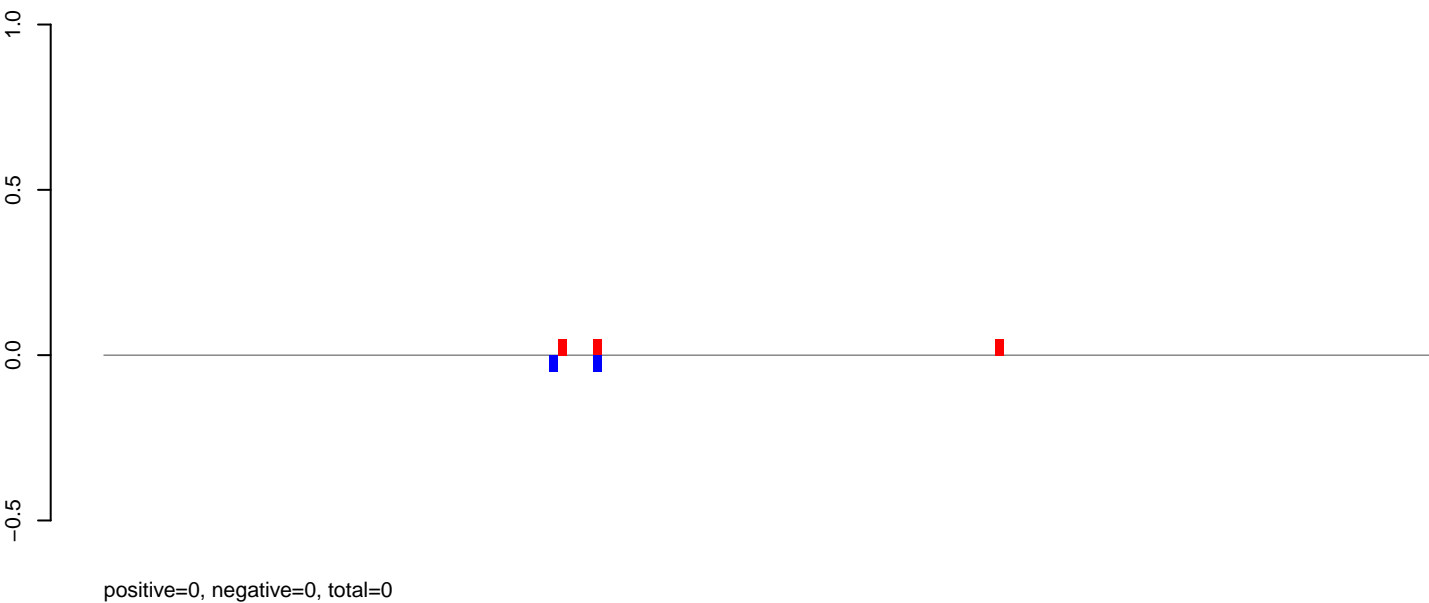
AeAeg_Aag2_cells.rep



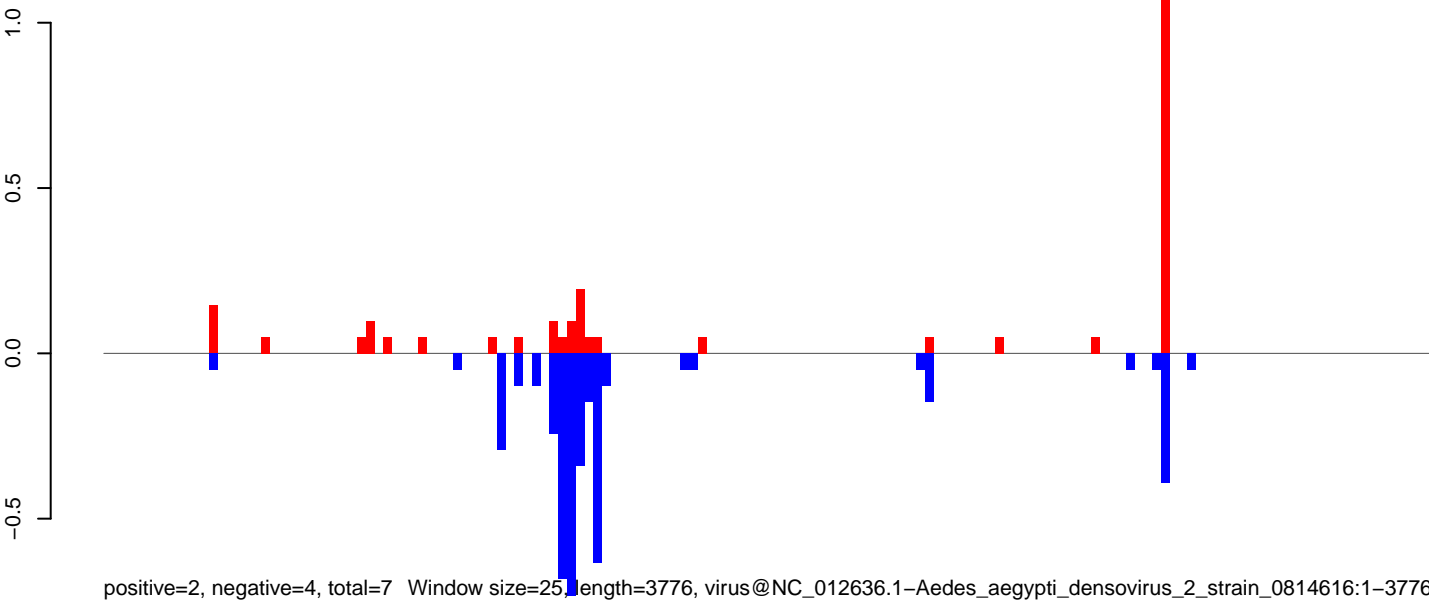
AeAeg_Aag2_cells.18_23.rep



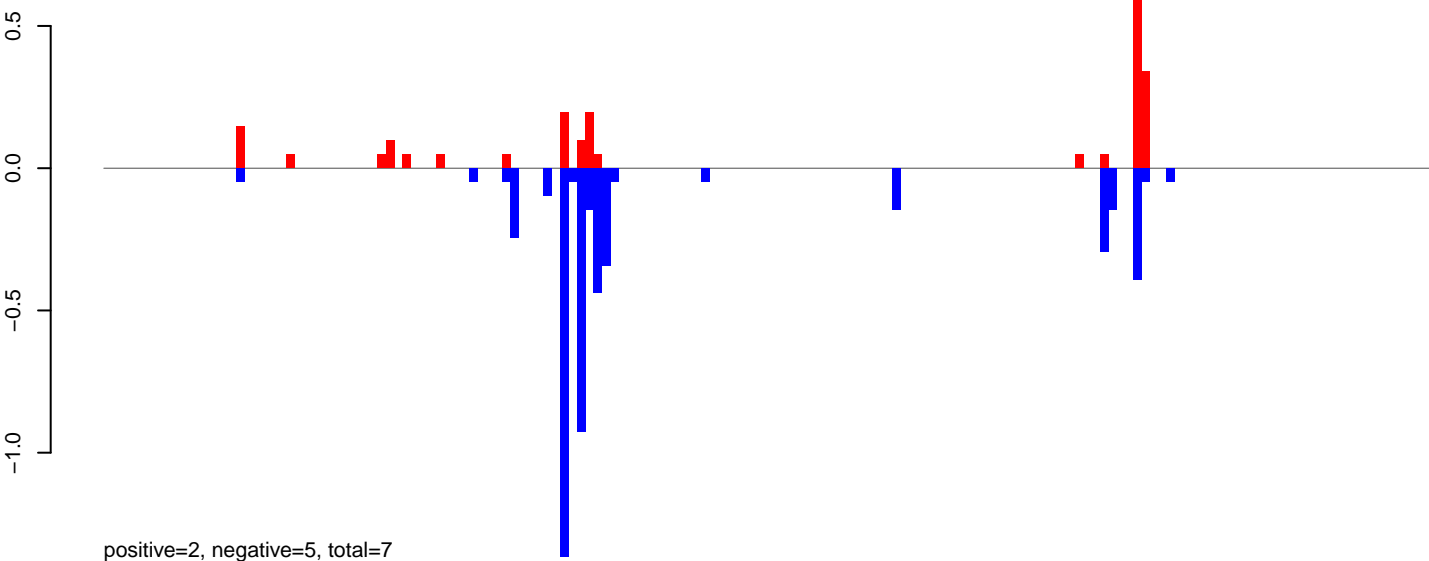
AeAeg_Aag2_cells.24_35.rep



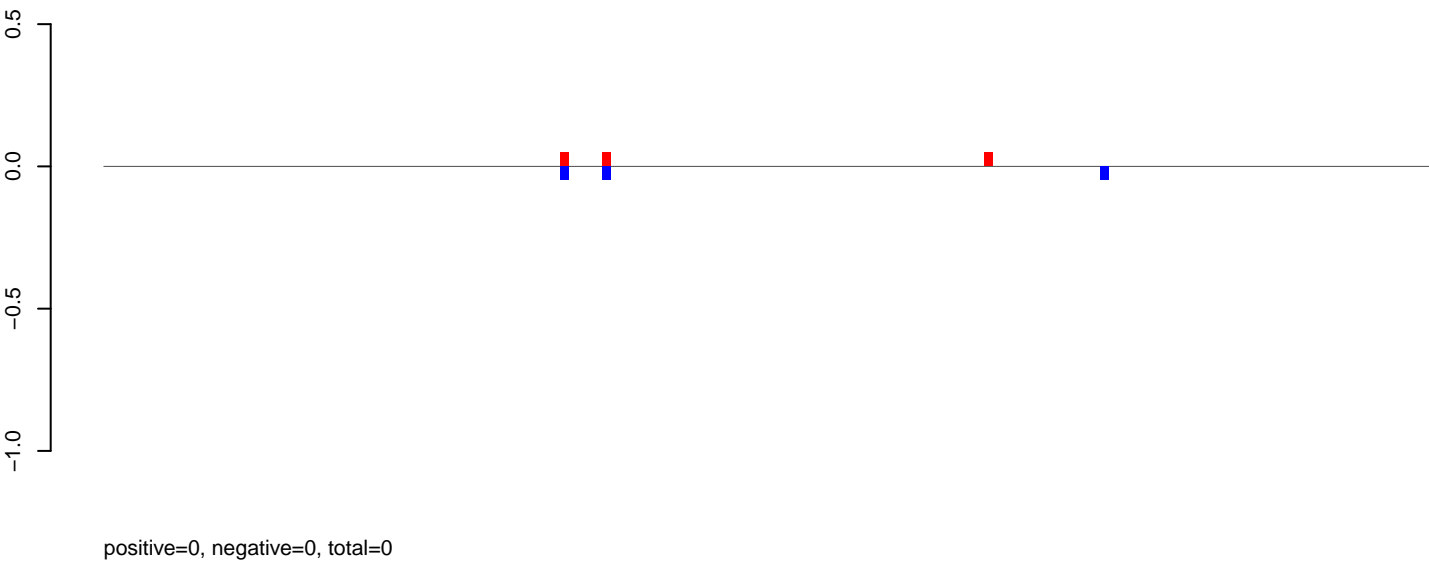
AeAeg_Aag2_cells.rep



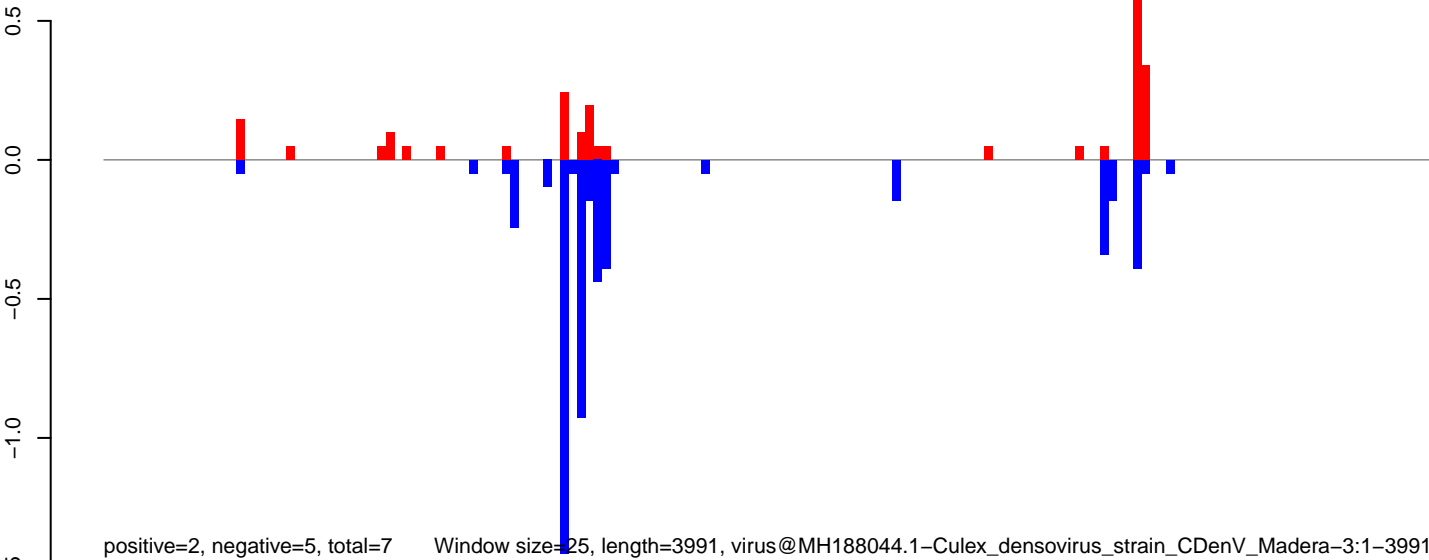
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



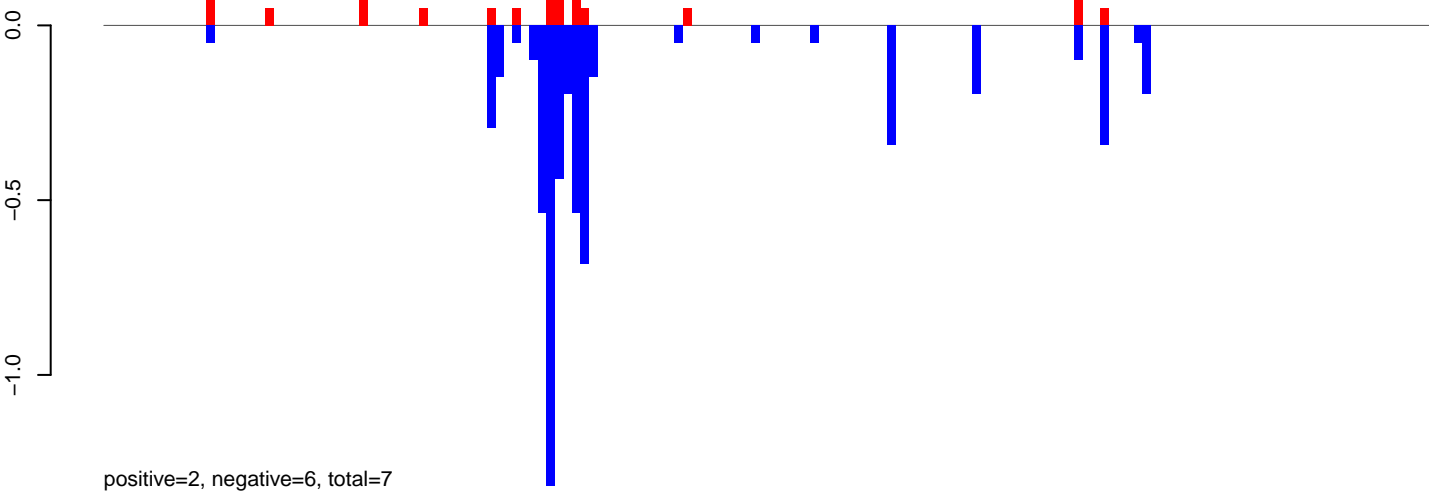
AeAeg_Aag2_cells.rep



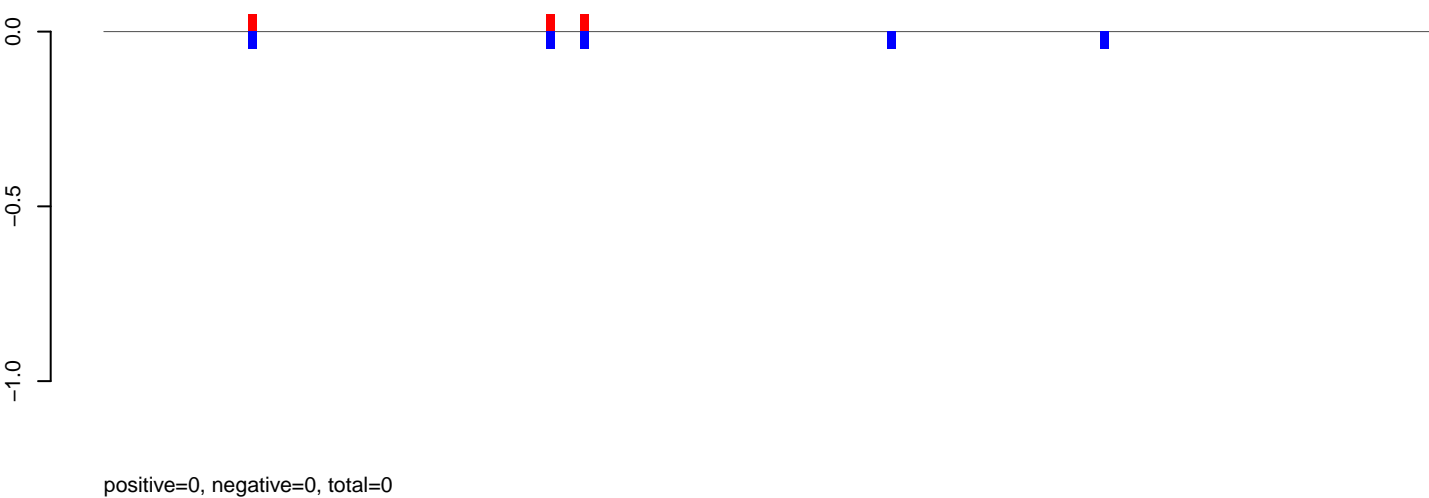
Window size=25, length=3991, virus@MH188044.1-Culex_densovirus_strain_CDenV_Madera-3:1-3991

0 1000 2000 3000 4000

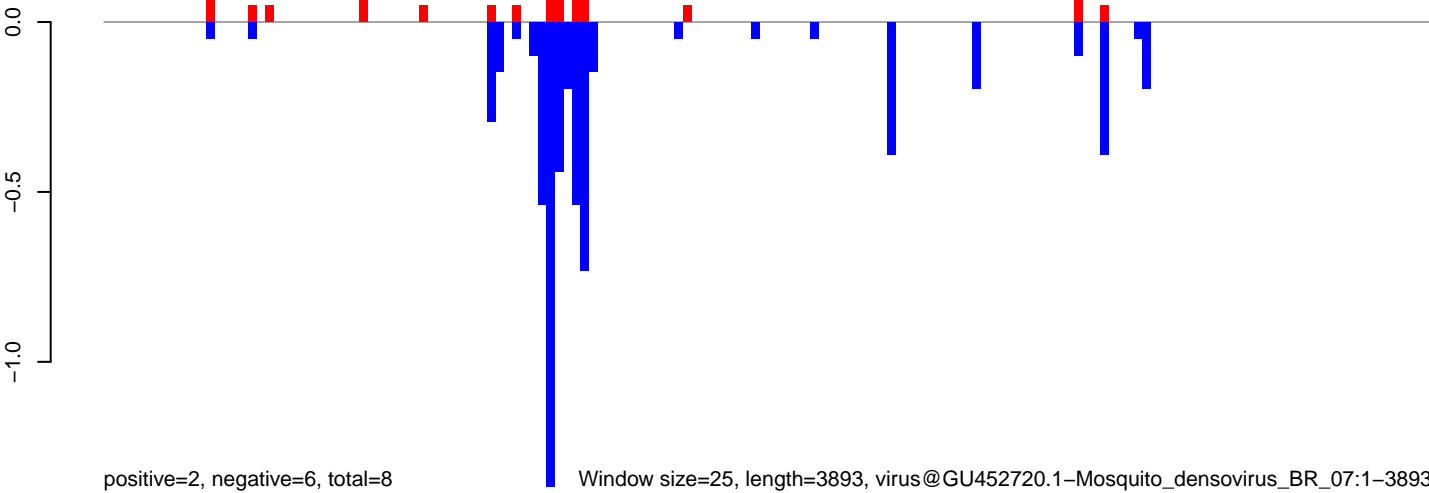
AeAeg_Aag2_cells.18_23.rep



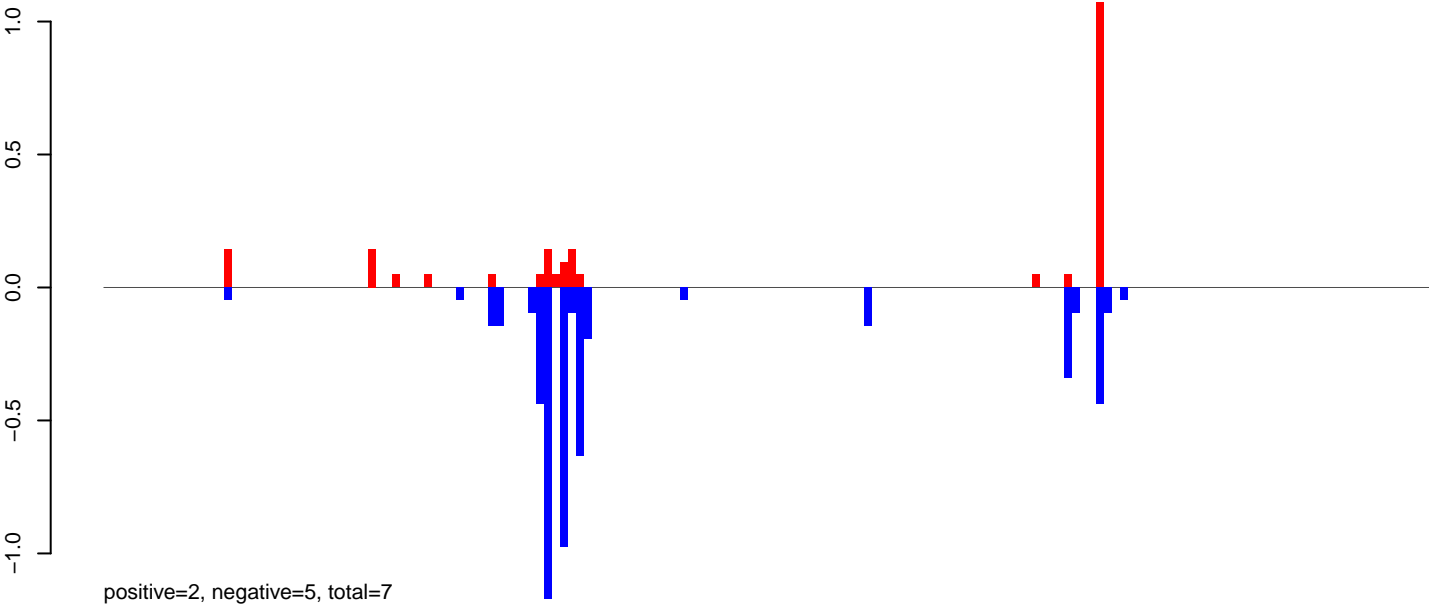
AeAeg_Aag2_cells.24_35.rep



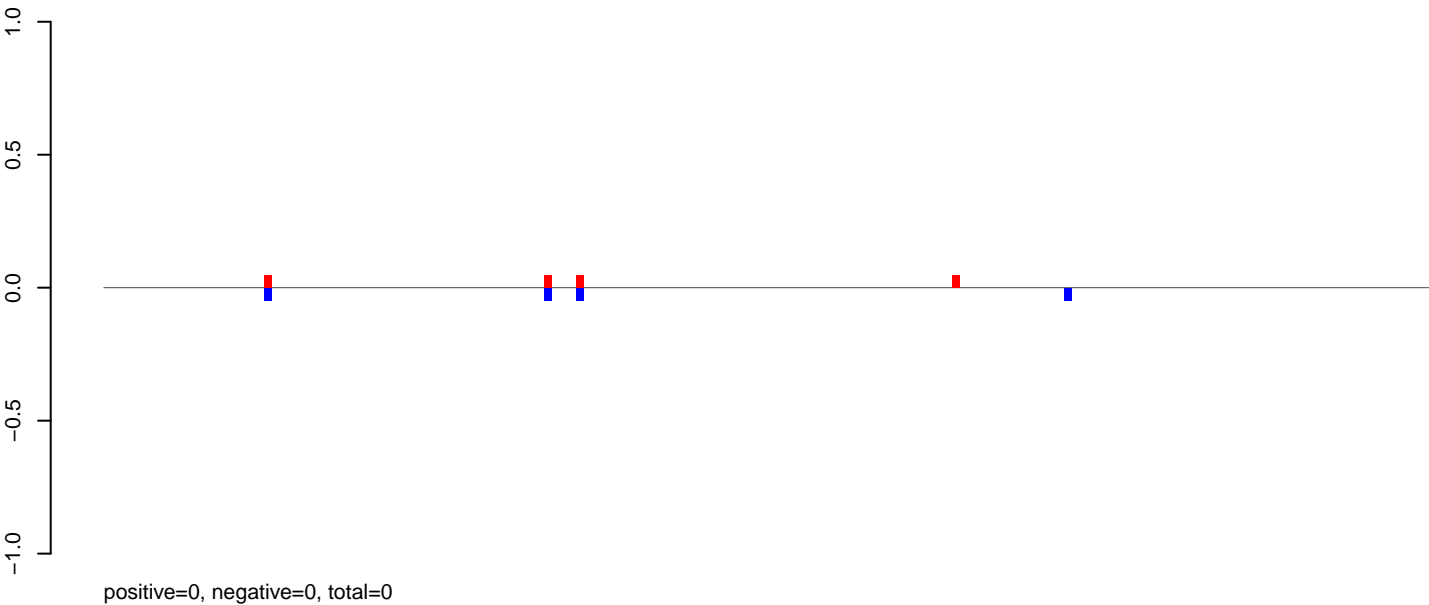
AeAeg_Aag2_cells.rep



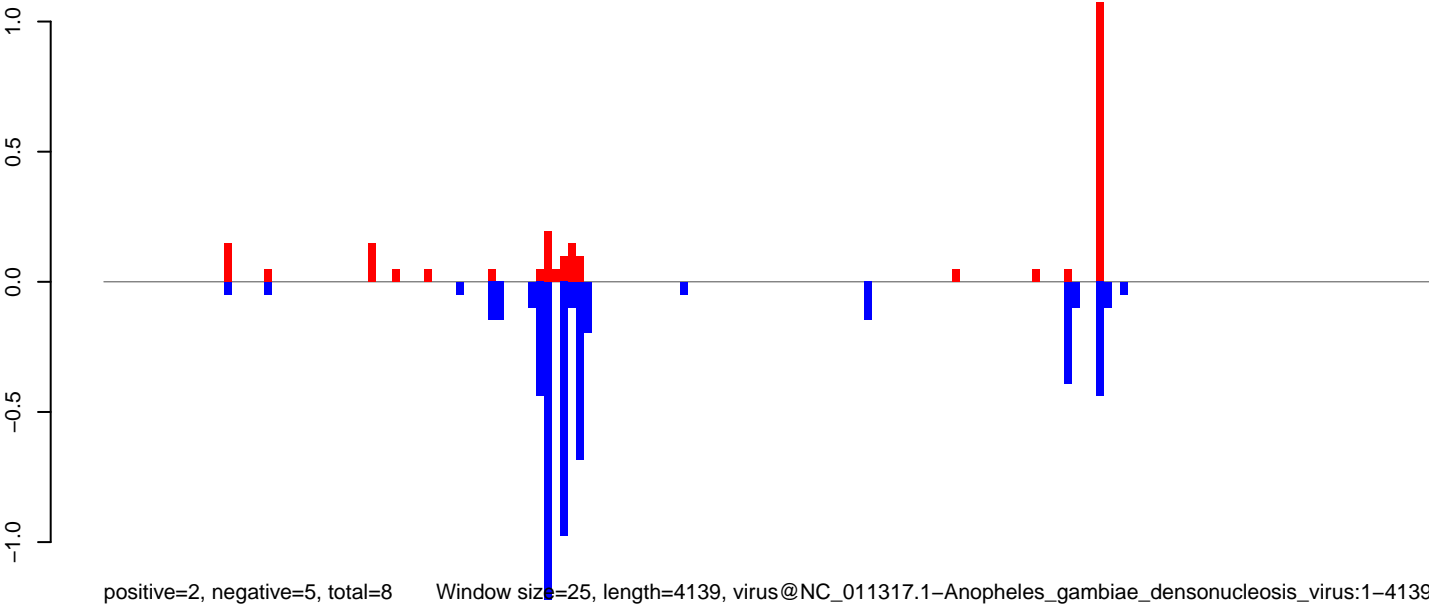
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

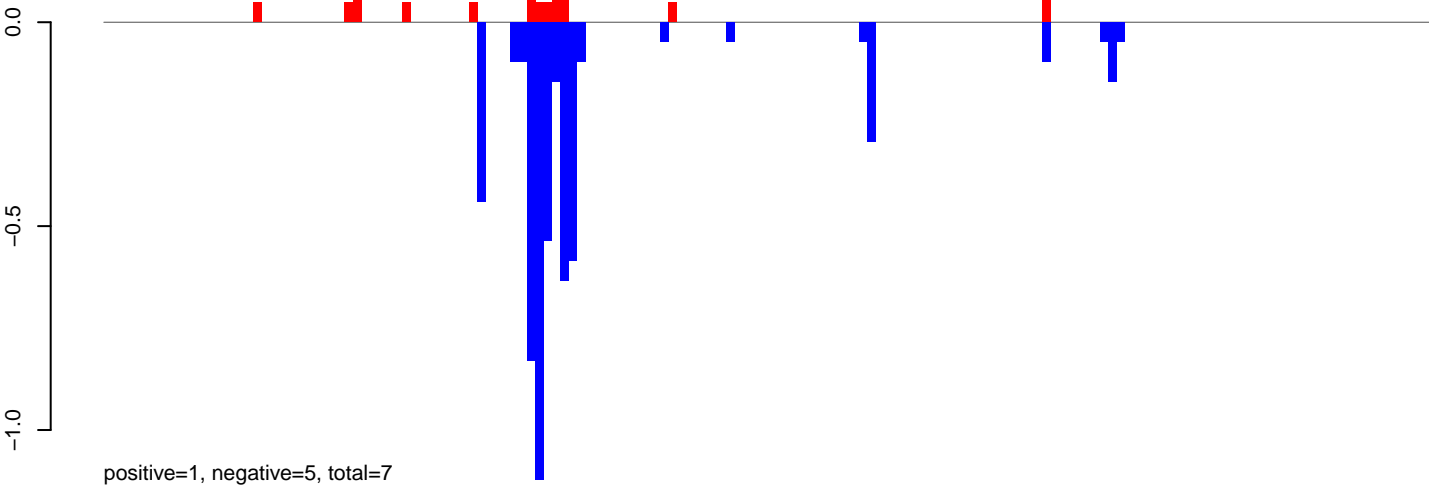


AeAeg_Aag2_cells.rep

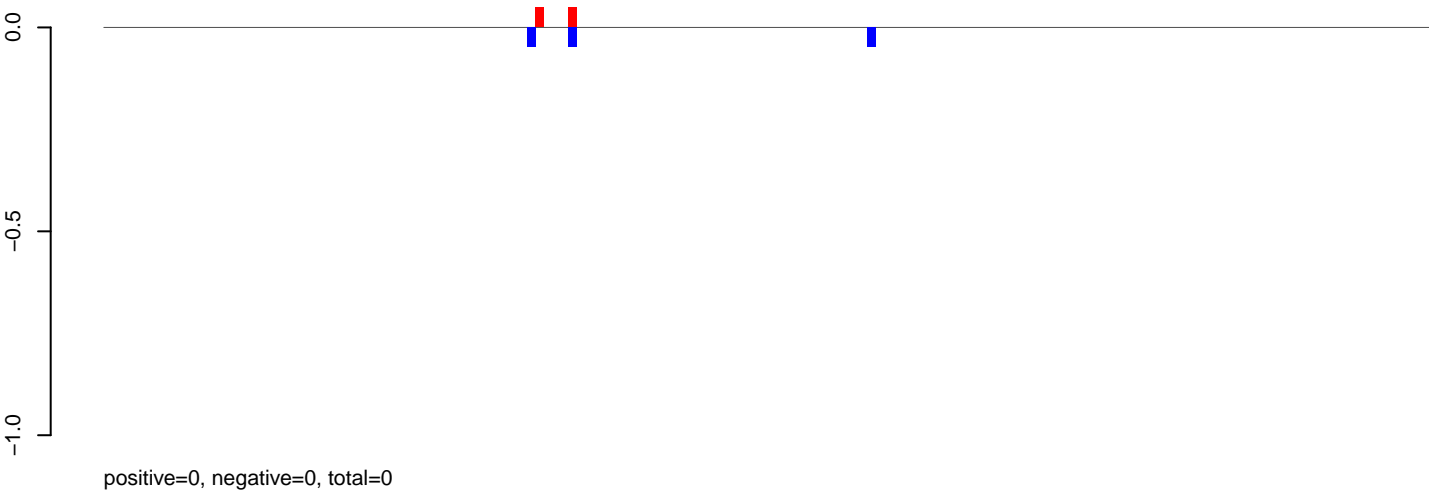


0 1000 2000 3000 4000

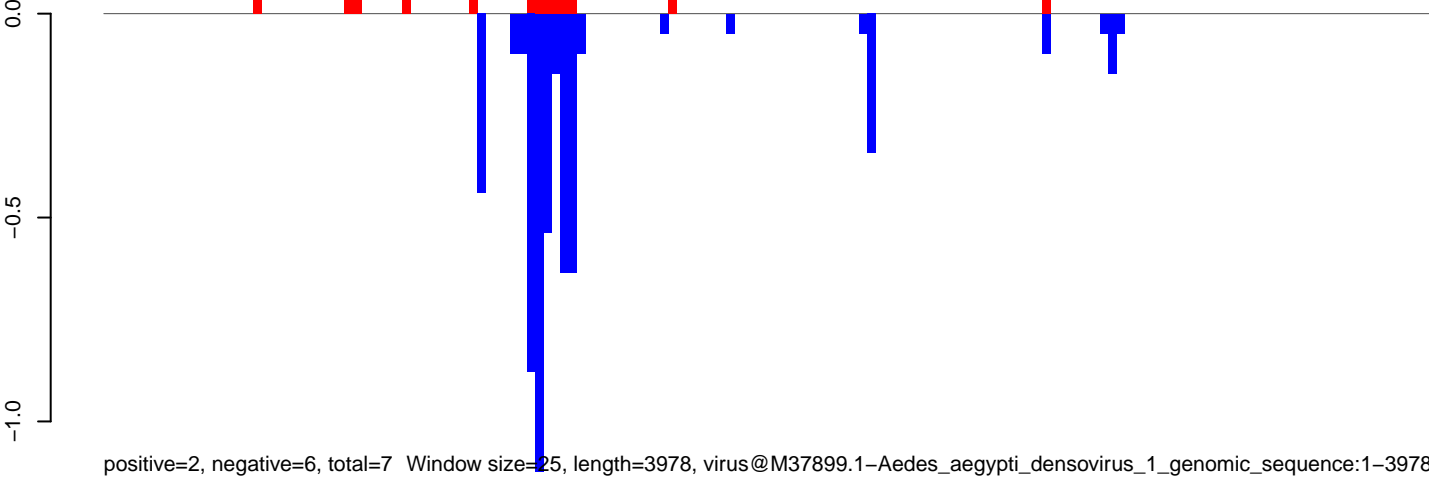
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

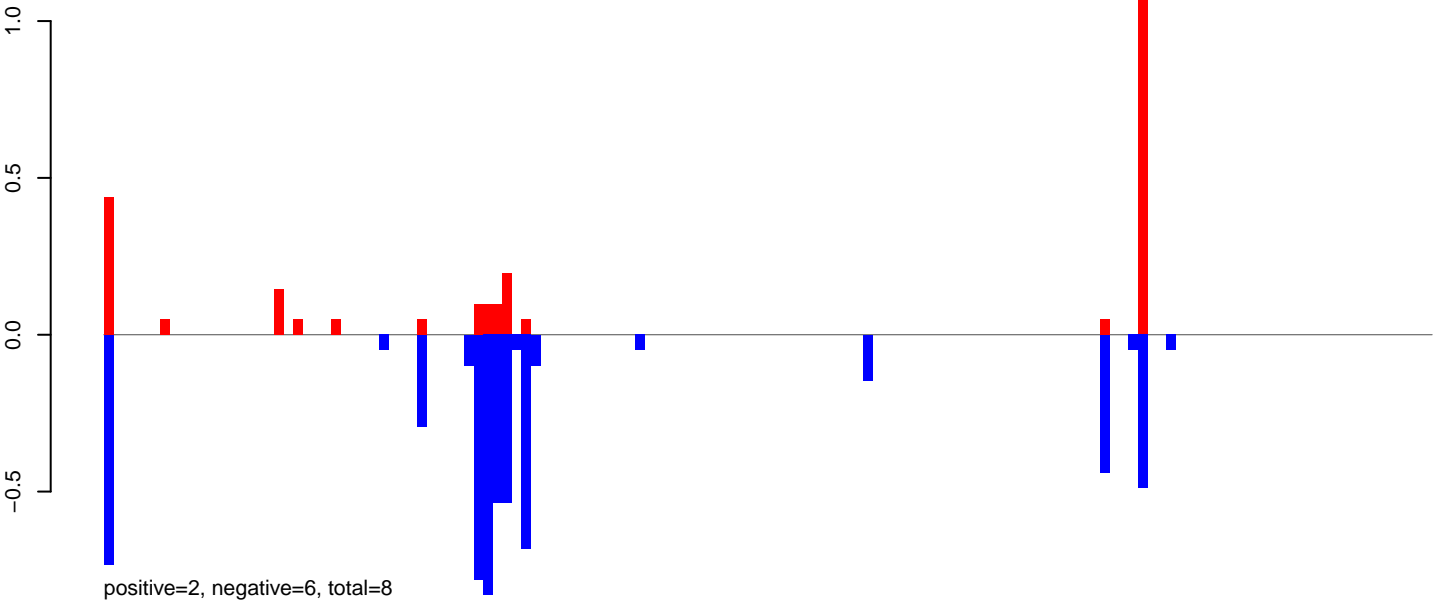


AeAeg_Aag2_cells.rep

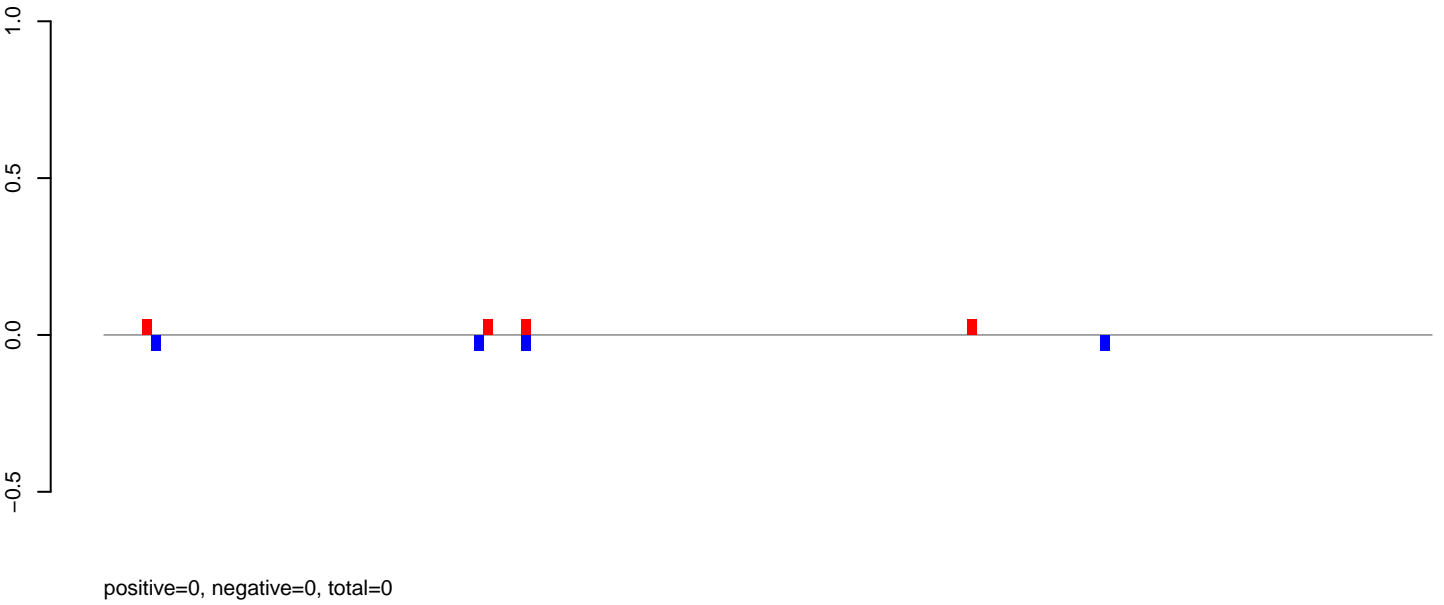


Window size=25, length=3978, virus@M37899.1-Aedes_aegypti_densovirus_1_genomic_sequence:1-3978

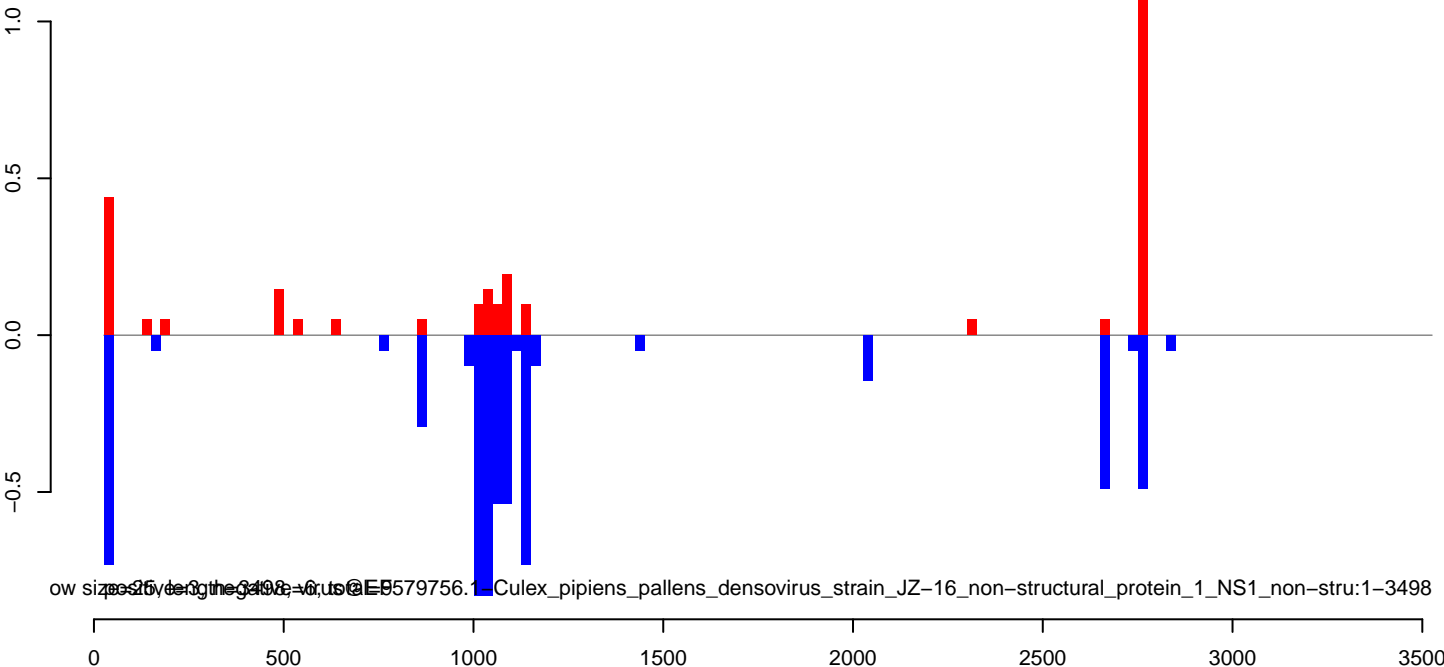
AeAeg_Aag2_cells.18_23.rep



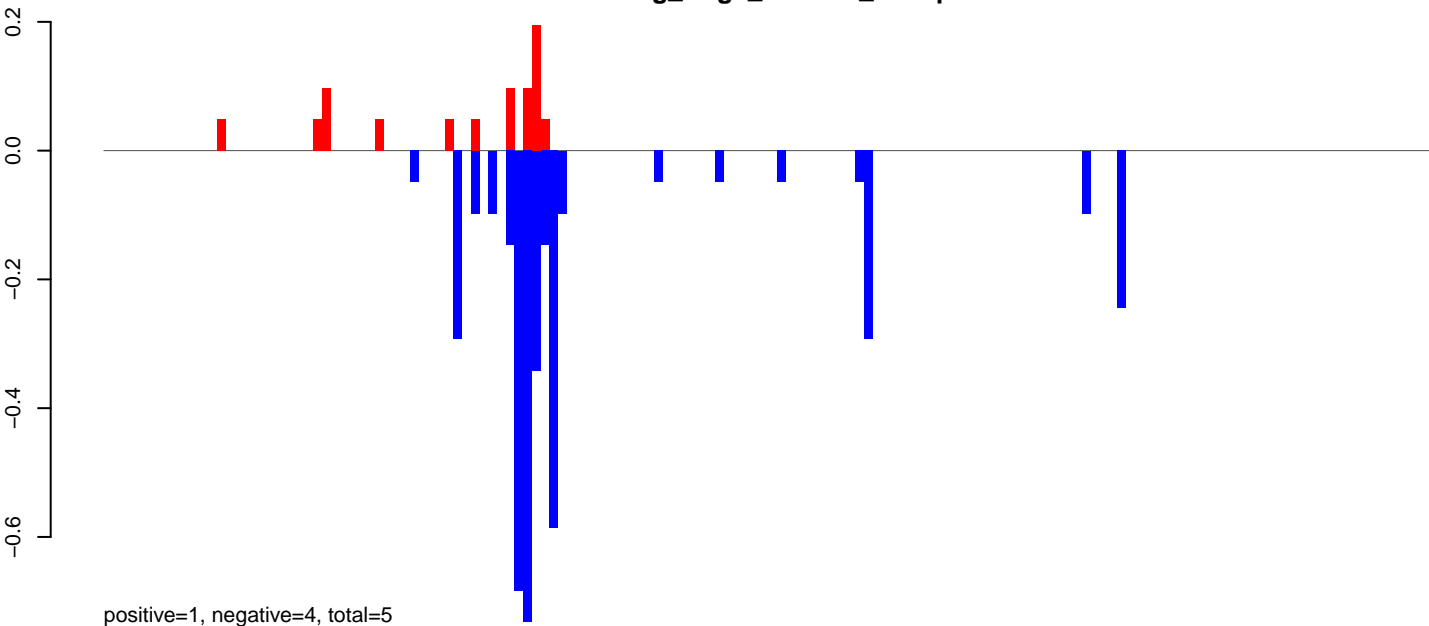
AeAeg_Aag2_cells.24_35.rep



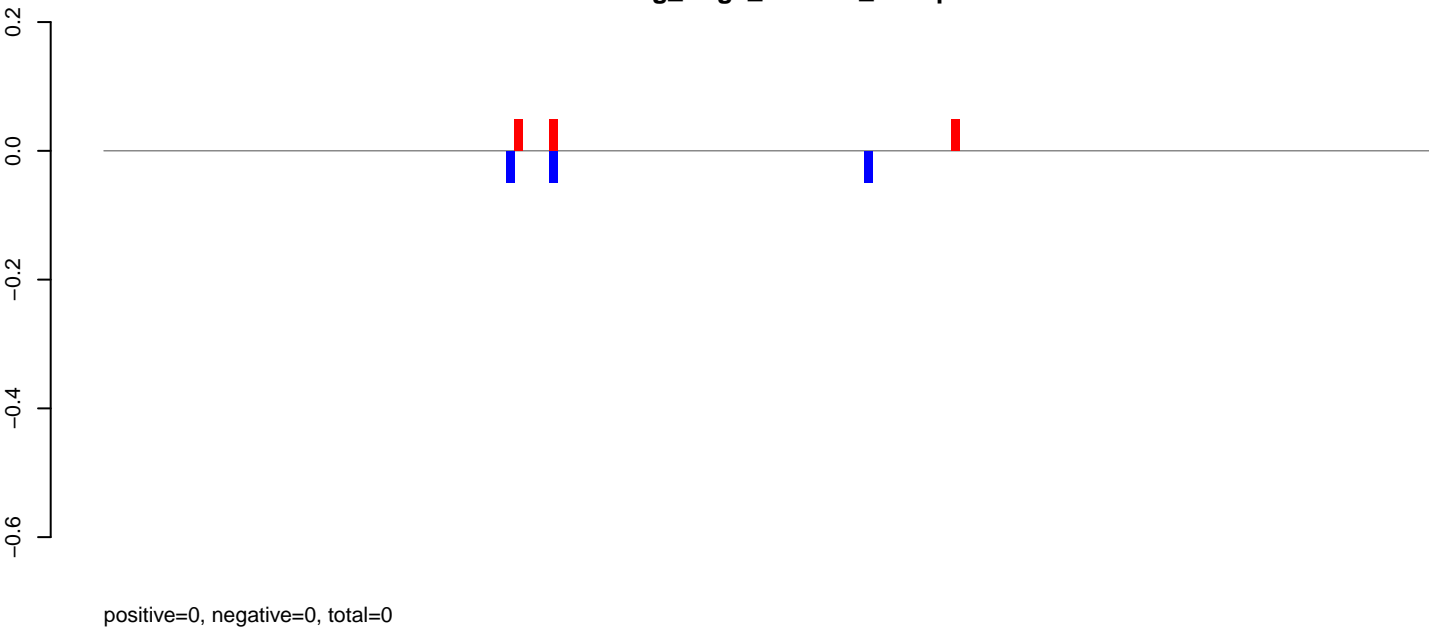
AeAeg_Aag2_cells.rep



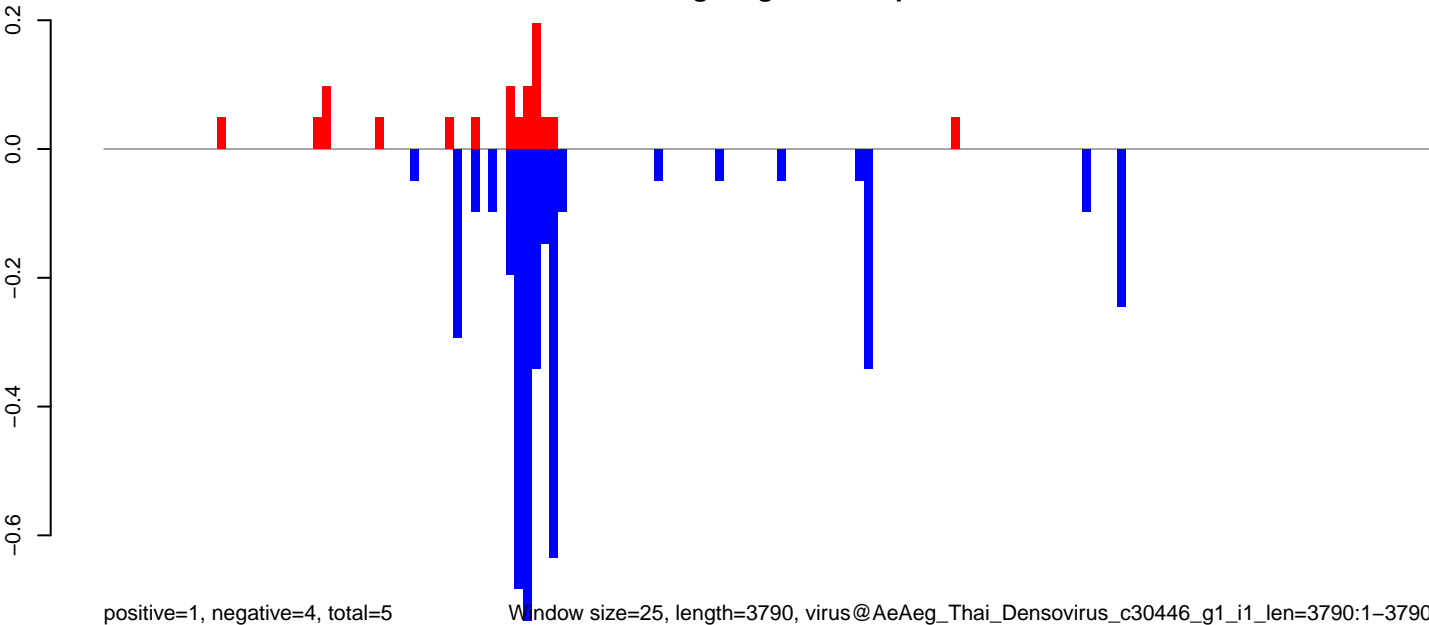
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

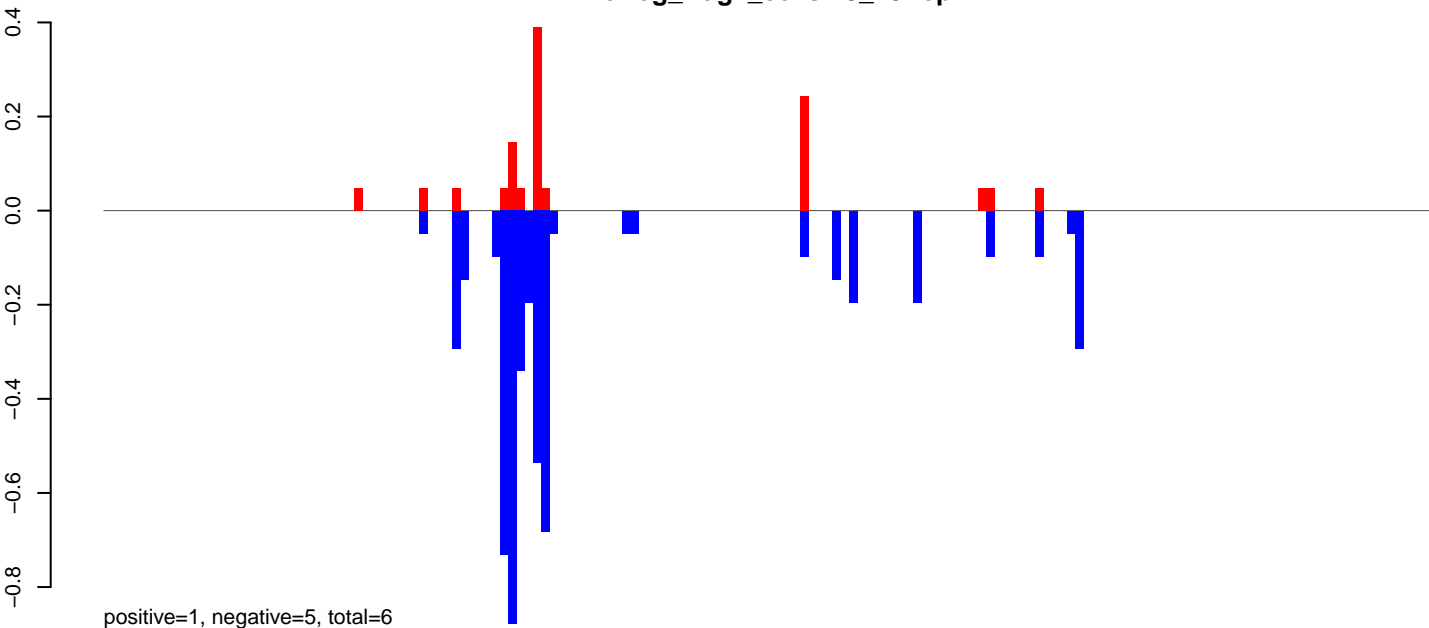


AeAeg_Aag2_cells.rep



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

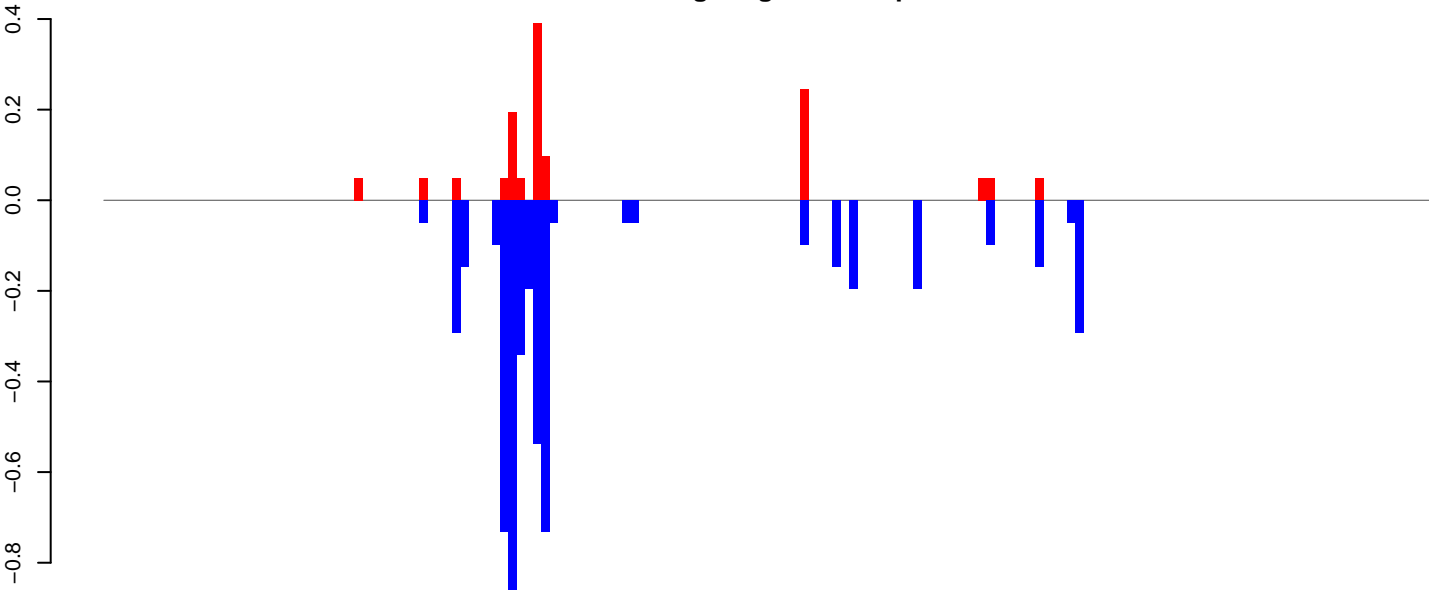
AeAeg_Aag2_cells.18_23.rep



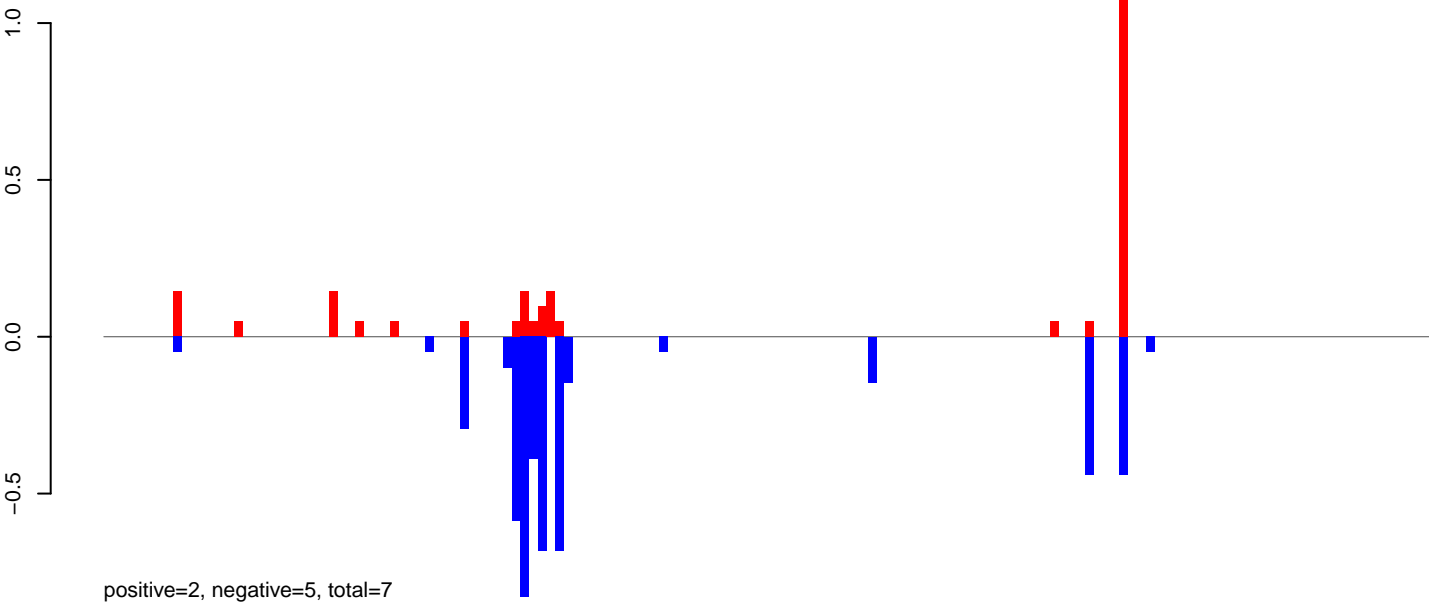
AeAeg_Aag2_cells.24_35.rep



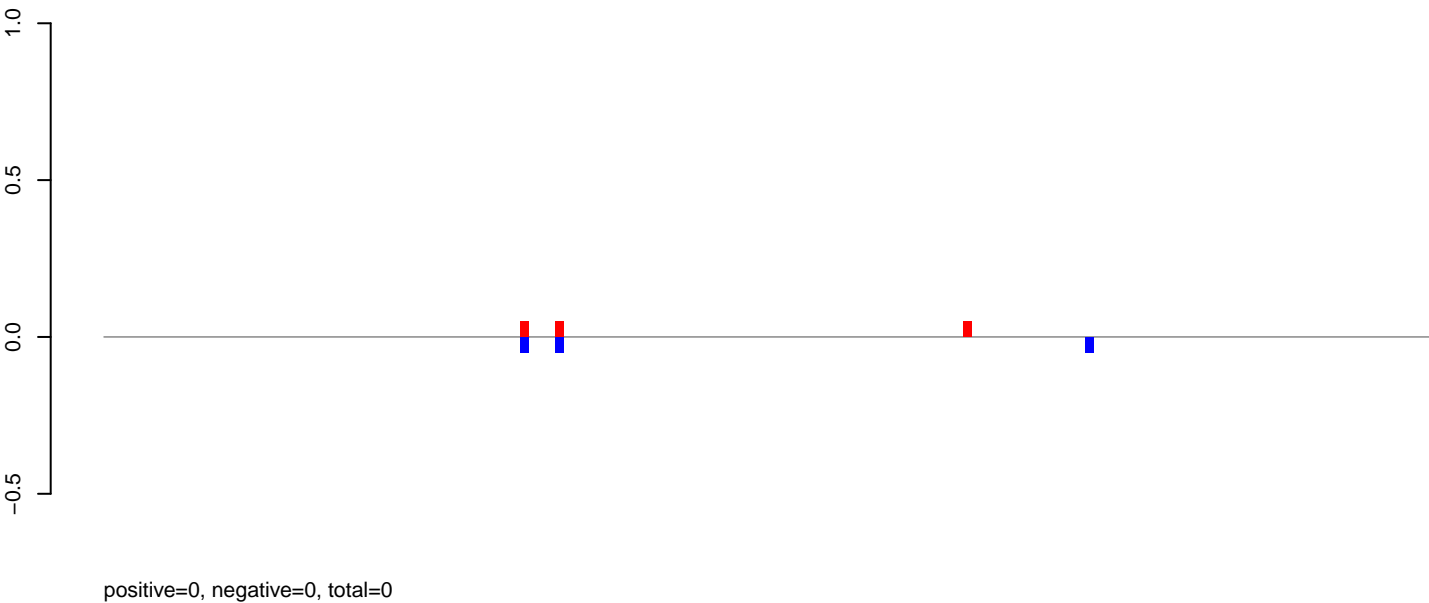
AeAeg_Aag2_cells.rep



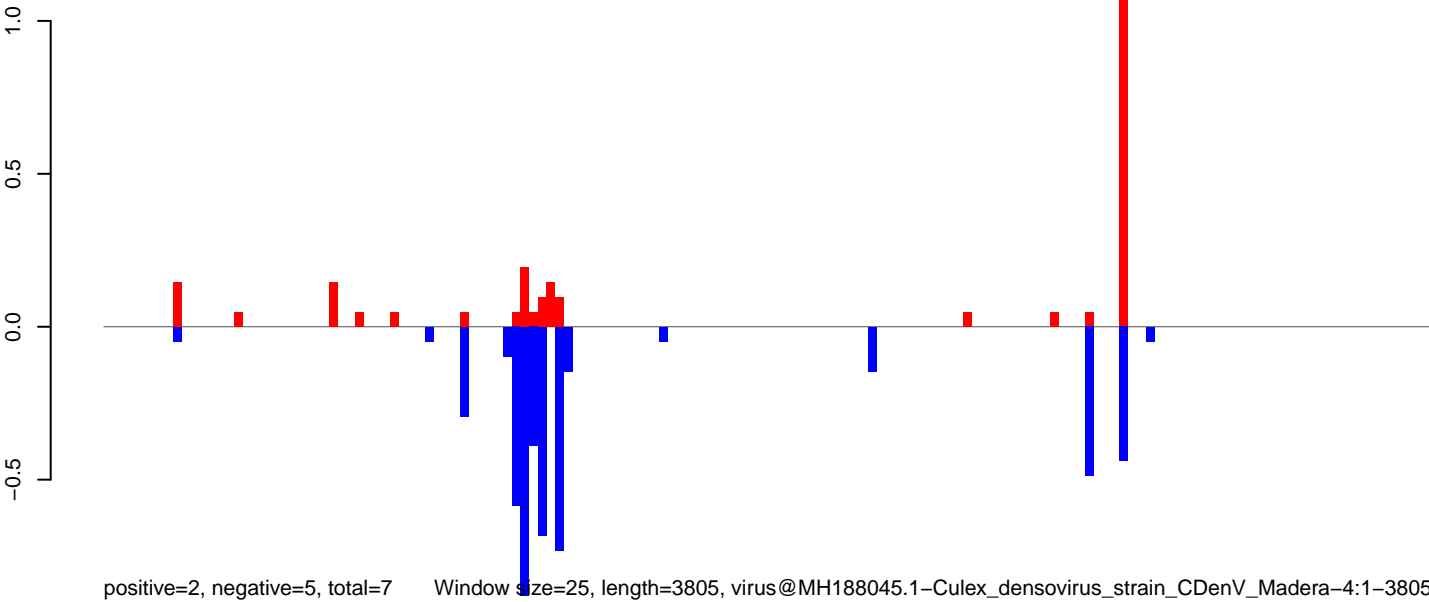
AeAeg_Aag2_cells.18_23.rep



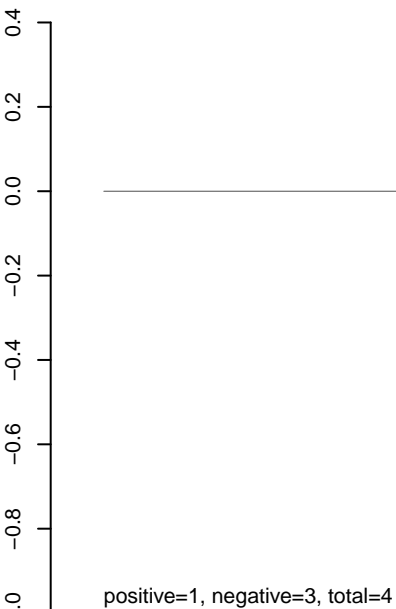
AeAeg_Aag2_cells.24_35.rep



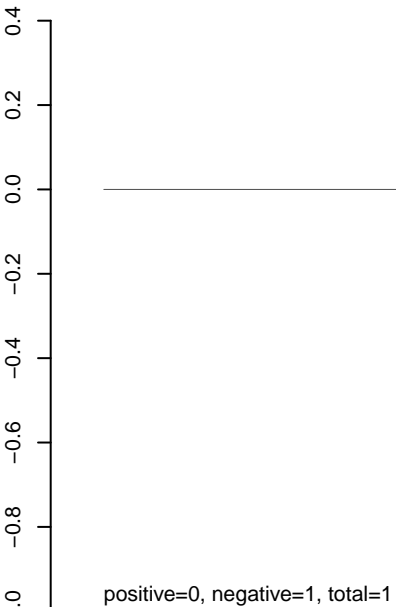
AeAeg_Aag2_cells.rep



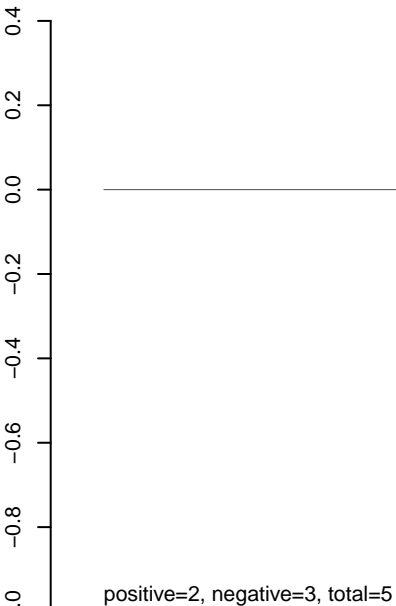
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

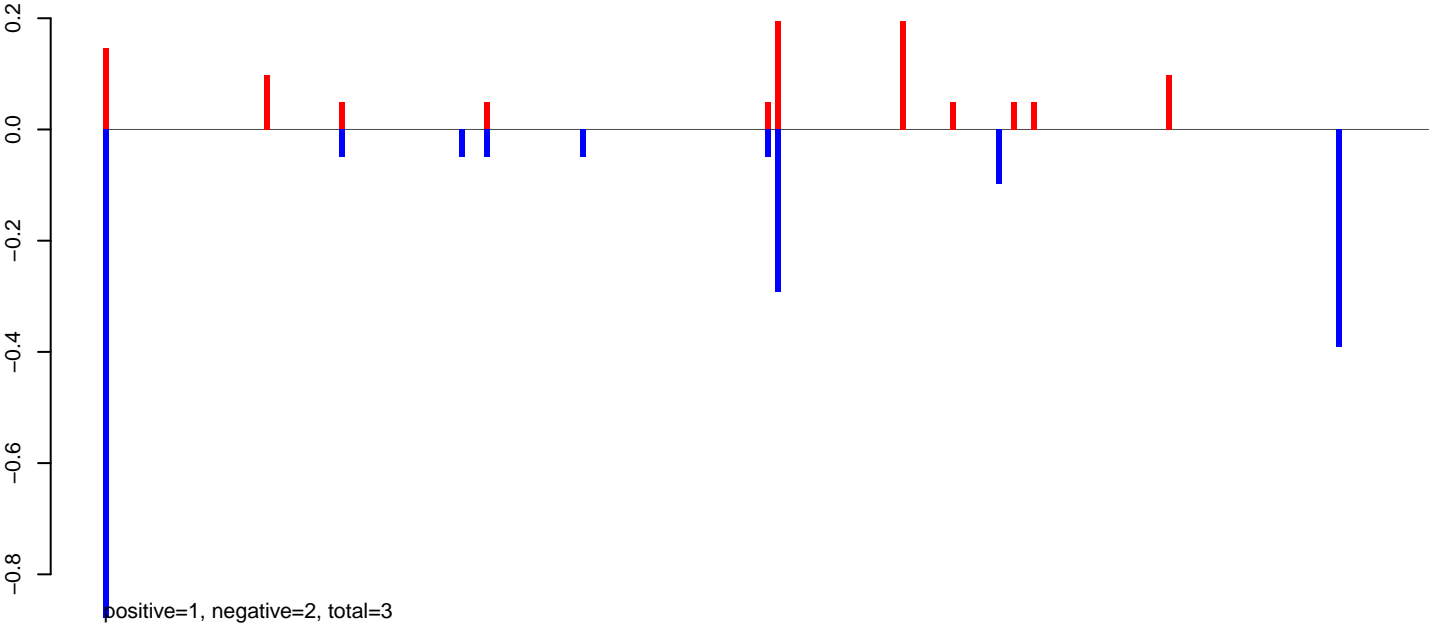


AeAeg_Aag2_cells.rep

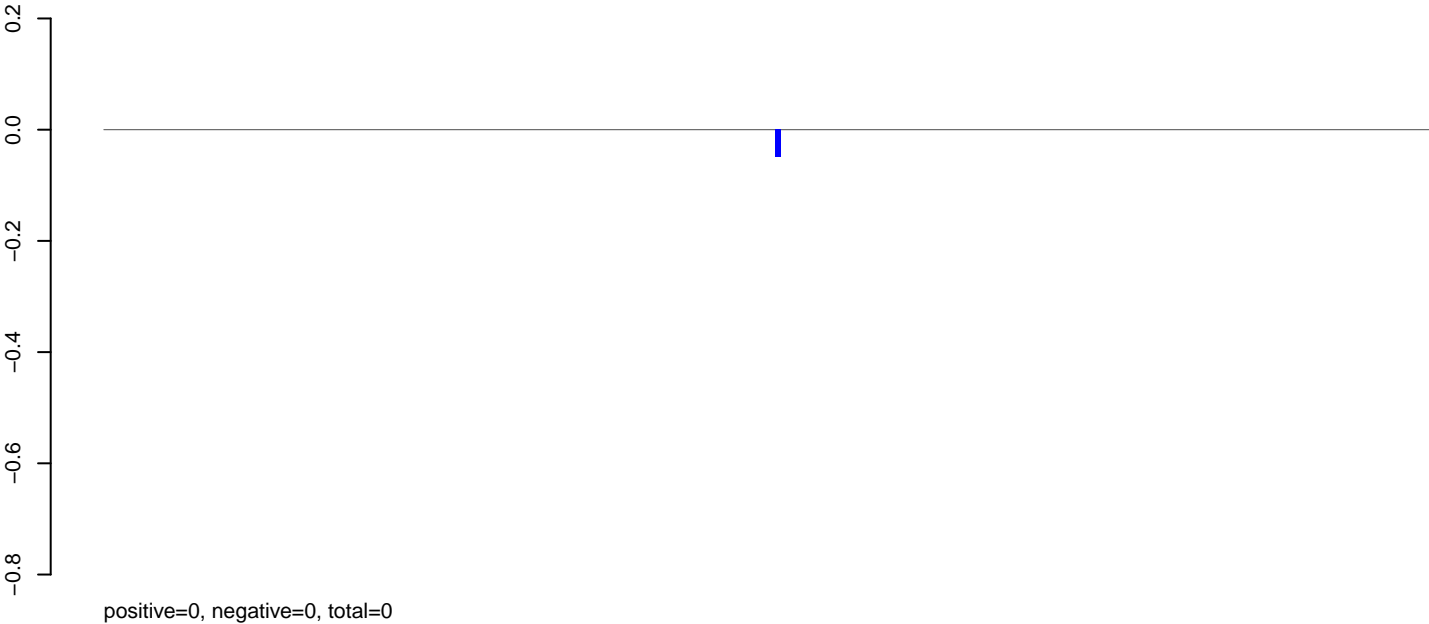


Window size=25, length=11375, virus@NC_005064.1-Kamiti_River_virus:1-11375

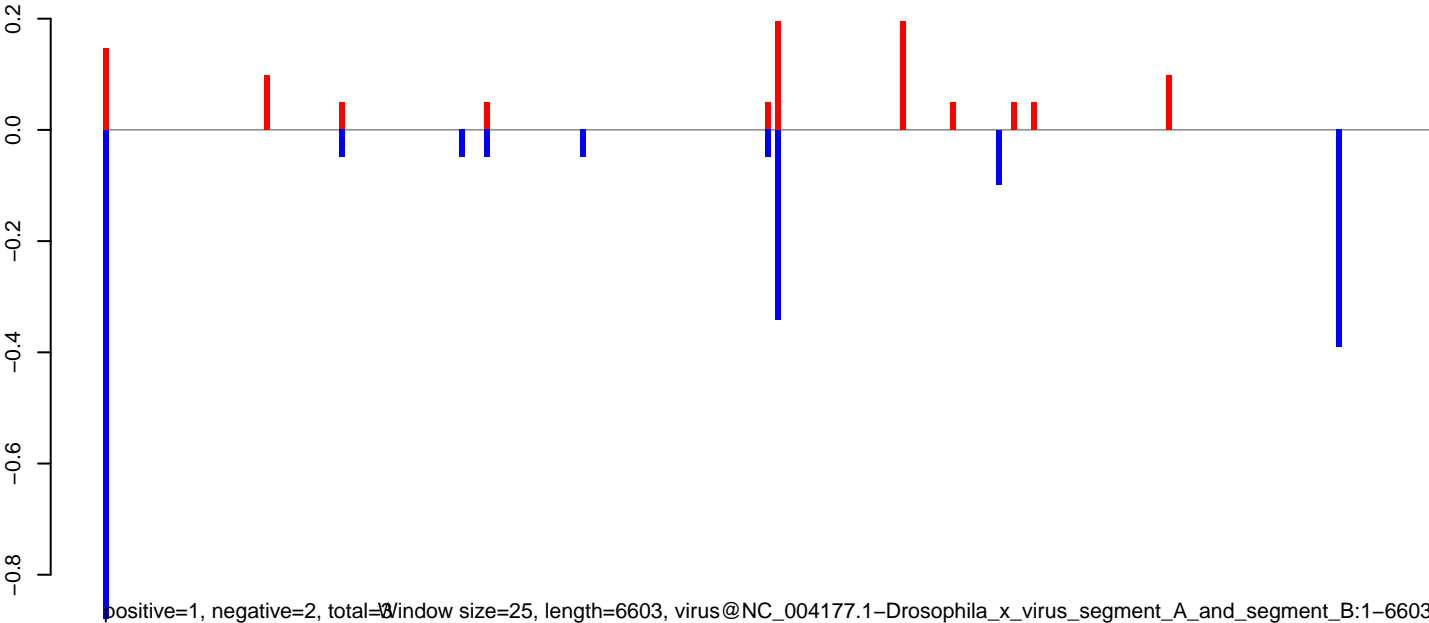
AeAeg_Aag2_cells.18_23.rep



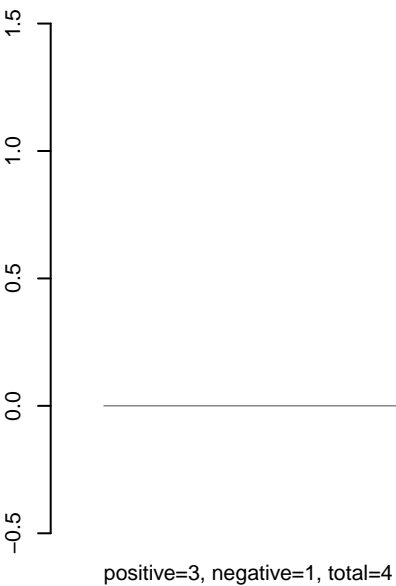
AeAeg_Aag2_cells.24_35.rep



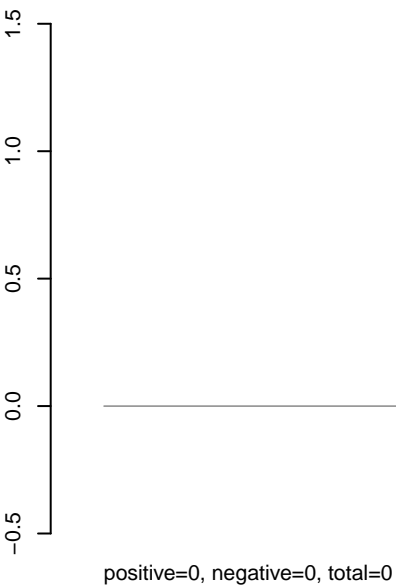
AeAeg_Aag2_cells.rep



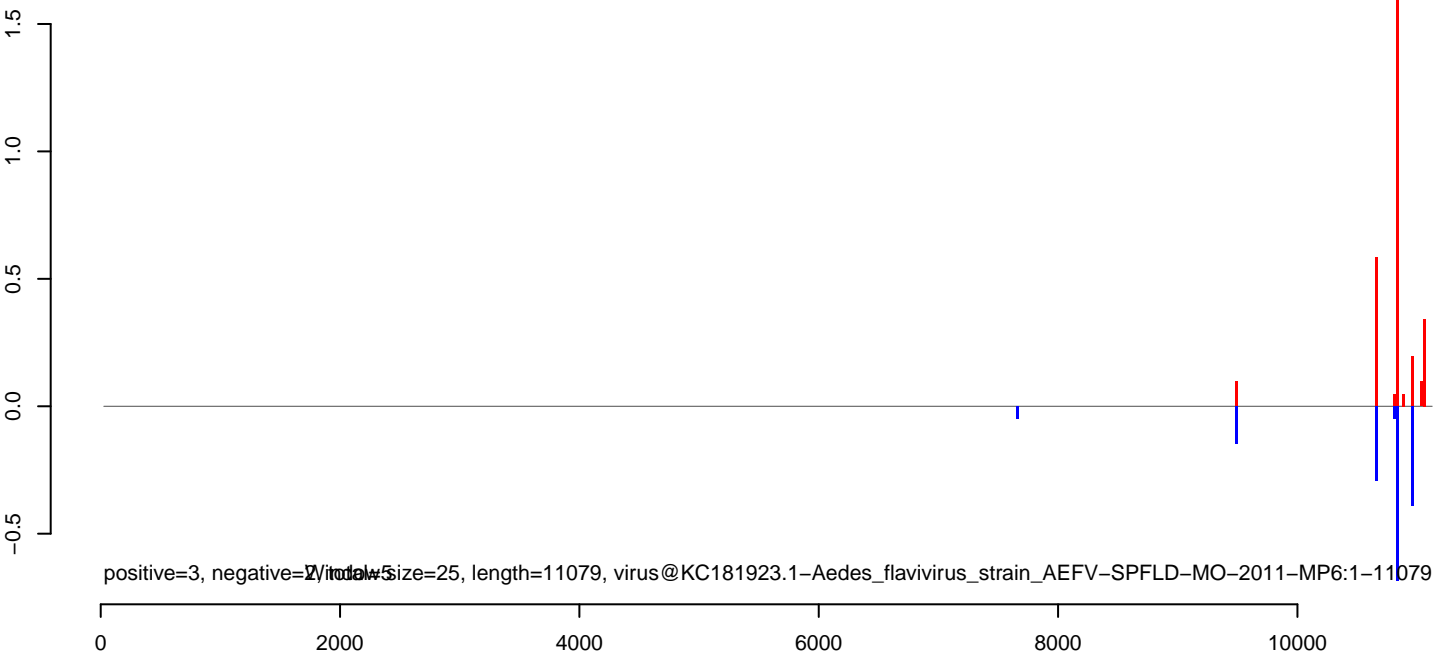
AeAeg_Aag2_cells.18_23.rep



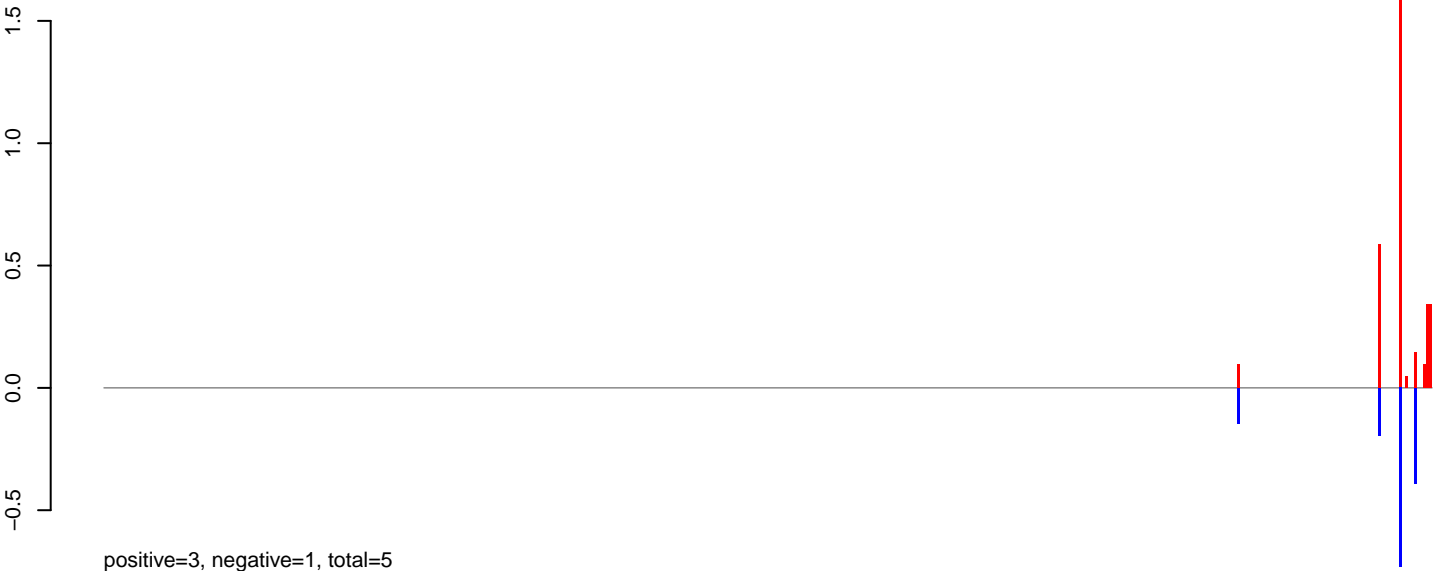
AeAeg_Aag2_cells.24_35.rep



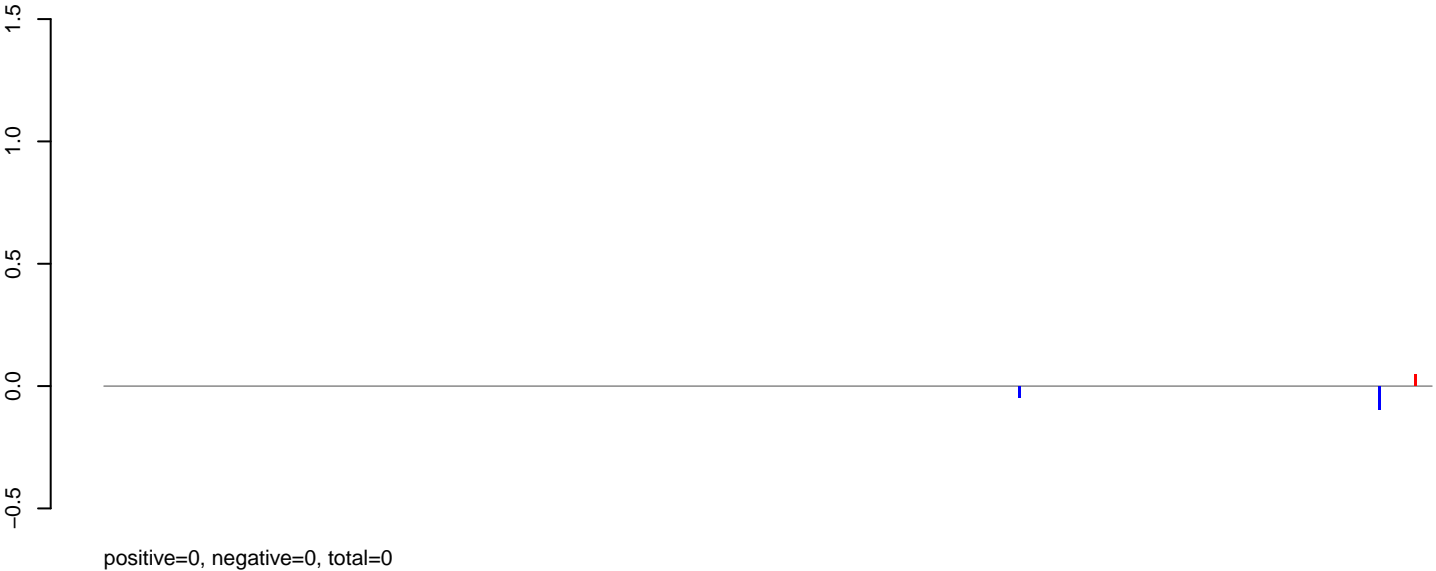
AeAeg_Aag2_cells.rep



AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

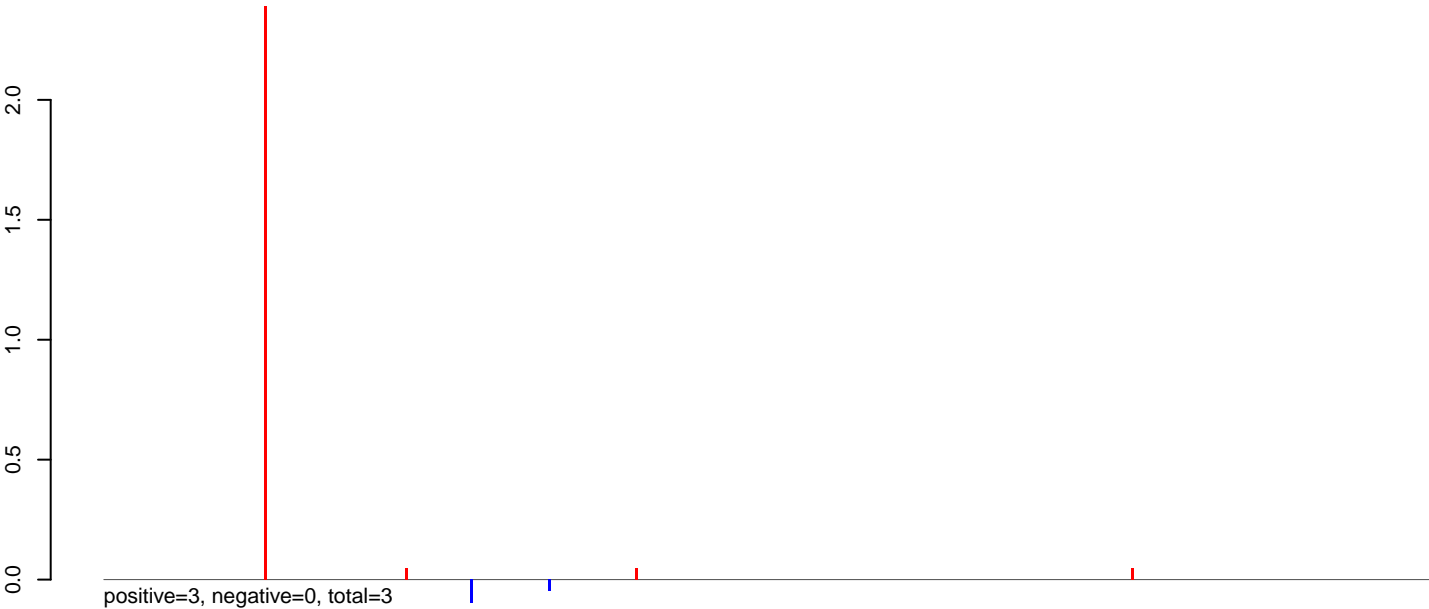


AeAeg_Aag2_cells.rep

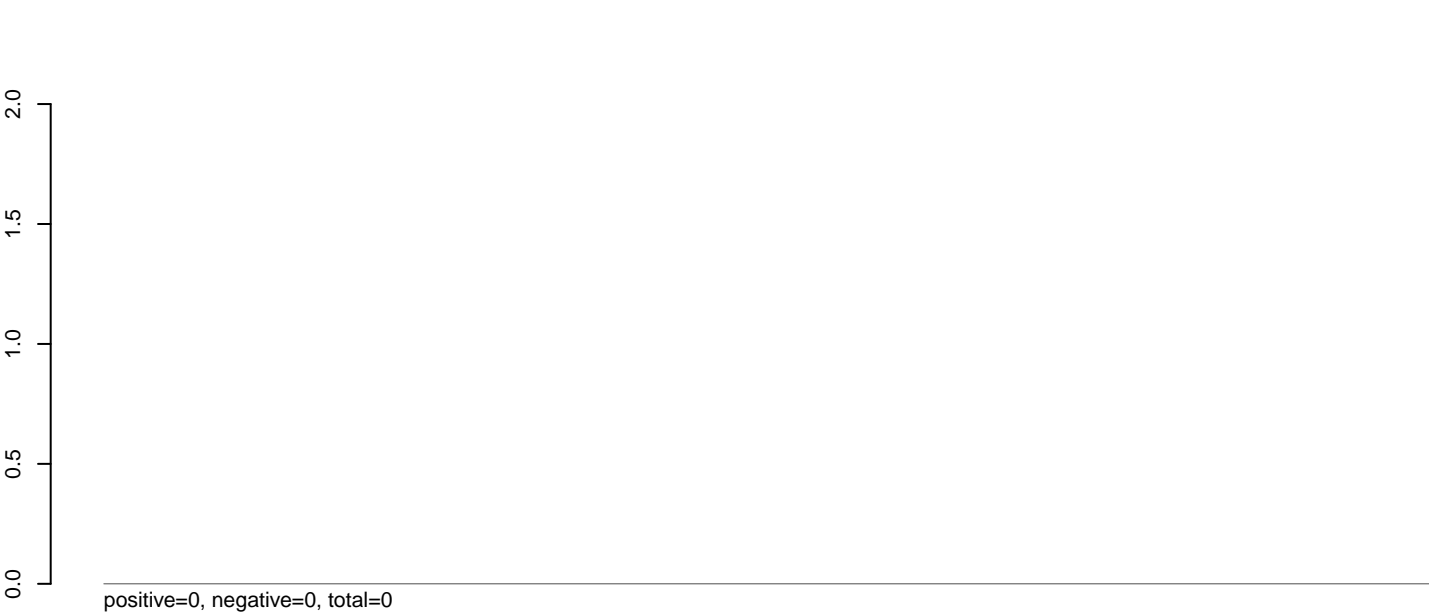


0 2000 4000 6000 8000 10000

AeAeg_Aag2_cells.18_23.rep



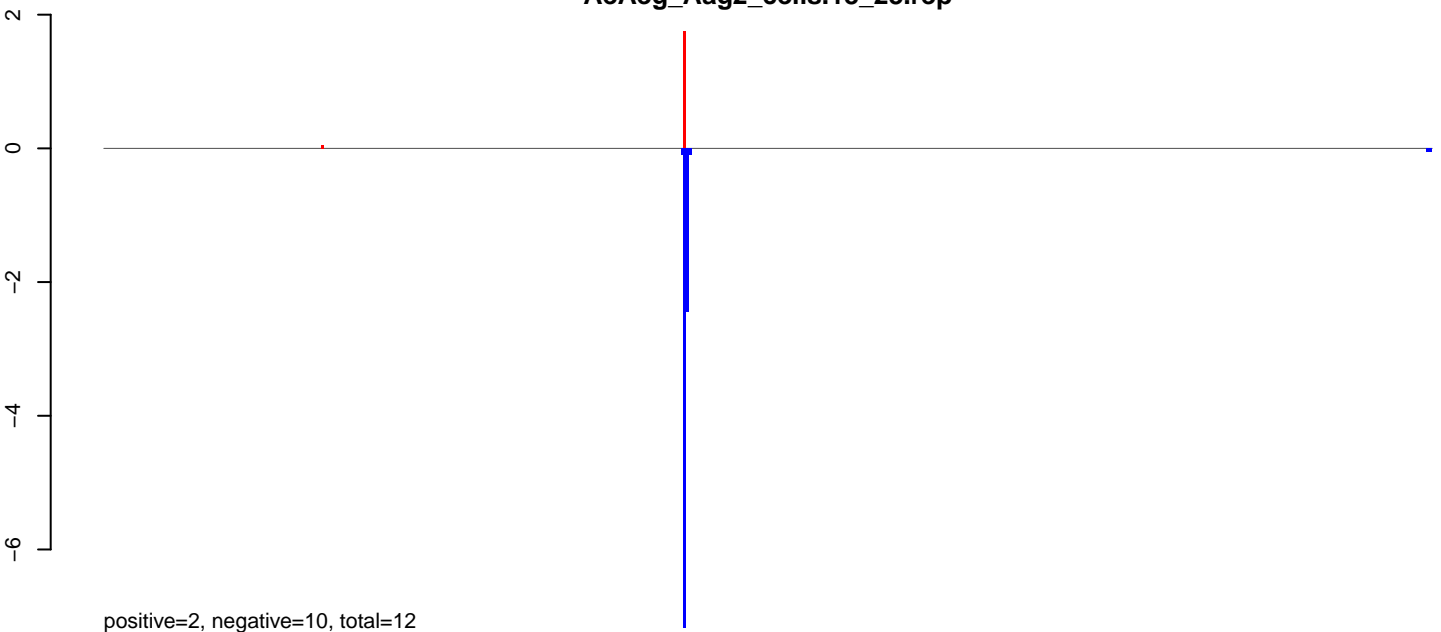
AeAeg_Aag2_cells.24_35.rep



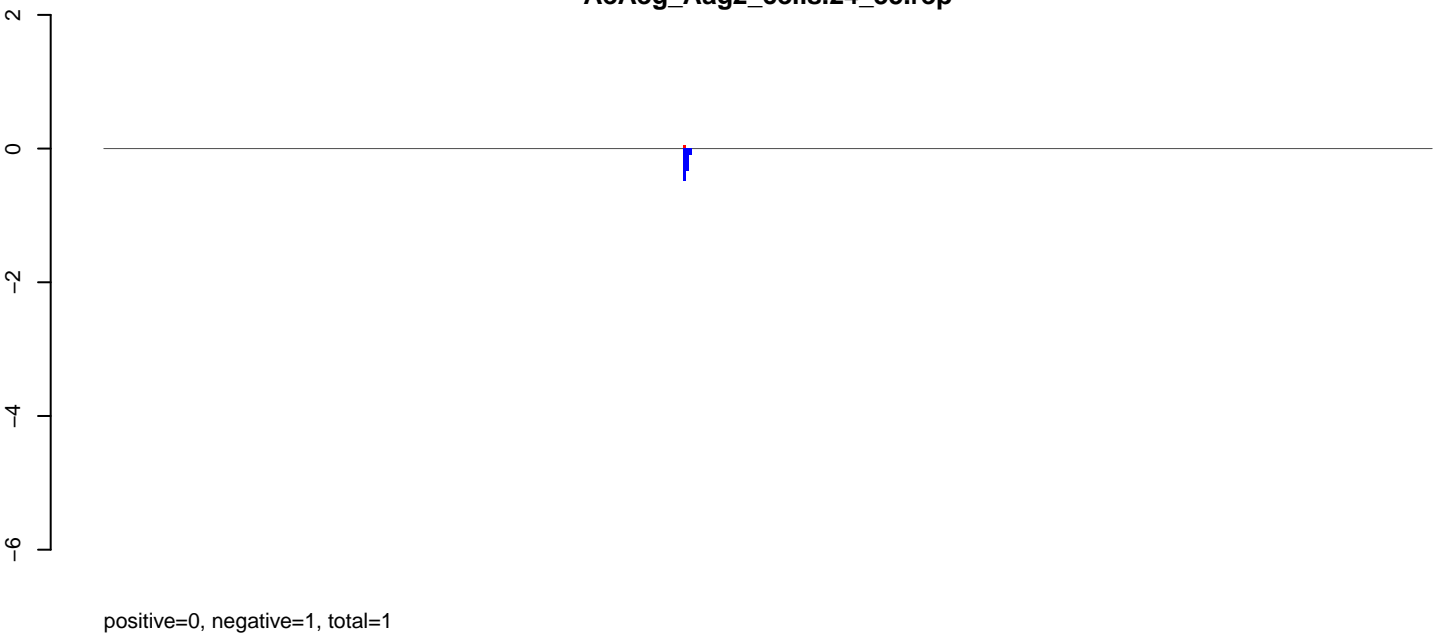
AeAeg_Aag2_cells.rep



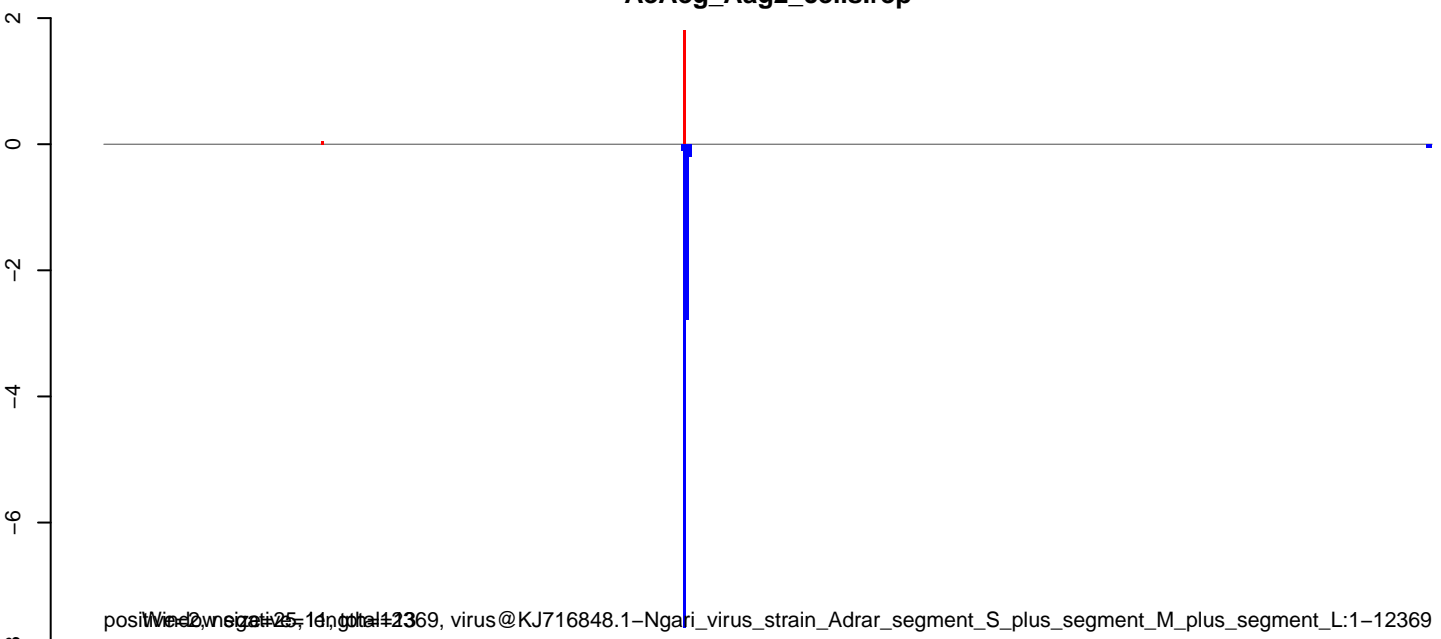
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



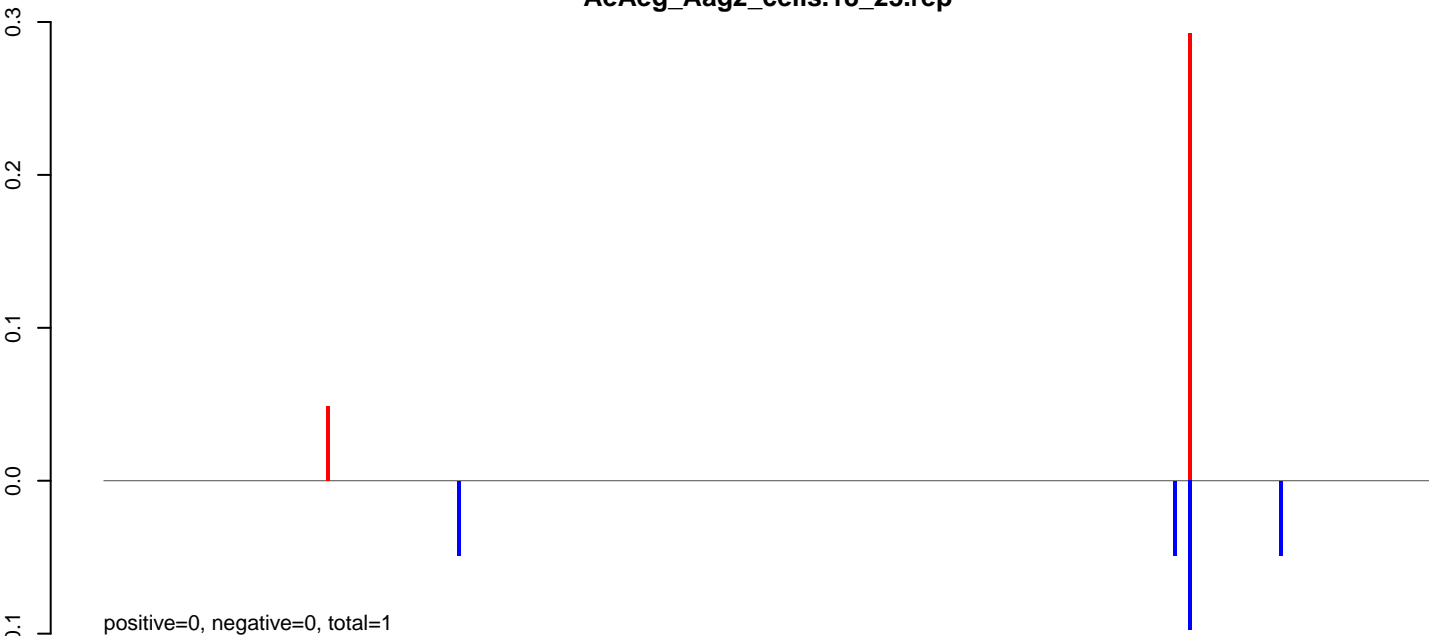
AeAeg_Aag2_cells.rep



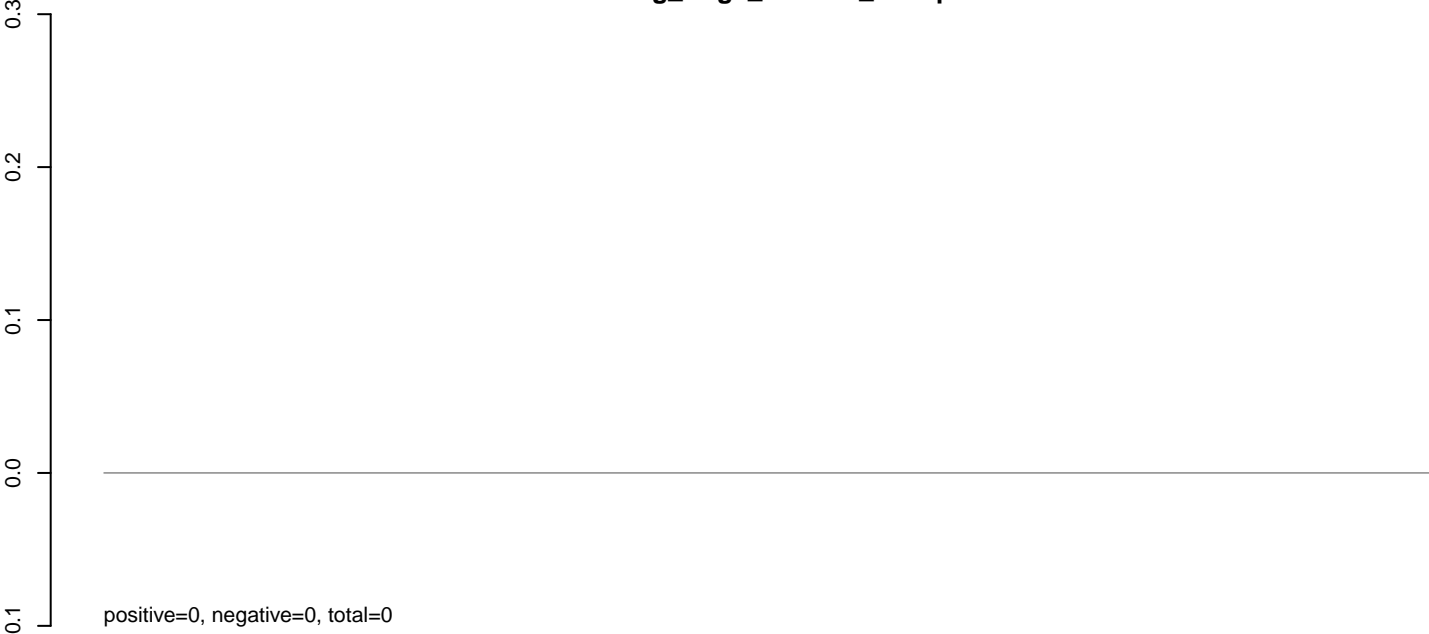
Wendy, size=25, length=2369, virus@KJ716848.1-Ngari_virus_strain_Adrar_segment_S_plus_segment_M_plus_segment_L:1-12369

0 2000 4000 6000 8000 10000 12000

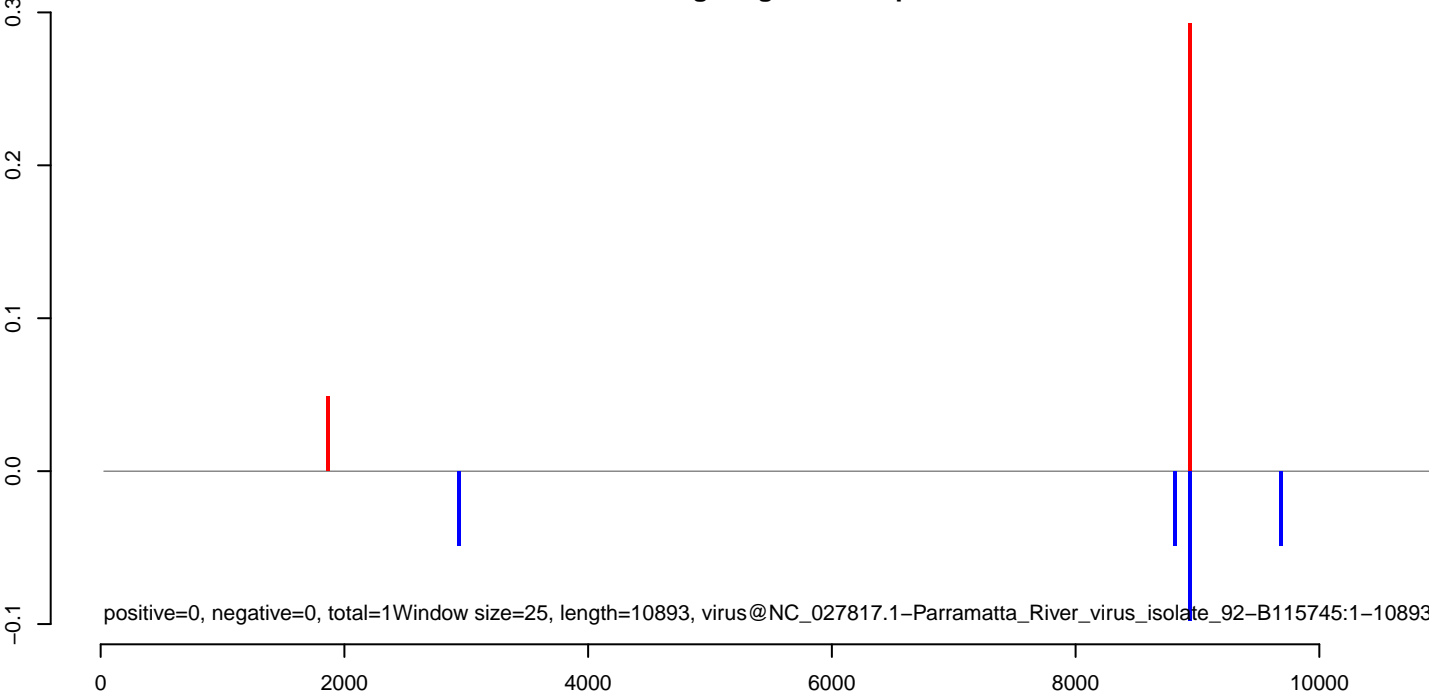
AeAeg_Aag2_cells.18_23.rep



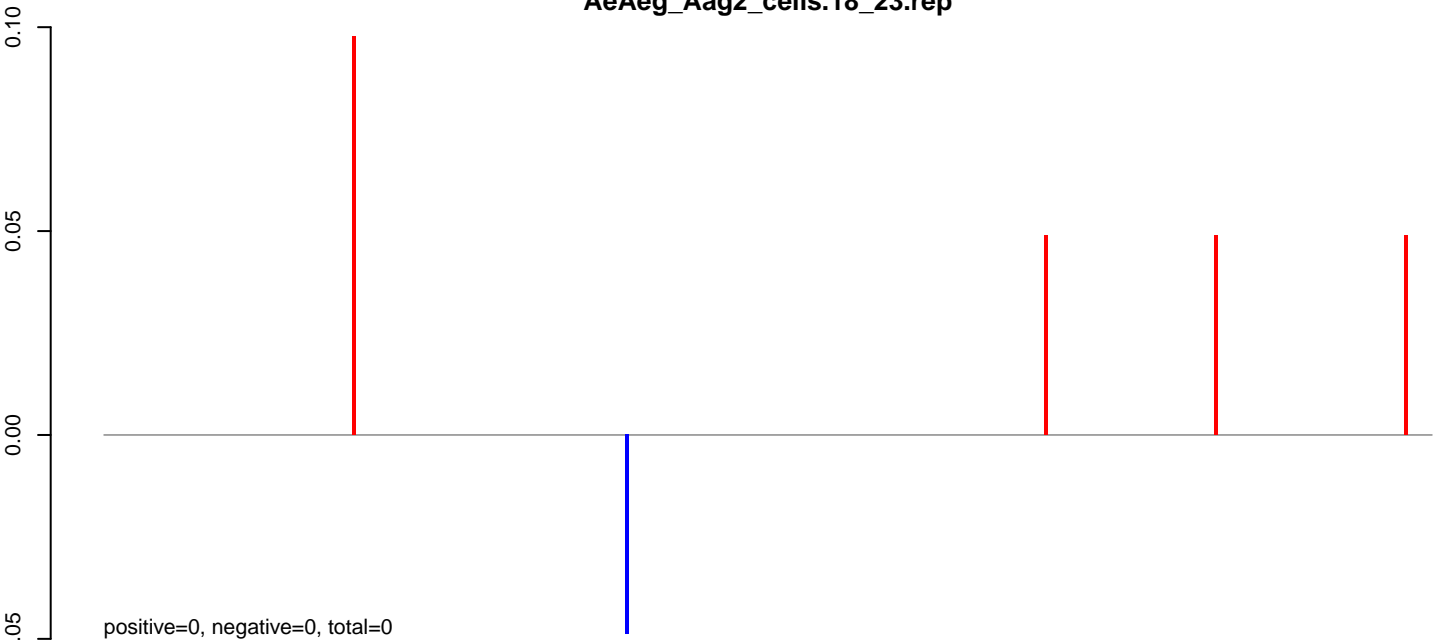
AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep

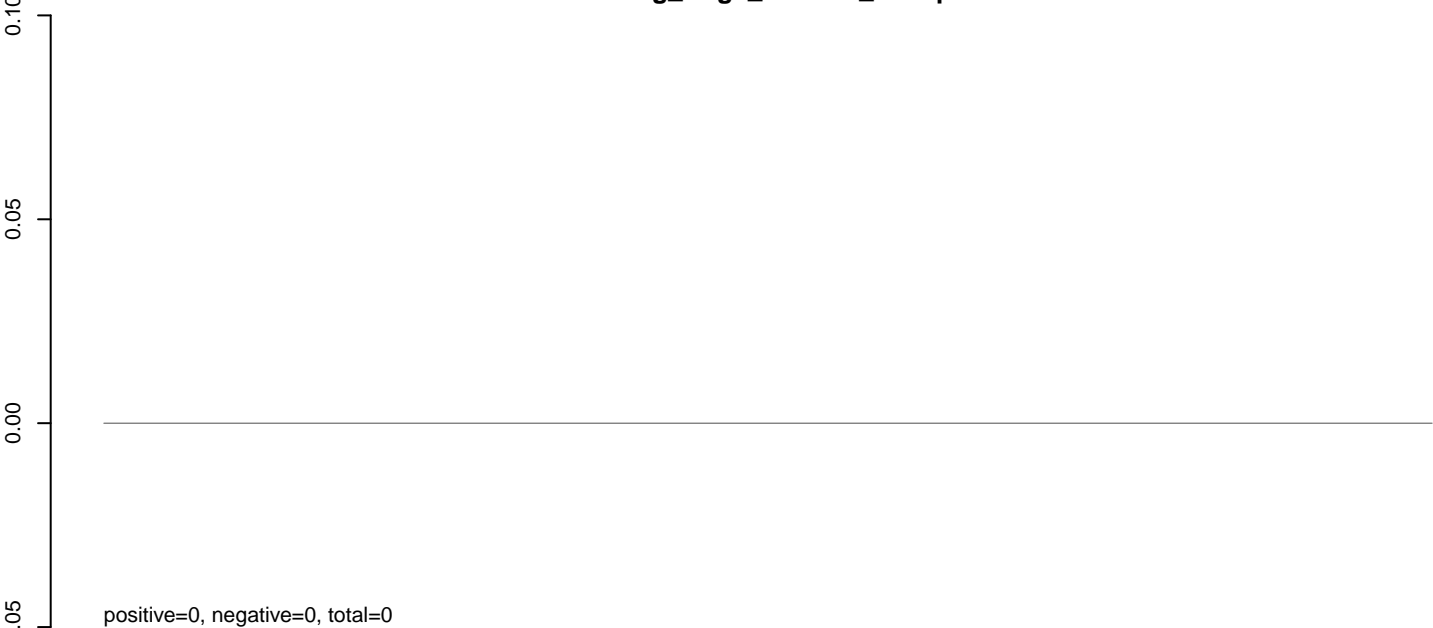


AeAeg_Aag2_cells.18_23.rep



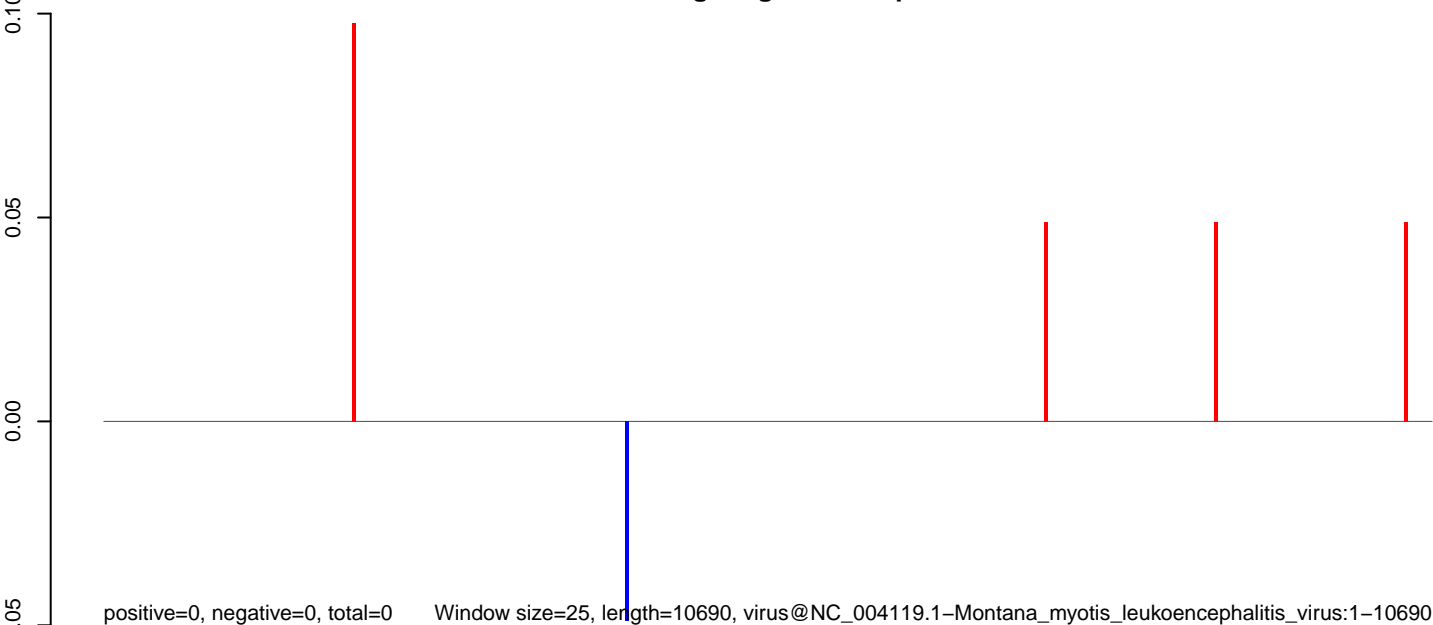
positive=0, negative=0, total=0

AeAeg_Aag2_cells.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_cells.rep

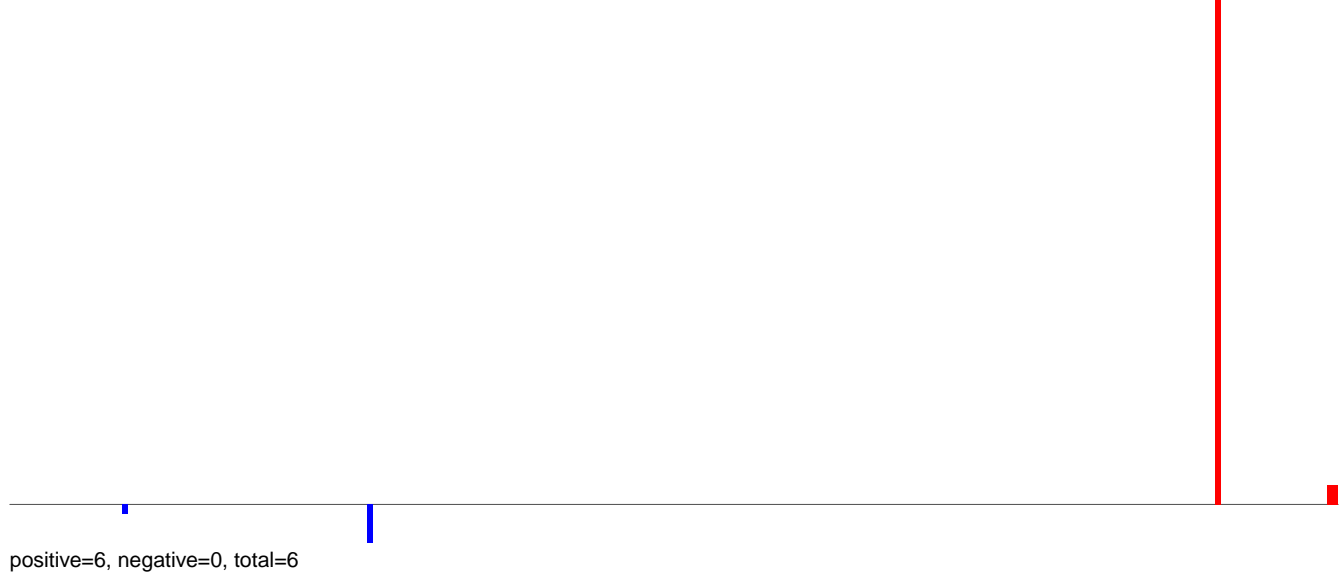


positive=0, negative=0, total=0

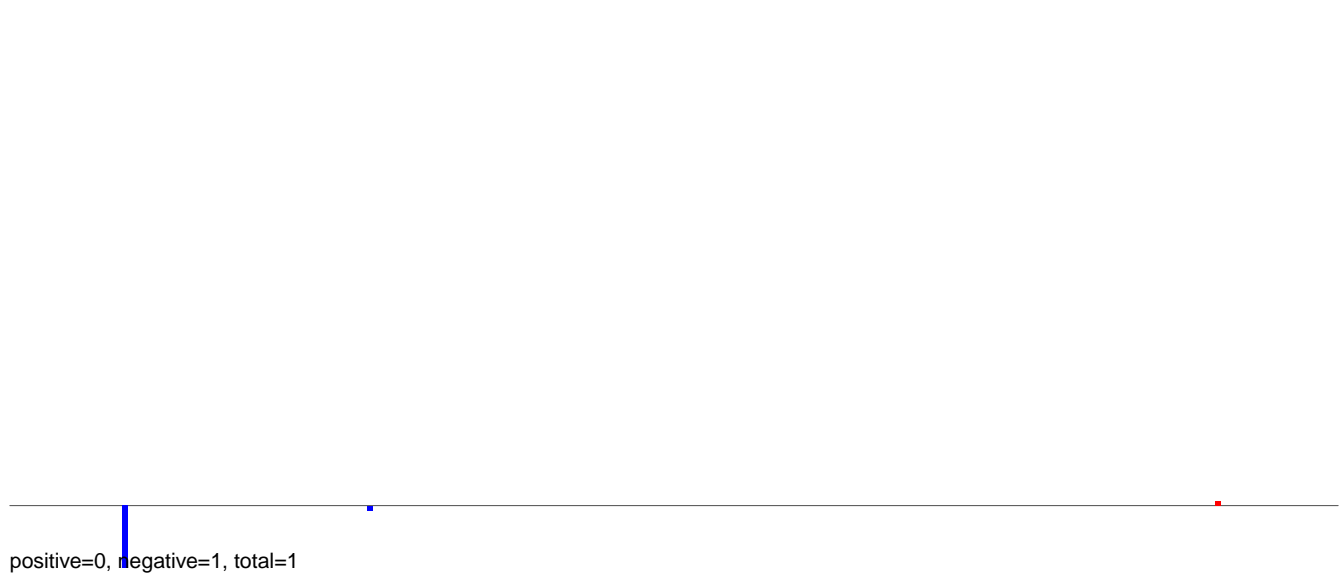
Window size=25, length=10690, virus@NC_004119.1-Montana_myotis_leukoencephalitis_virus:1-10690

0 2000 4000 6000 8000 10000

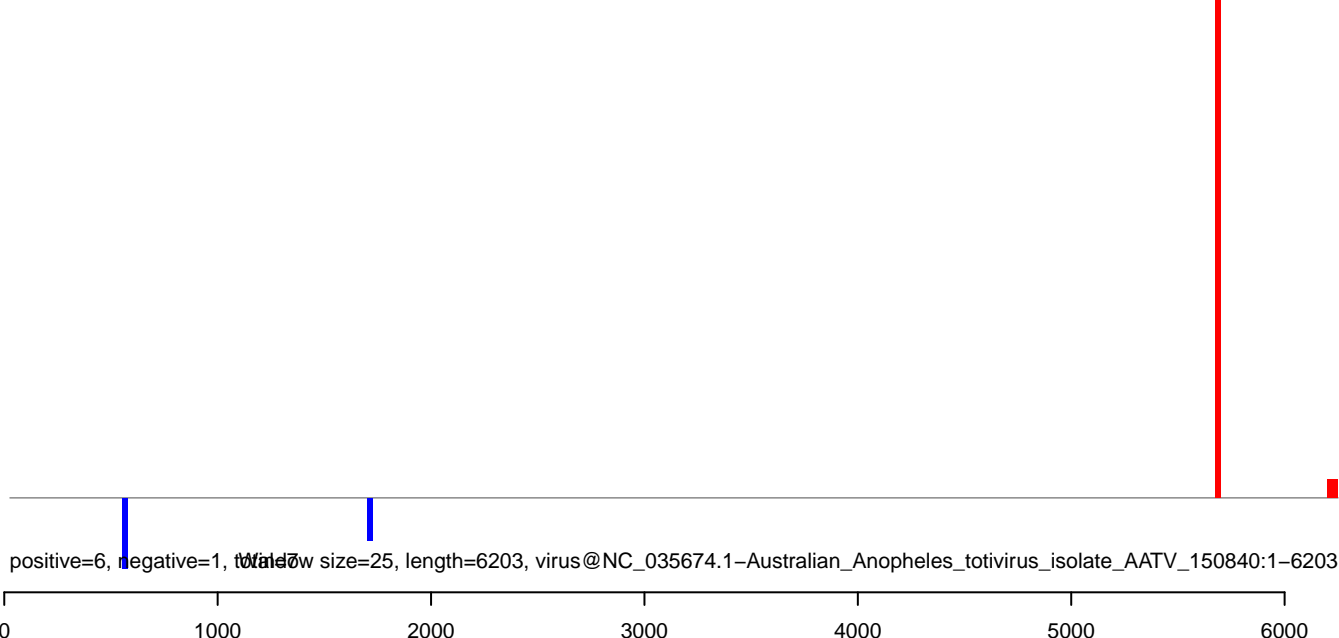
AeAeg_Aag2_cells.18_23.rep



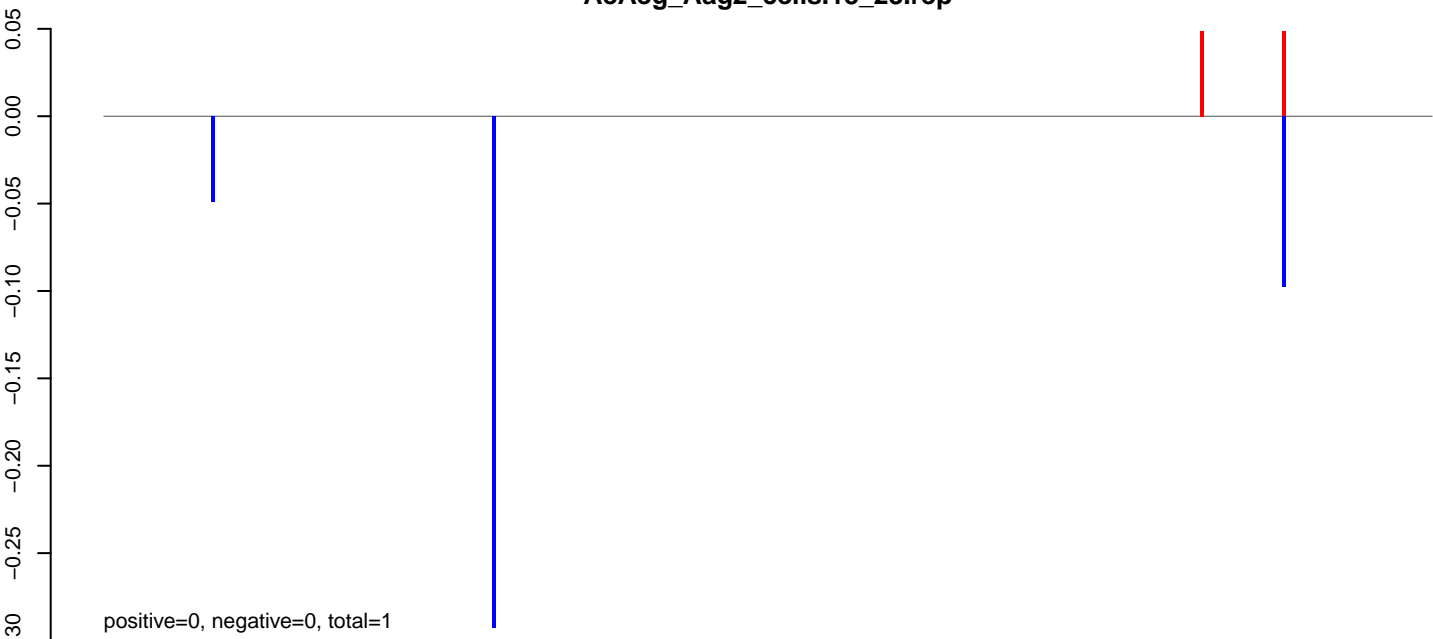
AeAeg_Aag2_cells.24_35.rep



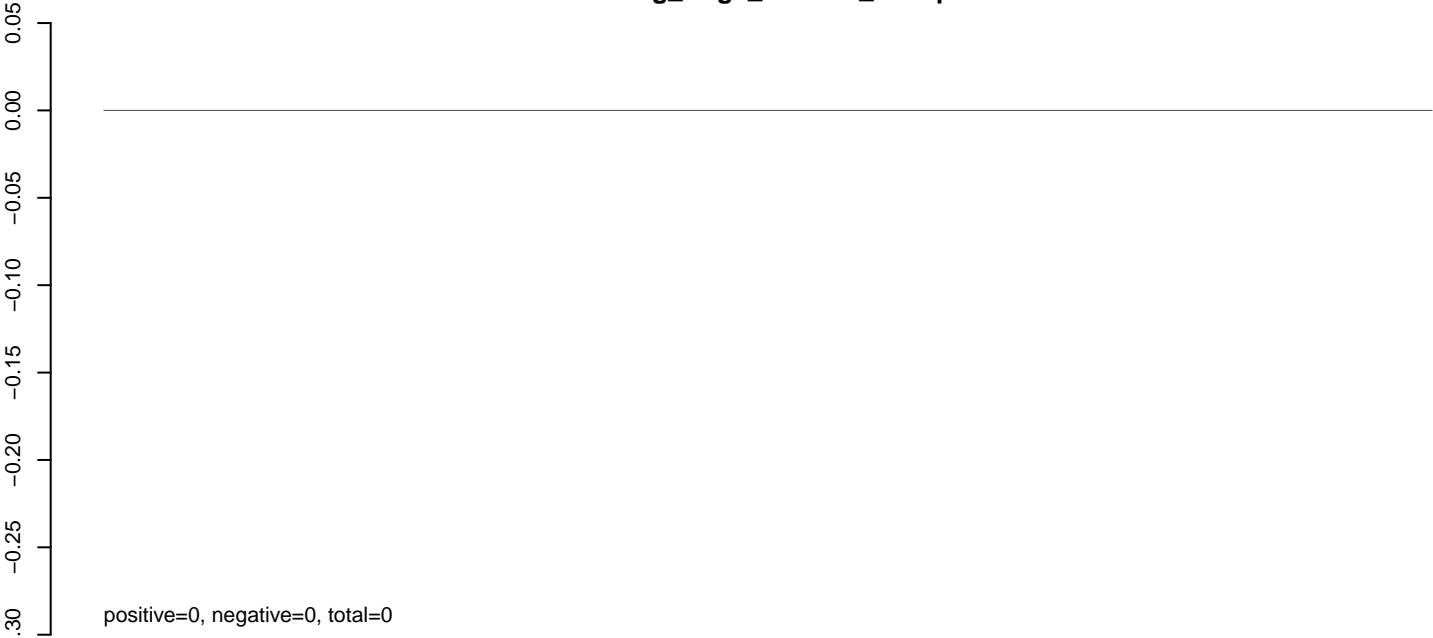
AeAeg_Aag2_cells.rep



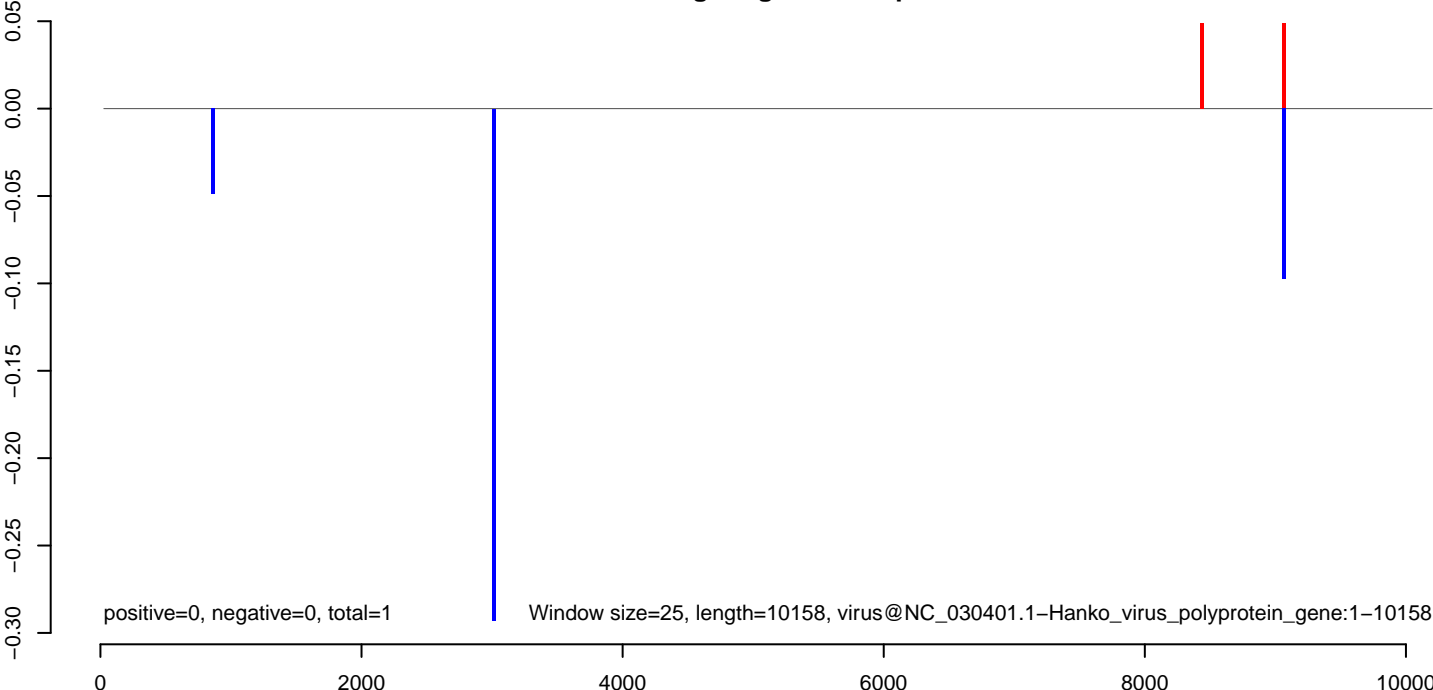
AeAeg_Aag2_cells.18_23.rep



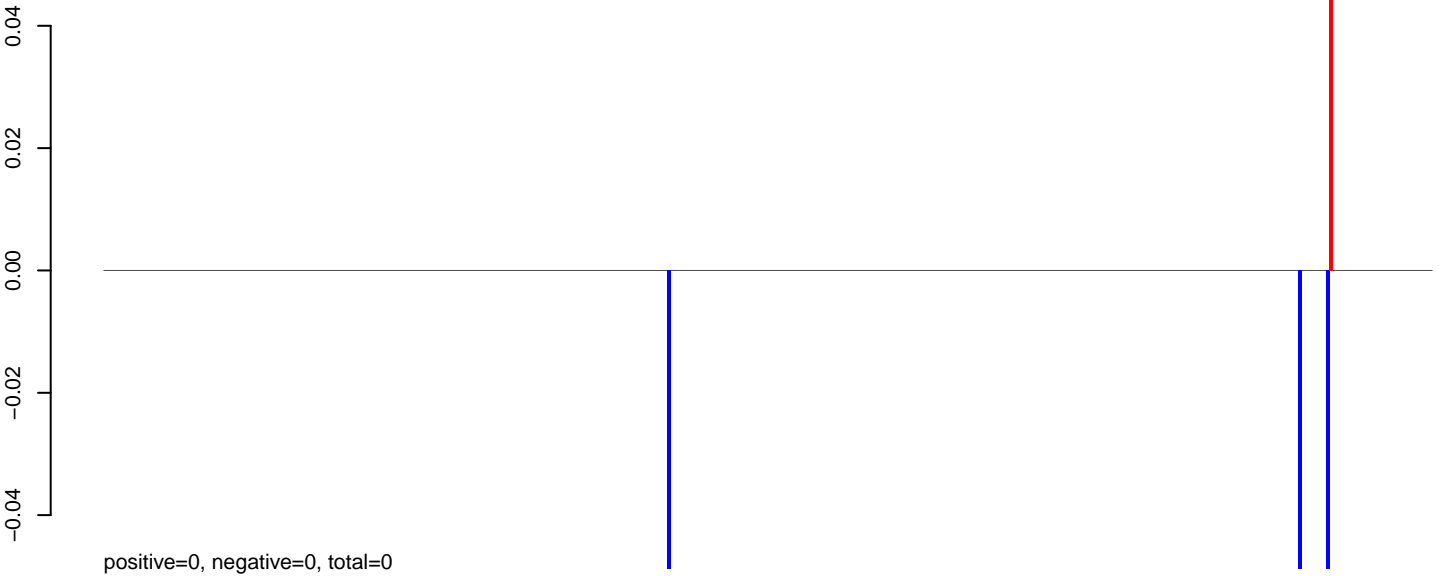
AeAeg_Aag2_cells.24_35.rep



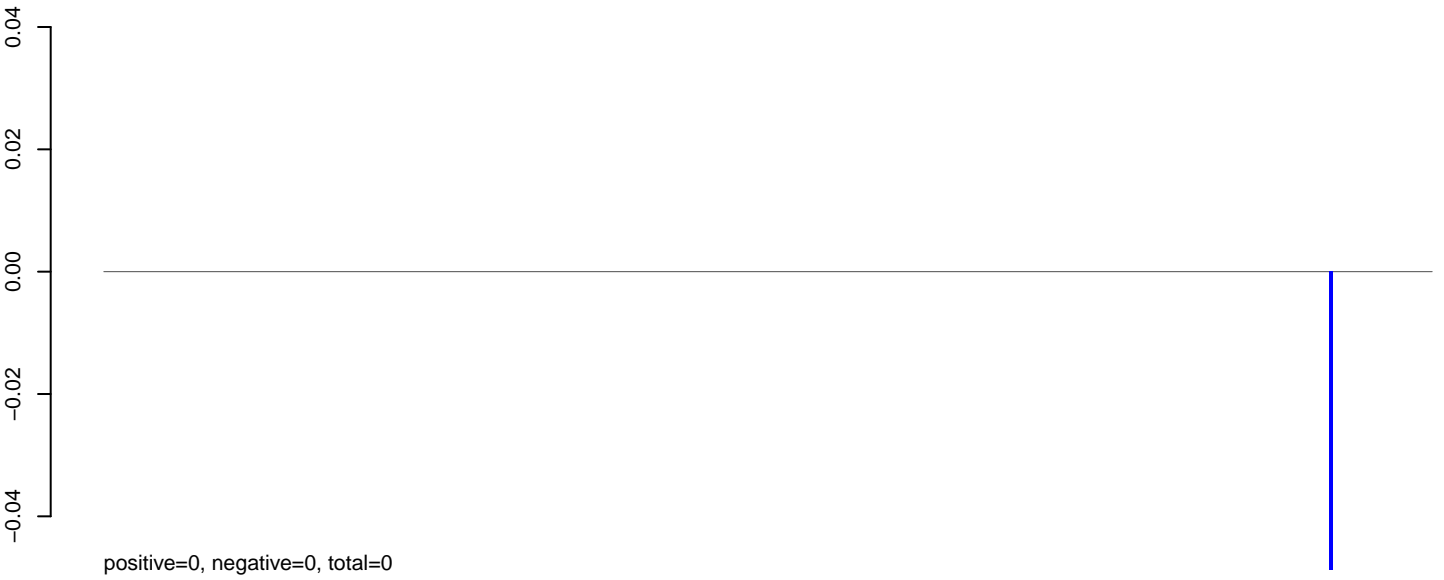
AeAeg_Aag2_cells.rep



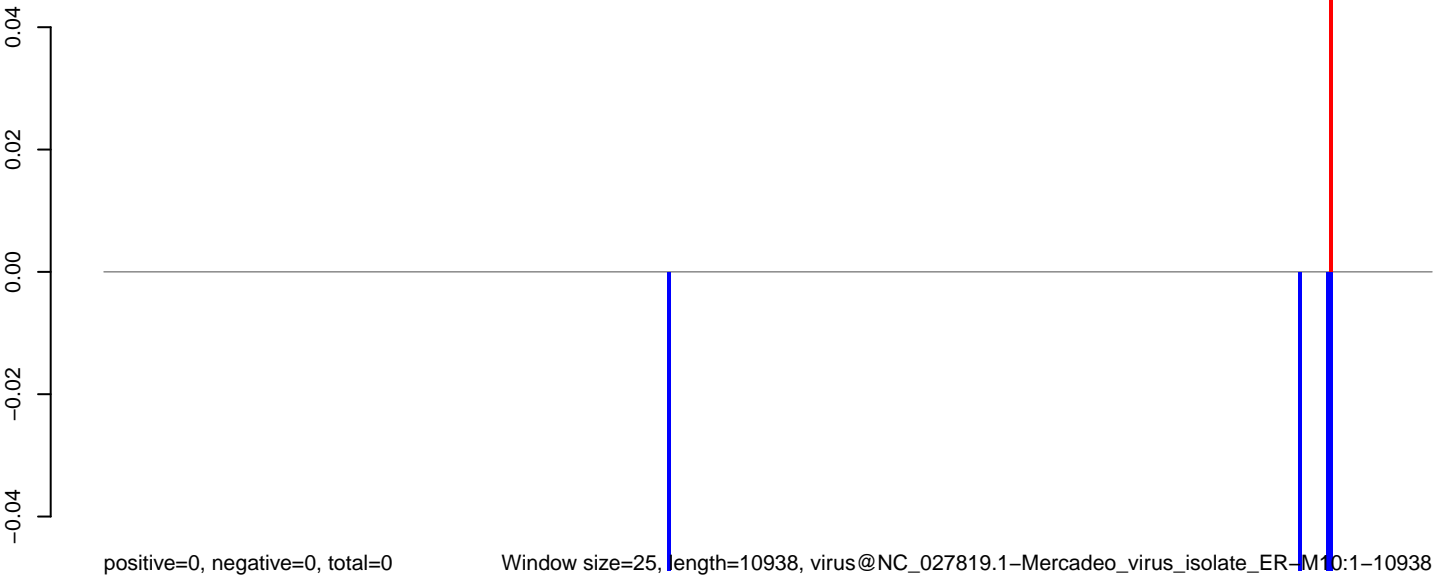
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep



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

0 2000 4000 6000 8000 10000

AeAeg_Aag2_cells.18_23.rep

0.04
0.02
0.00
-0.02
-0.04

positive=0, negative=0, total=0

AeAeg_Aag2_cells.24_35.rep

0.04
0.02
0.00
-0.02
-0.04

positive=0, negative=0, total=0

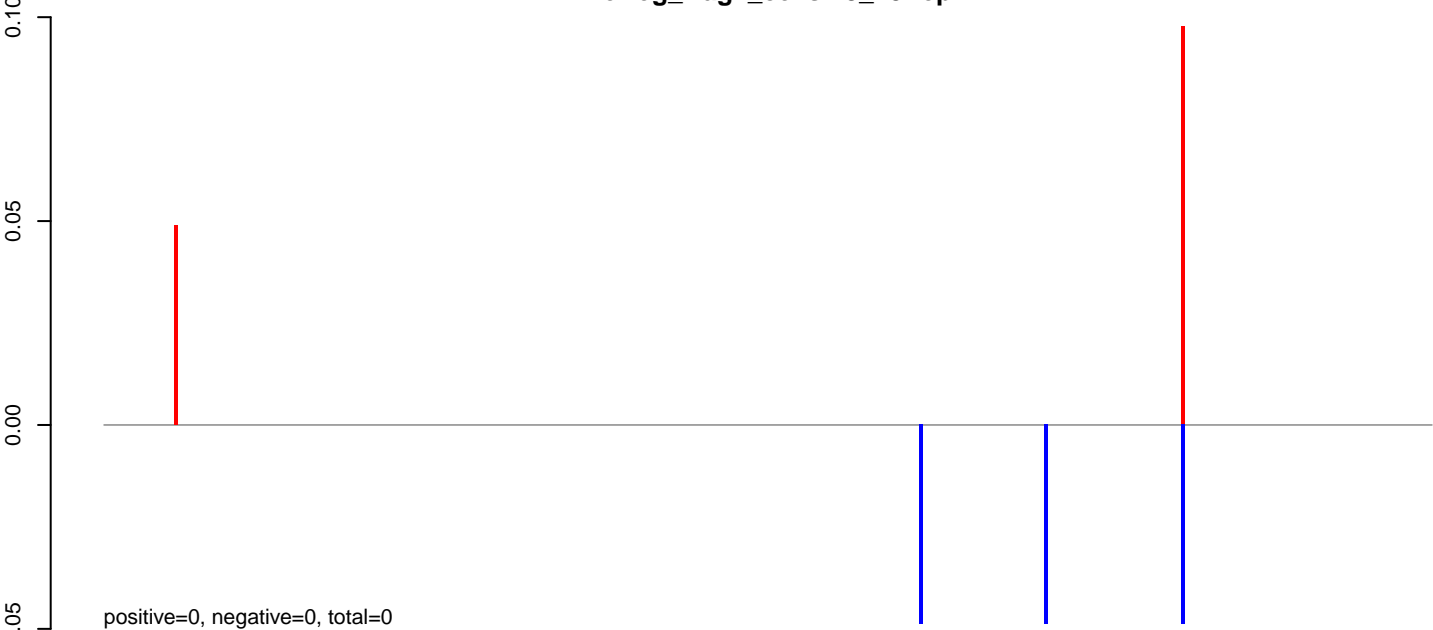
AeAeg_Aag2_cells.rep

0.04
0.02
0.00
-0.02
-0.04

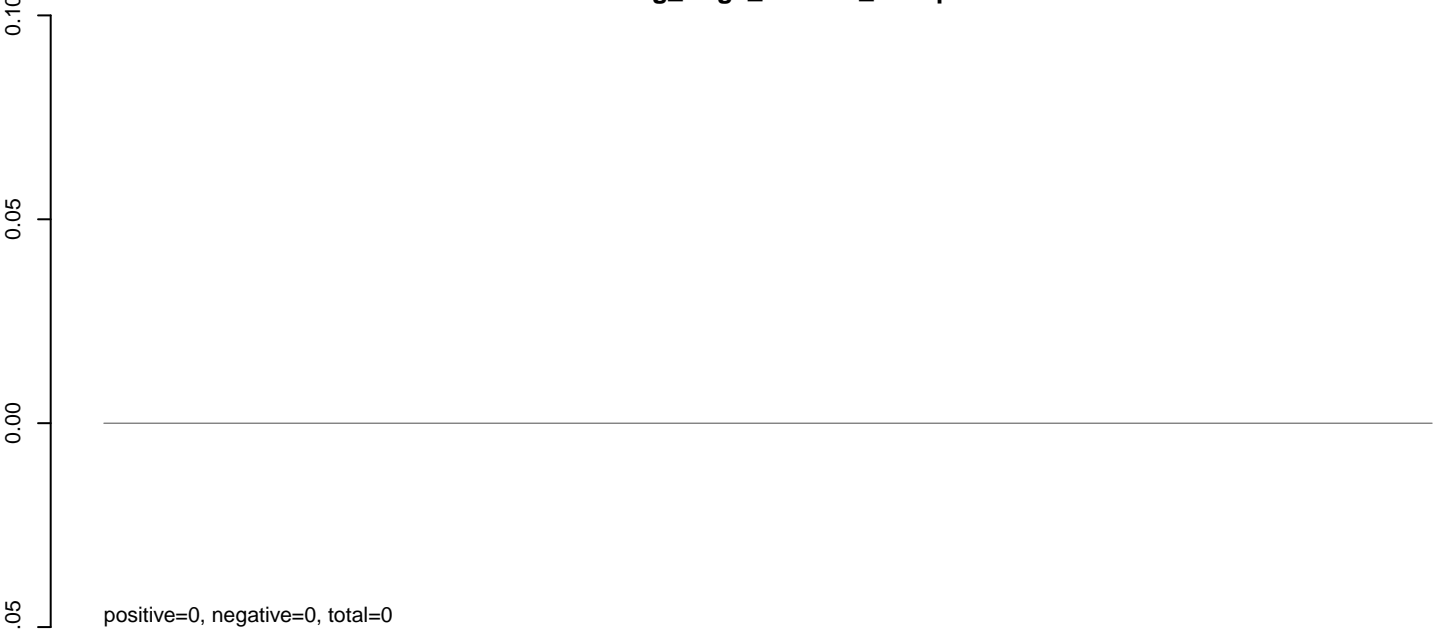
positive=0, negative=25, total=11624, virus@NC_023812.1-Madariaga_virus_strain_MADV_Cebus_apella_BRA_BEAN5122_1956:1-11624

0 2000 4000 6000 8000 10000 12000

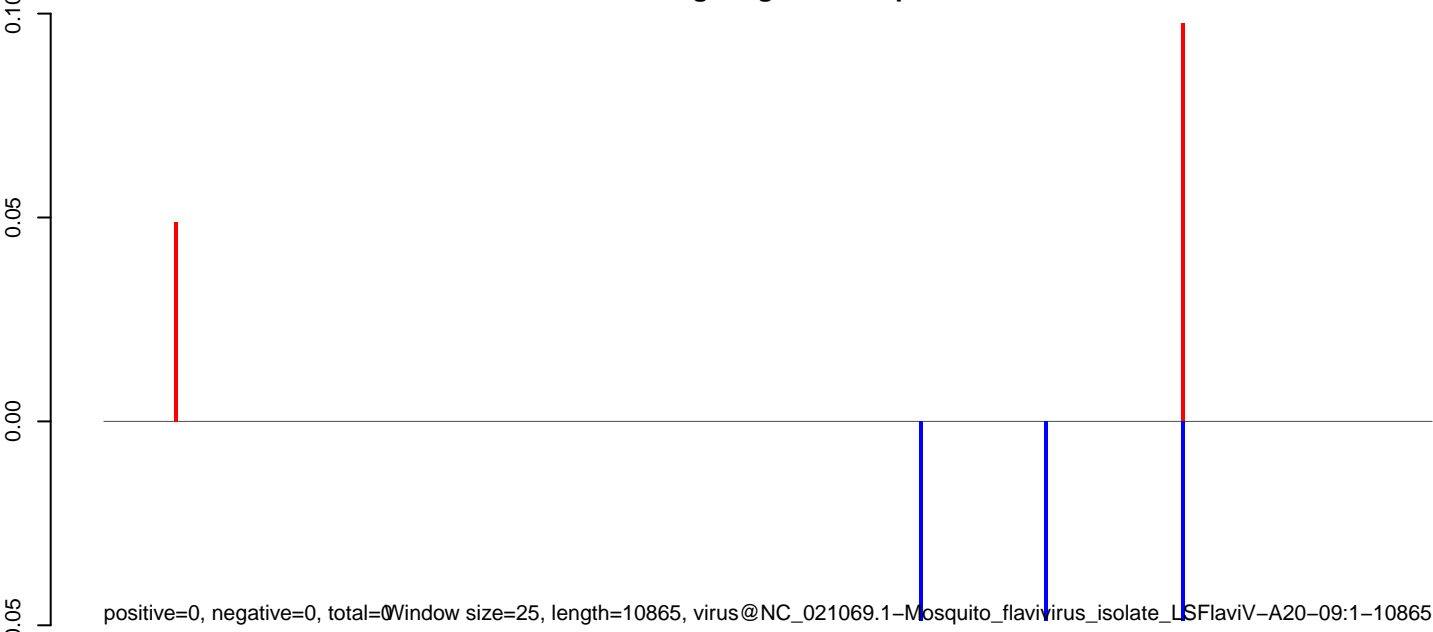
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

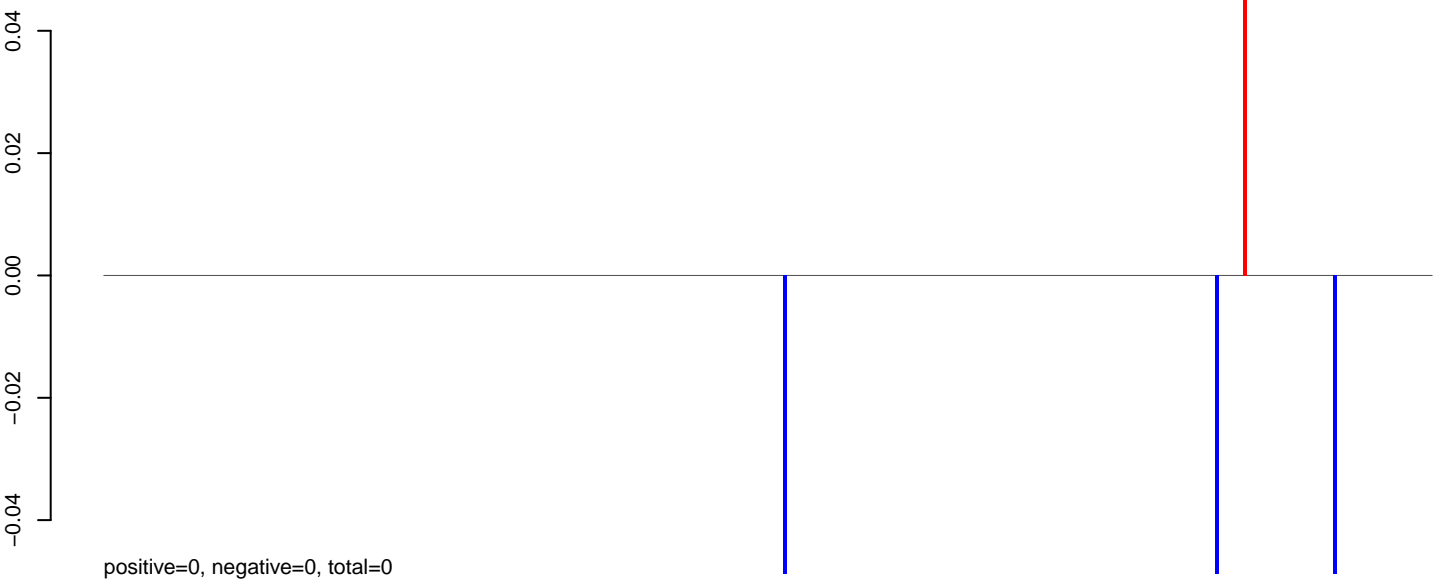


AeAeg_Aag2_cells.rep

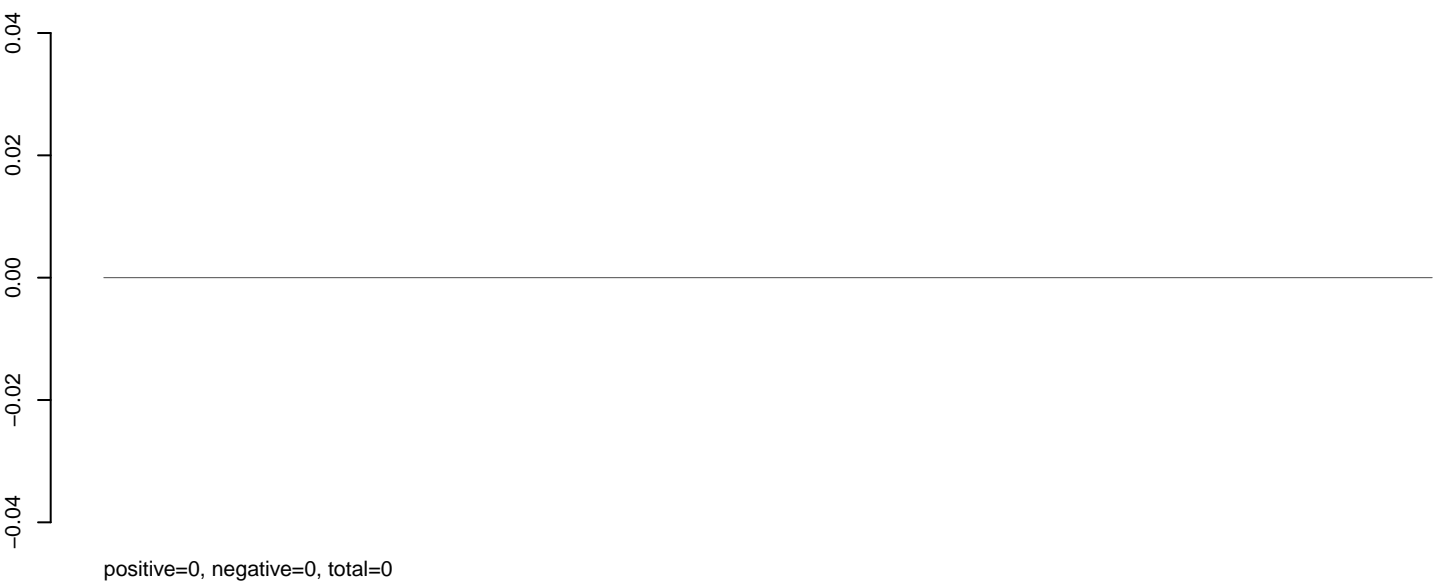


Window size=25, length=10865, virus@NC_021069.1-Mosquito_flavivirus_isolate_LSFlaviV-A20-09:1-10865

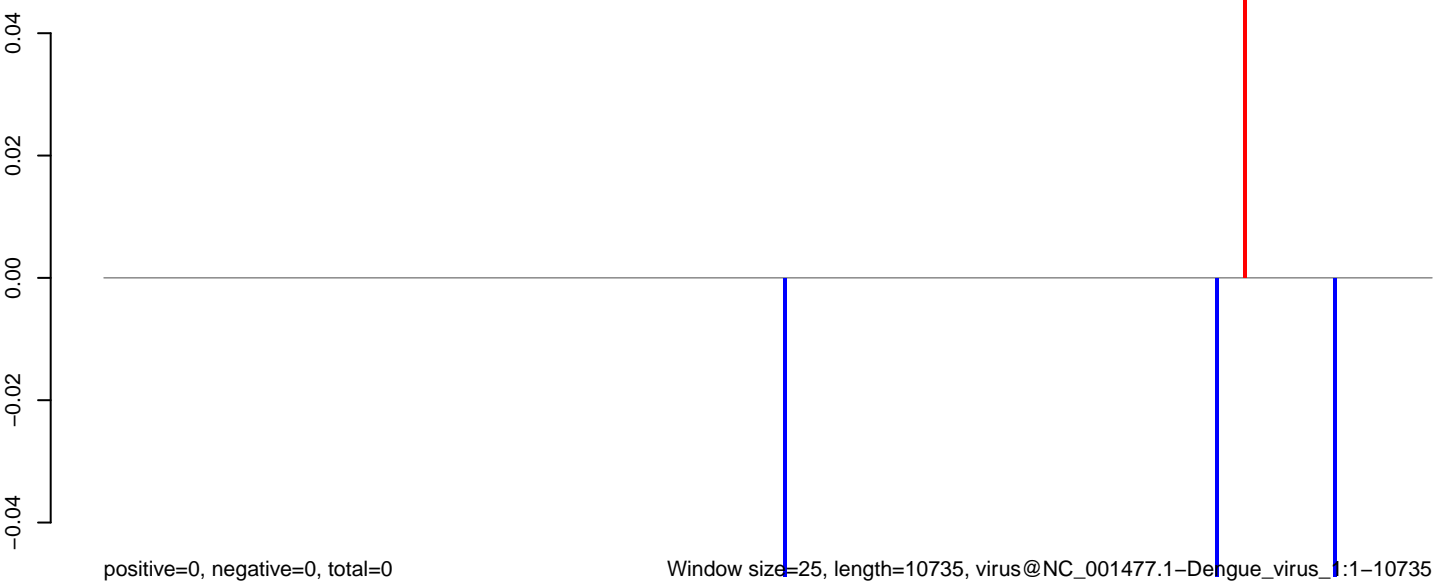
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

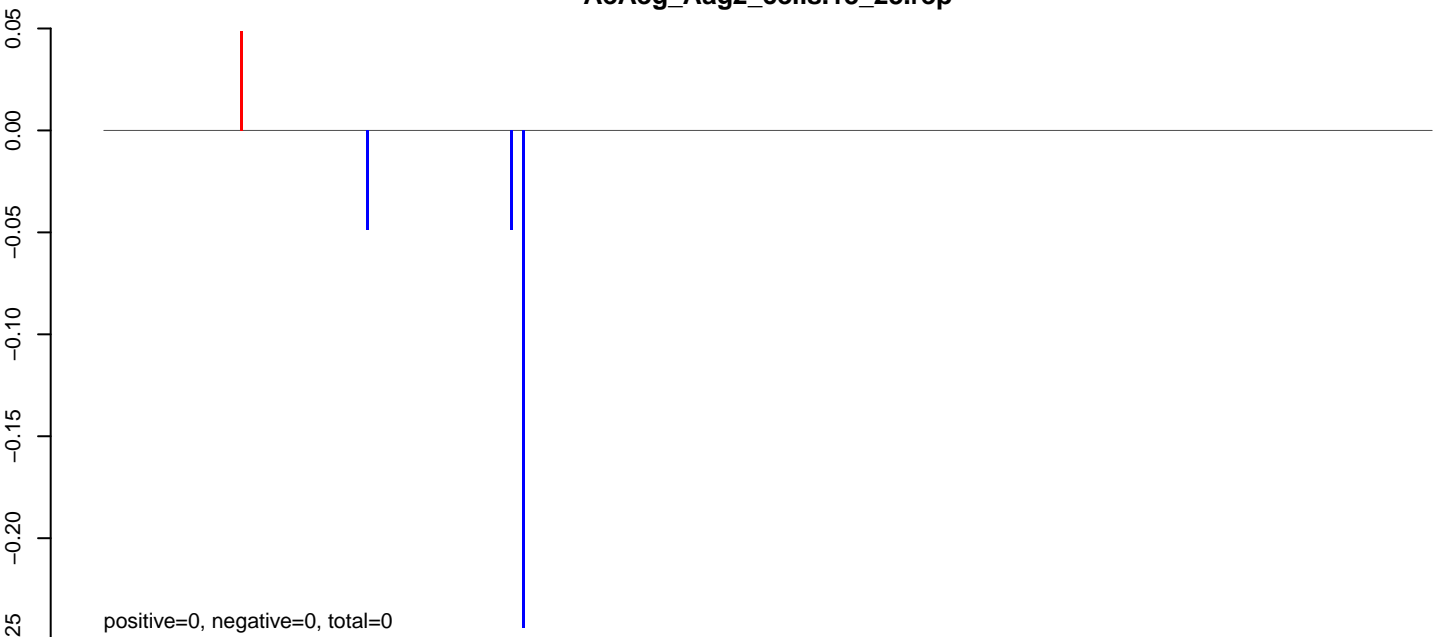


AeAeg_Aag2_cells.rep

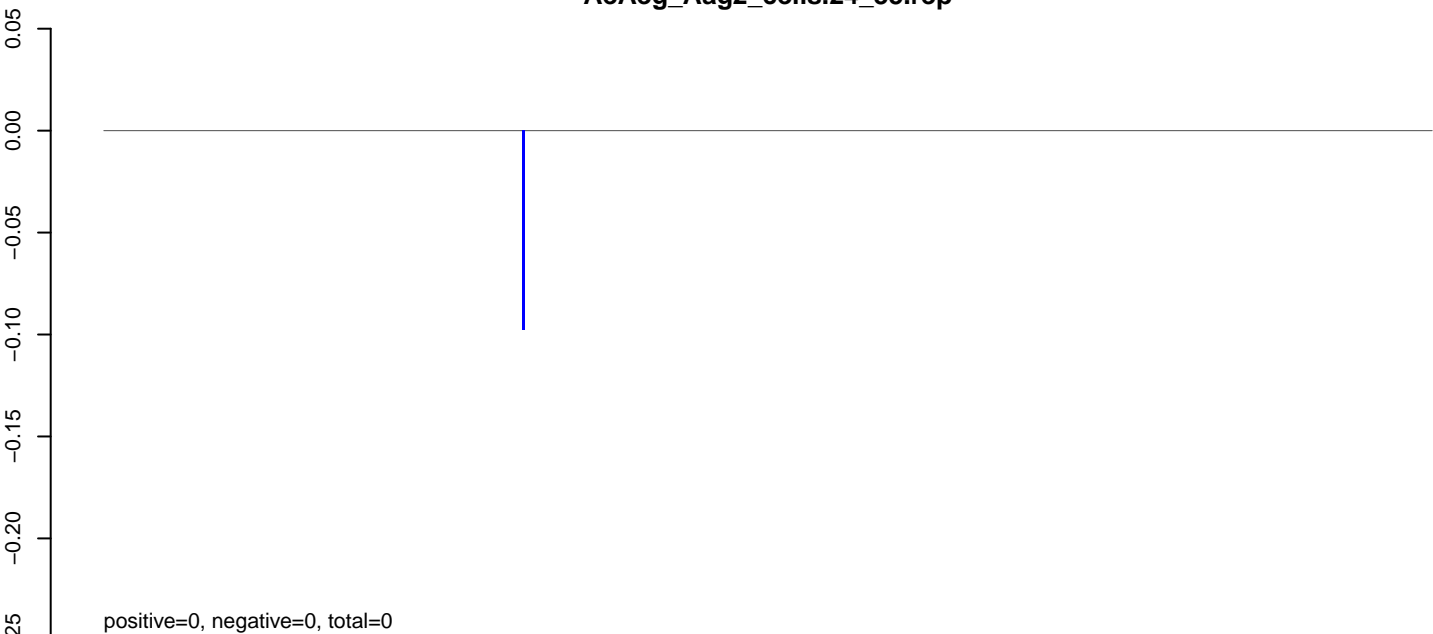


0 2000 4000 6000 8000 10000

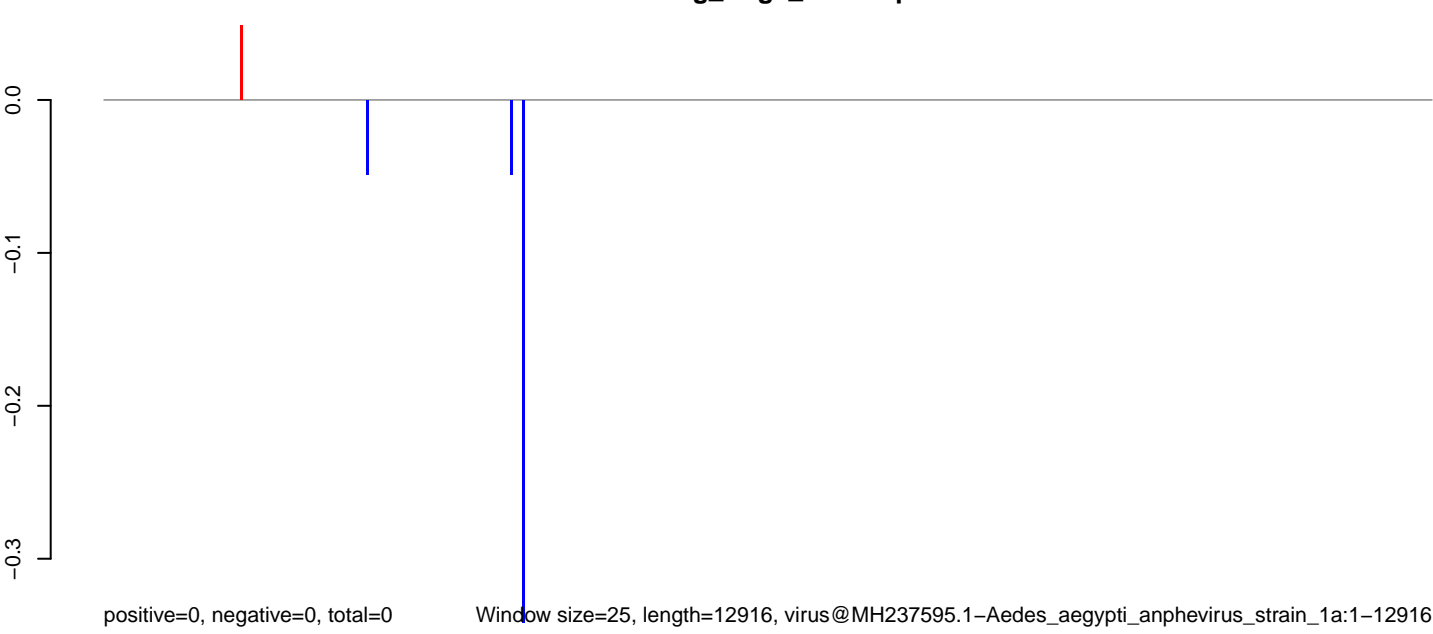
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

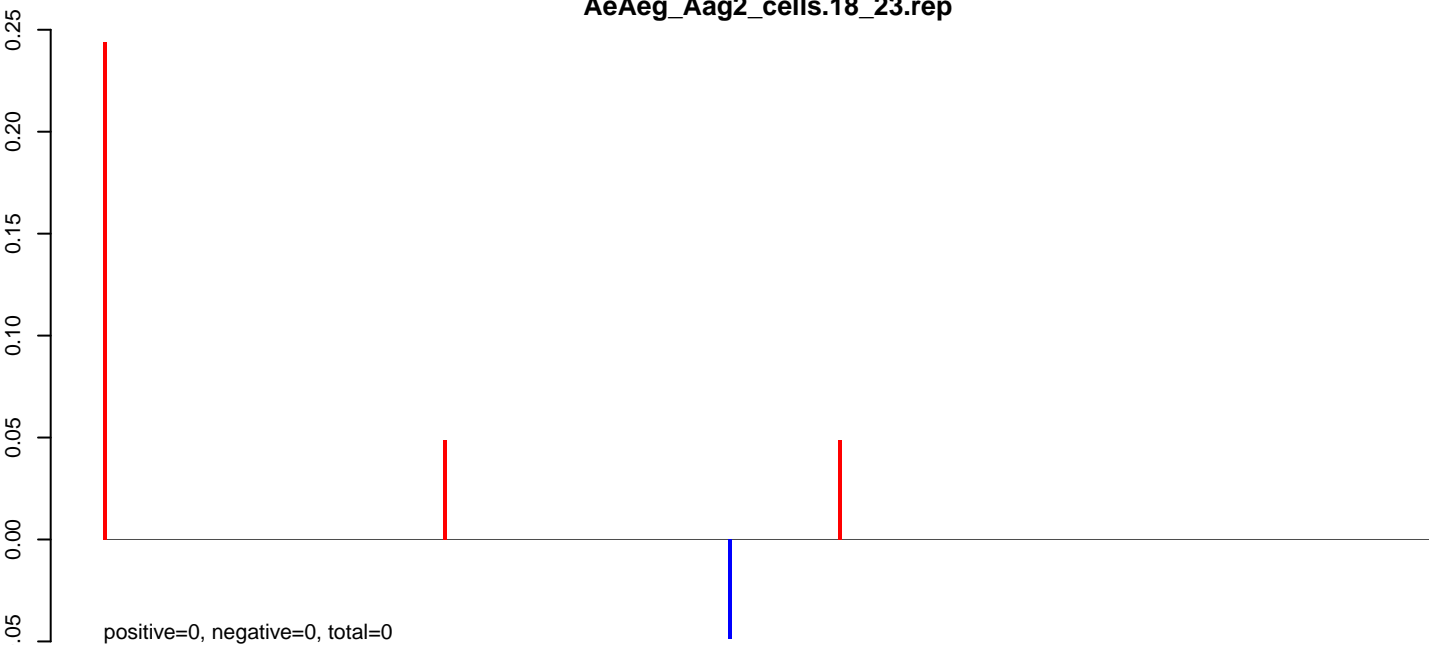


AeAeg_Aag2_cells.rep

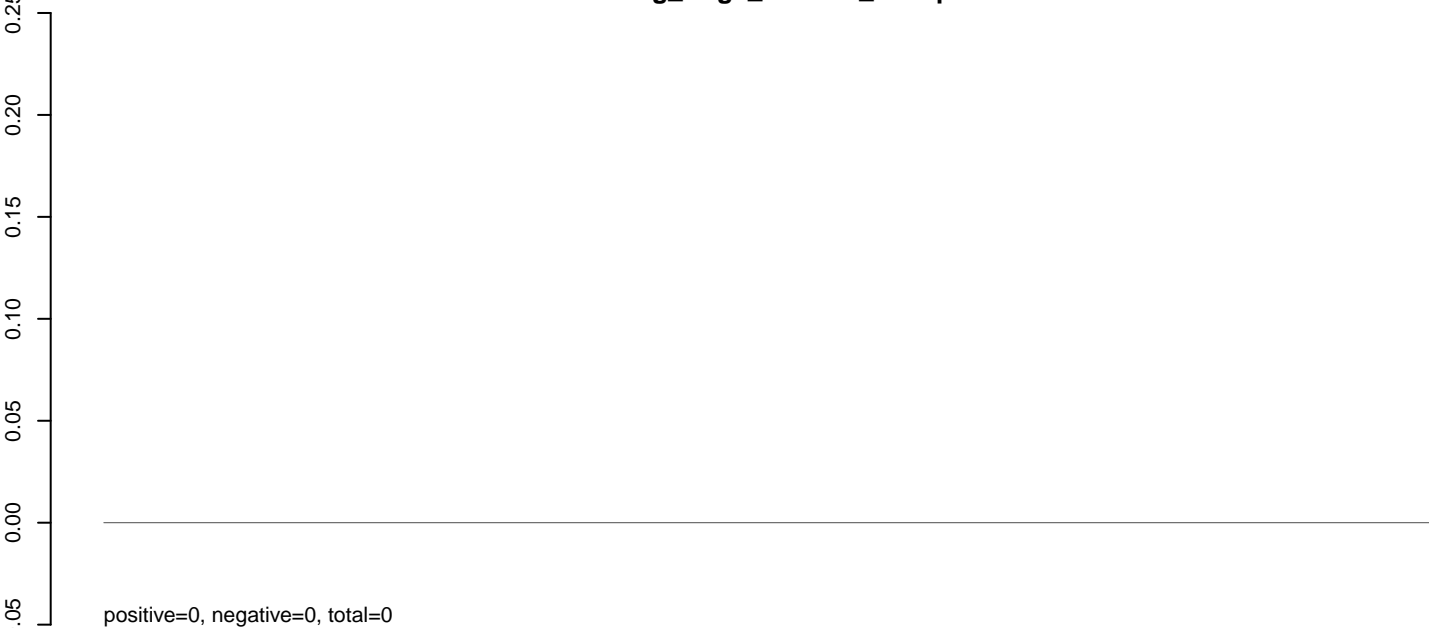


0 2000 4000 6000 8000 10000 12000

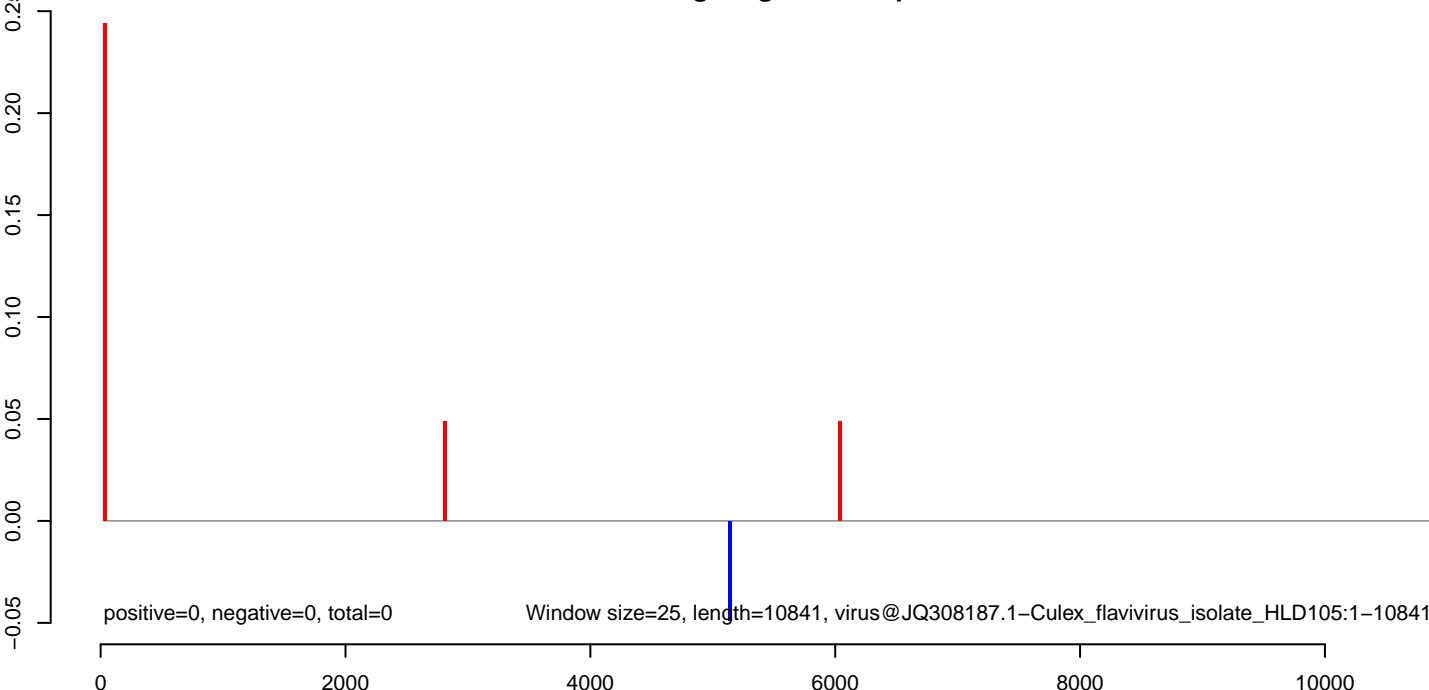
AeAeg_Aag2_cells.18_23.rep



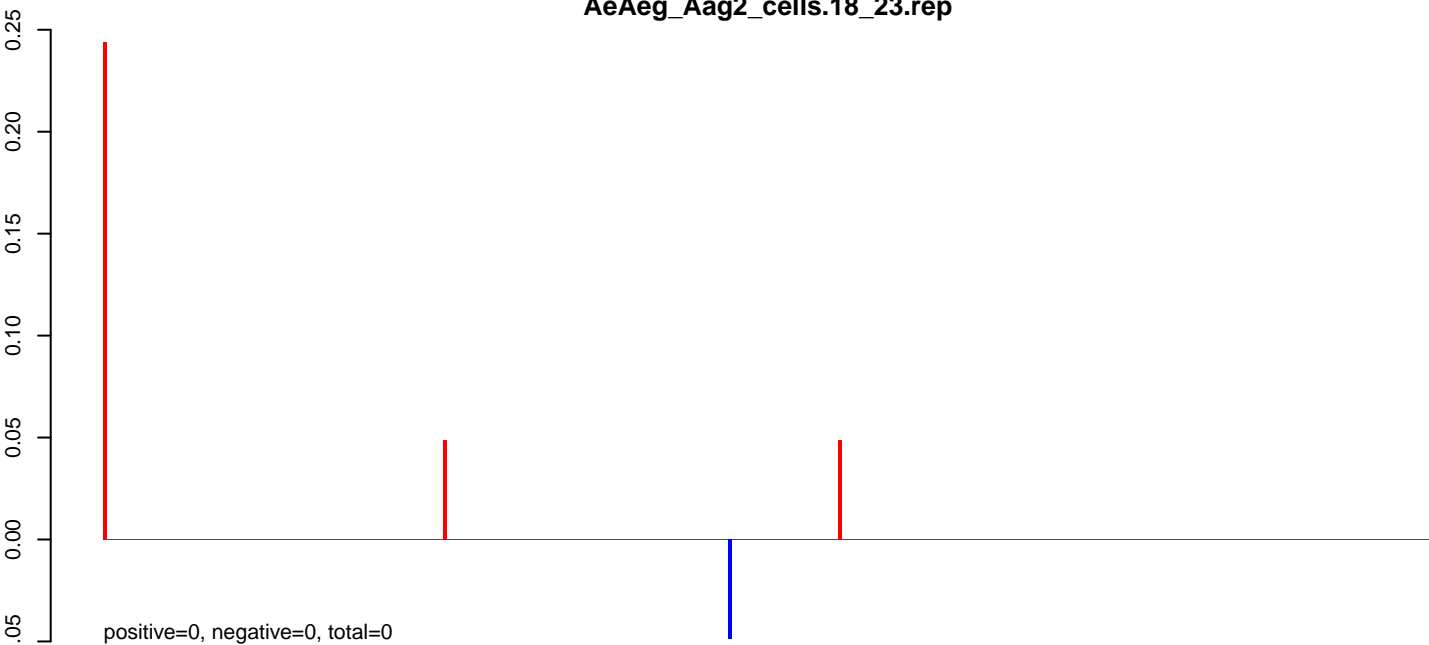
AeAeg_Aag2_cells.24_35.rep



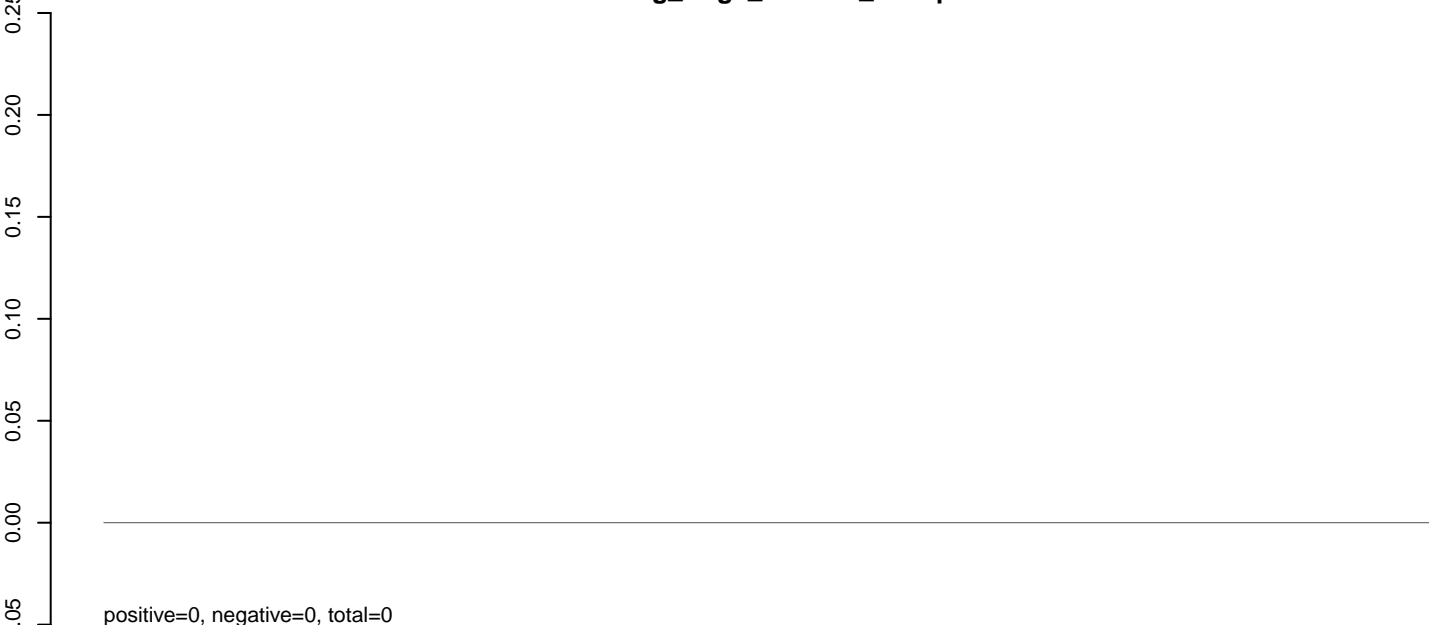
AeAeg_Aag2_cells.rep



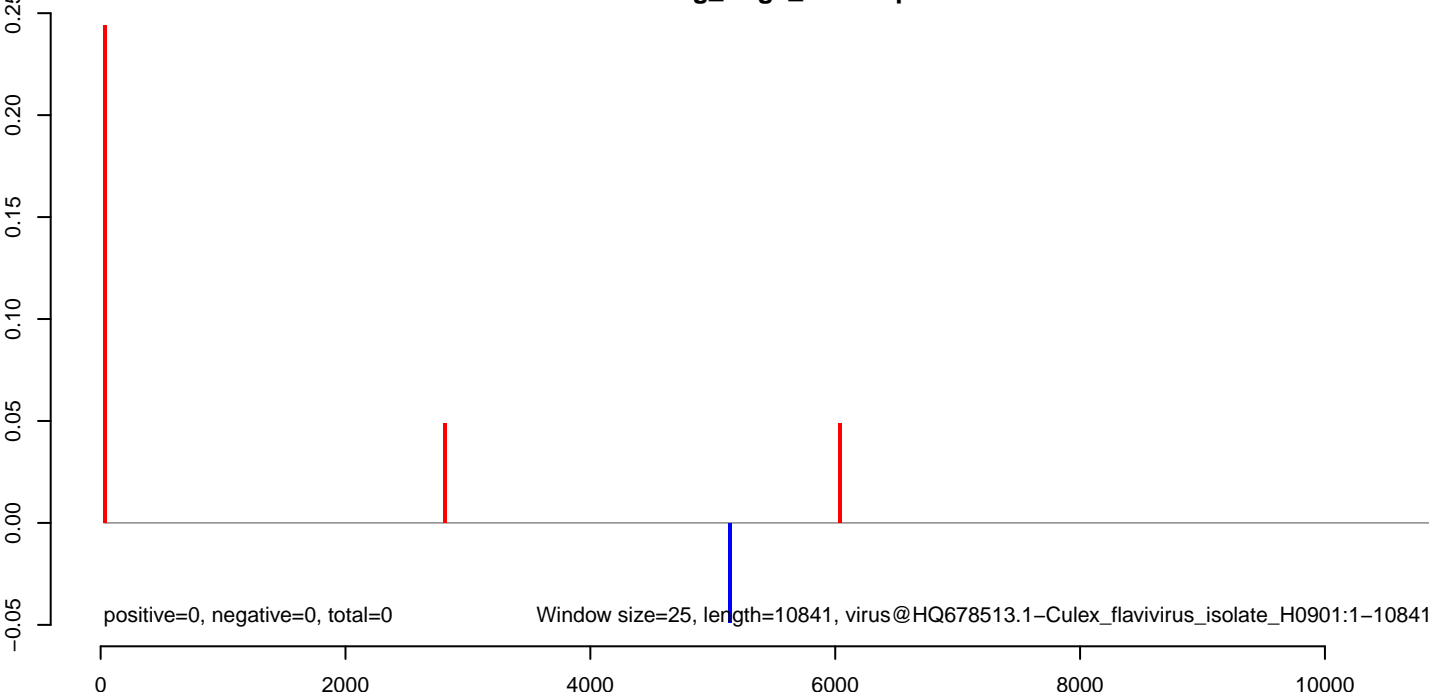
AeAeg_Aag2_cells.18_23.rep



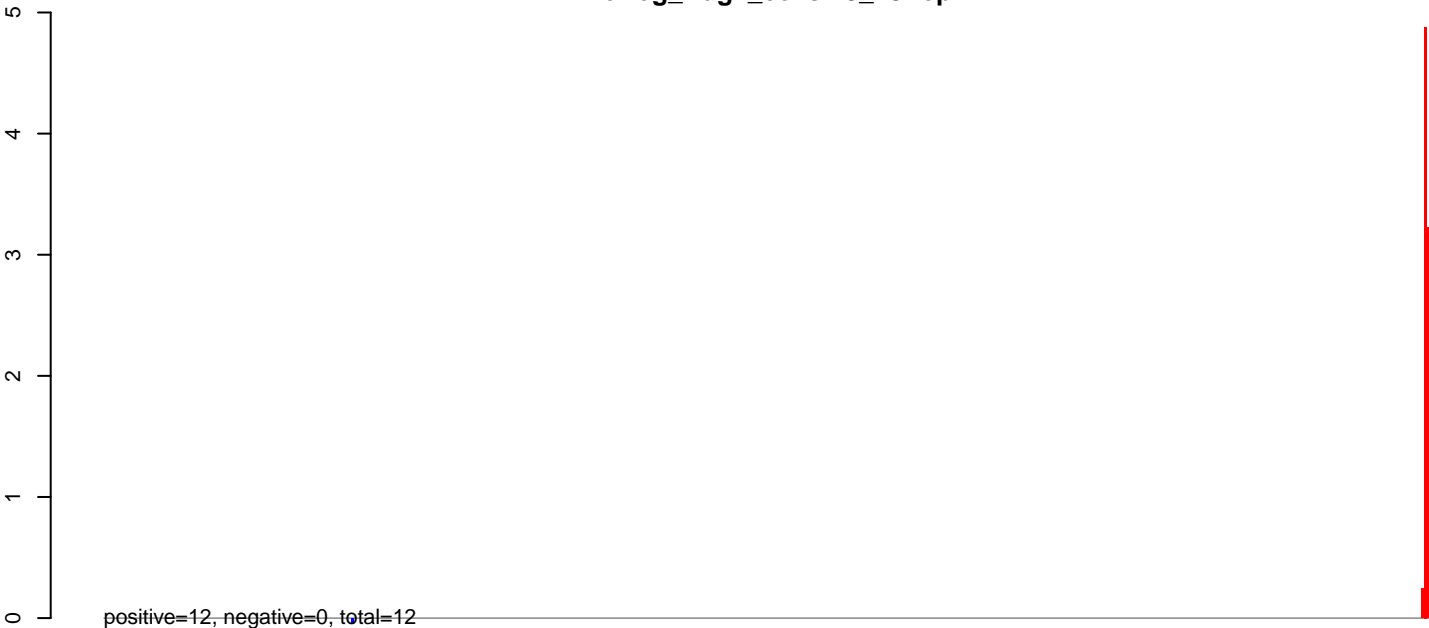
AeAeg_Aag2_cells.24_35.rep



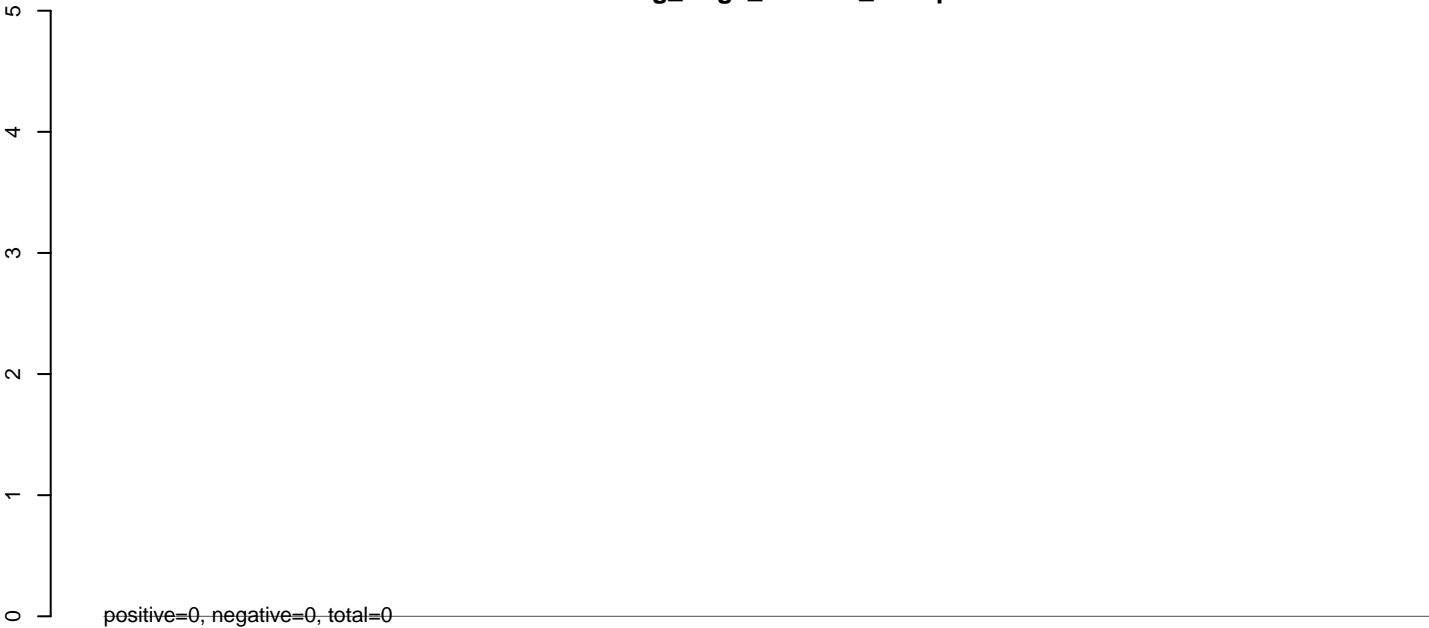
AeAeg_Aag2_cells.rep



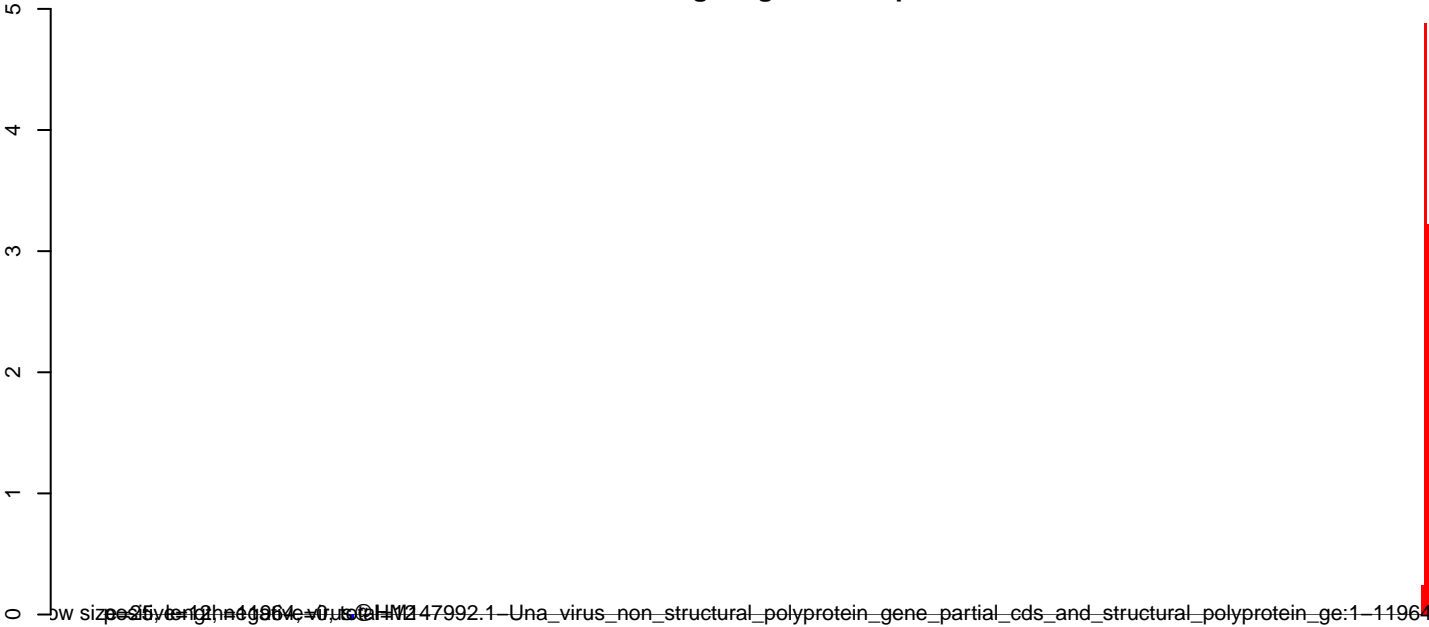
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



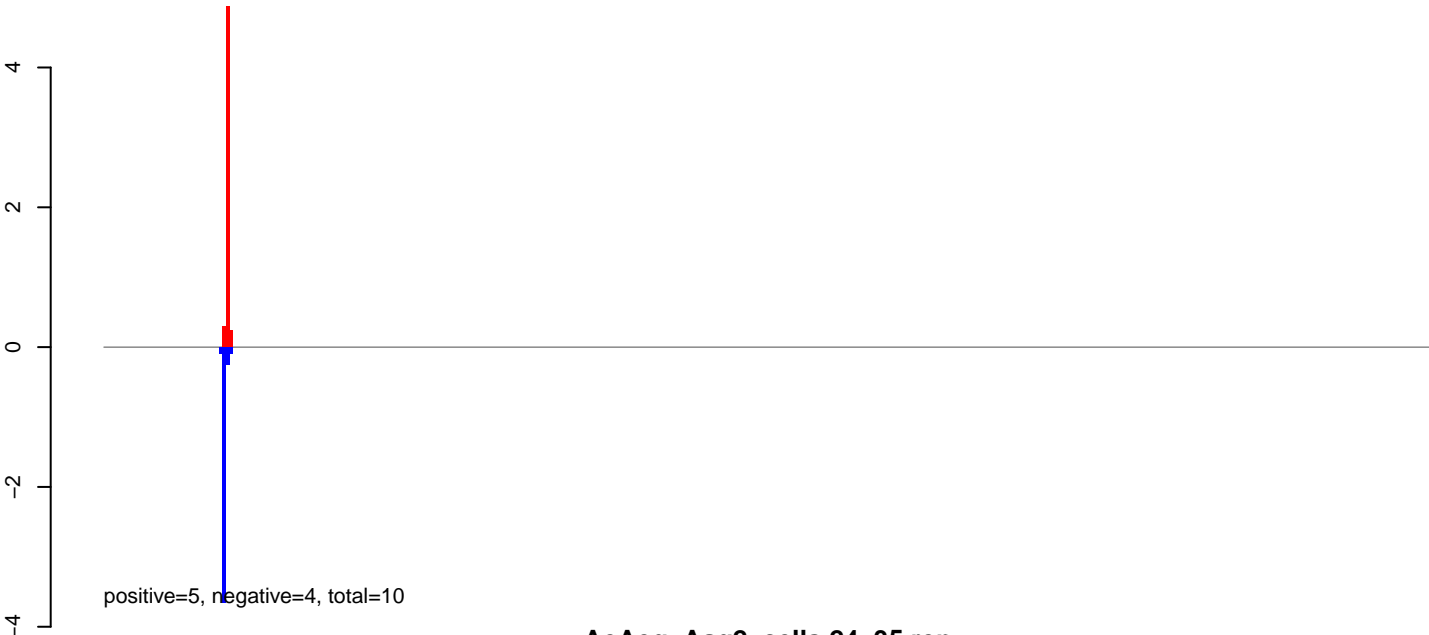
AeAeg_Aag2_cells.rep



bow size=12, length=12, negative=0, total=12
1-Una_virus_non_structural_polyprotein_gene_partial_cds_and_structural_polyprotein_ge:1-11964

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_cells.18_23.rep



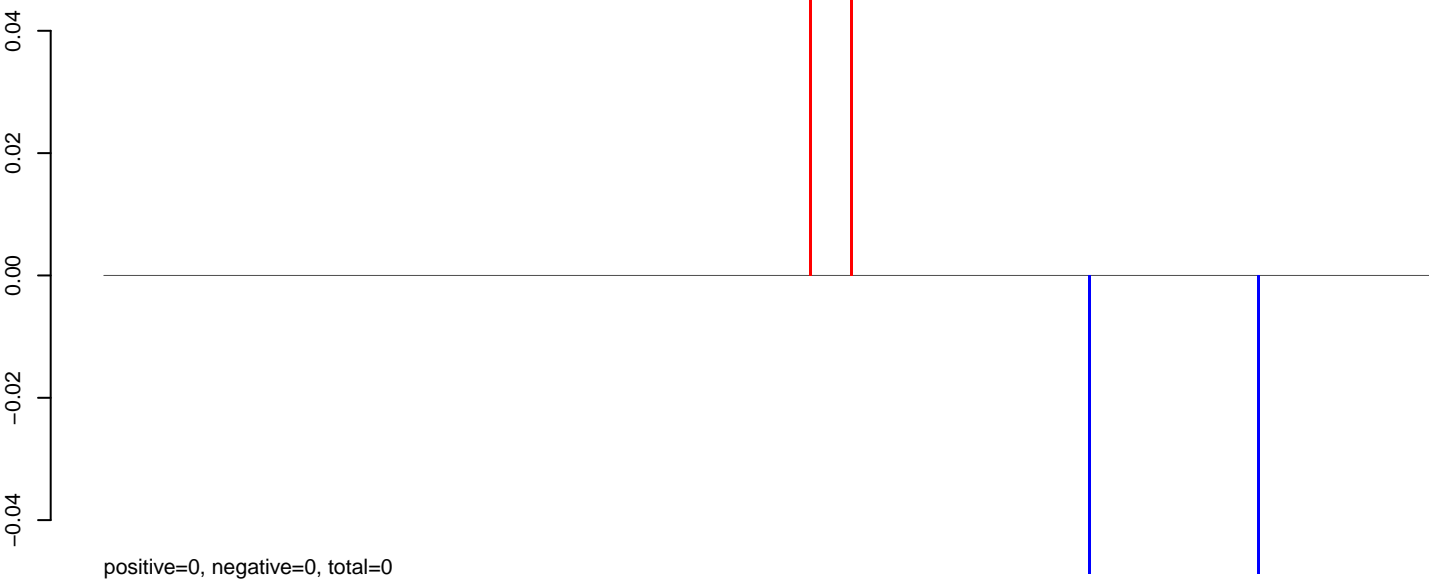
AeAeg_Aag2_cells.24_35.rep



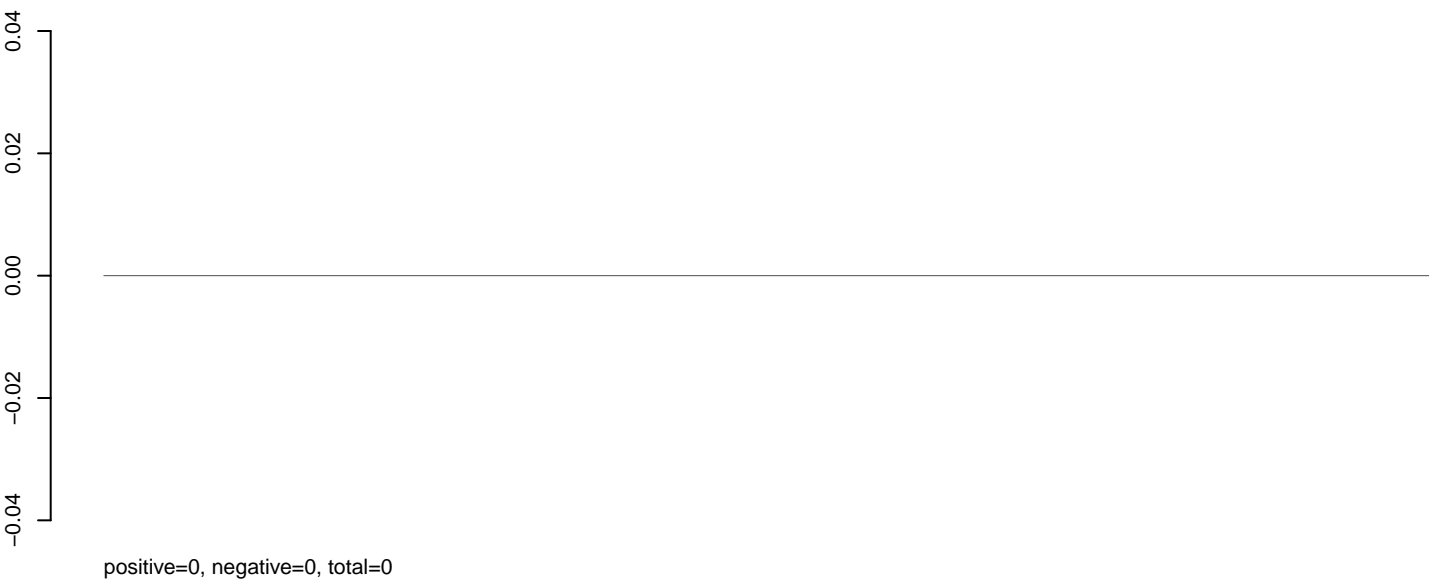
AeAeg_Aag2_cells.rep



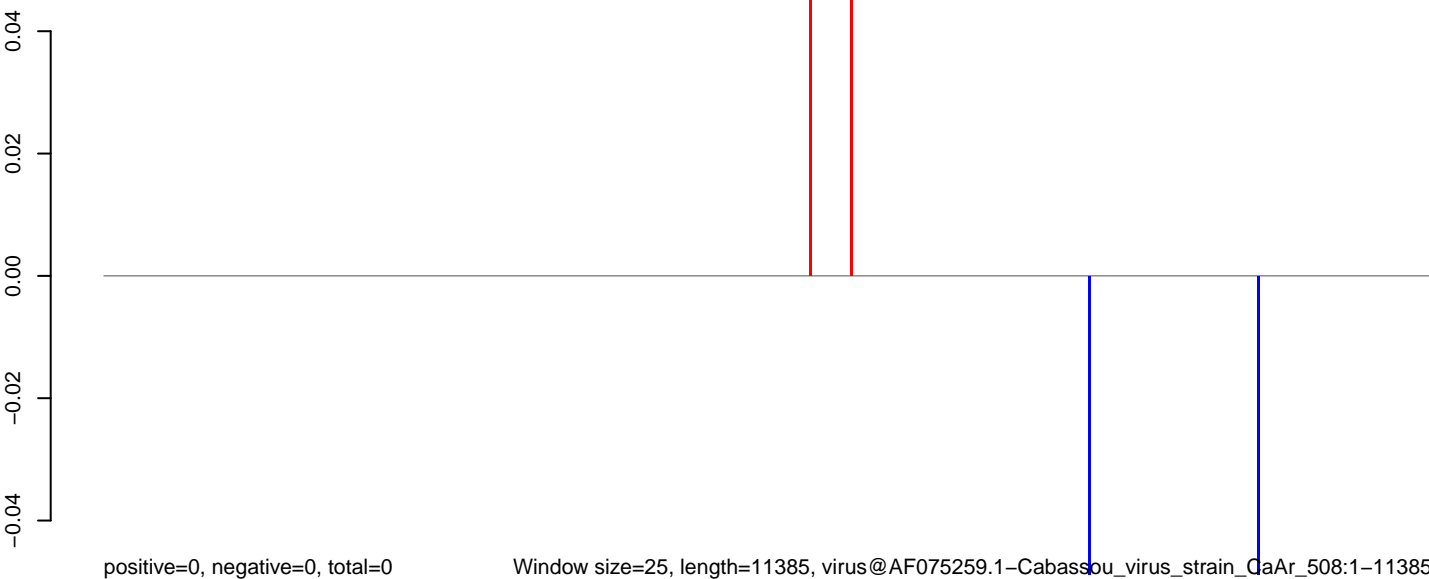
AeAeg_Aag2_cells.18_23.rep



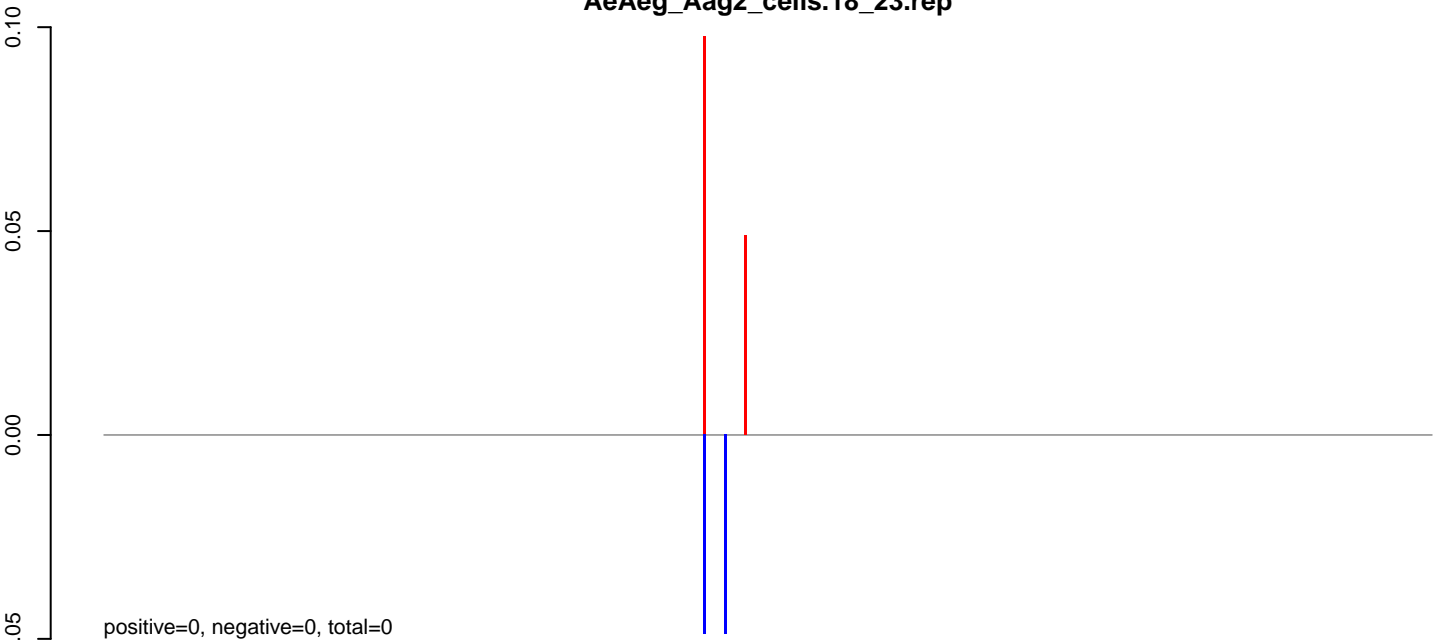
AeAeg_Aag2_cells.24_35.rep



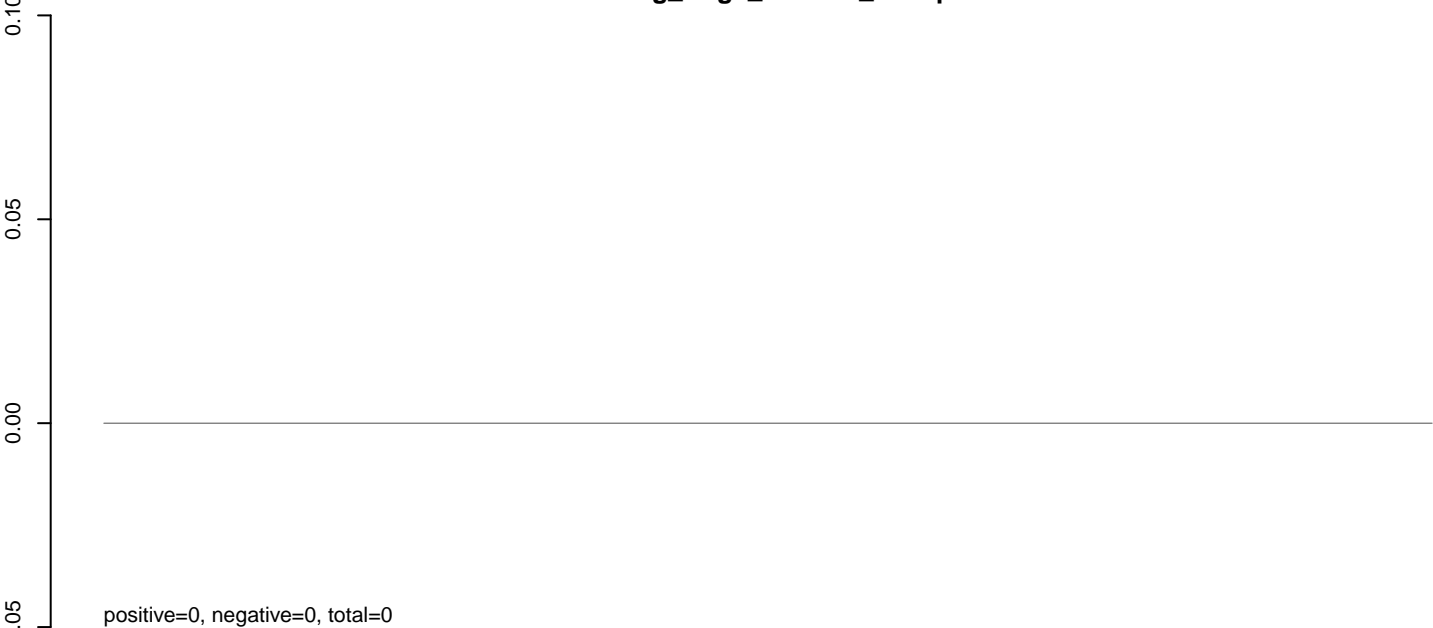
AeAeg_Aag2_cells.rep



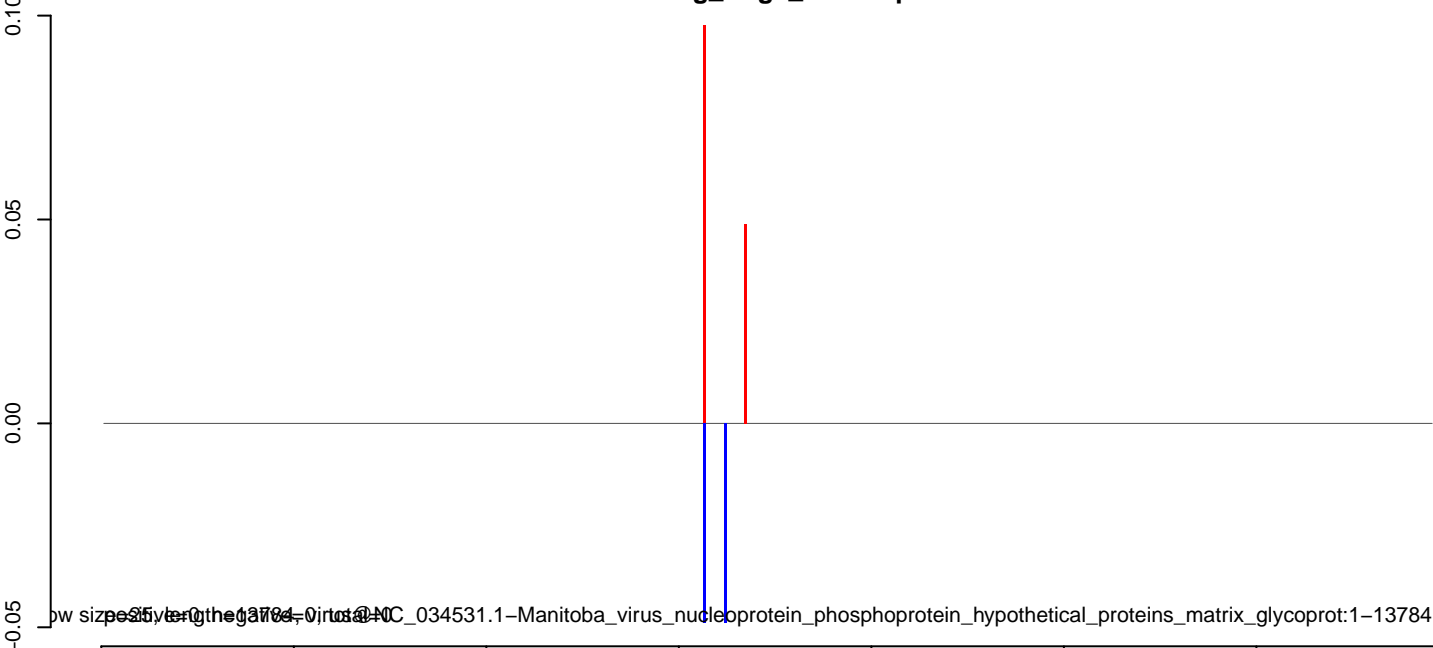
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

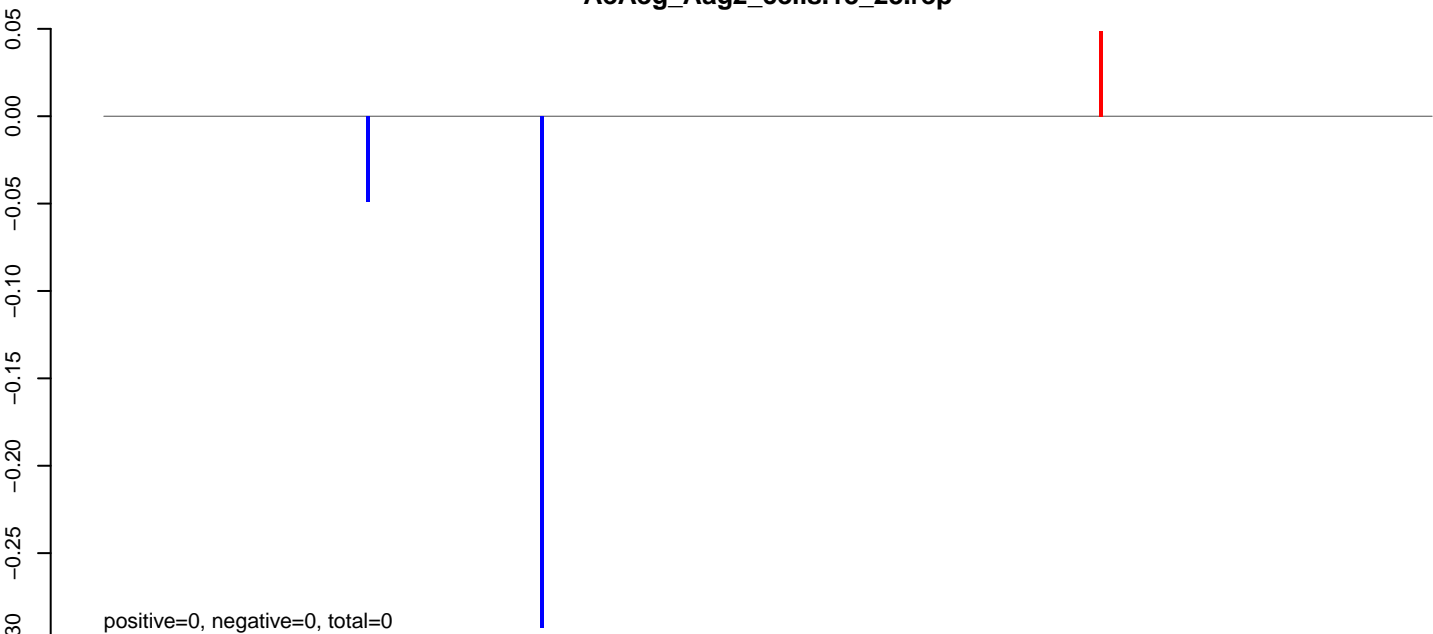


AeAeg_Aag2_cells.rep

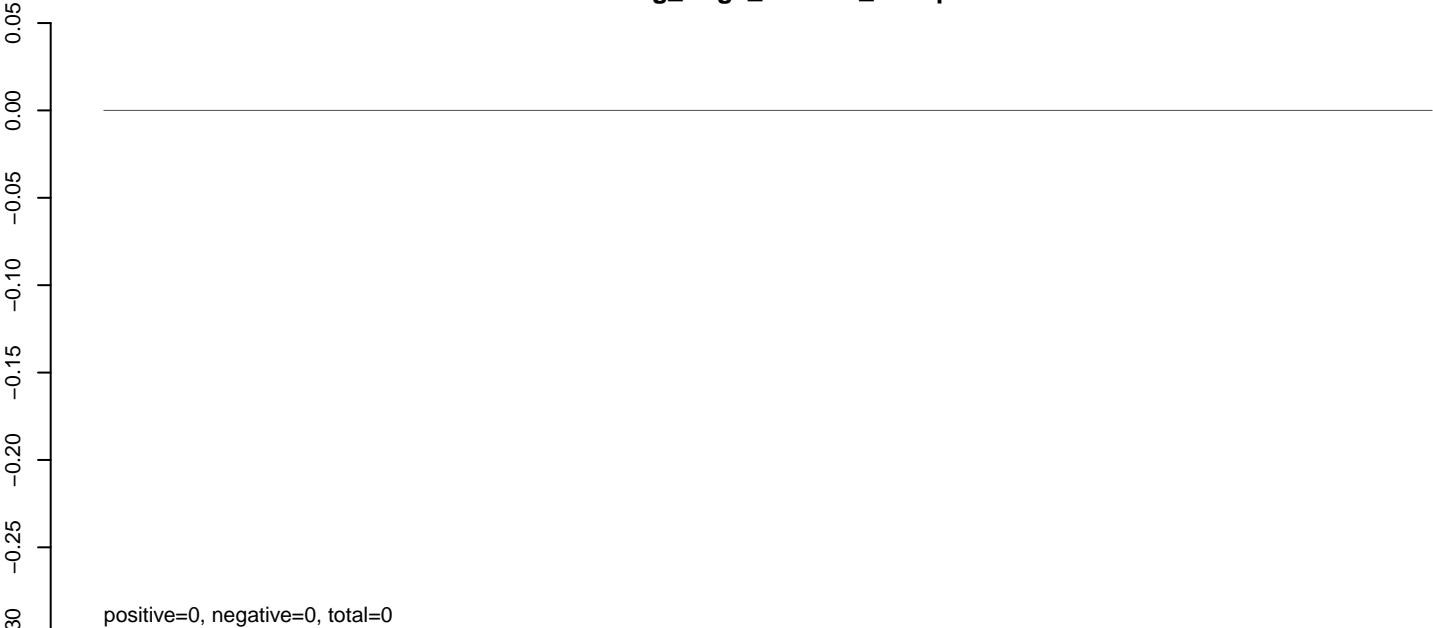


bow size=25, length=1784, int=40
NC_034531.1-Manitoba_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycoprot:1-13784

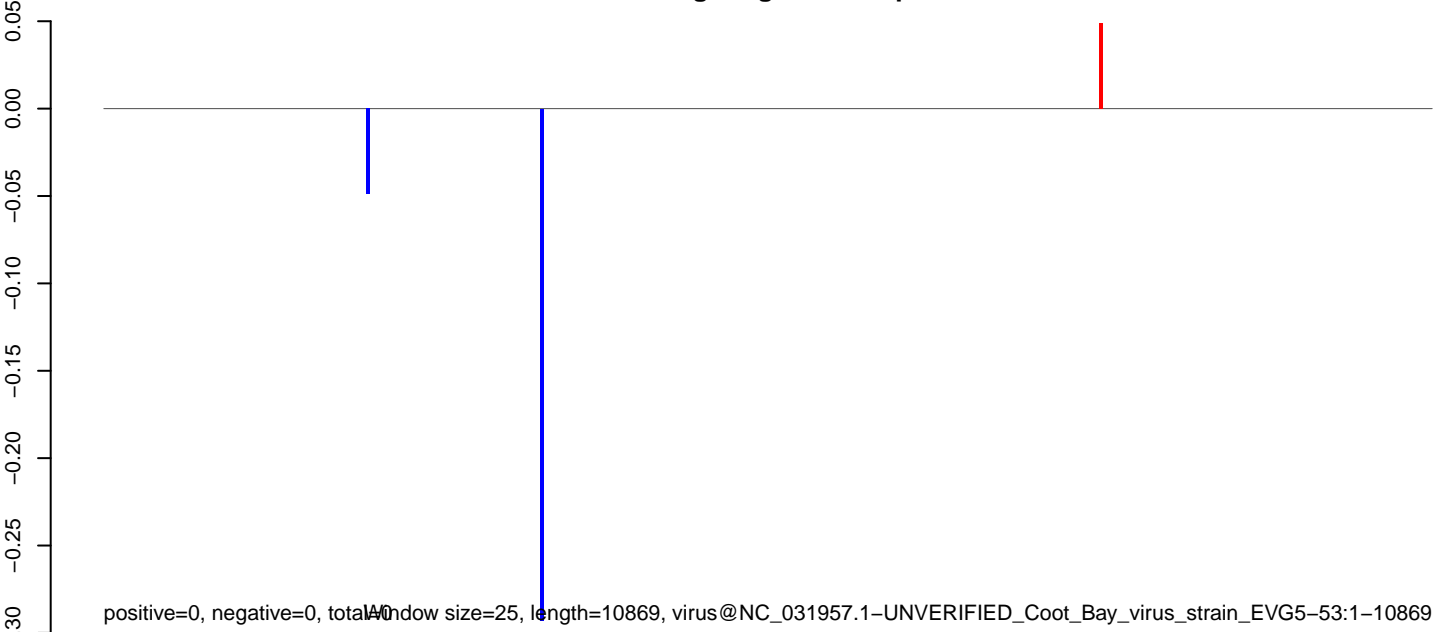
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



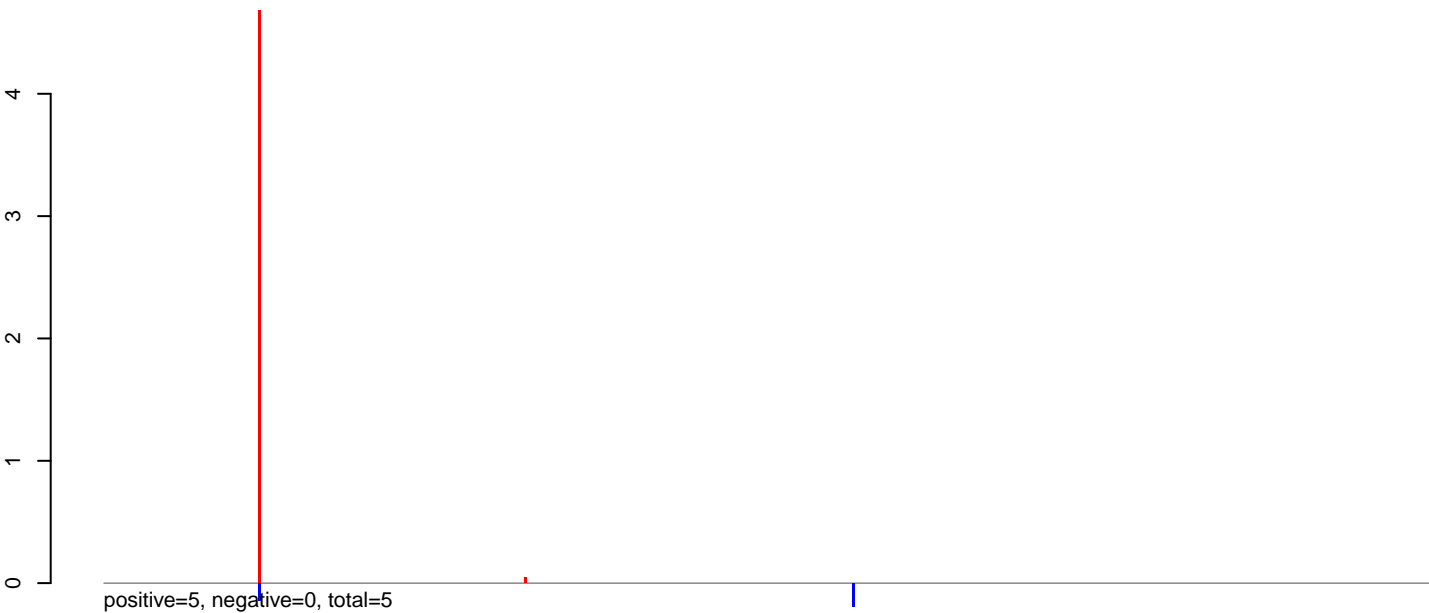
AeAeg_Aag2_cells.rep



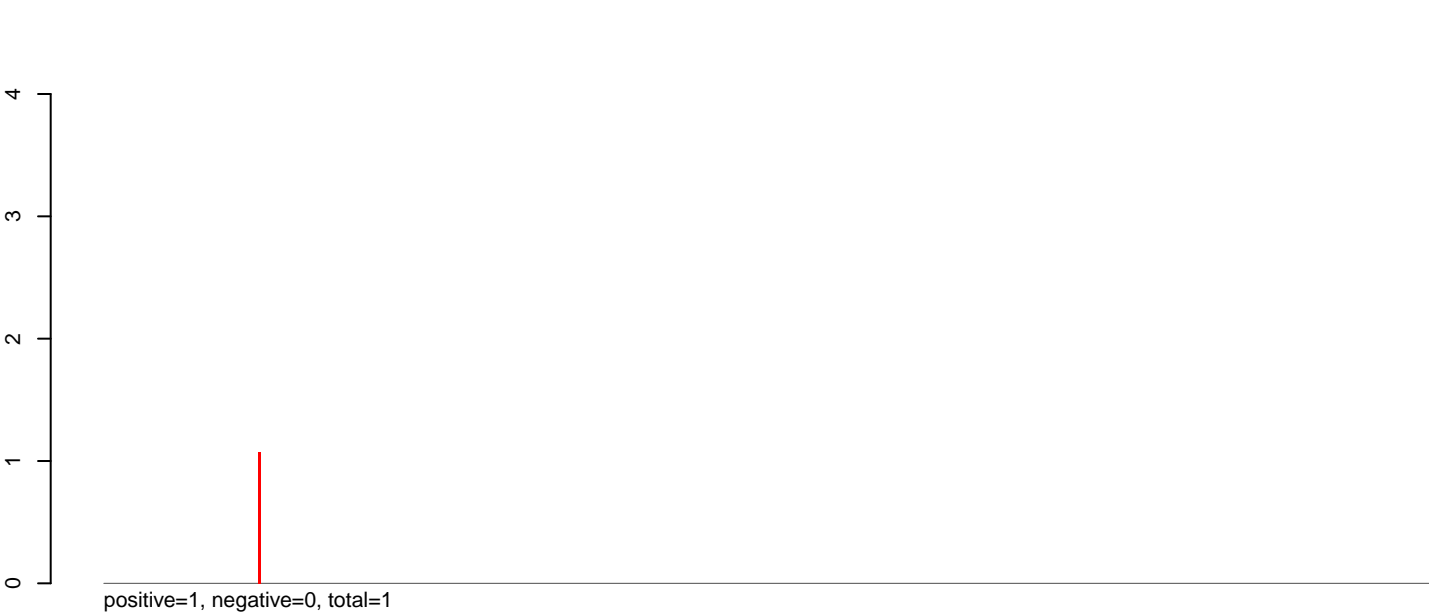
Window size=25, length=10869, virus@NC_031957.1-UNVERIFIED_Coot_Bay_virus_strain_EVG5-53:1-10869

0 2000 4000 6000 8000 10000

AeAeg_Aag2_cells.18_23.rep



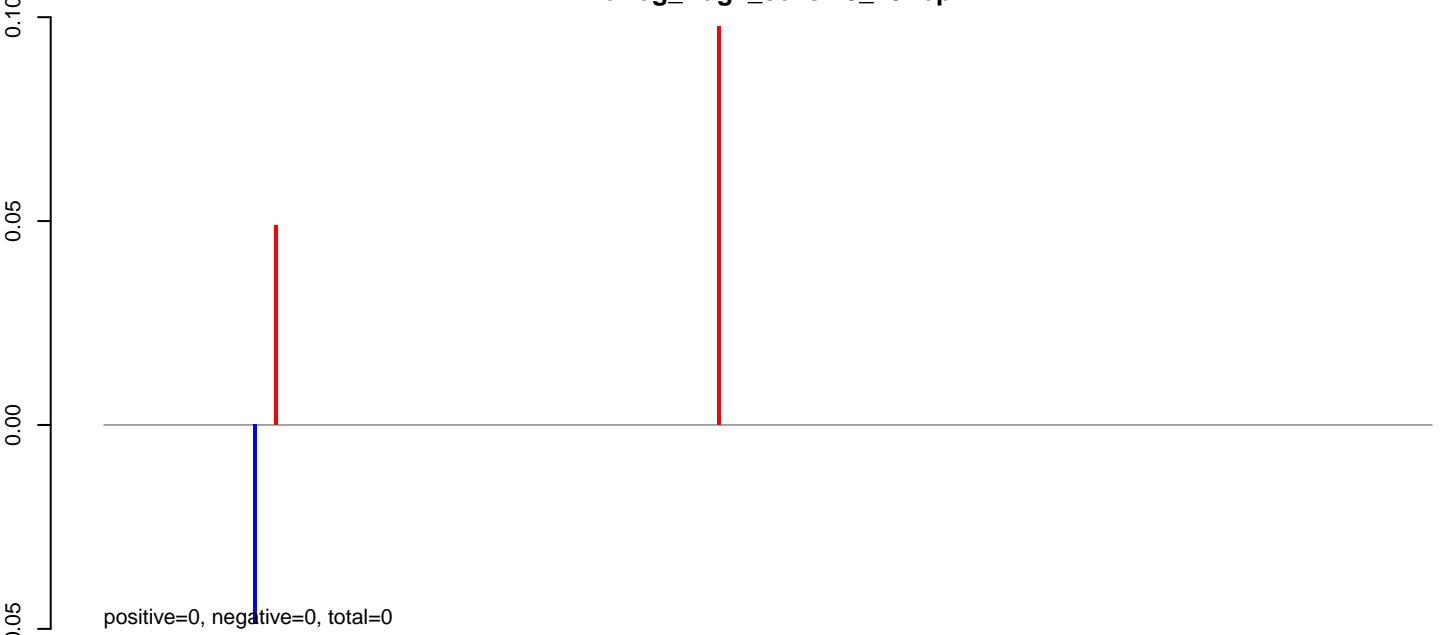
AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep



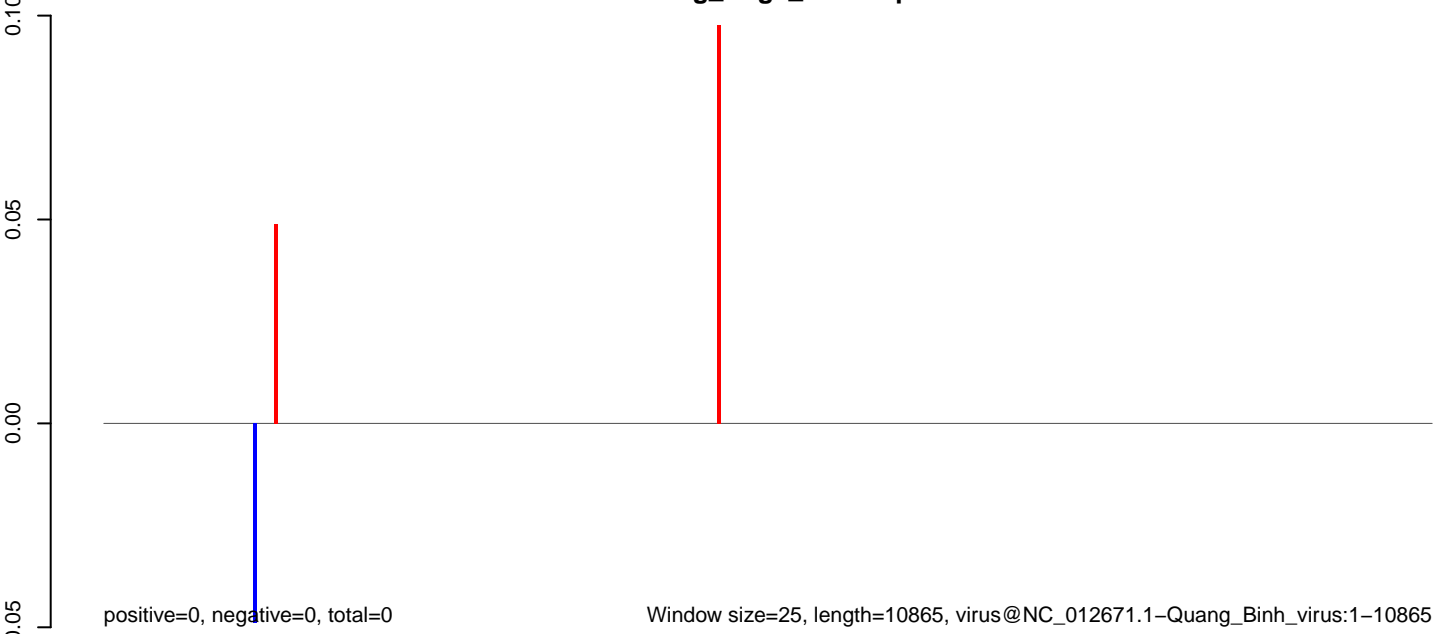
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

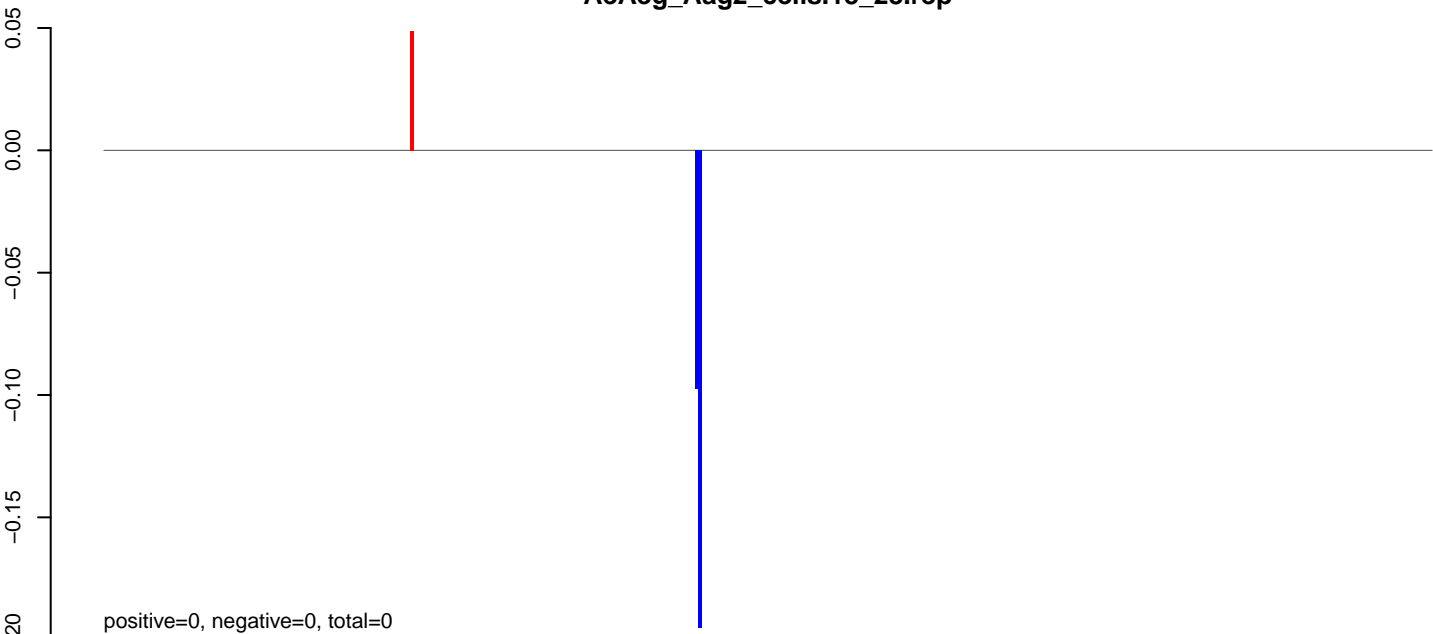


AeAeg_Aag2_cells.rep

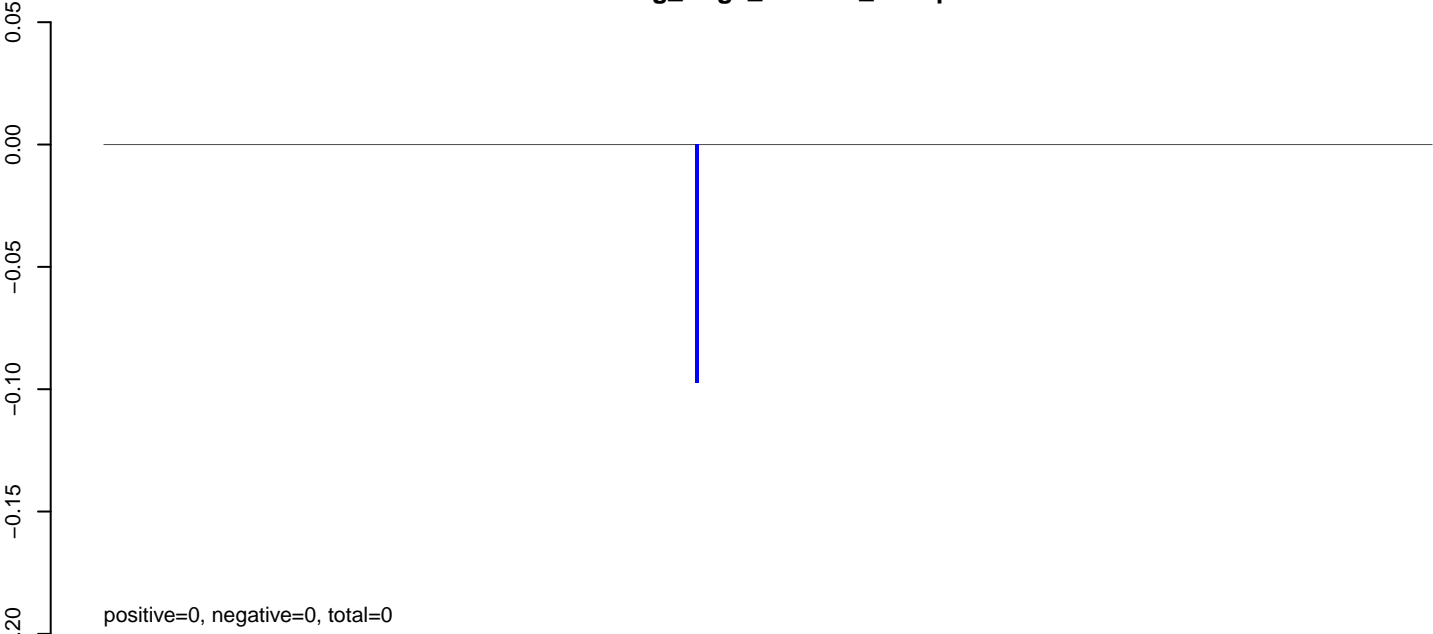


Window size=25, length=10865, virus@NC_012671.1-Quang_Binh_virus:1-10865

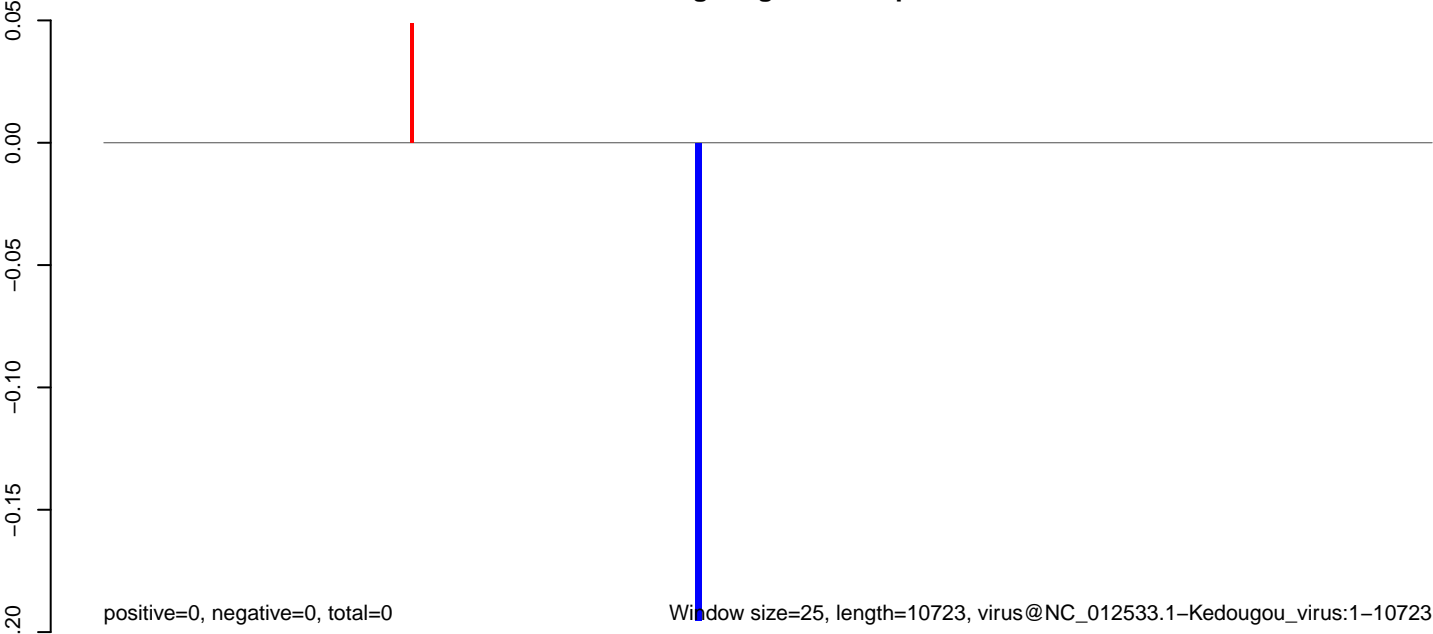
AeAeg_Aag2_cells.18_23.rep



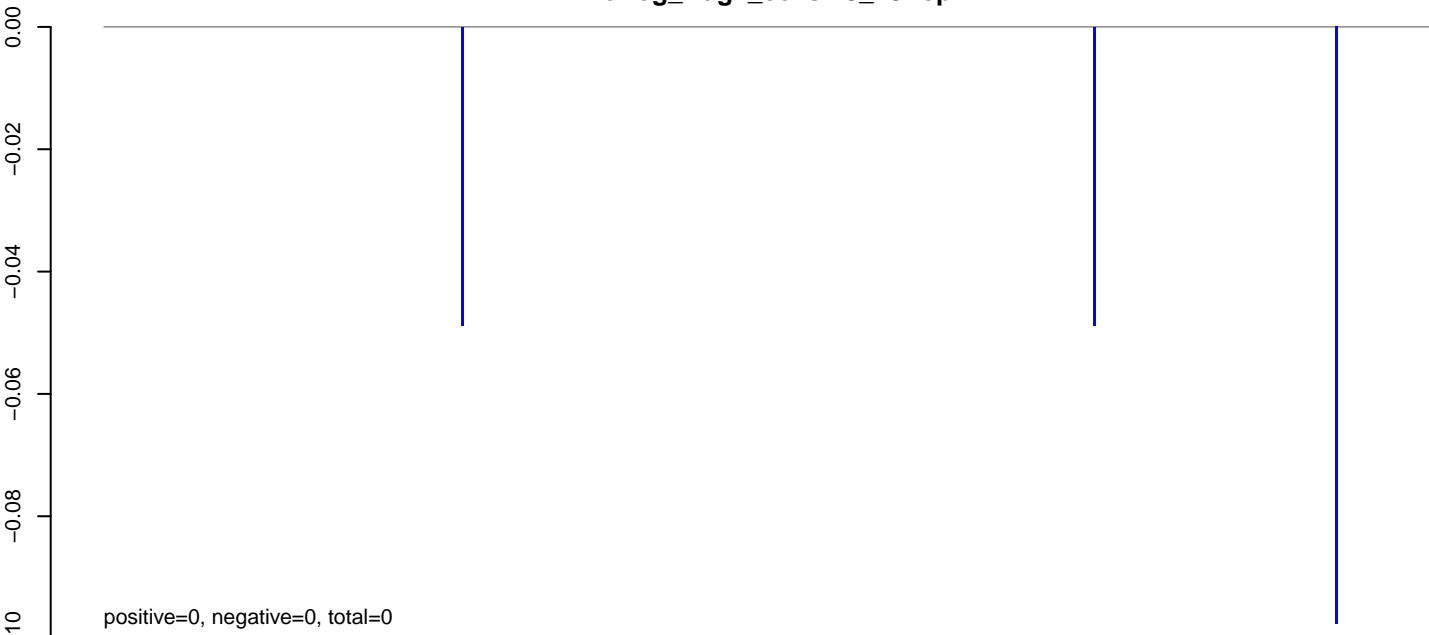
AeAeg_Aag2_cells.24_35.rep



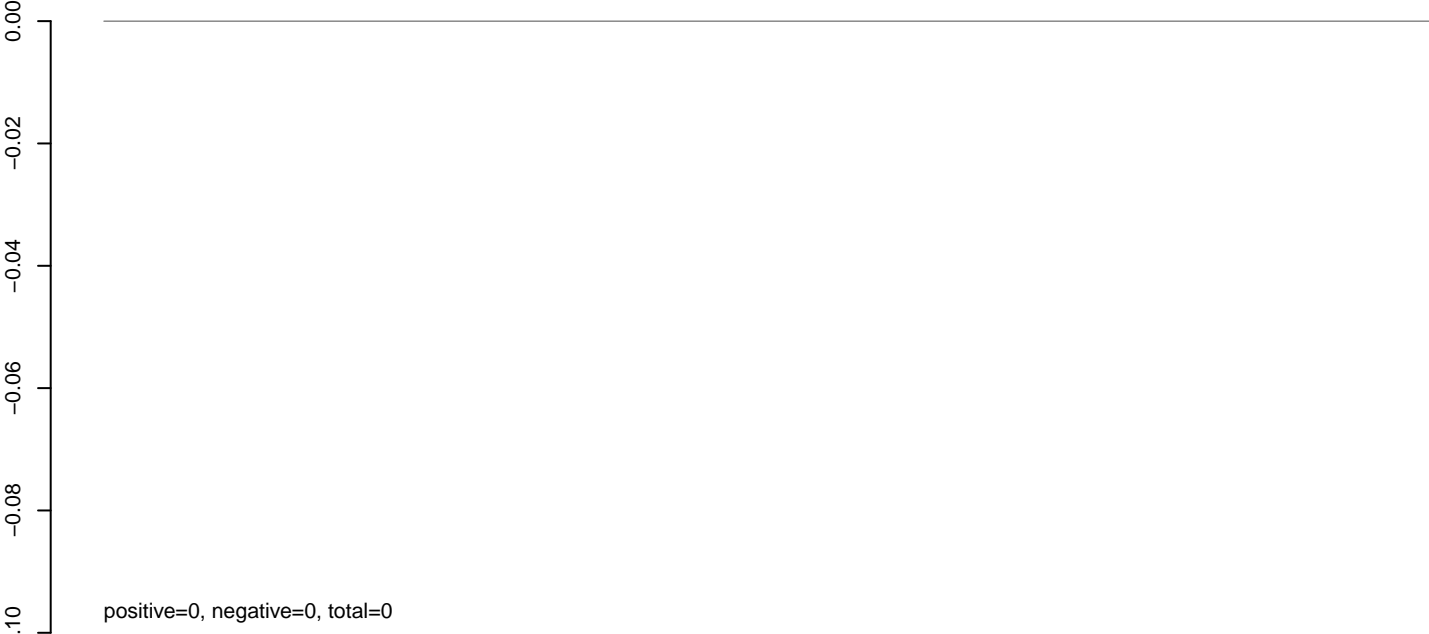
AeAeg_Aag2_cells.rep



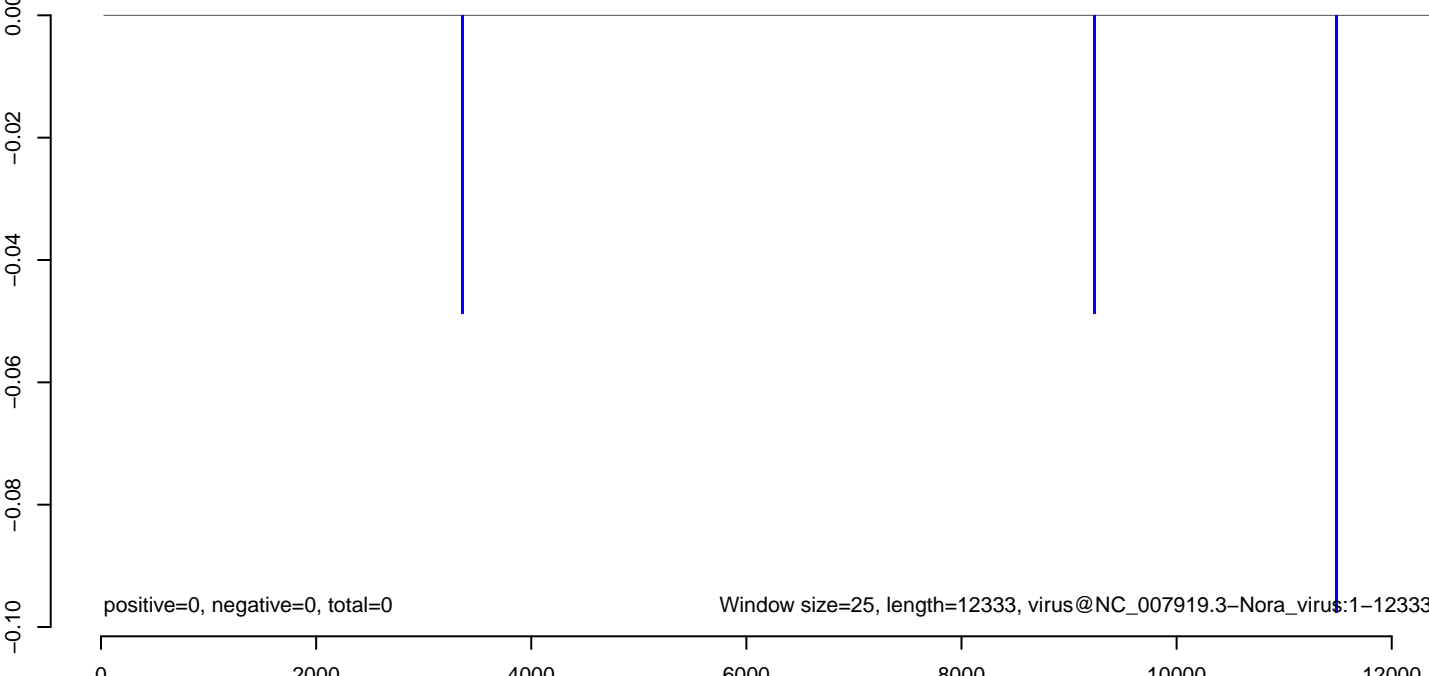
AeAeg_Aag2_cells.18_23.rep



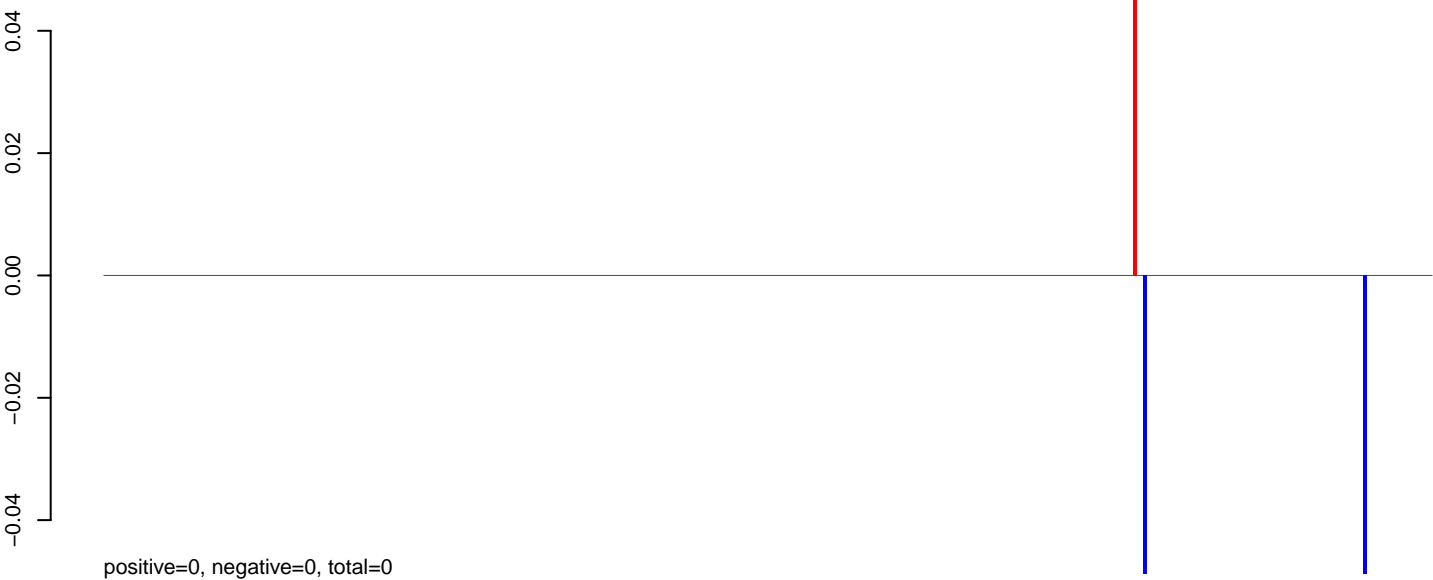
AeAeg_Aag2_cells.24_35.rep



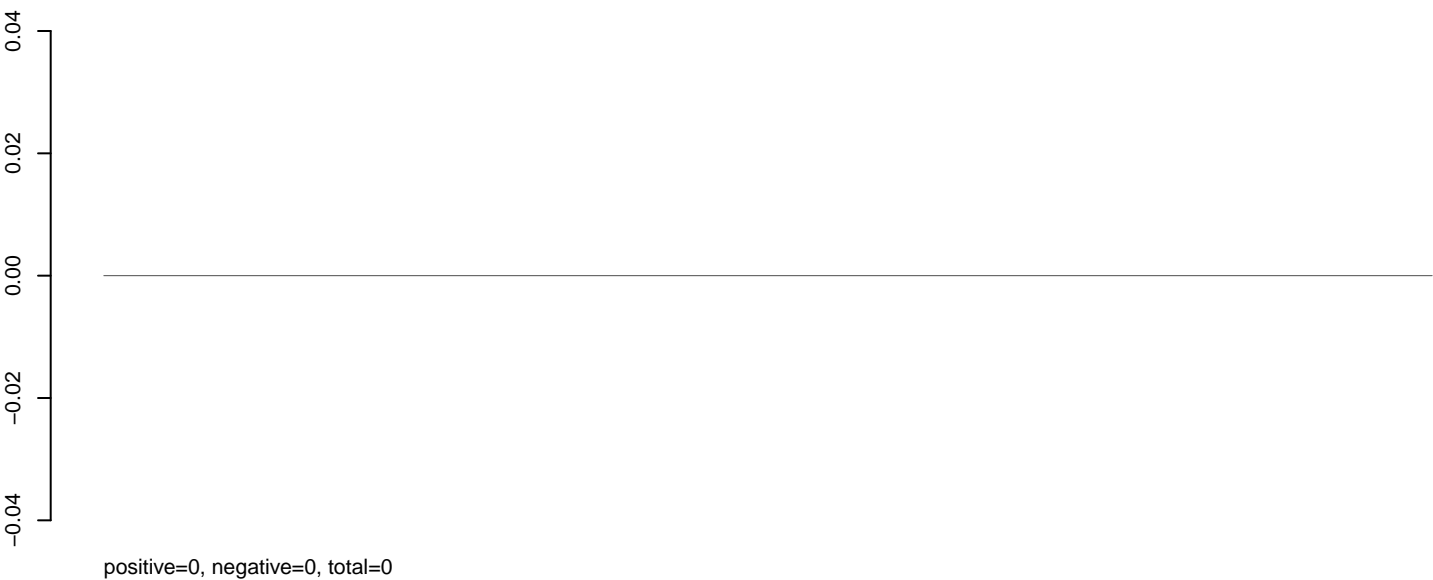
AeAeg_Aag2_cells.rep



AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

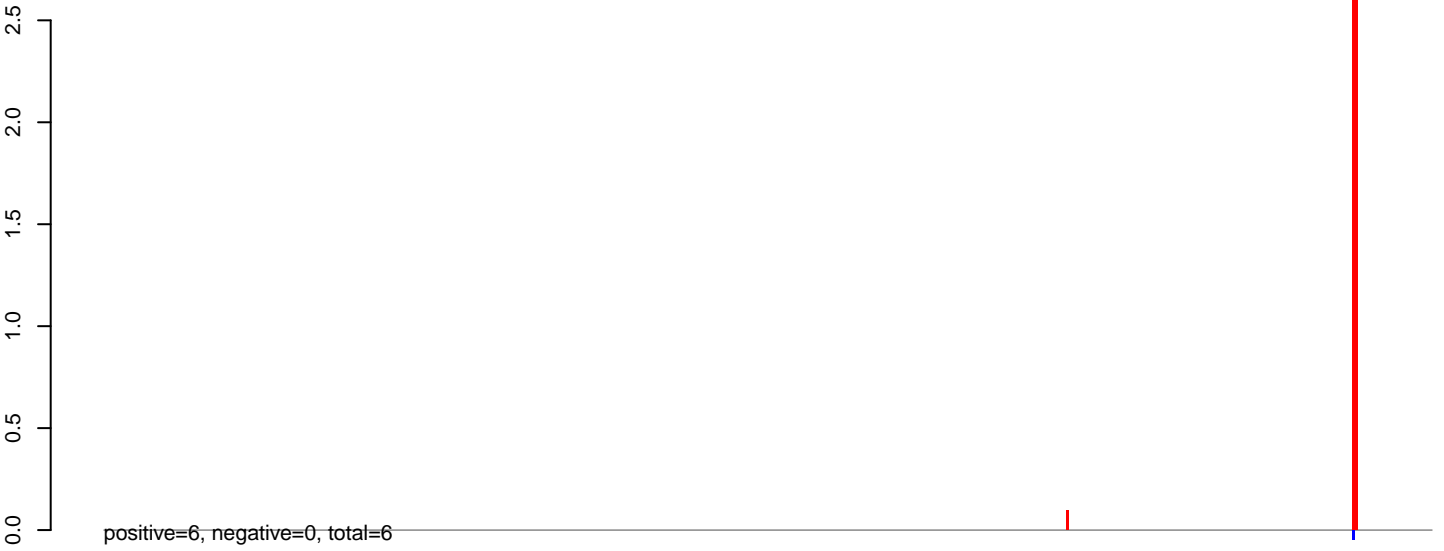


AeAeg_Aag2_cells.rep

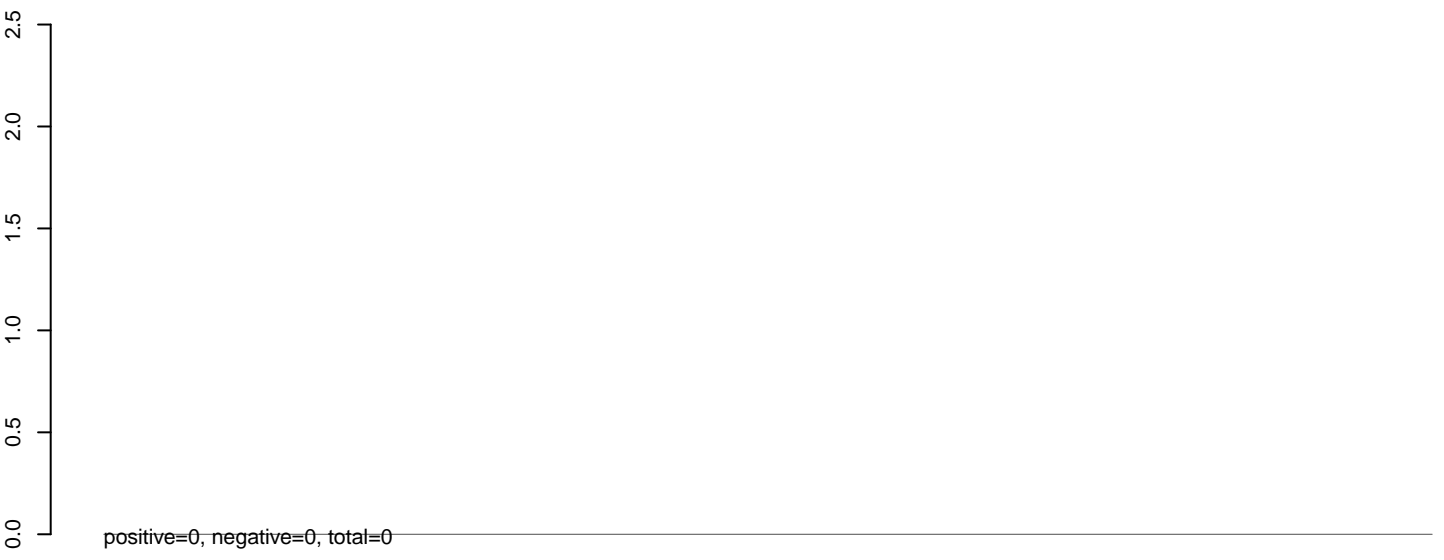


0 2000 4000 6000 8000 10000

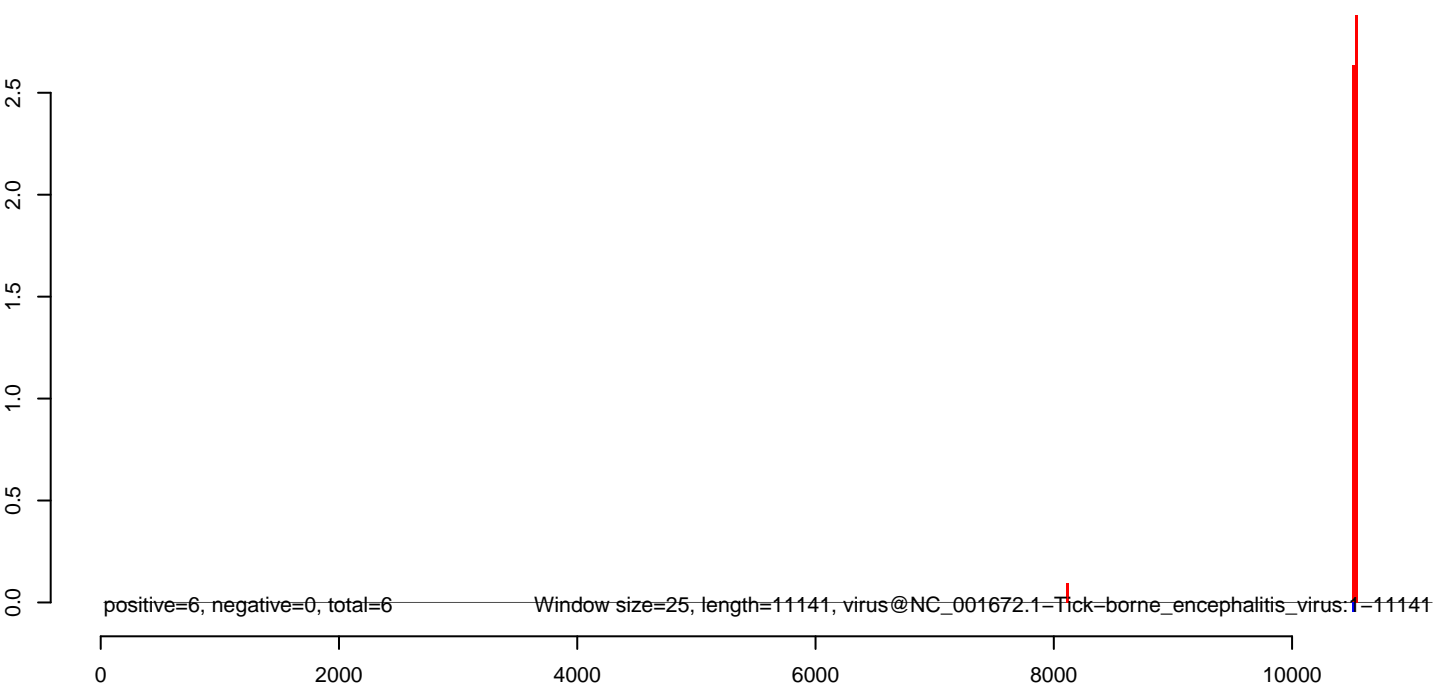
AeAeg_Aag2_cells.18_23.rep



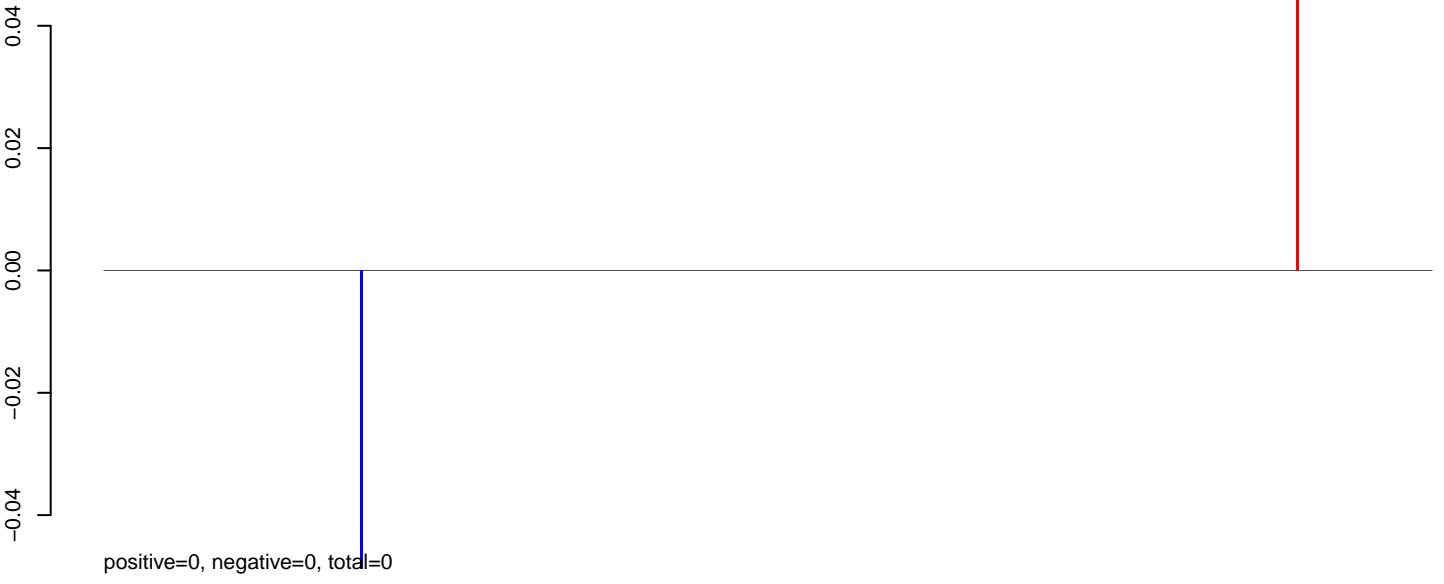
AeAeg_Aag2_cells.24_35.rep



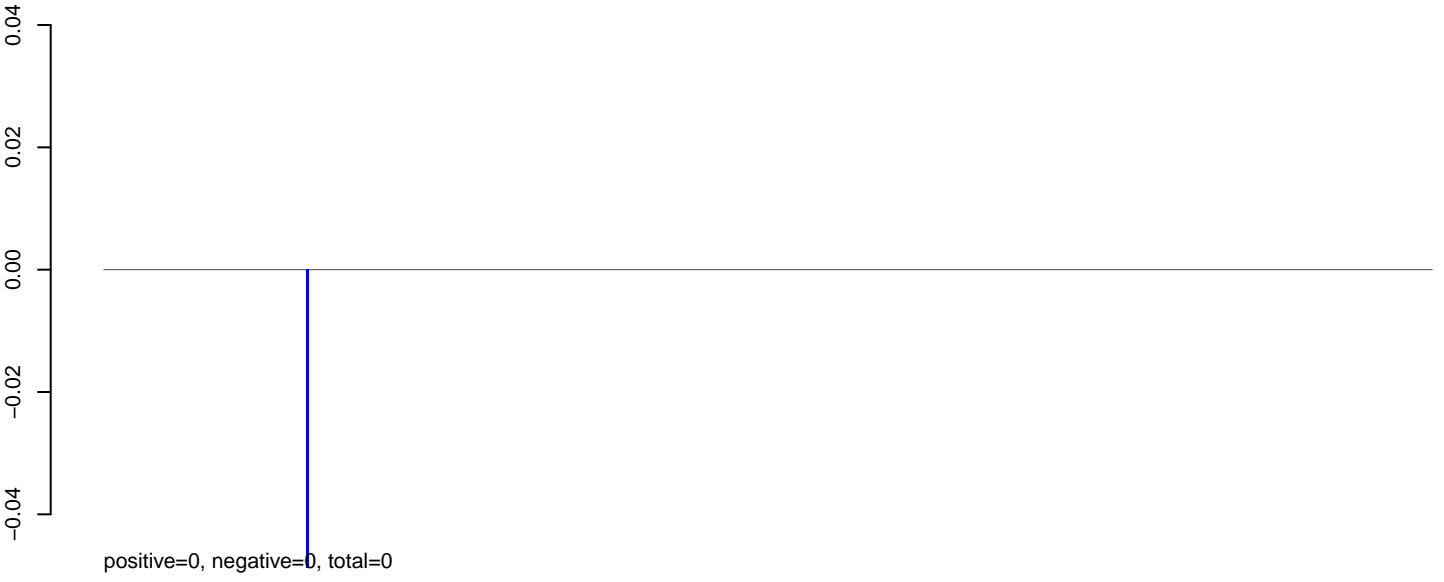
AeAeg_Aag2_cells.rep



AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

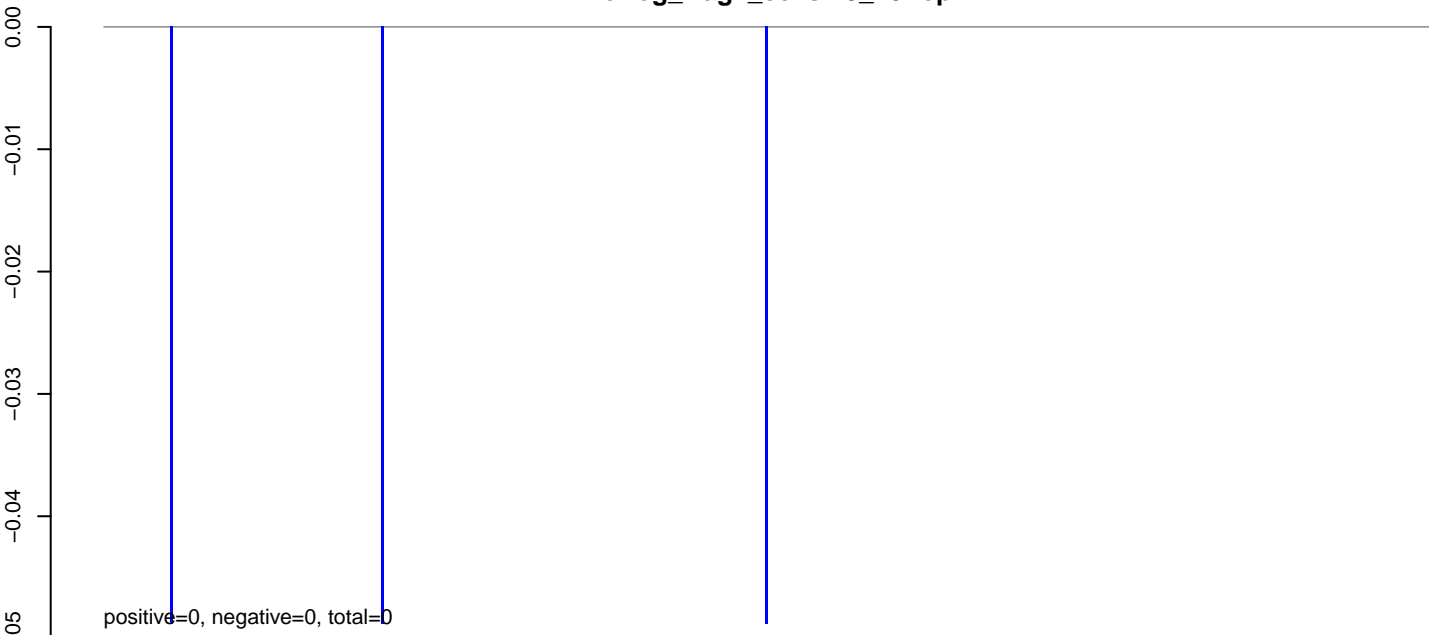


AeAeg_Aag2_cells.rep

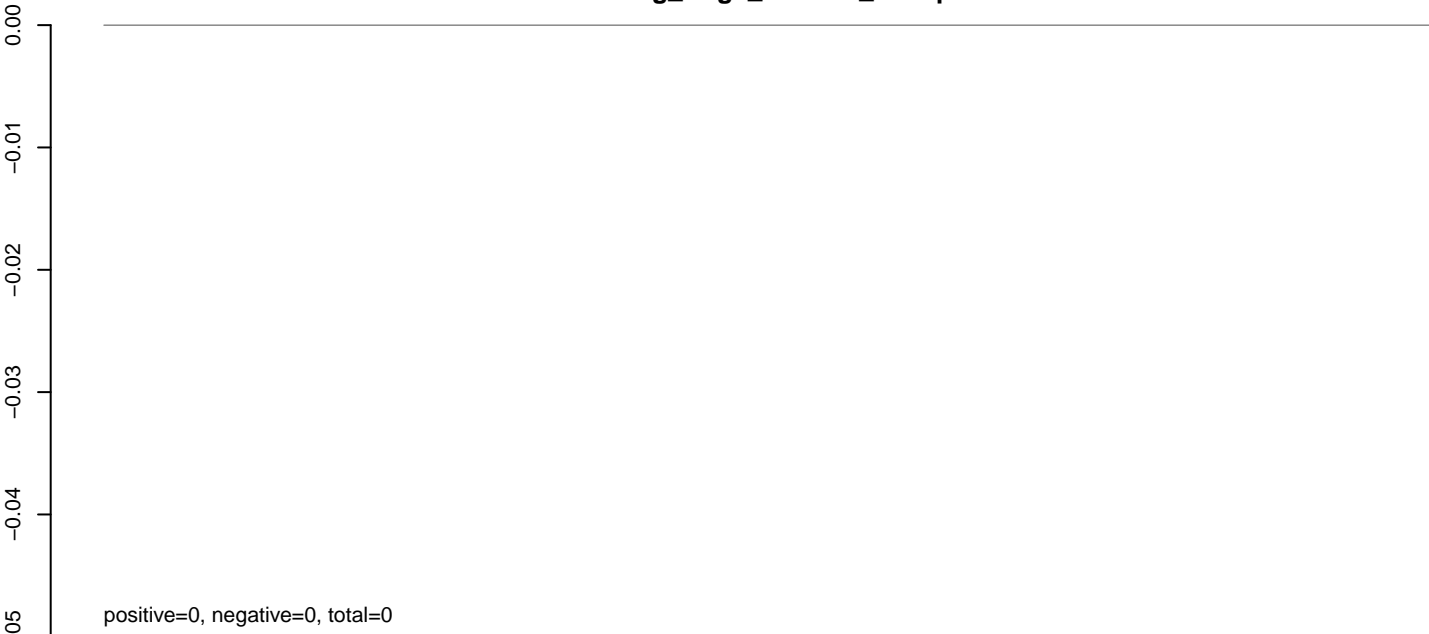


0 2000 4000 6000 8000 10000 12000

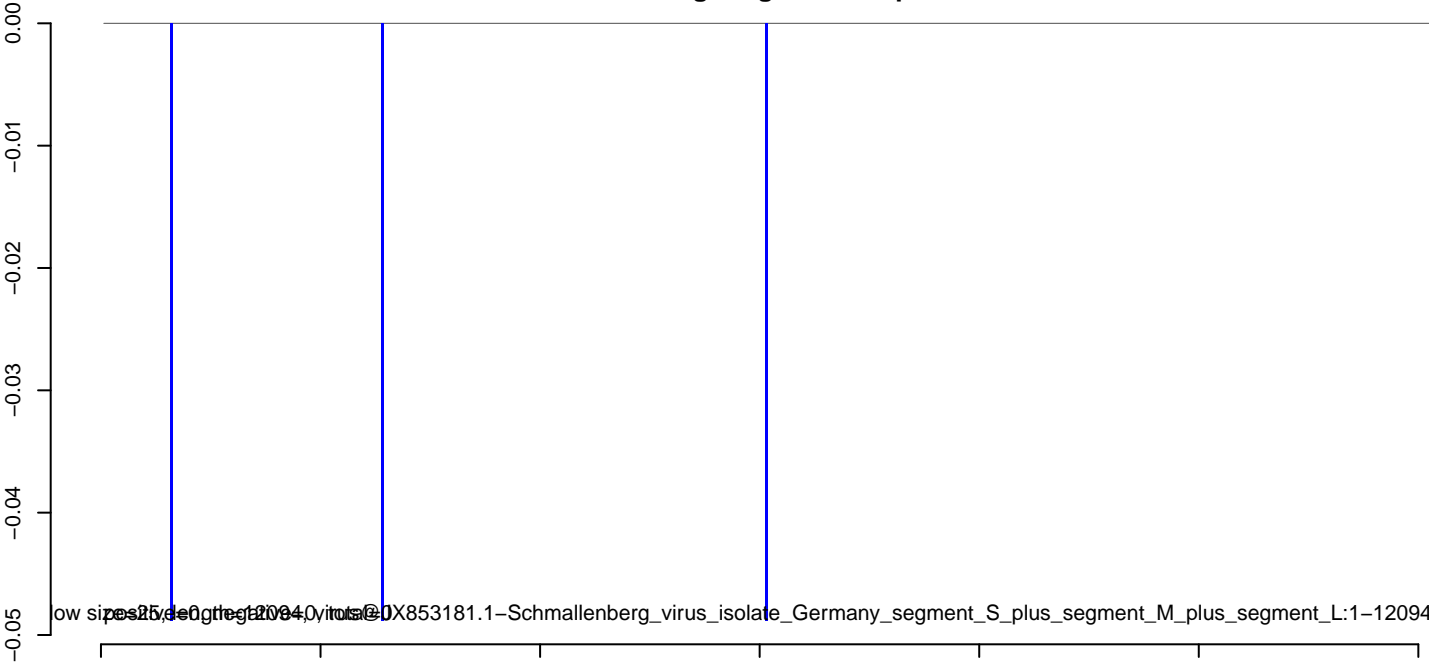
AeAeg_Aag2_cells.18_23.rep



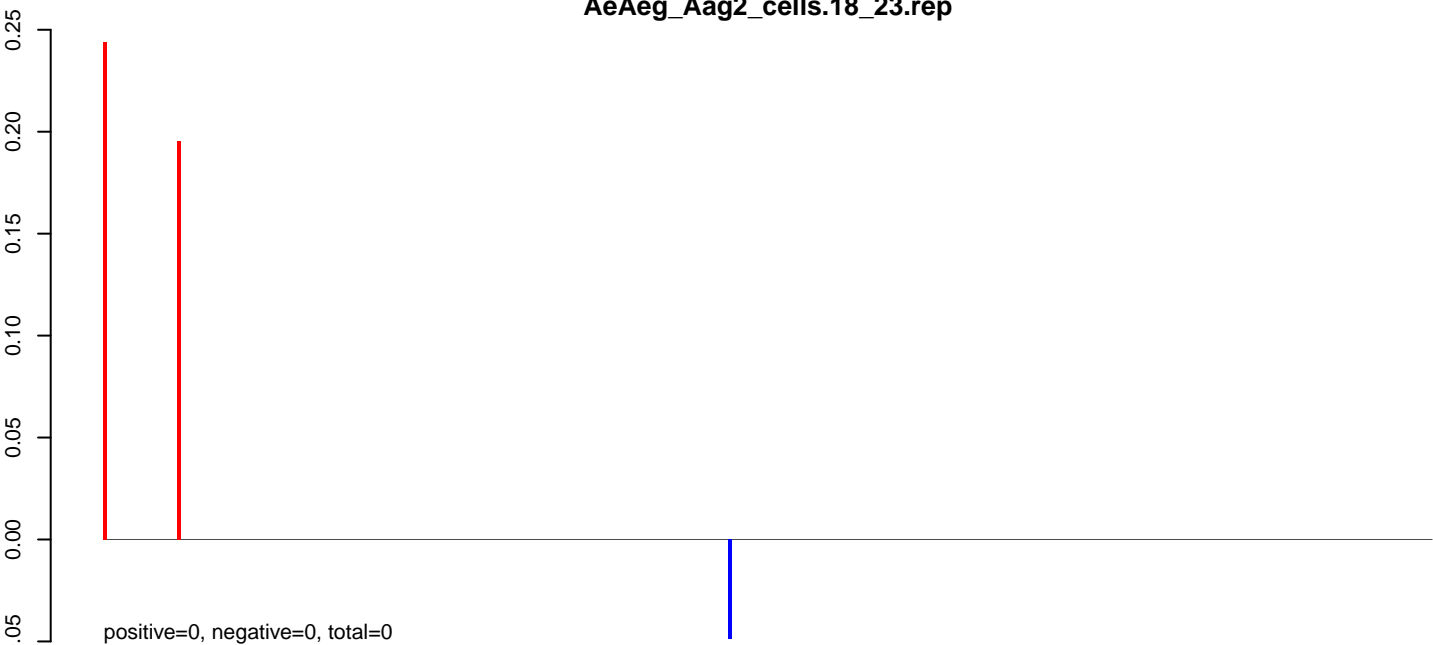
AeAeg_Aag2_cells.24_35.rep



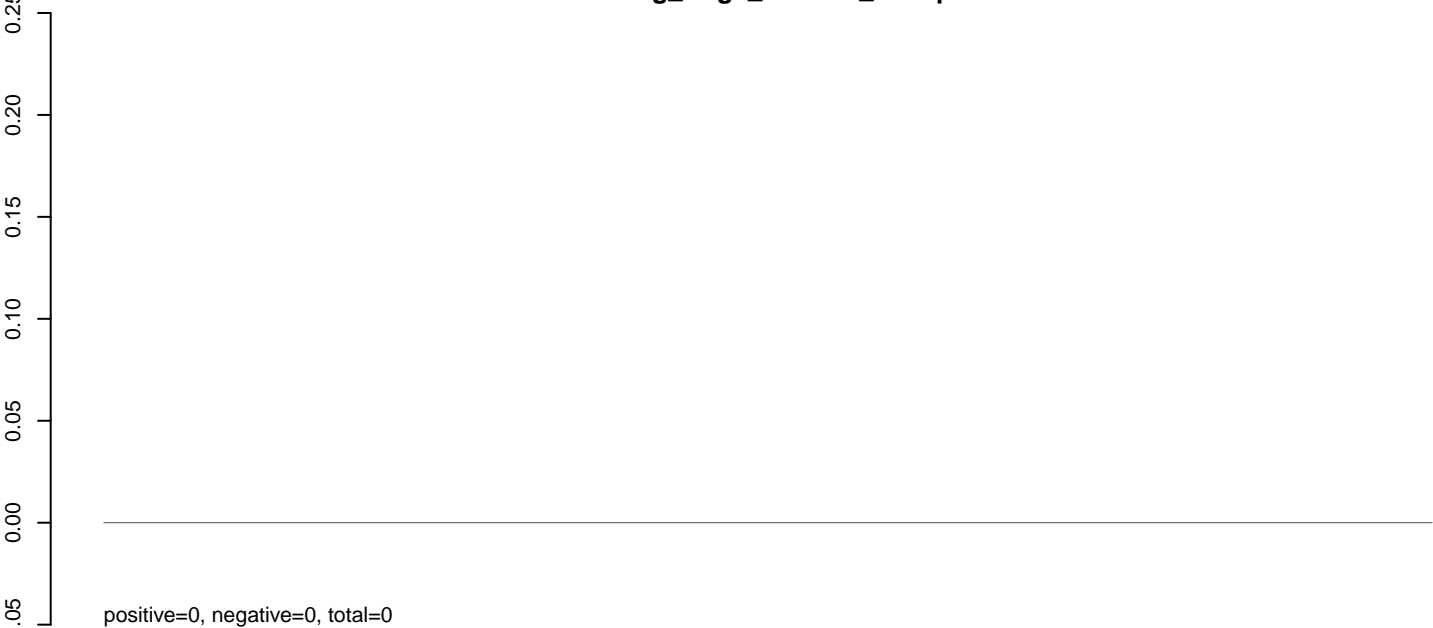
AeAeg_Aag2_cells.rep



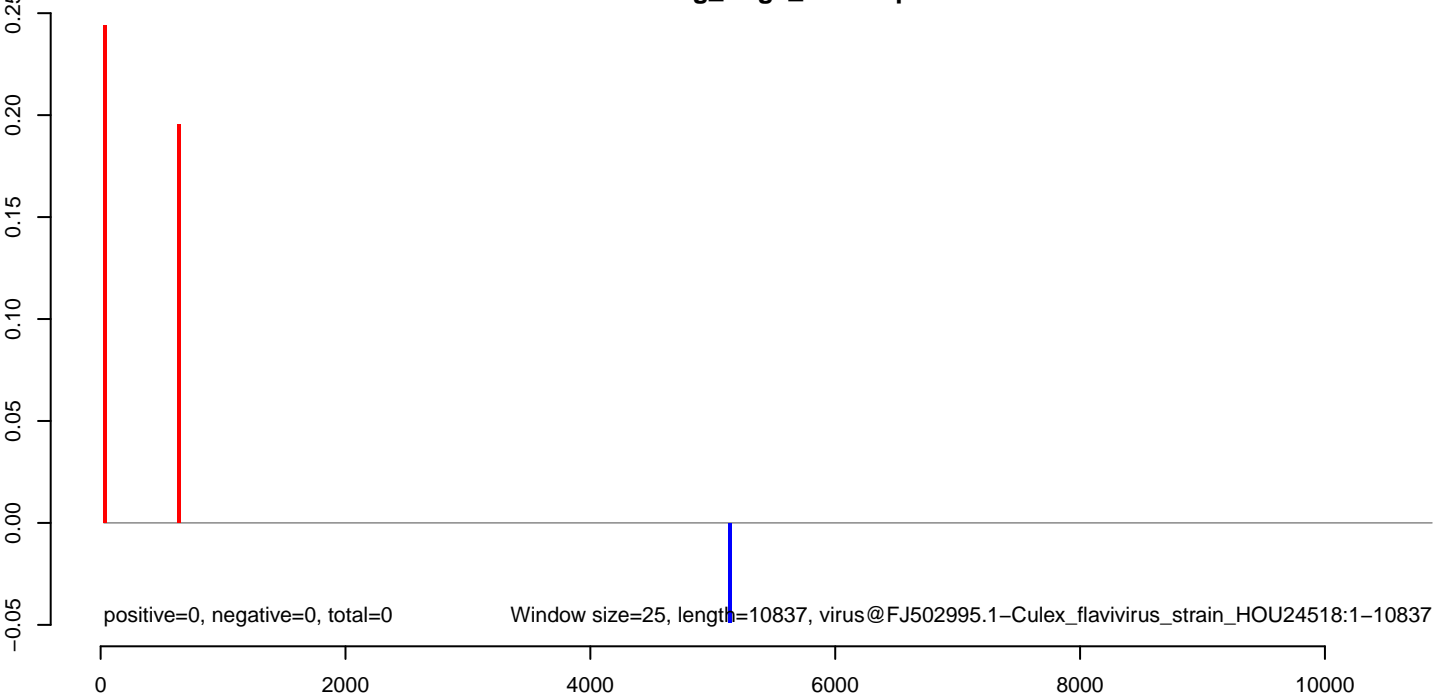
AeAeg_Aag2_cells.18_23.rep



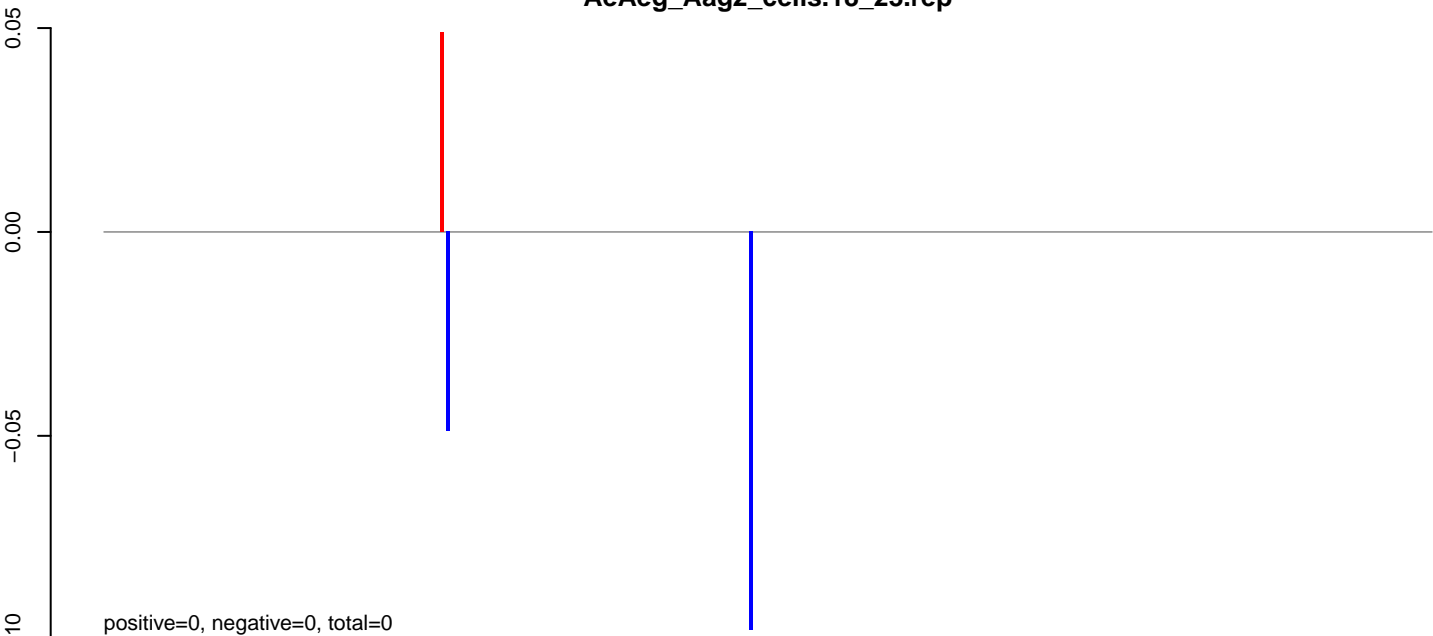
AeAeg_Aag2_cells.24_35.rep



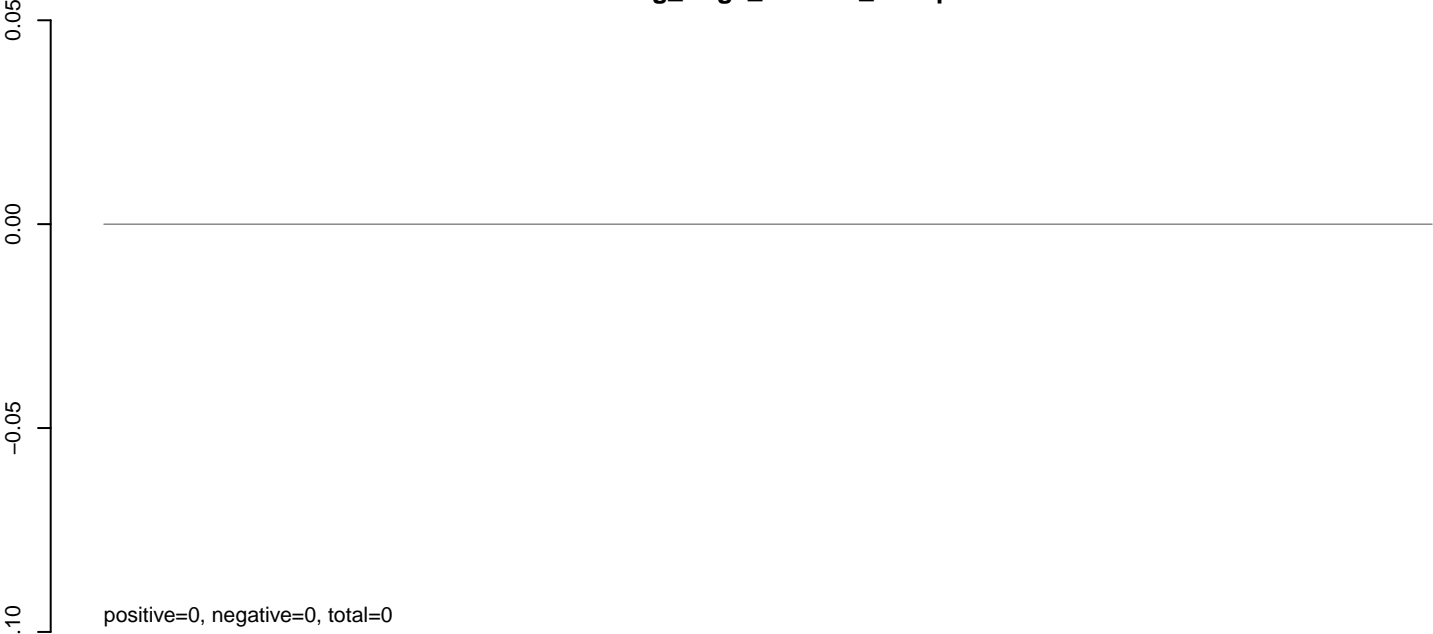
AeAeg_Aag2_cells.rep



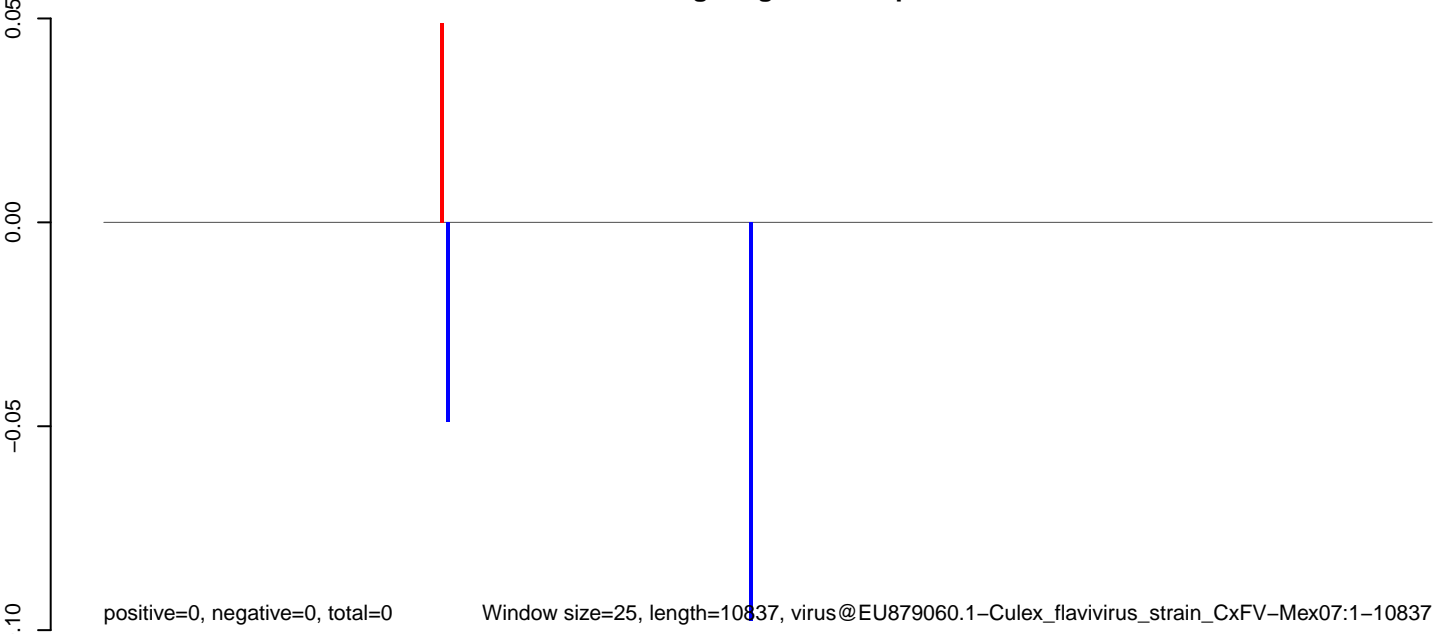
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

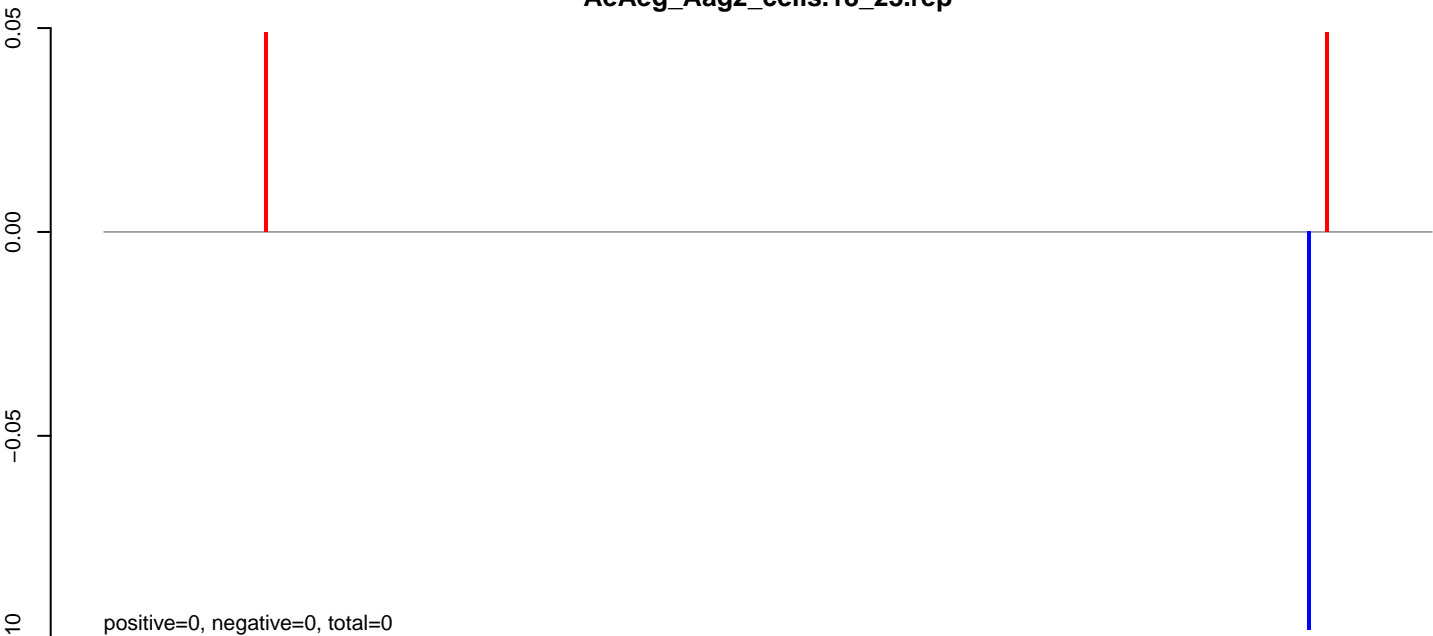


AeAeg_Aag2_cells.rep



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

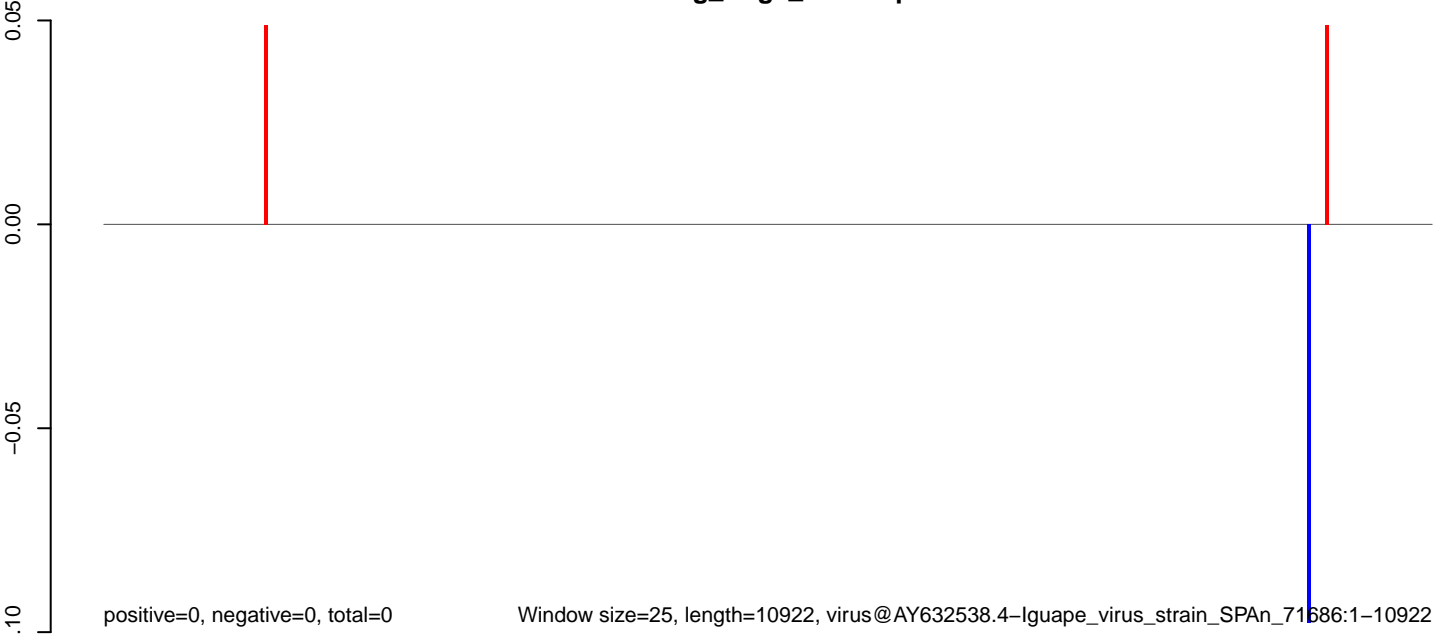
AeAeg_Aag2_cells.18_23.rep



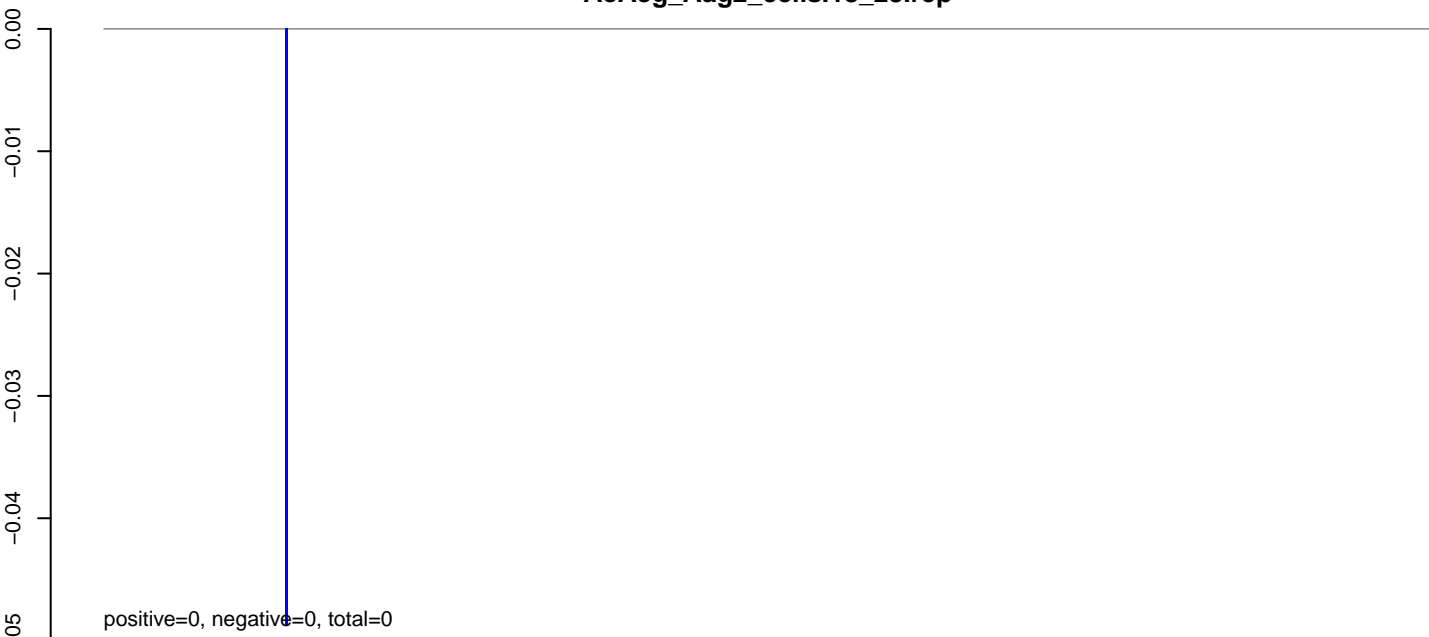
AeAeg_Aag2_cells.24_35.rep



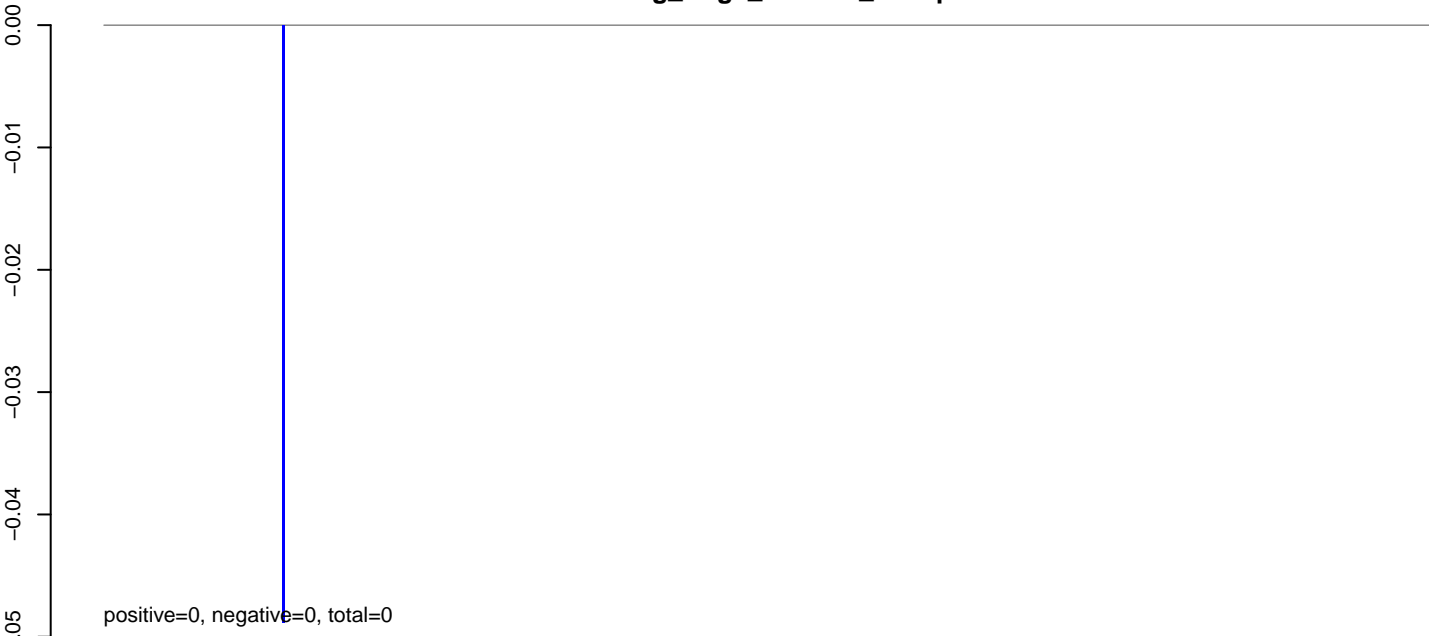
AeAeg_Aag2_cells.rep



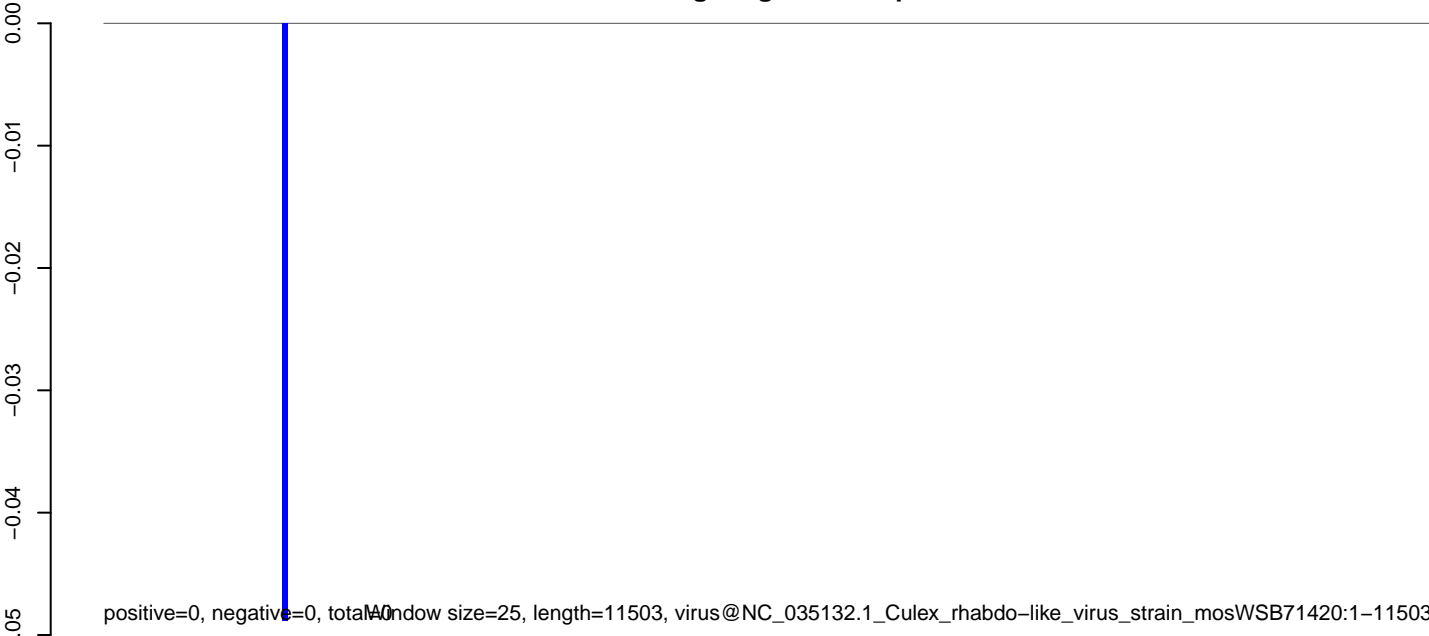
AeAeg_Aag2_cells.18_23.rep



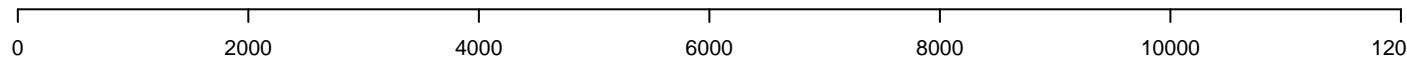
AeAeg_Aag2_cells.24_35.rep



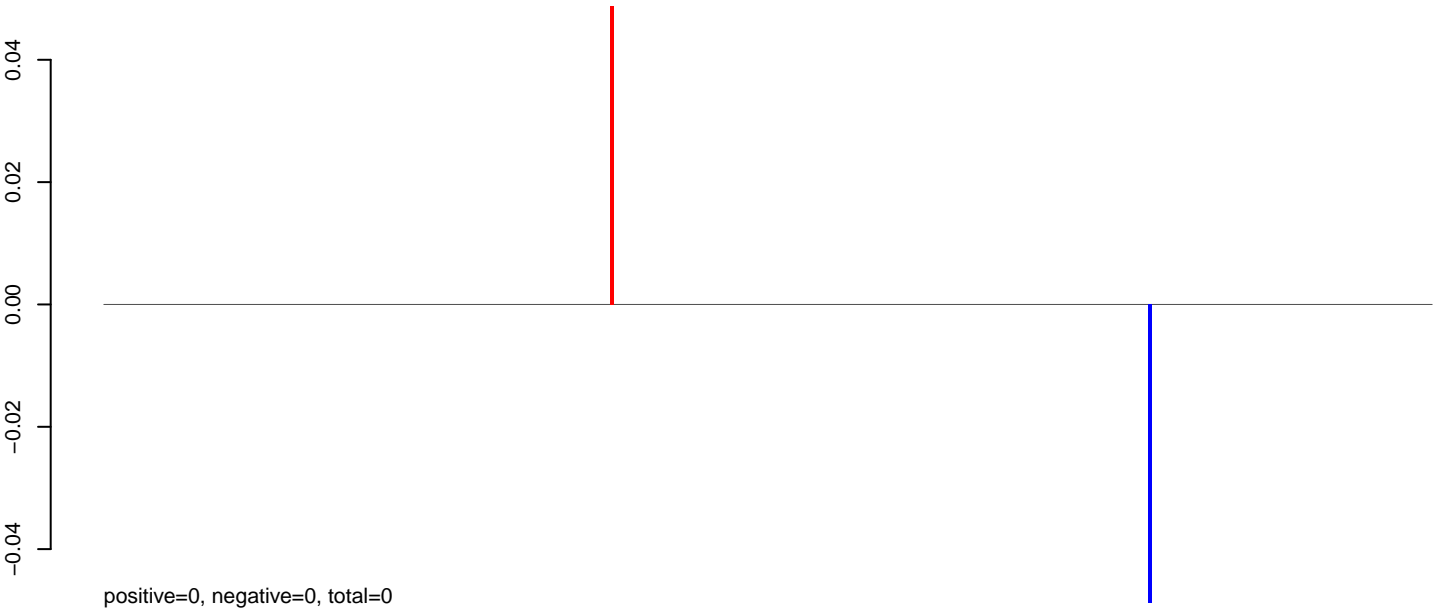
AeAeg_Aag2_cells.rep



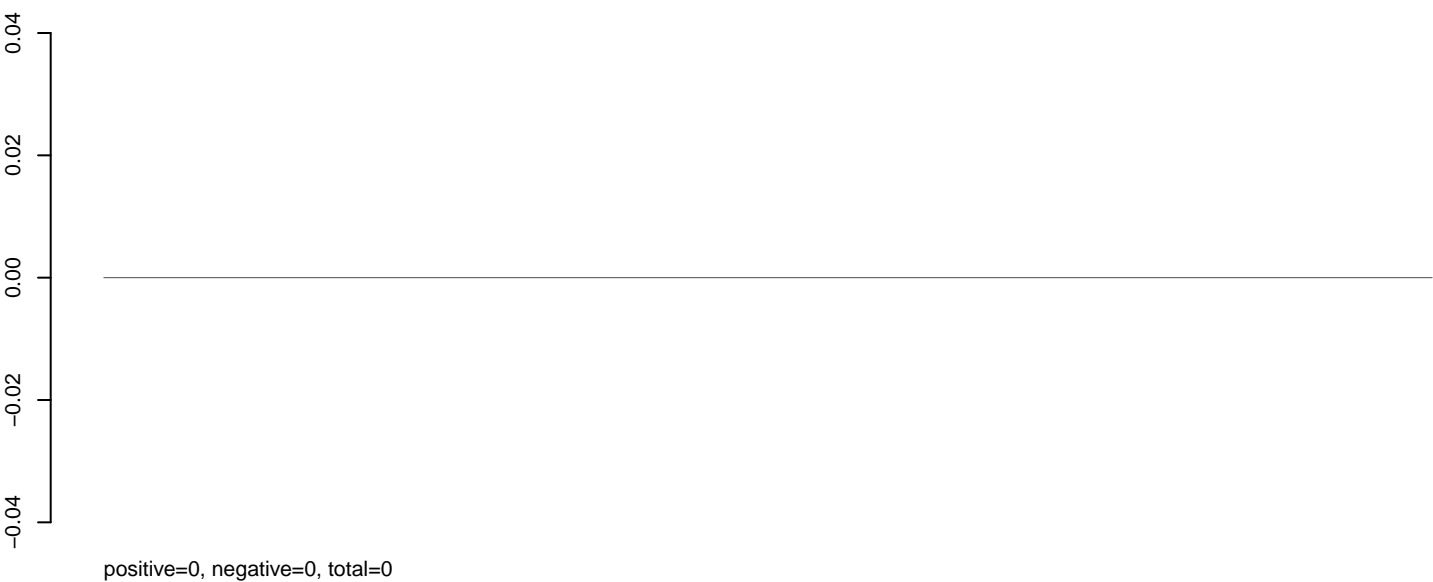
Window size=25, length=11503, virus@NC_035132.1_Culex_rhabdo-like_virus_strain_mosWSB71420:1-11503



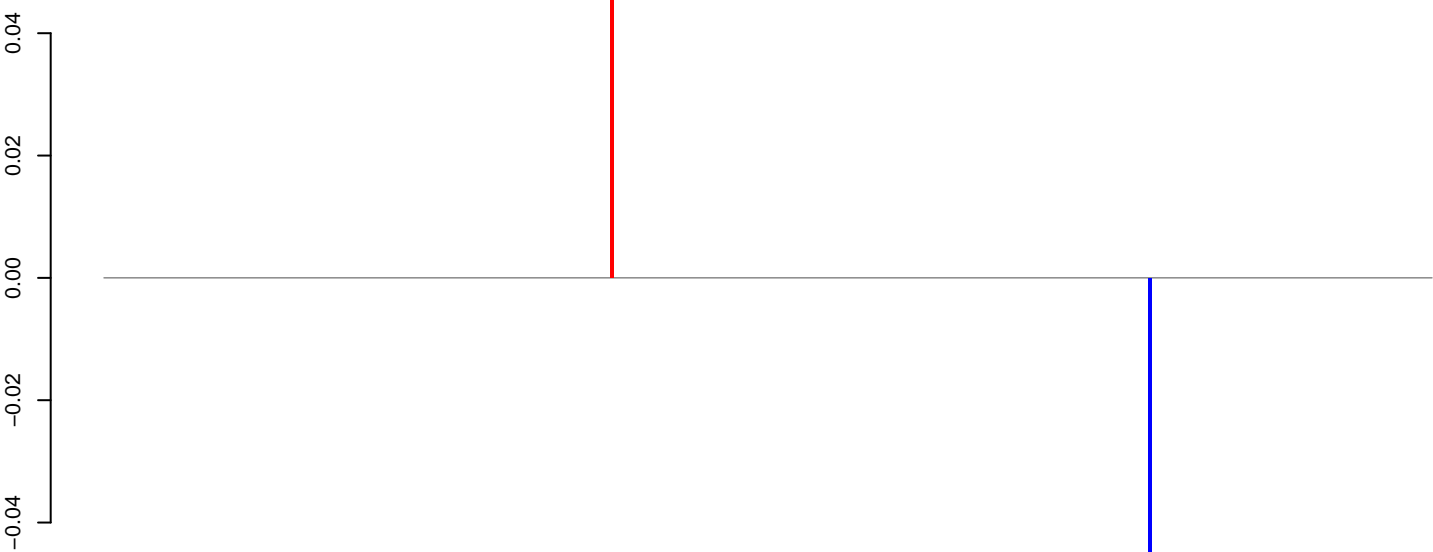
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



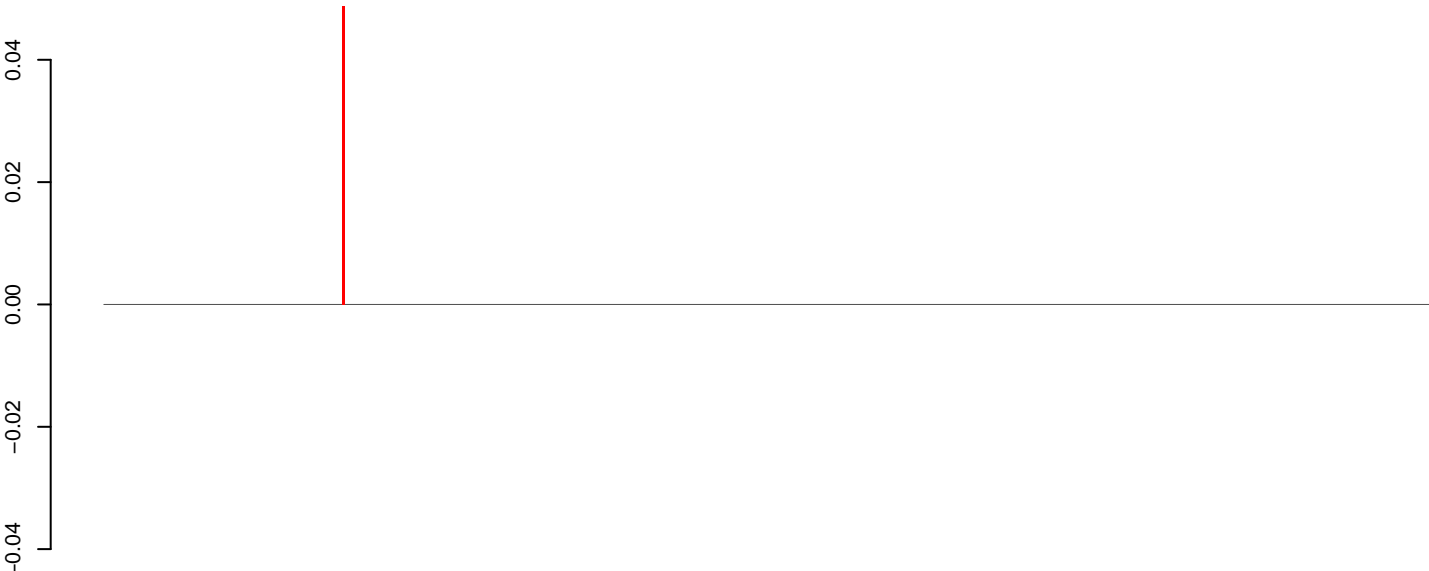
AeAeg_Aag2_cells.rep



bow size=25, length=1000, interval=40 NC_034454.1-Fukuoka_virus_nucleoprotein_phosphoprotein_matrix_hypothetical_protein_glycoprotei:1-10863

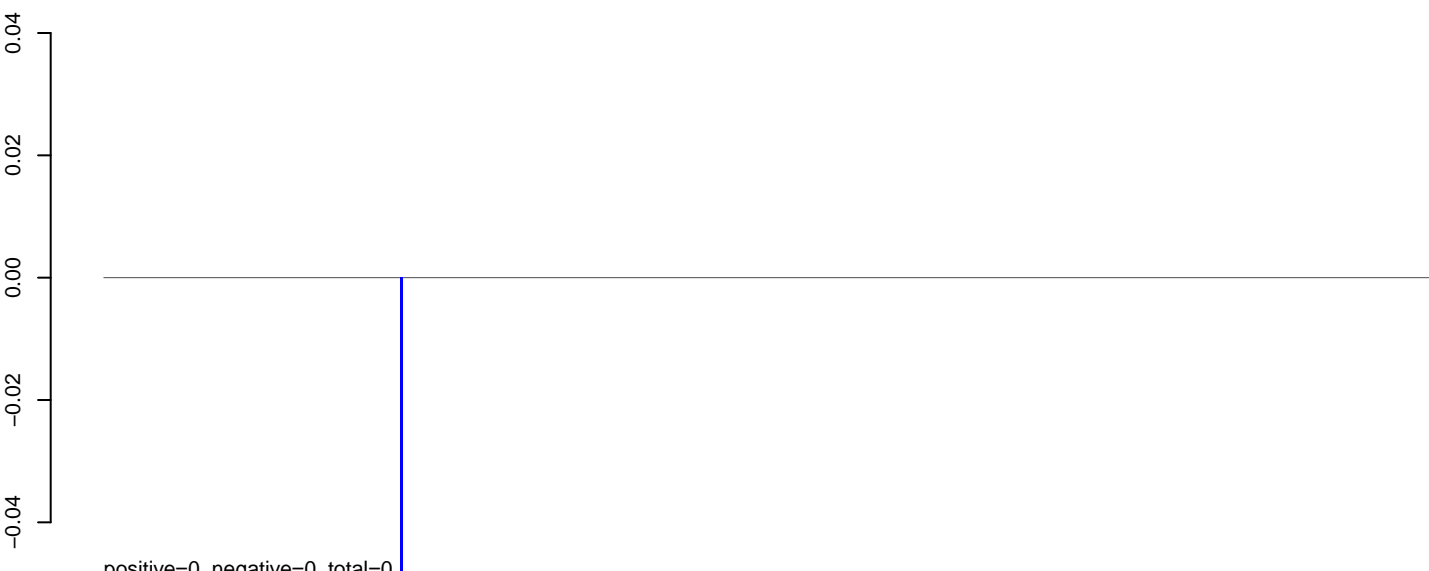
0 2000 4000 6000 8000 10000

AeAeg_Aag2_cells.18_23.rep



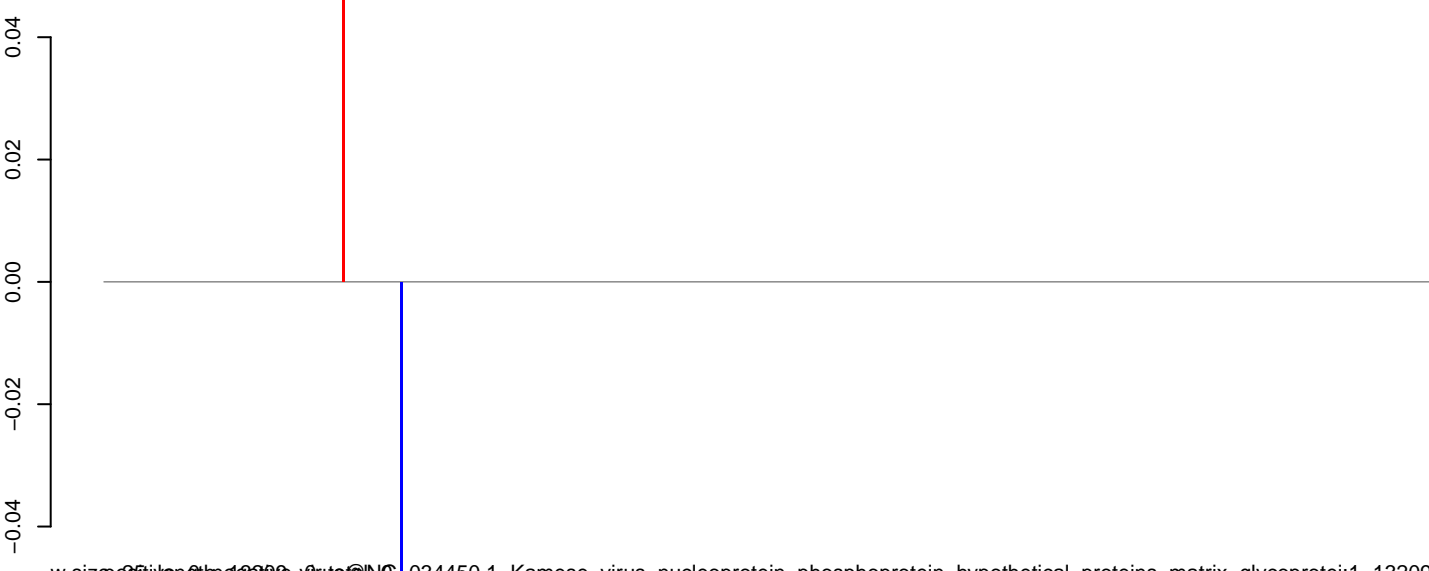
positive=0, negative=0, total=0

AeAeg_Aag2_cells.24_35.rep



positive=0, negative=0, total=0

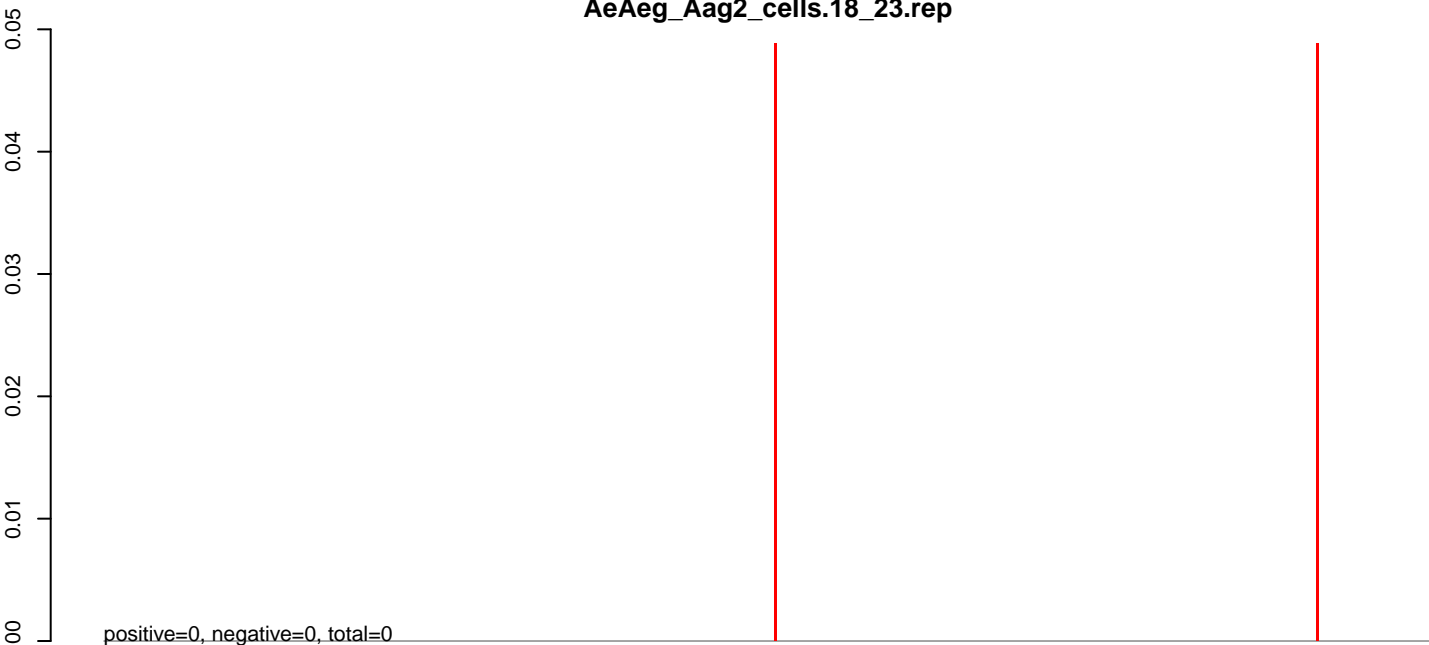
AeAeg_Aag2_cells.rep



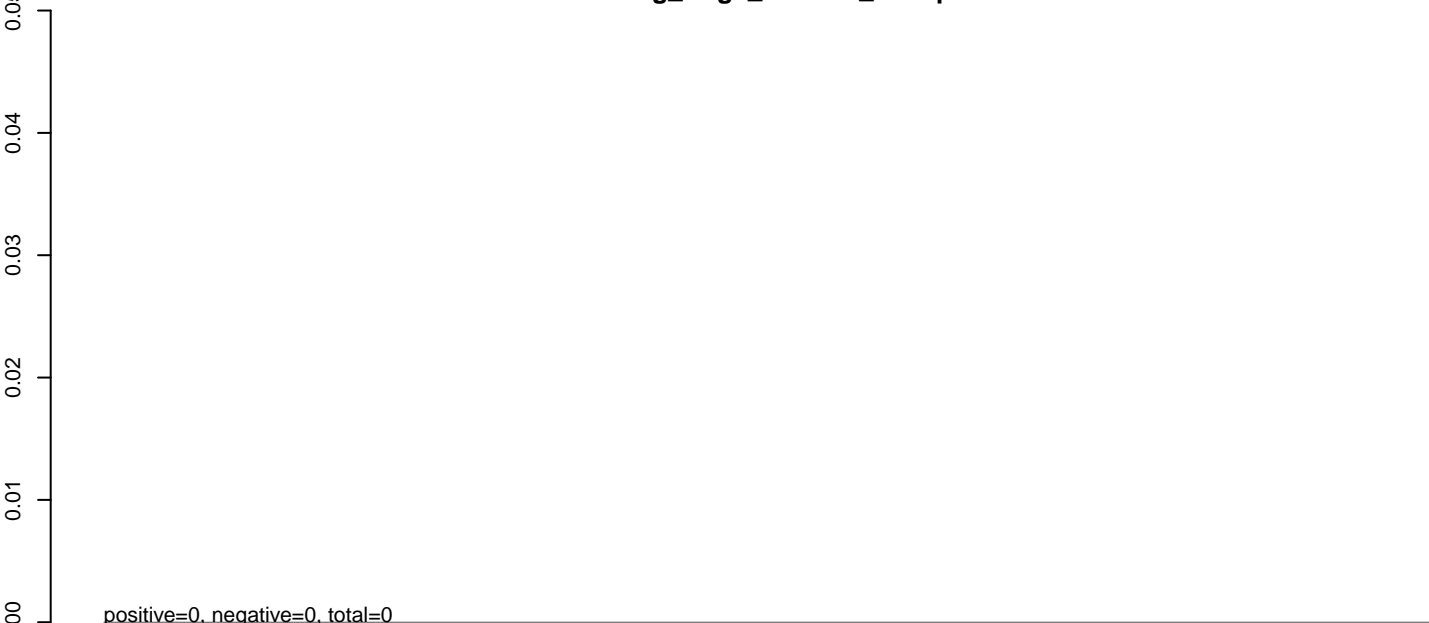
w size=25, neg=0, pos=0, total=0, use@NC_034450.1-Kamese_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycoprotei:1-13209

0 2000 4000 6000 8000 10000 12000

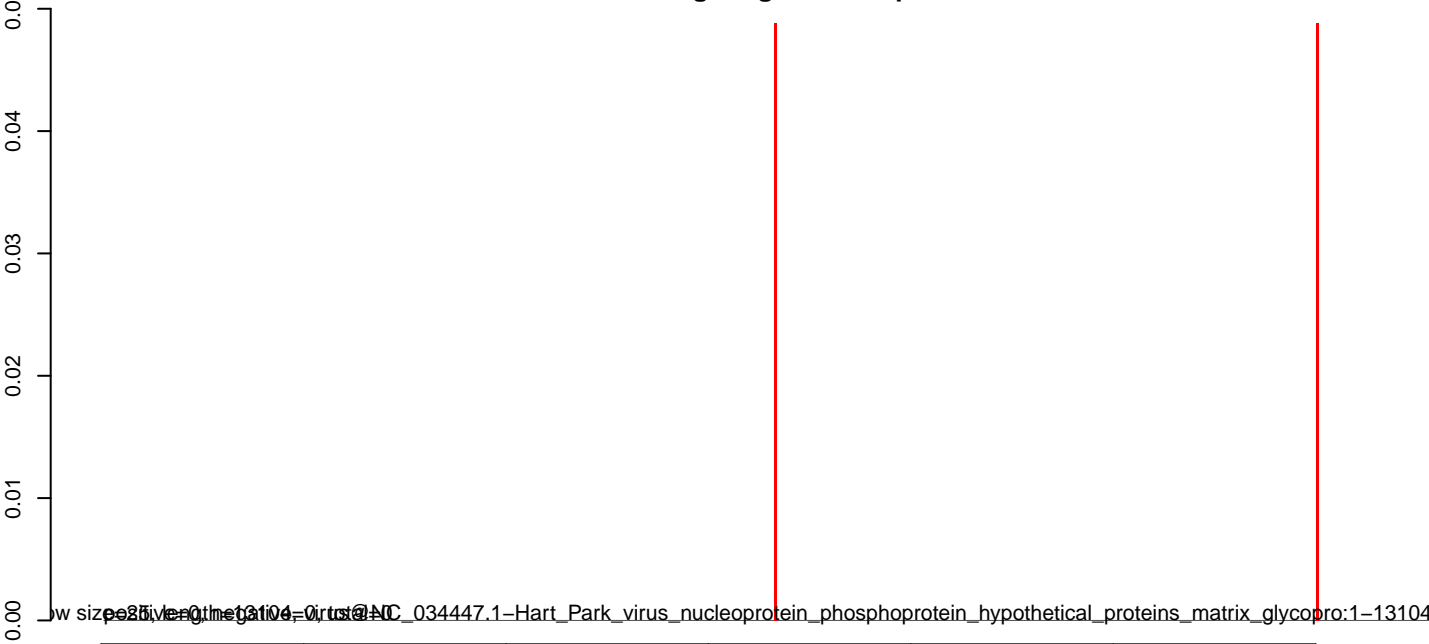
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

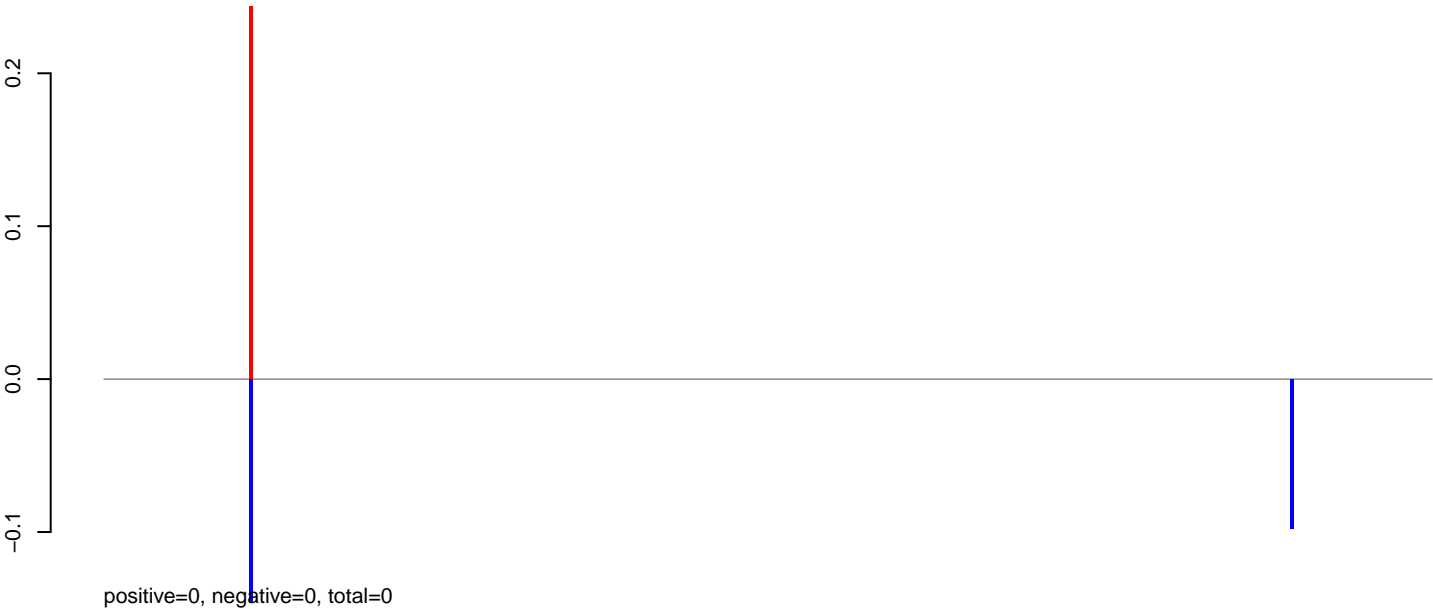


AeAeg_Aag2_cells.rep

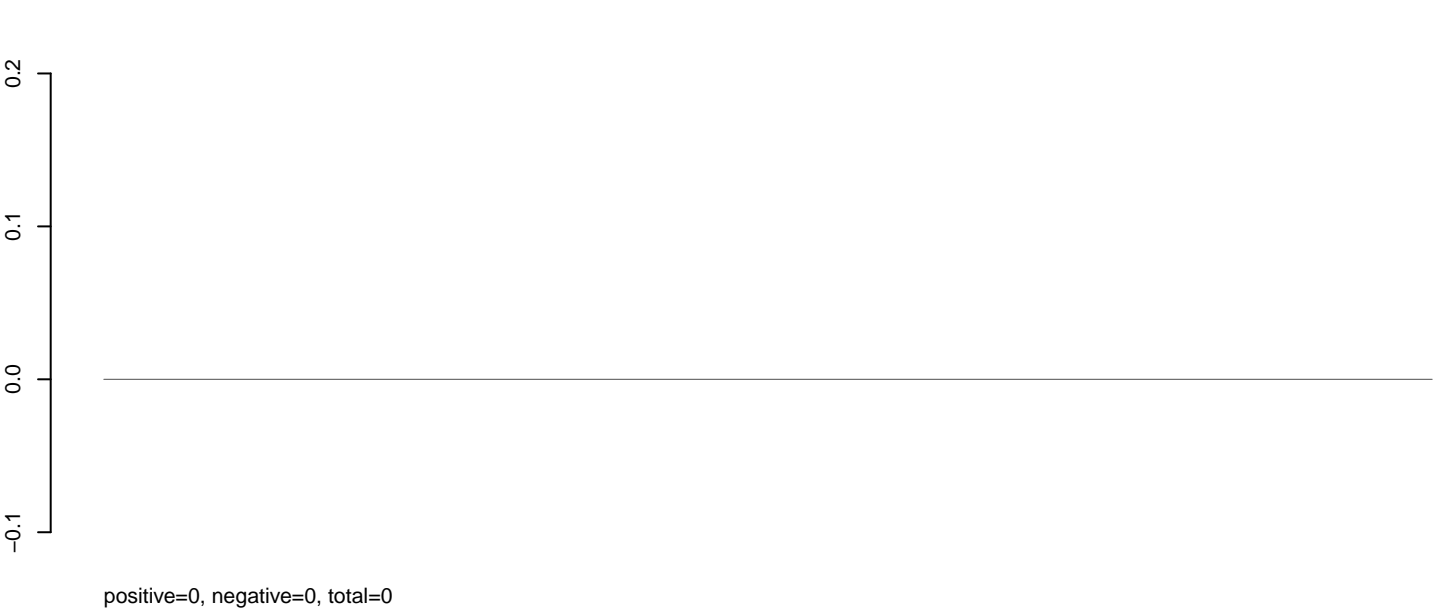


low size positive=0, negative=0, total=0
NC_034447.1-Hart_Park_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopro:1-13104

AeAeg_Aag2_cells.18_23.rep



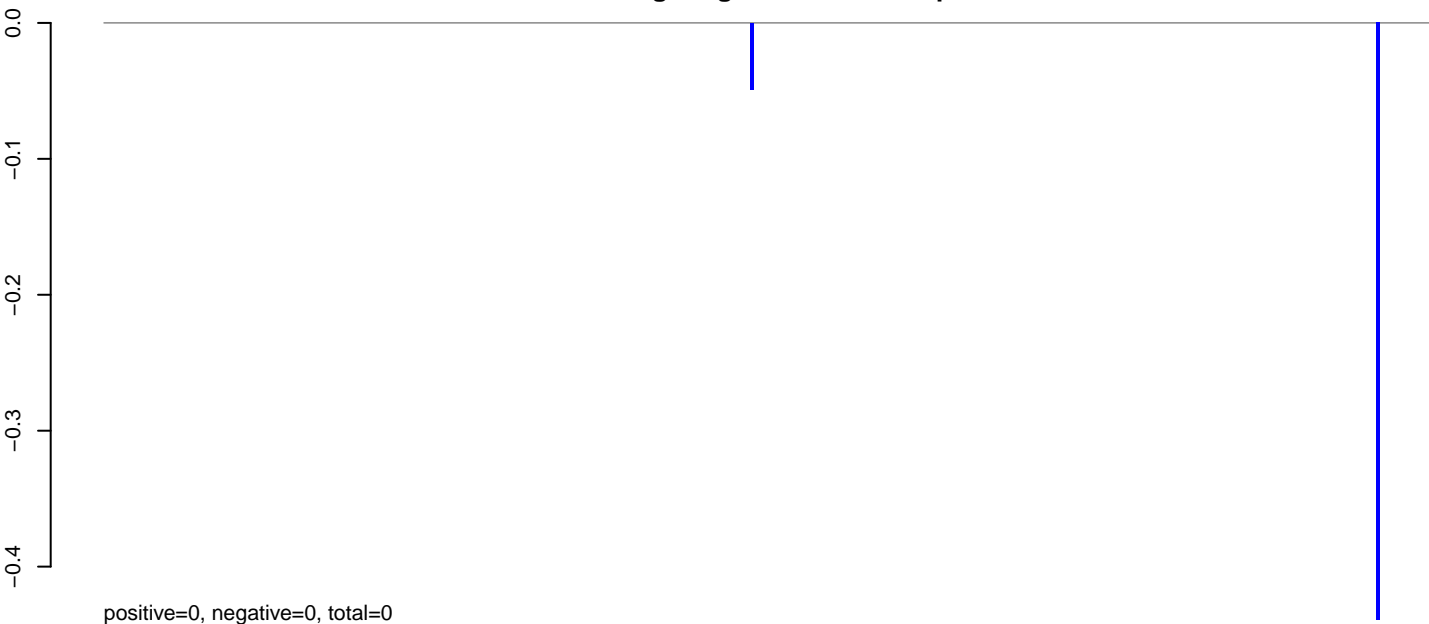
AeAeg_Aag2_cells.24_35.rep



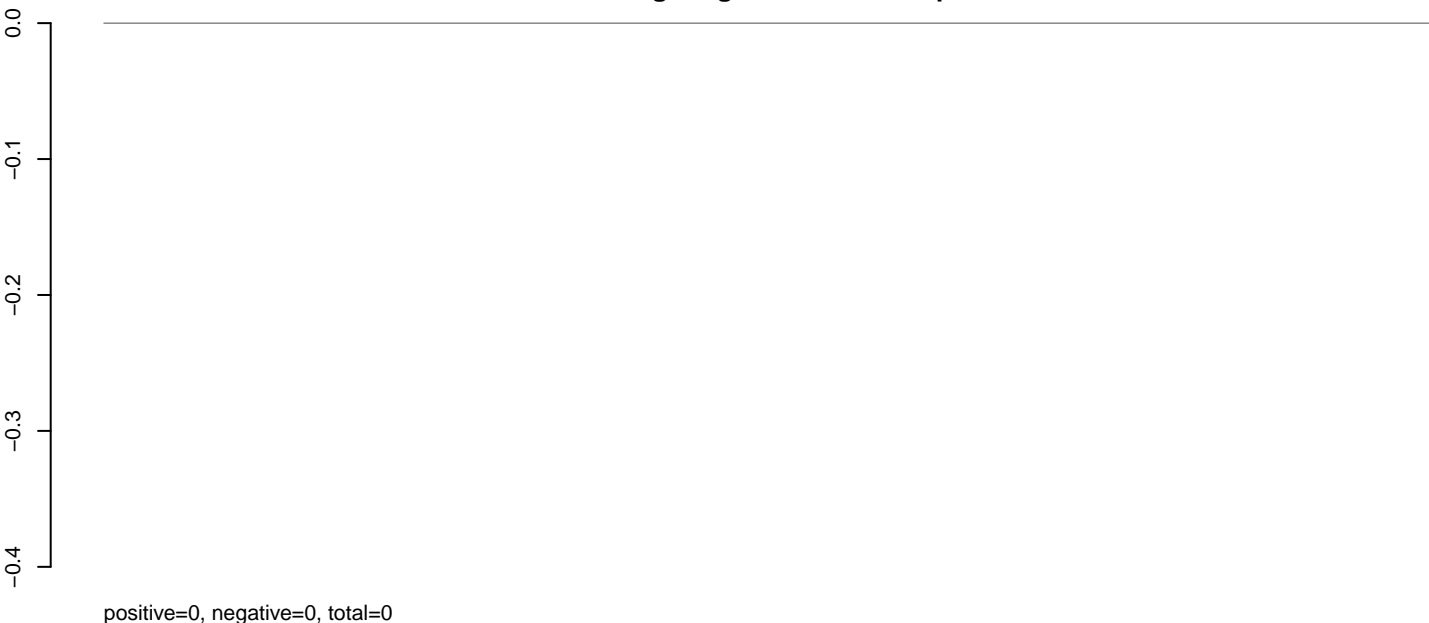
AeAeg_Aag2_cells.rep



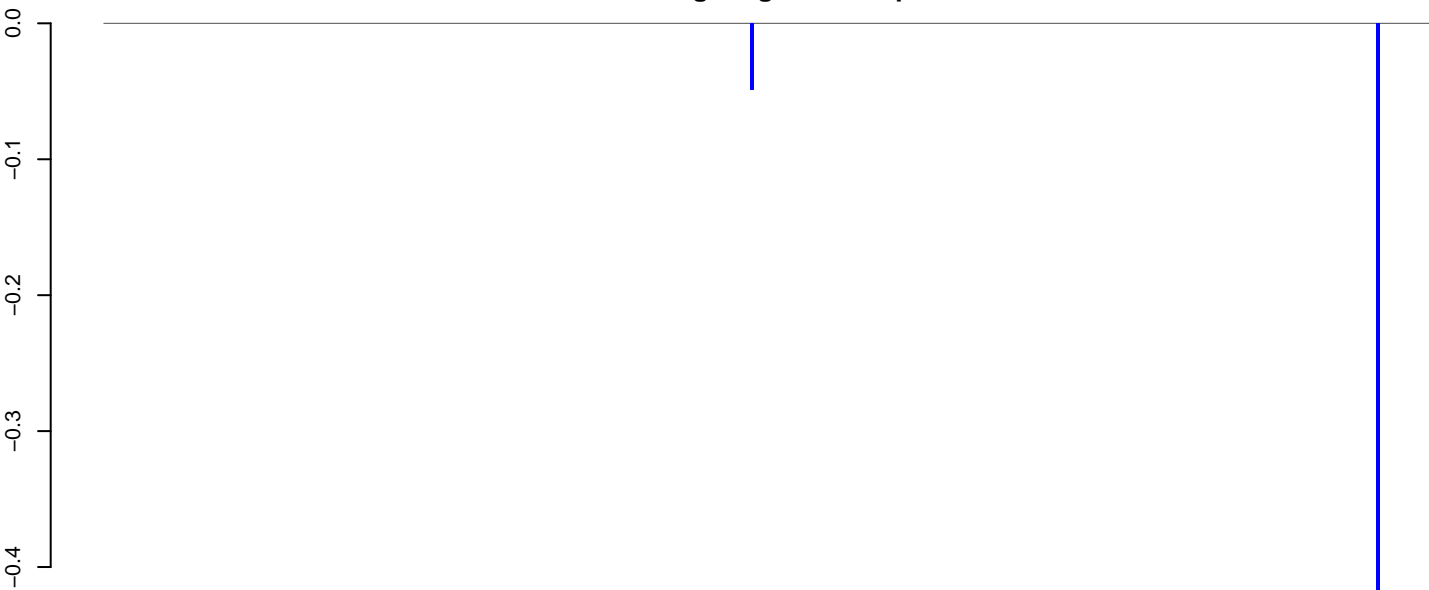
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



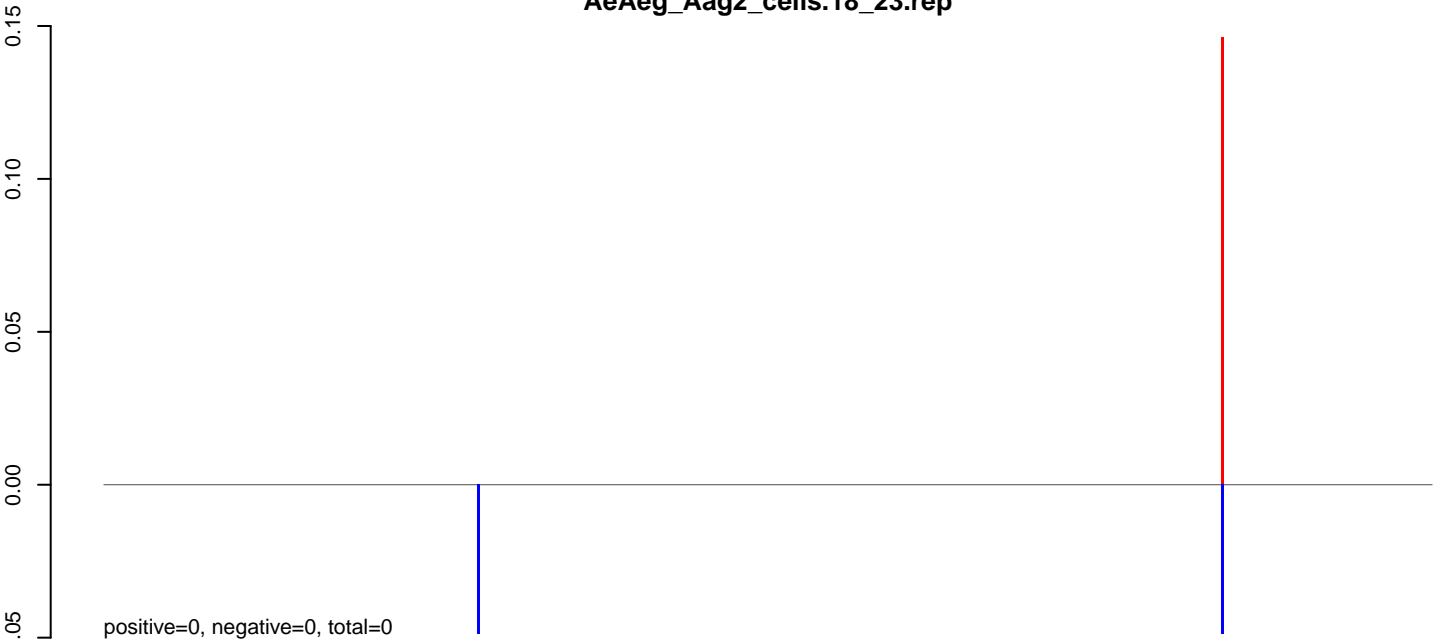
AeAeg_Aag2_cells.rep



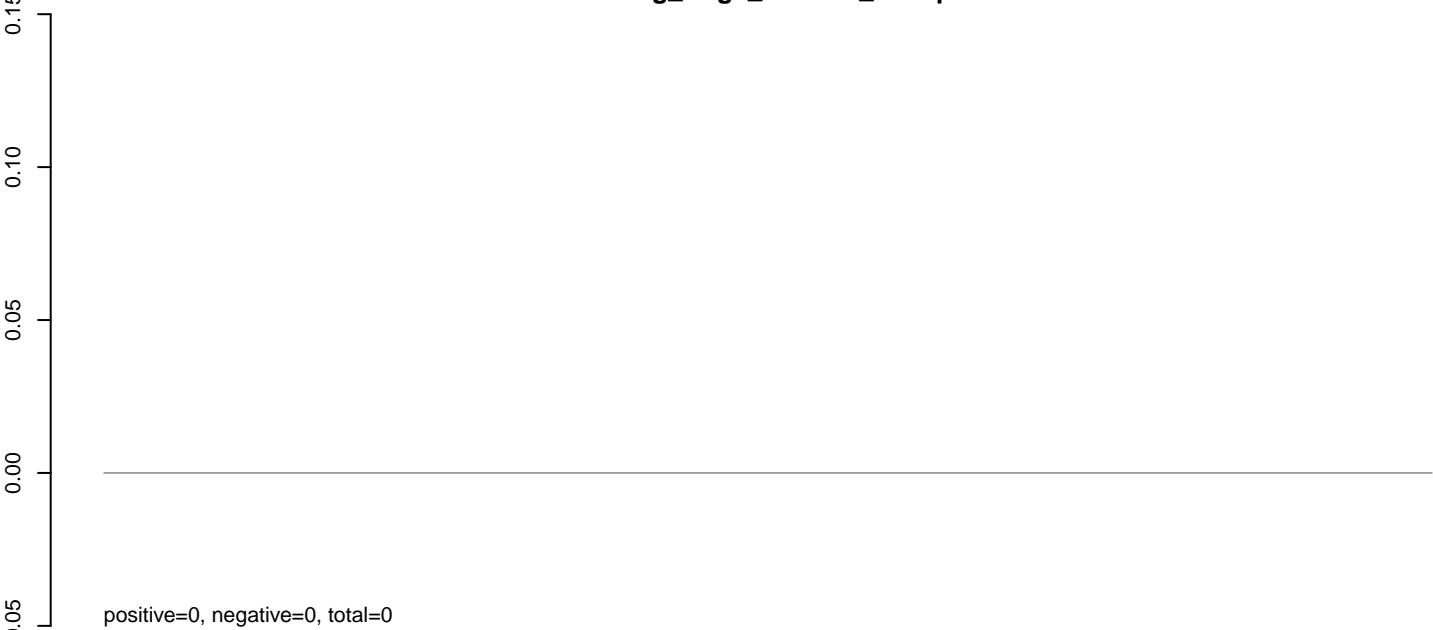
Window size=25, length=10173, virus@NC_033699.1-Jugra_virus_strain_P-9-314_polyprotein_gene:1-10173

0 2000 4000 6000 8000 10000

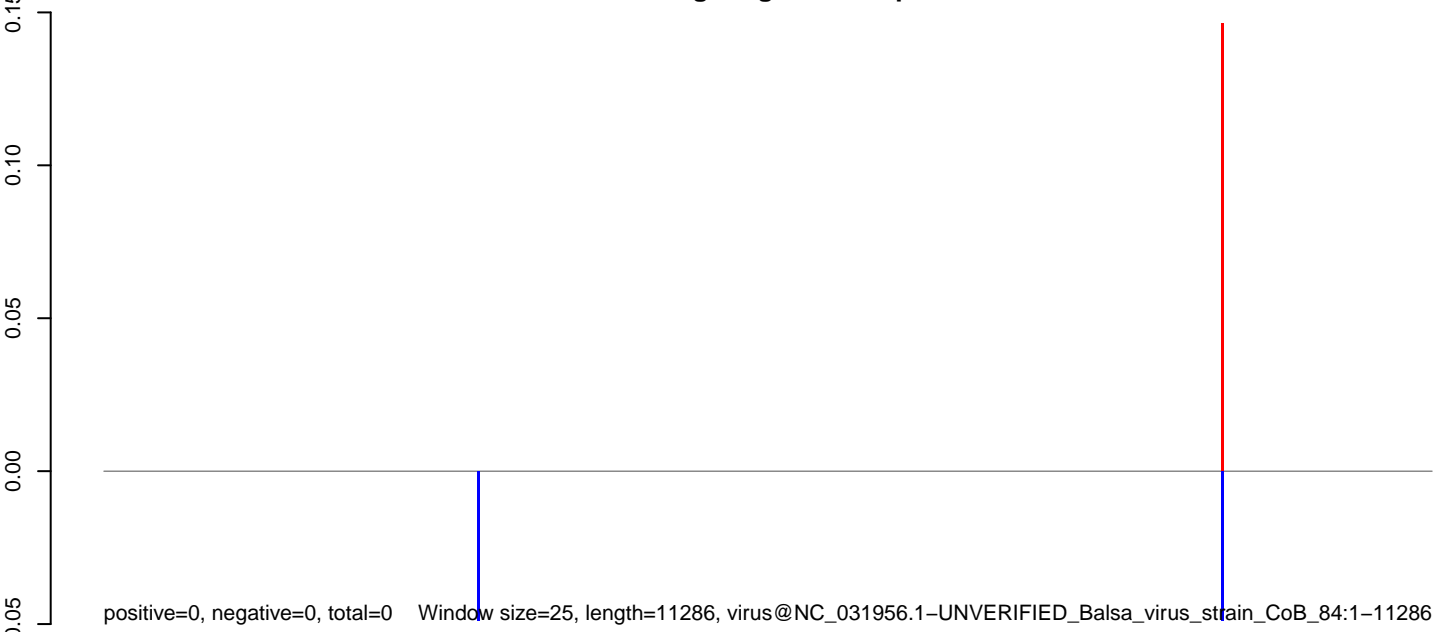
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep

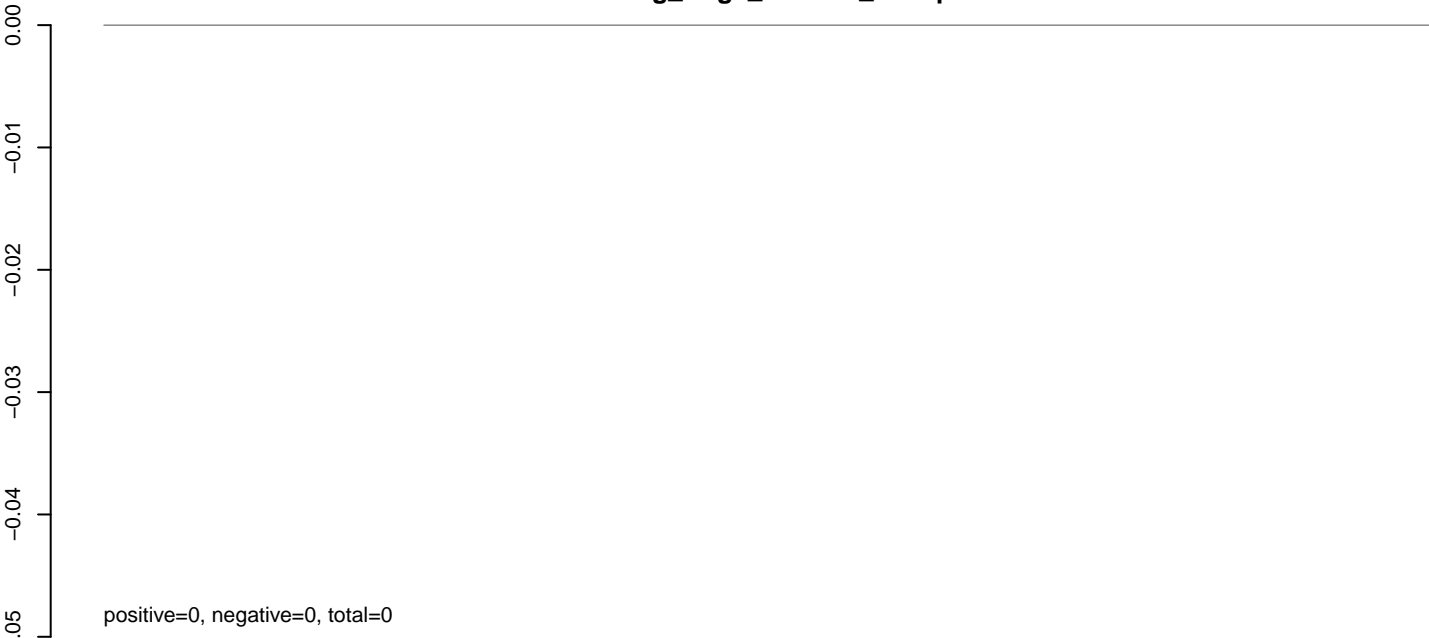


0 2000 4000 6000 8000 10000

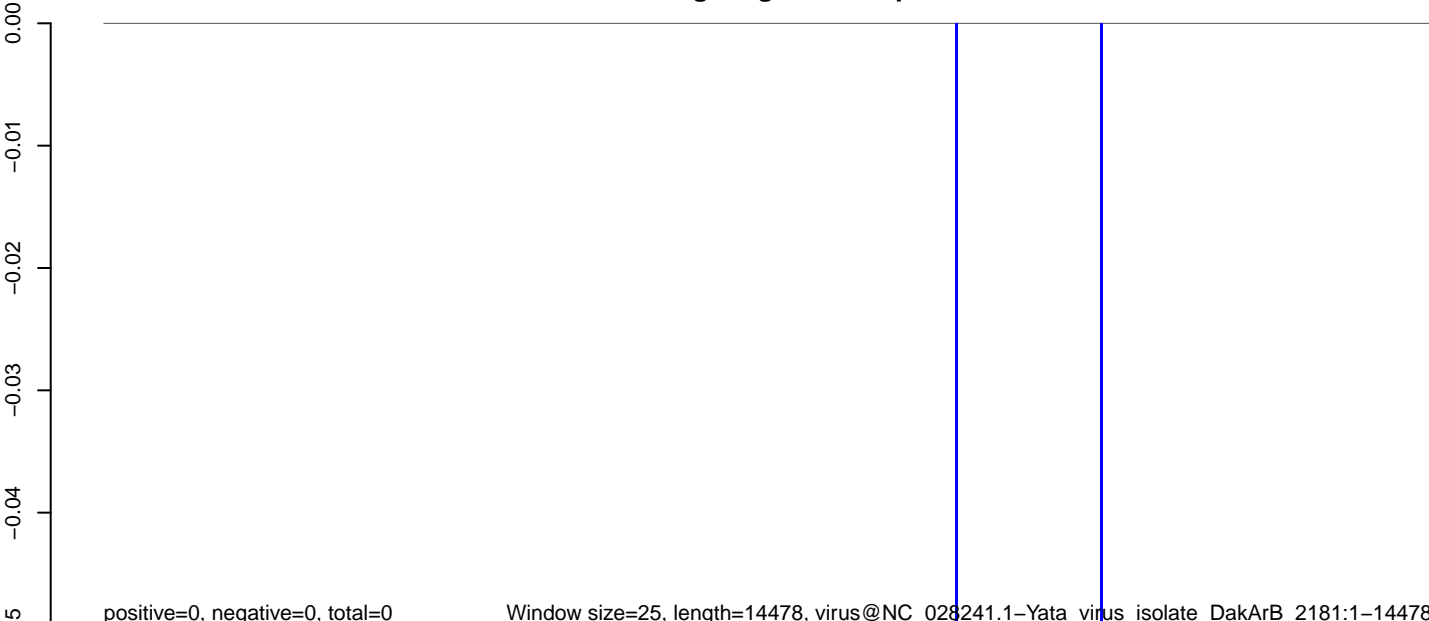
AeAeg_Aag2_cells.18_23.rep



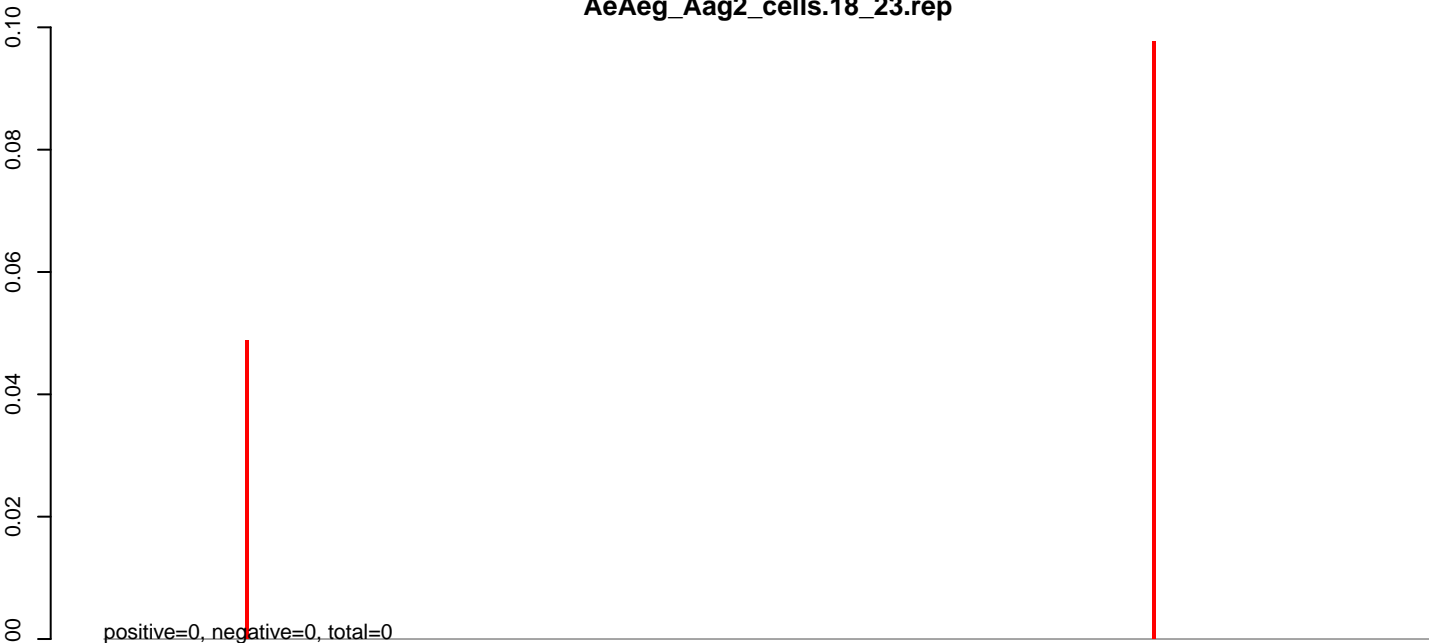
AeAeg_Aag2_cells.24_35.rep



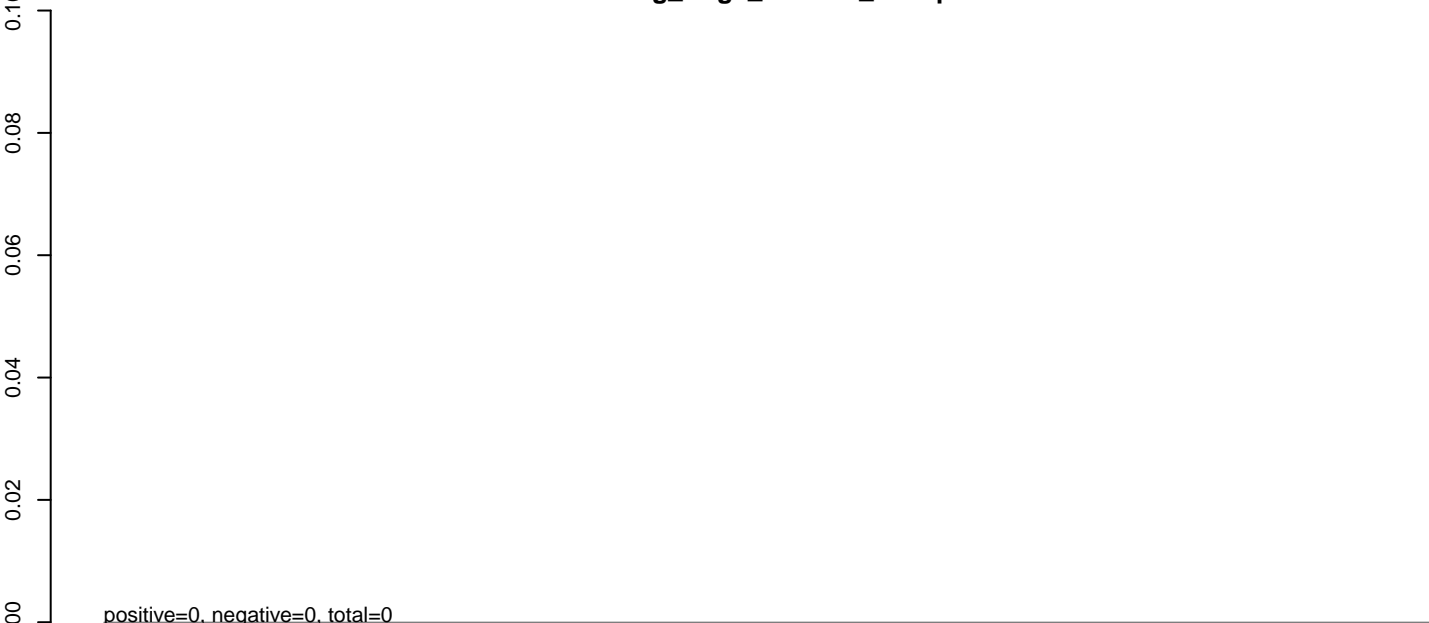
AeAeg_Aag2_cells.rep



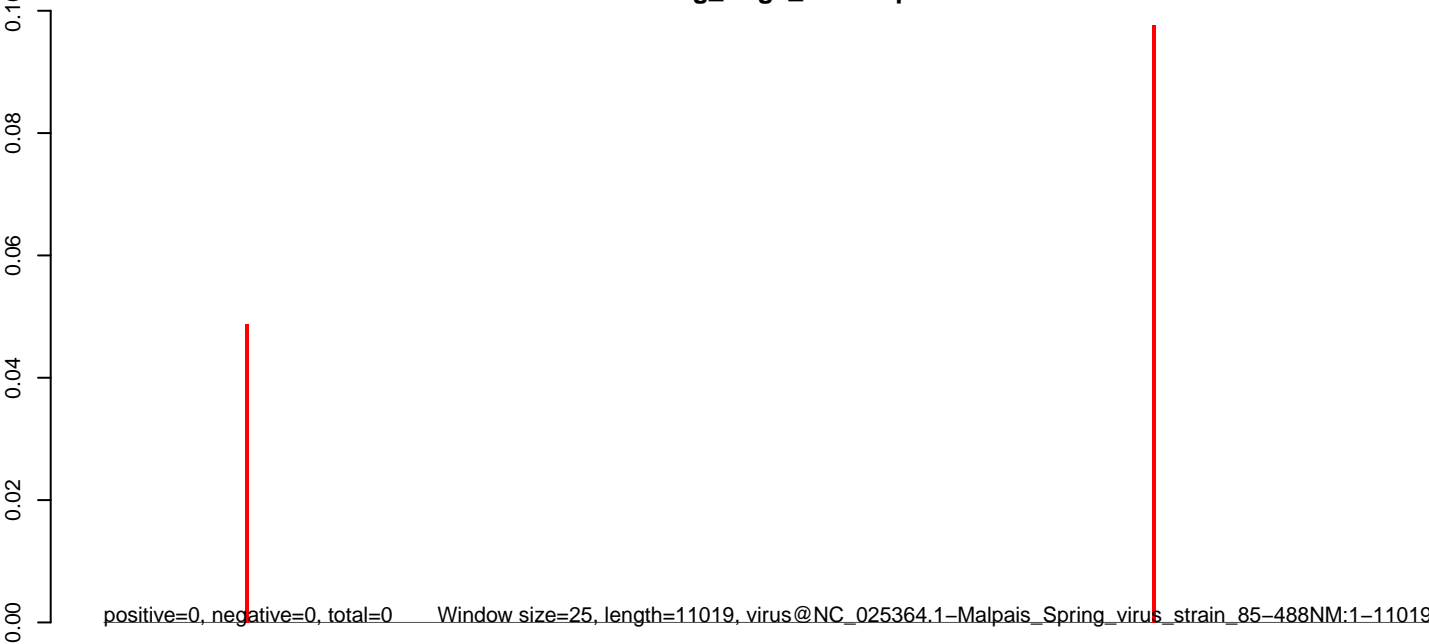
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



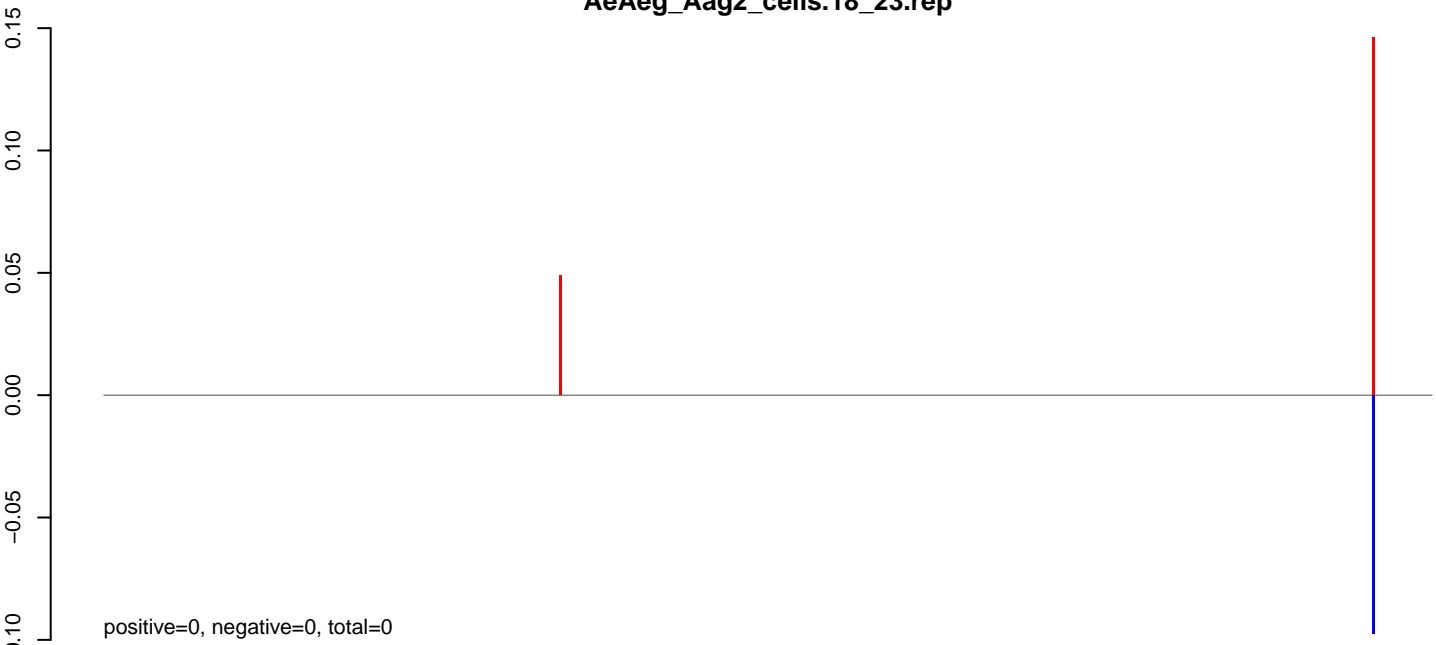
AeAeg_Aag2_cells.rep



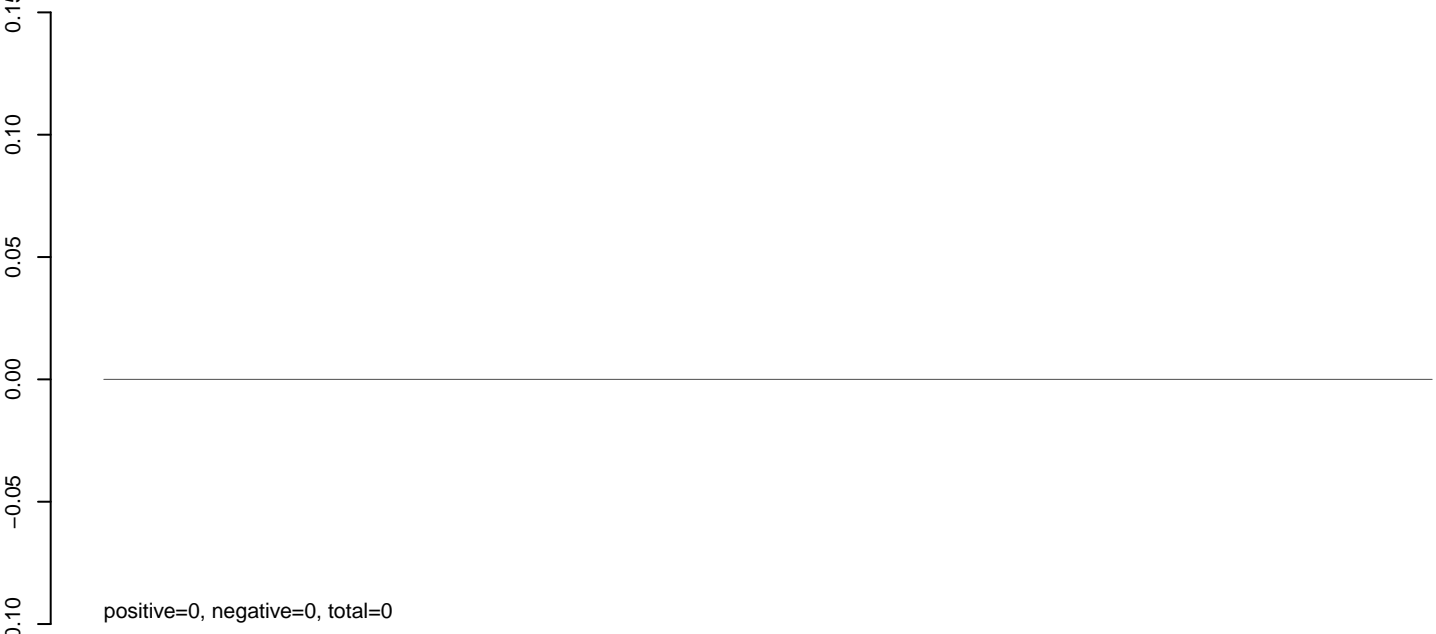
0 2000 4000 6000 8000 10000

Window size=25, length=11019, virus@NC_025364.1-Malpais_Spring_virus_strain_85-488NM:1-11019

AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



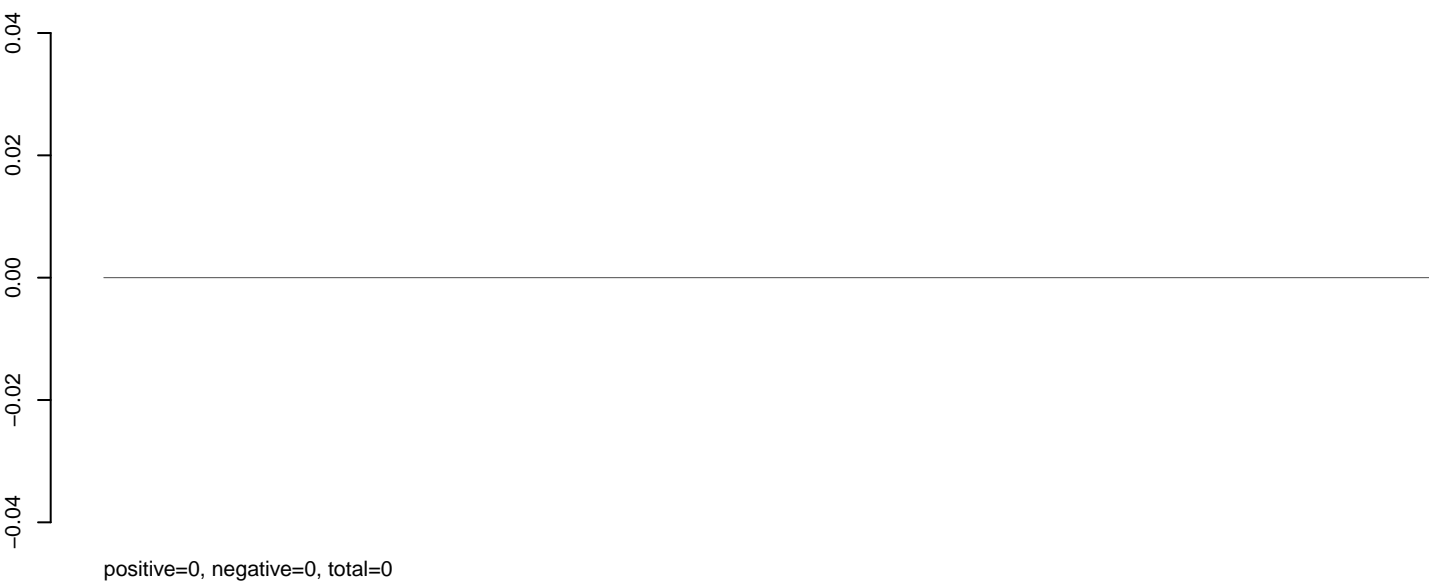
AeAeg_Aag2_cells.rep



AeAeg_Aag2_cells.18_23.rep



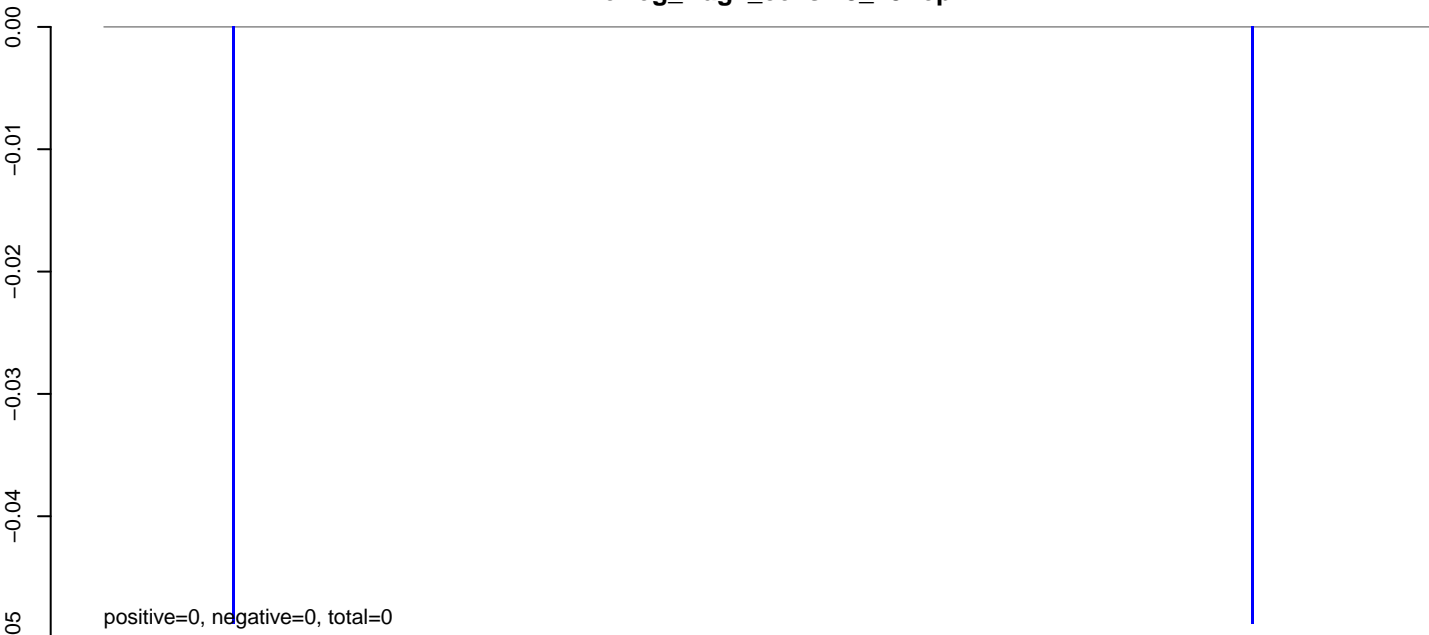
AeAeg_Aag2_cells.24_35.rep



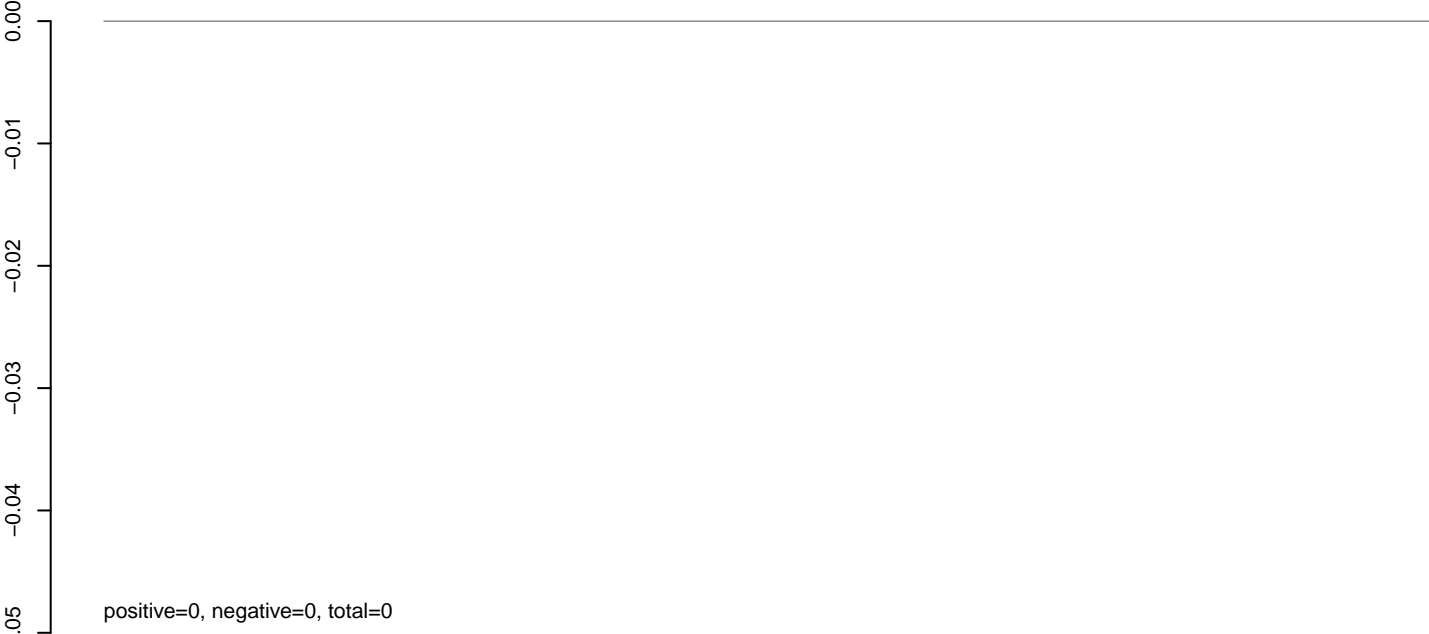
AeAeg_Aag2_cells.rep



AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



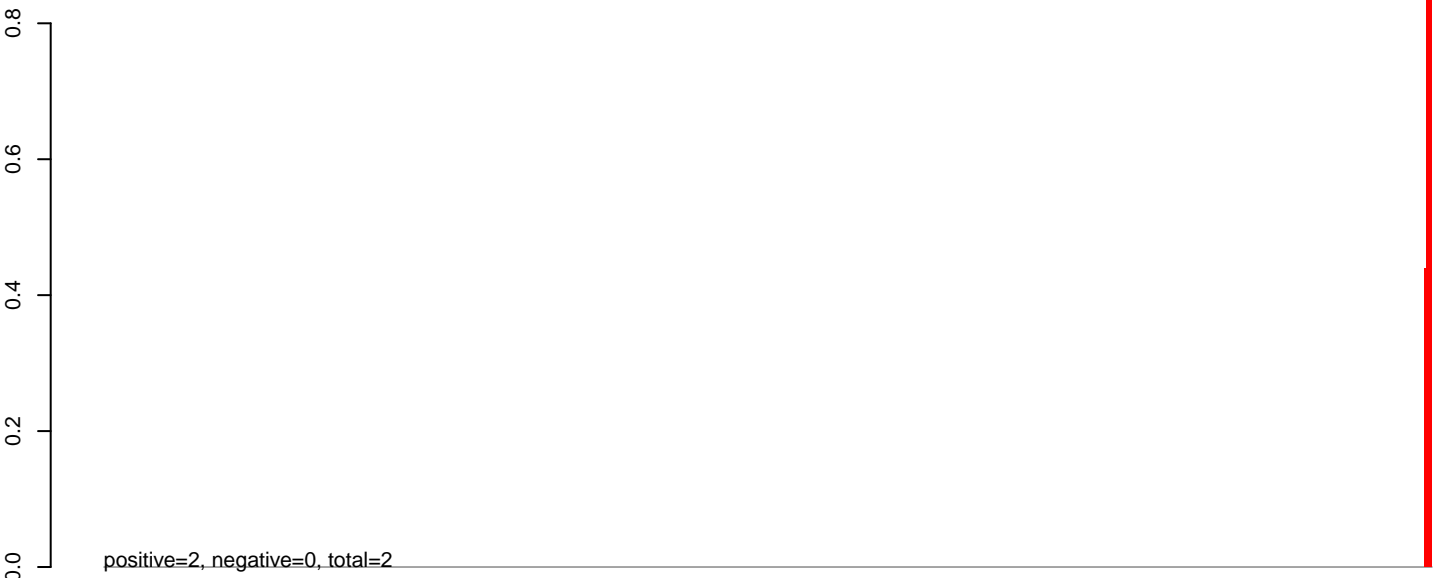
AeAeg_Aag2_cells.rep



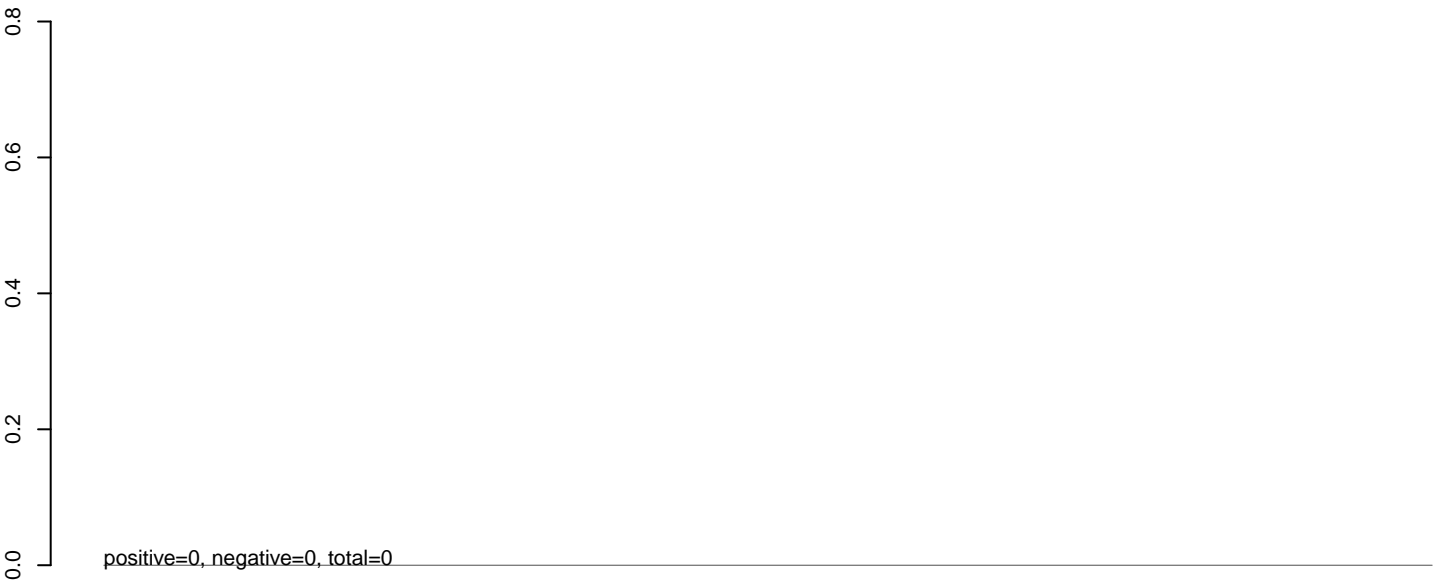
Window size=25, length=14717, virus@NC_017685.1-Obodhiang_virus:1-14717

0 5000 10000 15000

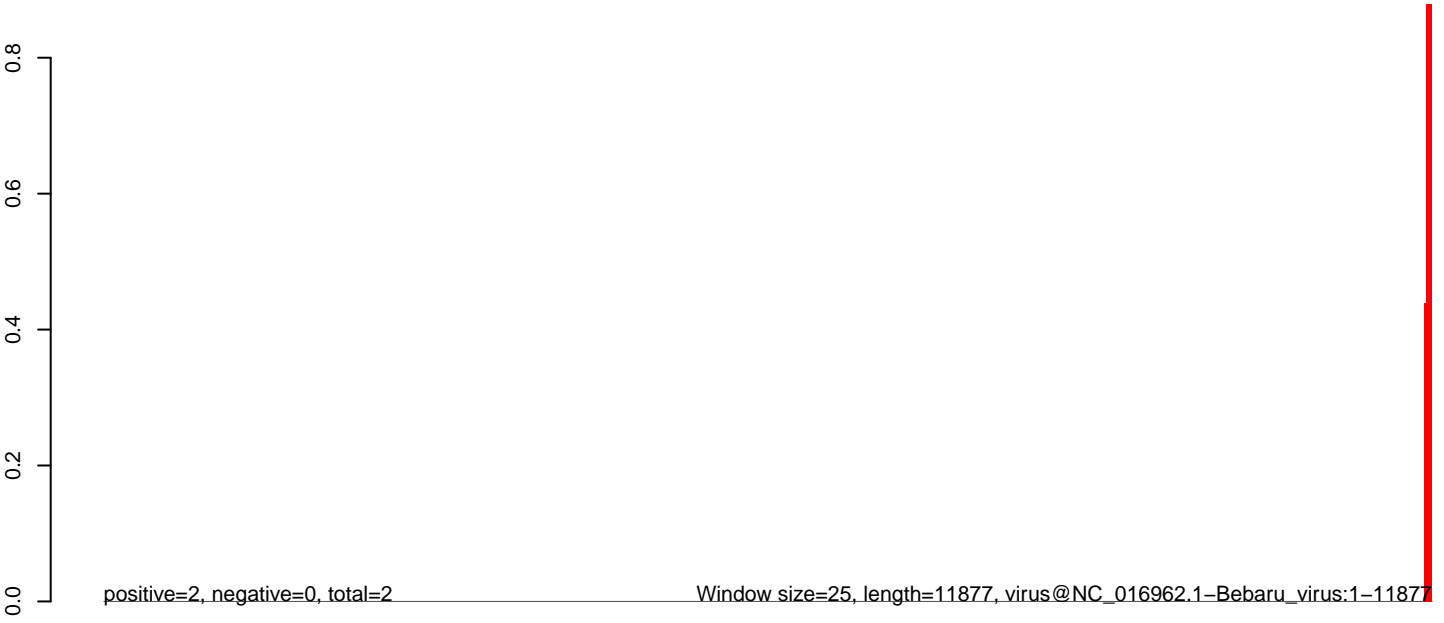
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

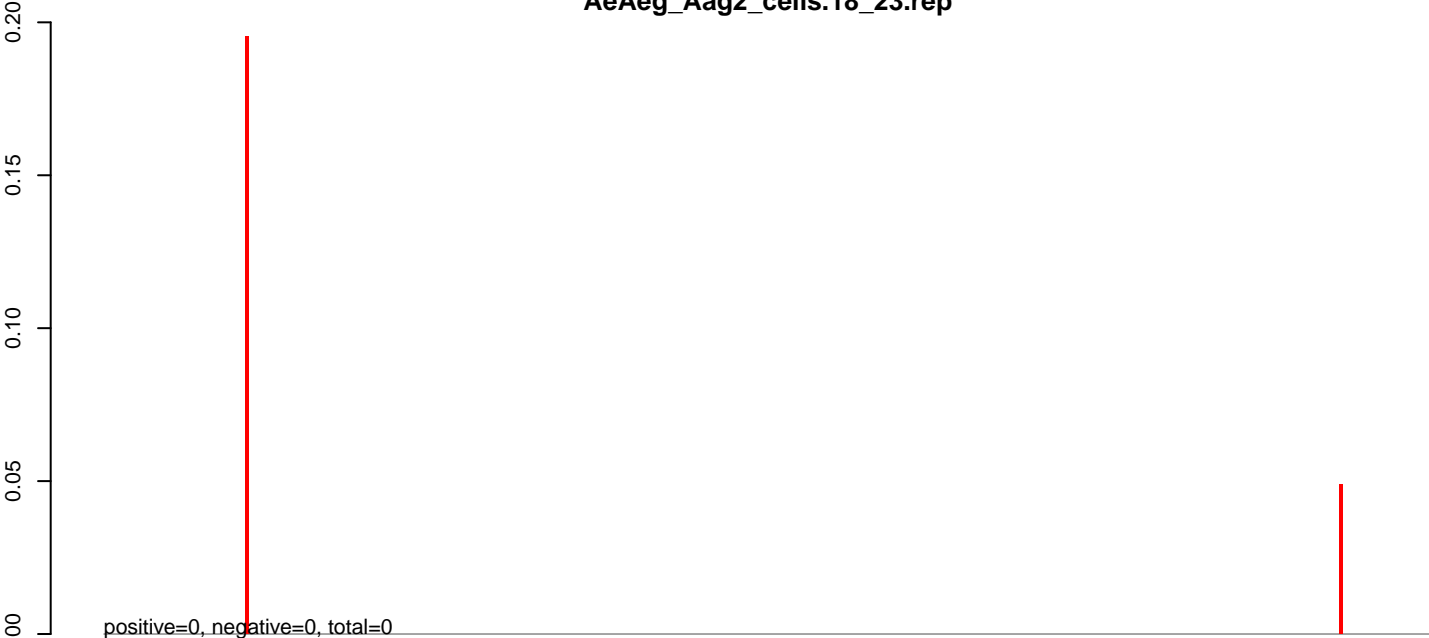


AeAeg_Aag2_cells.rep

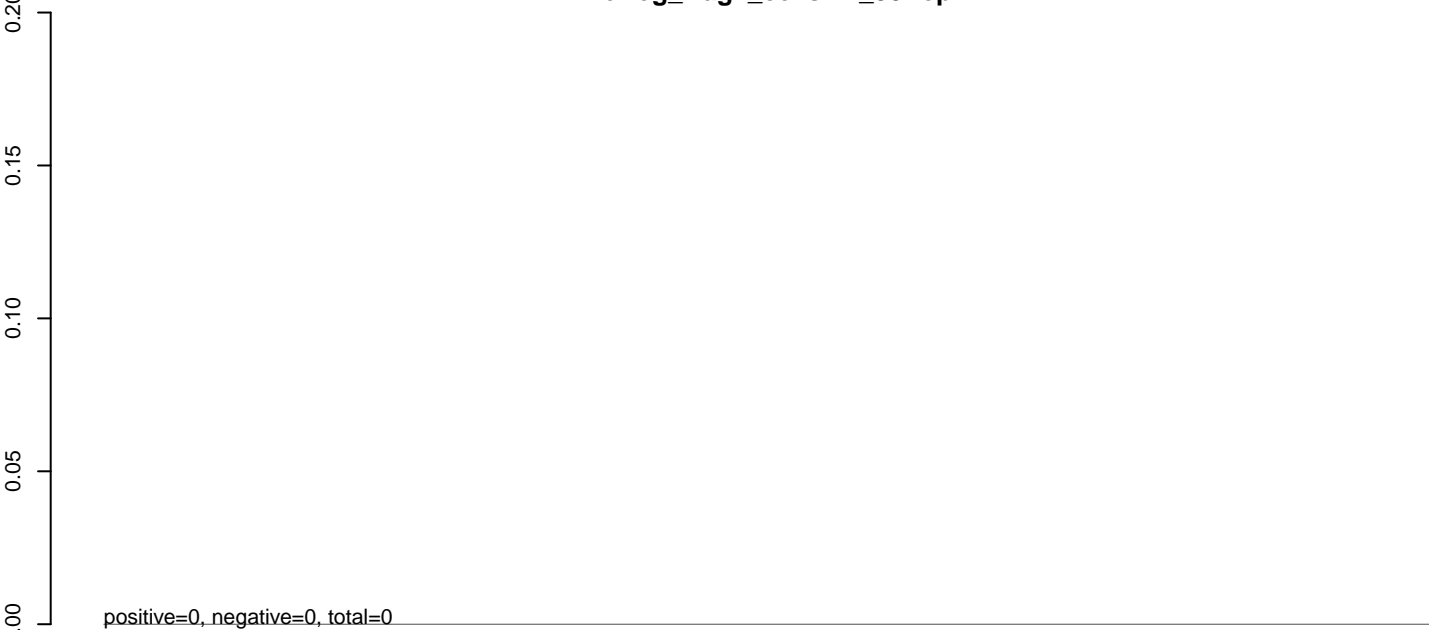


0 2000 4000 6000 8000 10000 12000

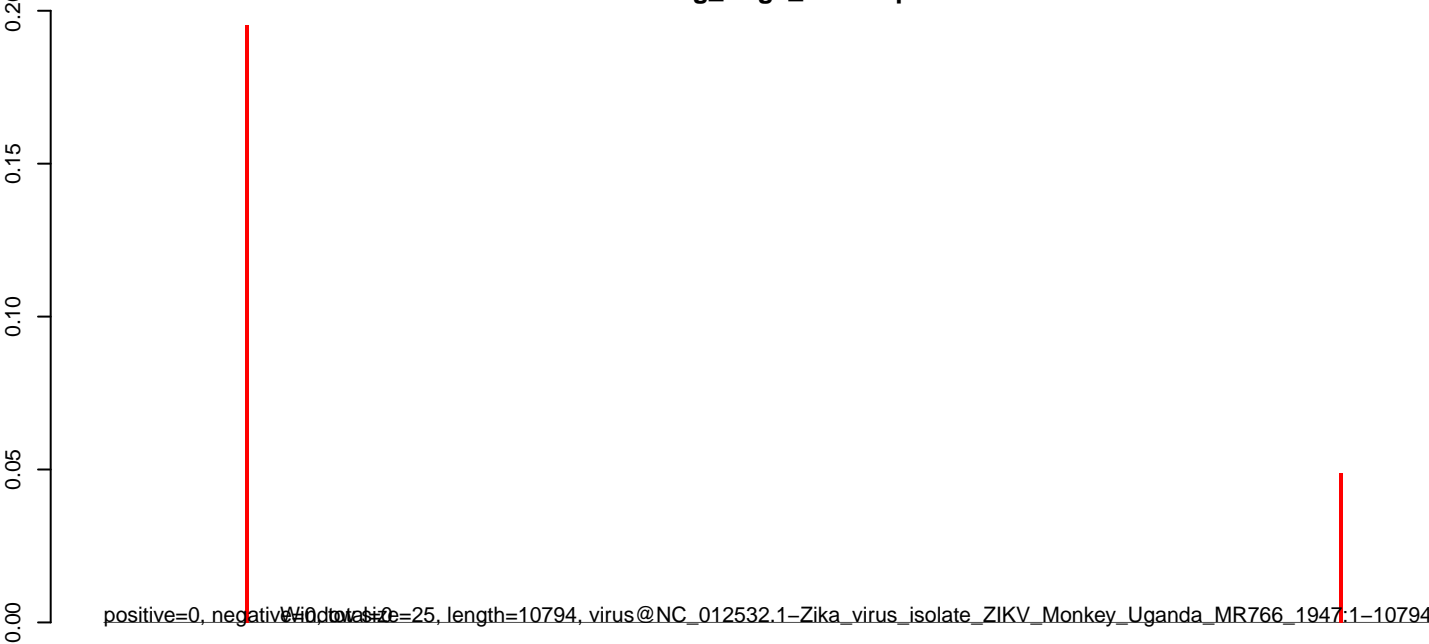
AeAeg_Aag2_cells.18_23.rep



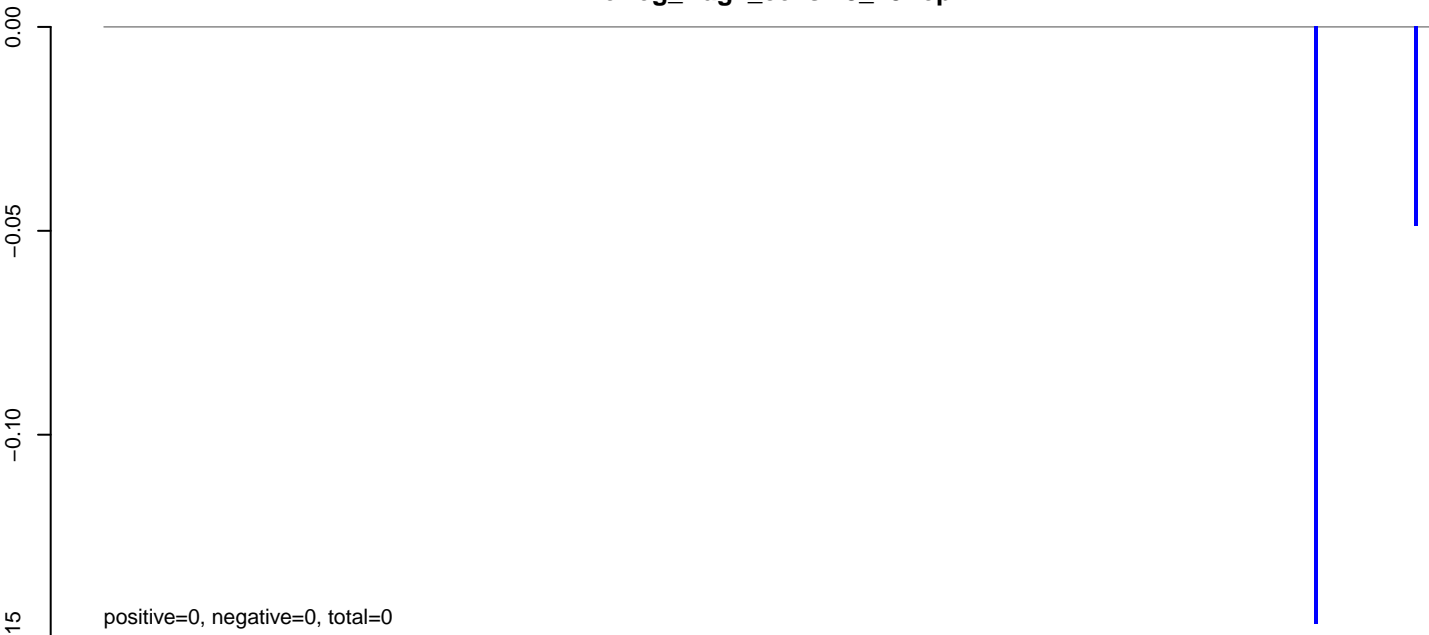
AeAeg_Aag2_cells.24_35.rep



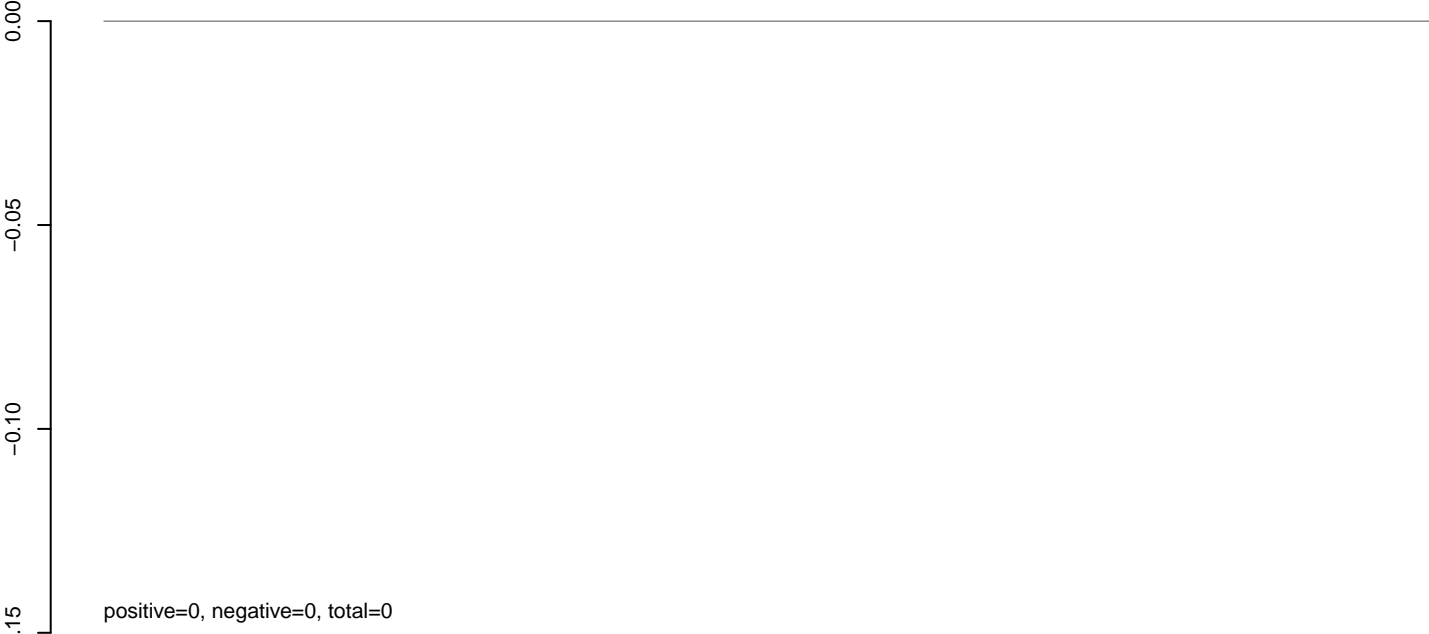
AeAeg_Aag2_cells.rep



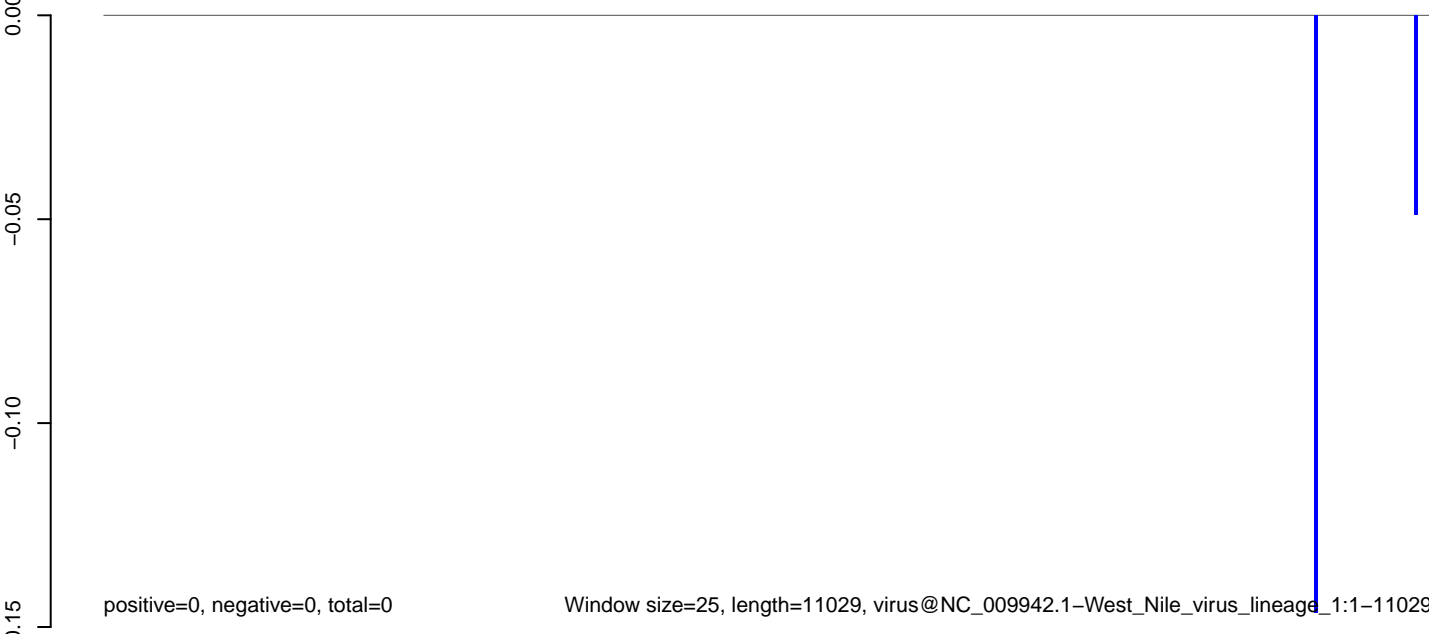
AeAeg_Aag2_cells.18_23.rep



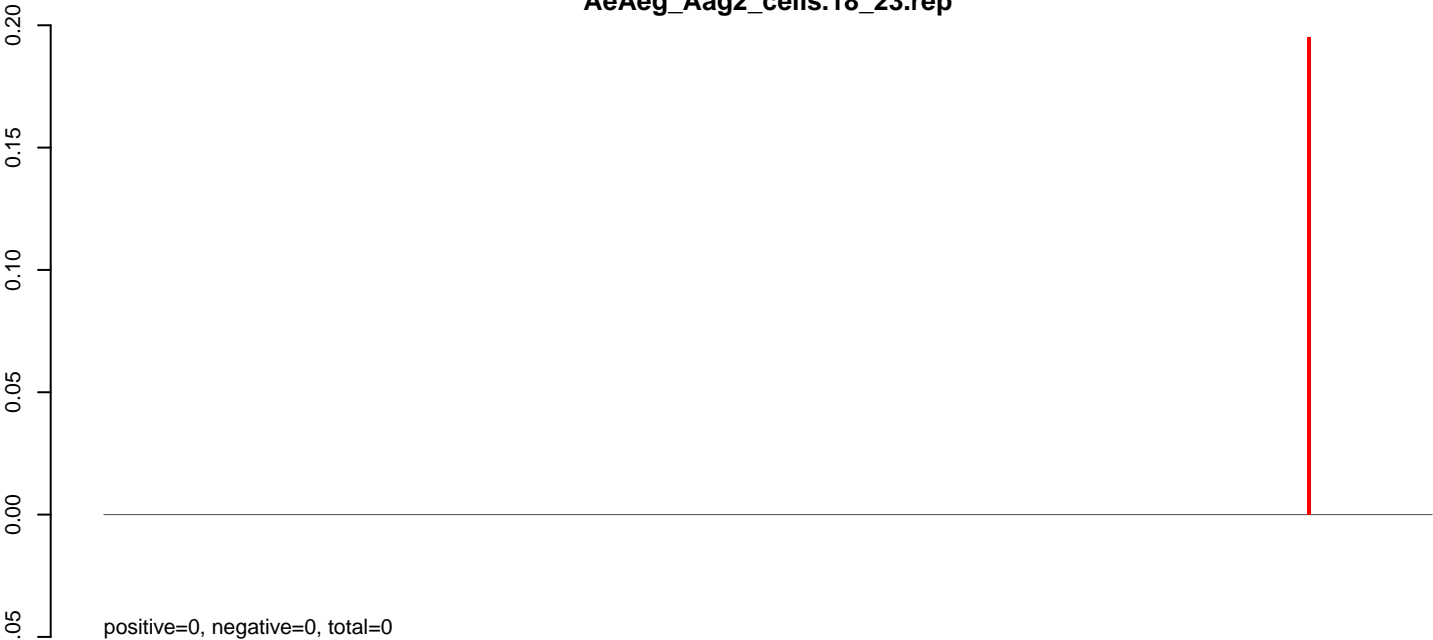
AeAeg_Aag2_cells.24_35.rep



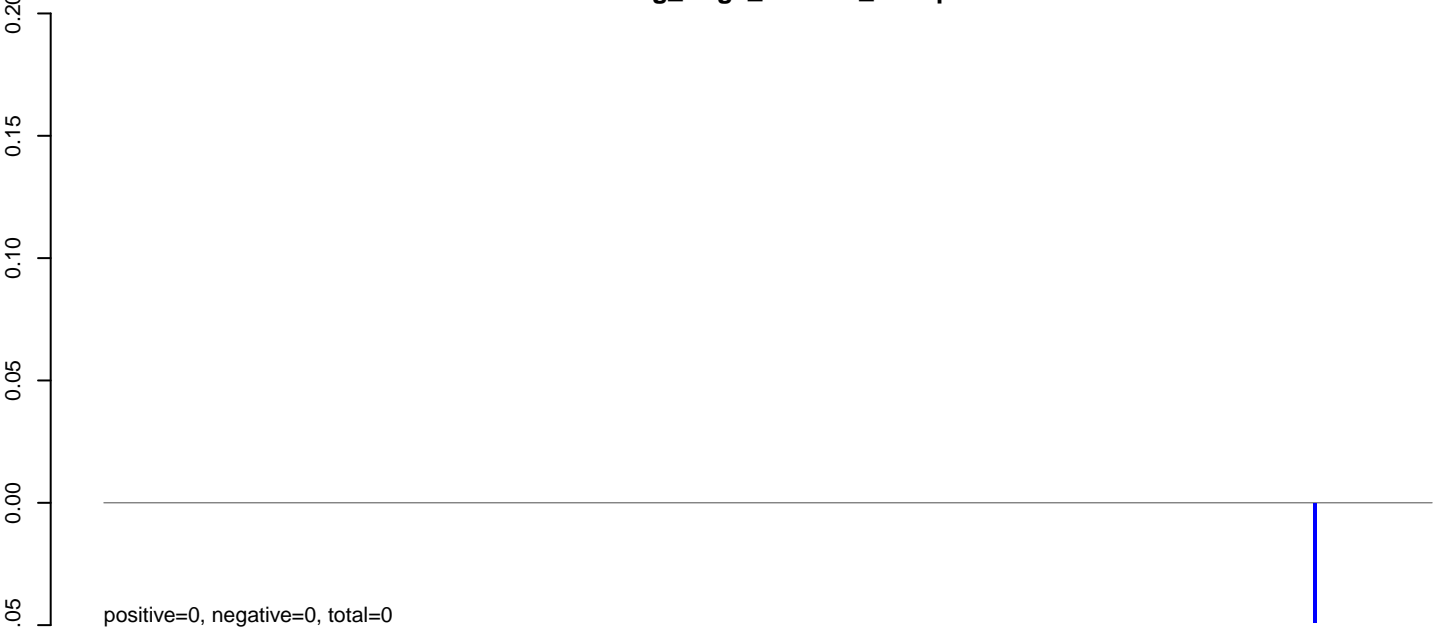
AeAeg_Aag2_cells.rep



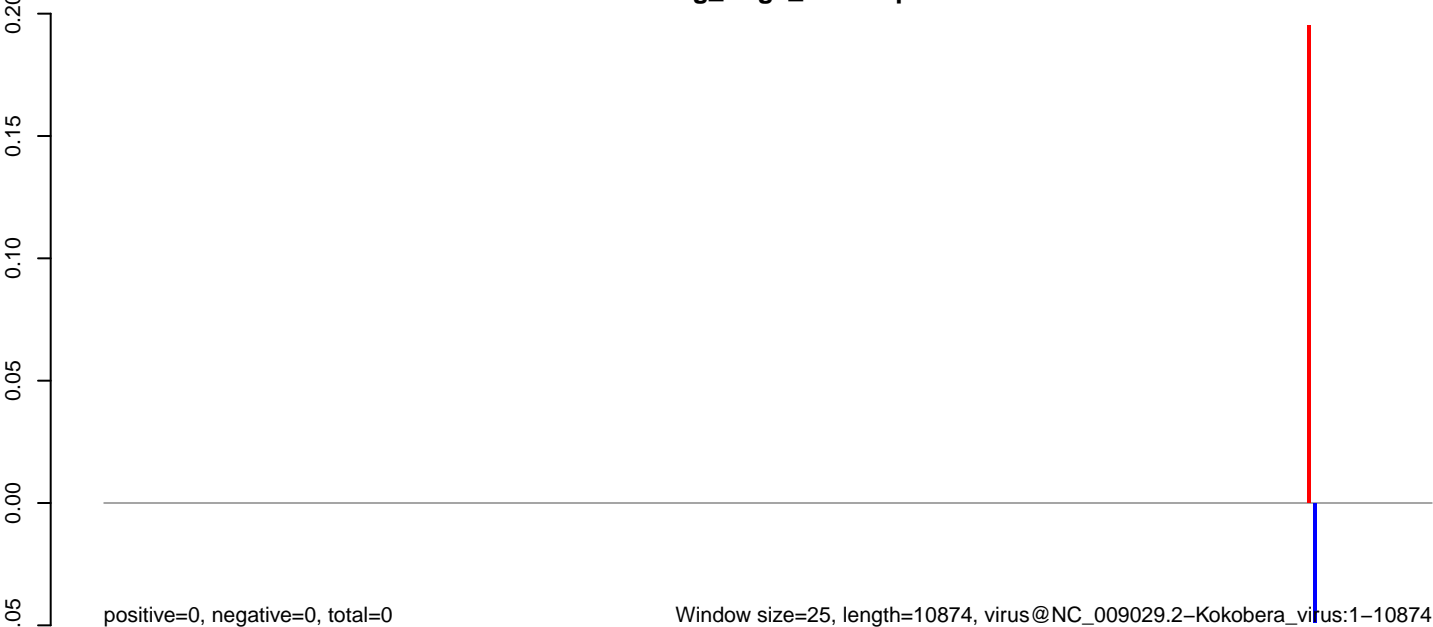
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



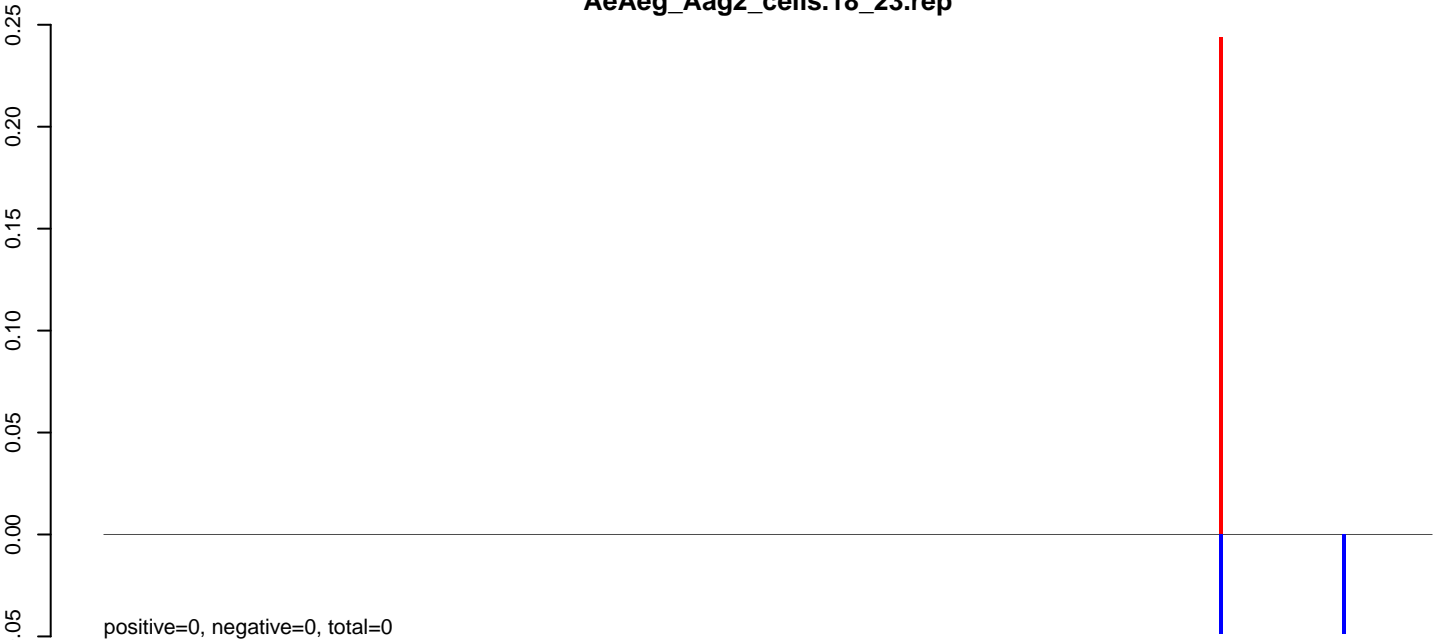
AeAeg_Aag2_cells.rep



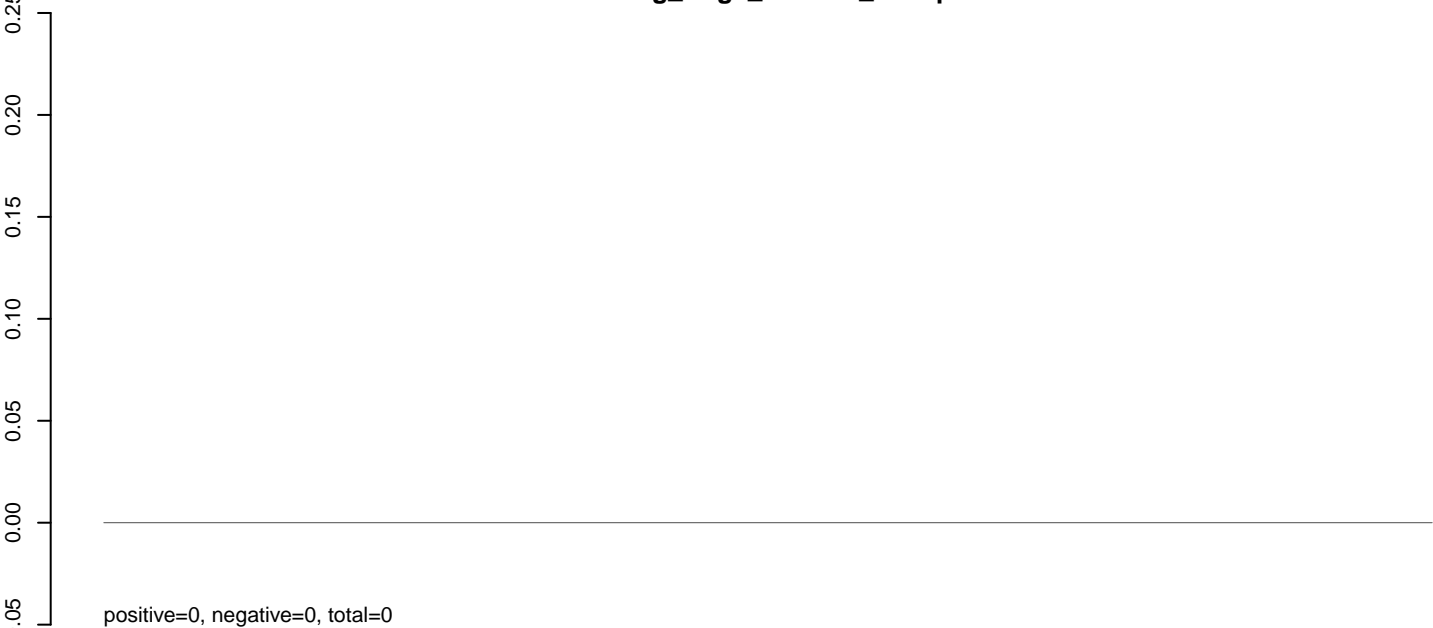
Window size=25, length=10874, virus@NC_009029.2-Kokobera_virus:1-10874

0 2000 4000 6000 8000 10000

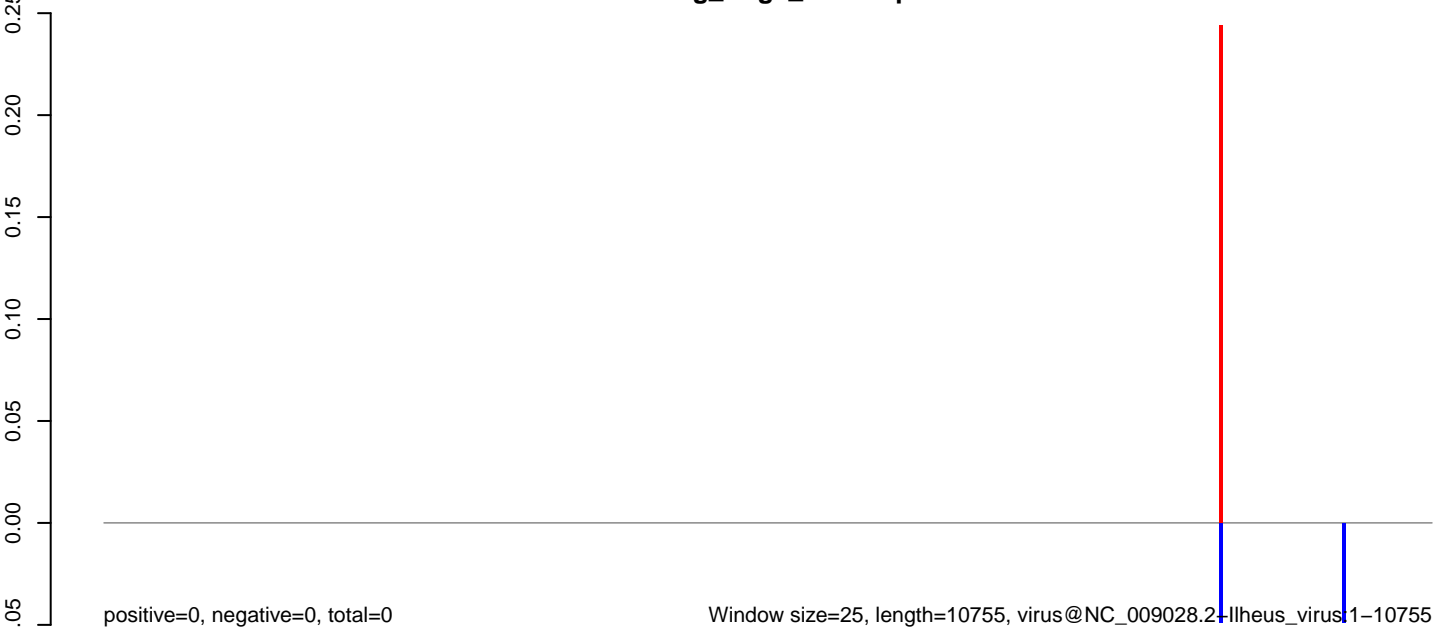
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



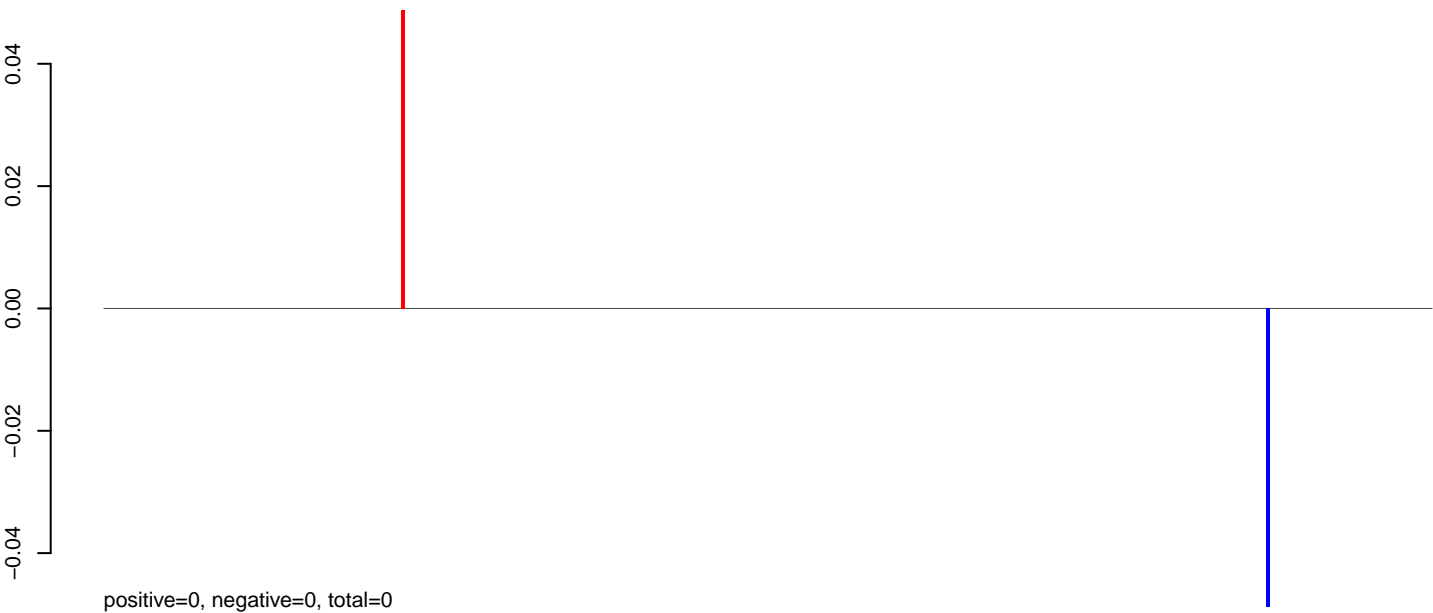
AeAeg_Aag2_cells.rep



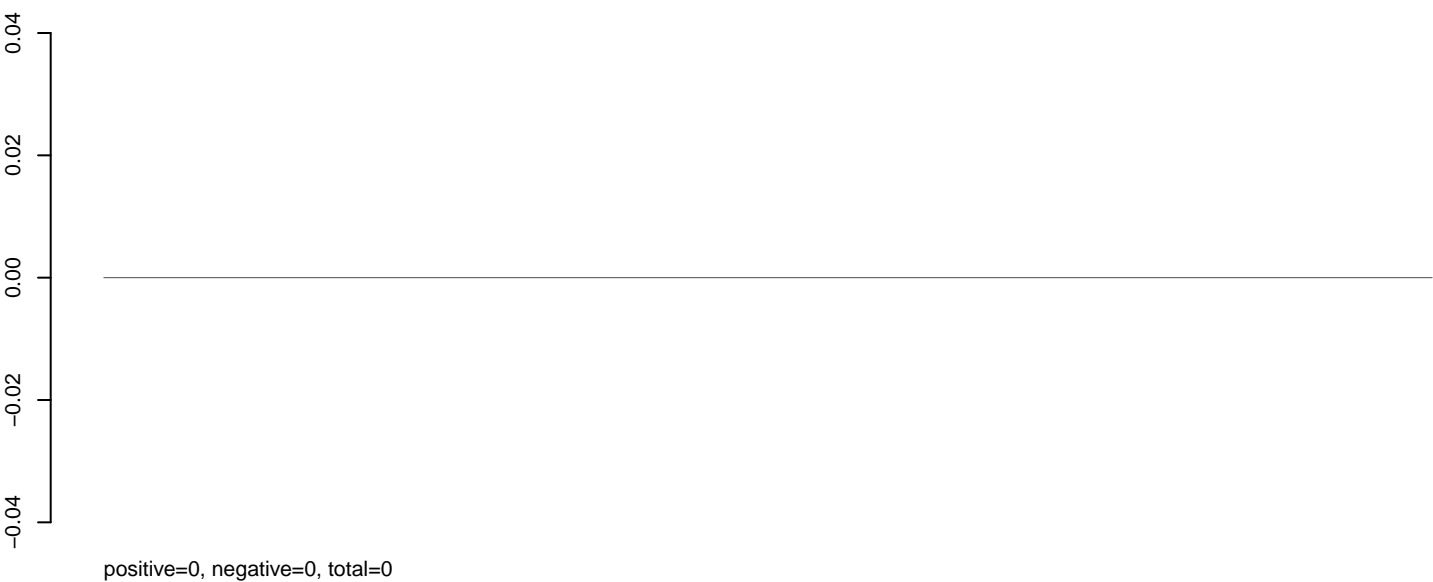
Window size=25, length=10755, virus@NC_009028.2-Illheus_virus1-10755

0 2000 4000 6000 8000 10000

AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

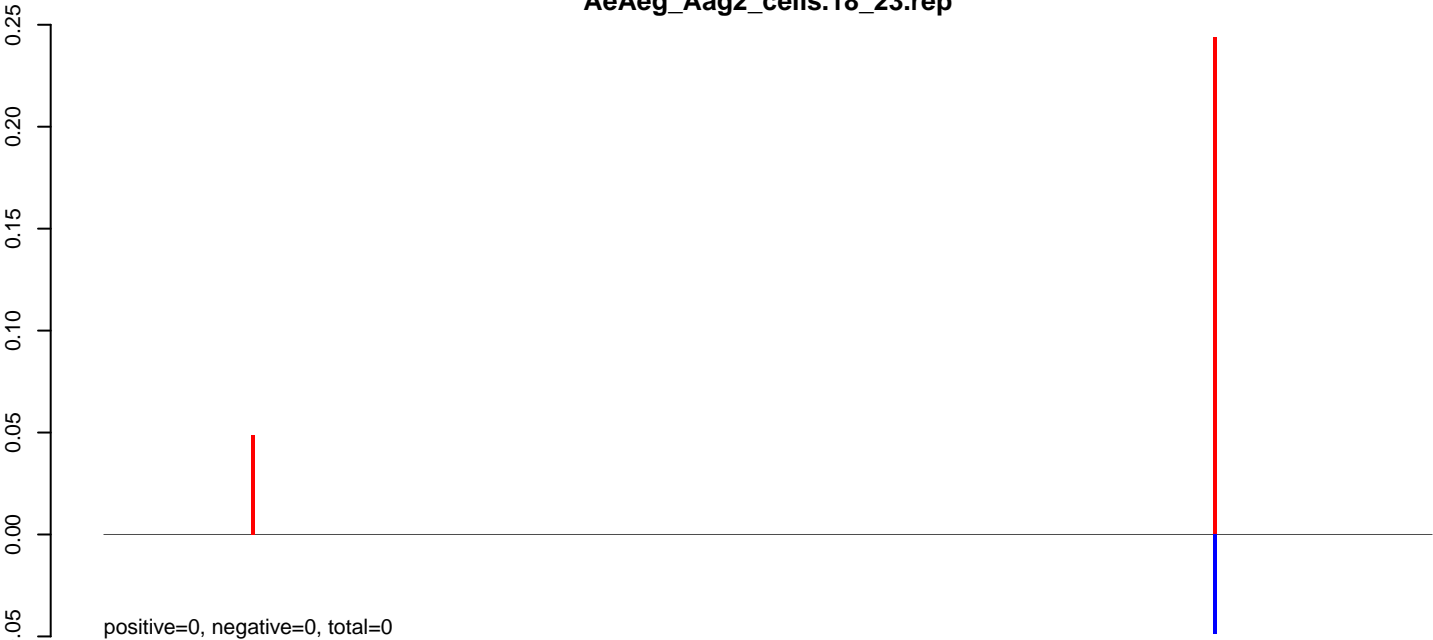


AeAeg_Aag2_cells.rep



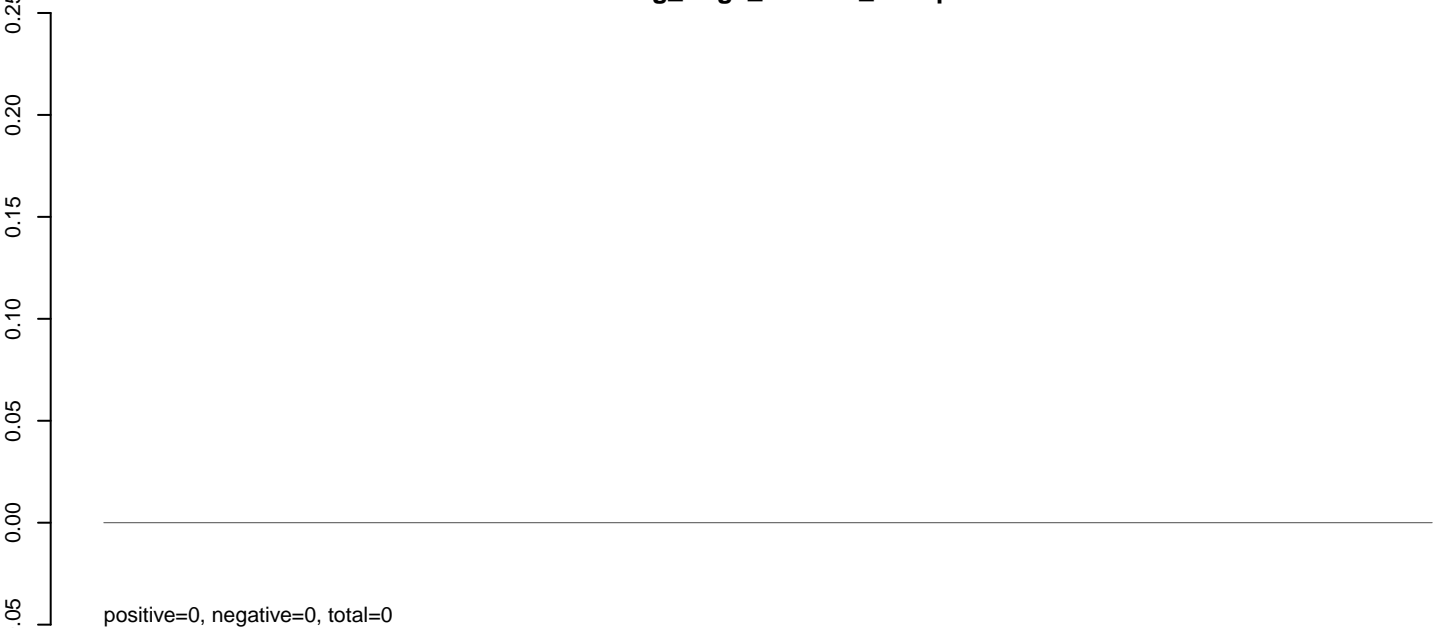
0 2000 4000 6000 8000 10000

AeAeg_Aag2_cells.18_23.rep



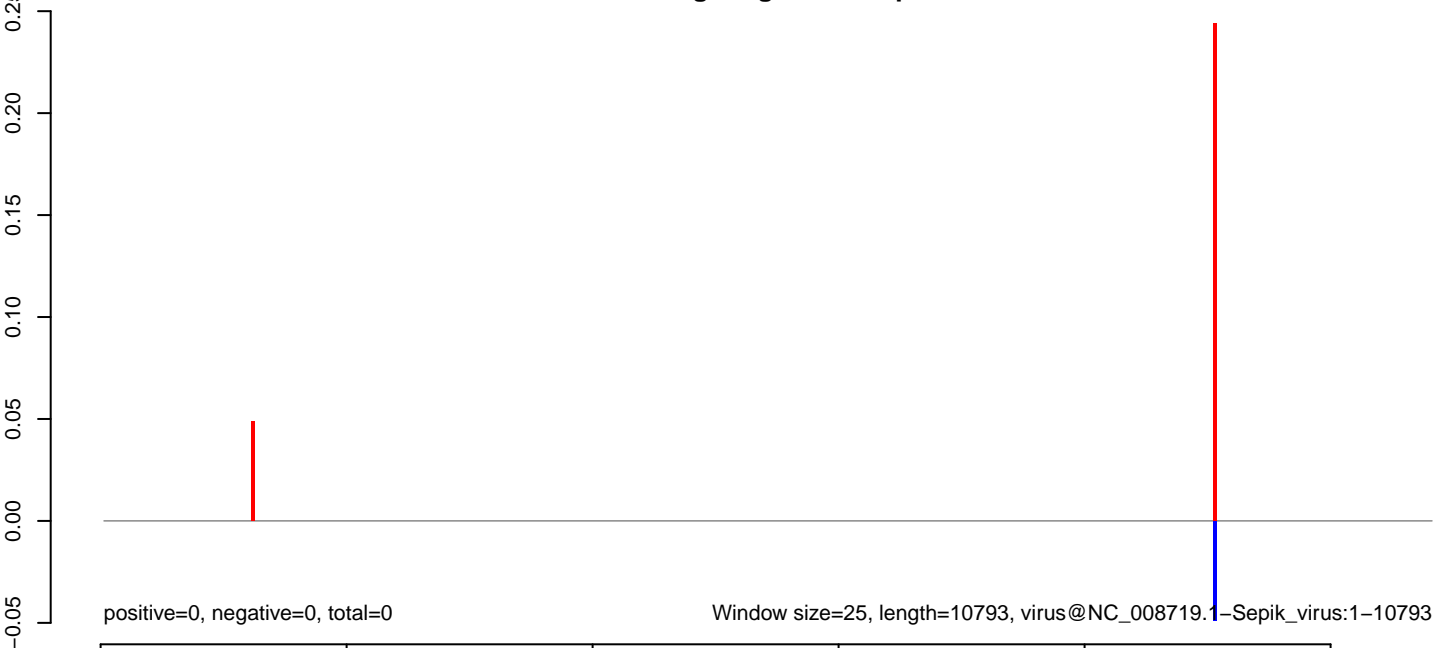
positive=0, negative=0, total=0

AeAeg_Aag2_cells.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_cells.rep

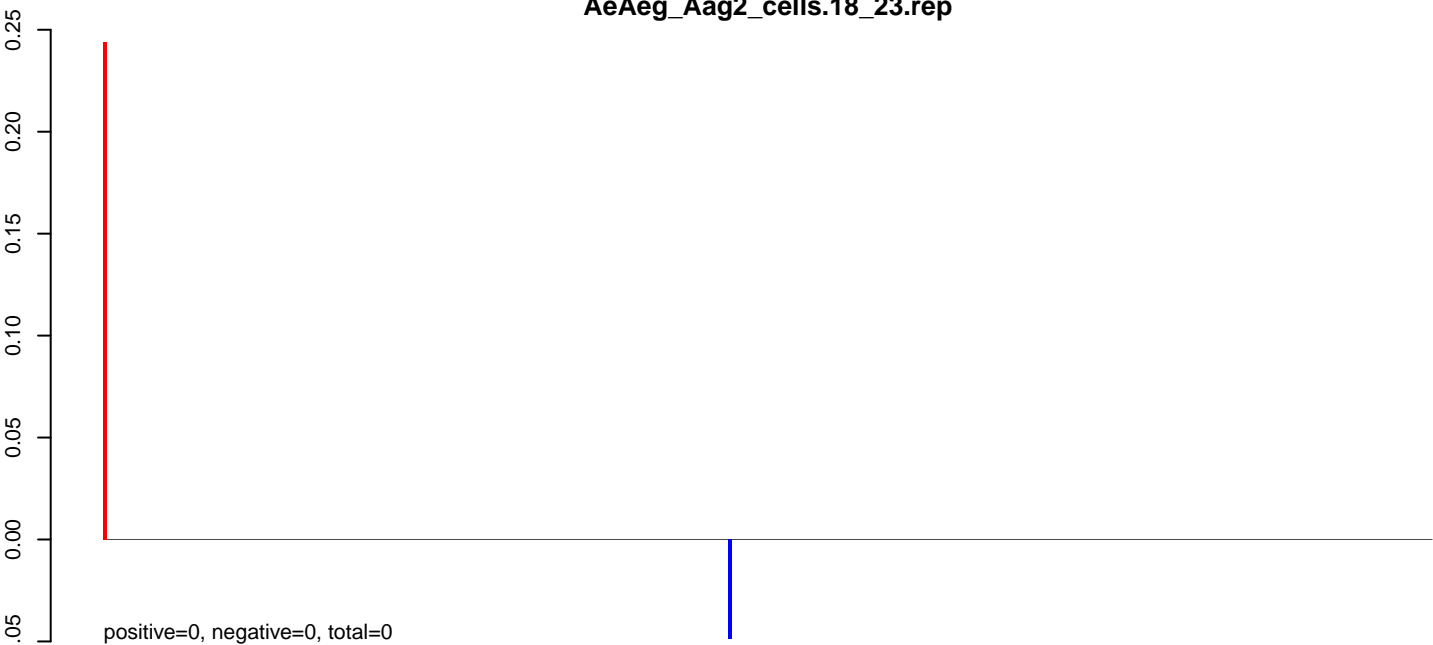


positive=0, negative=0, total=0

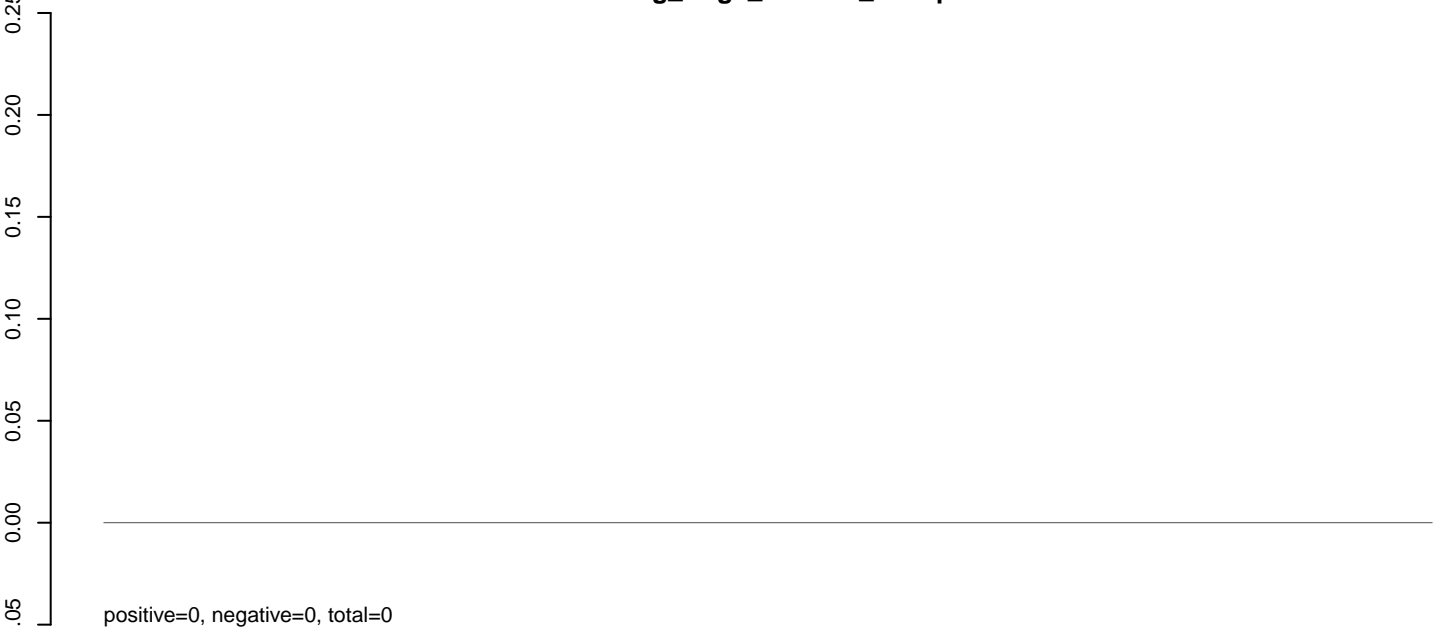
Window size=25, length=10793, virus@NC_008719.1-Sepik_virus:1-10793

0 2000 4000 6000 8000 10000

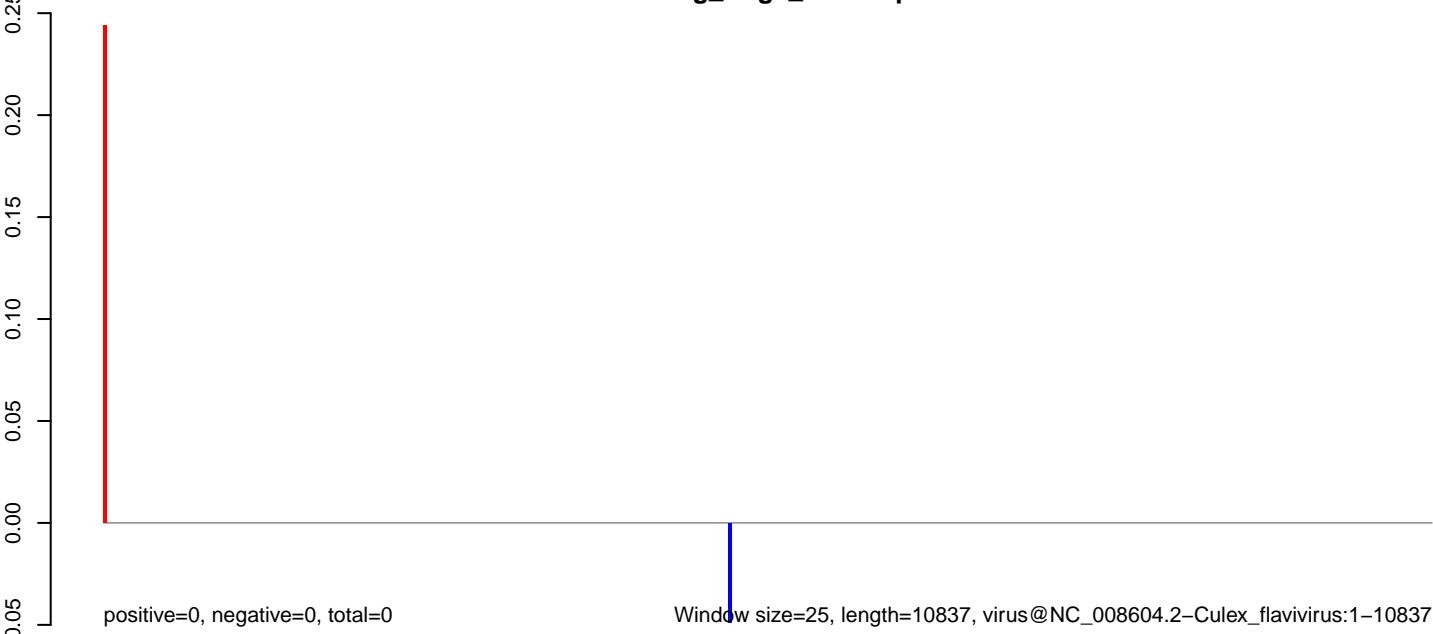
AeAeg_Aag2_cells.18_23.rep



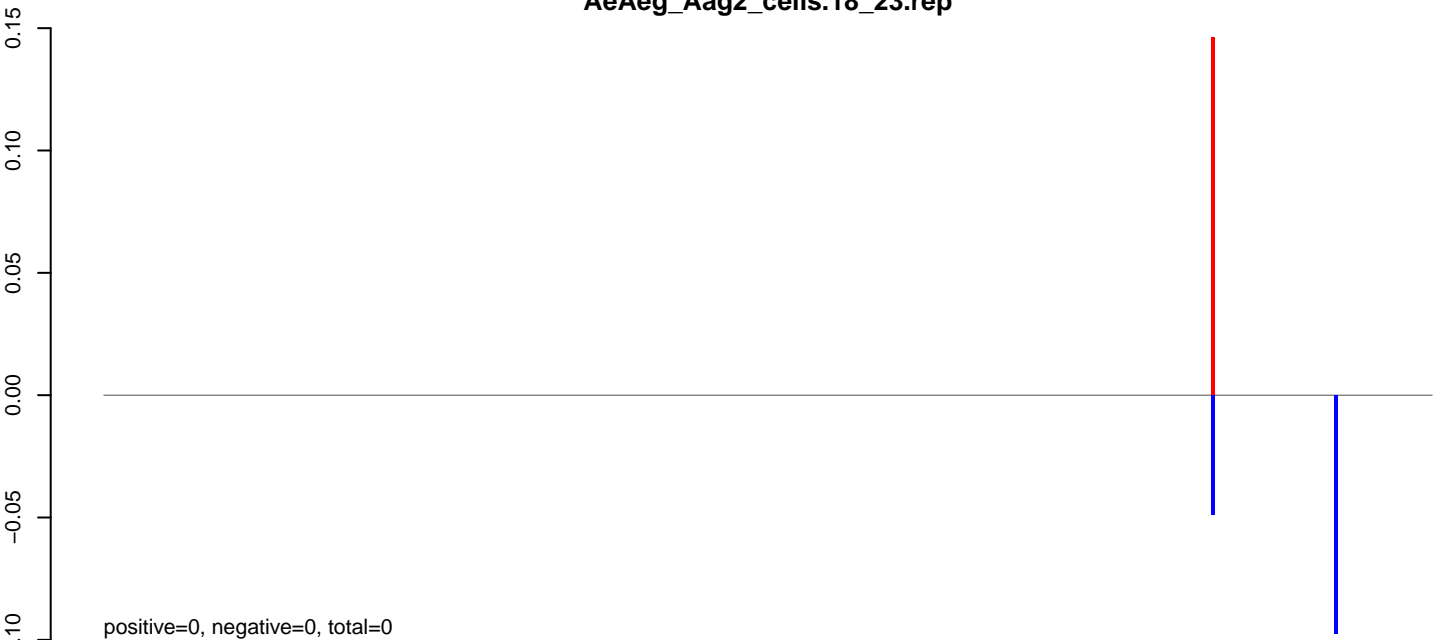
AeAeg_Aag2_cells.24_35.rep



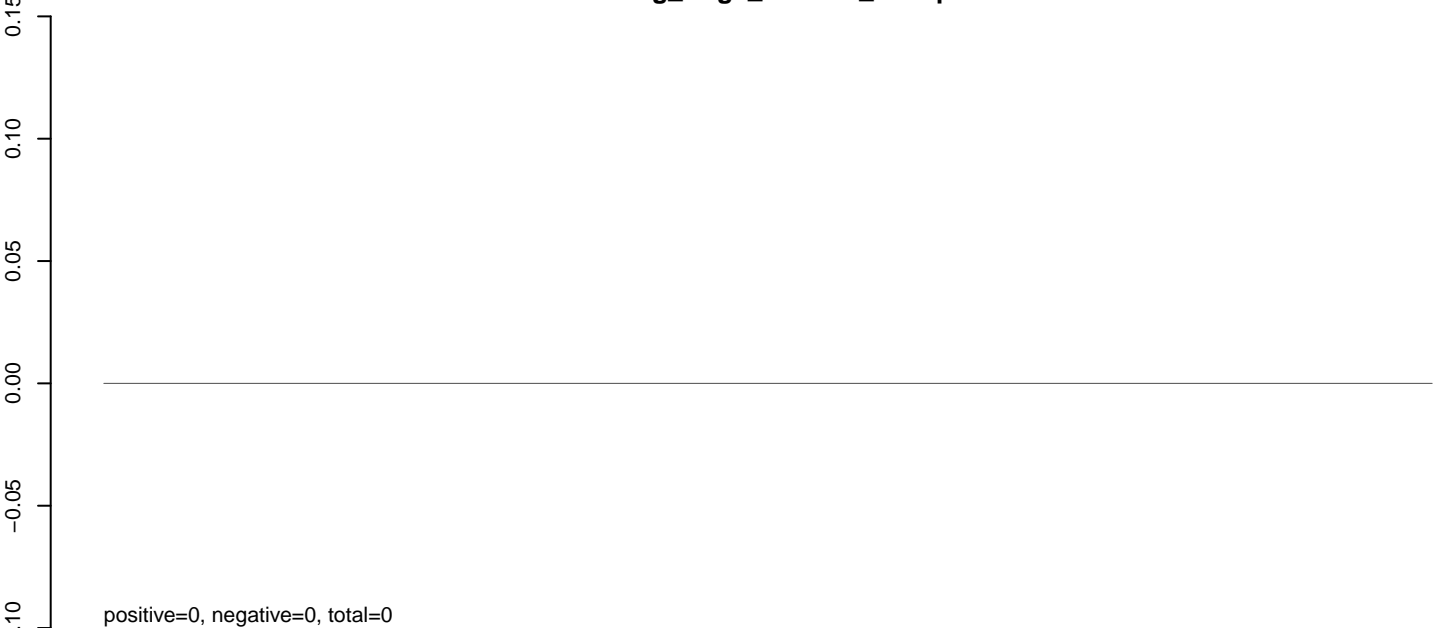
AeAeg_Aag2_cells.rep



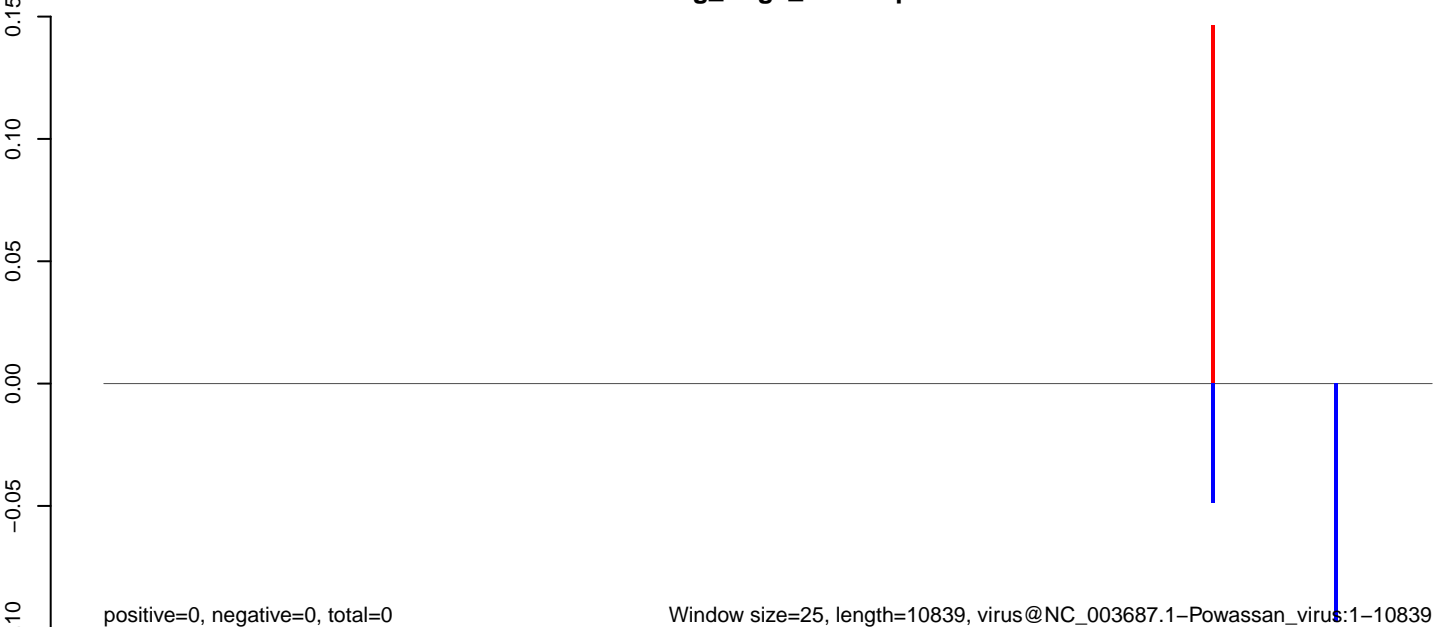
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

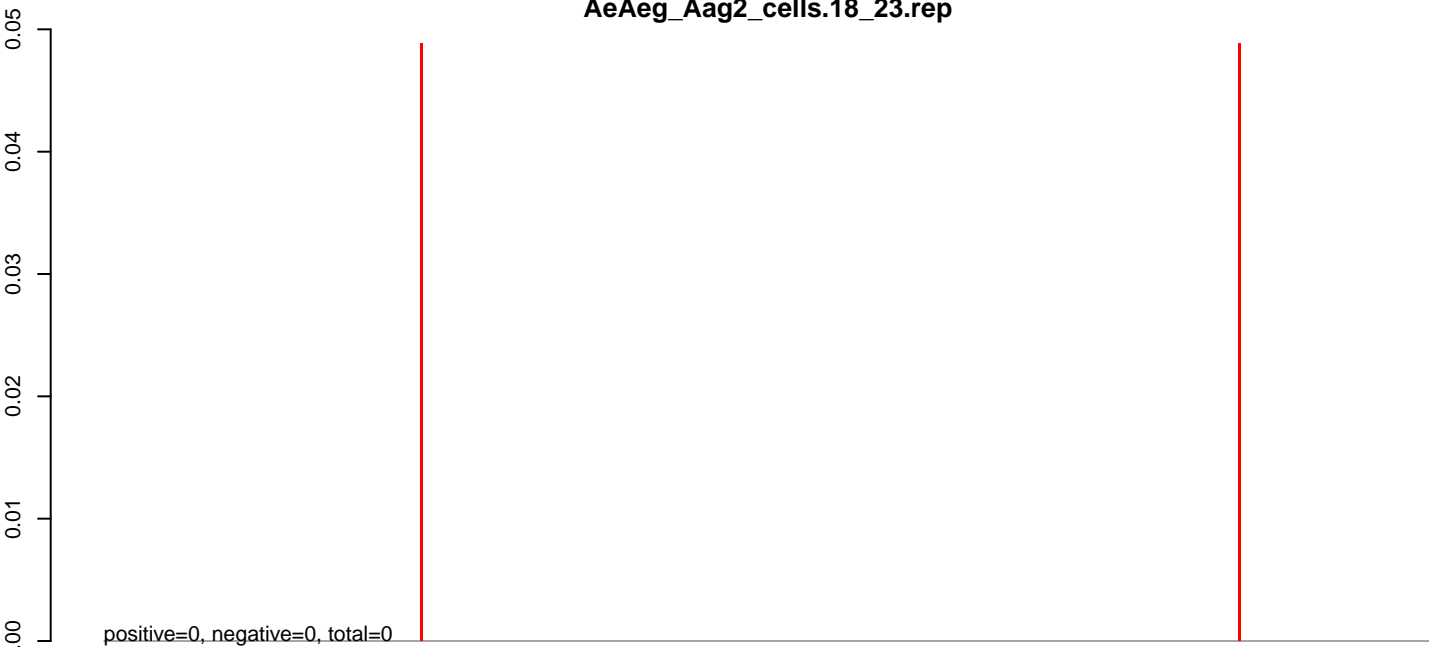


AeAeg_Aag2_cells.rep

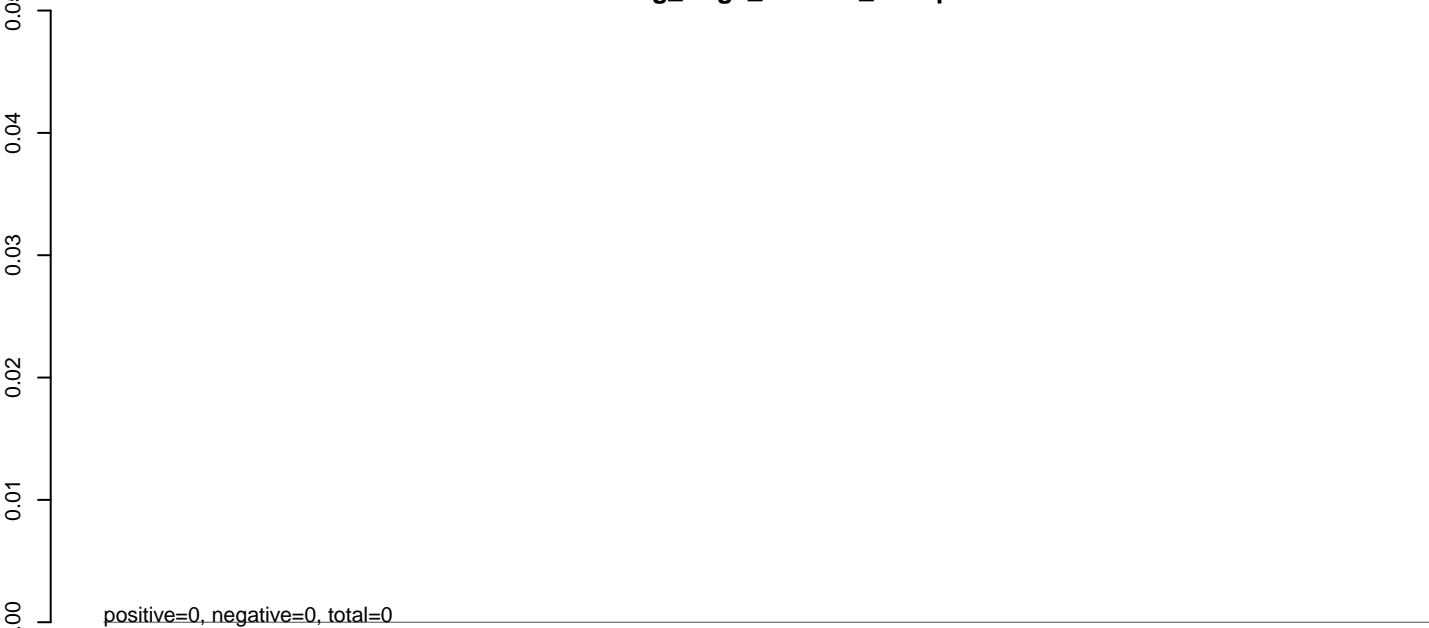


0 2000 4000 6000 8000 10000

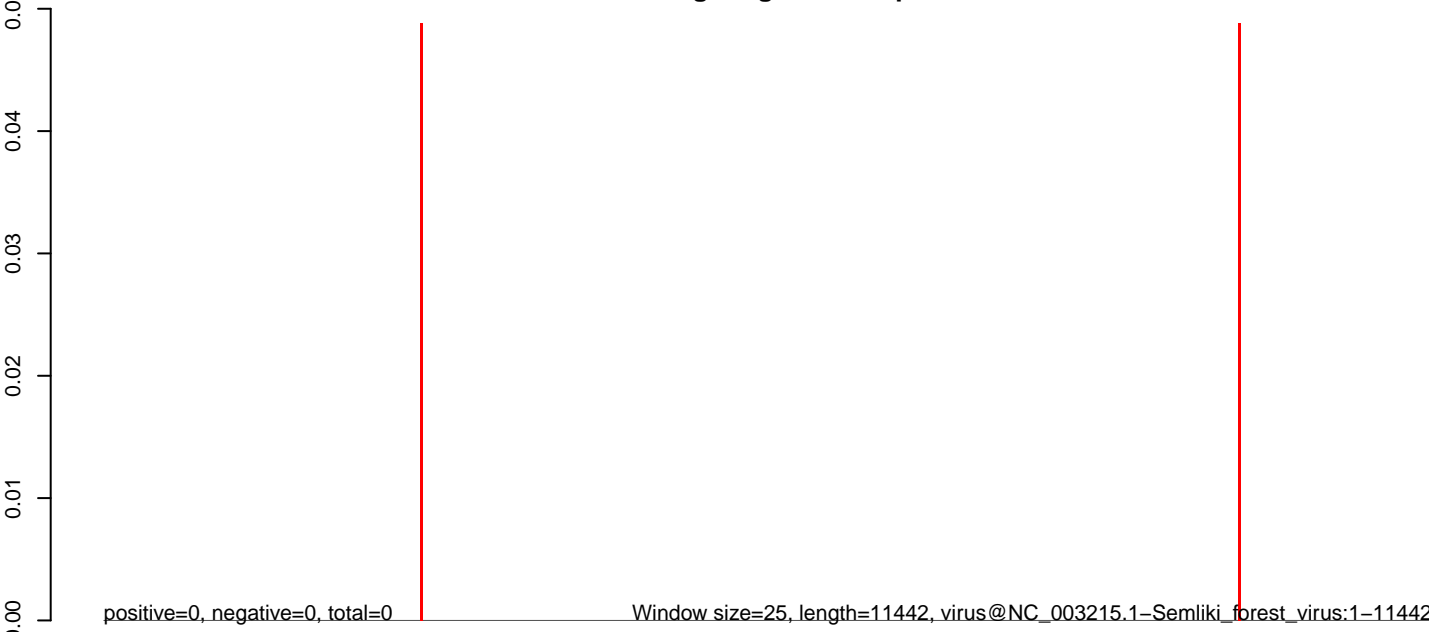
AeAeg_Aag2_cells.18_23.rep



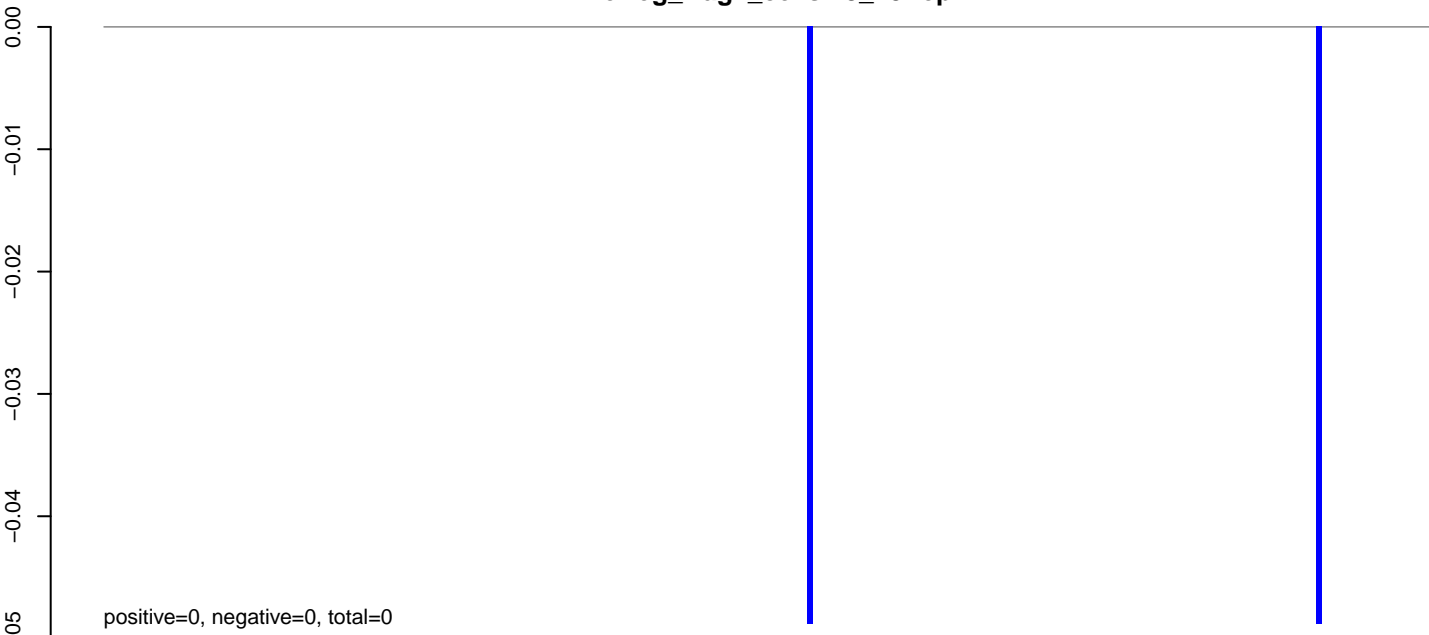
AeAeg_Aag2_cells.24_35.rep



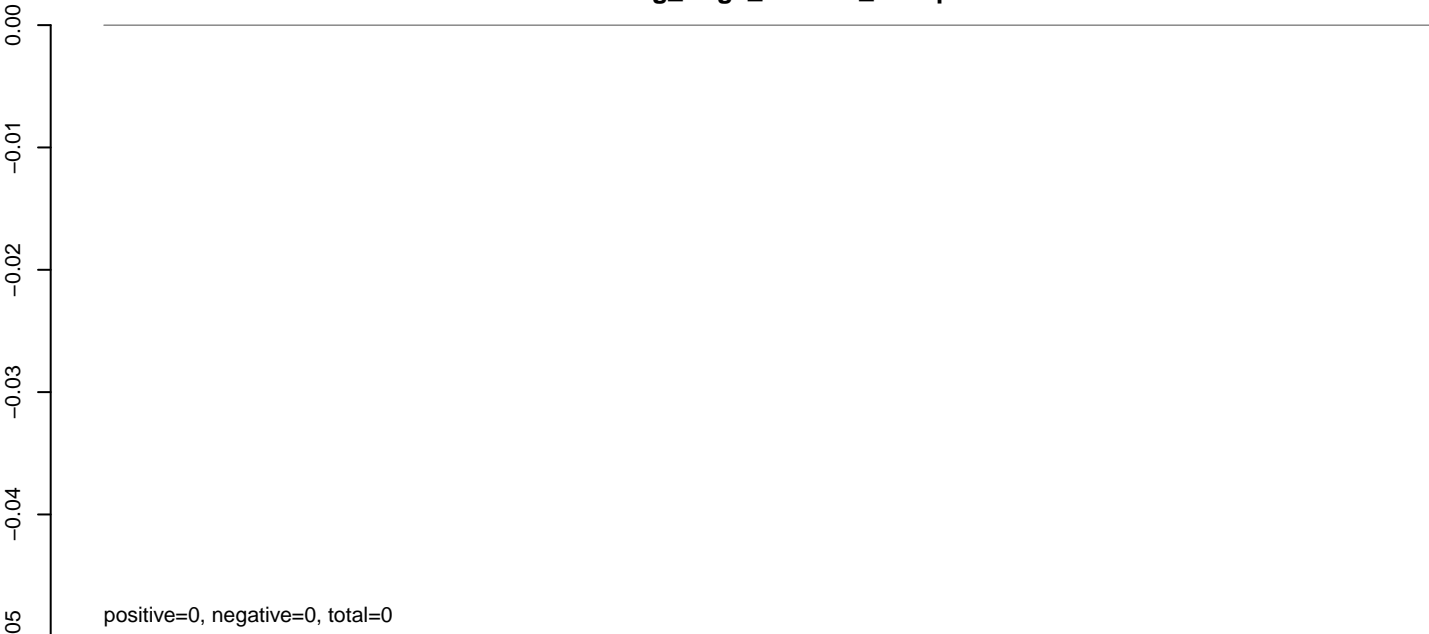
AeAeg_Aag2_cells.rep



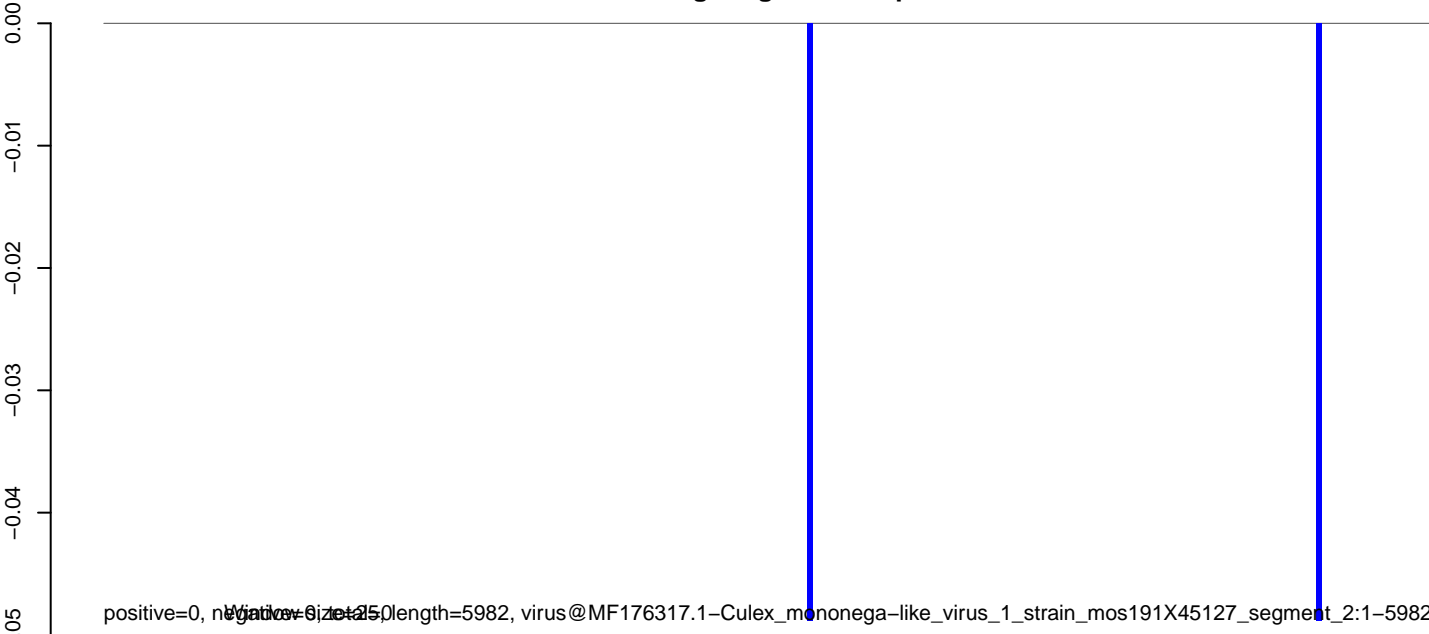
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



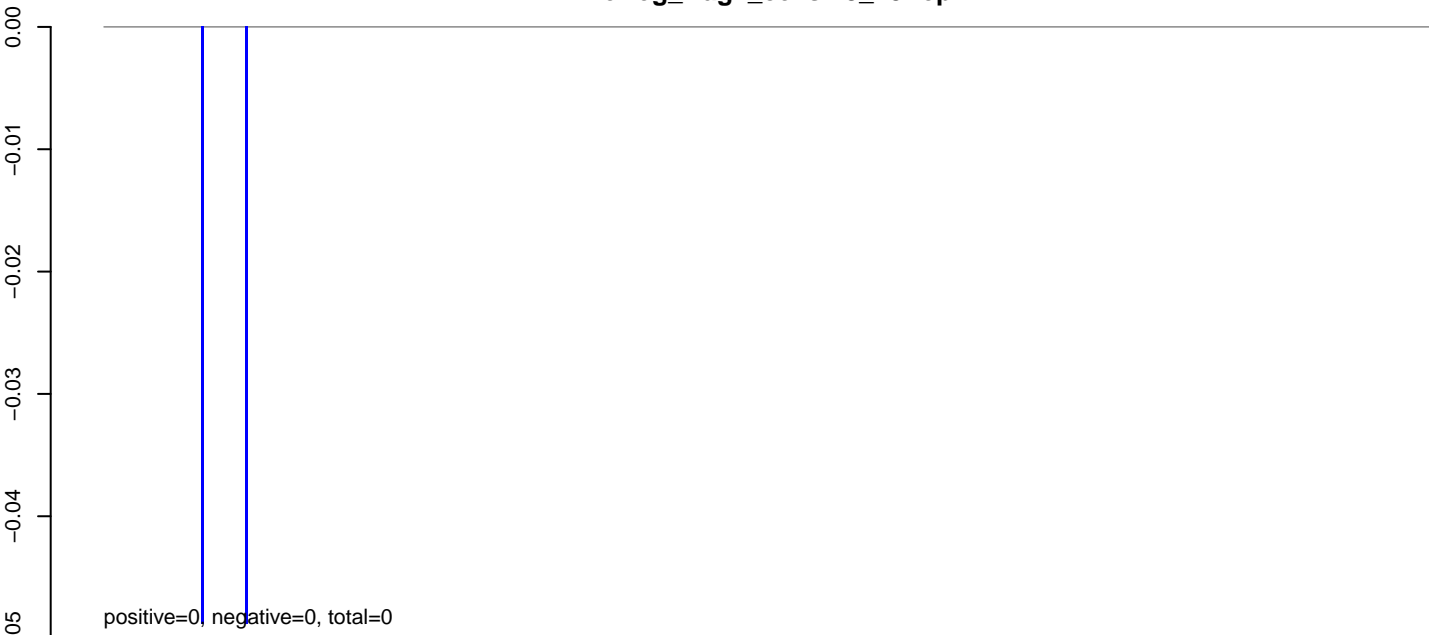
AeAeg_Aag2_cells.rep



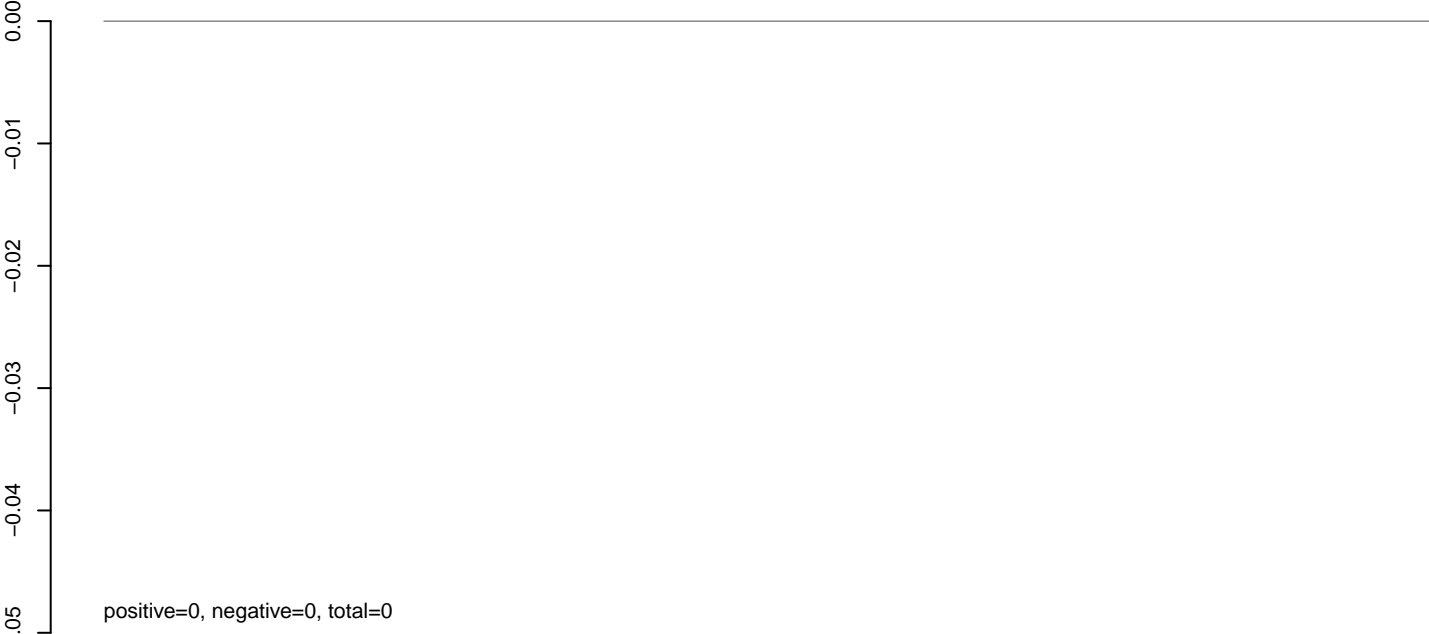
0 1000 2000 3000 4000 5000 6000

negative size=25 length=5982, virus@MF176317.1-Culex_moronega-like_virus_1_strain_mos191X45127_segment_2:1-5982

AeAeg_Aag2_cells.18_23.rep



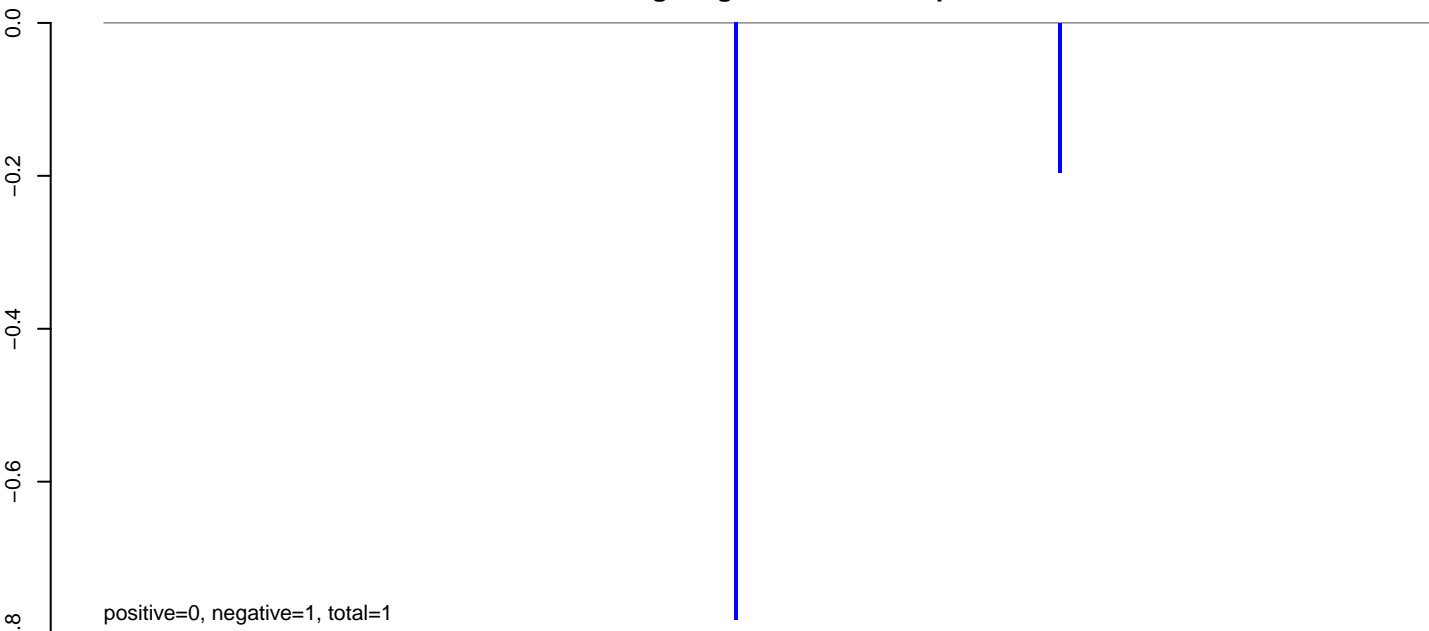
AeAeg_Aag2_cells.24_35.rep



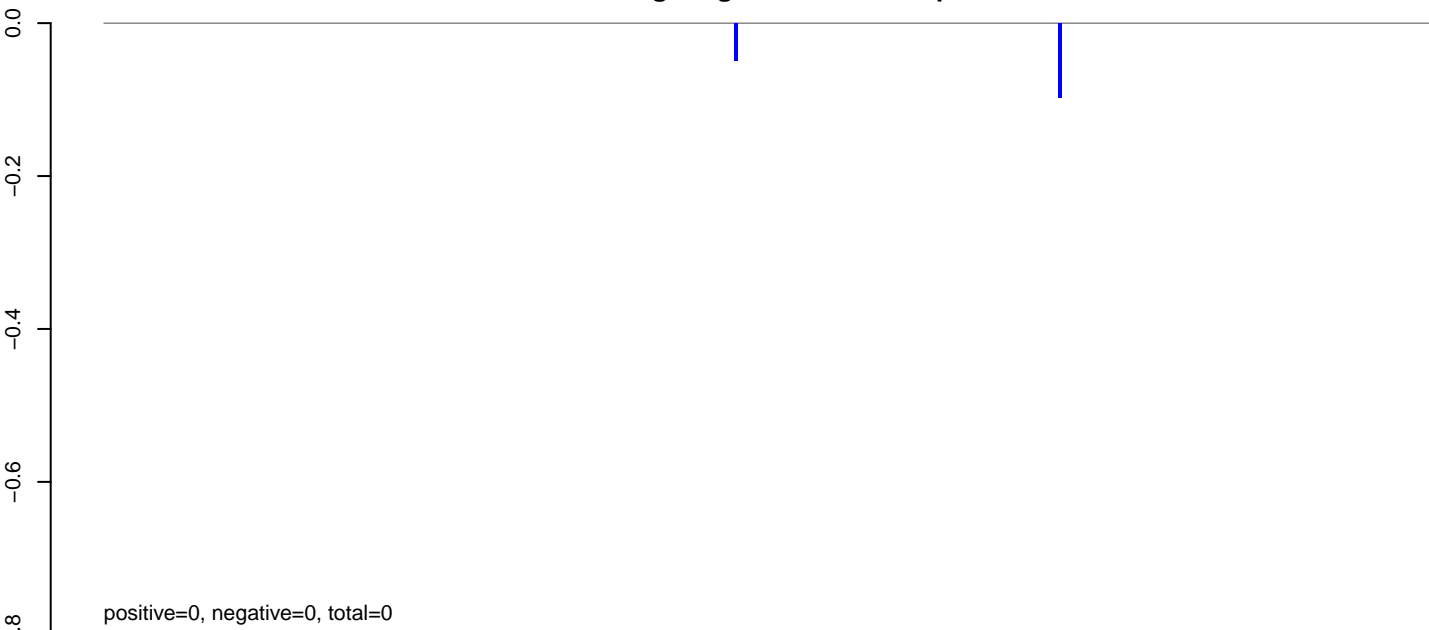
AeAeg_Aag2_cells.rep



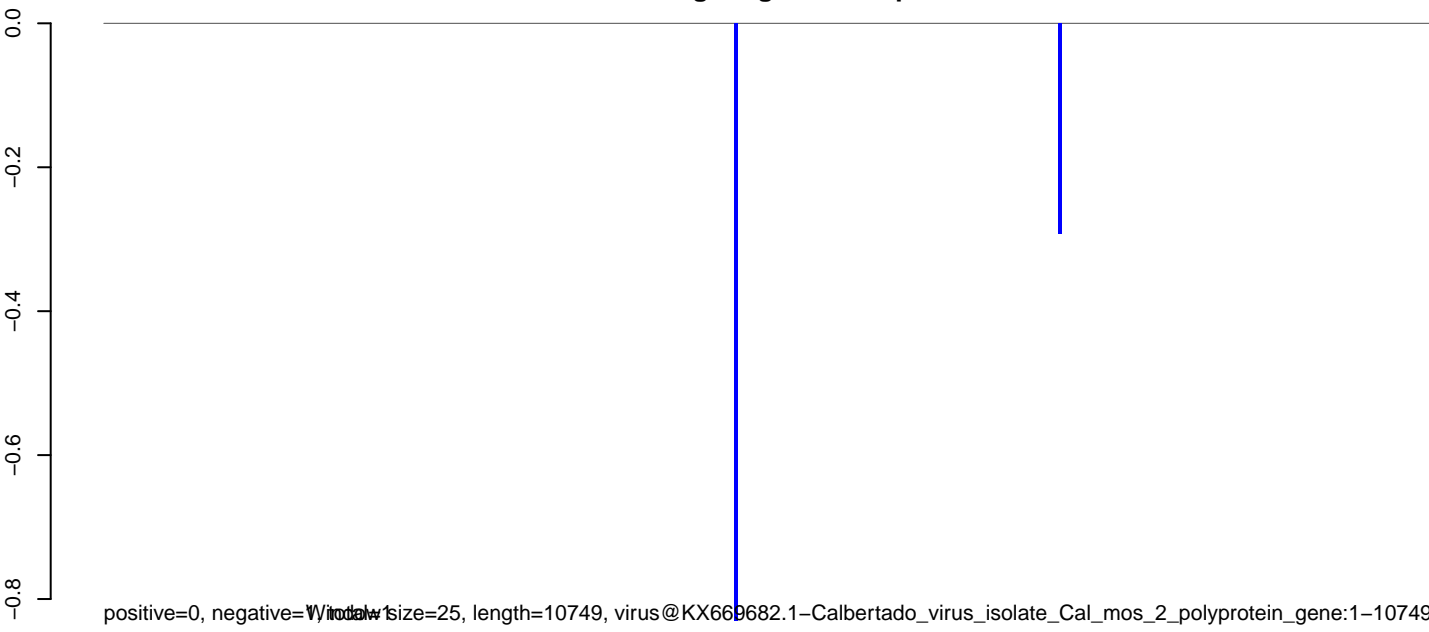
AeAeg_Aag2_cells.18_23.rep



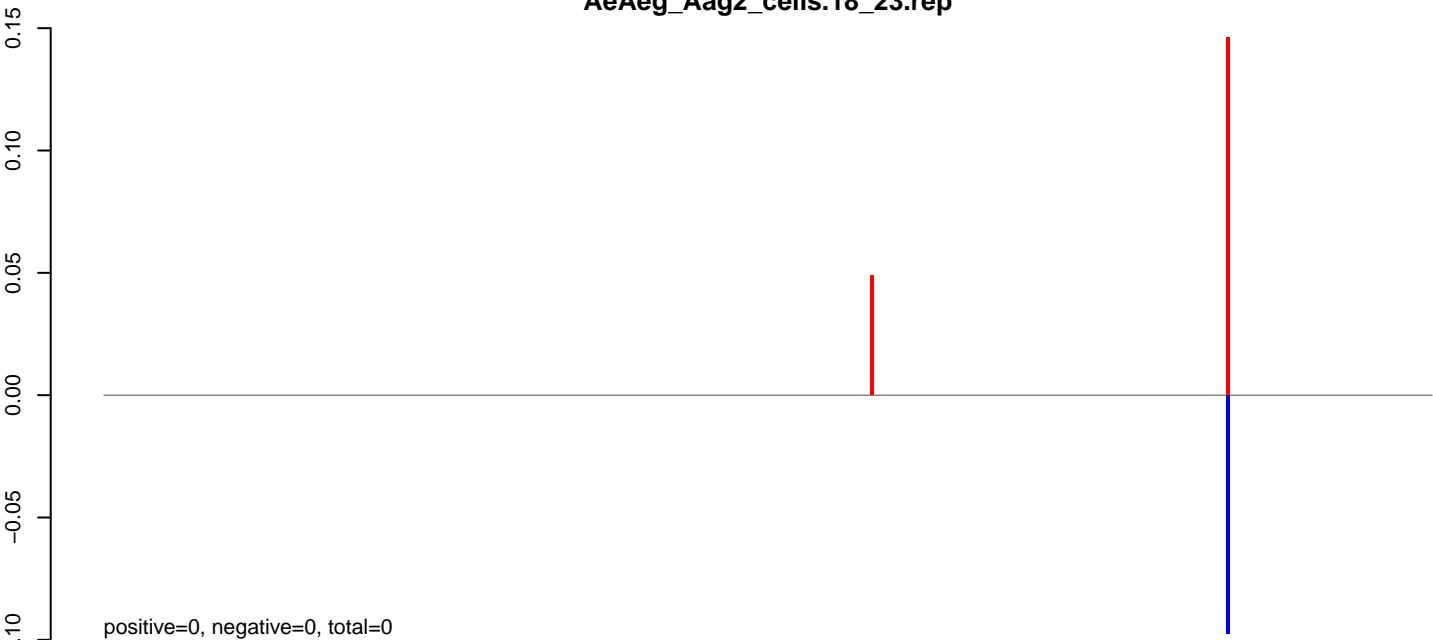
AeAeg_Aag2_cells.24_35.rep



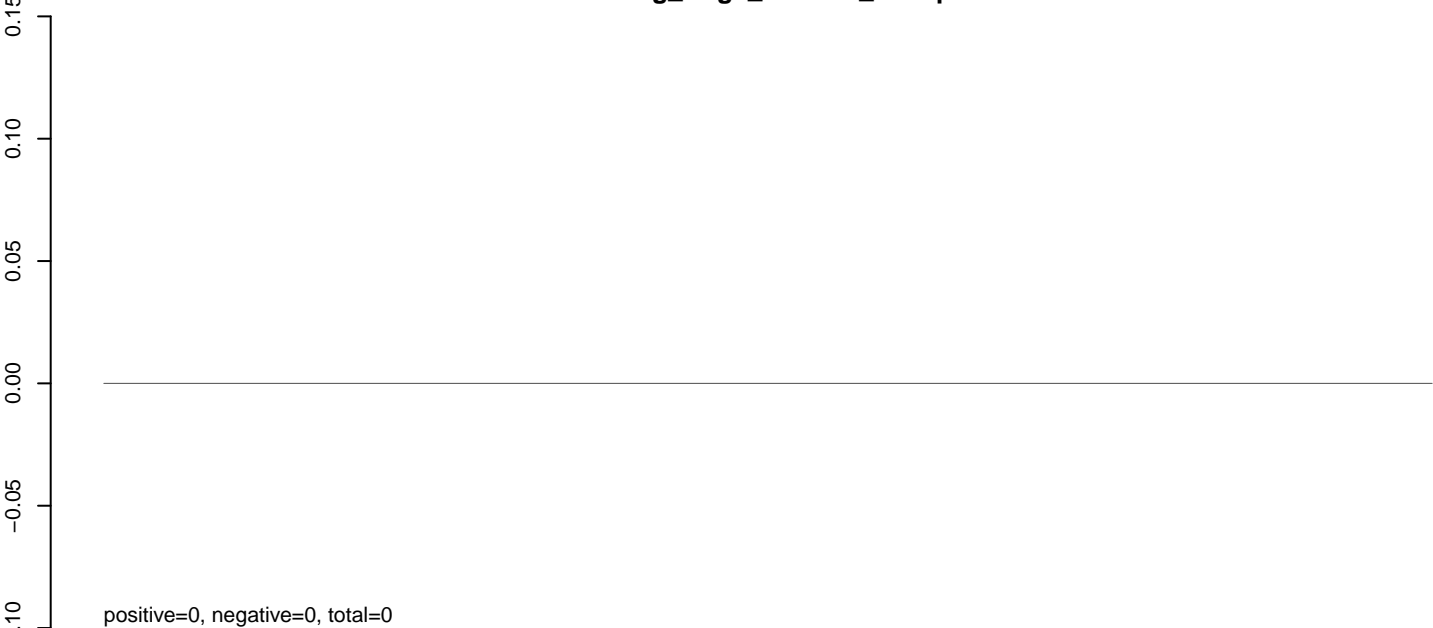
AeAeg_Aag2_cells.rep



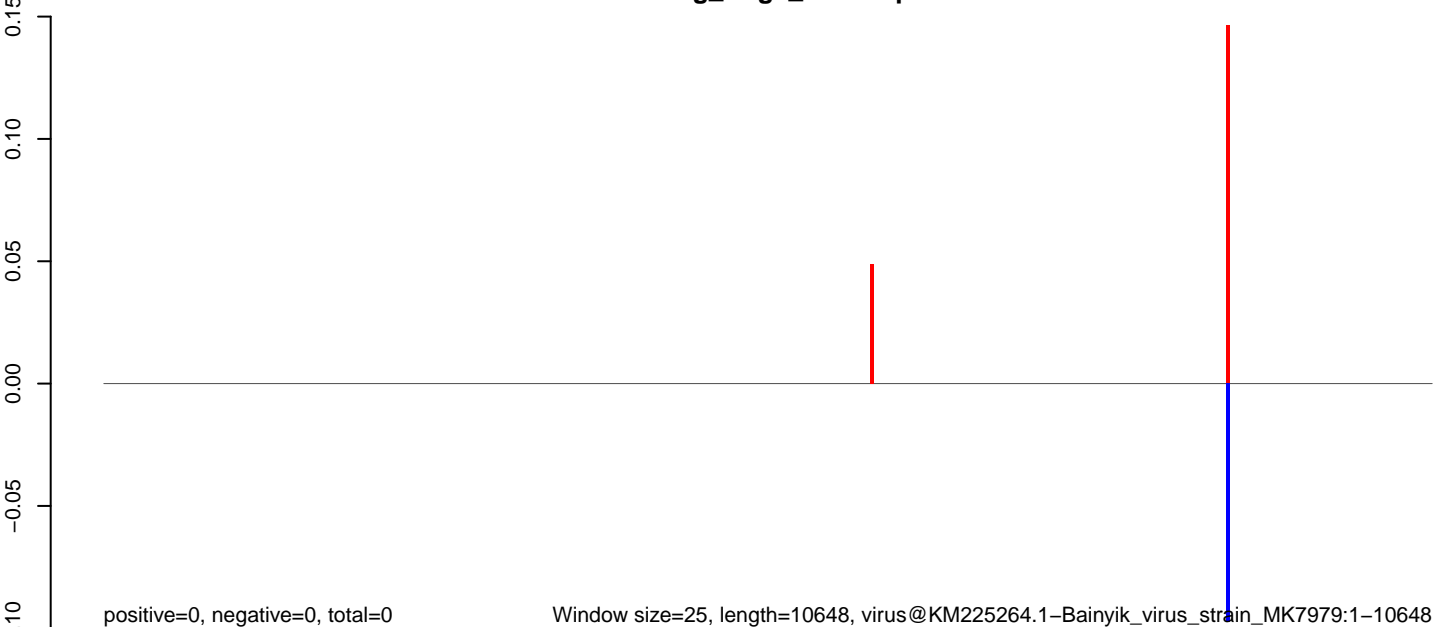
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



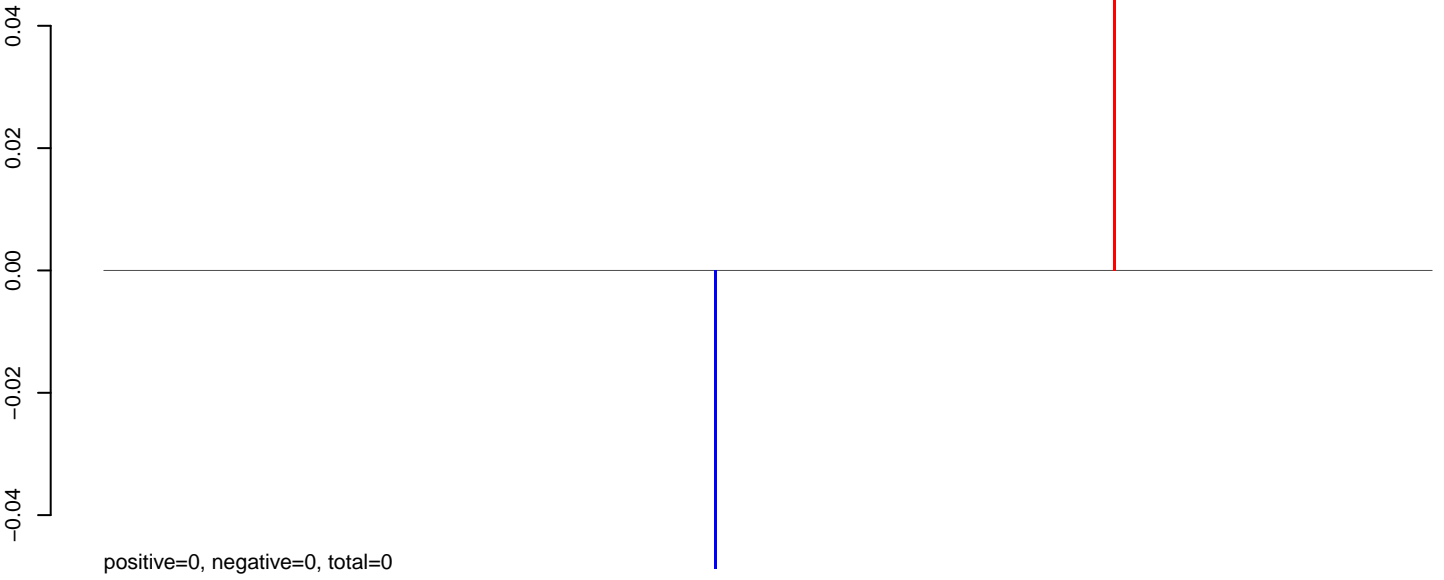
AeAeg_Aag2_cells.rep



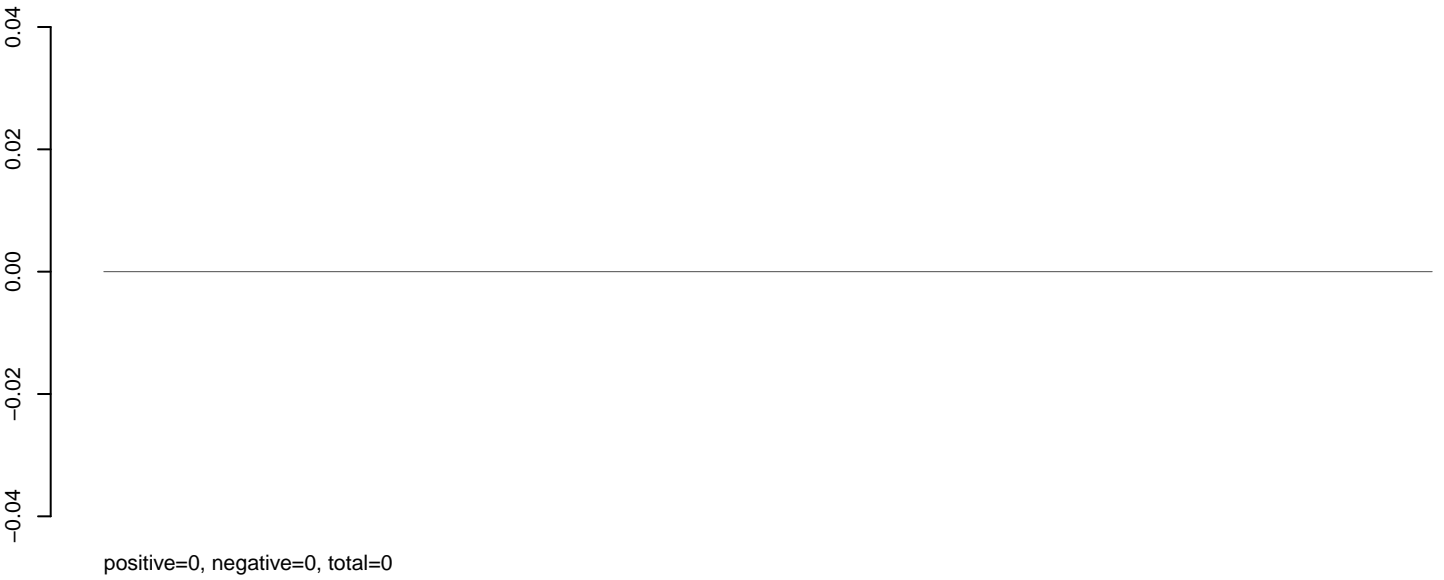
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0 2000 4000 6000 8000 10000

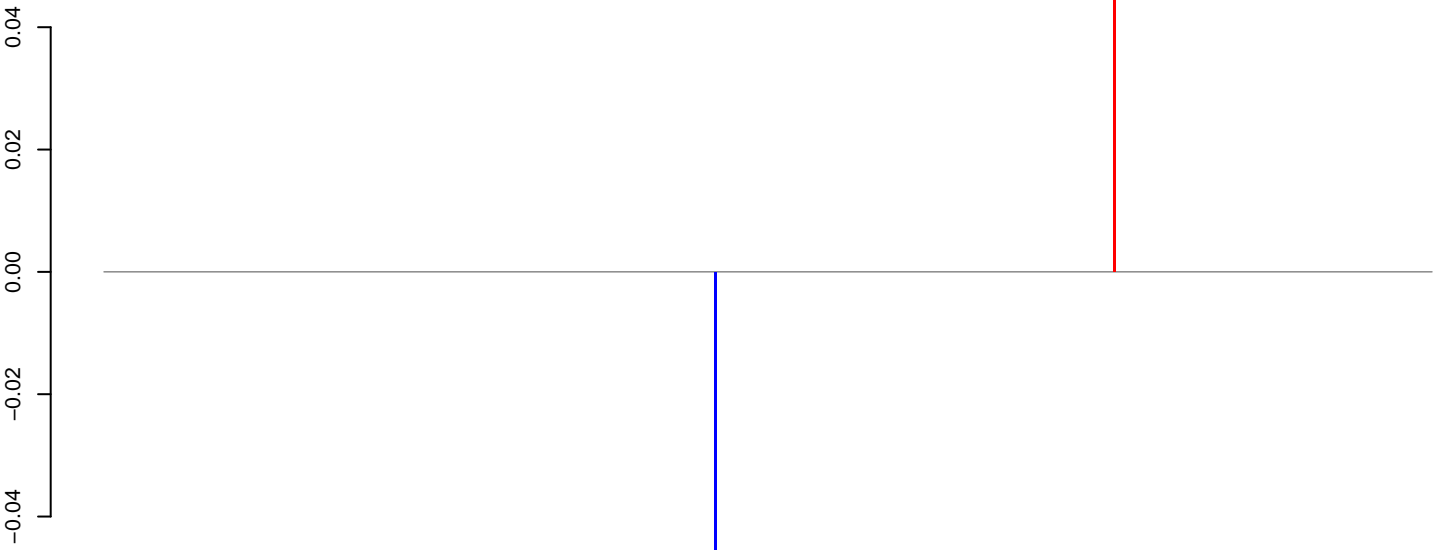
AeAeg_Aag2_cells.18_23.rep



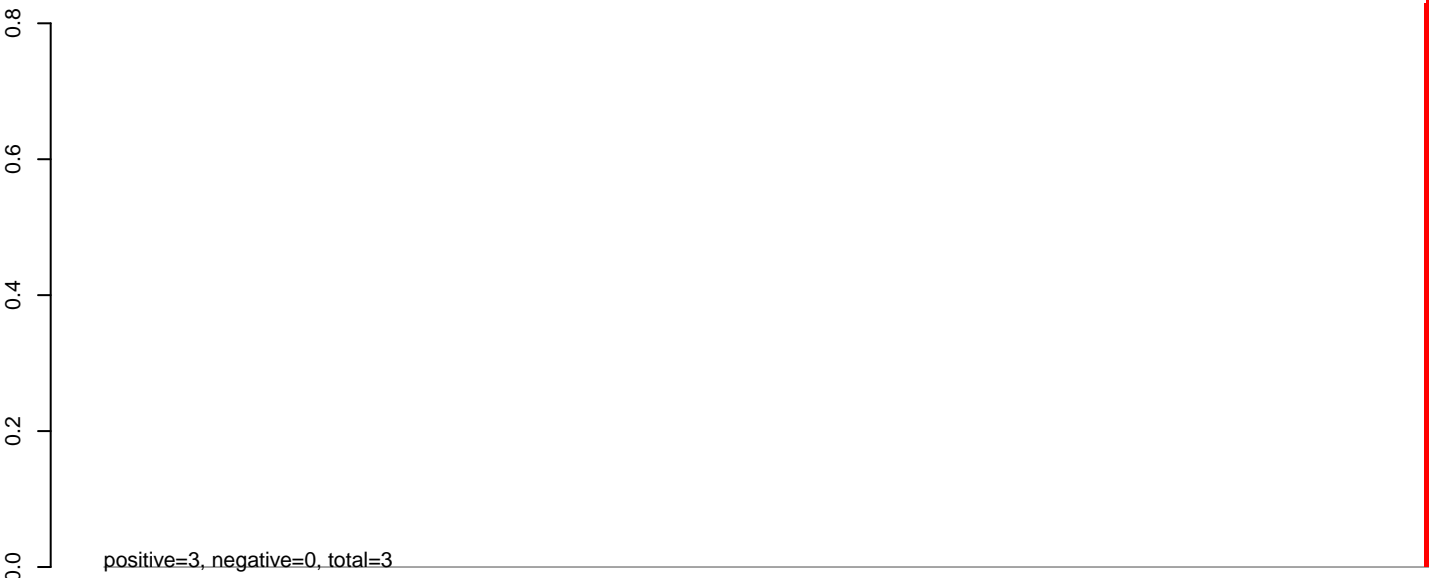
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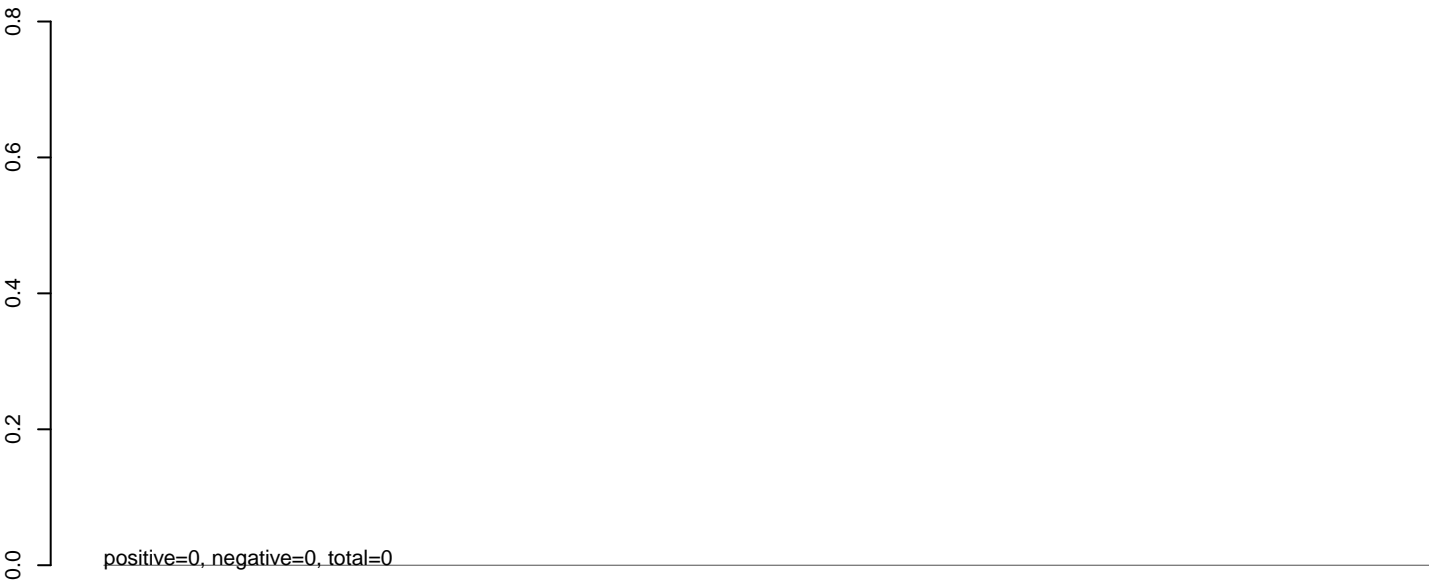
AeAeg_Aag2_cells.rep



AeAeg_Aag2_cells.18_23.rep



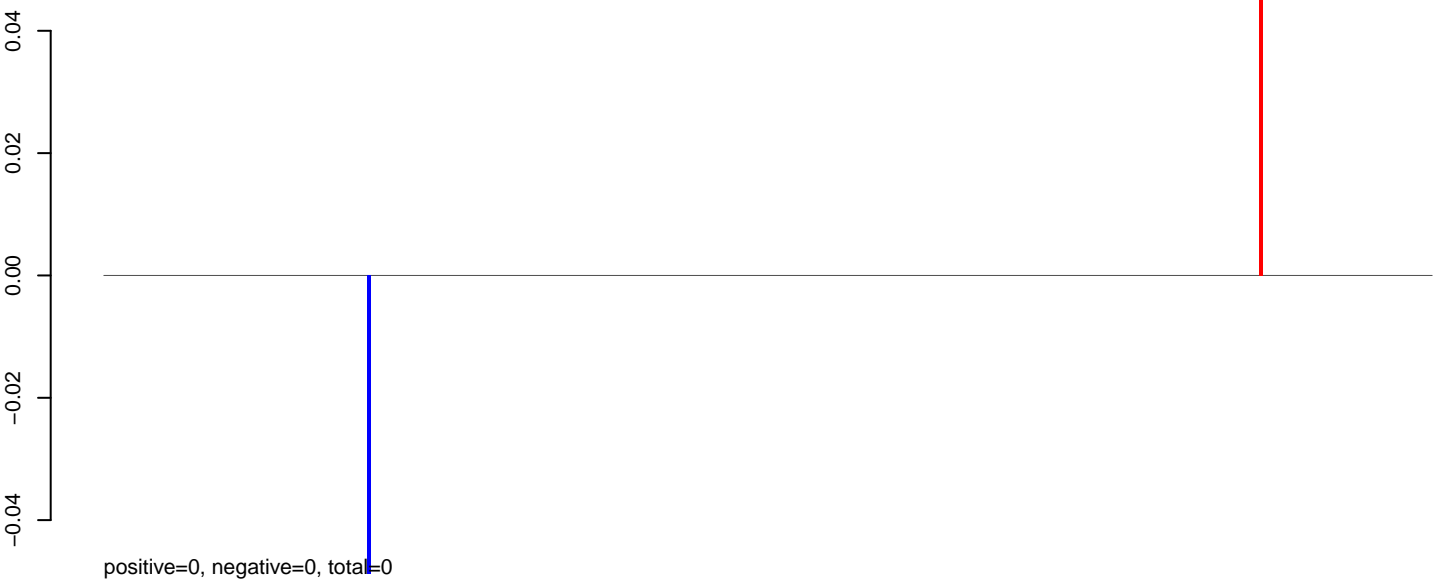
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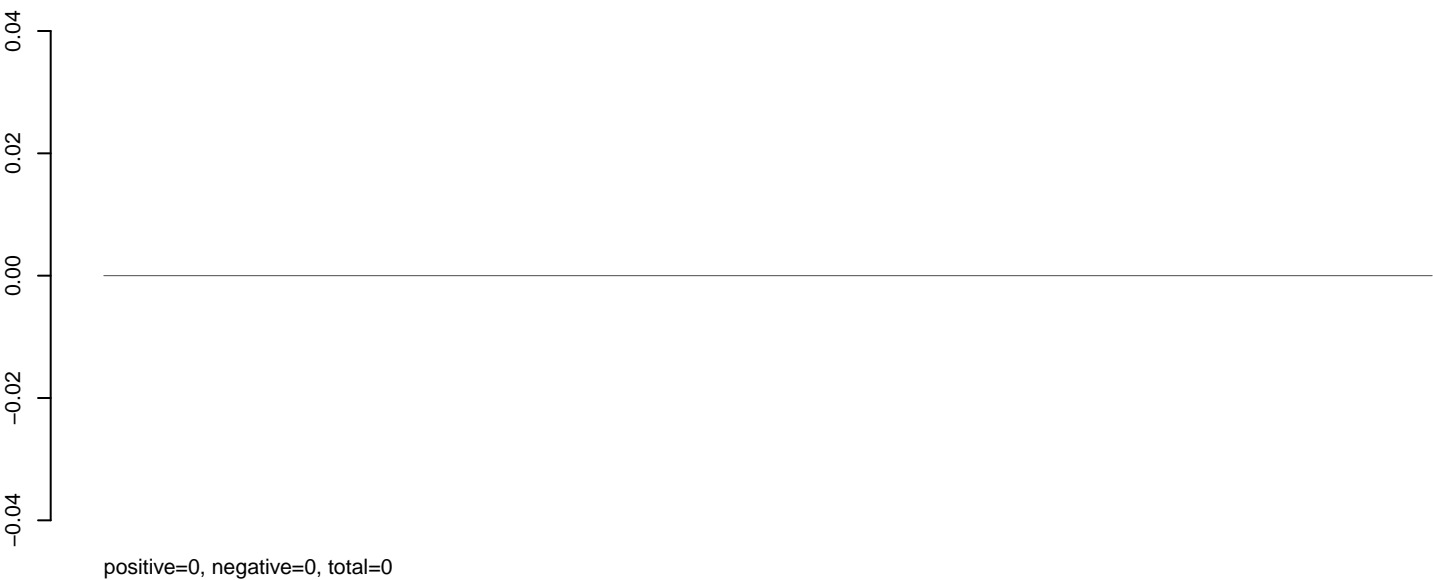
AeAeg_Aag2_cells.rep



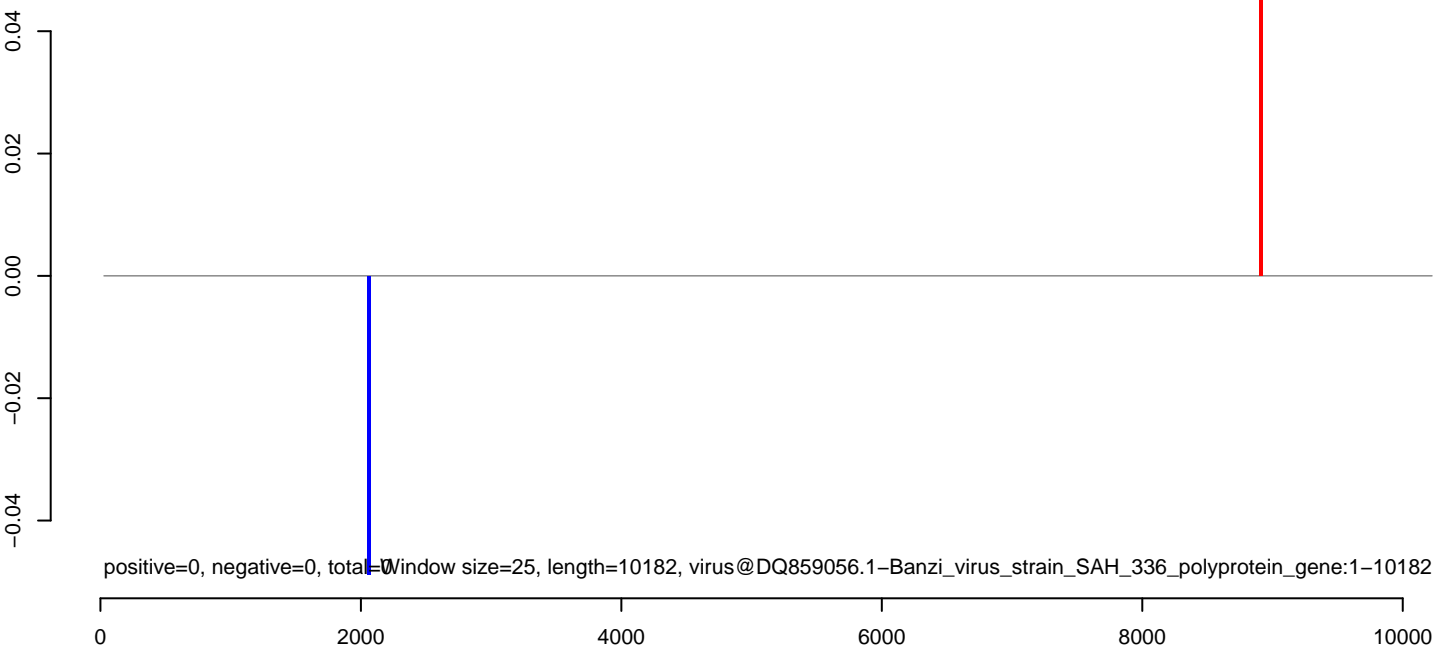
AeAeg_Aag2_cells.18_23.rep



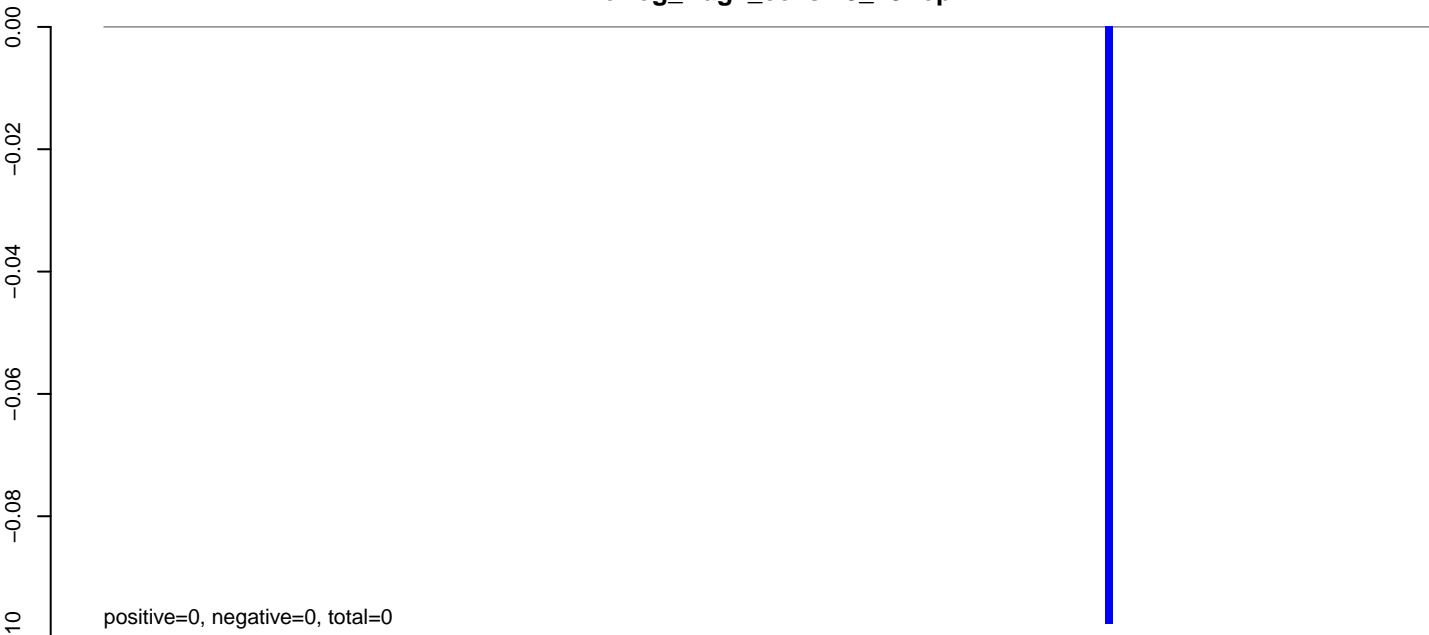
AeAeg_Aag2_cells.24_35.rep



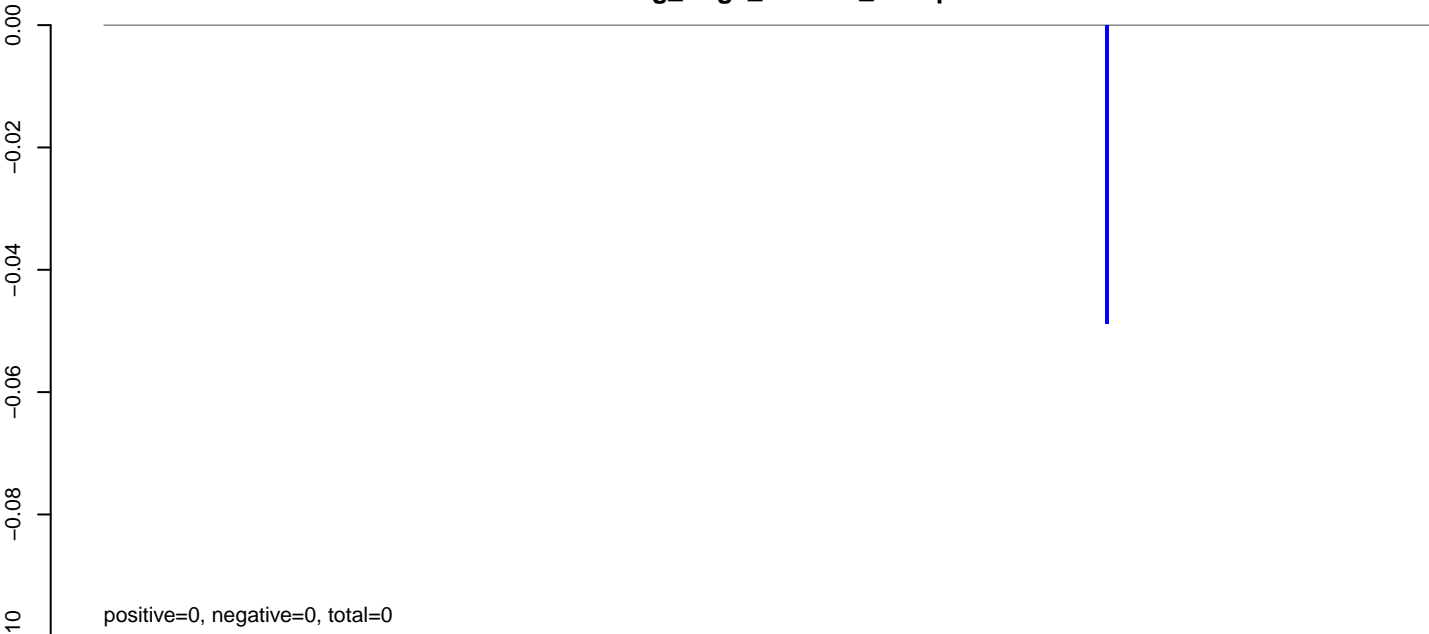
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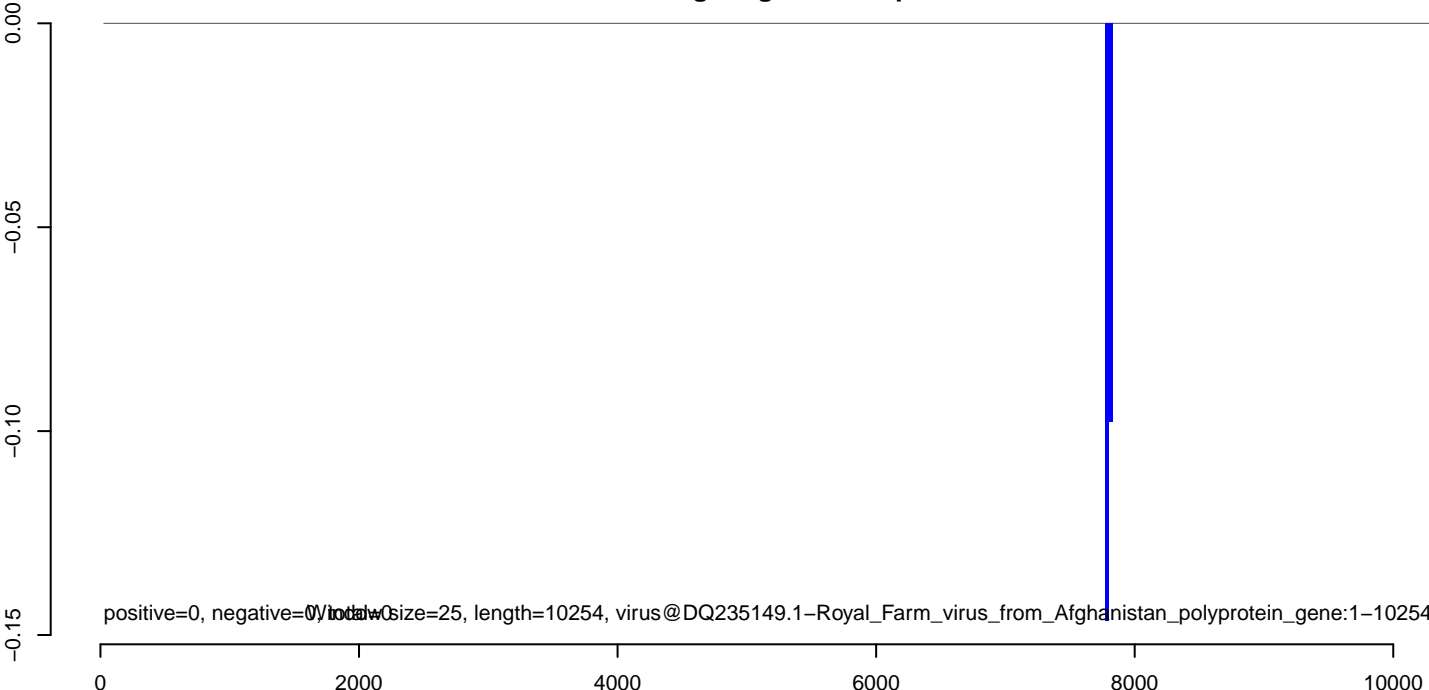
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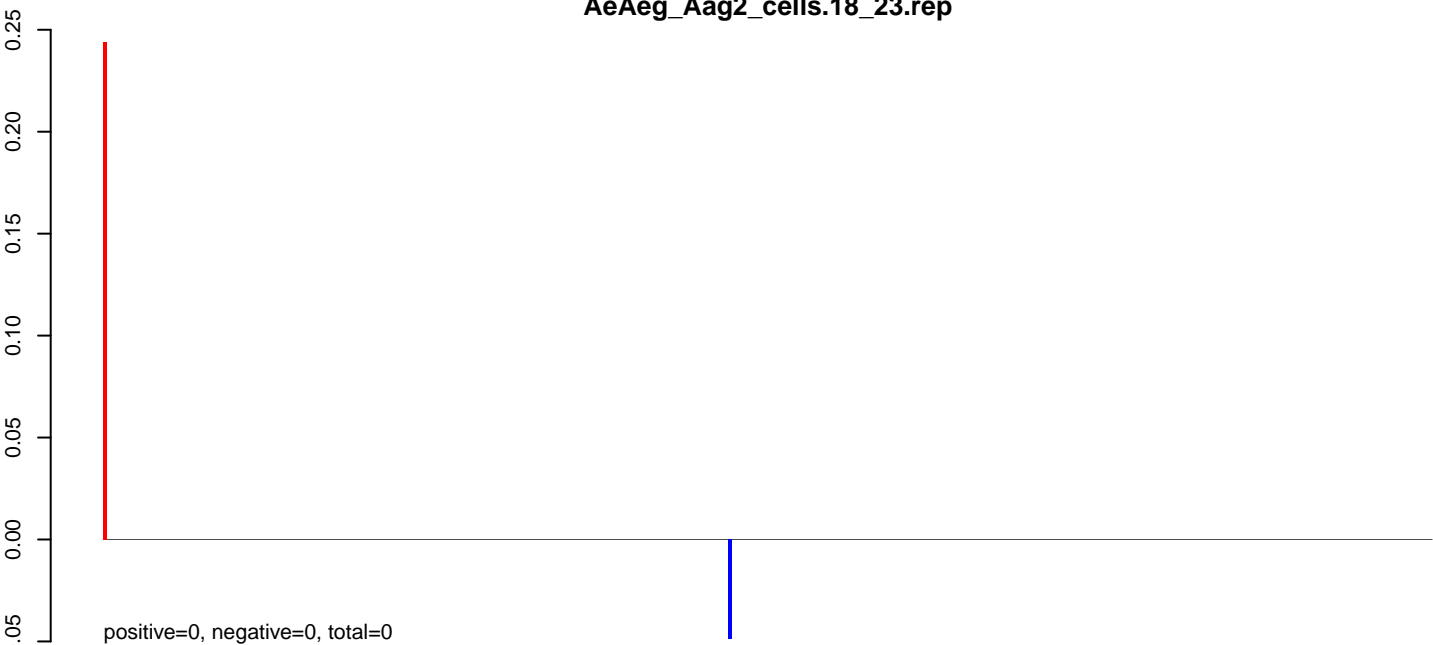
AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep

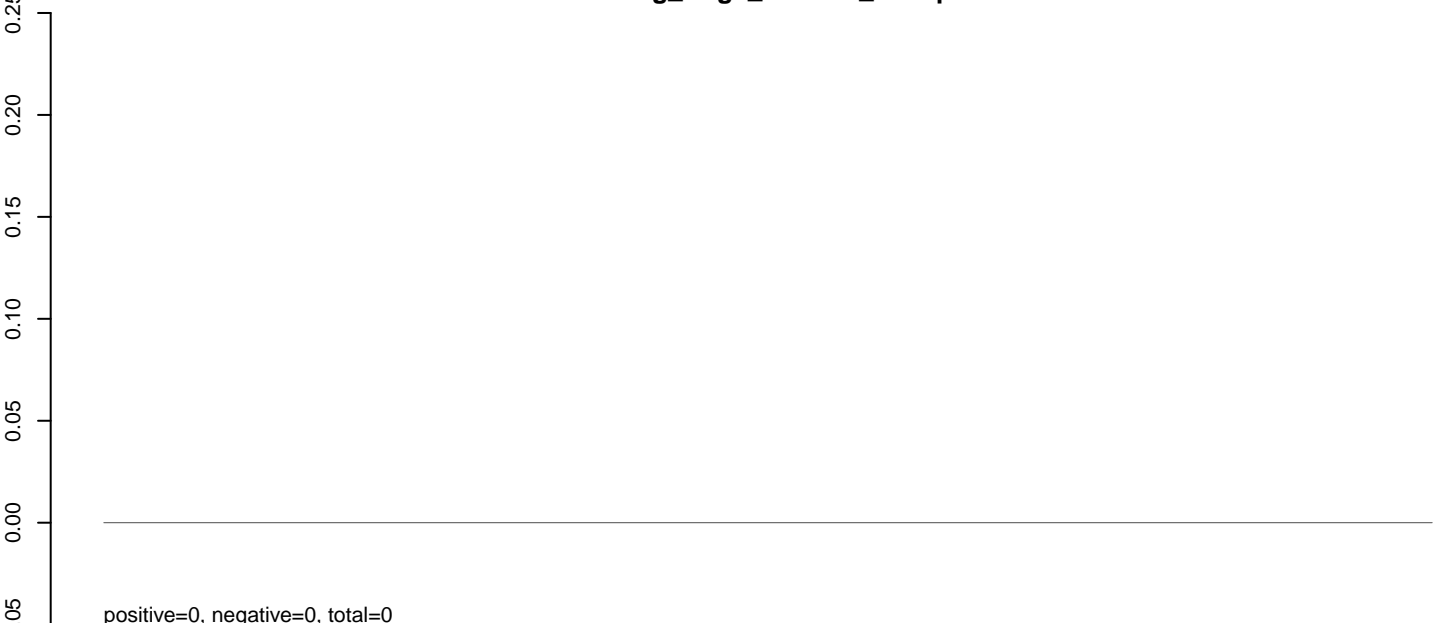


AeAeg_Aag2_cells.18_23.rep



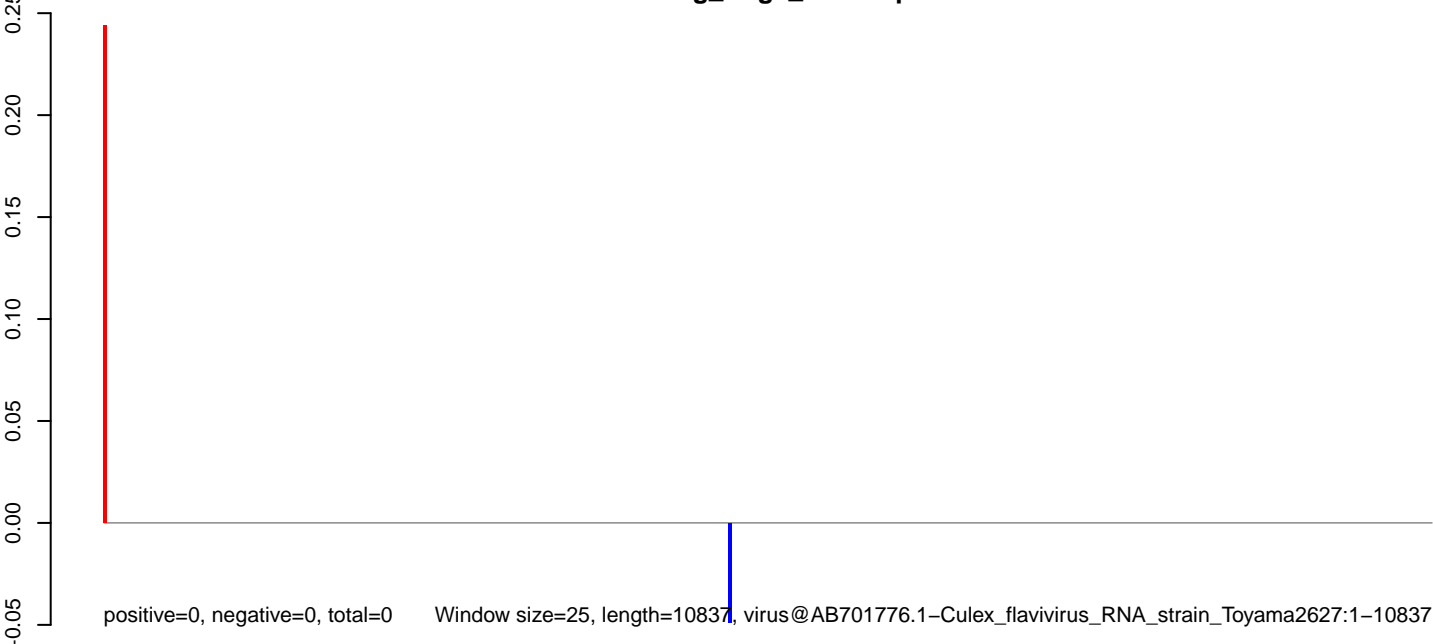
positive=0, negative=0, total=0

AeAeg_Aag2_cells.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_cells.rep

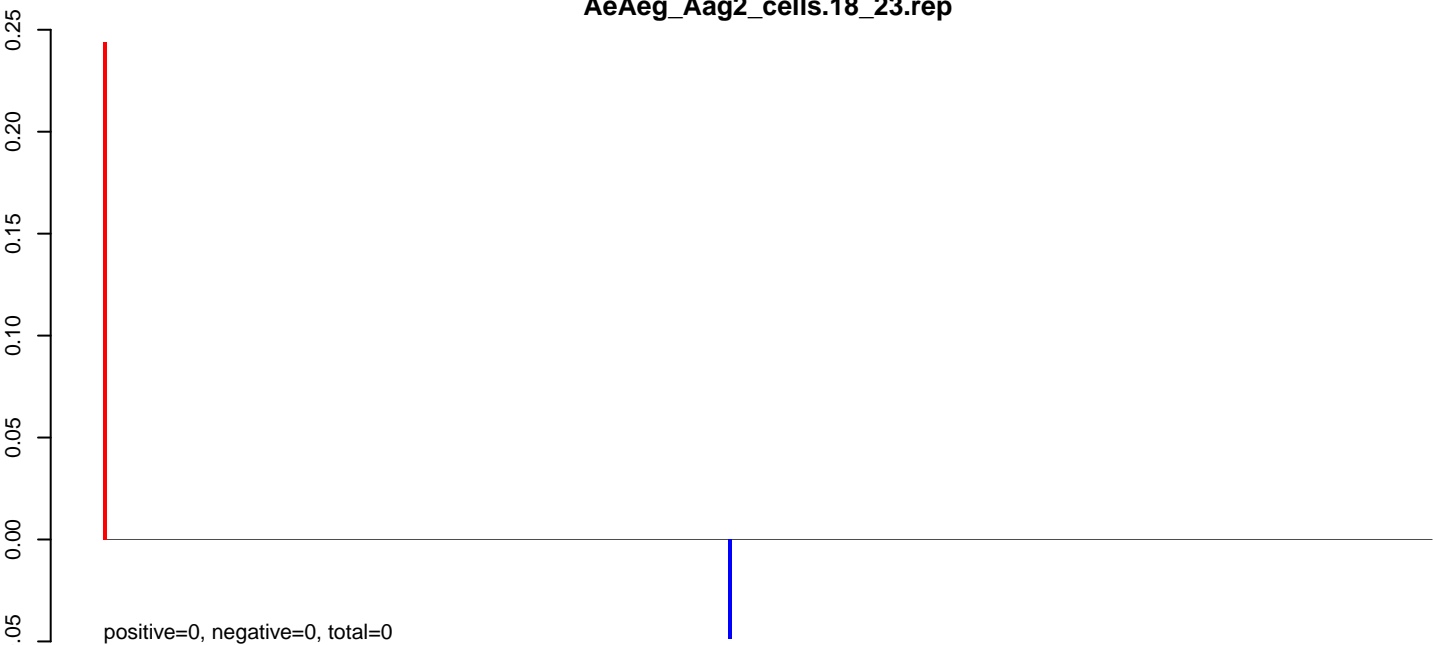


positive=0, negative=0, total=0

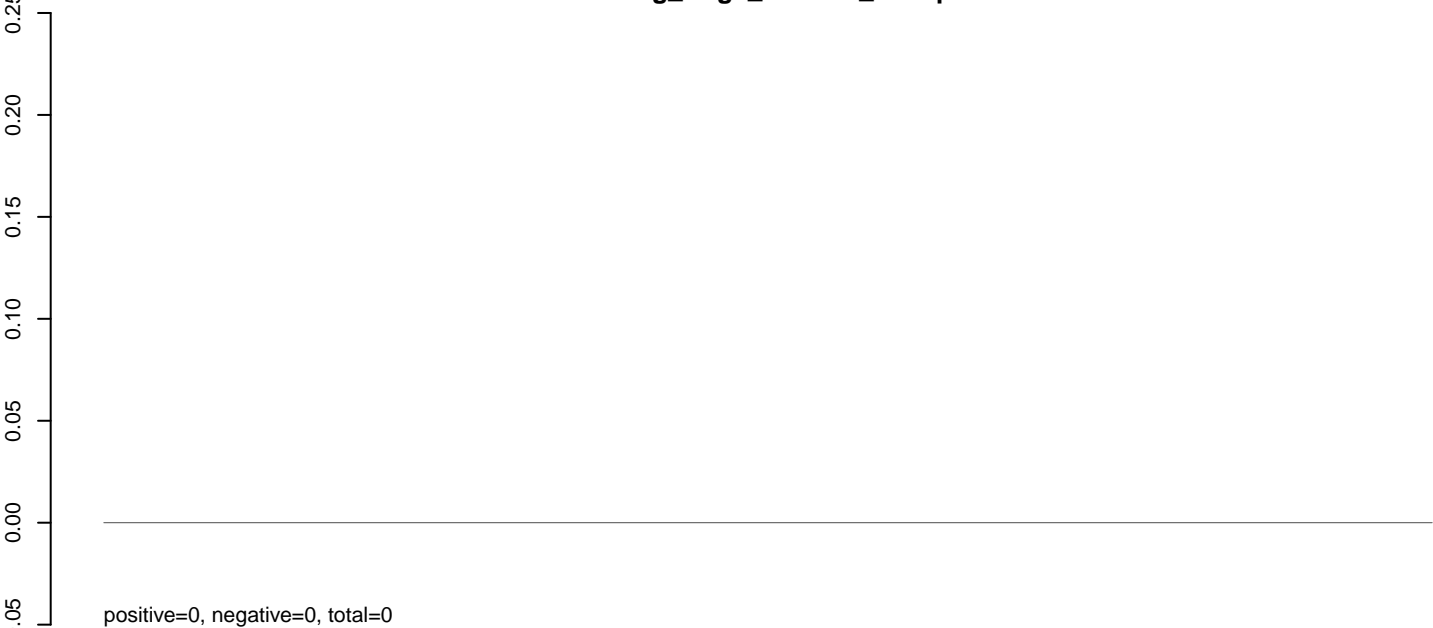
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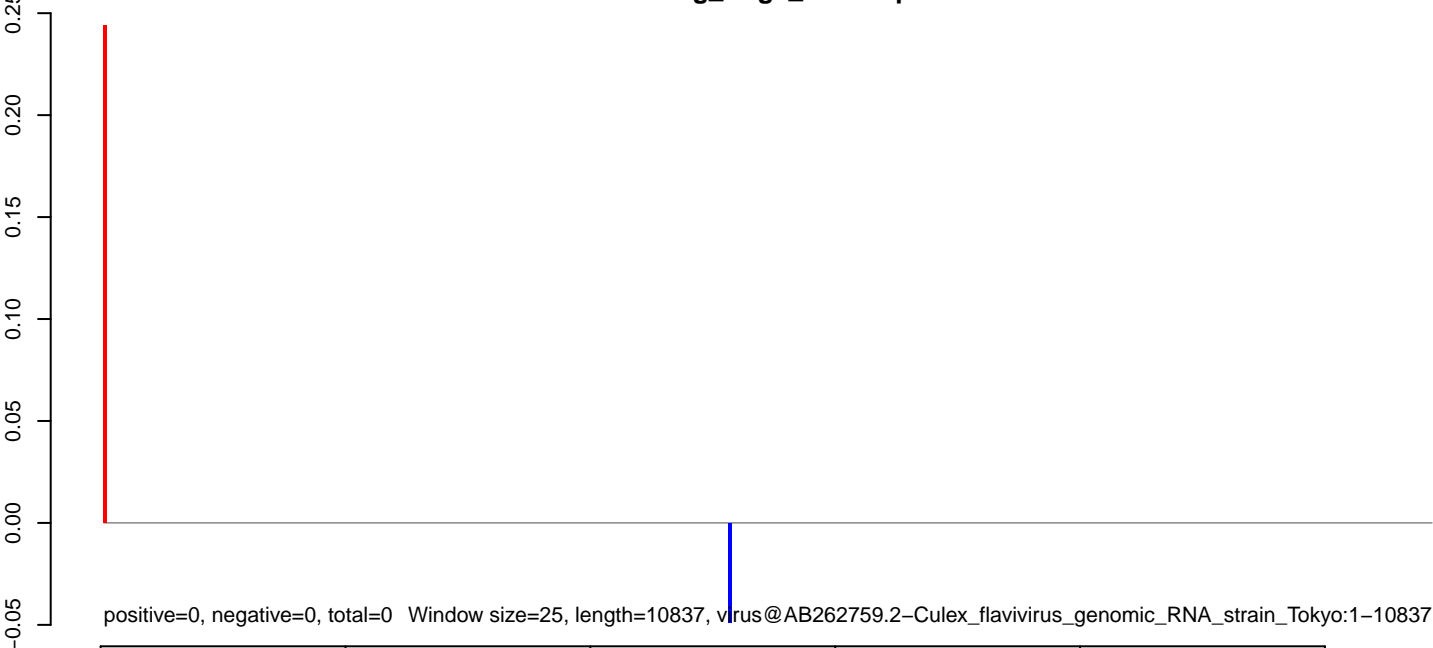
AeAeg_Aag2_cells.18_23.rep



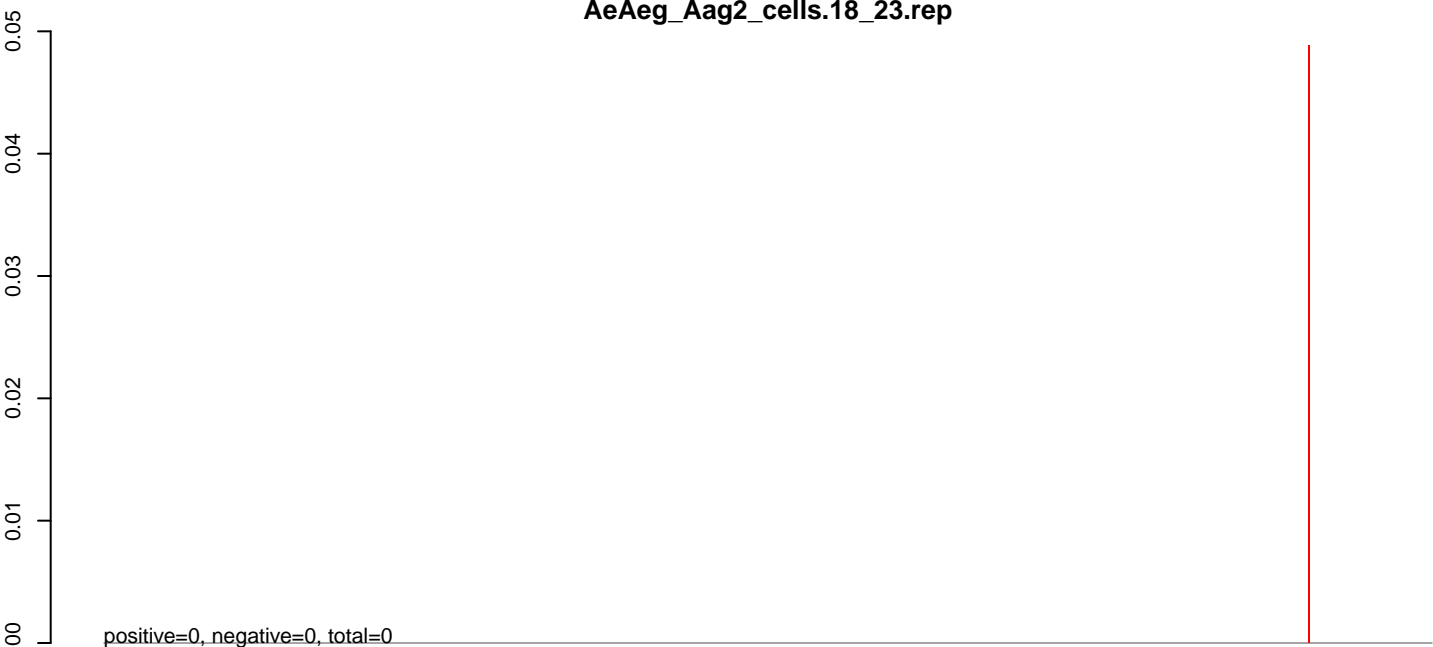
AeAeg_Aag2_cells.24_35.rep



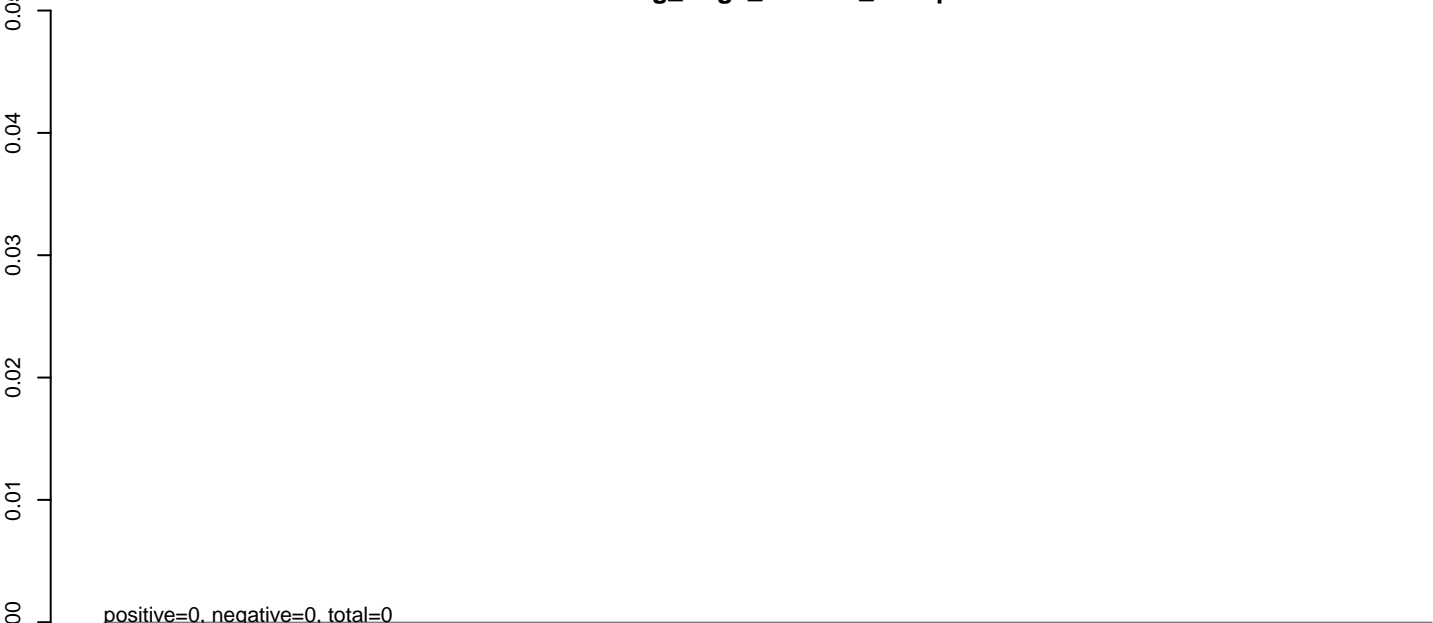
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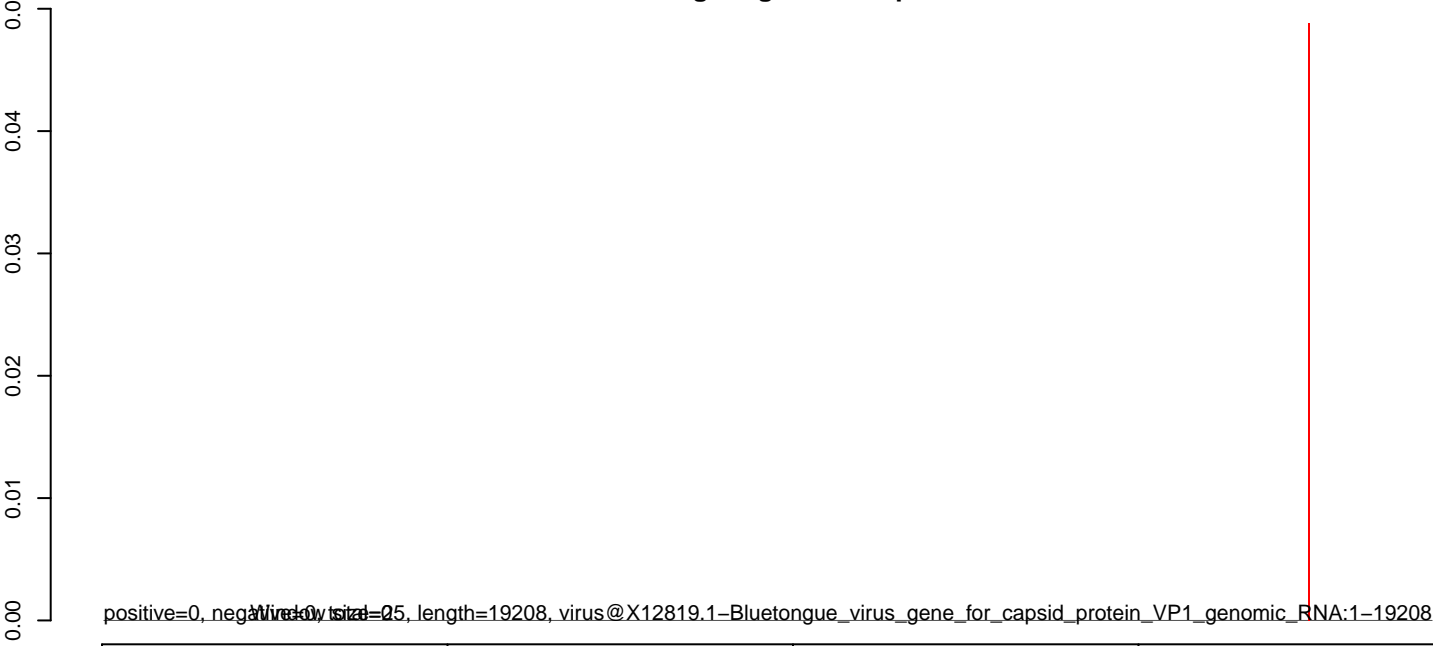
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AeAeg_Aag2_cells.24_35.rep

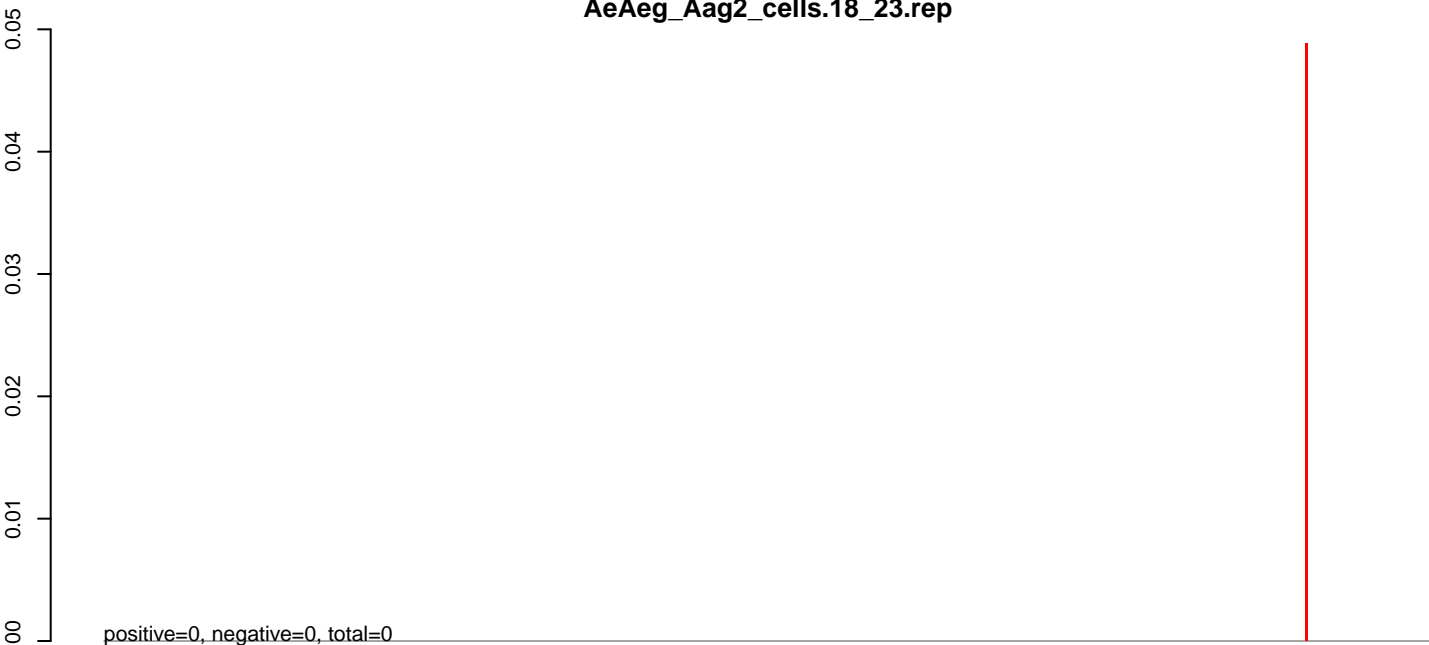


AeAeg_Aag2_cells.rep

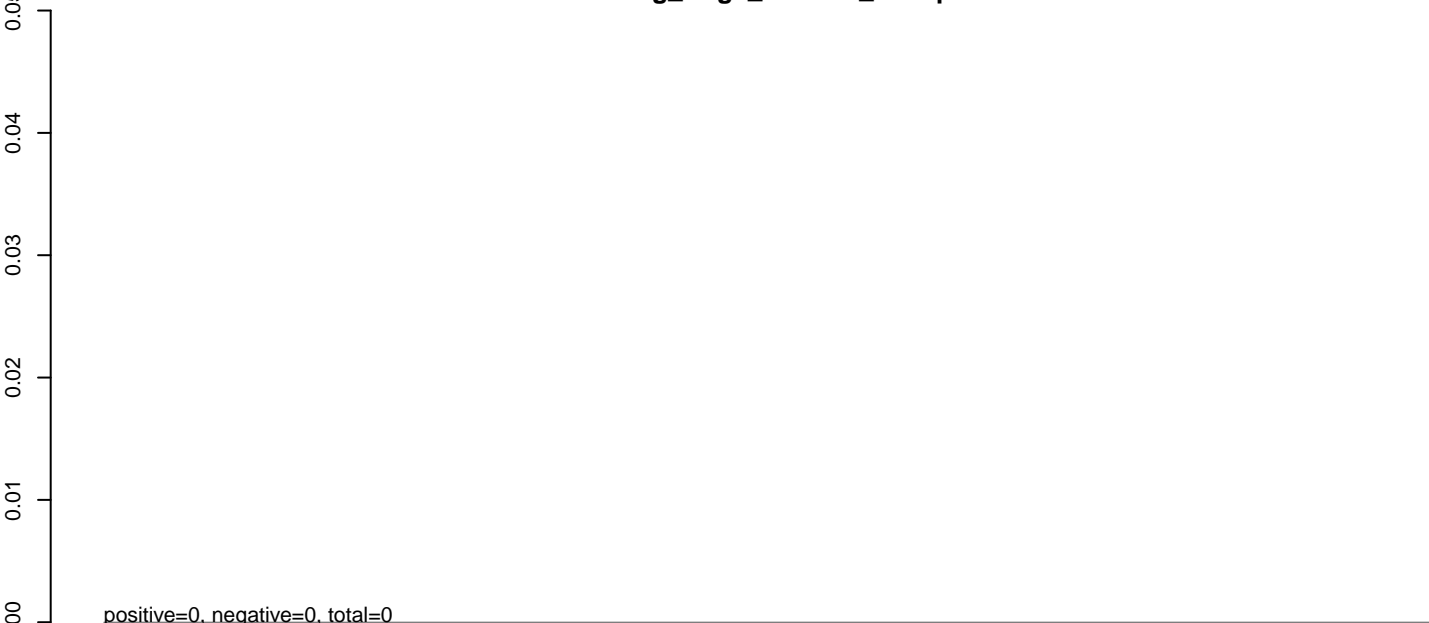


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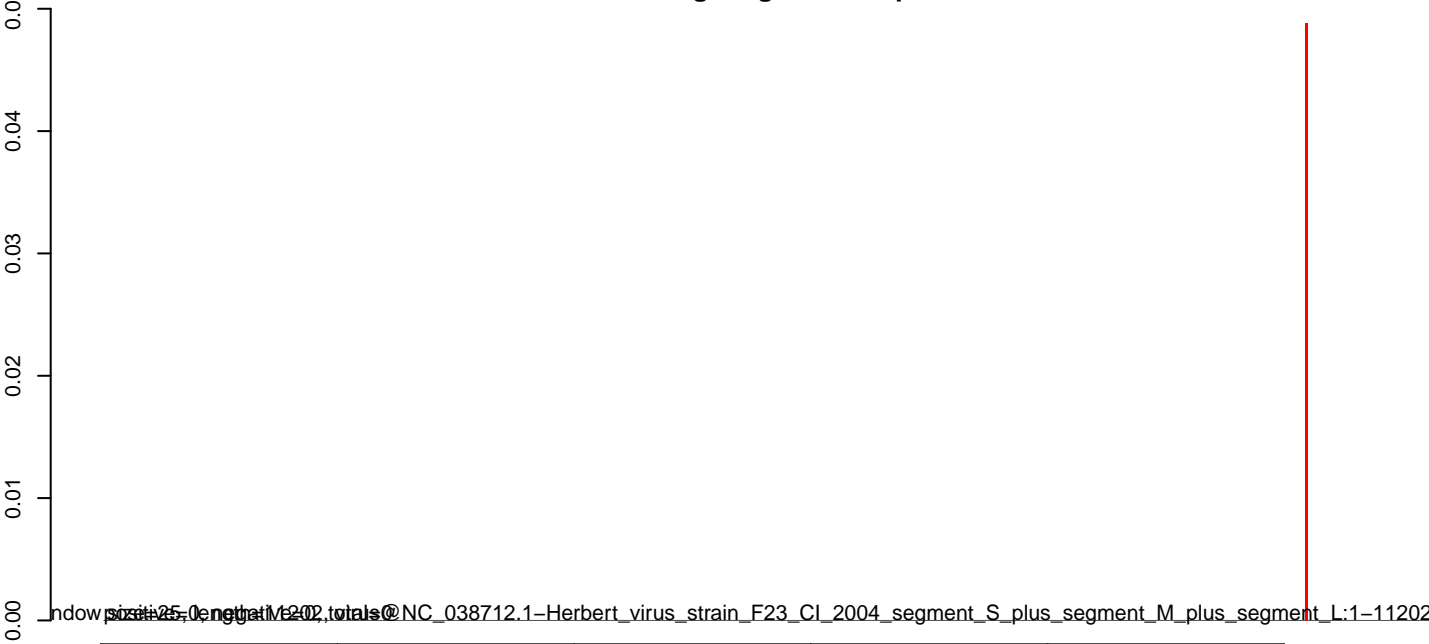
AeAeg_Aag2_cells.18_23.rep



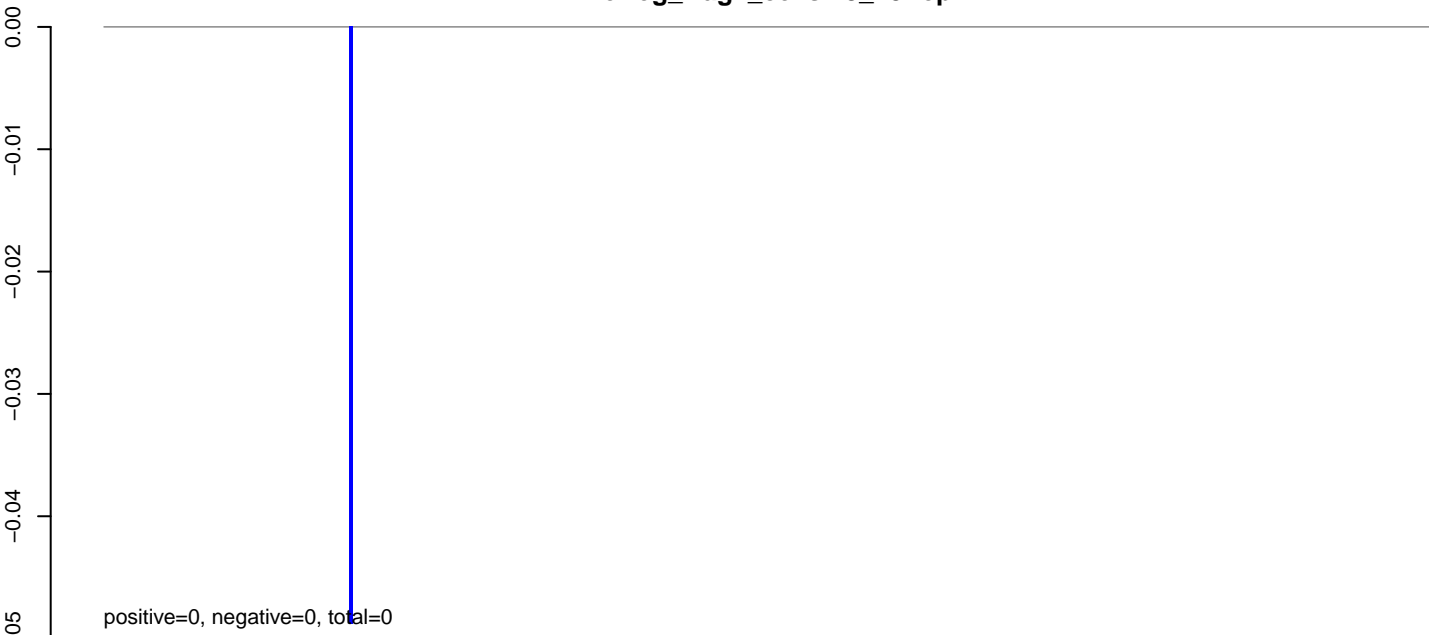
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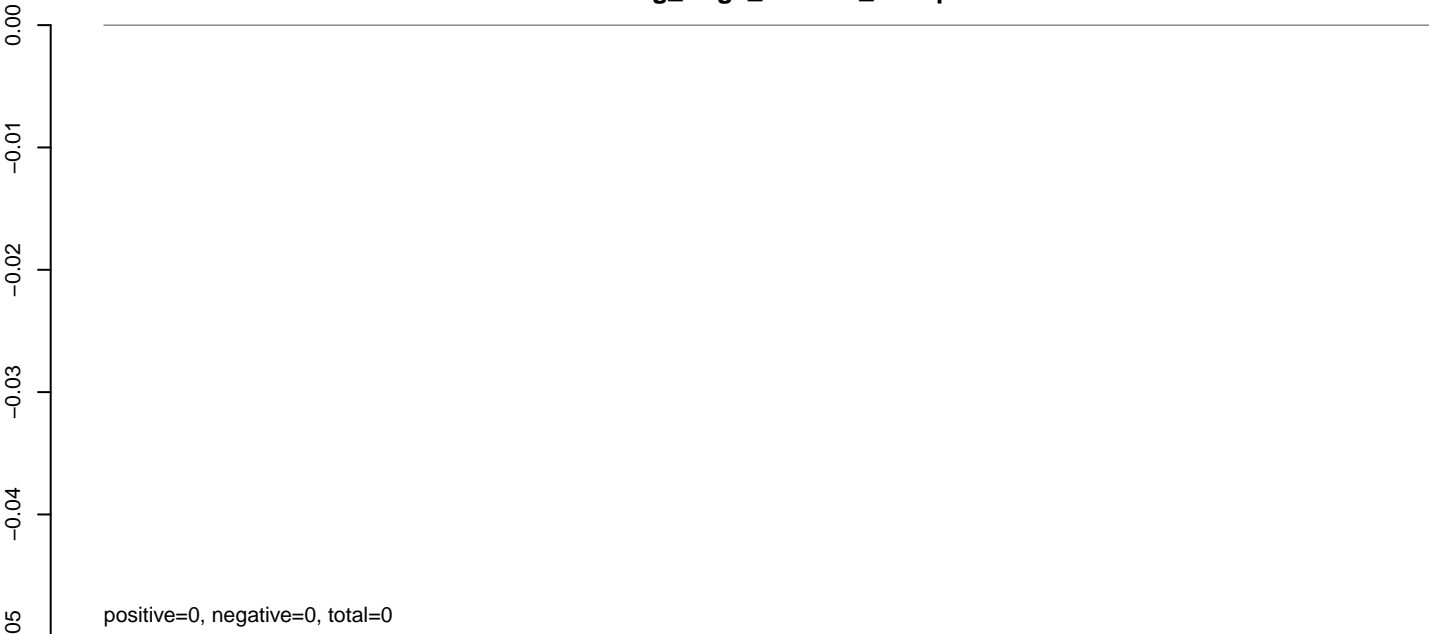
AeAeg_Aag2_cells.rep



AeAeg_Aag2_cells.18_23.rep



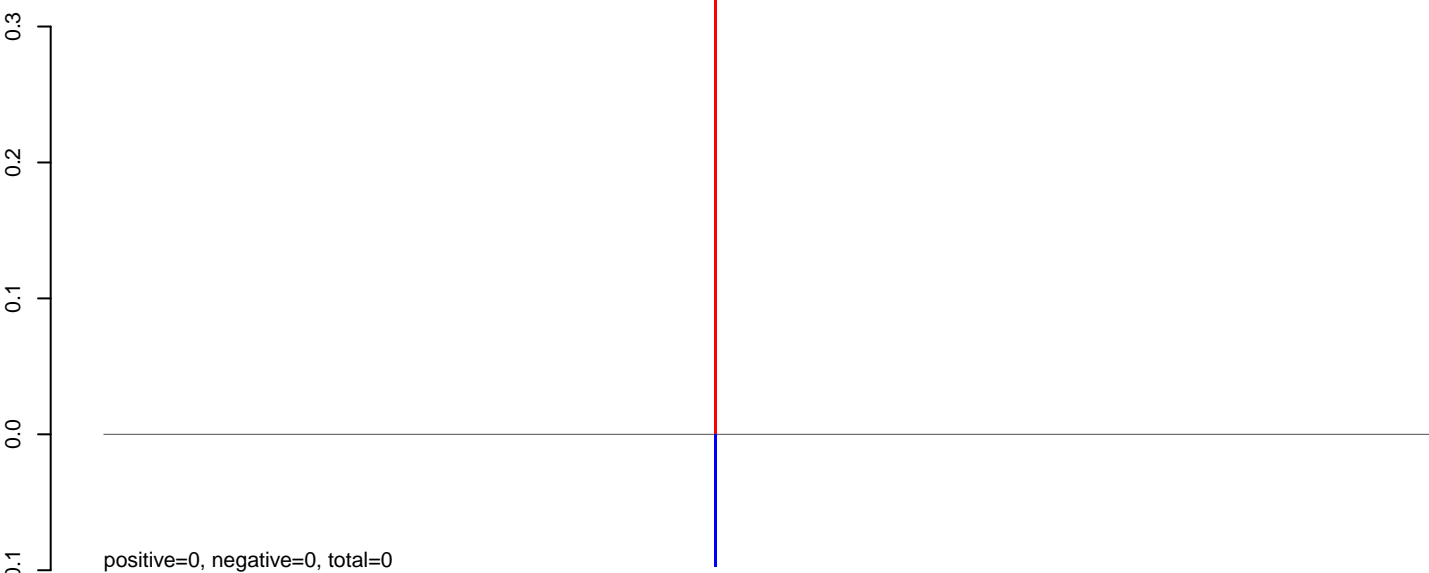
AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep



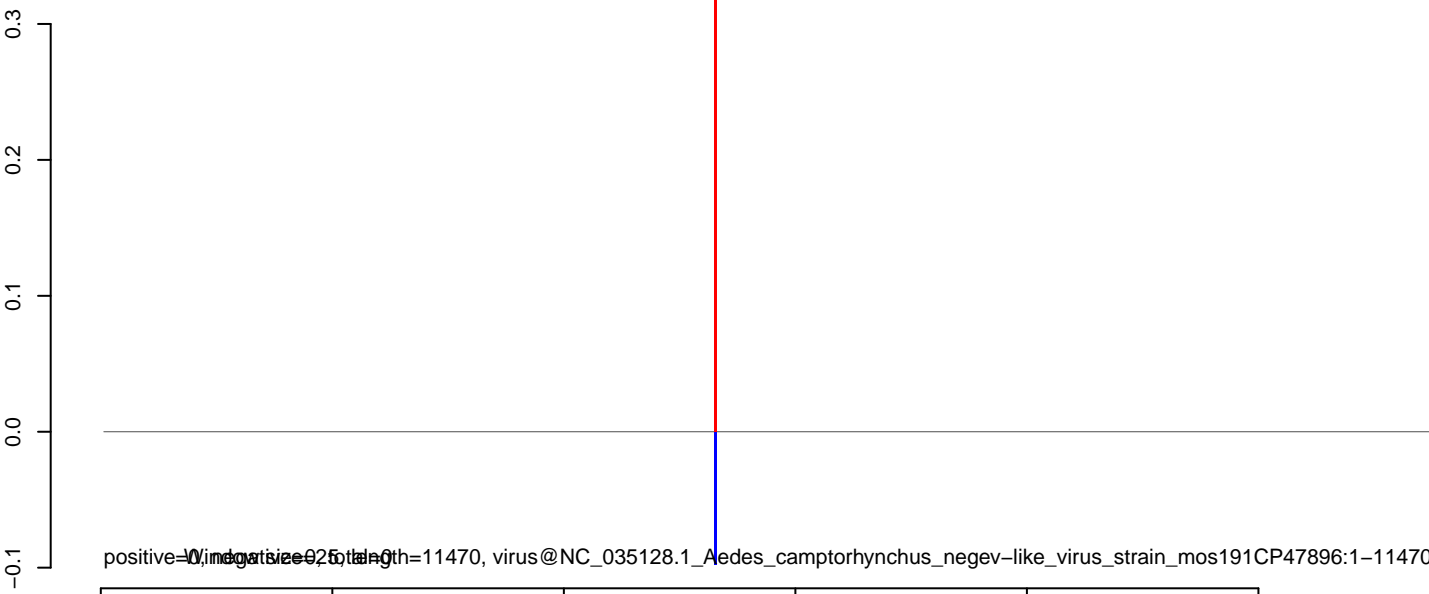
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

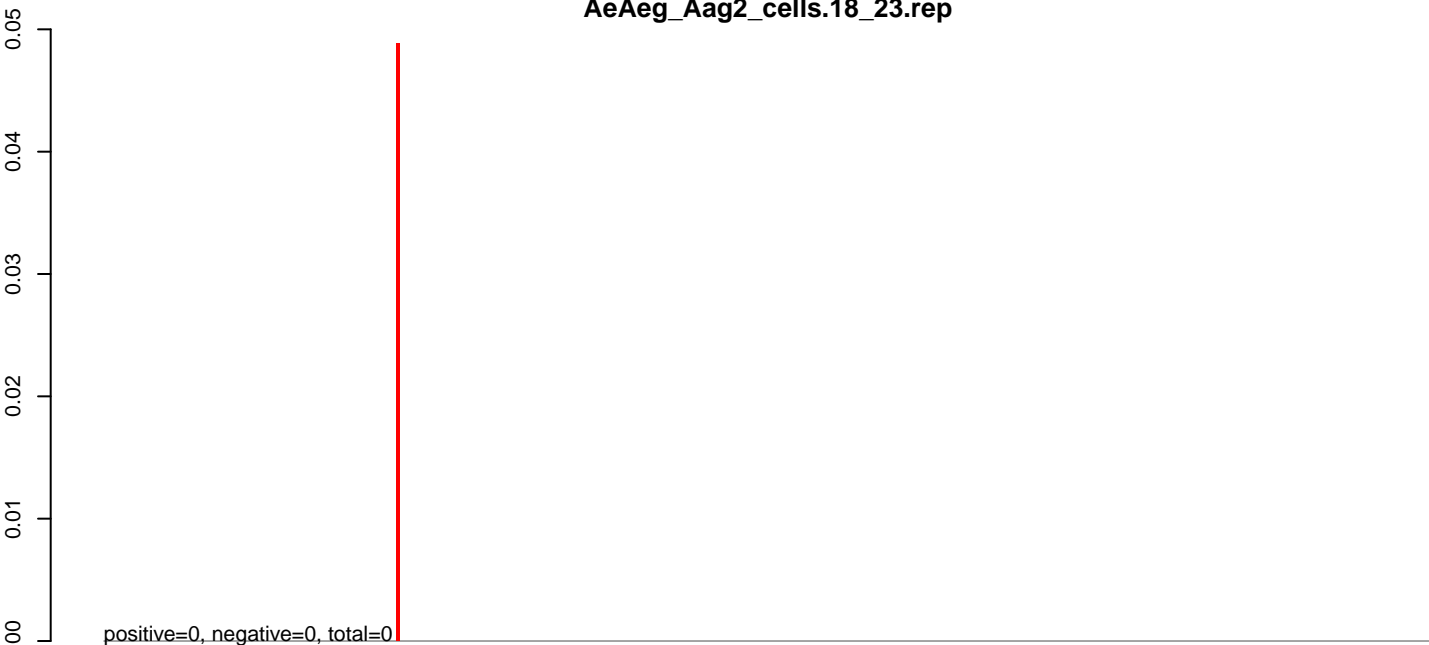


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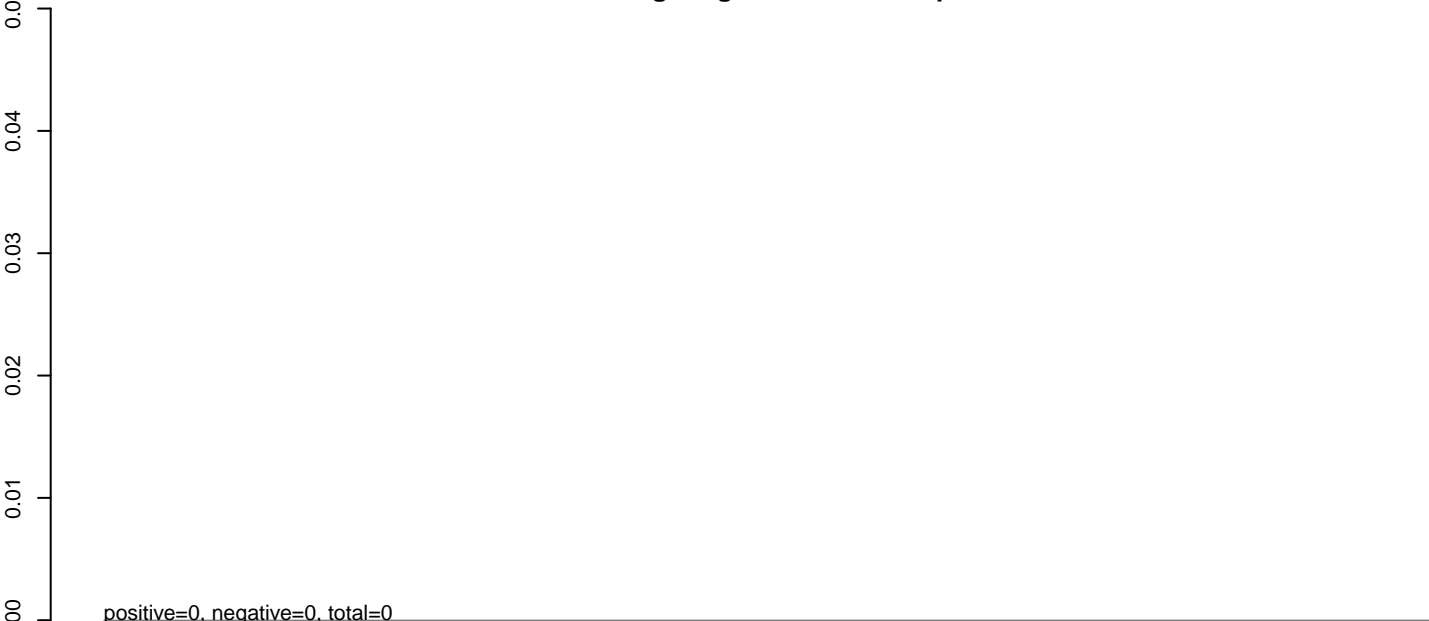


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AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

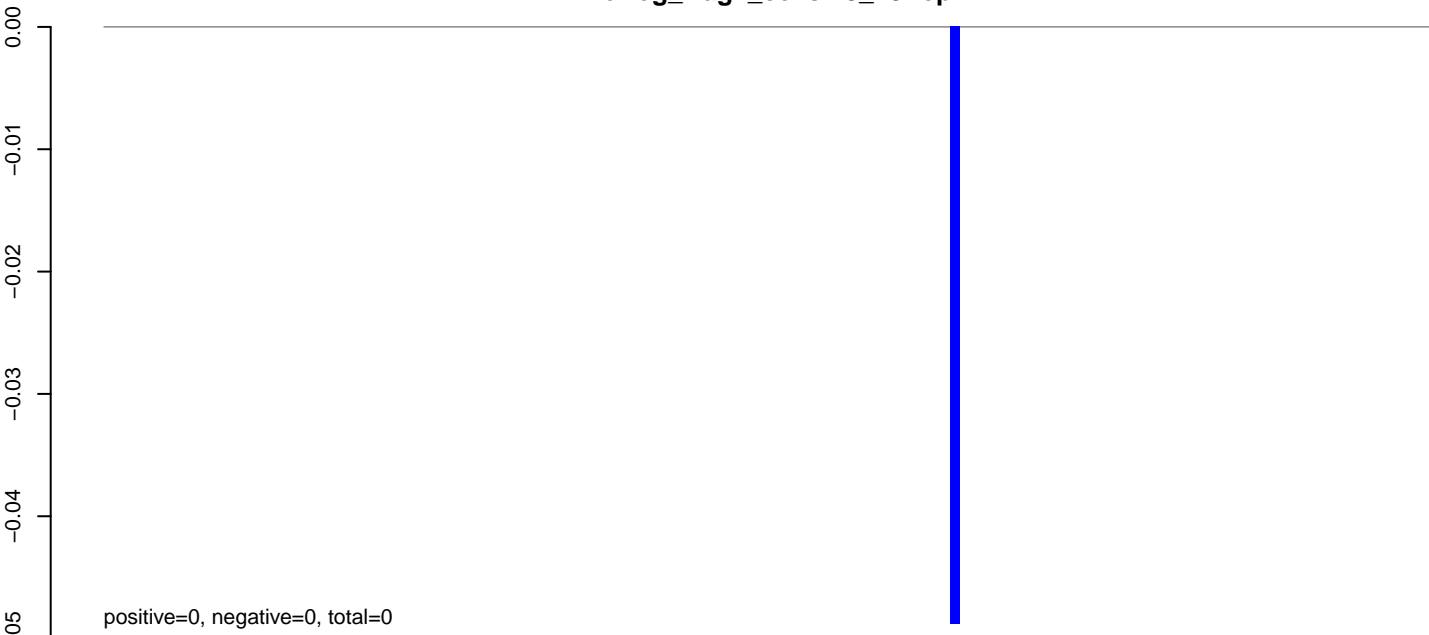


AeAeg_Aag2_cells.rep

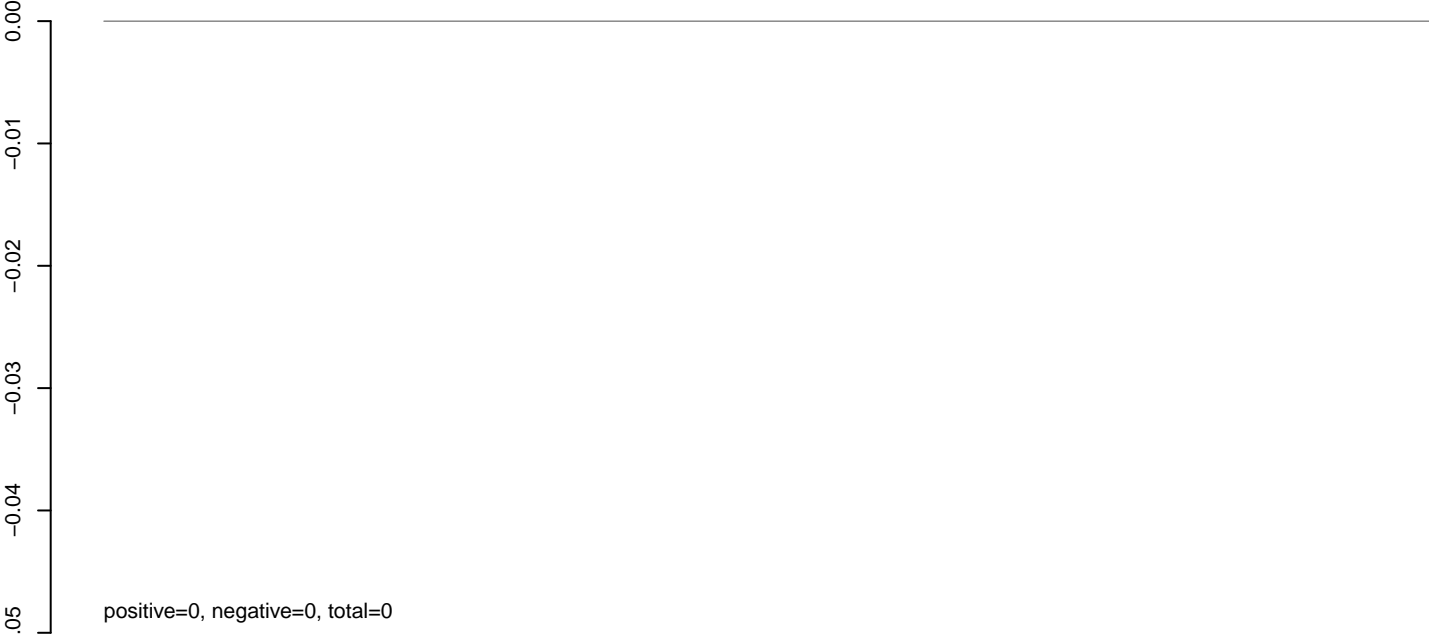


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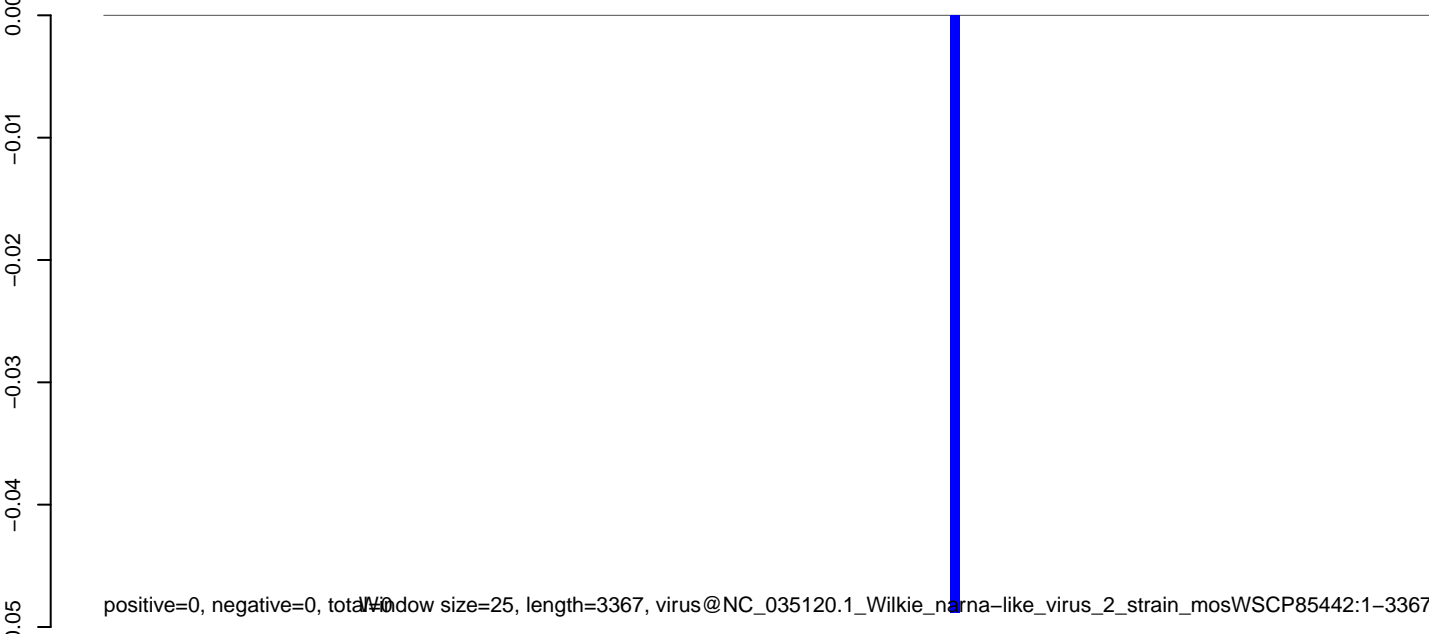
AeAeg_Aag2_cells.18_23.rep



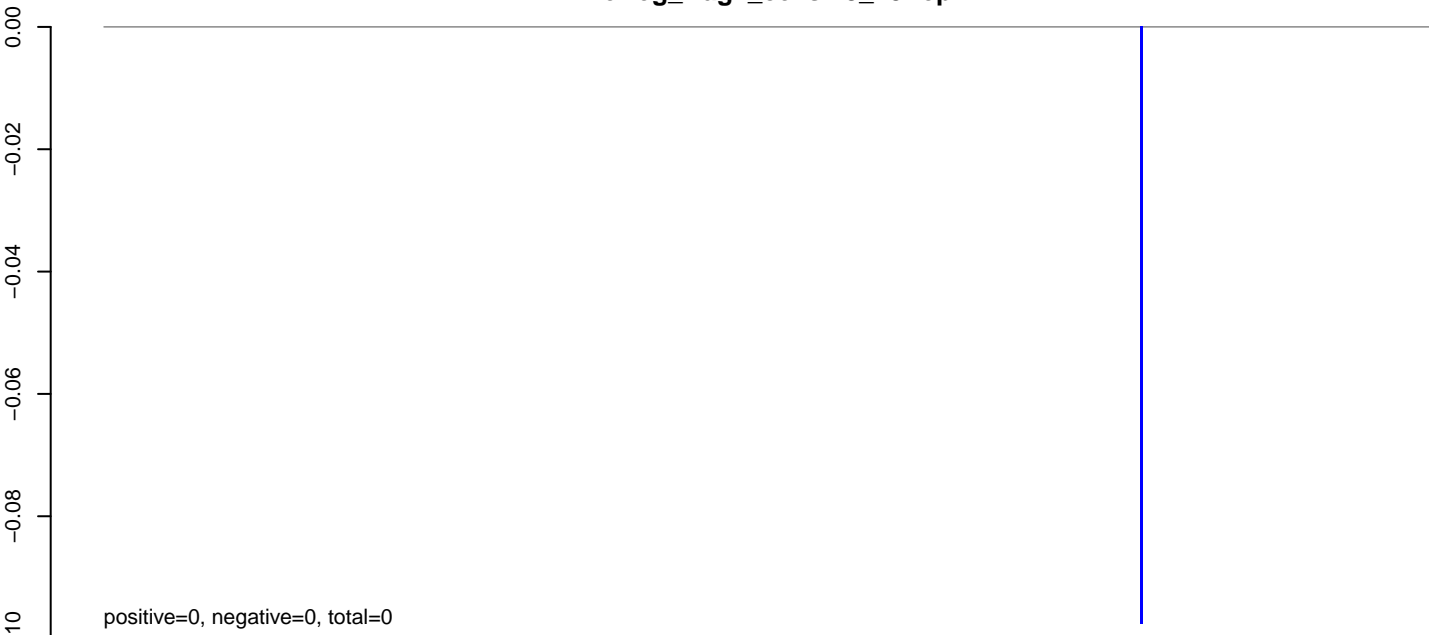
AeAeg_Aag2_cells.24_35.rep



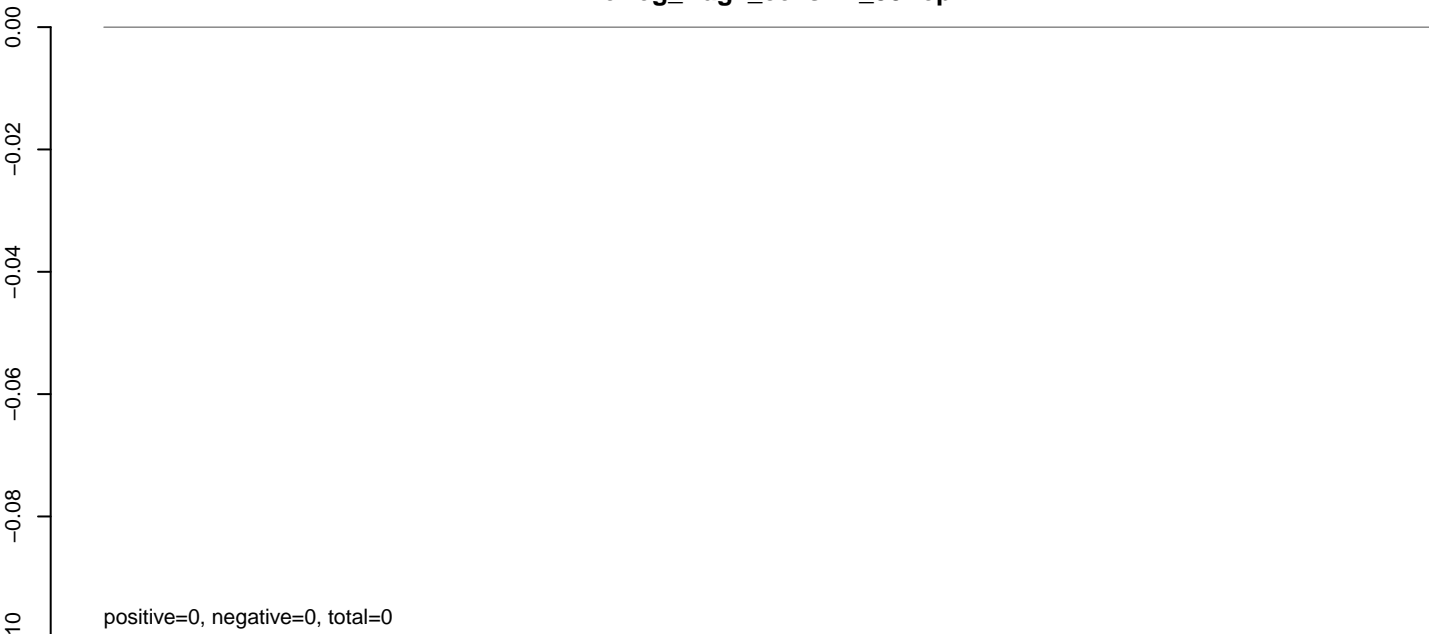
AeAeg_Aag2_cells.rep



AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



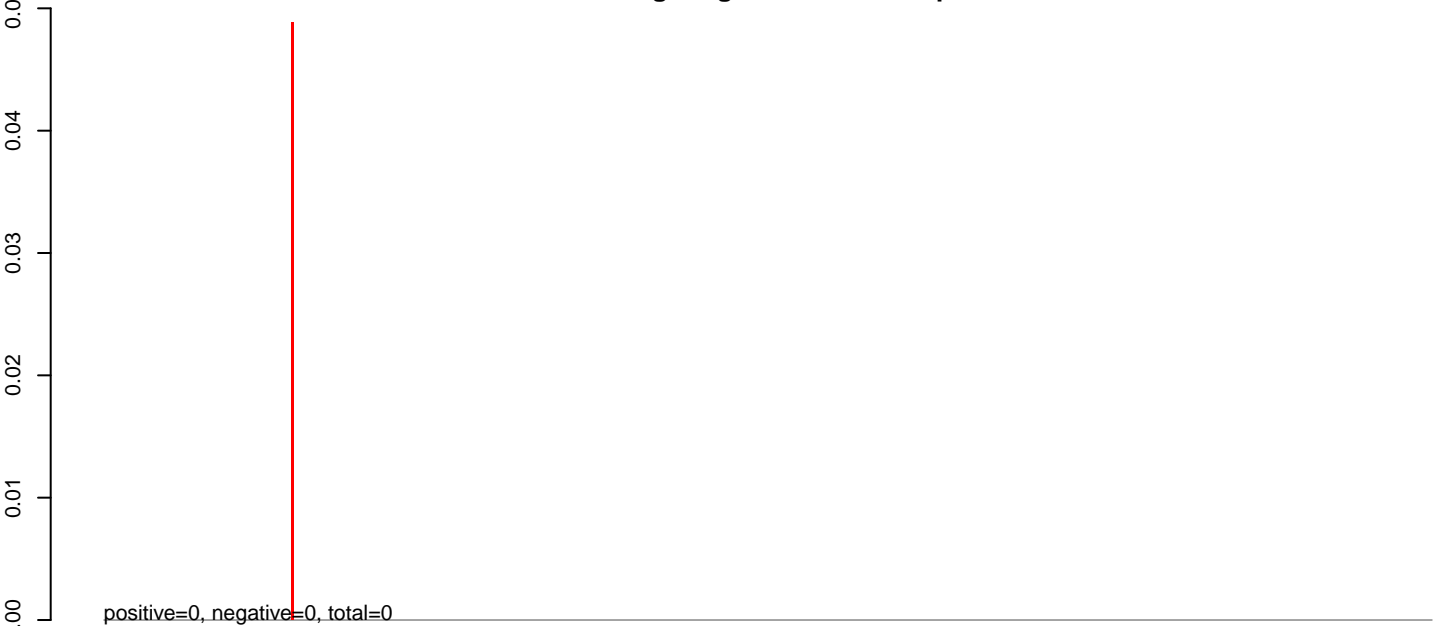
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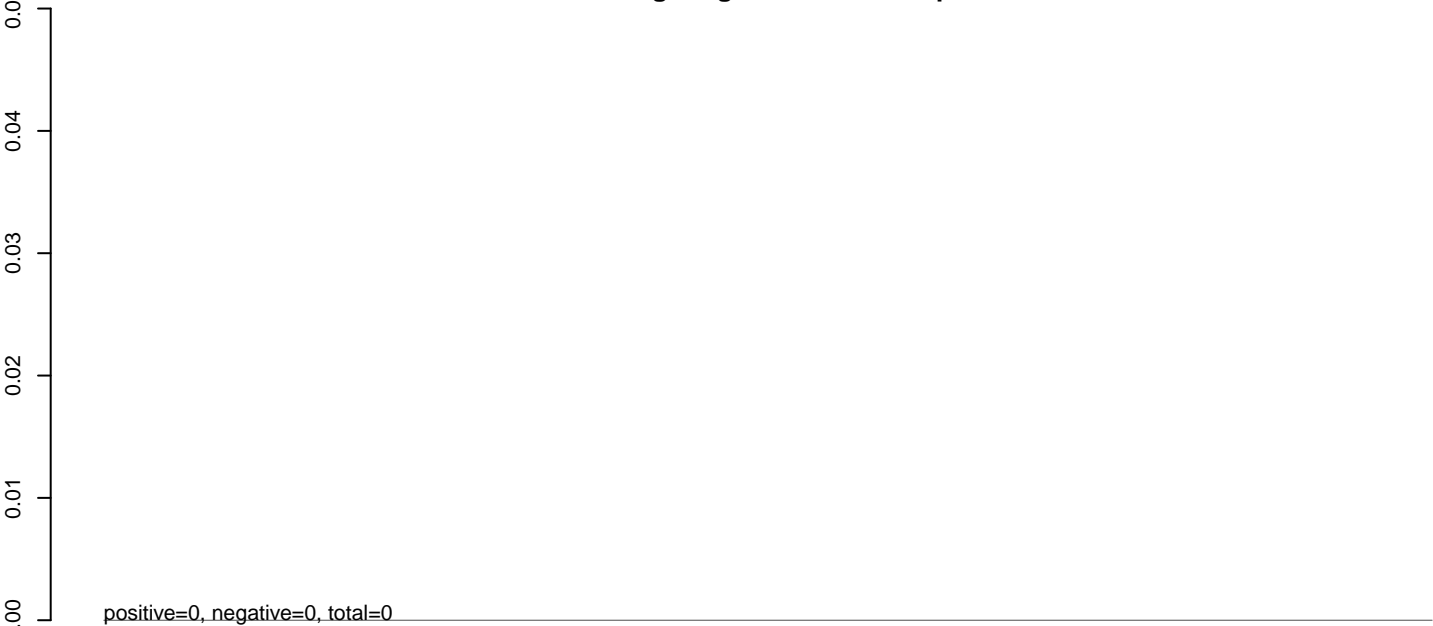
bow size=25, left=13189, right=0, total=0 NC_034543.1-Ord_River_virus_nucleoprotein_hypothetical_protein_phosphoprotein_hypothetical_pro:1-13189

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_cells.18_23.rep

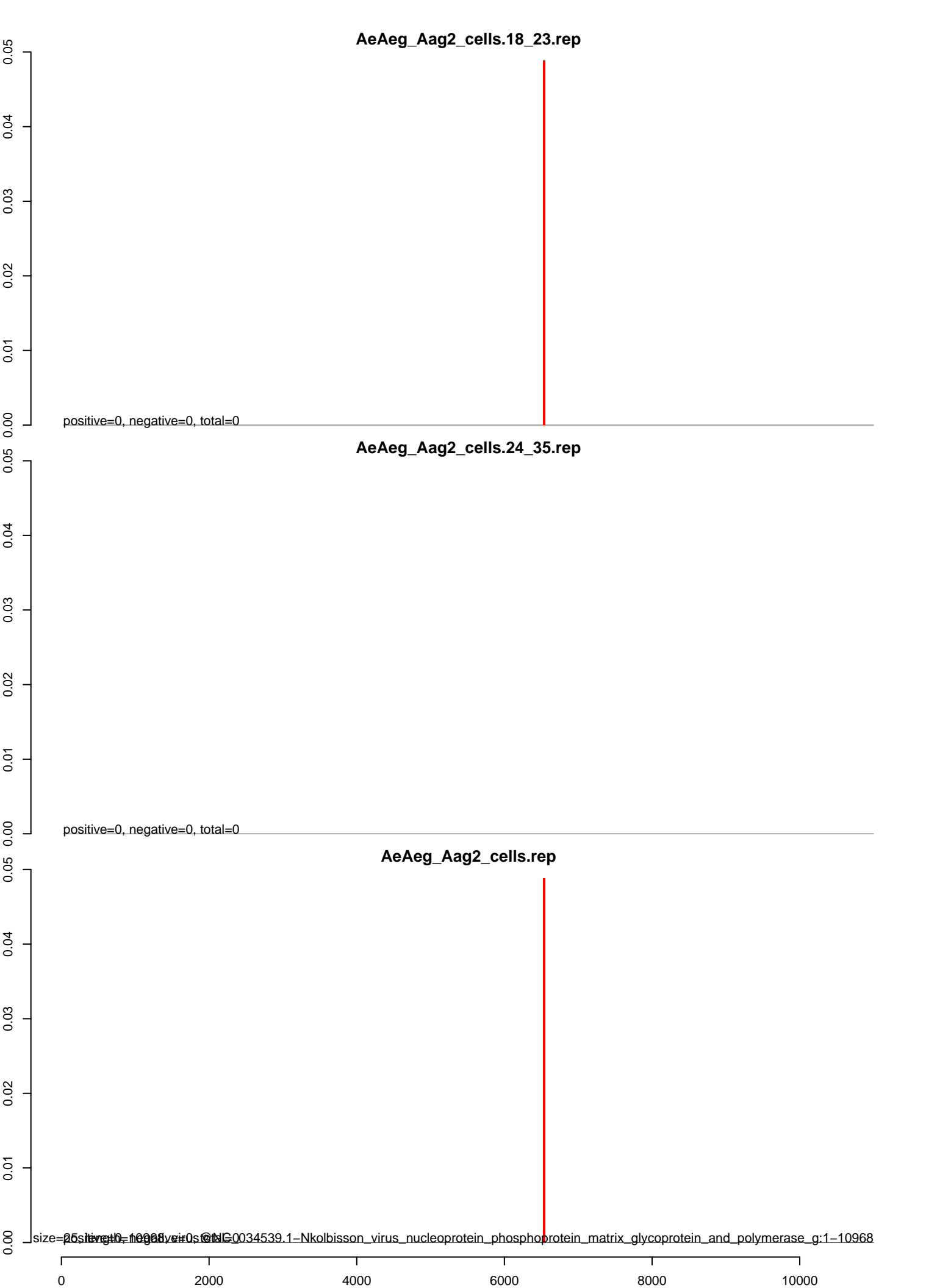


AeAeg_Aag2_cells.24_35.rep

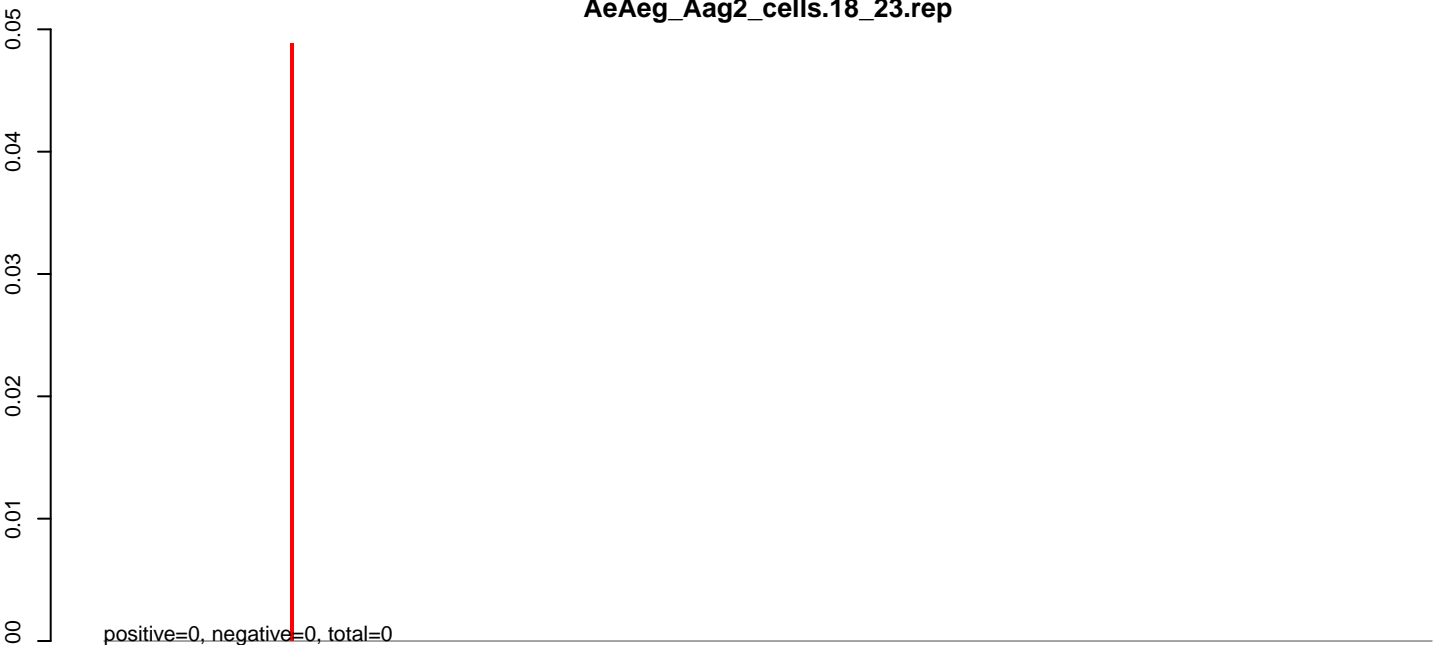


AeAeg_Aag2_cells.rep

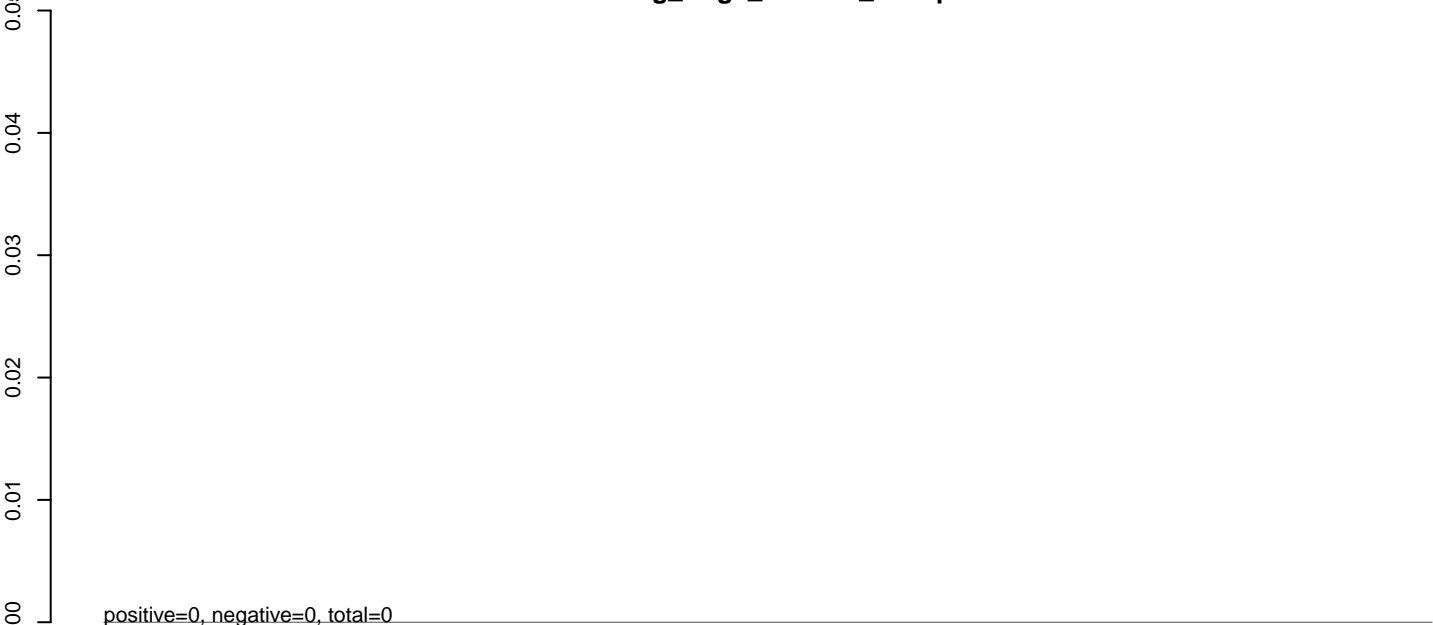




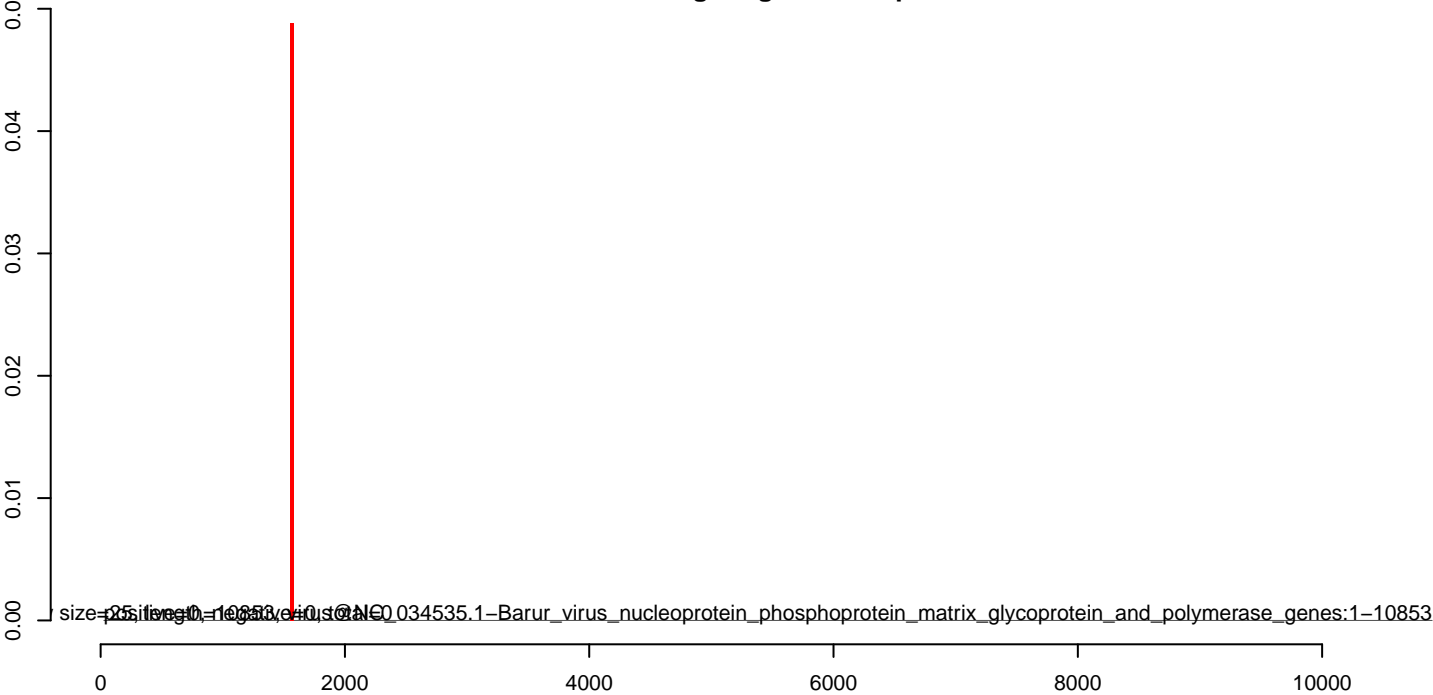
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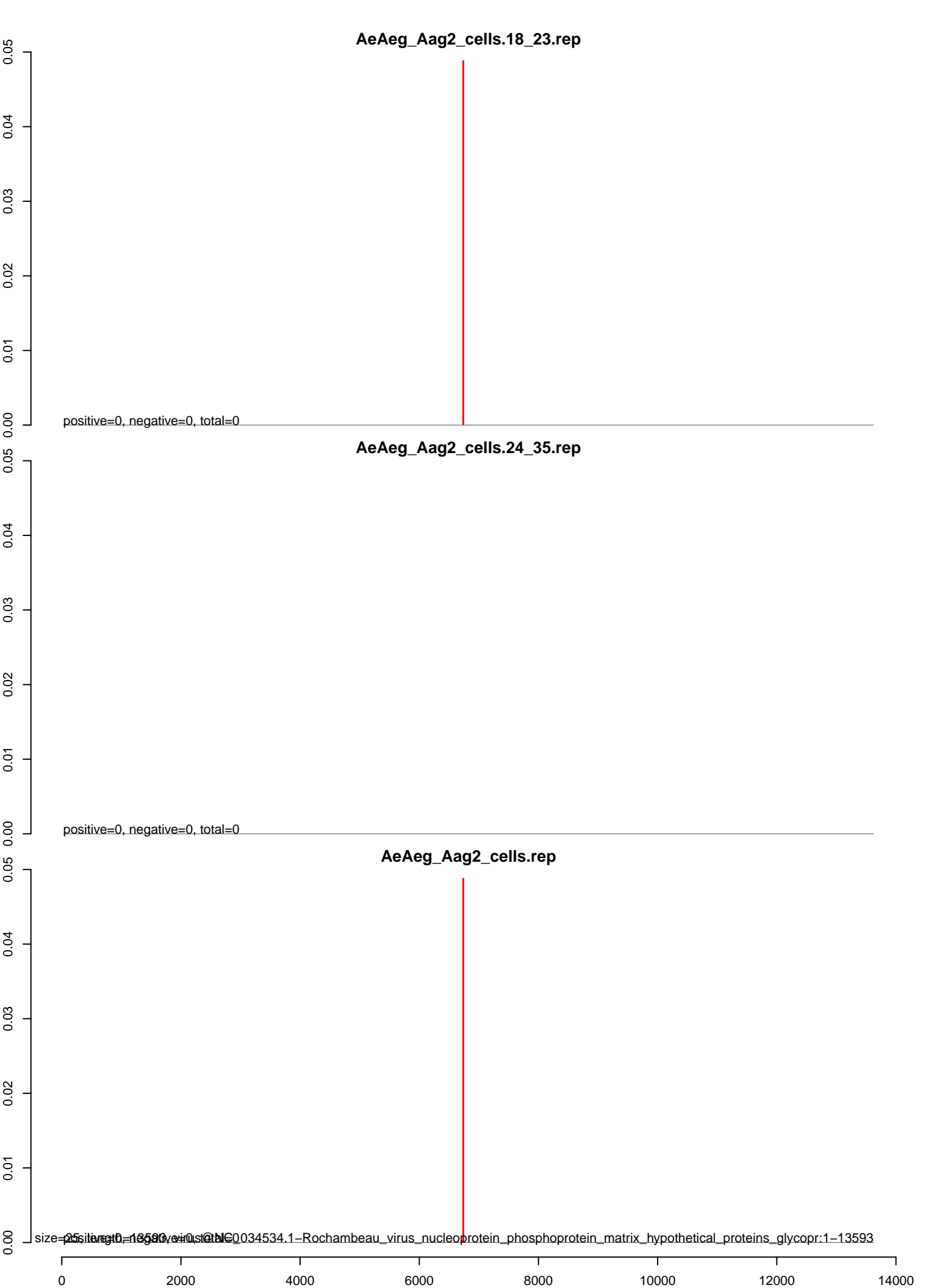


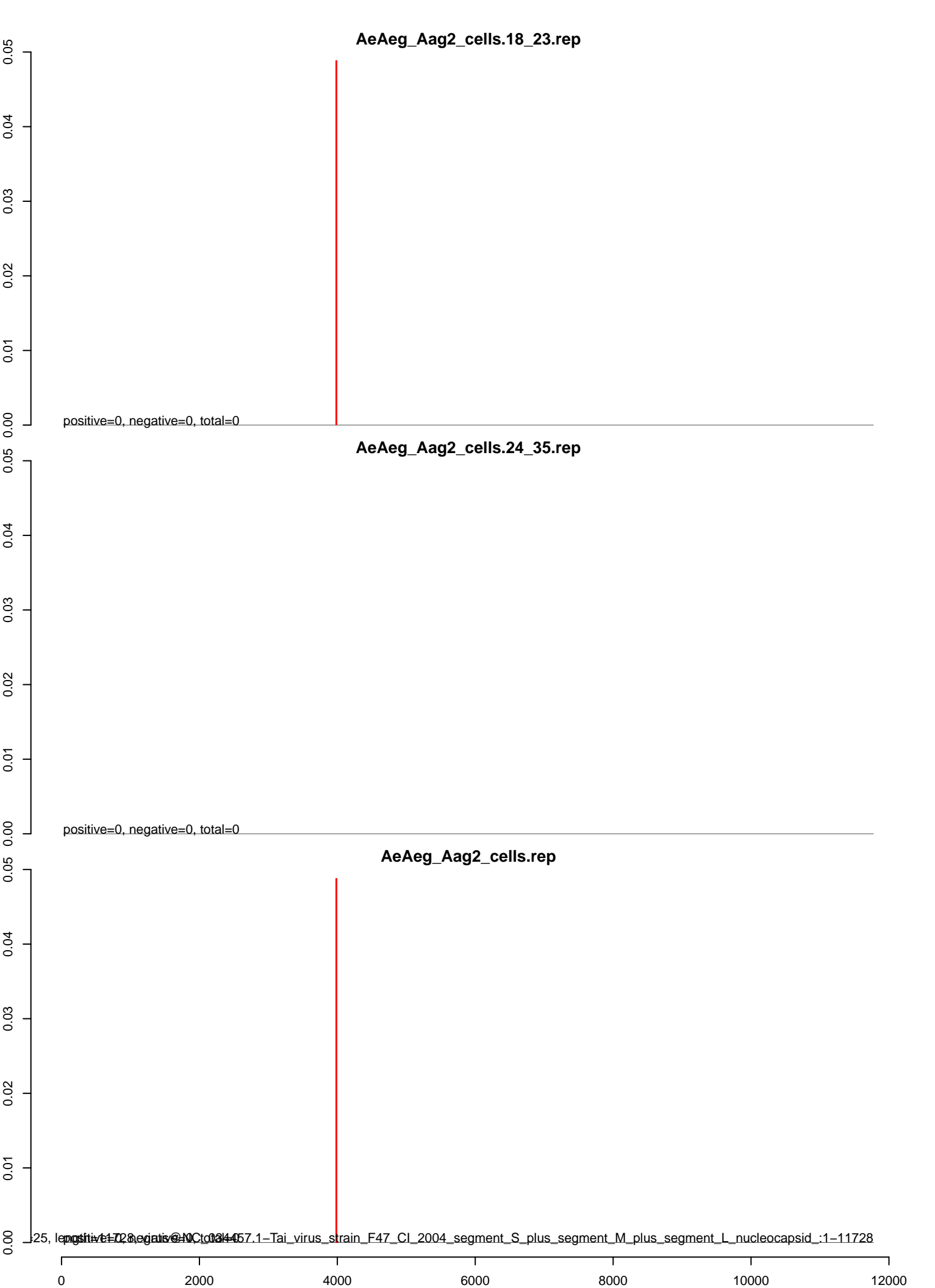
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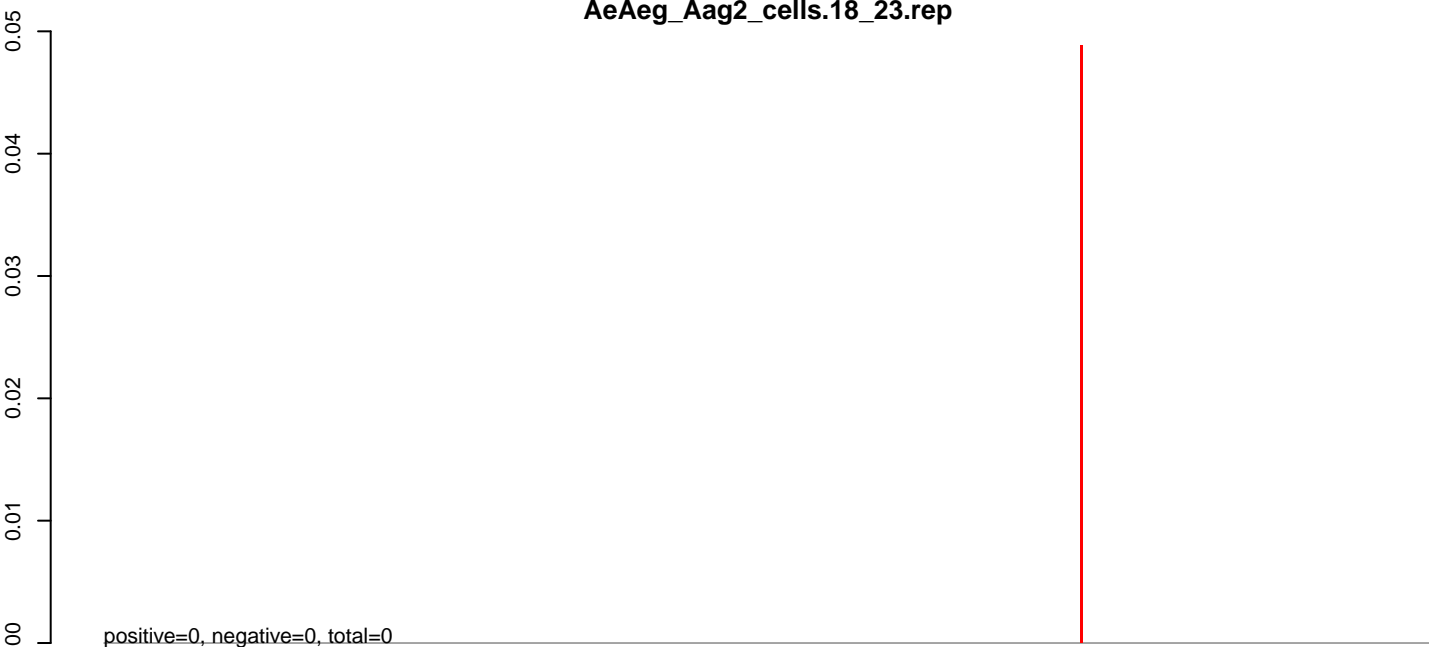
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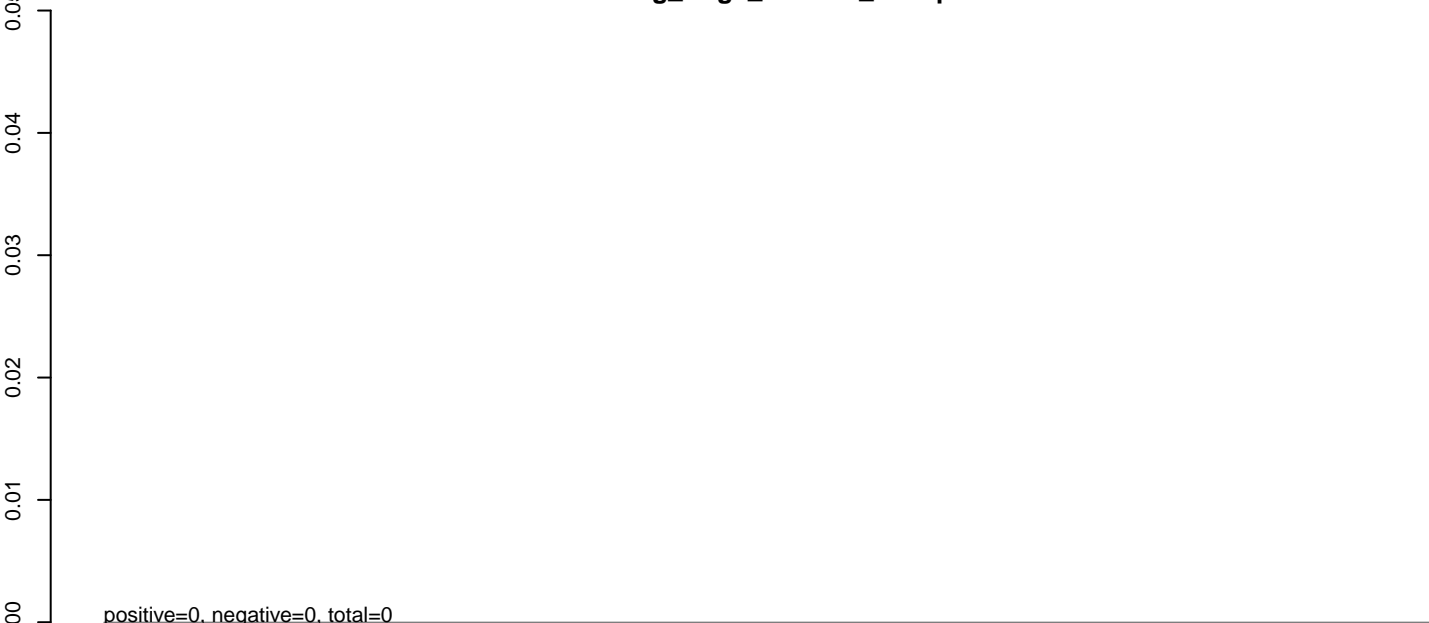




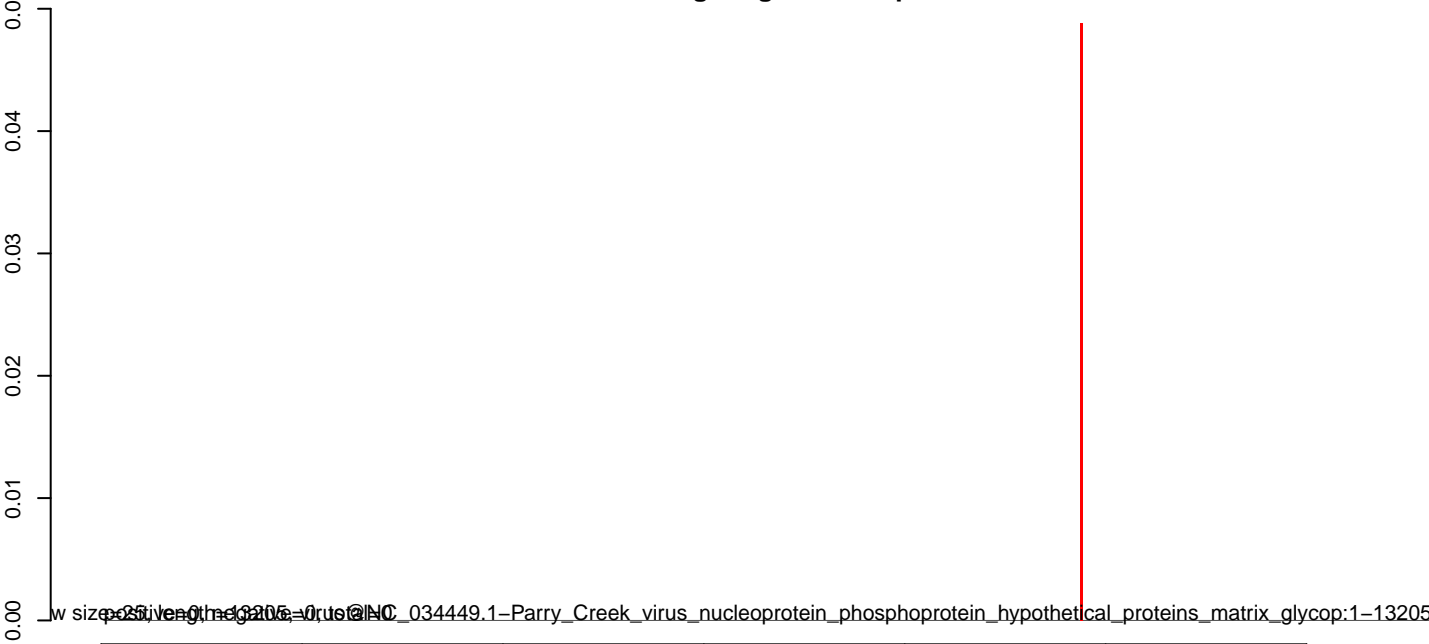
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

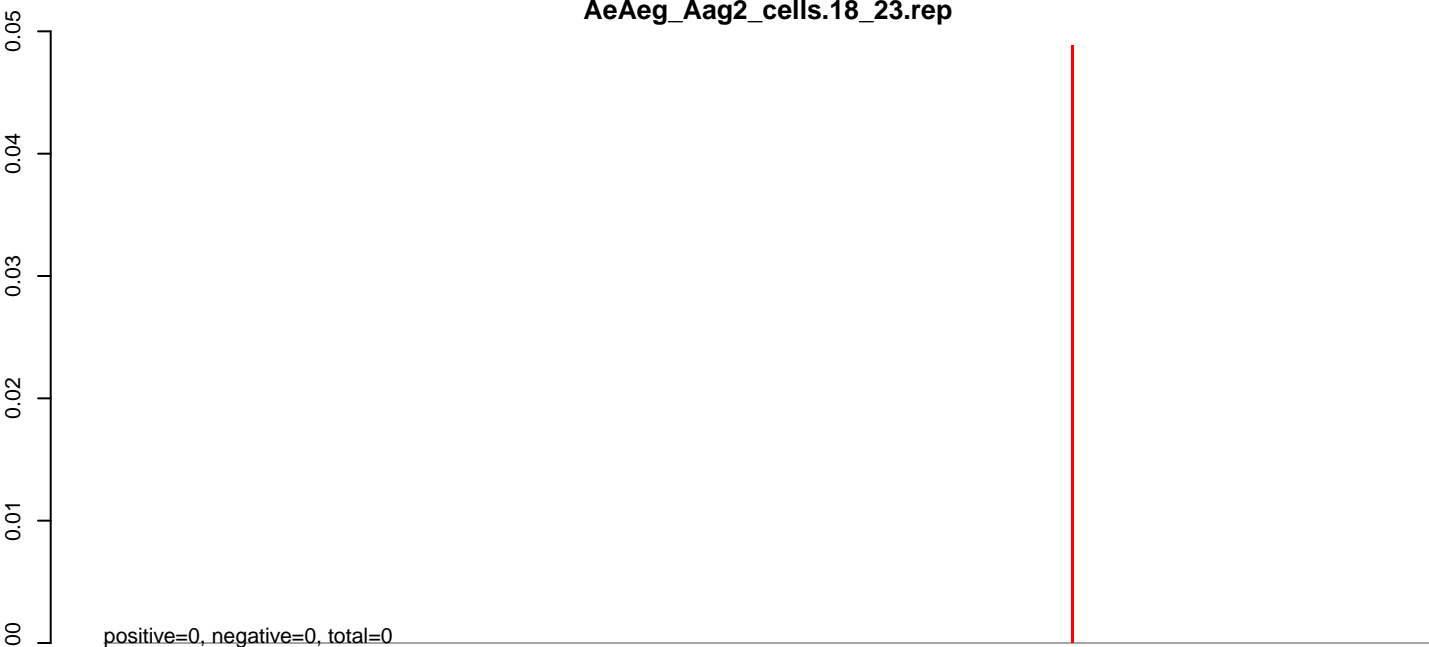


AeAeg_Aag2_cells.rep

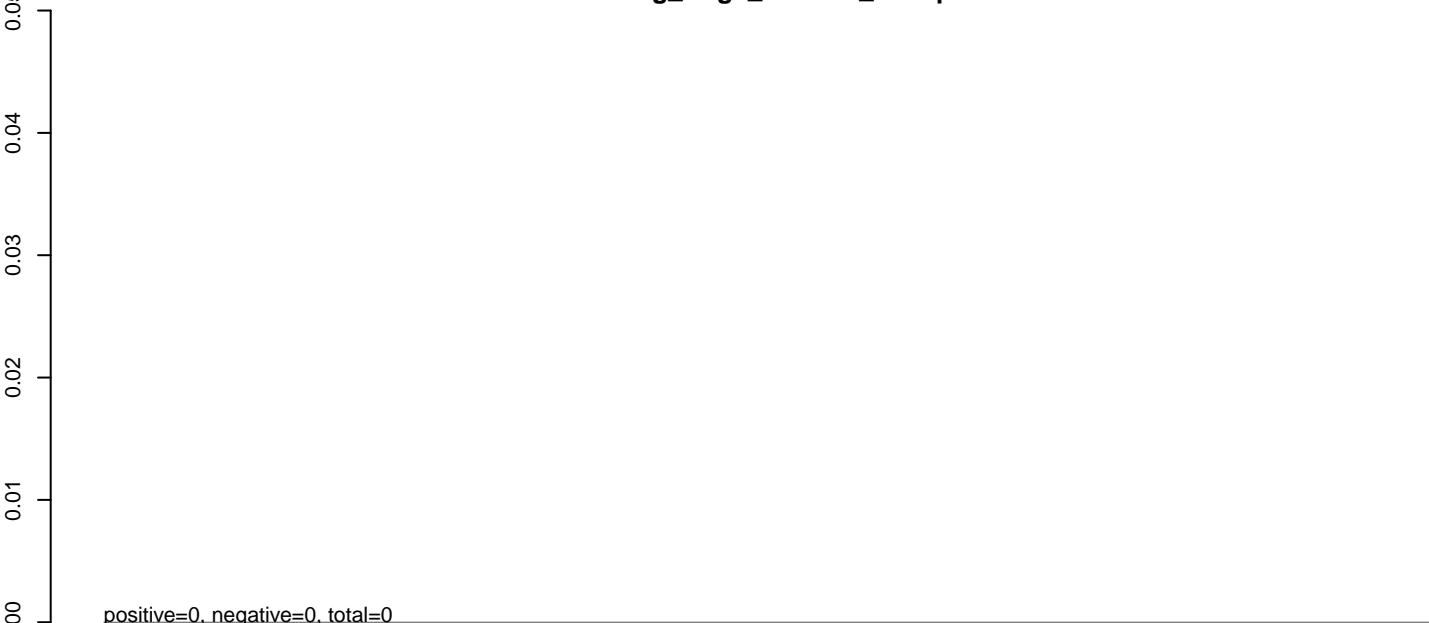


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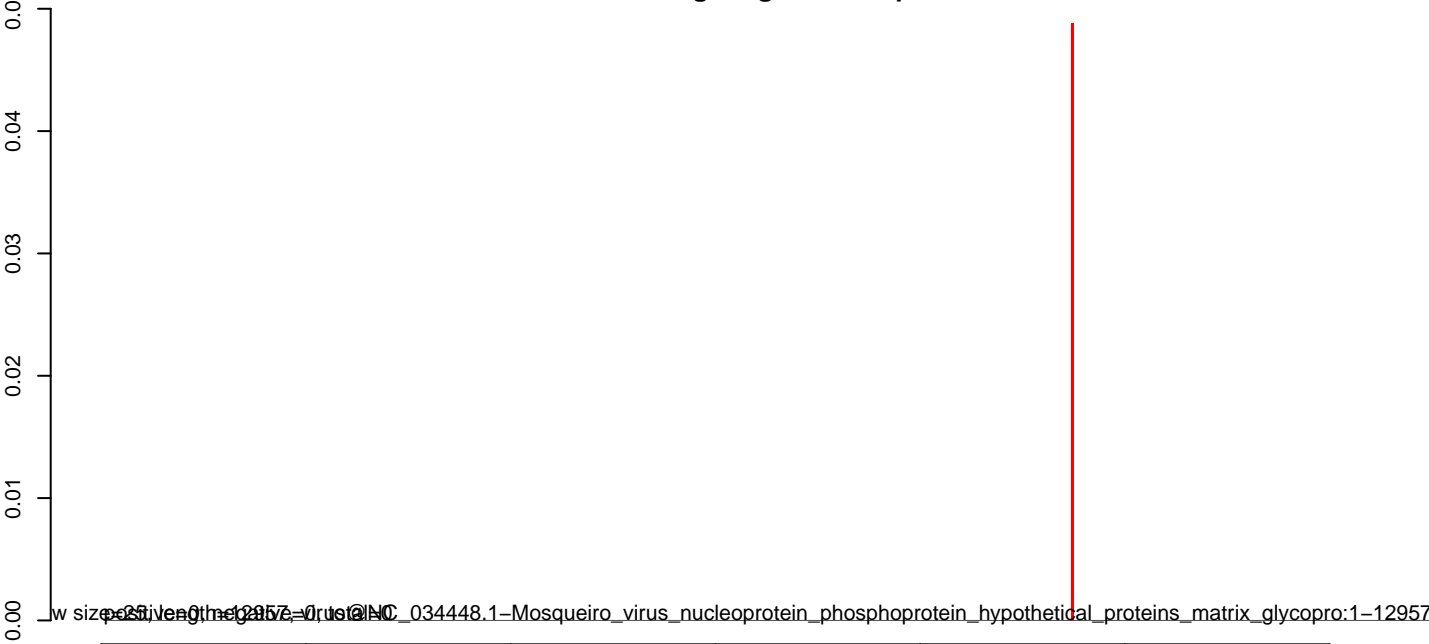
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

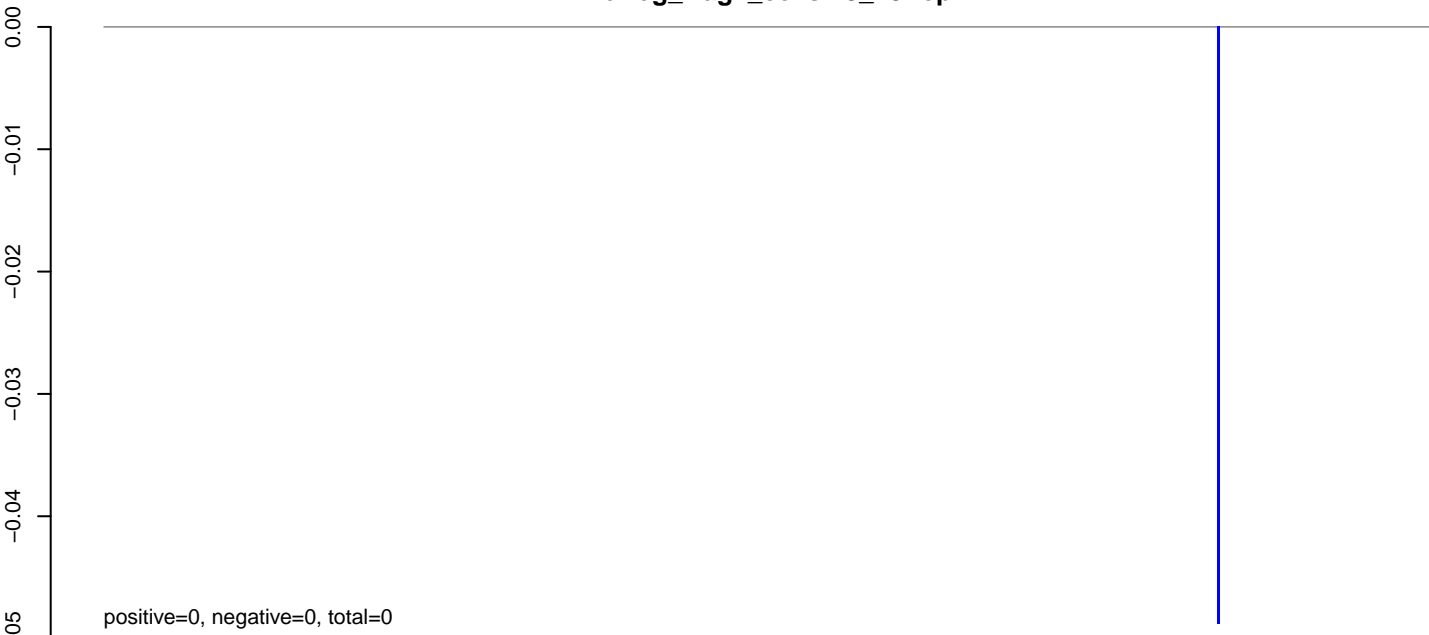


AeAeg_Aag2_cells.rep

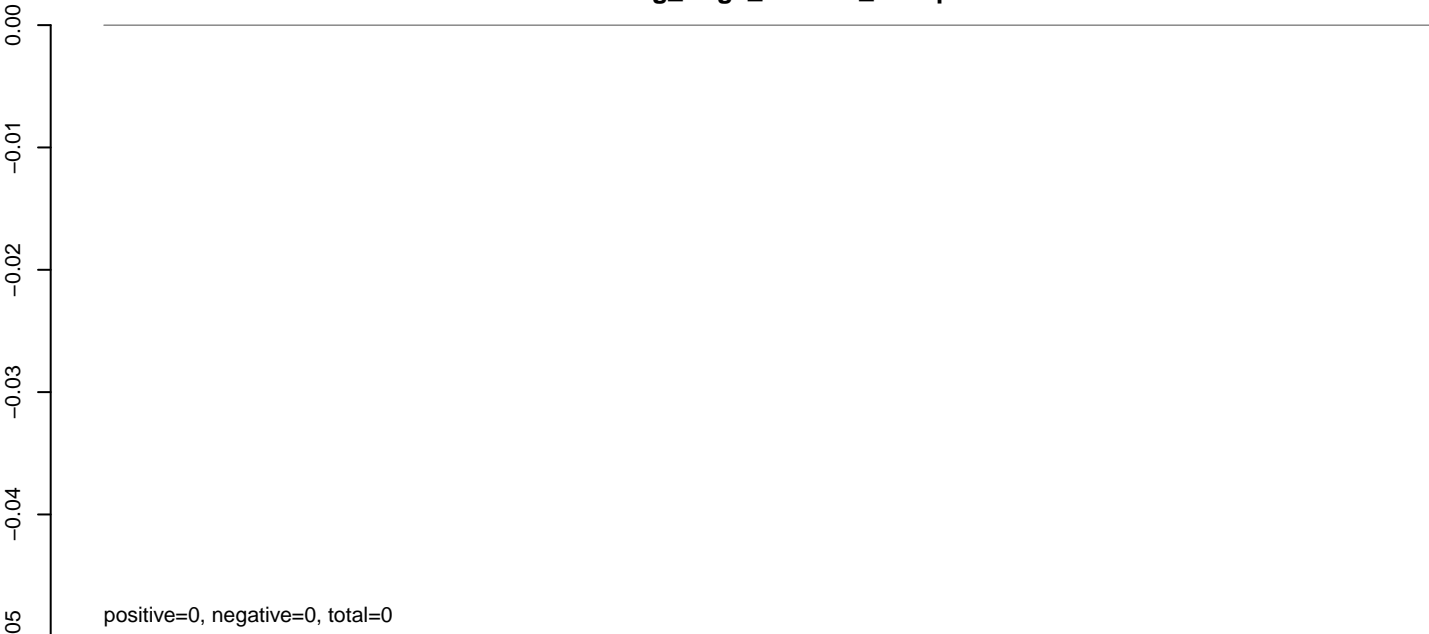


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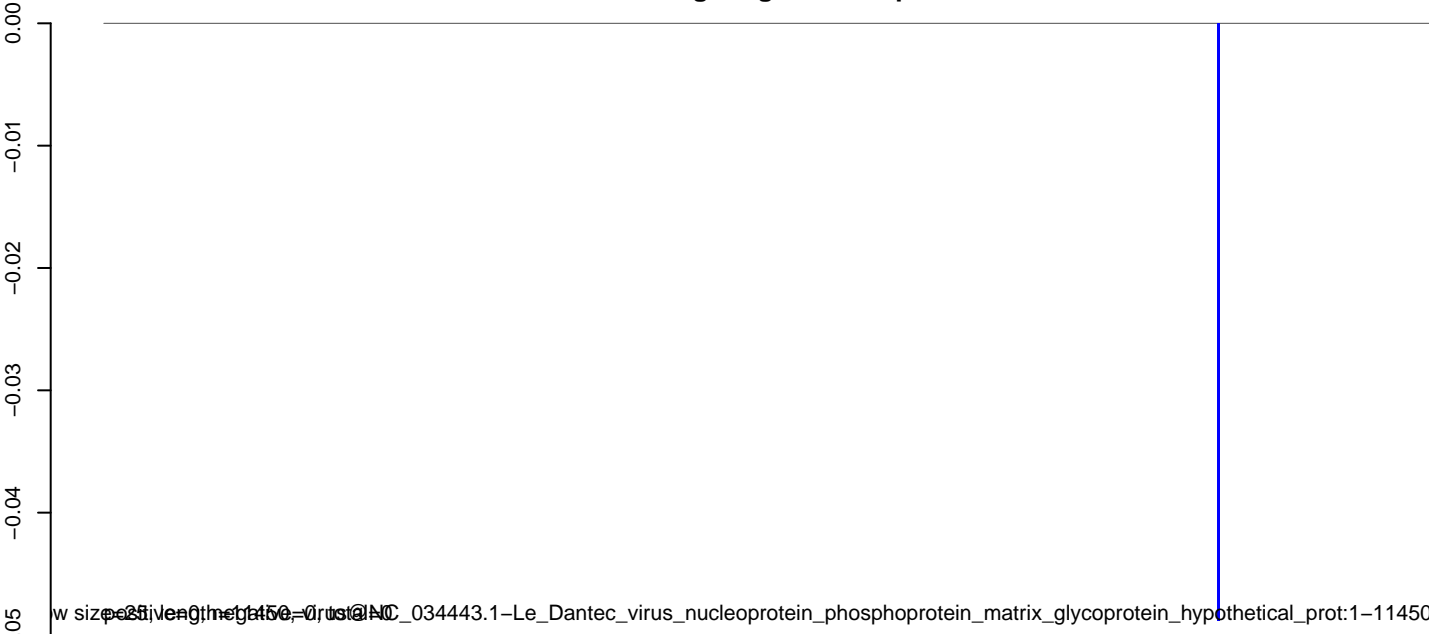
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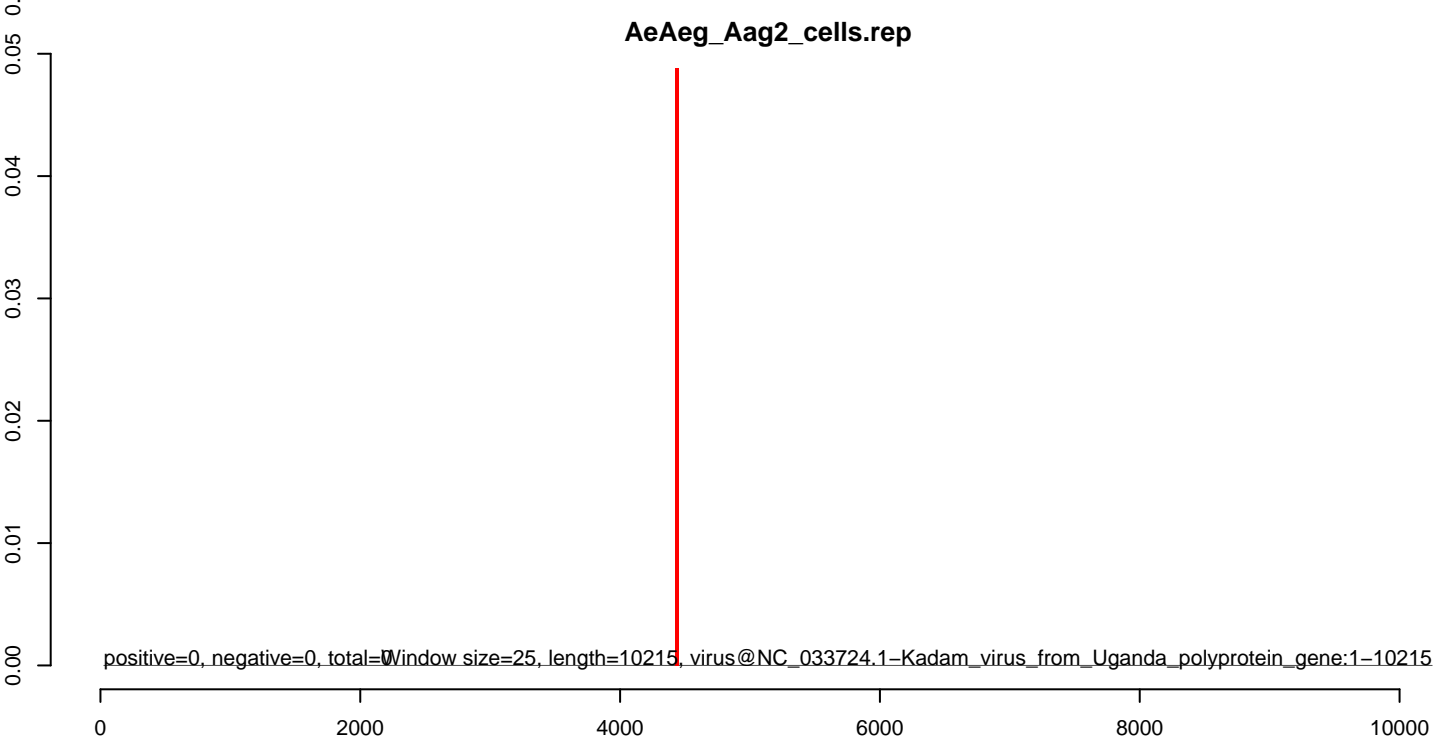
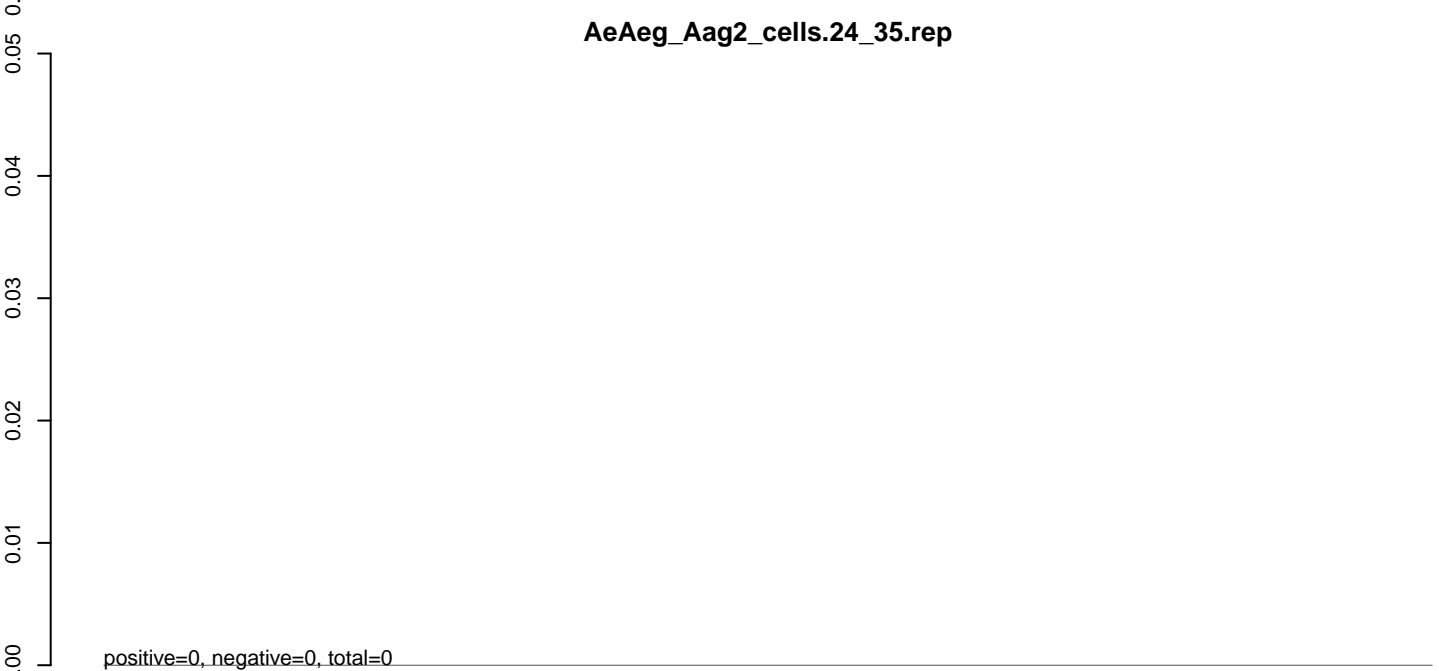
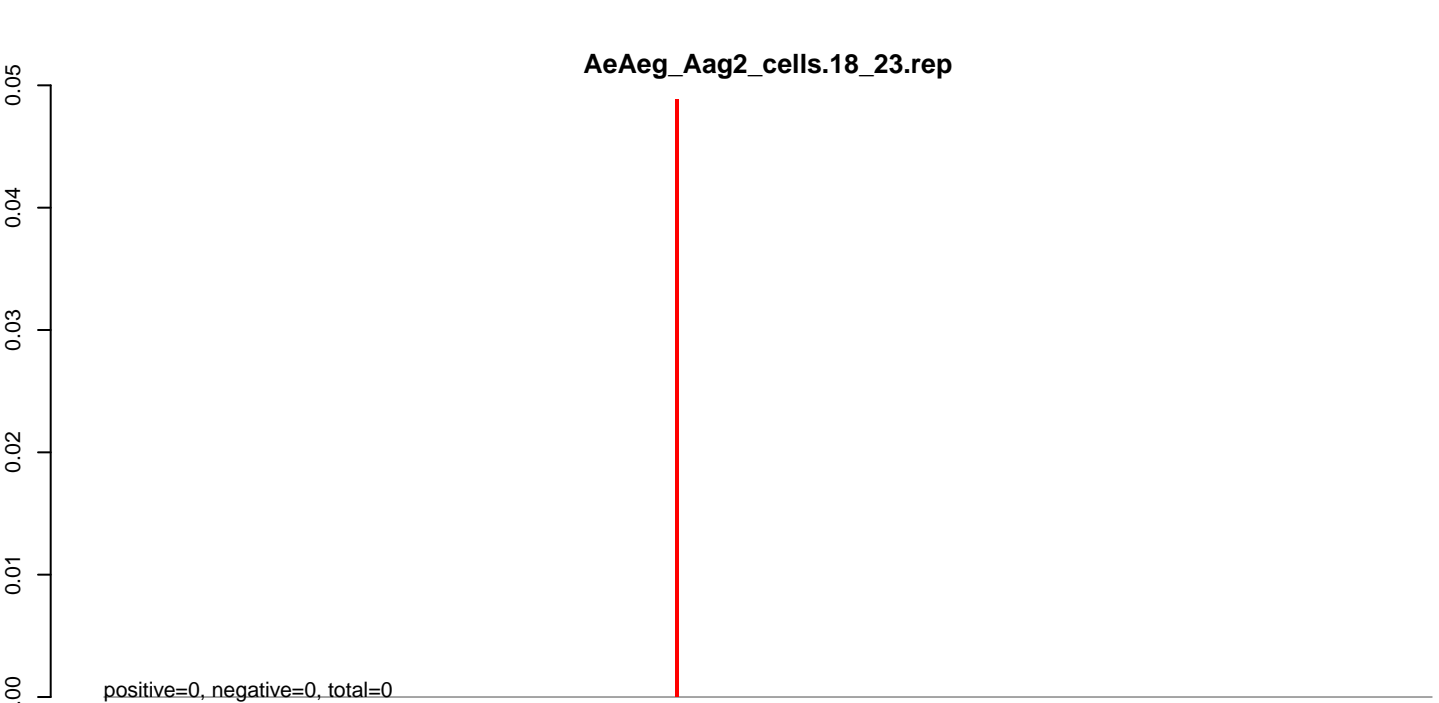
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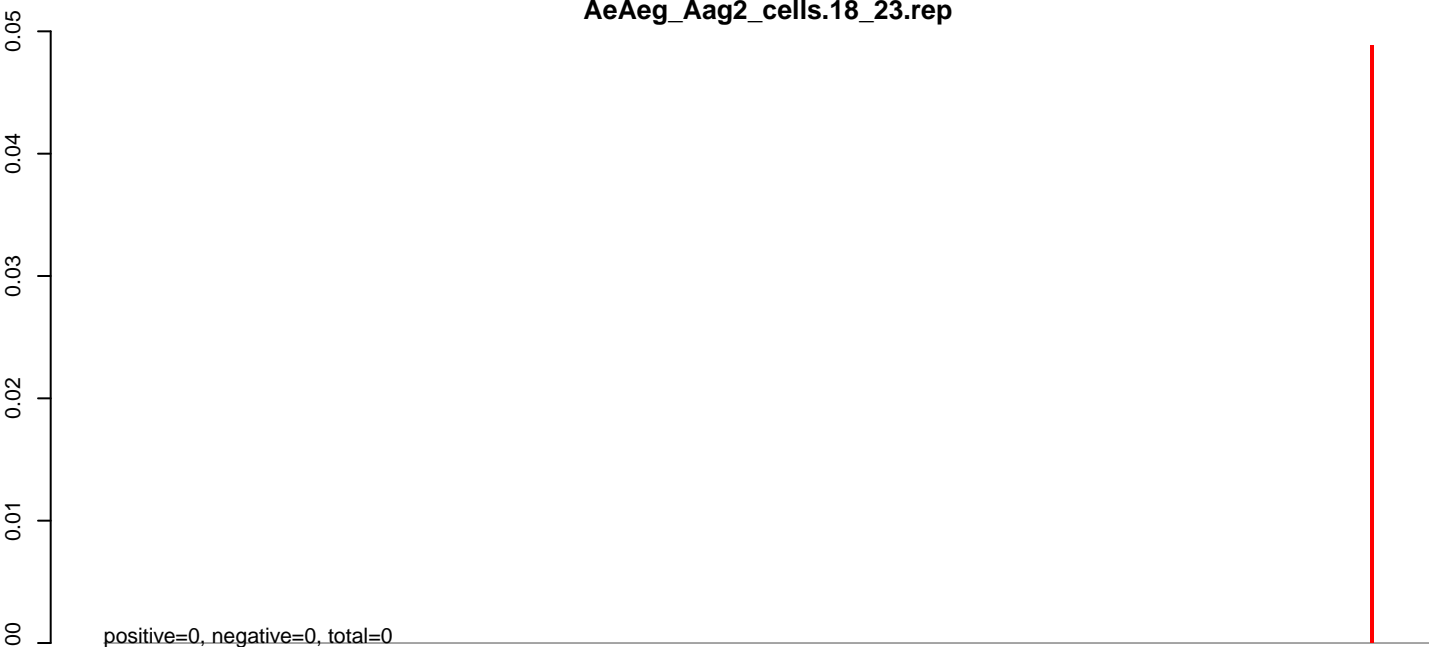
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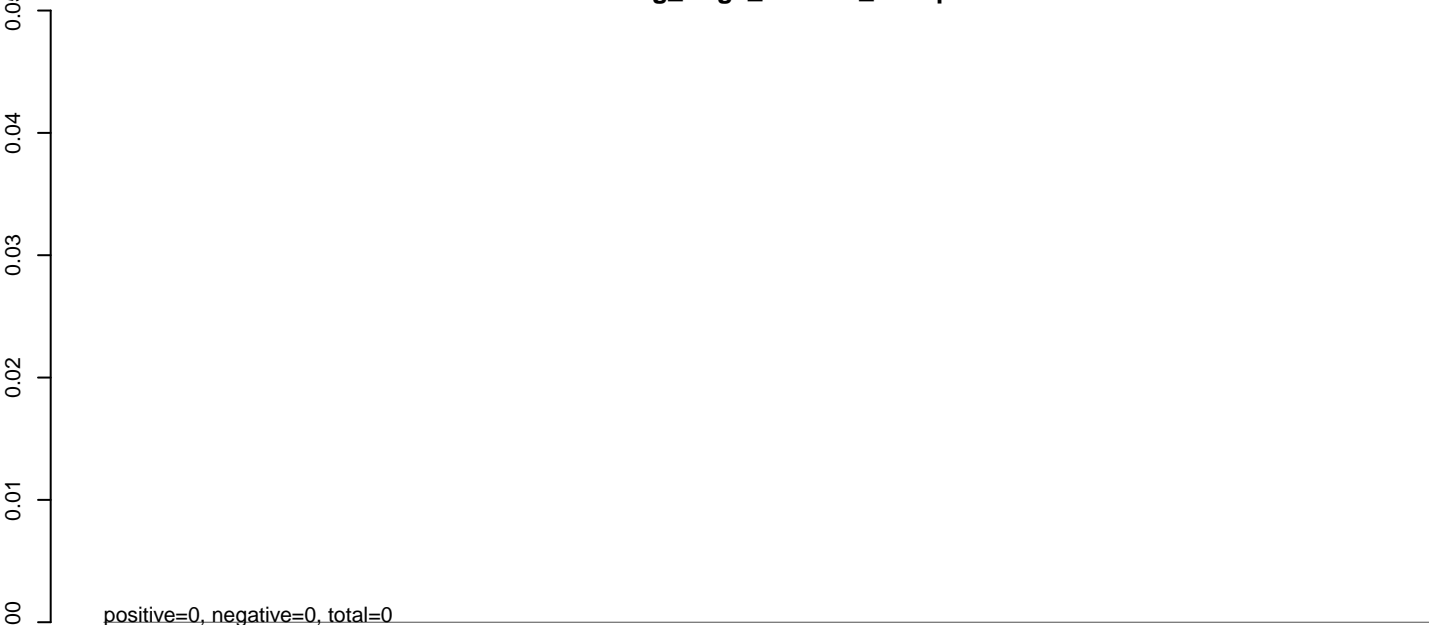
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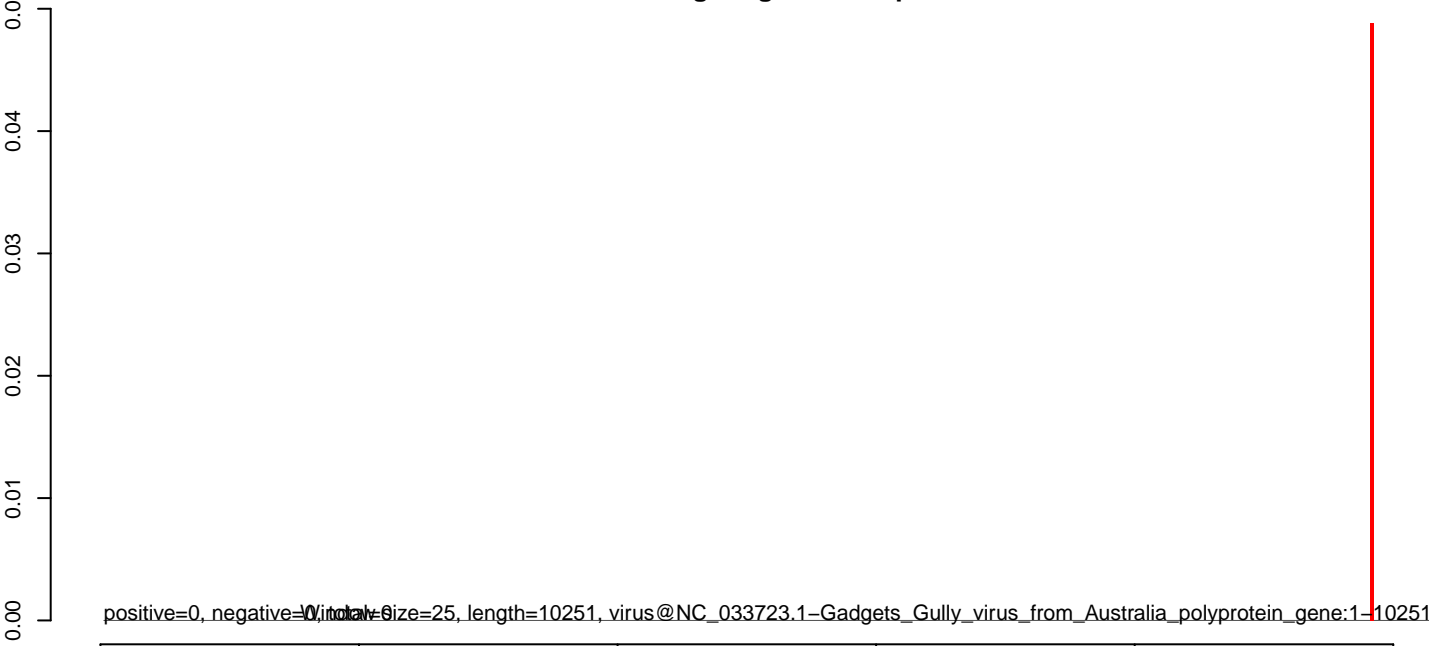
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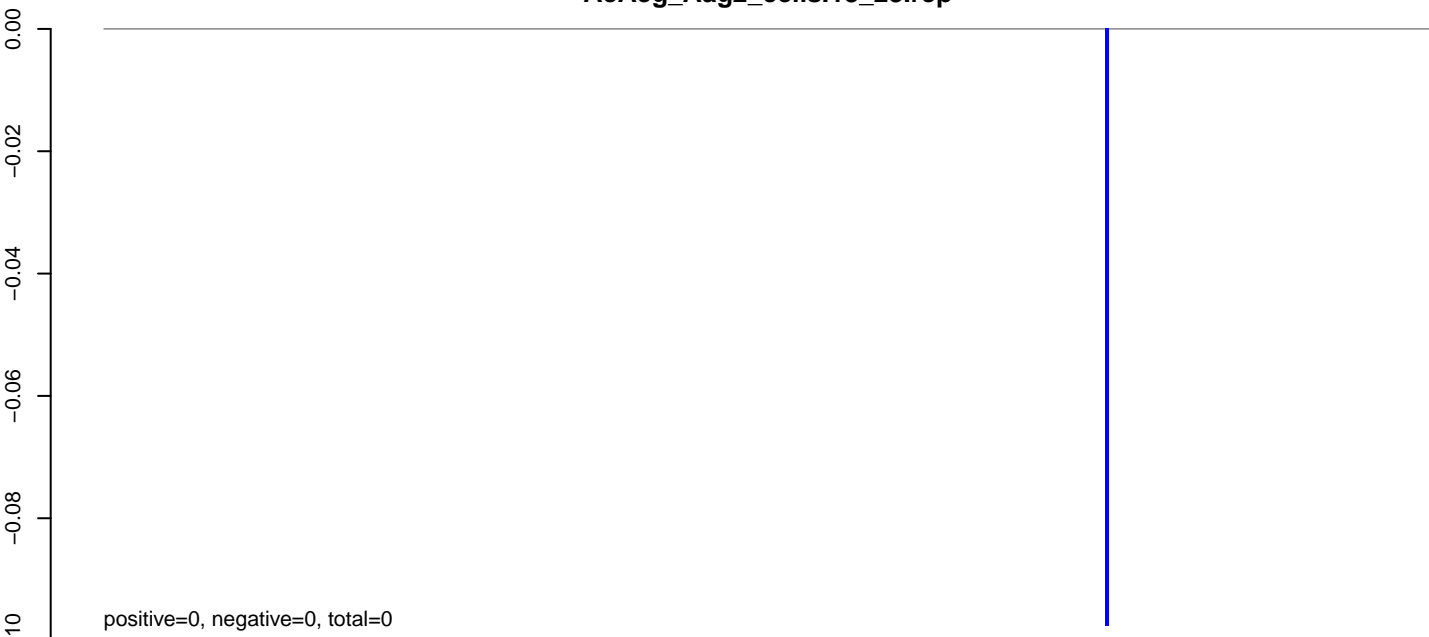
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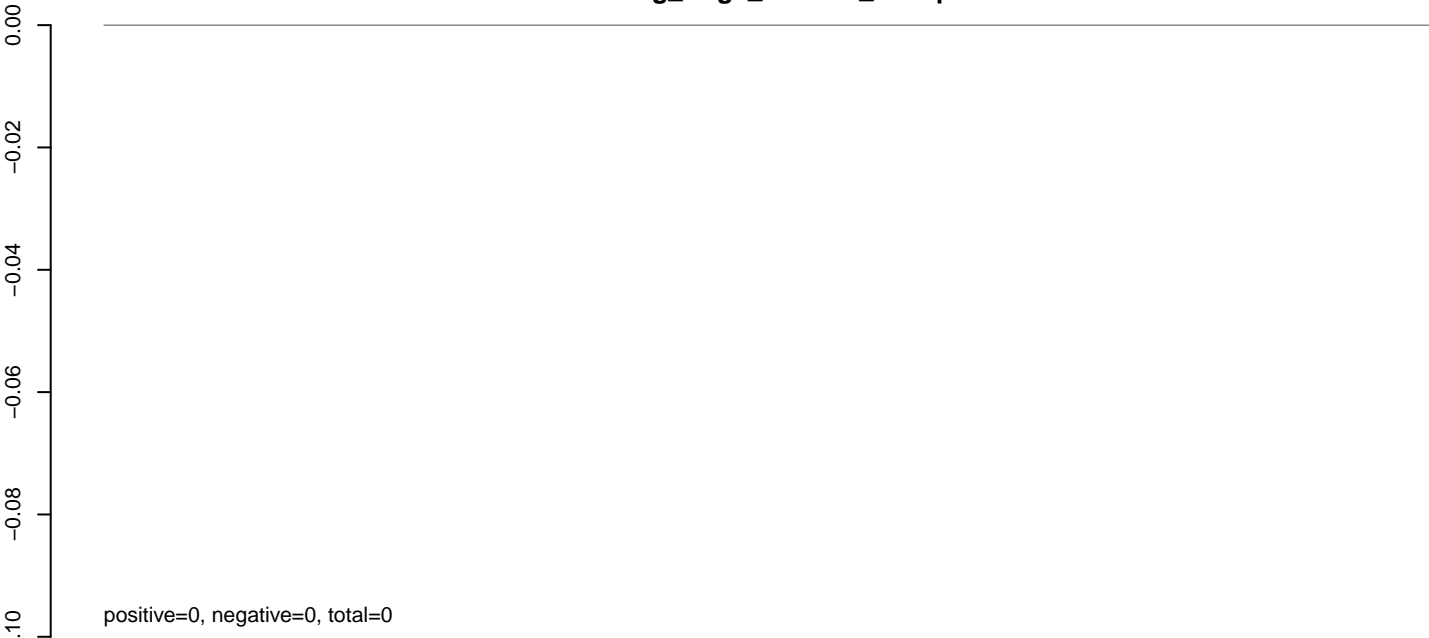
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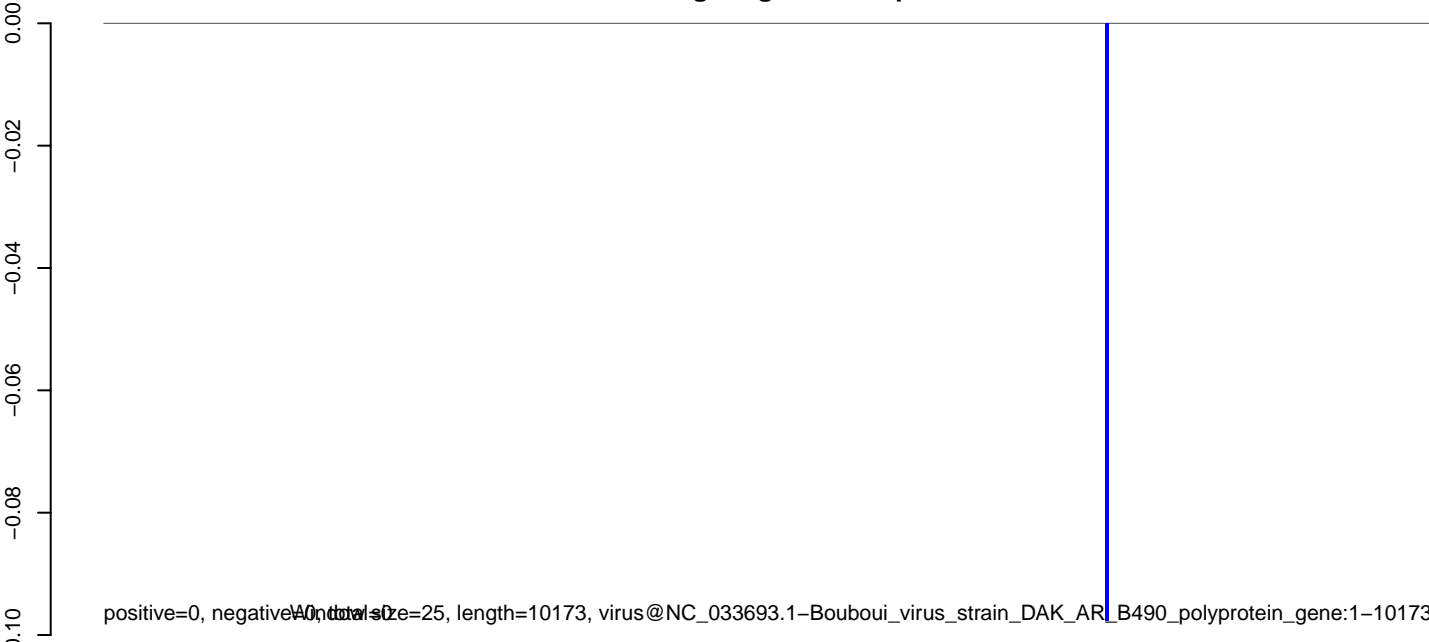
AeAeg_Aag2_cells.18_23.rep



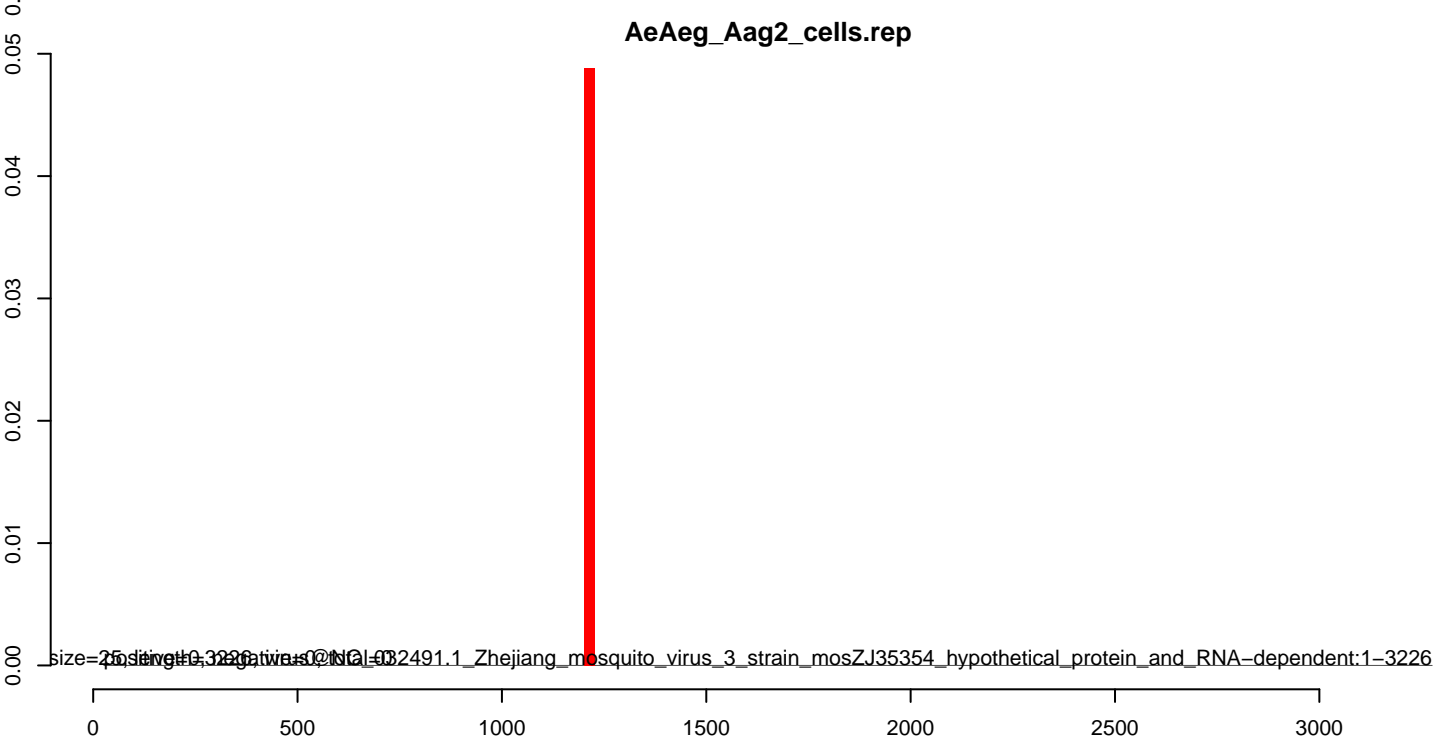
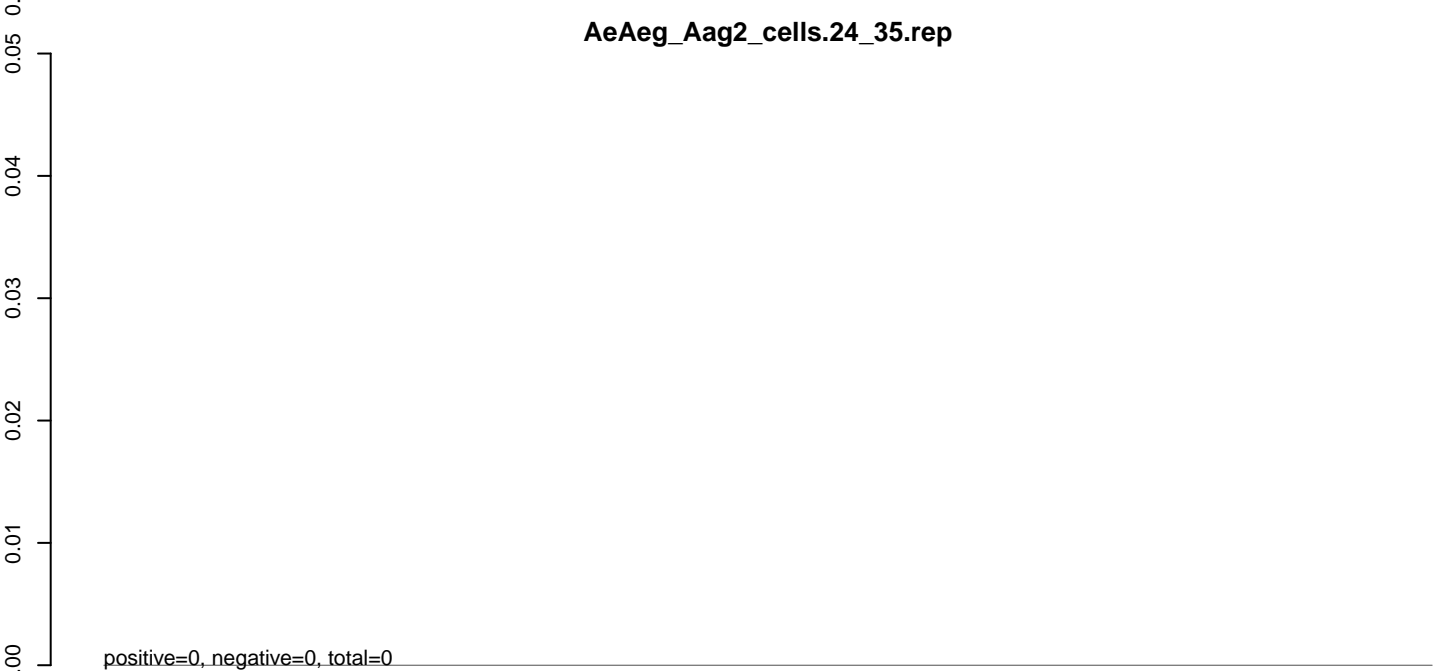
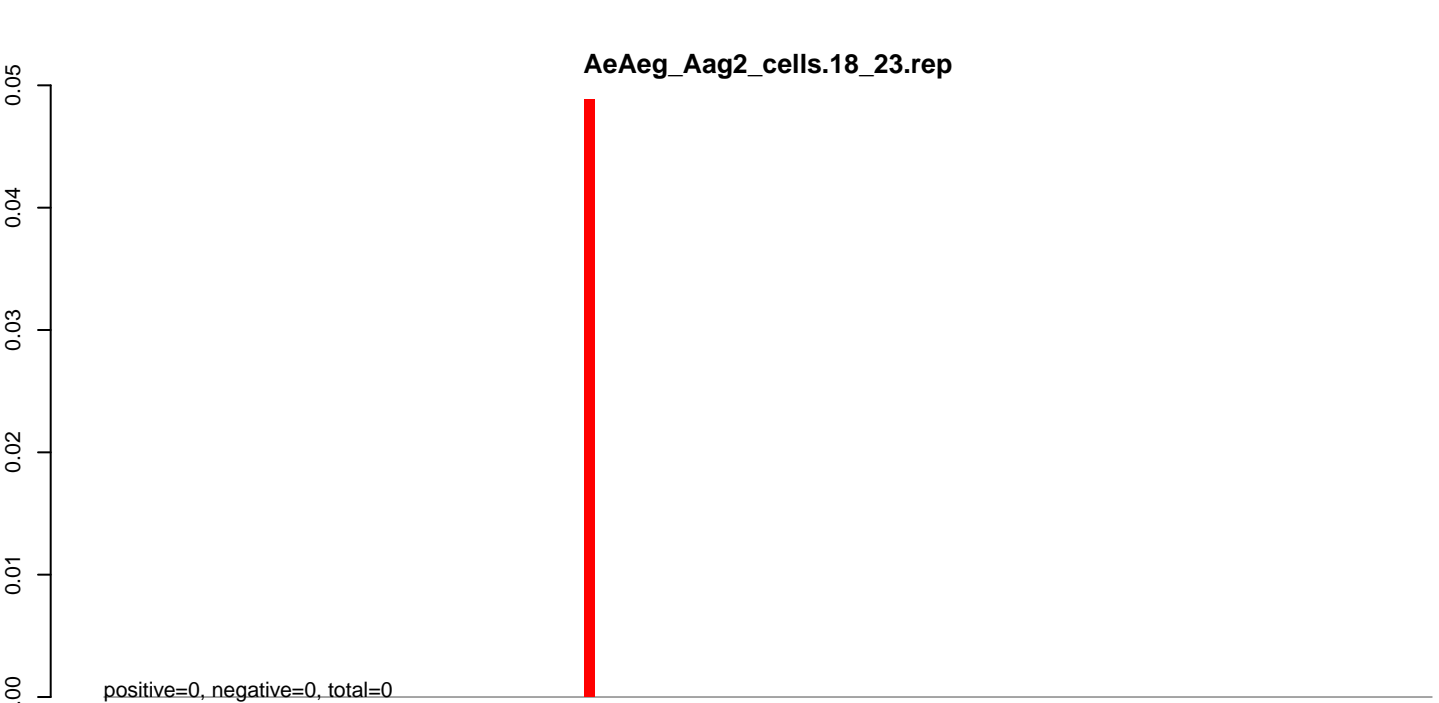
AeAeg_Aag2_cells.24_35.rep



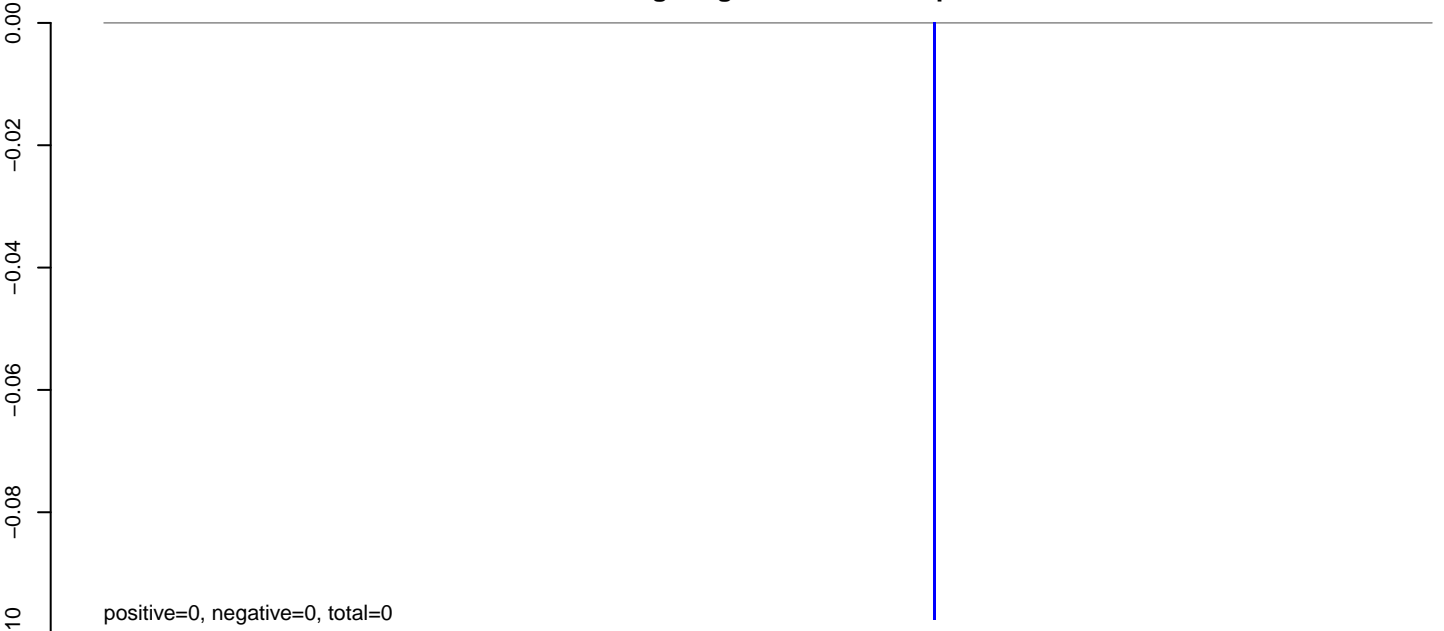
AeAeg_Aag2_cells.rep



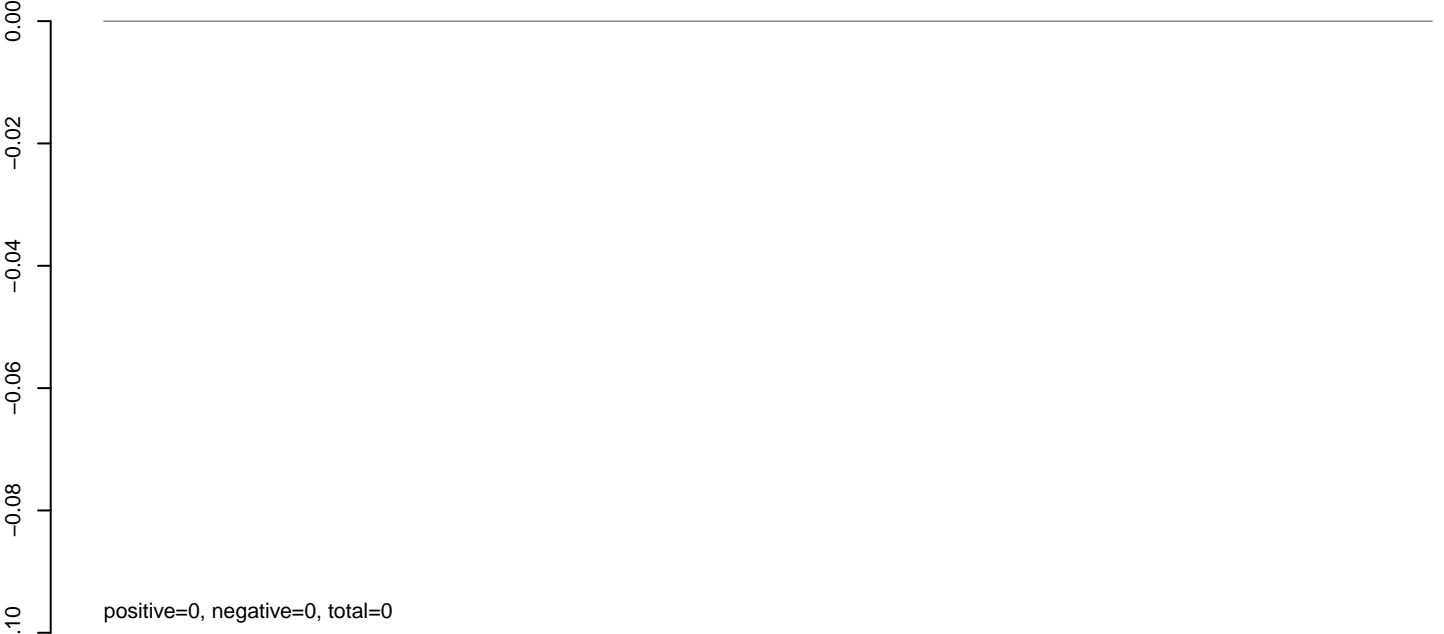
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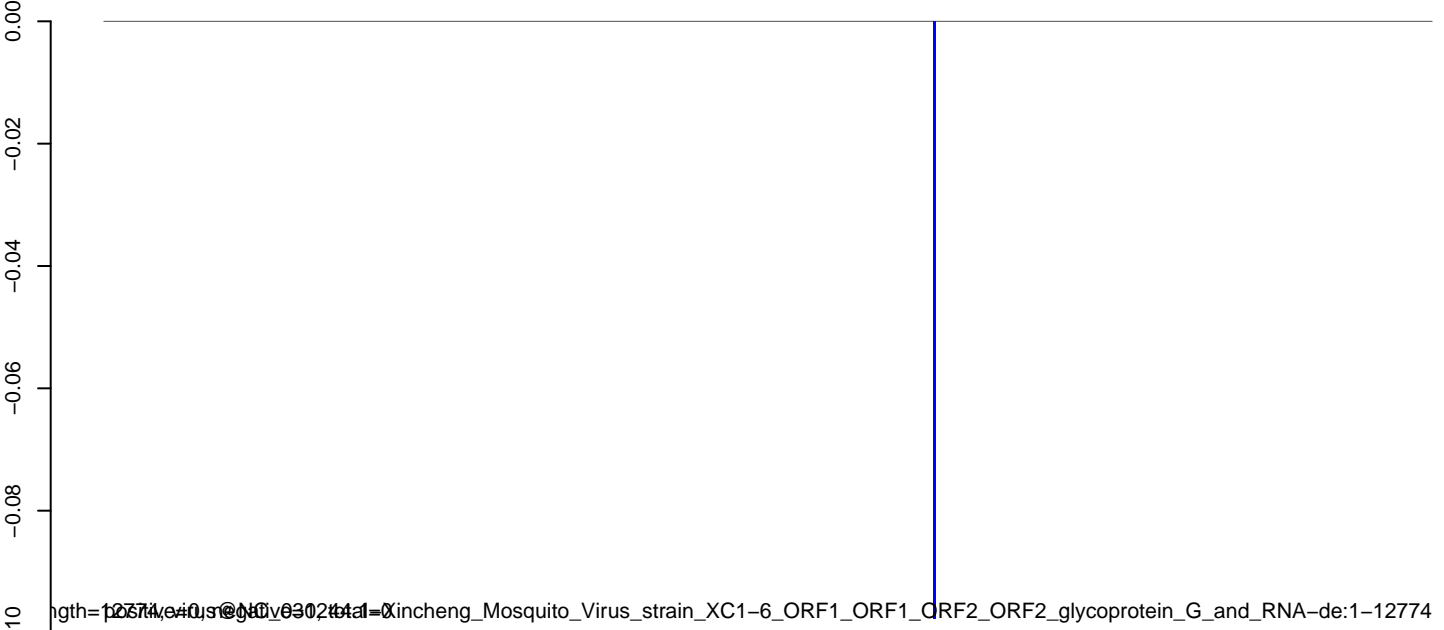
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AeAeg_Aag2_cells.24_35.rep

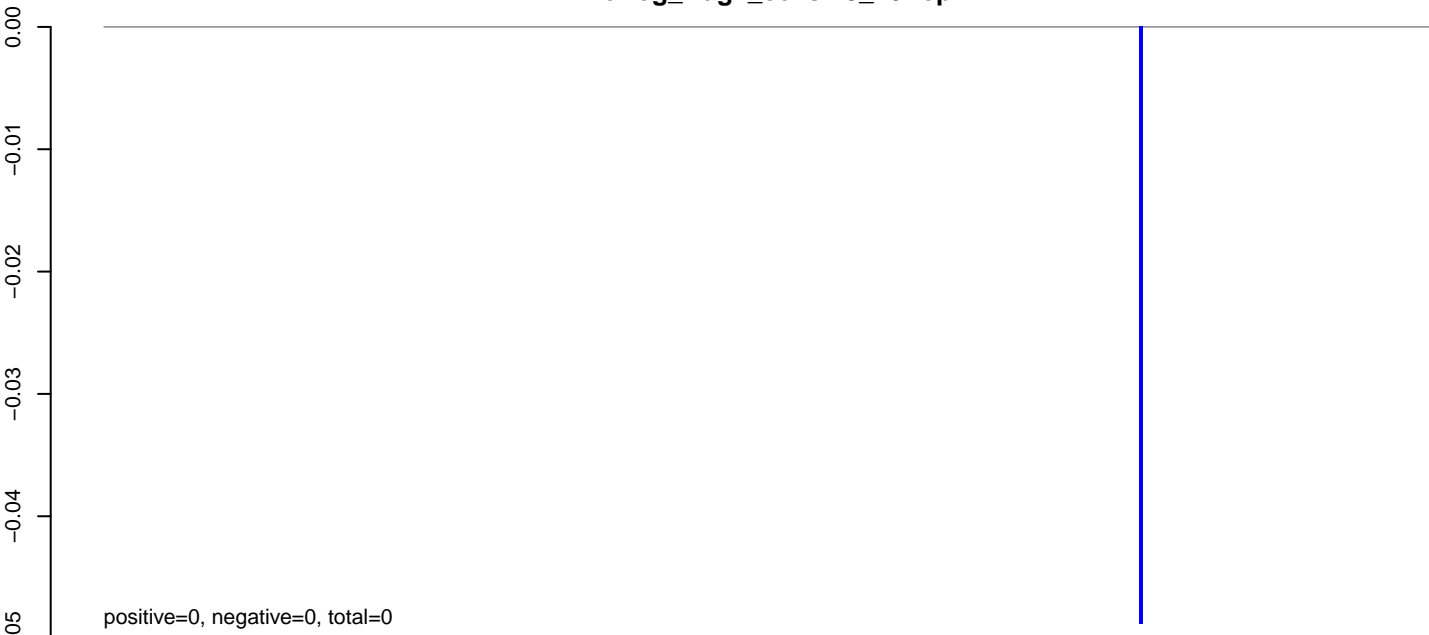


AeAeg_Aag2_cells.rep

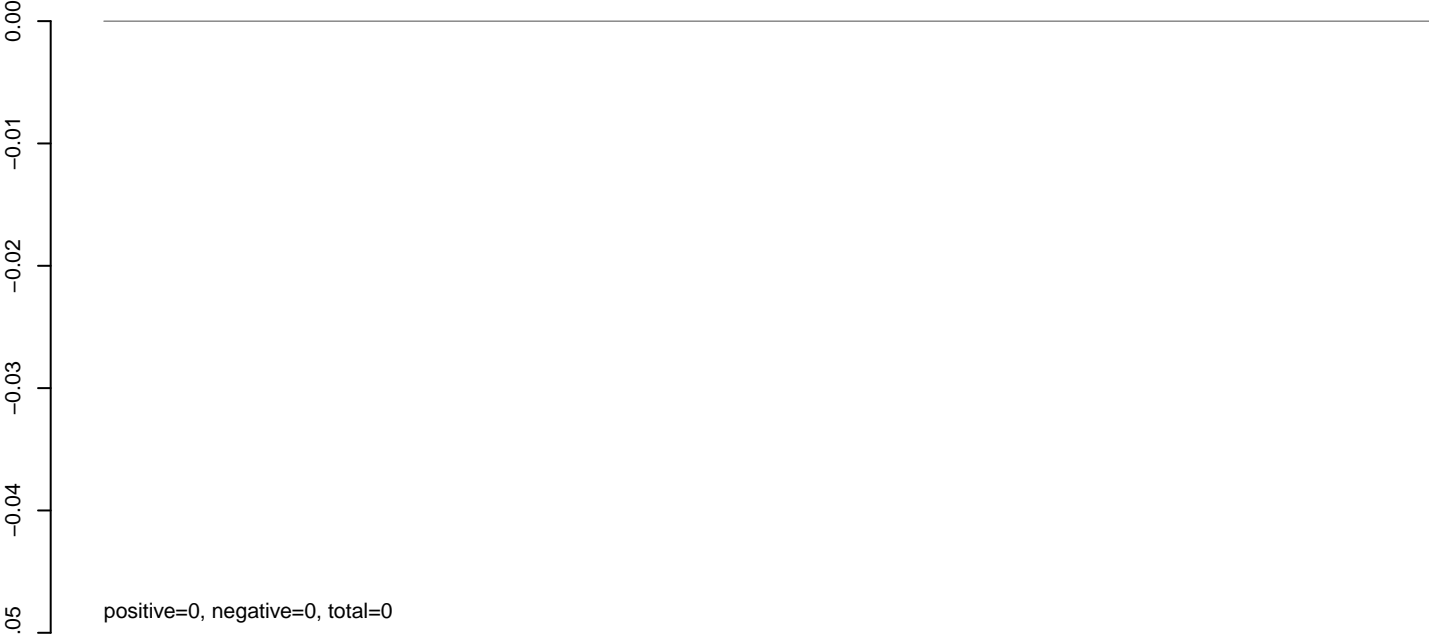


Xincheng_Mosquito_Virus_strain_XC1-6_ORF1_ORF1_ORF2_ORF2_glycoprotein_G_and_RNA-de:1-12774

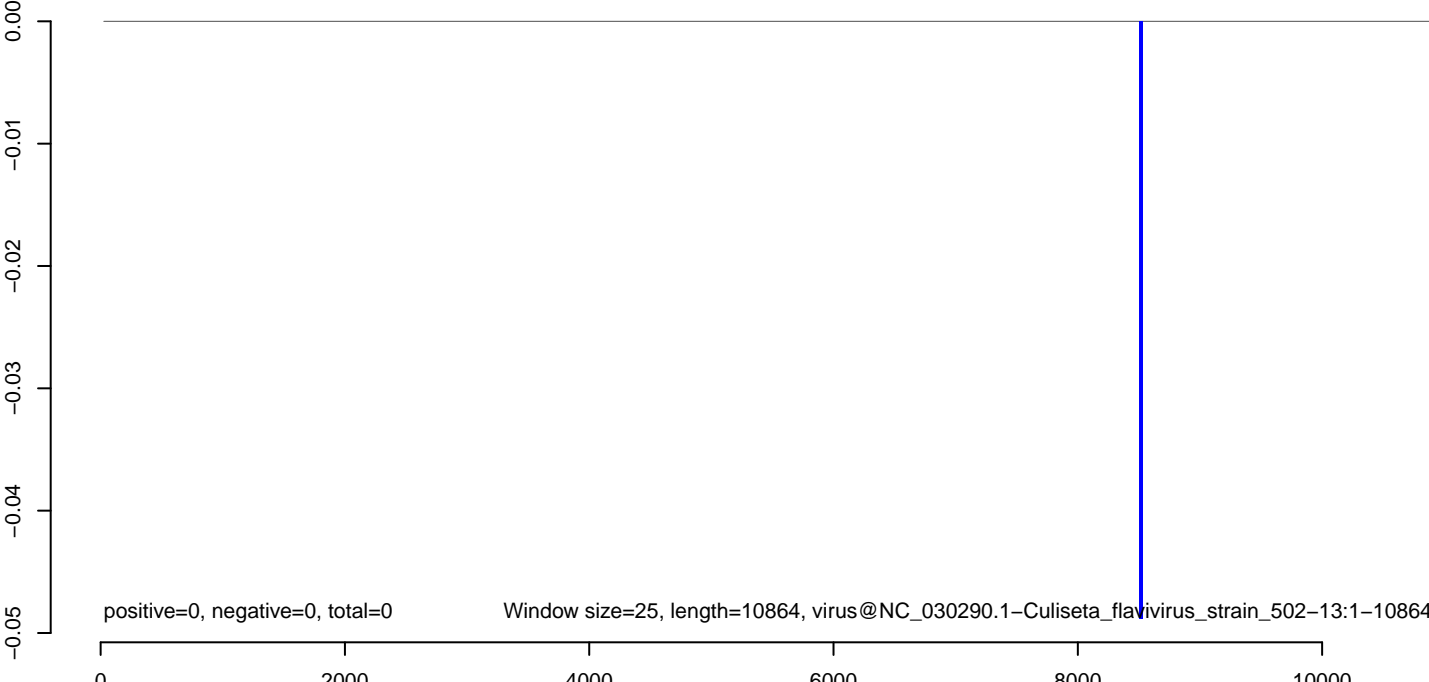
AeAeg_Aag2_cells.18_23.rep



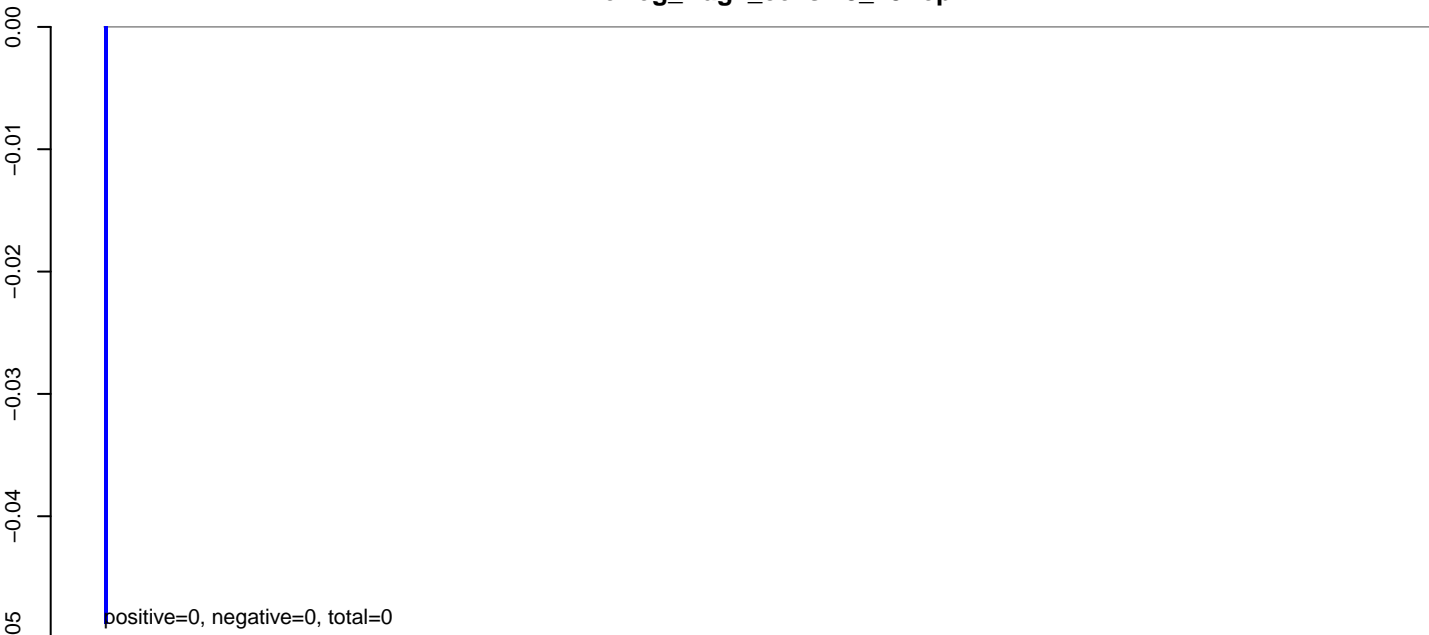
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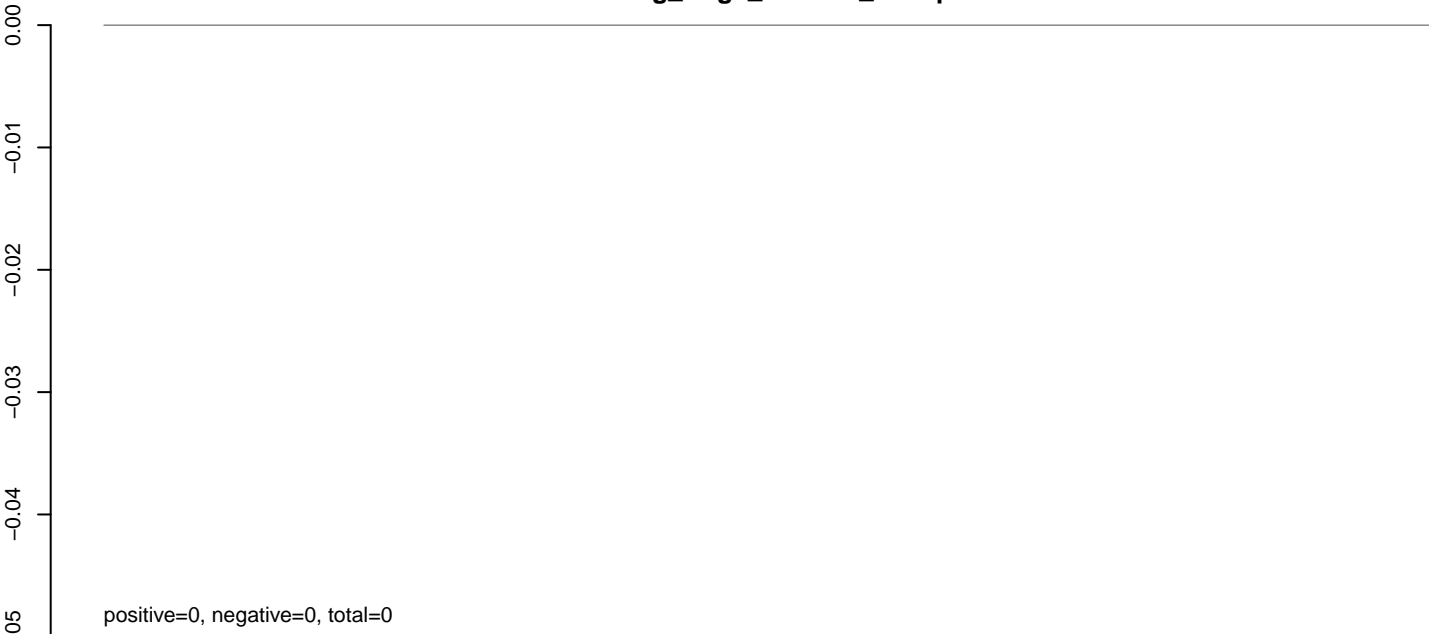
AeAeg_Aag2_cells.rep



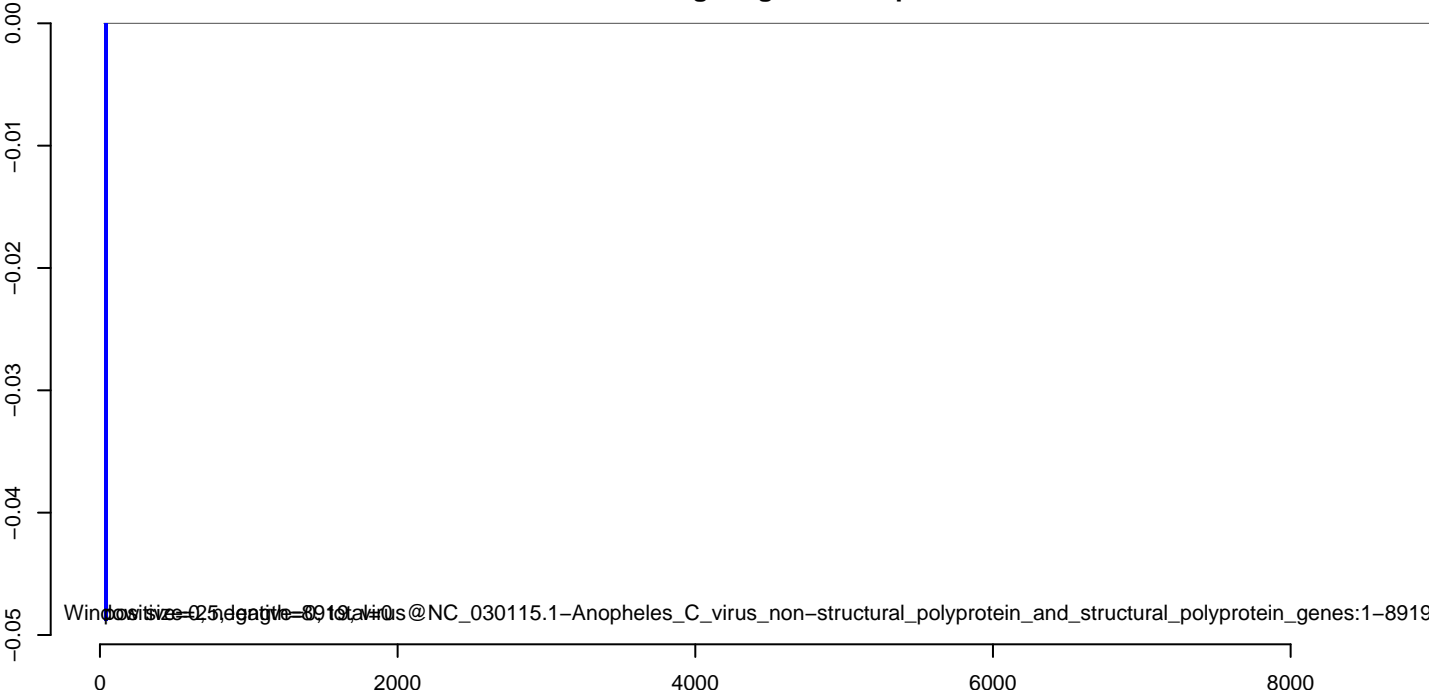
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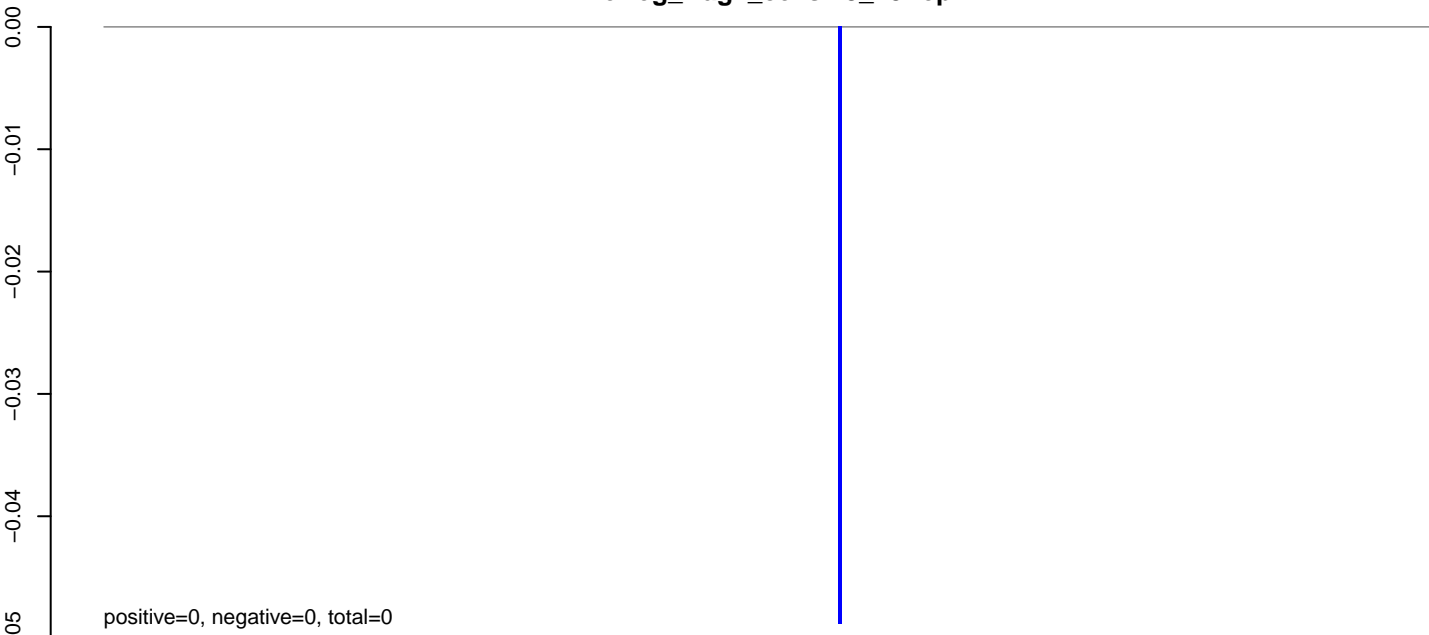
AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep

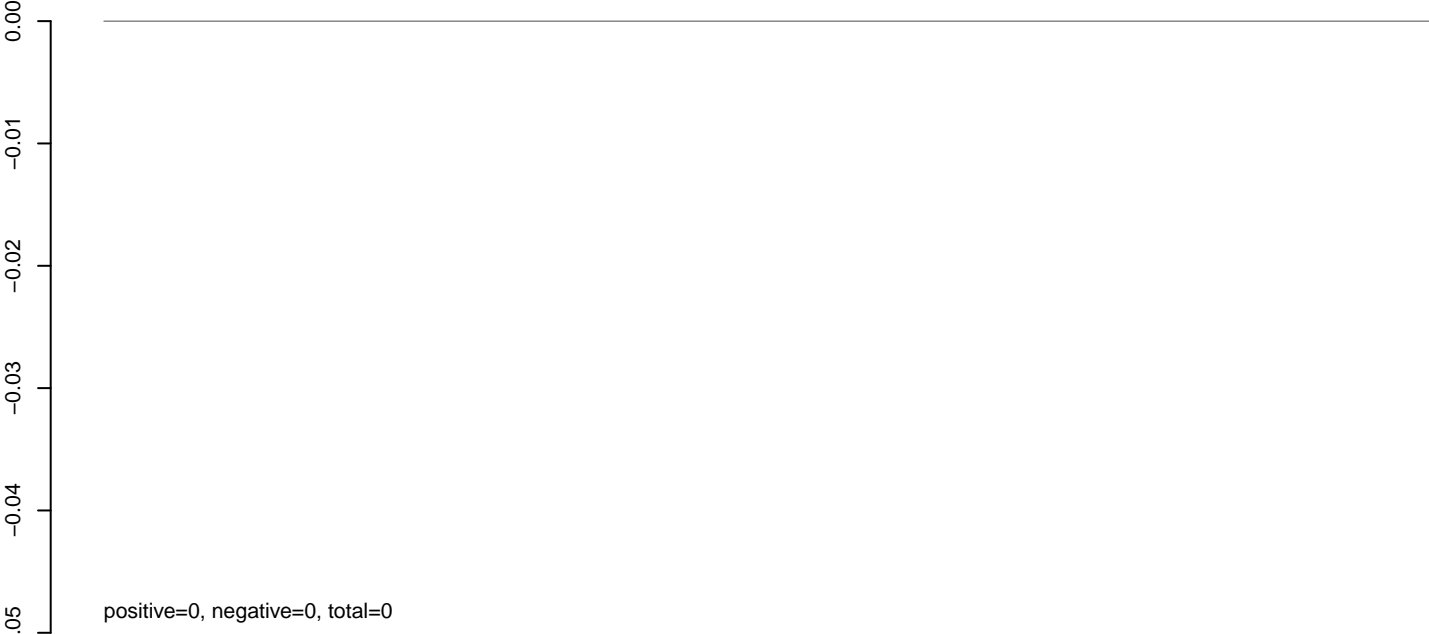


AeAeg_Aag2_cells.18_23.rep



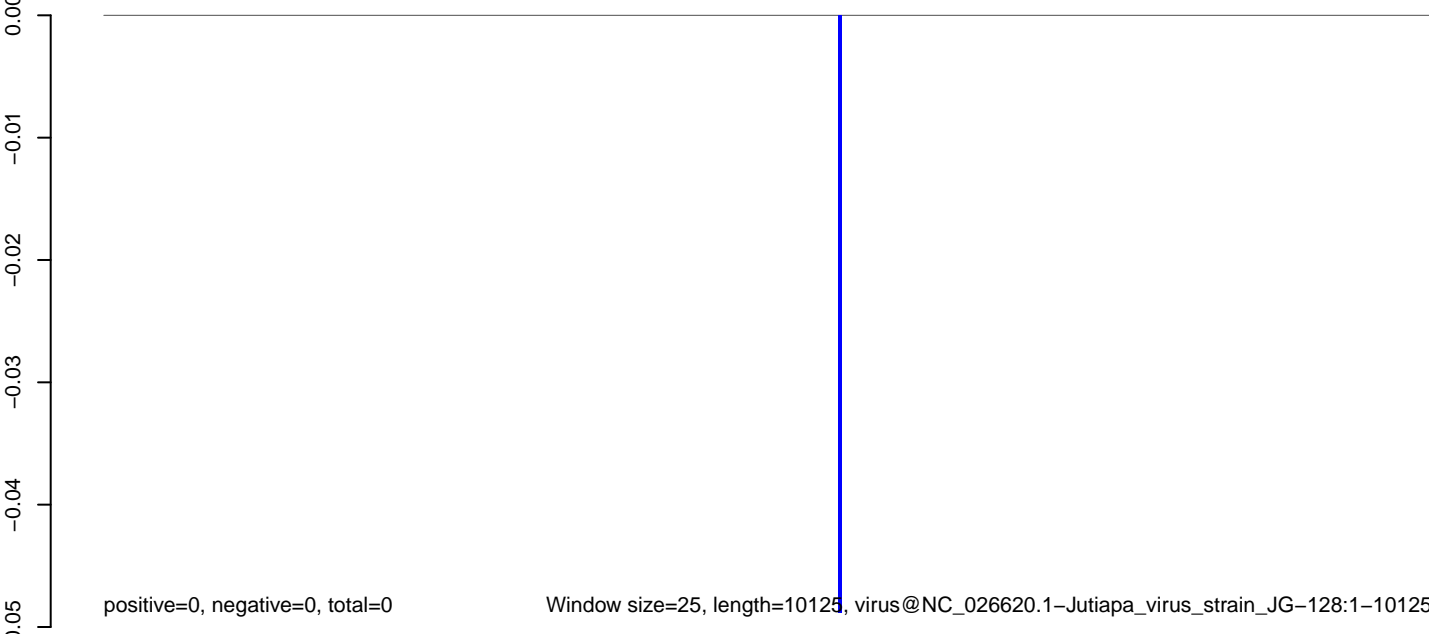
positive=0, negative=0, total=0

AeAeg_Aag2_cells.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_cells.rep

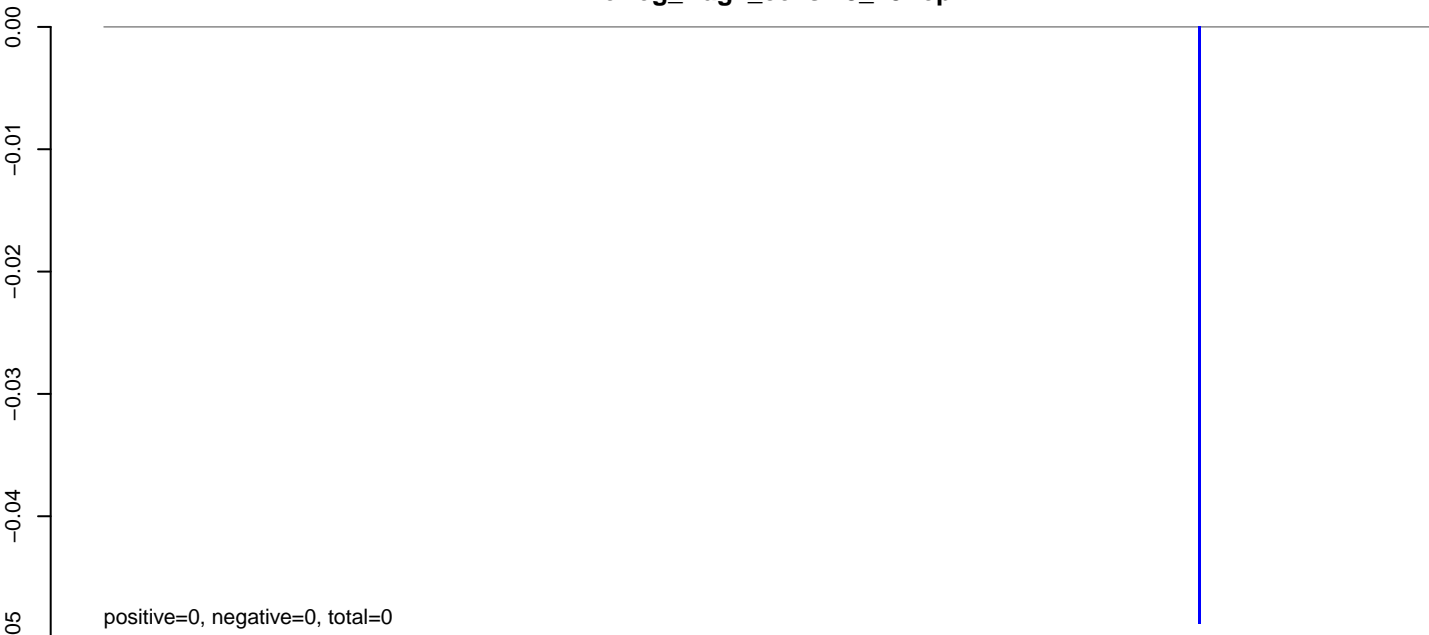


positive=0, negative=0, total=0

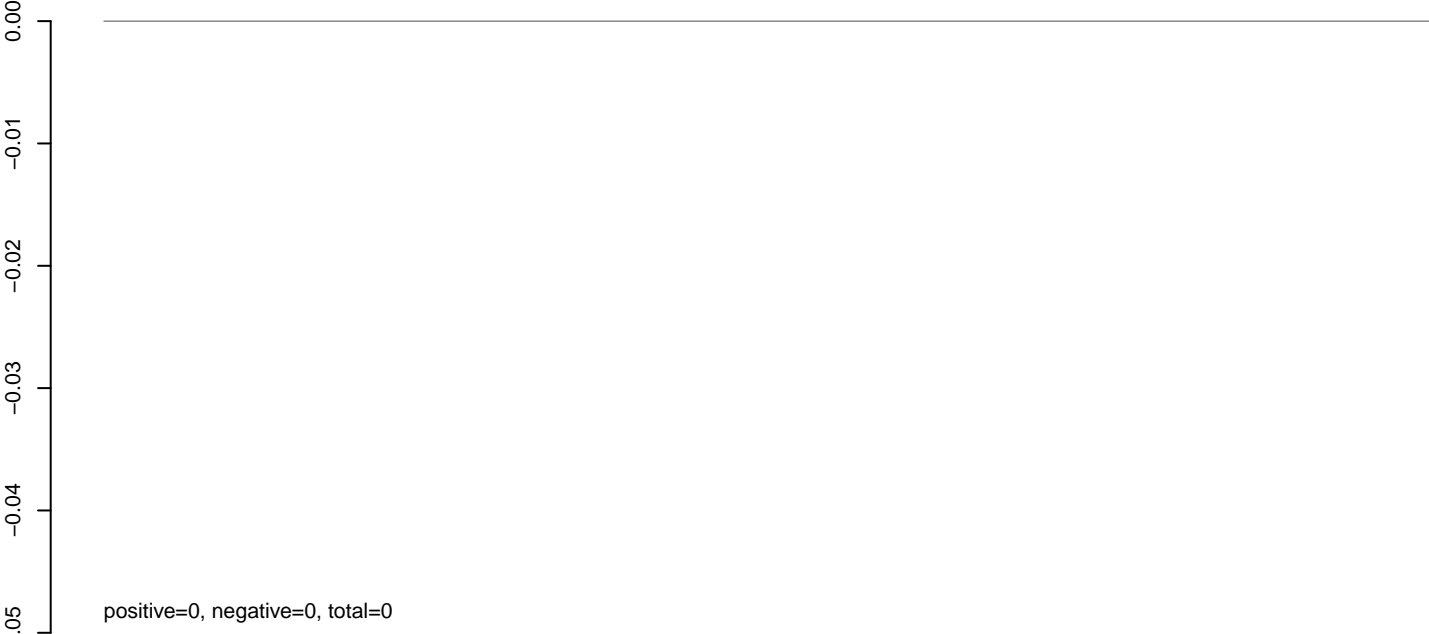
Window size=25, length=10125, virus@NC_026620.1-Jutiapa_virus_strain_JG-128:1-10125

0 2000 4000 6000 8000 10000

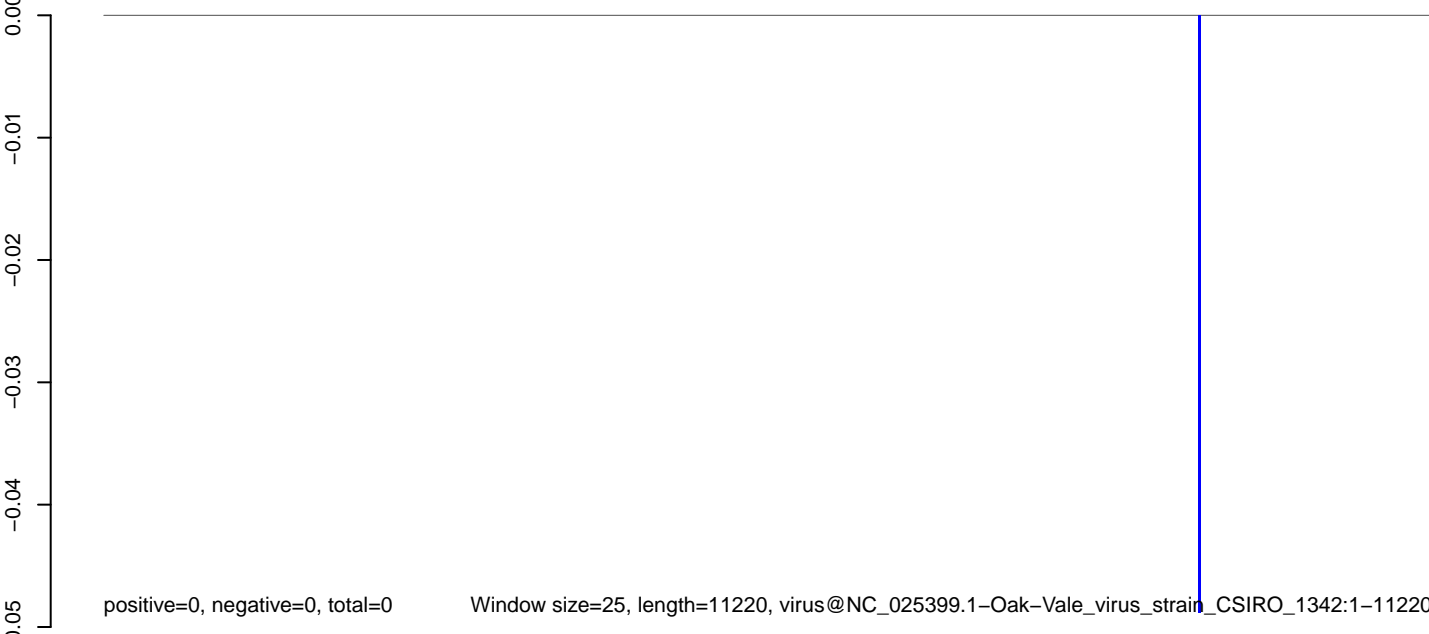
AeAeg_Aag2_cells.18_23.rep



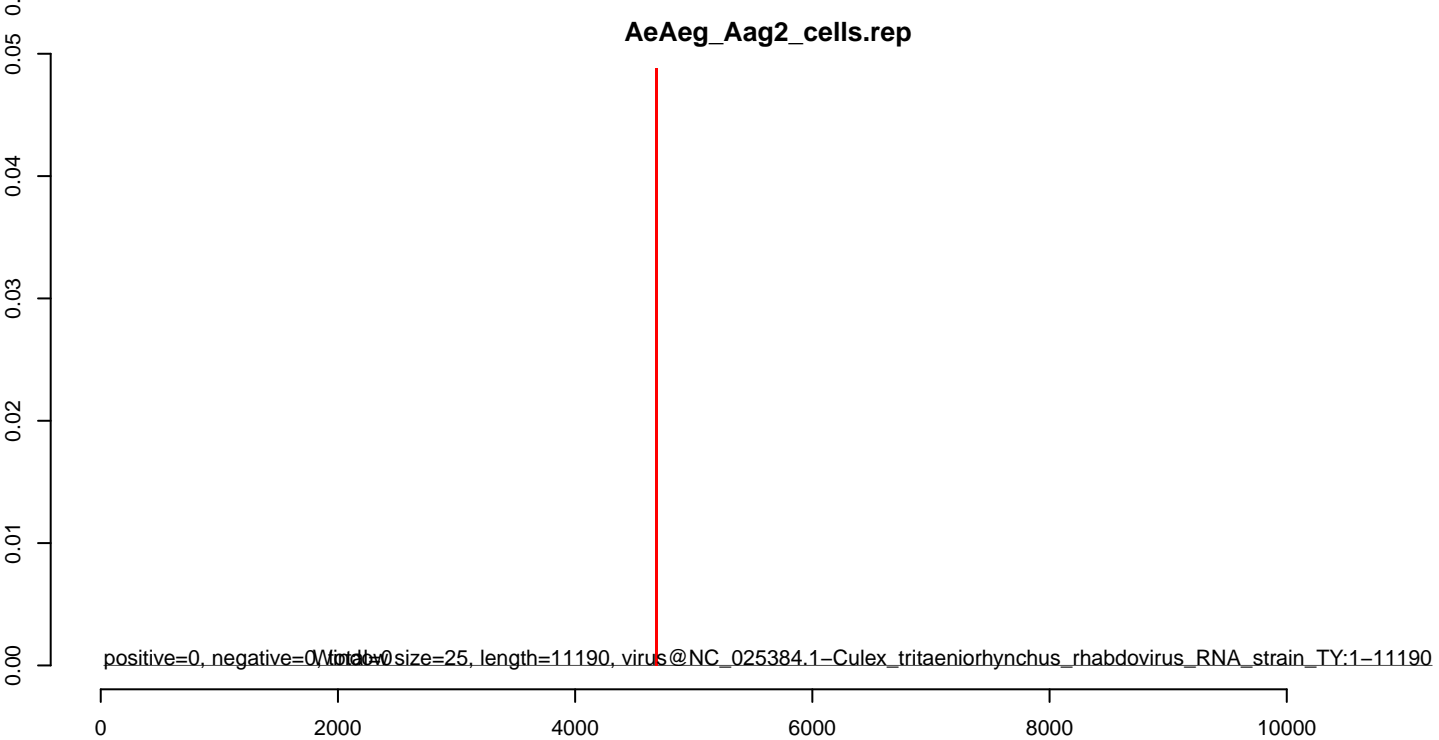
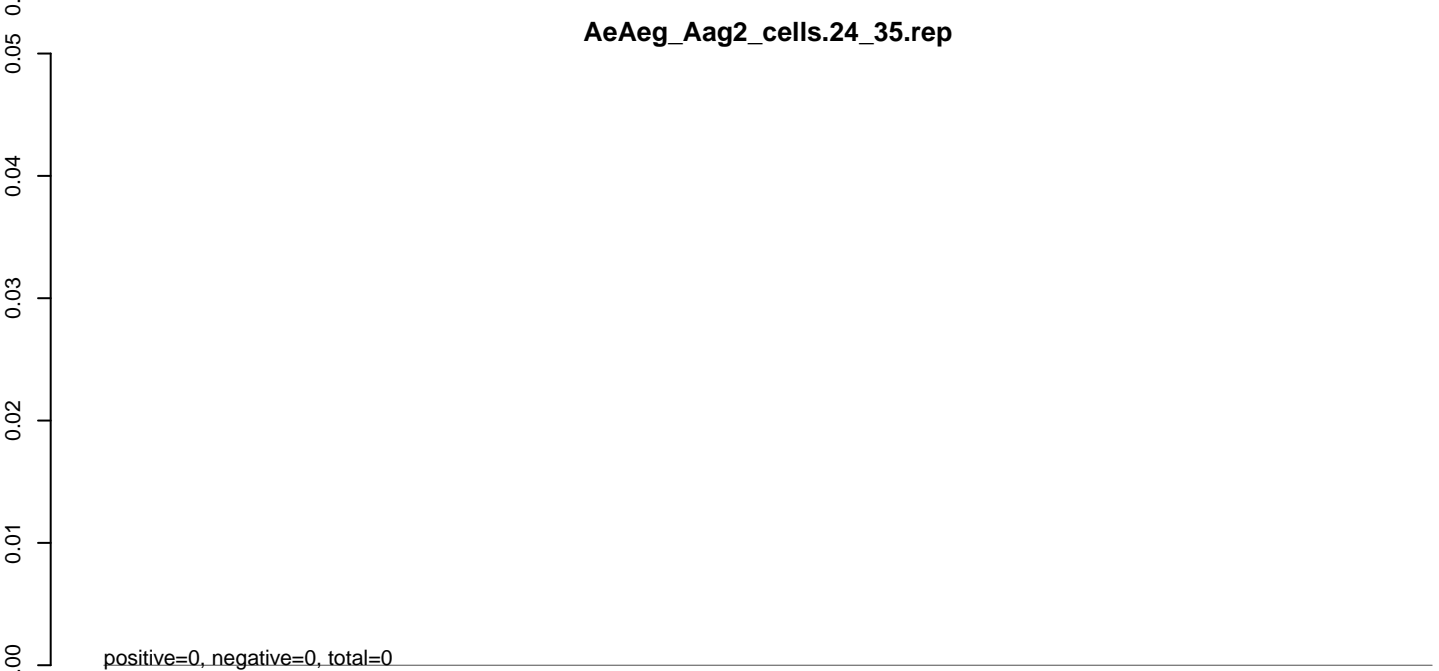
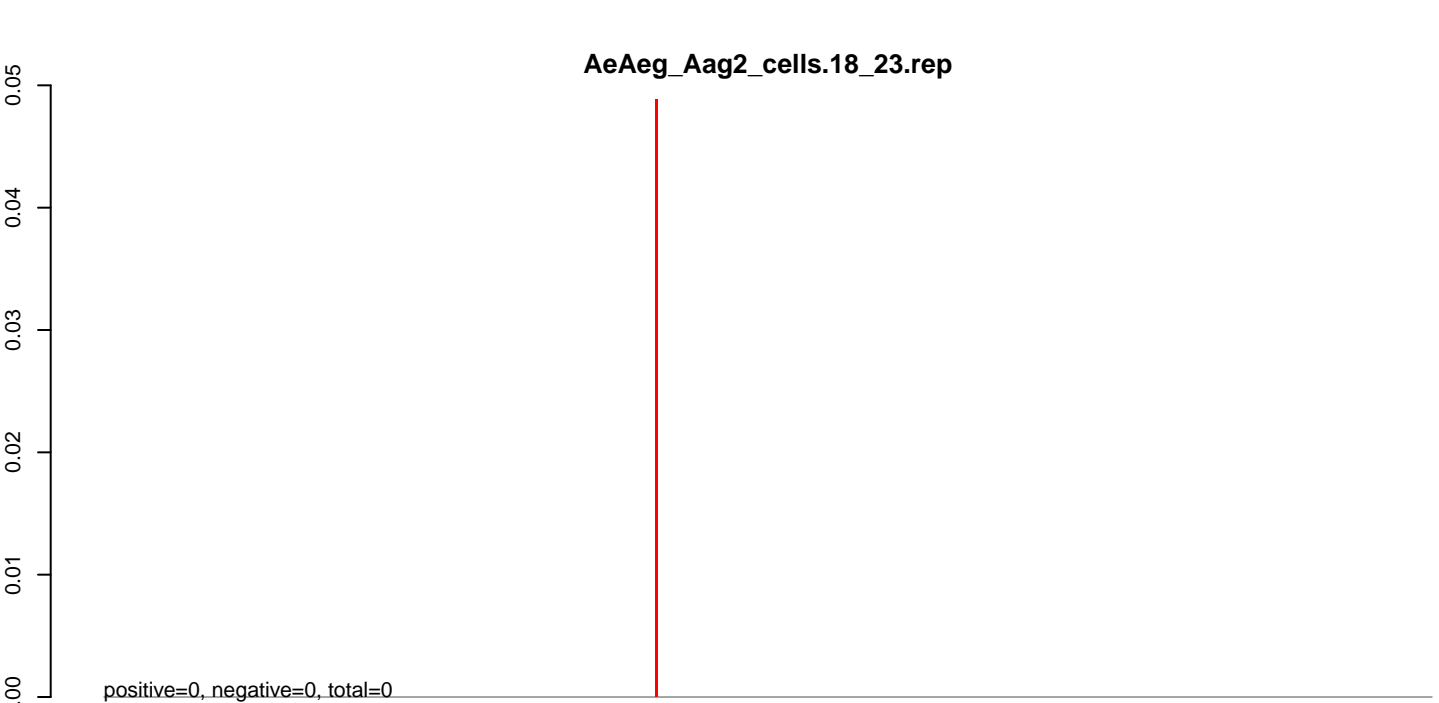
AeAeg_Aag2_cells.24_35.rep



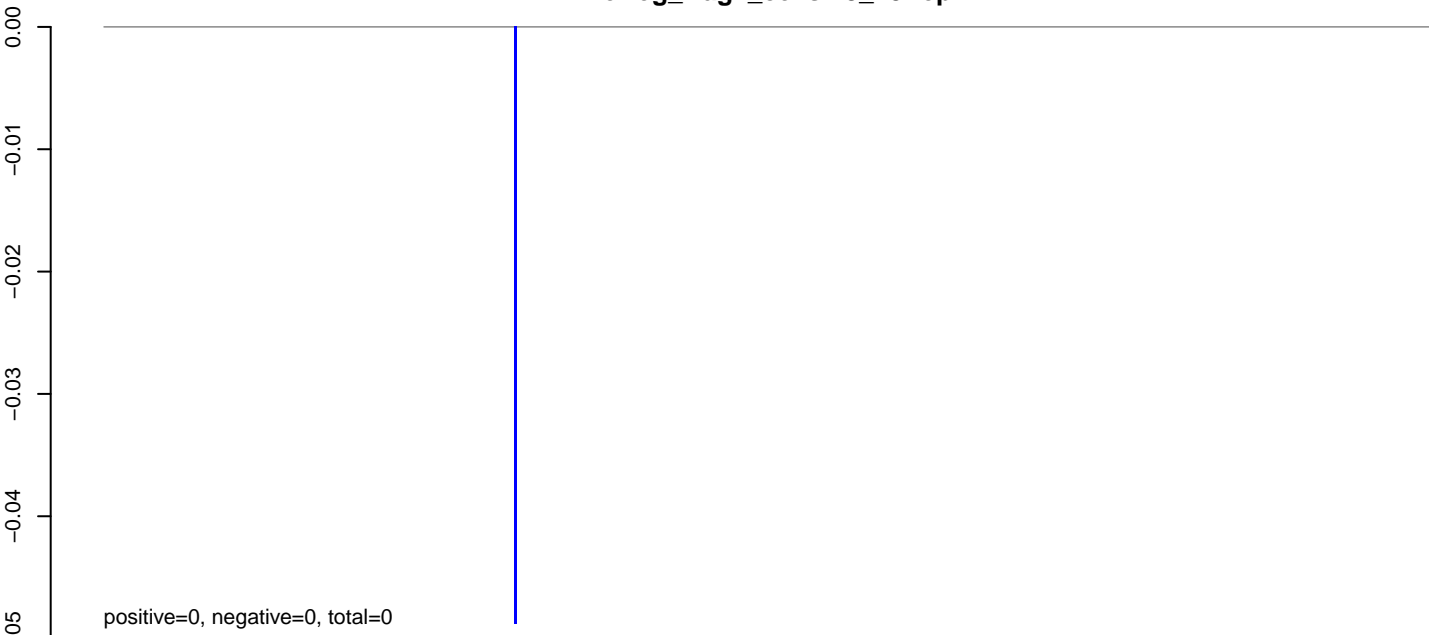
AeAeg_Aag2_cells.rep



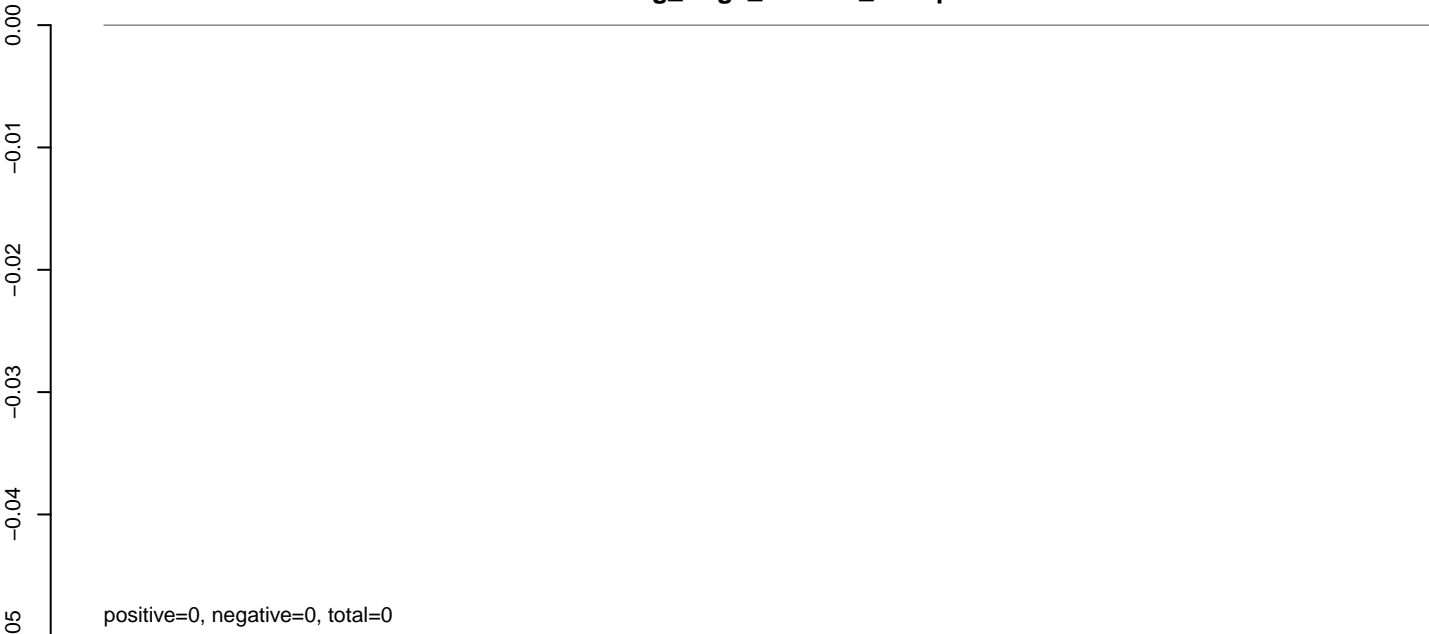
0 2000 4000 6000 8000 10000



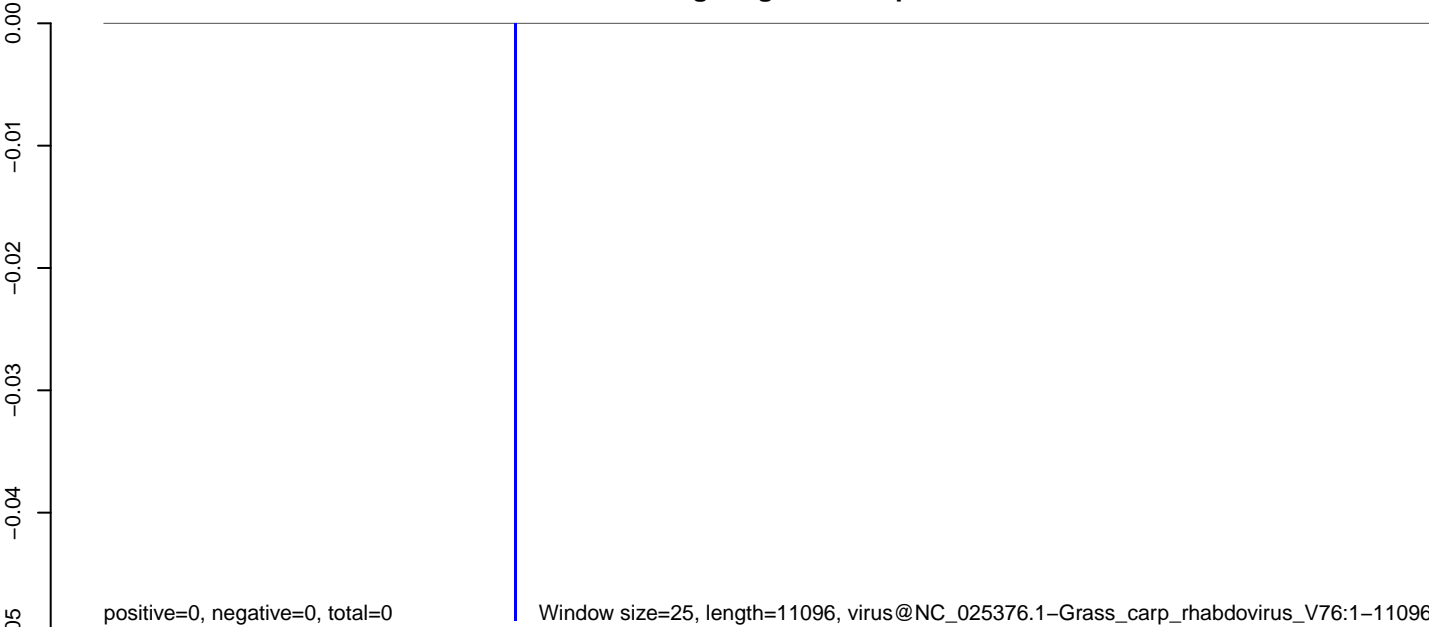
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

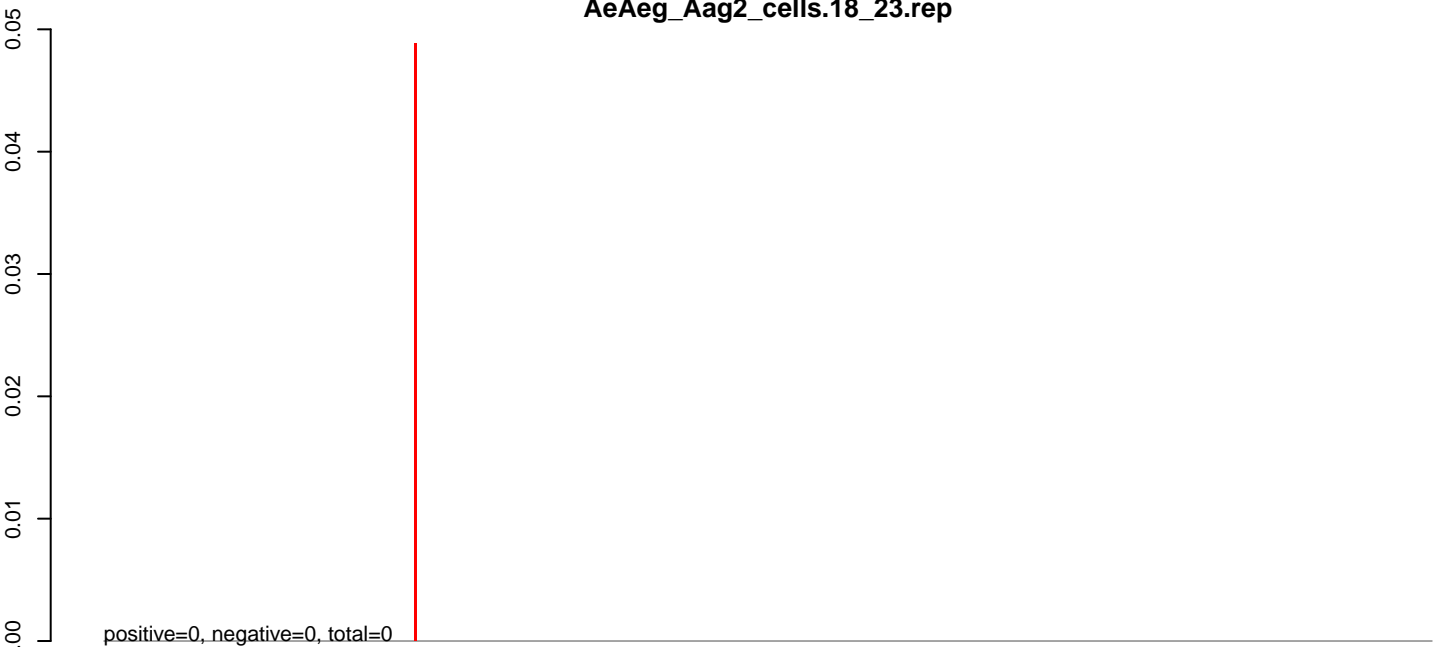


AeAeg_Aag2_cells.rep

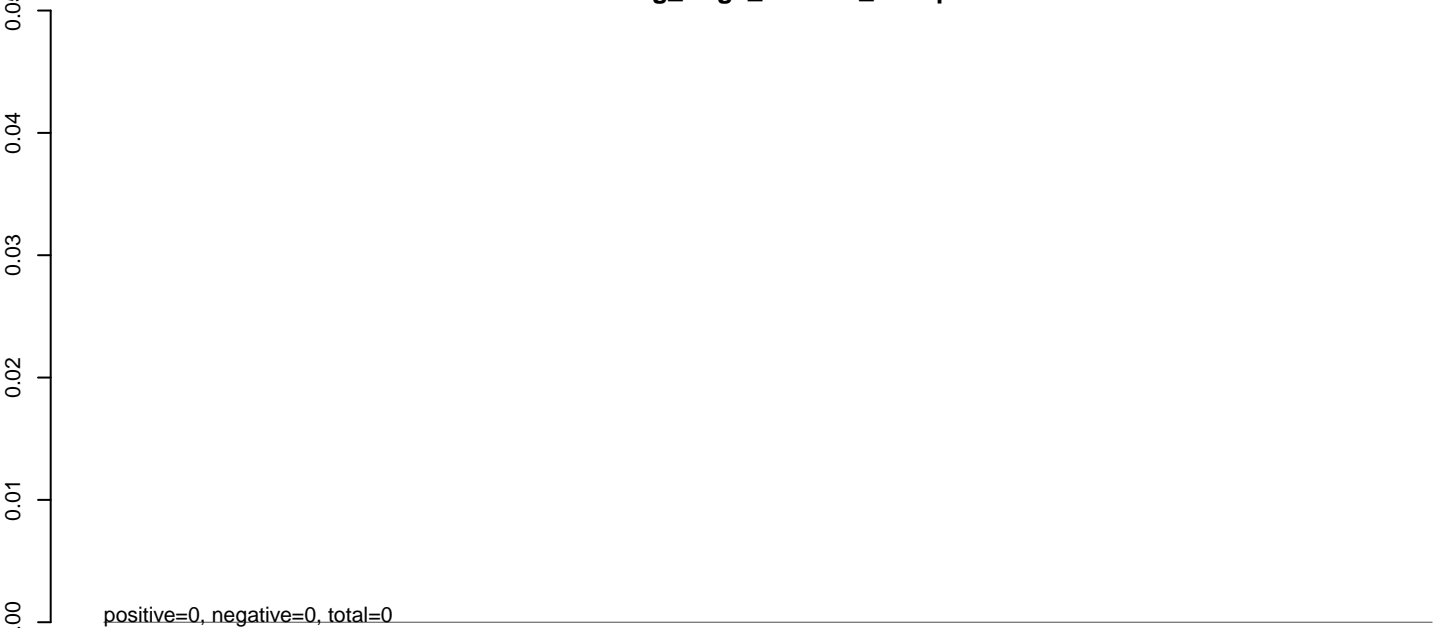


0 2000 4000 6000 8000 10000

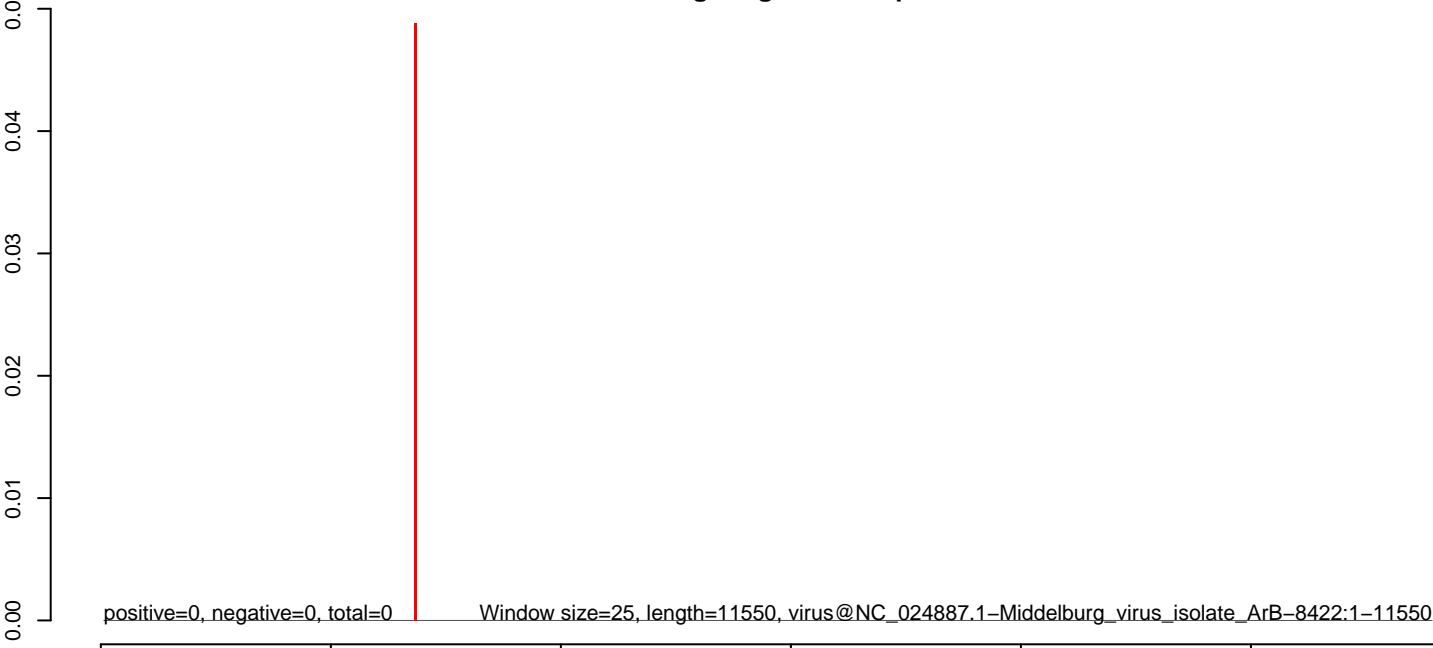
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

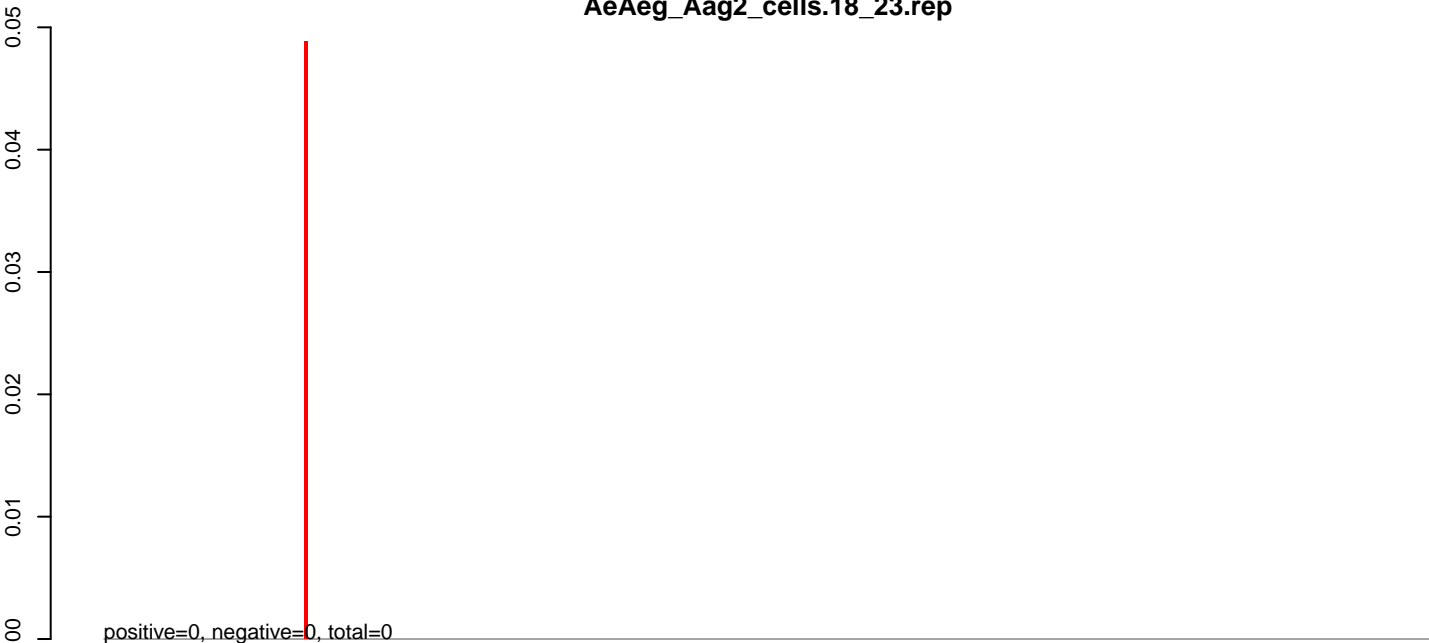


AeAeg_Aag2_cells.rep

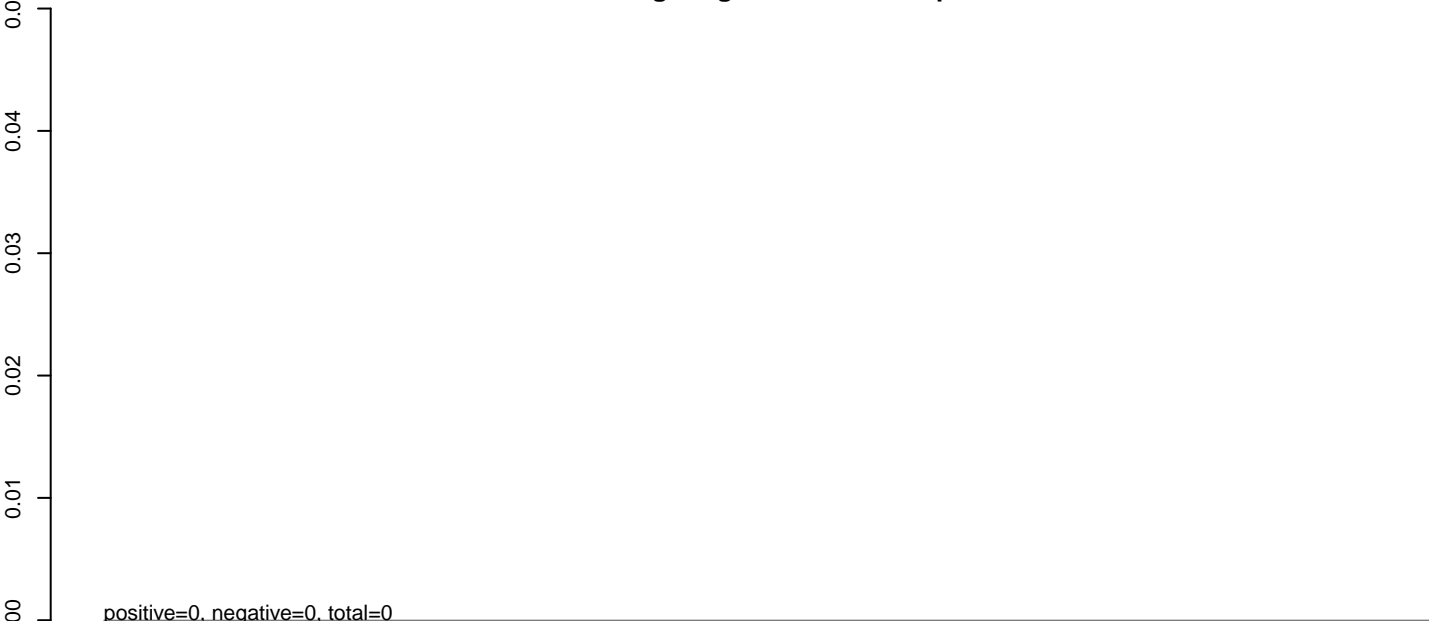


0 2000 4000 6000 8000 10000 12000

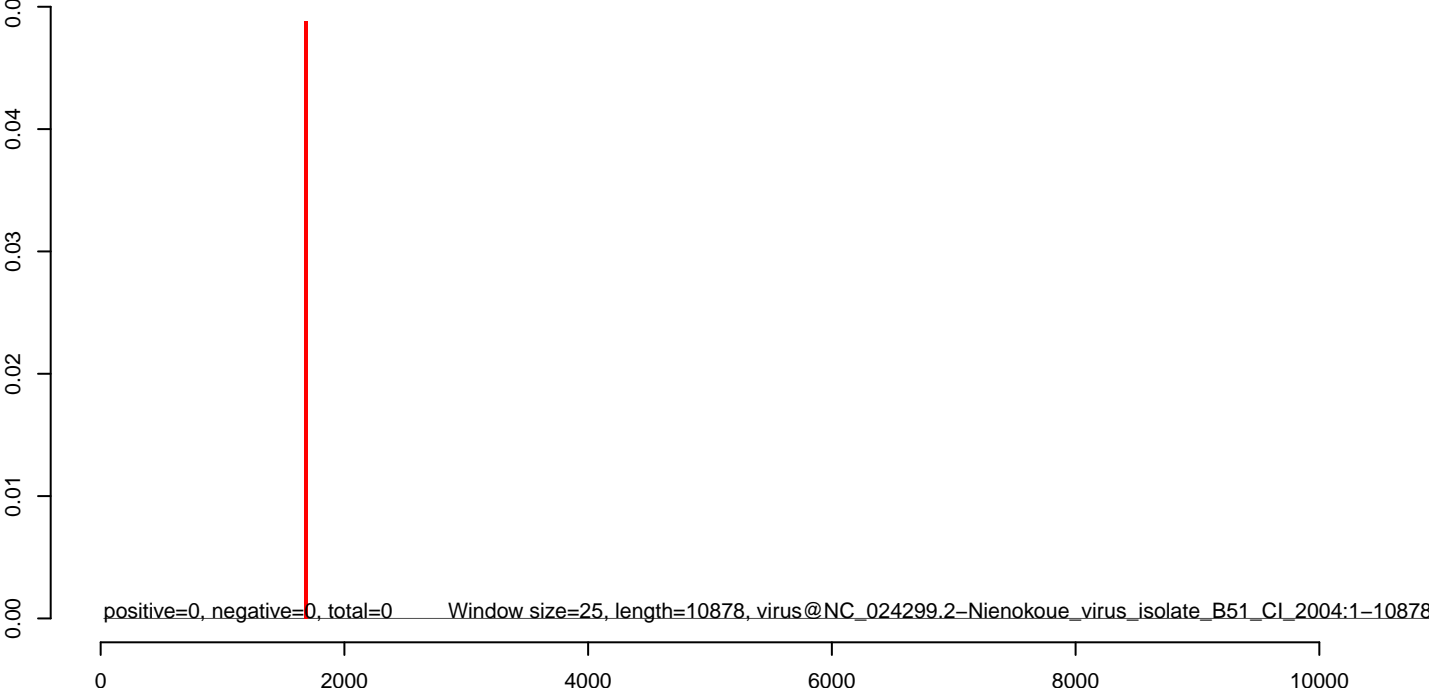
AeAeg_Aag2_cells.18_23.rep



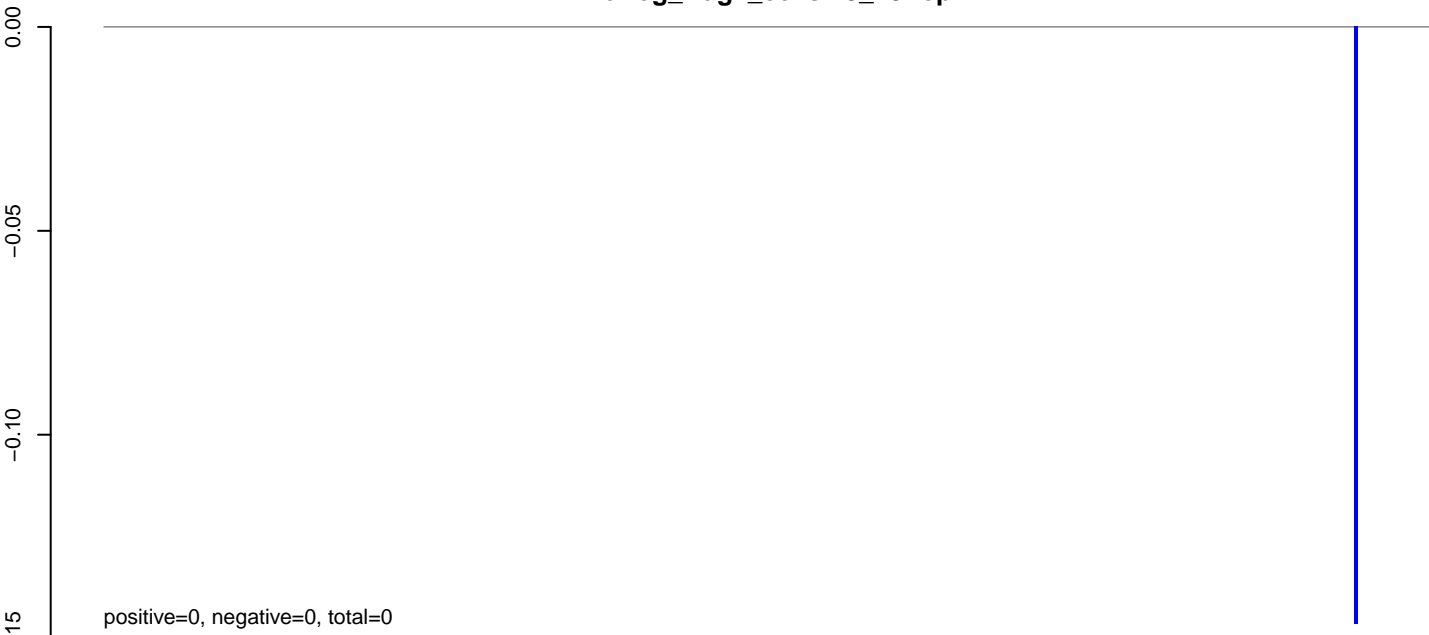
AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep

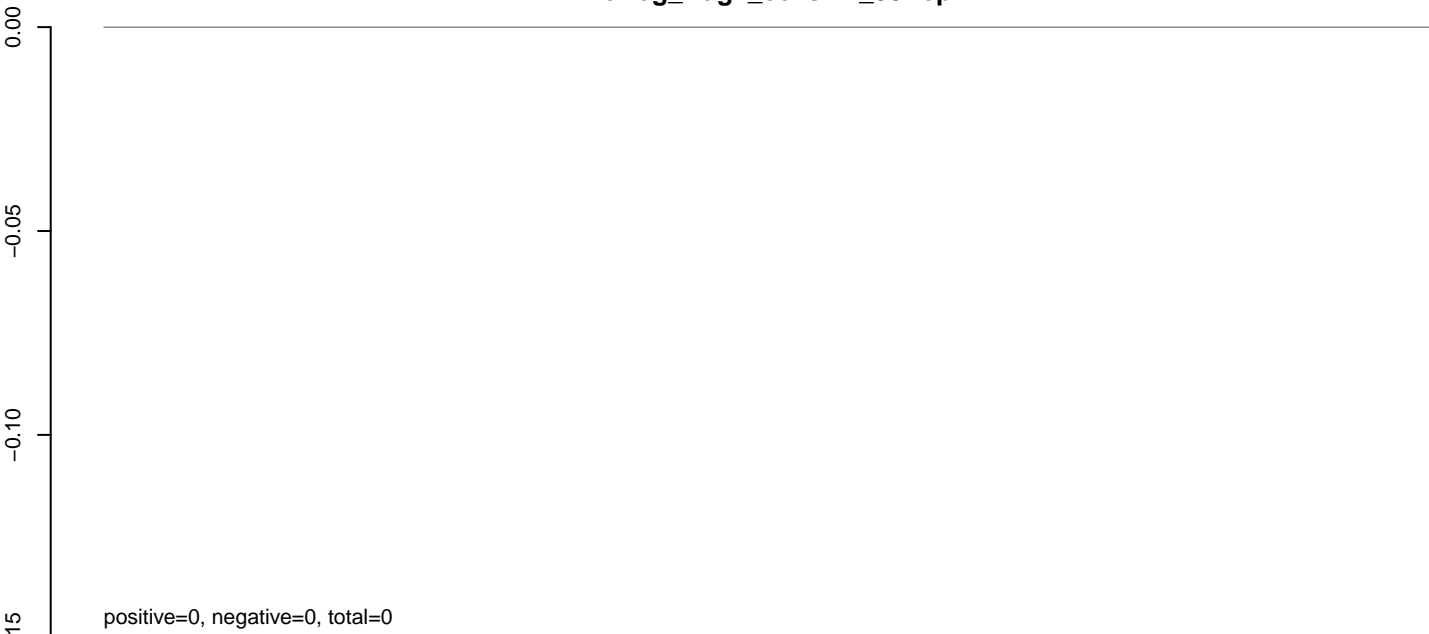


AeAeg_Aag2_cells.18_23.rep



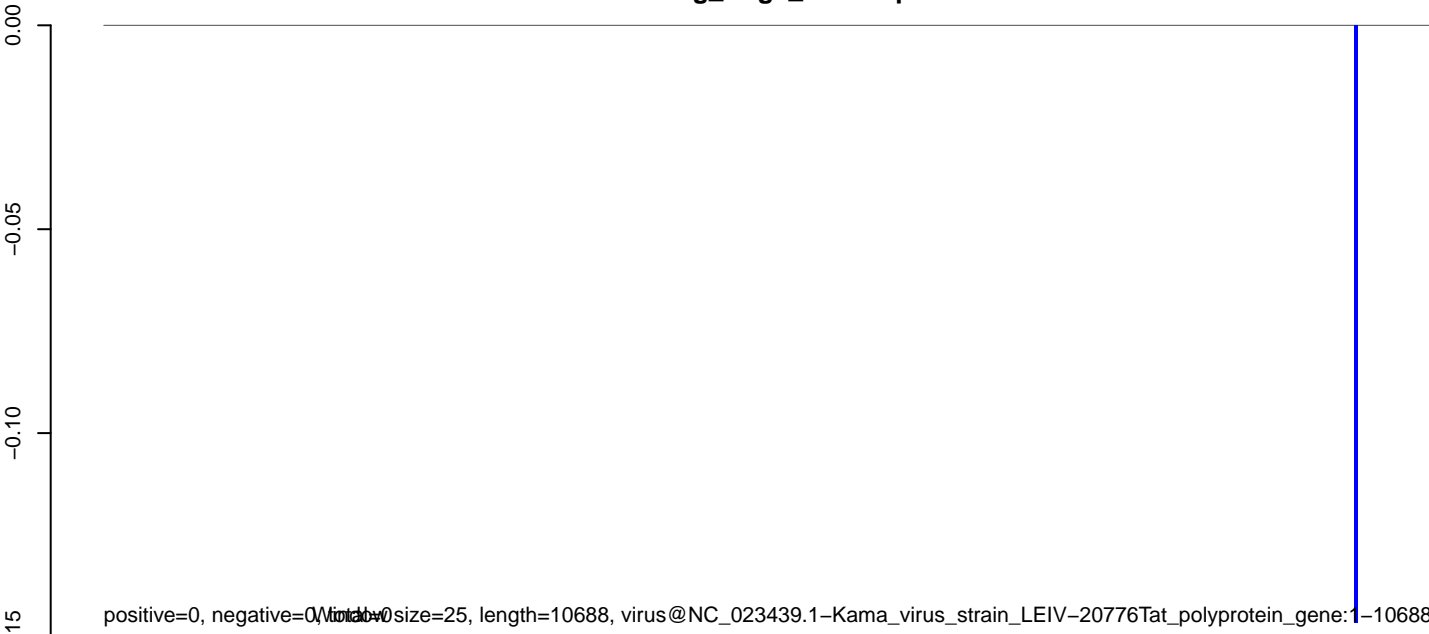
positive=0, negative=0, total=0

AeAeg_Aag2_cells.24_35.rep



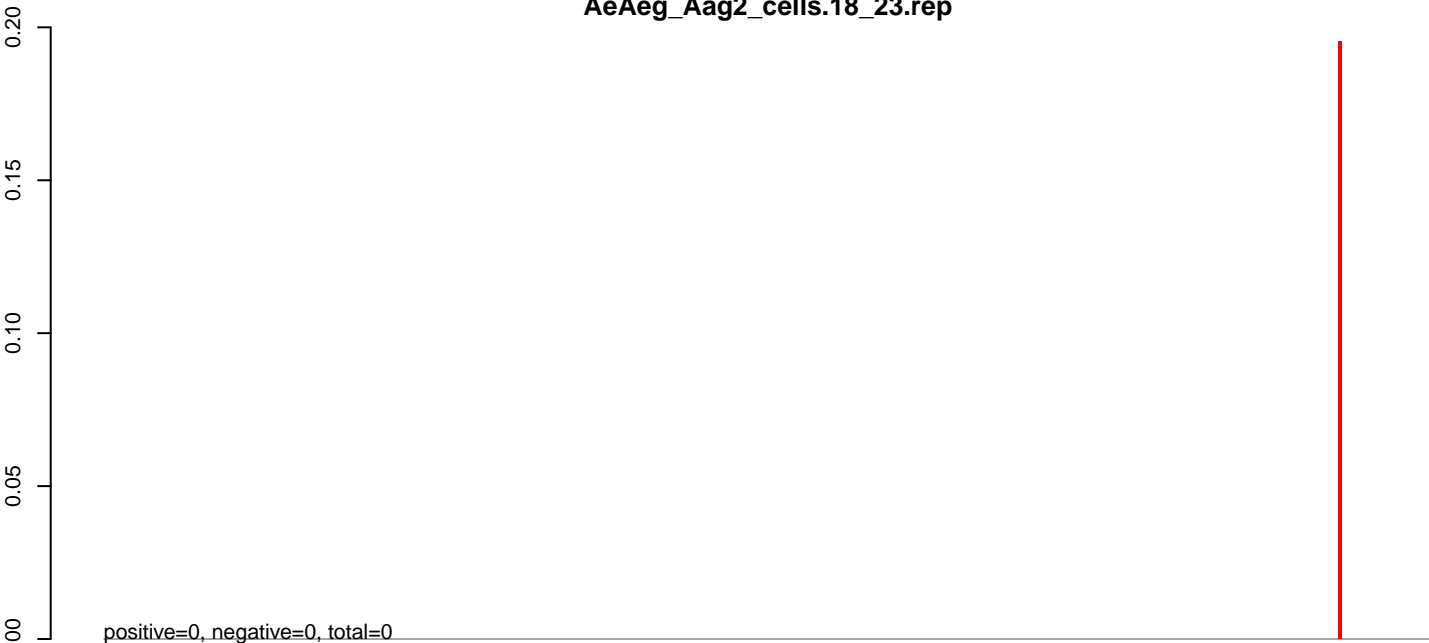
positive=0, negative=0, total=0

AeAeg_Aag2_cells.rep

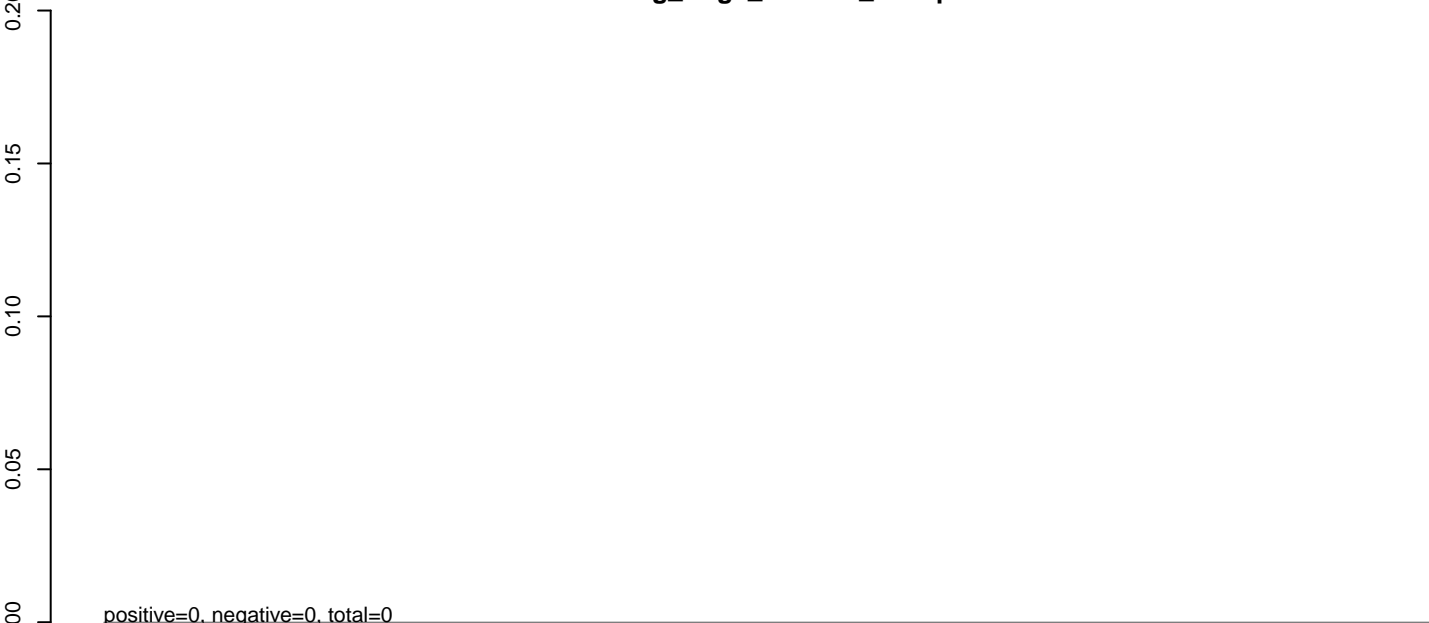


positive=0, negative=0, total=0, window size=25, length=10688, virus@NC_023439.1-Kama_virus_strain_LEIV-20776Tat_polyprotein_gene:1-10688

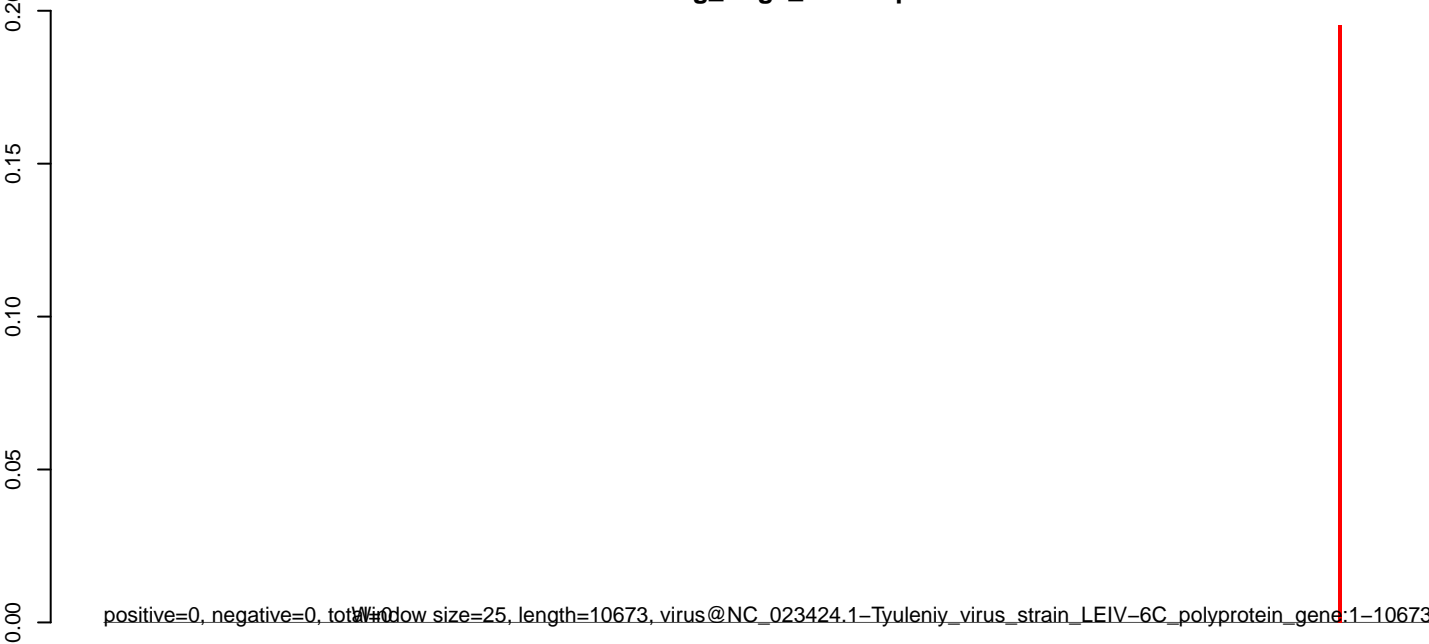
AeAeg_Aag2_cells.18_23.rep



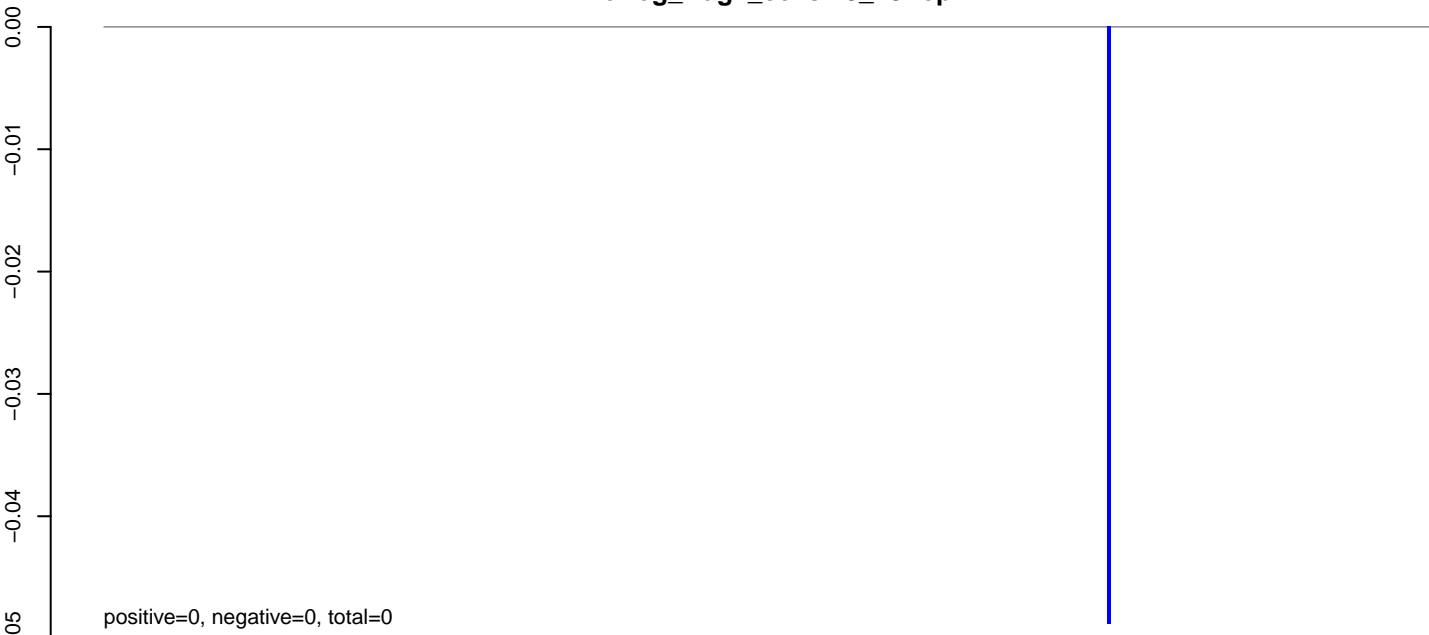
AeAeg_Aag2_cells.24_35.rep



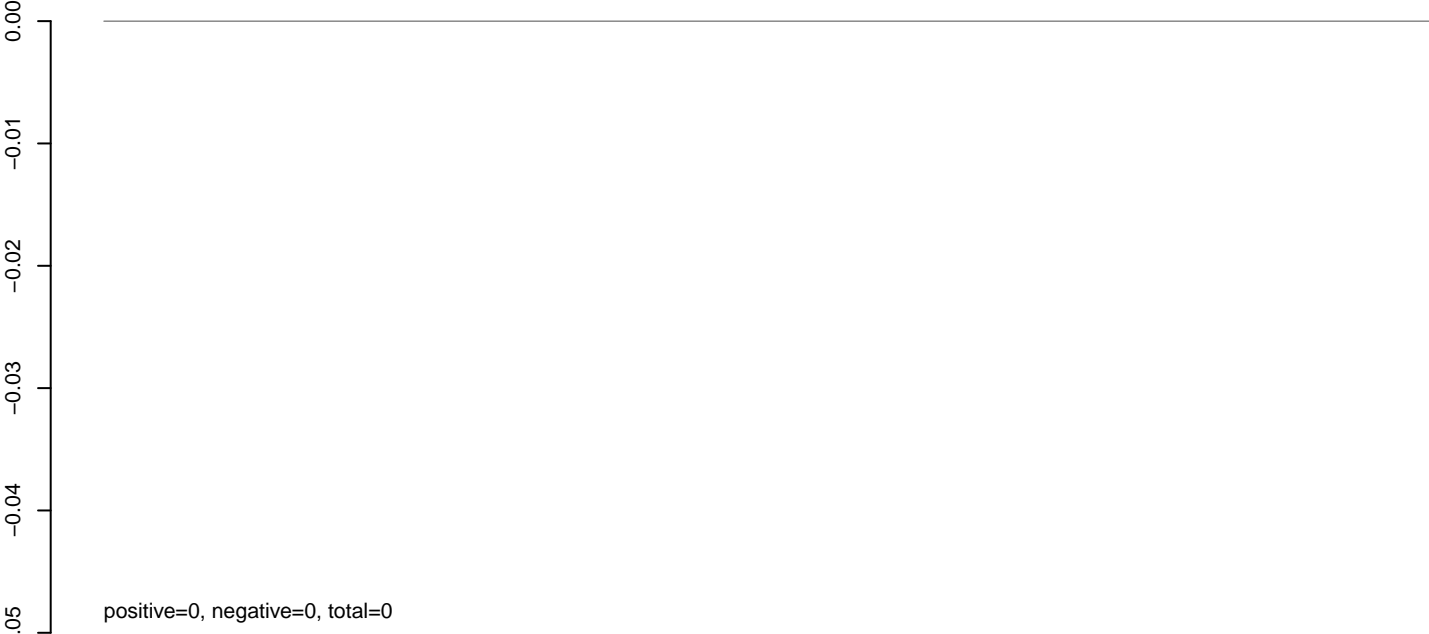
AeAeg_Aag2_cells.rep



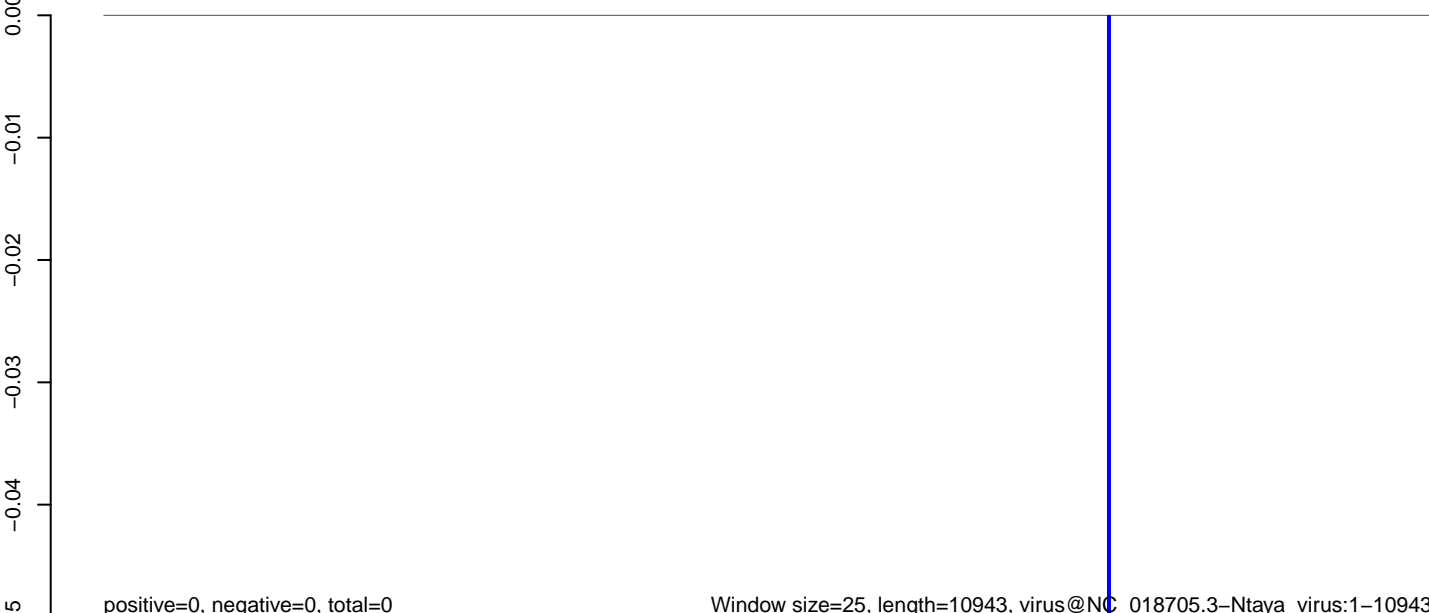
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



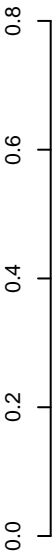
AeAeg_Aag2_cells.rep



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

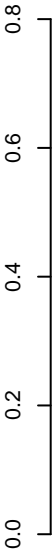
0 2000 4000 6000 8000 10000

AeAeg_Aag2_cells.18_23.rep



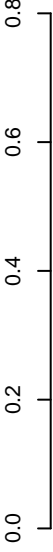
positive=2, negative=0, total=2

AeAeg_Aag2_cells.24_35.rep



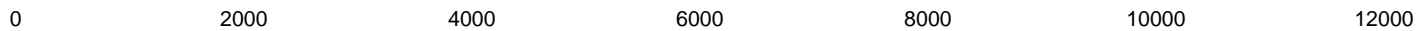
positive=0, negative=0, total=0

AeAeg_Aag2_cells.rep

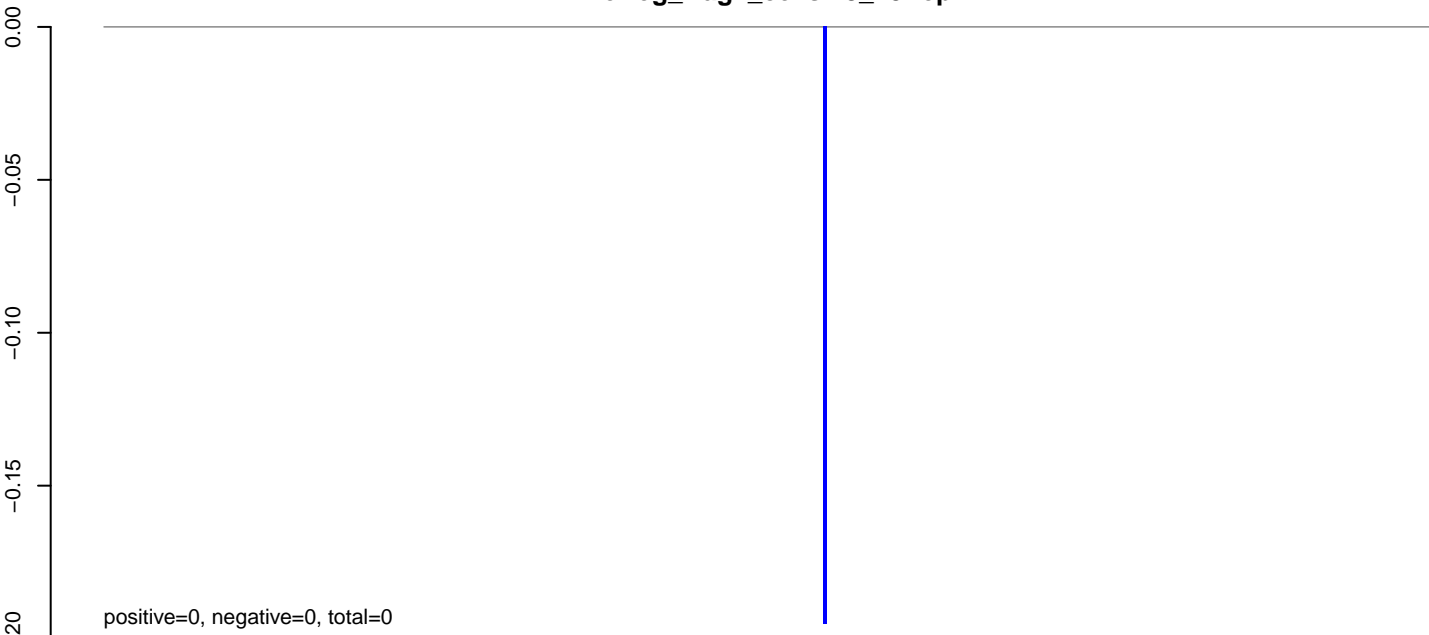


positive=2, negative=0, total=2

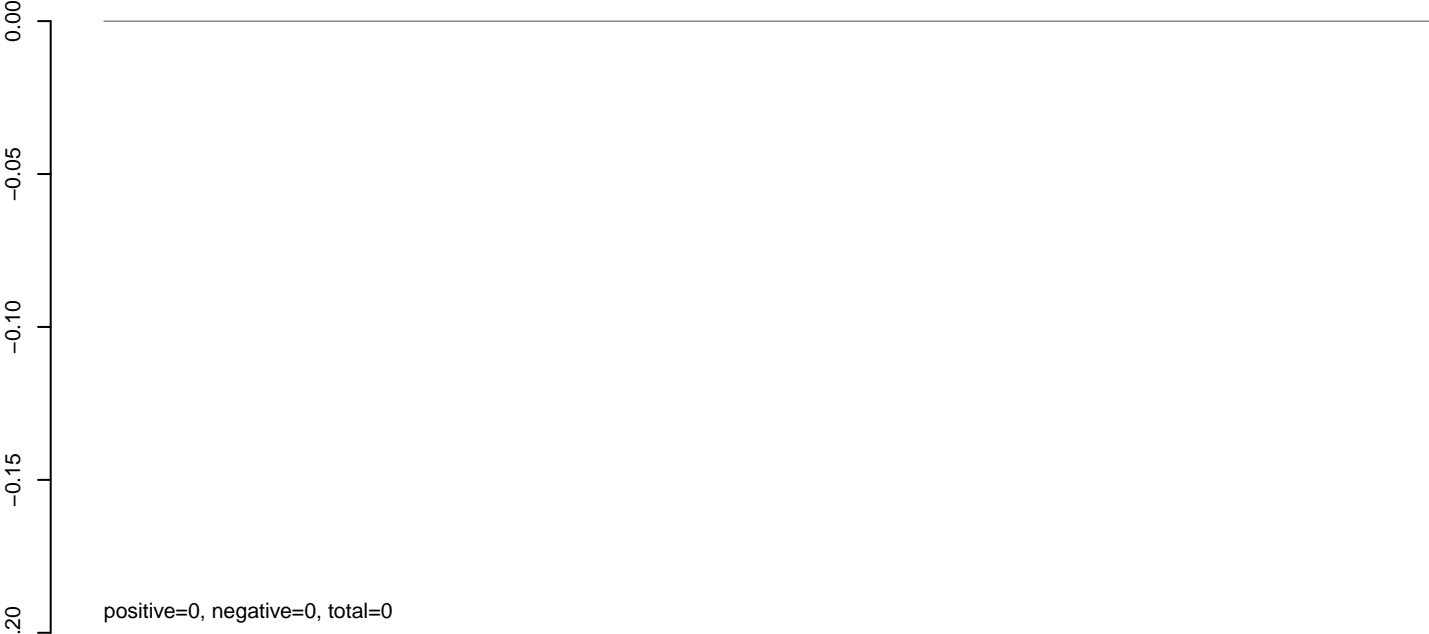
Window size=25, length=11634, virus@NC_018615.1-Eilat_virus_isolate_EO329:1-11634



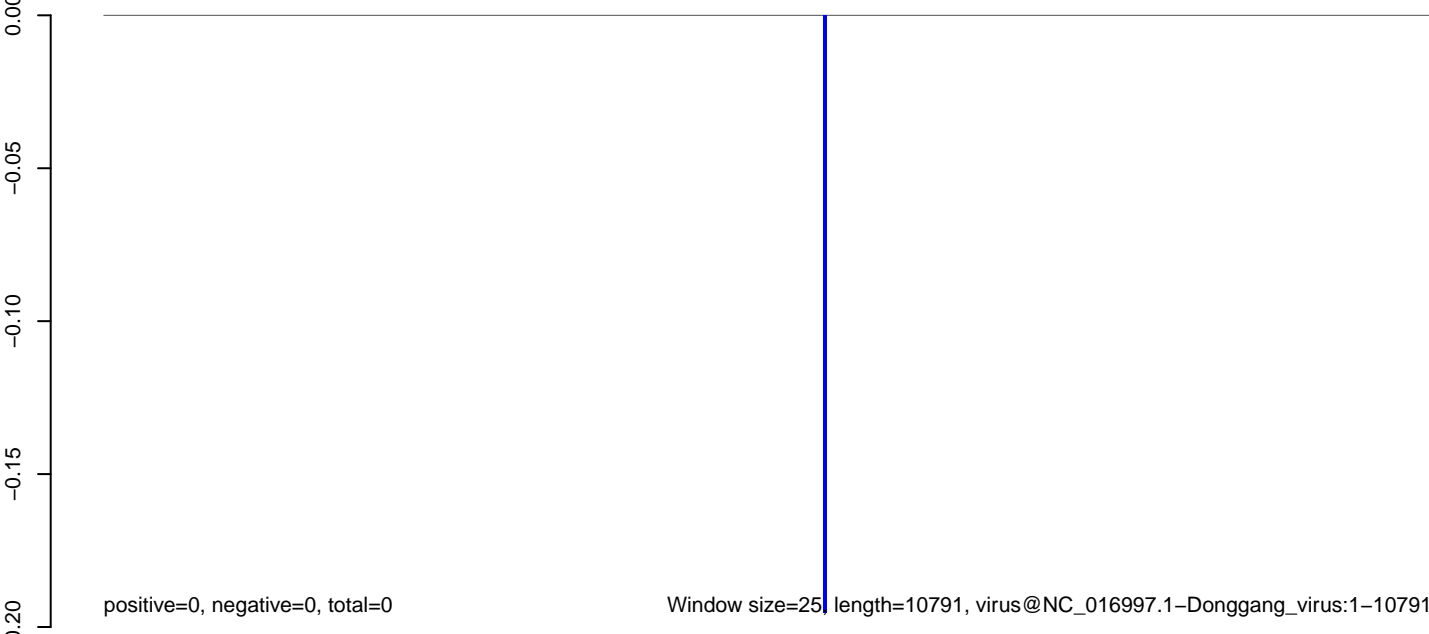
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

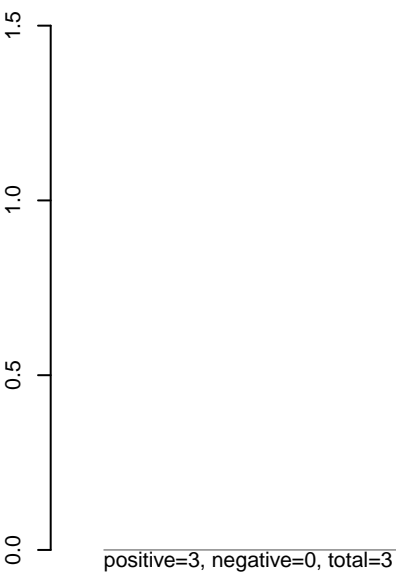


AeAeg_Aag2_cells.rep

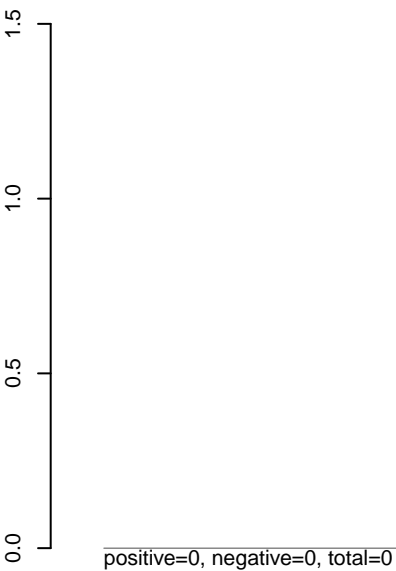


0 2000 4000 6000 8000 10000

AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

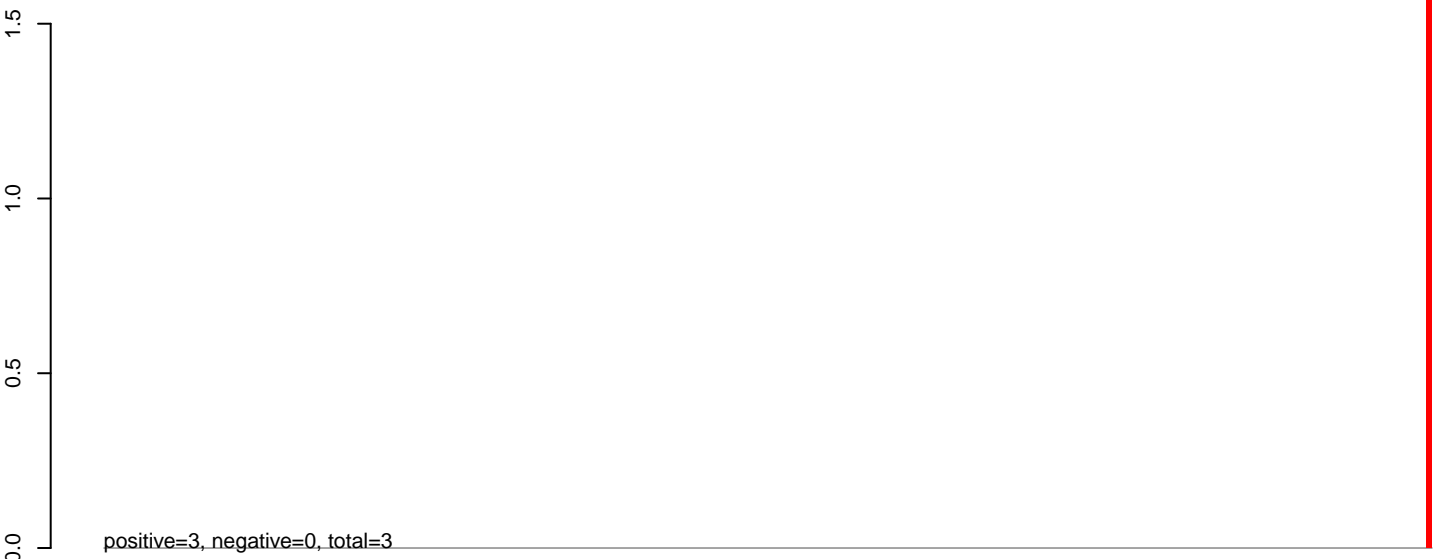


AeAeg_Aag2_cells.rep

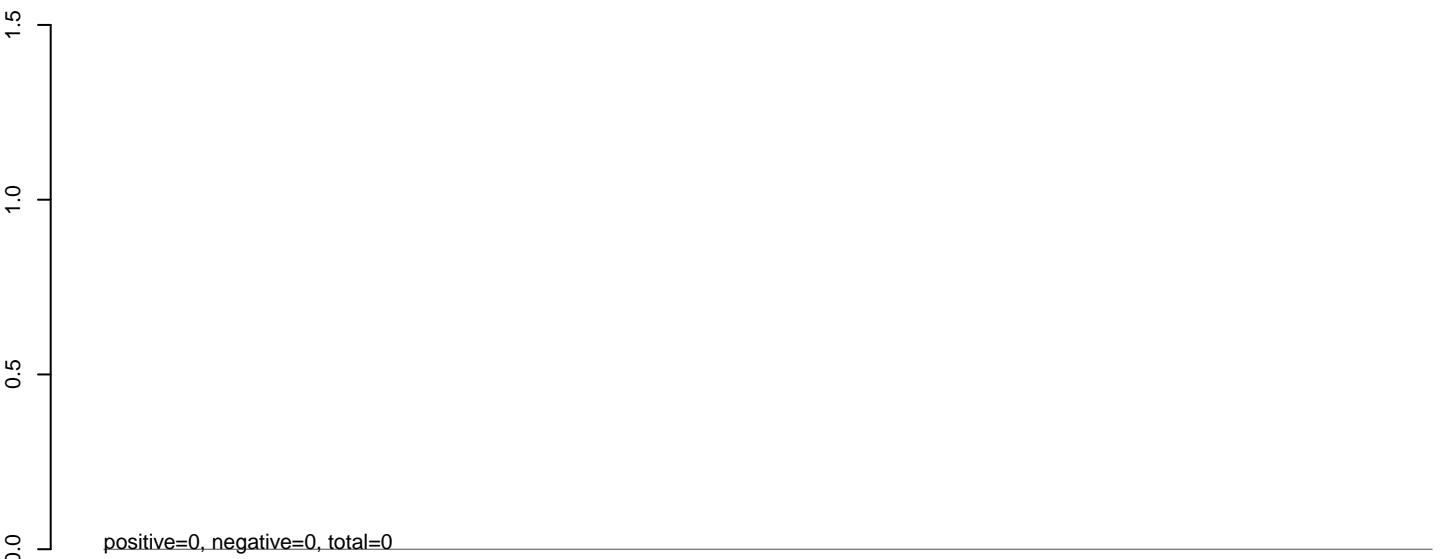


0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

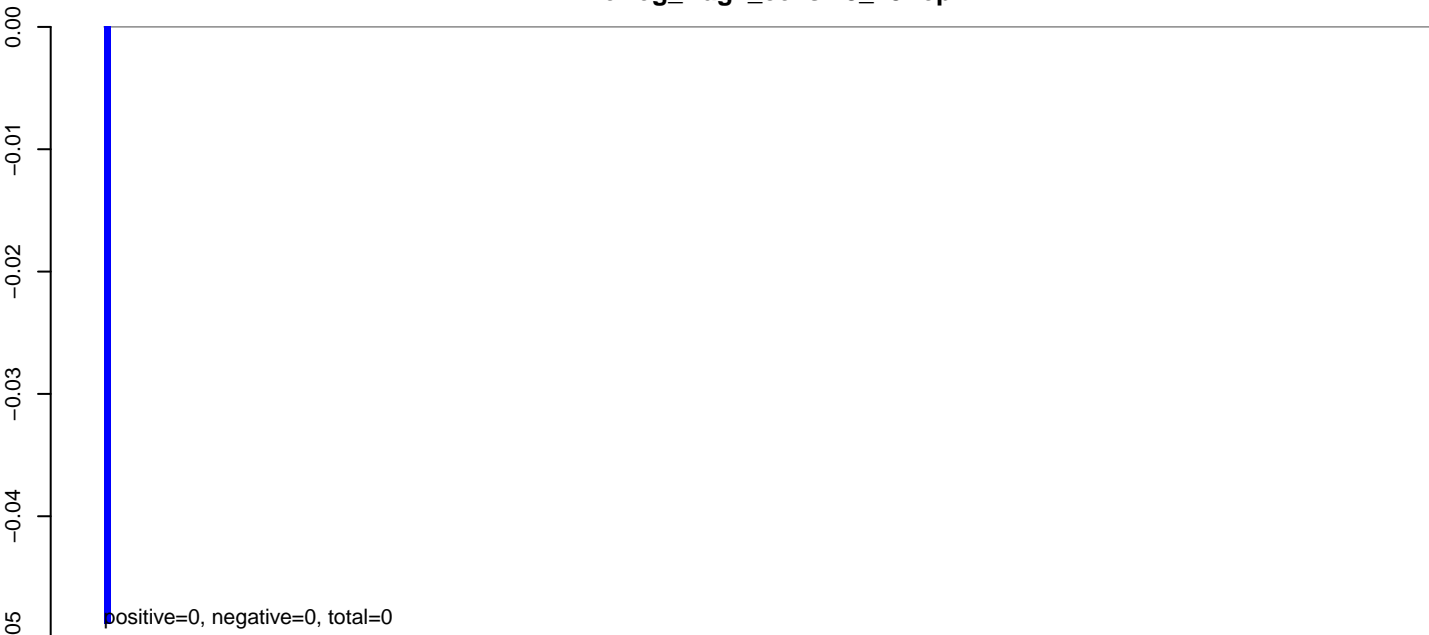


AeAeg_Aag2_cells.rep

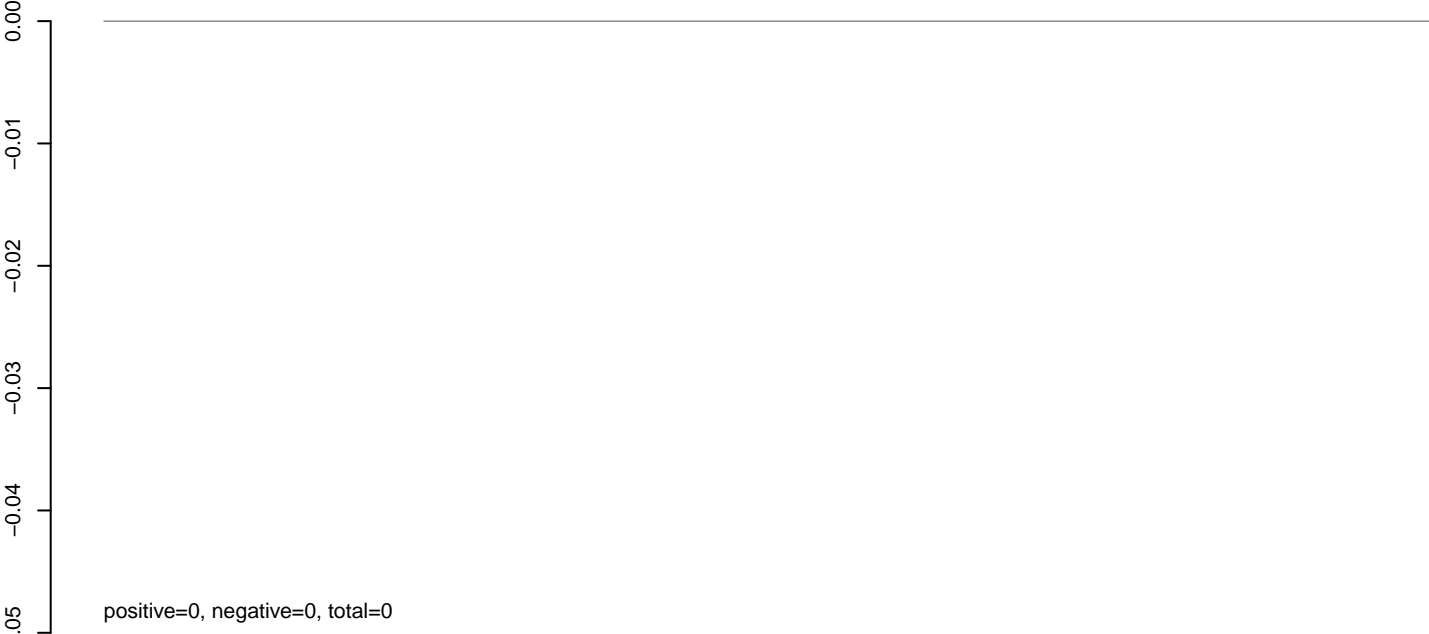


0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_cells.18_23.rep



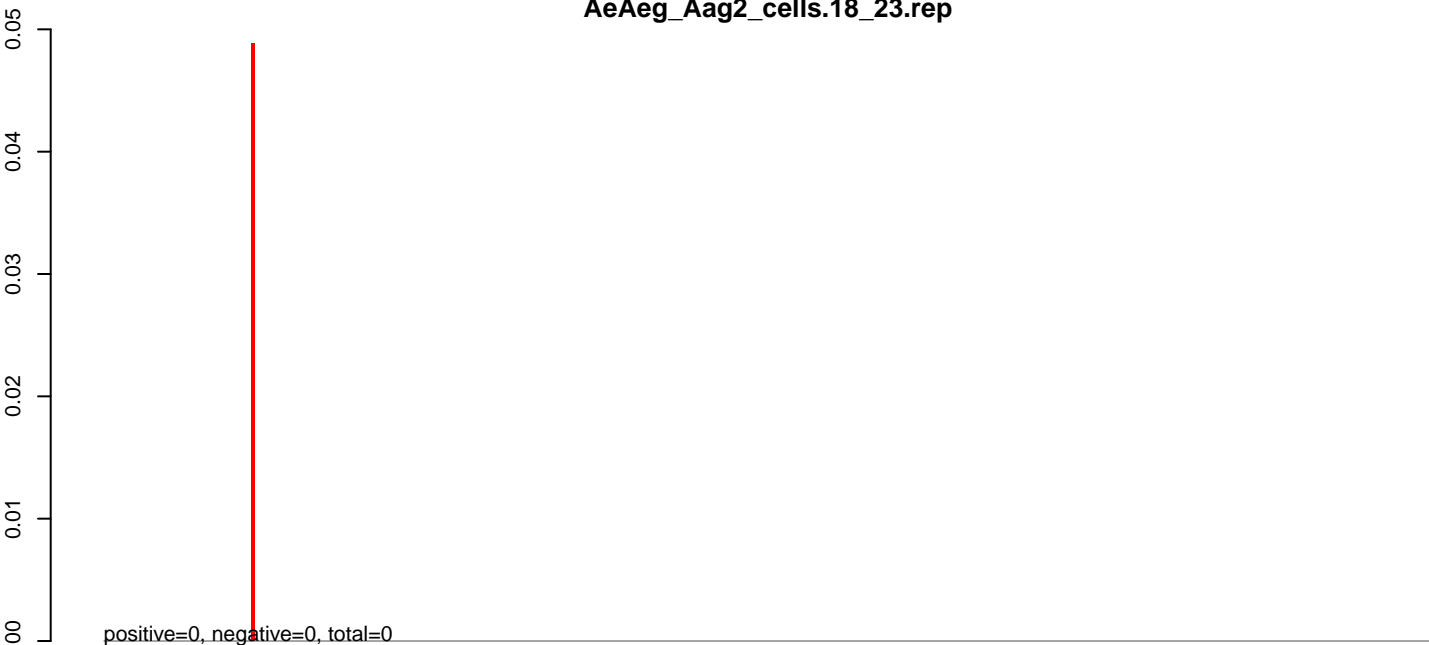
AeAeg_Aag2_cells.24_35.rep



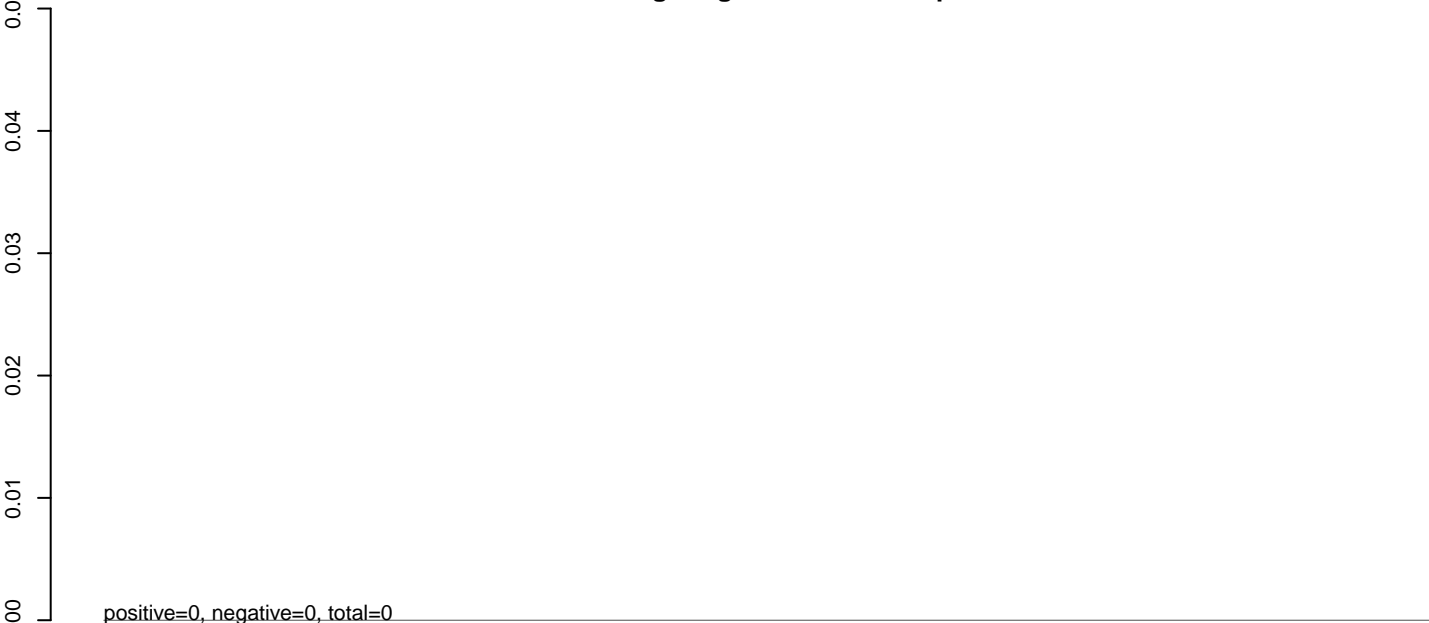
AeAeg_Aag2_cells.rep



AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

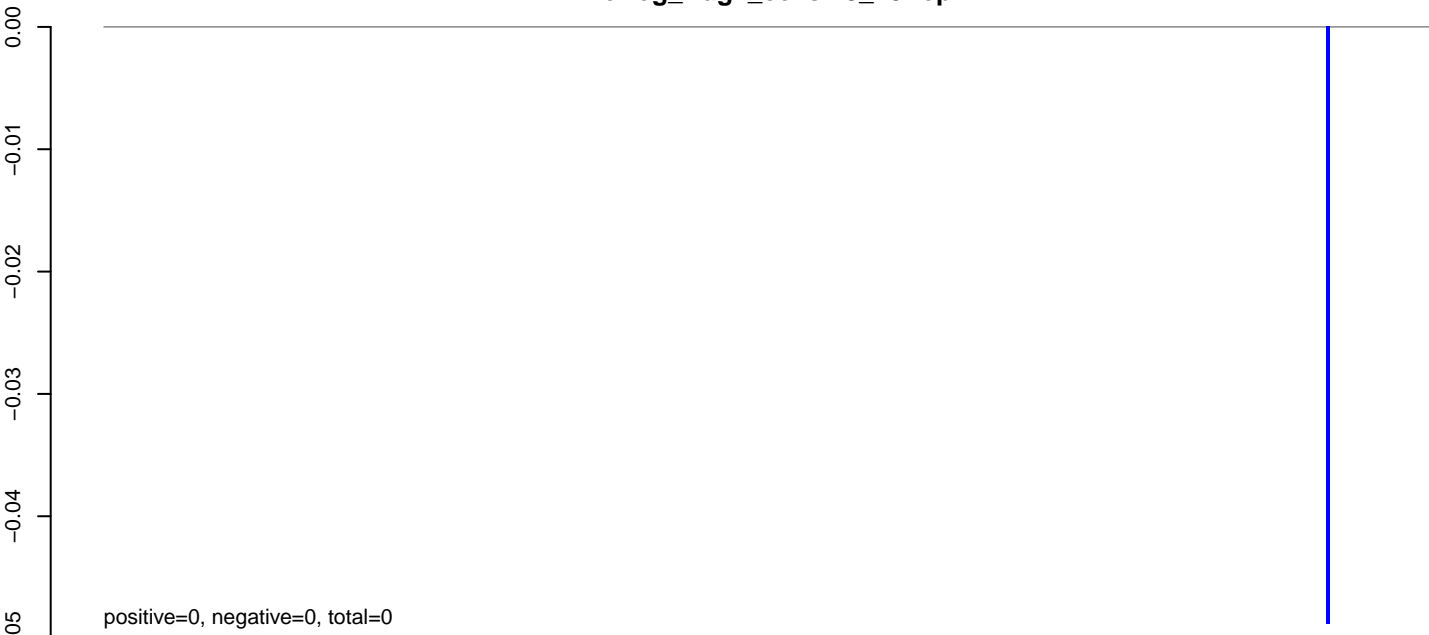


AeAeg_Aag2_cells.rep



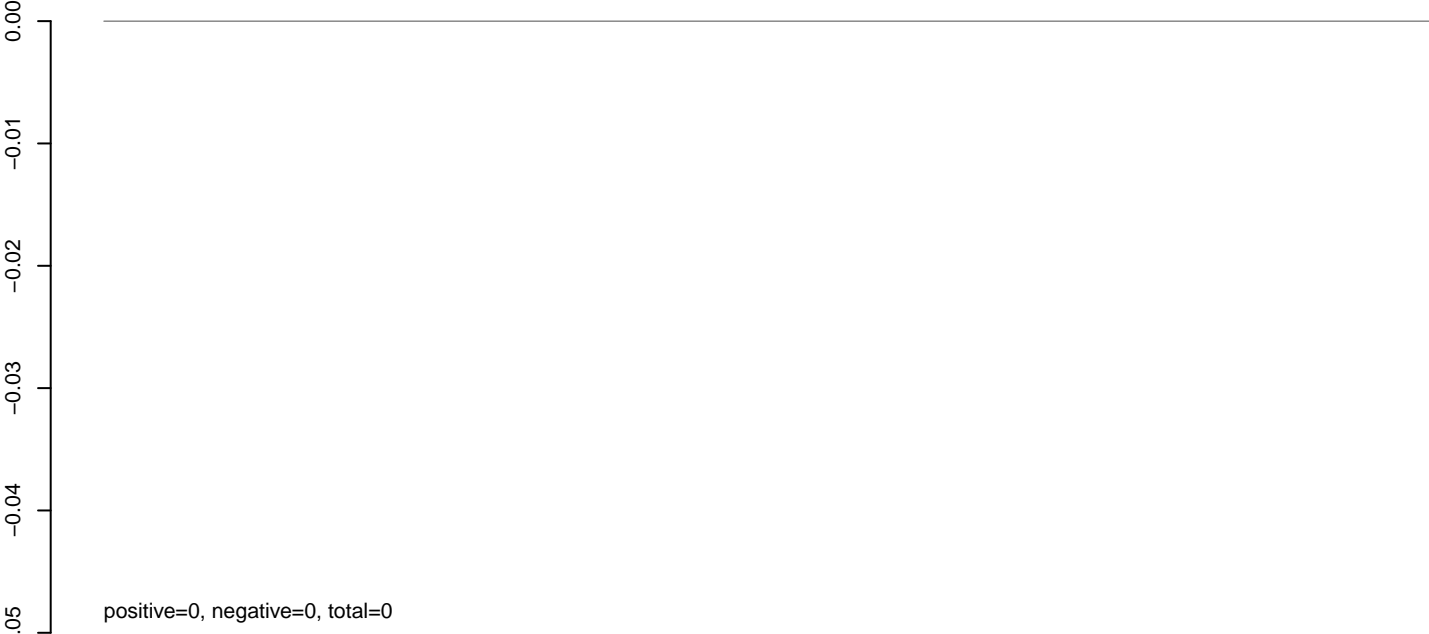
0 2000 4000 6000 8000 10000

AeAeg_Aag2_cells.18_23.rep



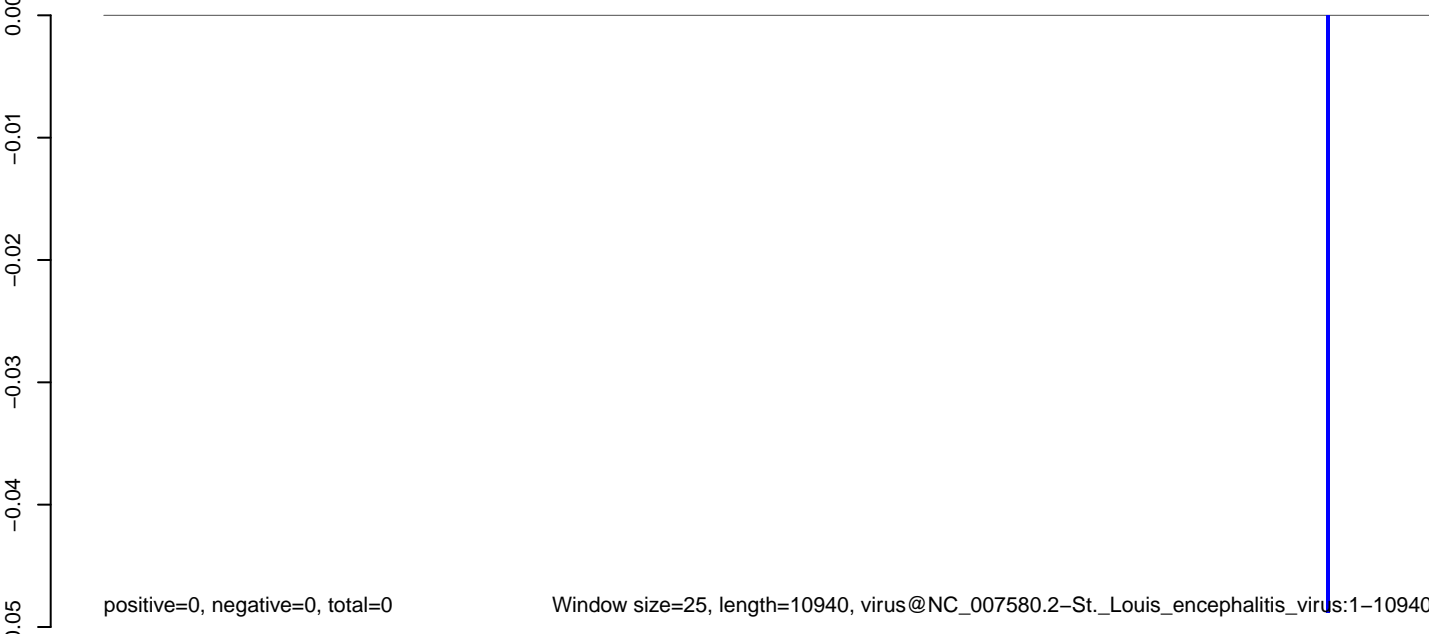
positive=0, negative=0, total=0

AeAeg_Aag2_cells.24_35.rep



positive=0, negative=0, total=0

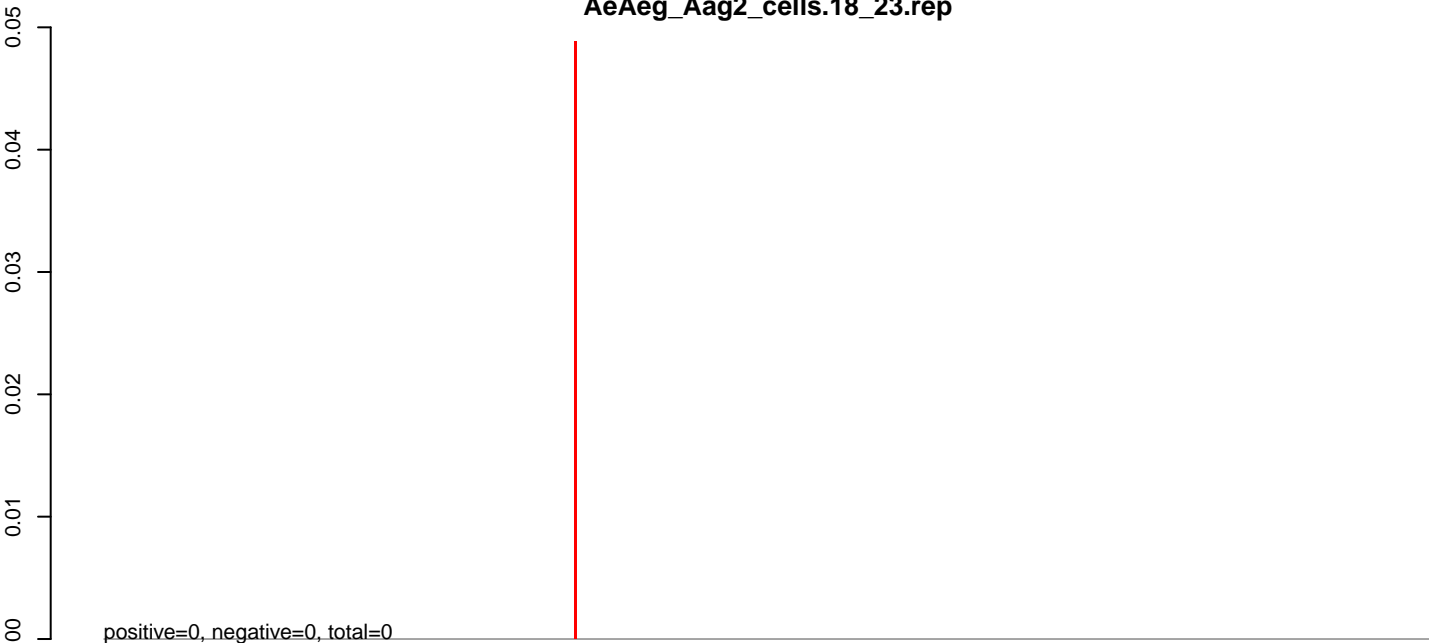
AeAeg_Aag2_cells.rep



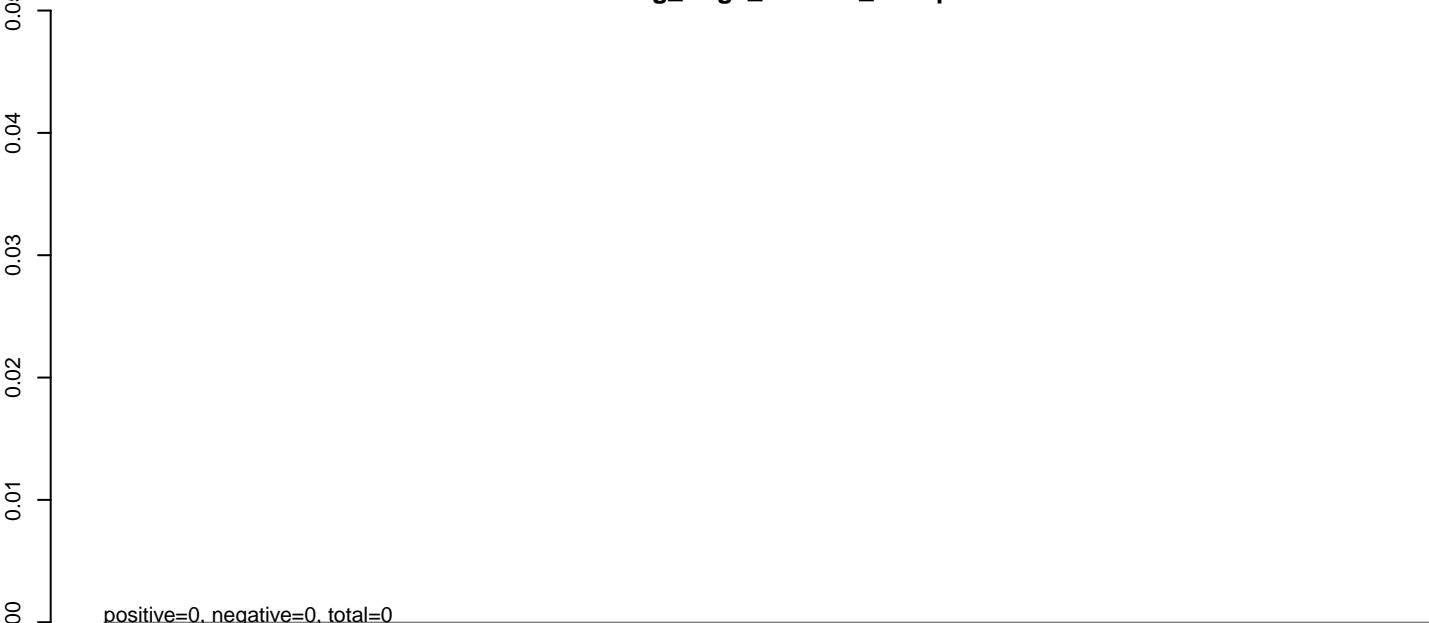
positive=0, negative=0, total=0

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

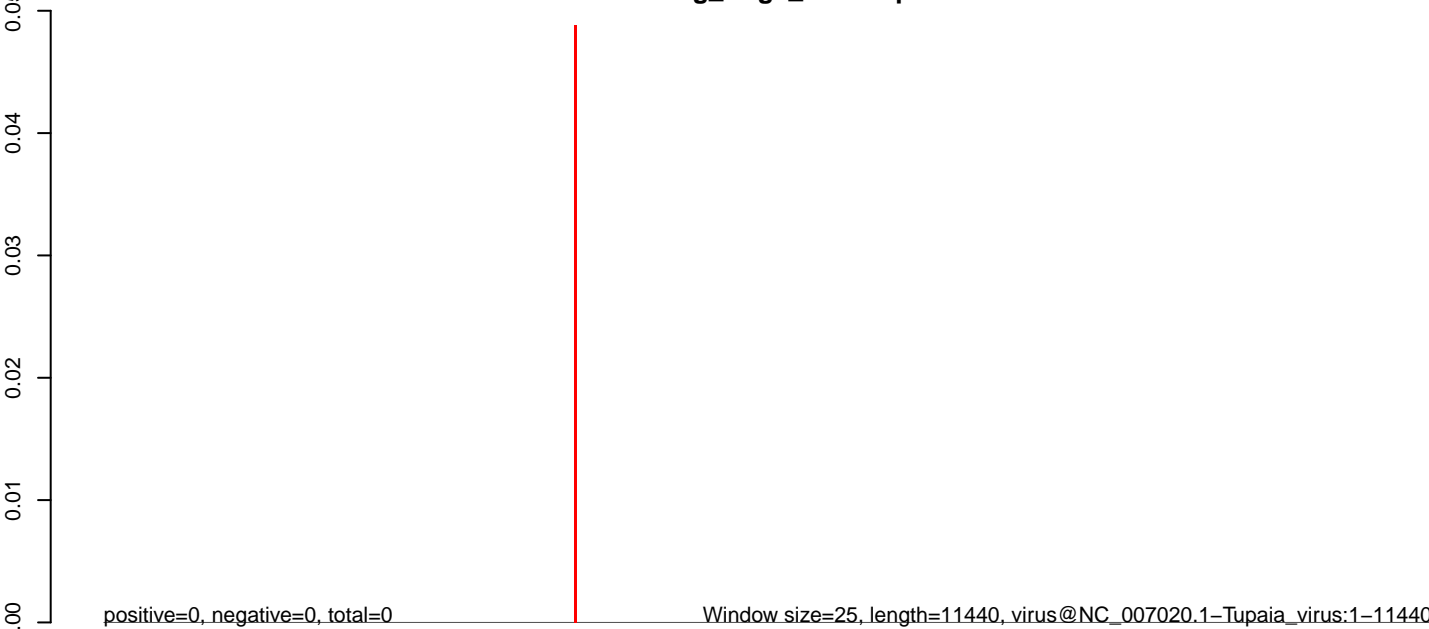
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

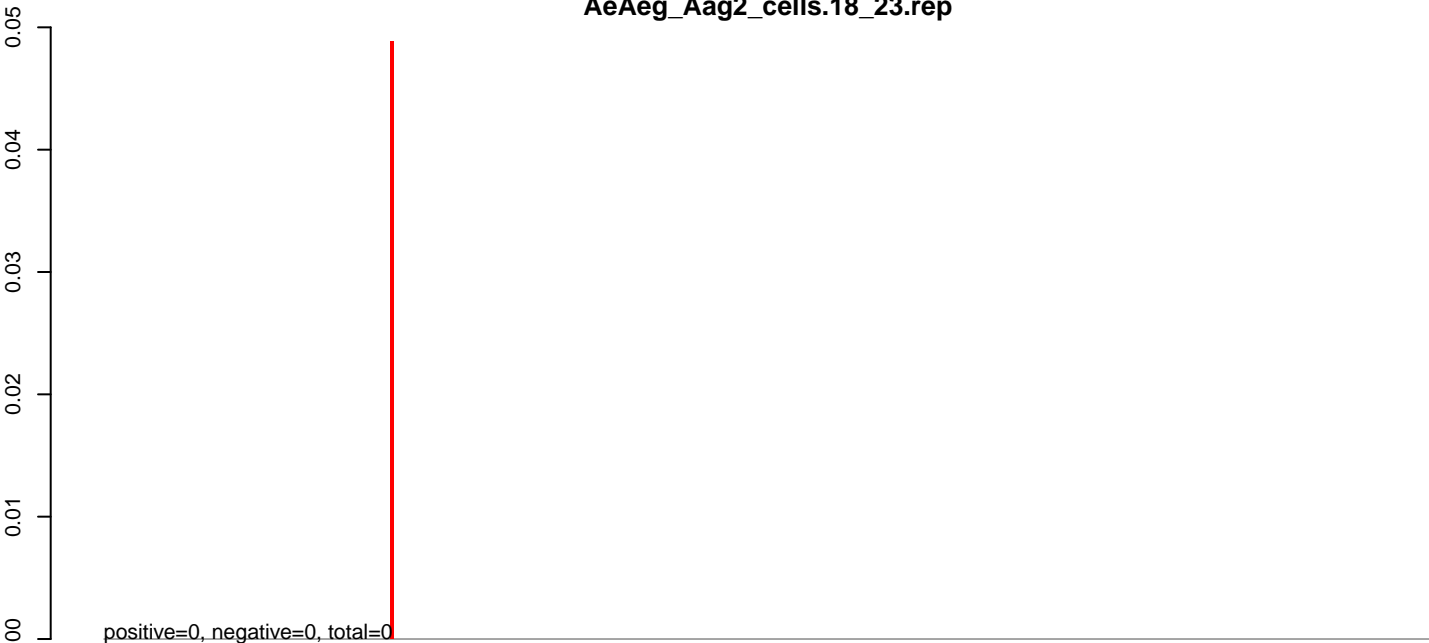


AeAeg_Aag2_cells.rep

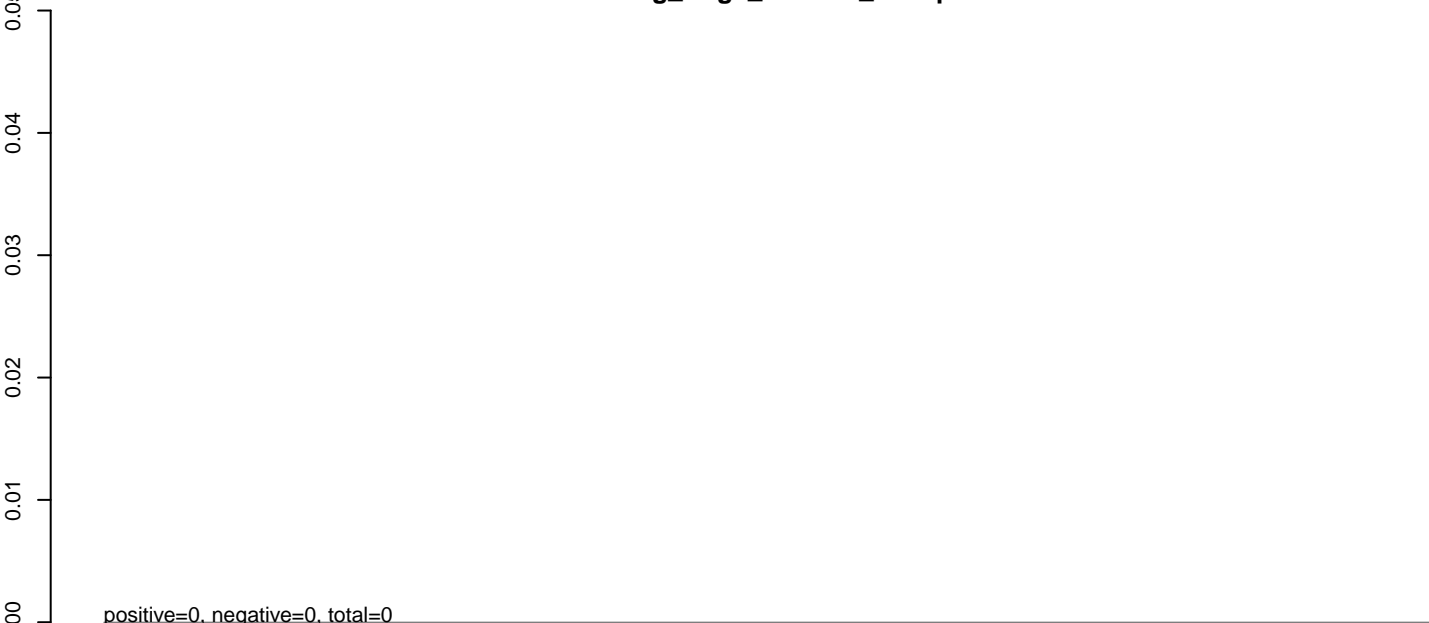


0 2000 4000 6000 8000 10000

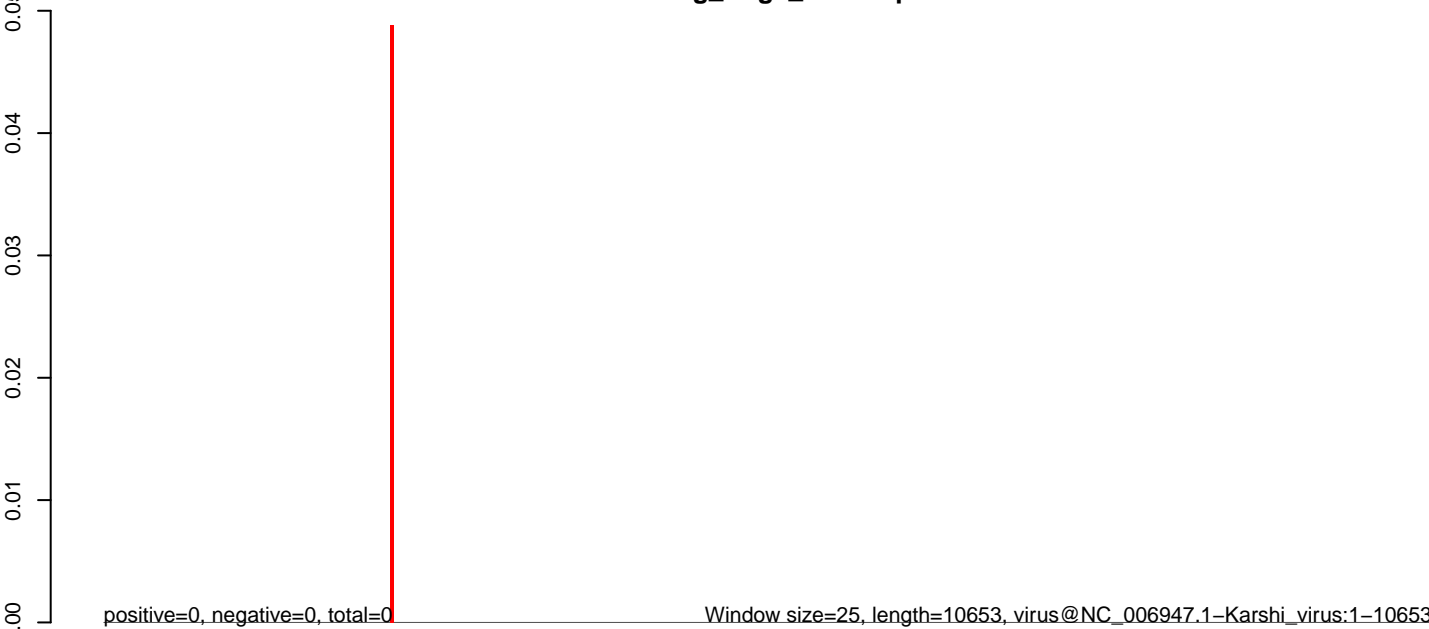
AeAeg_Aag2_cells.18_23.rep



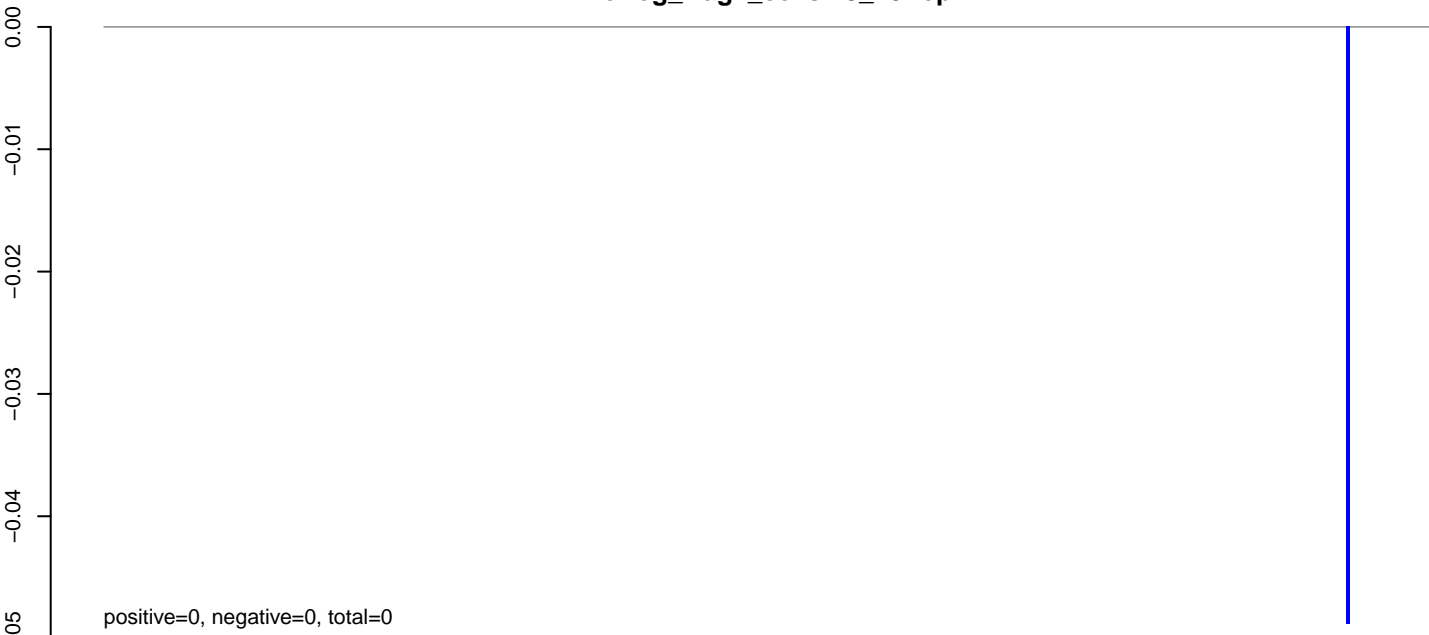
AeAeg_Aag2_cells.24_35.rep



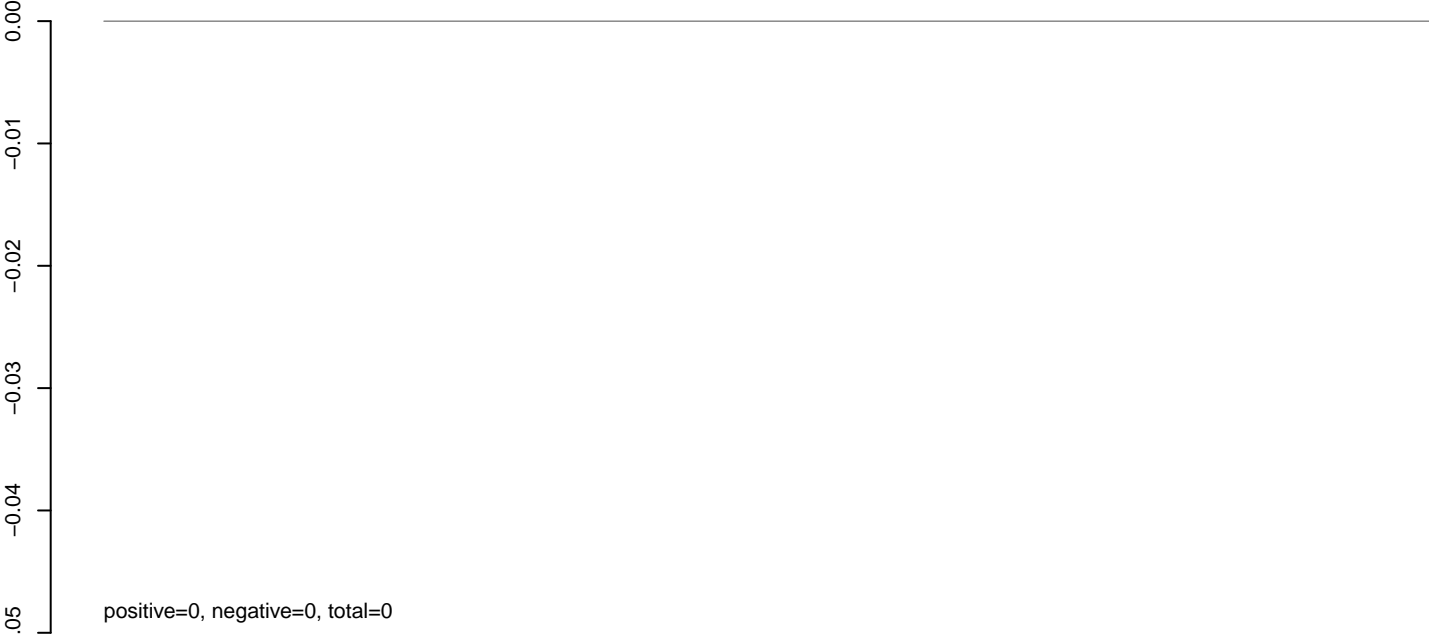
AeAeg_Aag2_cells.rep



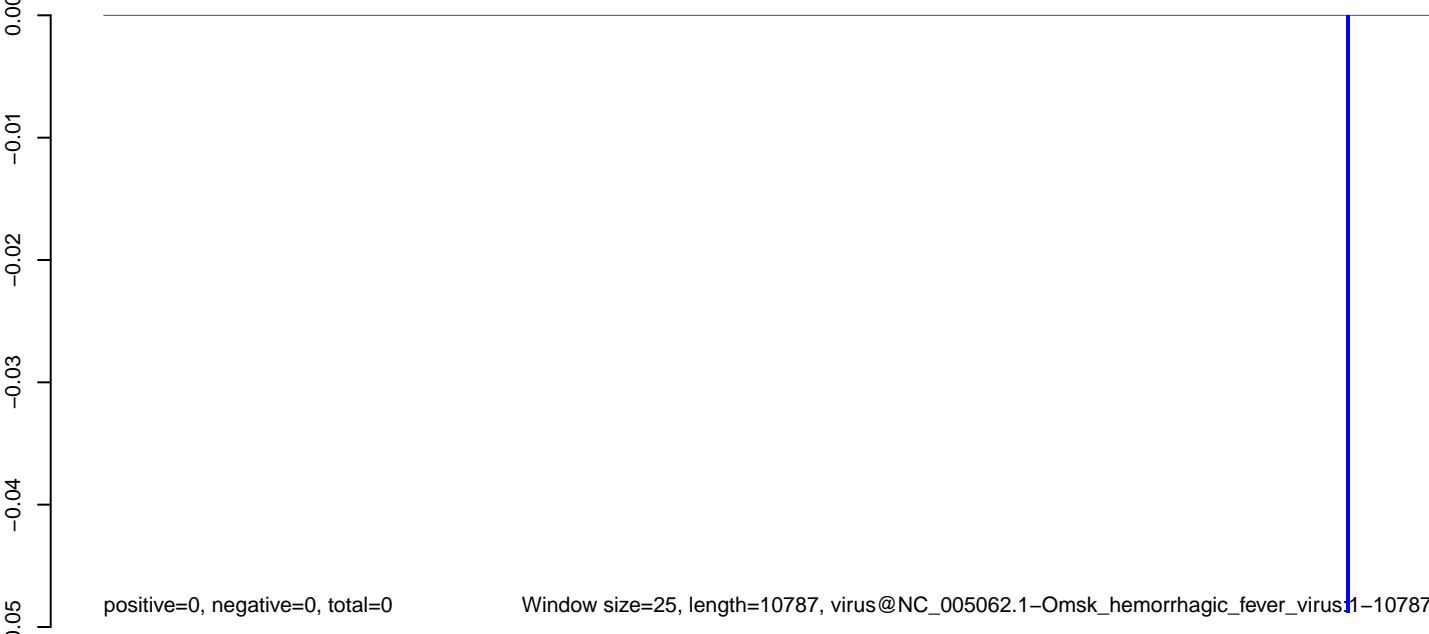
AeAeg_Aag2_cells.18_23.rep



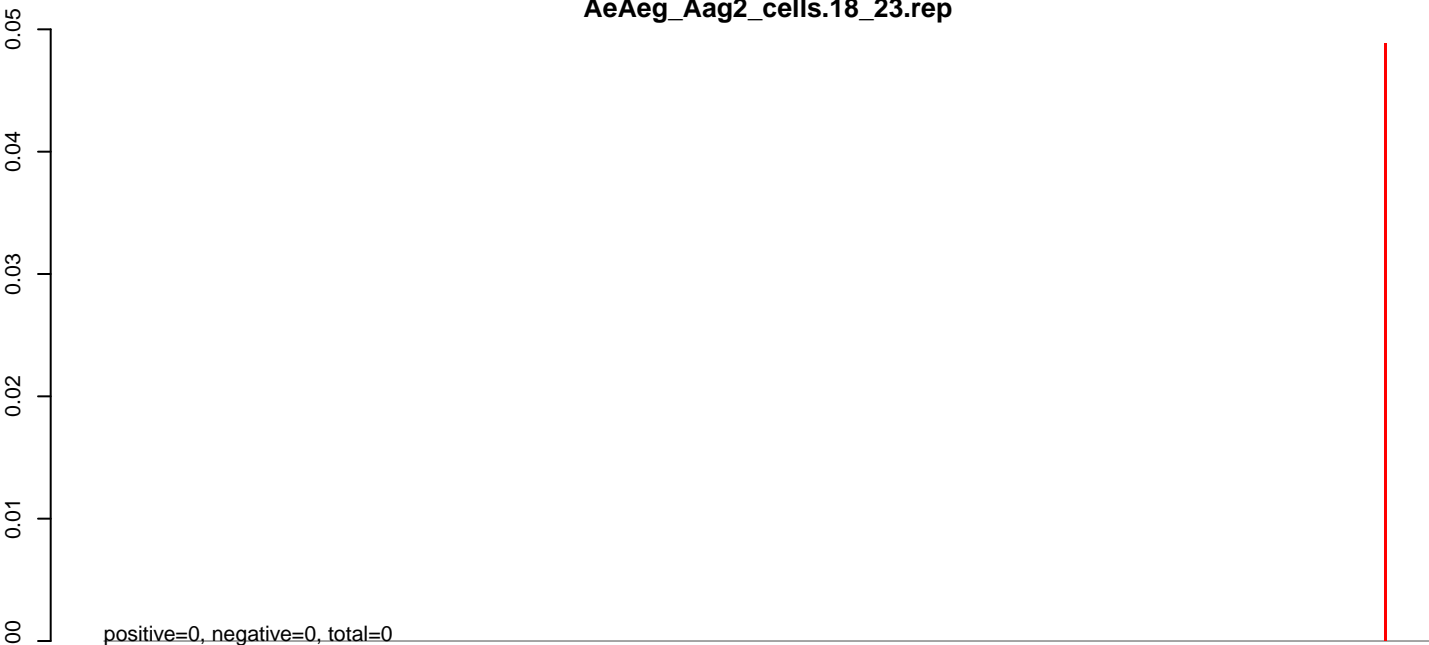
AeAeg_Aag2_cells.24_35.rep



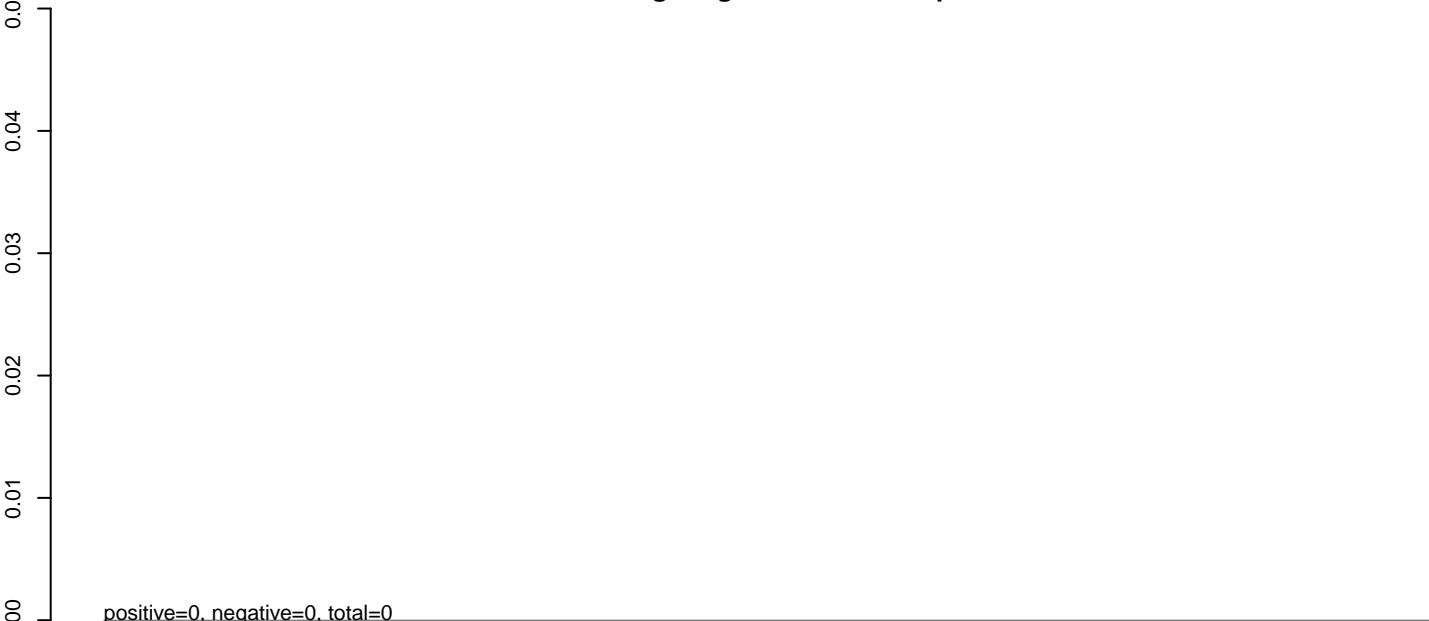
AeAeg_Aag2_cells.rep



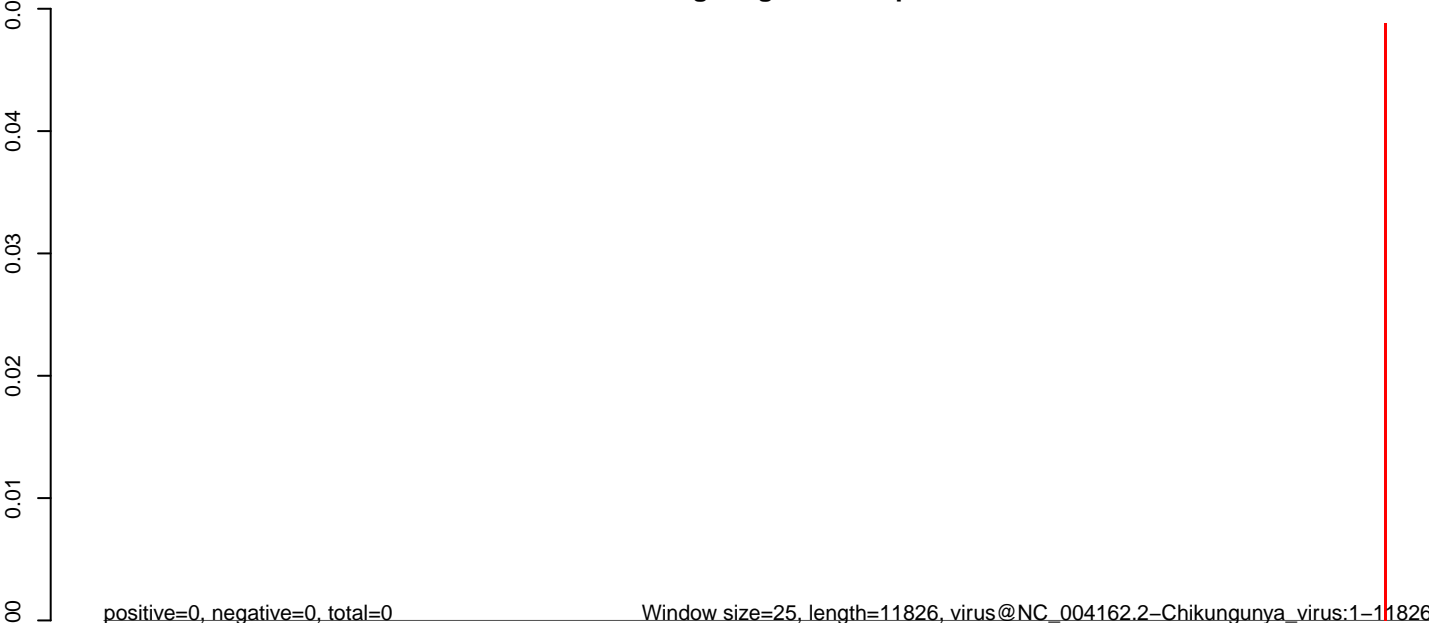
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

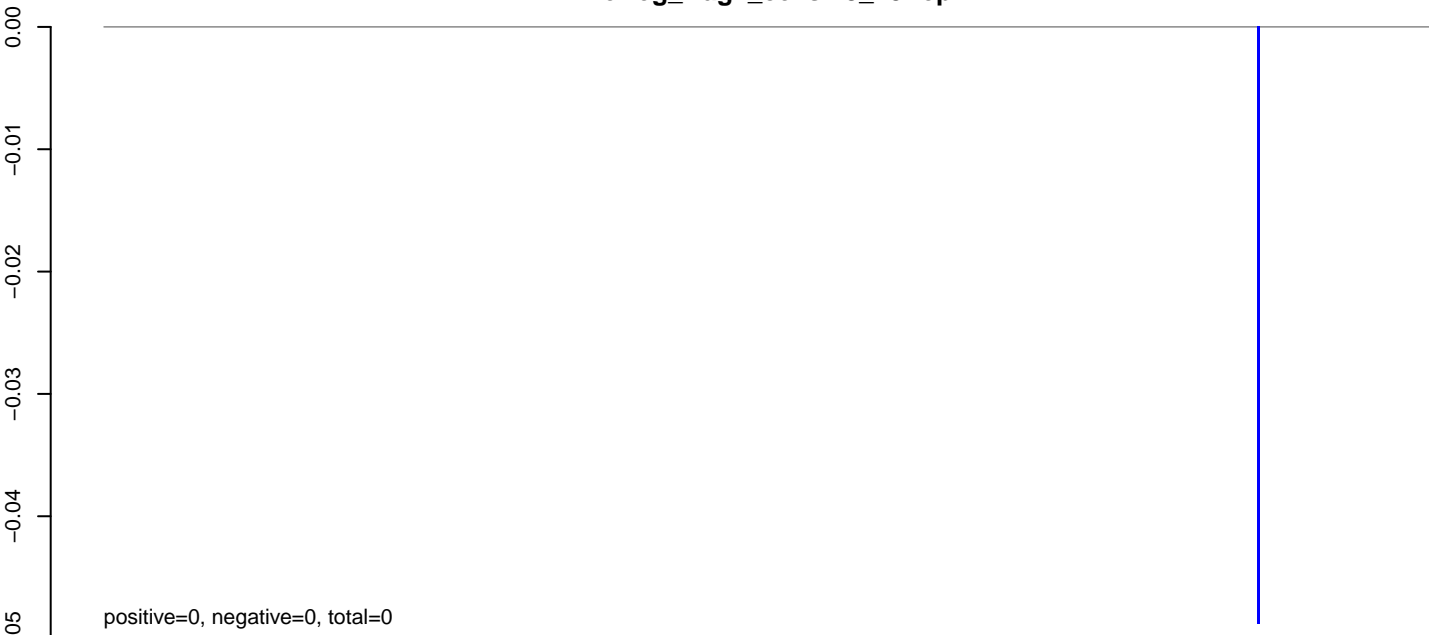


AeAeg_Aag2_cells.rep

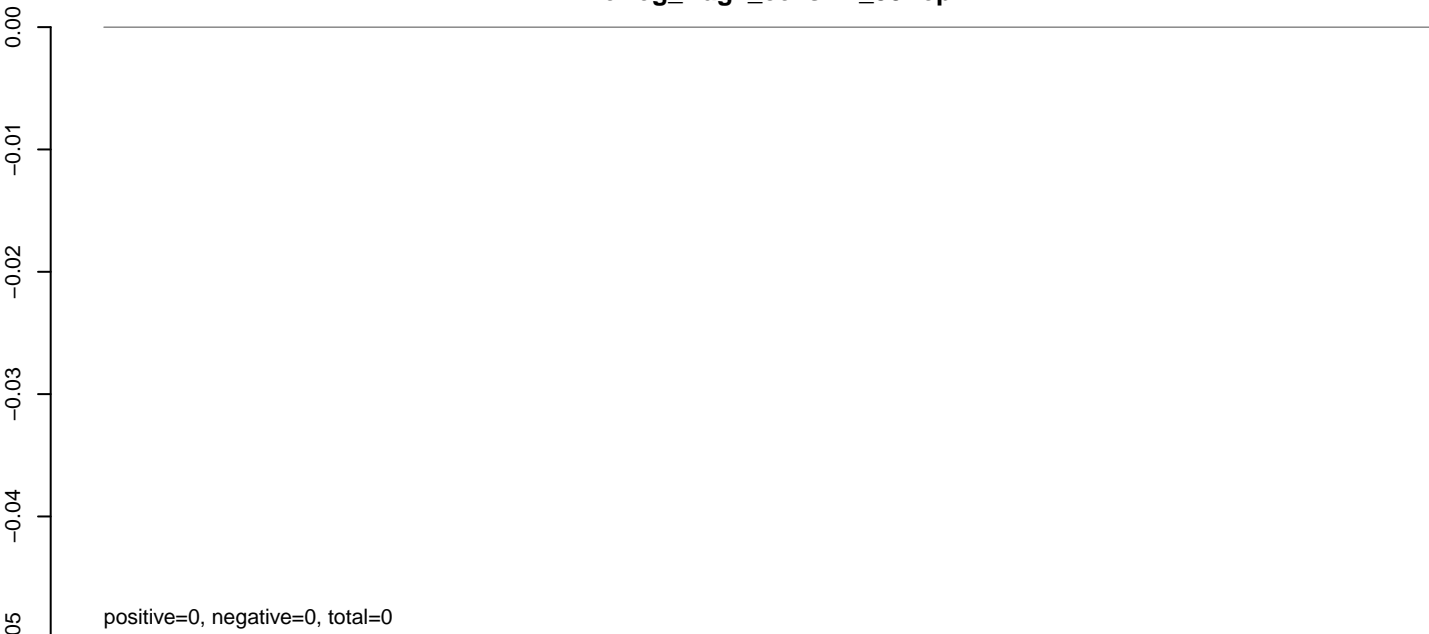


0 2000 4000 6000 8000 10000 12000

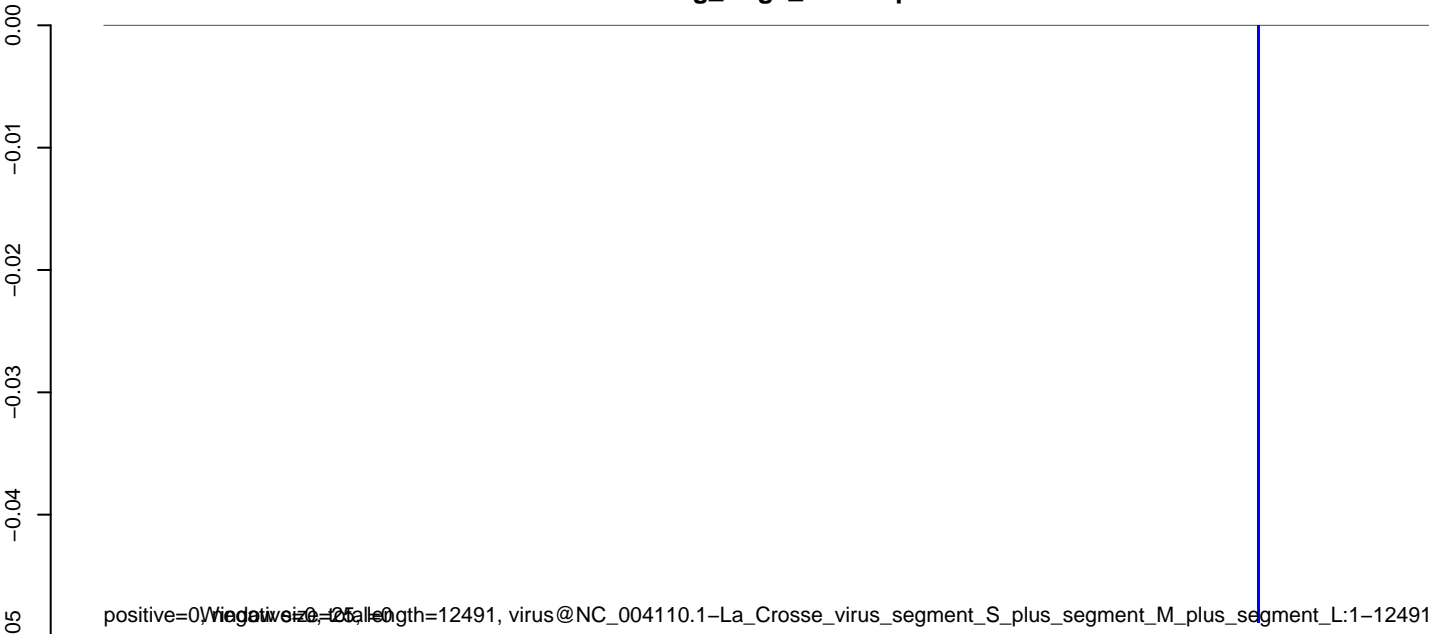
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

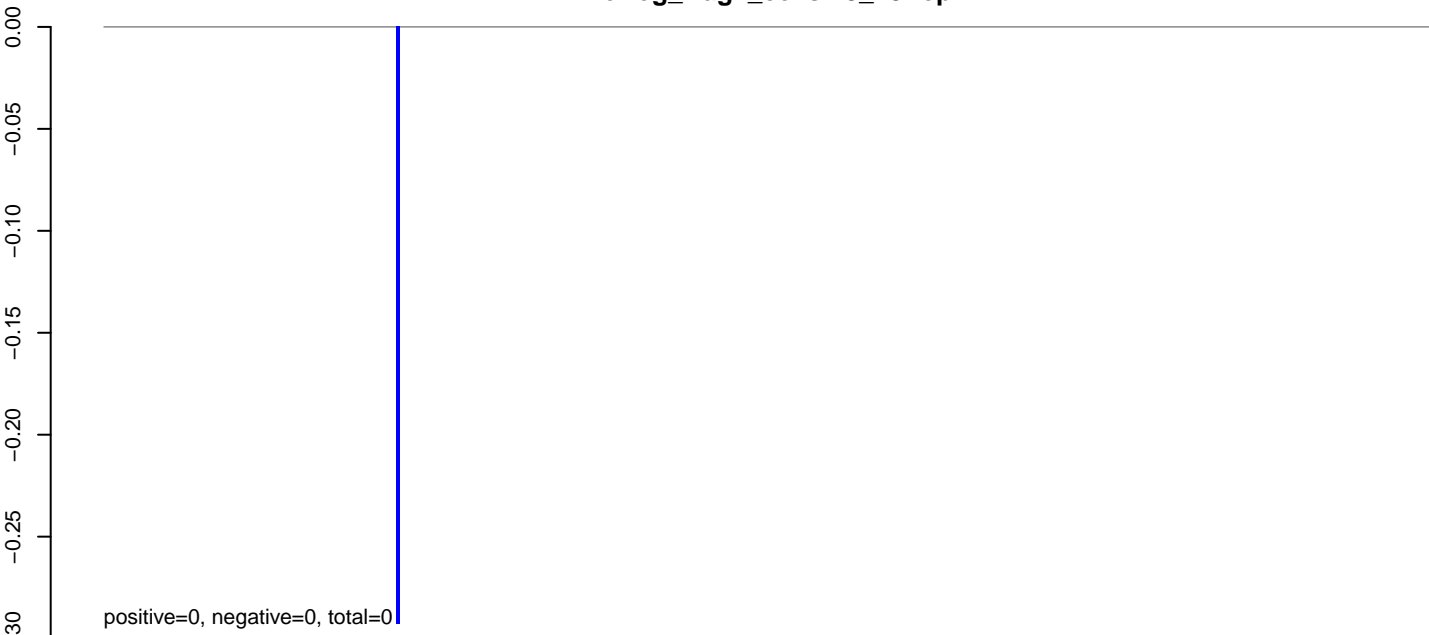


AeAeg_Aag2_cells.rep

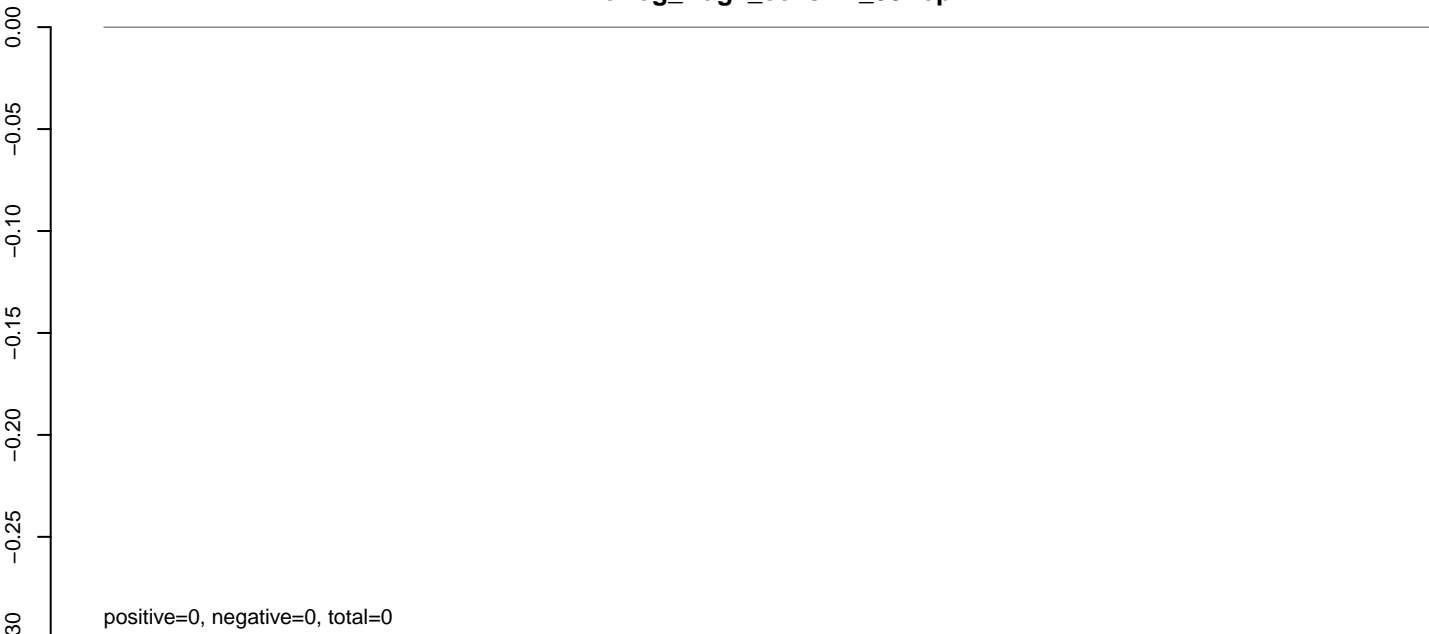


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positive=0, negative=0, total=0, virus@NC_004110.1-La_Crosse_virus_segment_S_plus_segment_M_plus_segment_L:1-12491

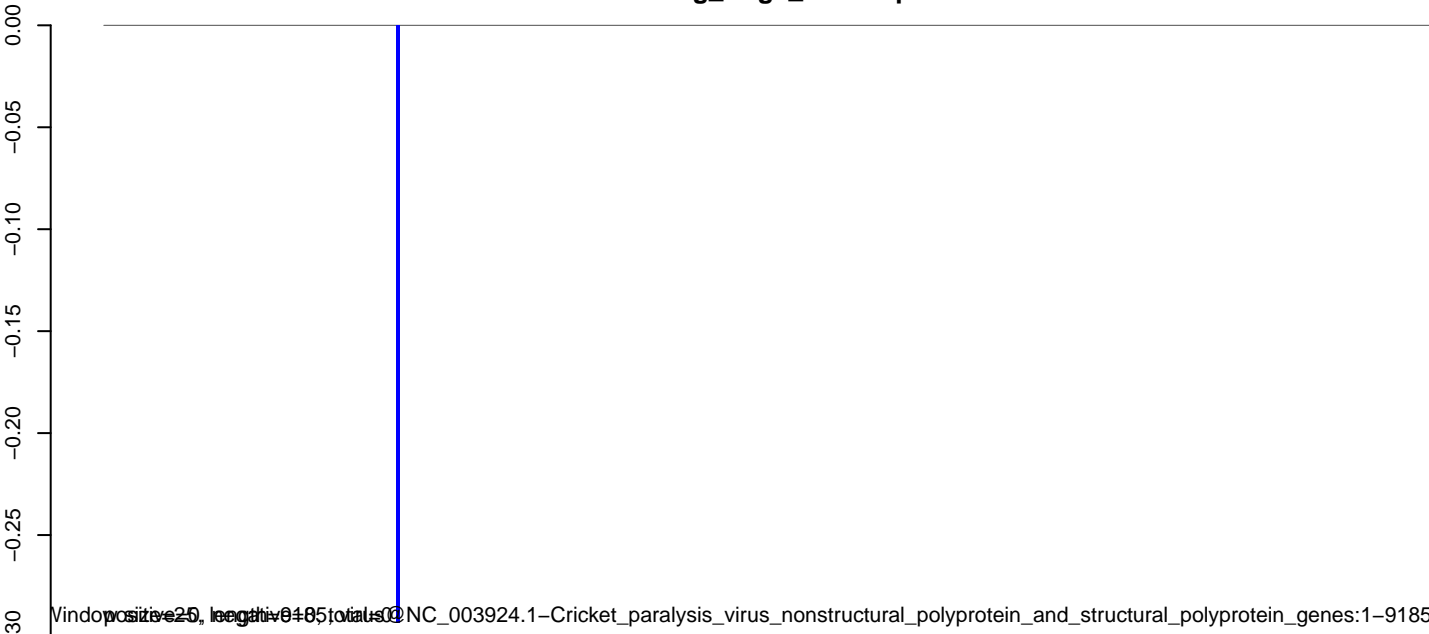
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

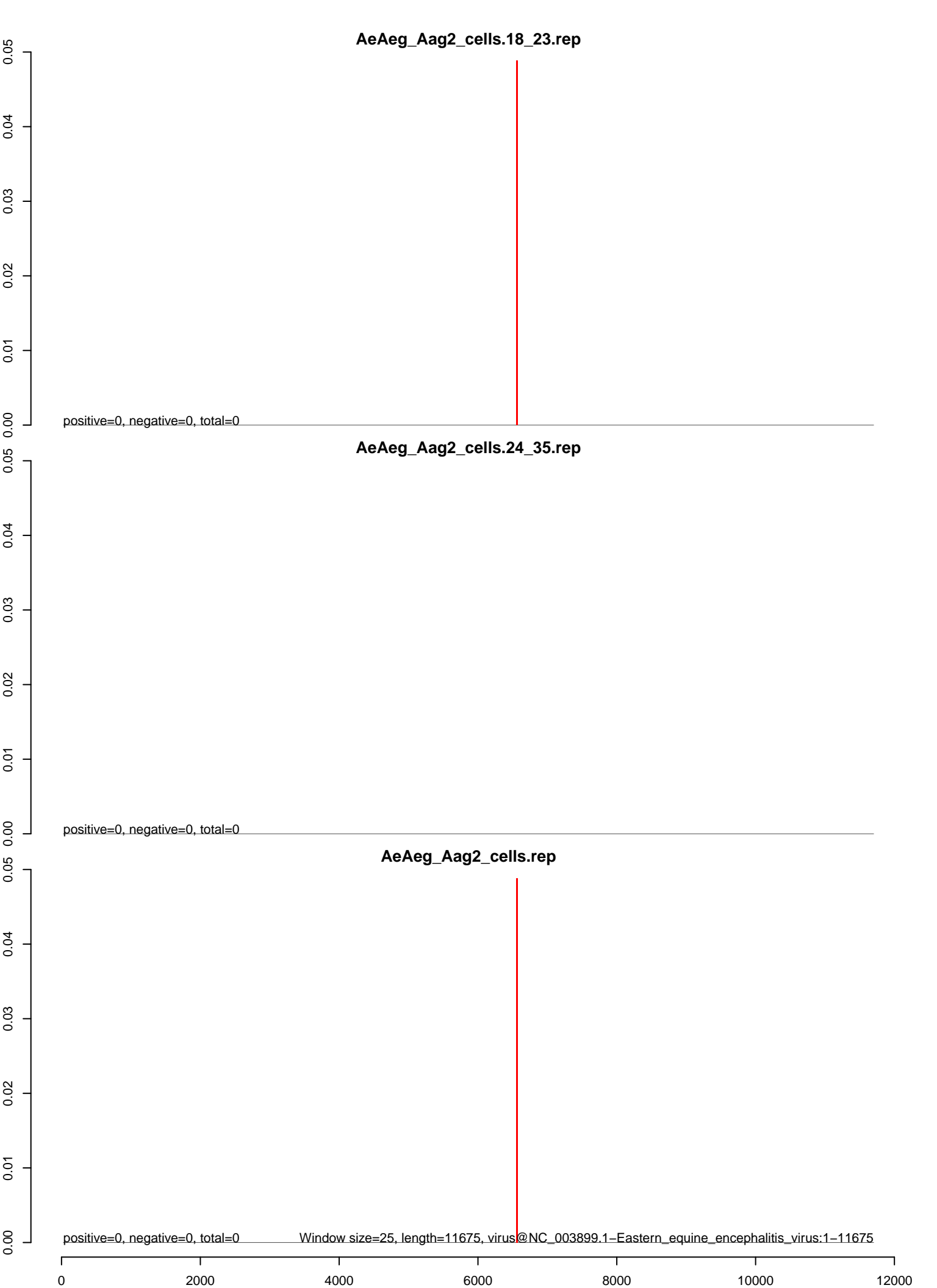


AeAeg_Aag2_cells.rep

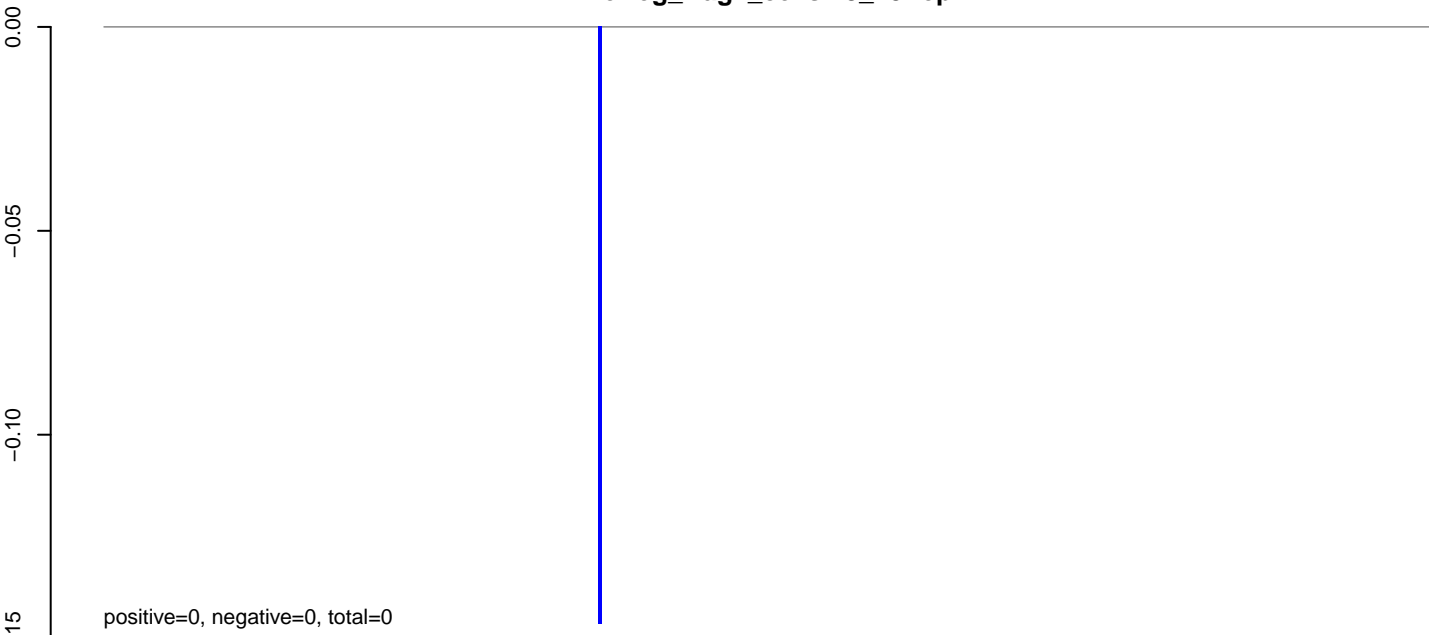


Window size: 25, length: 6485, total: 0 NC_003924.1-Cricket_paralysis_virus_nonstructural_polyprotein_and_structural_polyprotein_genes:1-9185

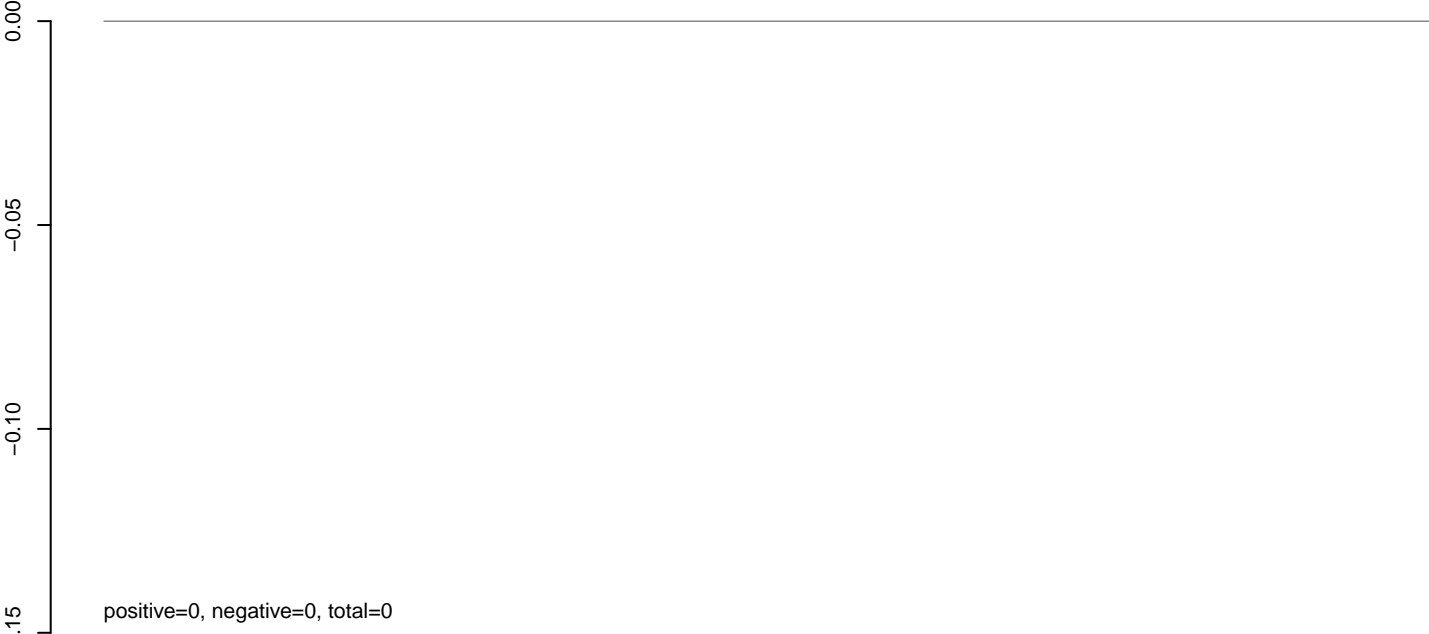
0 2000 4000 6000 8000



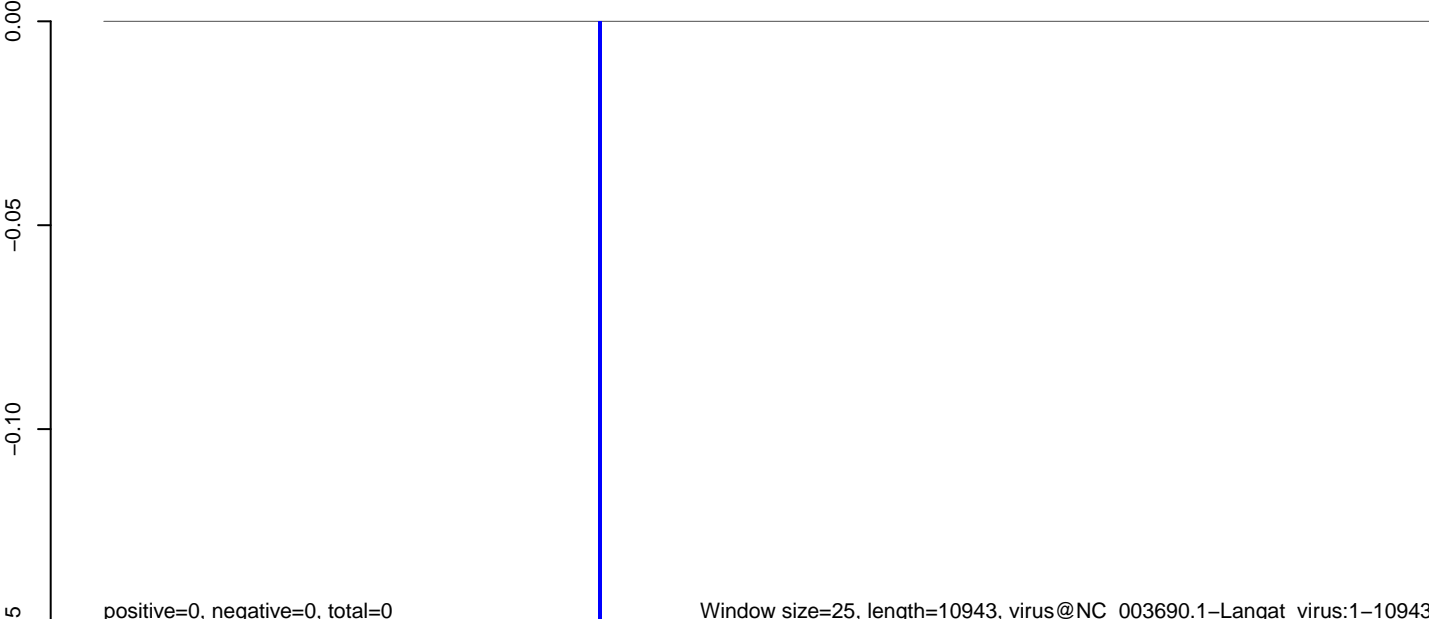
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



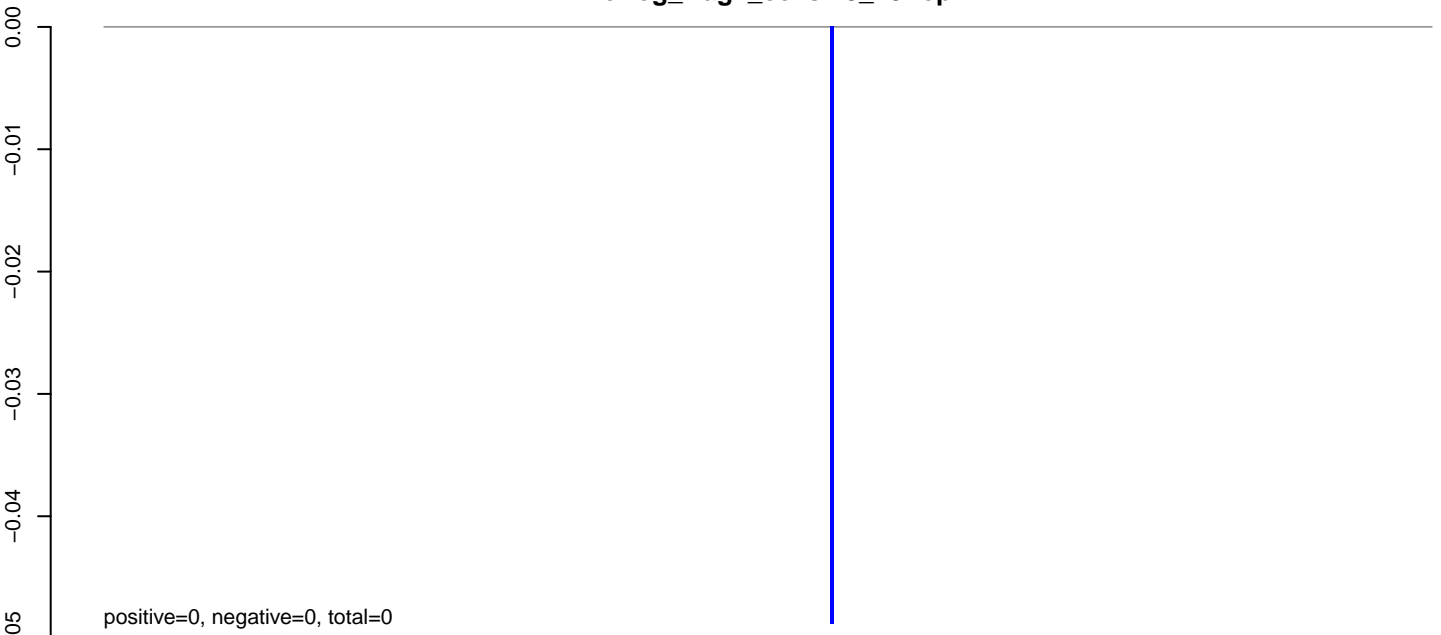
AeAeg_Aag2_cells.rep



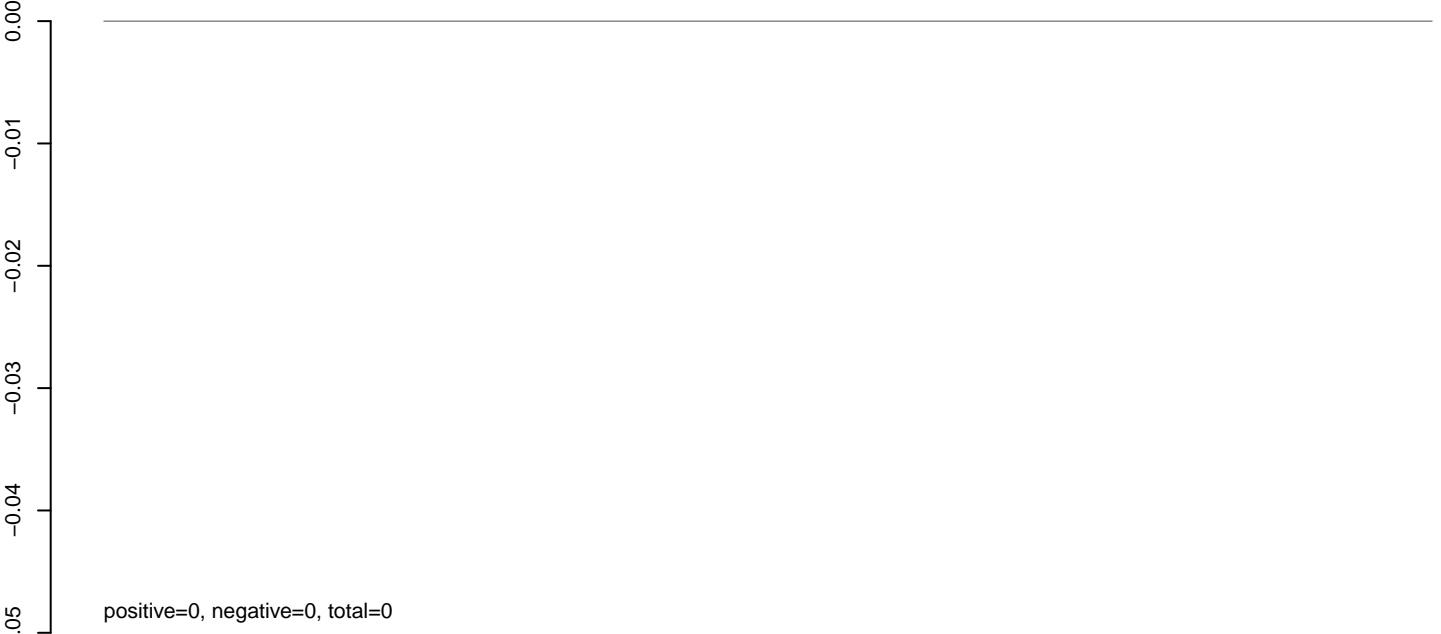
Window size=25, length=10943, virus@NC_003690.1-Langat_virus:1-10943

0 2000 4000 6000 8000 10000

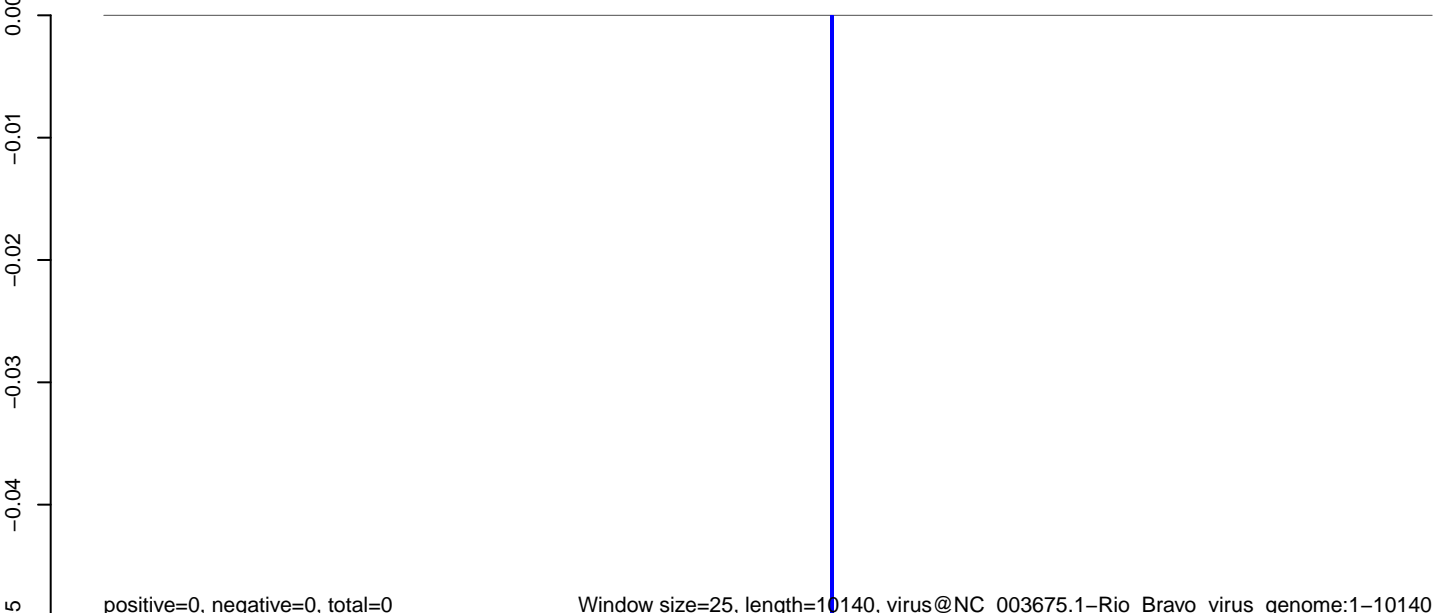
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



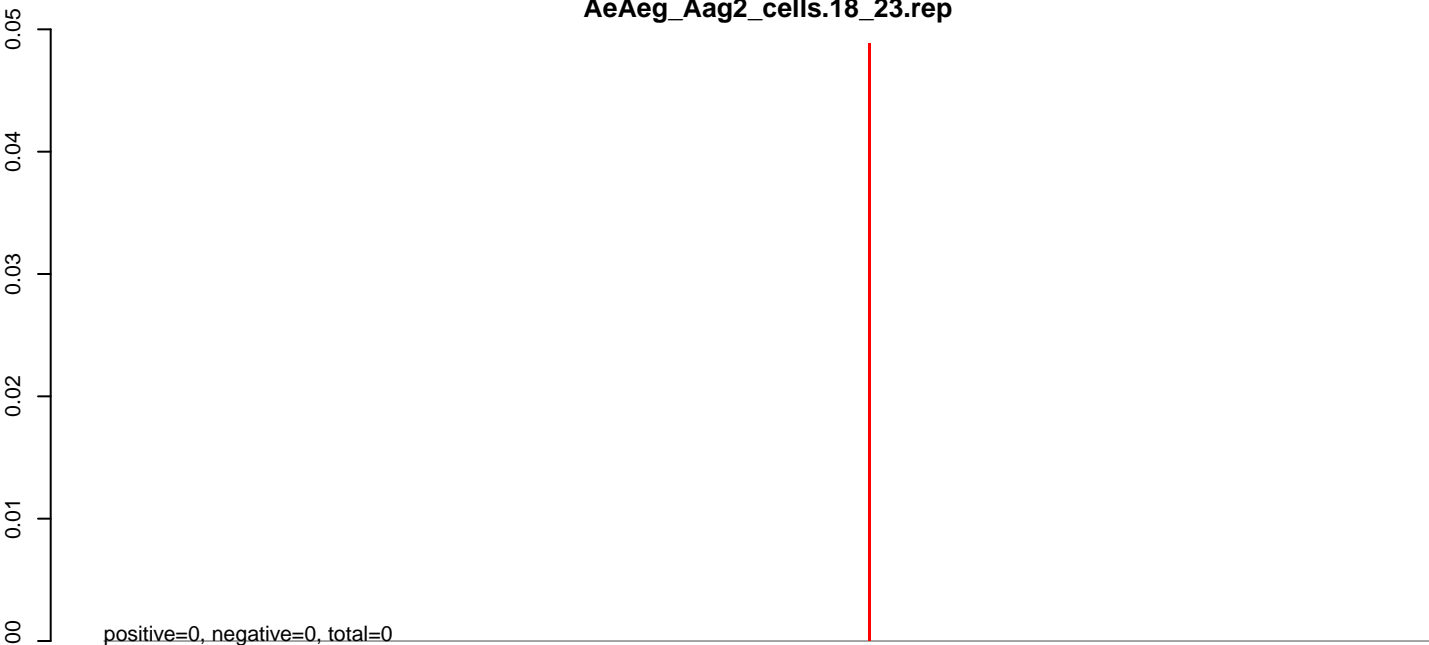
AeAeg_Aag2_cells.rep



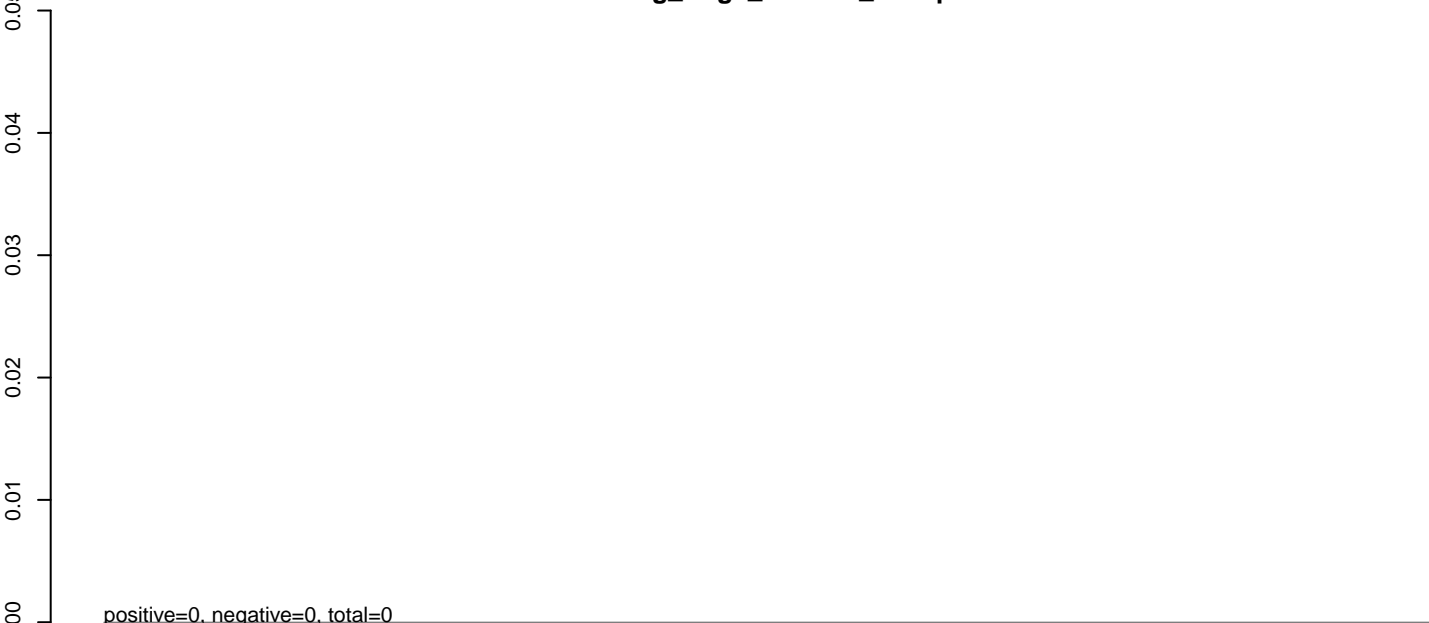
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0 2000 4000 6000 8000 10000

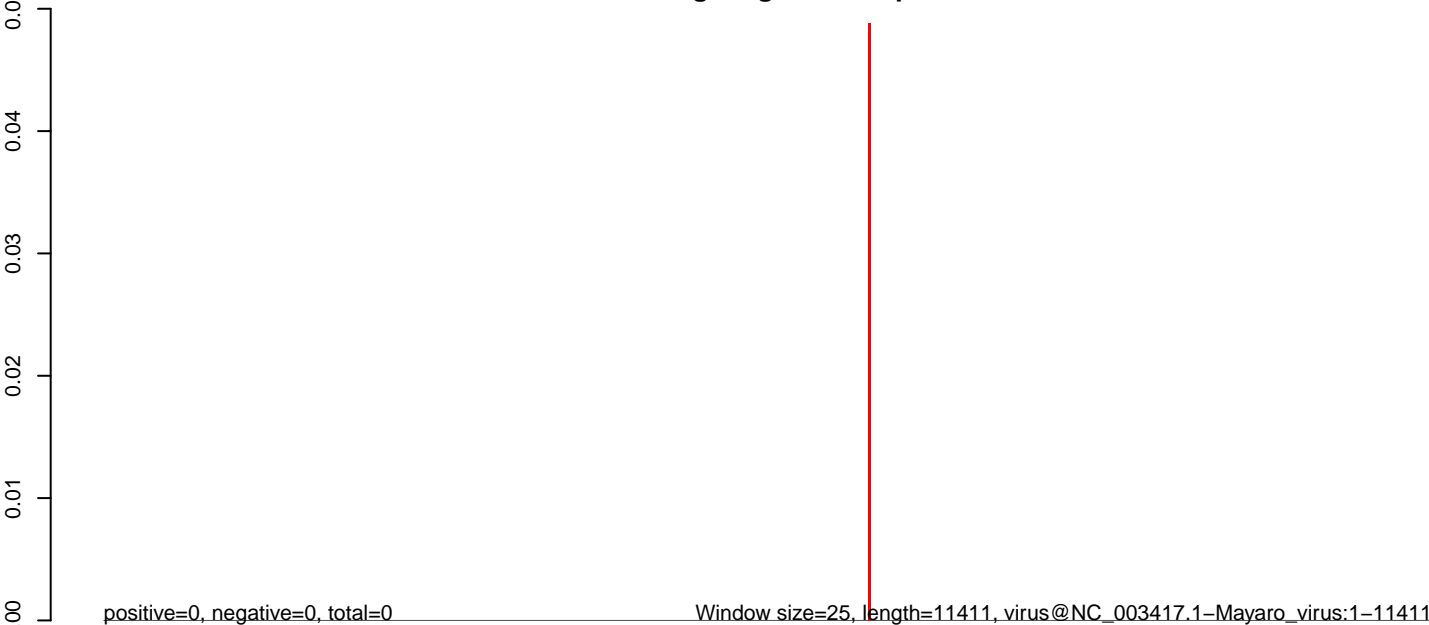
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

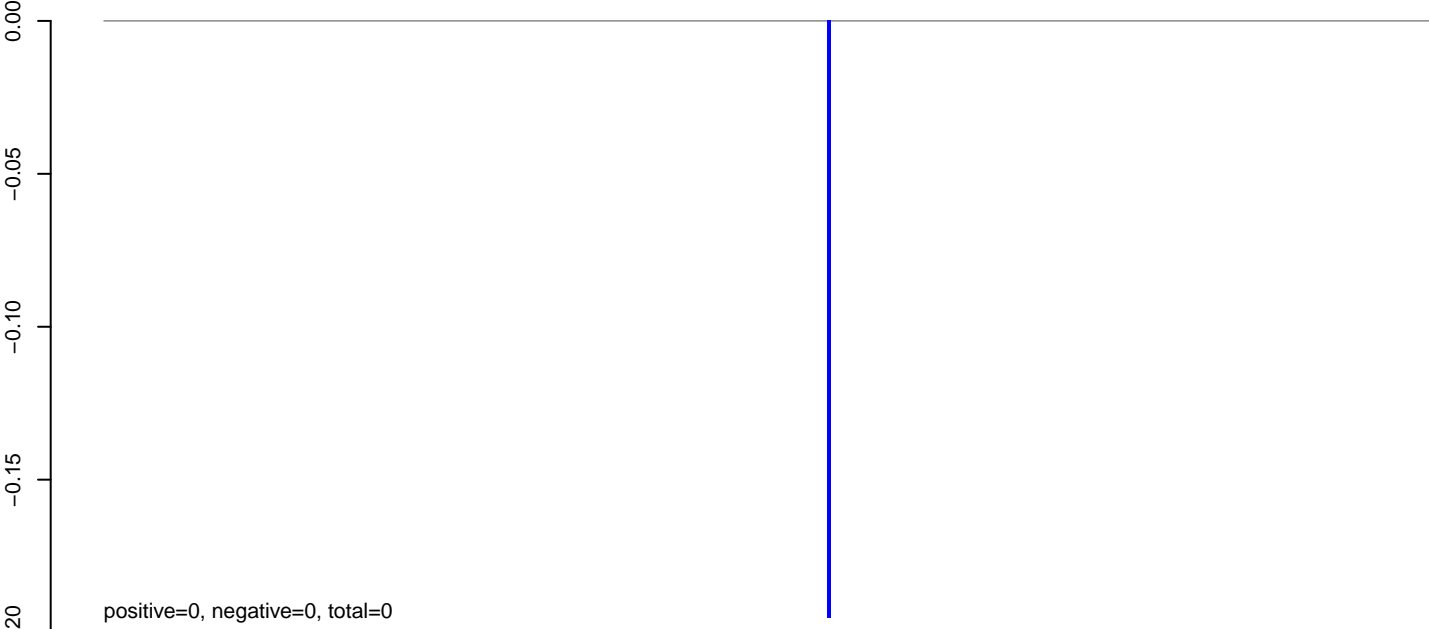


AeAeg_Aag2_cells.rep



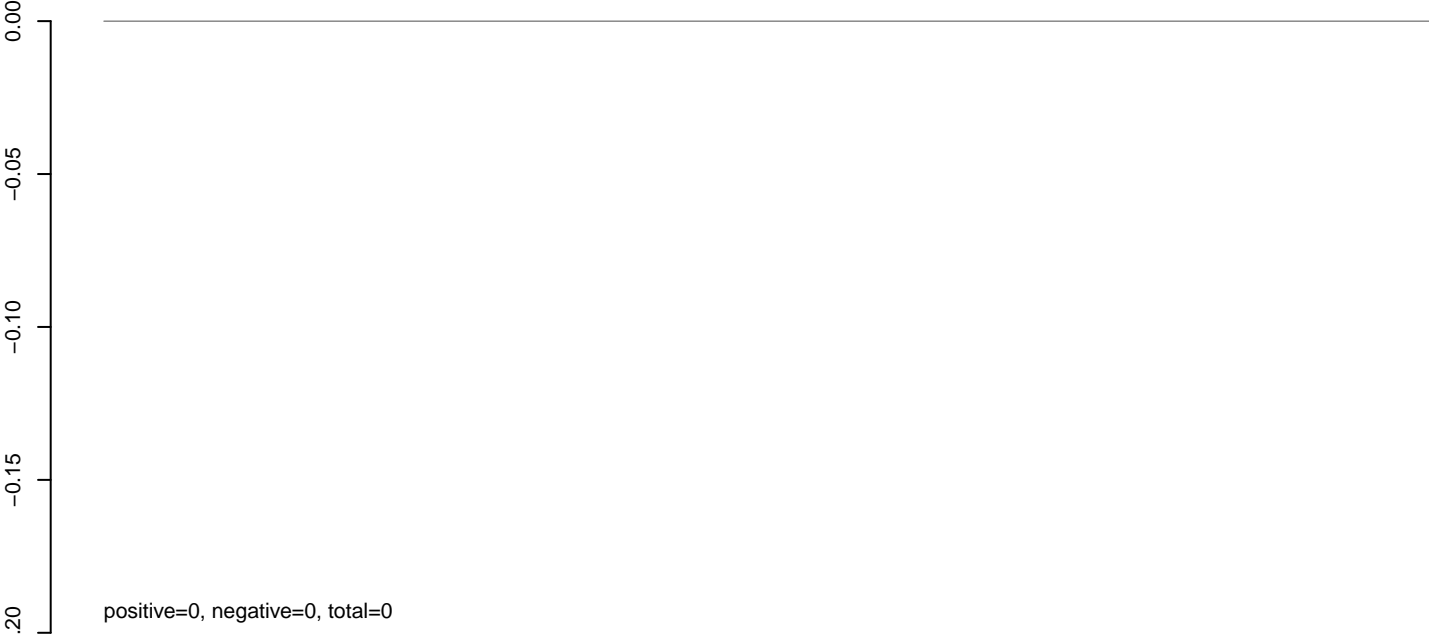
0 2000 4000 6000 8000 10000

AeAeg_Aag2_cells.18_23.rep



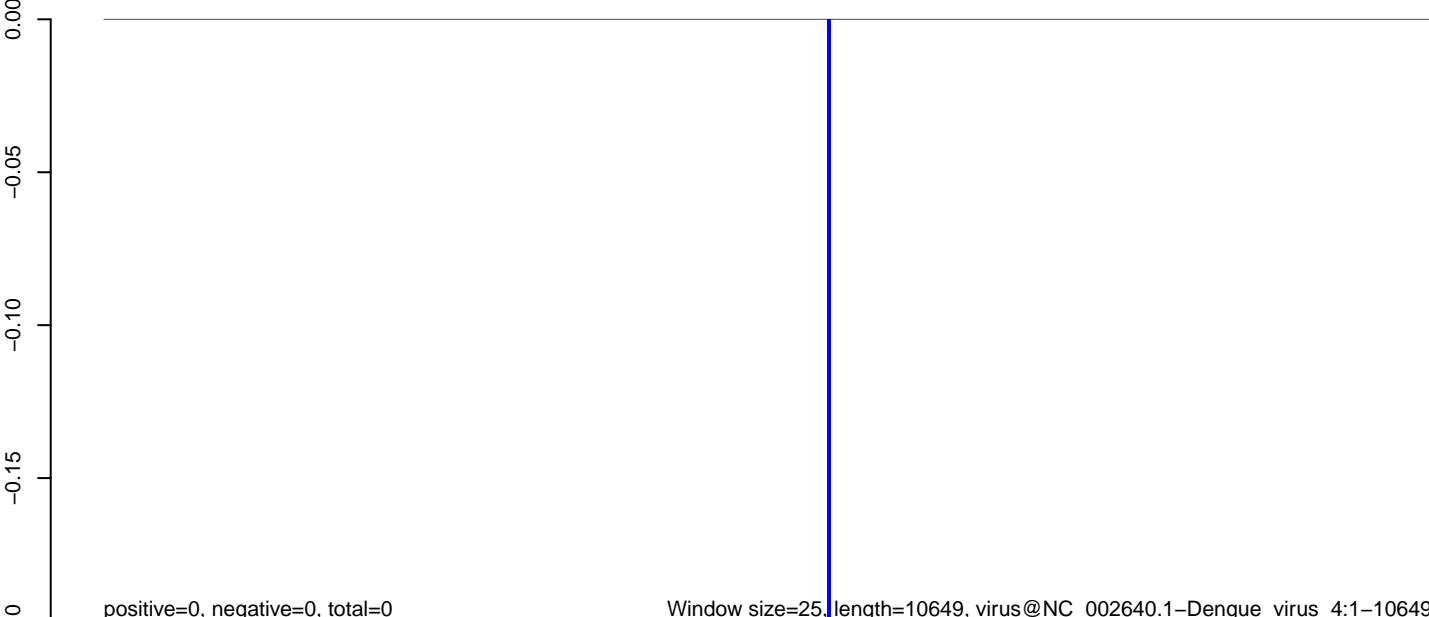
positive=0, negative=0, total=0

AeAeg_Aag2_cells.24_35.rep



positive=0, negative=0, total=0

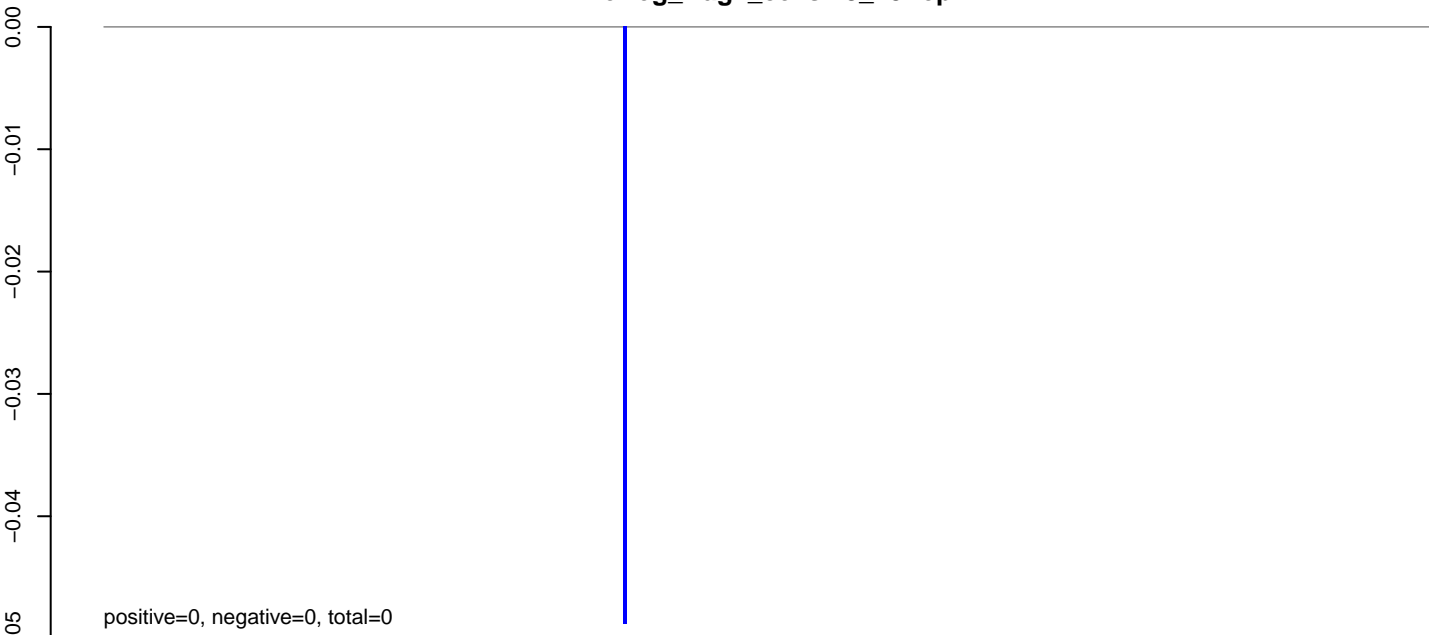
AeAeg_Aag2_cells.rep



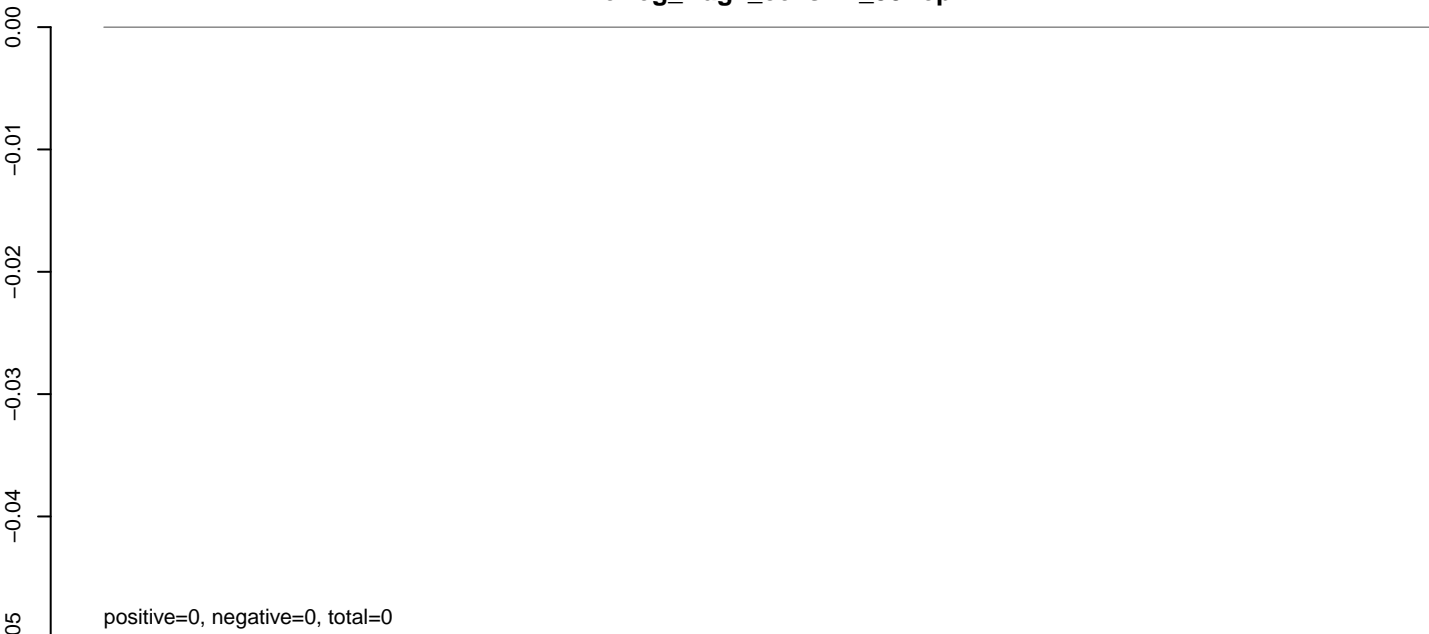
positive=0, negative=0, total=0

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

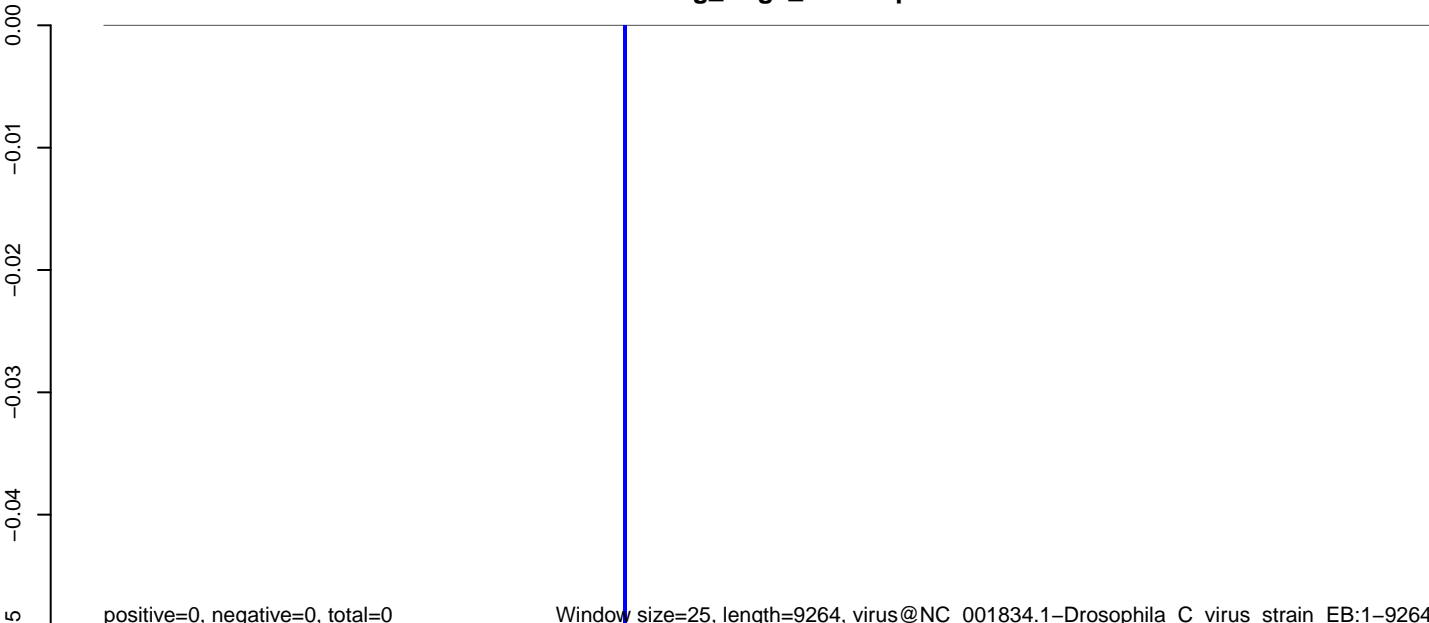
AeAeg_Aag2_cells.18_23.rep



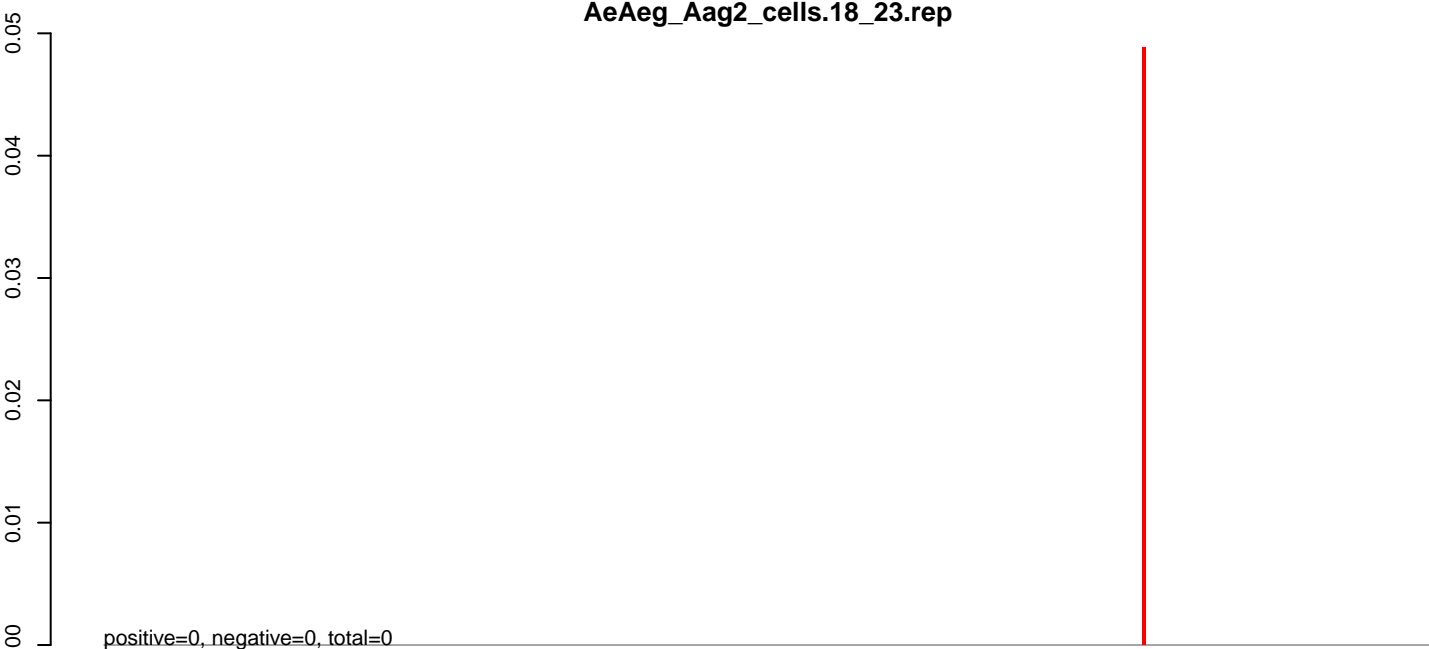
AeAeg_Aag2_cells.24_35.rep



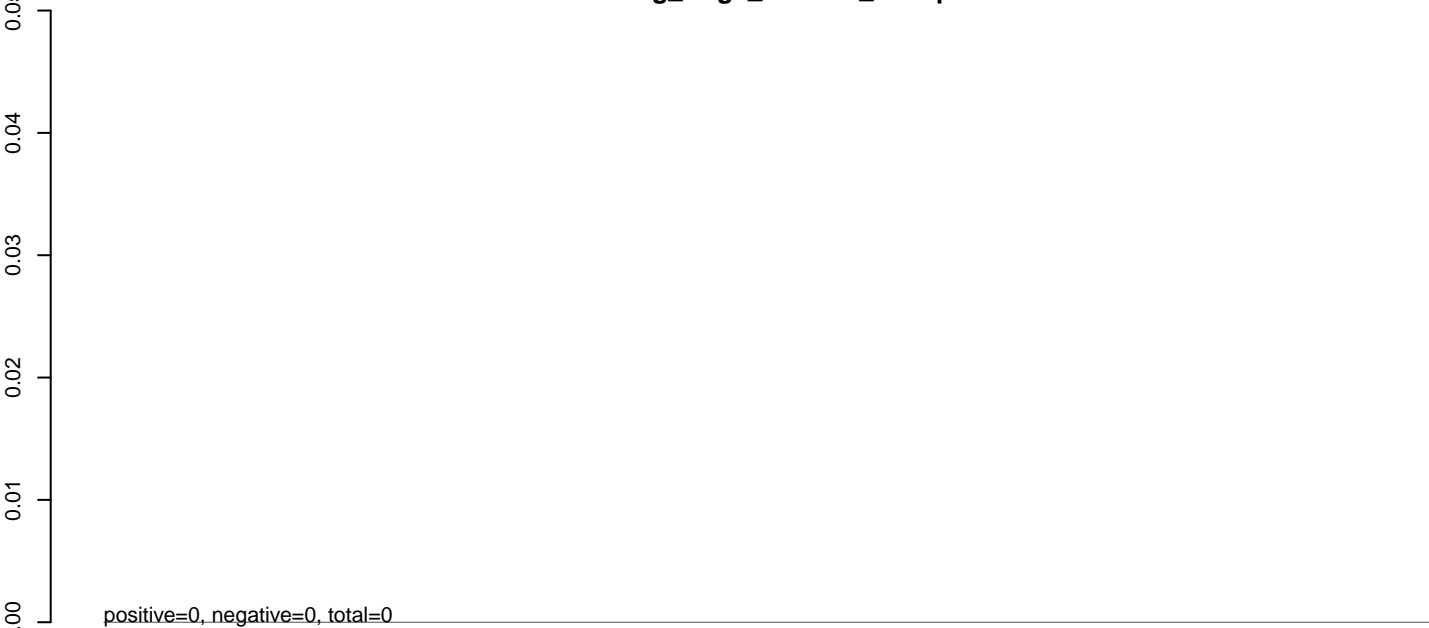
AeAeg_Aag2_cells.rep



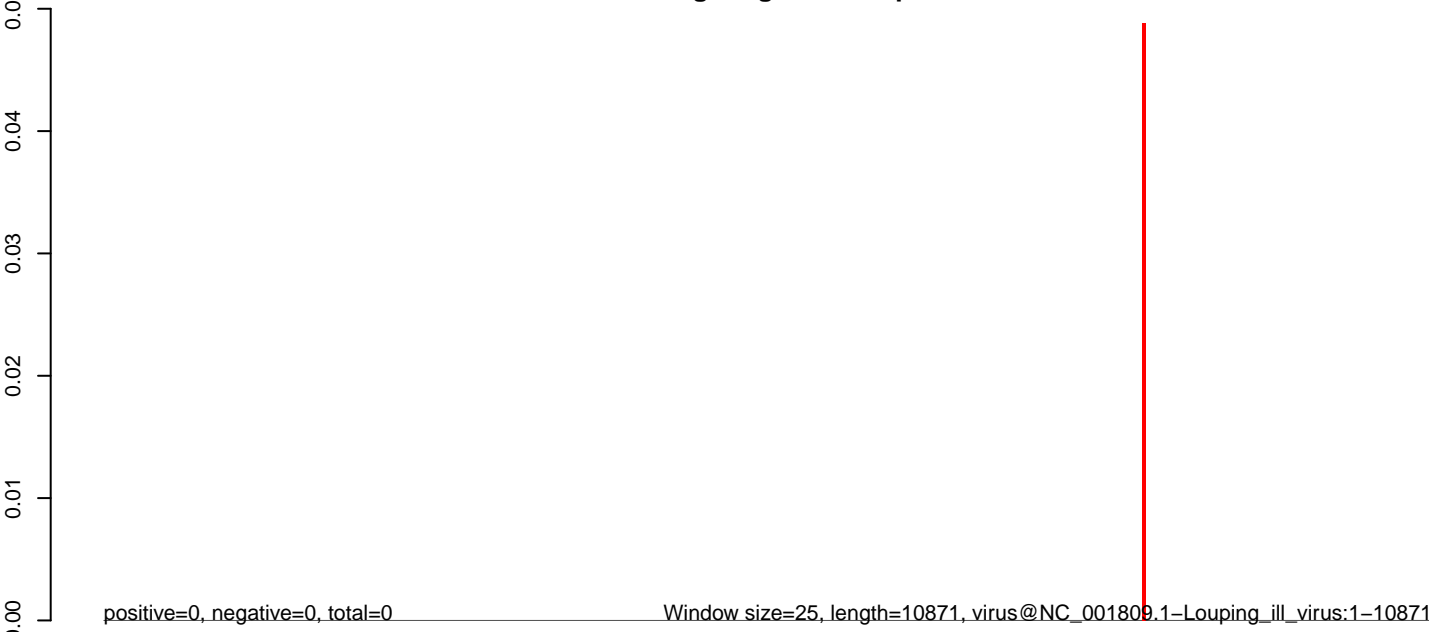
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

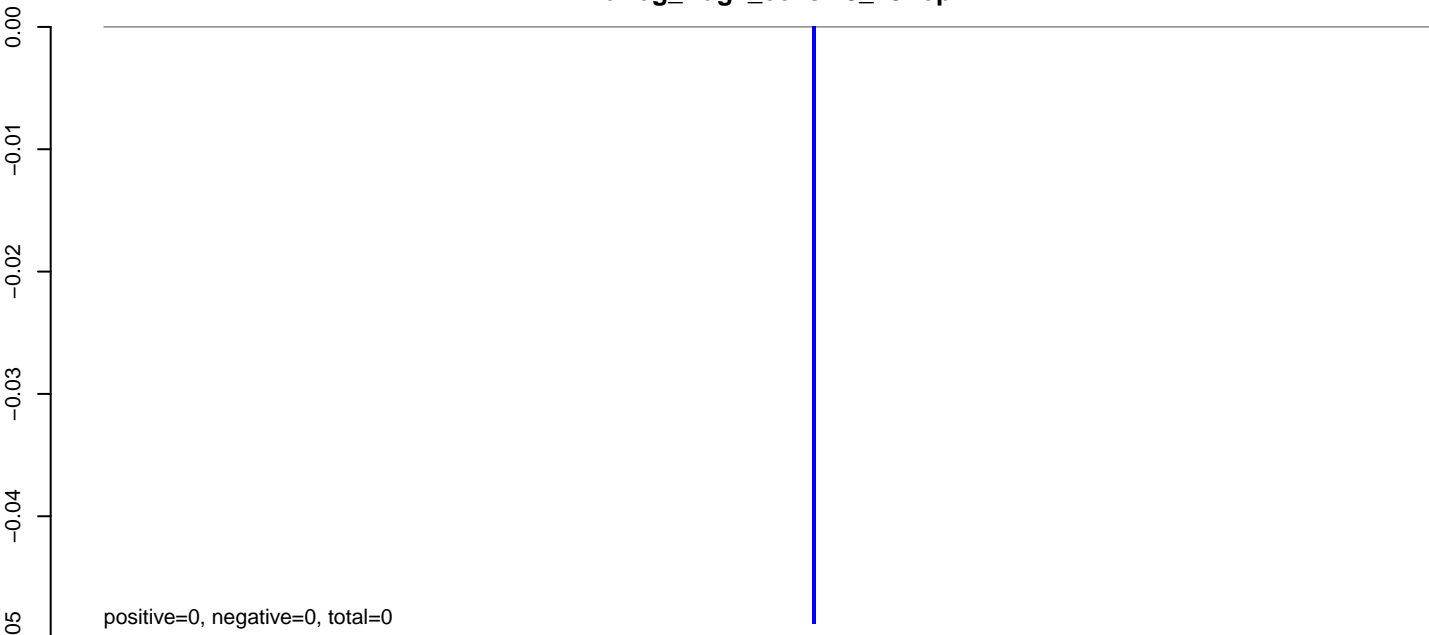


AeAeg_Aag2_cells.rep

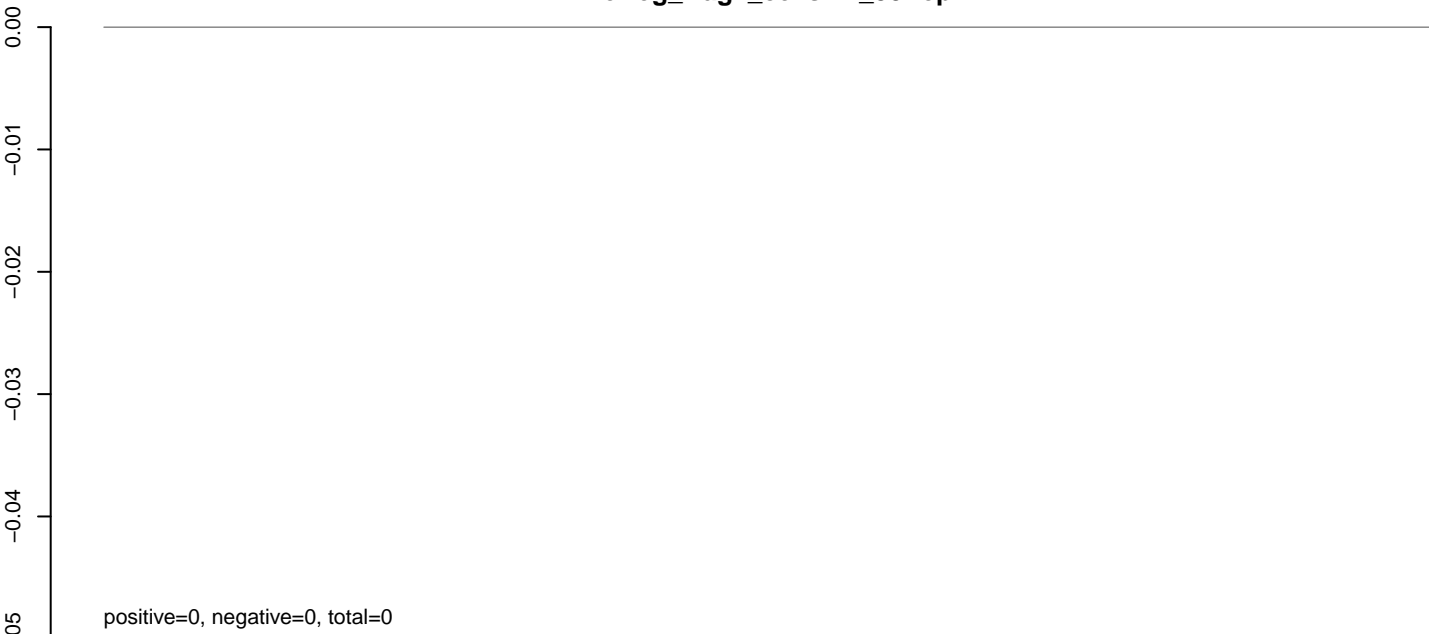


0 2000 4000 6000 8000 10000

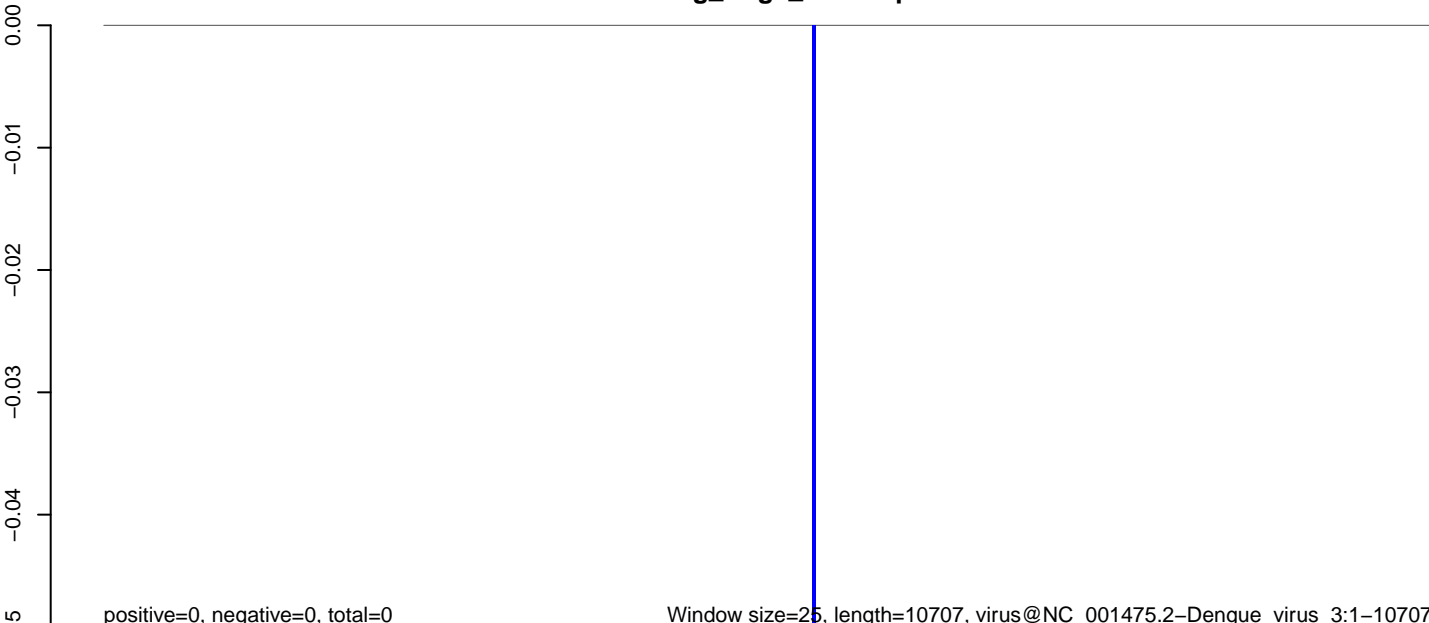
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



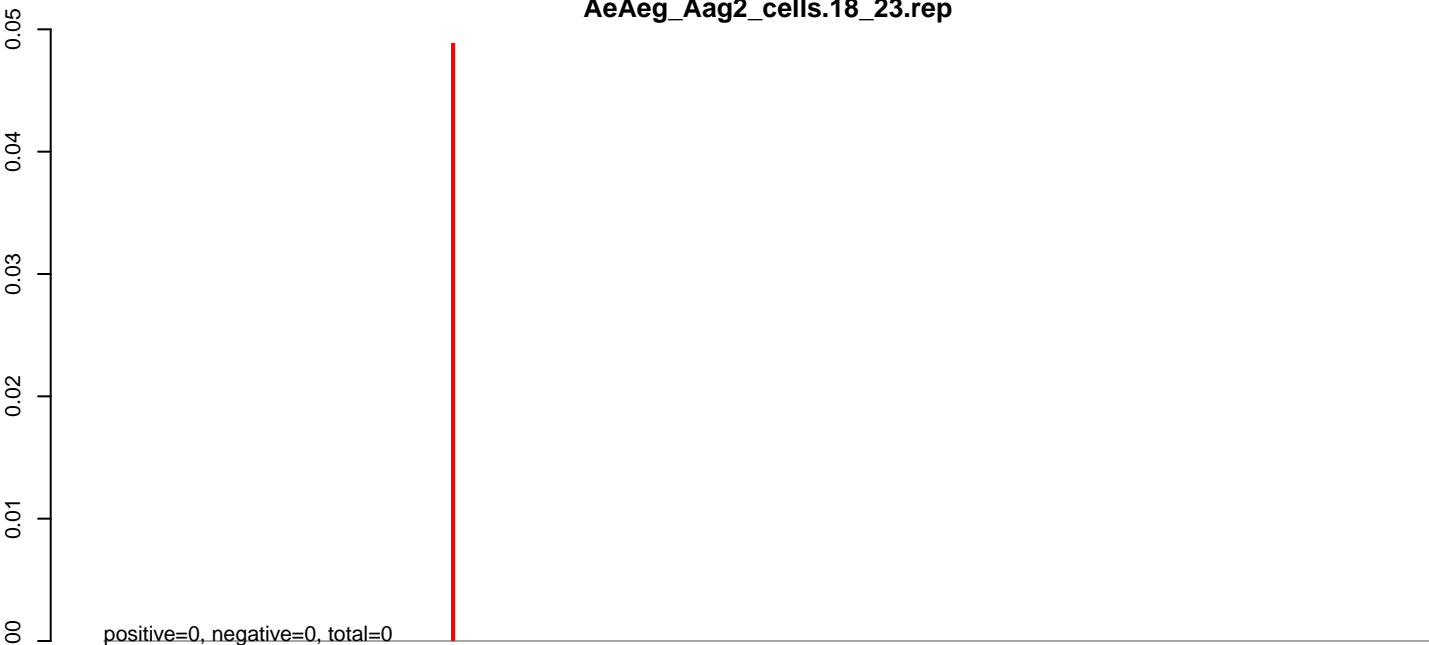
AeAeg_Aag2_cells.rep



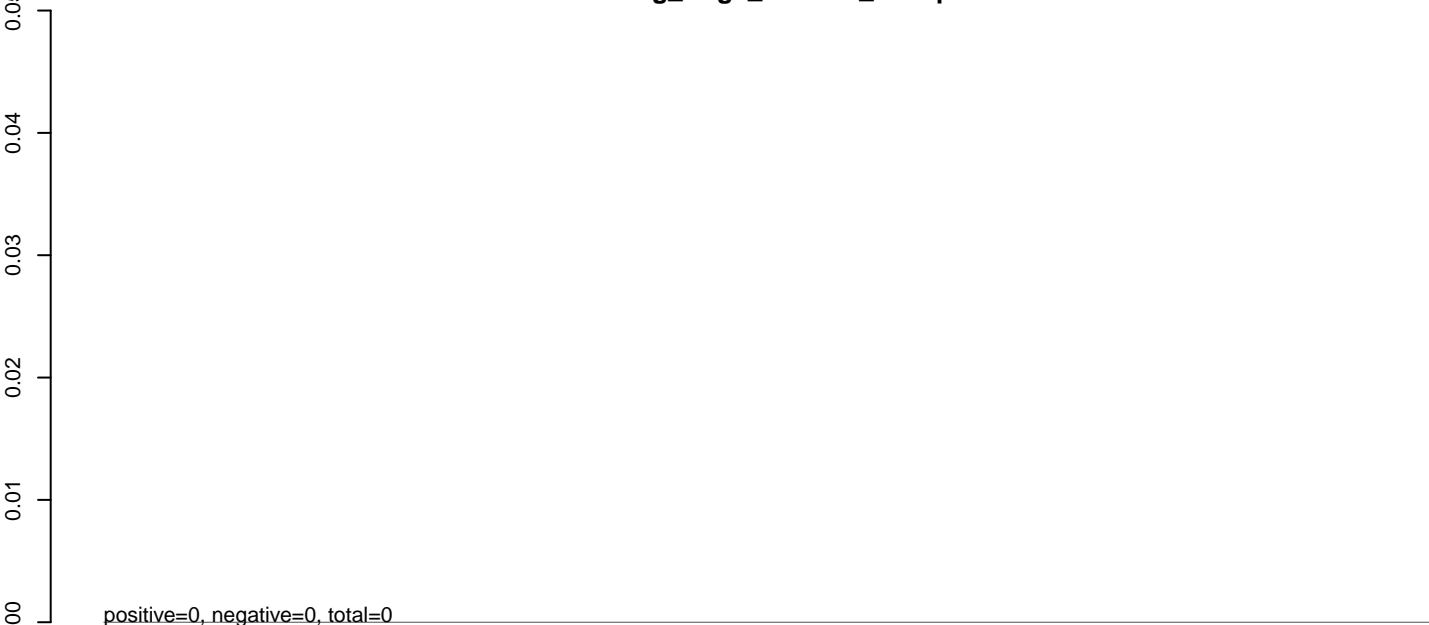
Window size=25, length=10707, virus@NC_001475.2-Dengue_virus_3:1-10707

0 2000 4000 6000 8000 10000

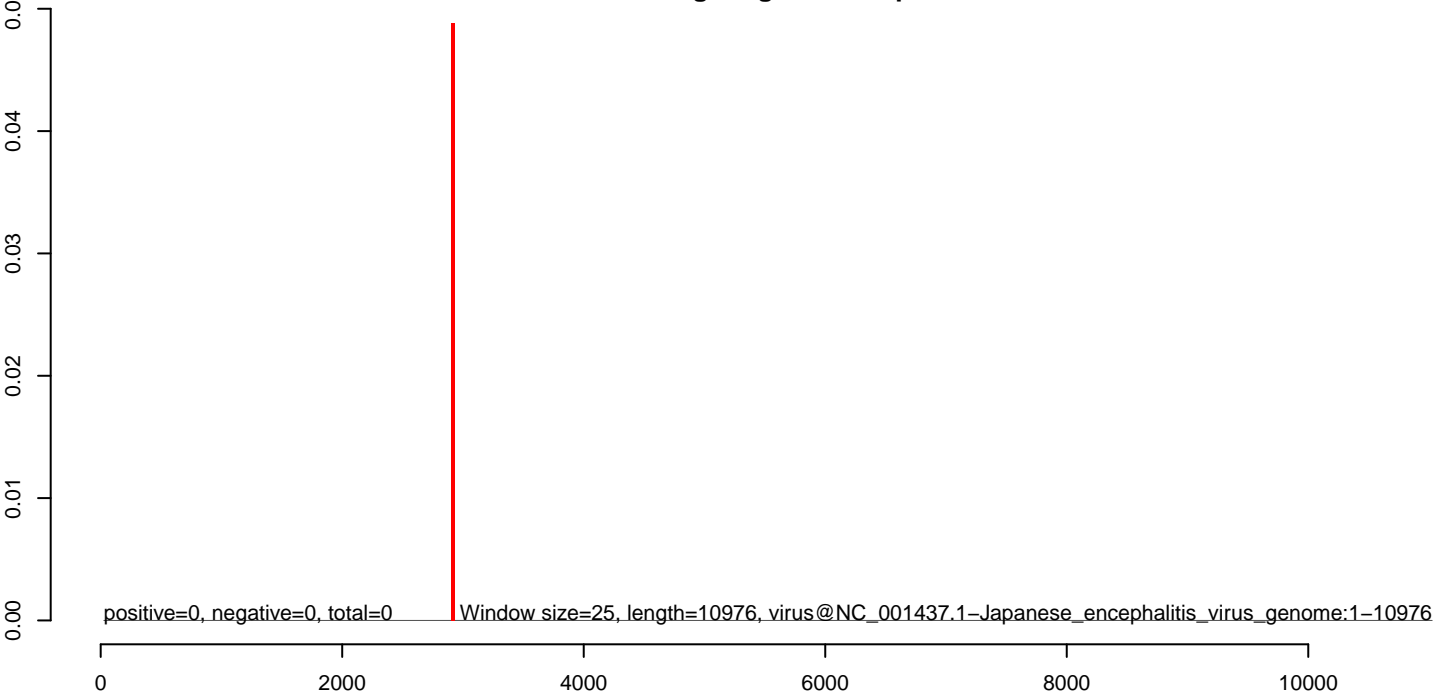
AeAeg_Aag2_cells.18_23.rep



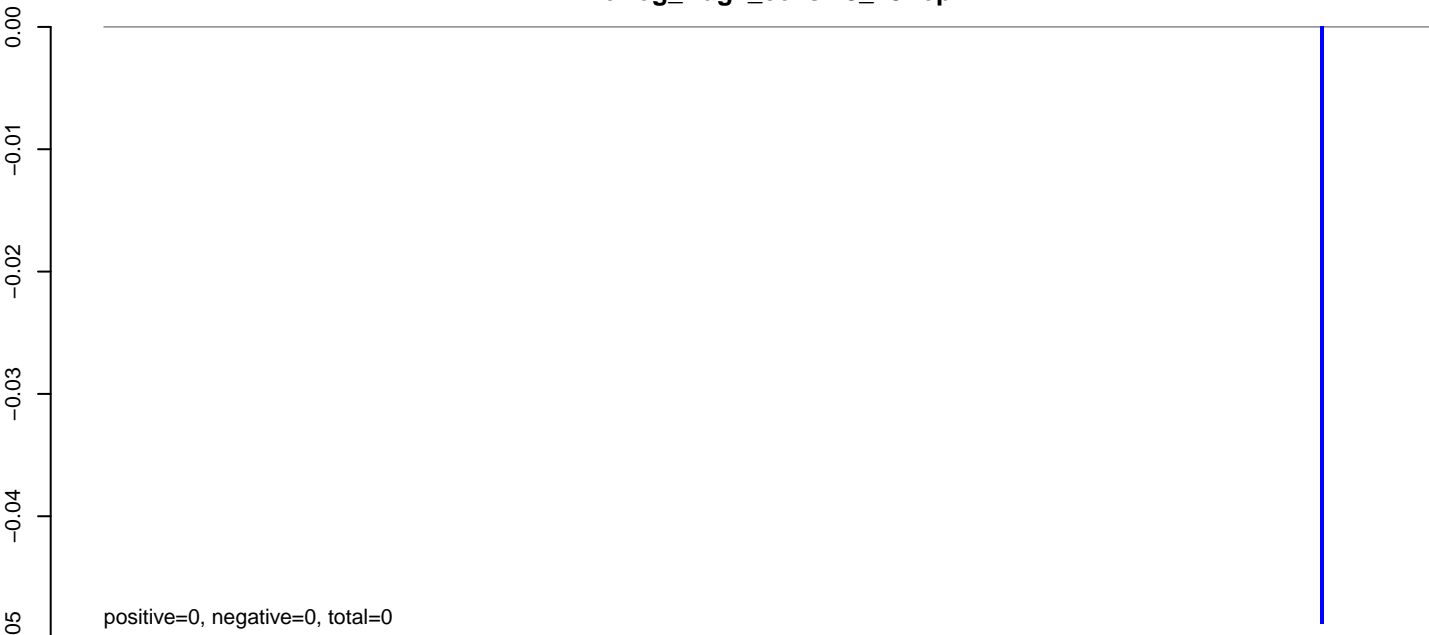
AeAeg_Aag2_cells.24_35.rep



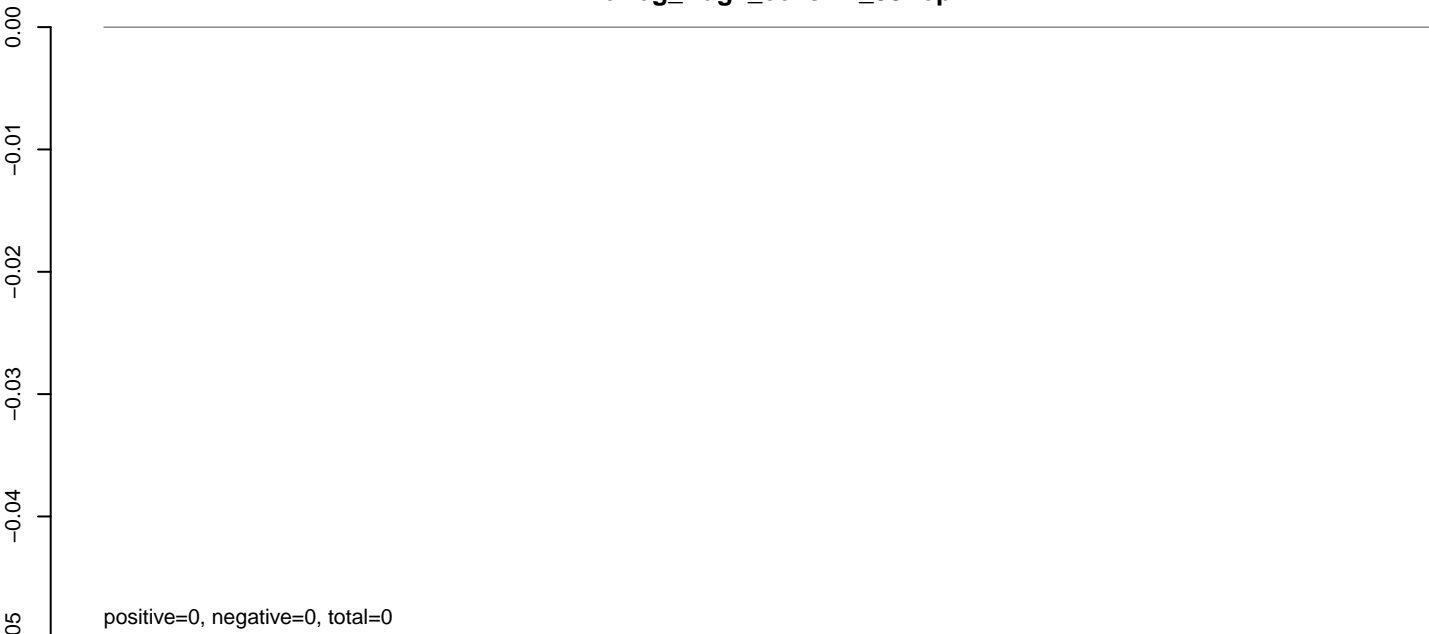
AeAeg_Aag2_cells.rep



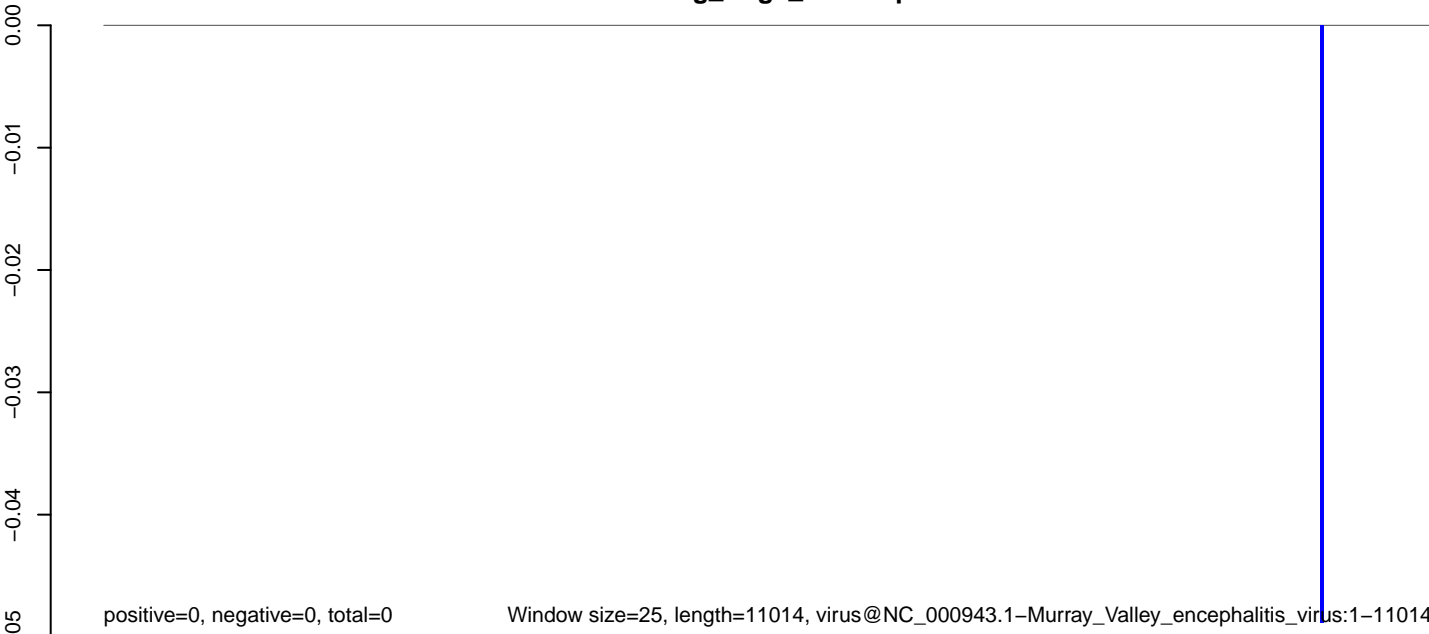
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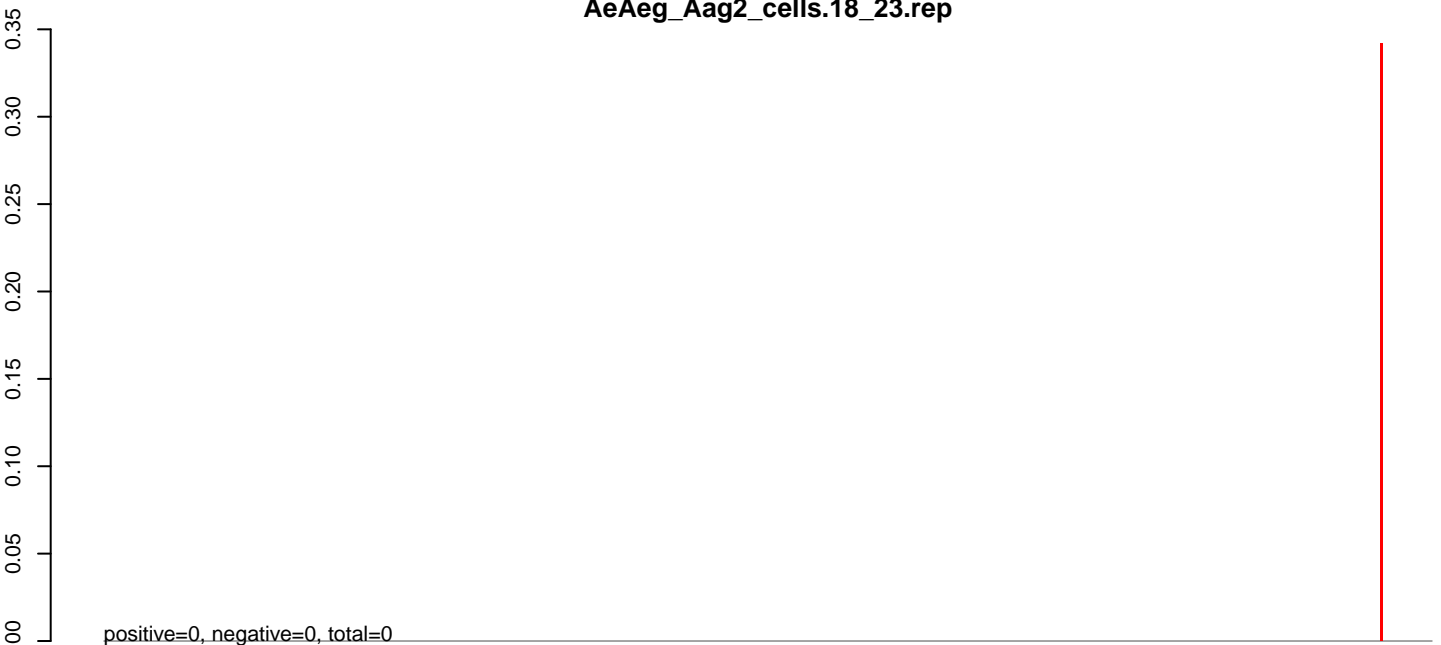
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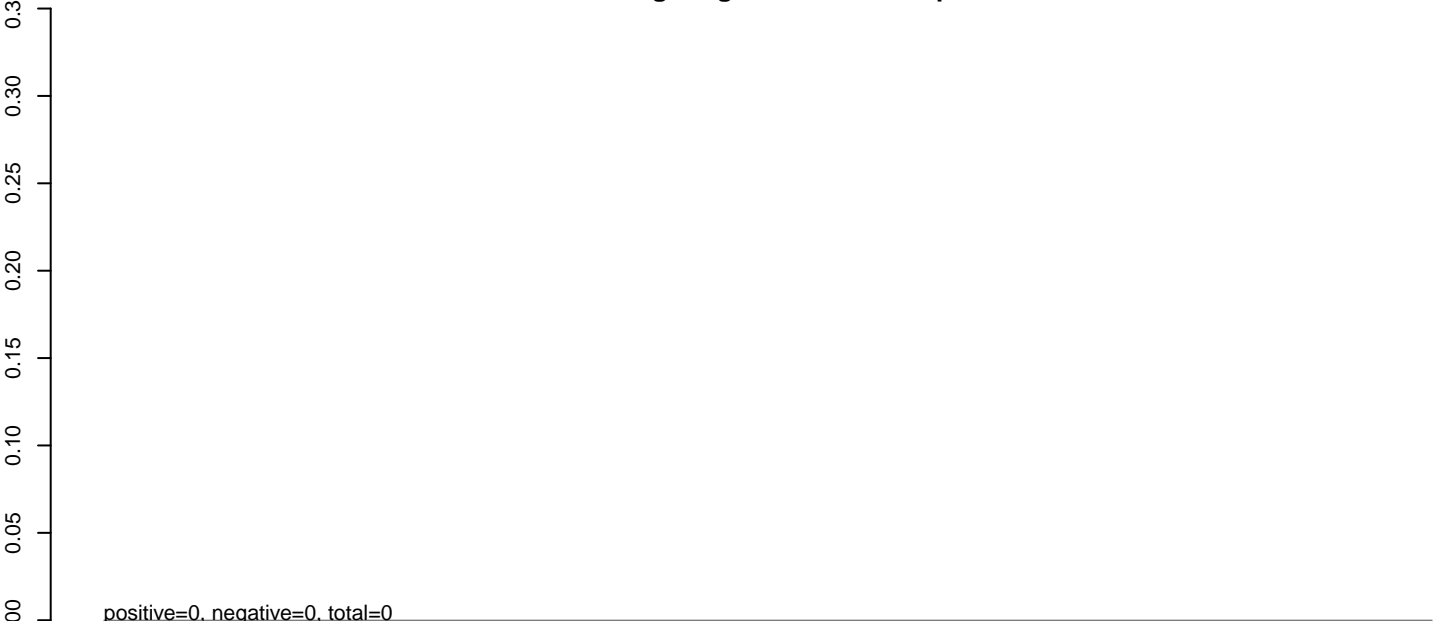
AeAeg_Aag2_cells.rep



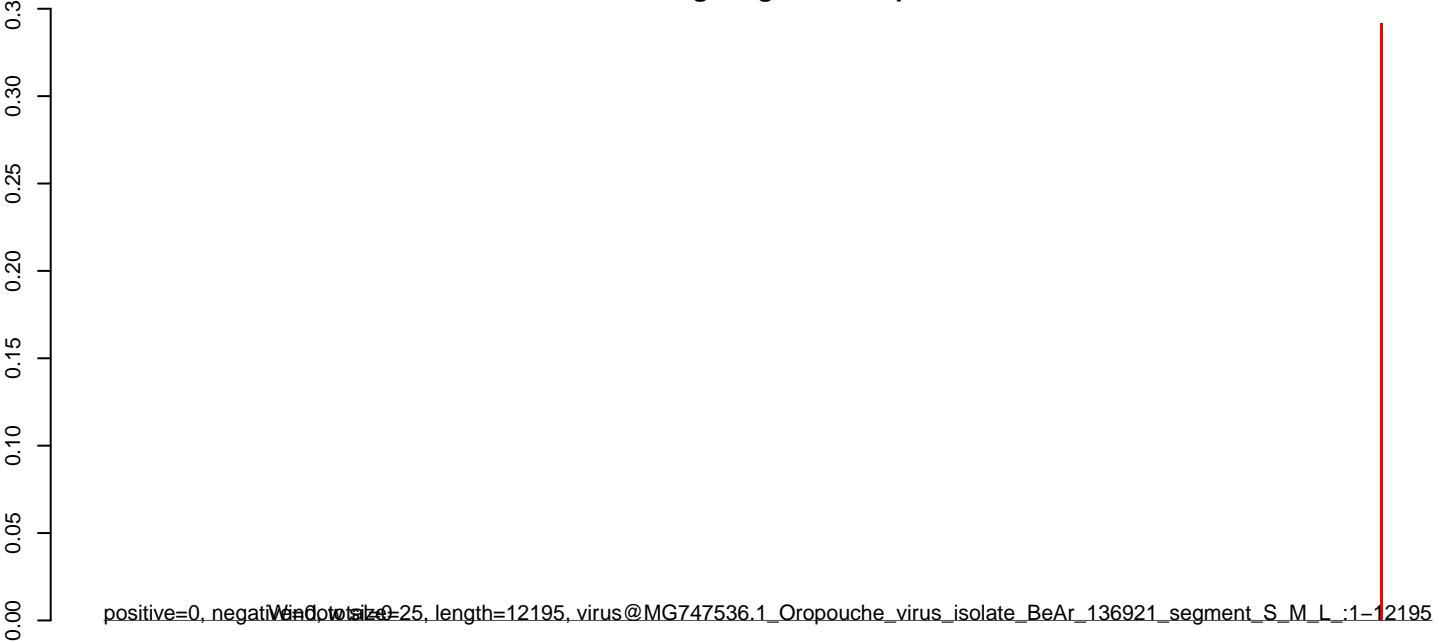
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



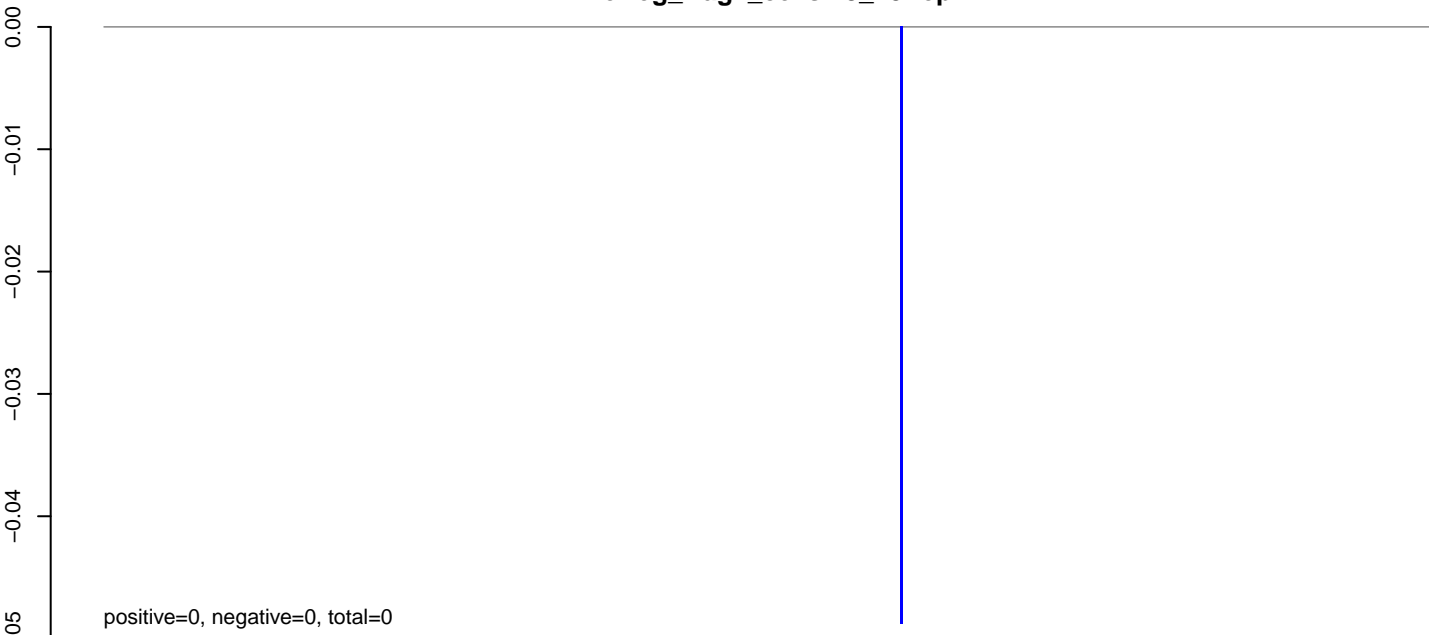
AeAeg_Aag2_cells.rep



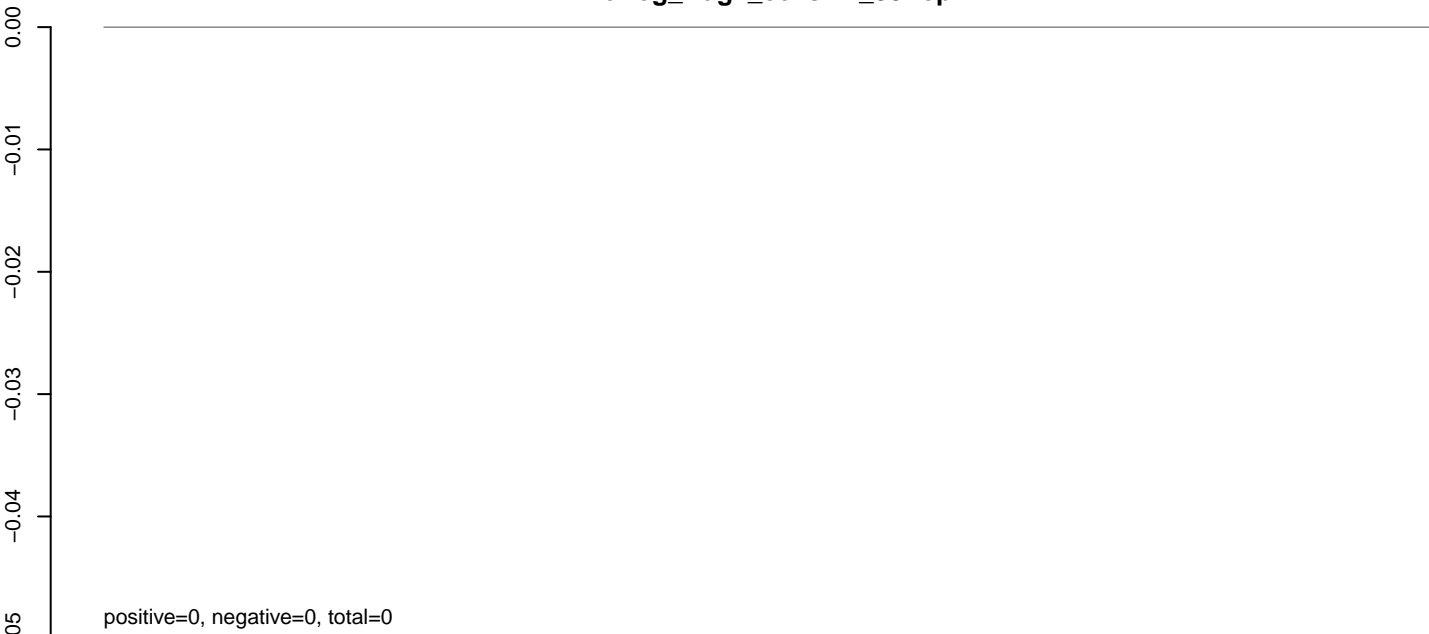
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0 2000 4000 6000 8000 10000 12000

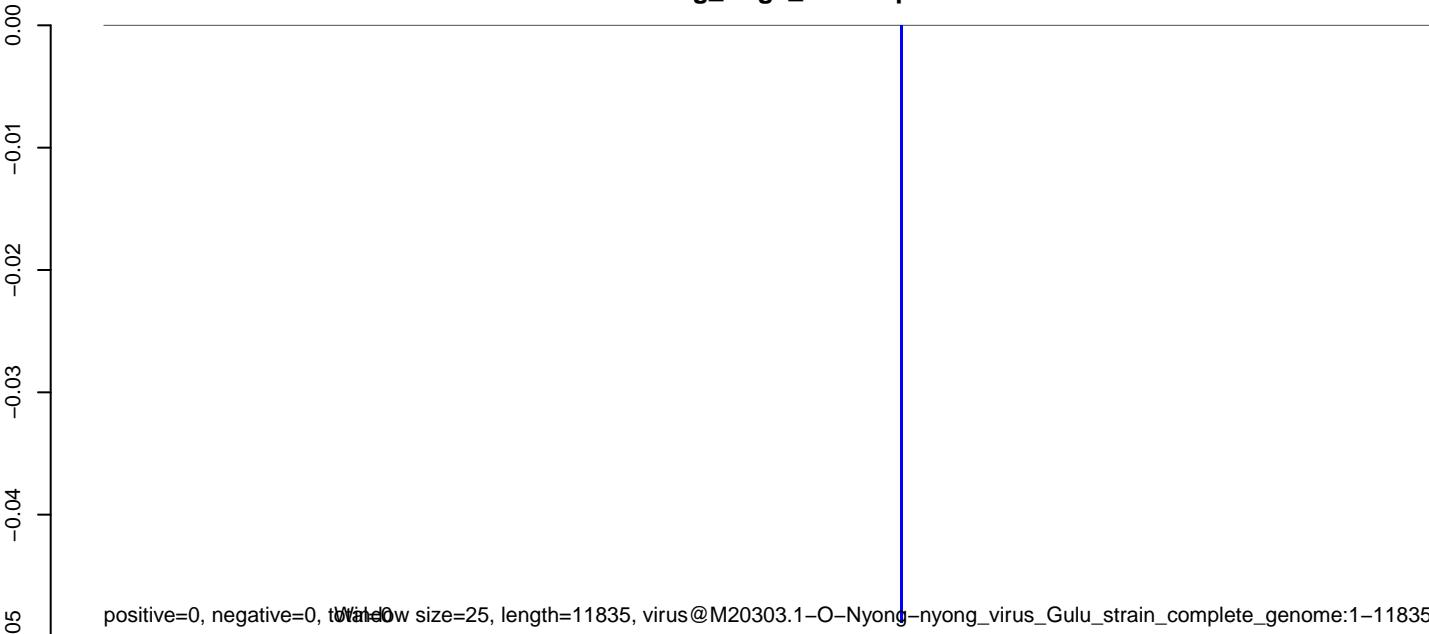
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



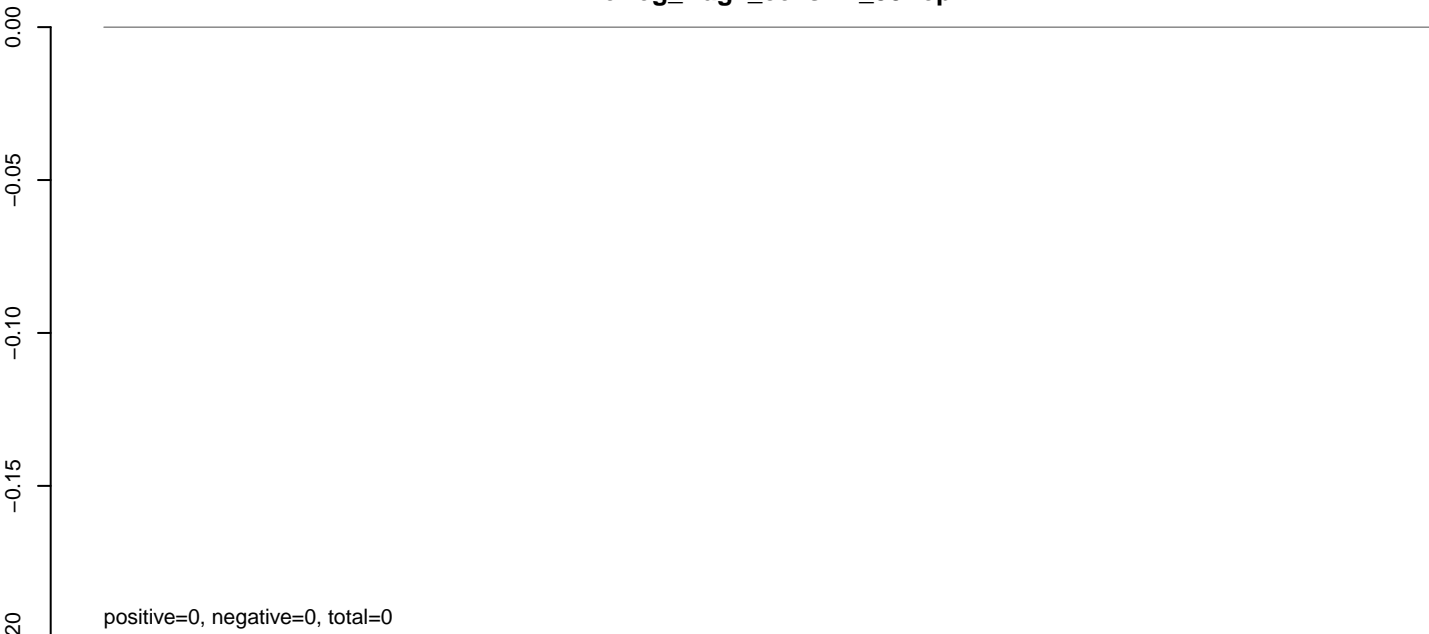
AeAeg_Aag2_cells.rep



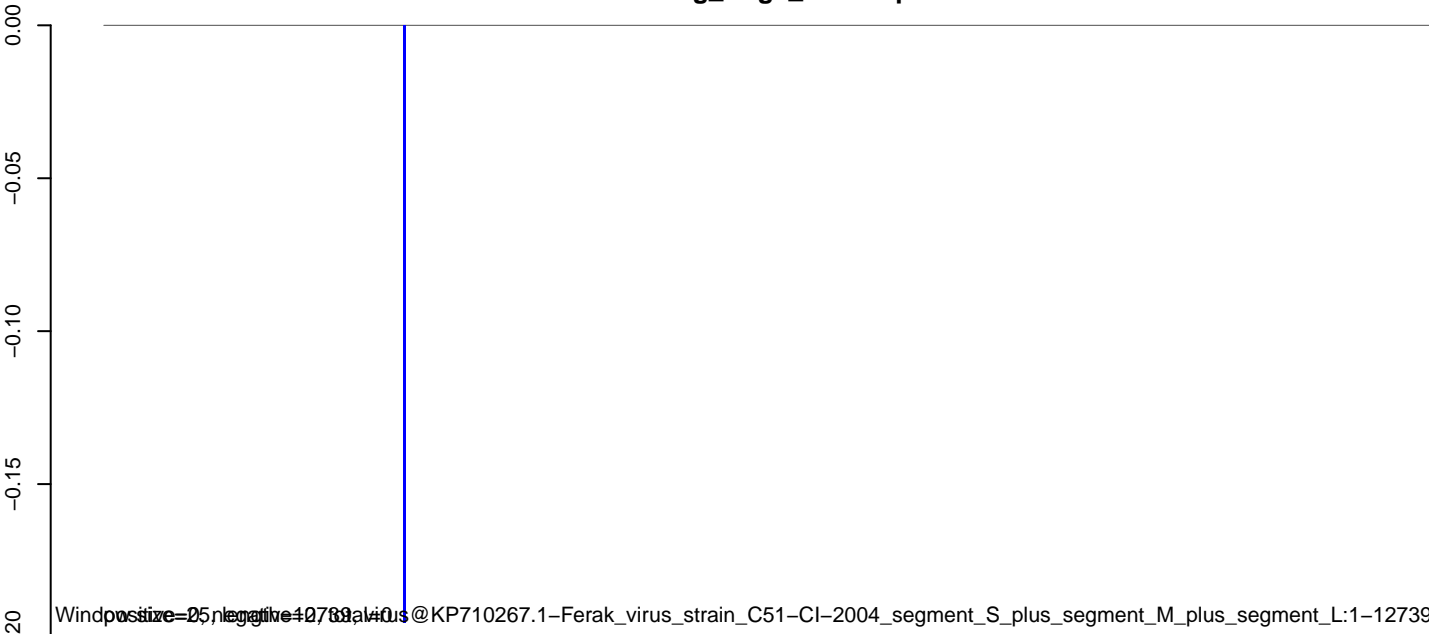
AeAeg_Aag2_cells.18_23.rep



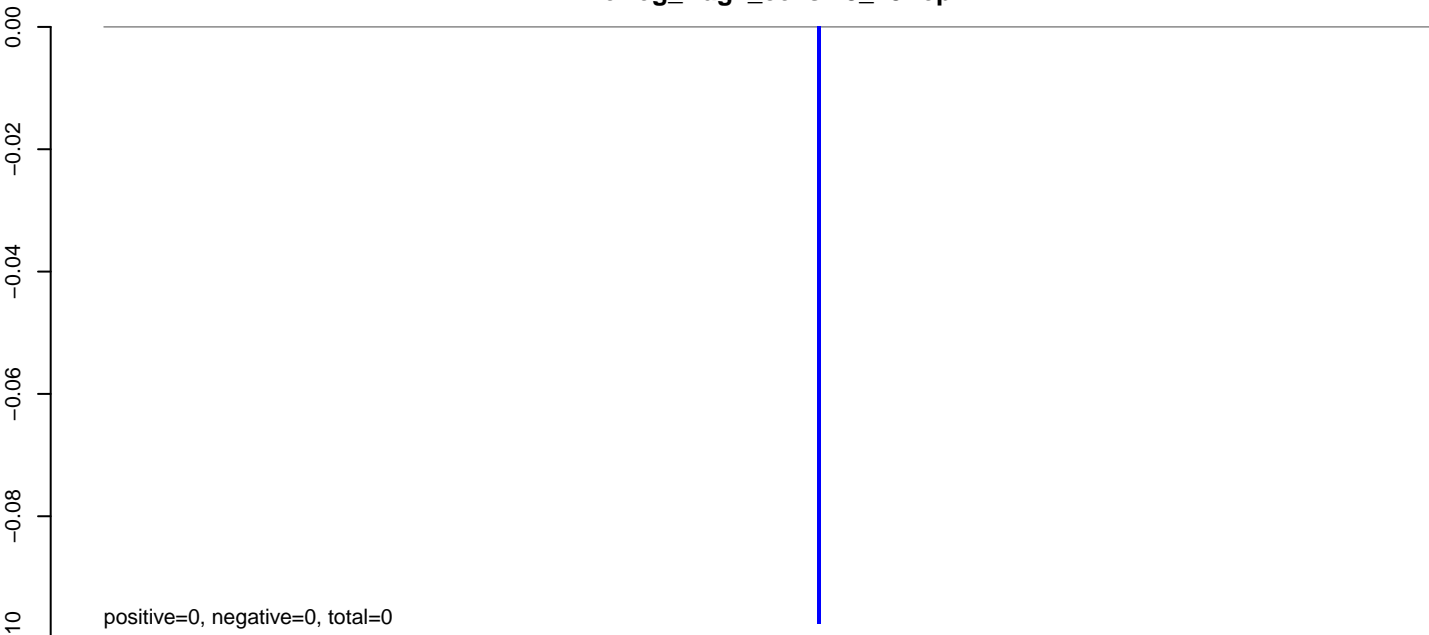
AeAeg_Aag2_cells.24_35.rep



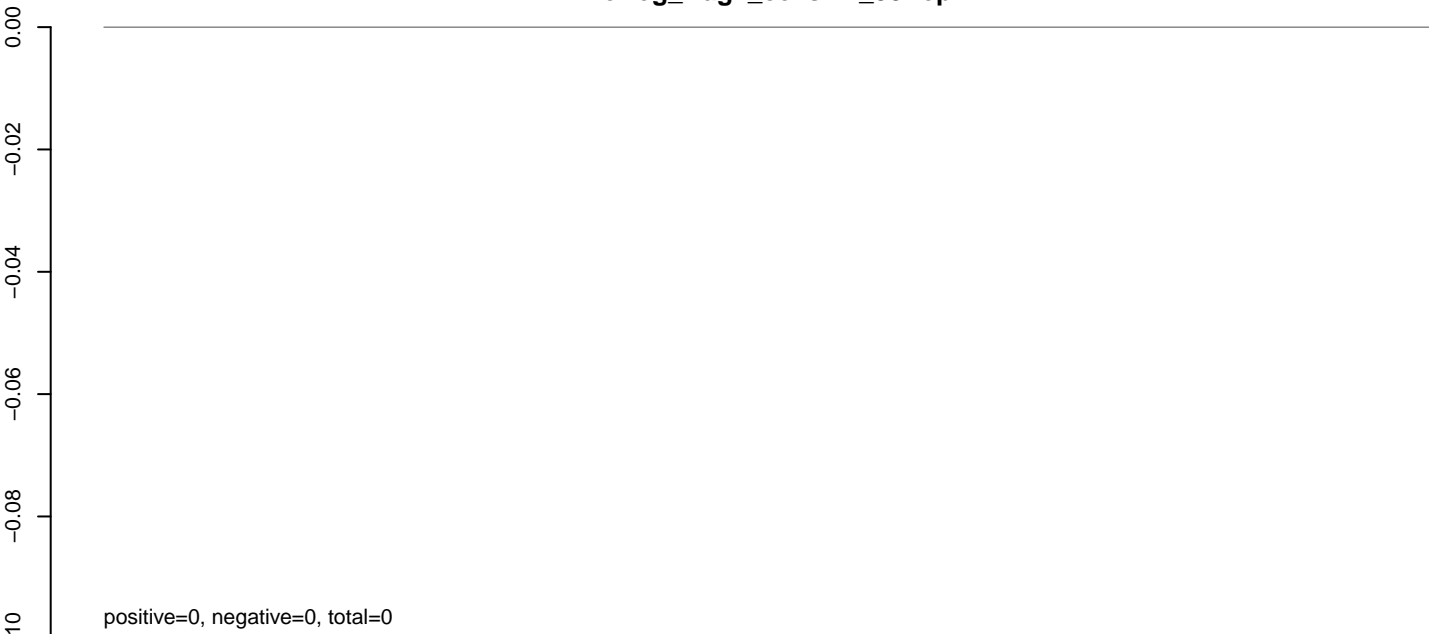
AeAeg_Aag2_cells.rep



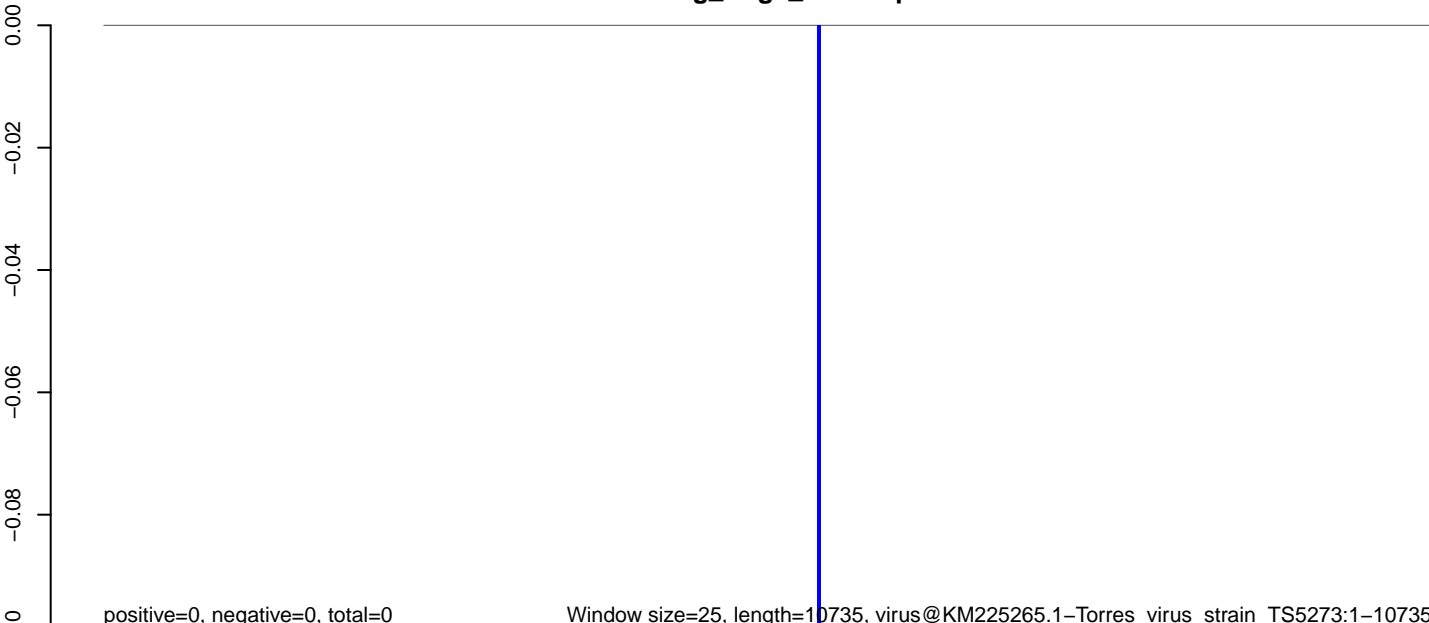
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

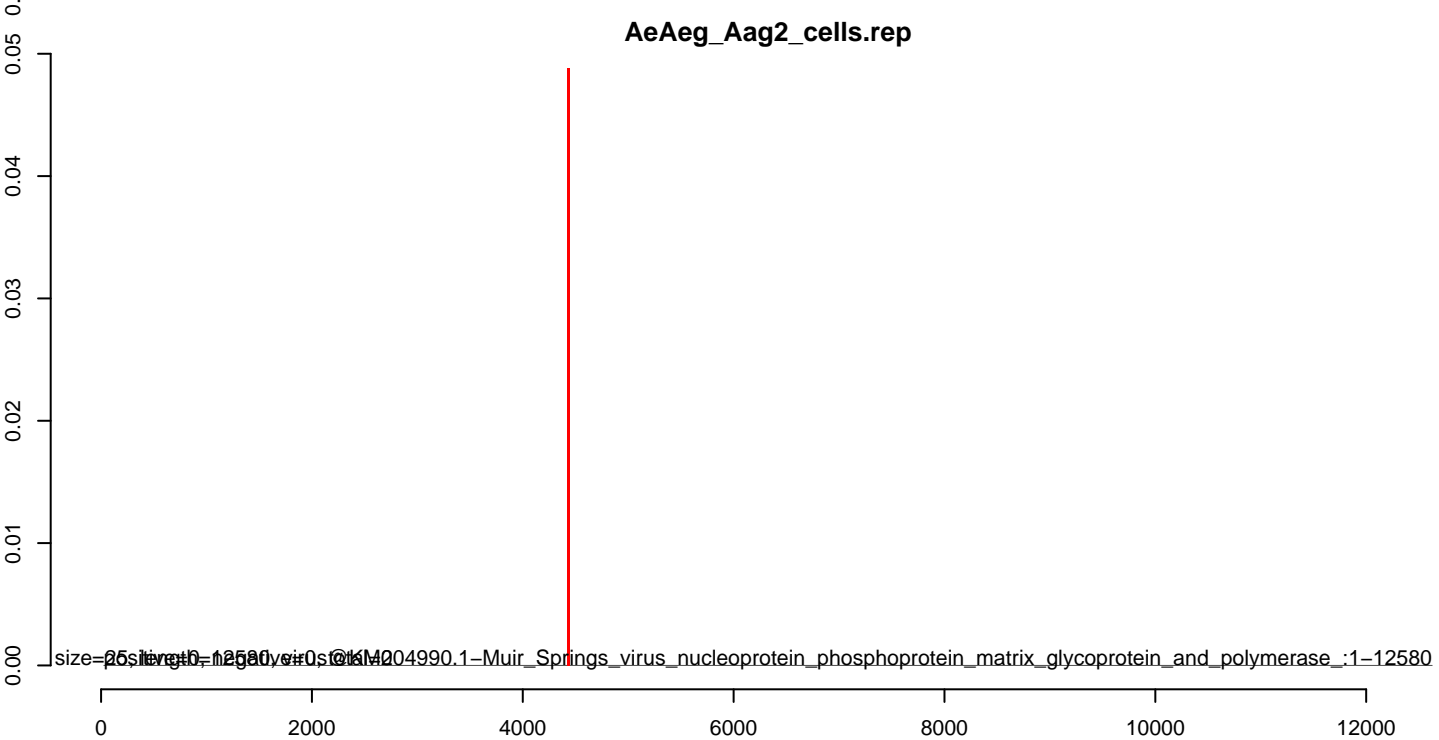
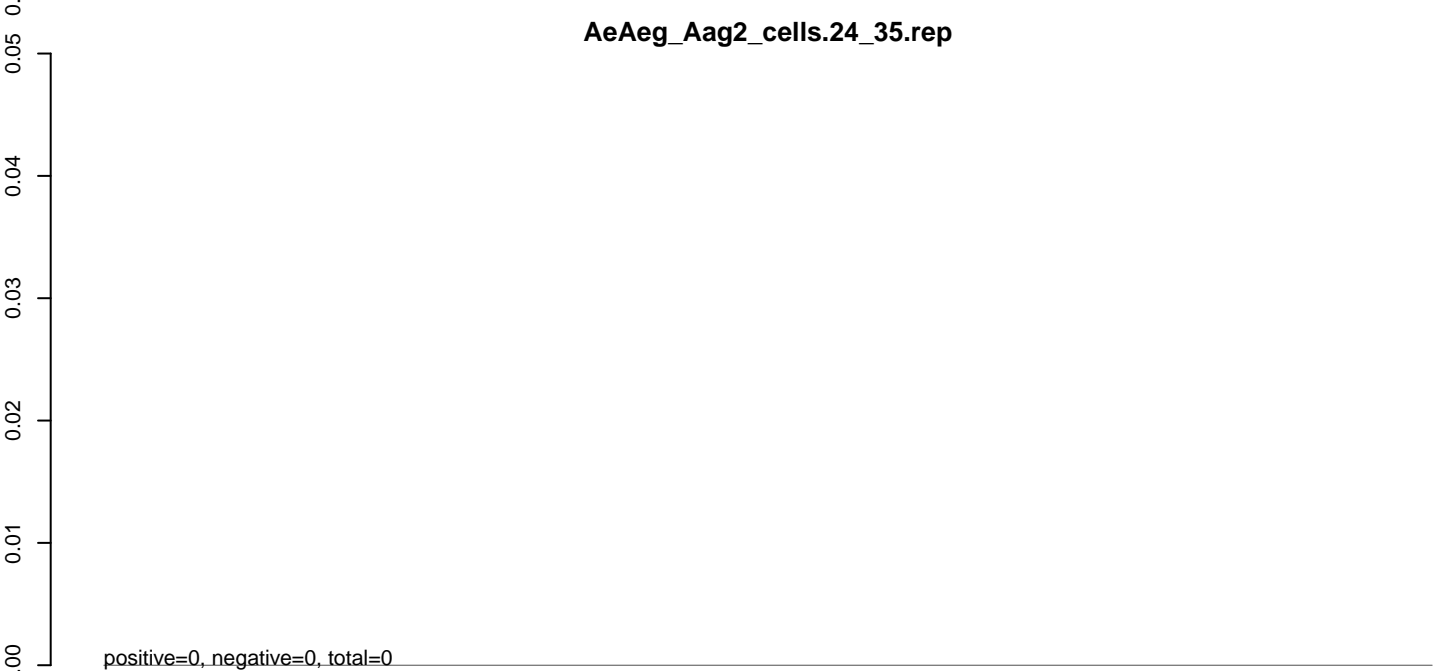
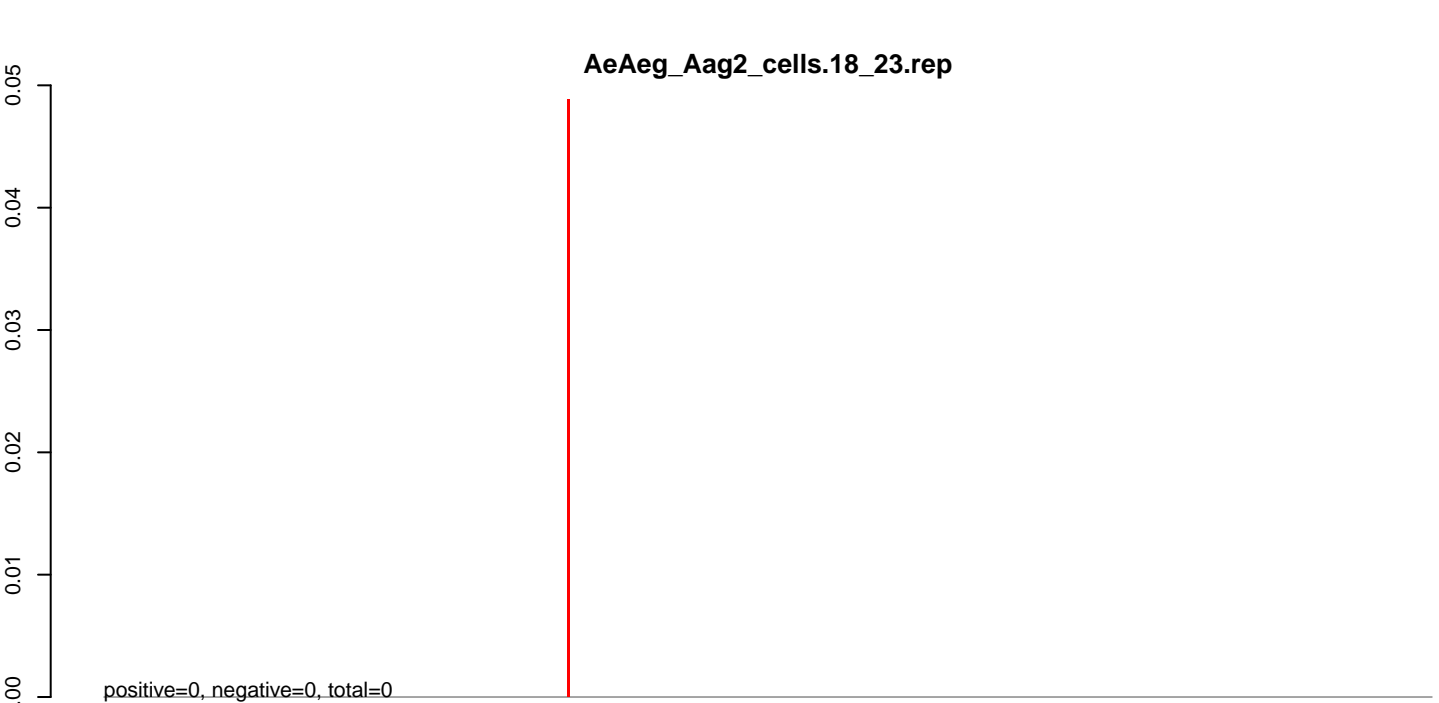


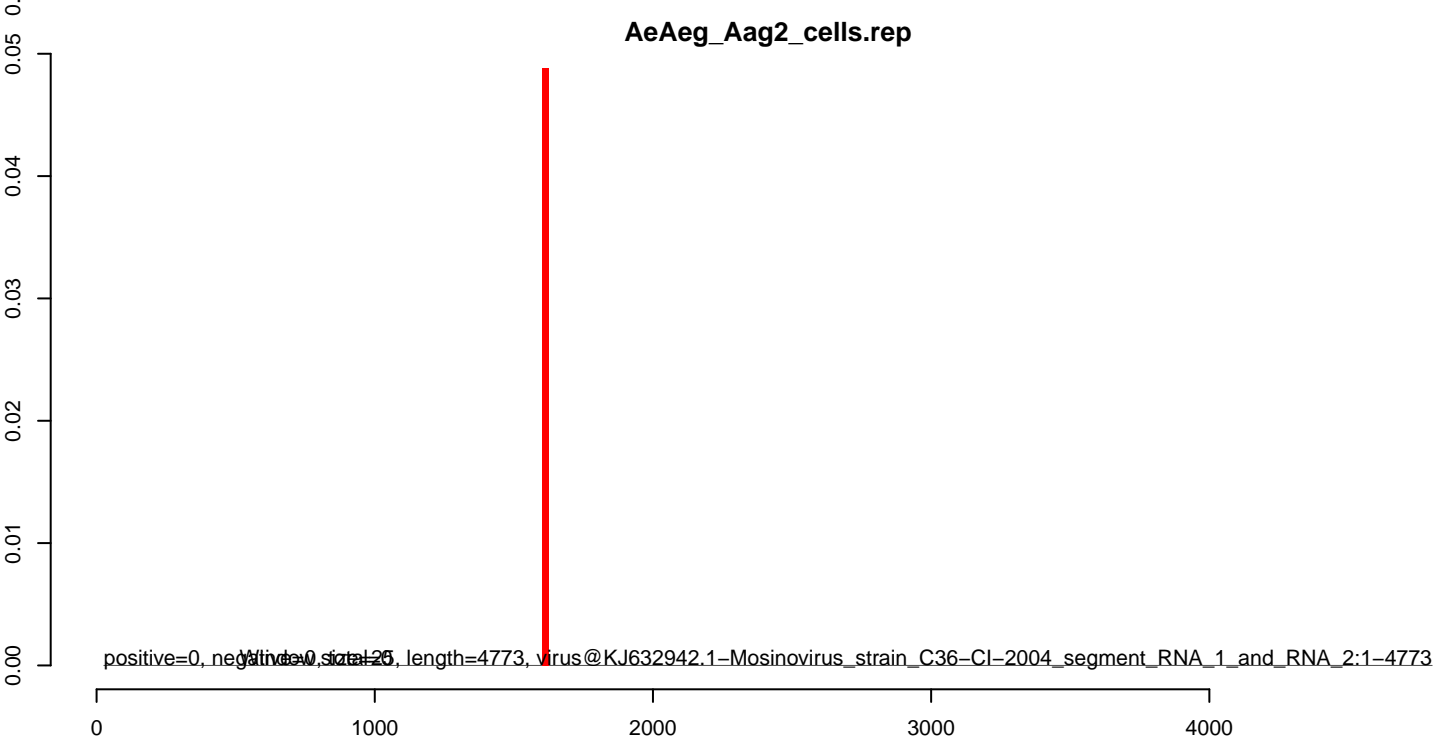
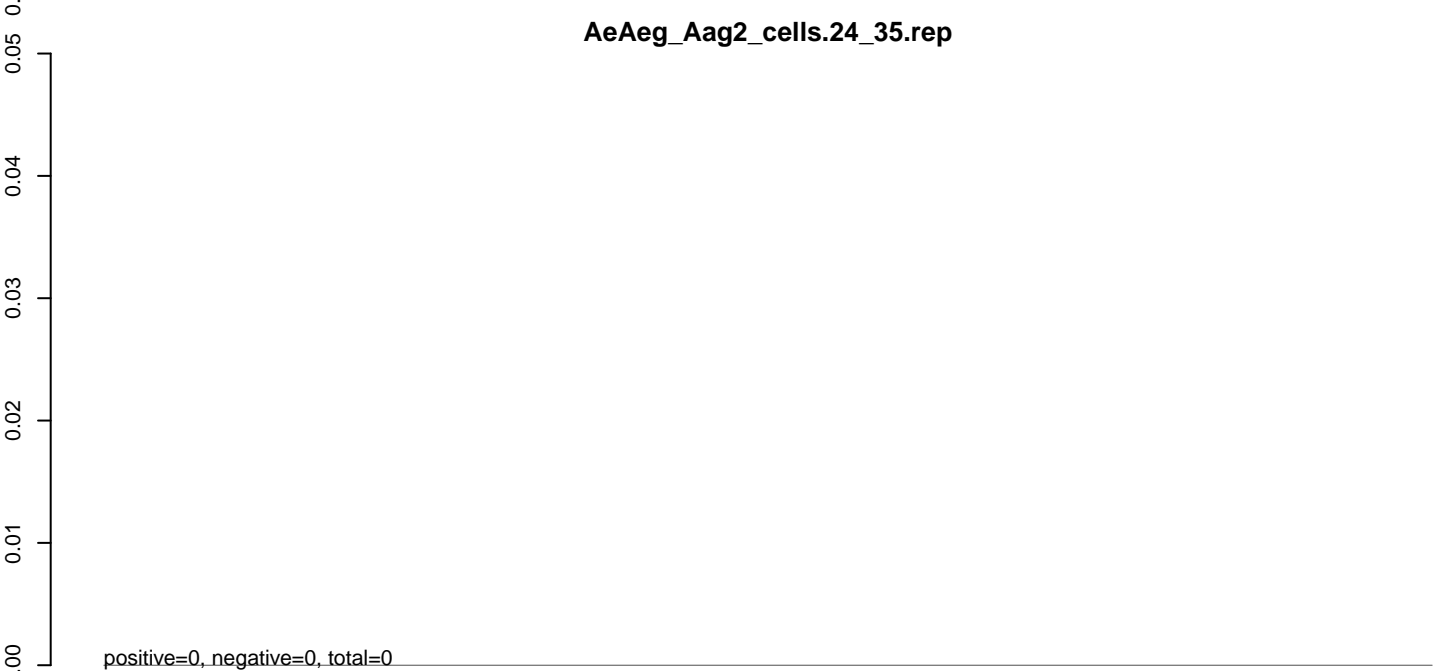
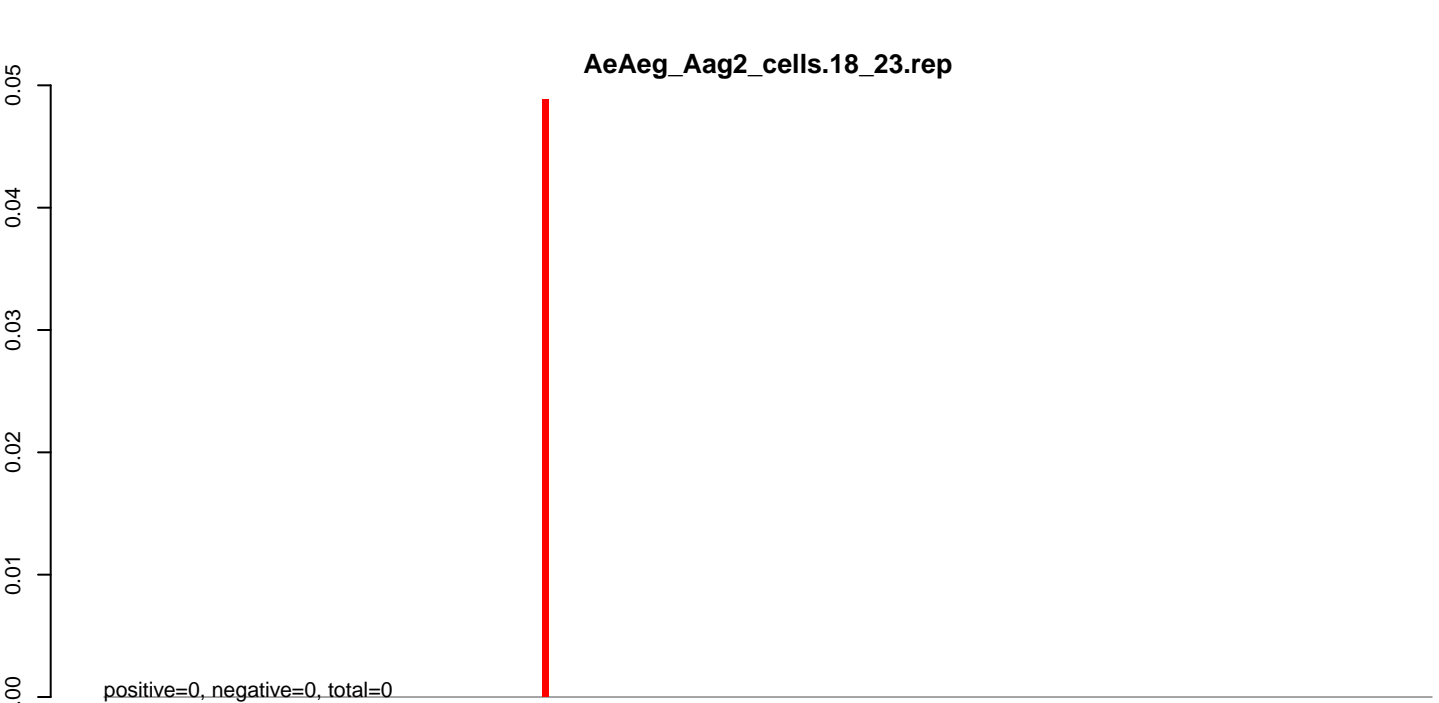
AeAeg_Aag2_cells.rep



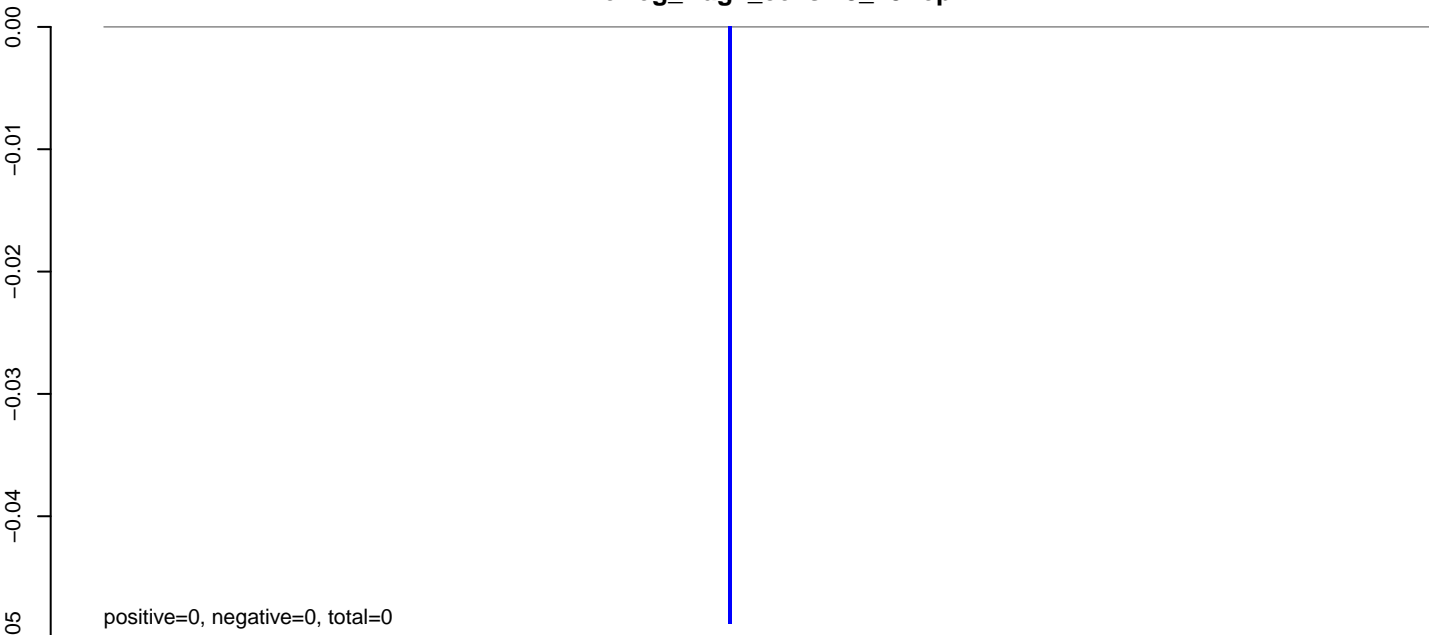
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0 2000 4000 6000 8000 10000



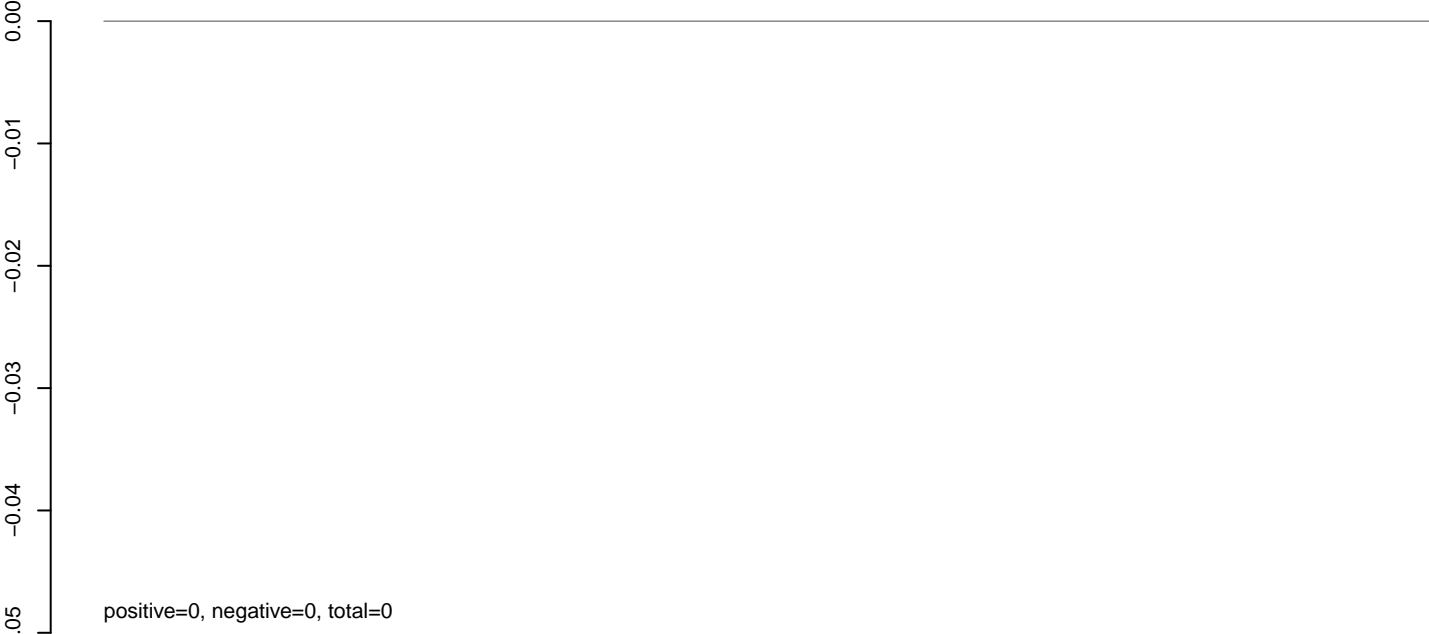


AeAeg_Aag2_cells.18_23.rep



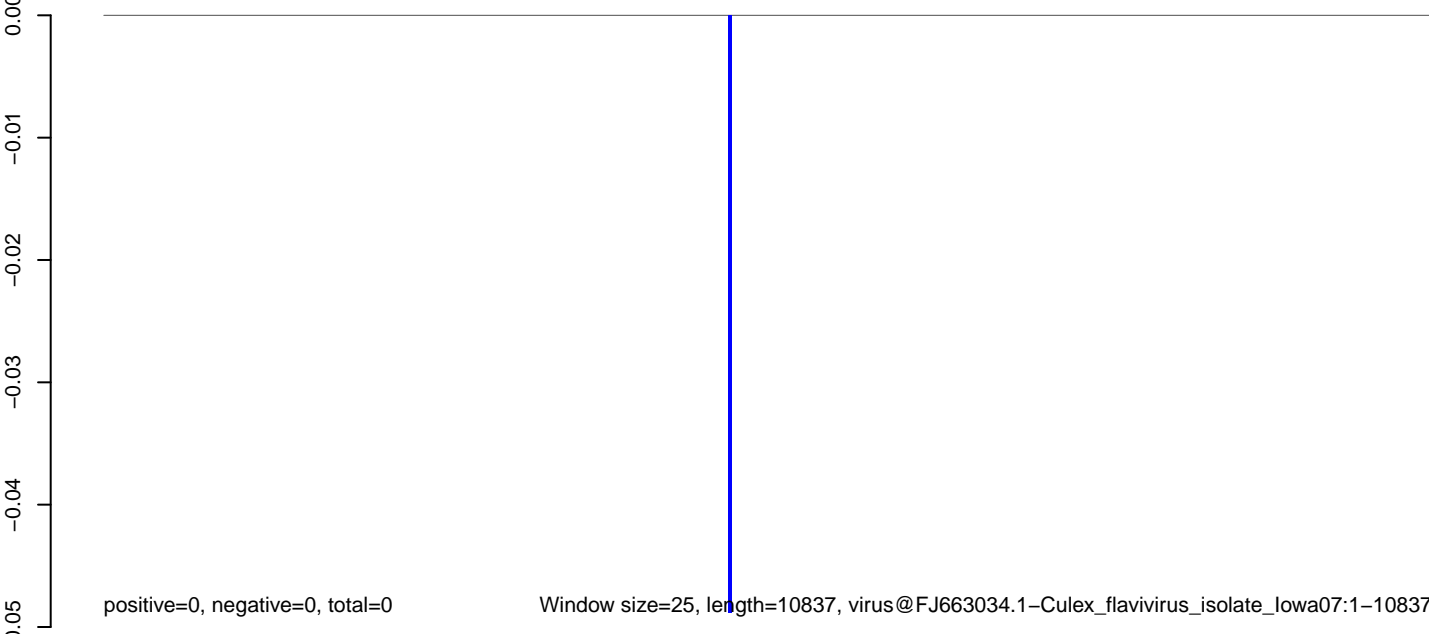
positive=0, negative=0, total=0

AeAeg_Aag2_cells.24_35.rep



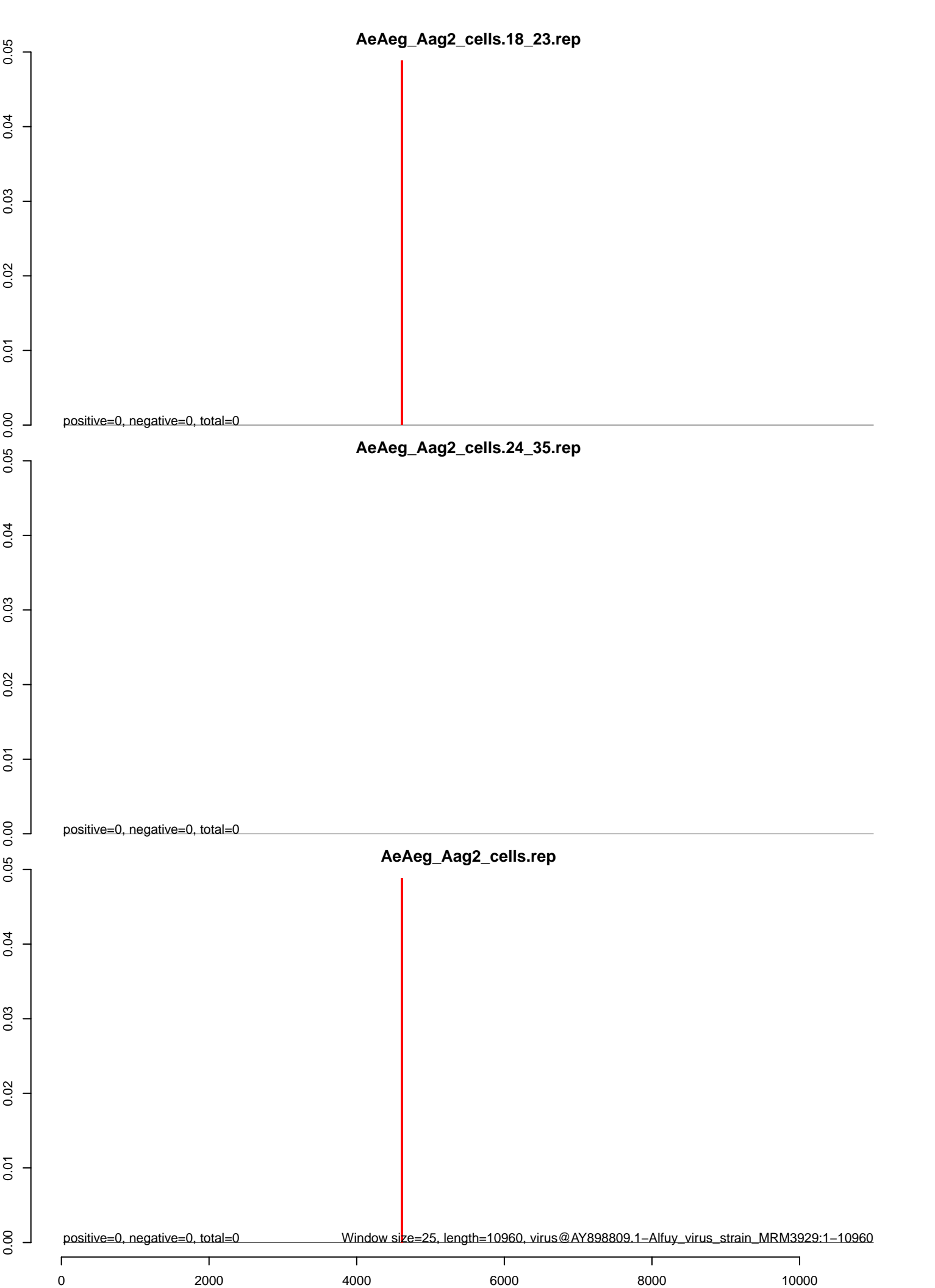
positive=0, negative=0, total=0

AeAeg_Aag2_cells.rep



positive=0, negative=0, total=0

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



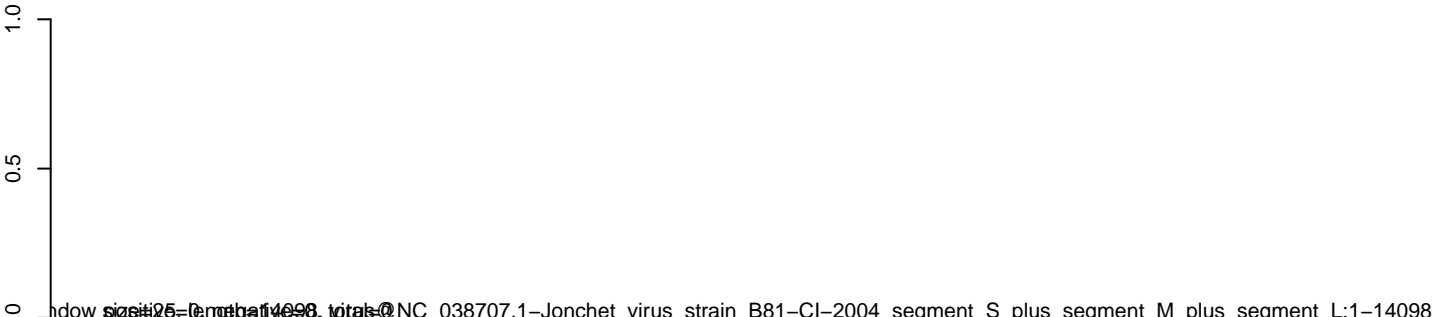
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep



0 2000 4000 6000 8000 10000 12000 14000

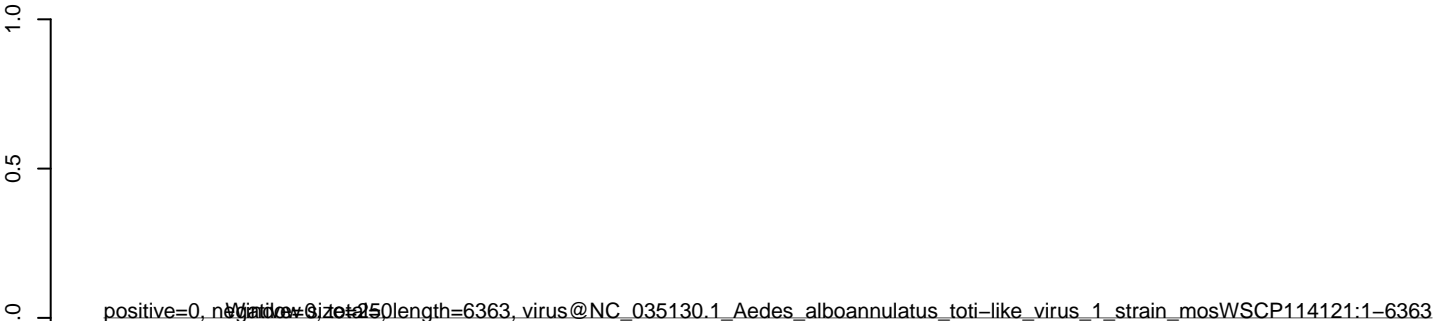
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AeAeg_Aag2_cells.24_35.rep

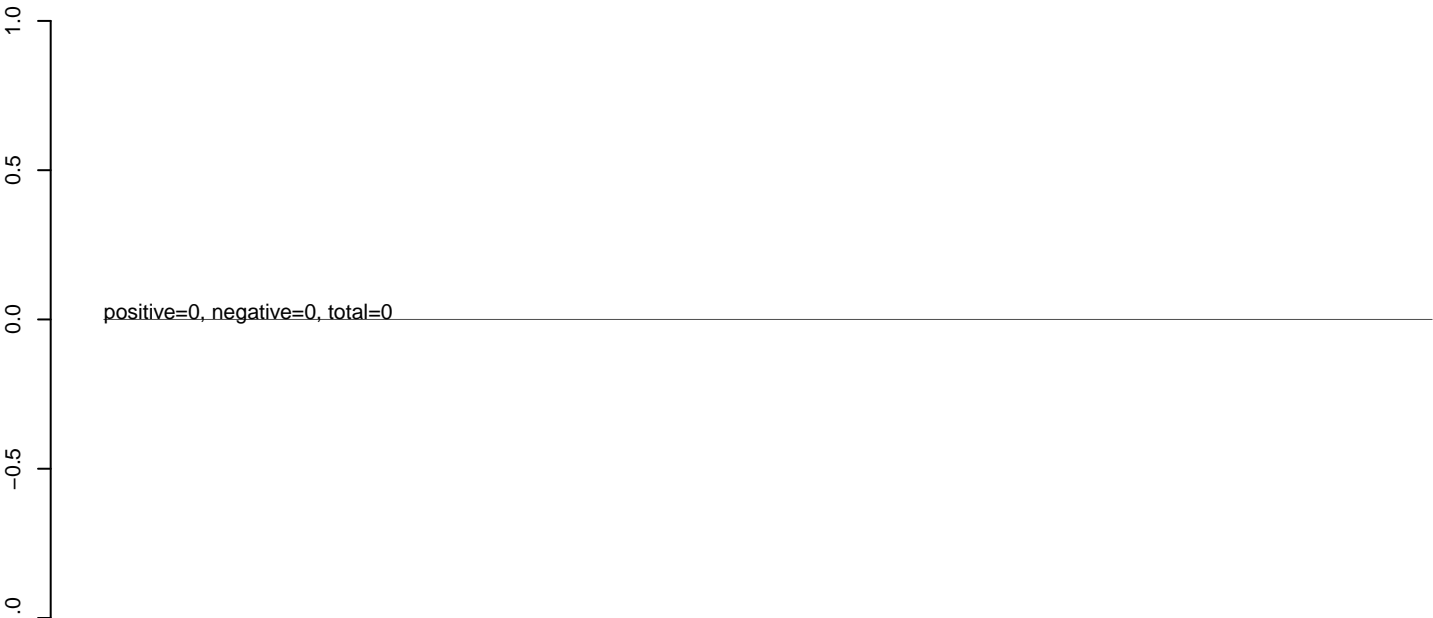


AeAeg_Aag2_cells.rep

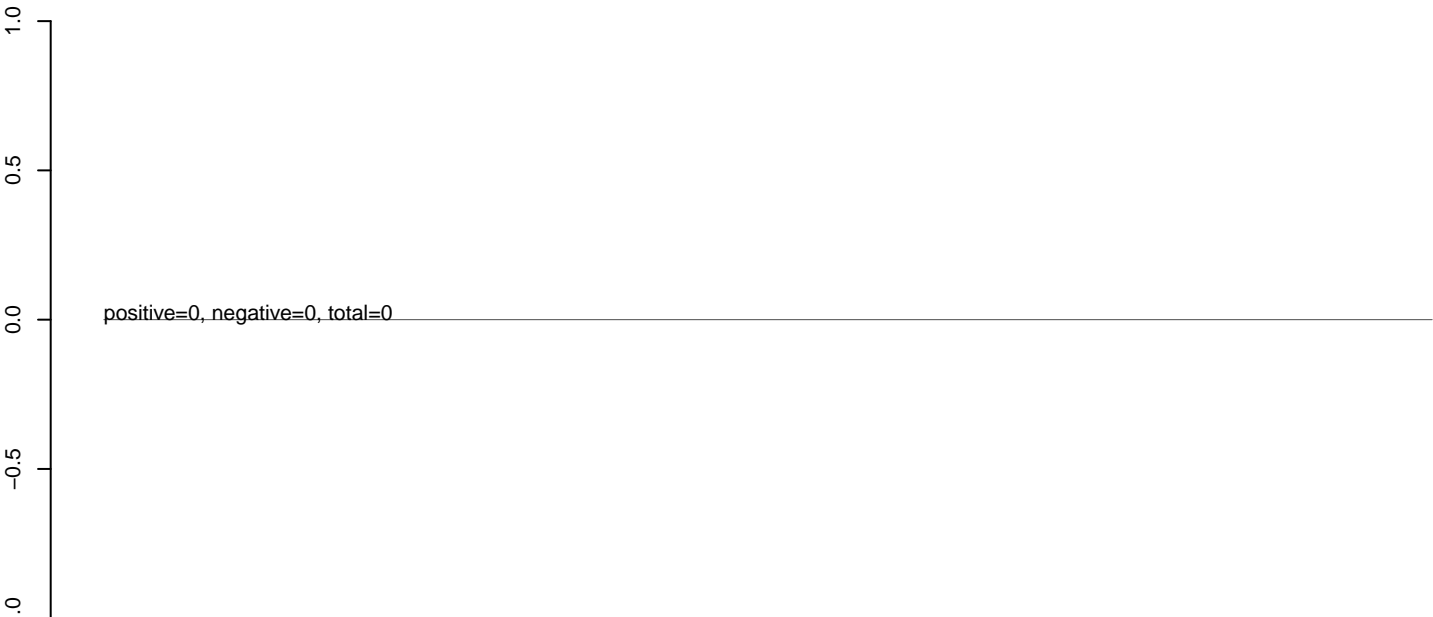


0 1000 2000 3000 4000 5000 6000

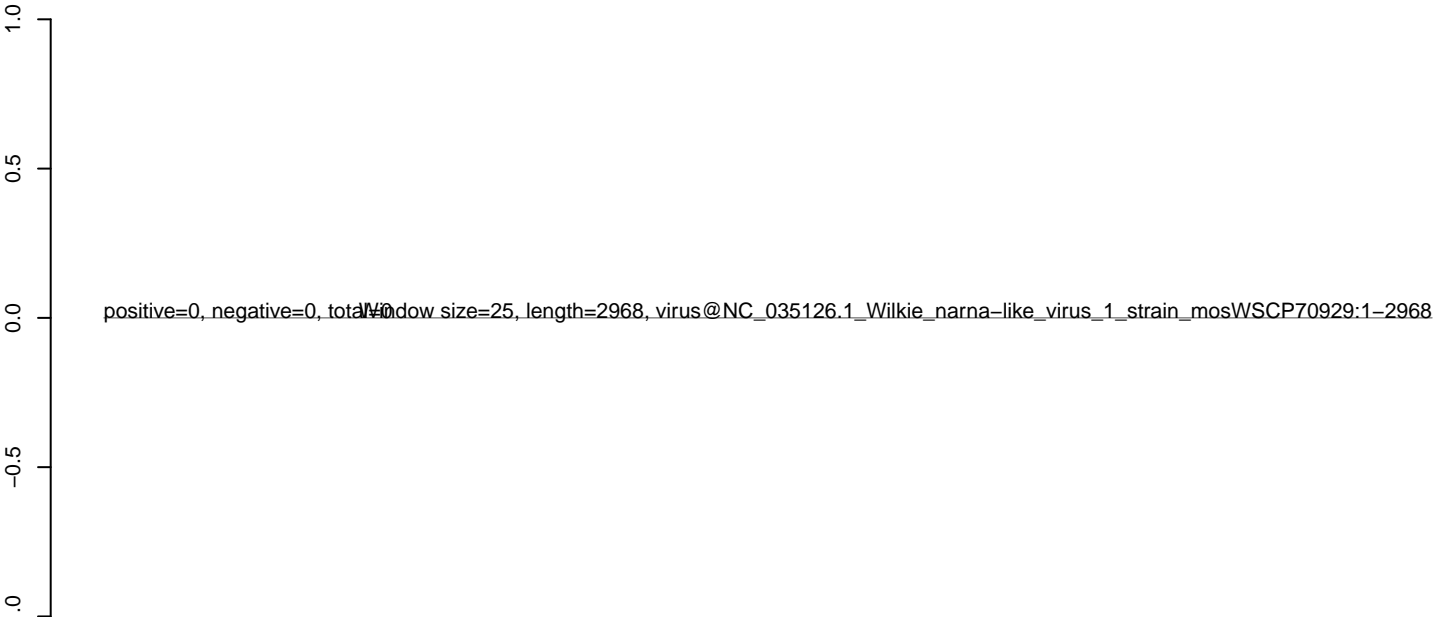
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep



0 500 1000 1500 2000 2500 3000

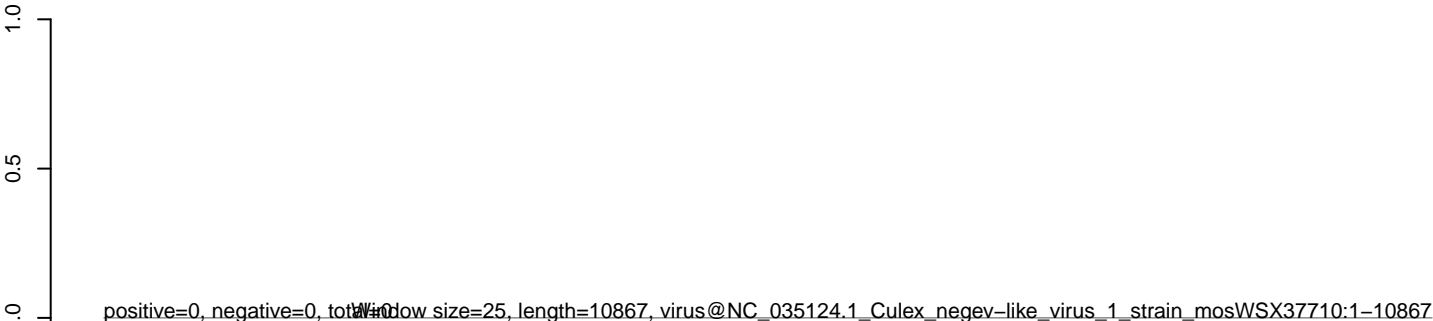
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep



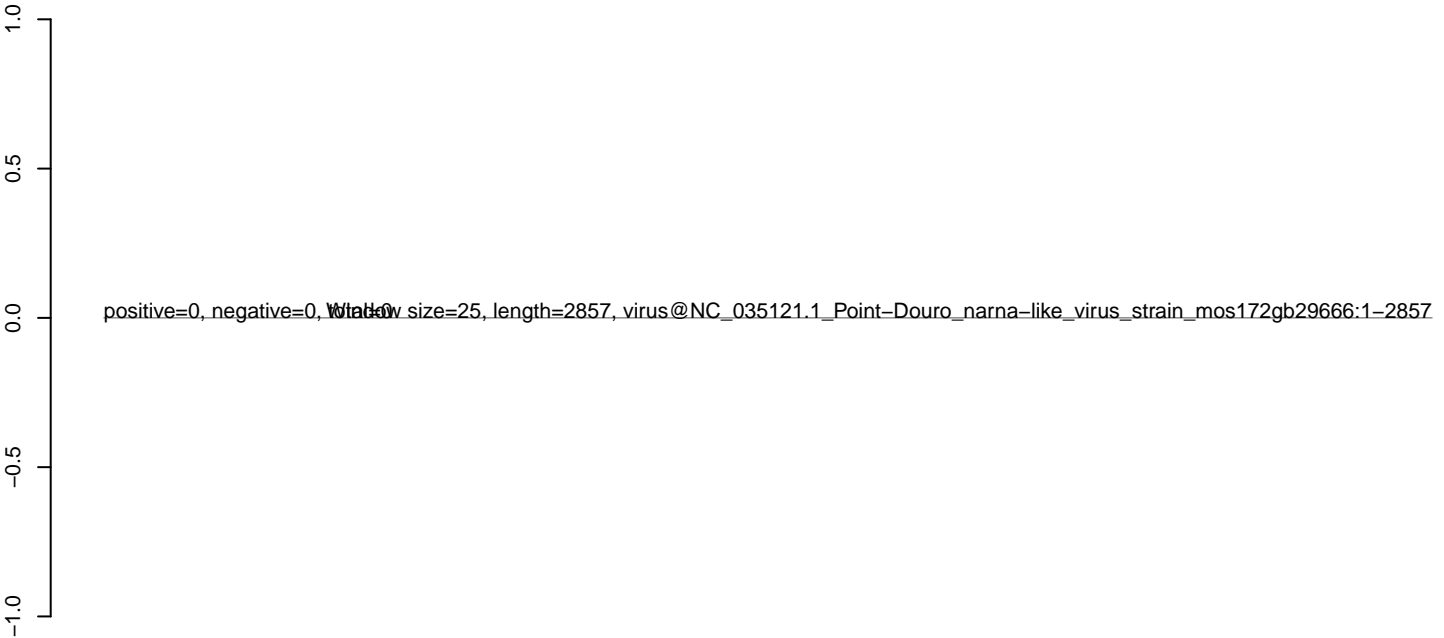
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

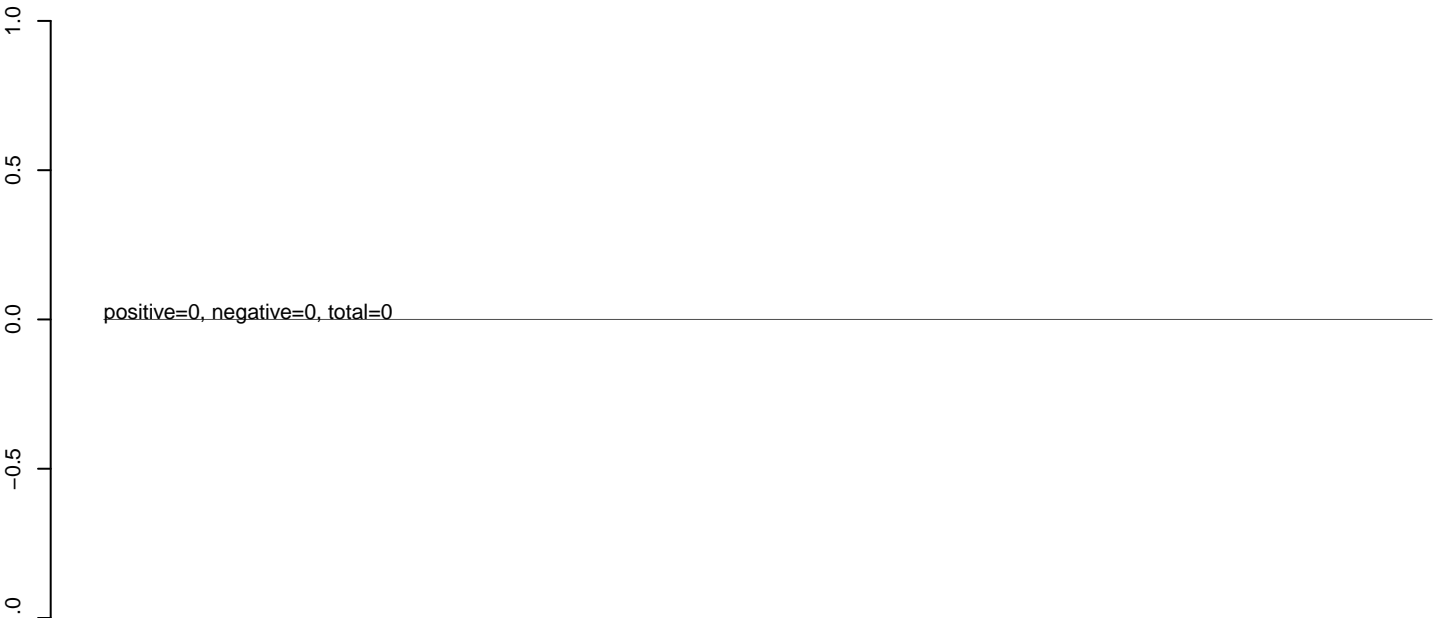


AeAeg_Aag2_cells.rep

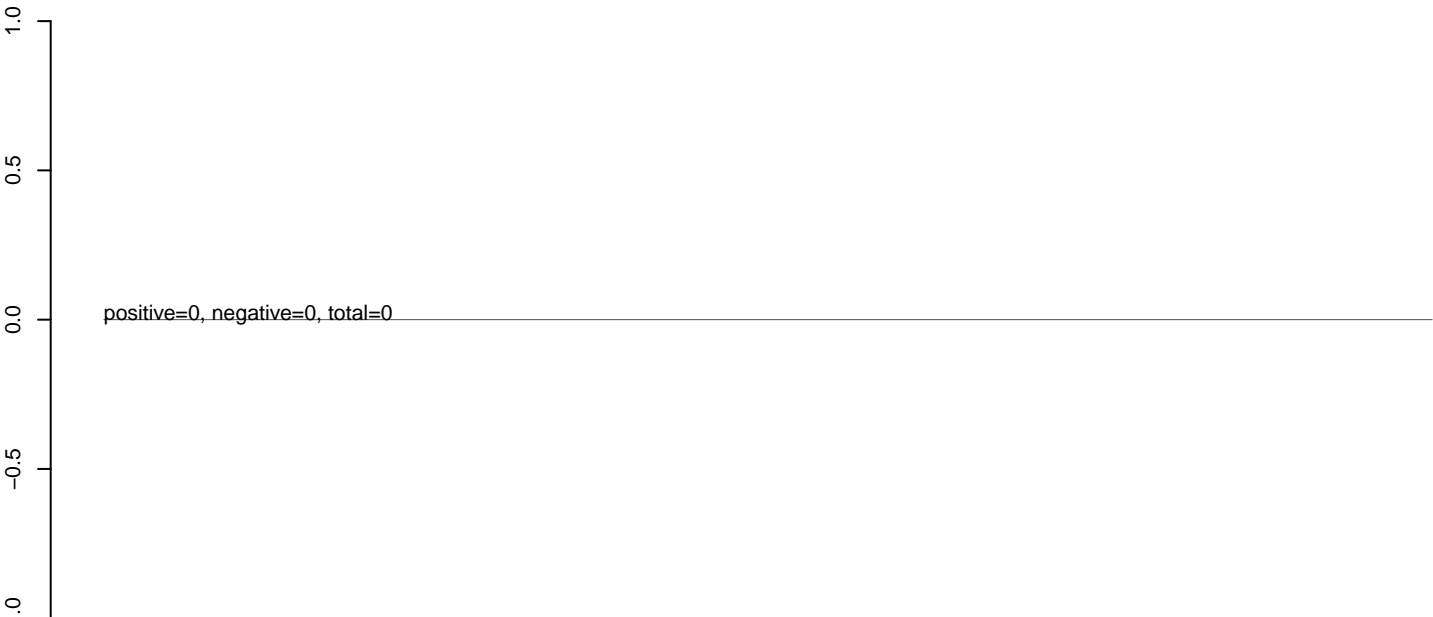


0 500 1000 1500 2000 2500 3000

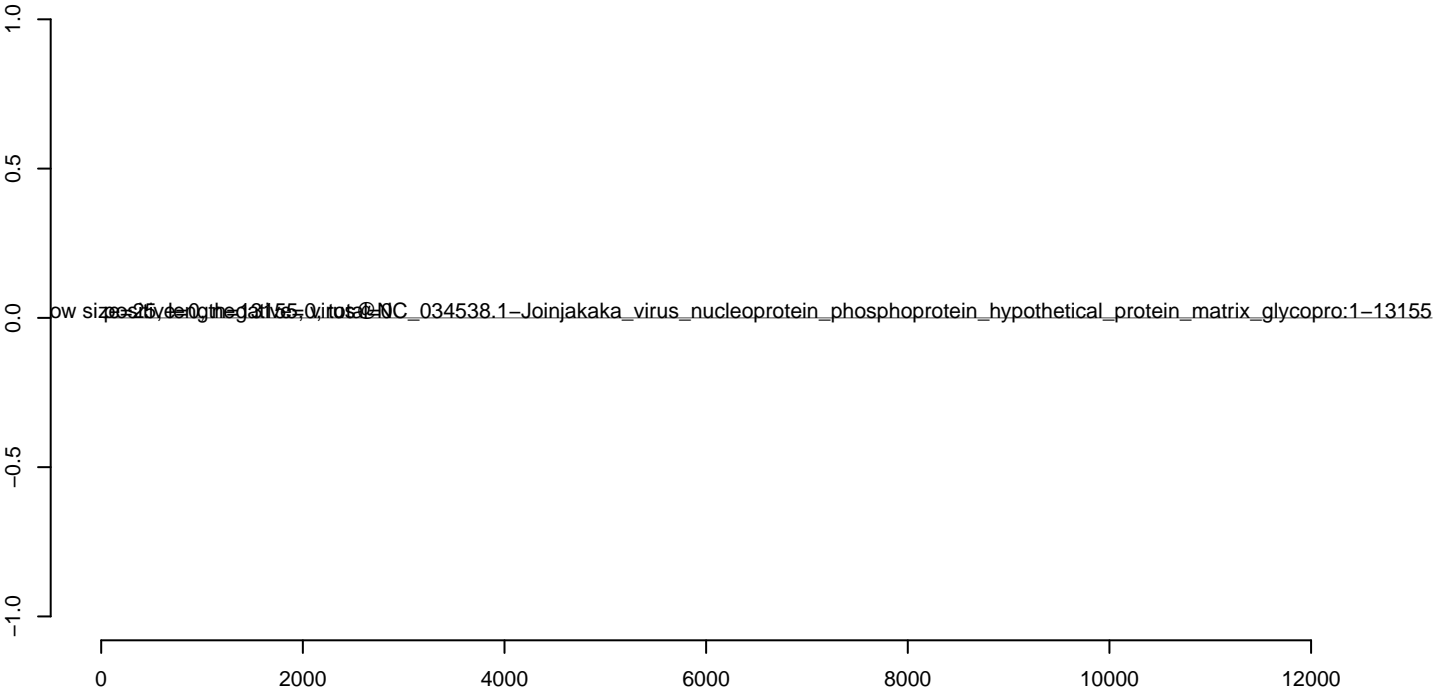
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep



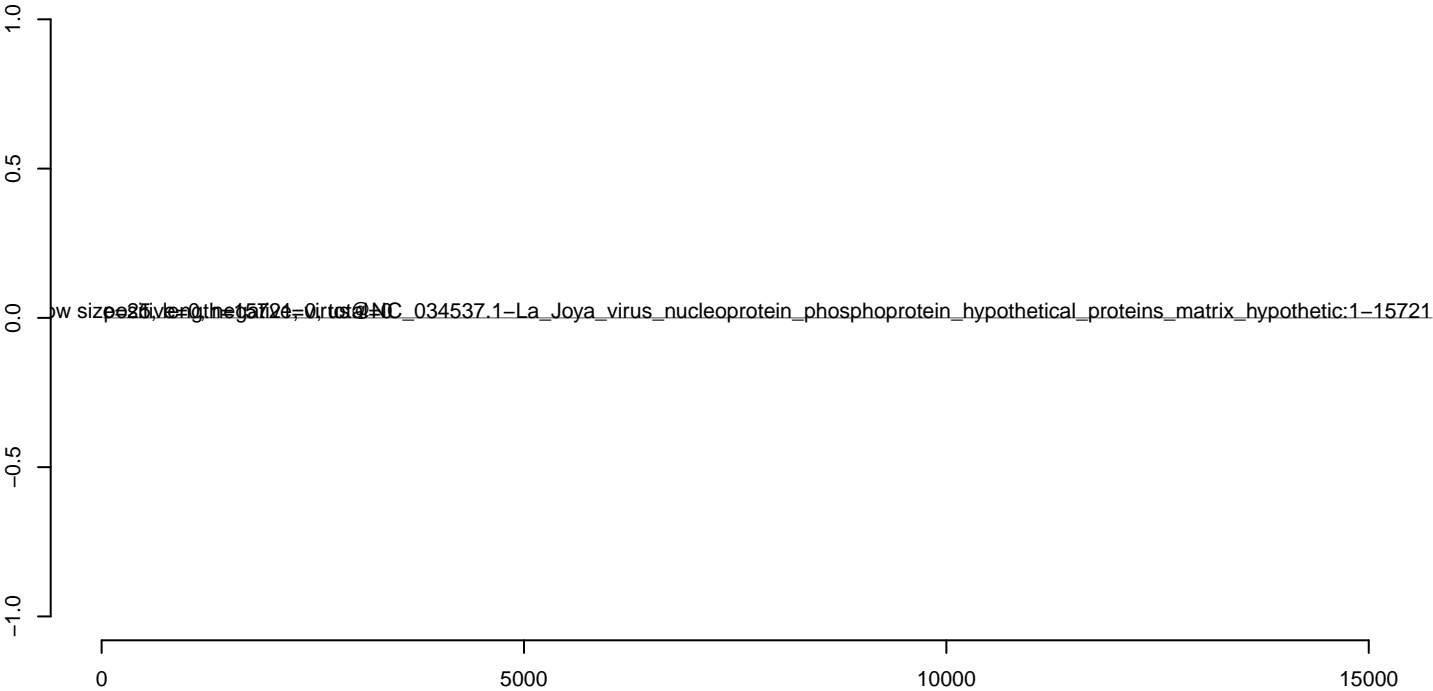
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AeAeg_Aag2_cells.24_35.rep



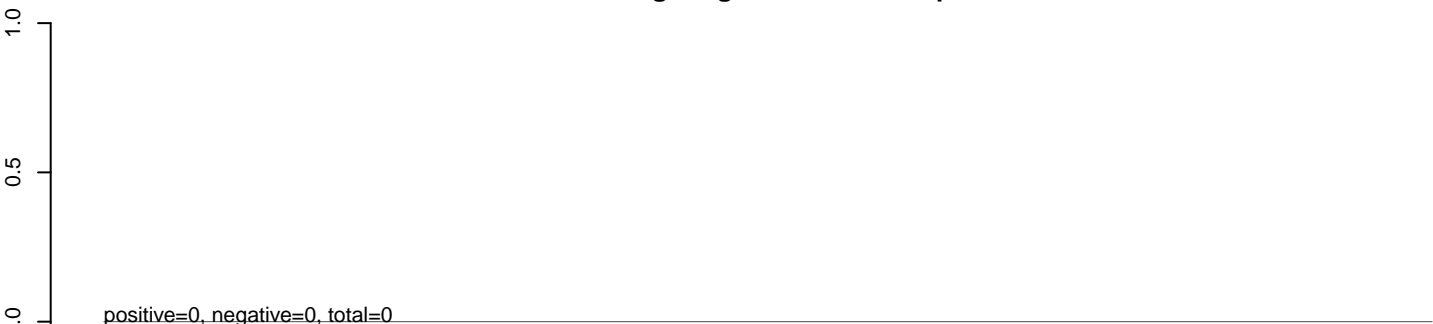
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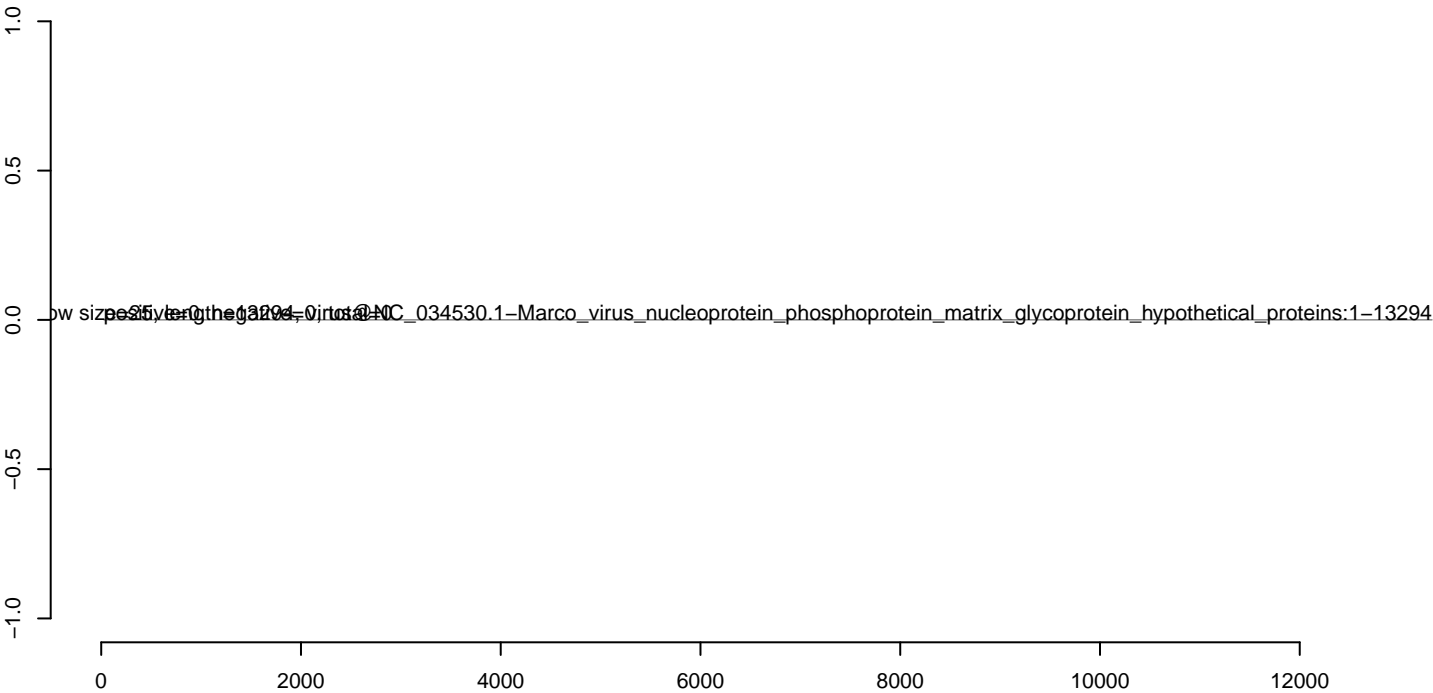
AeAeg_Aag2_cells.18_23.rep



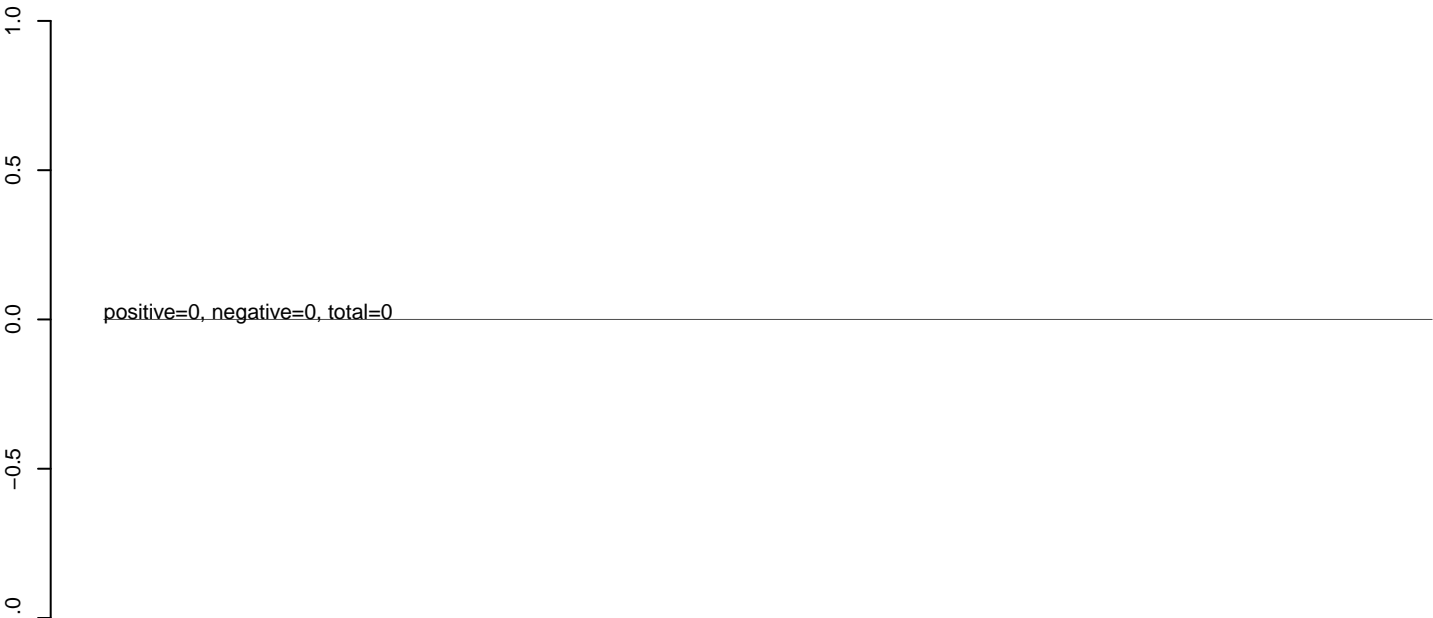
AeAeg_Aag2_cells.24_35.rep



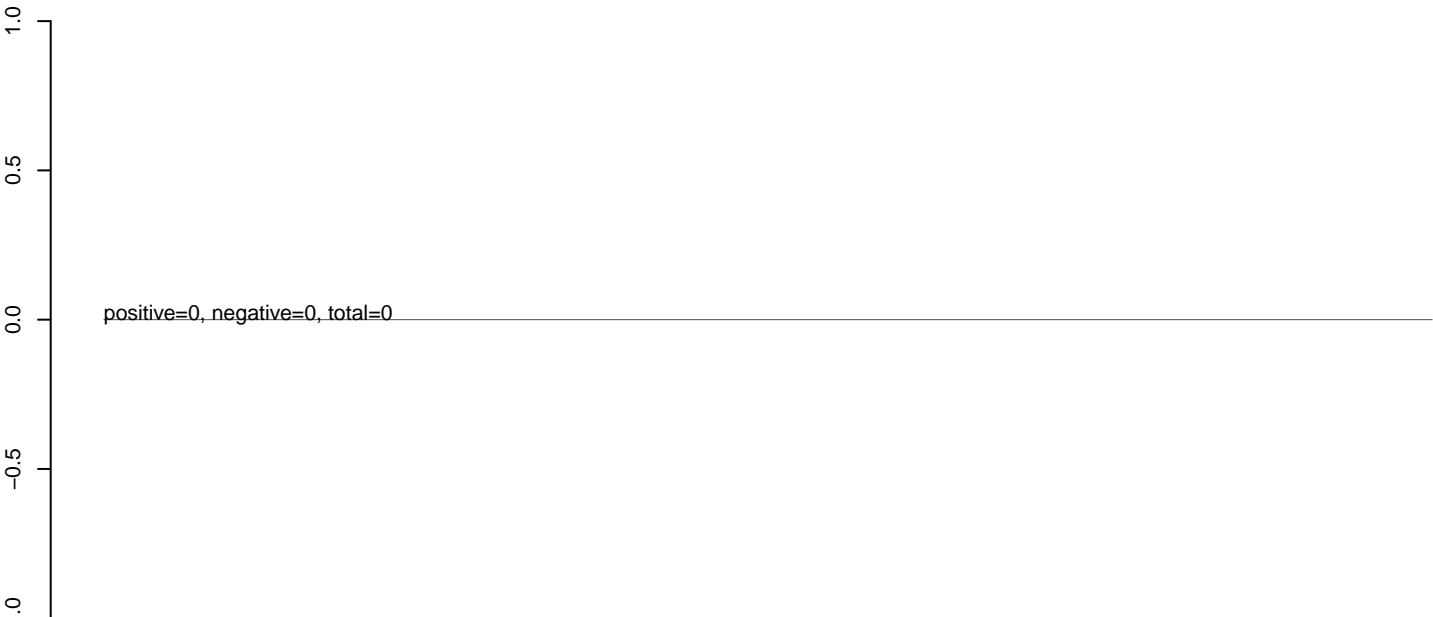
AeAeg_Aag2_cells.rep



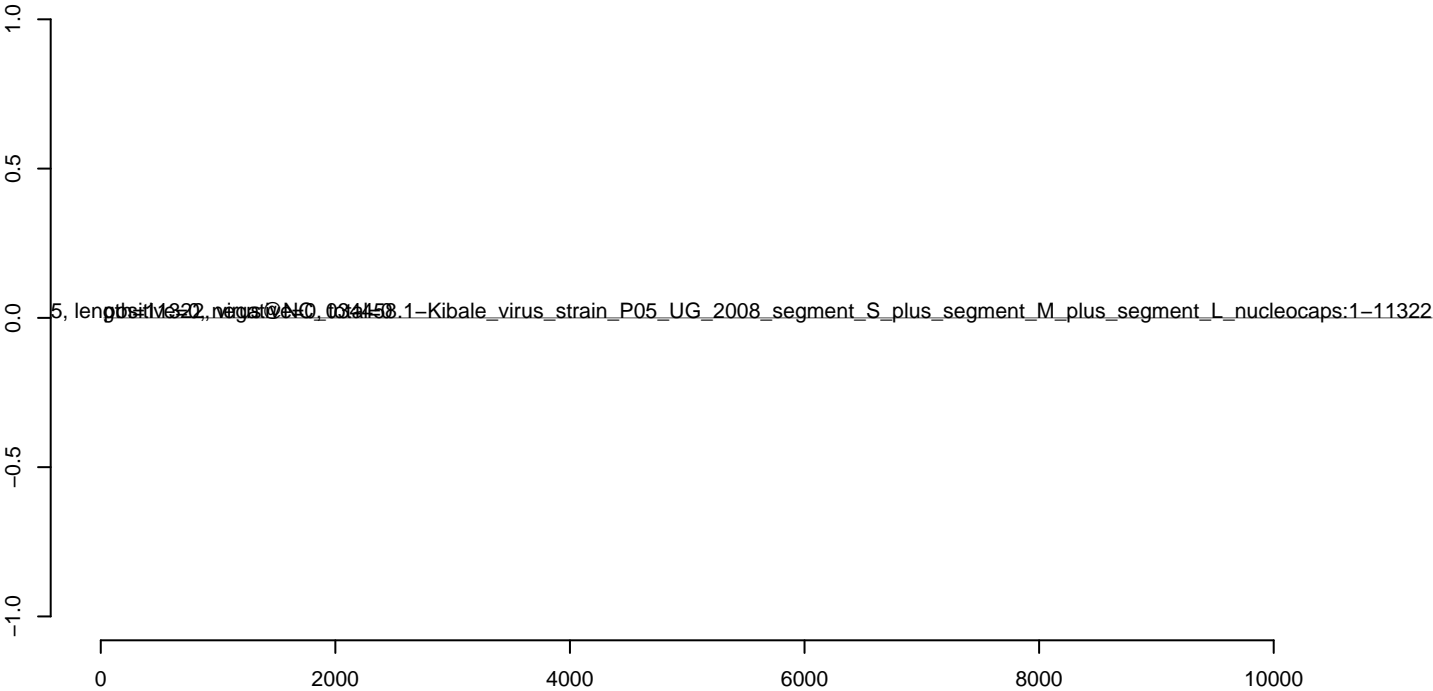
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep



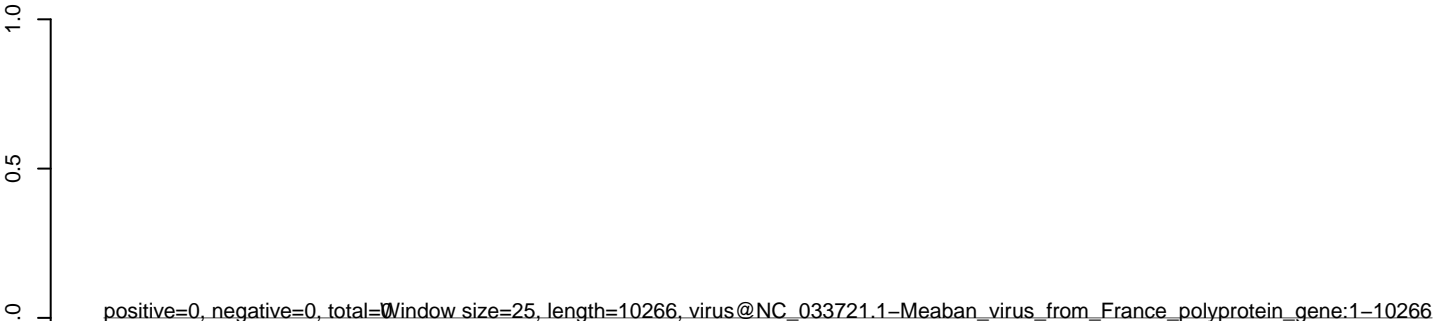
AeAeg_Aag2_cells.18_23.rep



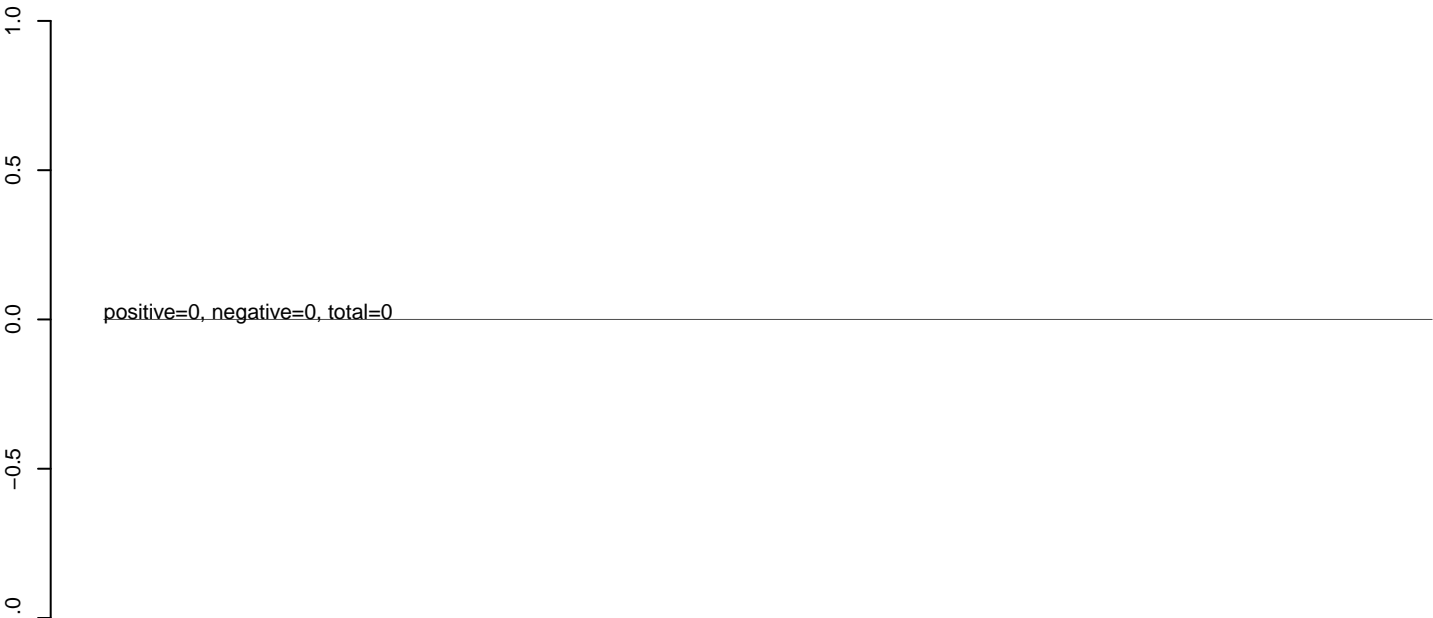
AeAeg_Aag2_cells.24_35.rep



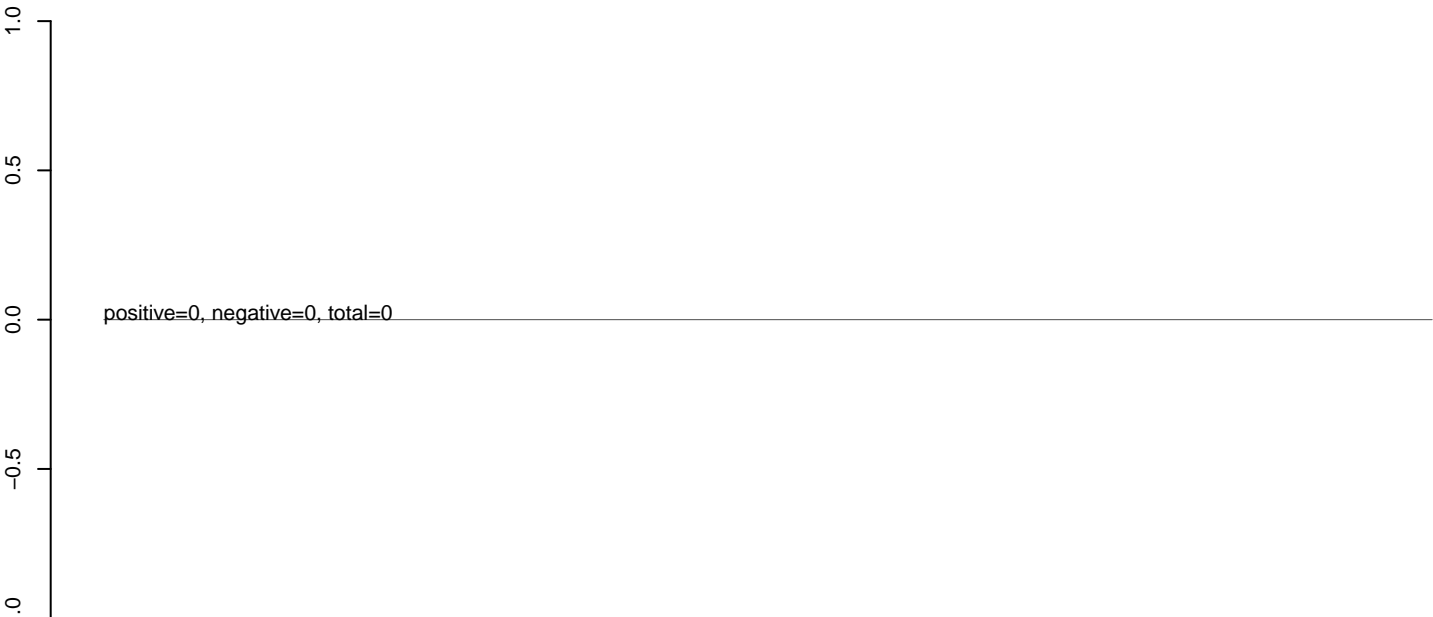
AeAeg_Aag2_cells.rep



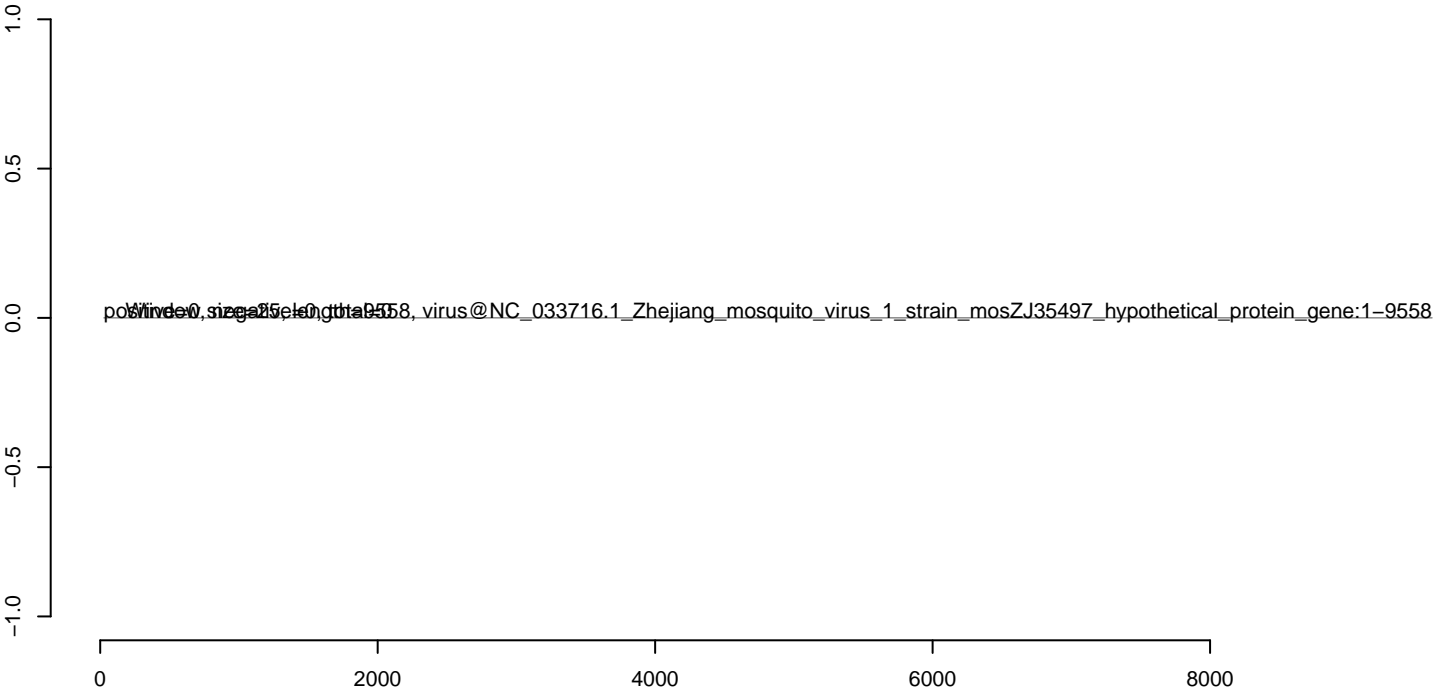
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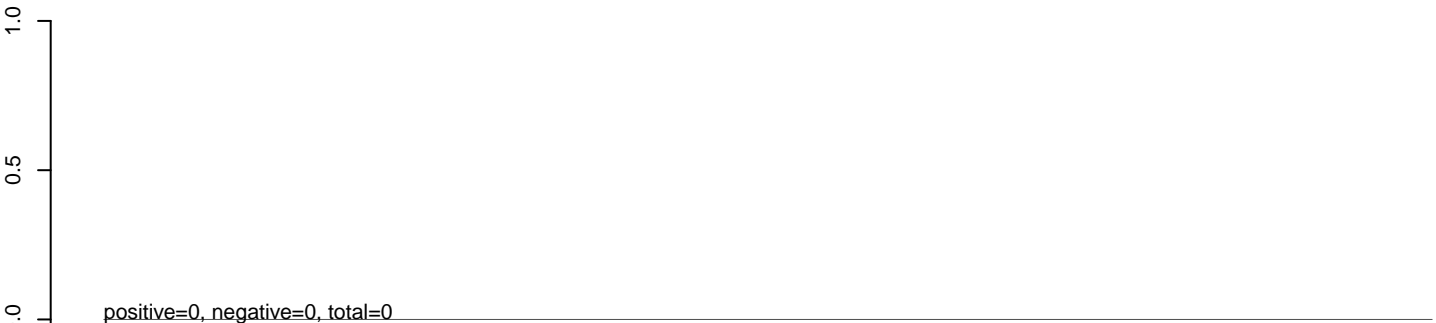
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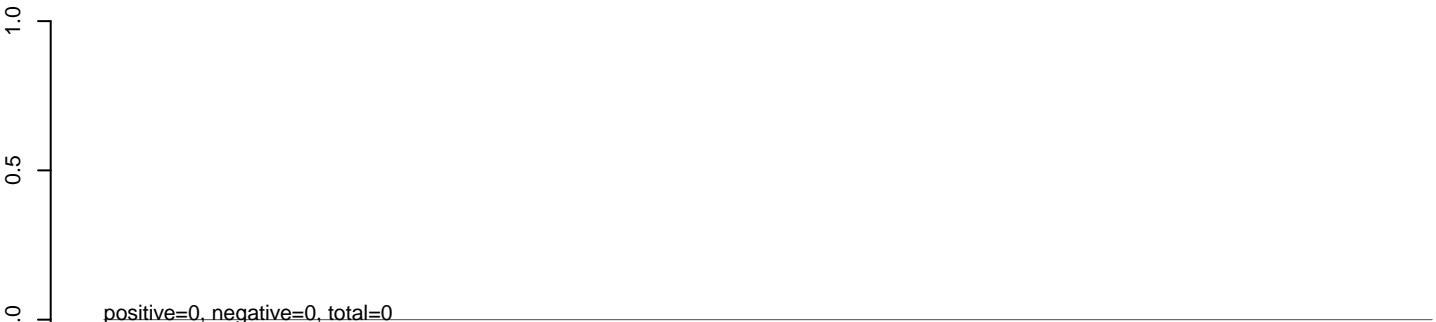
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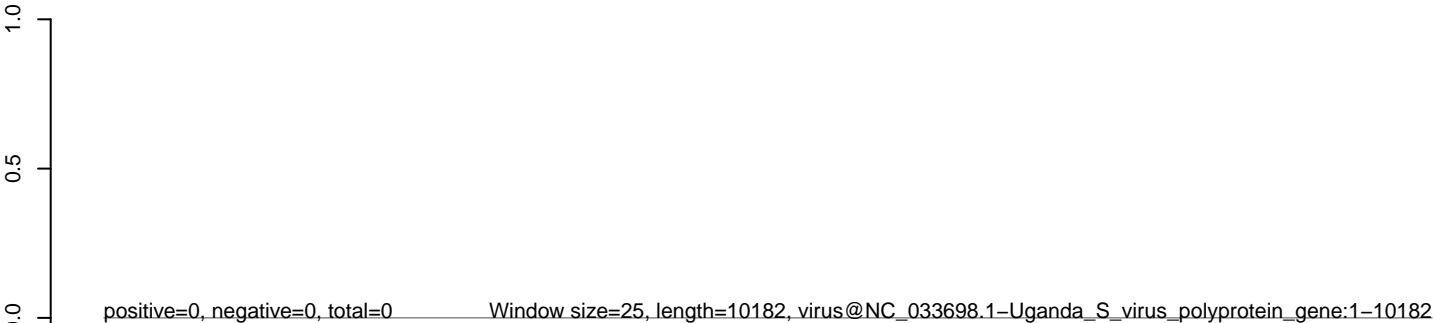
AeAeg_Aag2_cells.18_23.rep



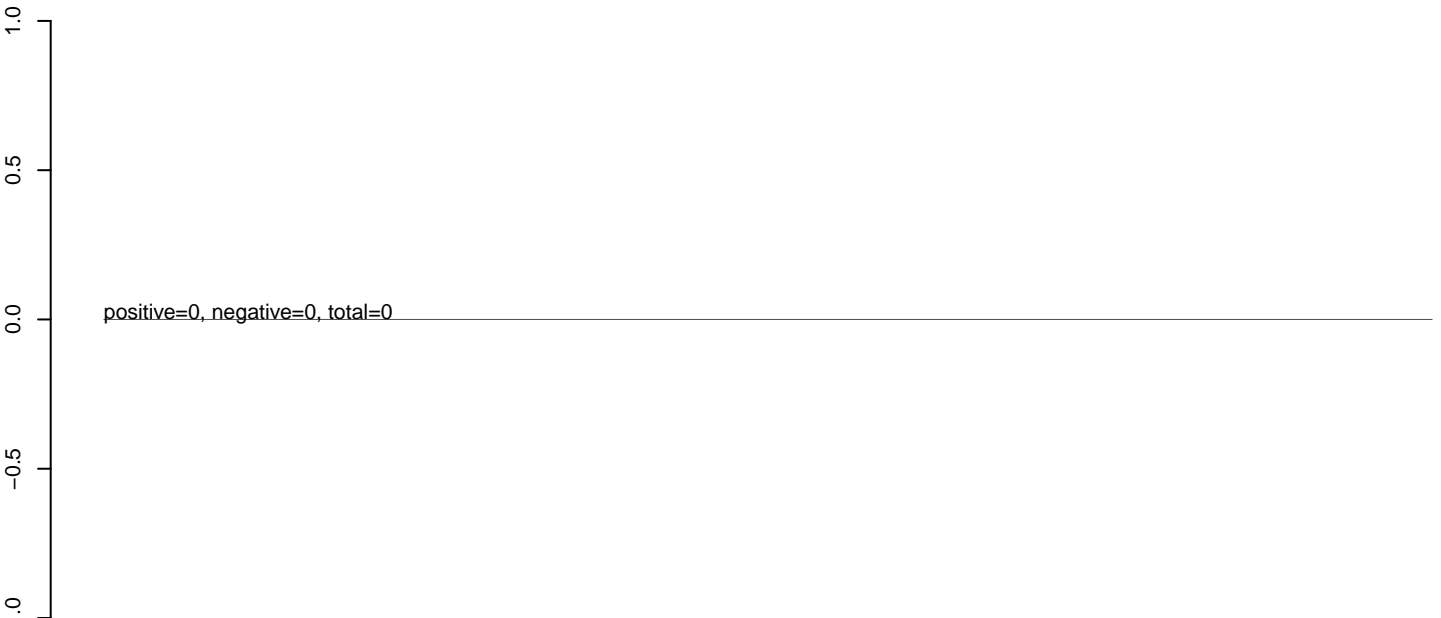
AeAeg_Aag2_cells.24_35.rep



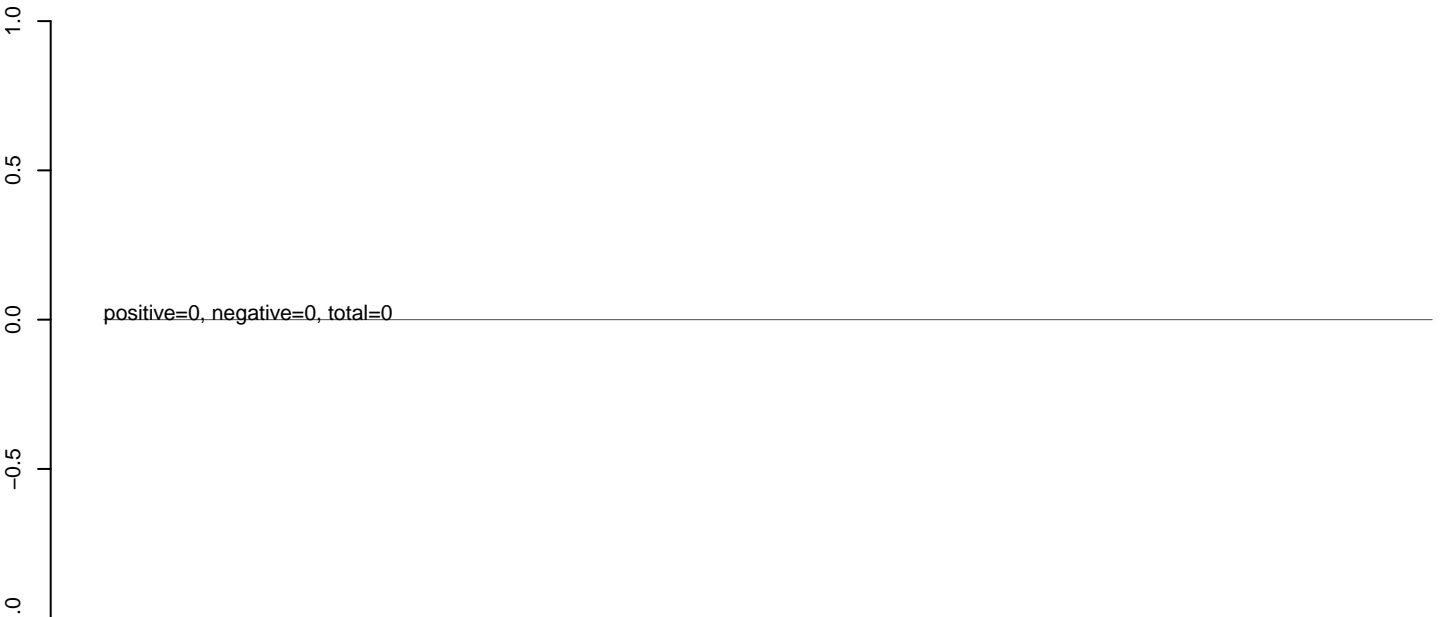
AeAeg_Aag2_cells.rep



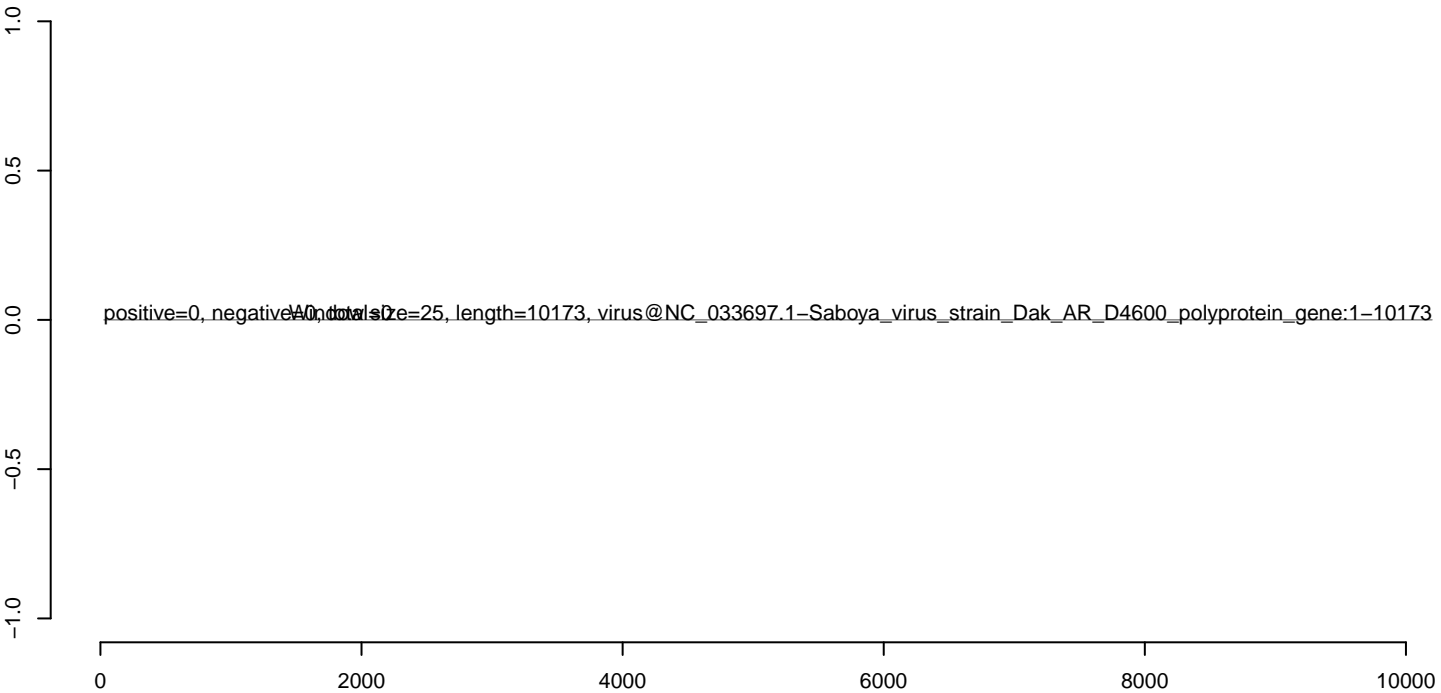
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep



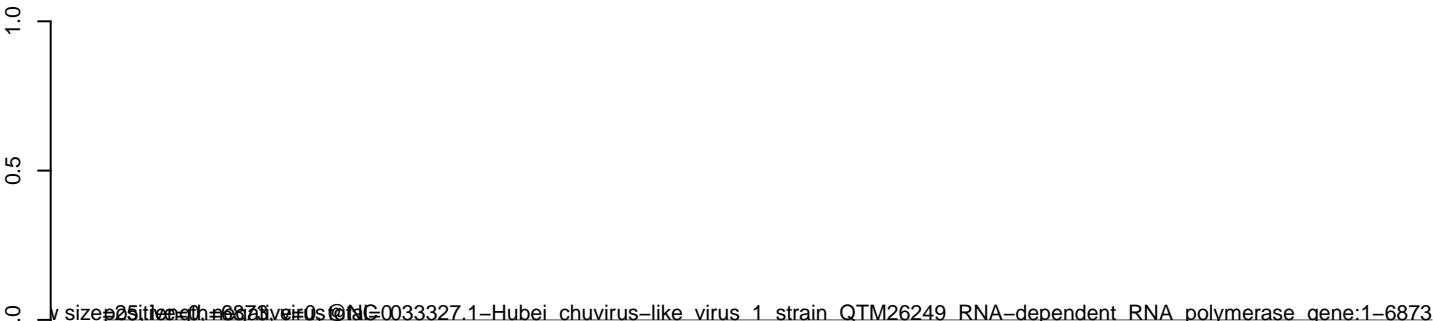
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep

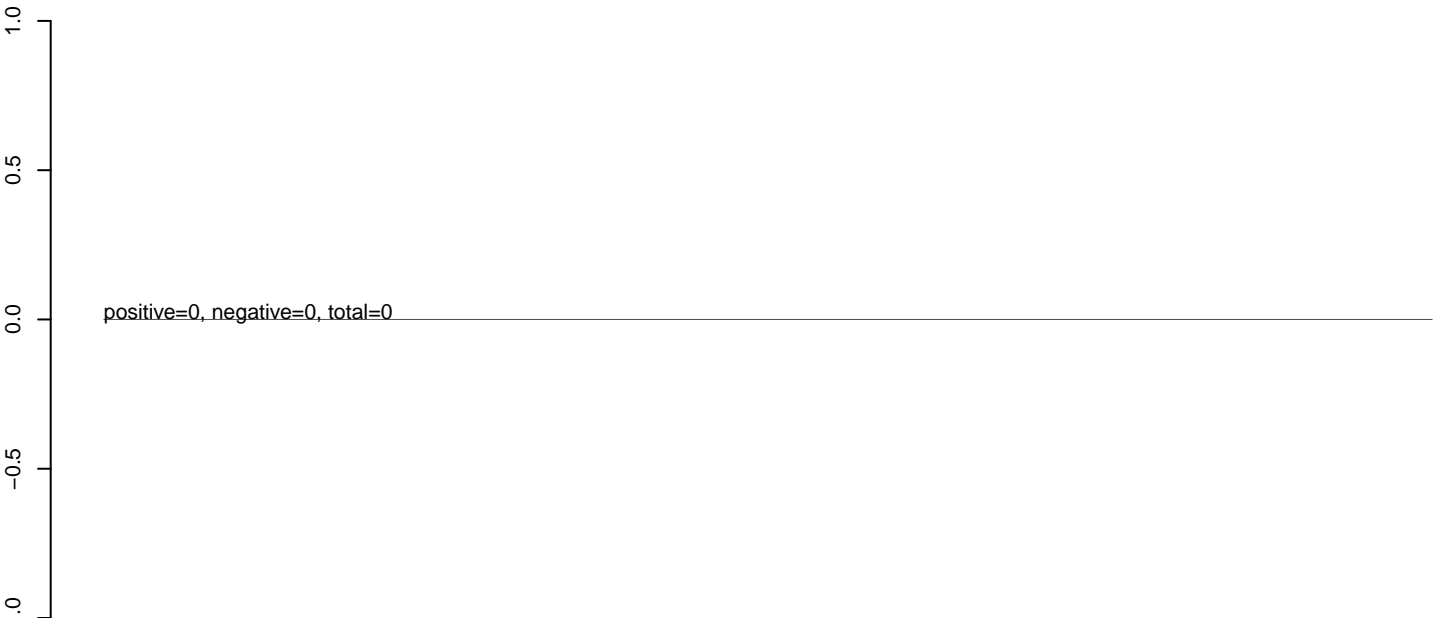


AeAeg_Aag2_cells.rep

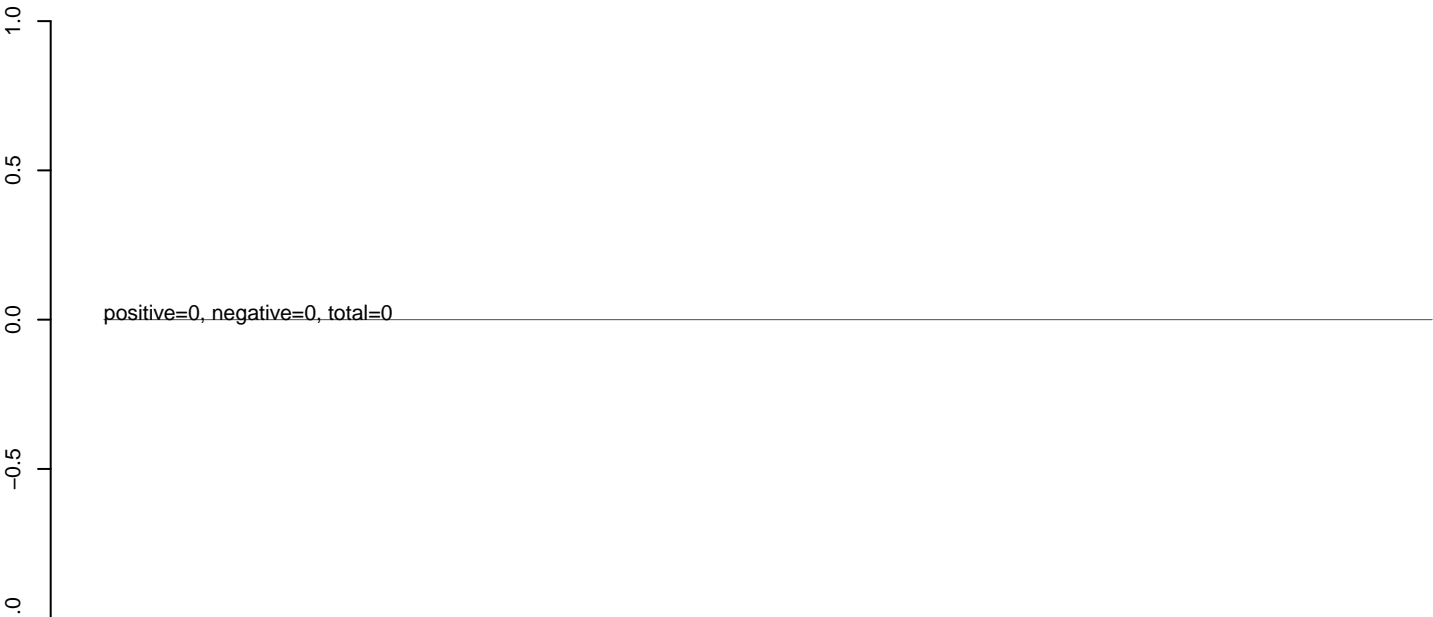


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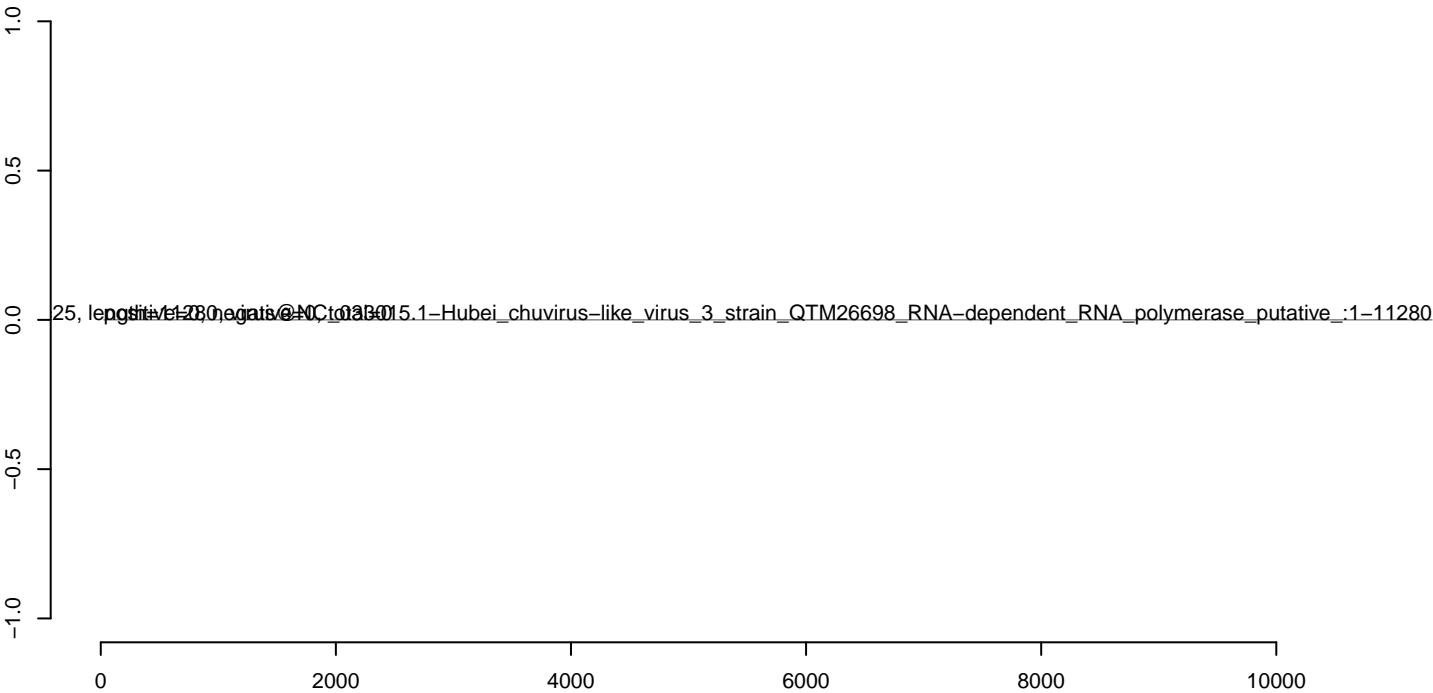
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep



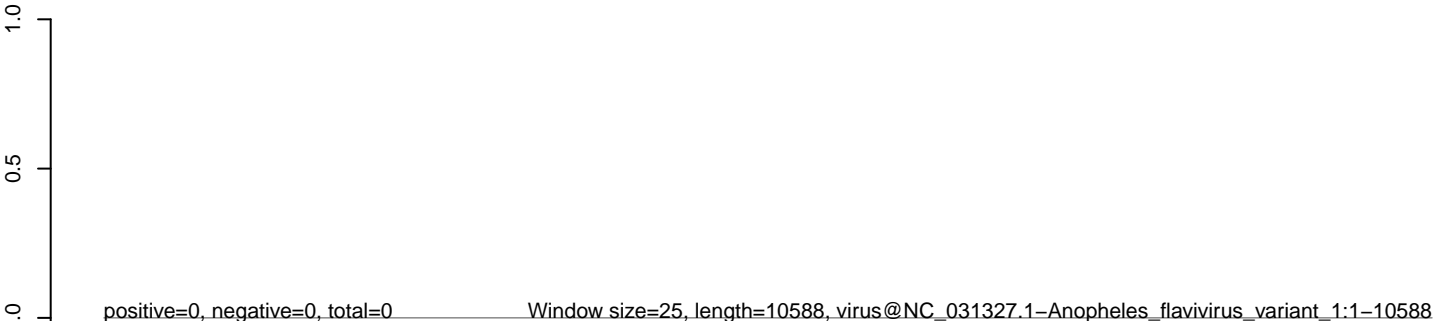
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep



0 2000 4000 6000 8000 10000

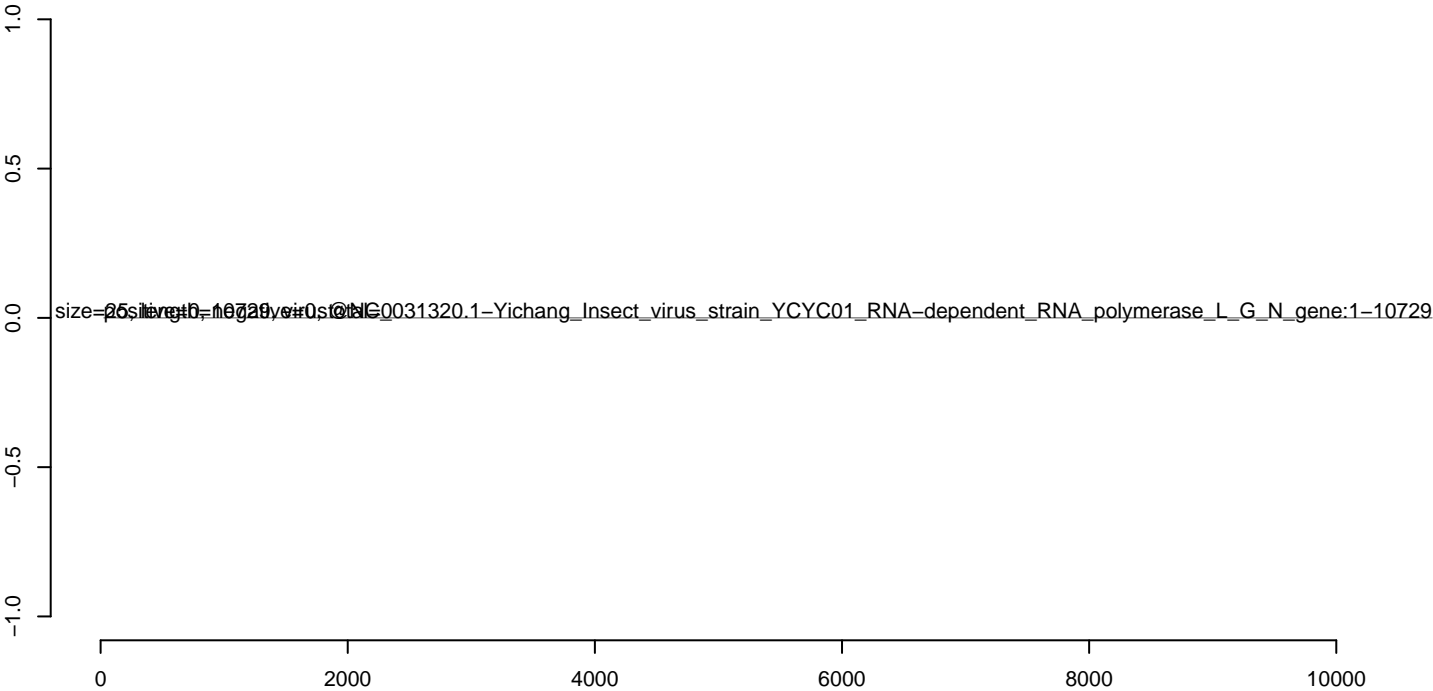
AeAeg_Aag2_cells.18_23.rep



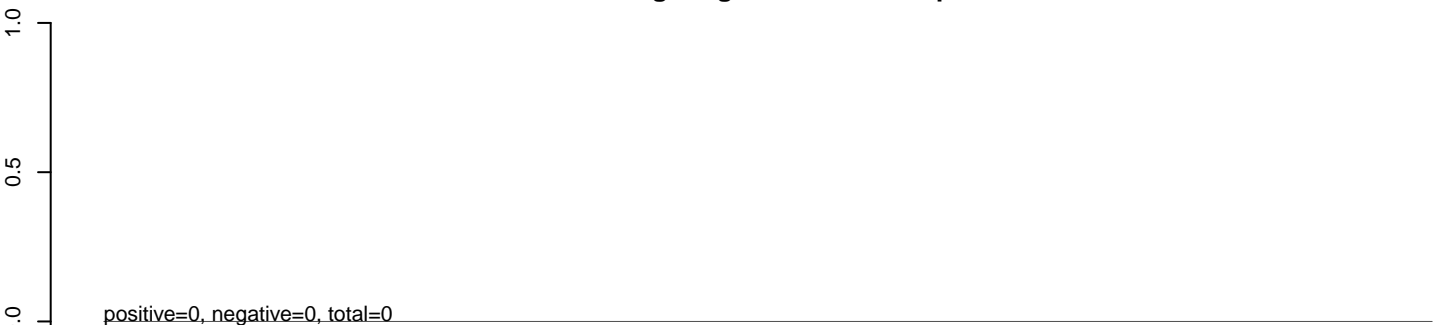
AeAeg_Aag2_cells.24_35.rep



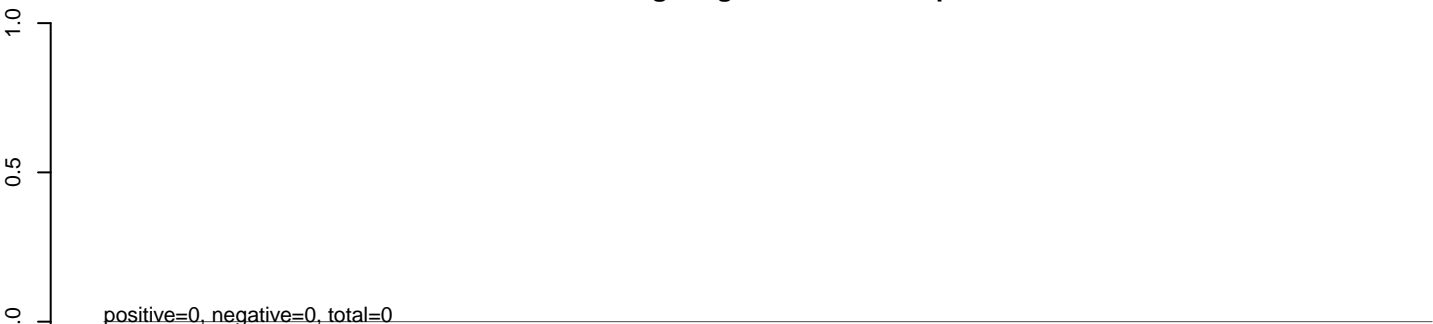
AeAeg_Aag2_cells.rep



AeAeg_Aag2_cells.18_23.rep



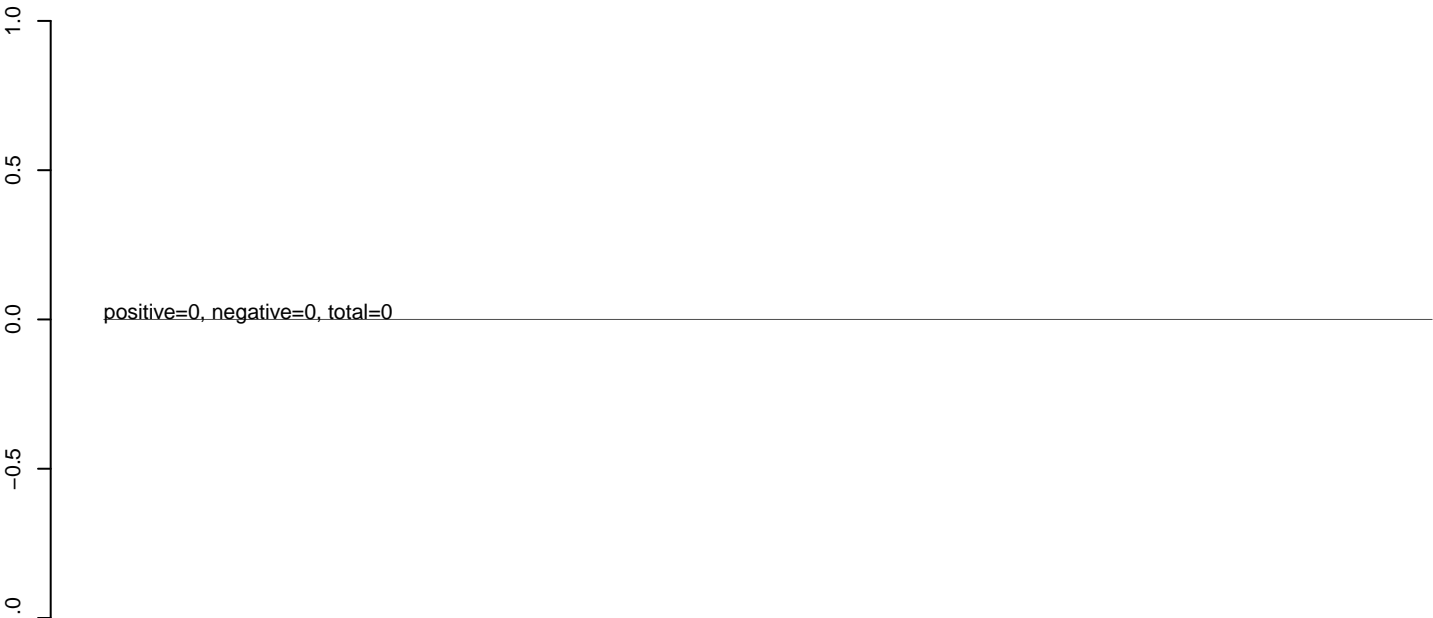
AeAeg_Aag2_cells.24_35.rep



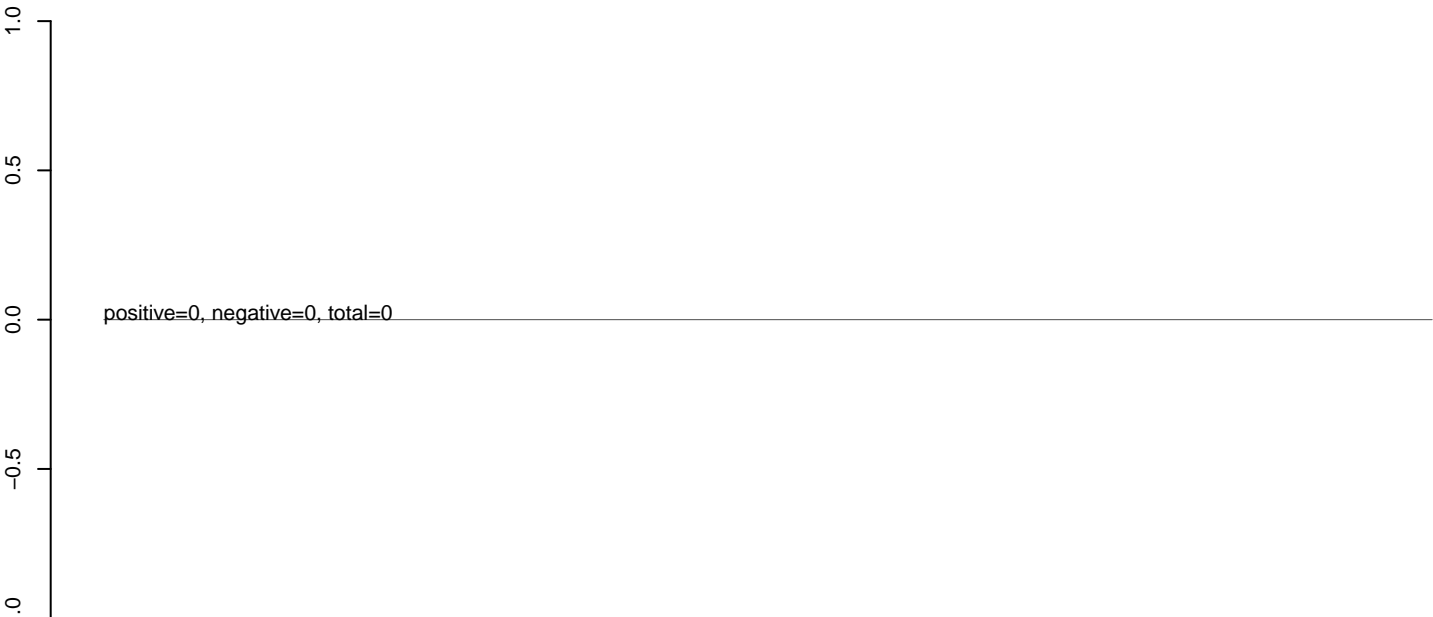
AeAeg_Aag2_cells.rep



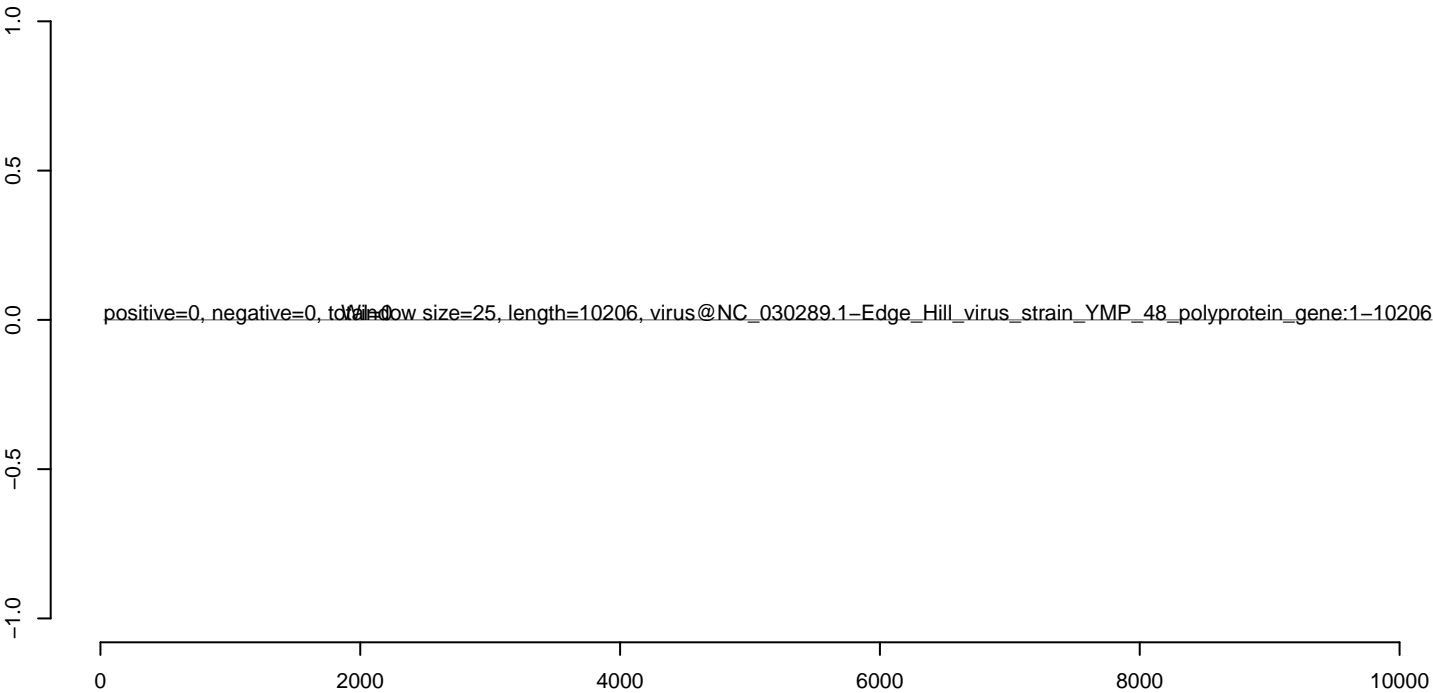
AeAeg_Aag2_cells.18_23.rep



AeAeg_Aag2_cells.24_35.rep



AeAeg_Aag2_cells.rep



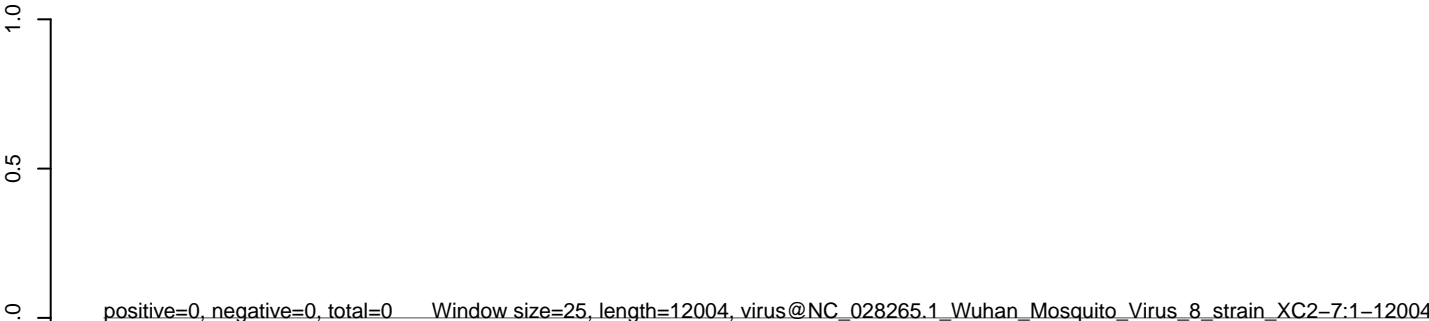
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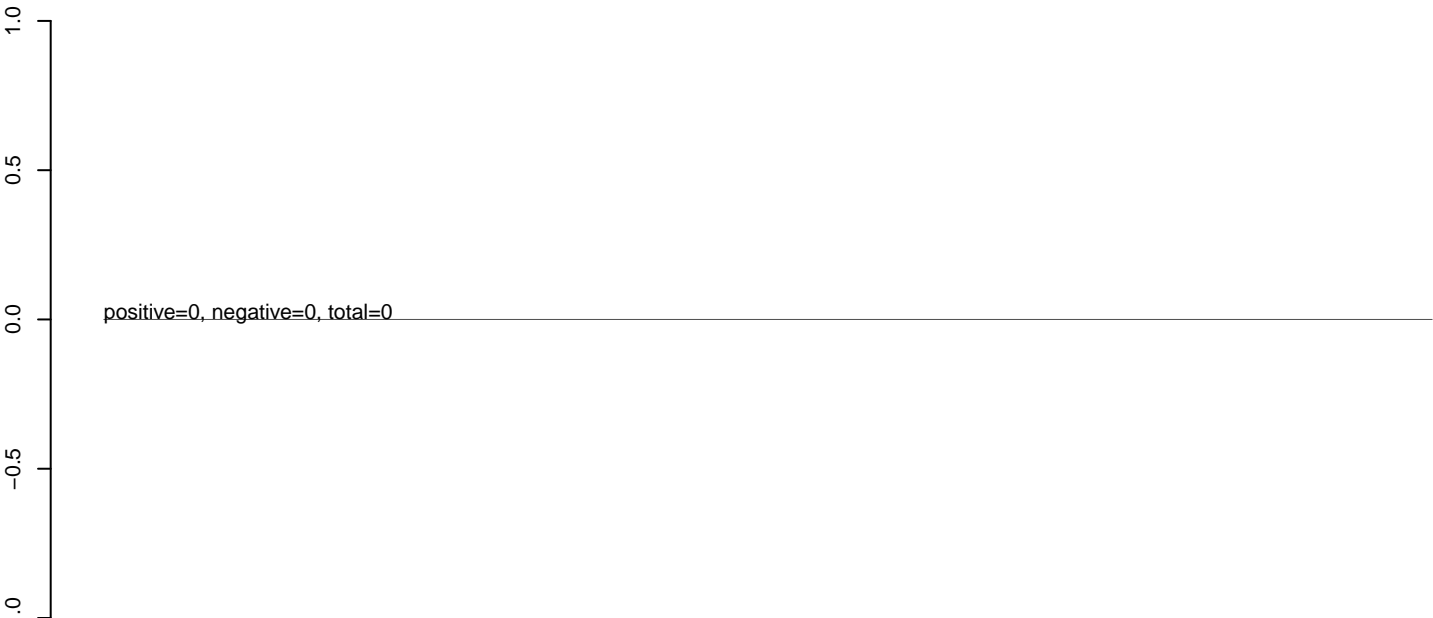
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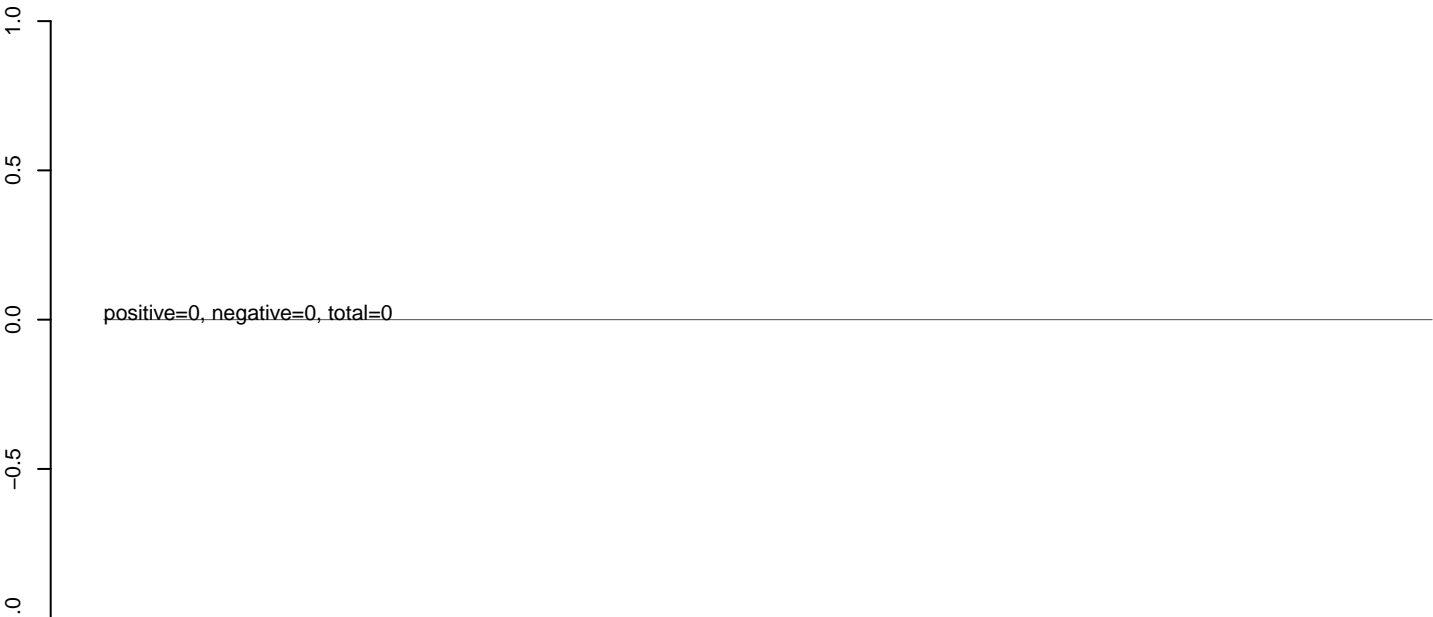
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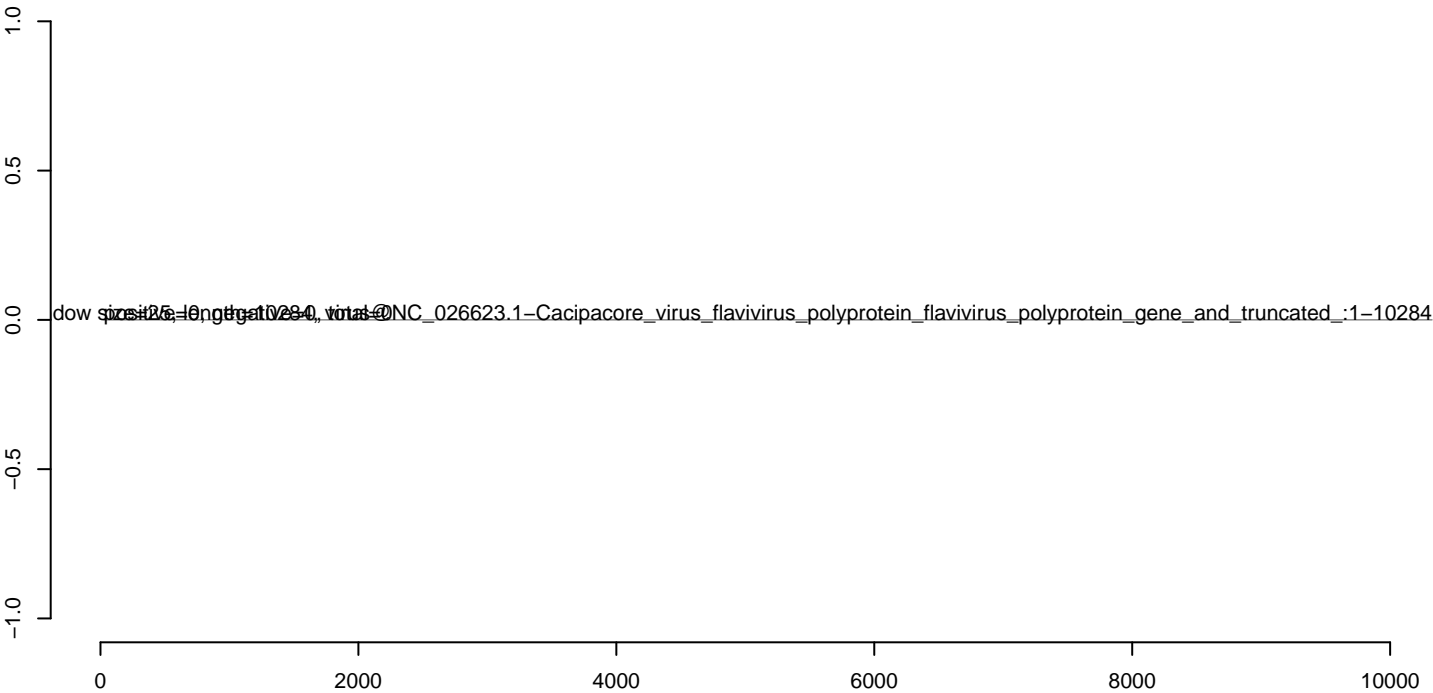
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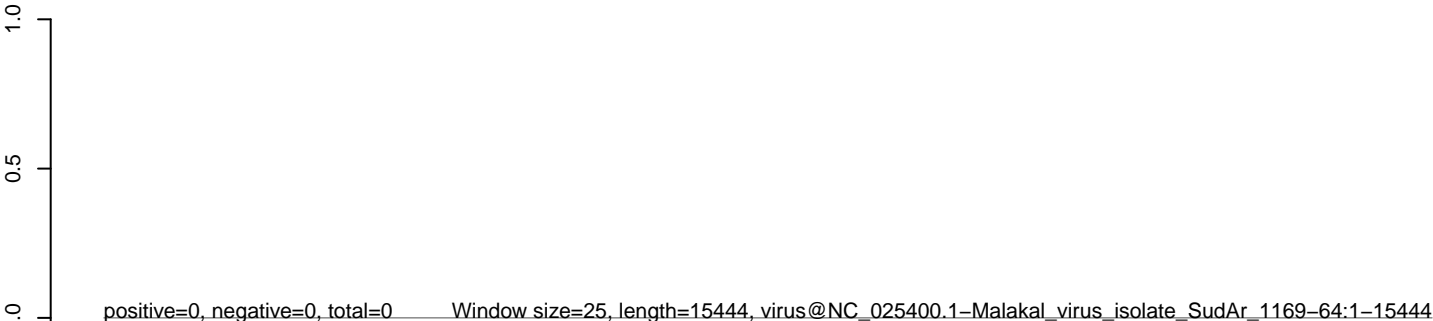
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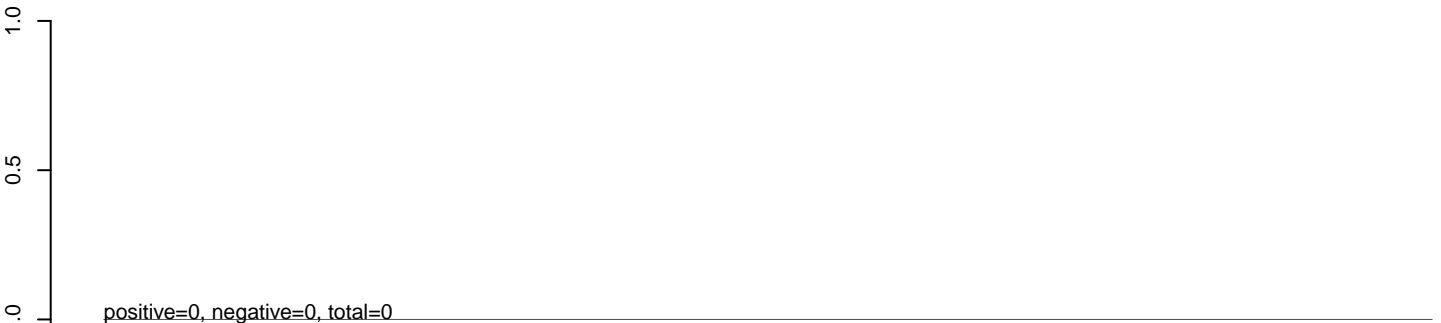
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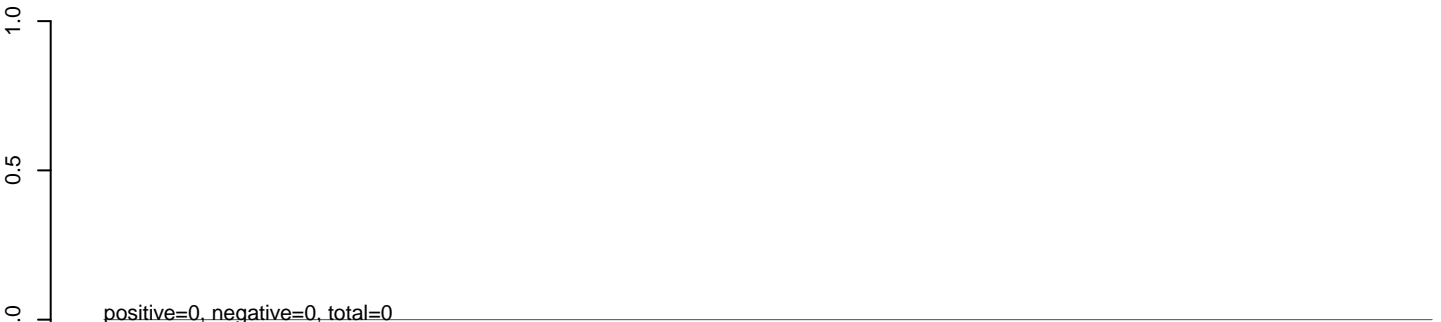
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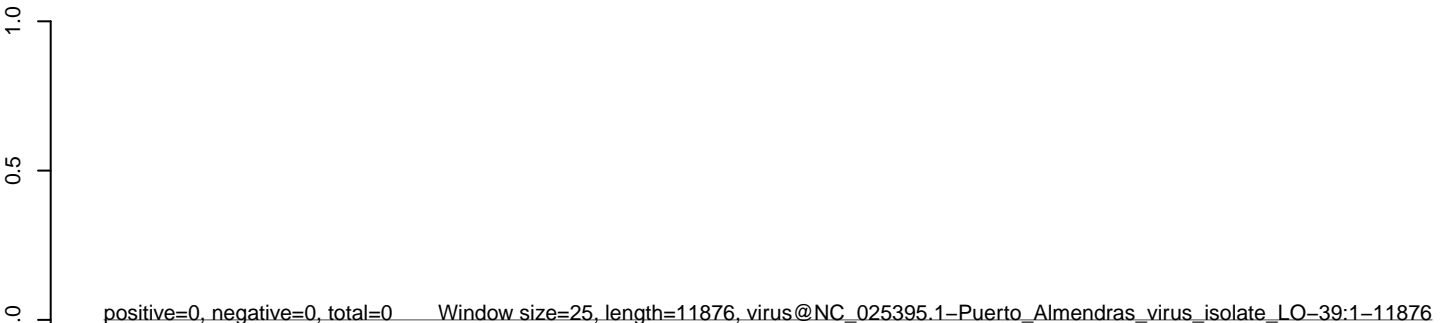
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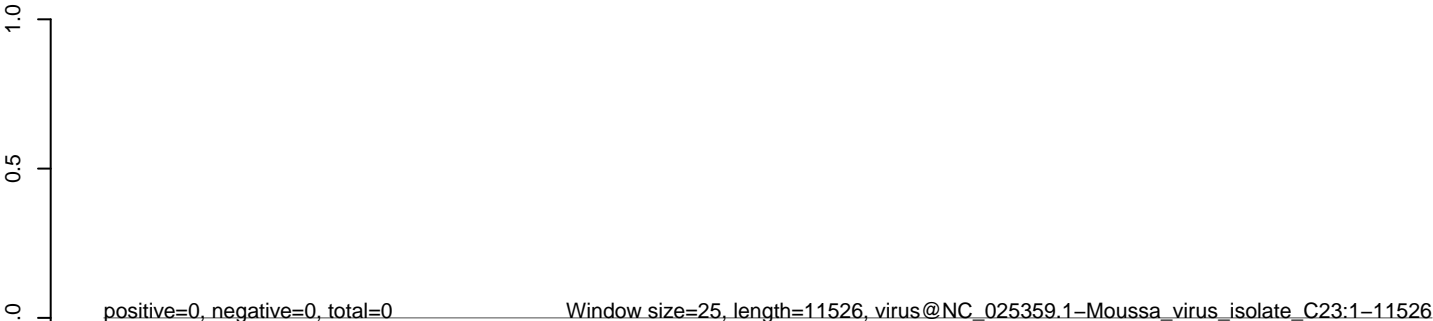
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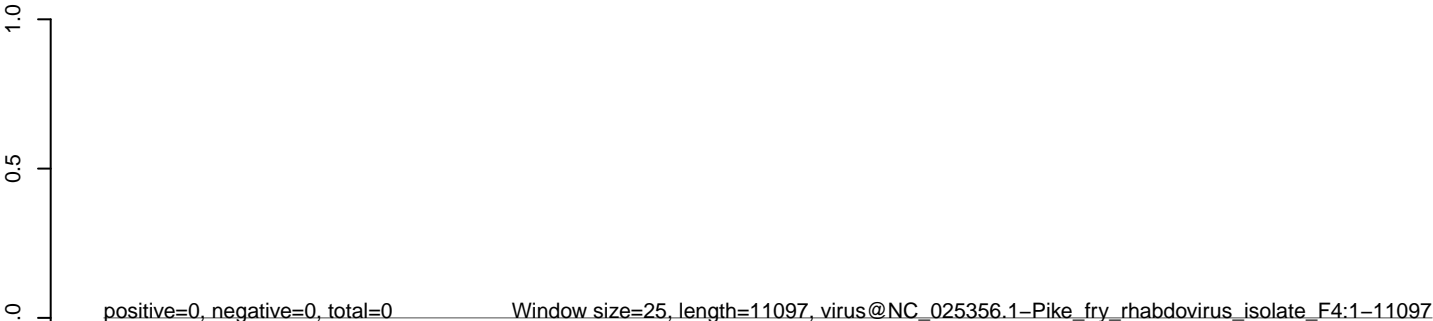
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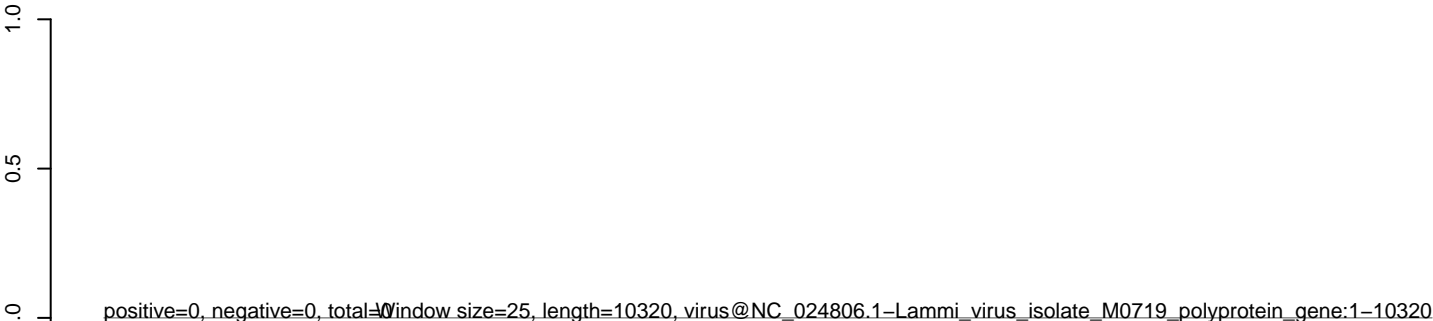
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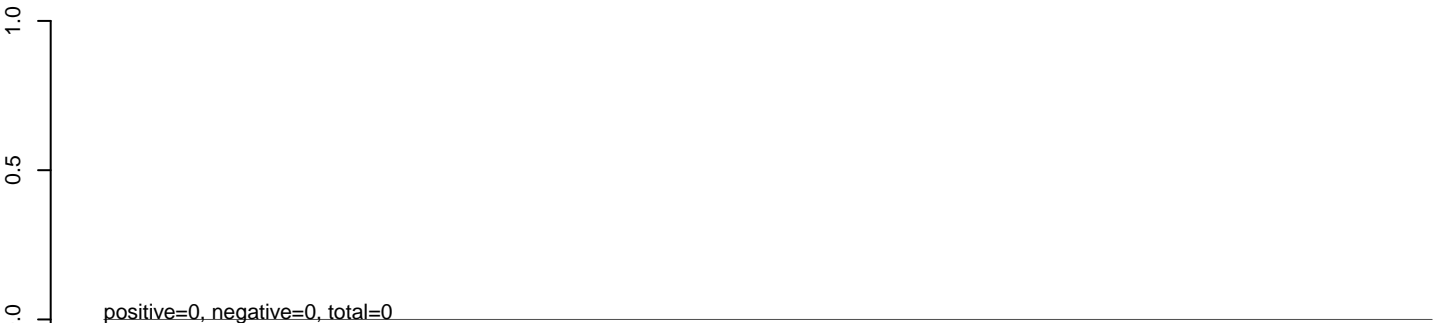
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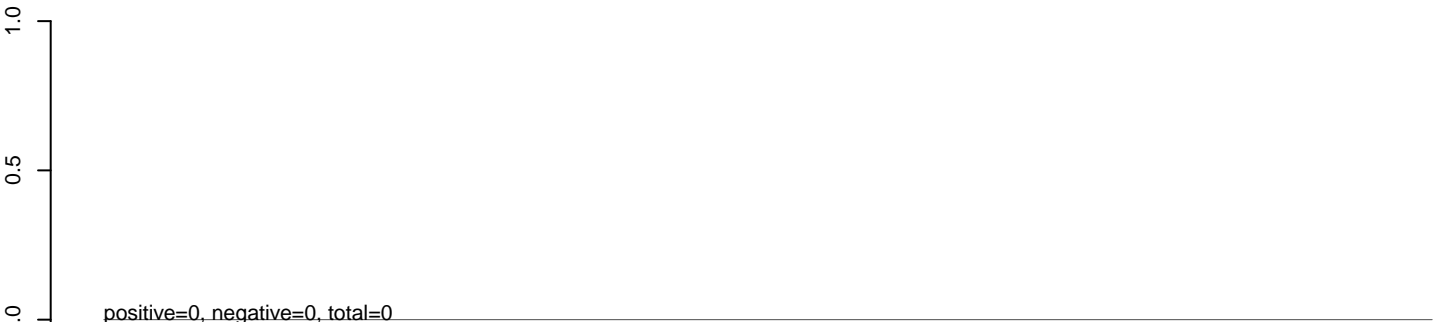
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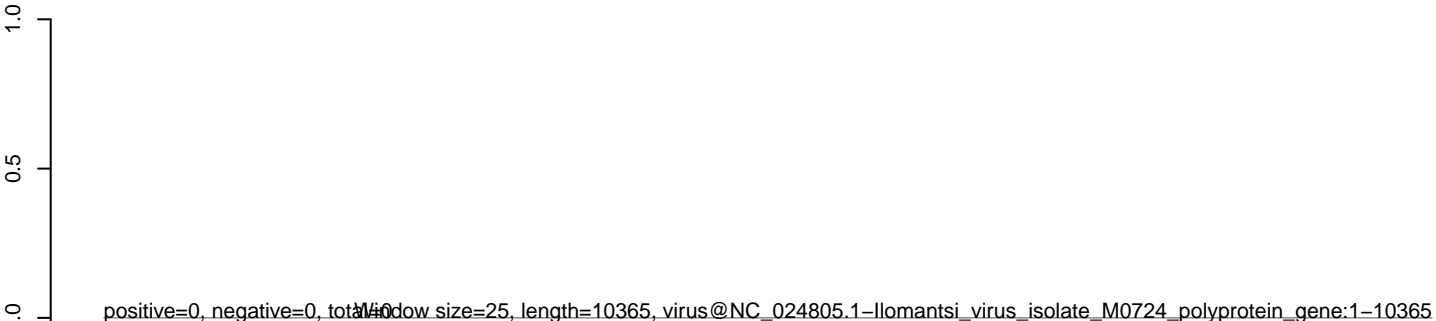
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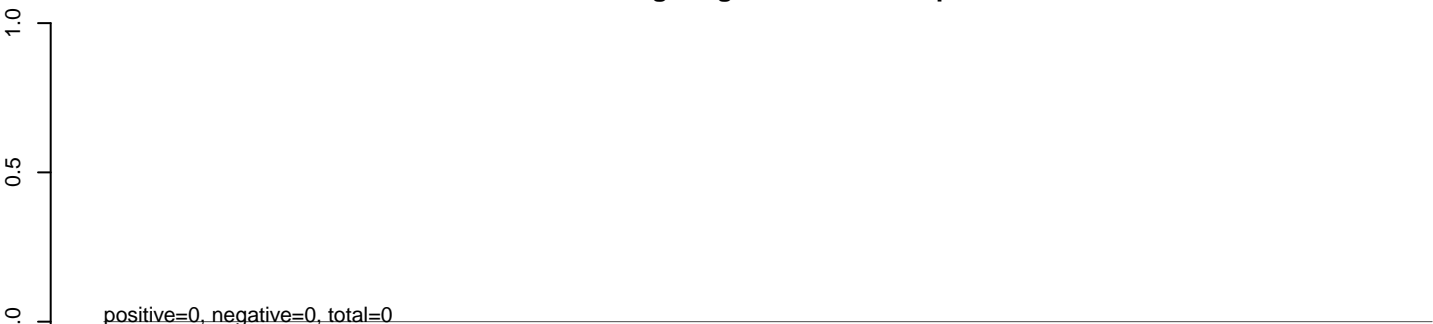


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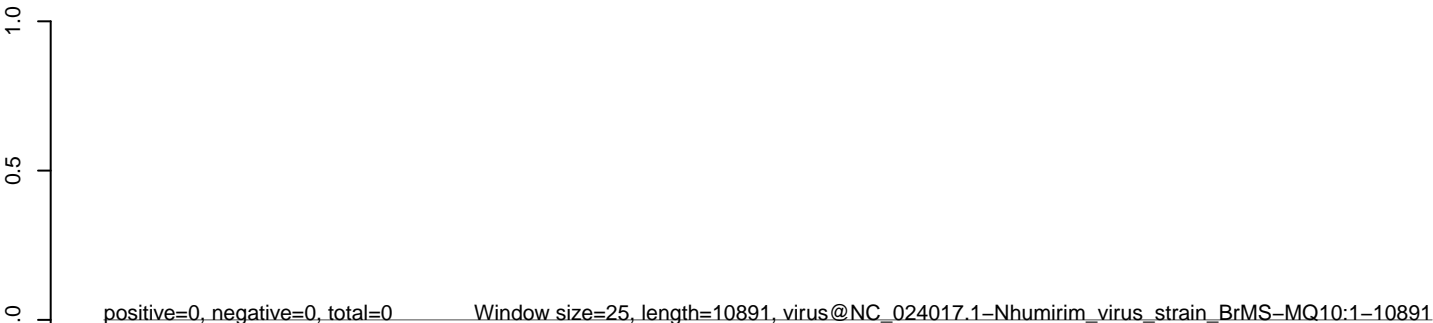
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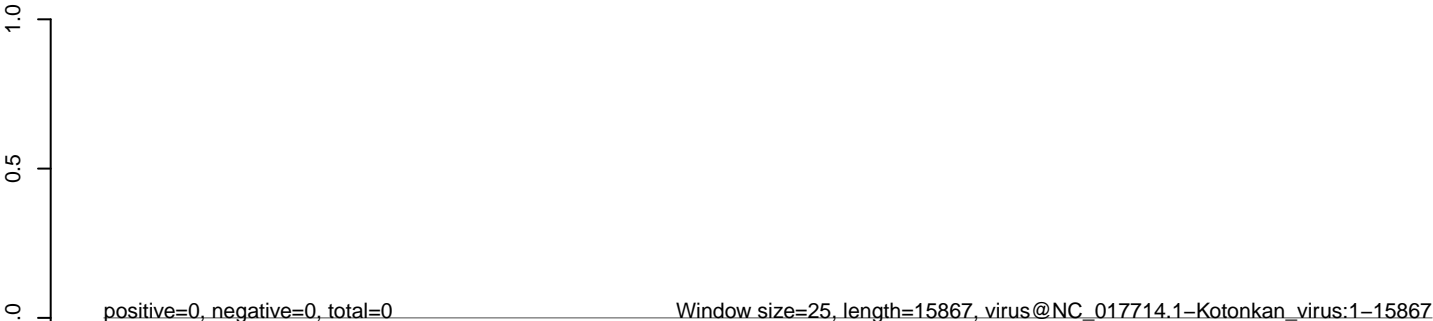
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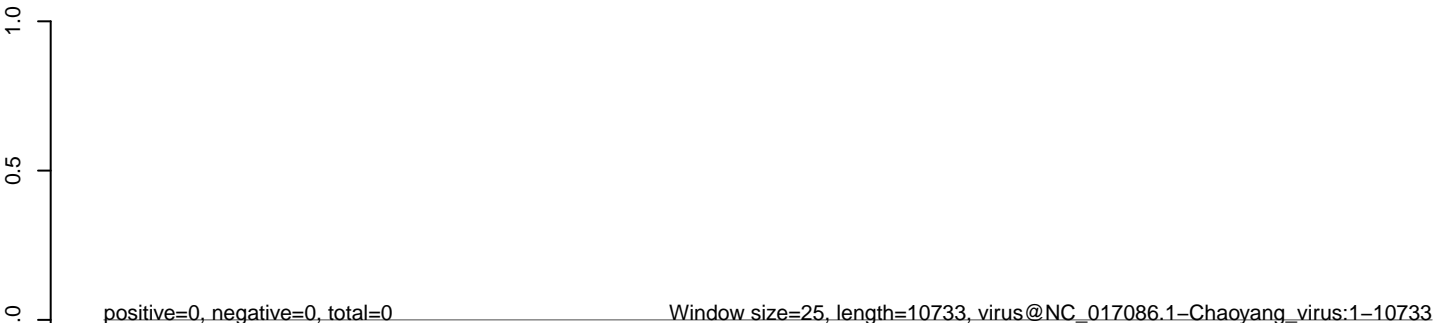
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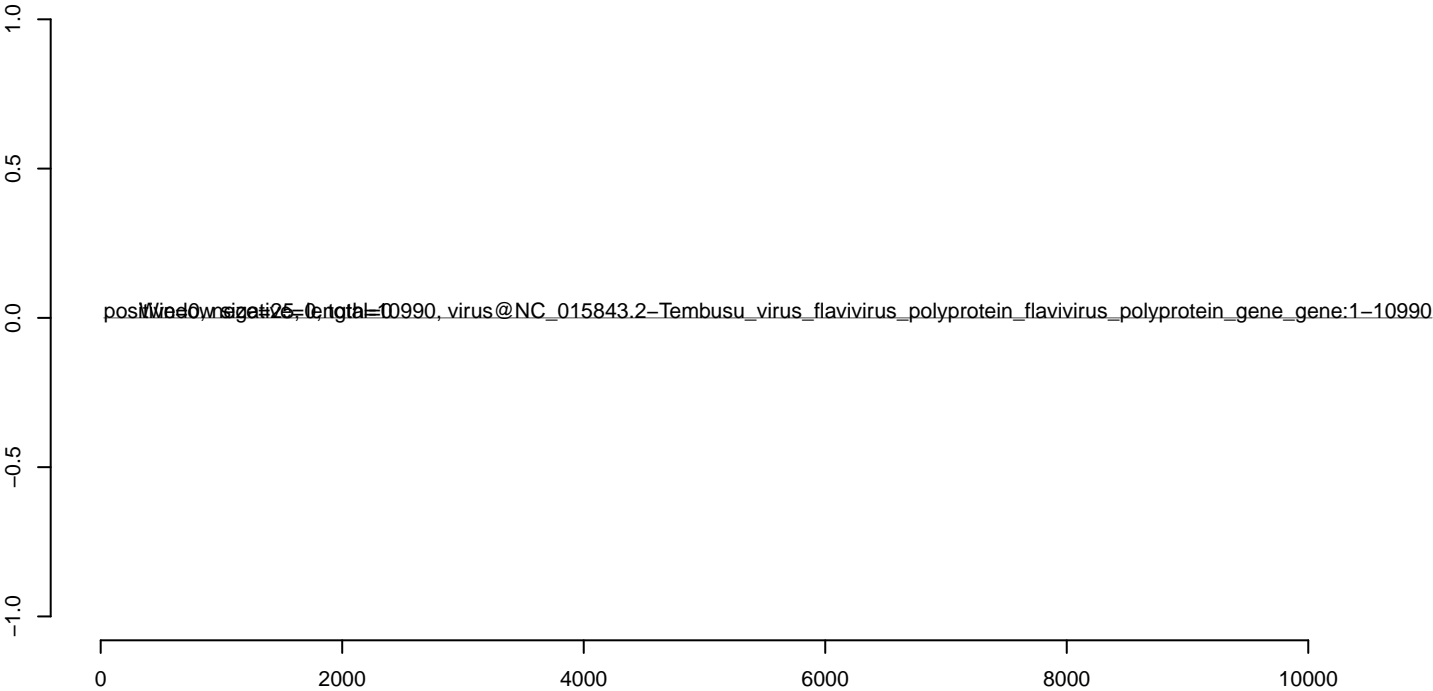
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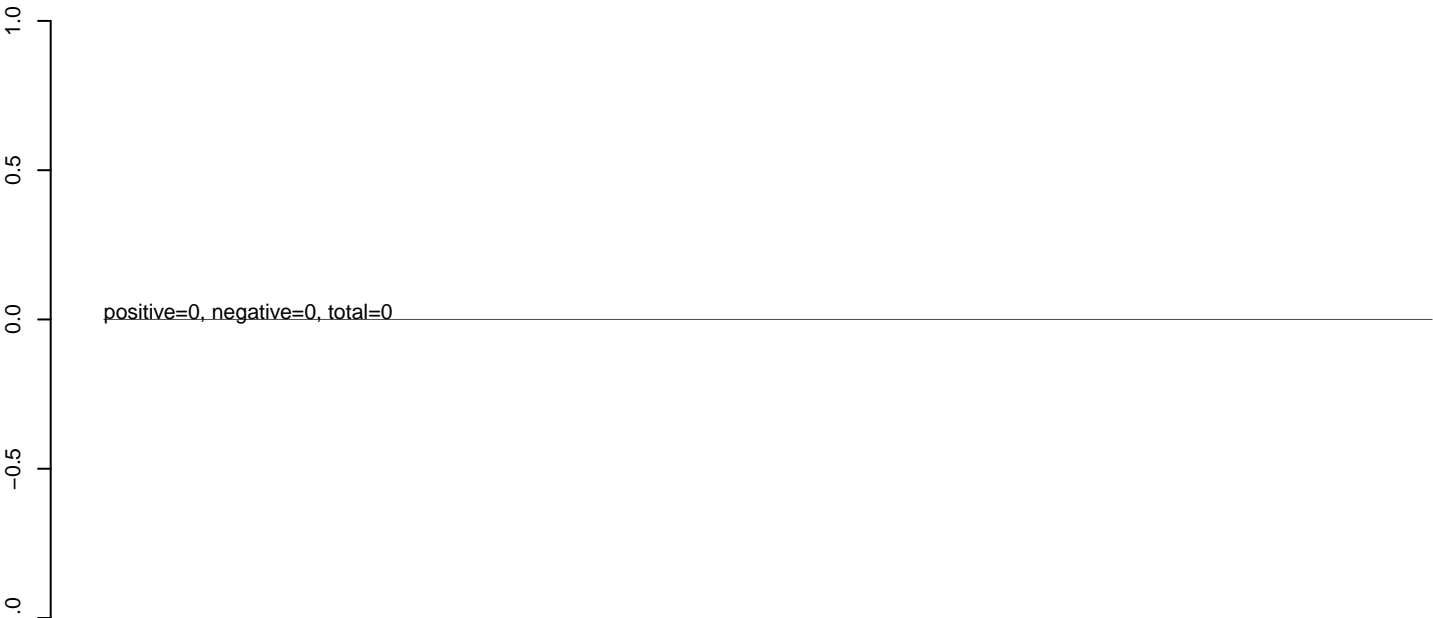
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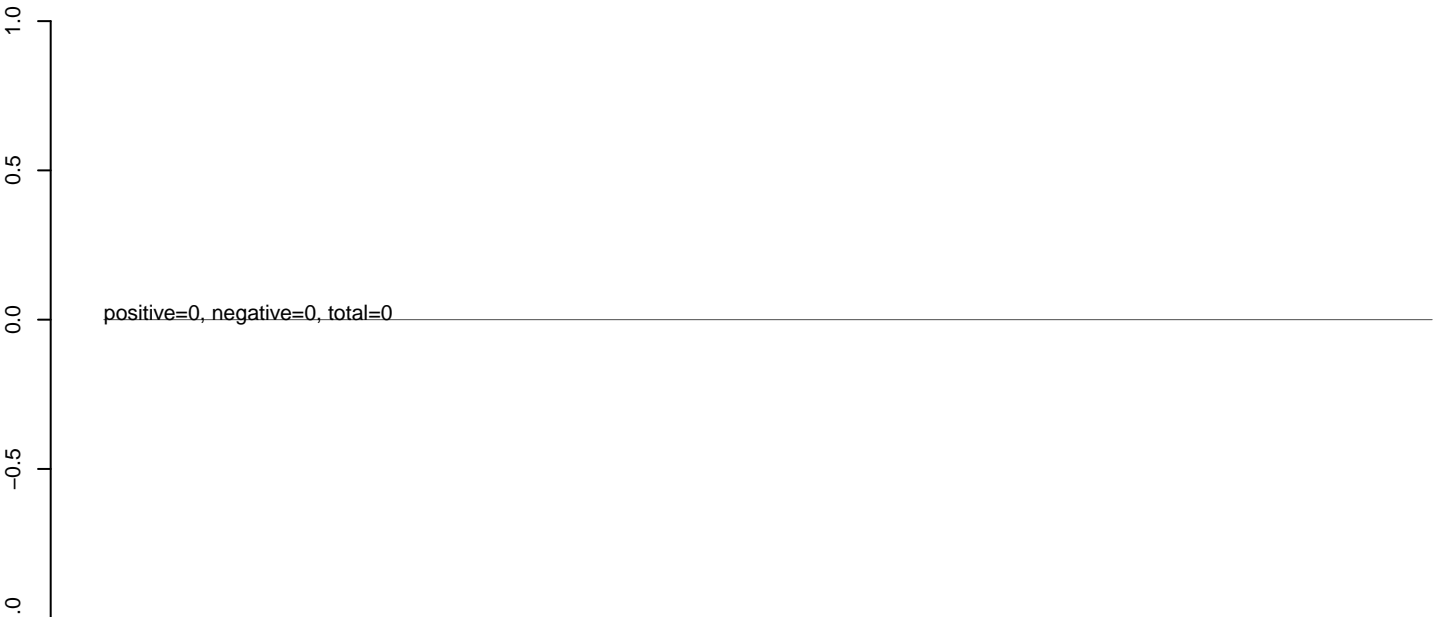
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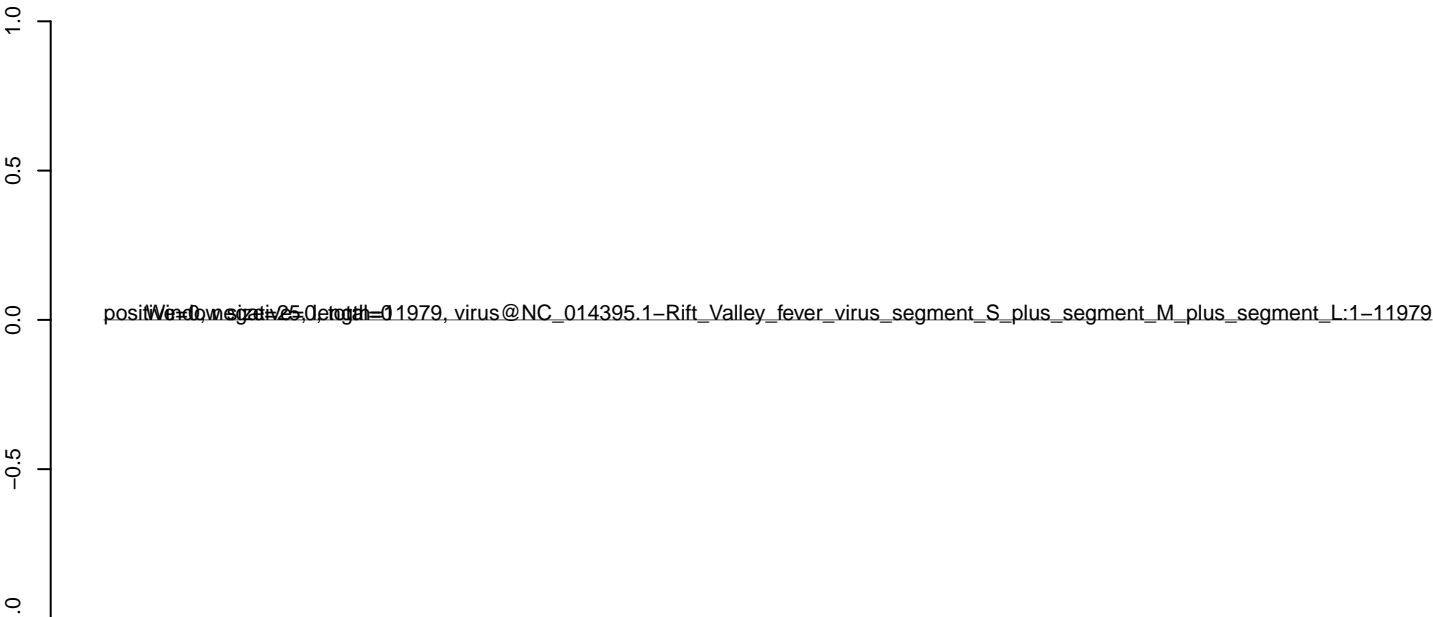
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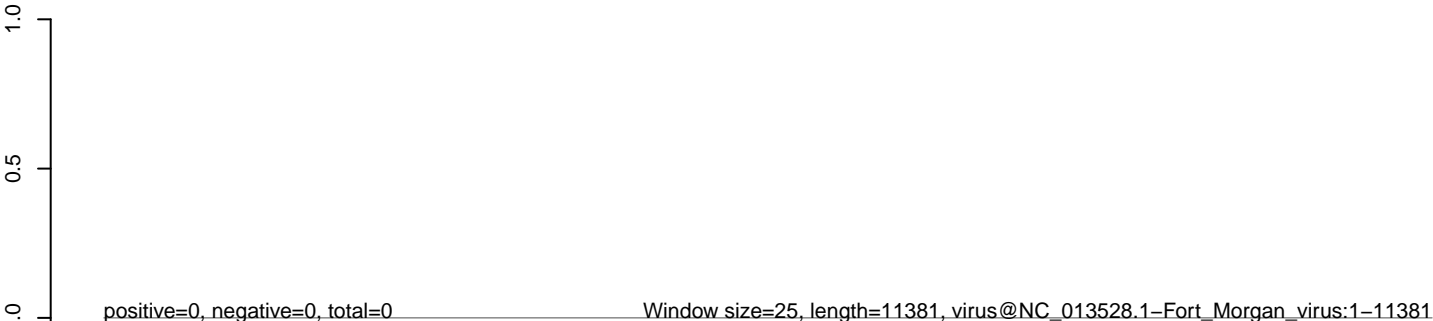
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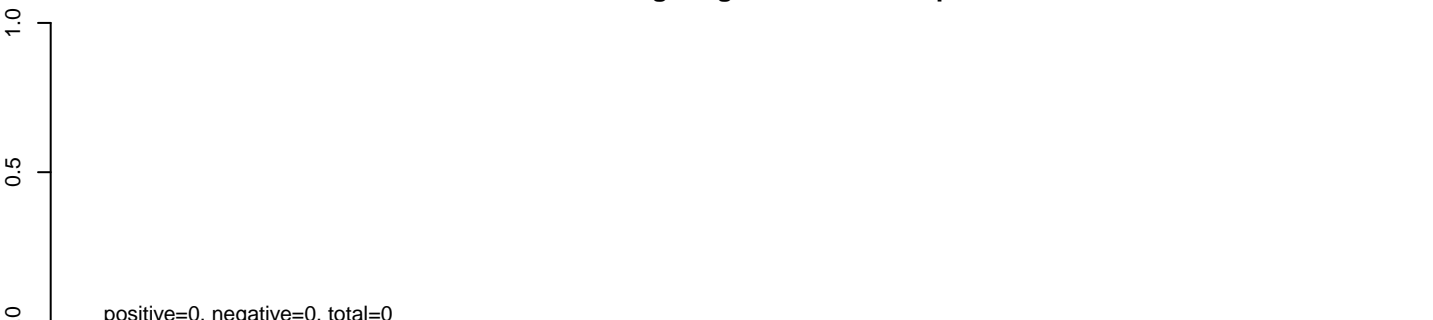


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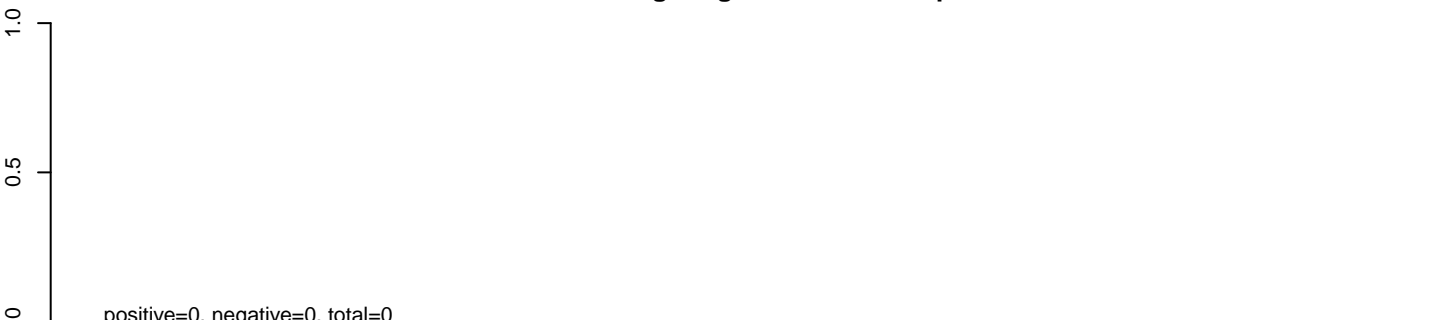


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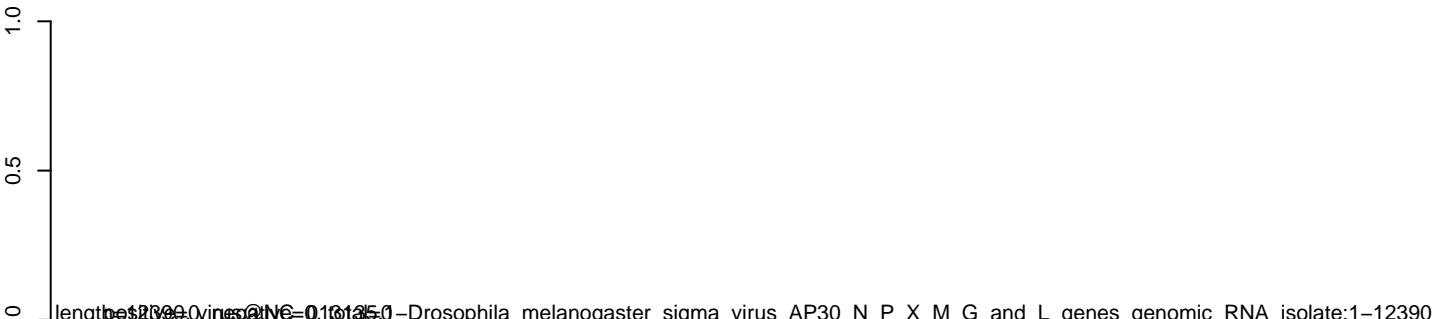
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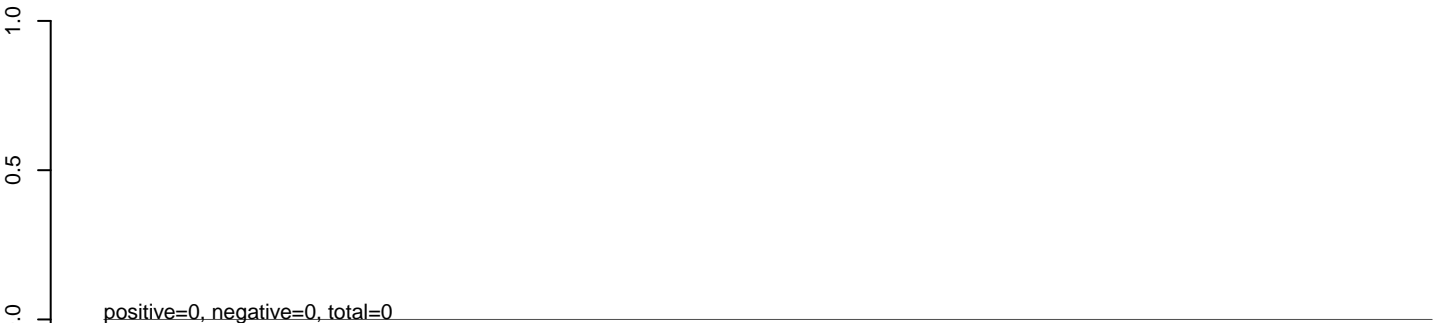


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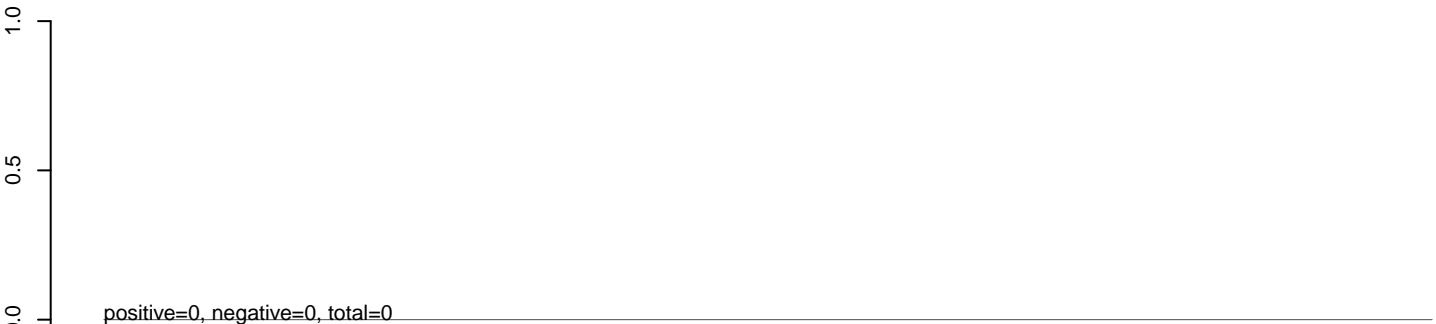


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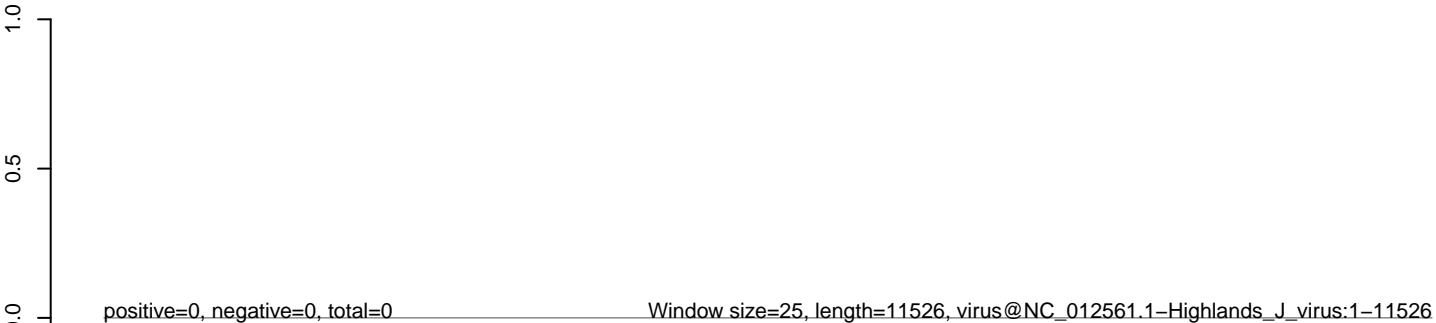
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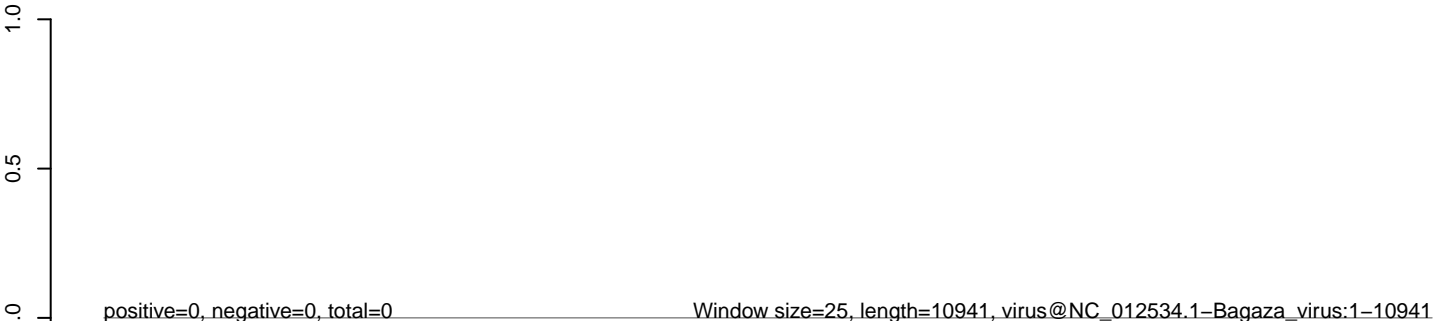
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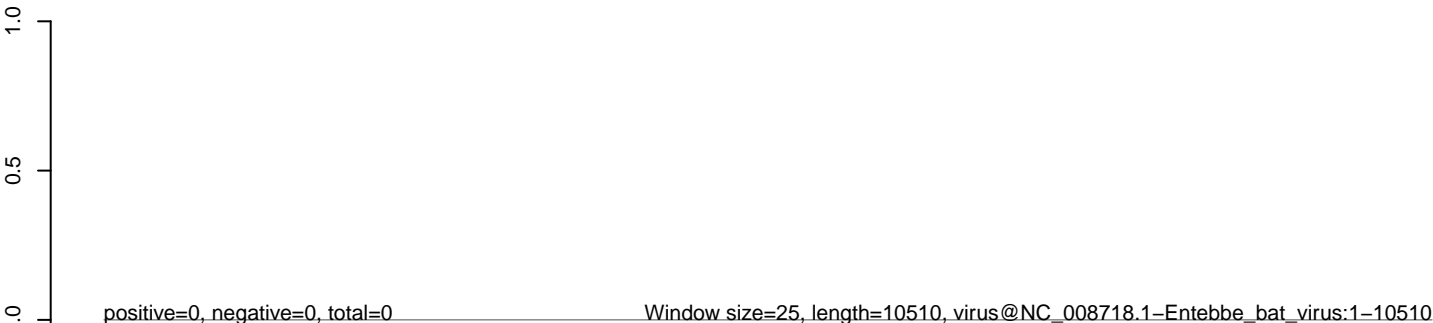
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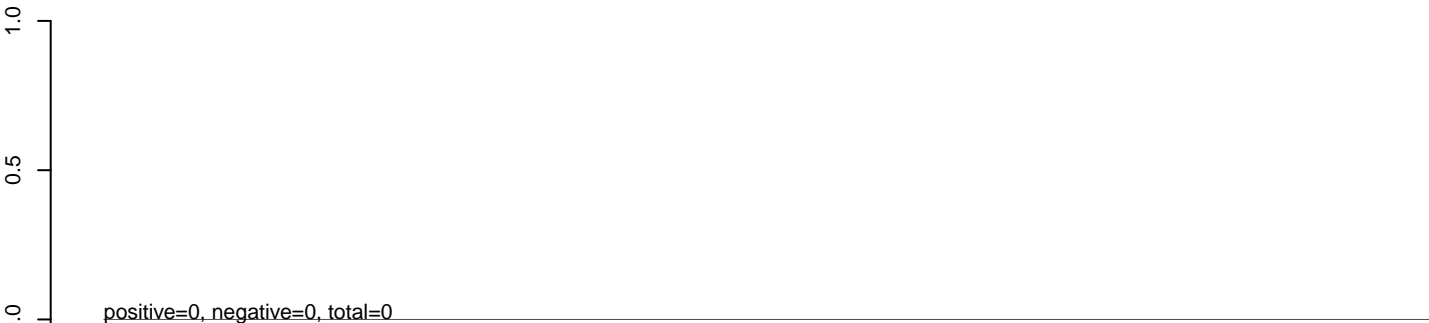


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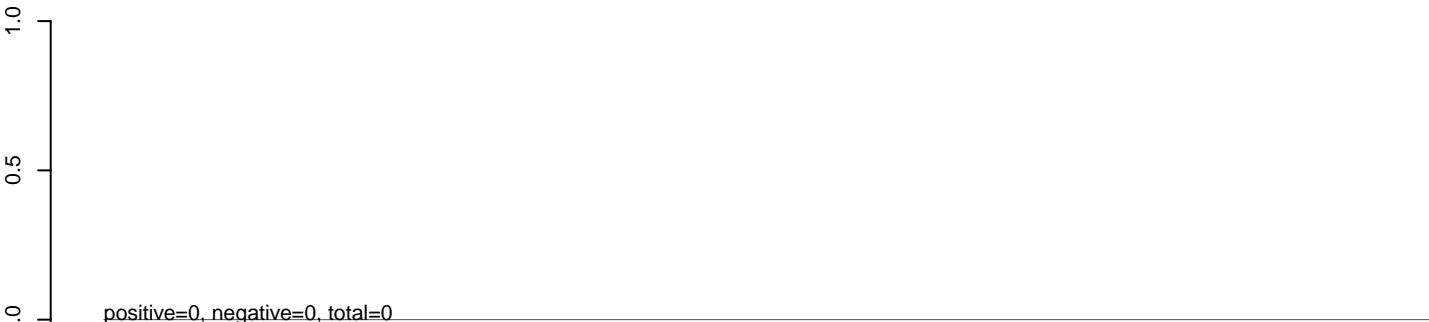


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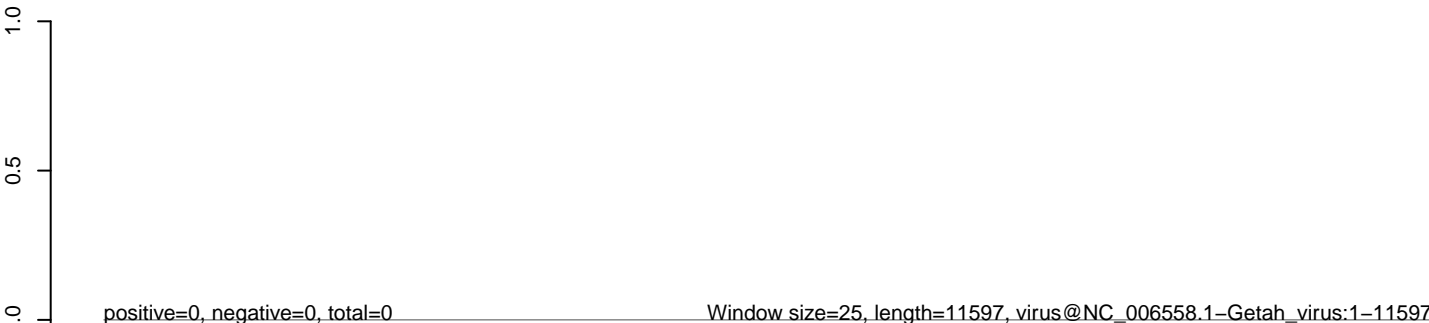
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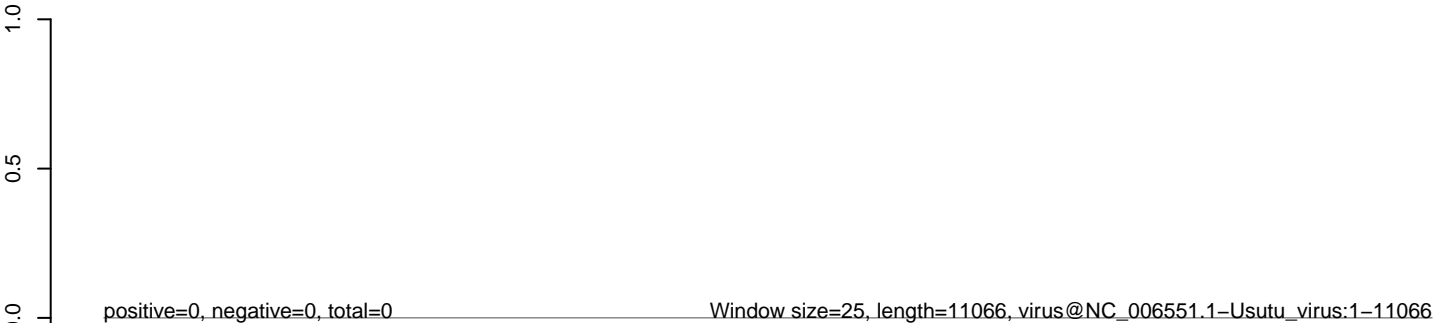
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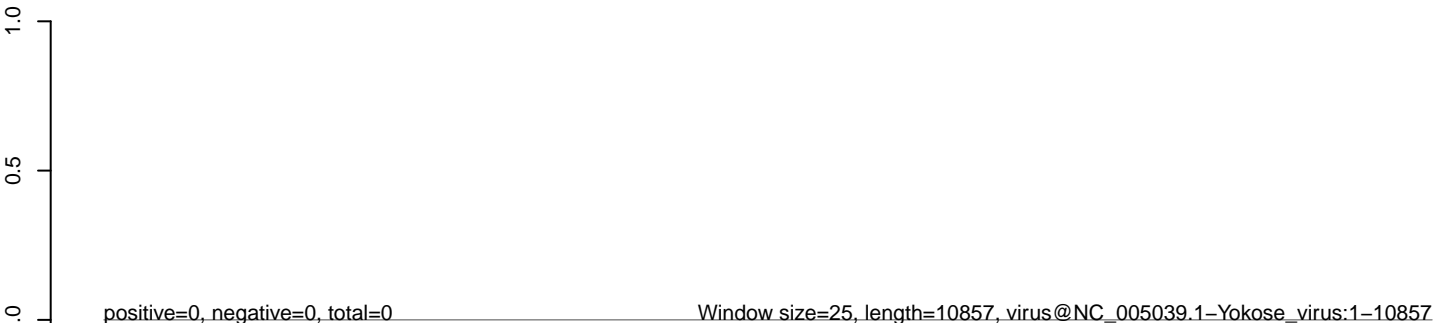
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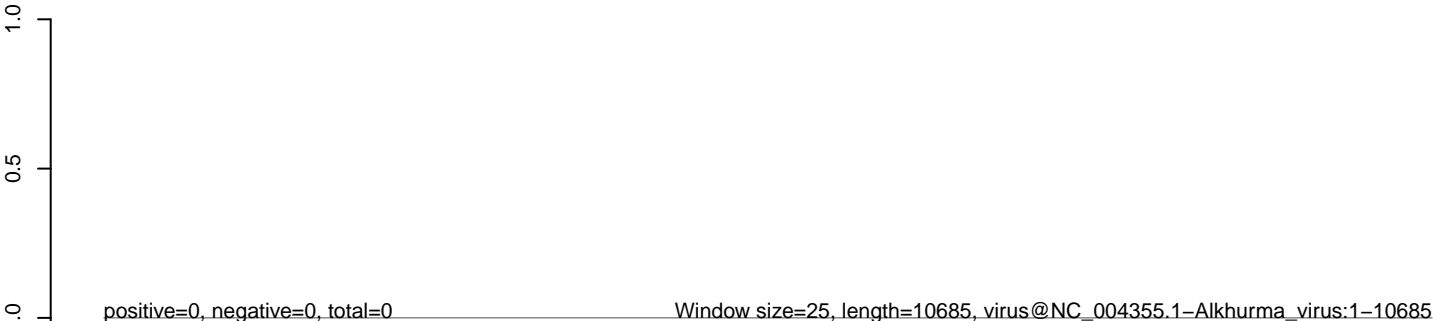
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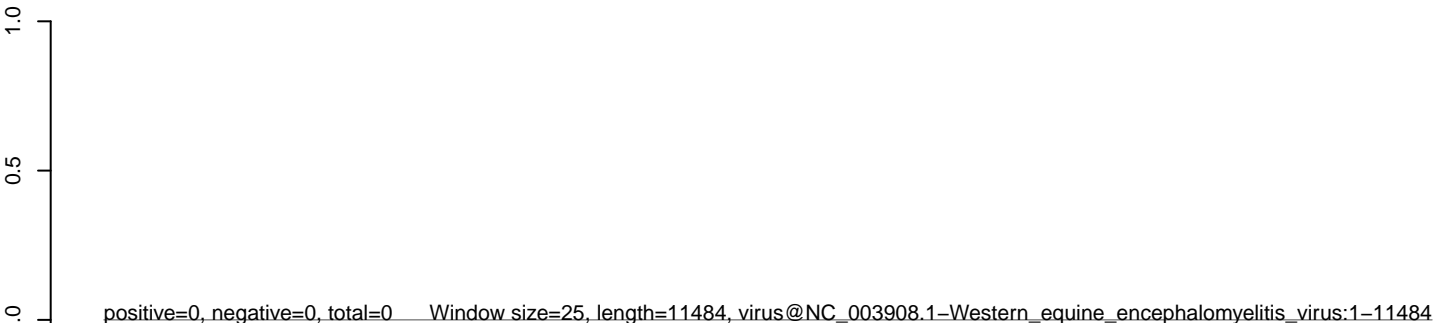
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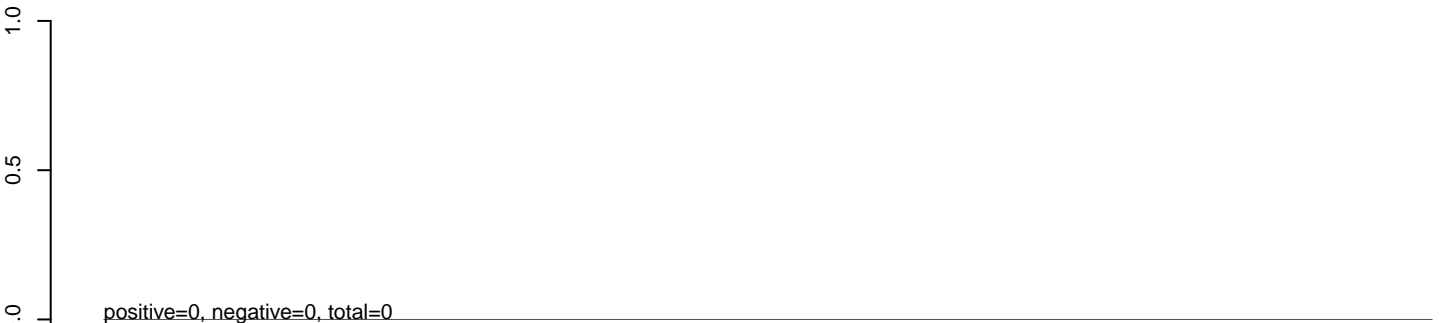


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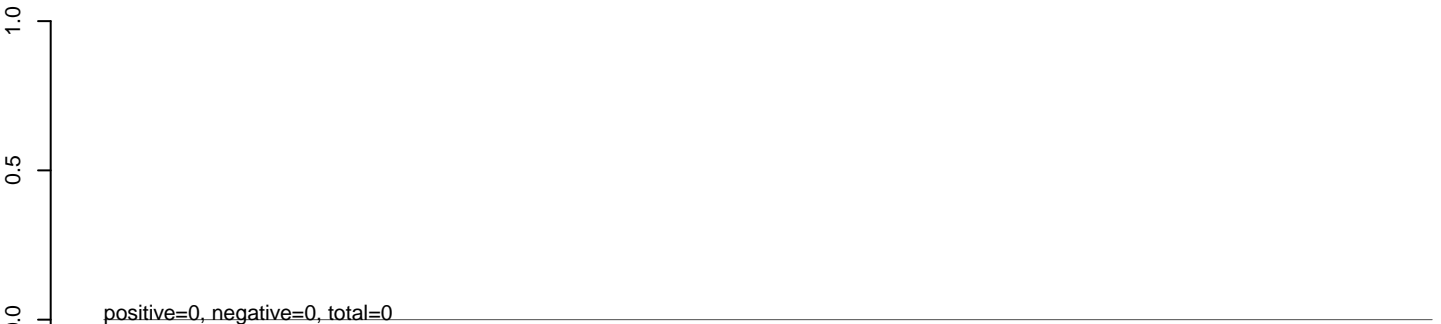


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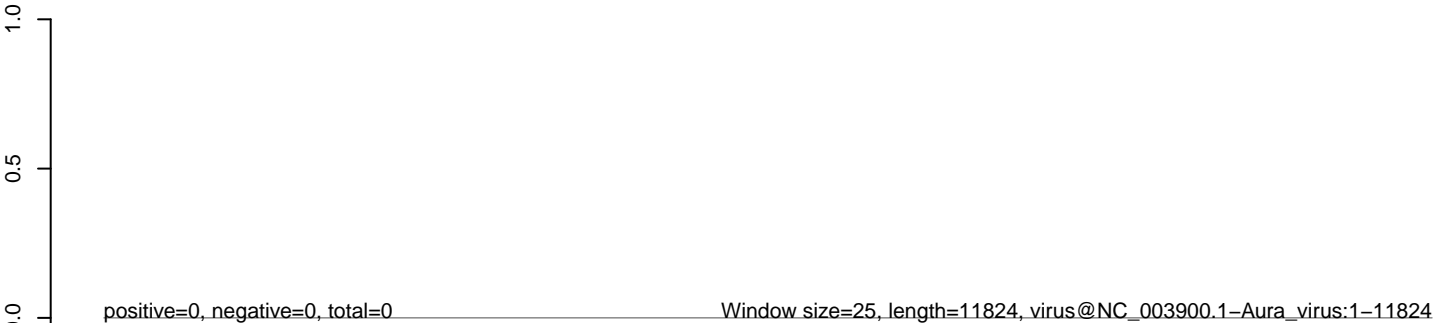
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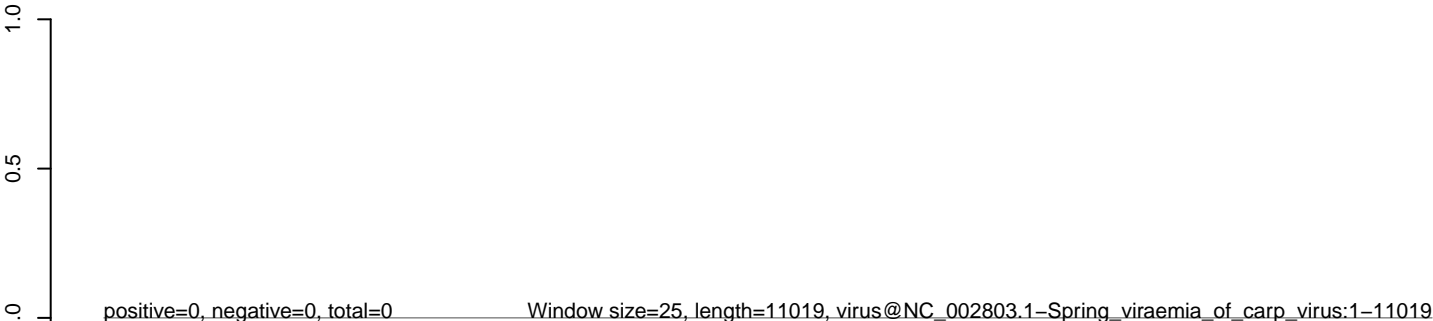
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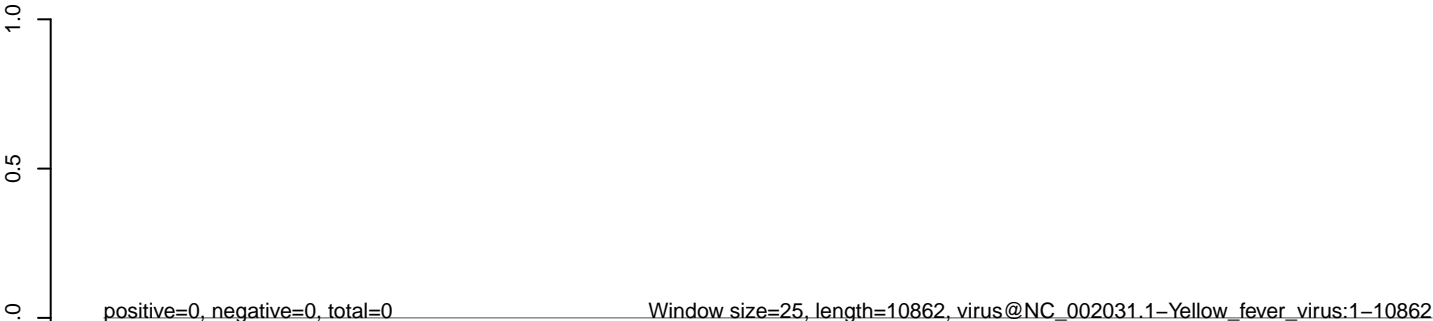
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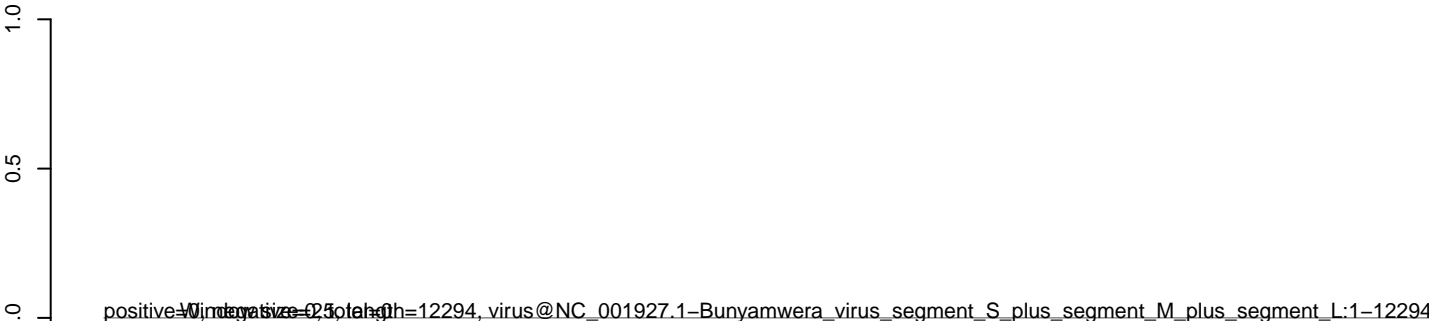
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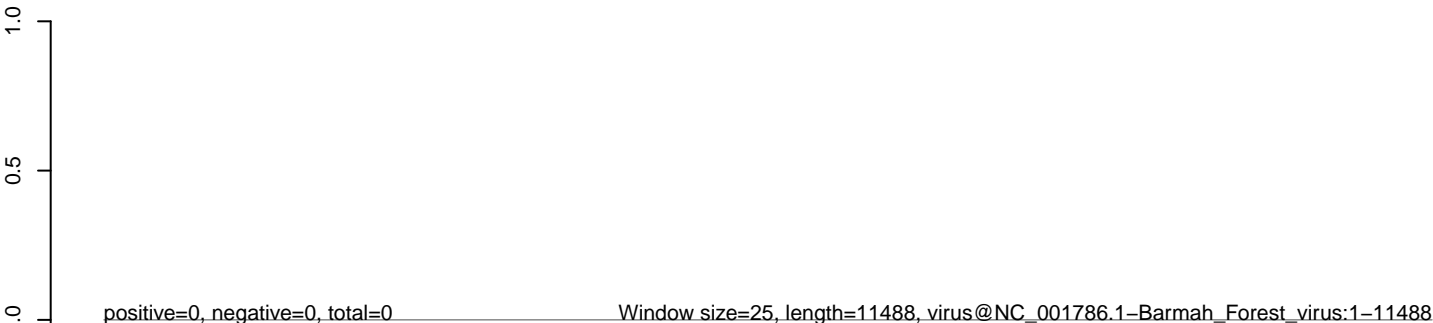
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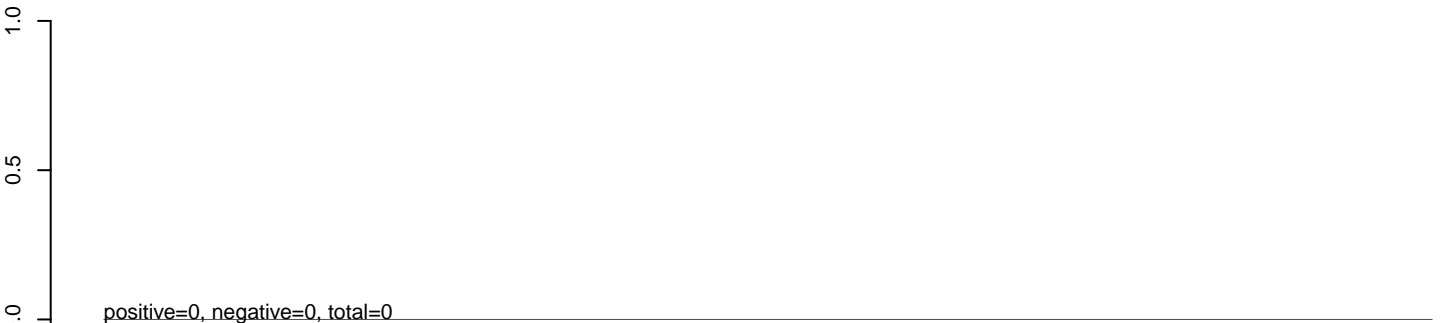
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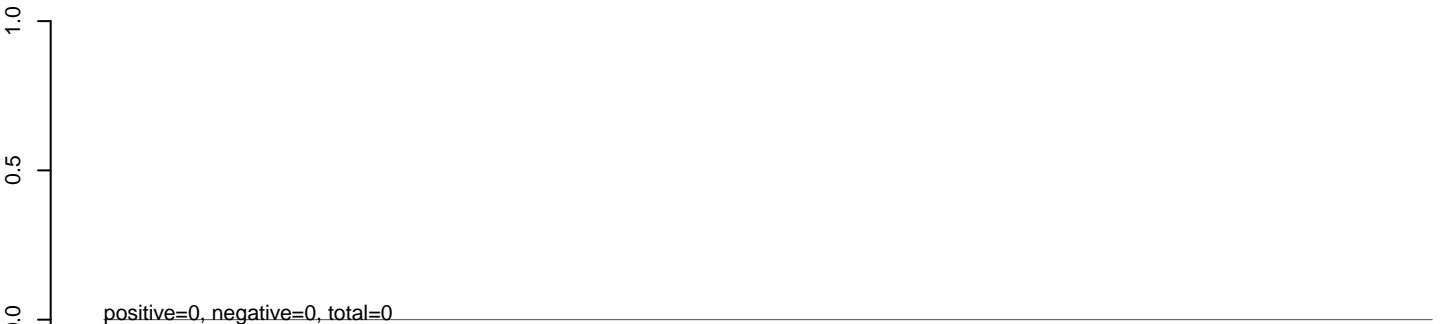
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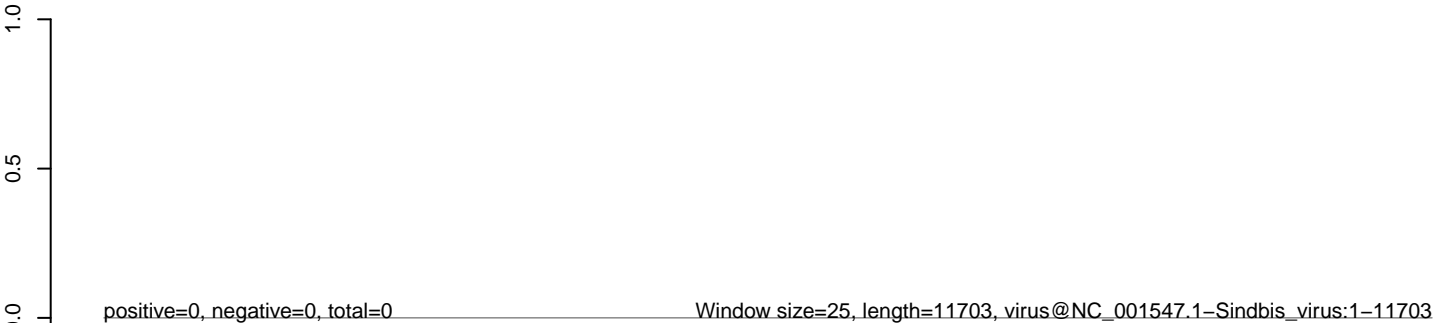
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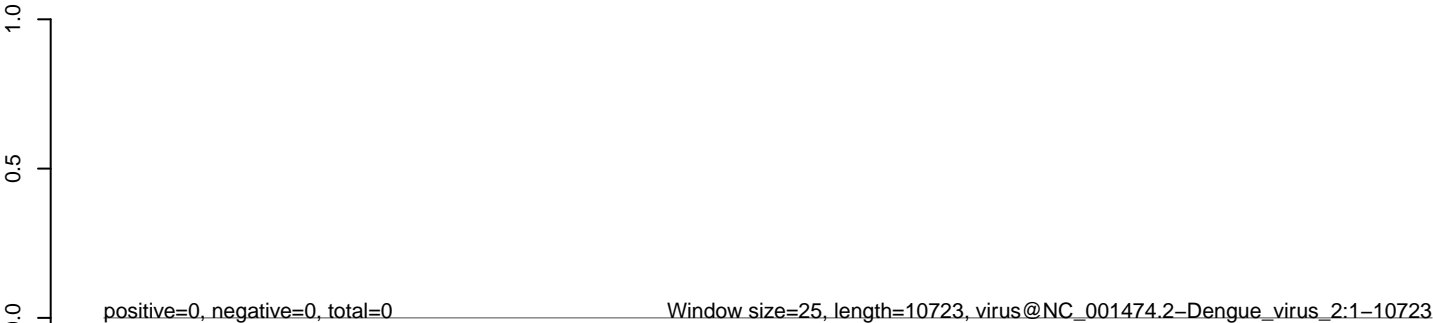
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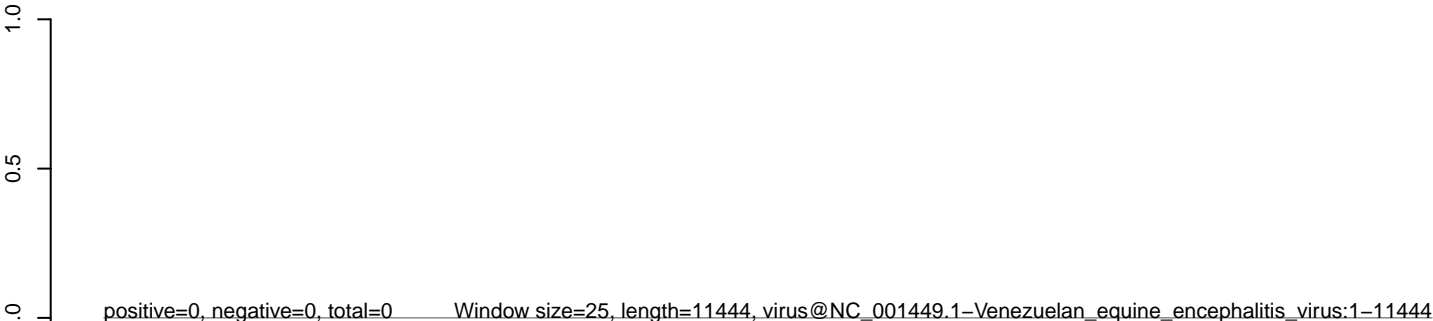
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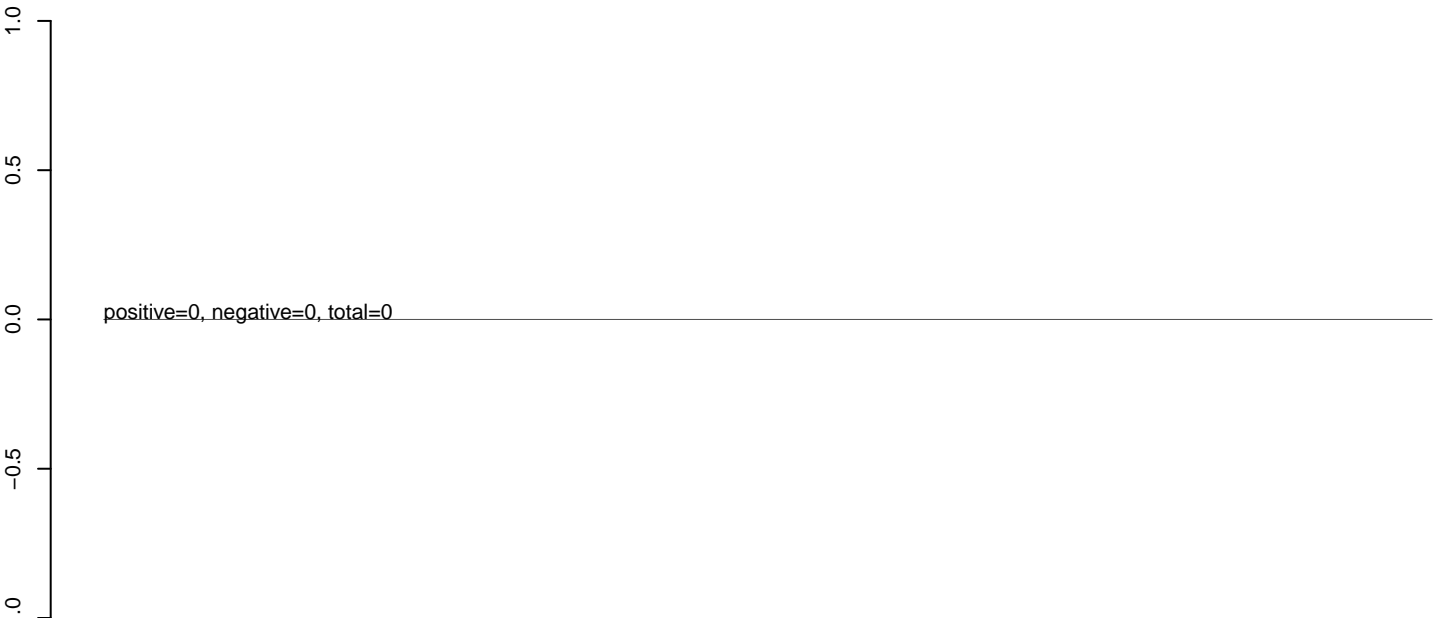


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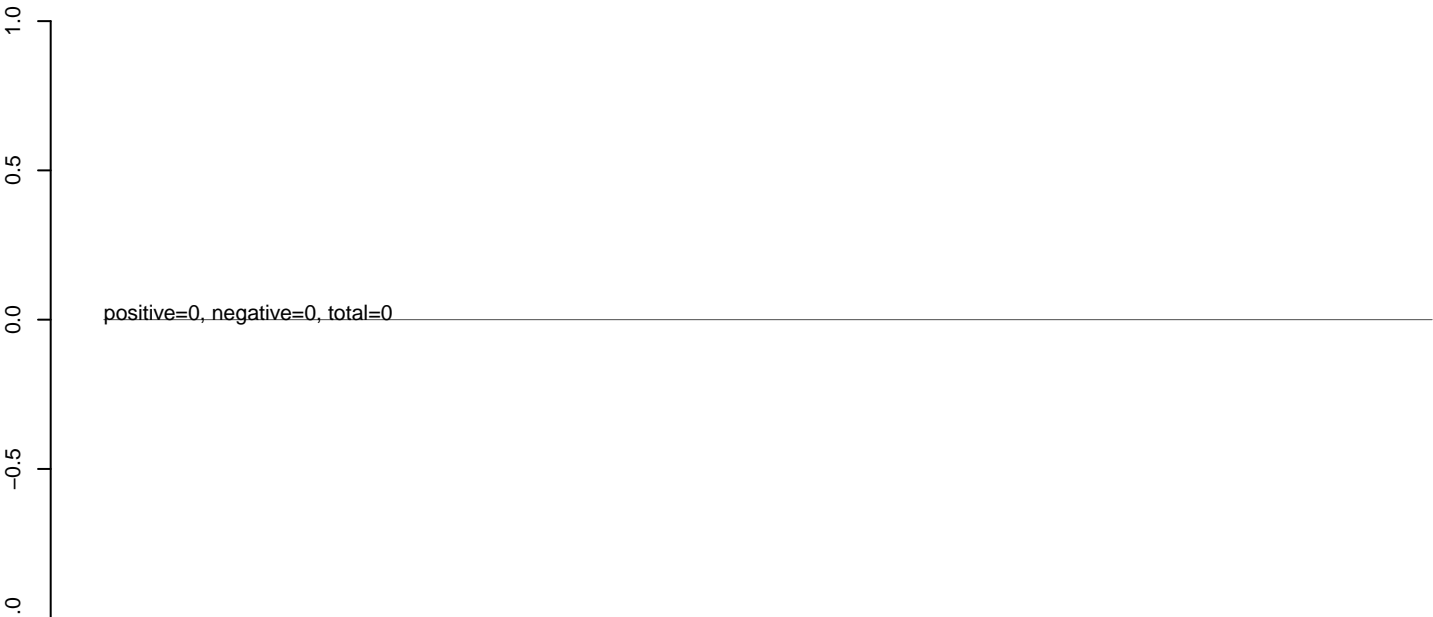


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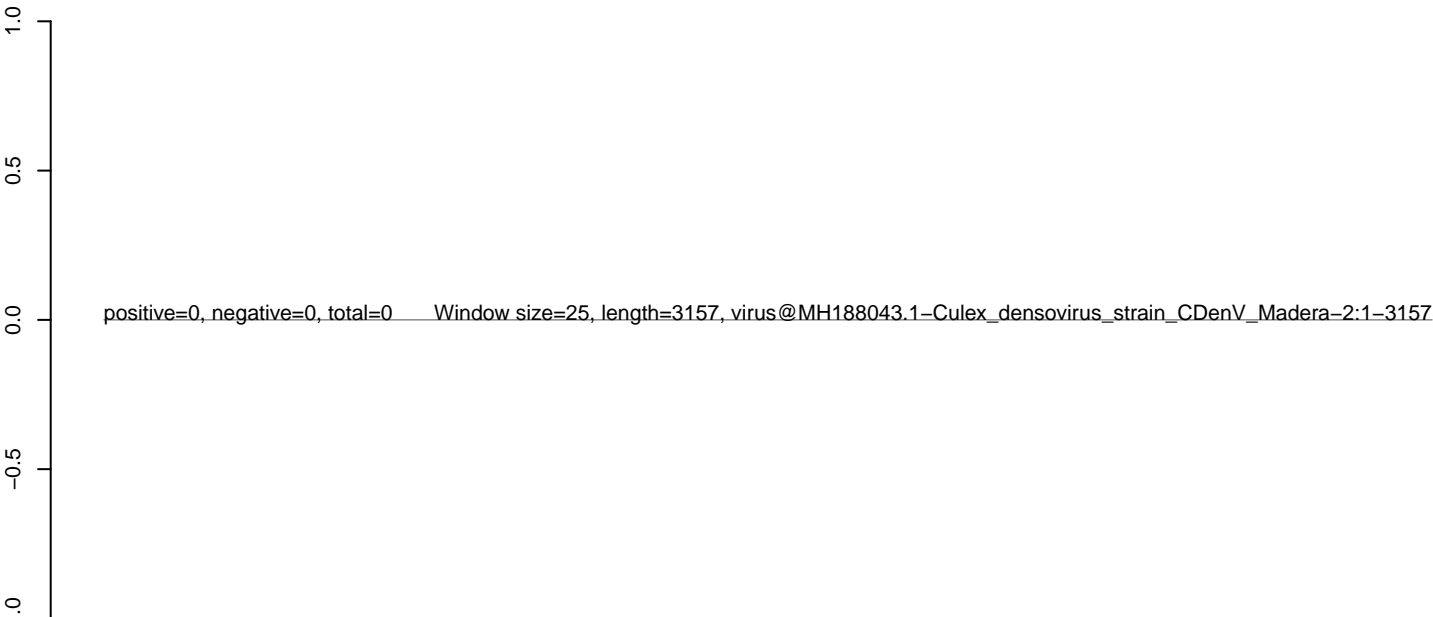
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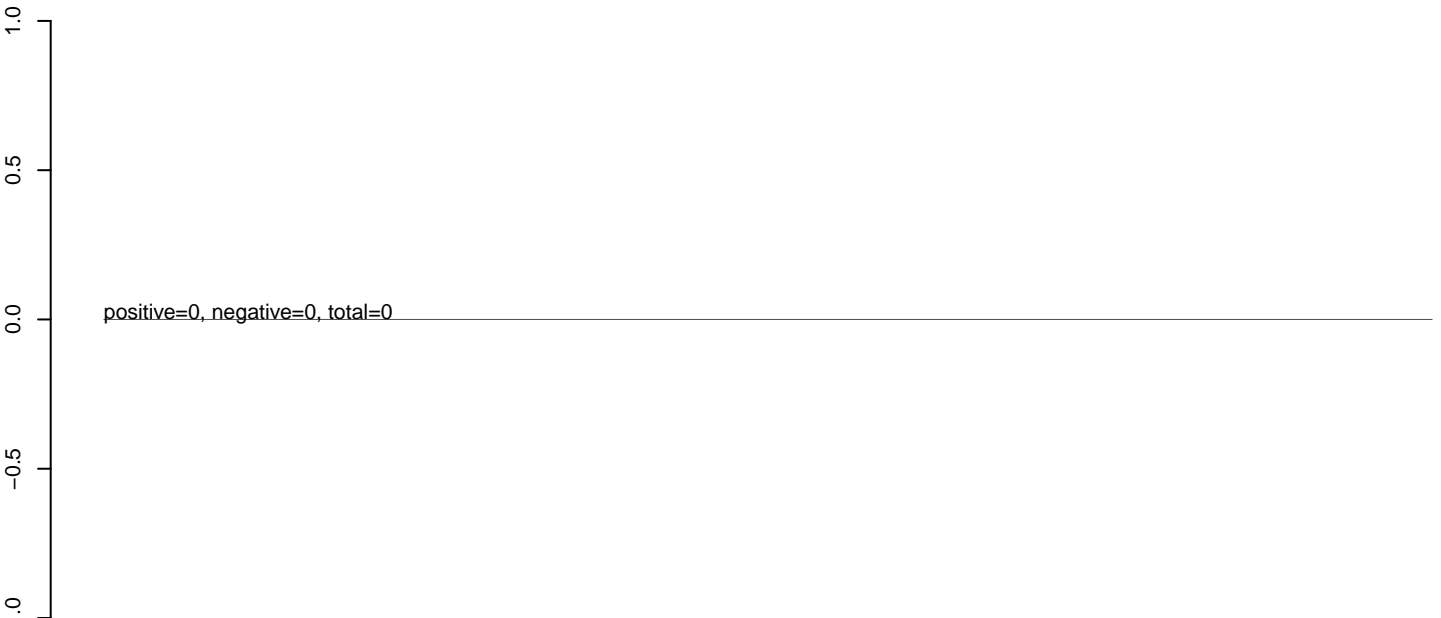


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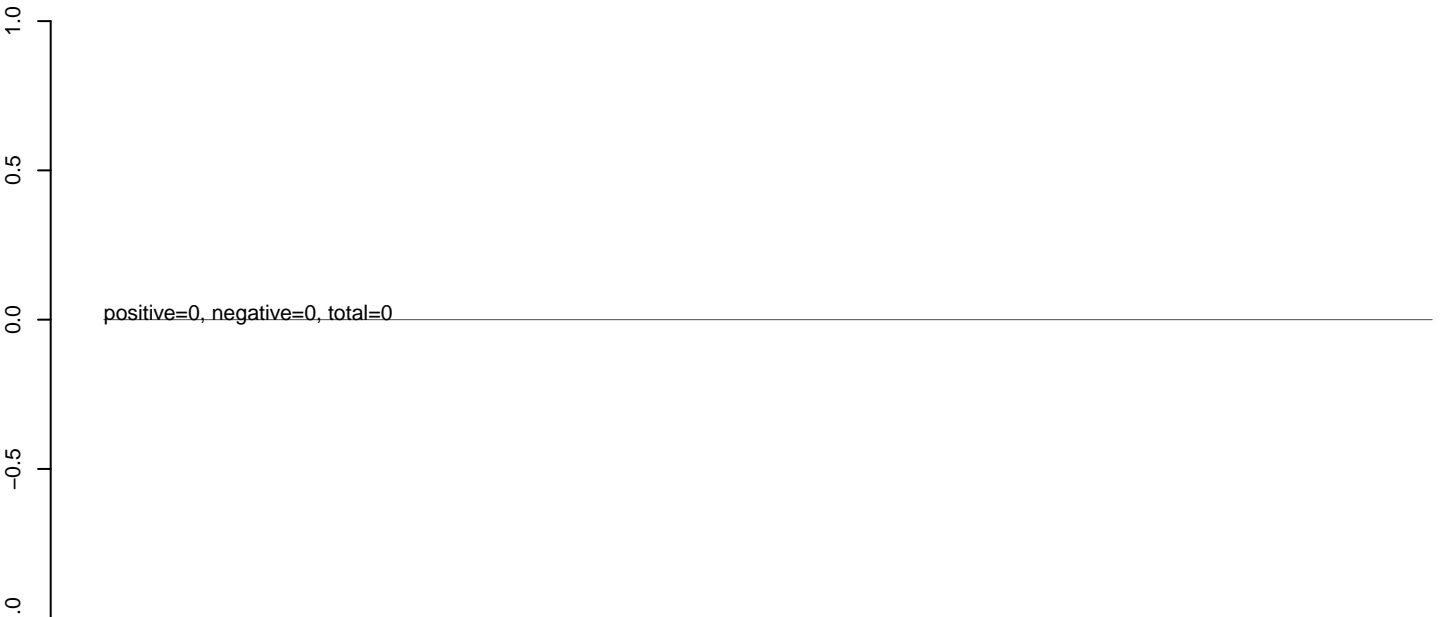


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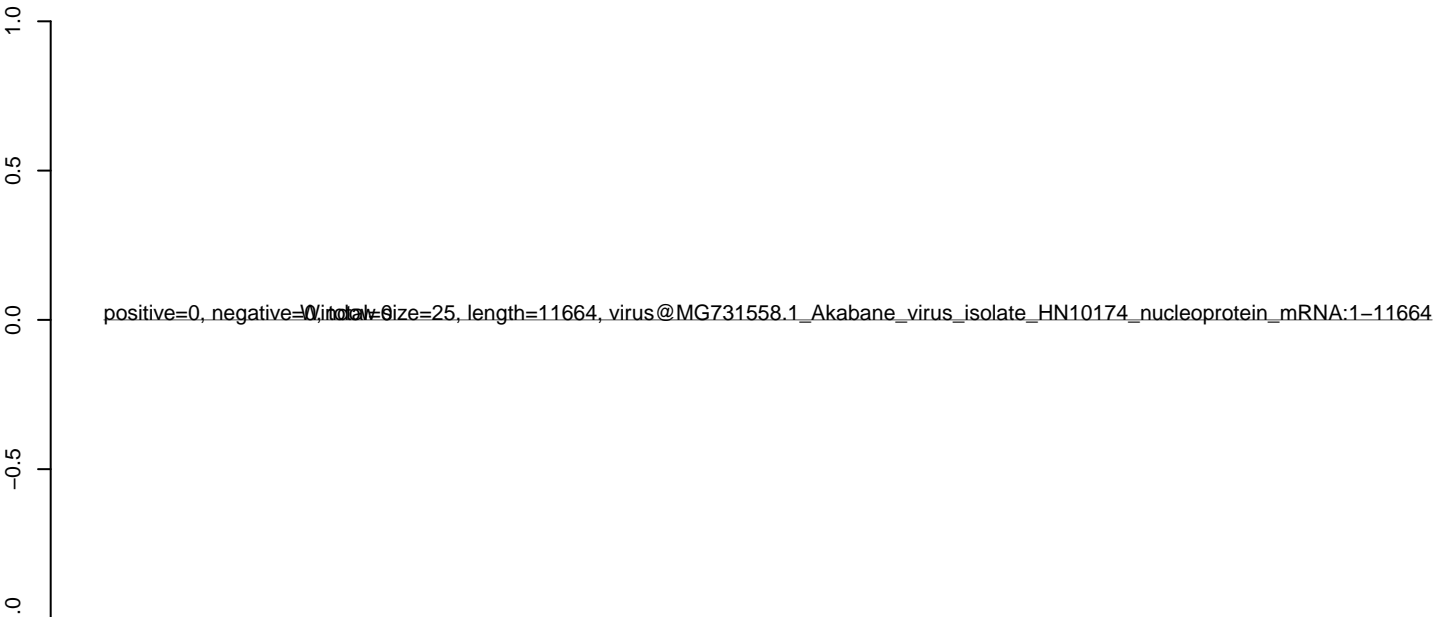
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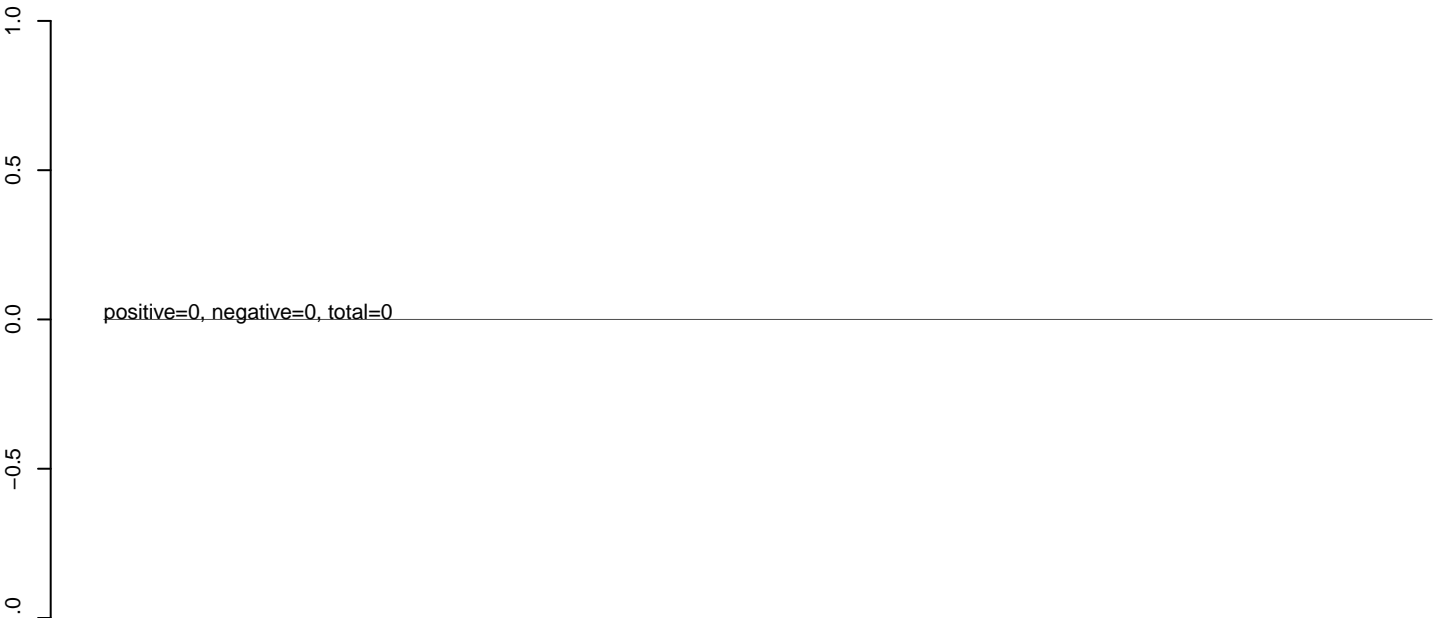


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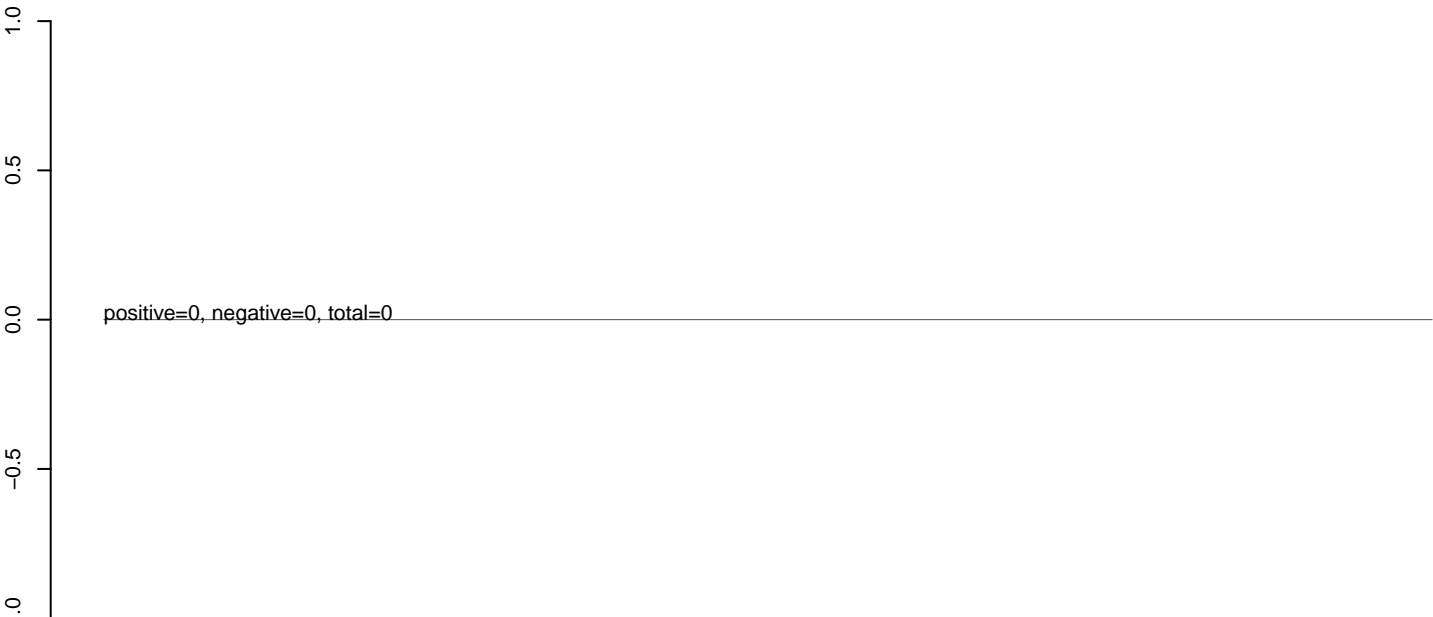


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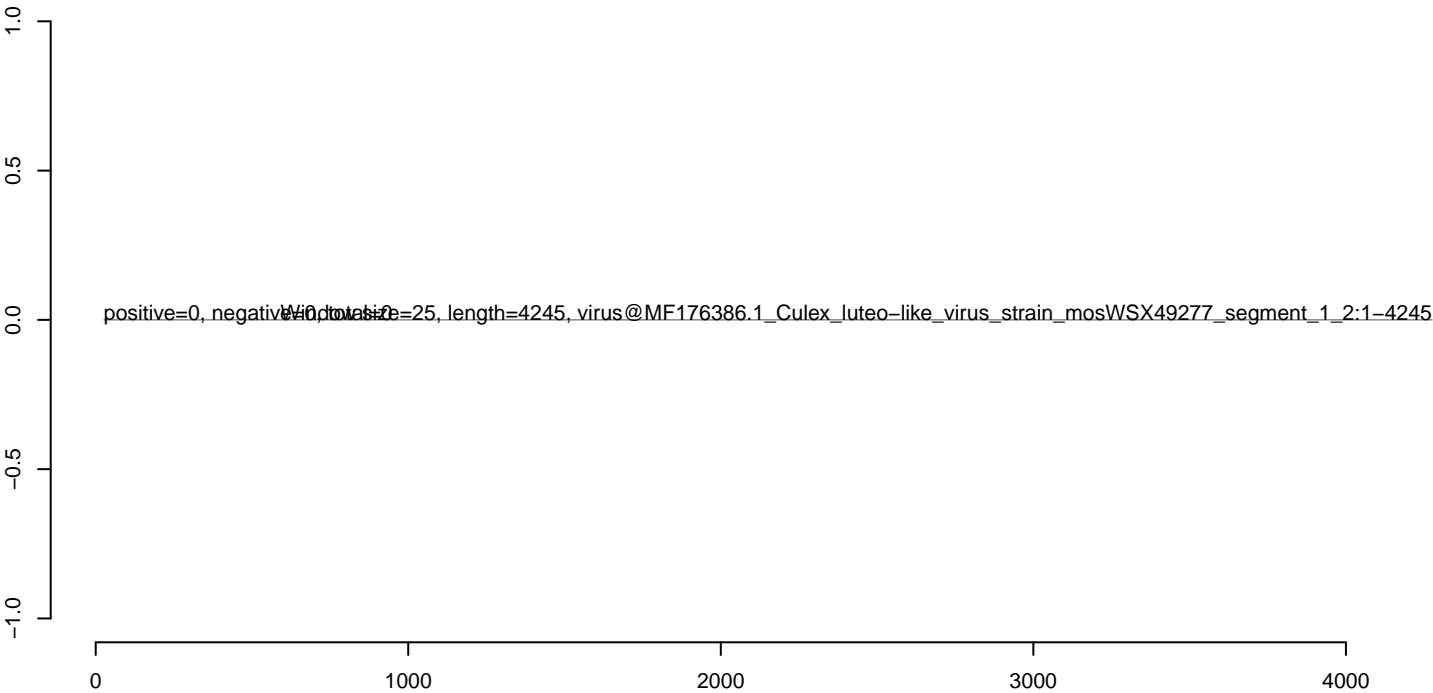
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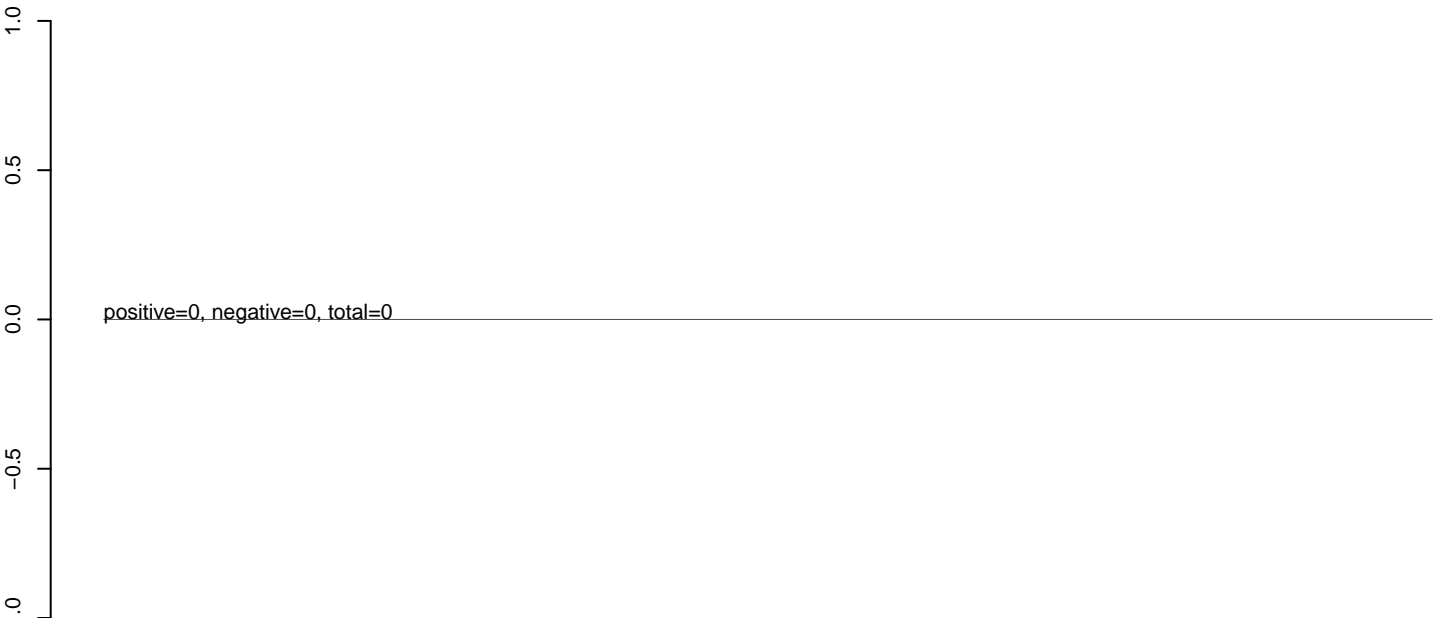
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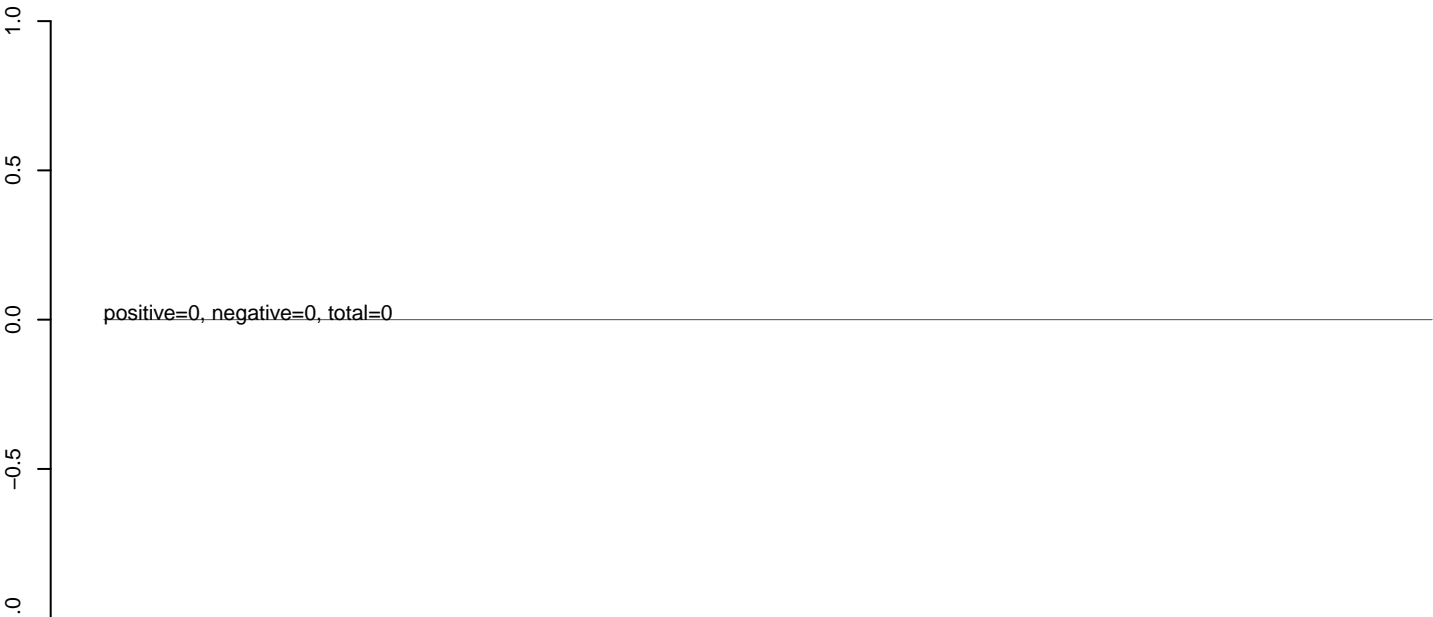
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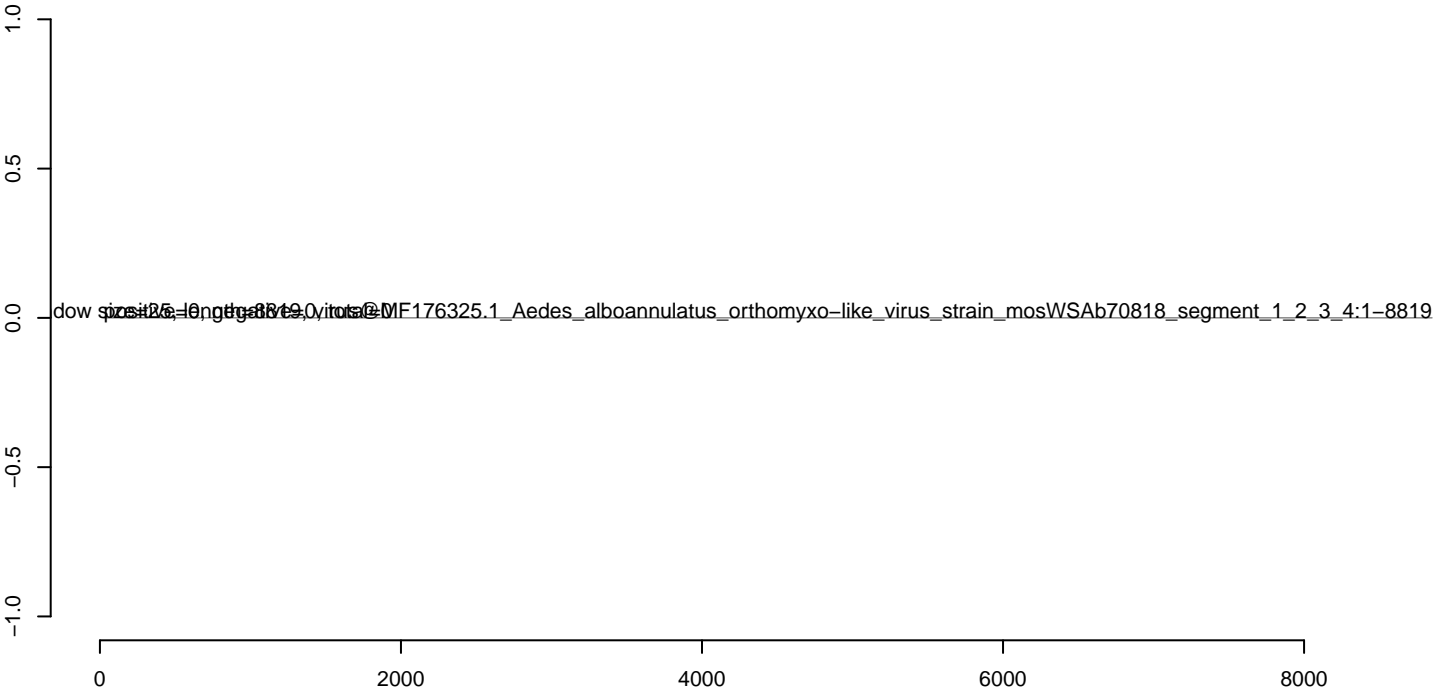
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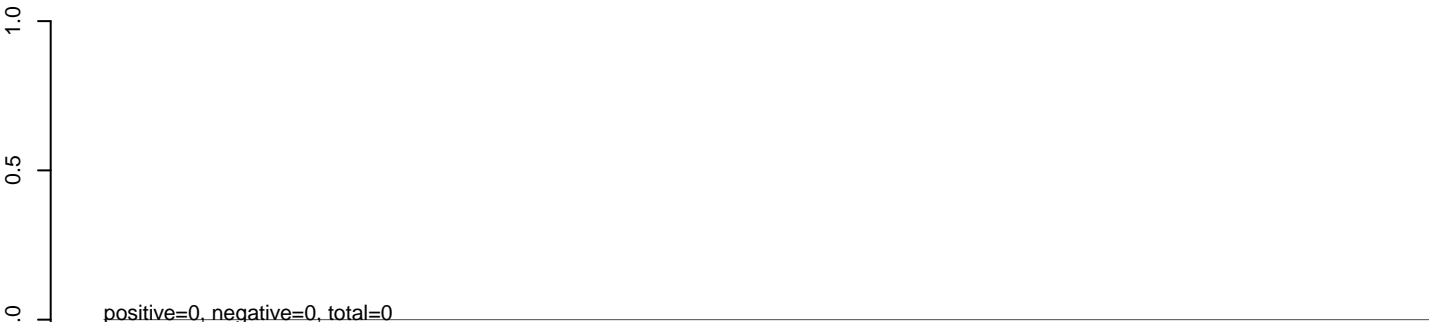
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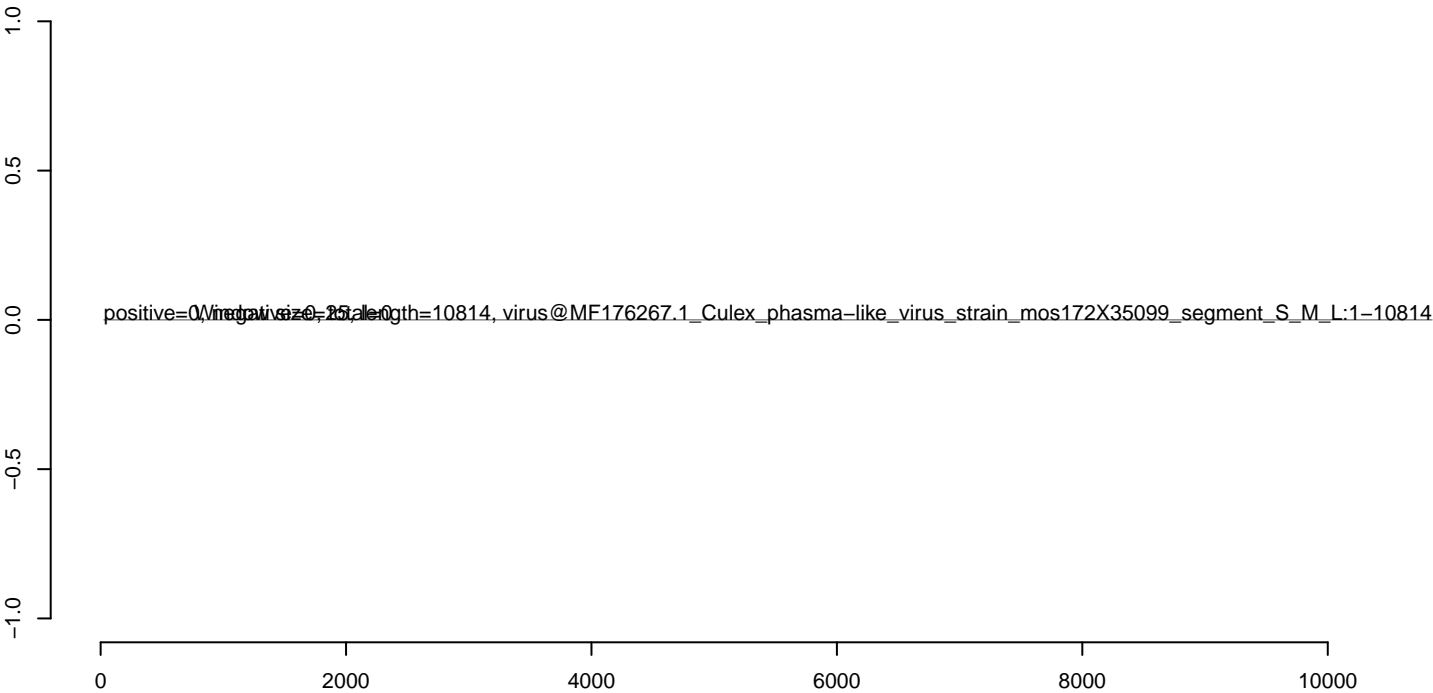
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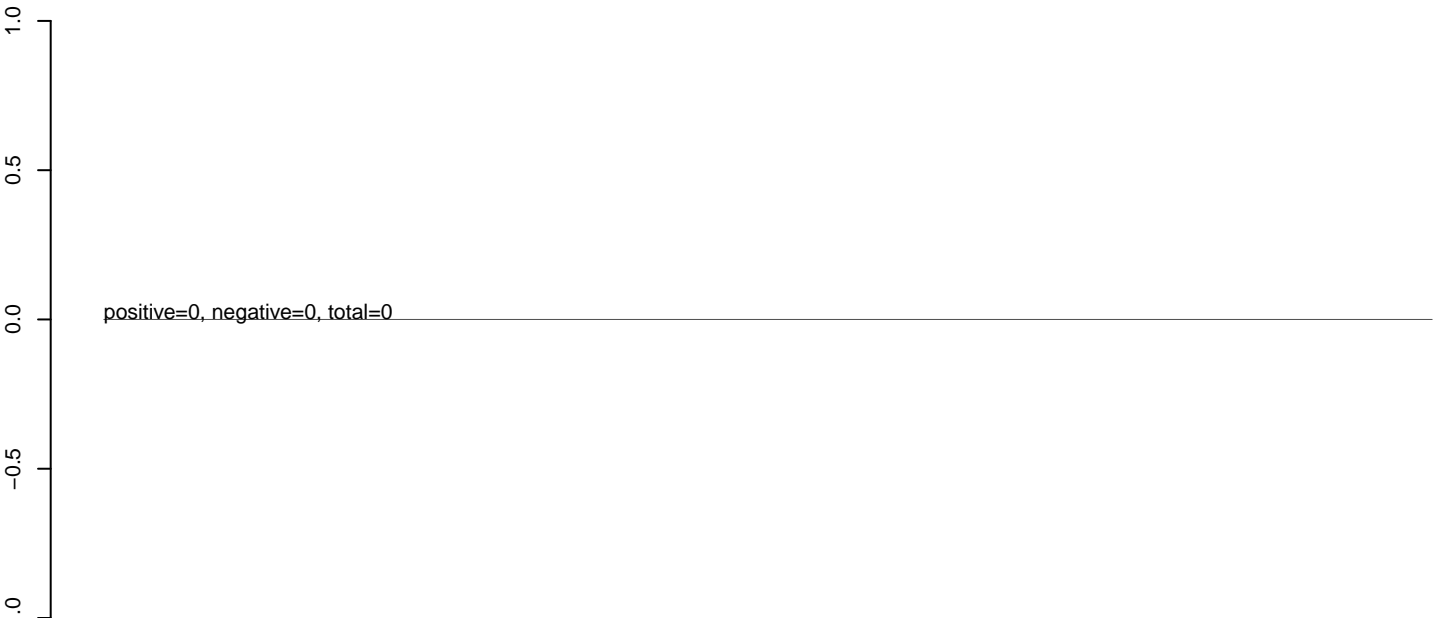
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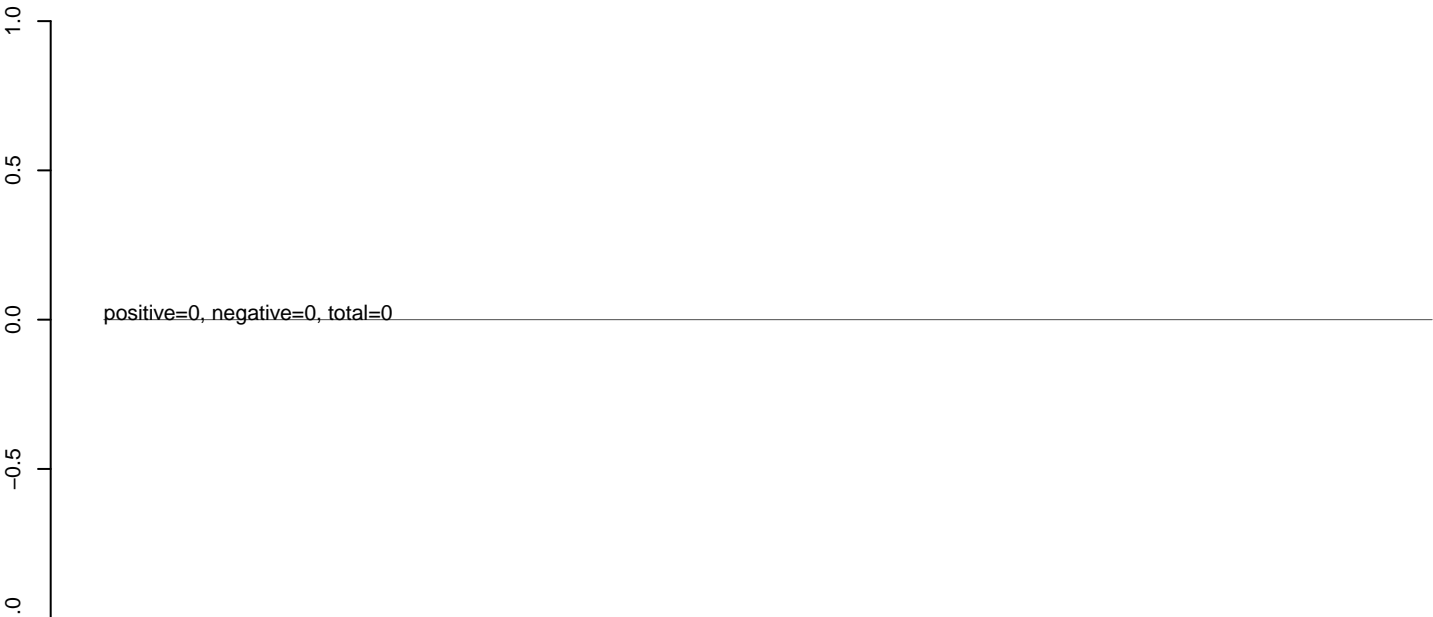
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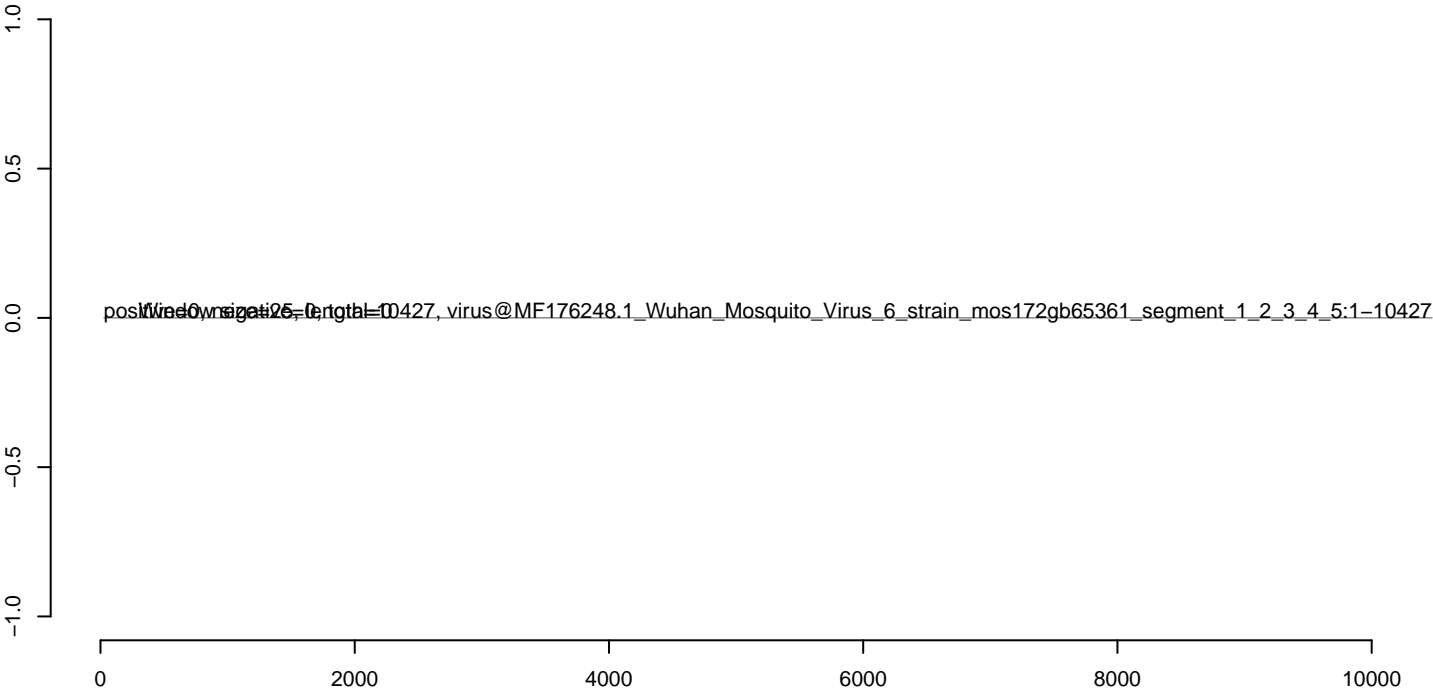
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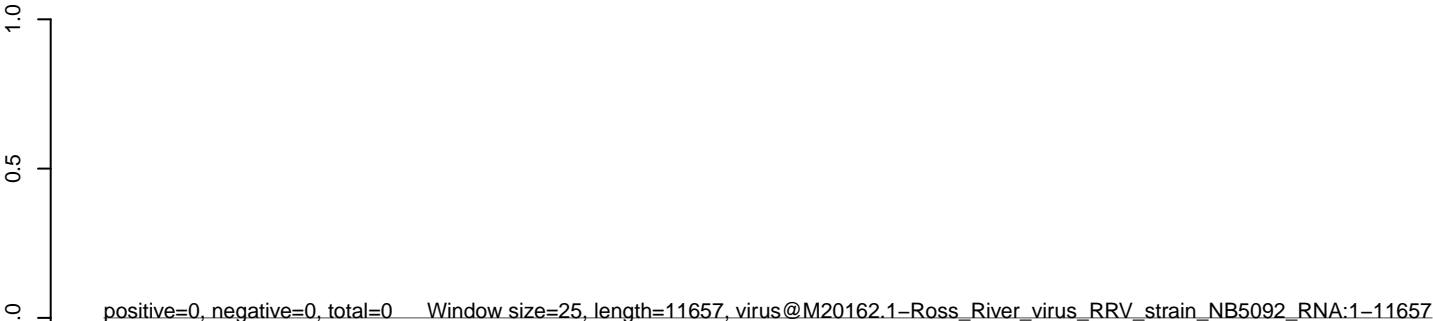
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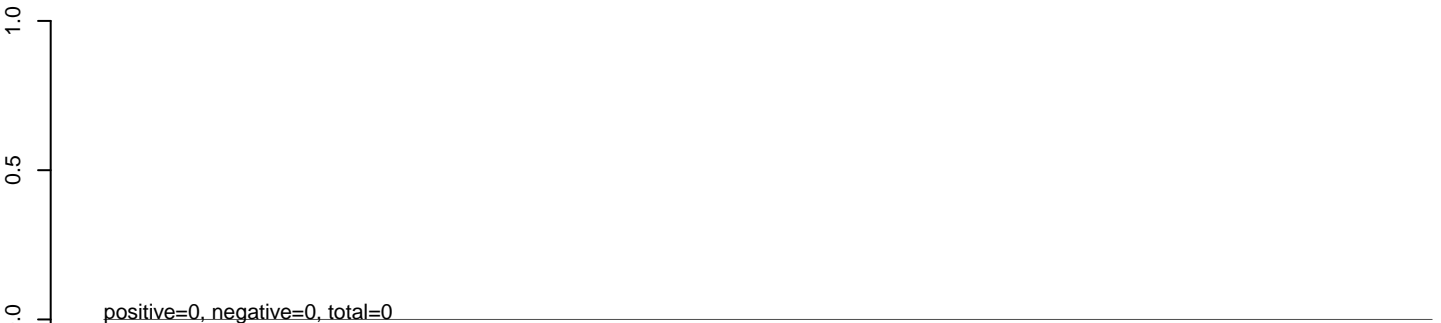


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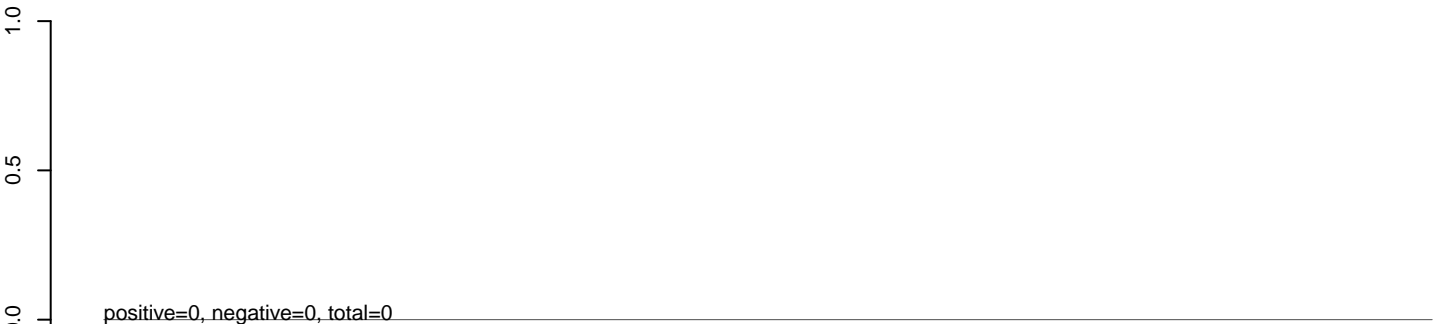


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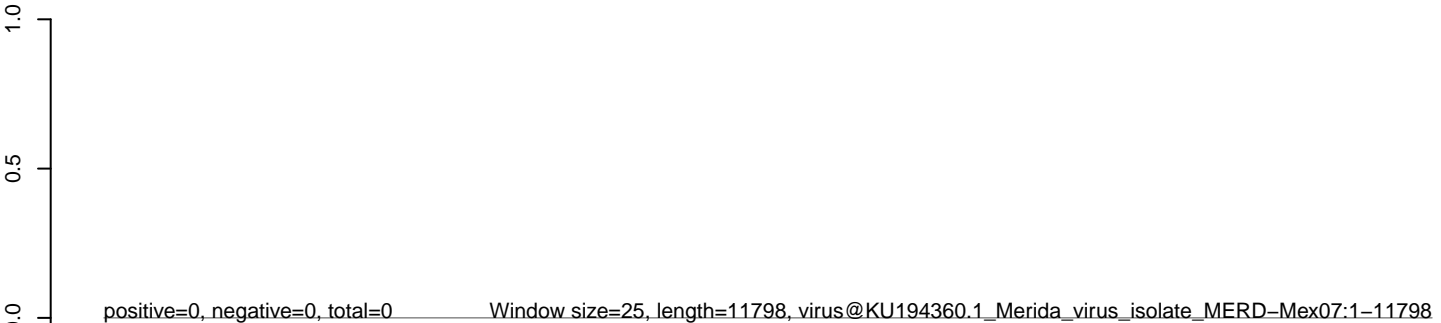
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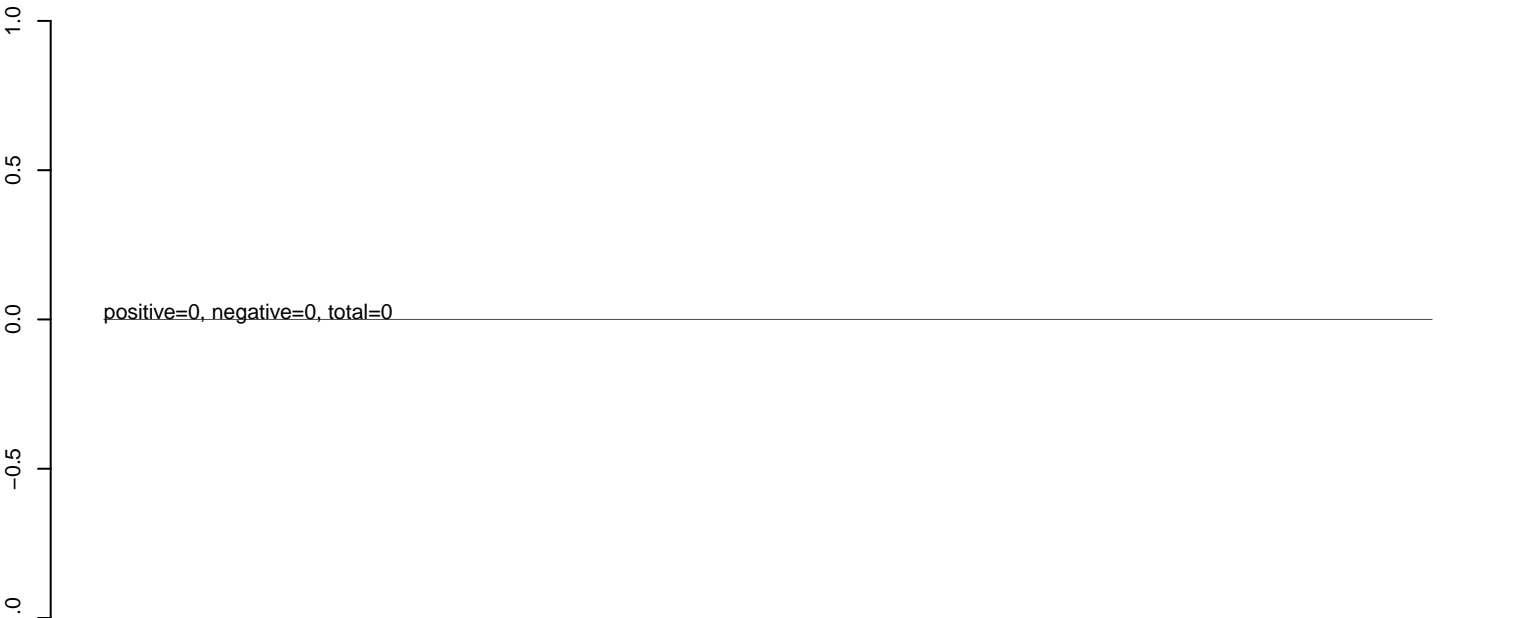


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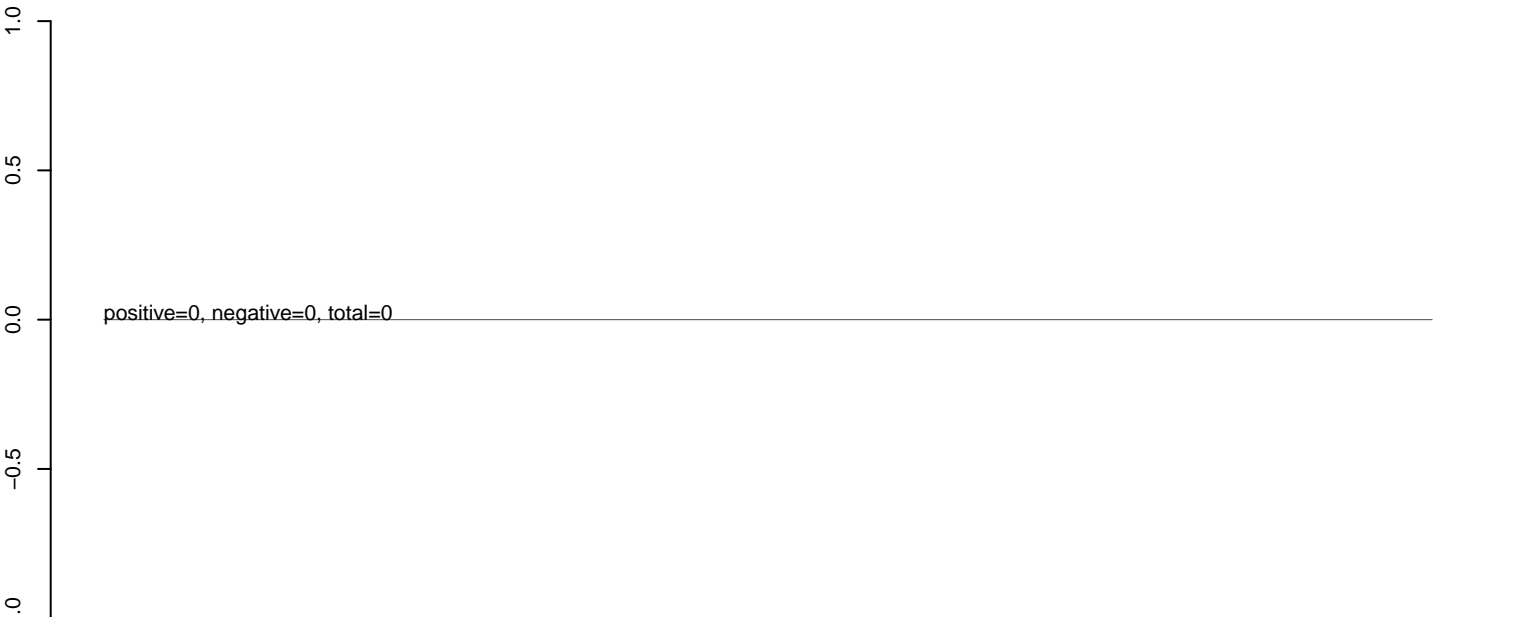


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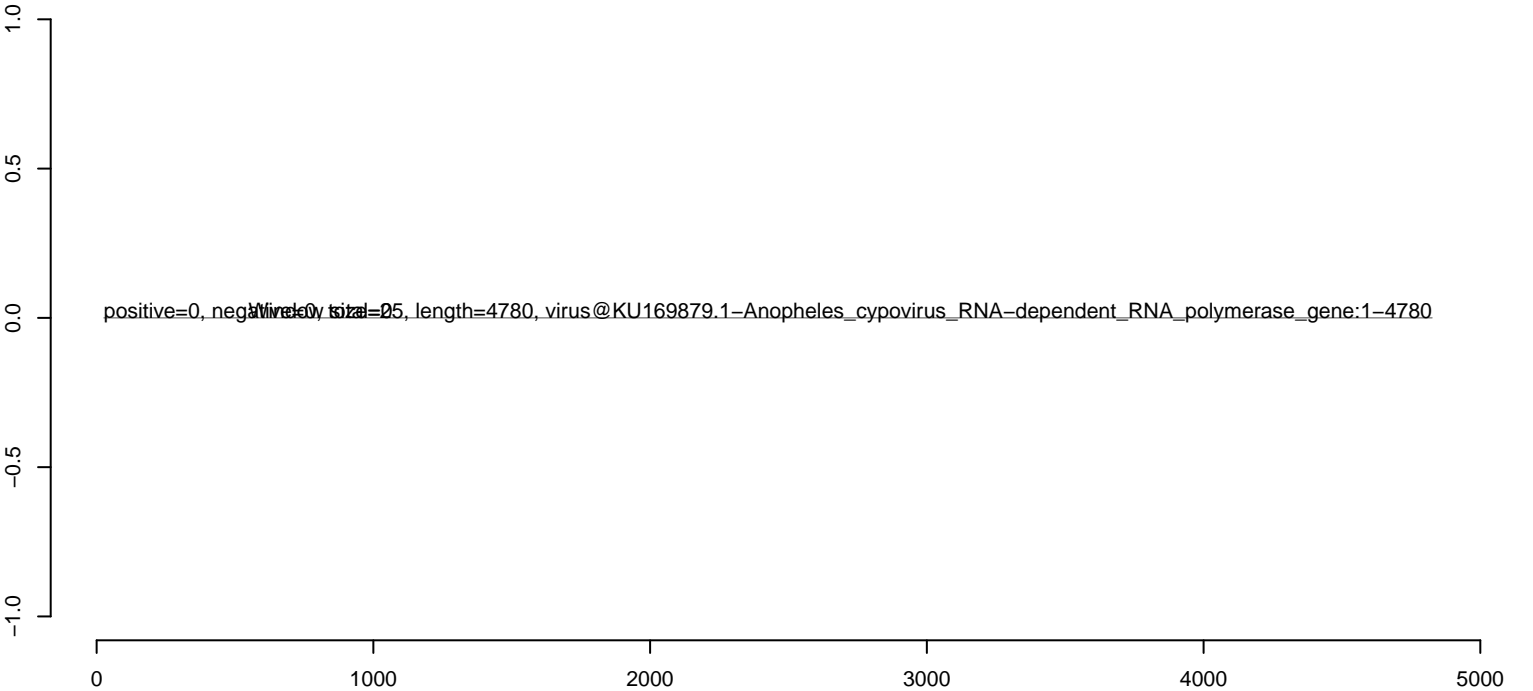
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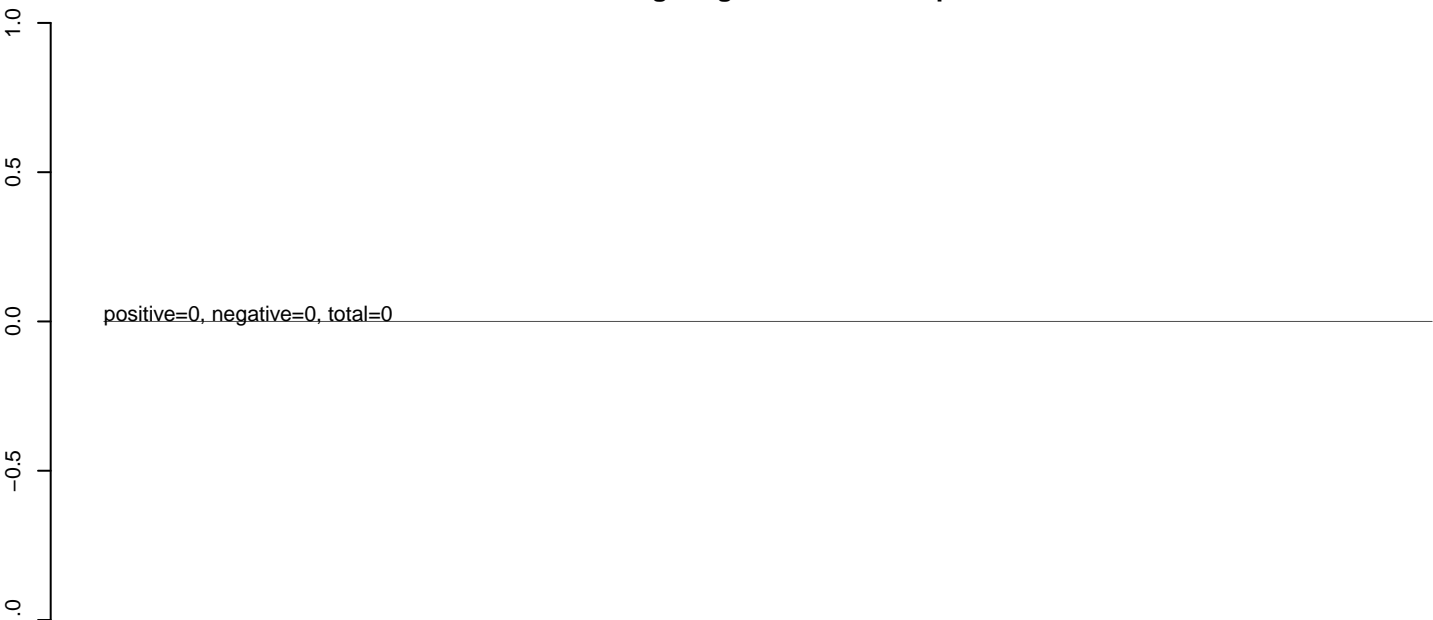
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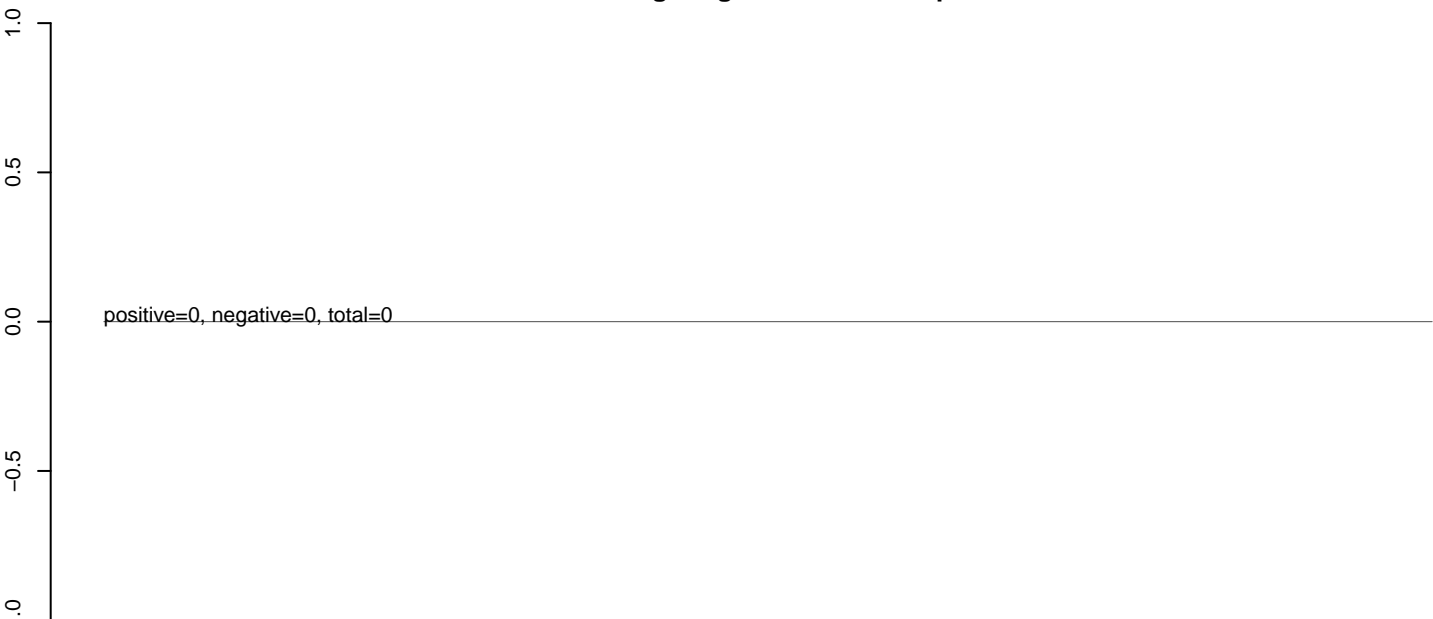
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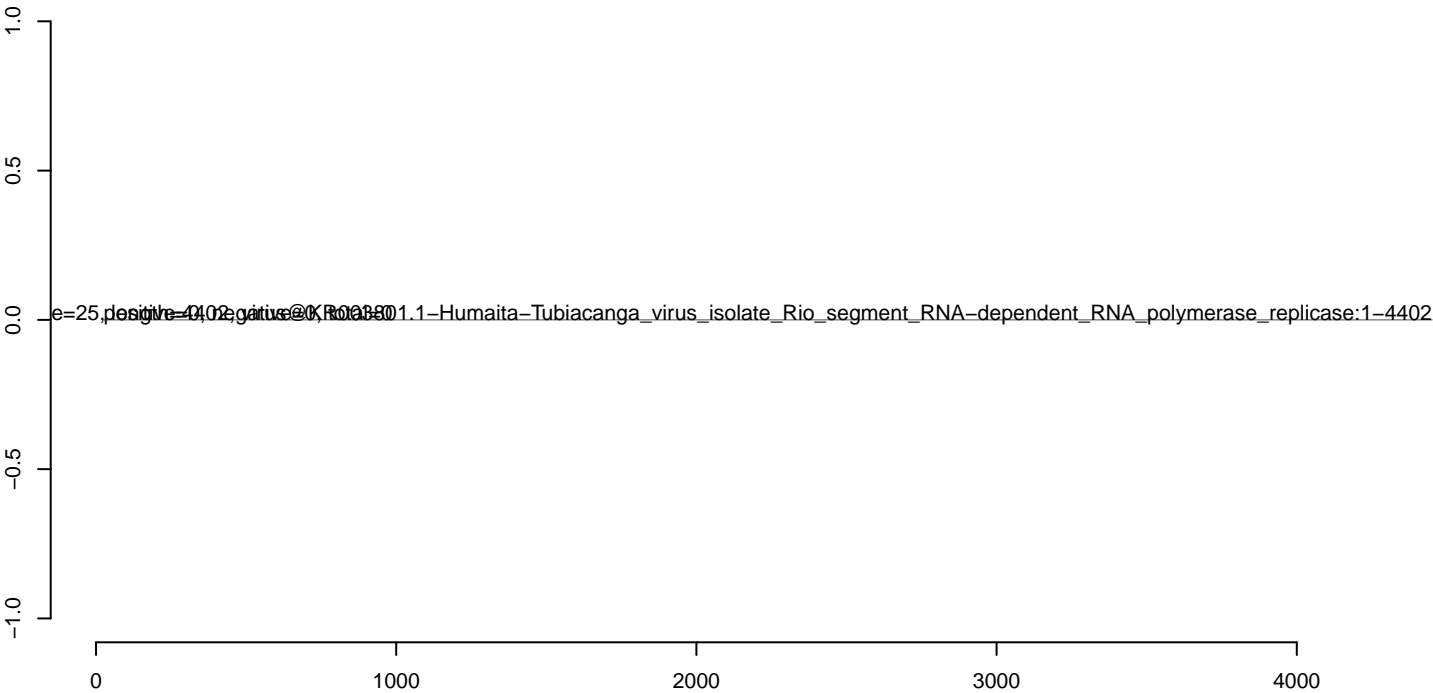
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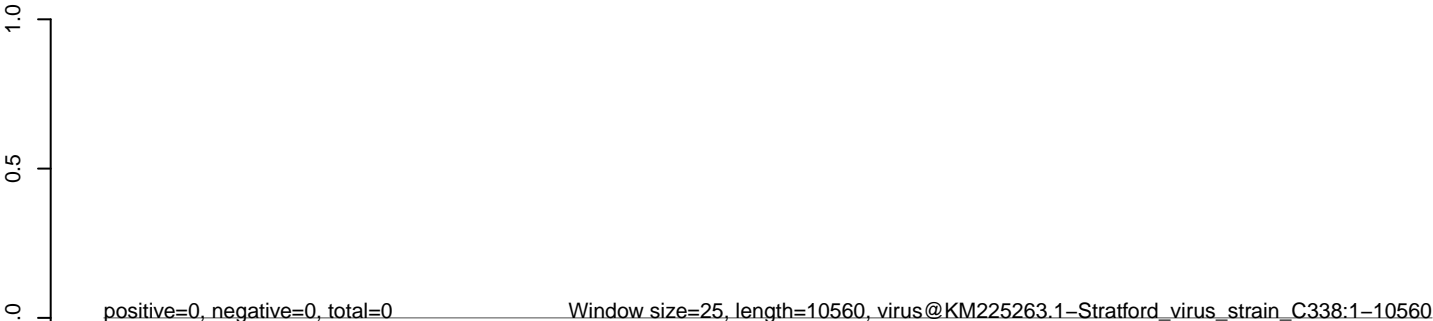
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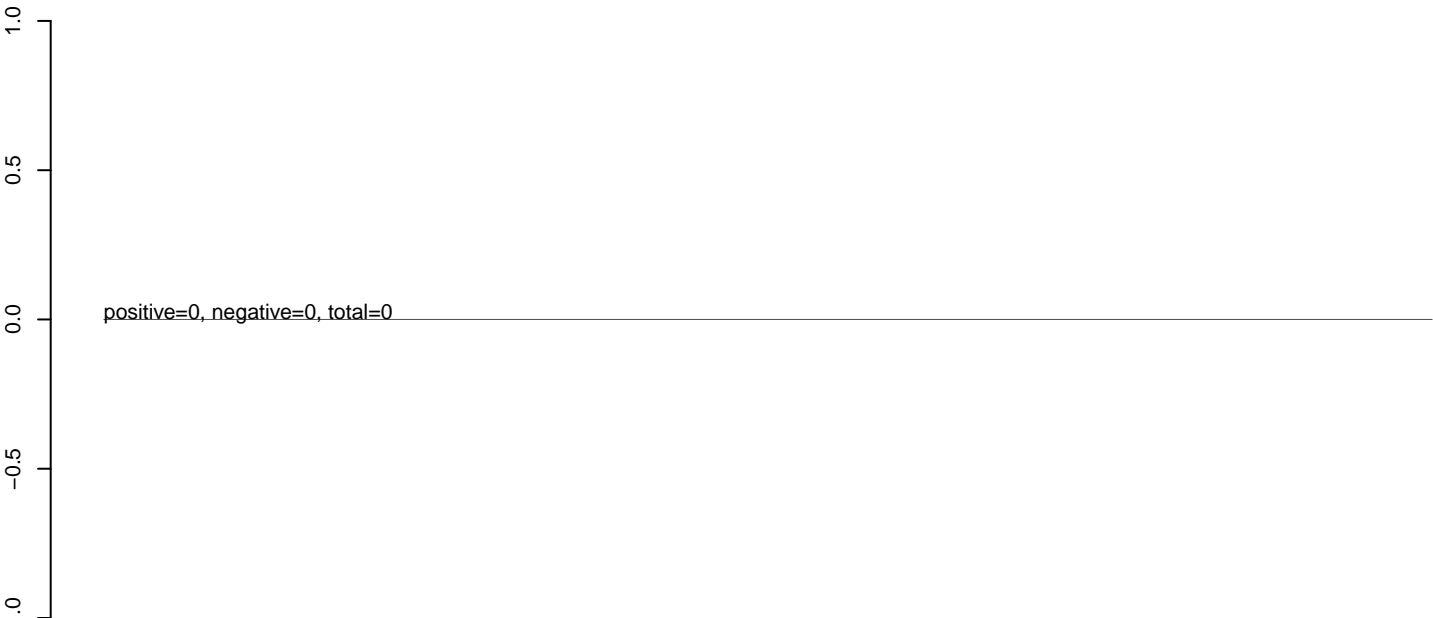


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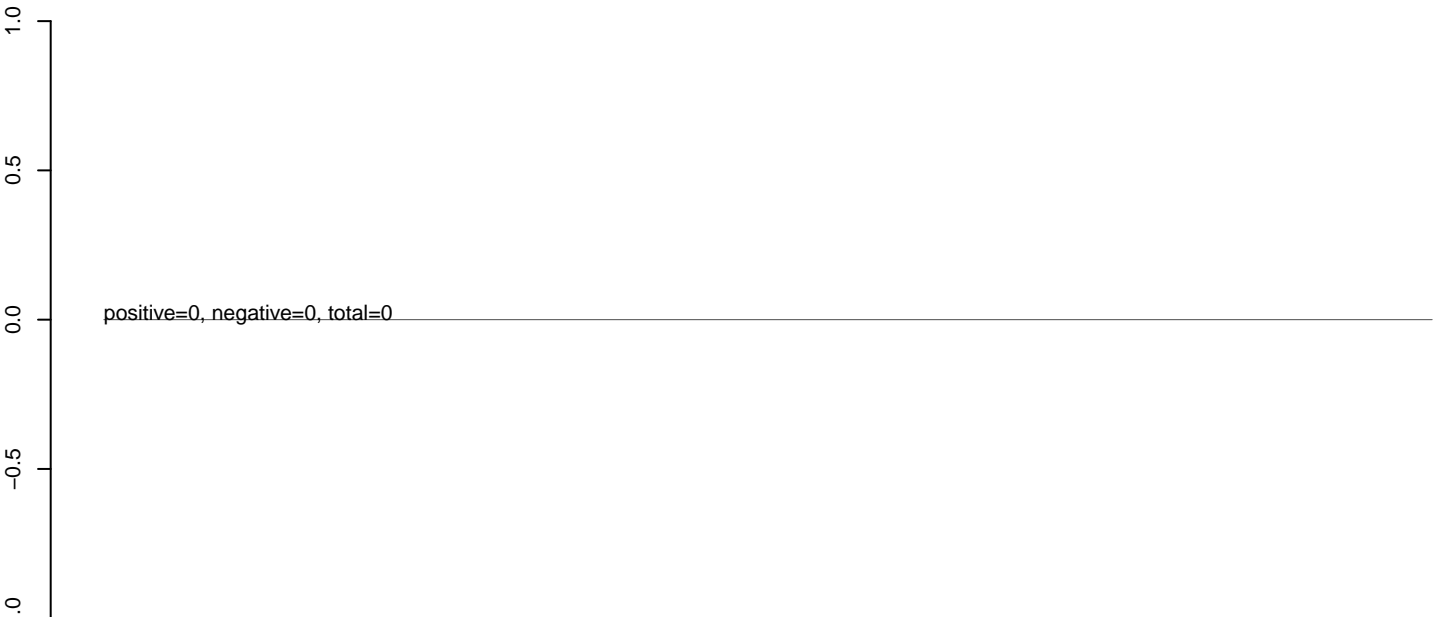


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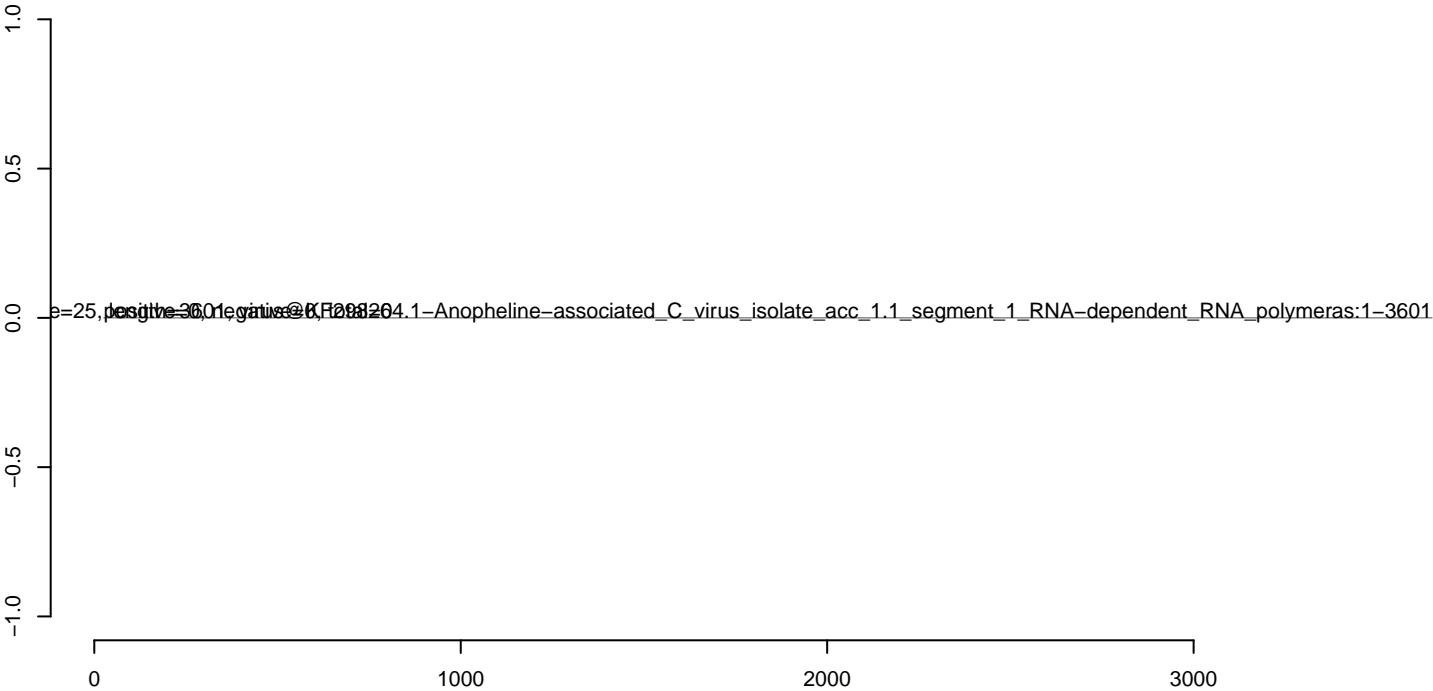
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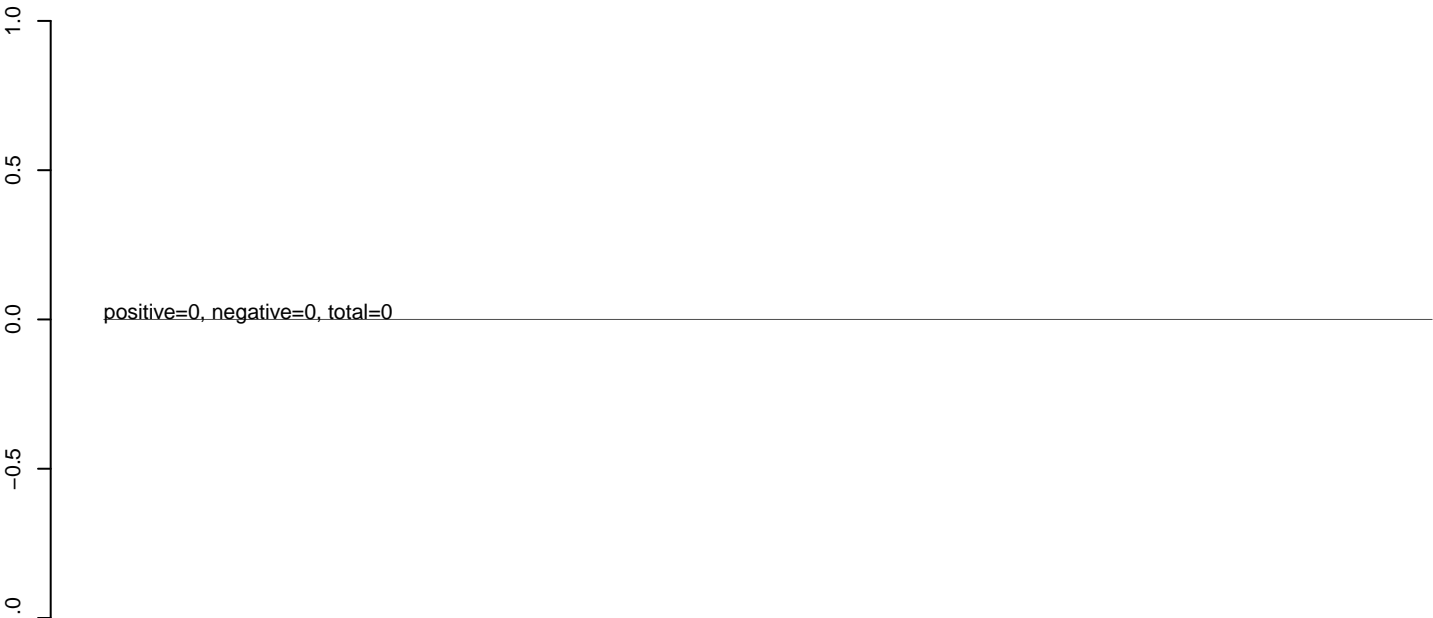
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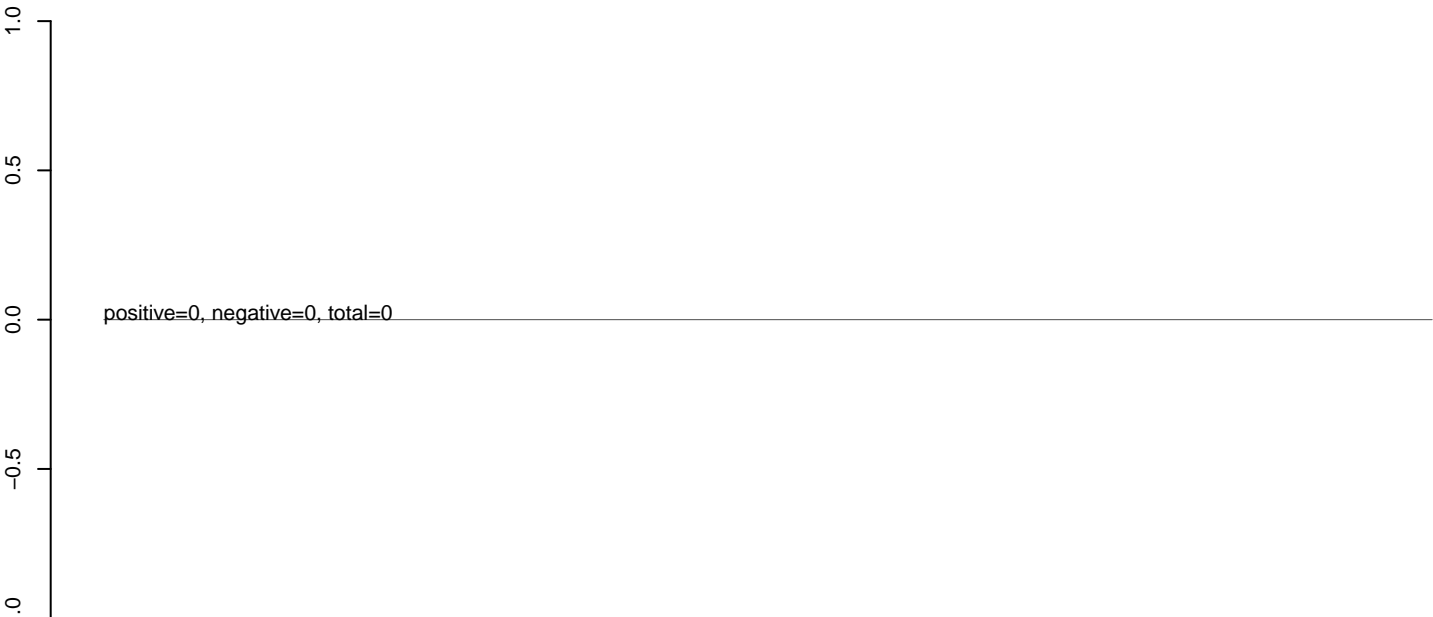
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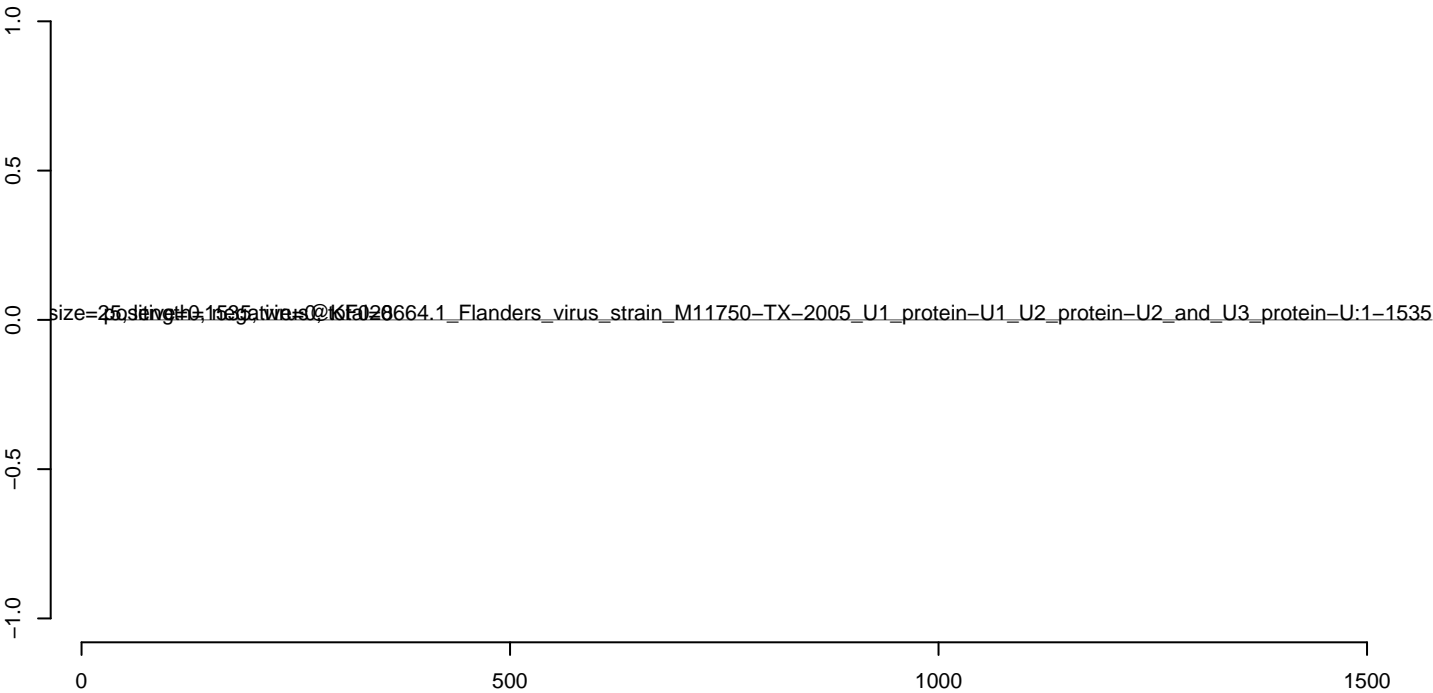
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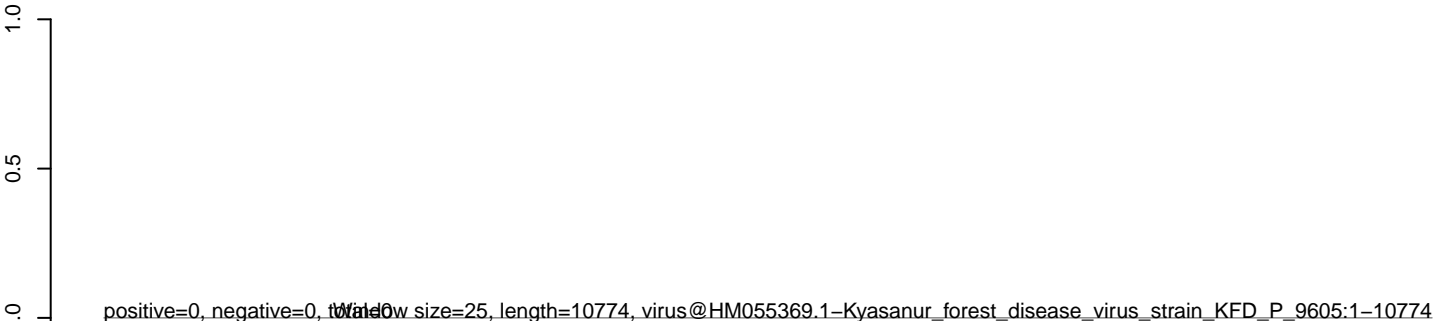
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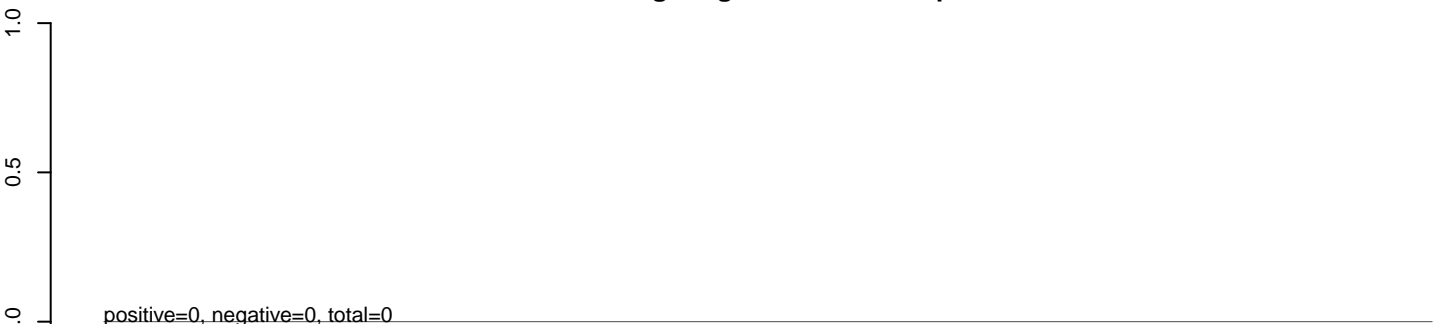
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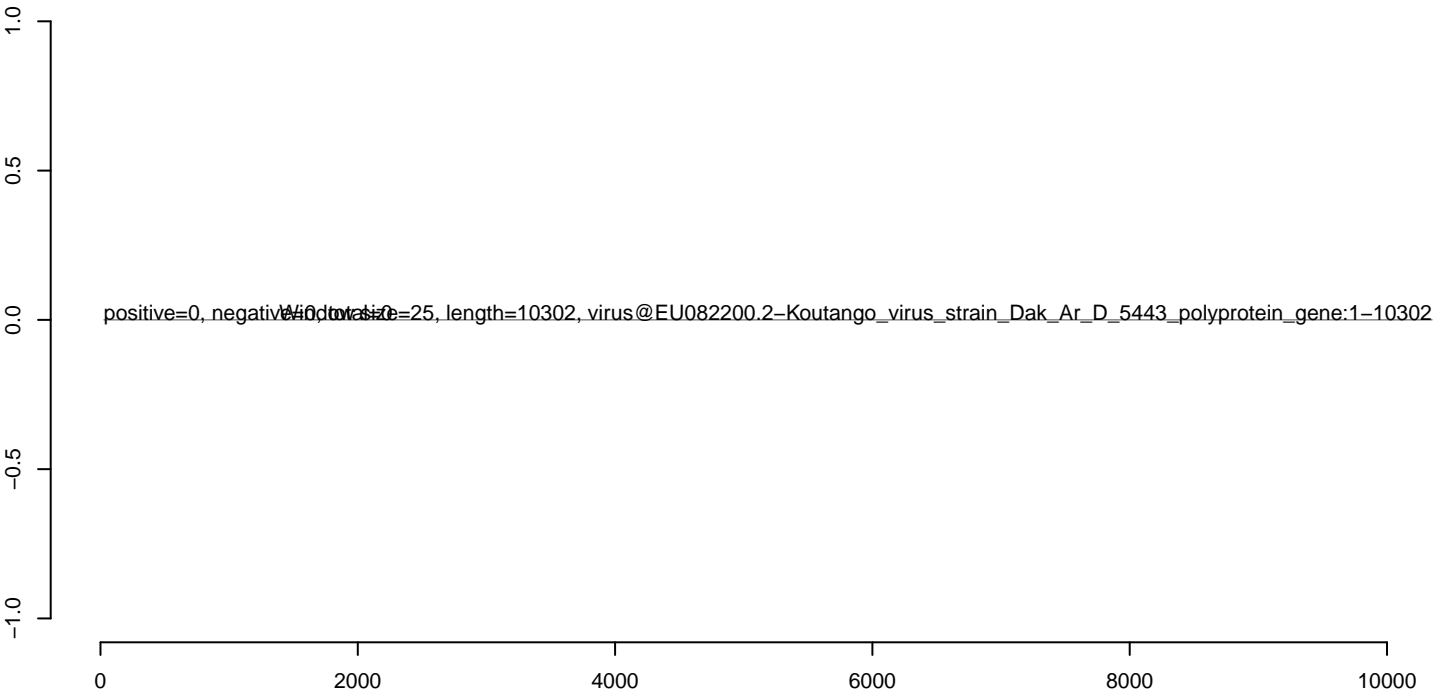
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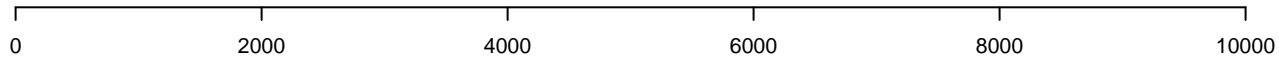
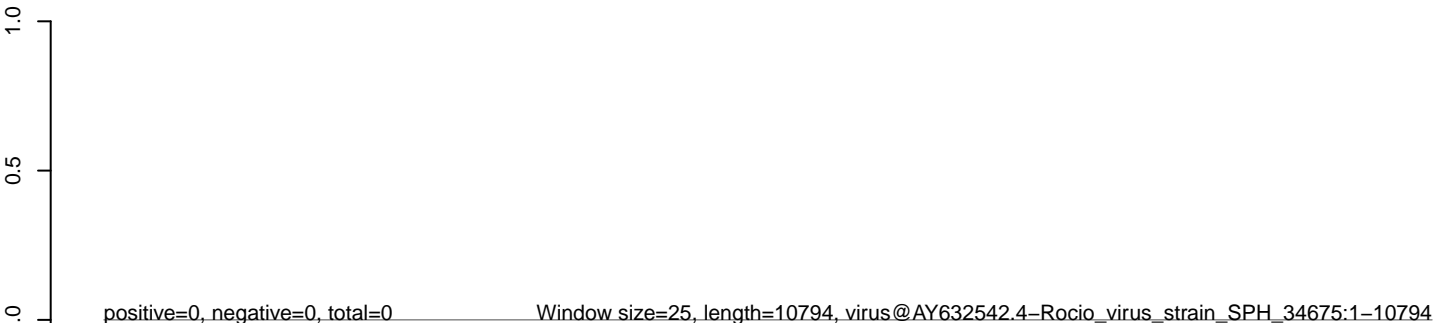
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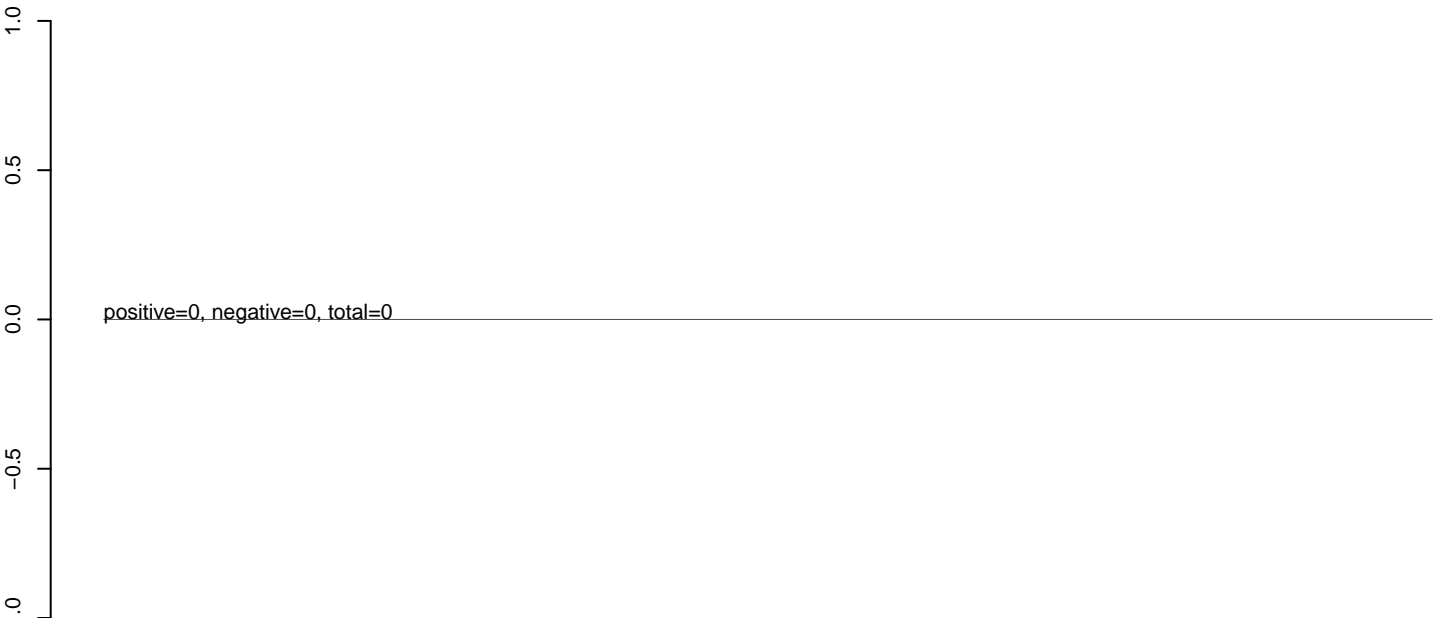
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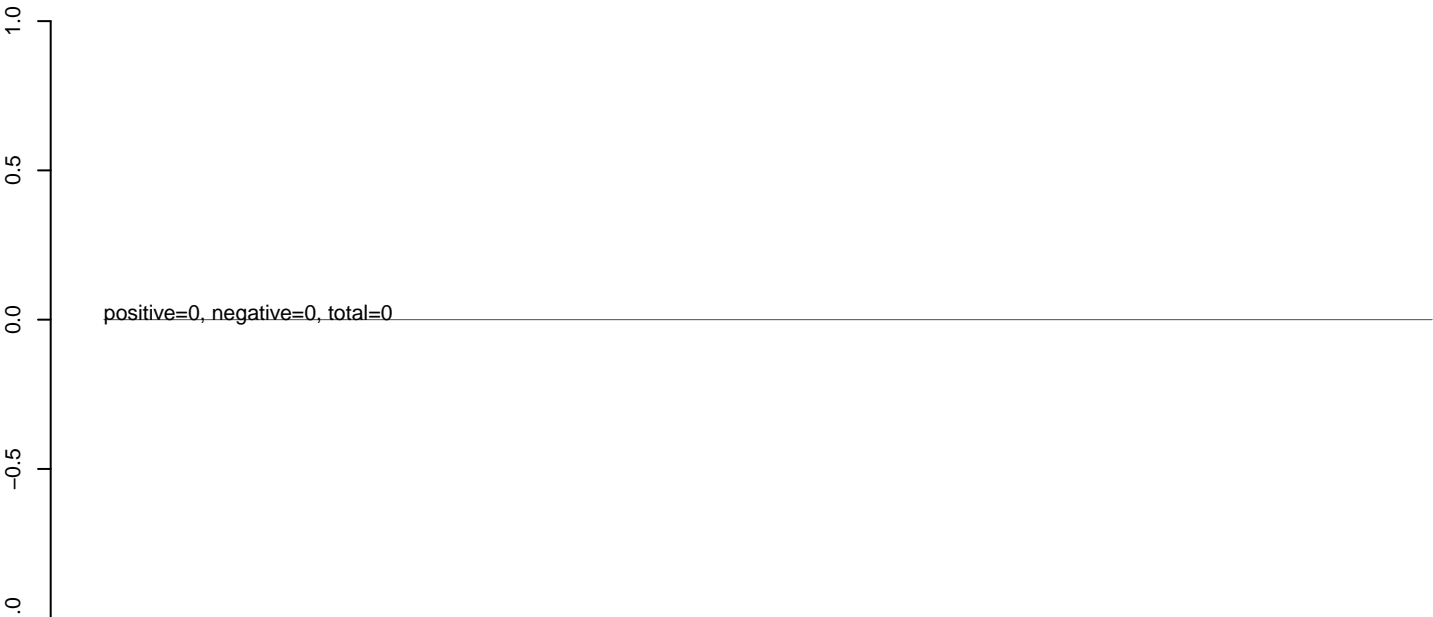
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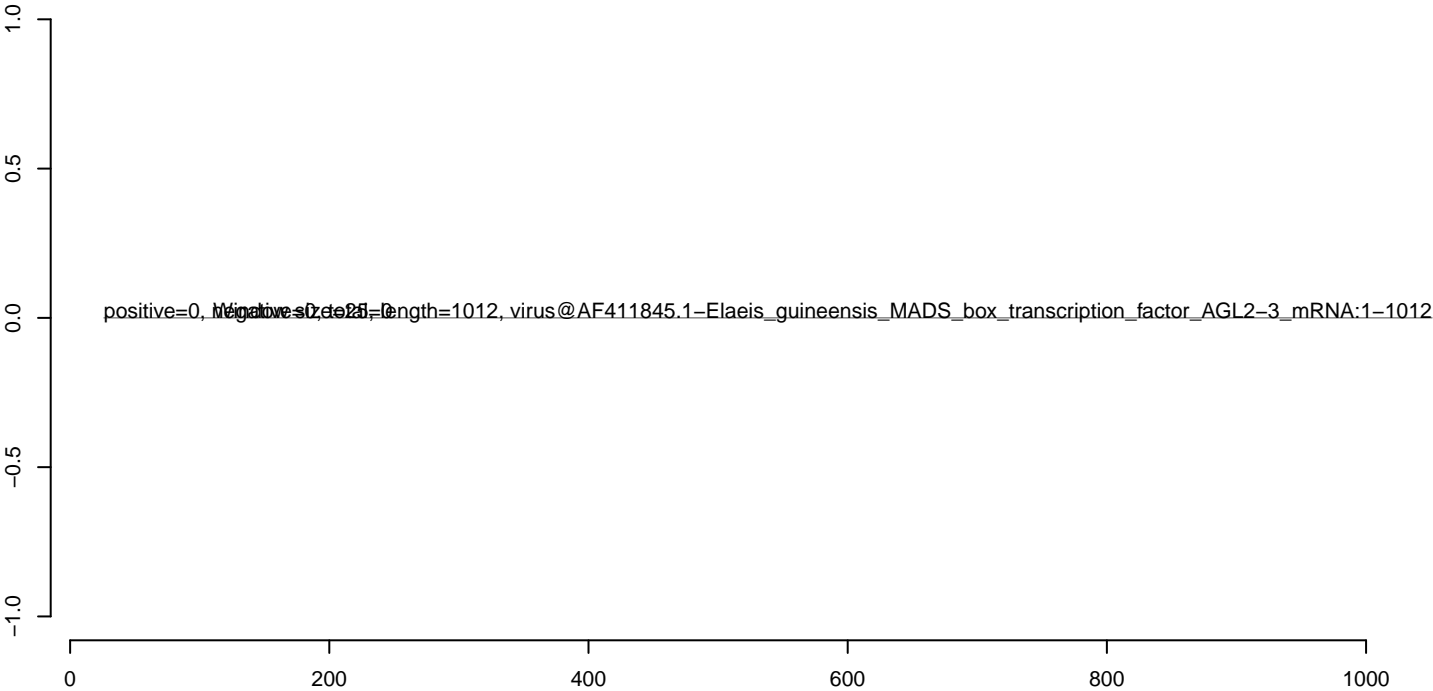
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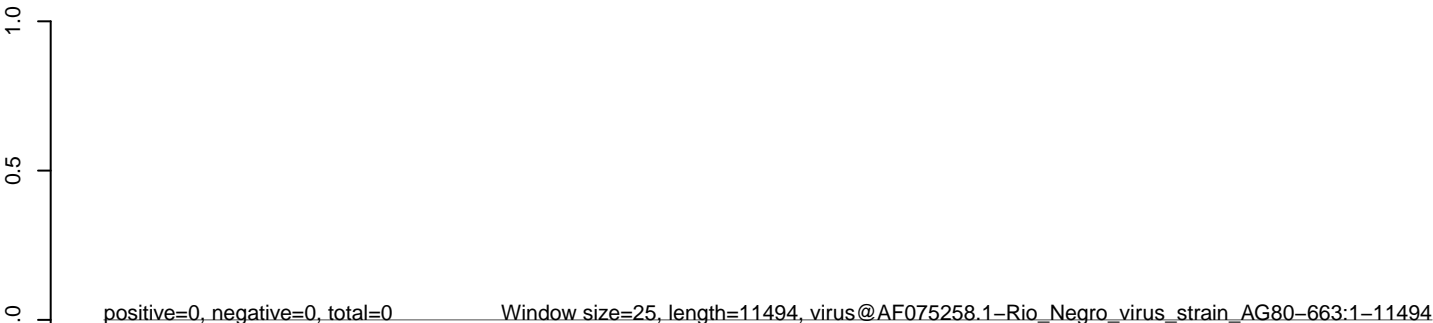
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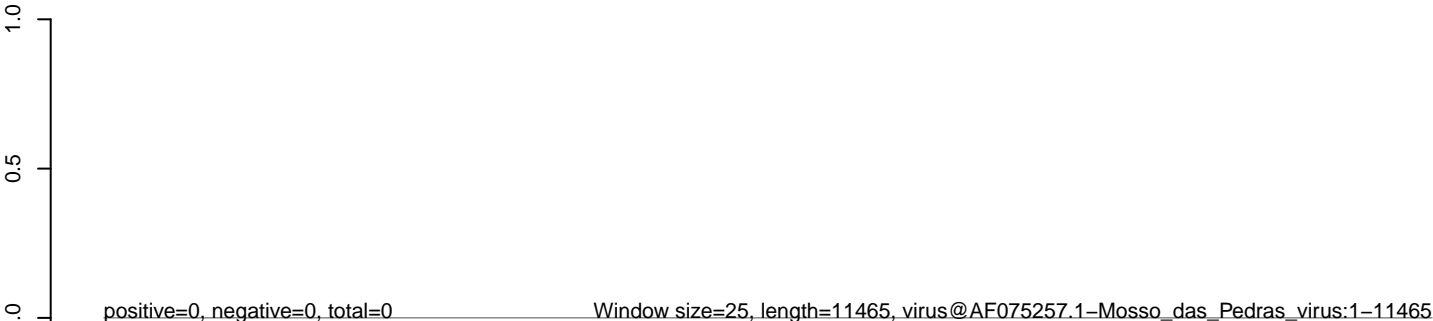
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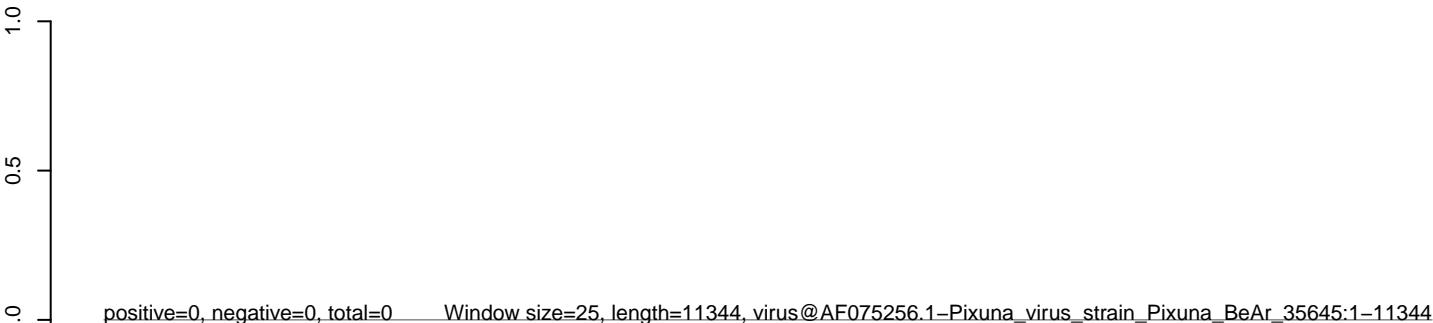
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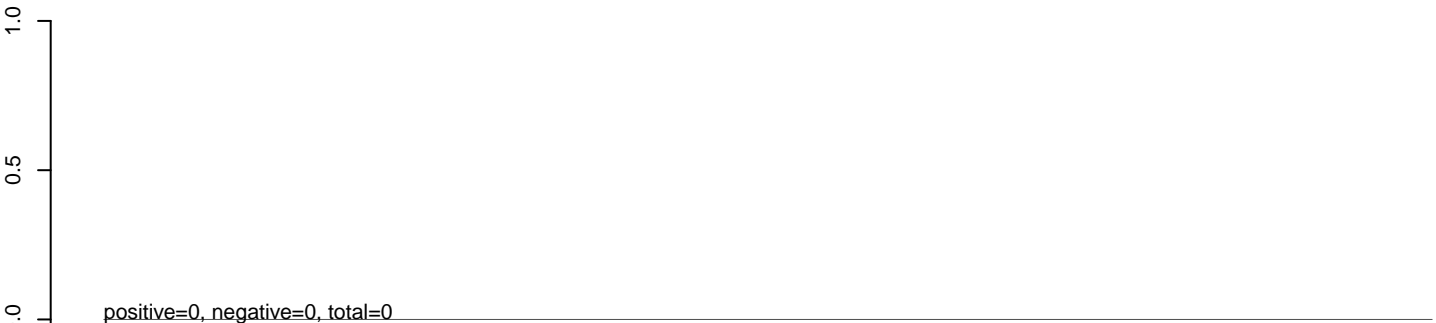


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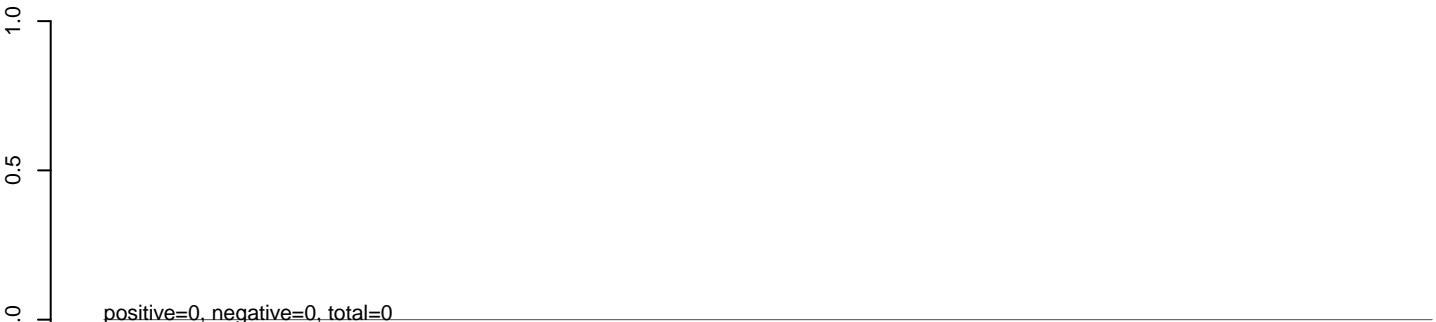


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