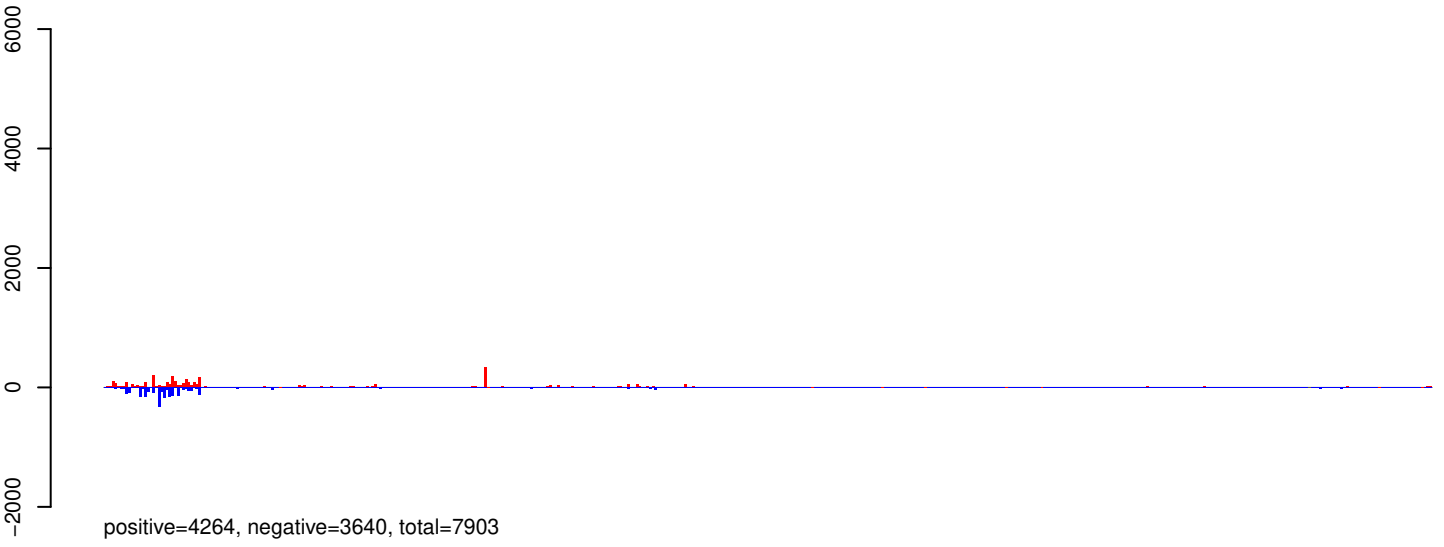
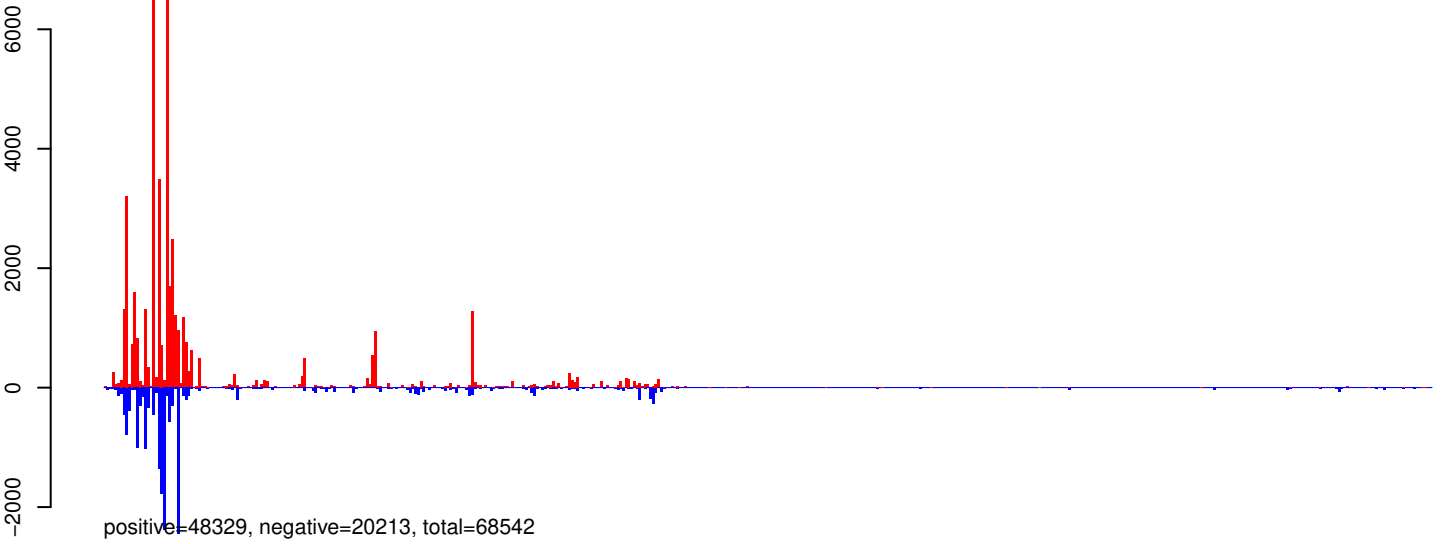


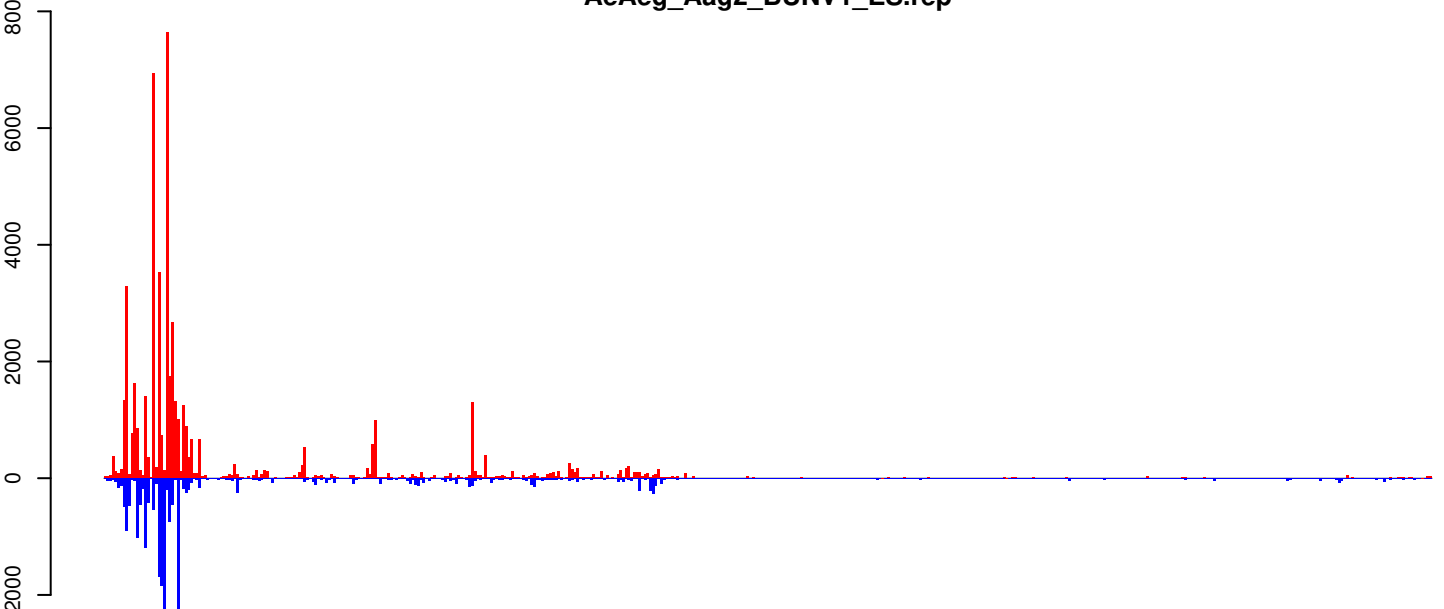
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



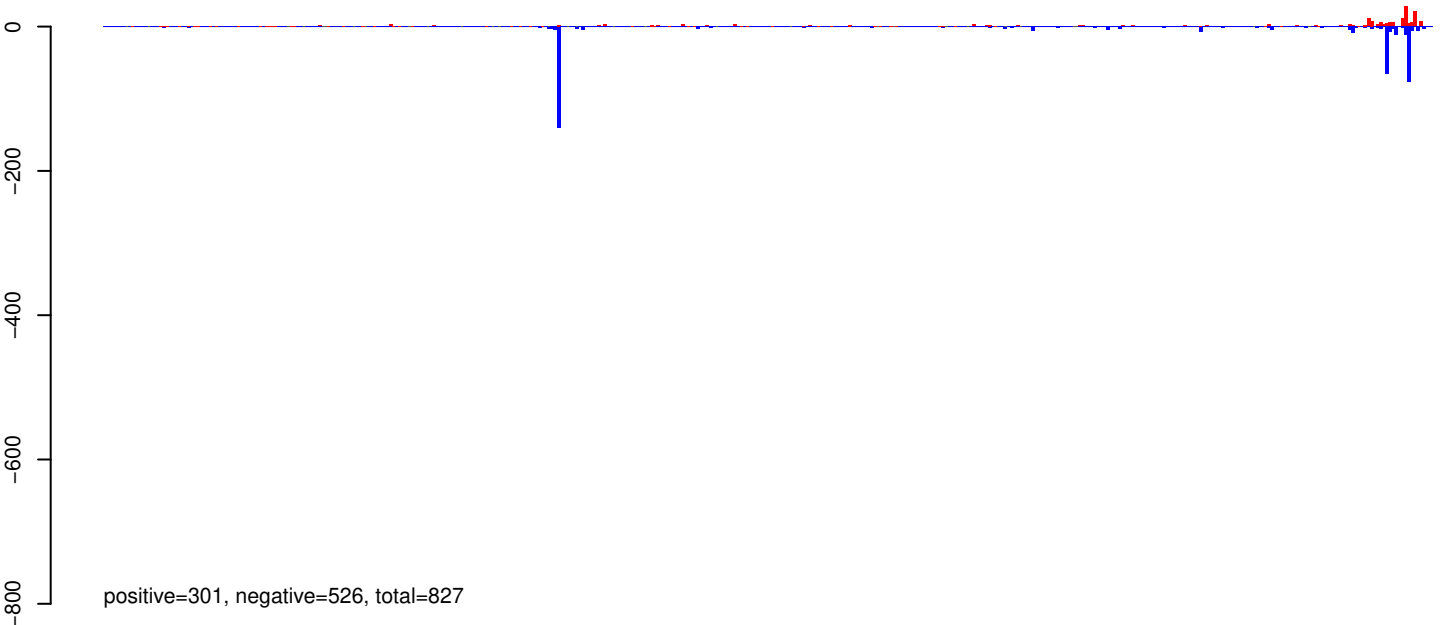
AeAeg_Aag2_BUNV1_ES.rep



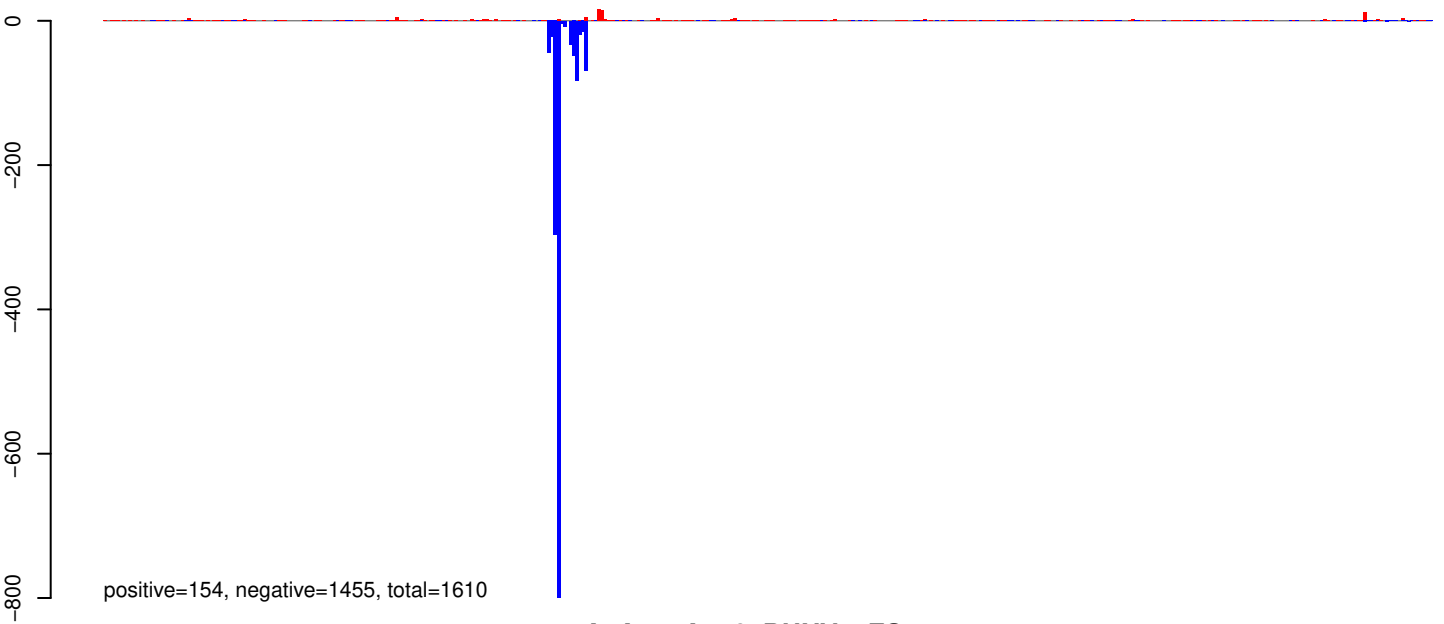
52589, 26652, 79241 @ NC_001927.1-Bunyamwera_virus_segment_S_plus_segment_M_plus_segment_L:1-12294

0 2000 4000 6000 8000 10000 12000

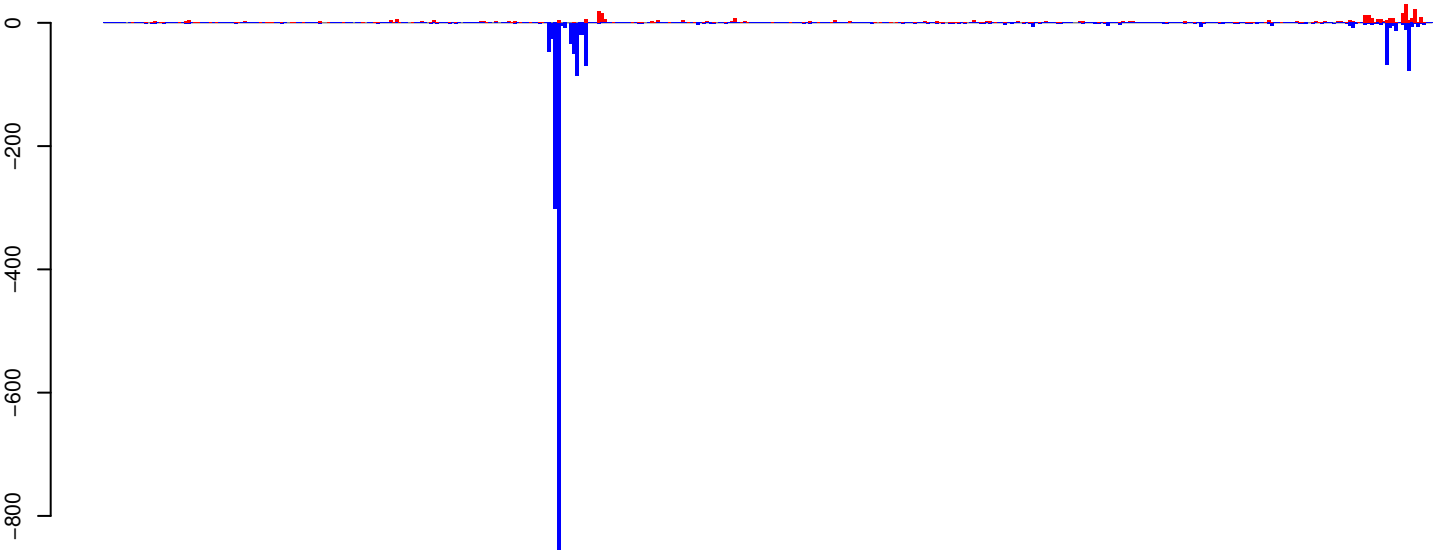
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AeAeg_Aag2_BUNV1_ES.24_35.rep



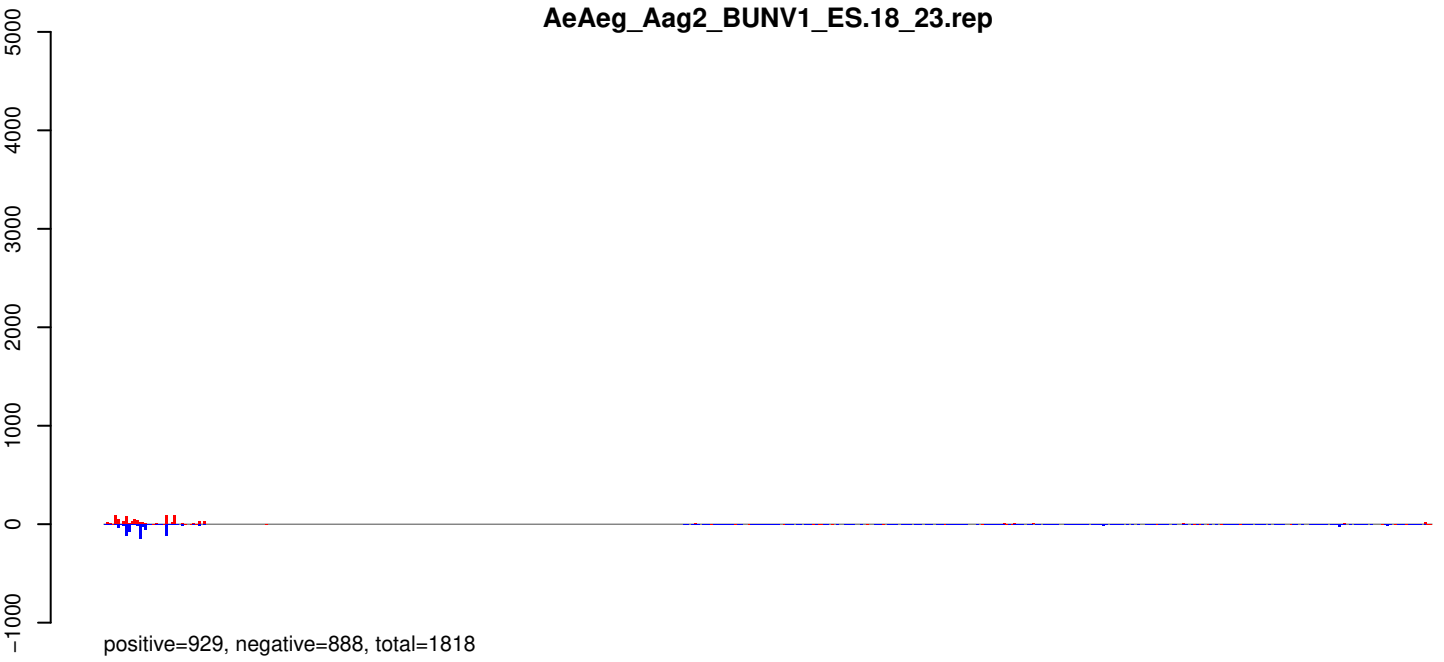
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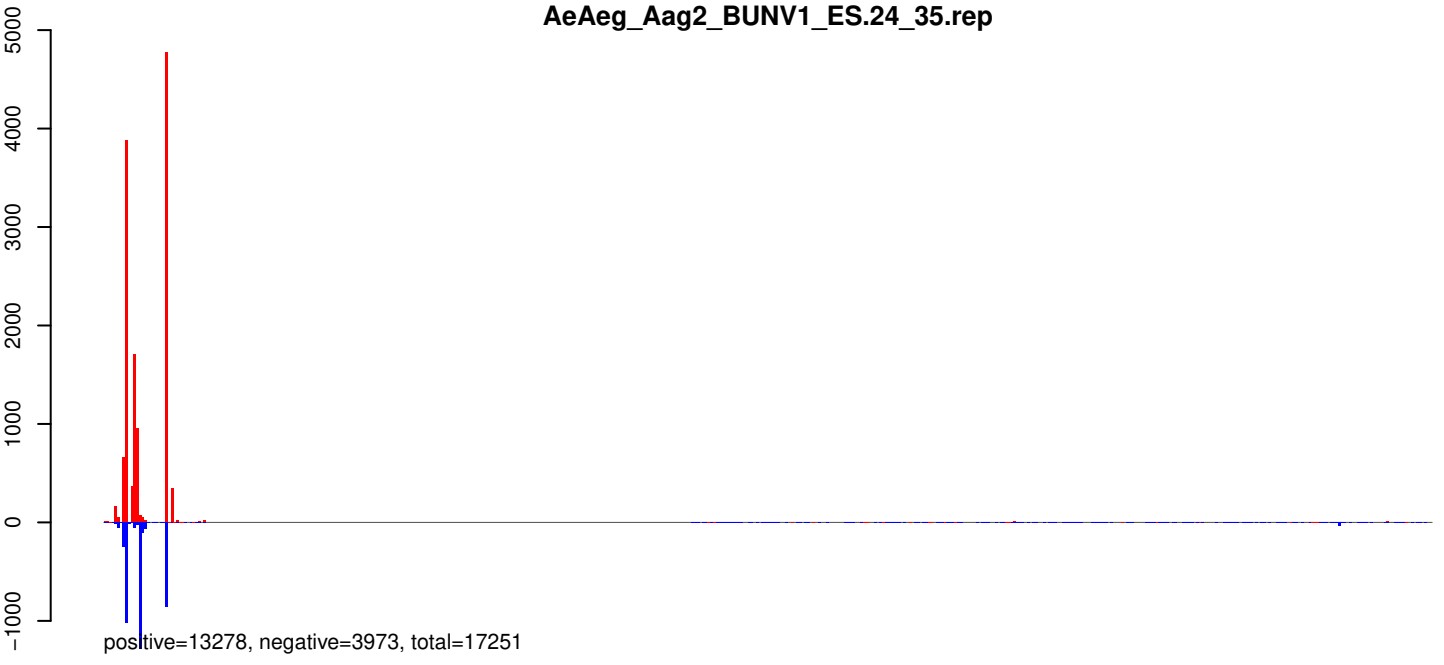
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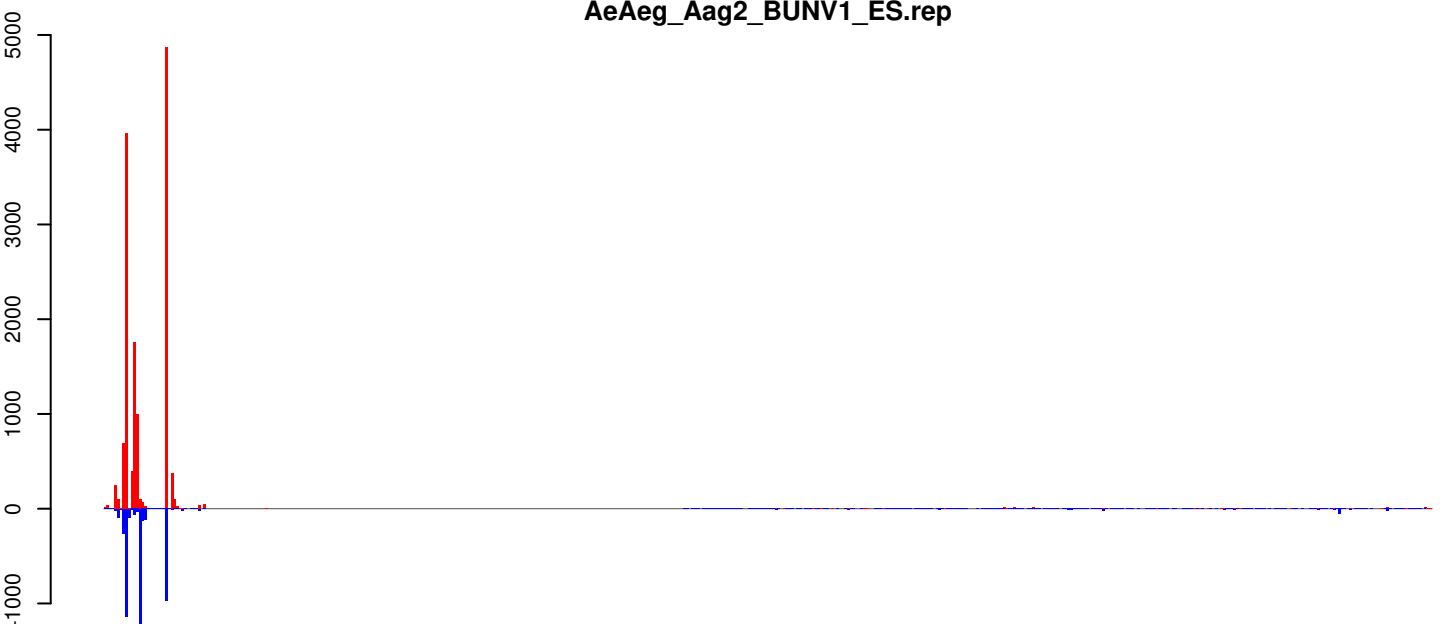
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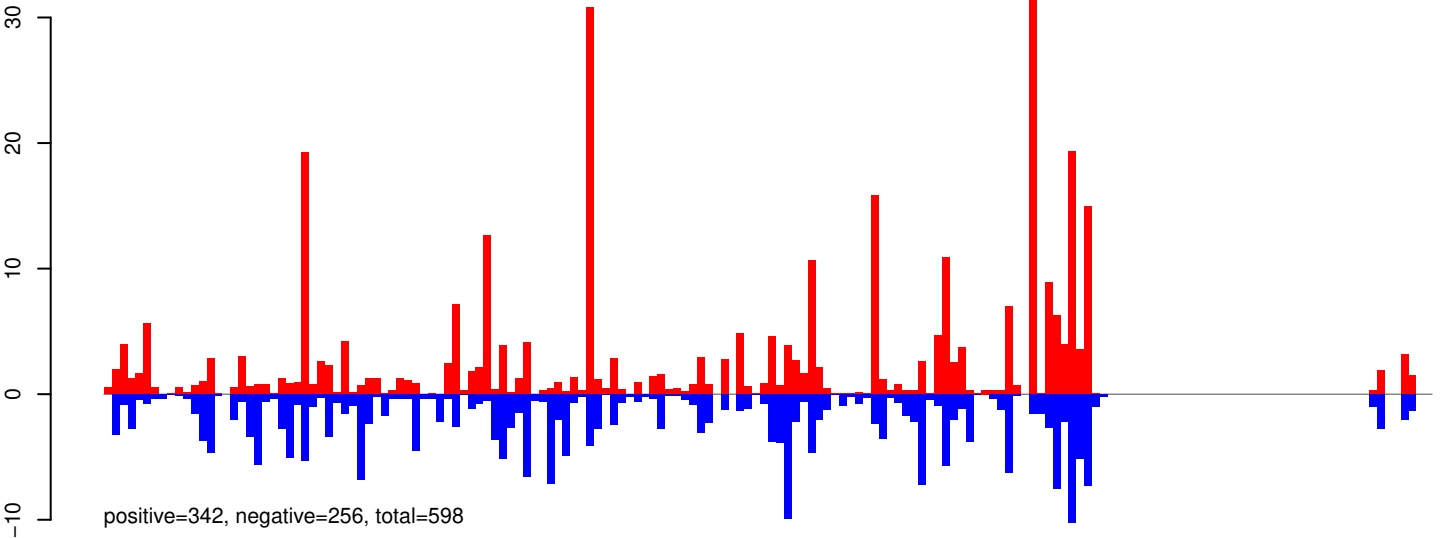
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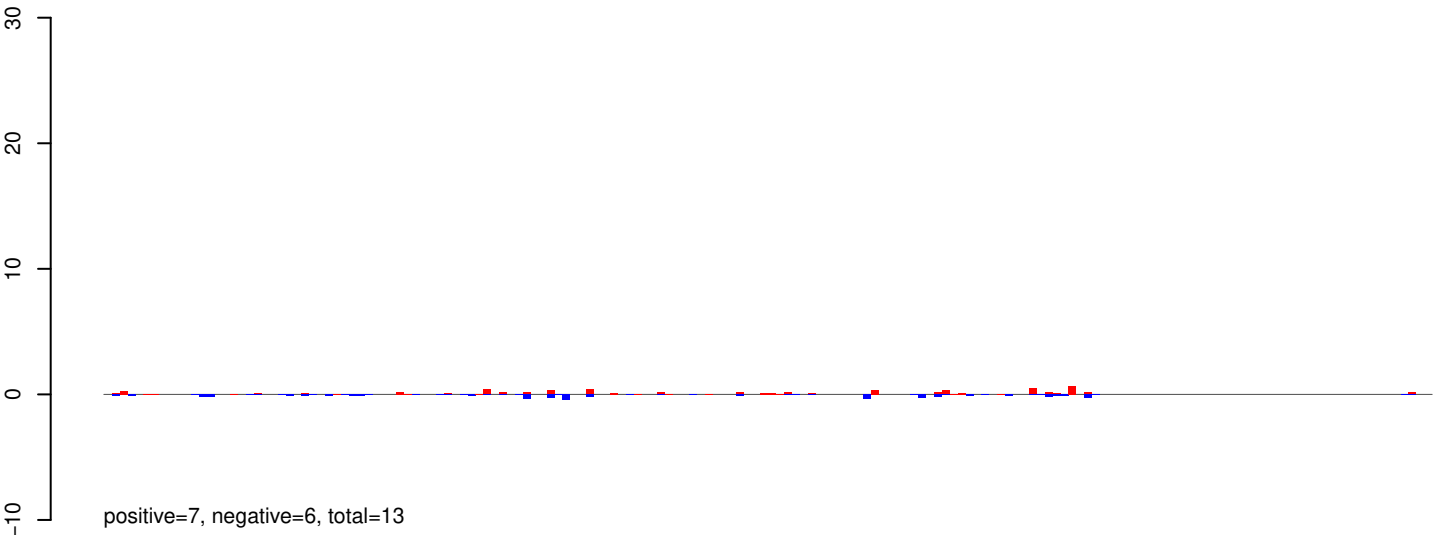
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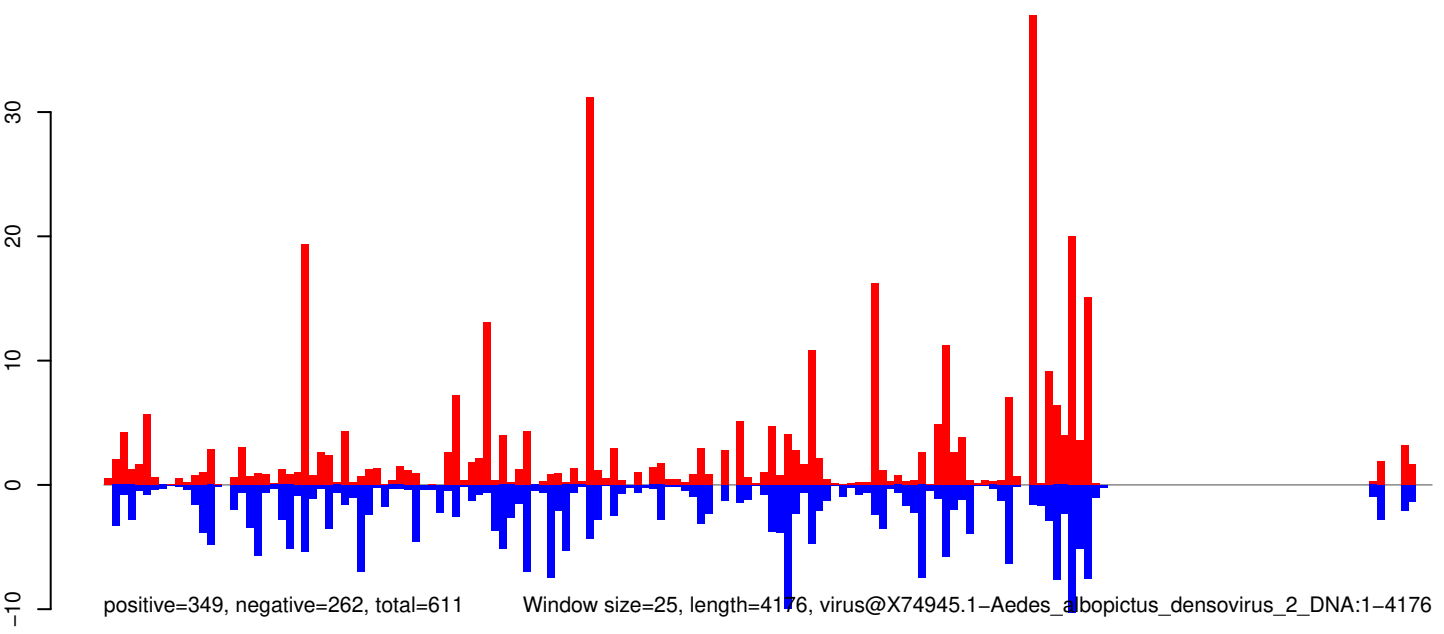
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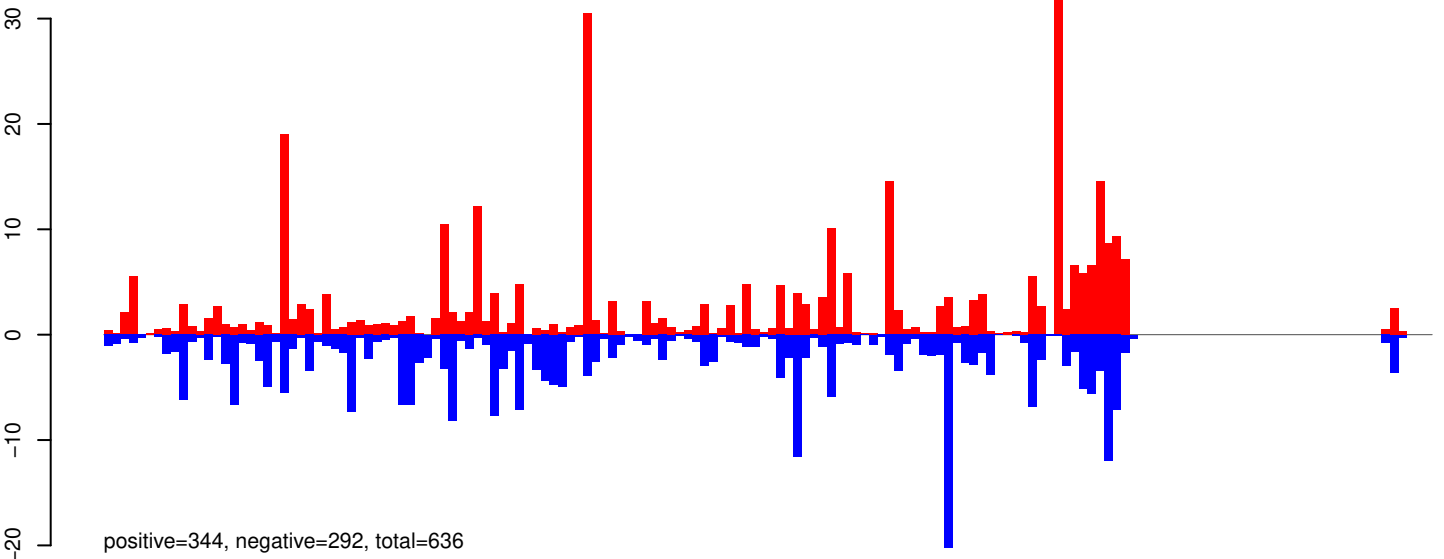
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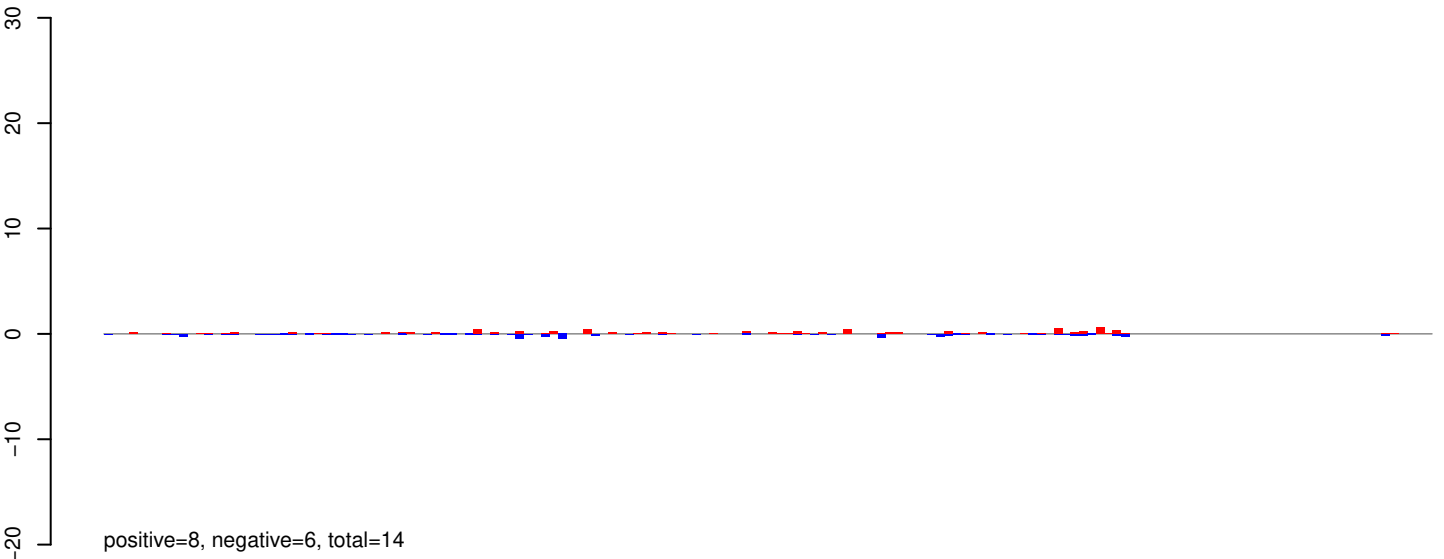
AeAeg_Aag2_BUNV1_ES.rep



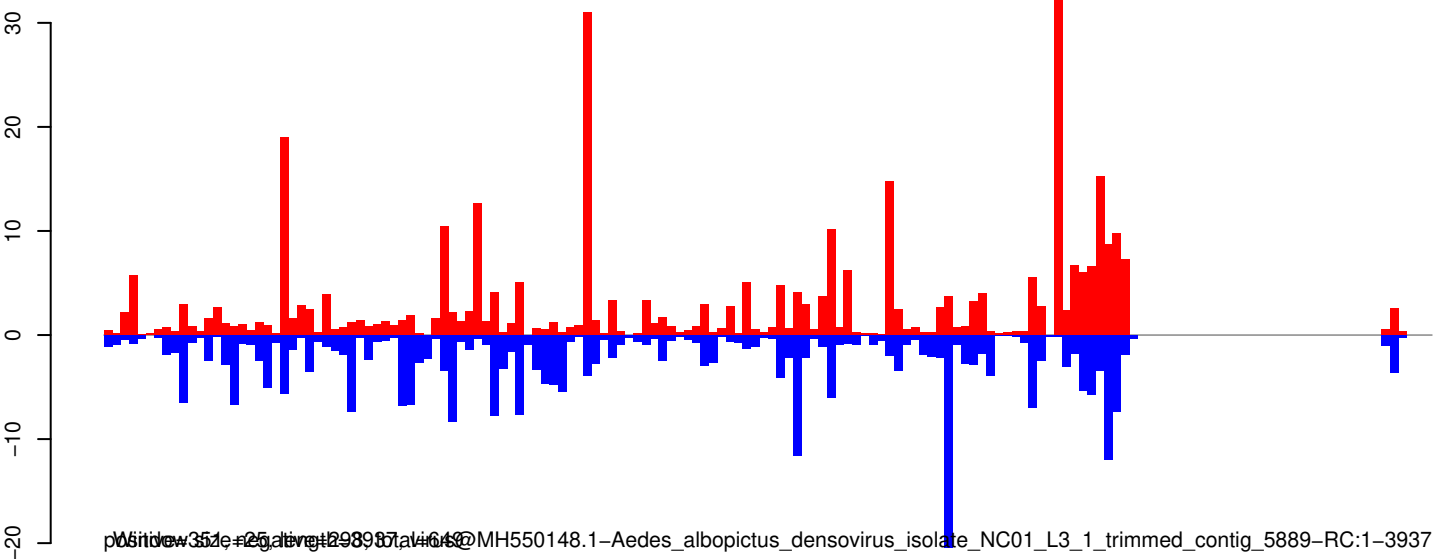
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



AeAeg_Aag2_BUNV1_ES.rep

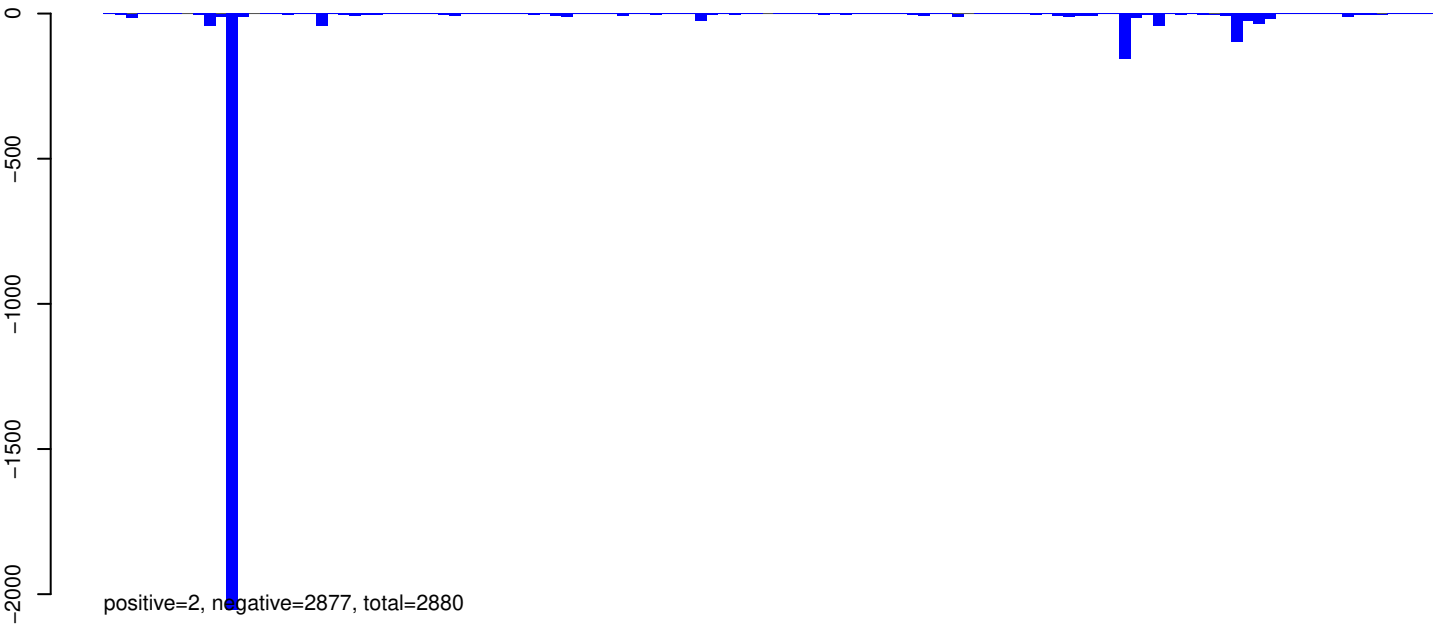


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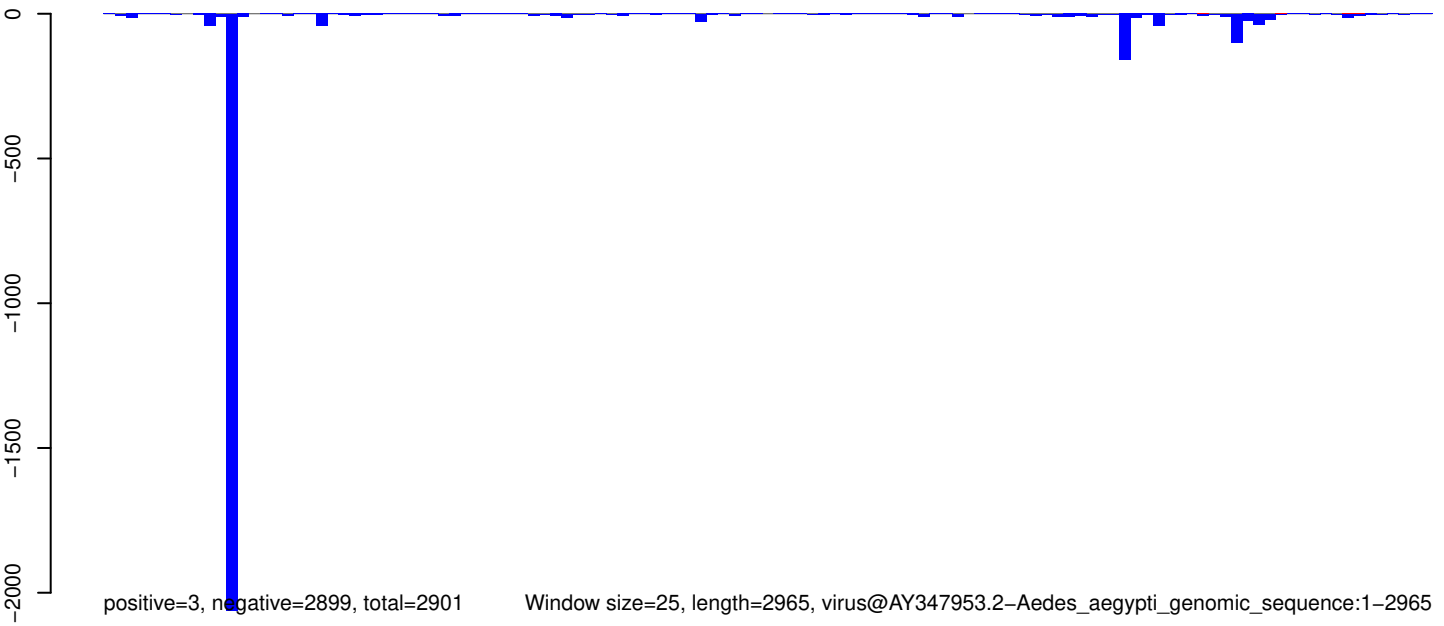
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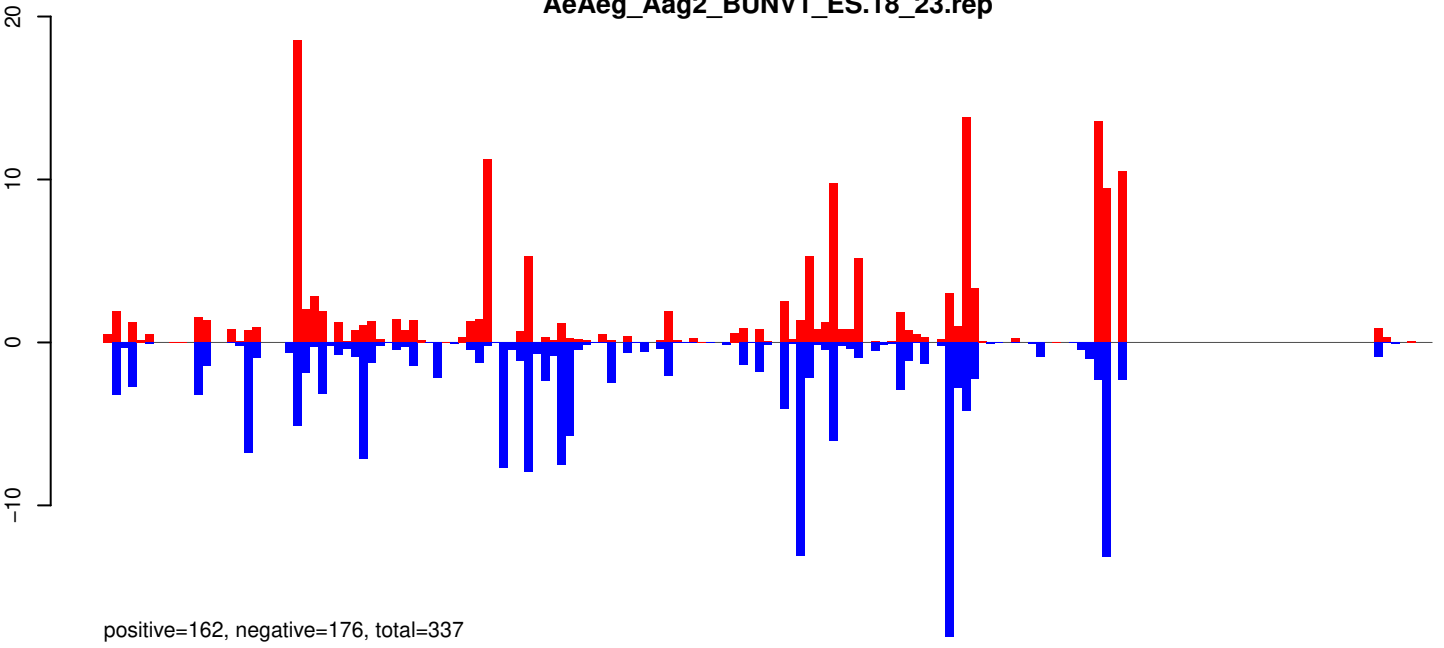
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AeAeg_Aag2_BUNV1_ES.rep

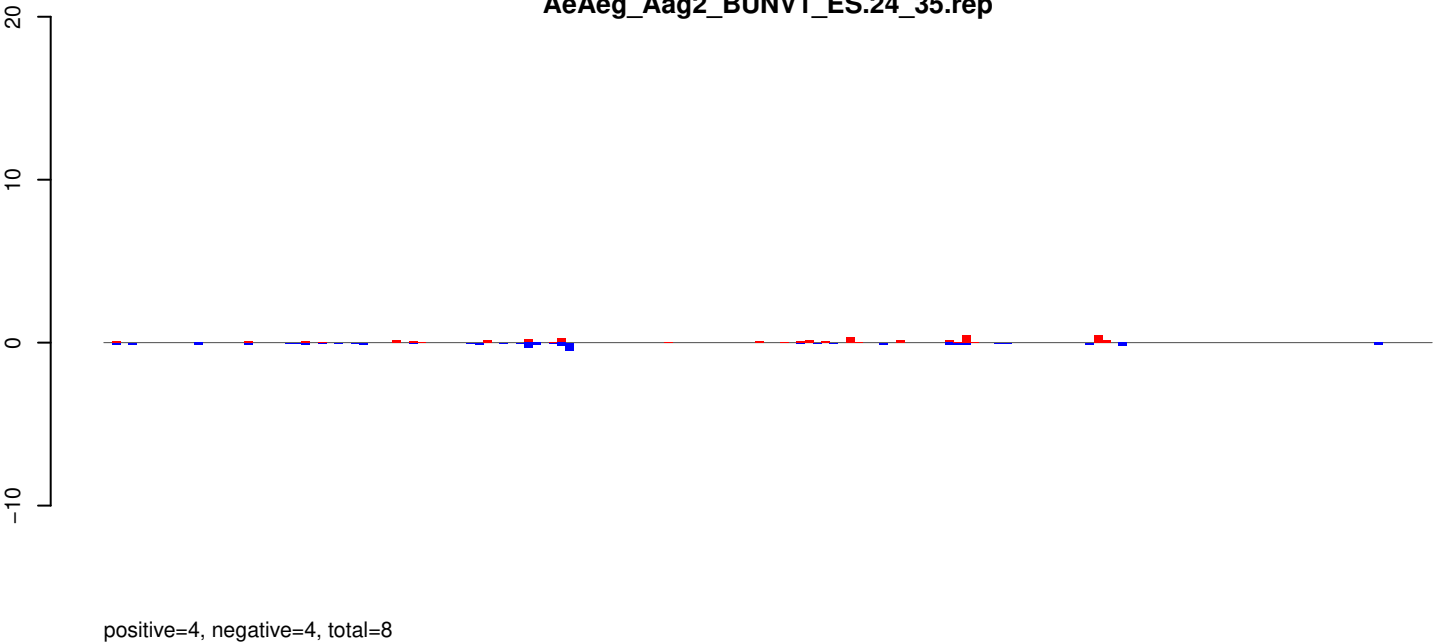


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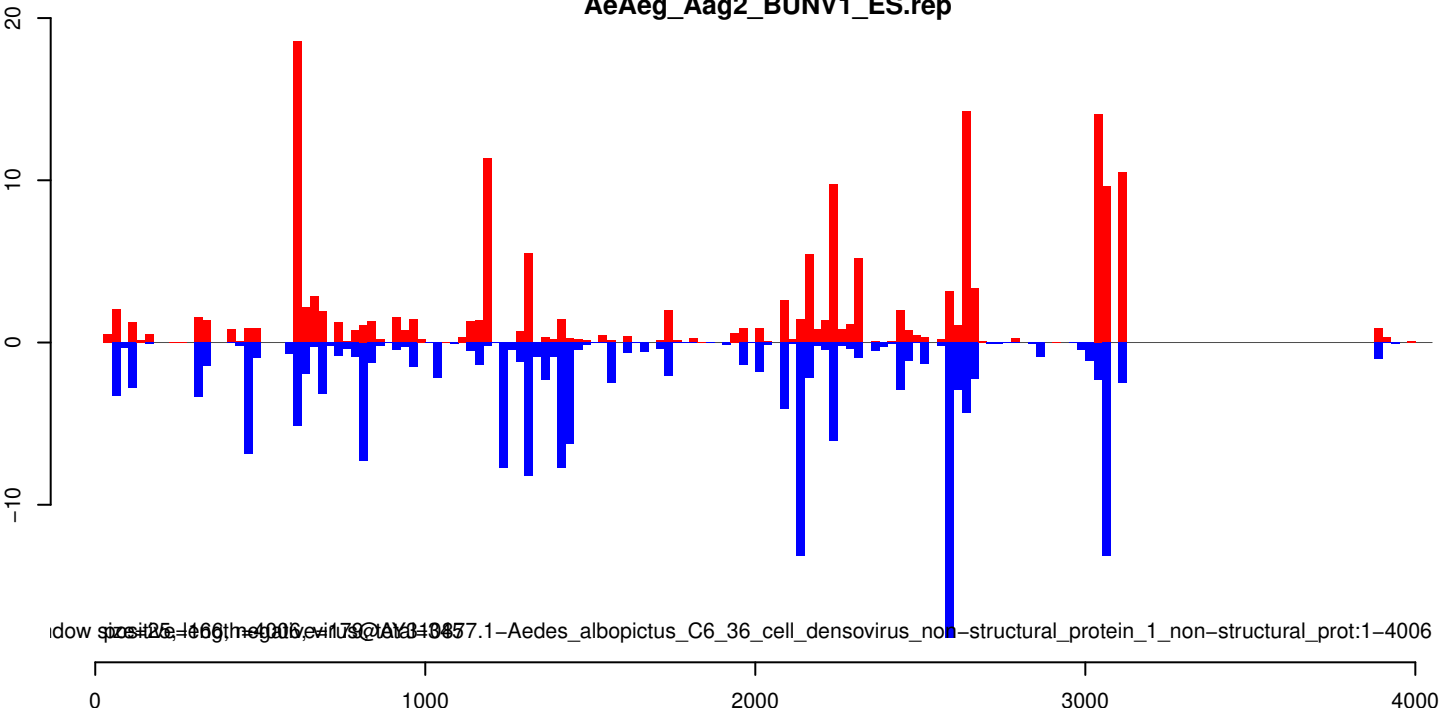
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AeAeg_Aag2_BUNV1_ES.24_35.rep

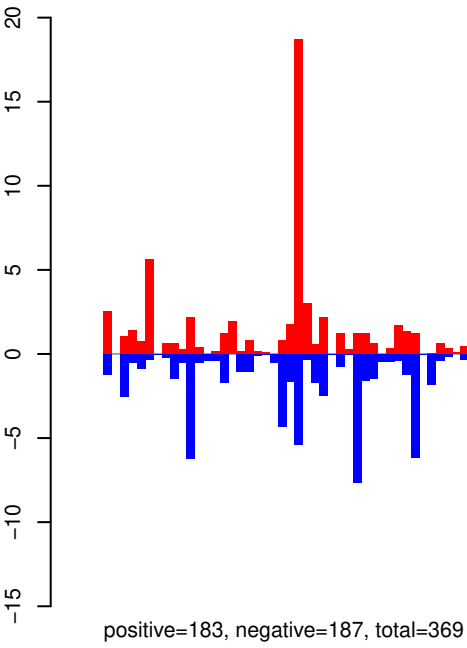


positive=4, negative=4, total=8

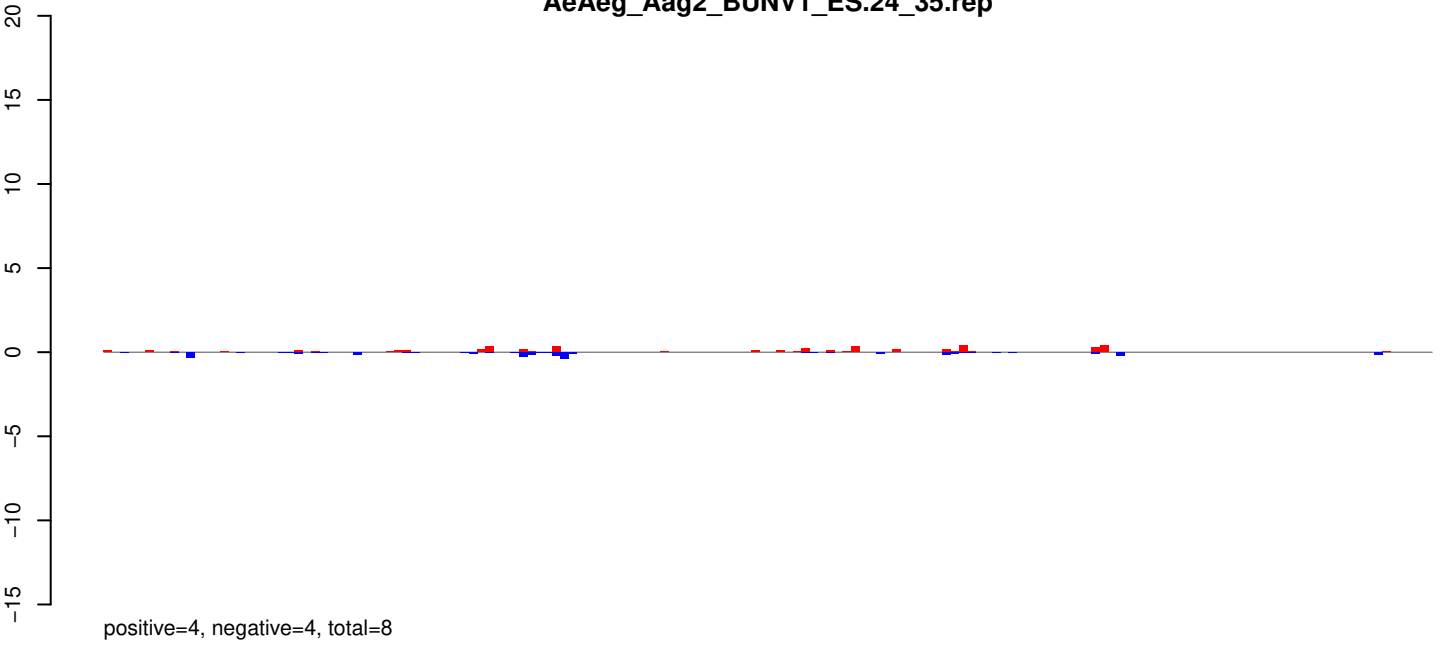
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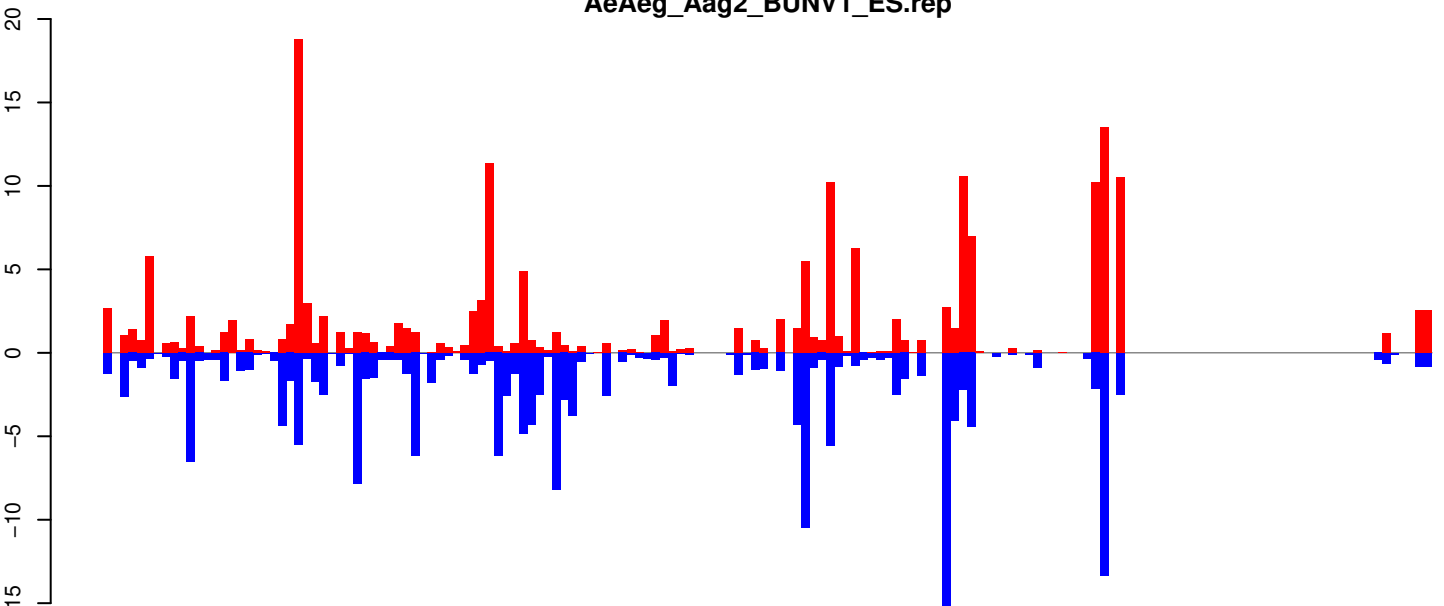
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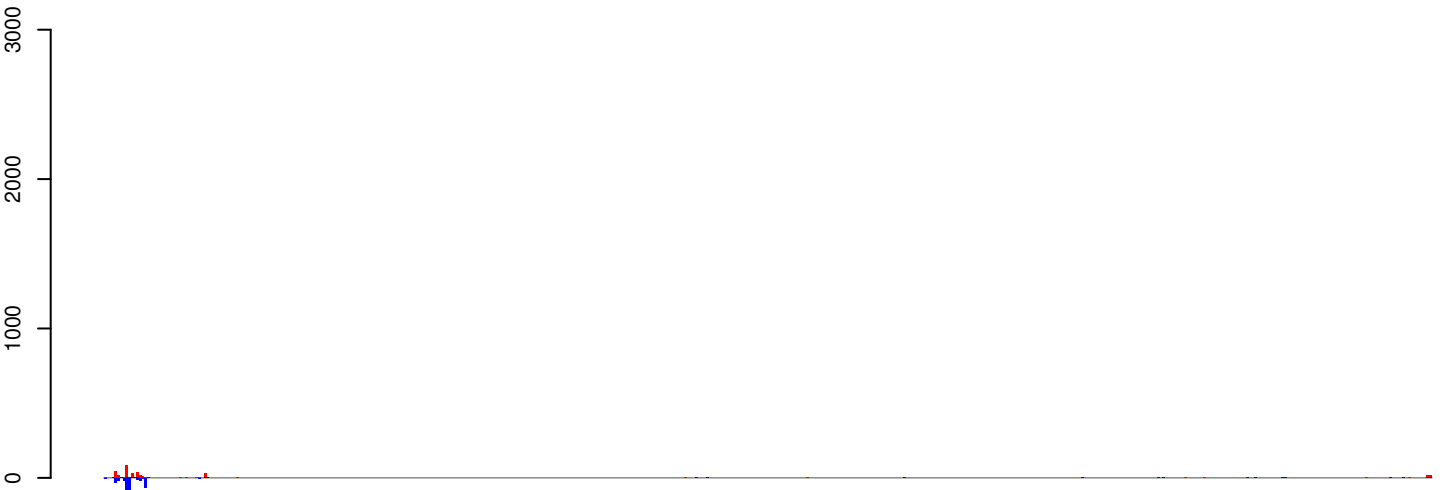
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AeAeg_Aag2_BUNV1_ES.rep

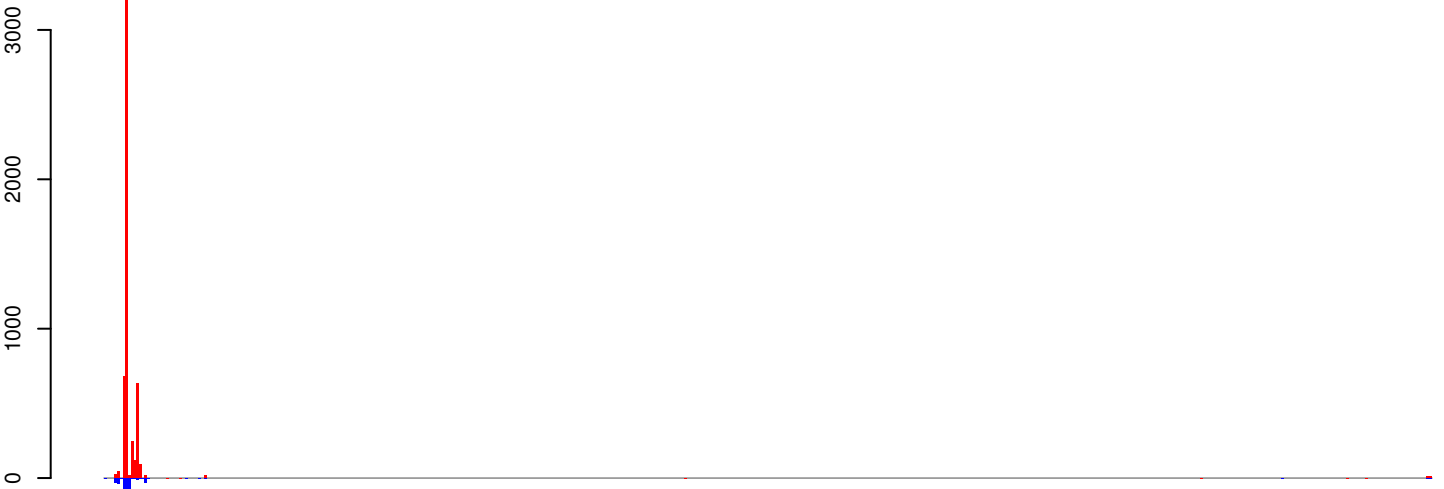


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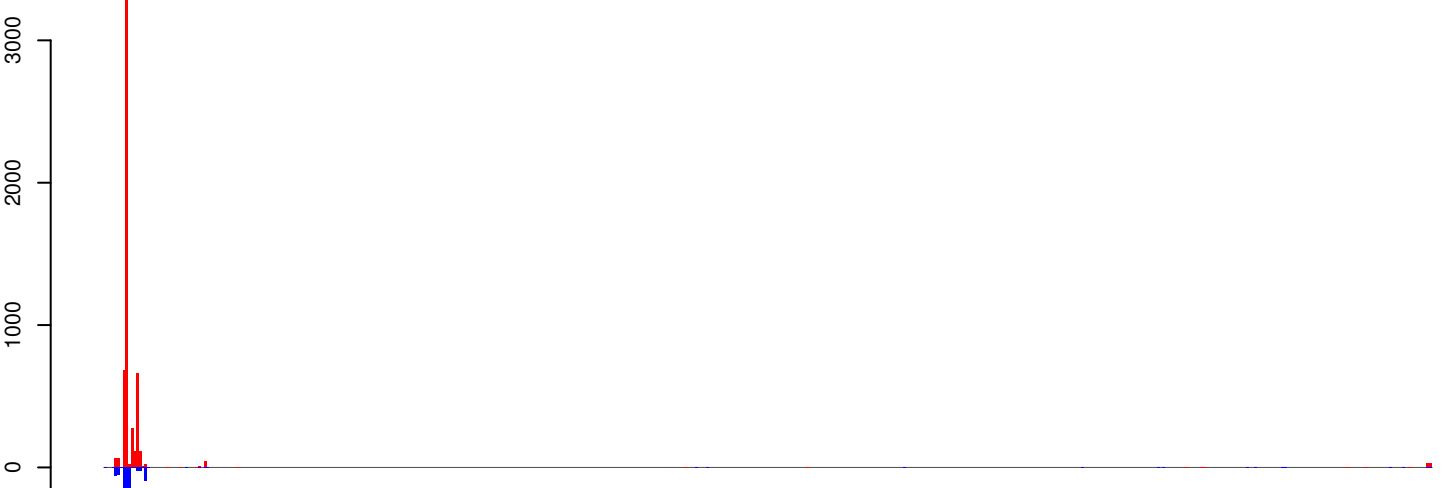
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AeAeg_Aag2_BUNV1_ES.24_35.rep

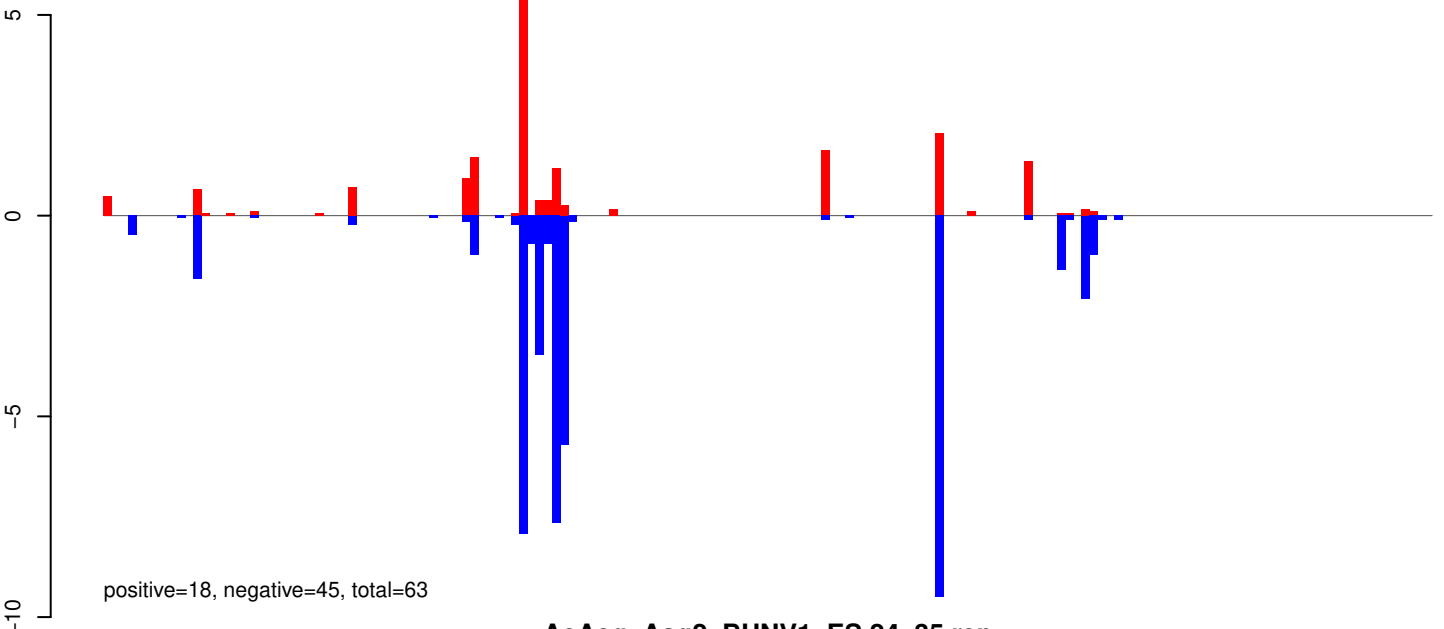


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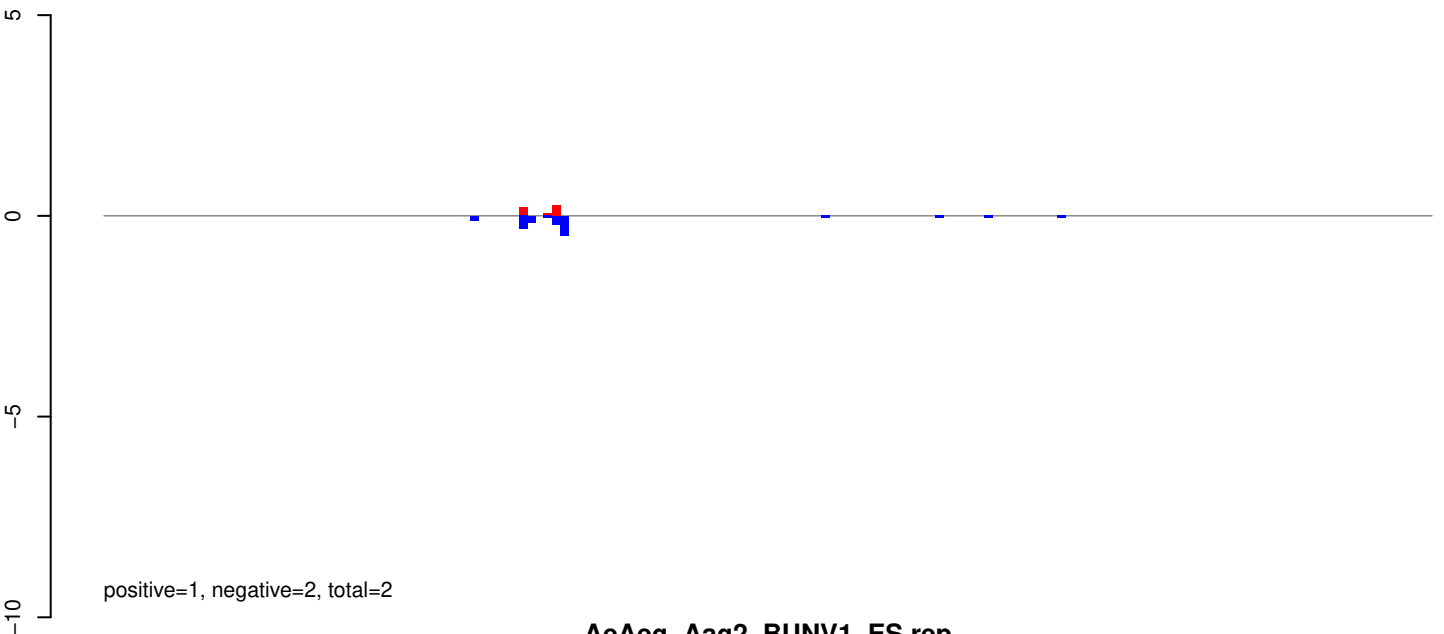
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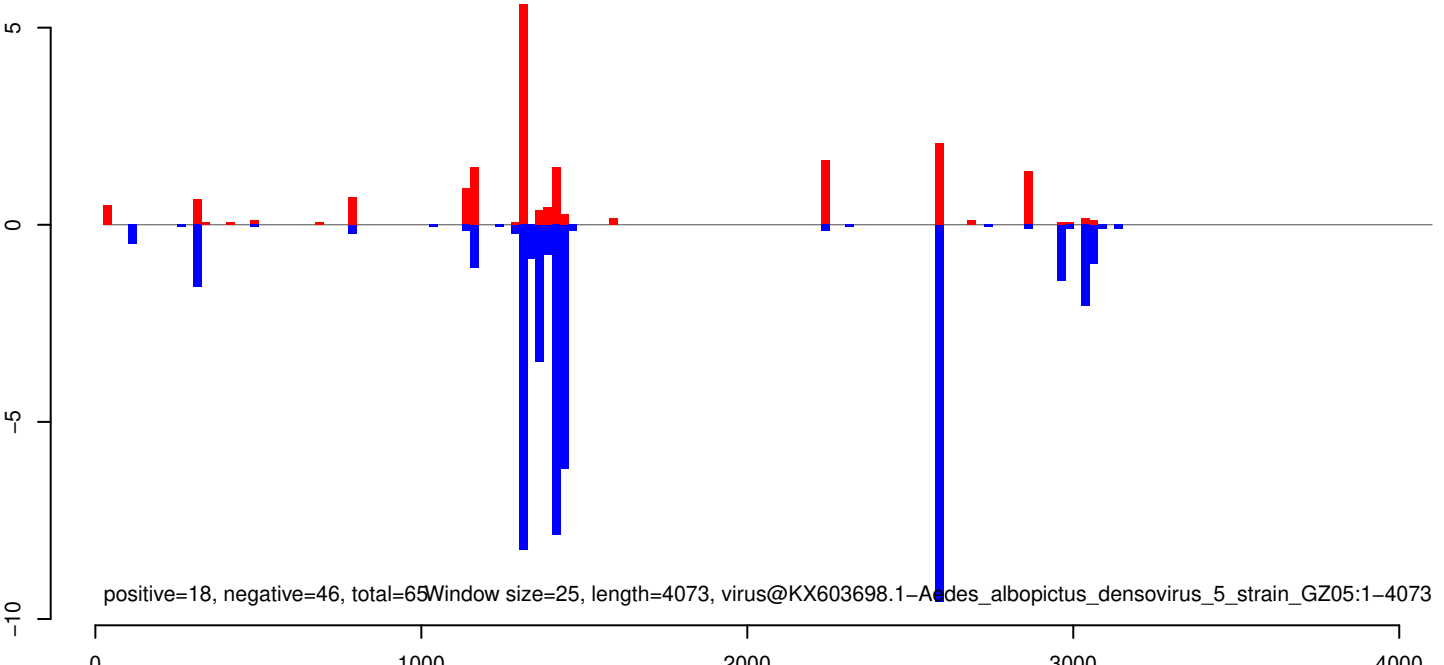
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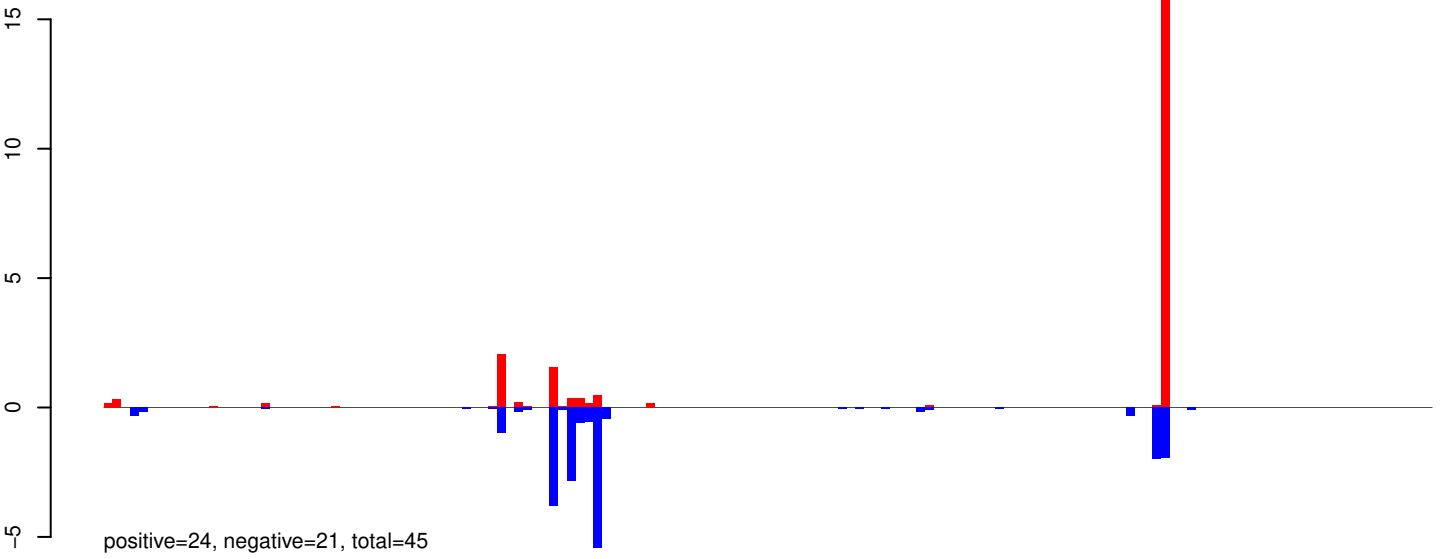
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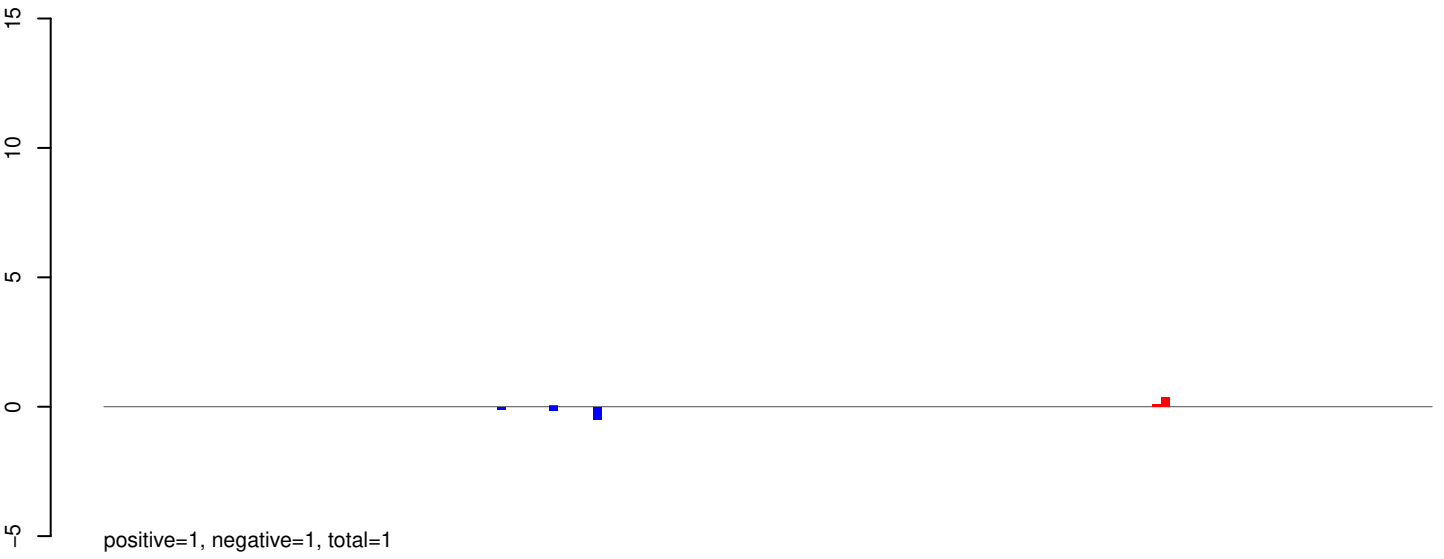
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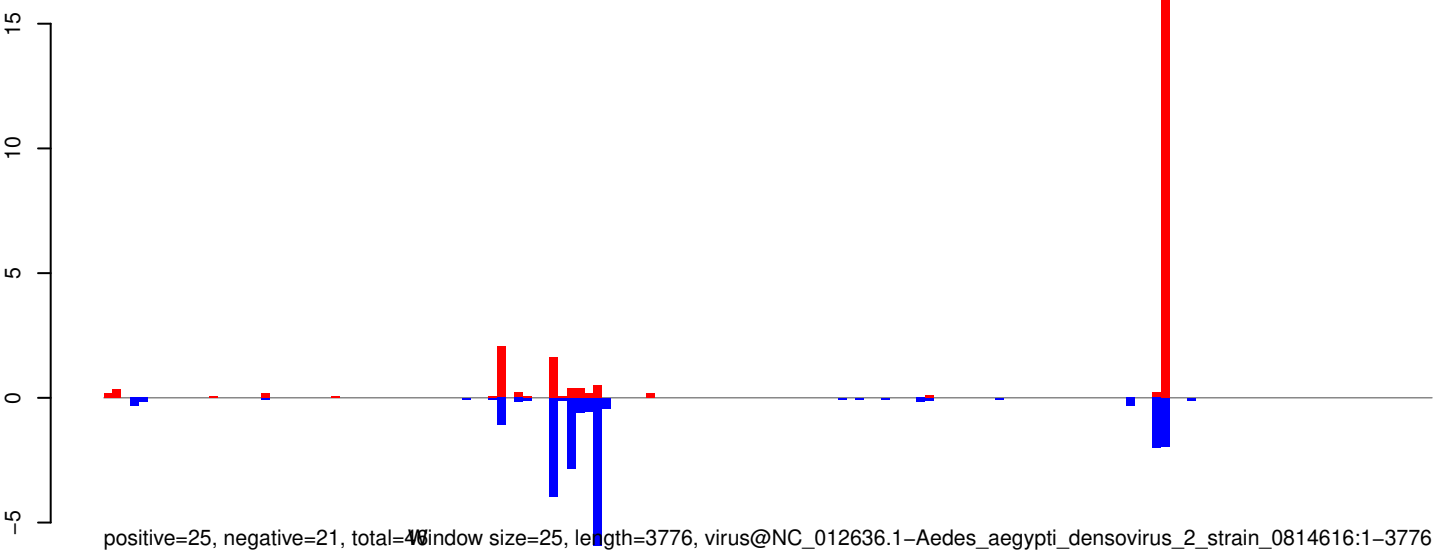
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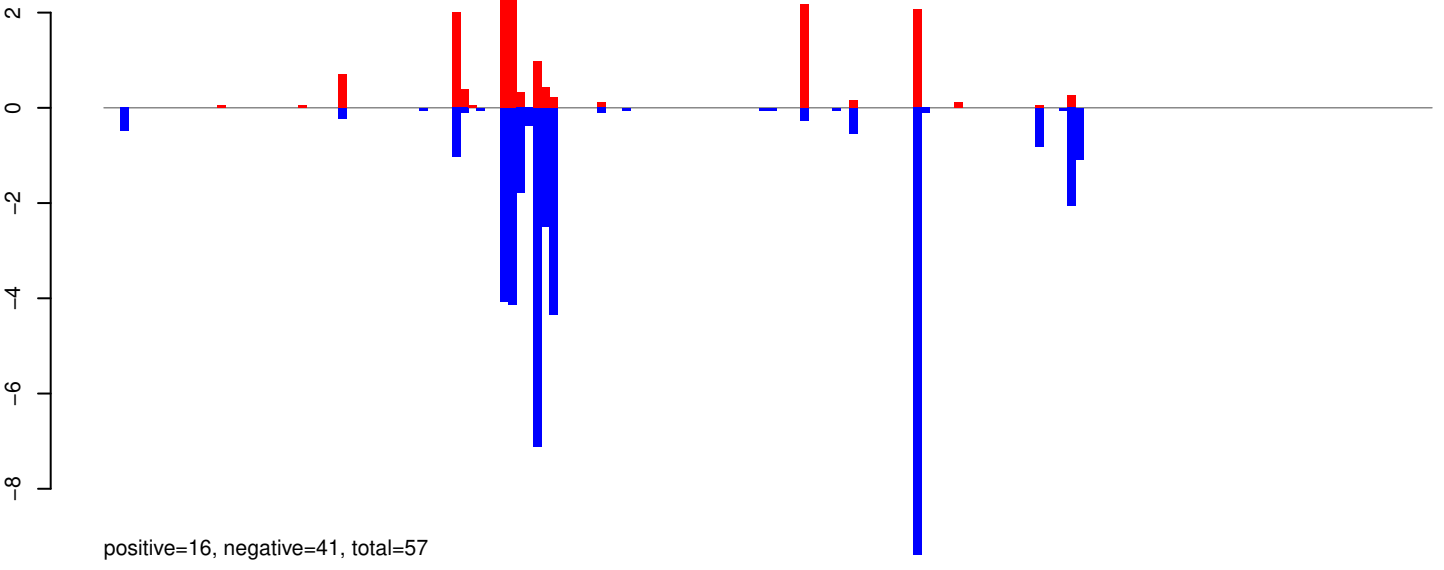
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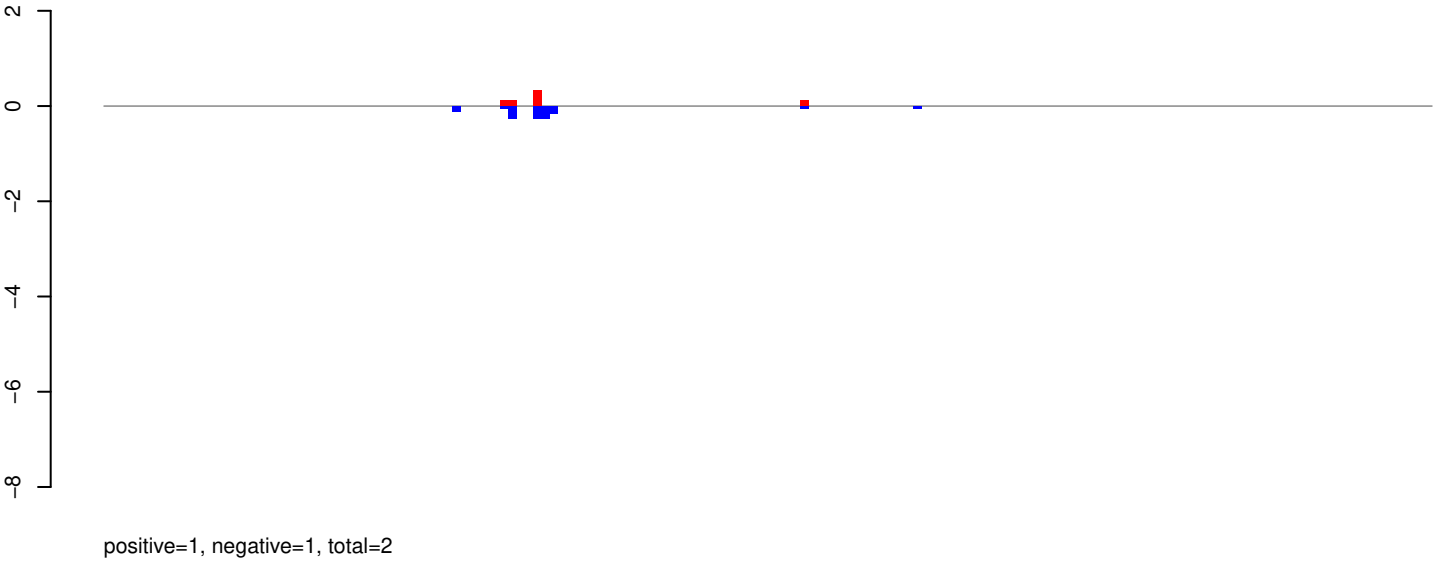
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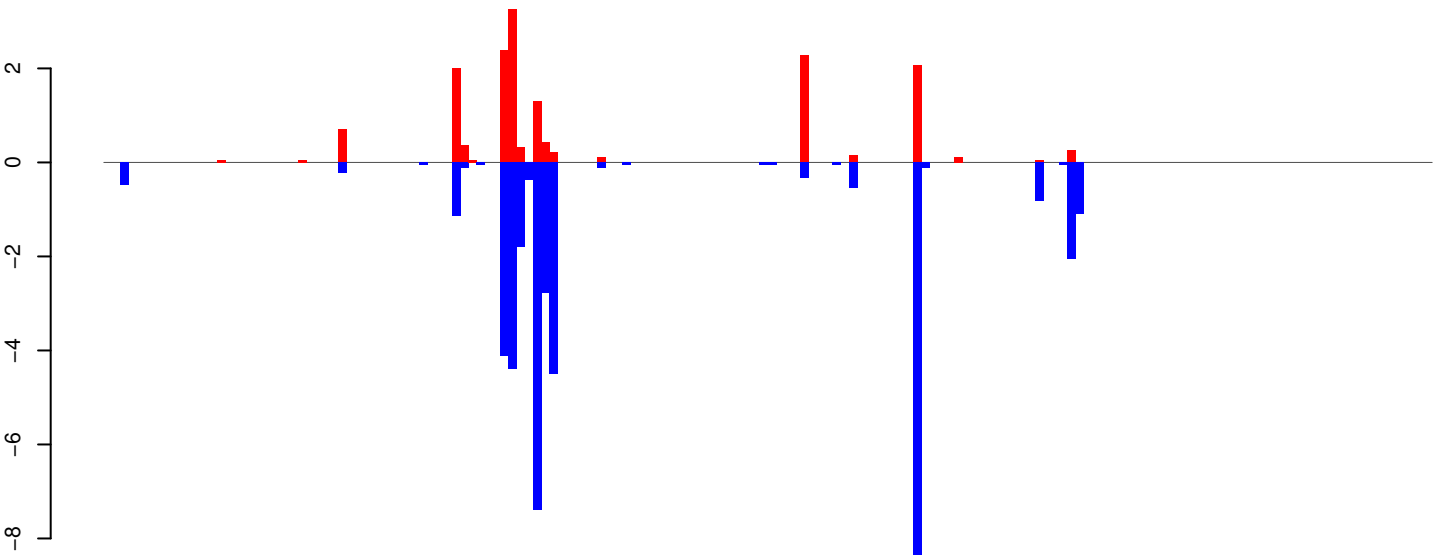
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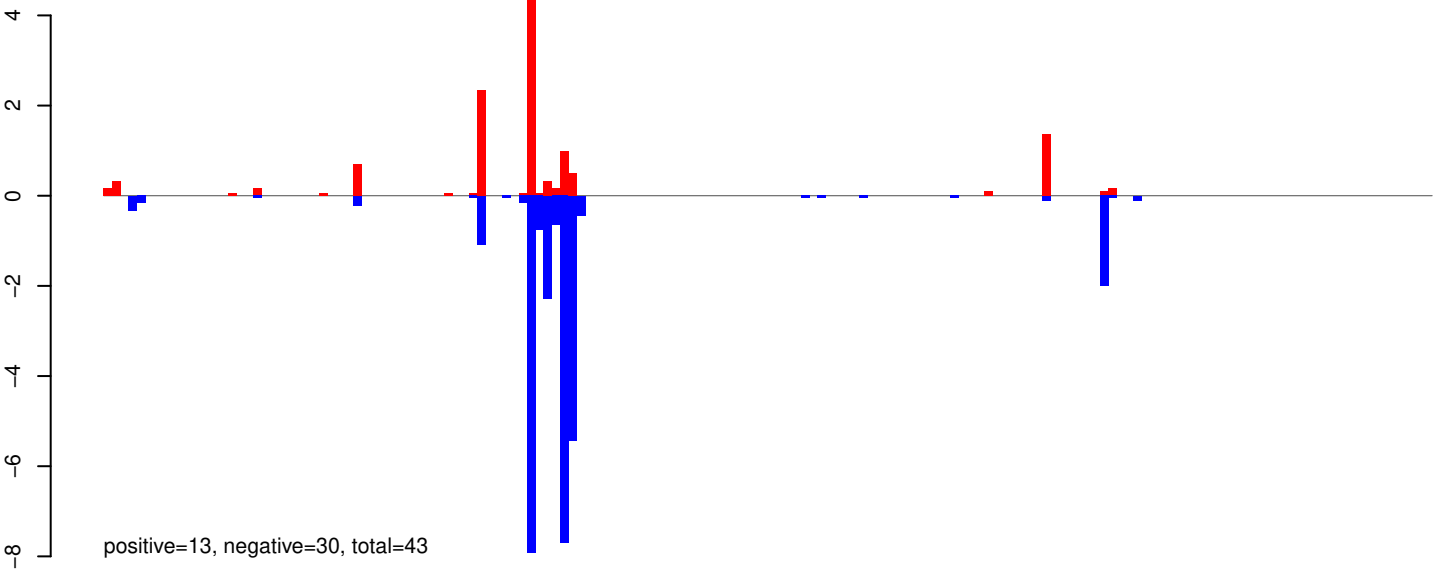
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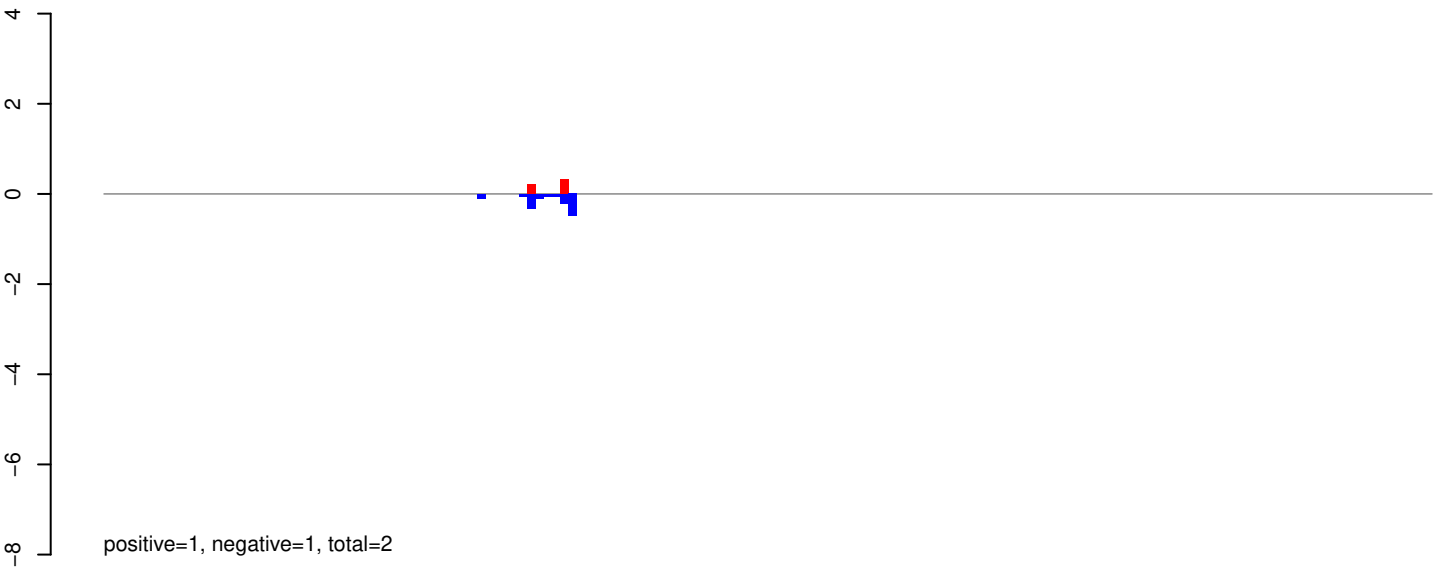
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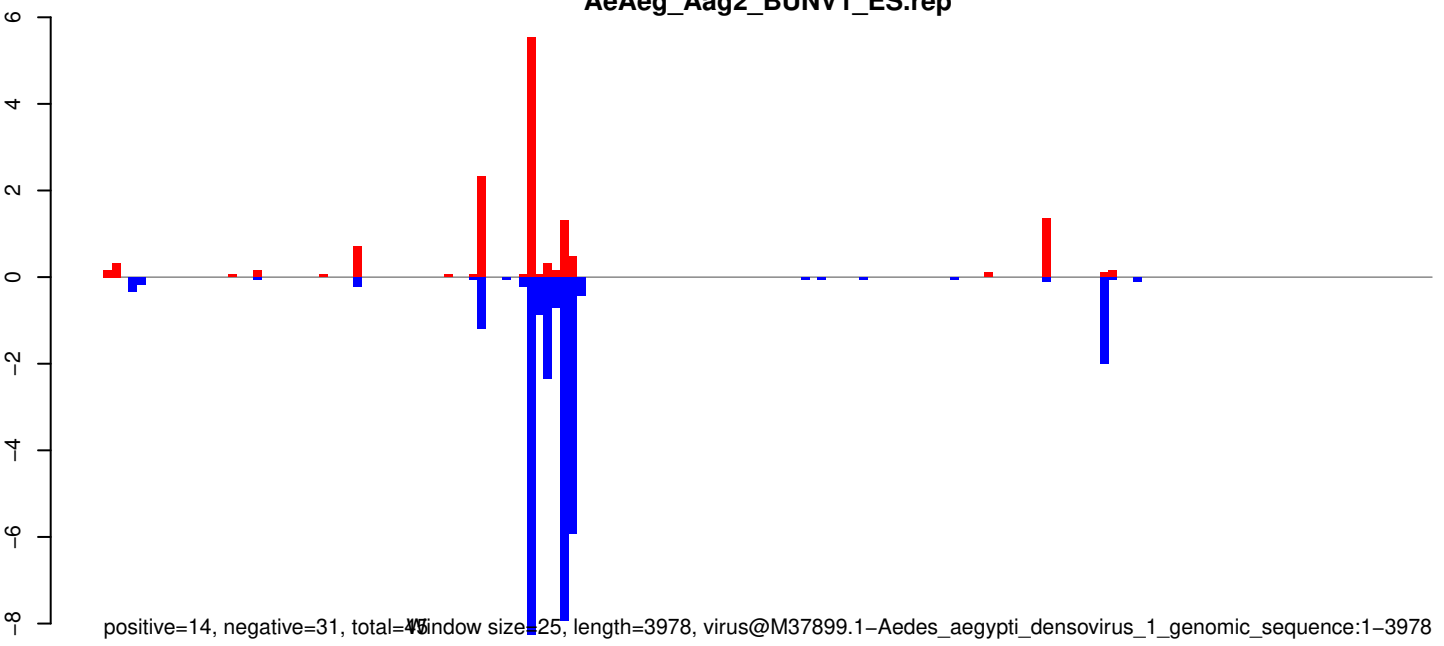
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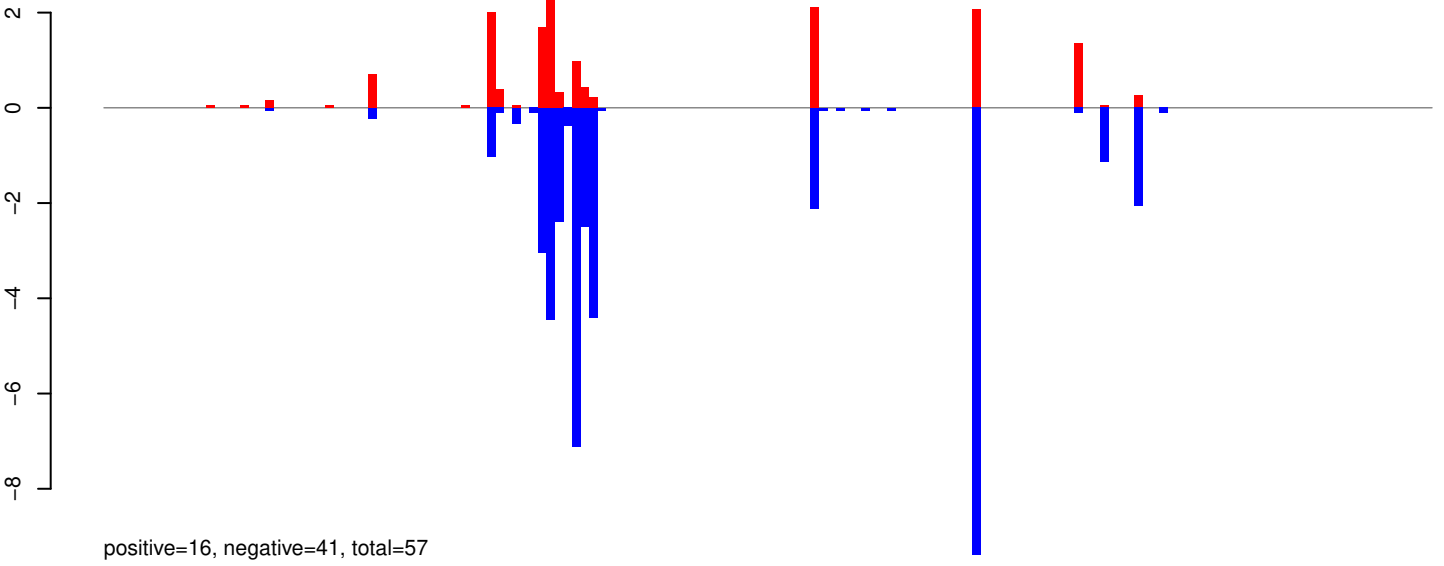
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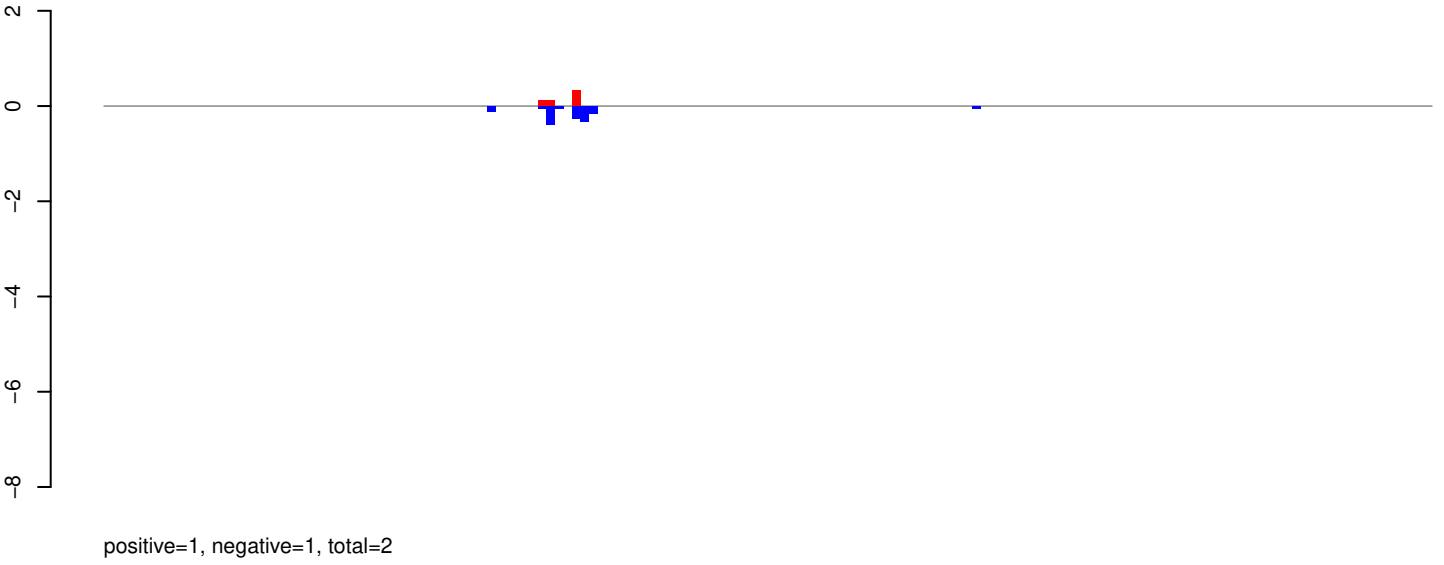
AeAeg_Aag2_BUNV1_ES.rep



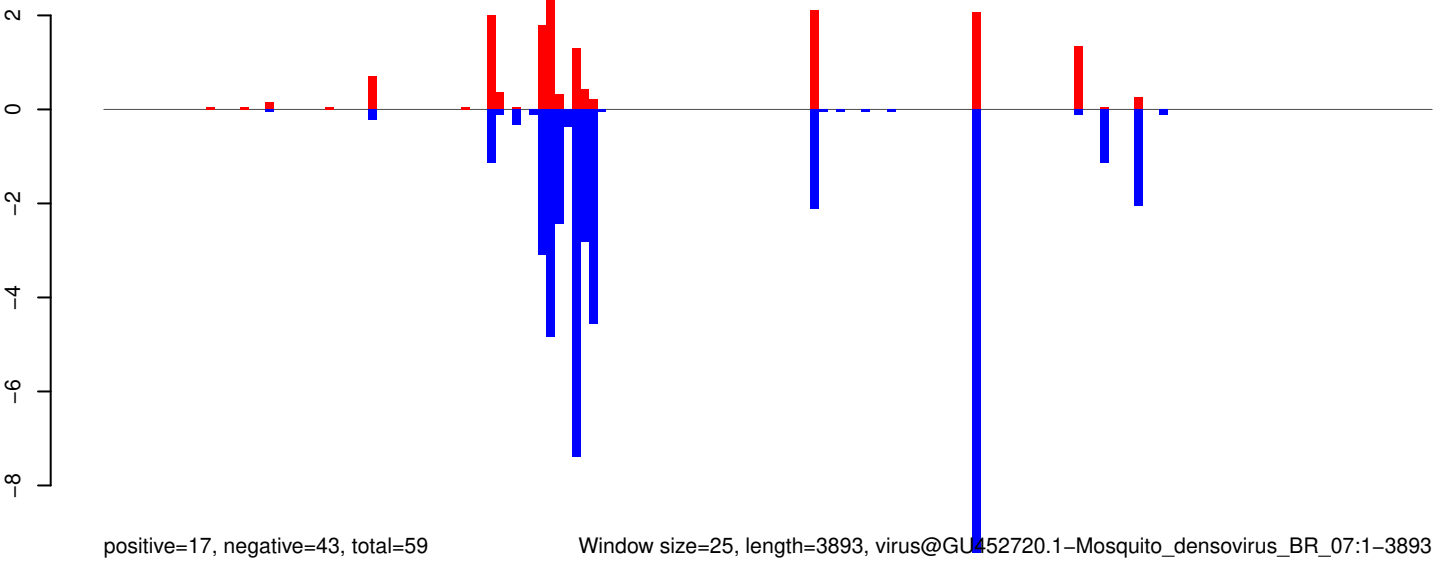
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AeAeg_Aag2_BUNV1_ES.24_35.rep



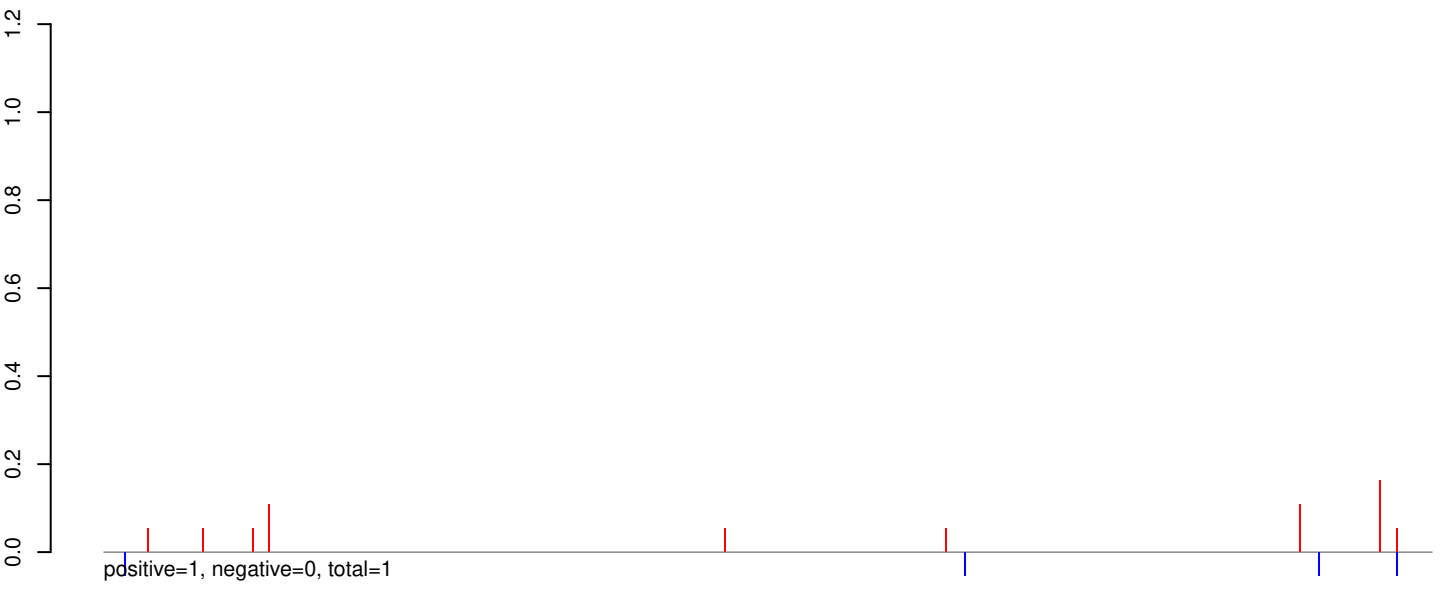
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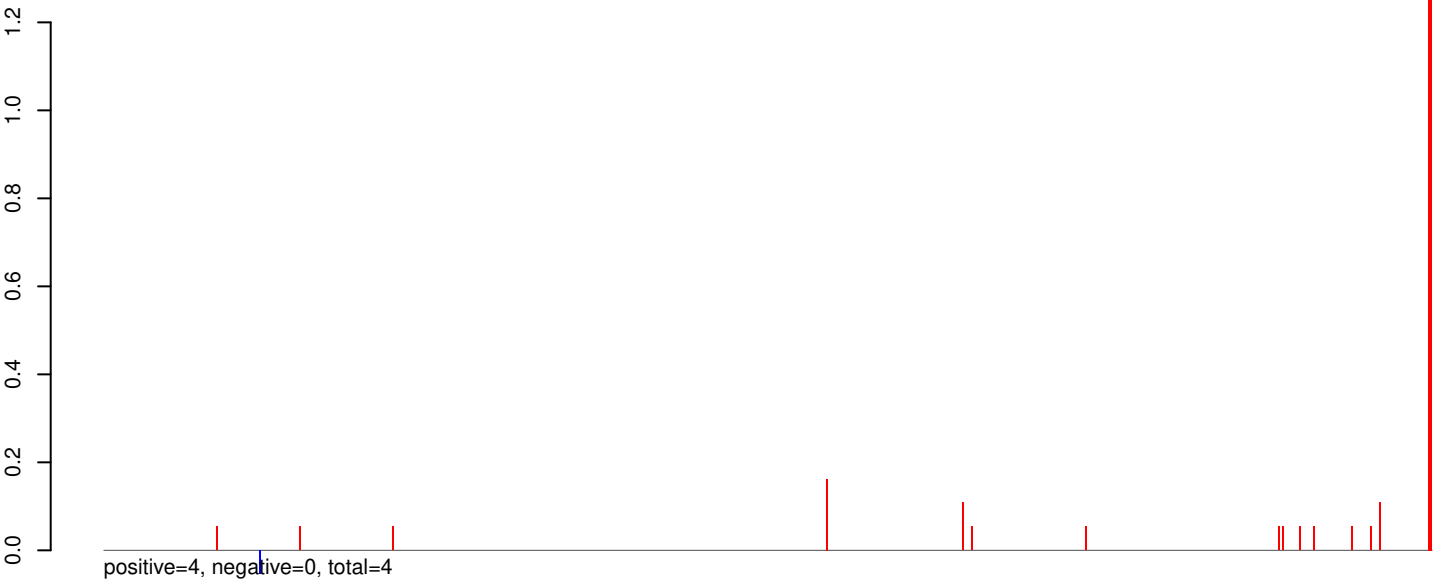
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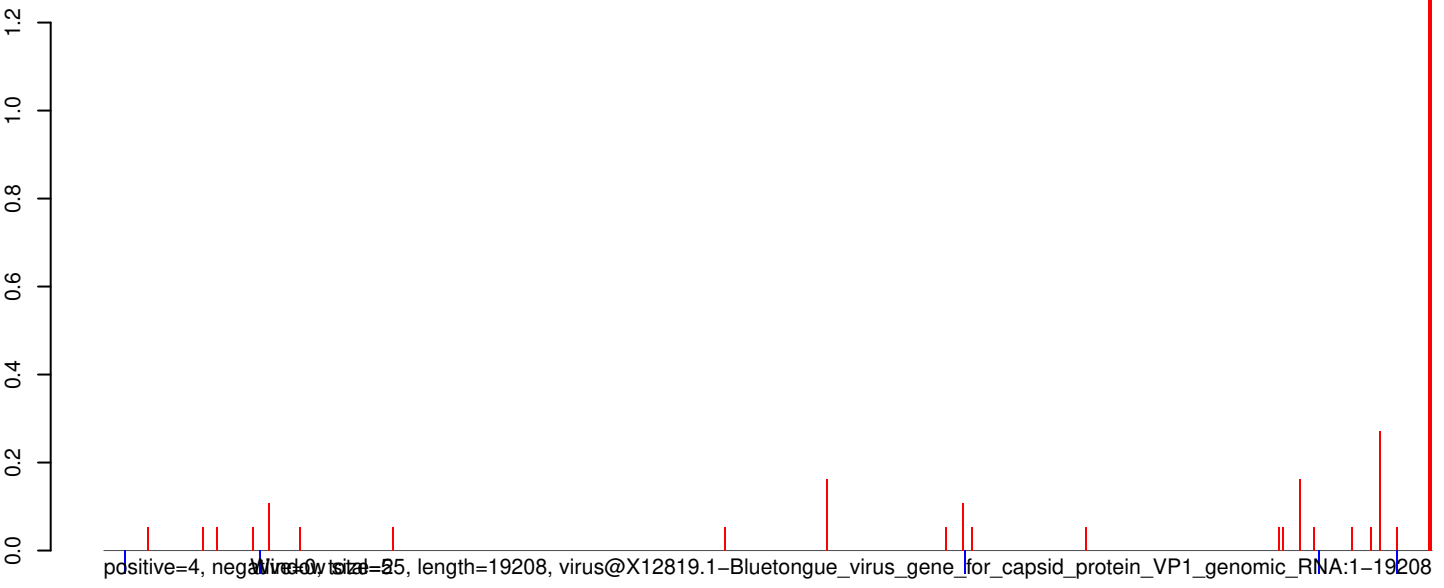
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AeAeg_Aag2_BUNV1_ES.24_35.rep

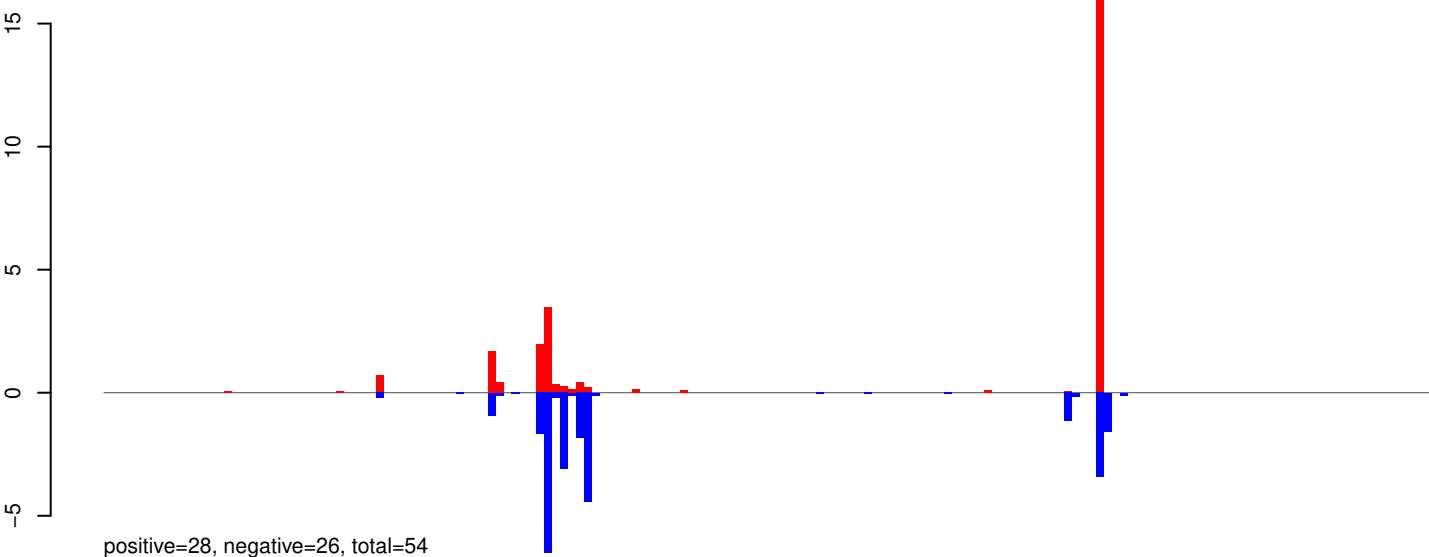


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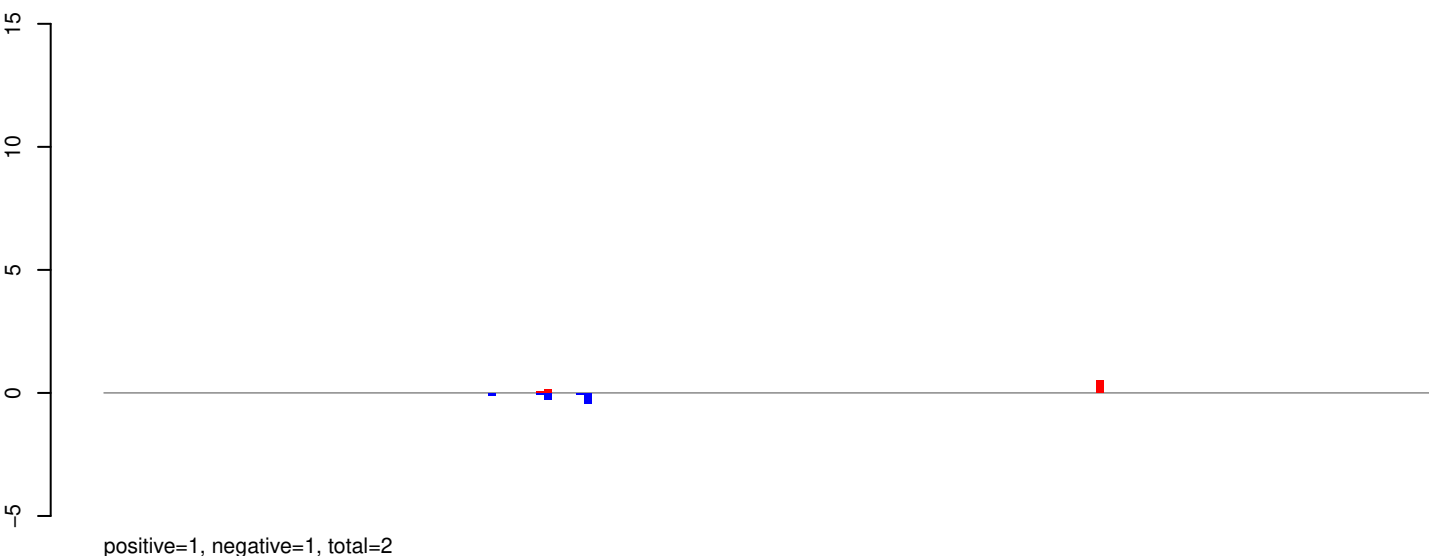


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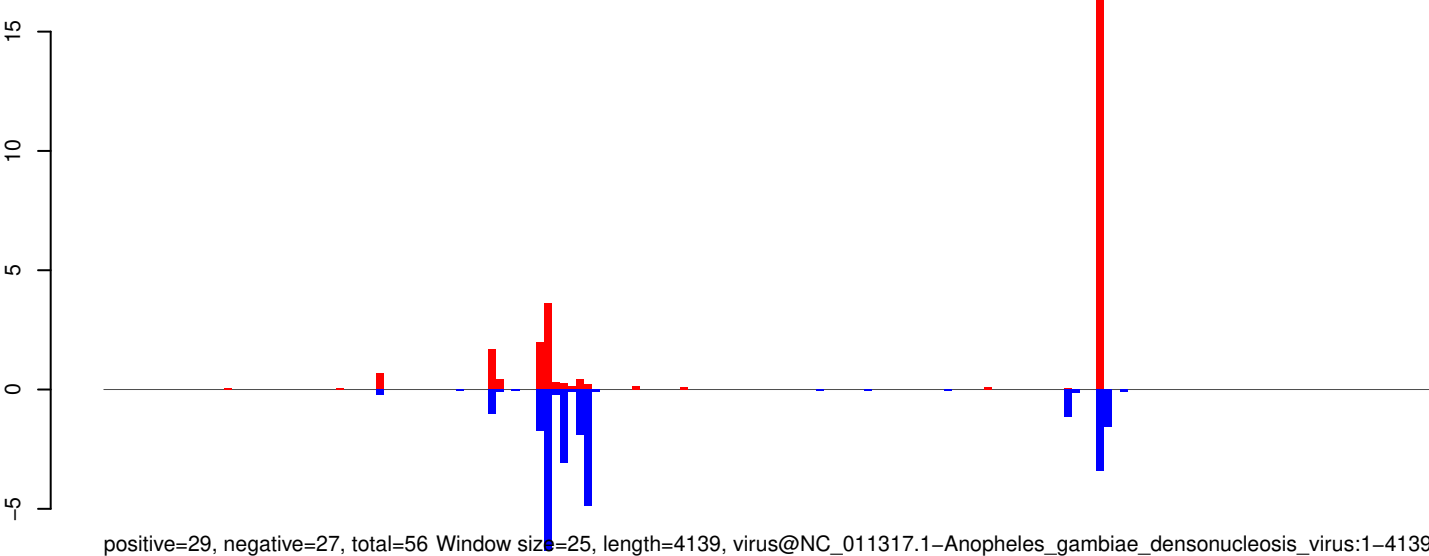
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep

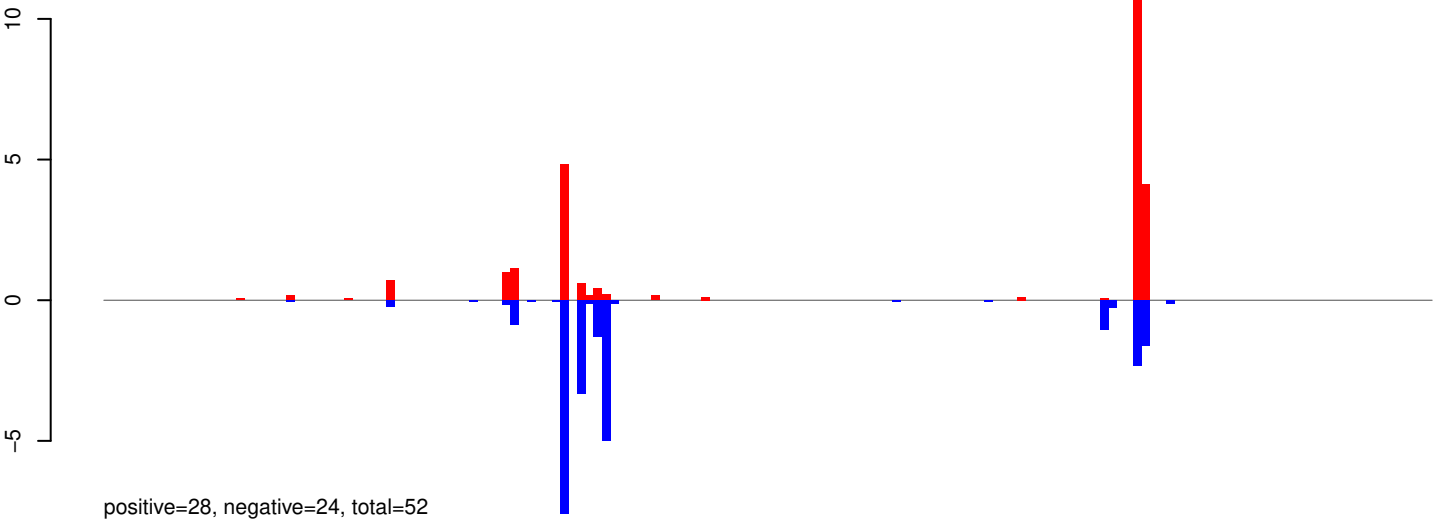


AeAeg_Aag2_BUNV1_ES.rep

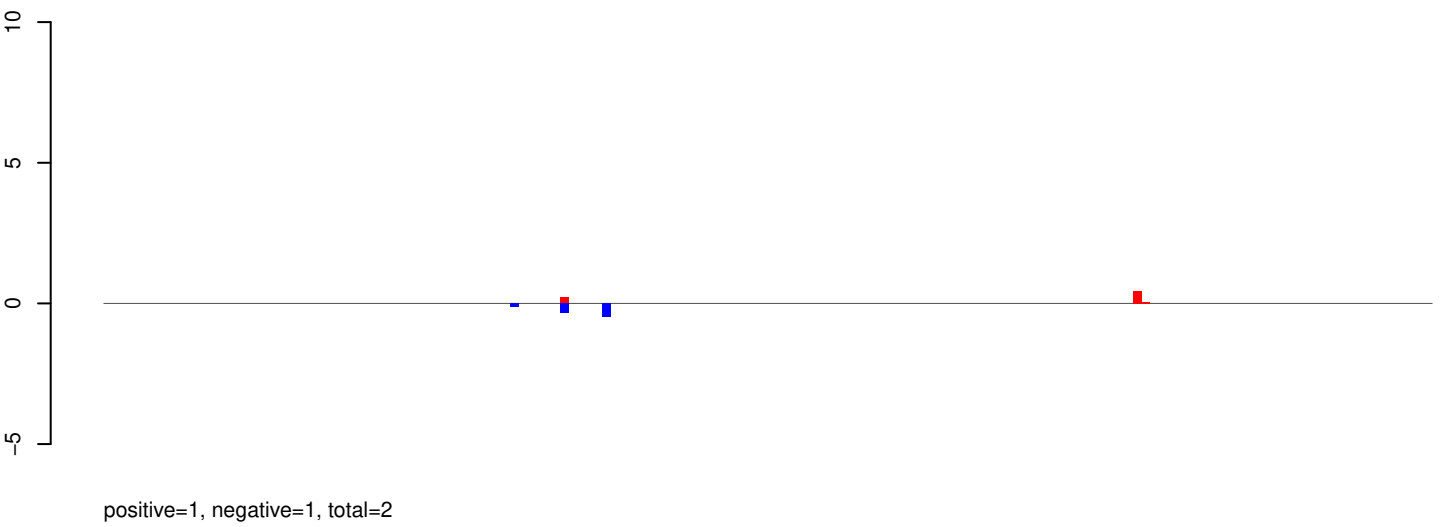


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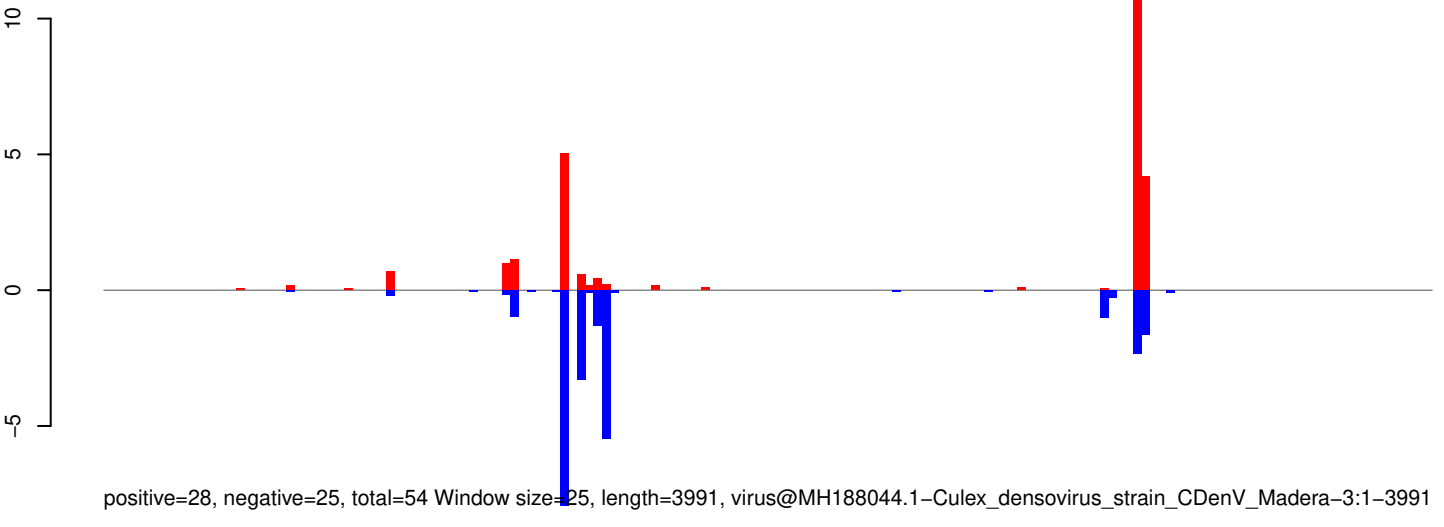
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AeAeg_Aag2_BUNV1_ES.24_35.rep

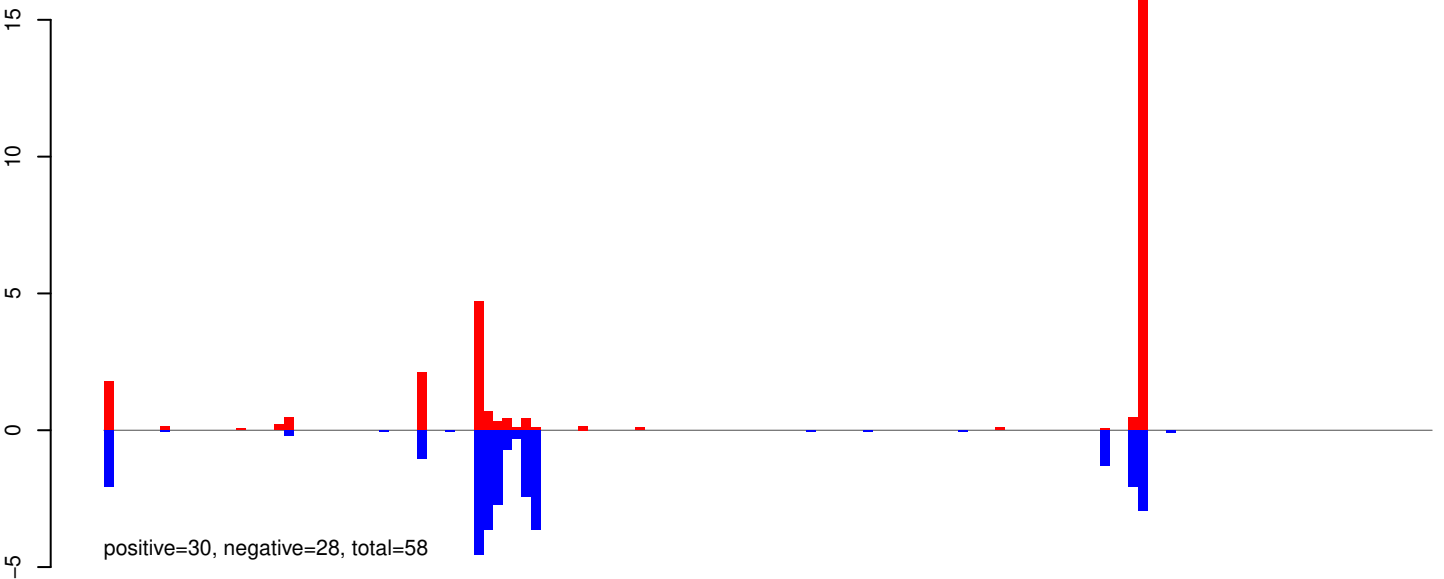


AeAeg_Aag2_BUNV1_ES.rep

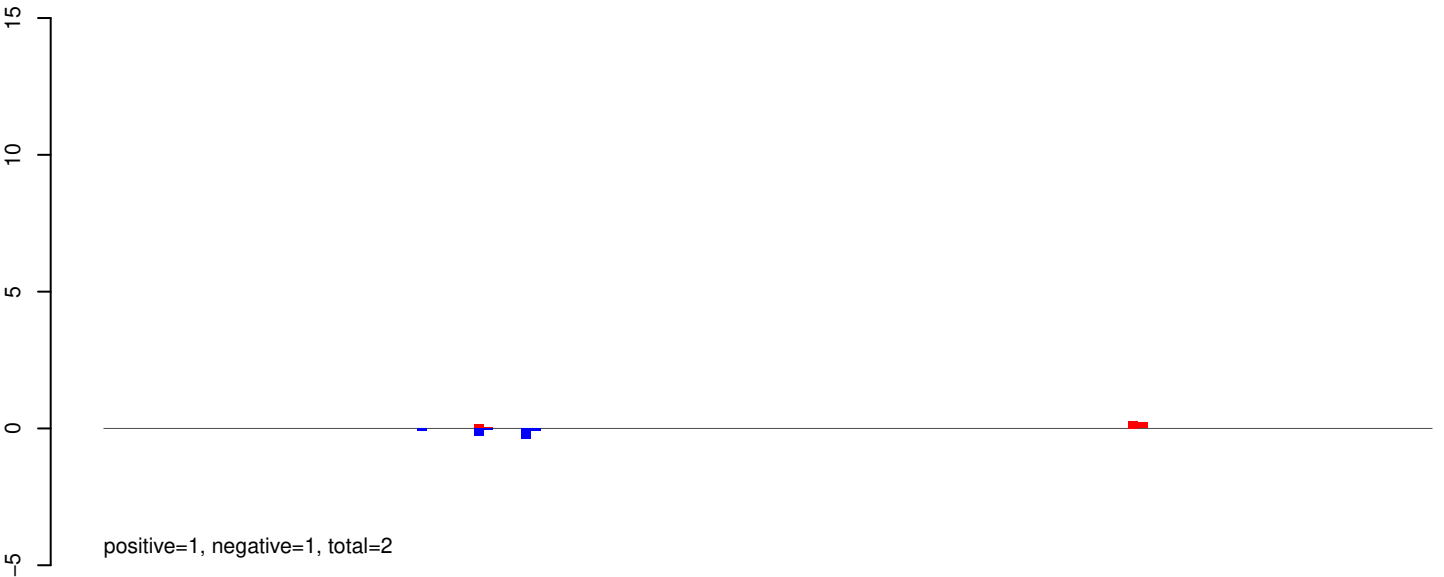


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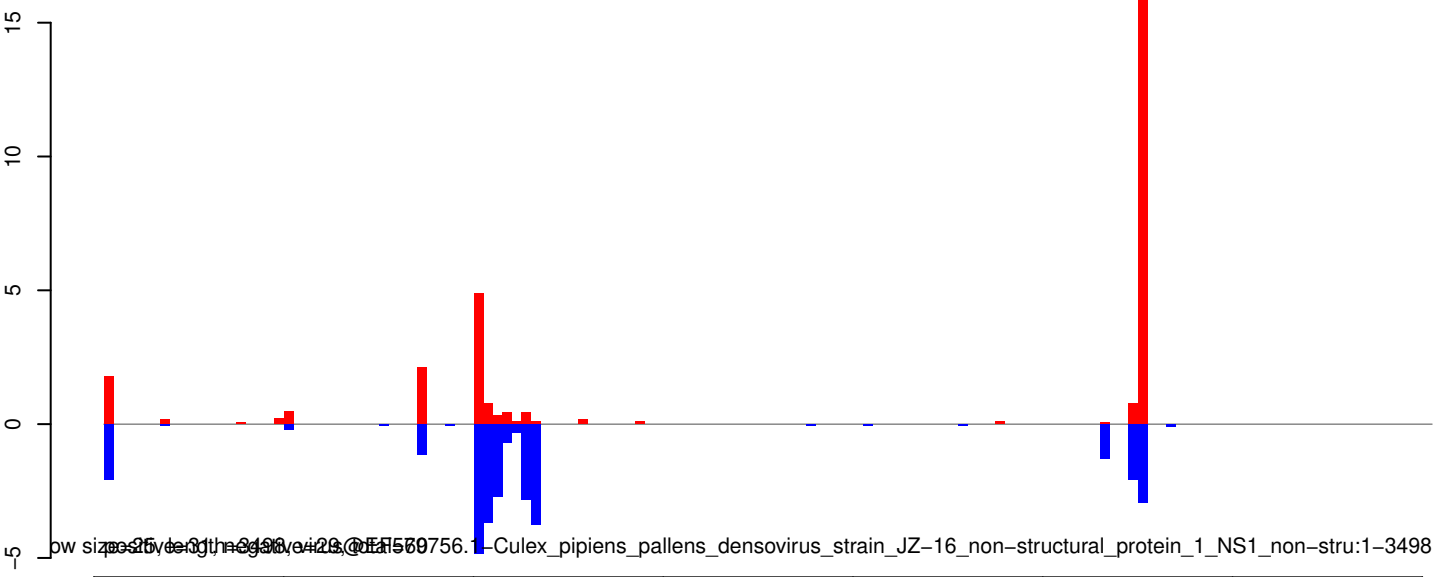
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AeAeg_Aag2_BUNV1_ES.24_35.rep

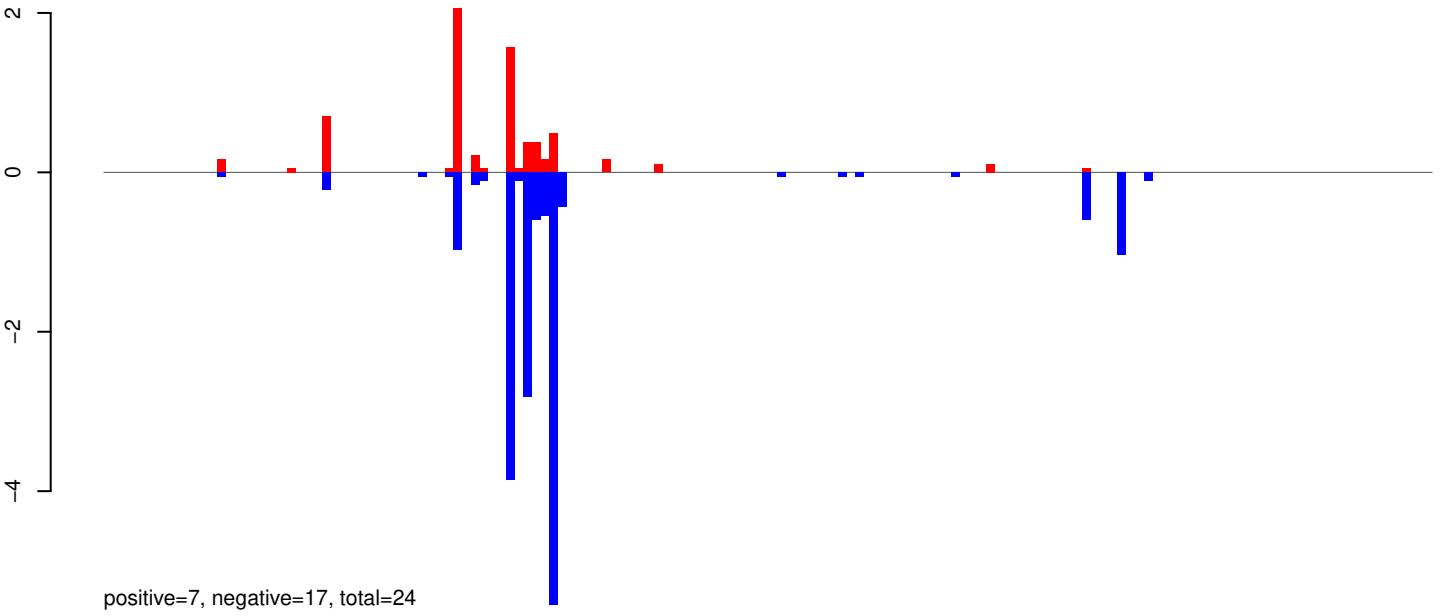


AeAeg_Aag2_BUNV1_ES.rep



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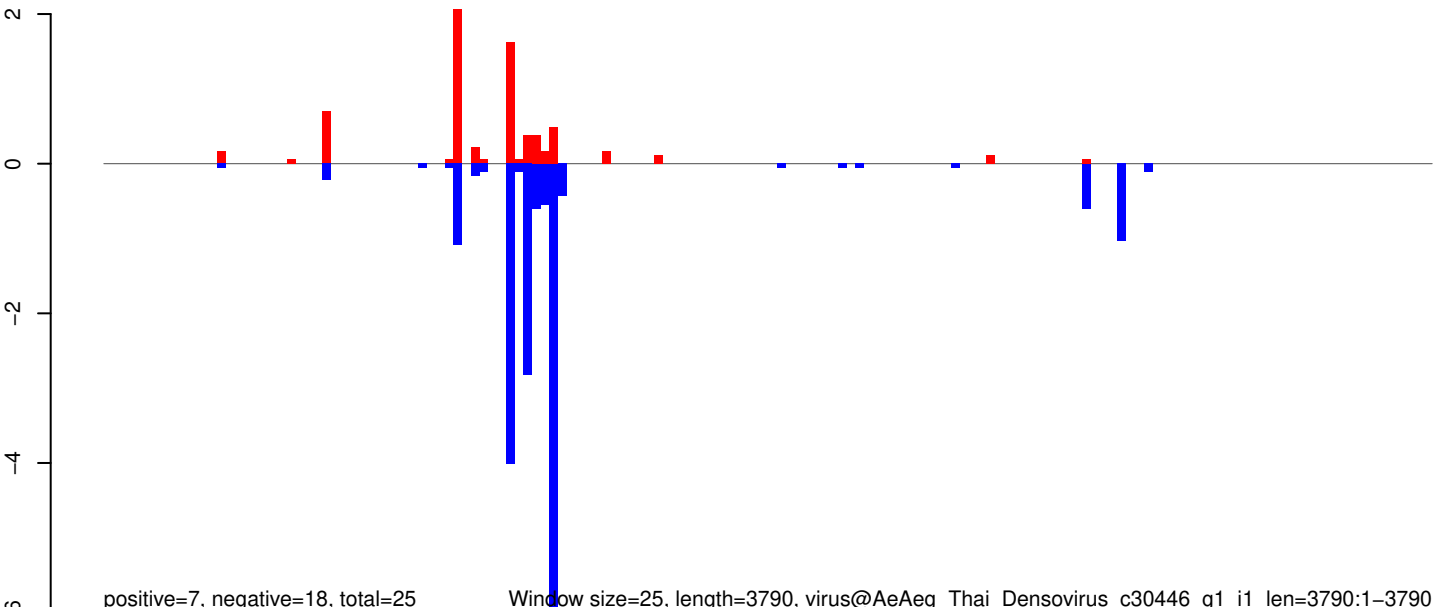
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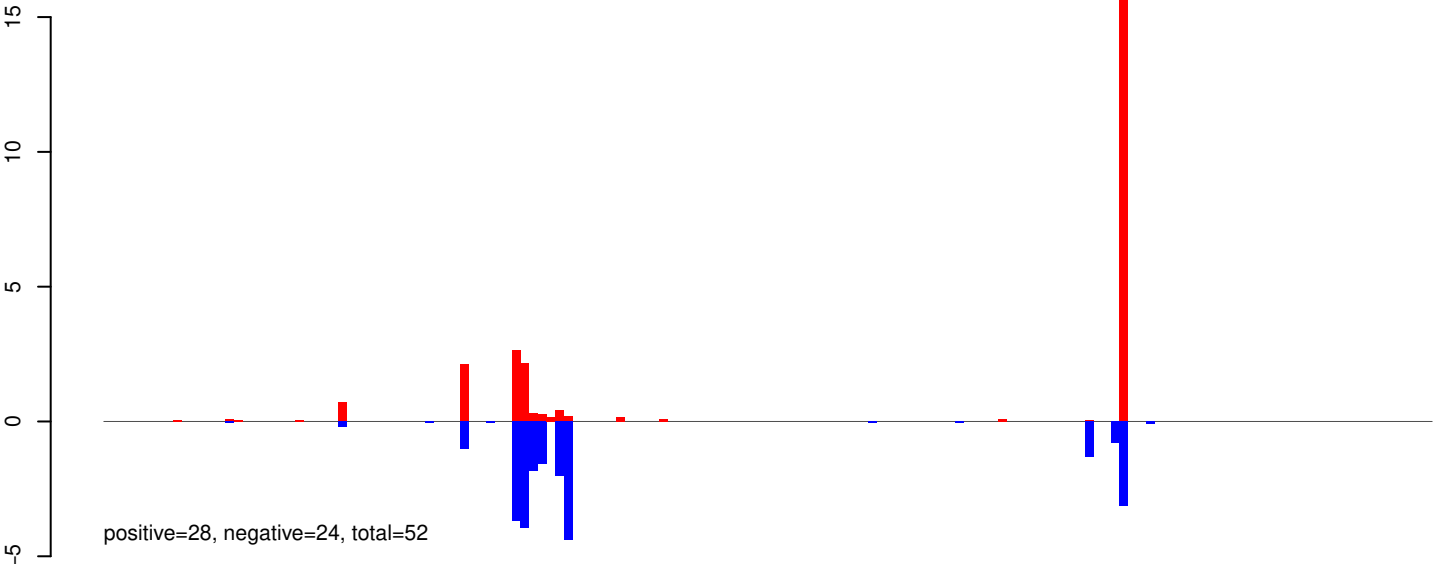


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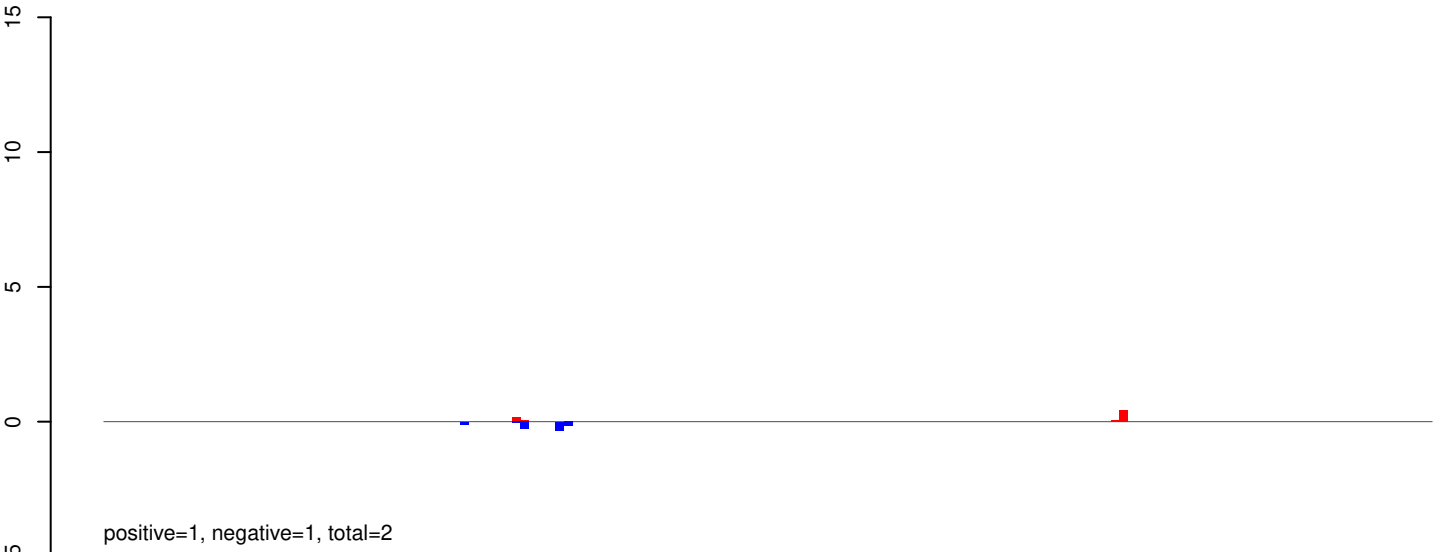


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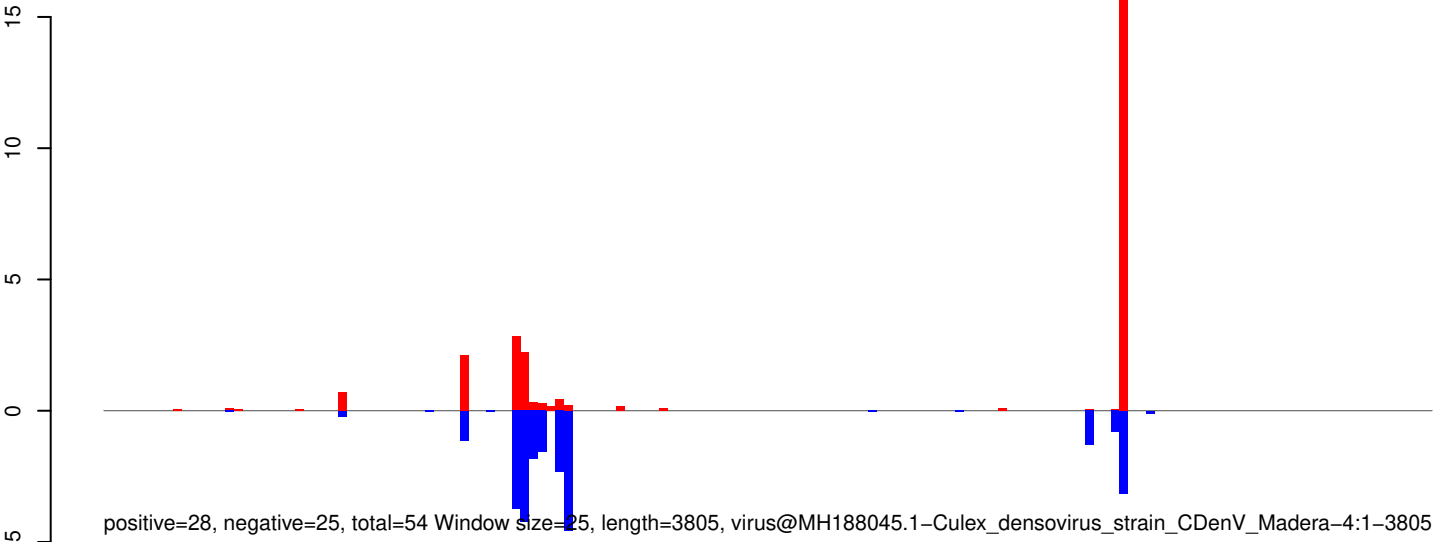
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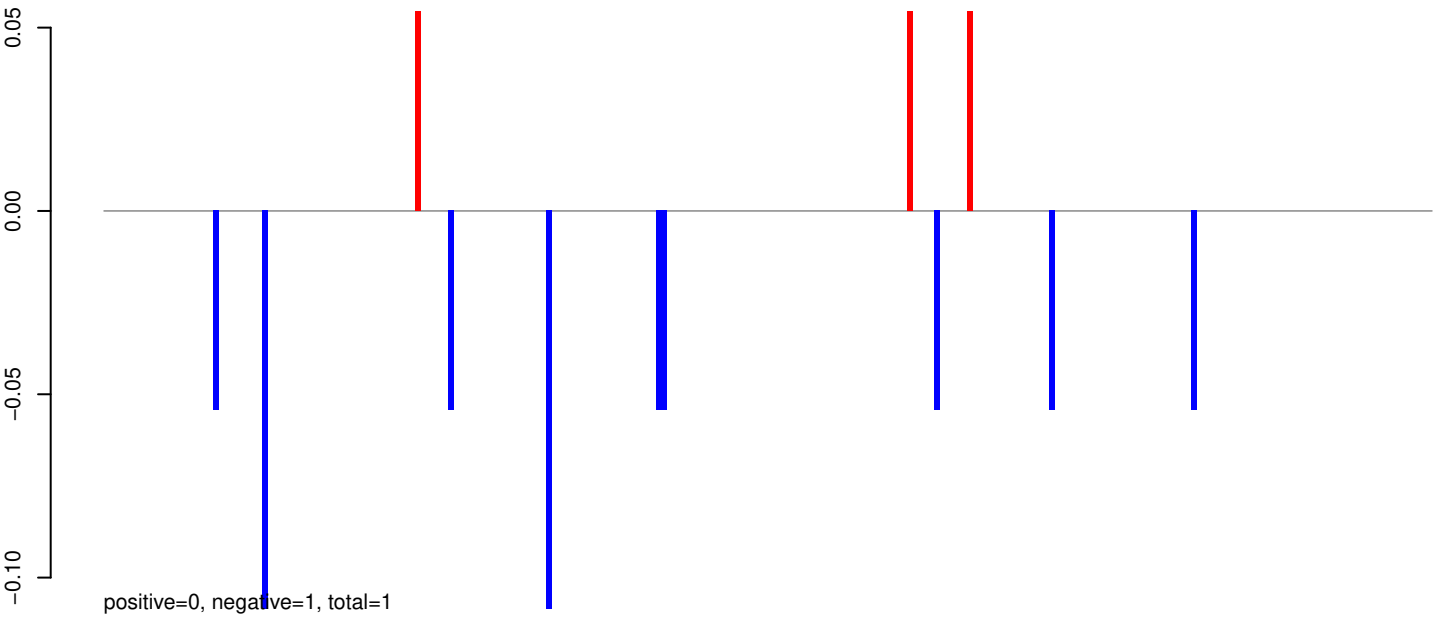
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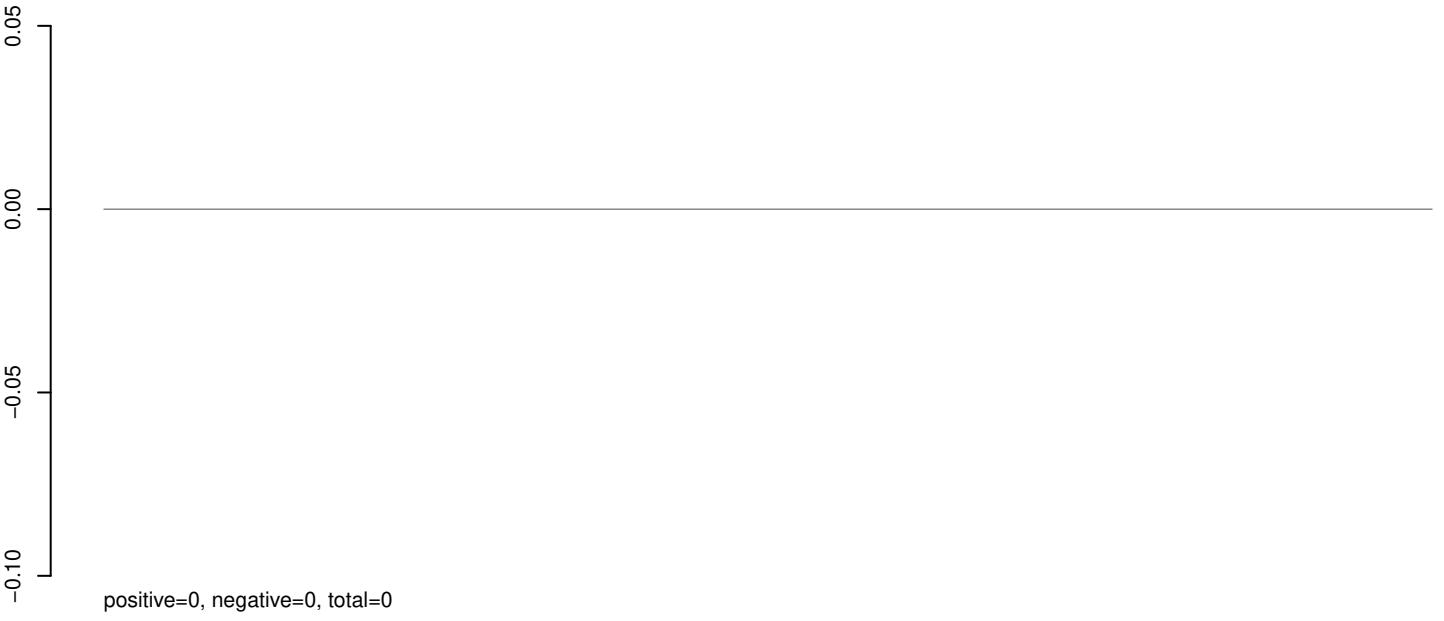
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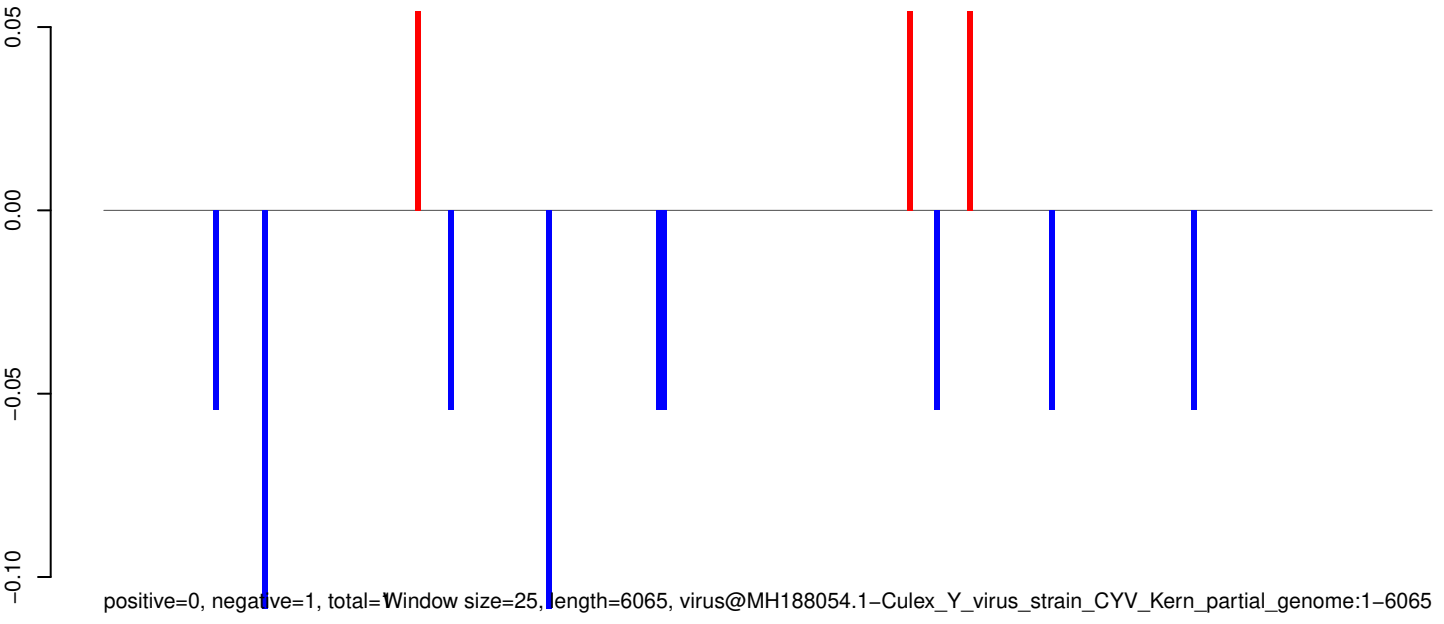
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



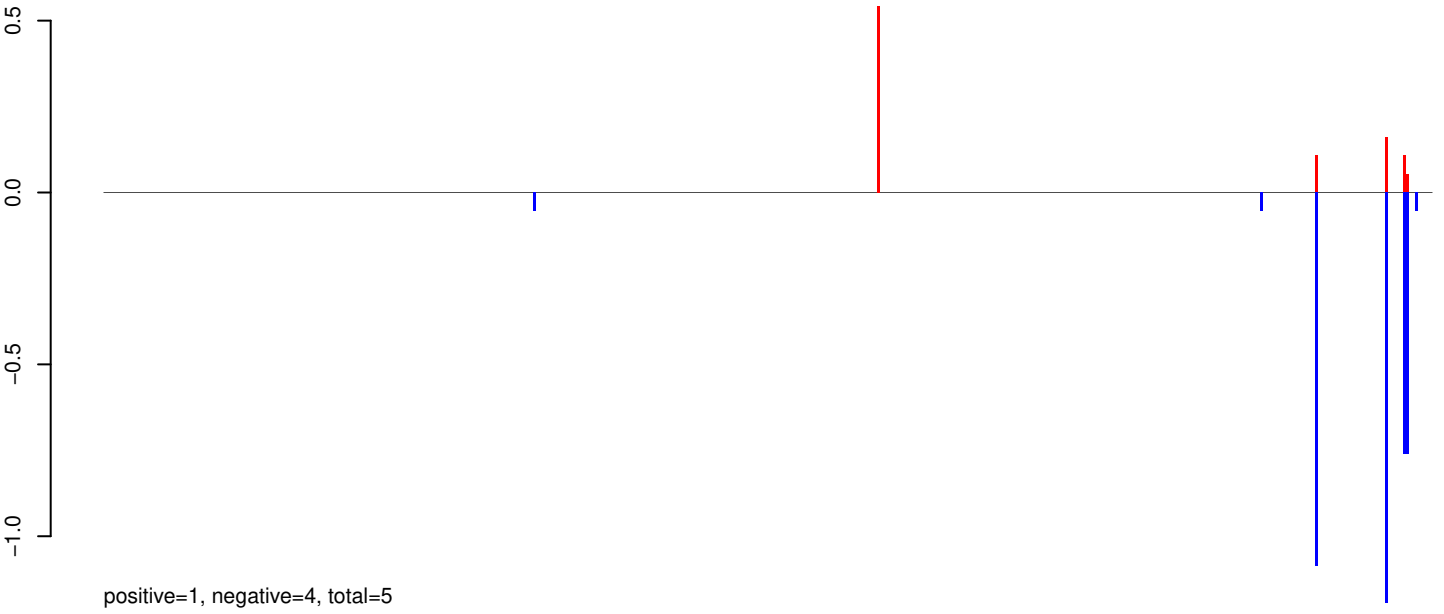
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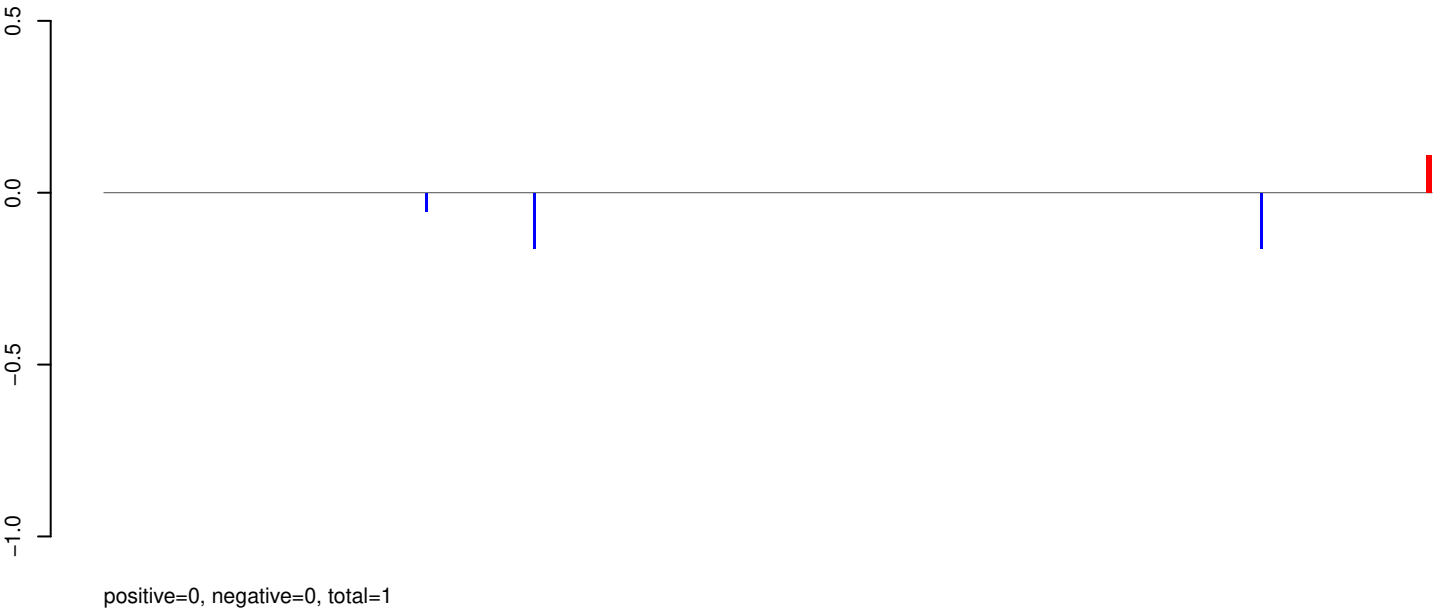
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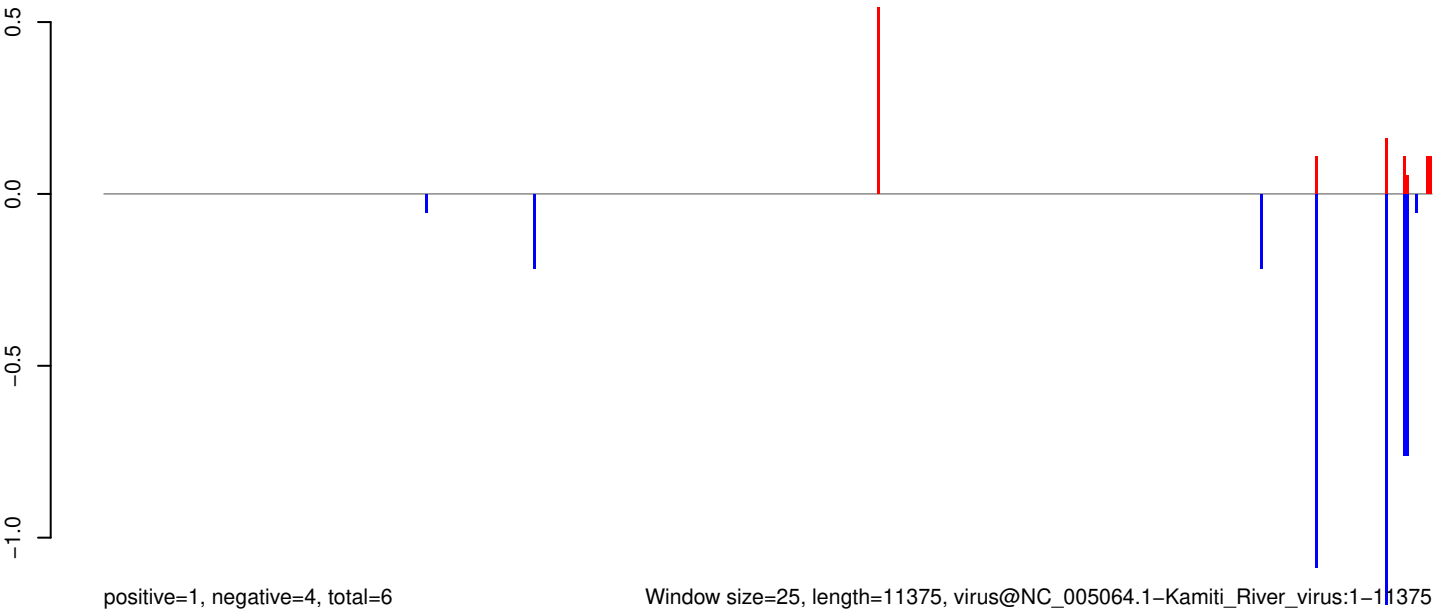
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



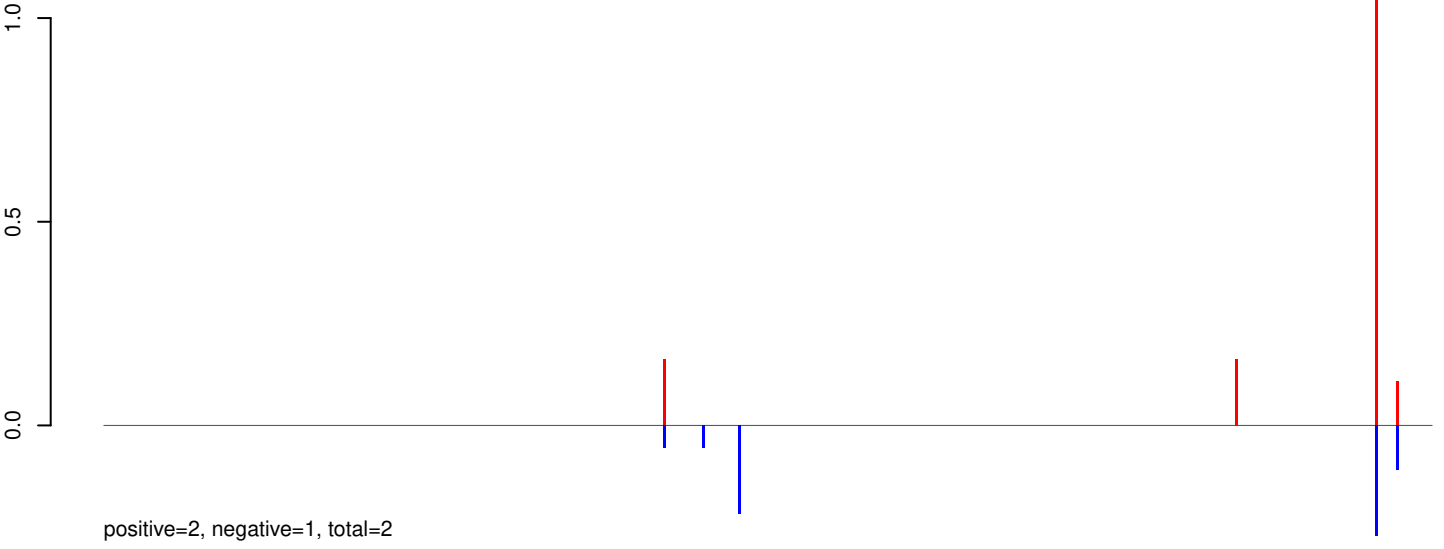
AeAeg_Aag2_BUNV1_ES.rep



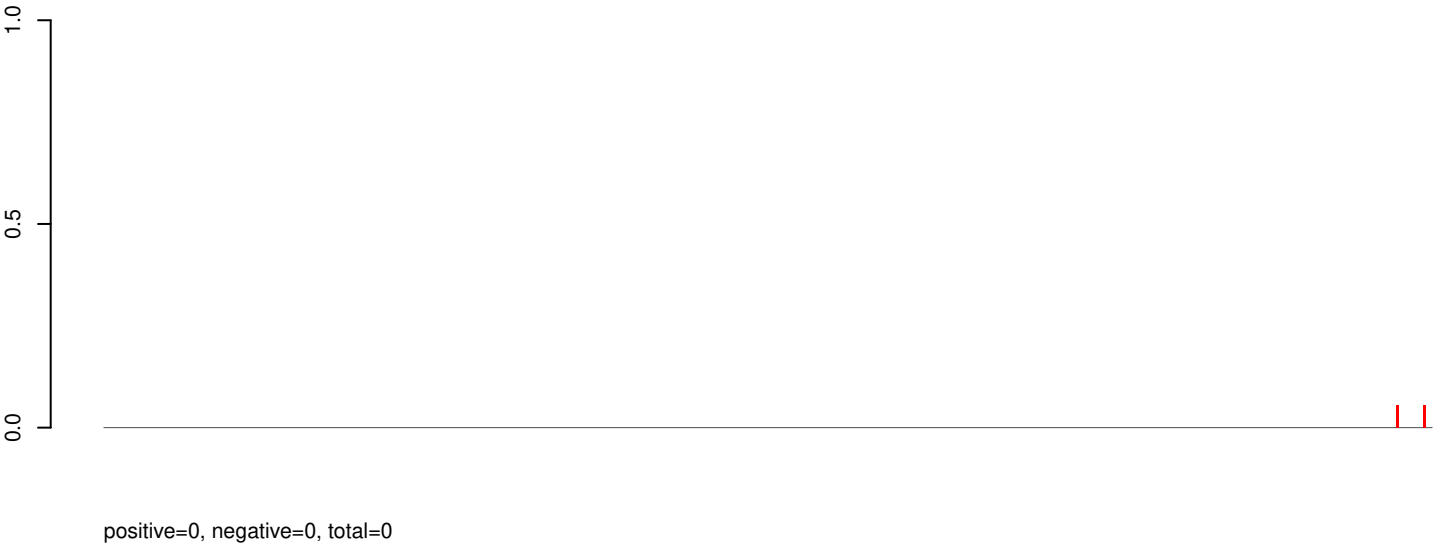
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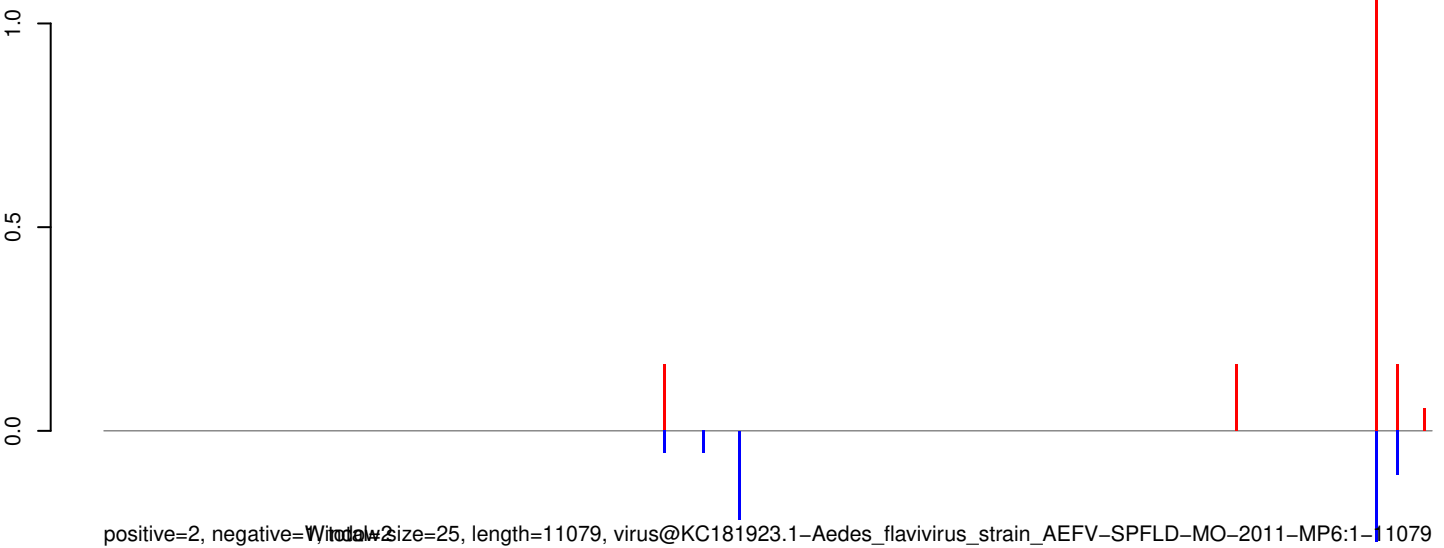
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep

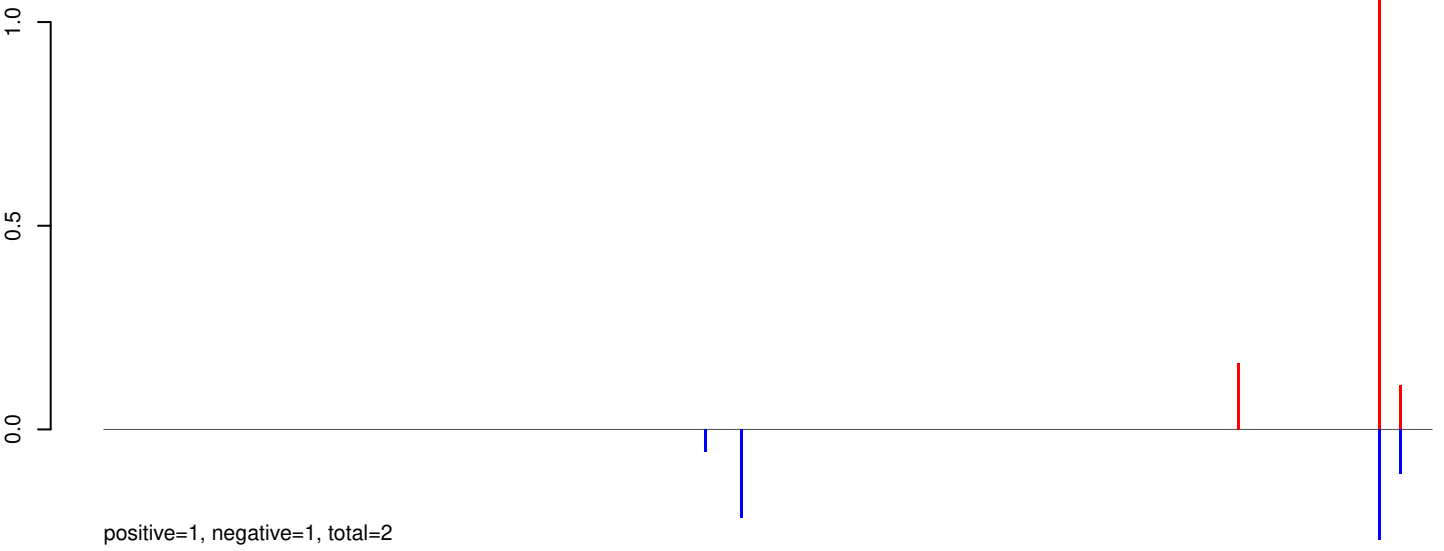


AeAeg_Aag2_BUNV1_ES.rep

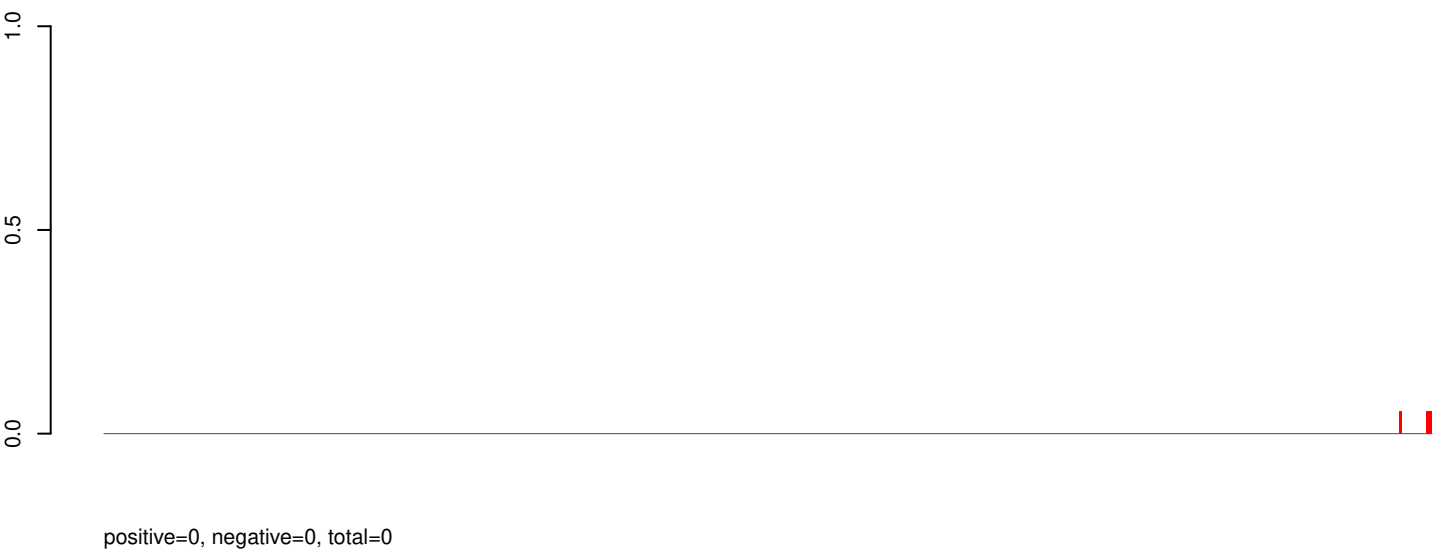


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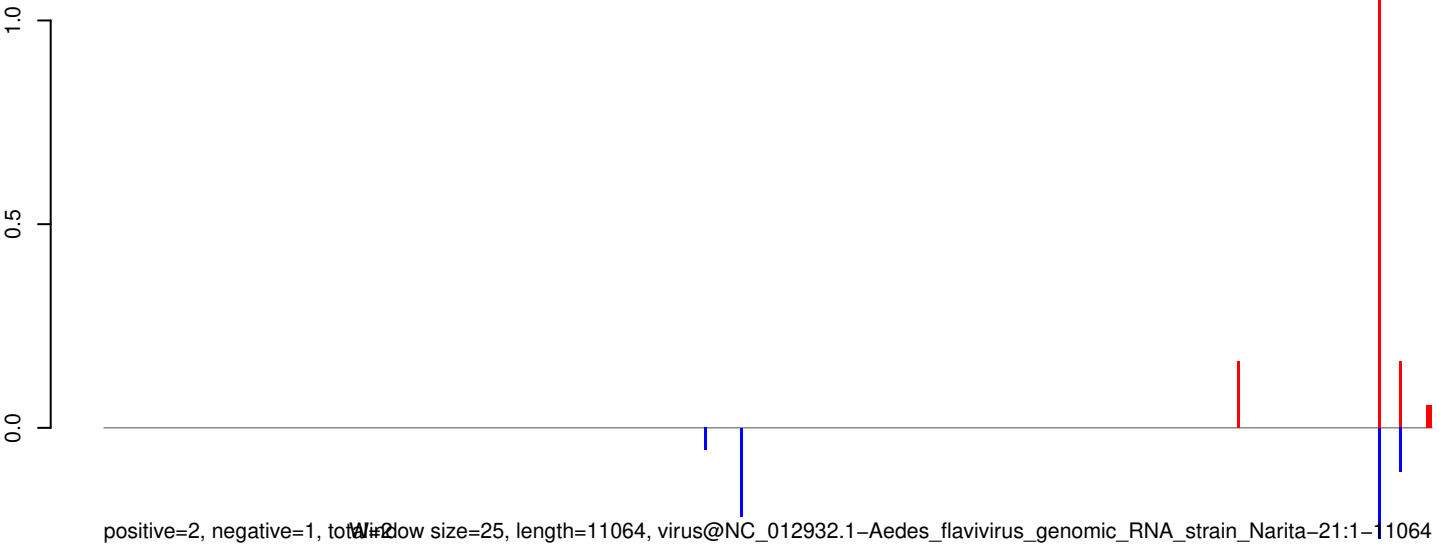
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AeAeg_Aag2_BUNV1_ES.24_35.rep



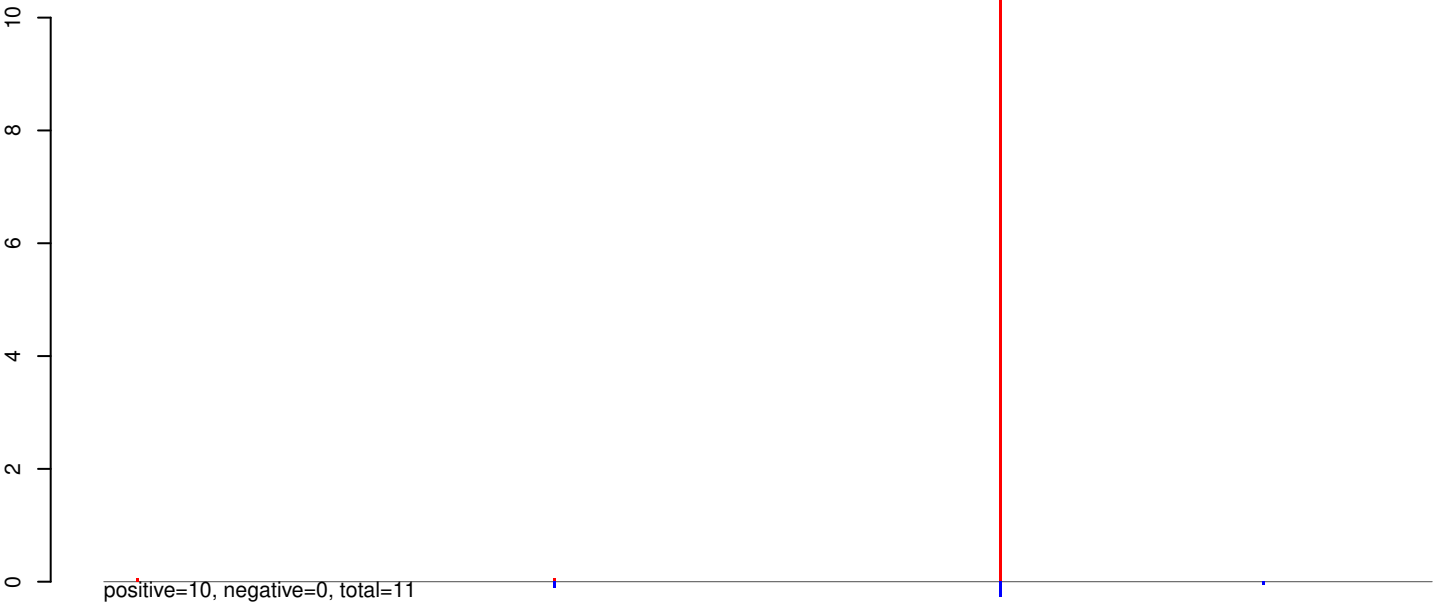
AeAeg_Aag2_BUNV1_ES.rep



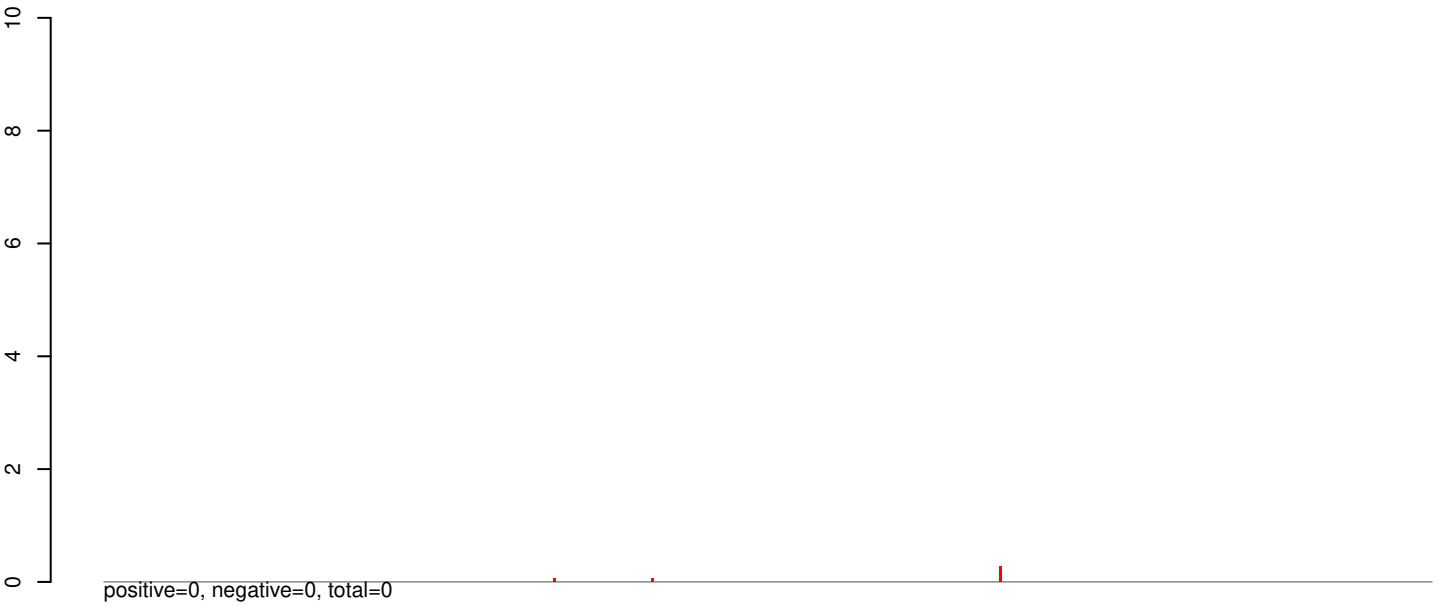
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0 2000 4000 6000 8000 10000

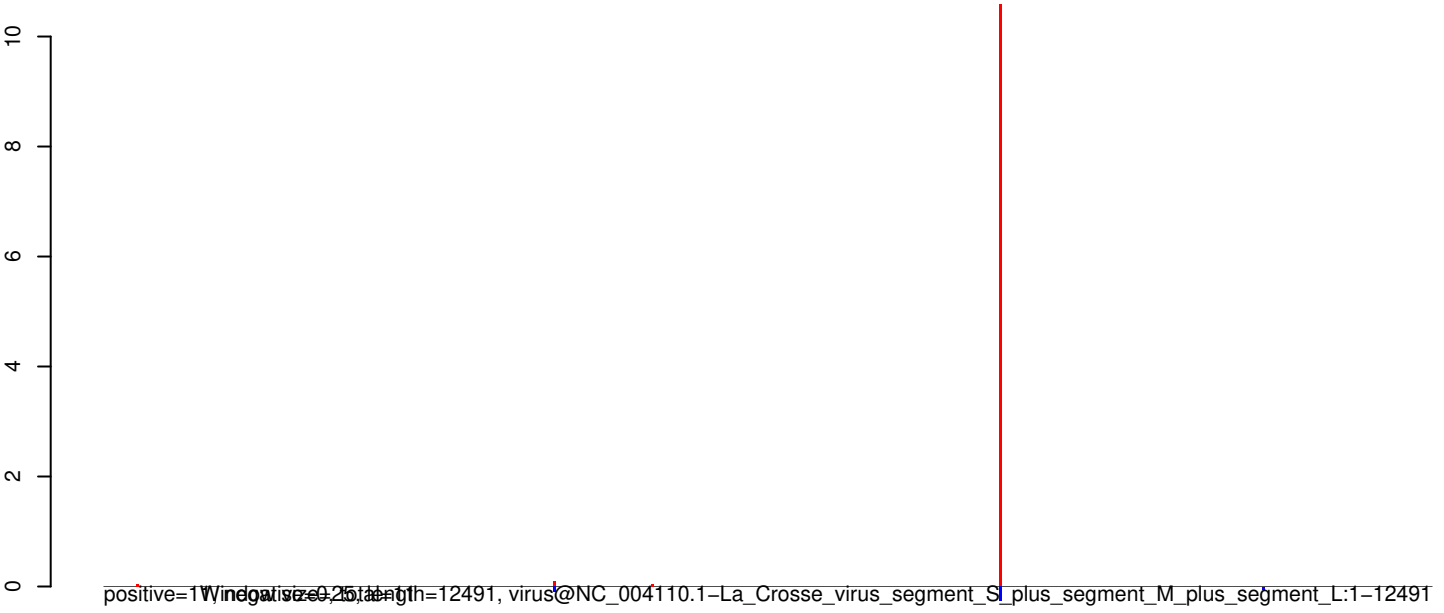
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



AeAeg_Aag2_BUNV1_ES.rep



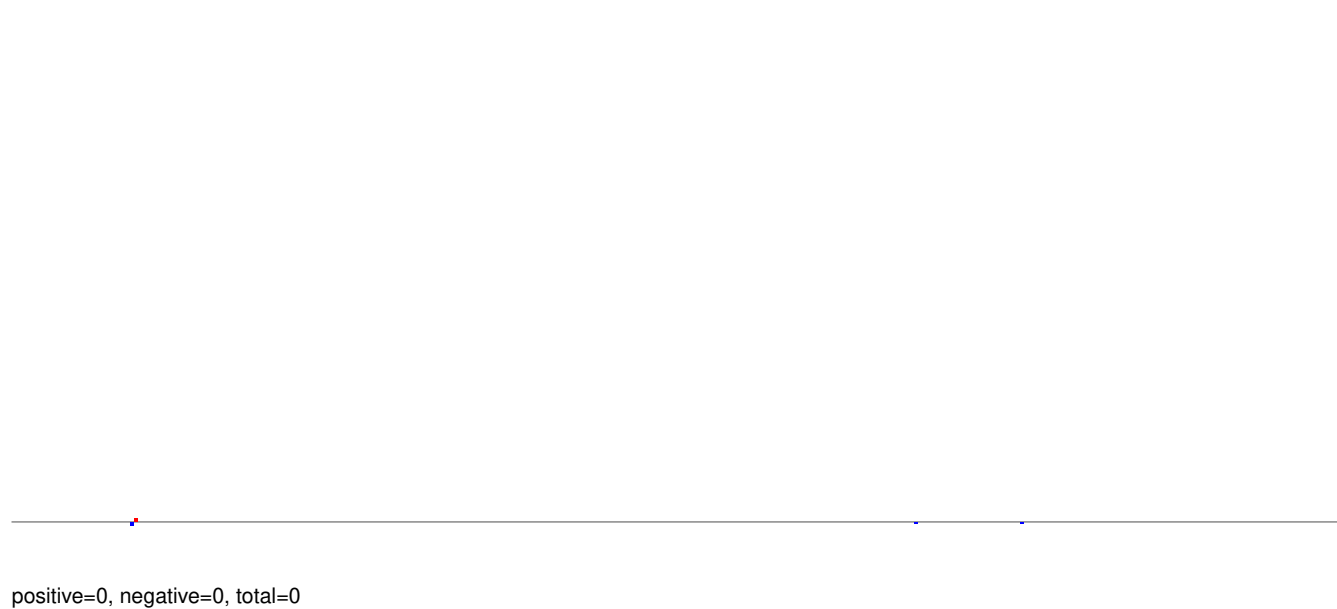
Window size: 25, step: 1, width=12491, virus@NC_004110.1-La_Crosse_virus_segment_S_plus_segment_M_plus_segment_L:1-12491

0 2000 4000 6000 8000 10000 12000

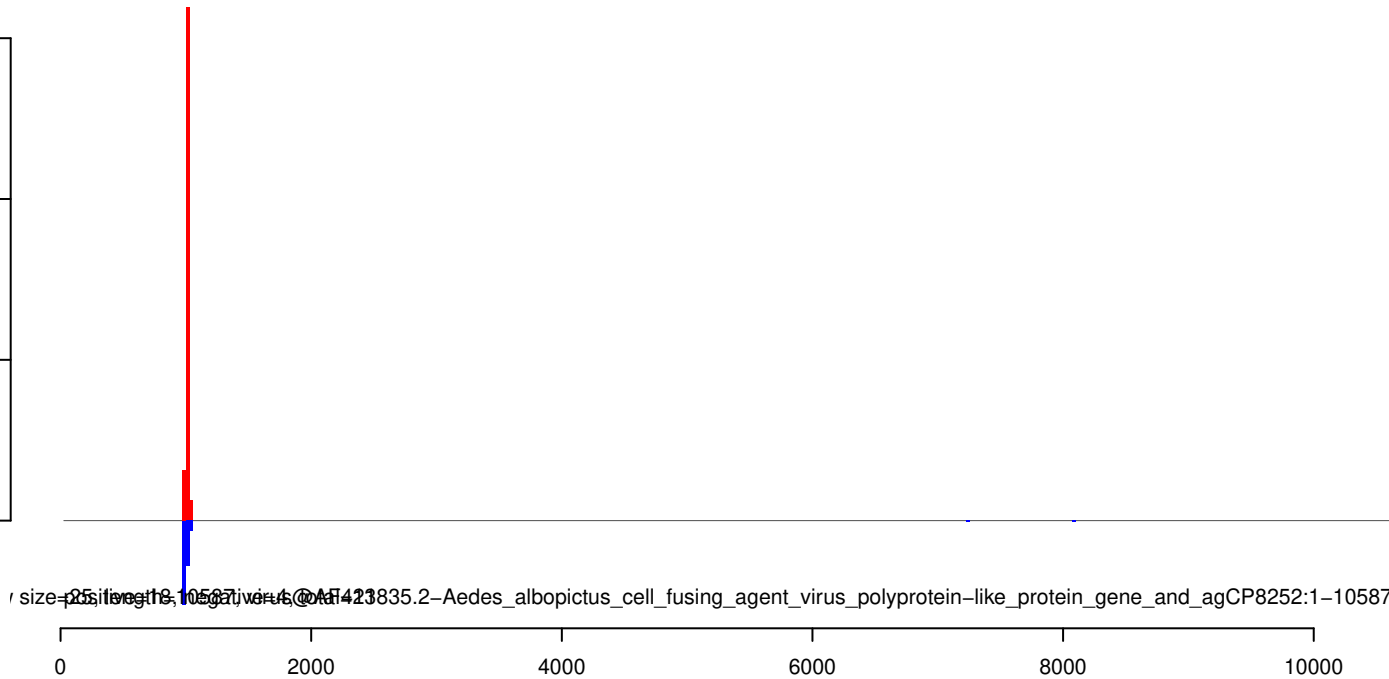
AeAeg_Aag2_BUNV1_ES.18_23.rep



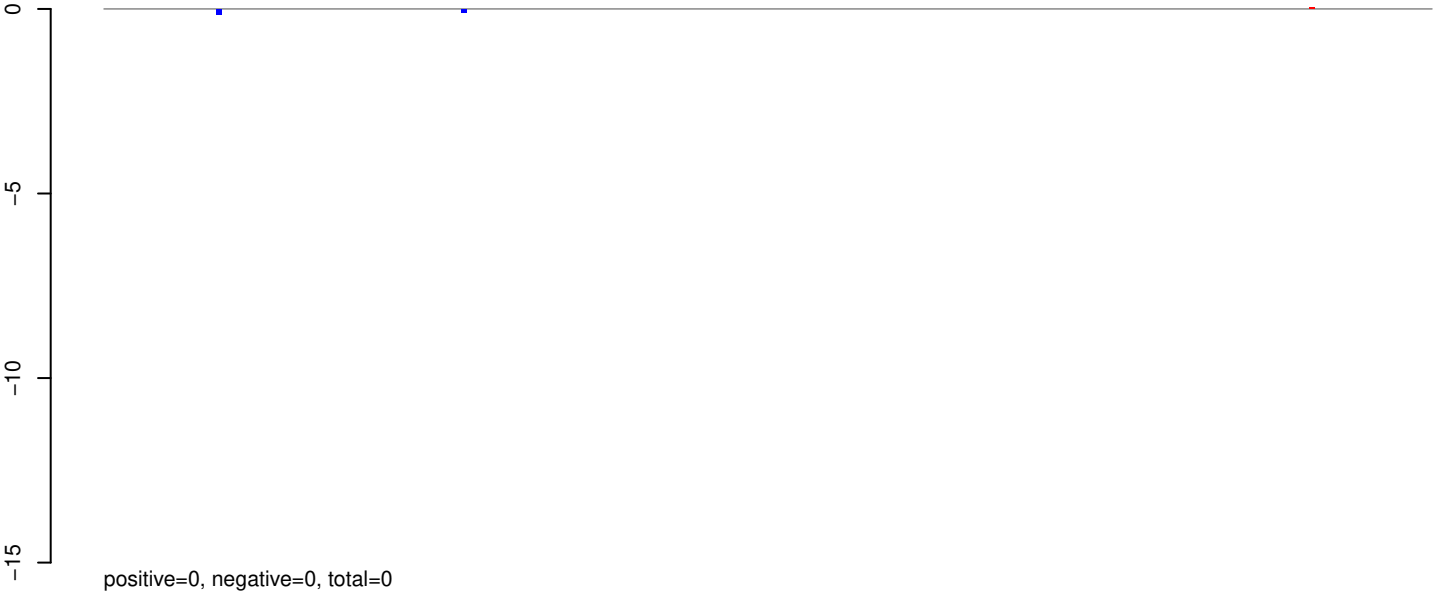
AeAeg_Aag2_BUNV1_ES.24_35.rep



AeAeg_Aag2_BUNV1_ES.rep



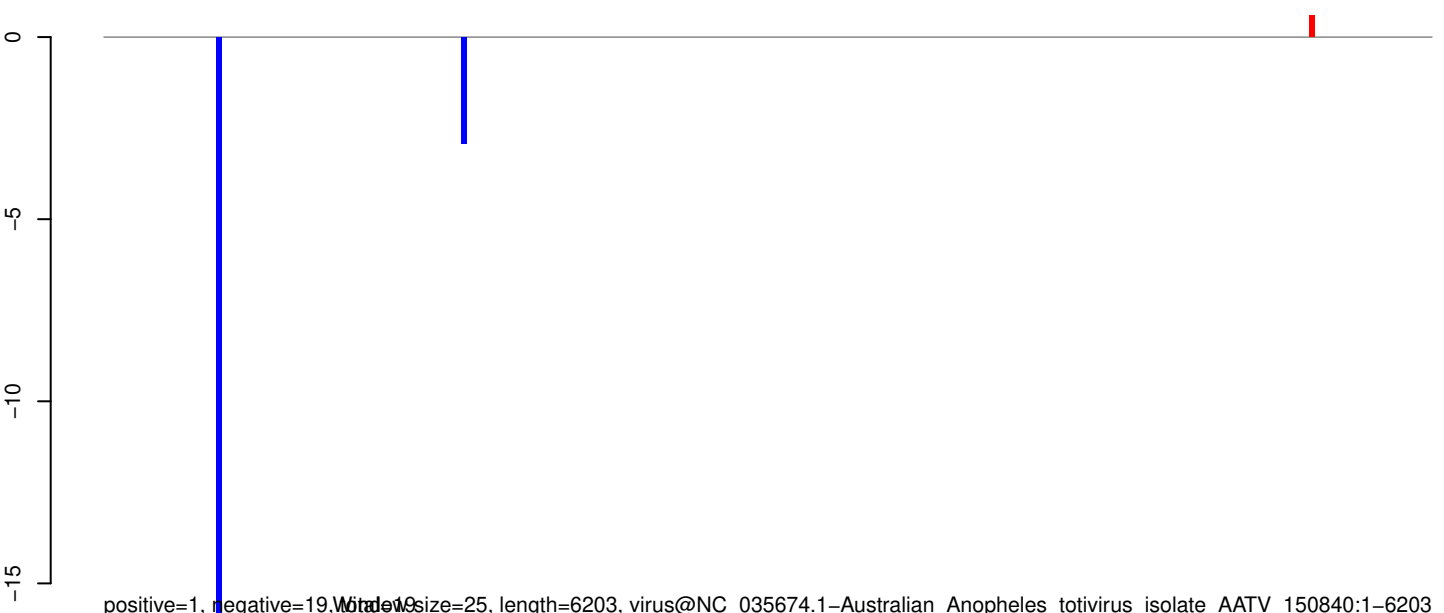
AeAeg_Aag2_BUNV1_ES.18_23.rep



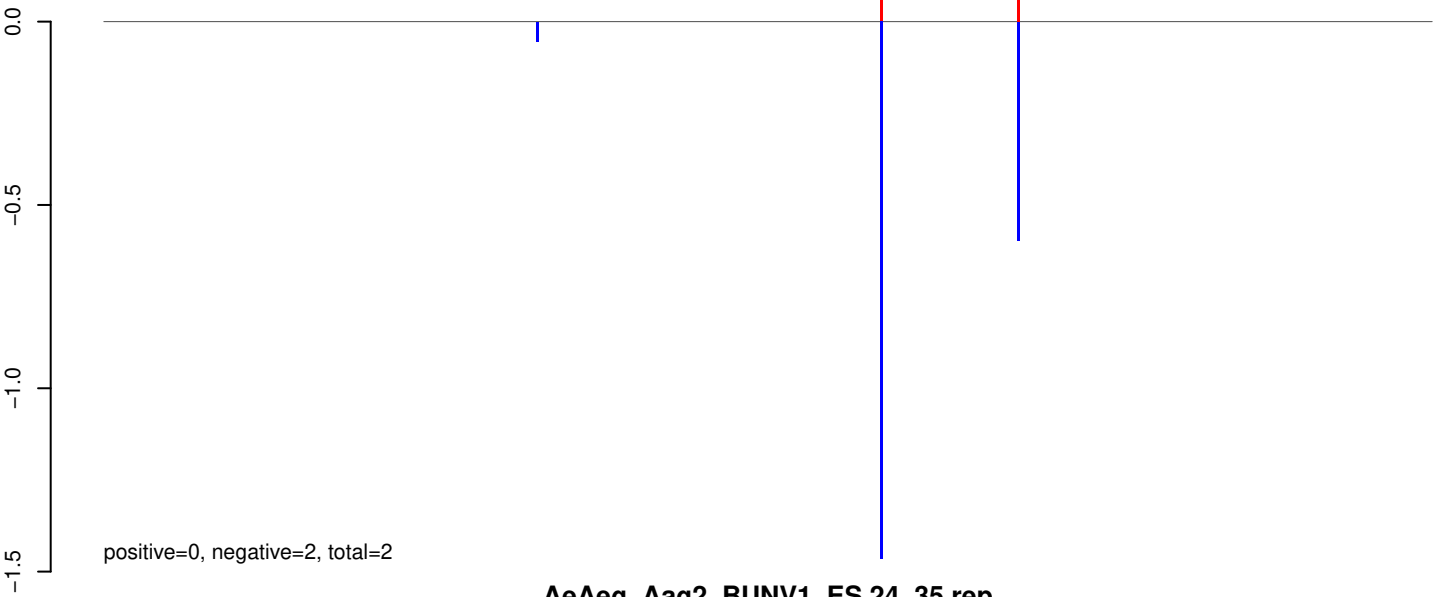
AeAeg_Aag2_BUNV1_ES.24_35.rep



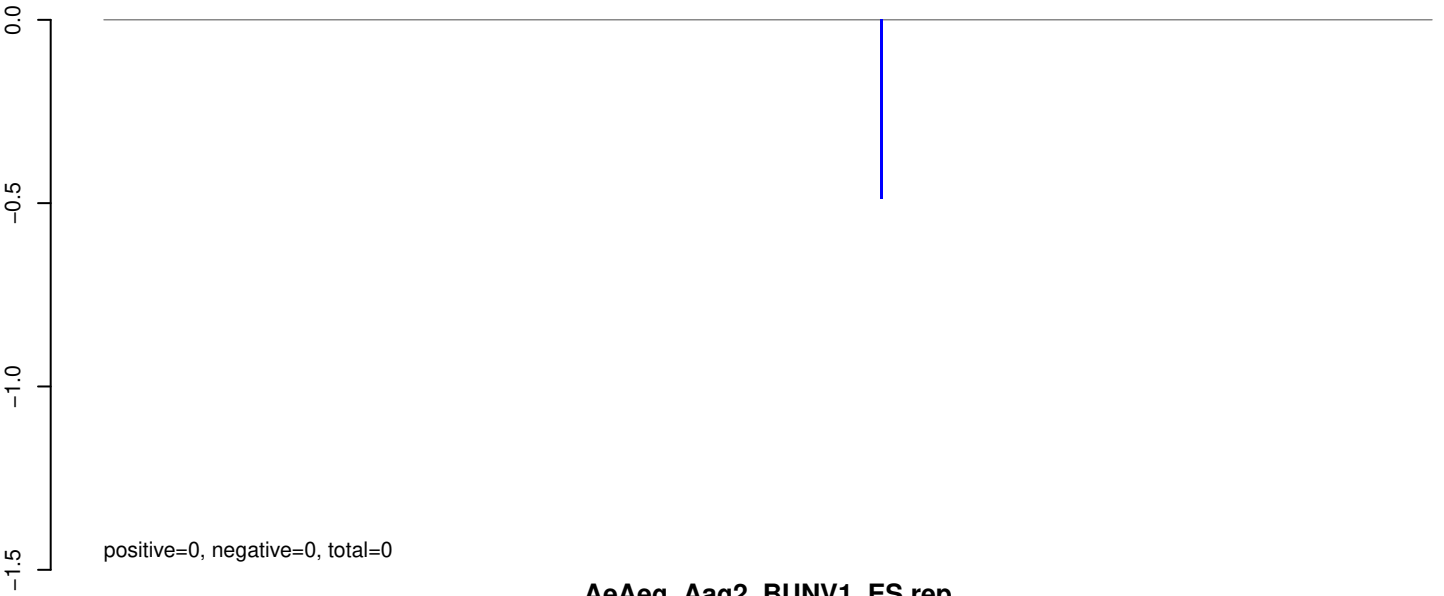
AeAeg_Aag2_BUNV1_ES.rep



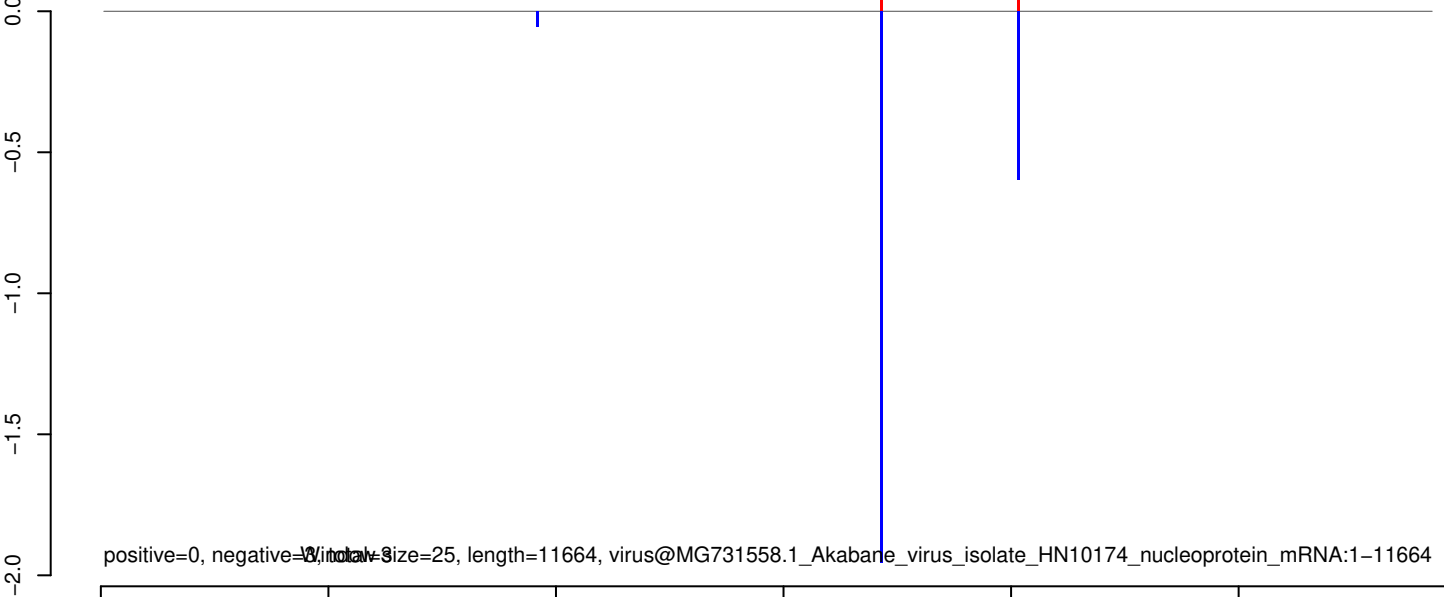
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



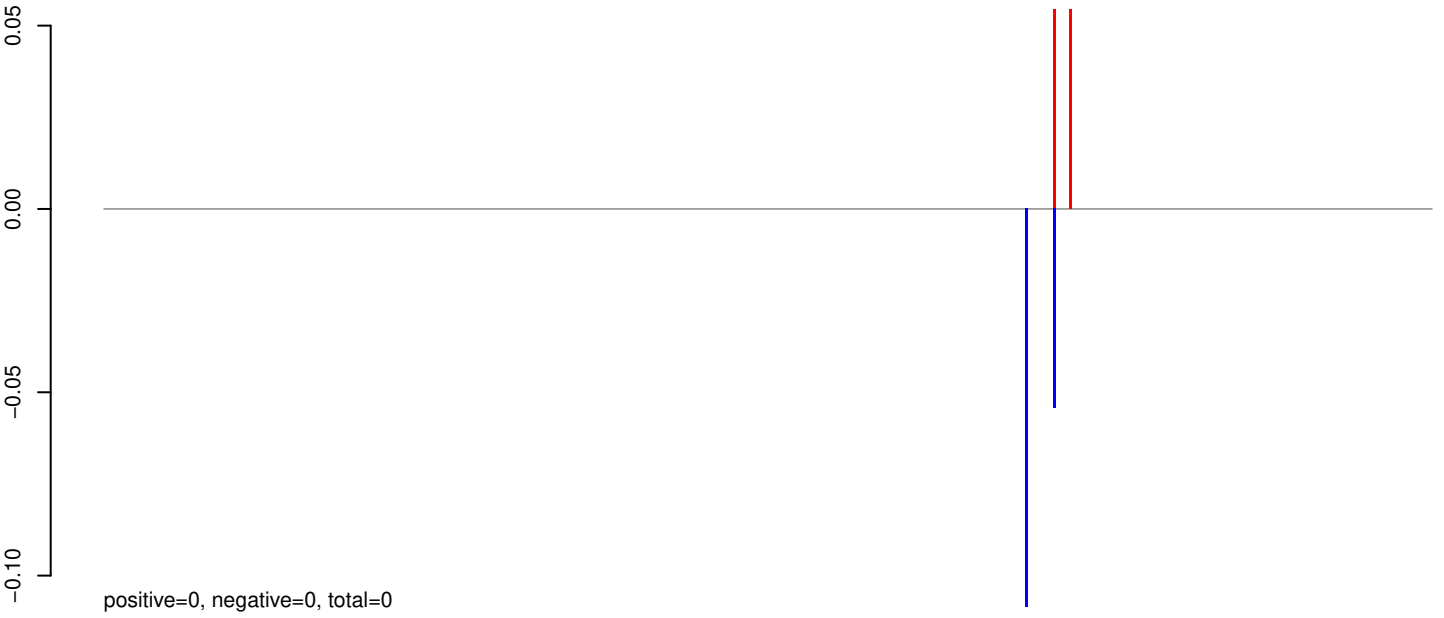
AeAeg_Aag2_BUNV1_ES.rep



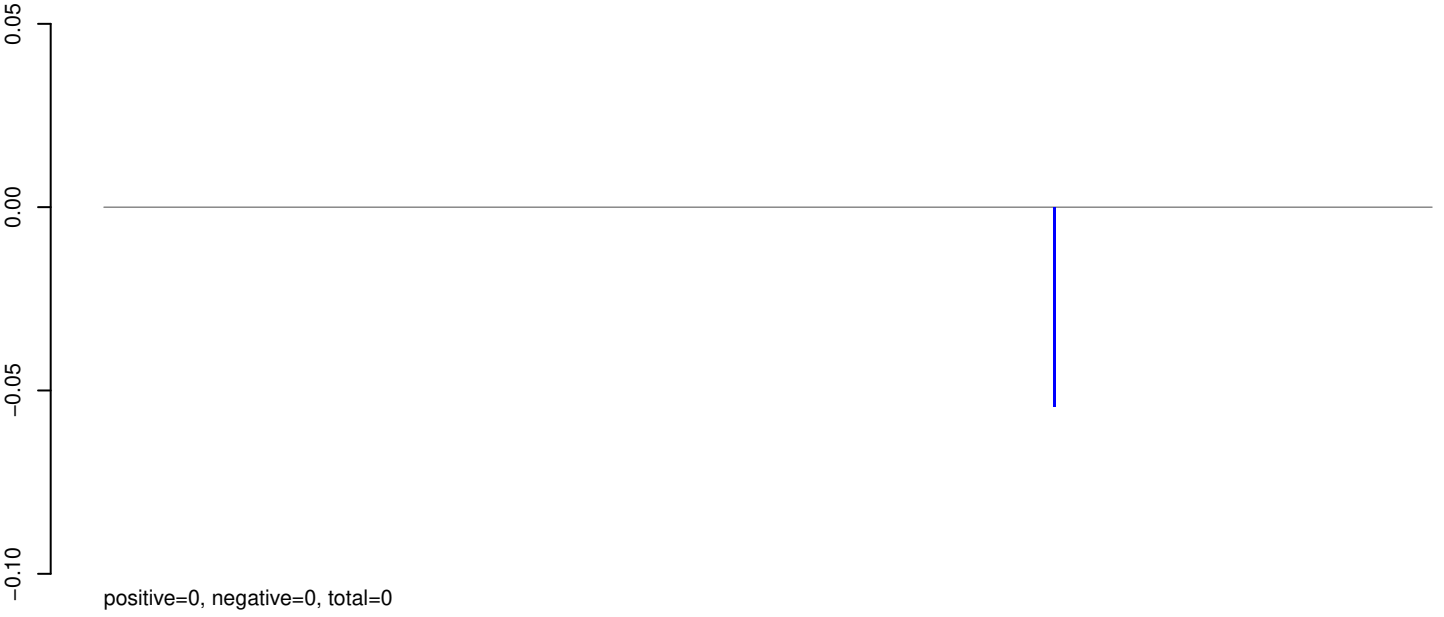
size=25, length=11664, virus@MG731558.1_Akabane_virus_isolate_HN10174_nucleoprotein_mRNA:1-11664

0 2000 4000 6000 8000 10000 12000

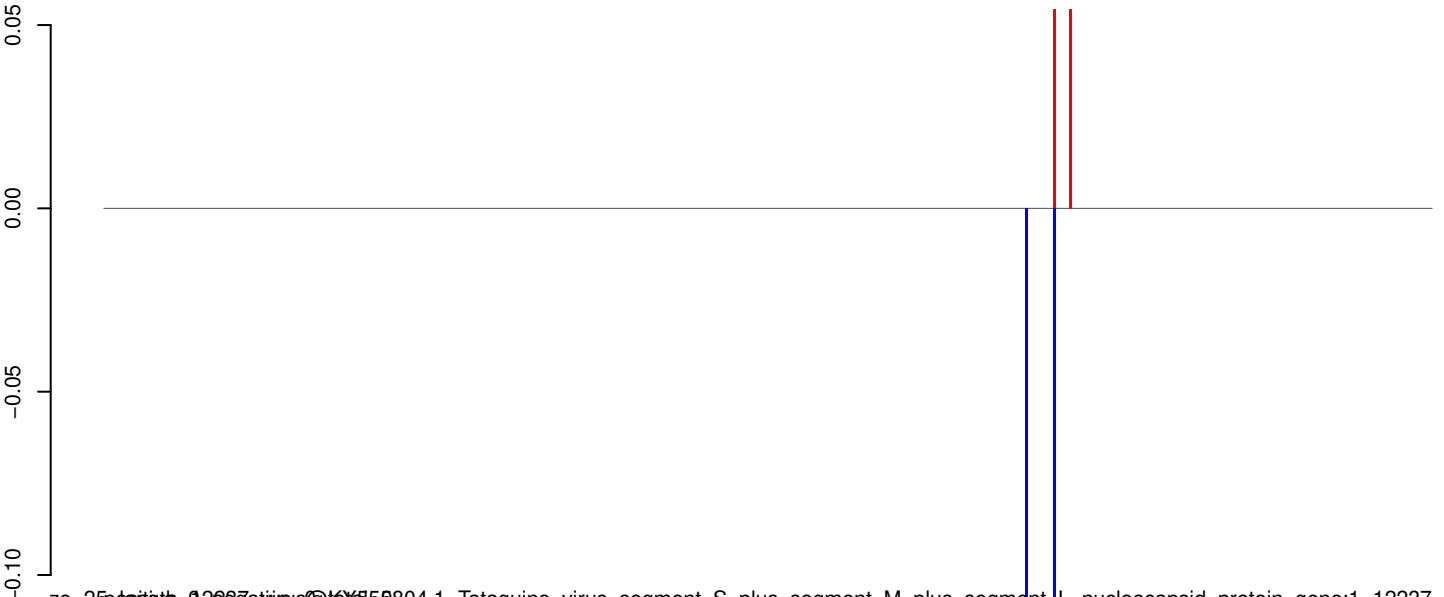
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



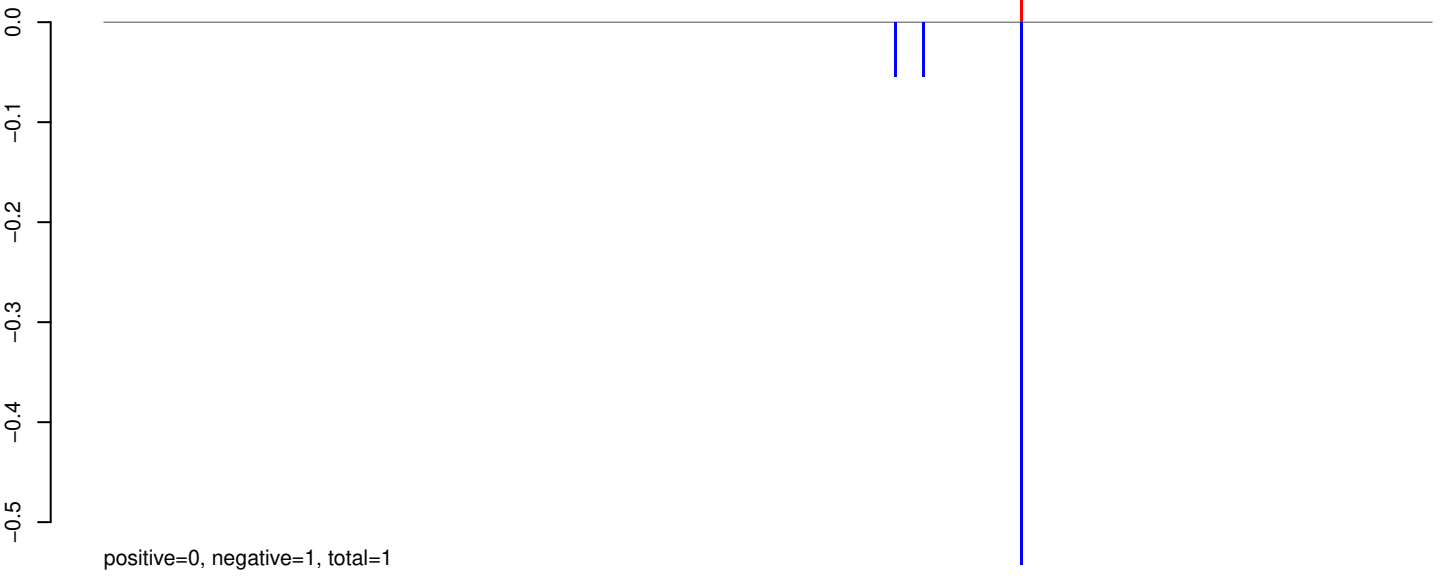
AeAeg_Aag2_BUNV1_ES.rep



ze=25 path=1287 avic@1656804.1-Tataguine_virus_segment_S_plus_segment_M_plus_segment_L_nucleocapsid_protein_gene:1-12237

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_BUNV1_ES.18_23.rep



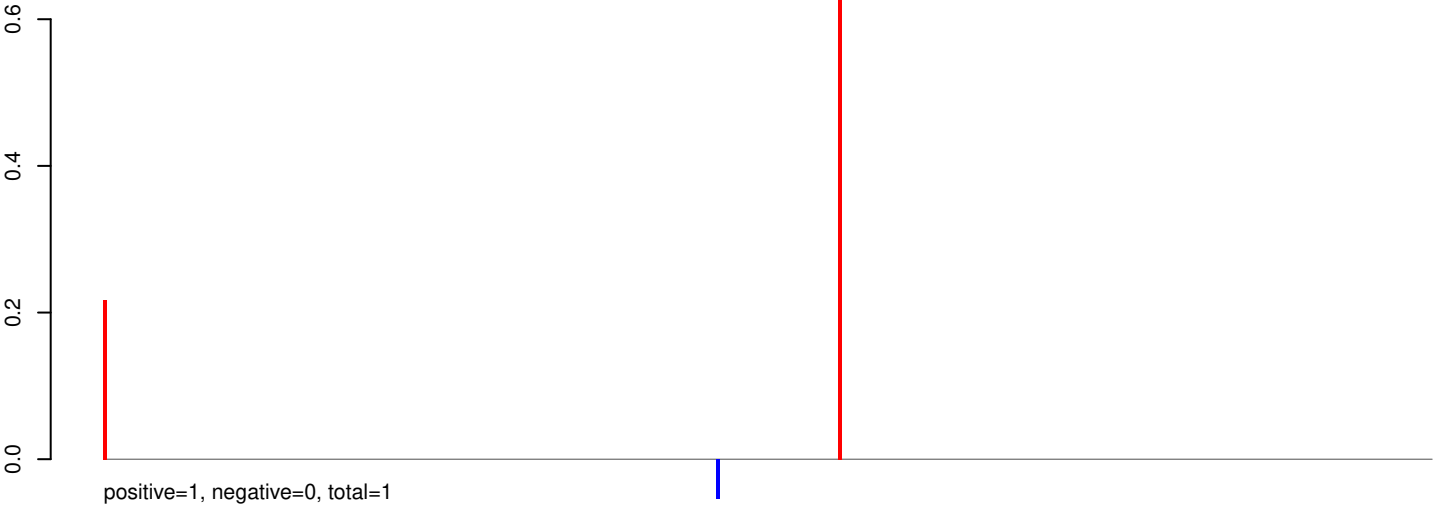
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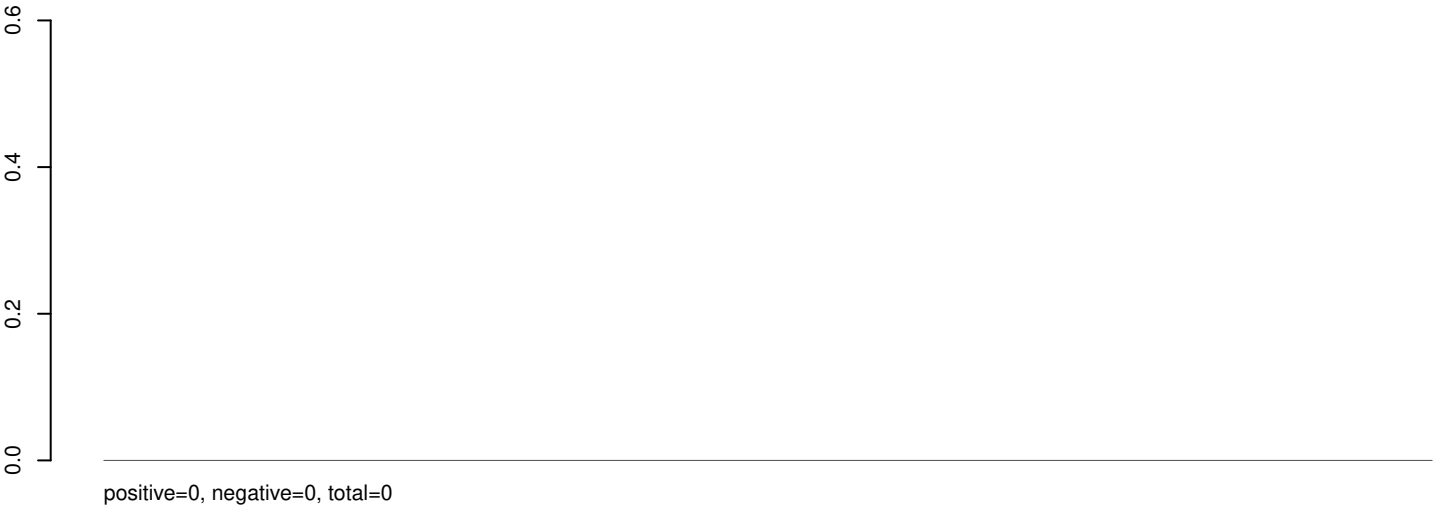
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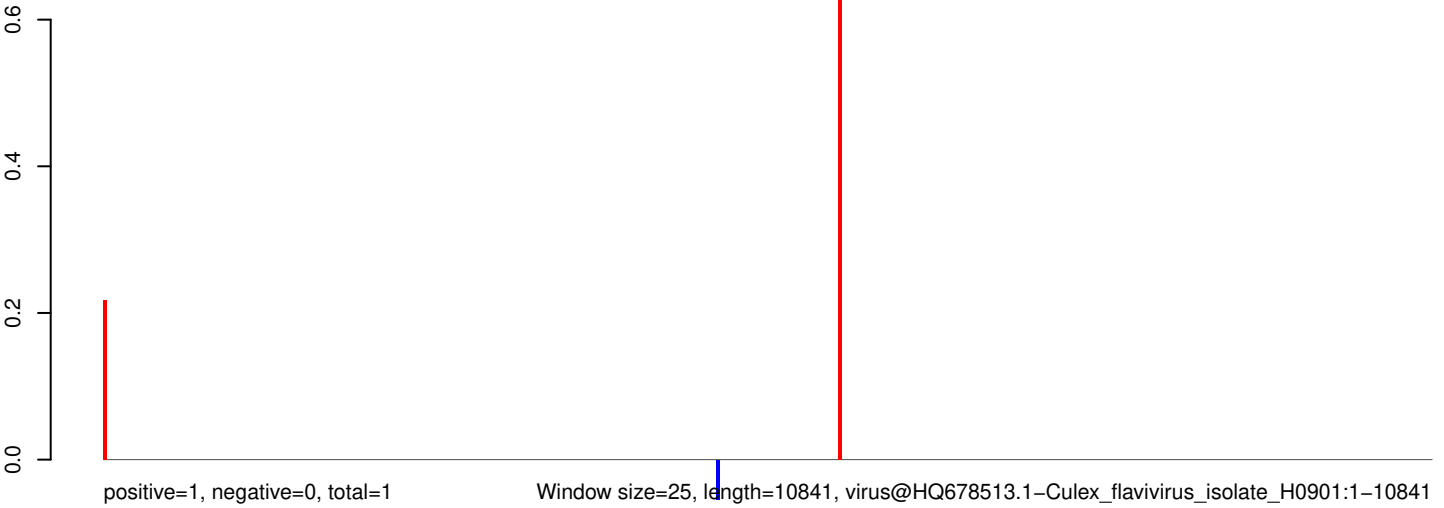
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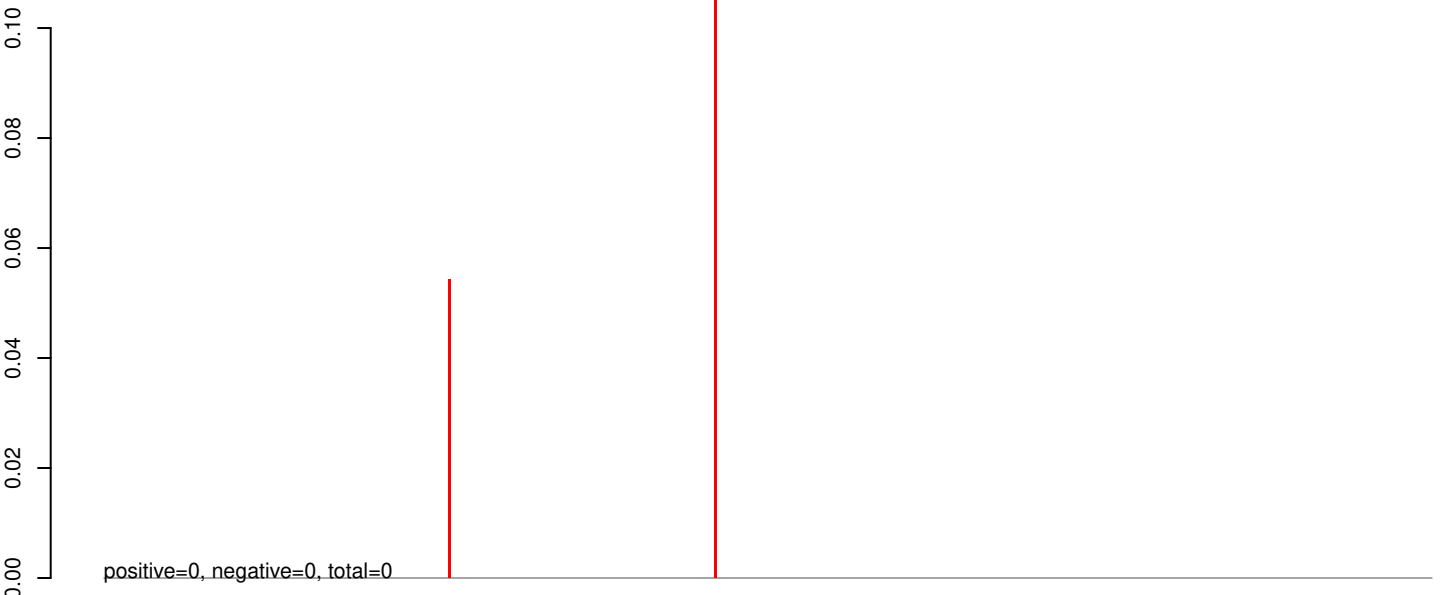
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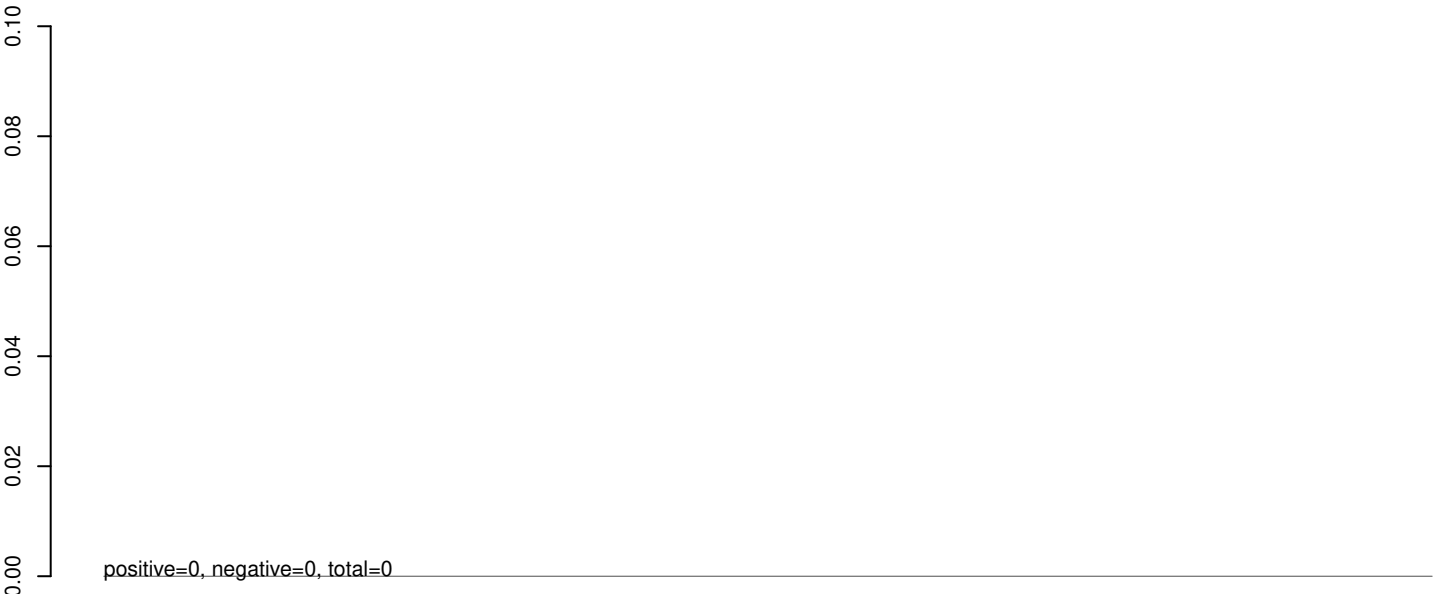
AeAeg_Aag2_BUNV1_ES.rep



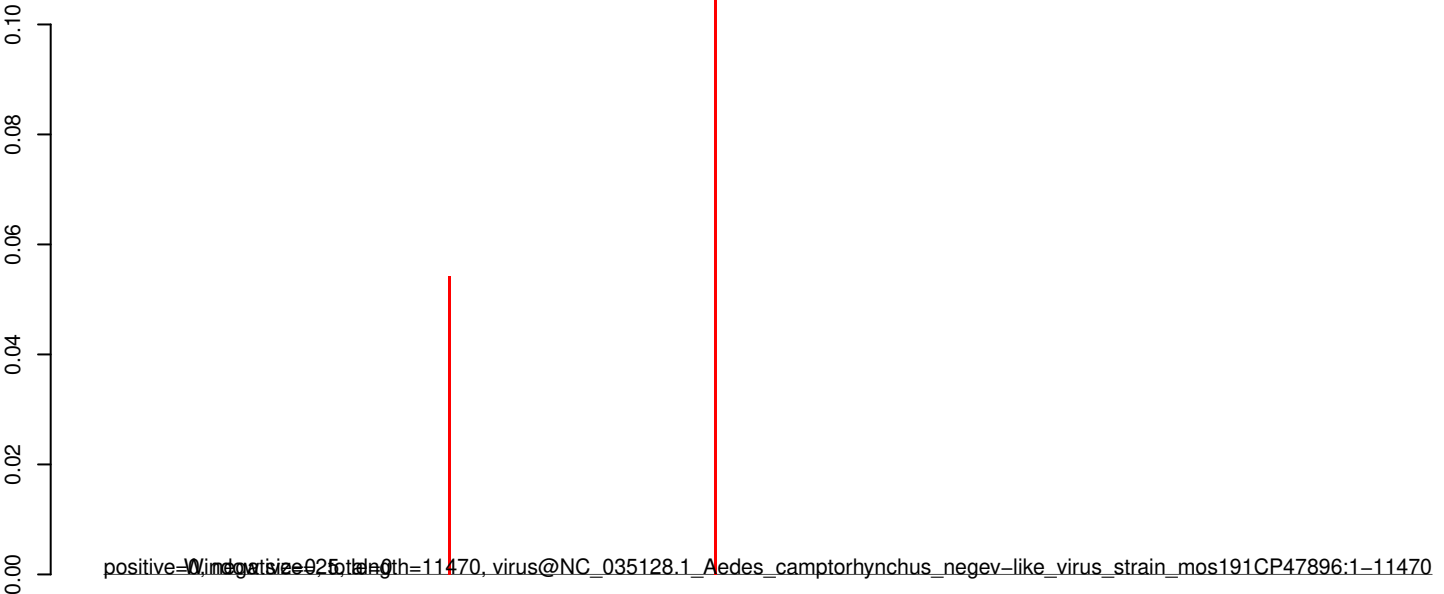
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep

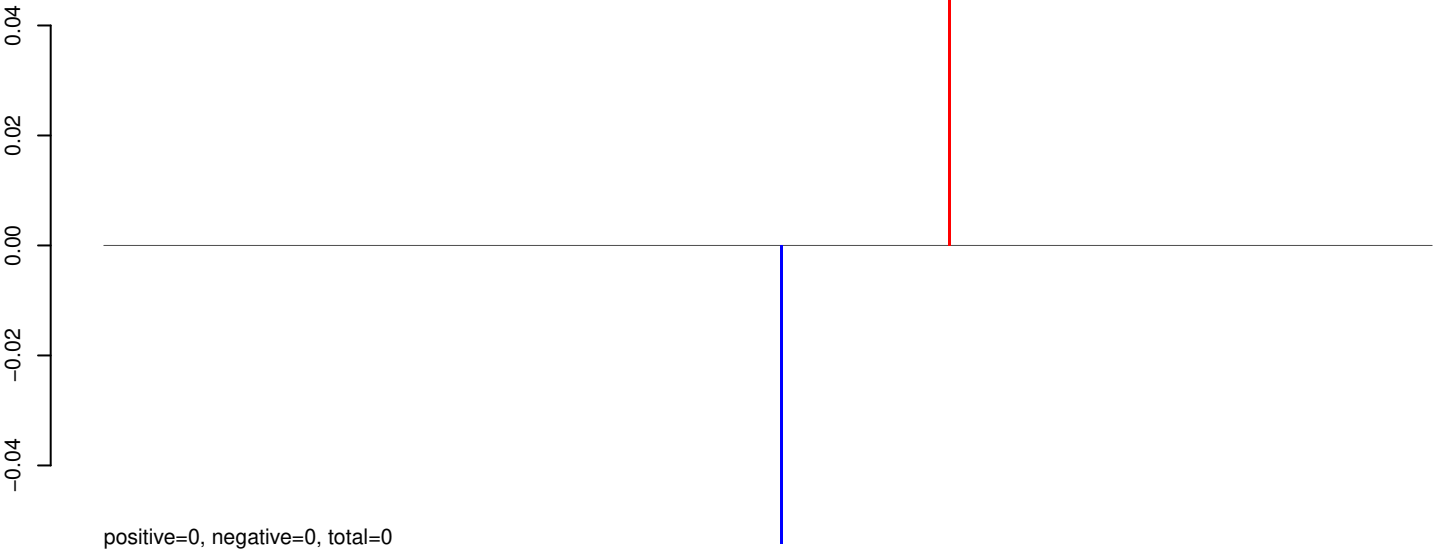


AeAeg_Aag2_BUNV1_ES.rep

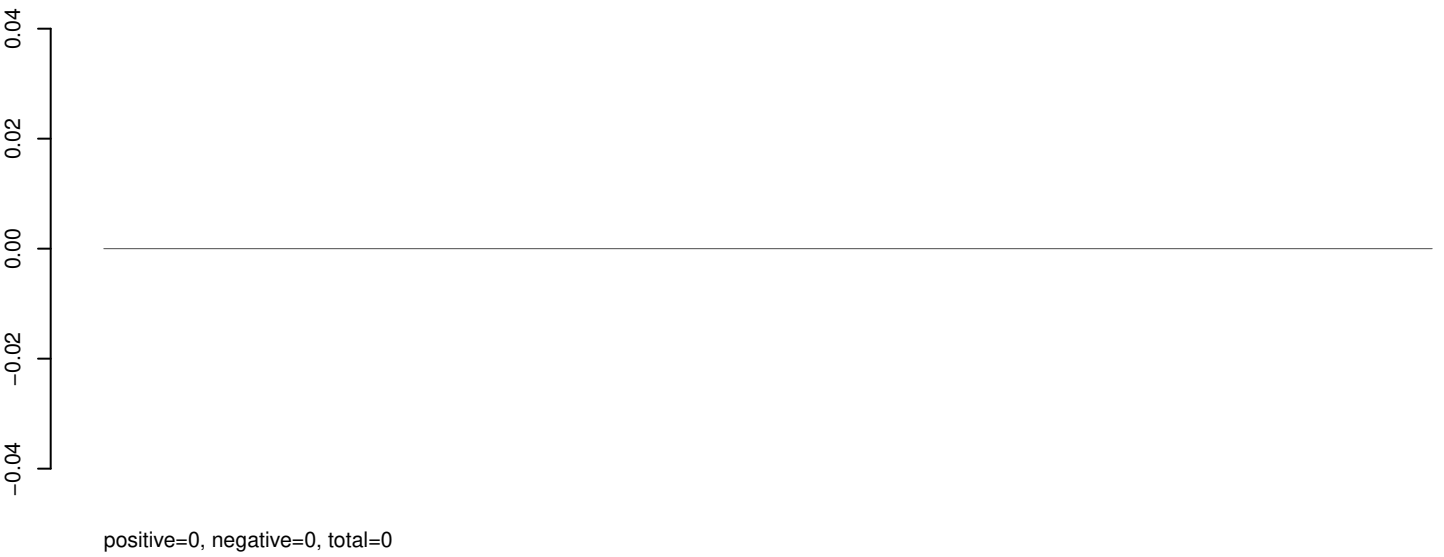


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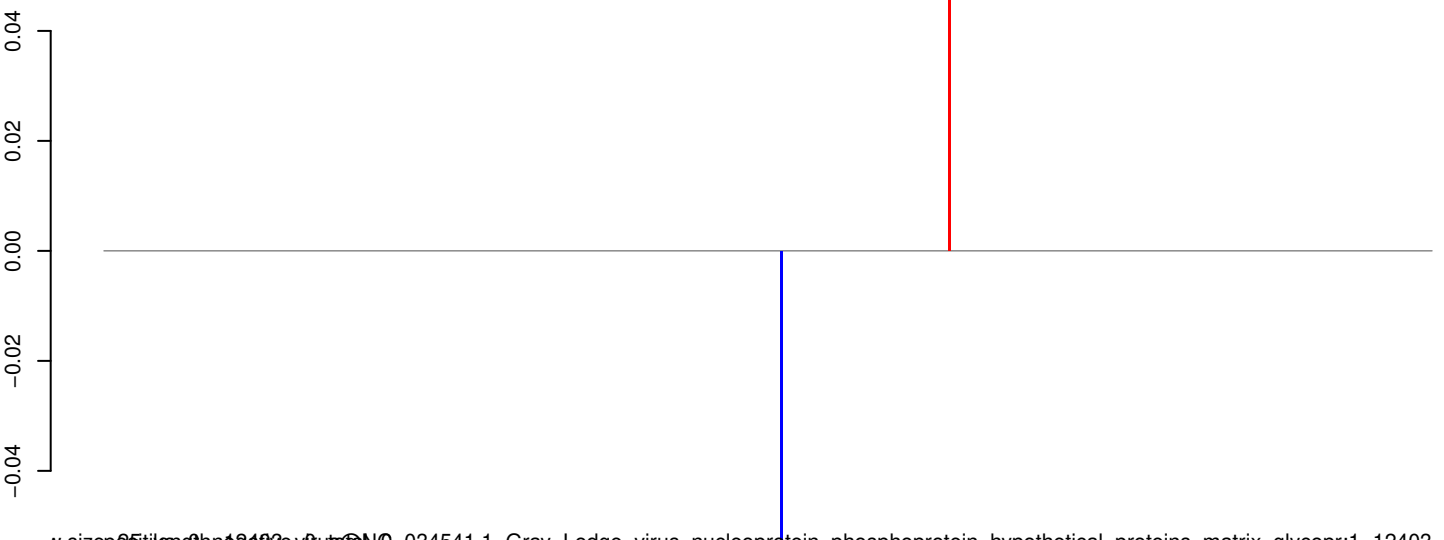
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



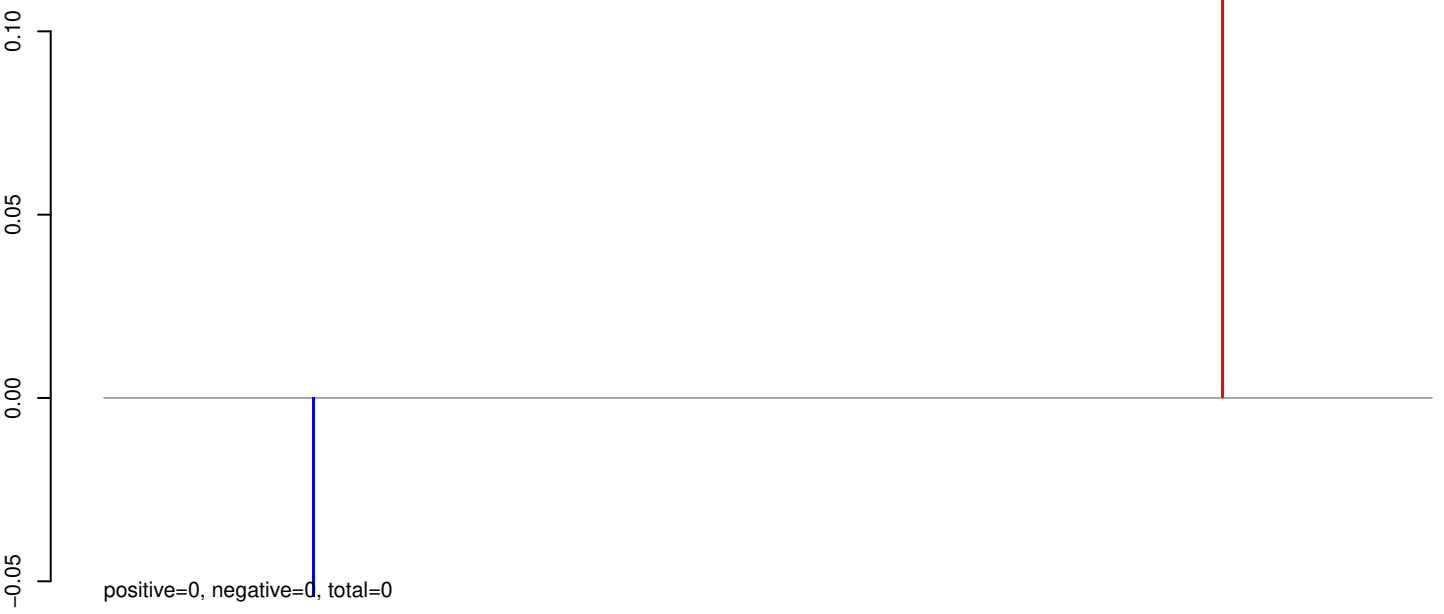
AeAeg_Aag2_BUNV1_ES.rep



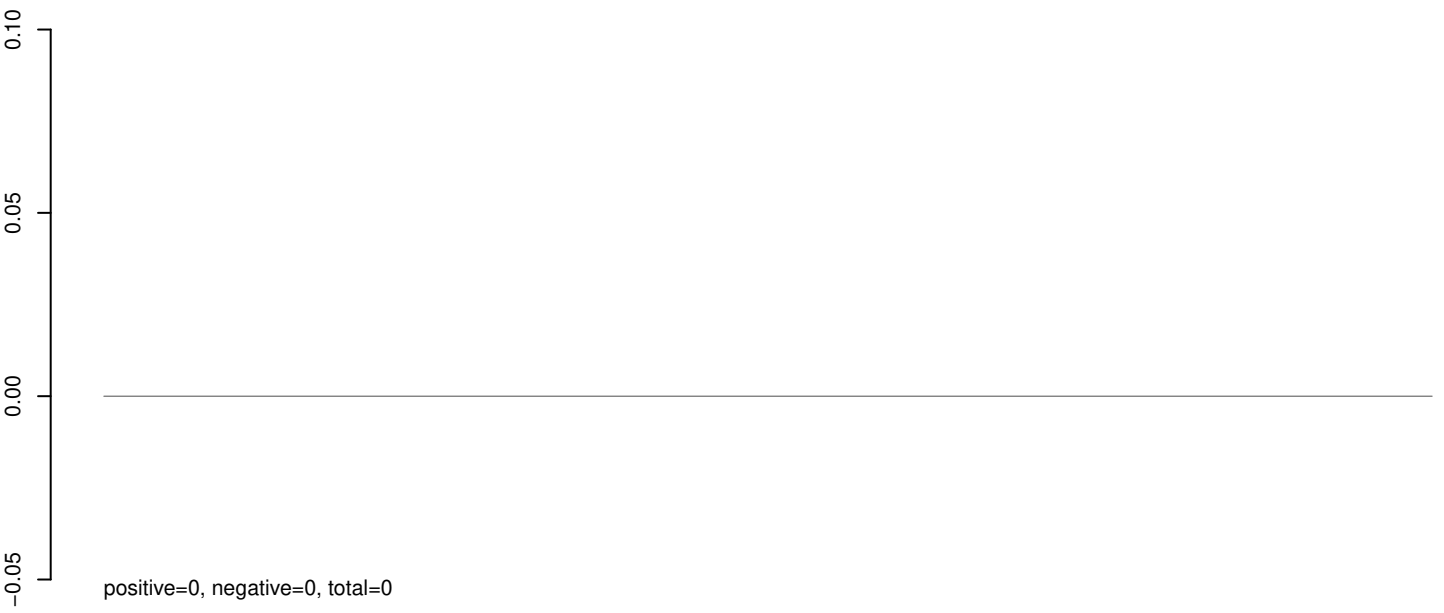
v size=25, len=11440, tuc=140, uc=140, 034541.1-Gray_Lodge_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopr:1-12403

0 2000 4000 6000 8000 10000 12000

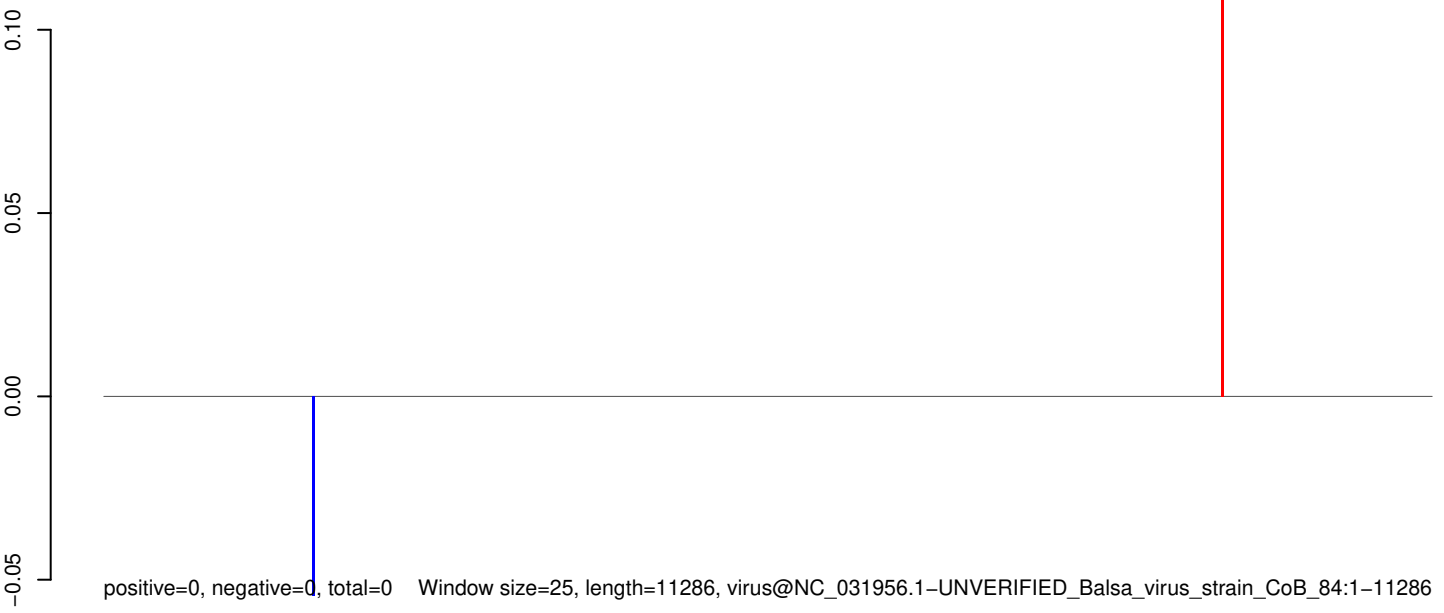
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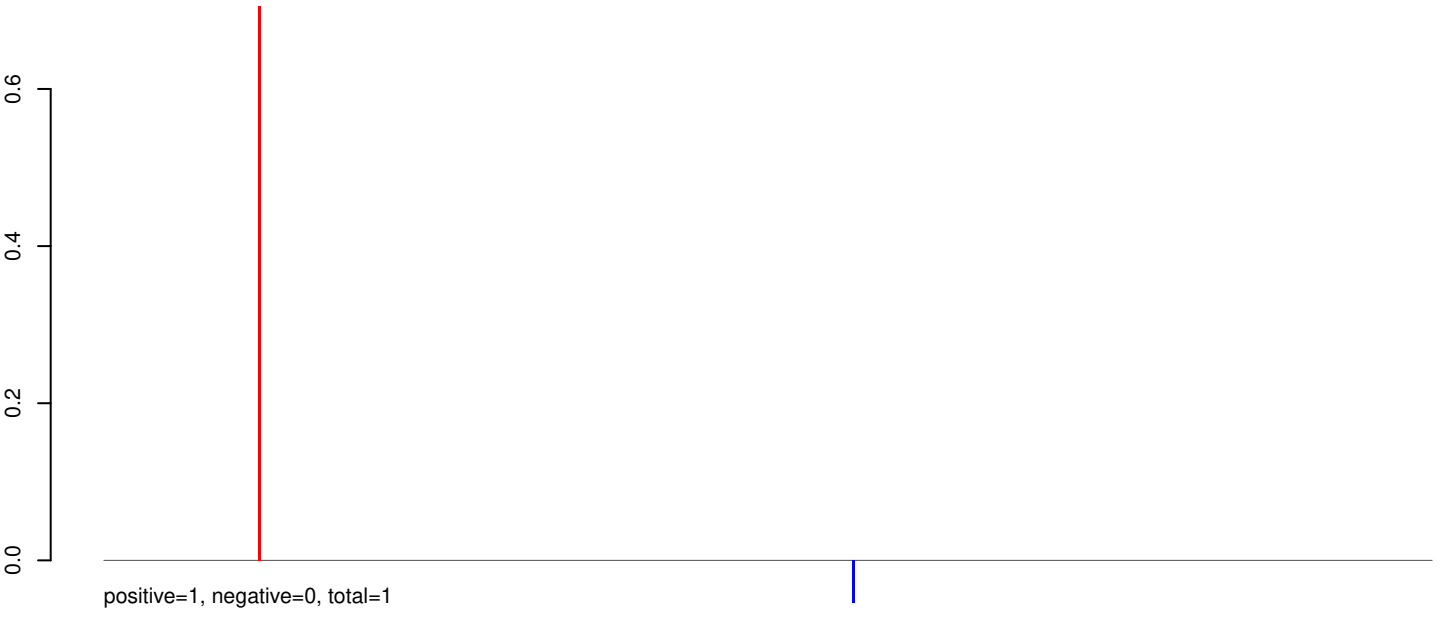
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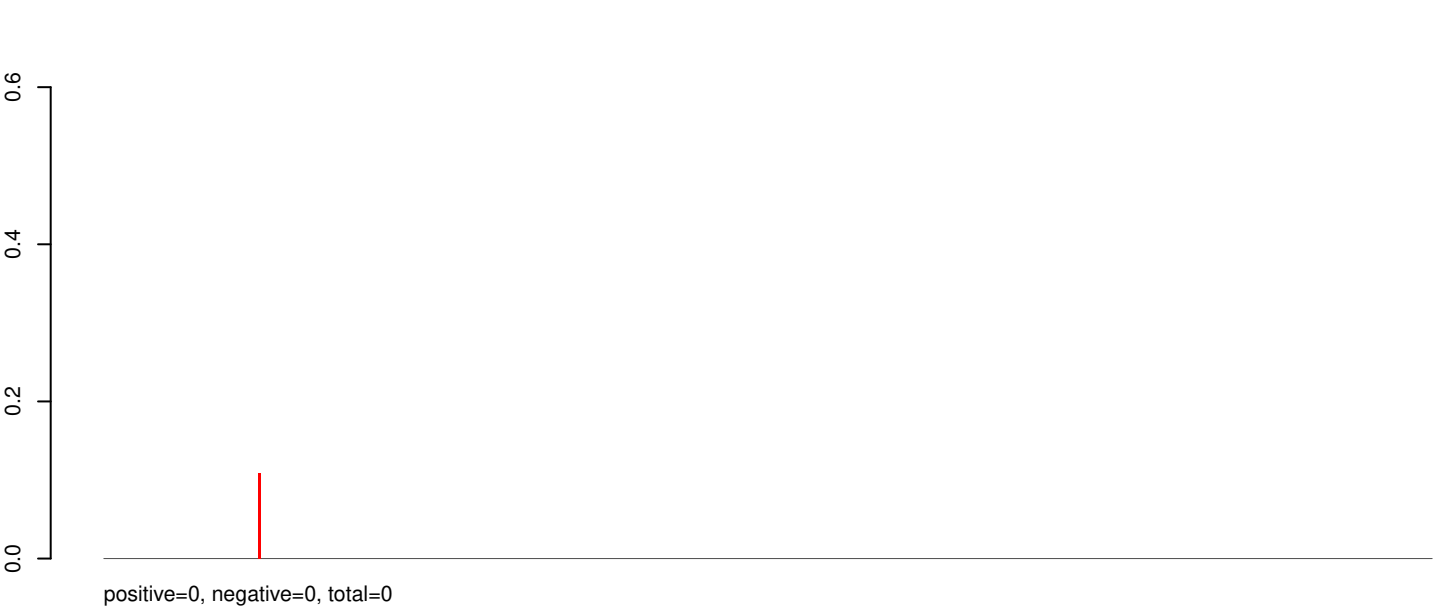
AeAeg_Aag2_BUNV1_ES.rep



AeAeg_Aag2_BUNV1_ES.18_23.rep



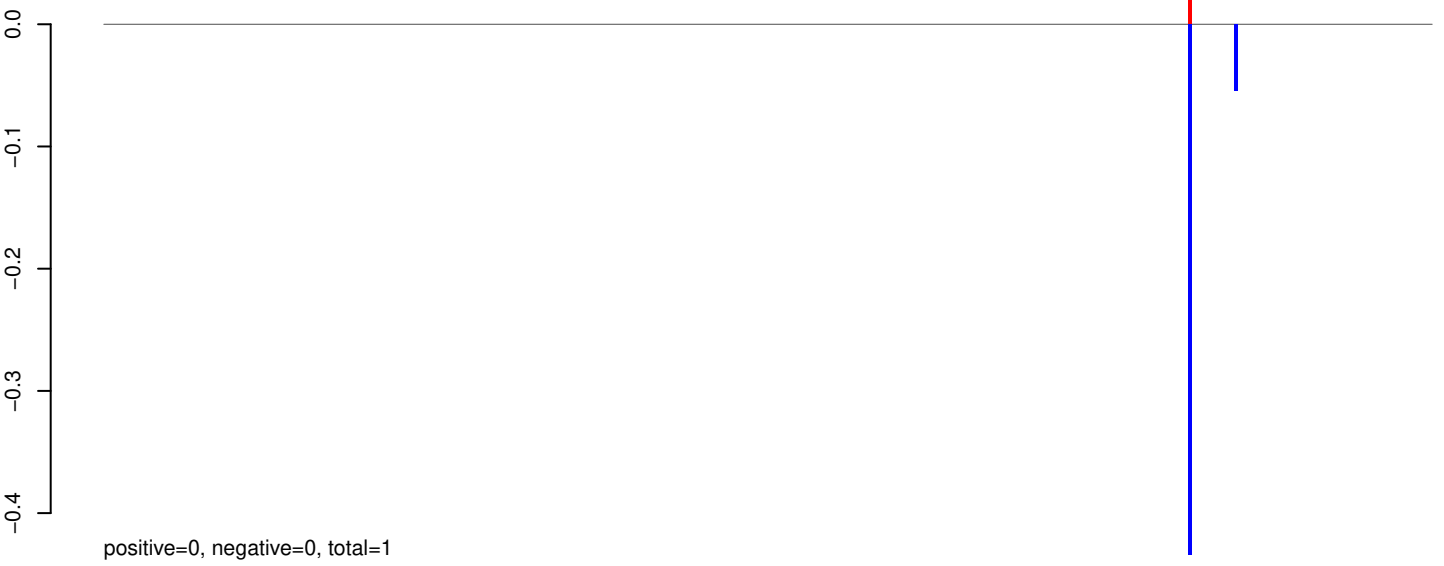
AeAeg_Aag2_BUNV1_ES.24_35.rep



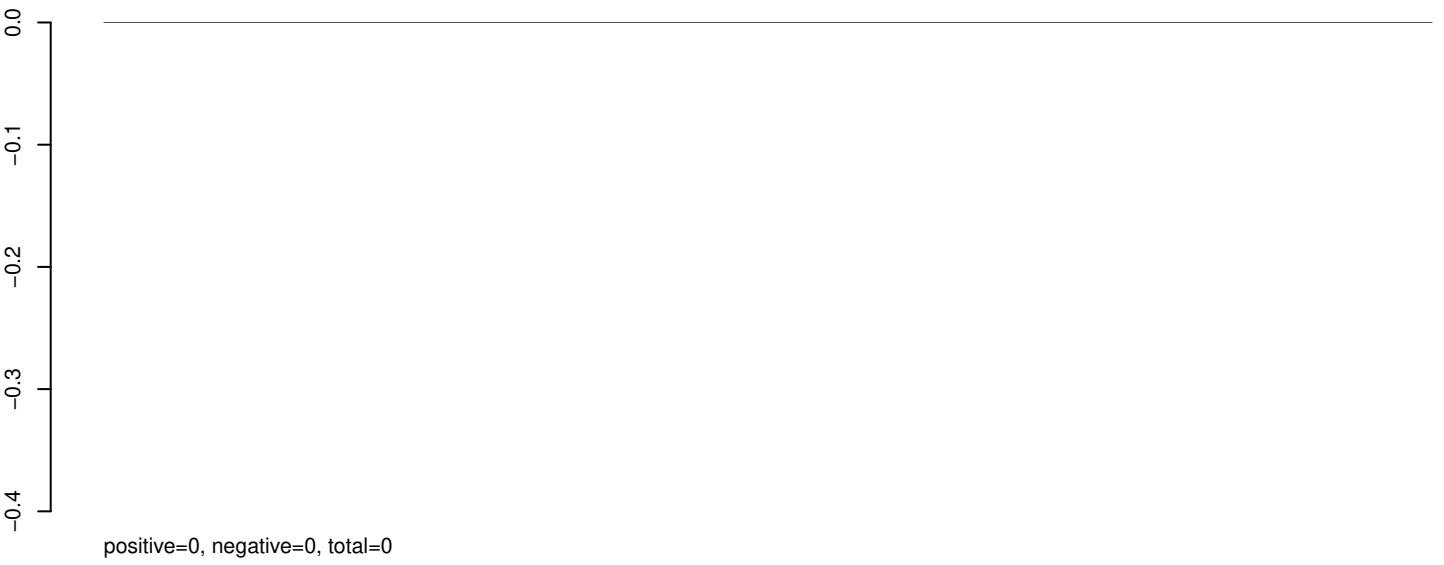
AeAeg_Aag2_BUNV1_ES.rep



AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



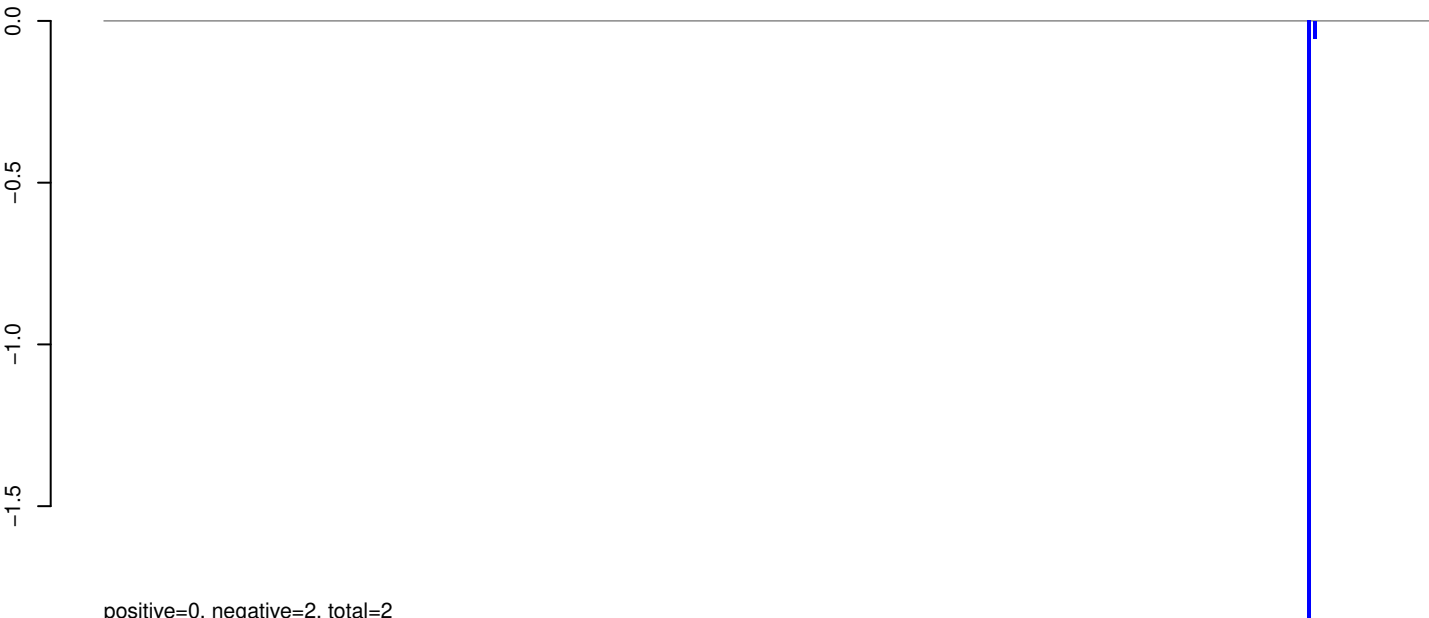
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Window size=25, length=10893, virus@NC_027817.1-Parramatta_River_virus_isolate_92-B115745:1-10893

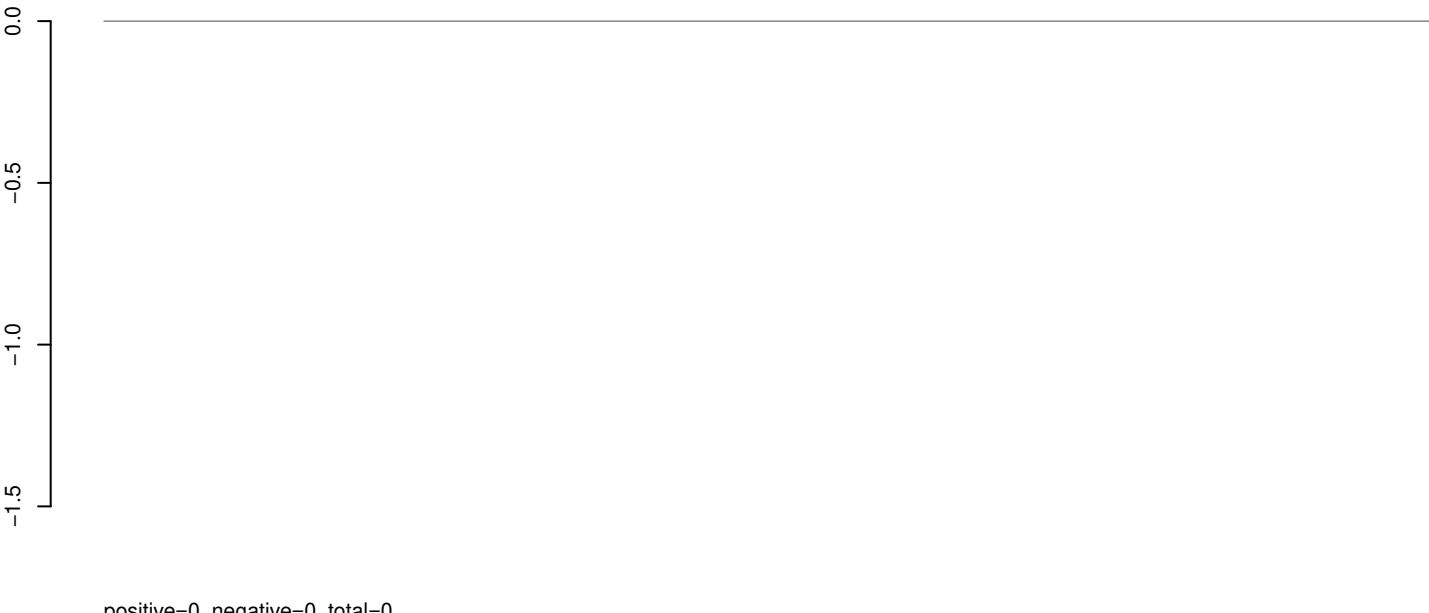
0 2000 4000 6000 8000 10000

AeAeg_Aag2_BUNV1_ES.18_23.rep



positive=0, negative=2, total=2

AeAeg_Aag2_BUNV1_ES.24_35.rep



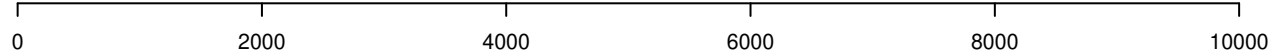
positive=0, negative=0, total=0

AeAeg_Aag2_BUNV1_ES.rep

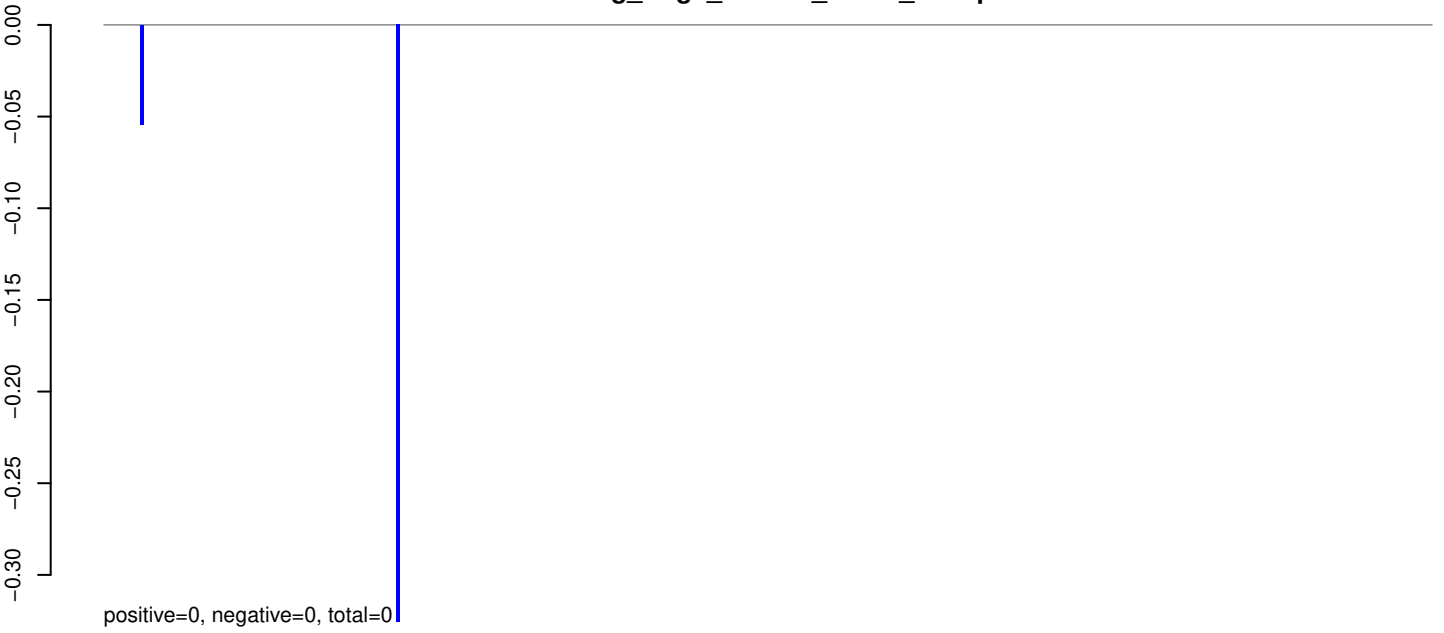


positive=0, negative=2, total=2

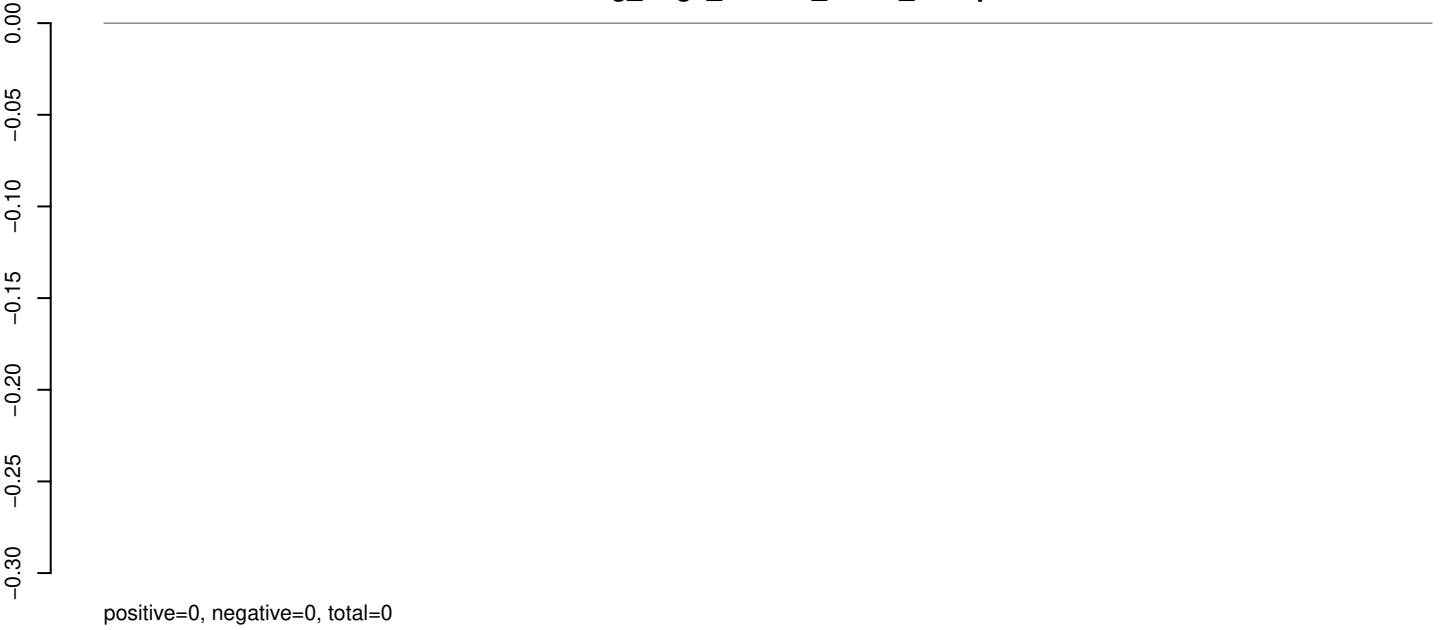
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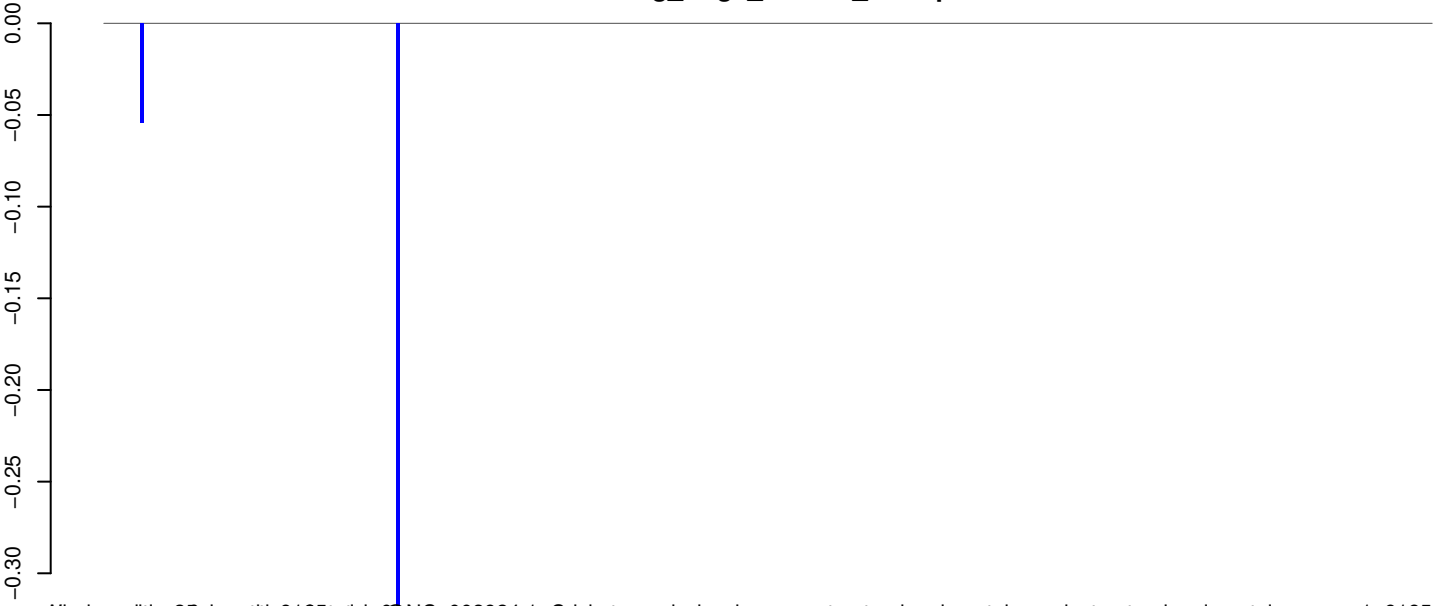
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



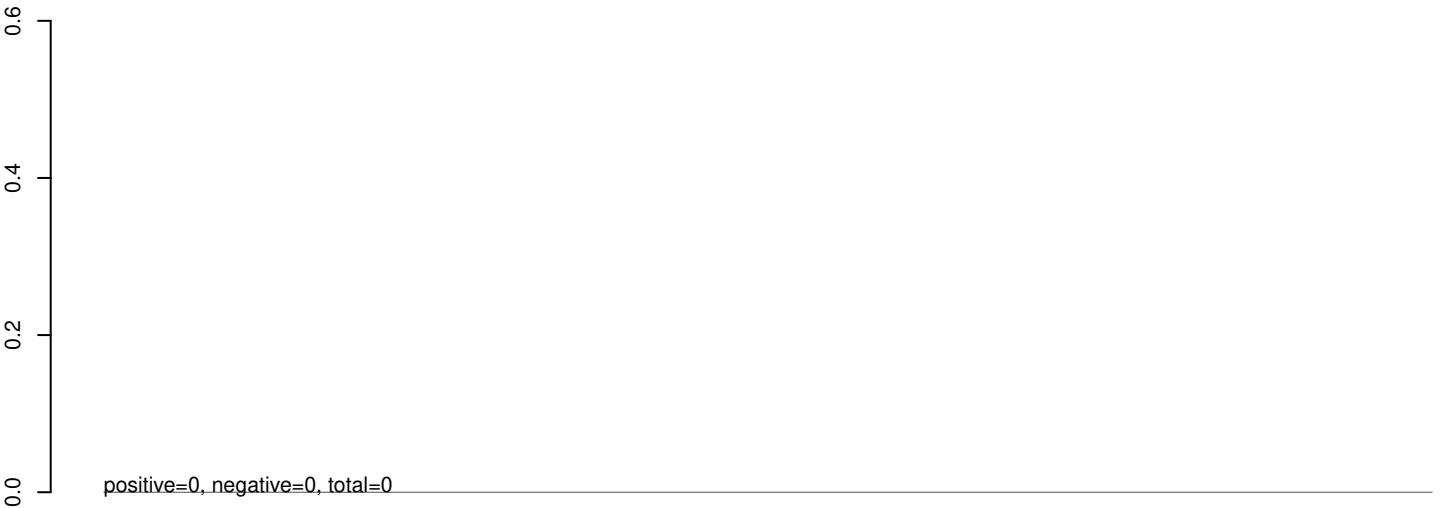
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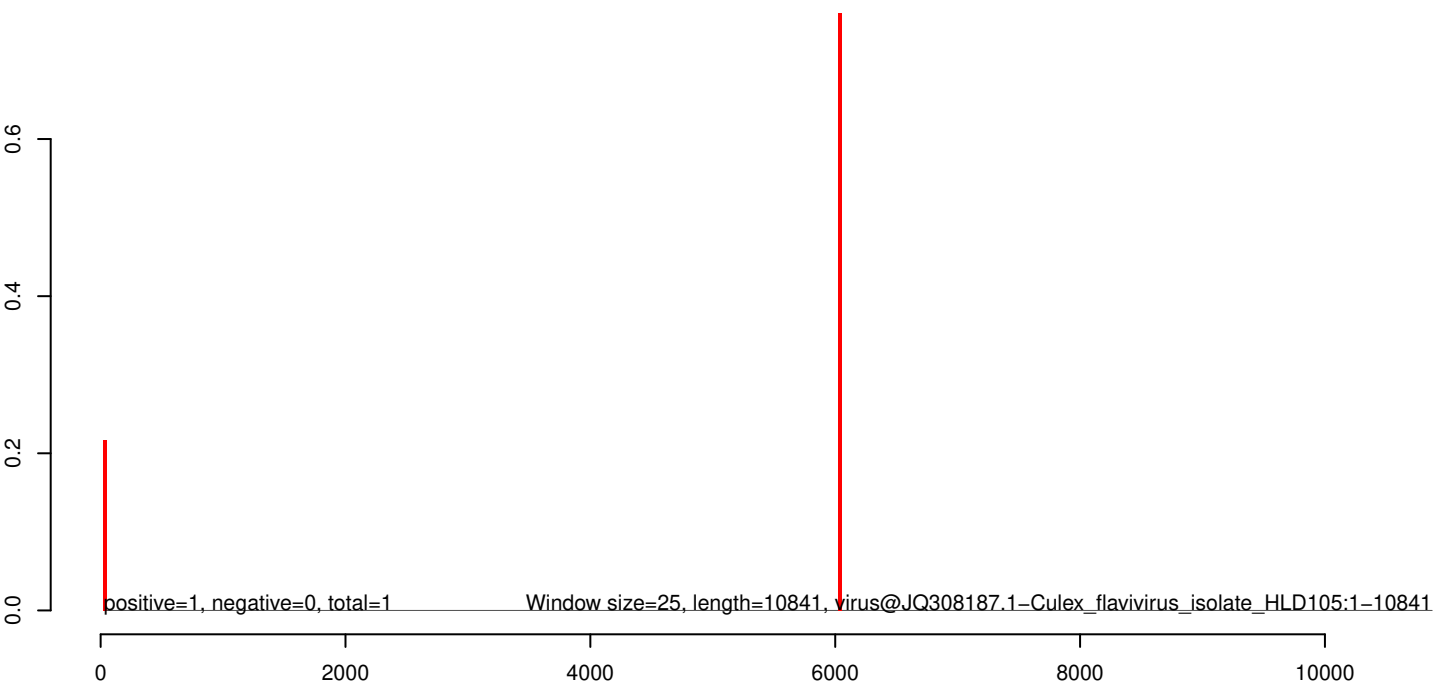
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AeAeg_Aag2_BUNV1_ES.24_35.rep



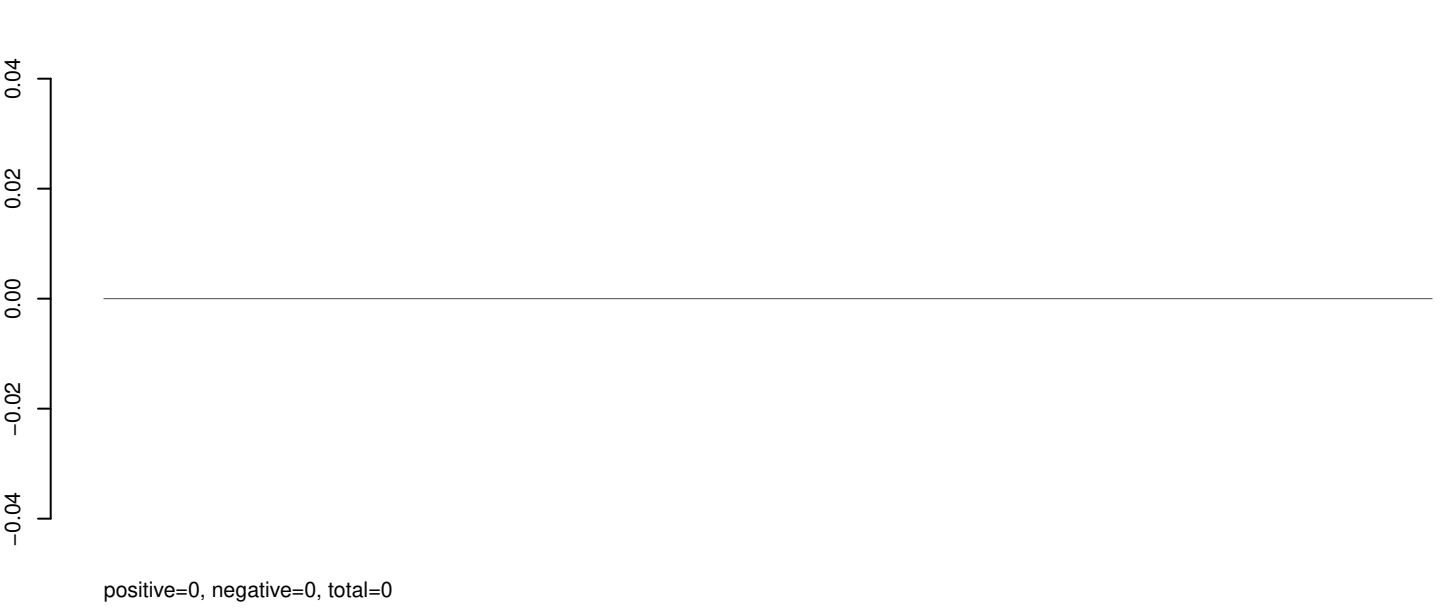
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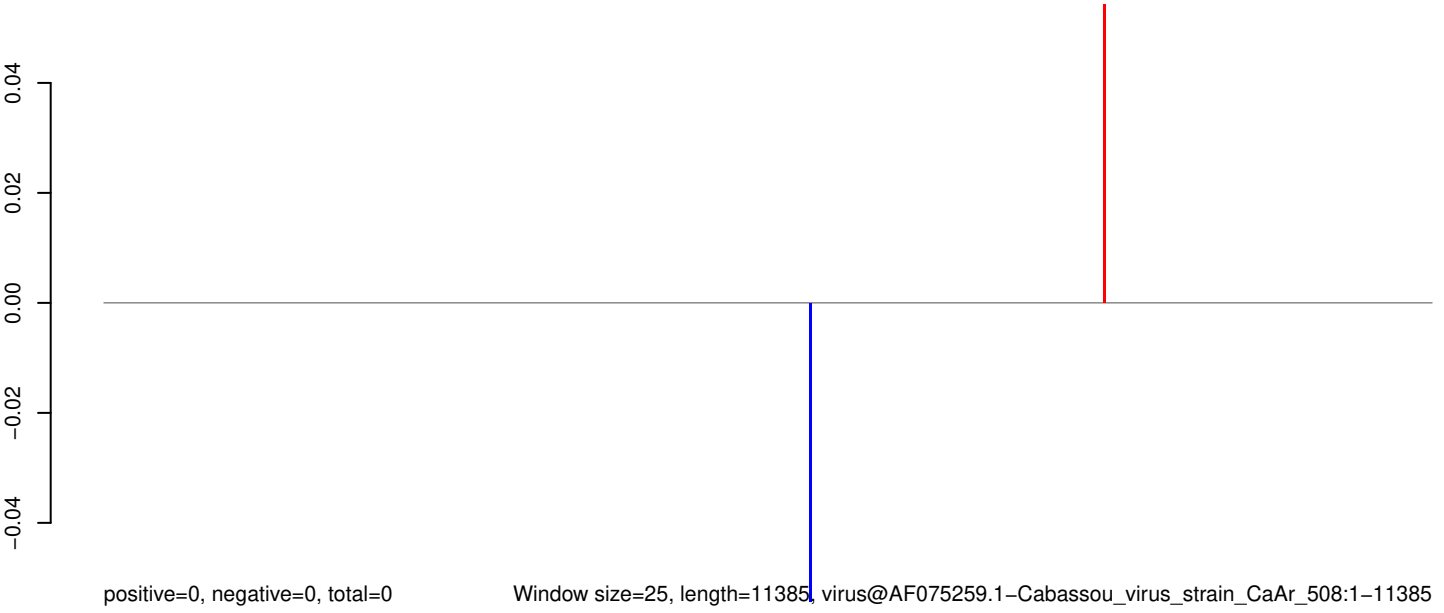
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AeAeg_Aag2_BUNV1_ES.24_35.rep

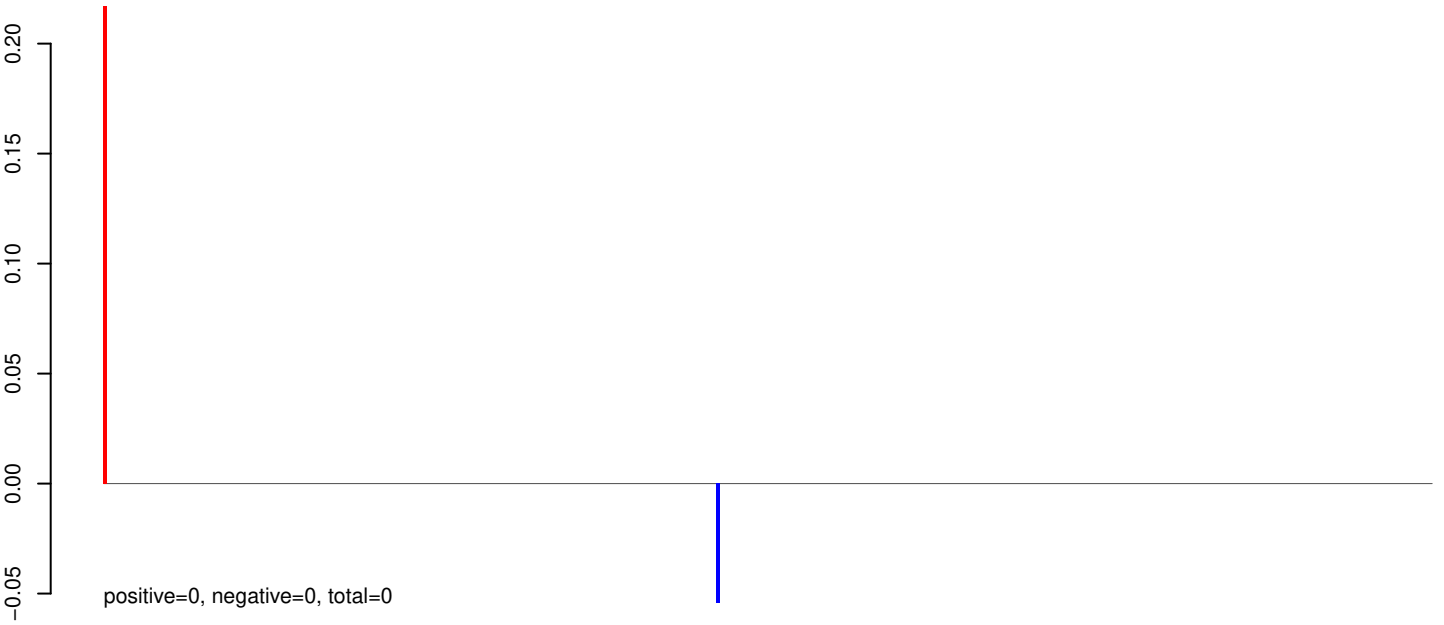


AeAeg_Aag2_BUNV1_ES.rep

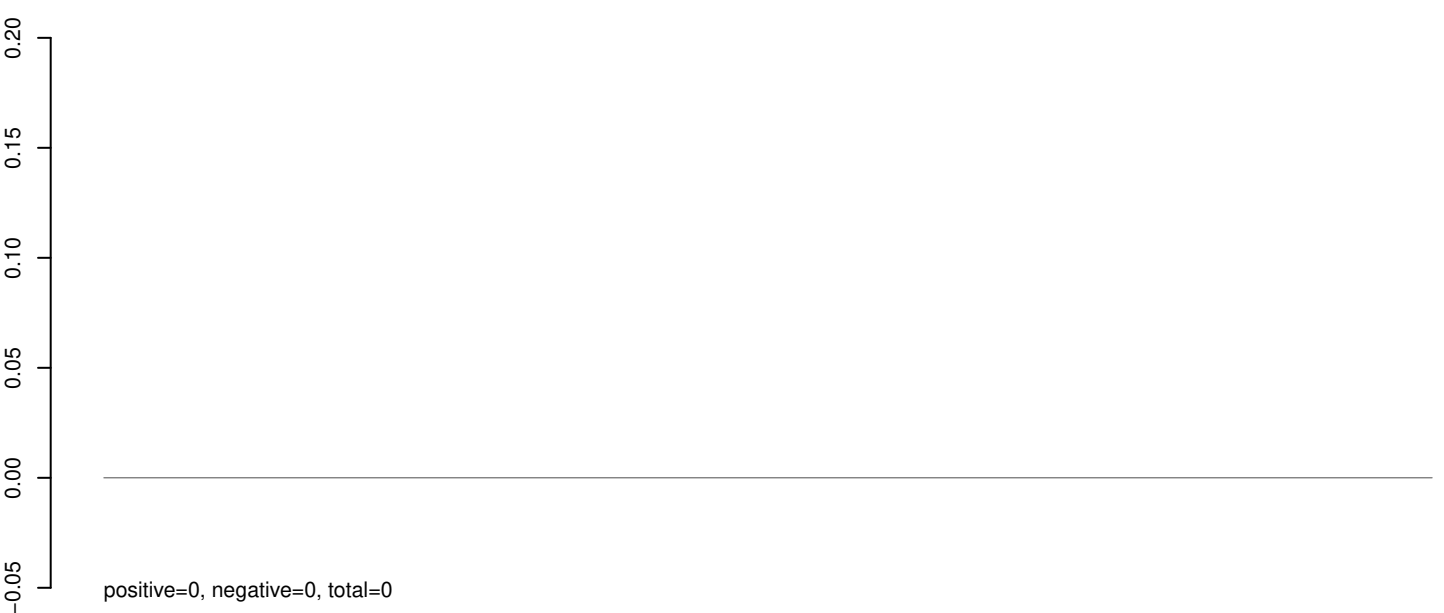


0 2000 4000 6000 8000 10000

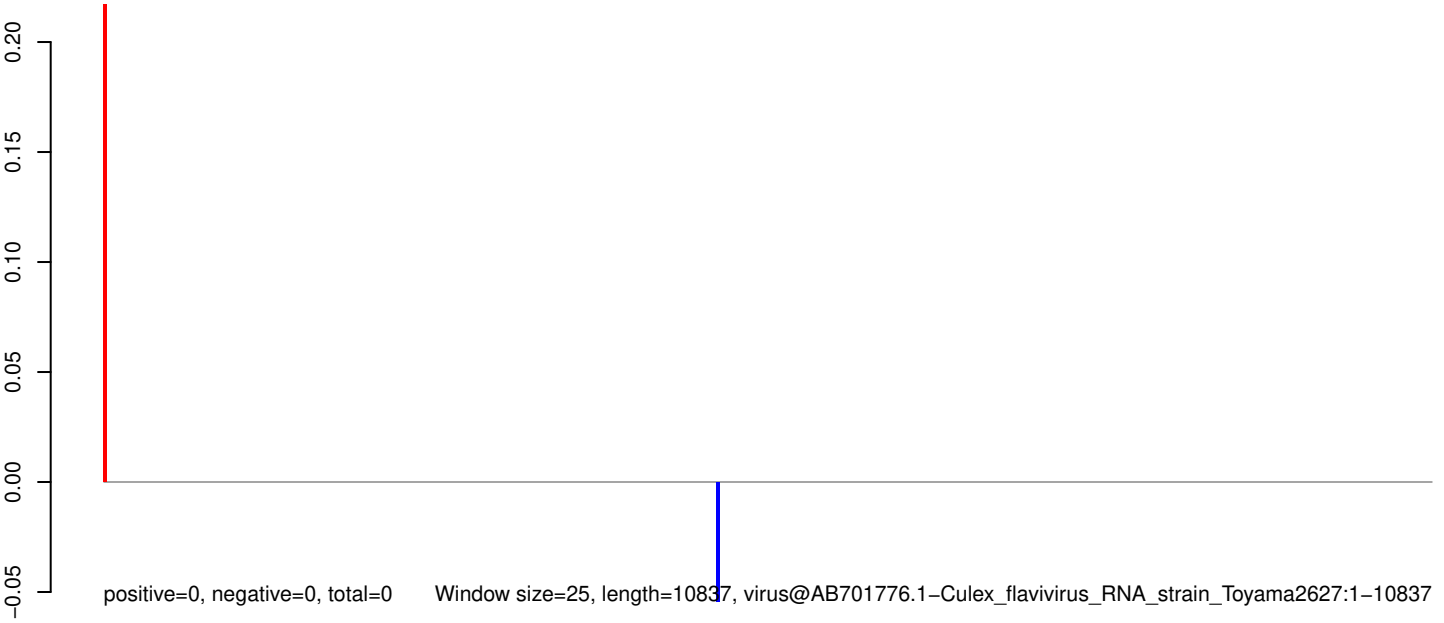
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep

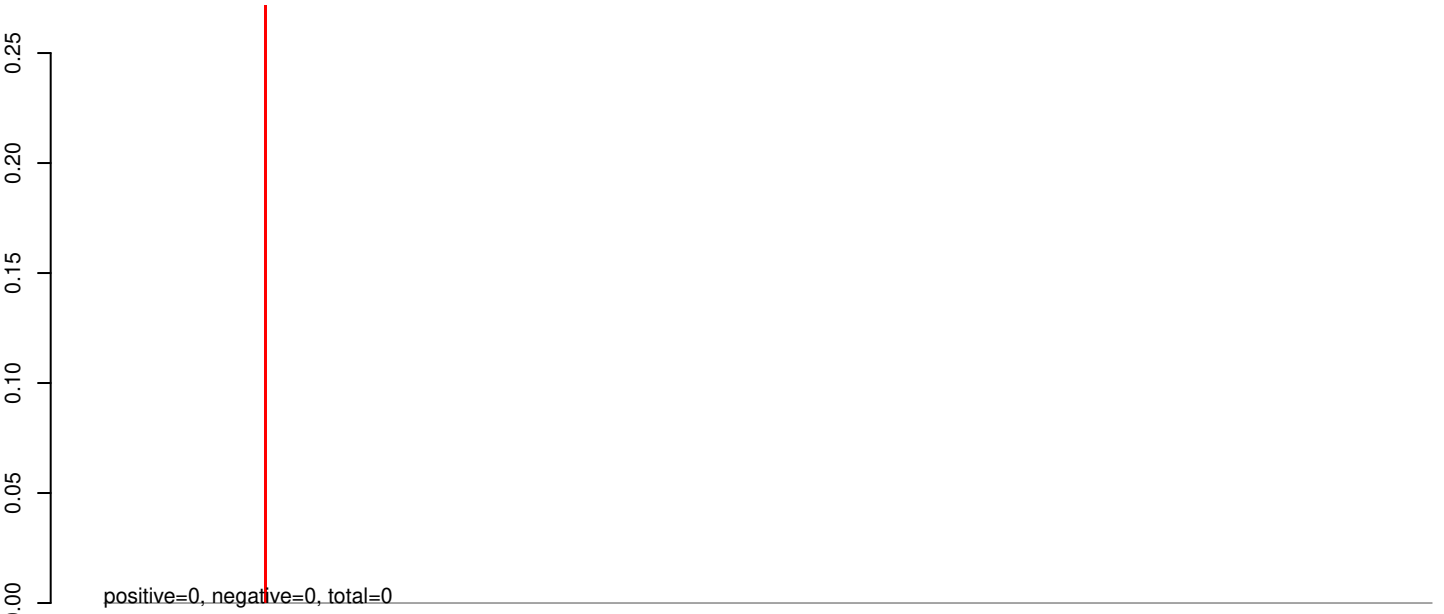


AeAeg_Aag2_BUNV1_ES.rep

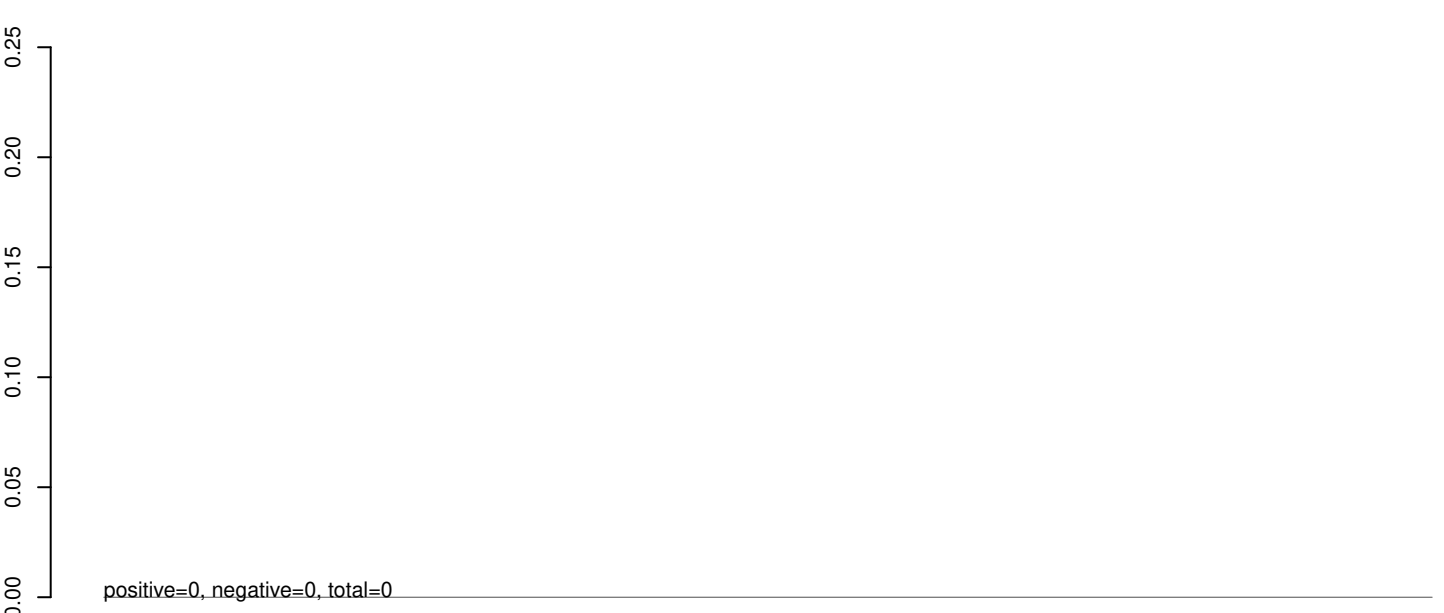


0 2000 4000 6000 8000 10000

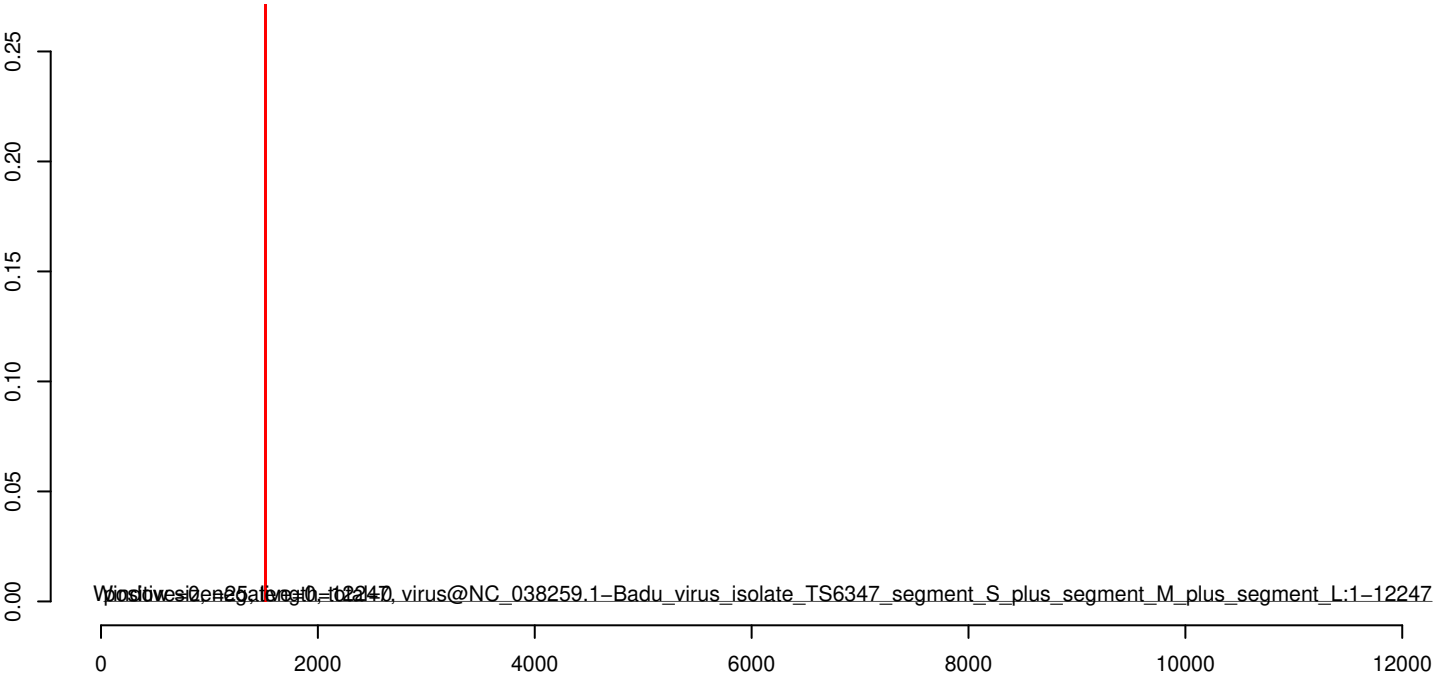
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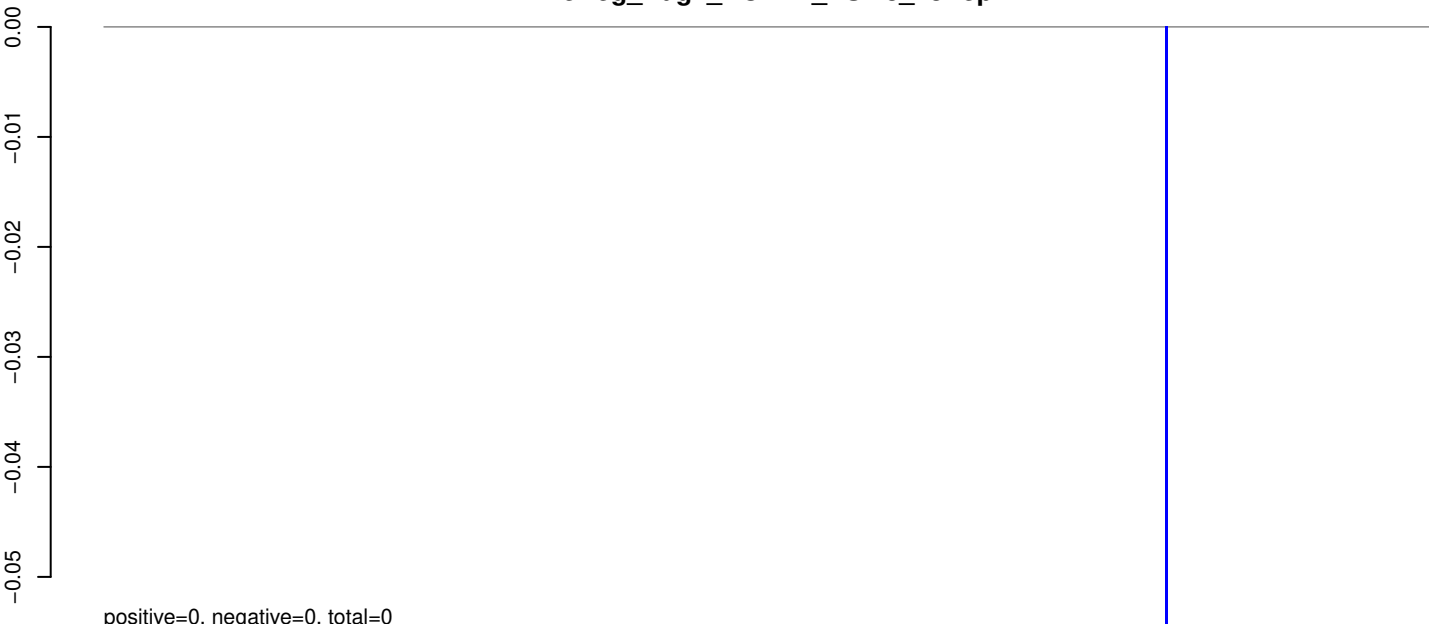
AeAeg_Aag2_BUNV1_ES.24_35.rep



AeAeg_Aag2_BUNV1_ES.rep

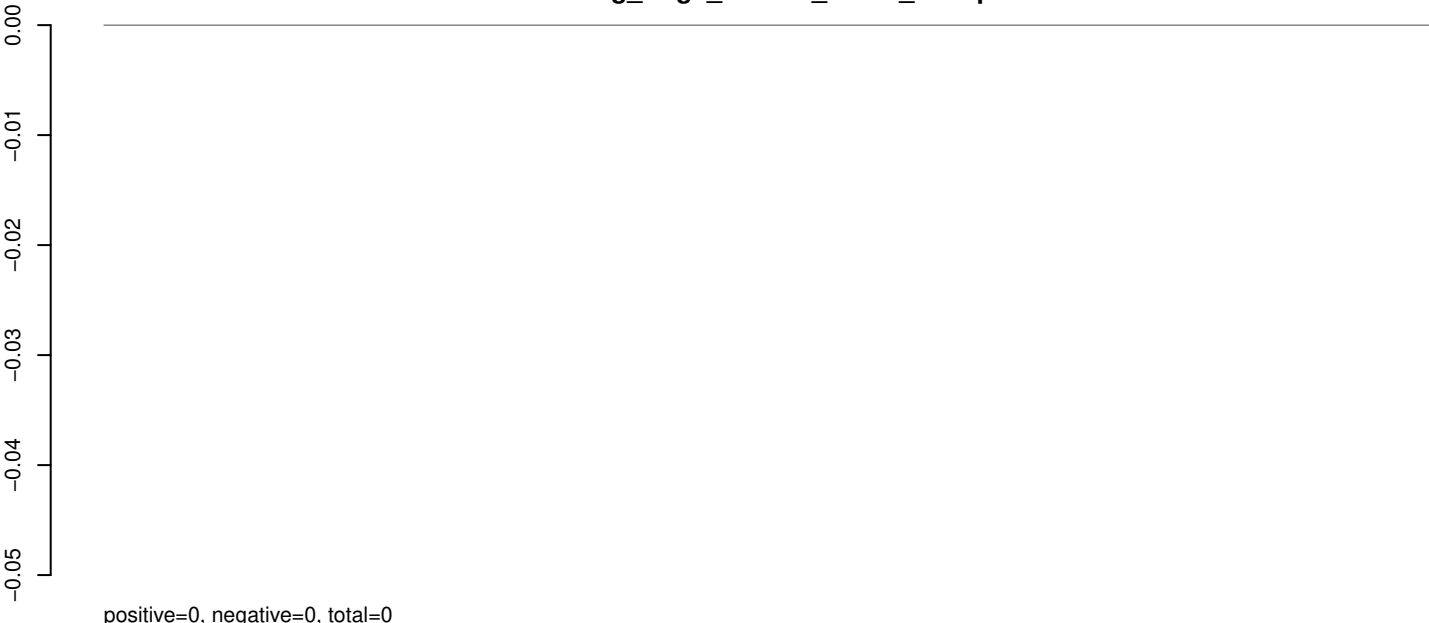


AeAeg_Aag2_BUNV1_ES.18_23.rep



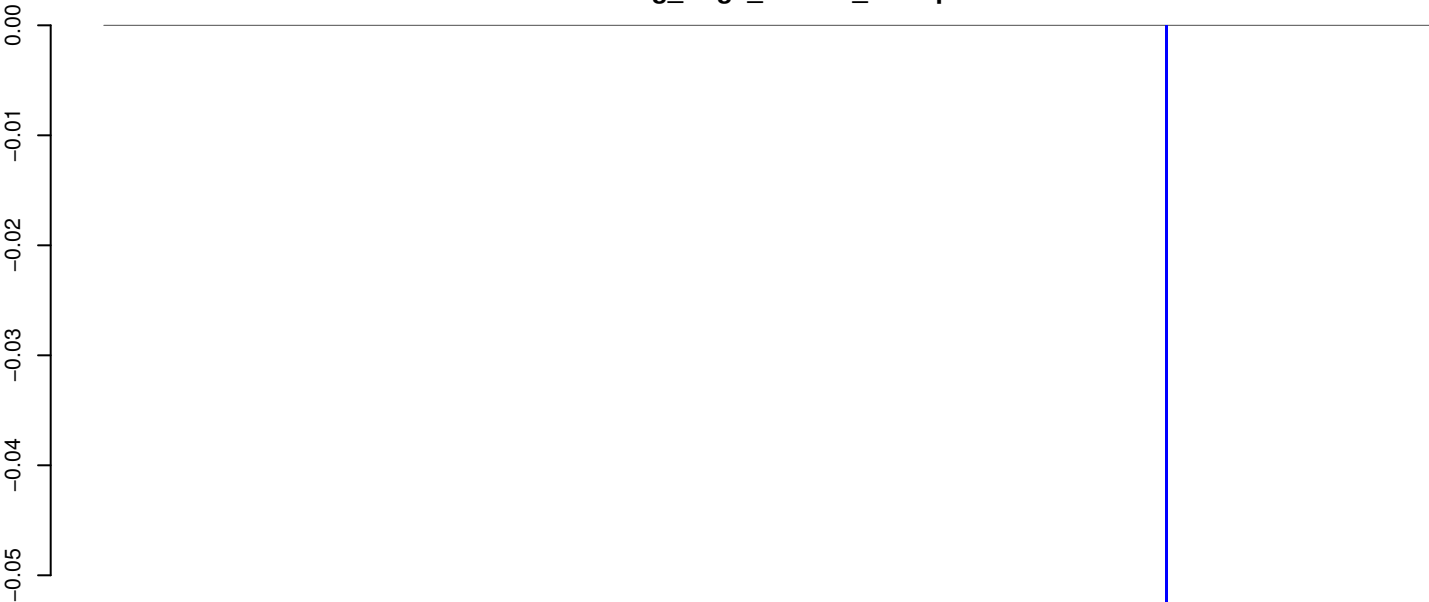
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AeAeg_Aag2_BUNV1_ES.24_35.rep



positive=0, negative=0, total=0

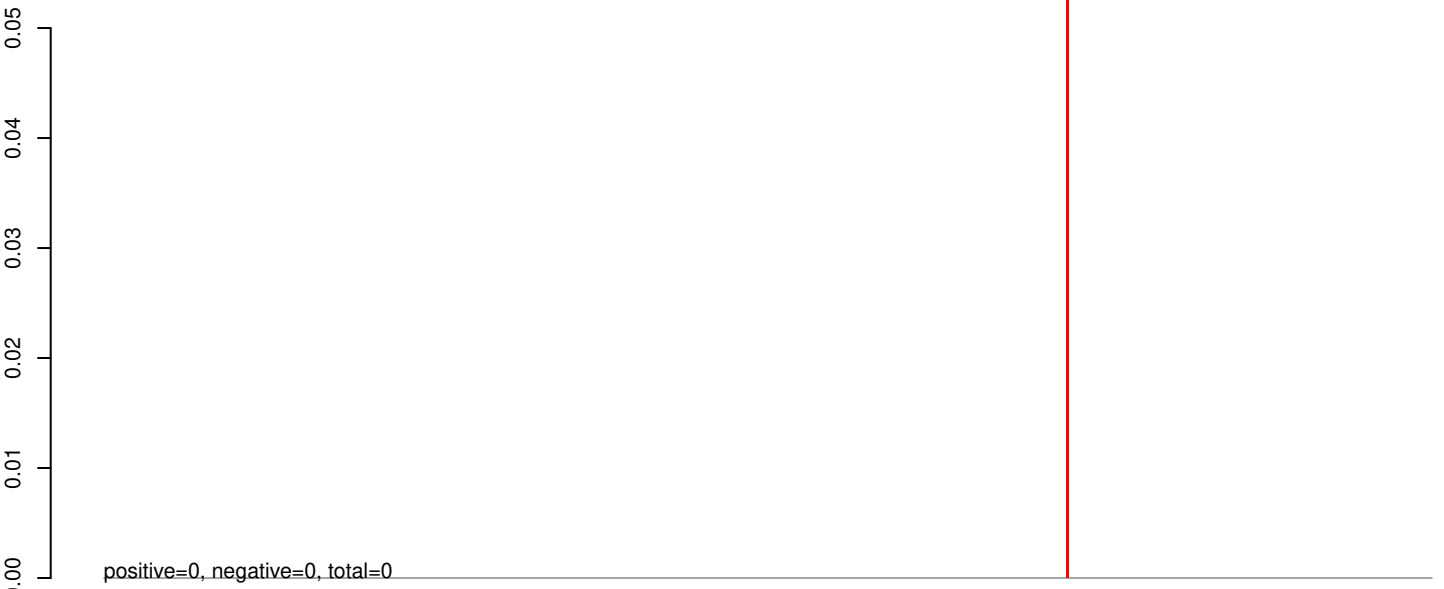
AeAeg_Aag2_BUNV1_ES.rep



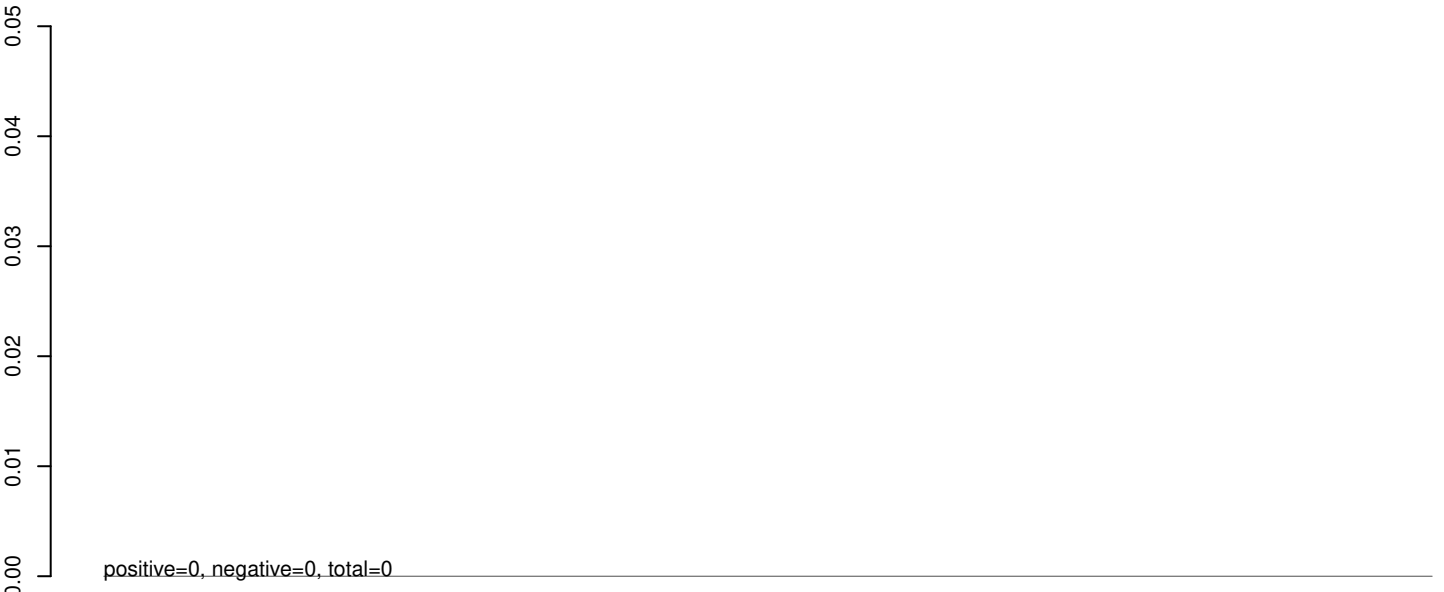
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0 2000 4000 6000 8000 10000 12000

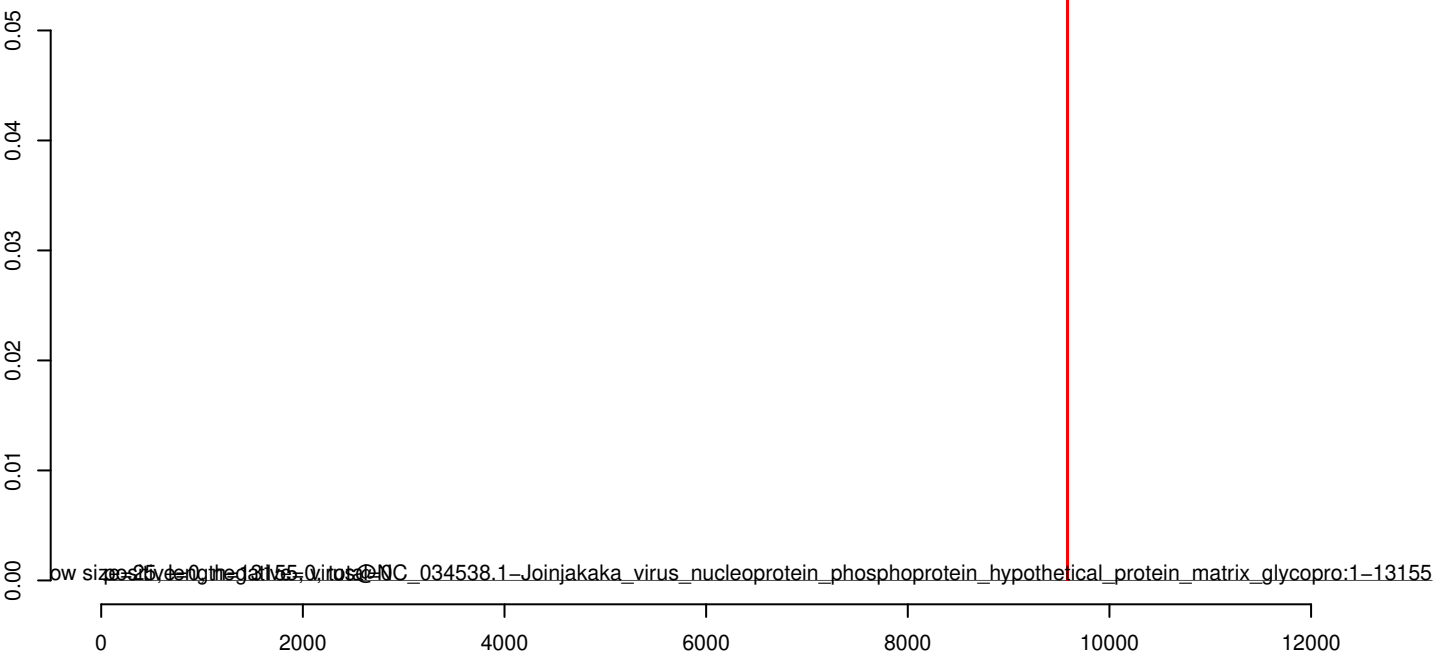
AeAeg_Aag2_BUNV1_ES.18_23.rep



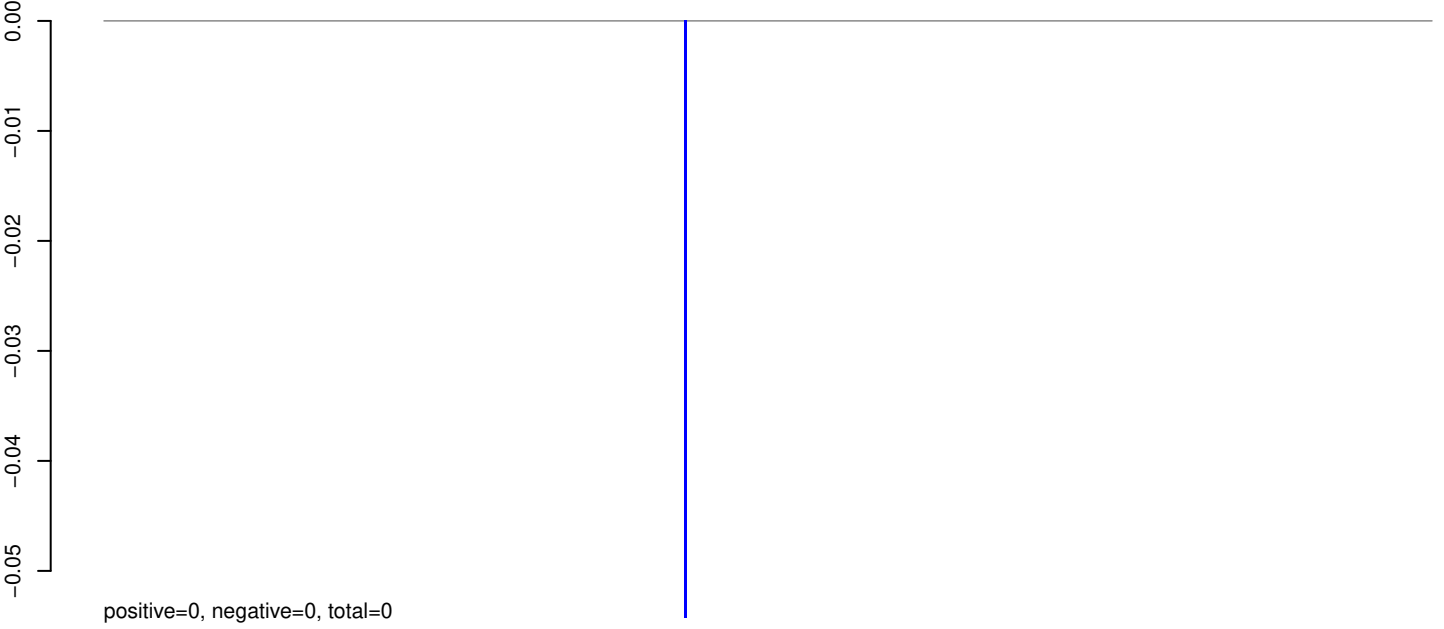
AeAeg_Aag2_BUNV1_ES.24_35.rep



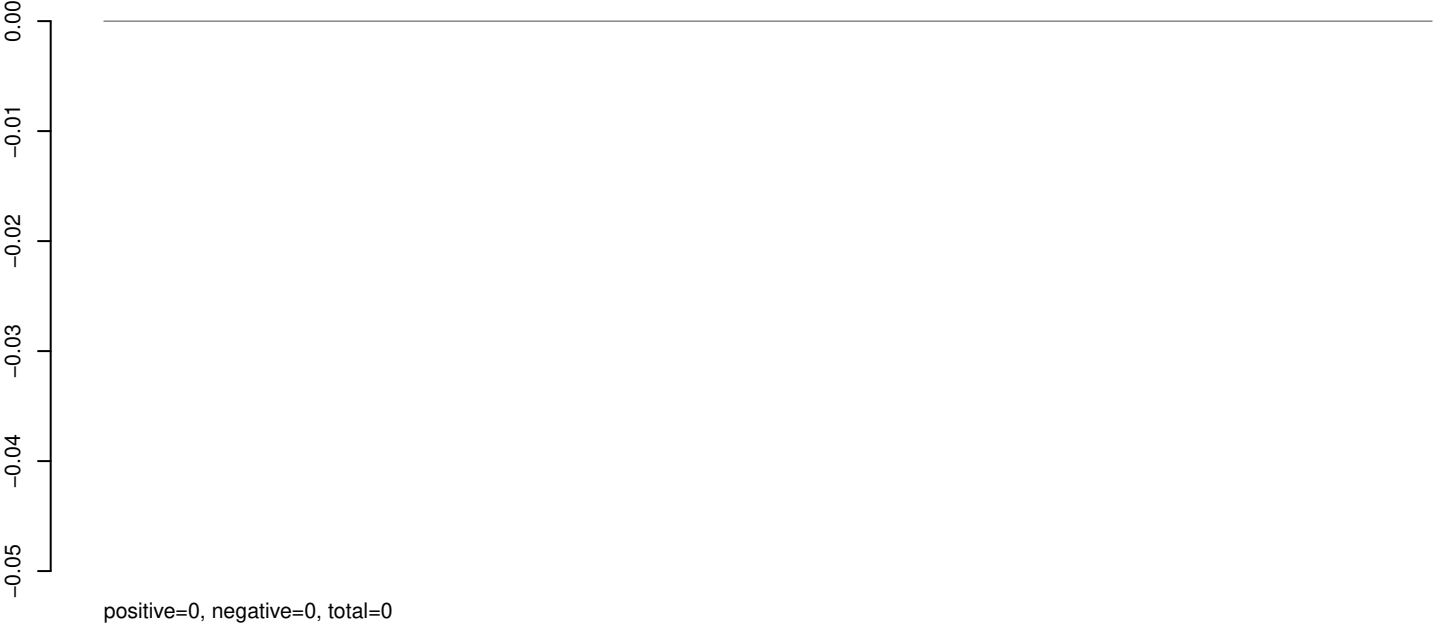
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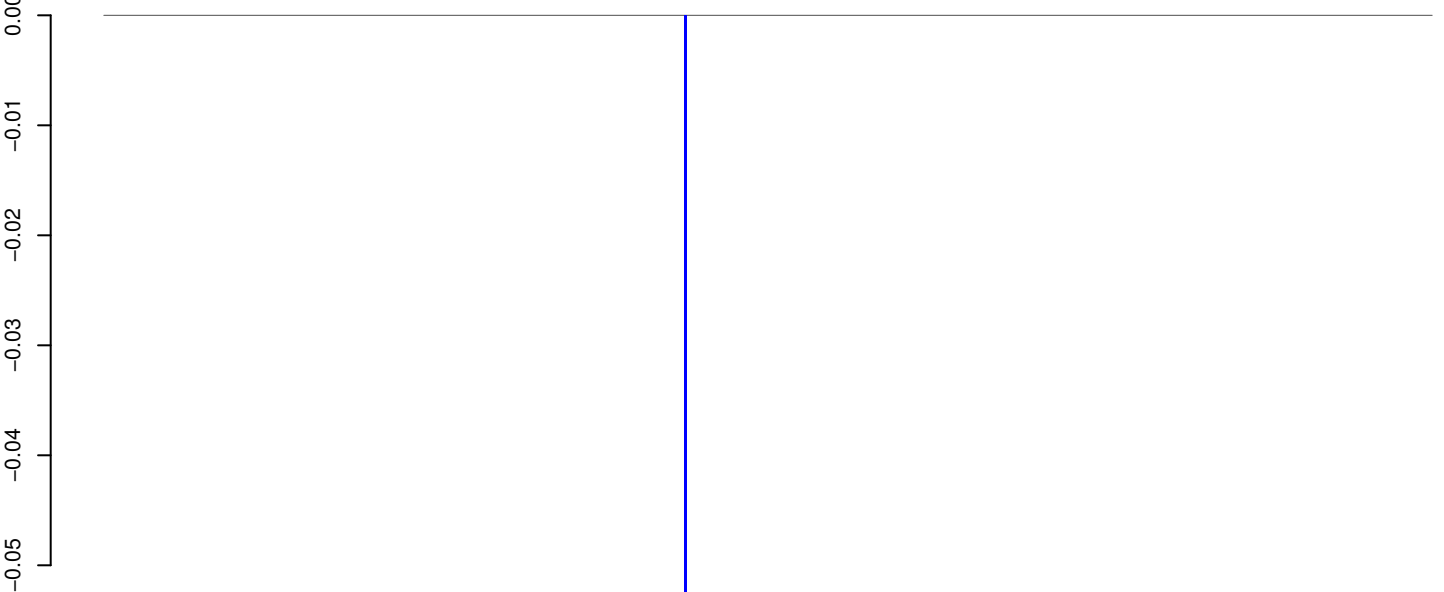
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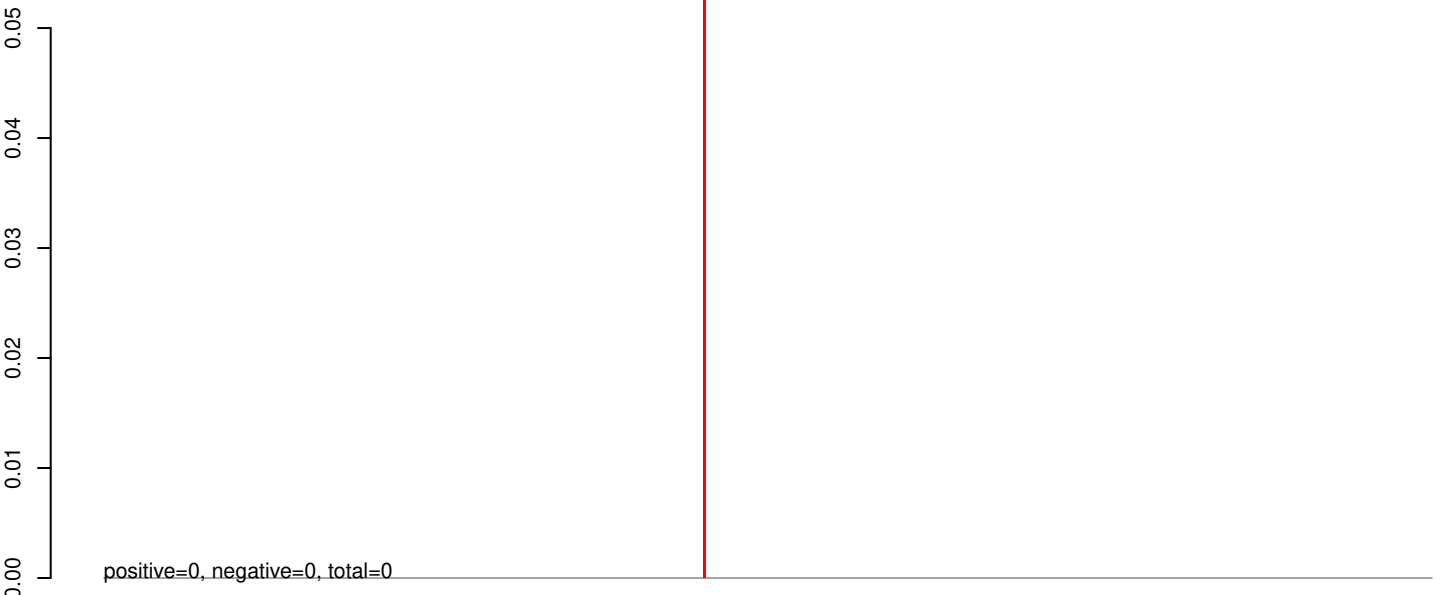
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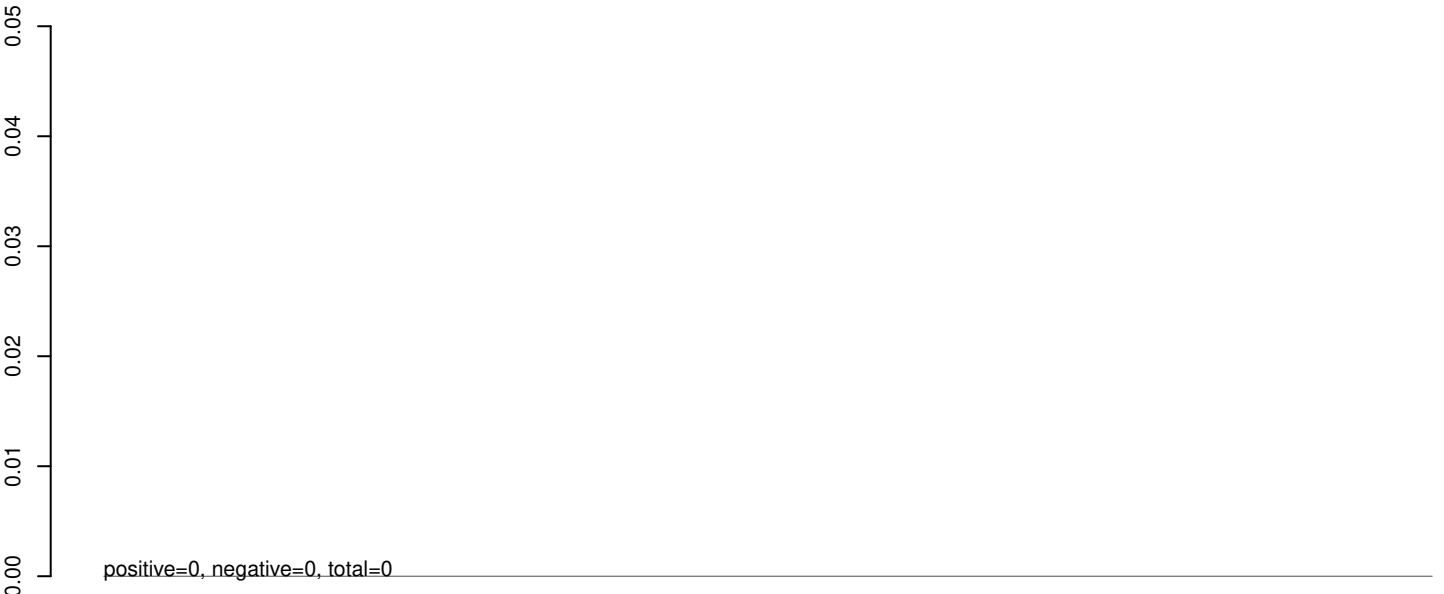
AeAeg_Aag2_BUNV1_ES.rep



AeAeg_Aag2_BUNV1_ES.18_23.rep



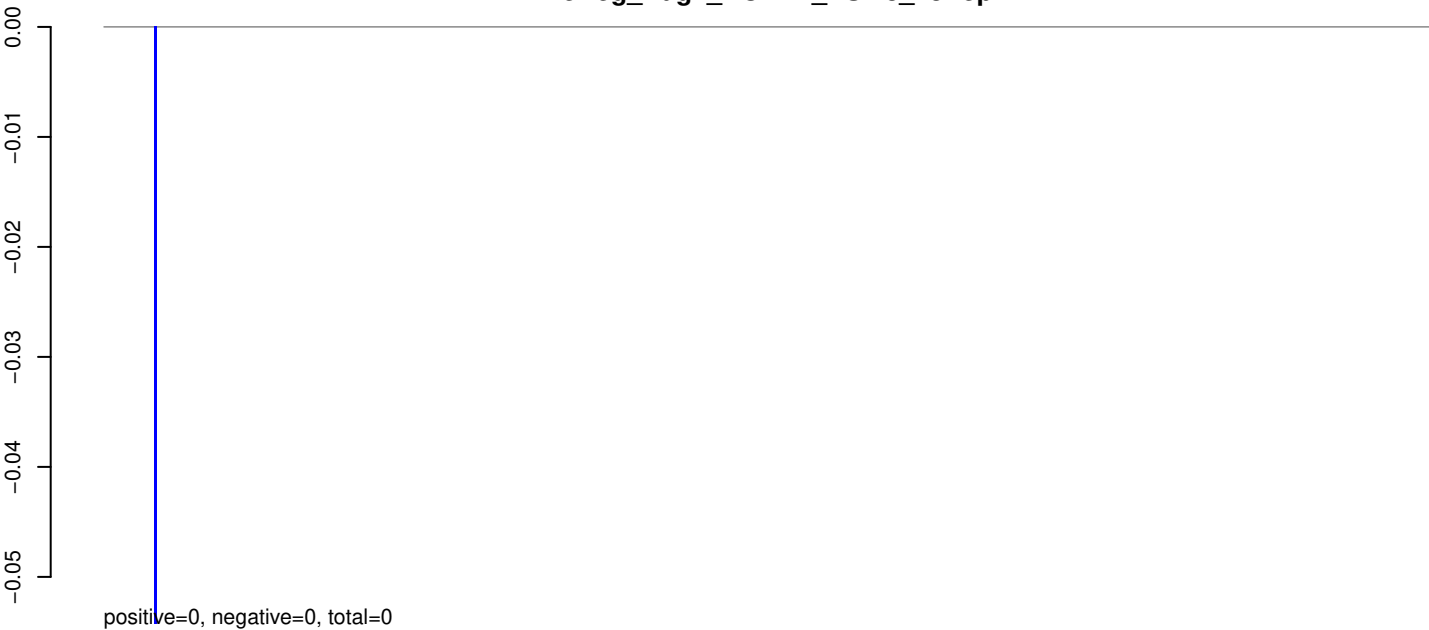
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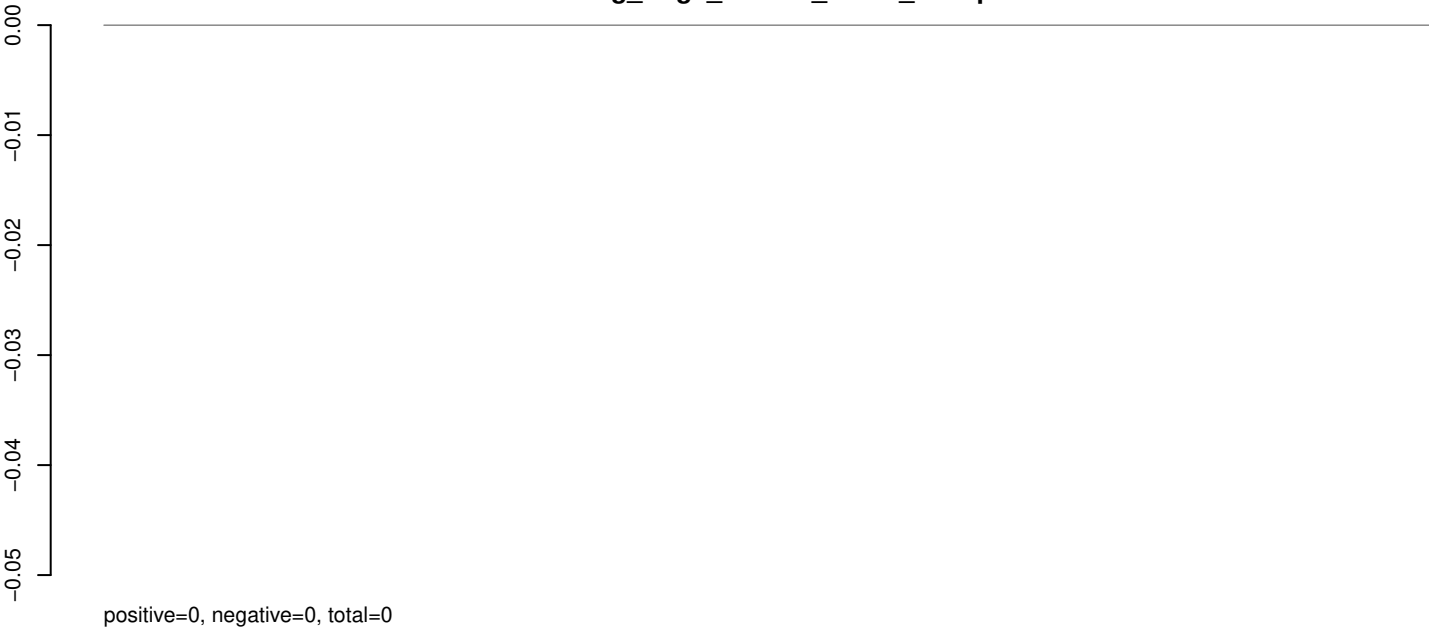
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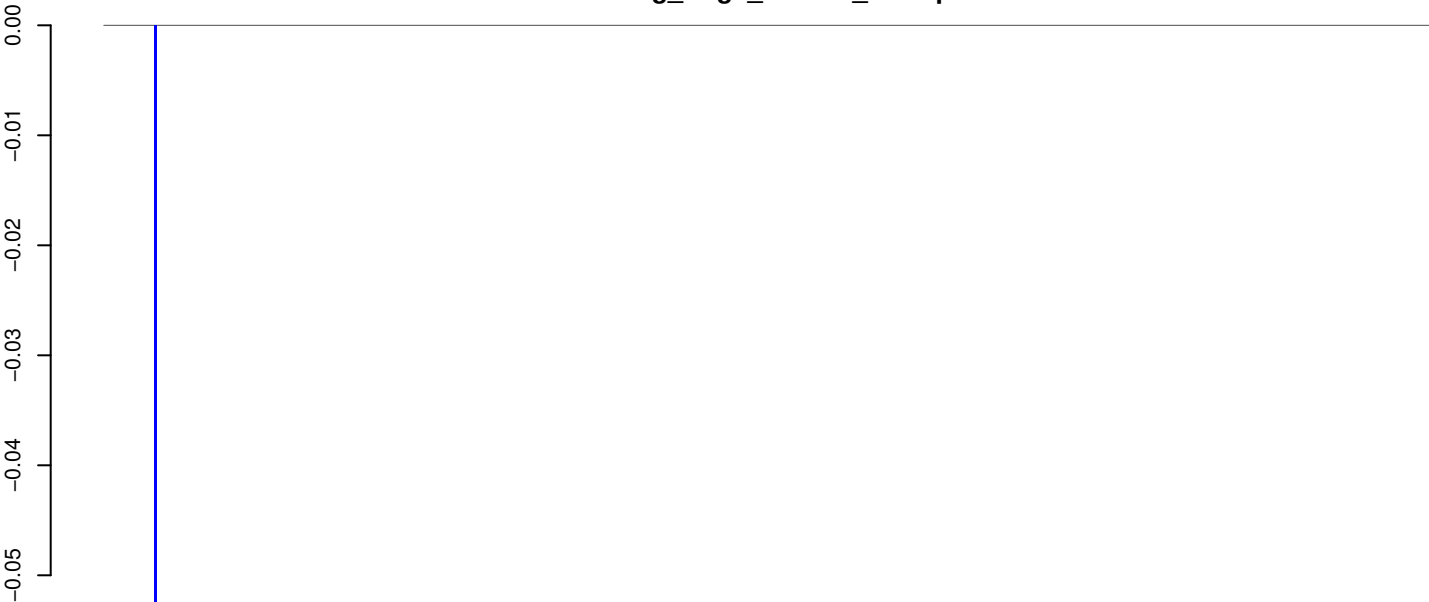
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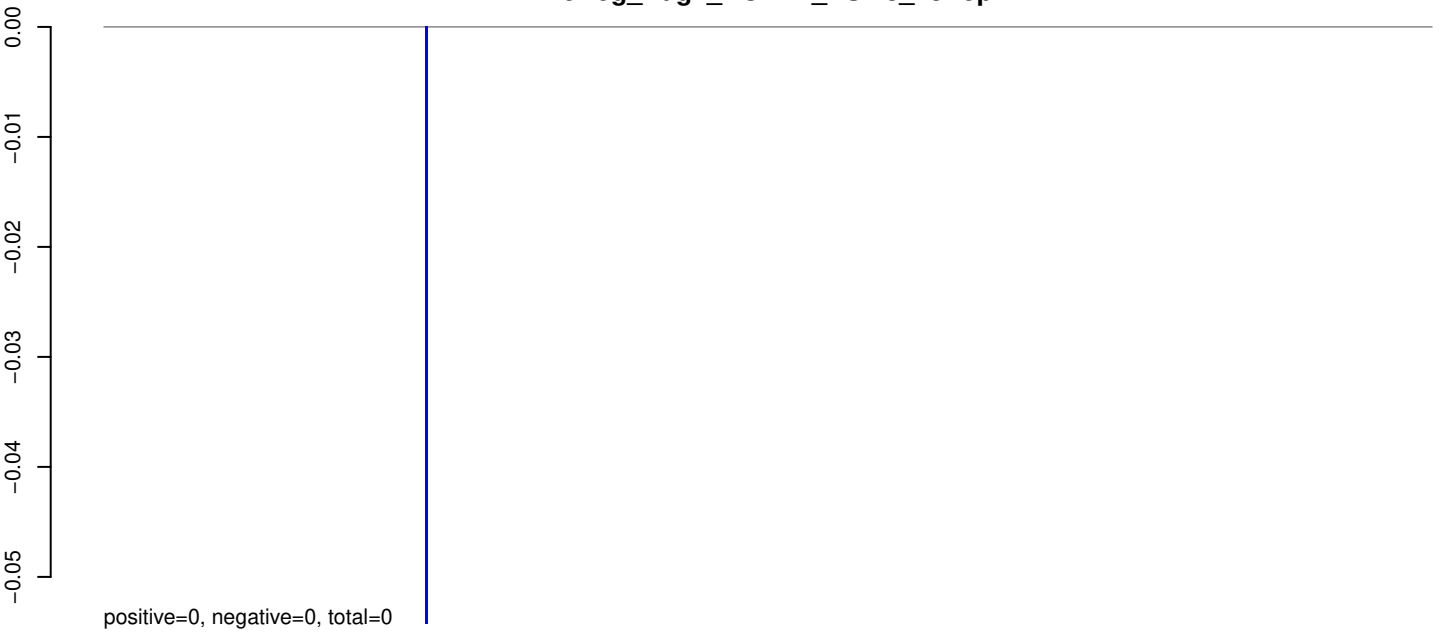
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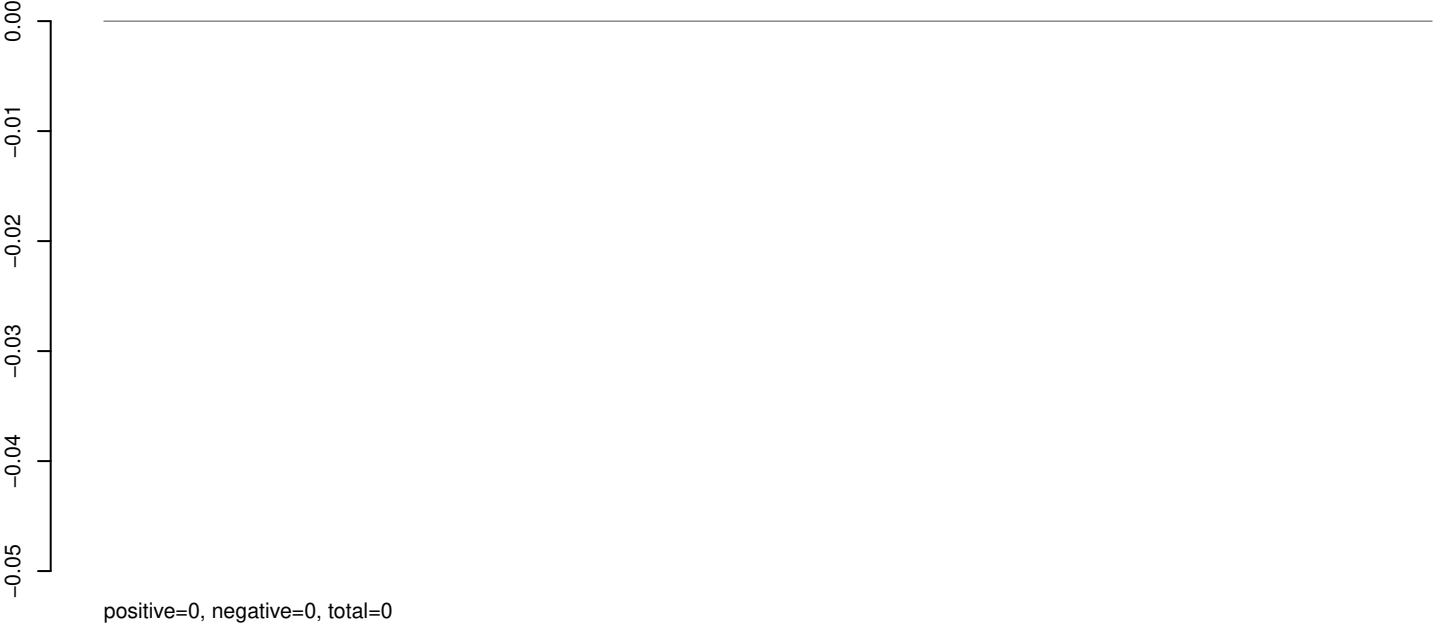
AeAeg_Aag2_BUNV1_ES.rep



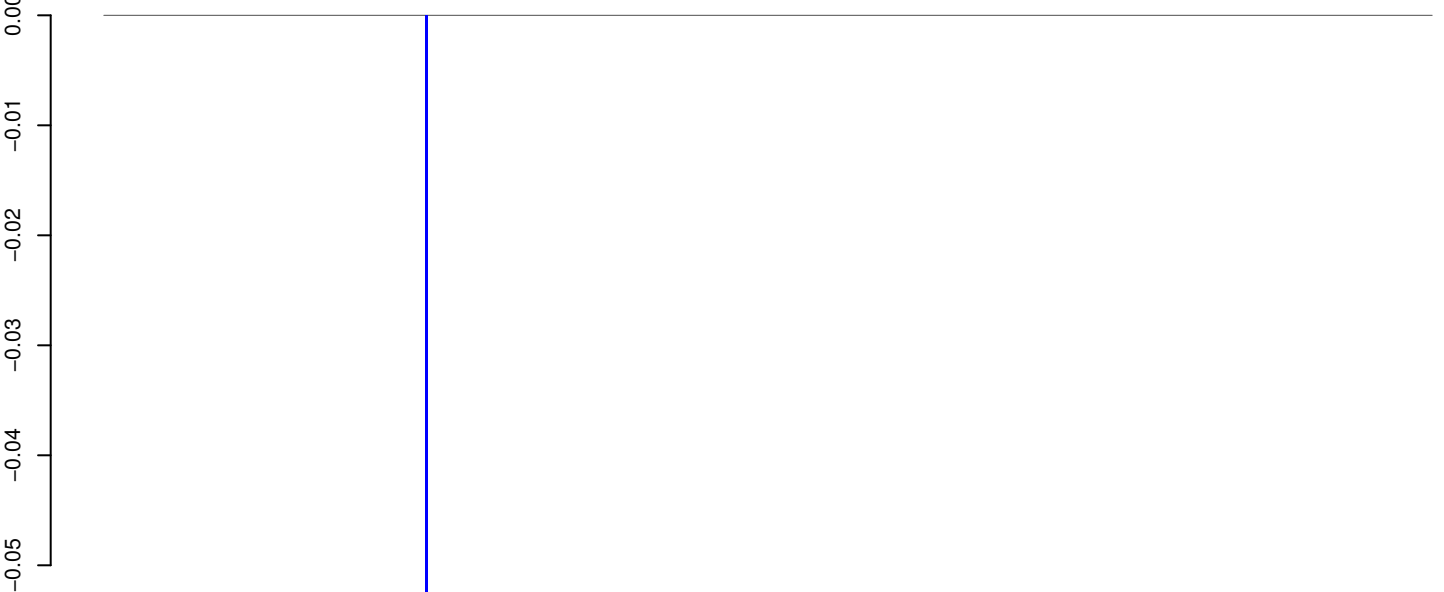
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



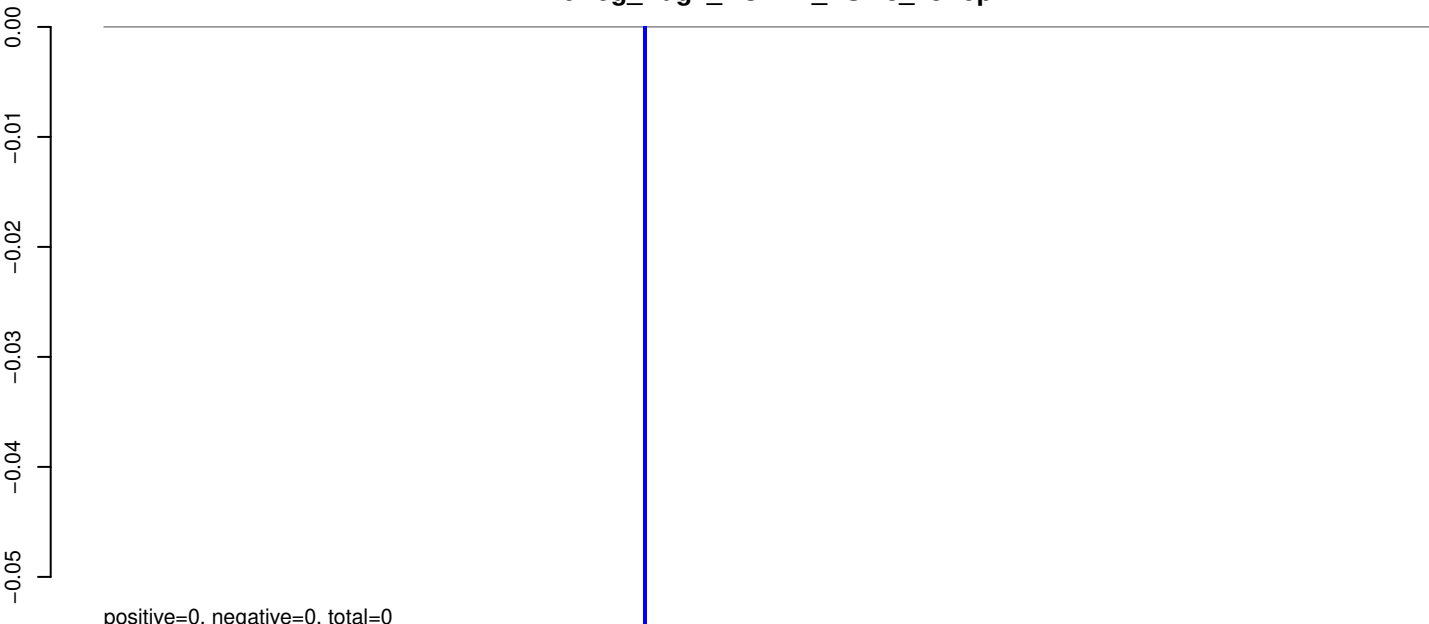
AeAeg_Aag2_BUNV1_ES.rep



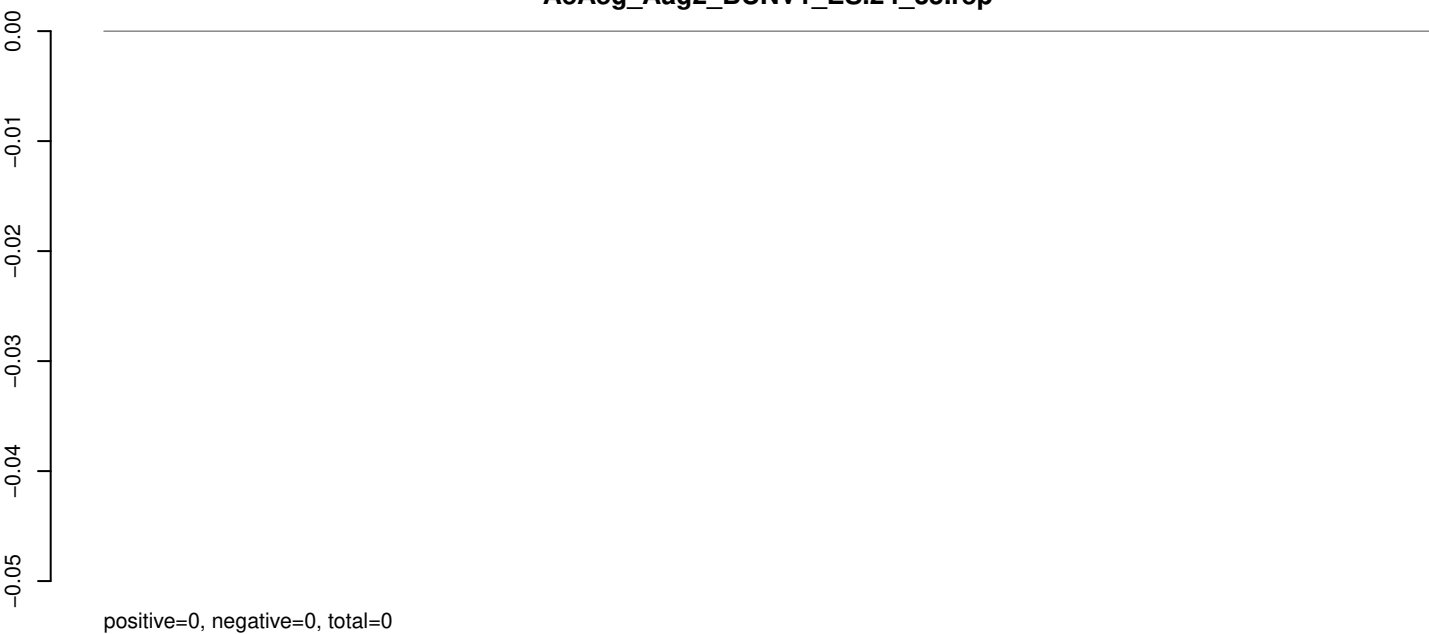
www.ebi.ac.uk/ena/browser/view/PRINCE_034447.1-Hart_Park_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopro:1-13104

0 2000 4000 6000 8000 10000 12000

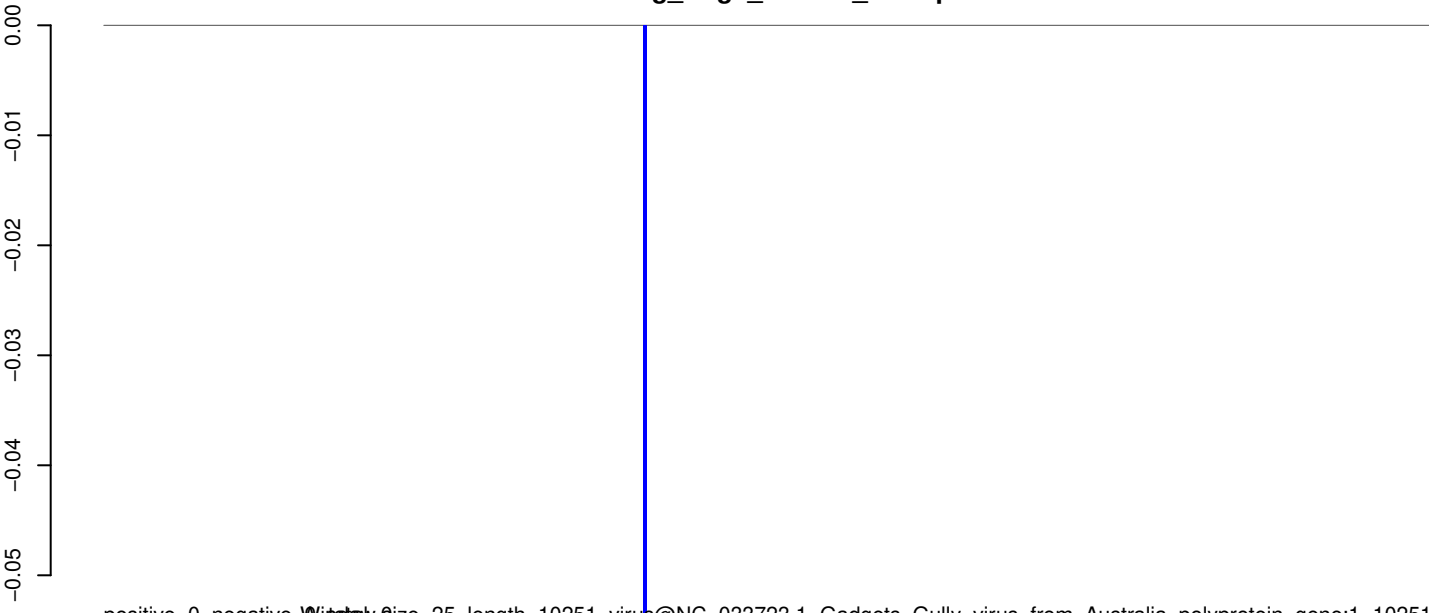
AeAeg_Aag2_BUNV1_ES.18_23.rep



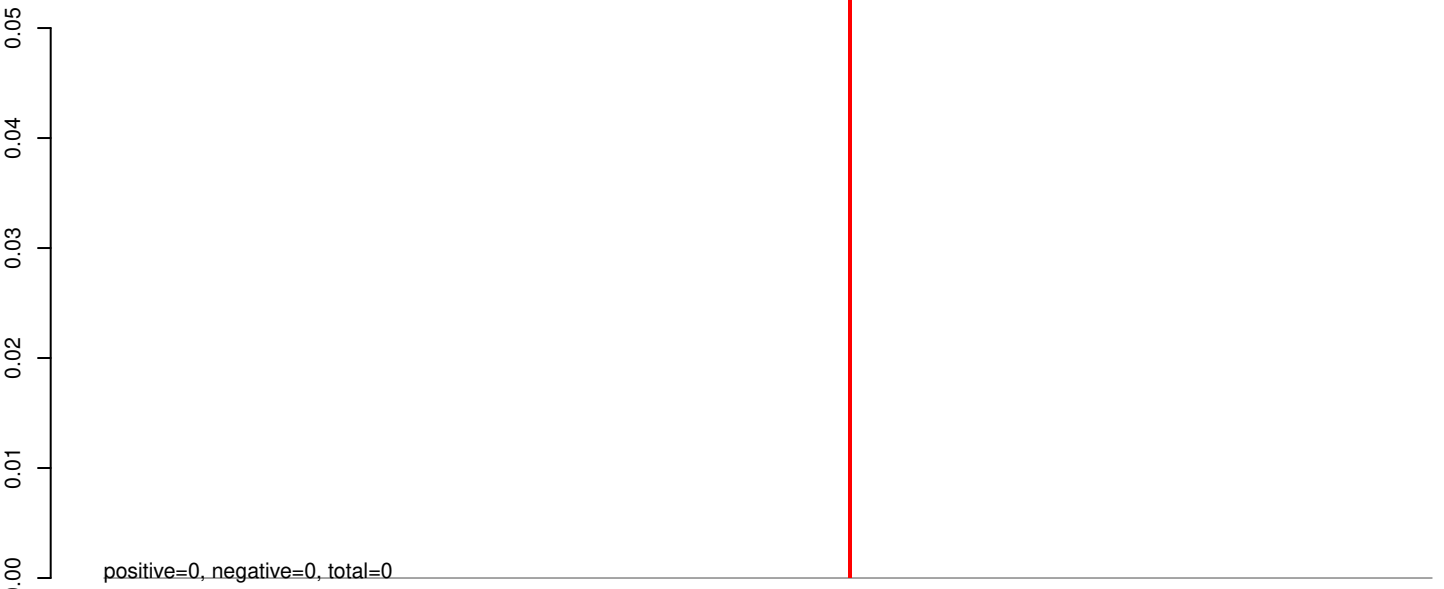
AeAeg_Aag2_BUNV1_ES.24_35.rep



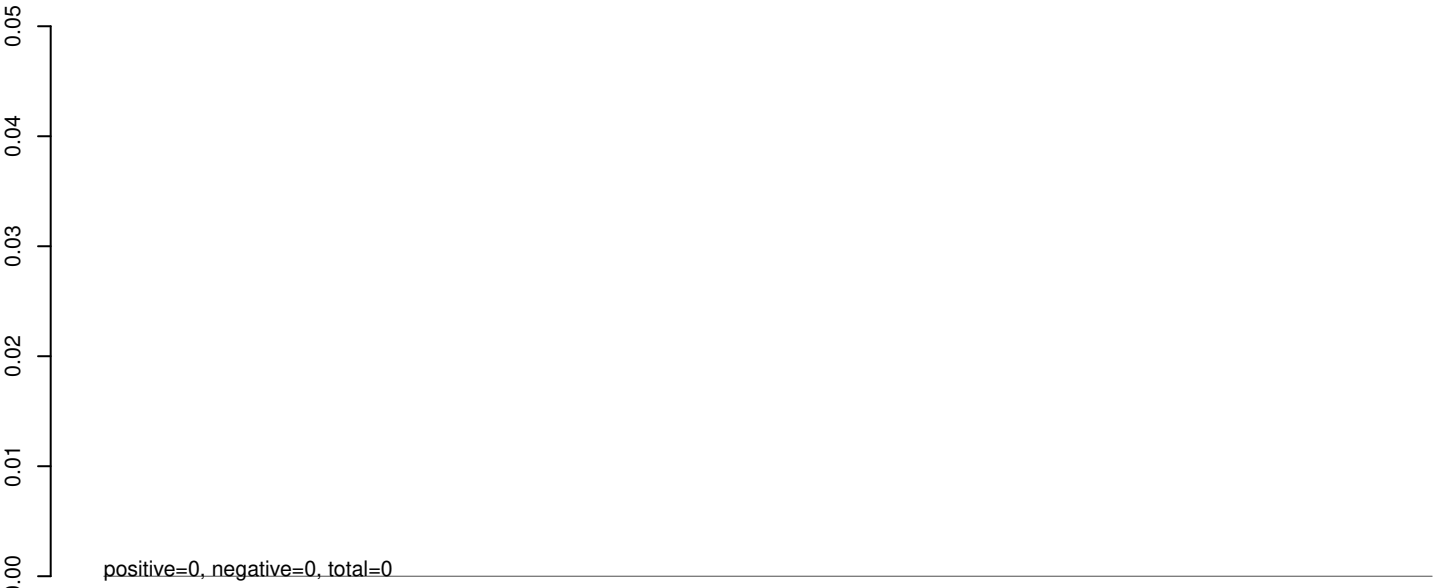
AeAeg_Aag2_BUNV1_ES.rep



AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep

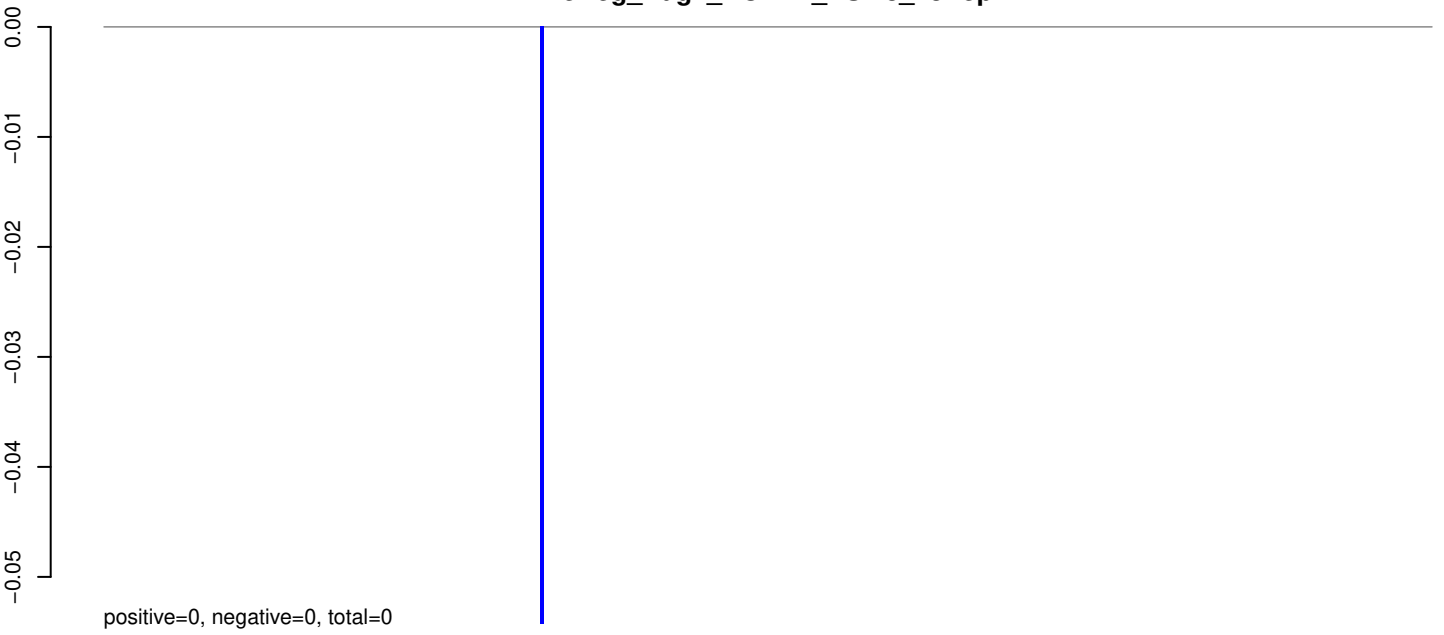


AeAeg_Aag2_BUNV1_ES.rep

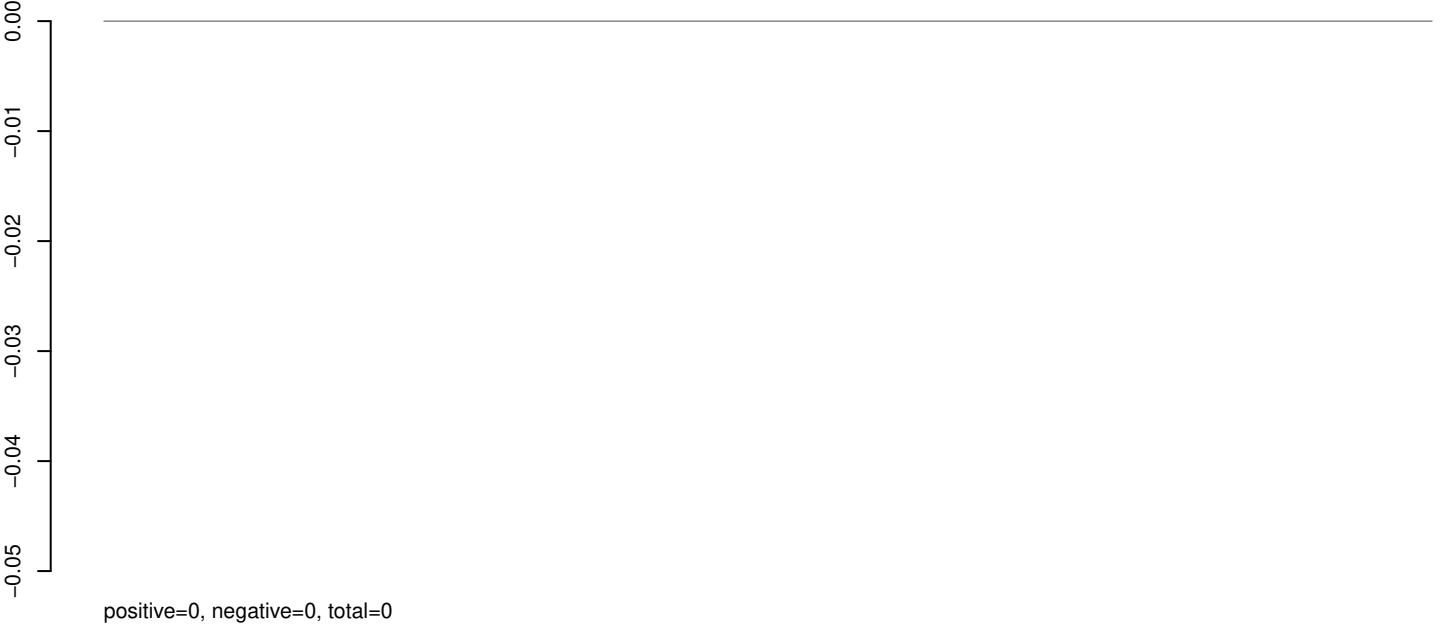


0 2000 4000 6000 8000 10000

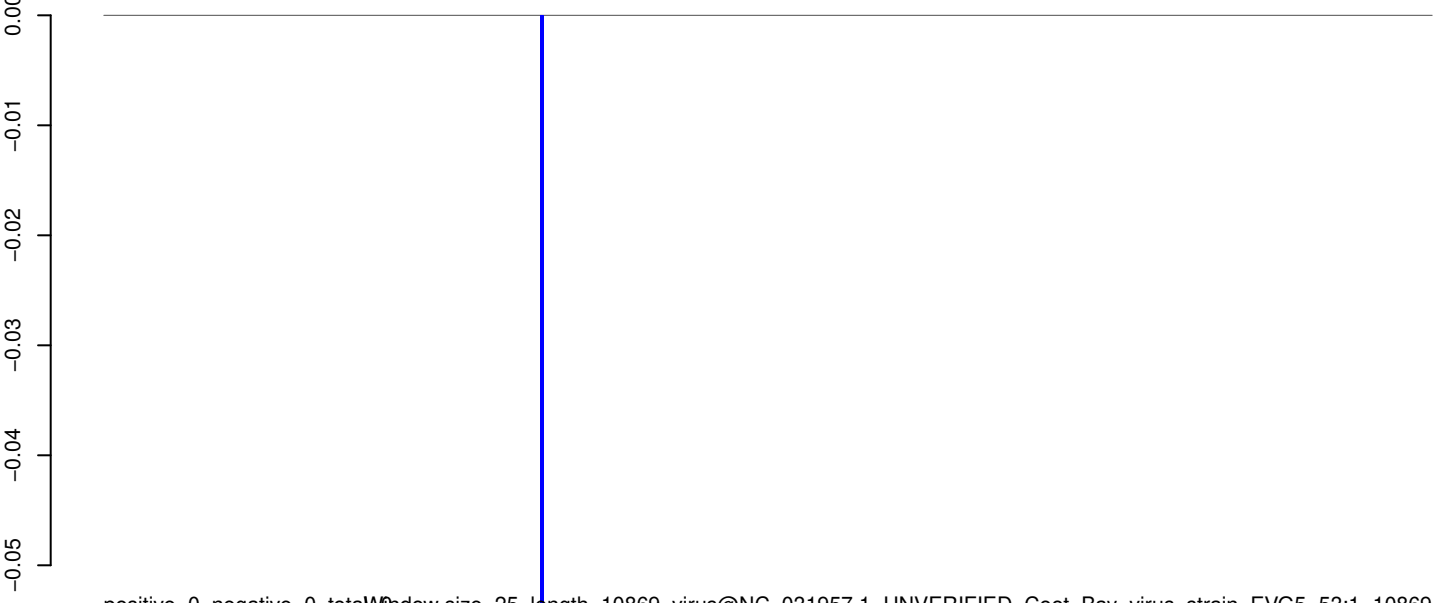
AeAeg_Aag2_BUNV1_ES.18_23.rep



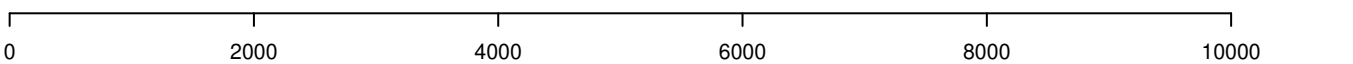
AeAeg_Aag2_BUNV1_ES.24_35.rep



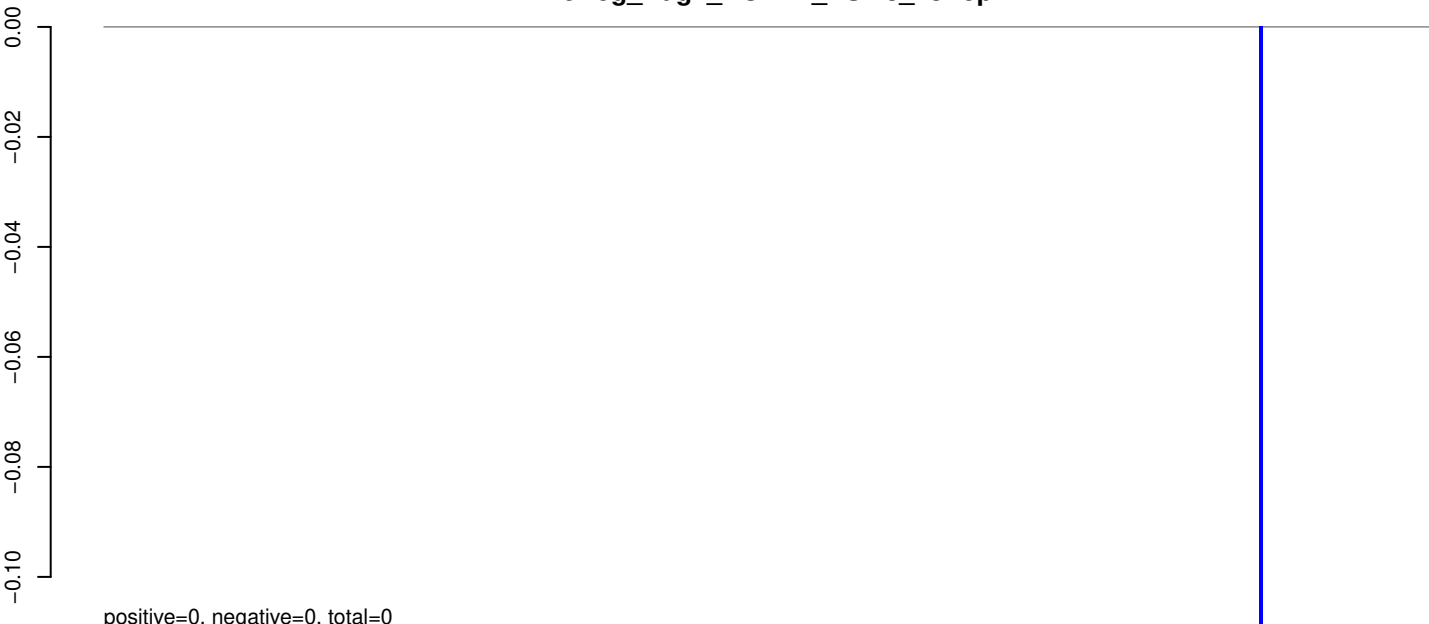
AeAeg_Aag2_BUNV1_ES.rep



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

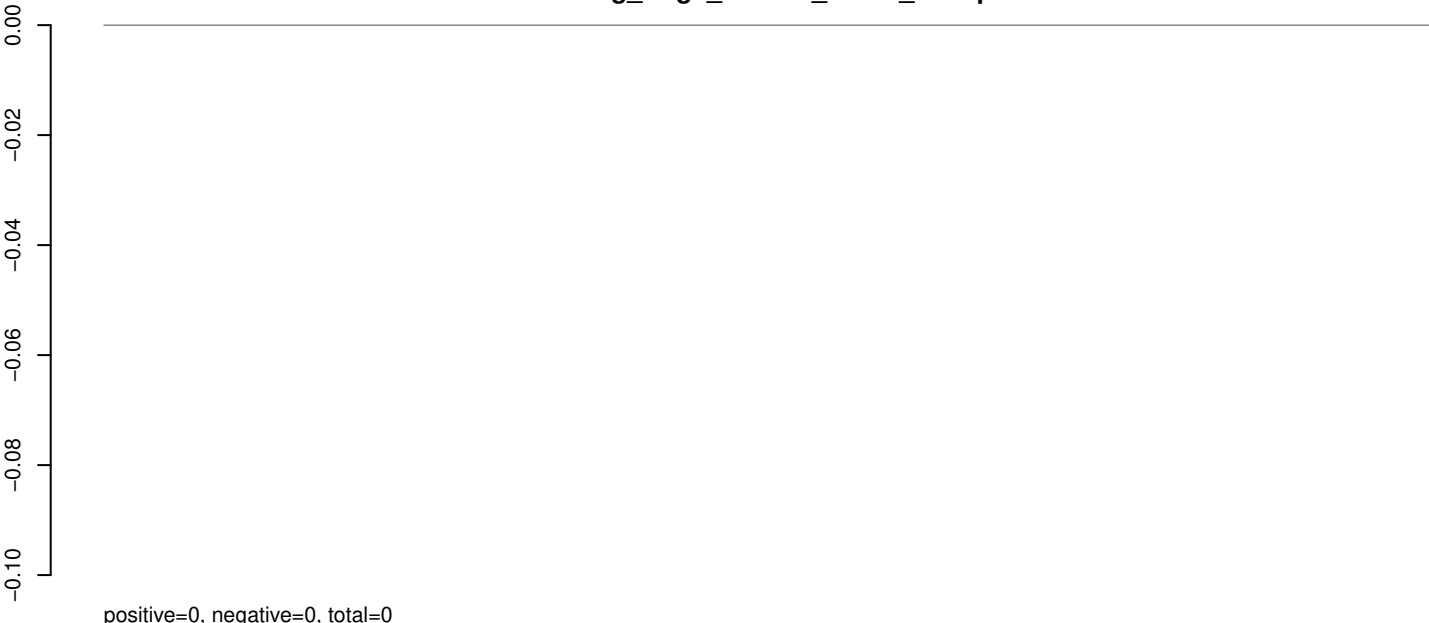


AeAeg_Aag2_BUNV1_ES.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BUNV1_ES.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BUNV1_ES.rep

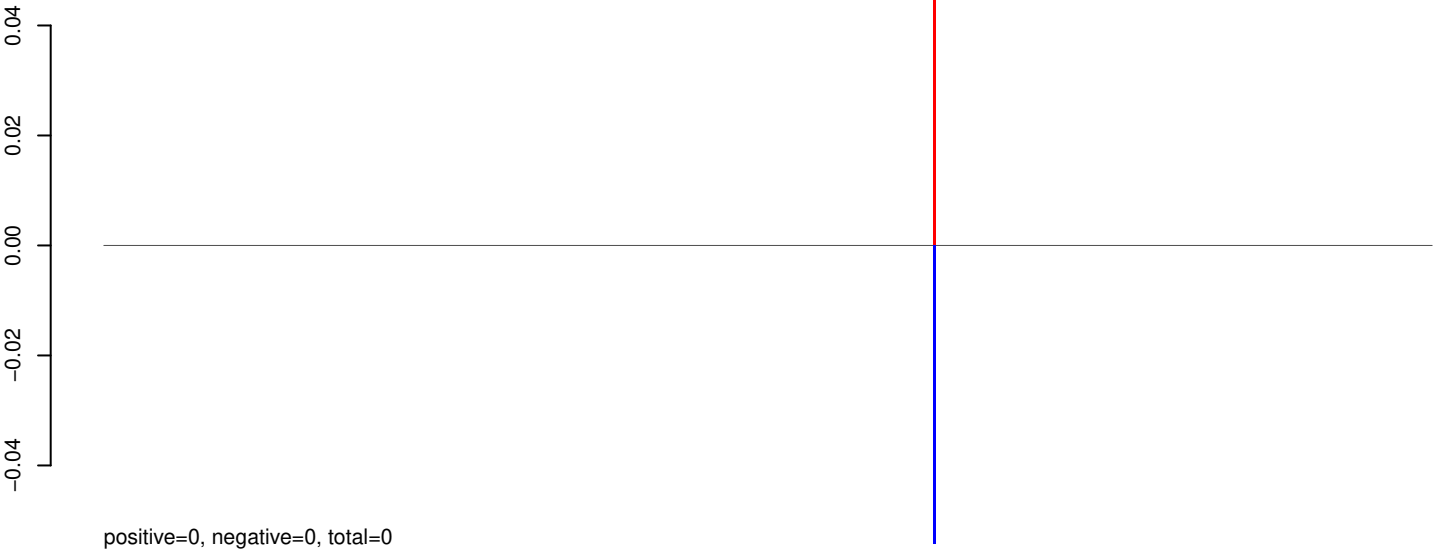


positive=0, negative=0, total=0

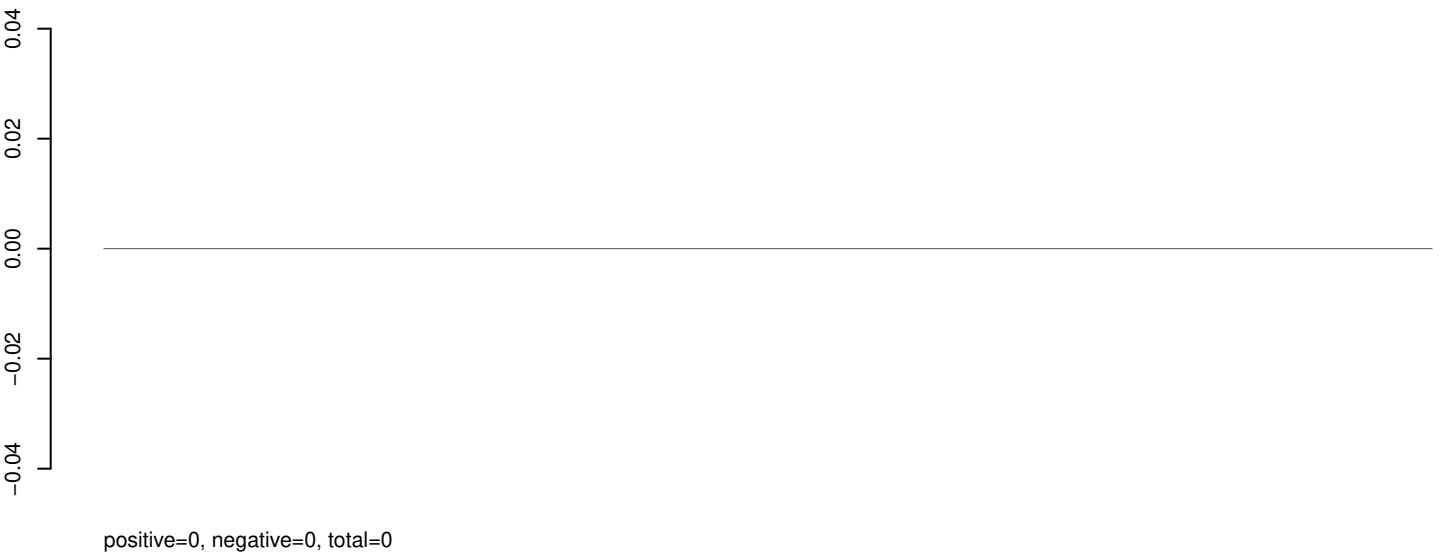
Window size=25, length=10588, virus@NC_031327.1-Anopheles_flavivirus_variant_1:1-10588

0 2000 4000 6000 8000 10000

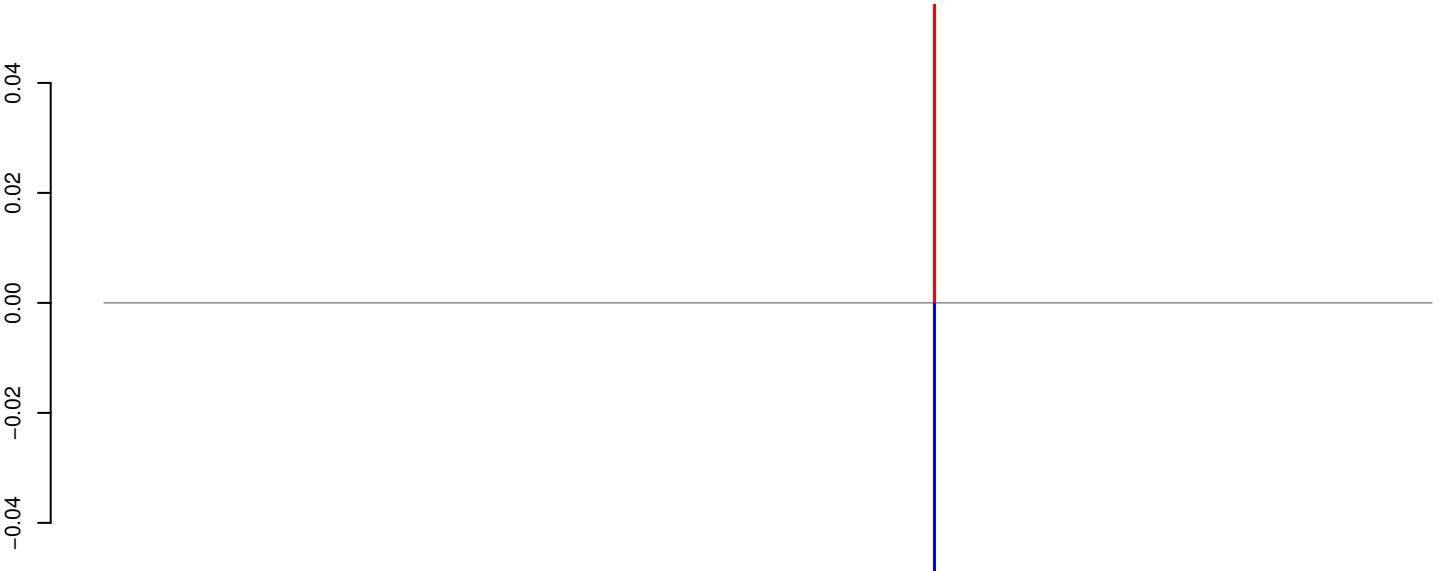
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



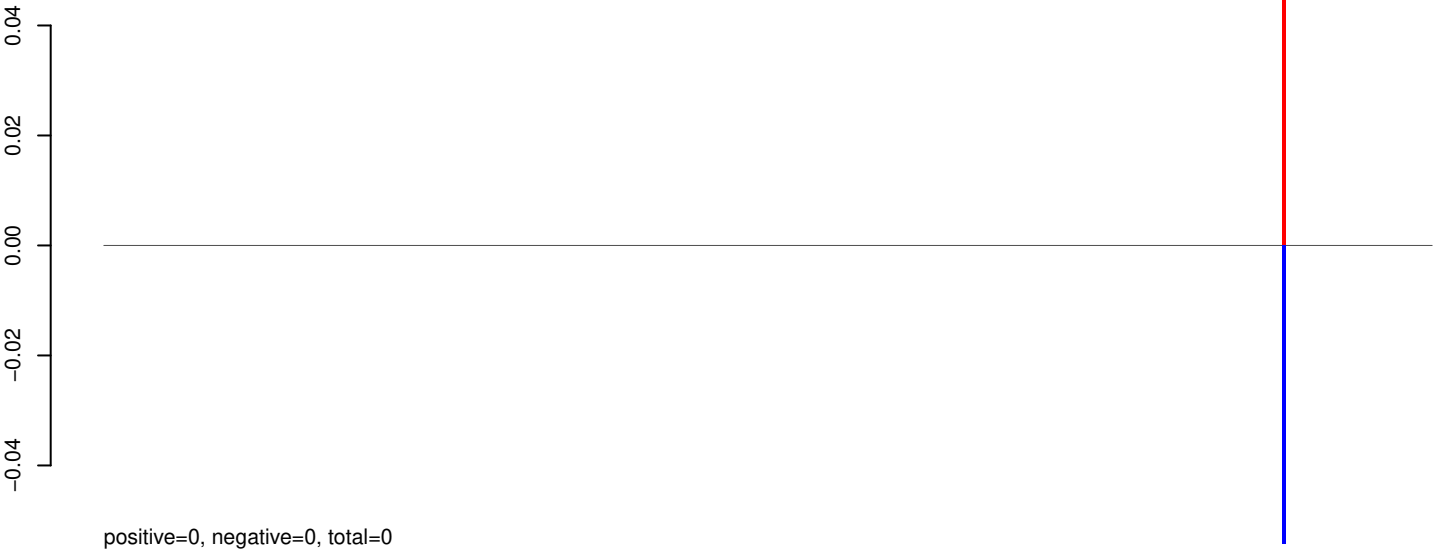
AeAeg_Aag2_BUNV1_ES.rep



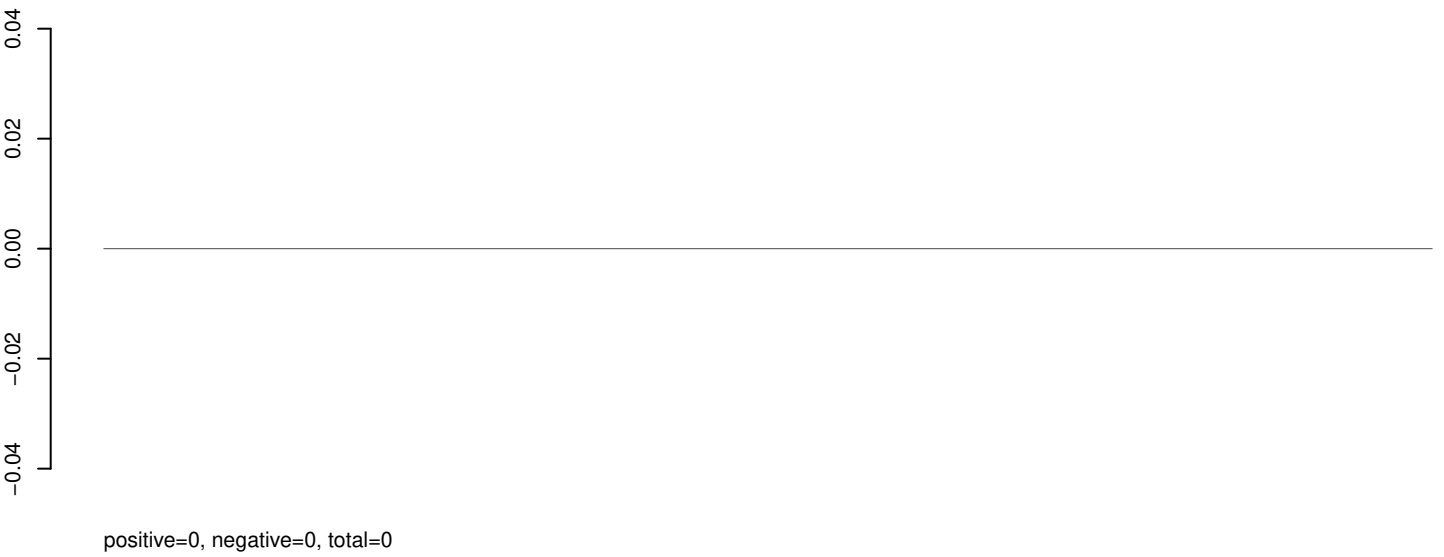
Xincheng_Mosquito_Virus_strain_XC1-6_ORF1_ORF1_ORF2_ORF2_glycoprotein_G_and_RNA-de:1-12774

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



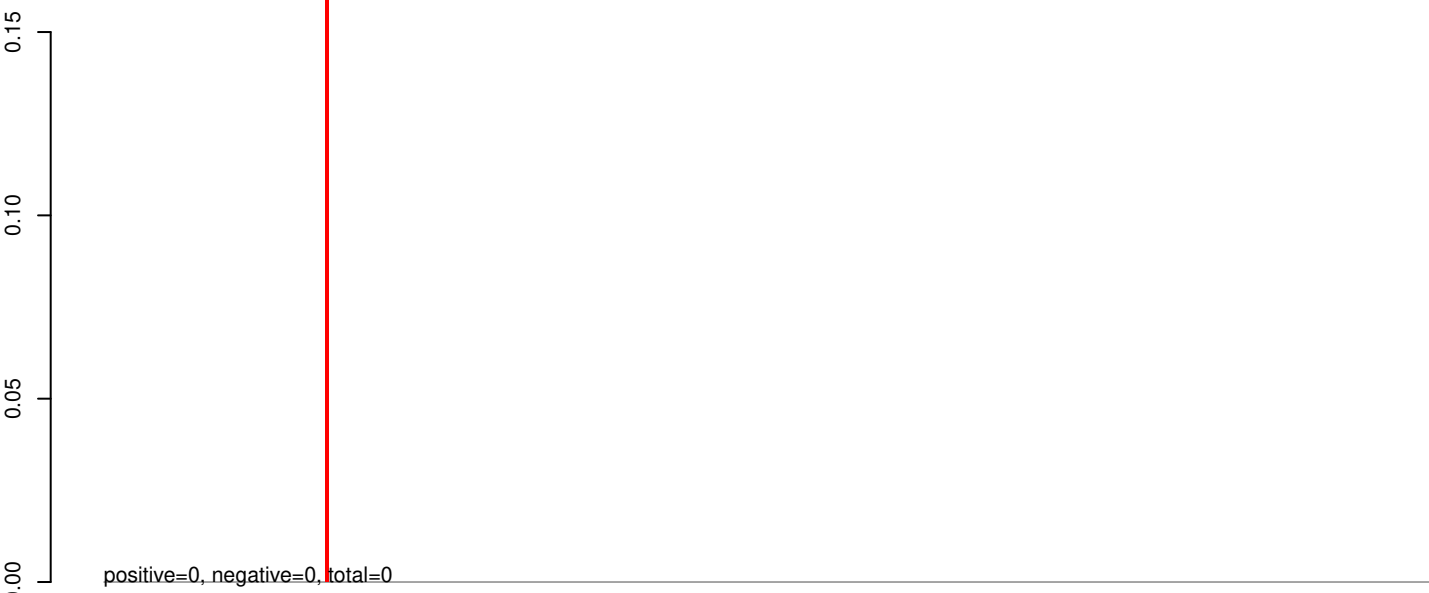
AeAeg_Aag2_BUNV1_ES.rep



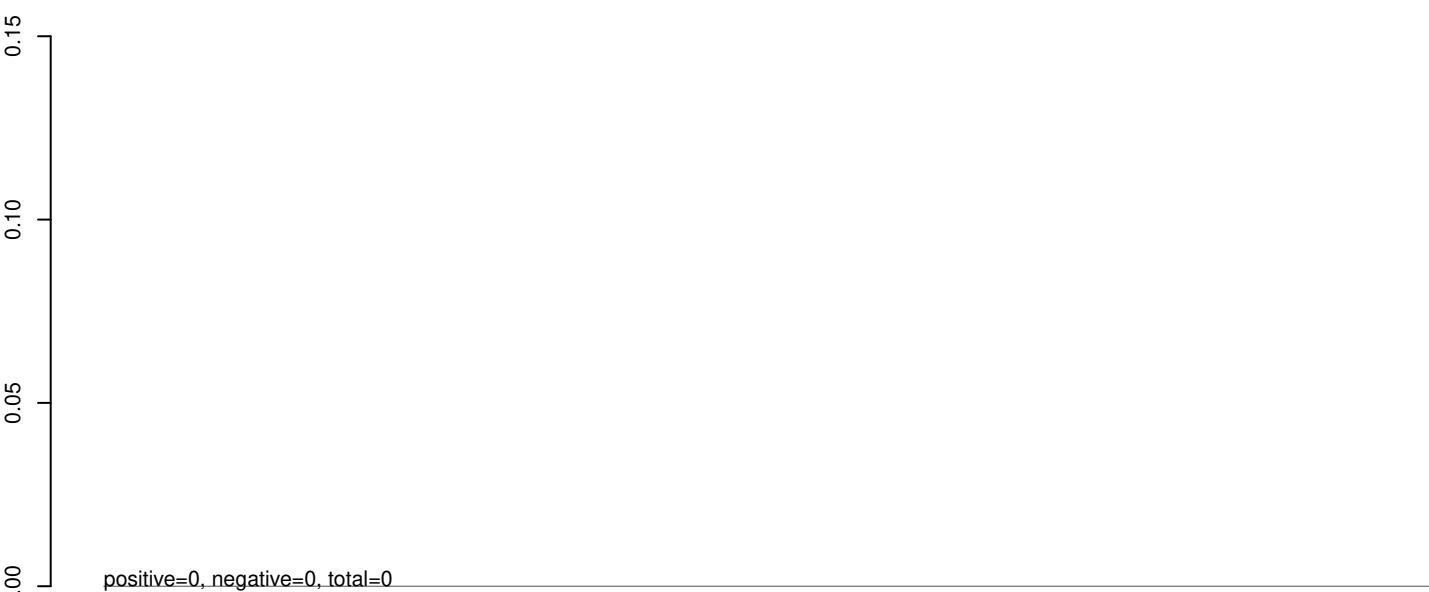
Window size=25, length=10158, virus@NC_030401.1-Hanko_virus_polyprotein_gene:1-10158

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



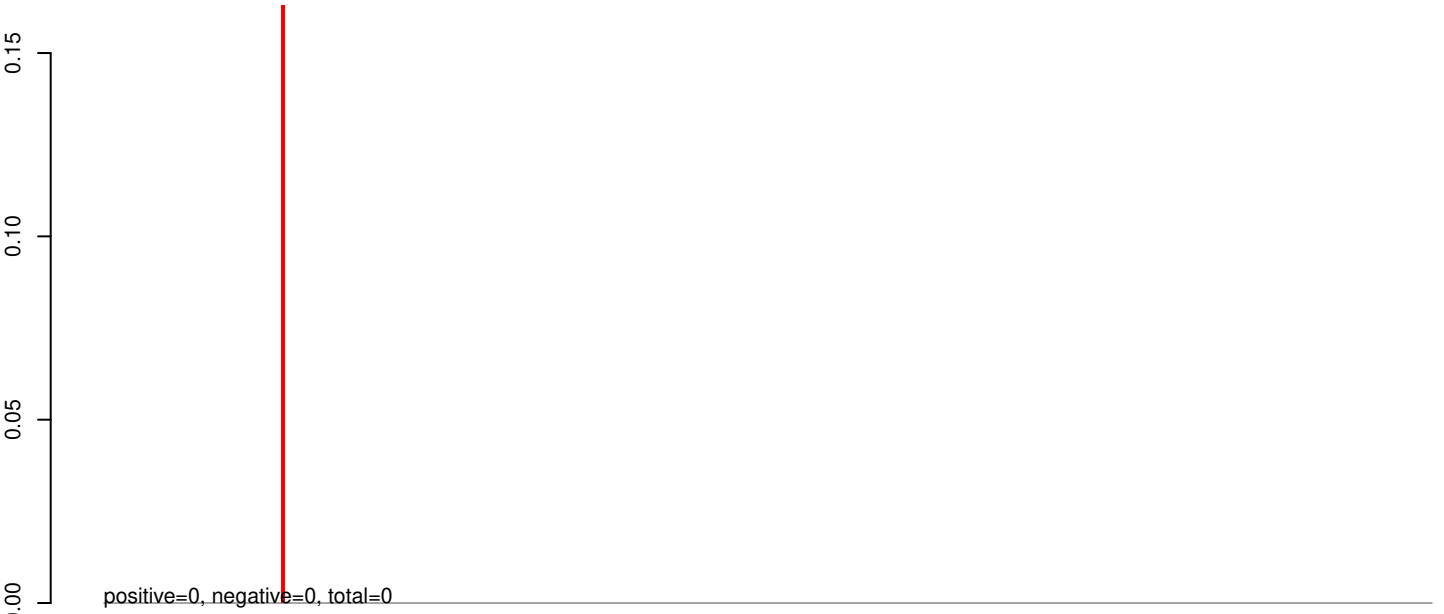
AeAeg_Aag2_BUNV1_ES.rep



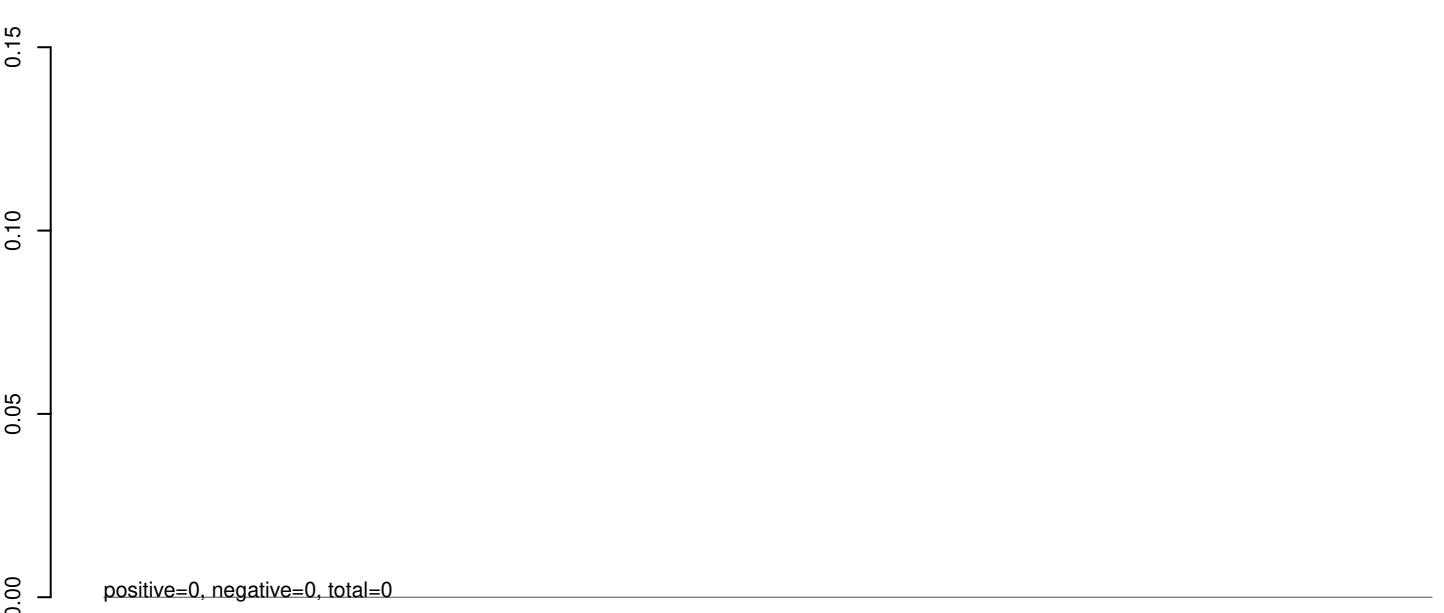
Window size=25, length=10938, virus@NC_027819.1-Mercadeo_virus_isolate_ER-M10:1-10938

0 2000 4000 6000 8000 10000

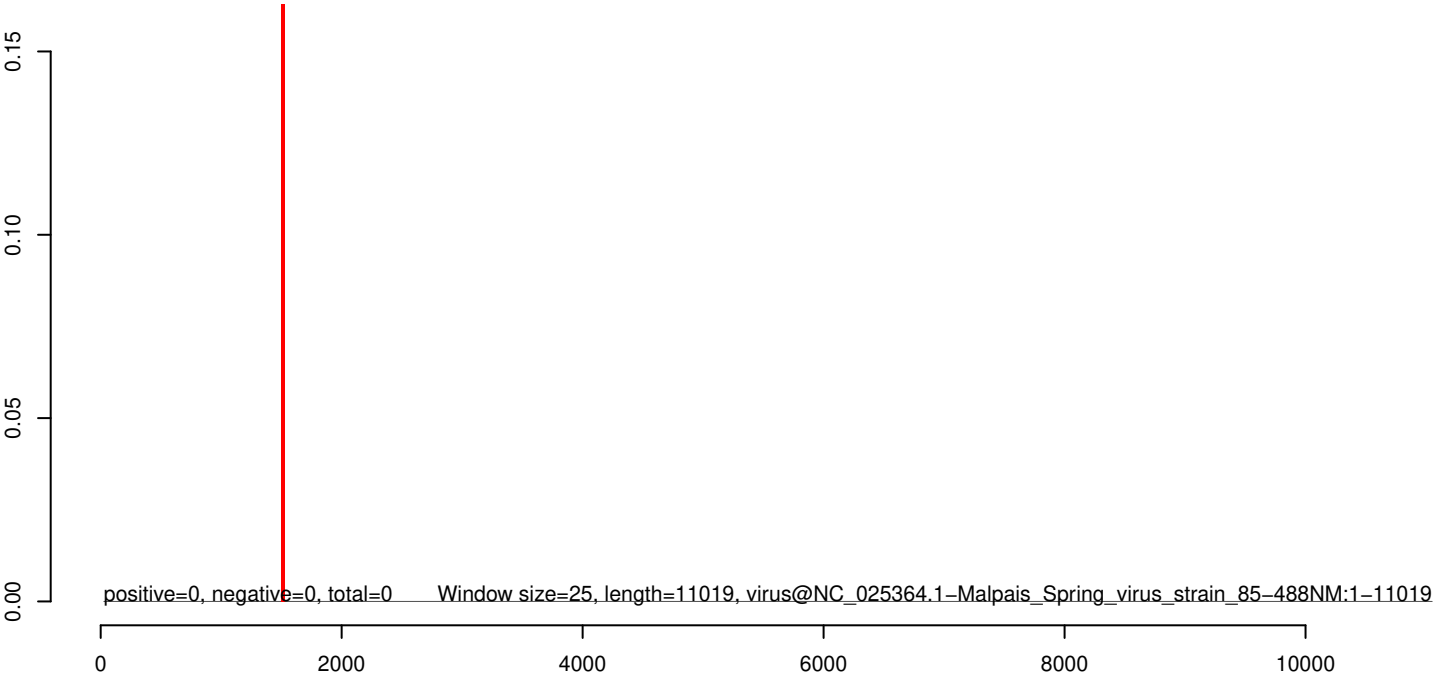
AeAeg_Aag2_BUNV1_ES.18_23.rep



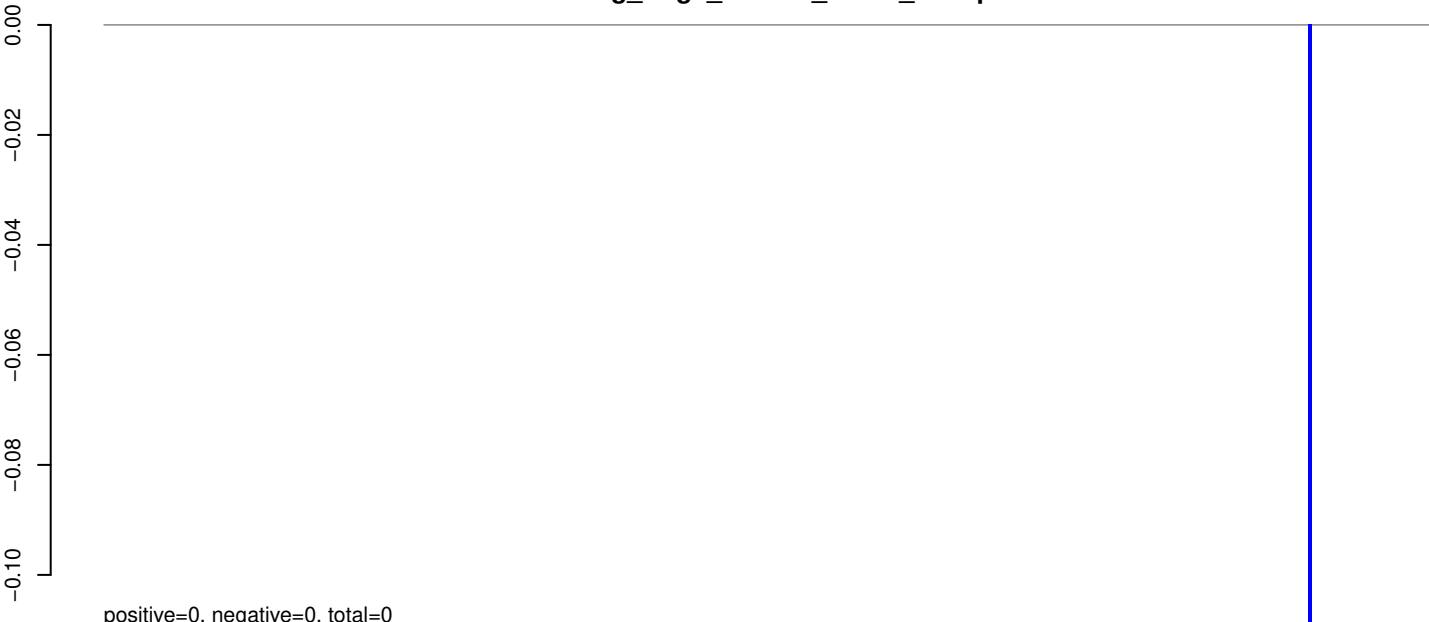
AeAeg_Aag2_BUNV1_ES.24_35.rep



AeAeg_Aag2_BUNV1_ES.rep

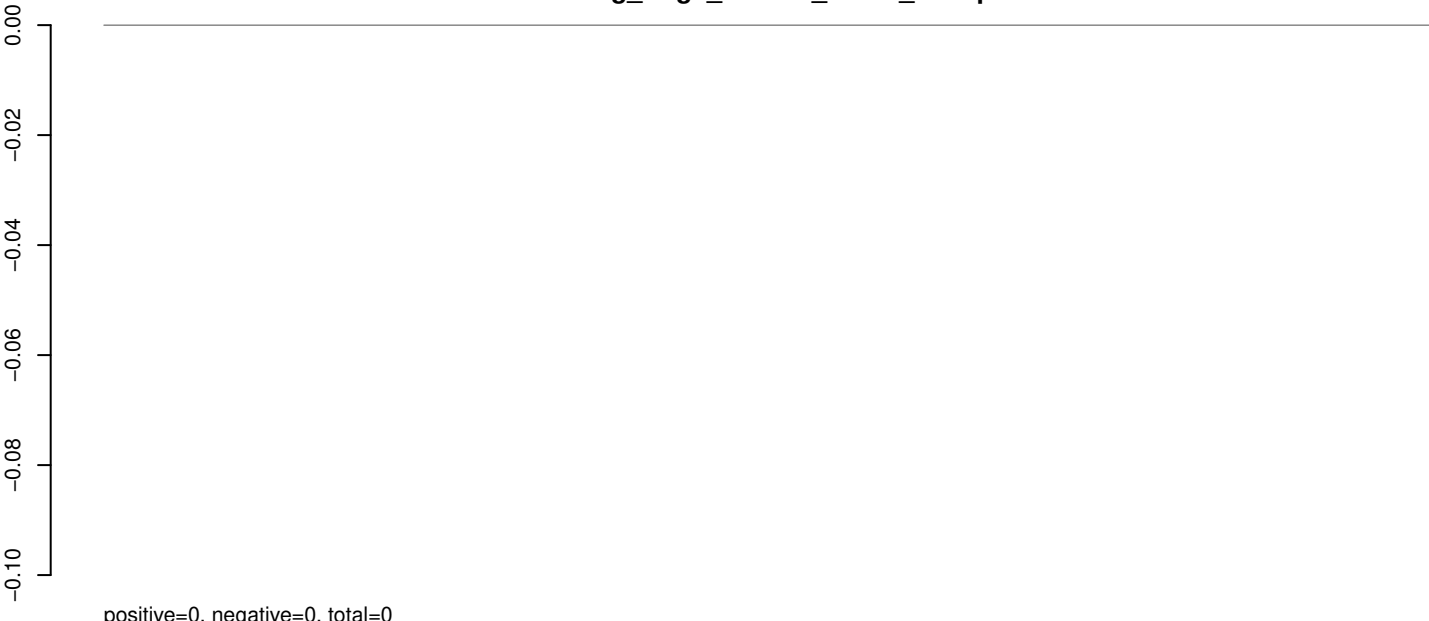


AeAeg_Aag2_BUNV1_ES.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BUNV1_ES.24_35.rep



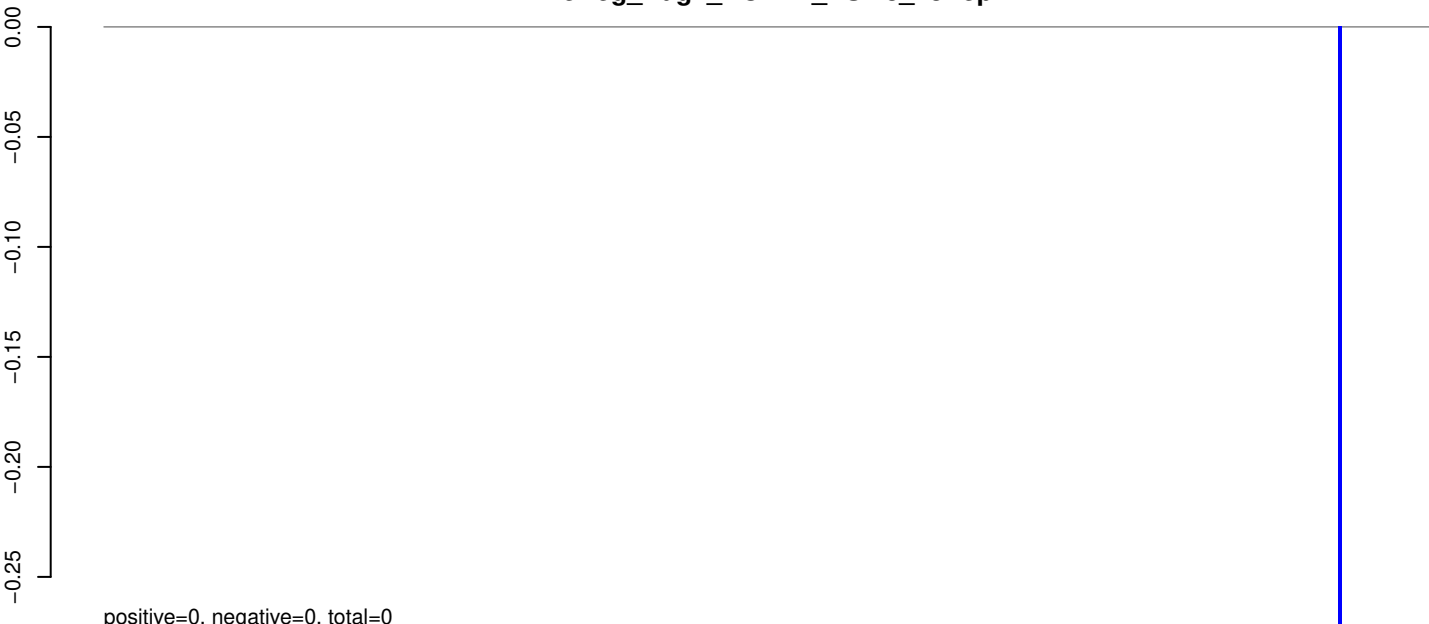
positive=0, negative=0, total=0

AeAeg_Aag2_BUNV1_ES.rep



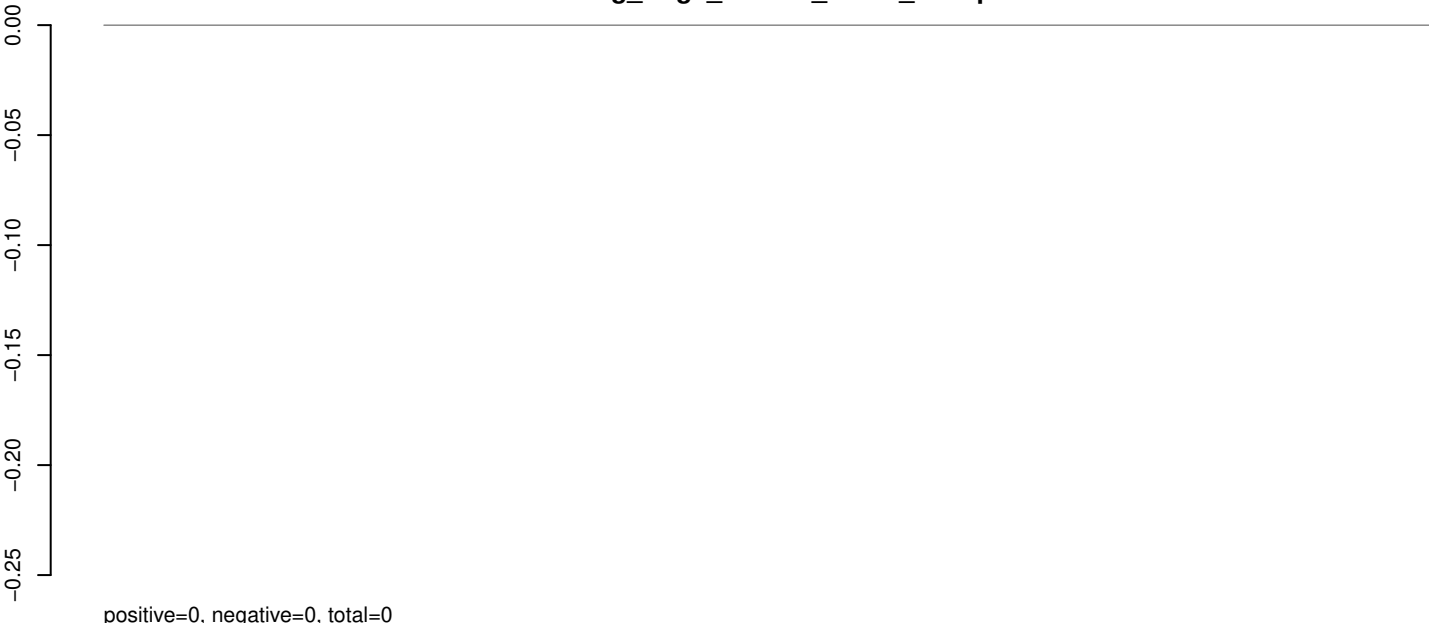
positive=0, negative=0, total=0, width=25, length=10688, virus@NC_023439.1-Kama_virus_strain_LEIV-20776Tat_polyprotein_gene:1-10688

AeAeg_Aag2_BUNV1_ES.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BUNV1_ES.24_35.rep



positive=0, negative=0, total=0

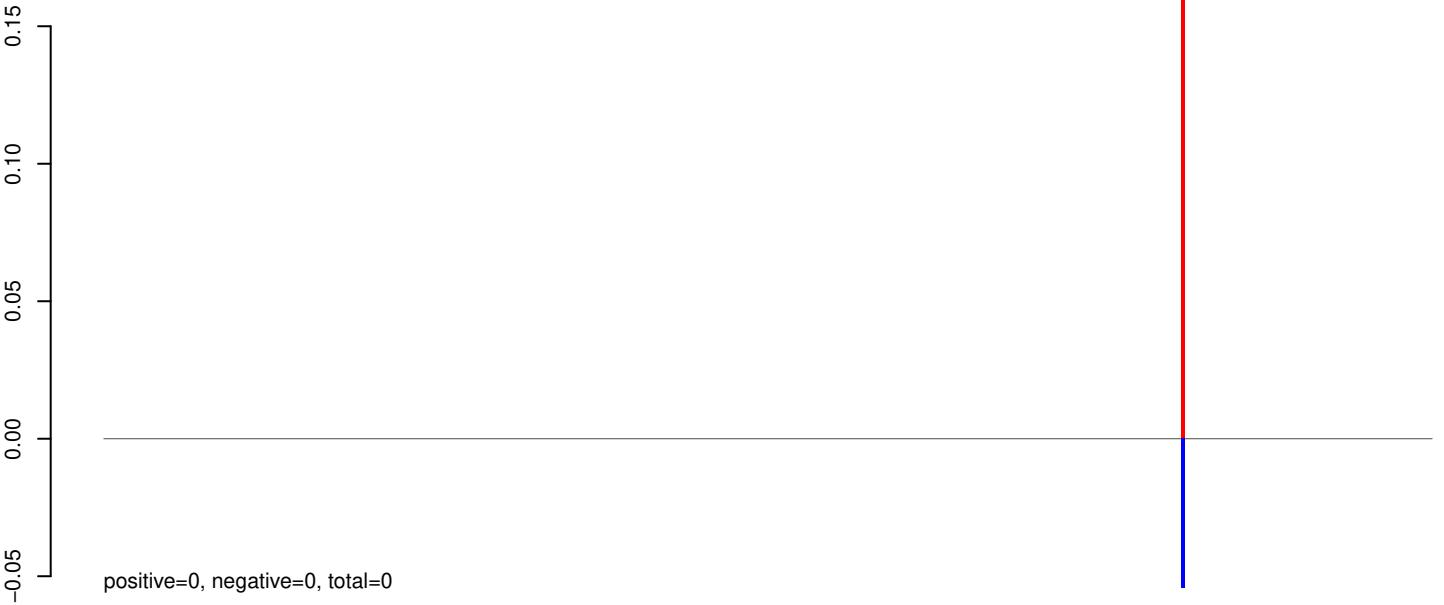
AeAeg_Aag2_BUNV1_ES.rep



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

0 2000 4000 6000 8000 10000

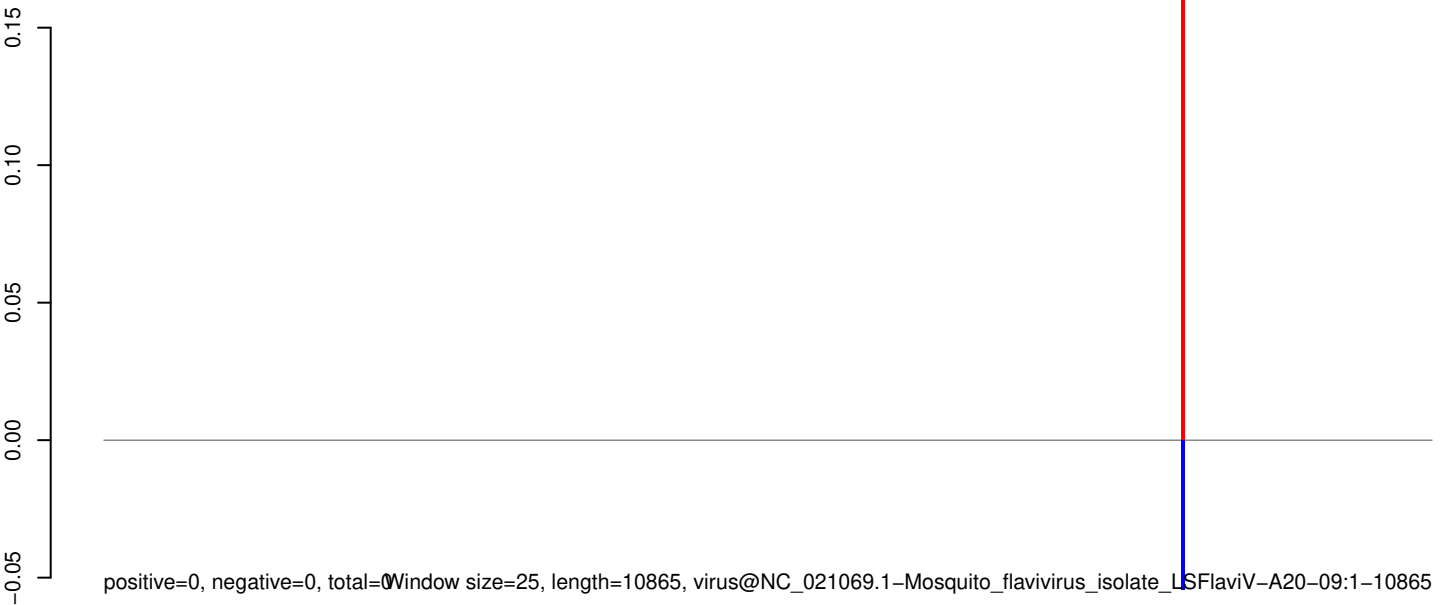
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



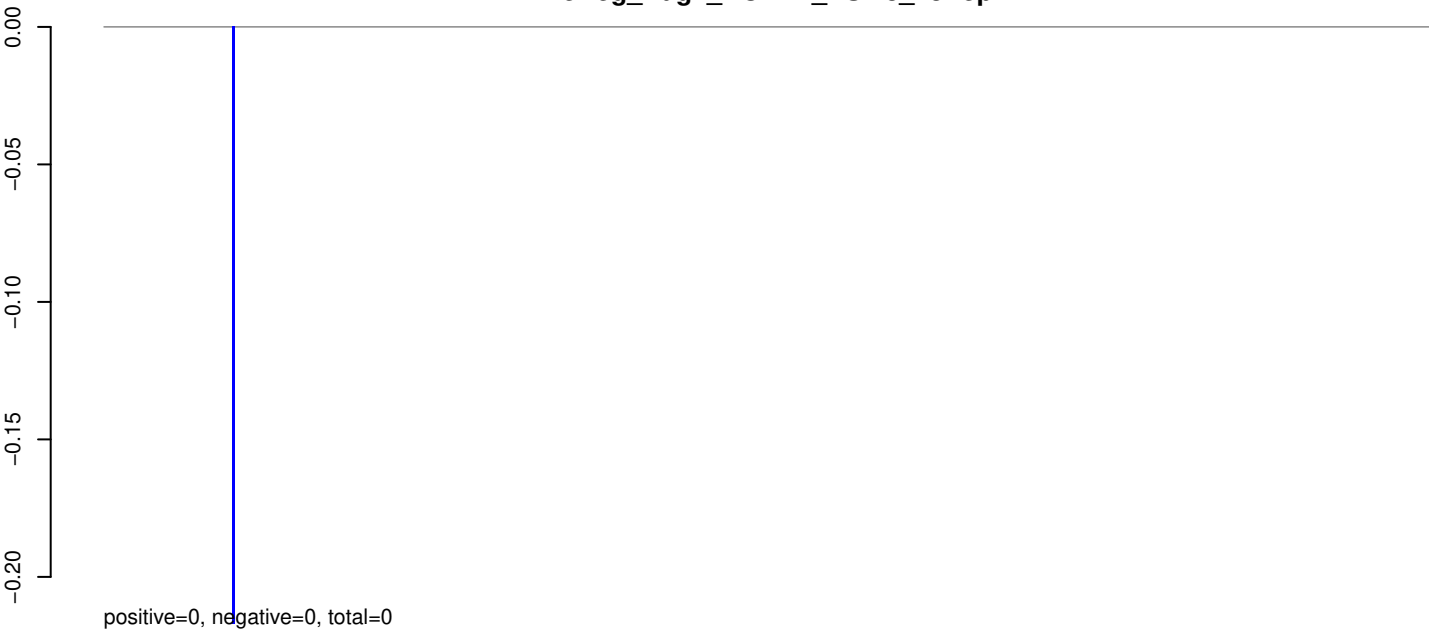
AeAeg_Aag2_BUNV1_ES.rep



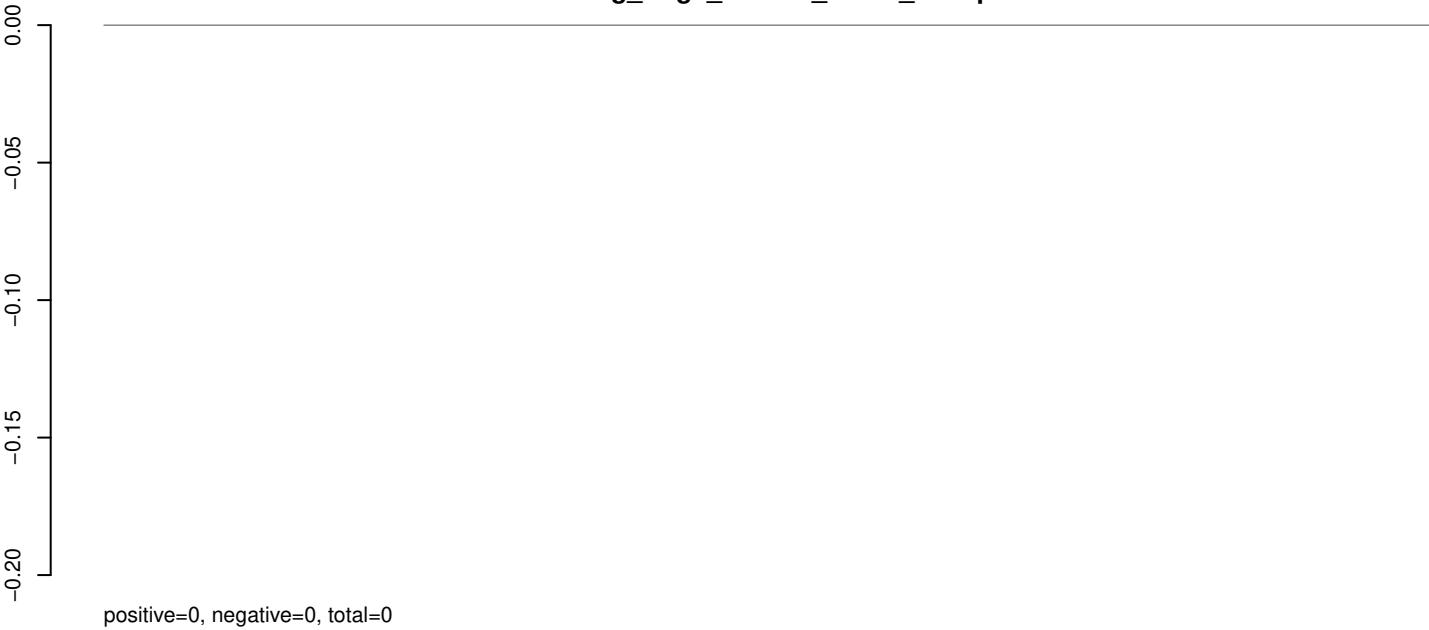
Window size=25, length=10865, virus@NC_021069.1-Mosquito_flavivirus_isolate_LSFlaviV-A20-09:1-10865

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep

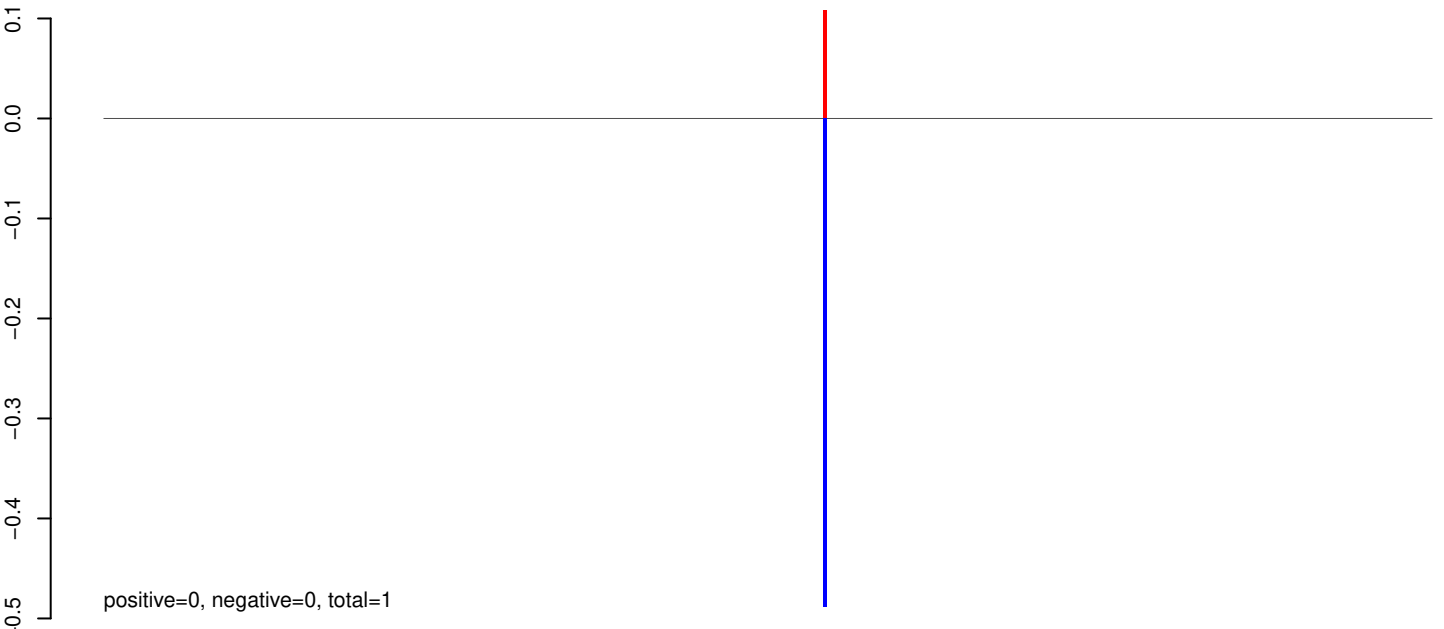


AeAeg_Aag2_BUNV1_ES.rep

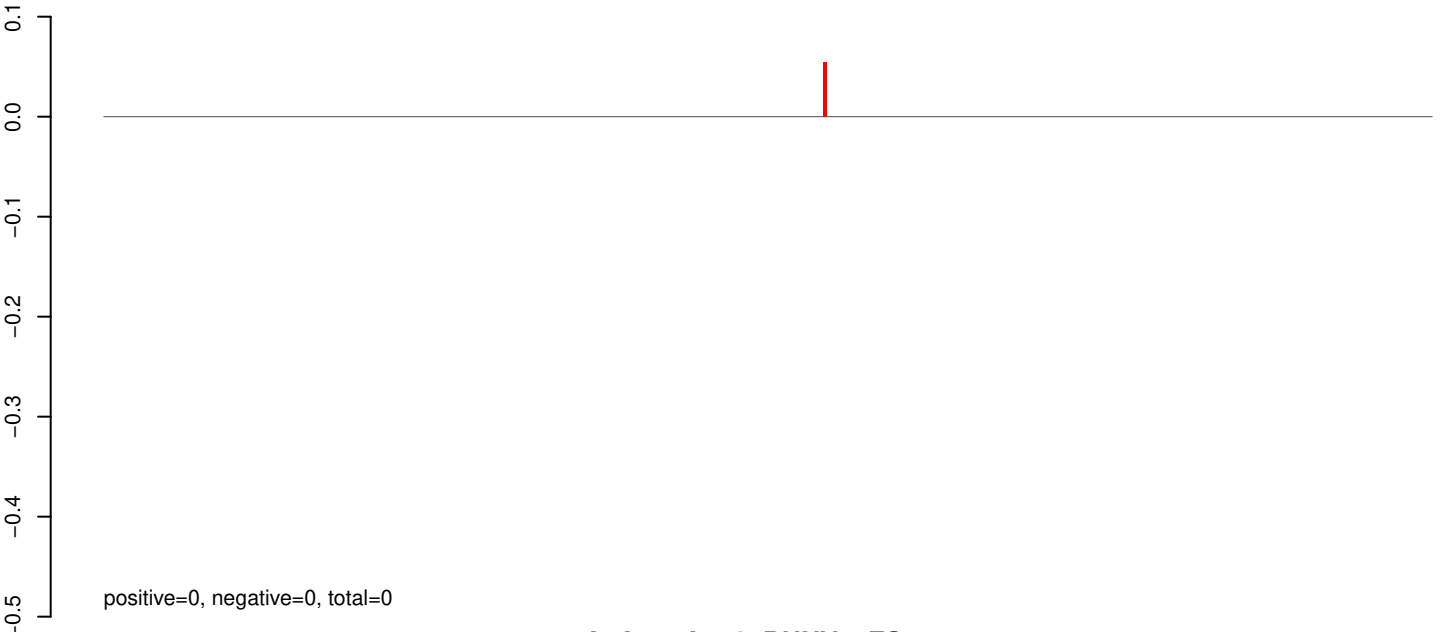


Window size=25, length=14717, virus@NC_017685.1-Obodhiang_virus:1-14717

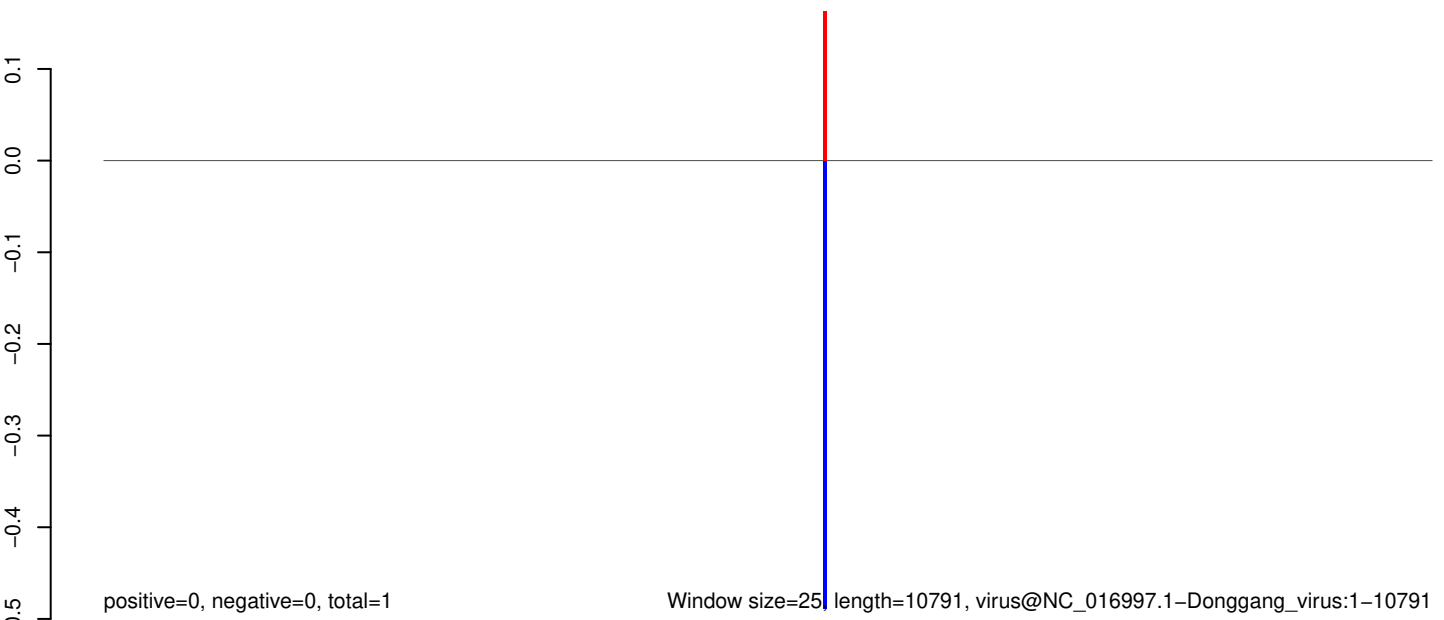
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



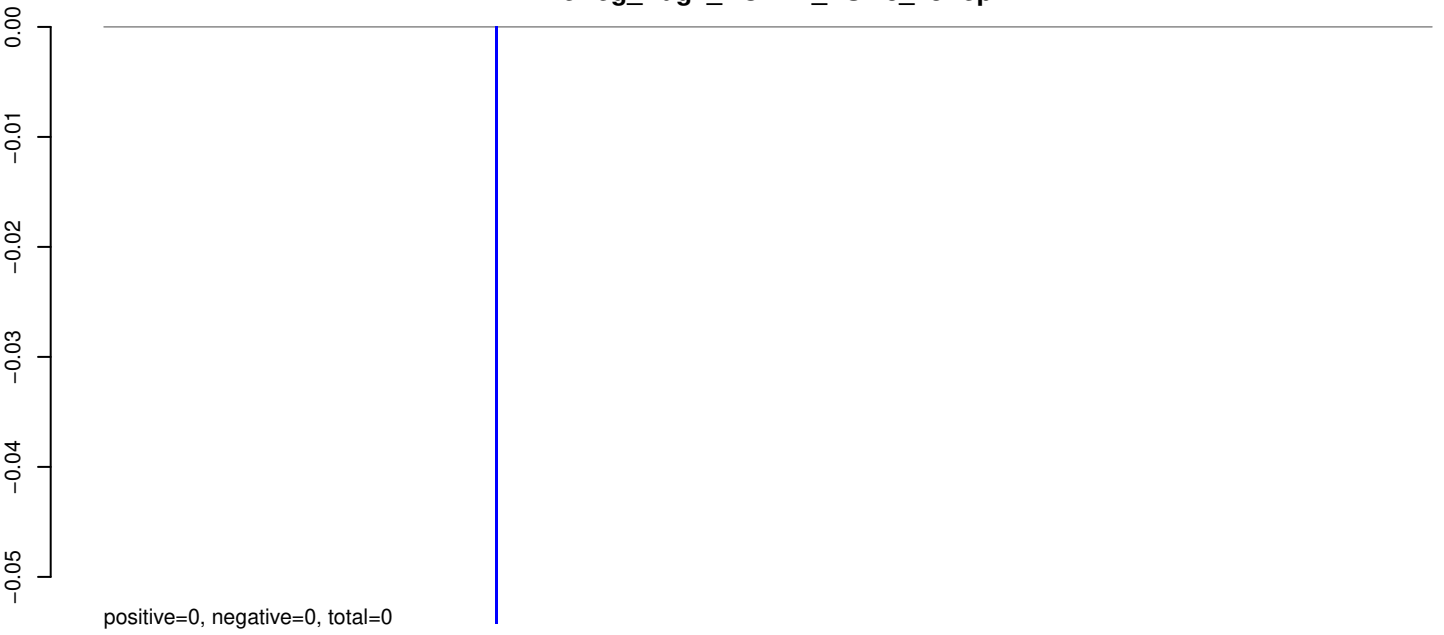
AeAeg_Aag2_BUNV1_ES.rep



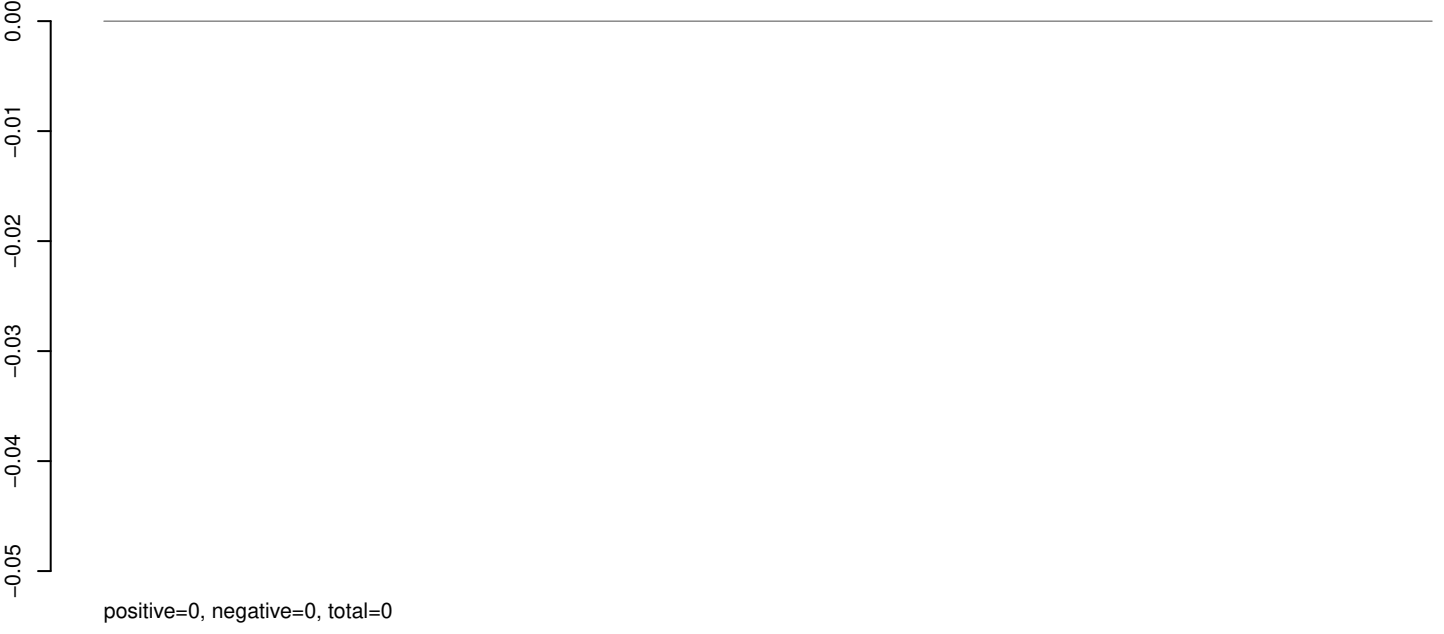
Window size=25, length=10791, virus@NC_016997.1-Donggang_virus:1-10791

0 2000 4000 6000 8000 10000

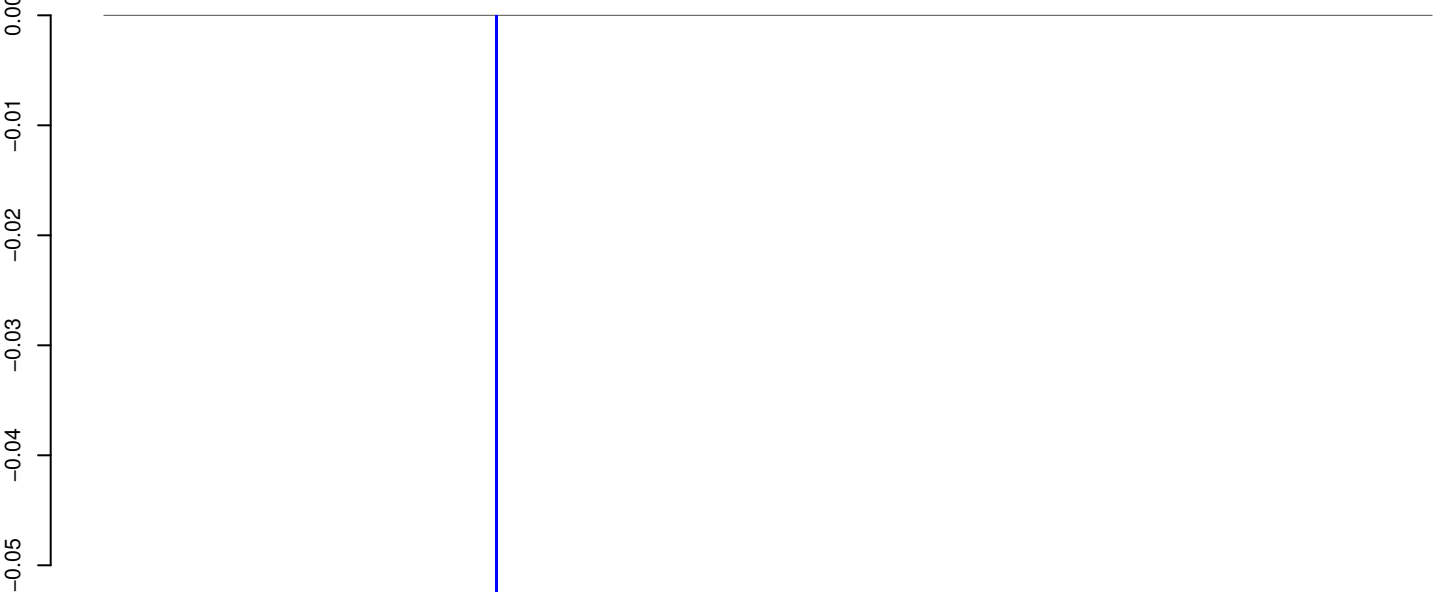
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



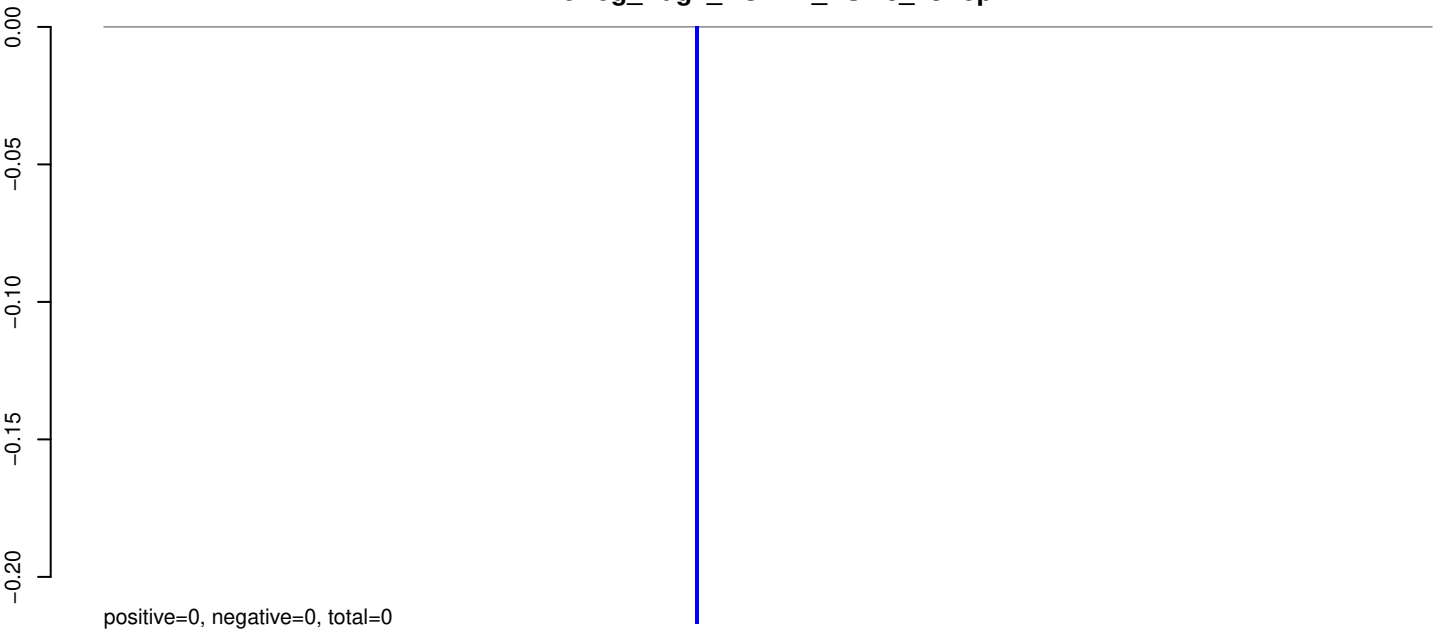
AeAeg_Aag2_BUNV1_ES.rep



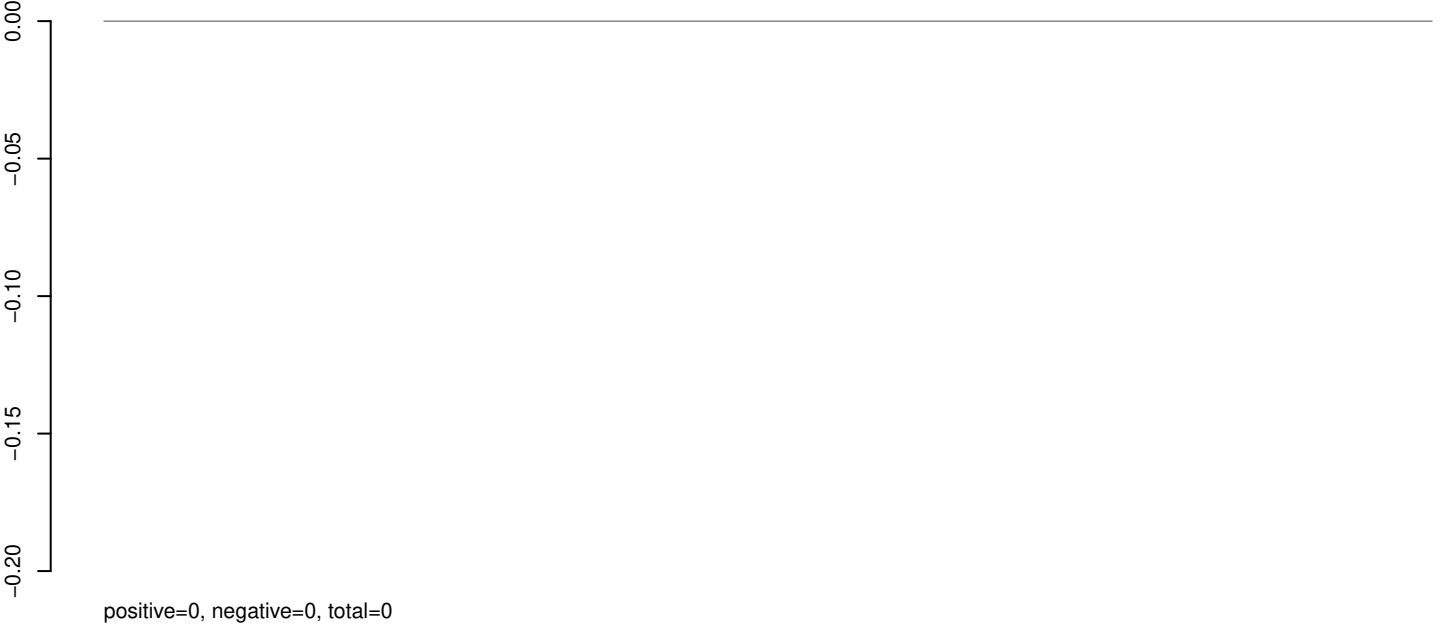
length=12390, coverage=0.161350 -Drosophila_melanogaster_sigma_virus_AP30_N_P_X_M_G_and_L_genes_genomic_RNA_isolate:1-12390

0 2000 4000 6000 8000 10000 12000

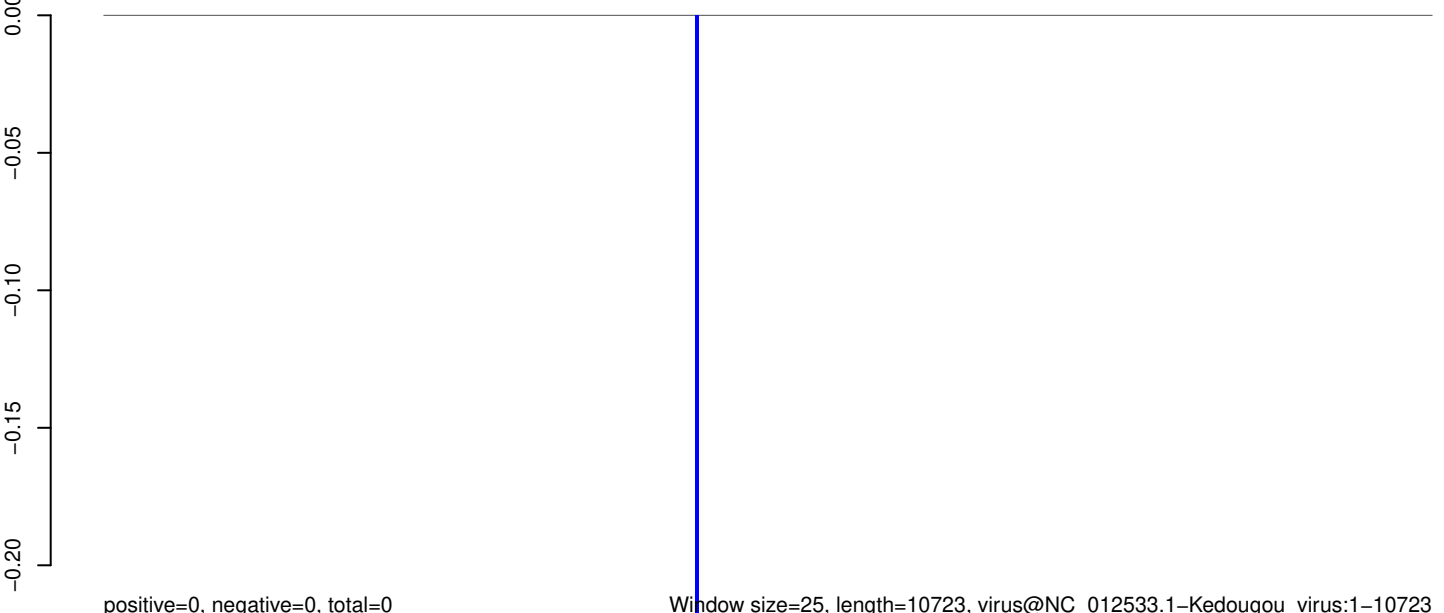
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



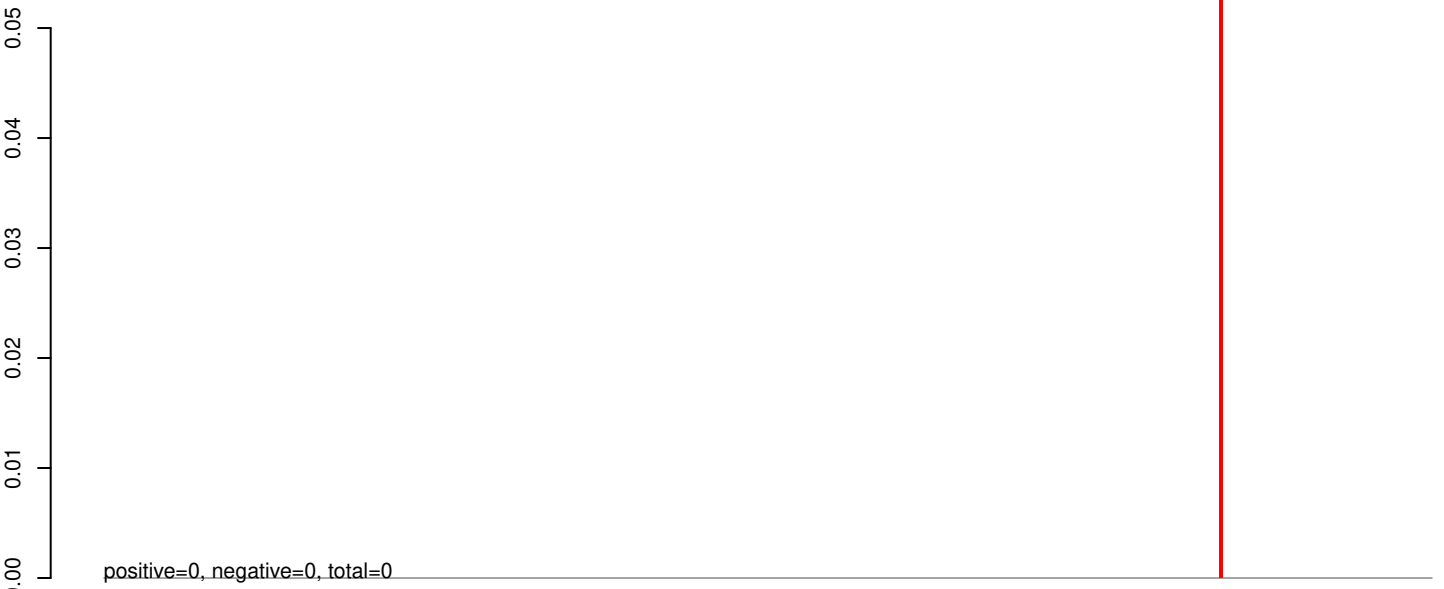
AeAeg_Aag2_BUNV1_ES.rep



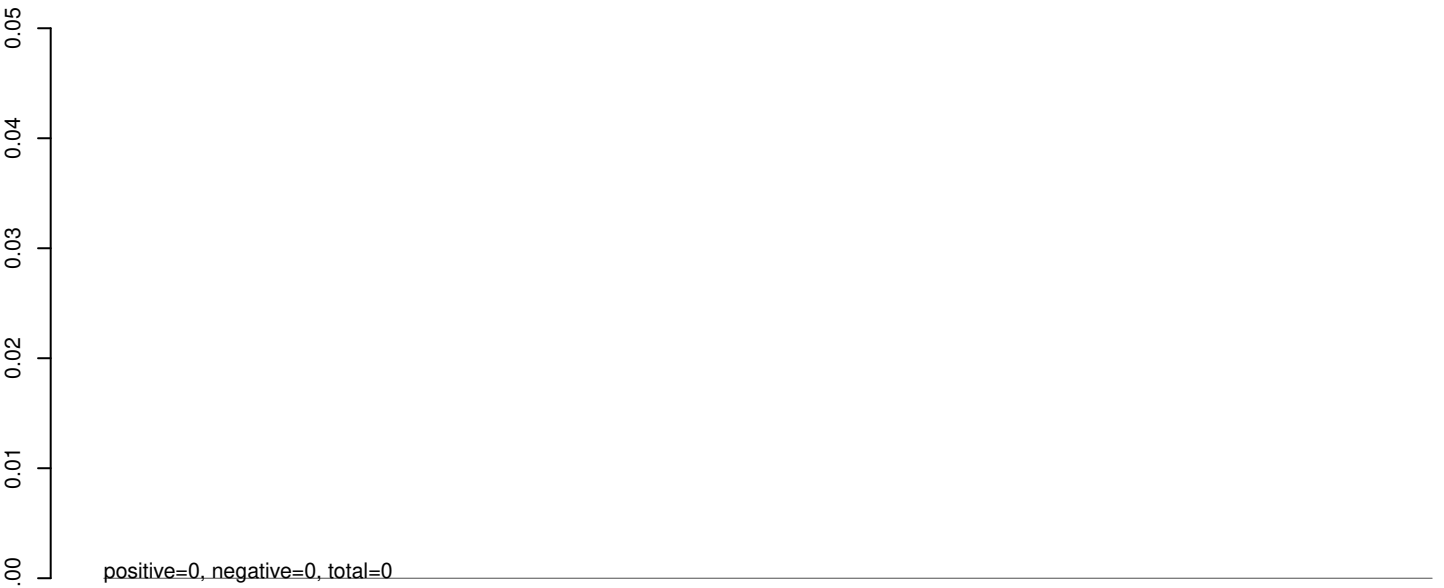
Window size=25, length=10723, virus@NC_012533.1-Kedougou_virus:1-10723

0 2000 4000 6000 8000 10000

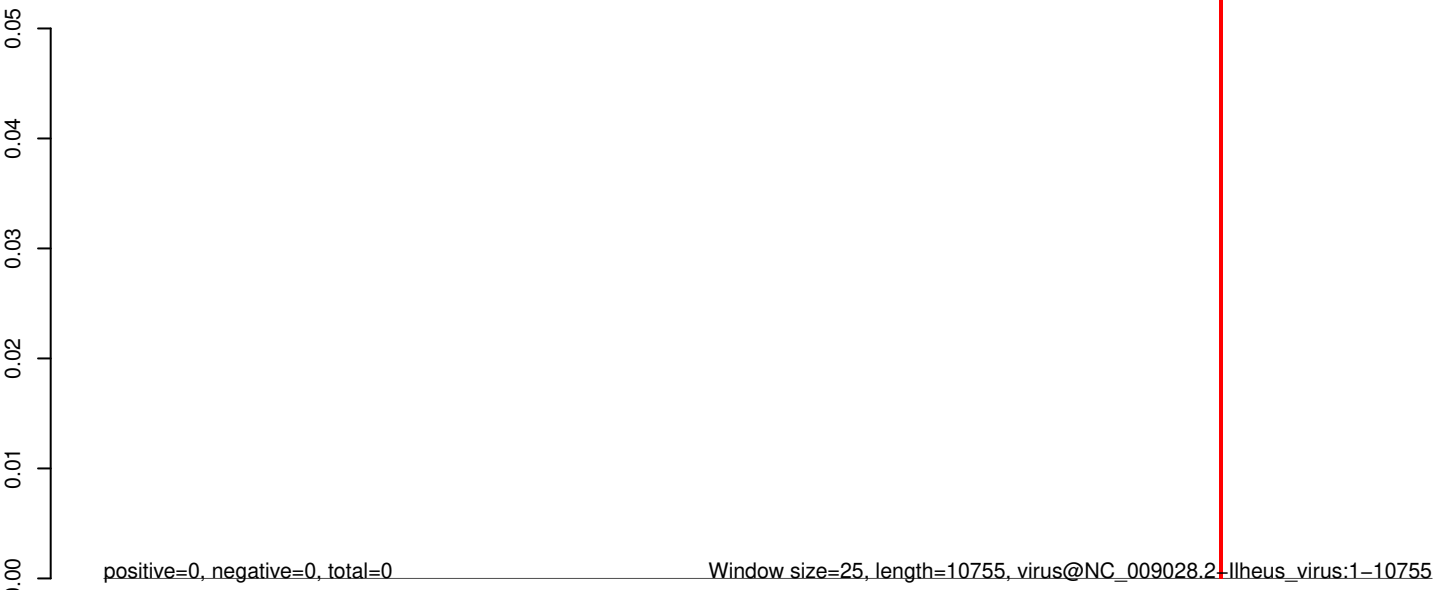
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep

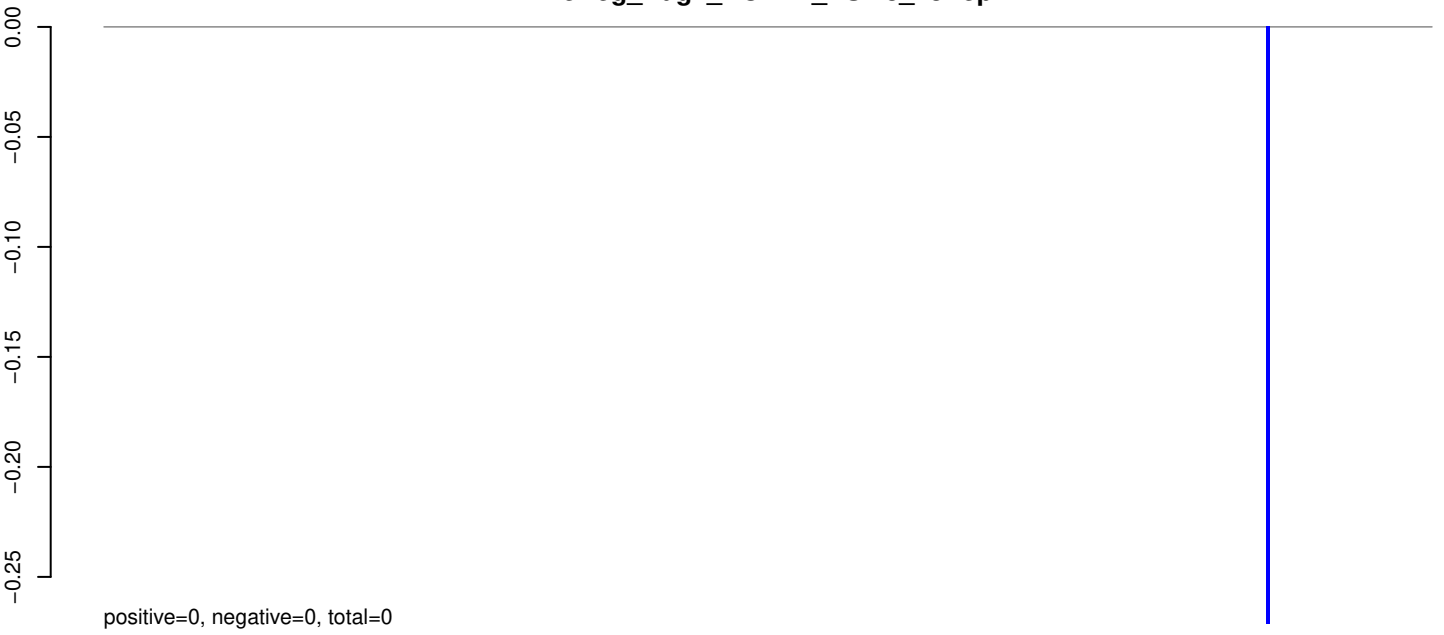


AeAeg_Aag2_BUNV1_ES.rep

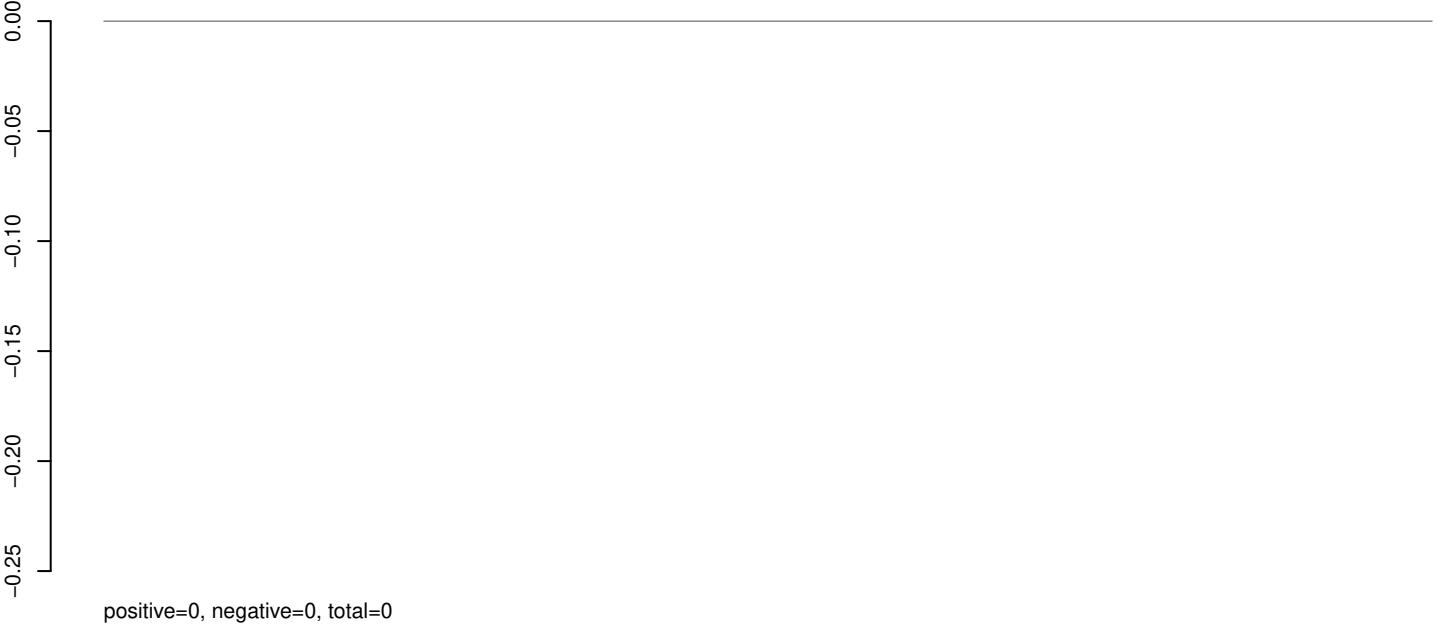


0 2000 4000 6000 8000 10000

AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



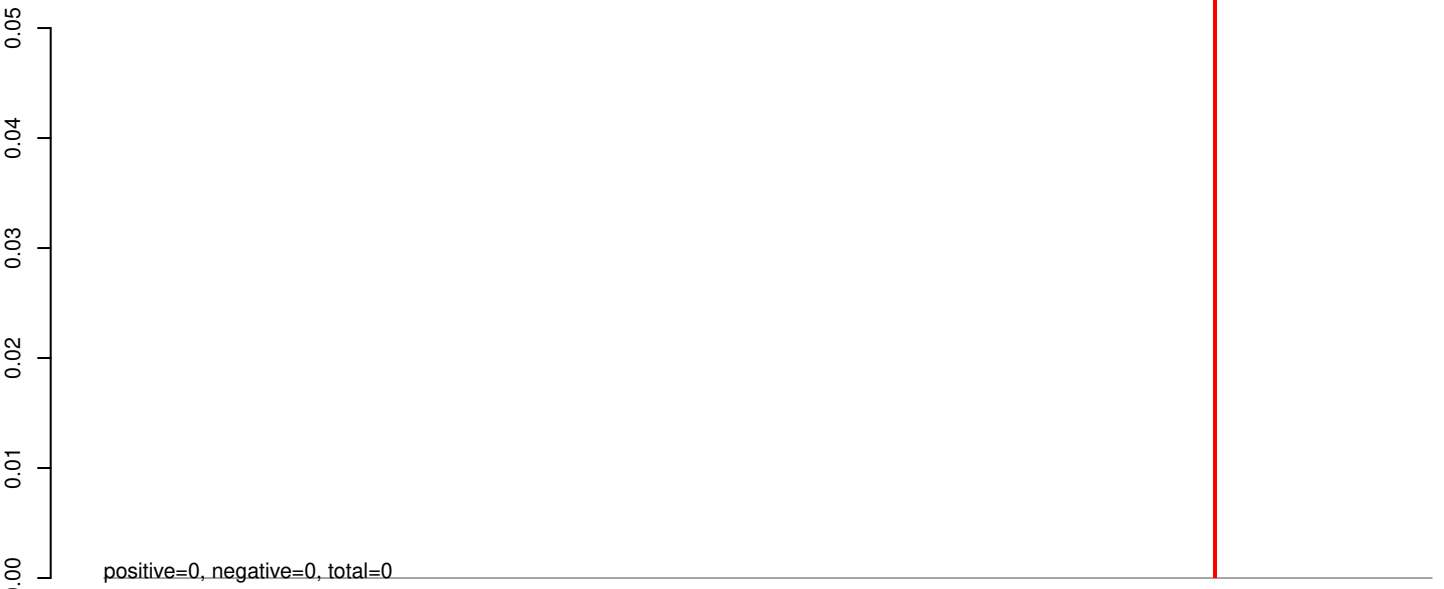
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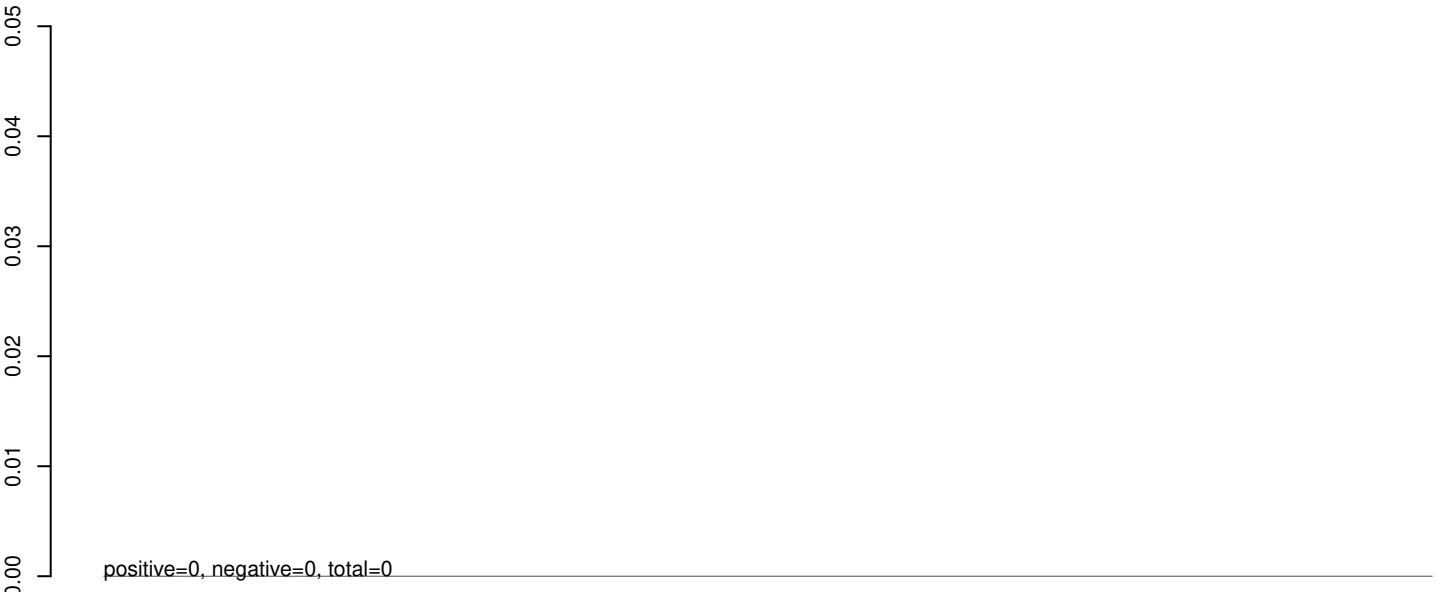
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0 2000 4000 6000 8000 10000

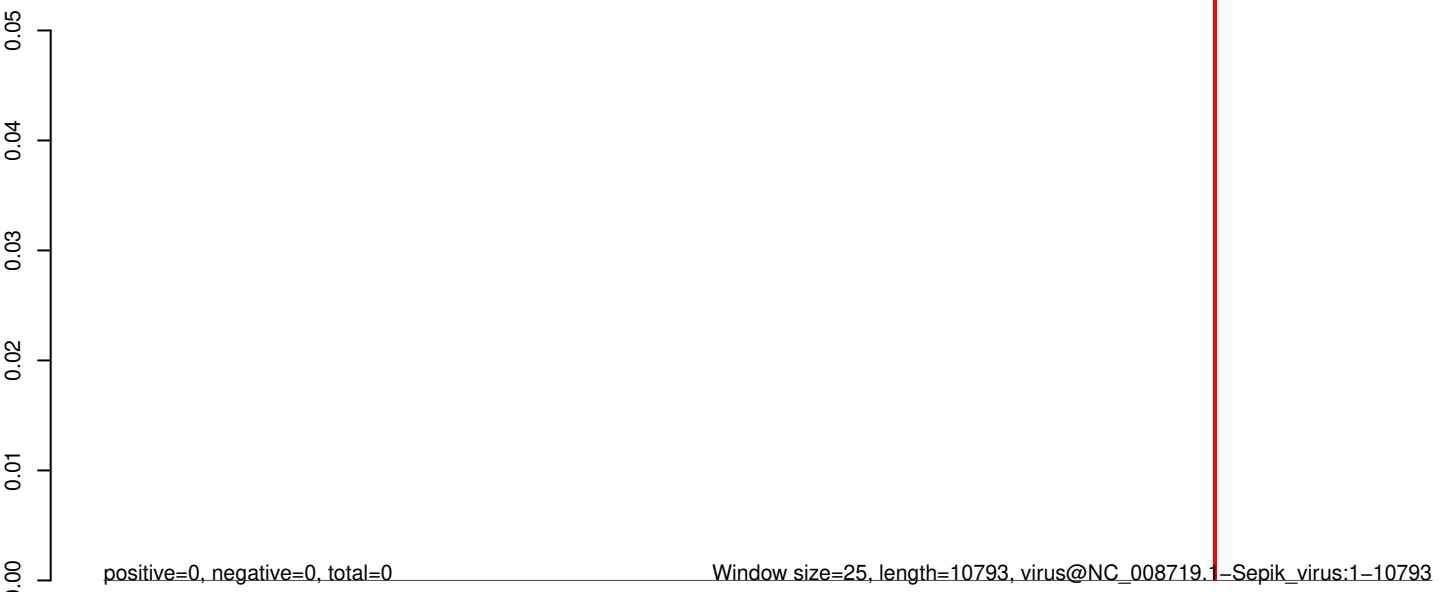
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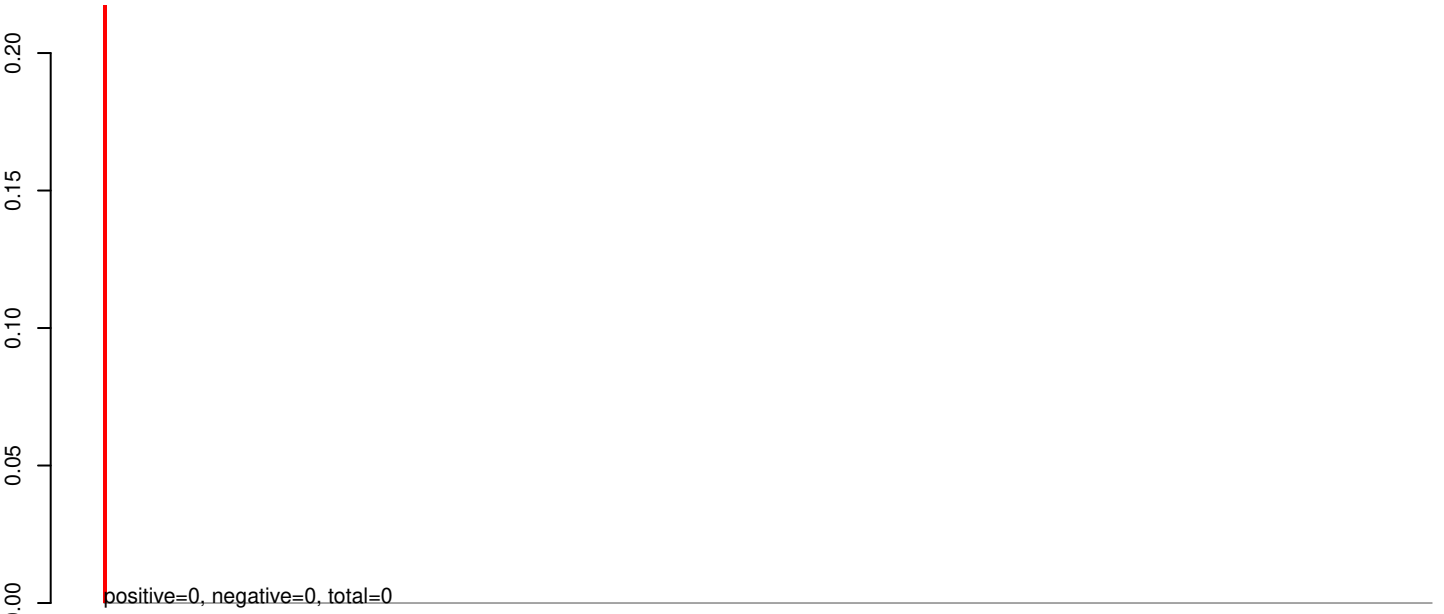
AeAeg_Aag2_BUNV1_ES.24_35.rep



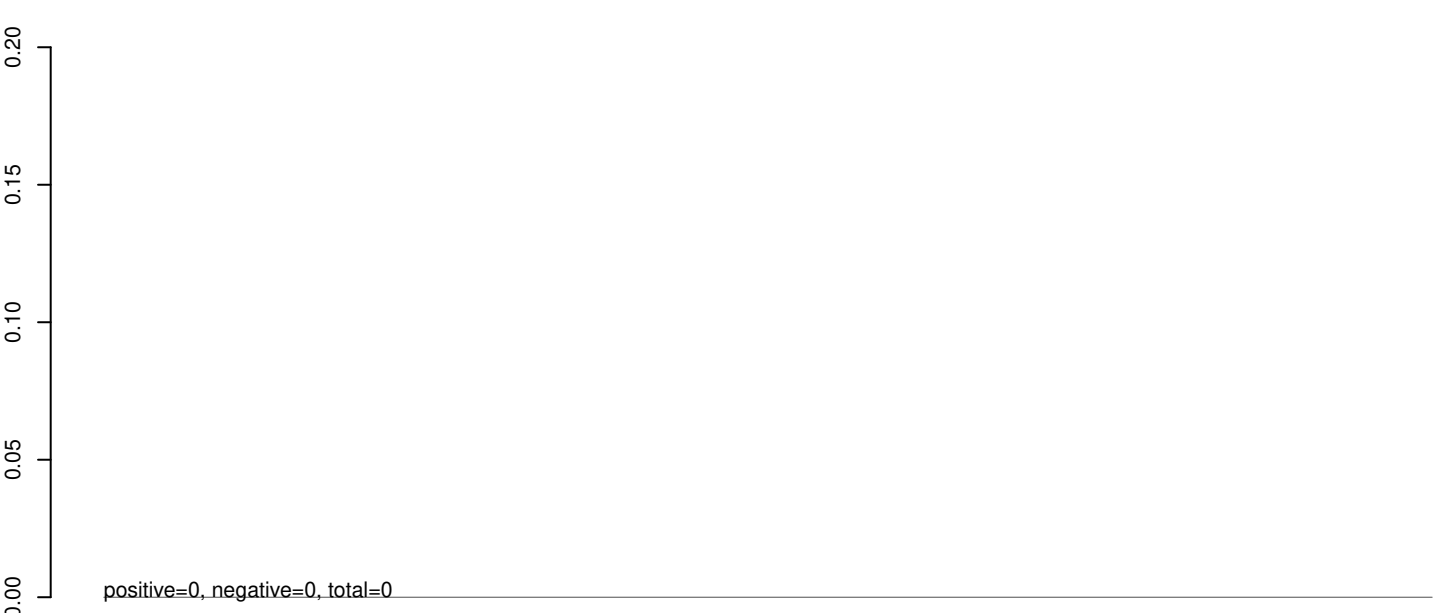
AeAeg_Aag2_BUNV1_ES.rep



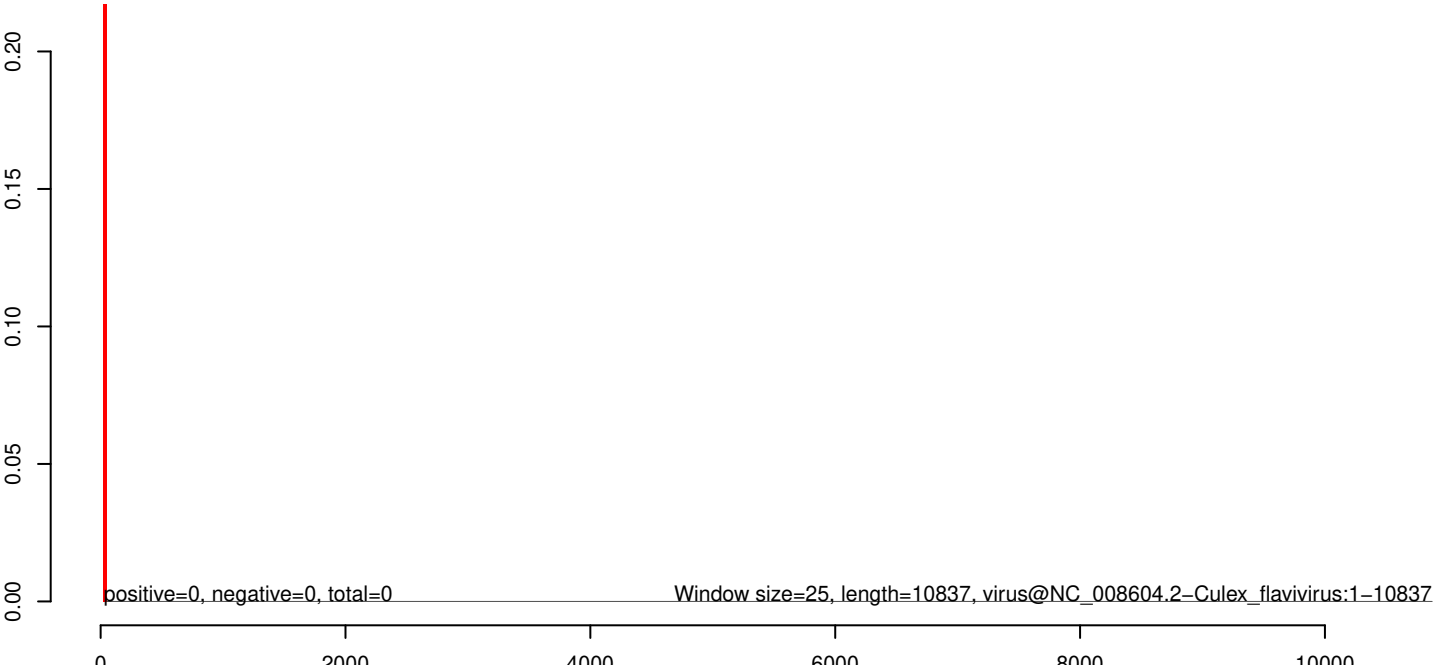
AeAeg_Aag2_BUNV1_ES.18_23.rep



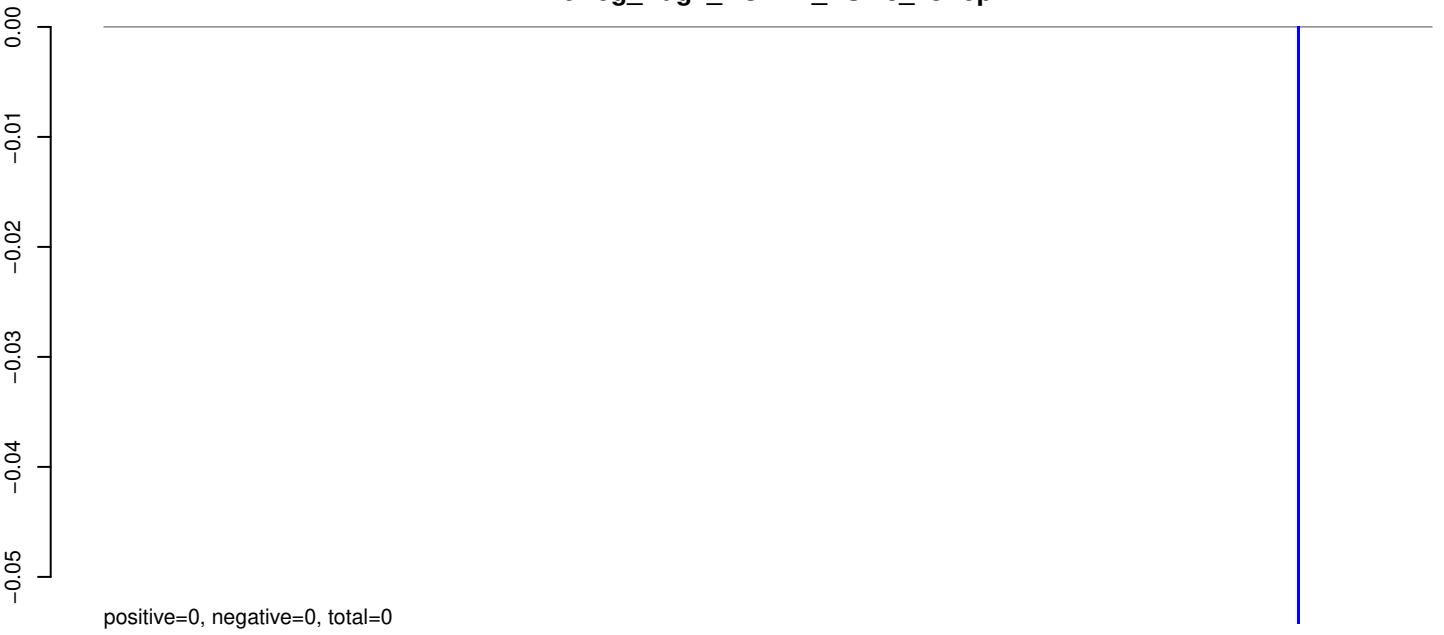
AeAeg_Aag2_BUNV1_ES.24_35.rep



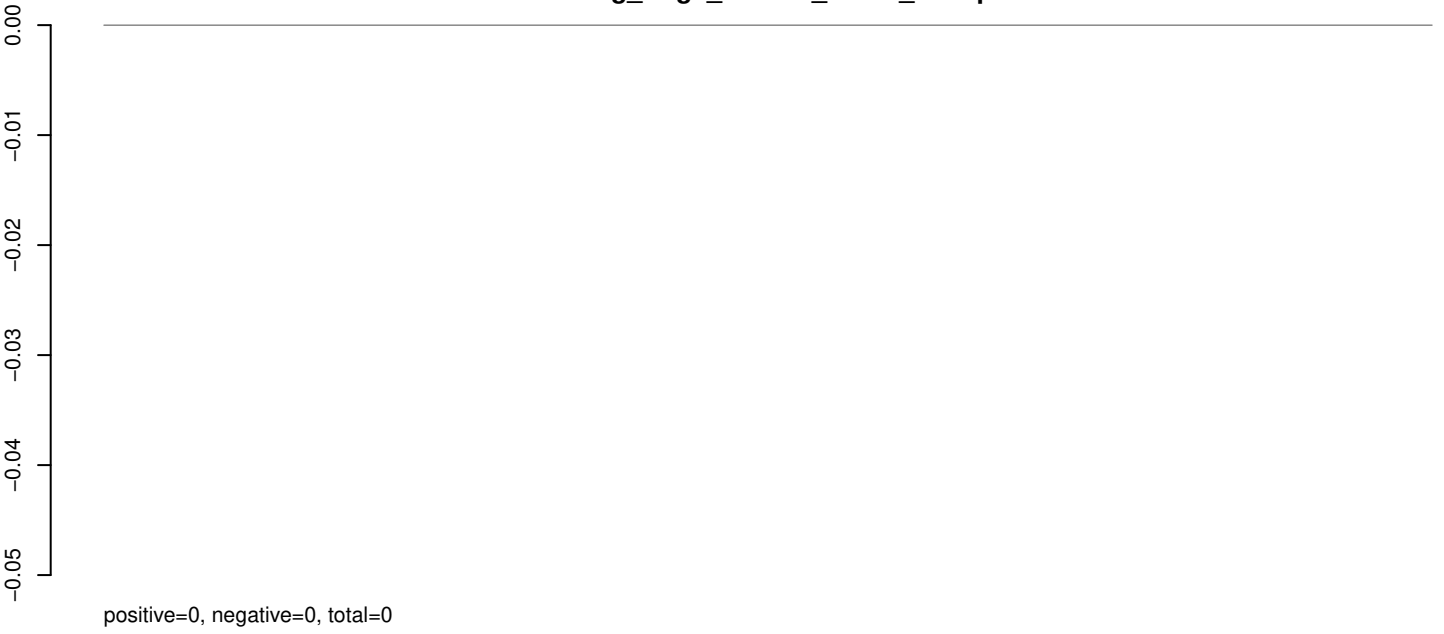
AeAeg_Aag2_BUNV1_ES.rep



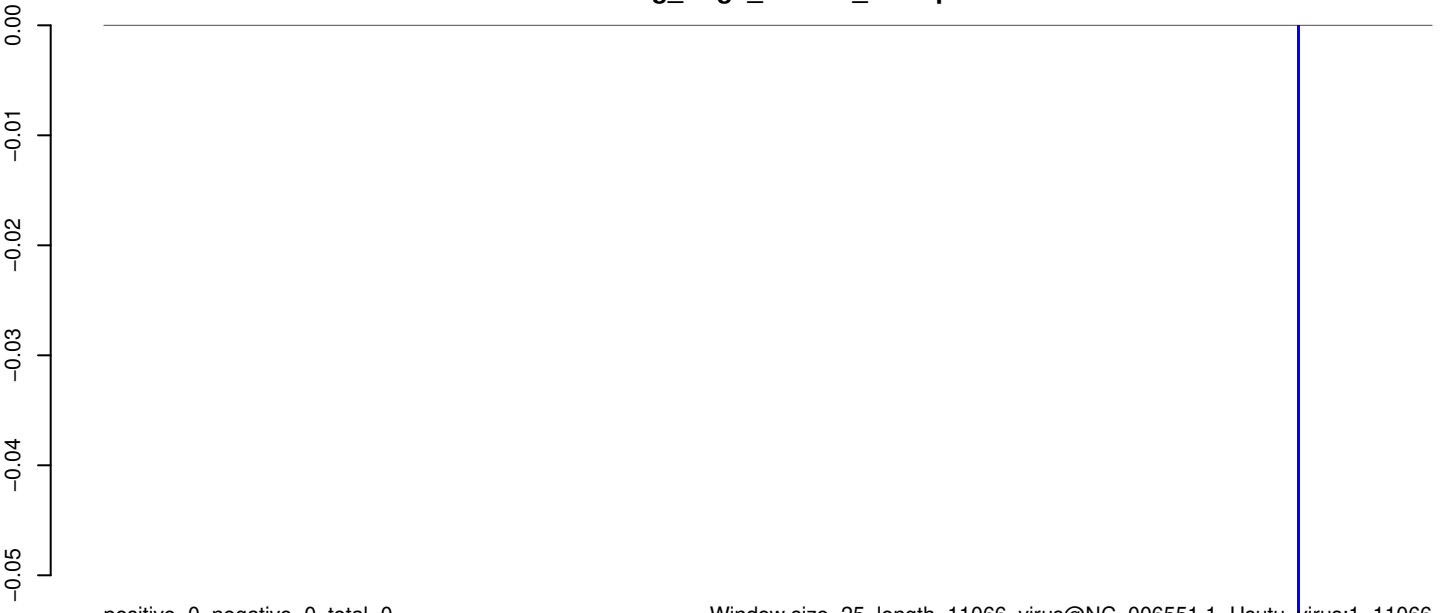
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



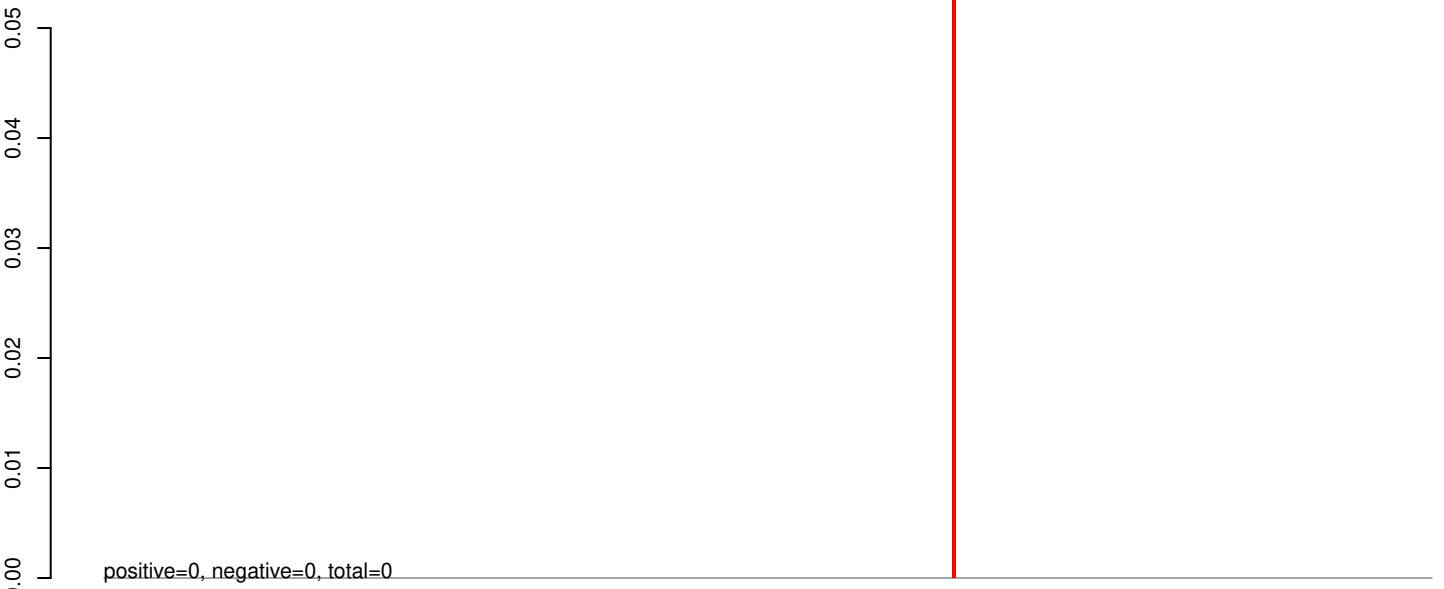
AeAeg_Aag2_BUNV1_ES.rep



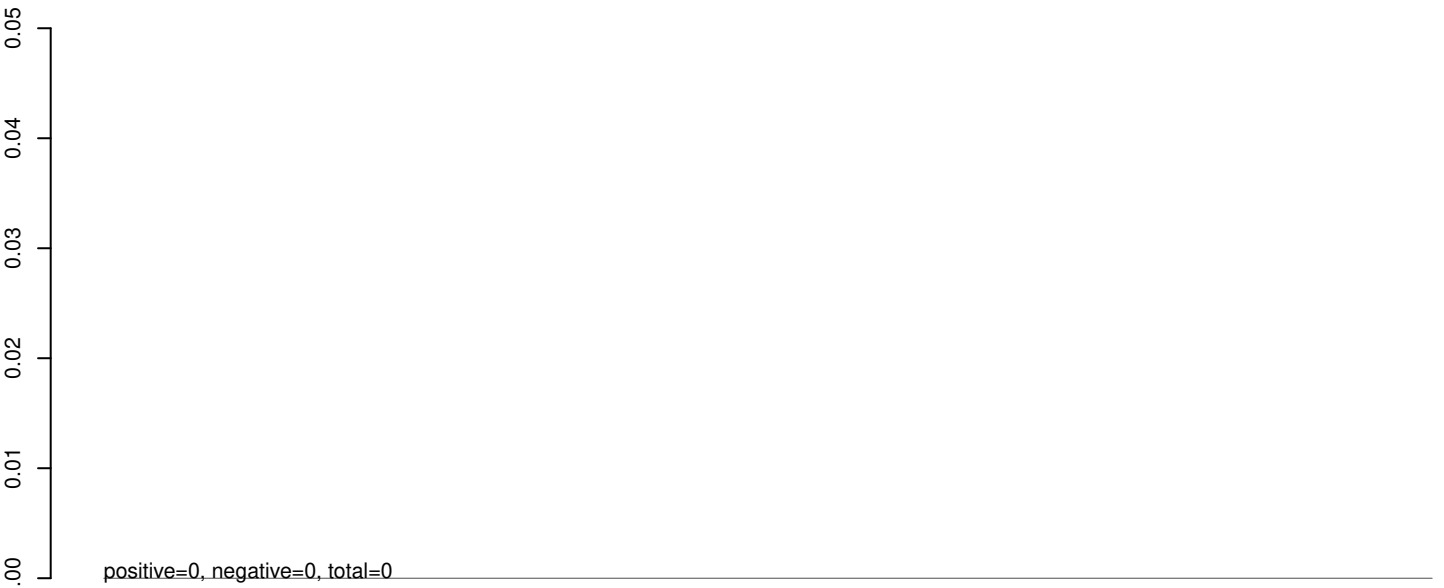
Window size=25, length=11066, virus@NC_006551.1-Usutu_virus:1-11066

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep

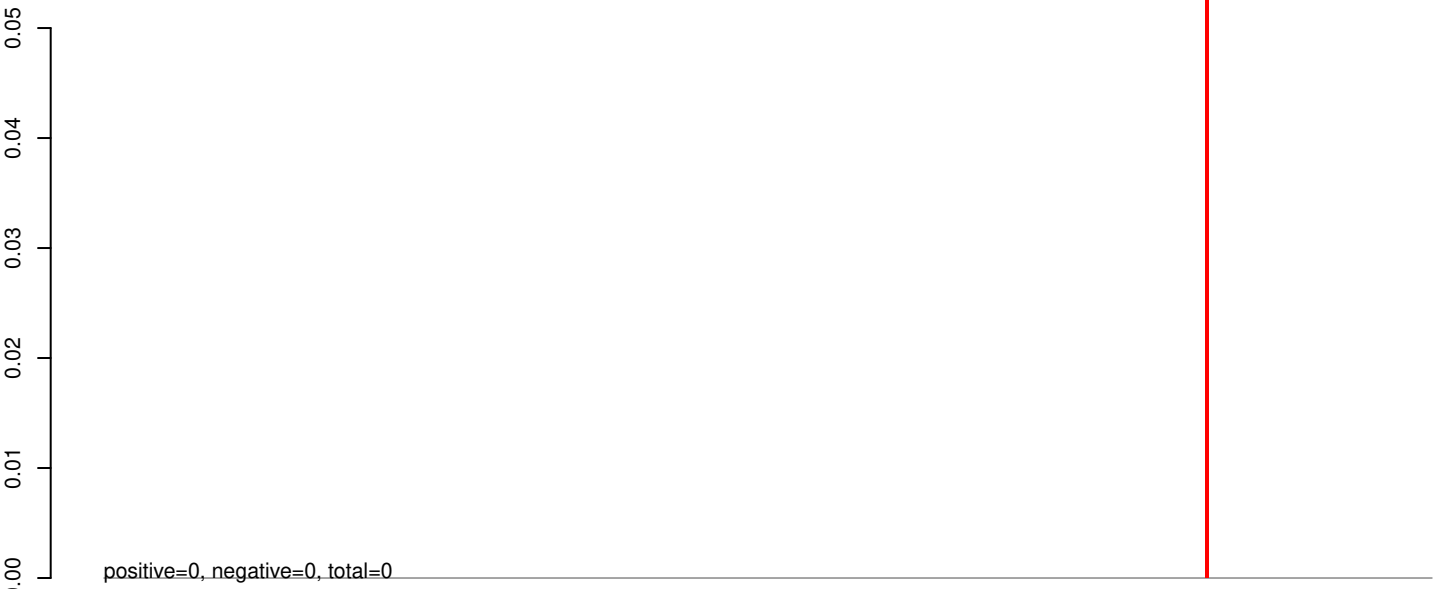


AeAeg_Aag2_BUNV1_ES.rep

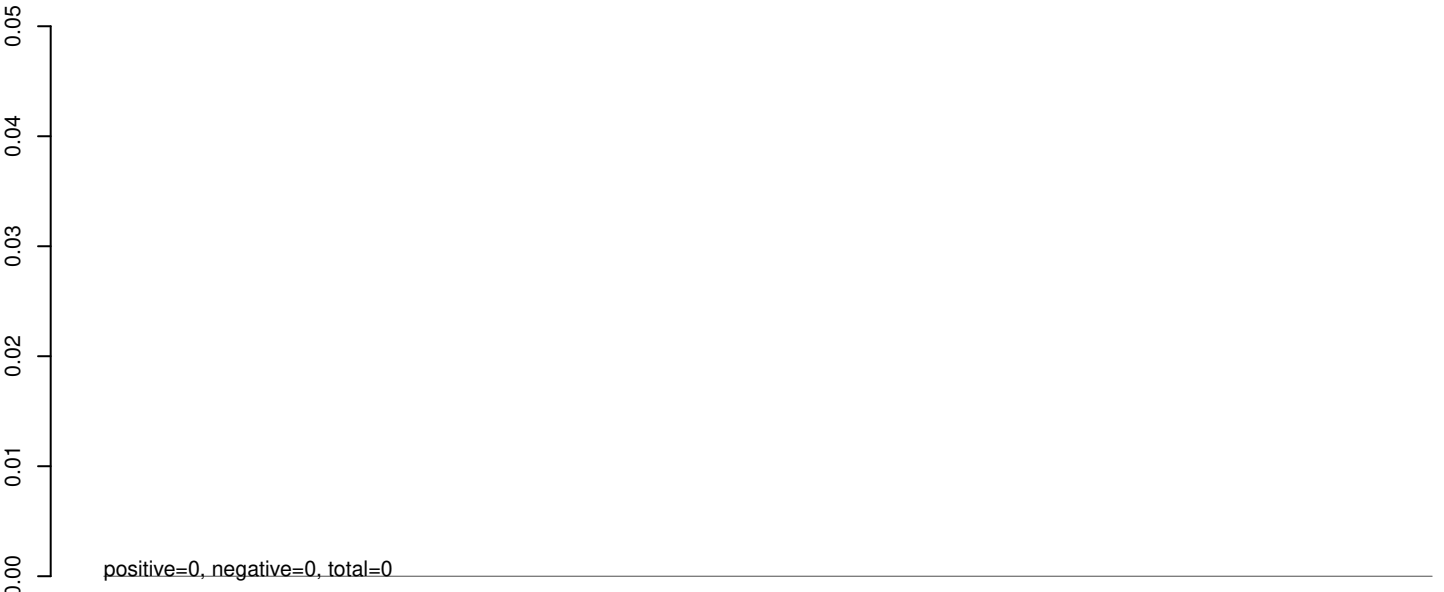


0 2000 4000 6000 8000 10000

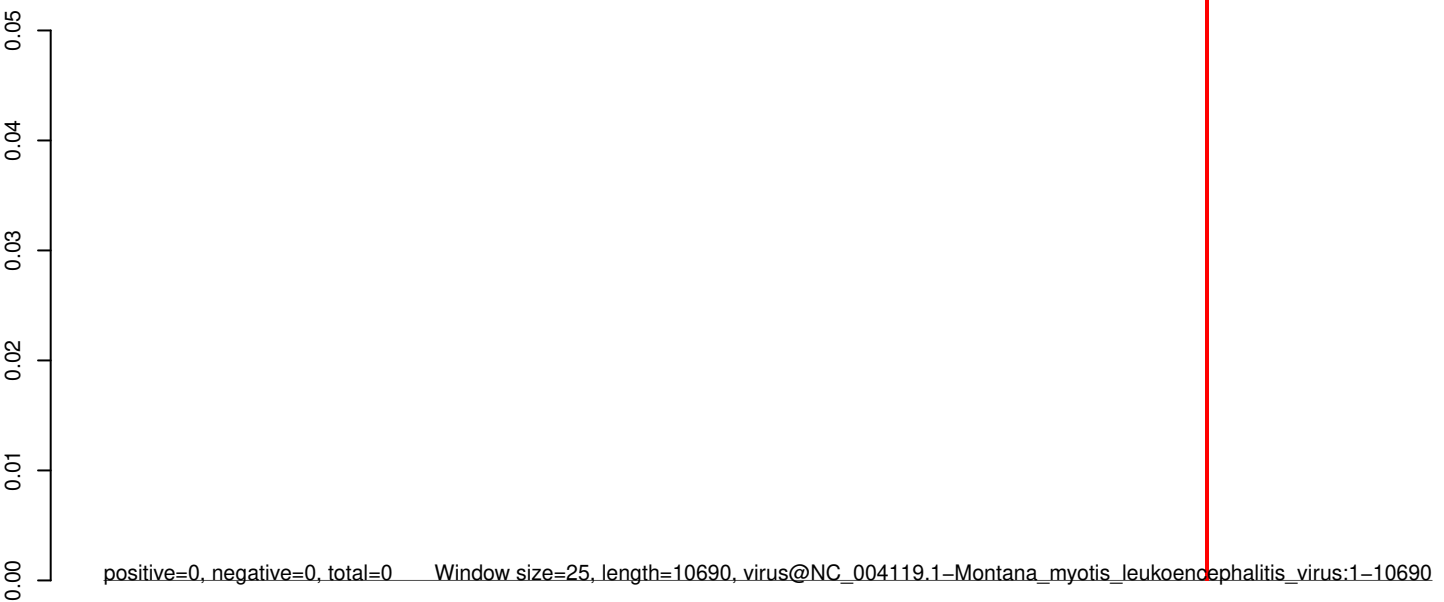
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep

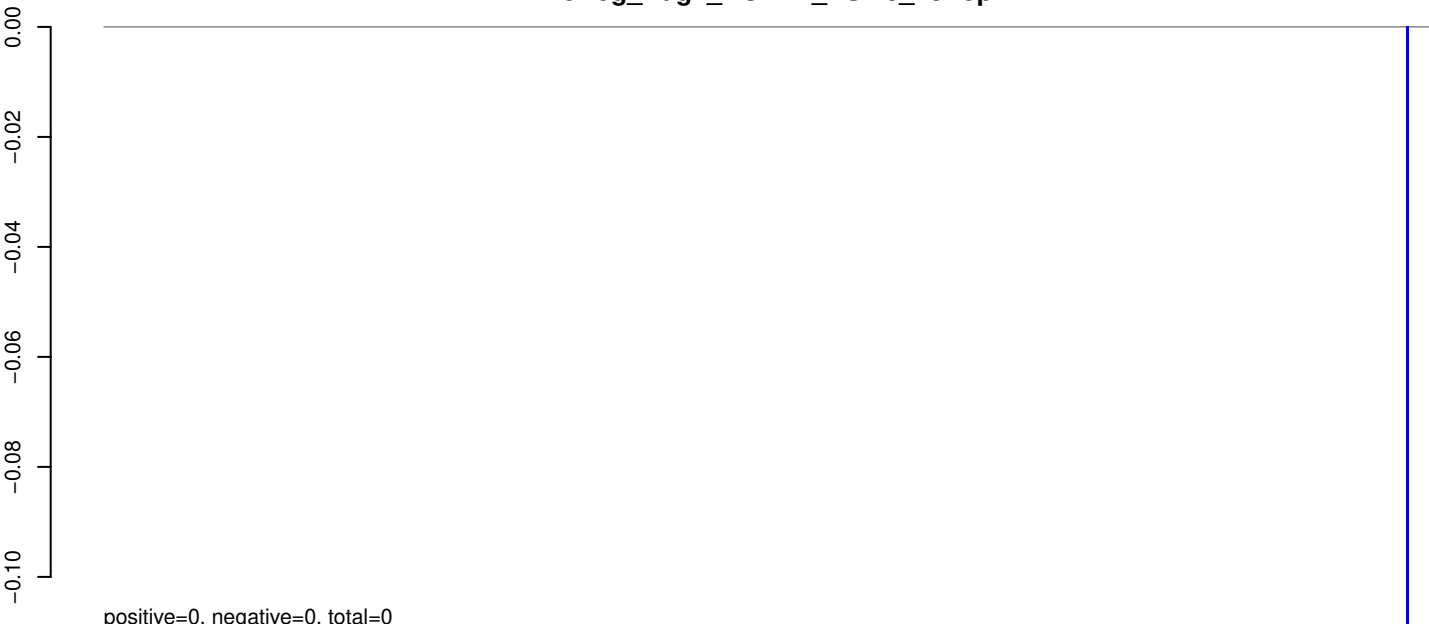


AeAeg_Aag2_BUNV1_ES.rep



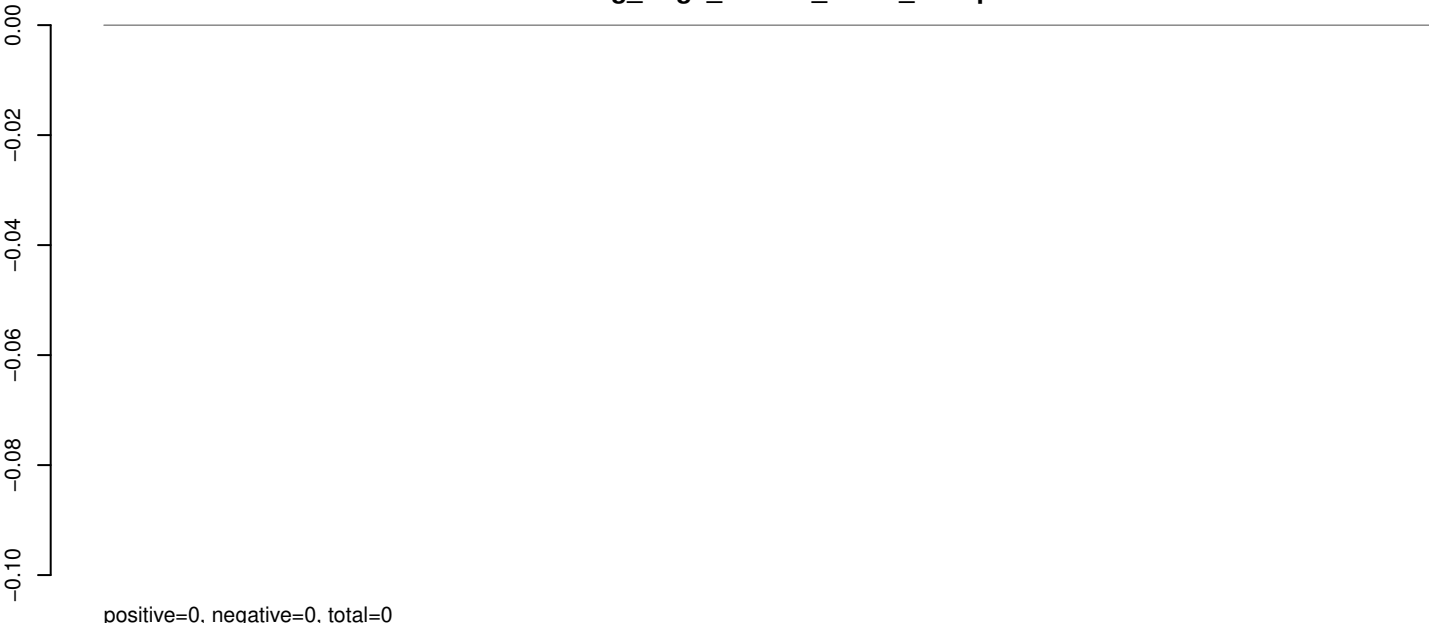
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AeAeg_Aag2_BUNV1_ES.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BUNV1_ES.24_35.rep



positive=0, negative=0, total=0

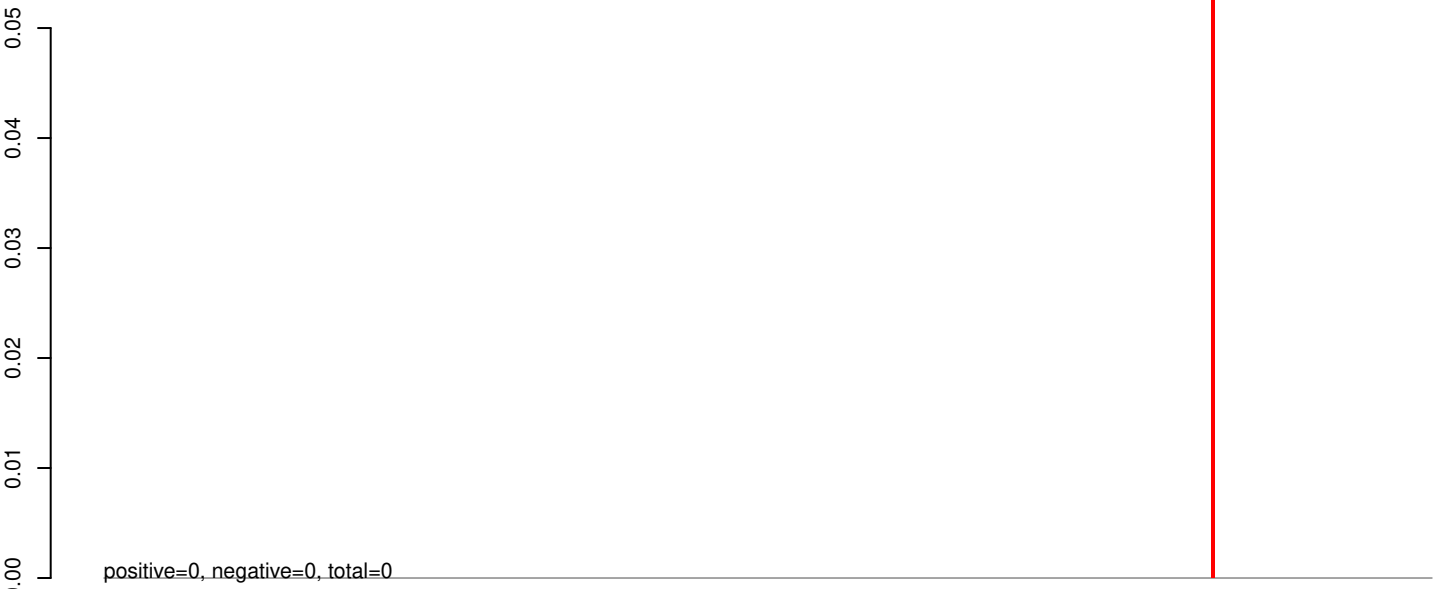
AeAeg_Aag2_BUNV1_ES.rep



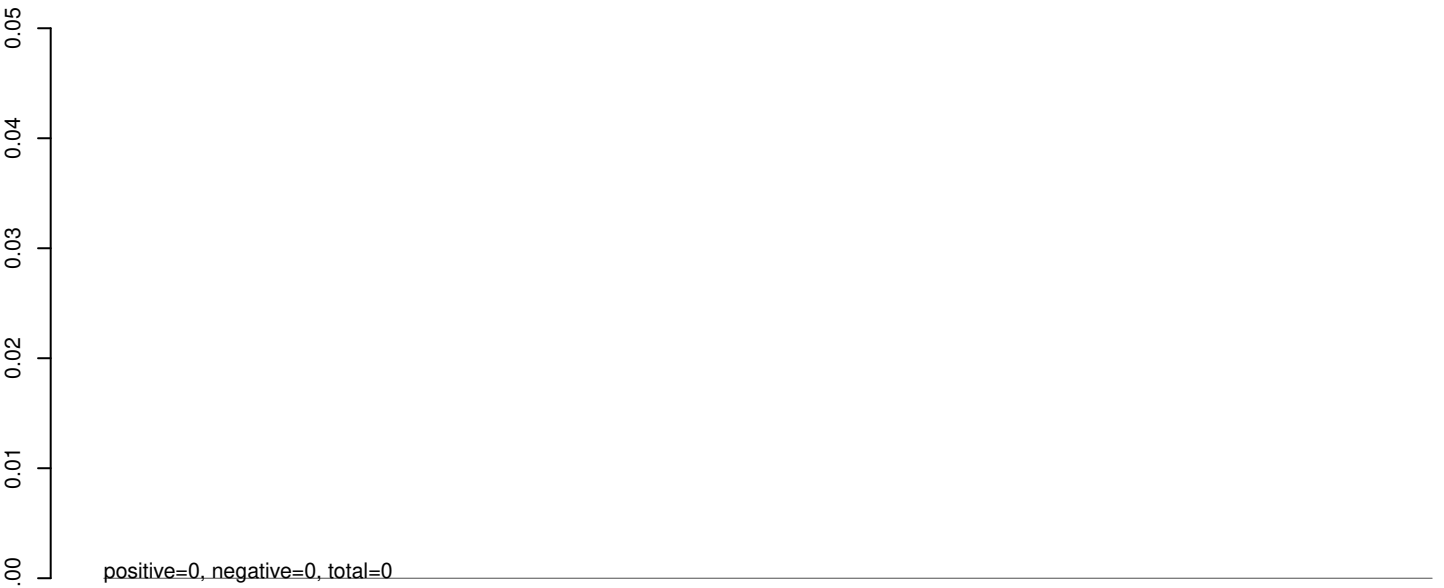
positive=0, negative=0, total=0 Window size=25, length=11484, virus@NC_003908.1-Western_equine_encephalomyelitis_virus:1-11484

0 2000 4000 6000 8000 10000

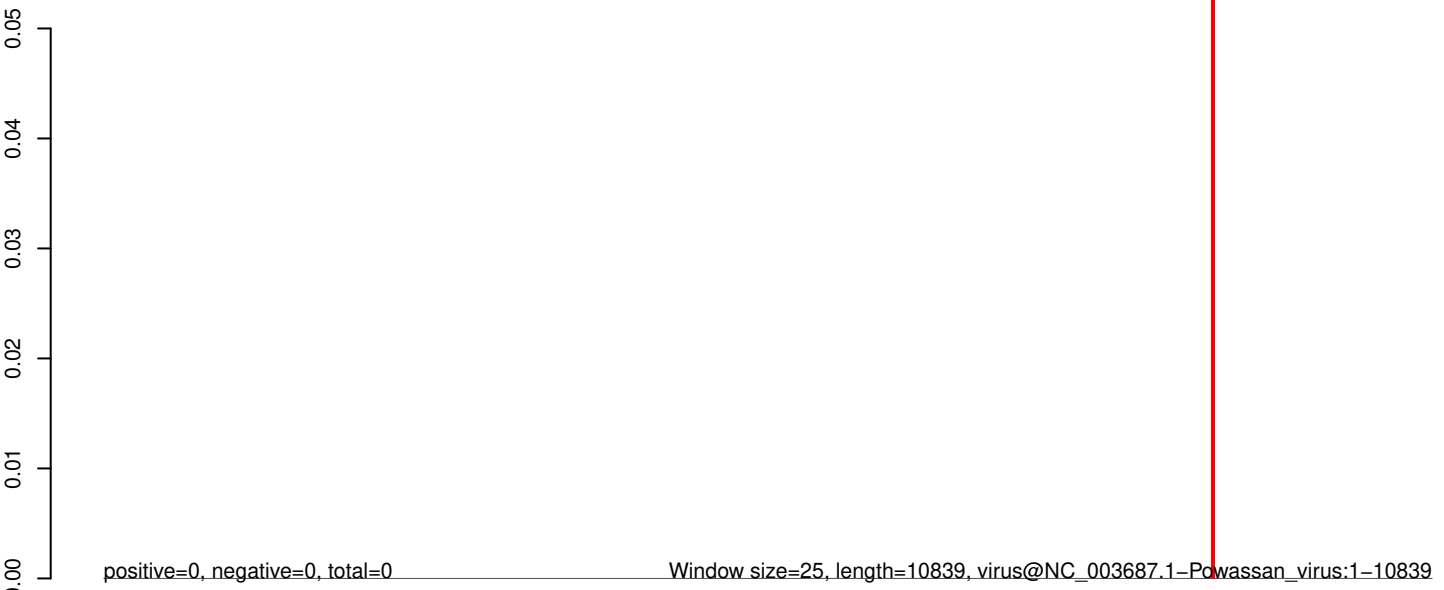
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



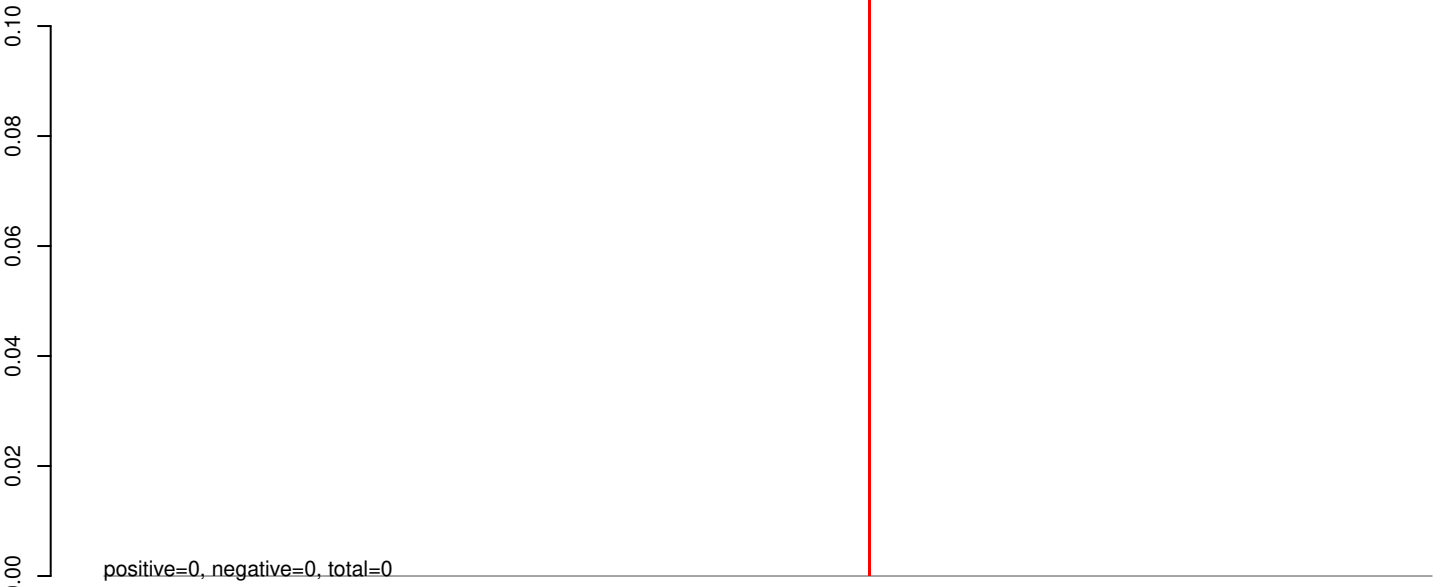
AeAeg_Aag2_BUNV1_ES.rep



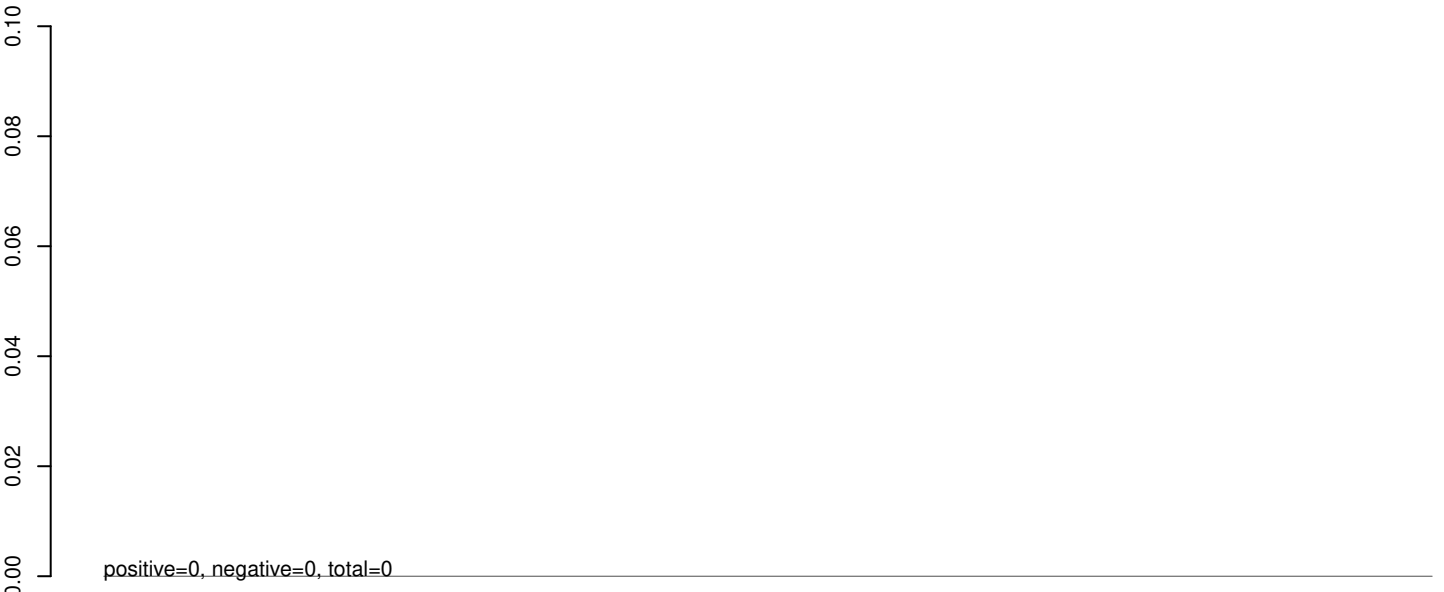
Window size=25, length=10839, virus@NC_003687.1-Powassan_virus:1-10839

0 2000 4000 6000 8000 10000

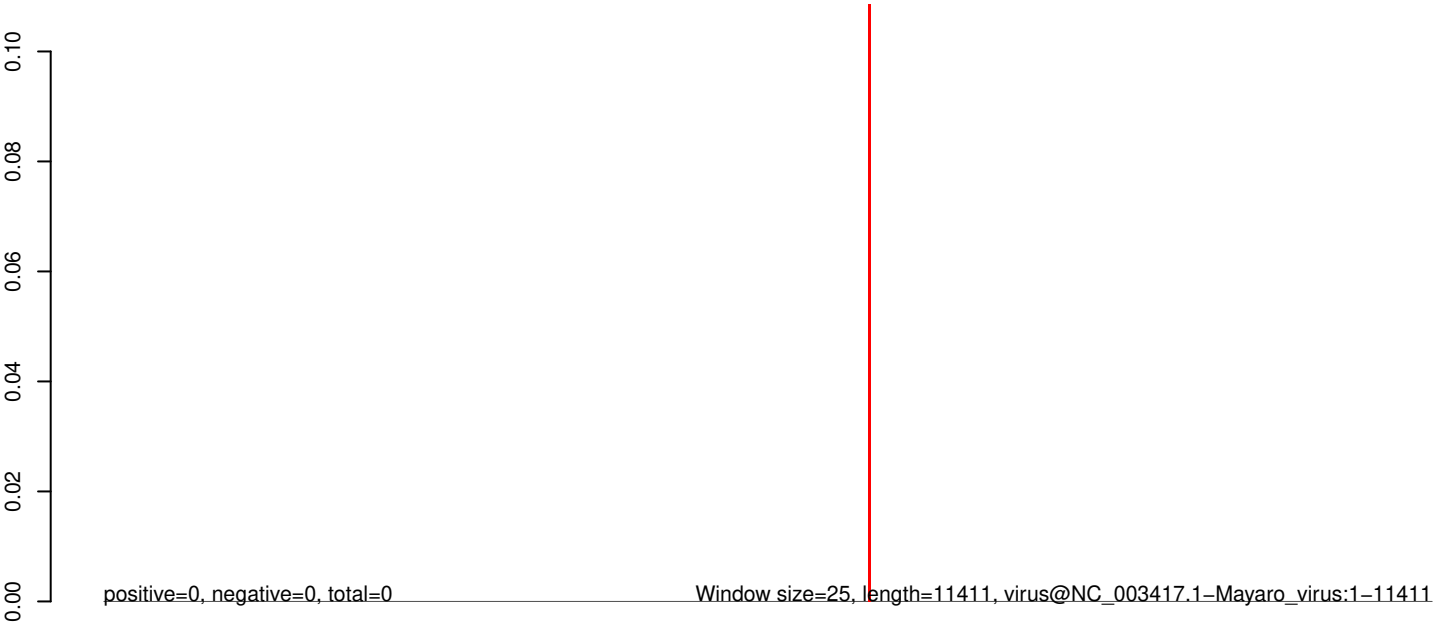
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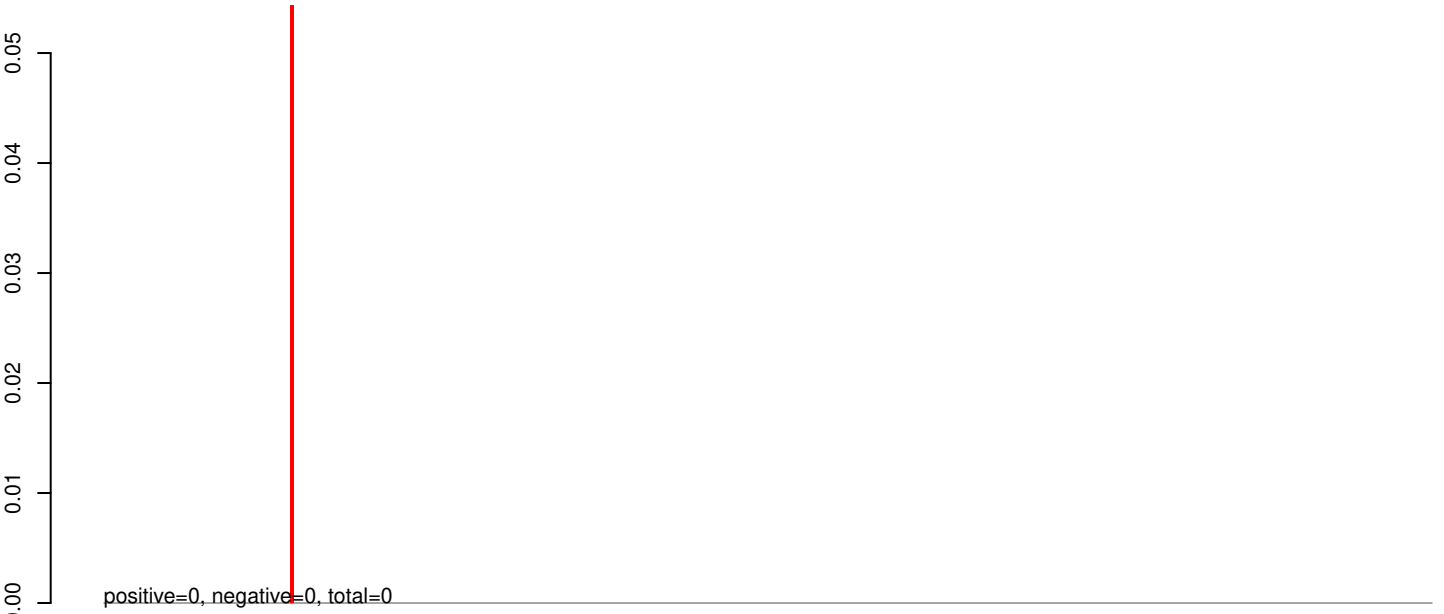
AeAeg_Aag2_BUNV1_ES.24_35.rep



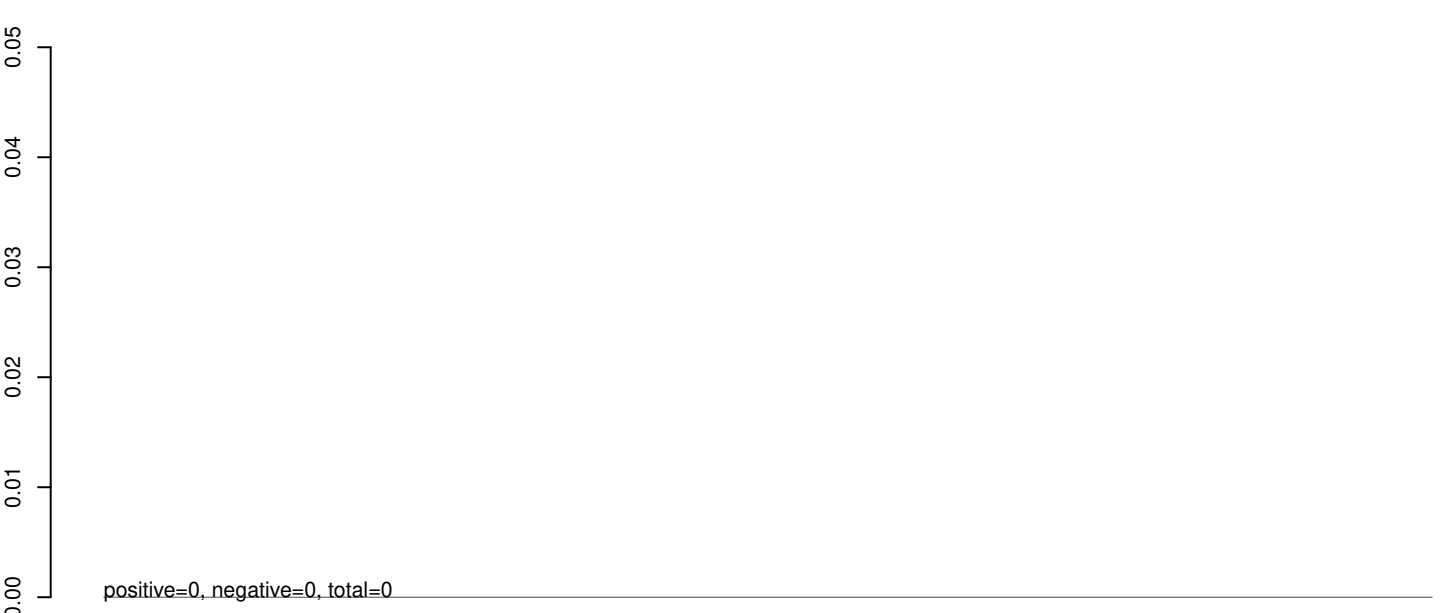
AeAeg_Aag2_BUNV1_ES.rep



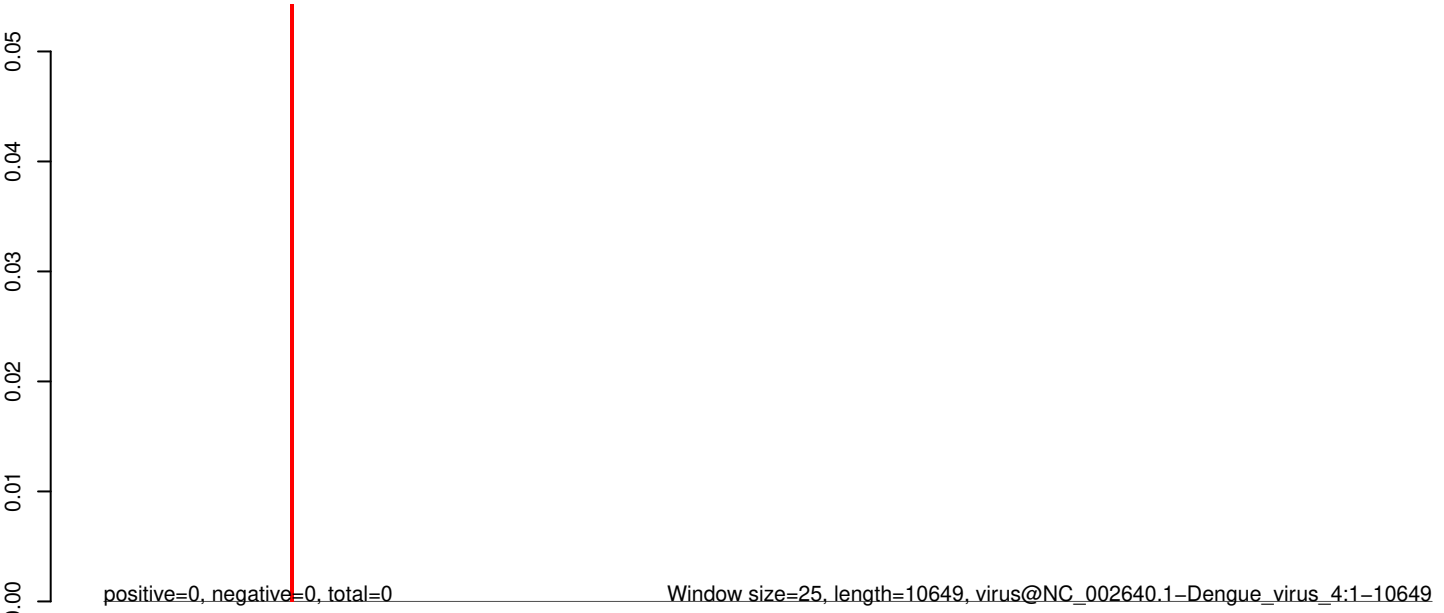
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep

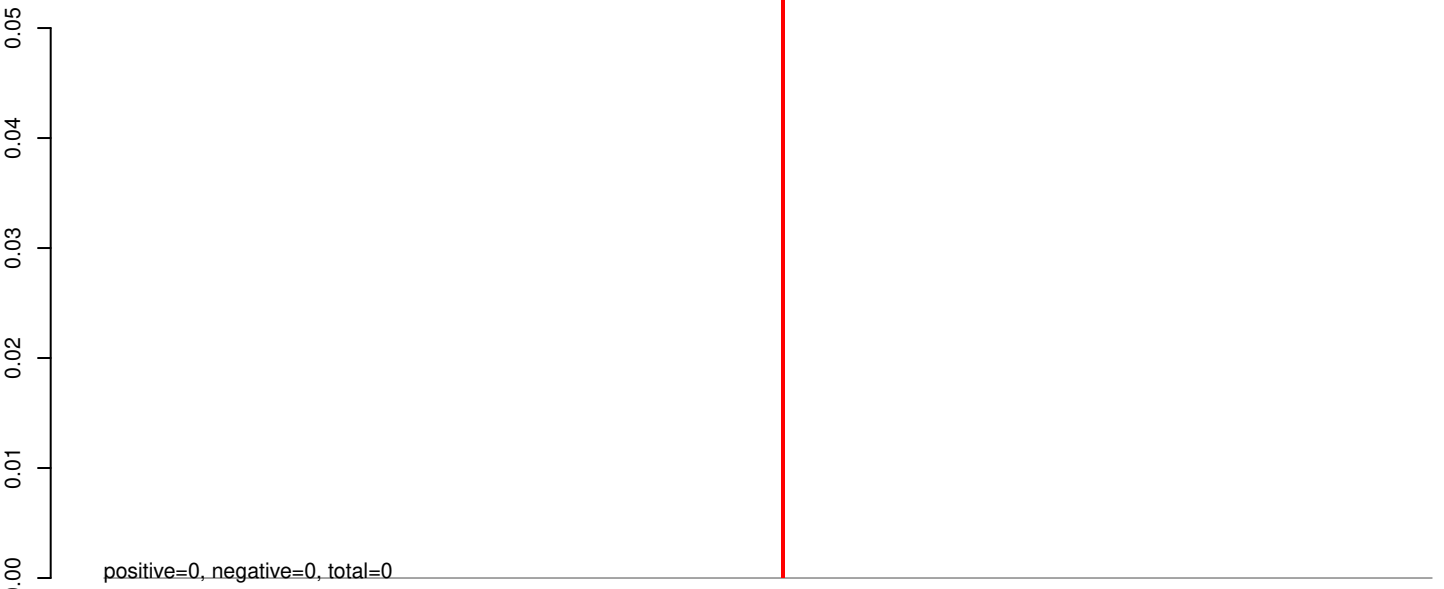


AeAeg_Aag2_BUNV1_ES.rep

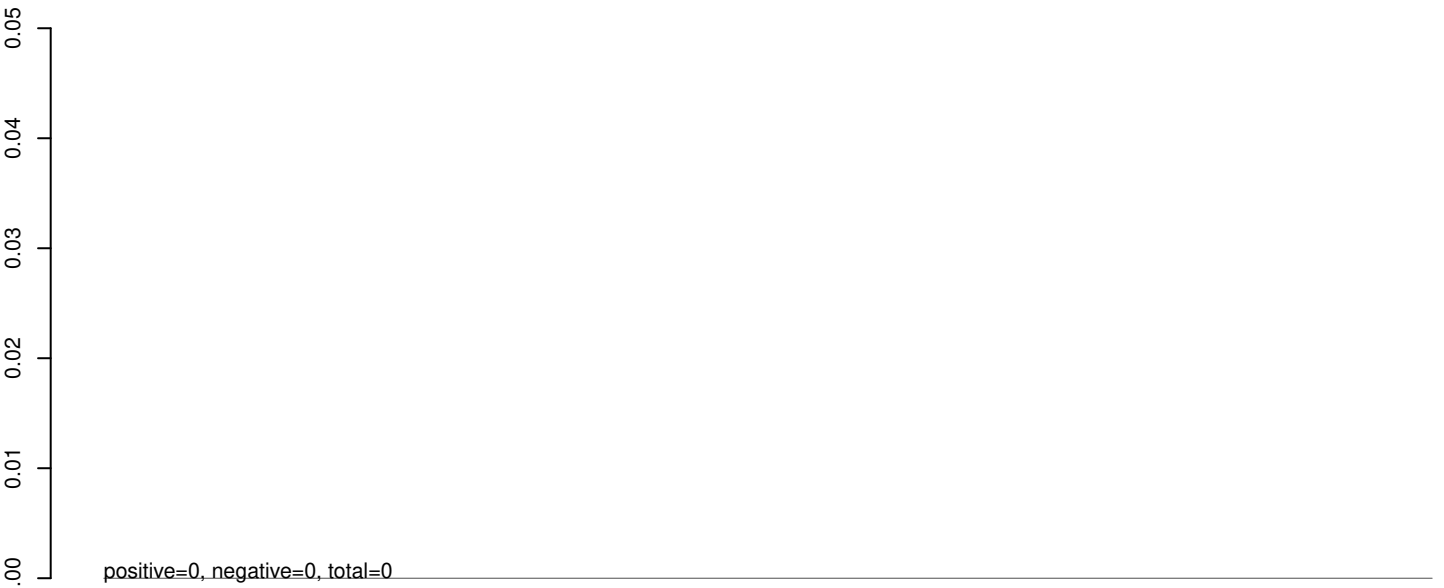


0 2000 4000 6000 8000 10000

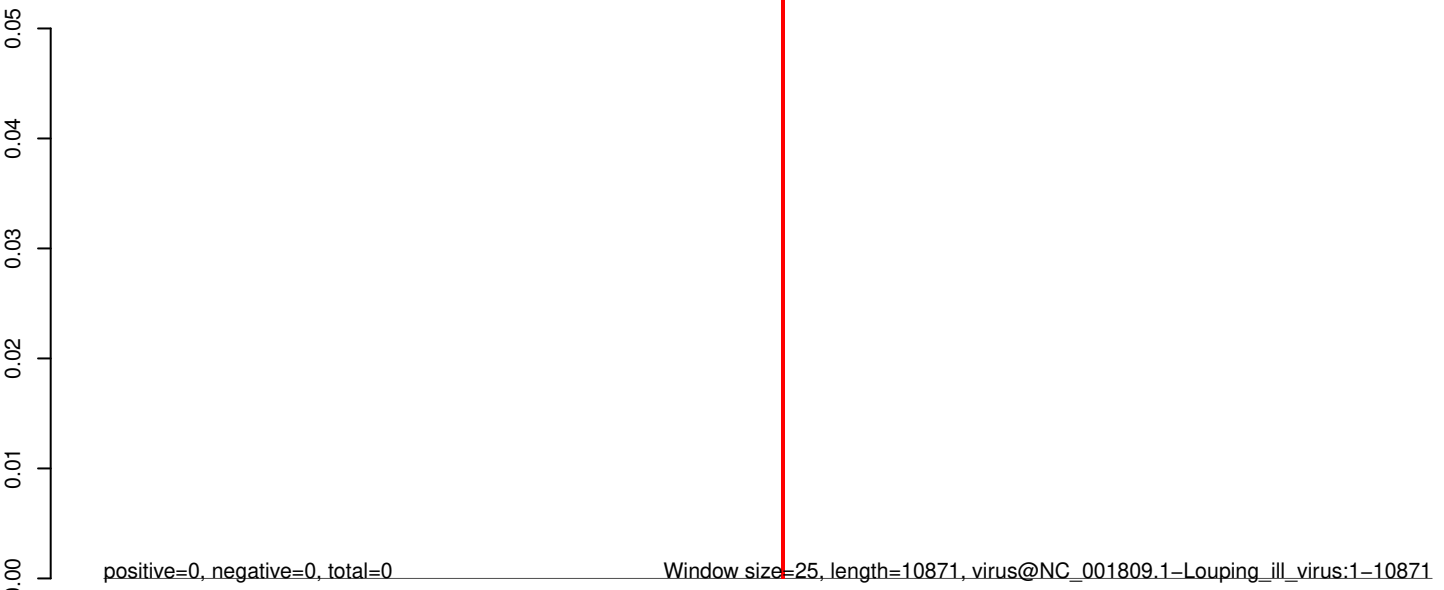
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



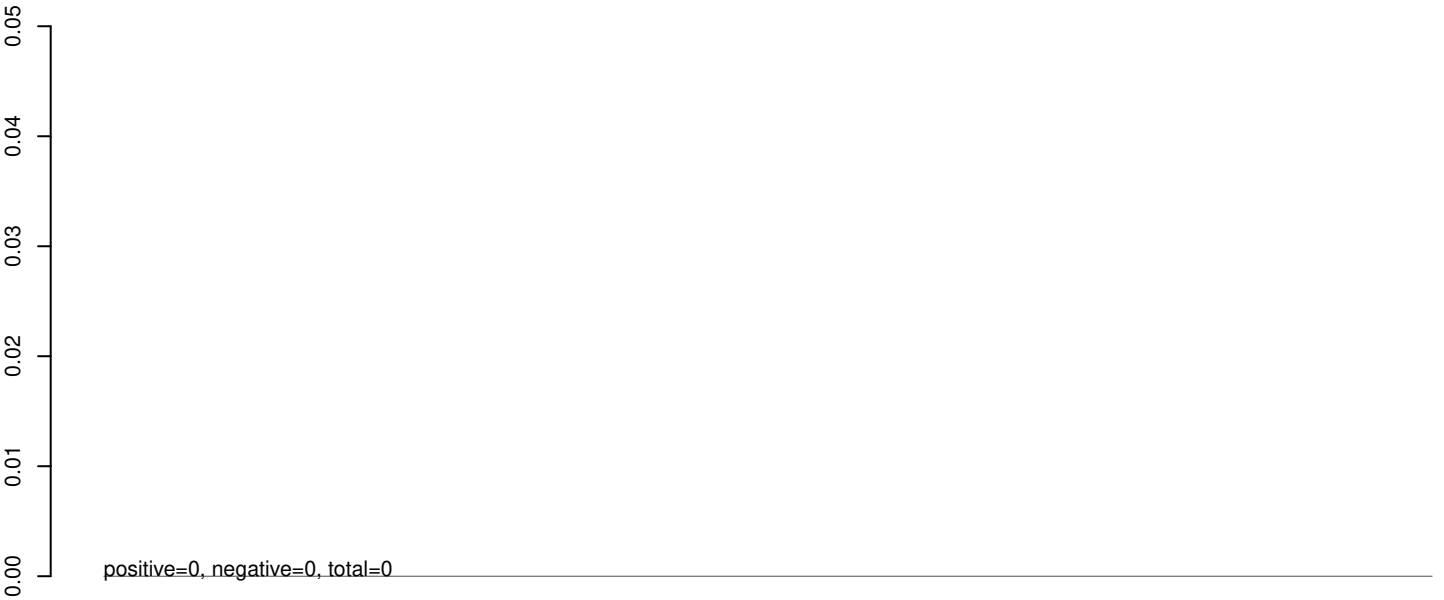
AeAeg_Aag2_BUNV1_ES.rep



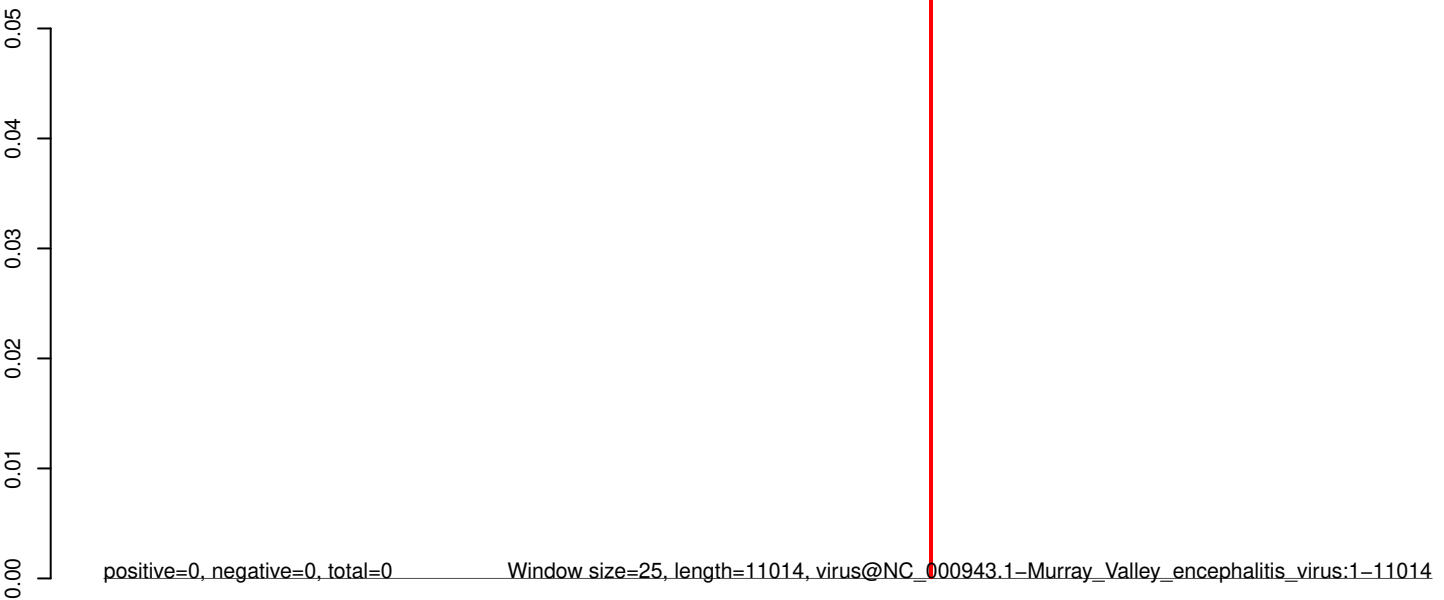
AeAeg_Aag2_BUNV1_ES.18_23.rep



AeAeg_Aag2_BUNV1_ES.24_35.rep



AeAeg_Aag2_BUNV1_ES.rep



0 2000 4000 6000 8000 10000

AeAeg_Aag2_BUNV1_ES.18_23.rep

0.05
0.04
0.03
0.02
0.01
0.00

positive=0, negative=0, total=0



AeAeg_Aag2_BUNV1_ES.24_35.rep

0.05
0.04
0.03
0.02
0.01
0.00

positive=0, negative=0, total=0

AeAeg_Aag2_BUNV1_ES.rep

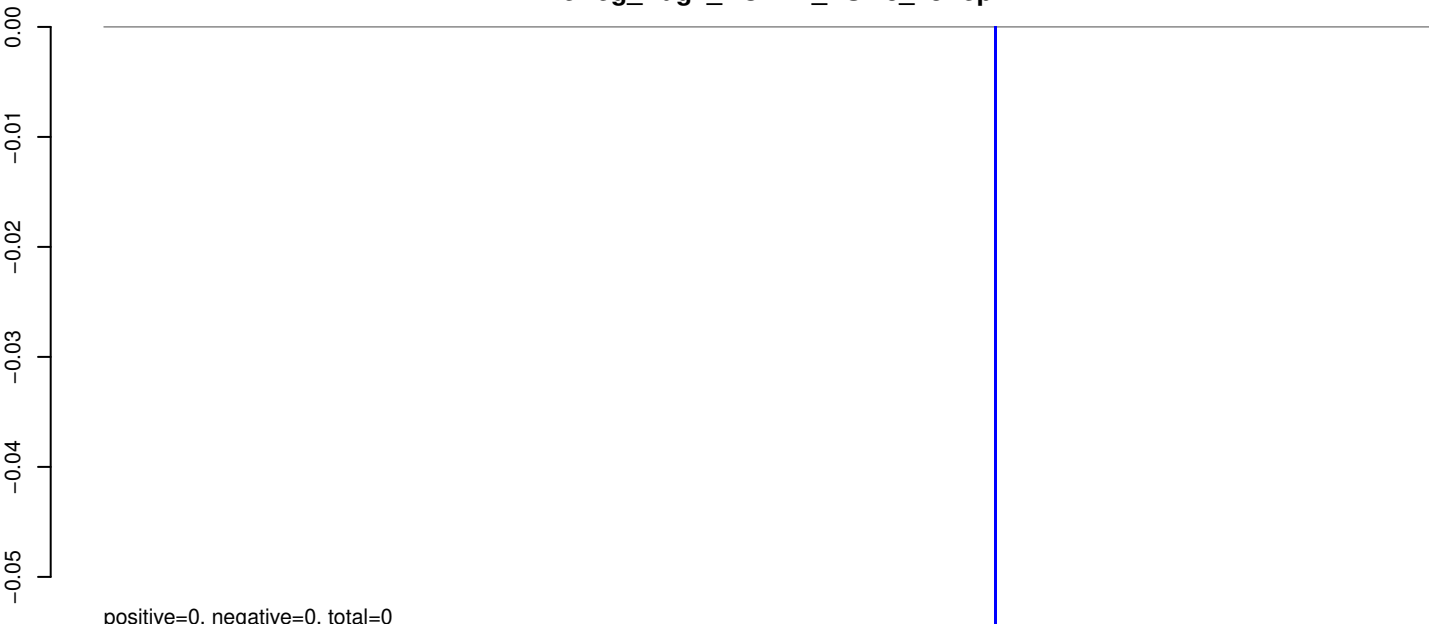
0.05
0.04
0.03
0.02
0.01
0.00

positive=0, negative=0, total=0, window size=25, length=12195, virus@MG747536.1_Oropouche_virus_isolate_BeAr_136921_segment_S_M_L :1-12195

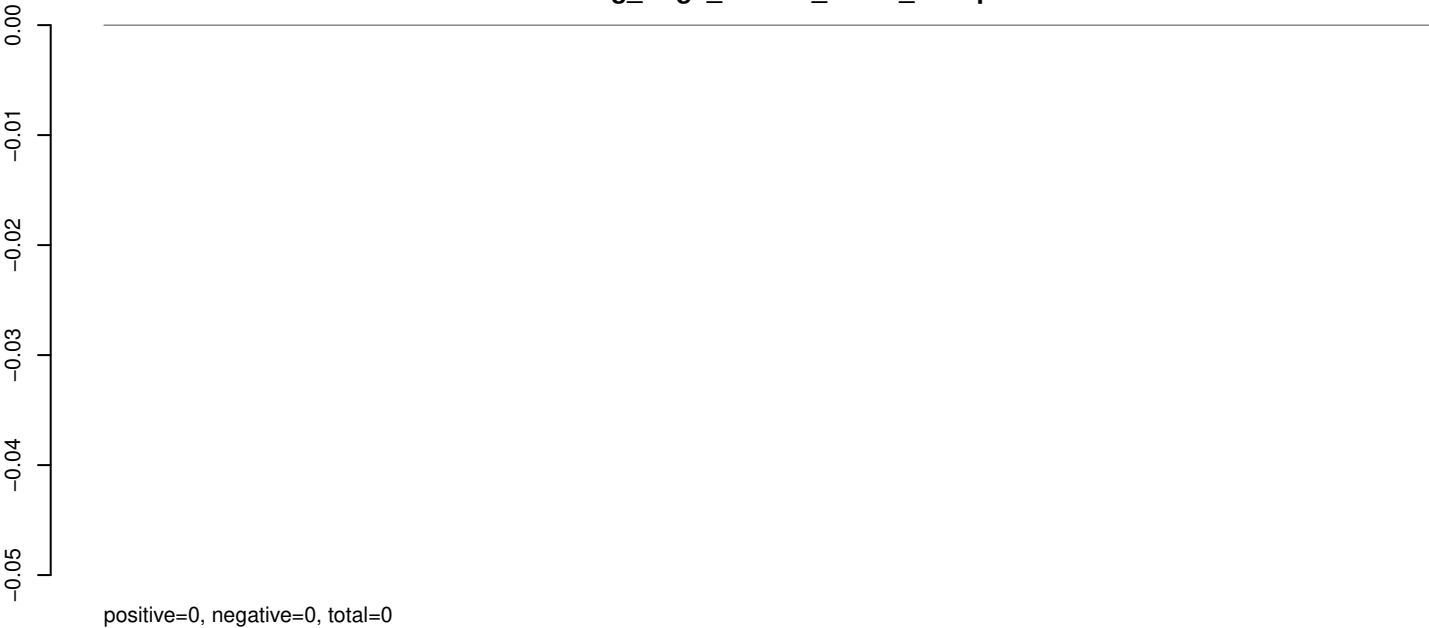


0 2000 4000 6000 8000 10000 12000

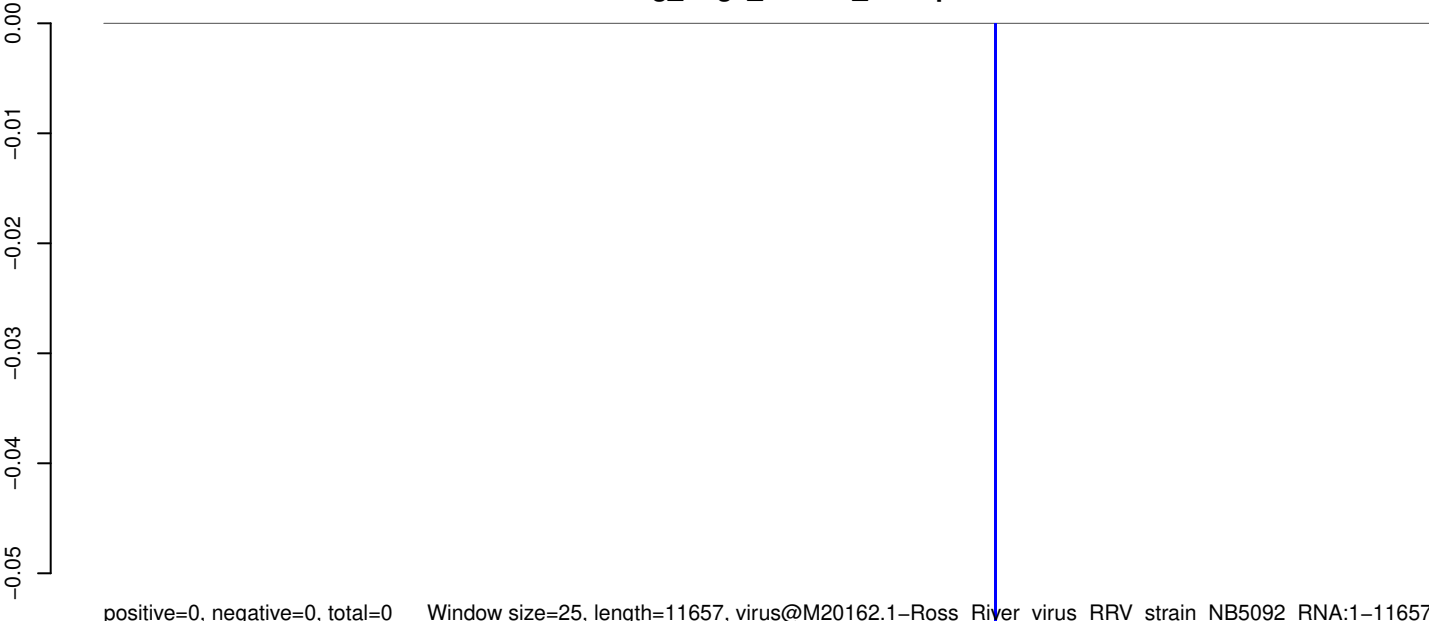
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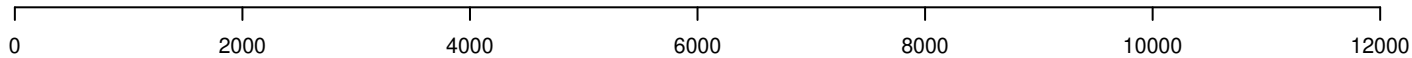
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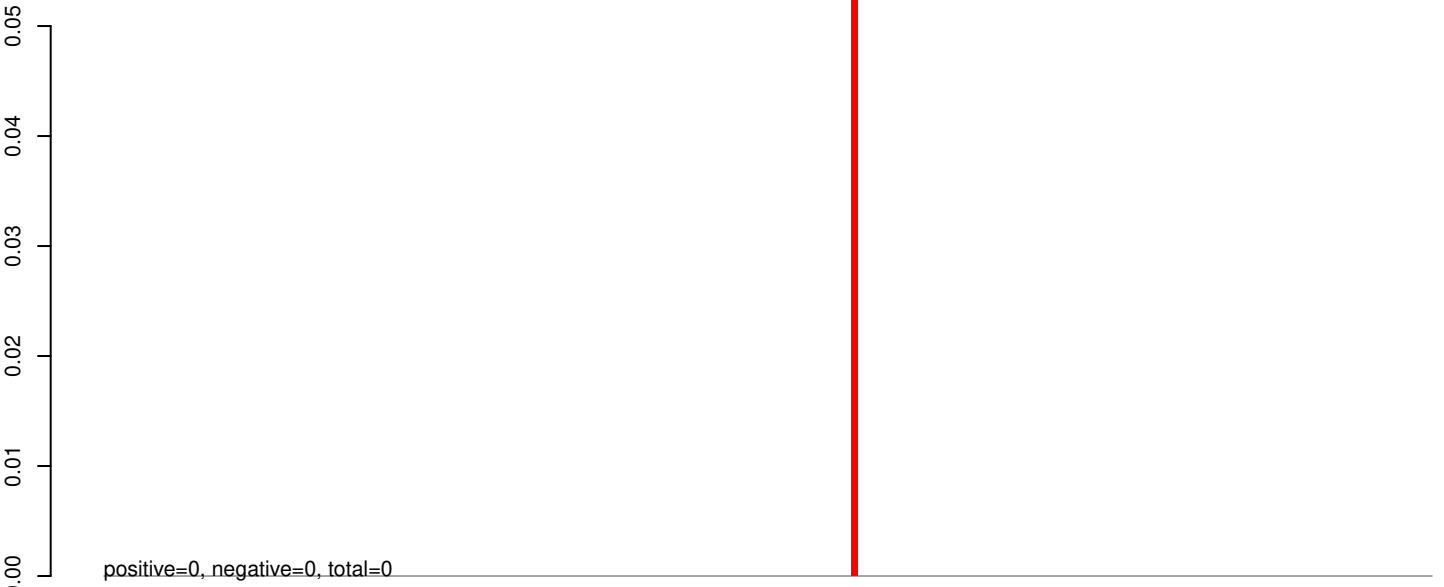
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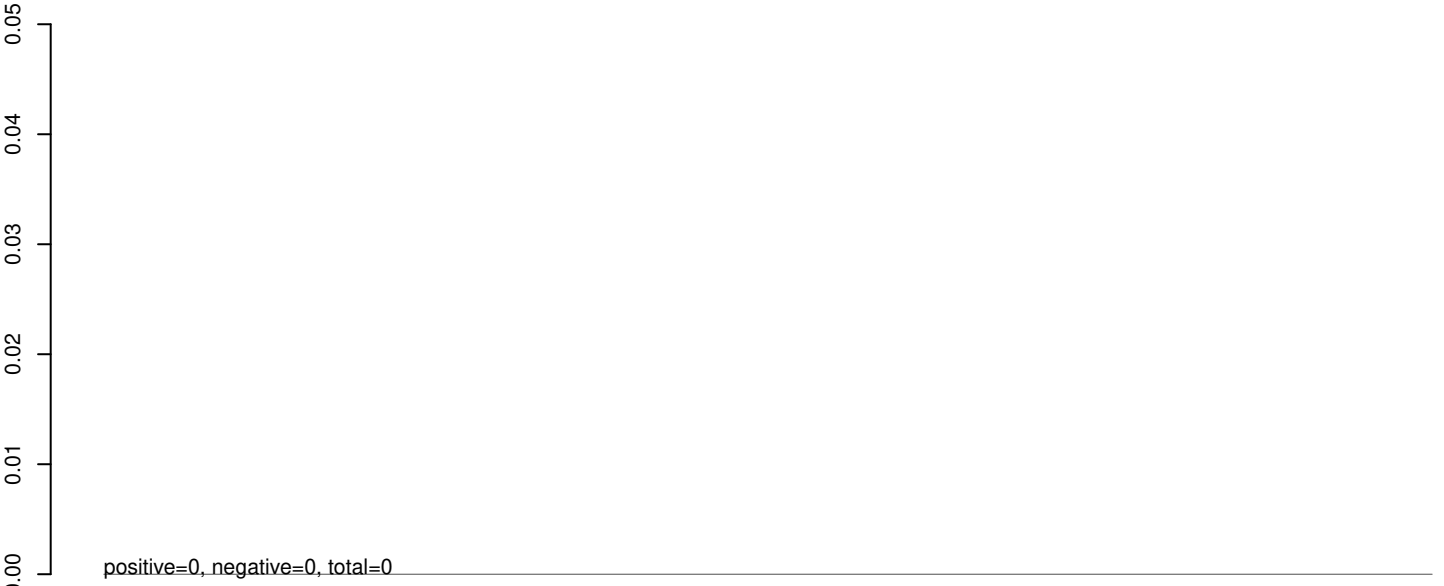
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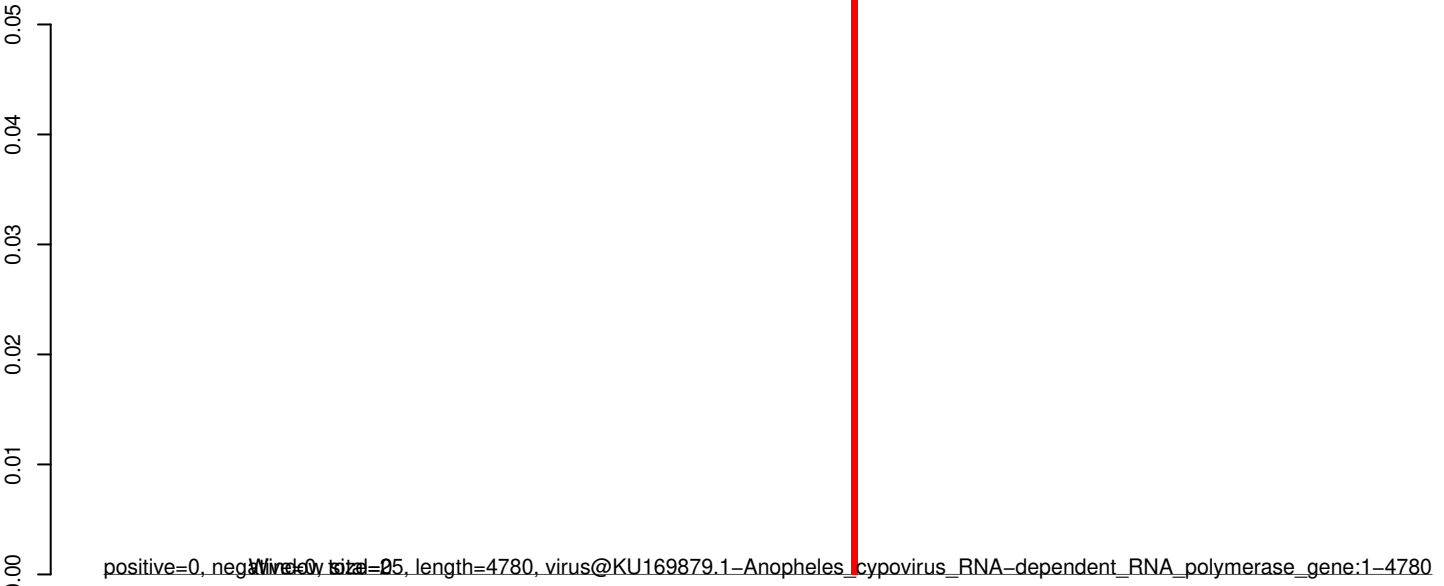
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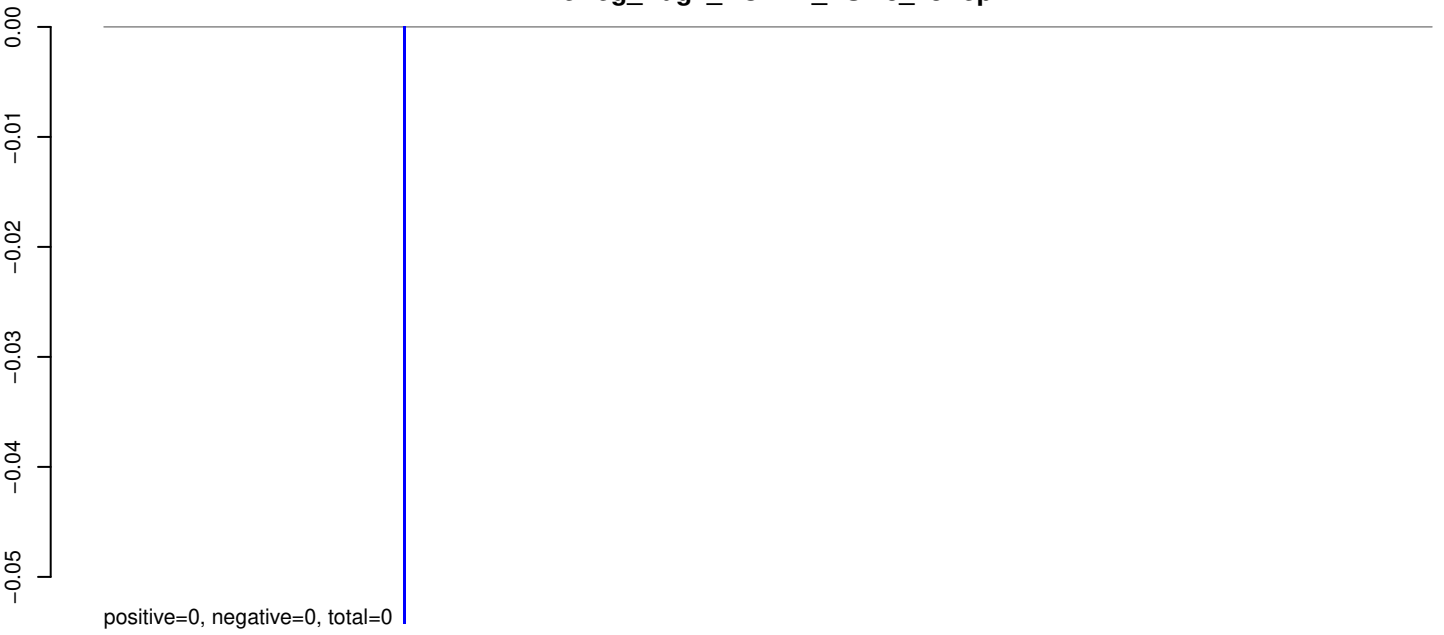
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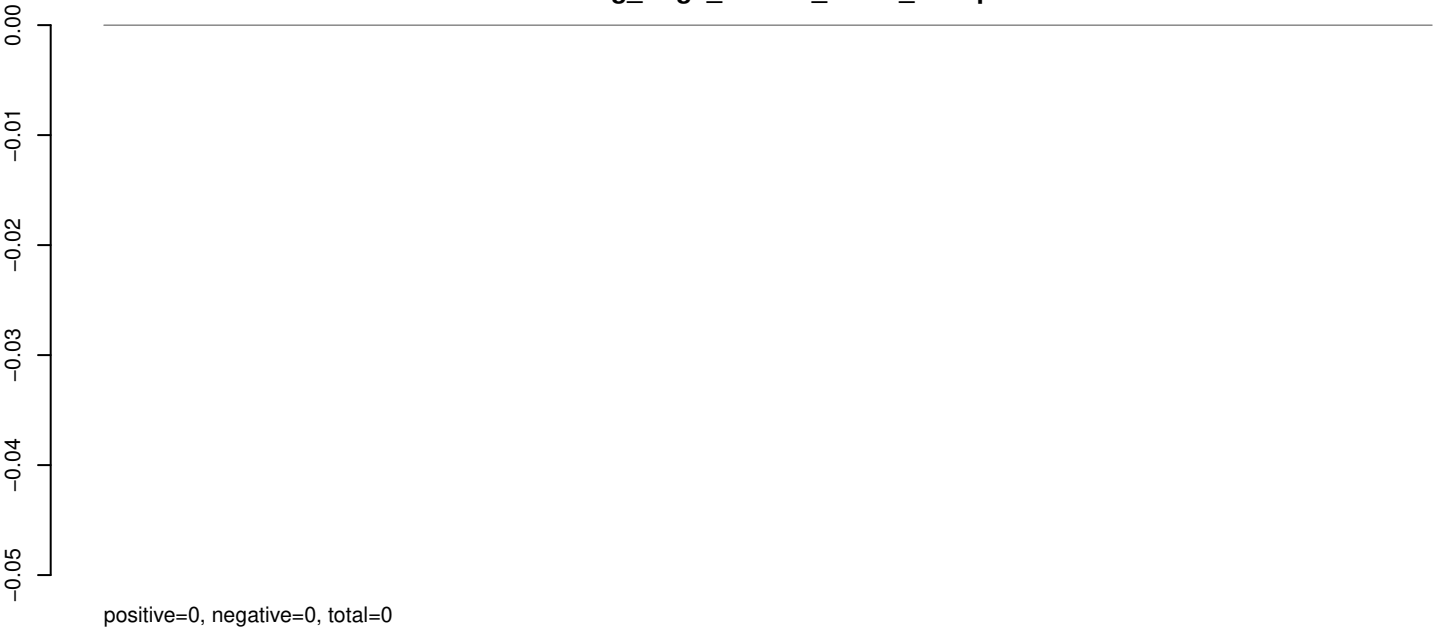
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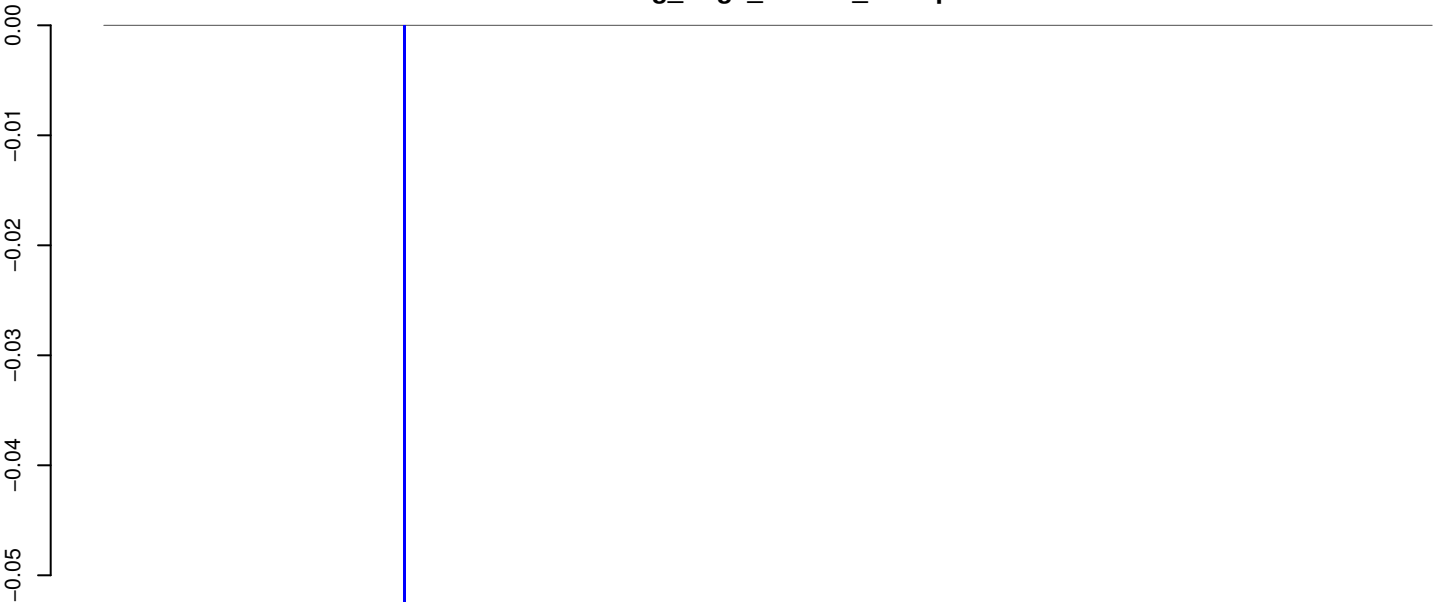
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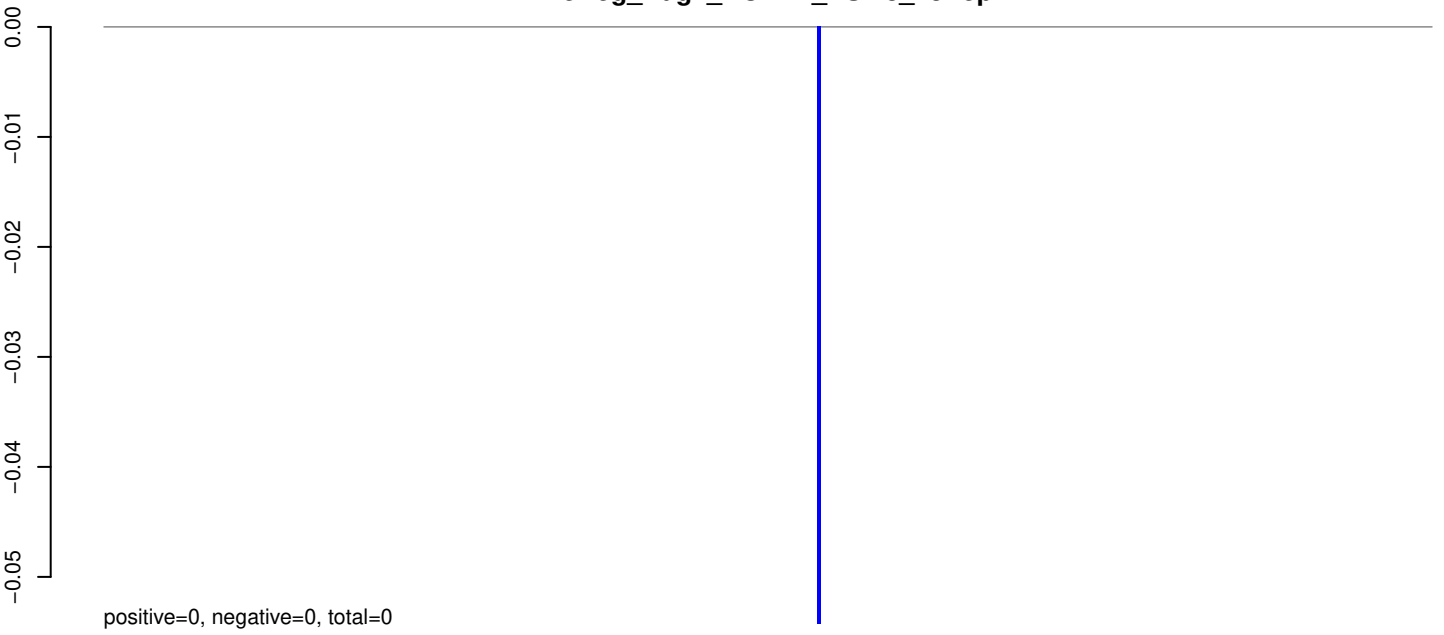
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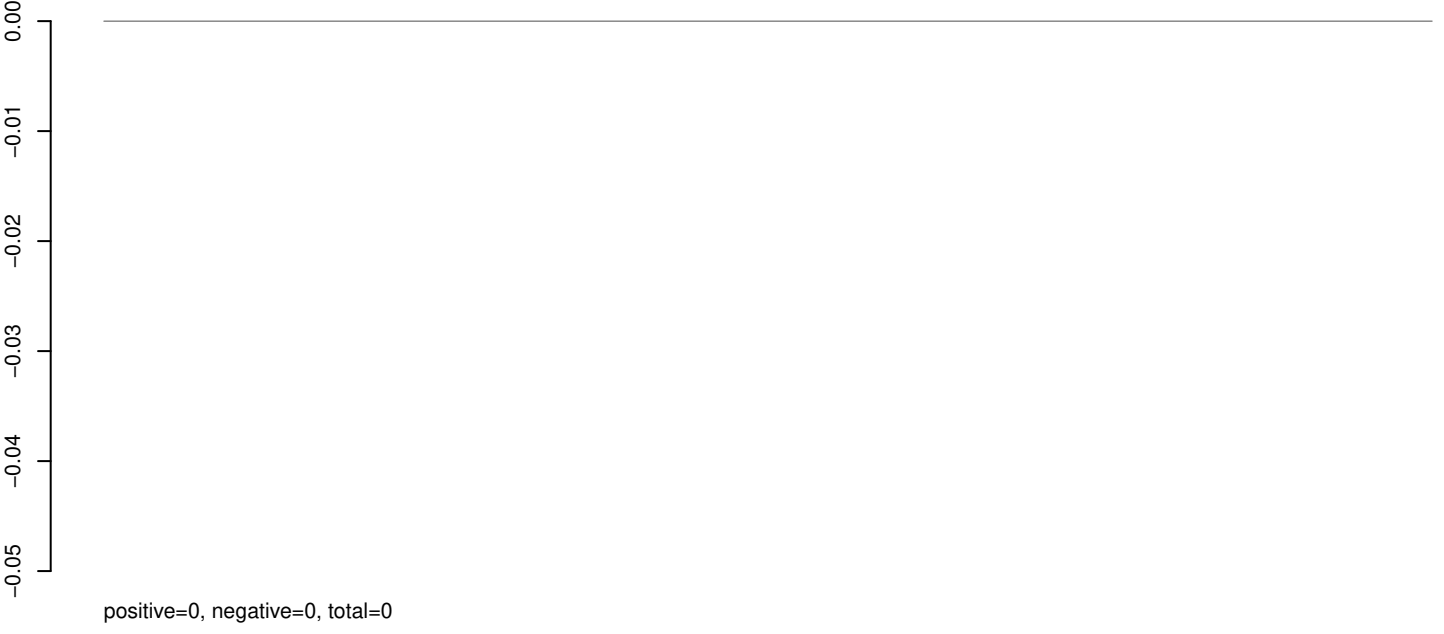
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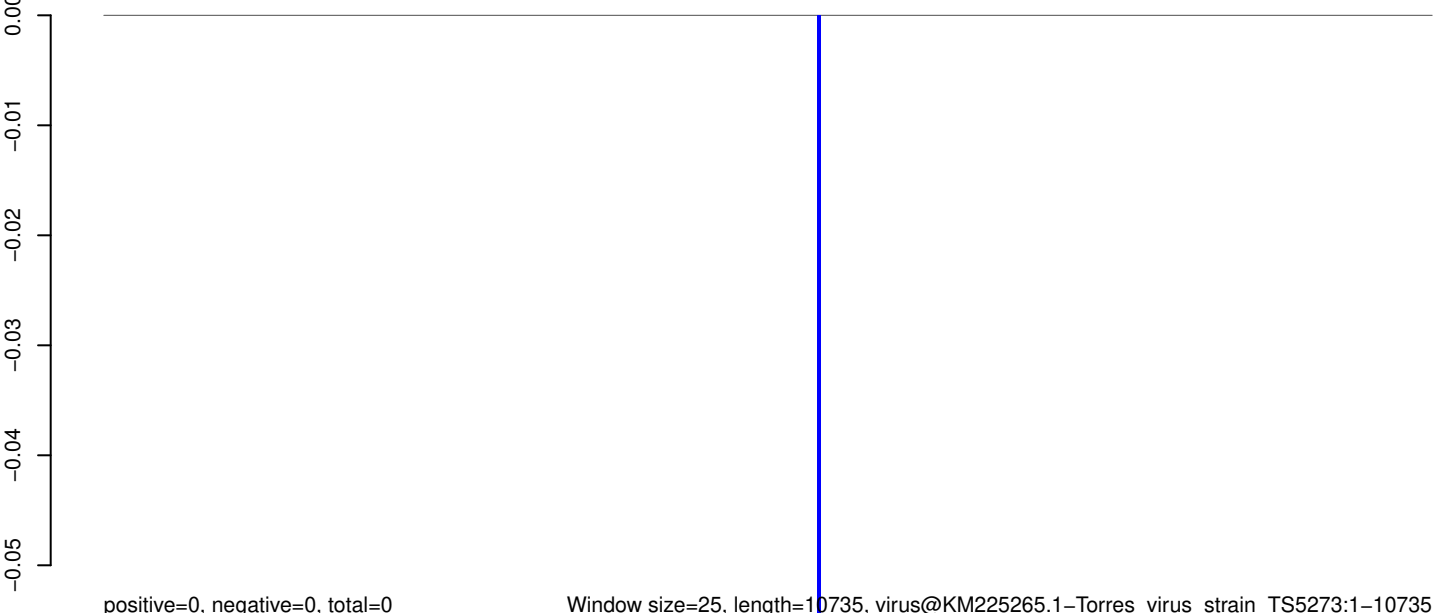
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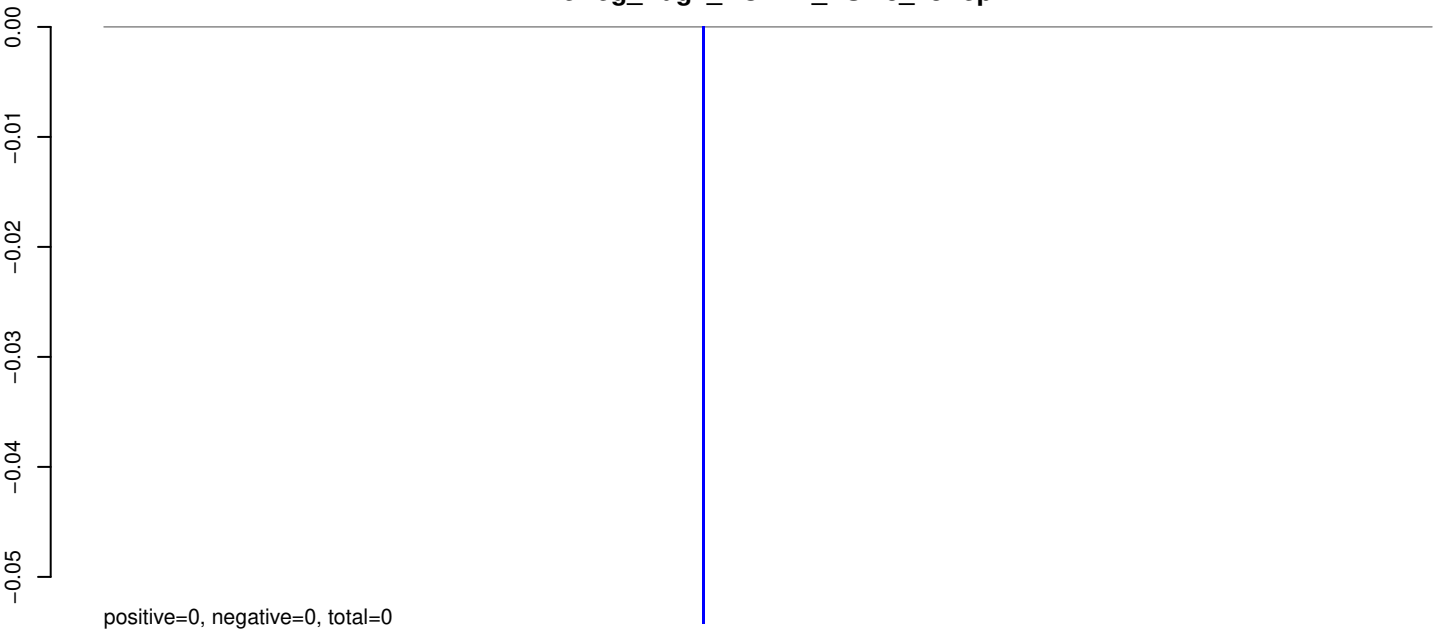
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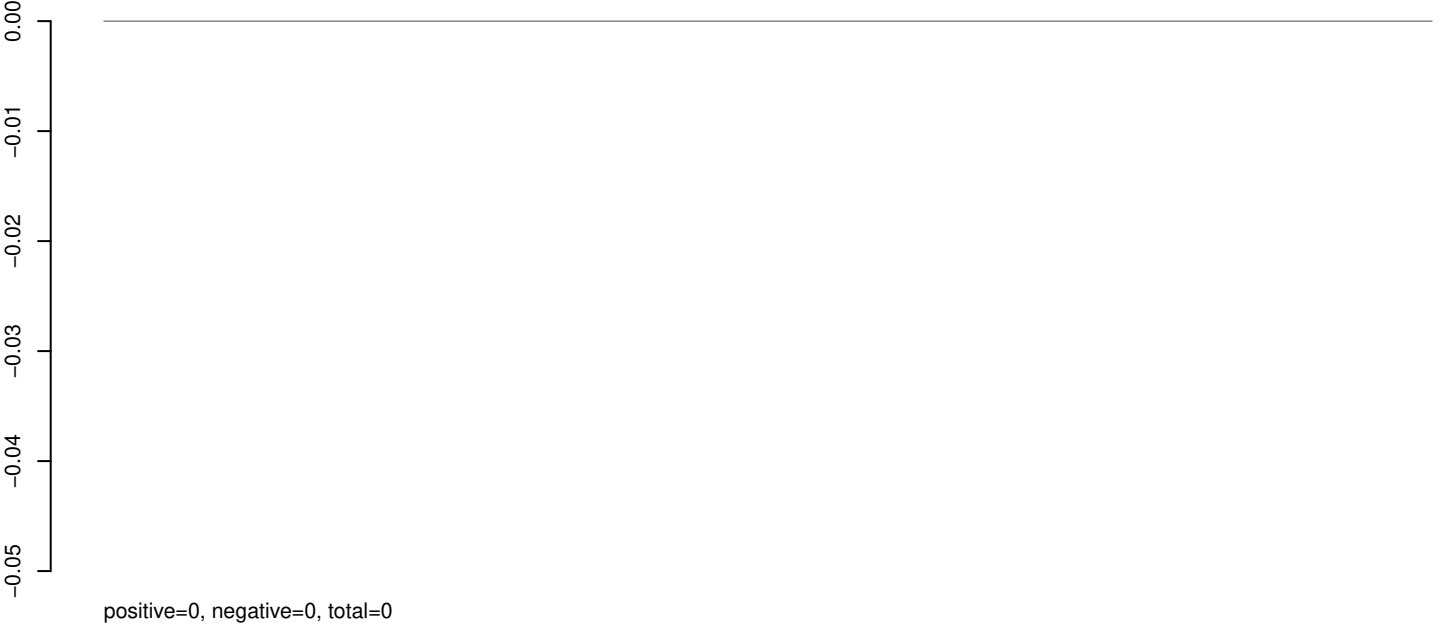
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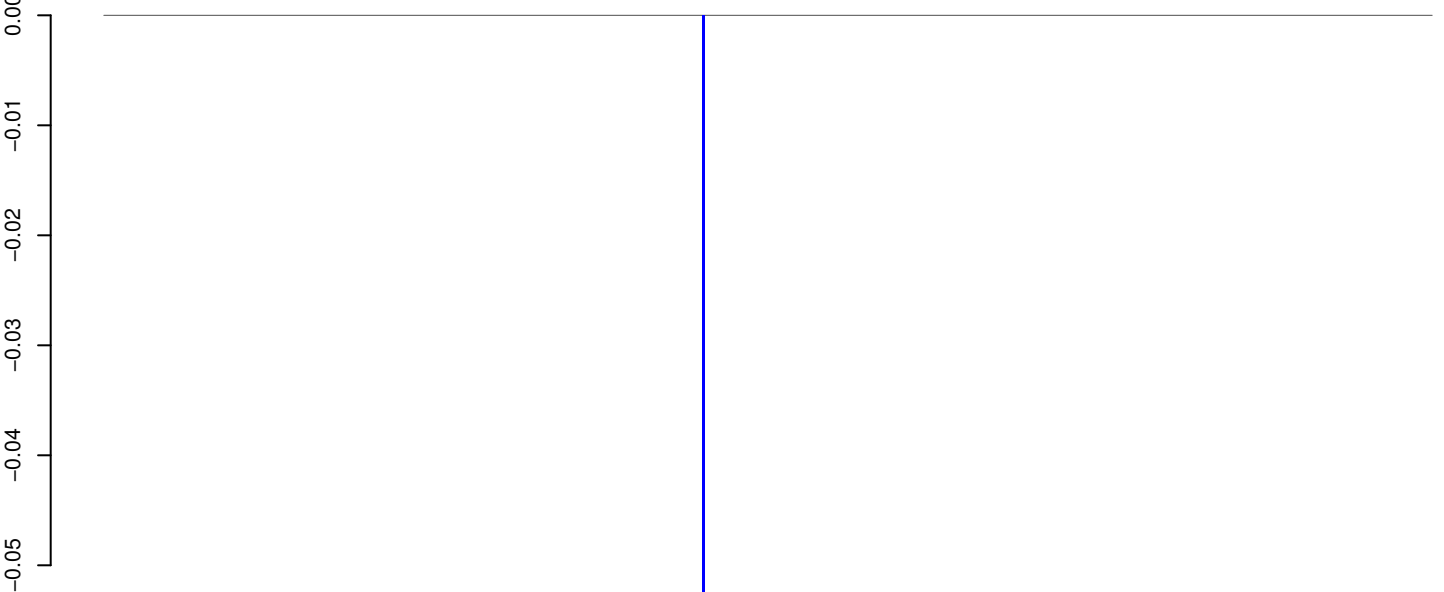
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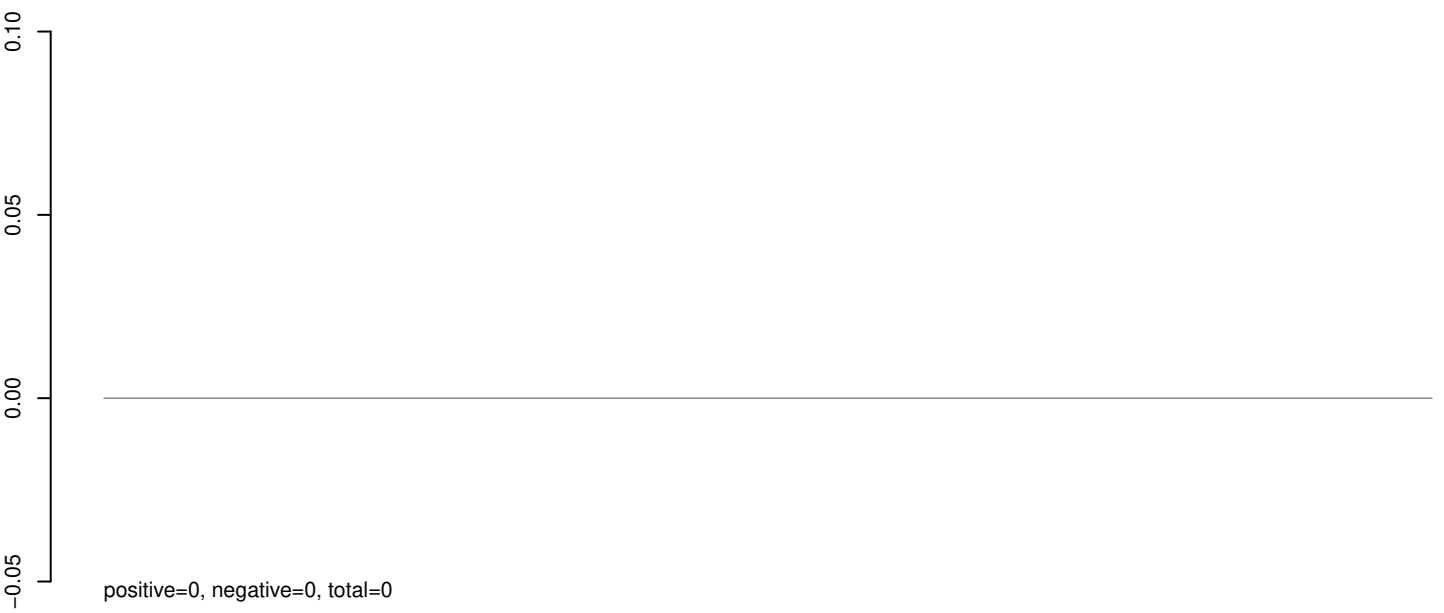
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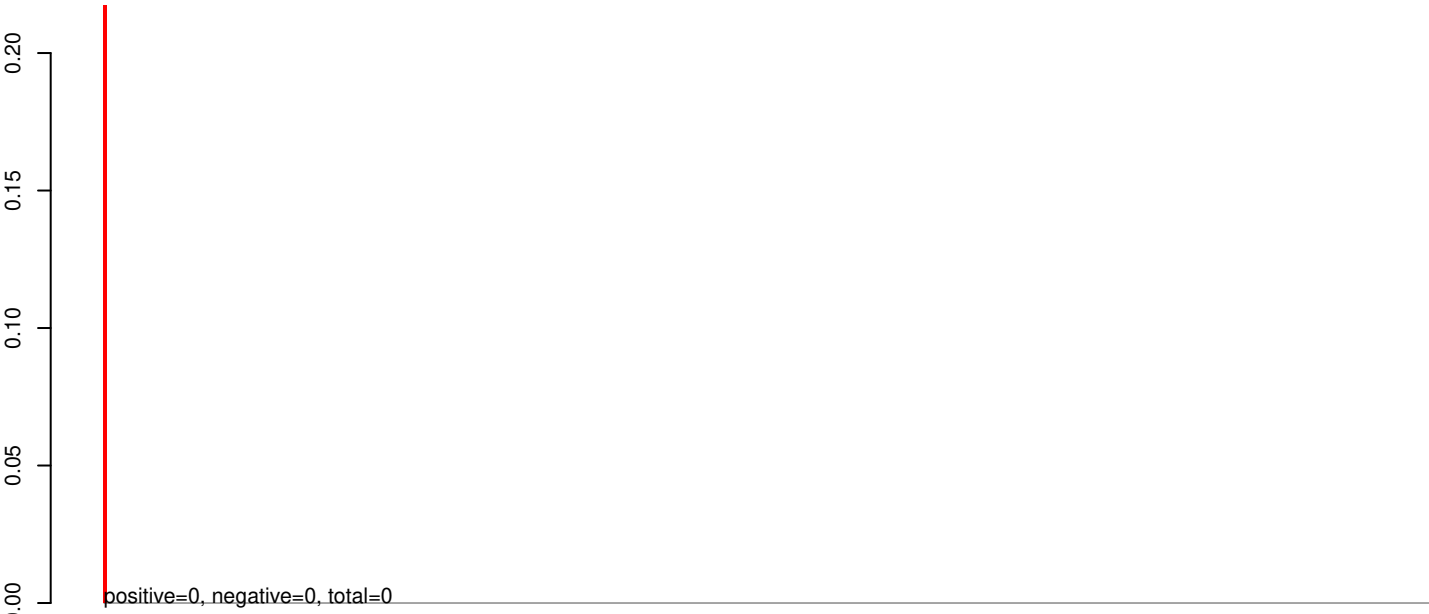
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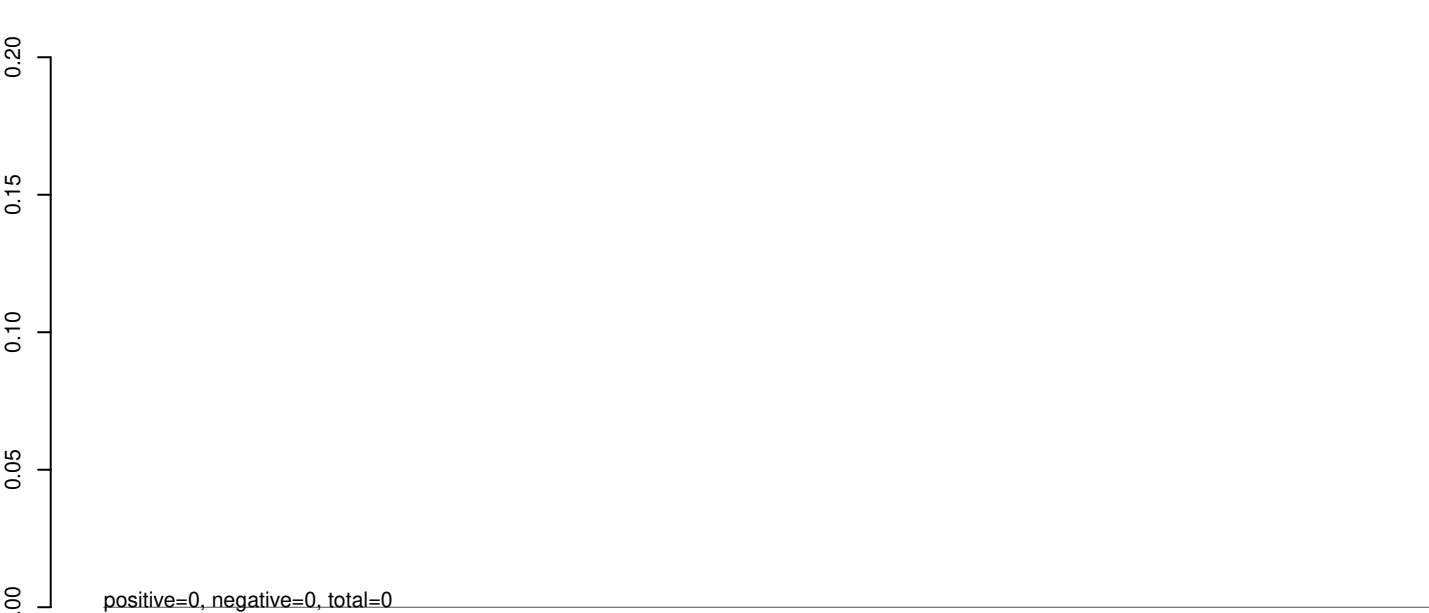
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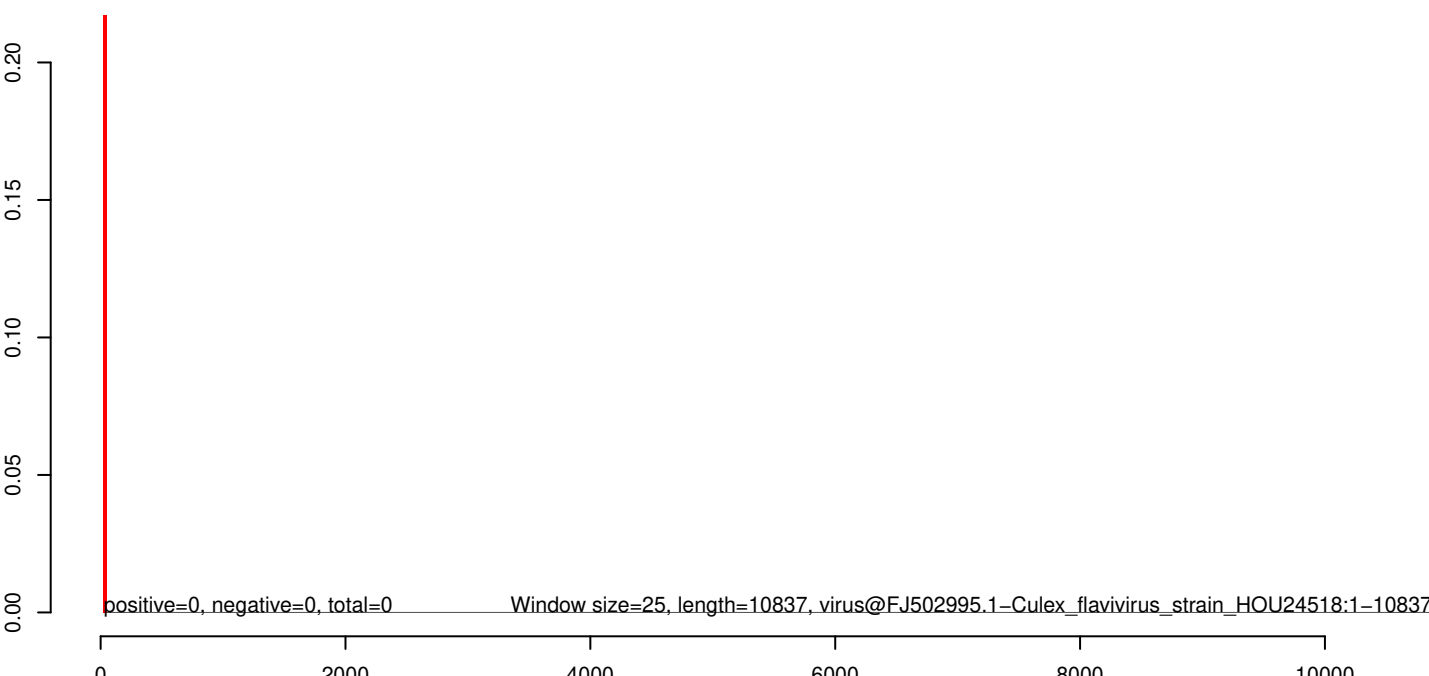
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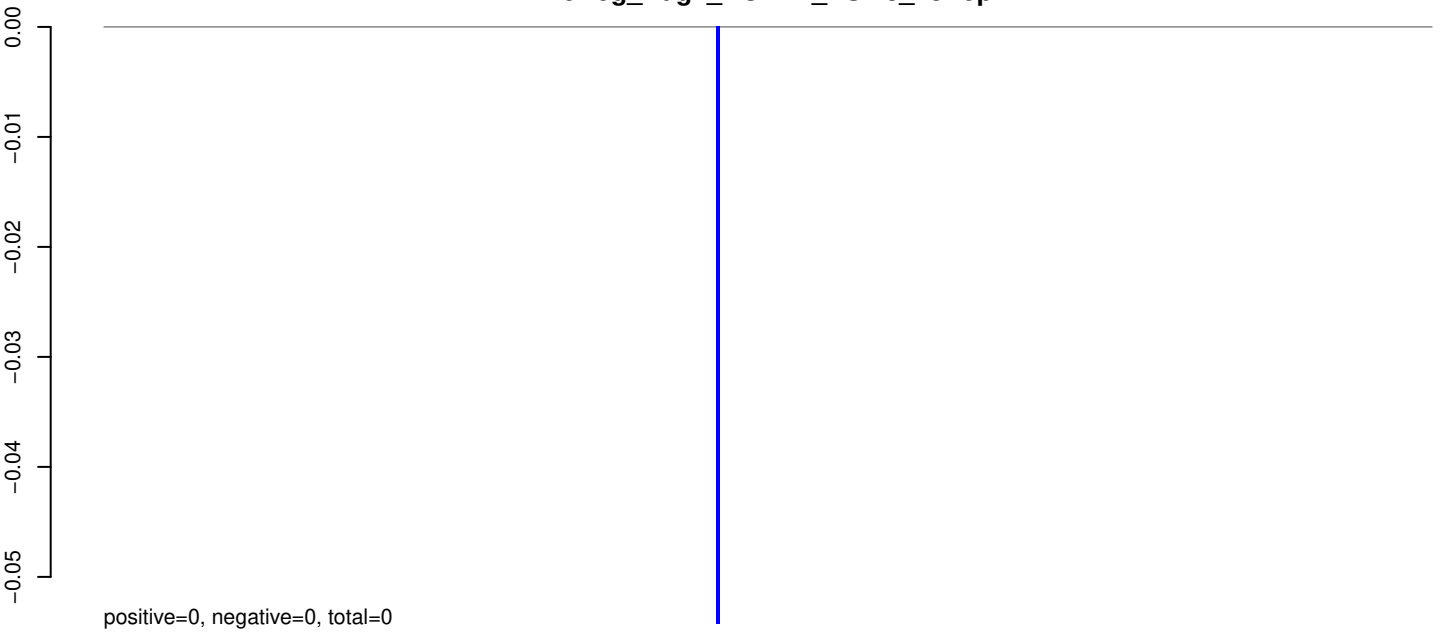
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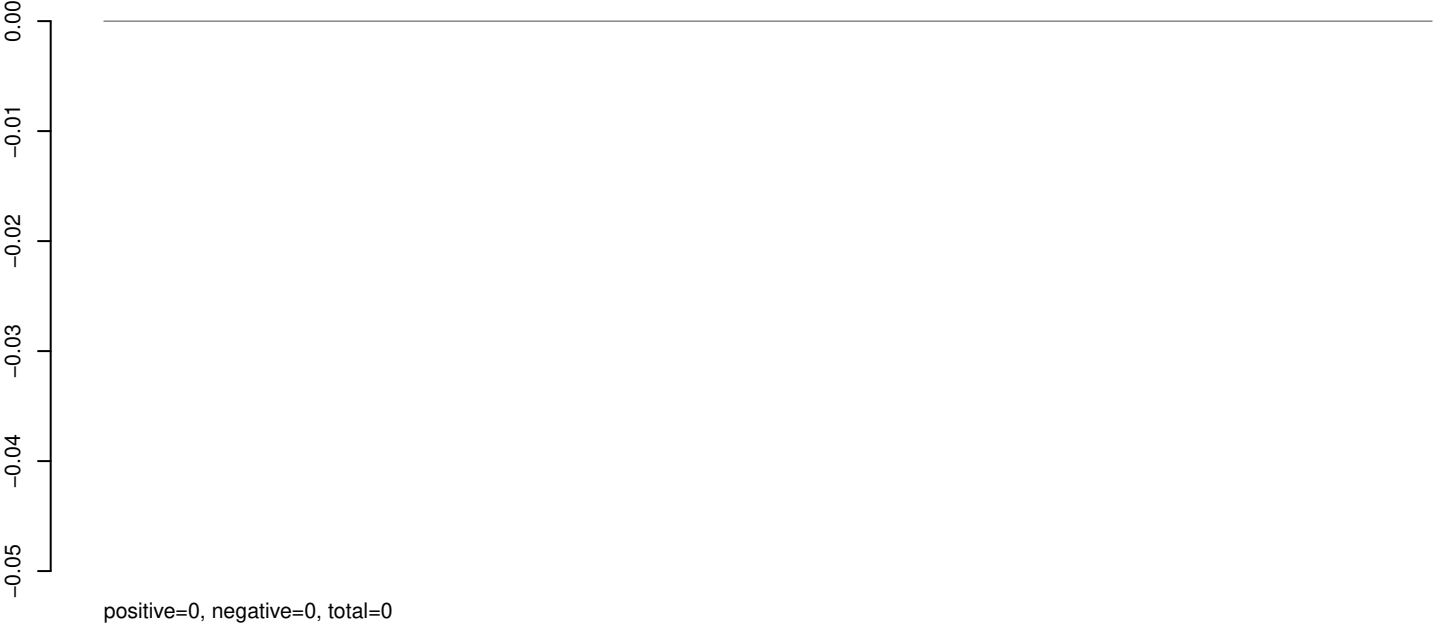
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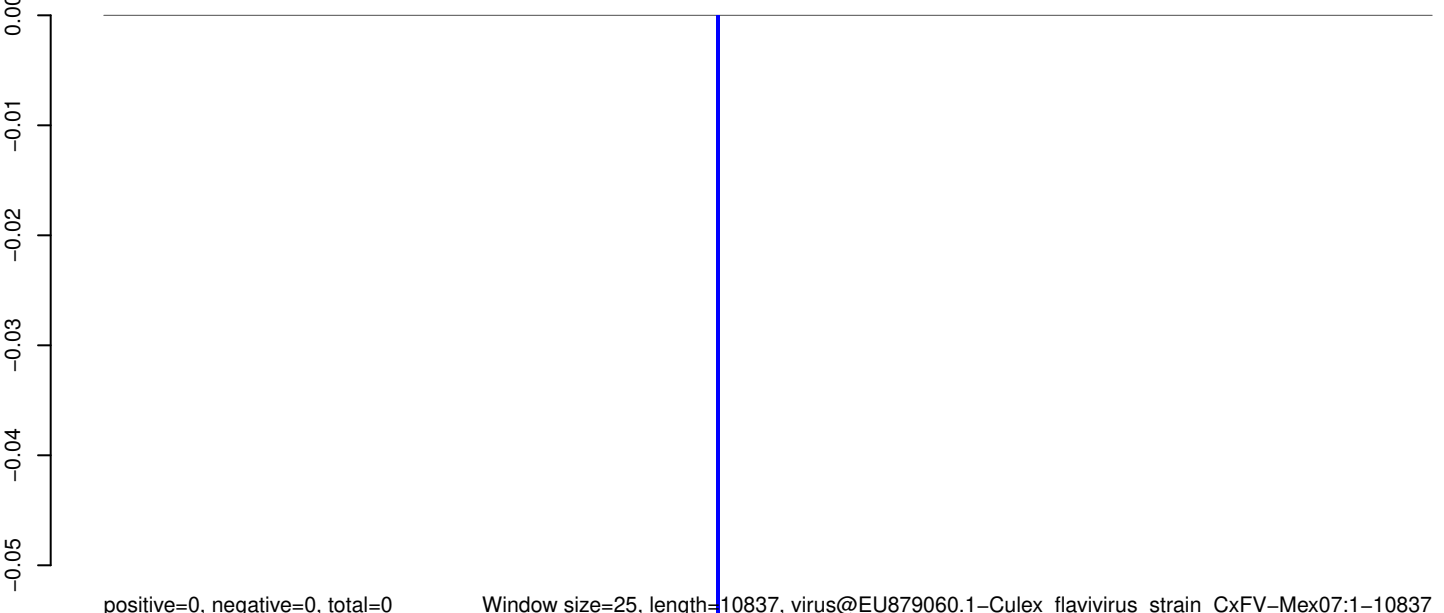
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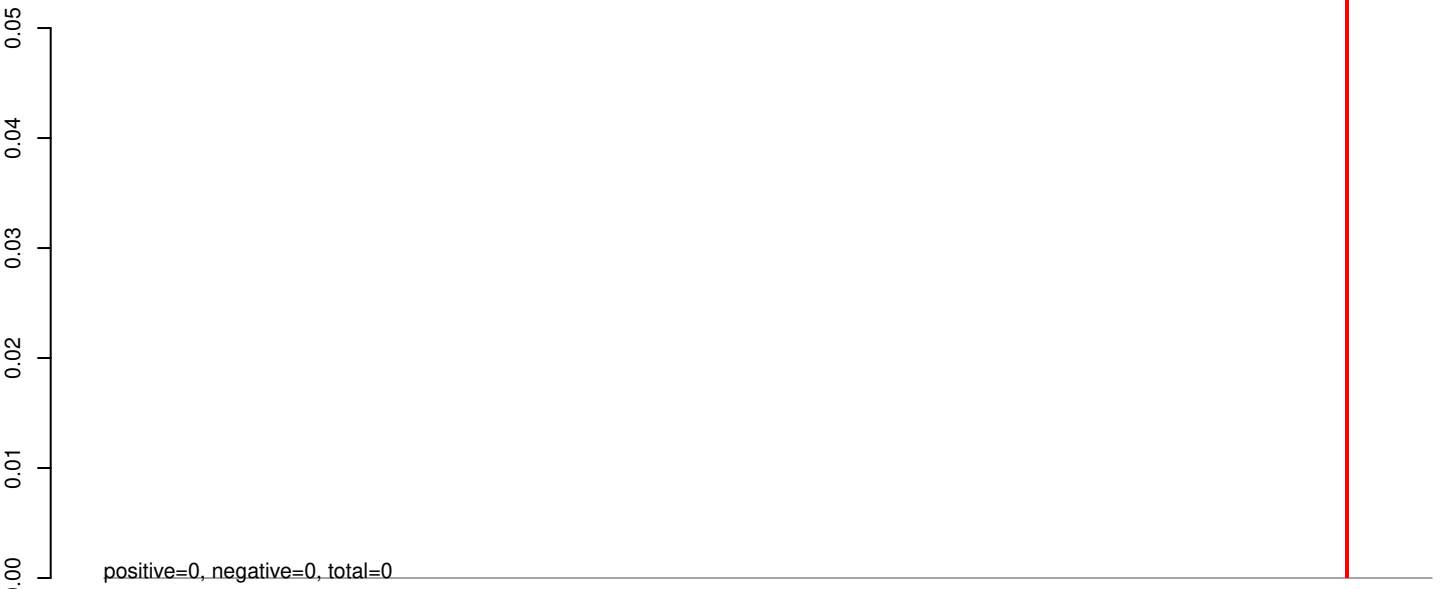


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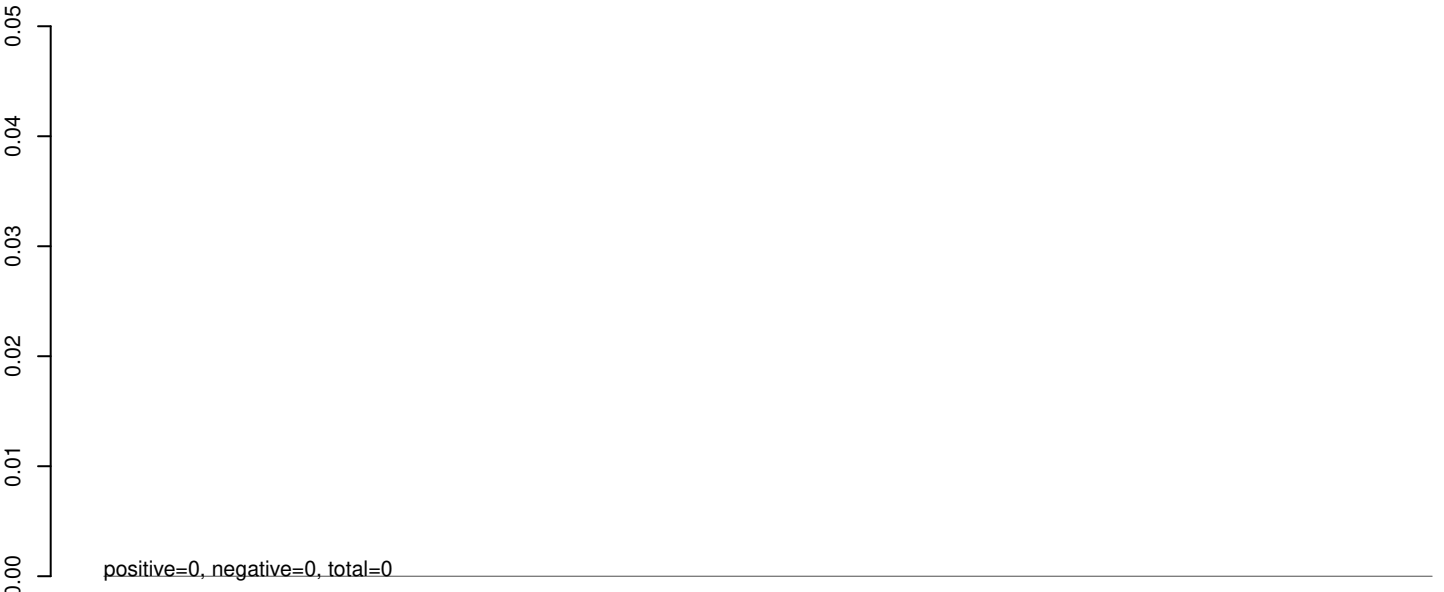


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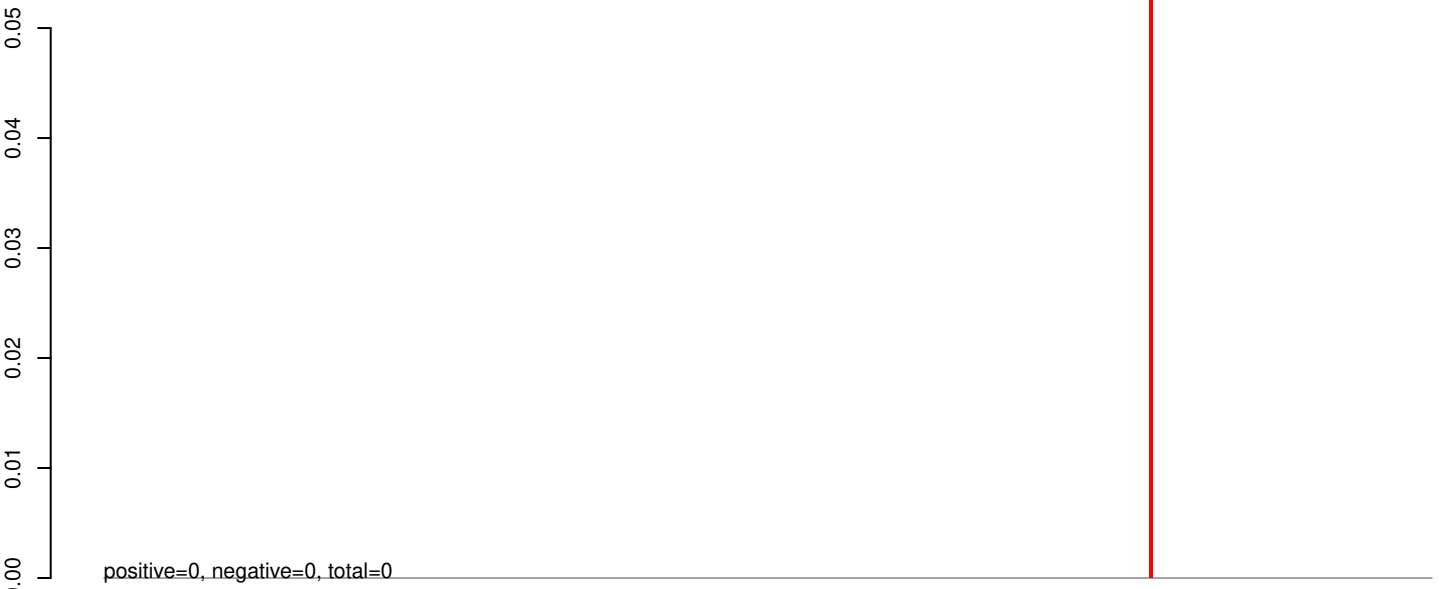
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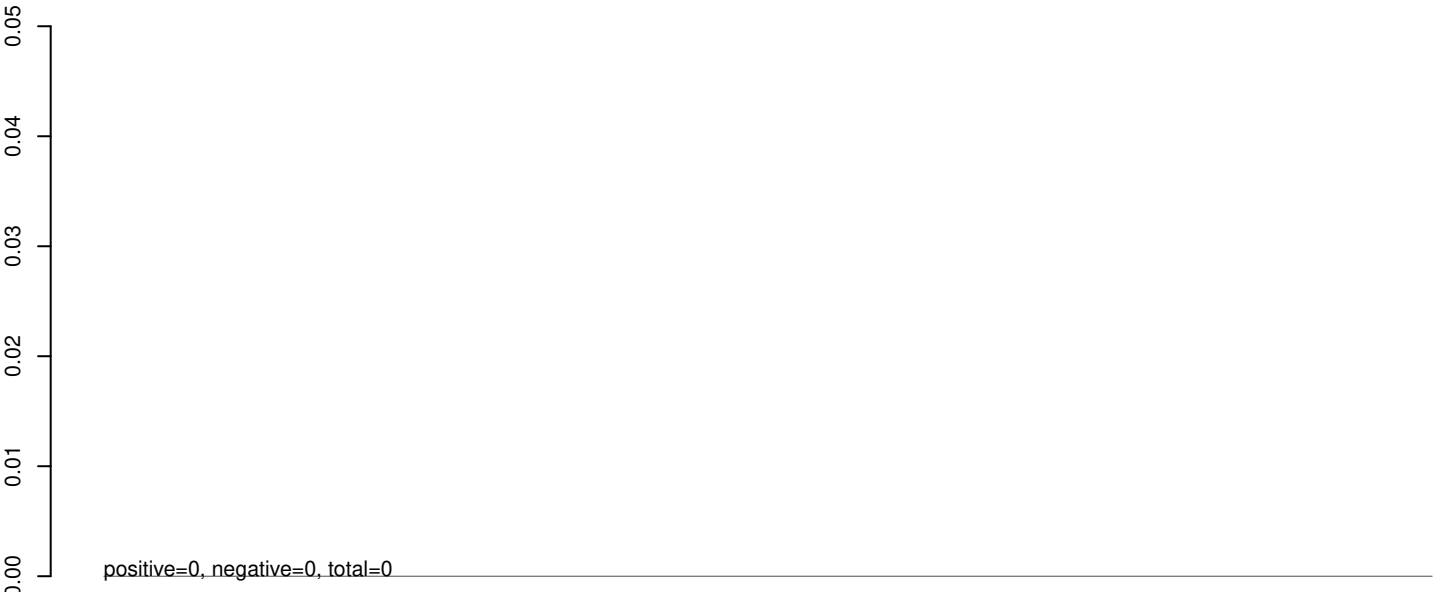
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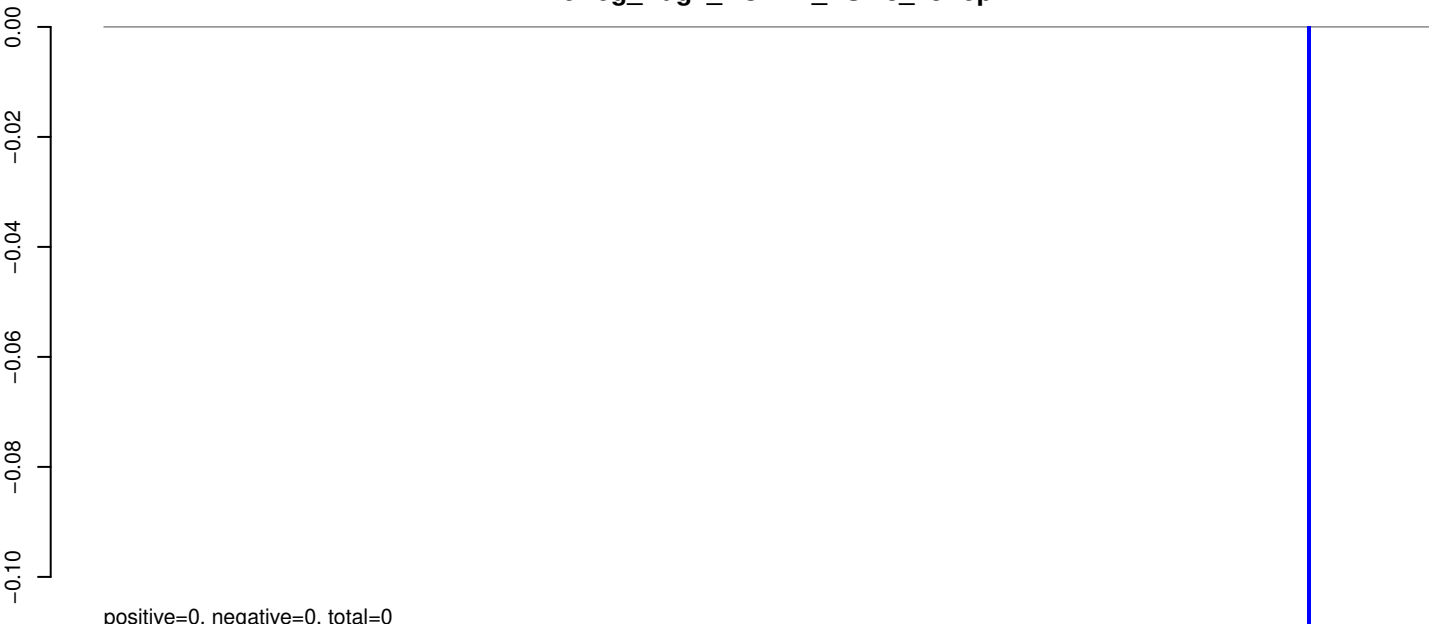


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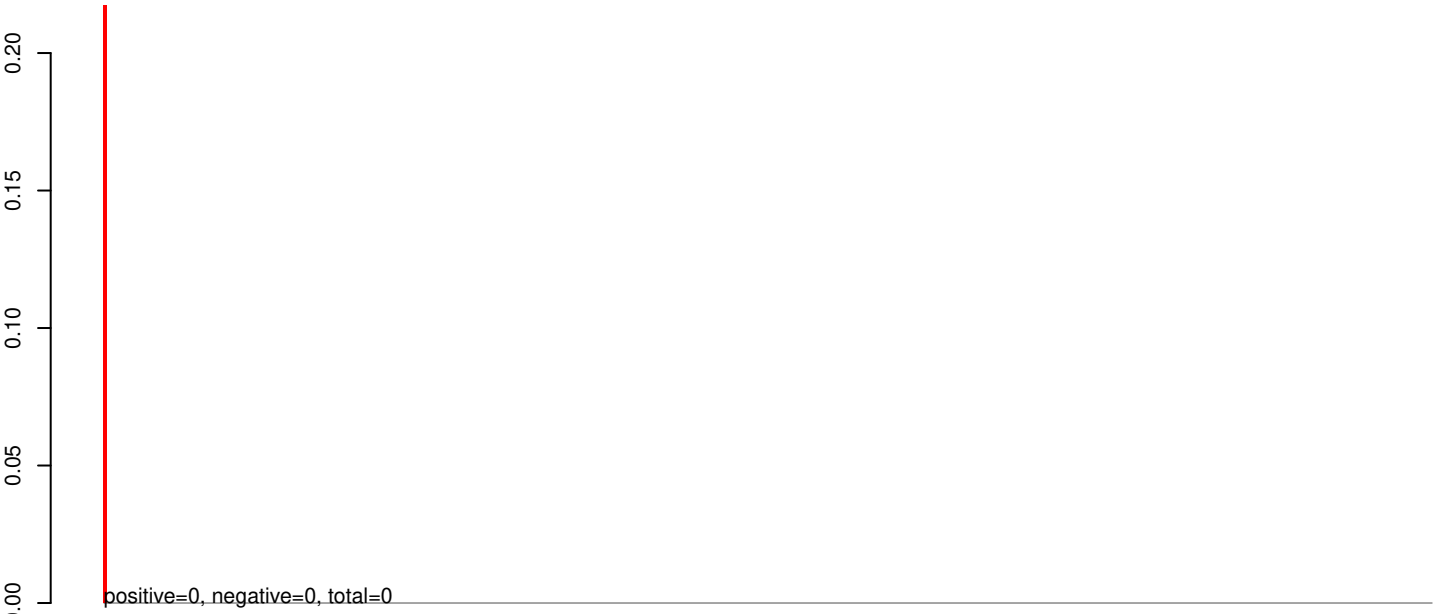
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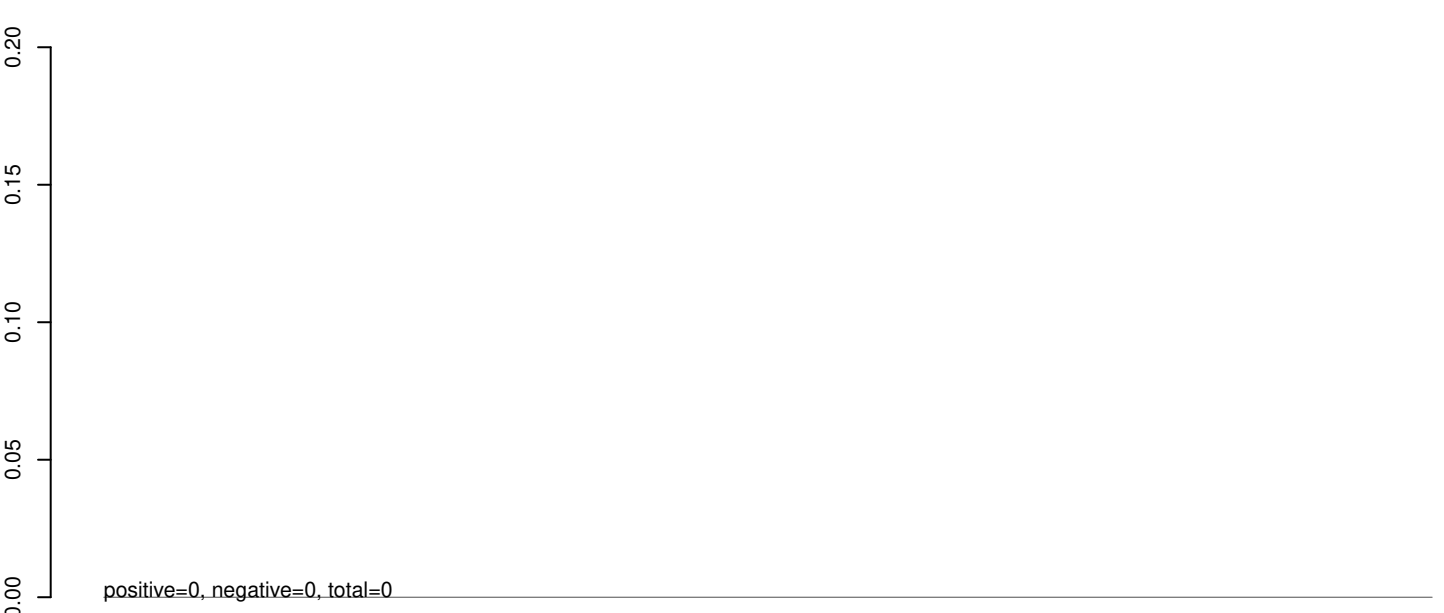
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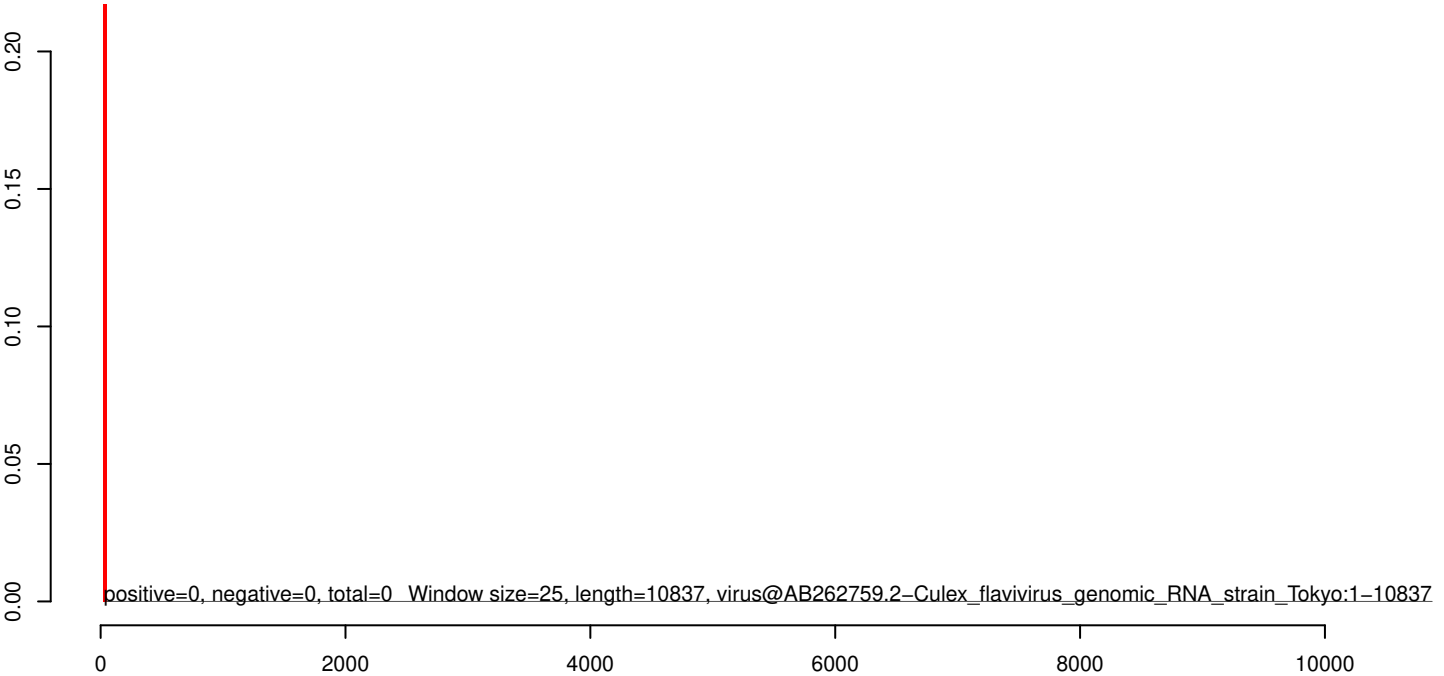
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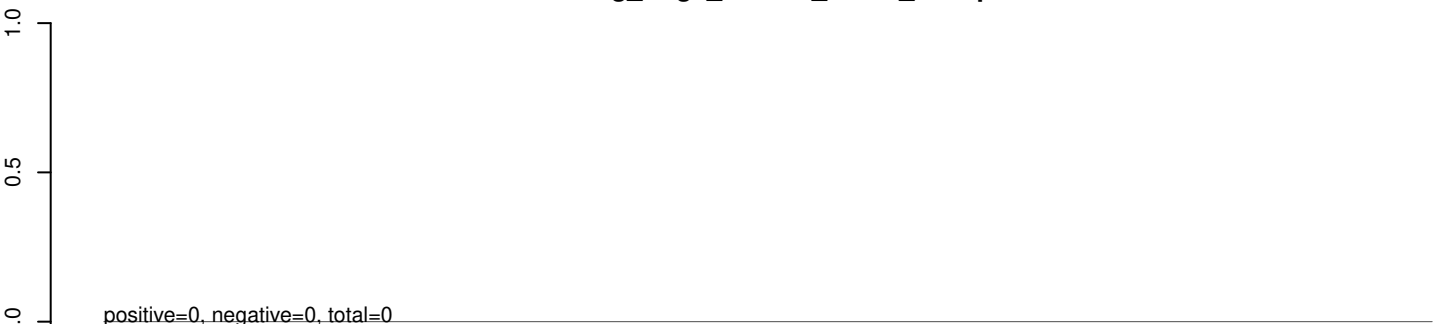
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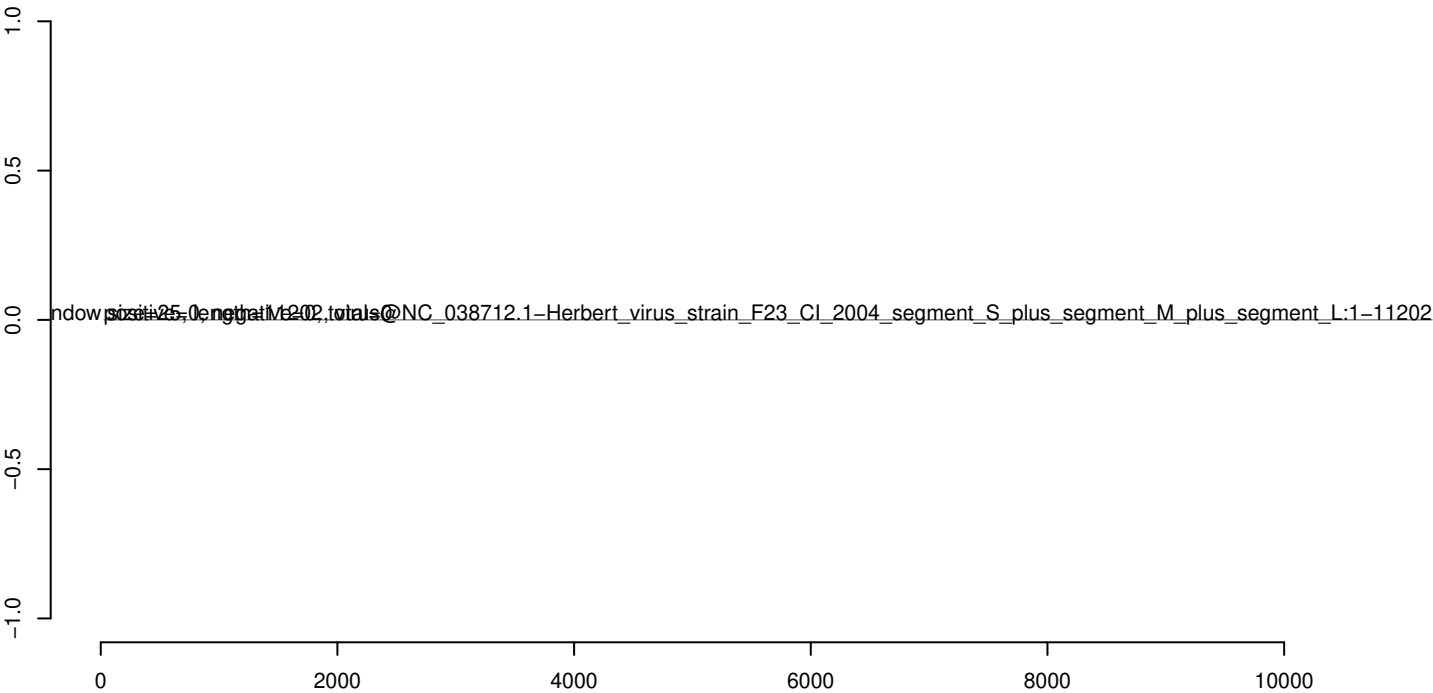
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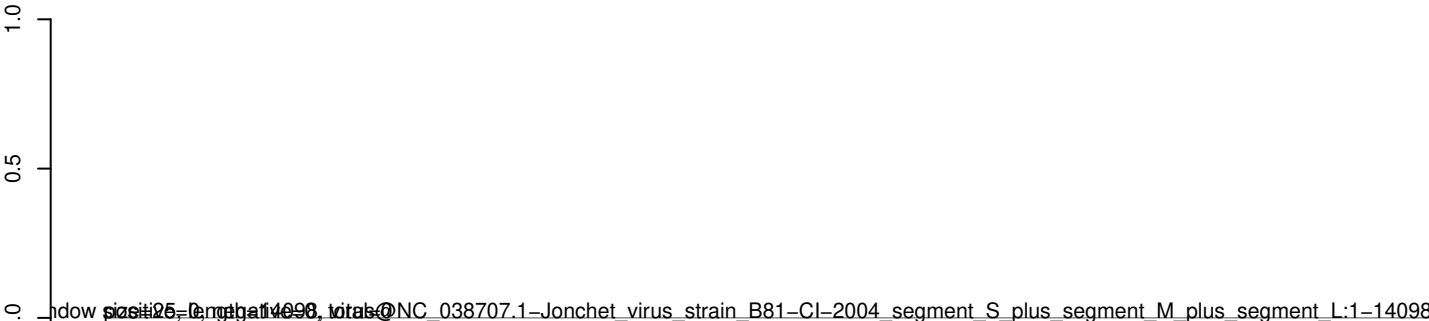
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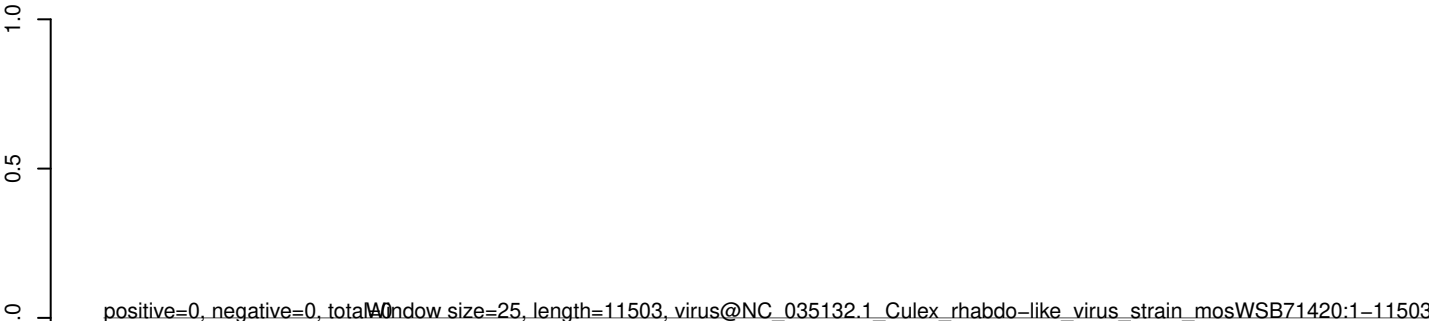
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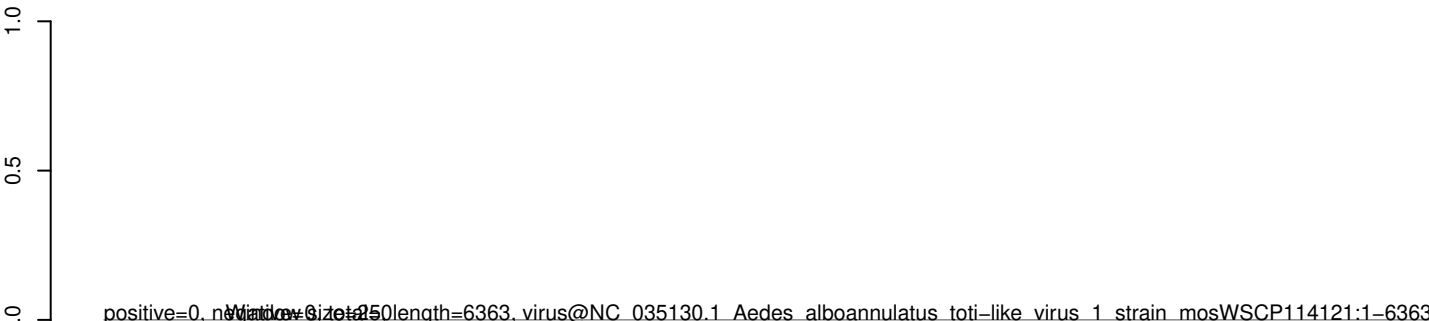
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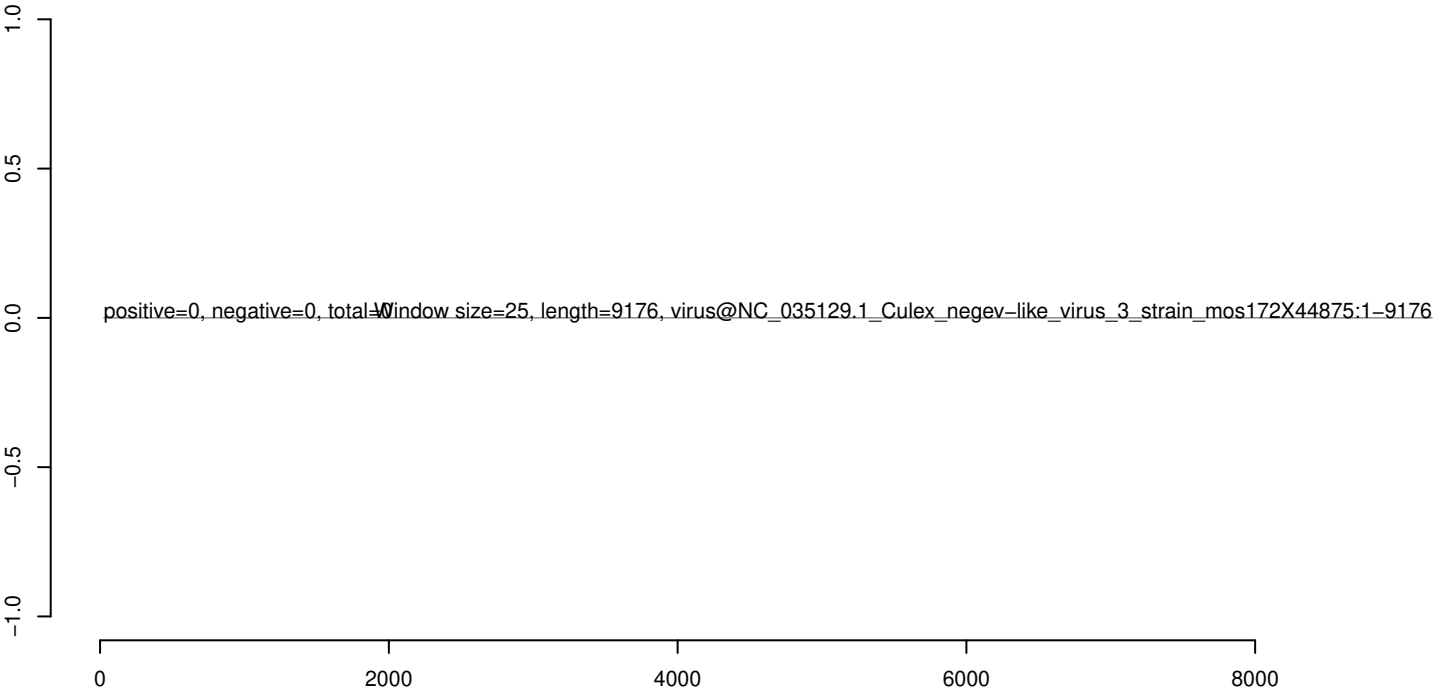
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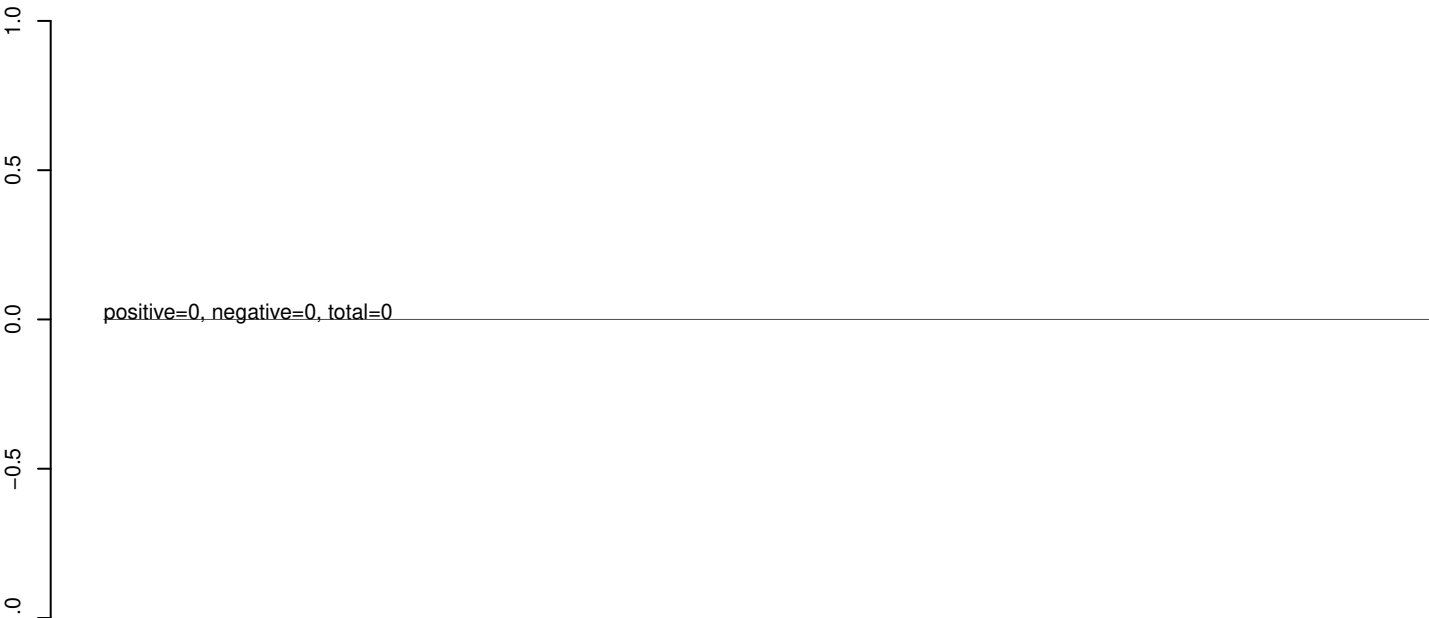
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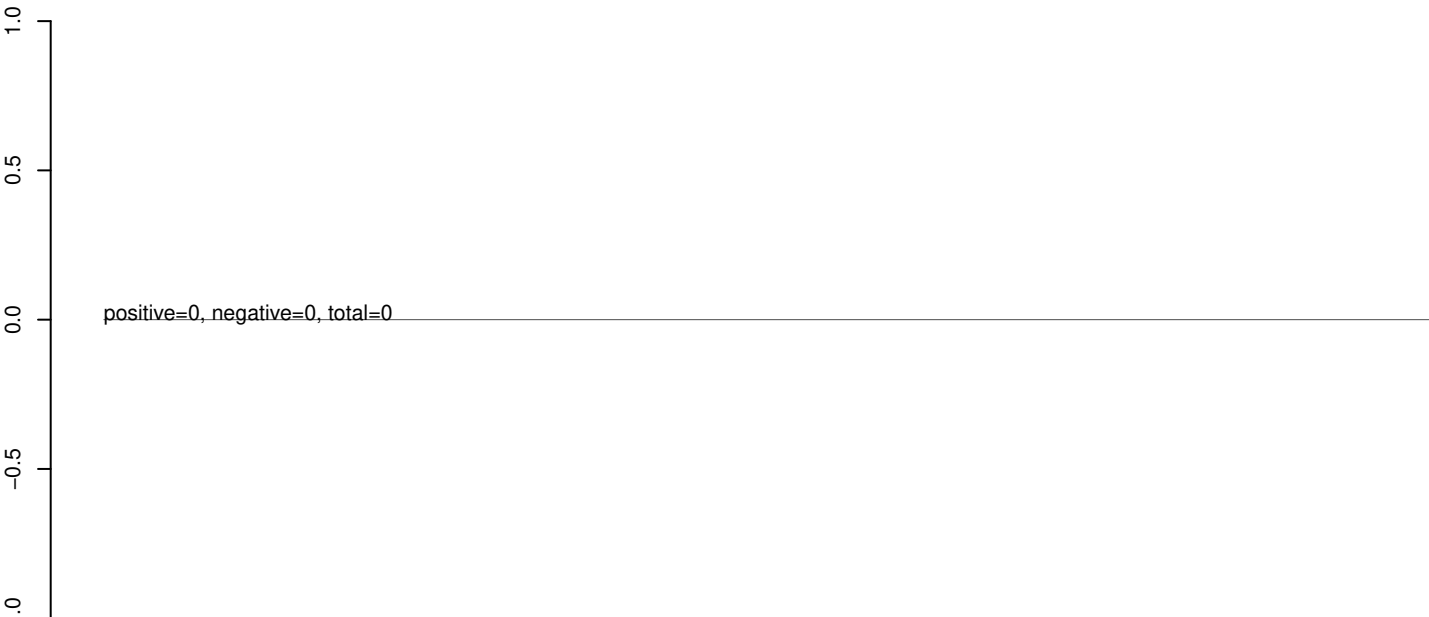
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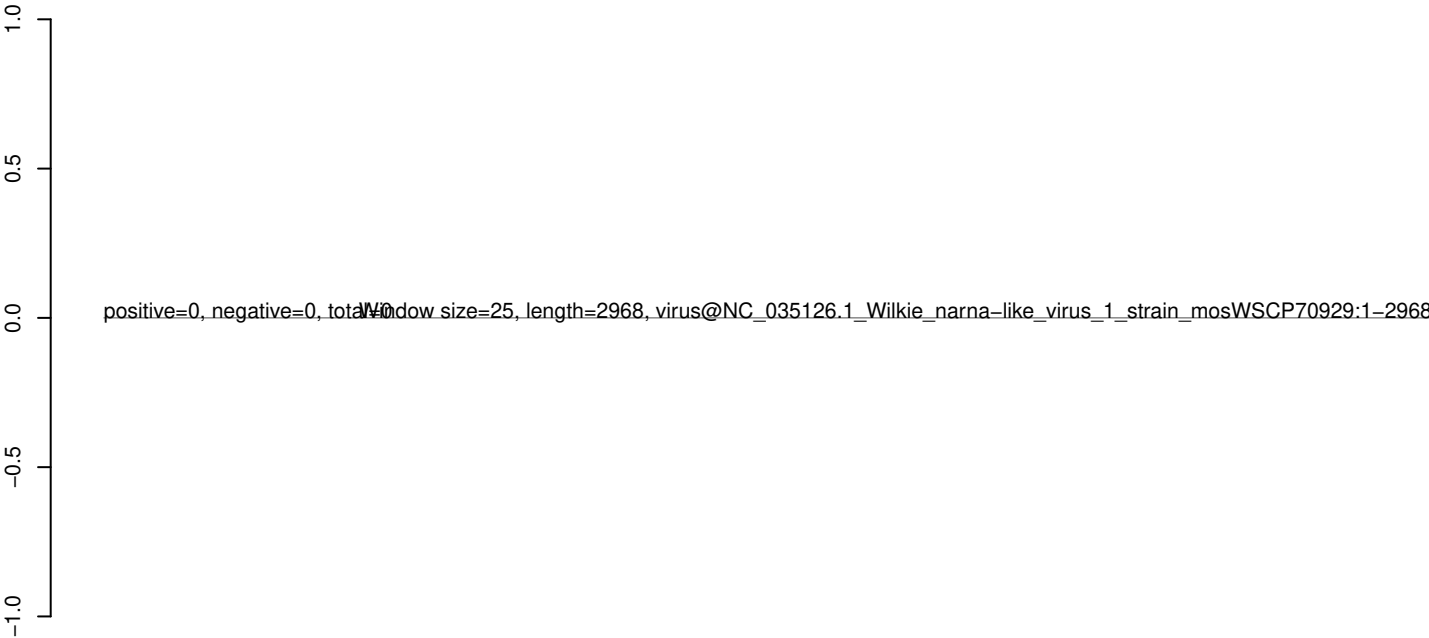
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AeAeg_Aag2_BUNV1_ES.rep



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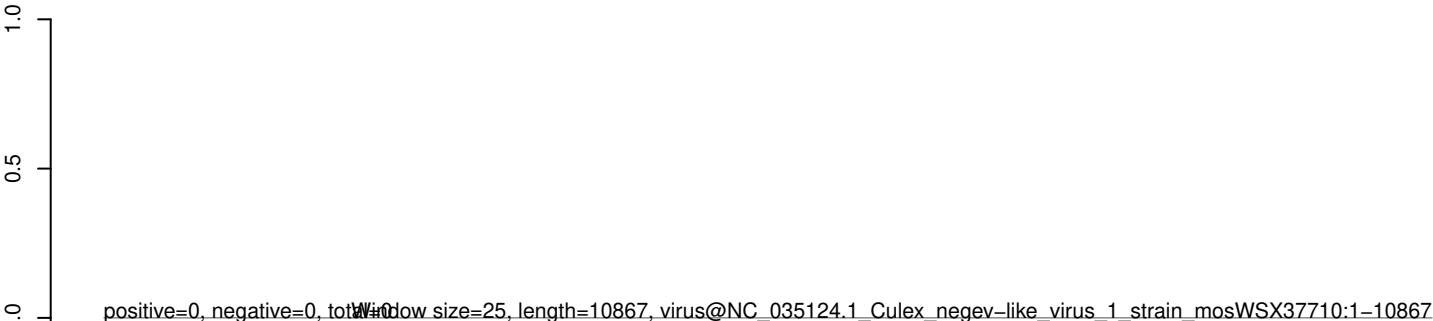
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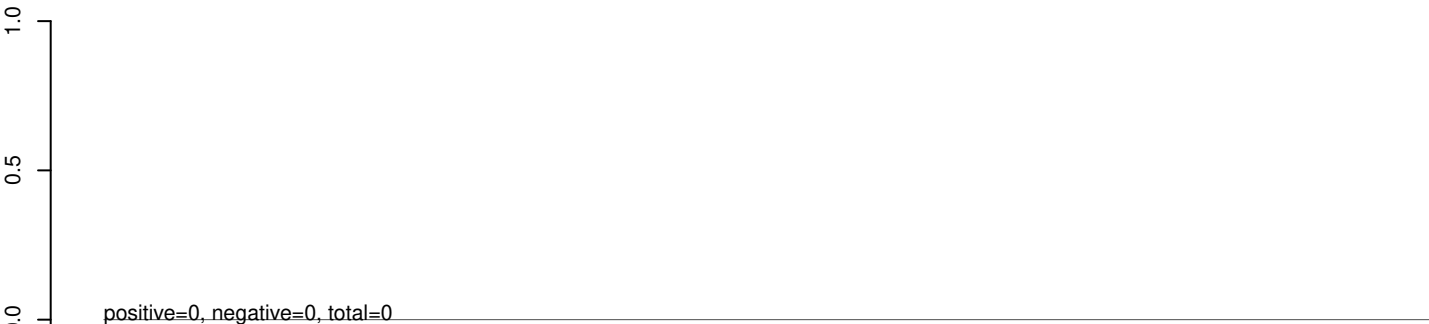


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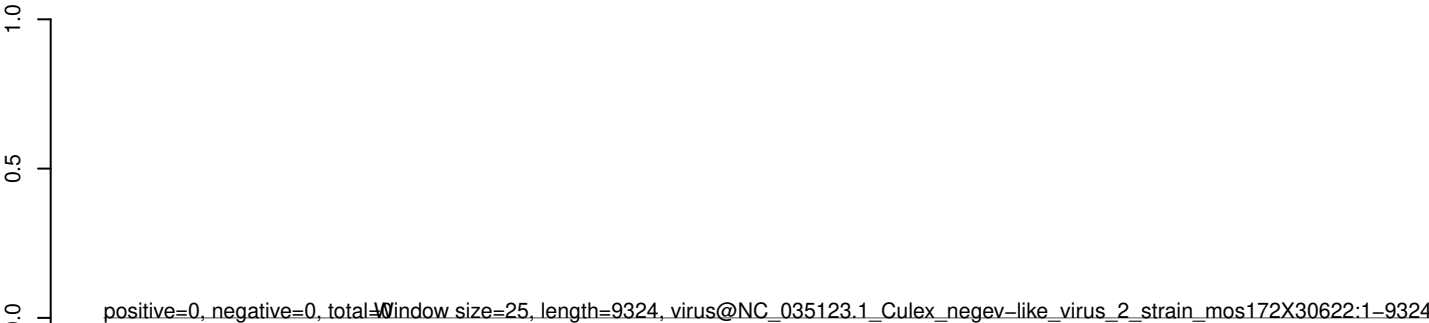
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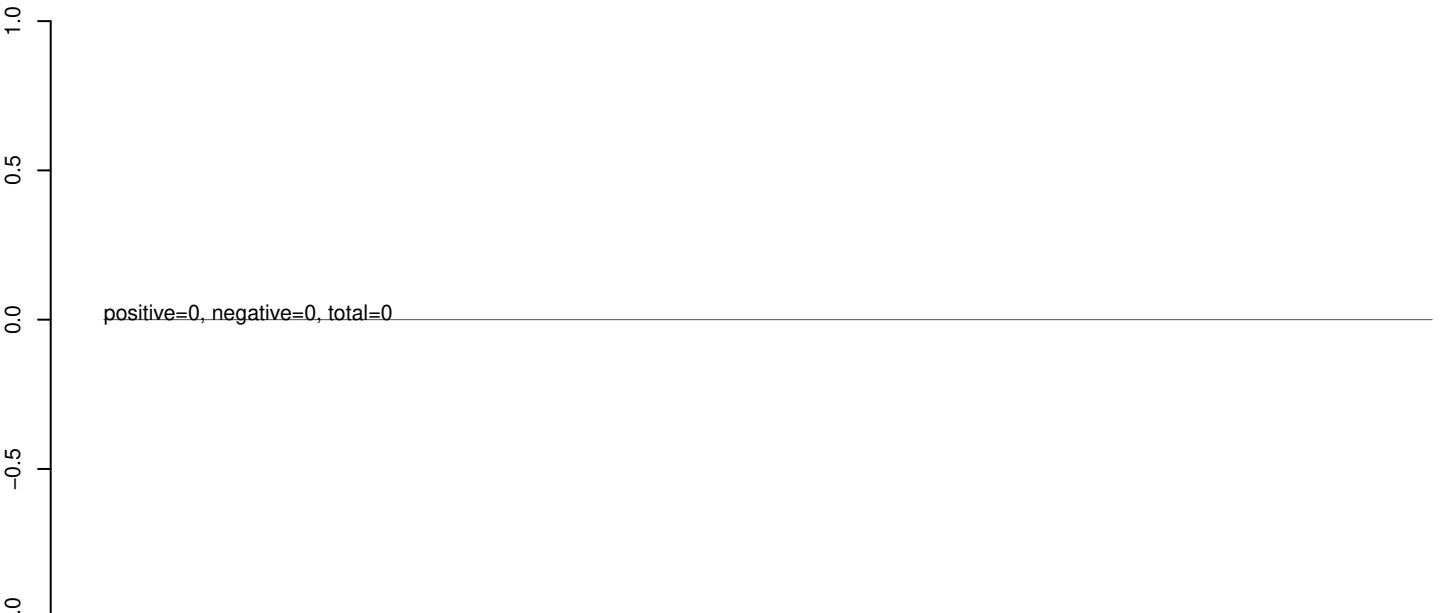


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AeAeg_Aag2_BUNV1_ES.rep



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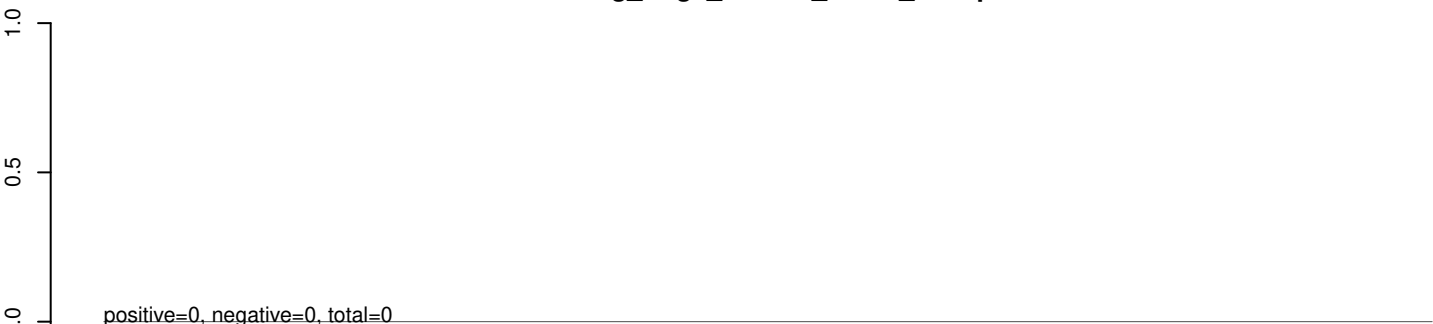
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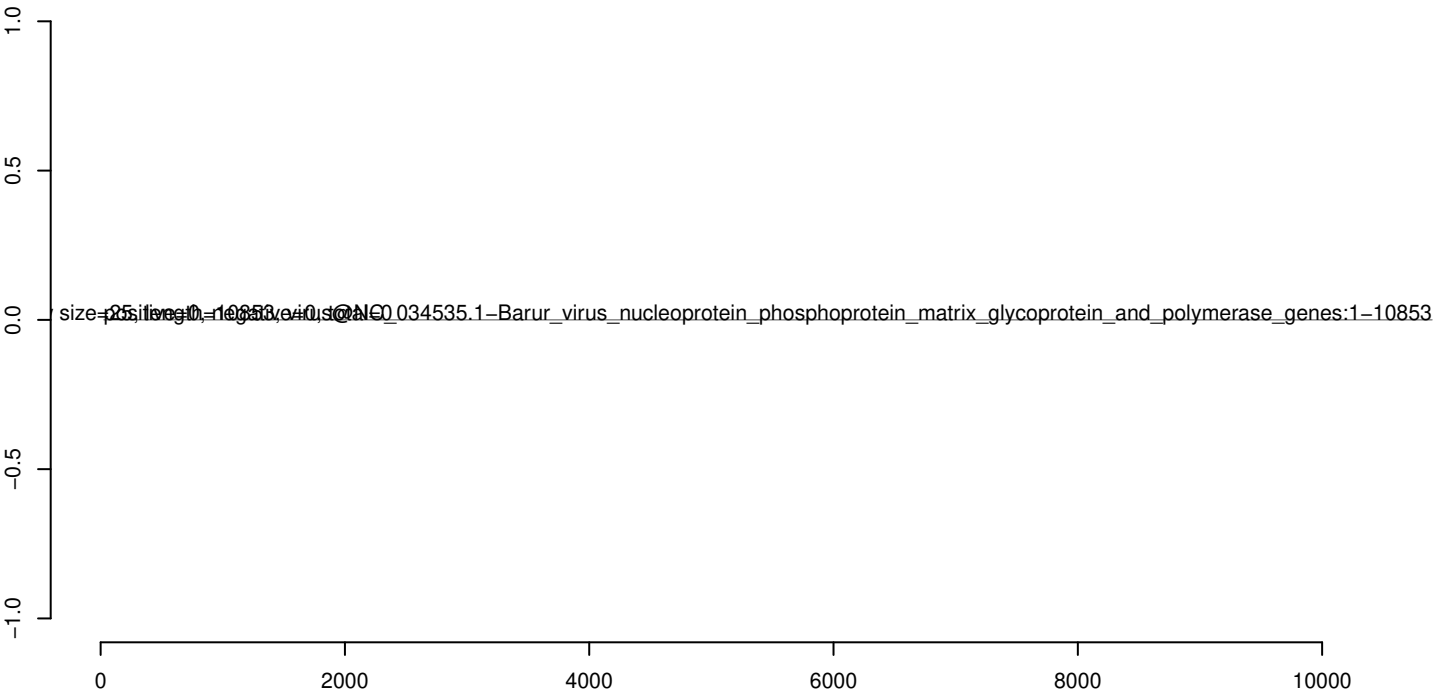
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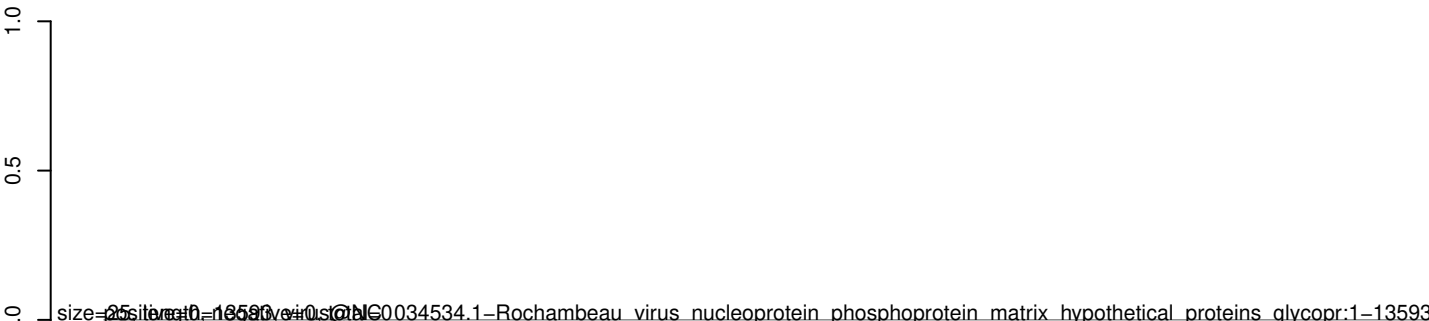
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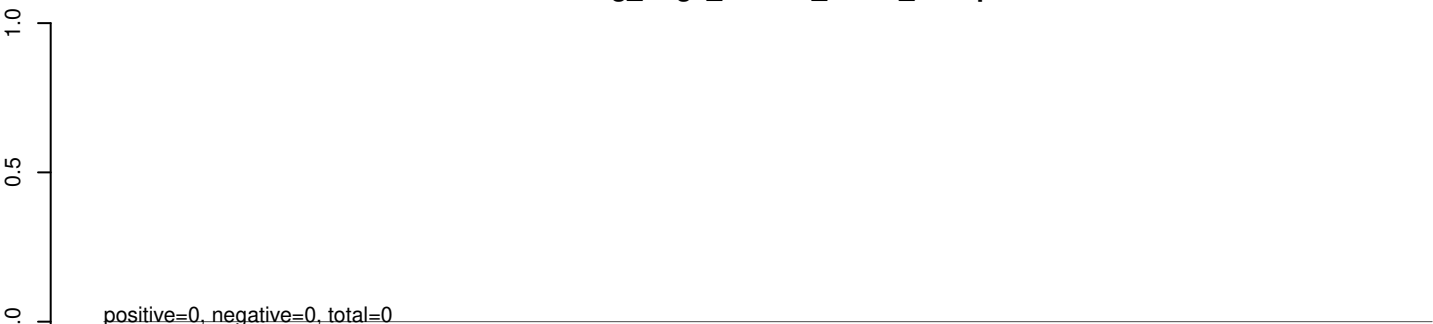


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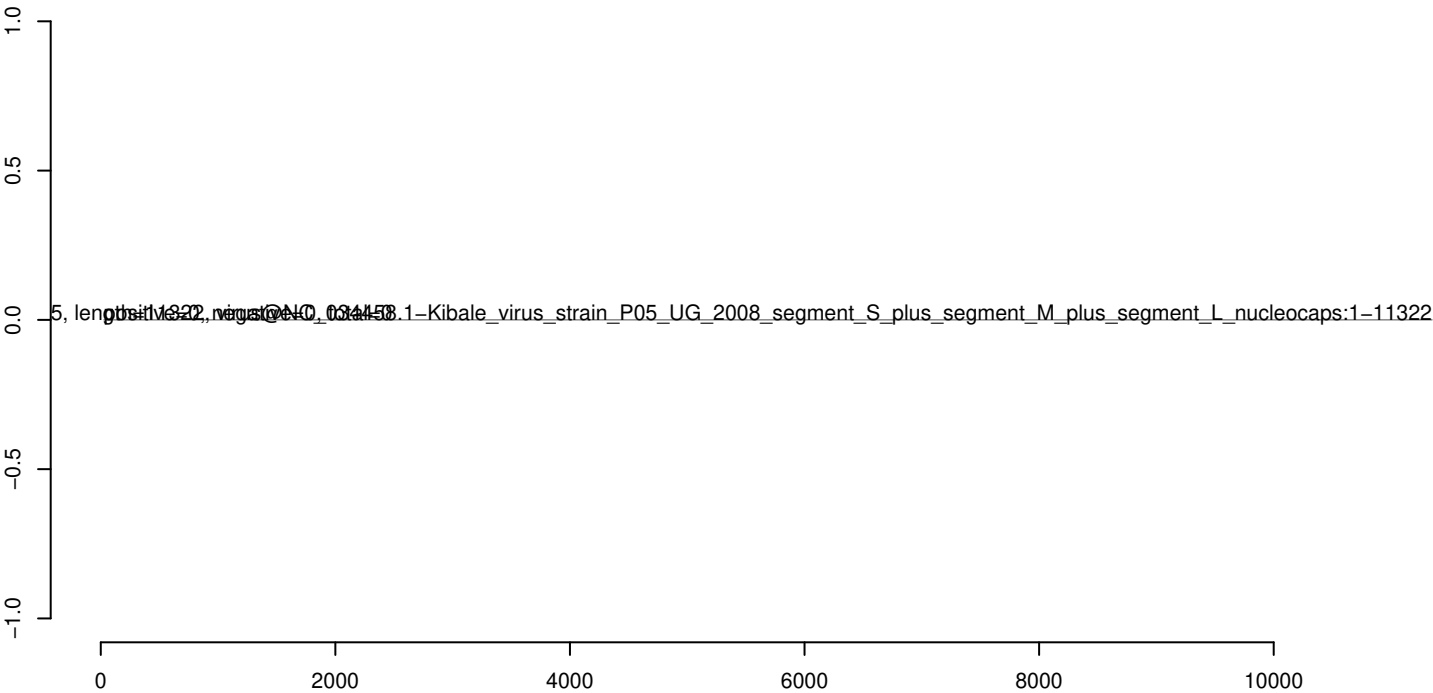
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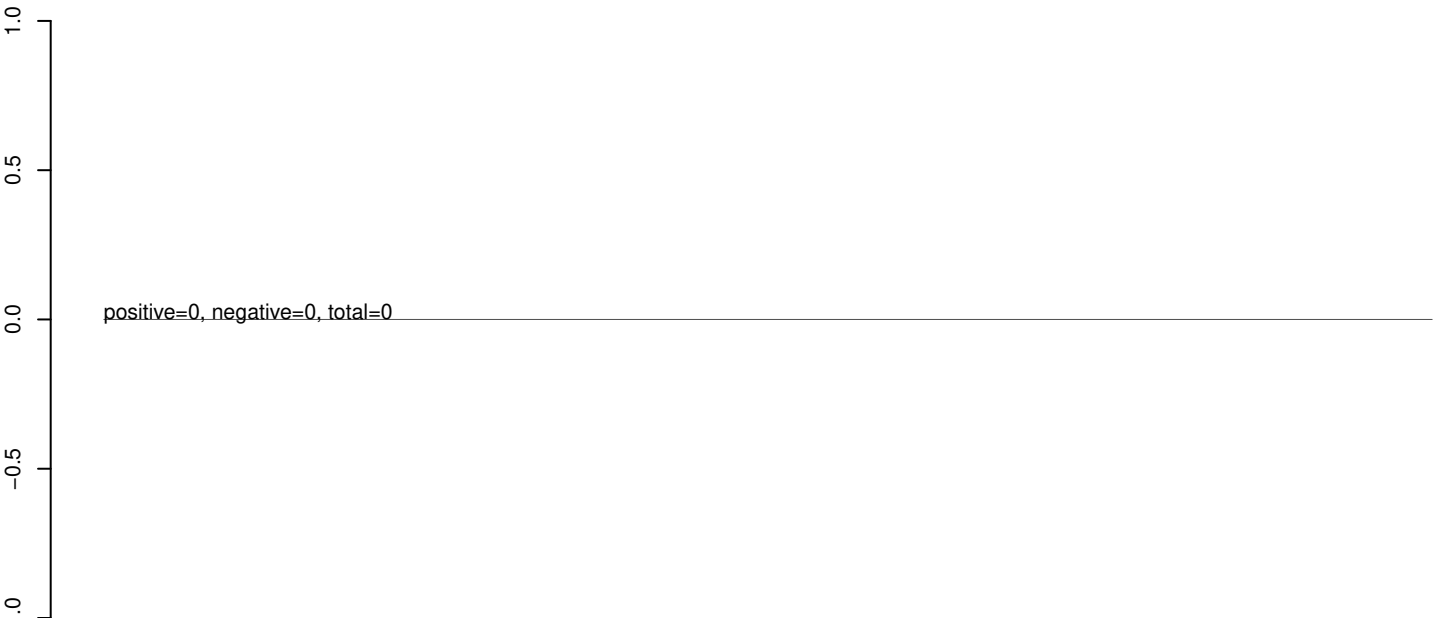
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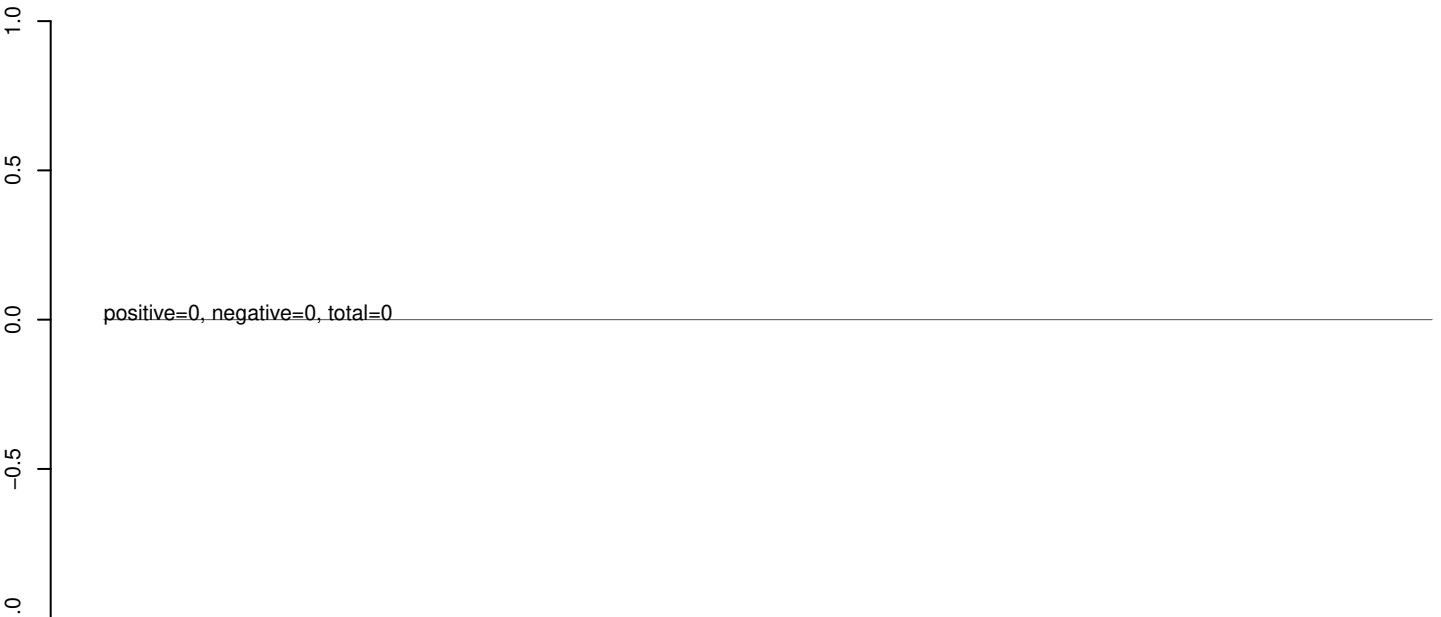
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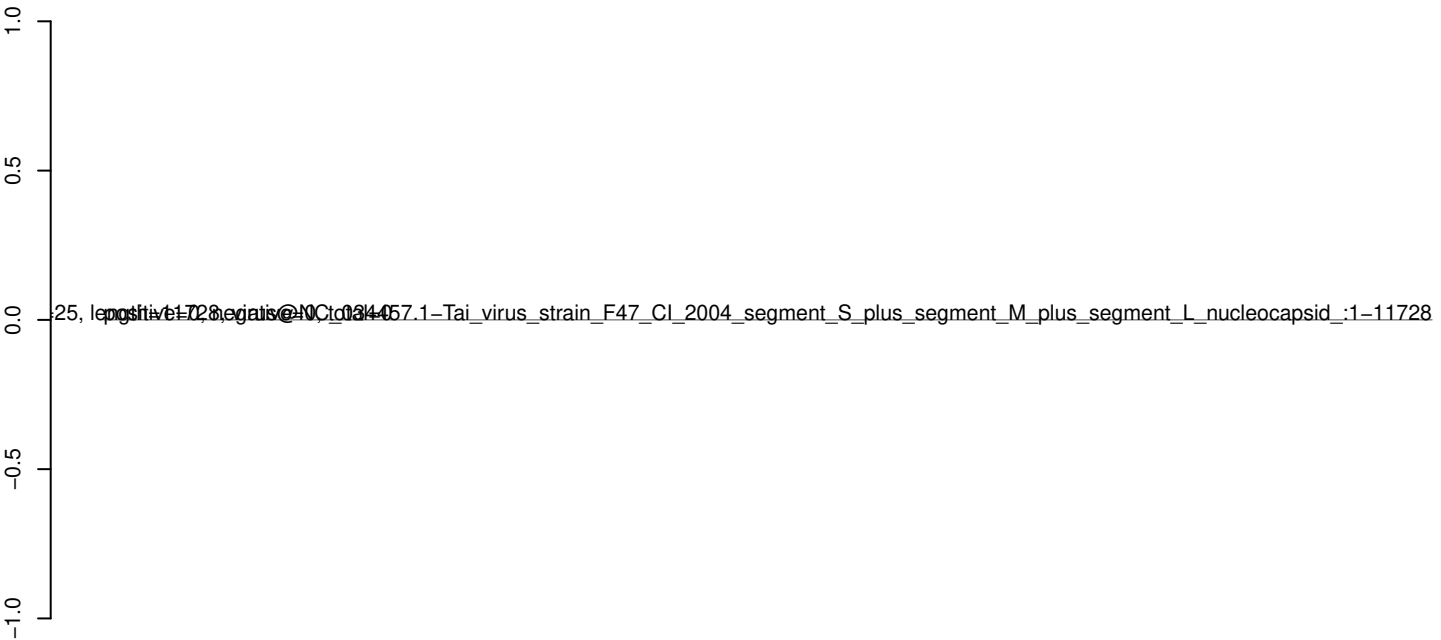
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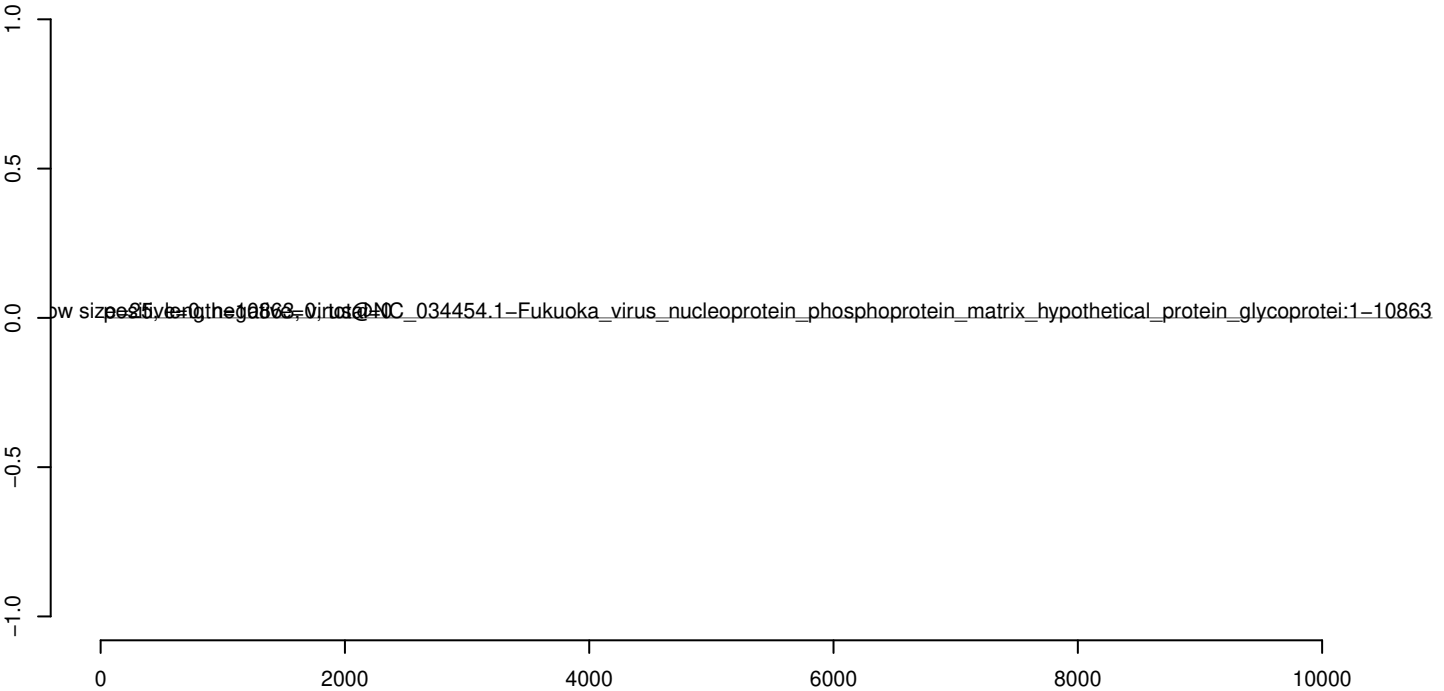
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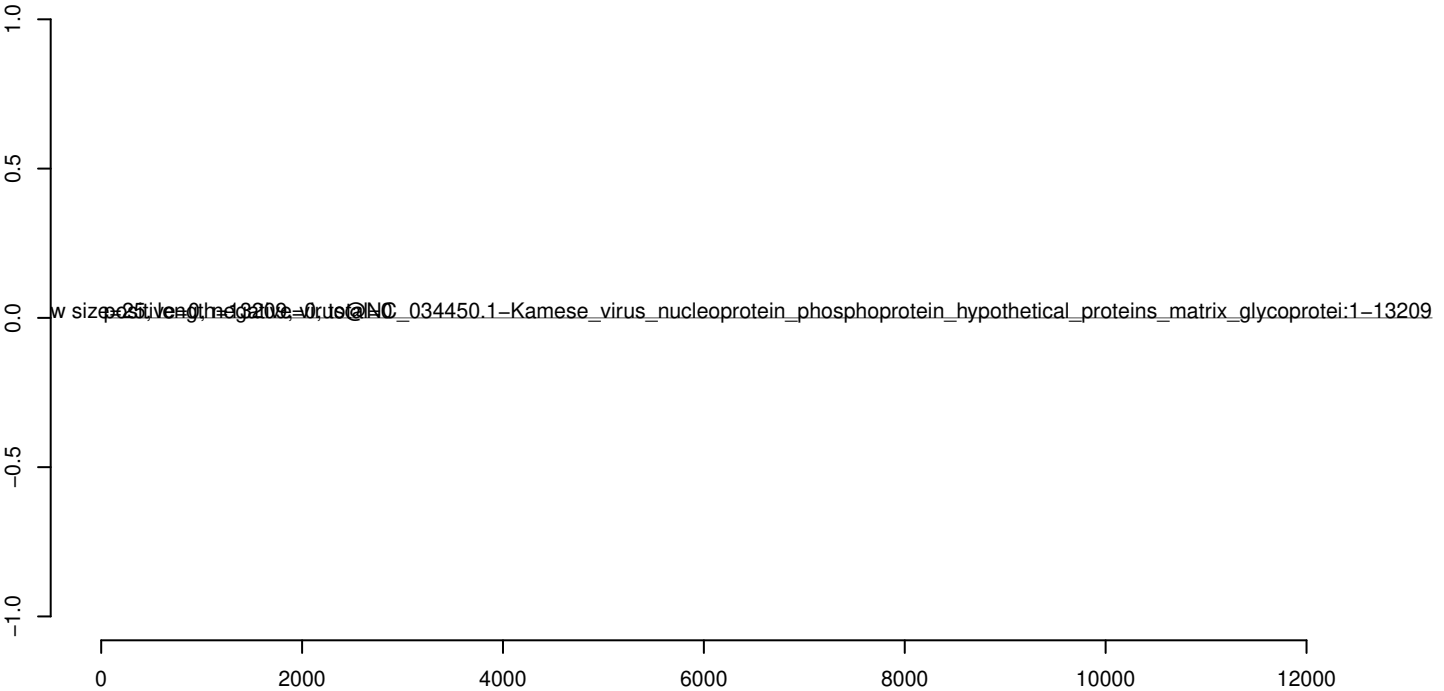
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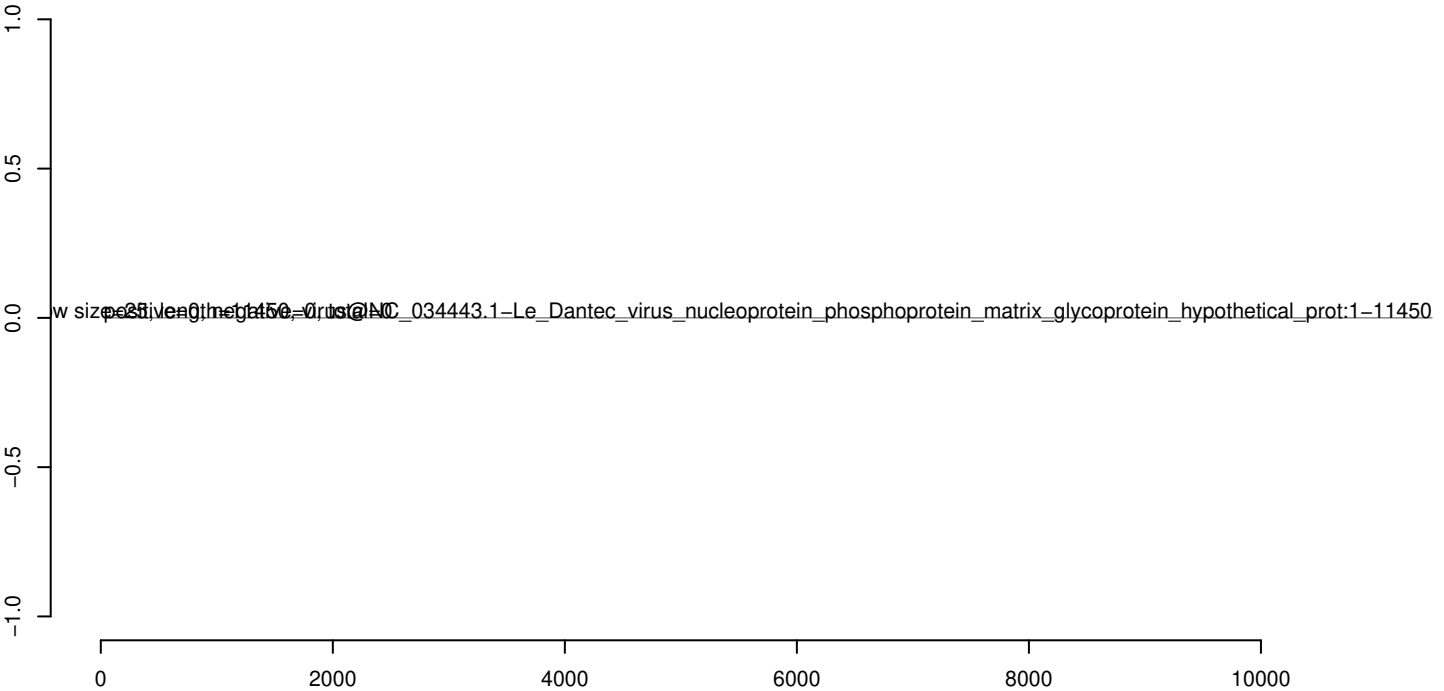
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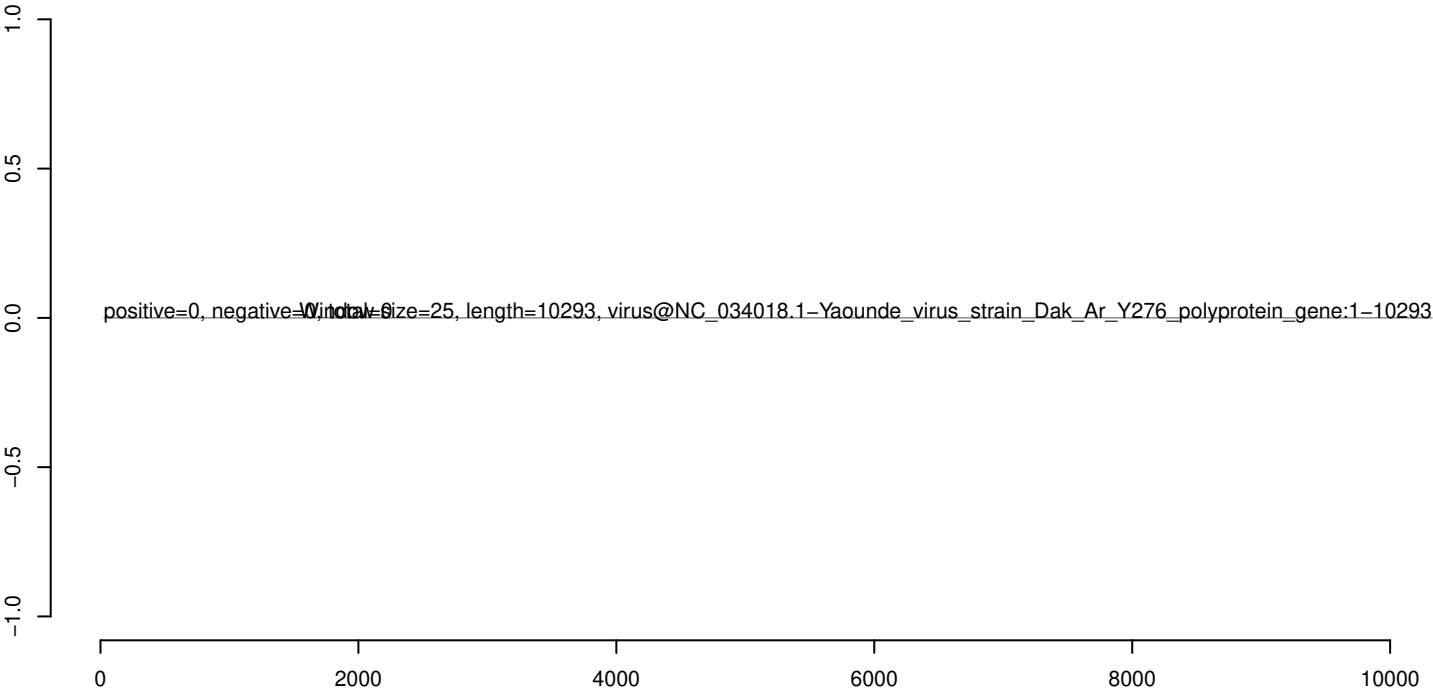
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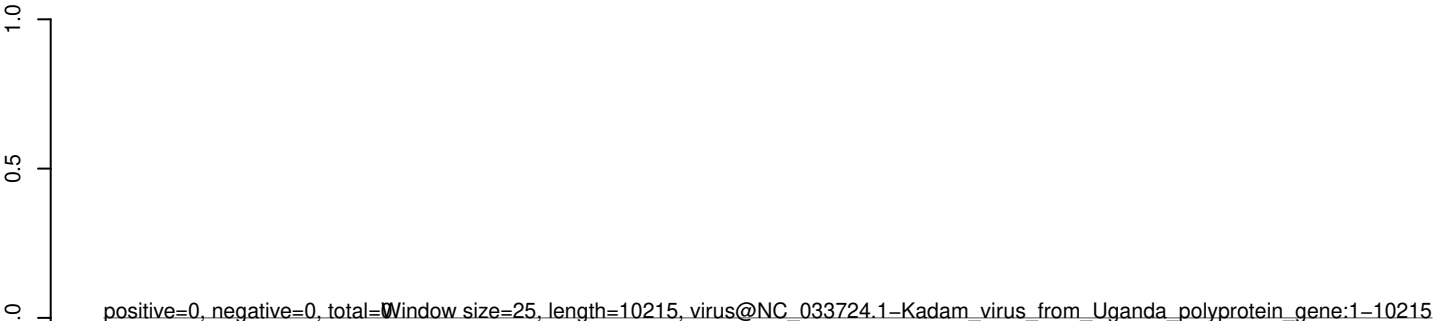
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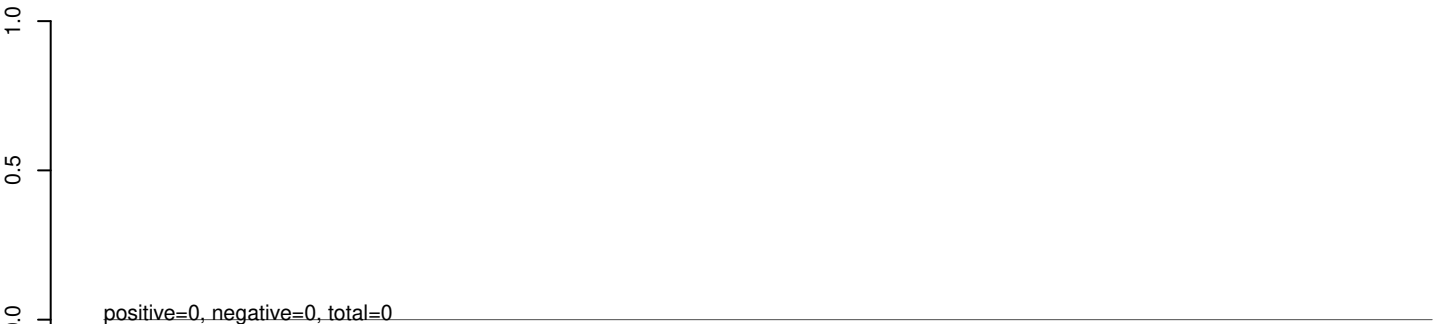
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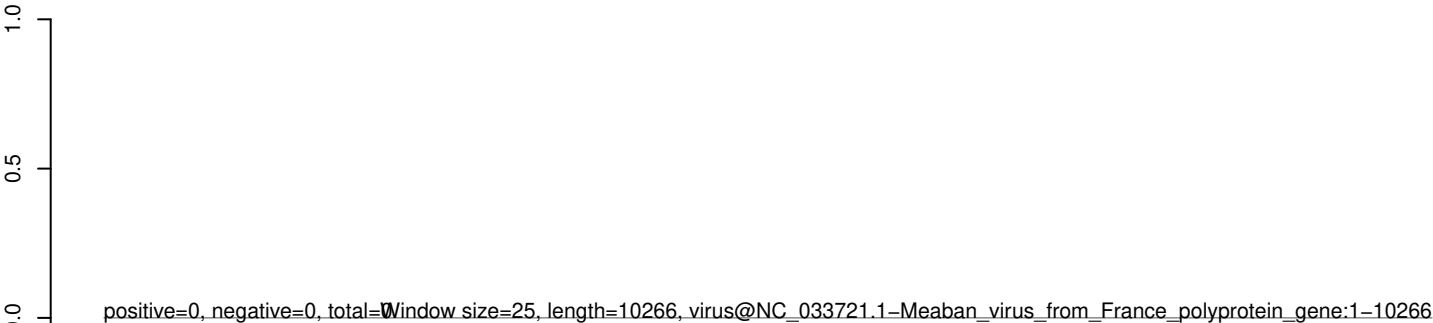
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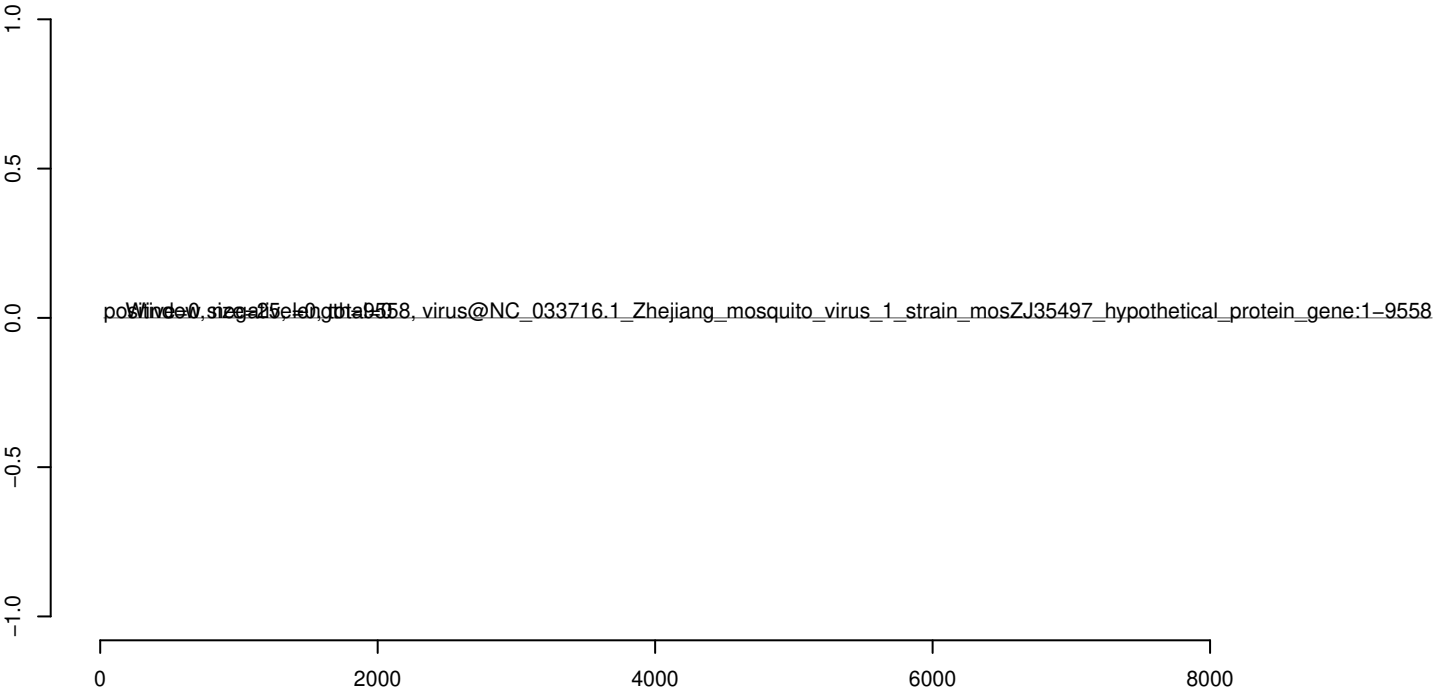
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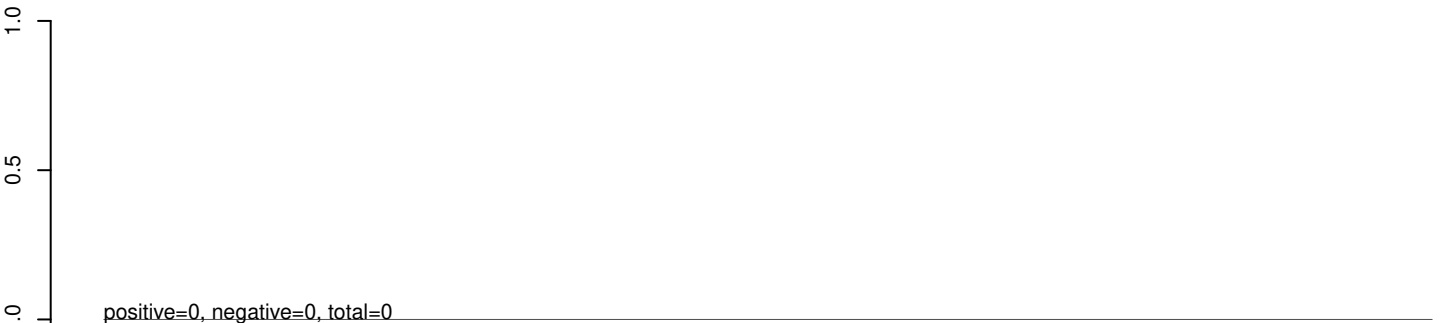
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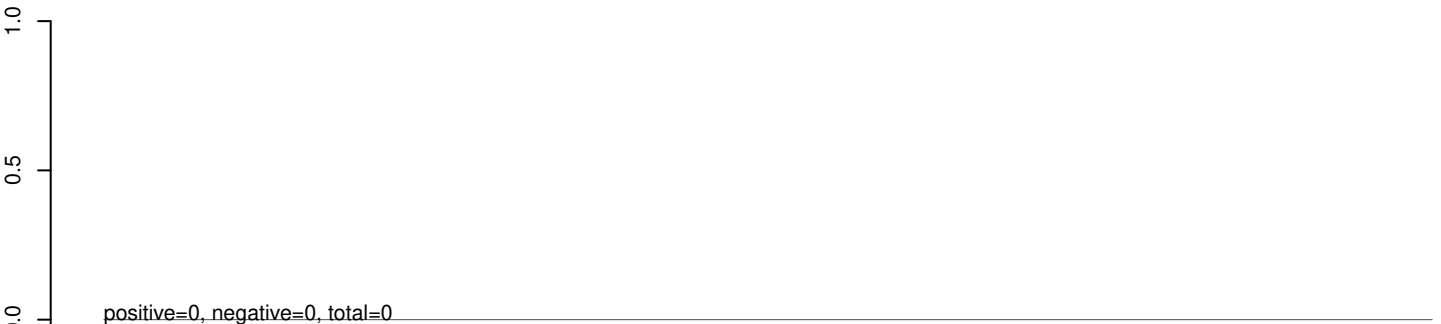
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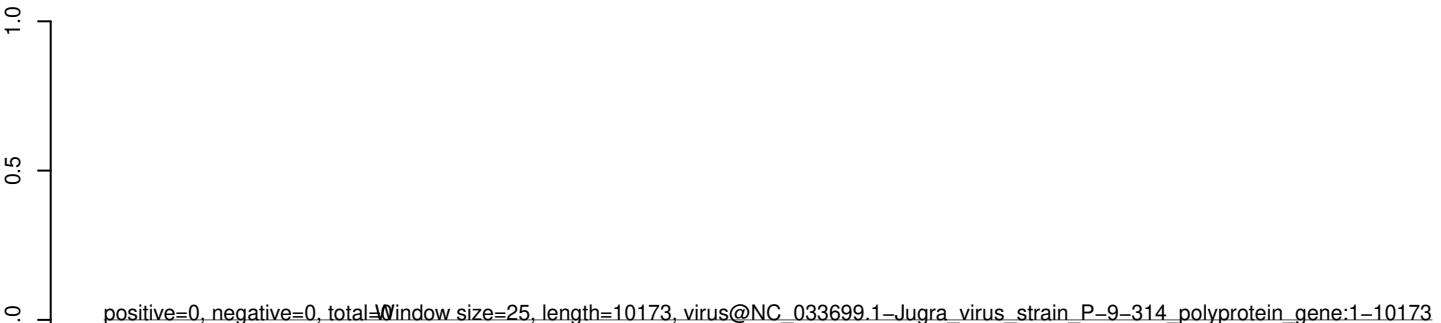
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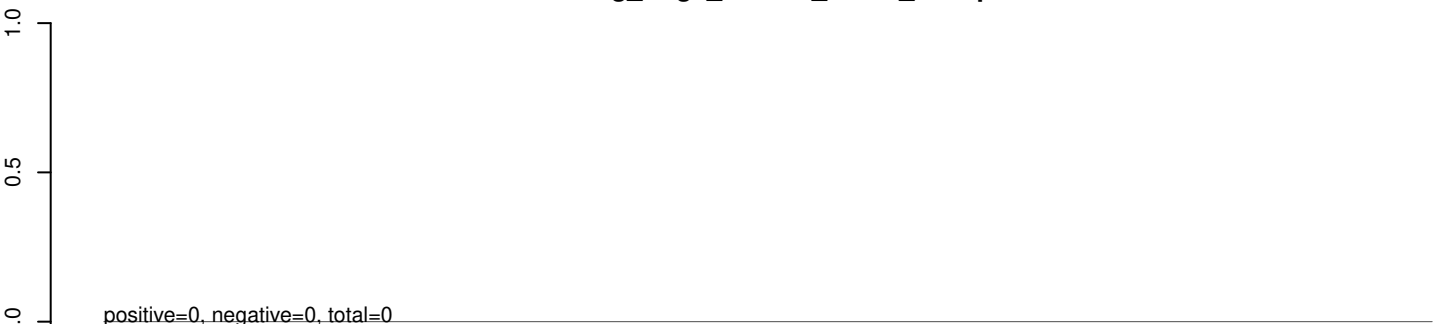
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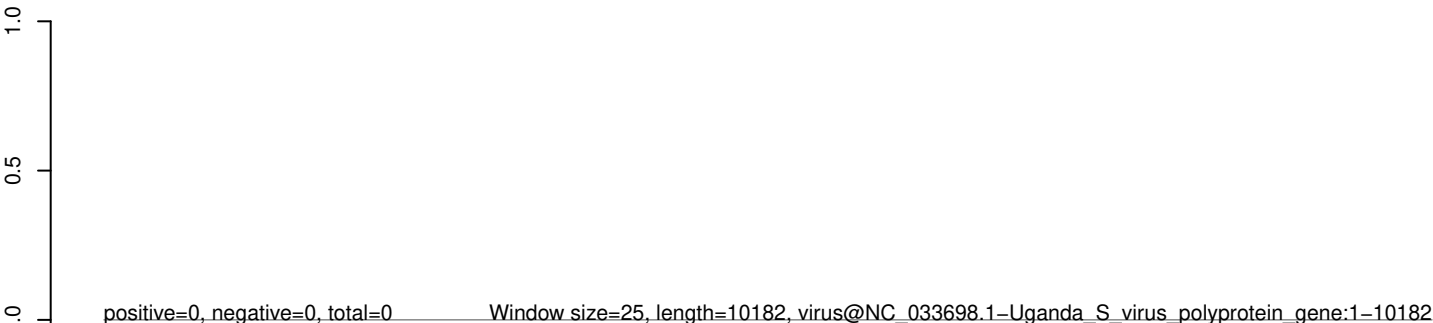
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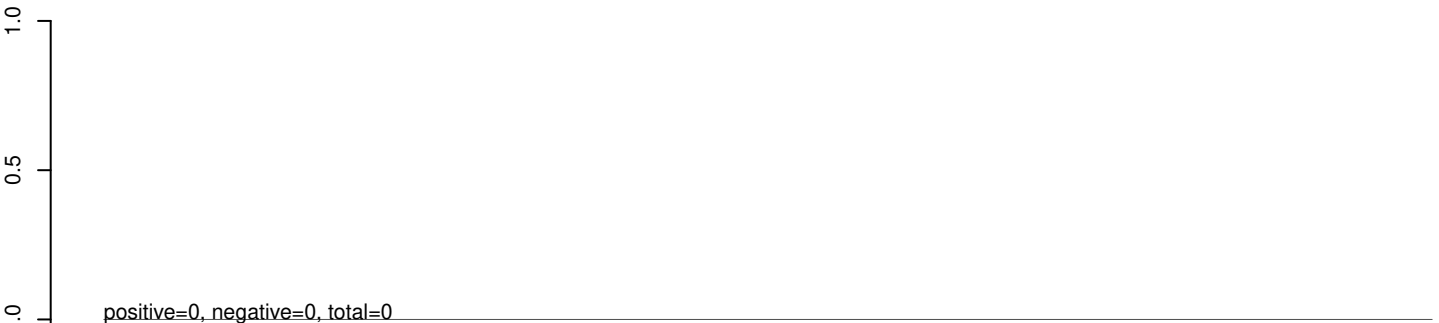
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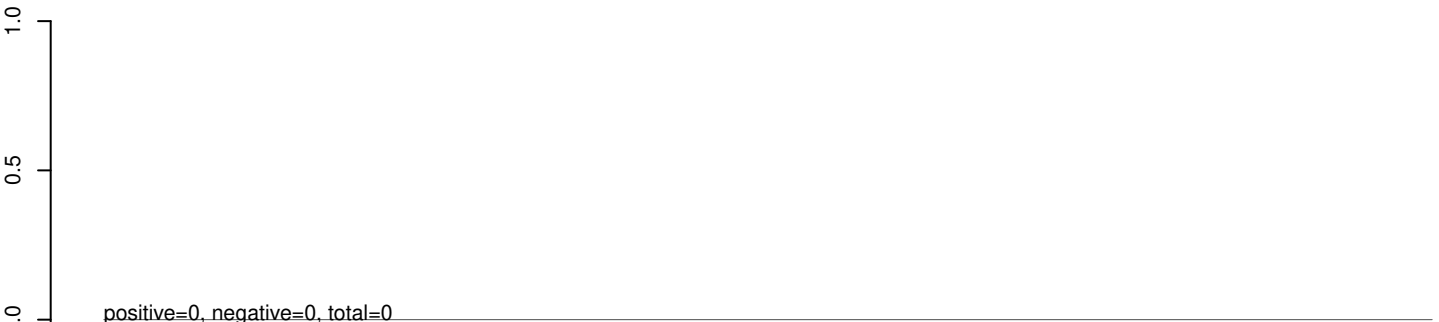
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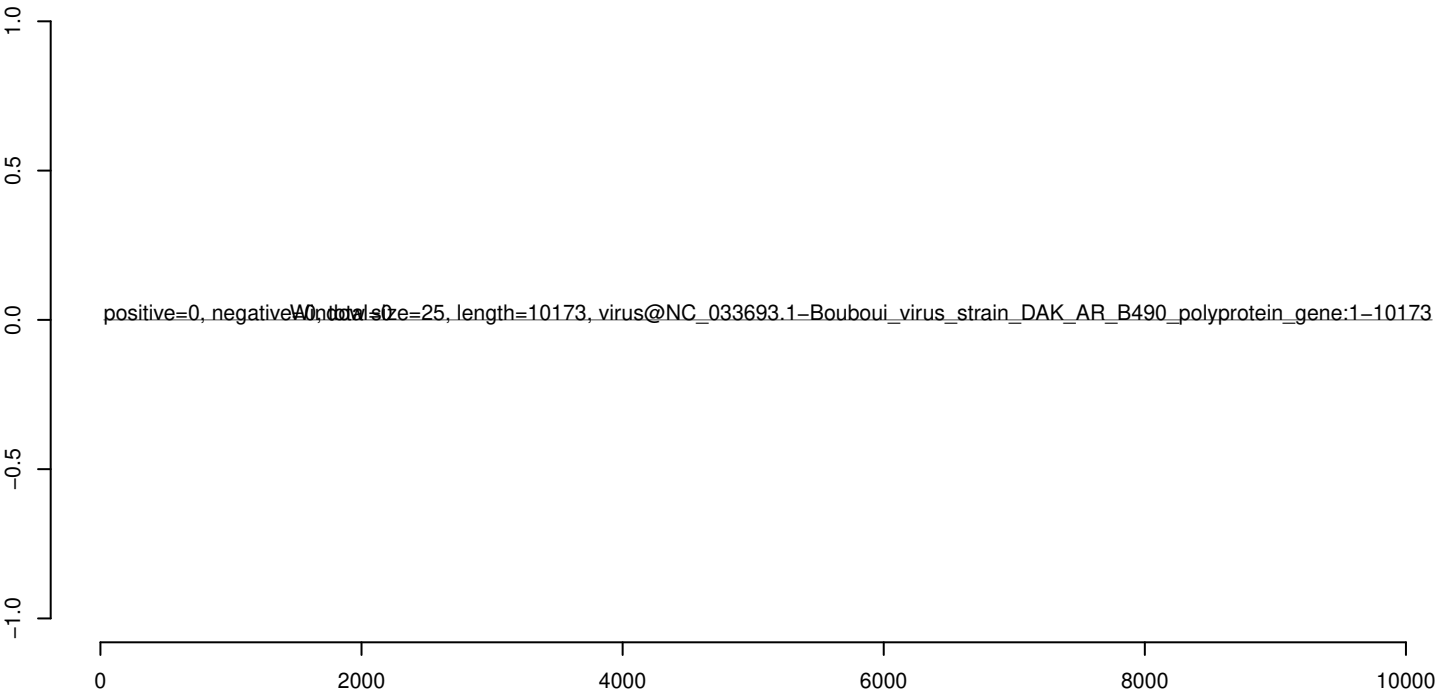
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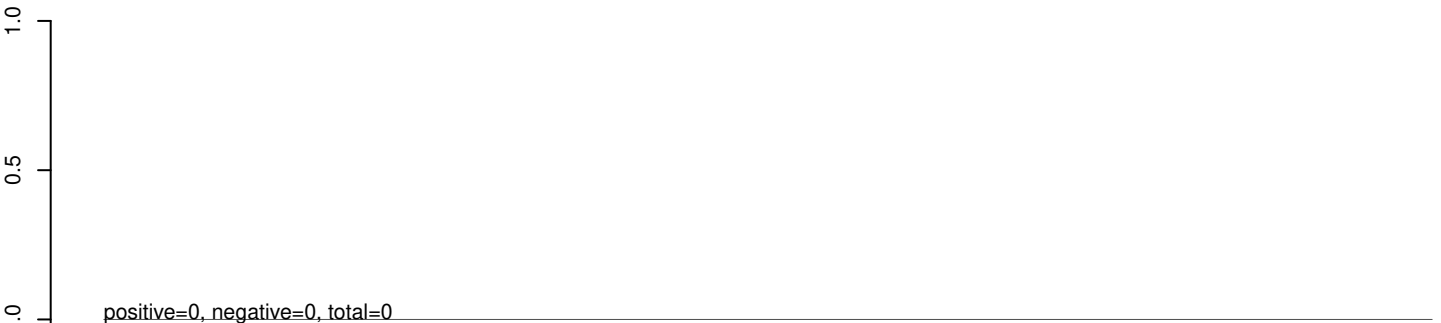
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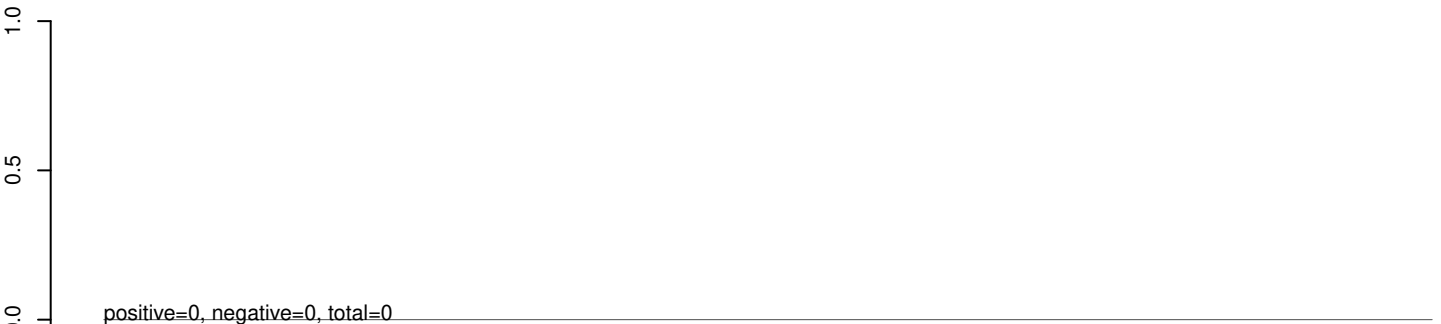
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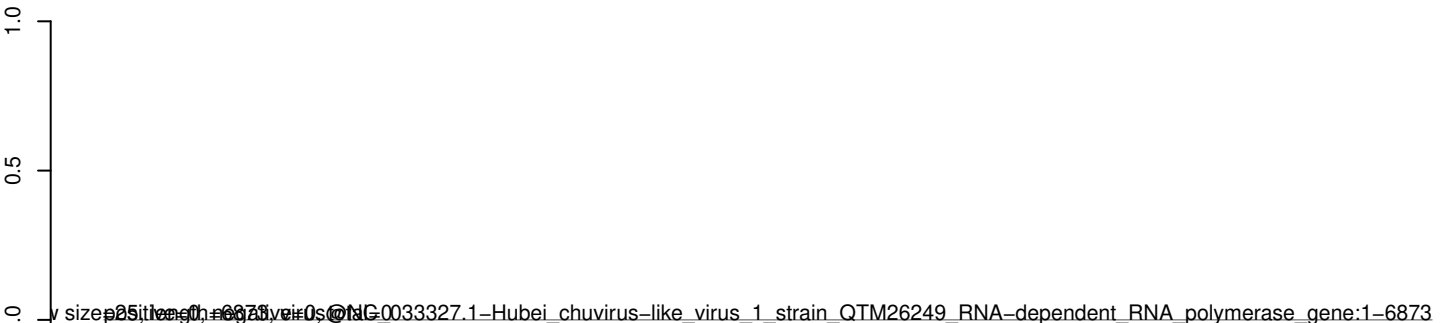
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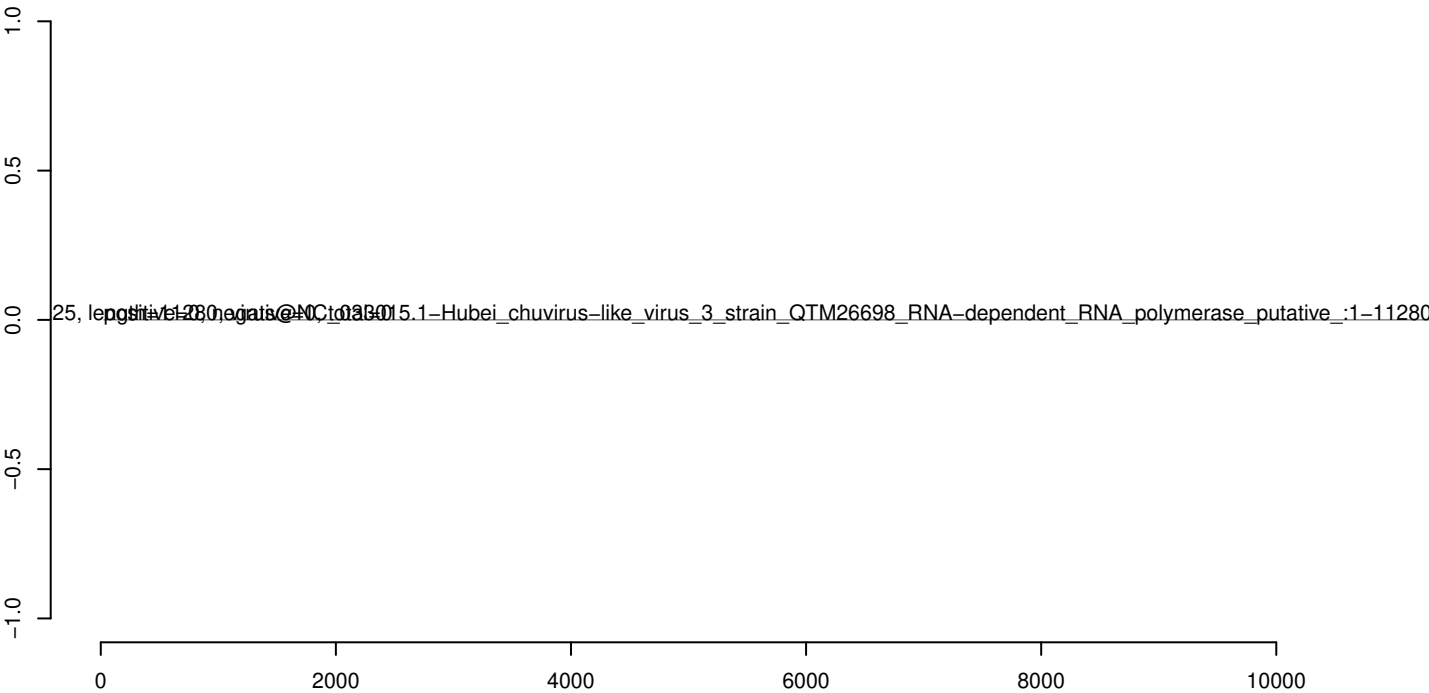
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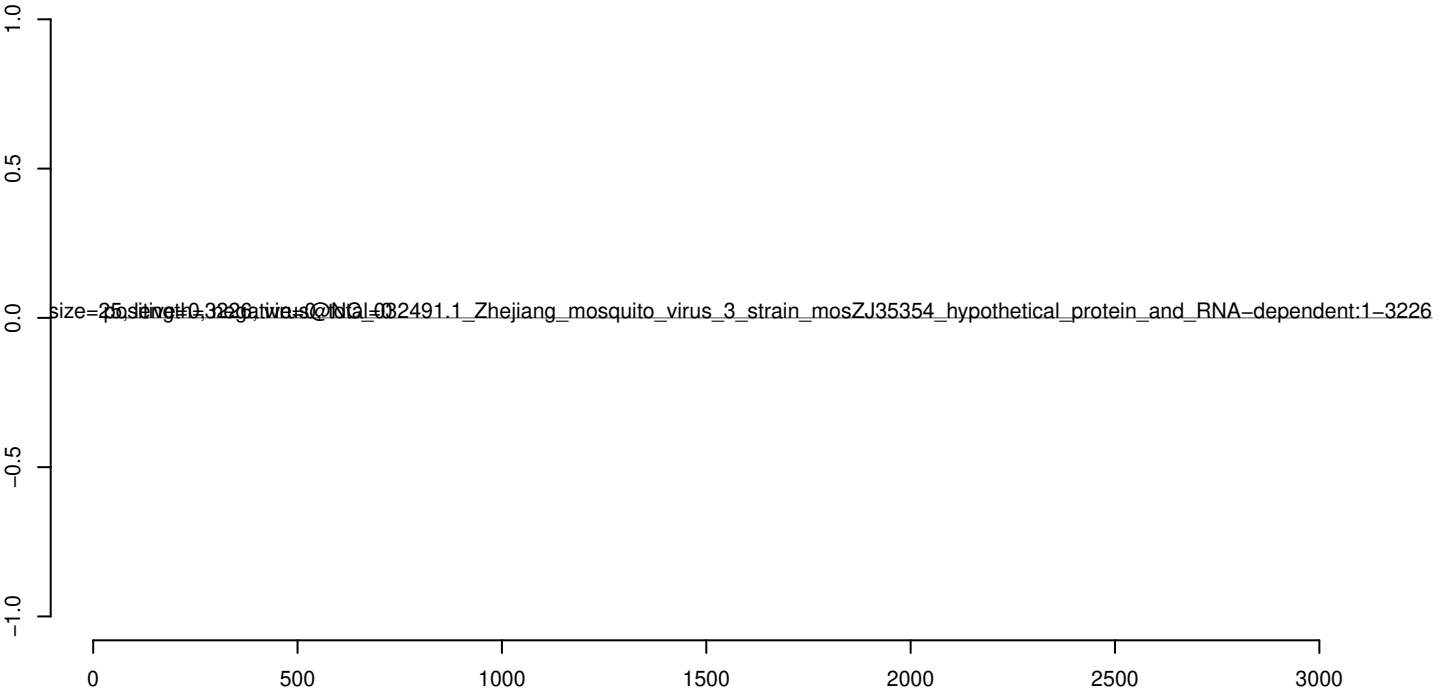
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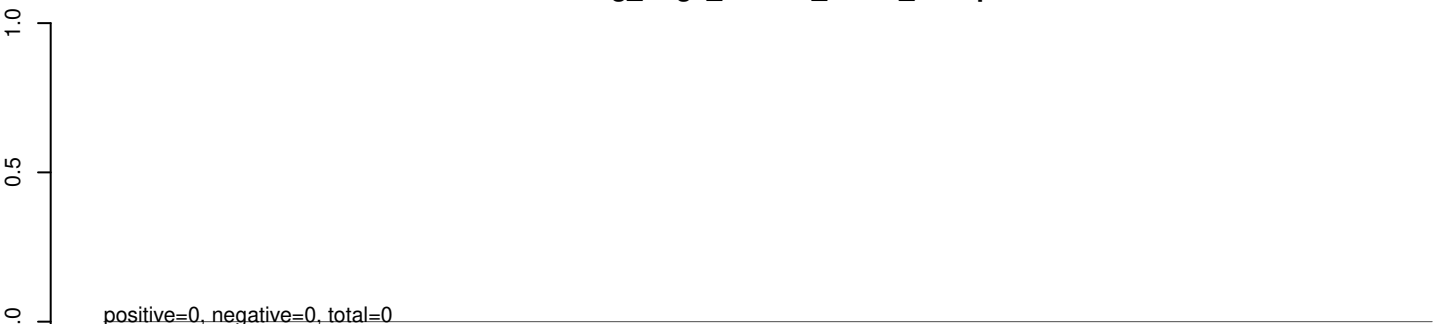
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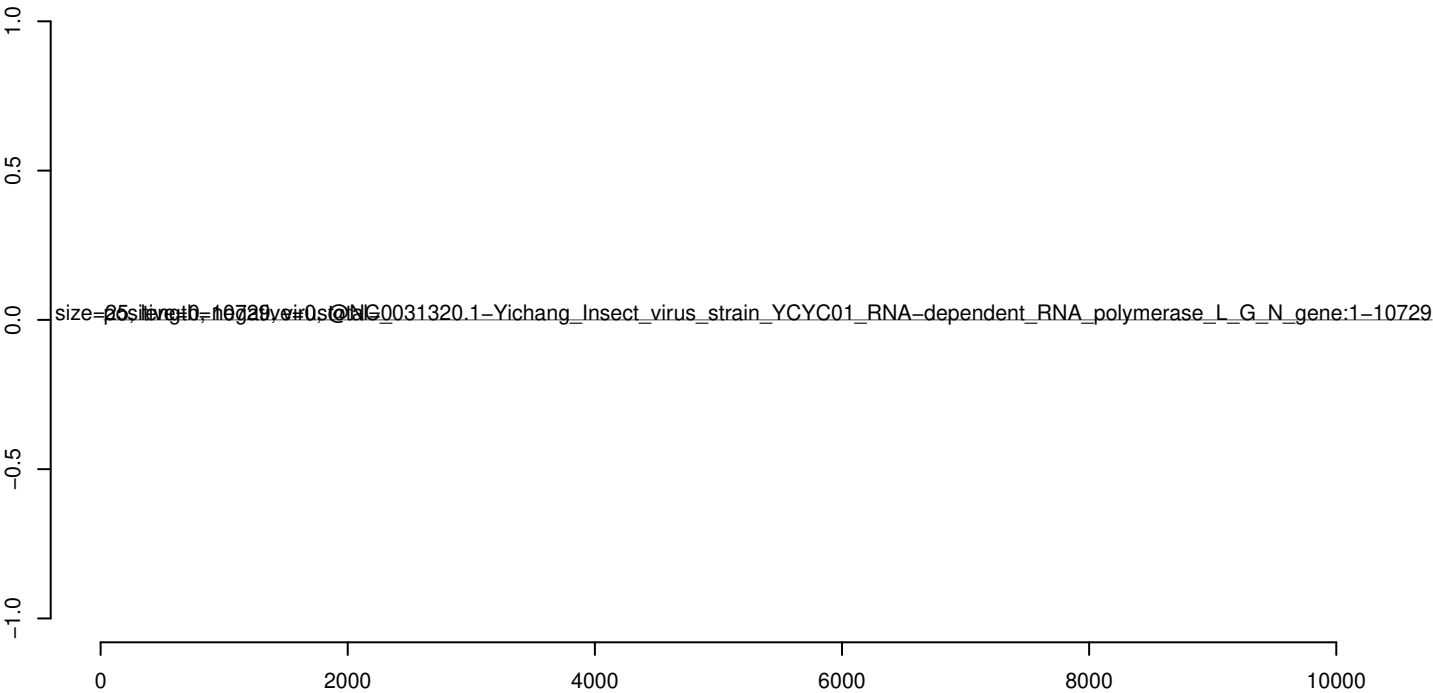
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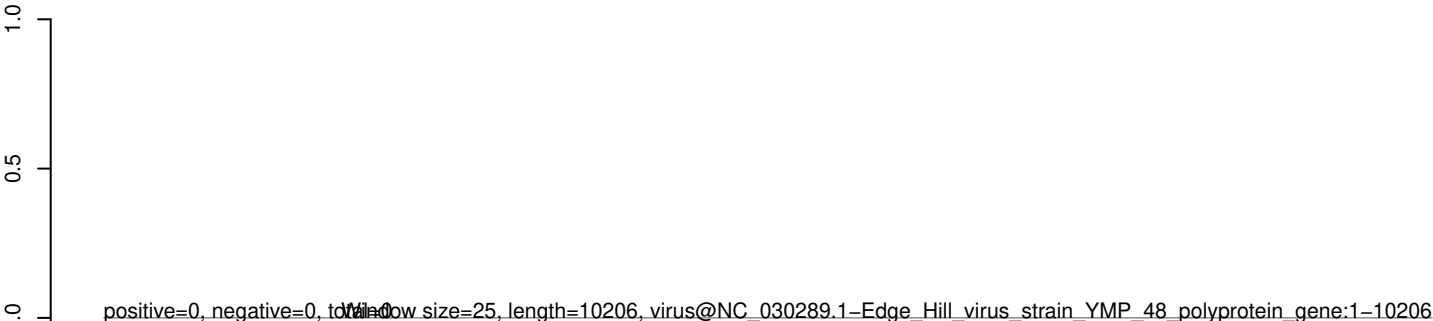
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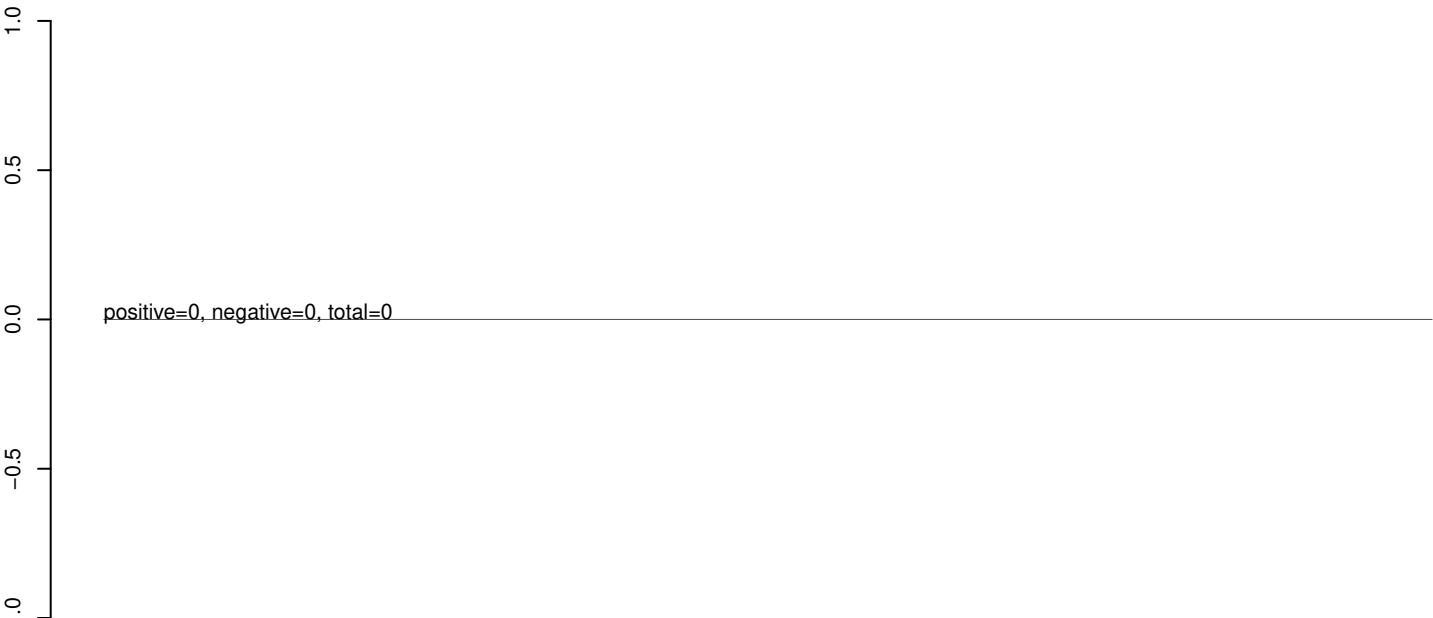
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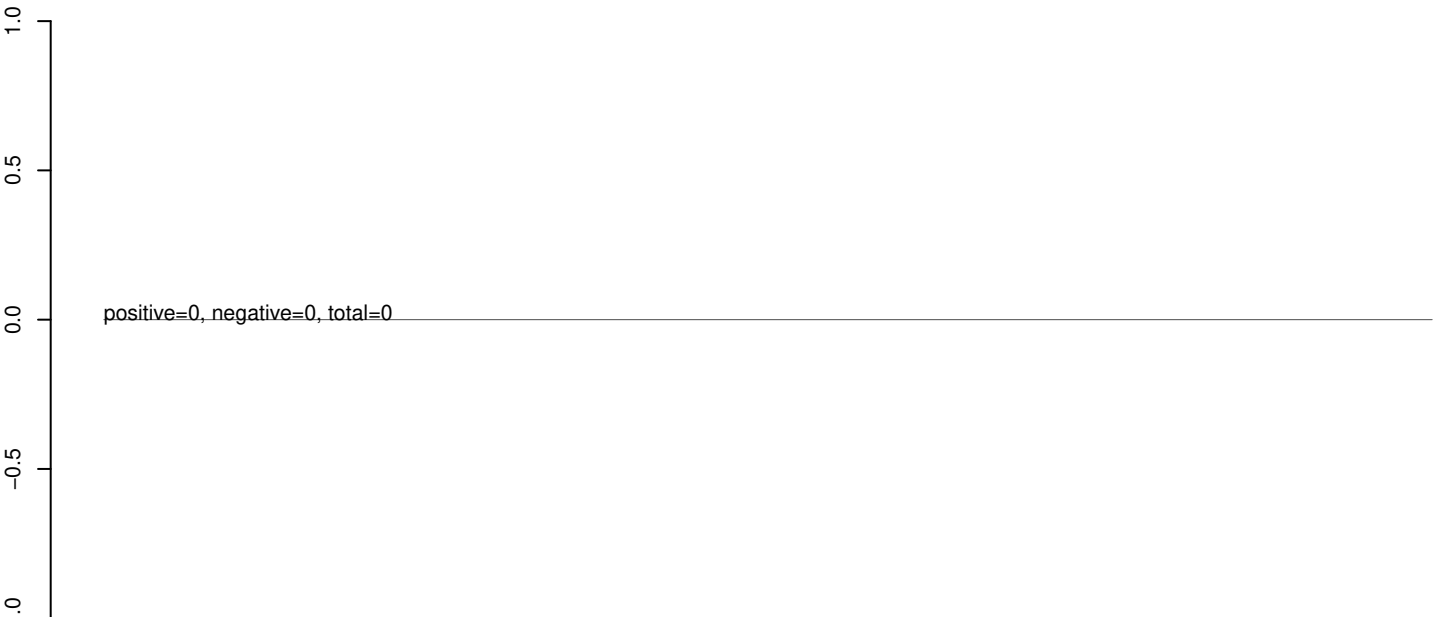
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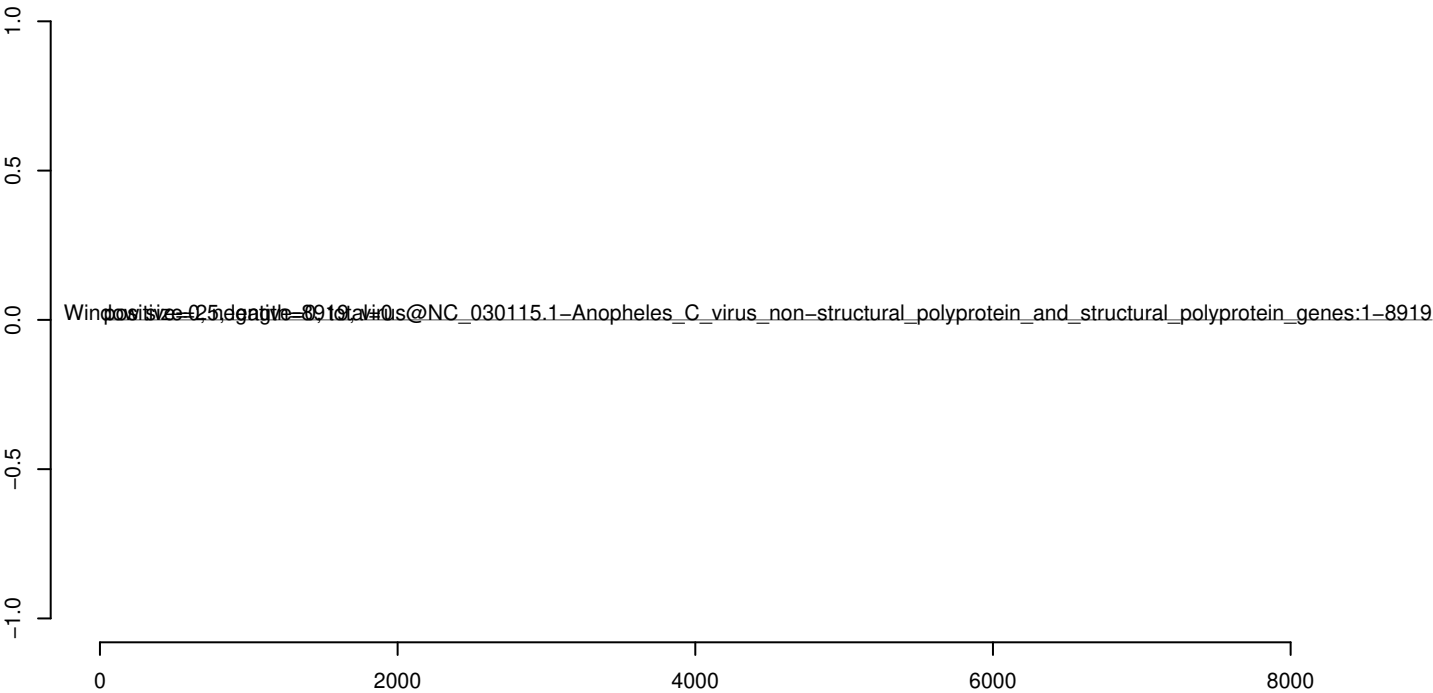
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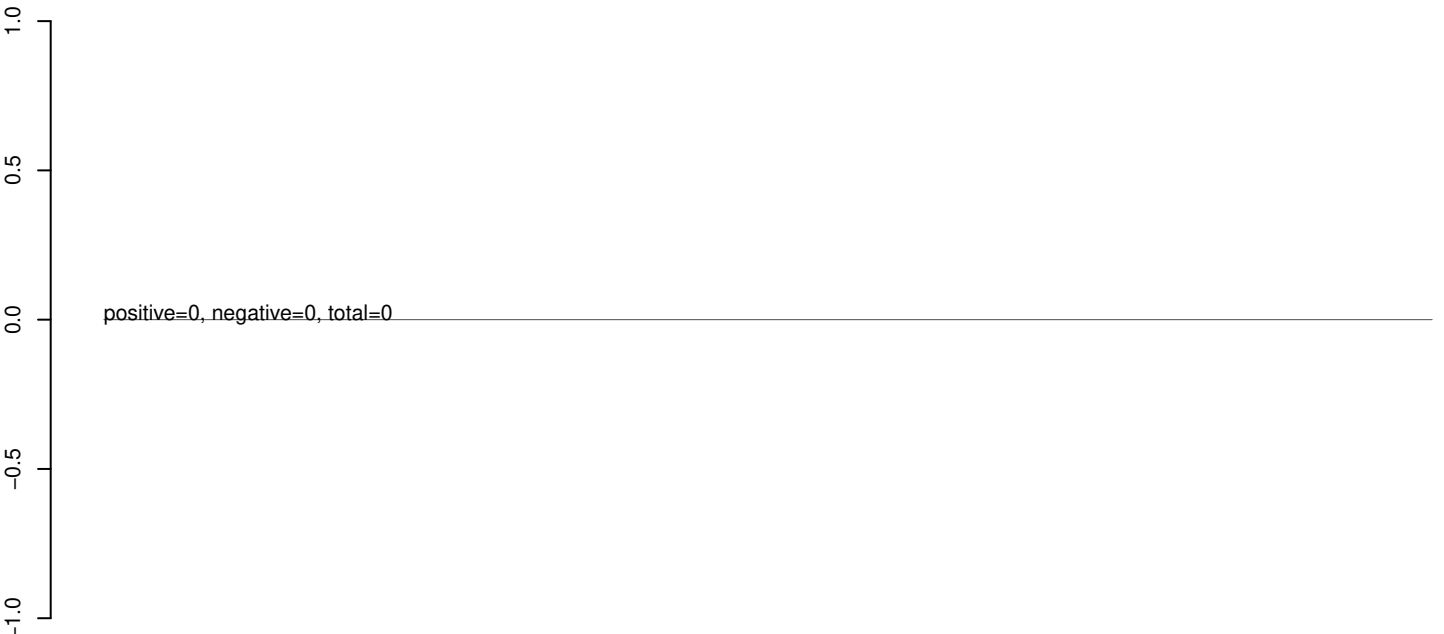
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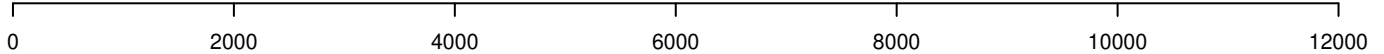
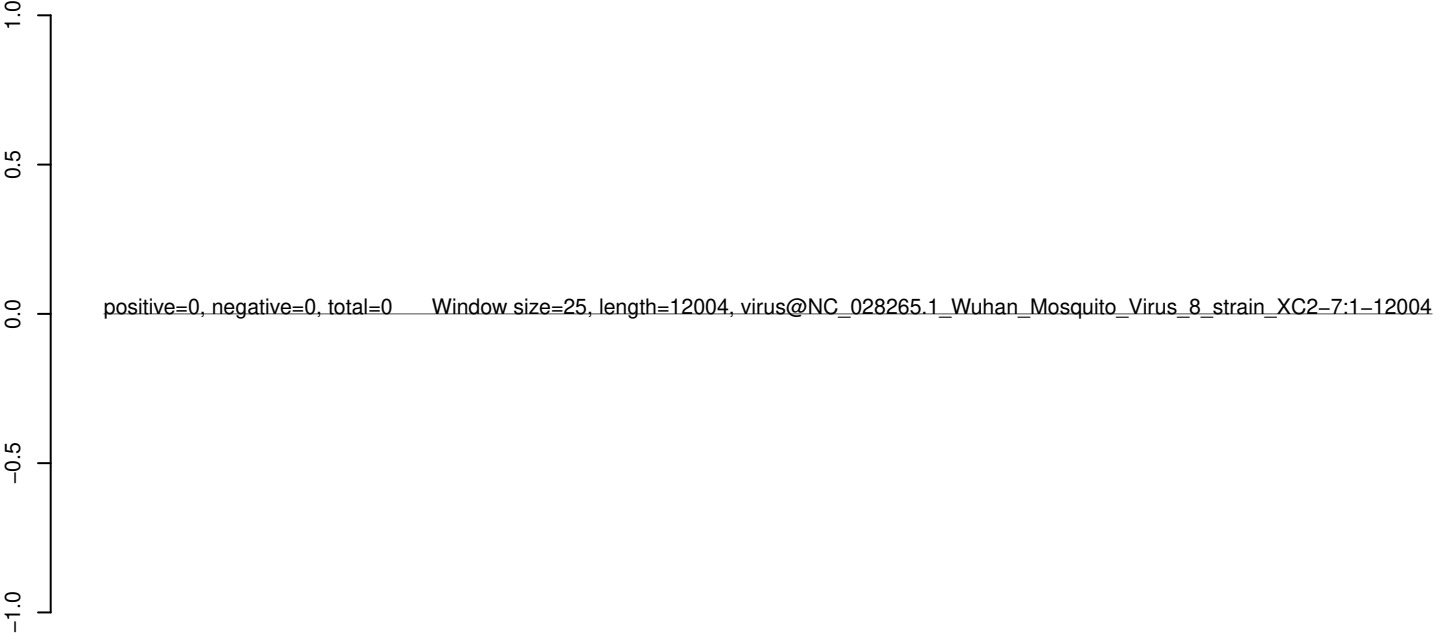
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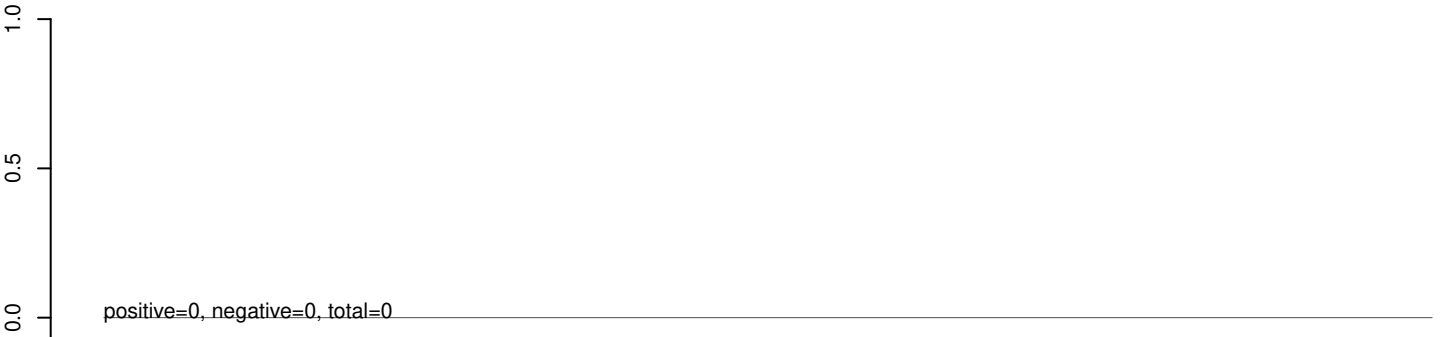
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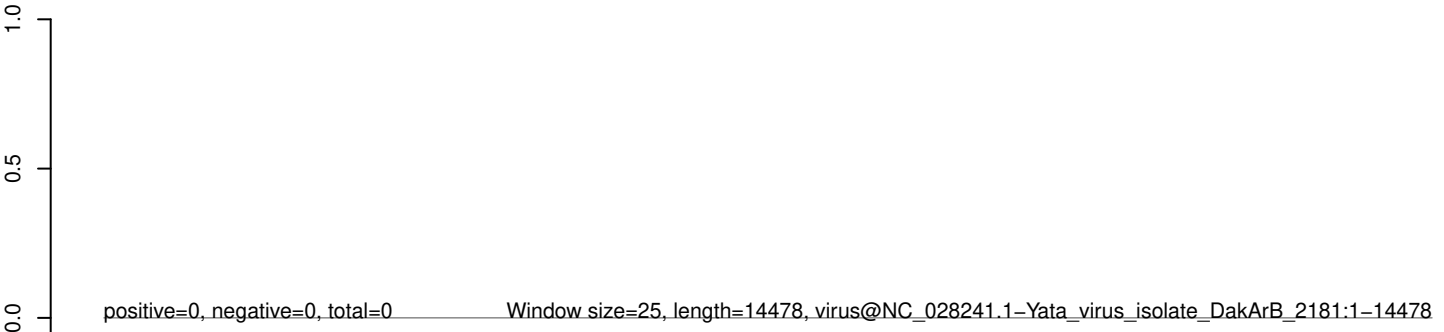
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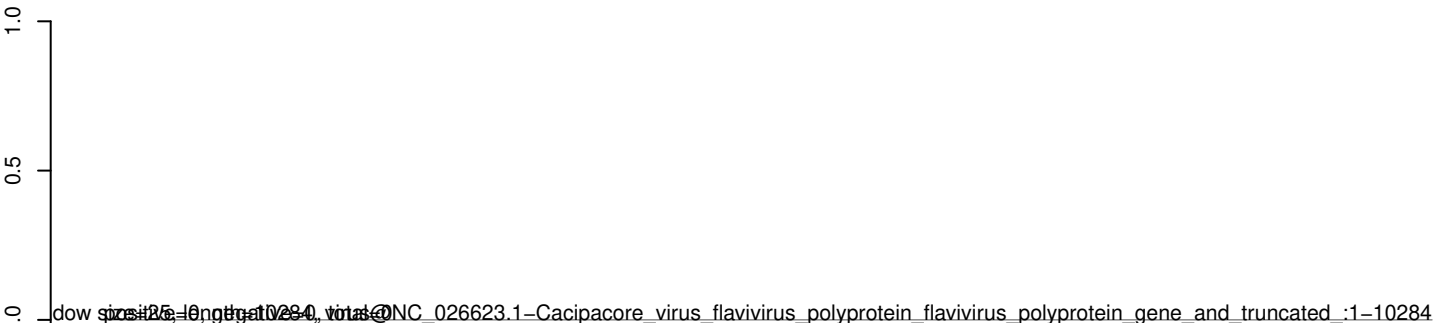
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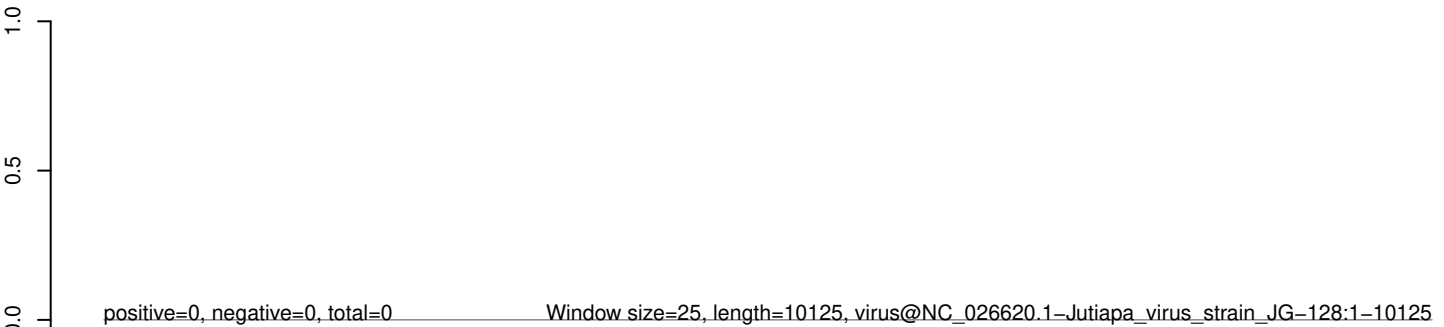
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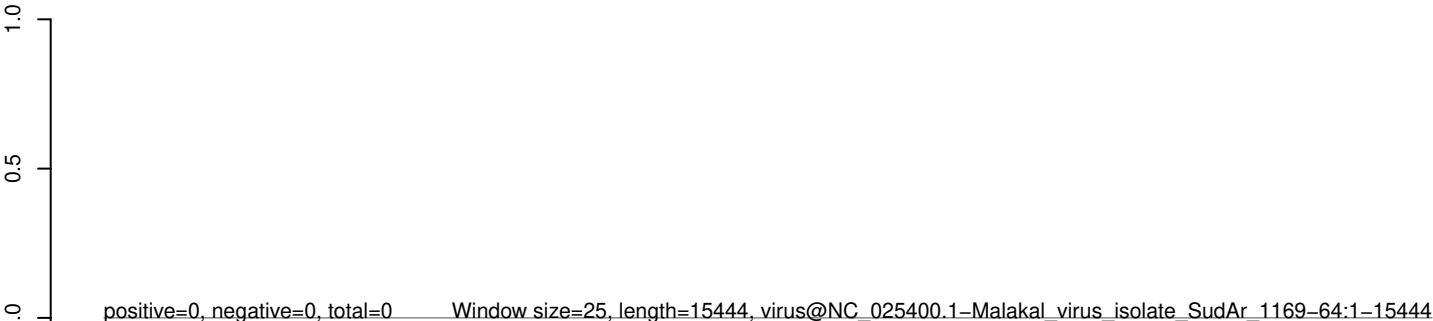
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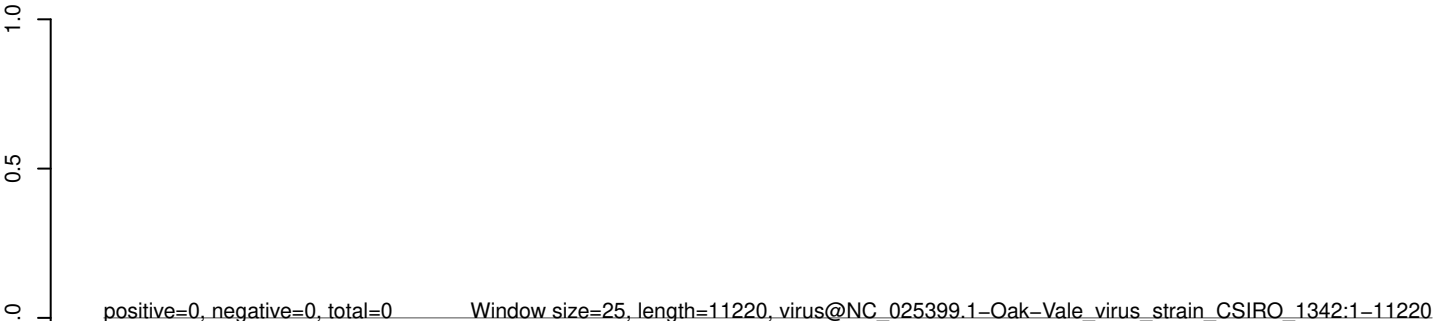
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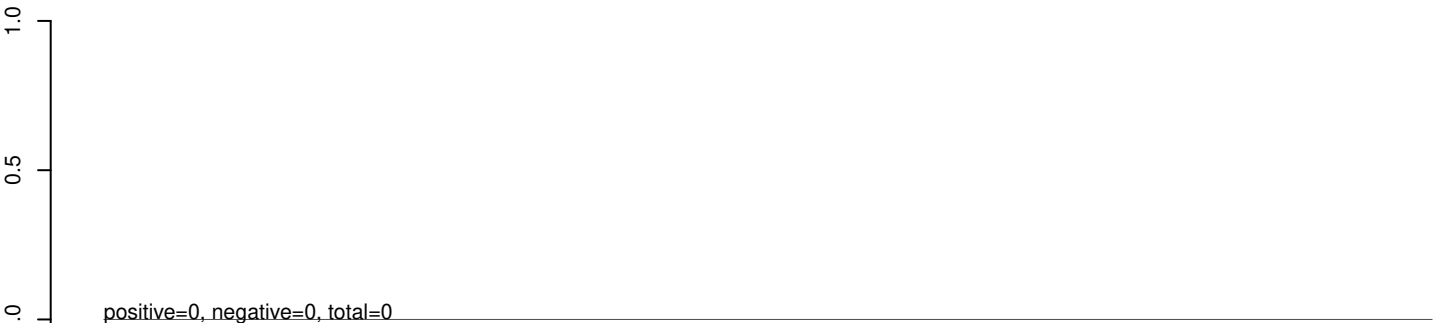


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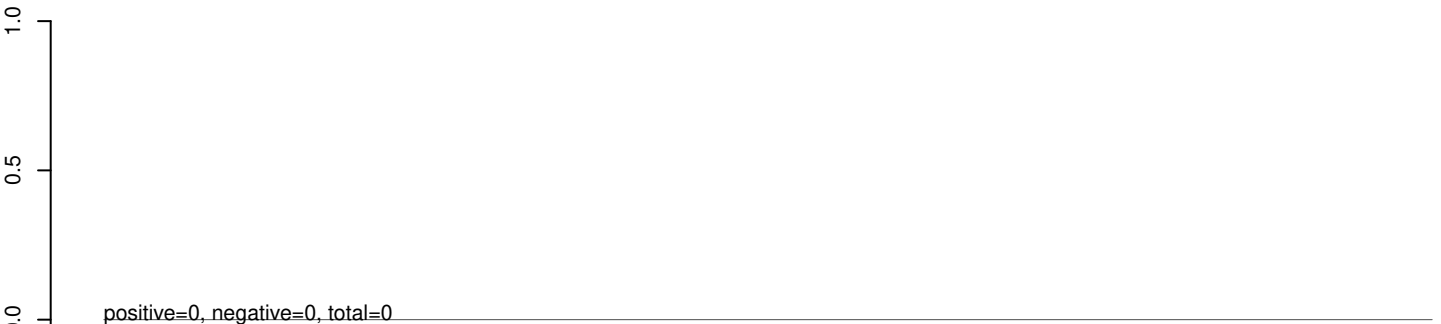


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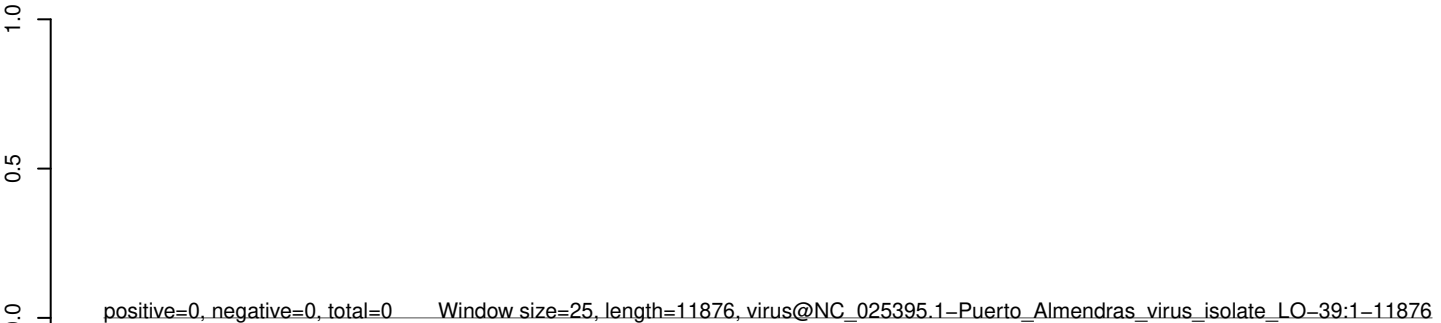
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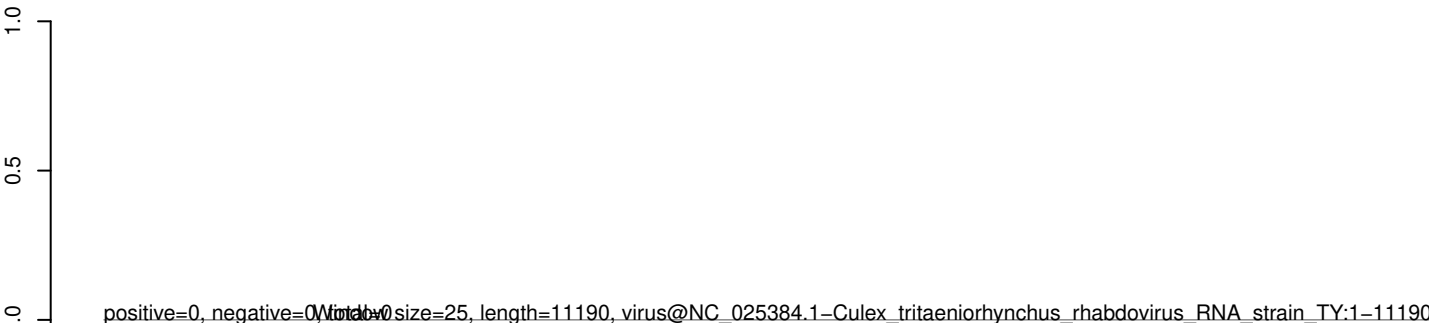
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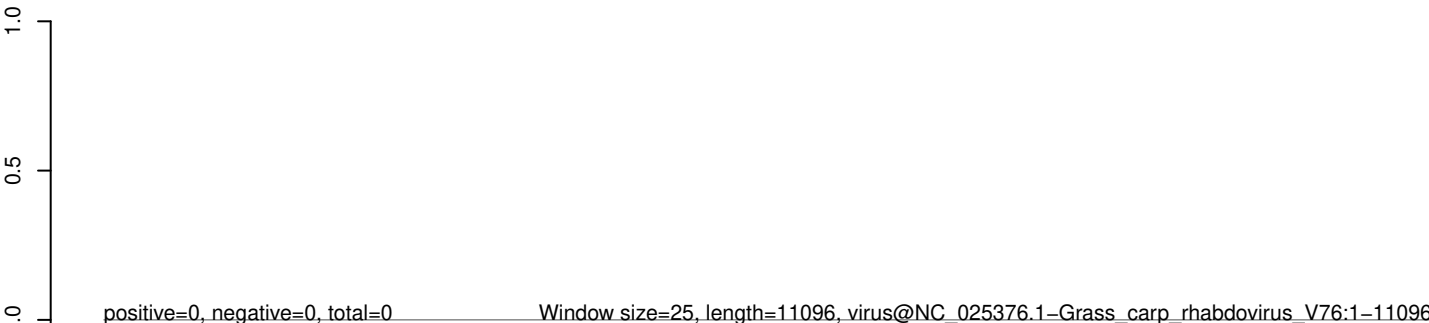
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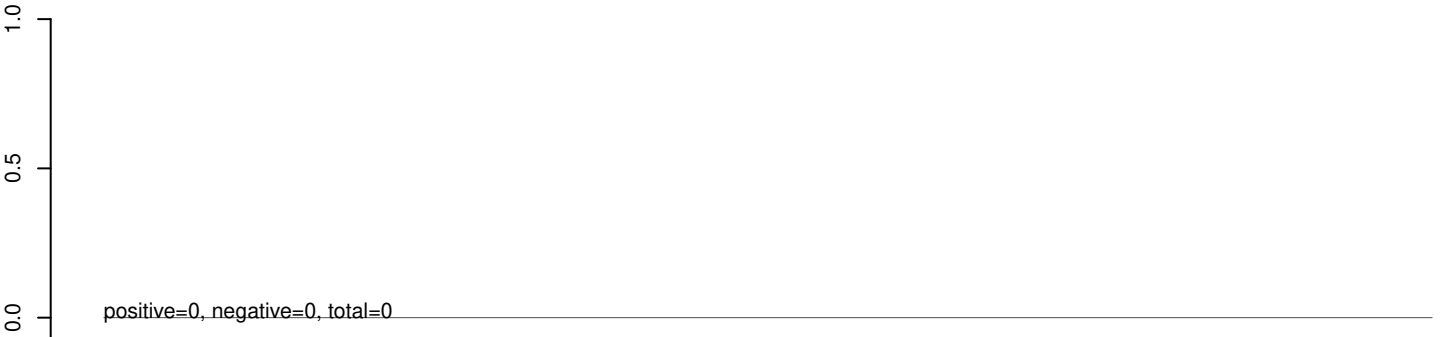


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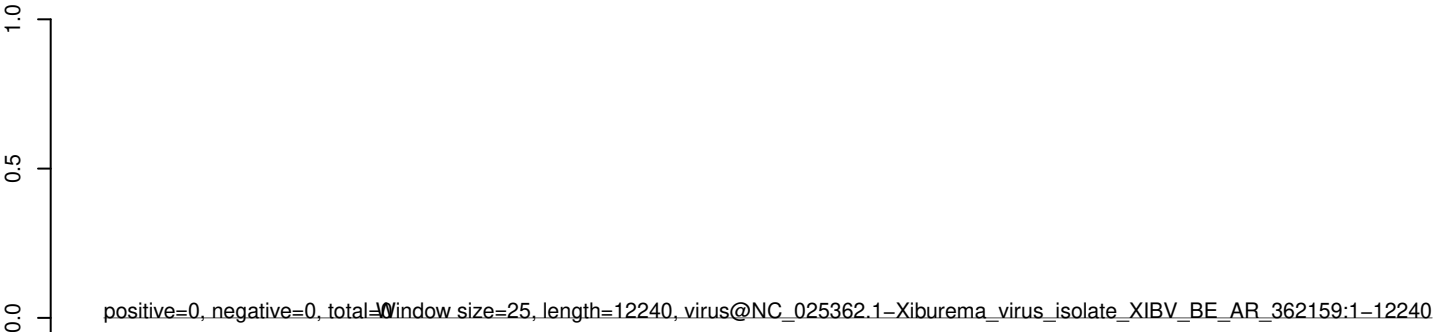
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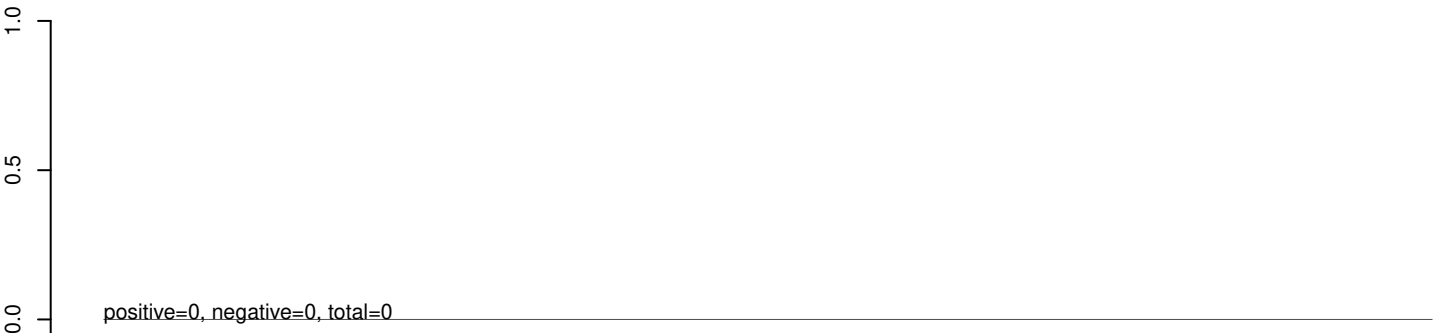
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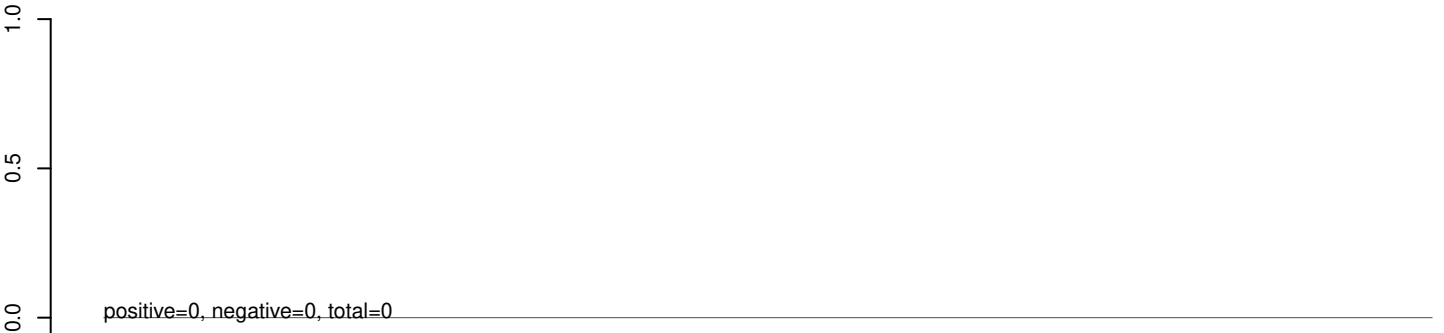
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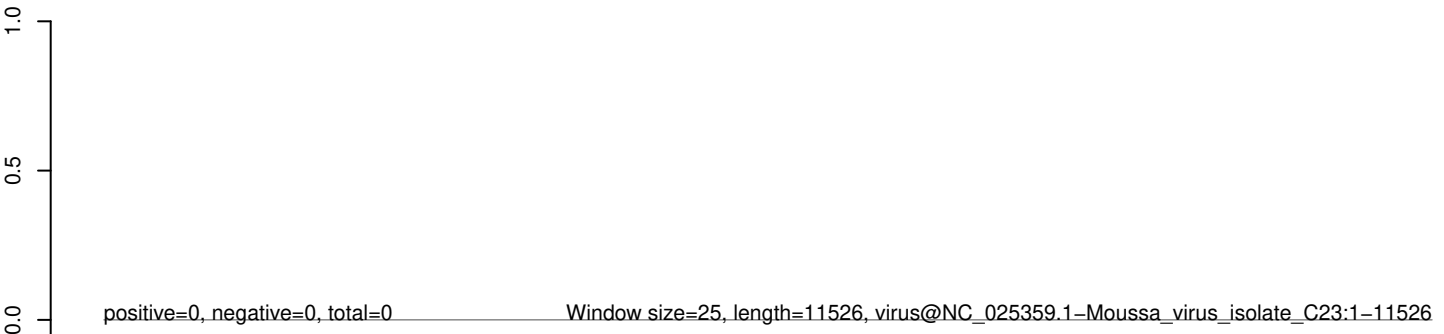
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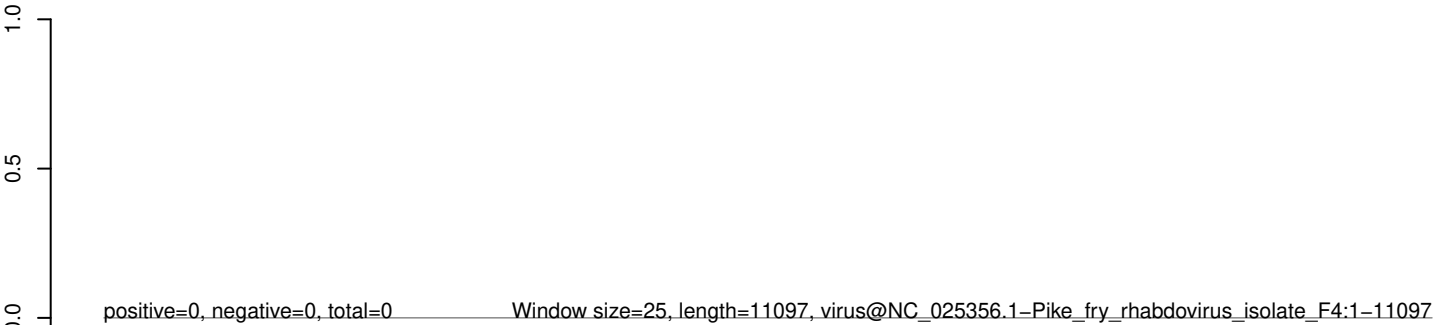
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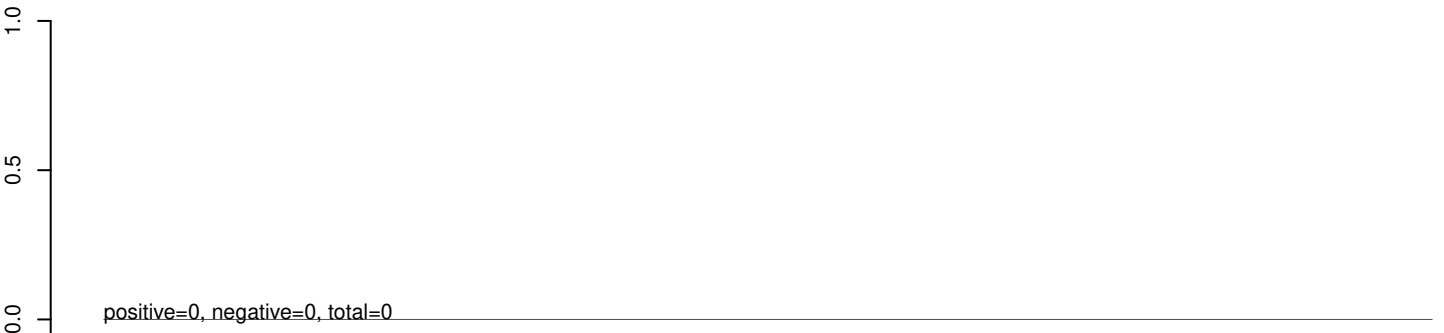
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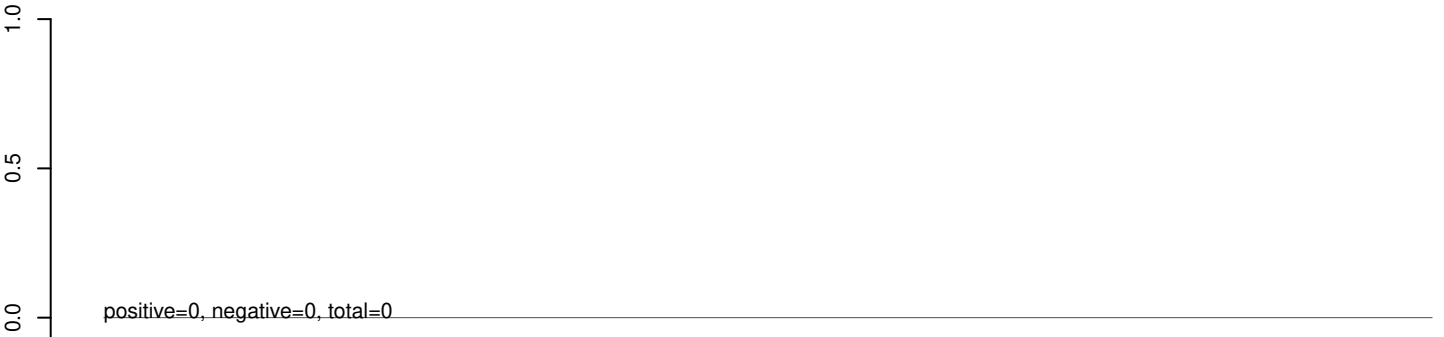
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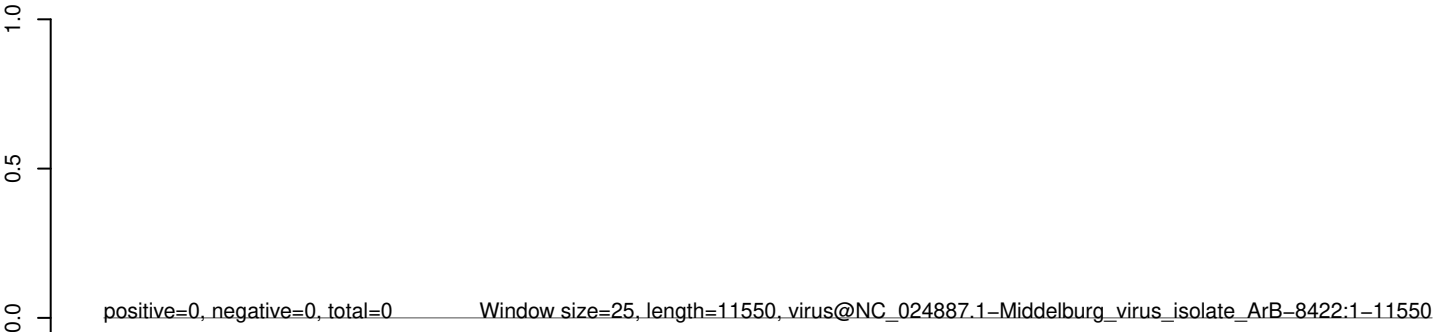
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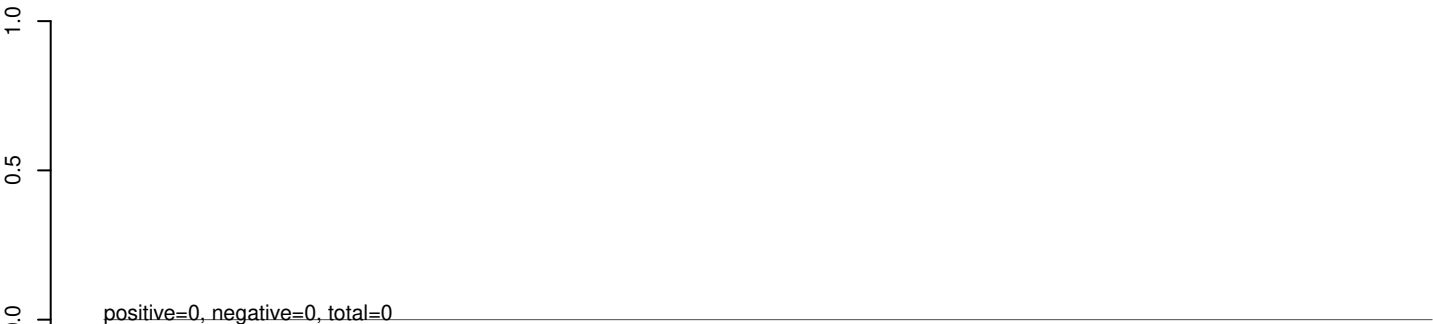


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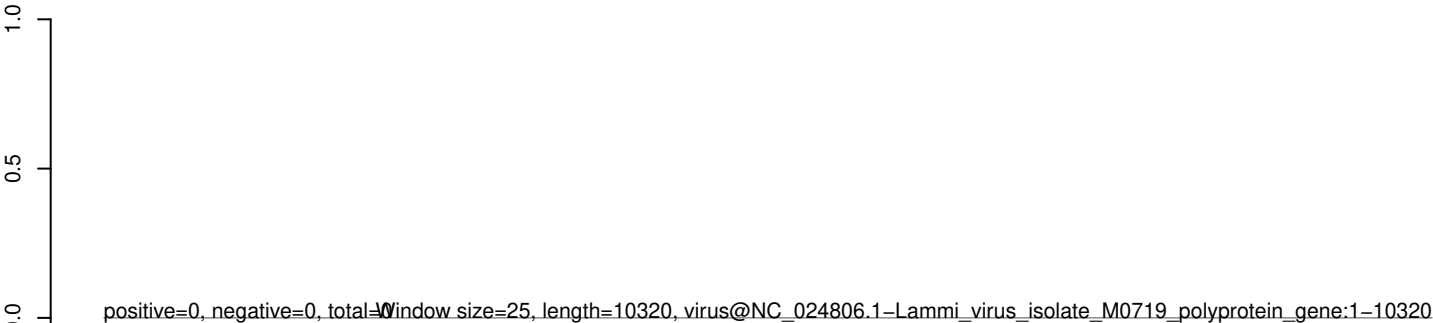
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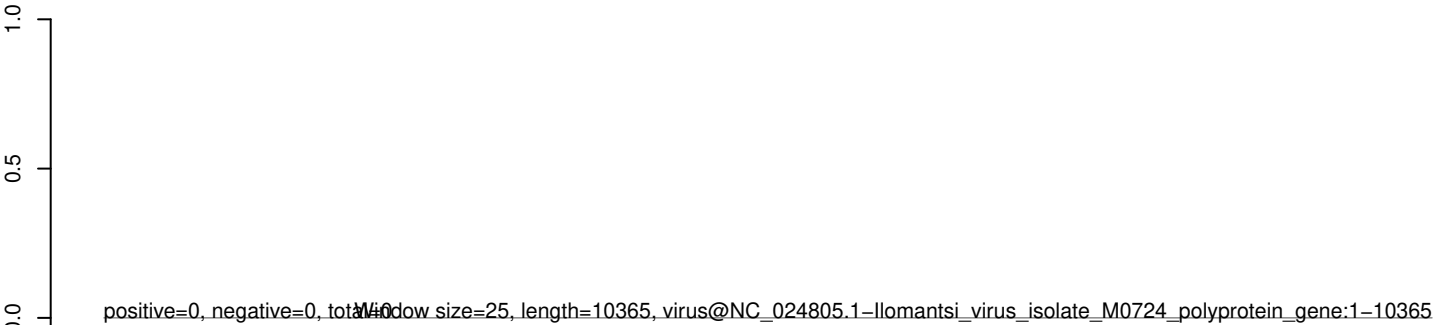
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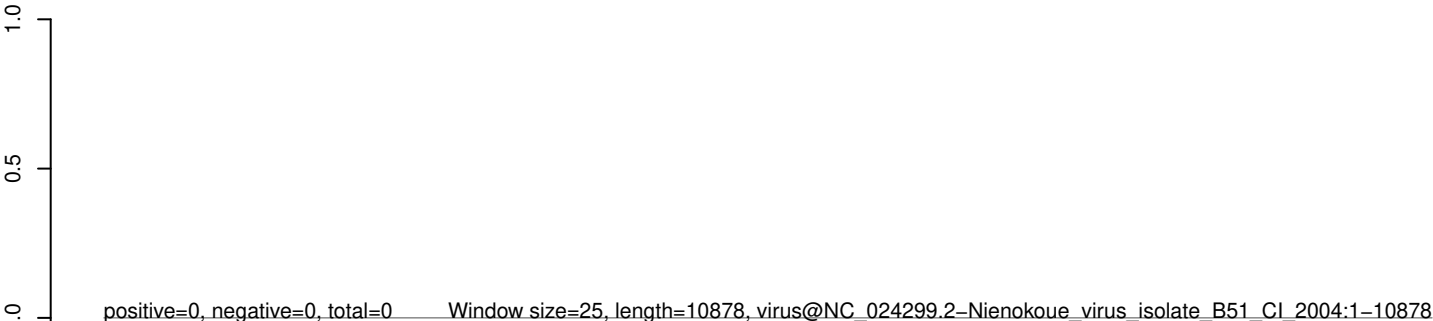
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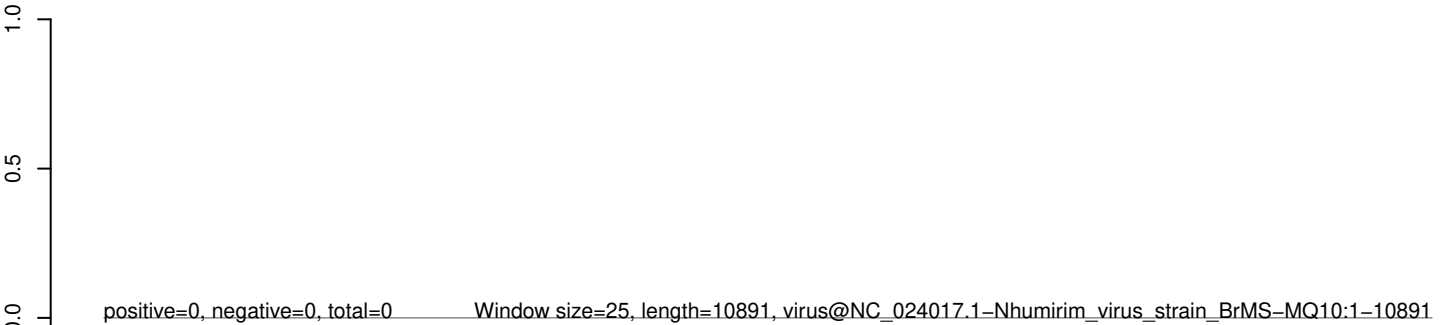
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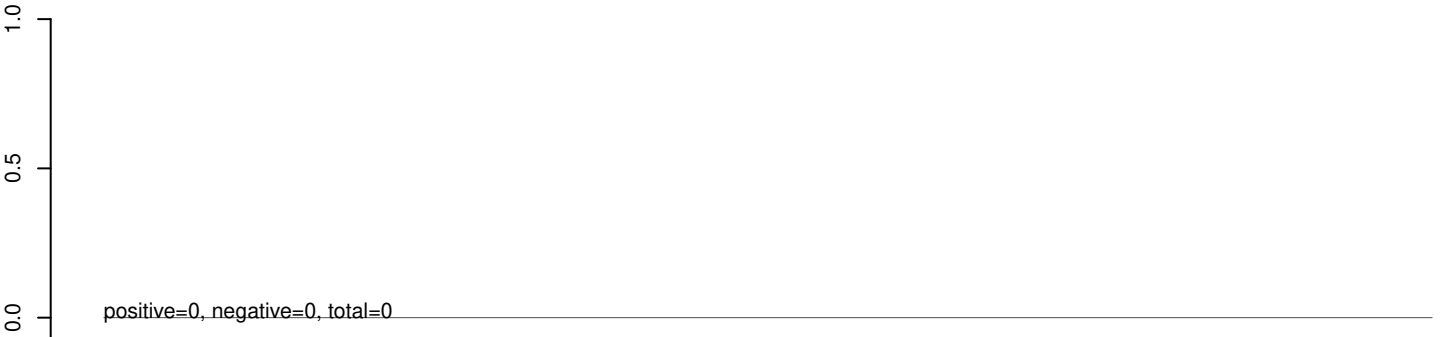


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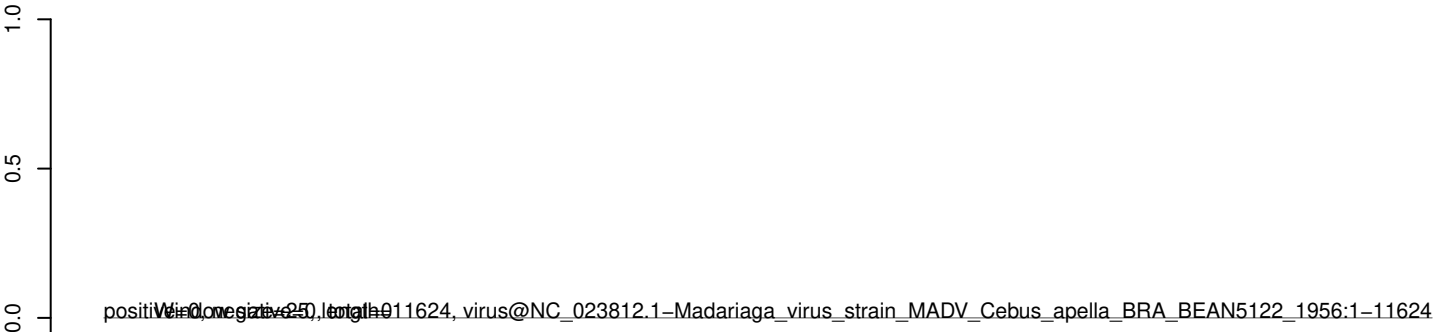
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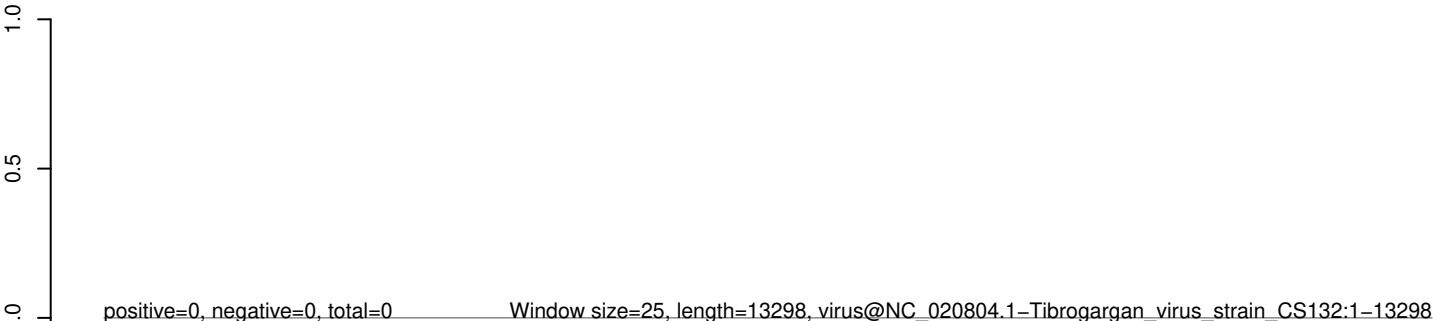
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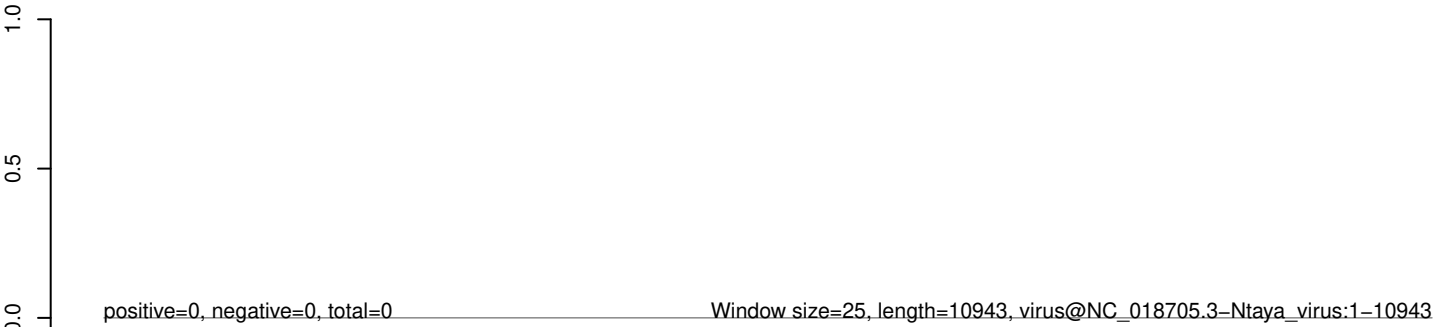
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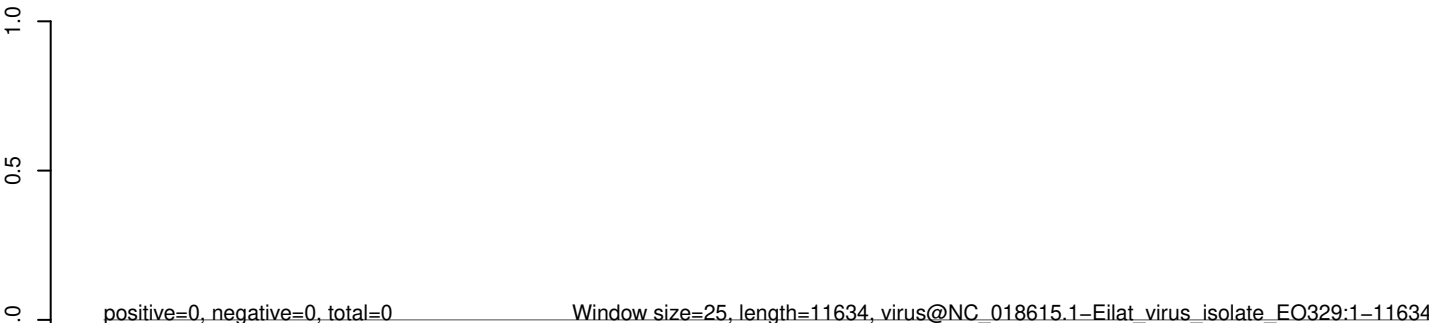
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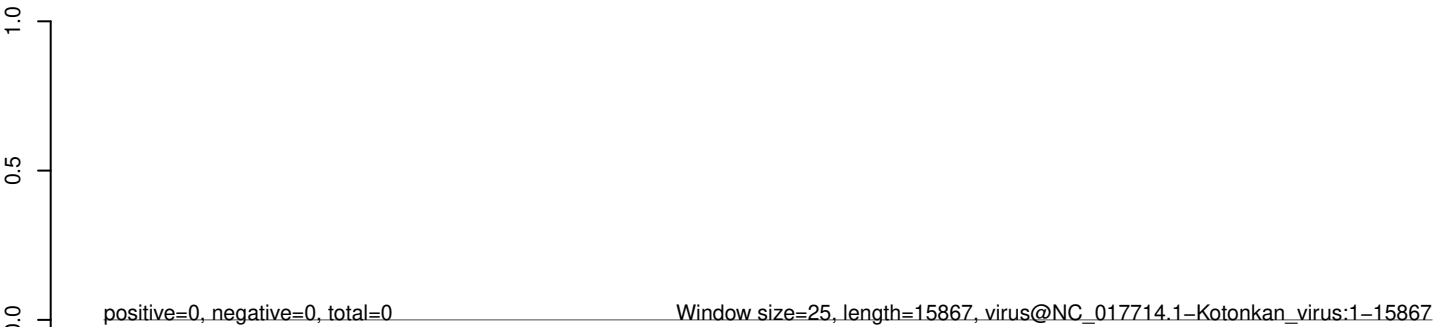
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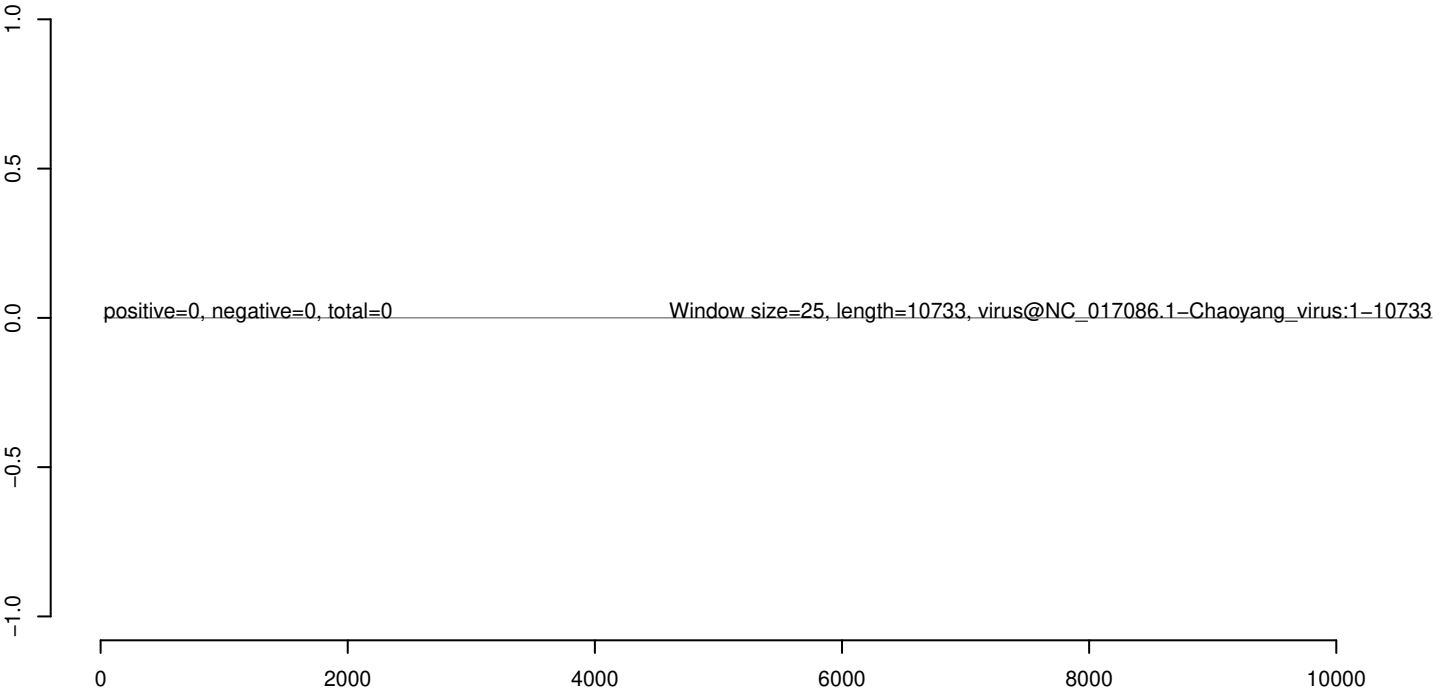
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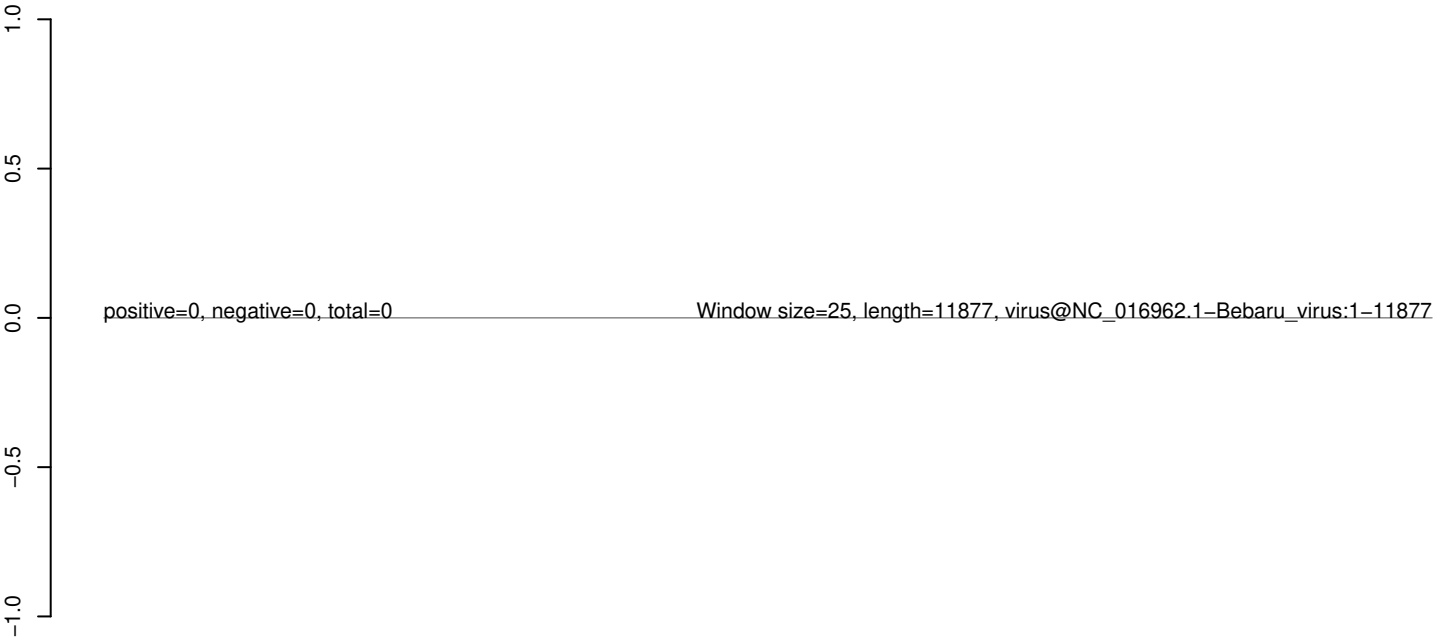
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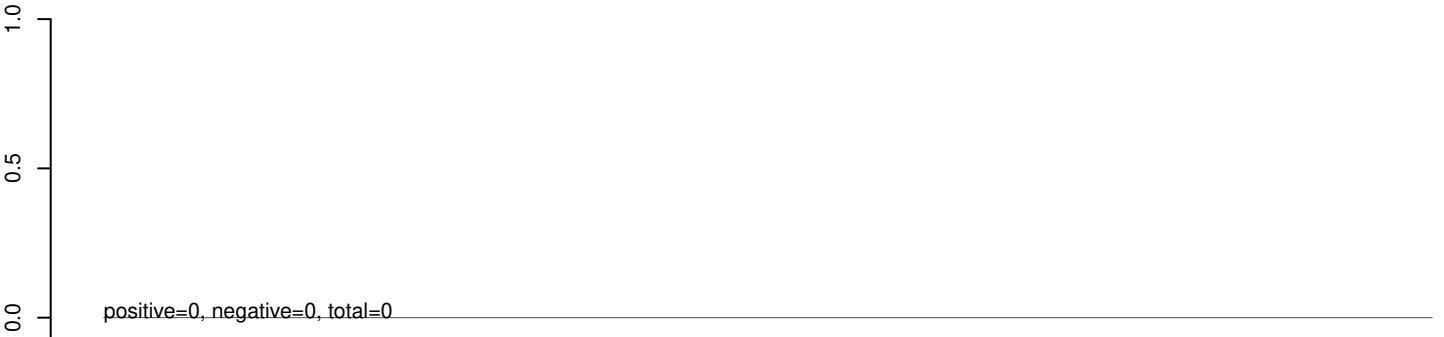
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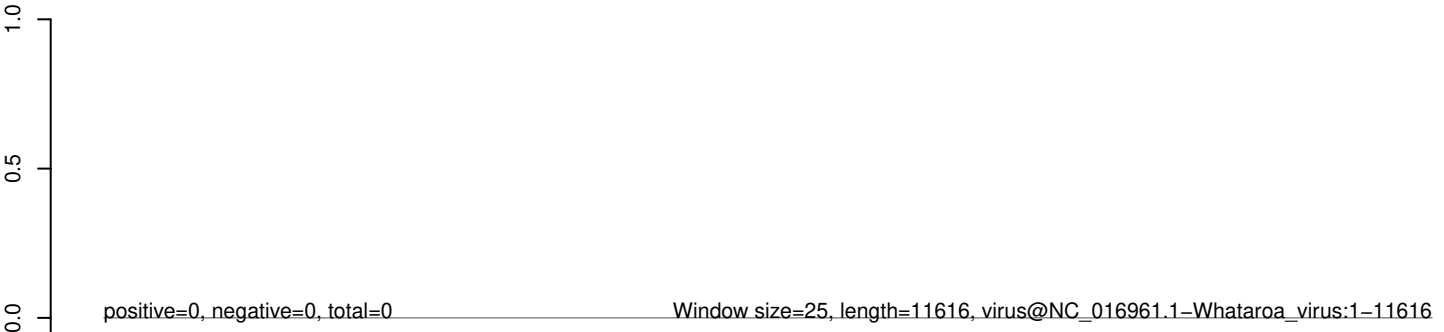
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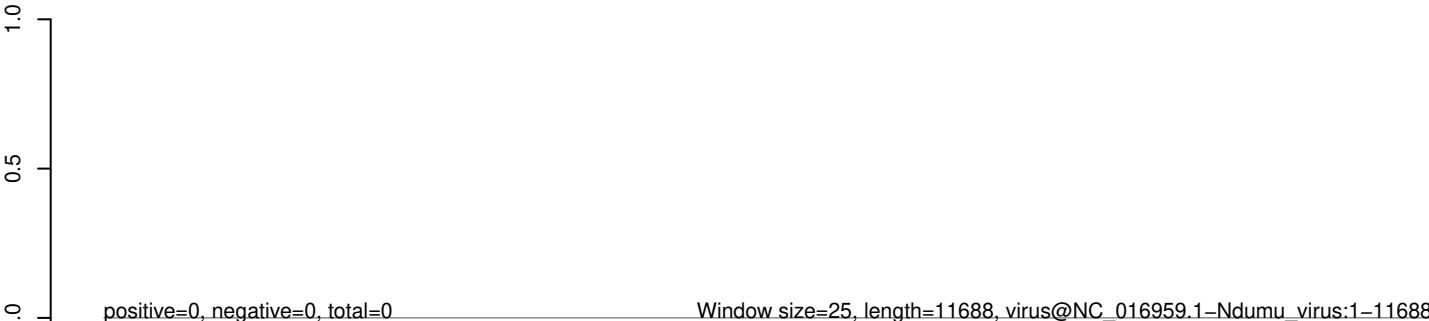
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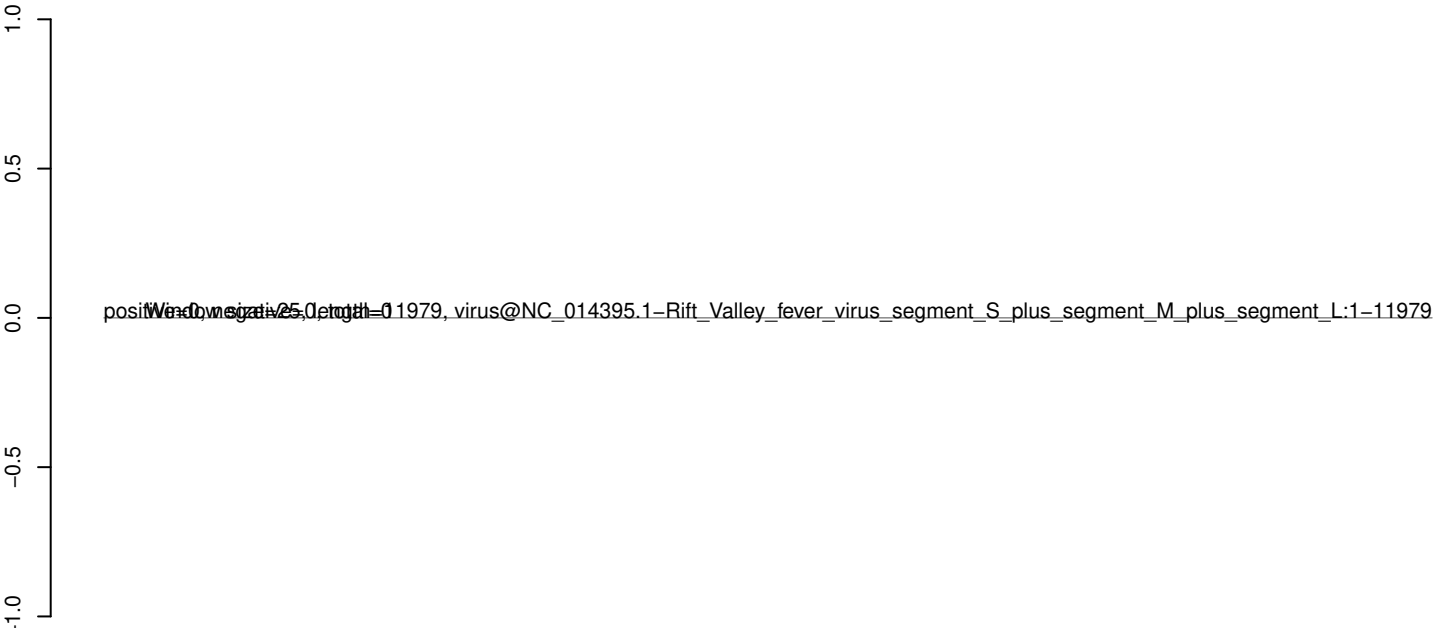
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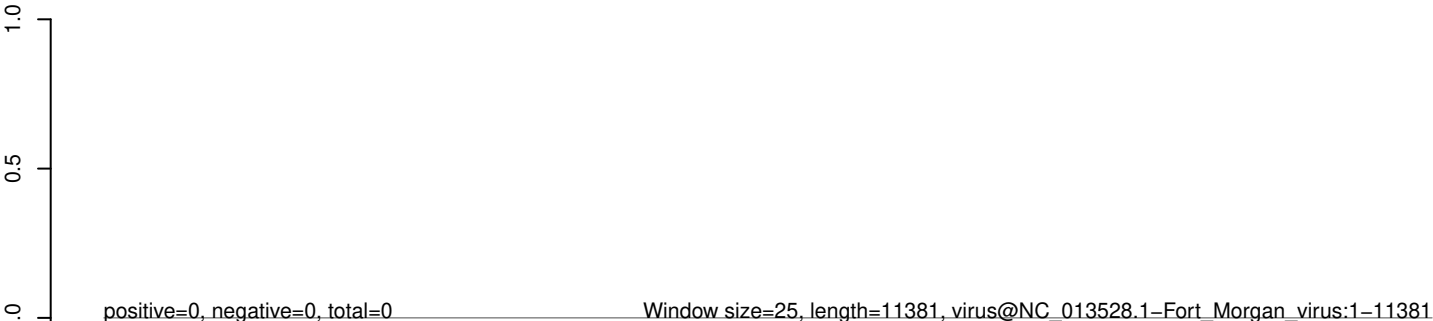
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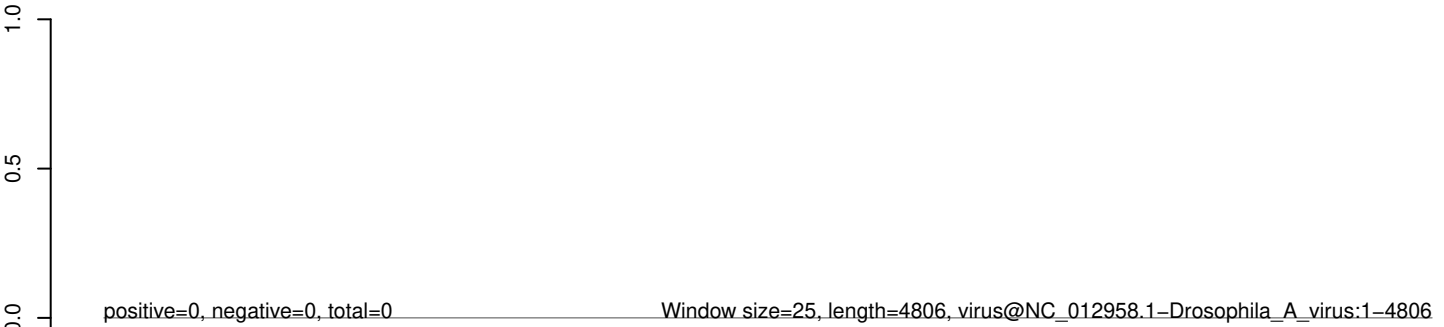
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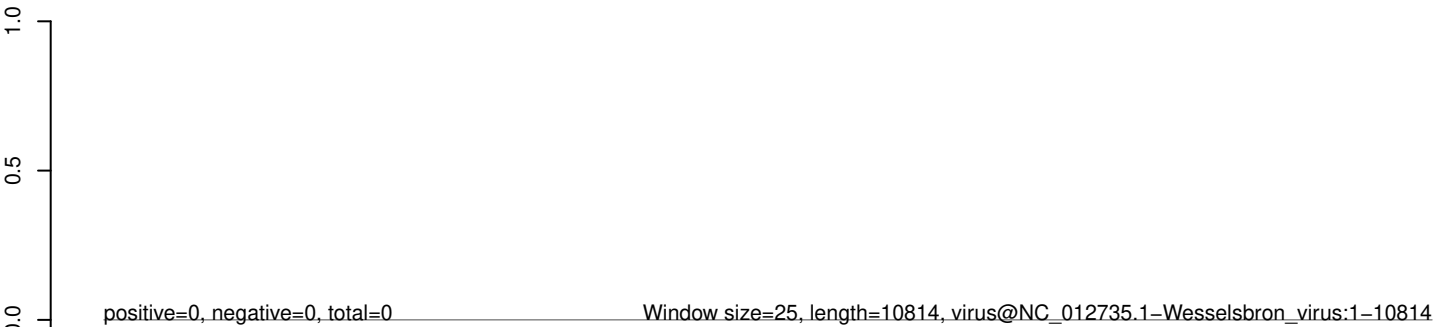
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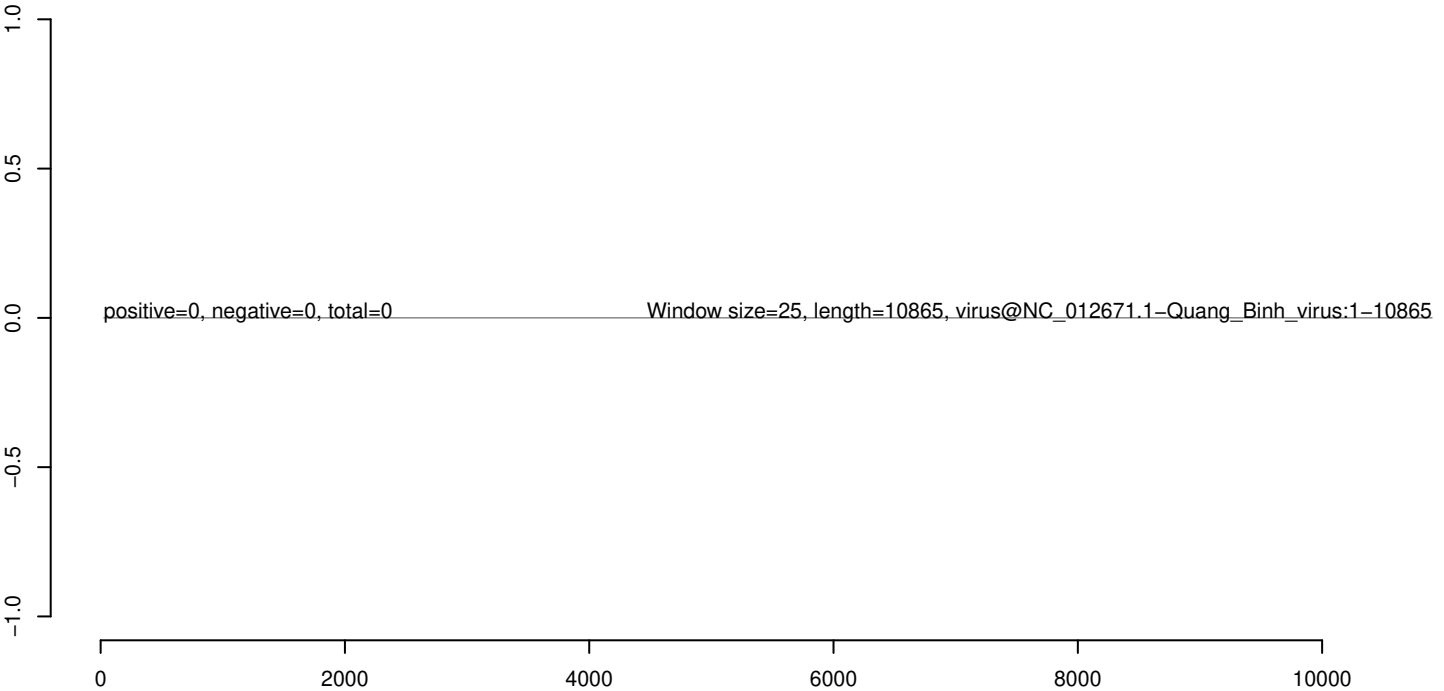
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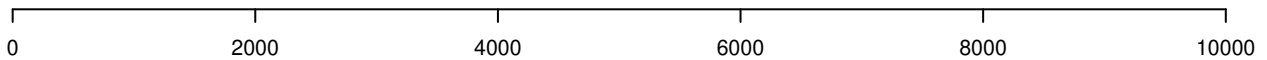
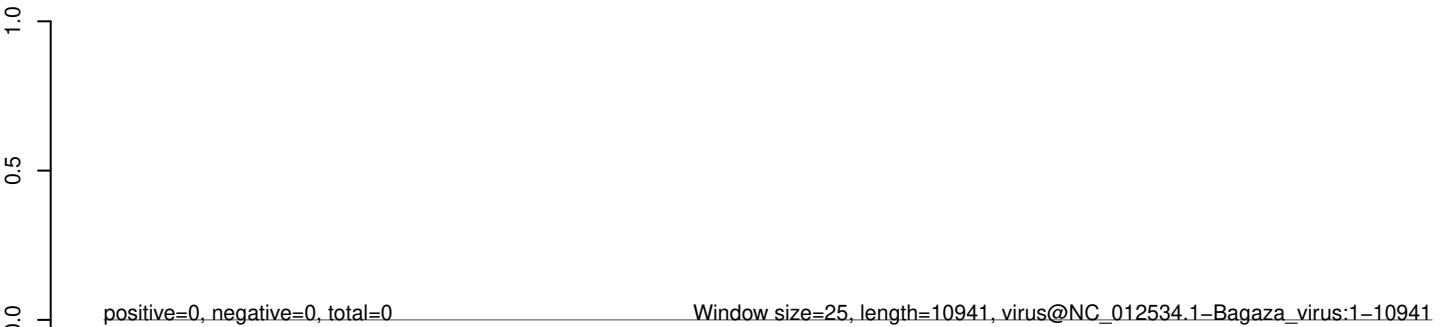
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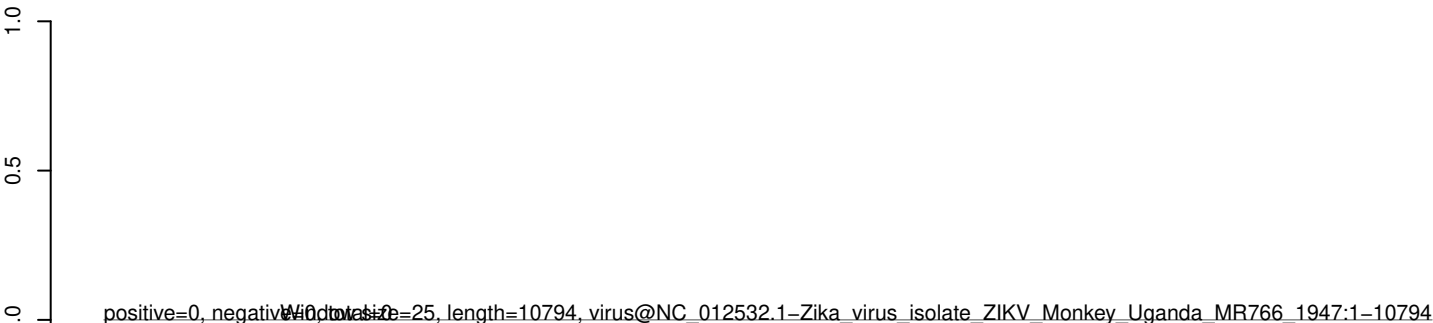
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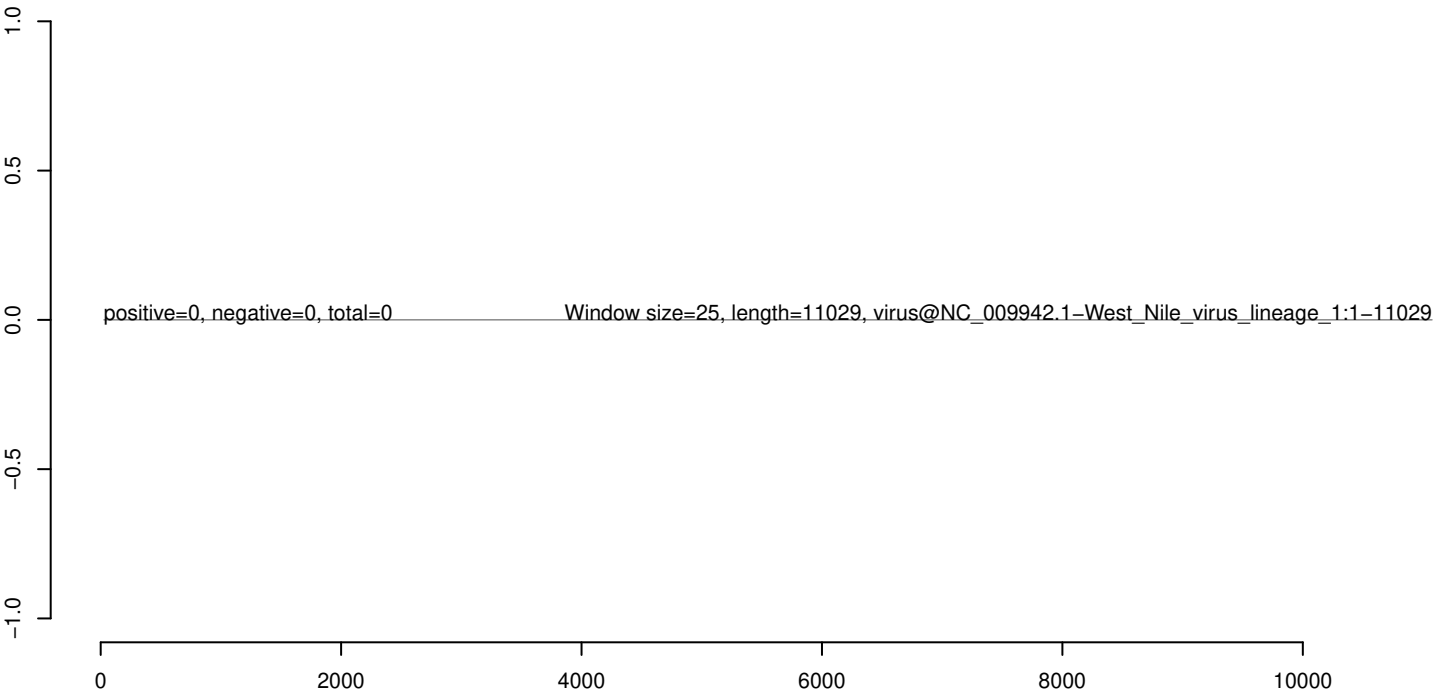
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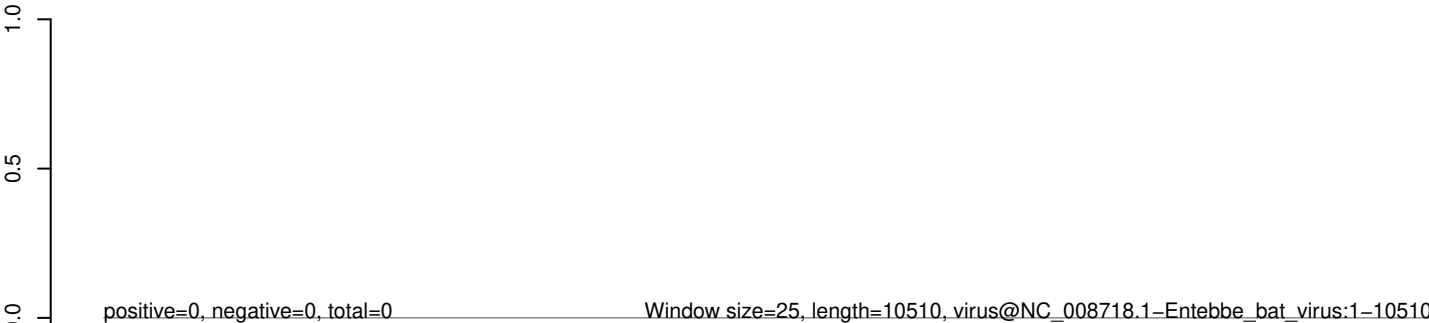
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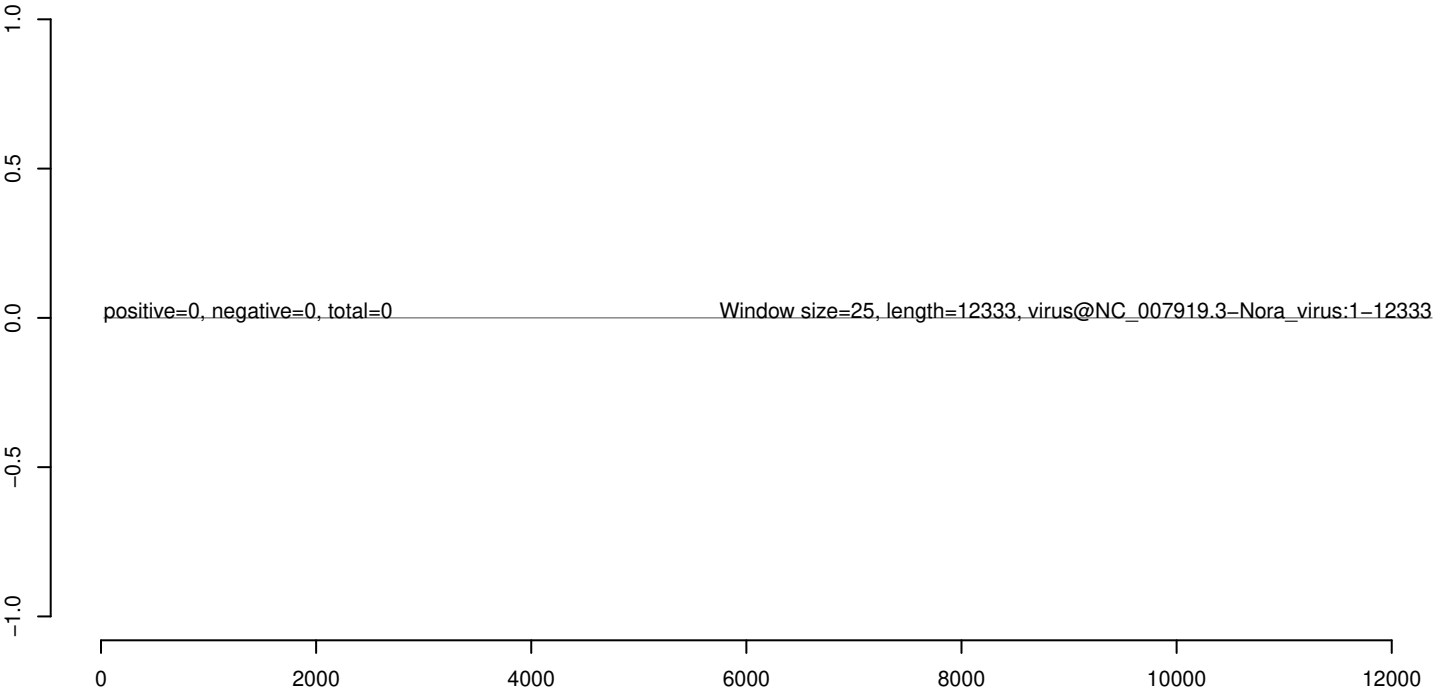
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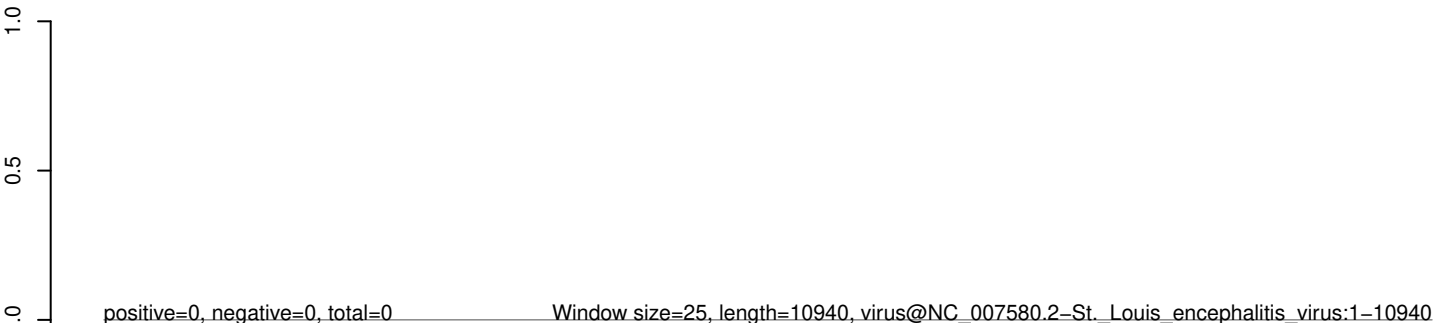
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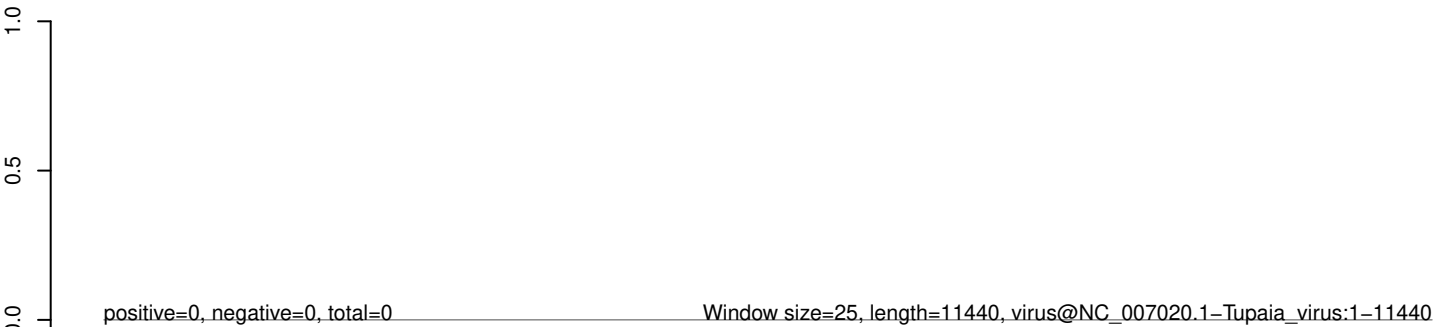
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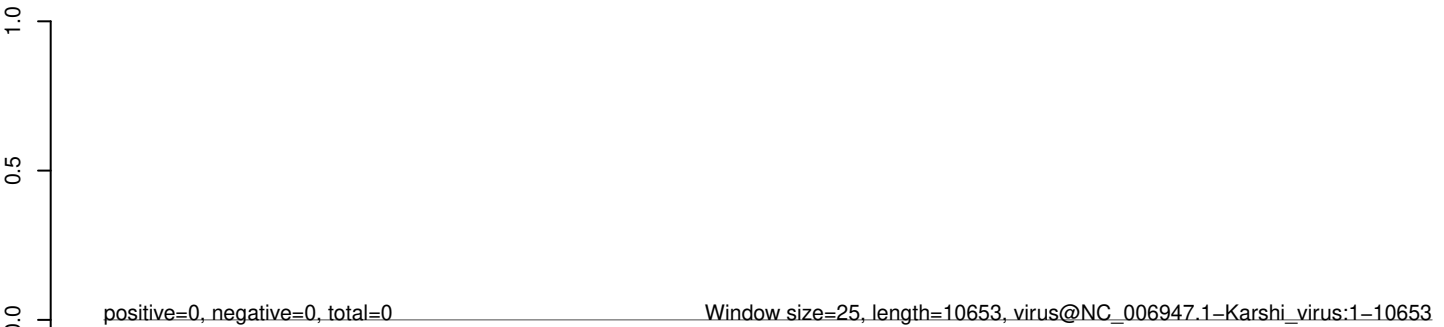
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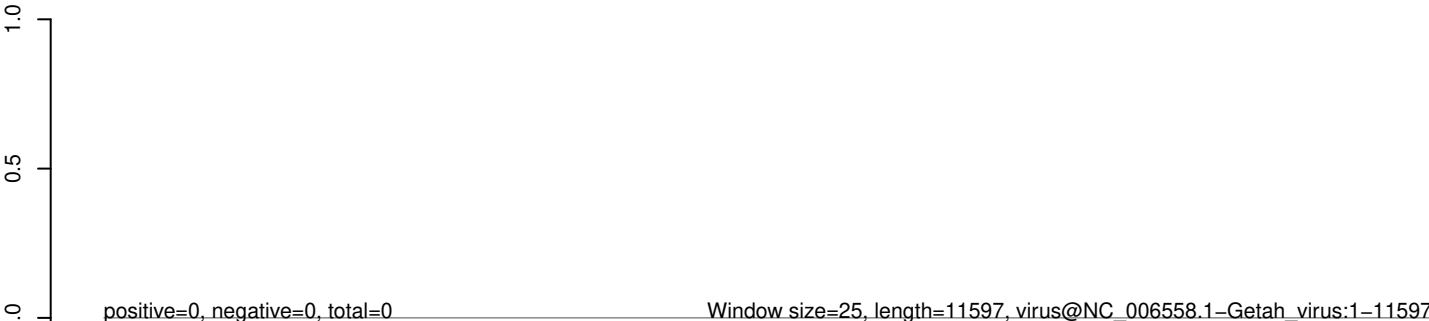
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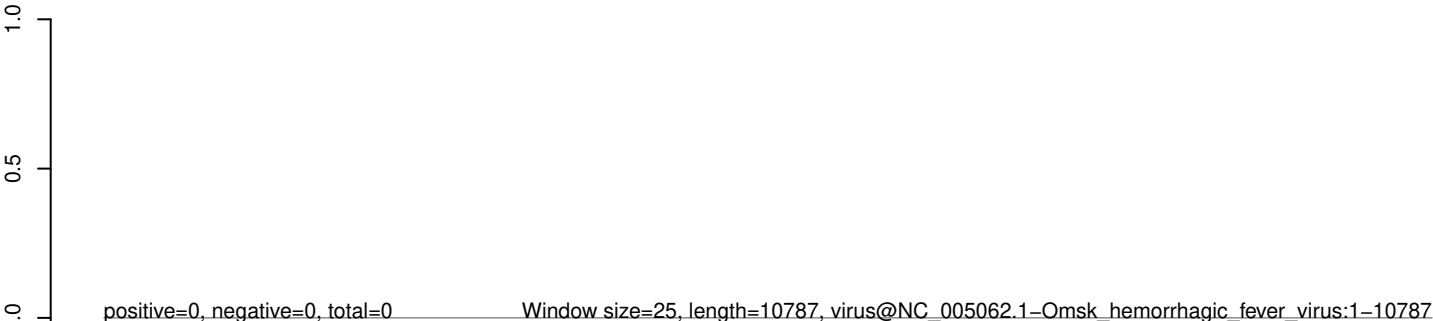
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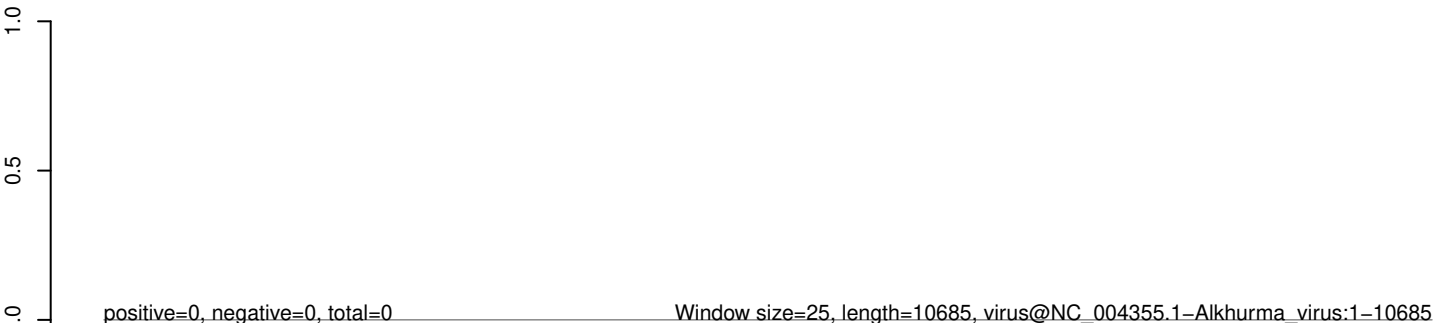
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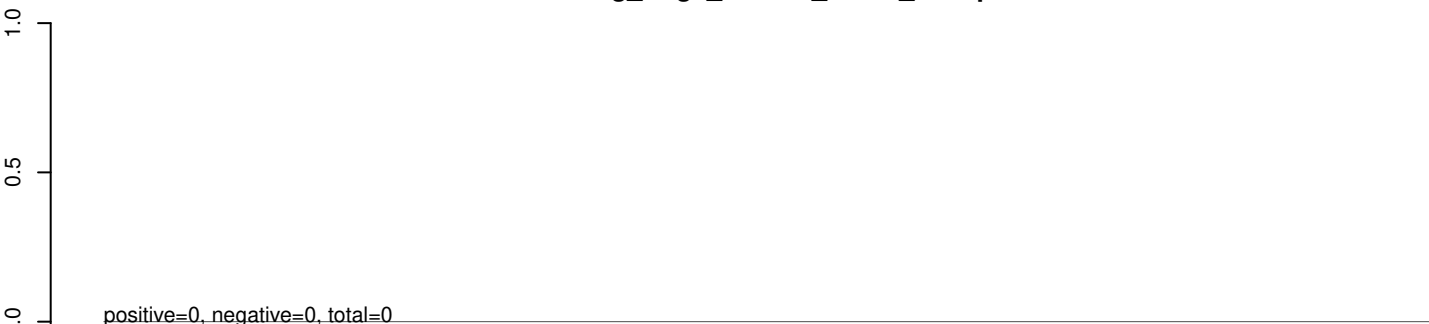
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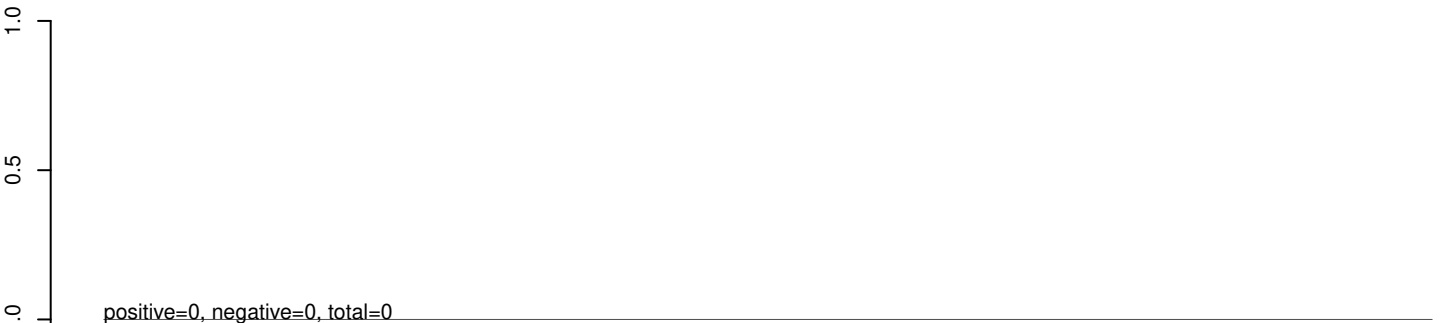


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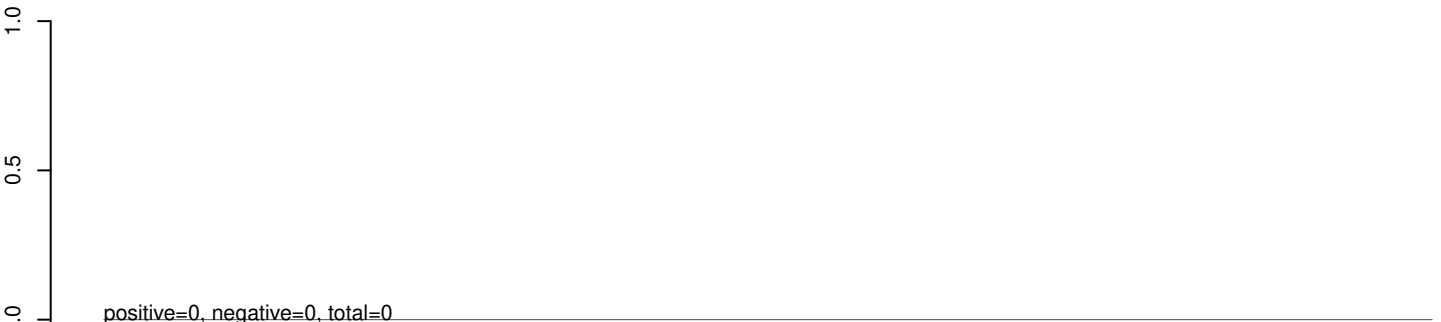


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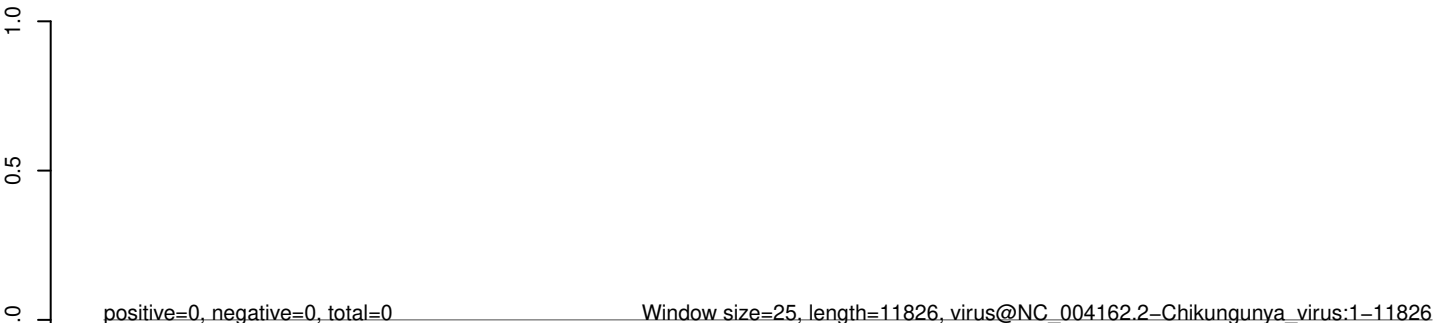
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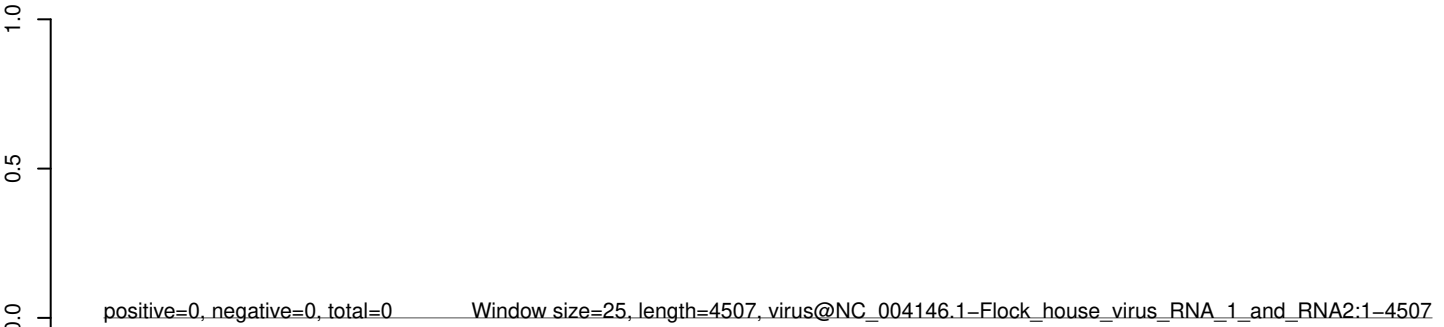
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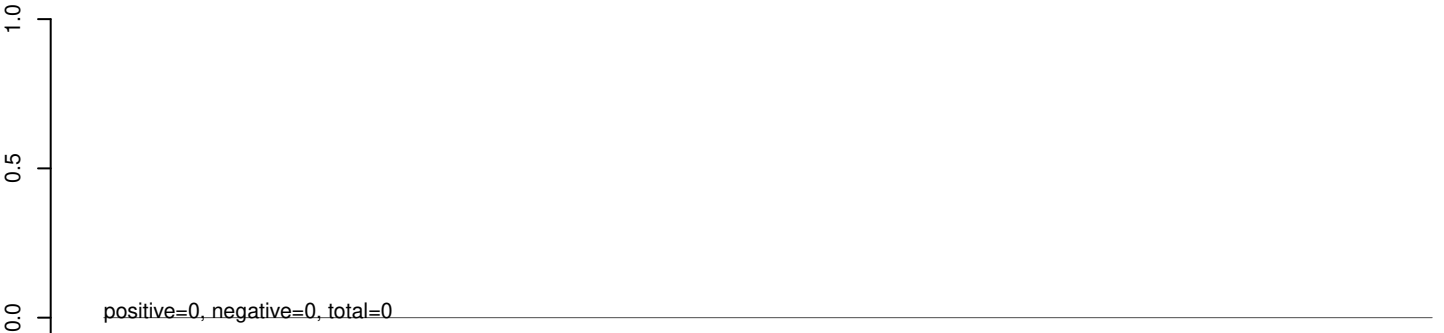


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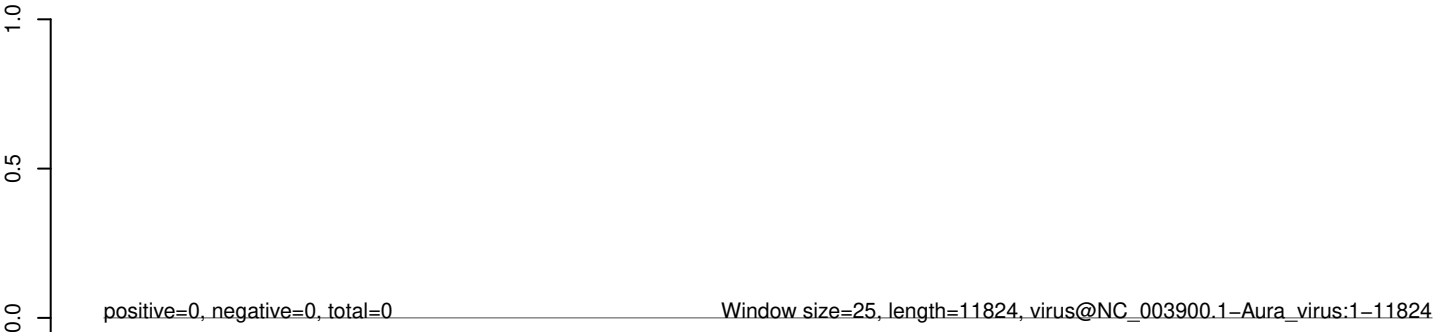
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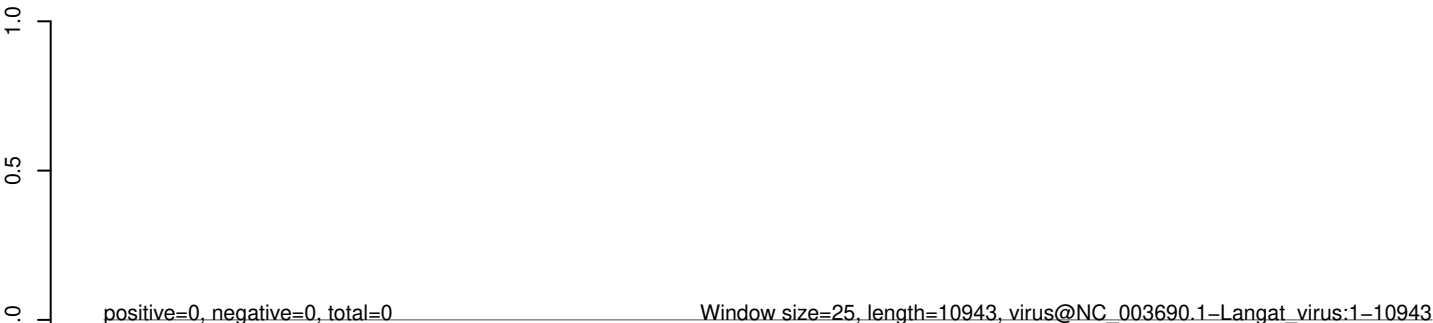
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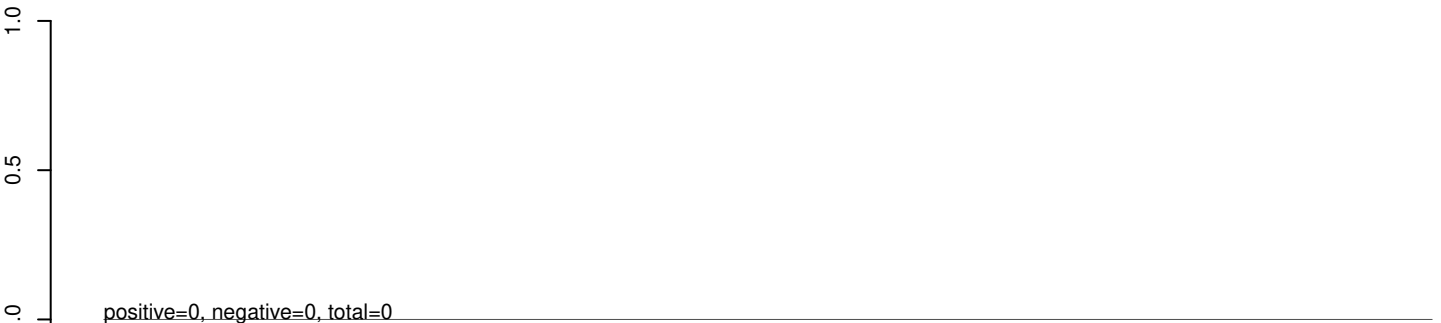
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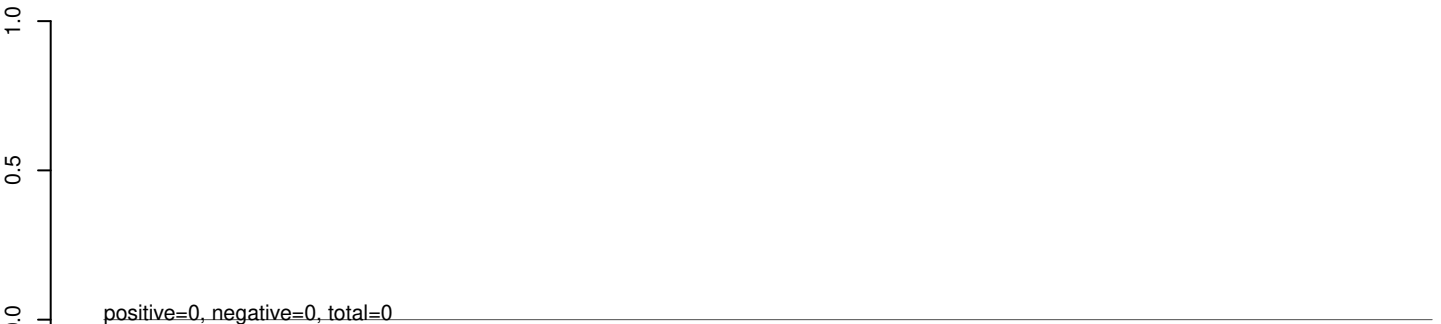
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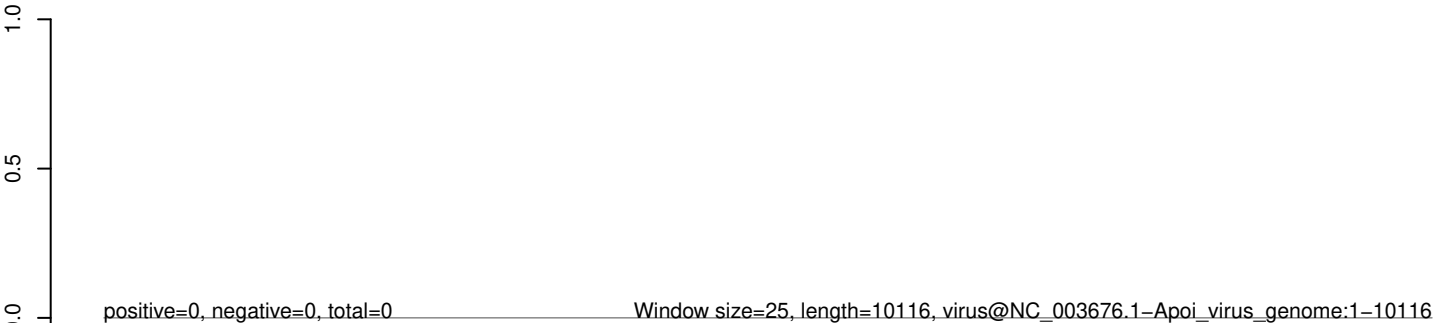
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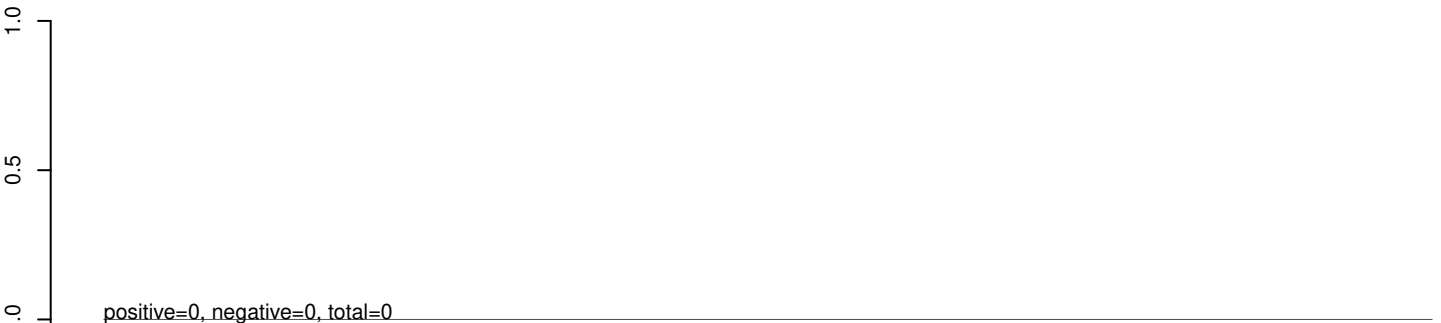
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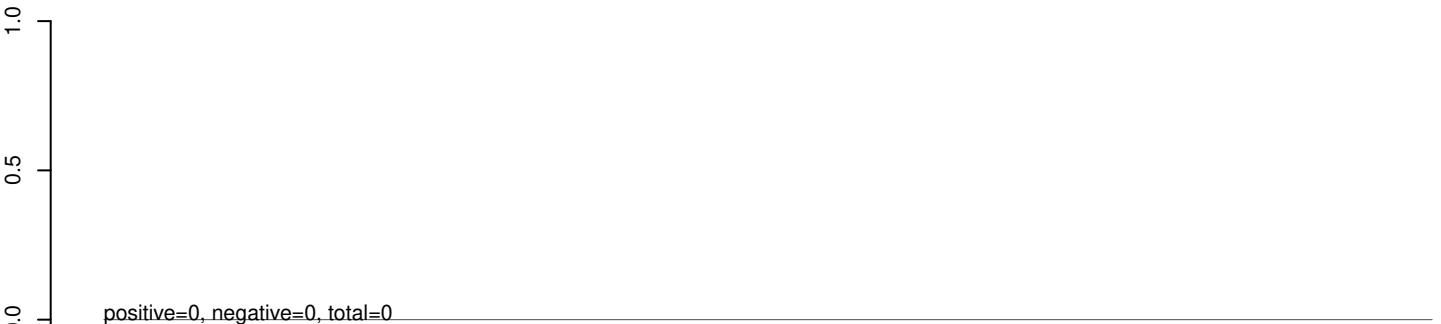
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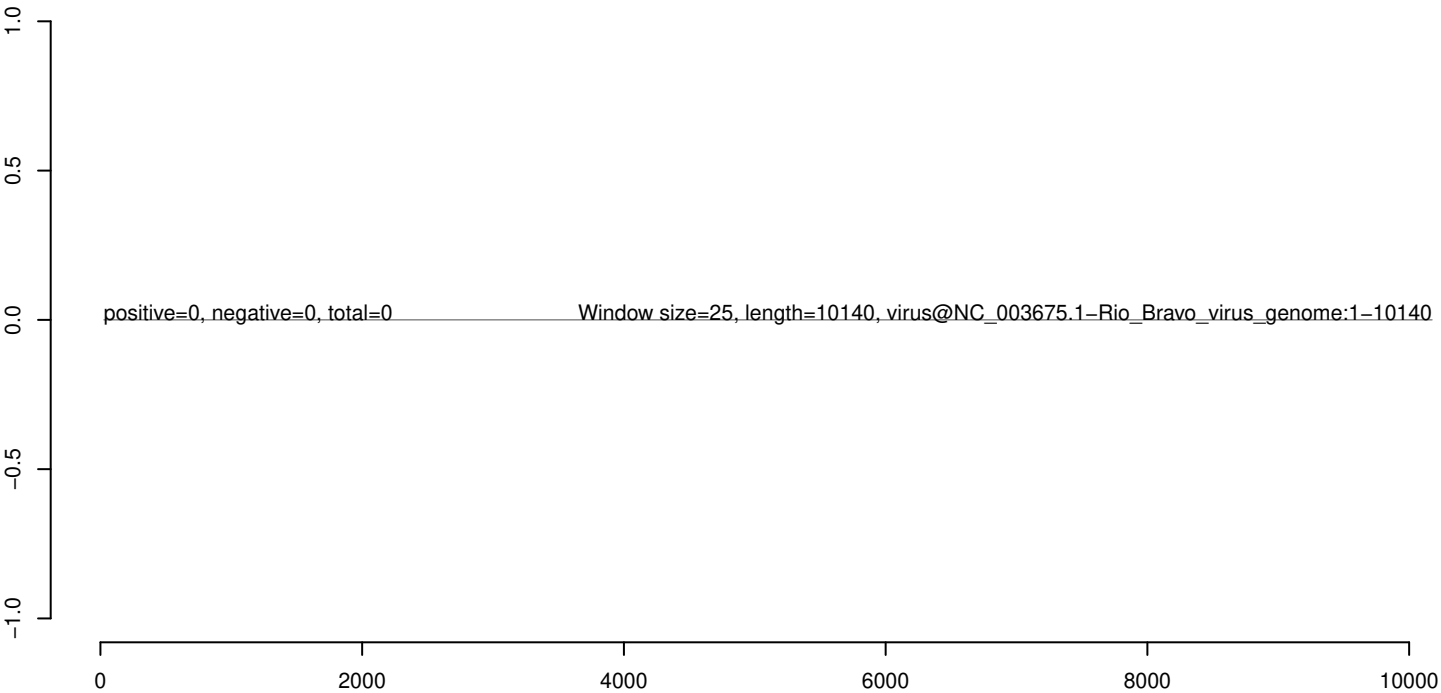
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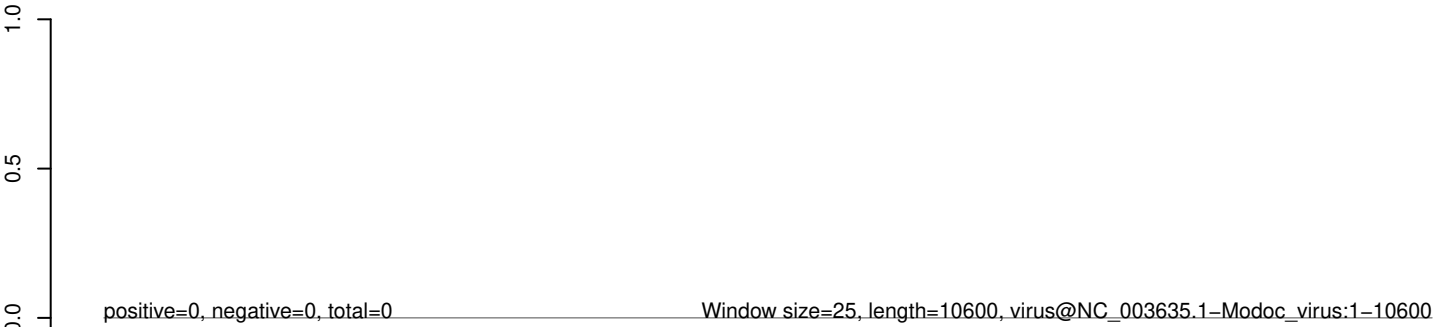
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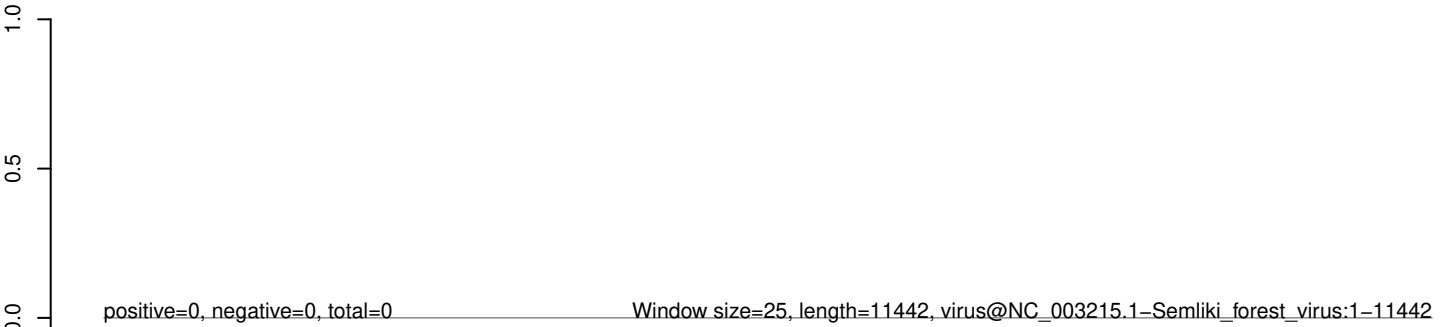
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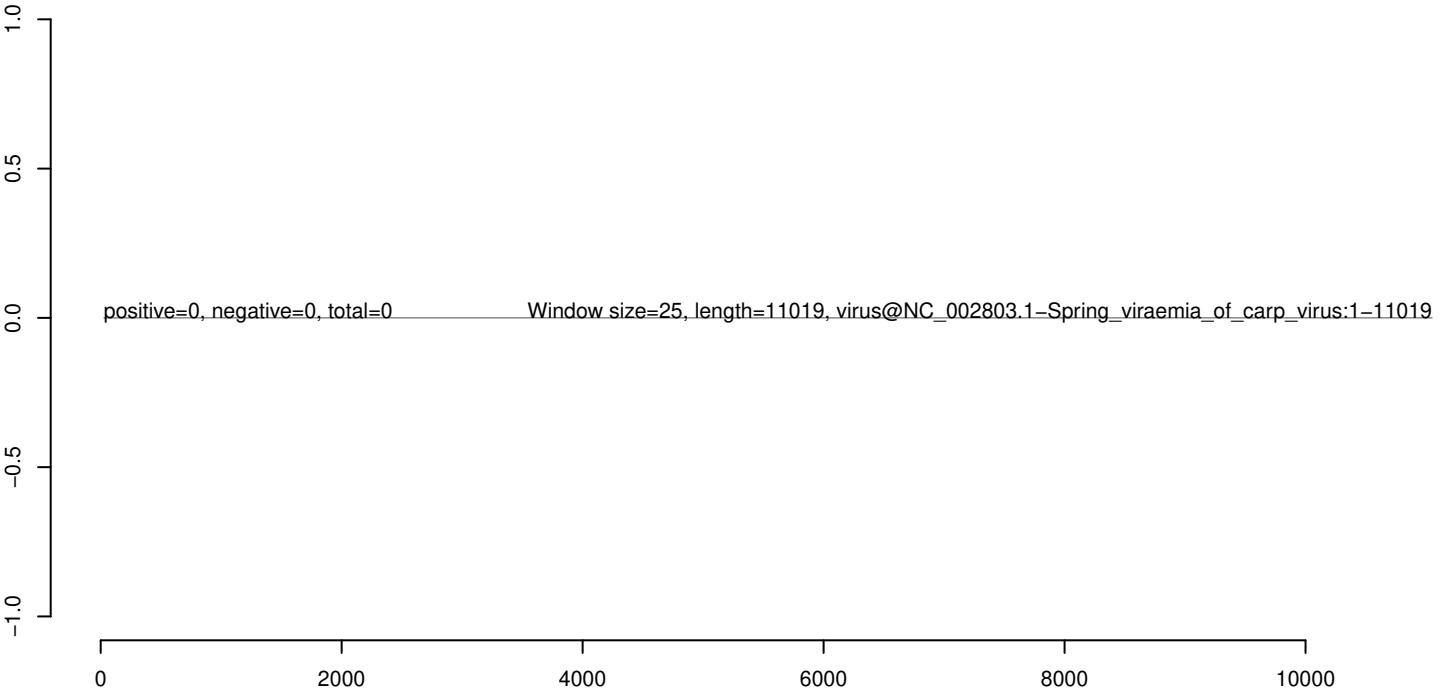
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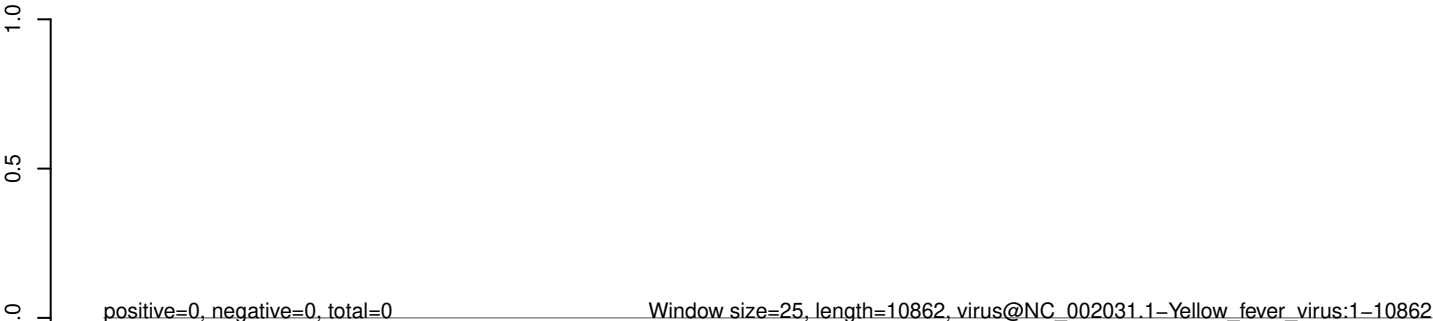
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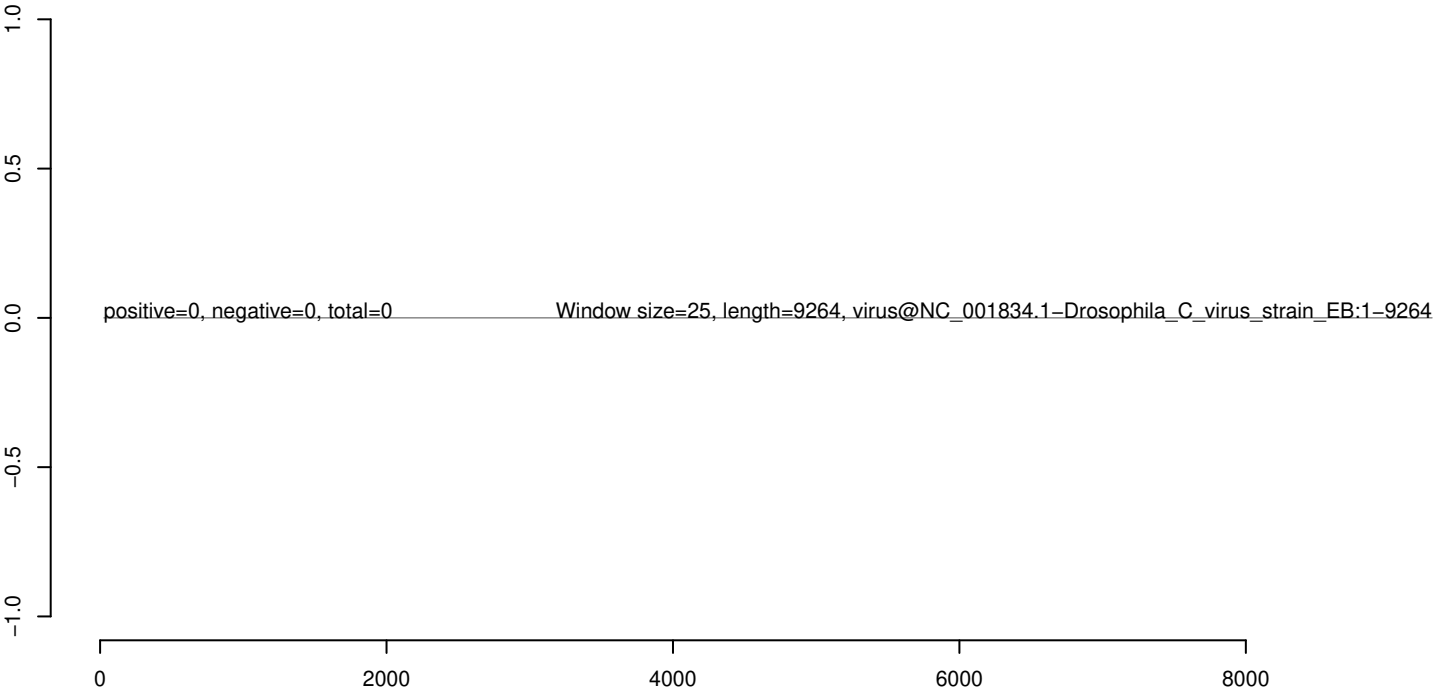
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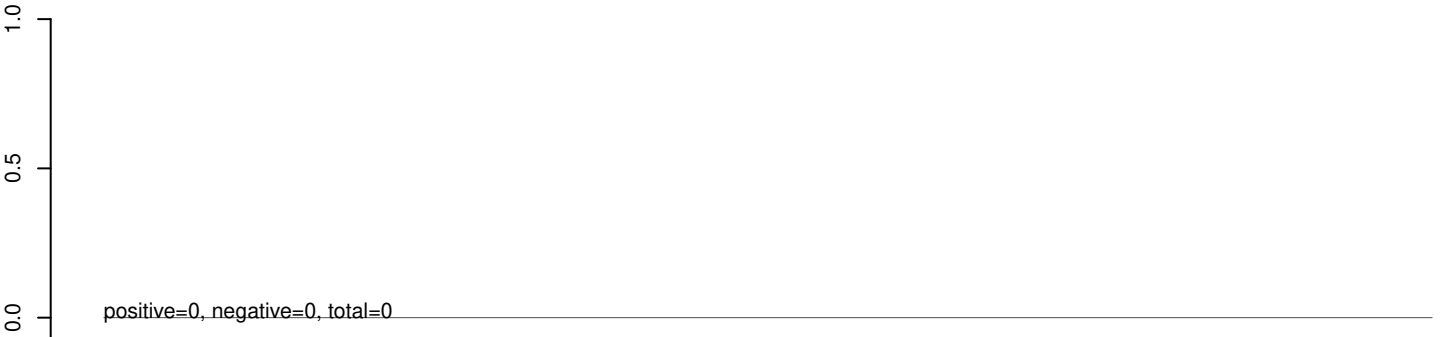
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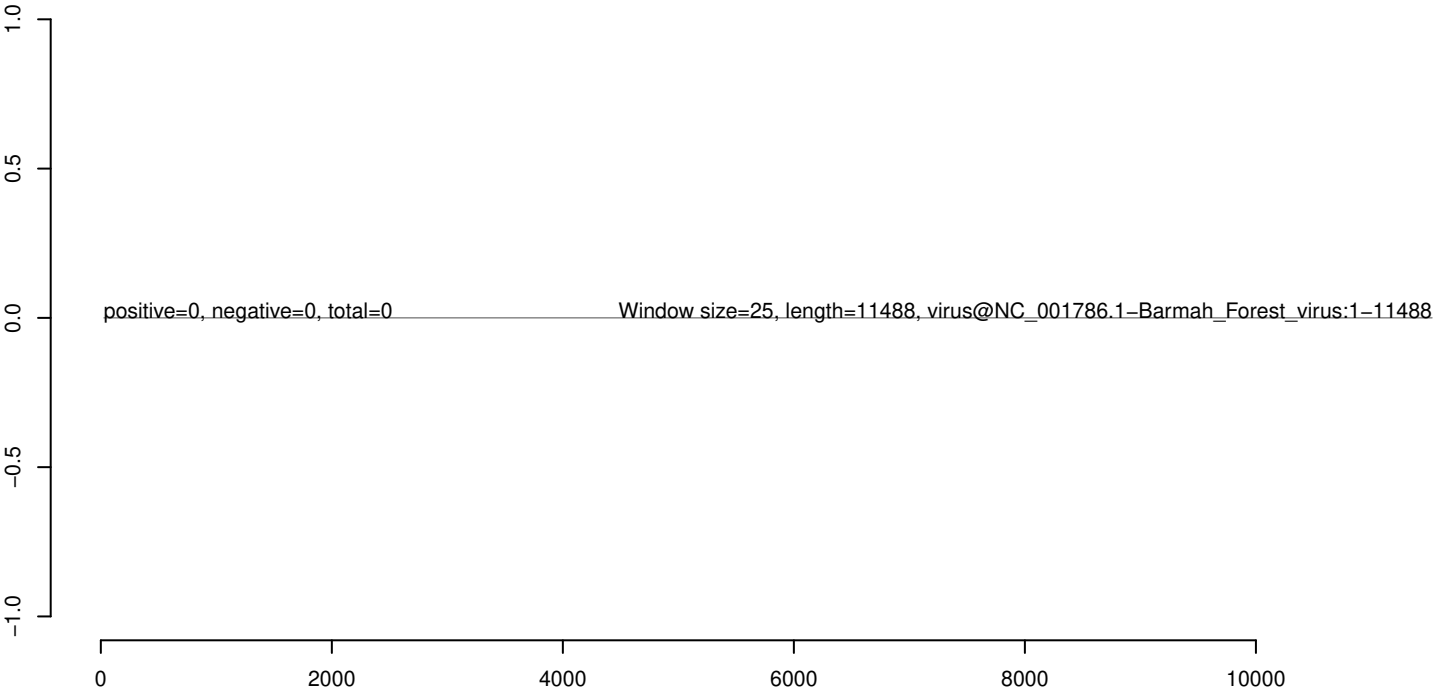
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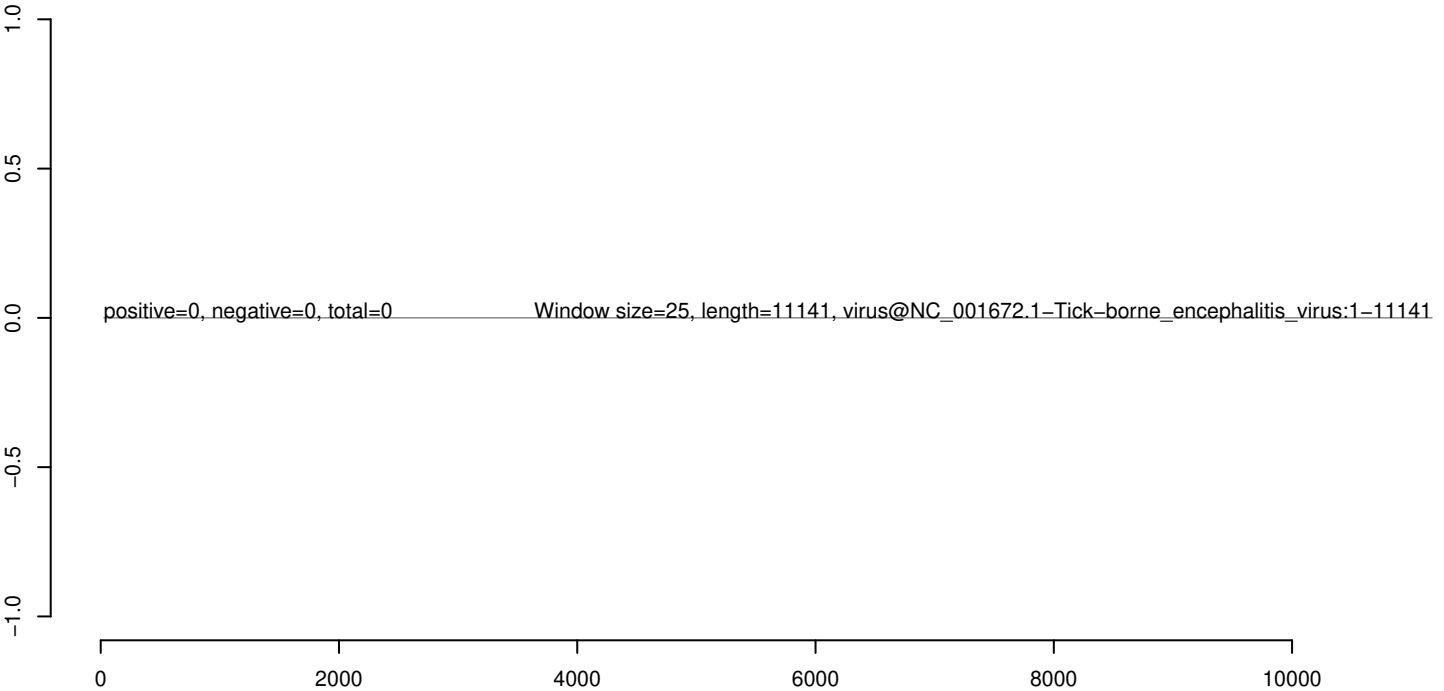
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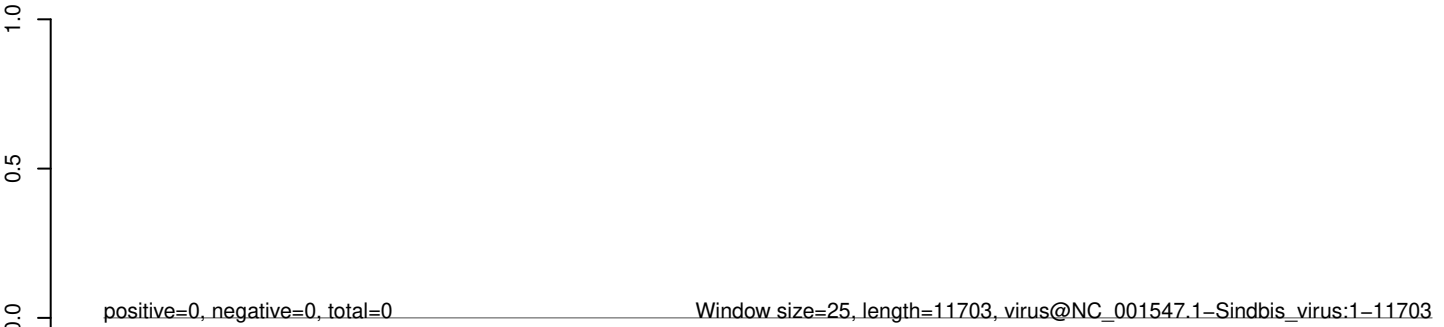
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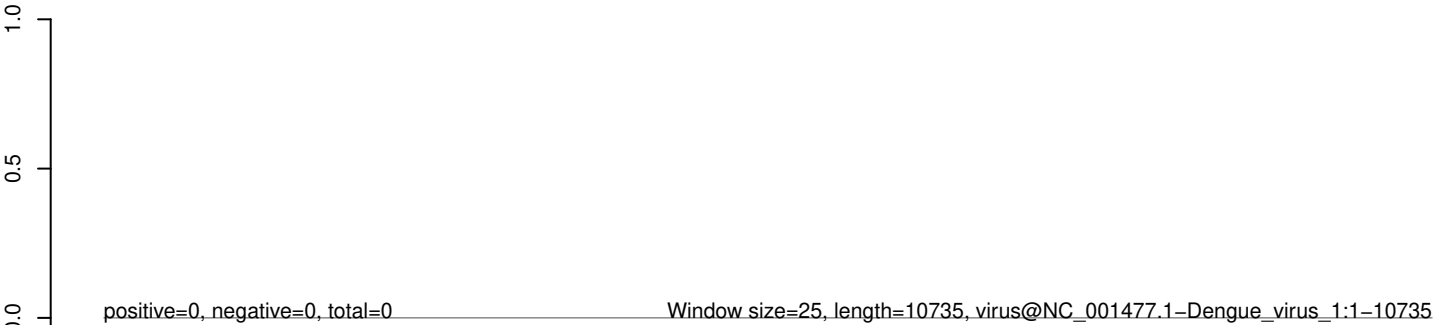
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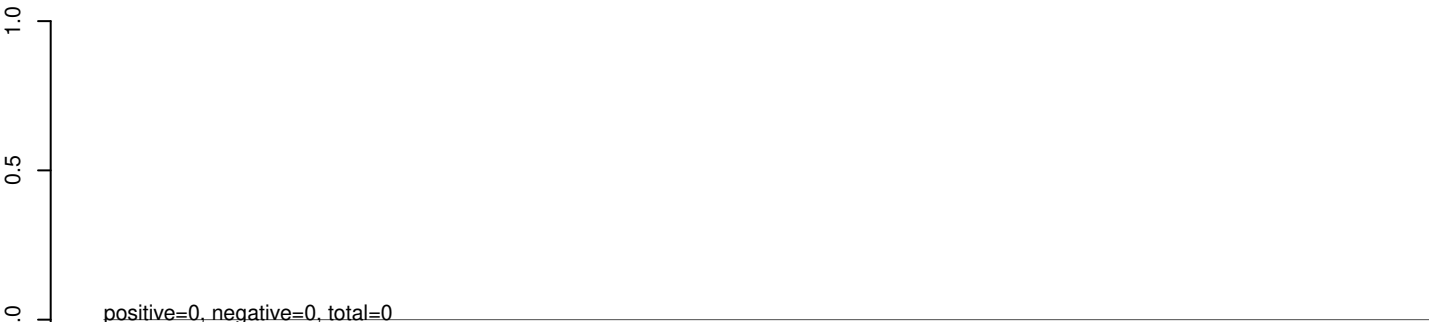


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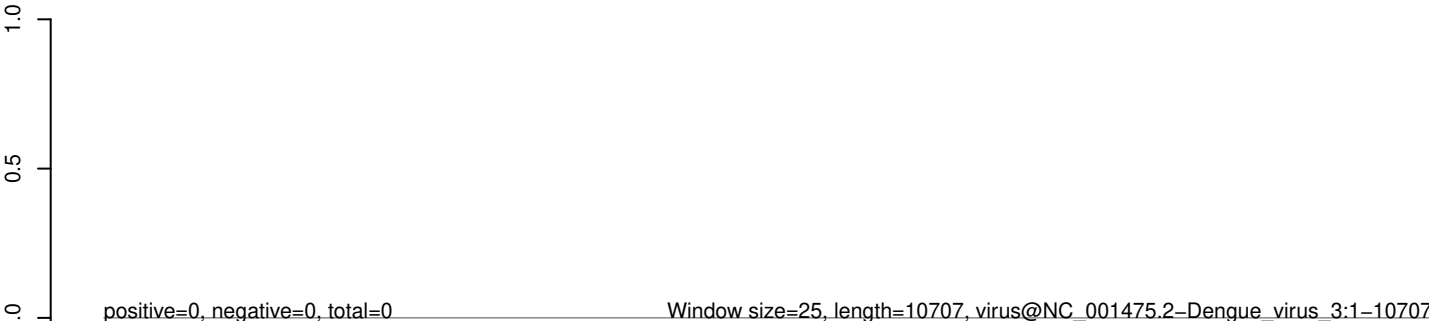
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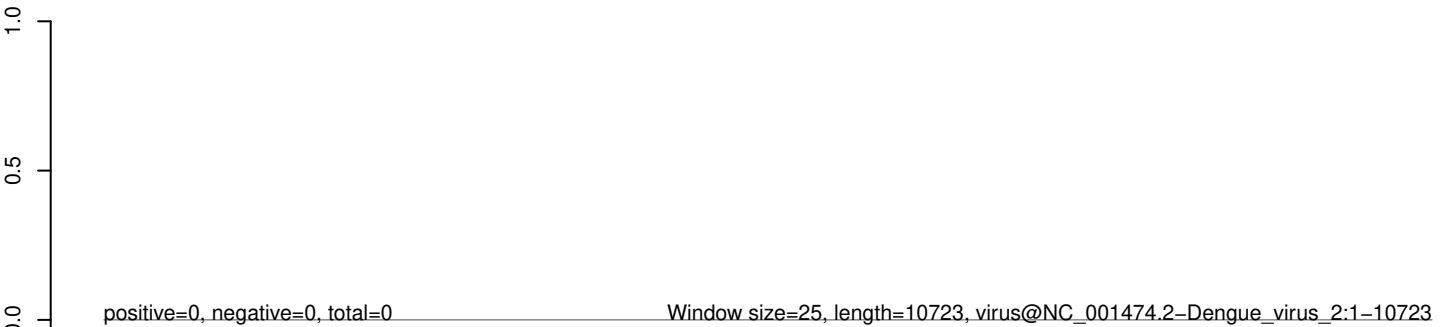
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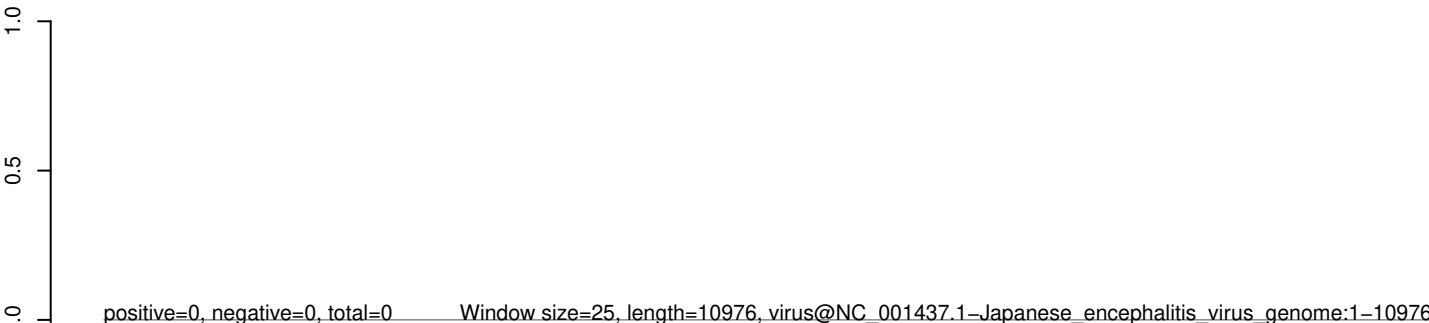
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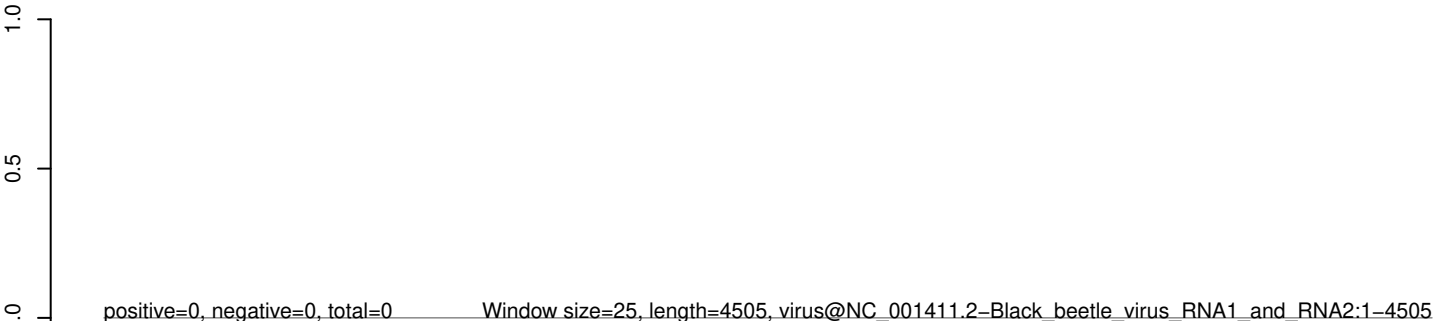
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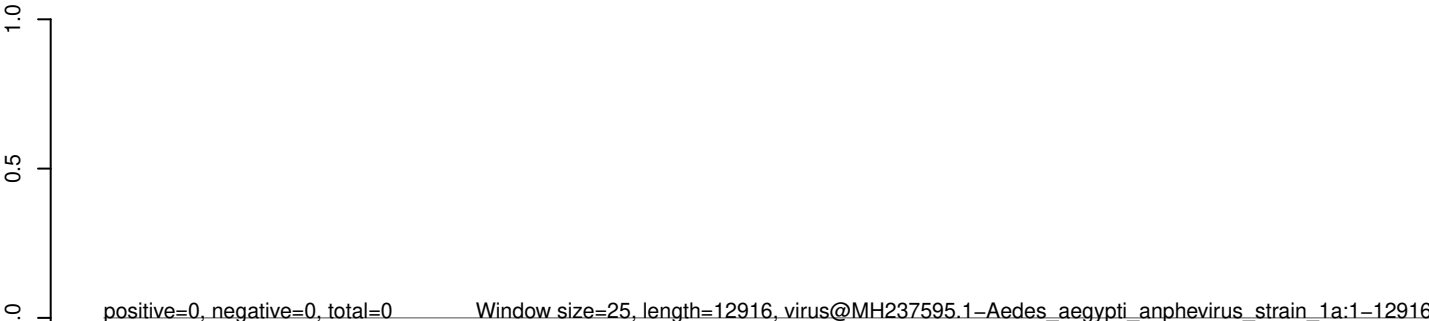
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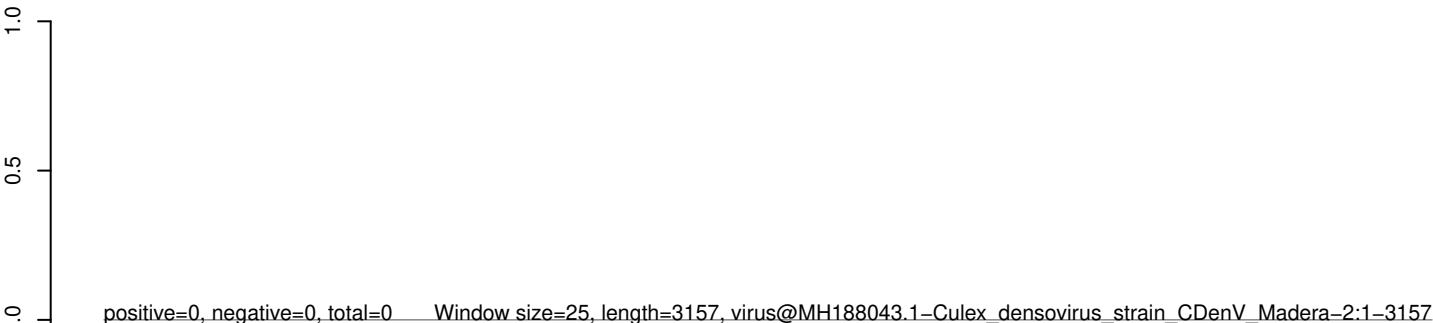
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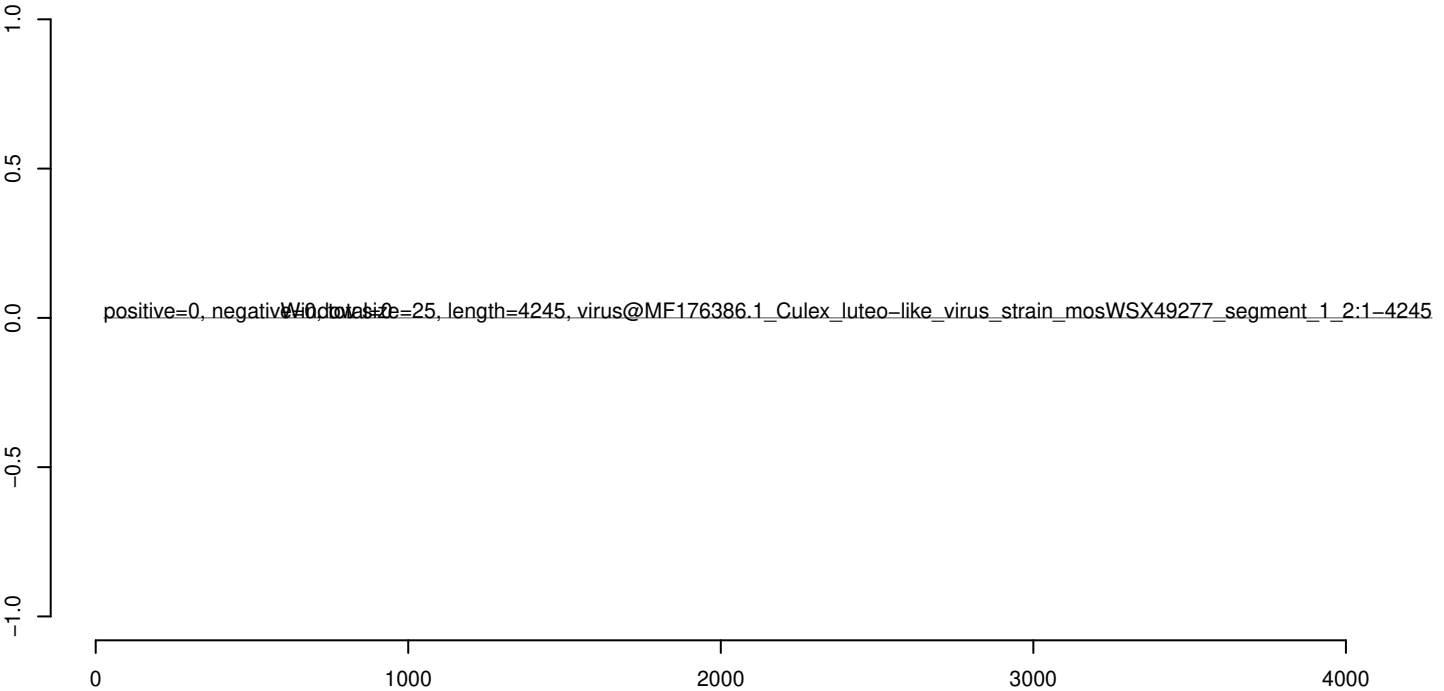
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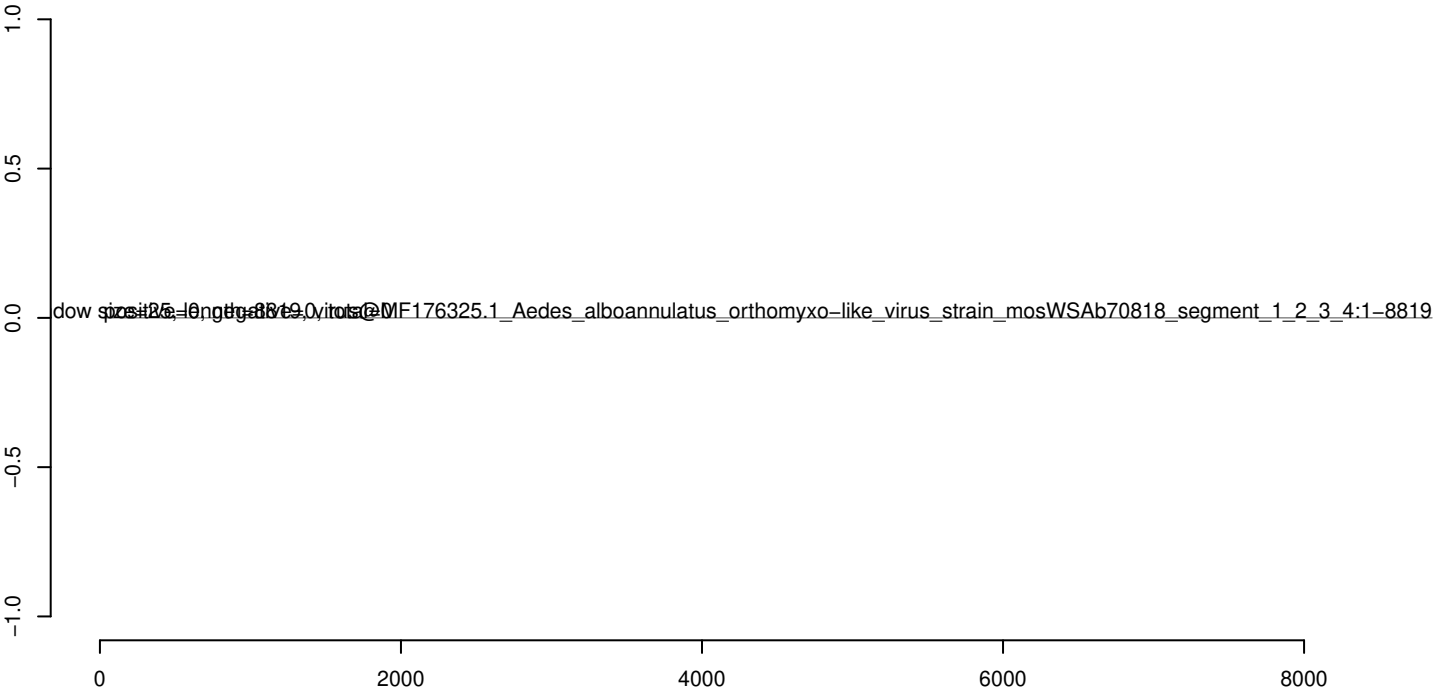
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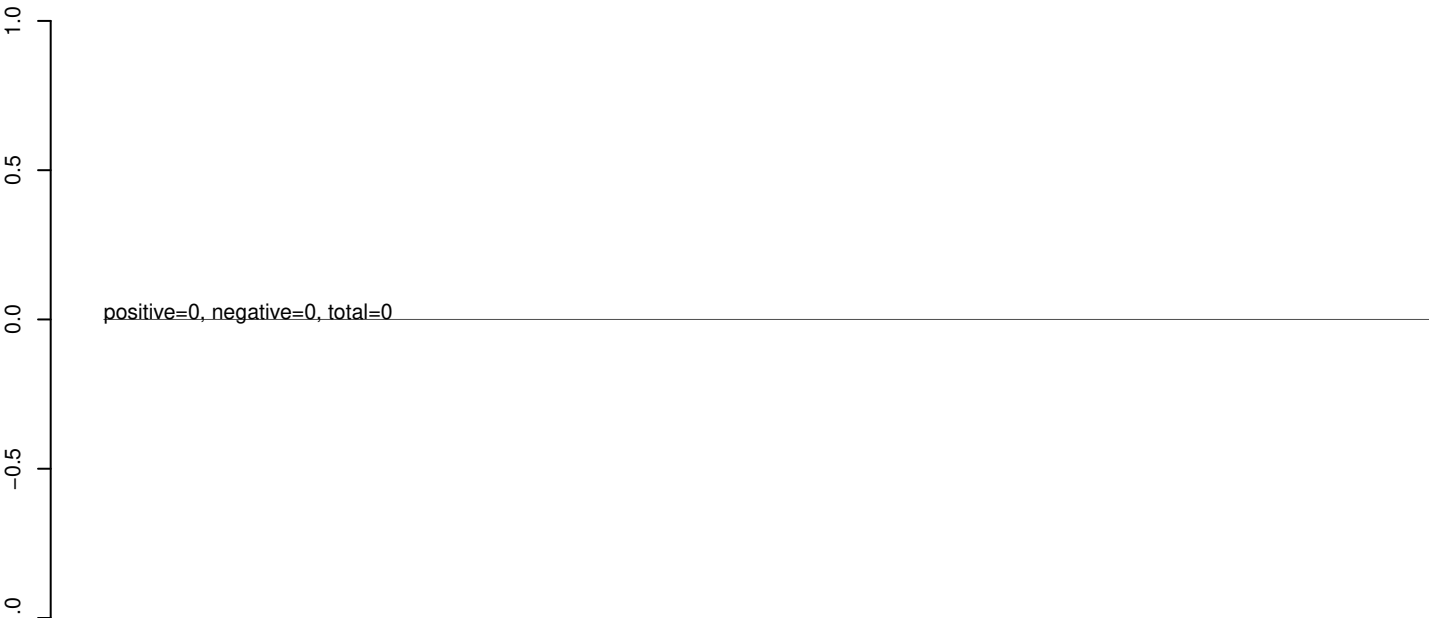
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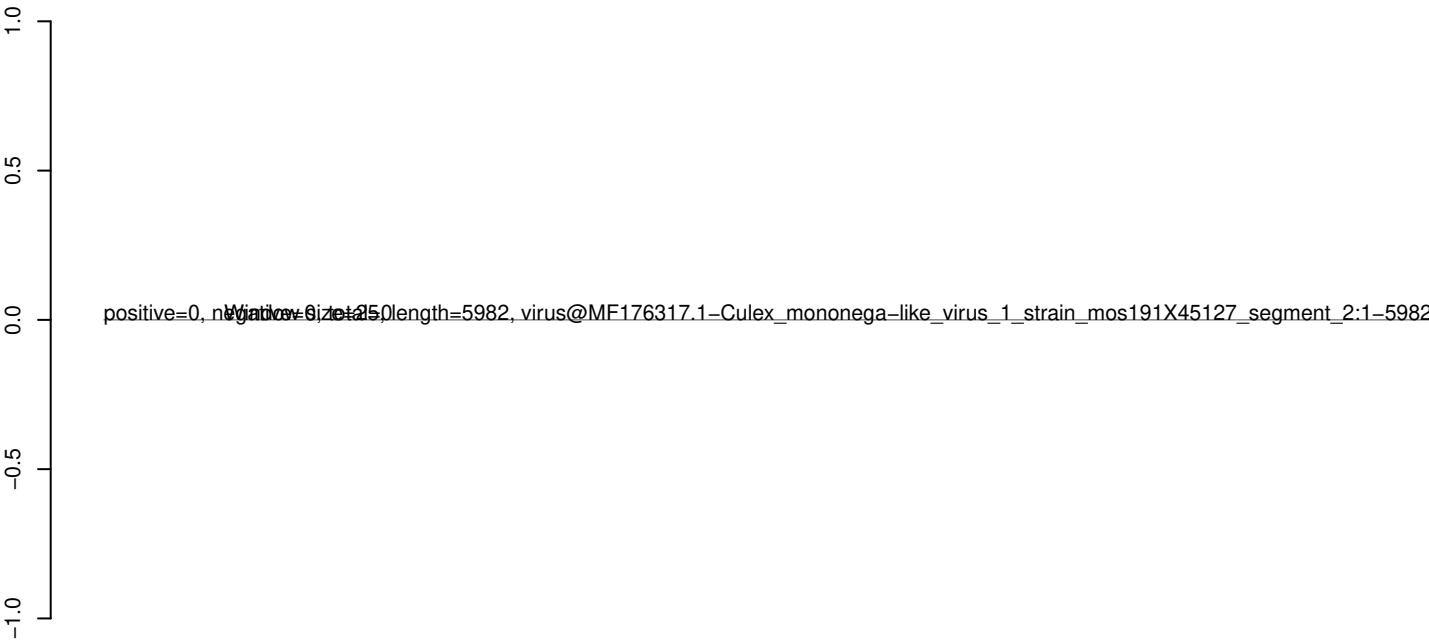
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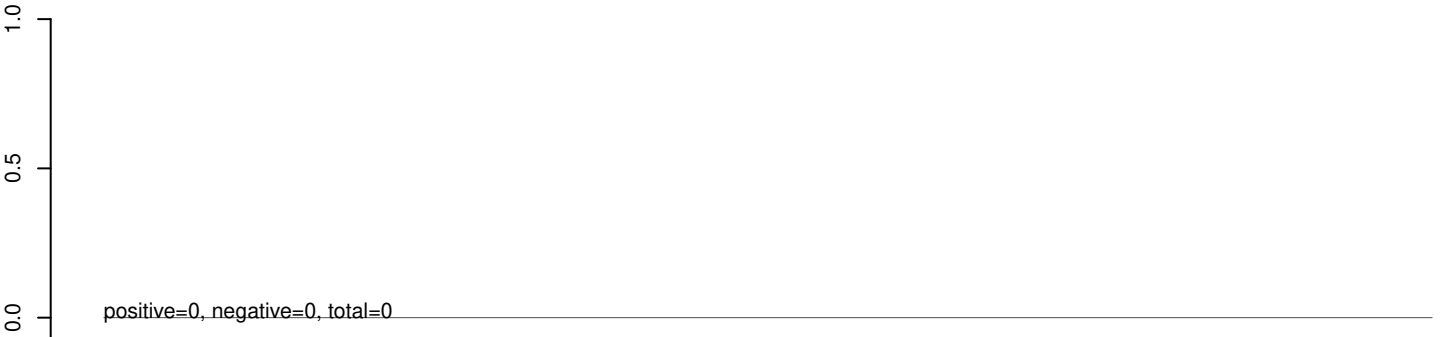
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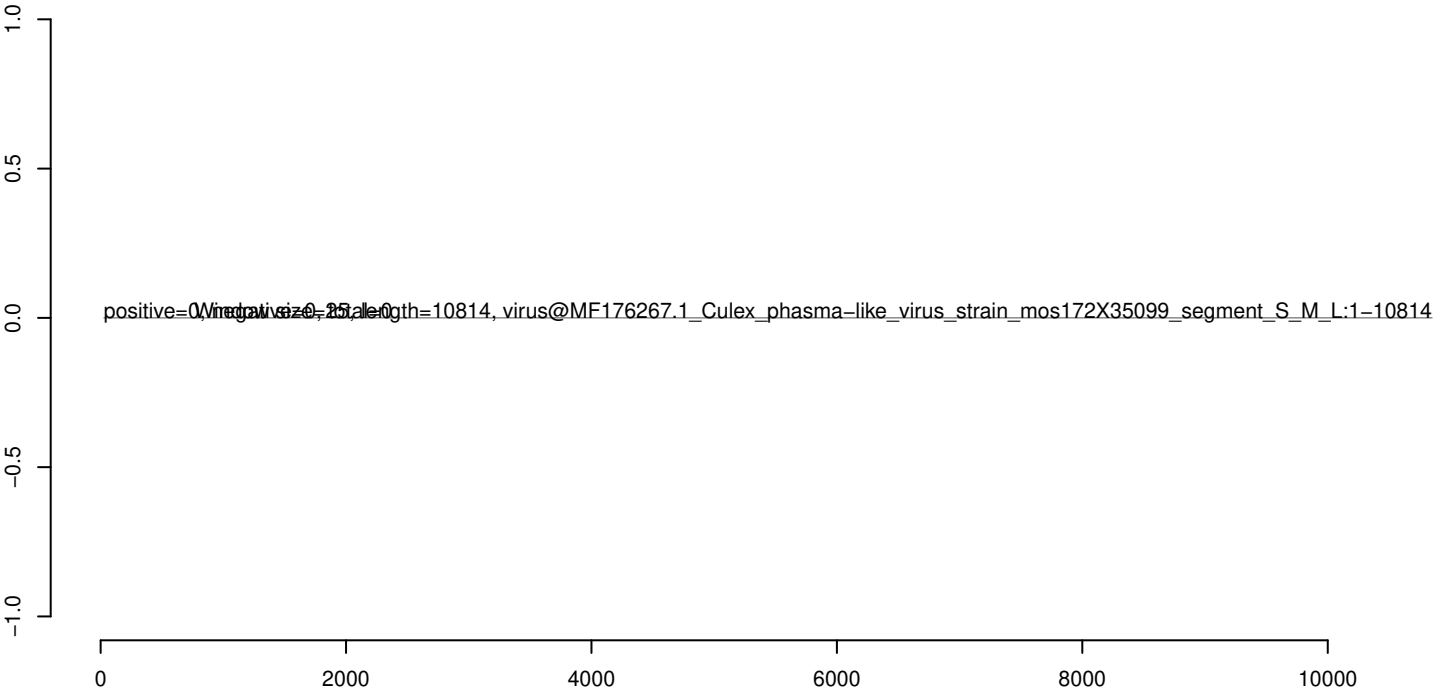
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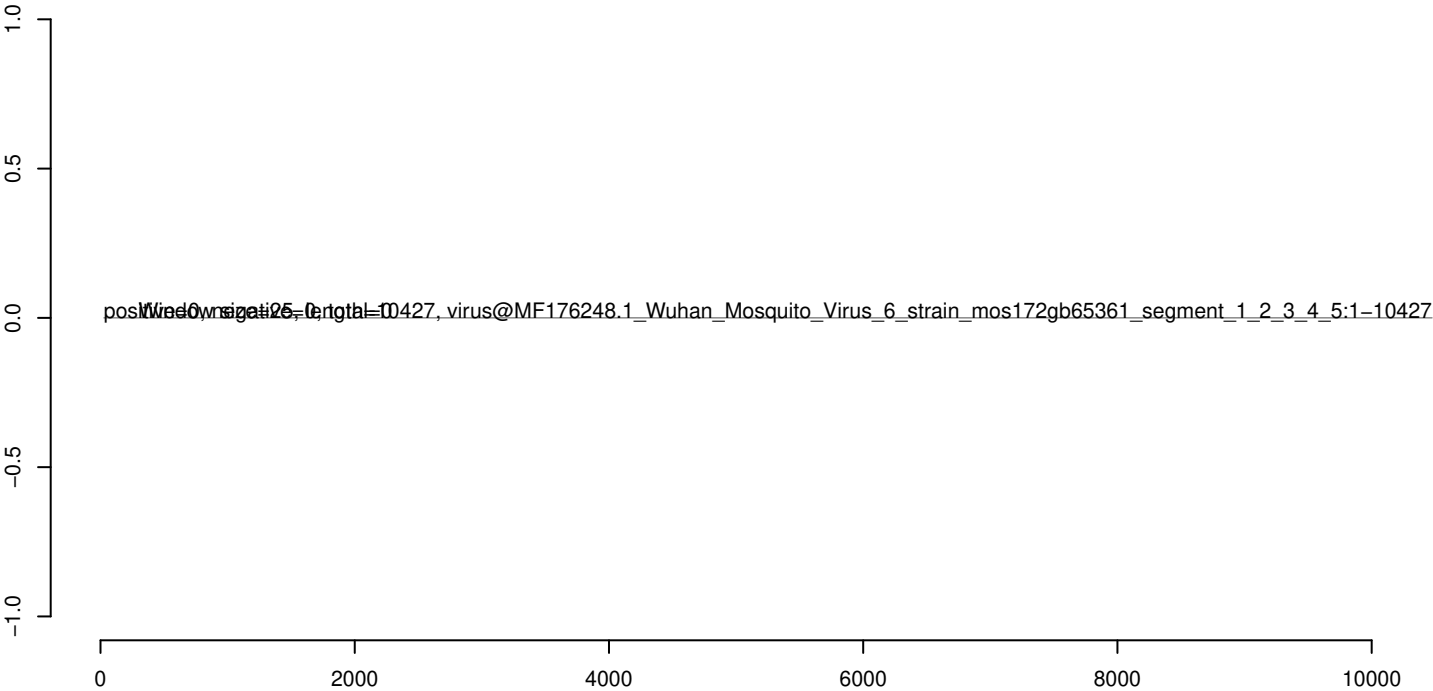
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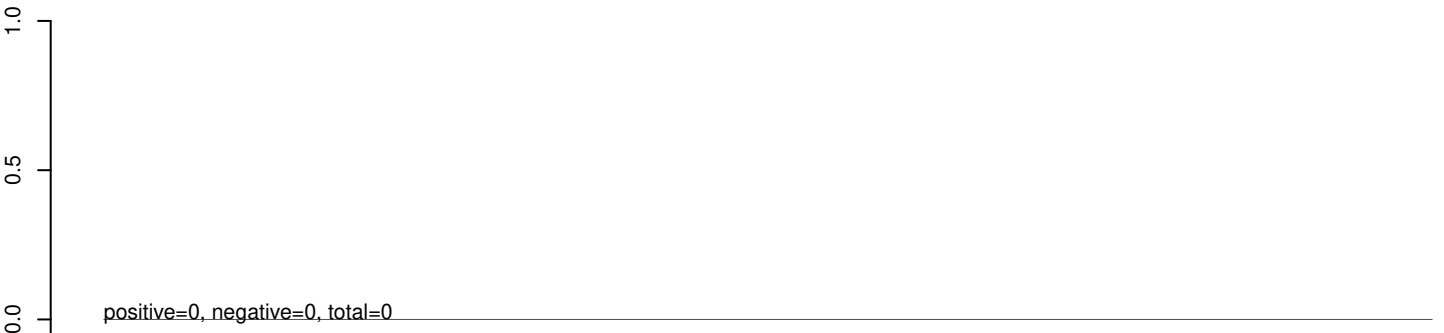
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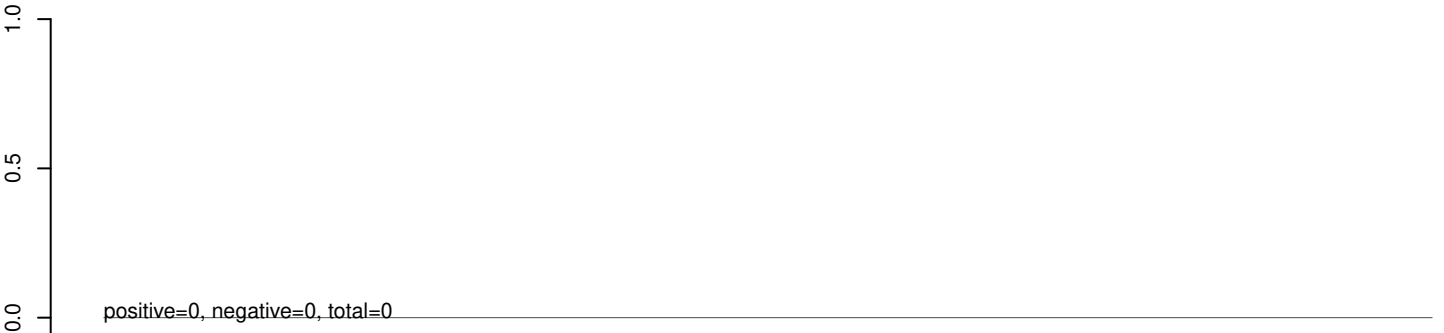
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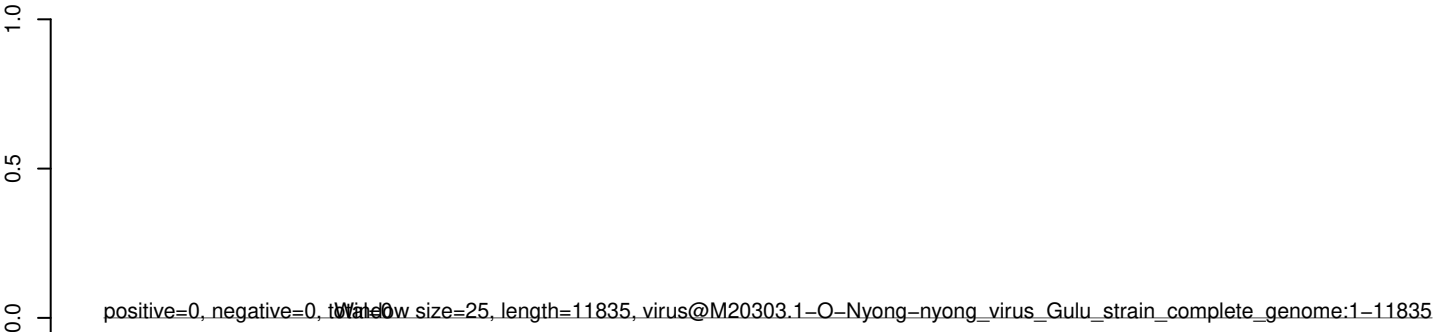
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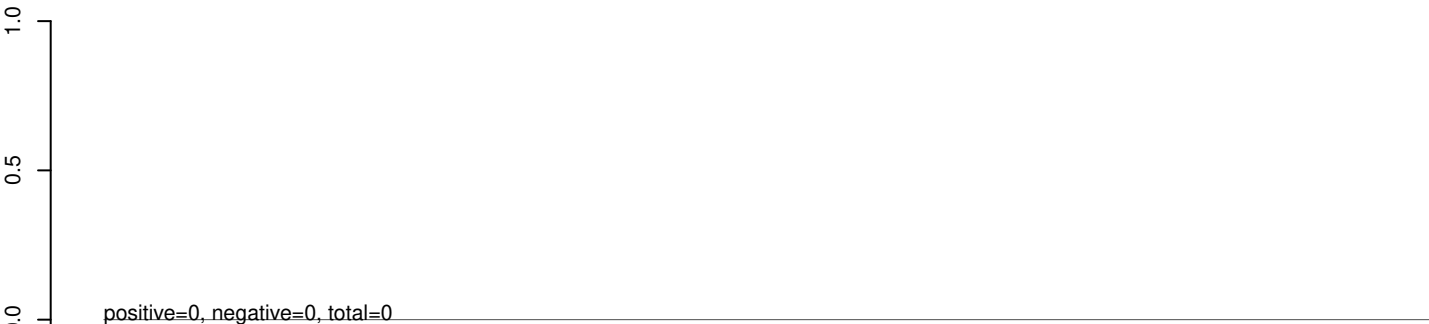
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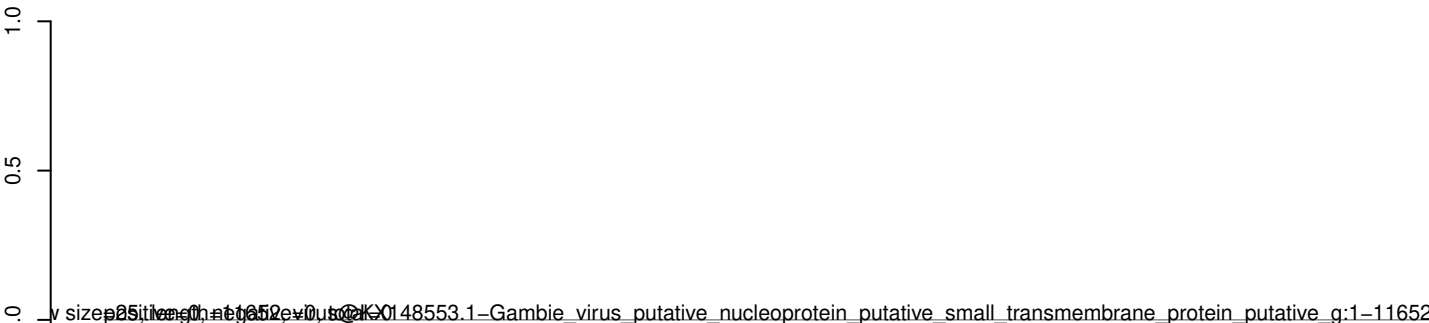
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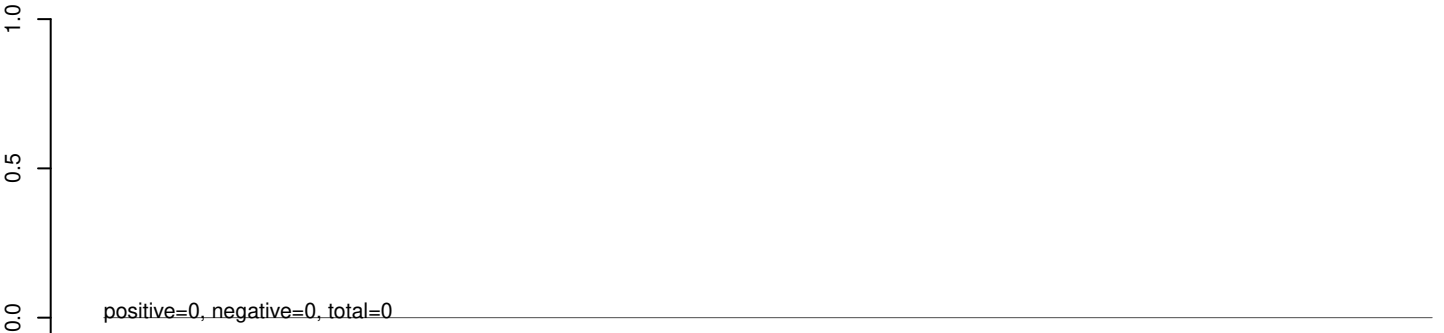


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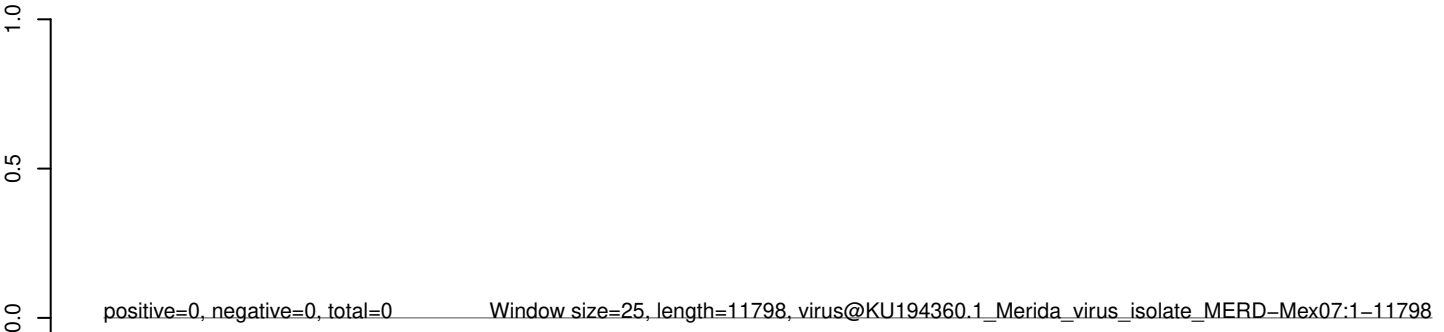
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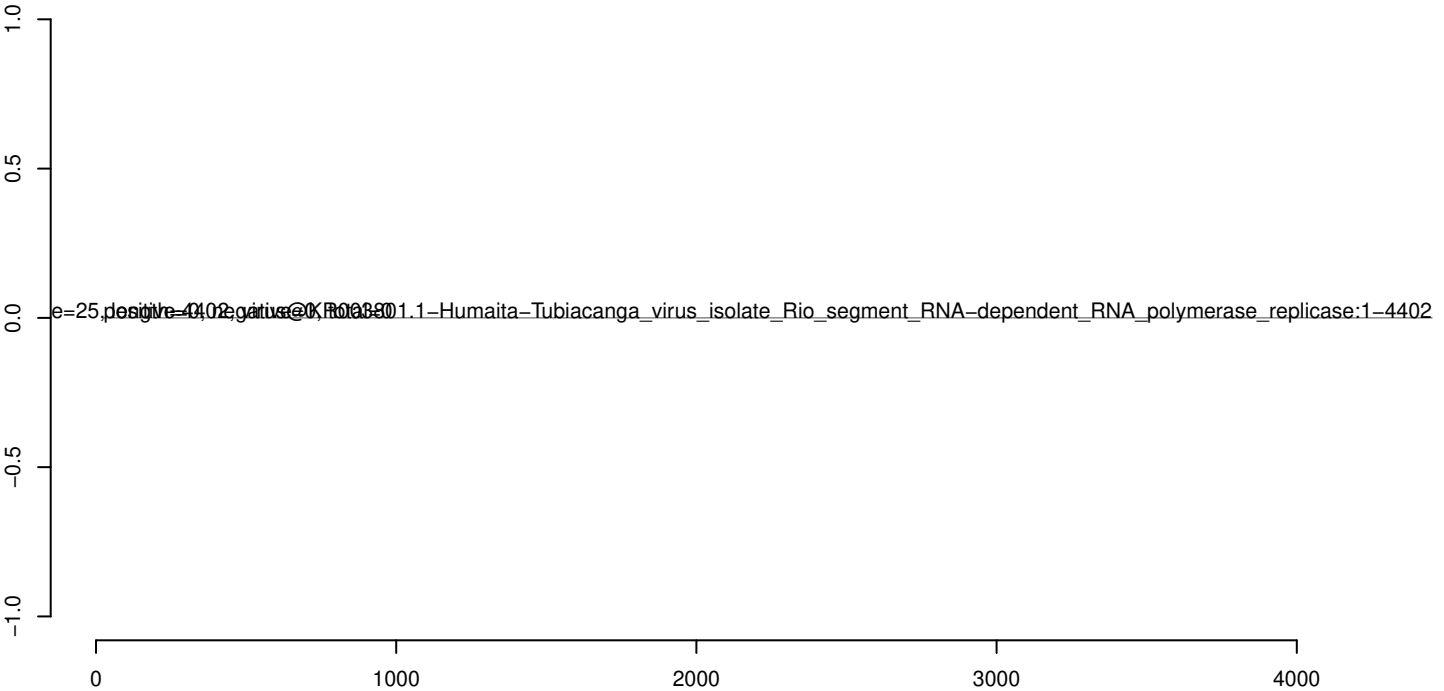
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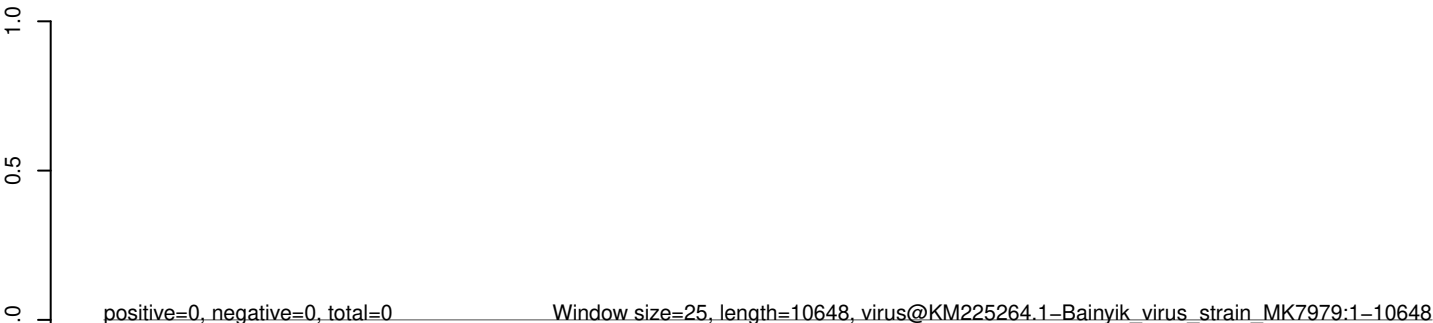
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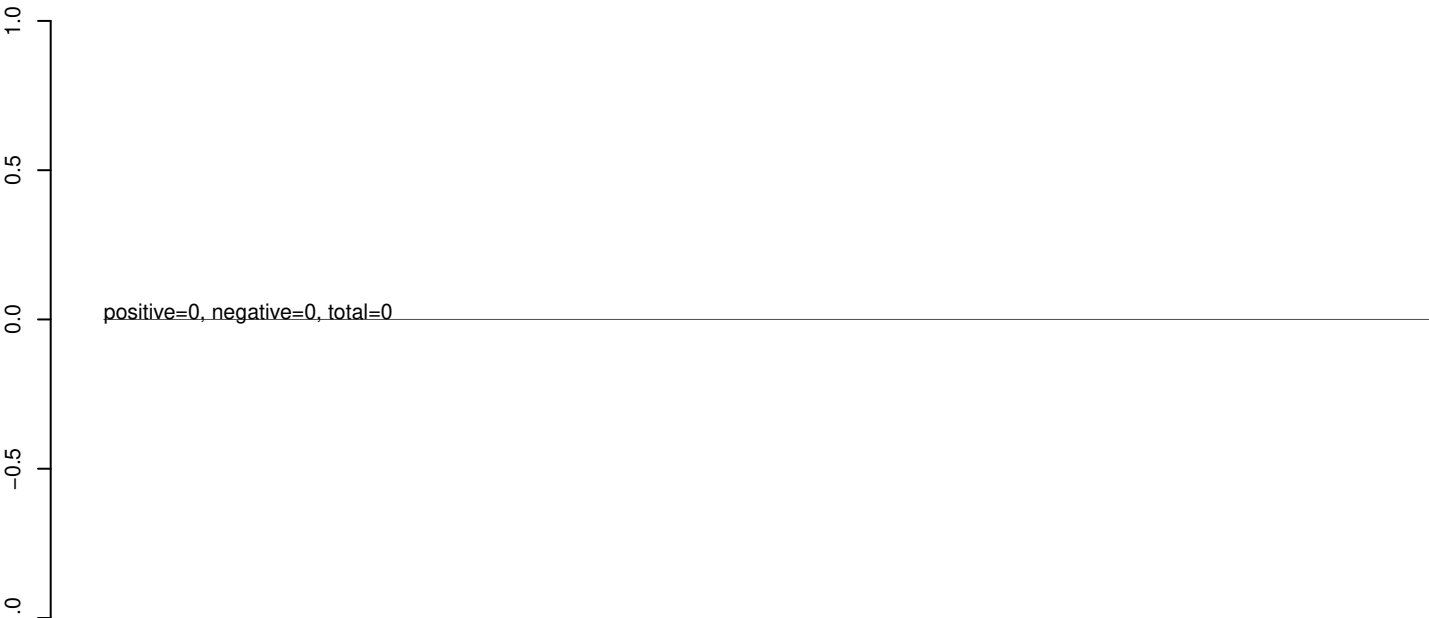
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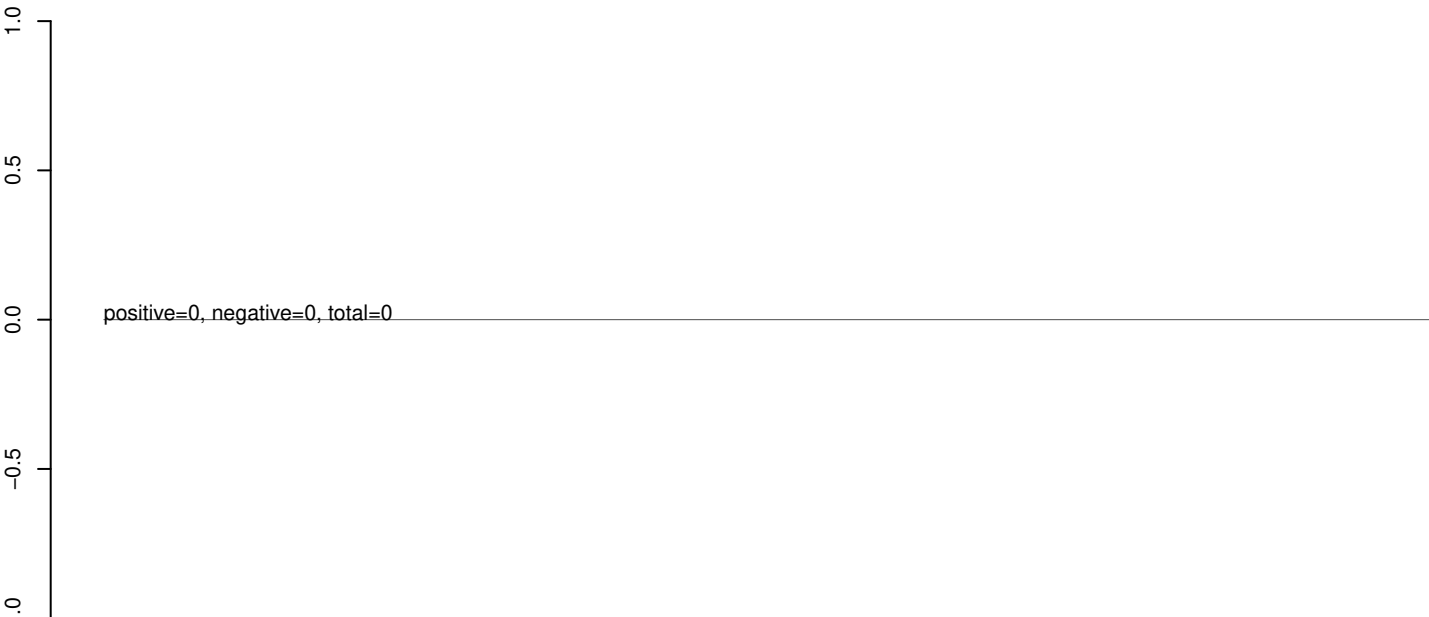
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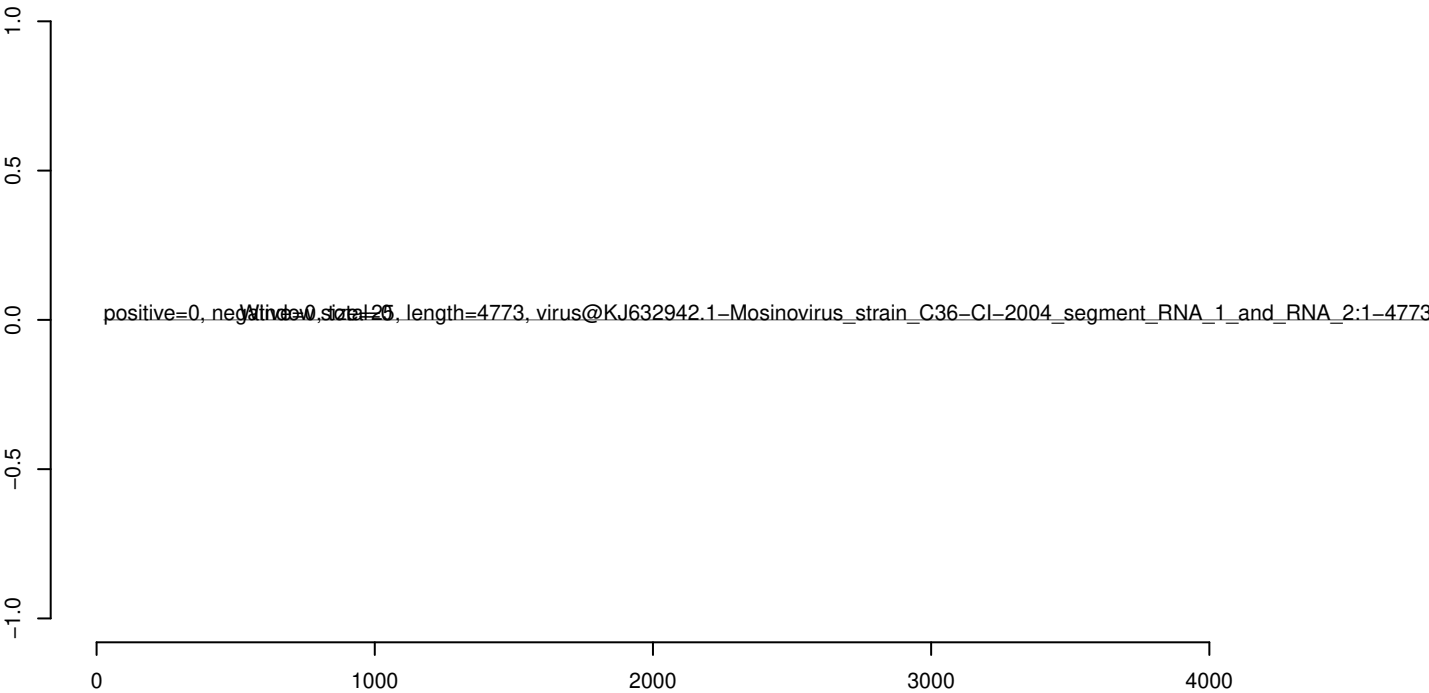
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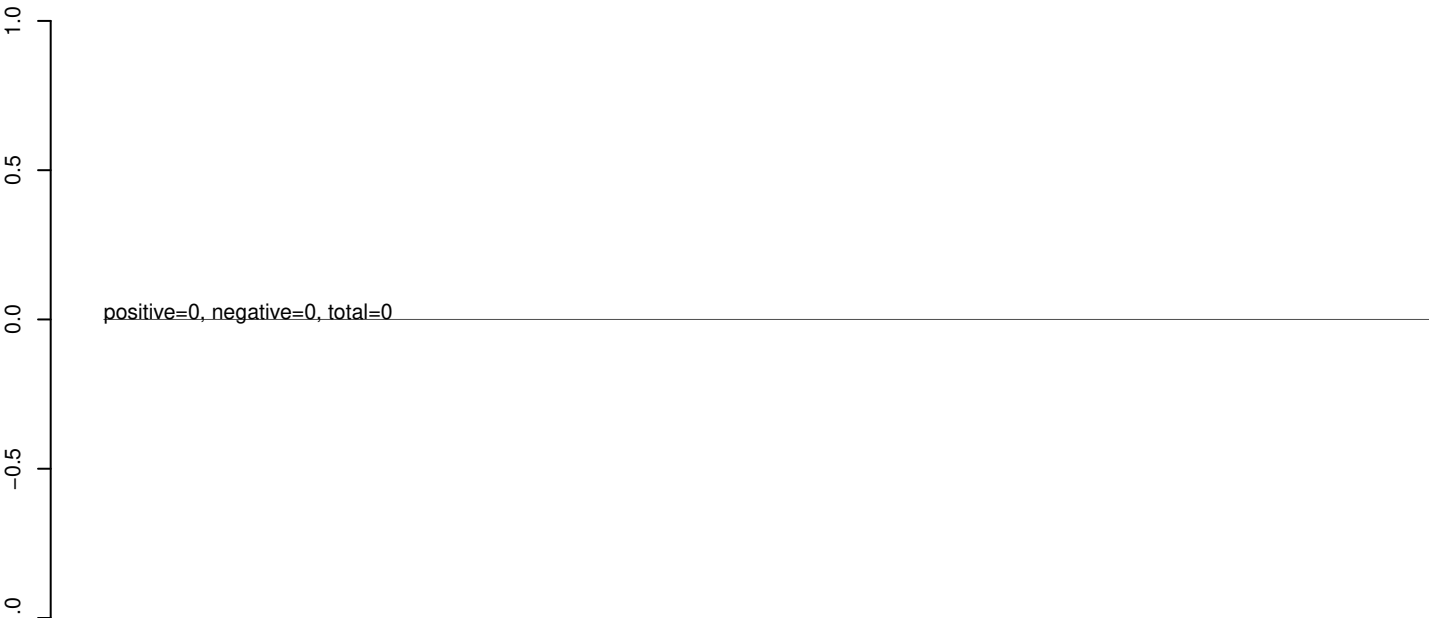
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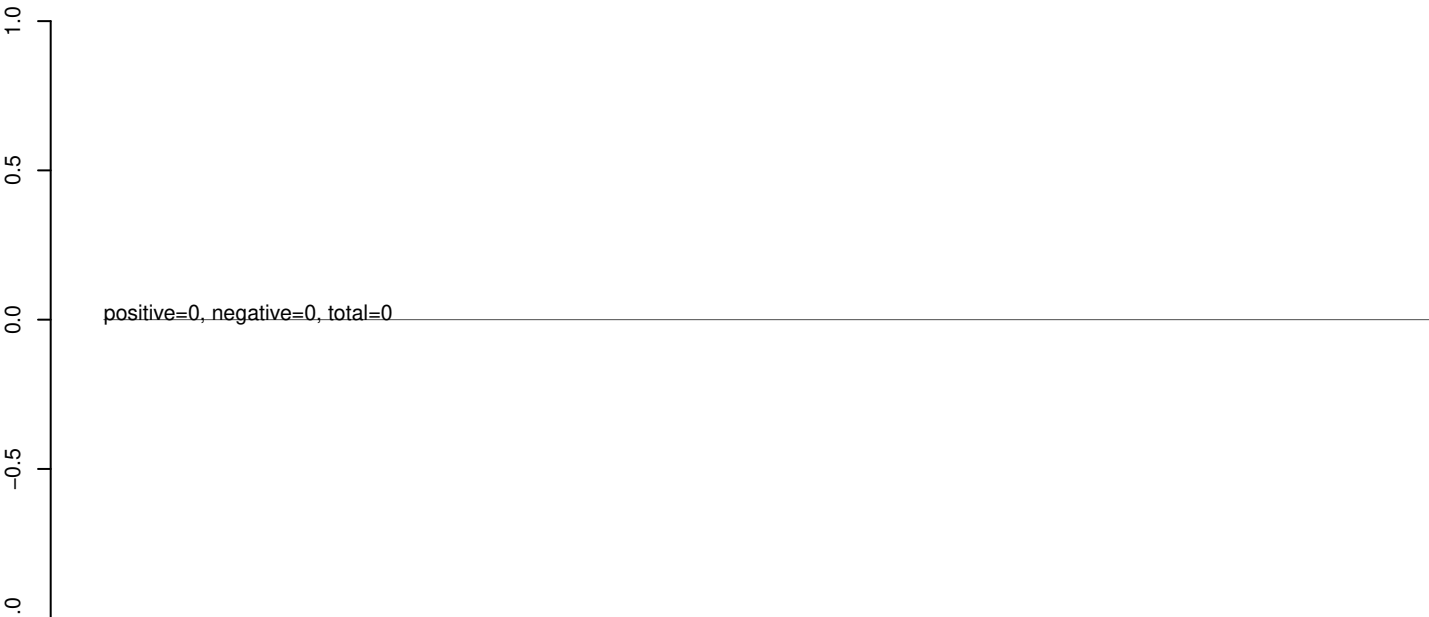
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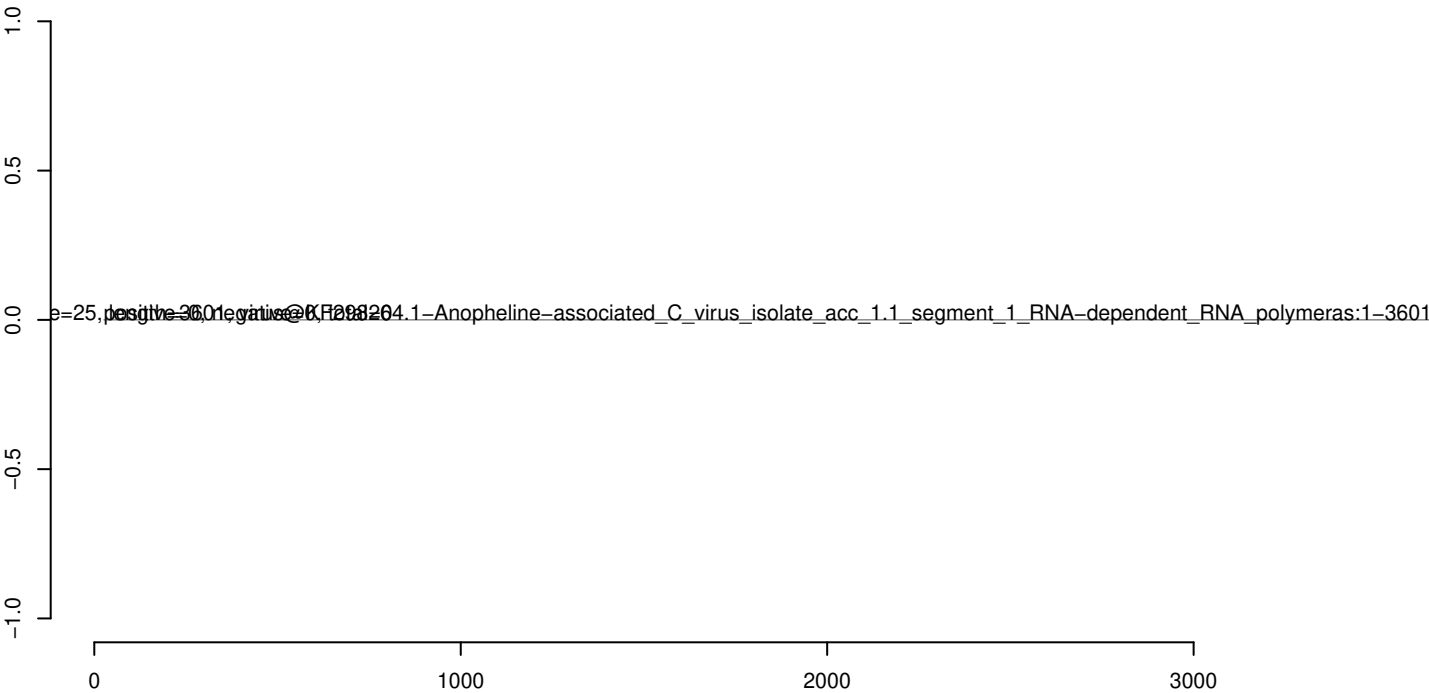
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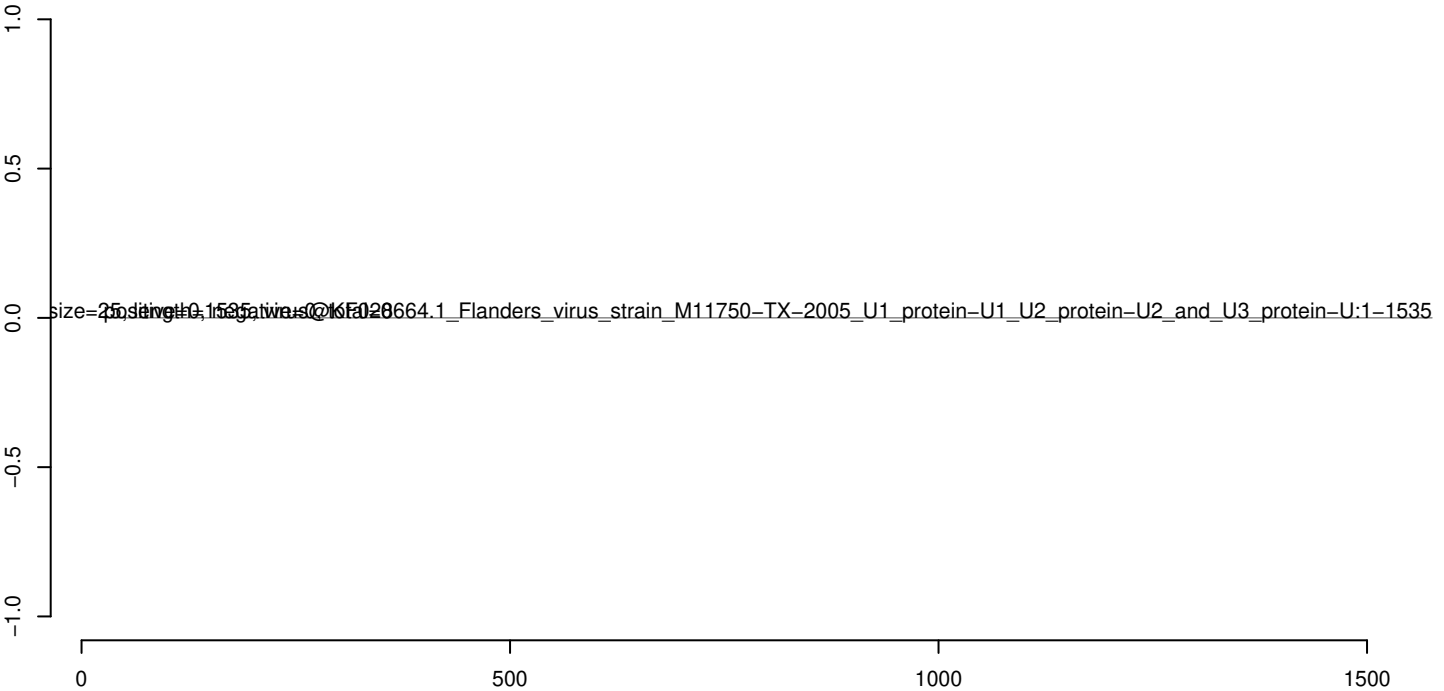
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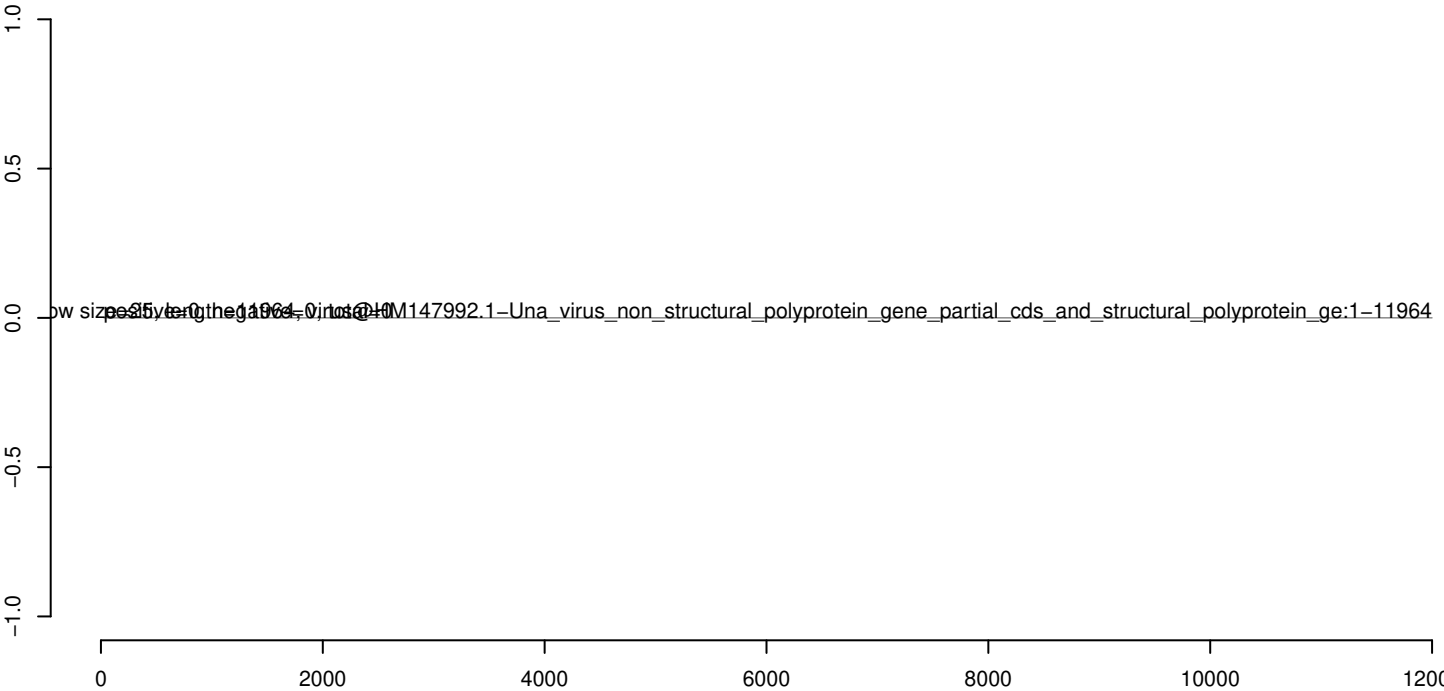
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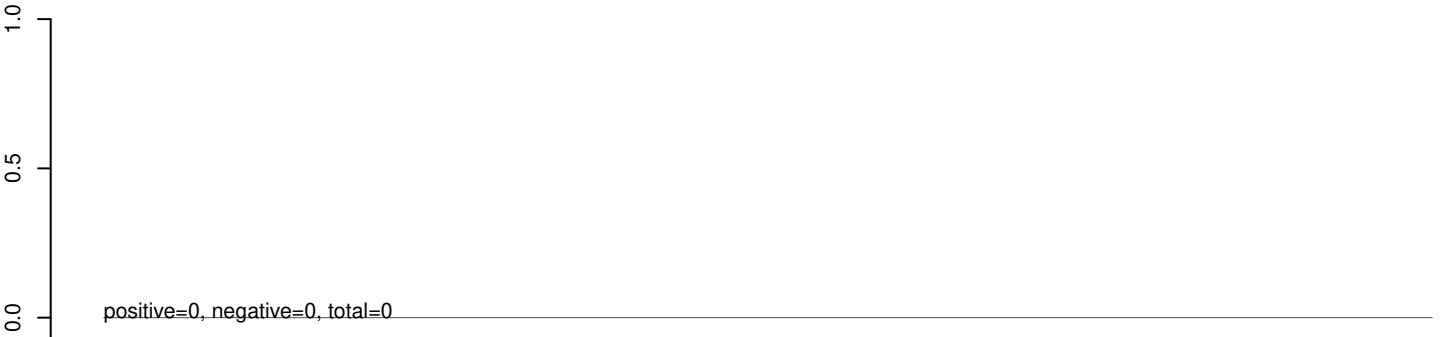
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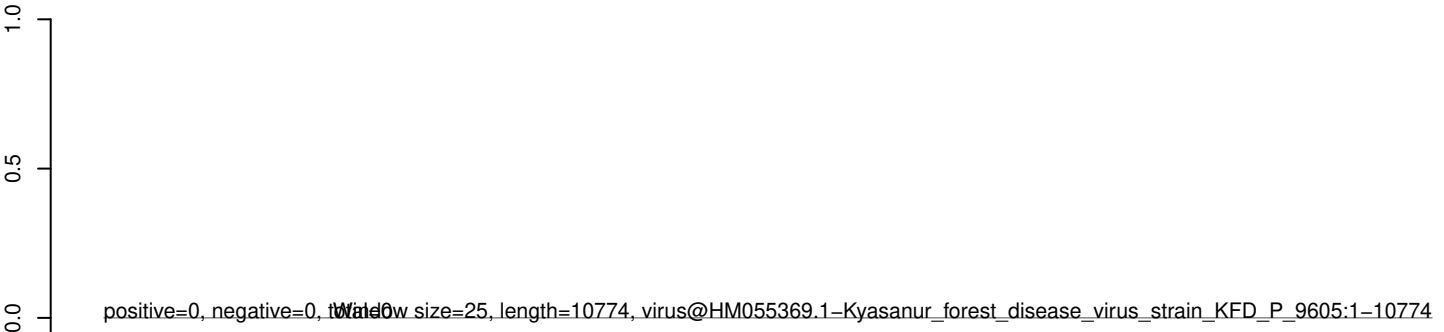
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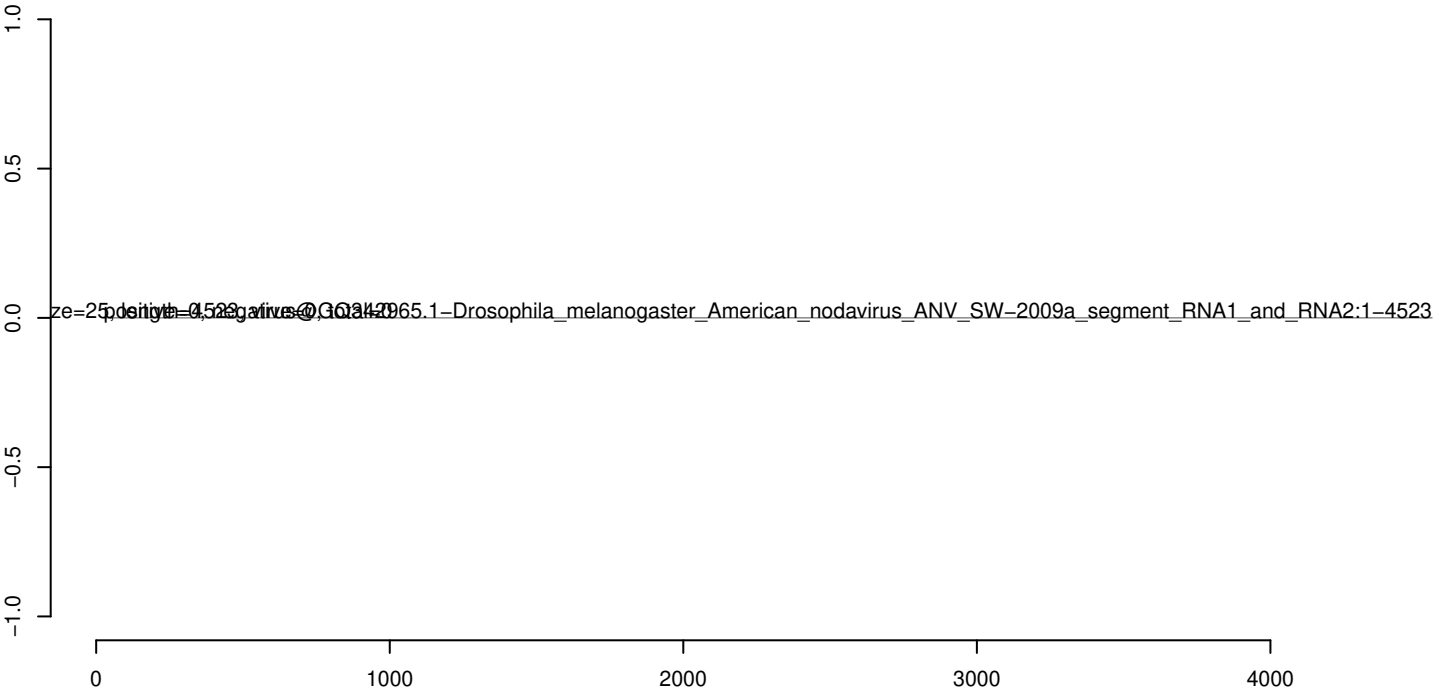
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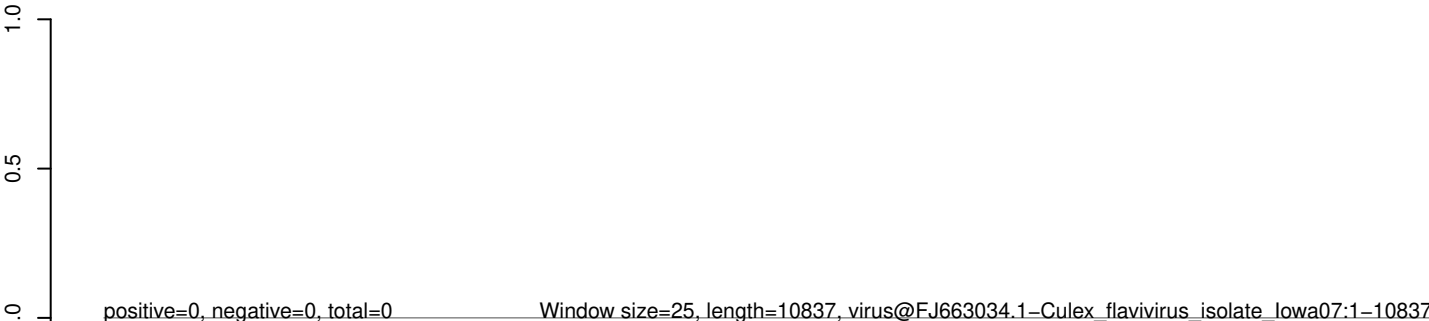
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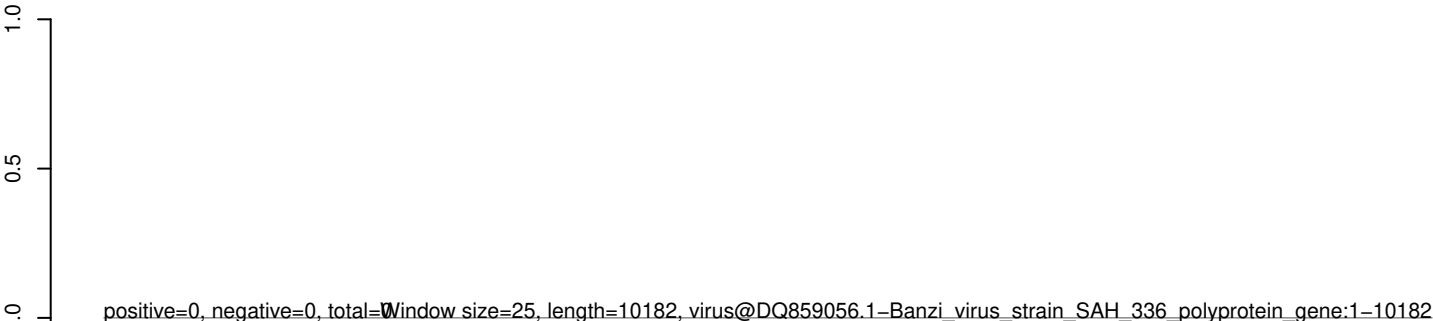
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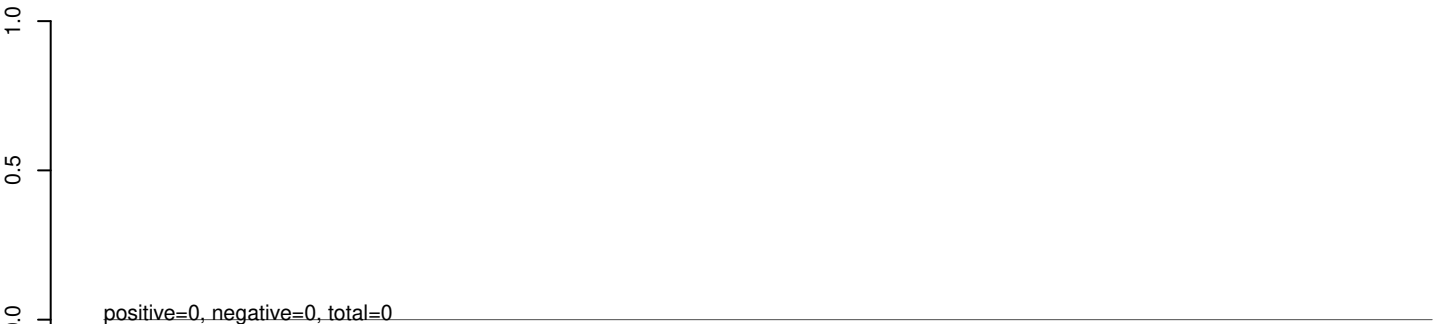
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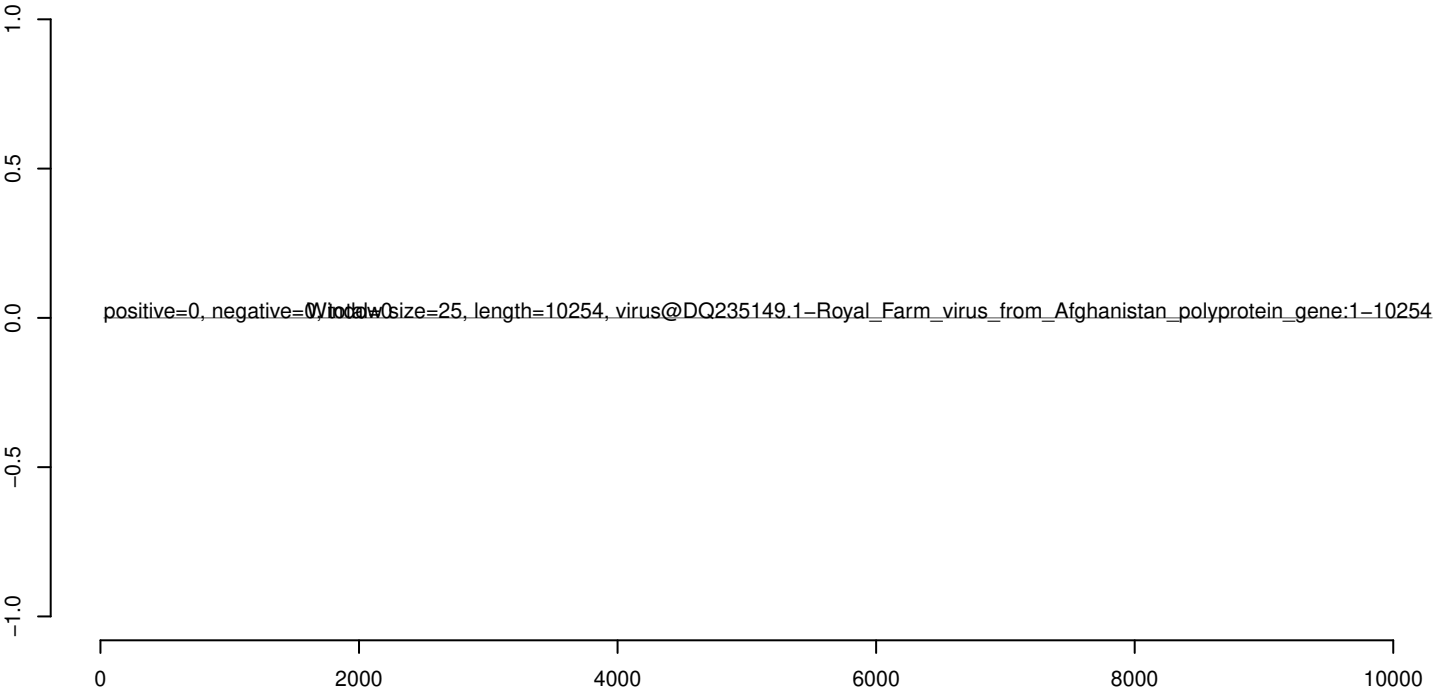
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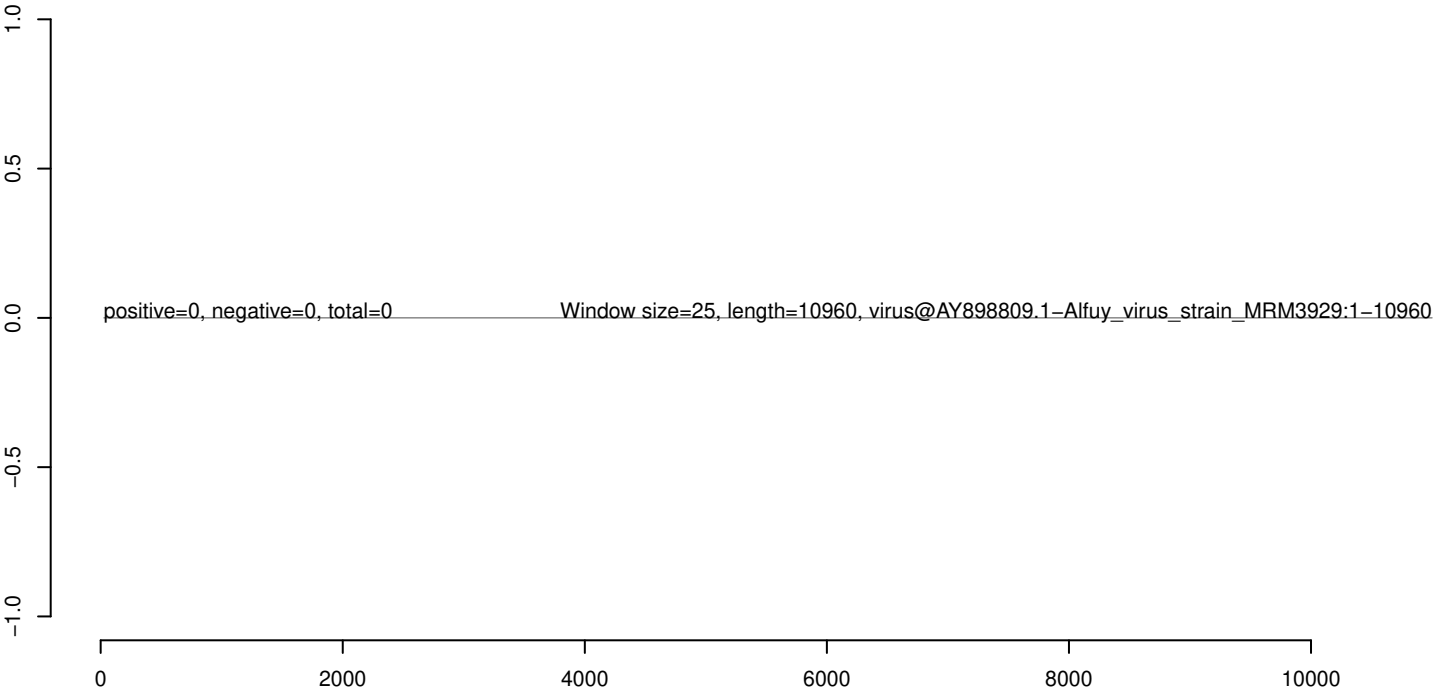
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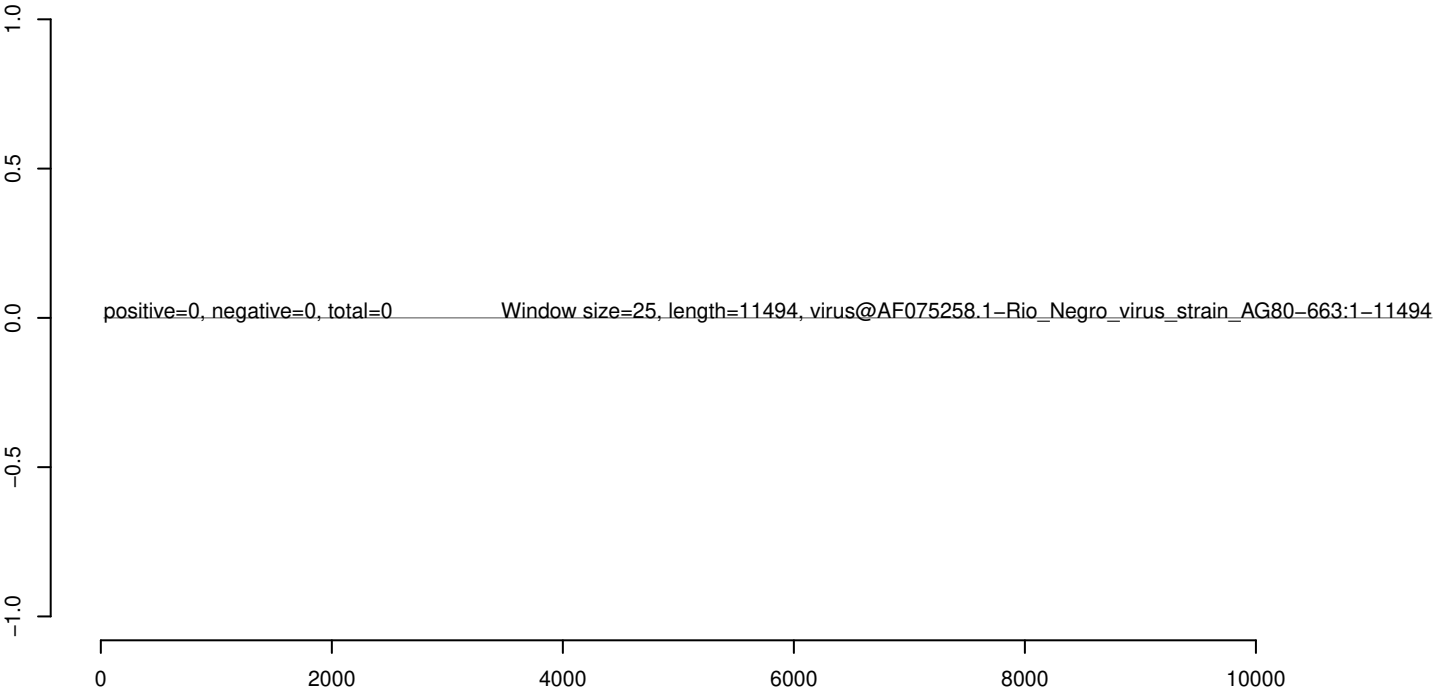
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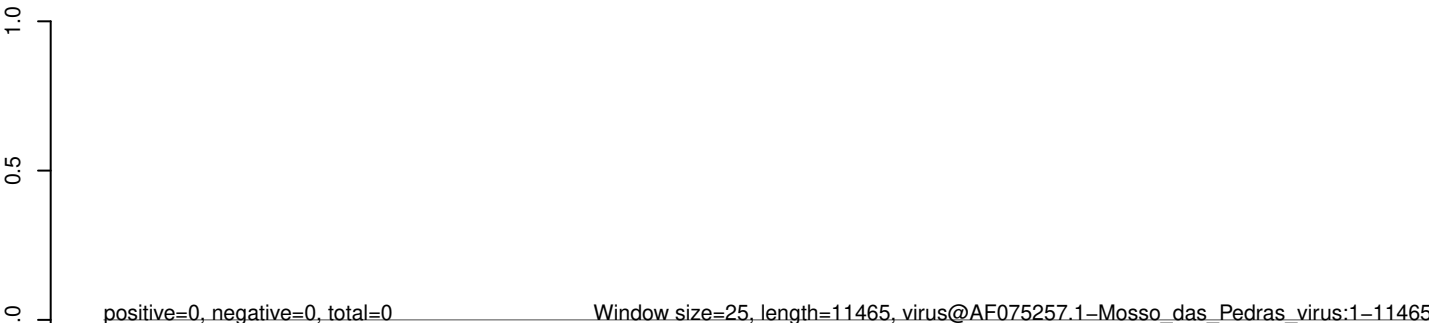
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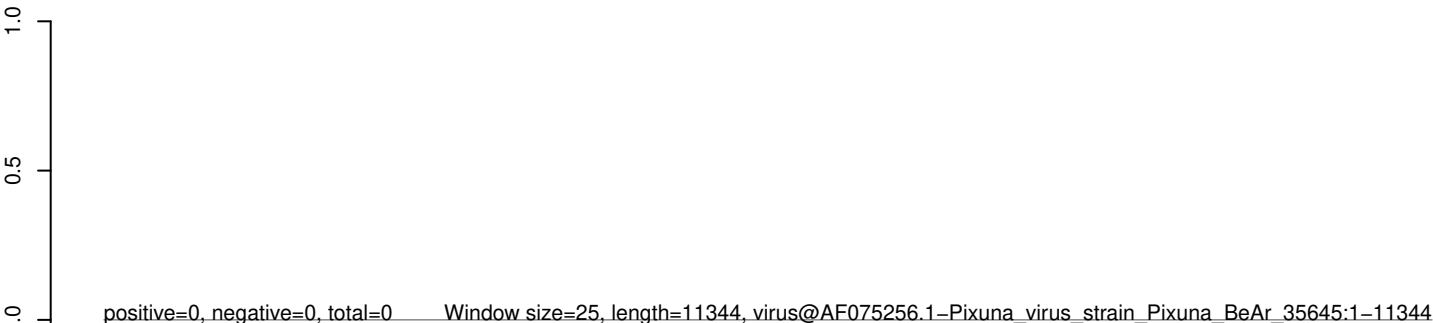
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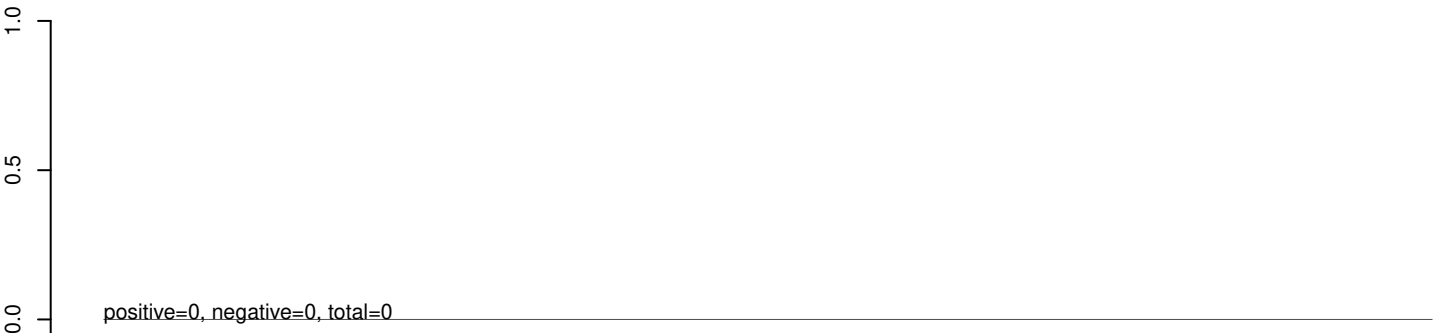


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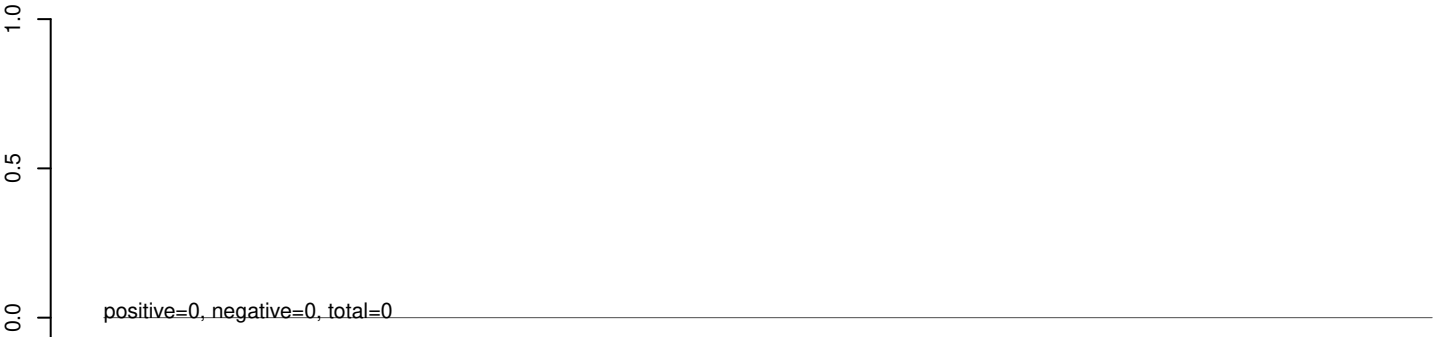


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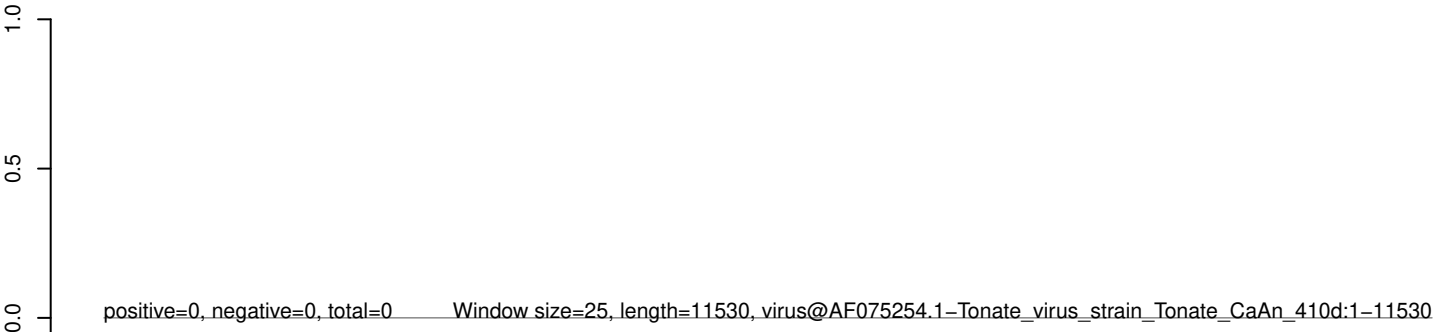
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AeAeg_Aag2_BUNV1_ES.rep



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AeAeg_Aag2_BUNV1_ES.rep

