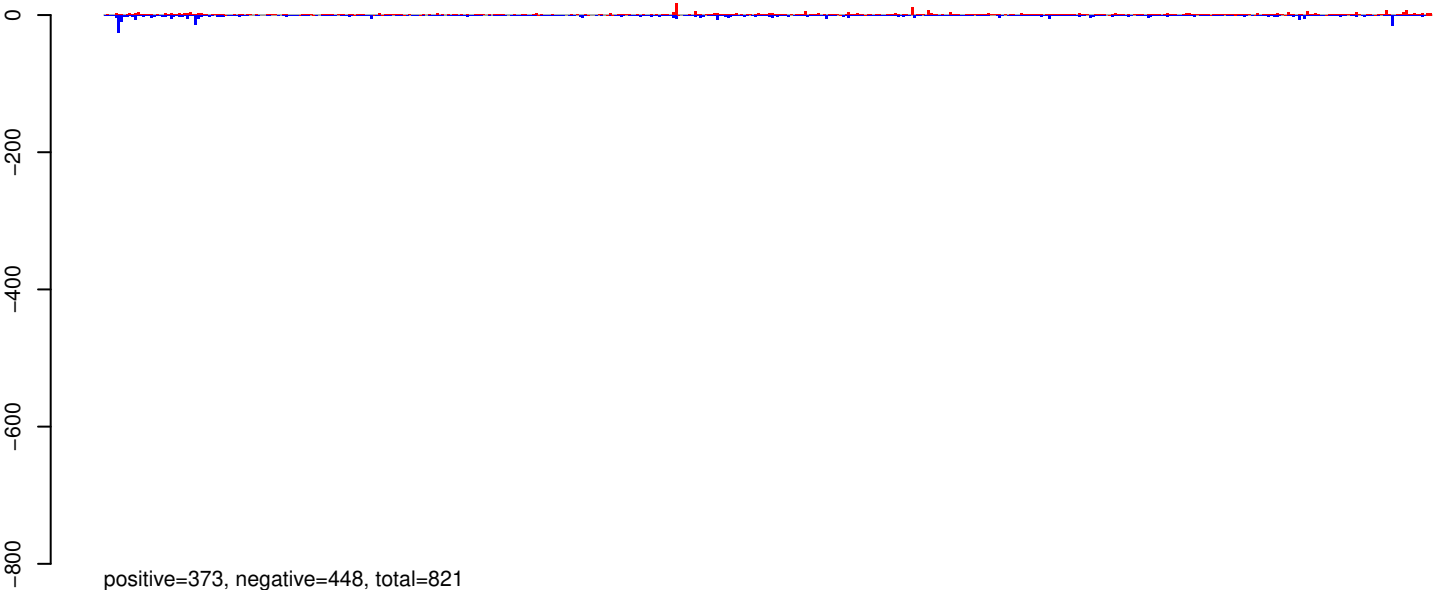
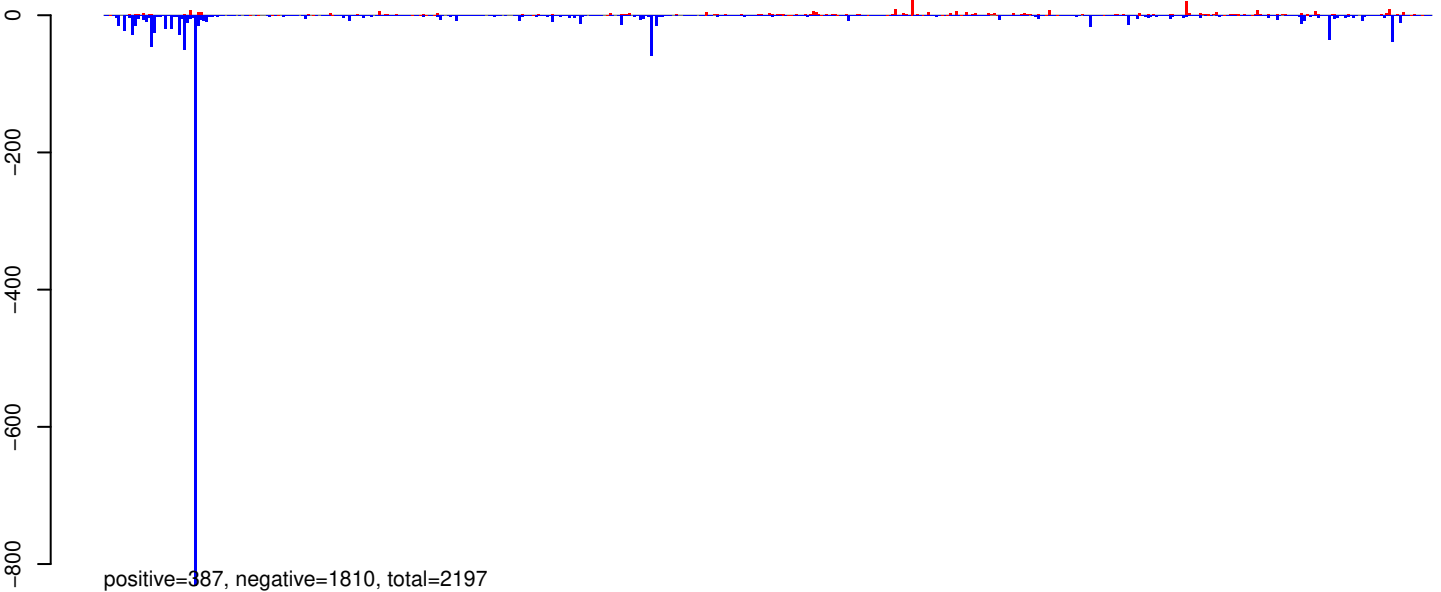


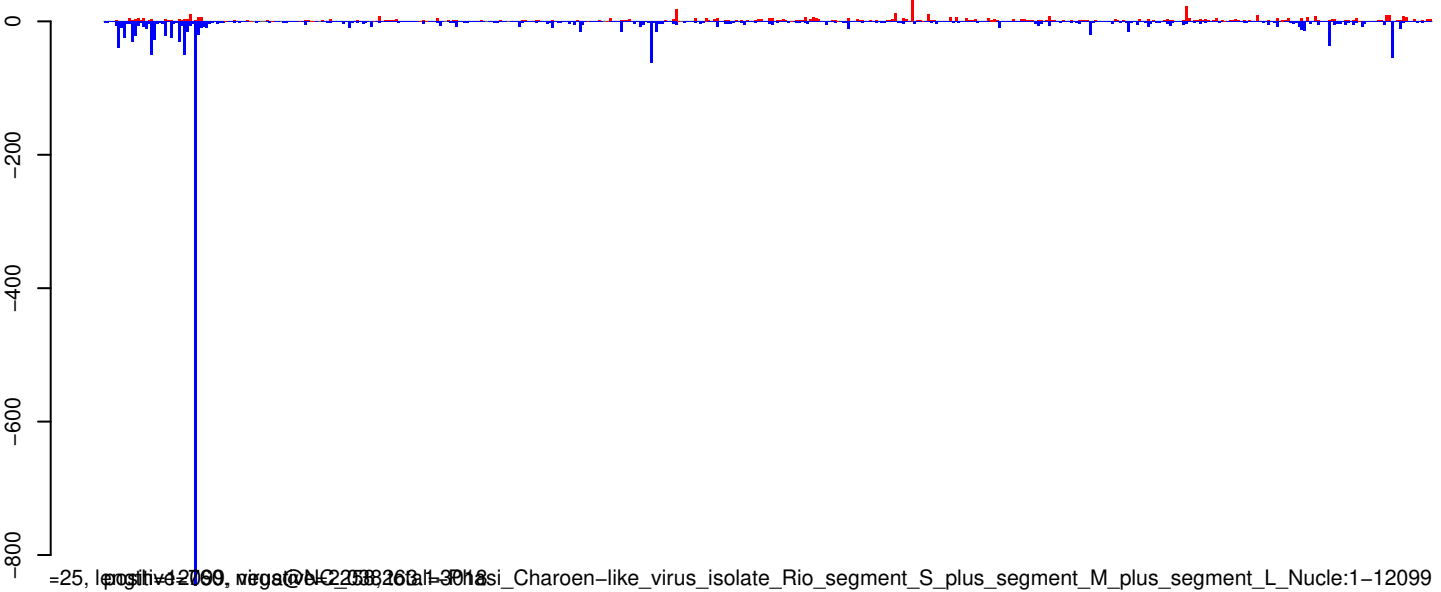
AeAeg_Aag2_BTV1_AK.18_23.rep



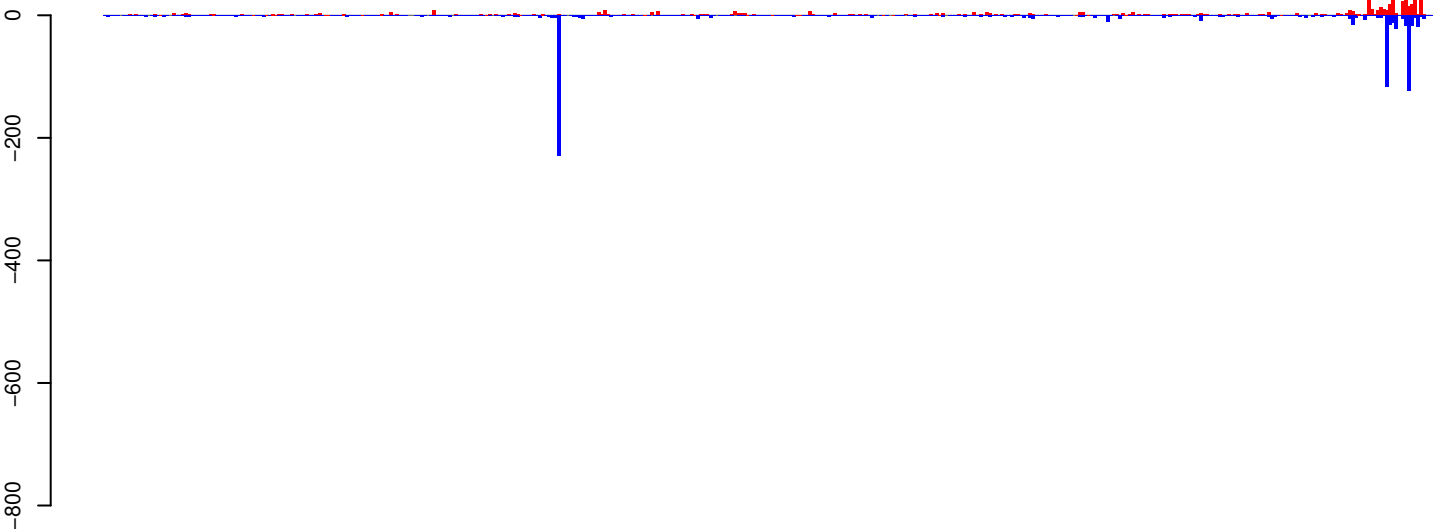
AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep

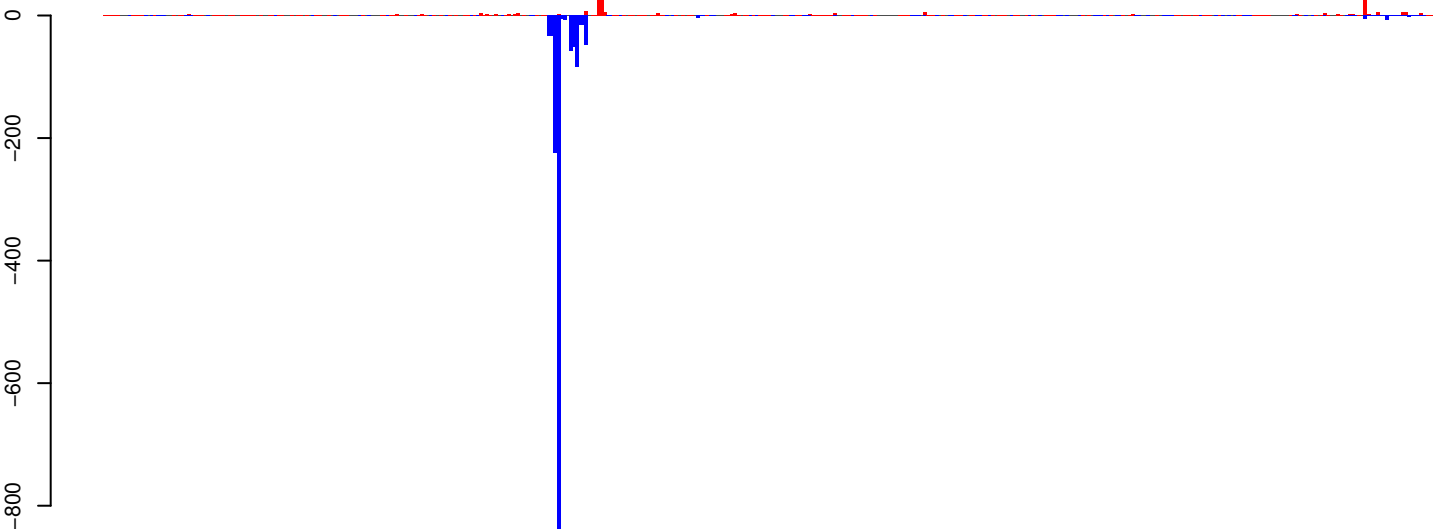


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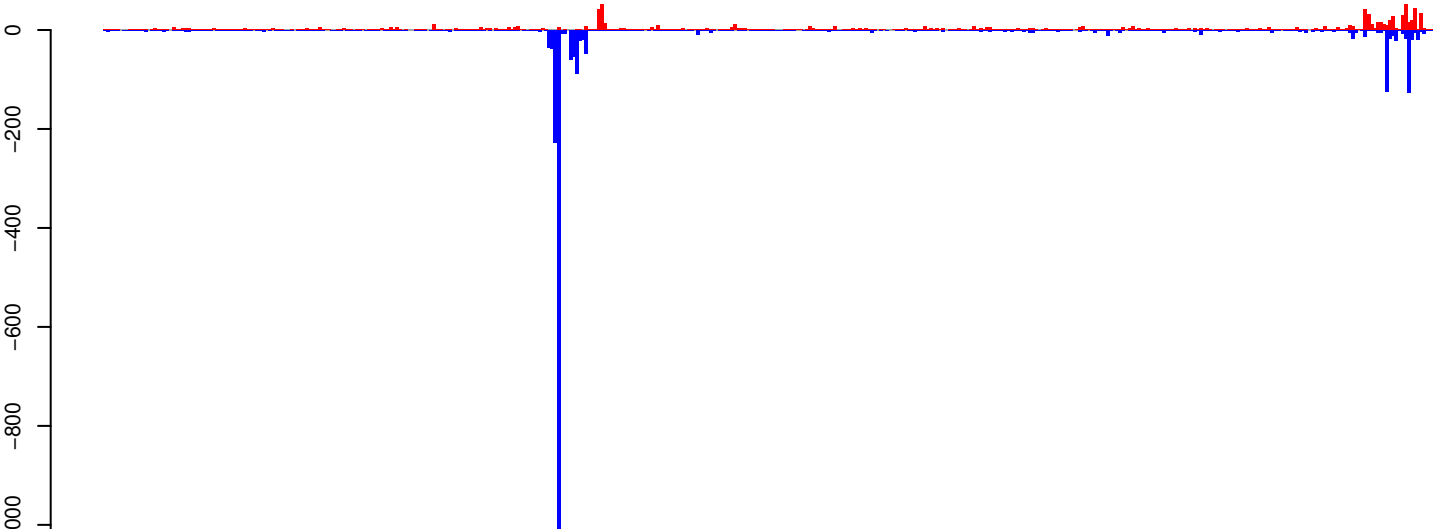
positive=692, negative=977, total=1668

AeAeg_Aag2_BTV1_AK.24_35.rep



positive=310, negative=1544, total=1854

AeAeg_Aag2_BTV1_AK.rep



positive=1002, negative=2520, total=3522, size=25, length=10682, virus@NC_001564.2-Cell_fusing_agent_virus_strain_Galveston:1-10682

AeAeg_Aag2_BTV1_AK.18_23.rep

positive=690, negative=676, total=1366

AeAeg_Aag2_BTV1_AK.24_35.rep

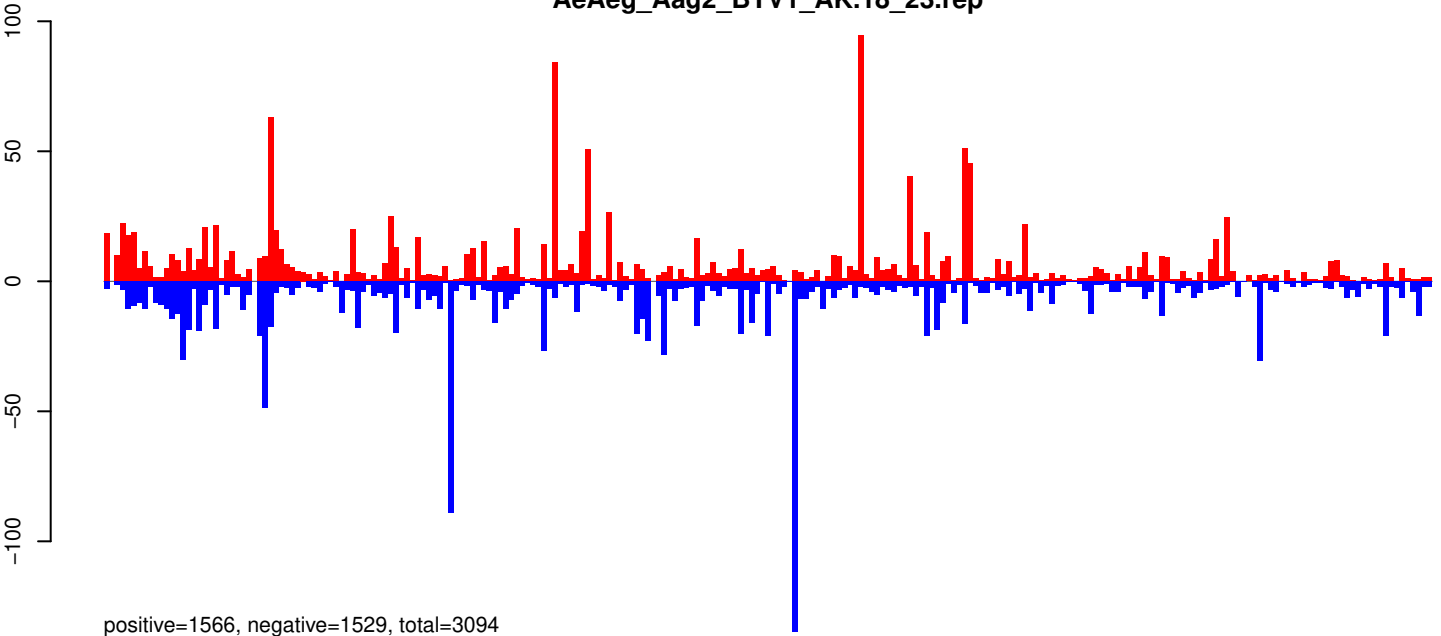
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AeAeg_Aag2_BTV1_AK.rep

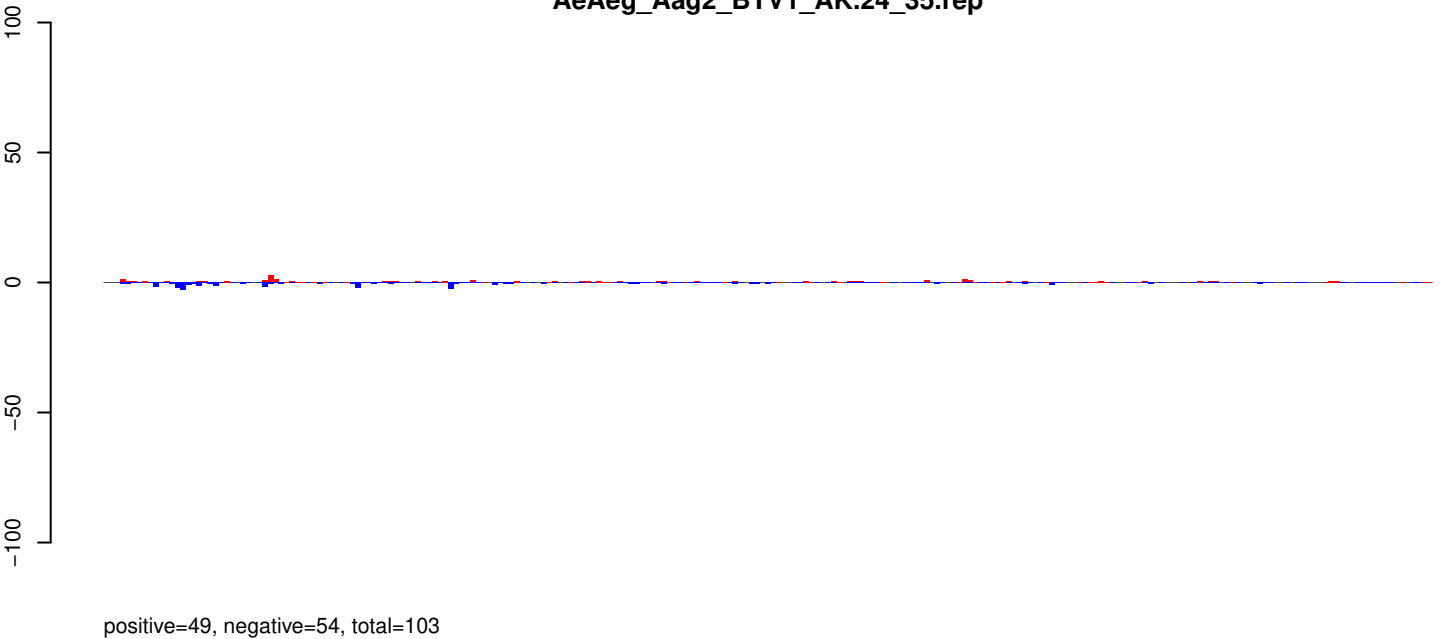
positive=15619, negative=723, total=16342, virus@X12819.1=Bluetongue_virus_gene_for_capsid_protein_VP1_genomic_RNA:1-19208

0 5000 10000 15000 20000

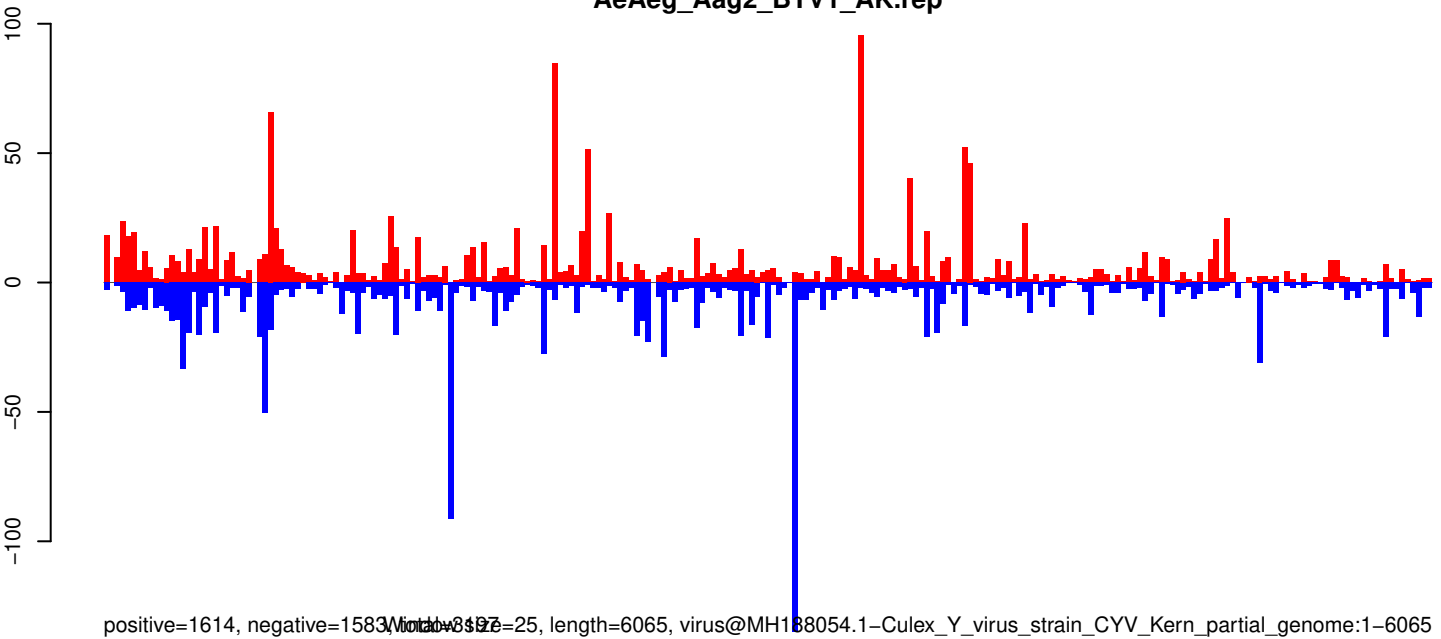
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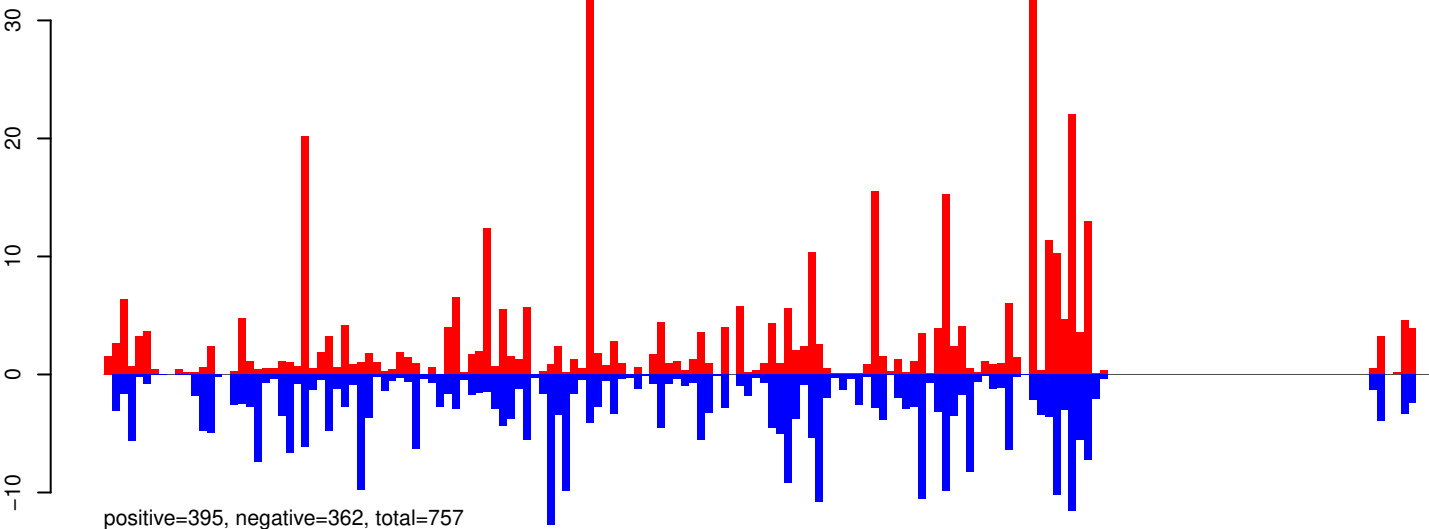
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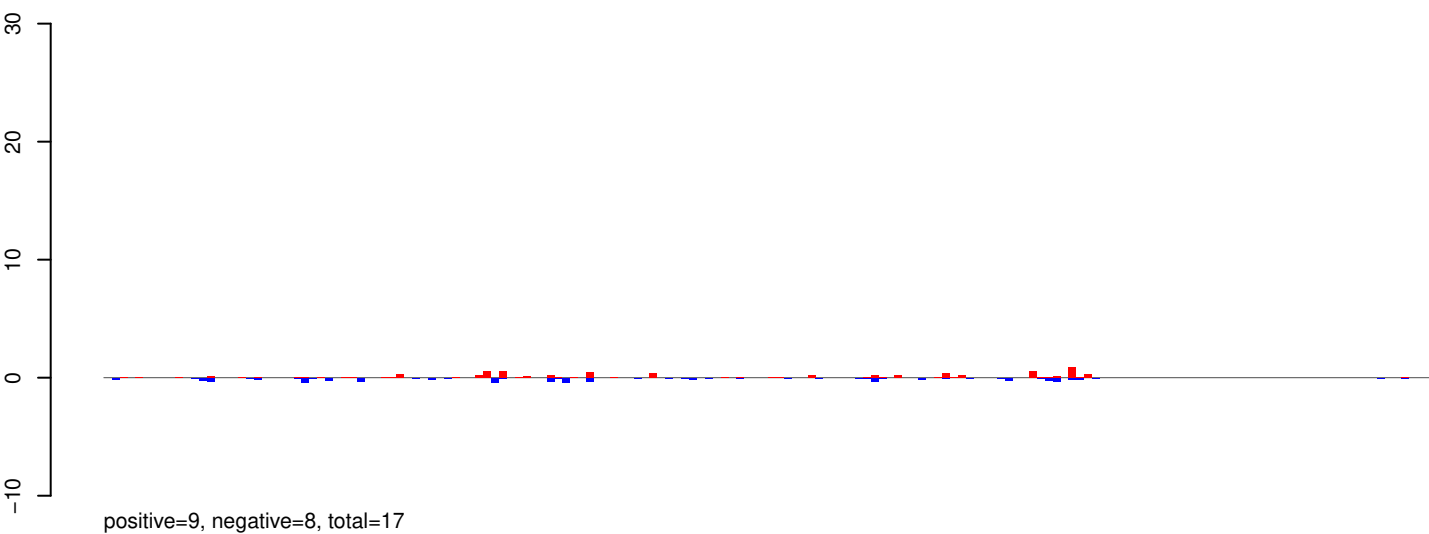
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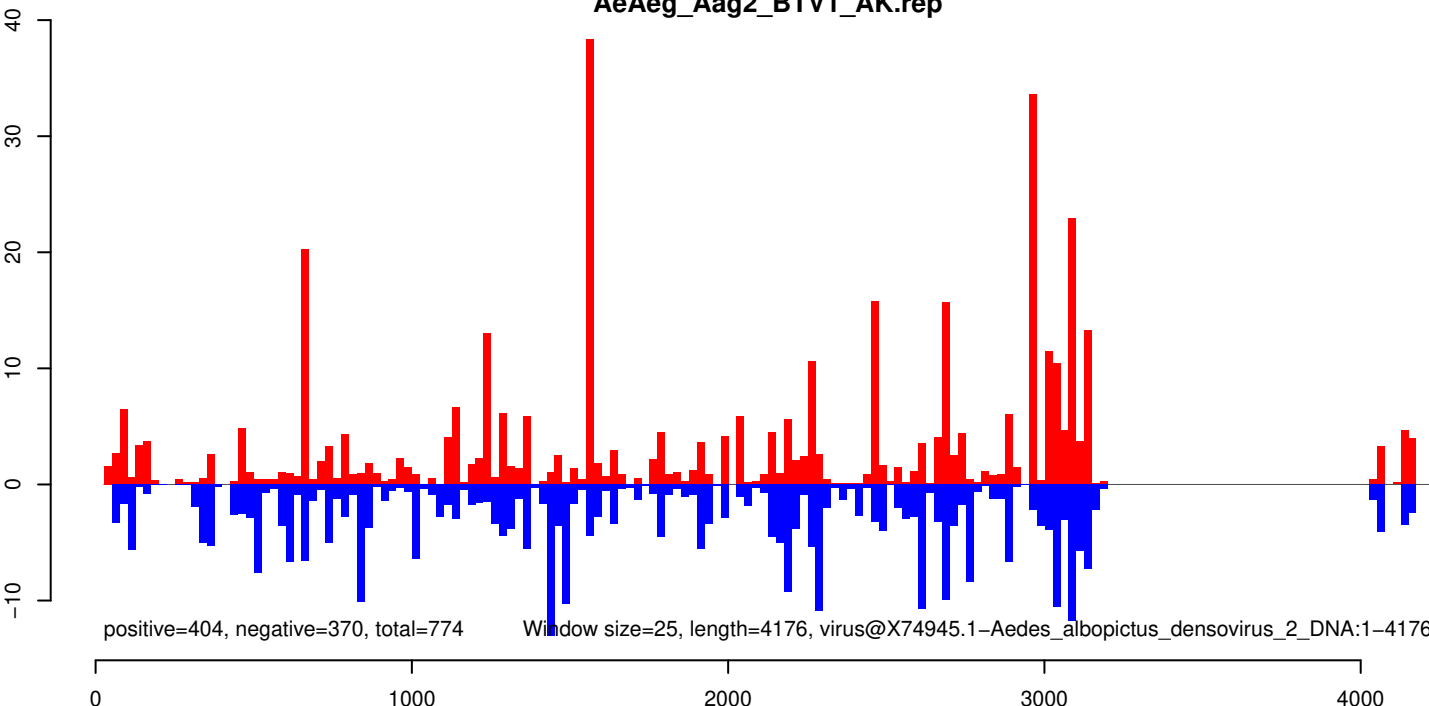
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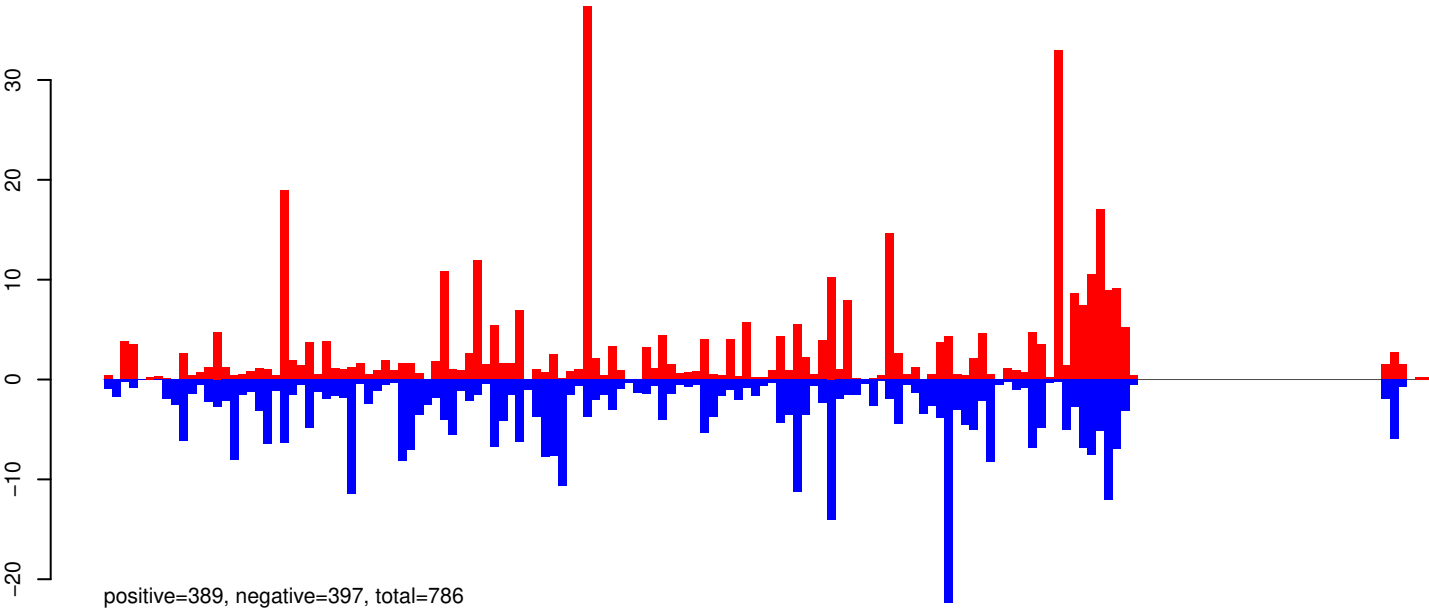
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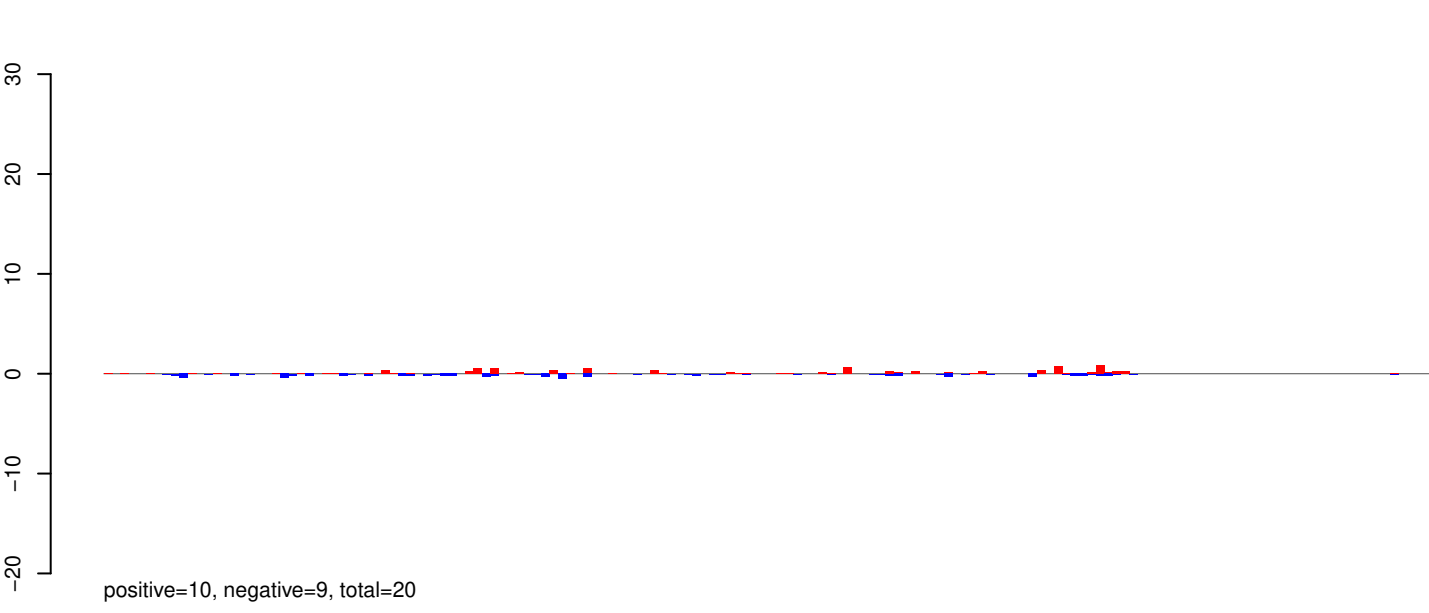
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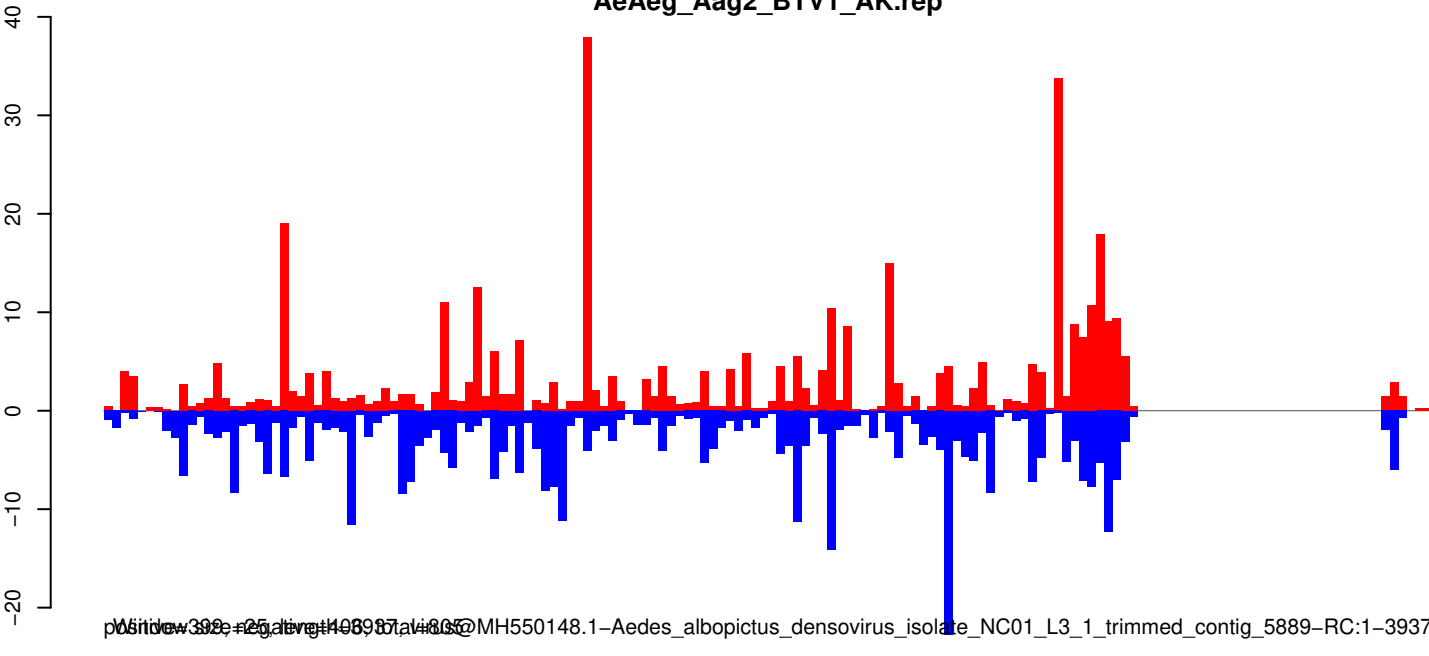
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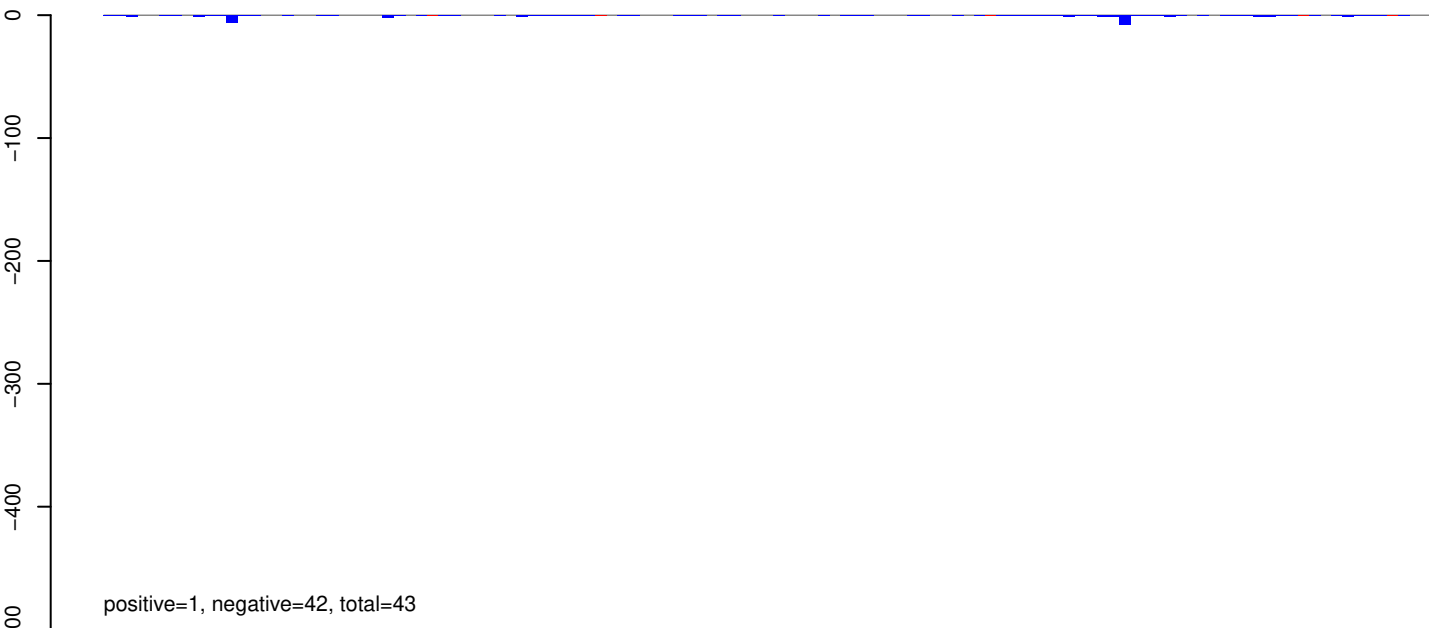
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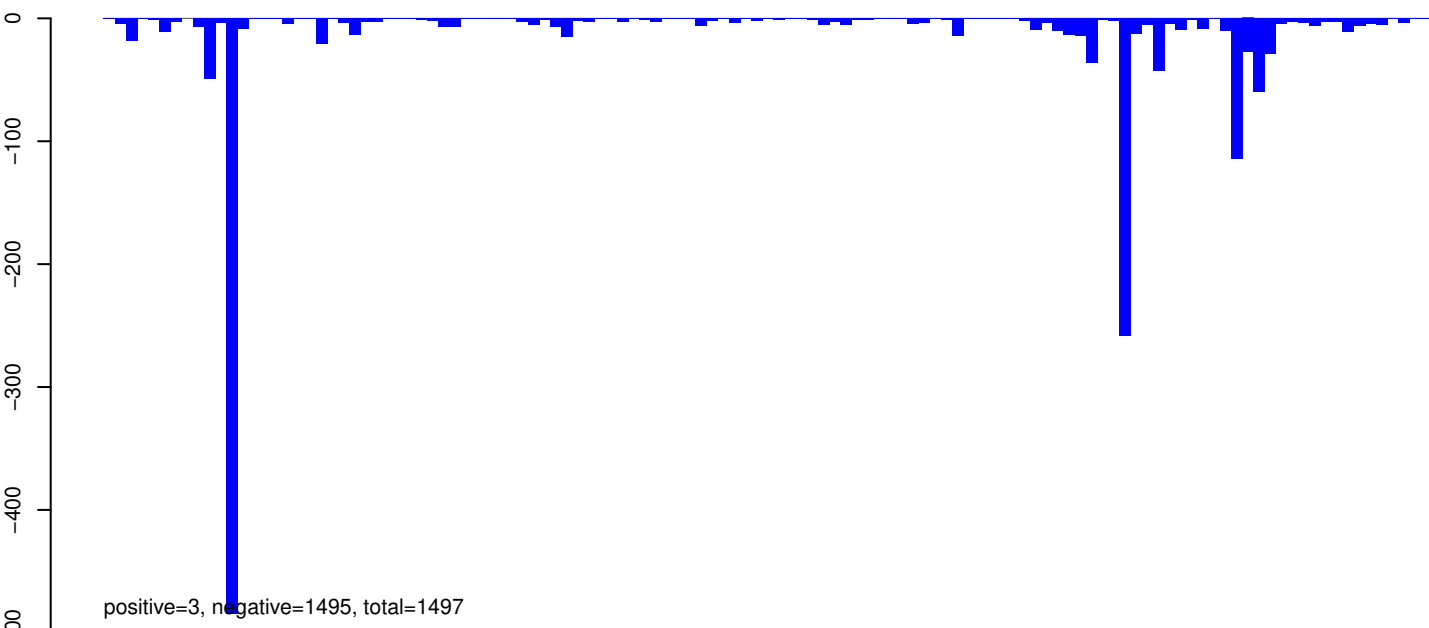
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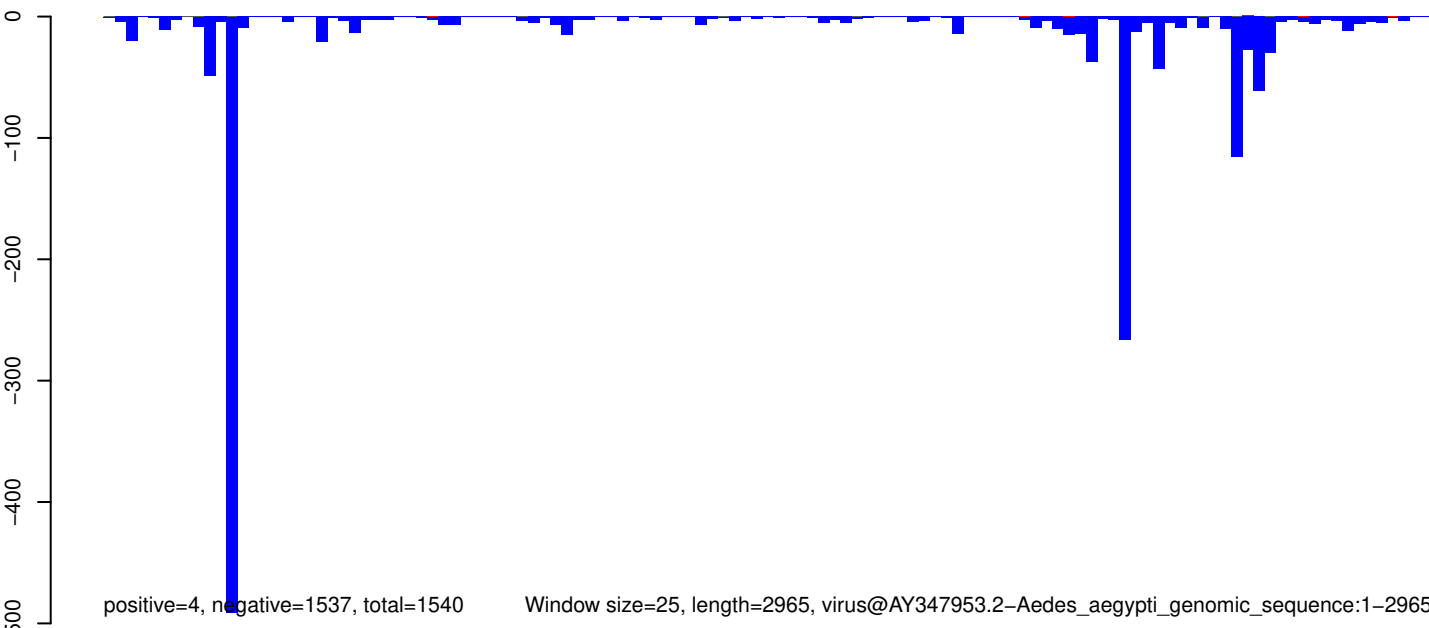
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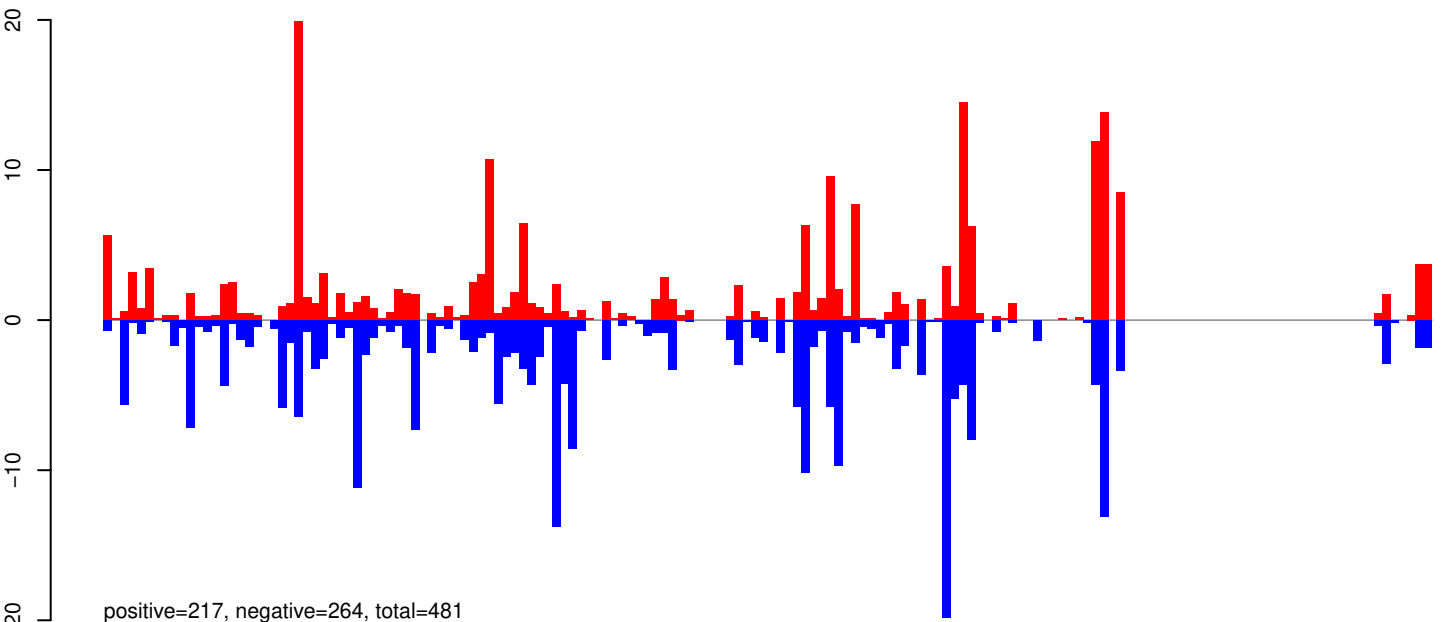
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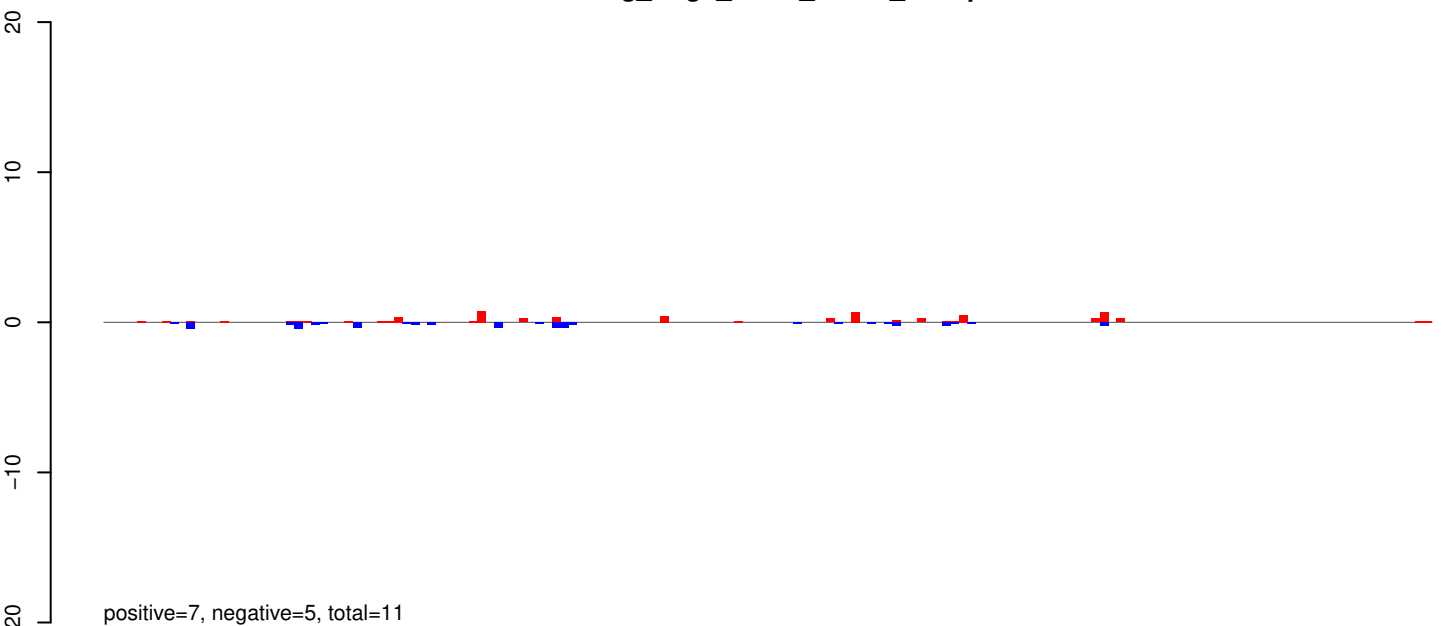
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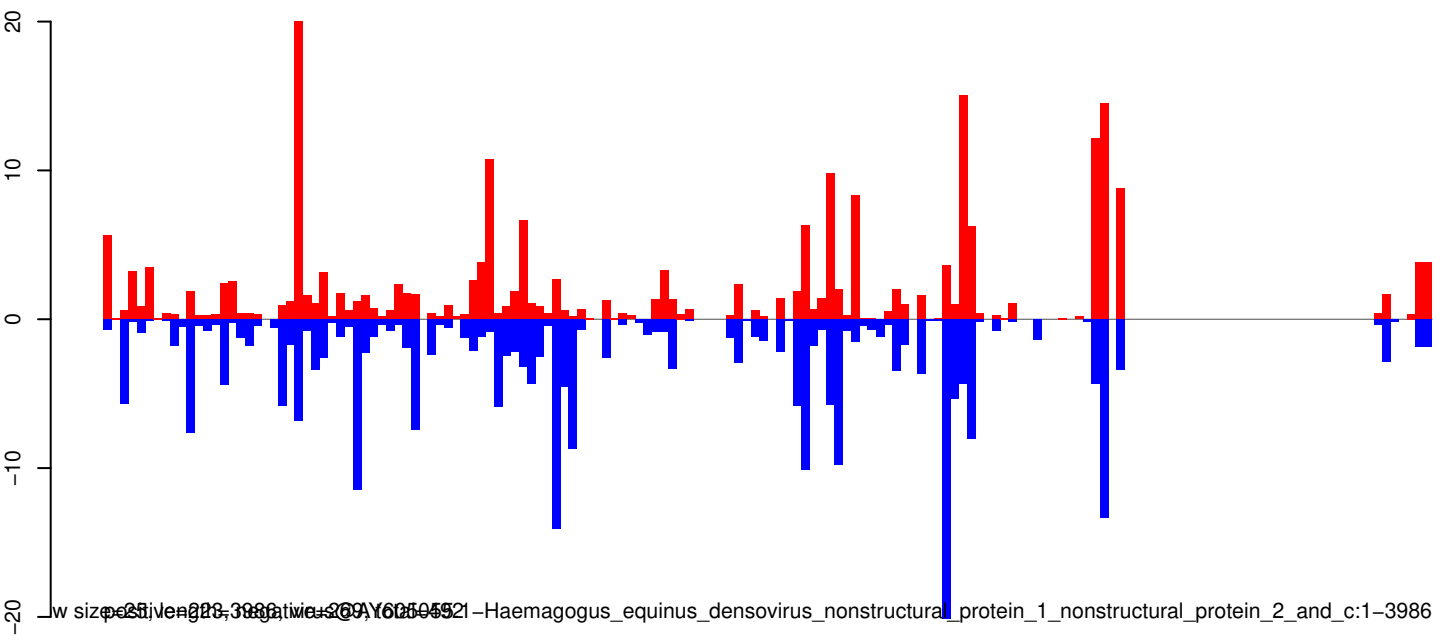
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AeAeg_Aag2_BTV1_AK.24_35.rep



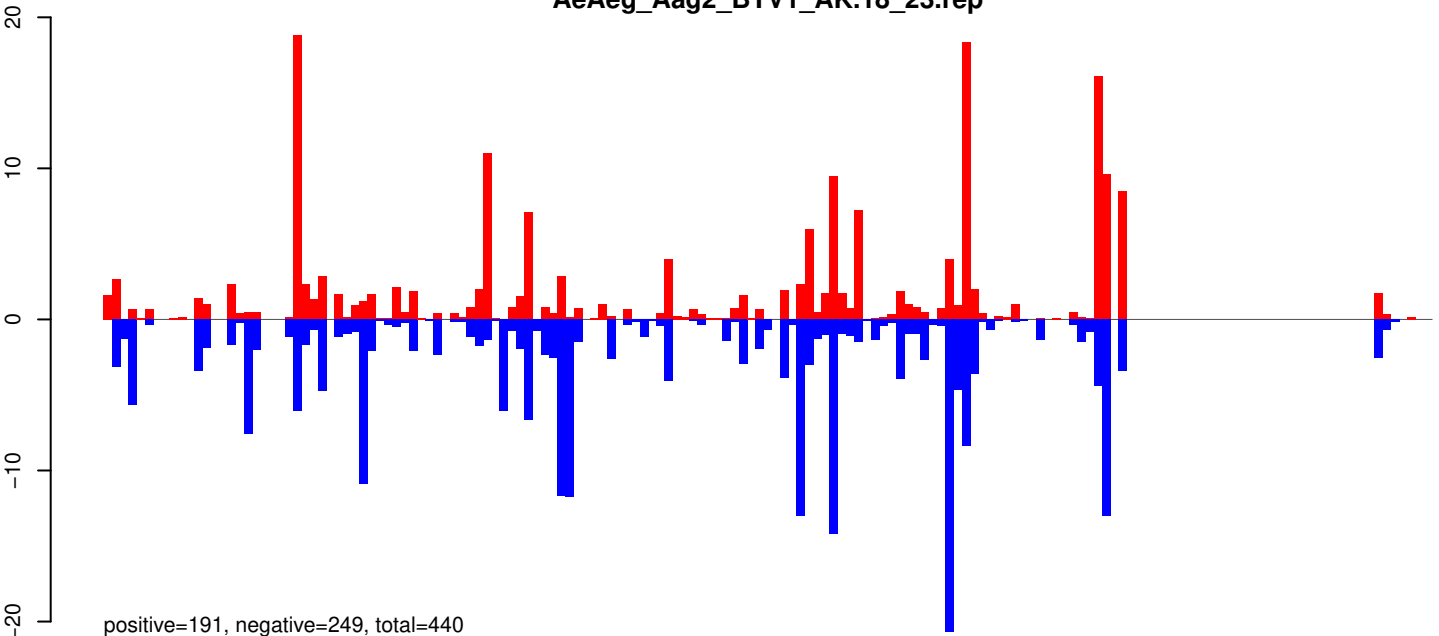
AeAeg_Aag2_BTV1_AK.rep



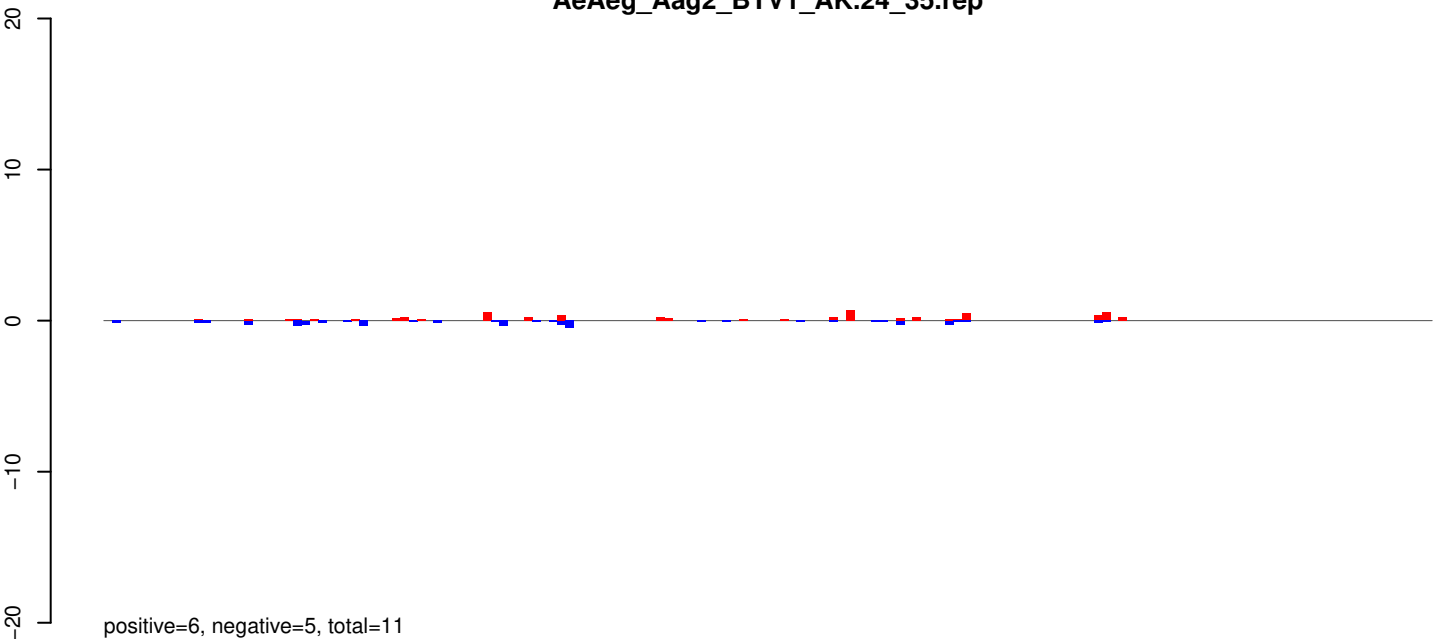
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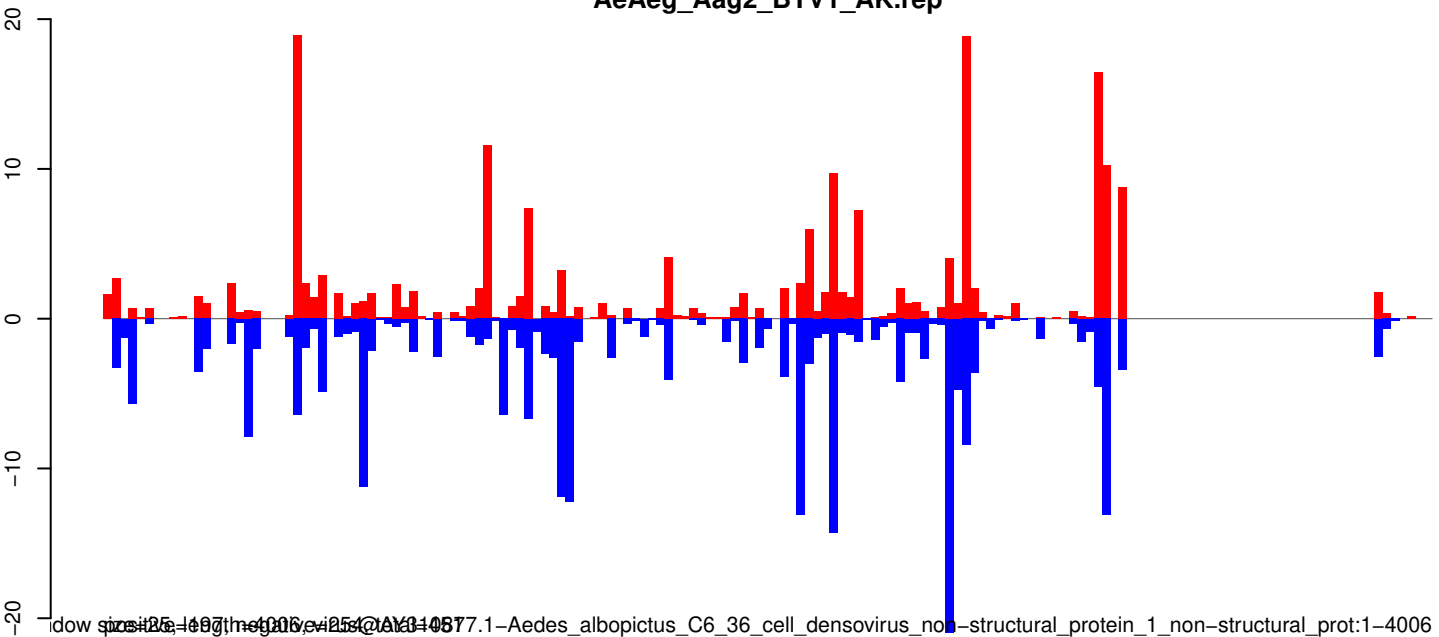
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AeAeg_Aag2_BTV1_AK.24_35.rep

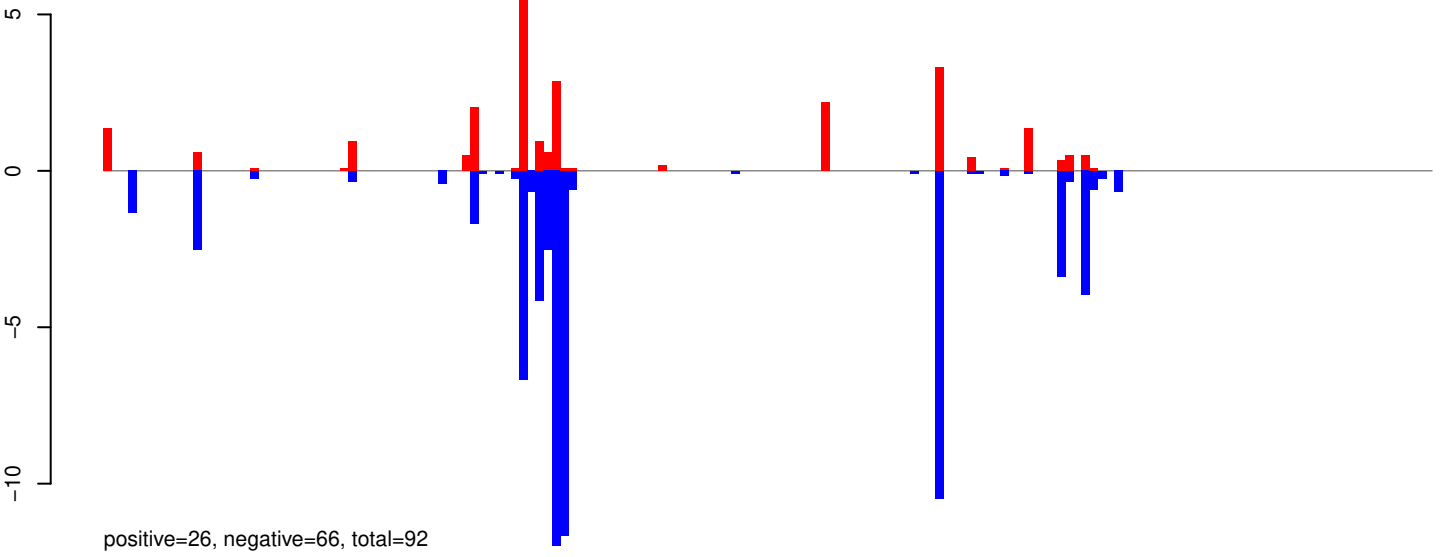


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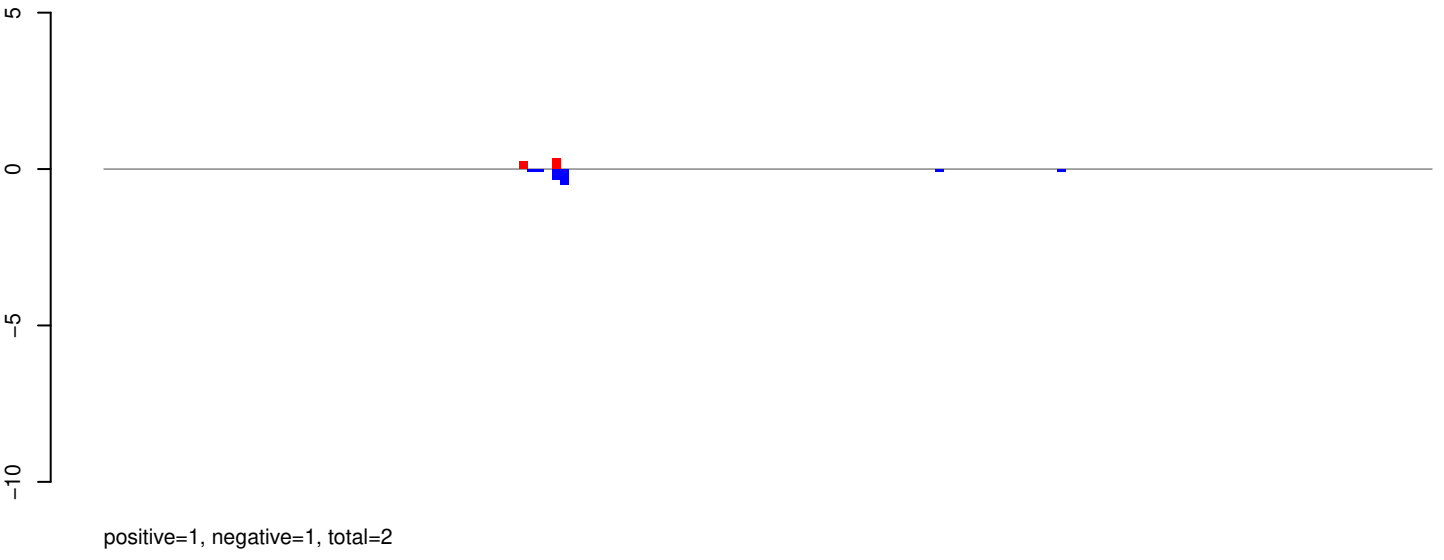


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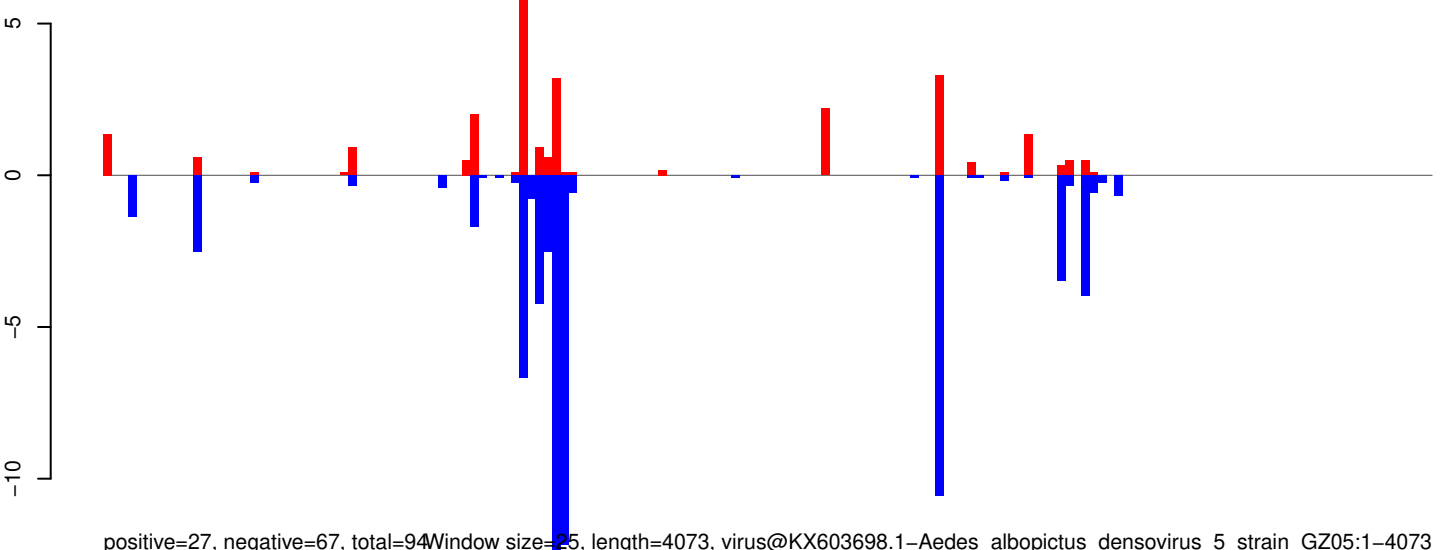
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AeAeg_Aag2_BTV1_AK.24_35.rep



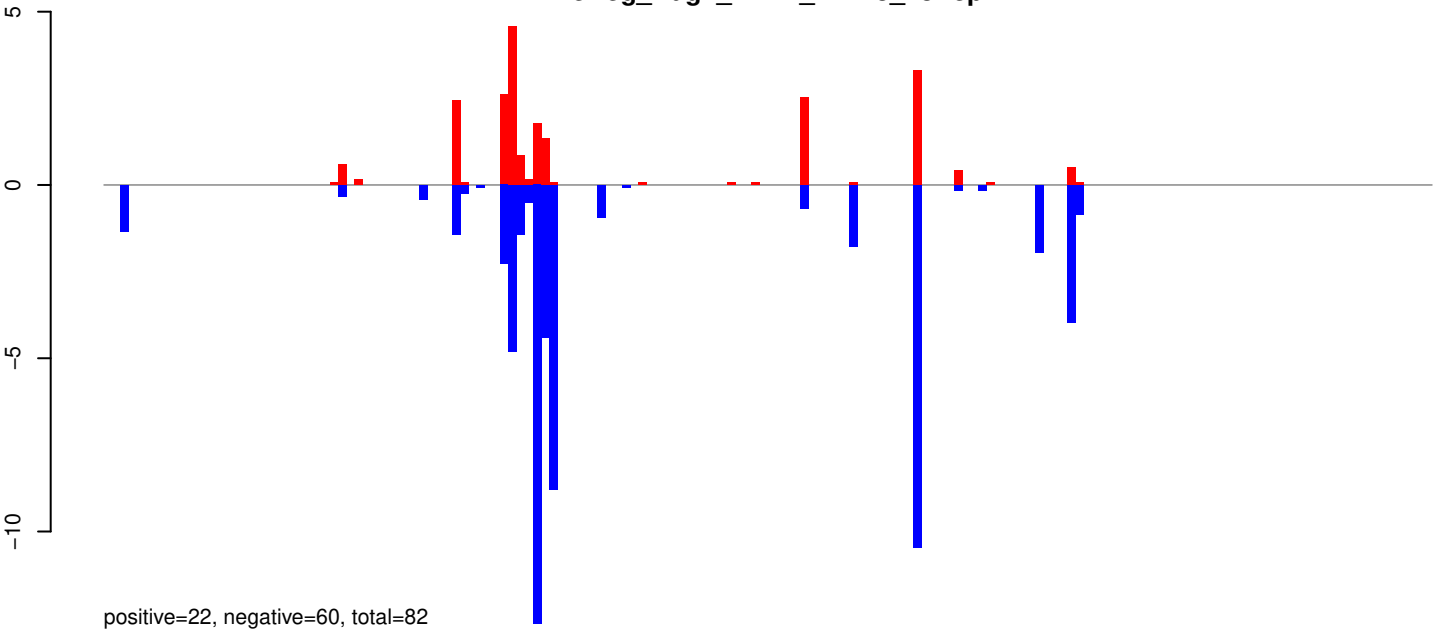
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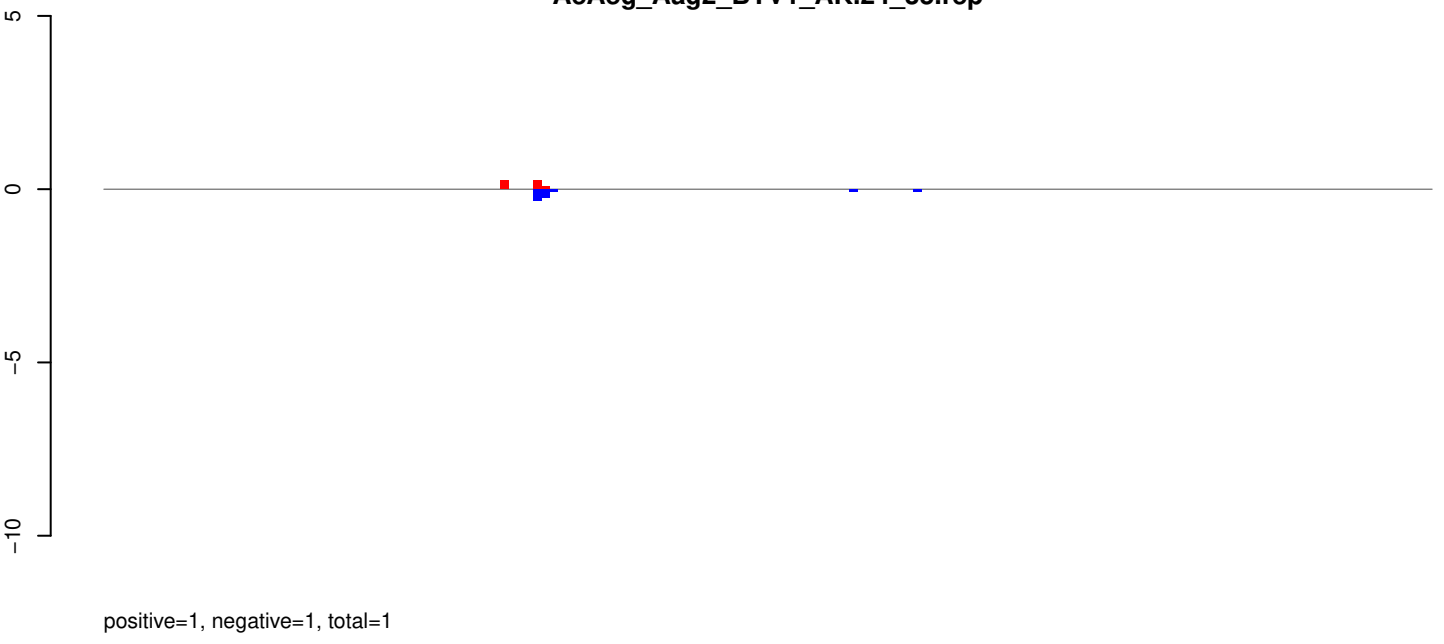
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0 1000 2000 3000 4000

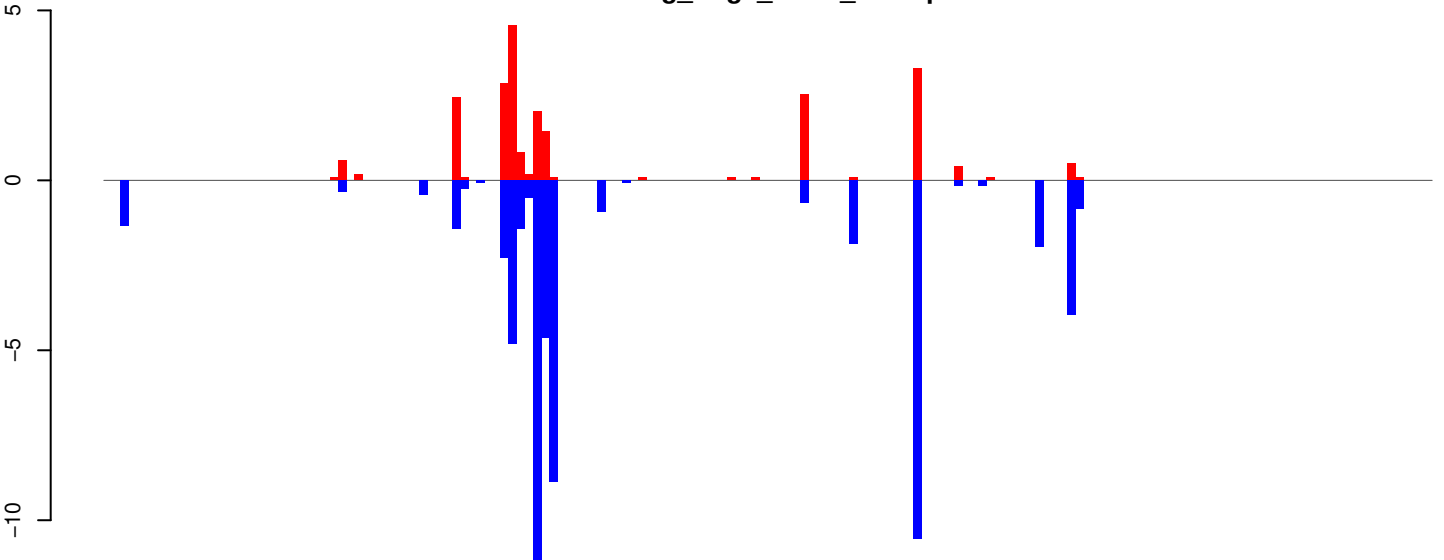
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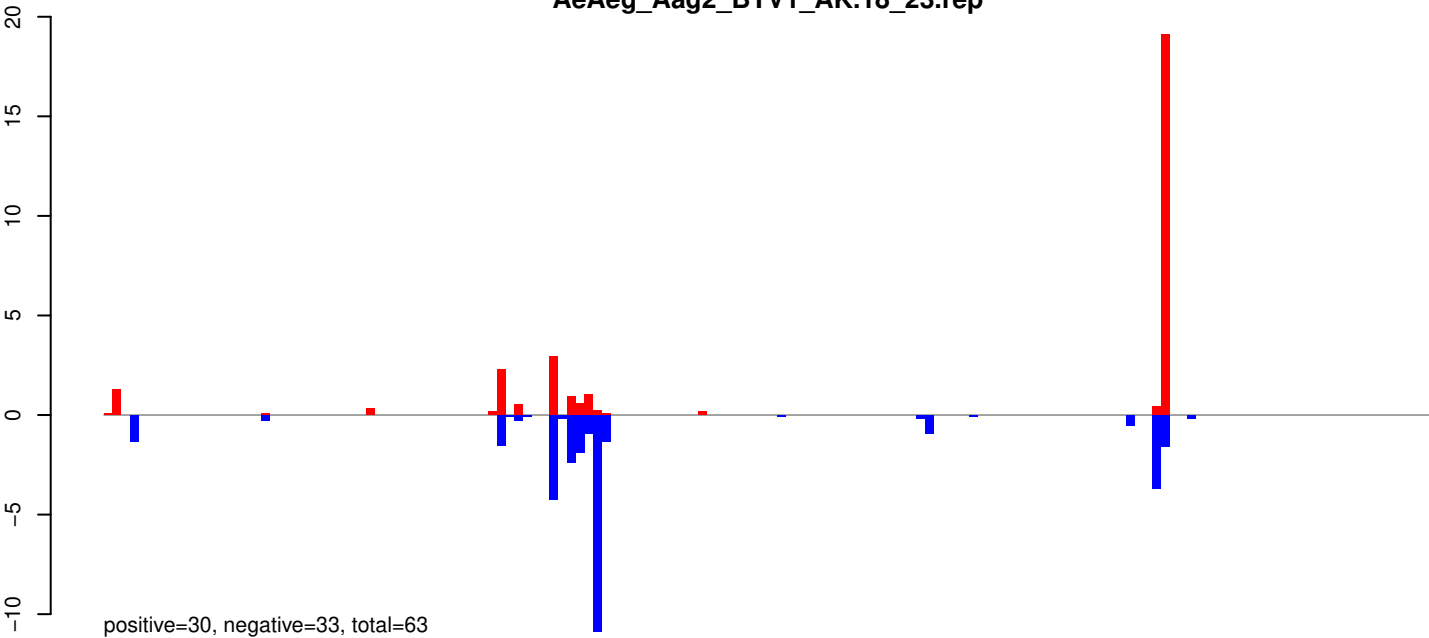
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AeAeg_Aag2_BTV1_AK.rep



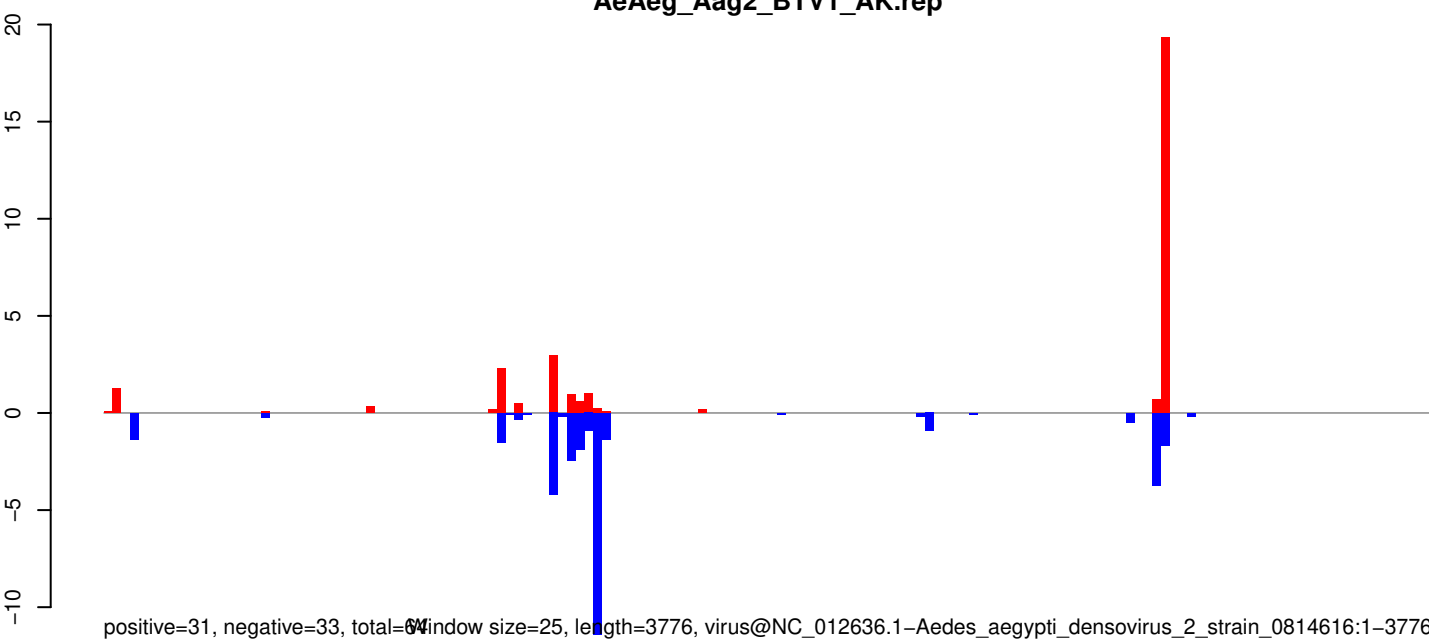
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AeAeg_Aag2_BTV1_AK.24_35.rep

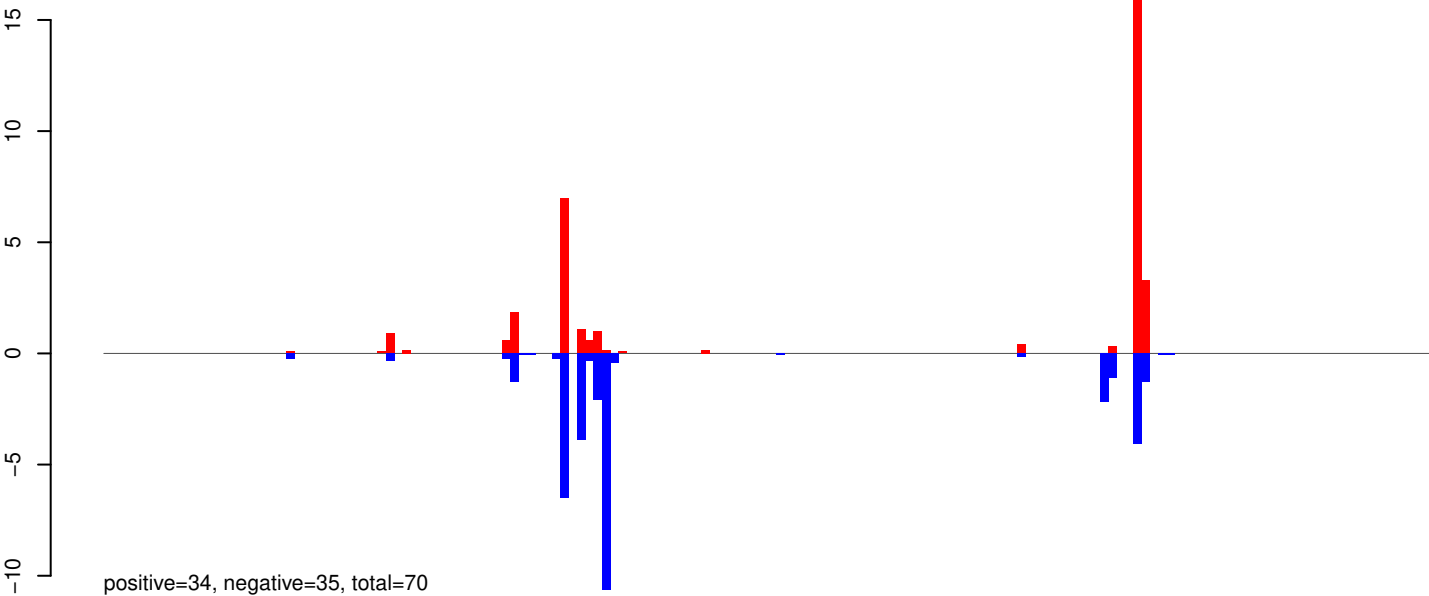


AeAeg_Aag2_BTV1_AK.rep

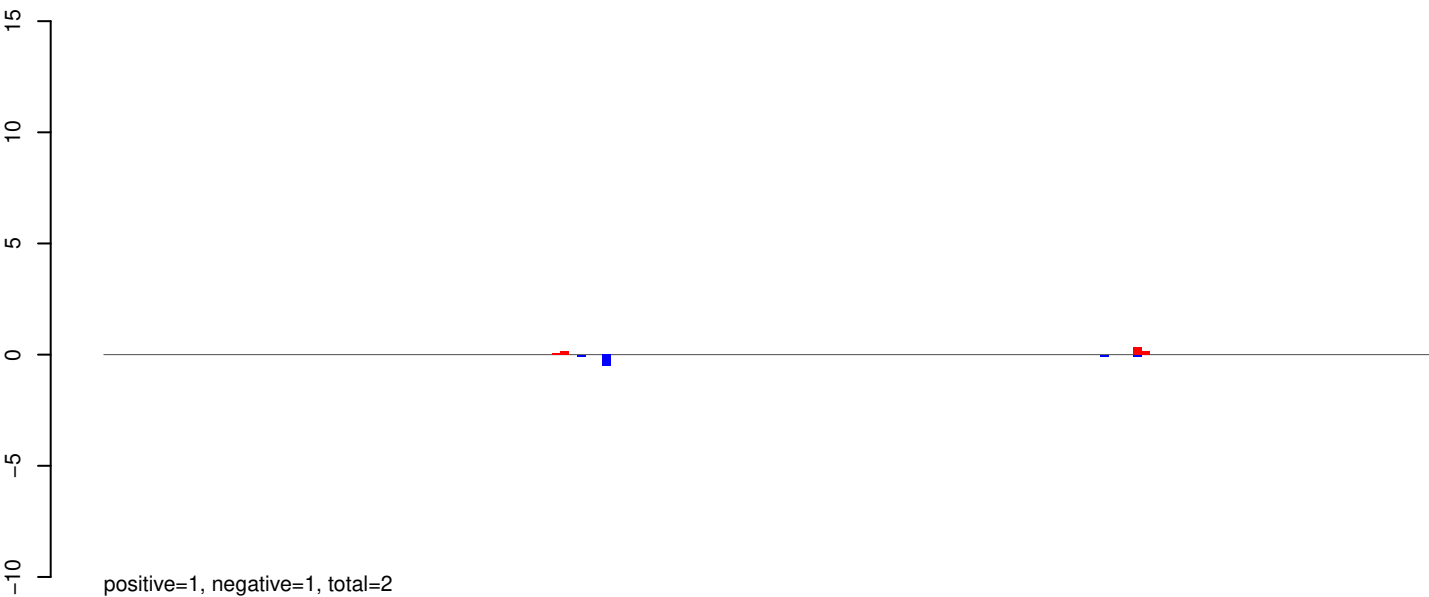


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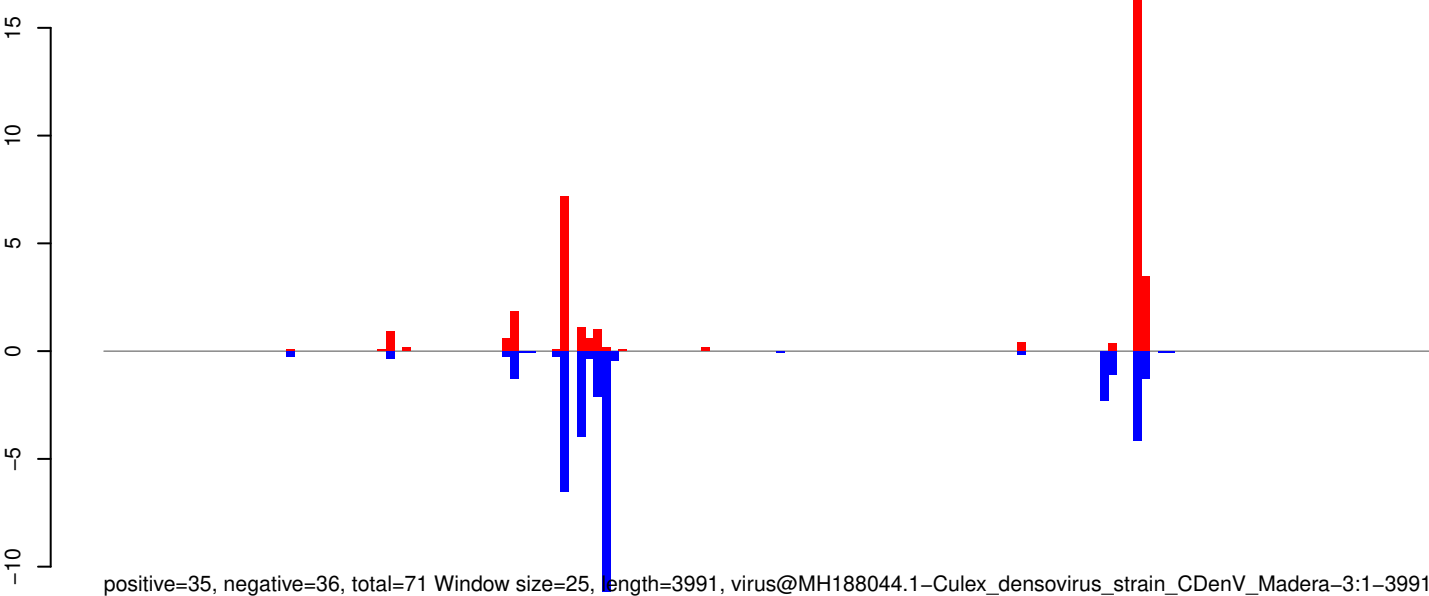
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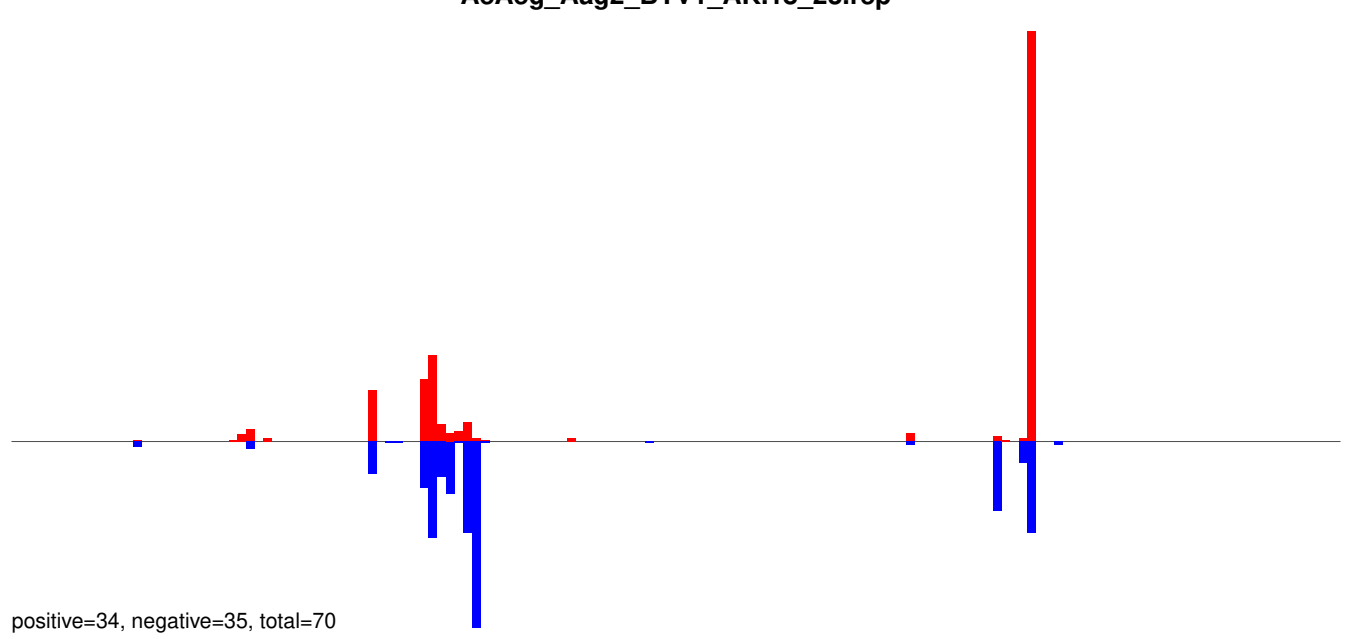
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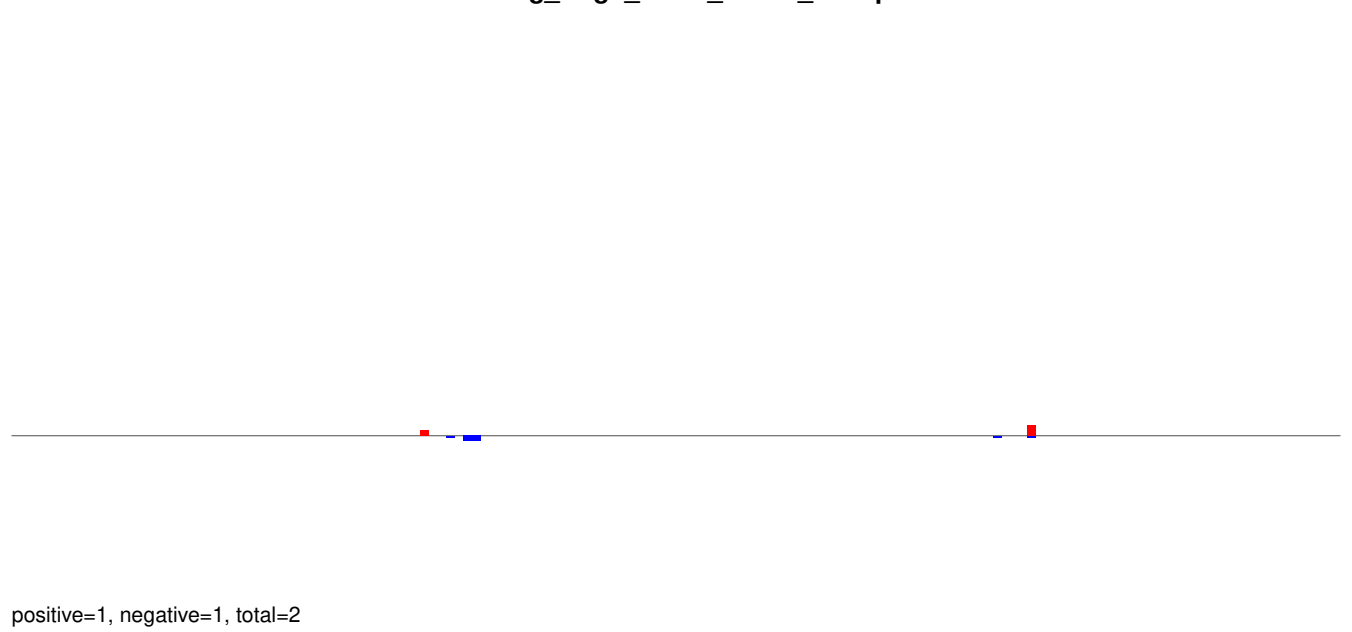
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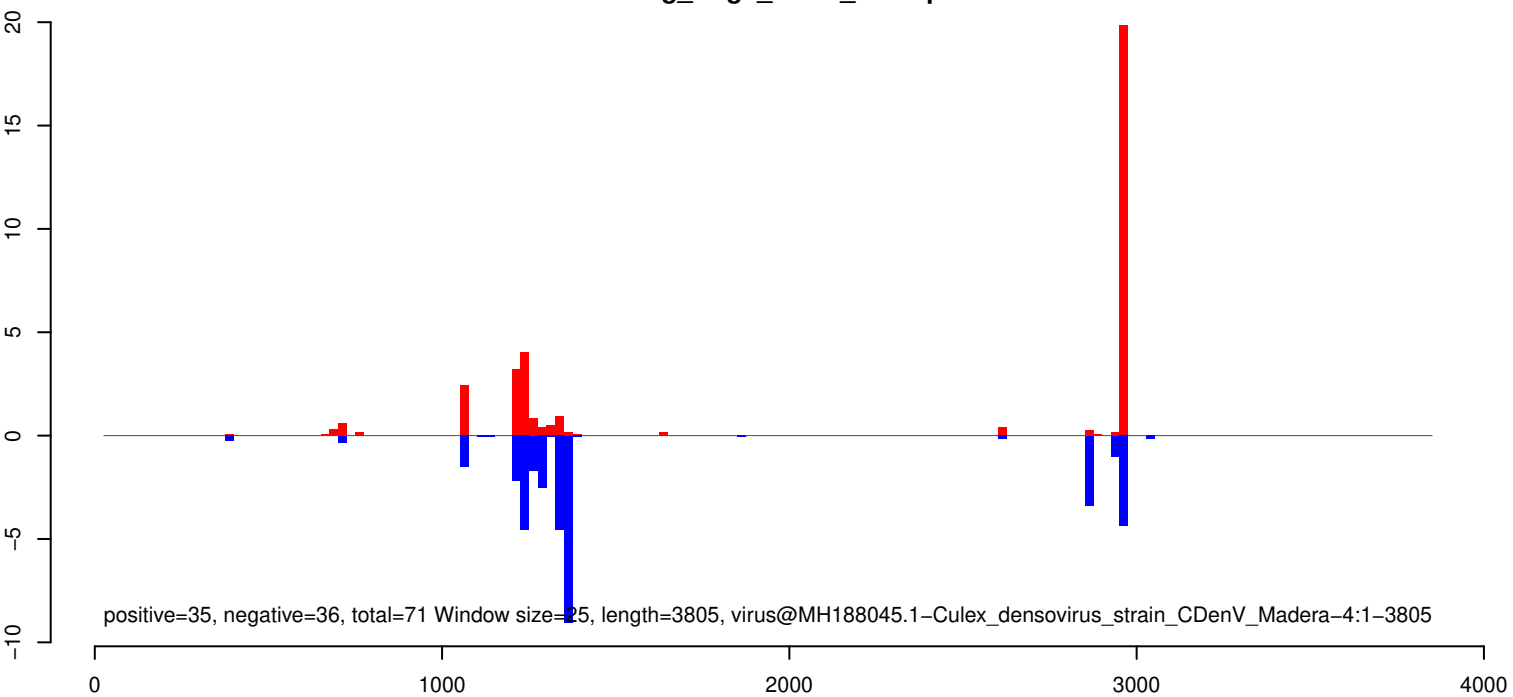
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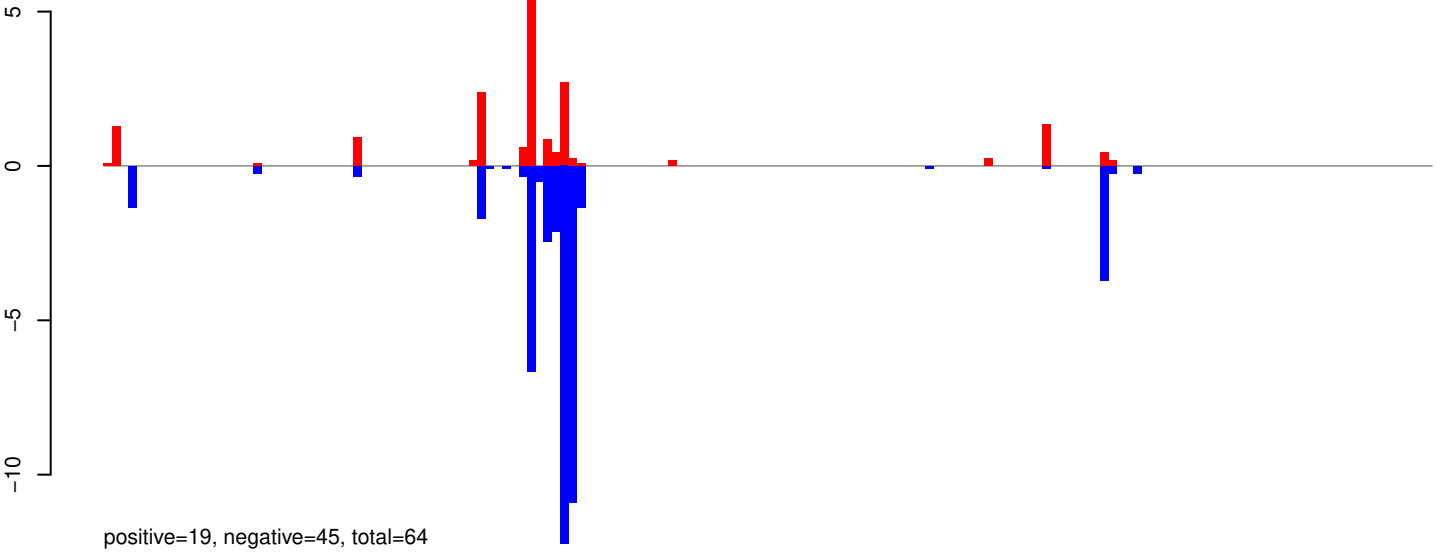
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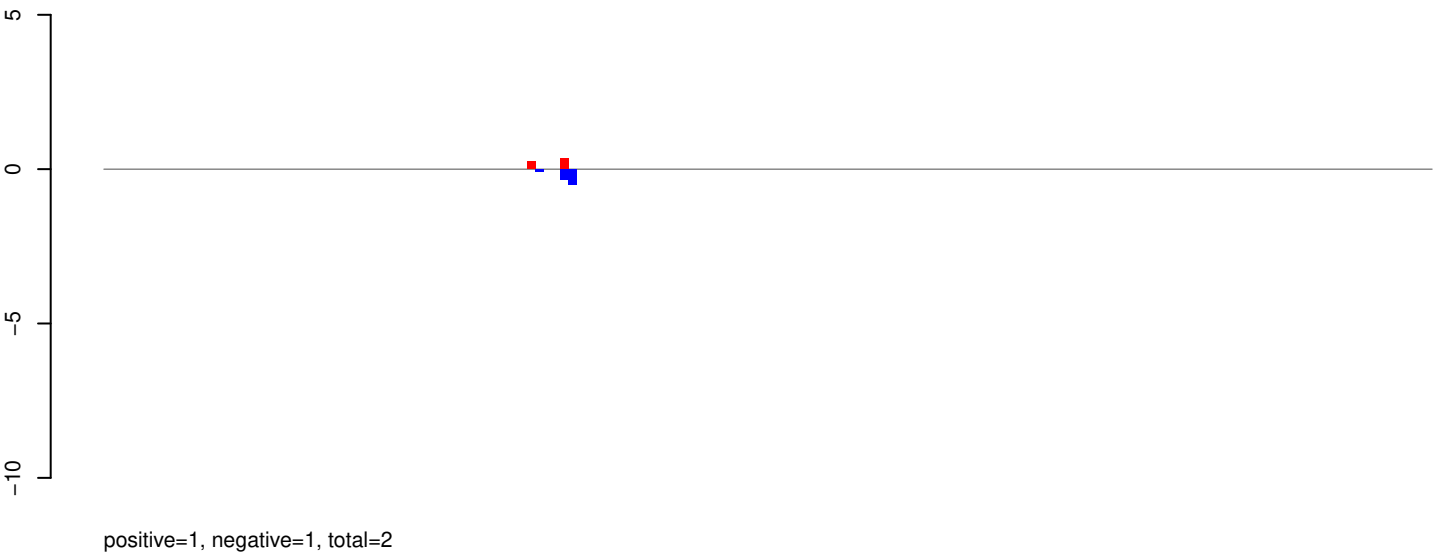
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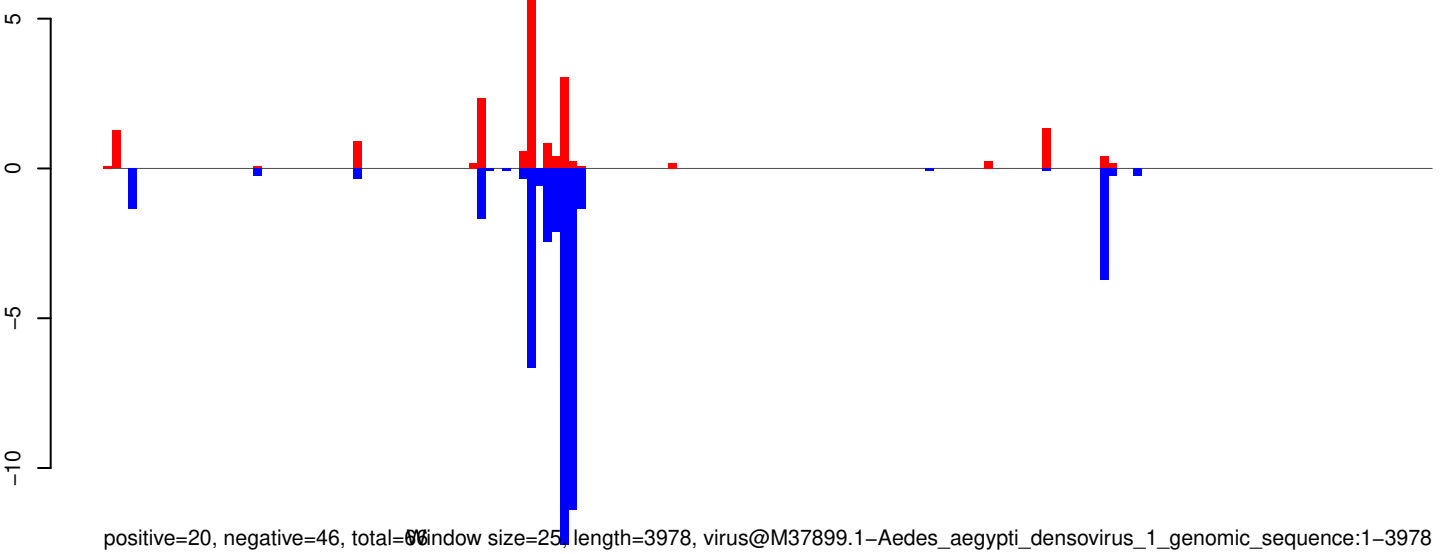
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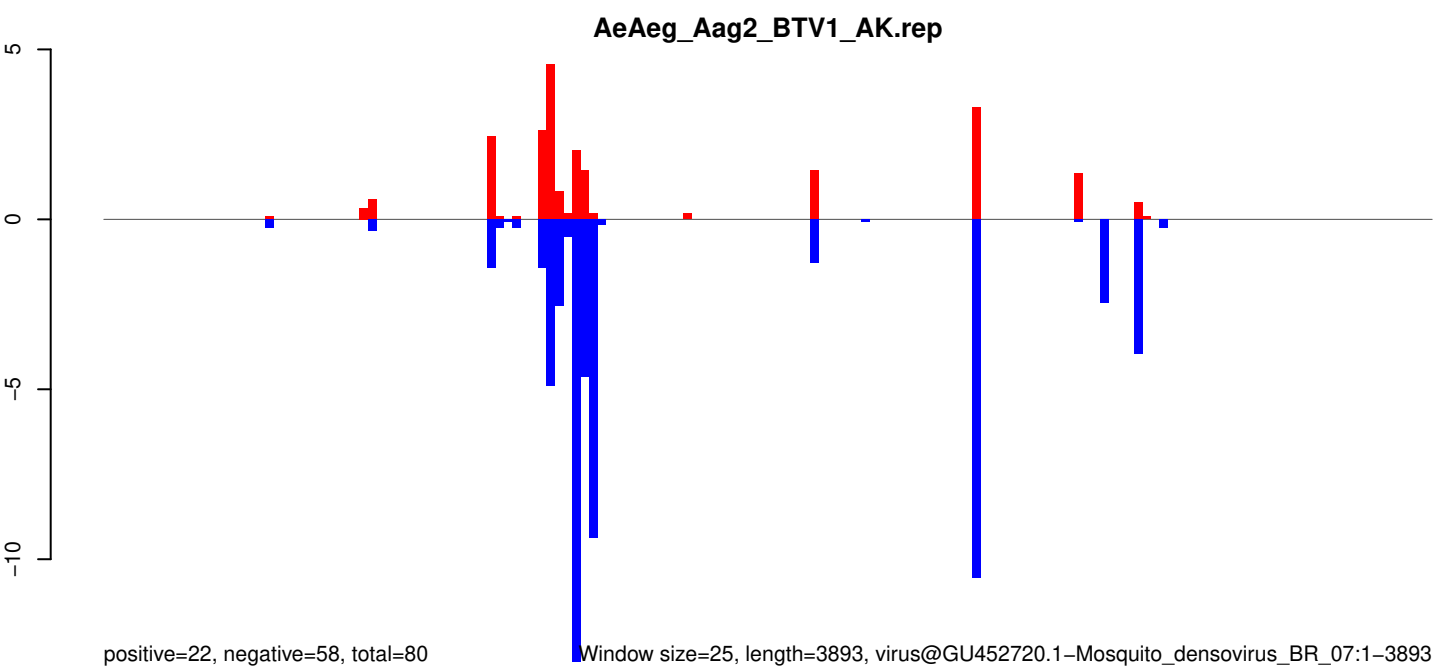
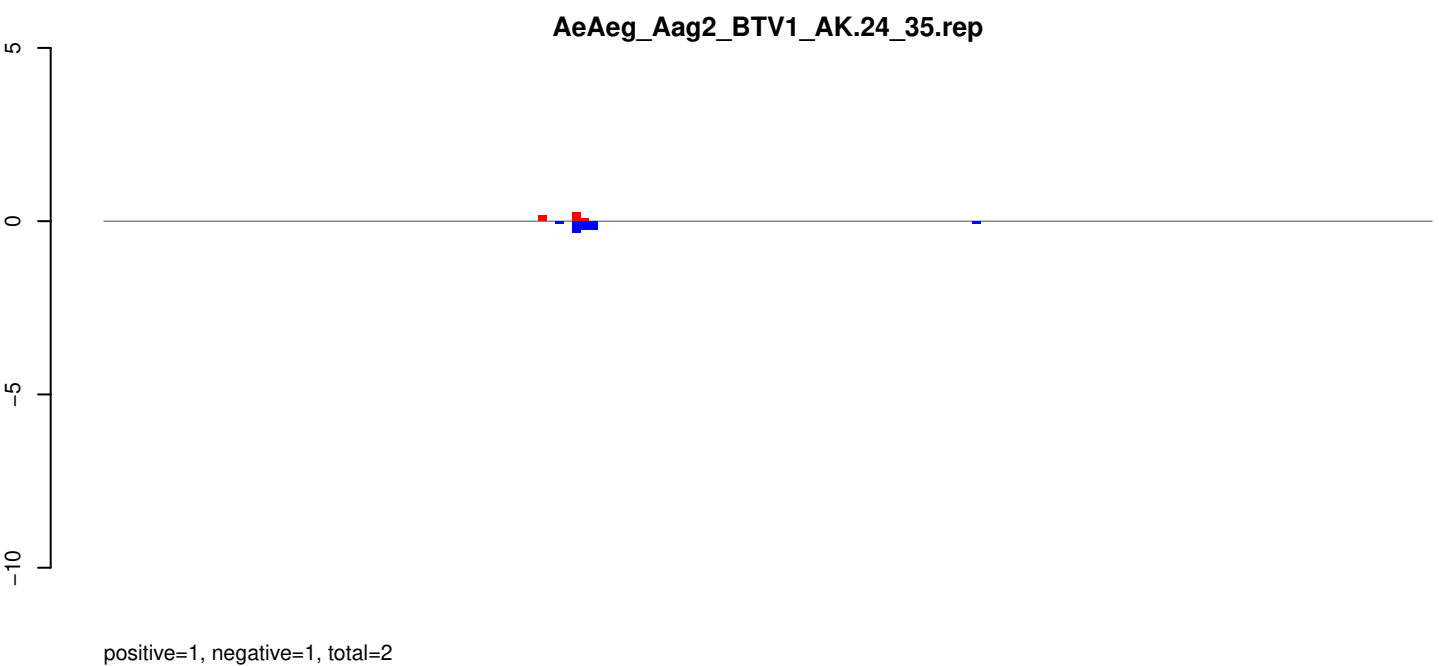
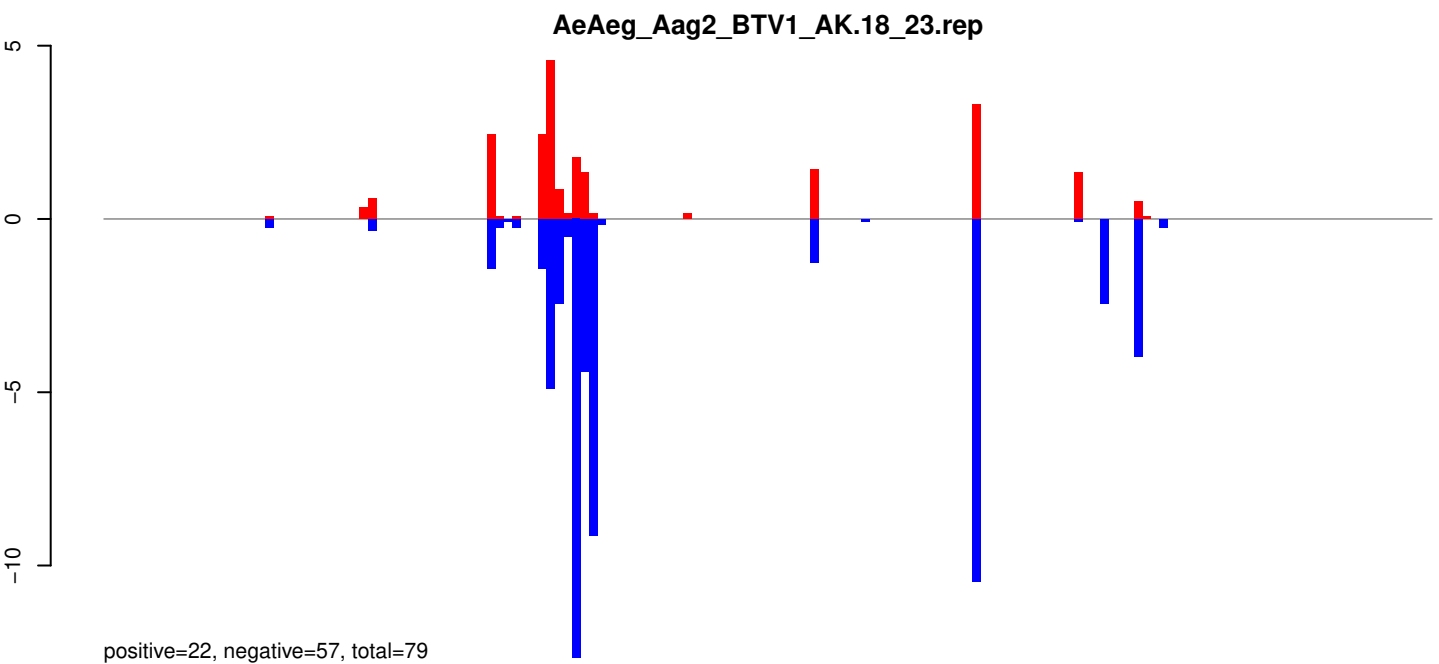
AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep

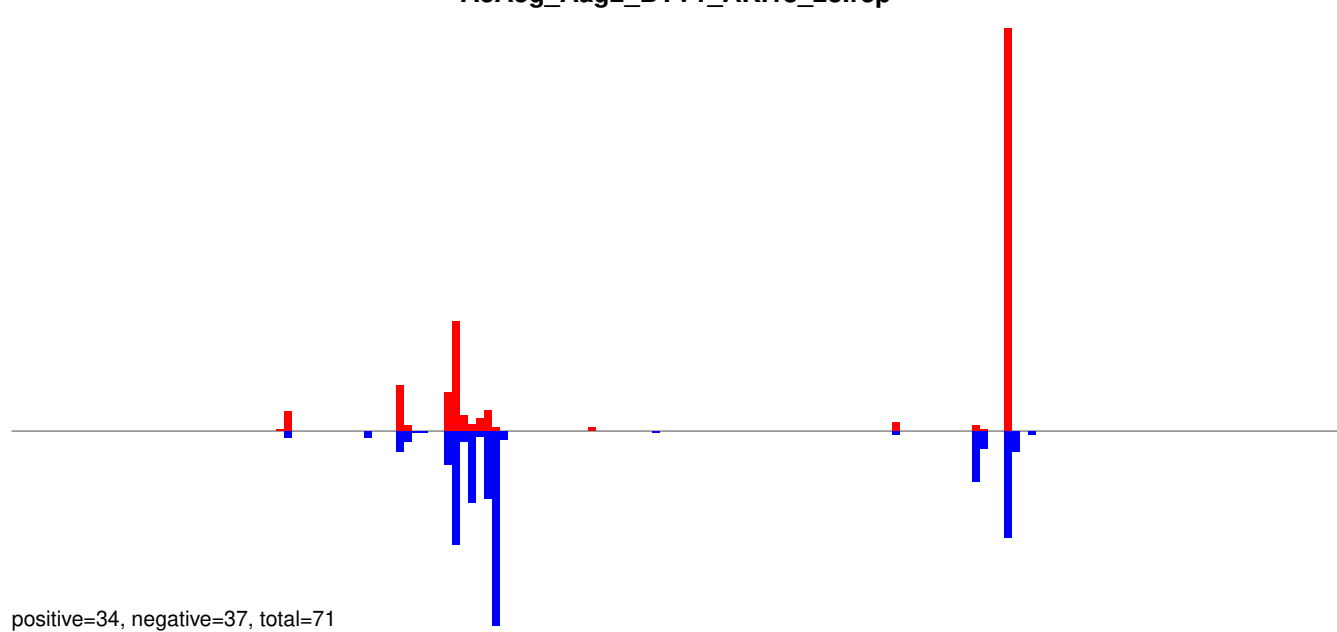


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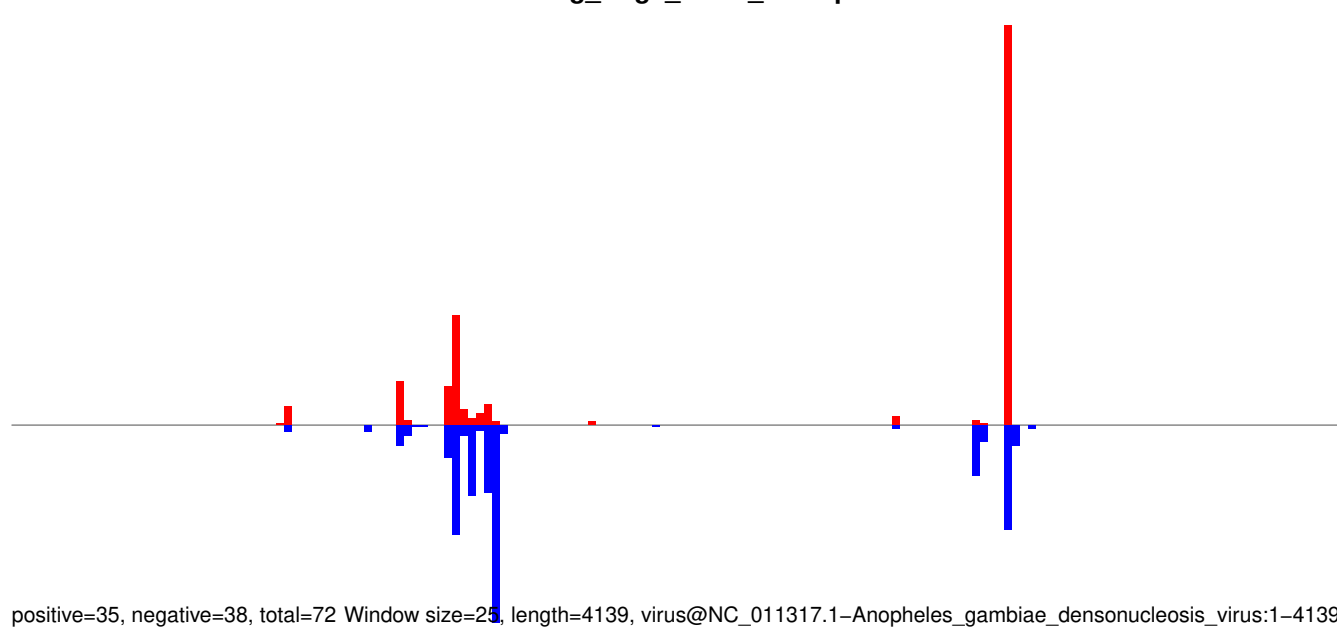
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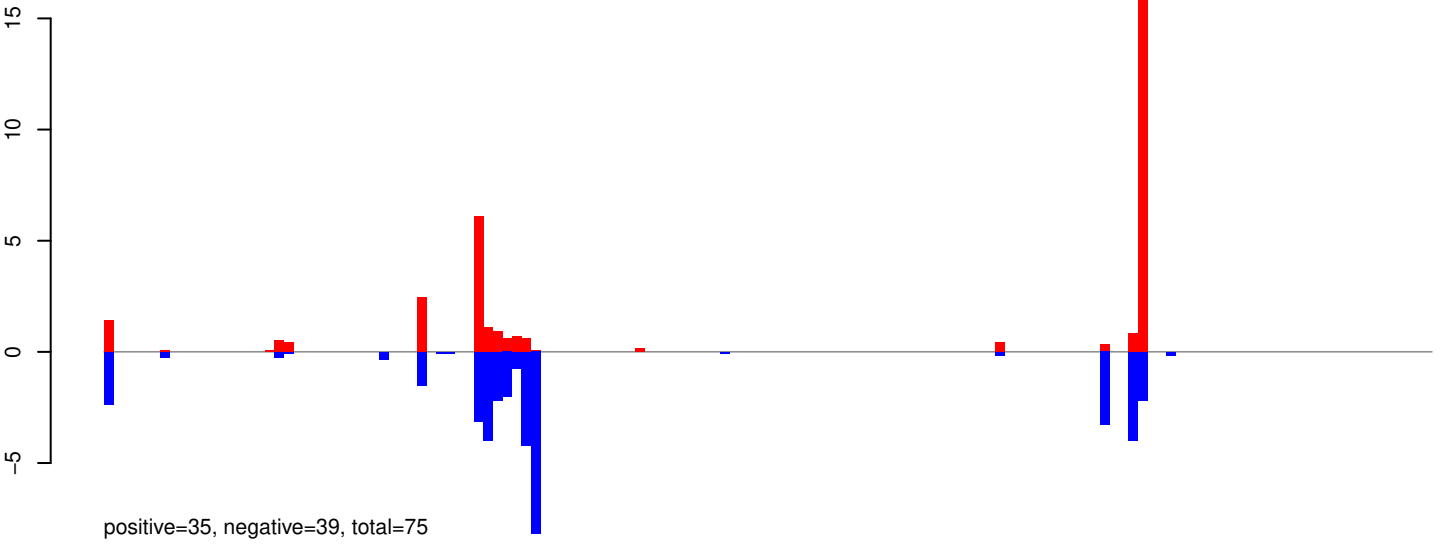
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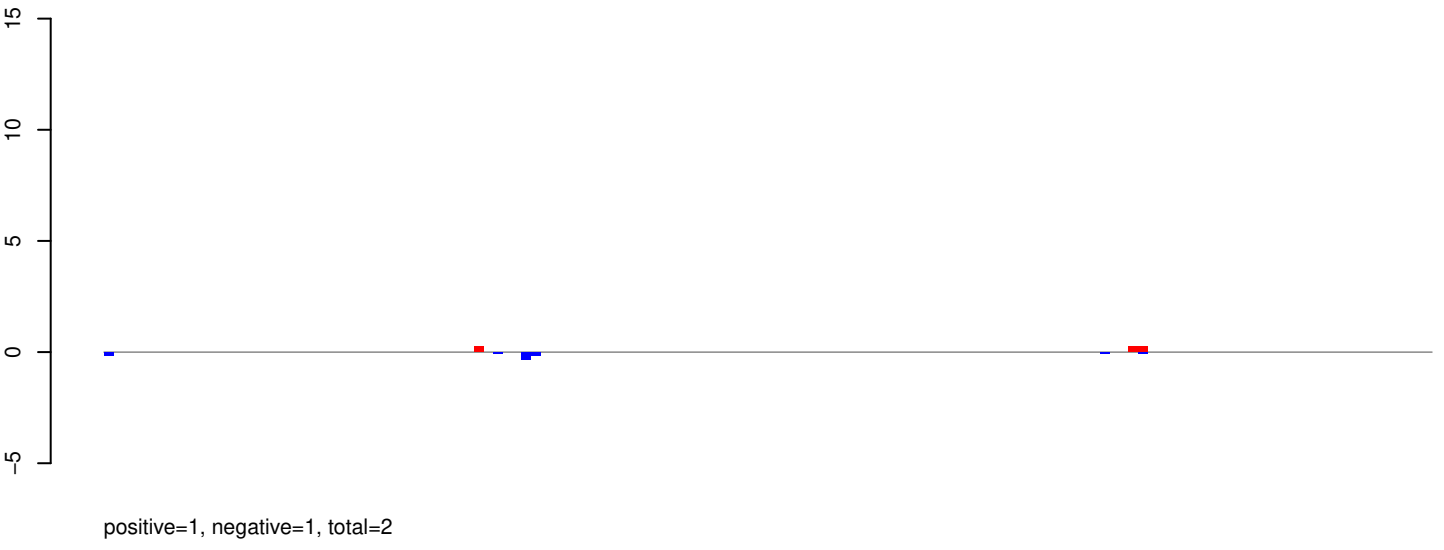
AeAeg_Aag2_BTV1_AK.rep



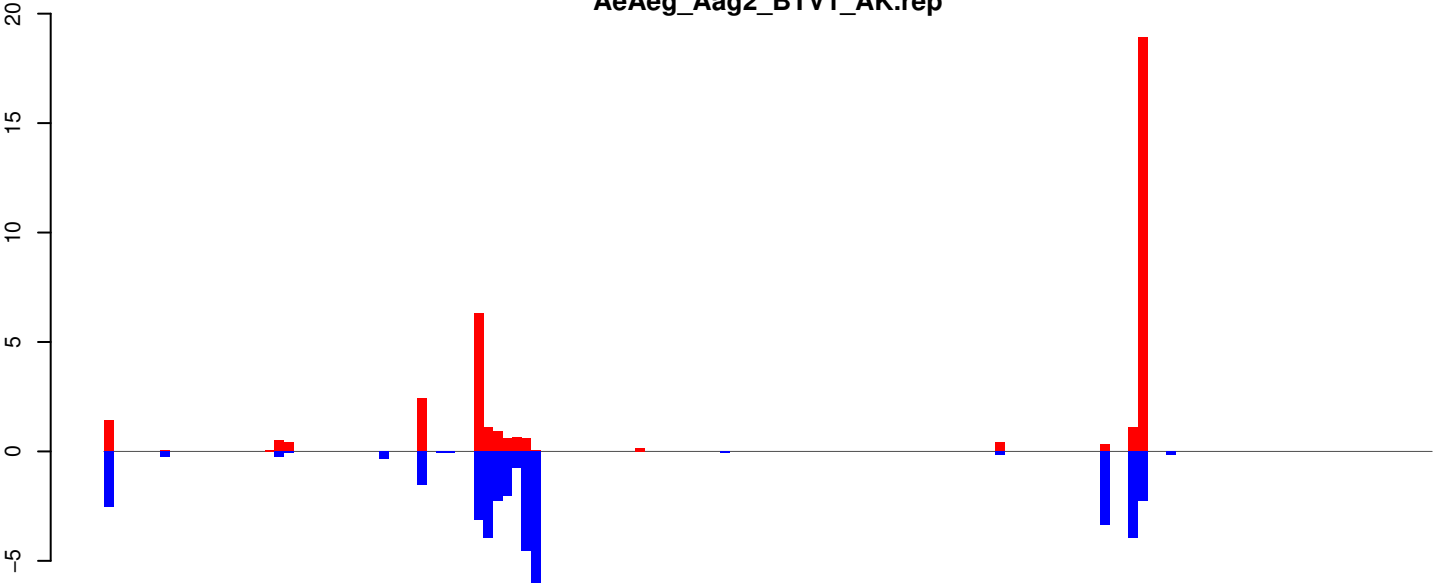
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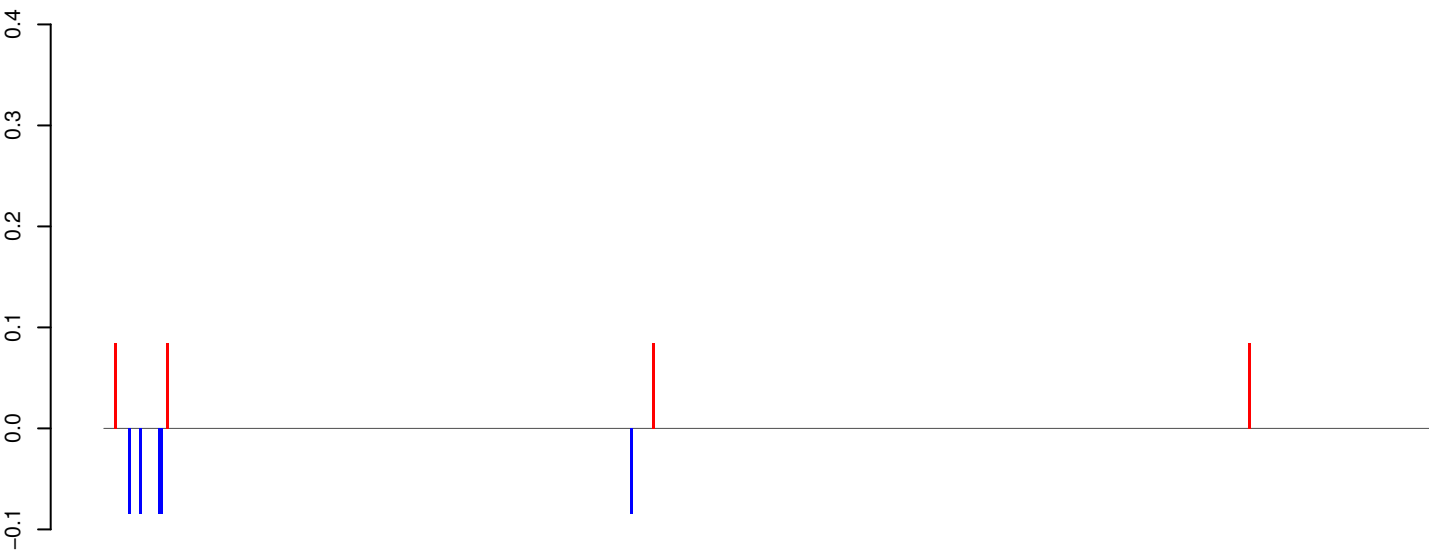
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AeAeg_Aag2_BTV1_AK.rep

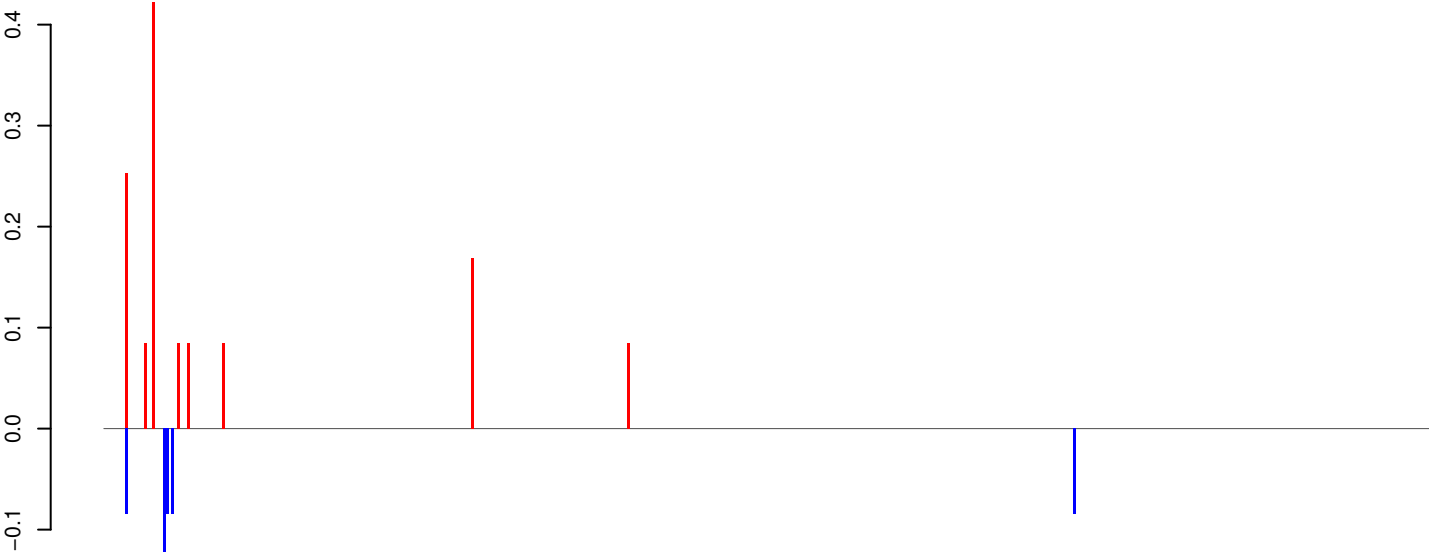


AeAeg_Aag2_BTV1_AK.18_23.rep



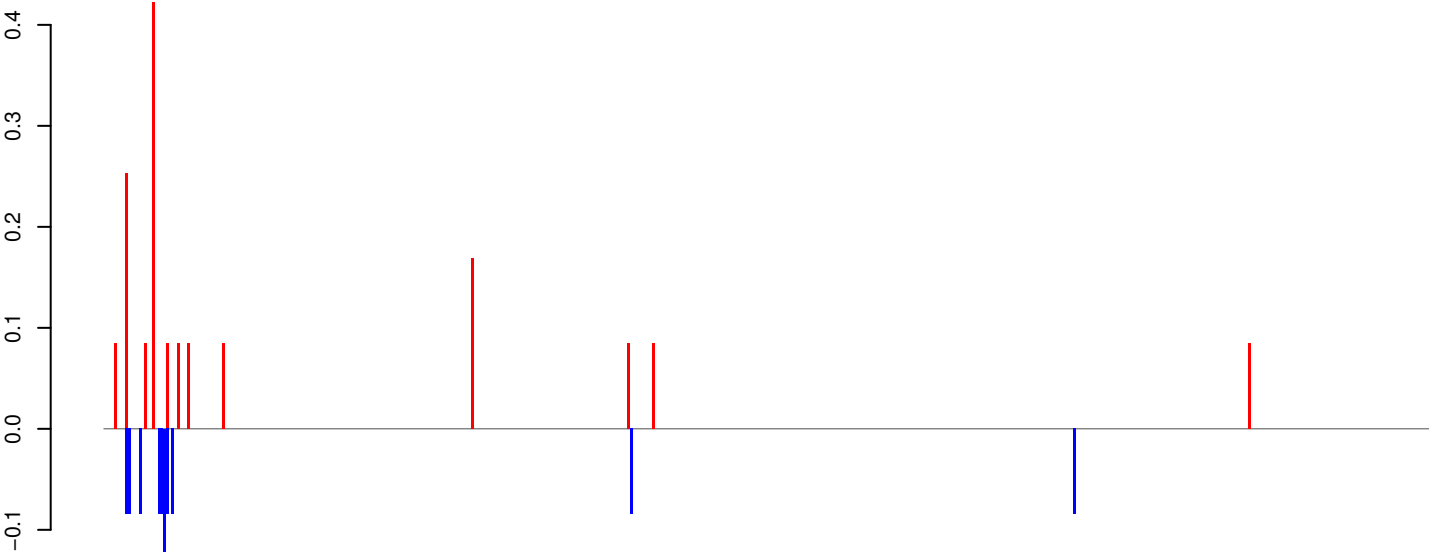
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AeAeg_Aag2_BTV1_AK.24_35.rep



positive=1, negative=1, total=2

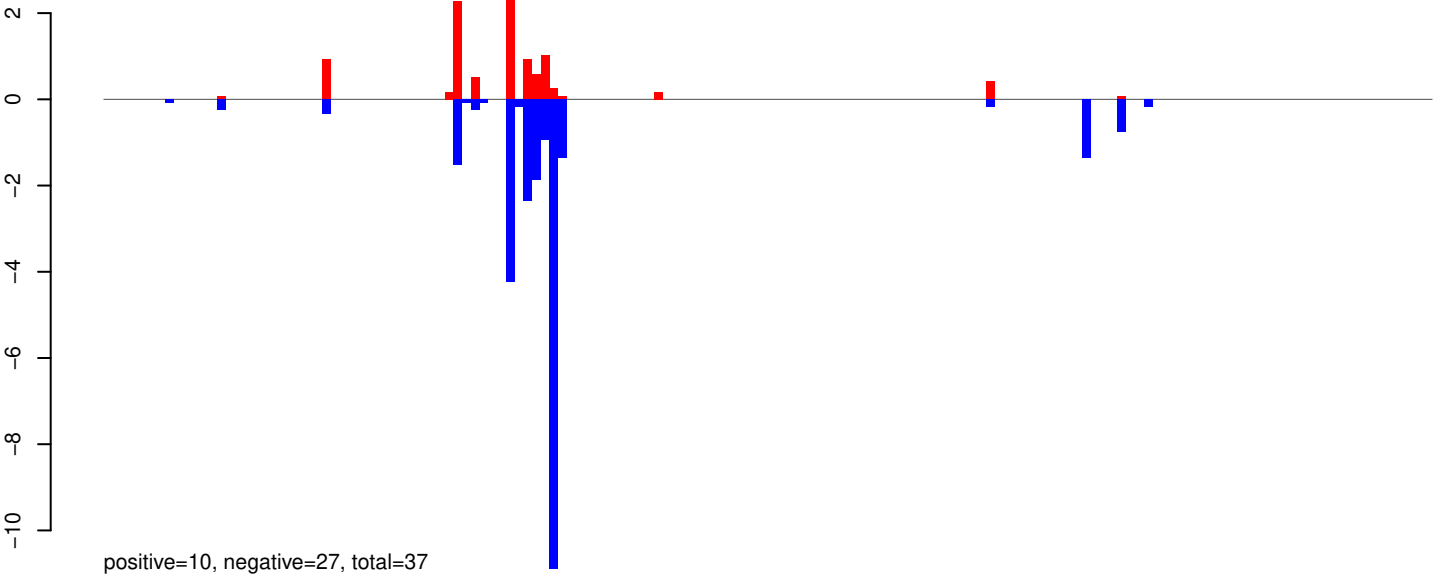
AeAeg_Aag2_BTV1_AK.rep



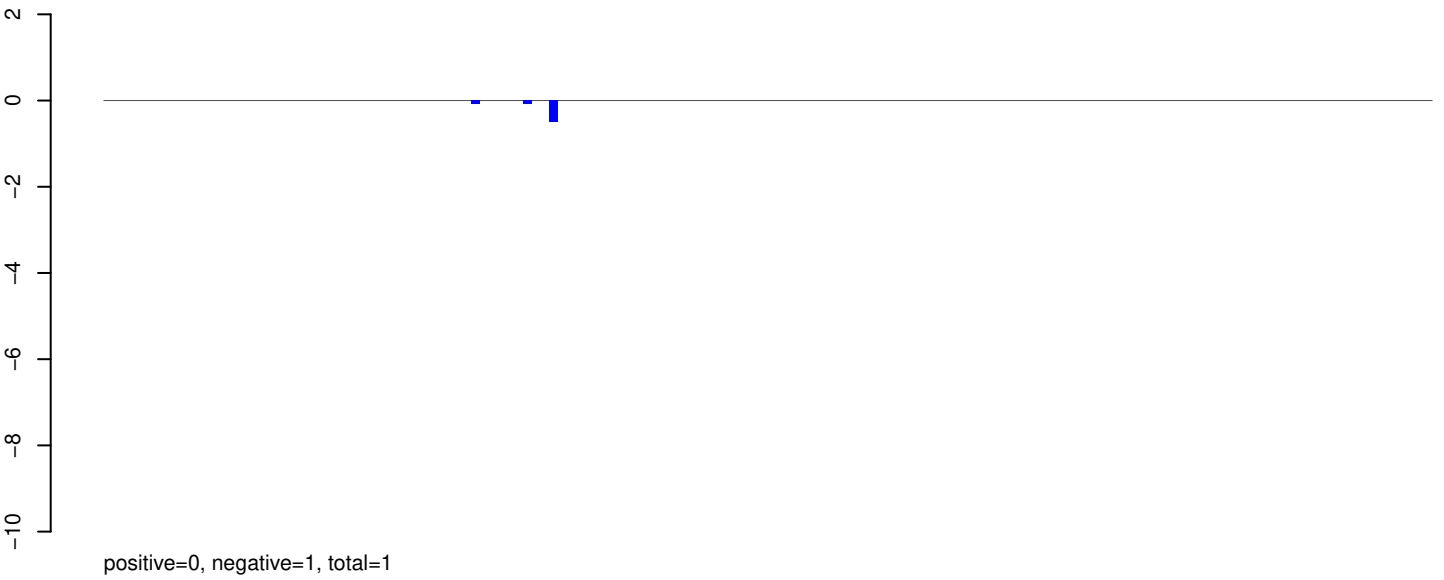
positive=2, negative=2, total=4
Window size=25, step=1, length=12294, virus@NC_001927.1-Bunyamwera_virus_segment_S_plus_segment_M_plus_segment_L:1-12294

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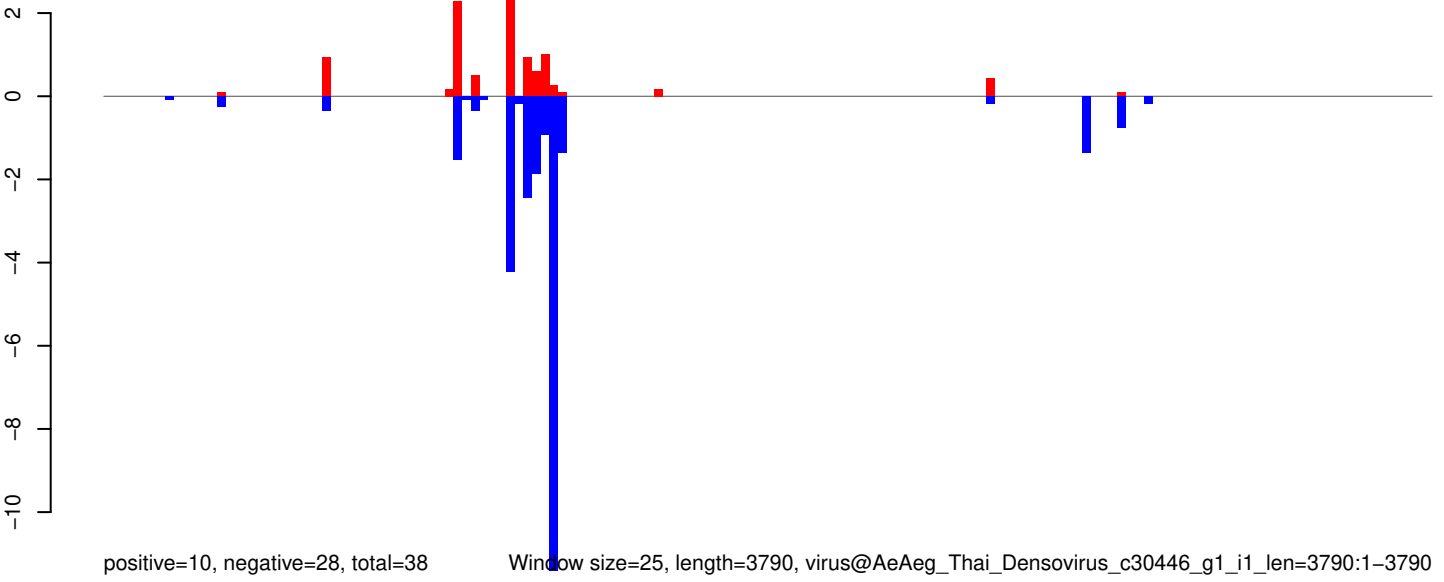
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AeAeg_Aag2_BTV1_AK.24_35.rep

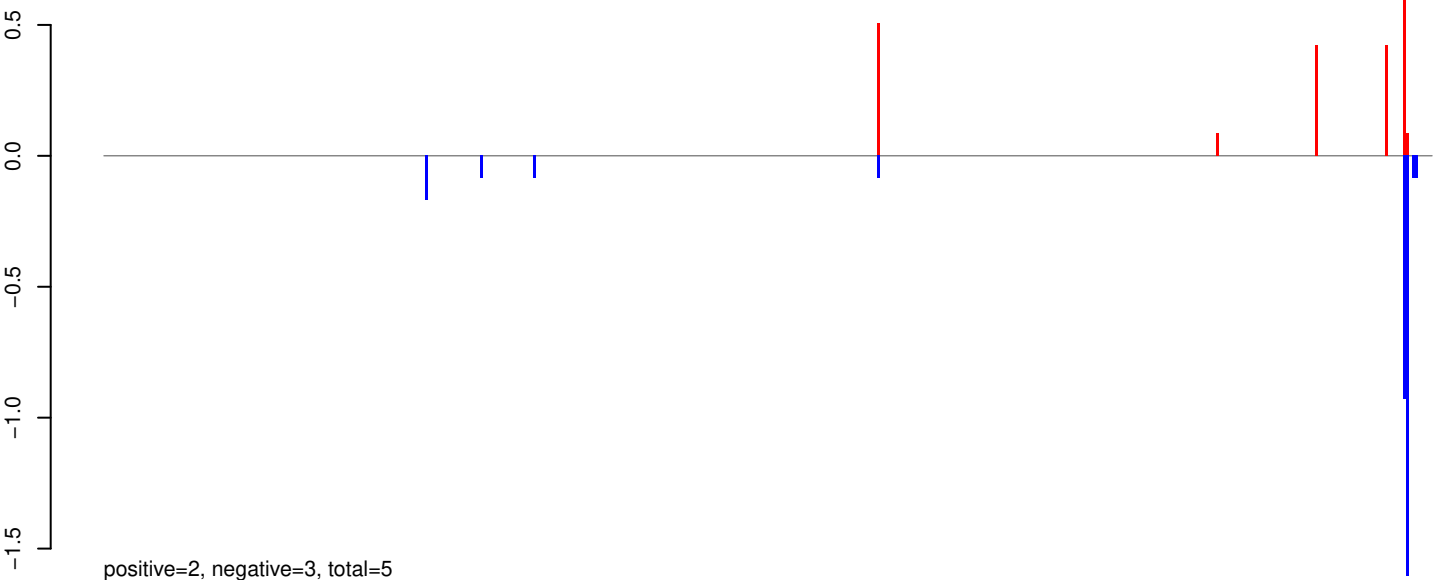


AeAeg_Aag2_BTV1_AK.rep

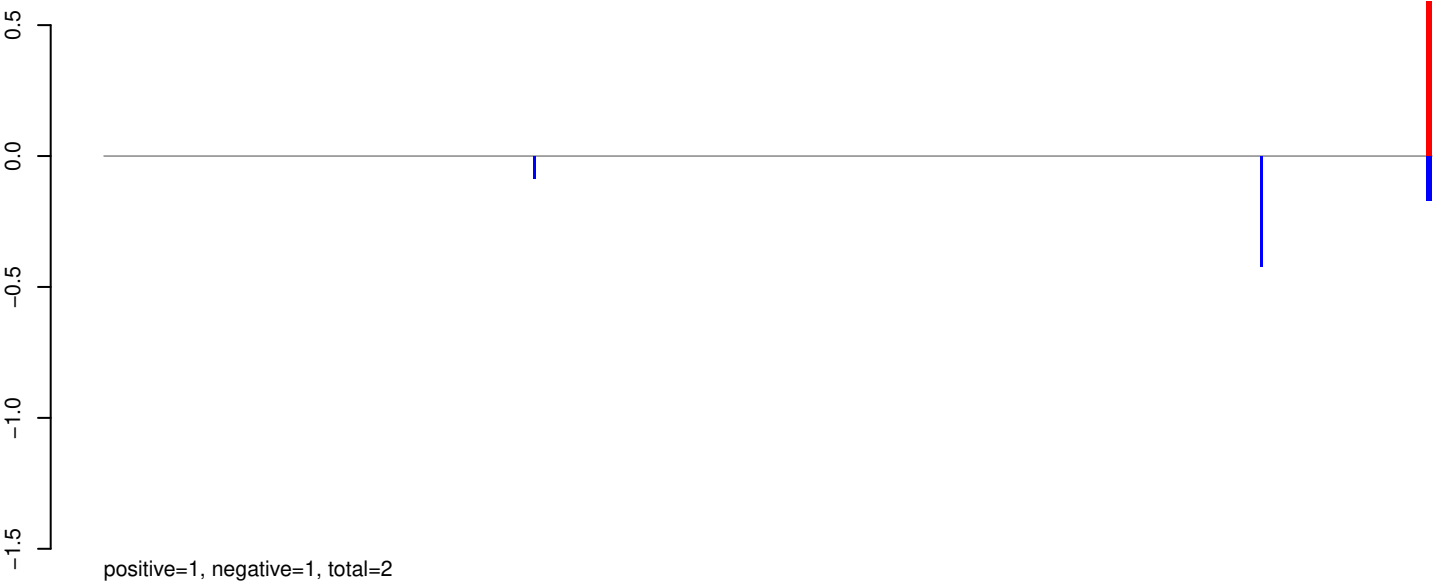


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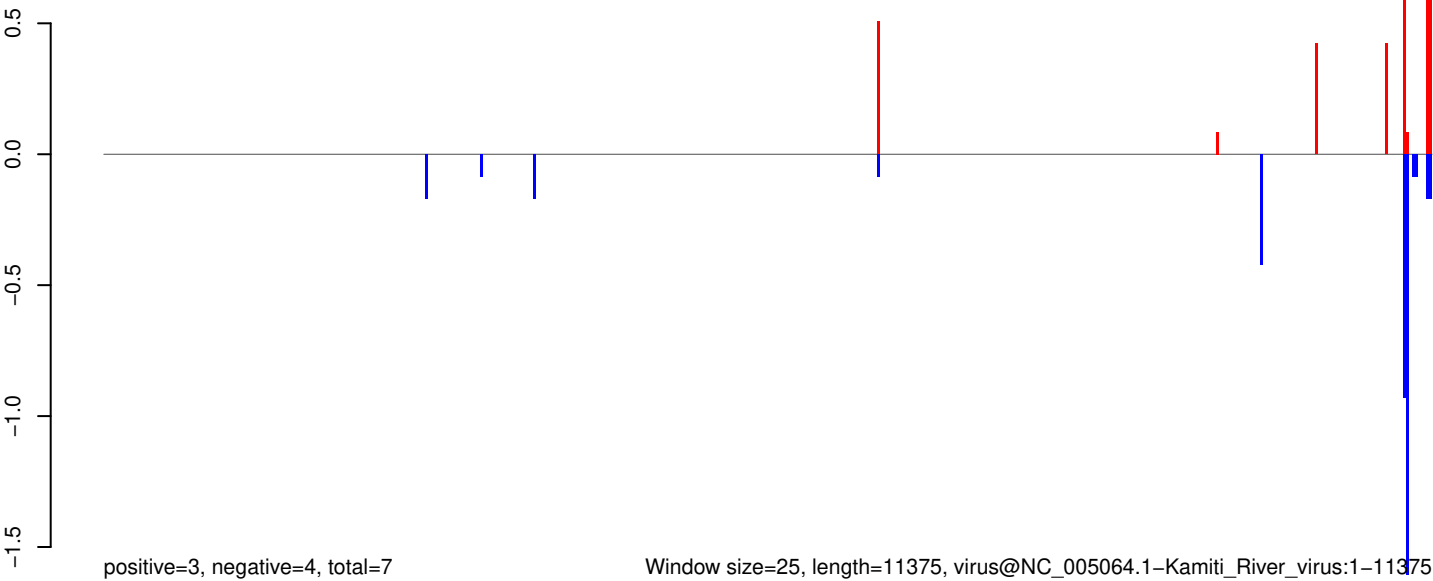
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AeAeg_Aag2_BTV1_AK.24_35.rep

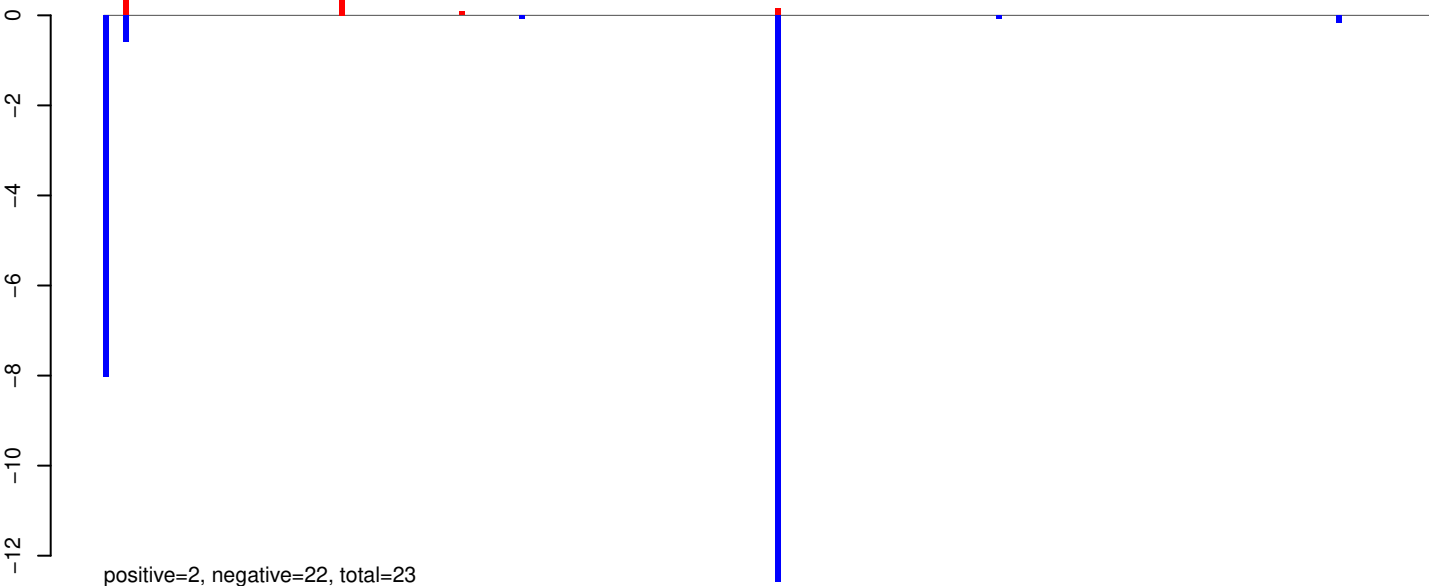


AeAeg_Aag2_BTV1_AK.rep

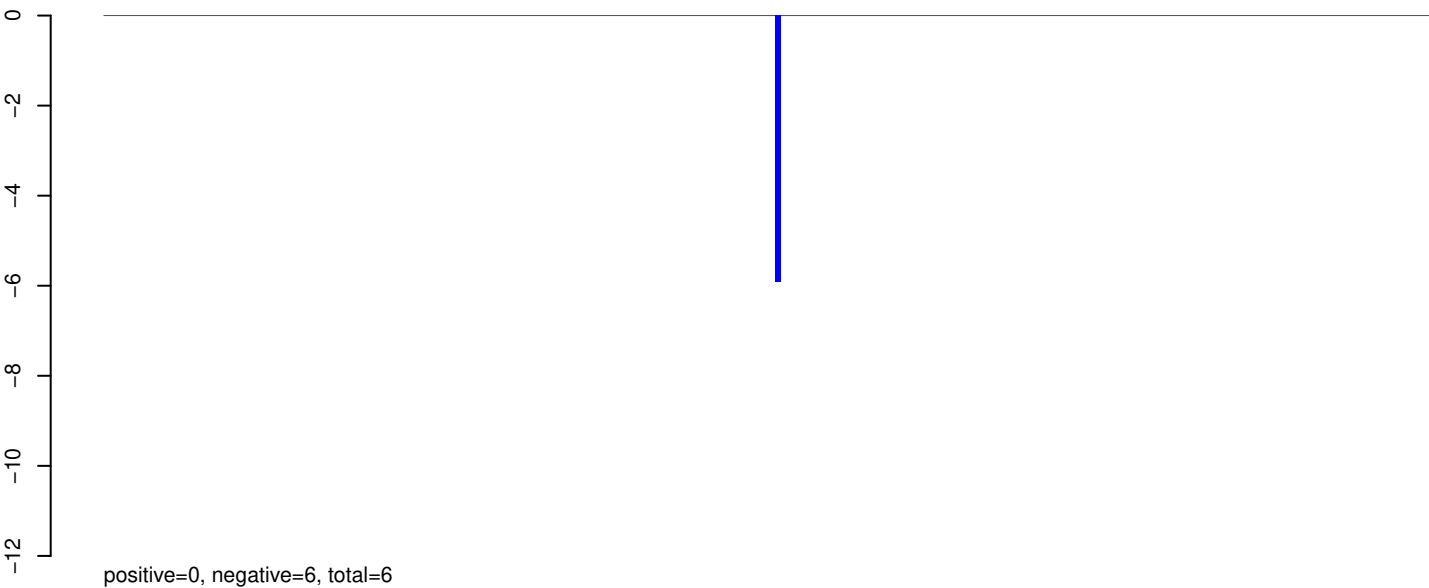


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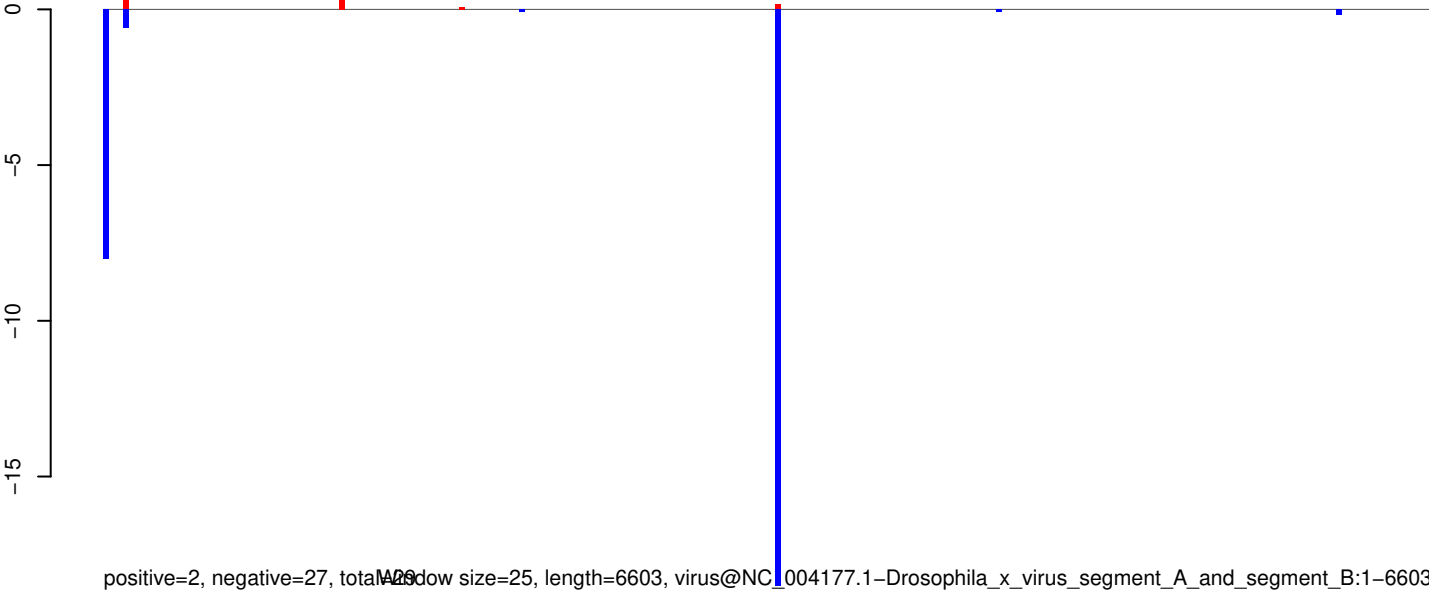
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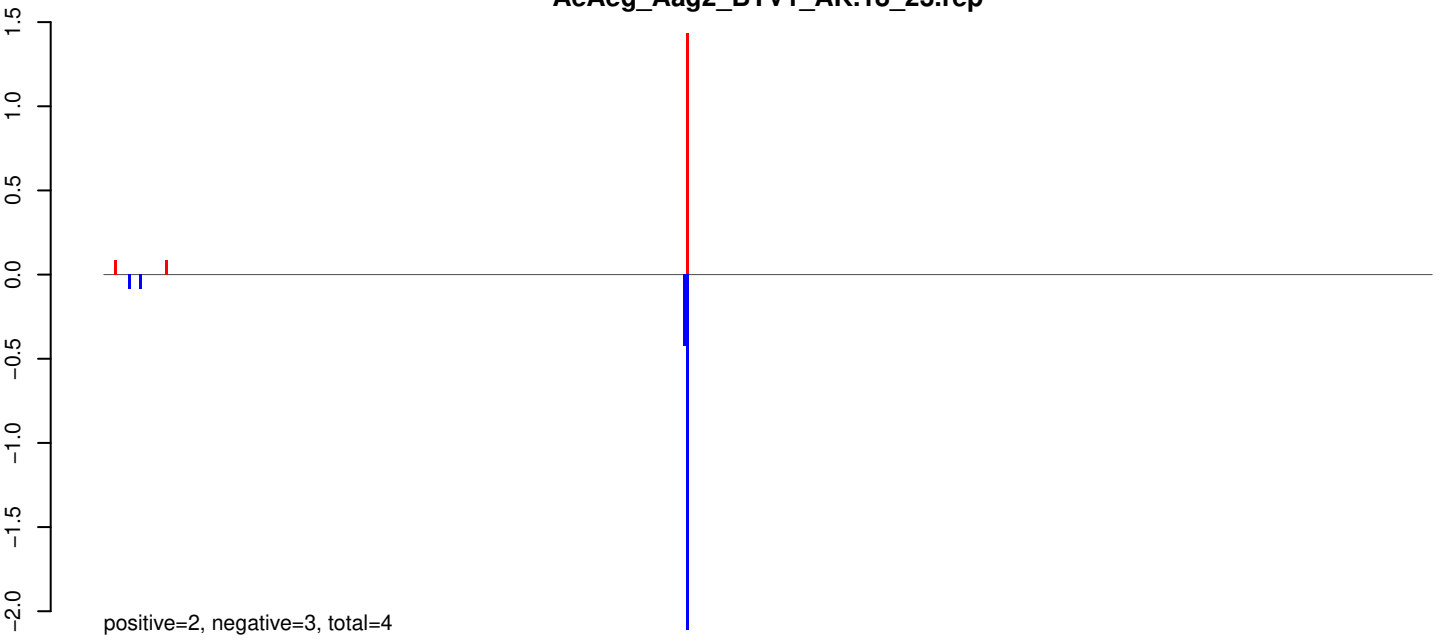
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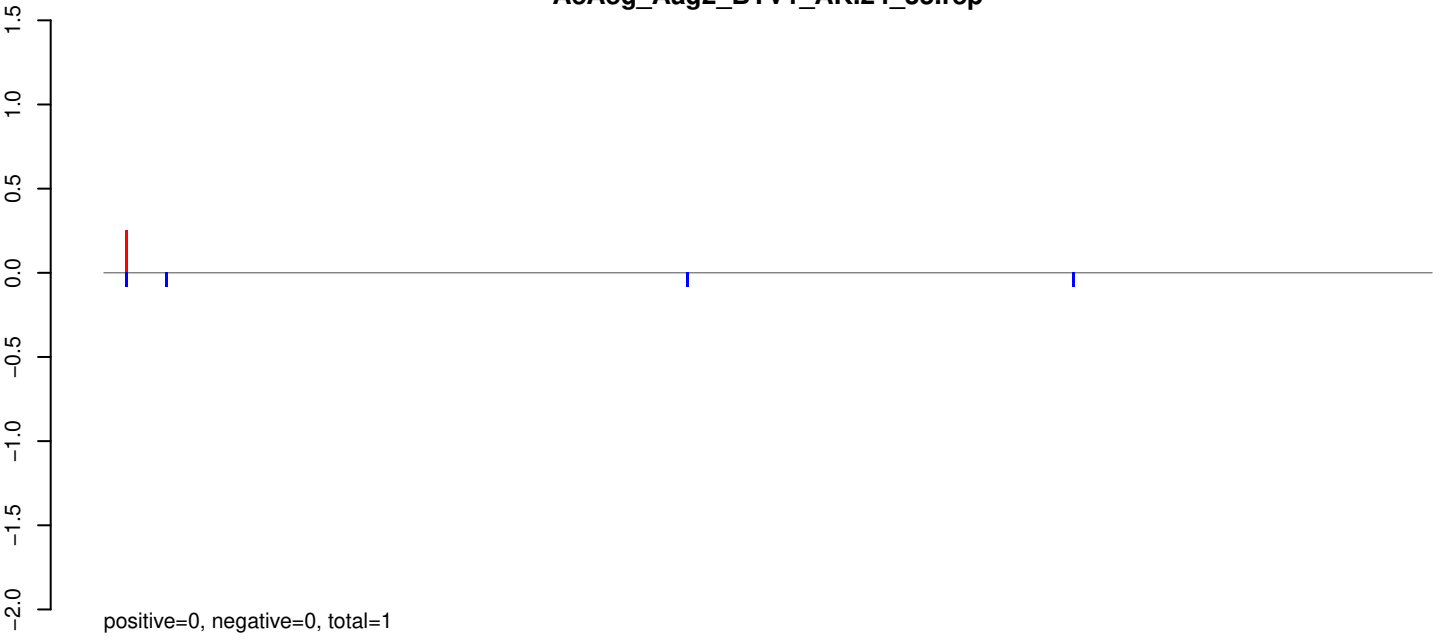
AeAeg_Aag2_BTV1_AK.rep



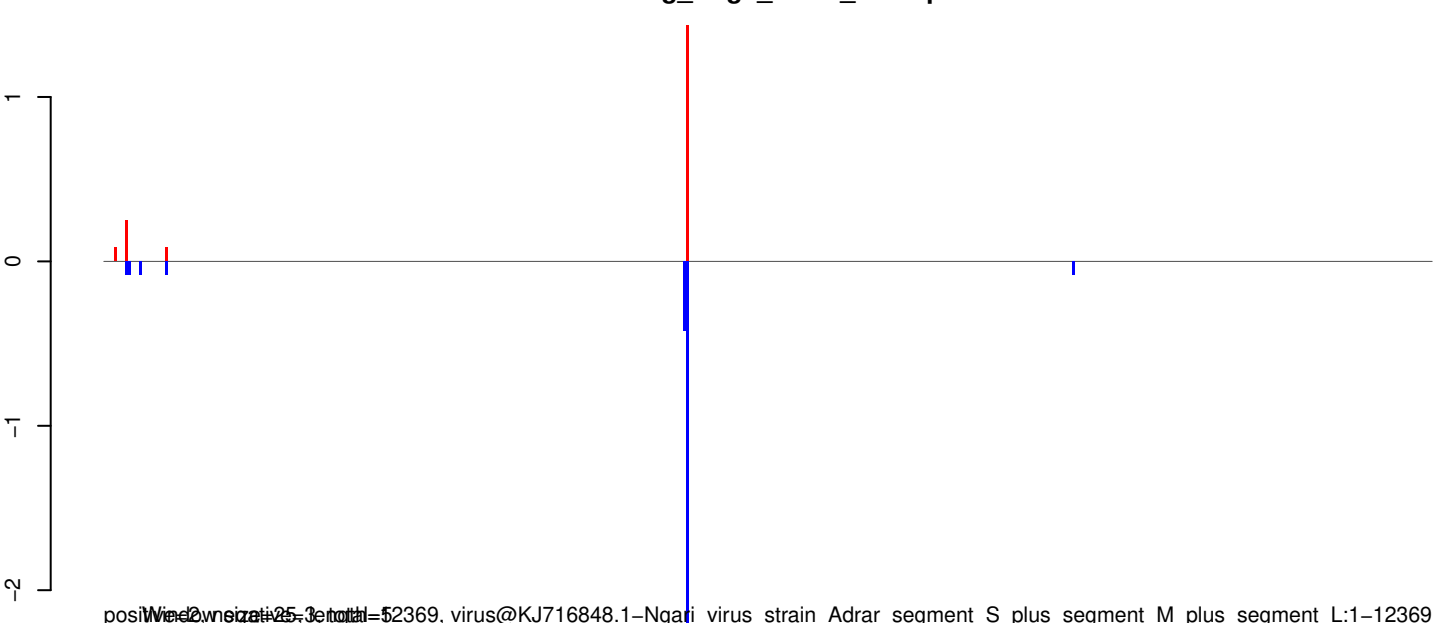
AeAeg_Aag2_BTV1_AK.18_23.rep



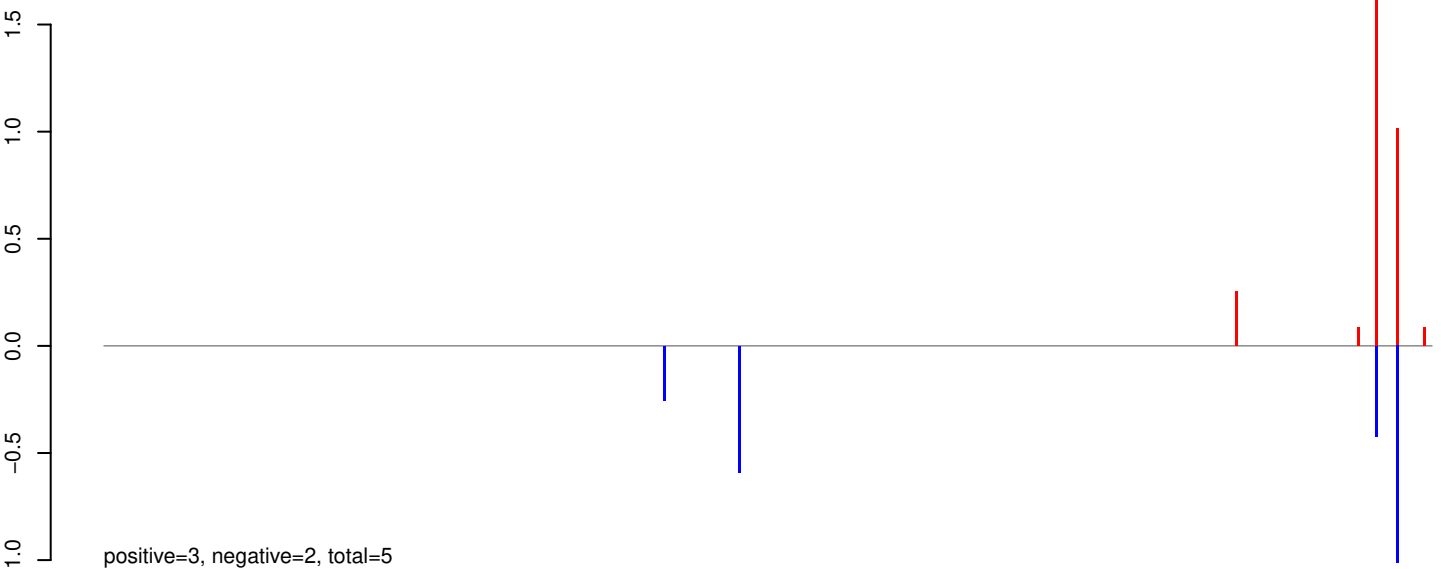
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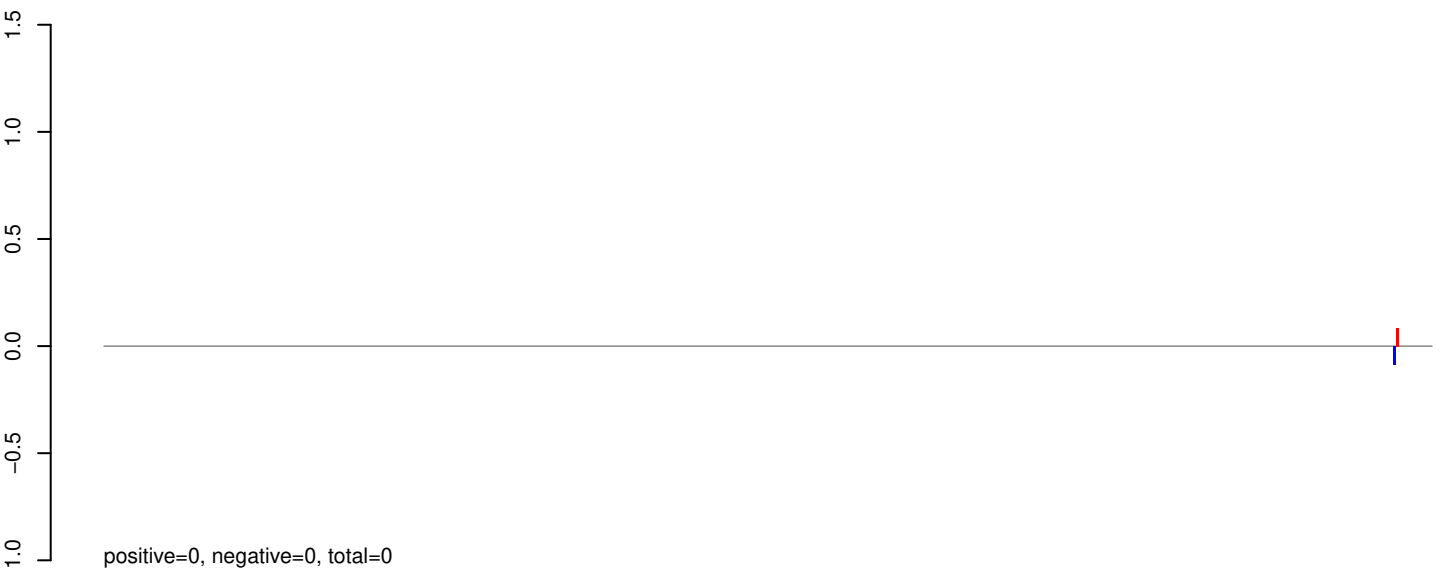
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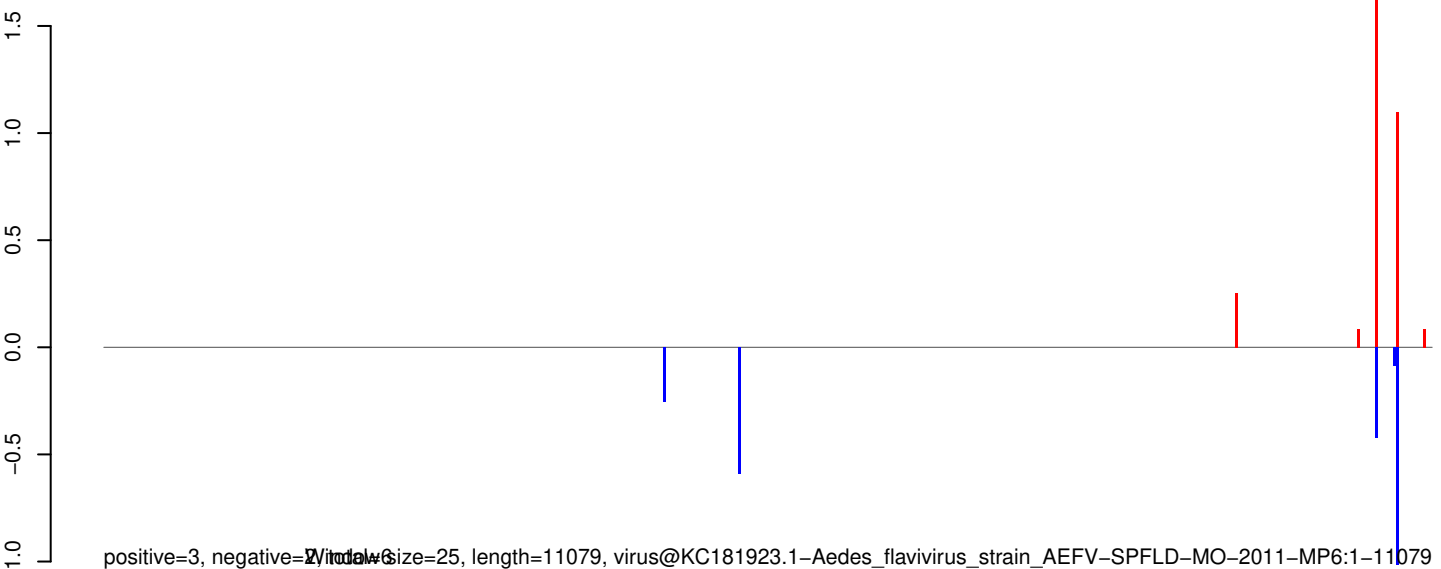
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AeAeg_Aag2_BTV1_AK.24_35.rep



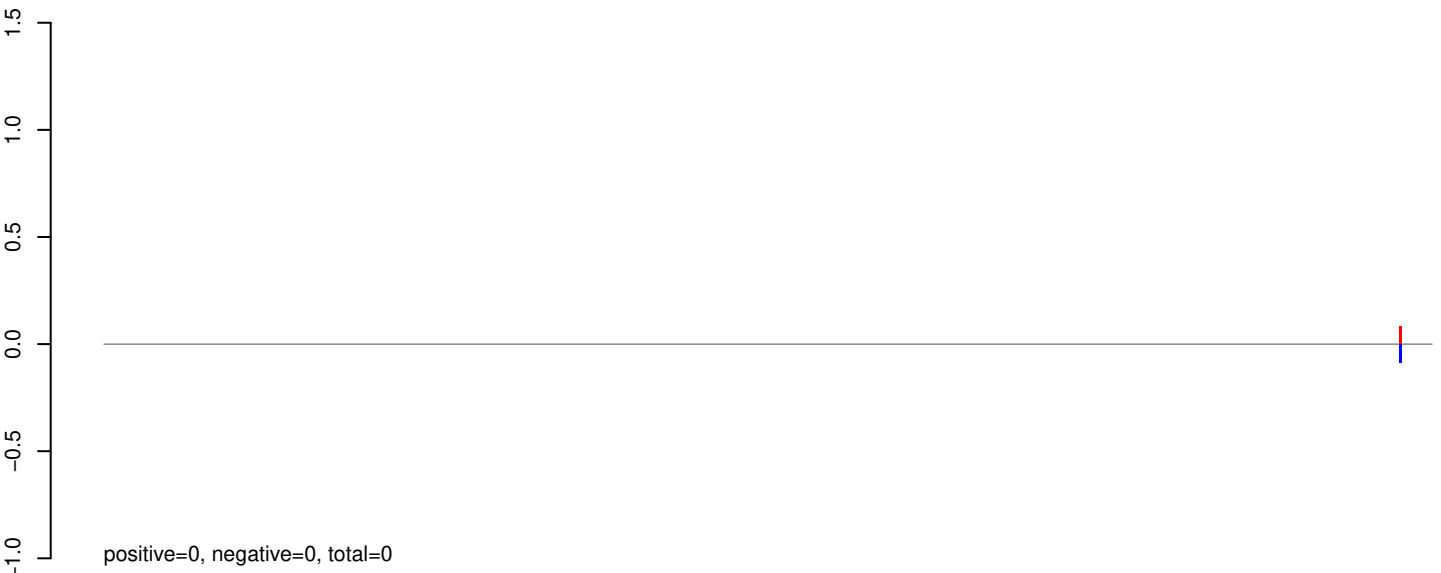
AeAeg_Aag2_BTV1_AK.rep



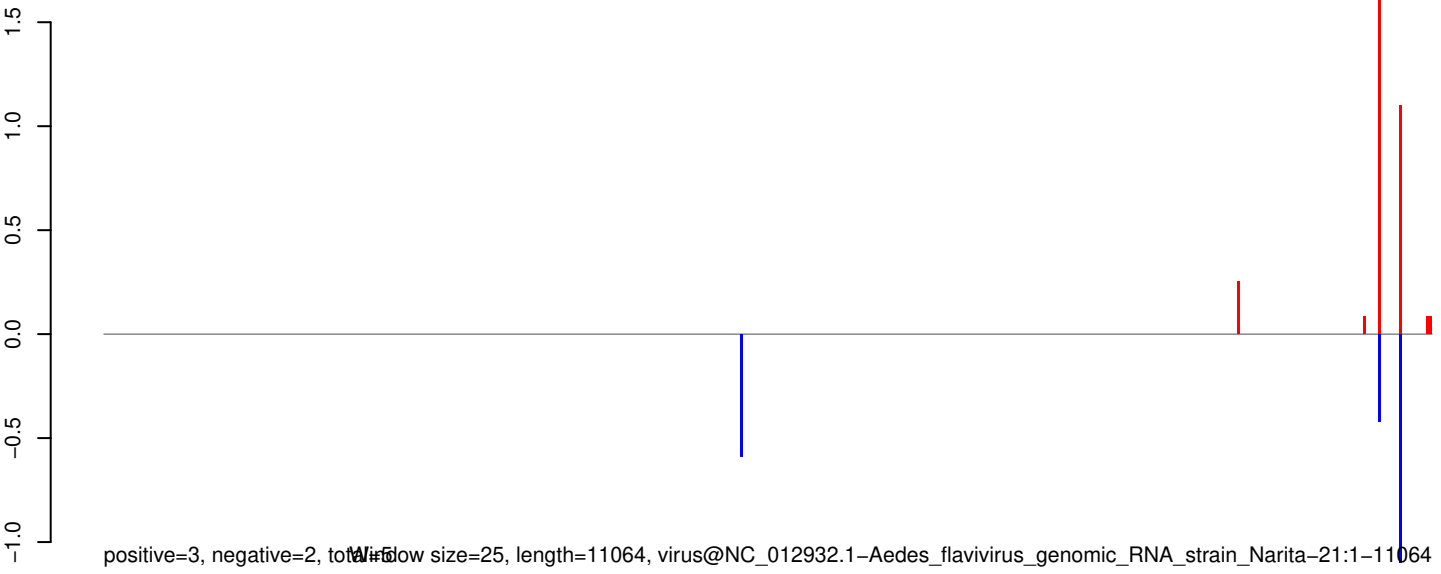
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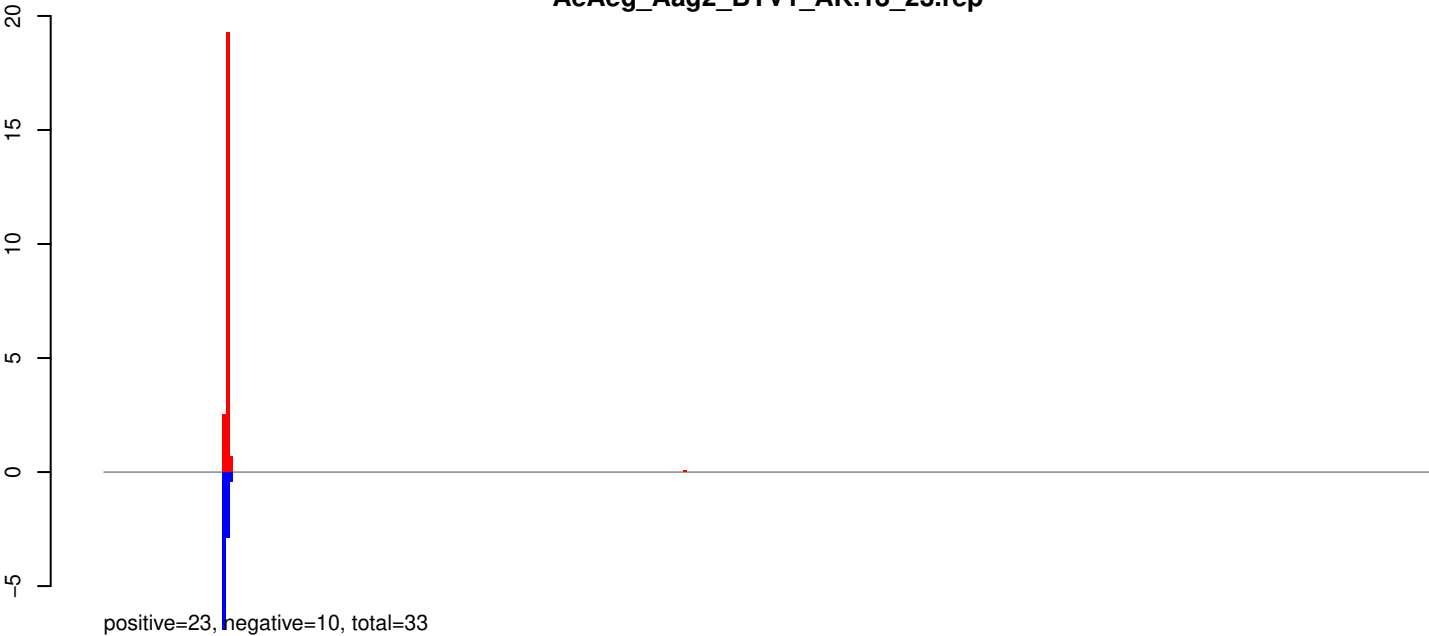
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AeAeg_Aag2_BTV1_AK.rep



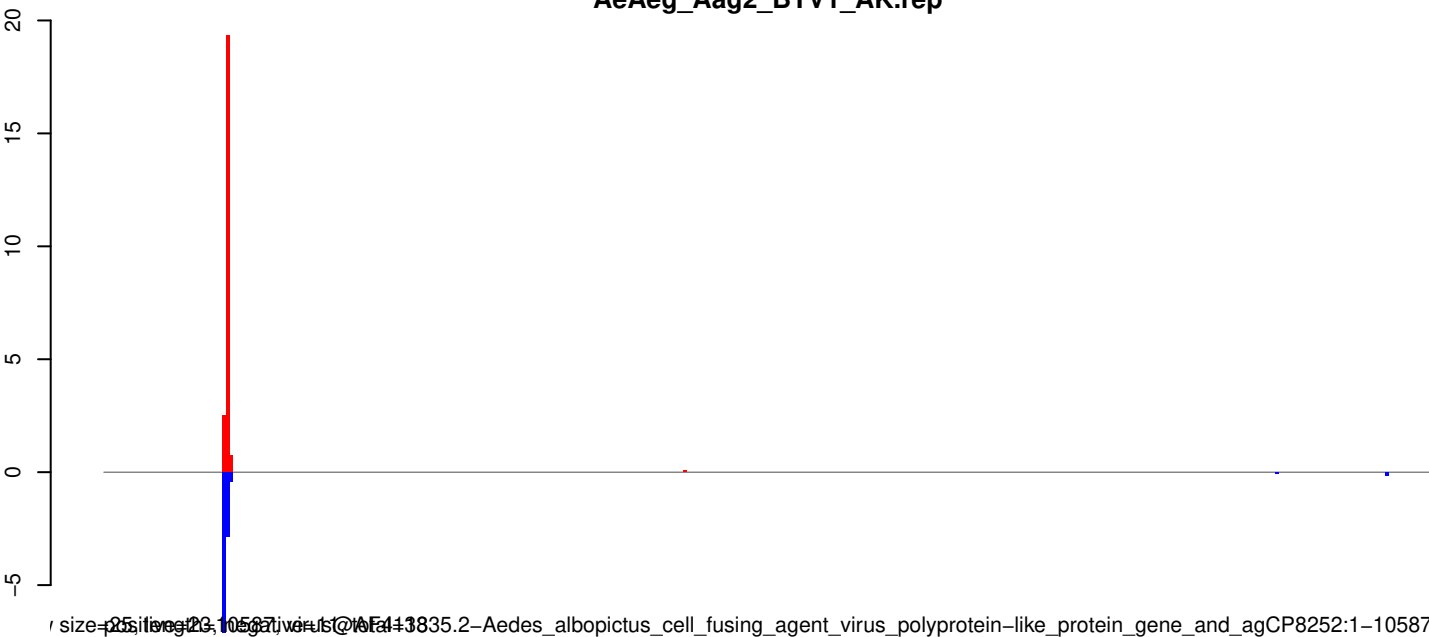
AeAeg_Aag2_BTV1_AK.18_23.rep



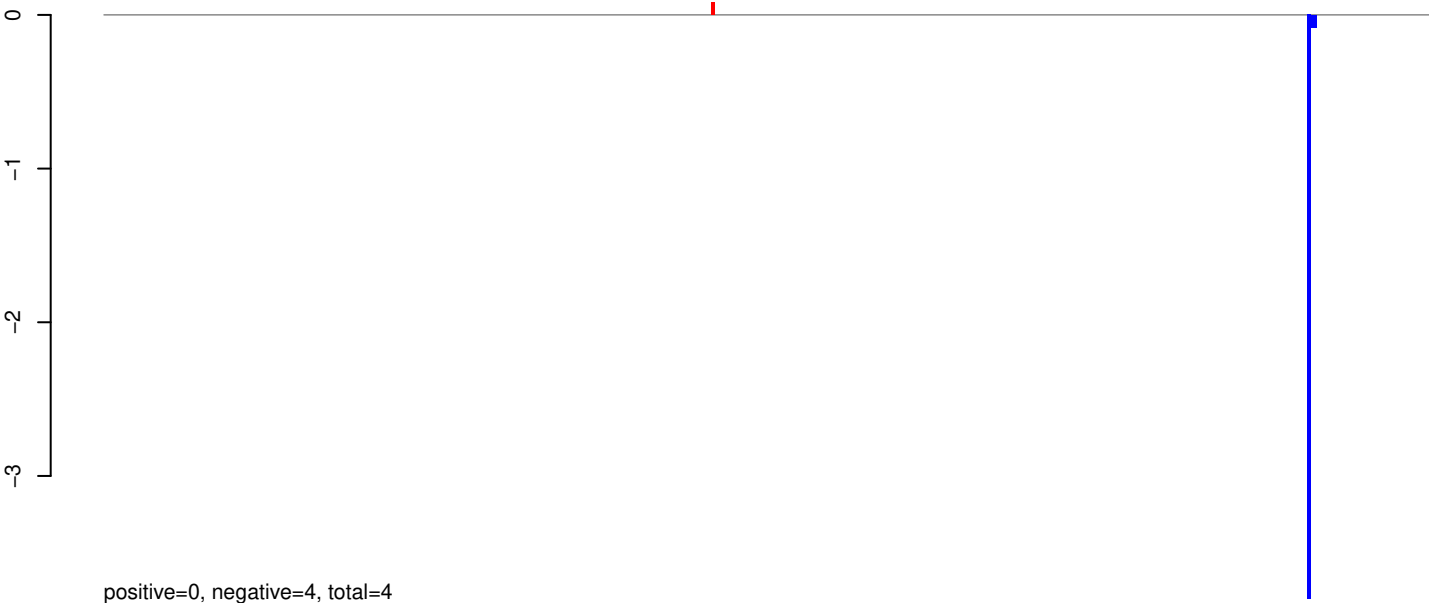
AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep

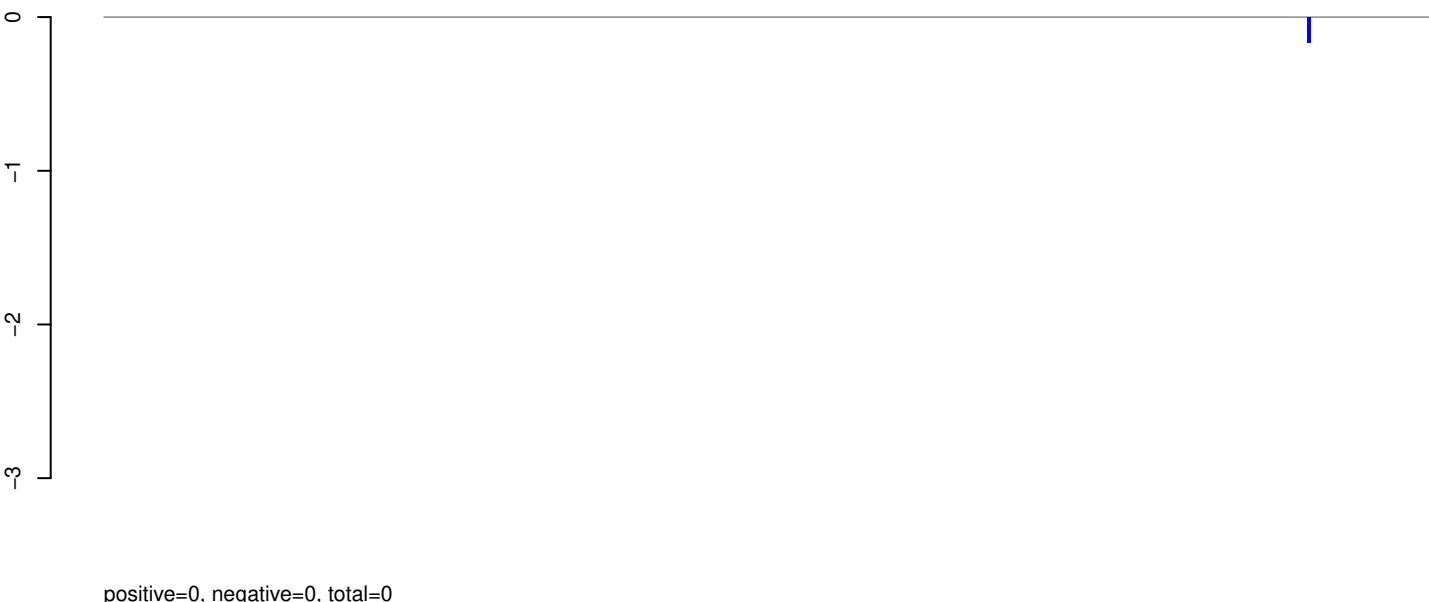


AeAeg_Aag2_BTV1_AK.18_23.rep



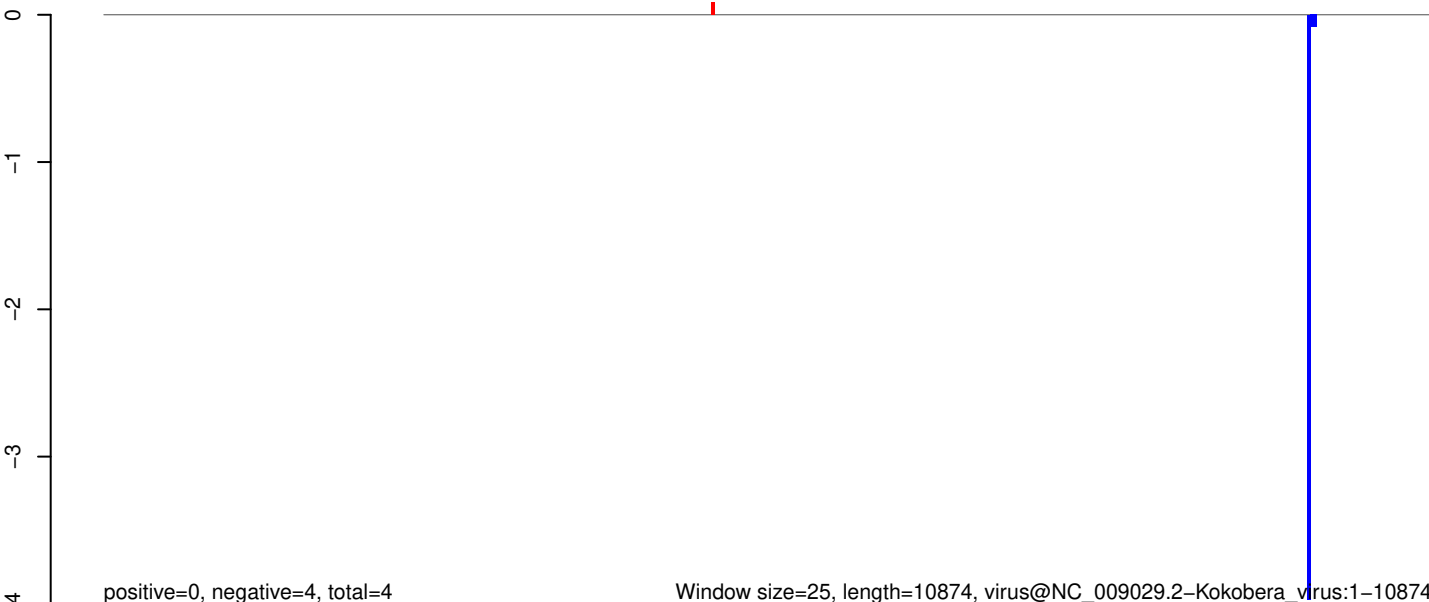
positive=0, negative=4, total=4

AeAeg_Aag2_BTV1_AK.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BTV1_AK.rep



positive=0, negative=4, total=4

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

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV1_AK.18_23.rep



positive=0, negative=1, total=1

AeAeg_Aag2_BTV1_AK.24_35.rep



positive=0, negative=11, total=11

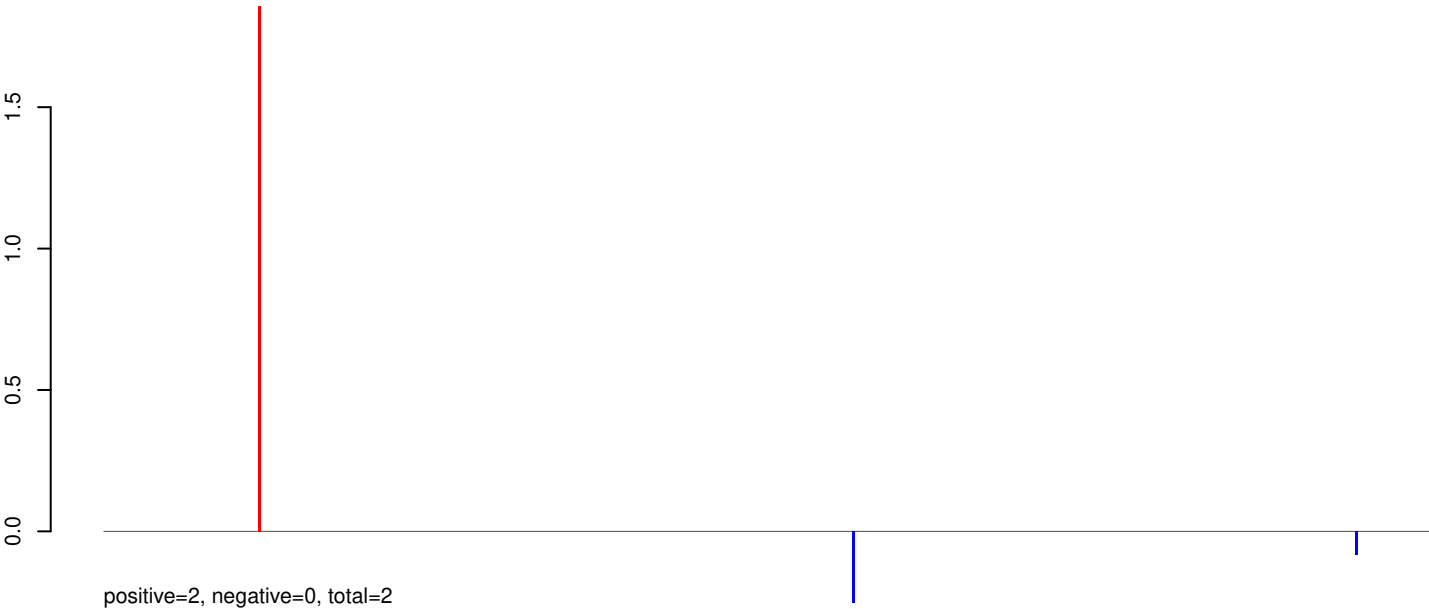
AeAeg_Aag2_BTV1_AK.rep



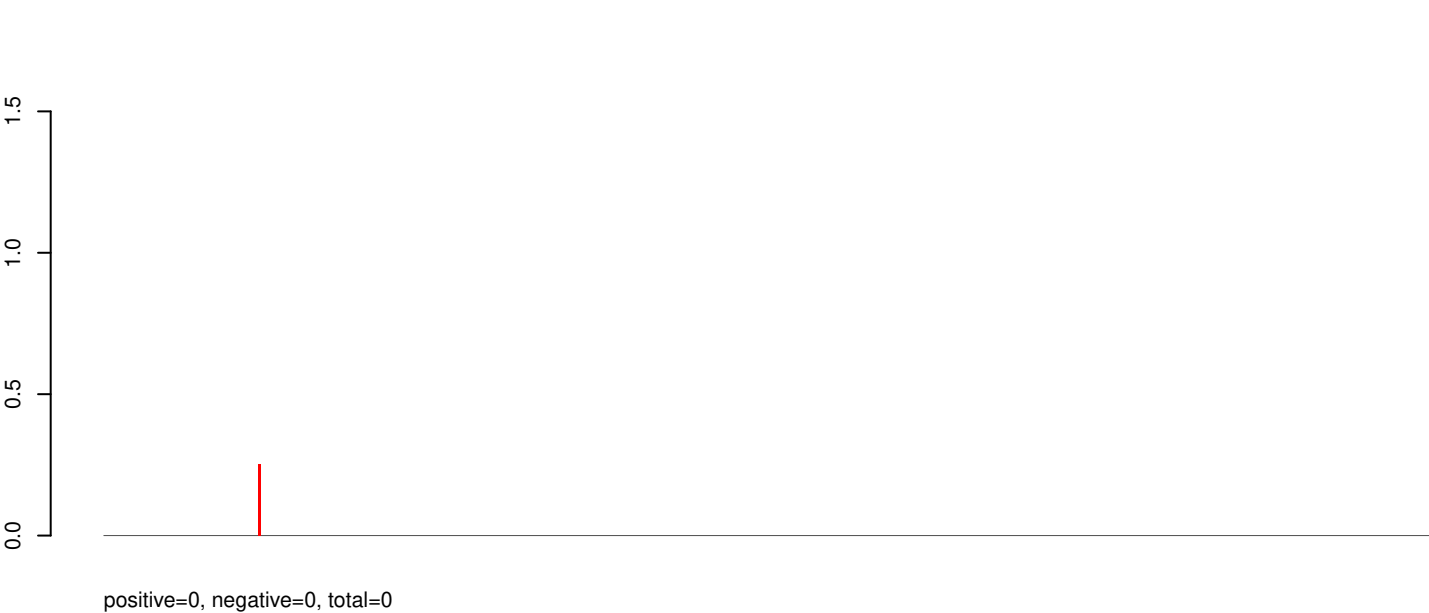
positive=0, negative=11, window size=25, length=6203, virus@NC_035674.1-Australian_Anopheles_totivirus_isolate_AATV_150840:1-6203

0 1000 2000 3000 4000 5000 6000

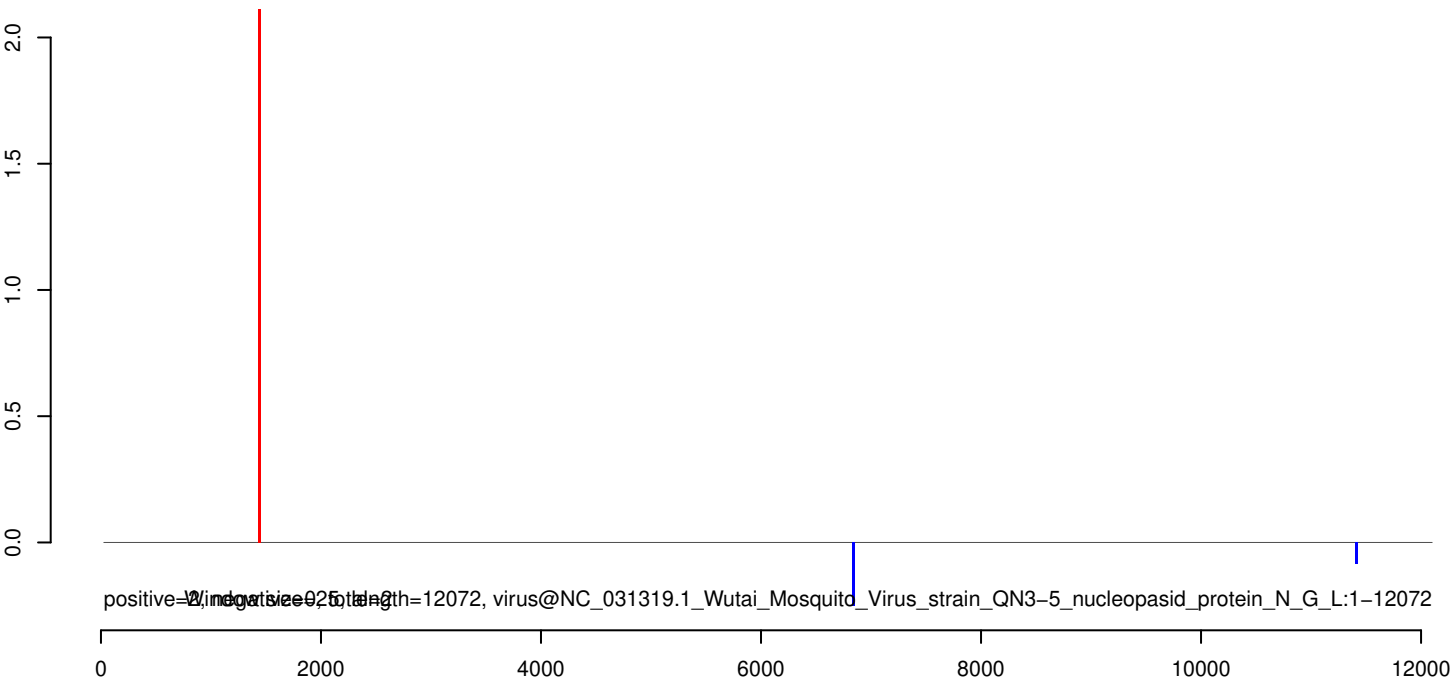
AeAeg_Aag2_BTV1_AK.18_23.rep



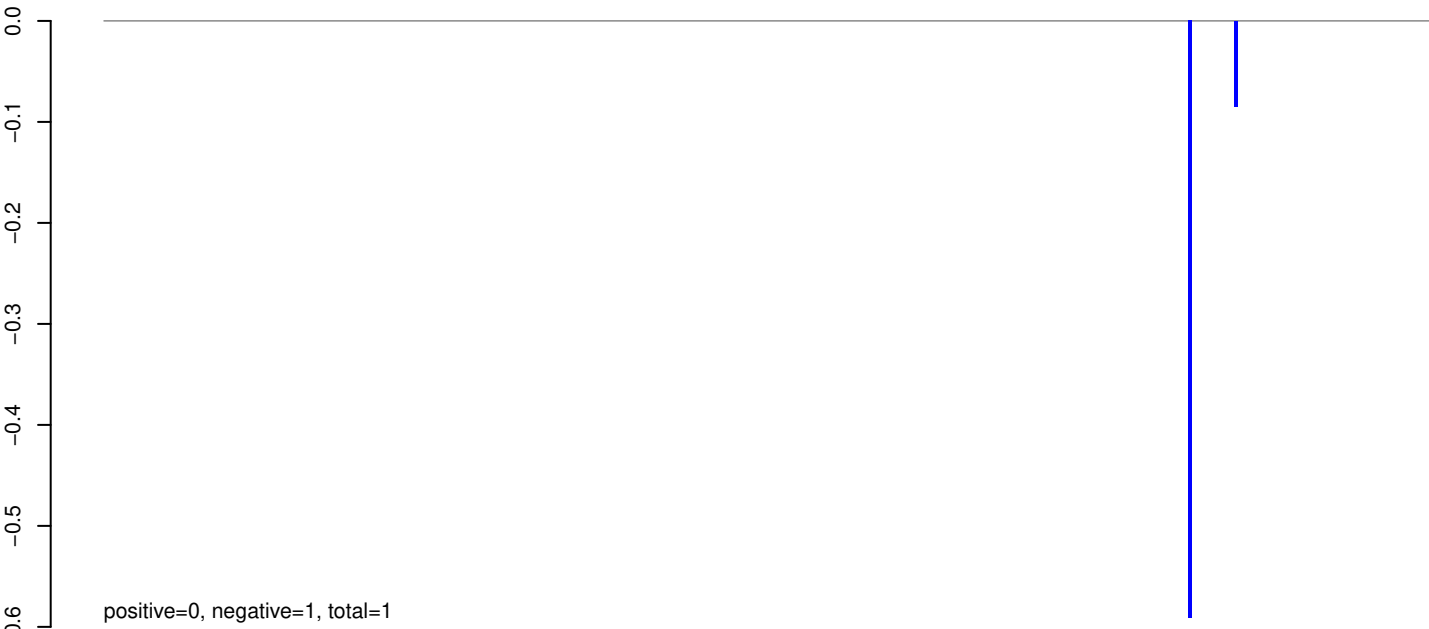
AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



AeAeg_Aag2_BTV1_AK.18_23.rep



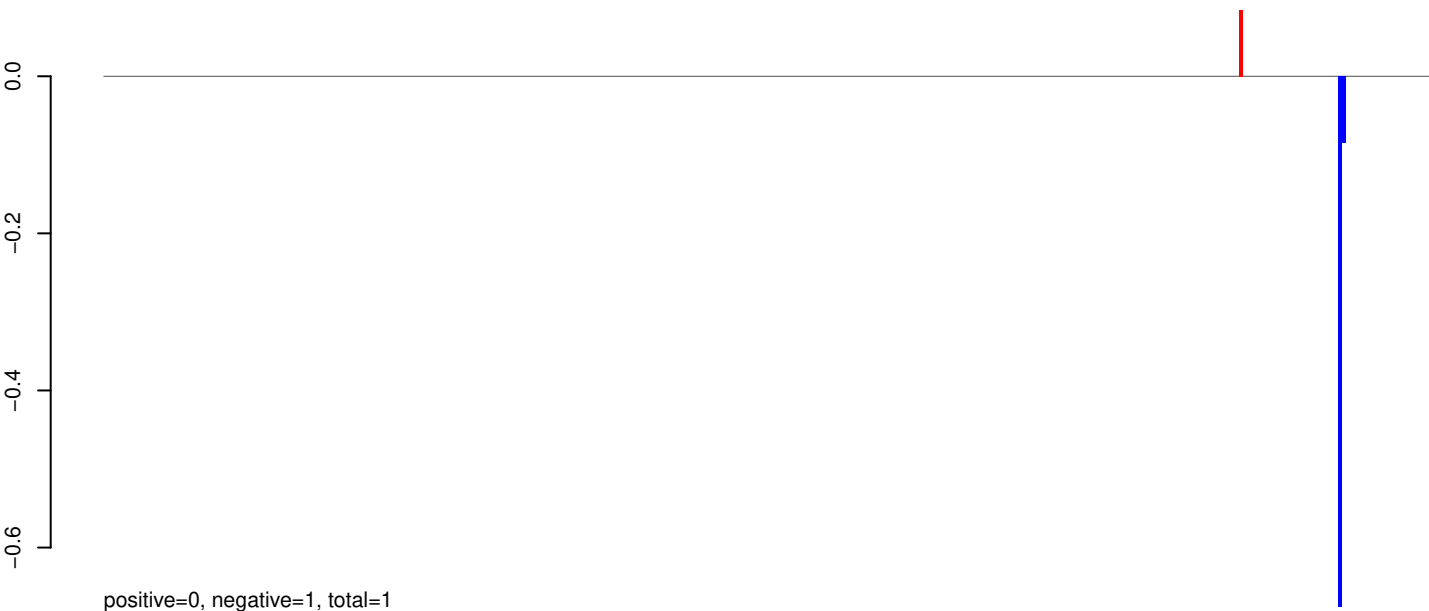
AeAeg_Aag2_BTV1_AK.24_35.rep



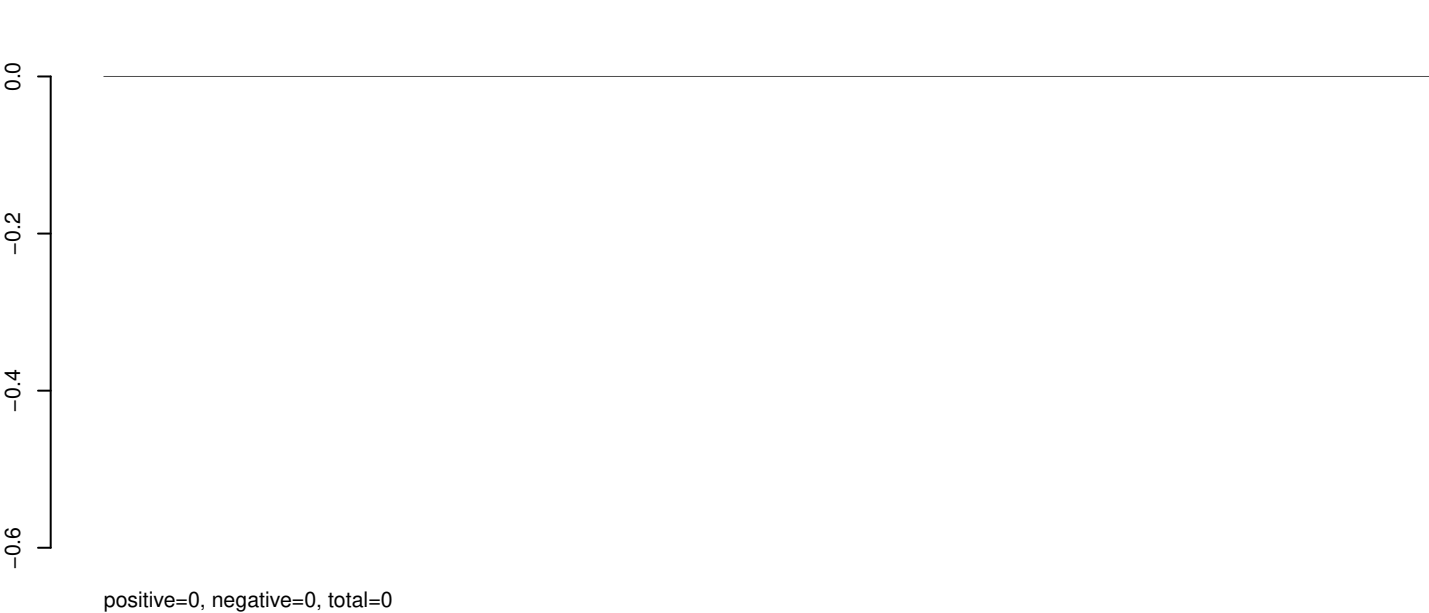
AeAeg_Aag2_BTV1_AK.rep



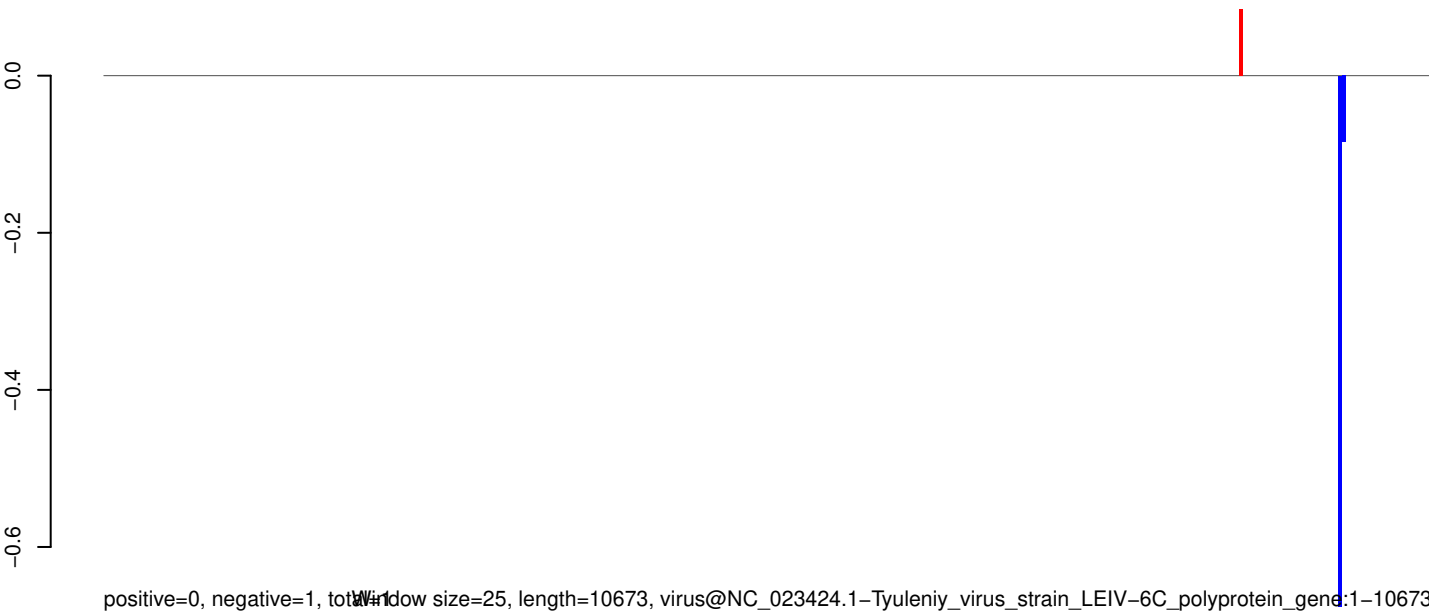
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



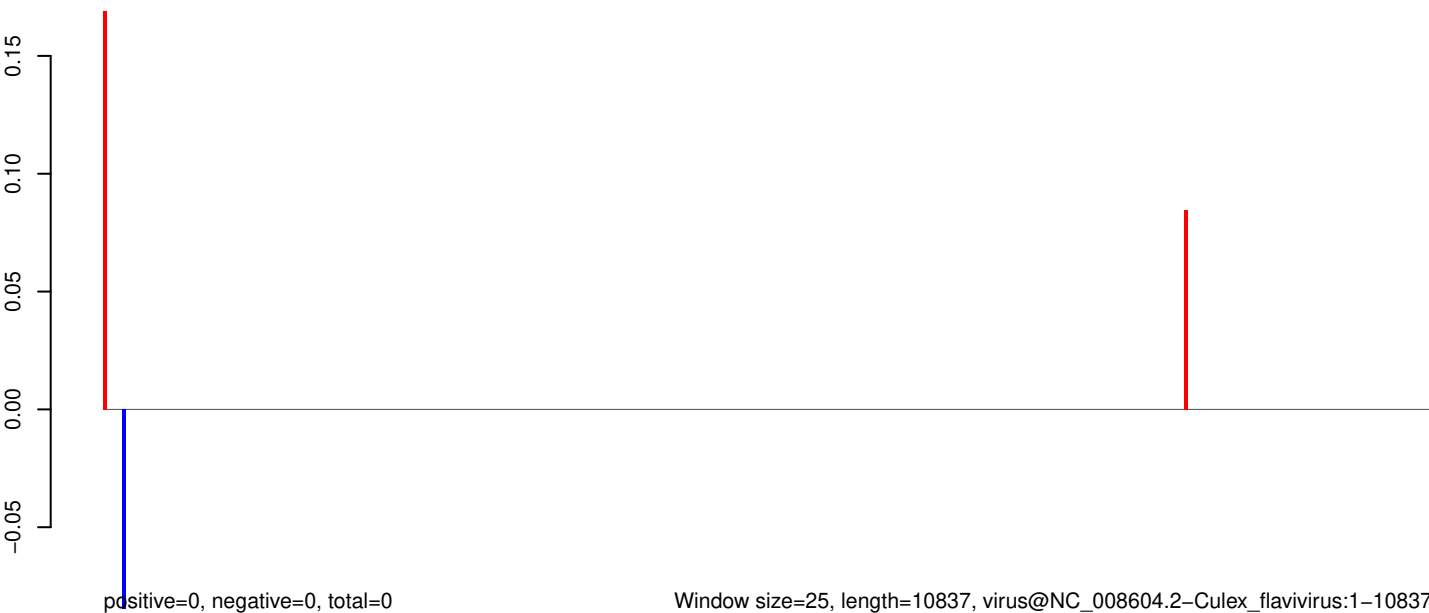
AeAeg_Aag2_BTV1_AK.18_23.rep



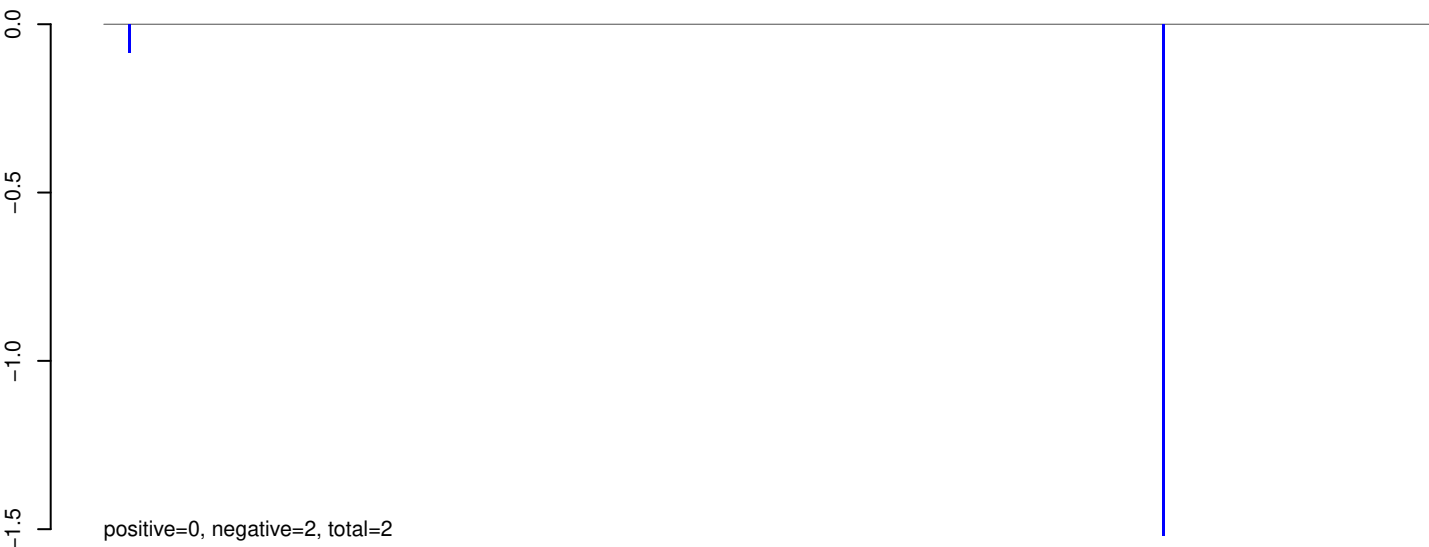
AeAeg_Aag2_BTV1_AK.24_35.rep



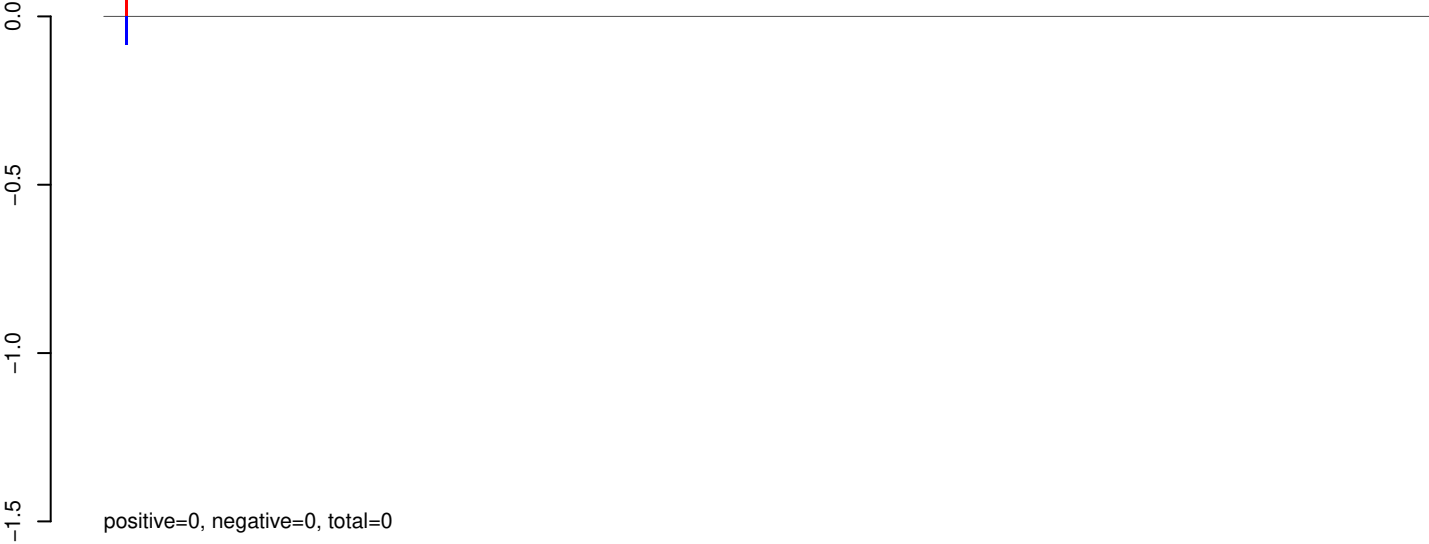
AeAeg_Aag2_BTV1_AK.rep



AeAeg_Aag2_BTV1_AK.18_23.rep



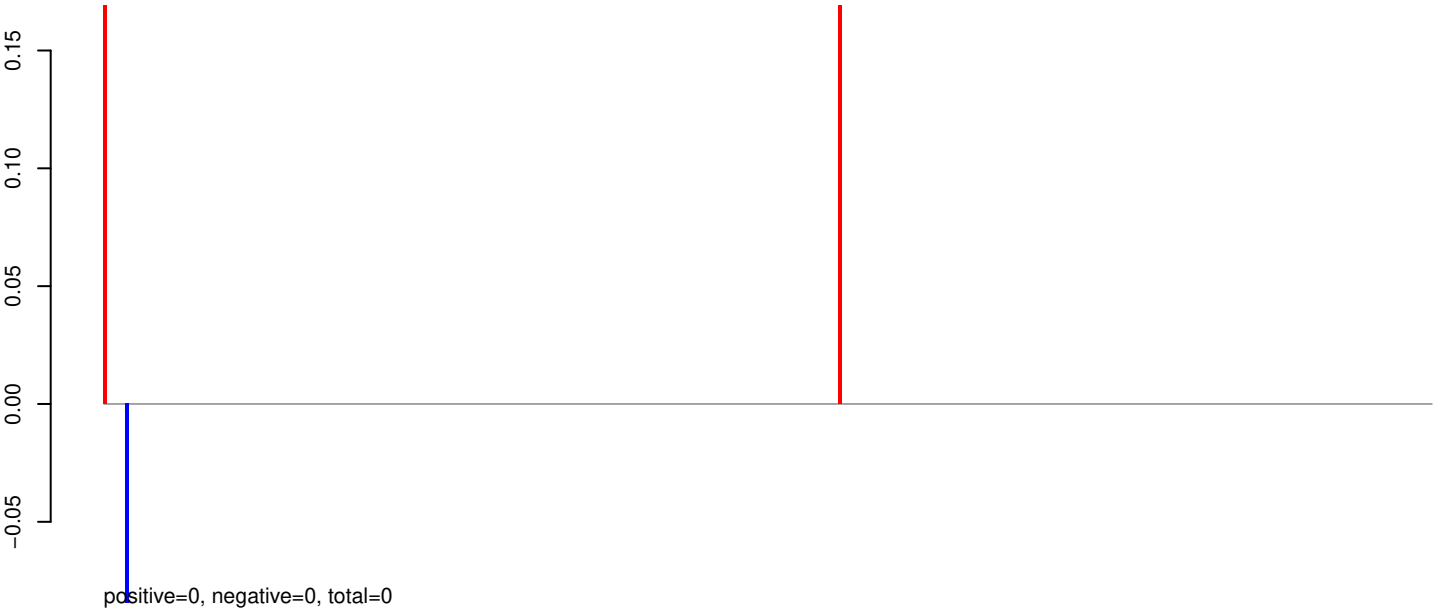
AeAeg_Aag2_BTV1_AK.24_35.rep



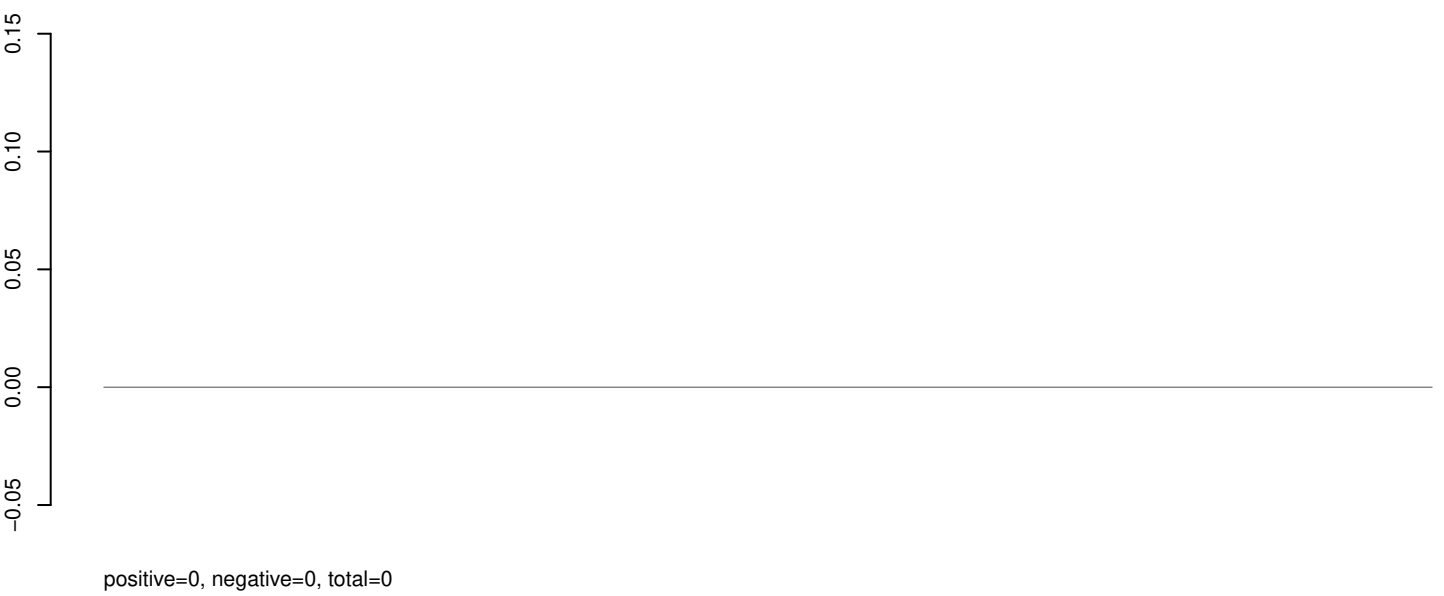
AeAeg_Aag2_BTV1_AK.rep



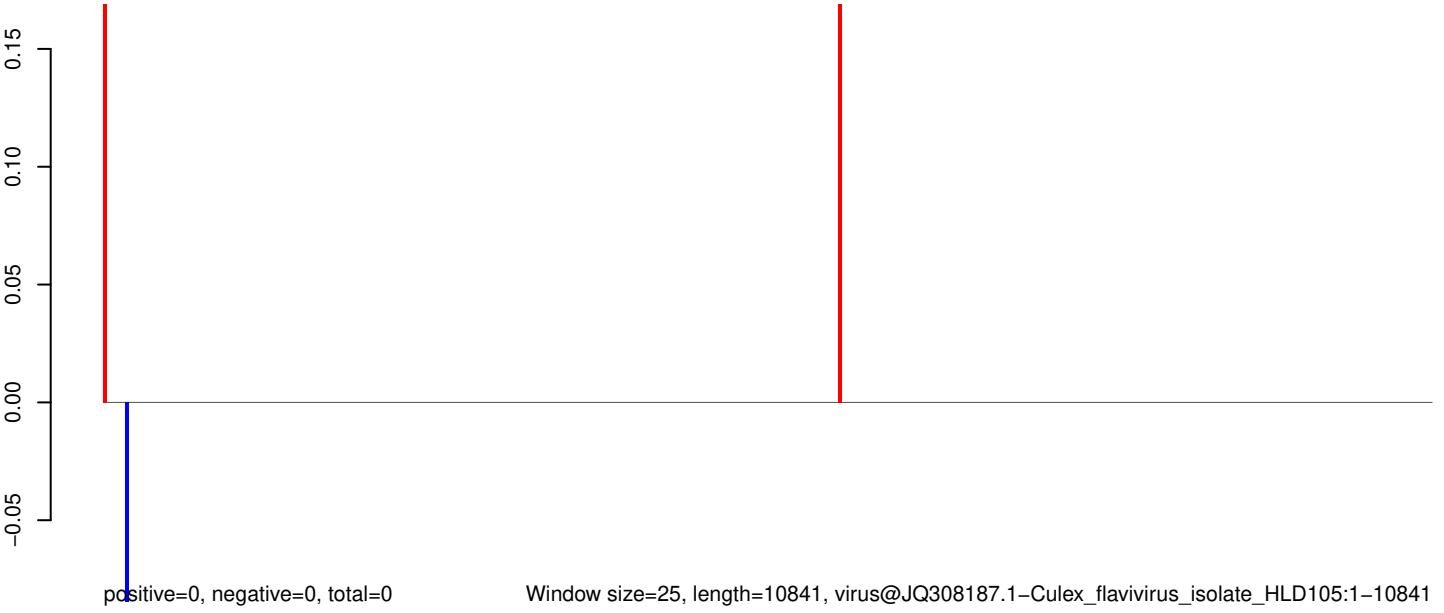
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



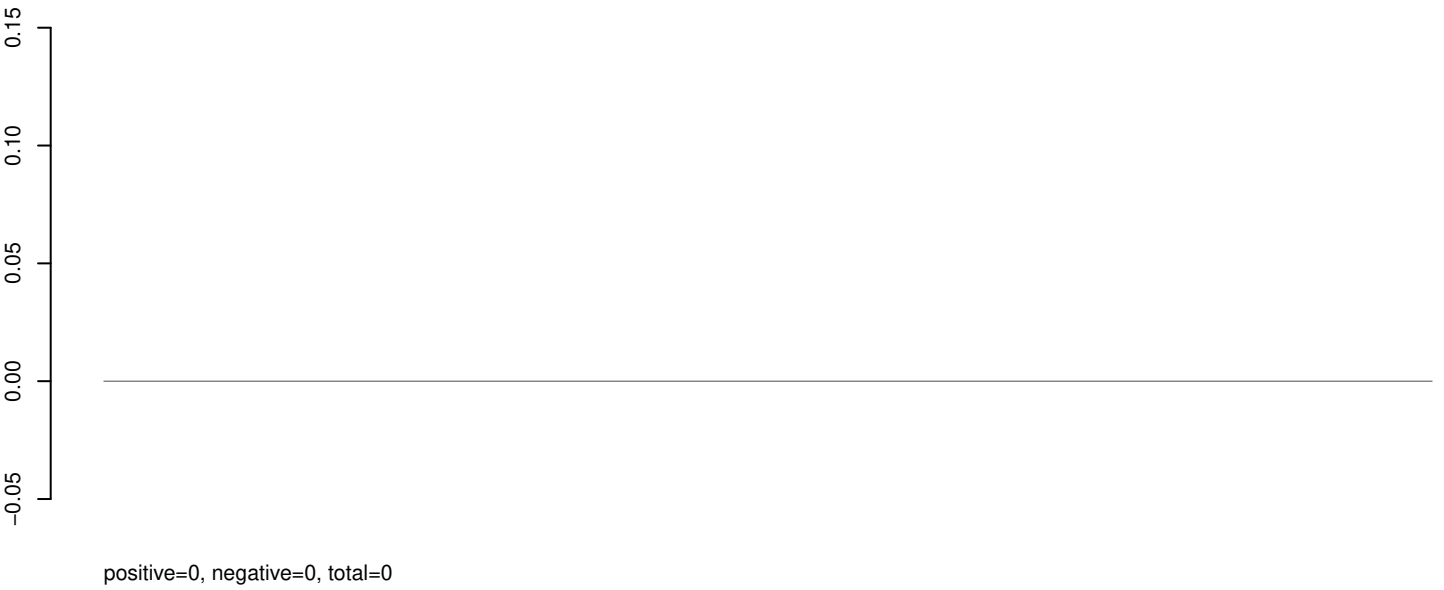
Window size=25, length=10841, virus@JQ308187.1-Culex_flavivirus_isolate_HLD105:1-10841

0 2000 4000 6000 8000 10000

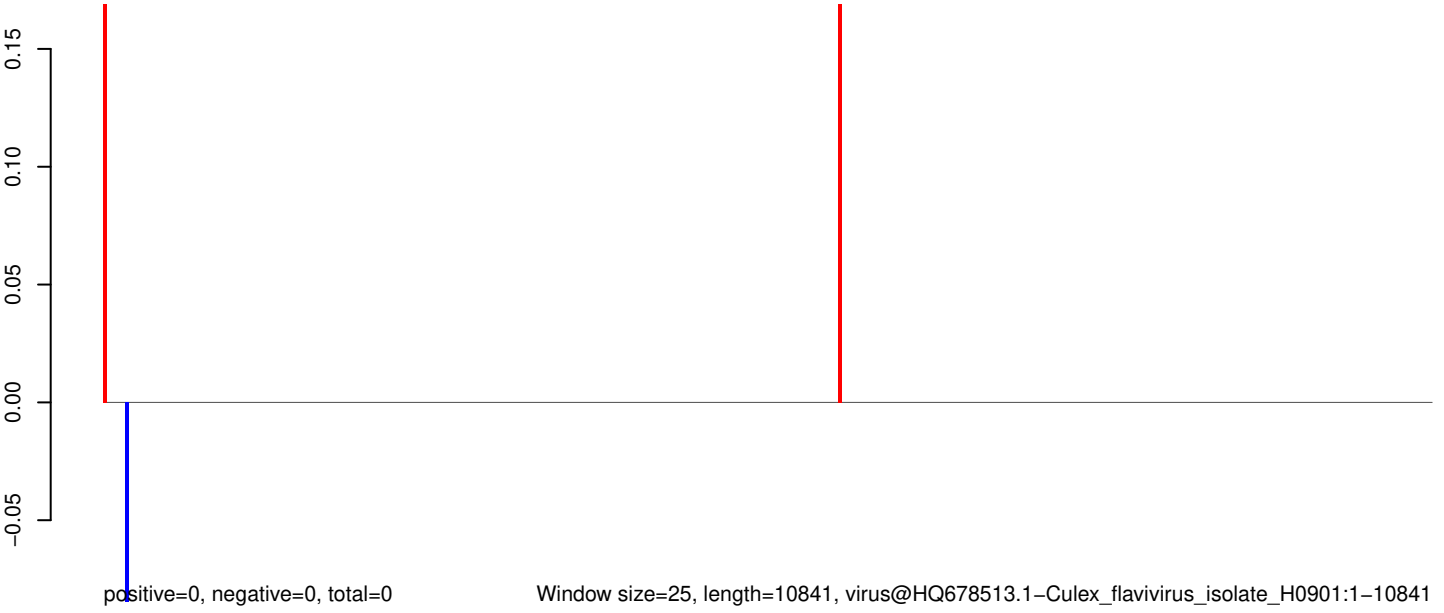
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



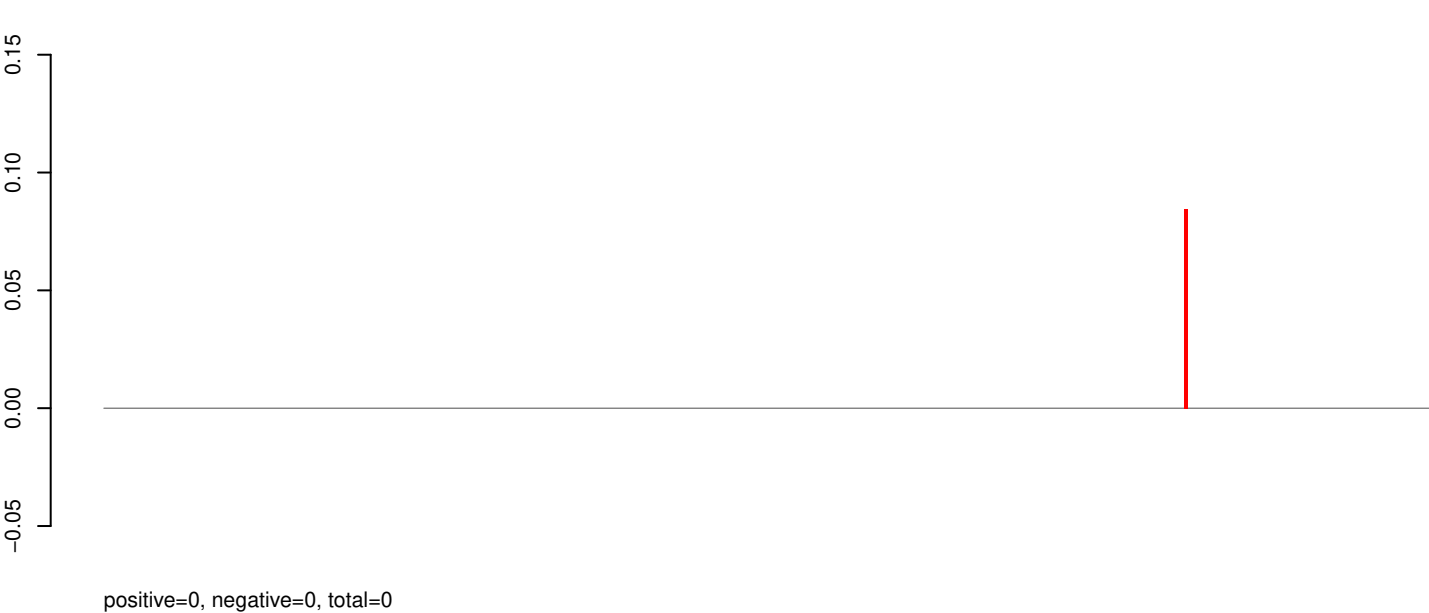
Window size=25, length=10841, virus@HQ678513.1-Culex_flavivirus_isolate_H0901:1-10841

0 2000 4000 6000 8000 10000

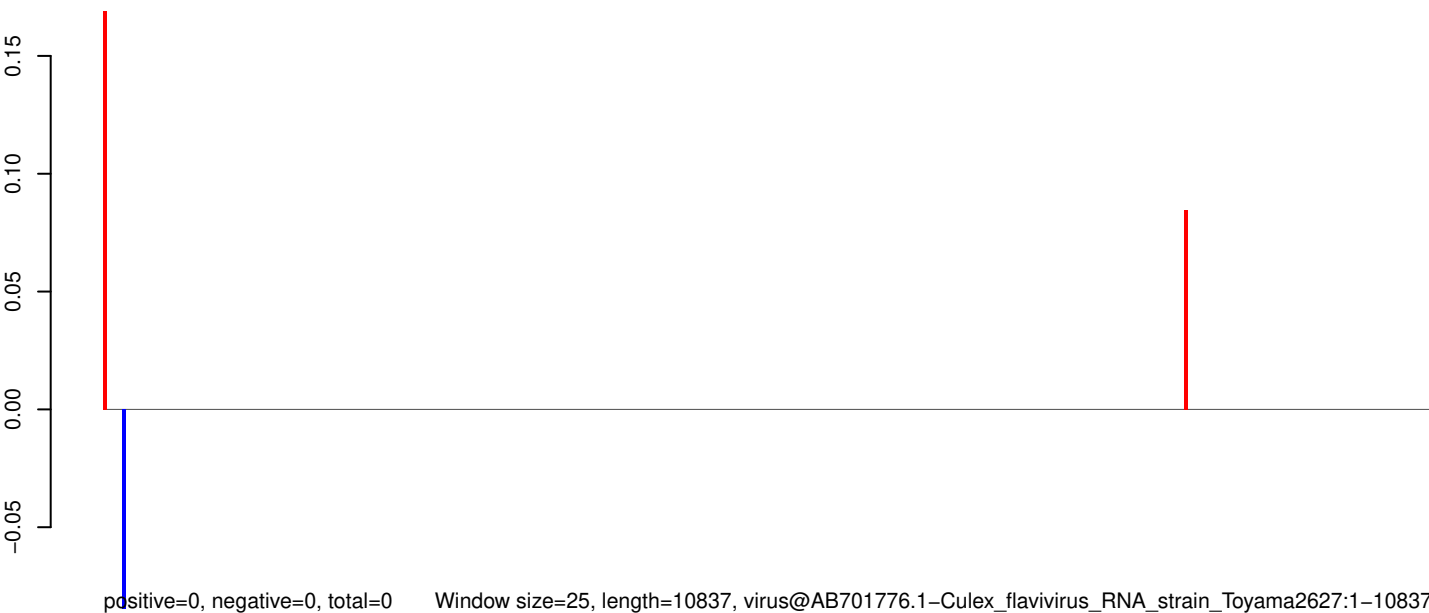
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



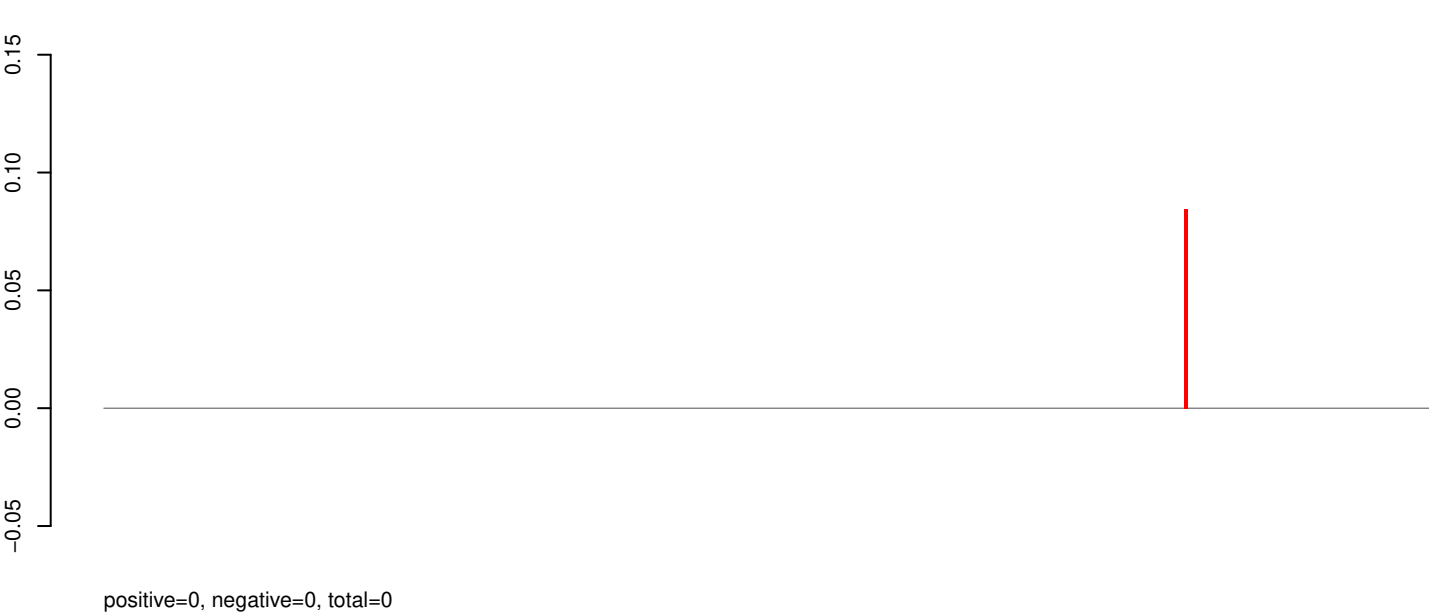
Window size=25, length=10837, virus@AB701776.1-Culex_flavivirus_RNA_strain_Toyama2627:1-10837

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



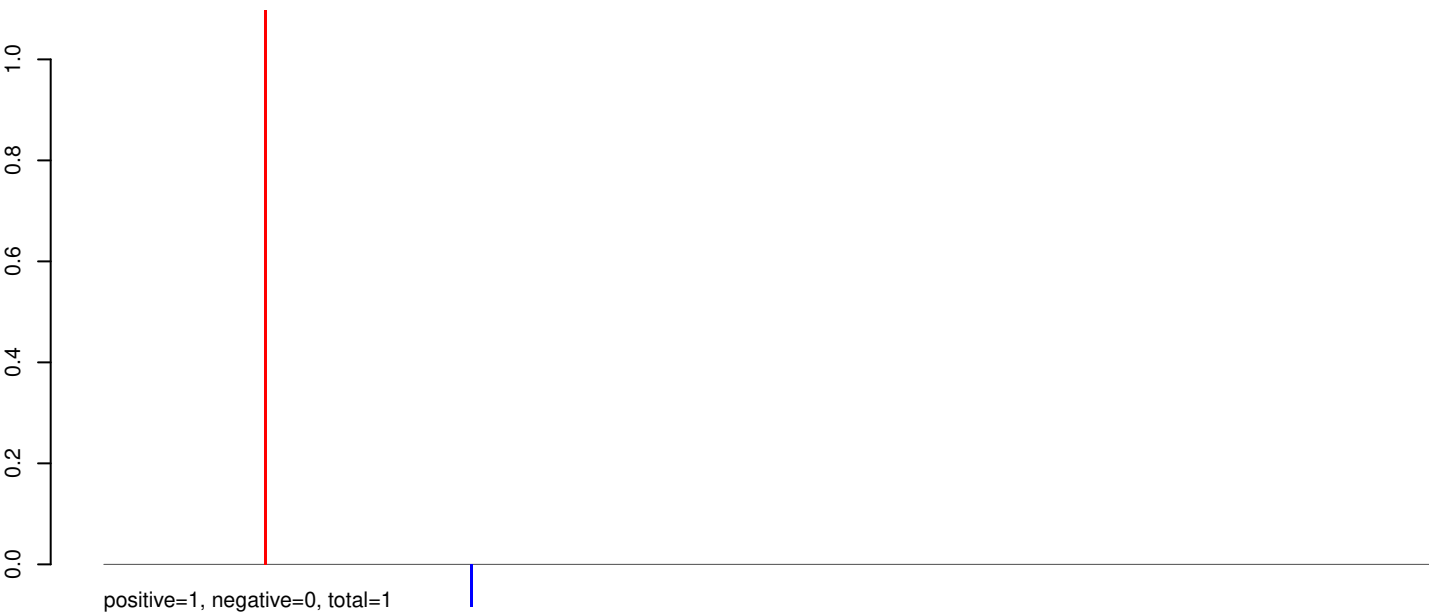
AeAeg_Aag2_BTV1_AK.rep



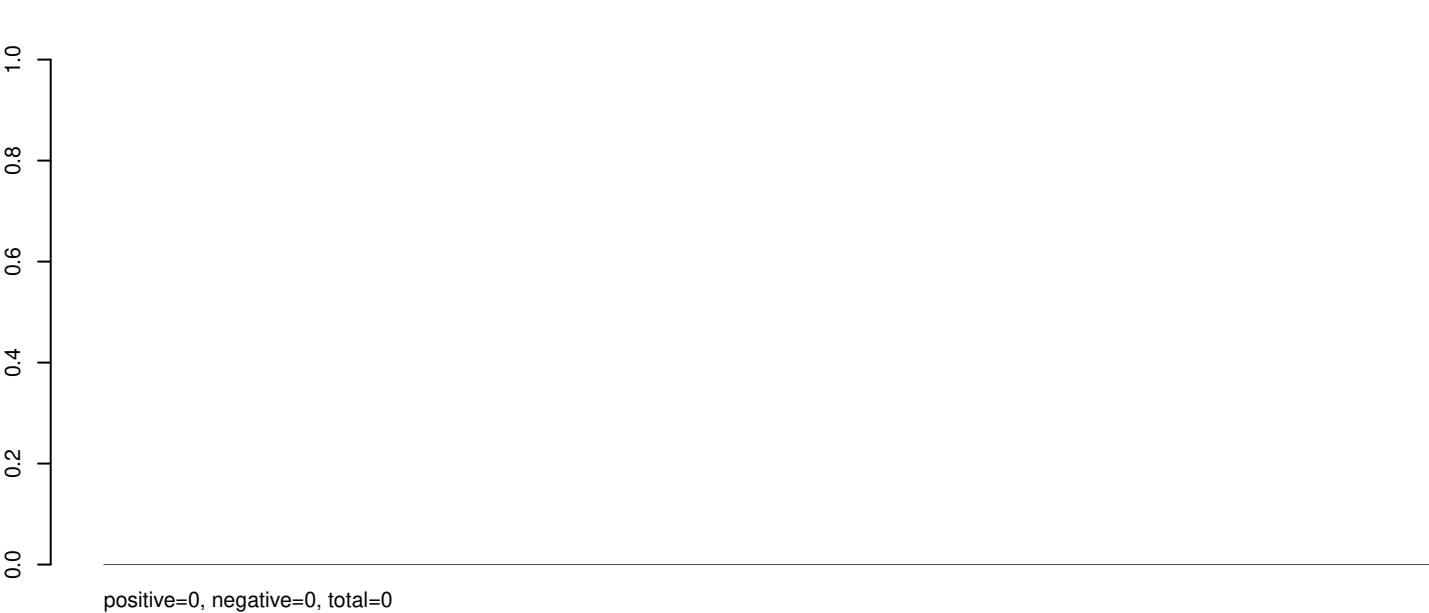
Window size=25, length=10837, virus@AB262759.2-Culex_flavivirus_genomic_RNA_strain_Tokyo:1-10837

0 2000 4000 6000 8000 10000

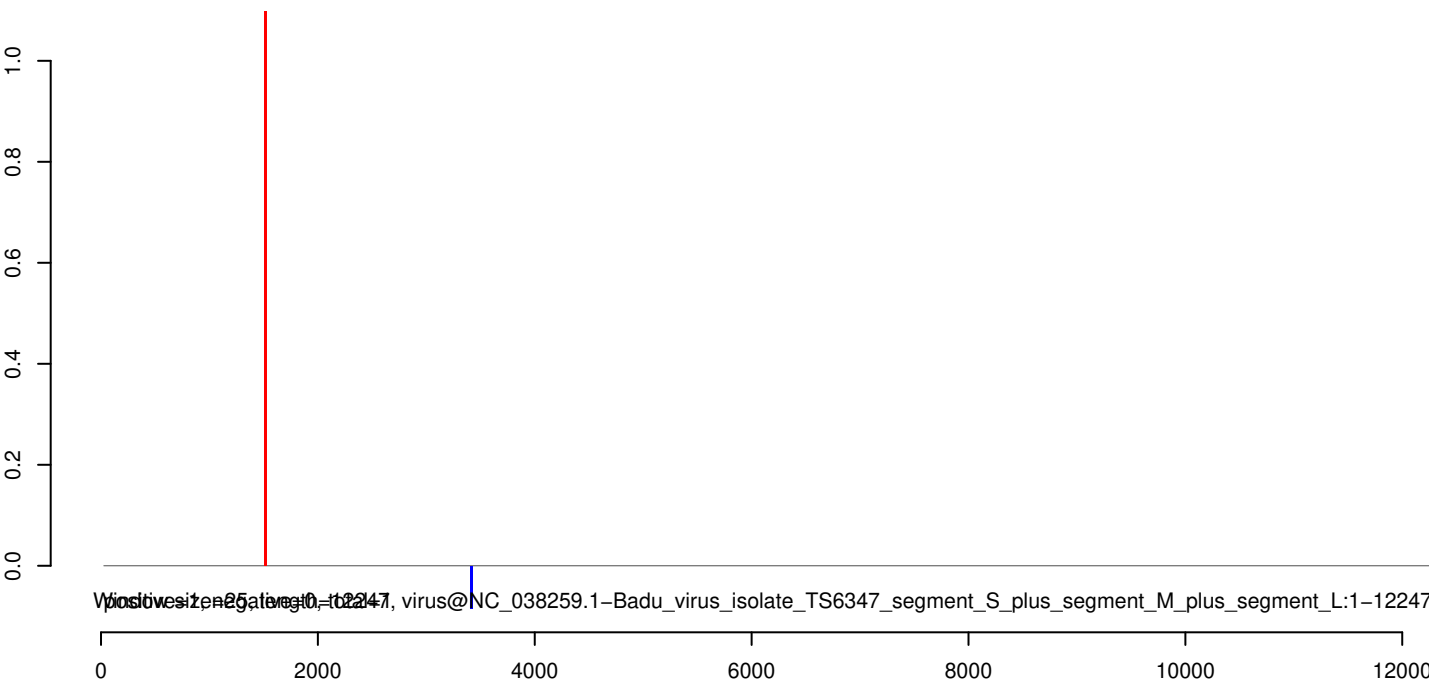
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



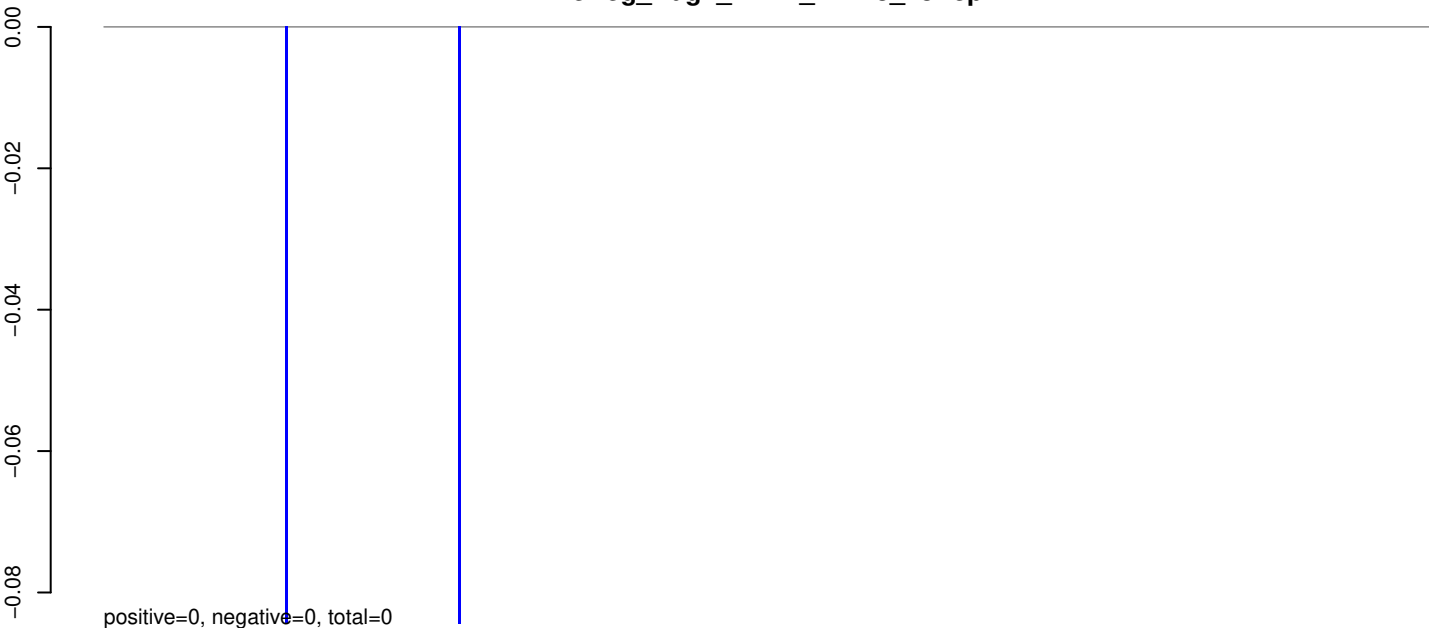
AeAeg_Aag2_BTV1_AK.rep



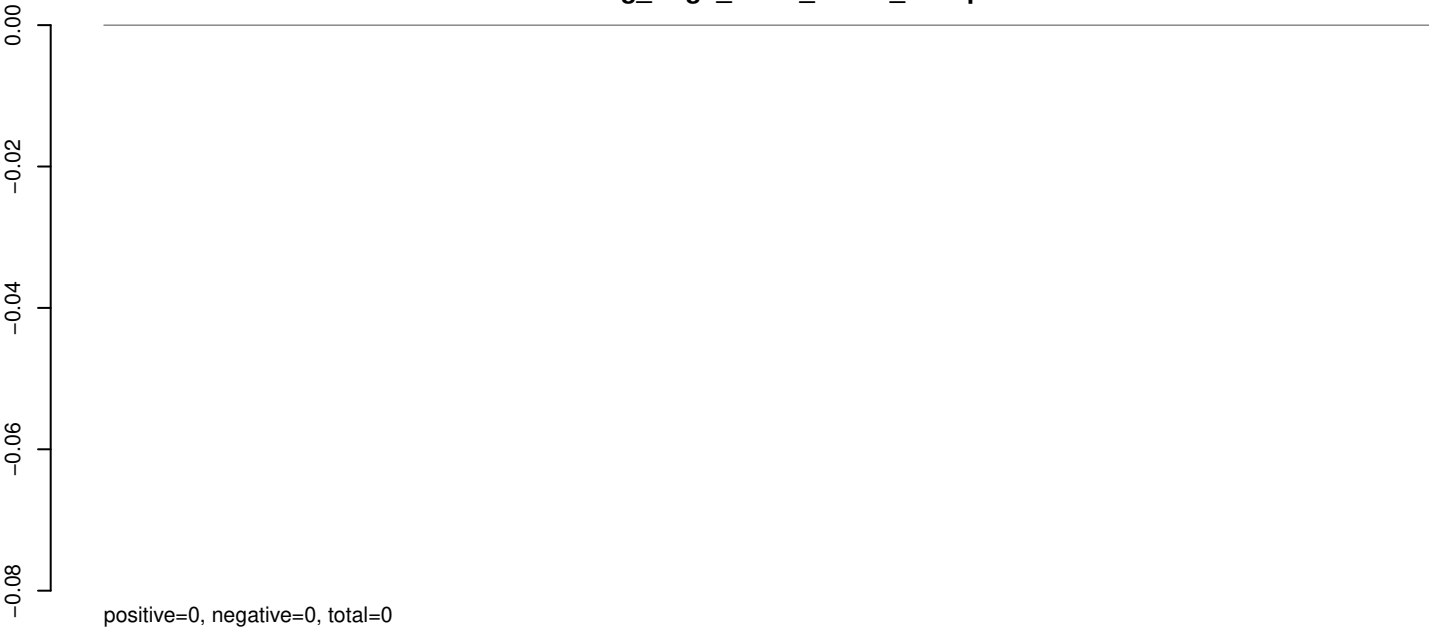
positive=1, negative=0, total=1, virus@NC_038259.1-Badu_virus_isolate_TS6347_segment_S_plus_segment_M_plus_segment_L:1-12247

0 2000 4000 6000 8000 10000 12000

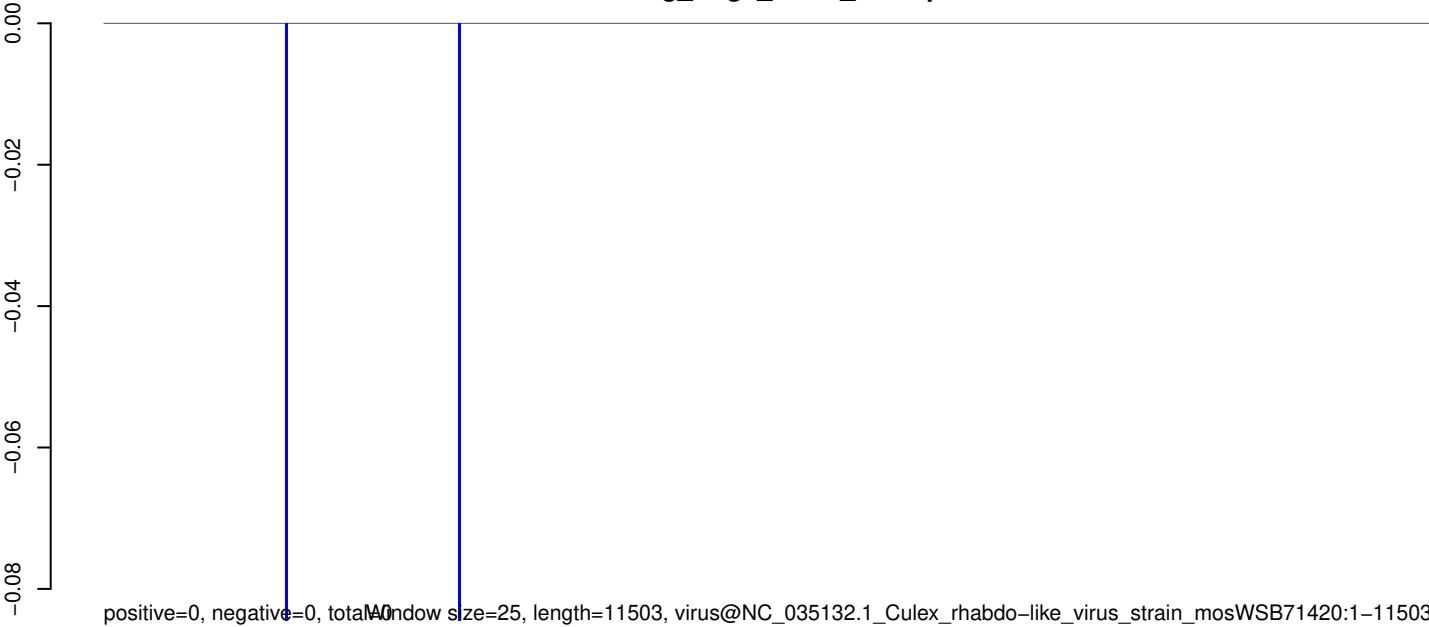
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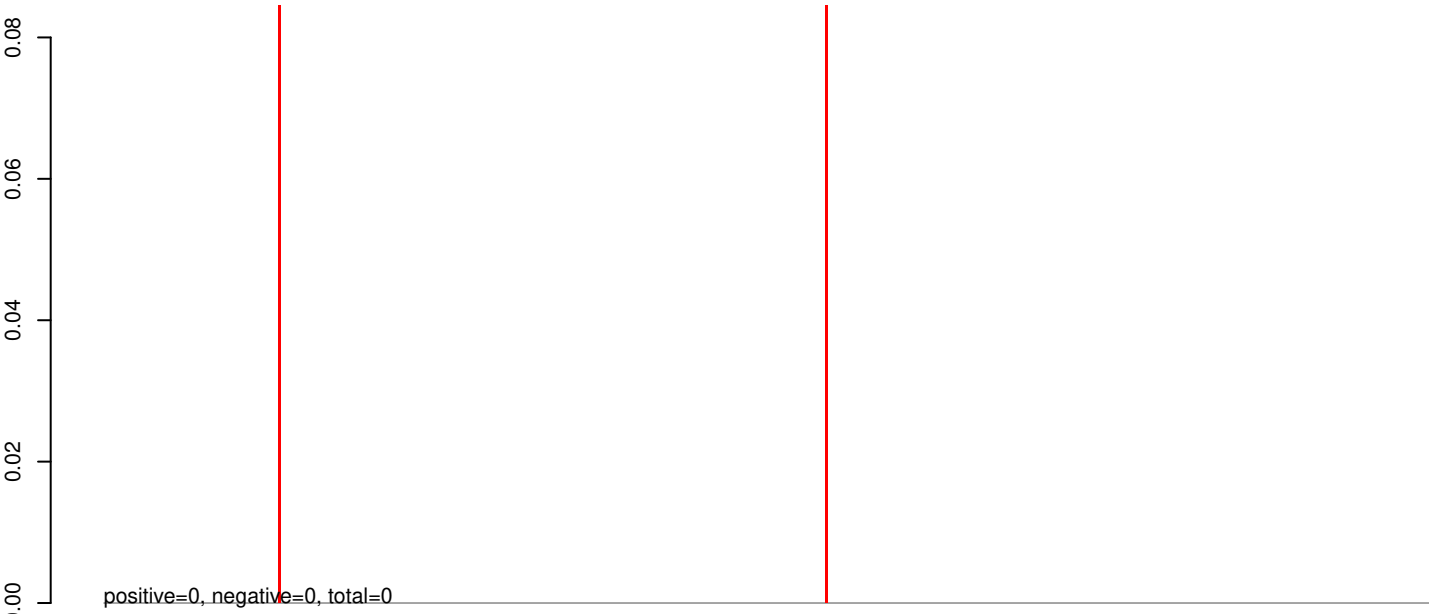
AeAeg_Aag2_BTV1_AK.24_35.rep



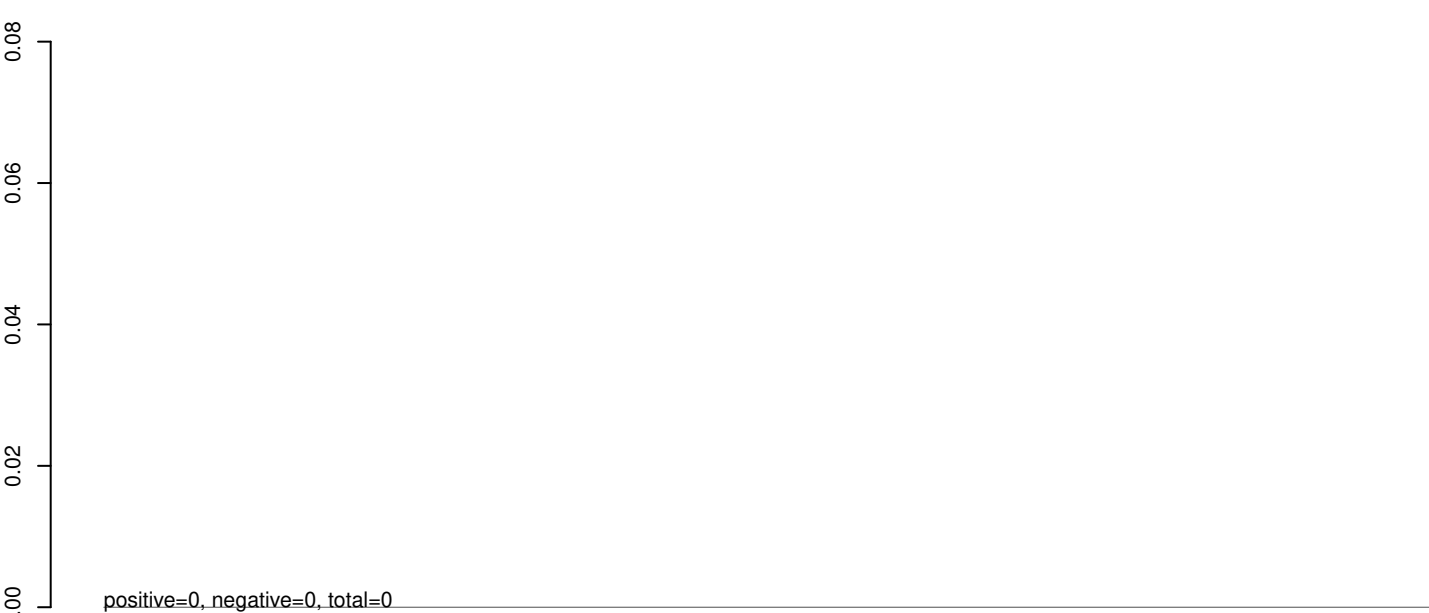
AeAeg_Aag2_BTV1_AK.rep



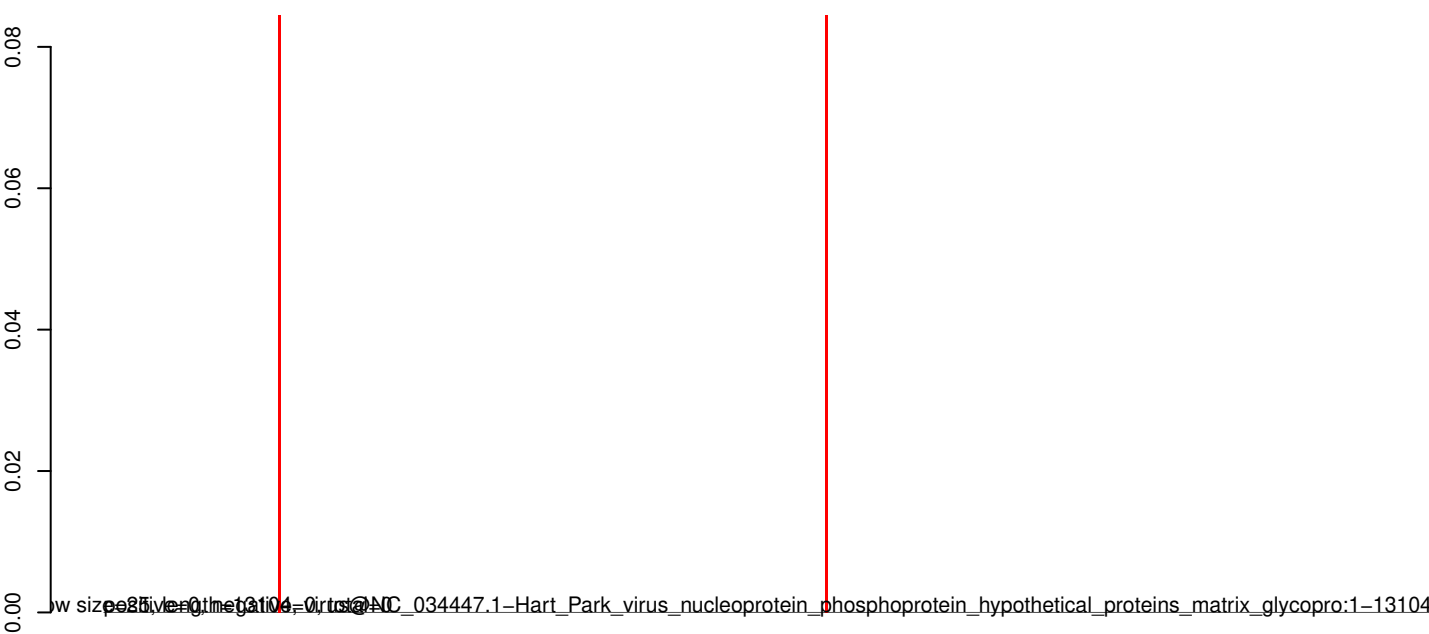
AeAeg_Aag2_BTV1_AK.18_23.rep



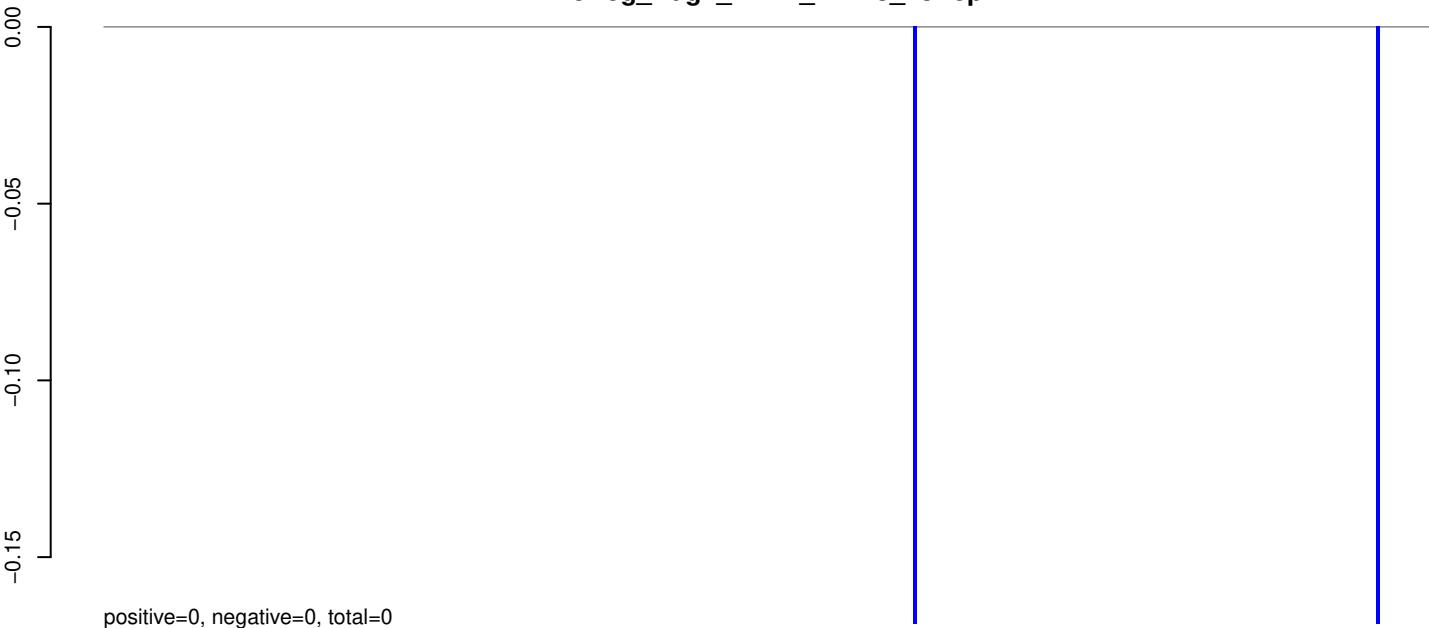
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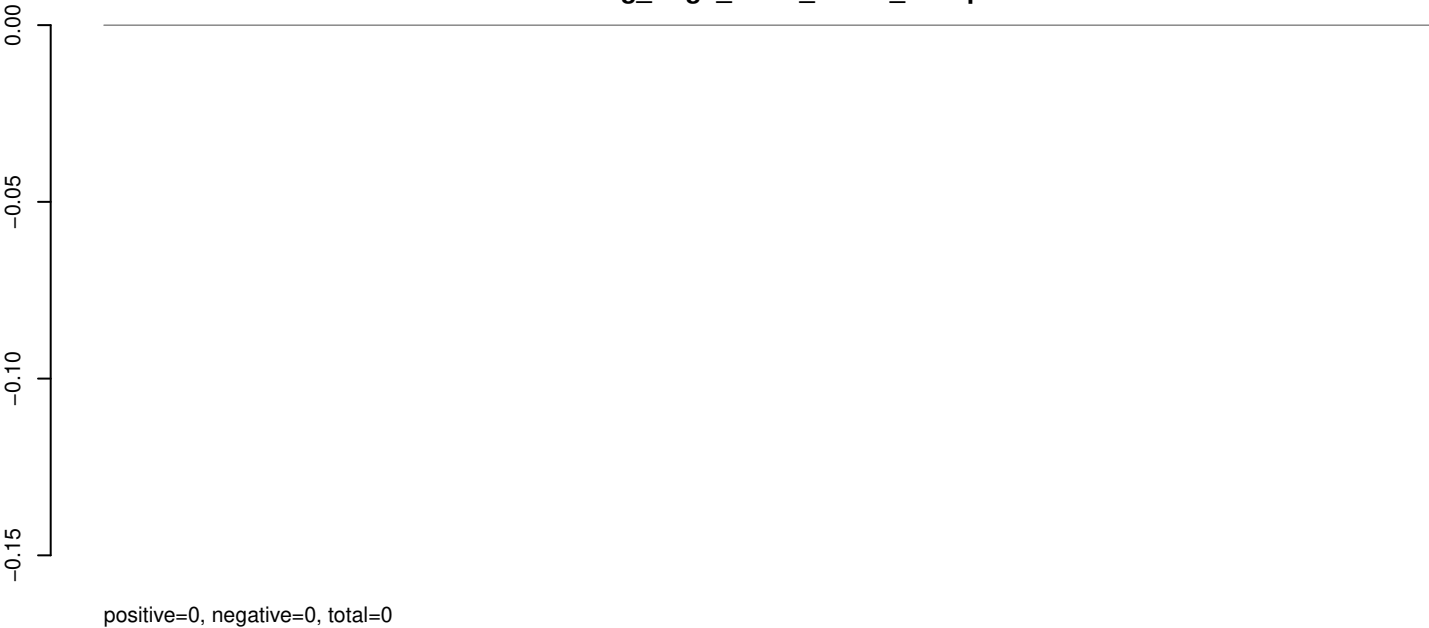
AeAeg_Aag2_BTV1_AK.rep



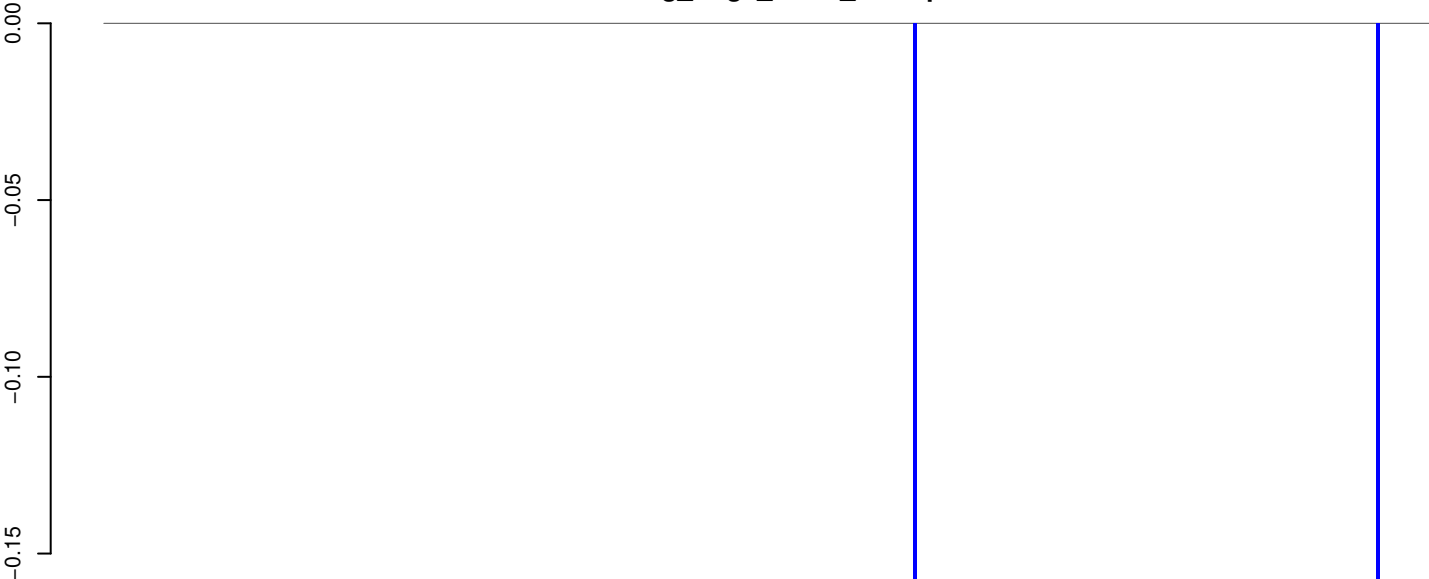
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



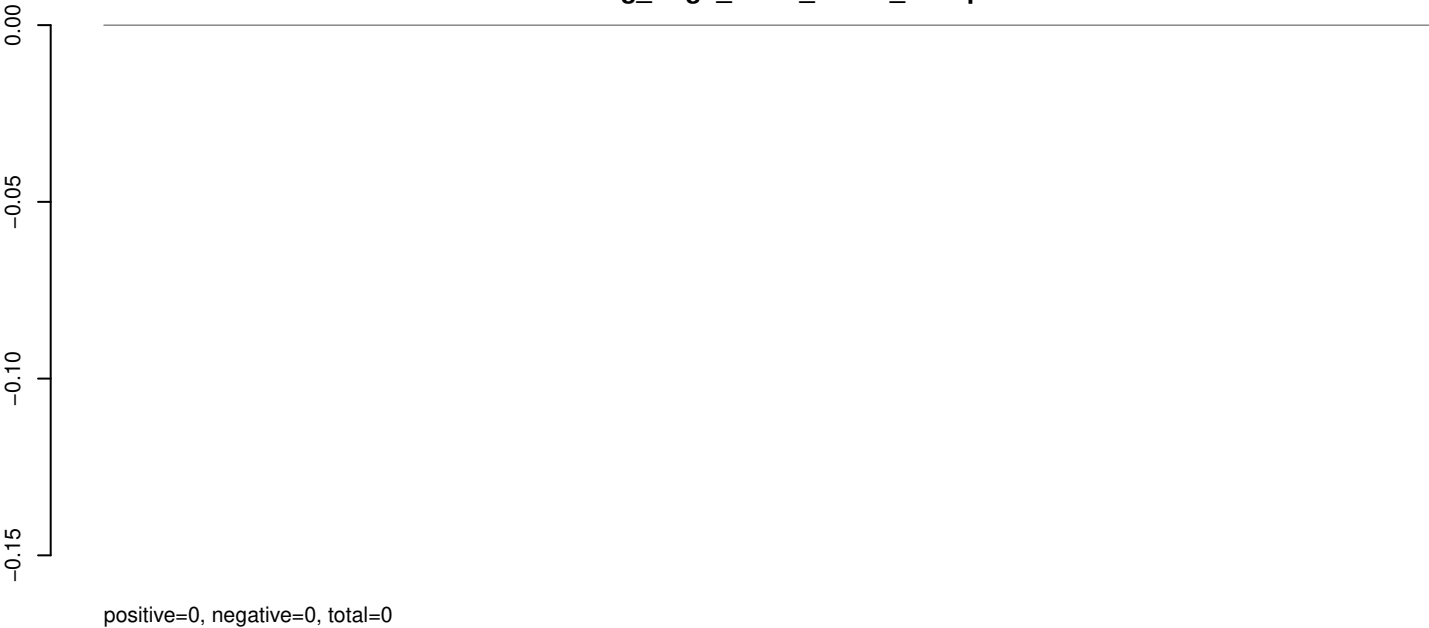
Window size=25, length=10173, virus@NC_033699.1-Jugra_virus_strain_P-9-314_polyprotein_gene:1-10173

0 2000 4000 6000 8000 10000

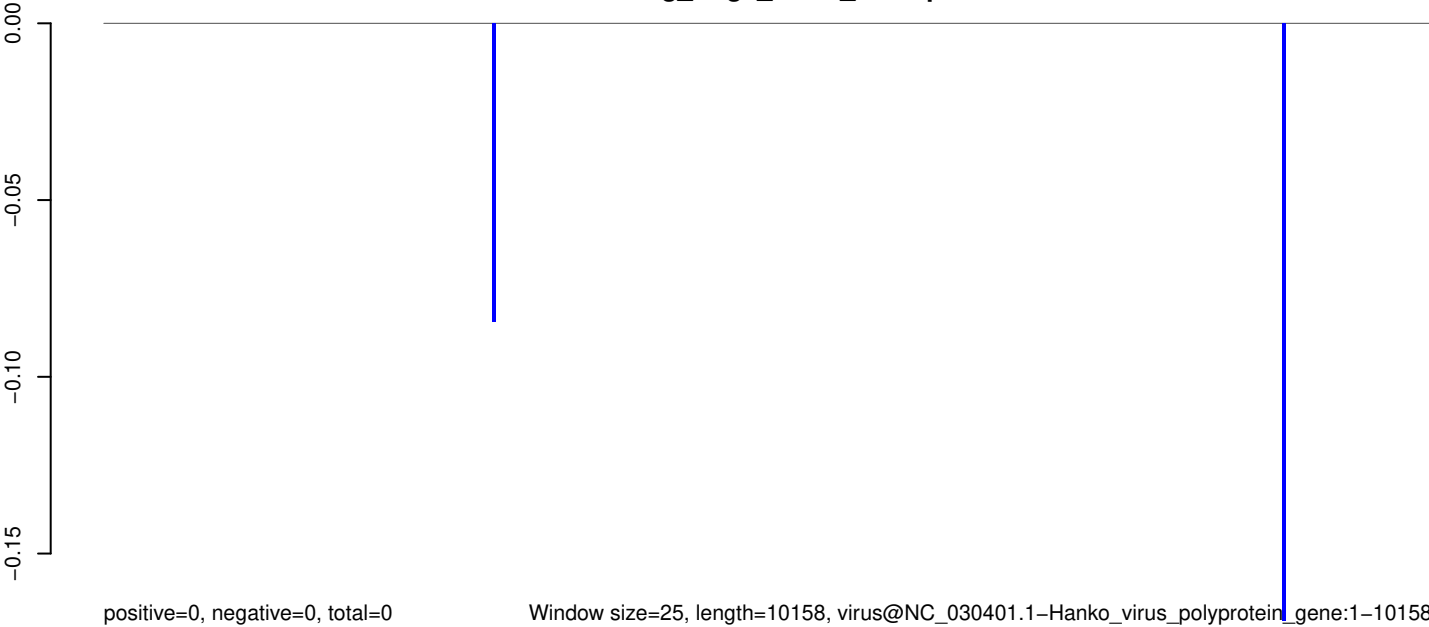
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



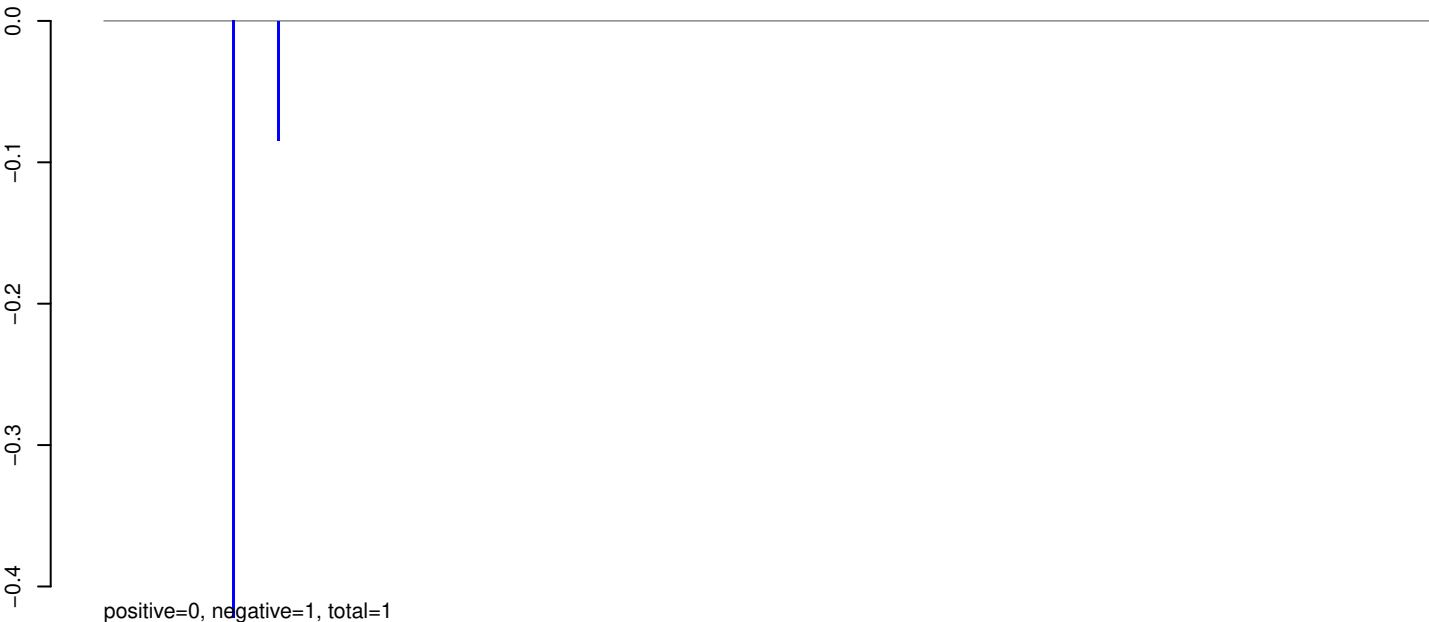
AeAeg_Aag2_BTV1_AK.rep



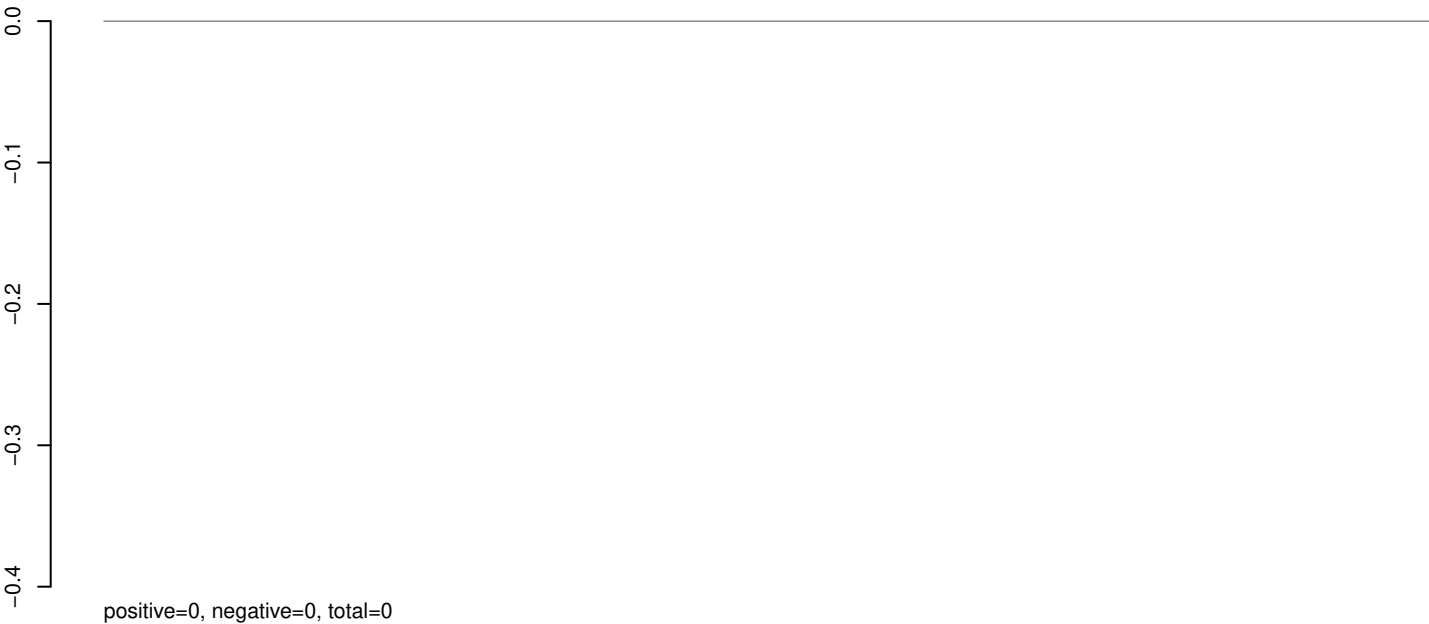
Window size=25, length=10158, virus@NC_030401.1-Hanko_virus_polyprotein_gene:1-10158

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep

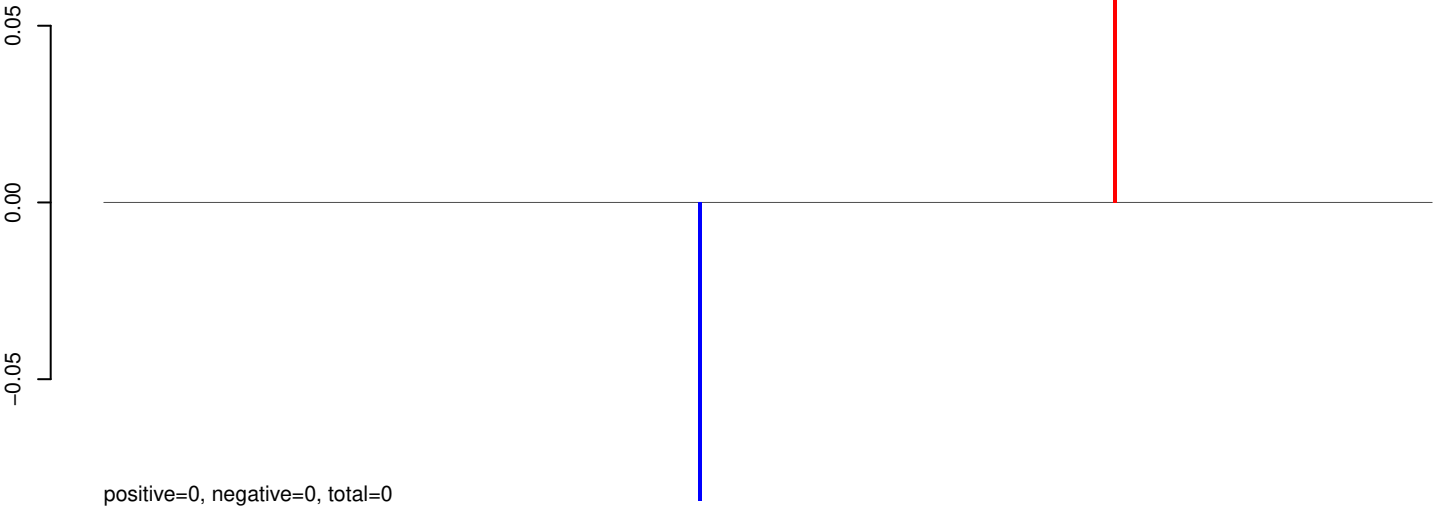


AeAeg_Aag2_BTV1_AK.rep

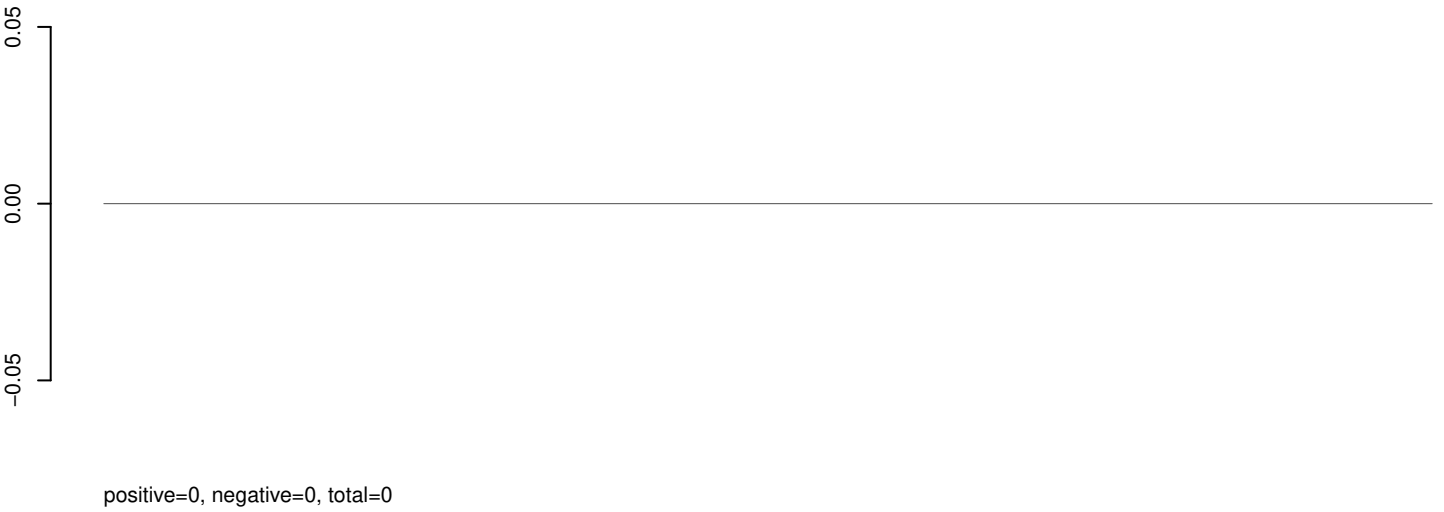


Window size=25, length=14717, virus@NC_017685.1-Obodhiang_virus:1-14717

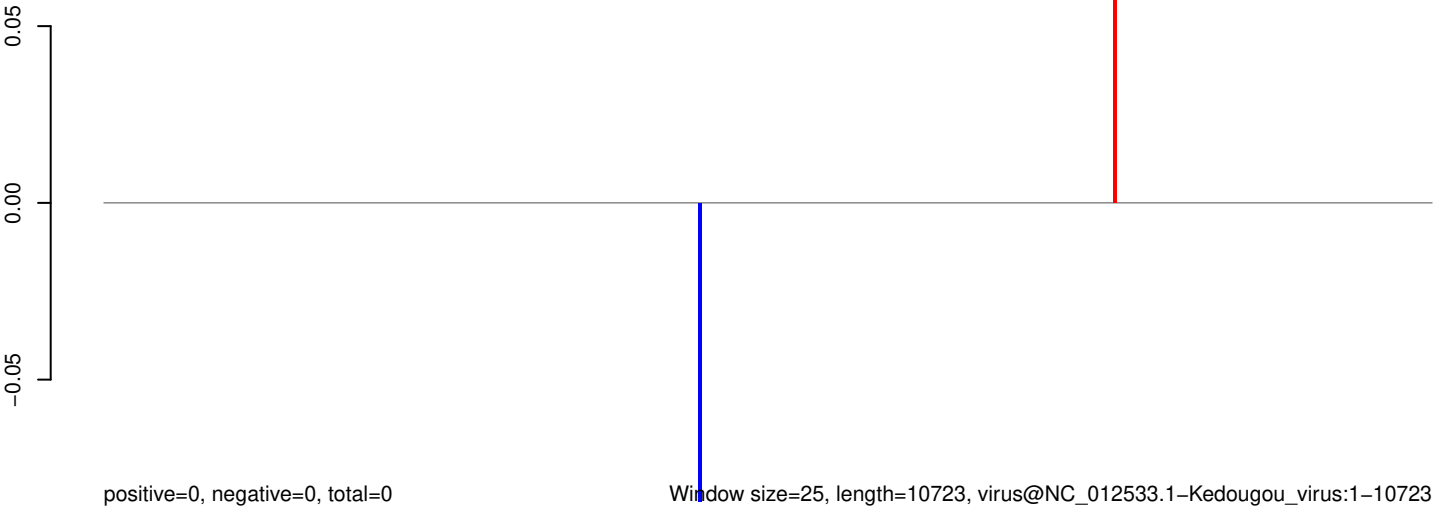
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



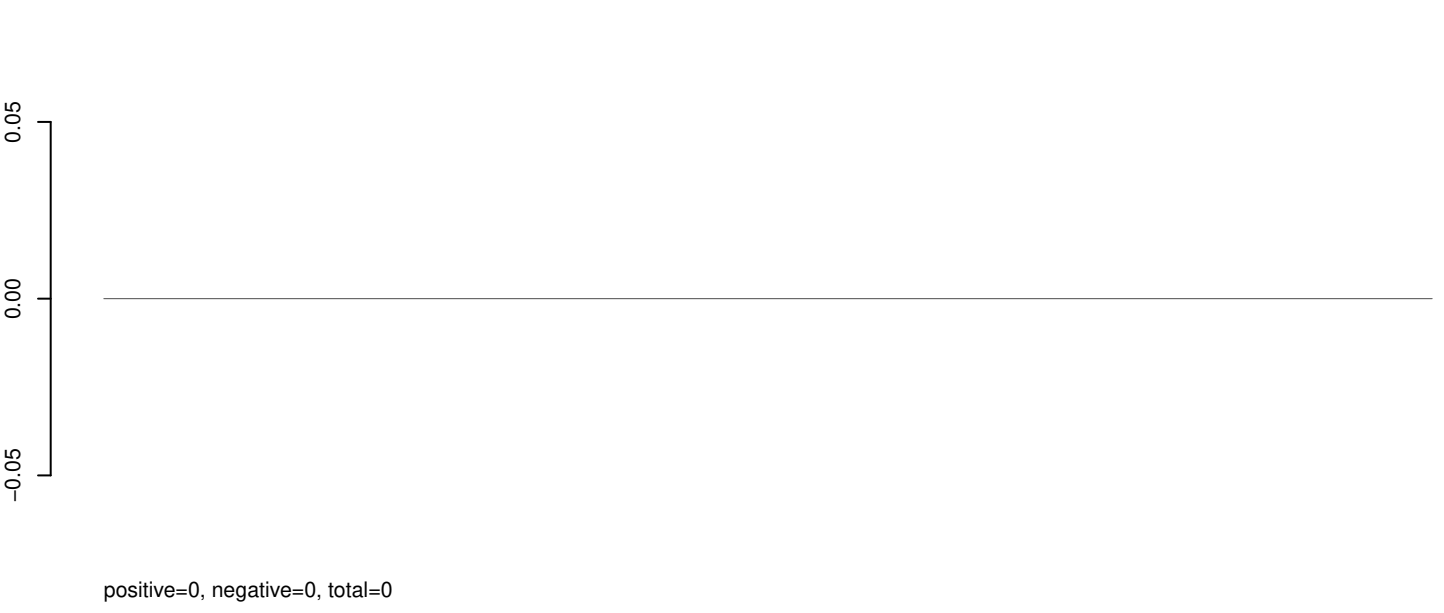
Window size=25, length=10723, virus@NC_012533.1-Kedougou_virus:1-10723

0 2000 4000 6000 8000 10000

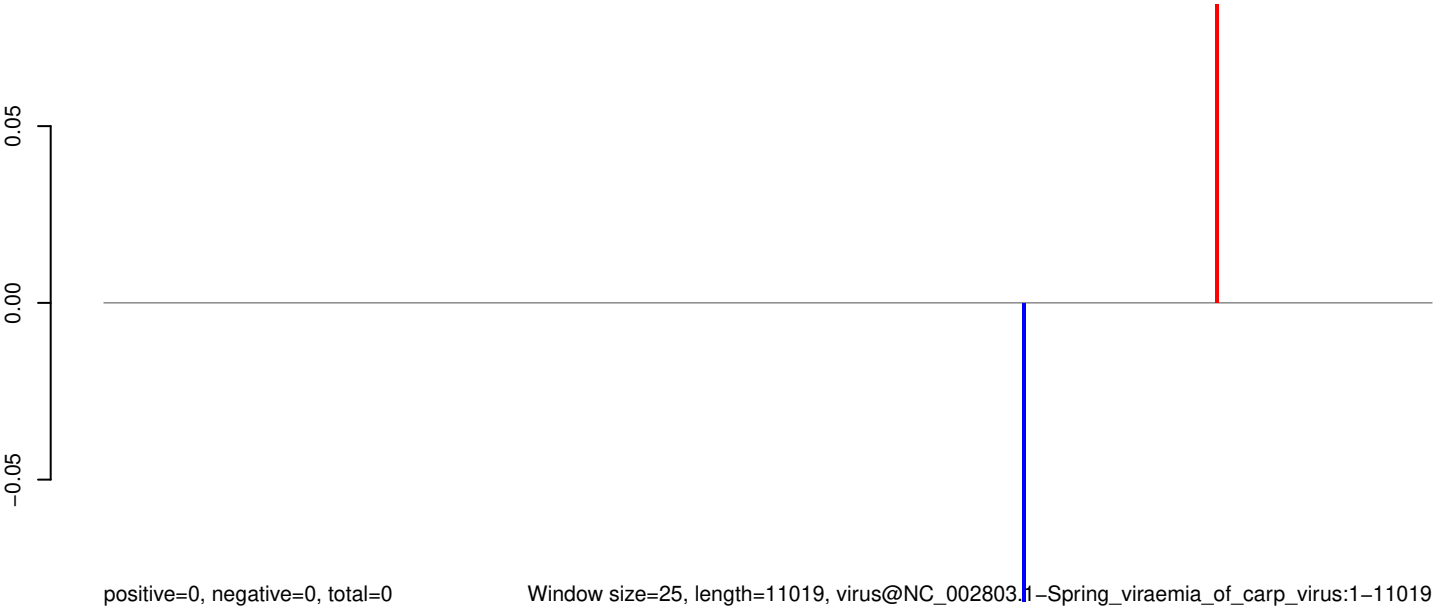
AeAeg_Aag2_BTV1_AK.18_23.rep



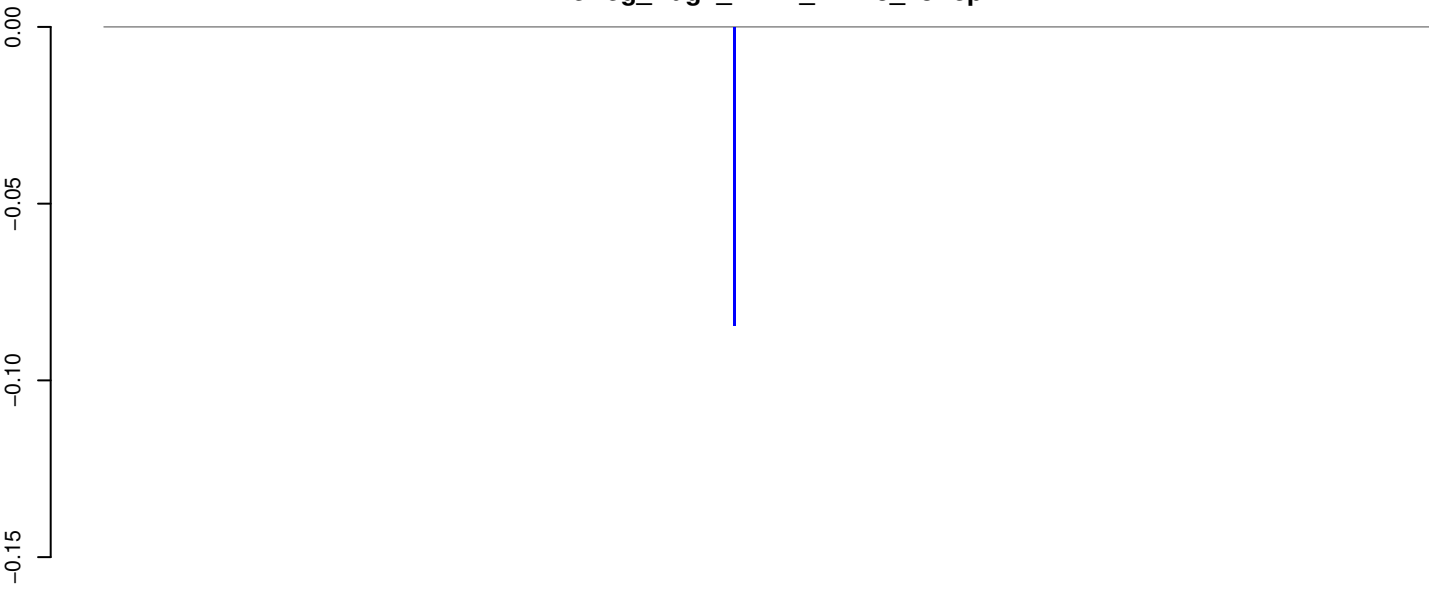
AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep

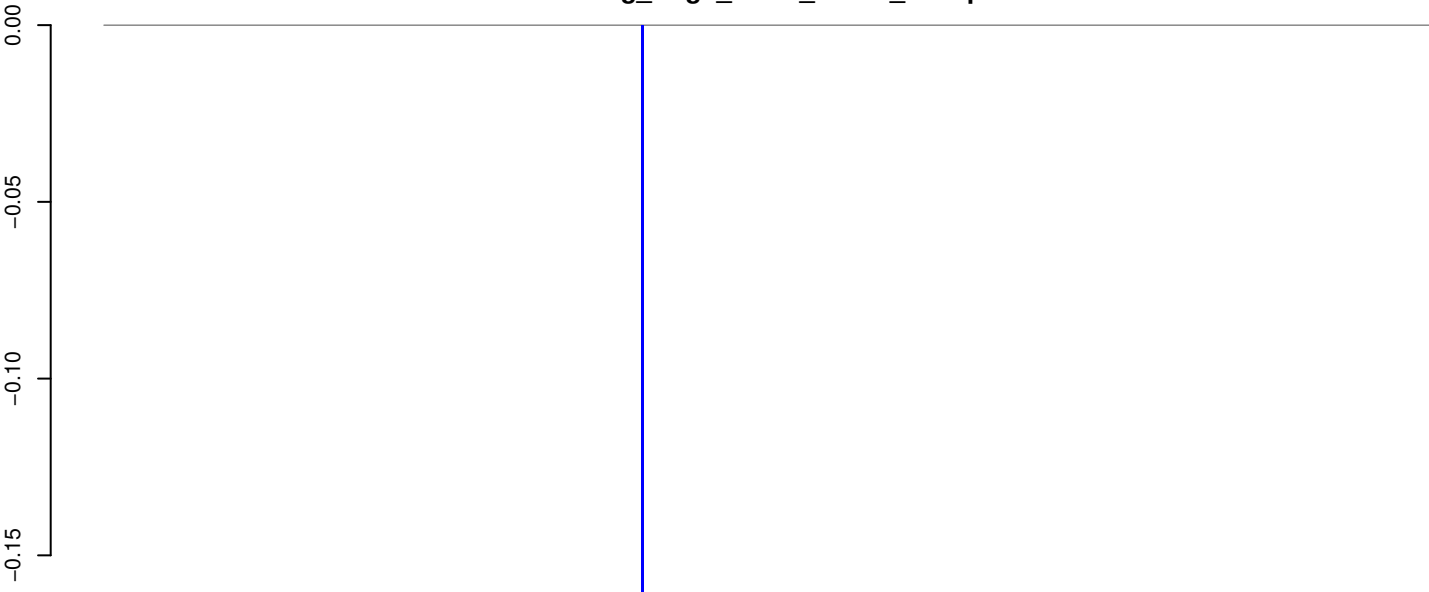


AeAeg_Aag2_BTV1_AK.18_23.rep



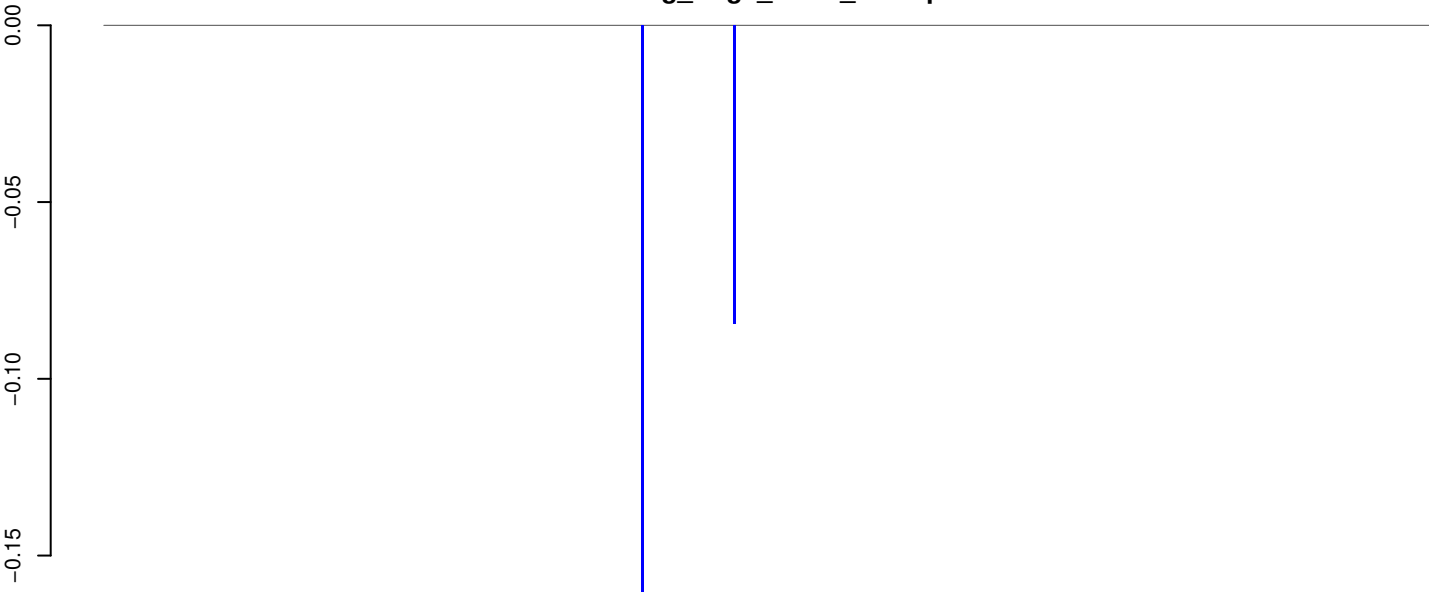
positive=0, negative=0, total=0

AeAeg_Aag2_BTV1_AK.24_35.rep



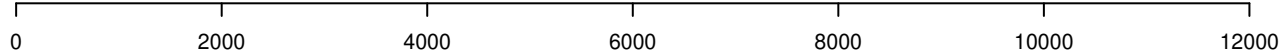
positive=0, negative=0, total=0

AeAeg_Aag2_BTV1_AK.rep

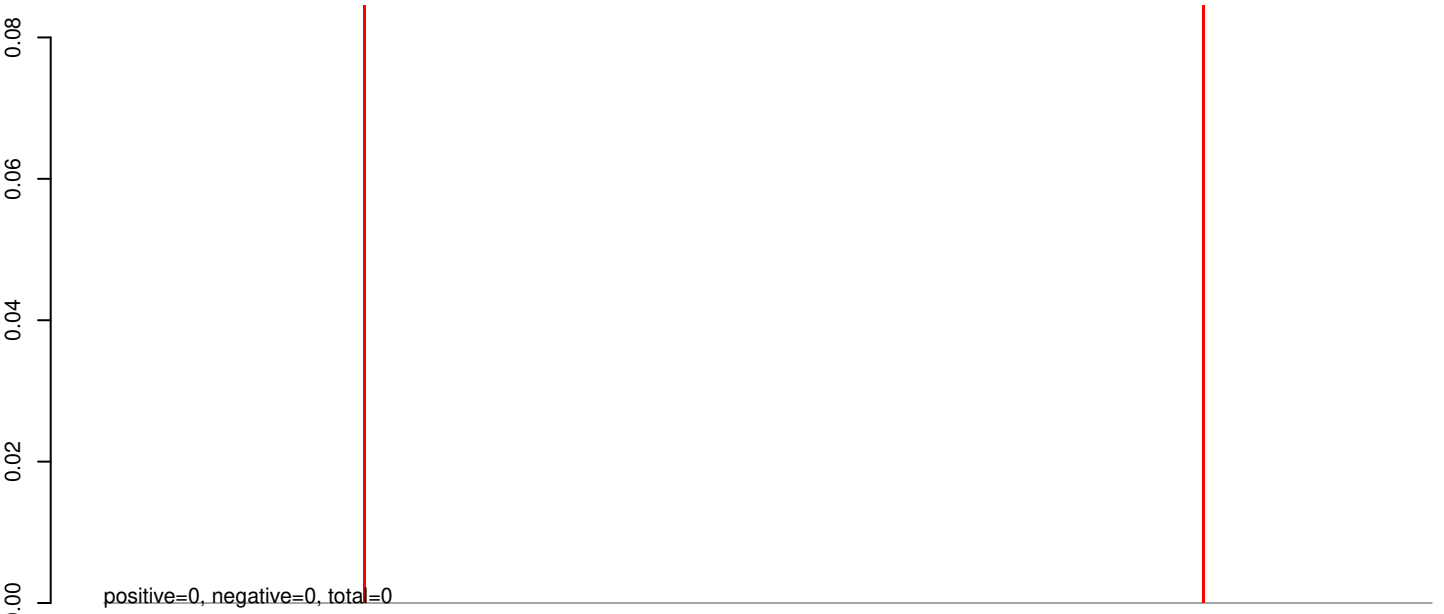


positive=0, negative=0, total=0

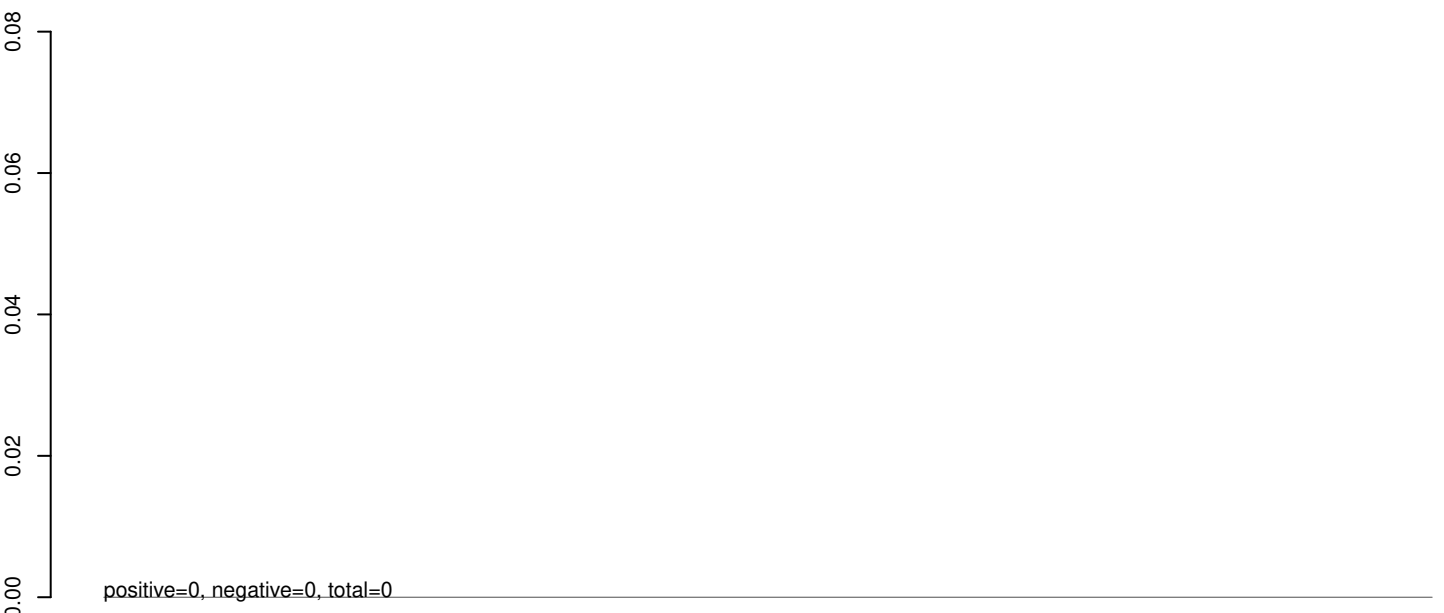
Window size=25, length=12916, virus@MH237595.1-Aedes_aegypti_anphevirus_strain_1a:1-12916



AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



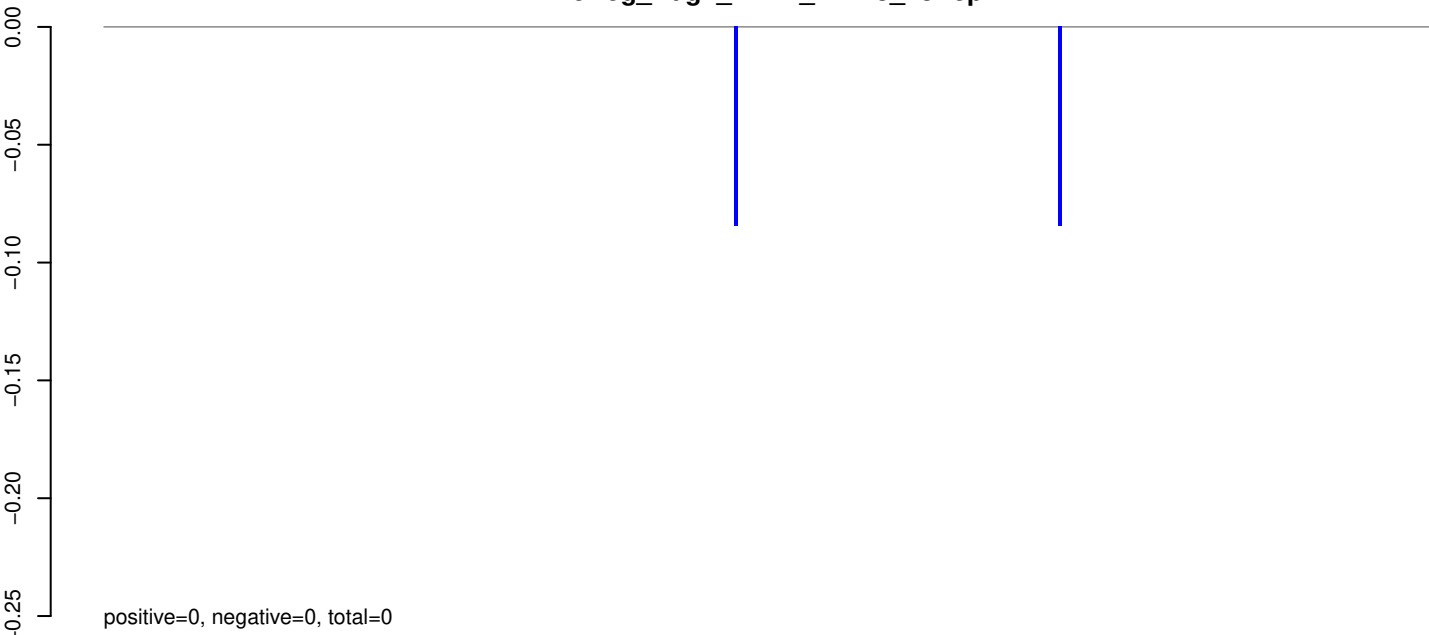
AeAeg_Aag2_BTV1_AK.rep



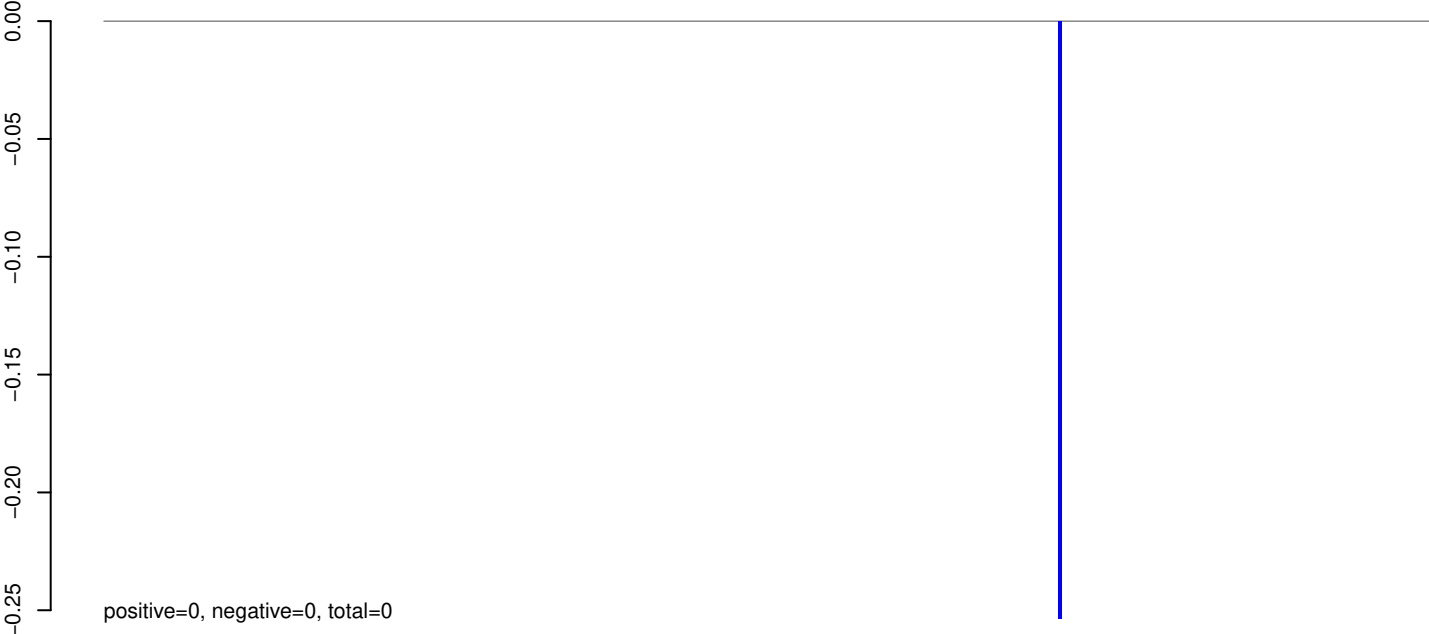
Window size=25, length=11664, virus@MG731558.1_Akabane_virus_isolate_HN10174_nucleoprotein_mRNA:1-11664

0 2000 4000 6000 8000 10000 12000

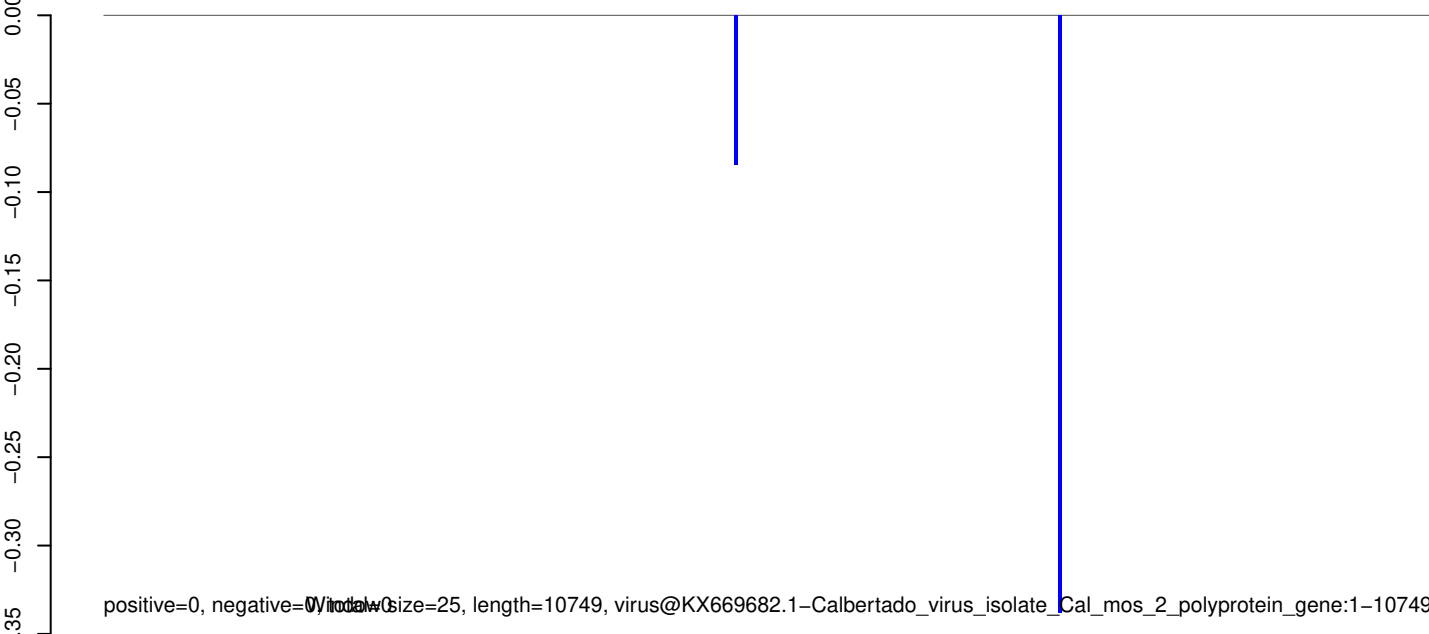
AeAeg_Aag2_BTV1_AK.18_23.rep



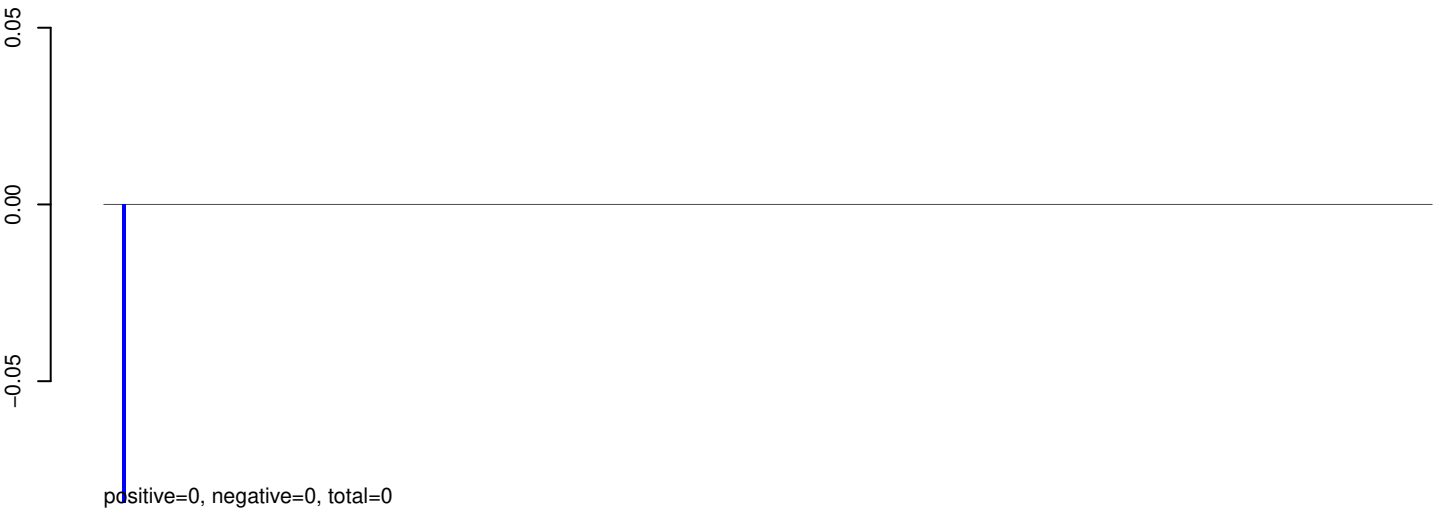
AeAeg_Aag2_BTV1_AK.24_35.rep



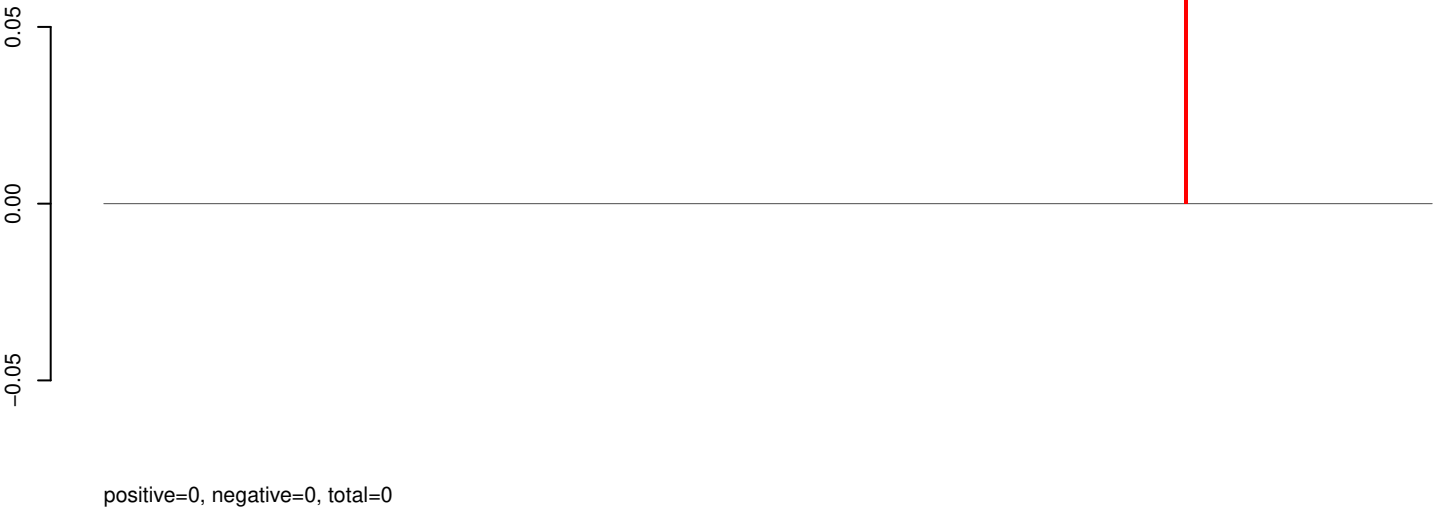
AeAeg_Aag2_BTV1_AK.rep



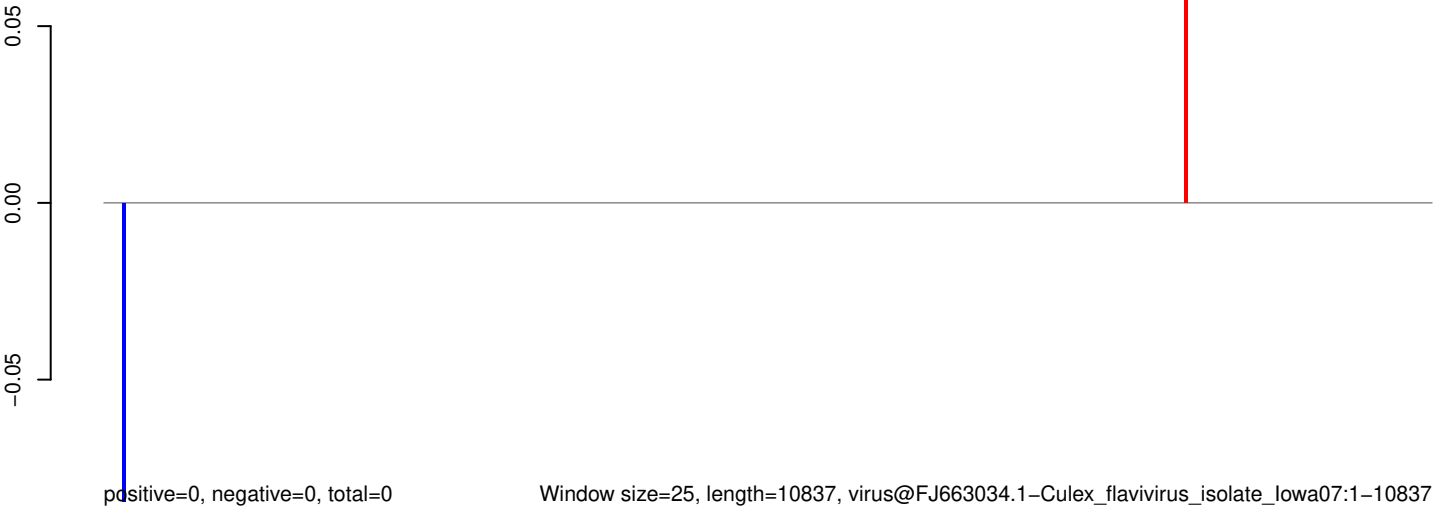
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



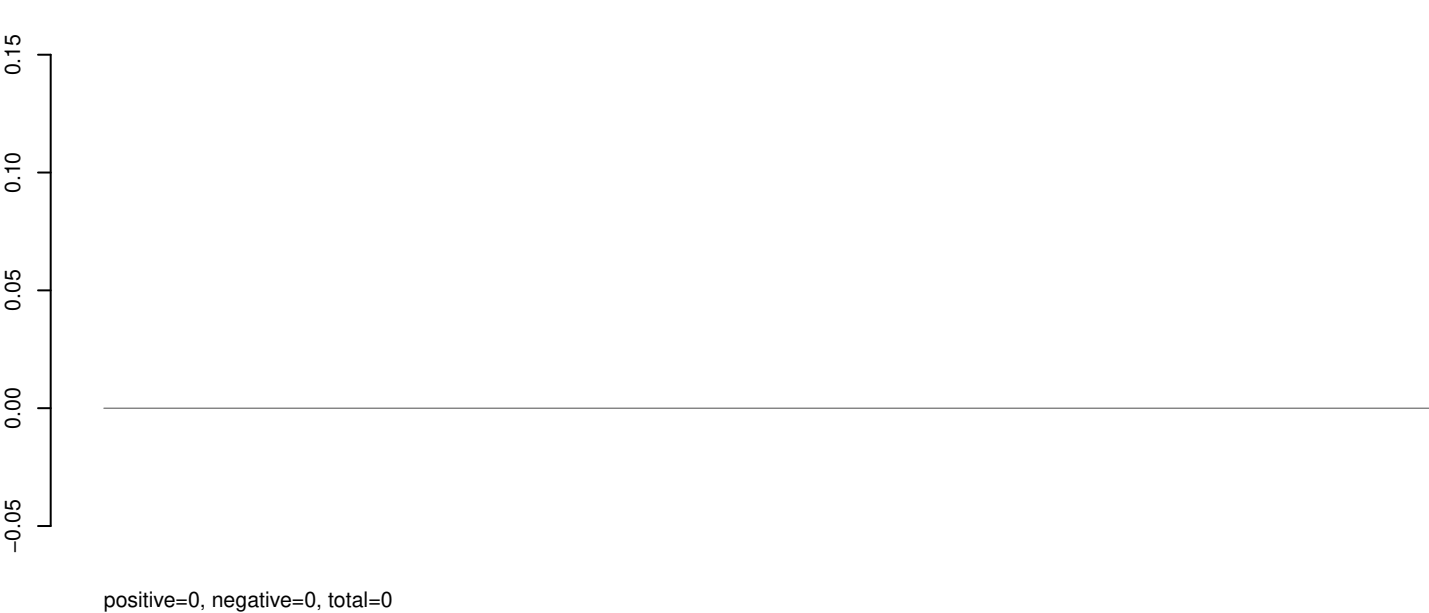
AeAeg_Aag2_BTV1_AK.rep



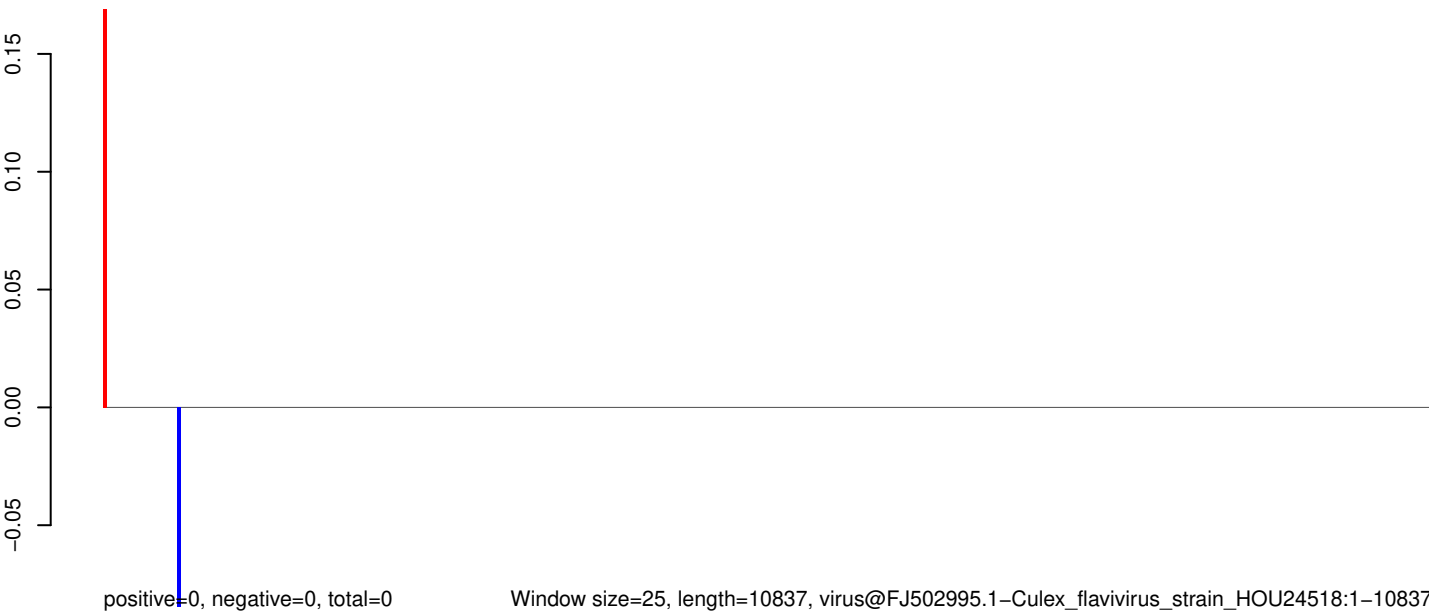
AeAeg_Aag2_BTV1_AK.18_23.rep



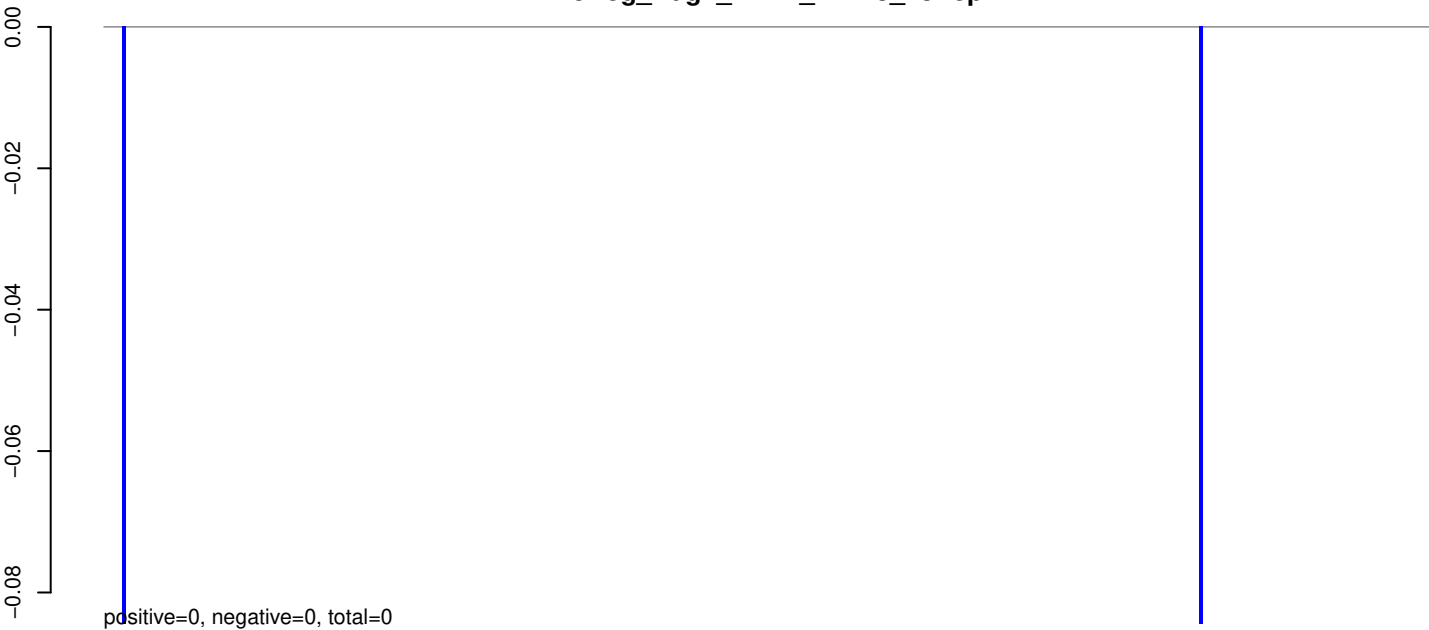
AeAeg_Aag2_BTV1_AK.24_35.rep



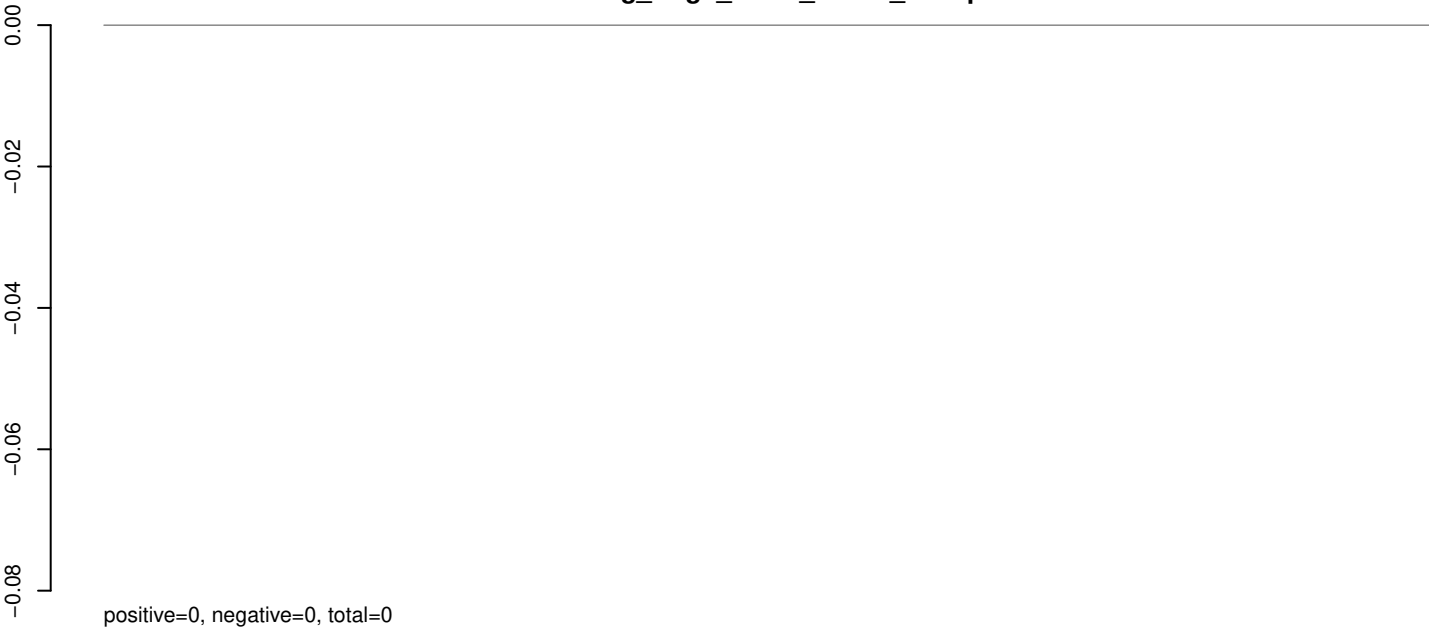
AeAeg_Aag2_BTV1_AK.rep



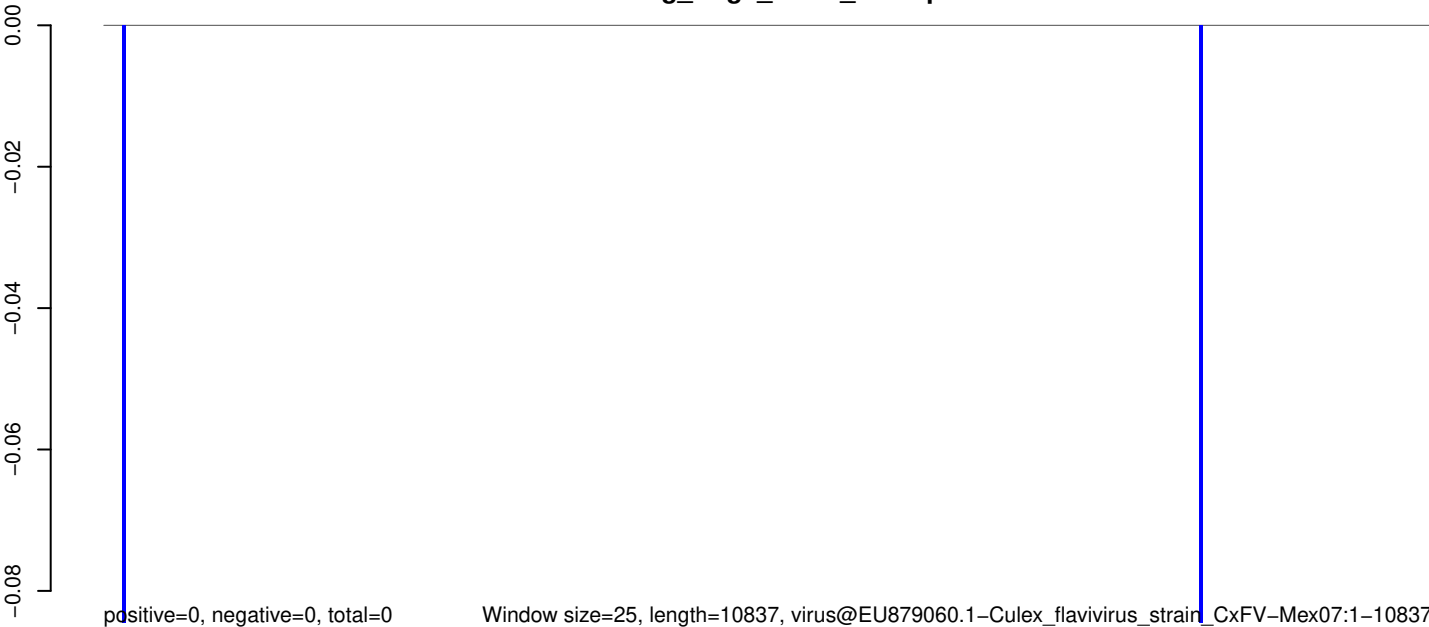
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep

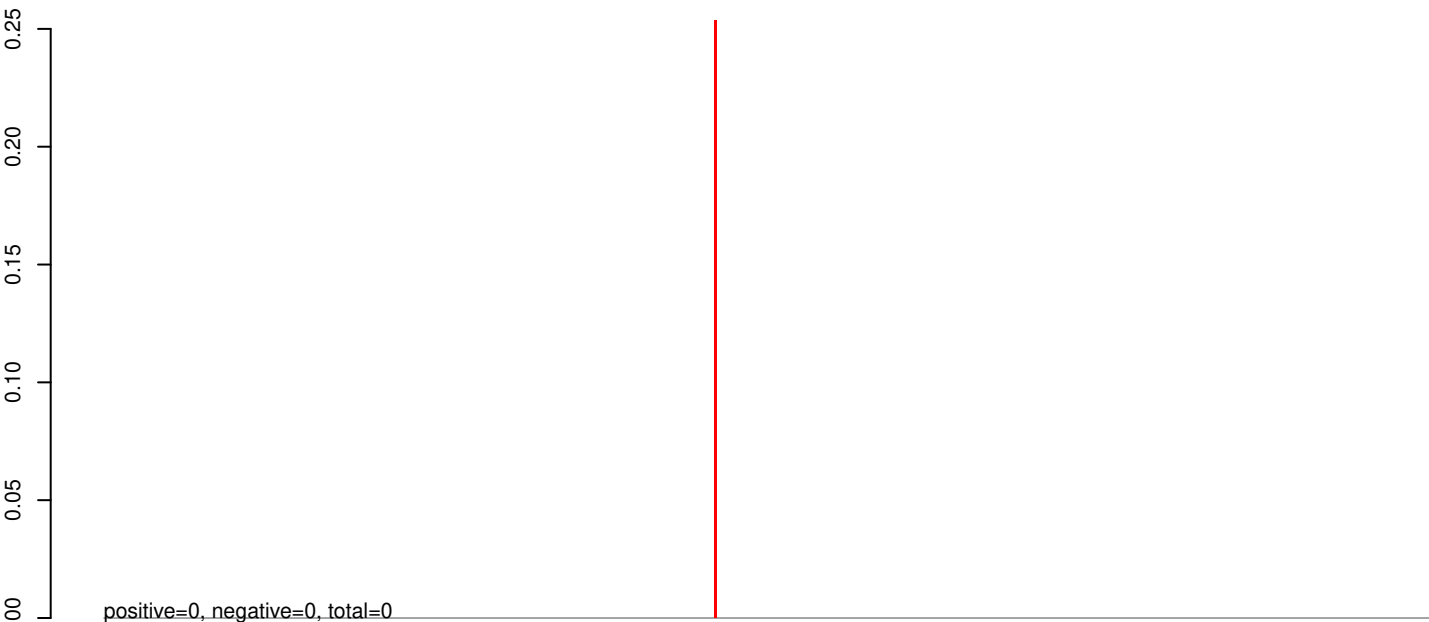


AeAeg_Aag2_BTV1_AK.rep

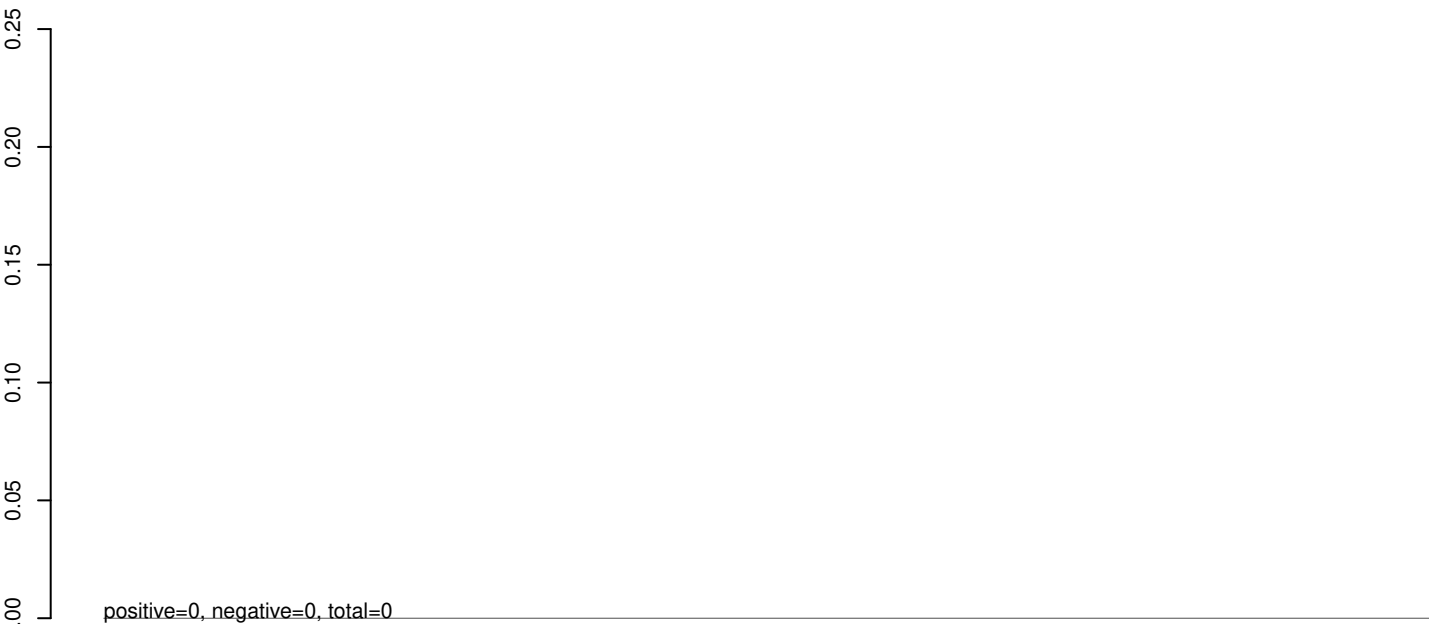


0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV1_AK.18_23.rep



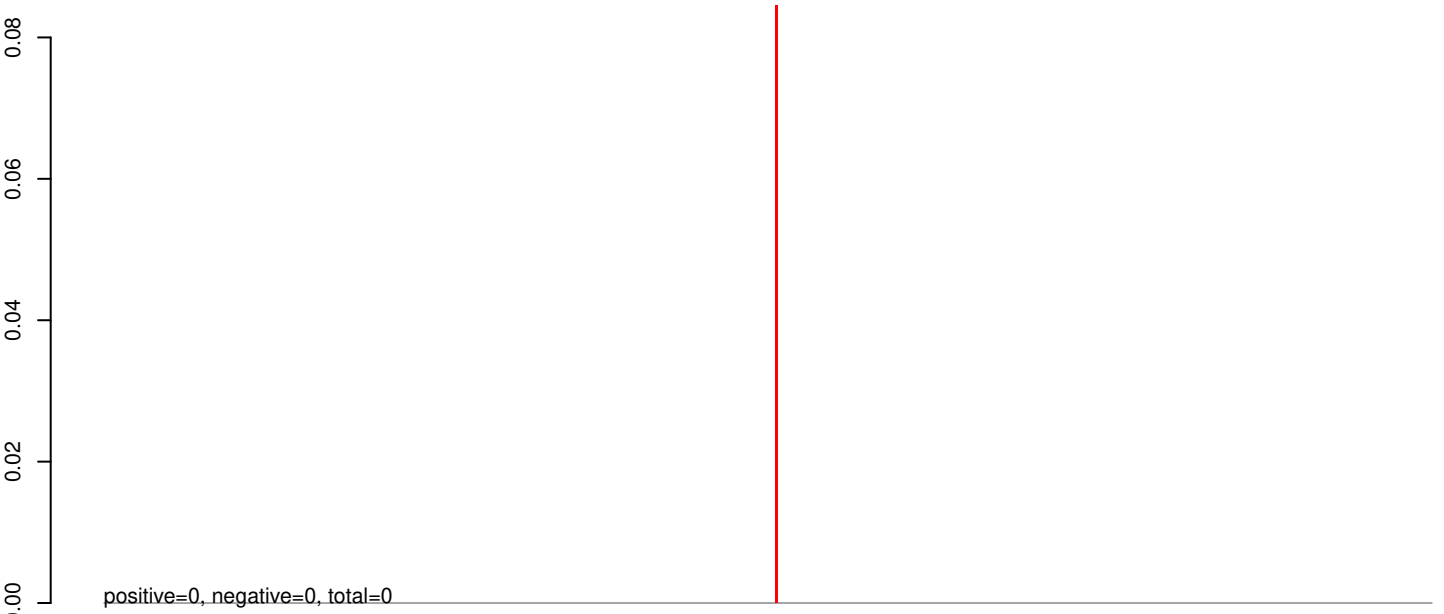
AeAeg_Aag2_BTV1_AK.24_35.rep



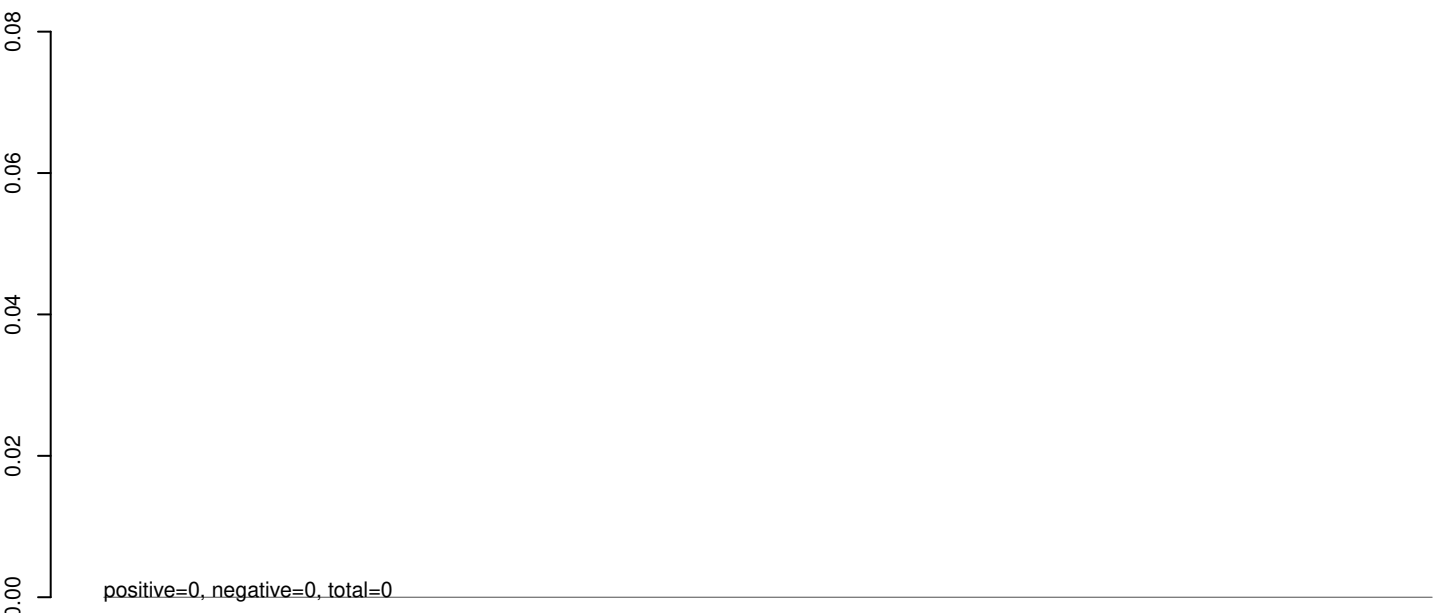
AeAeg_Aag2_BTV1_AK.rep



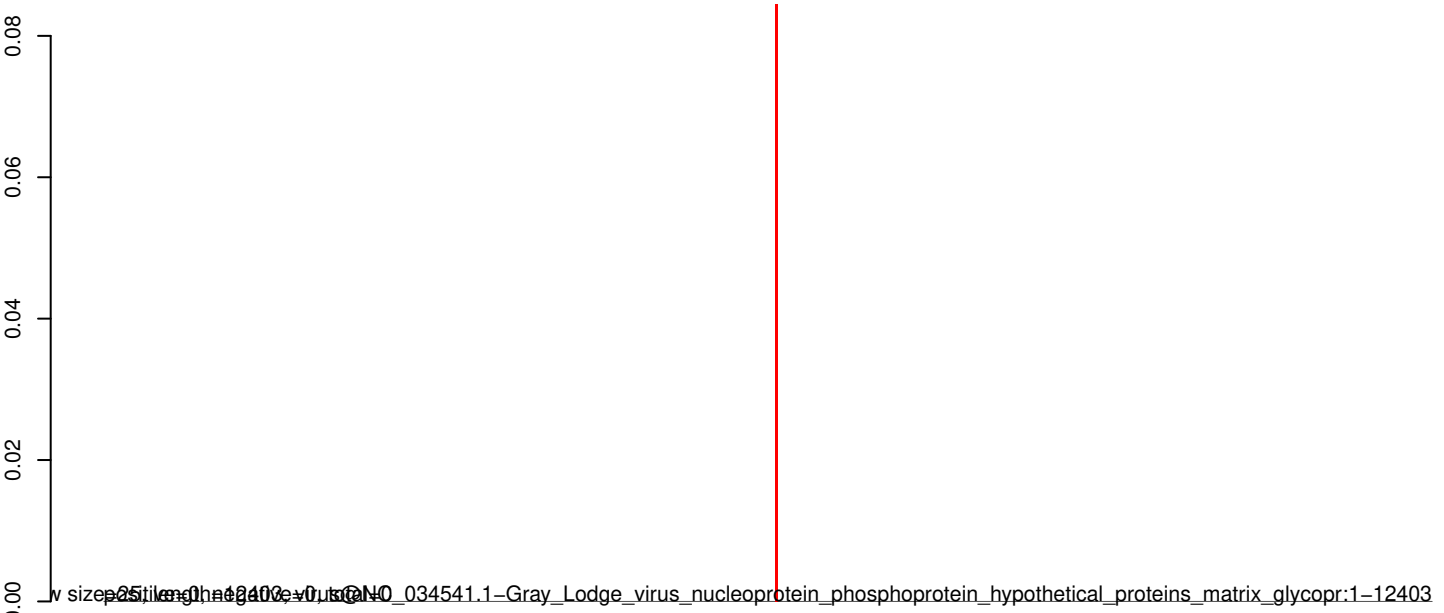
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep

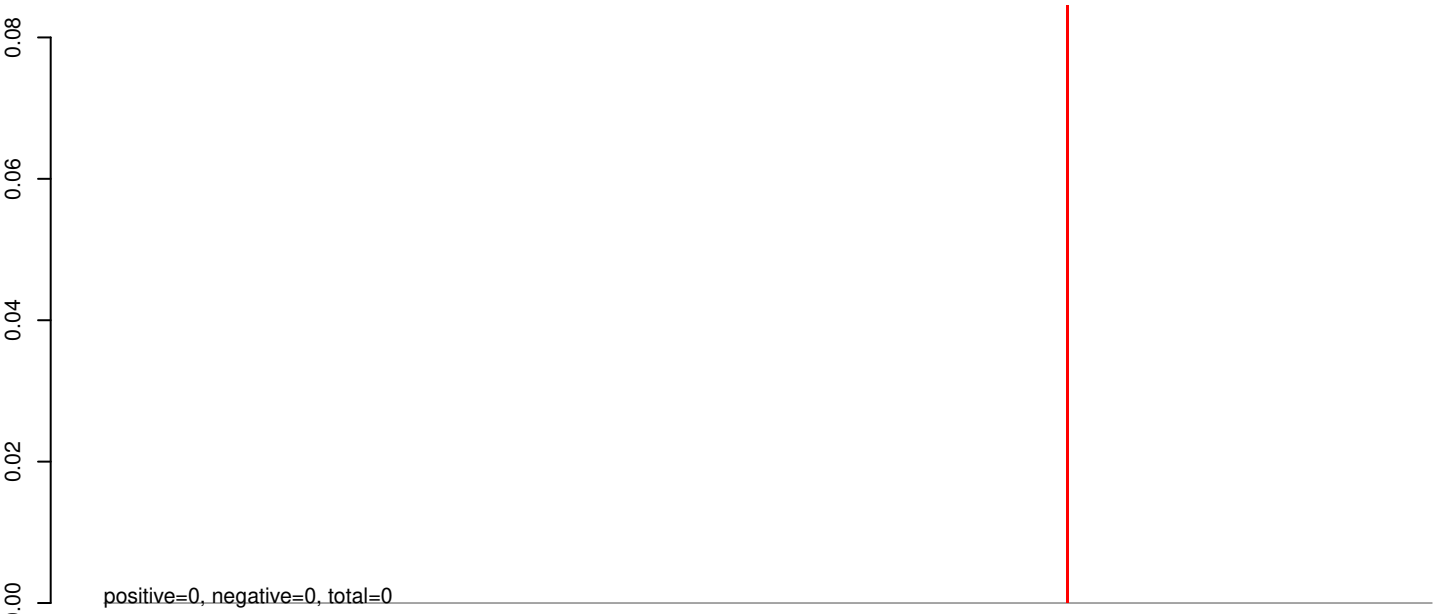


AeAeg_Aag2_BTV1_AK.rep

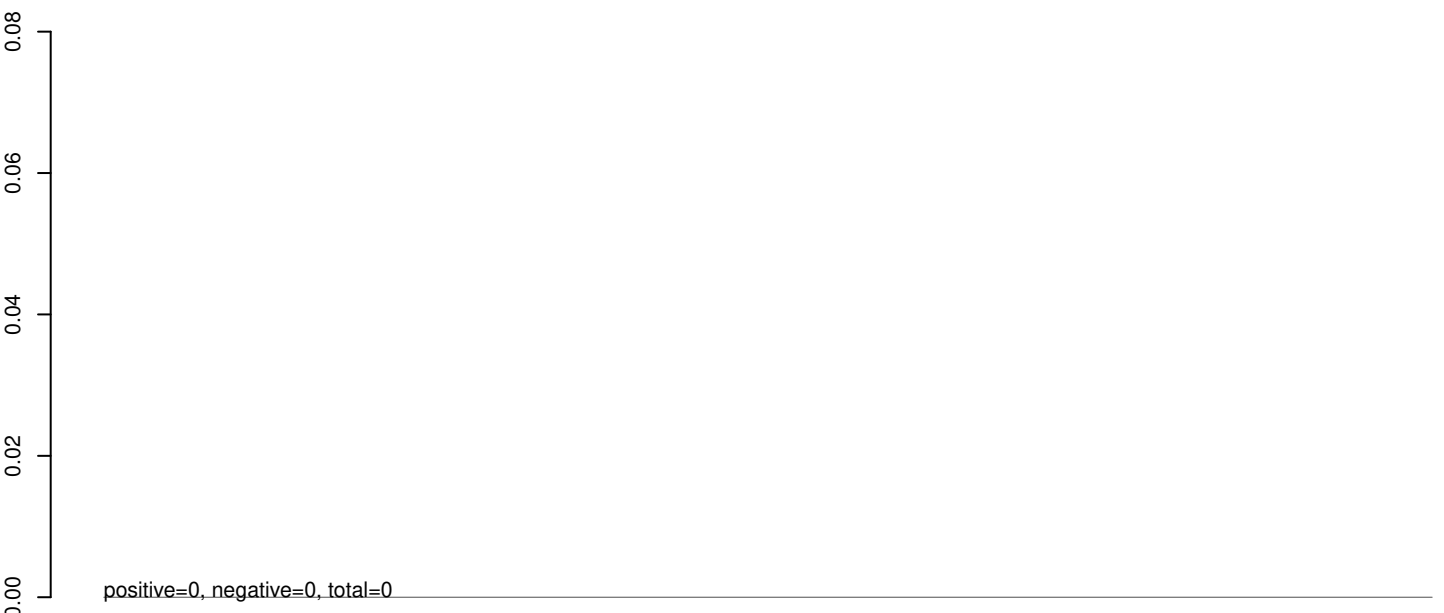


0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_BTV1_AK.18_23.rep



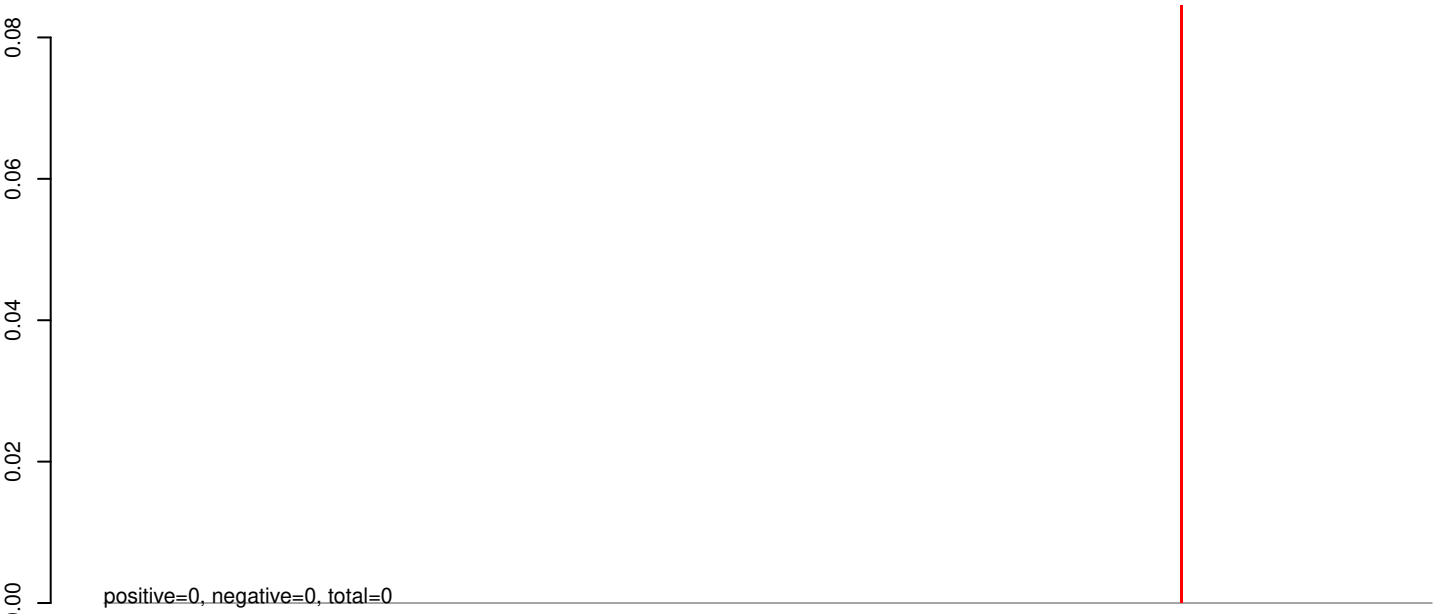
AeAeg_Aag2_BTV1_AK.24_35.rep



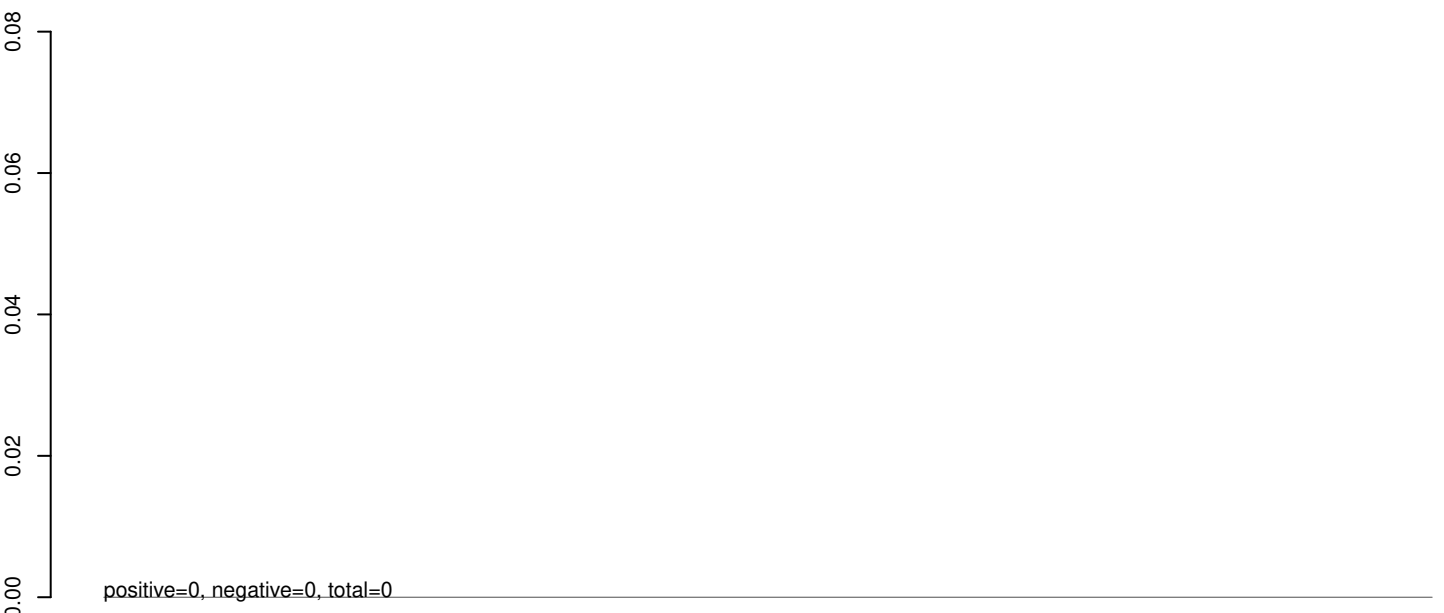
AeAeg_Aag2_BTV1_AK.rep



AeAeg_Aag2_BTV1_AK.18_23.rep



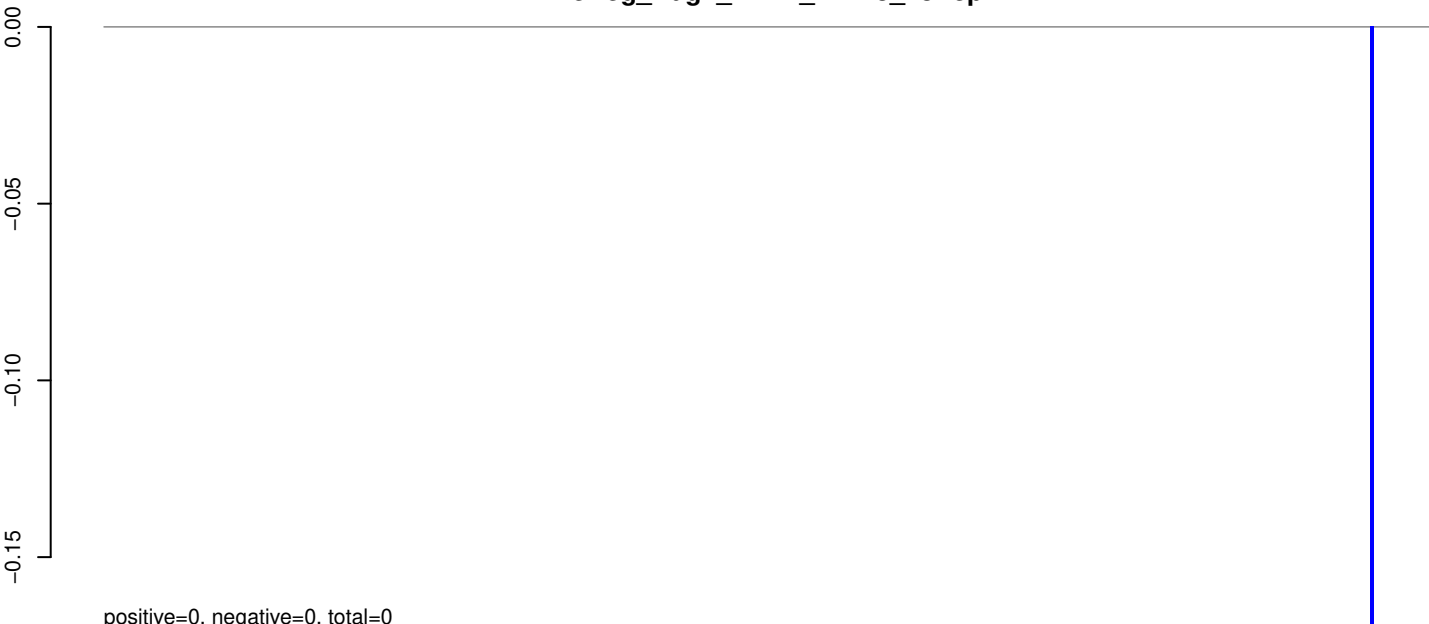
AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep

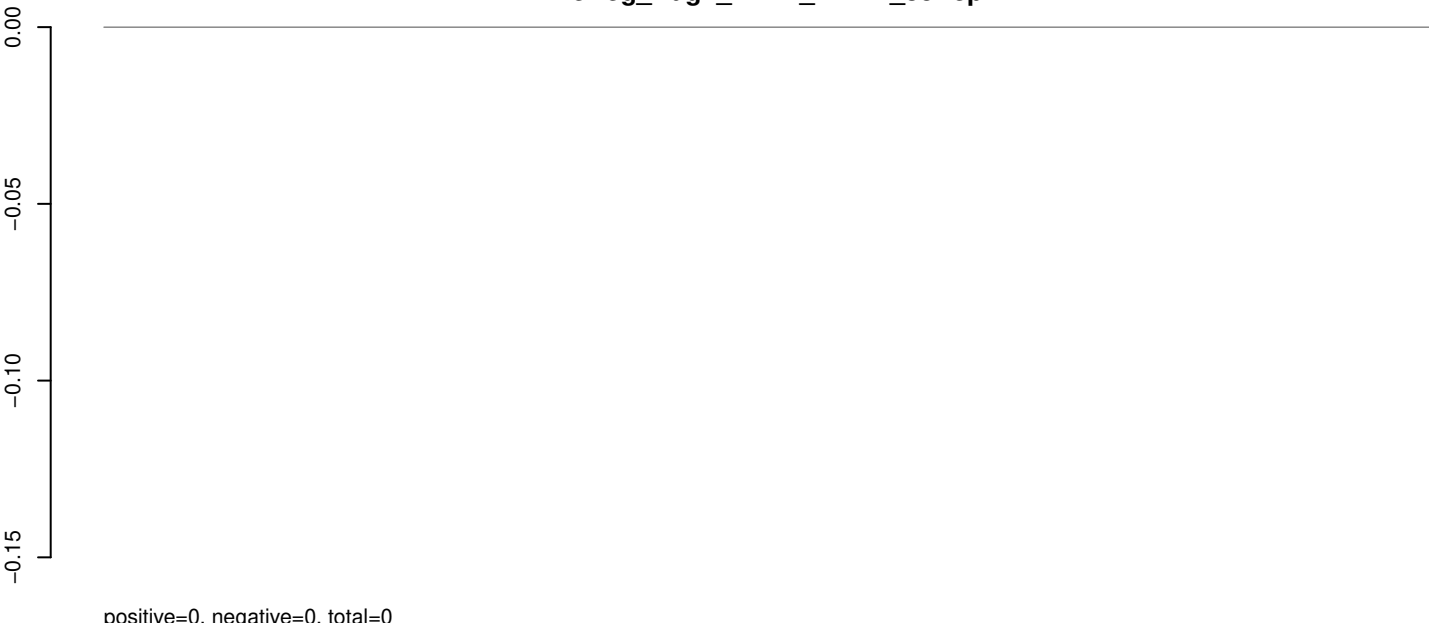


AeAeg_Aag2_BTV1_AK.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BTV1_AK.24_35.rep



positive=0, negative=0, total=0

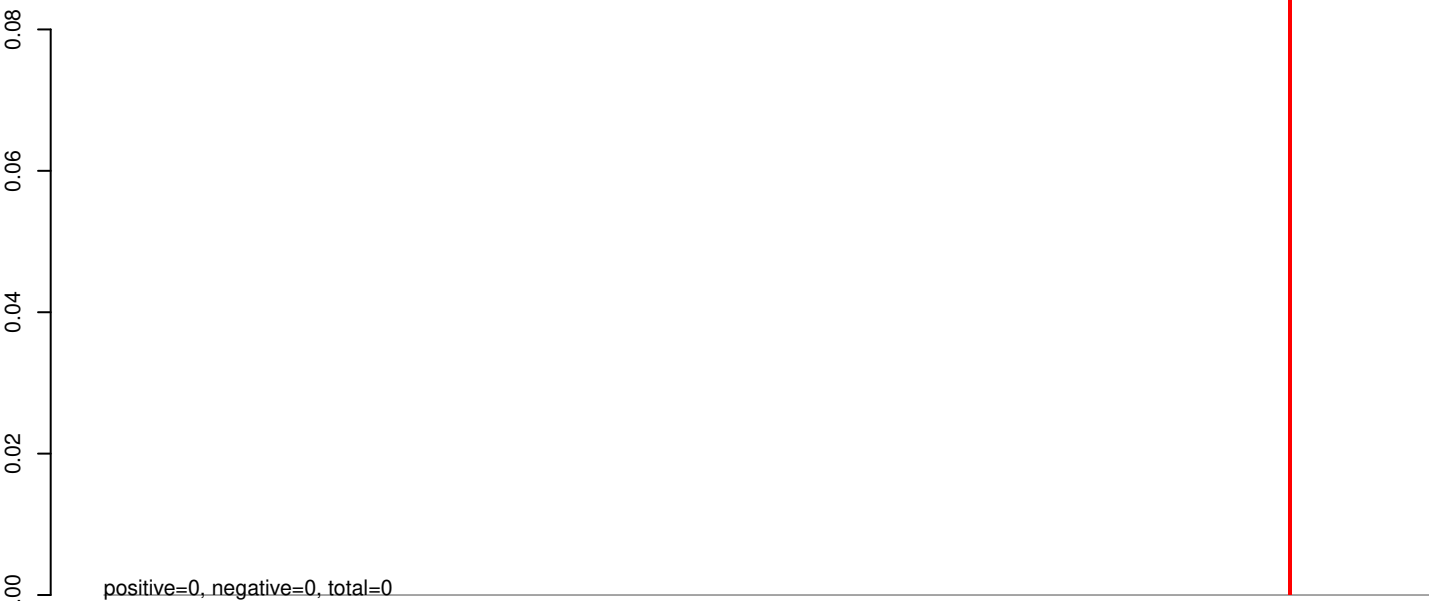
AeAeg_Aag2_BTV1_AK.rep



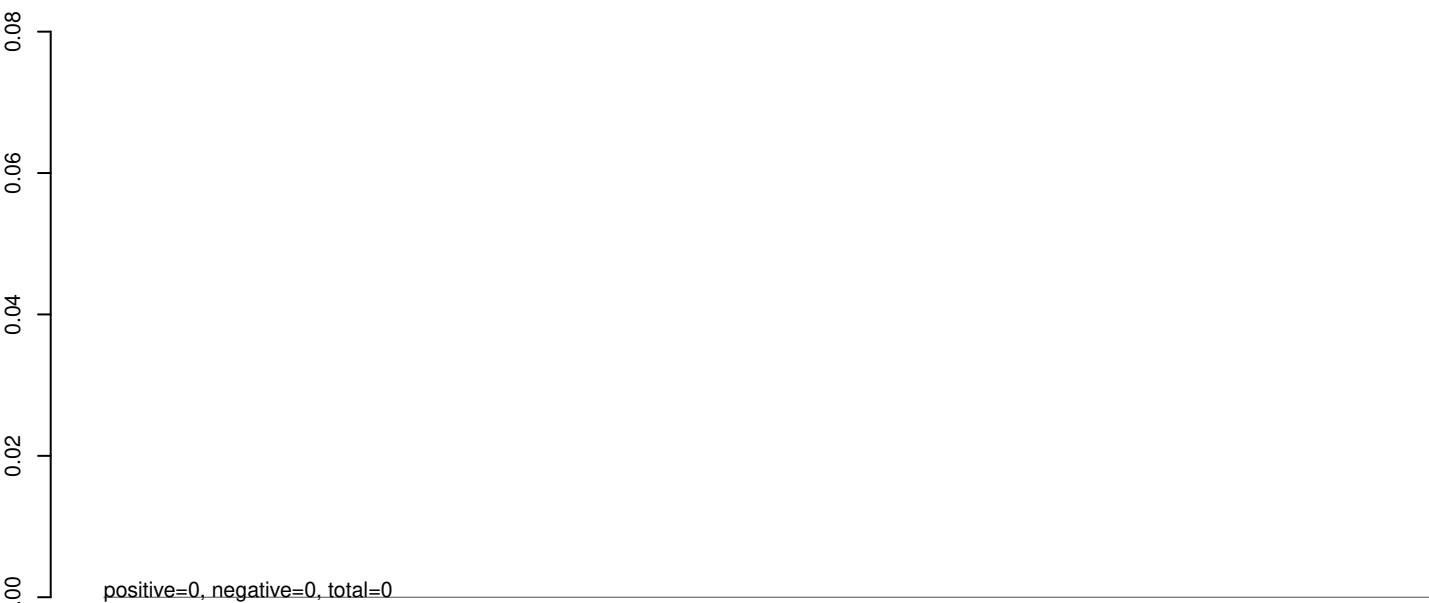
positive=0, negative=0, total=6, window-size=25, length=10251, virus@NC_033723.1-Gadgets_Gully_virus_from_Australia_polyprotein_gene:1-10251

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV1_AK.18_23.rep



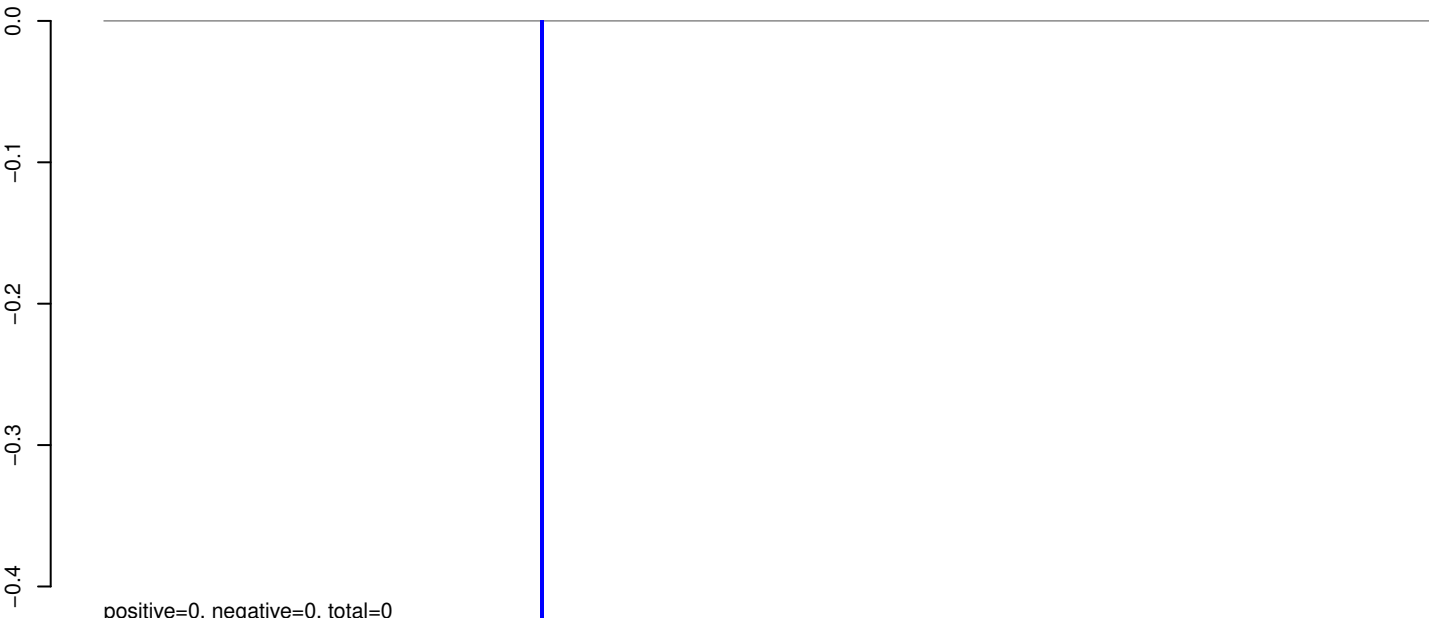
AeAeg_Aag2_BTV1_AK.24_35.rep



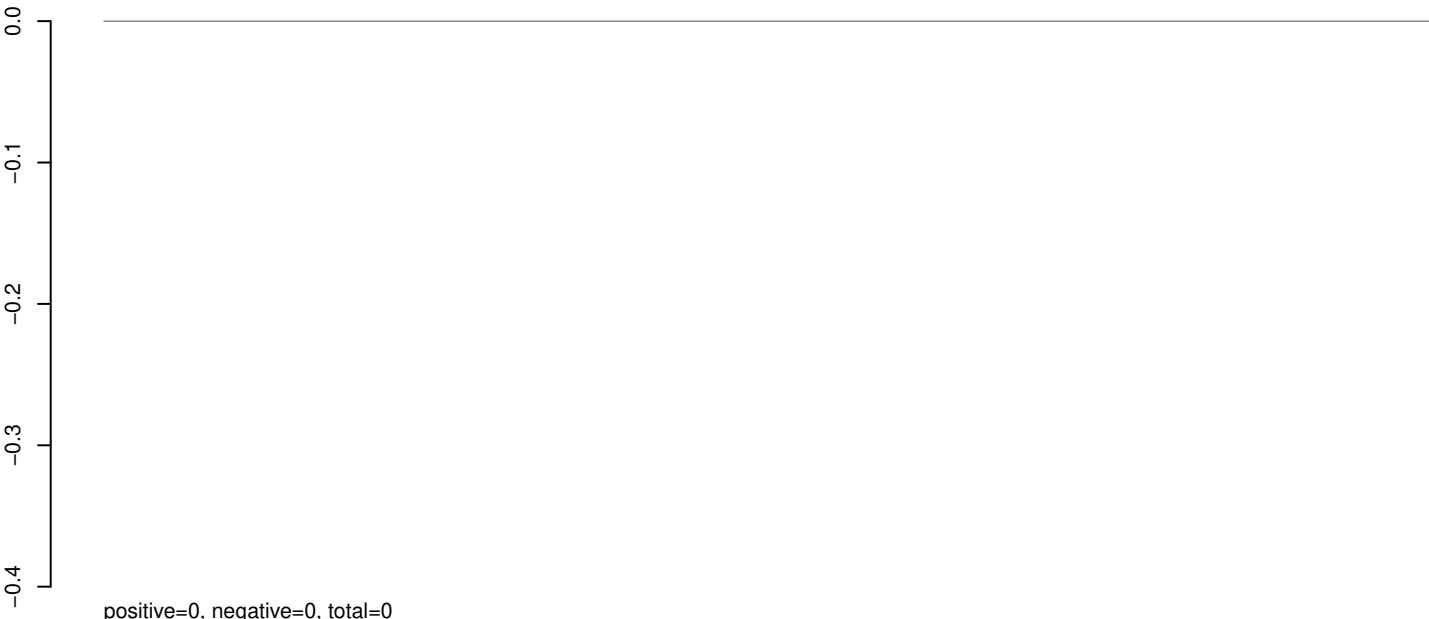
AeAeg_Aag2_BTV1_AK.rep



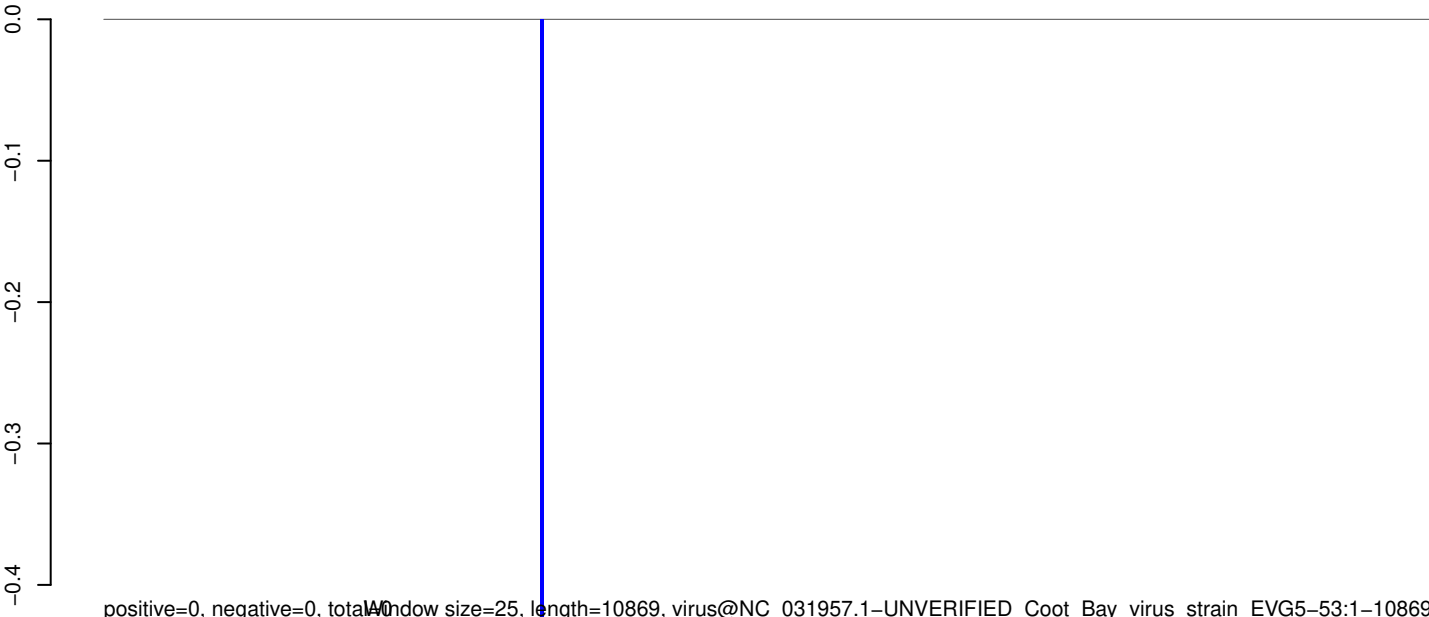
AeAeg_Aag2_BTV1_AK.18_23.rep



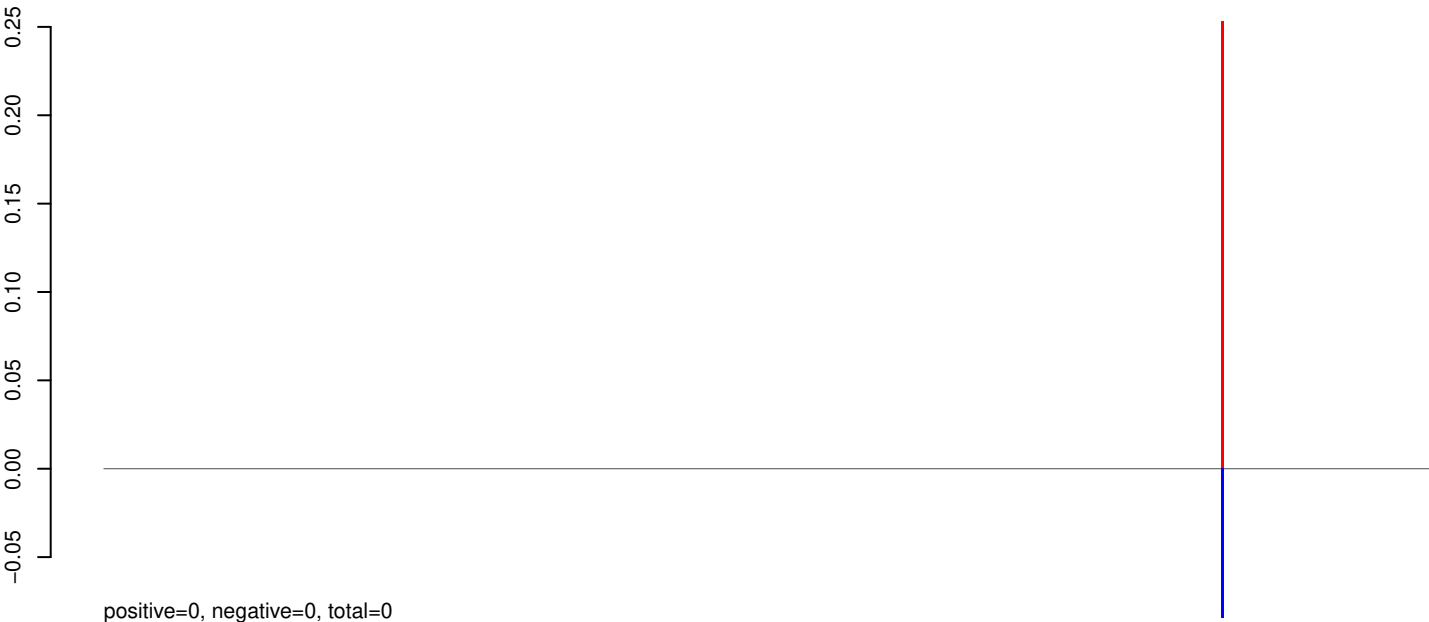
AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep

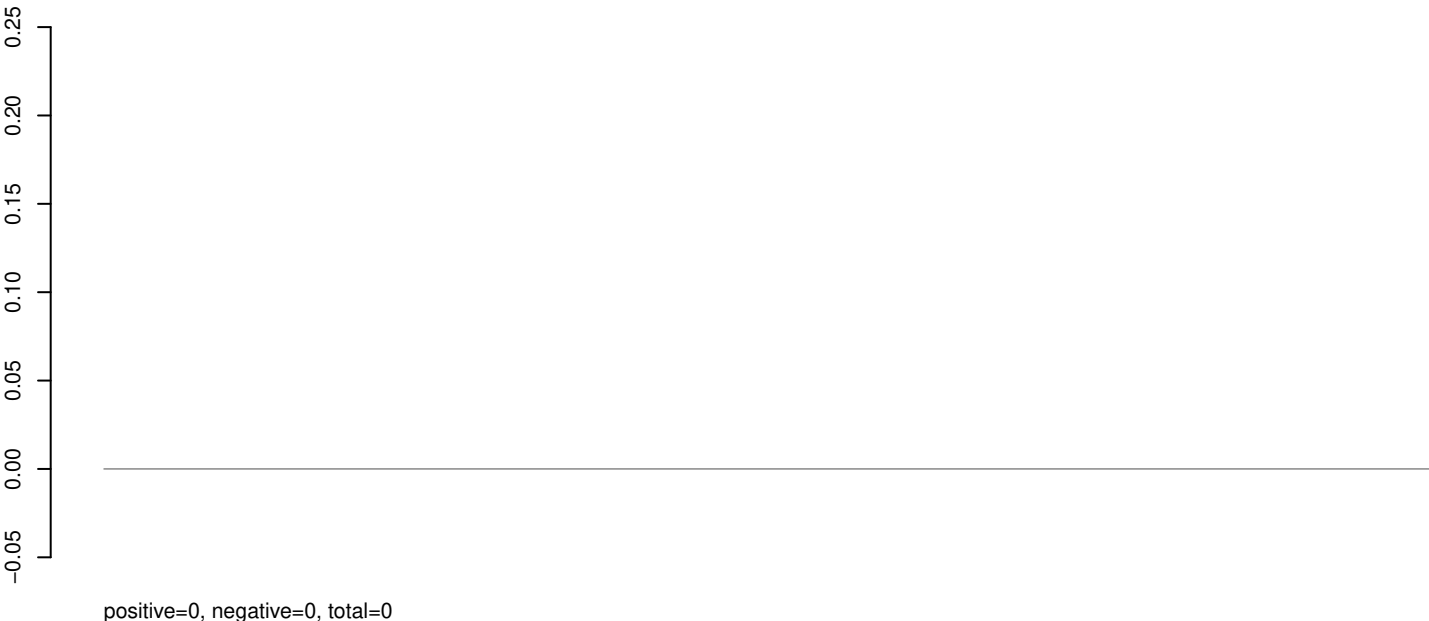


AeAeg_Aag2_BTV1_AK.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BTV1_AK.24_35.rep



positive=0, negative=0, total=0

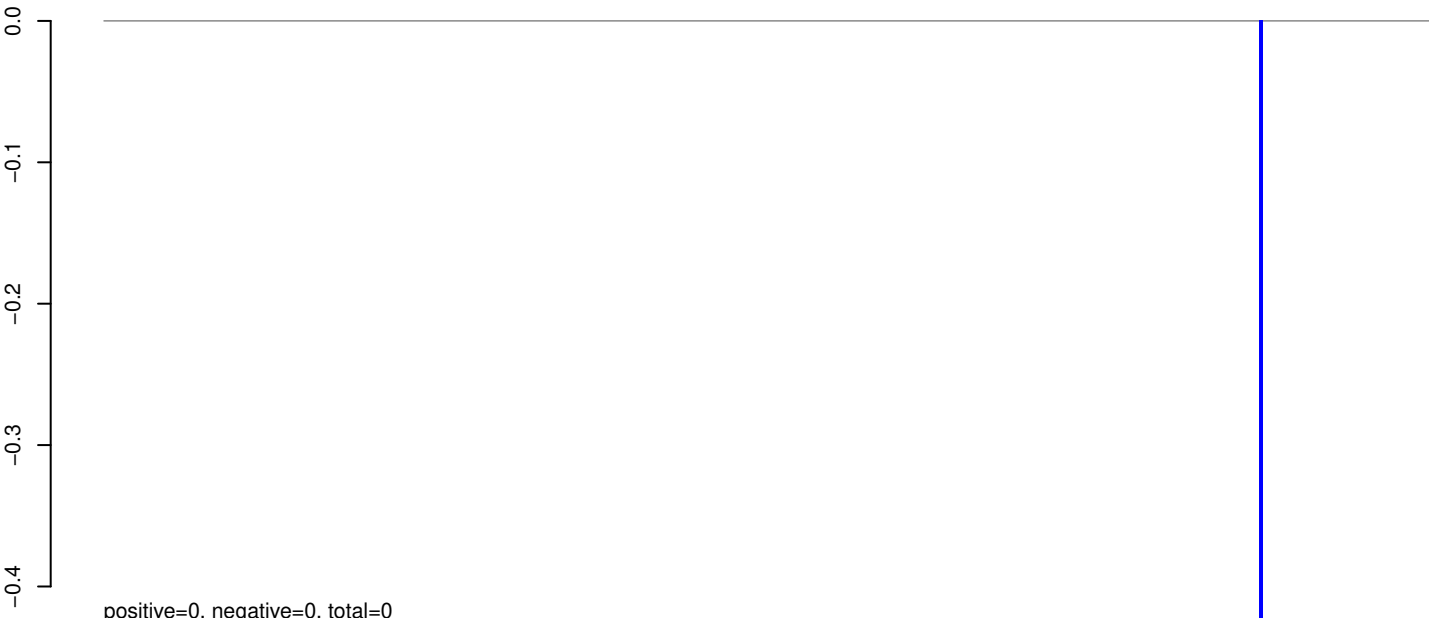
AeAeg_Aag2_BTV1_AK.rep



positive=0, negative=0, total=0 Window size=25, length=11286, virus@NC_031956.1-UNVERIFIED_Balsa_virus_strain_CoB_84:1-11286

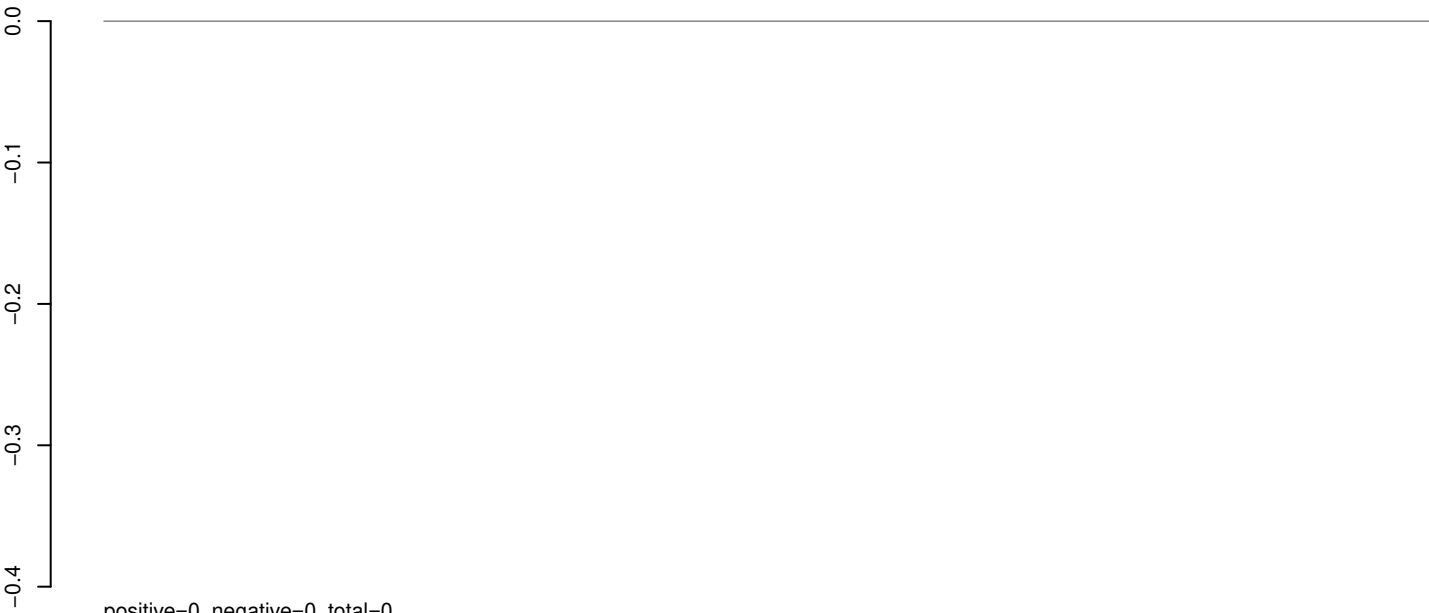
0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV1_AK.18_23.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BTV1_AK.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_BTV1_AK.rep

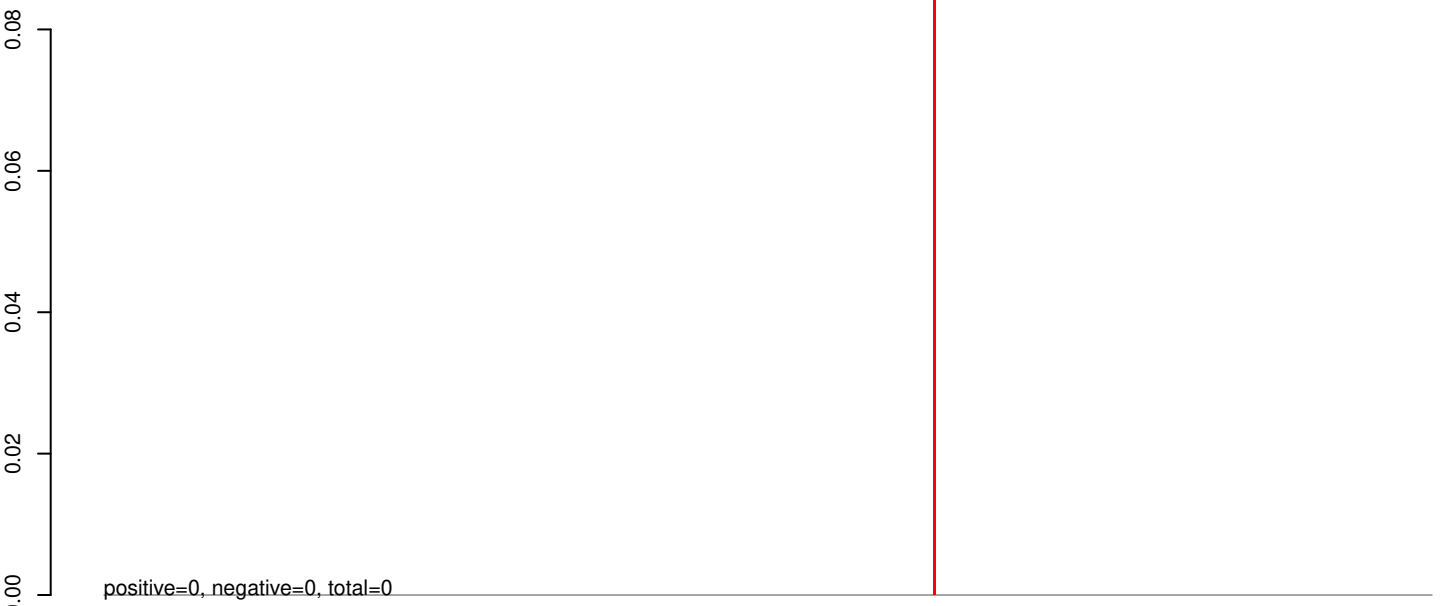


positive=0, negative=0, total=0

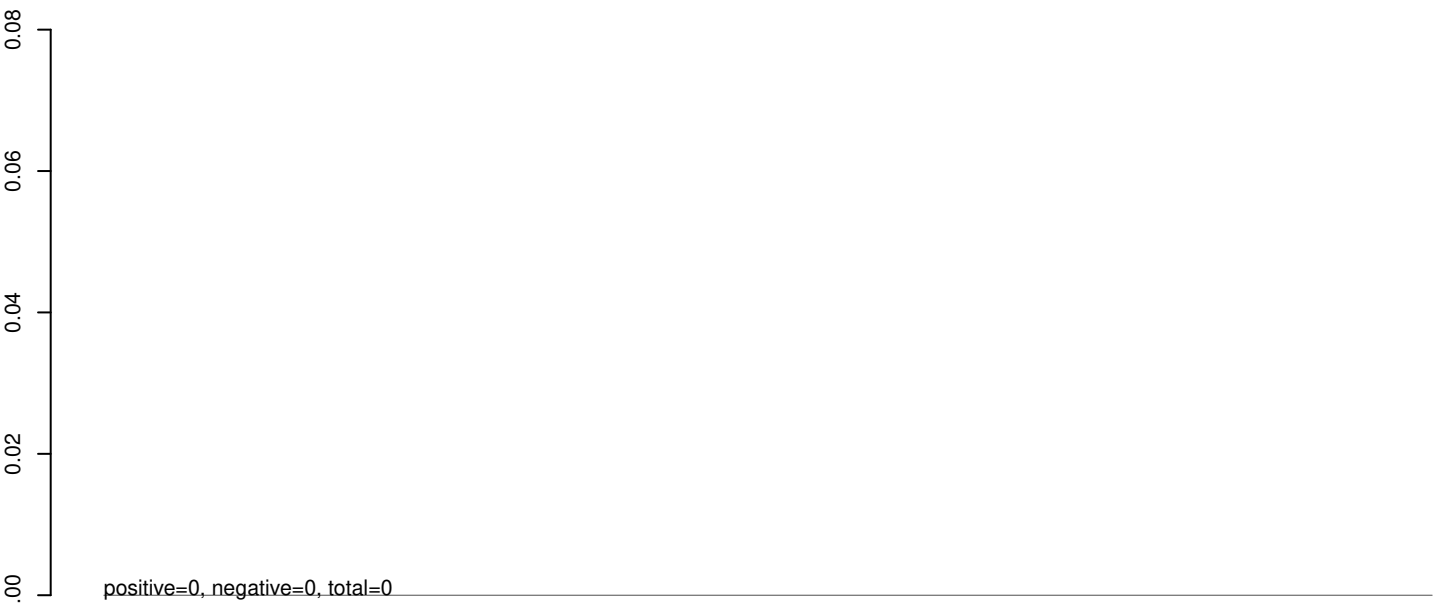
Window size=25, length=10588, virus@NC_031327.1-Anopheles_flavivirus_variant_1:1-10588

0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep

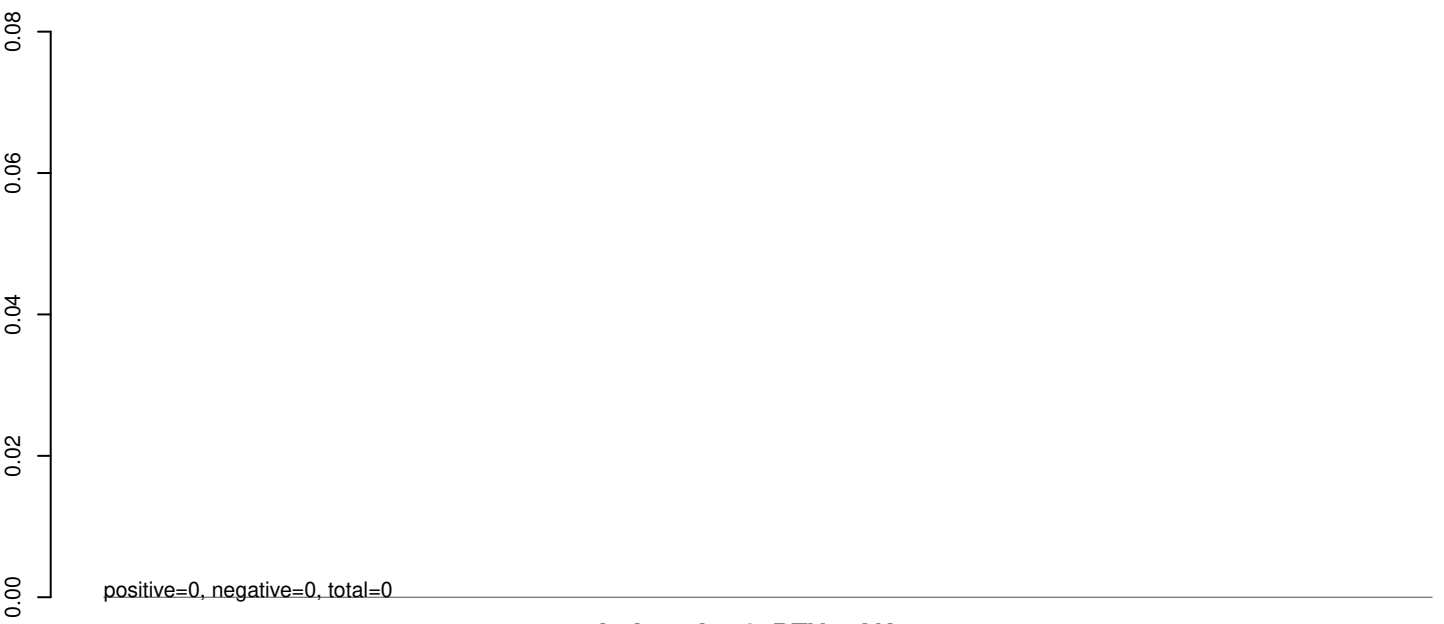


Xincheng_Mosquito_Virus_strain_XC1-6_ORF1_ORF1_ORF2_ORF2_glycoprotein_G_and_RNA-de:1-12774

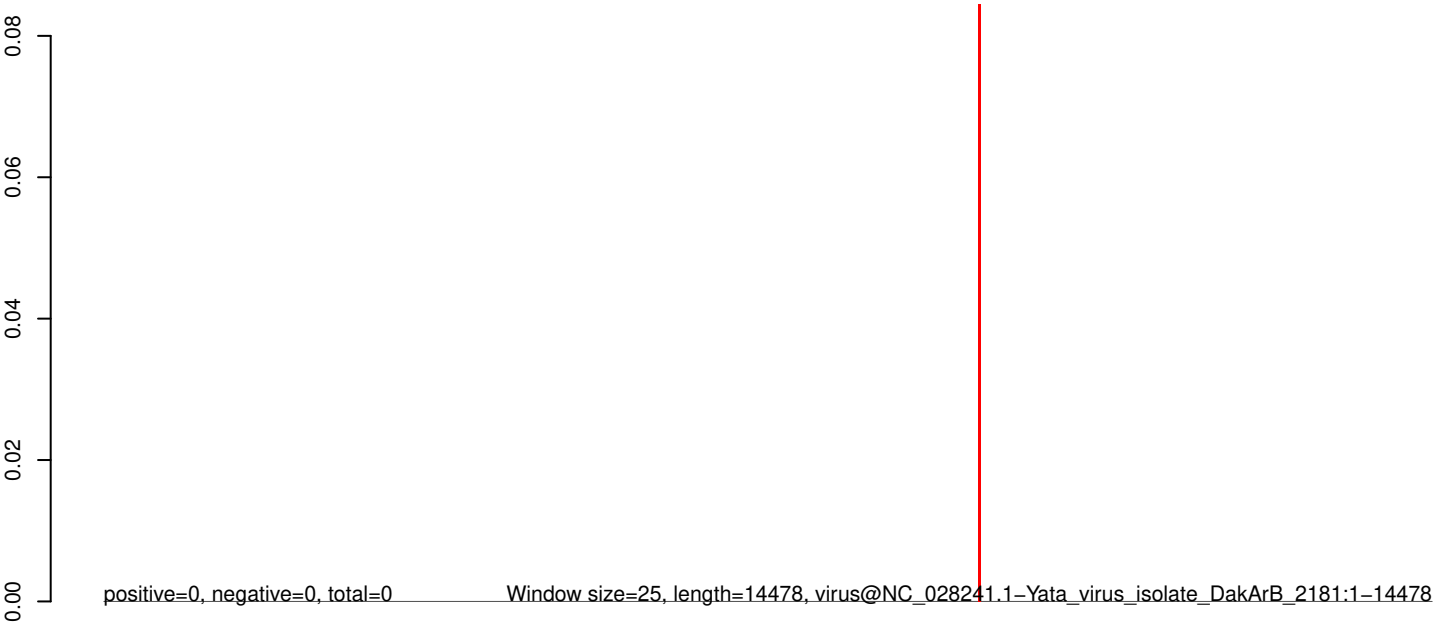
AeAeg_Aag2_BTV1_AK.18_23.rep



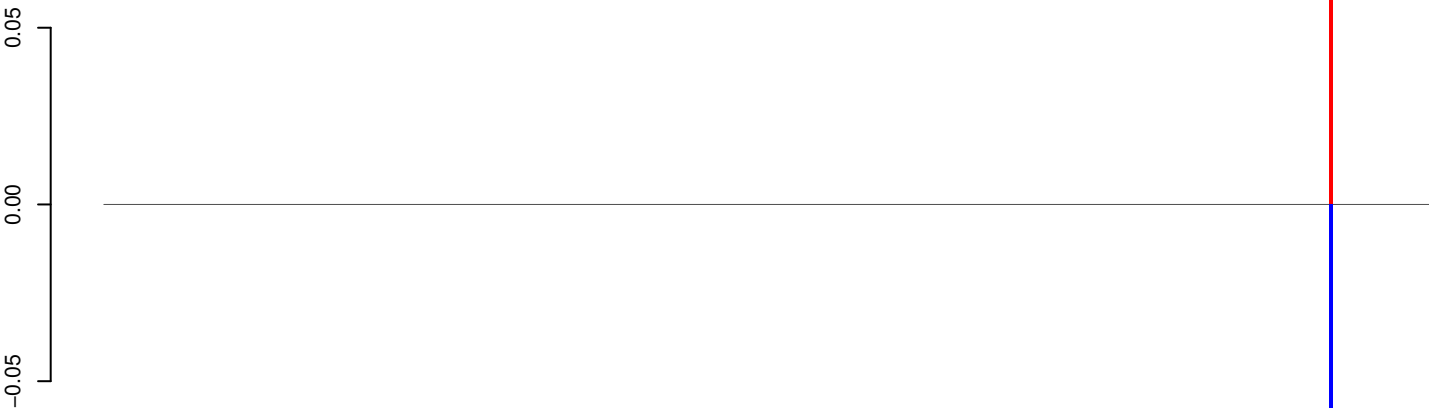
AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep

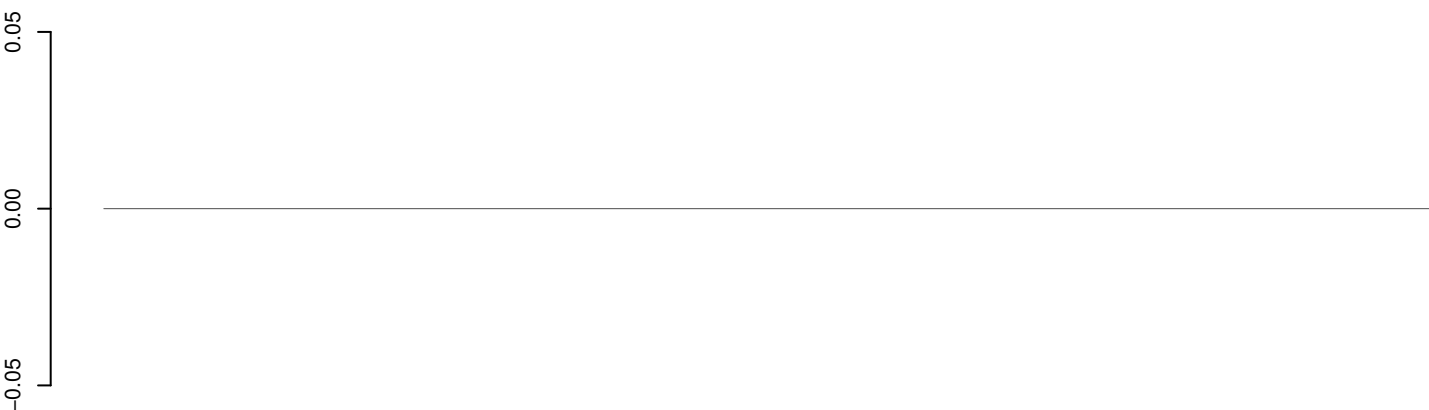


AeAeg_Aag2_BTV1_AK.18_23.rep



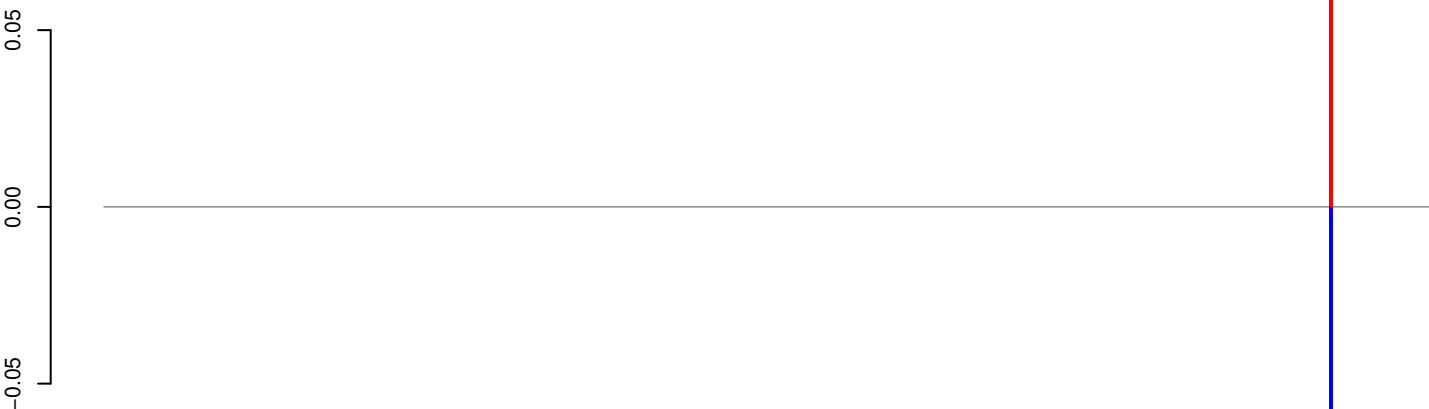
positive=0, negative=0, total=0

AeAeg_Aag2_BTV1_AK.24_35.rep



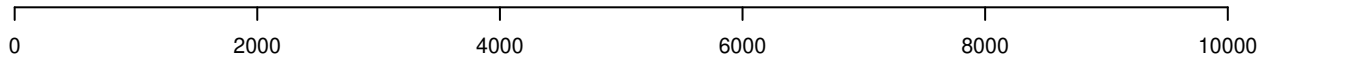
positive=0, negative=0, total=0

AeAeg_Aag2_BTV1_AK.rep

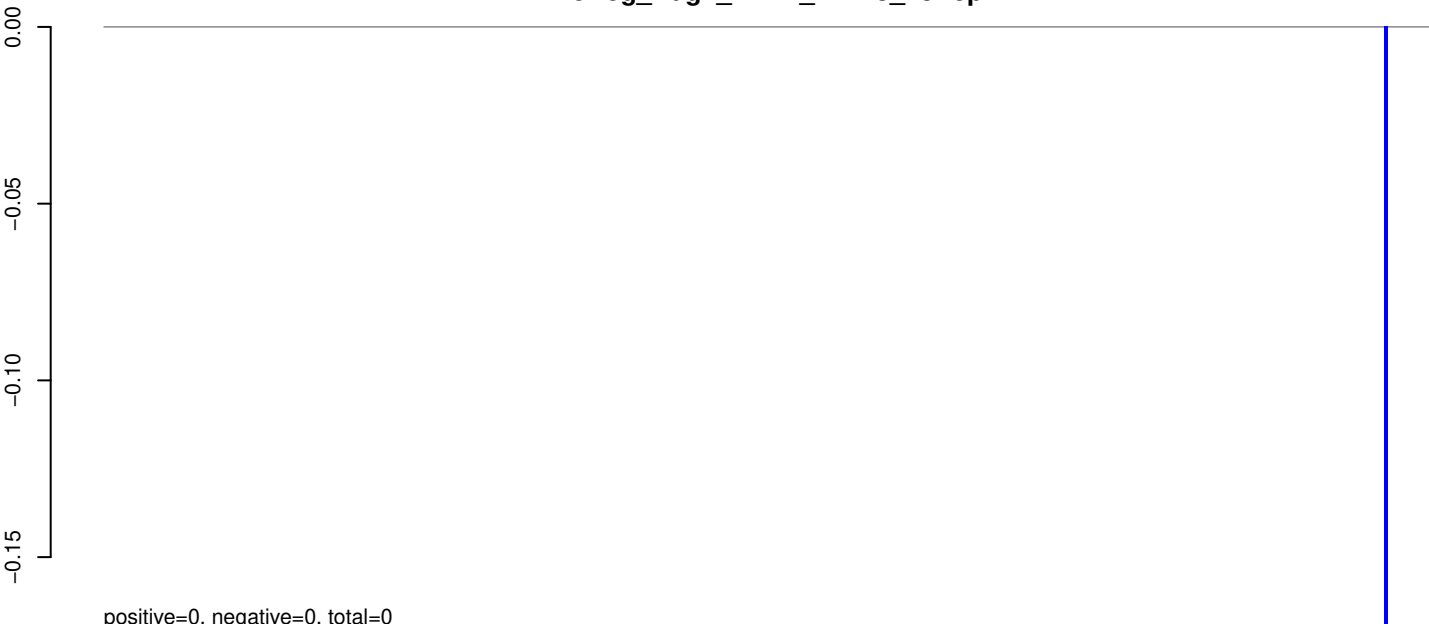


positive=0, negative=0, total=0

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



AeAeg_Aag2_BTV1_AK.18_23.rep



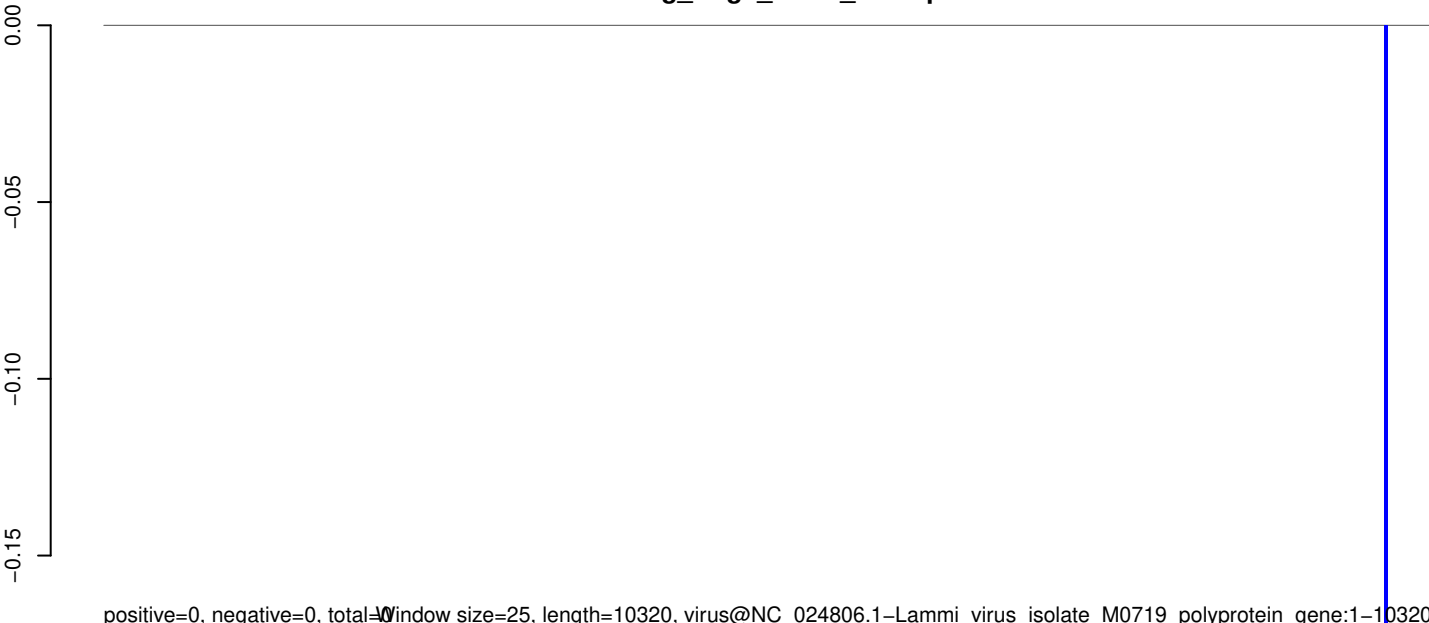
positive=0, negative=0, total=0

AeAeg_Aag2_BTV1_AK.24_35.rep



positive=0, negative=0, total=0

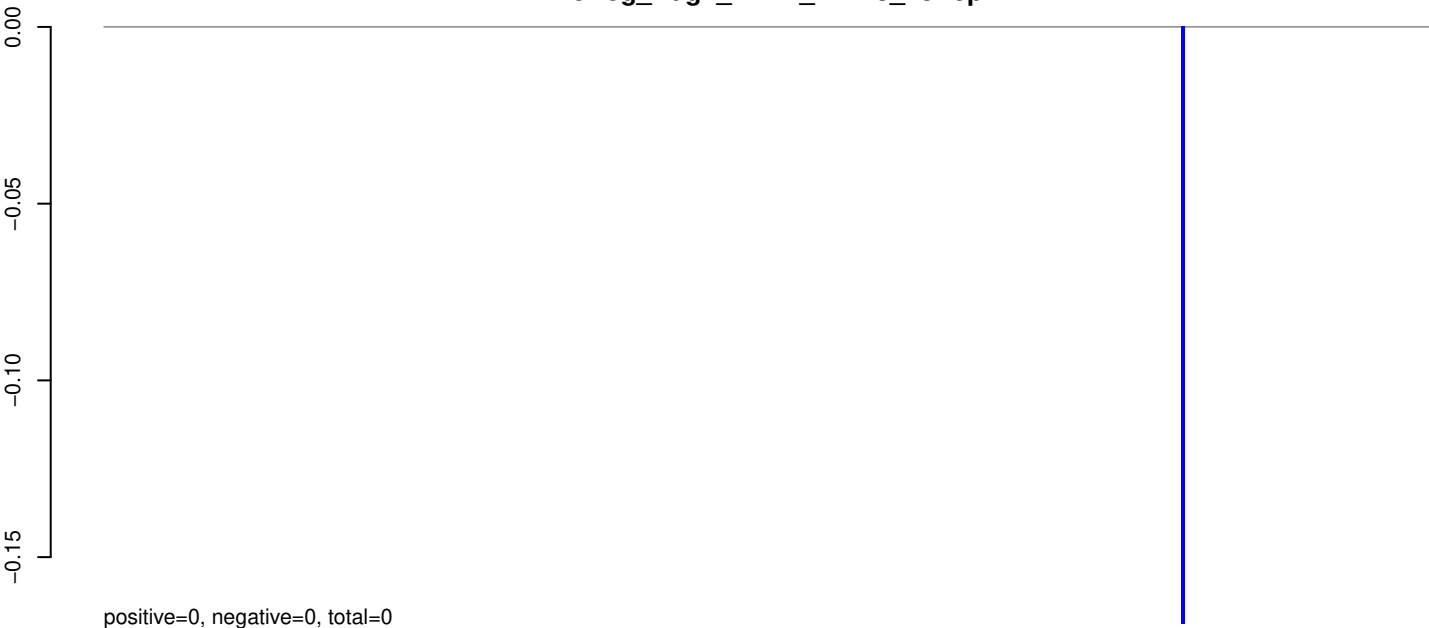
AeAeg_Aag2_BTV1_AK.rep



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

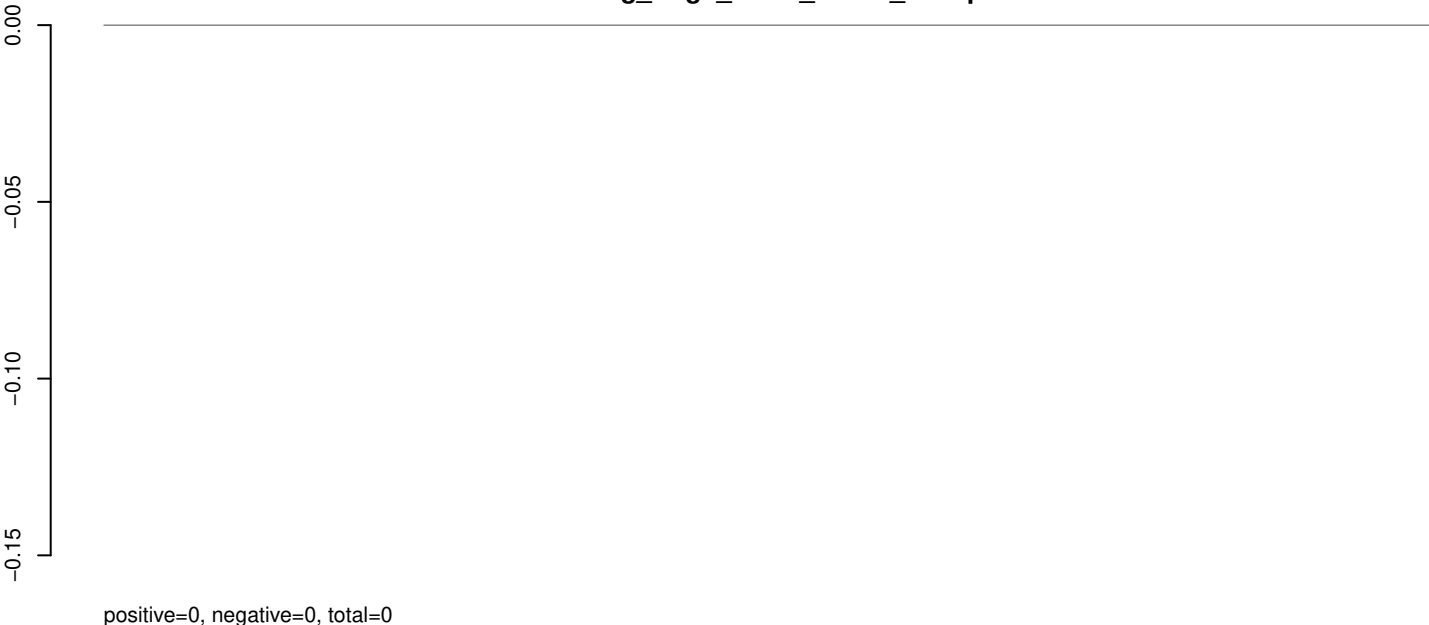
0 2000 4000 6000 8000 10000

AeAeg_Aag2_BTV1_AK.18_23.rep



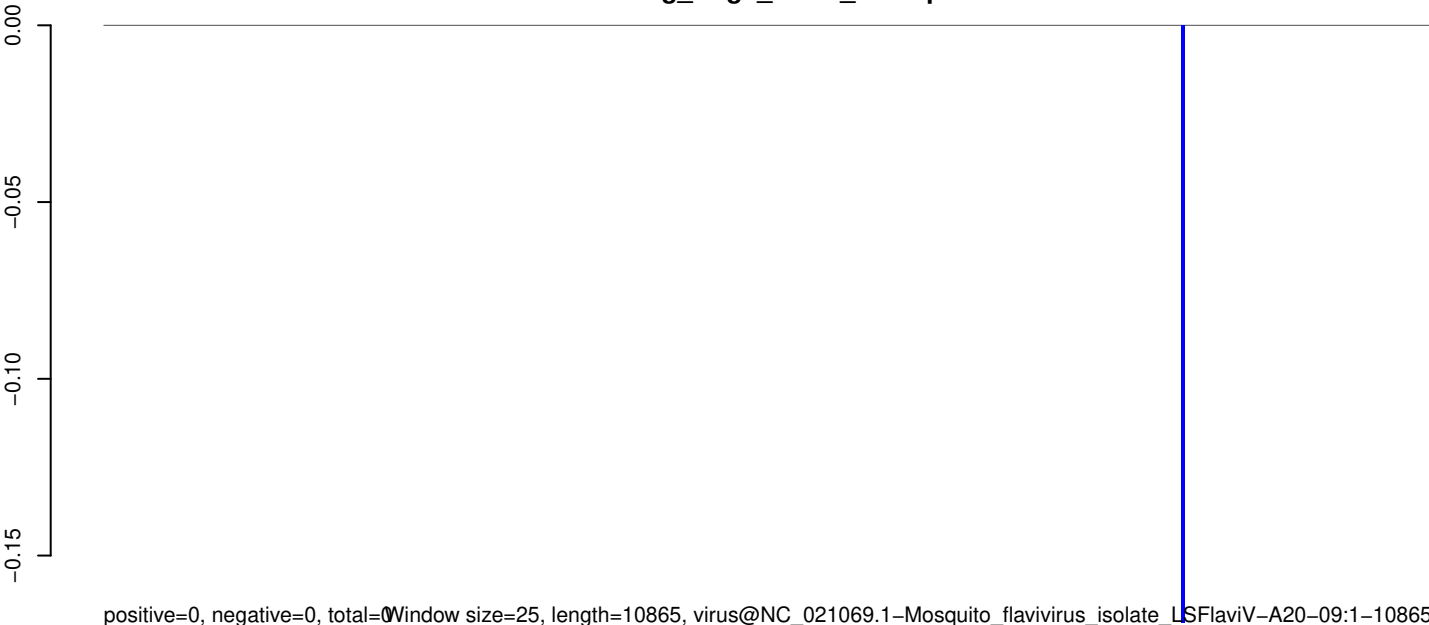
positive=0, negative=0, total=0

AeAeg_Aag2_BTV1_AK.24_35.rep



positive=0, negative=0, total=0

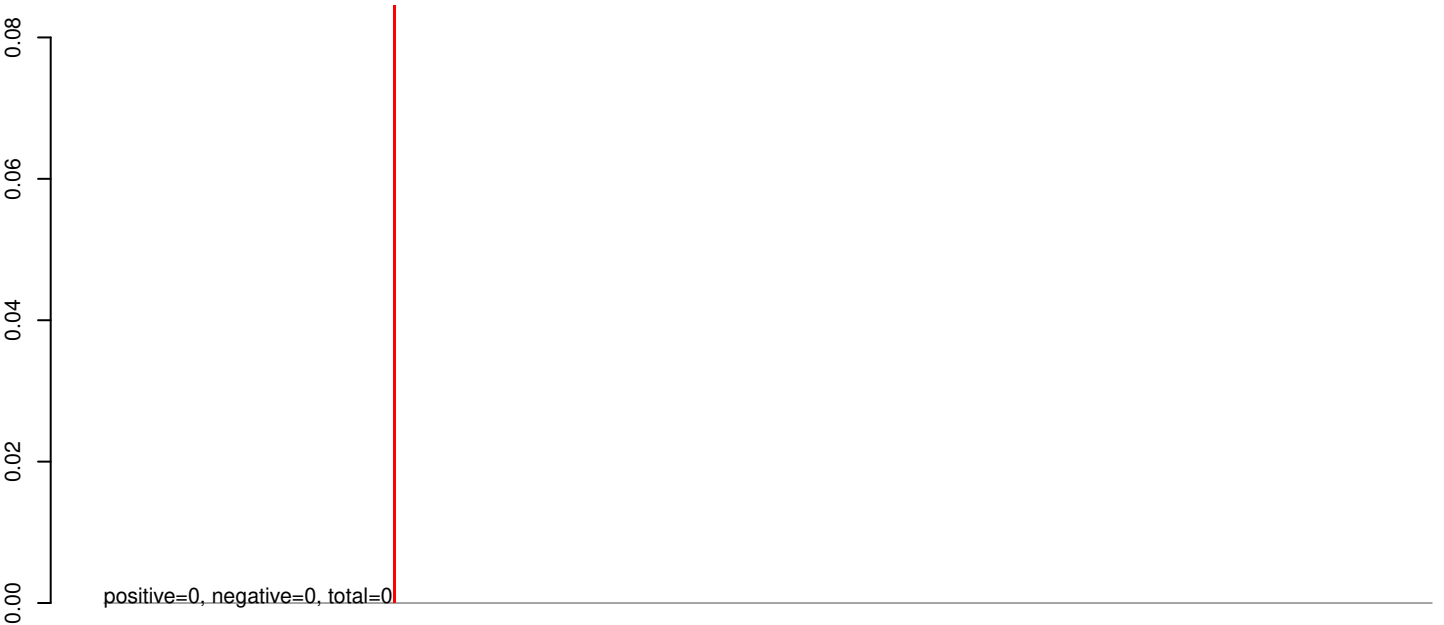
AeAeg_Aag2_BTV1_AK.rep



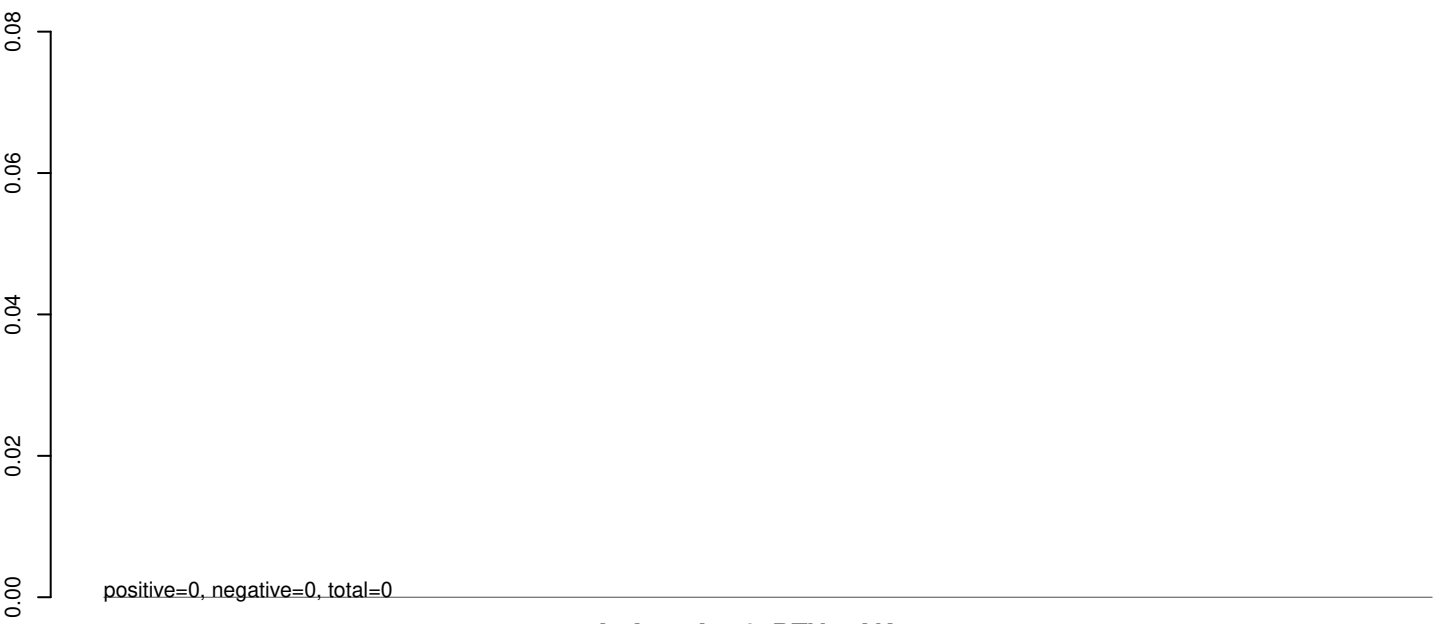
positive=0, negative=0, total=0 Window size=25, length=10865, virus@NC_021069.1-Mosquito_flavivirus_isolate_LSFlaviV-A20-09:1-10865

0 2000 4000 6000 8000 10000

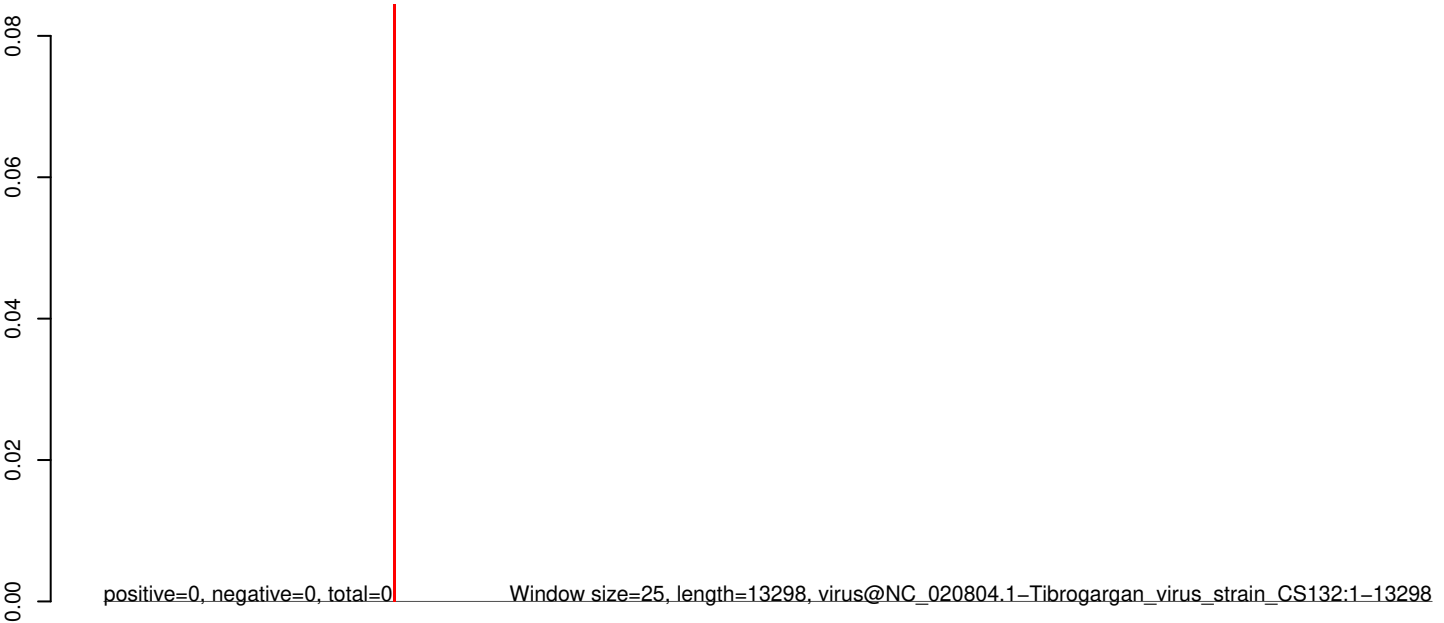
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep

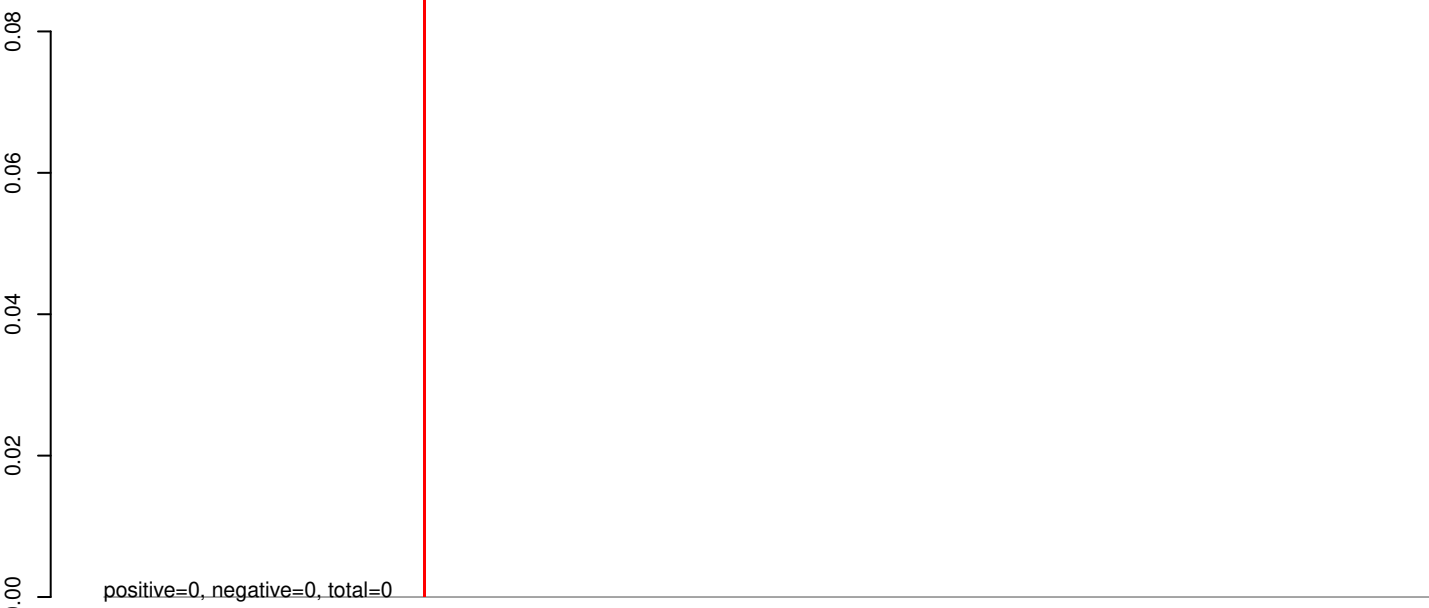


AeAeg_Aag2_BTV1_AK.rep

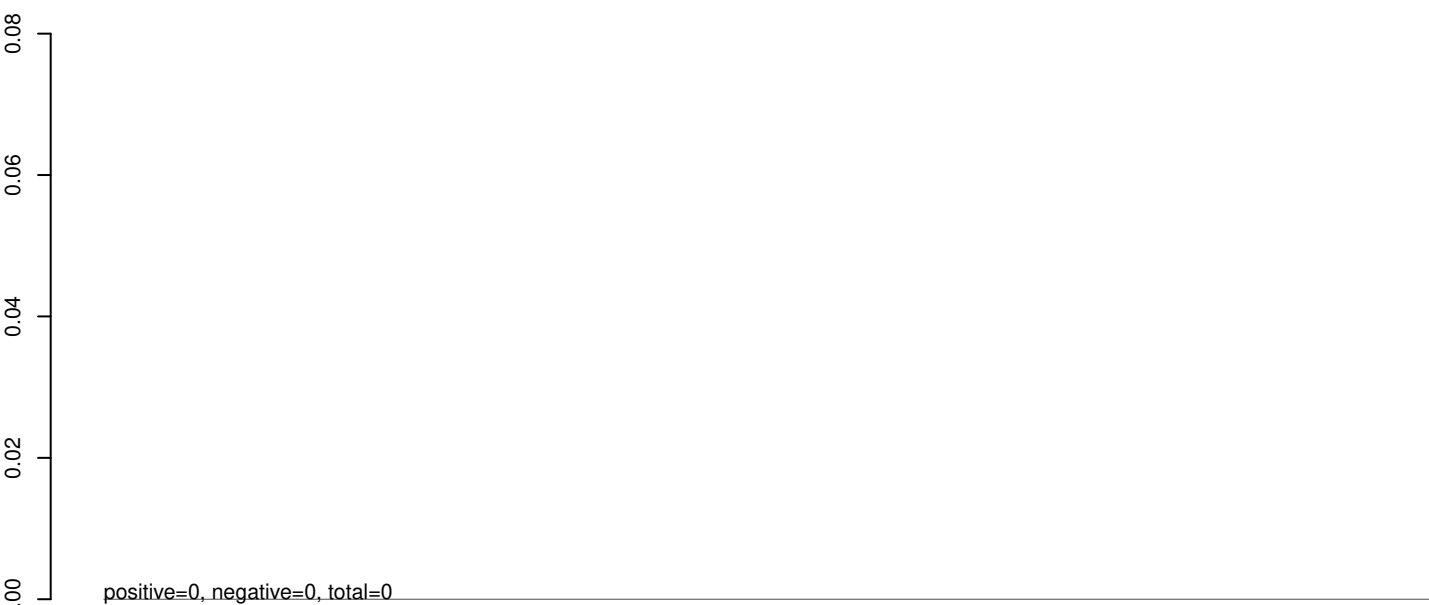


Window size=25, length=13298, virus@NC_020804.1-Tibrogargan_virus_strain_CS132:1-13298

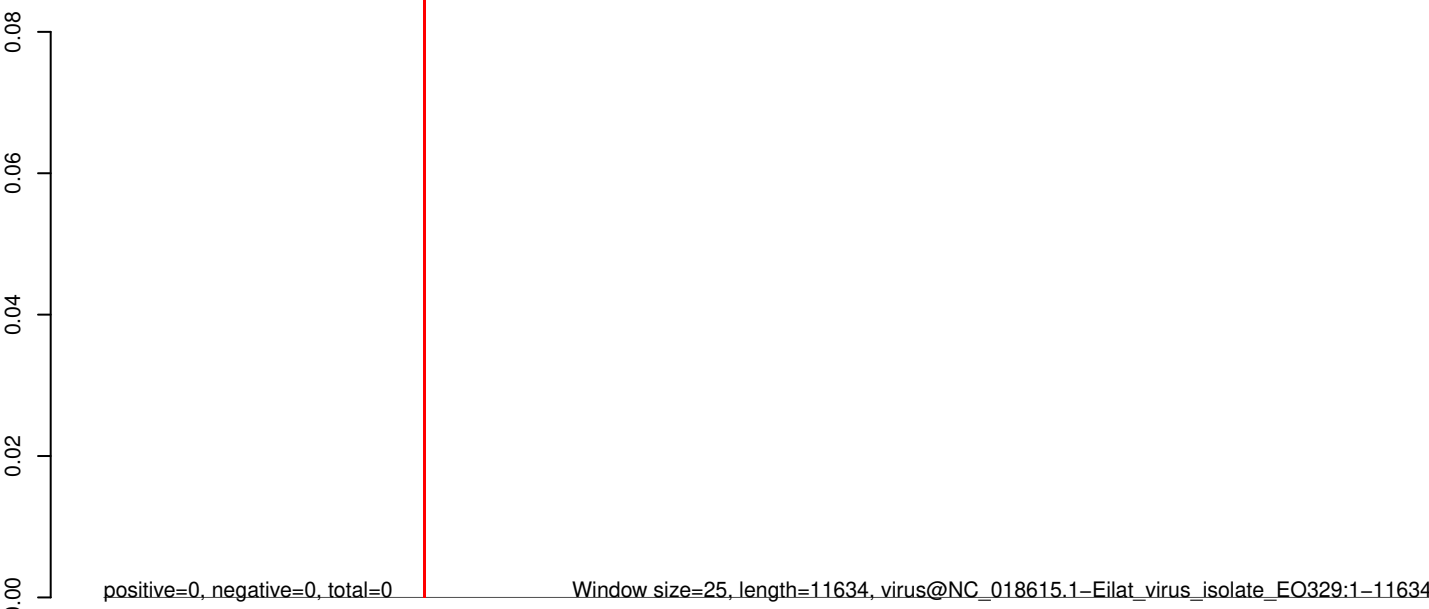
AeAeg_Aag2_BTV1_AK.18_23.rep



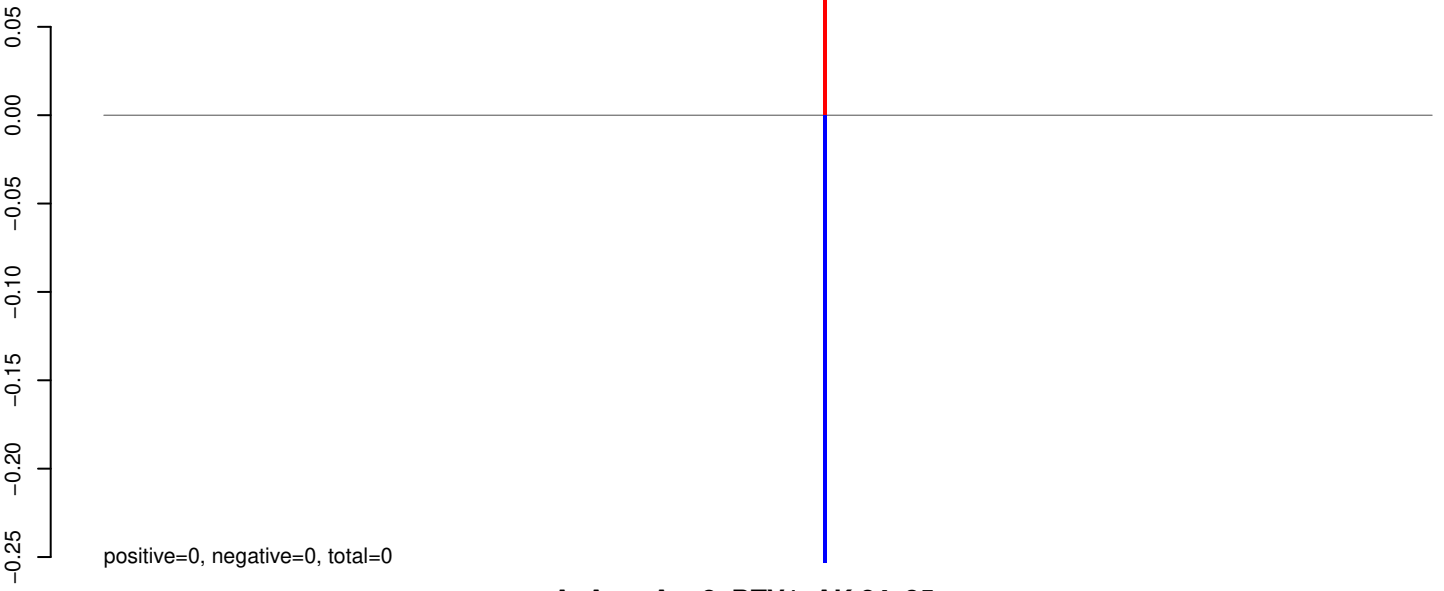
AeAeg_Aag2_BTV1_AK.24_35.rep



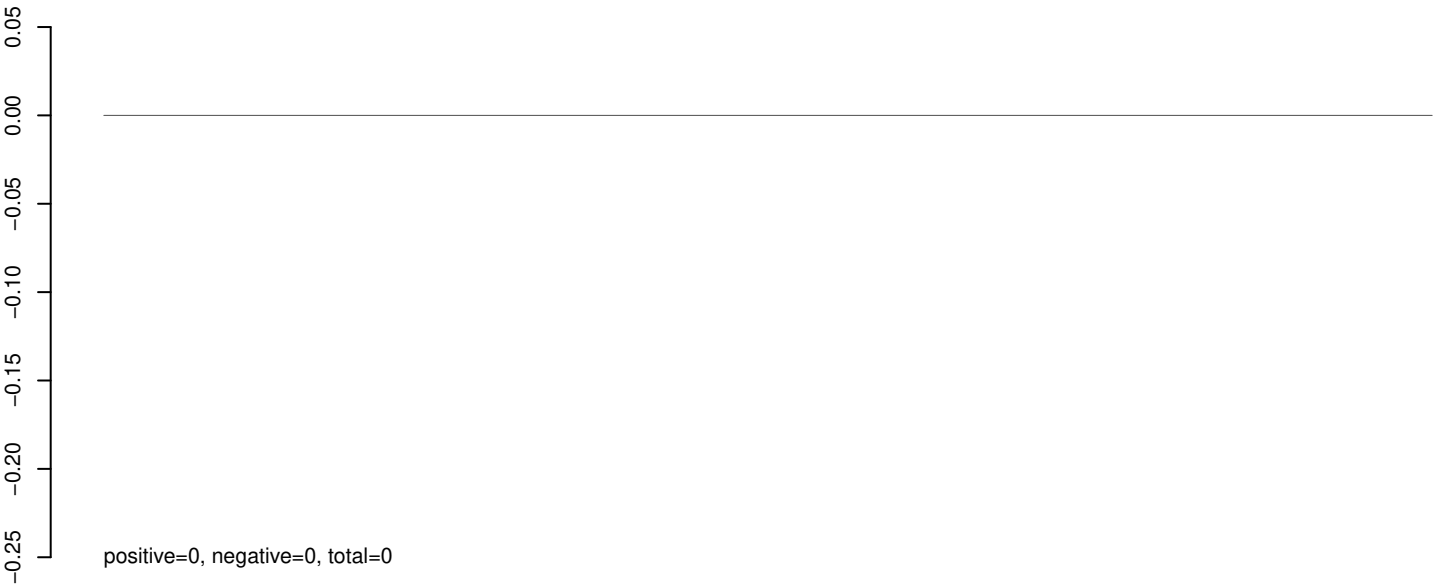
AeAeg_Aag2_BTV1_AK.rep



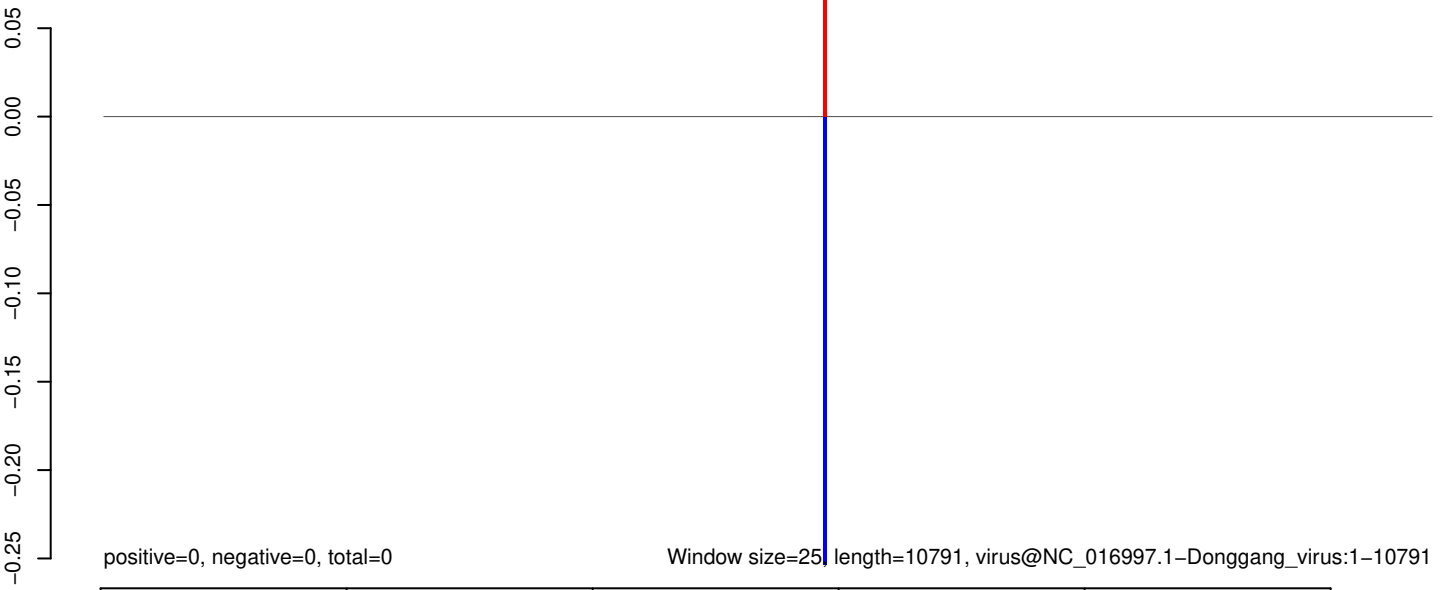
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AeAeg_Aag2_BTV1_AK.24_35.rep

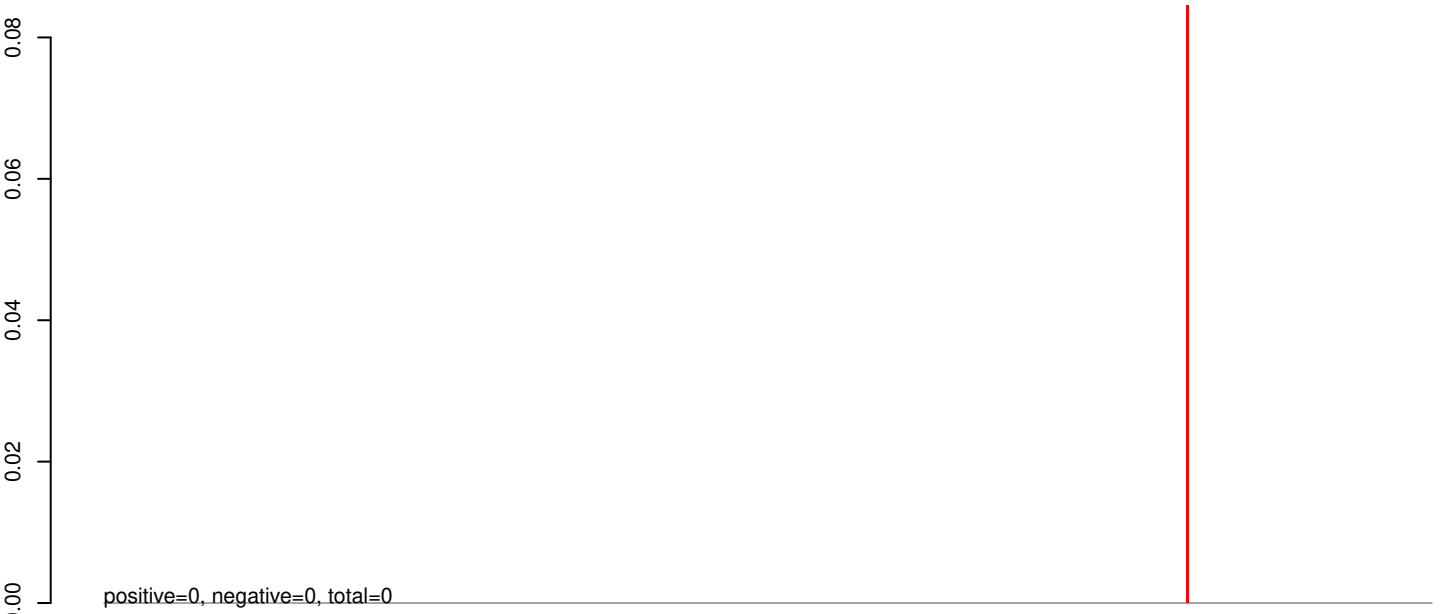


AeAeg_Aag2_BTV1_AK.rep

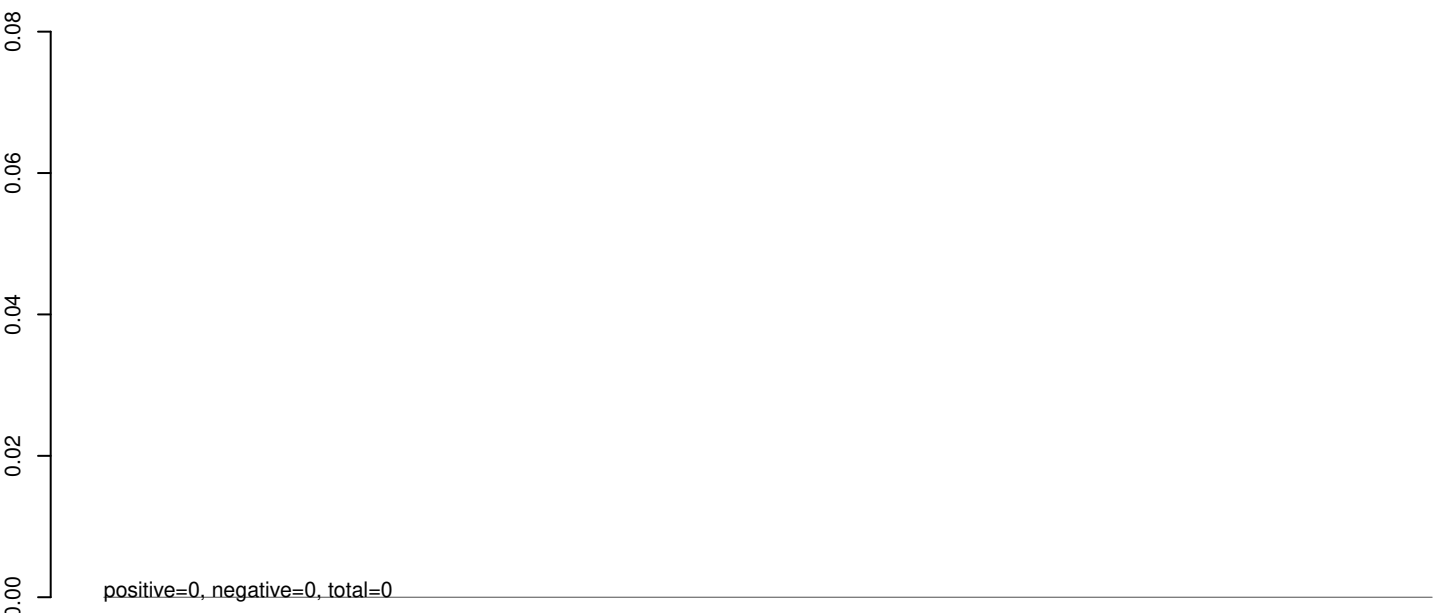


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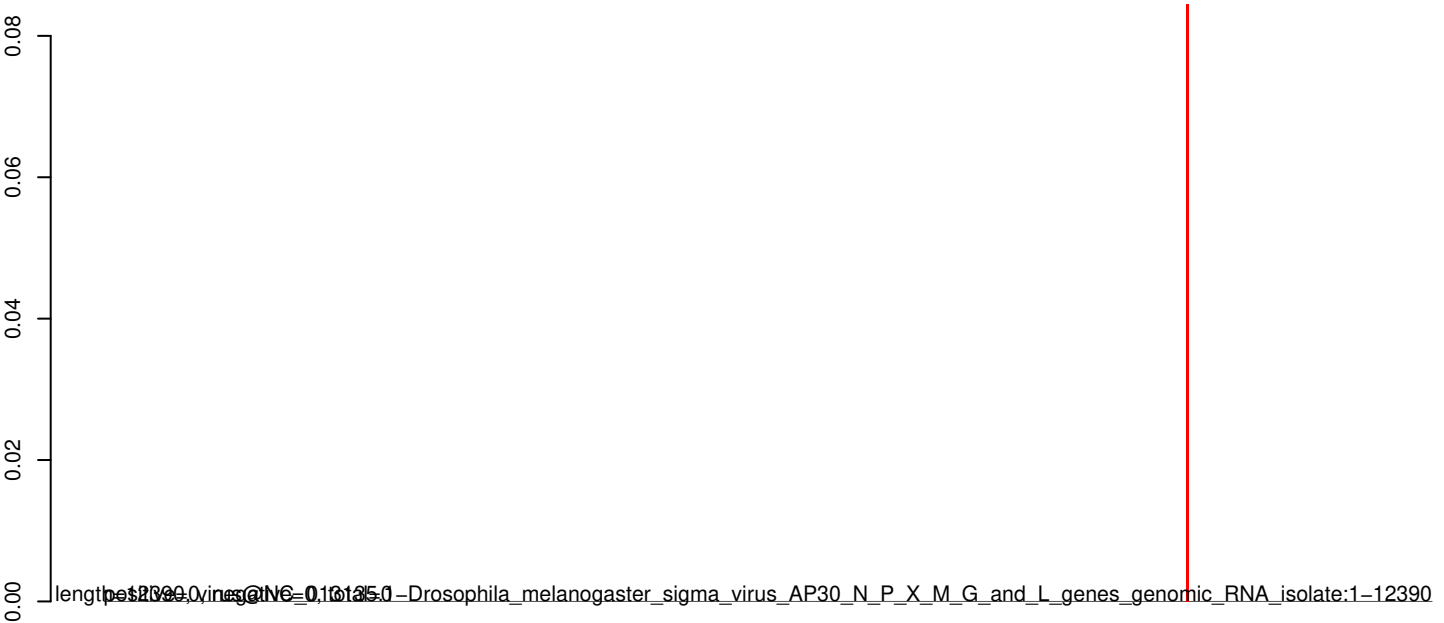
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep

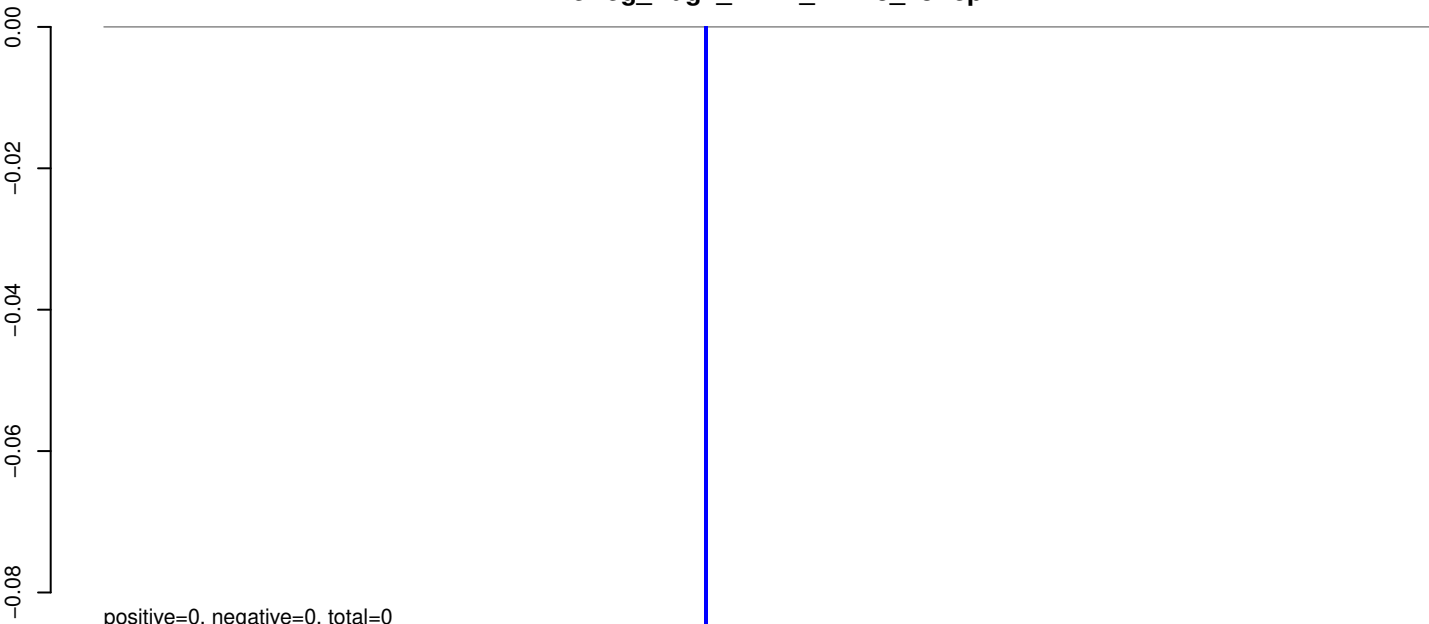


AeAeg_Aag2_BTV1_AK.rep

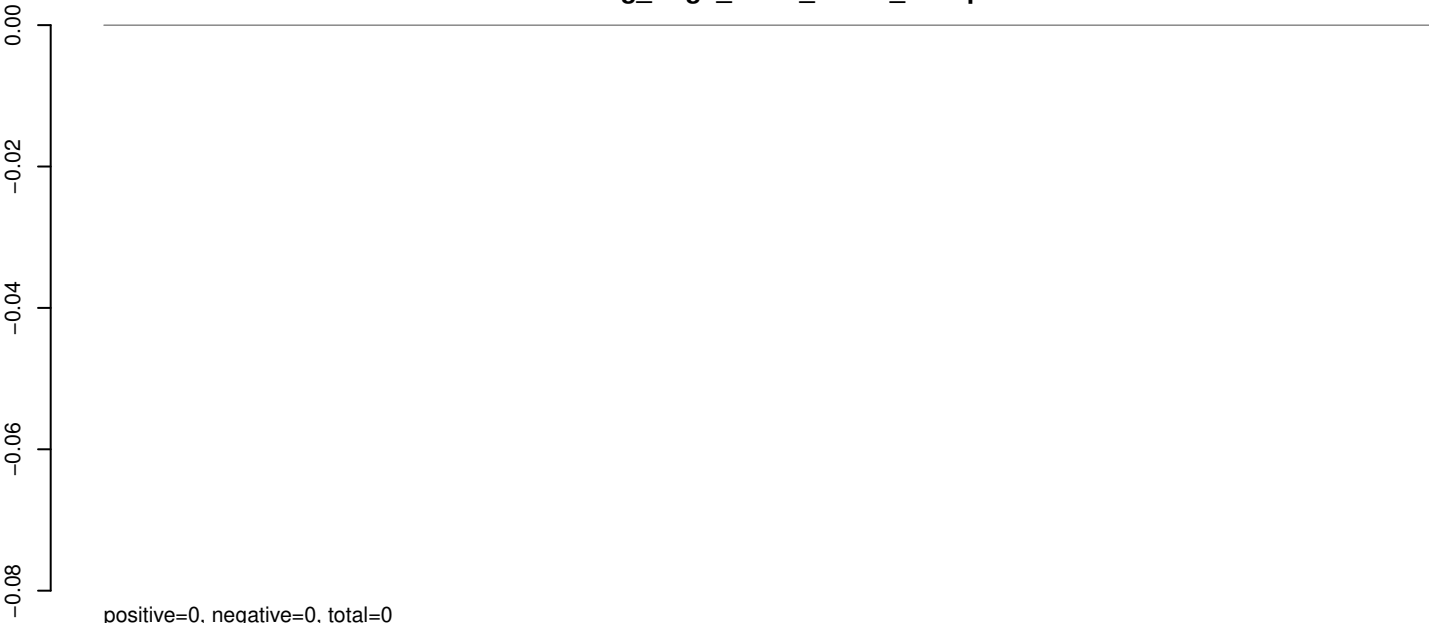


length=2090, coverage=0.16135 - Drosophila_melanogaster_sigma_virus_AP30_N_P_X_M_G_and_L_genes_genomic_RNA_isolate:1-12390

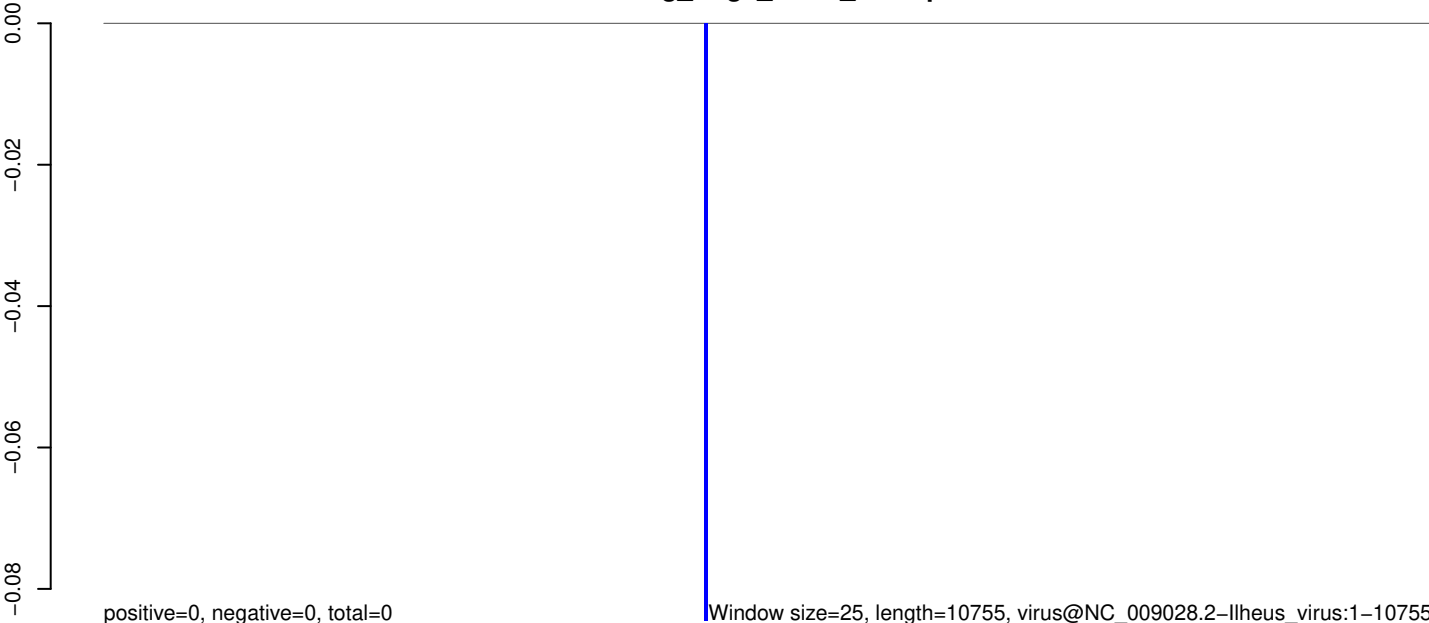
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep

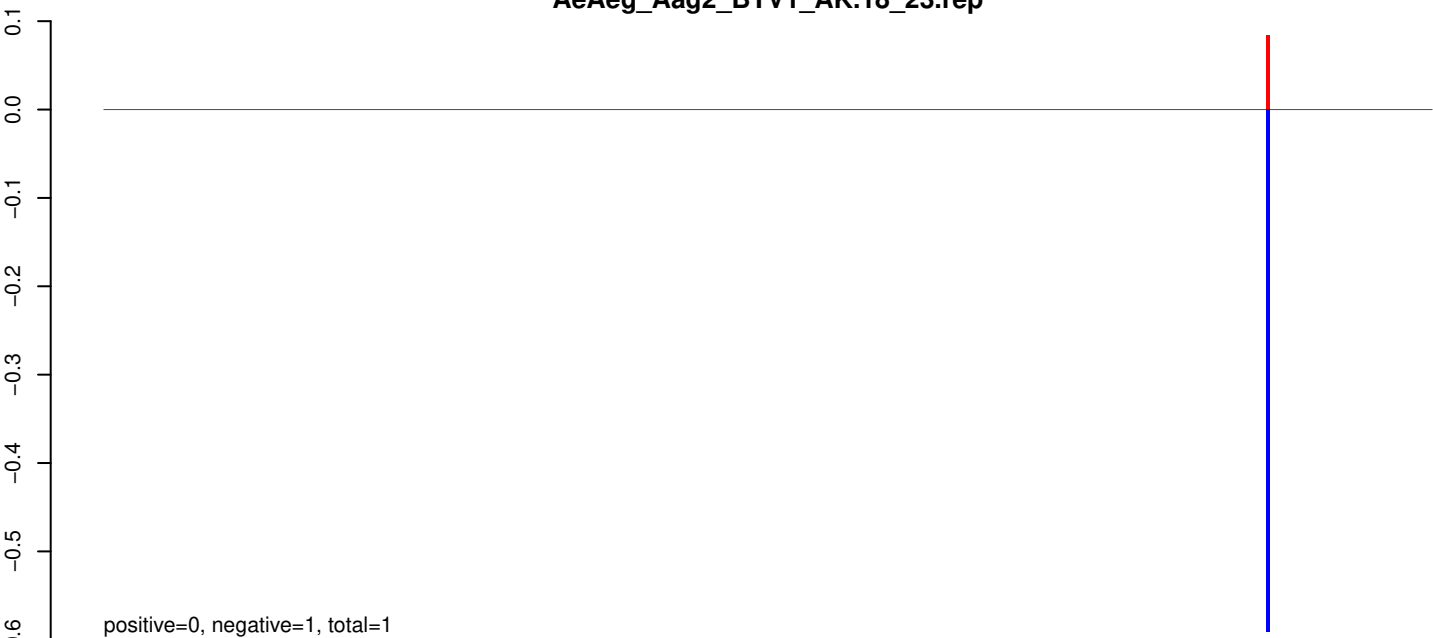


AeAeg_Aag2_BTV1_AK.rep

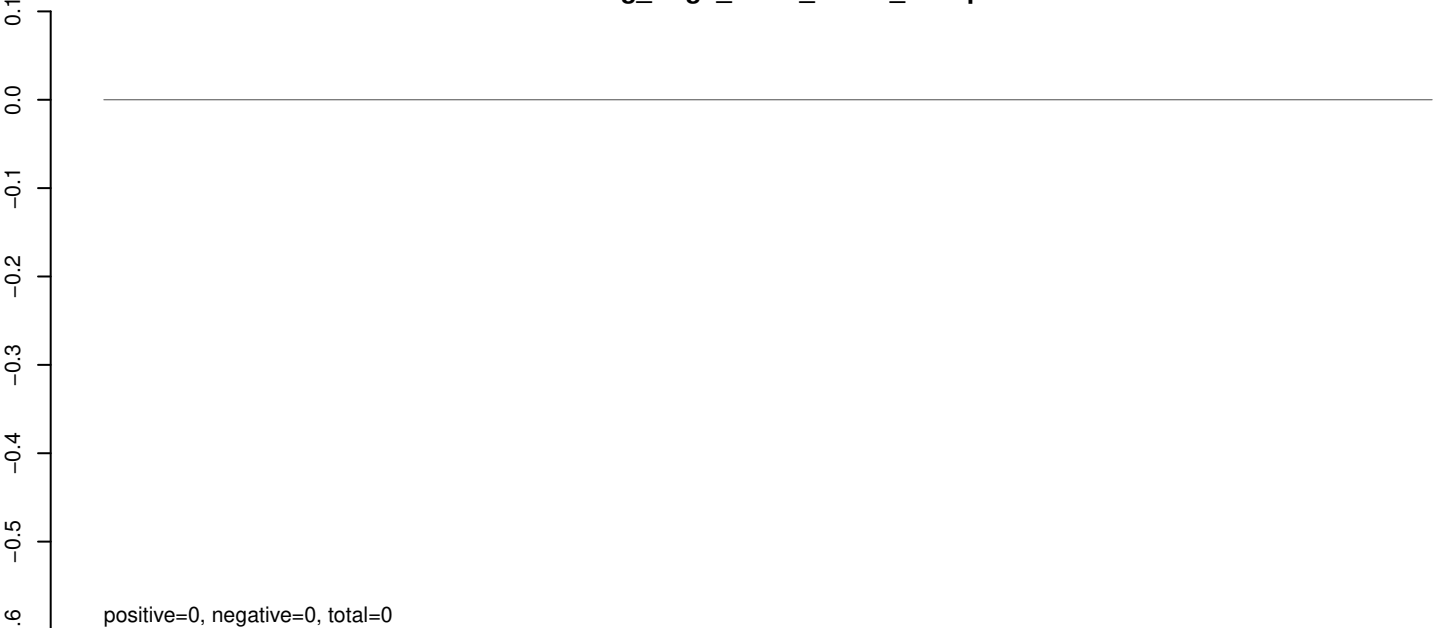


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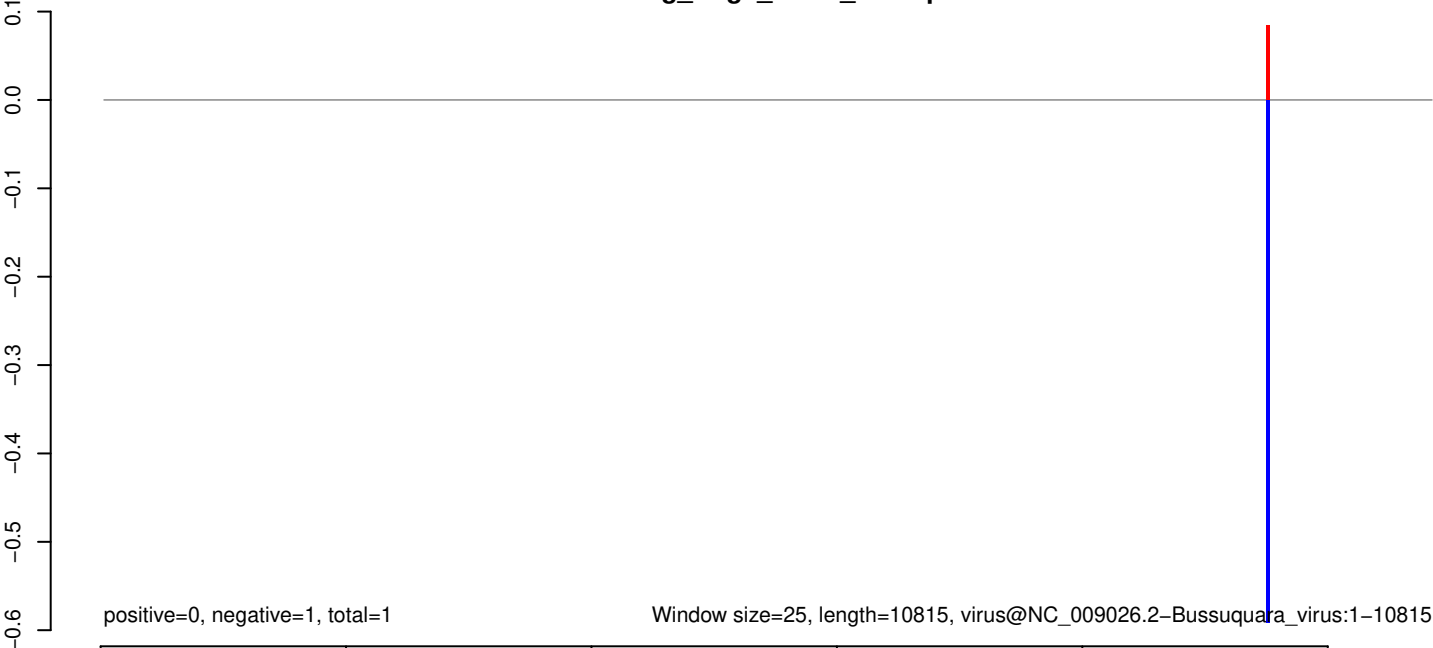
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep

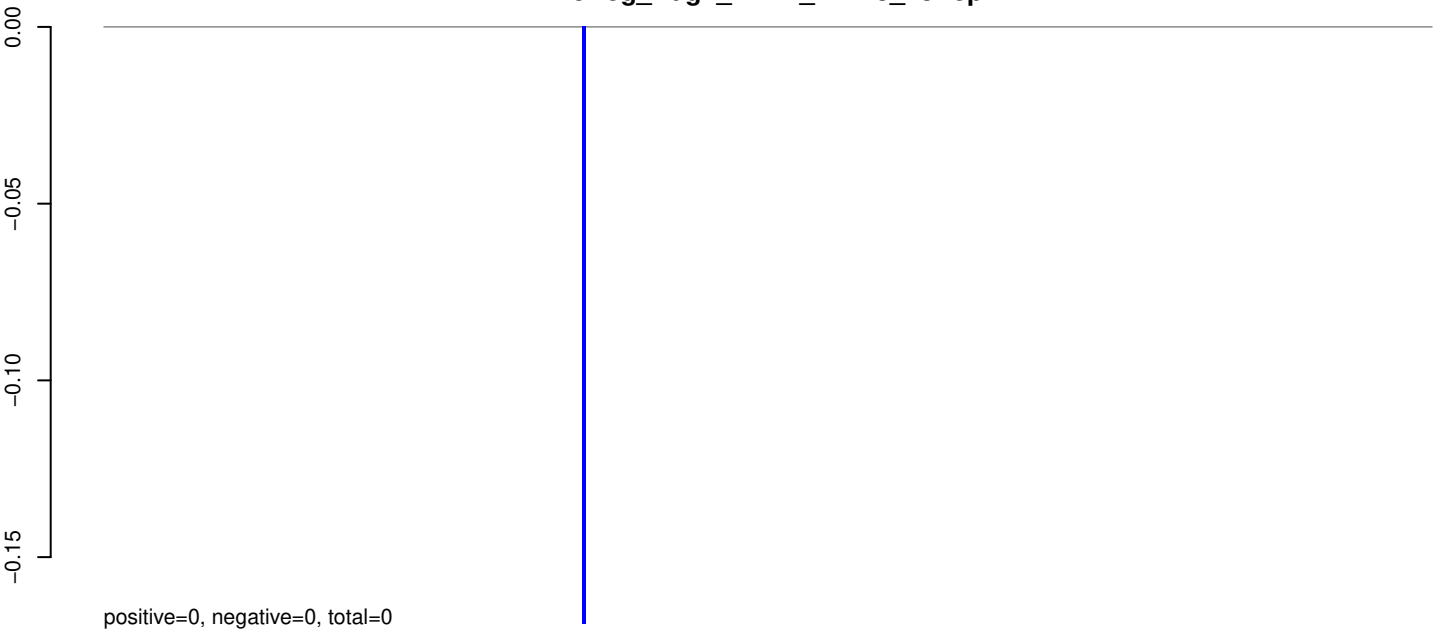


AeAeg_Aag2_BTV1_AK.rep

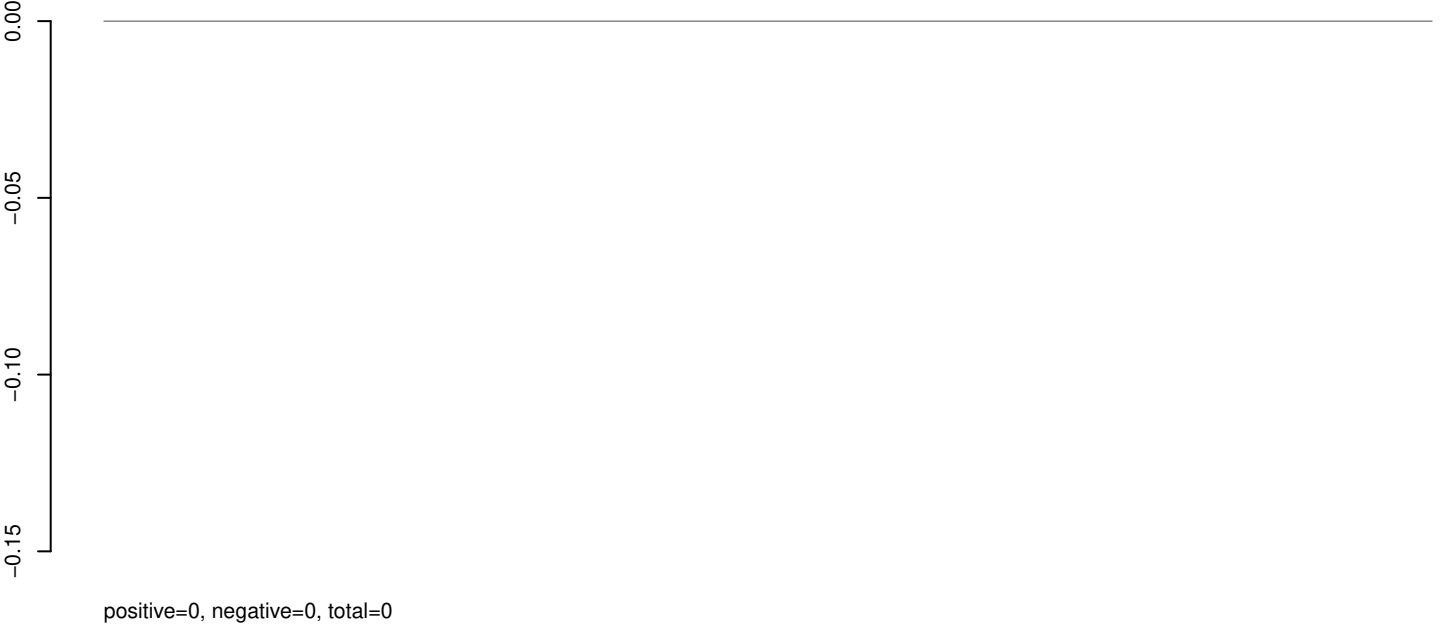


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AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



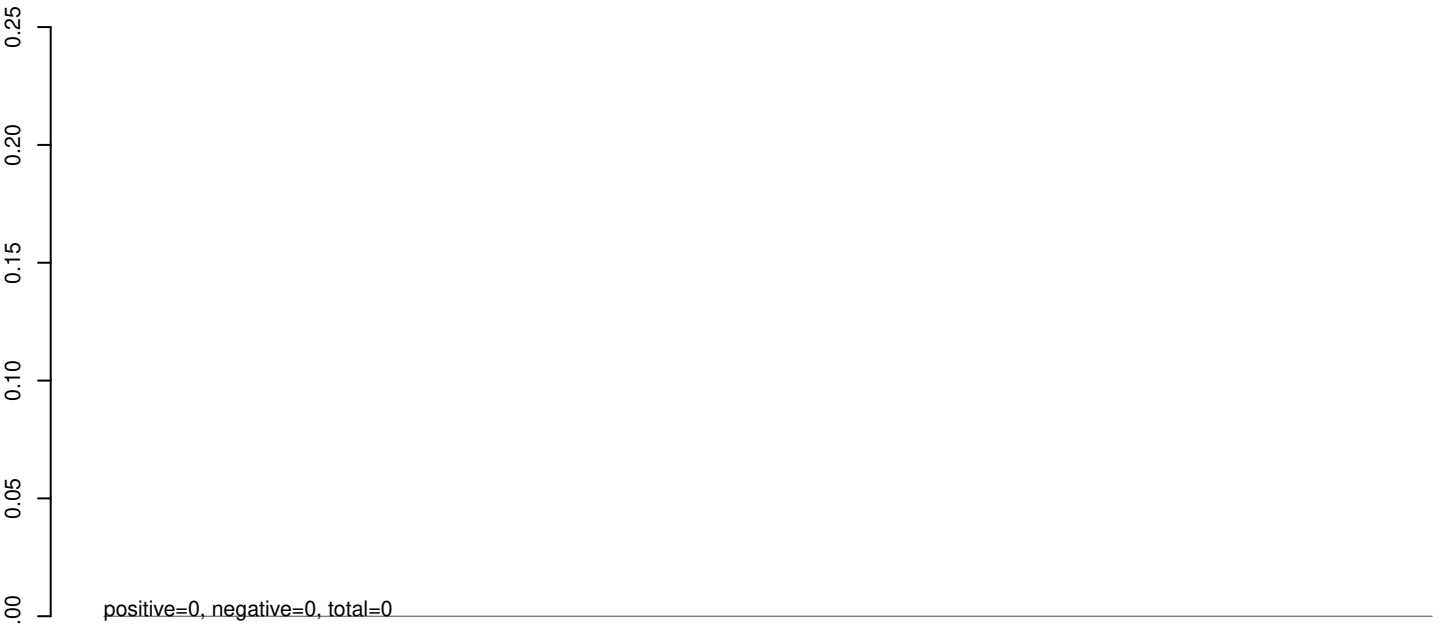
AeAeg_Aag2_BTV1_AK.rep



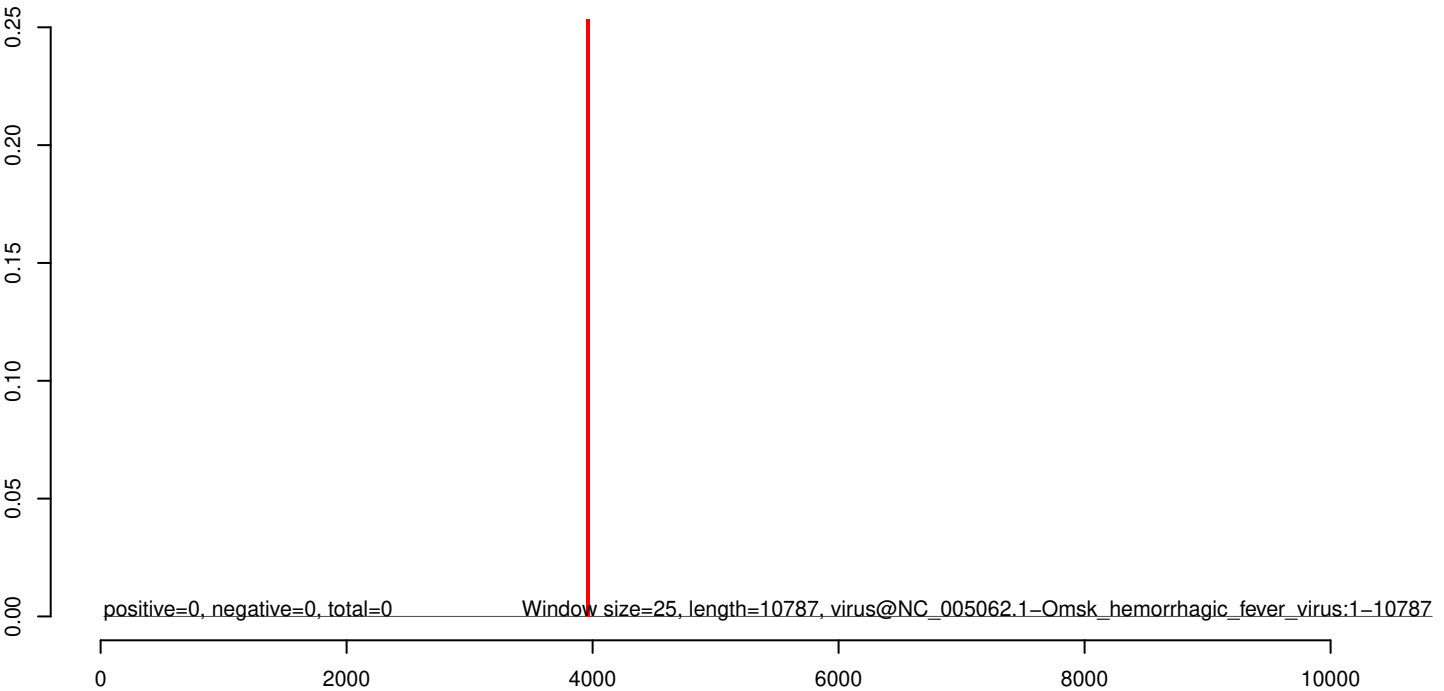
AeAeg_Aag2_BTV1_AK.18_23.rep



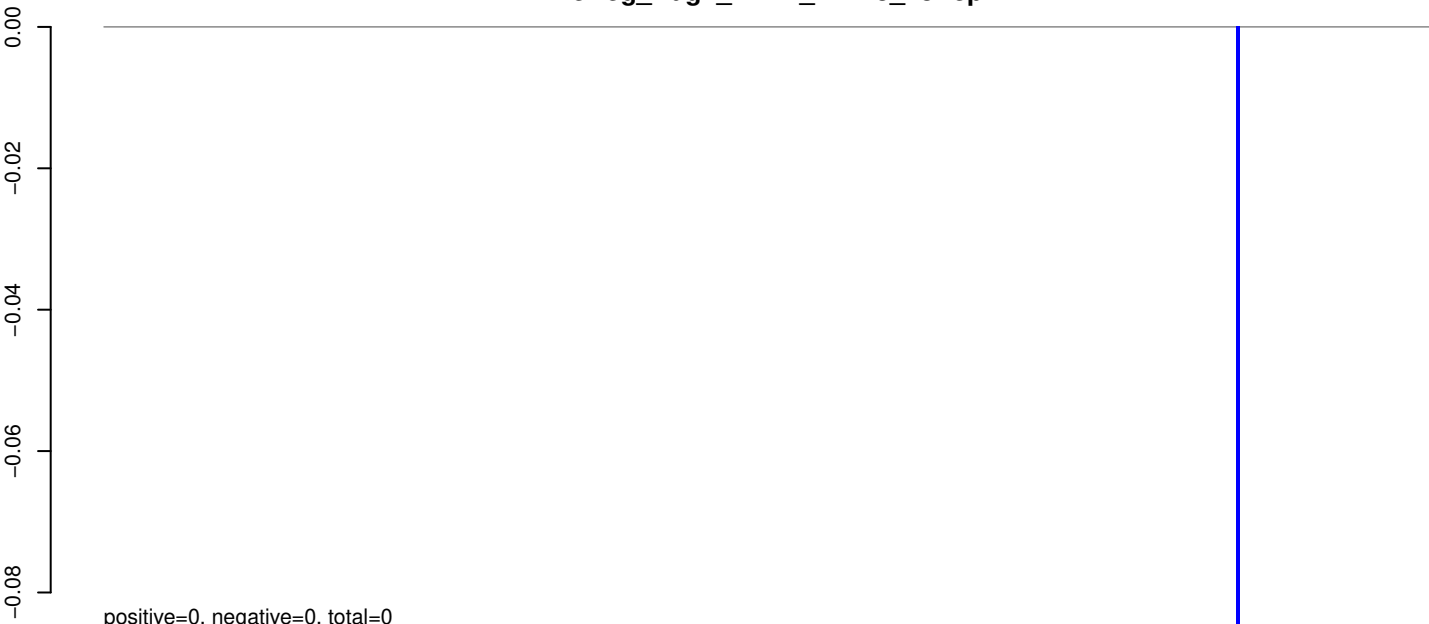
AeAeg_Aag2_BTV1_AK.24_35.rep



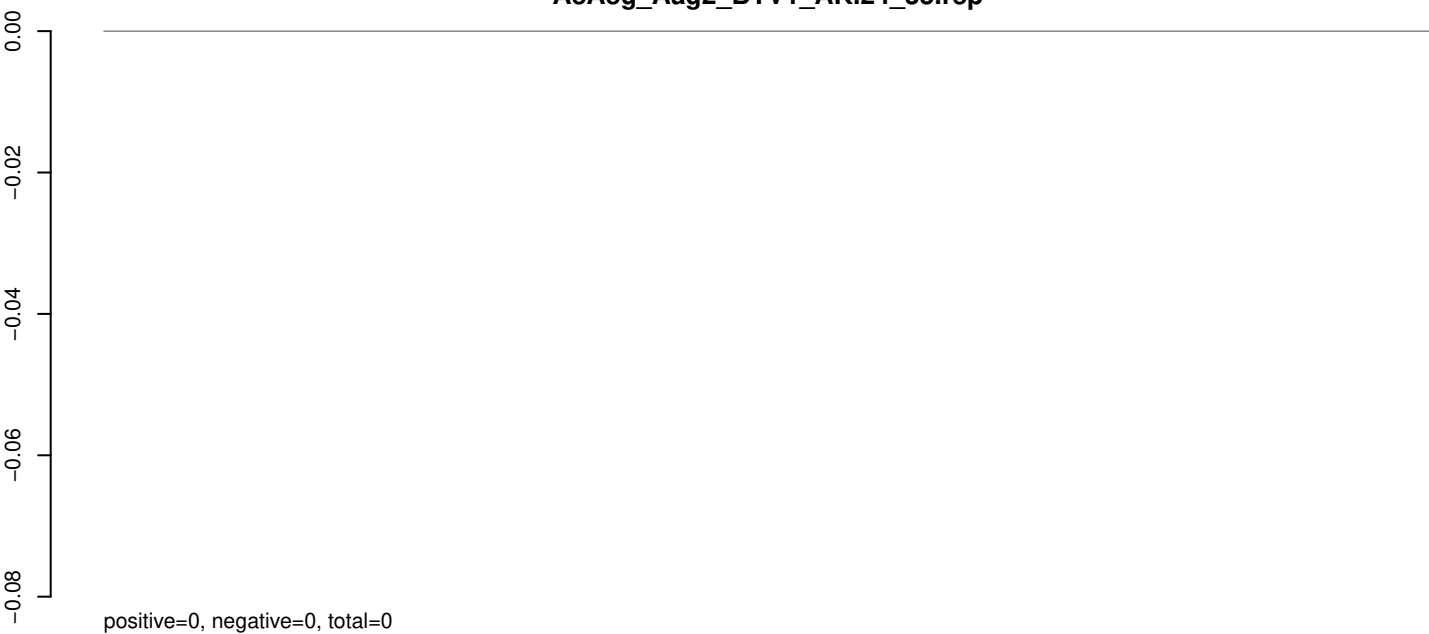
AeAeg_Aag2_BTV1_AK.rep



AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



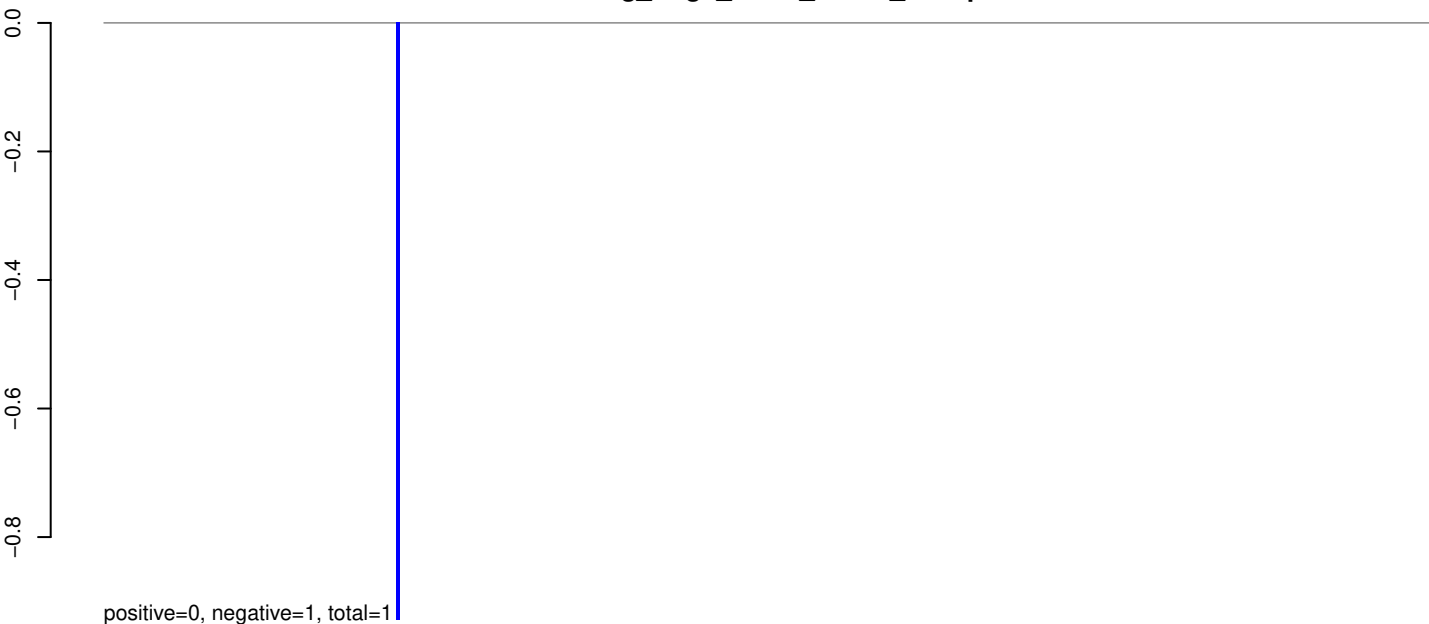
AeAeg_Aag2_BTV1_AK.rep



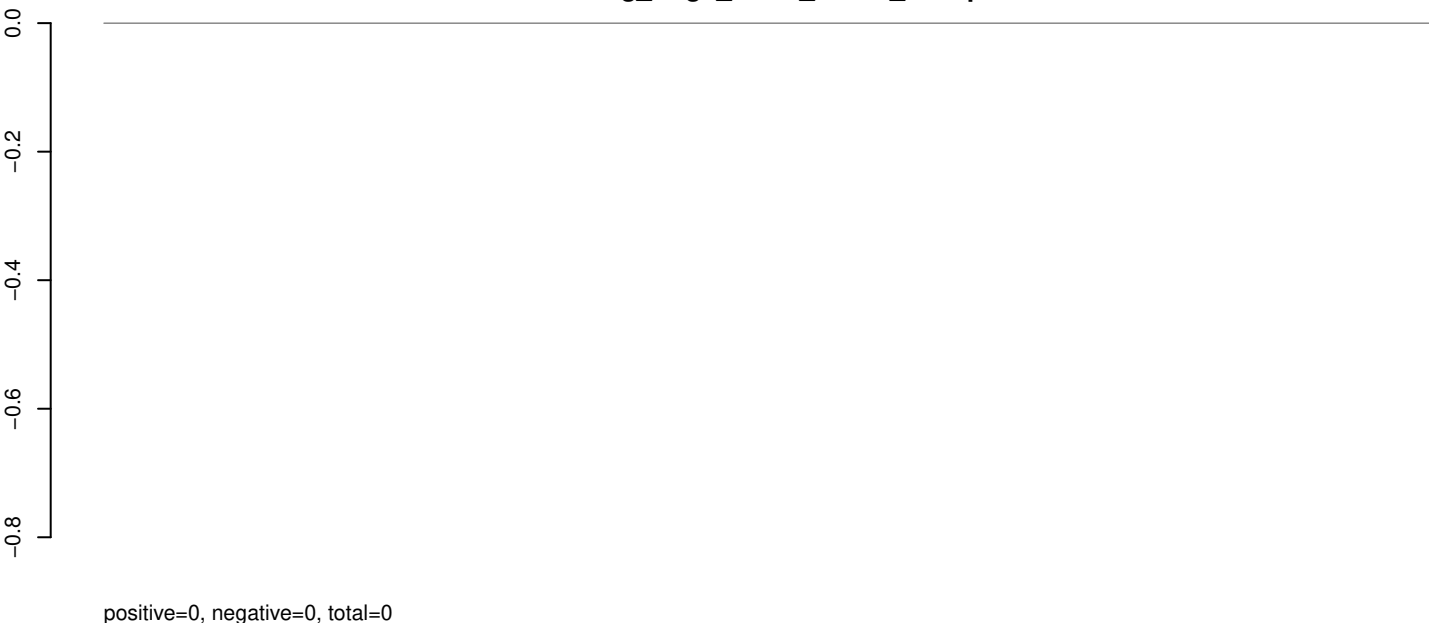
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0 2000 4000 6000 8000 10000

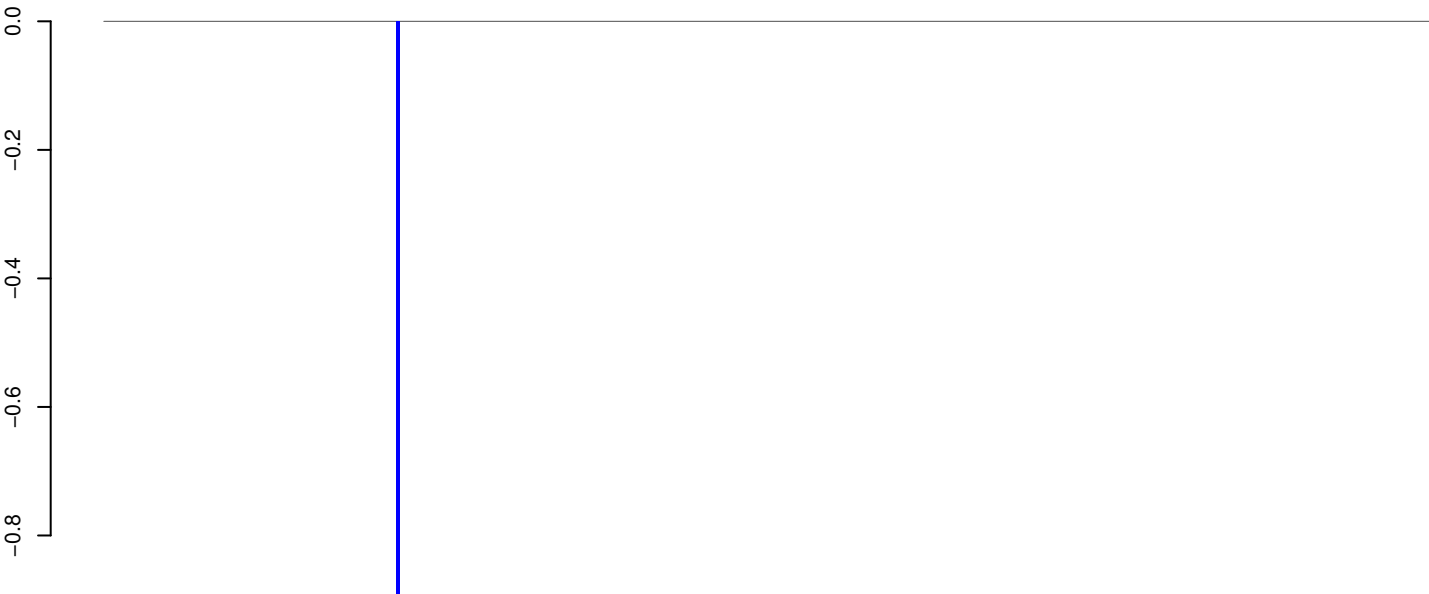
AeAeg_Aag2_BTV1_AK.18_23.rep



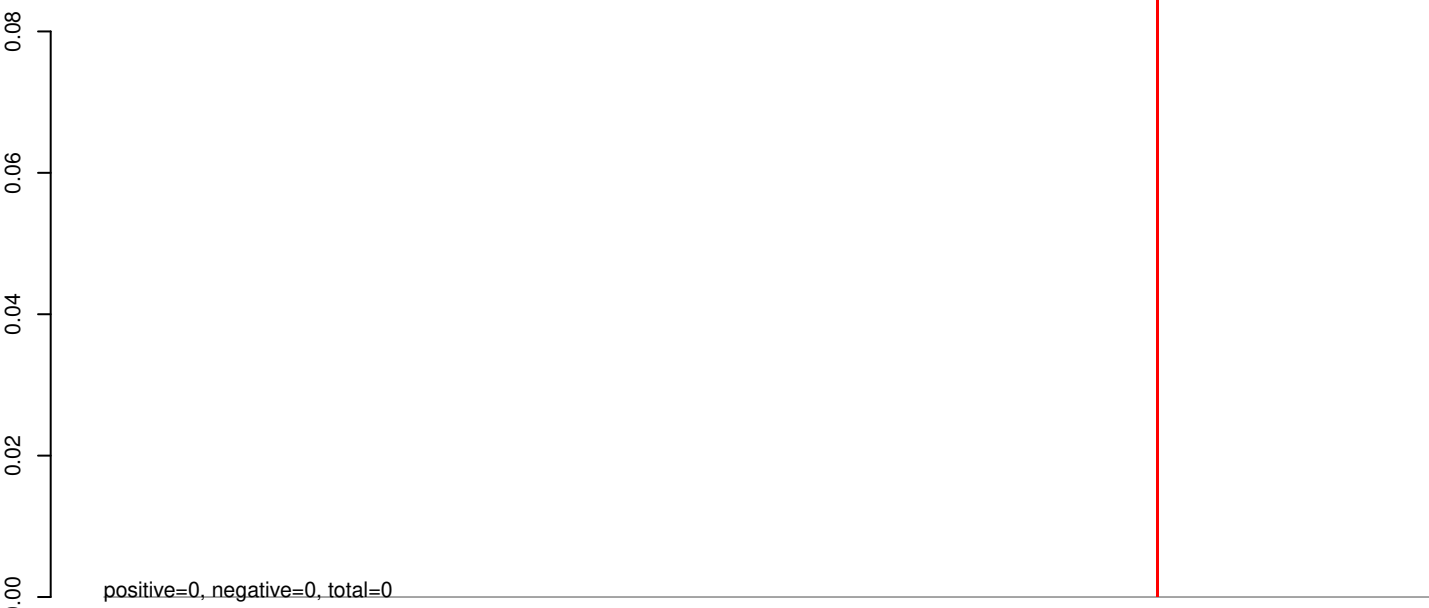
AeAeg_Aag2_BTV1_AK.24_35.rep



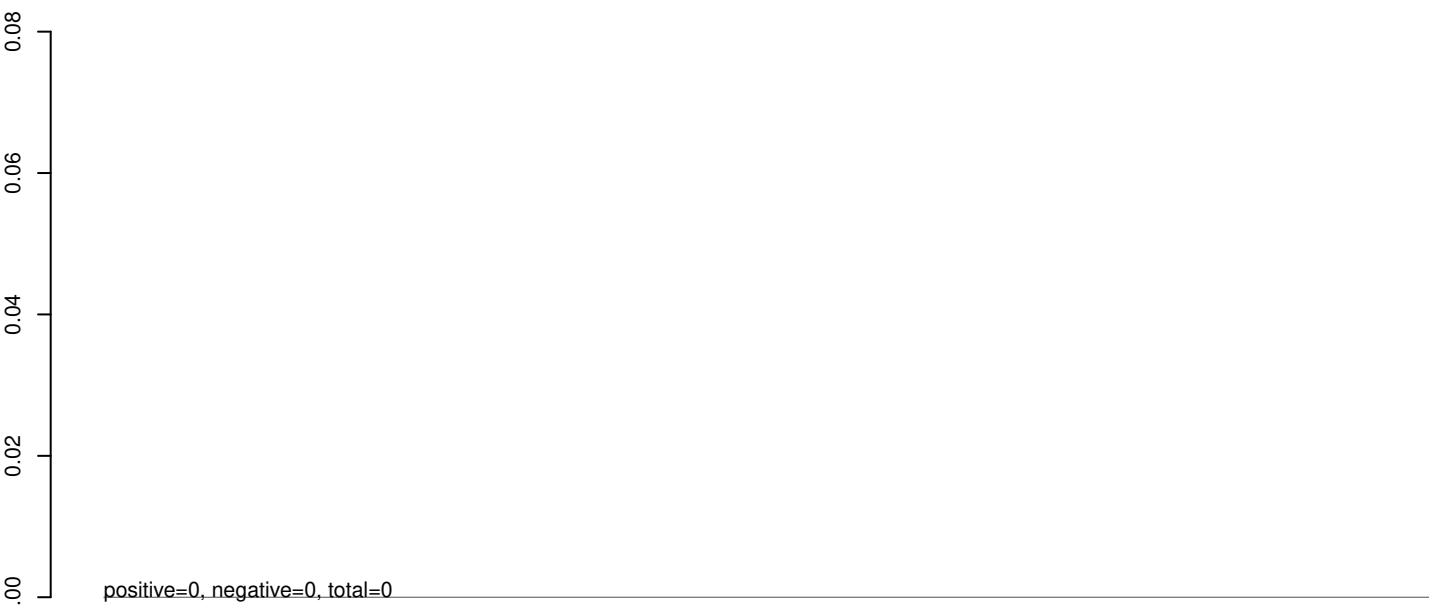
AeAeg_Aag2_BTV1_AK.rep



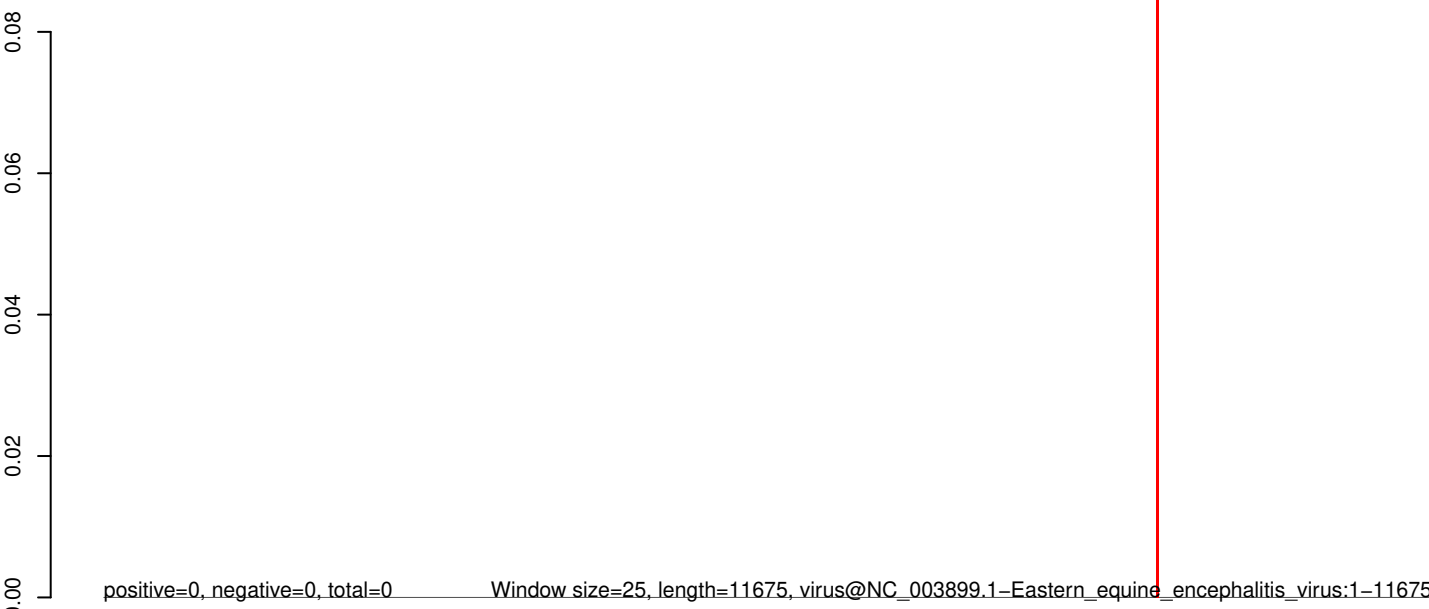
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep

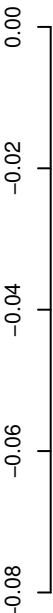


AeAeg_Aag2_BTV1_AK.rep



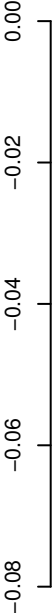
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AeAeg_Aag2_BTV1_AK.18_23.rep



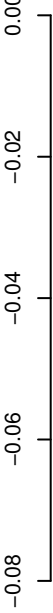
positive=0, negative=0, total=0

AeAeg_Aag2_BTV1_AK.24_35.rep



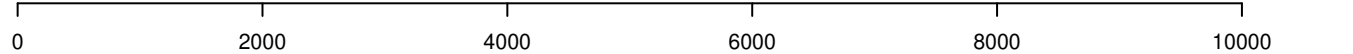
positive=0, negative=0, total=0

AeAeg_Aag2_BTV1_AK.rep

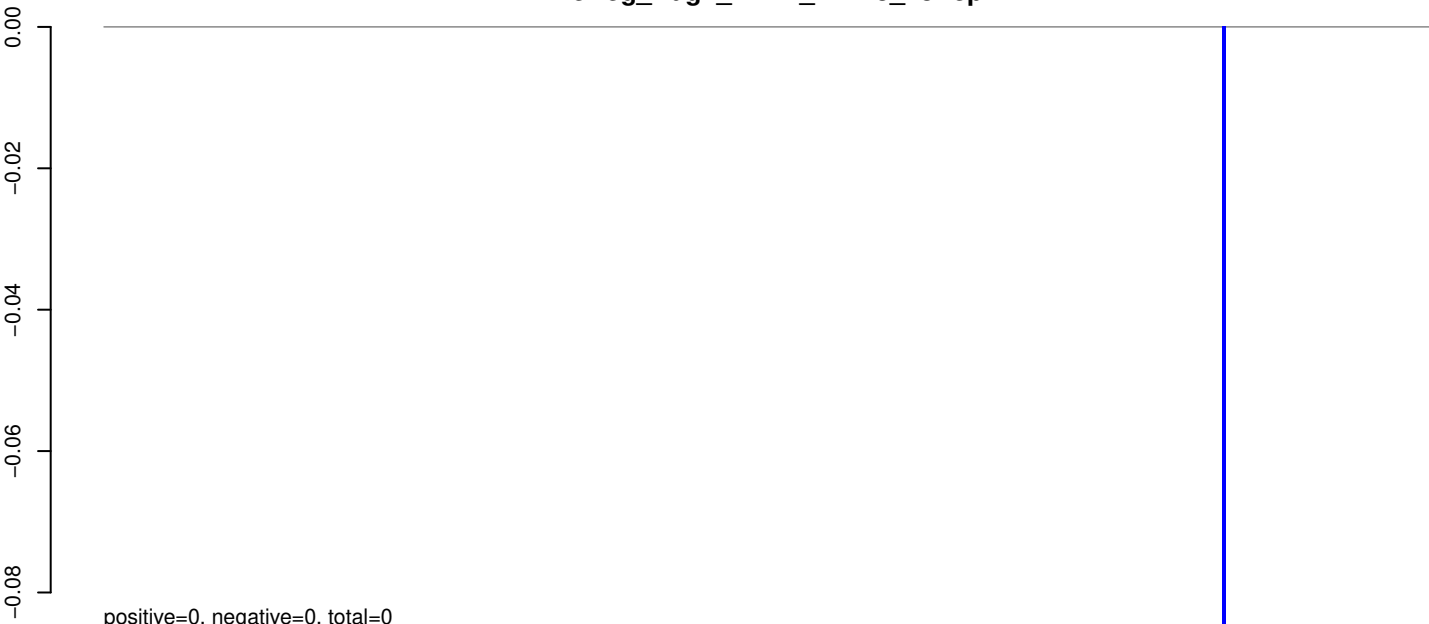


positive=0, negative=0, total=0

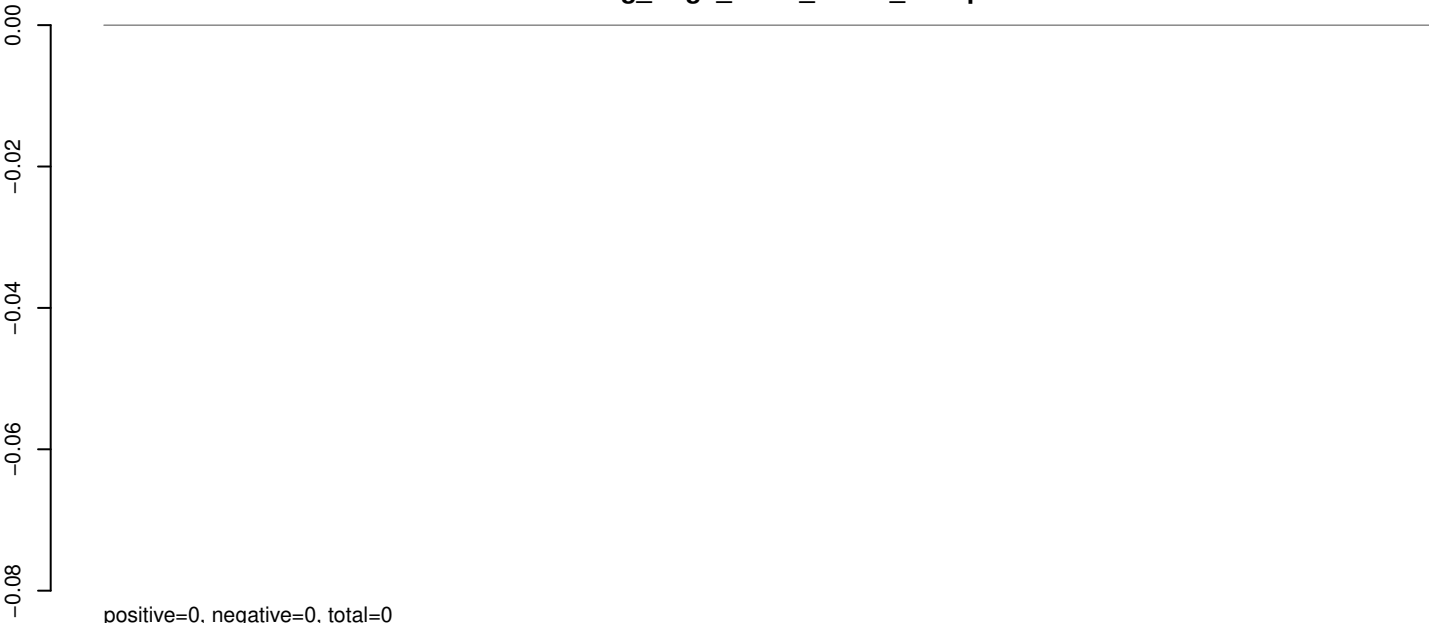
Window size=25, length=10839, virus@NC_003687.1-Powassan_virus:1-10839



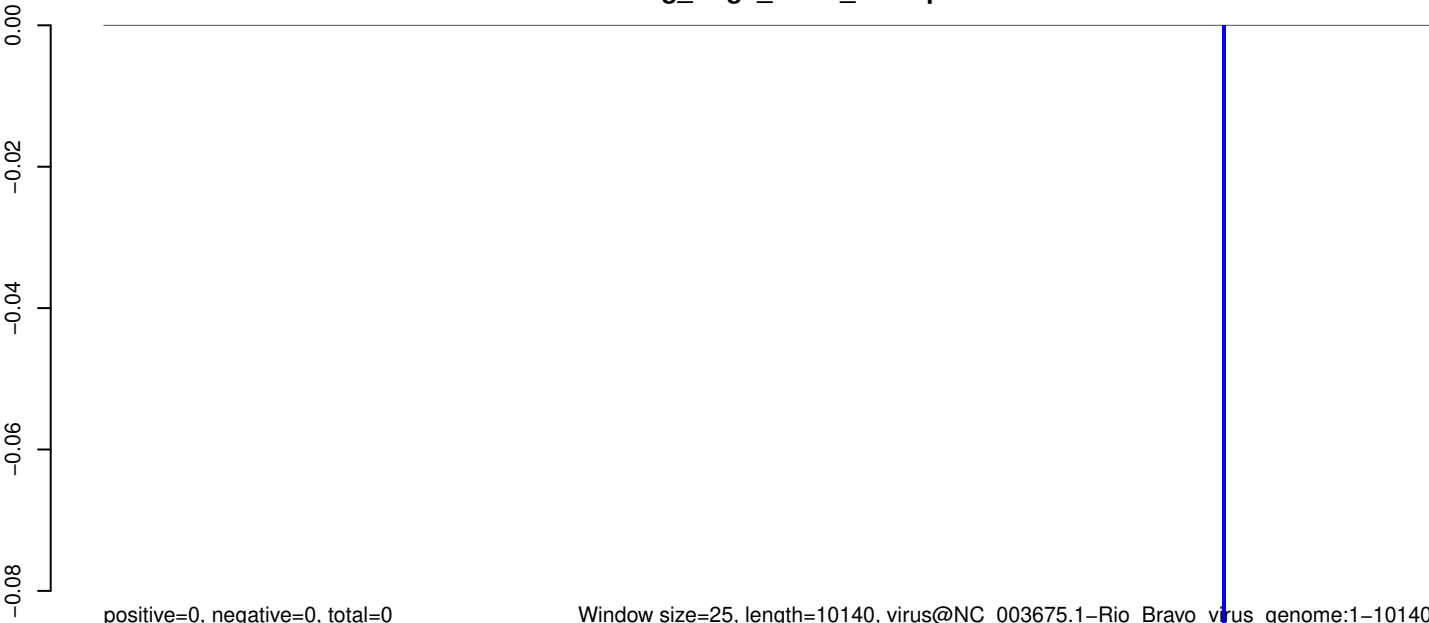
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



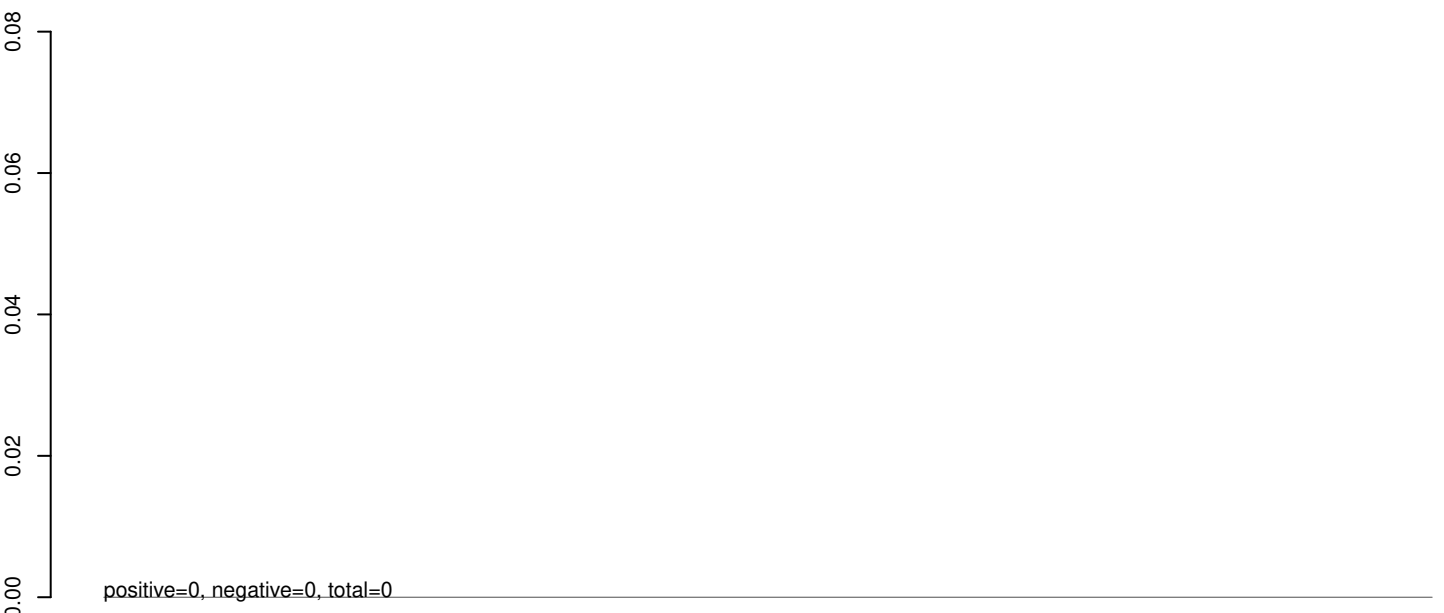
Window size=25, length=10140, virus@NC_003675.1-Rio_Bravo_virus_genome:1-10140

0 2000 4000 6000 8000 10000

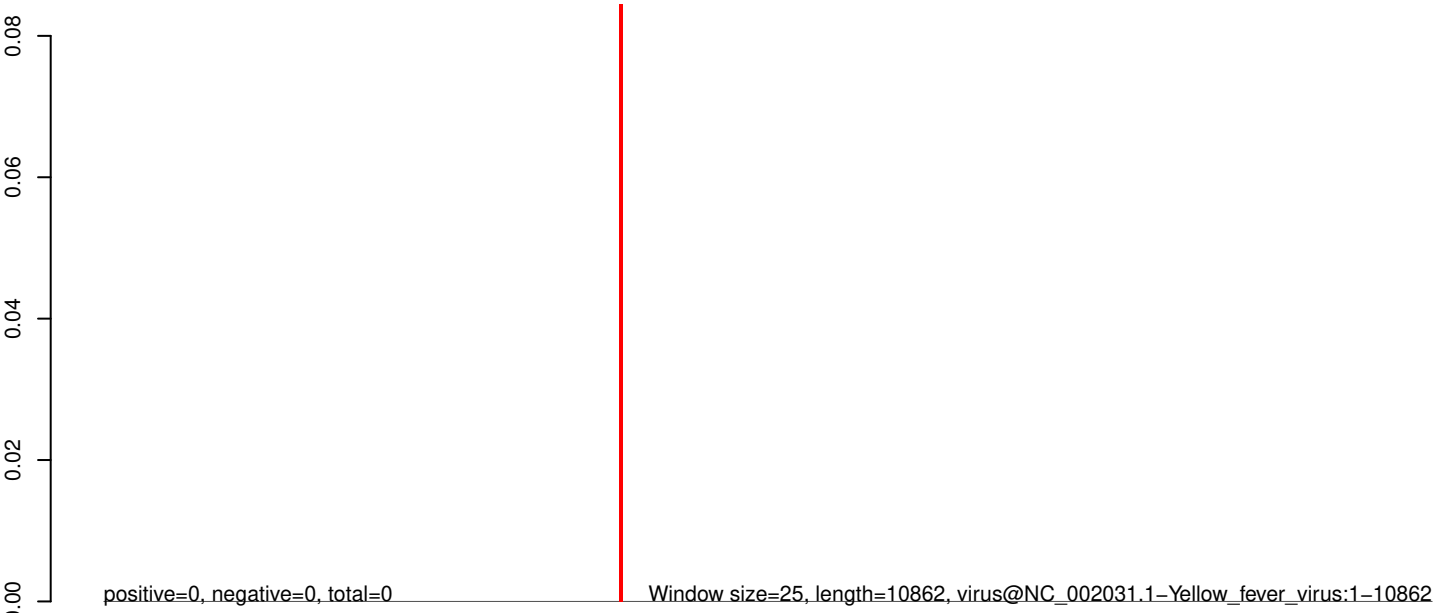
AeAeg_Aag2_BTV1_AK.18_23.rep



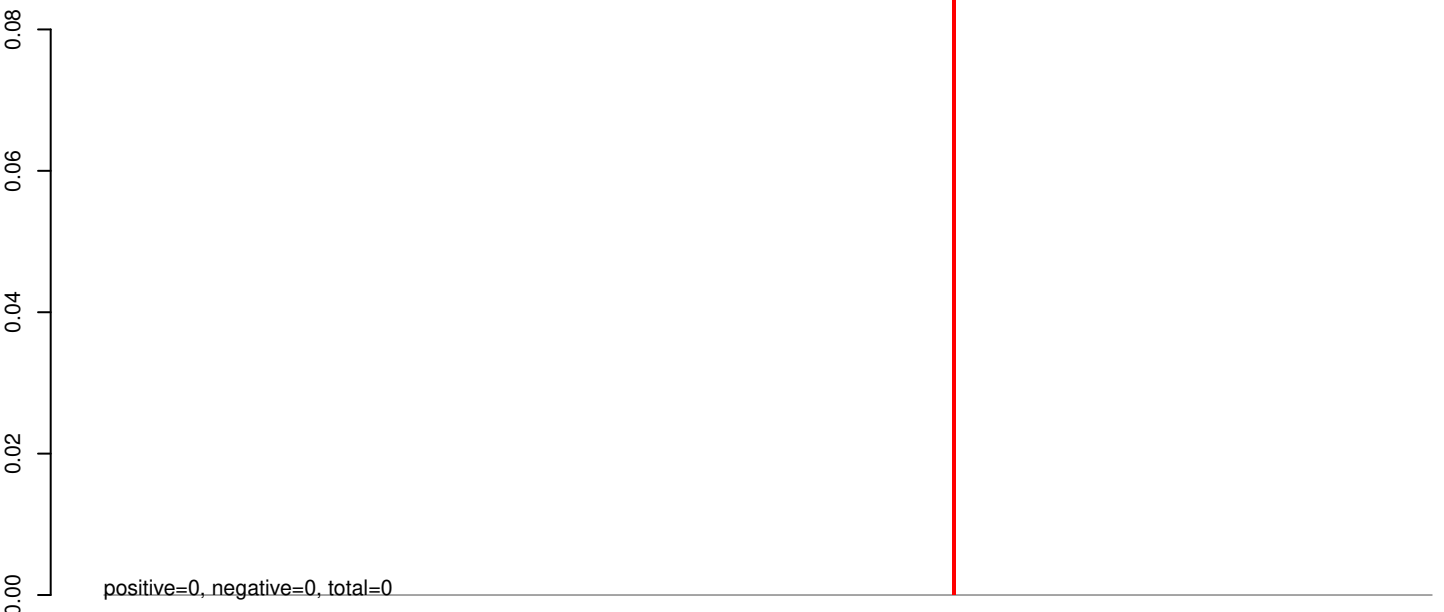
AeAeg_Aag2_BTV1_AK.24_35.rep



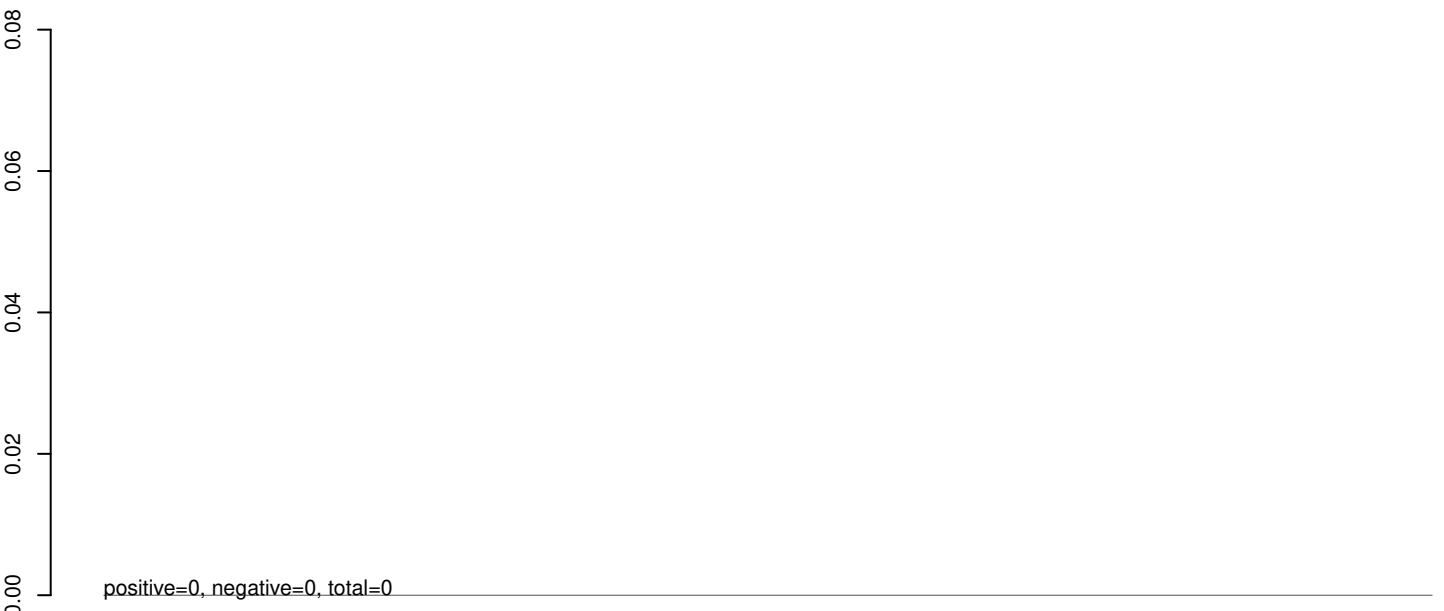
AeAeg_Aag2_BTV1_AK.rep



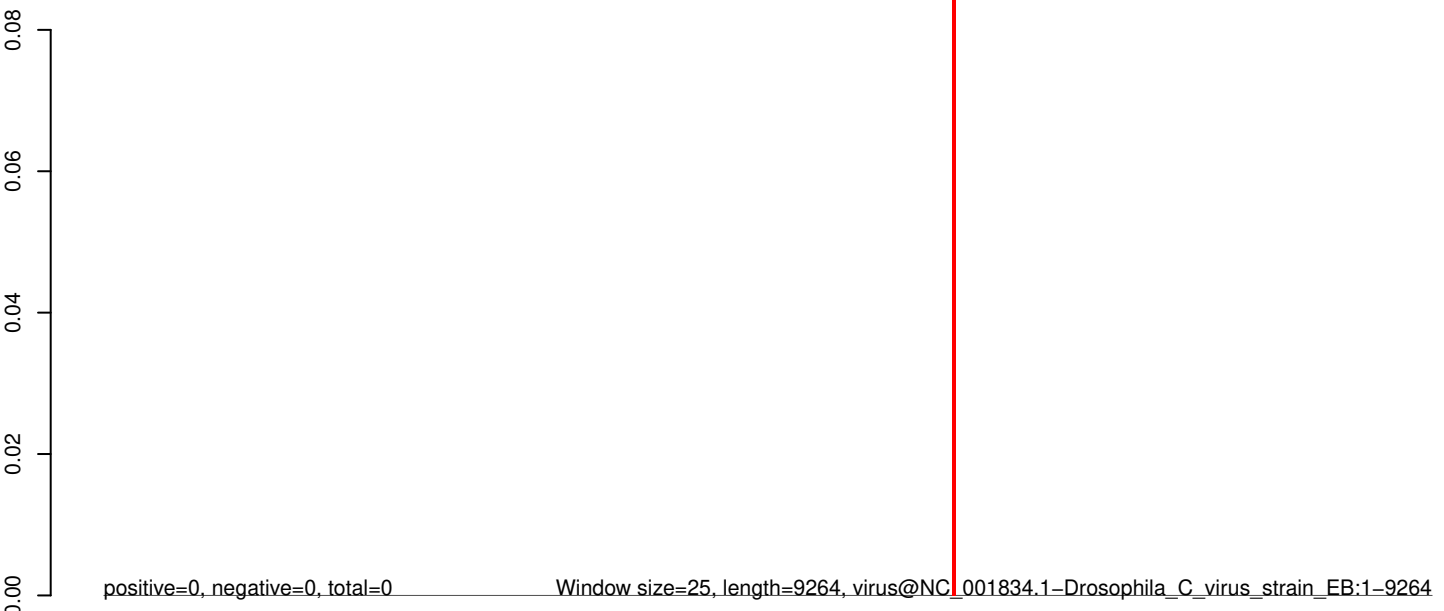
AeAeg_Aag2_BTV1_AK.18_23.rep



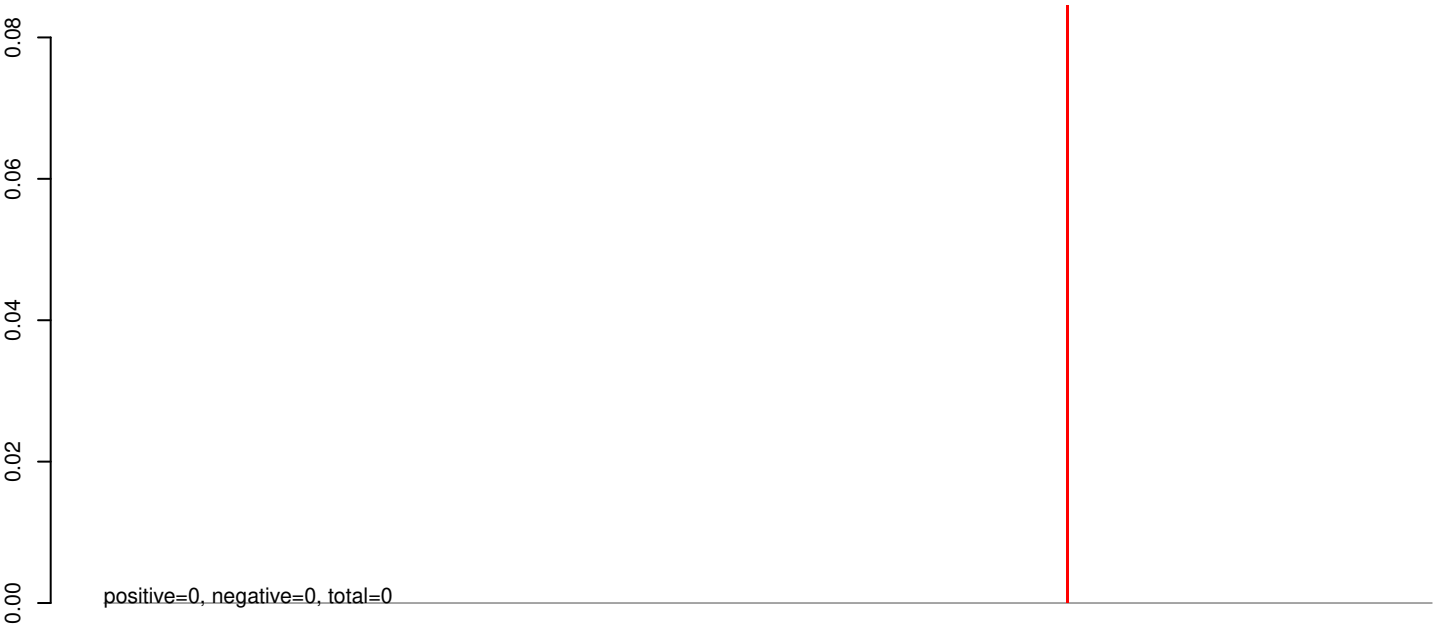
AeAeg_Aag2_BTV1_AK.24_35.rep



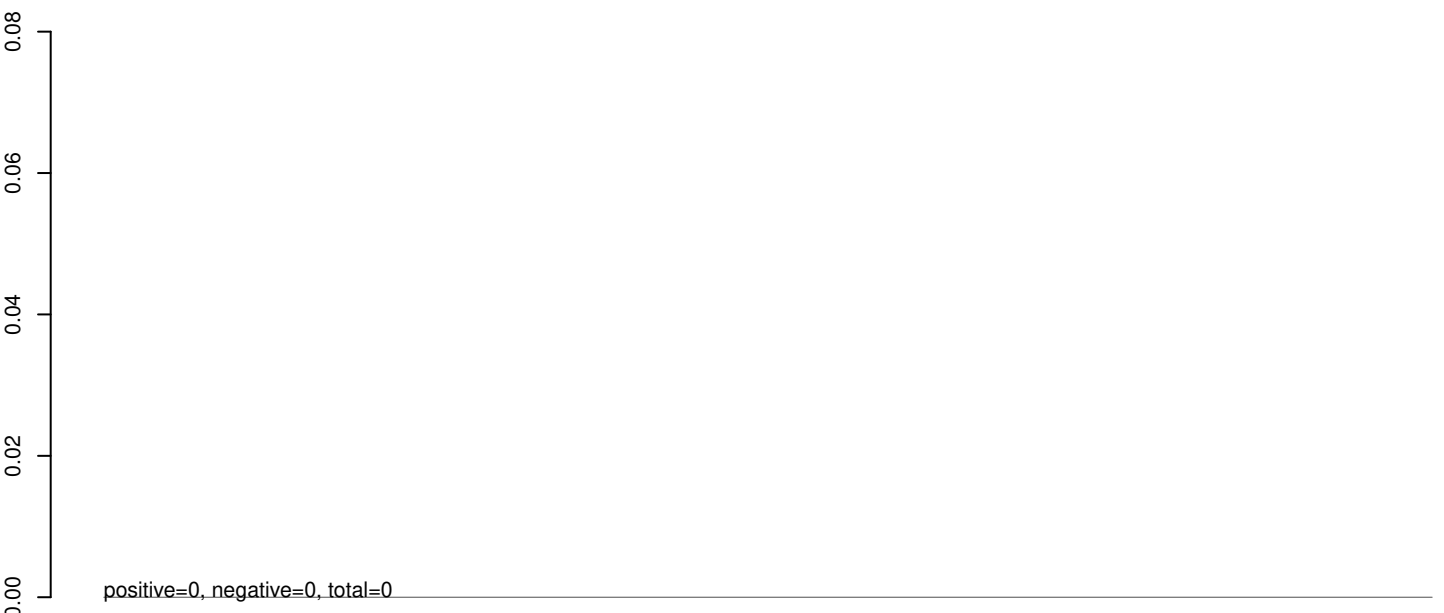
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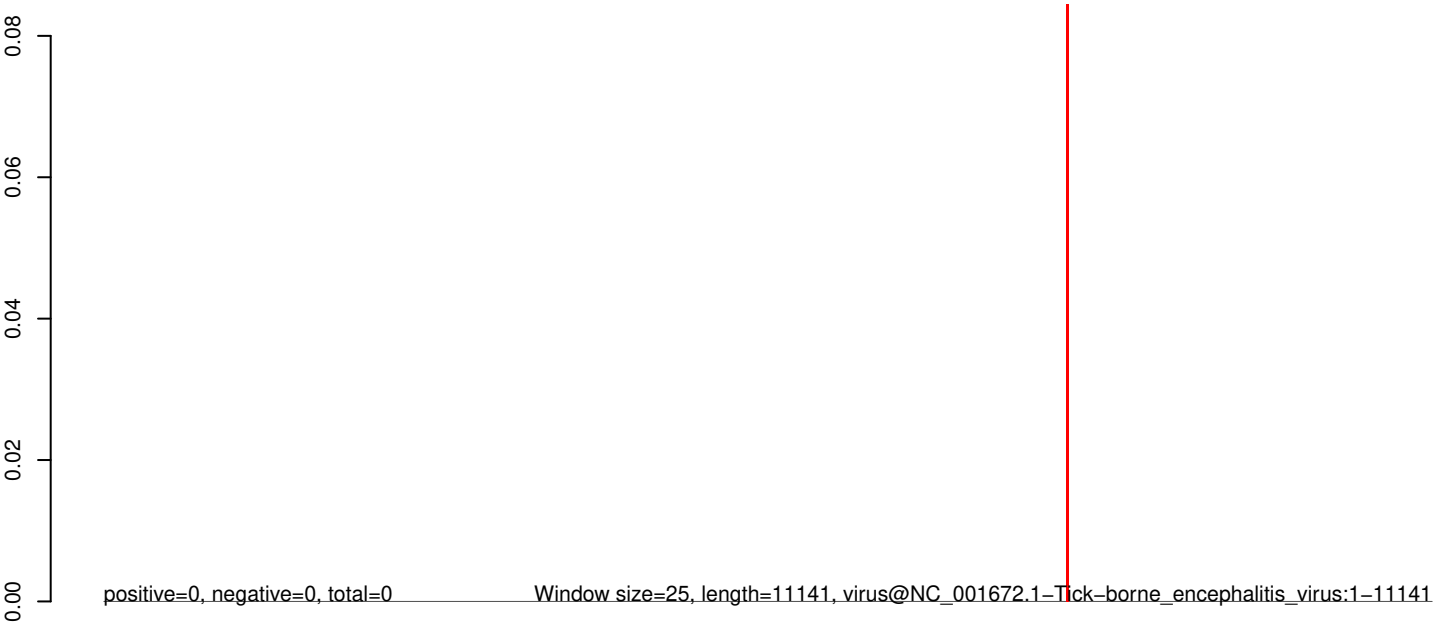
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



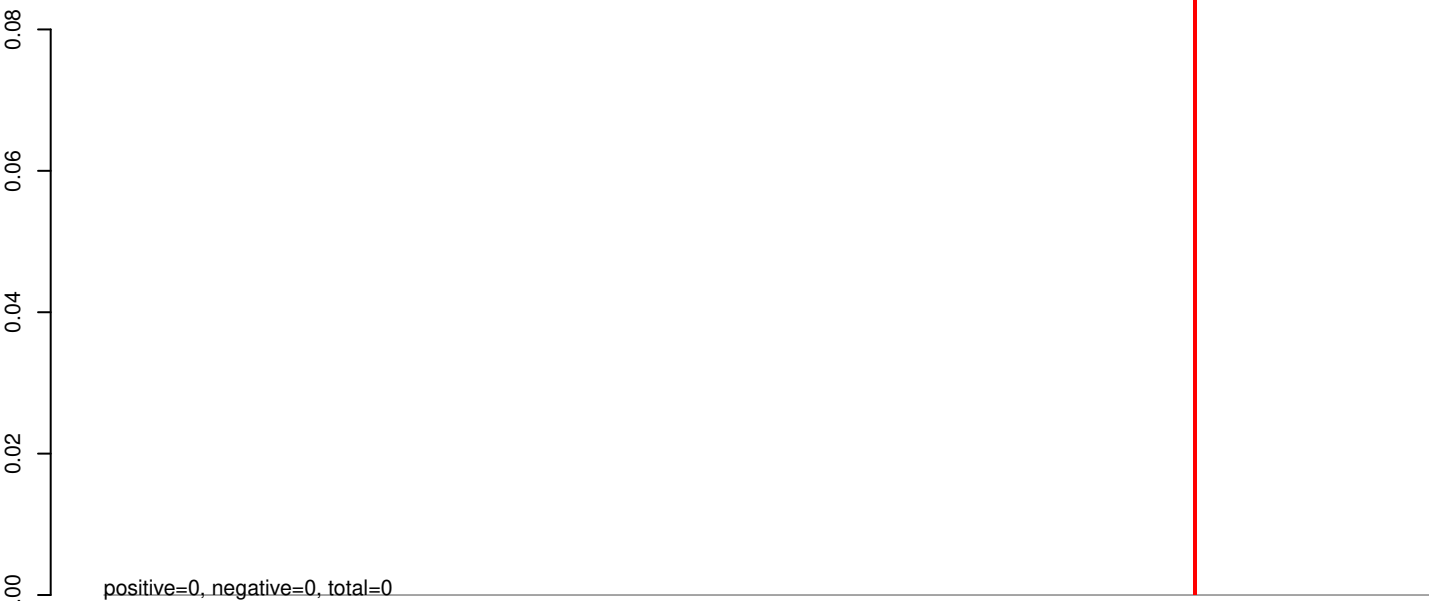
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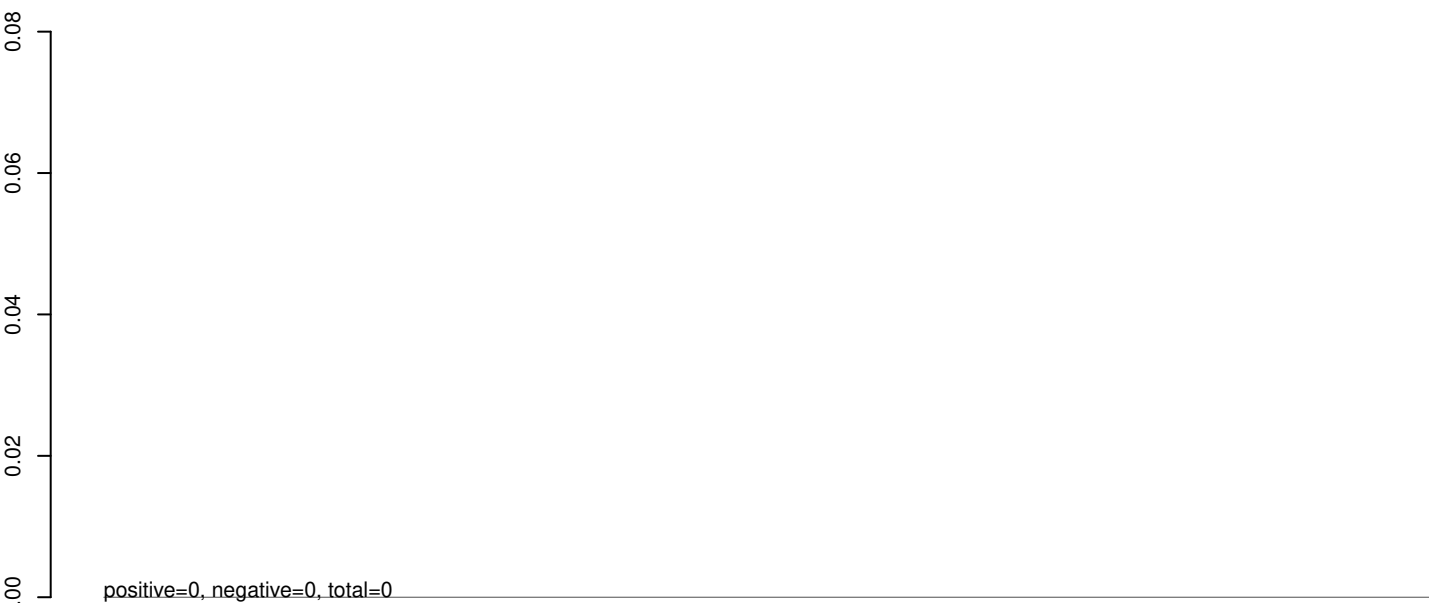
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0 2000 4000 6000 8000 10000

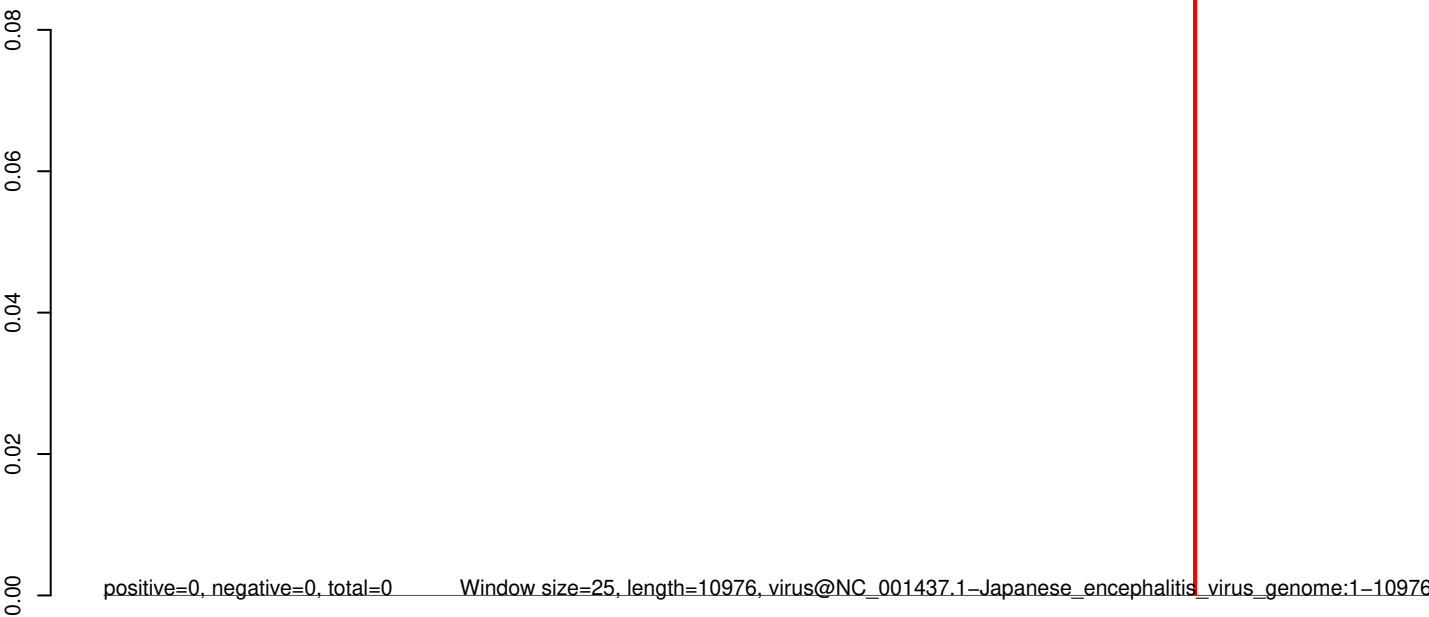
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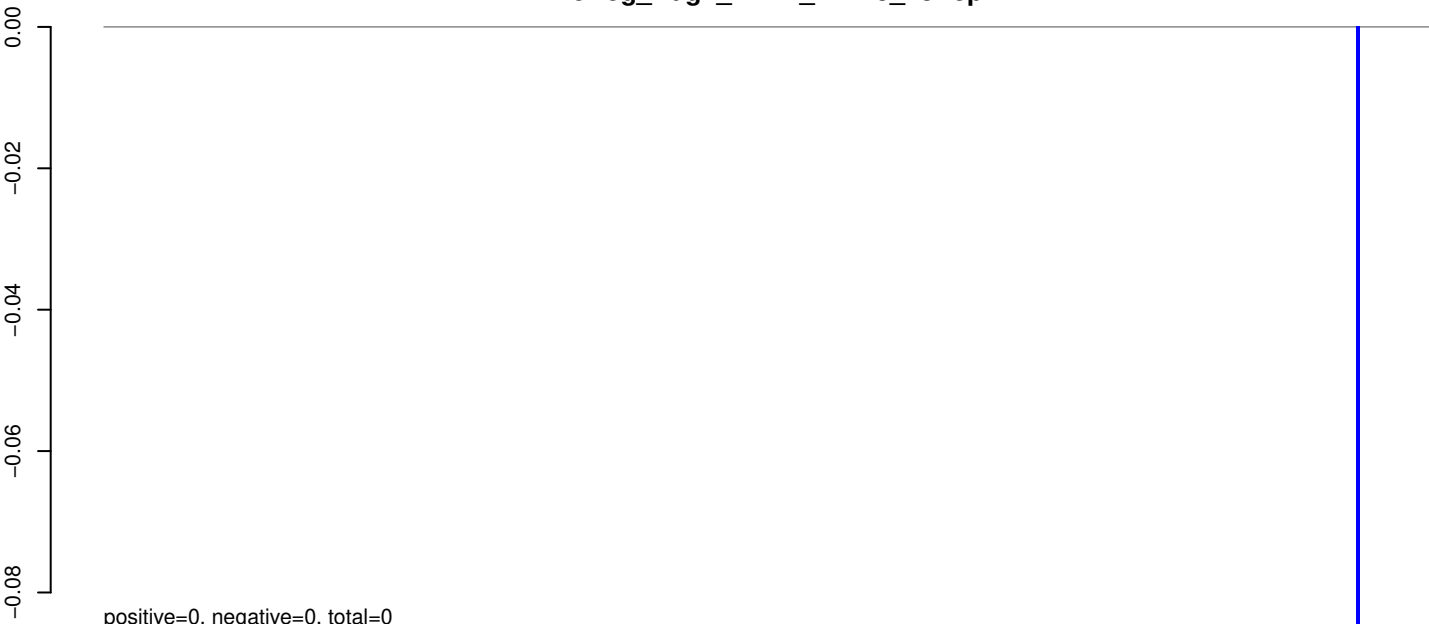
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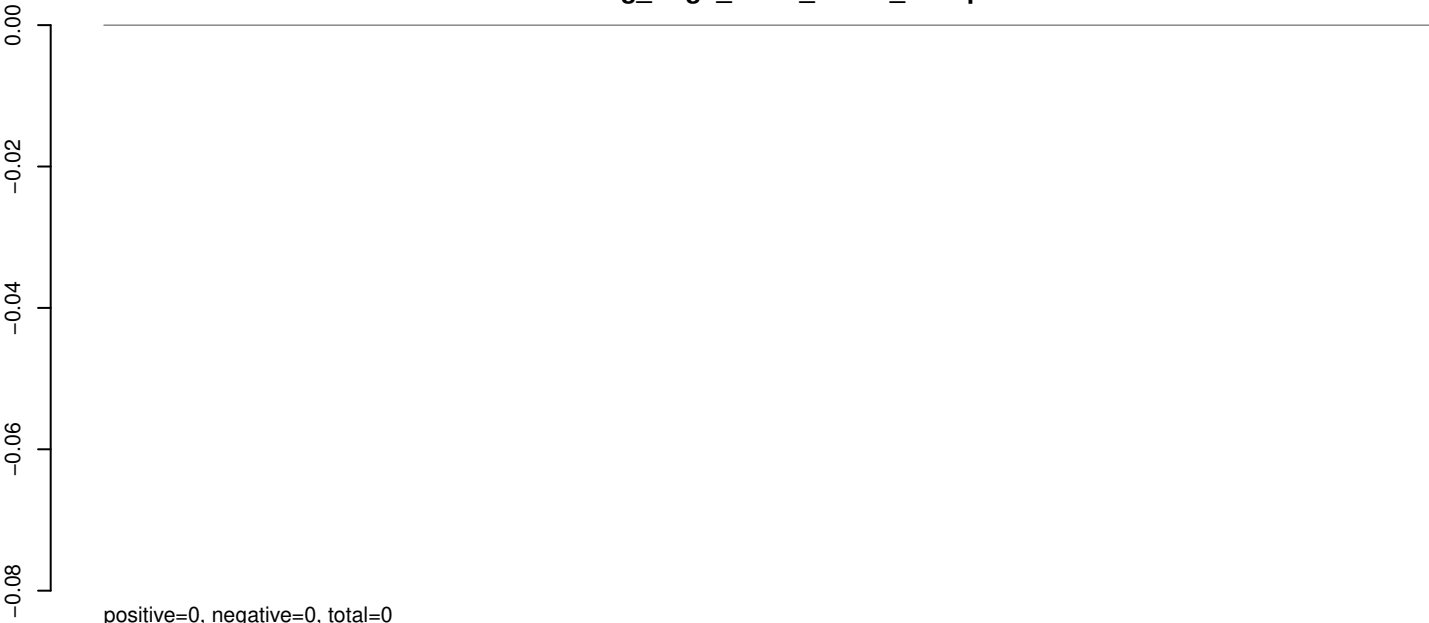
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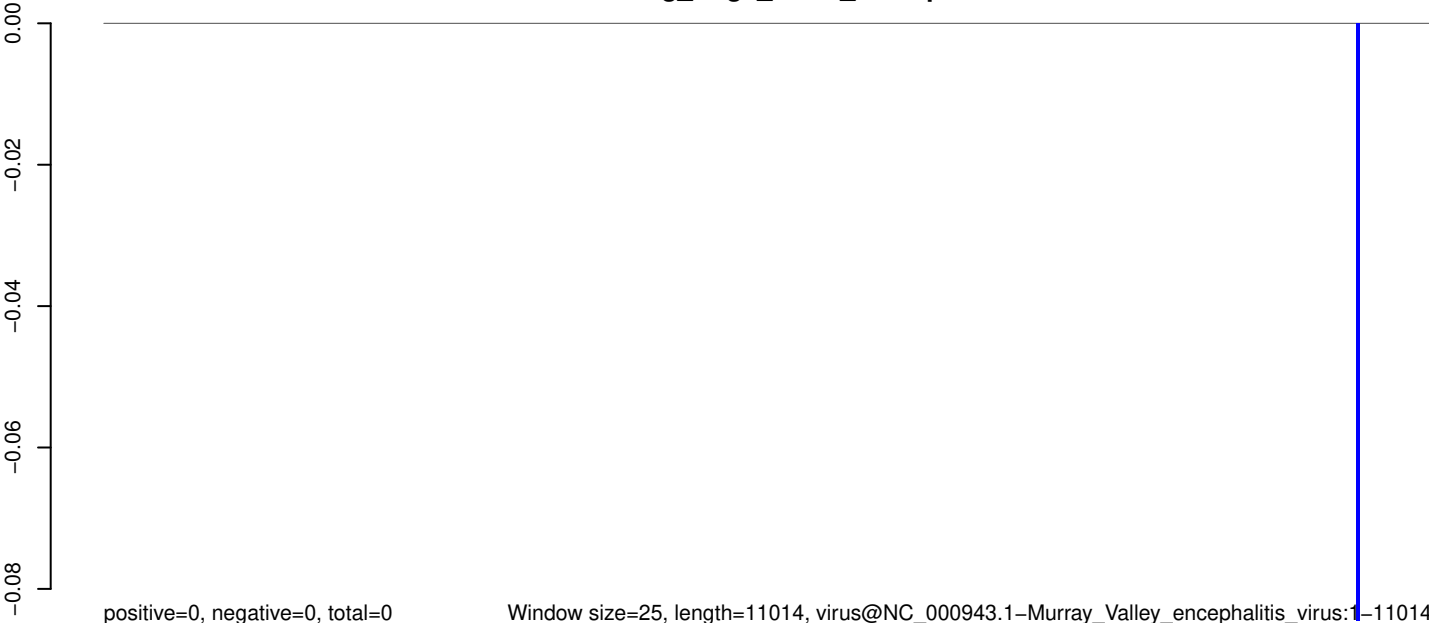
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AeAeg_Aag2_BTV1_AK.24_35.rep



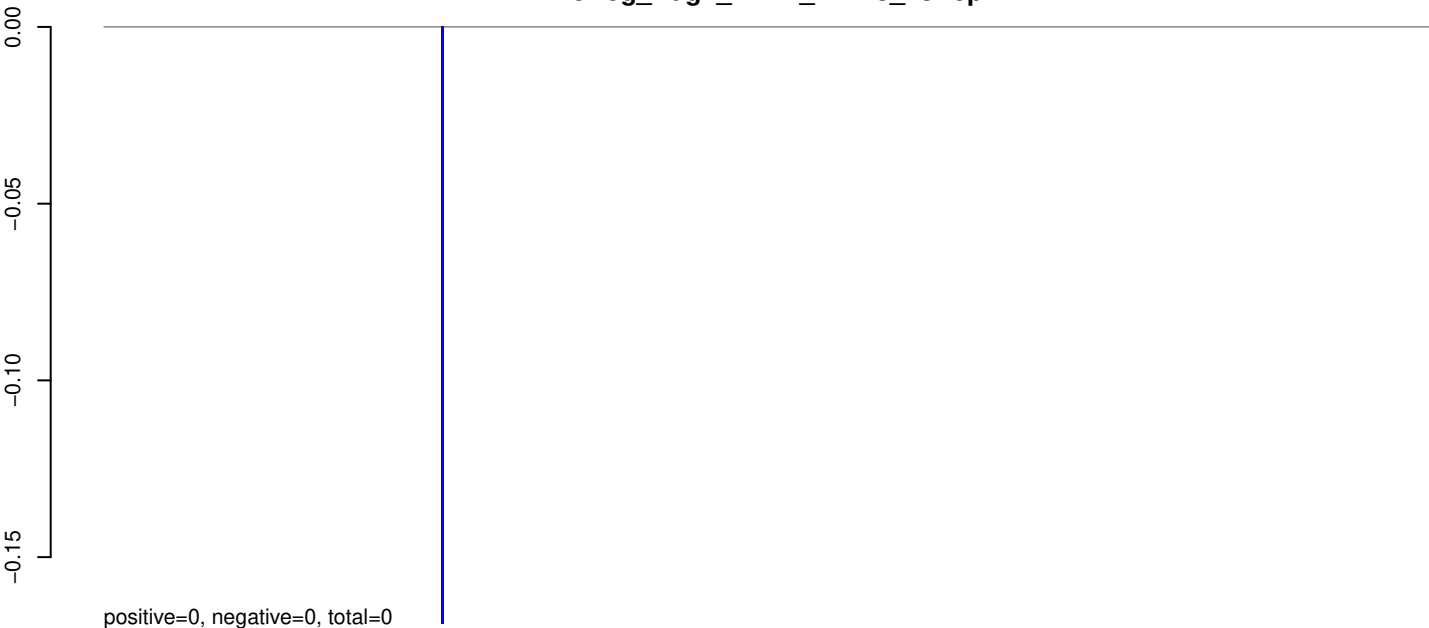
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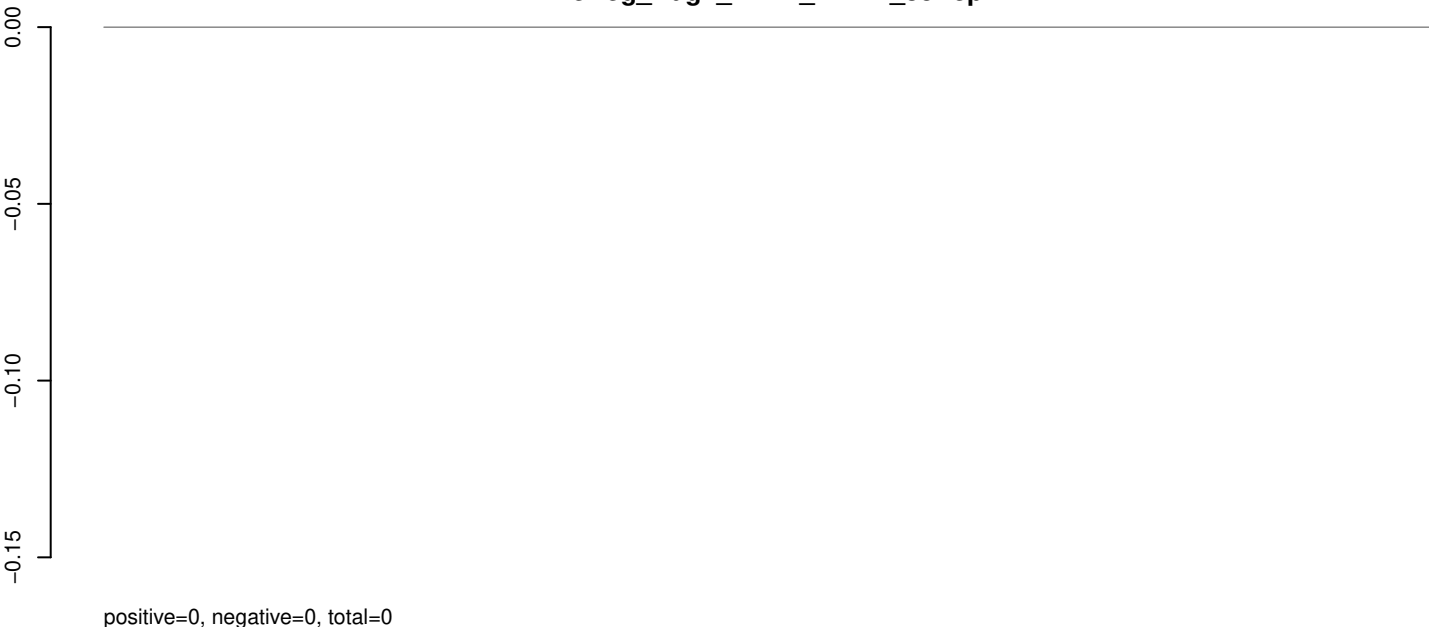
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0 2000 4000 6000 8000 10000

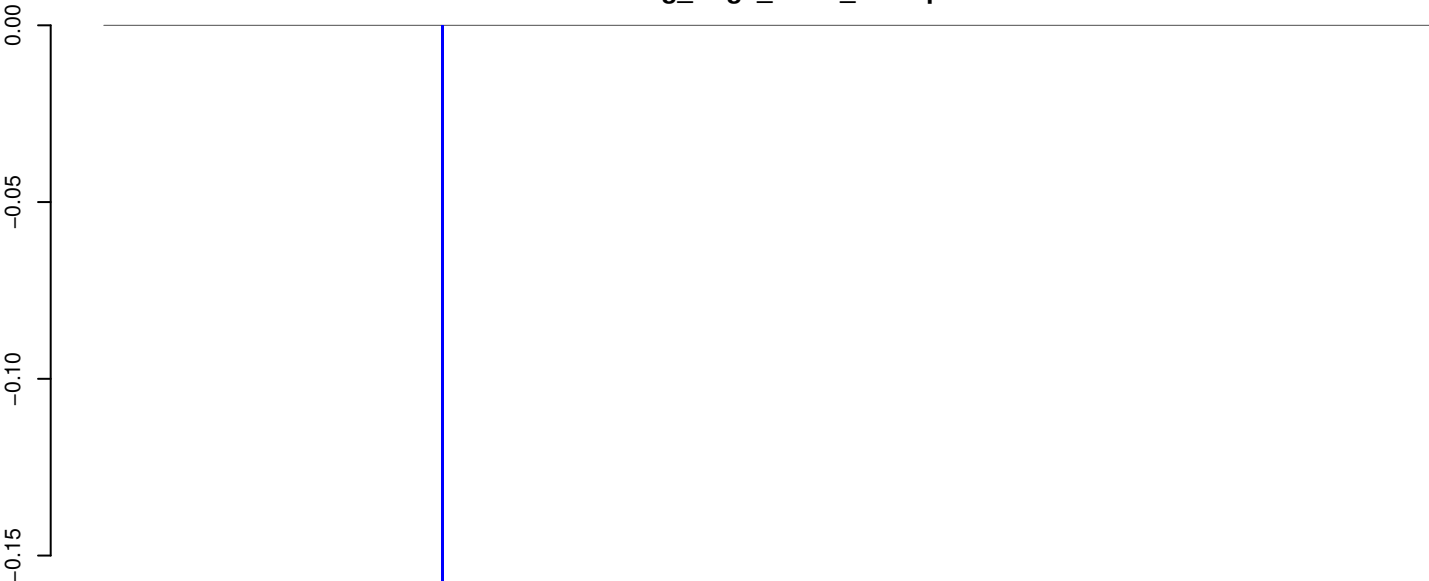
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AeAeg_Aag2_BTV1_AK.24_35.rep



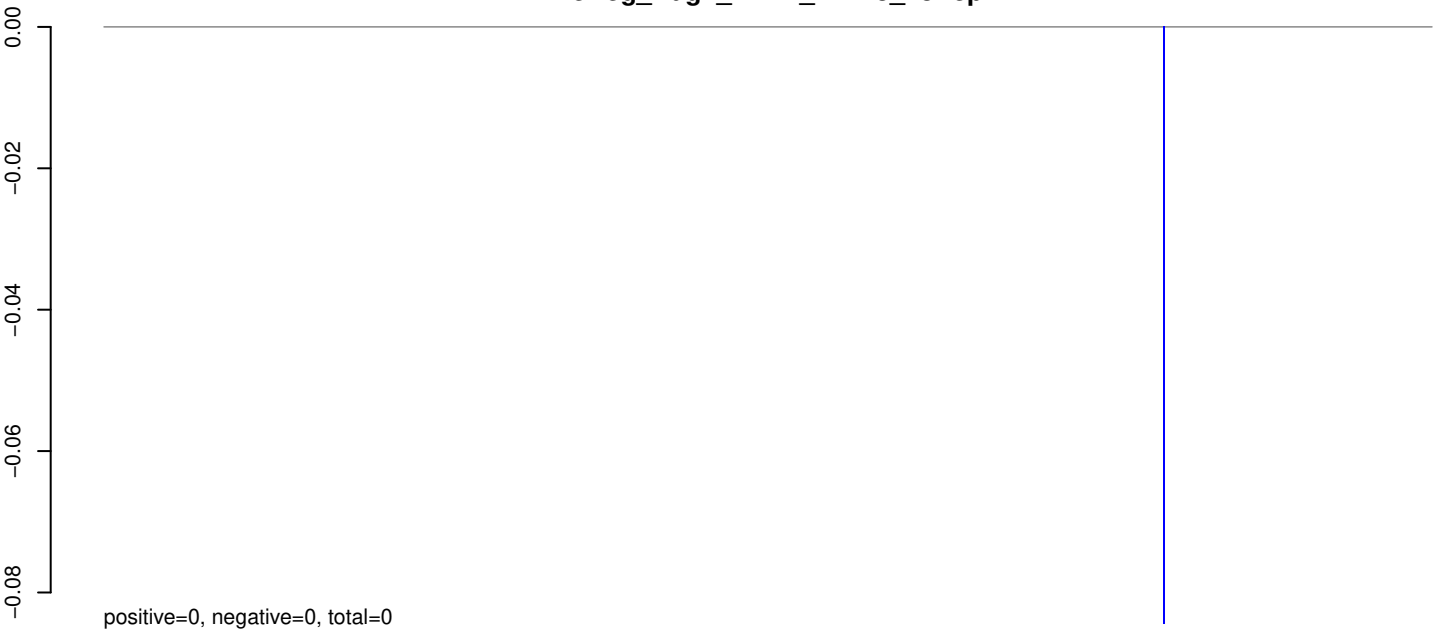
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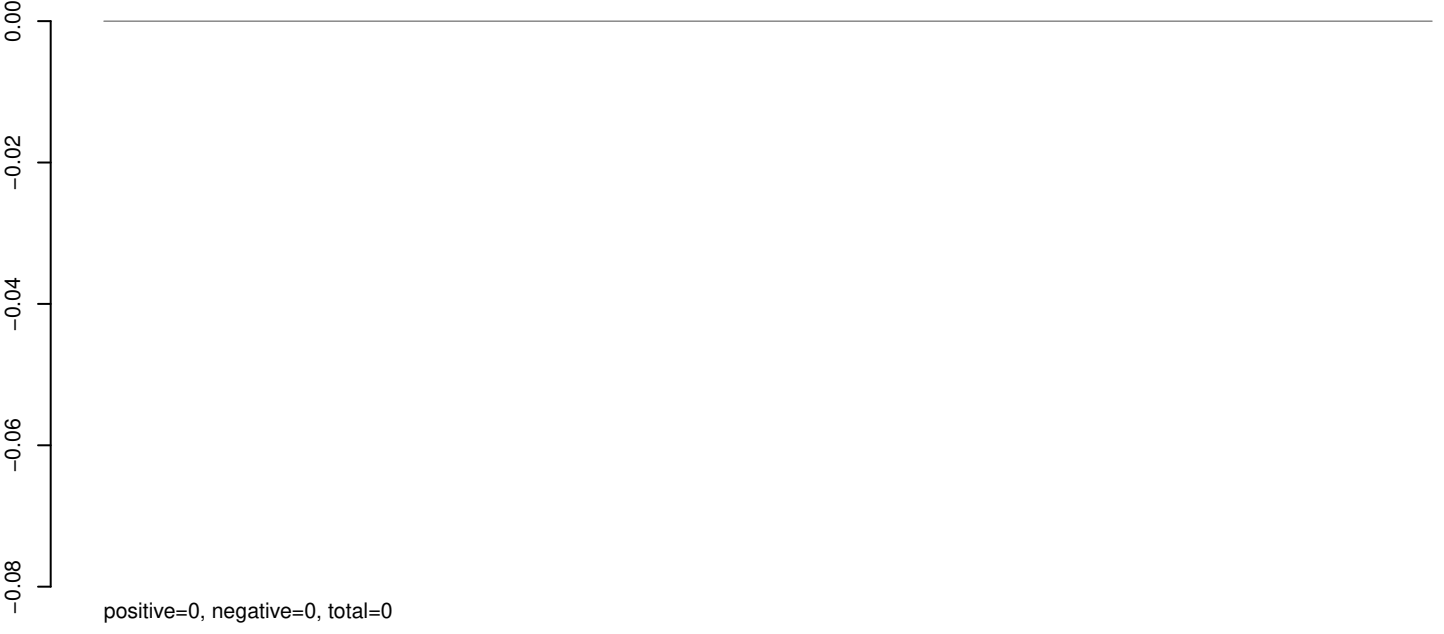
Window size=25, length=12195, virus@MG747536.1_Oropouche_virus_isolate_BeAr_136921_segment_S_M_L_:1-12195

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep

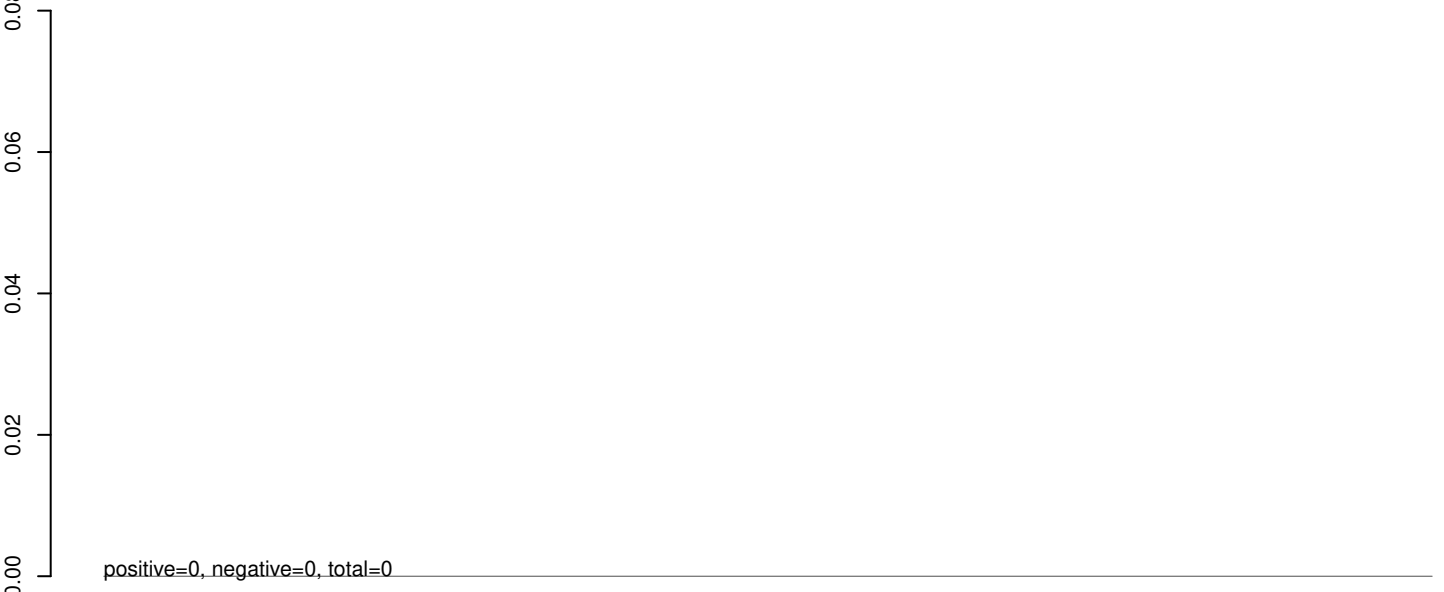


0 5000 10000 15000 20000

AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep

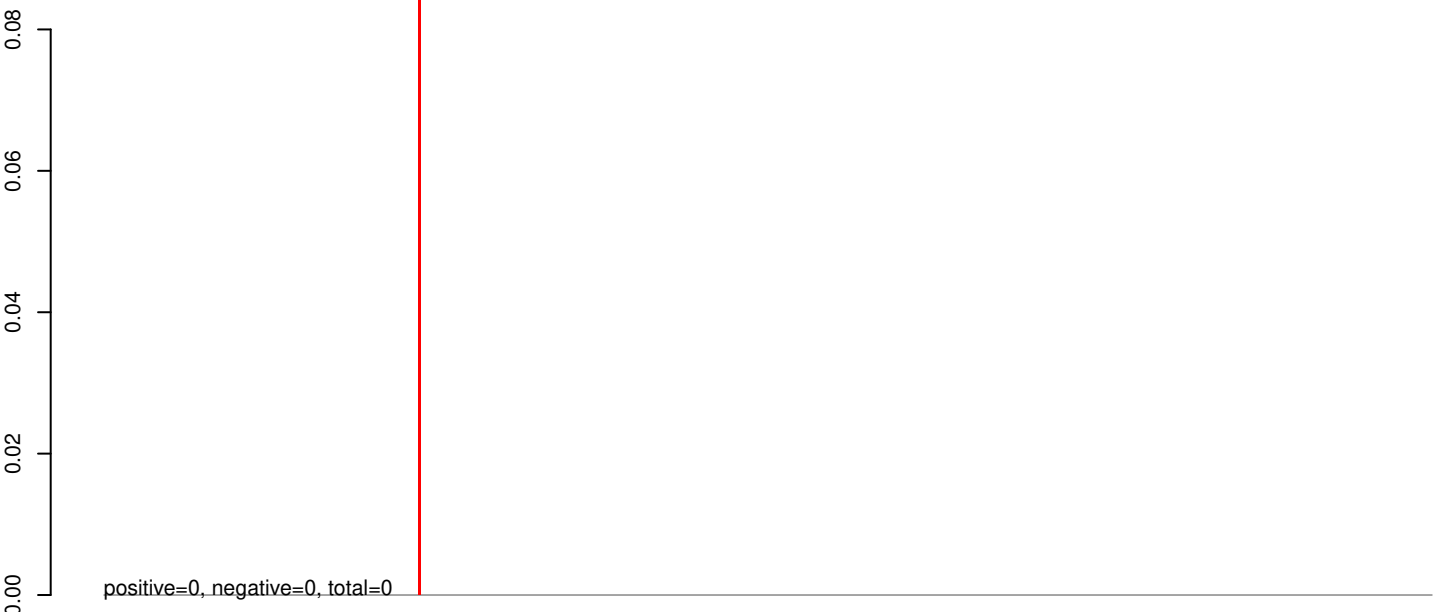


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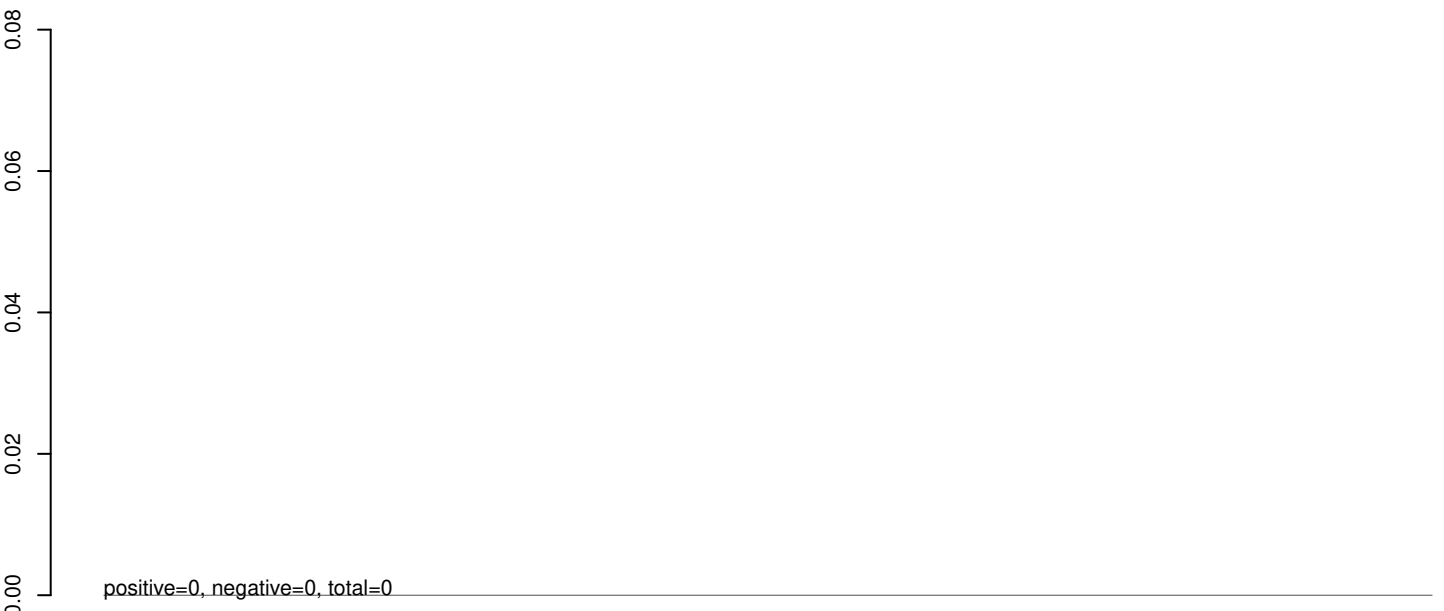


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virus@MF176267.1_Culex phasma-like virus_strain_mos172X35099_segment_S_M_L:1-10814

AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



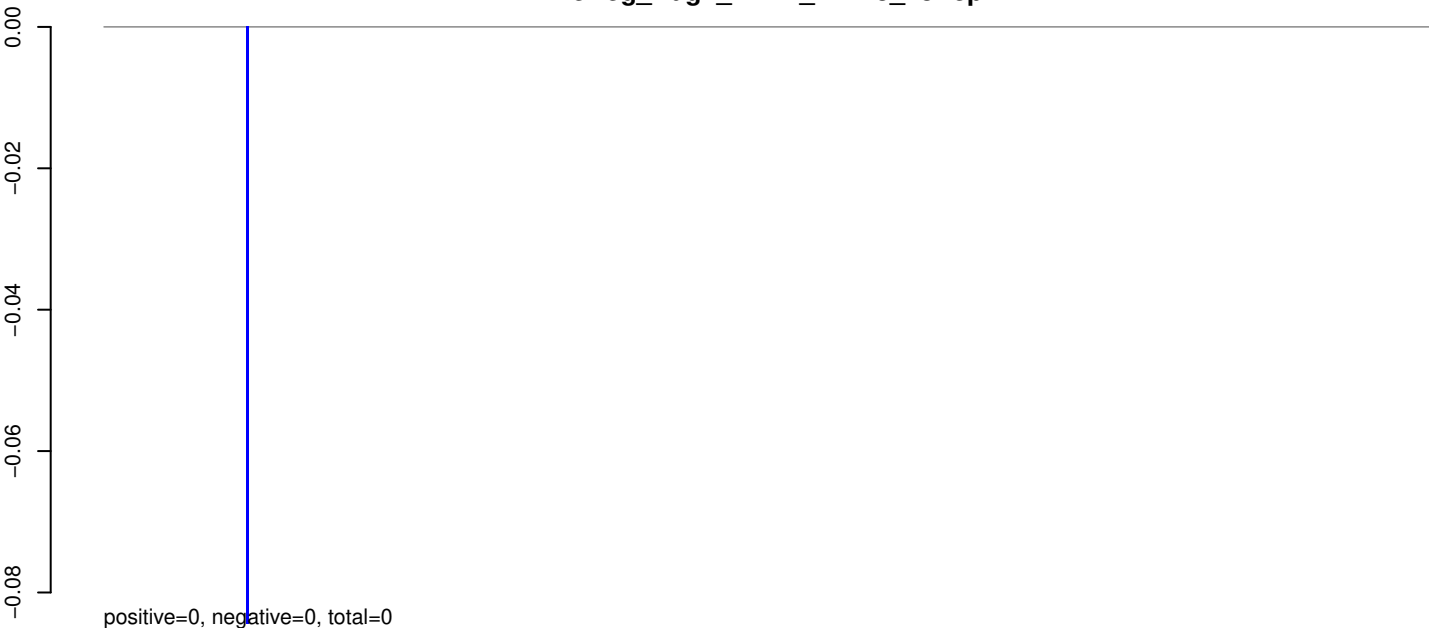
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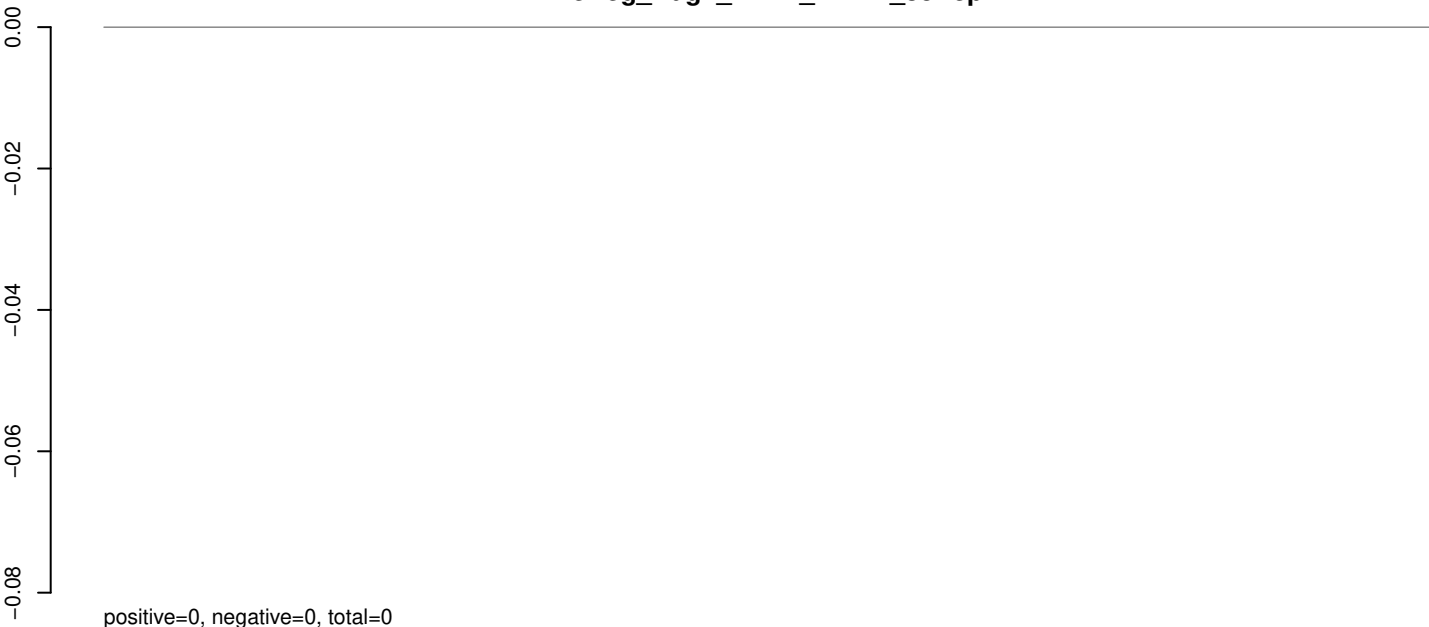
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0 2000 4000 6000 8000 10000 12000

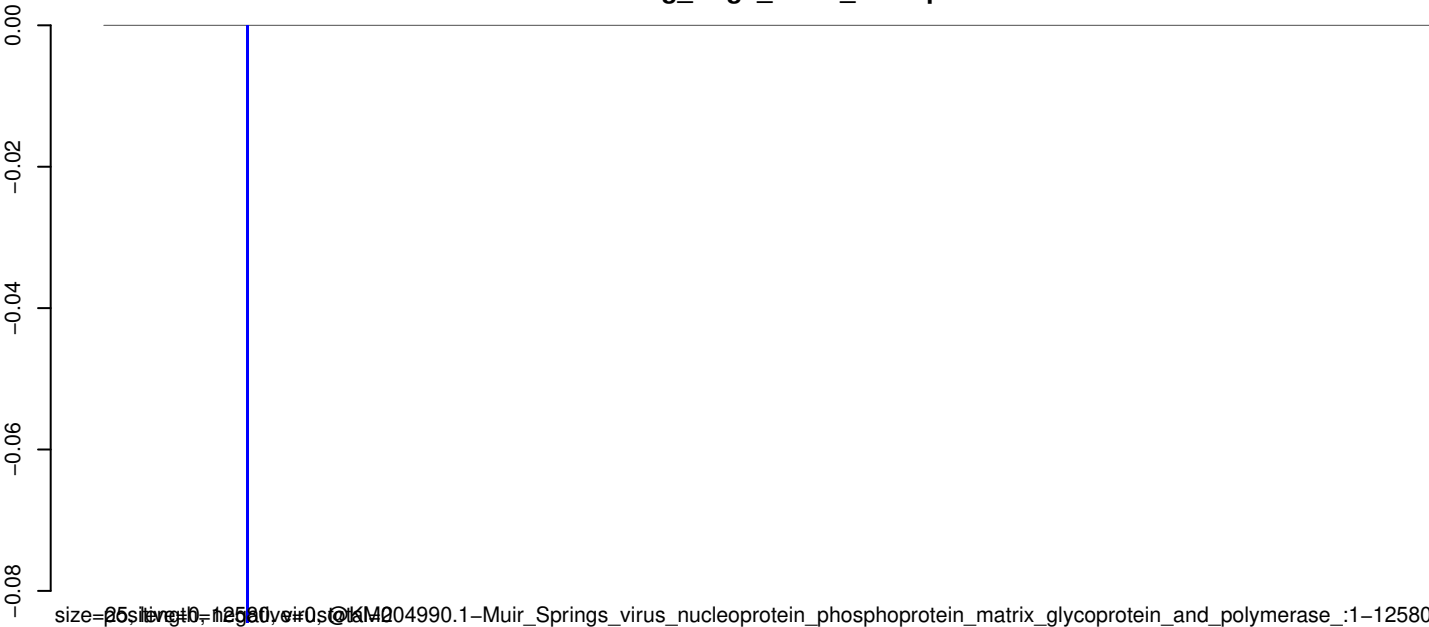
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AeAeg_Aag2_BTV1_AK.24_35.rep

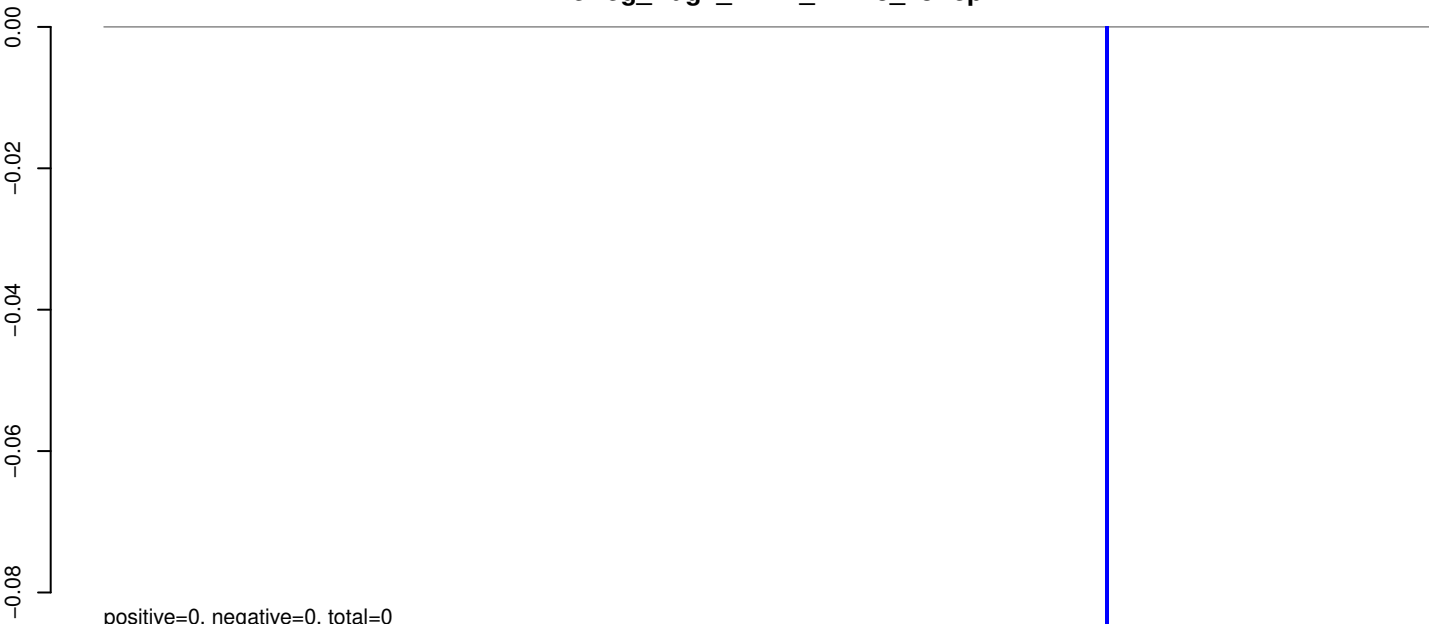


AeAeg_Aag2_BTV1_AK.rep

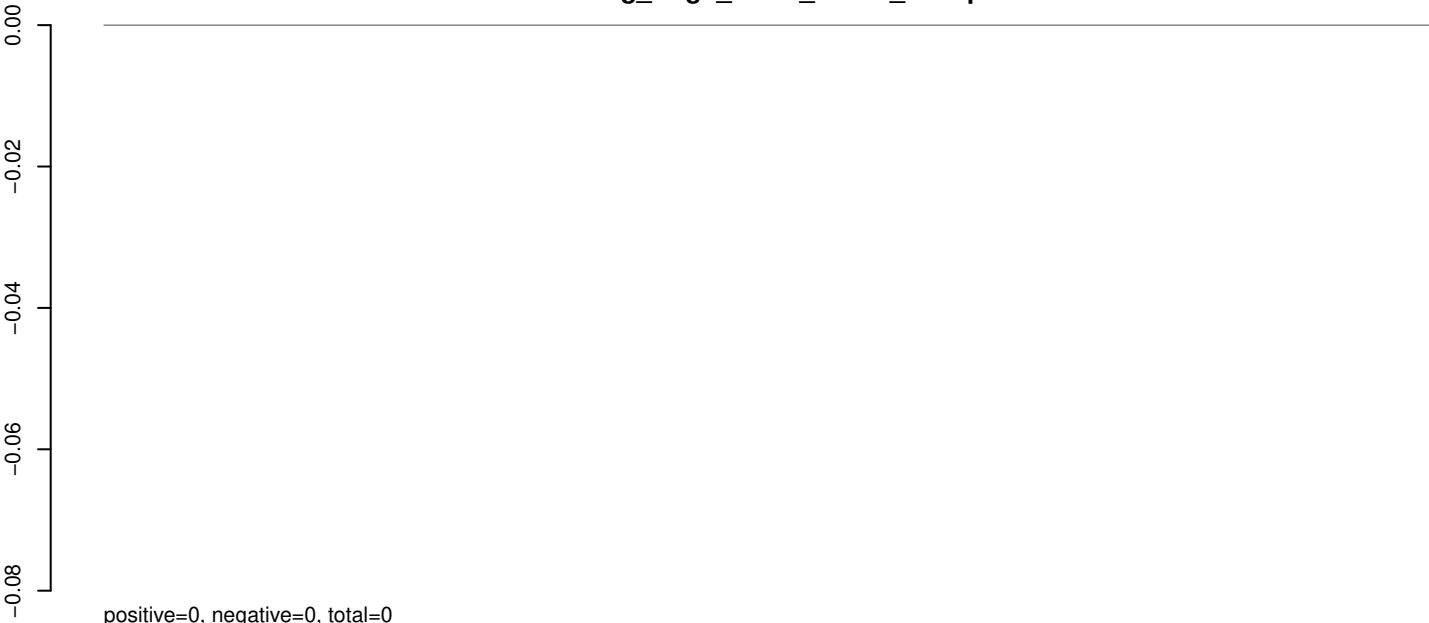


0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_BTV1_AK.18_23.rep



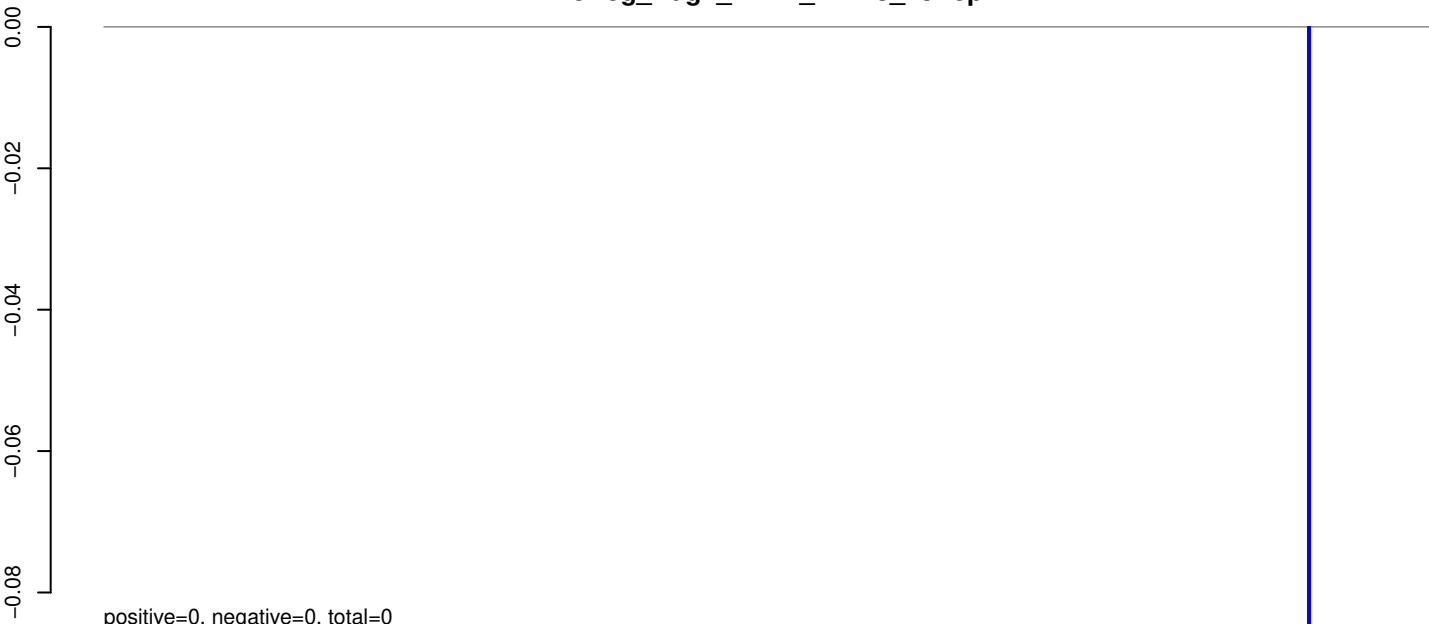
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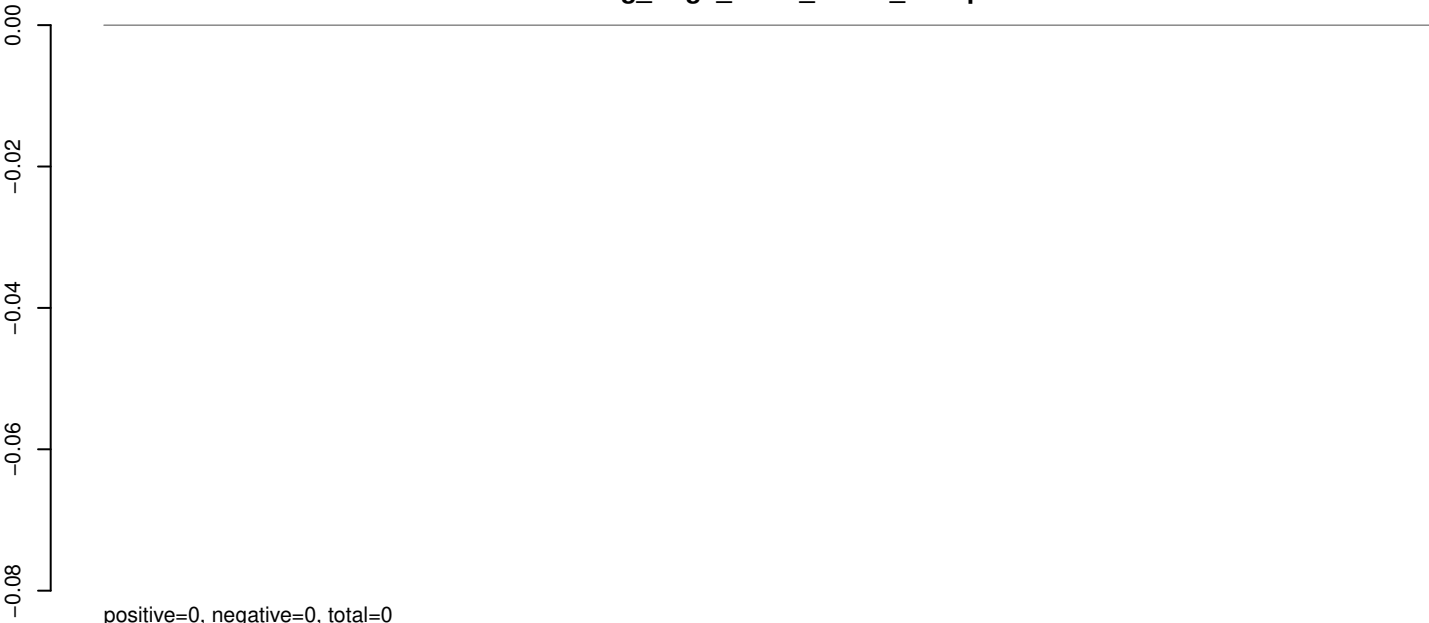
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AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



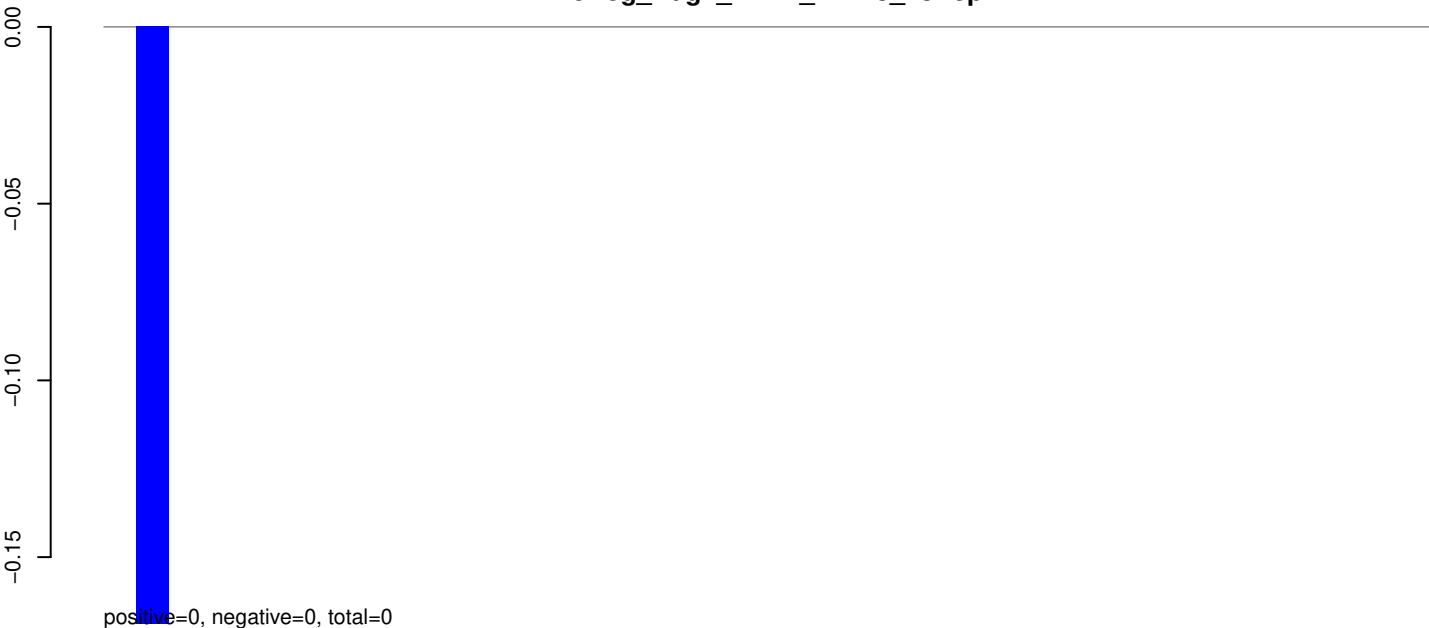
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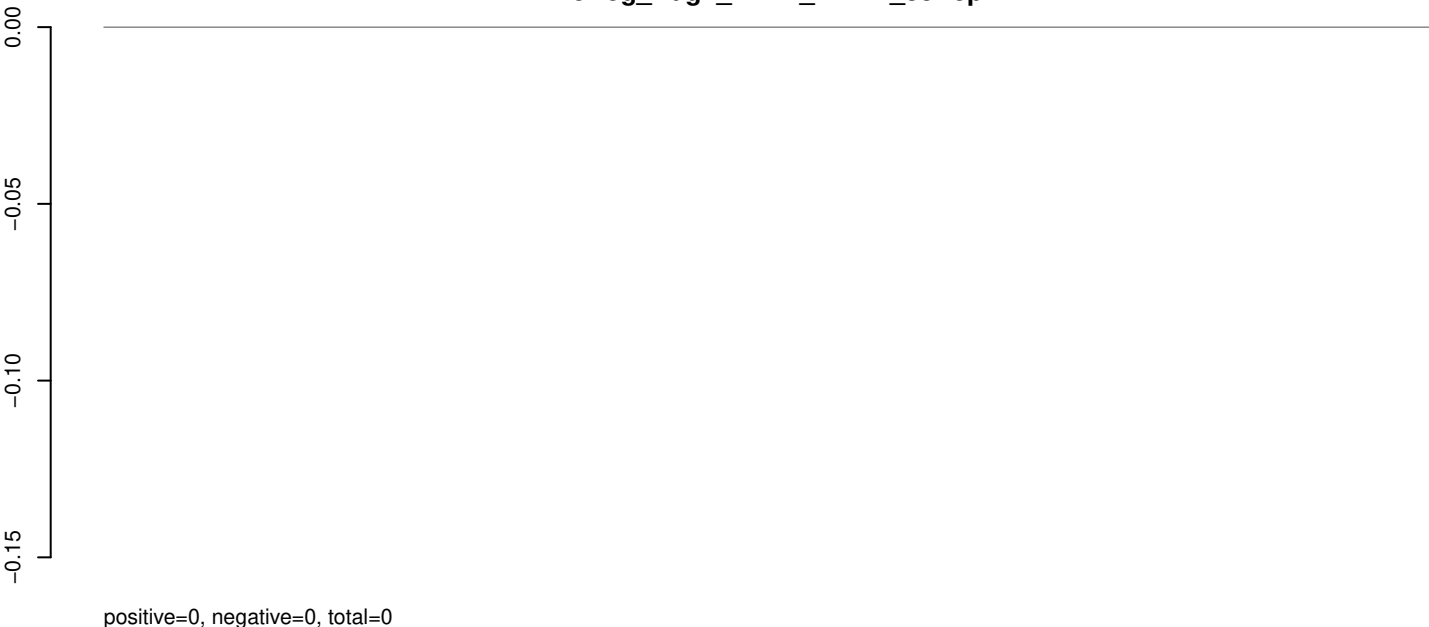
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0 2000 4000 6000 8000 10000

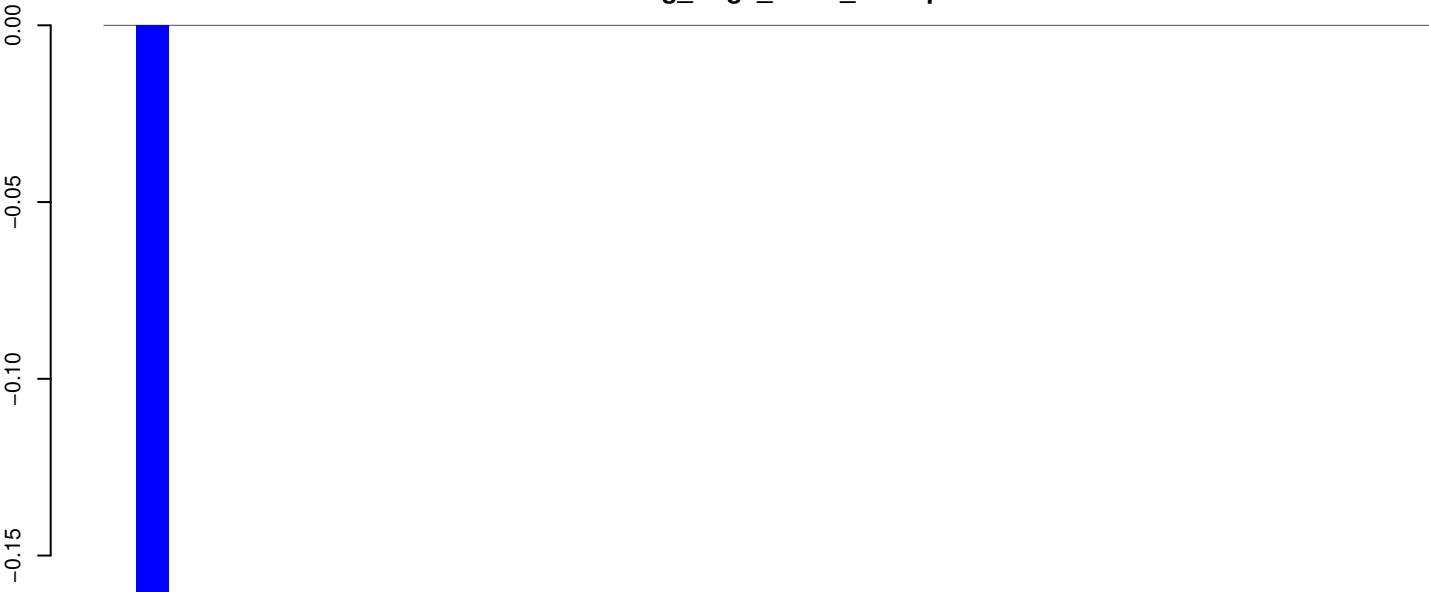
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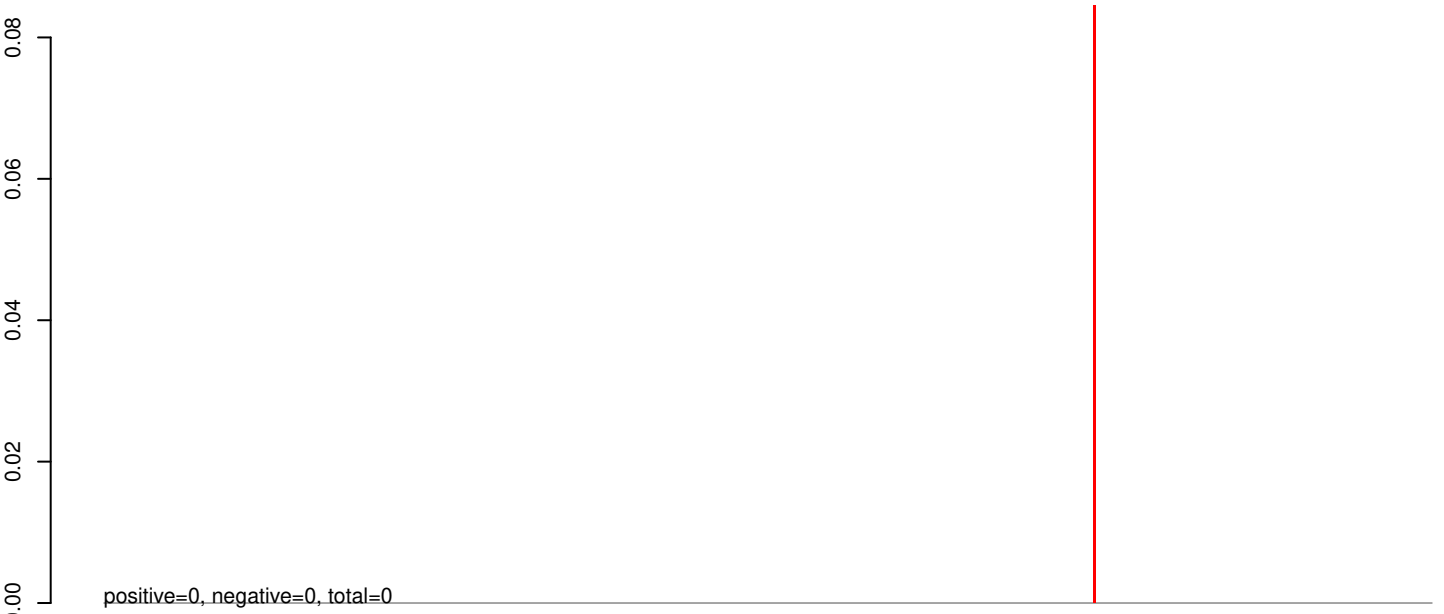
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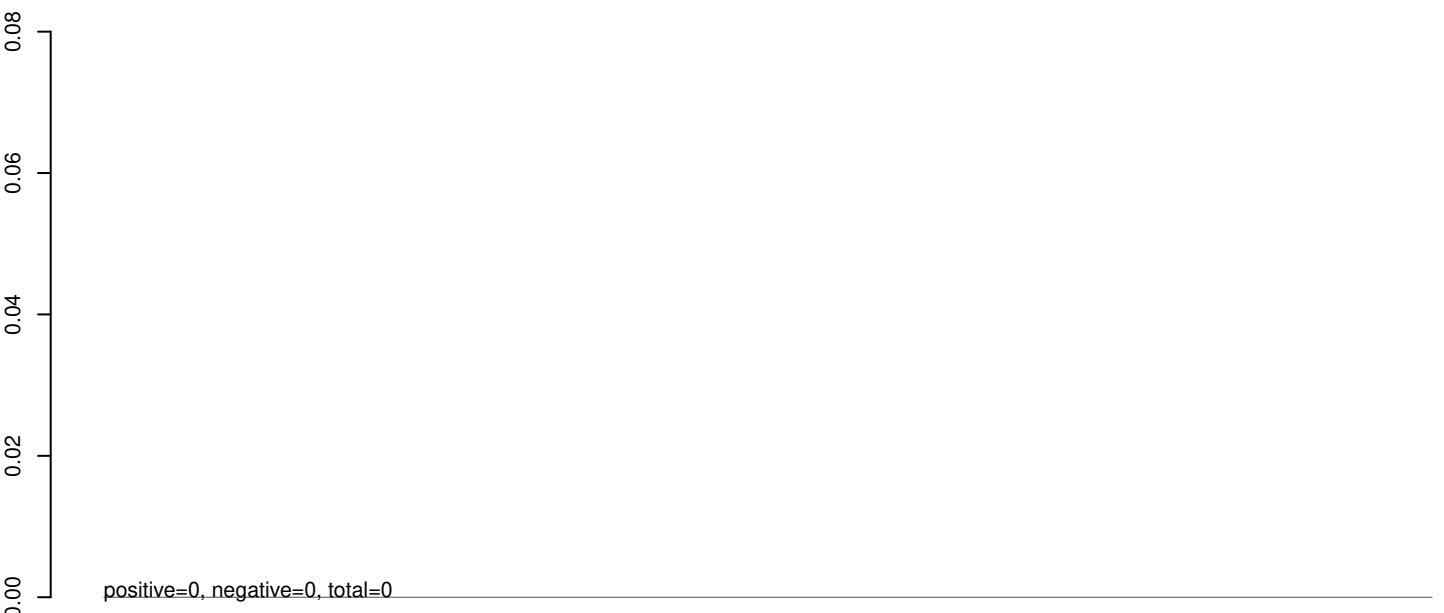
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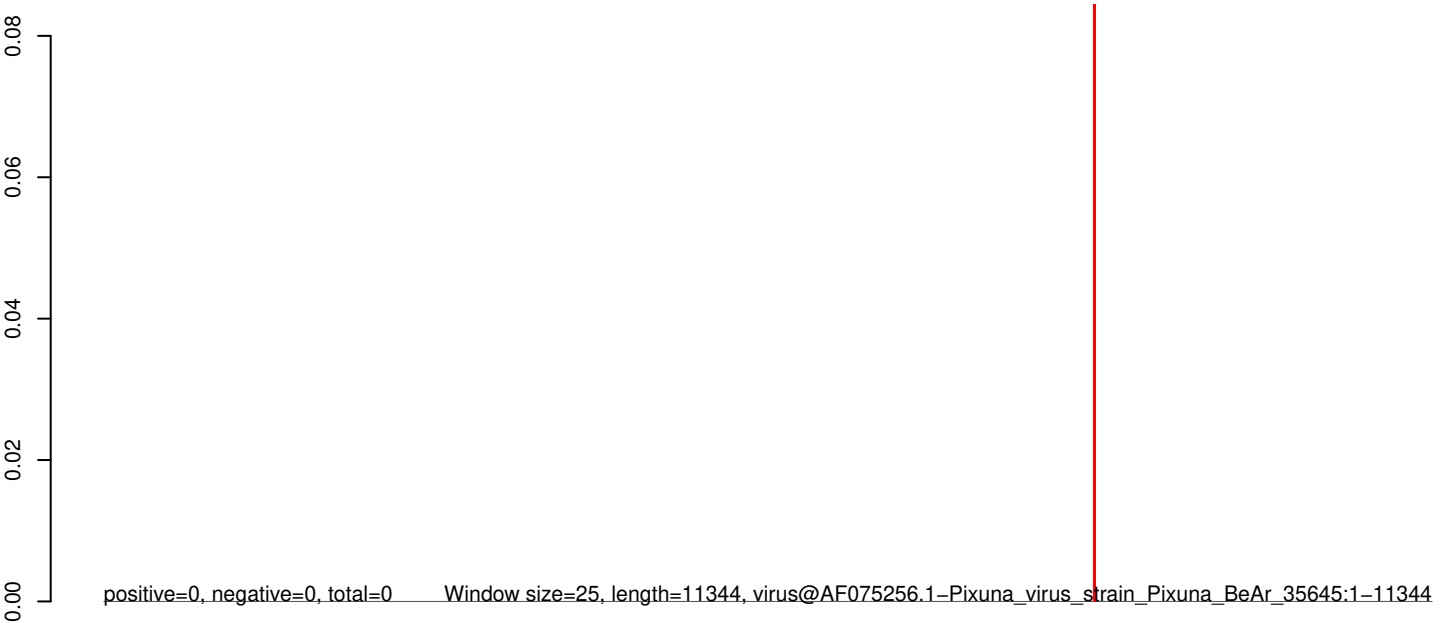
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AeAeg_Aag2_BTV1_AK.24_35.rep



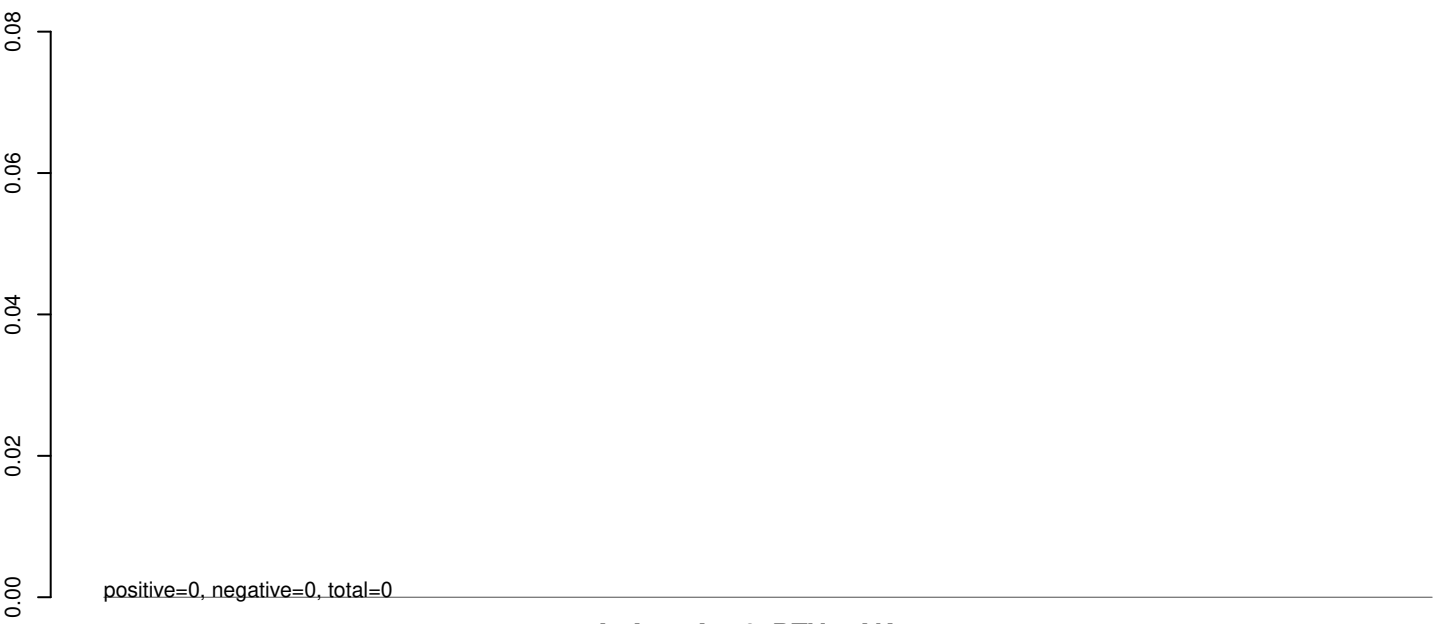
AeAeg_Aag2_BTV1_AK.rep



AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



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

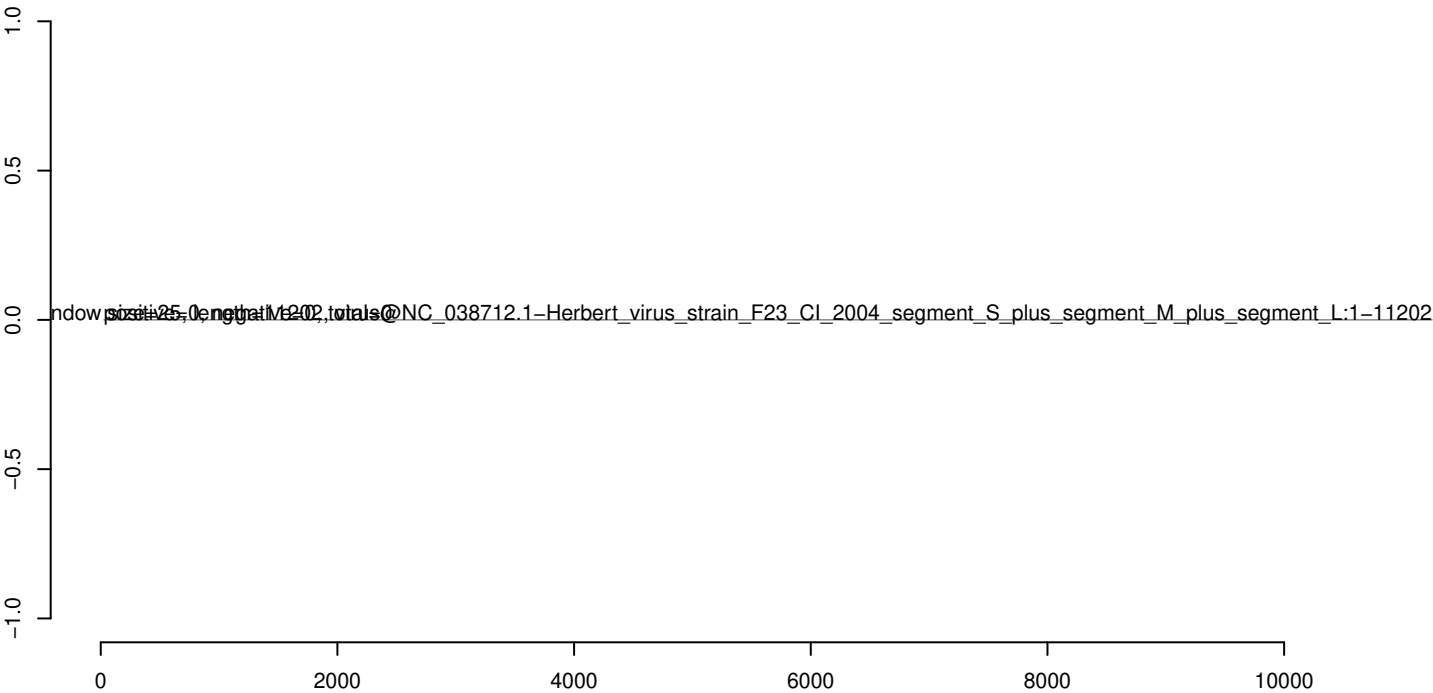
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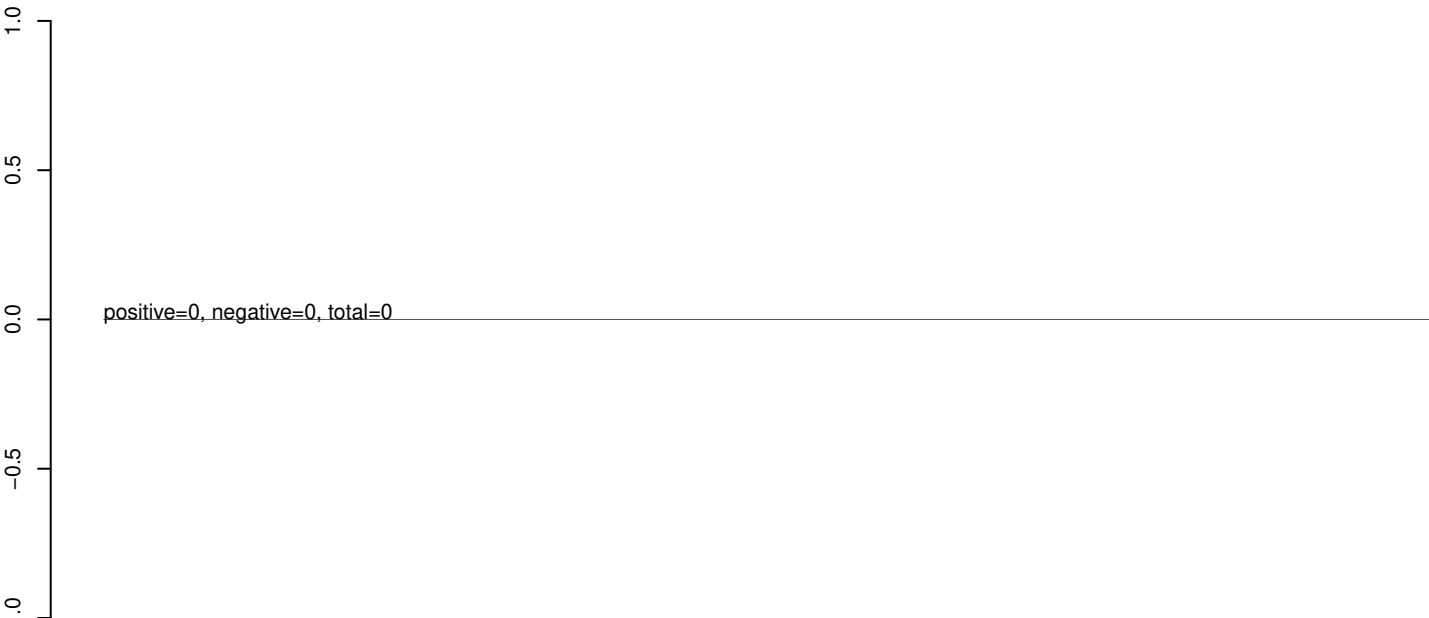
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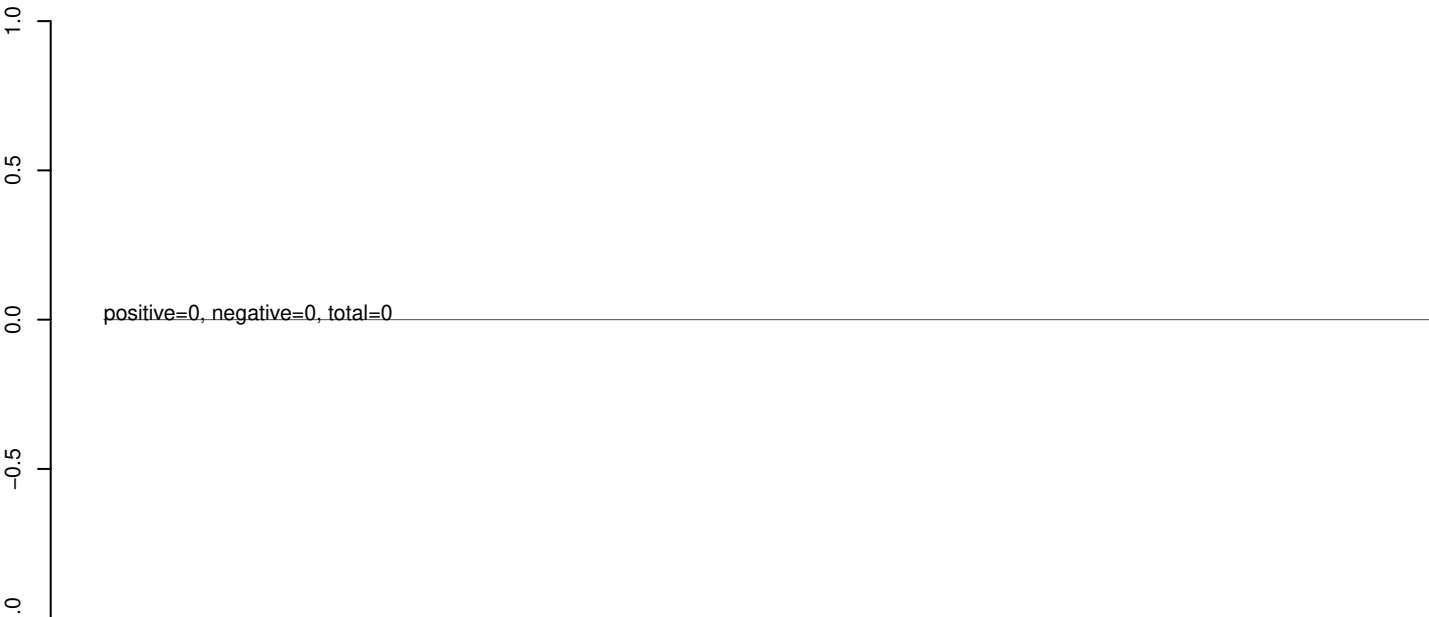
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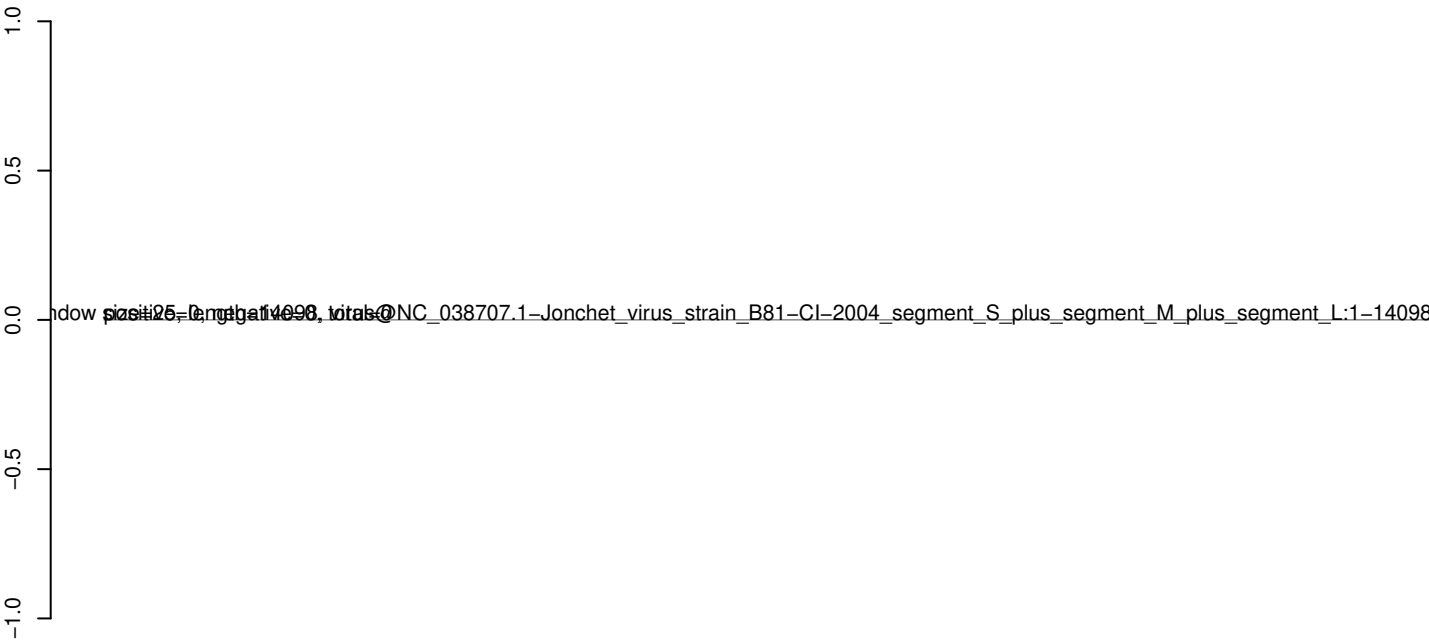
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



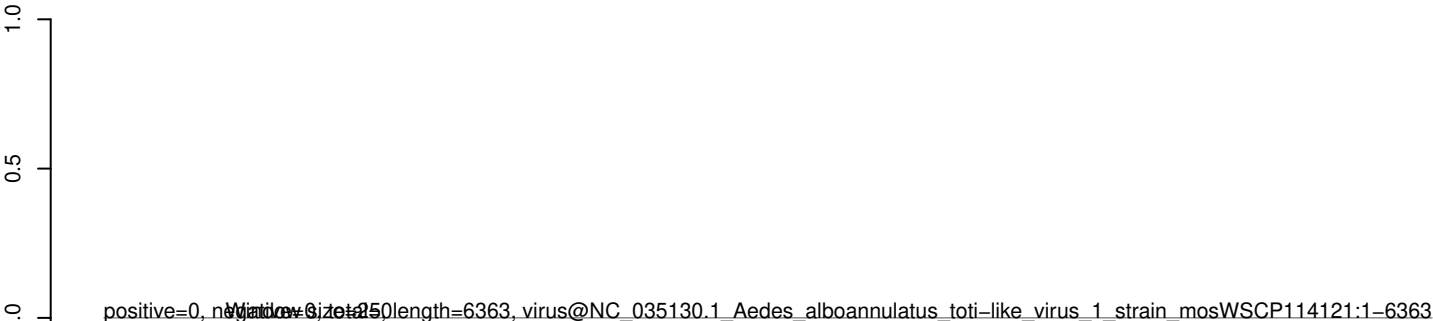
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AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep

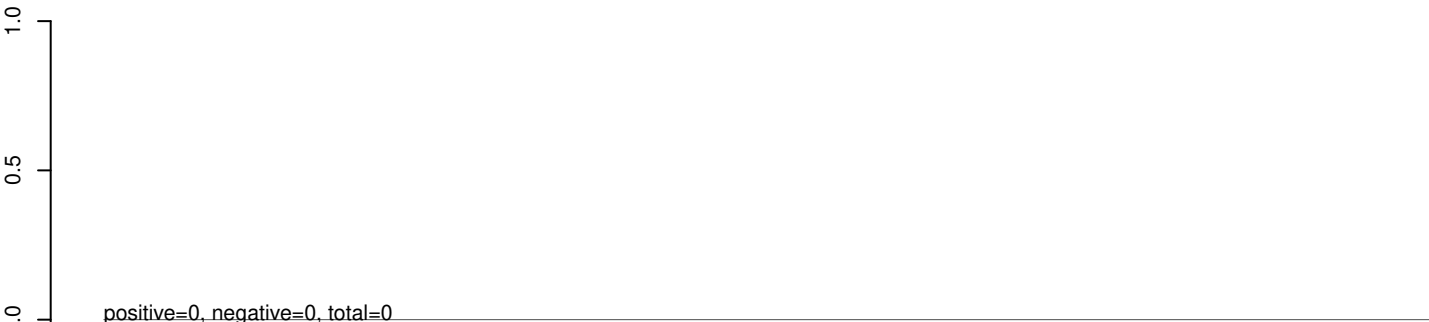


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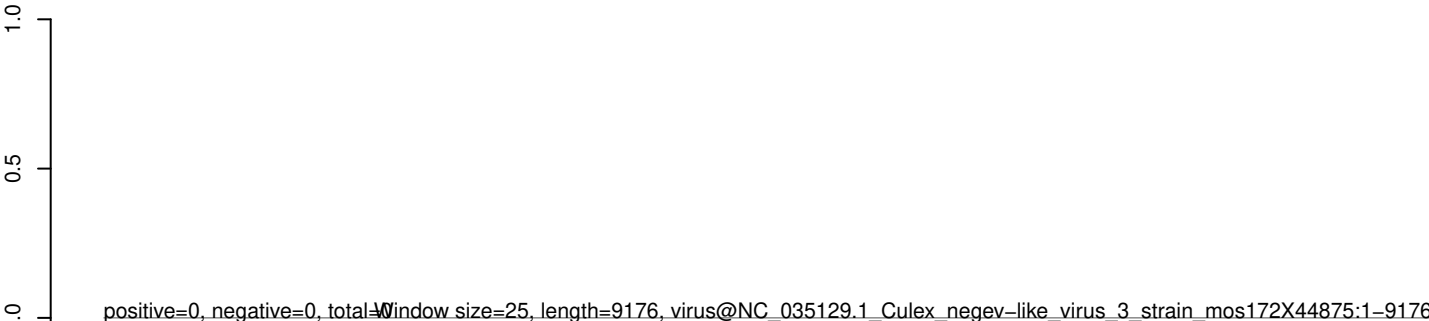
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



0 2000 4000 6000 8000

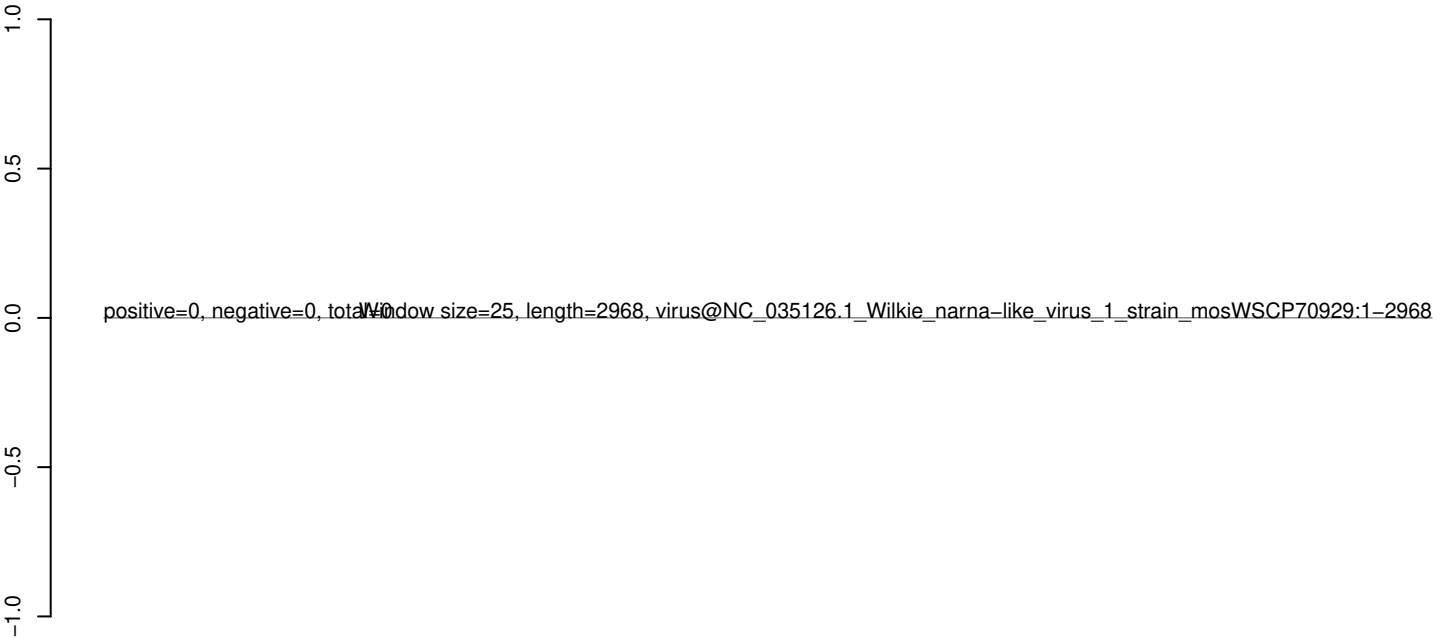
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



0 500 1000 1500 2000 2500 3000

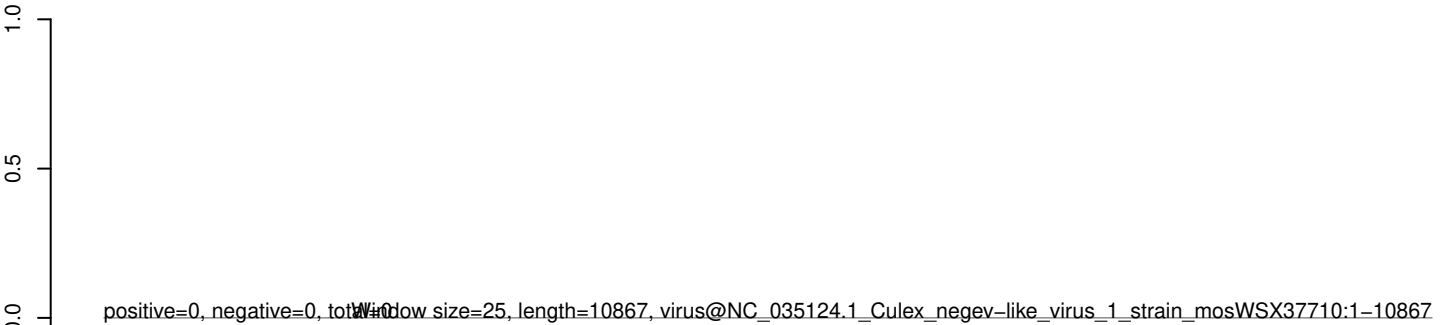
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AeAeg_Aag2_BTV1_AK.24_35.rep

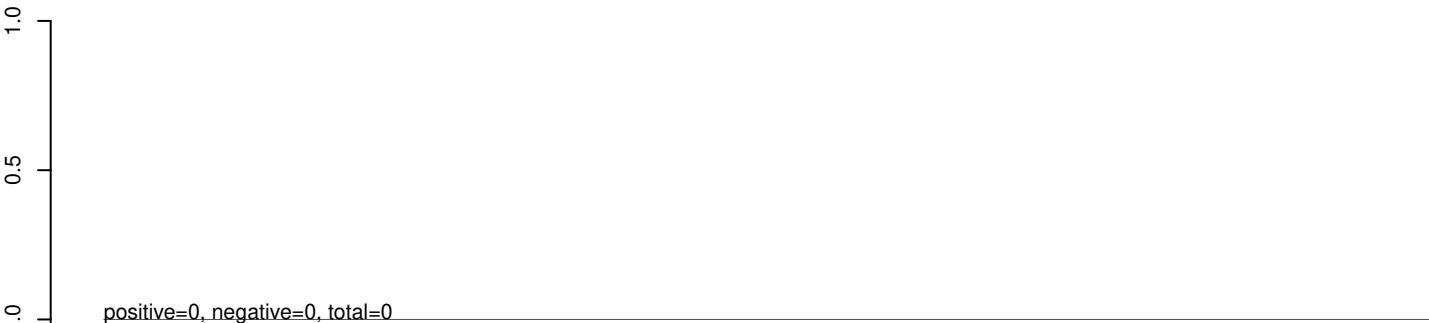


AeAeg_Aag2_BTV1_AK.rep

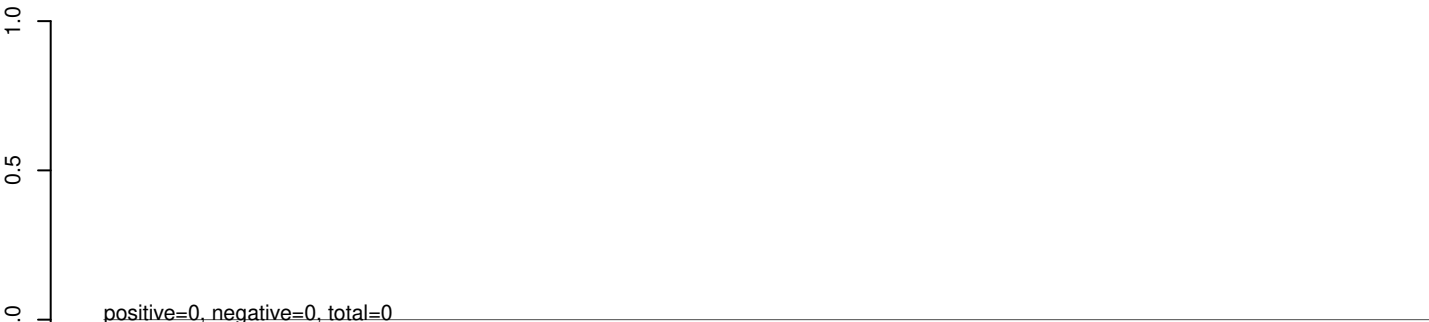


0 2000 4000 6000 8000 10000

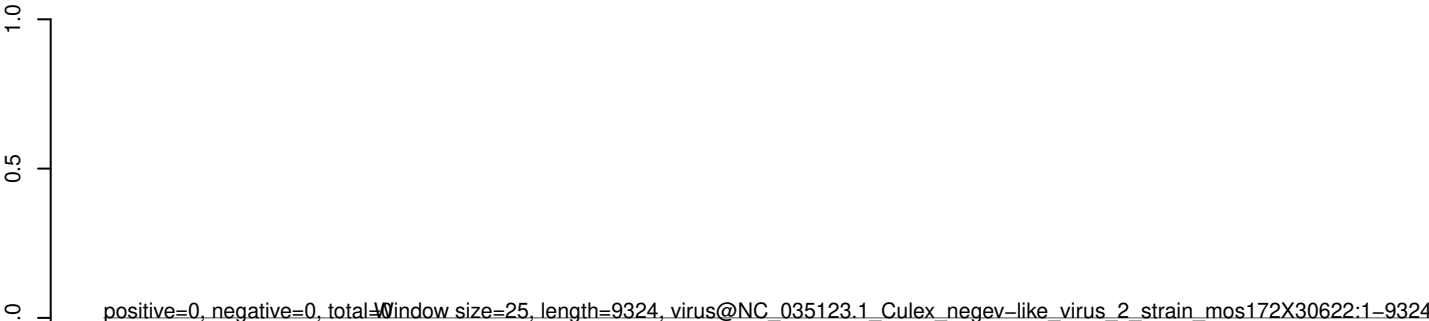
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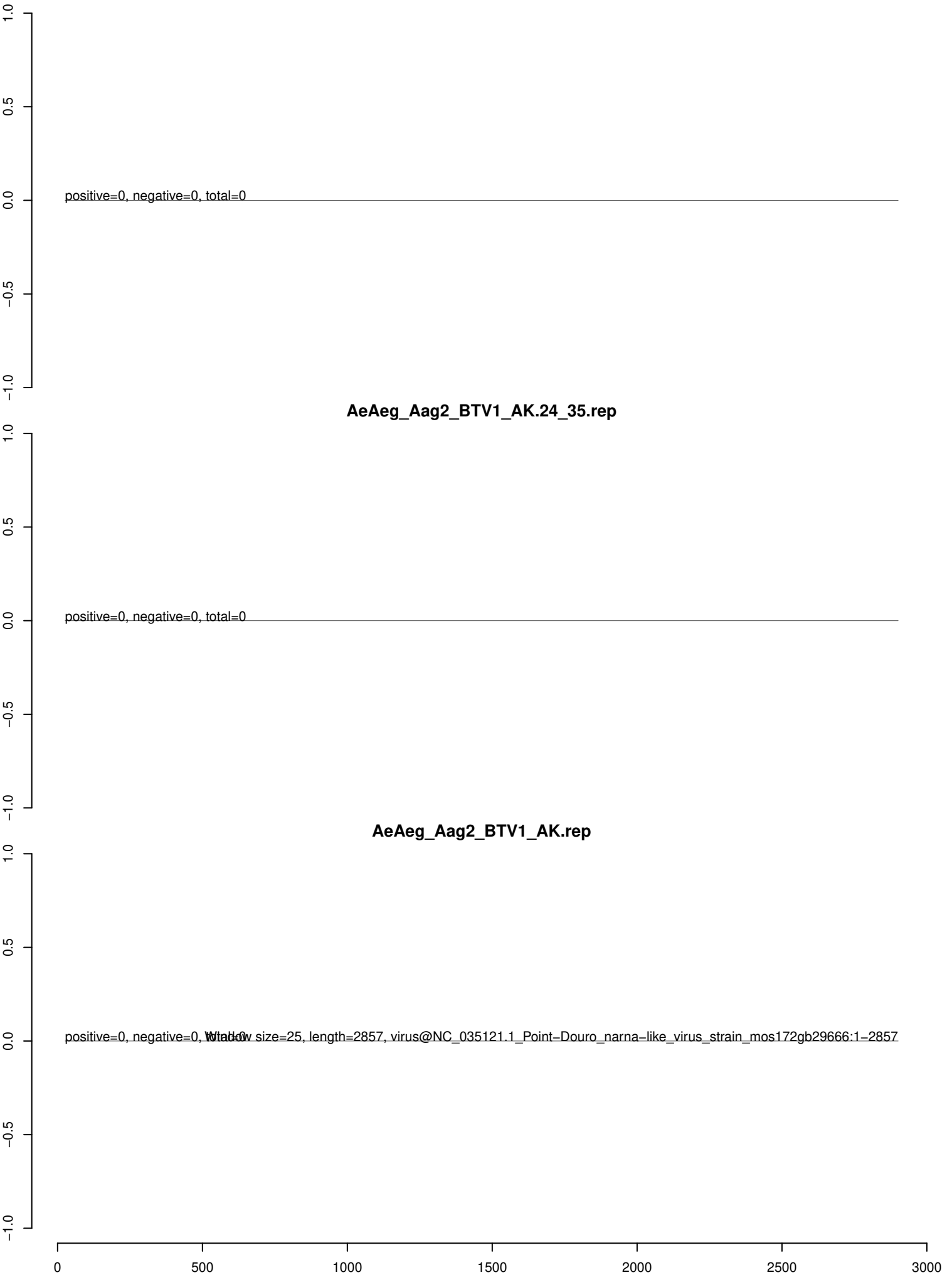


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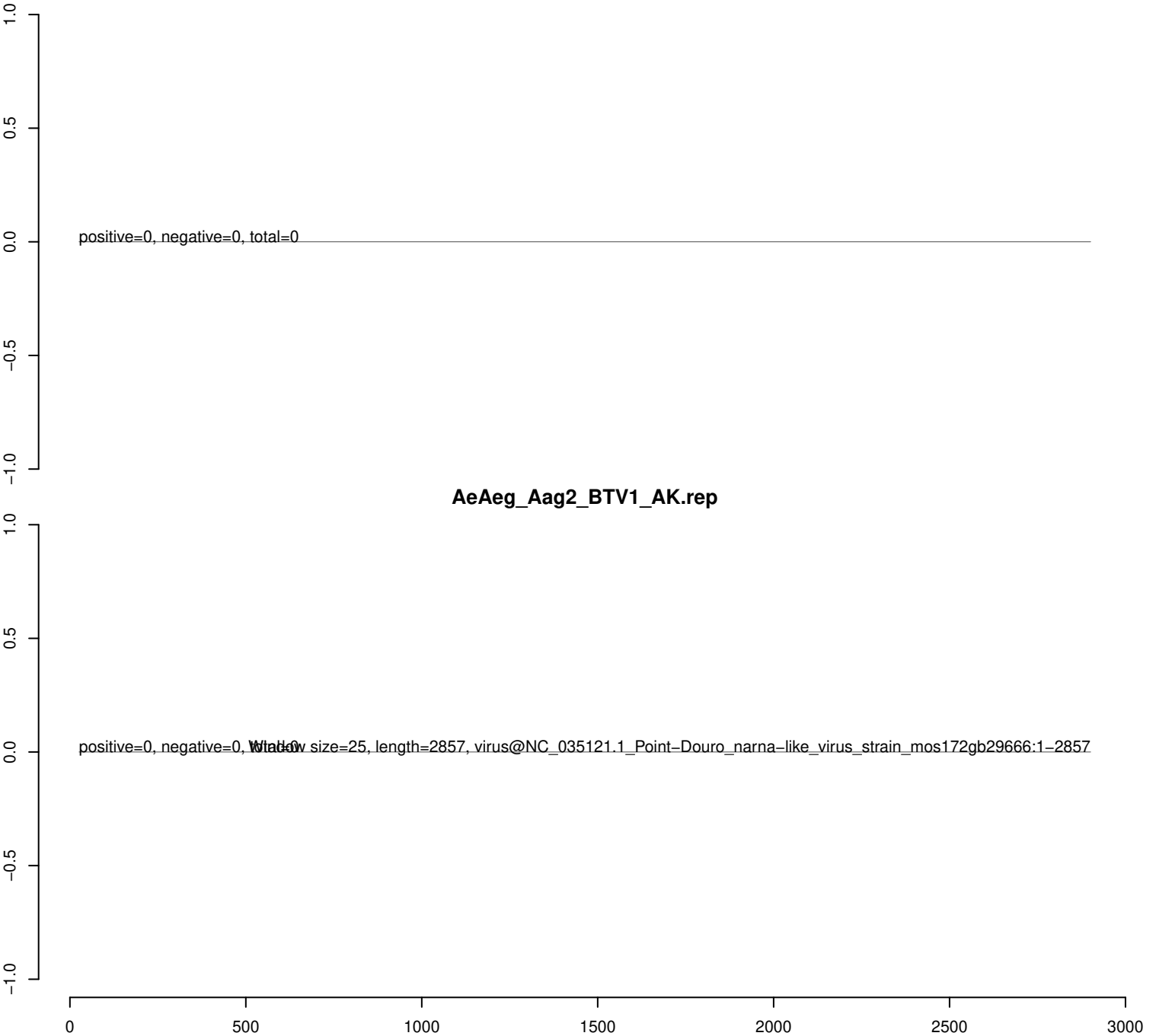


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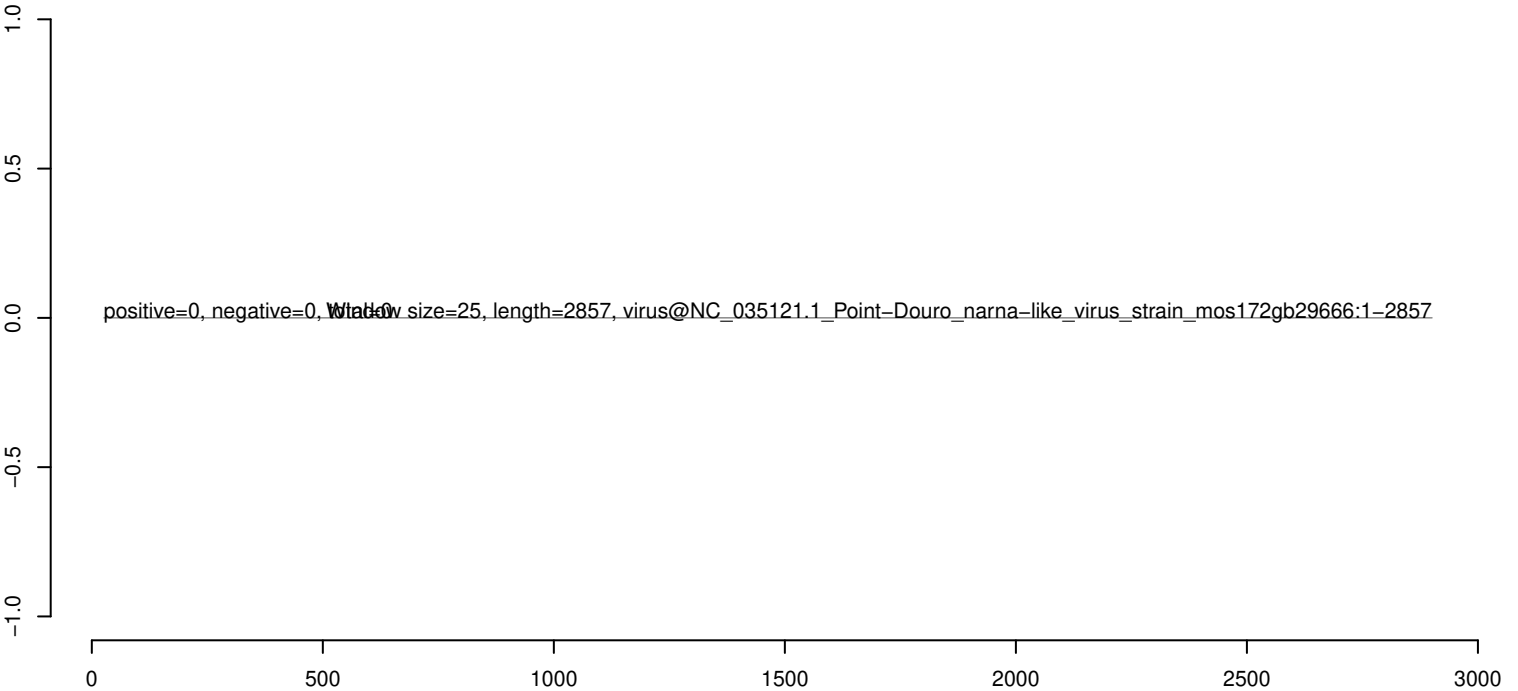
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AeAeg_Aag2_BTV1_AK.24_35.rep



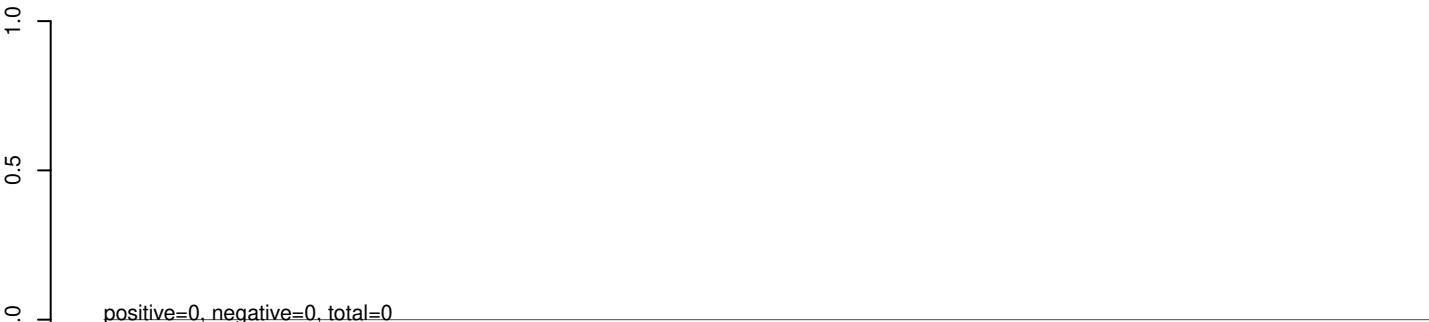
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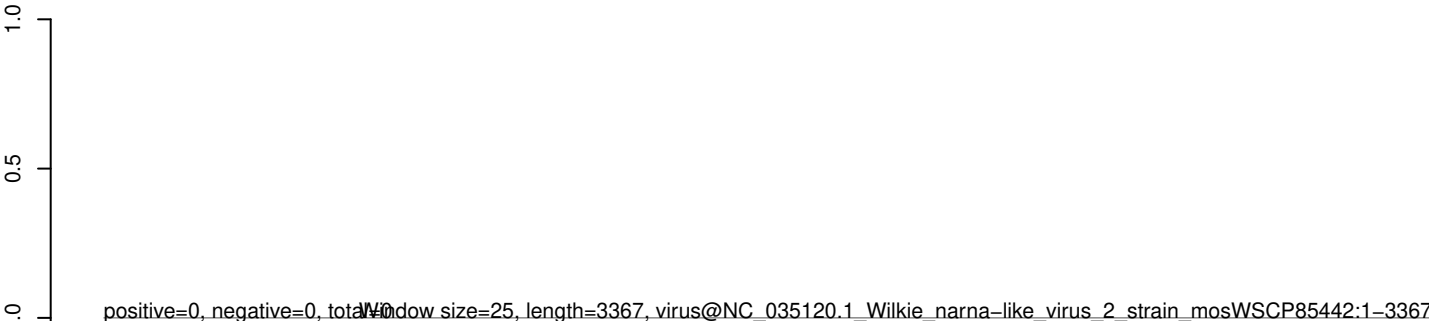
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



0 500 1000 1500 2000 2500 3000 3500

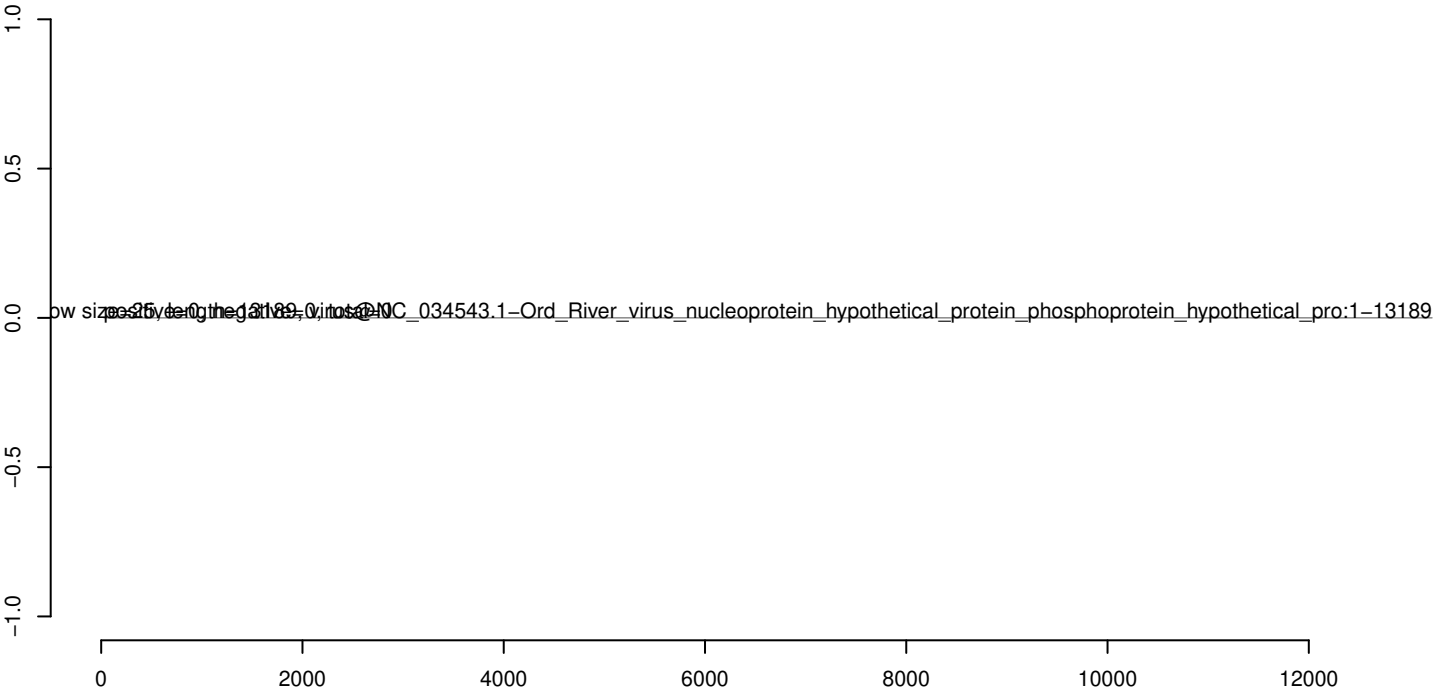
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AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



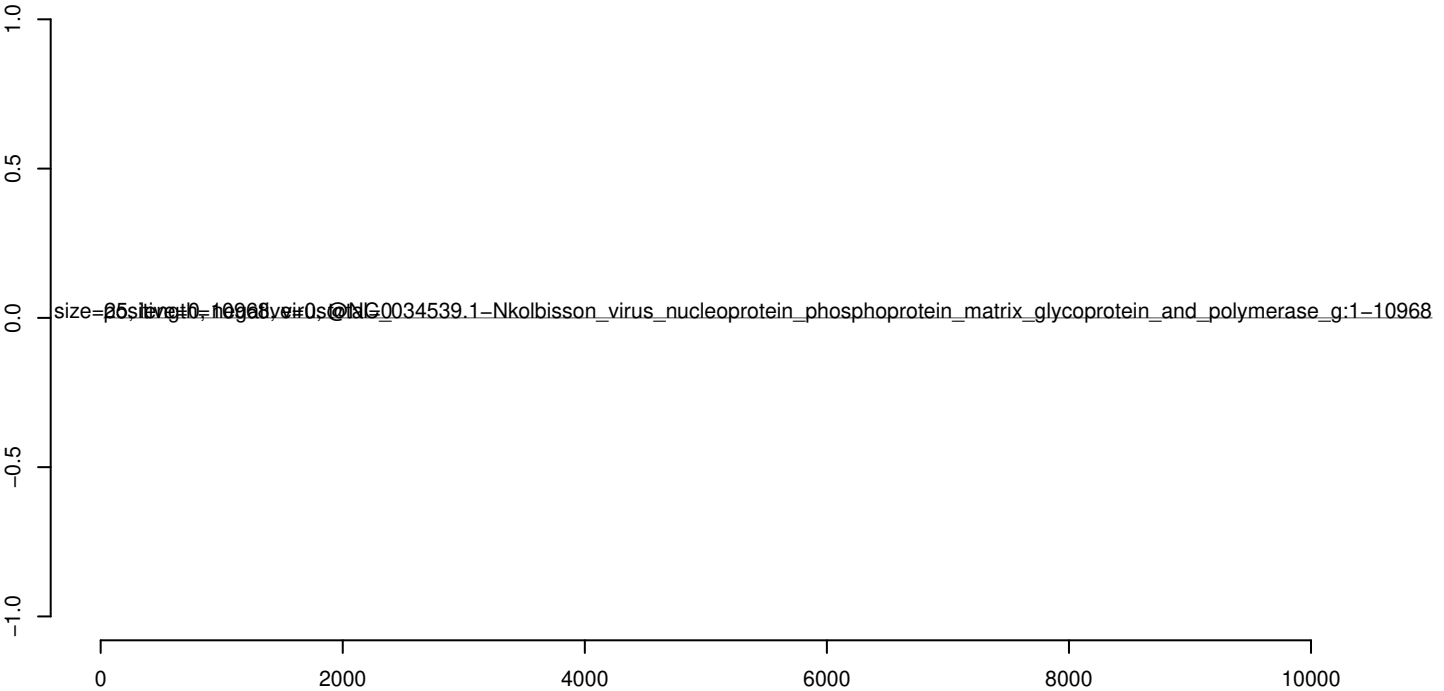
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AeAeg_Aag2_BTV1_AK.24_35.rep



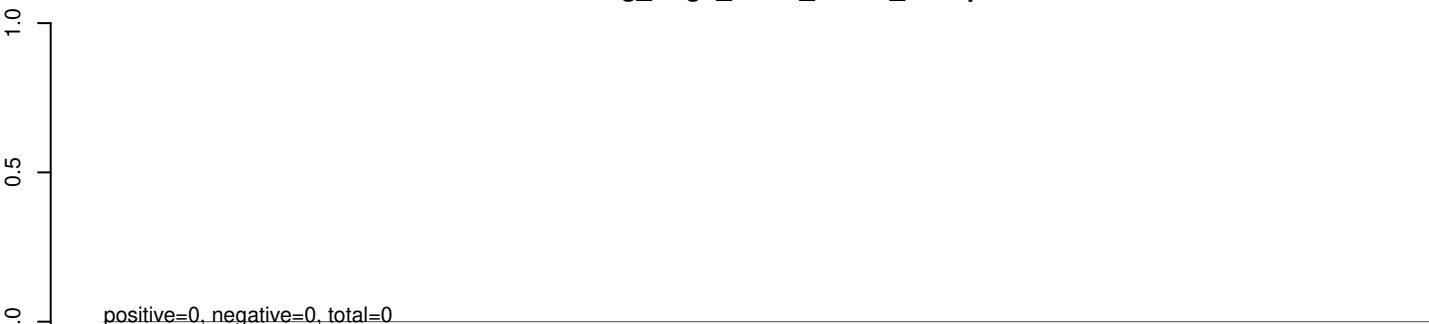
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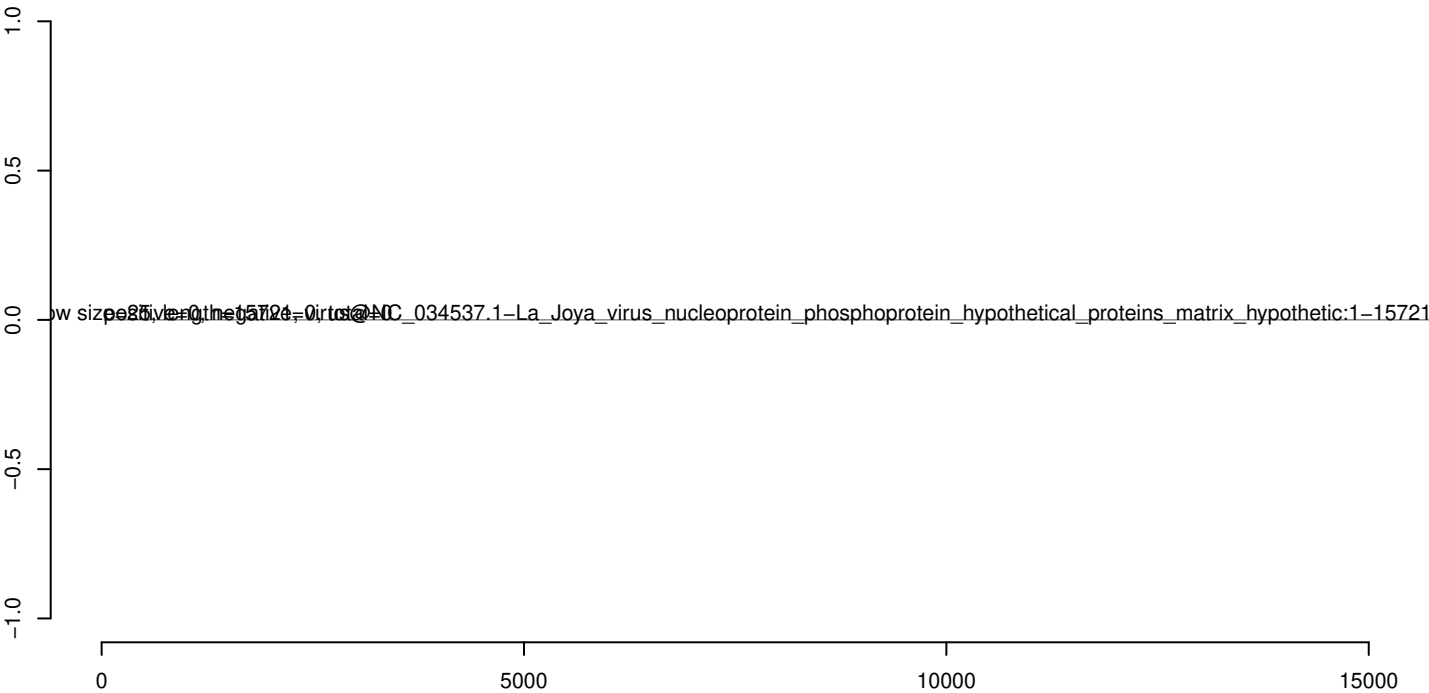
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



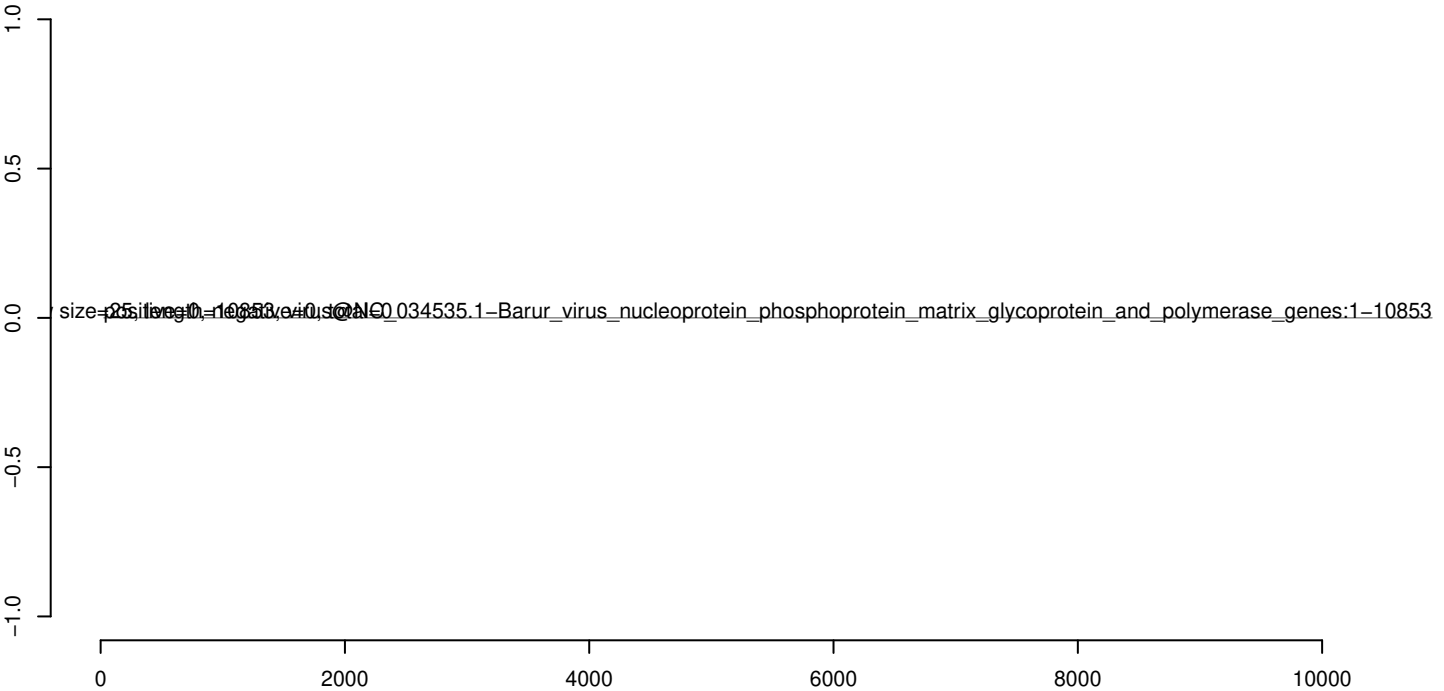
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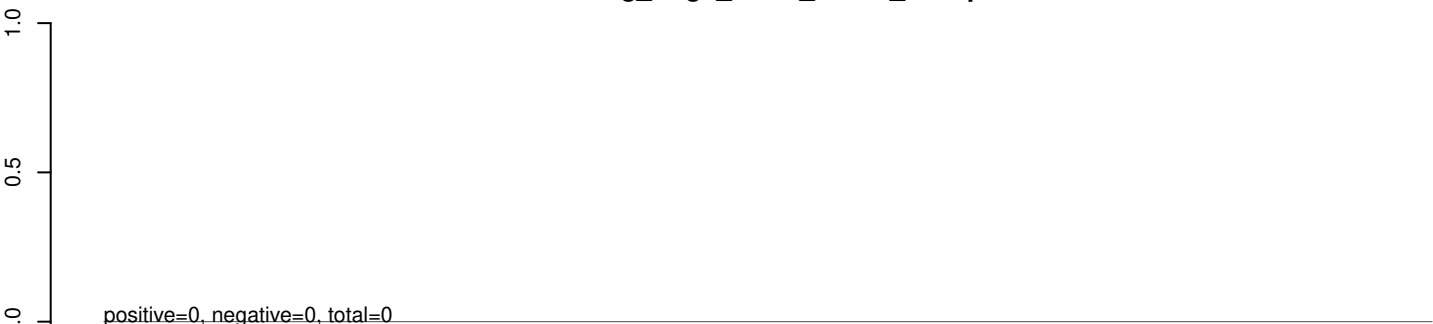
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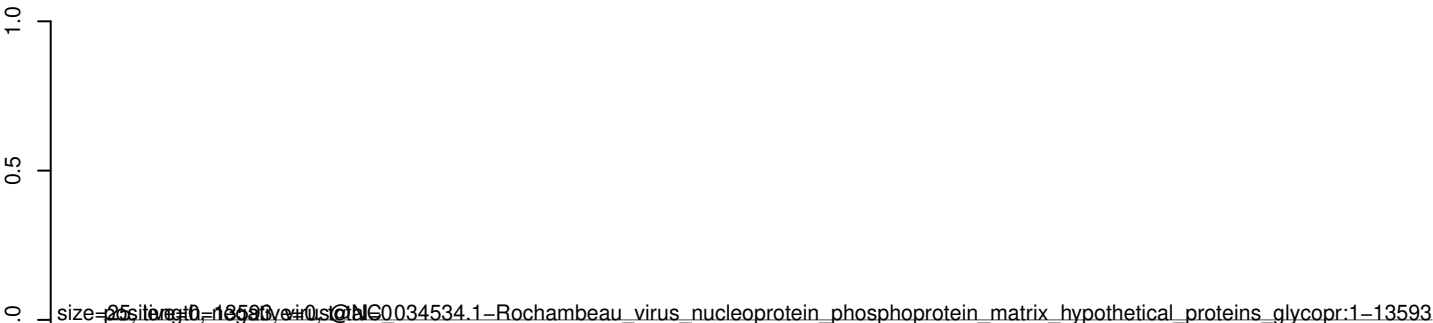
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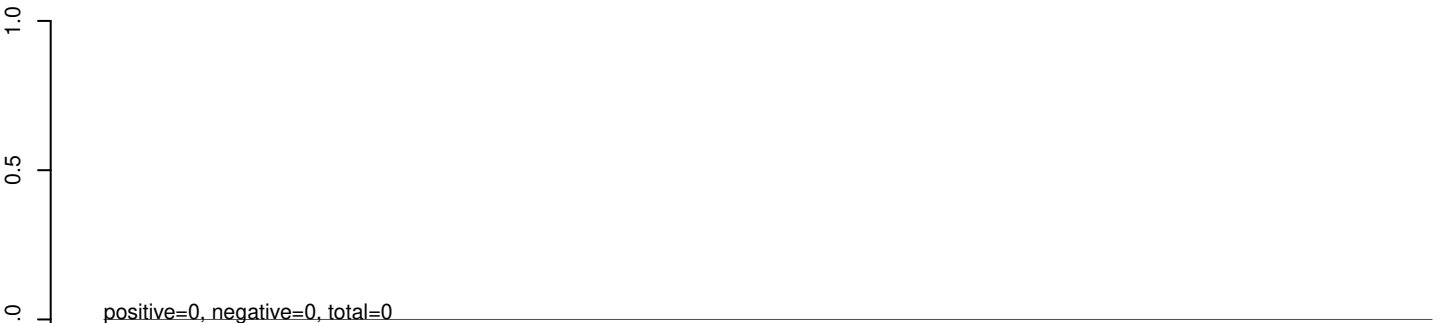
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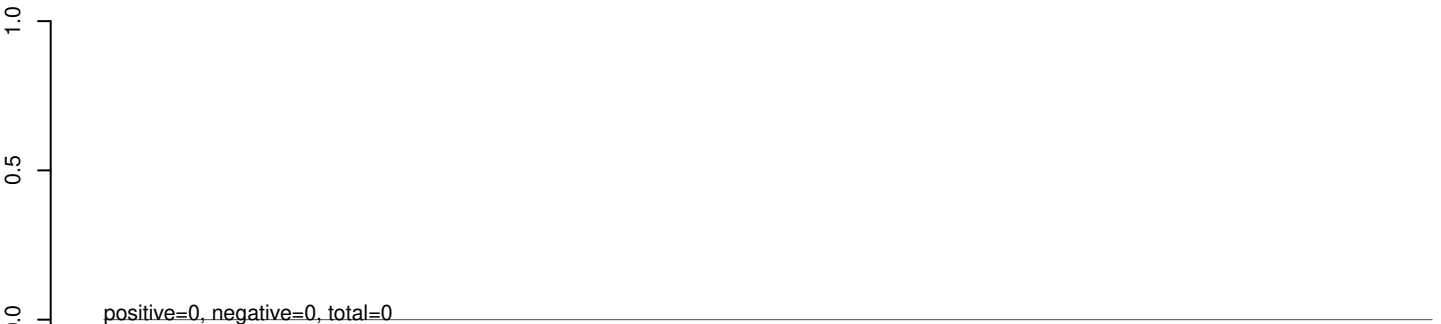
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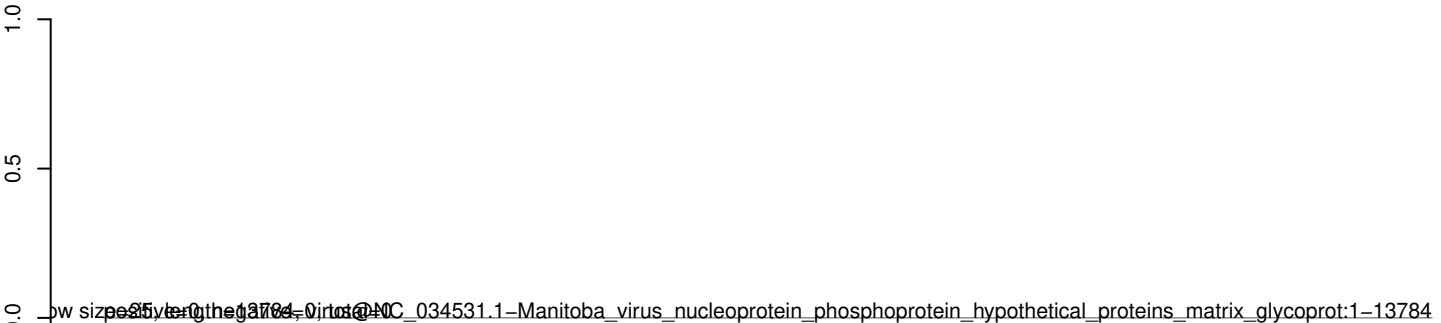
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AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



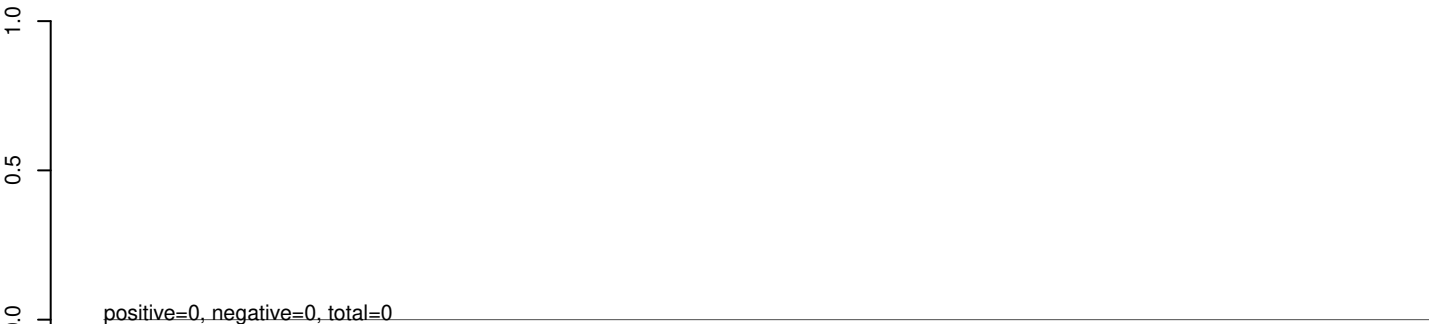
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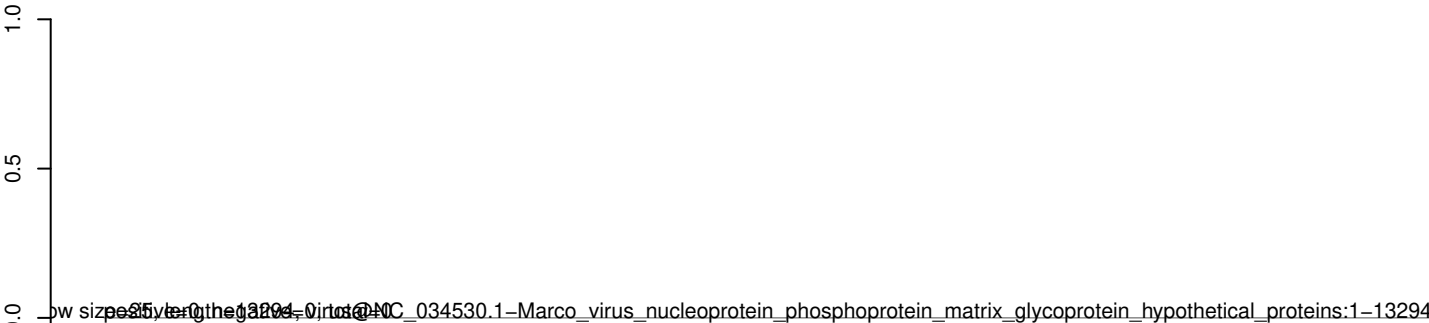
AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



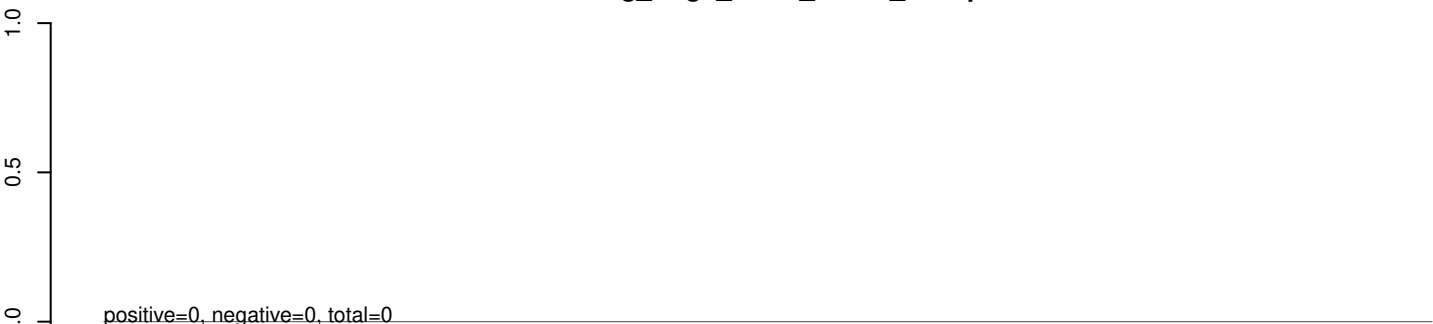
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0 2000 4000 6000 8000 10000 12000

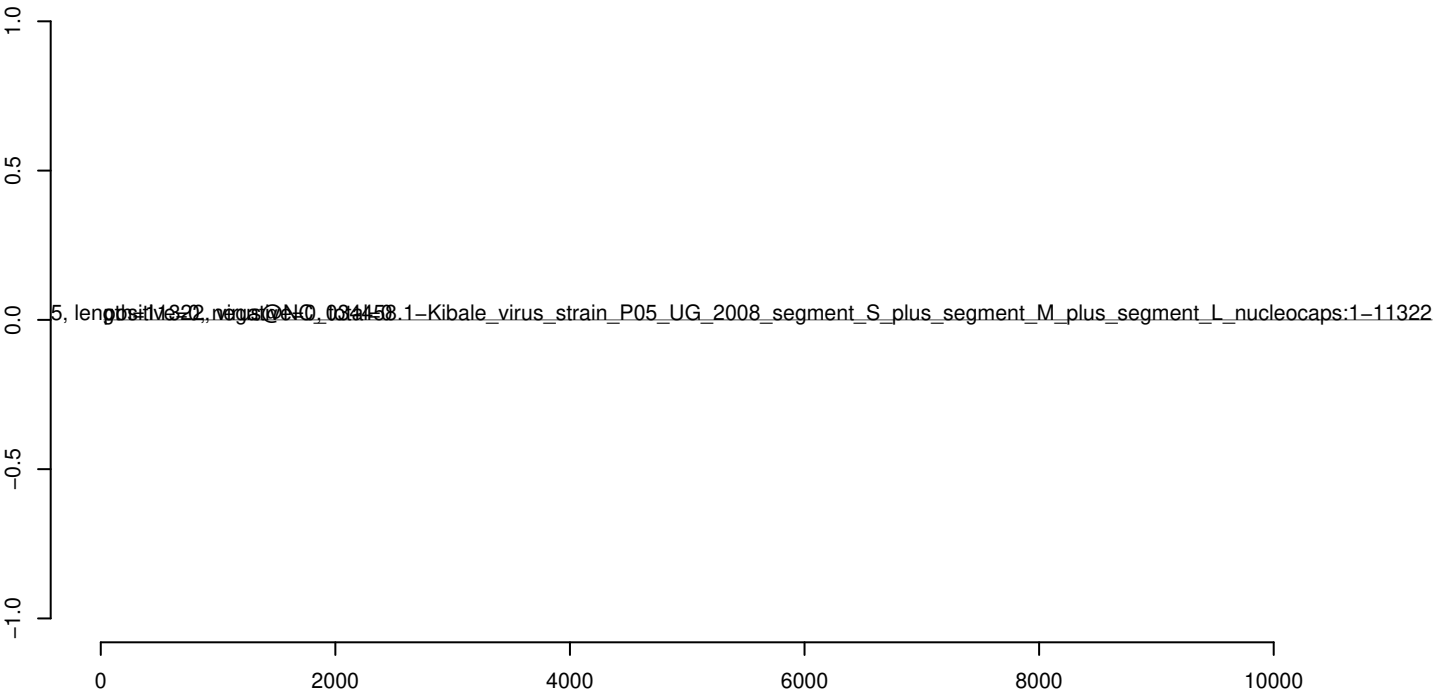
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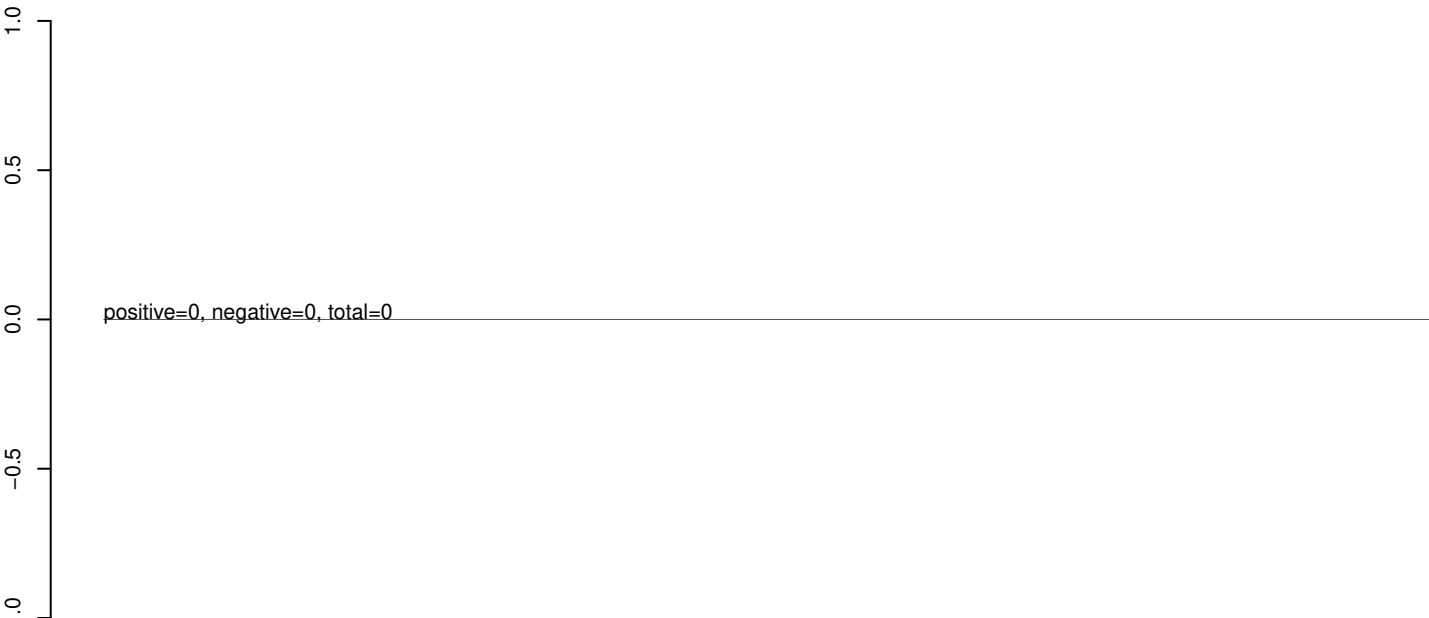
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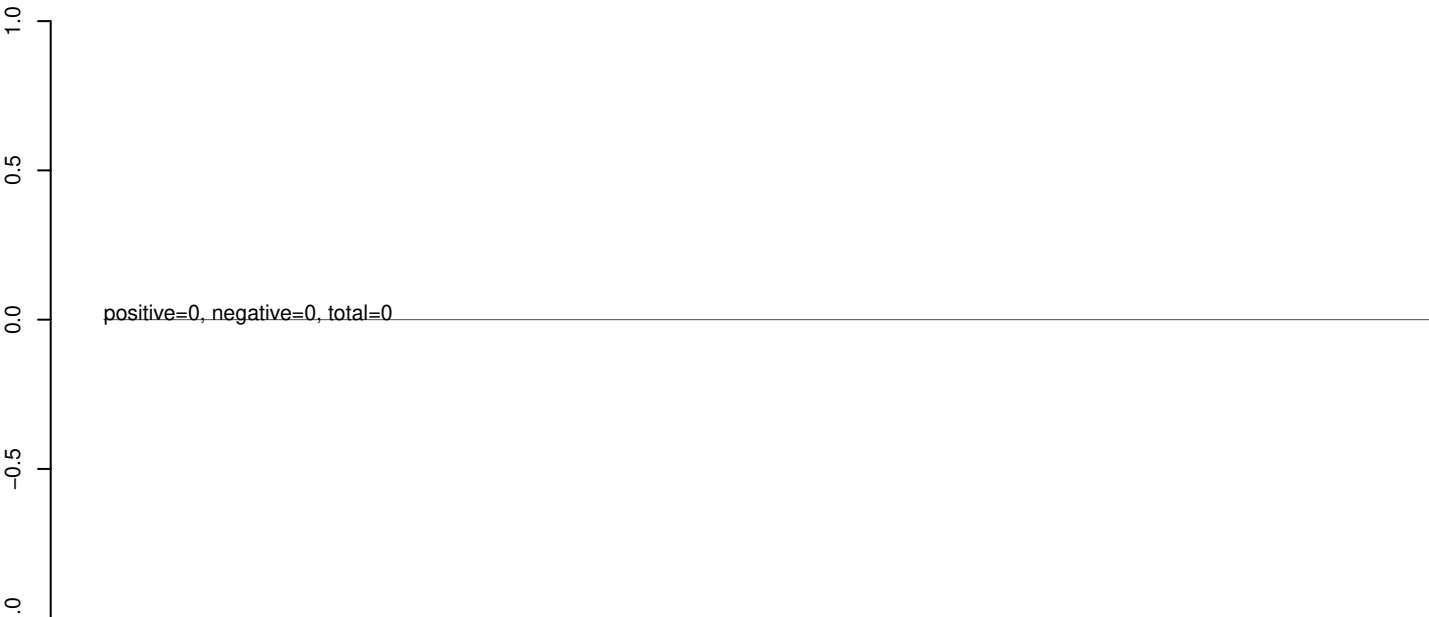
AeAeg_Aag2_BTV1_AK.rep



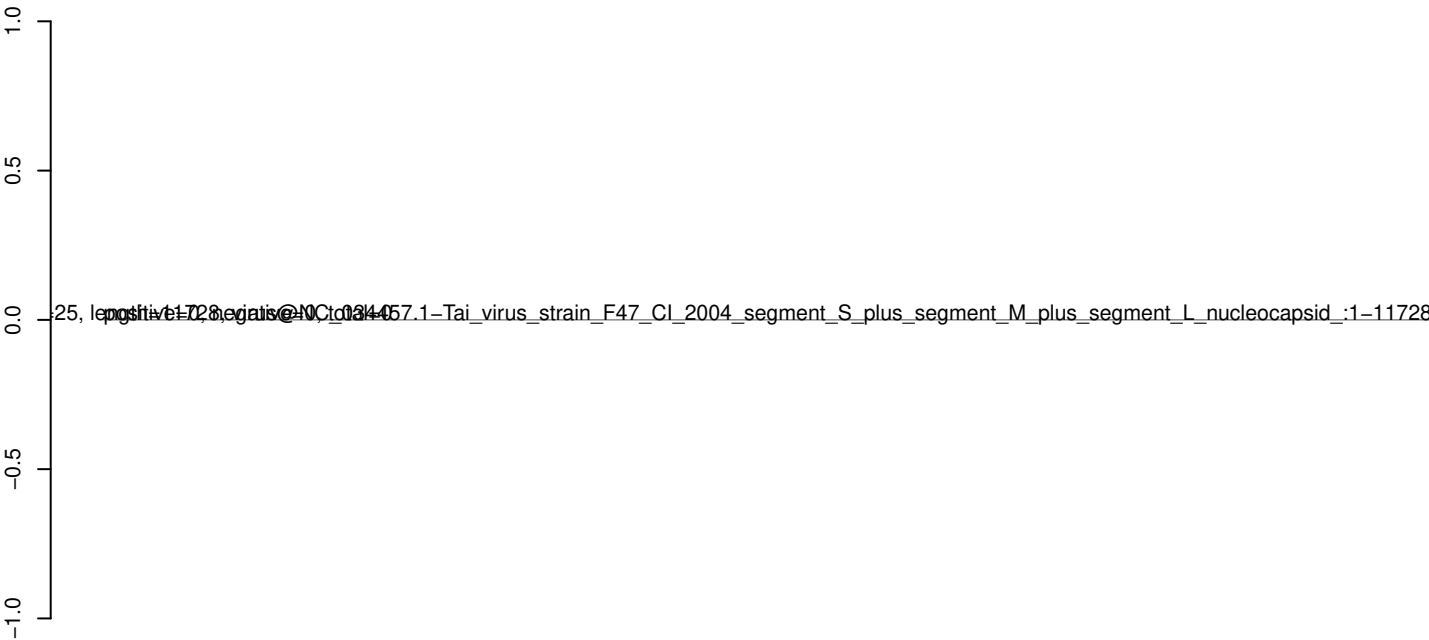
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AeAeg_Aag2_BTV1_AK.24_35.rep



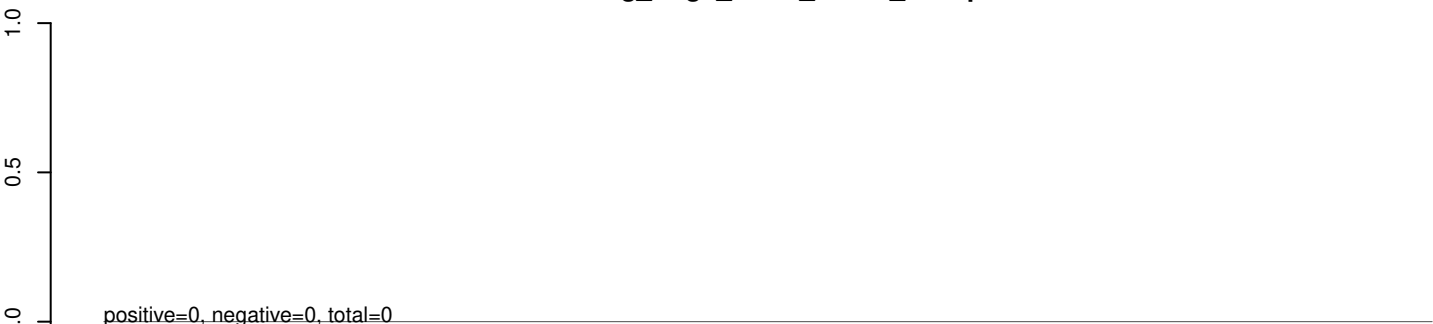
AeAeg_Aag2_BTV1_AK.rep



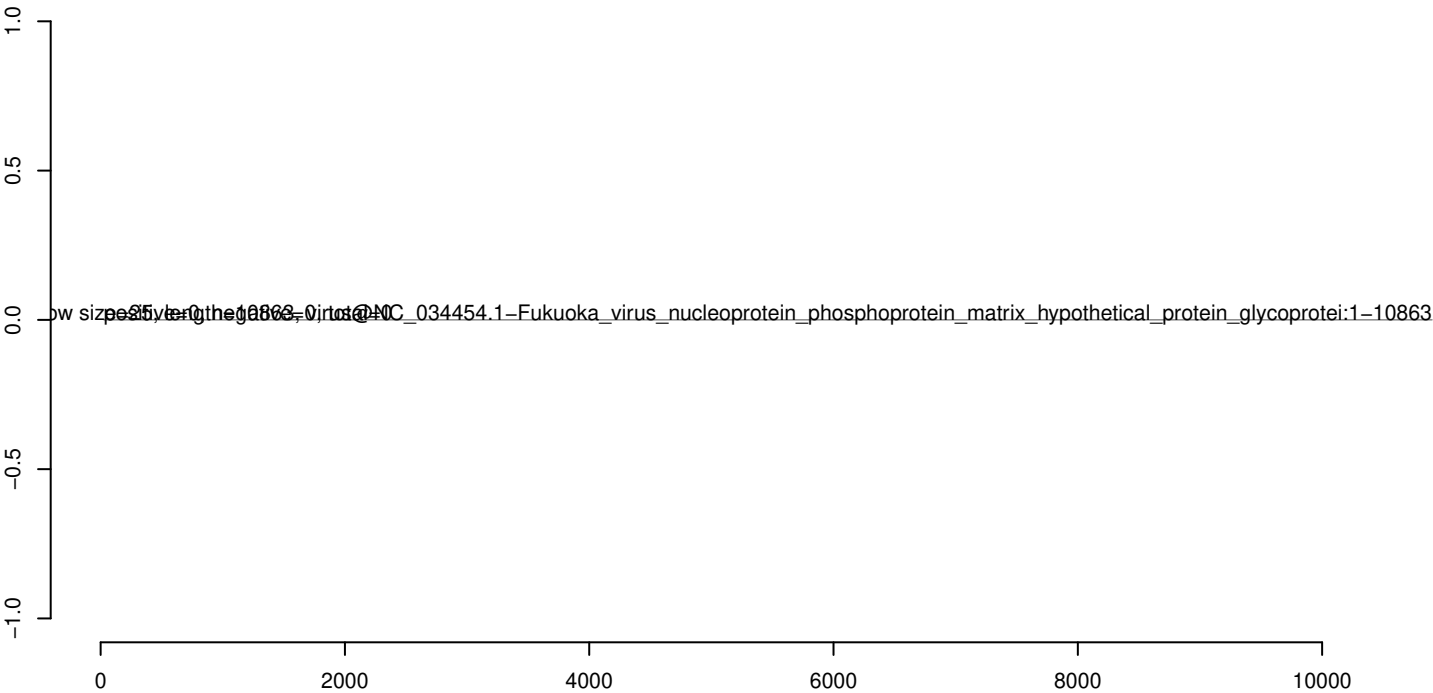
AeAeg_Aag2_BTV1_AK.18_23.rep



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AeAeg_Aag2_BTV1_AK.rep



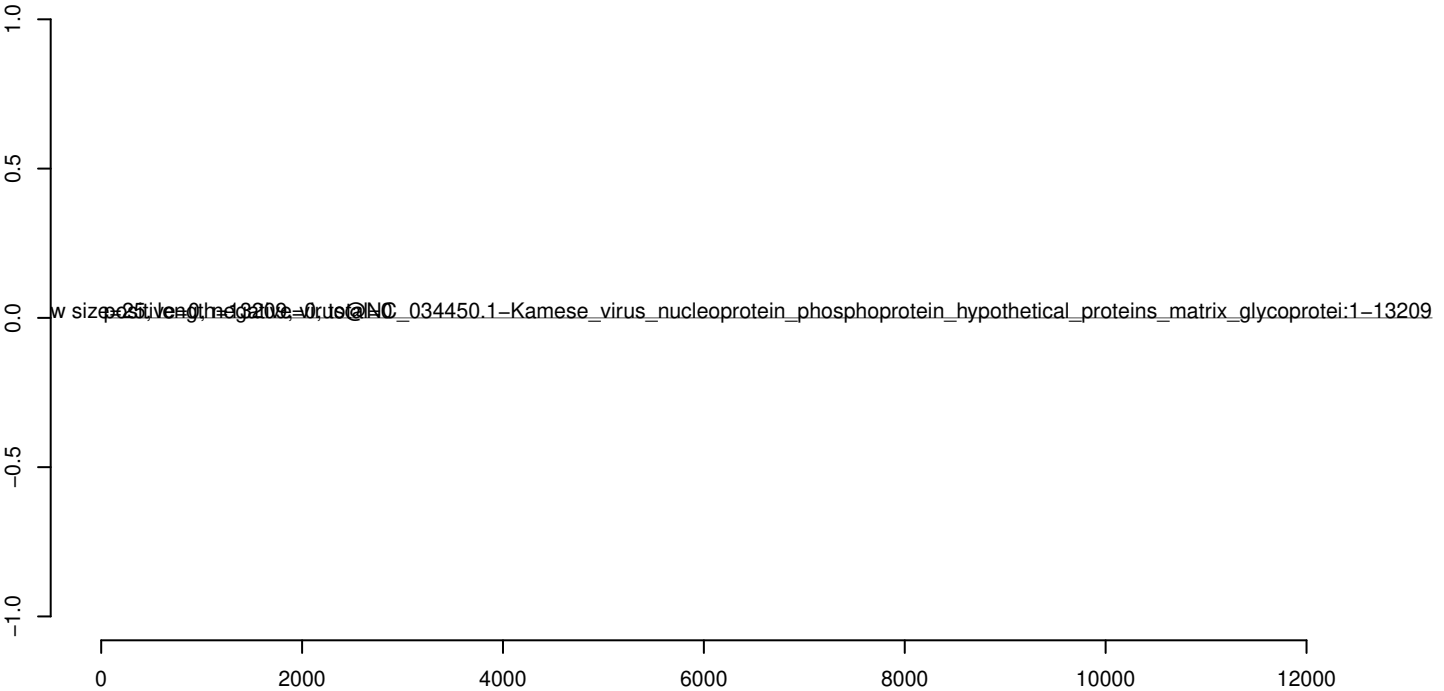
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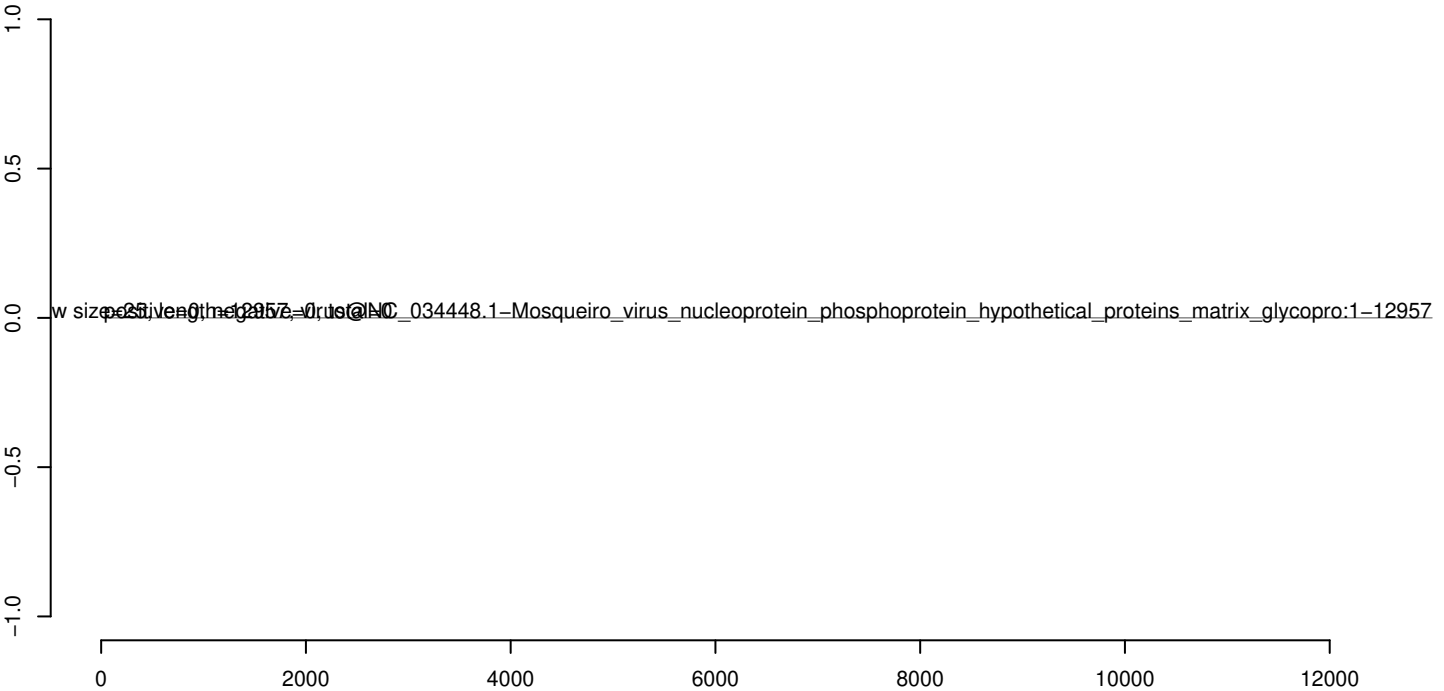
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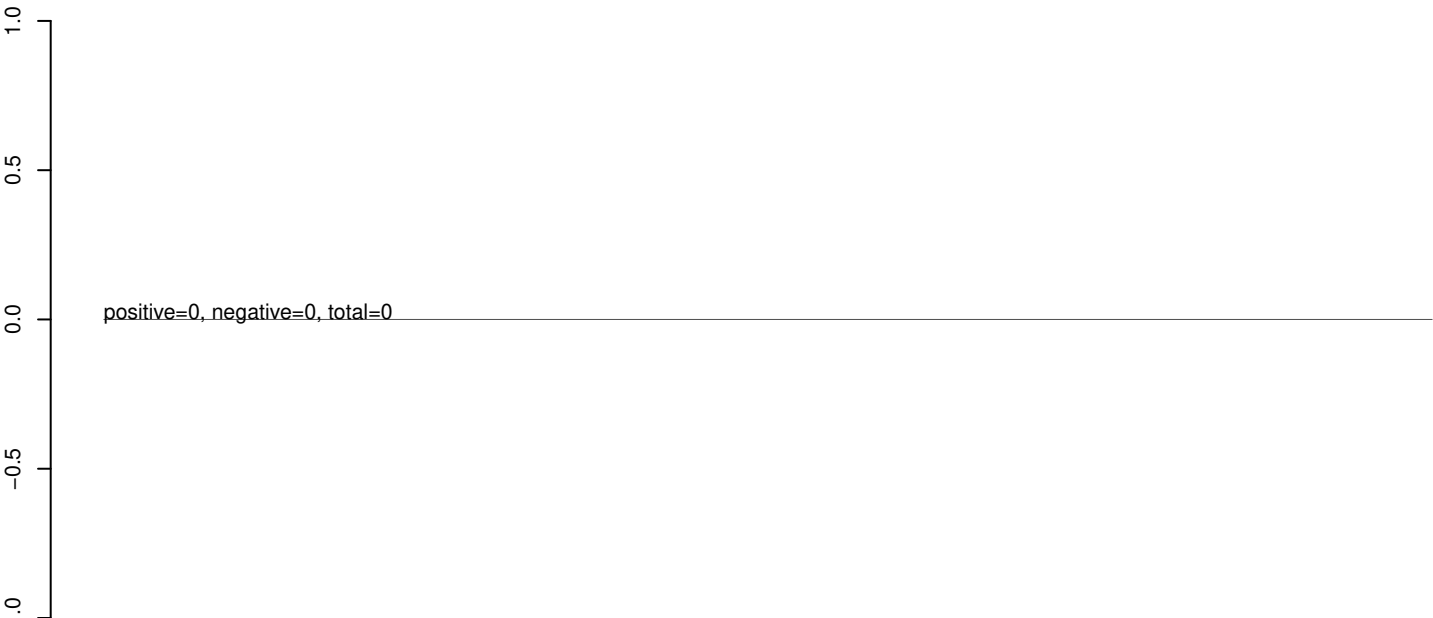
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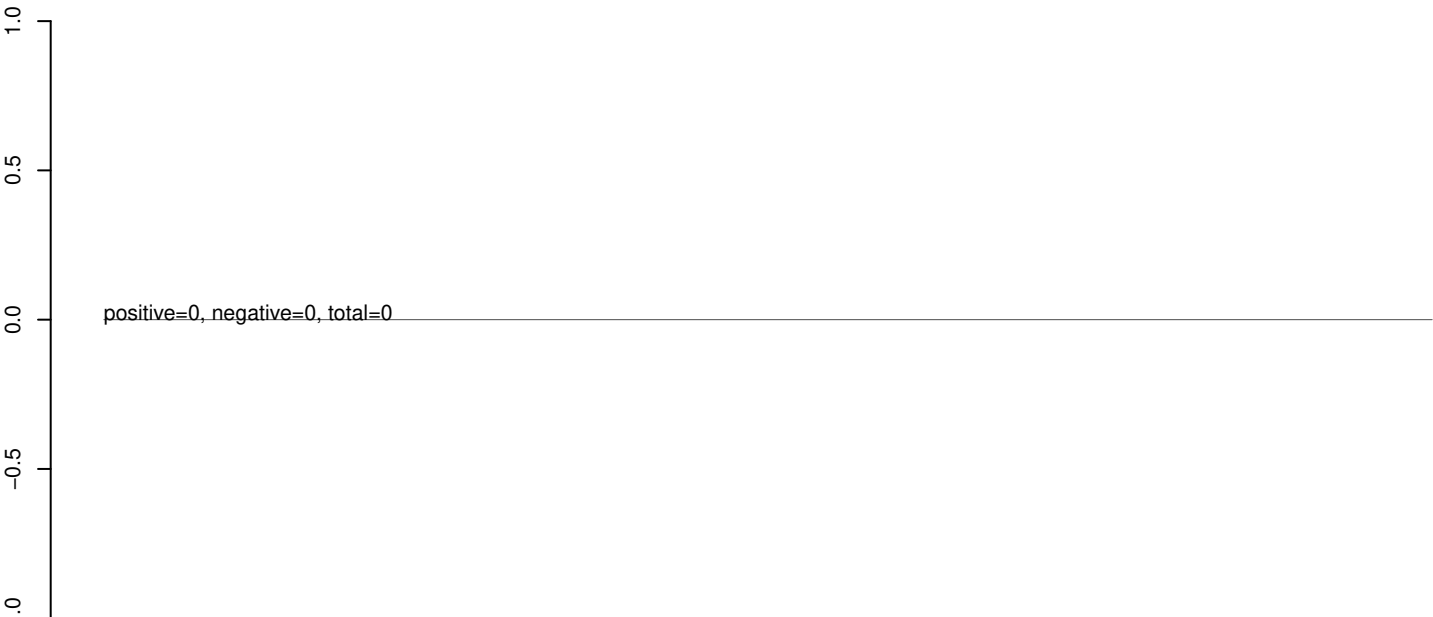
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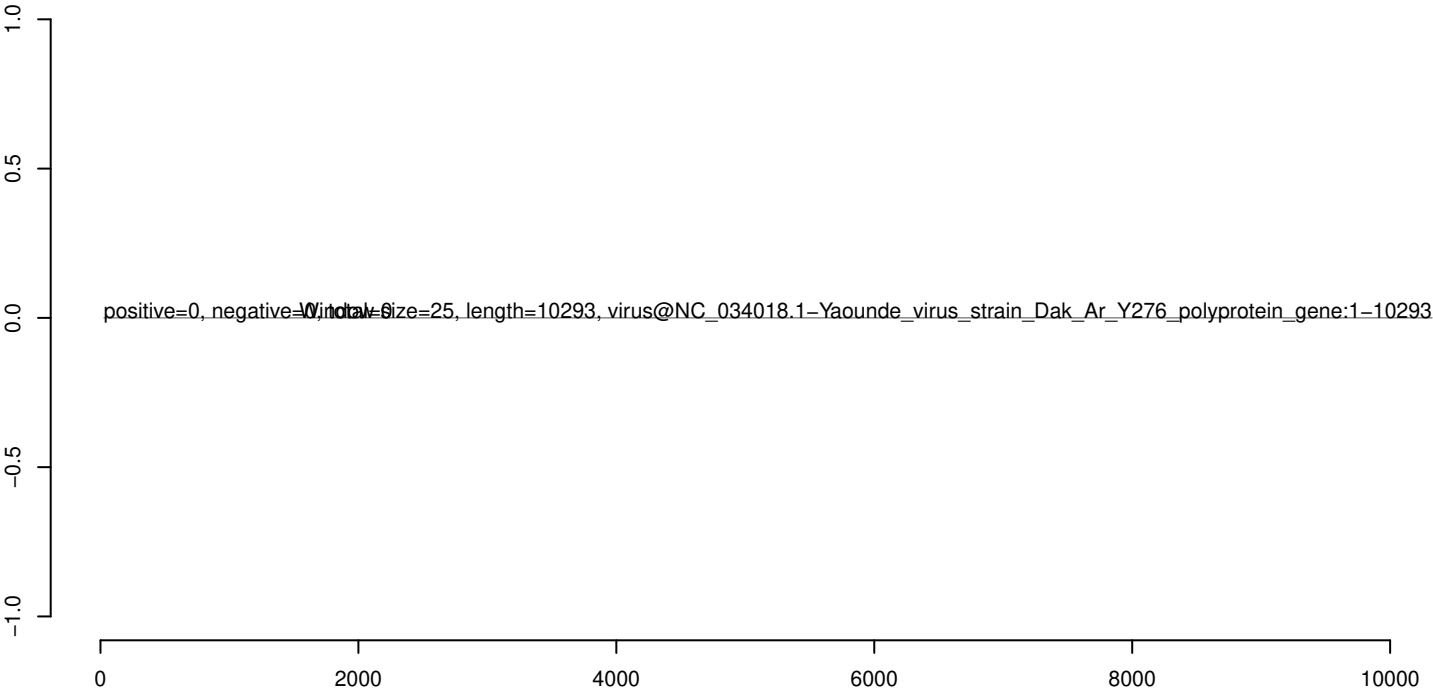
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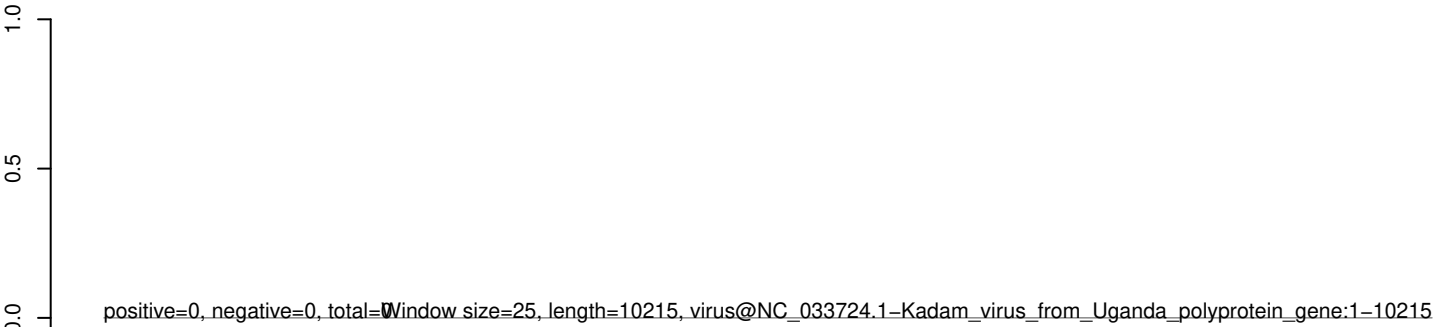
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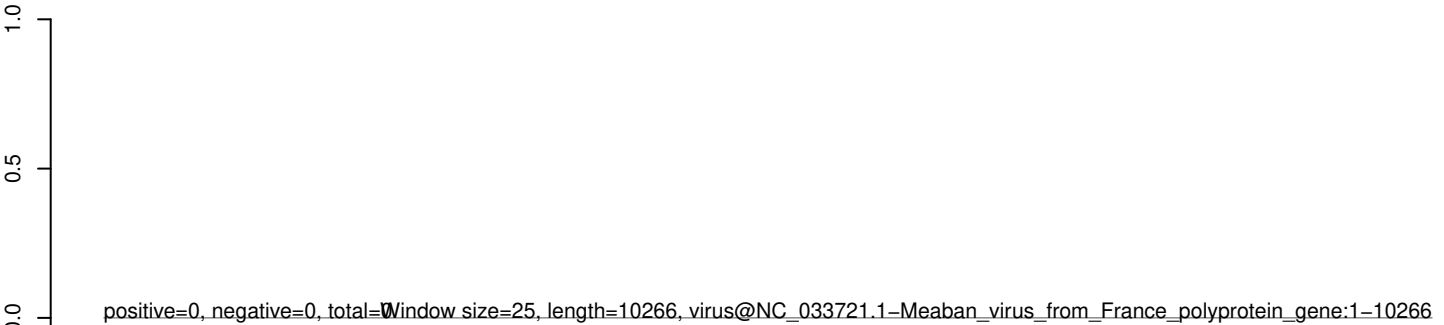
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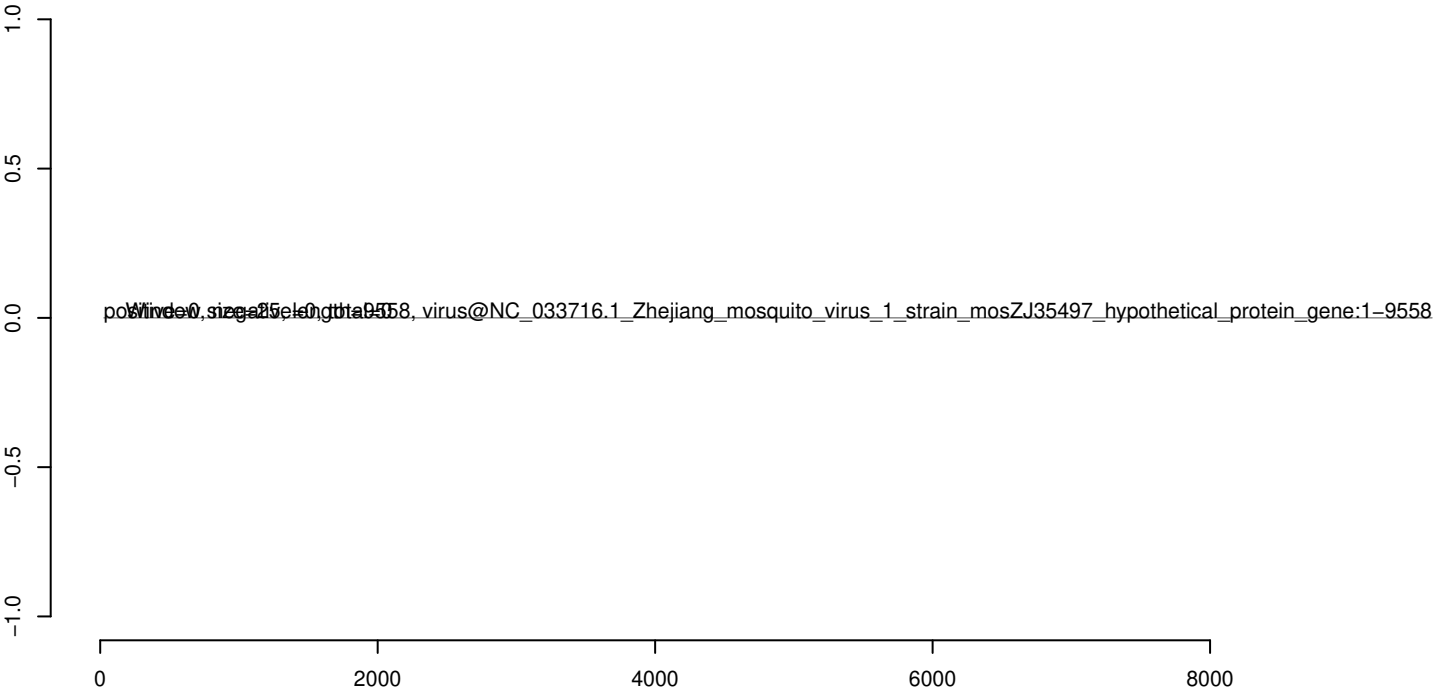
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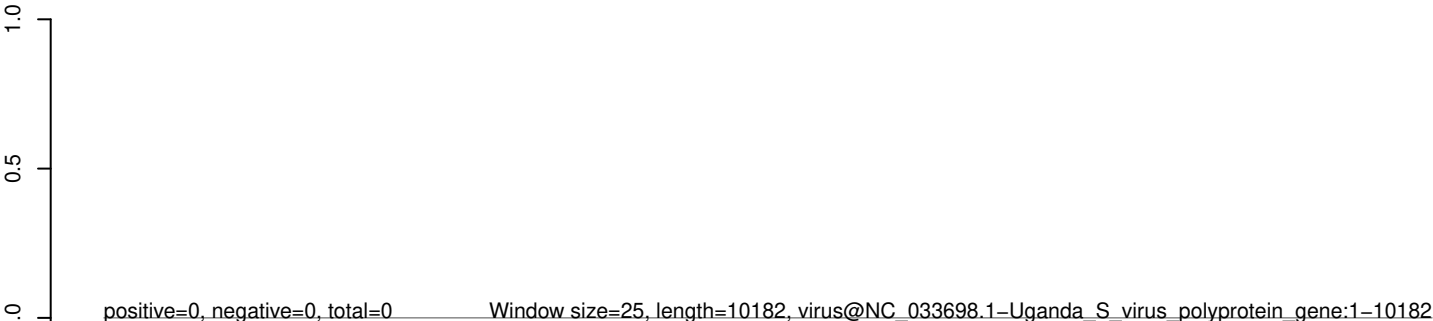
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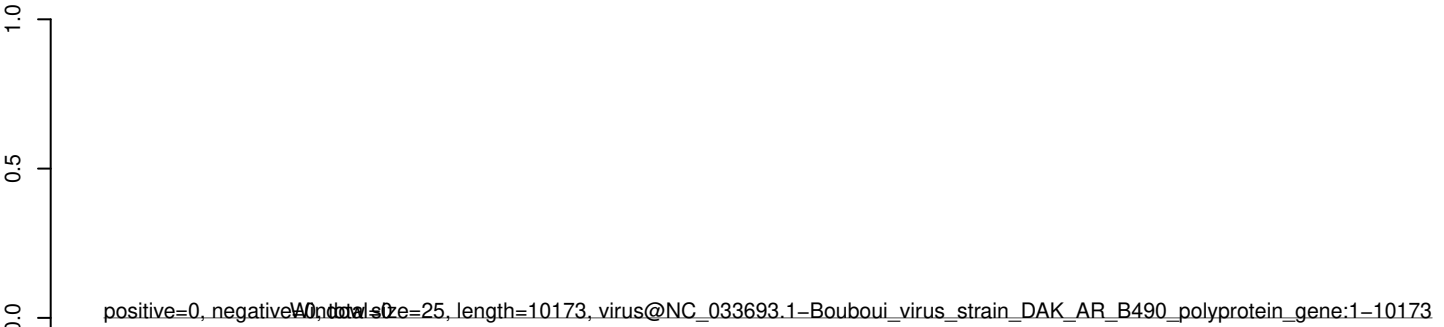
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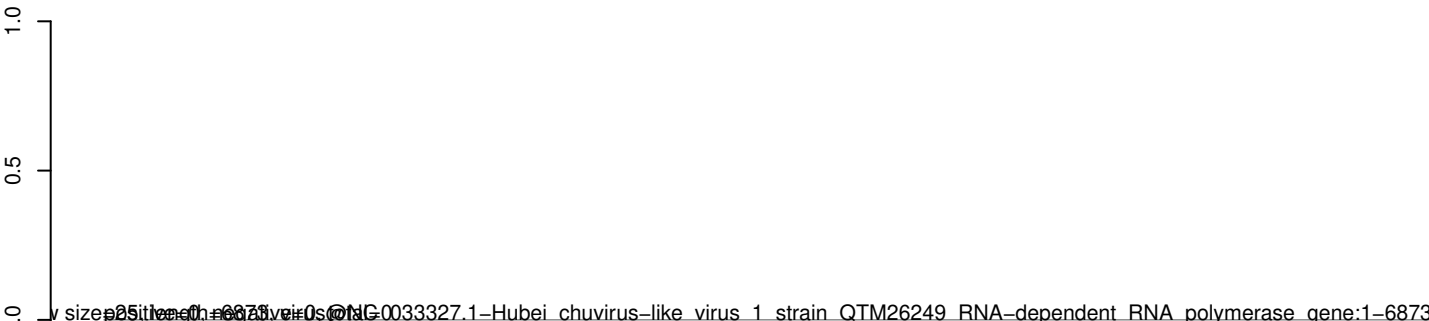
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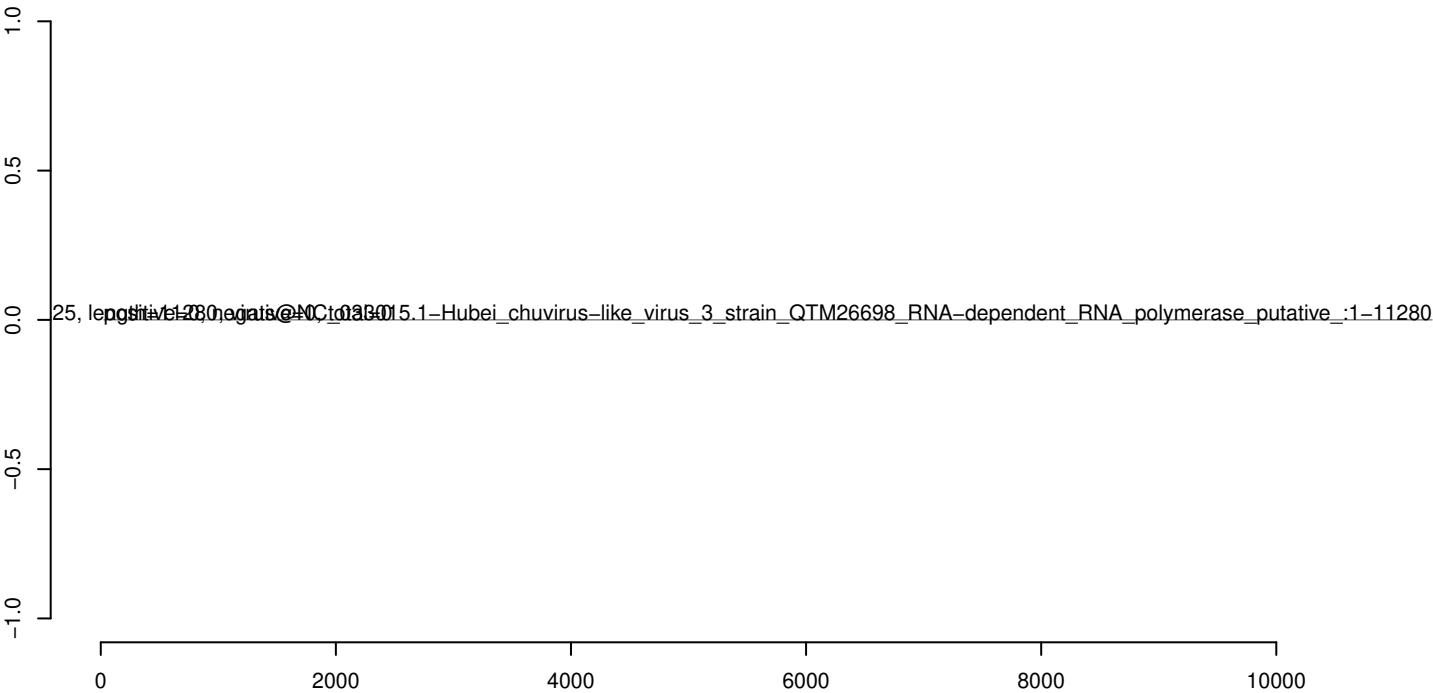
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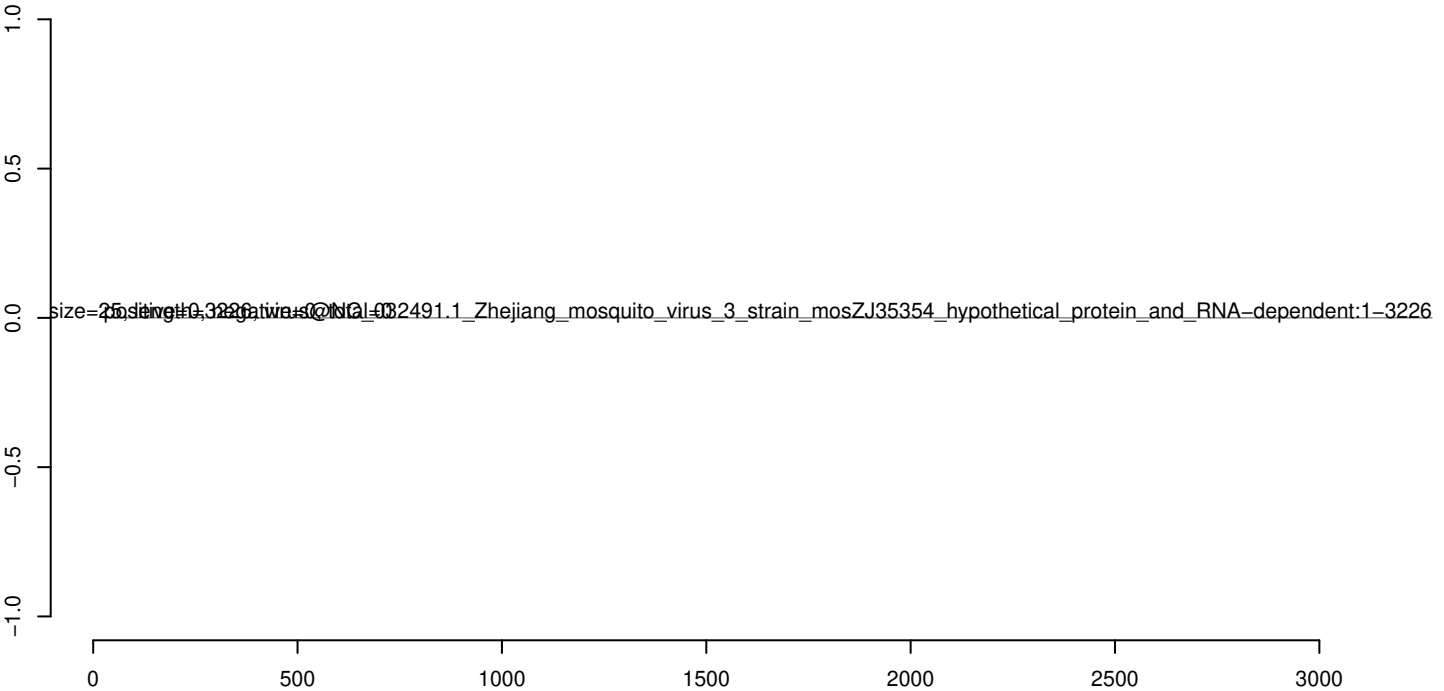
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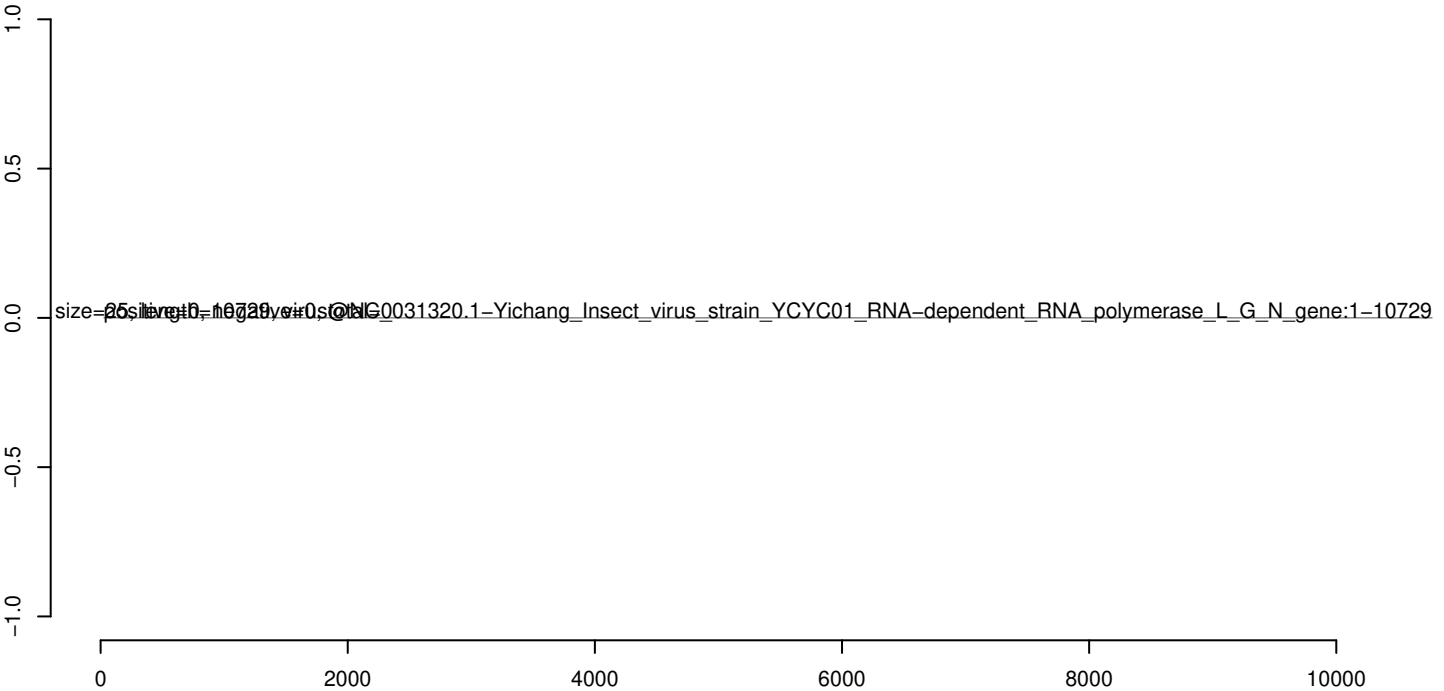
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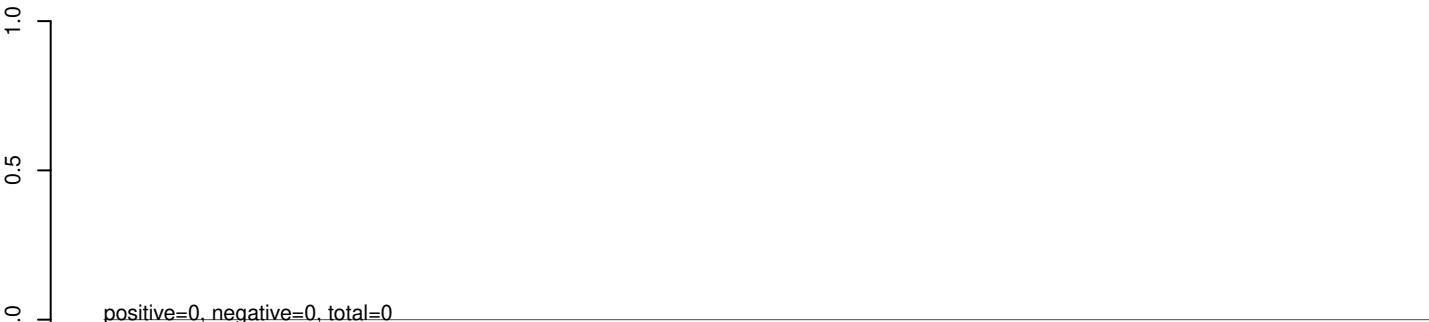
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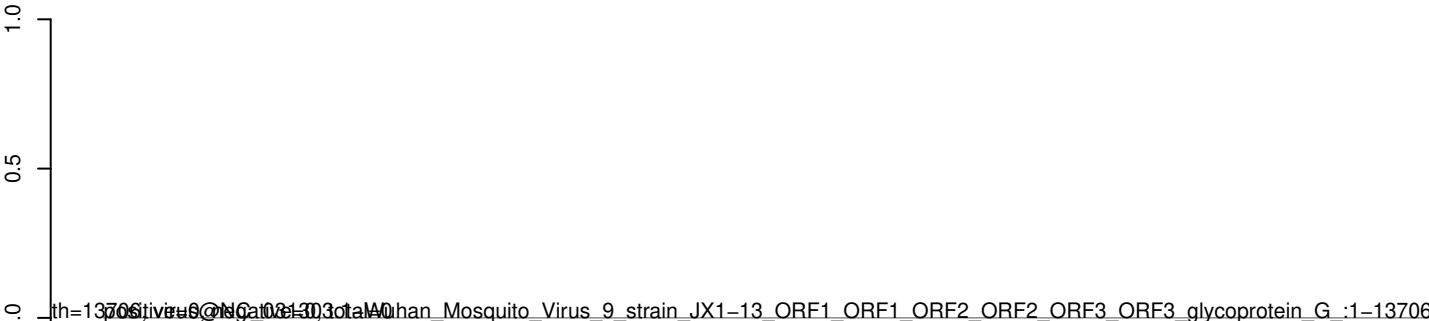
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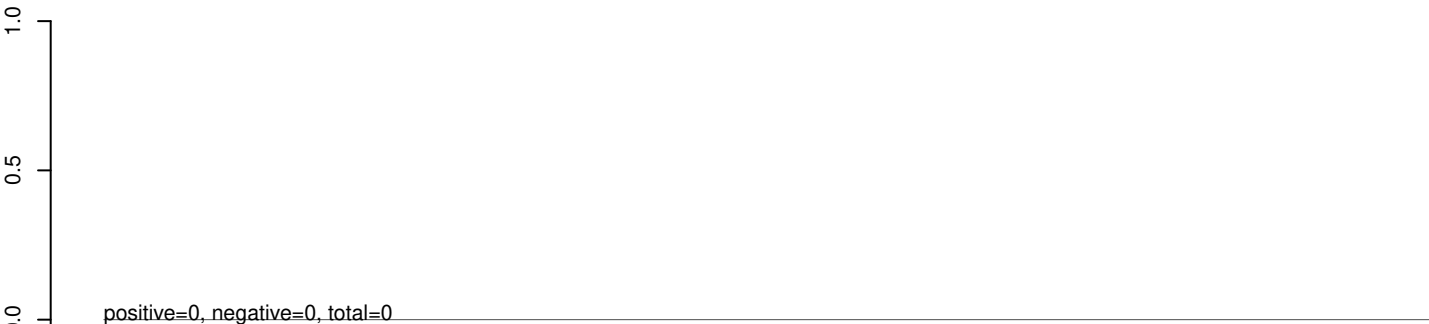


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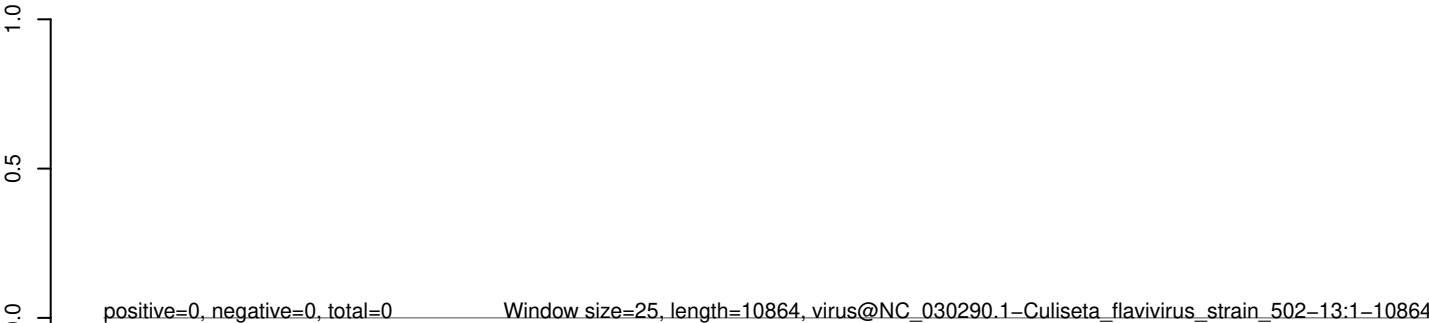
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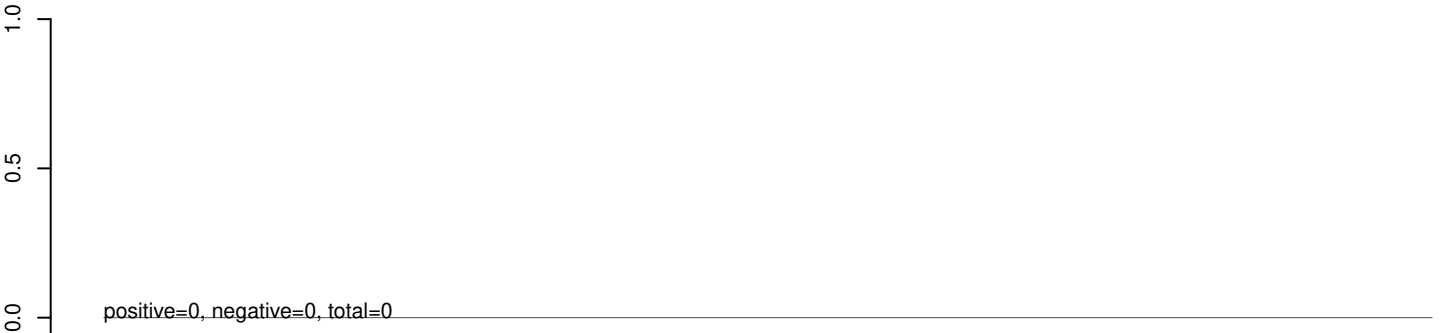


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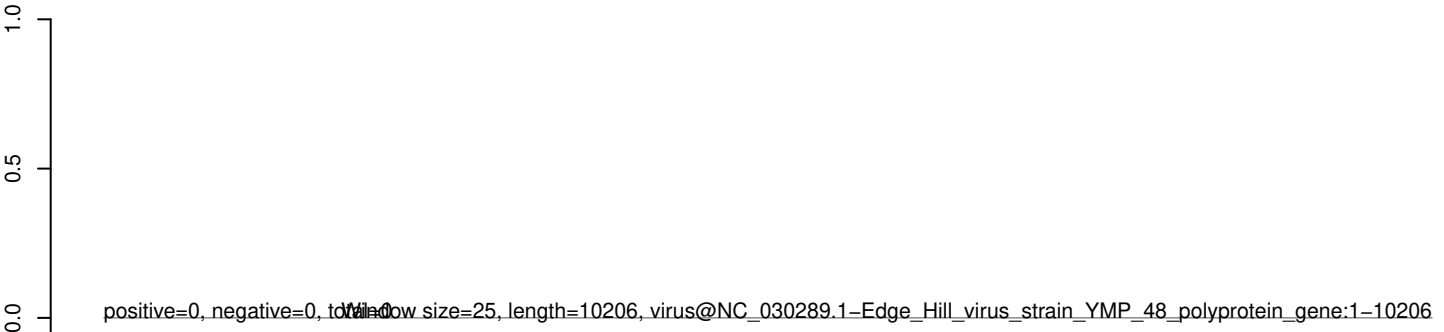
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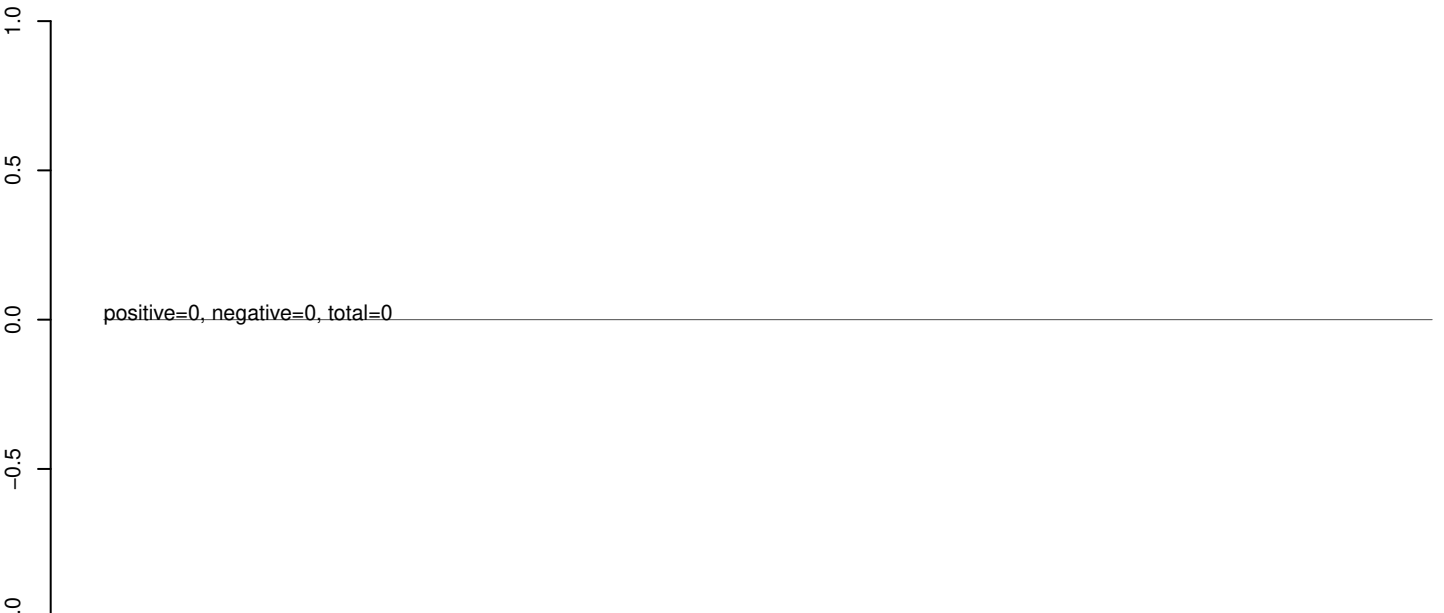
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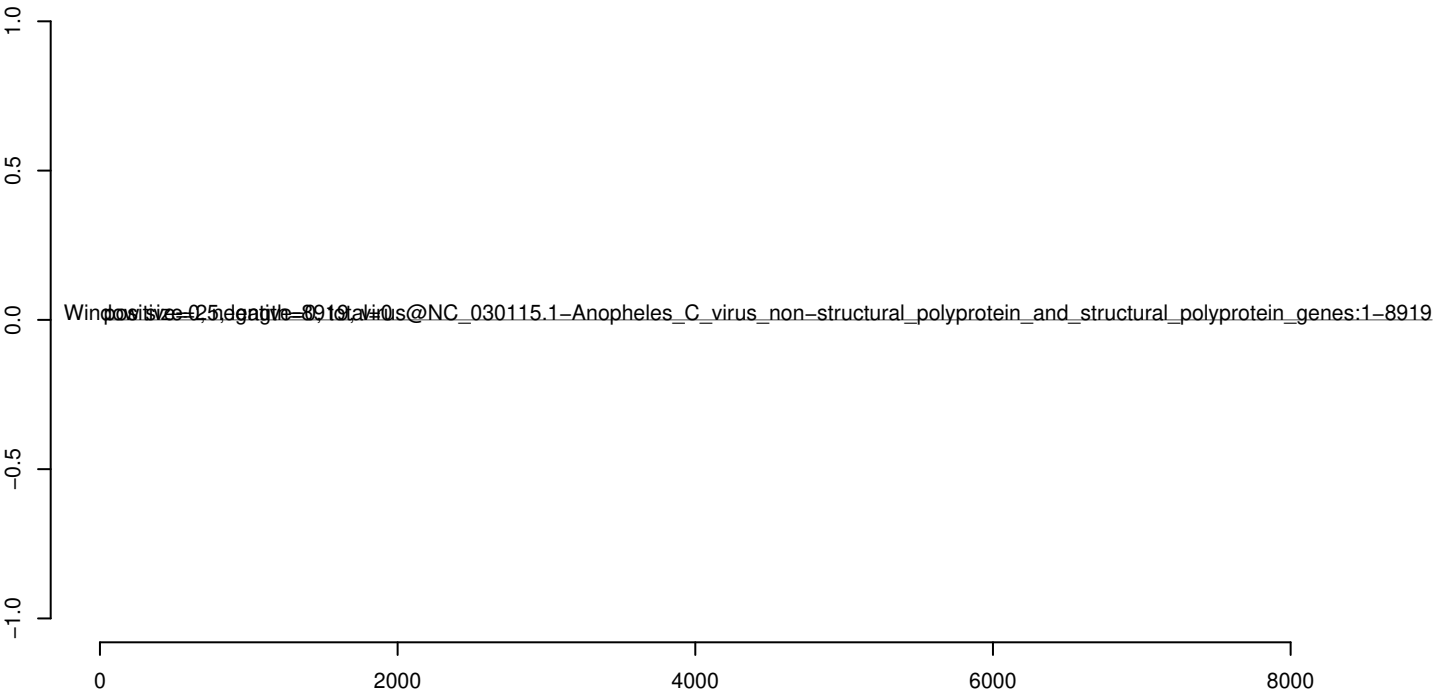
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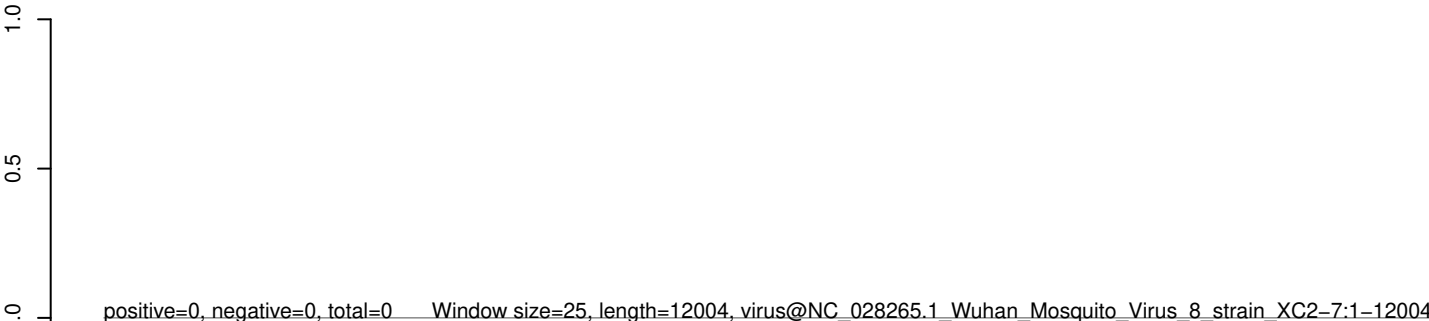
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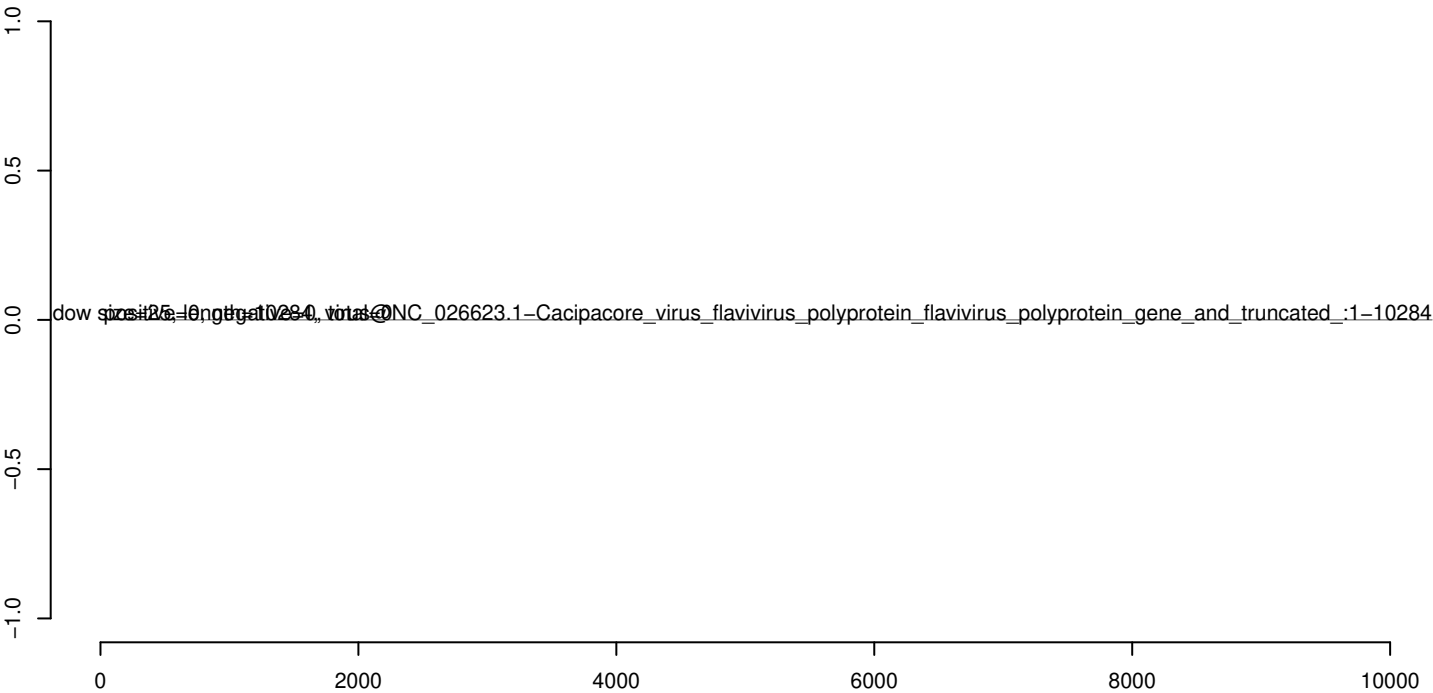
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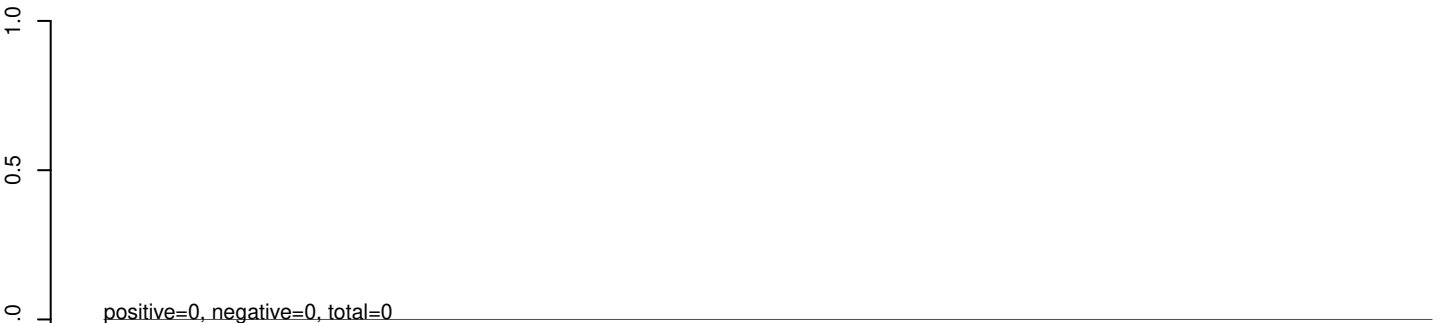
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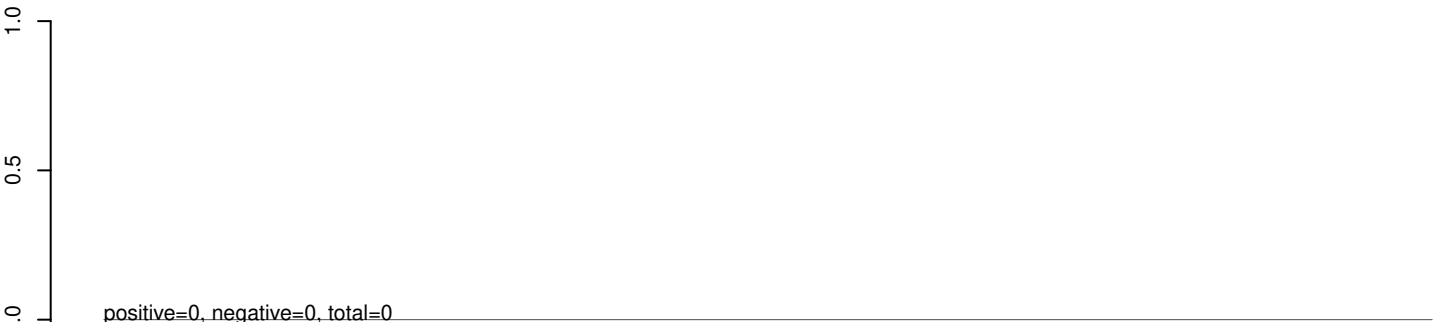
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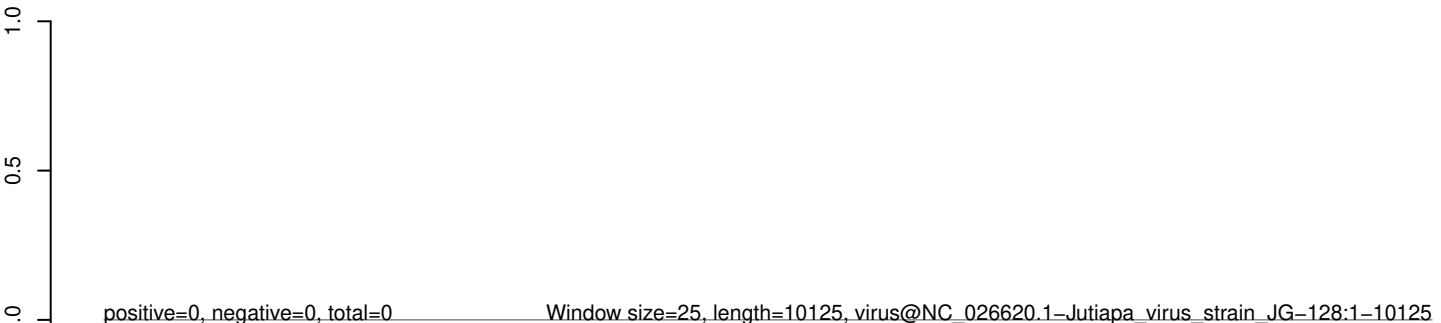
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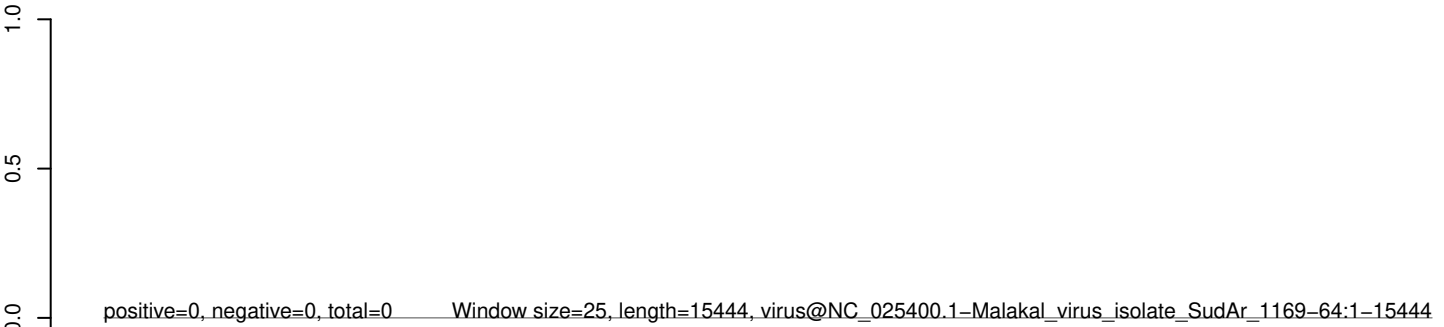
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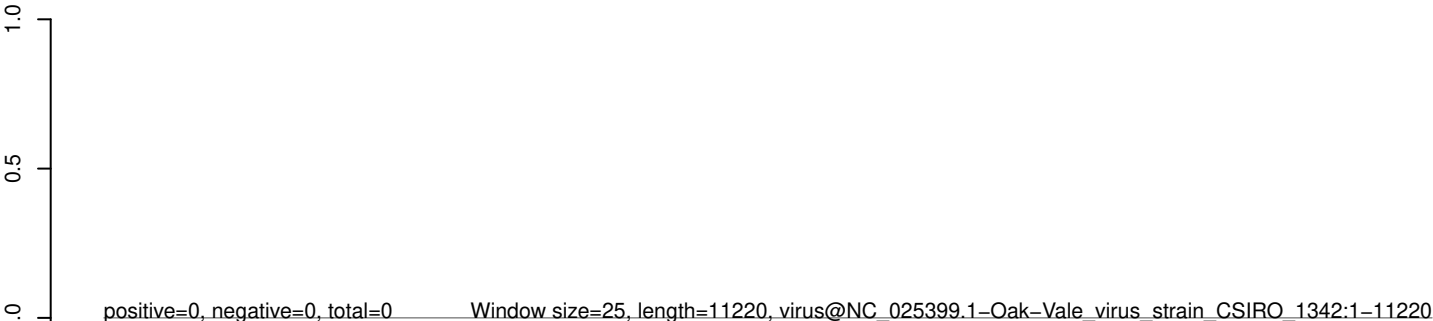
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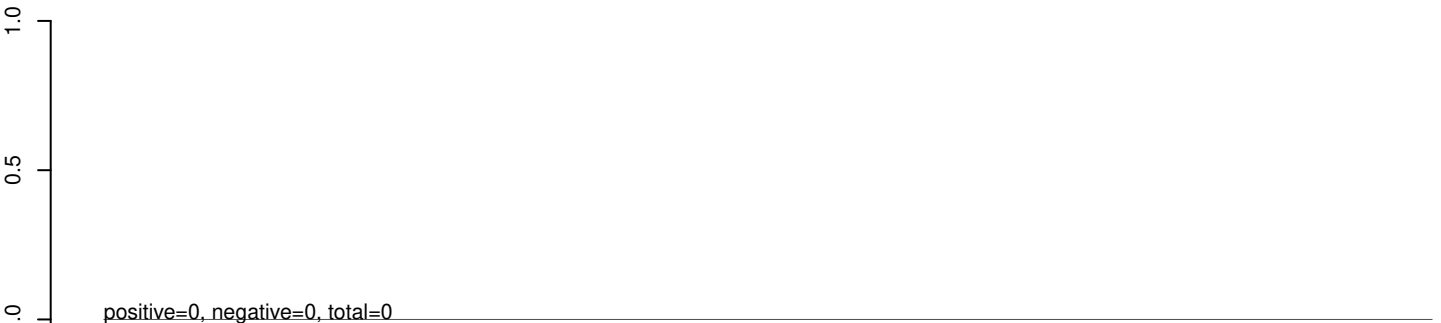


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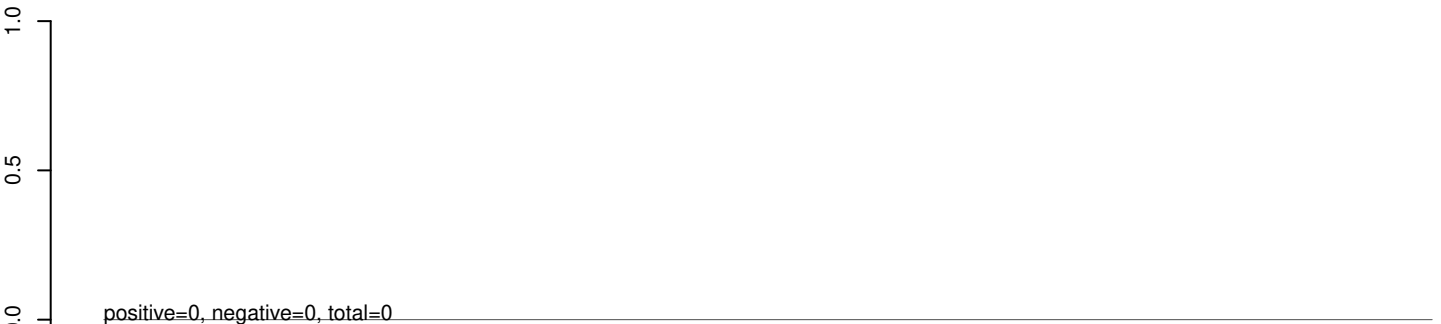


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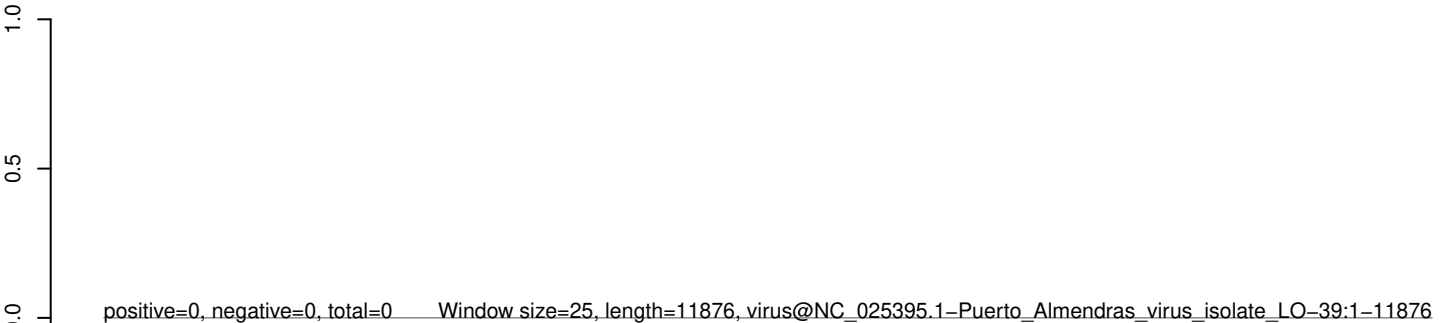
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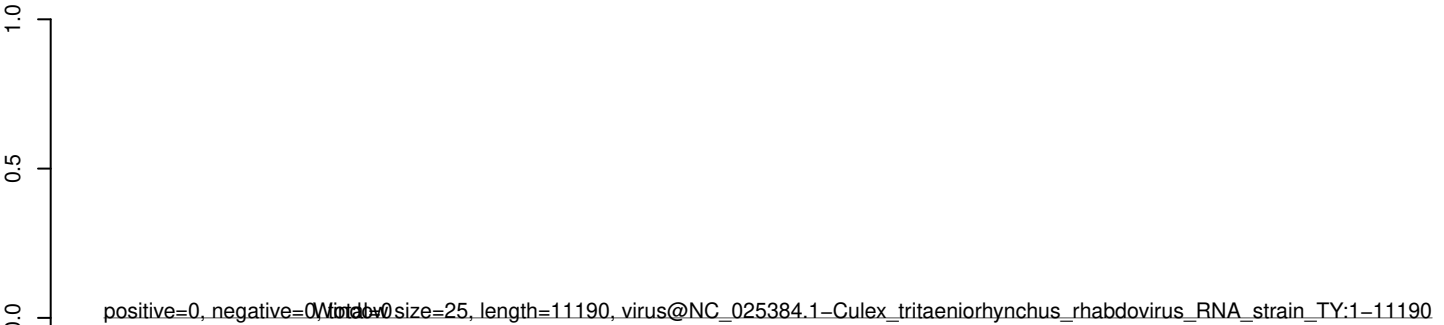
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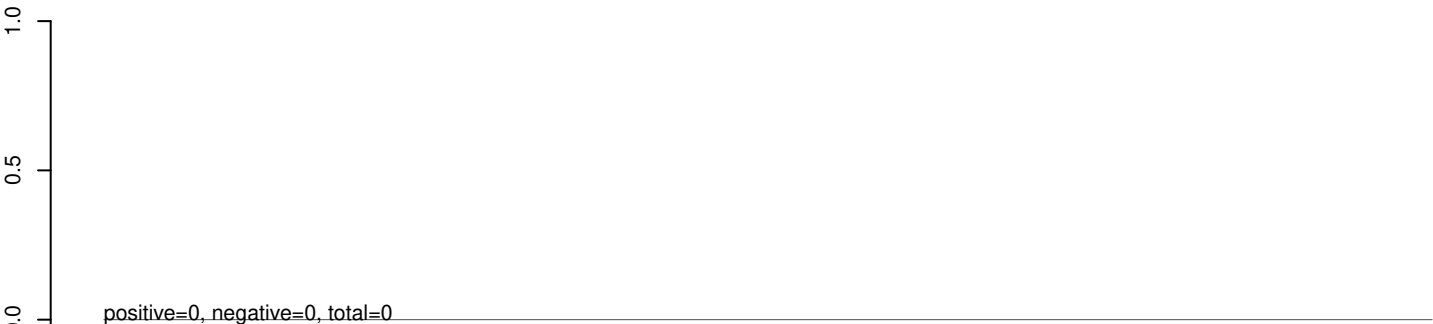
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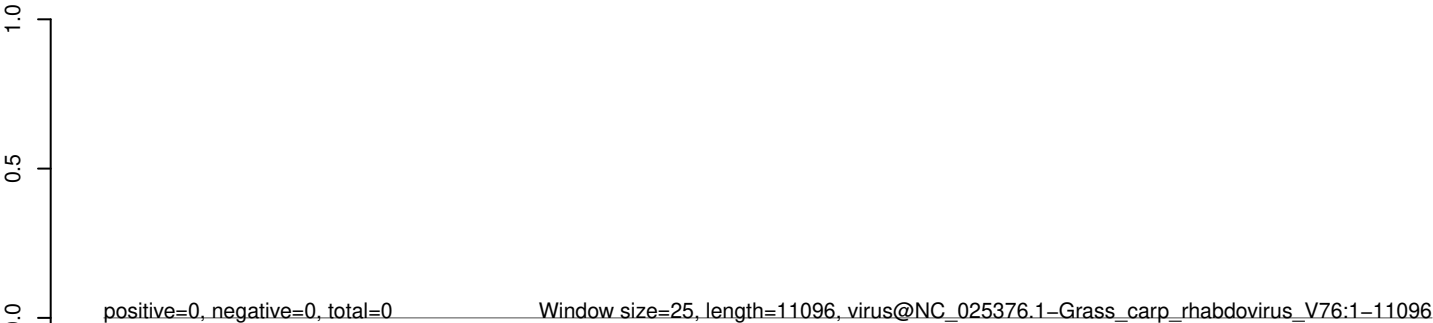
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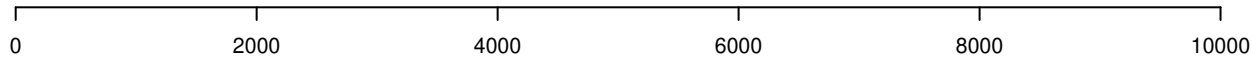
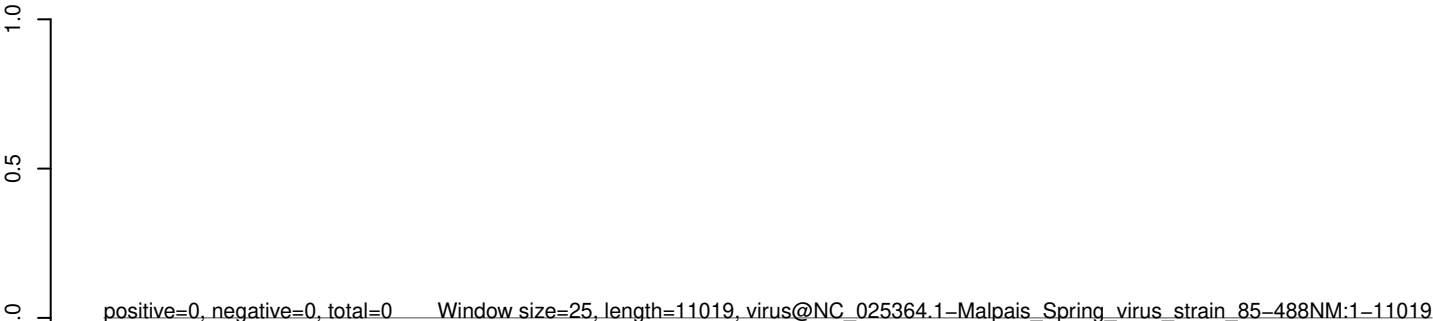
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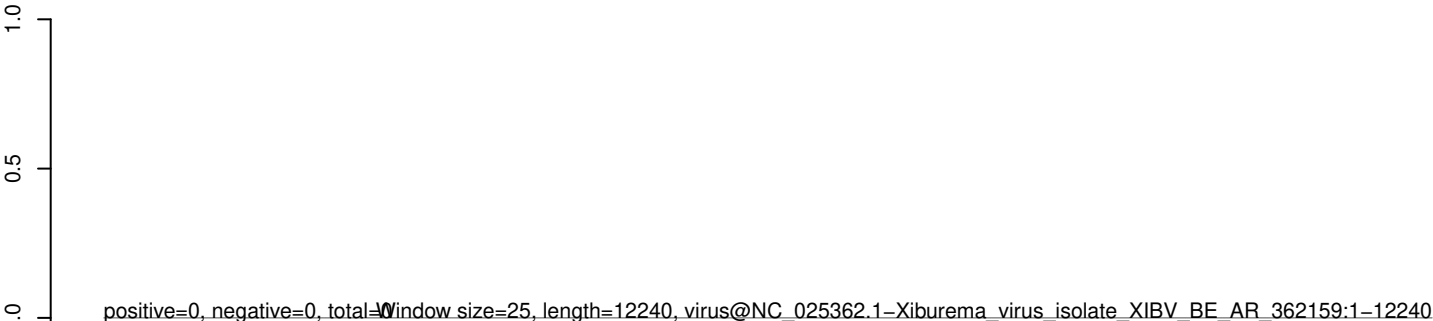
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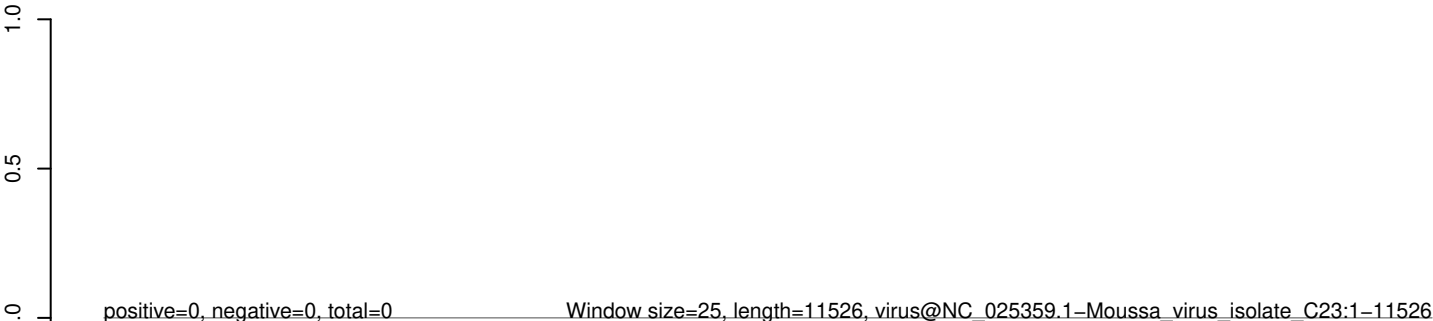
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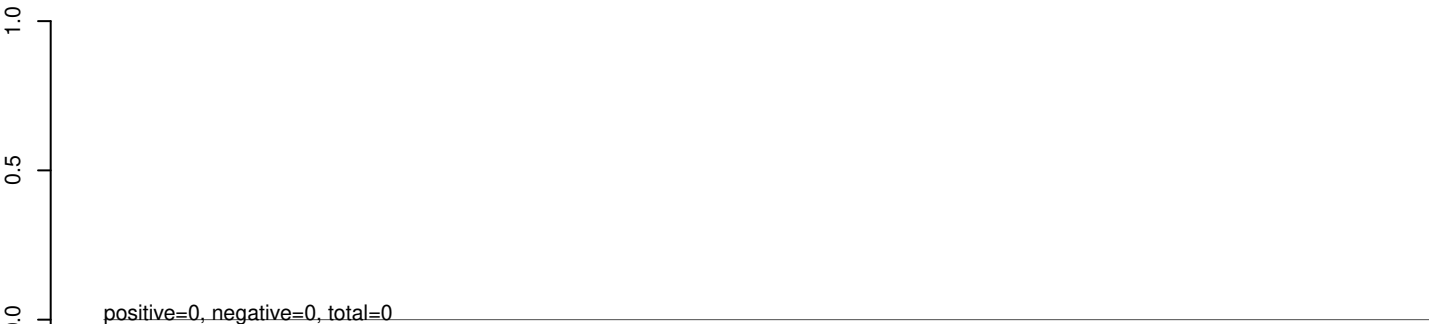


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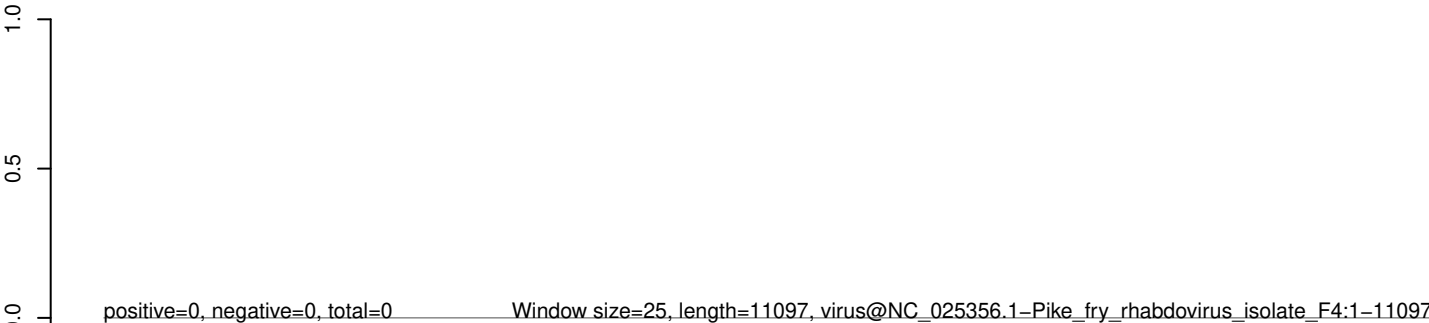
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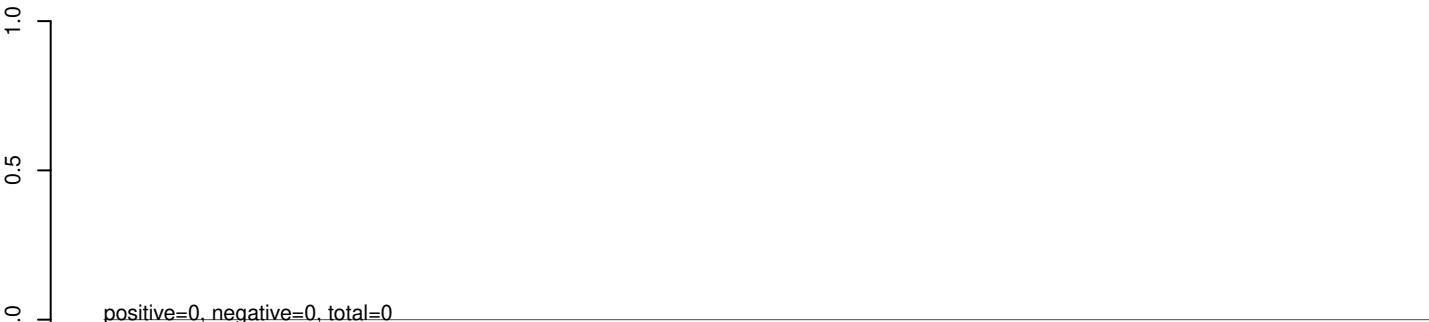


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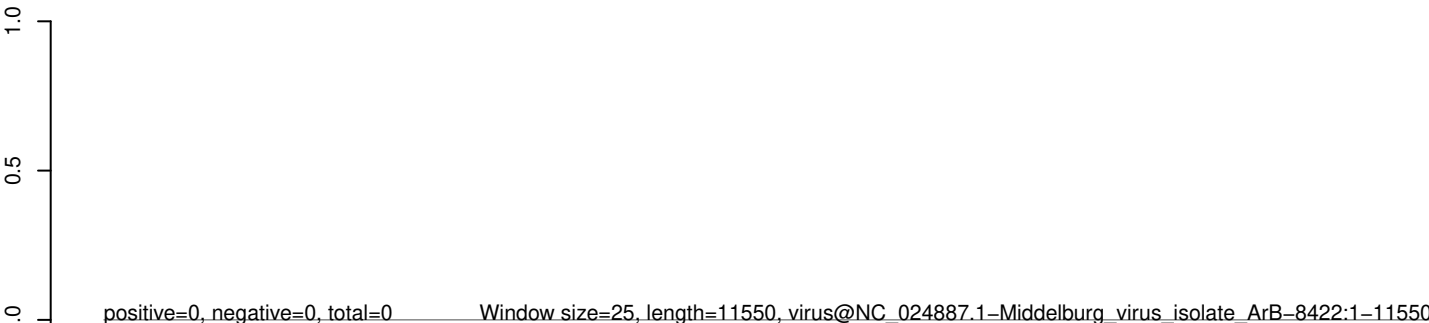
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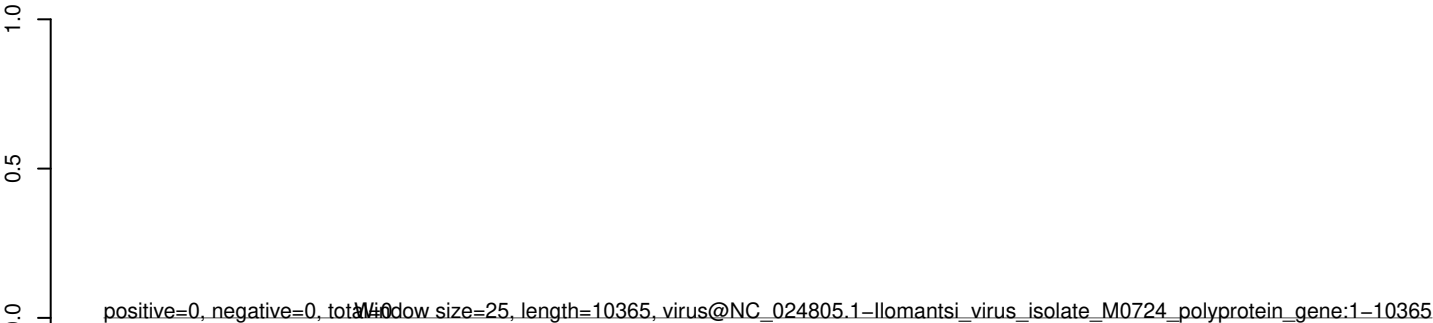
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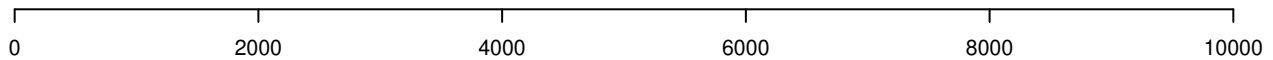
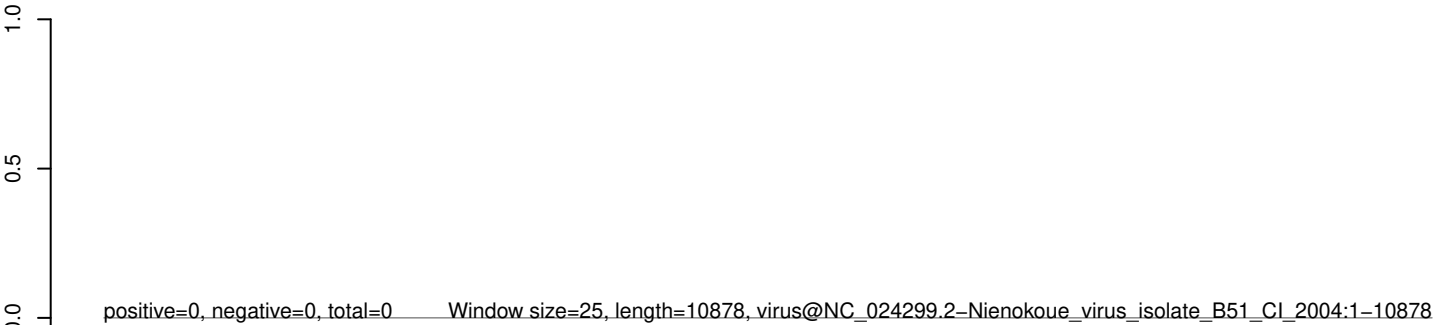
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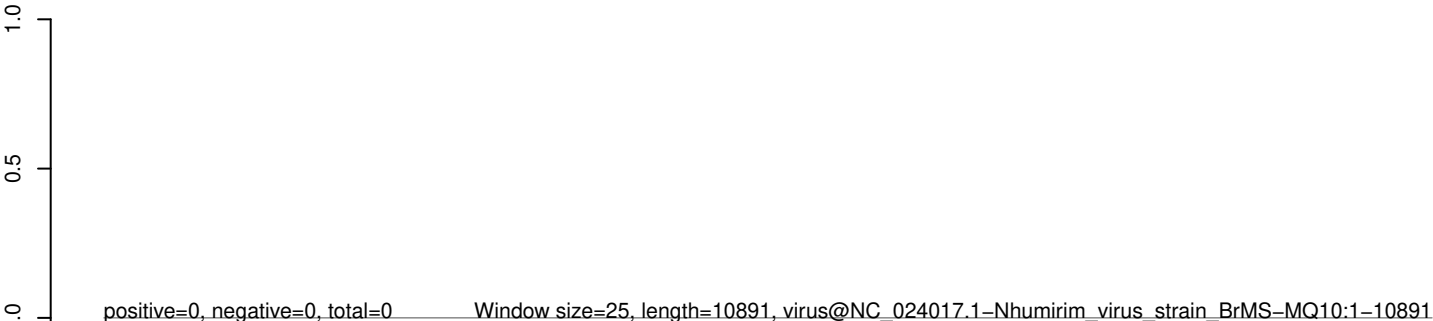
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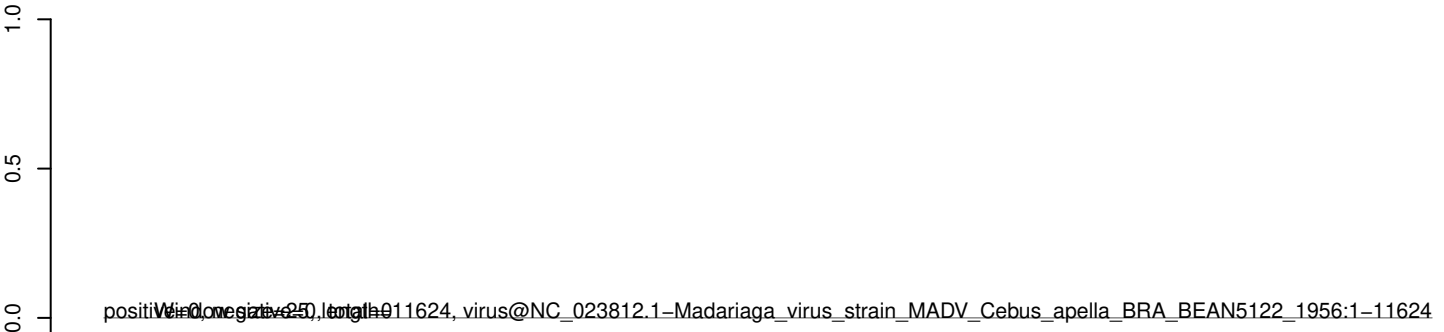
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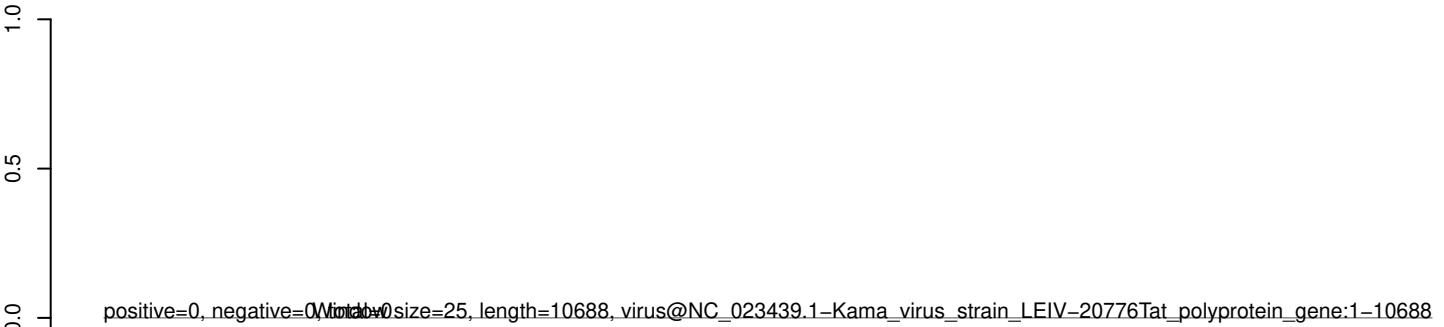
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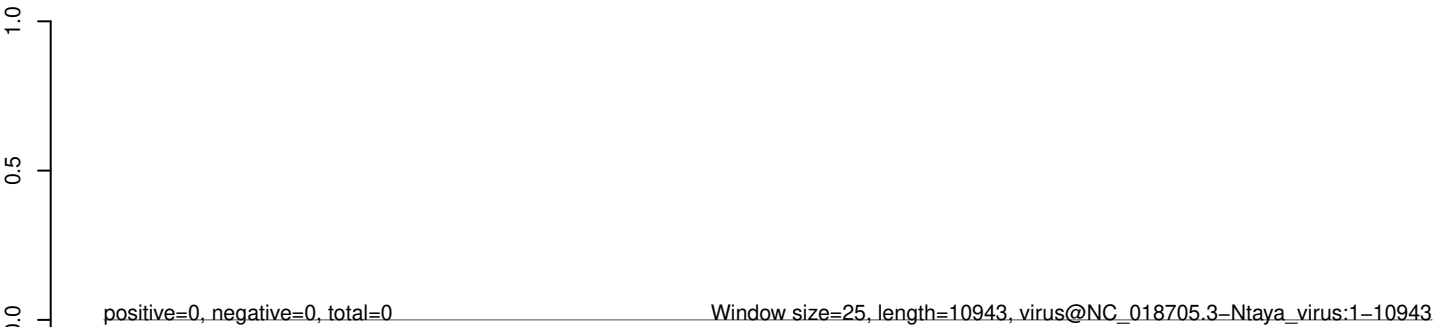
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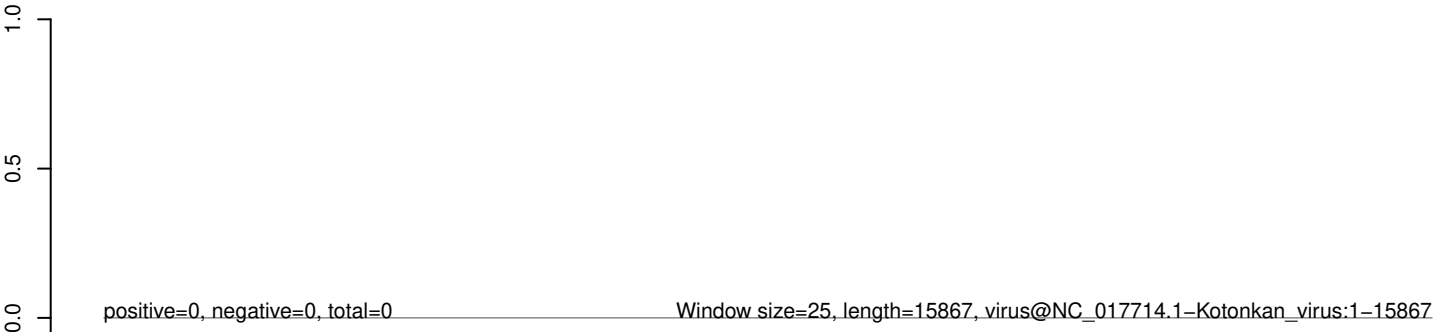
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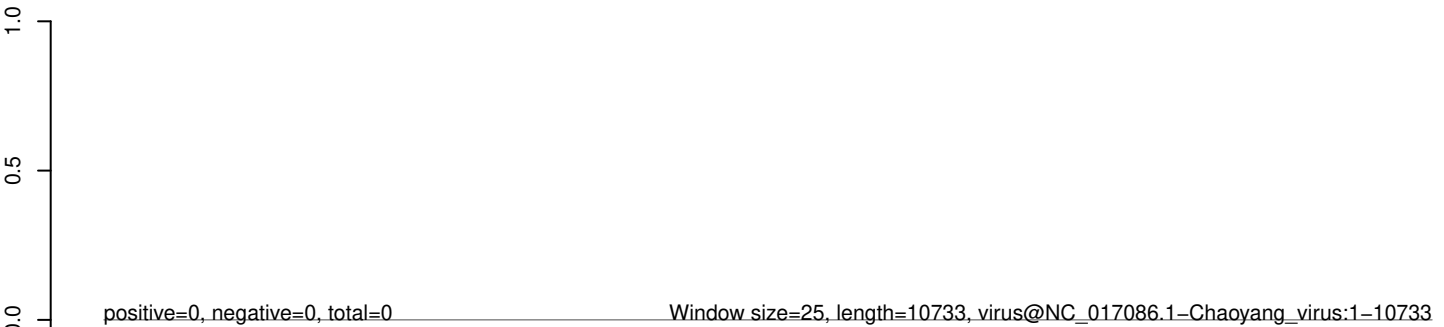
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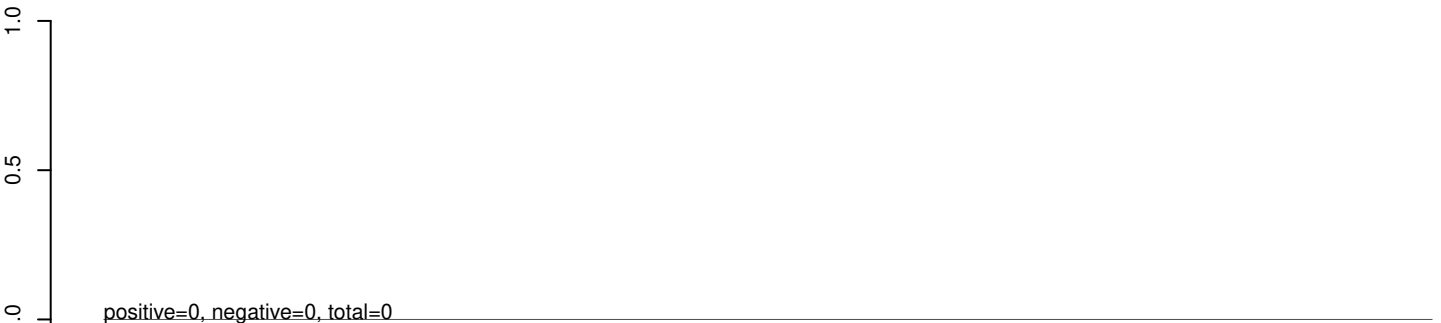


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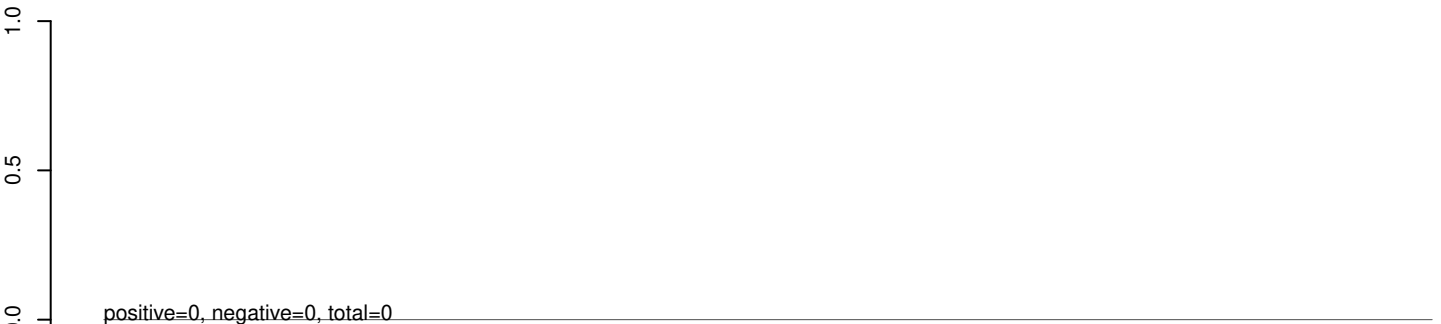


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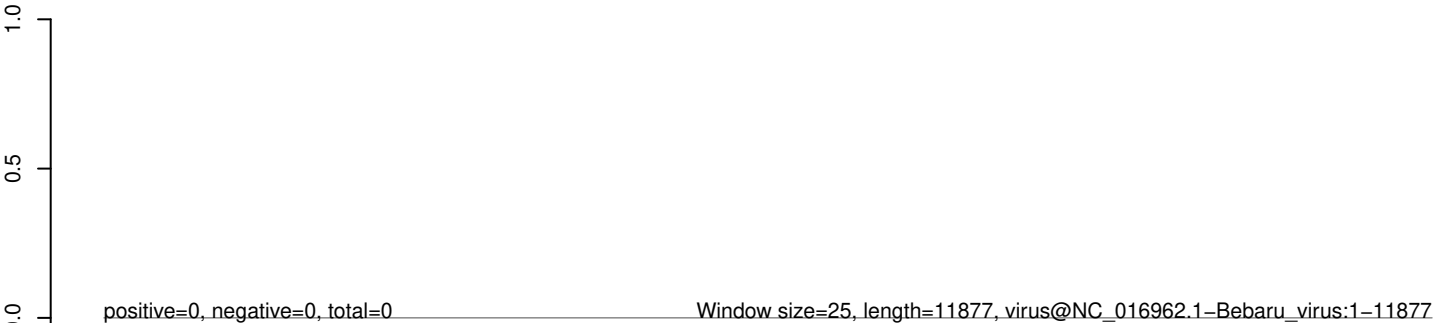
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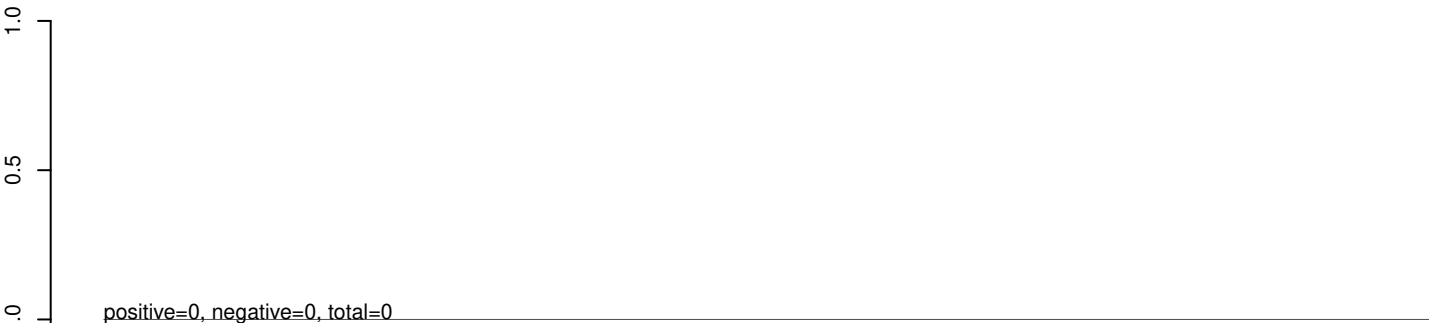


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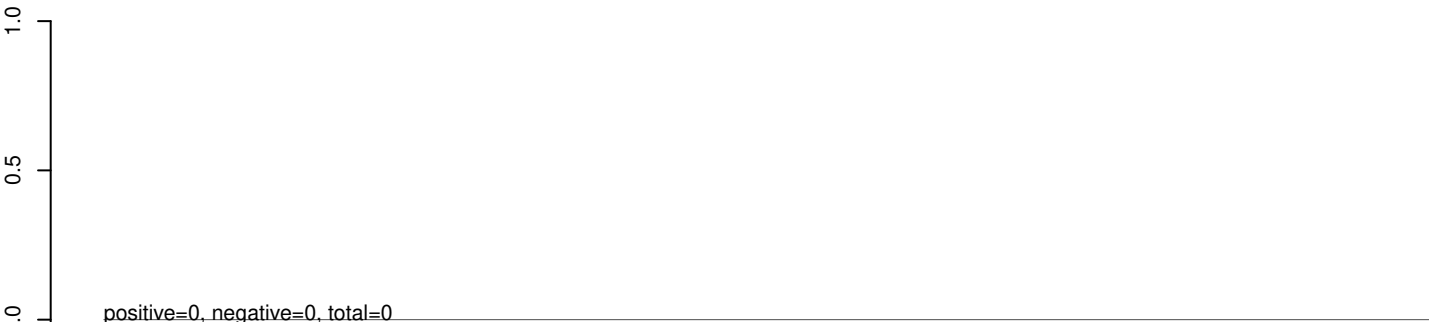


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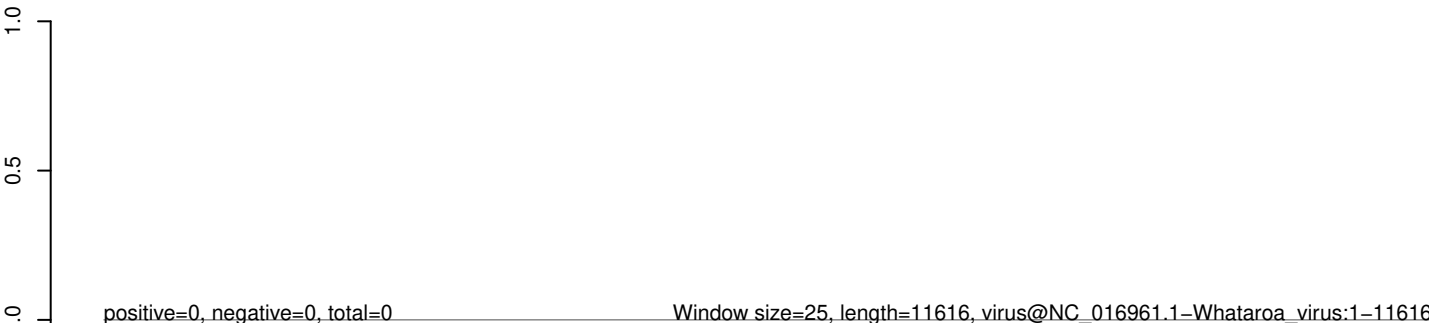
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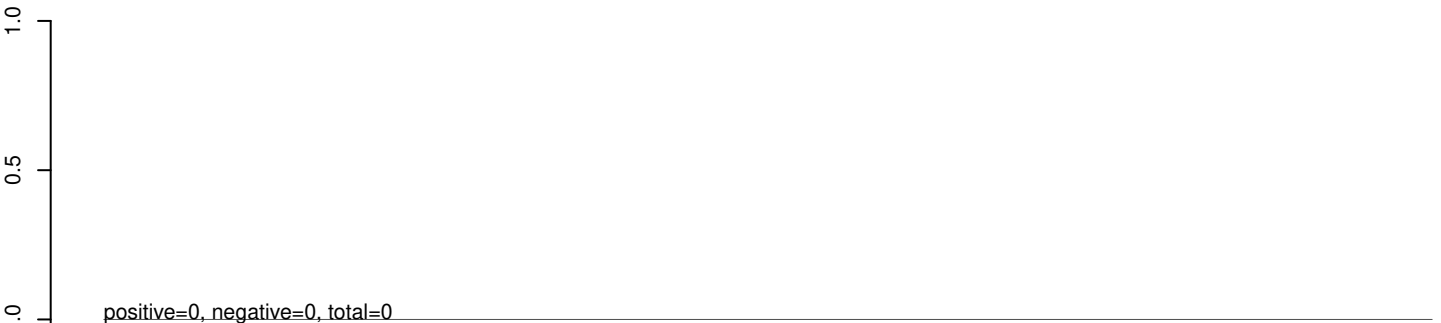
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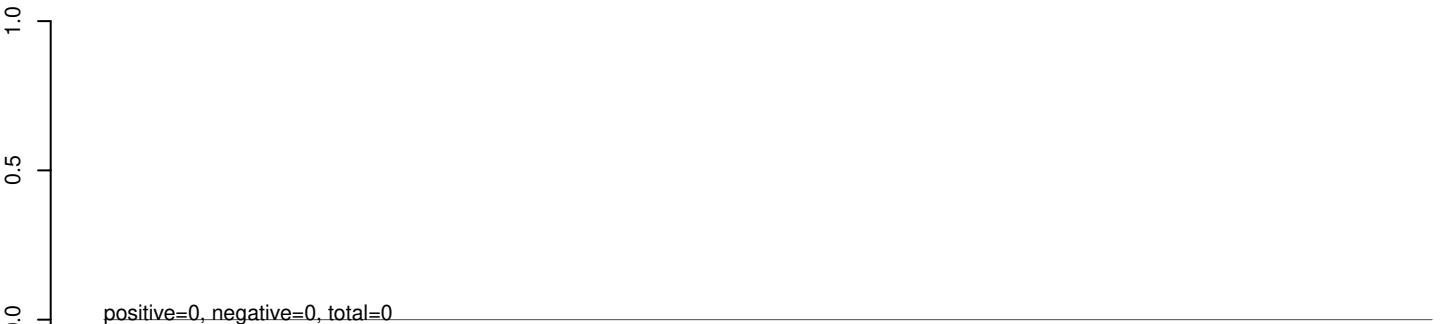
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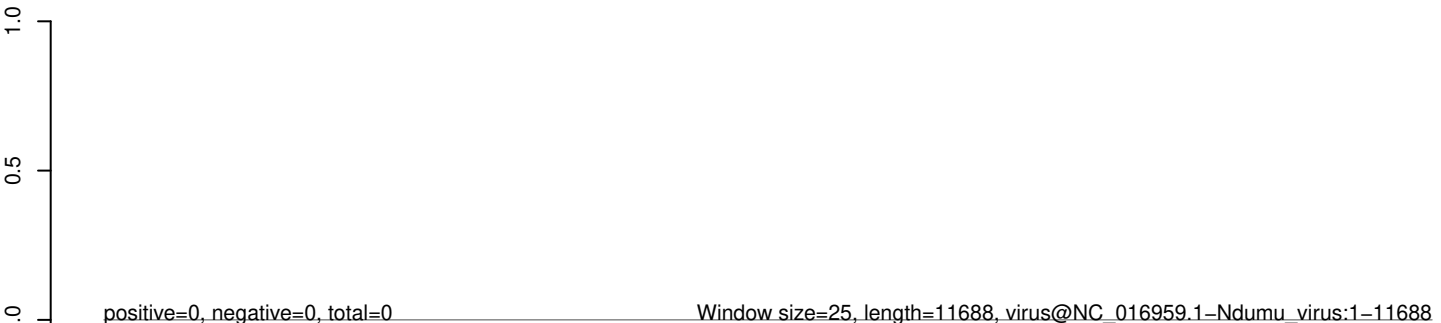
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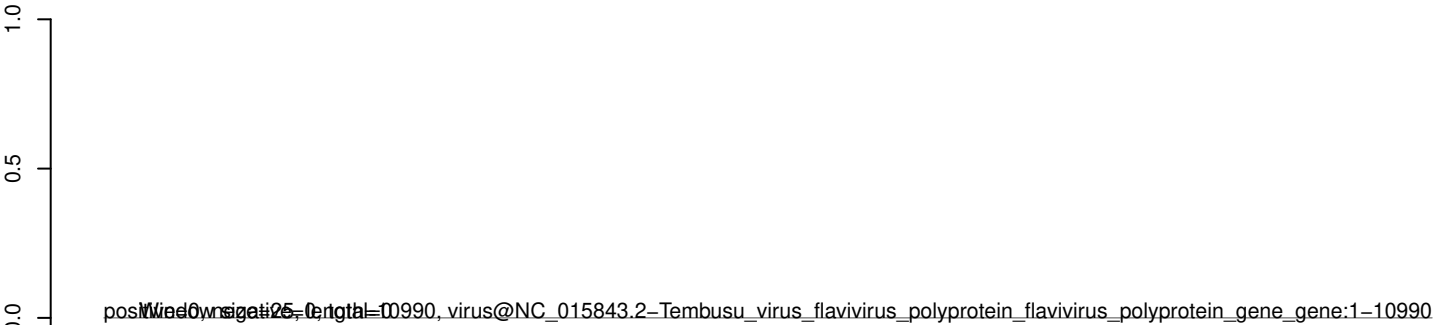
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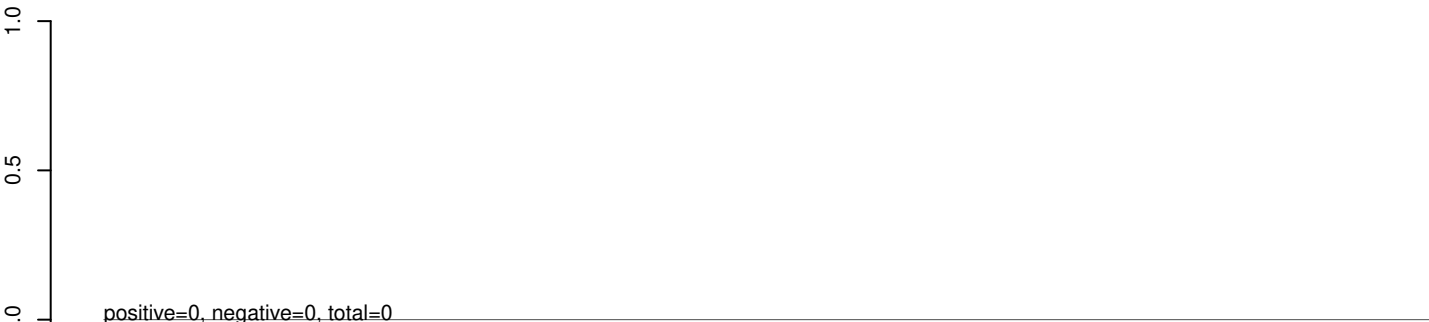


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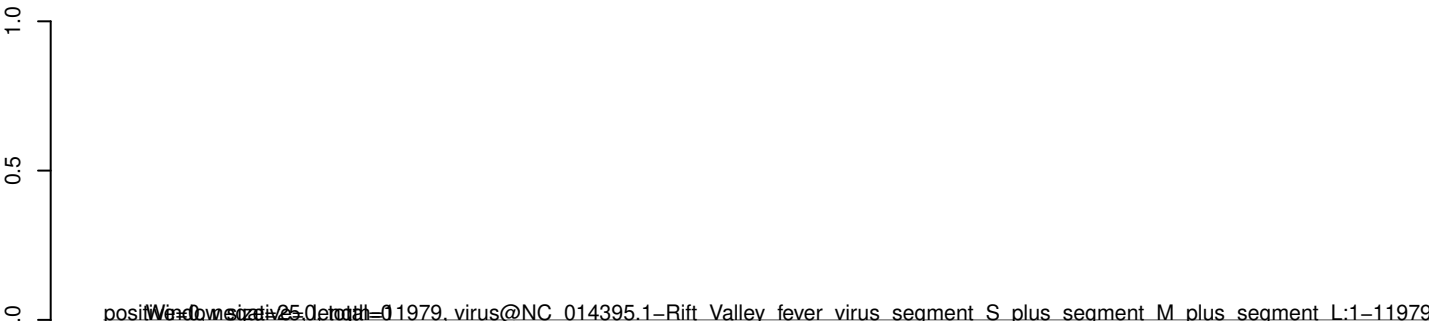
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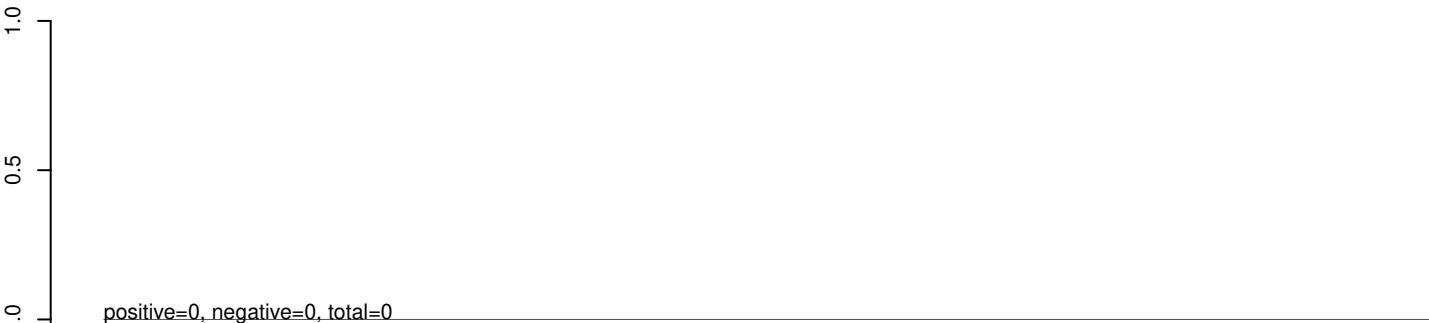


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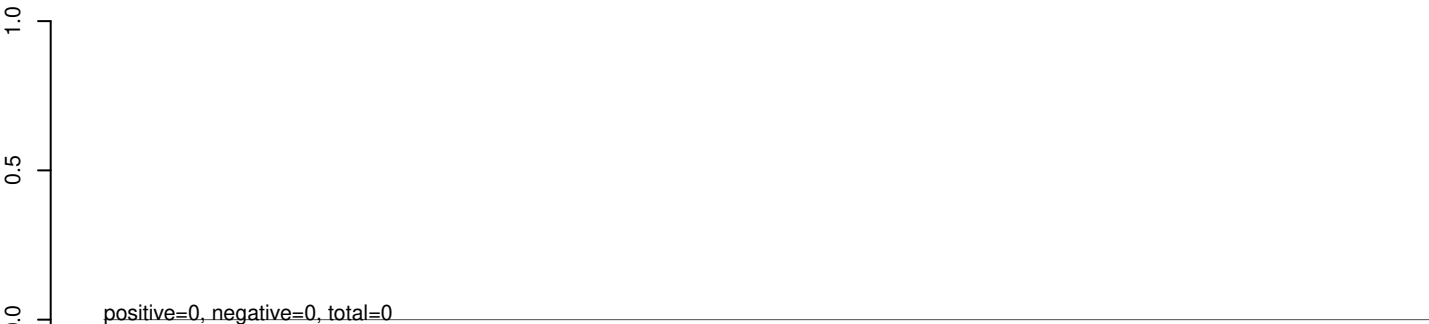


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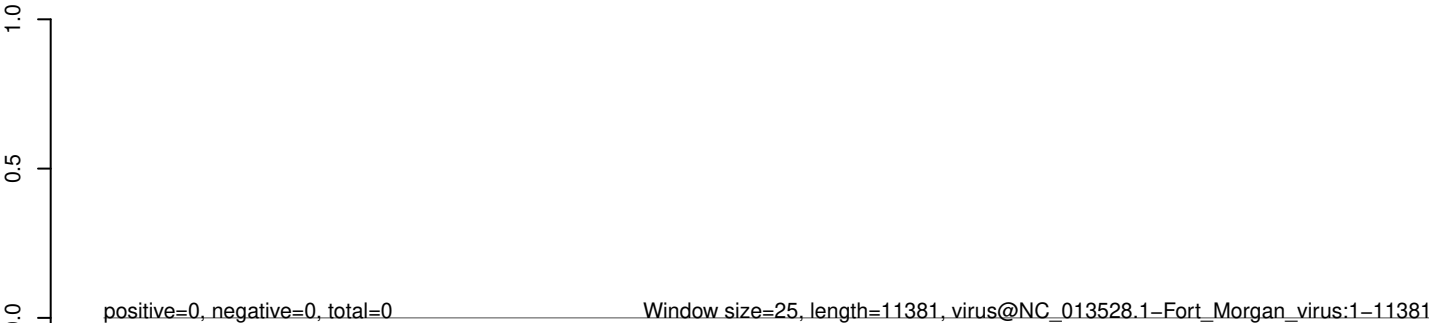
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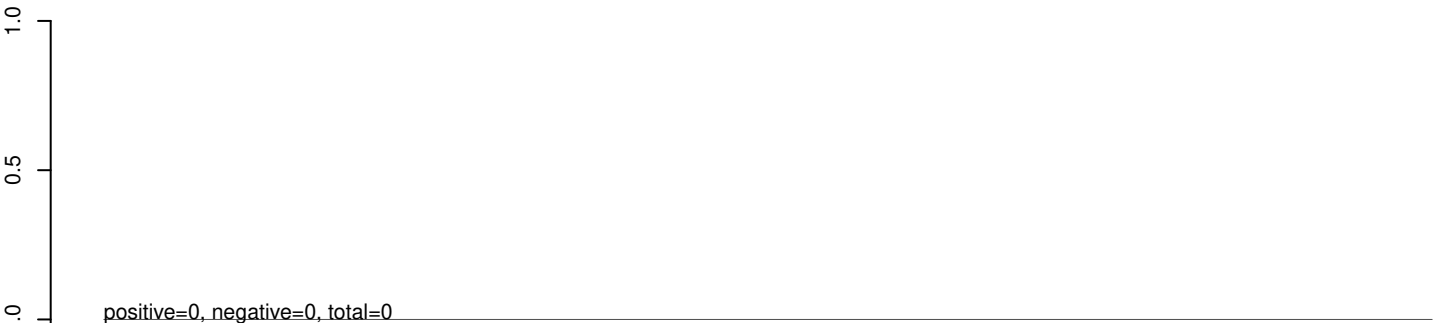


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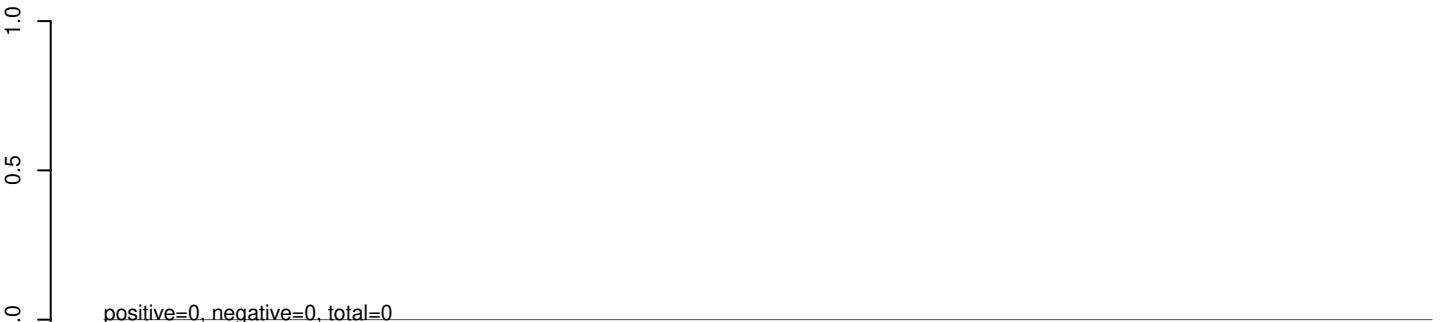


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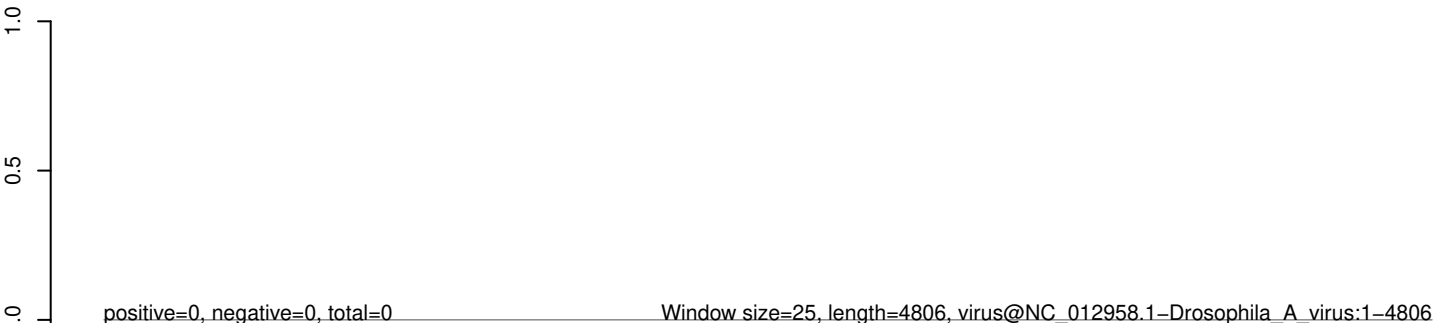
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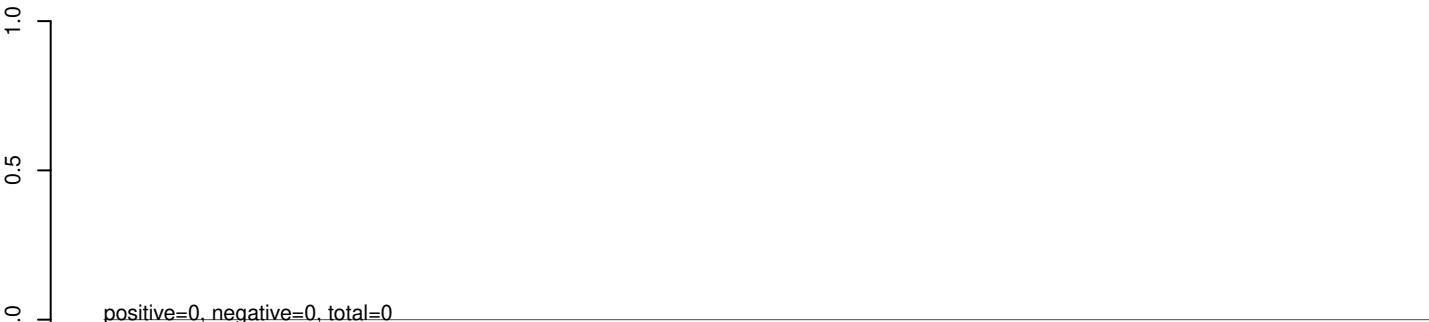
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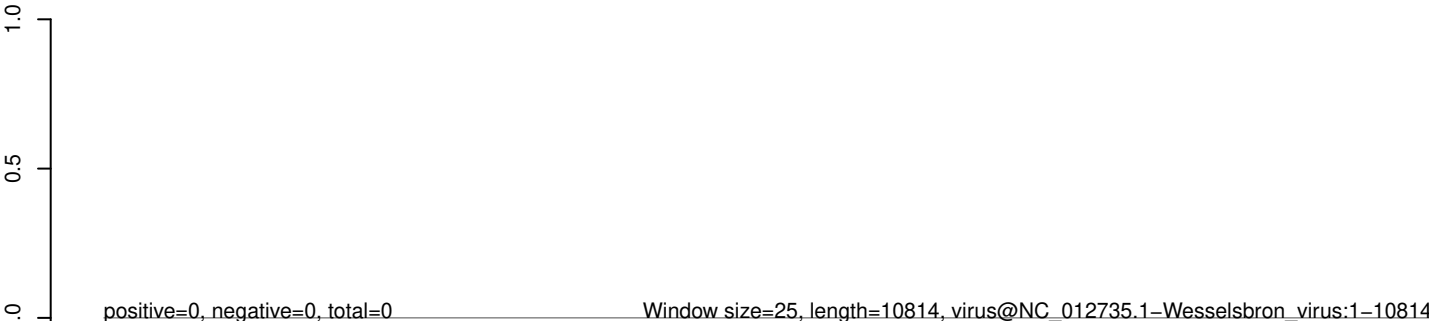
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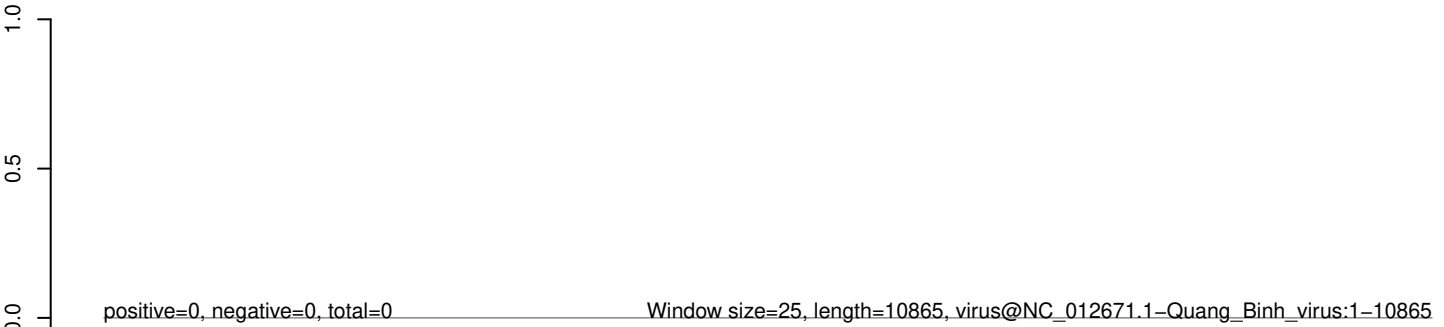
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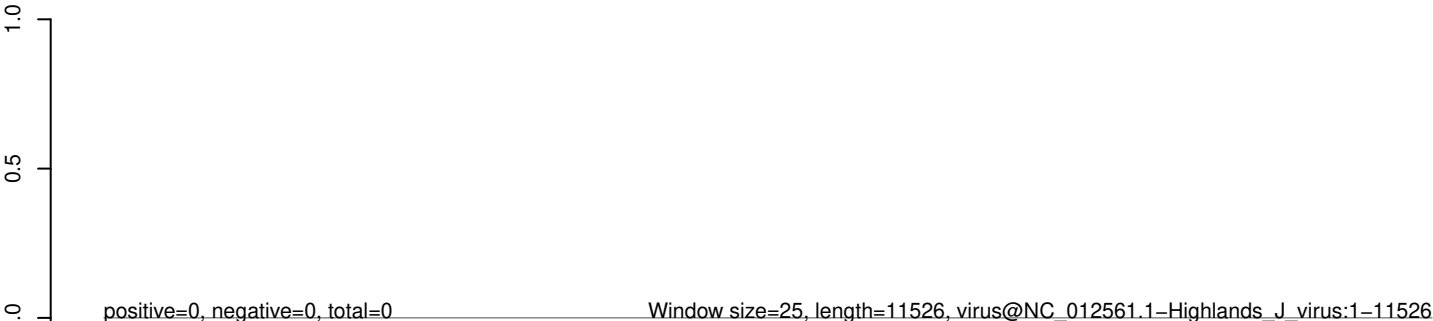
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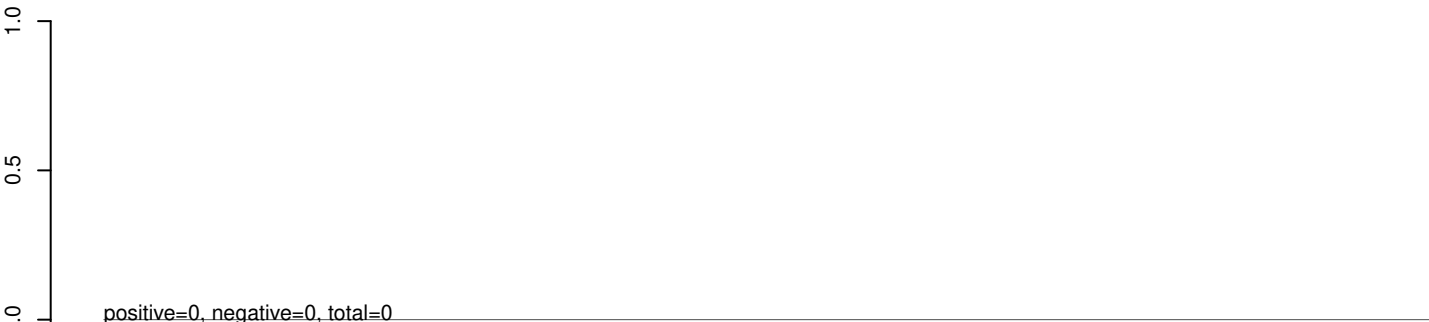


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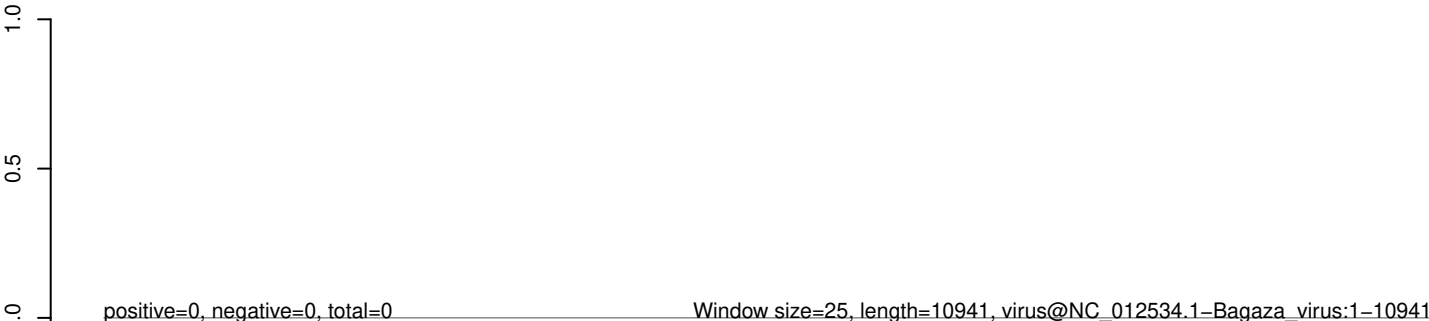
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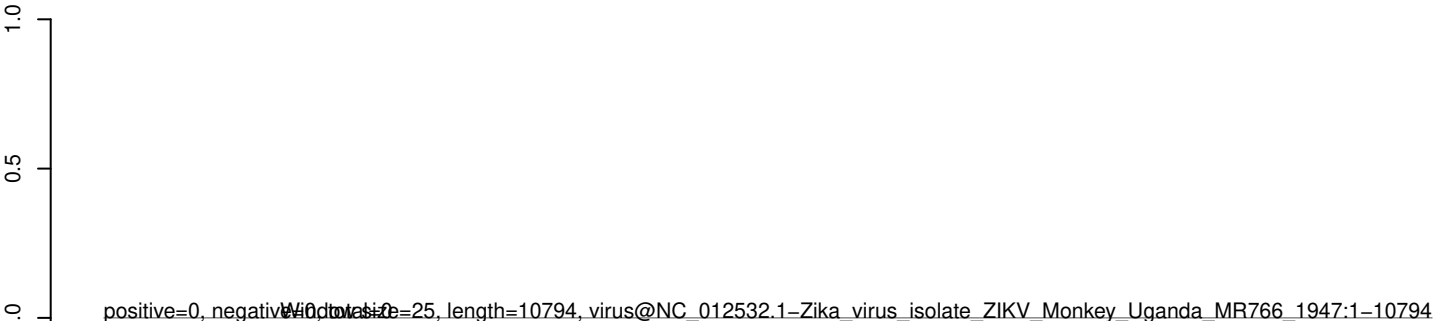
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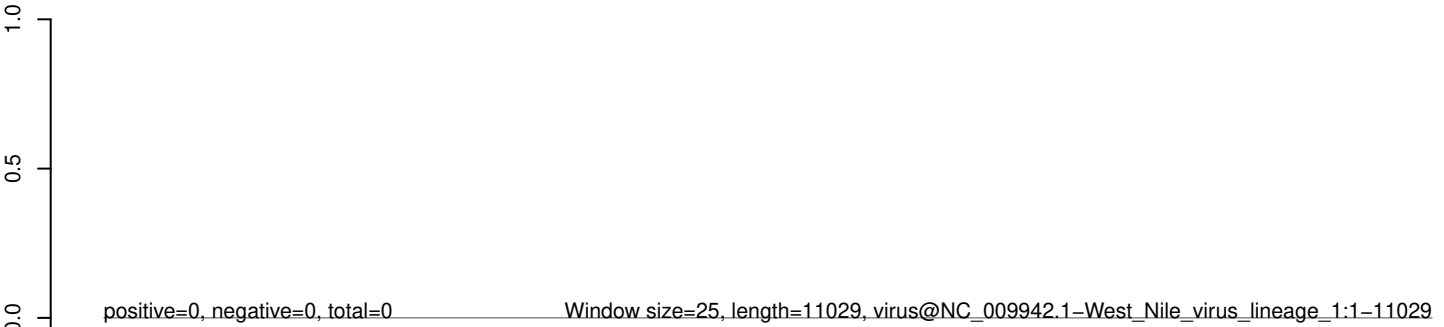
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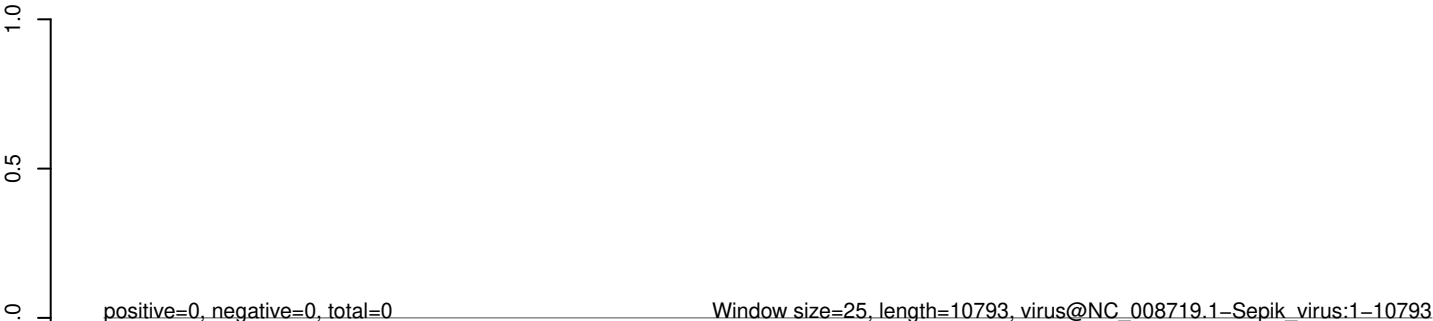
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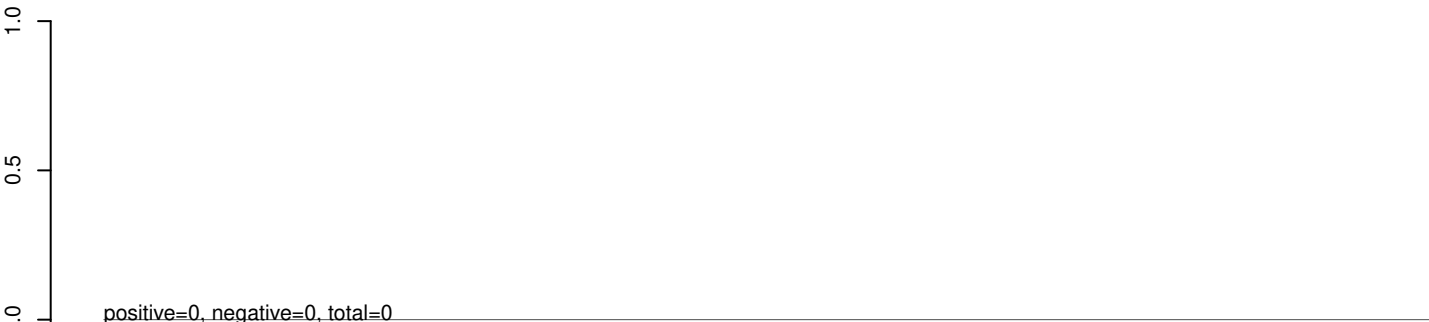


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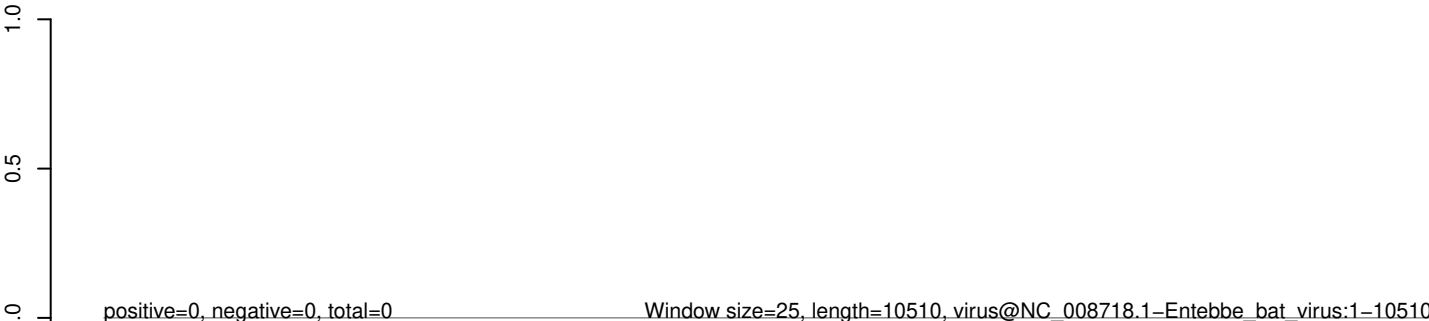
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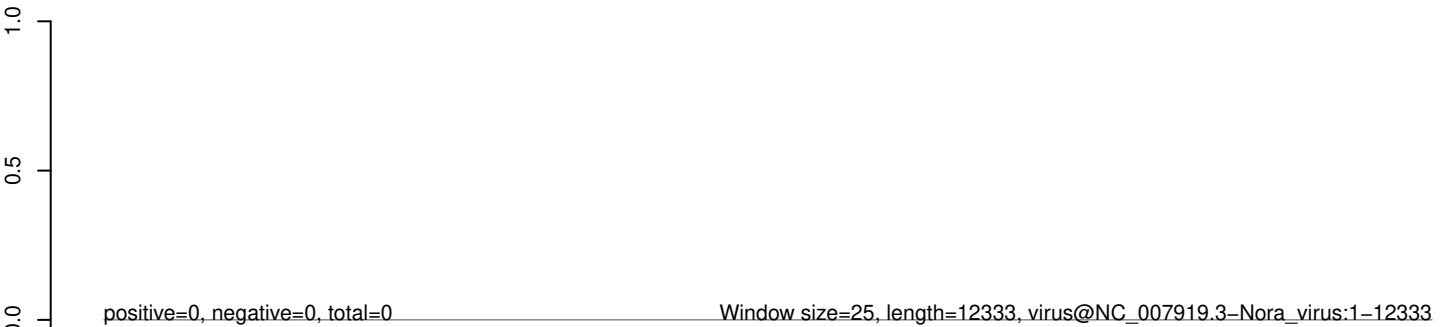
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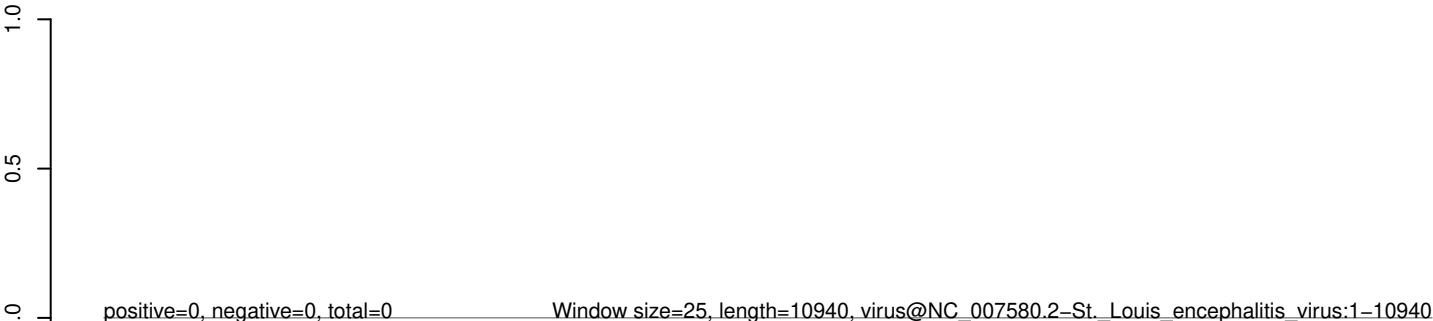
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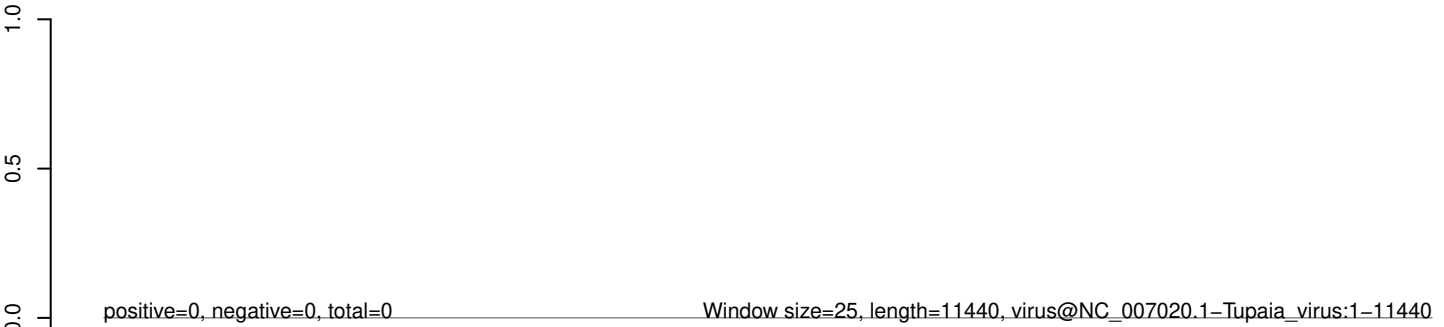
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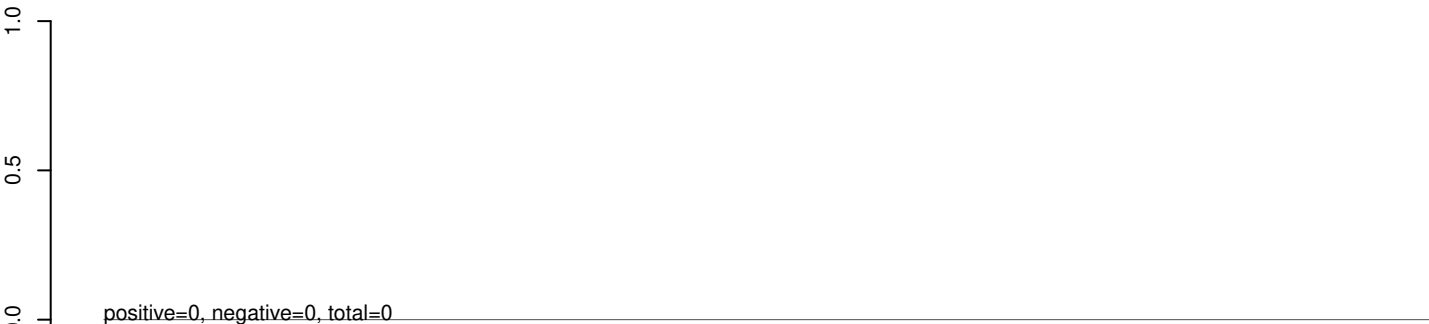


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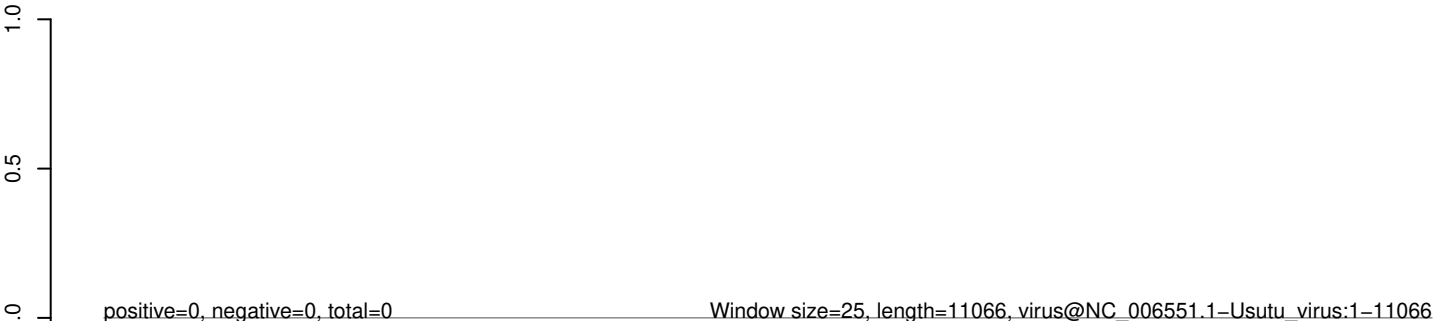
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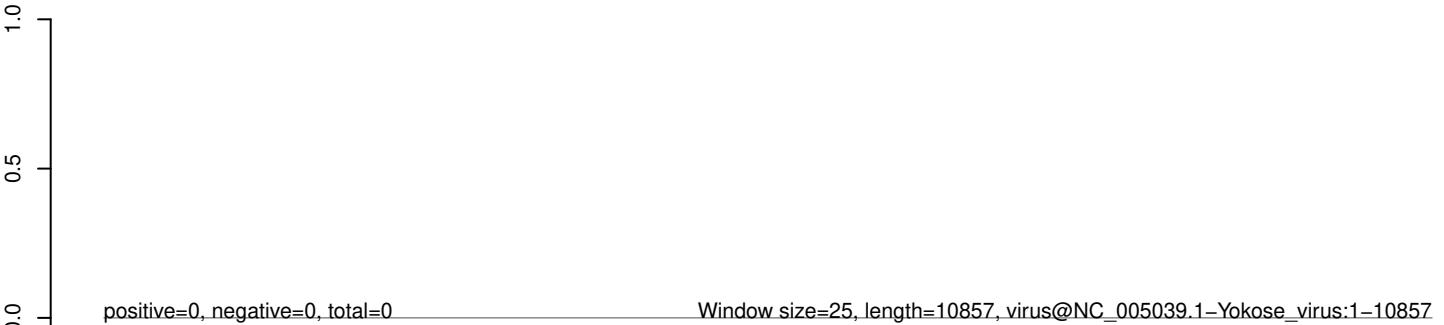
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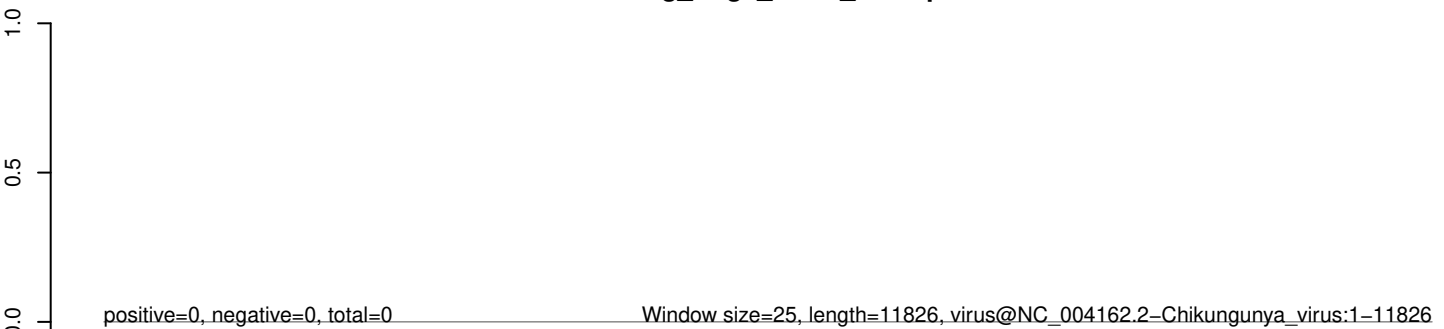
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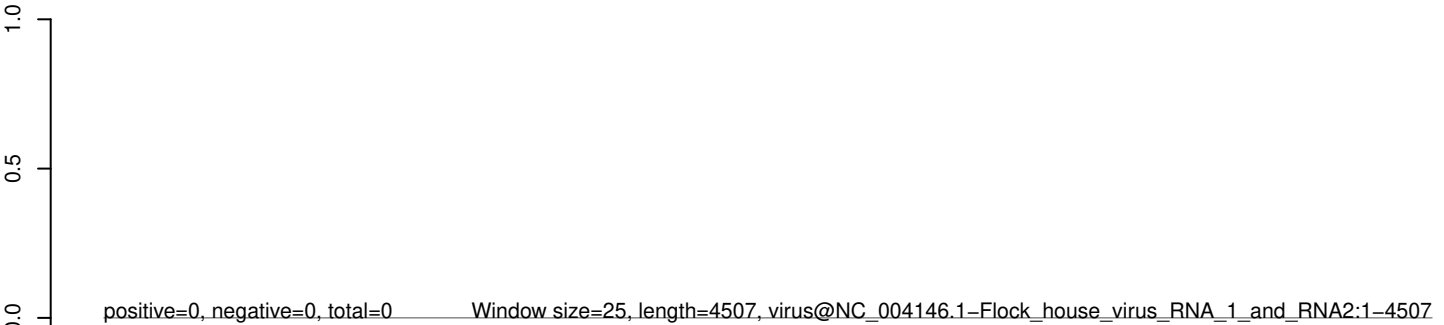
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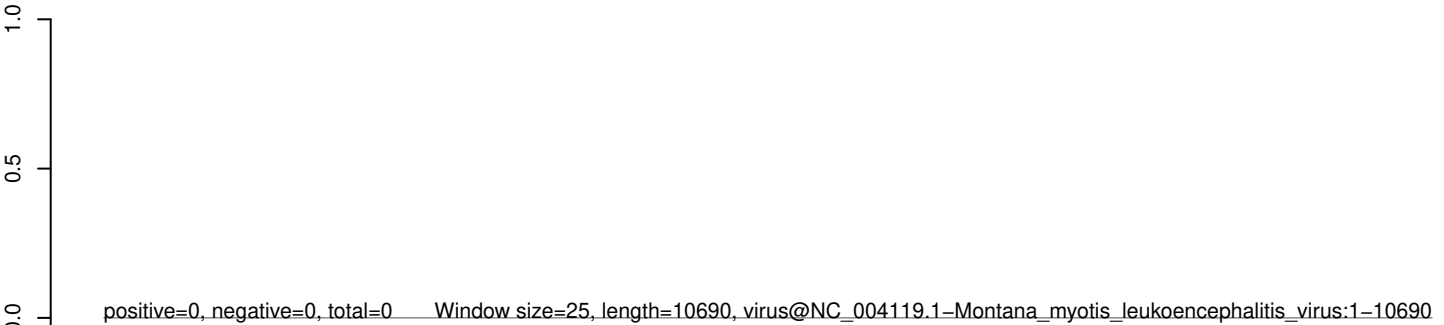
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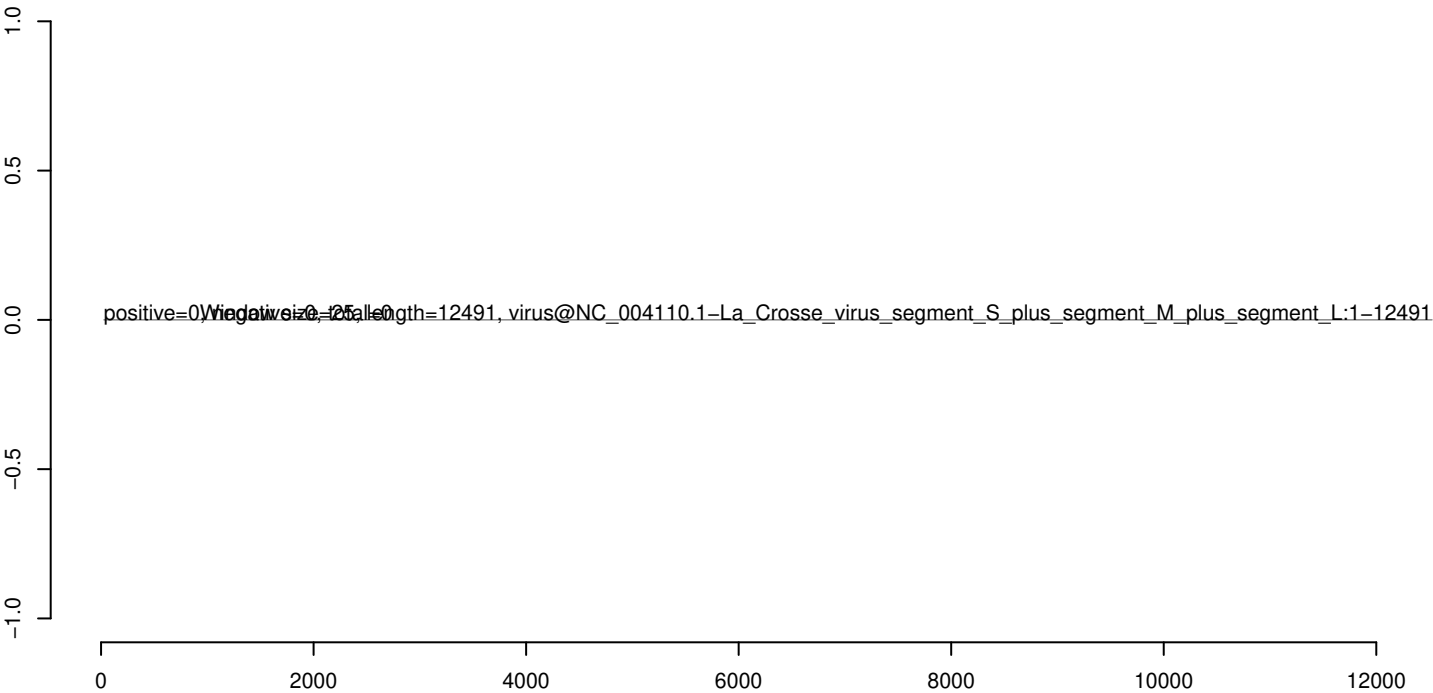
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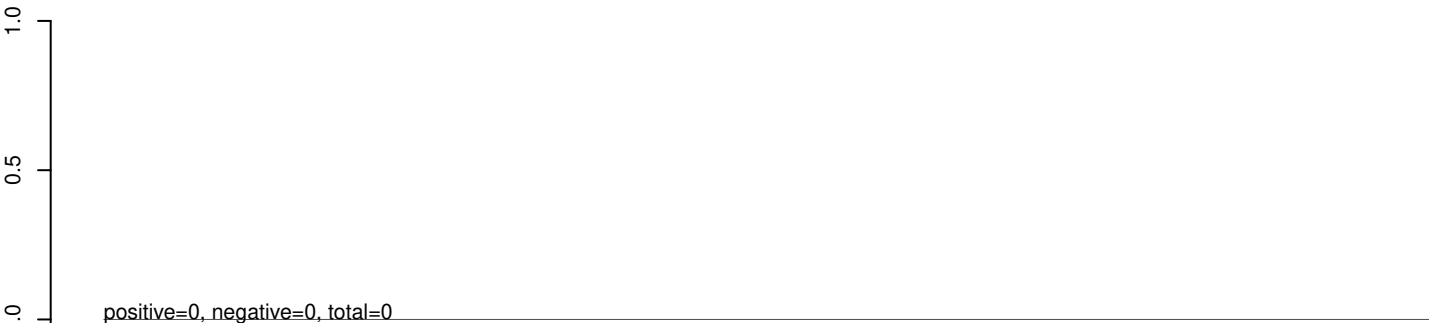
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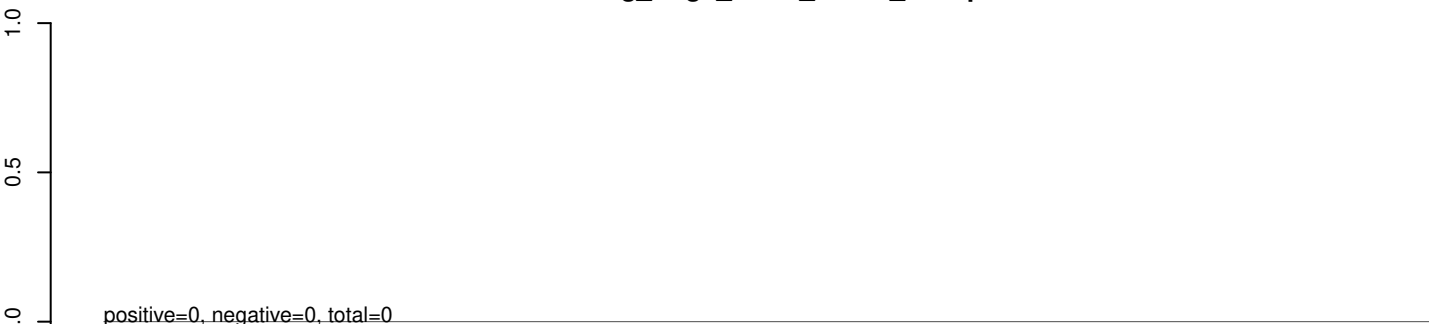
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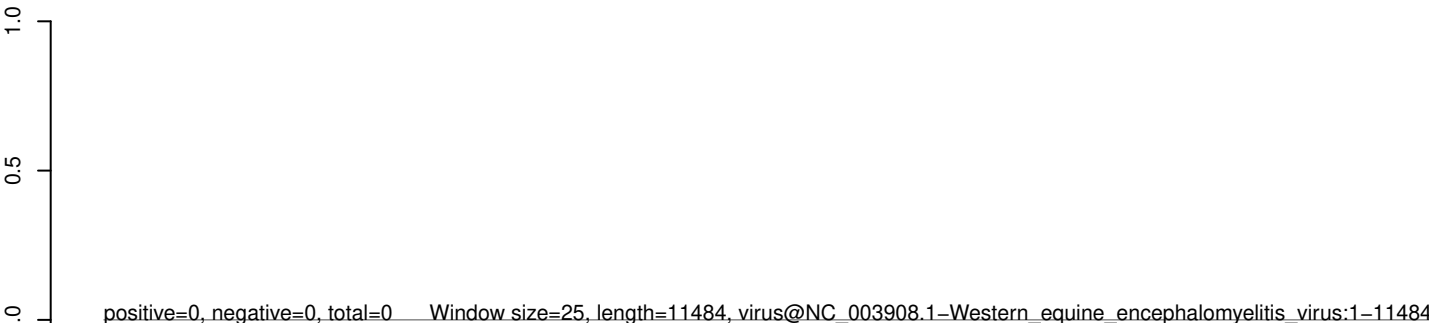
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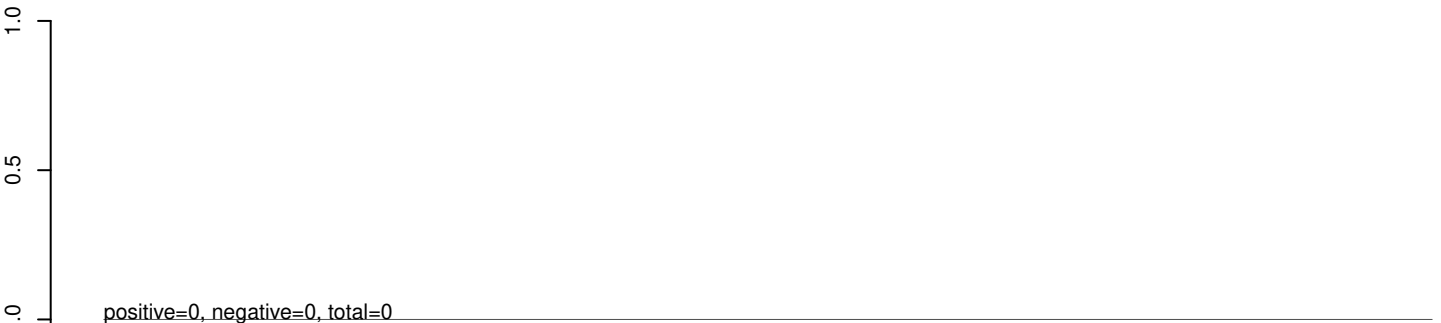


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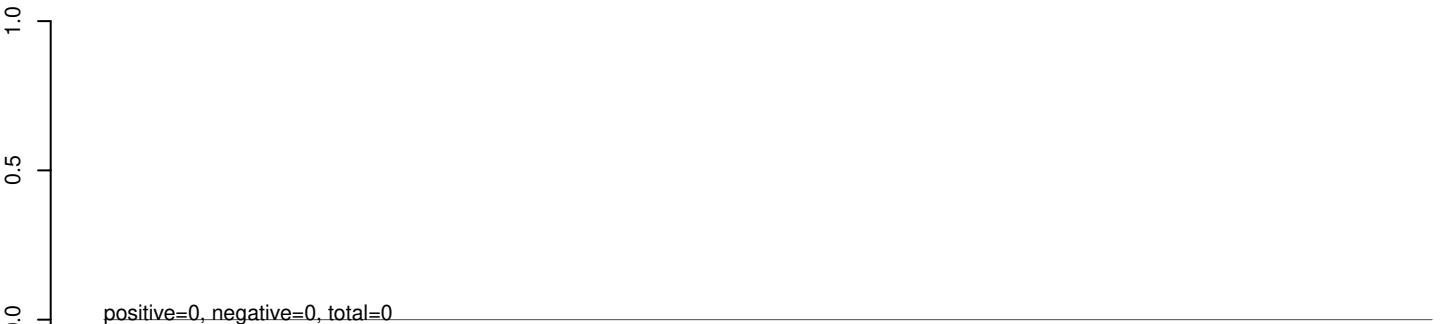


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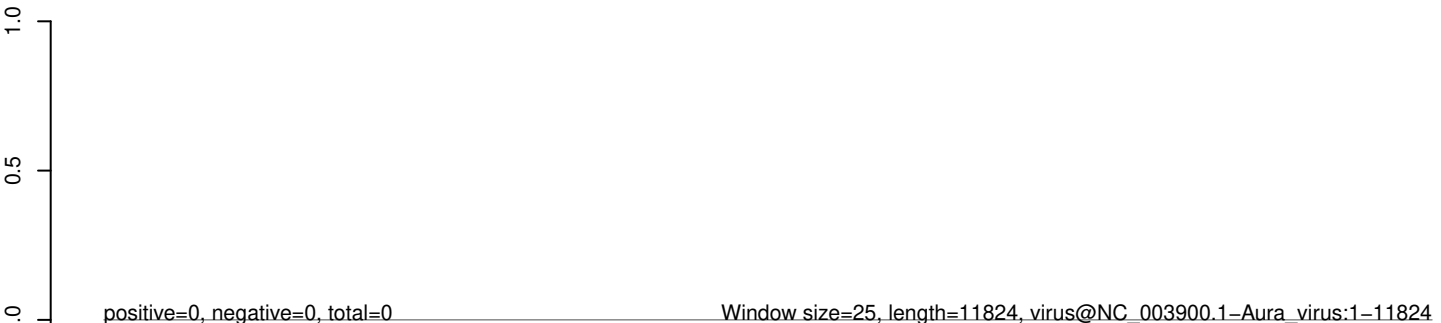
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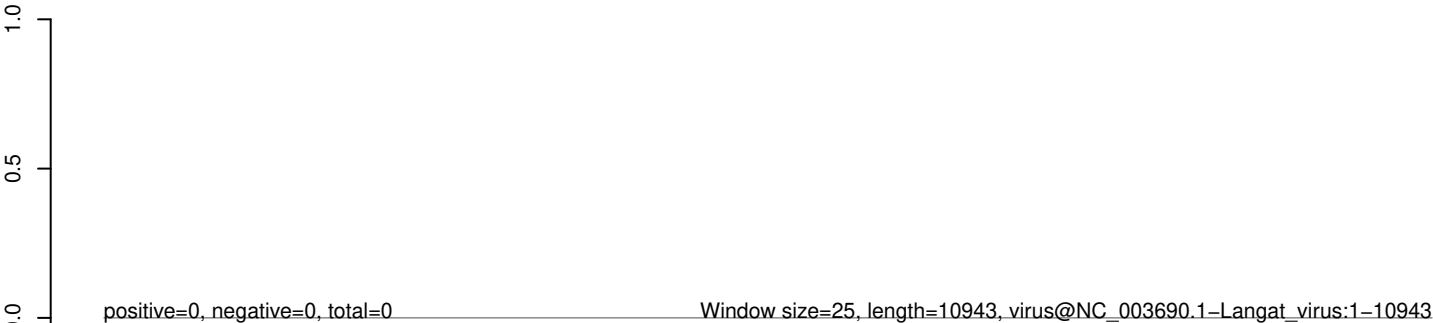
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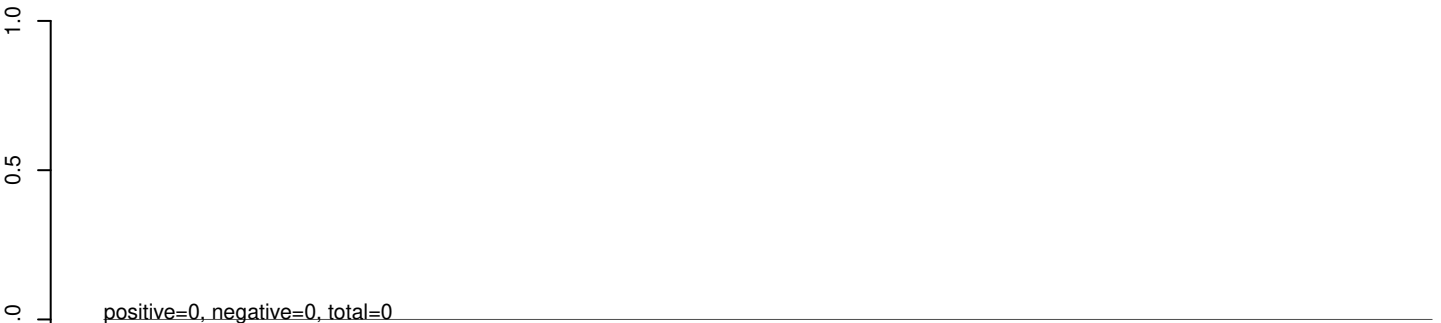


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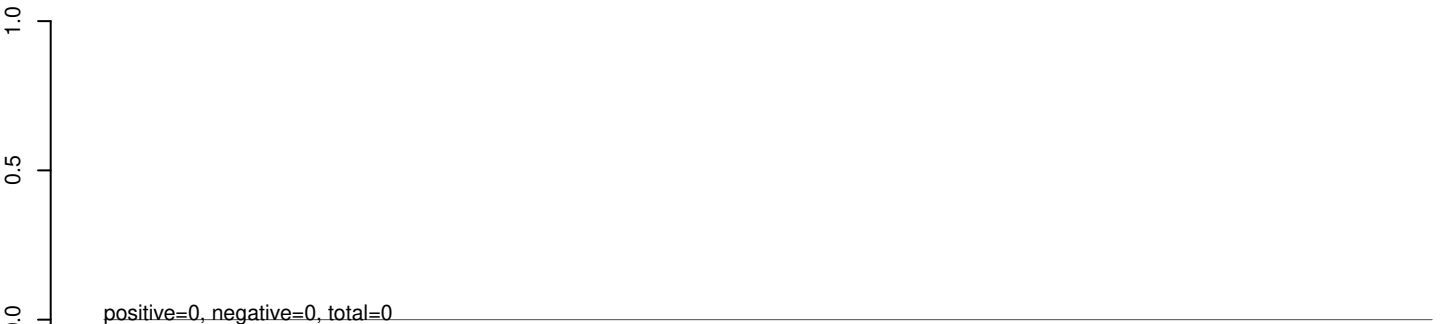


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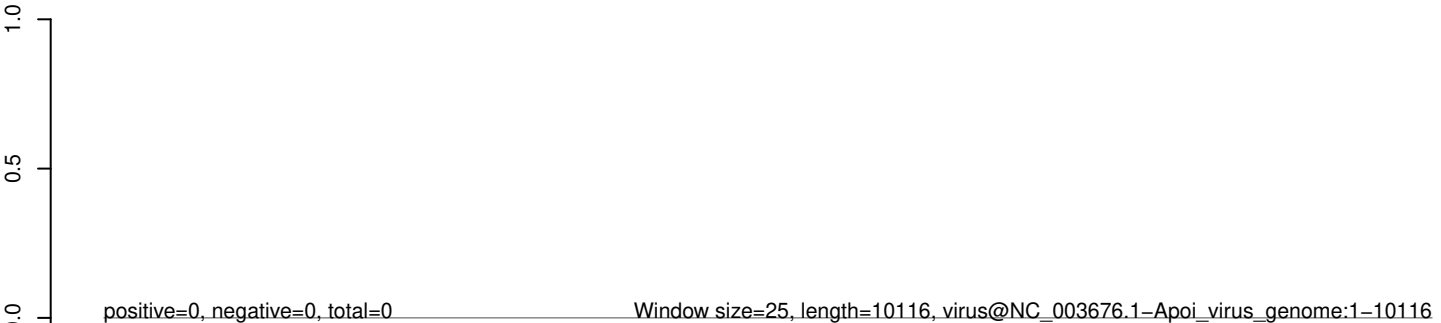
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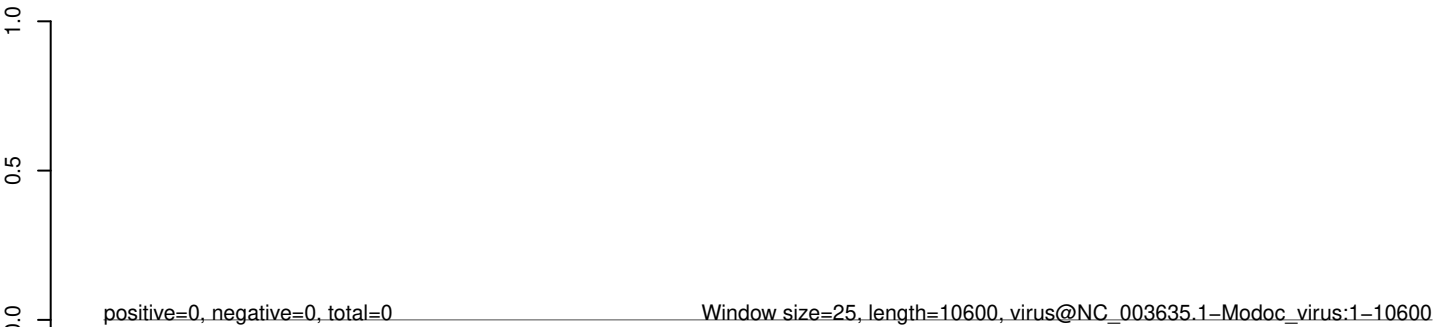
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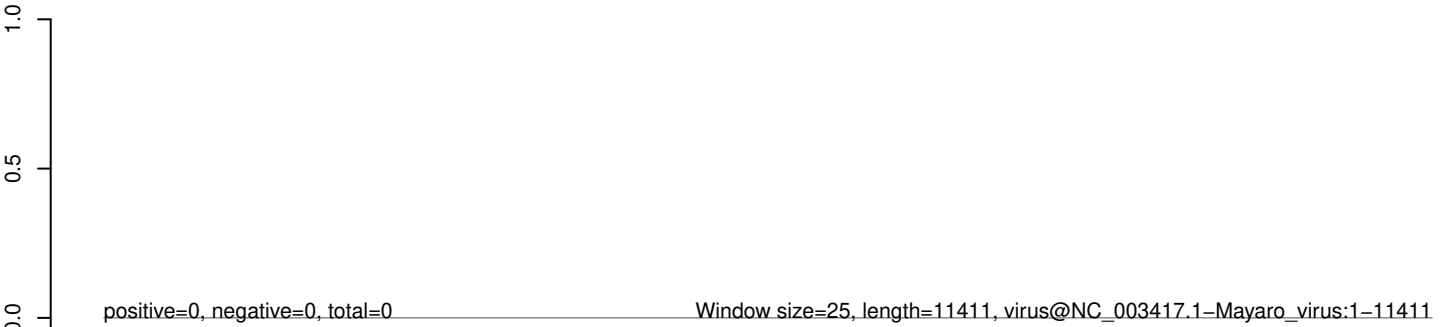
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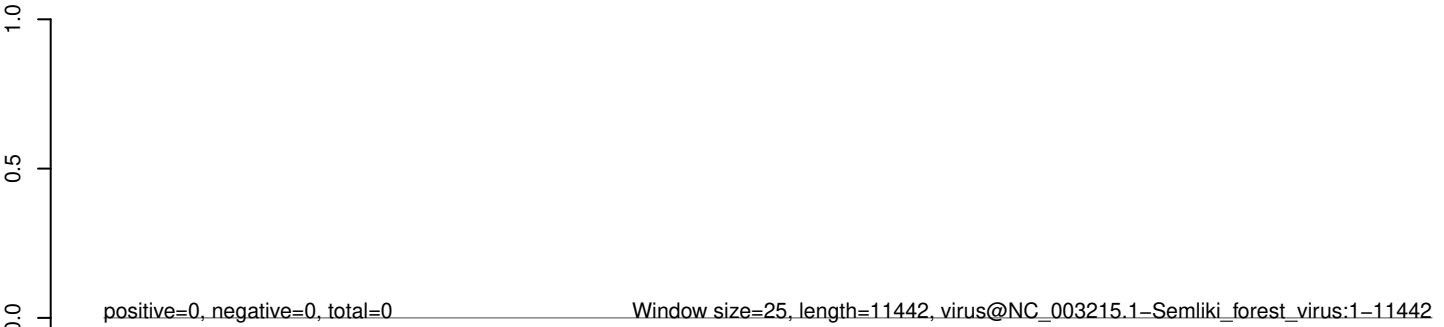
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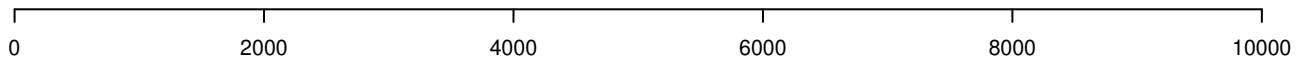
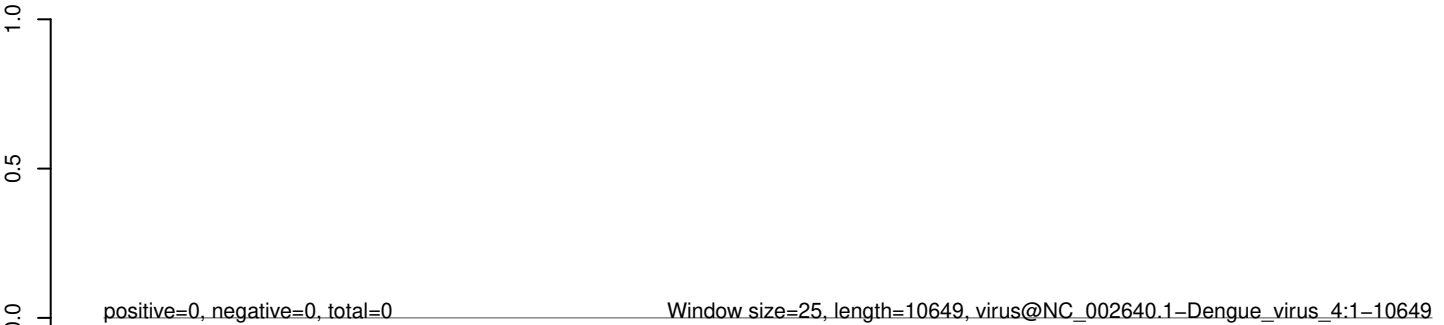
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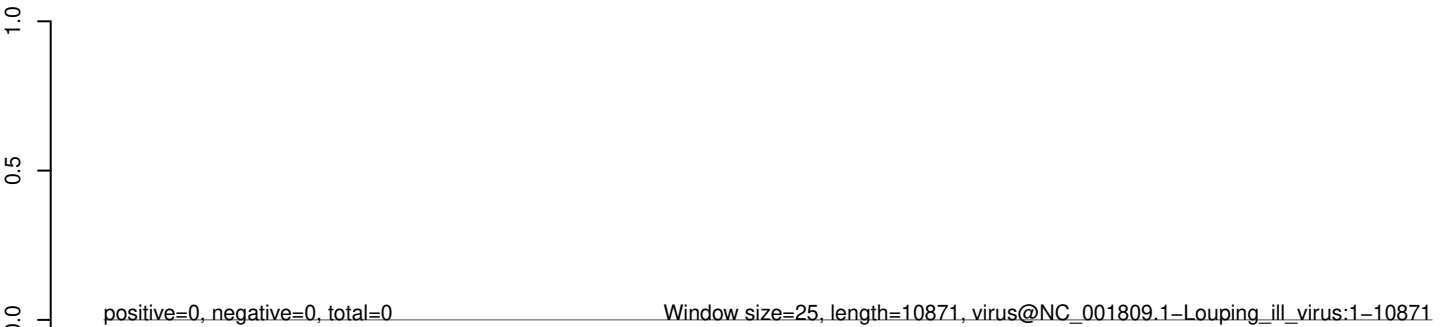
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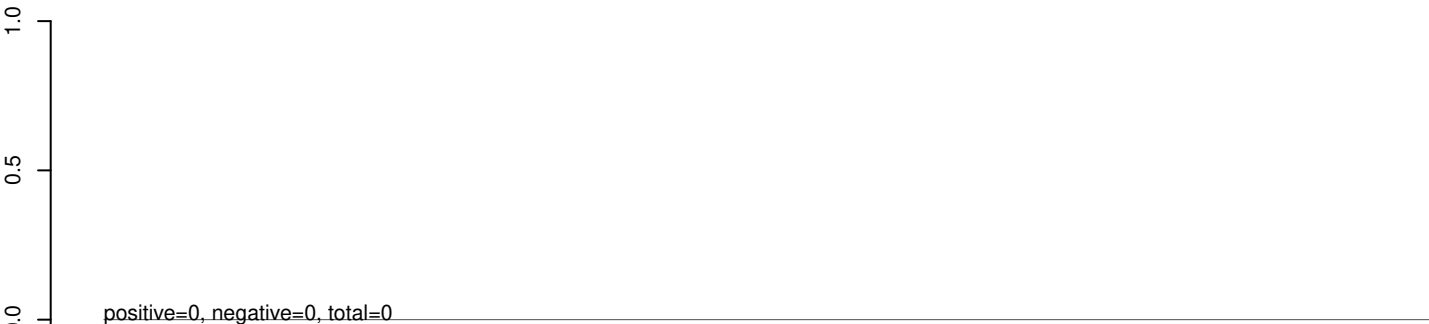


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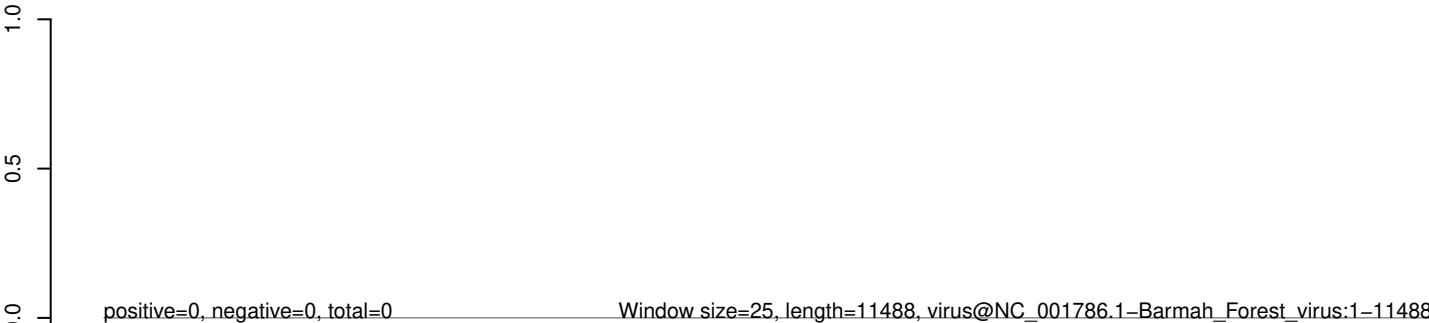
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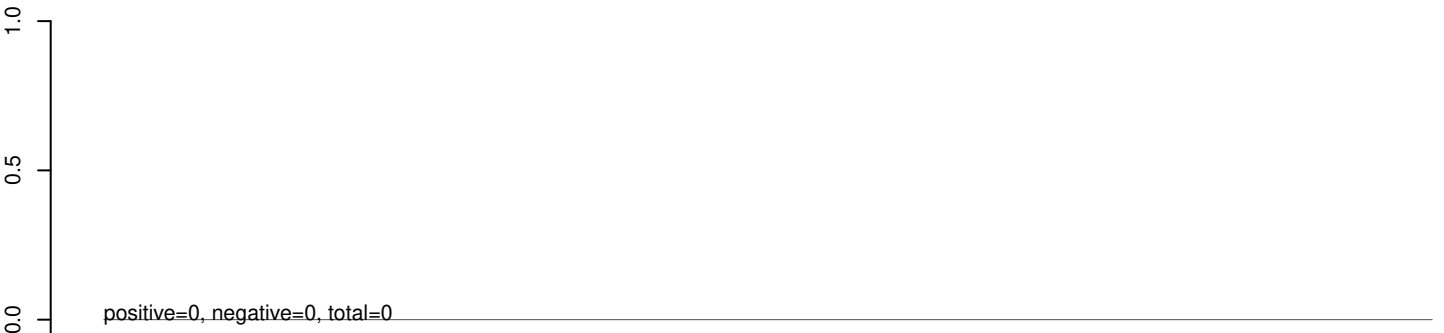


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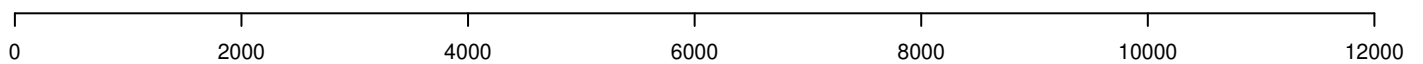
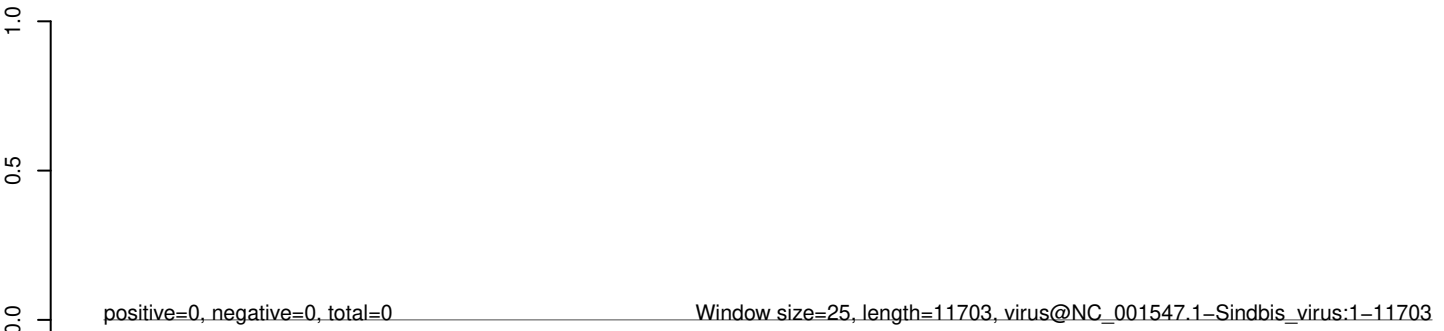
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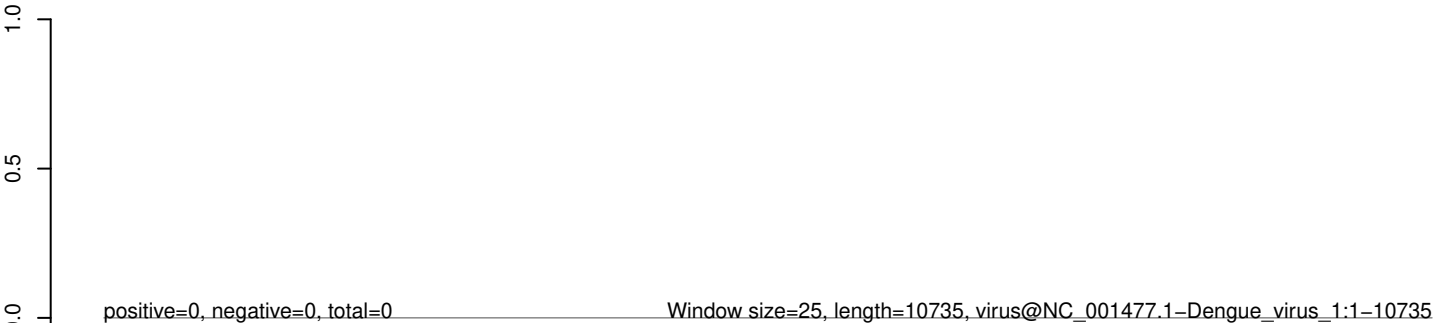
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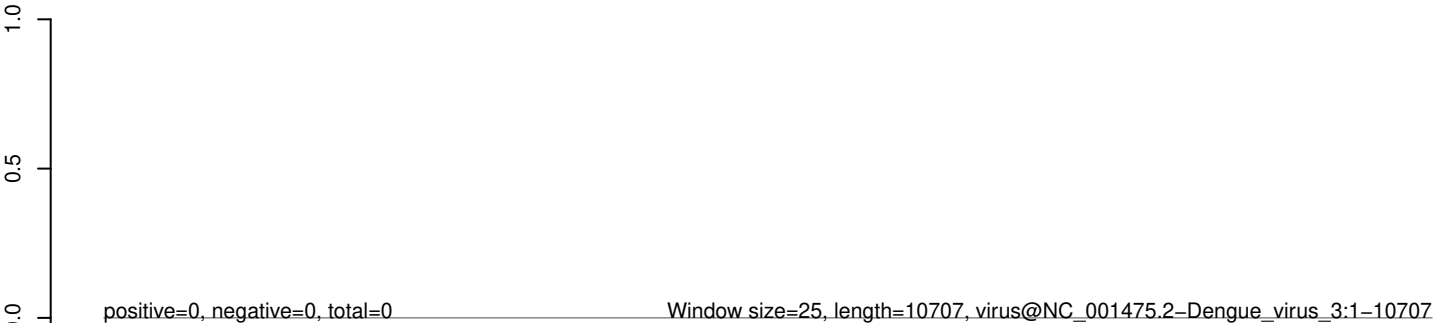
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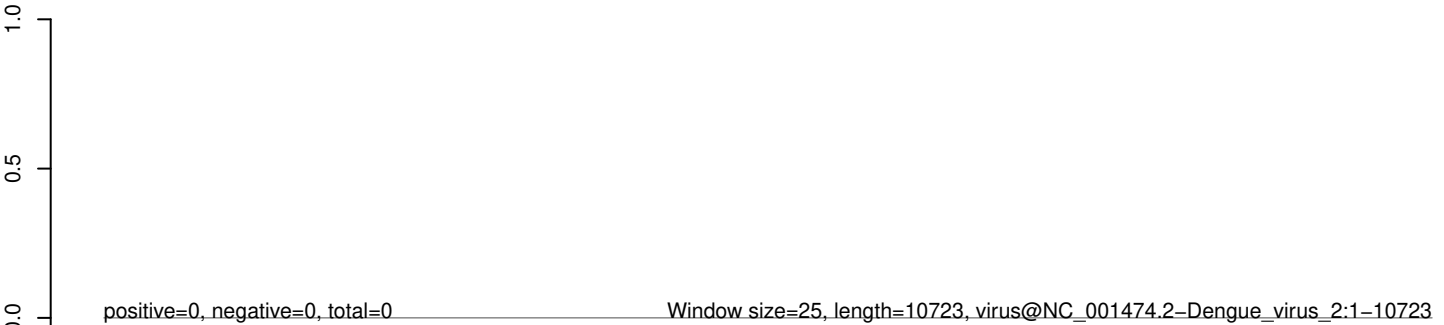
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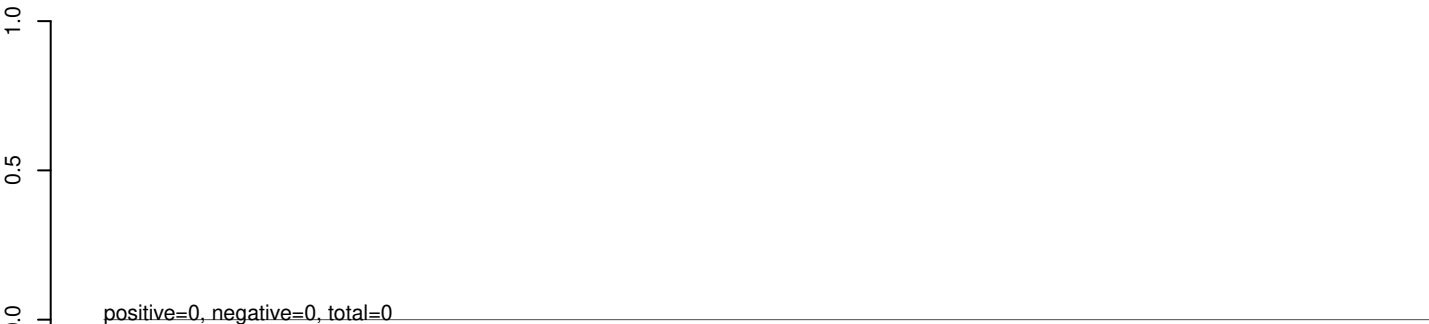


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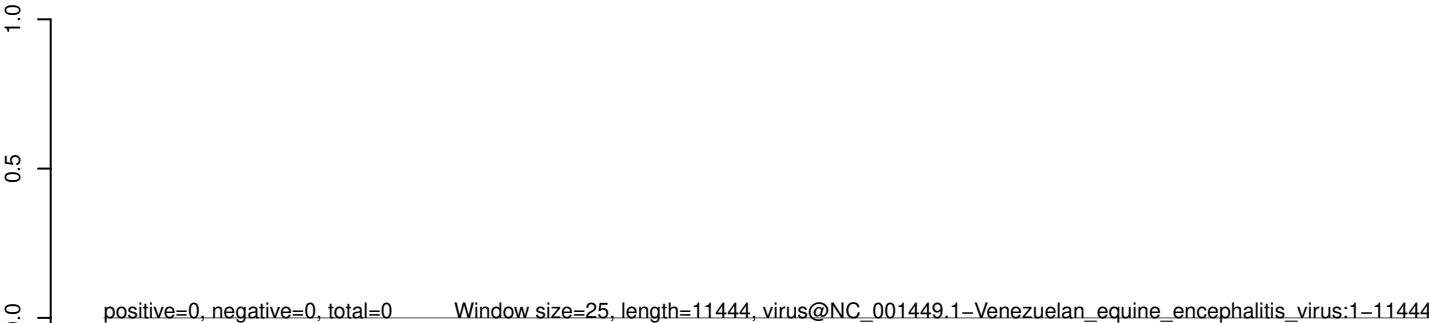
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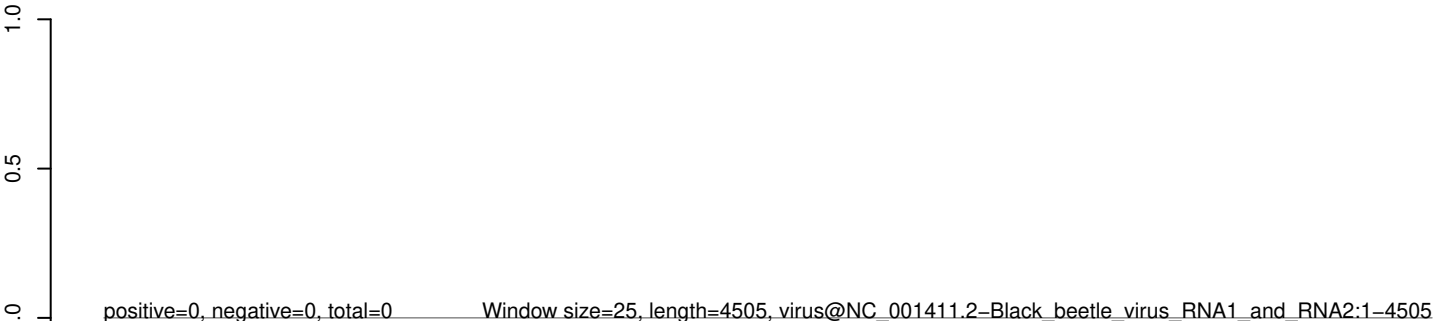
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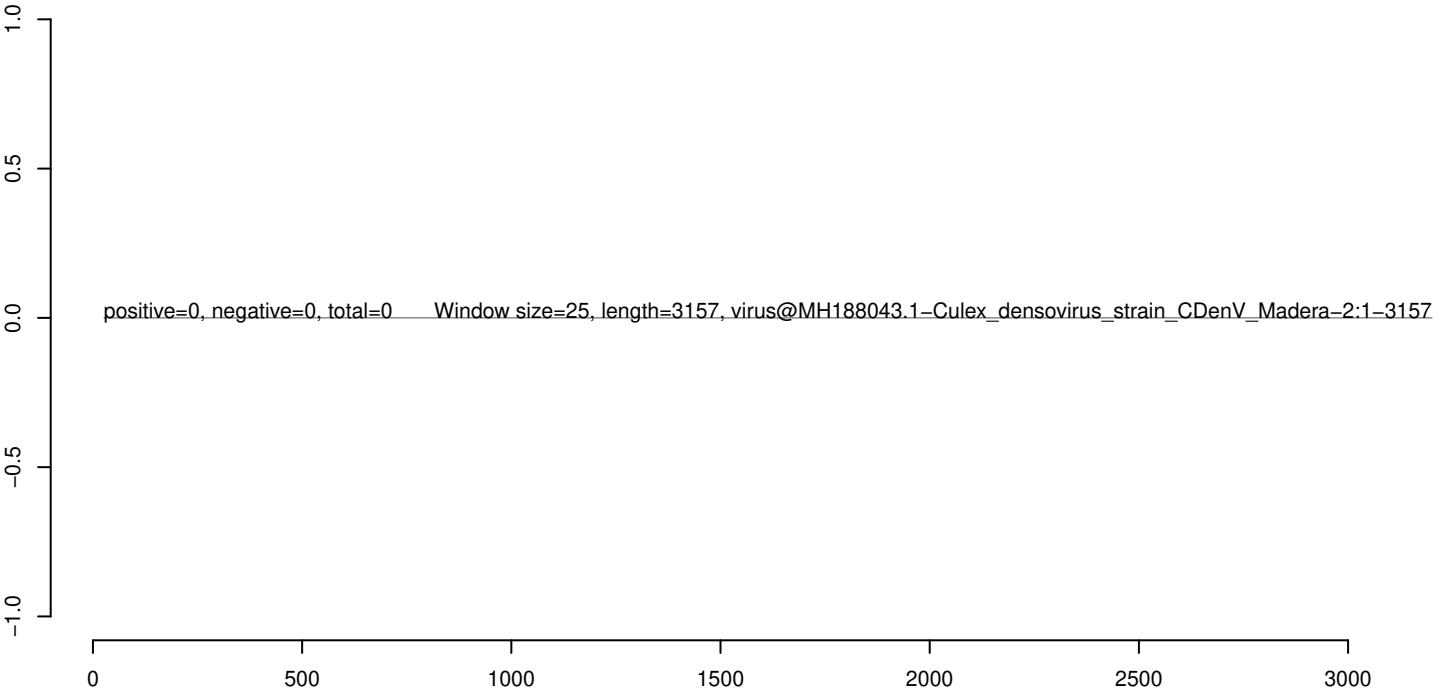
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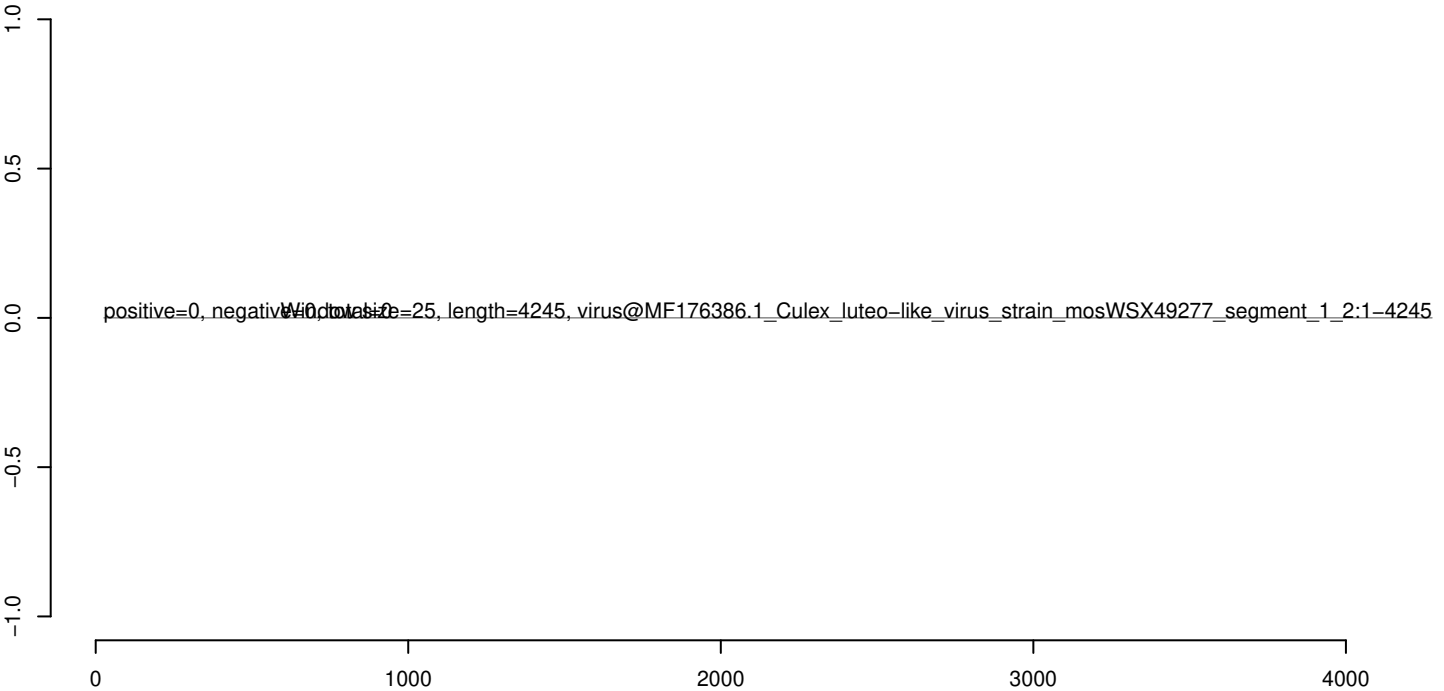
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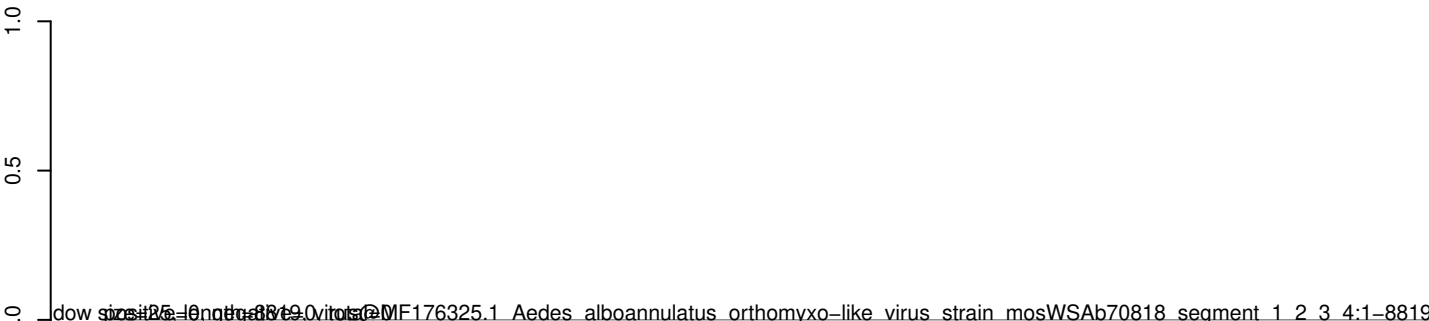
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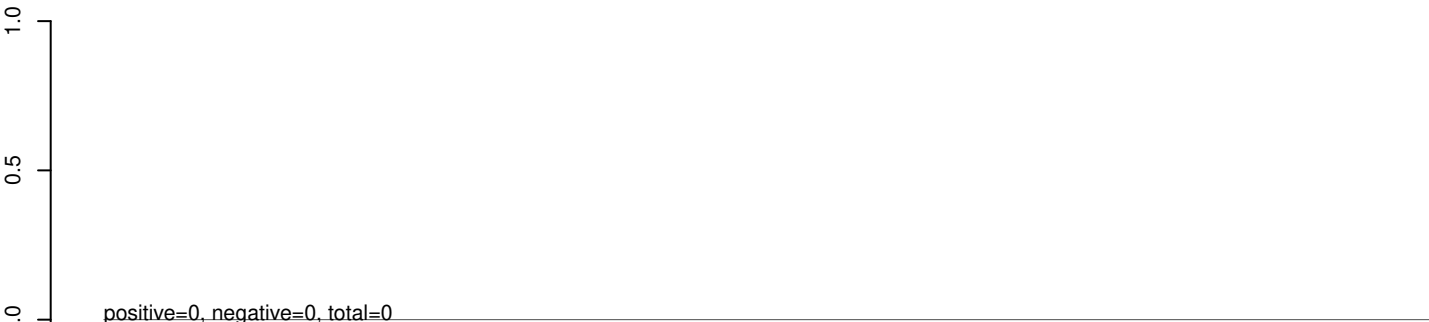


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AeAeg_Aag2_BTV1_AK.rep



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AeAeg_Aag2_BTV1_AK.18_23.rep



AeAeg_Aag2_BTV1_AK.24_35.rep



AeAeg_Aag2_BTV1_AK.rep



0 2000 4000 6000 8000 10000 12000

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AeAeg_Aag2_BTV1_AK.rep



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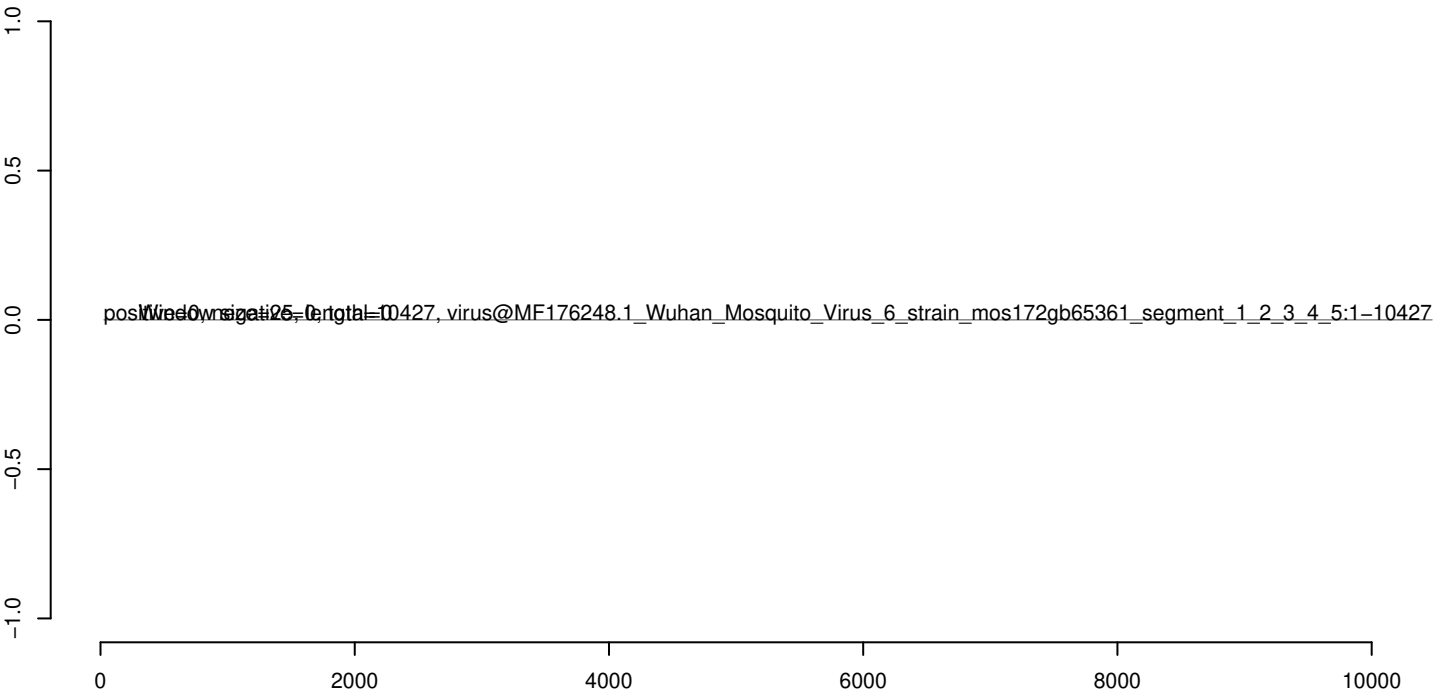
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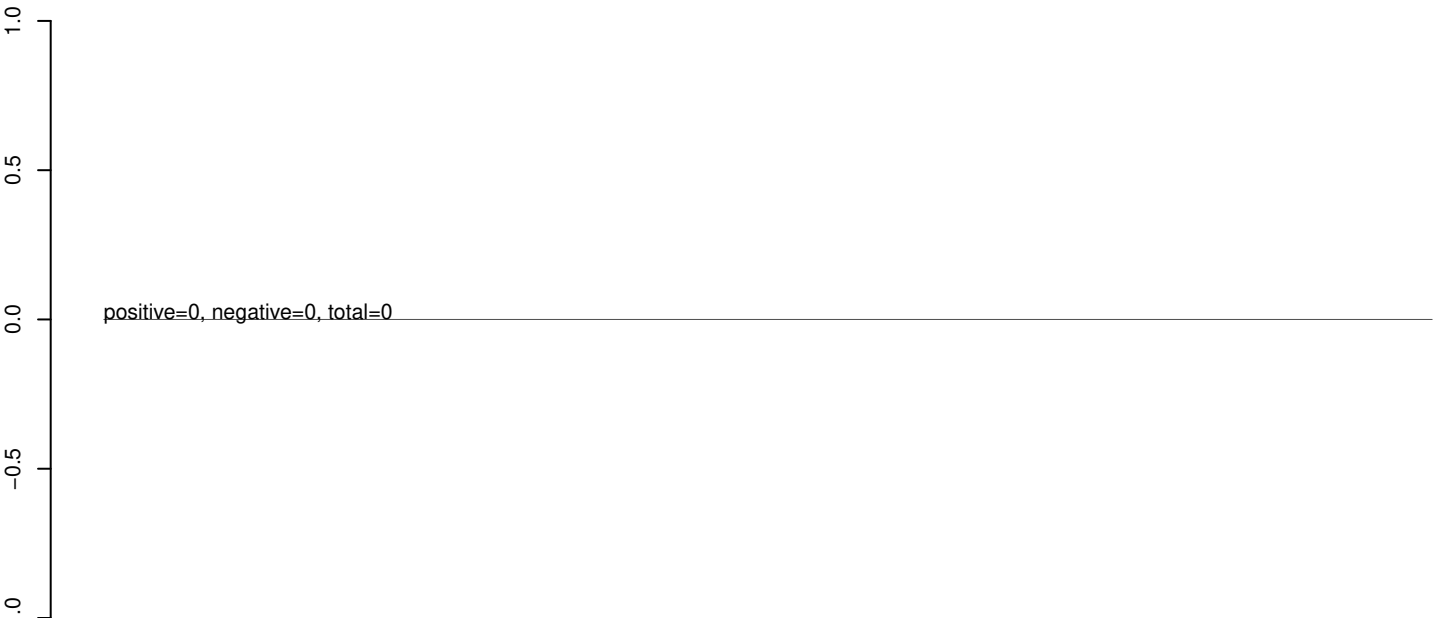
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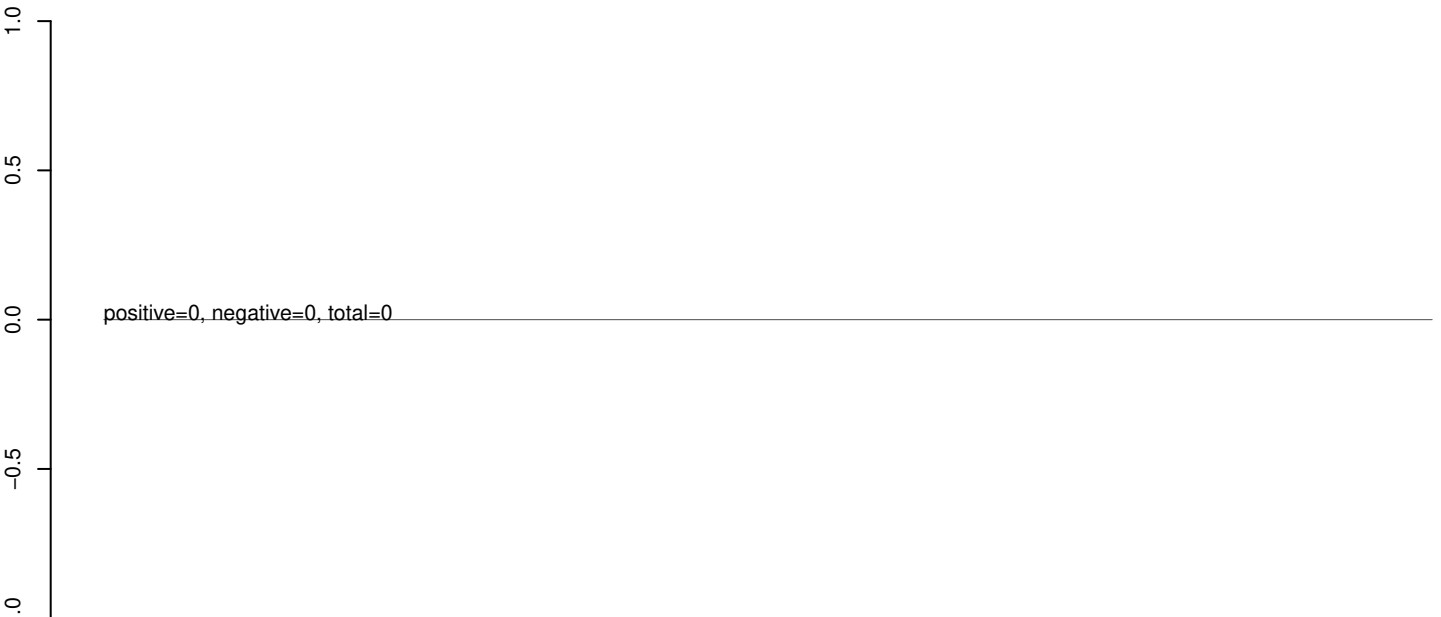
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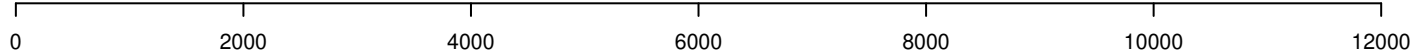
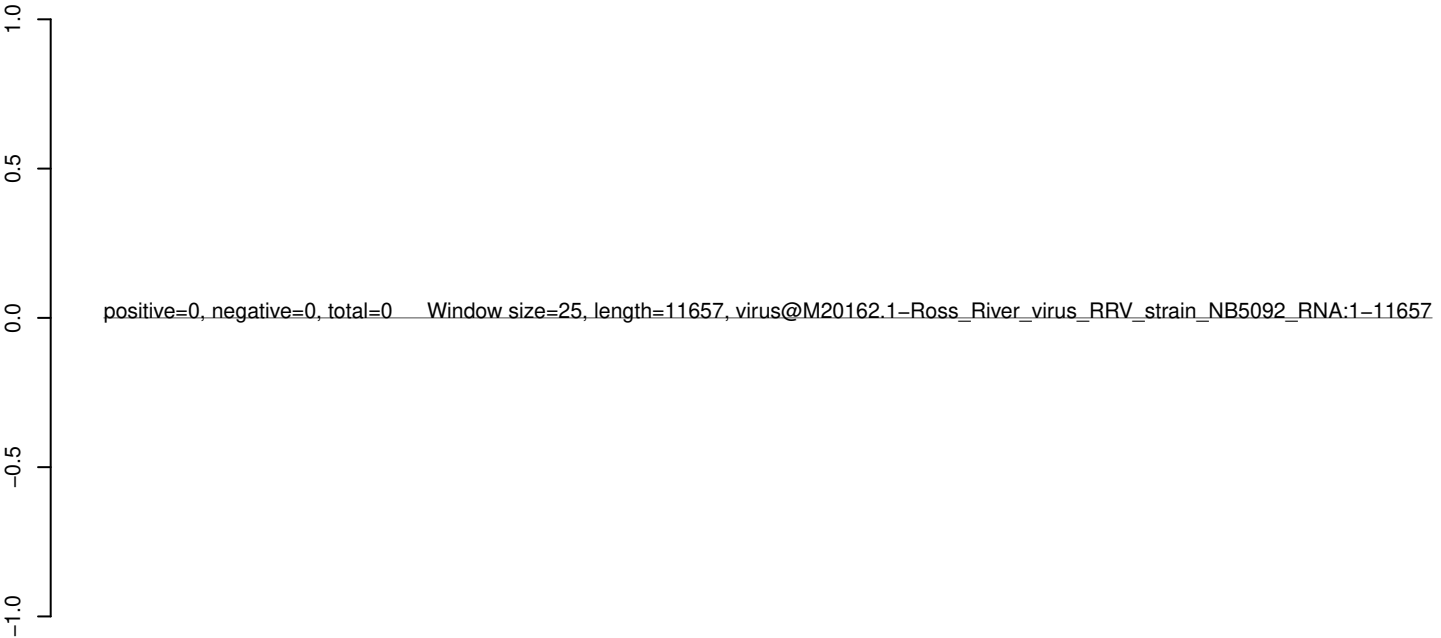
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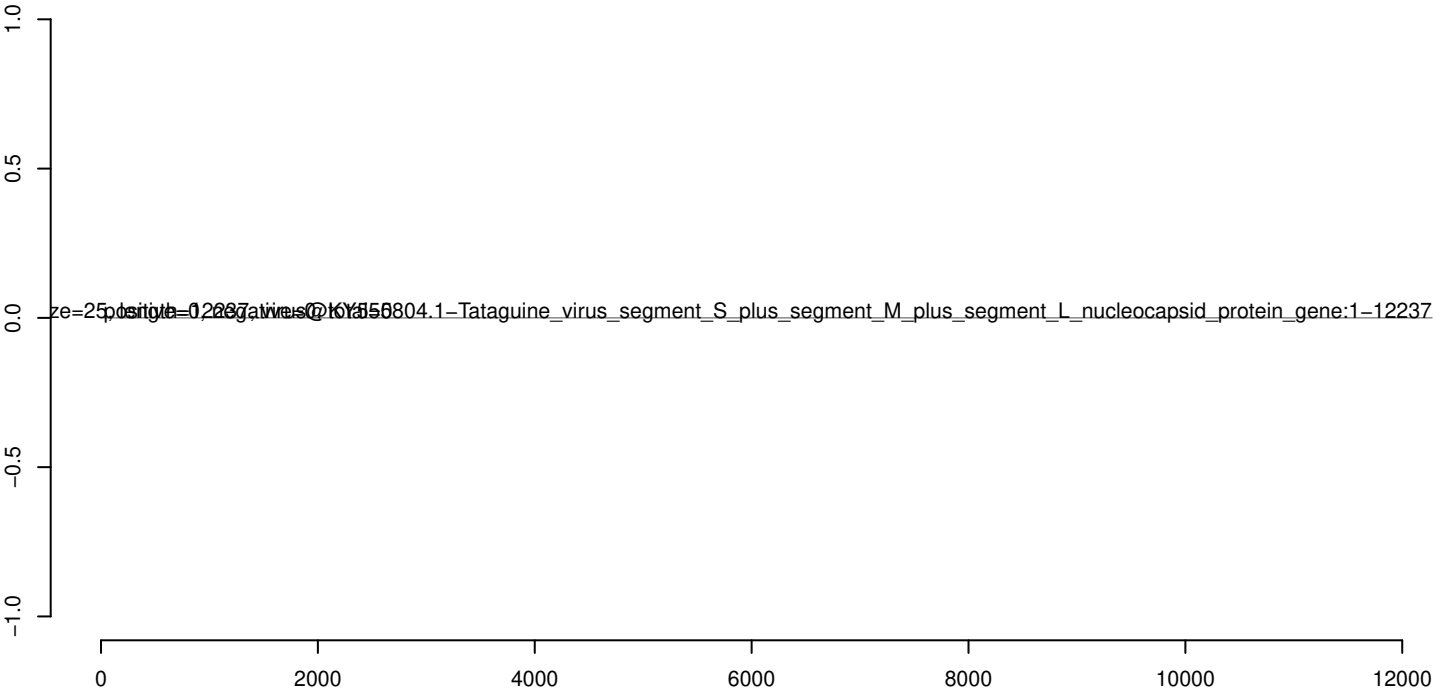
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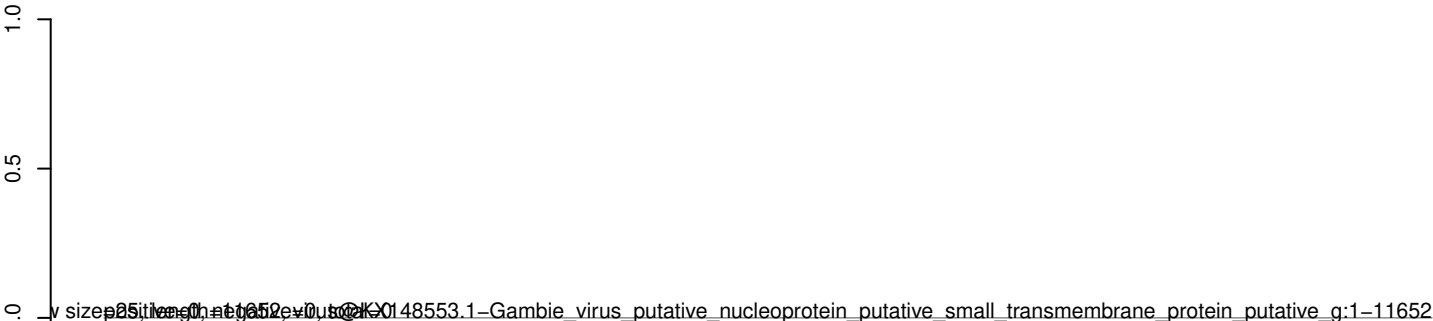
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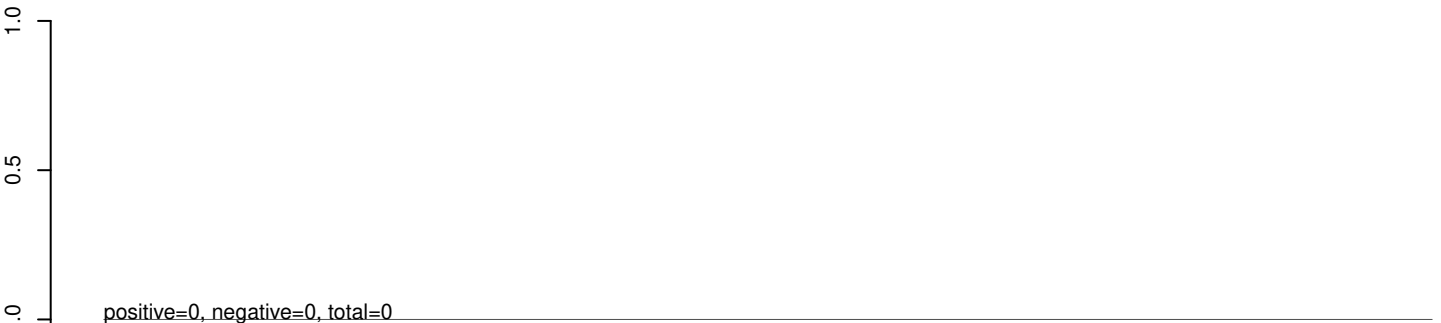
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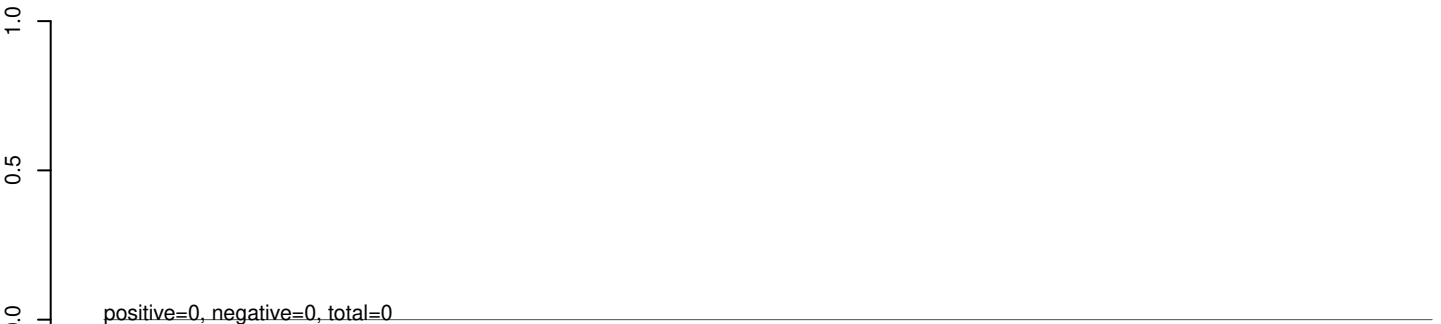
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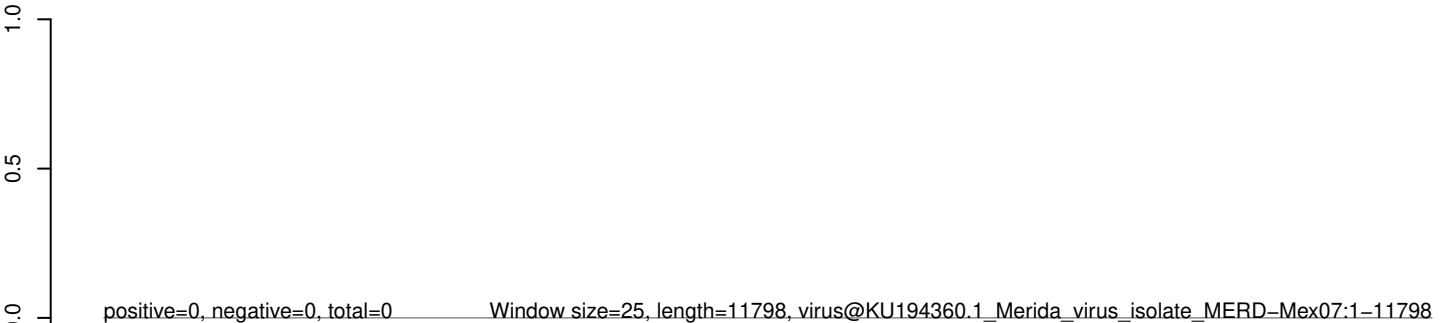
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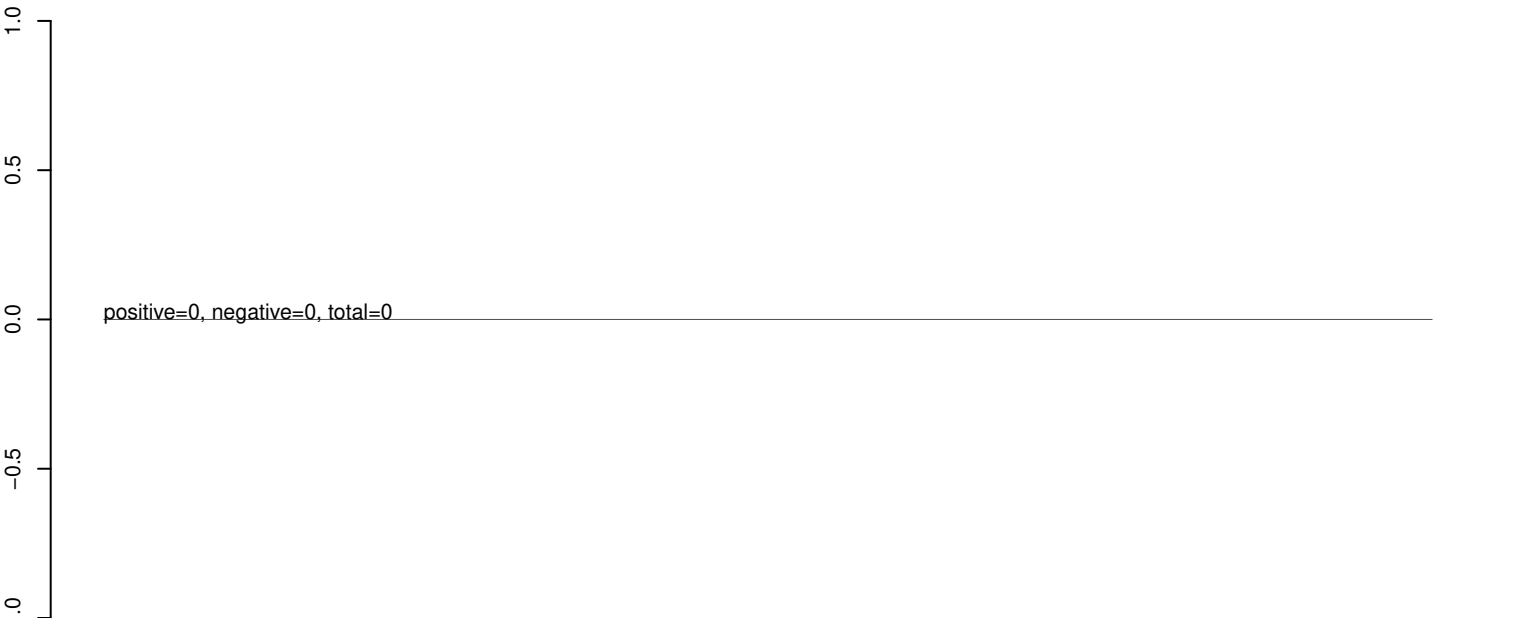


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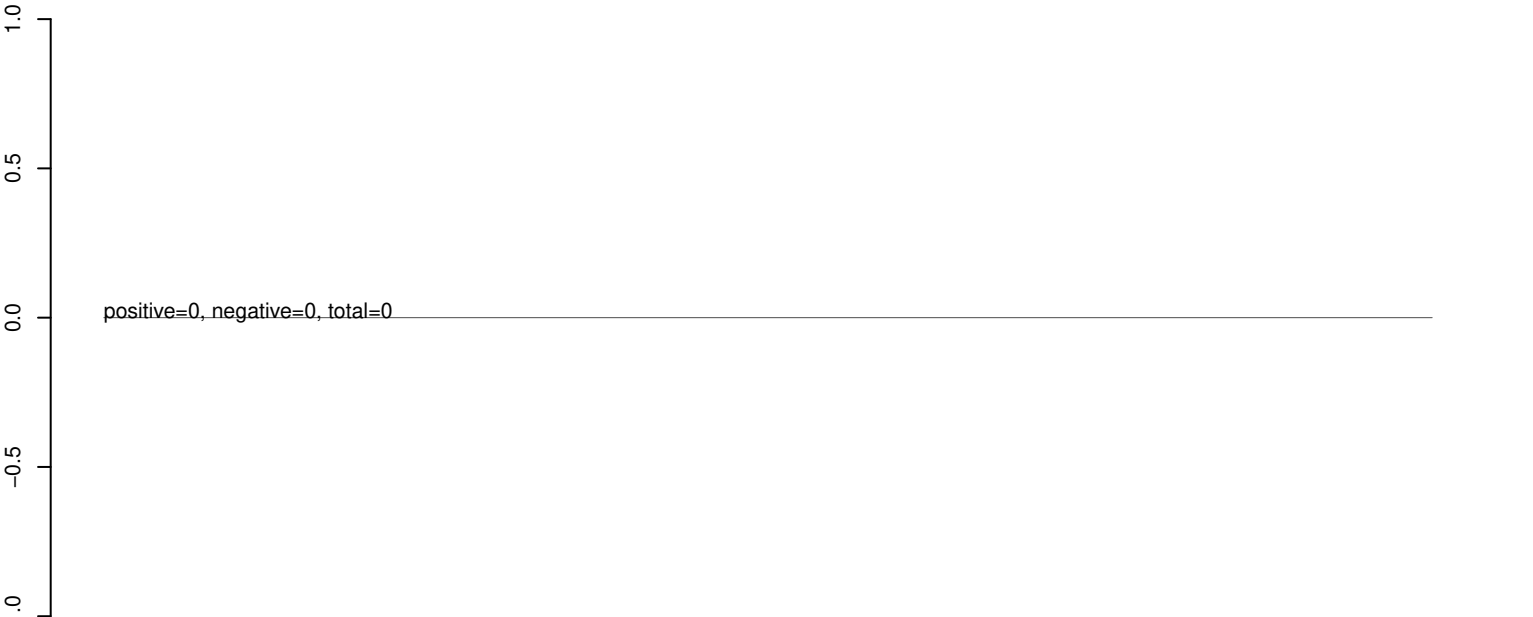


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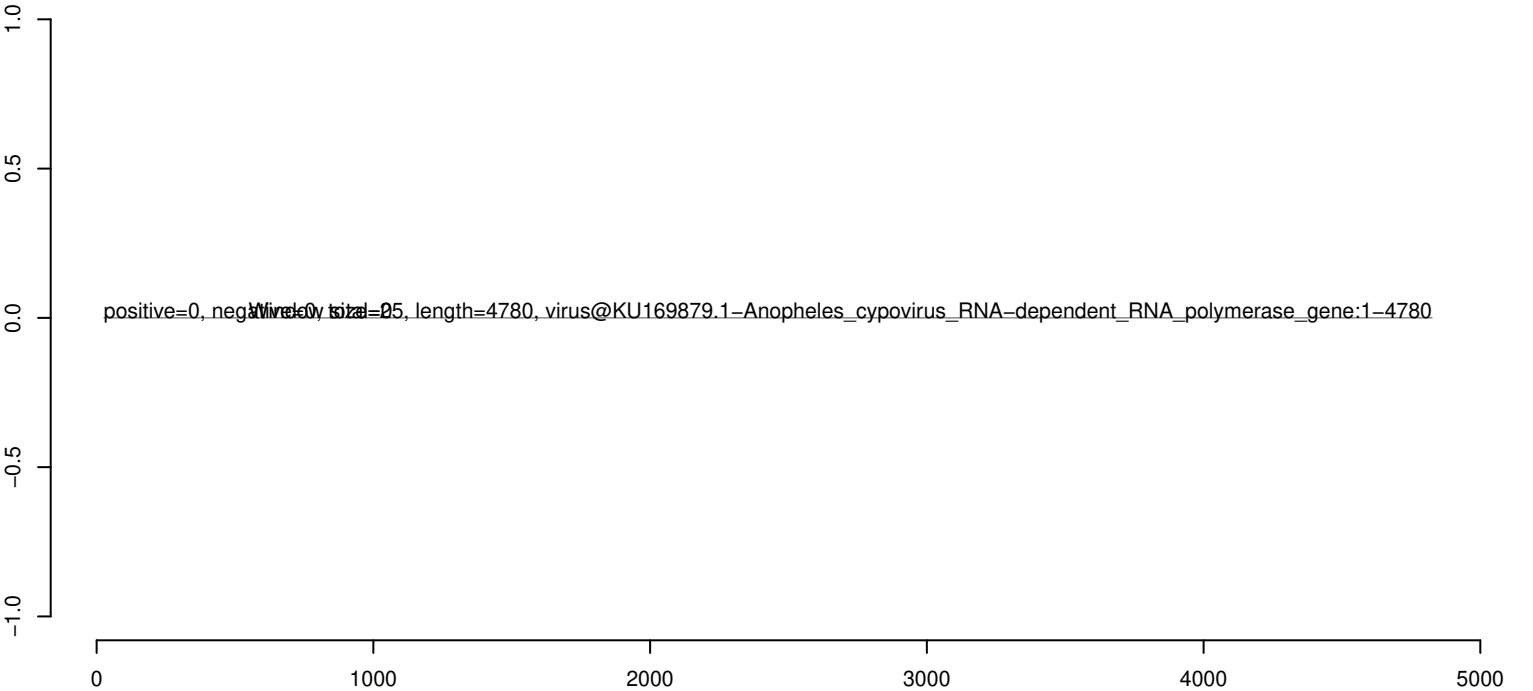
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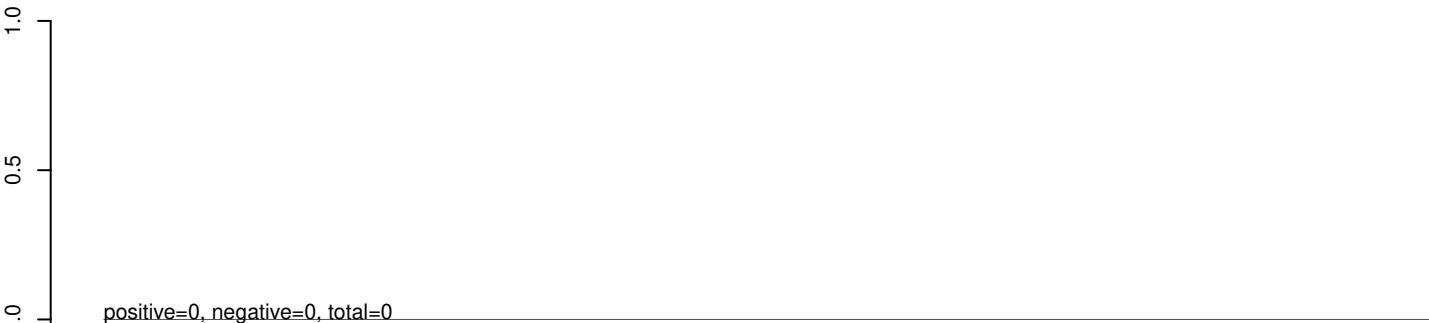
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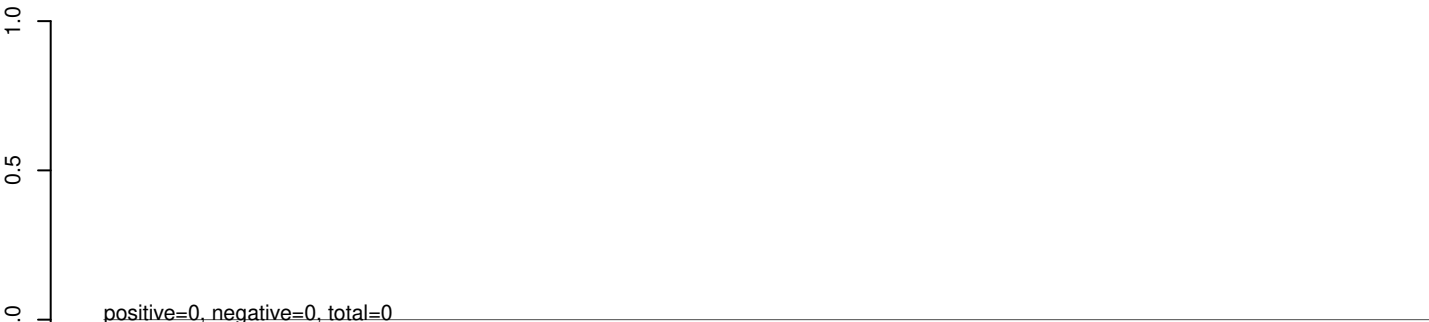
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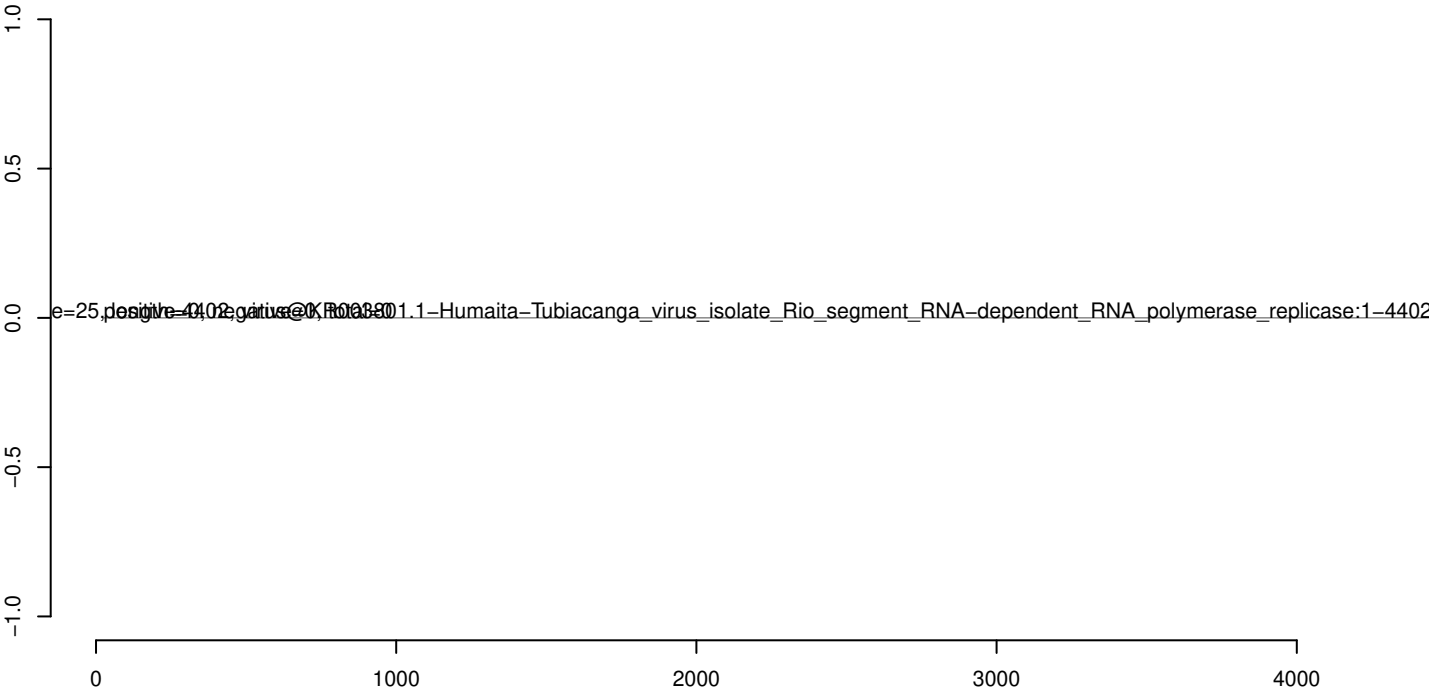
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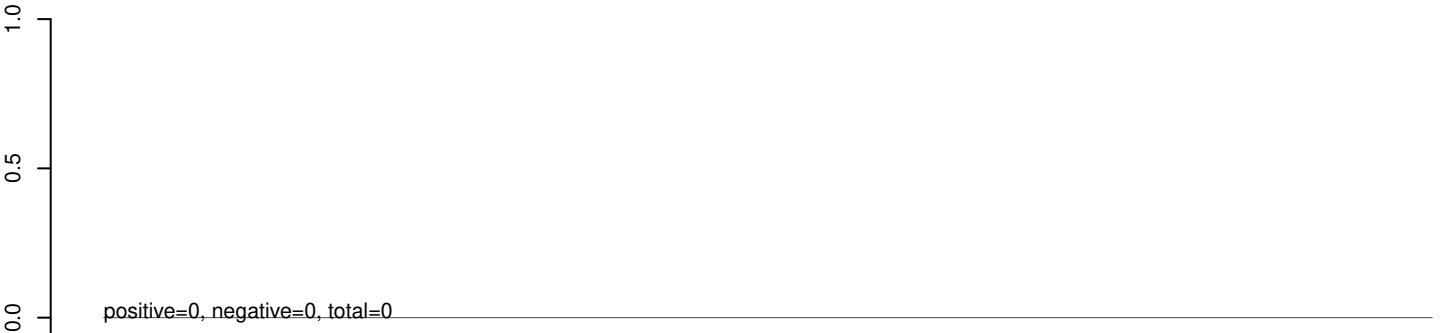
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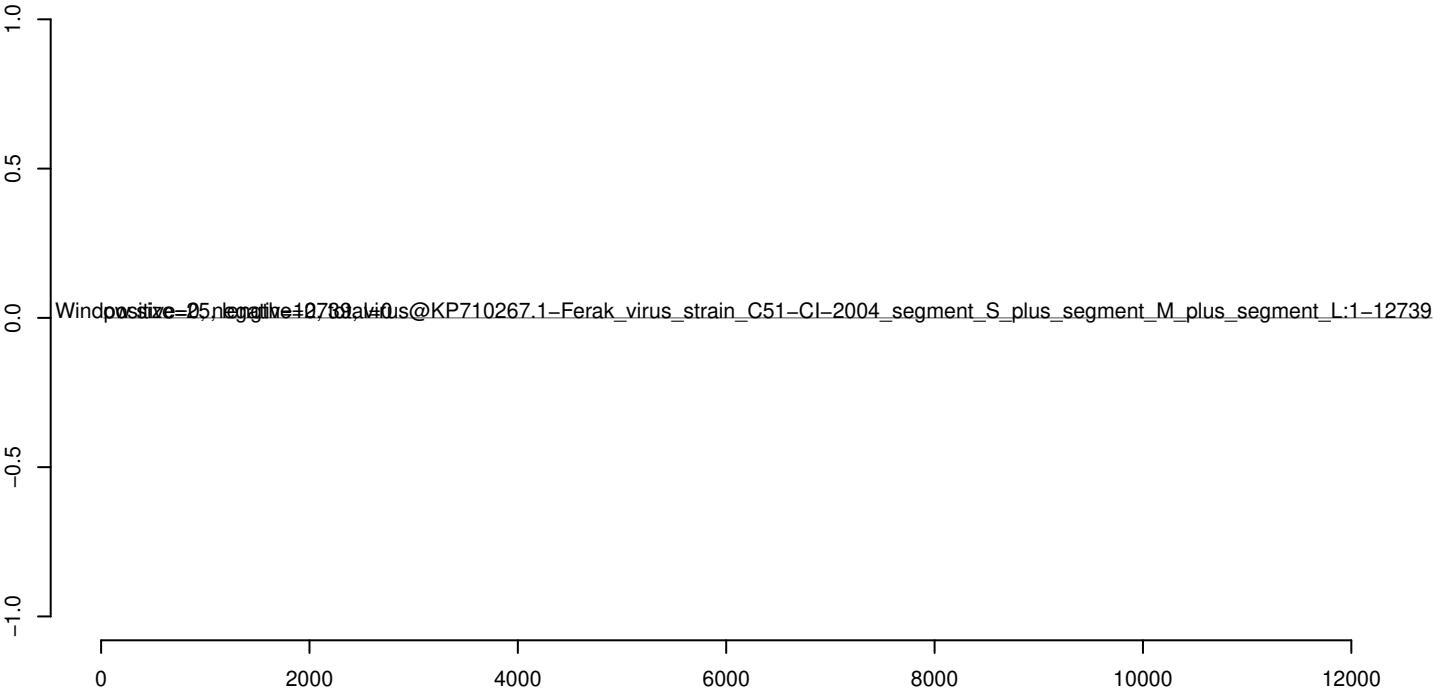
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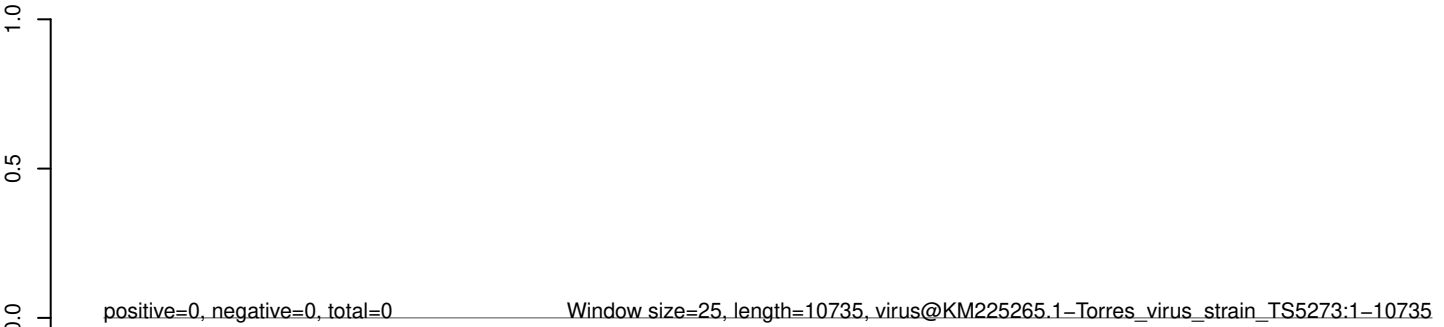
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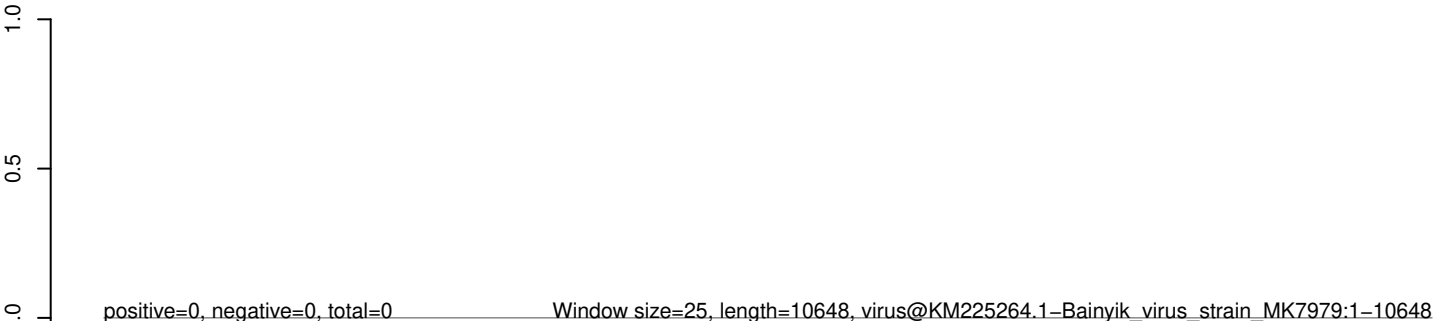
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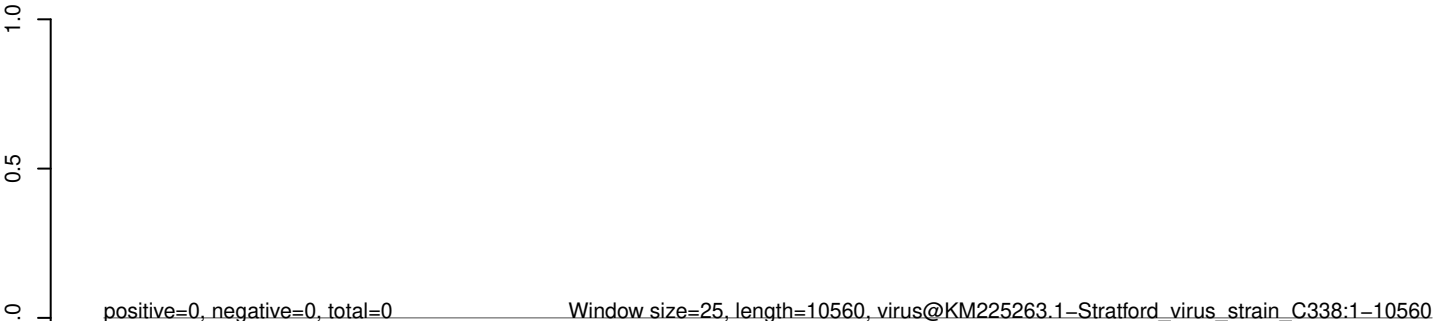
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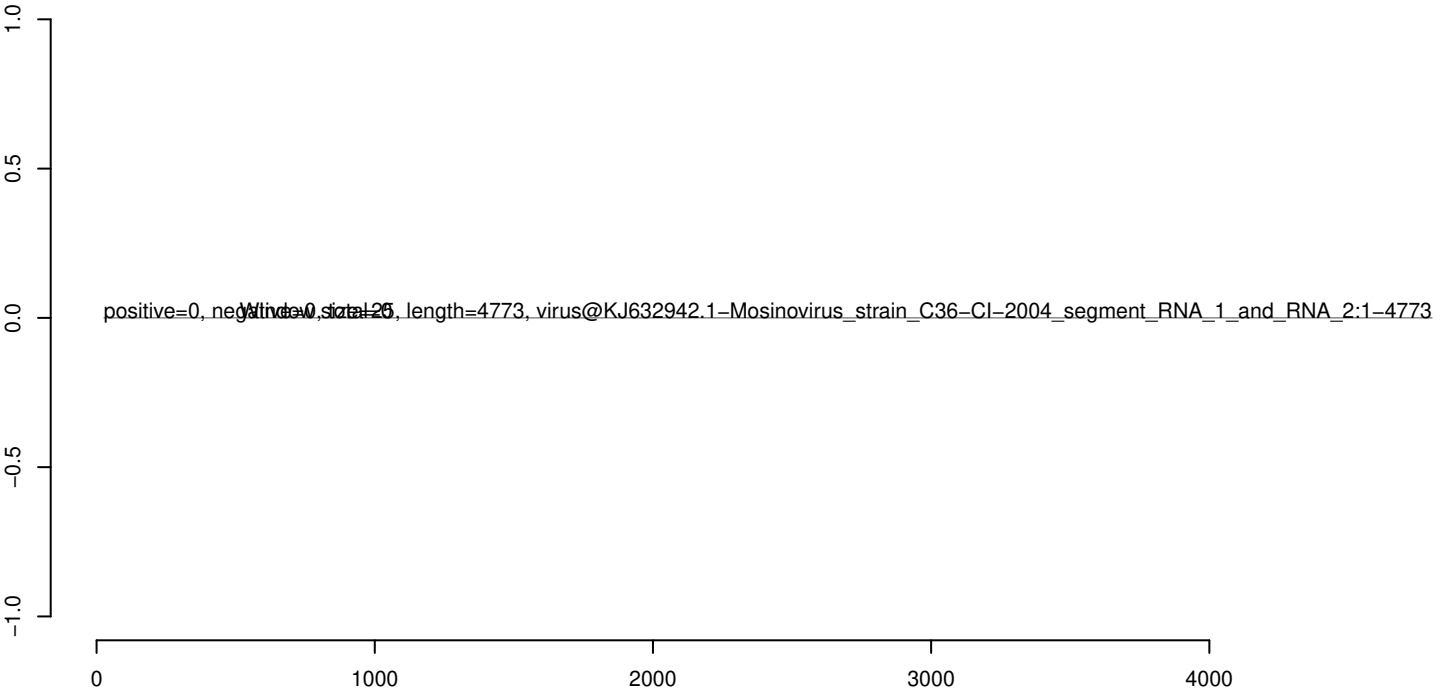
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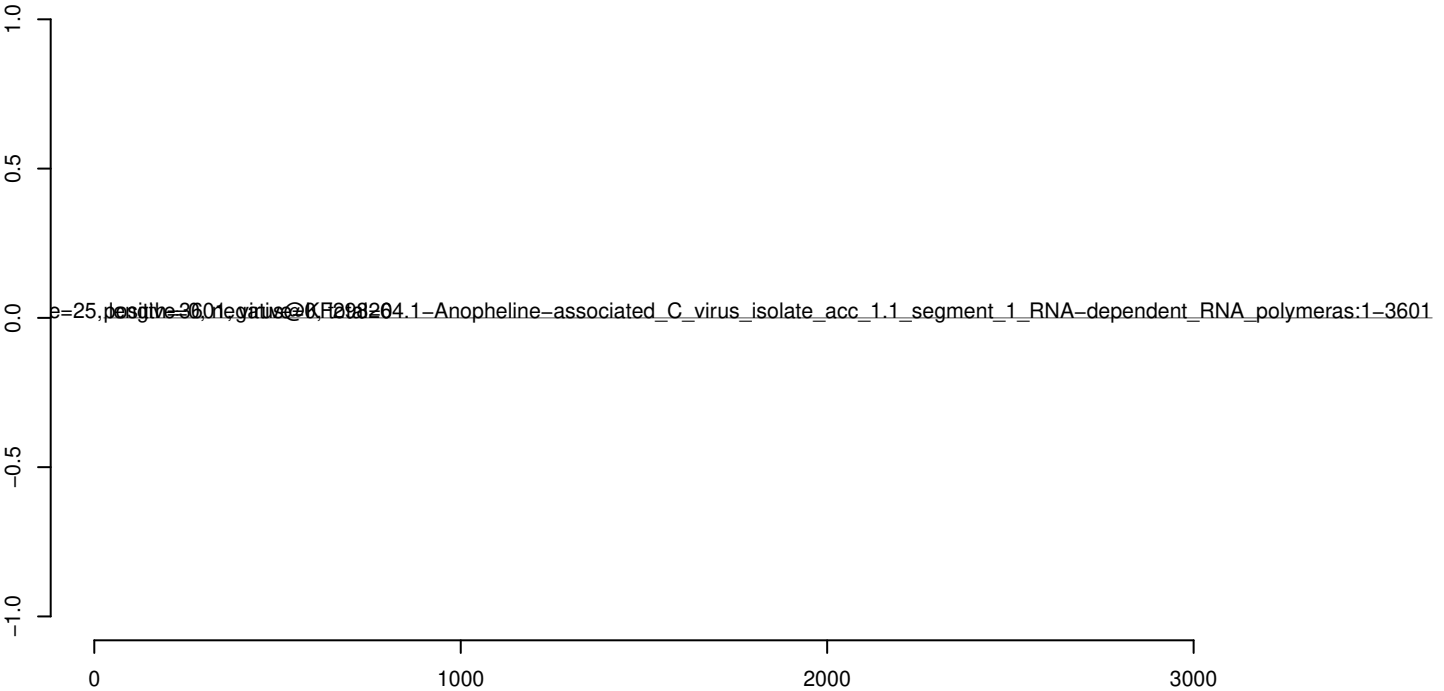
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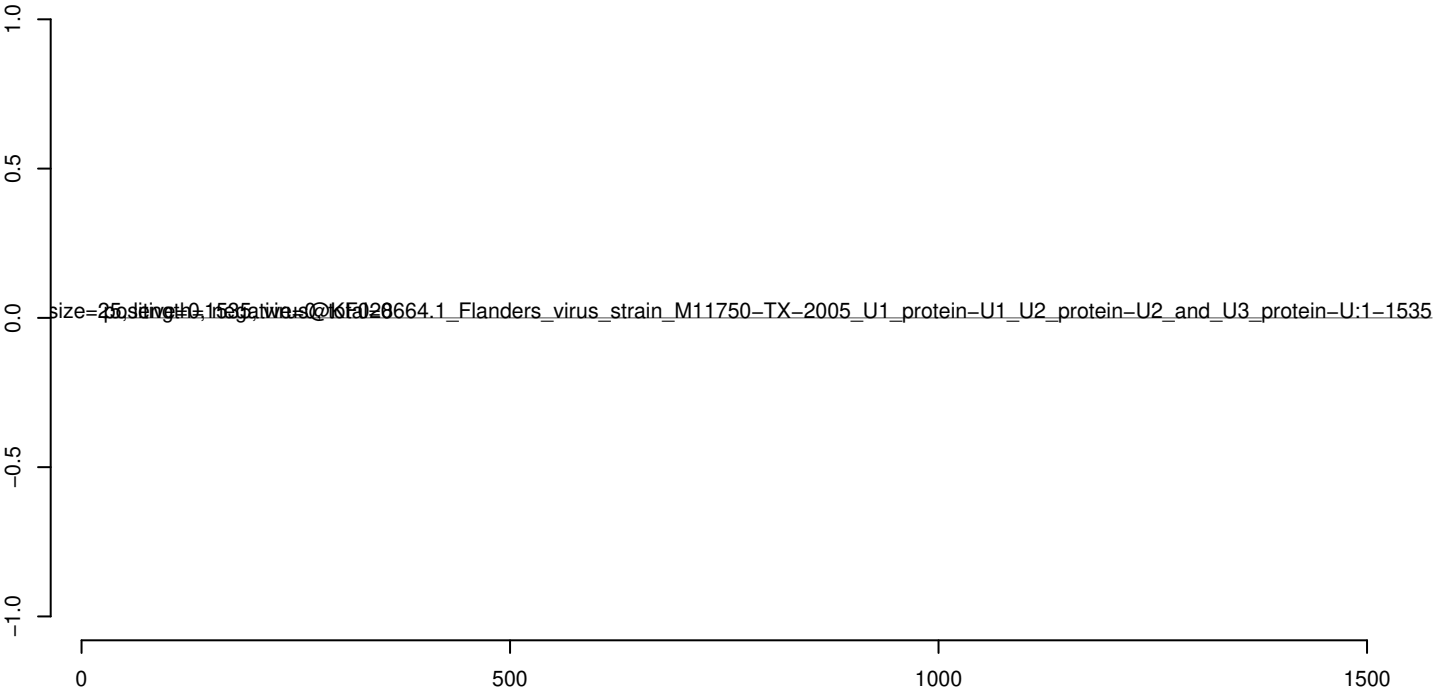
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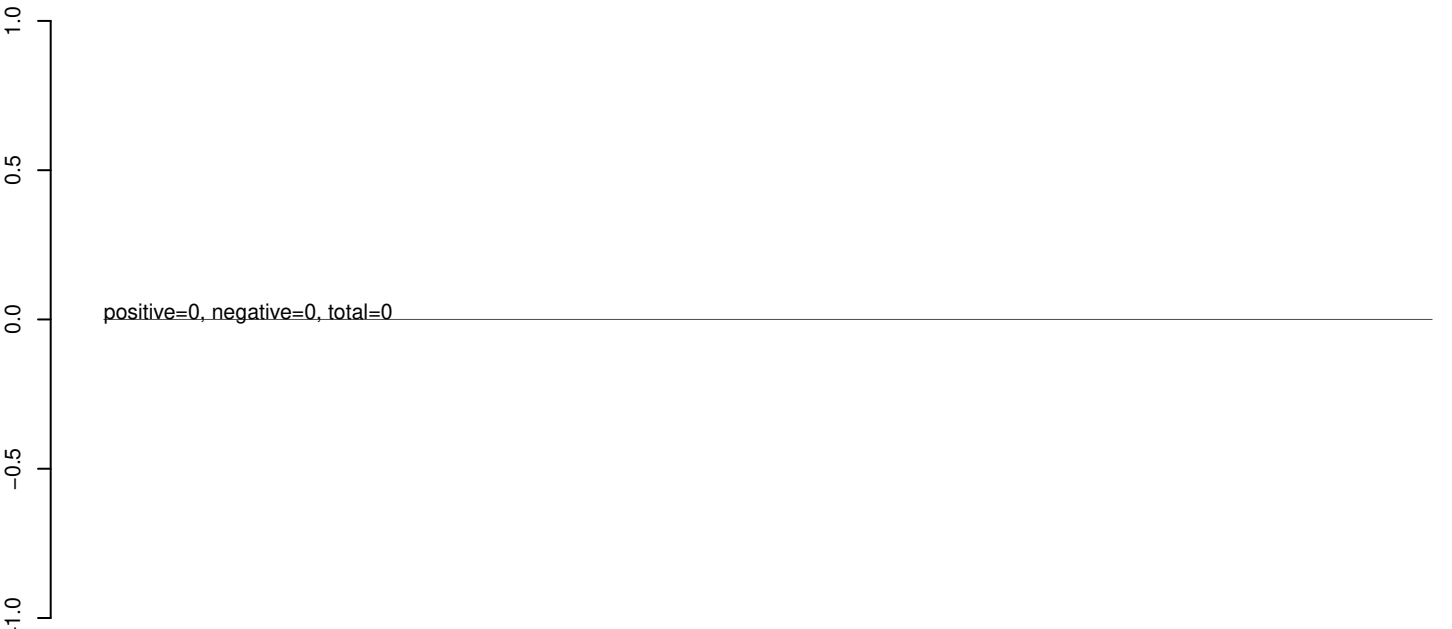
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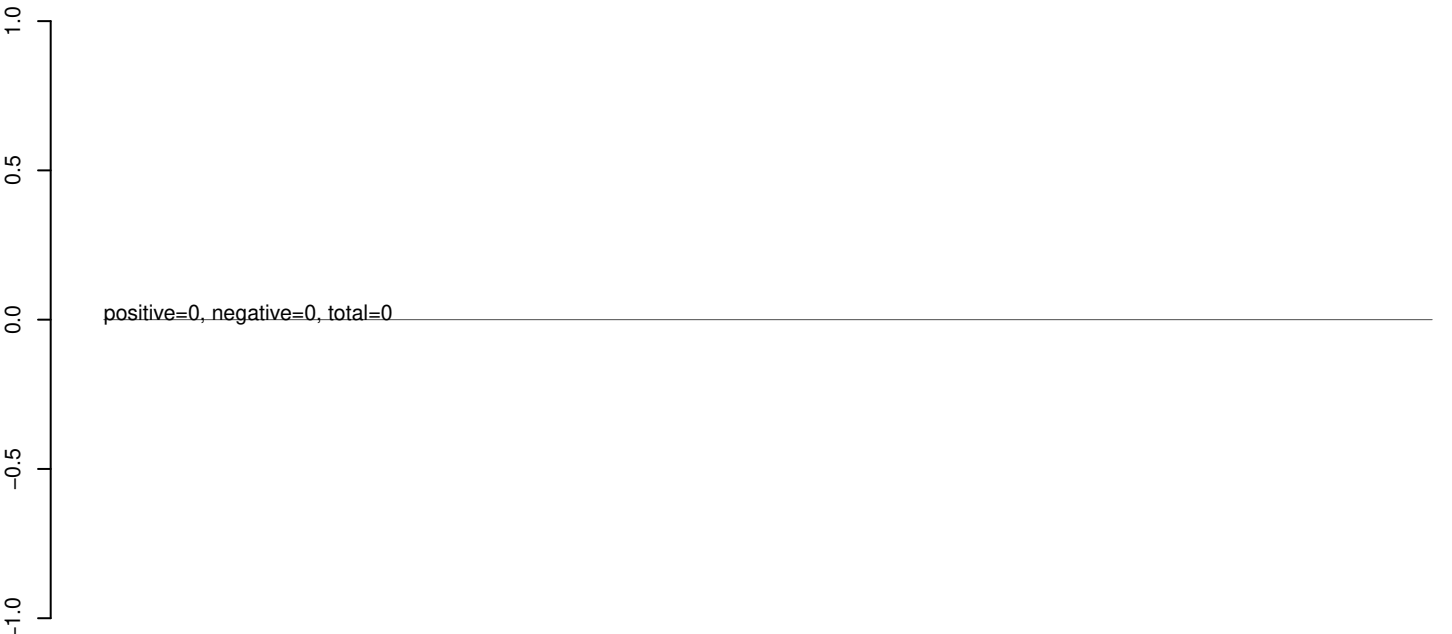
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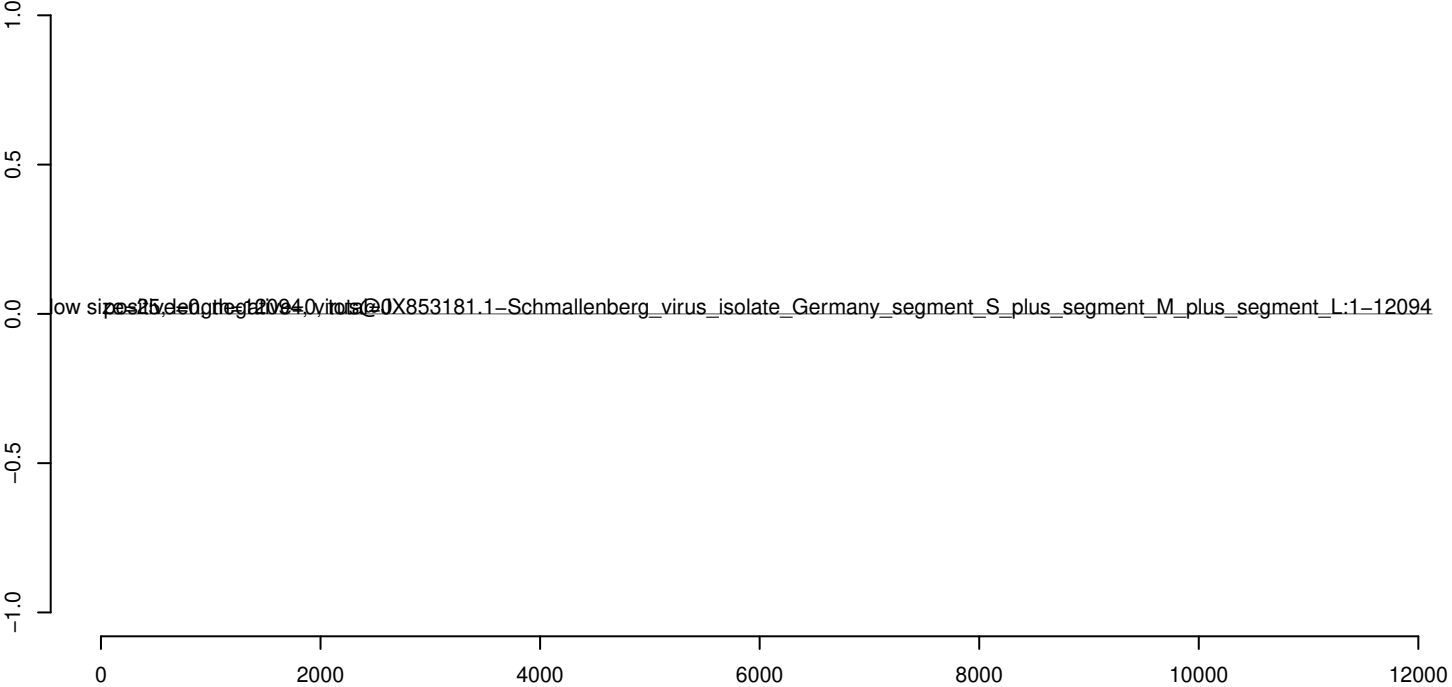
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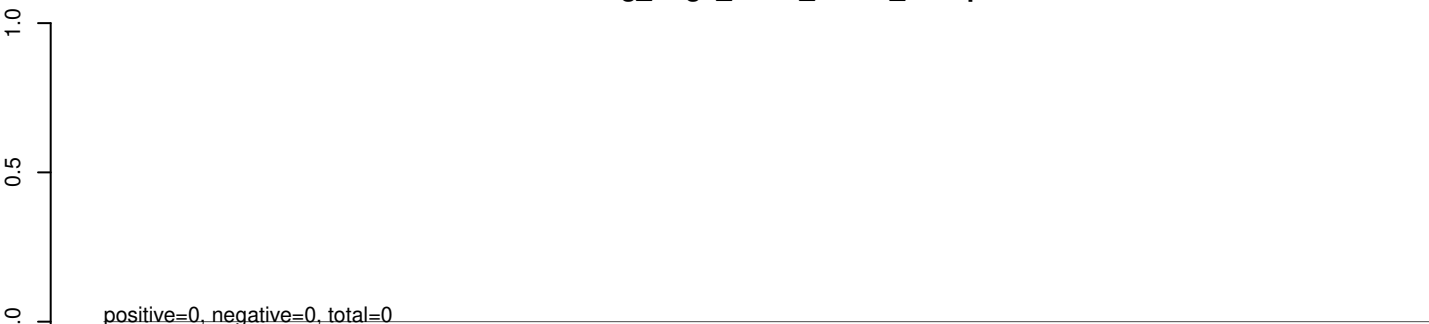
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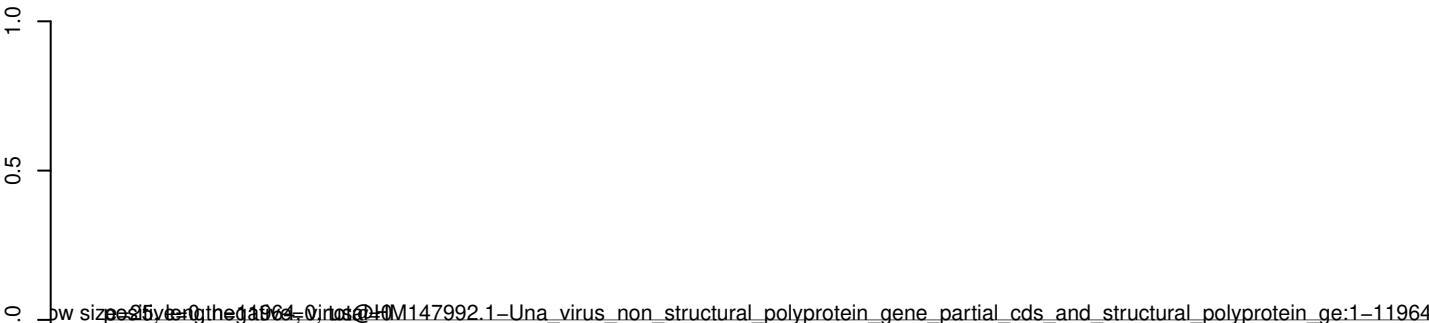
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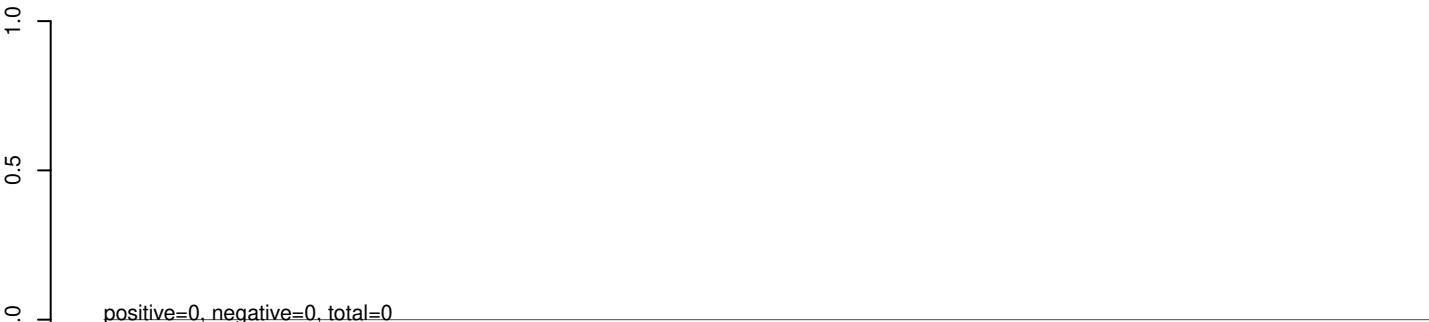
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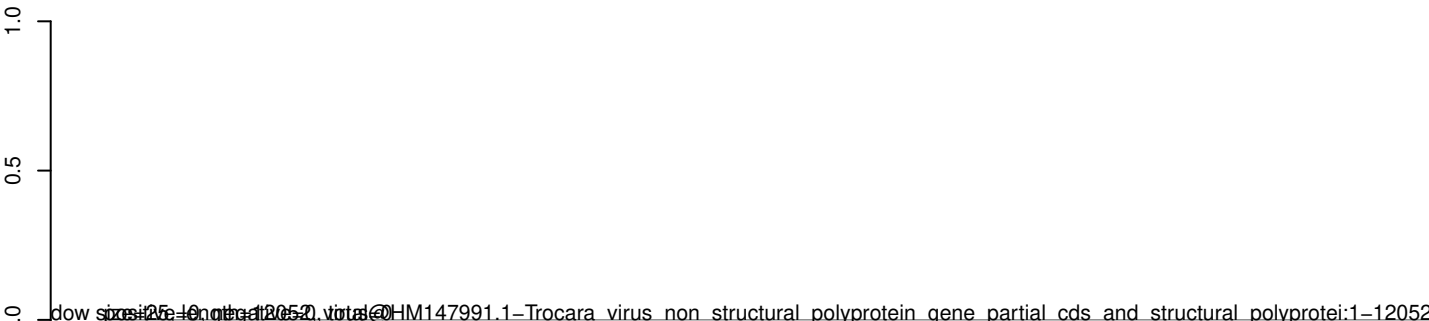
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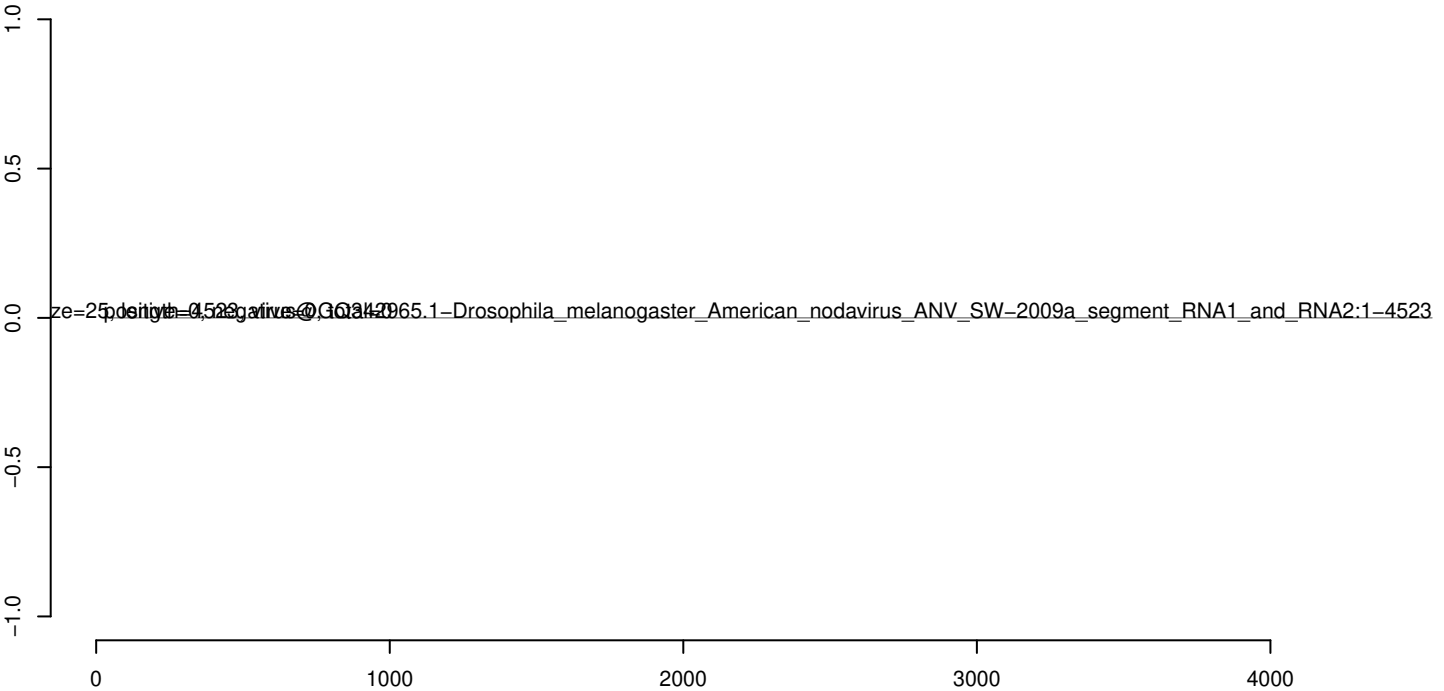
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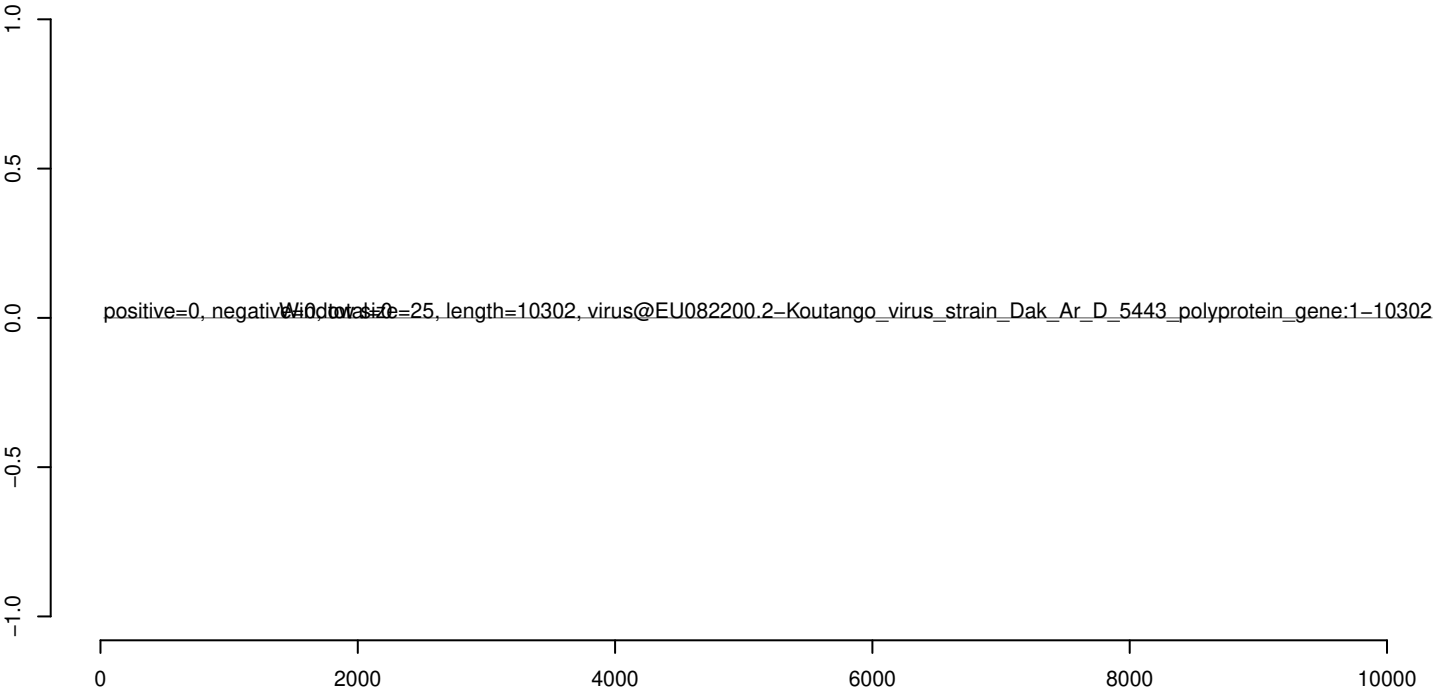
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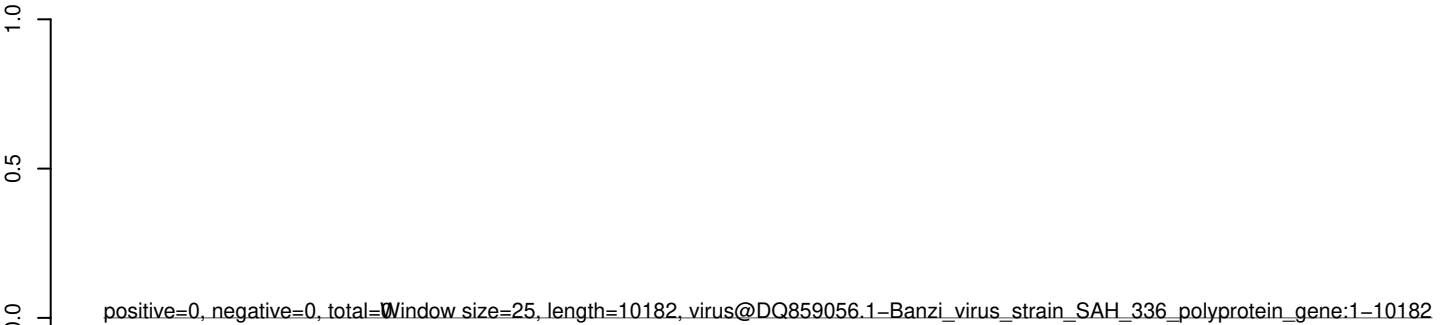
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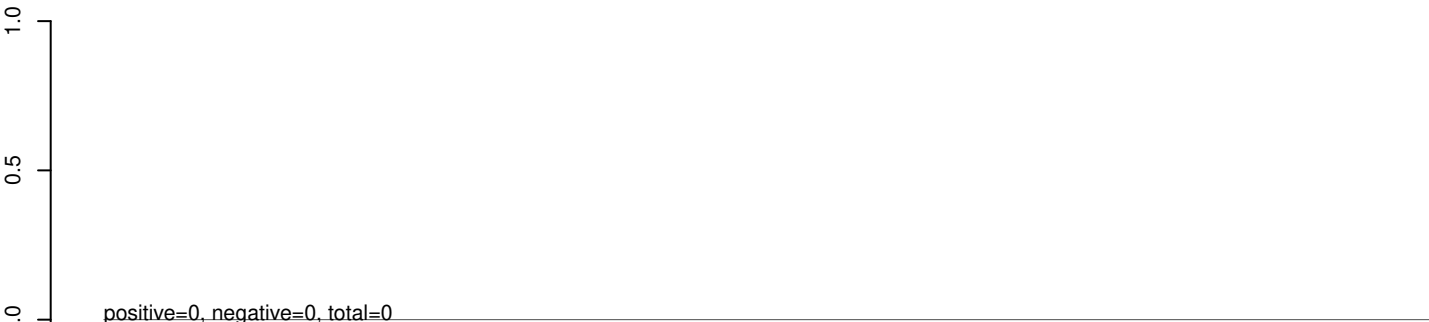
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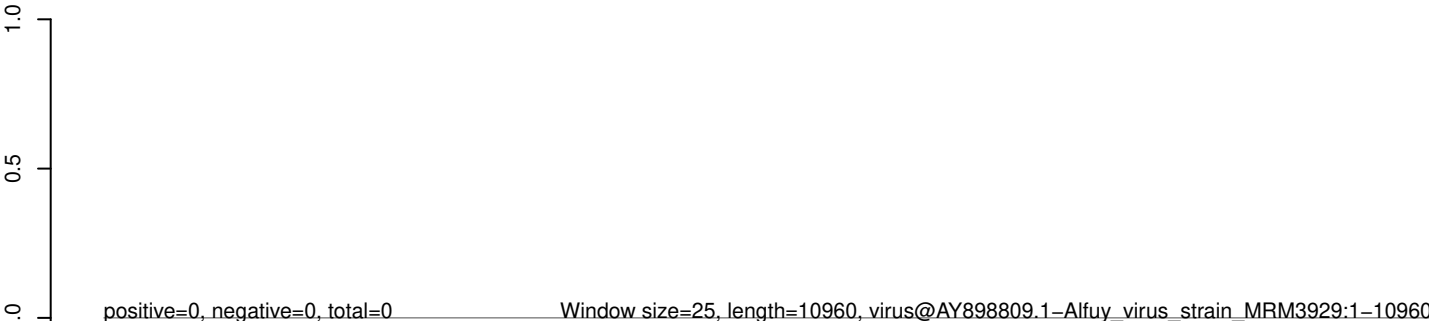
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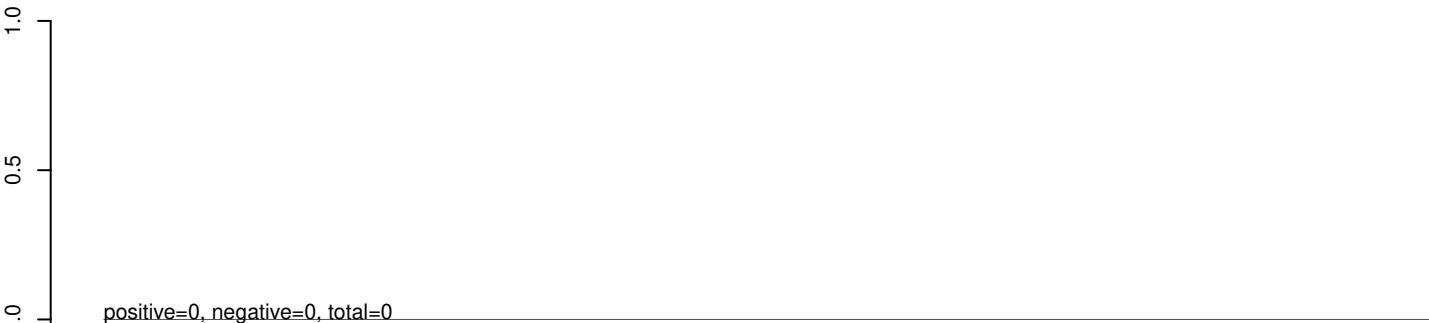


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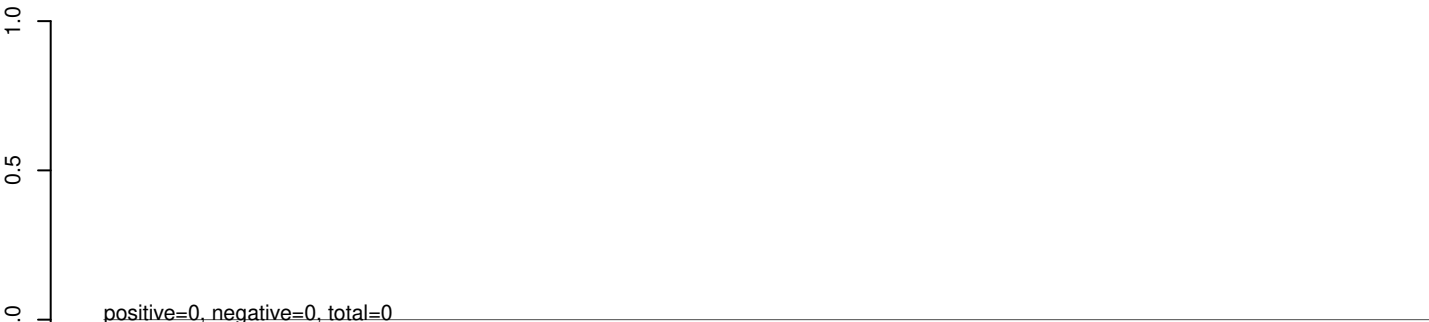


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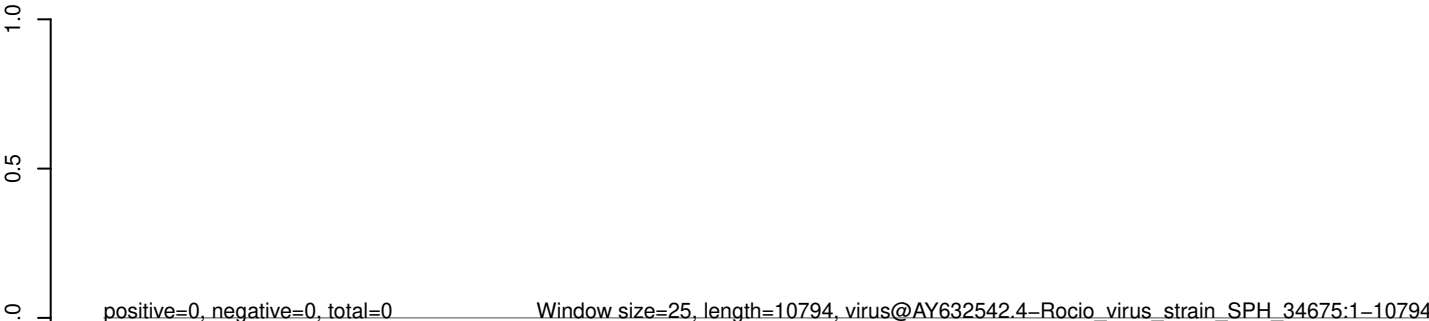
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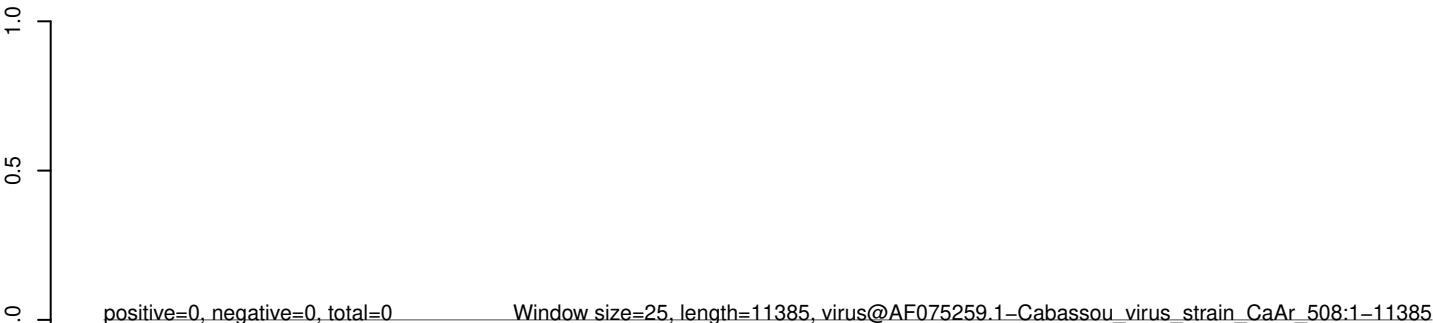
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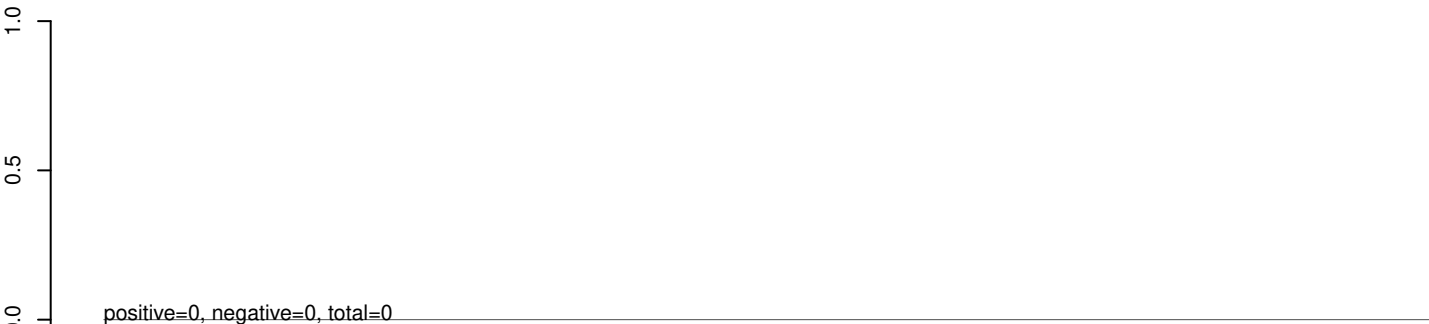


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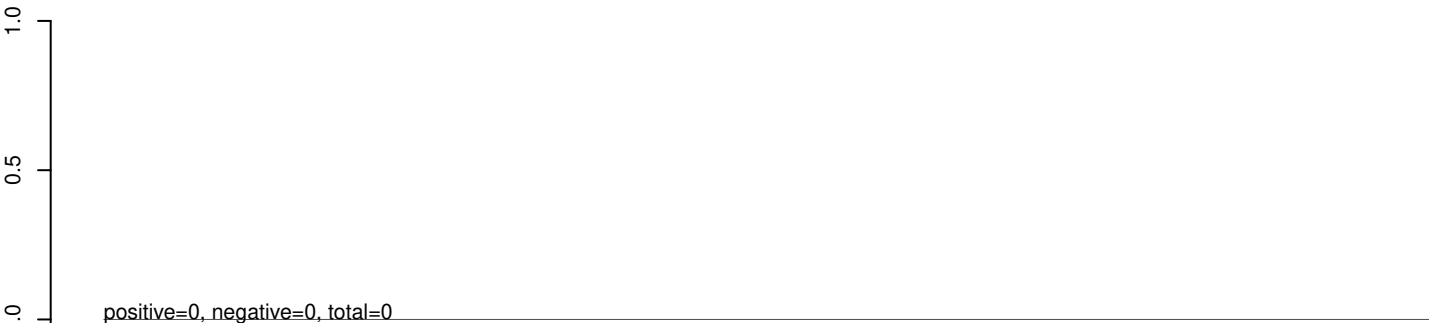


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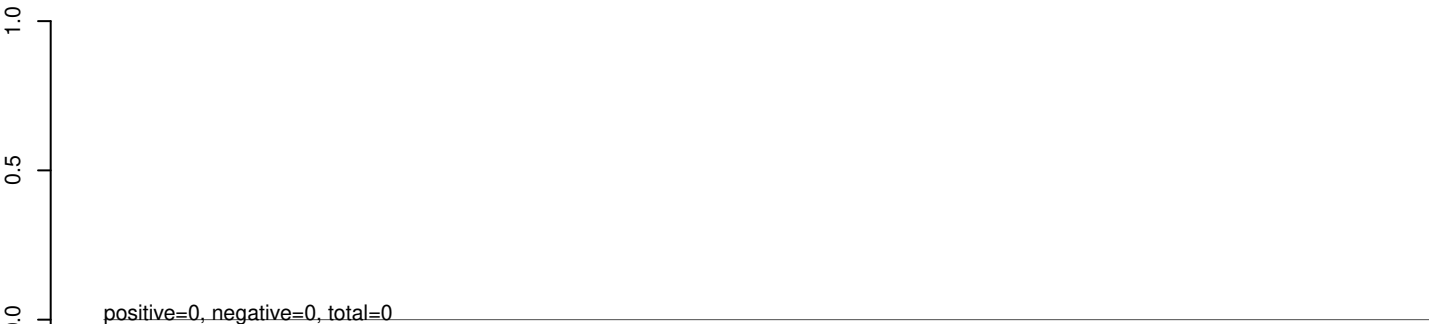


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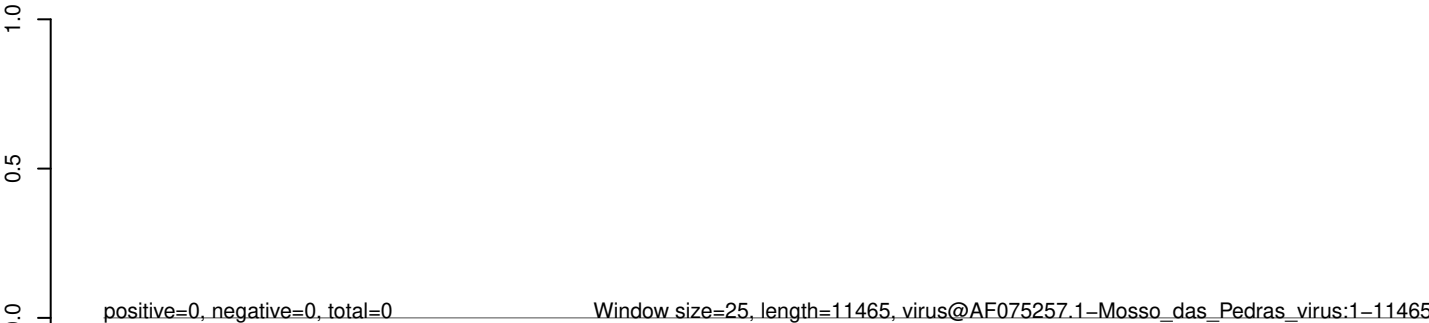
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