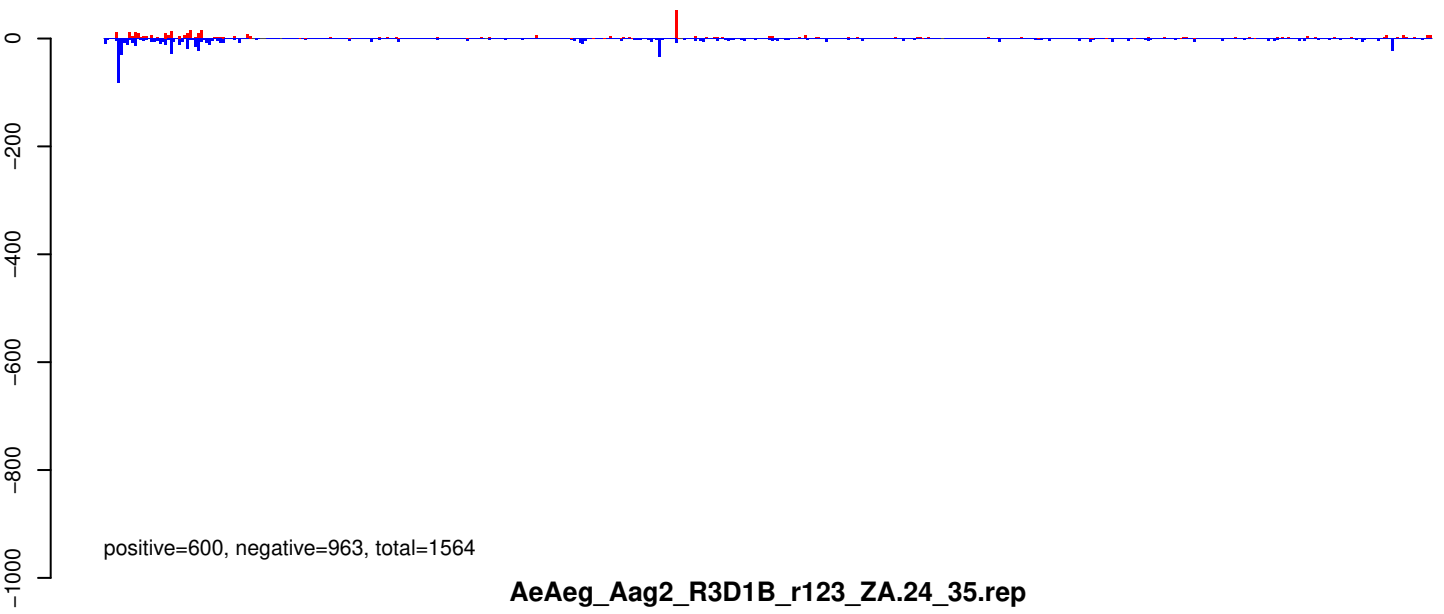
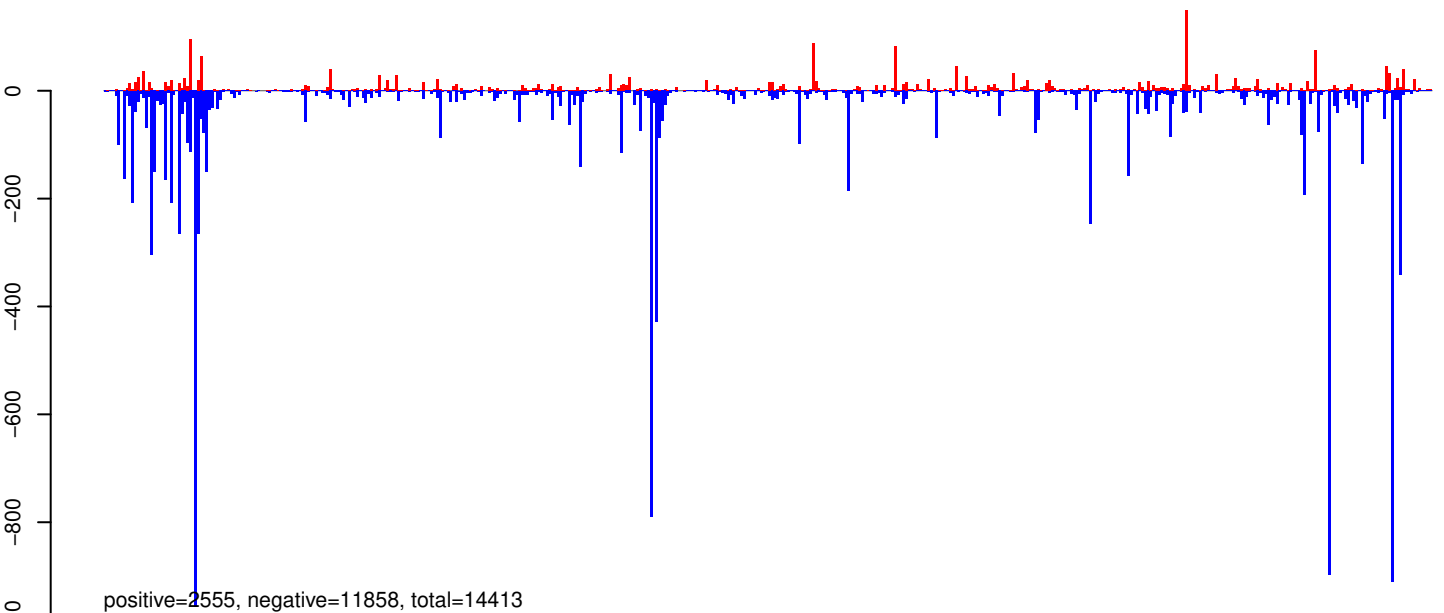


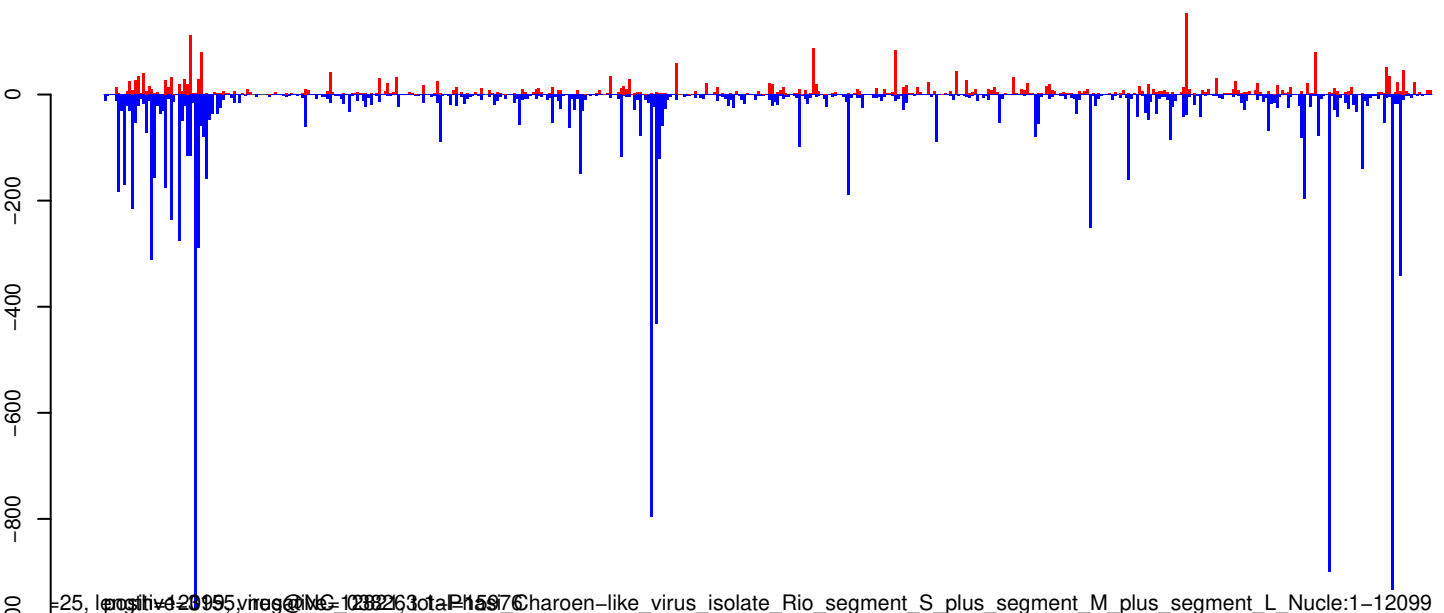
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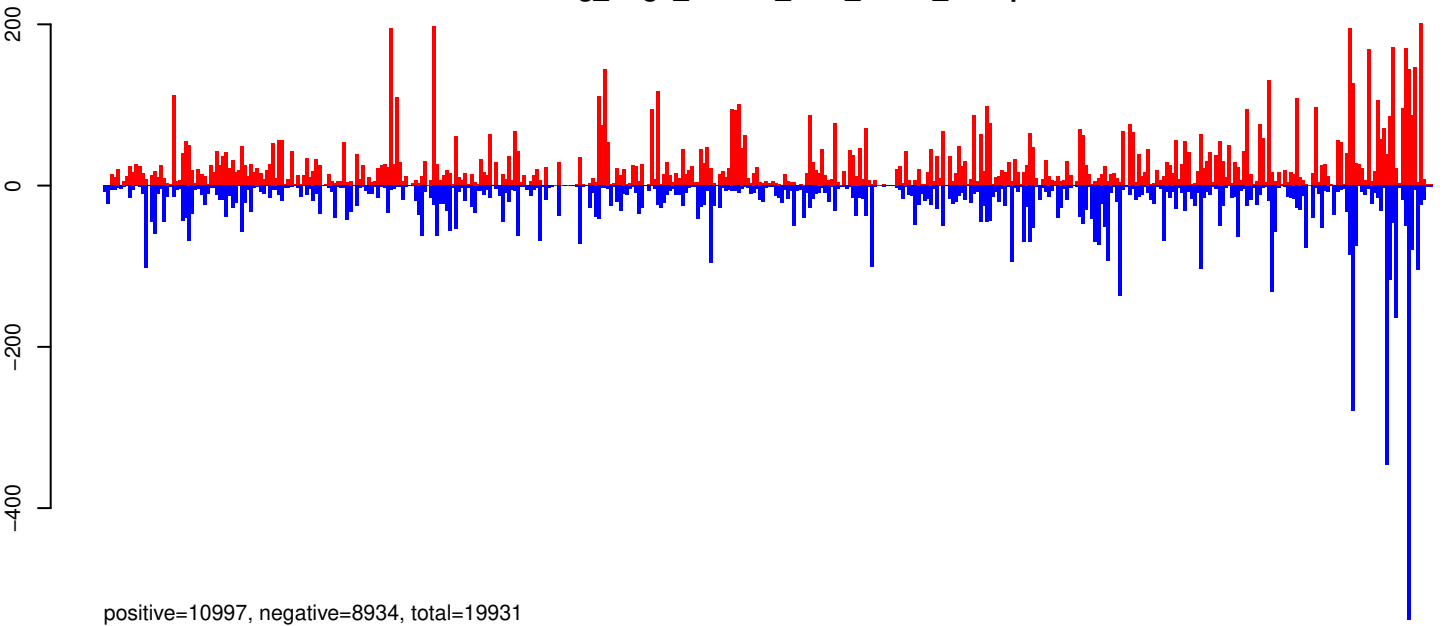
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AeAeg_Aag2_R3D1B_r123_ZA.rep

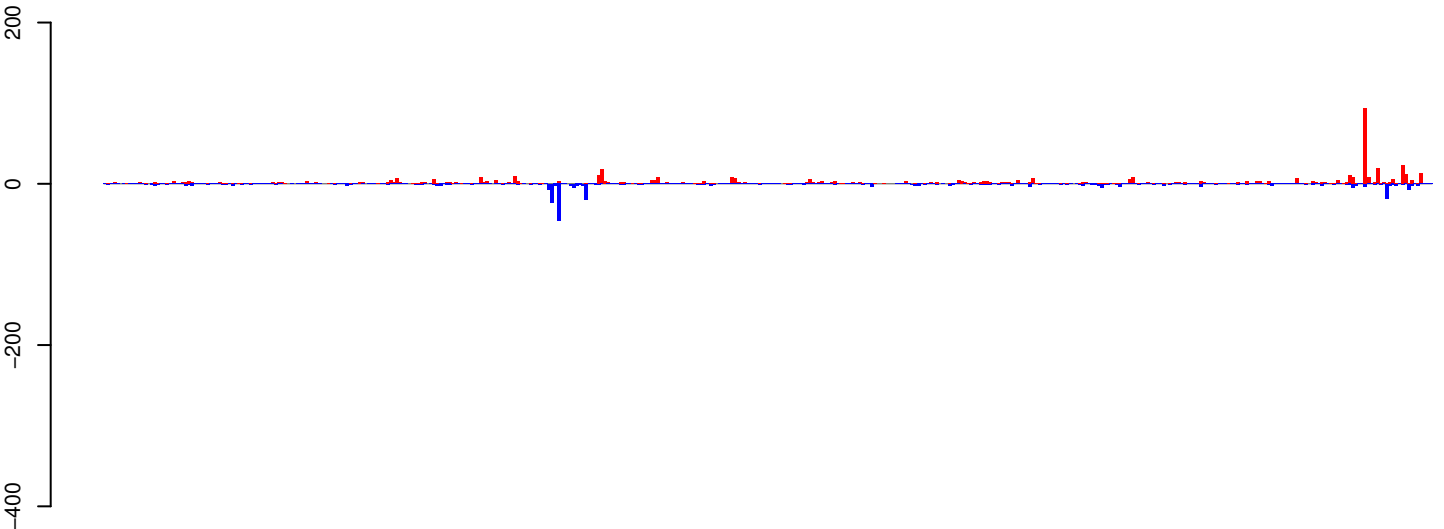


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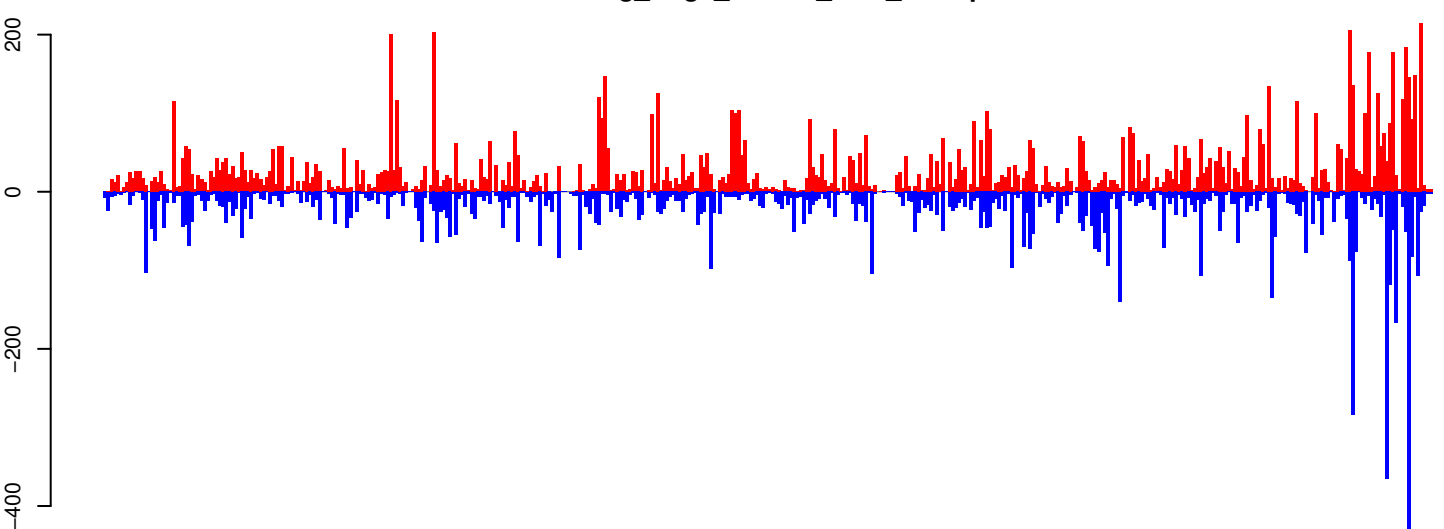
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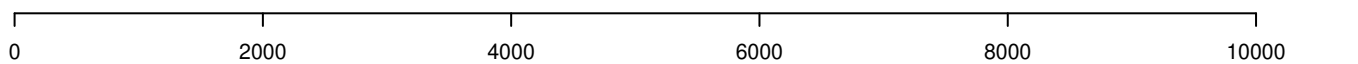


positive=617, negative=330, total=948

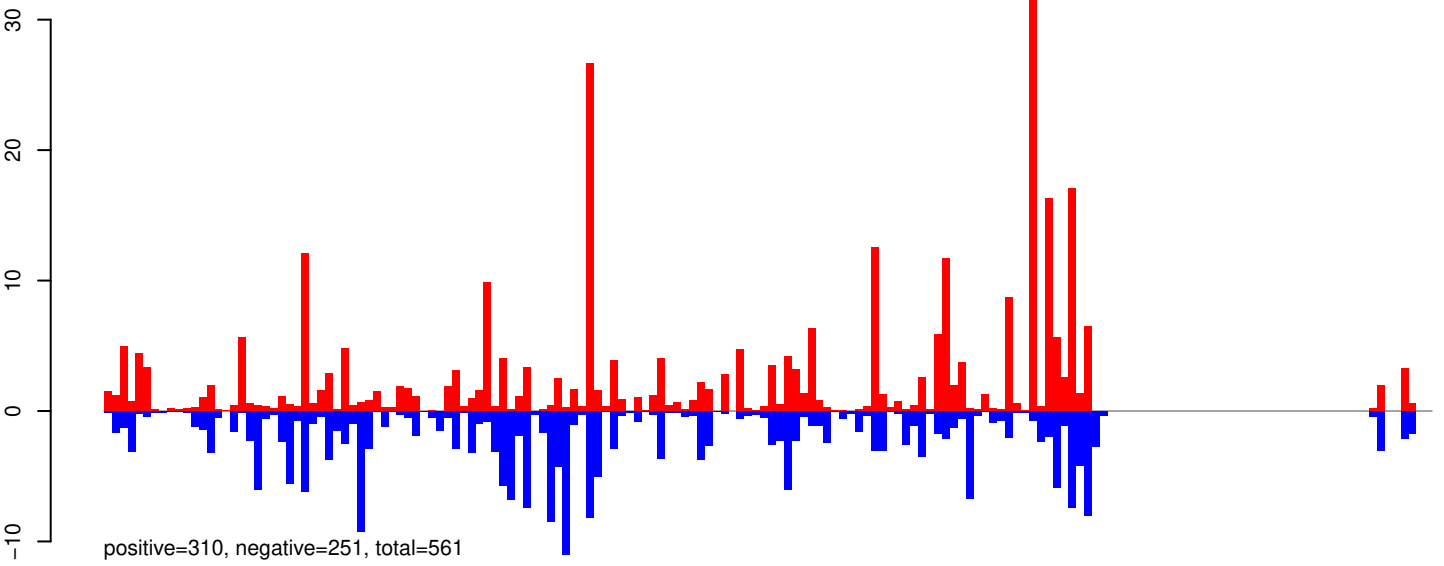
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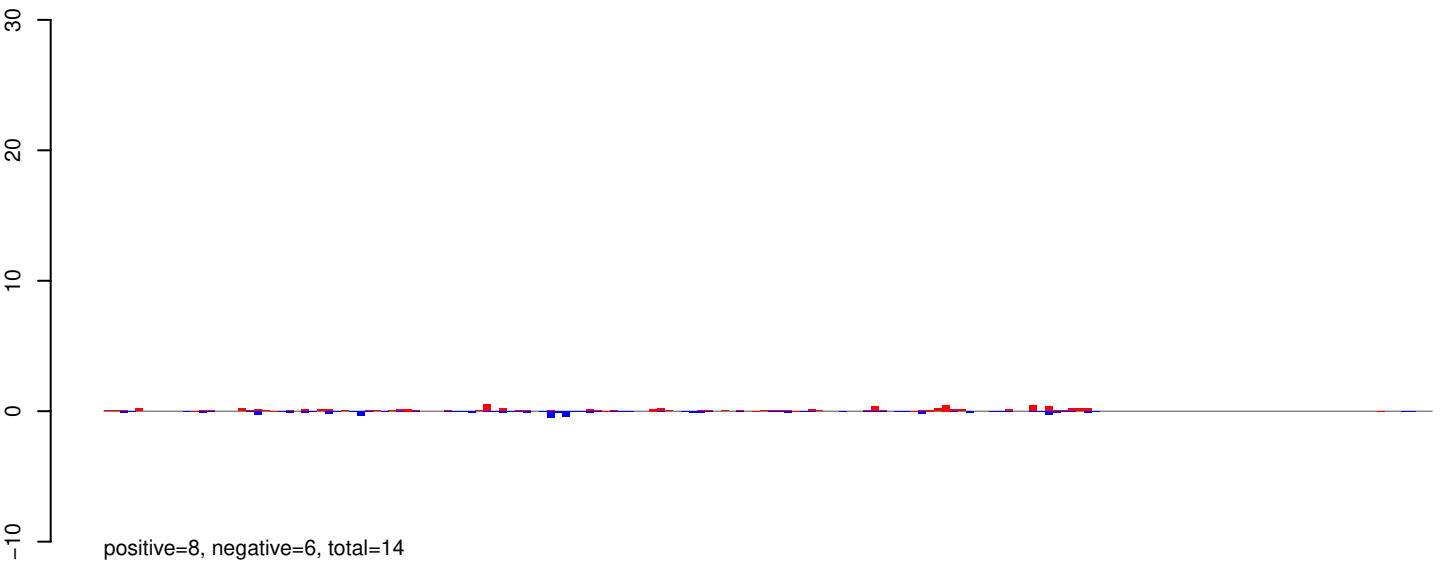
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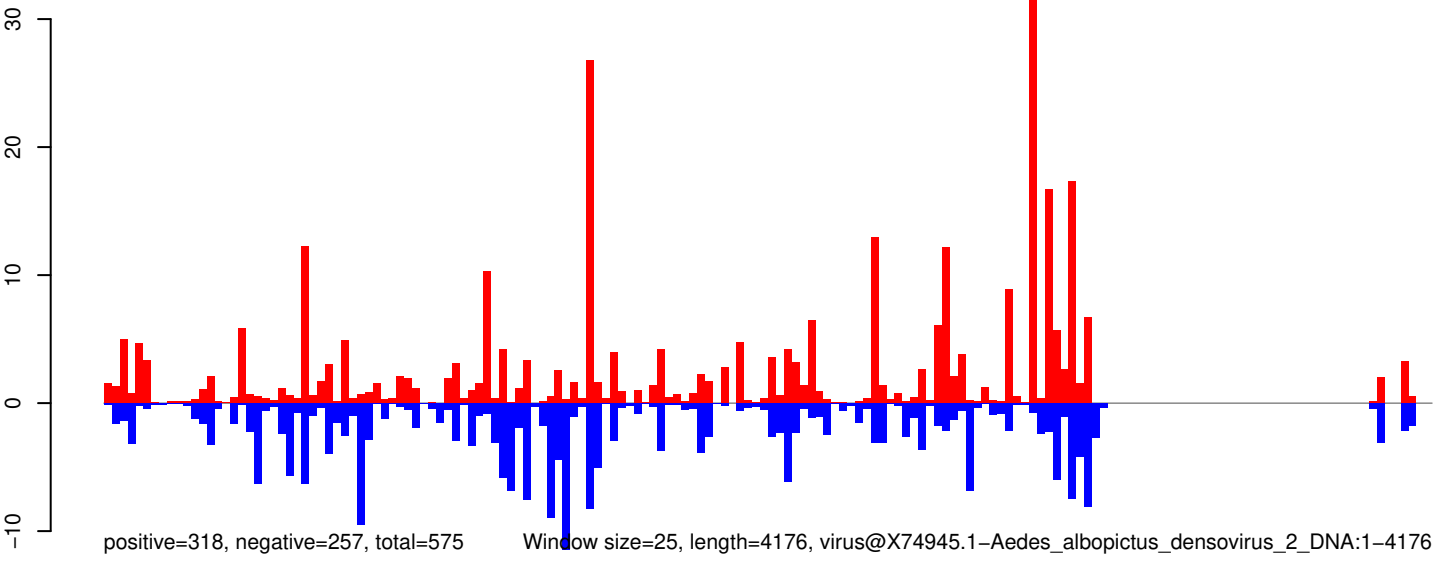
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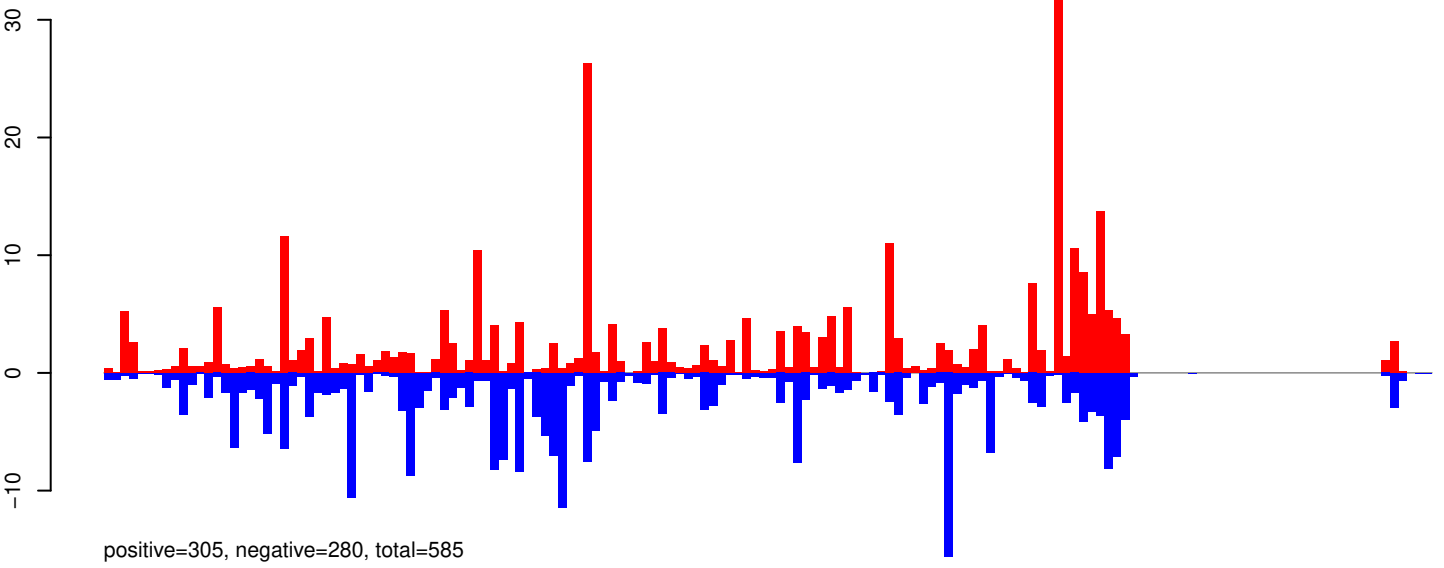
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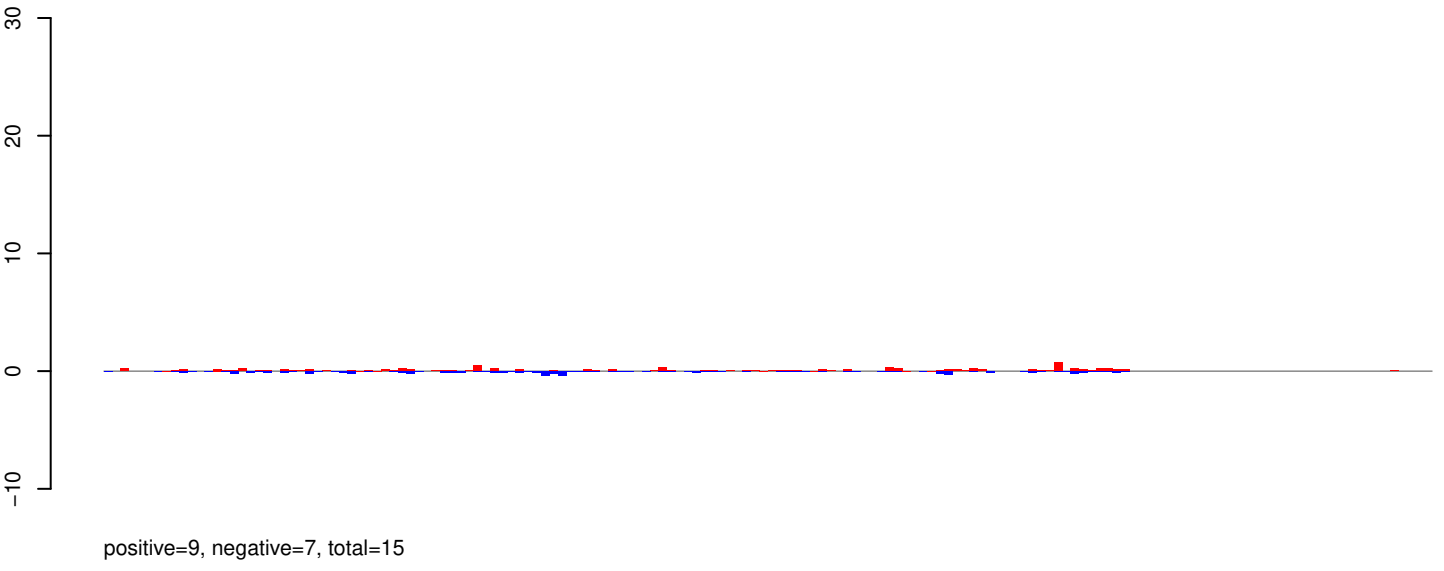
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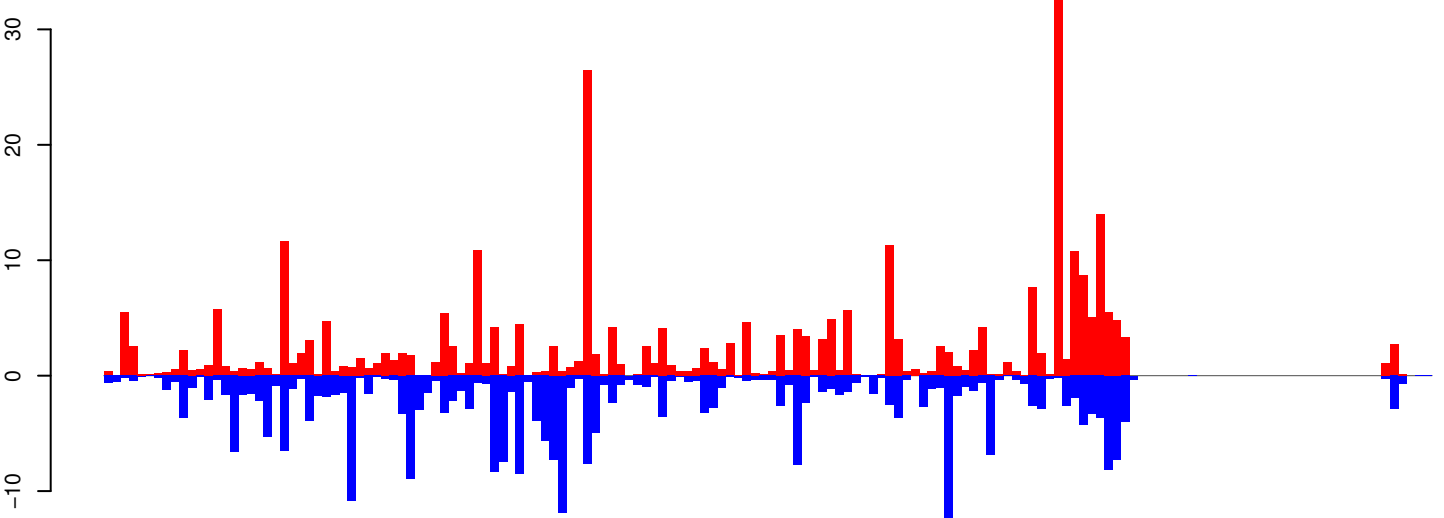
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



AeAeg_Aag2_R3D1B_r123_ZA.rep



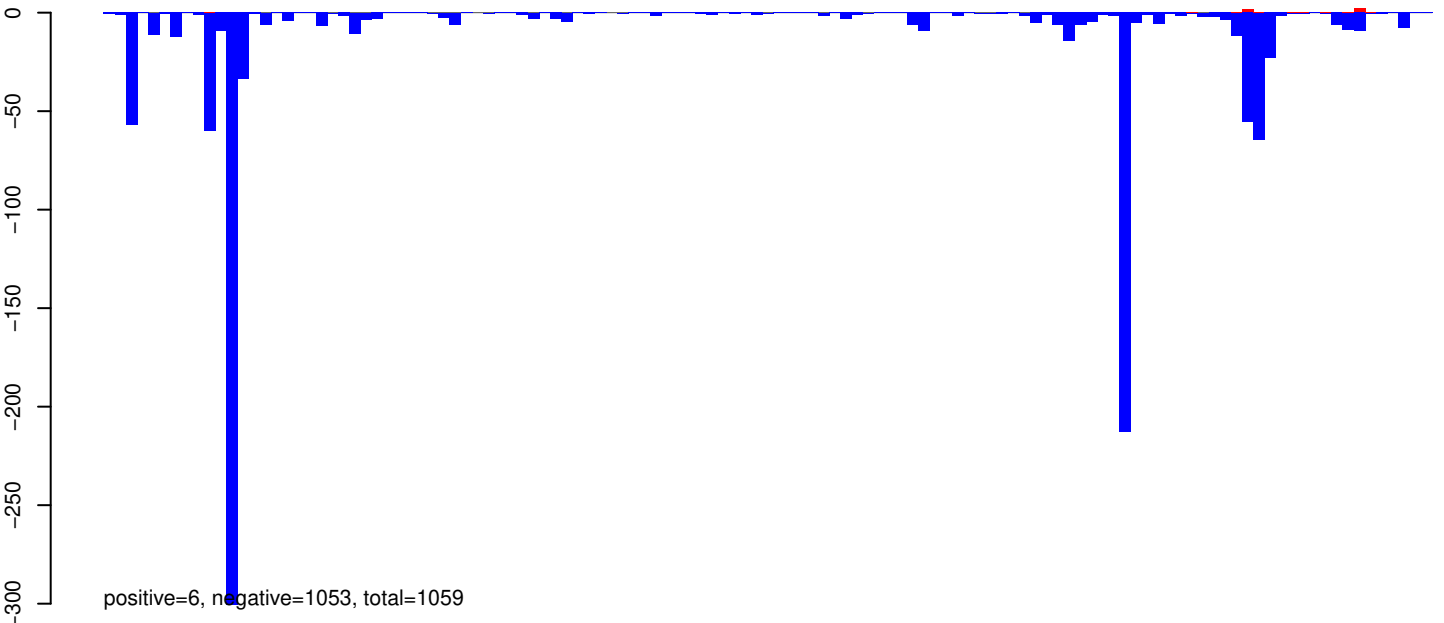
Window size=25, stride=1, step=1, @MH550148.1-Aedes albopictus_densovirus_isolate_NC01_L3_1_trimmed_contig_5889-RC:1-3937

0 1000 2000 3000 4000

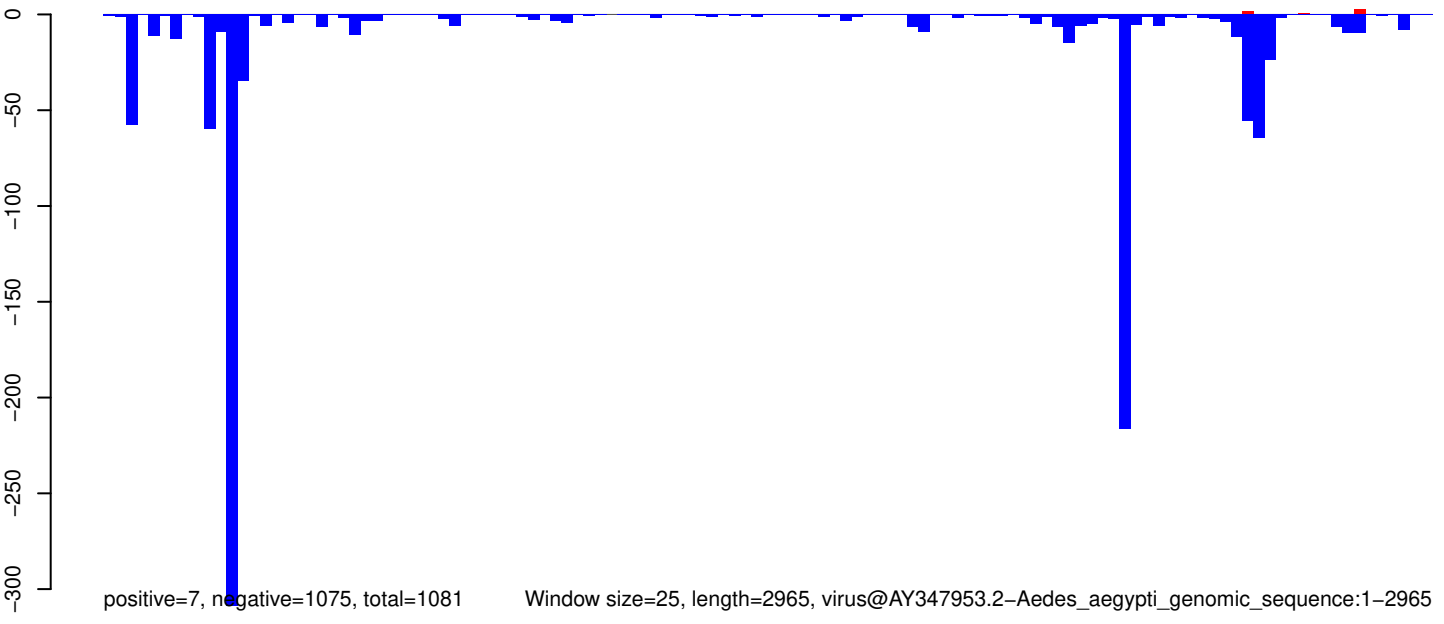
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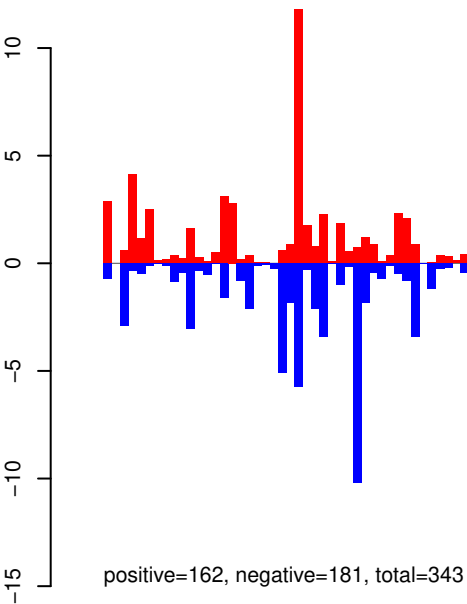
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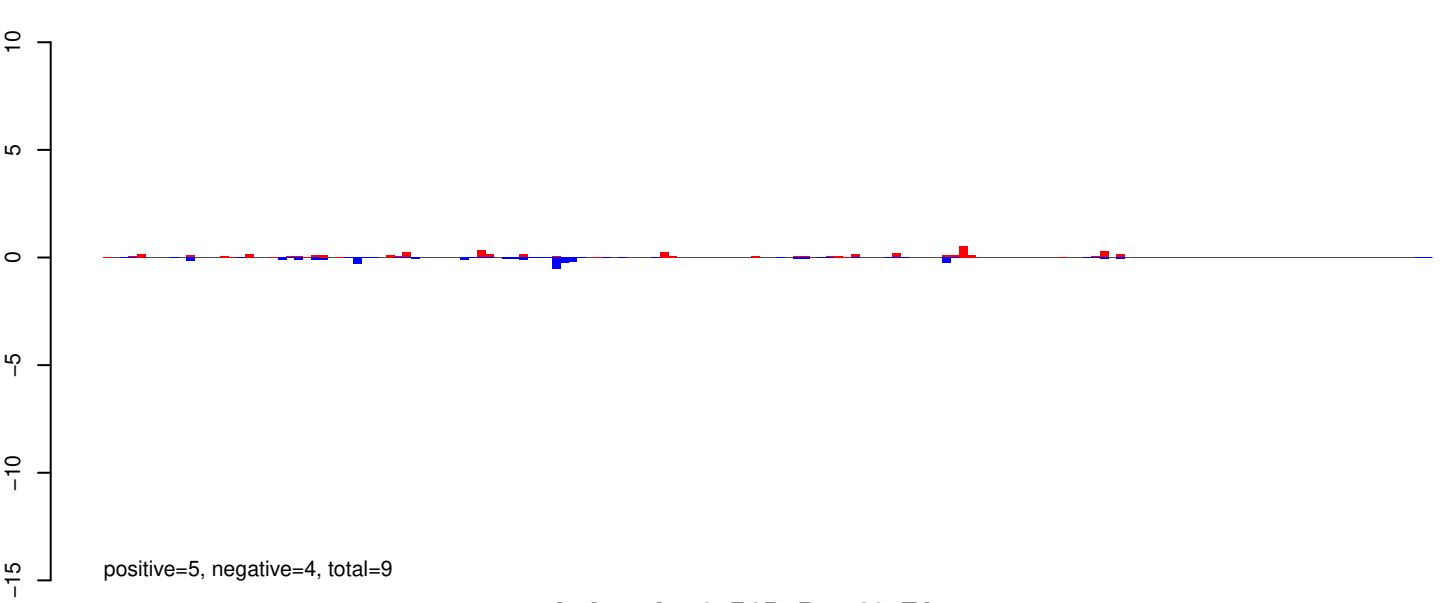
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0 500 1000 1500 2000 2500 3000

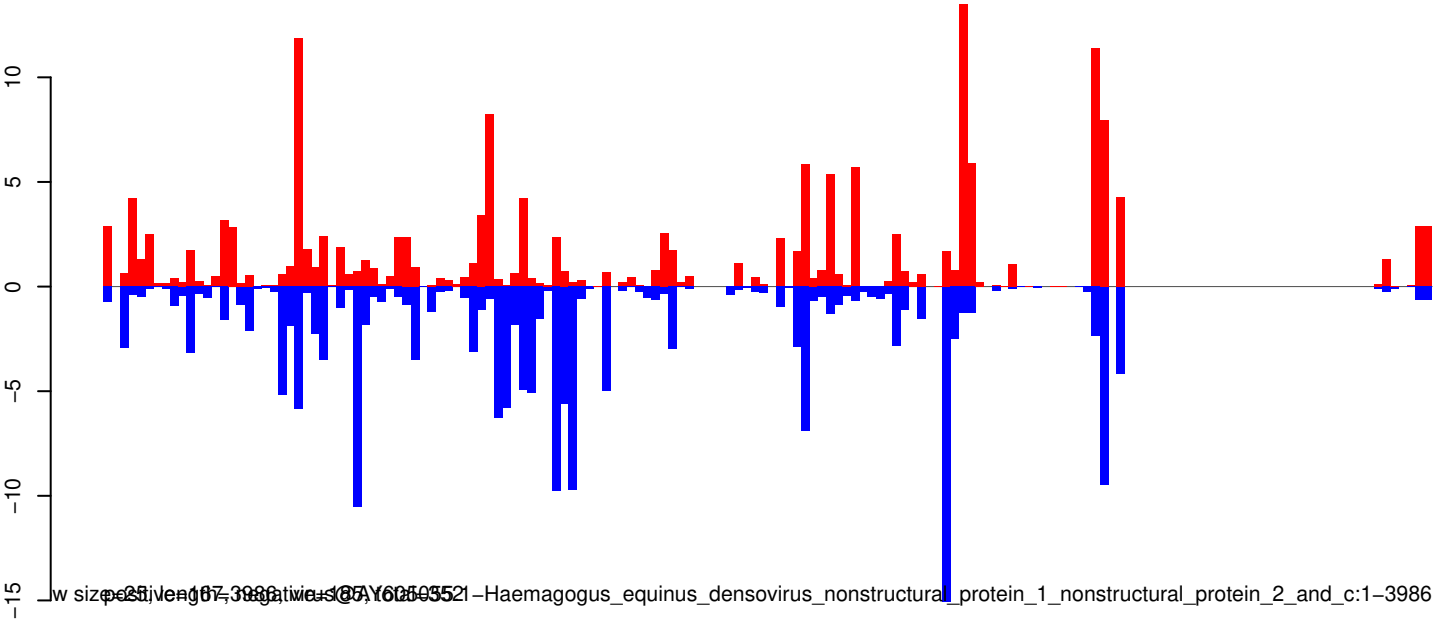
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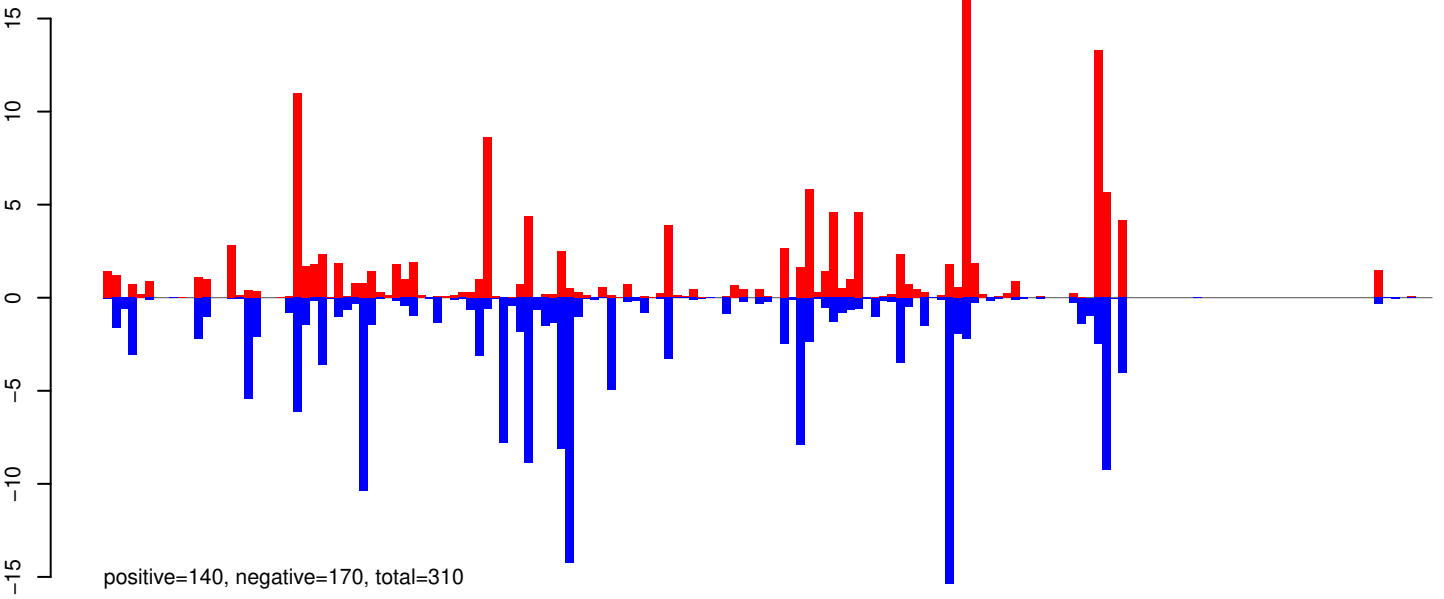


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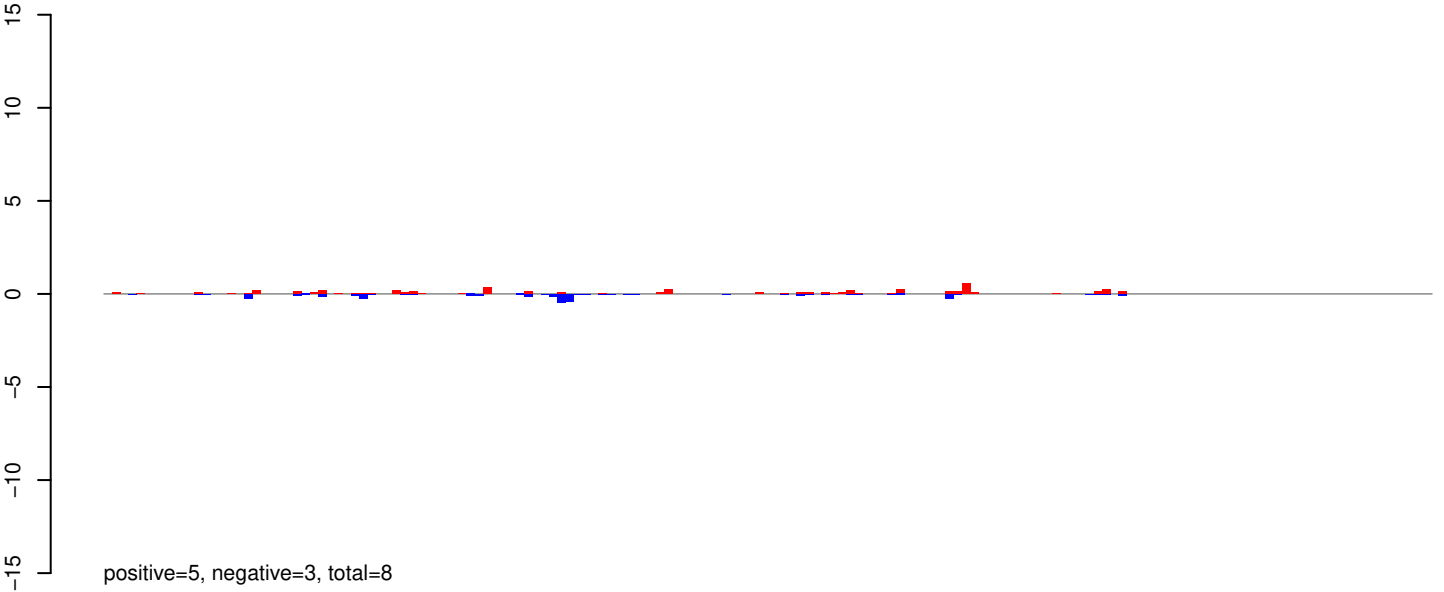


GenBank accession: AF162932.1 - Haemagogus equinus densovirus nonstructural protein 1 nonstructural protein 2 and c:1-3986

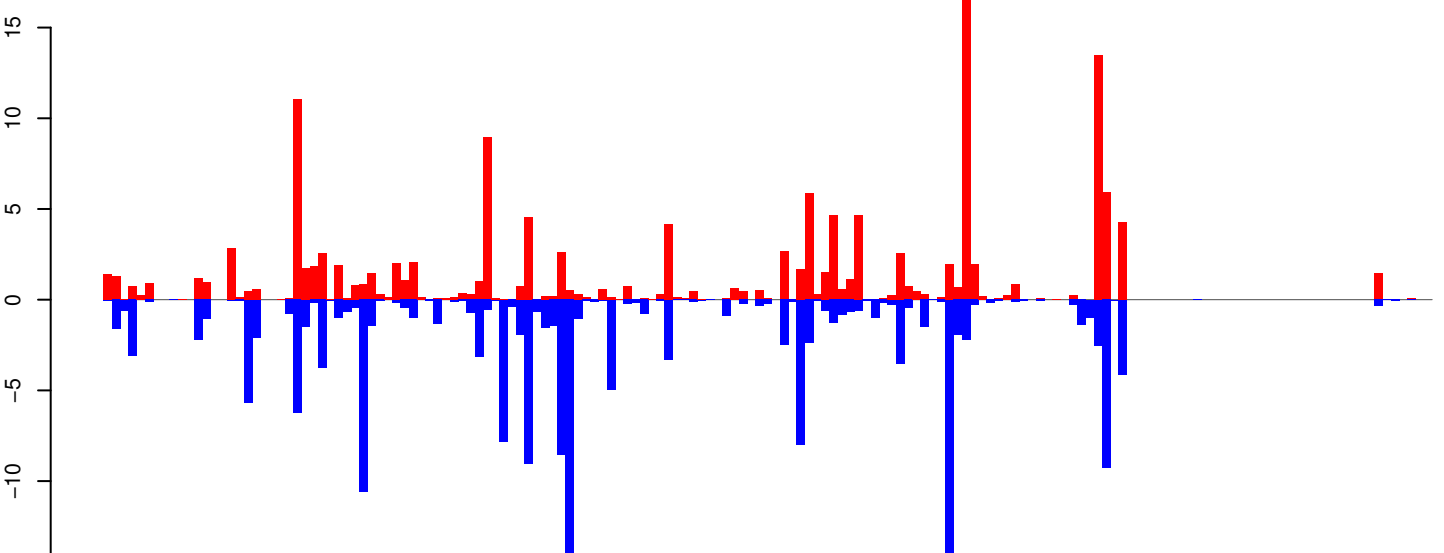
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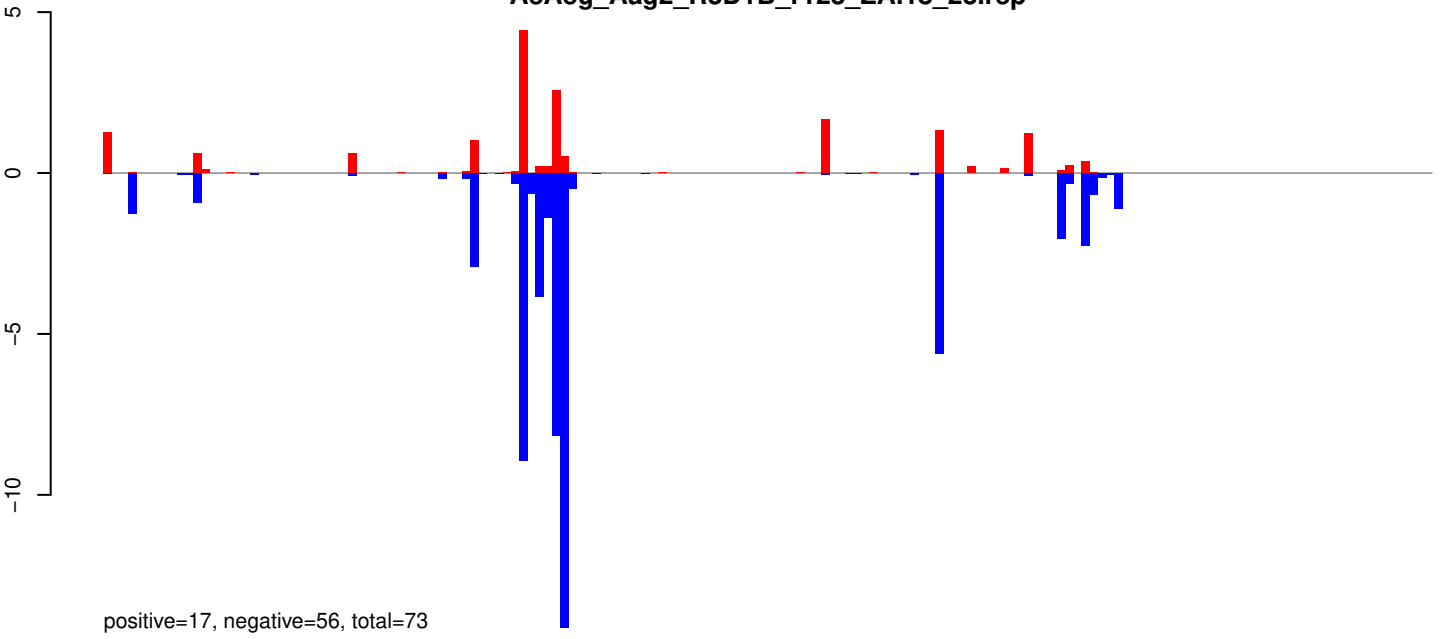
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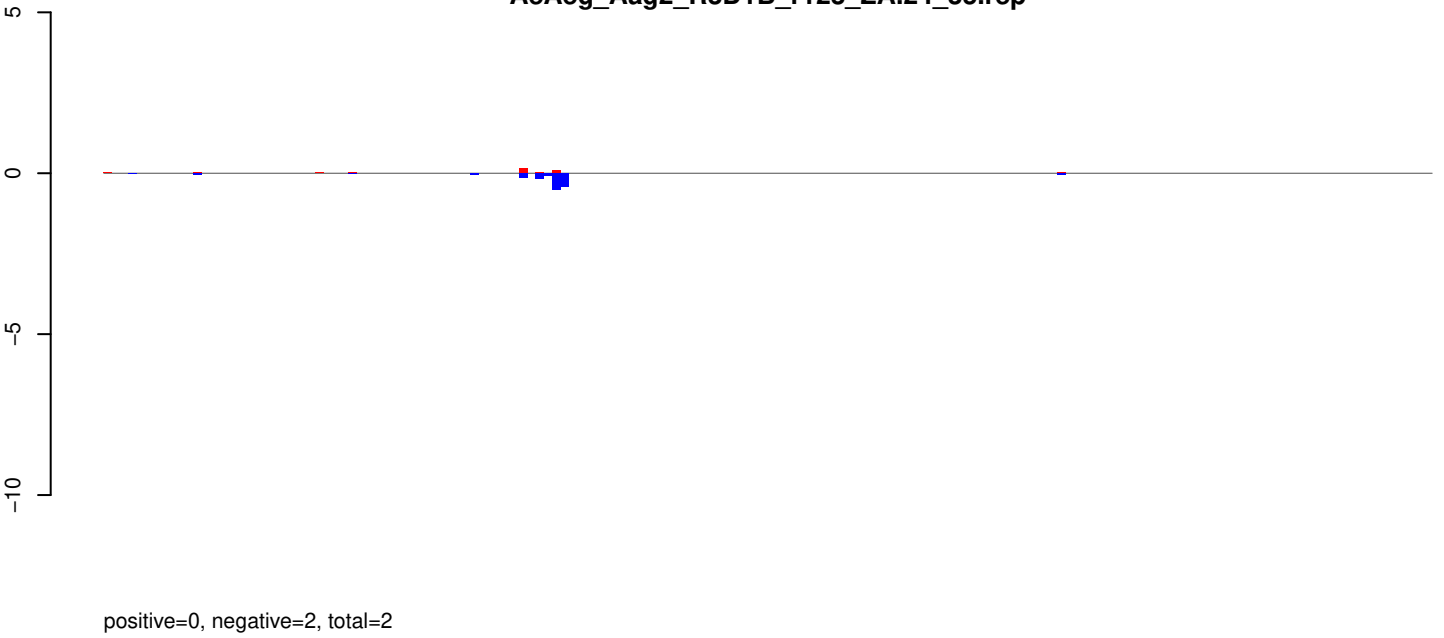
downloaded from <https://www.ncbi.nlm.nih.gov/Traces/trace/showTrace.cgi?db=GenBank&acc=AF198771> - Aedes albopictus_C6_36_cell_densovirus_non-structural_protein_1_non-structural_prot:1-4006

0 1000 2000 3000 4000

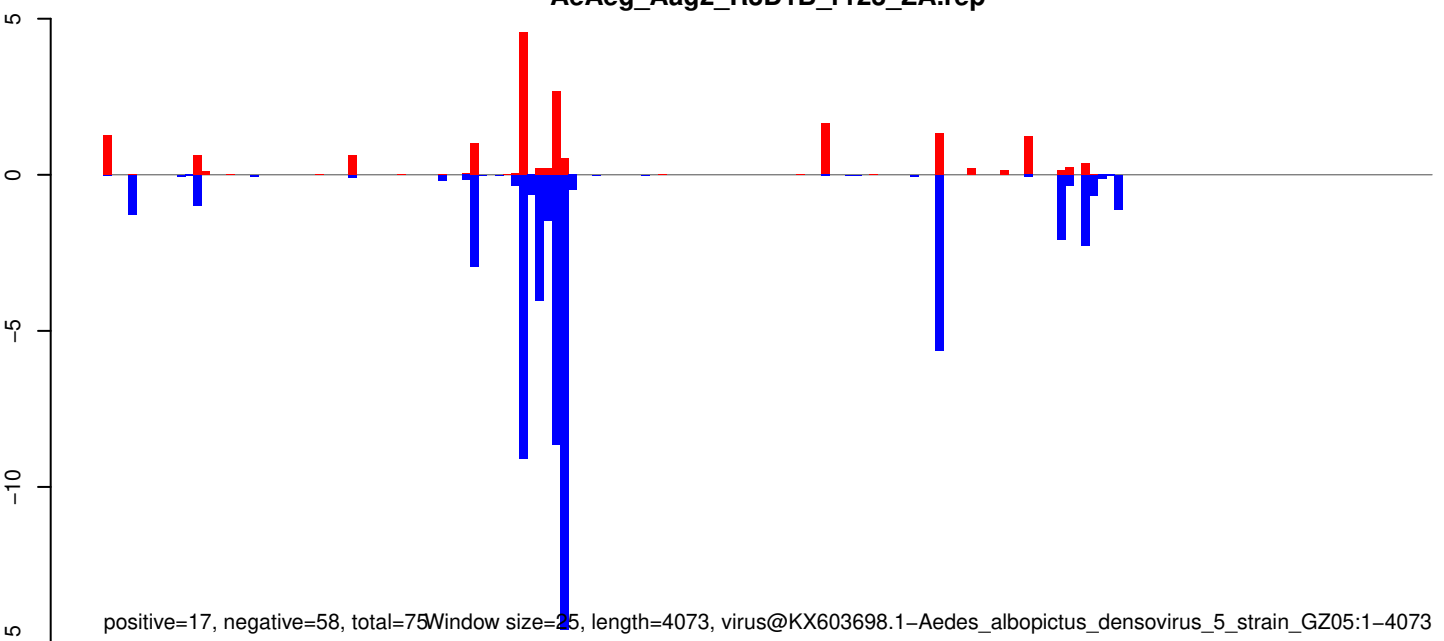
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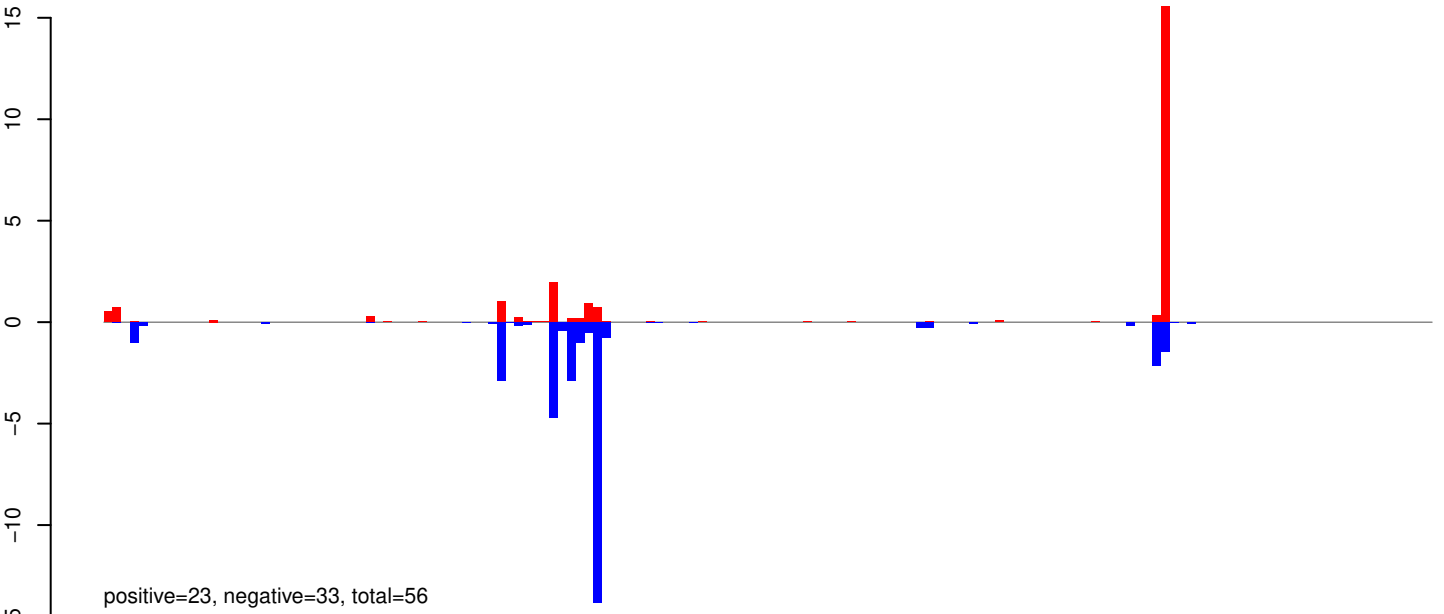


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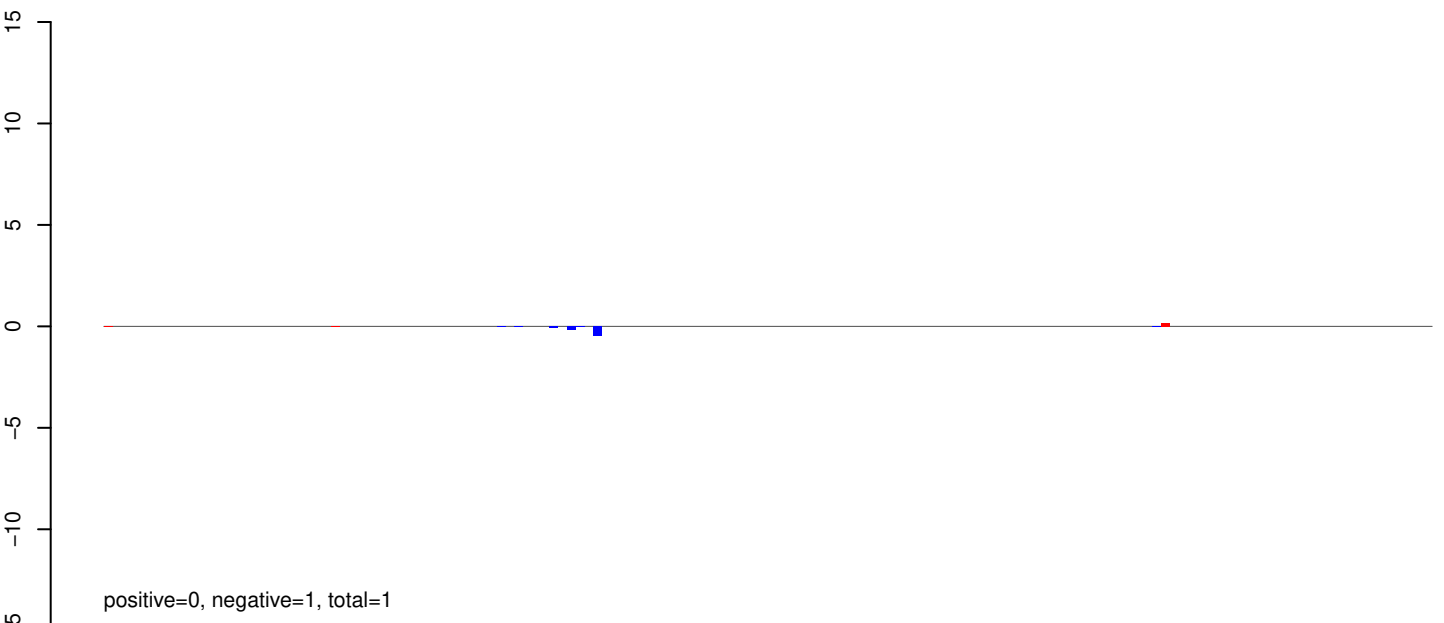


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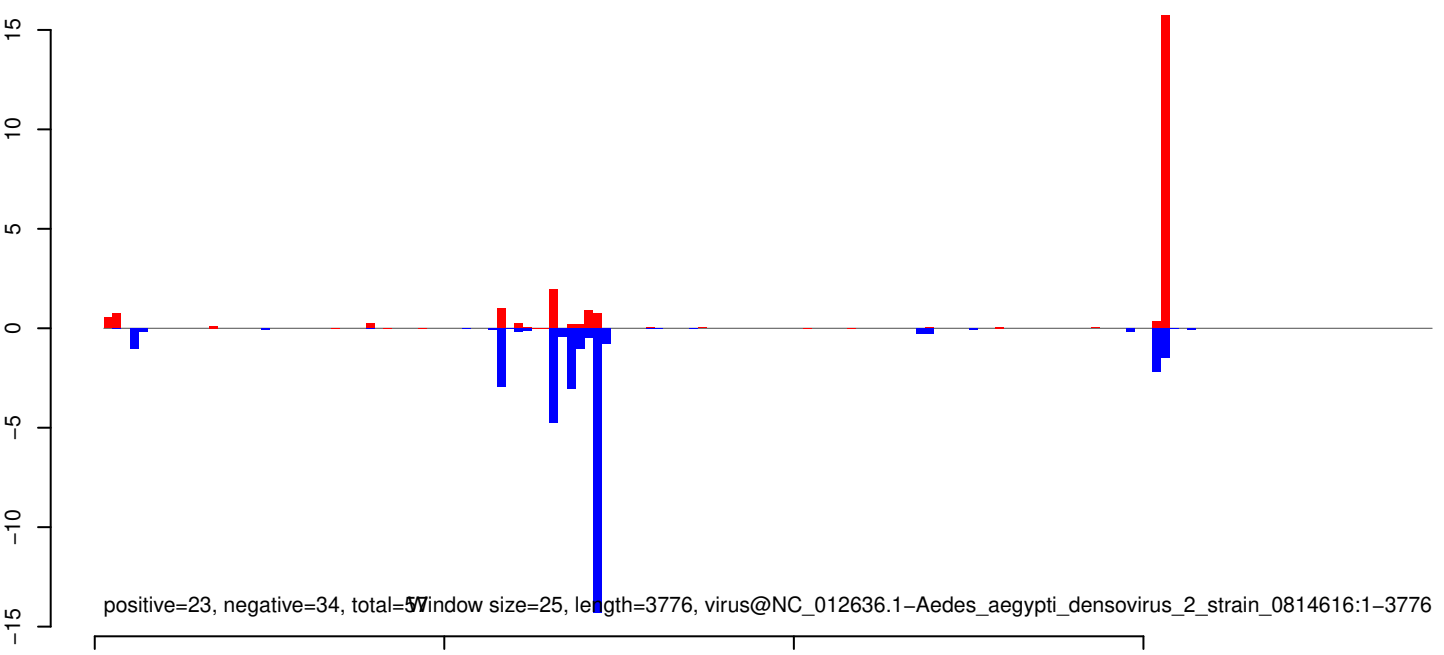
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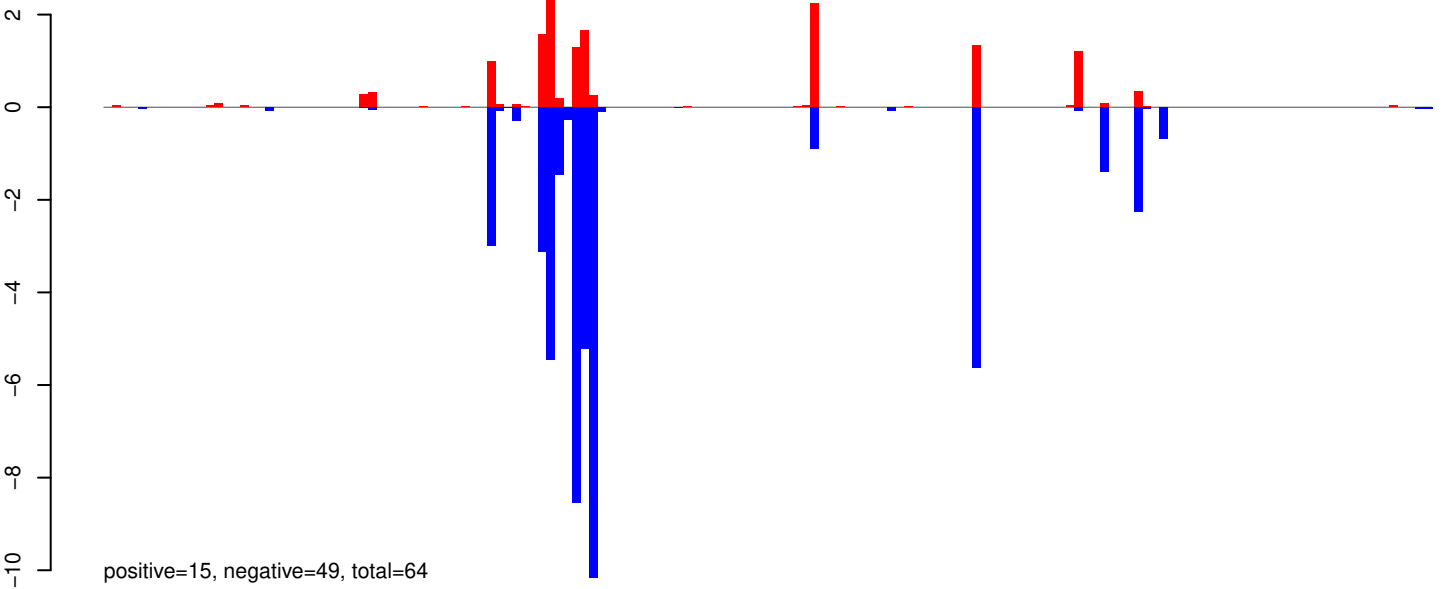
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AeAeg_Aag2_R3D1B_r123_ZA.rep



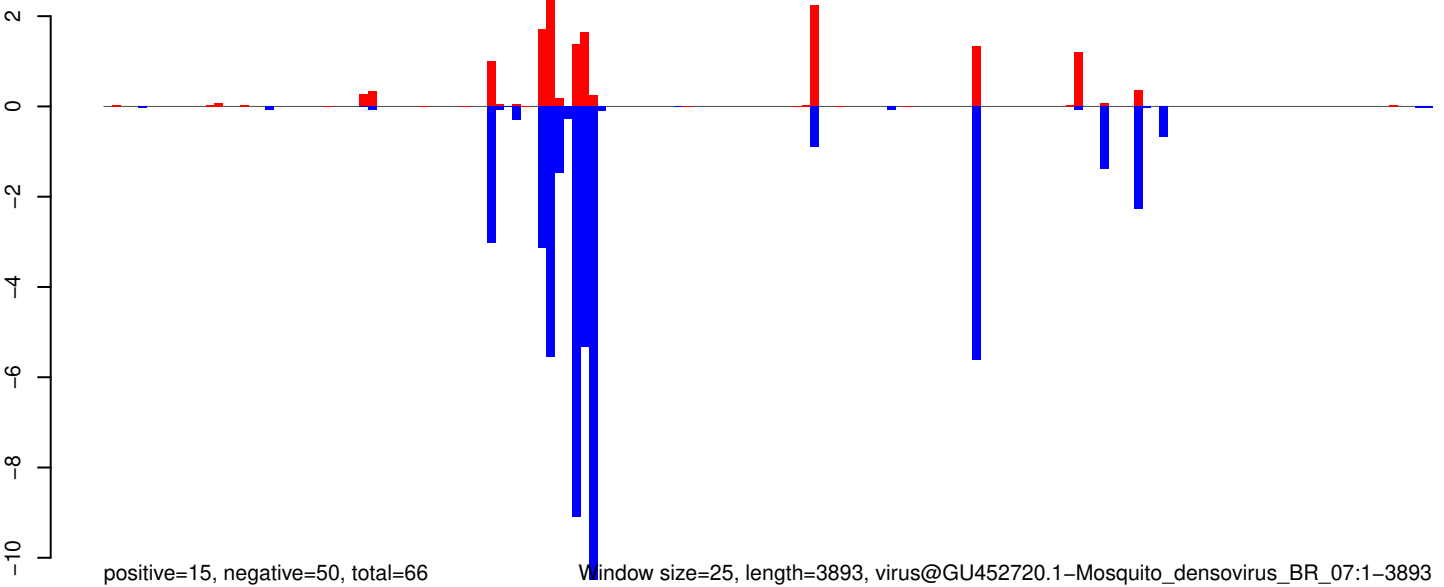
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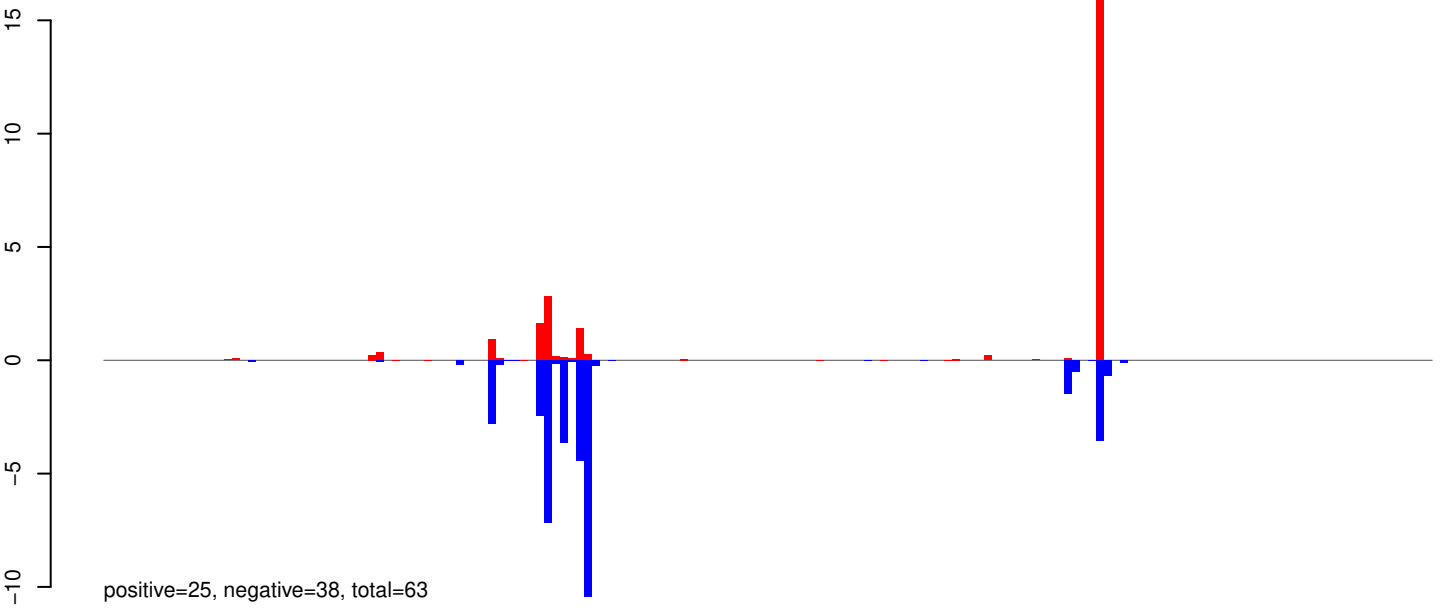
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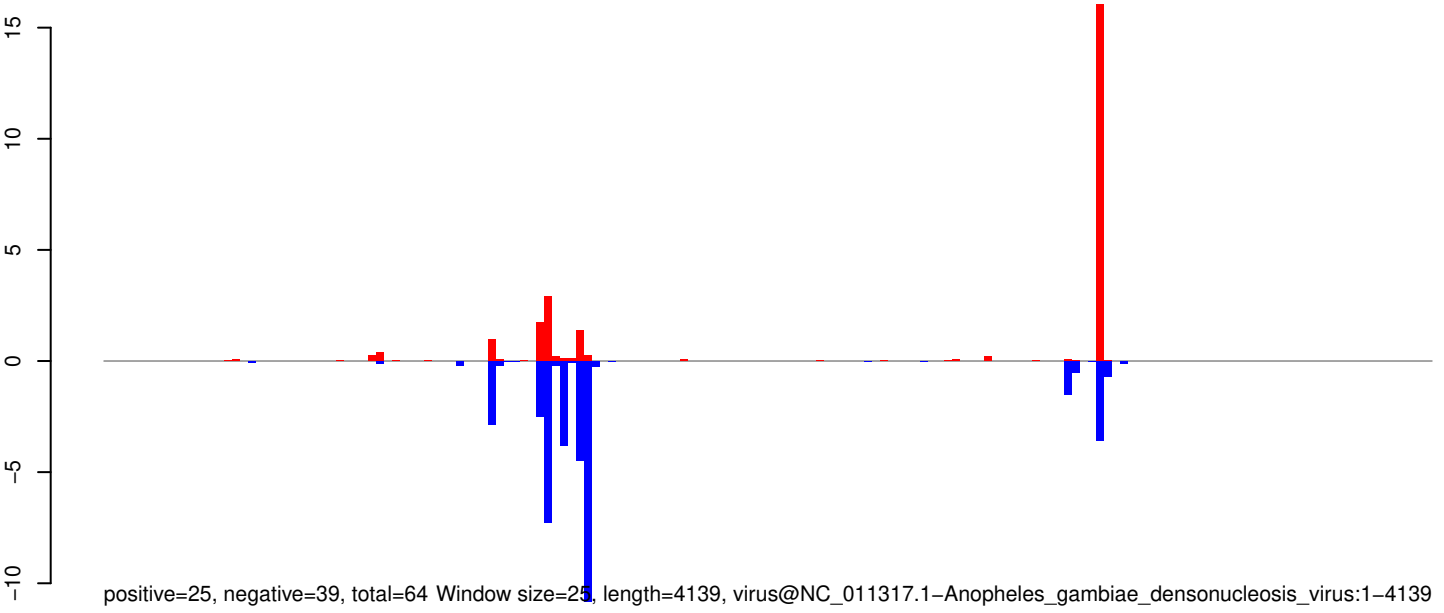
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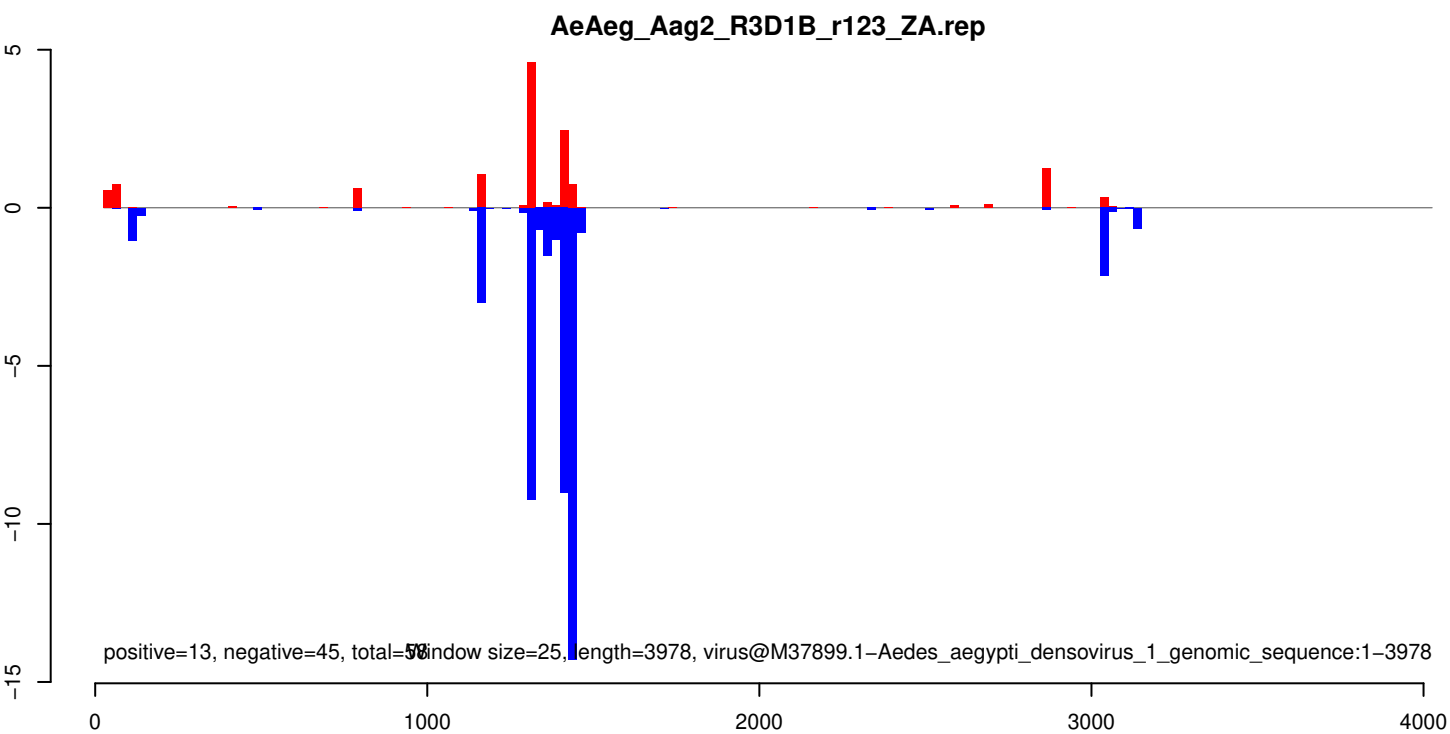
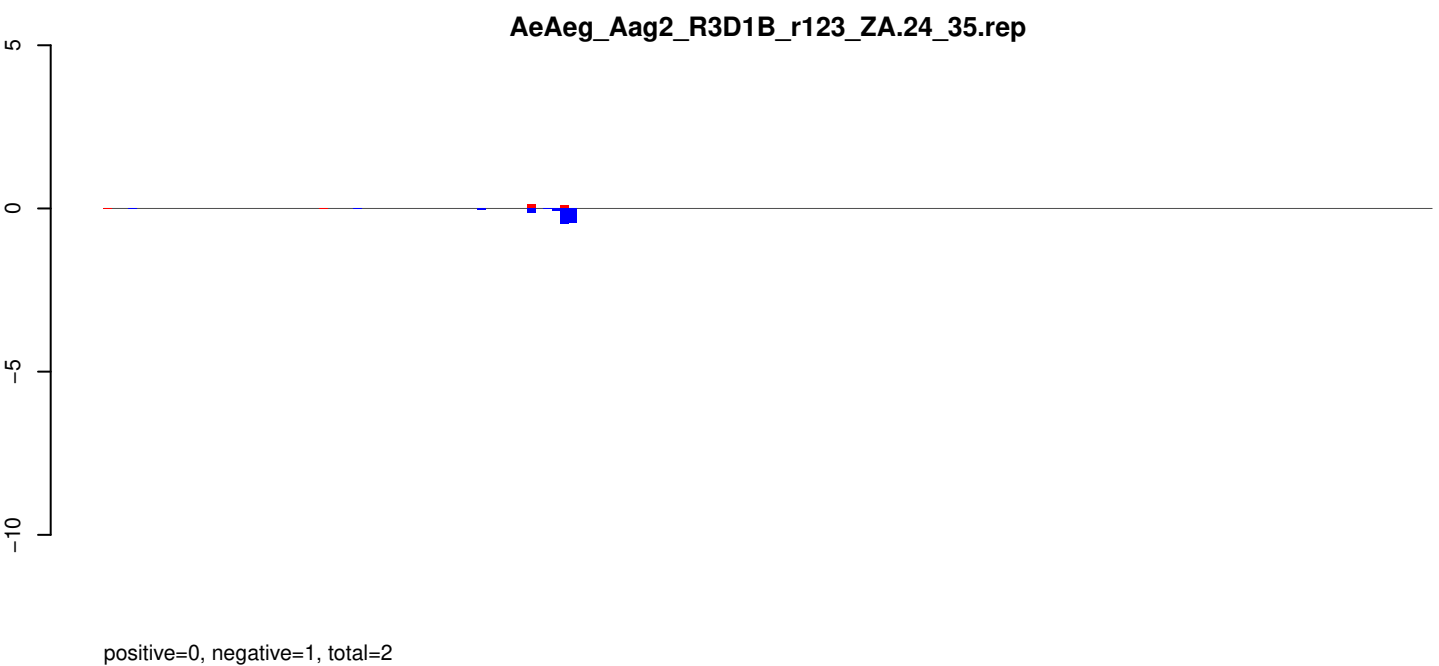
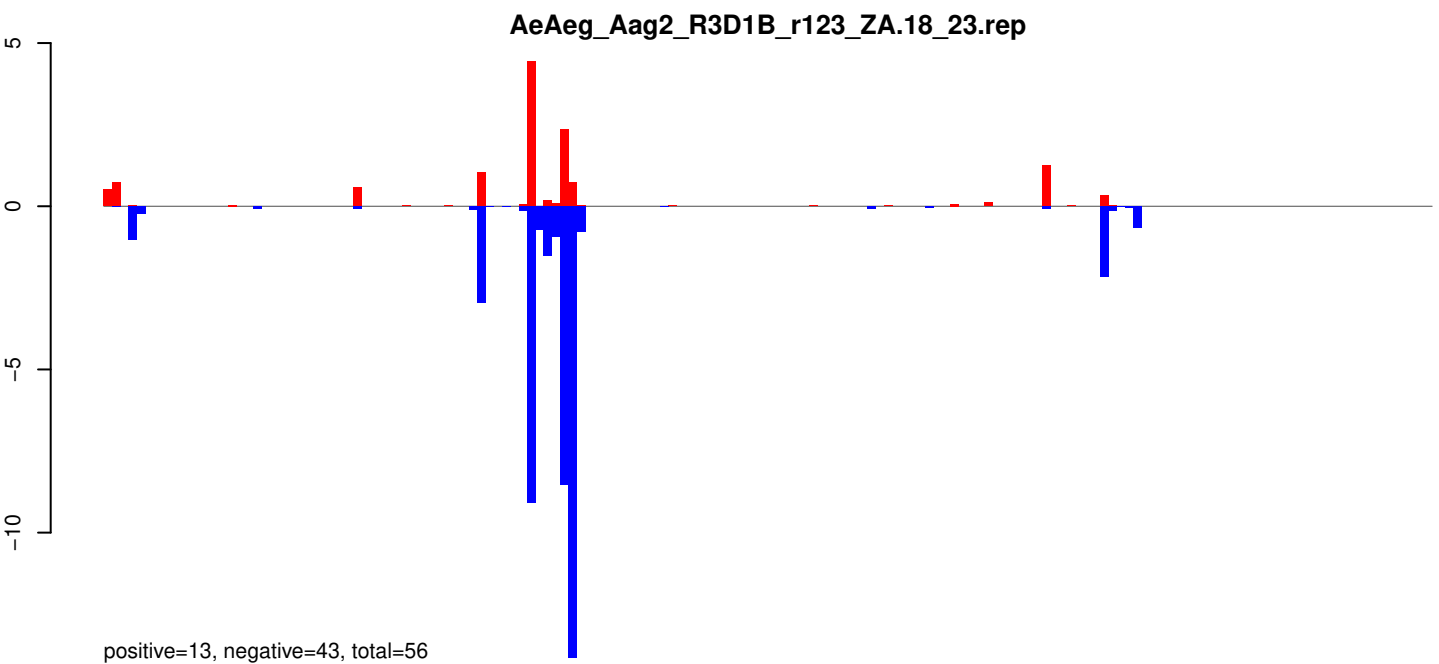


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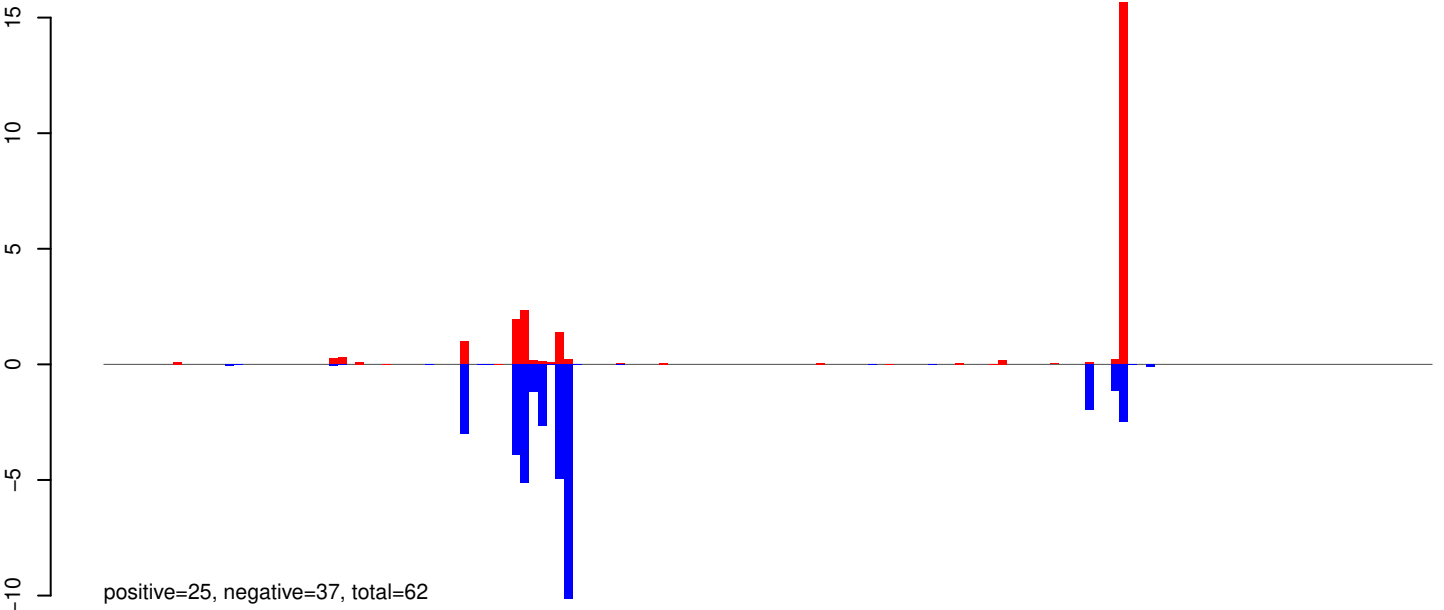


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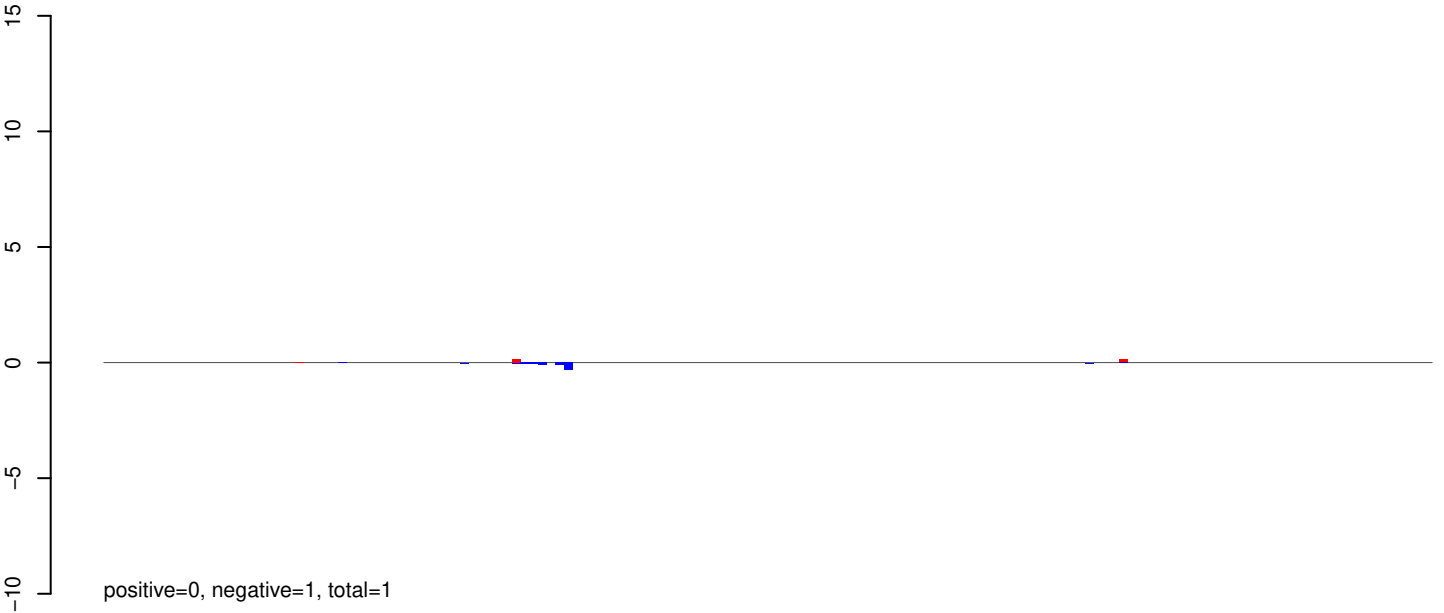




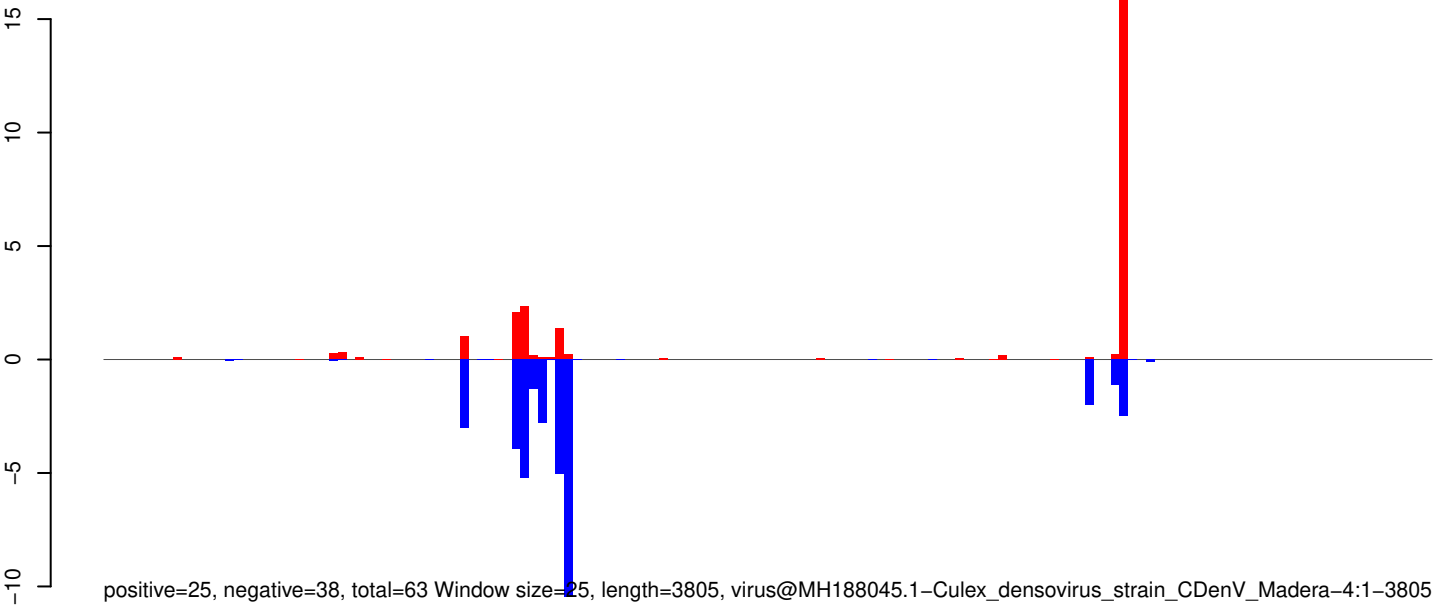
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep

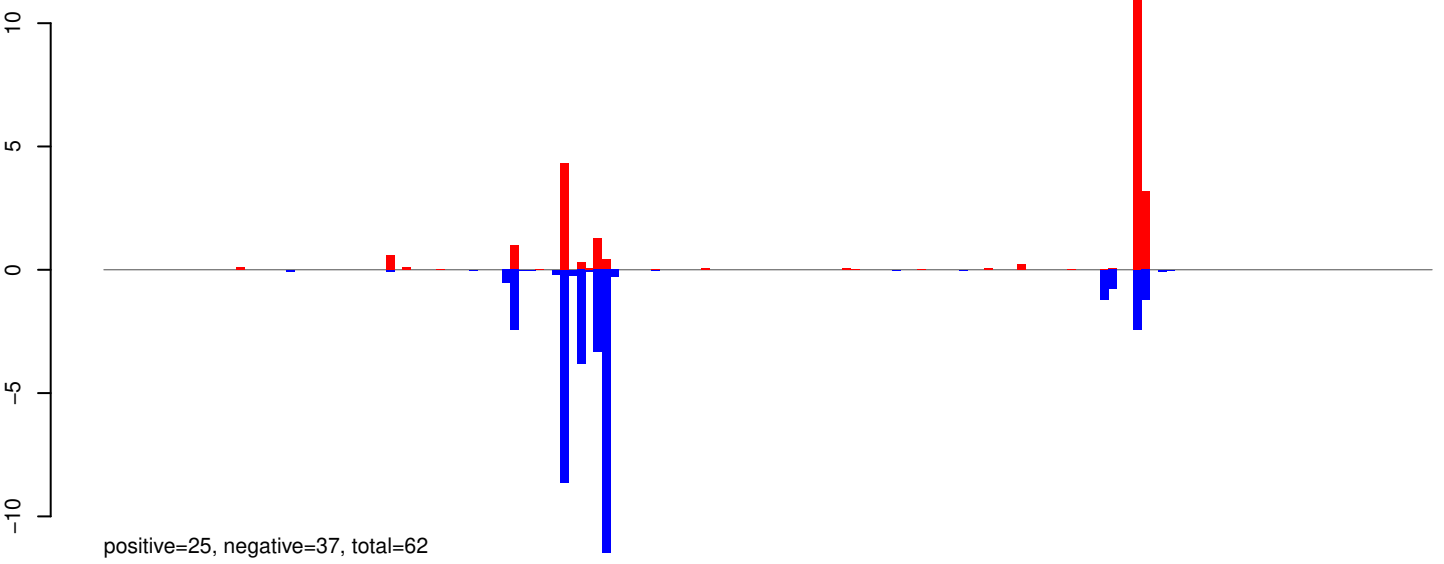


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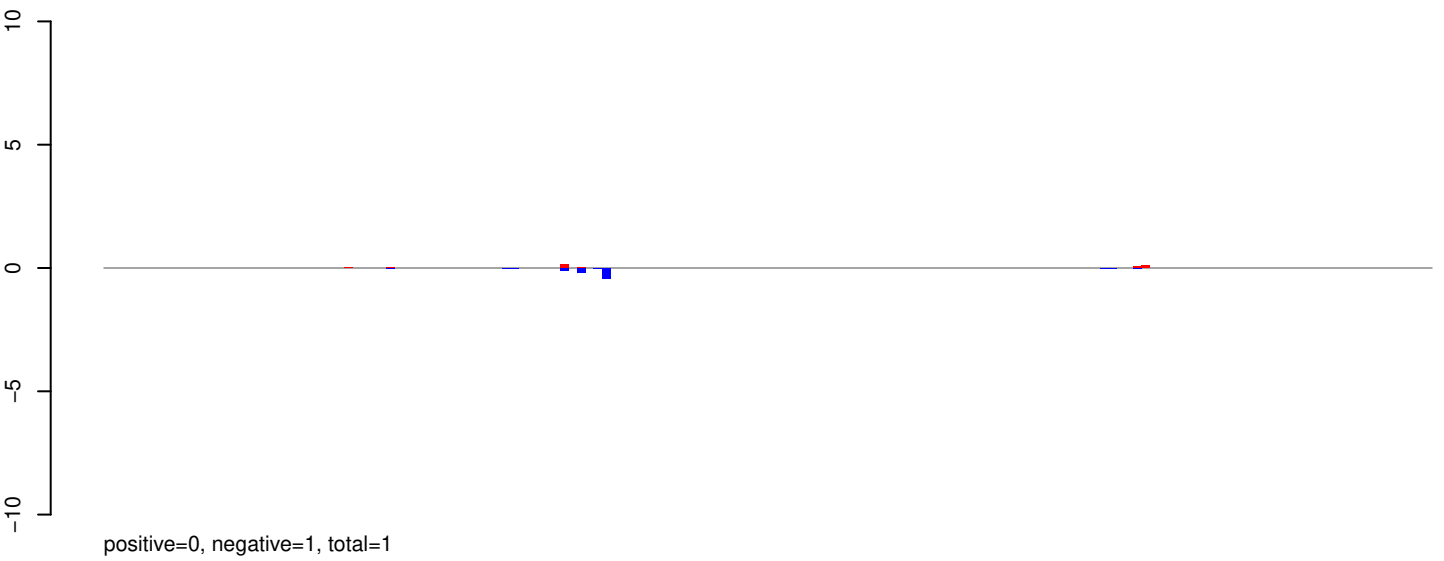


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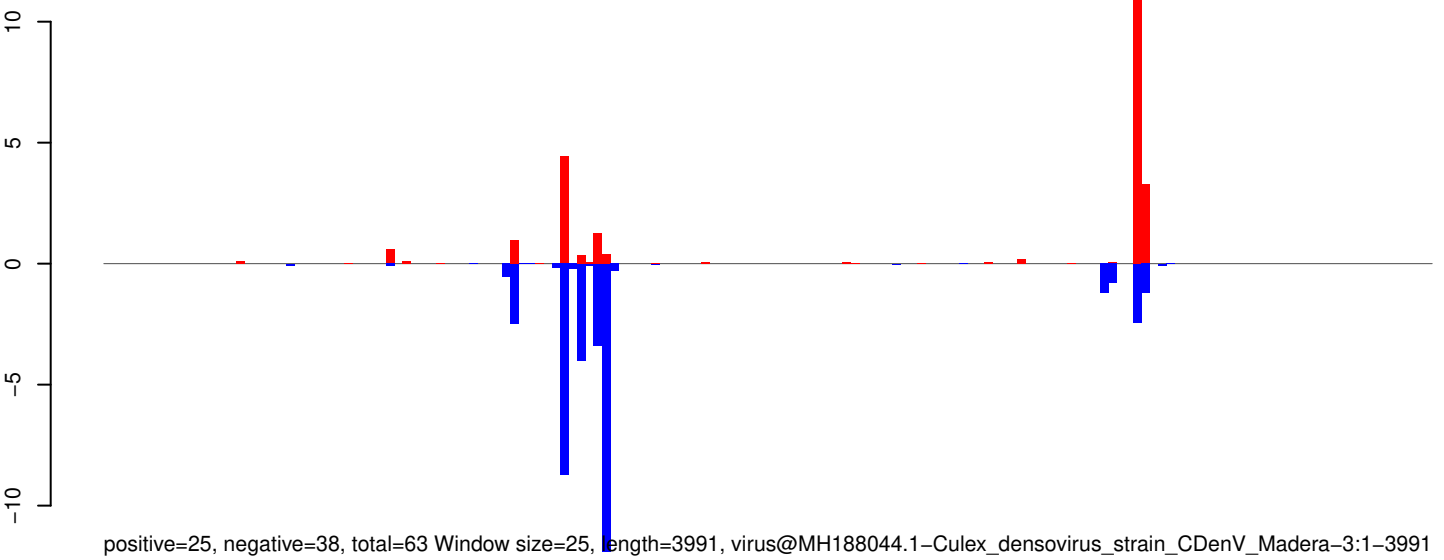
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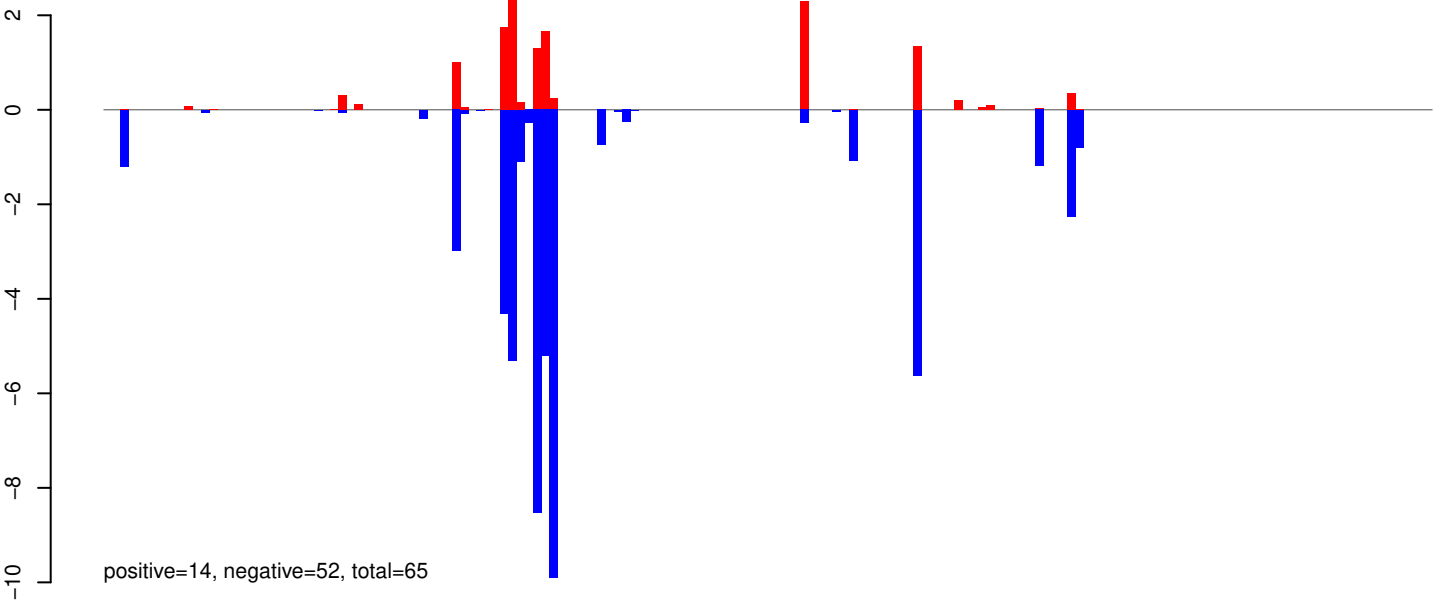


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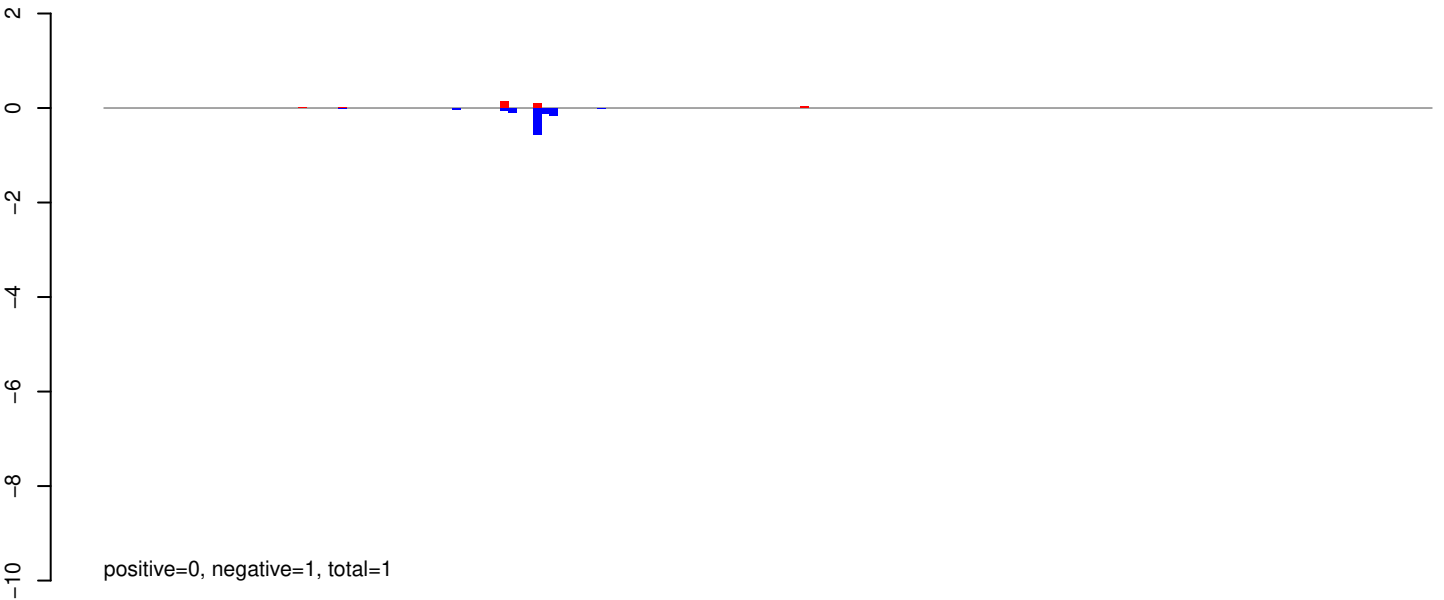


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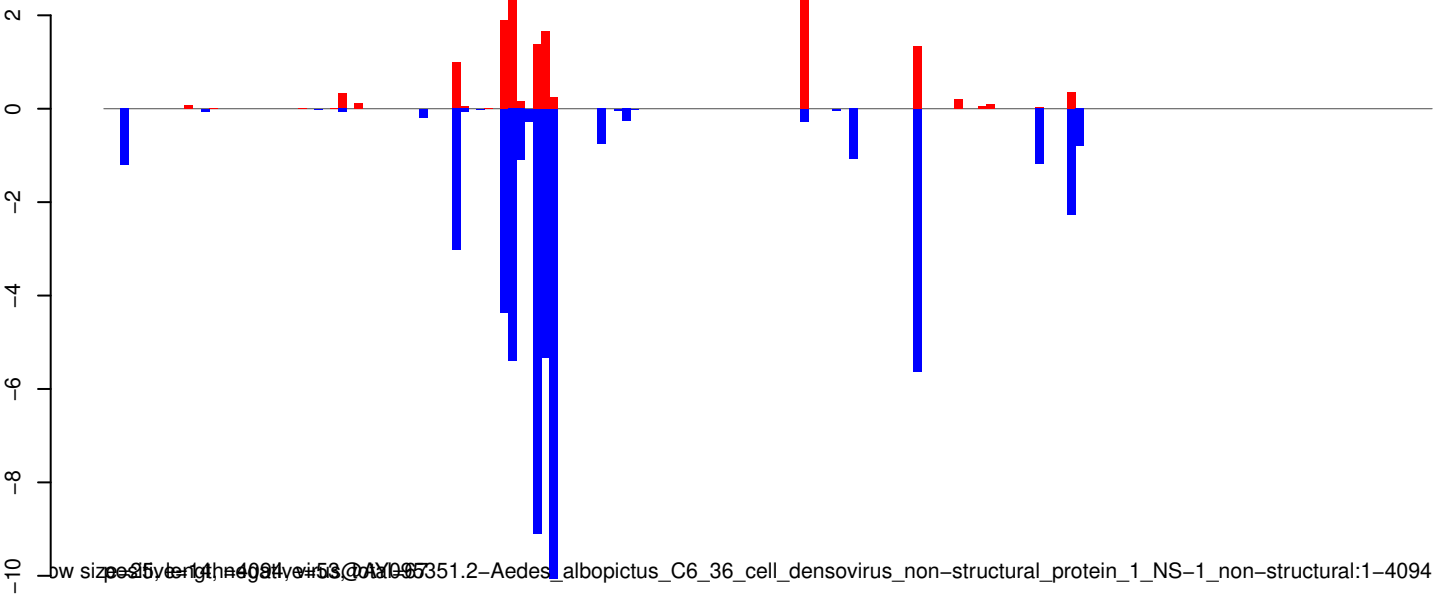
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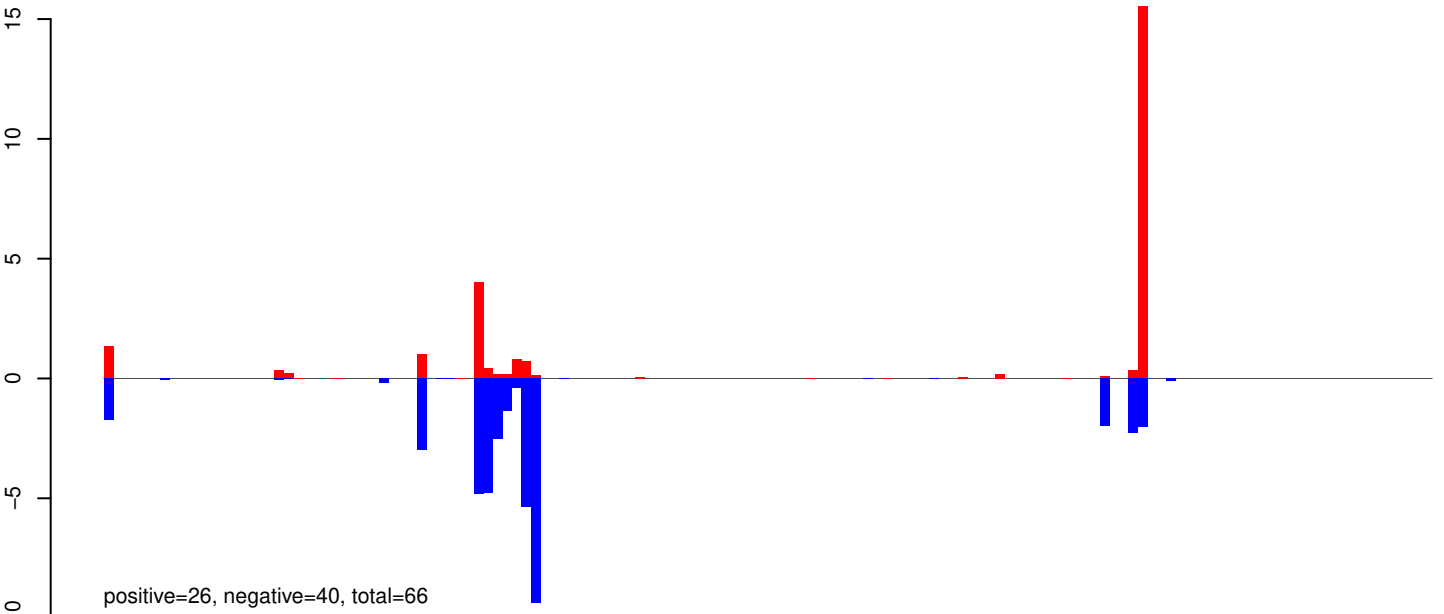


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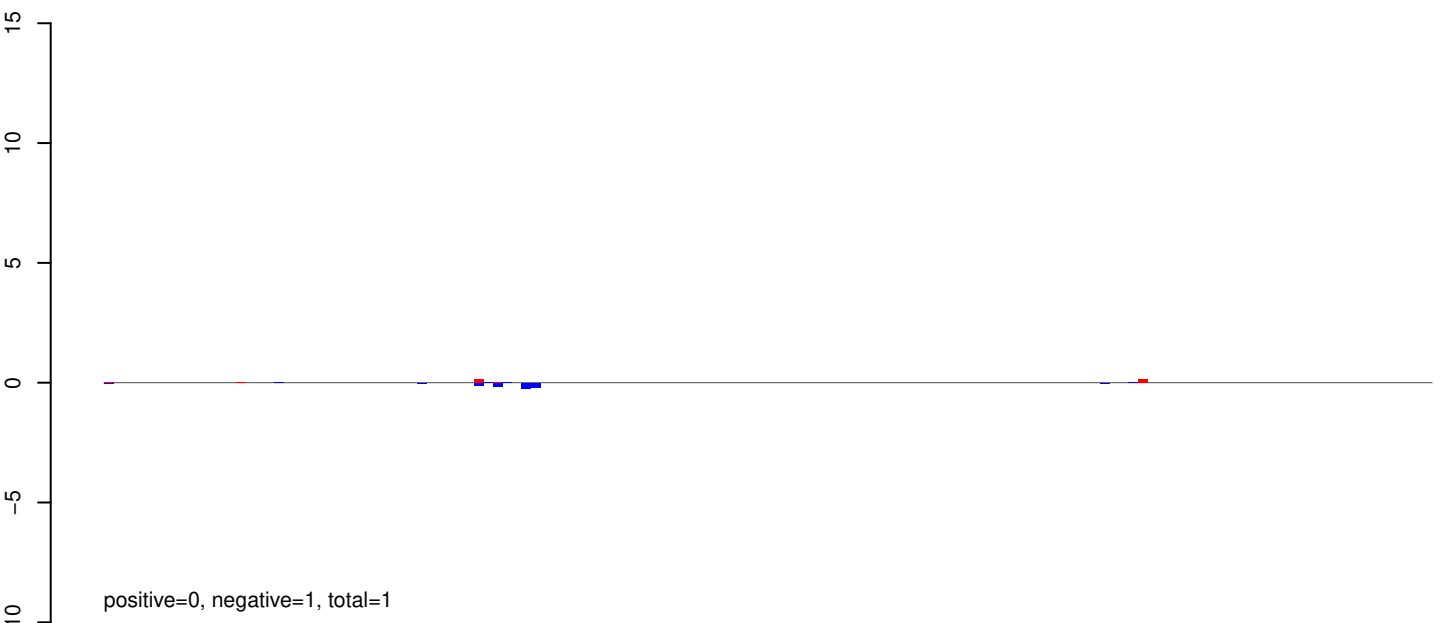


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Aedes albopictus_C6_36_cell_densovirus_non-structural_protein_1_NS-1_non-structural:1-4094

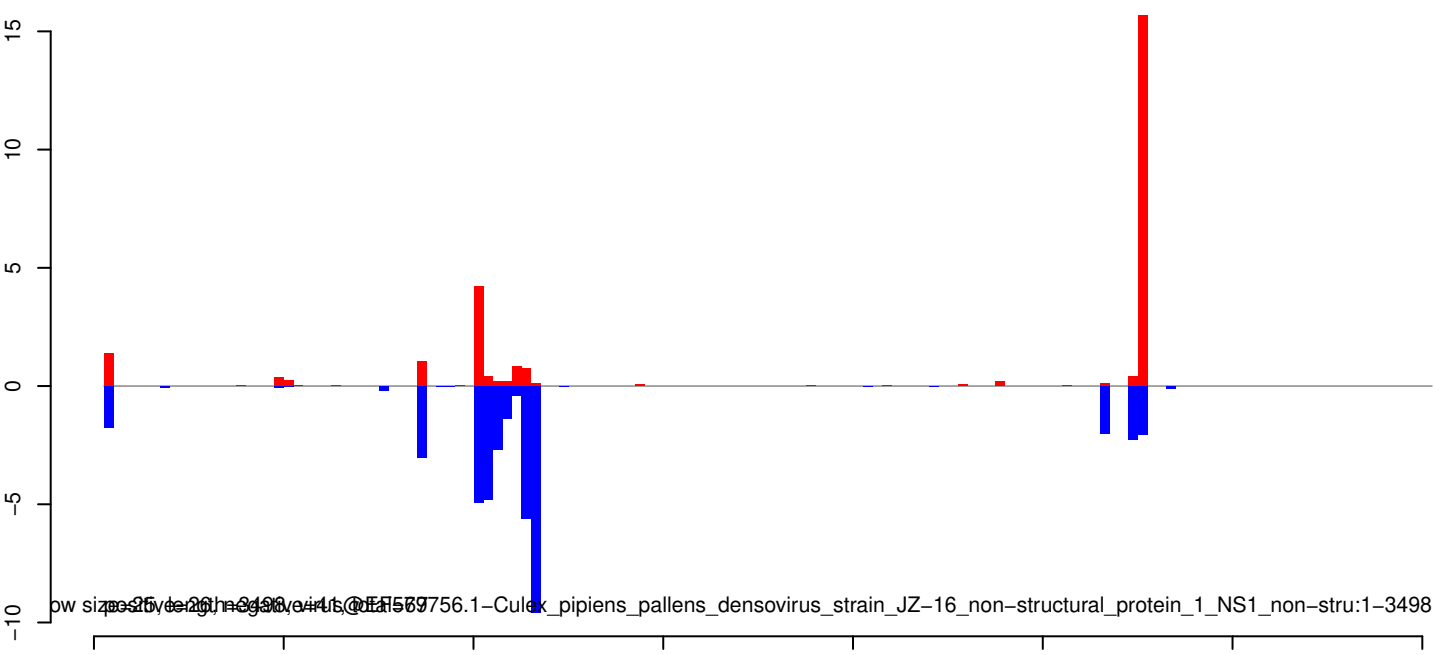
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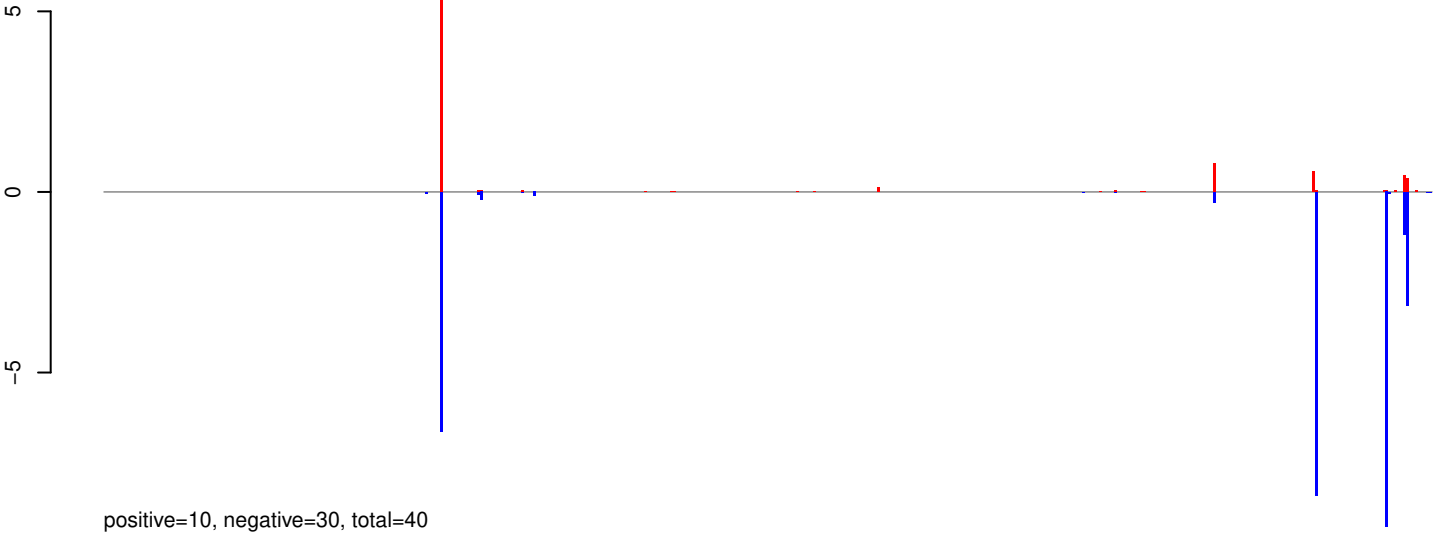
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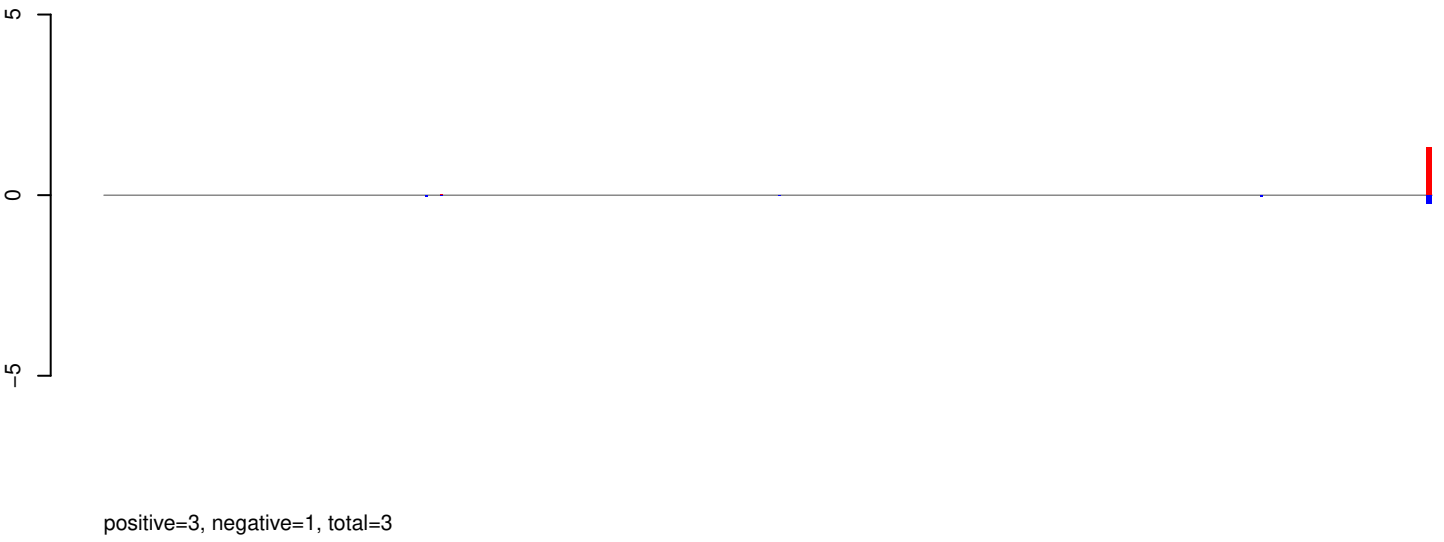
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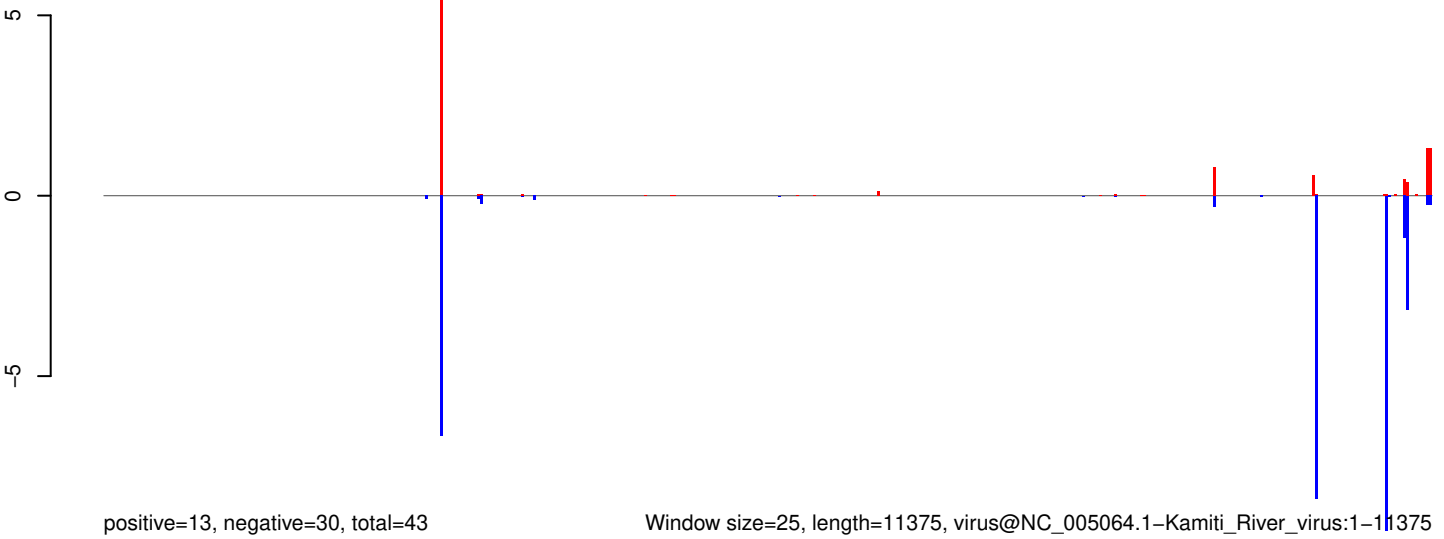
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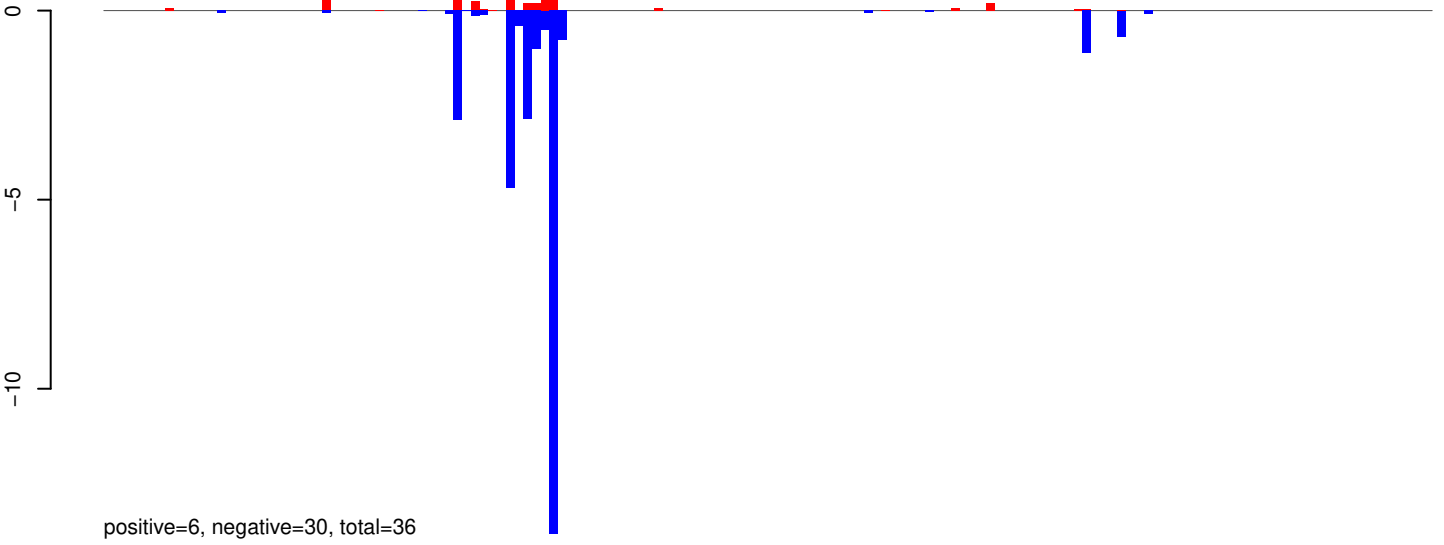
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Window size=25, length=11375, virus@NC_005064.1-Kamiti_River_virus:1-11375

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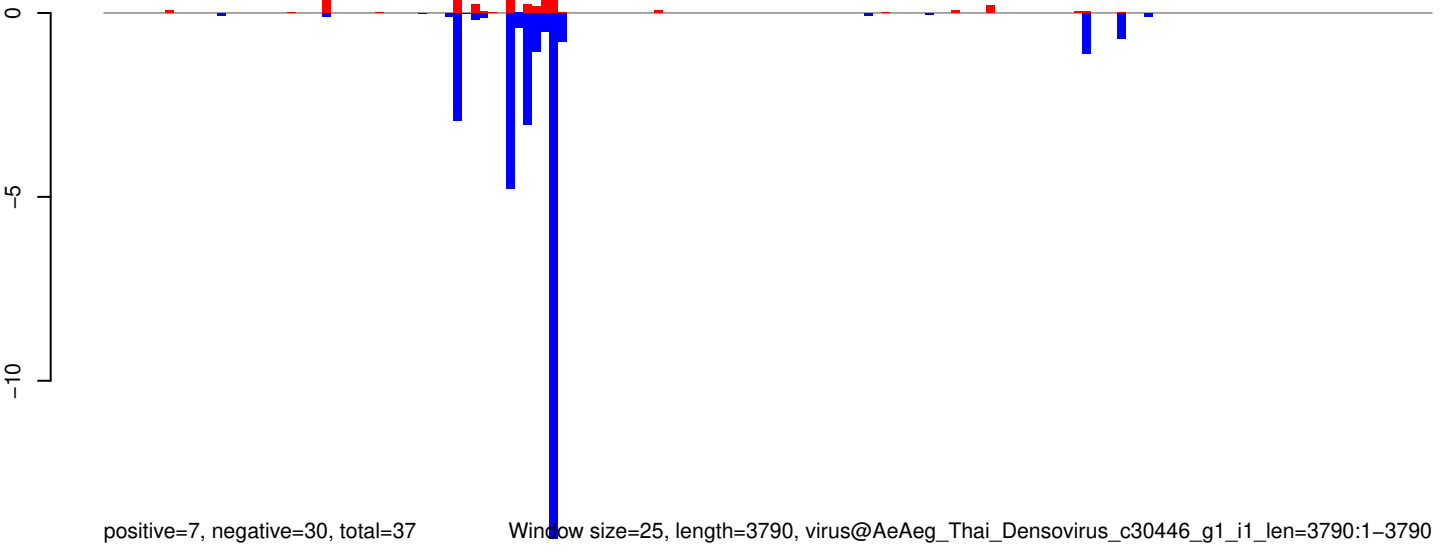
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



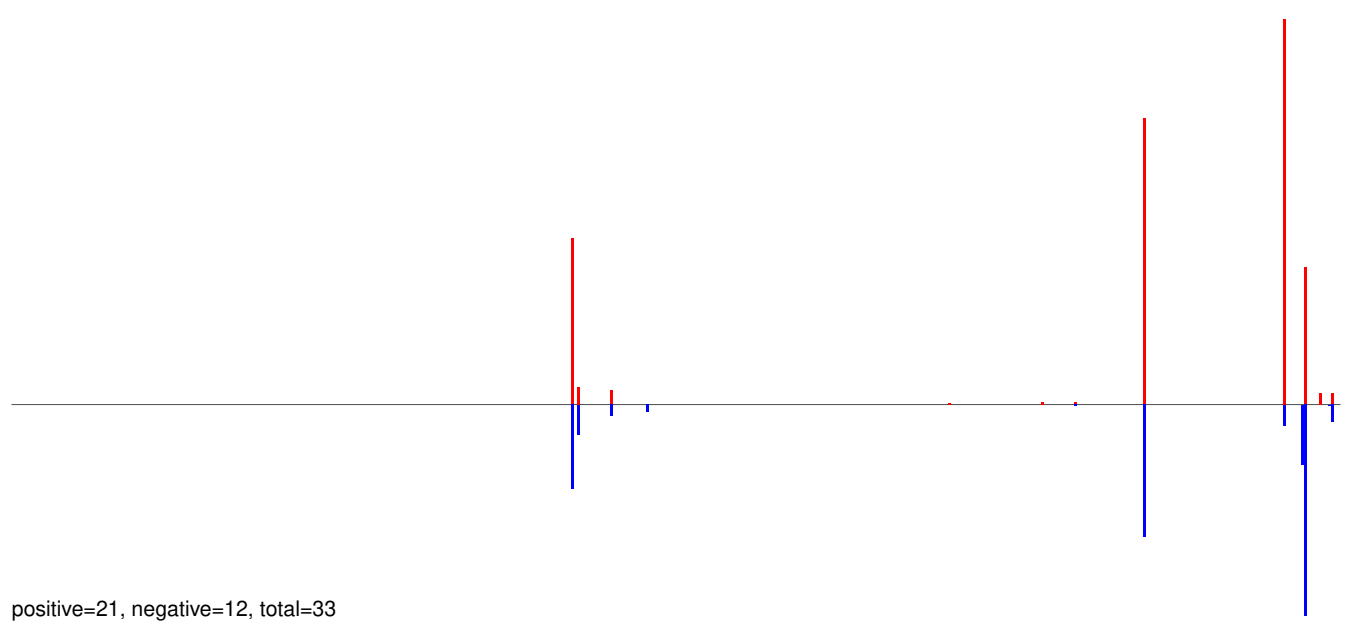
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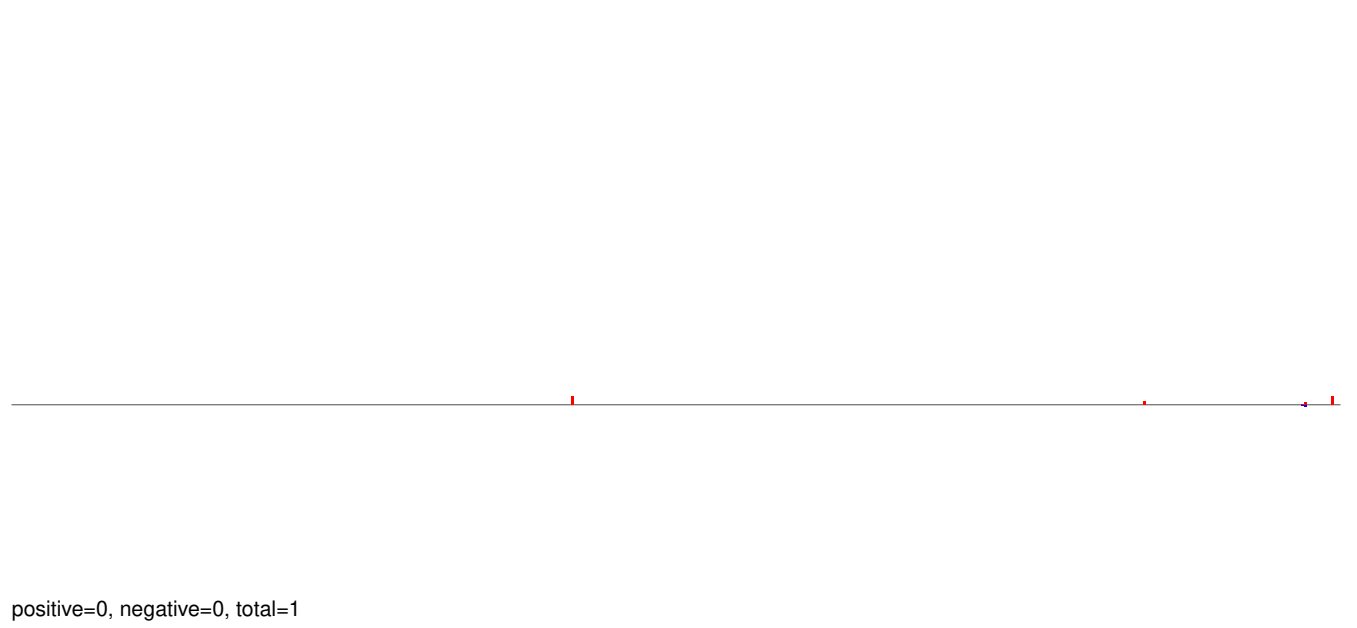
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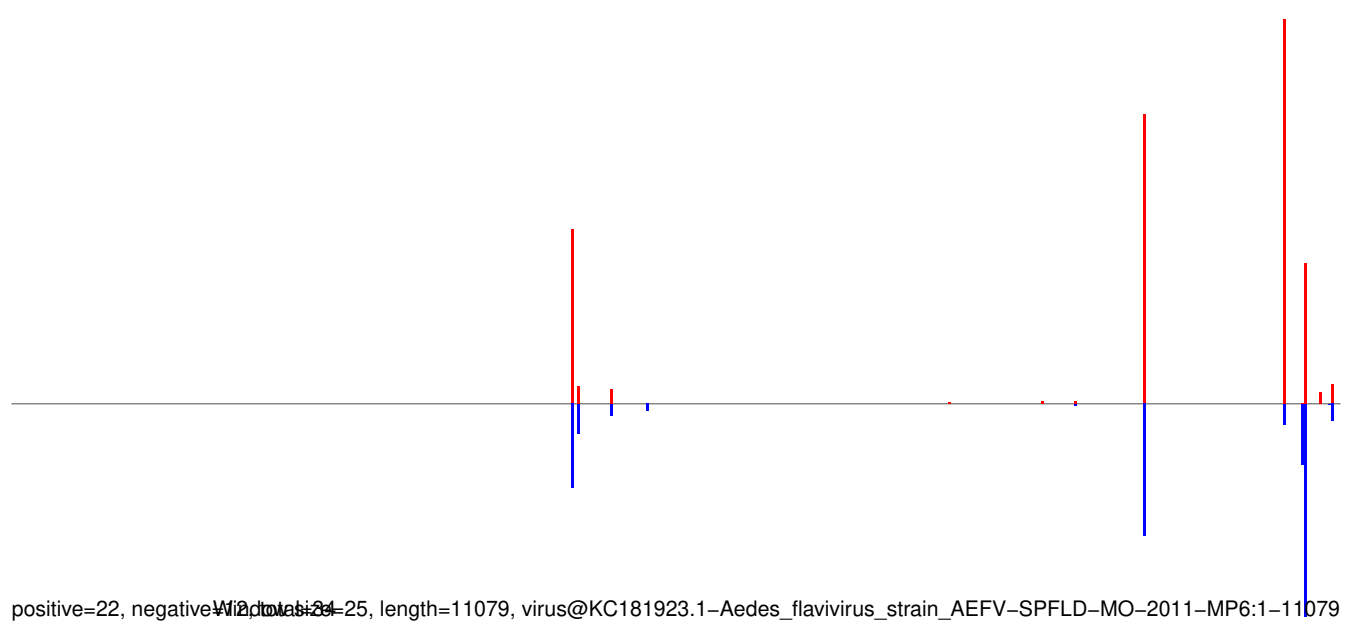
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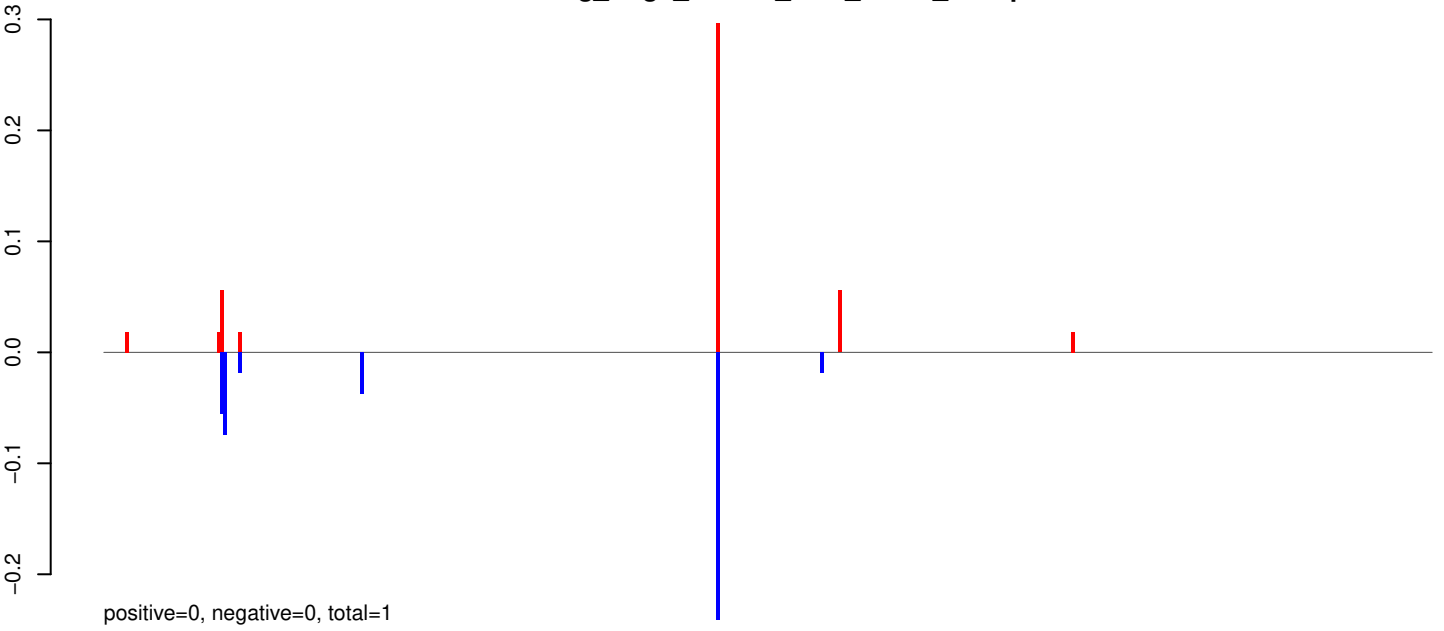


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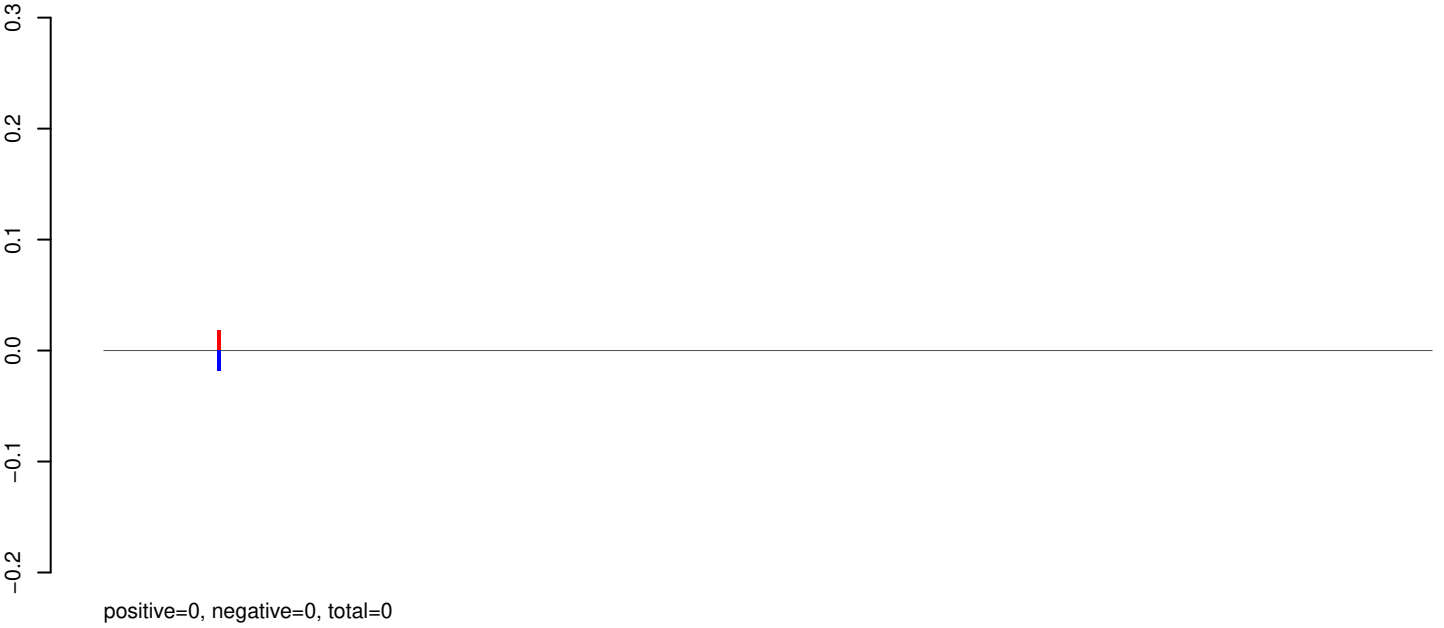


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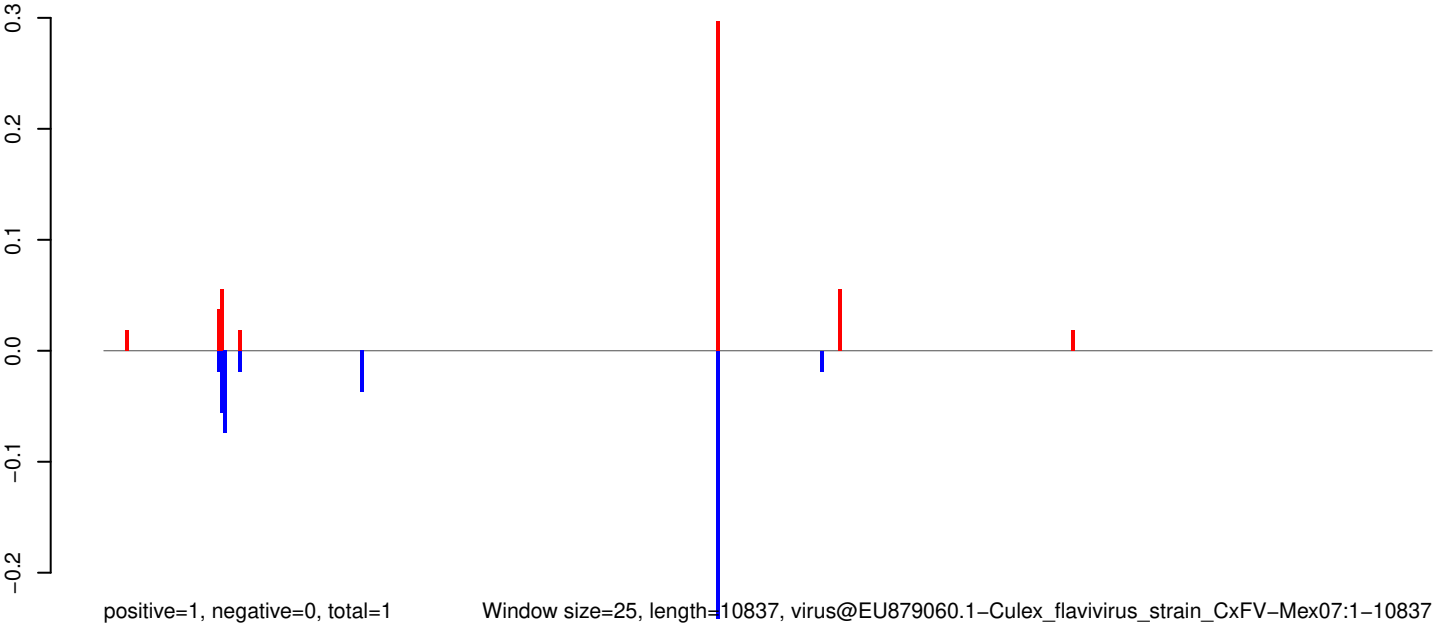
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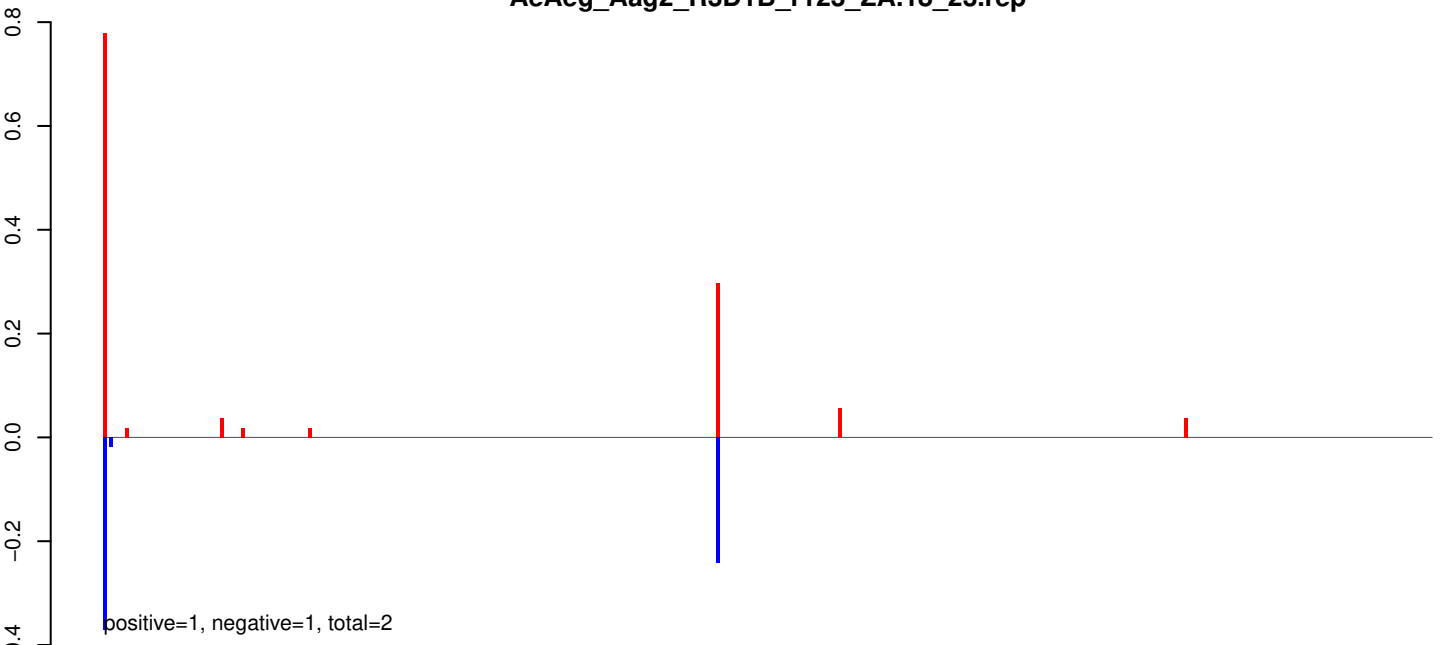


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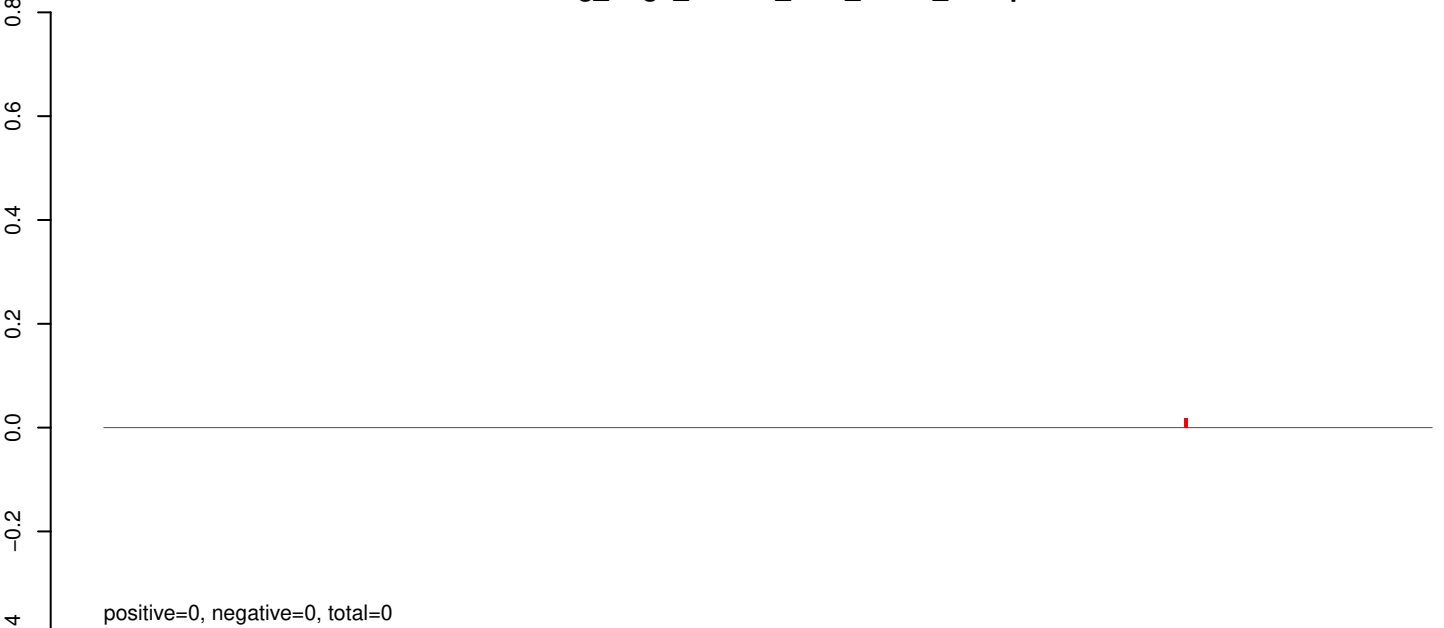


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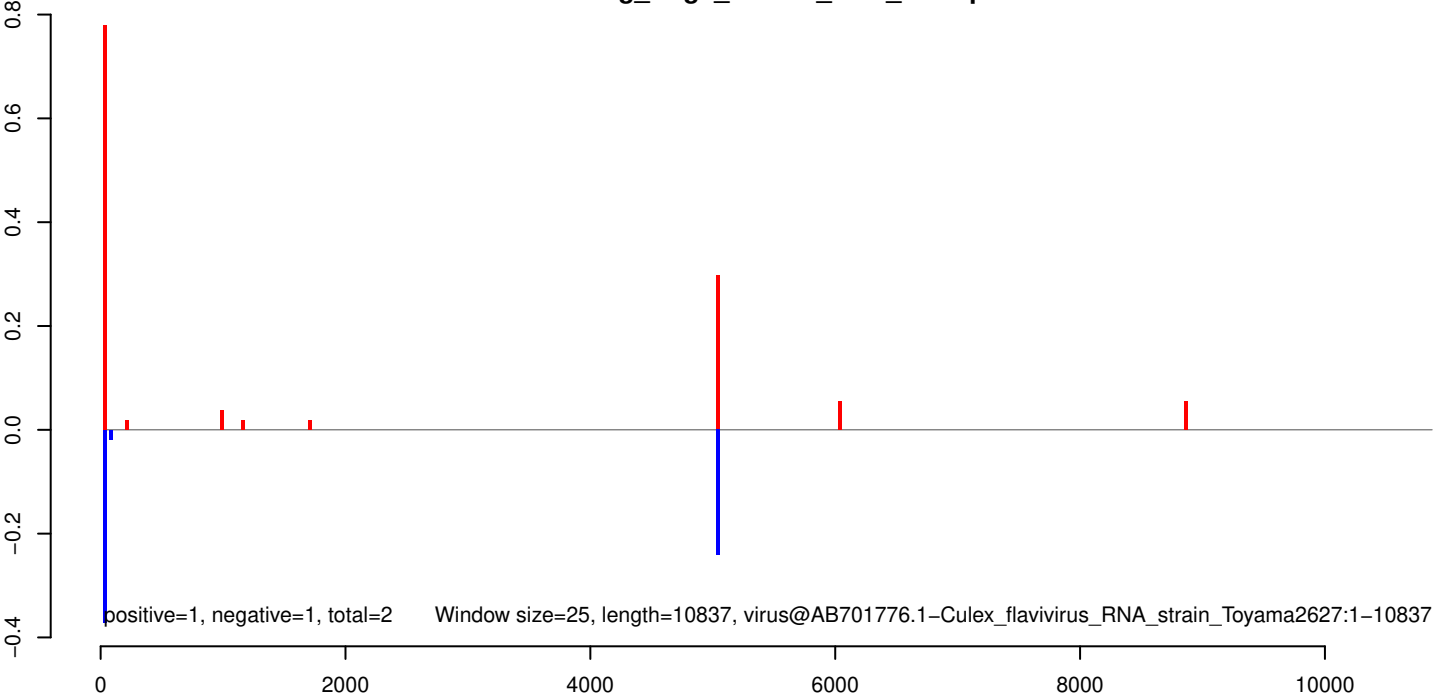
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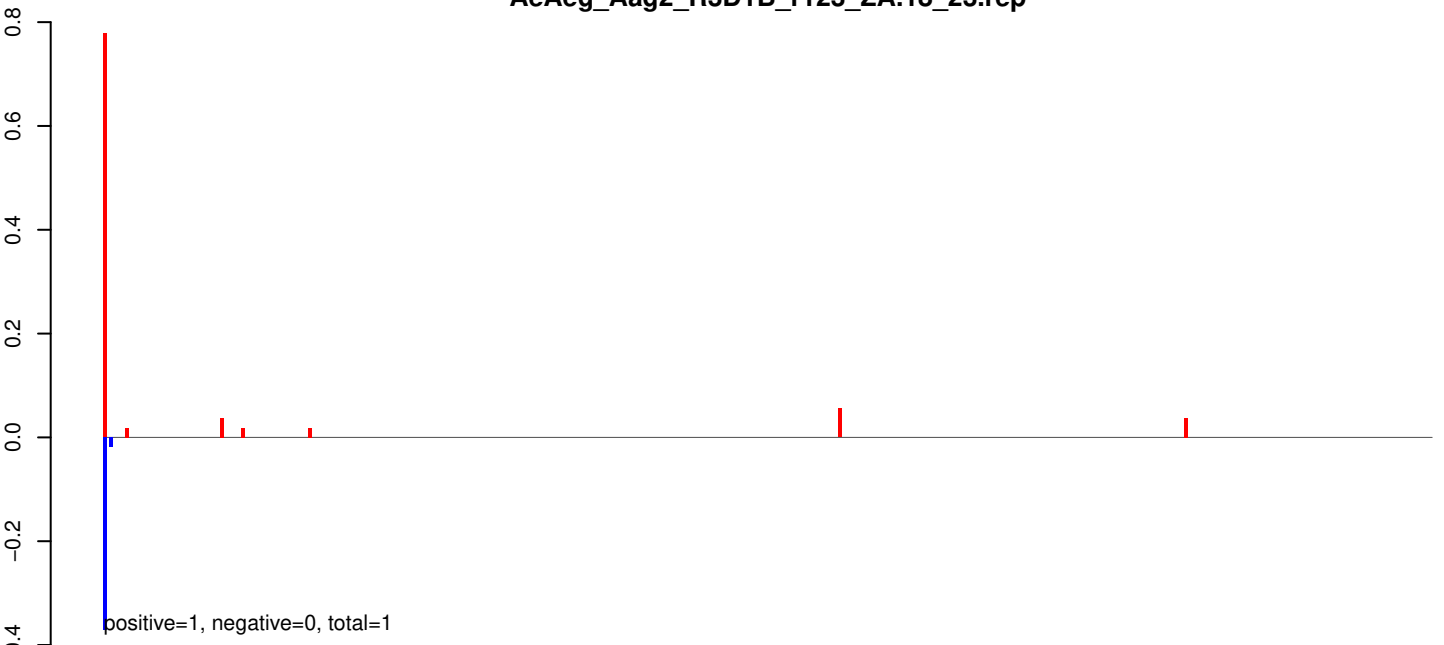
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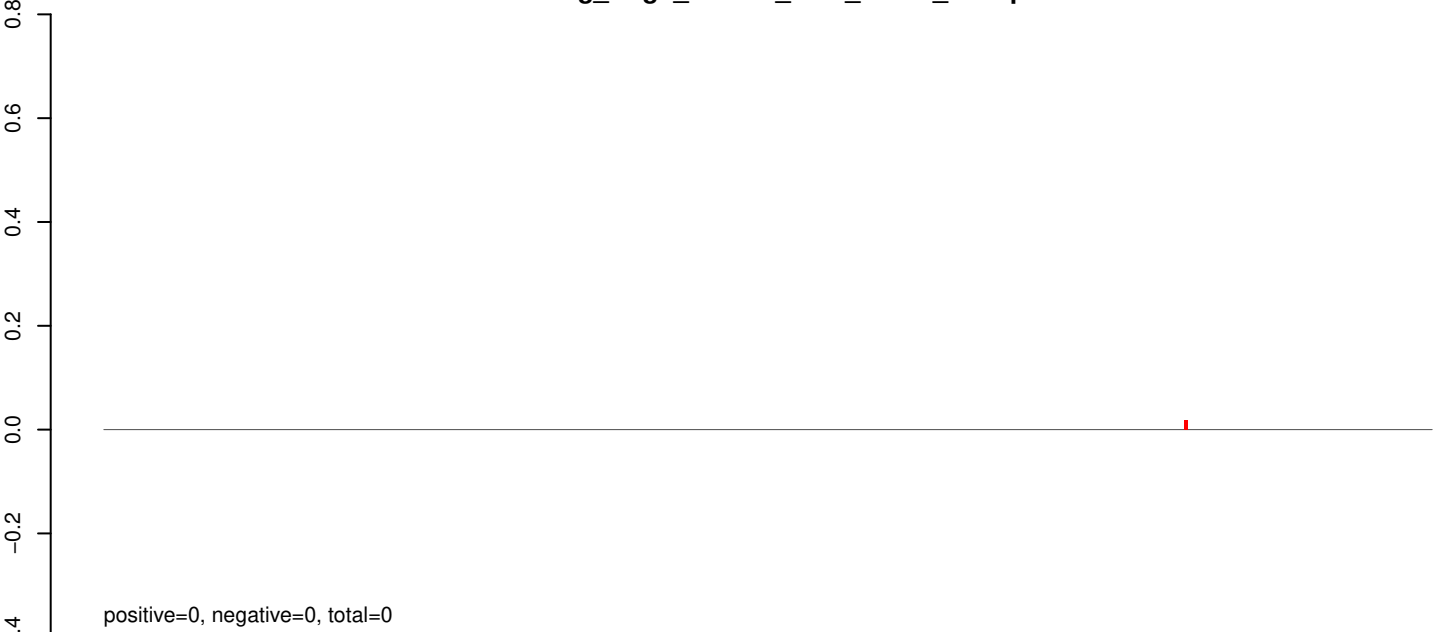
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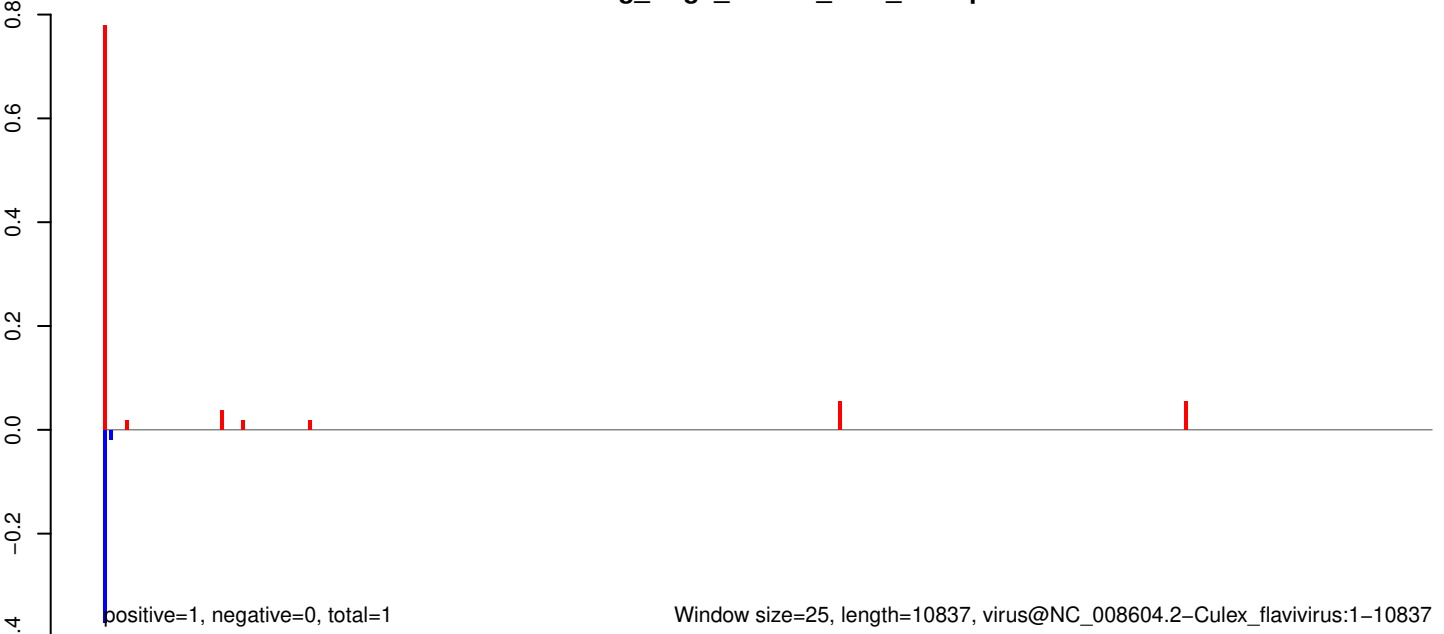
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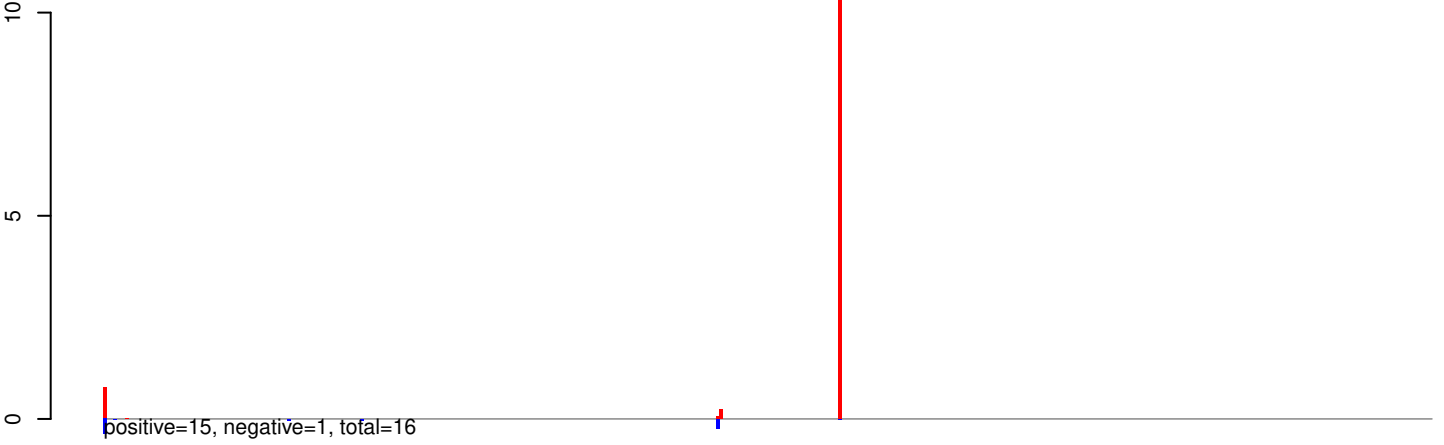
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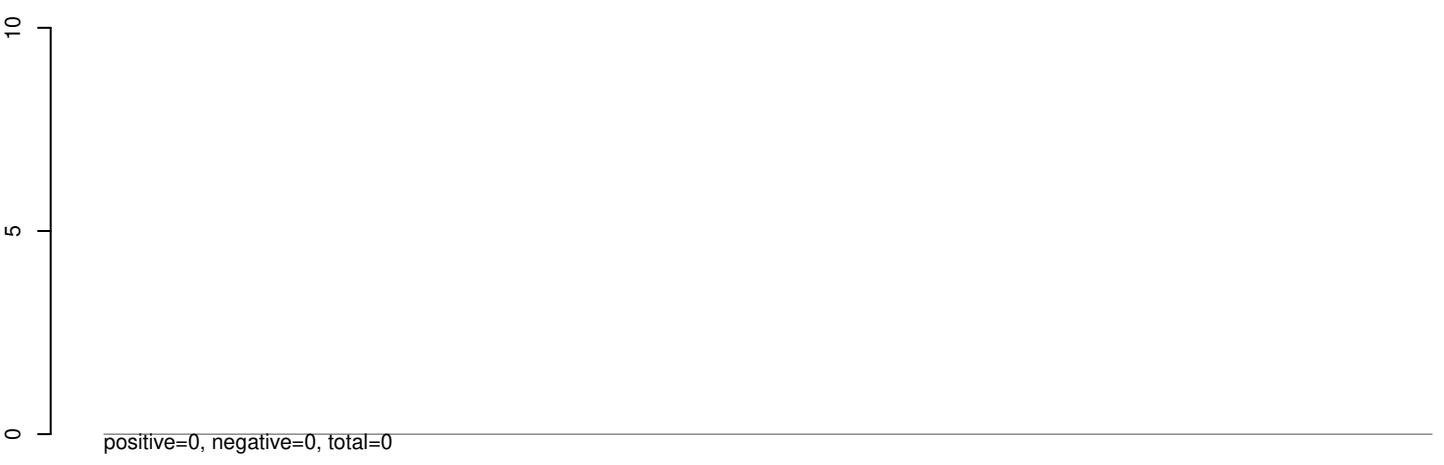
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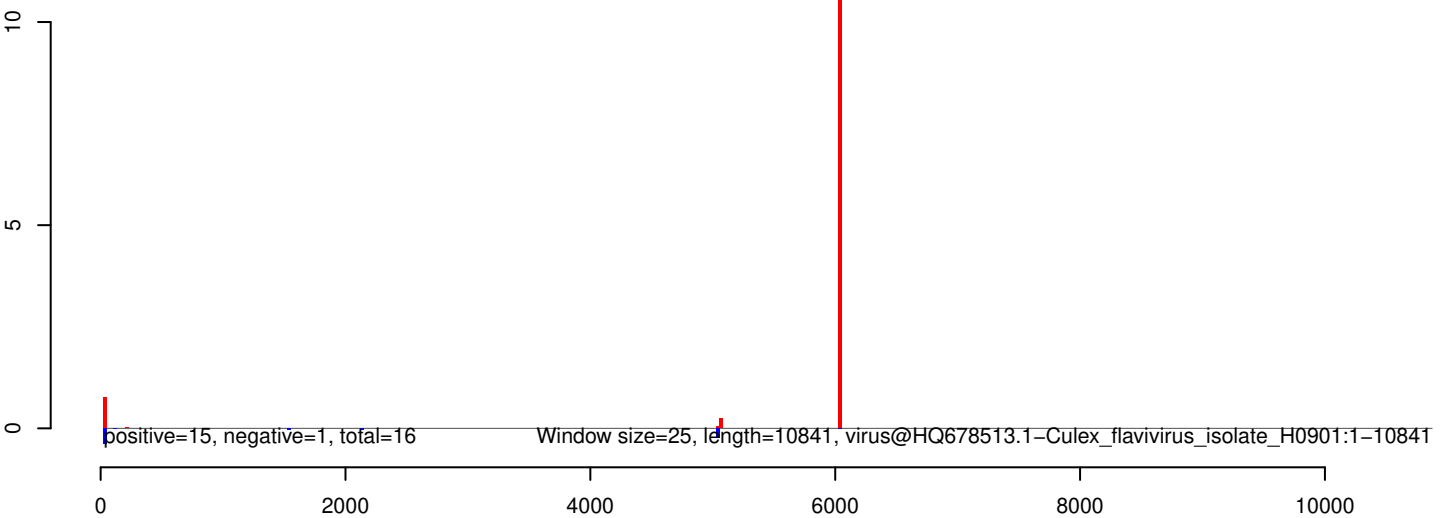
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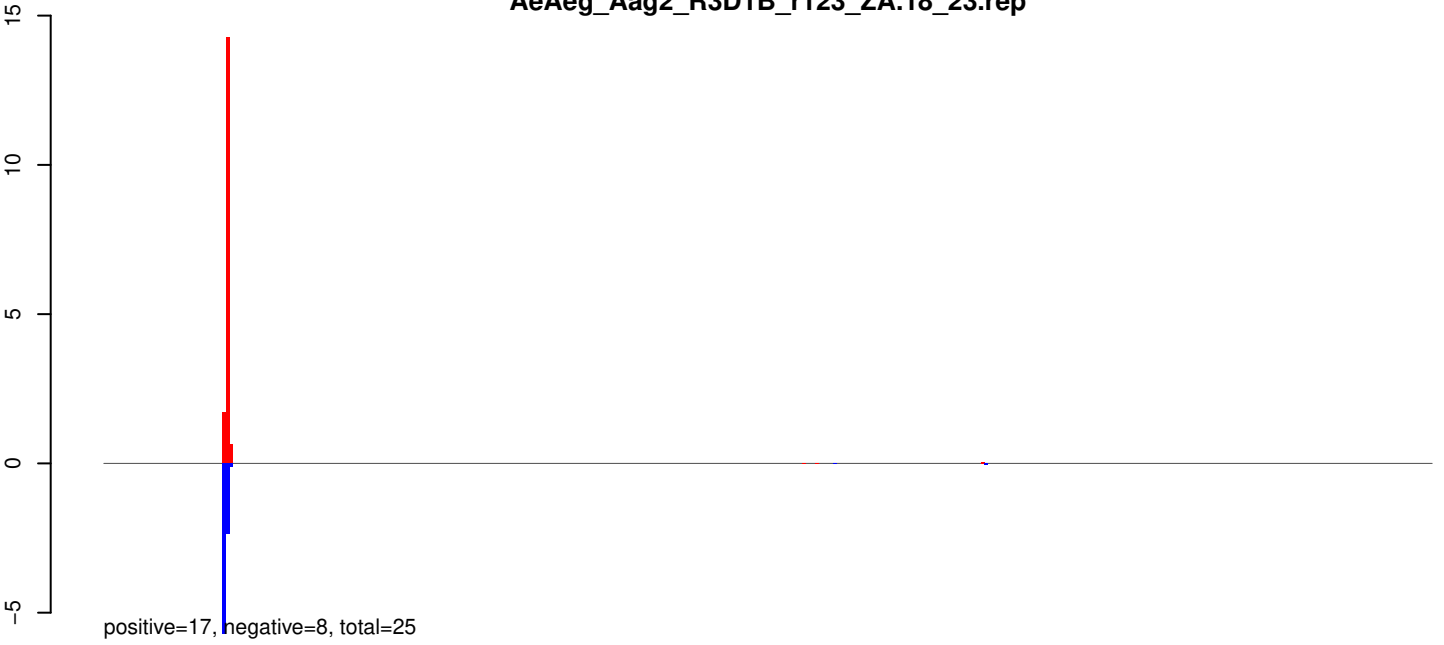
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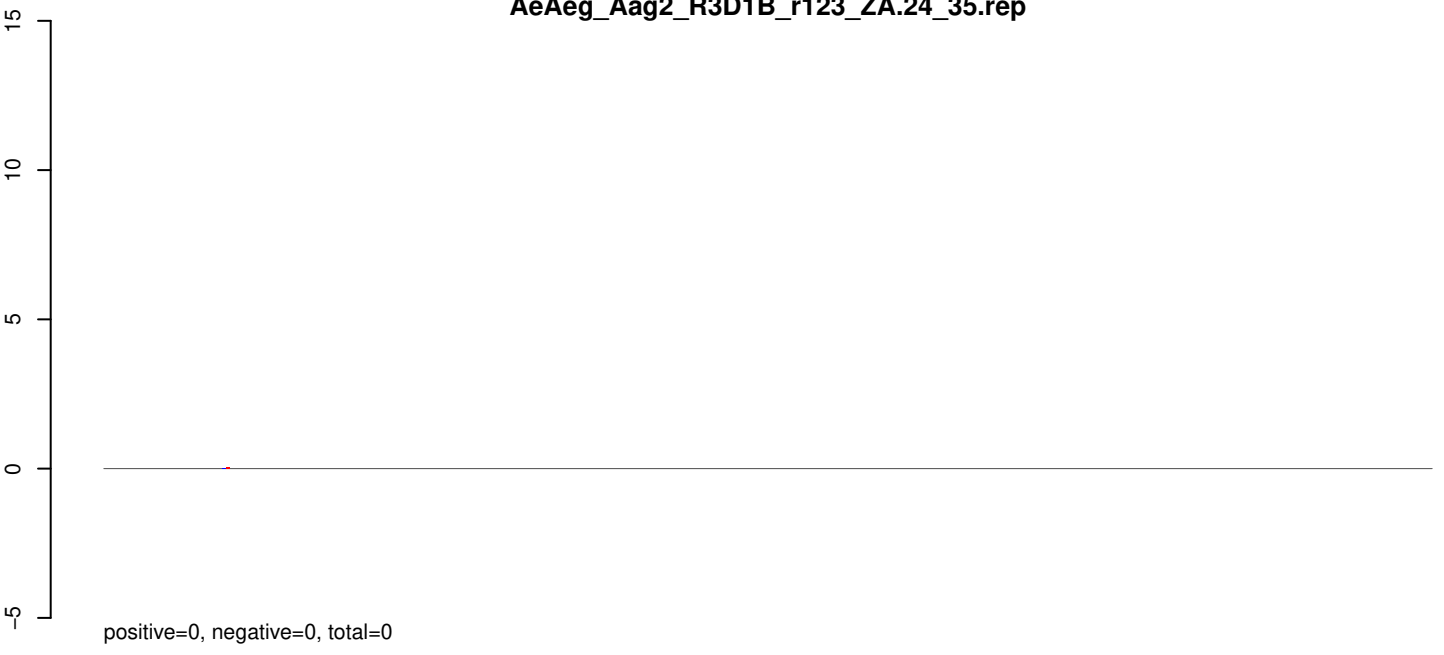
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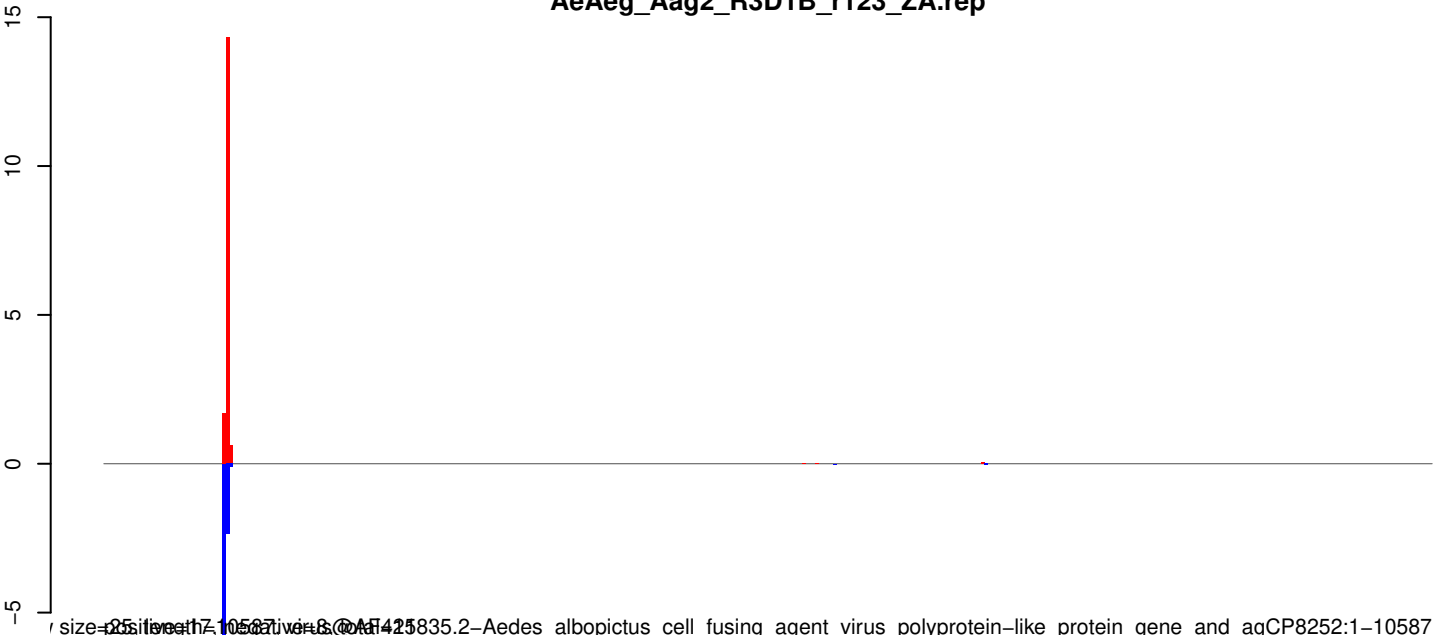
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



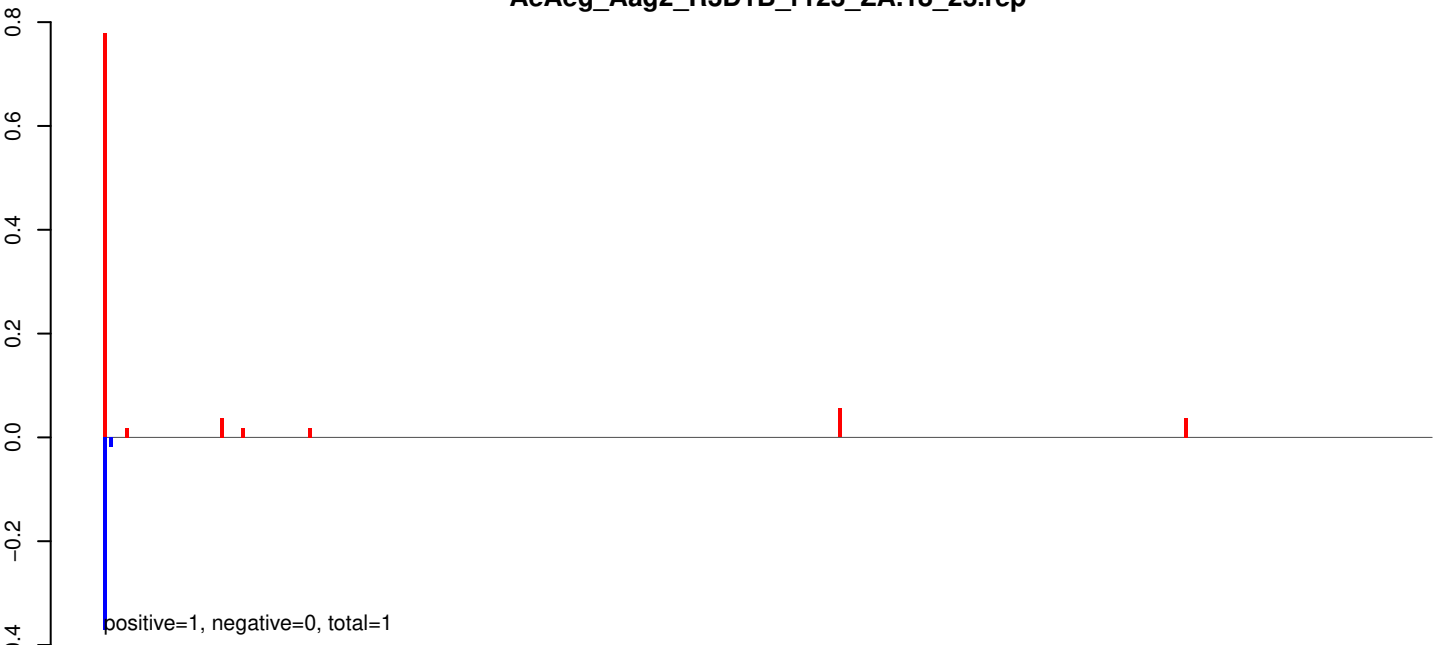
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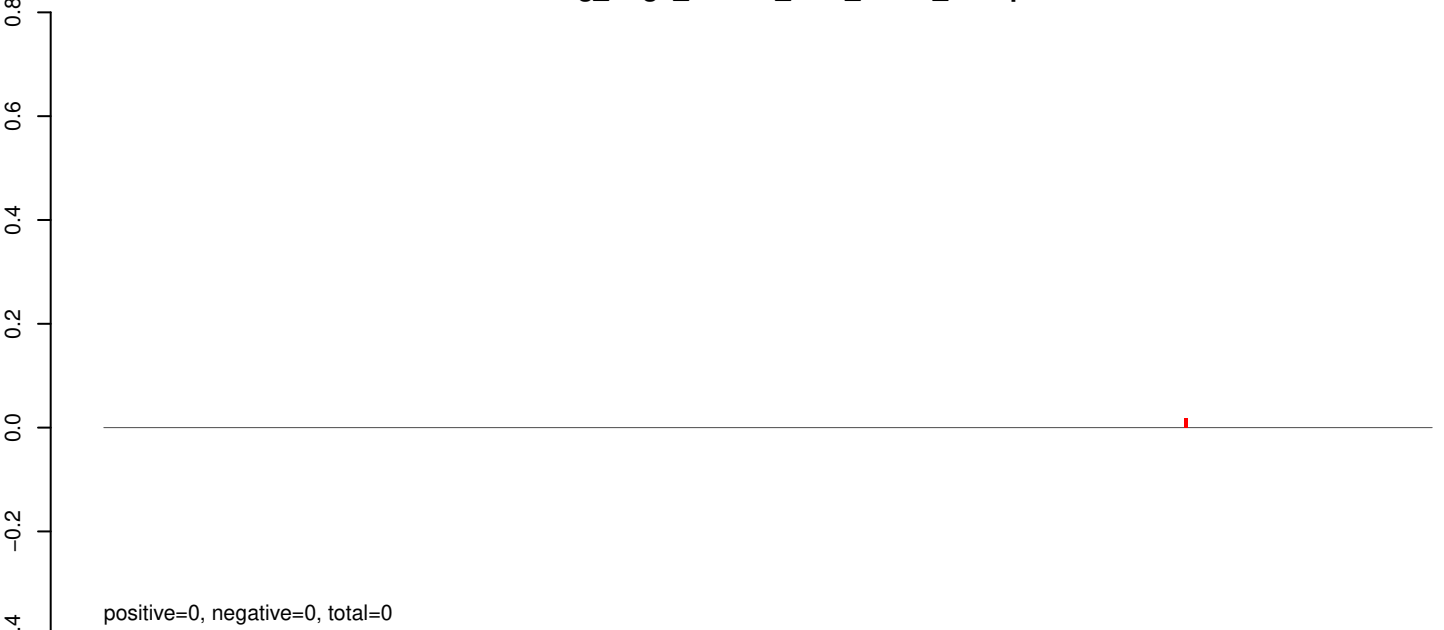
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0 2000 4000 6000 8000 10000

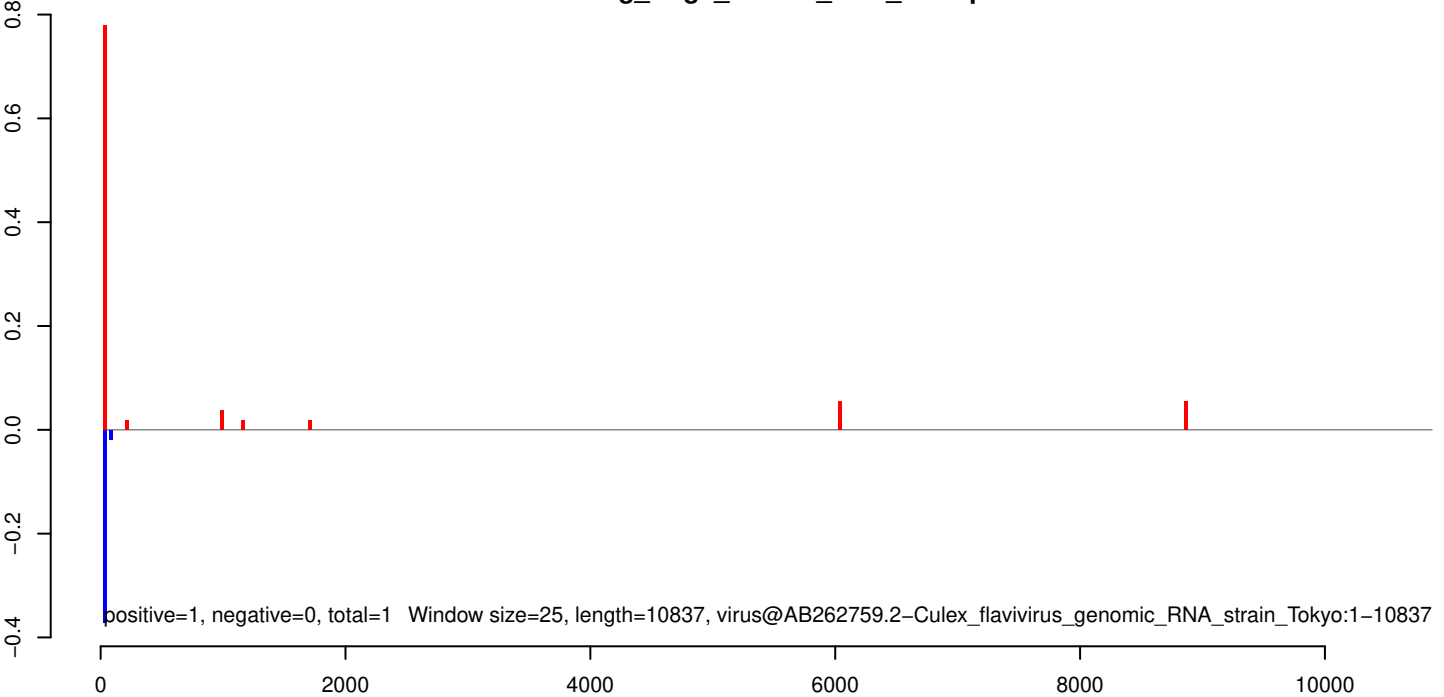
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



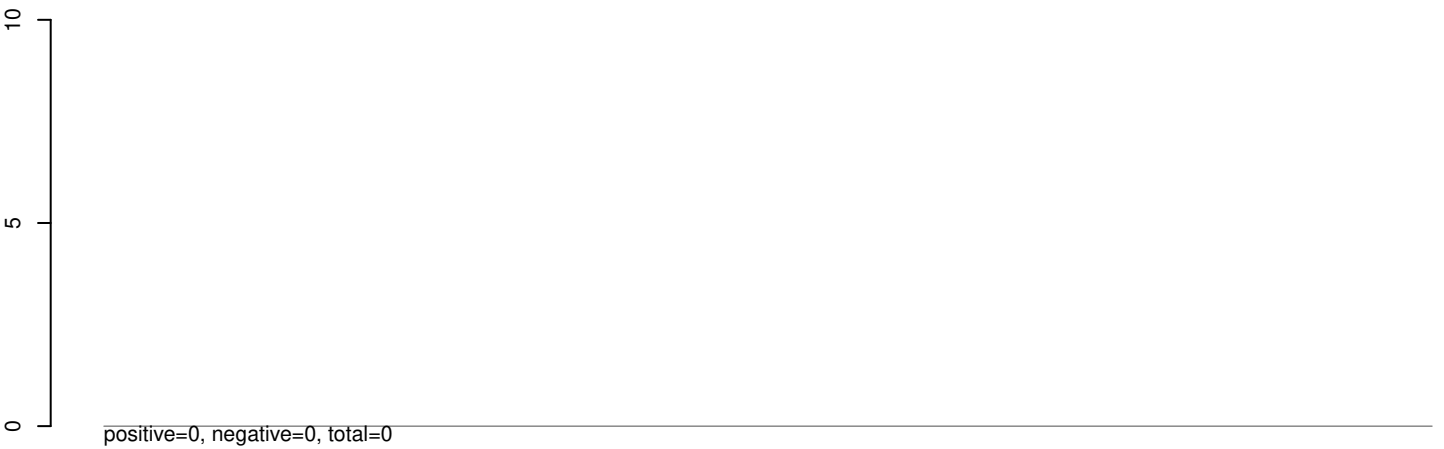
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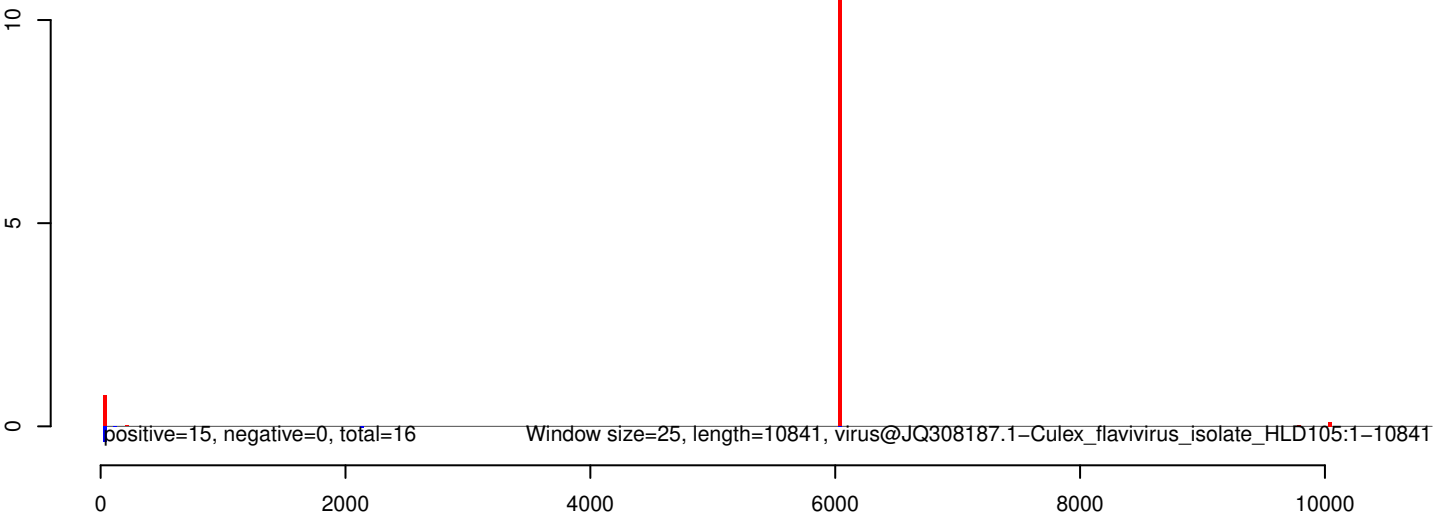
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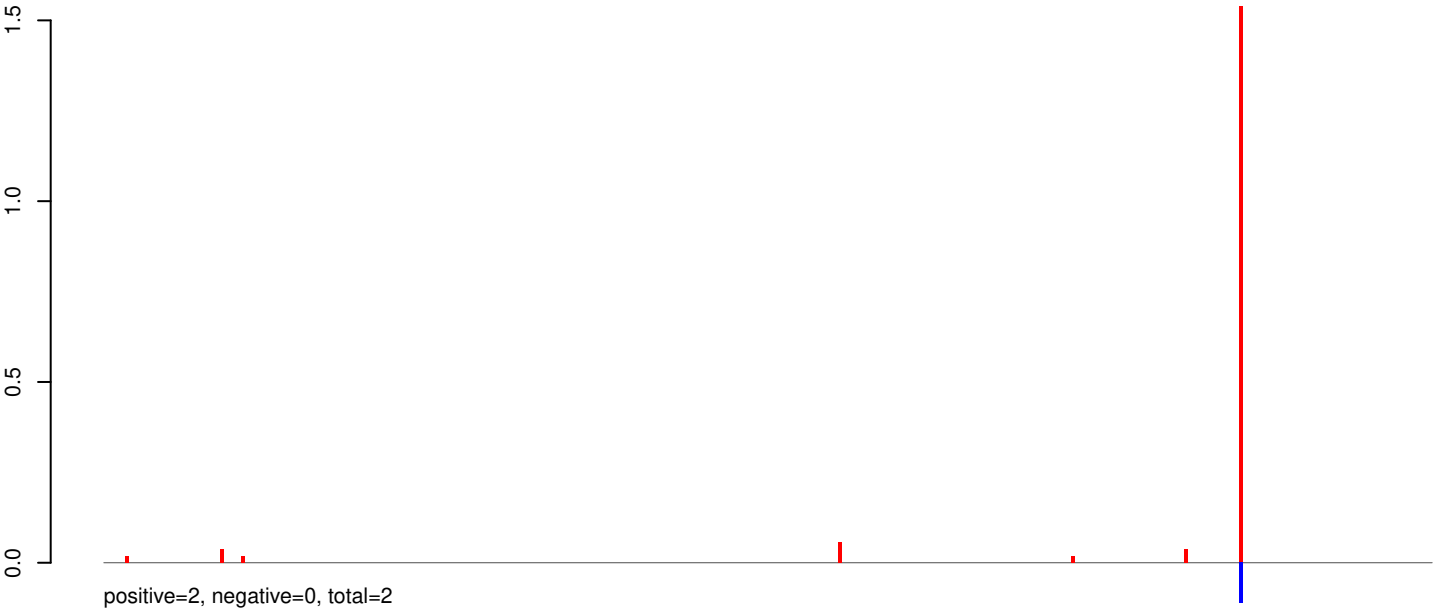
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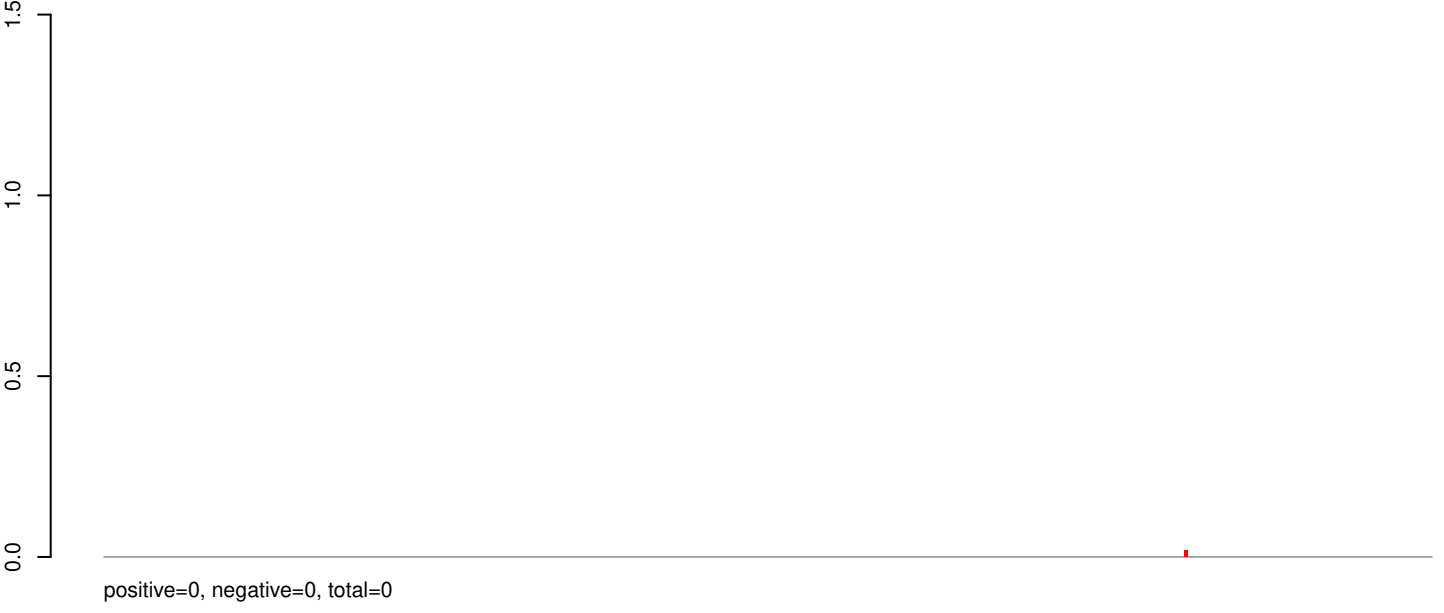
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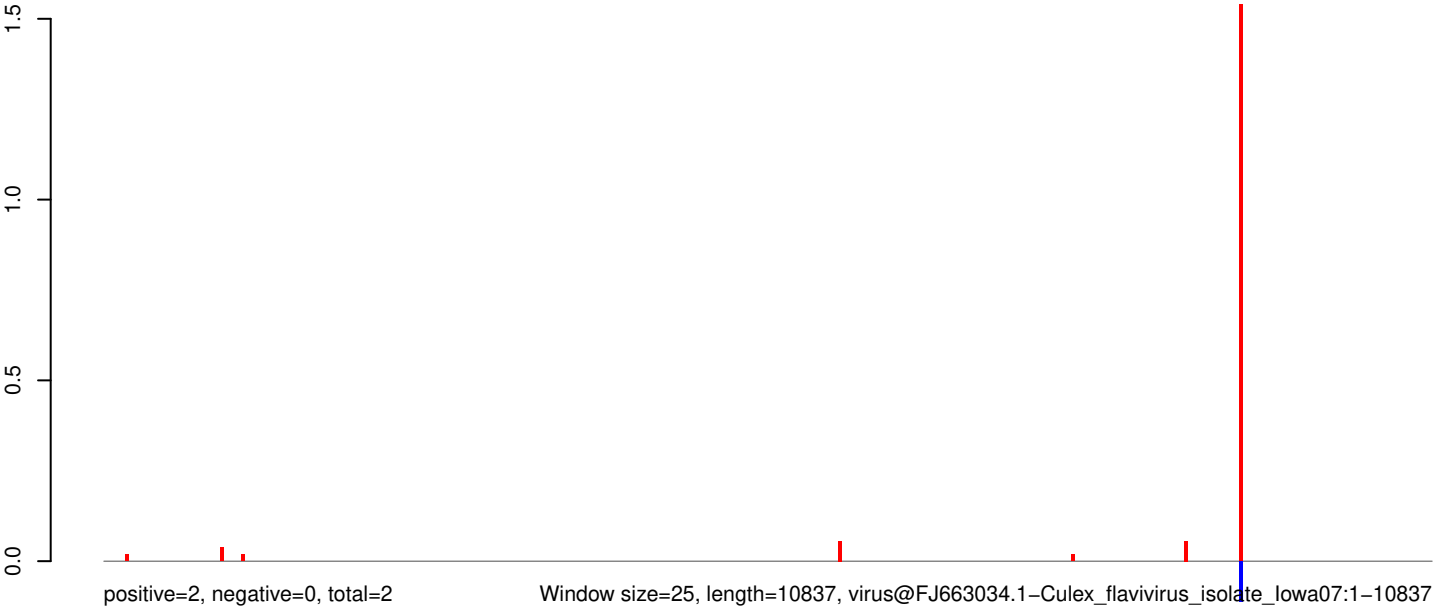
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



AeAeg_Aag2_R3D1B_r123_ZA.rep



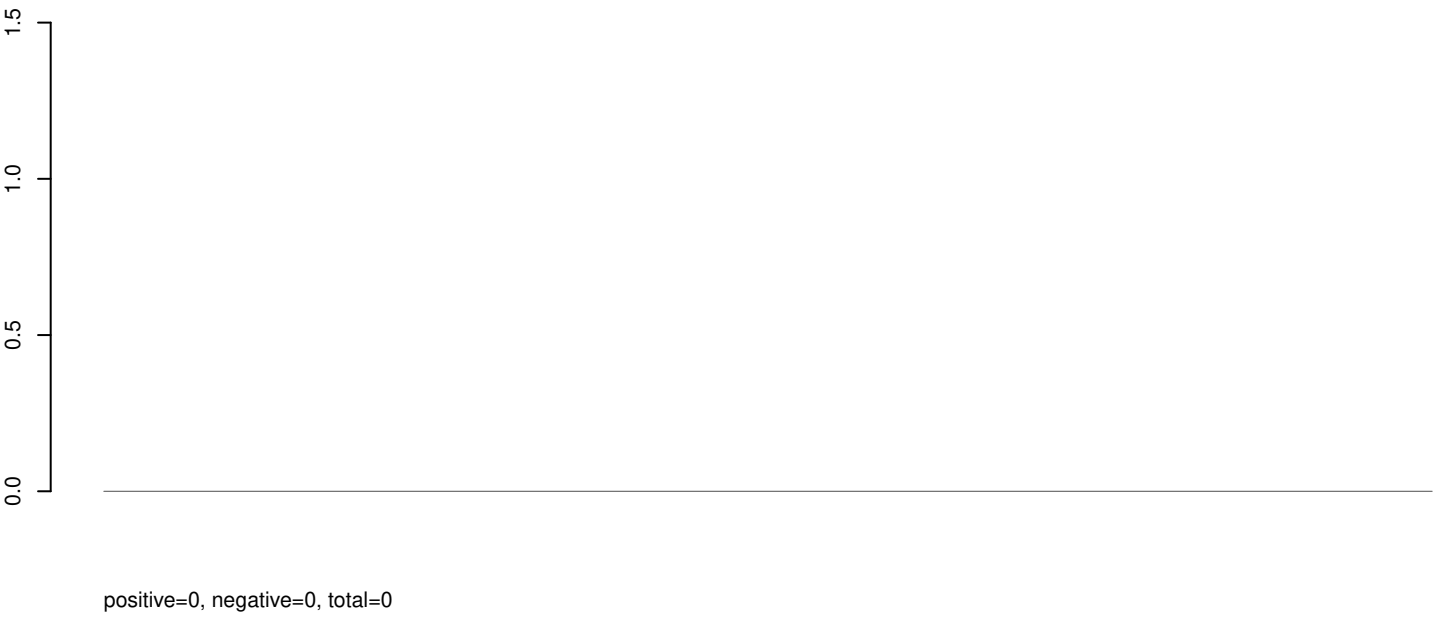
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0 2000 4000 6000 8000 10000

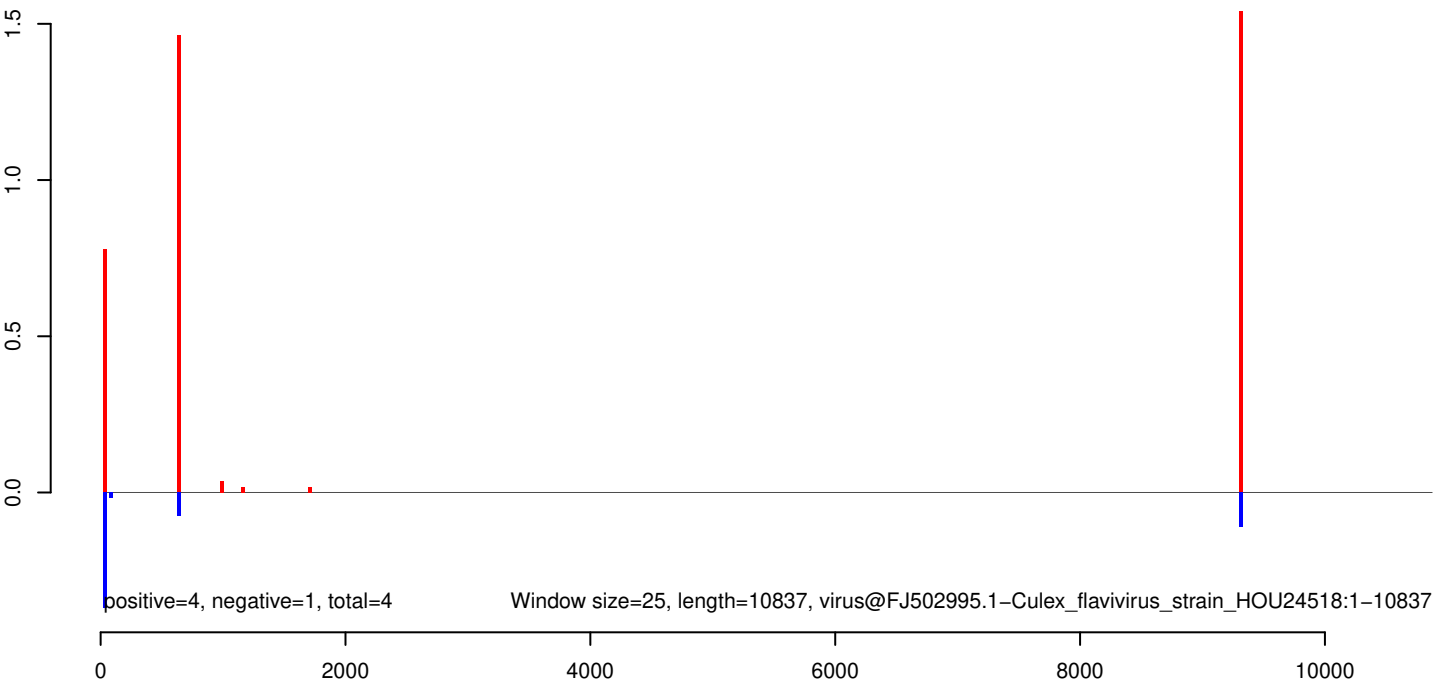
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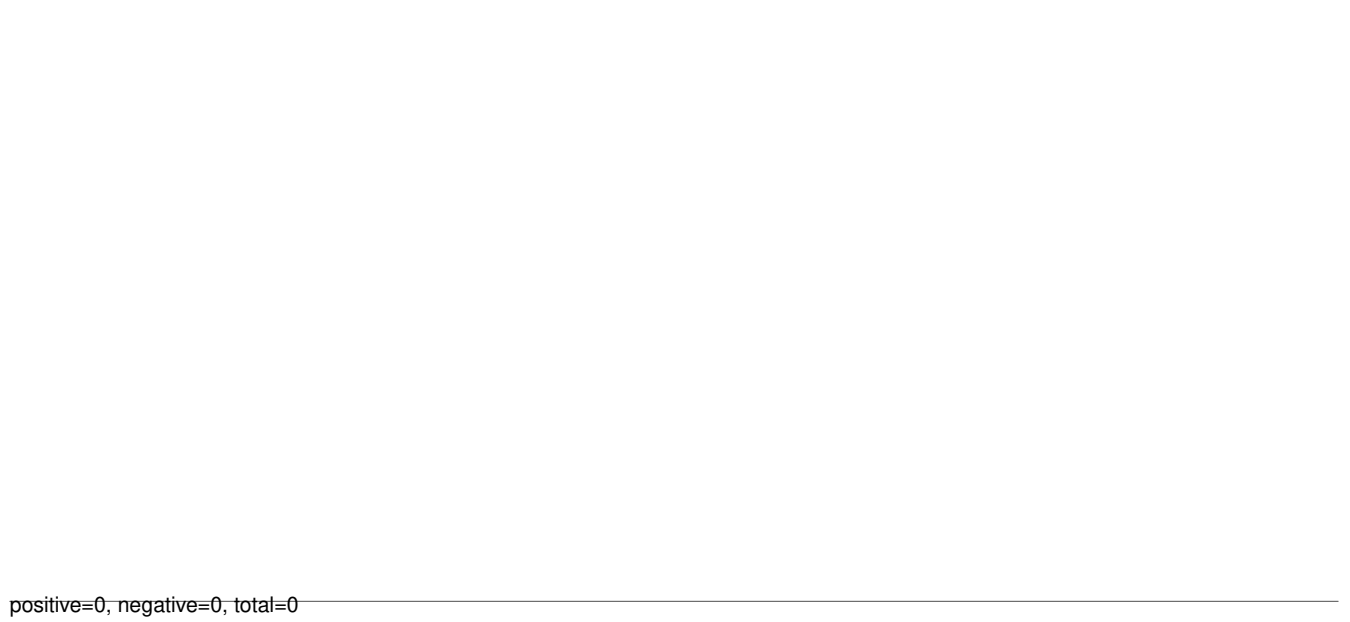
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AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



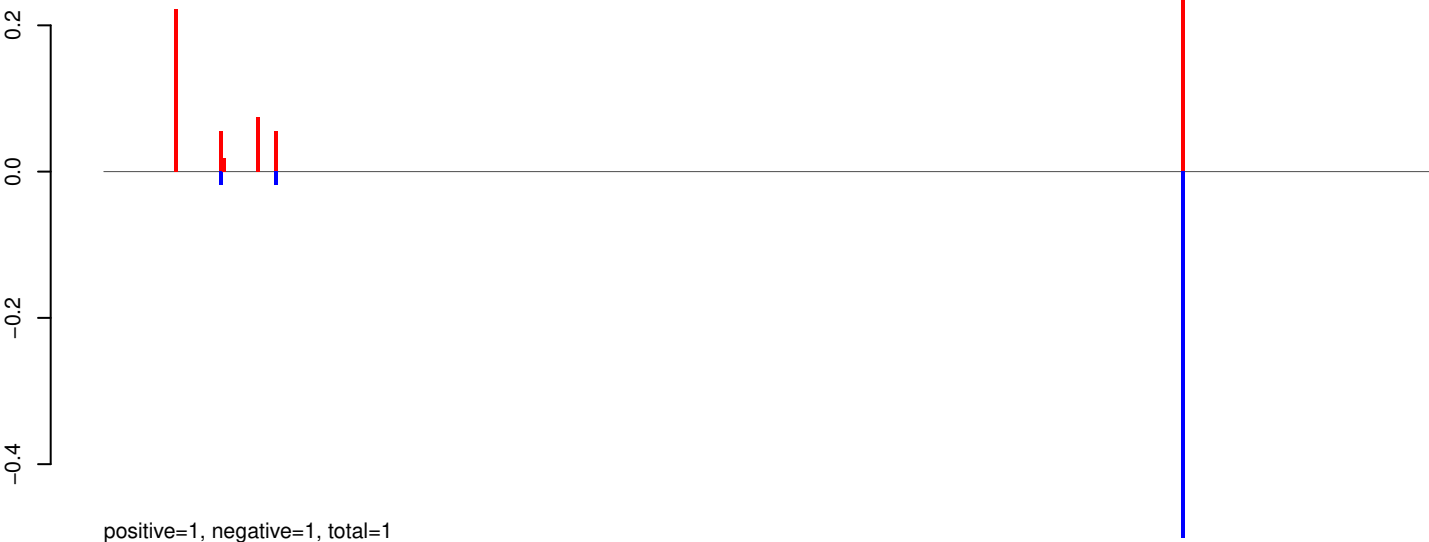
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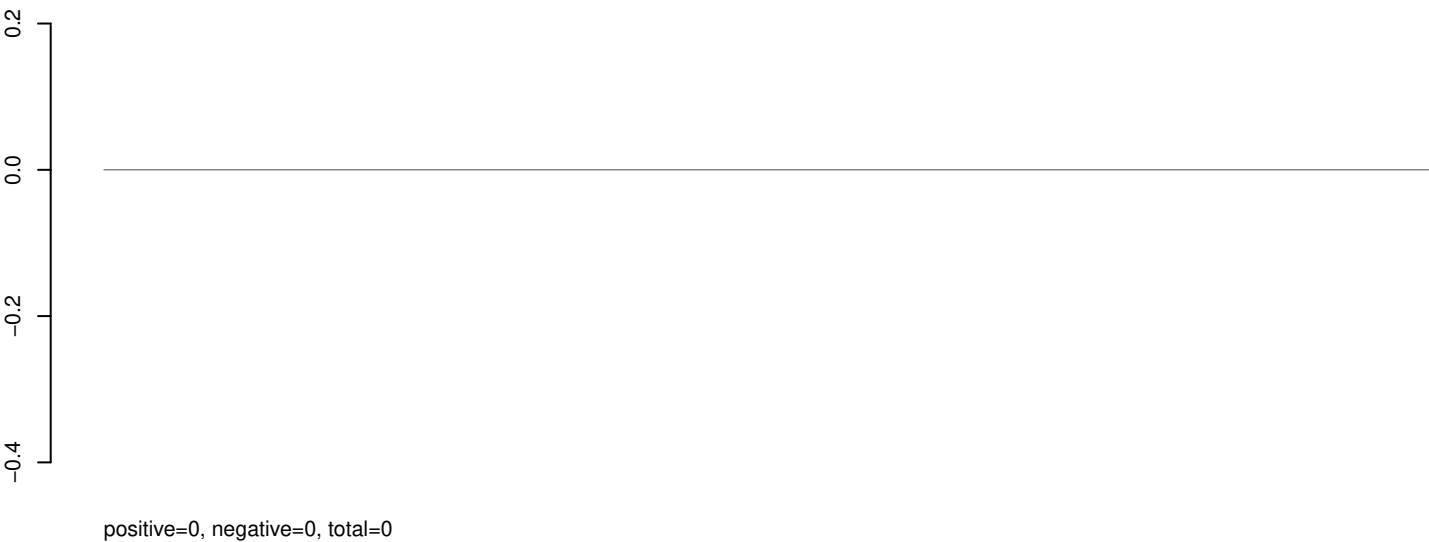
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0 2000 4000 6000 8000 10000

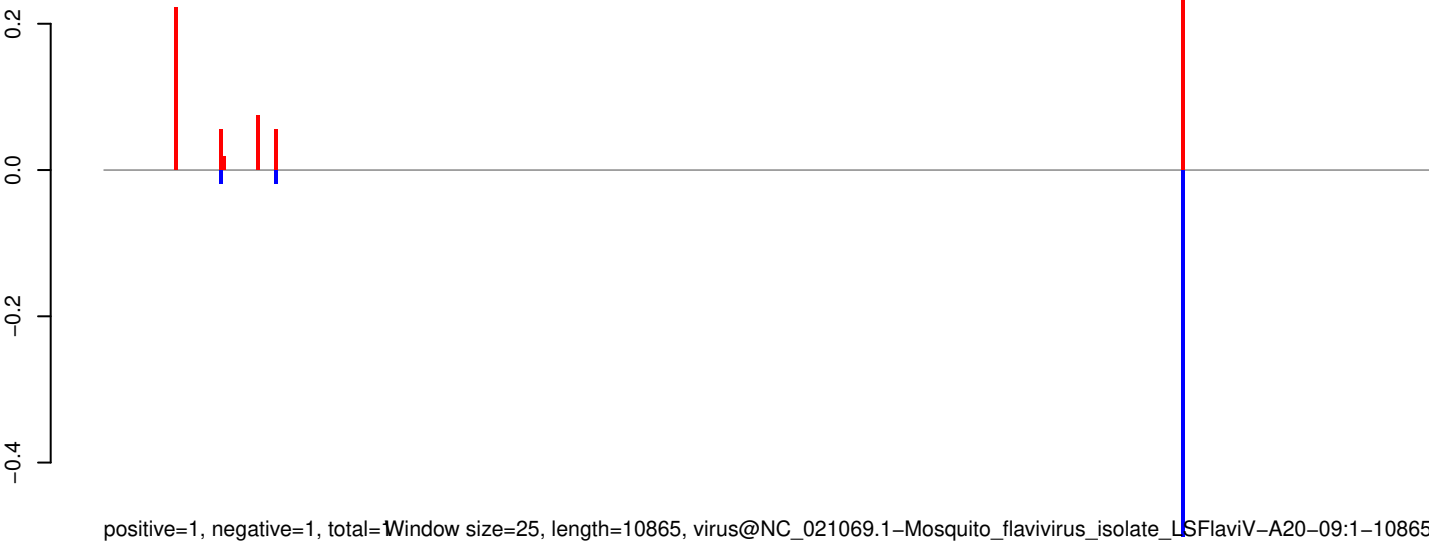
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep

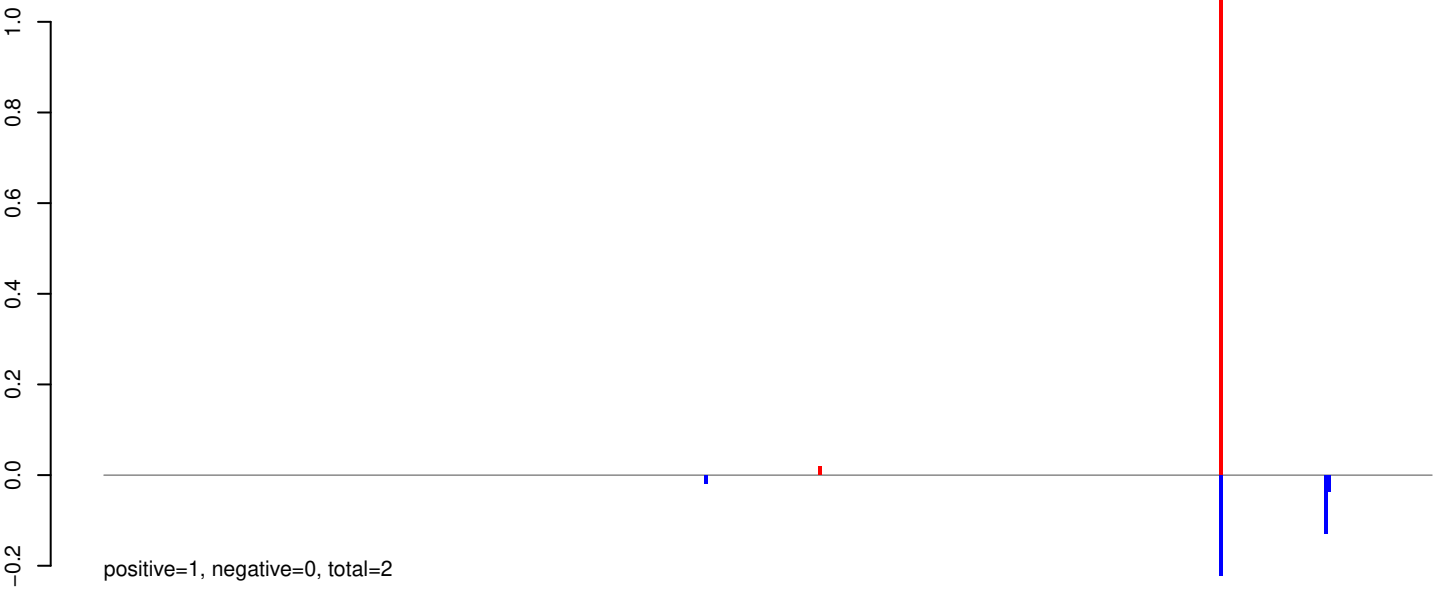


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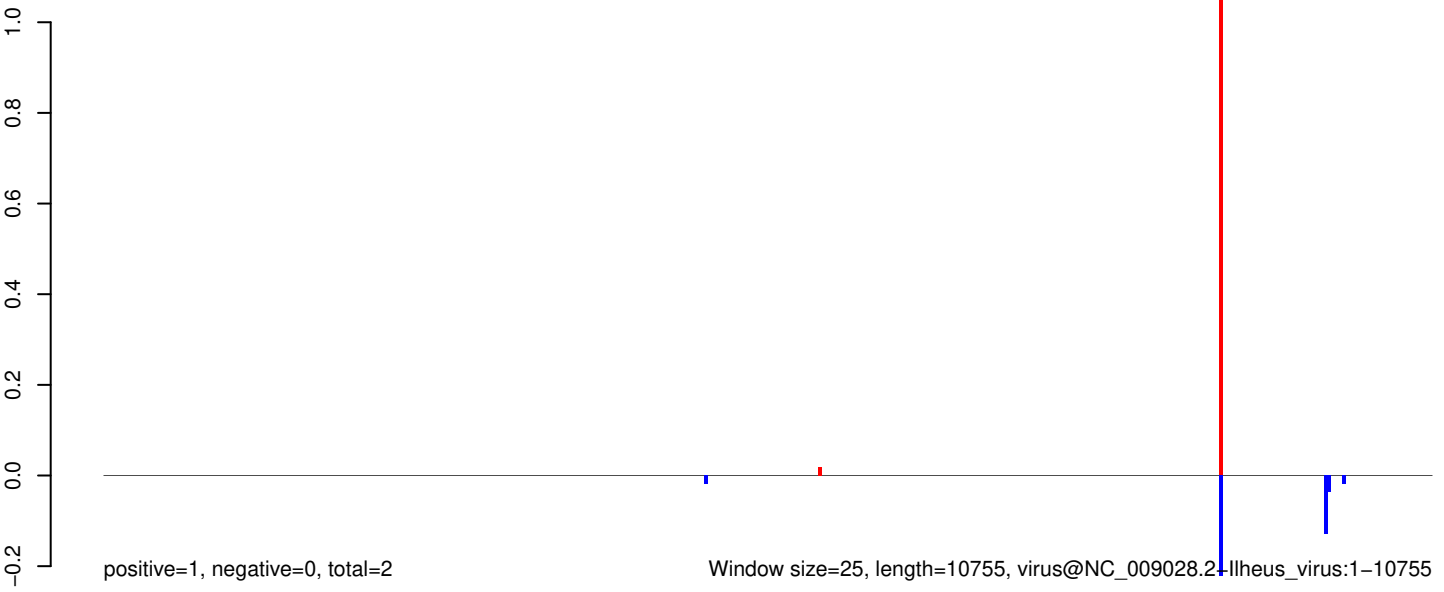
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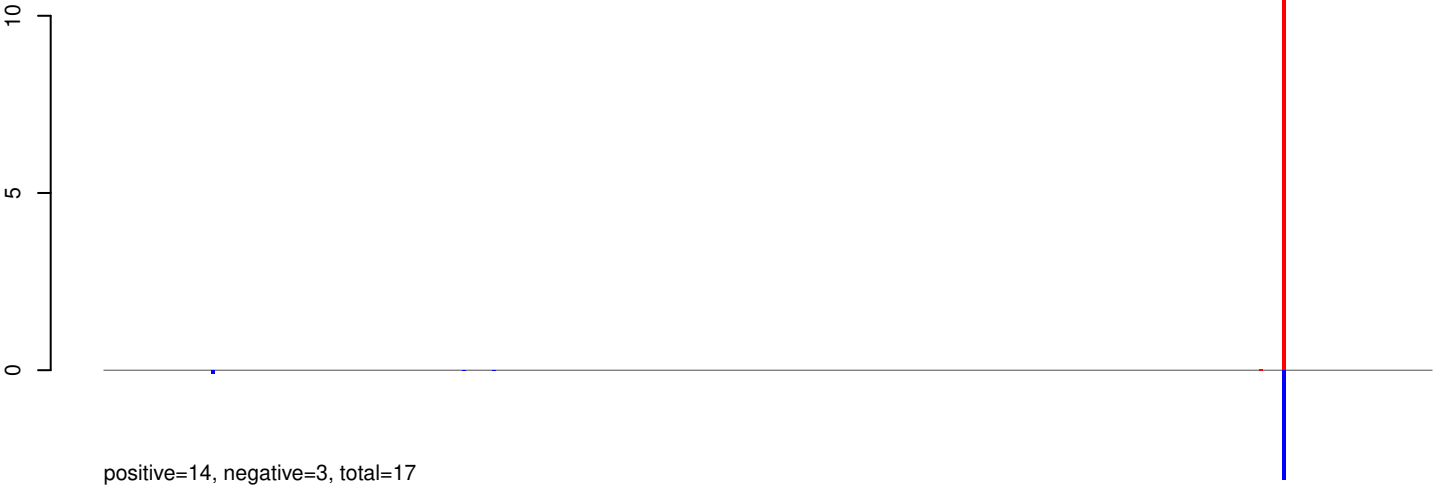
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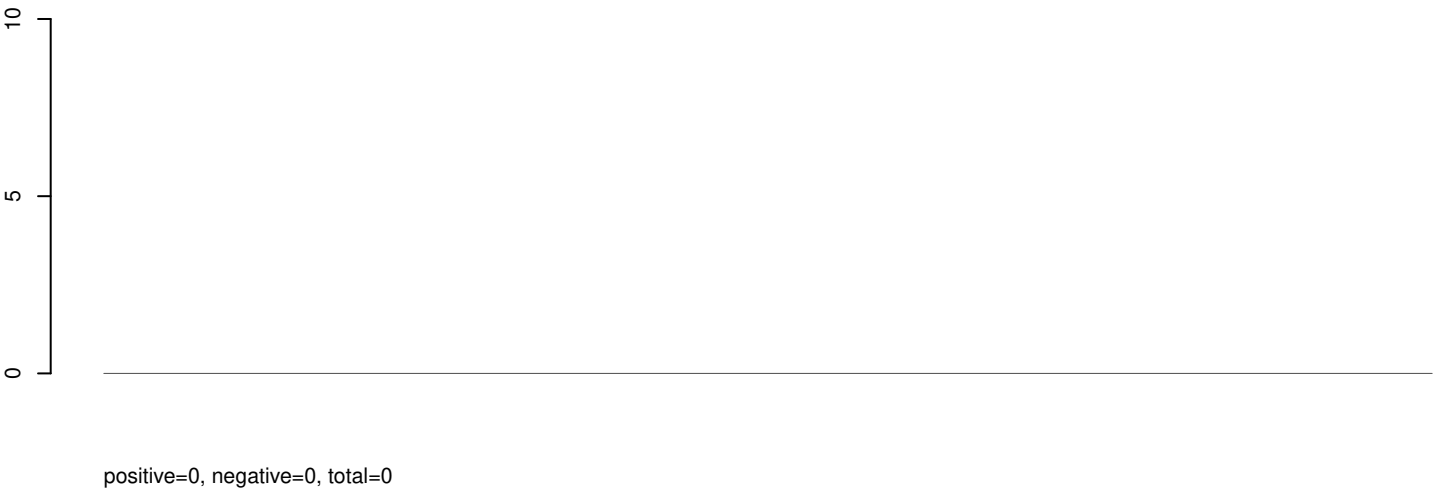
Window size=25, length=10755, virus@NC_009028.2-Ilheus_virus:1-10755

0 2000 4000 6000 8000 10000

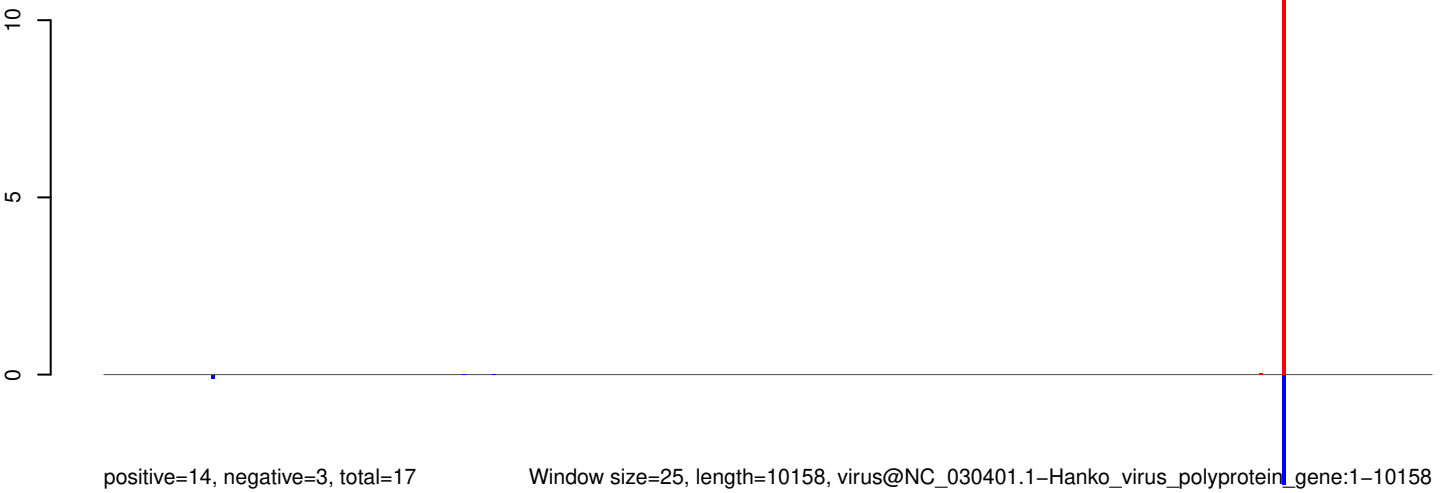
AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep

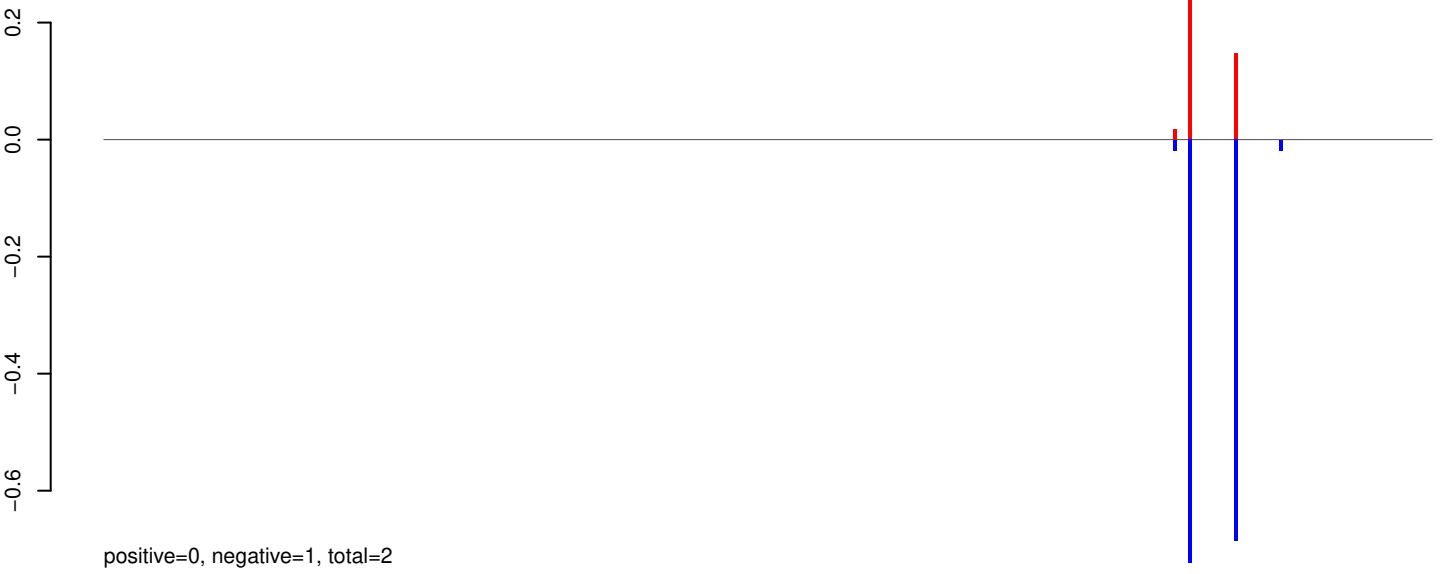


AeAeg_Aag2_R3D1B_r123_ZA.rep

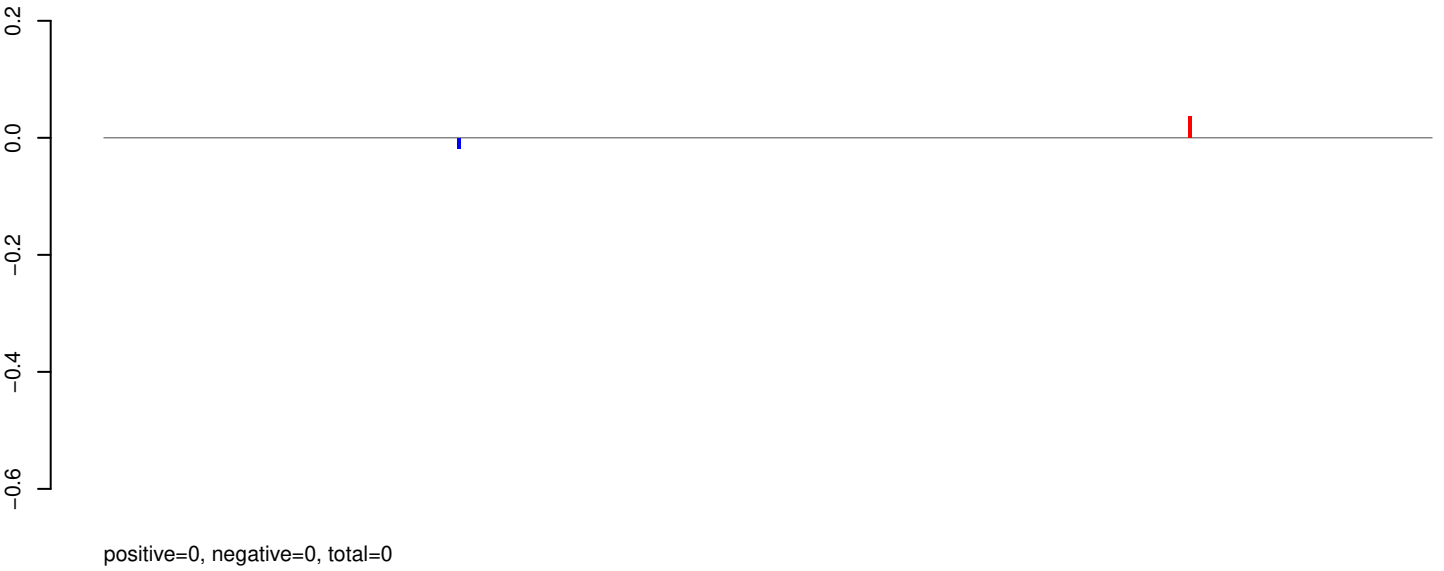


0 2000 4000 6000 8000 10000

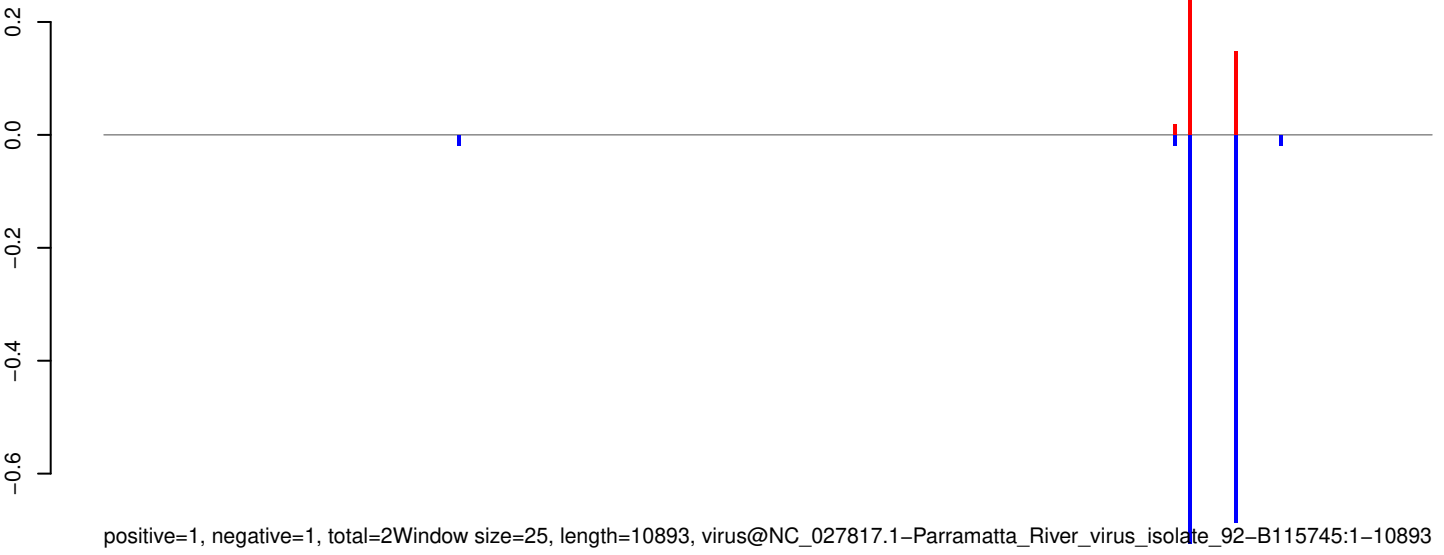
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



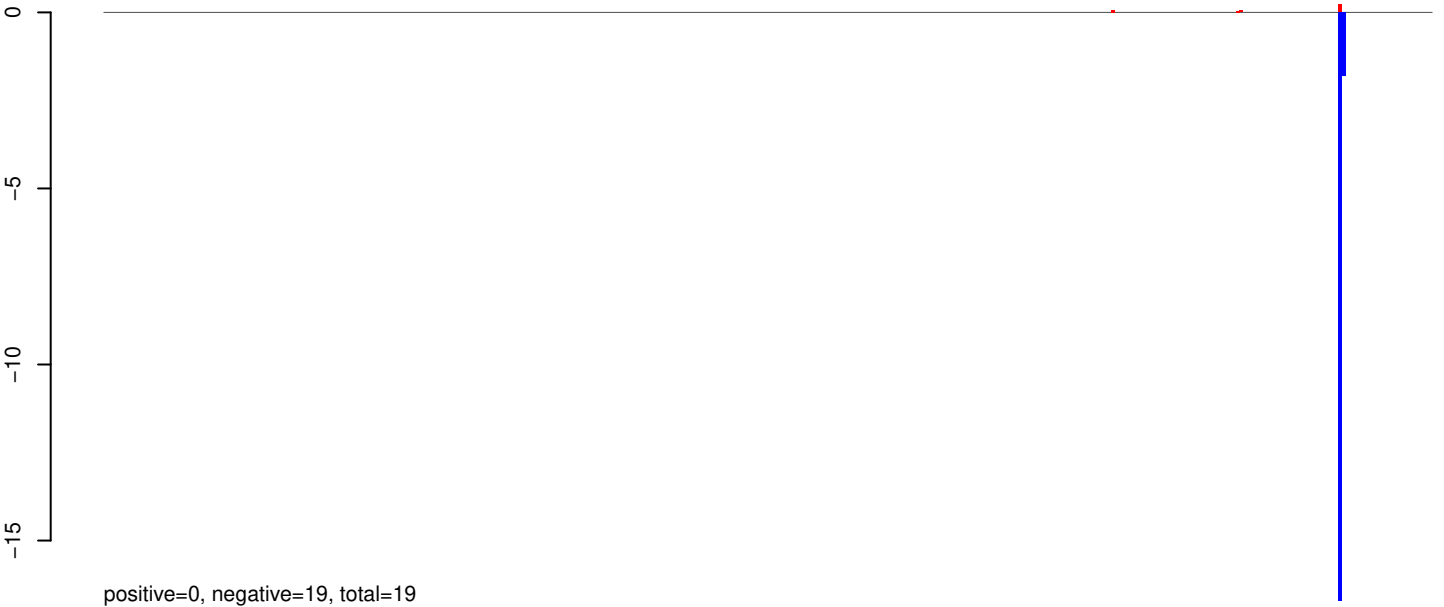
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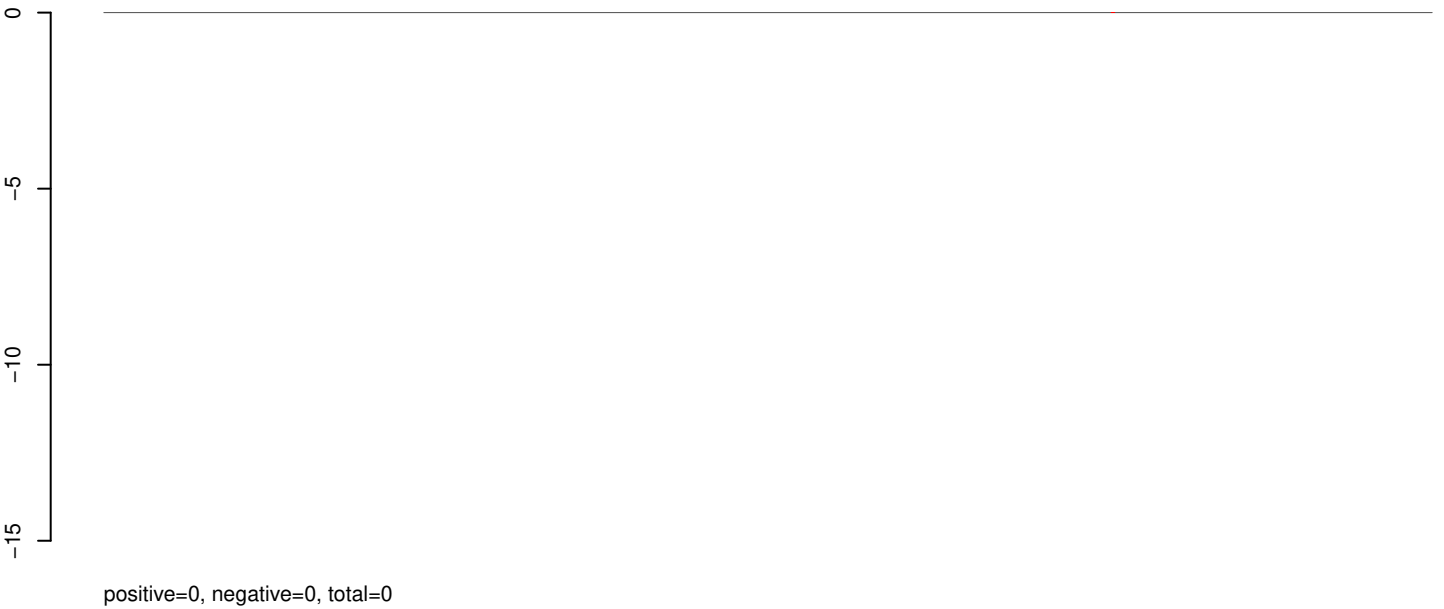
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0 2000 4000 6000 8000 10000

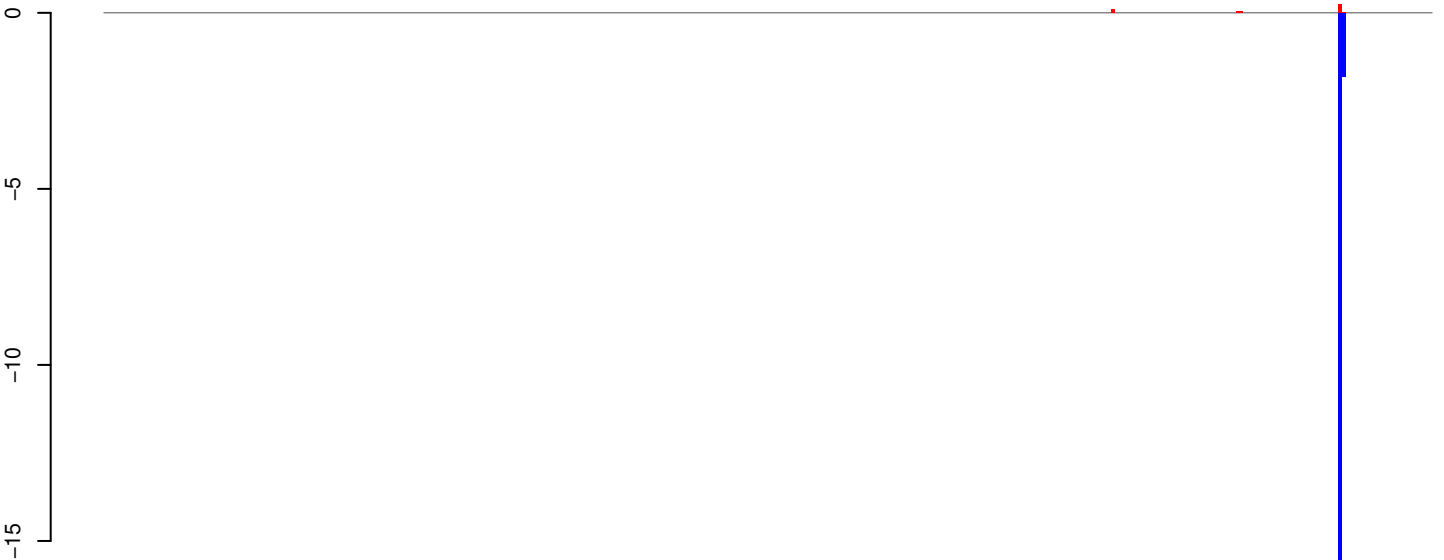
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



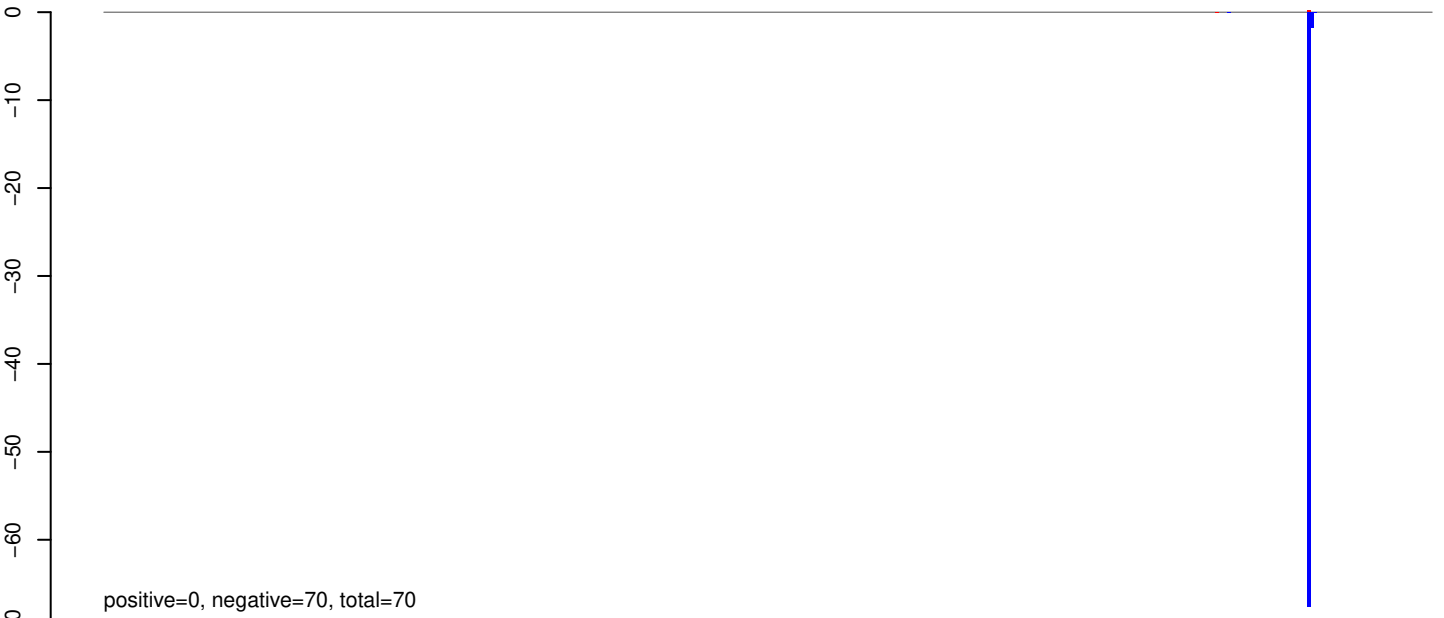
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Window size=25, length=10673, virus@NC_023424.1-Tyuleniy_virus_strain_LEIV-6C_polyprotein_gene:1-10673

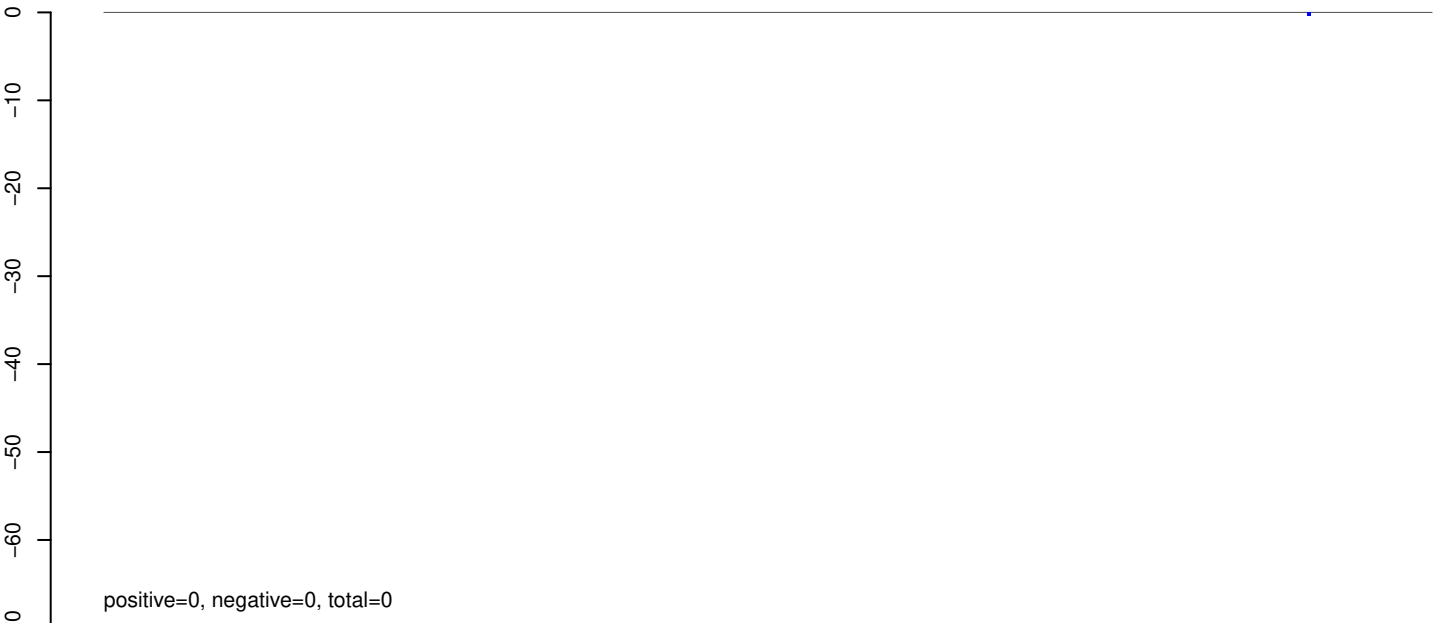
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AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



positive=0, negative=70, total=70

AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_R3D1B_r123_ZA.rep



positive=0, negative=70, total=70

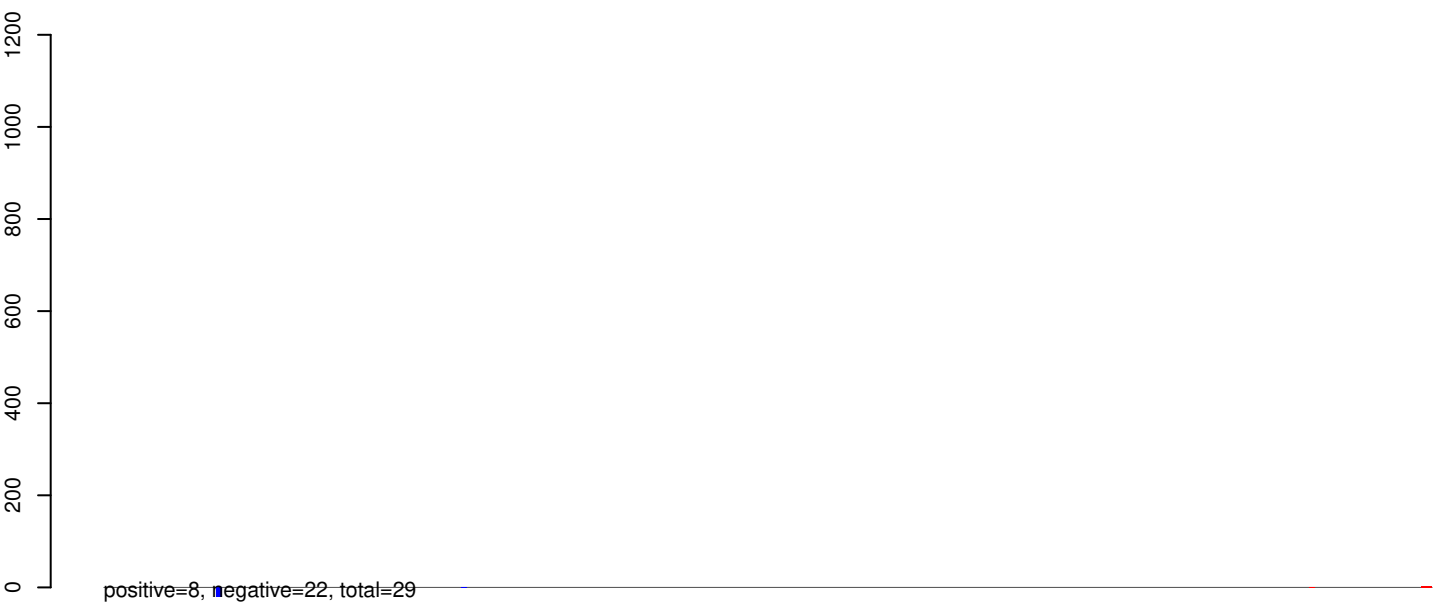
Window size=25, length=10874, virus@NC_009029.2-Kokobera_virus:1-10874

0 2000 4000 6000 8000 10000

AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep

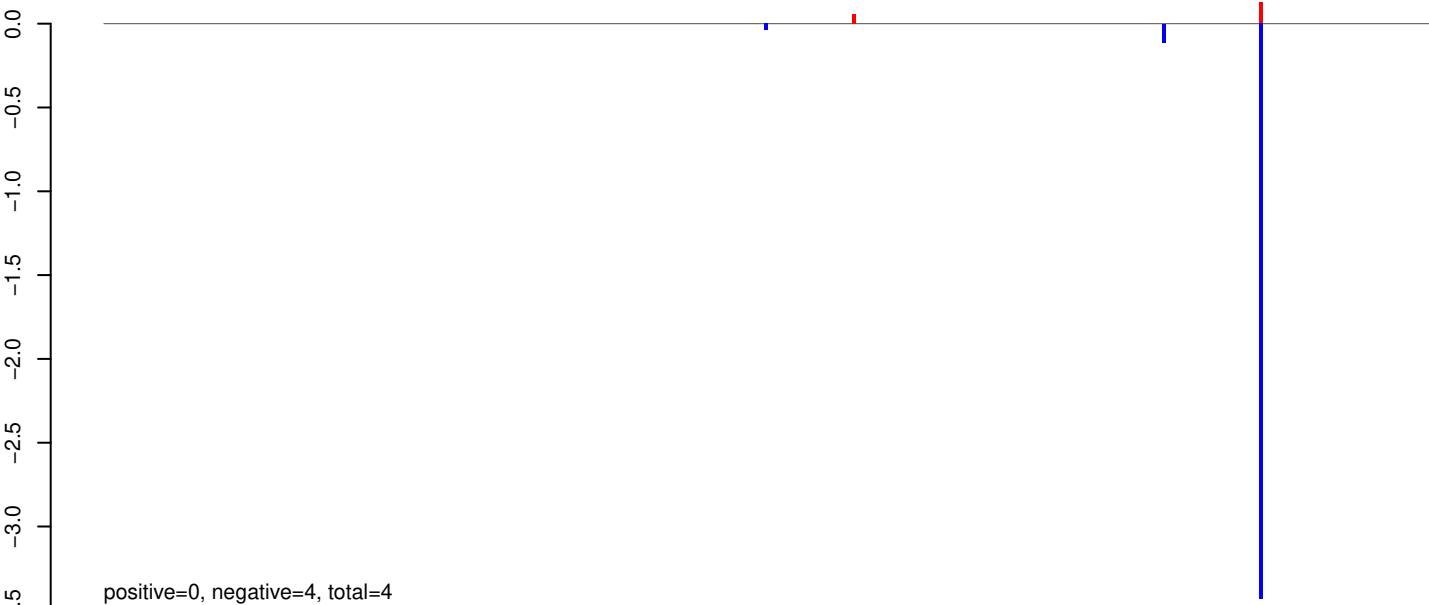


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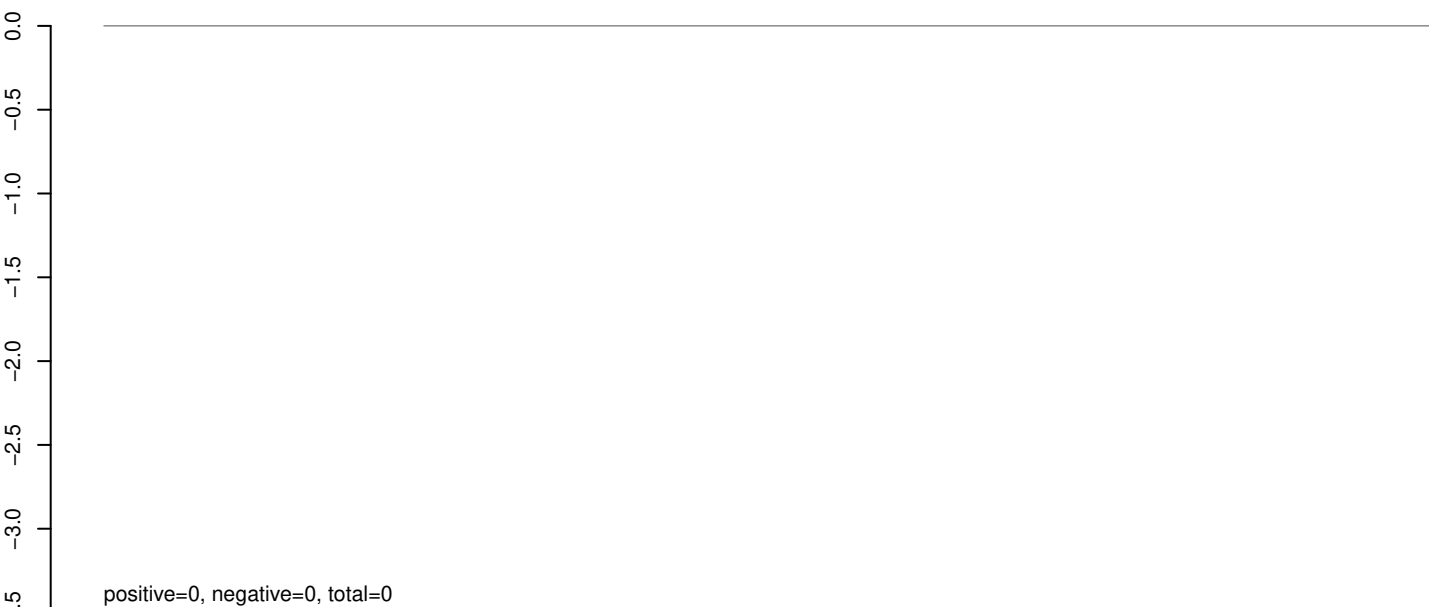


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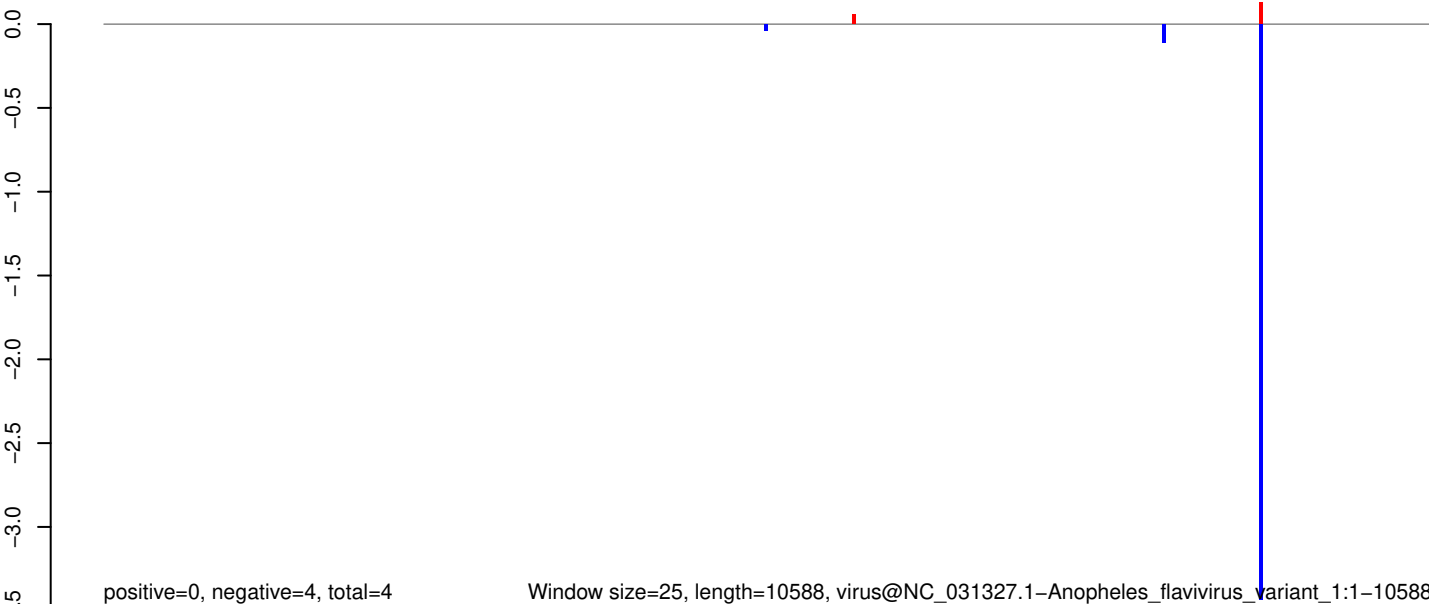
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



AeAeg_Aag2_R3D1B_r123_ZA.rep



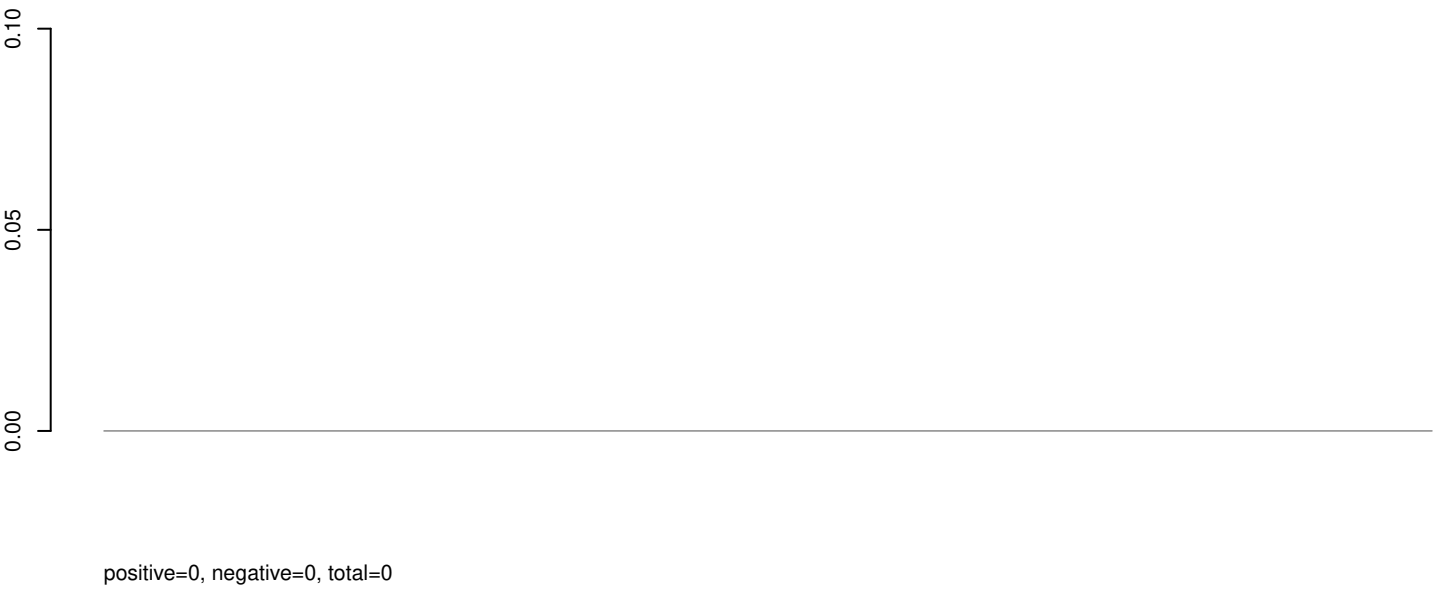
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0 2000 4000 6000 8000 10000

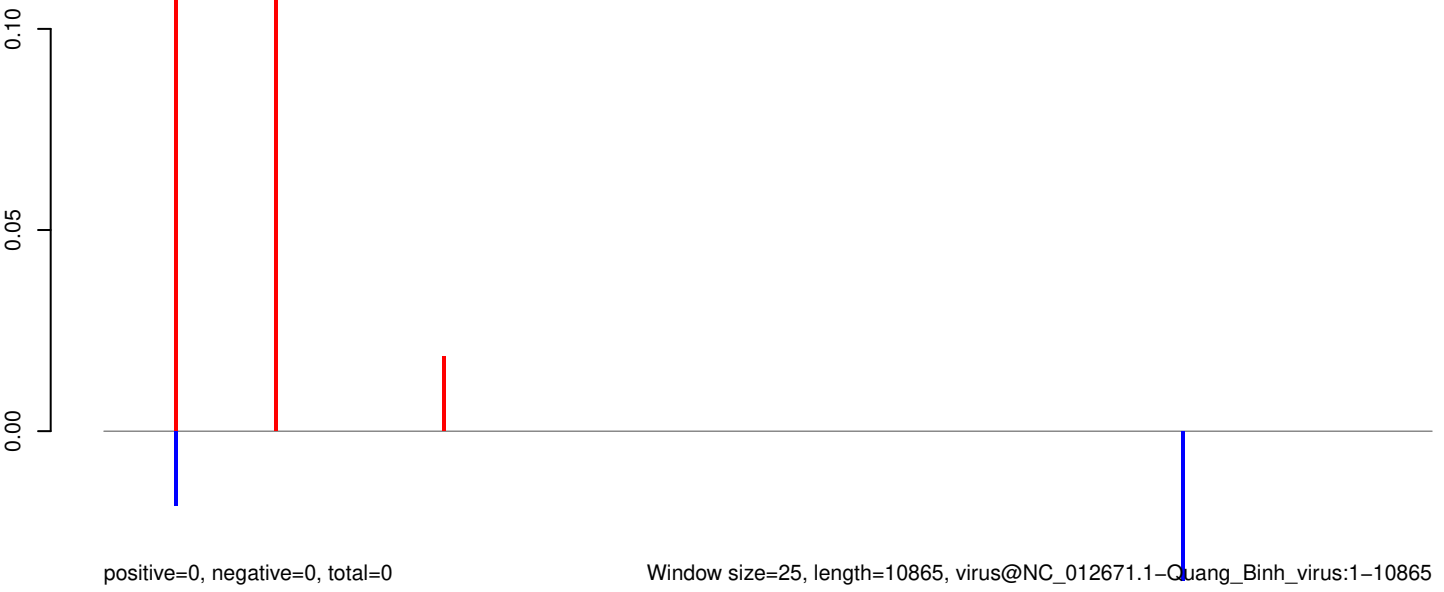
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



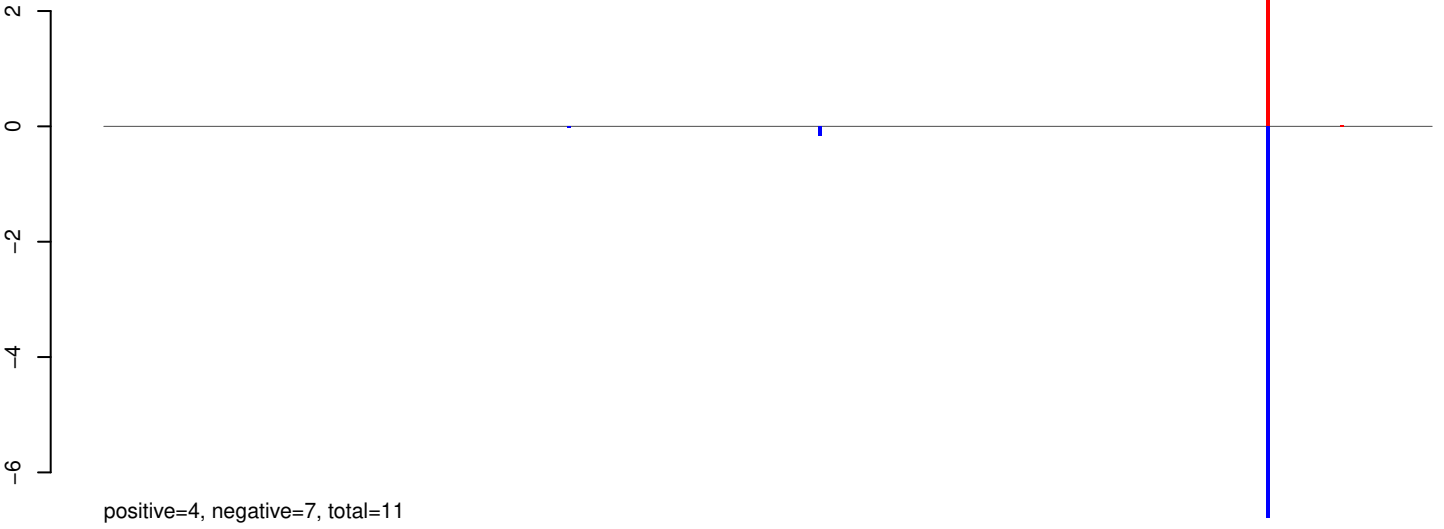
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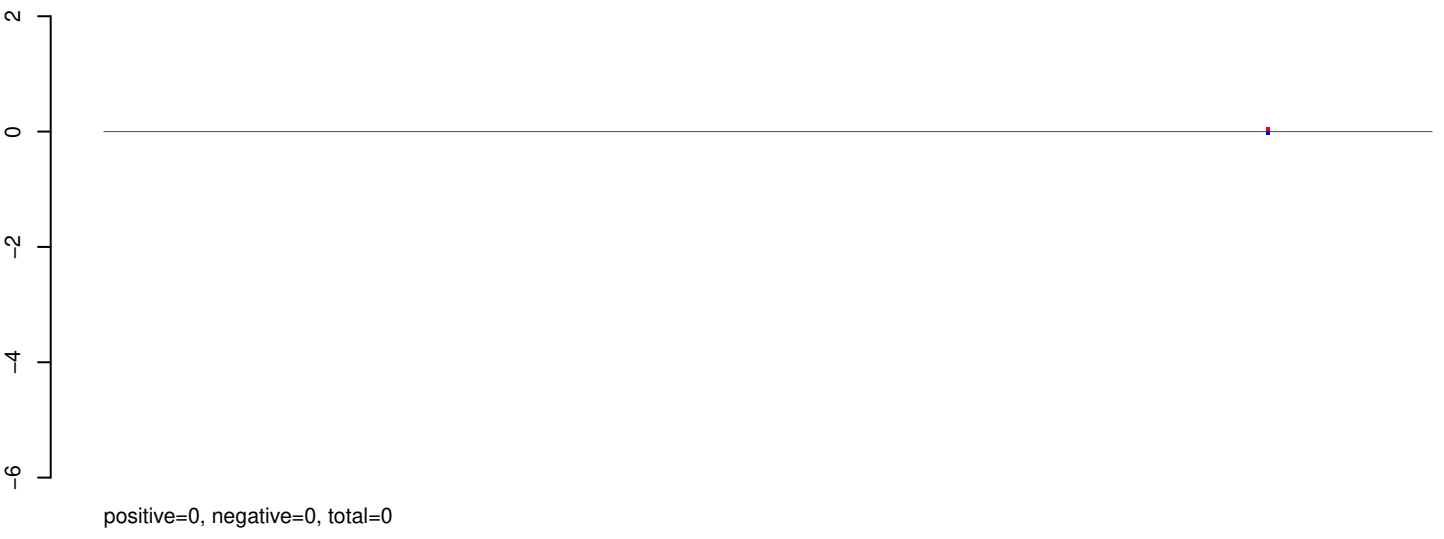
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0 2000 4000 6000 8000 10000

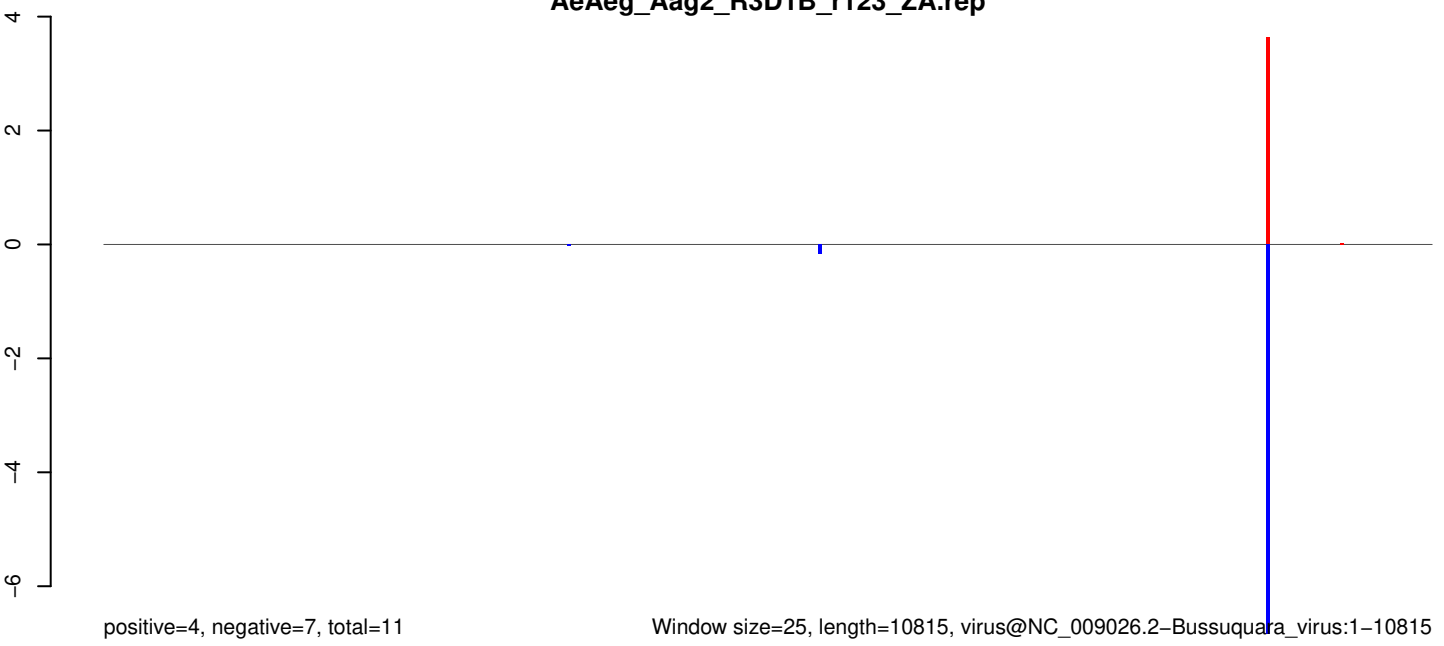
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



AeAeg_Aag2_R3D1B_r123_ZA.rep

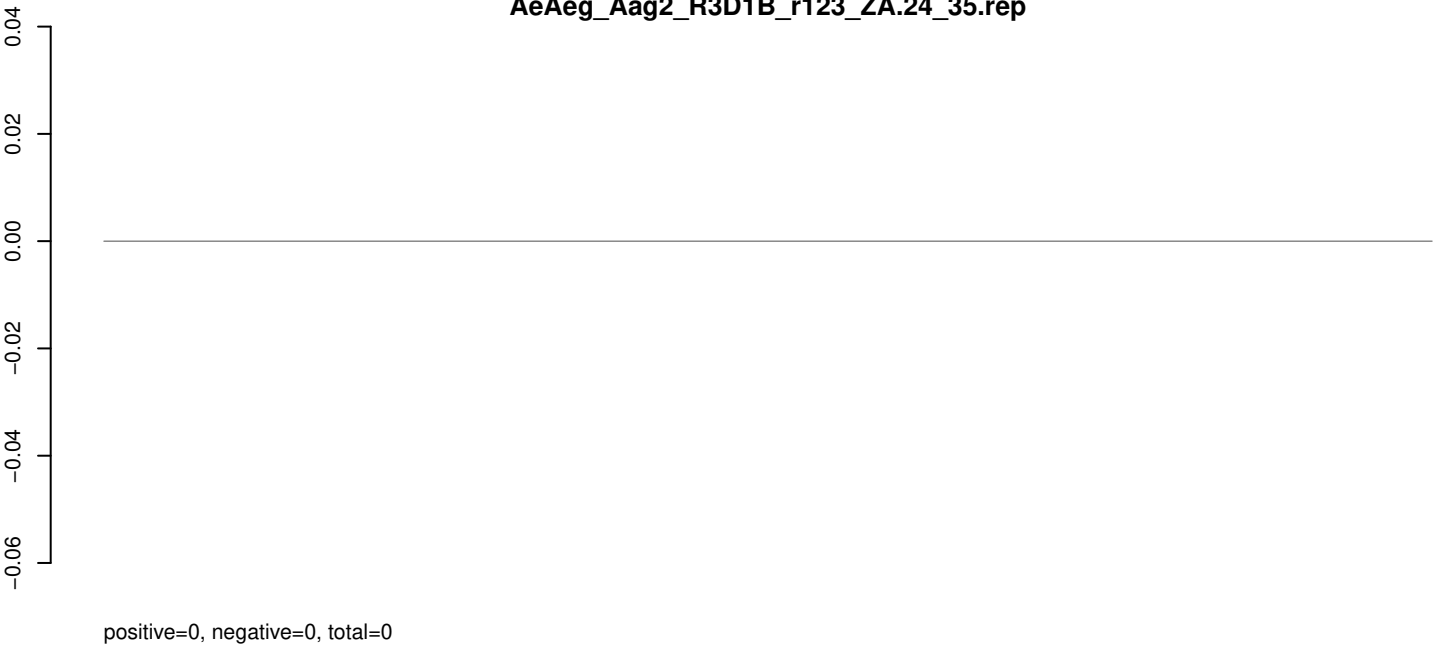


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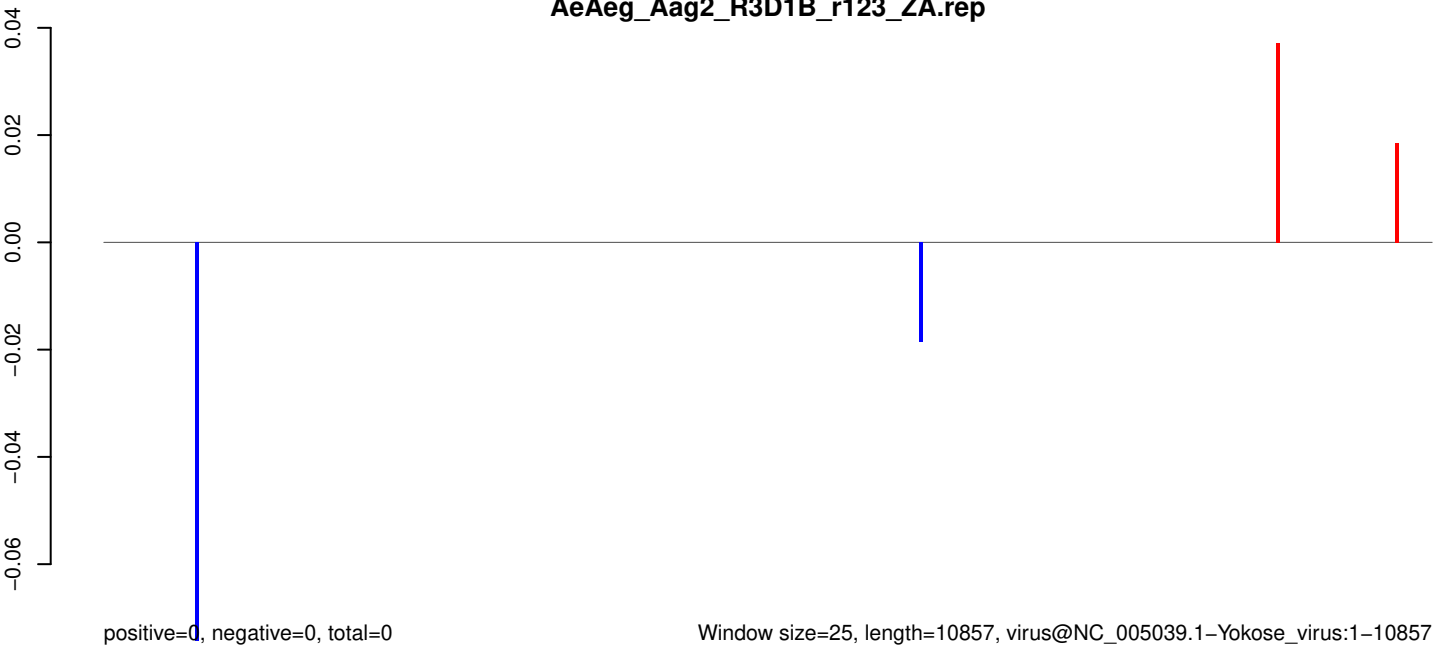
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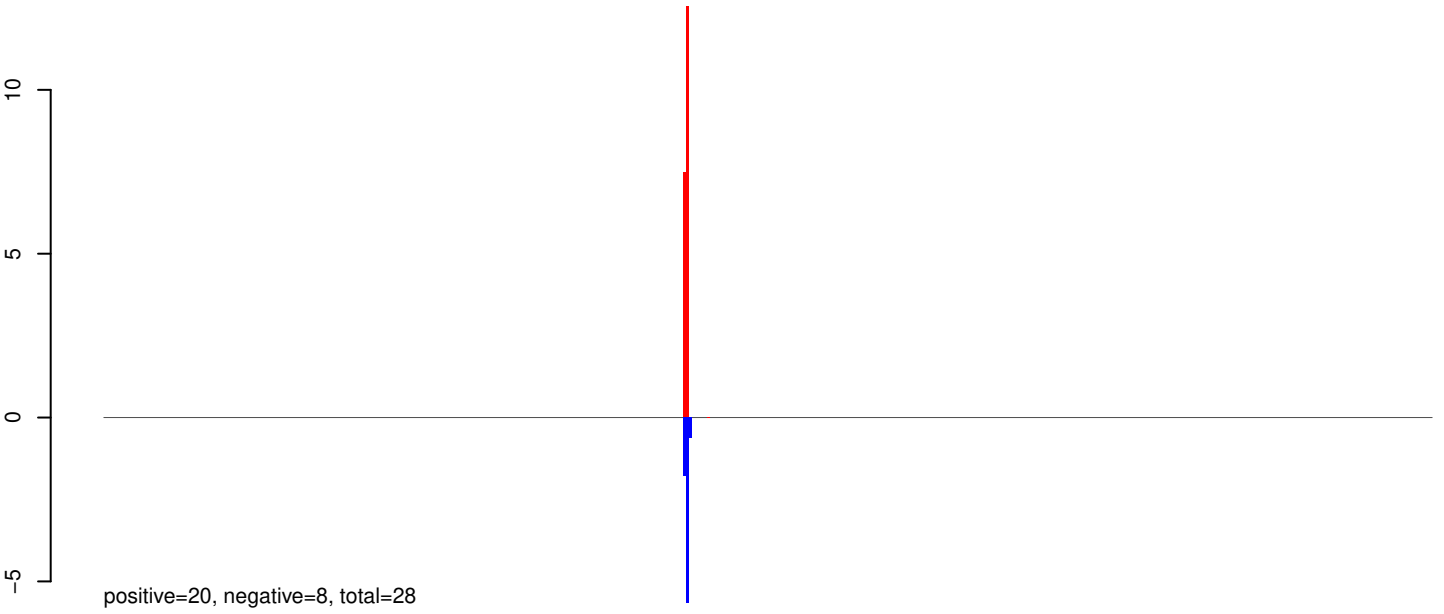
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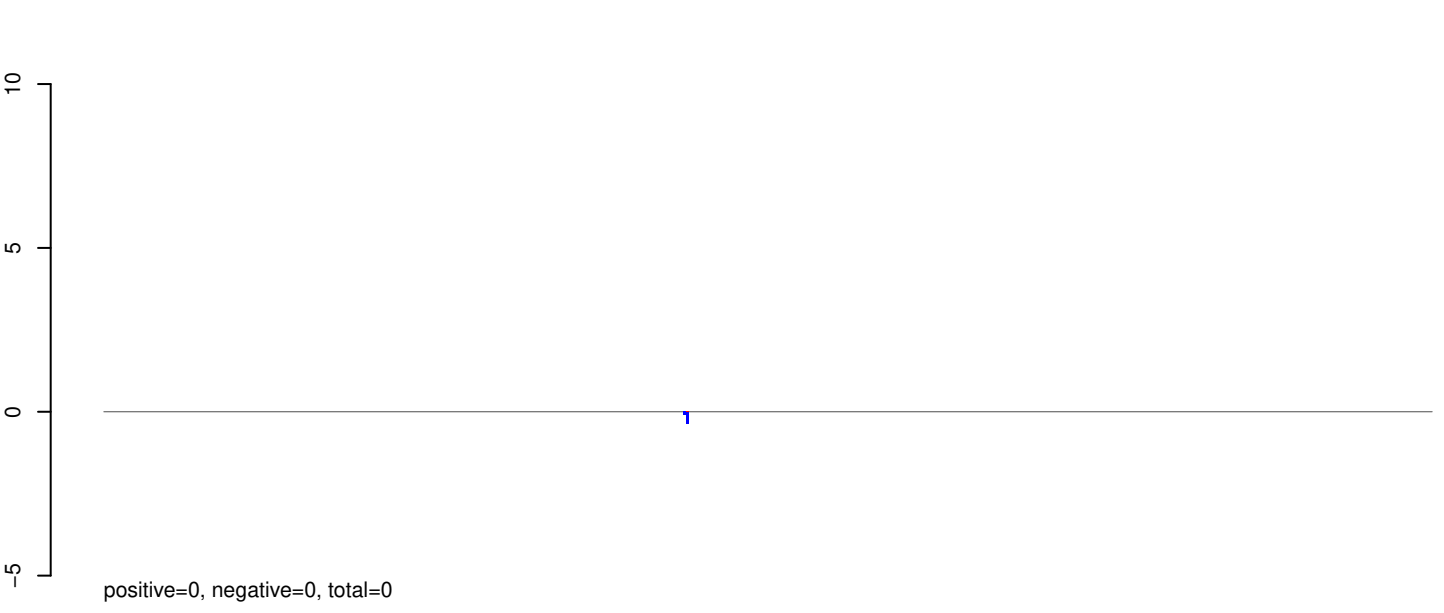
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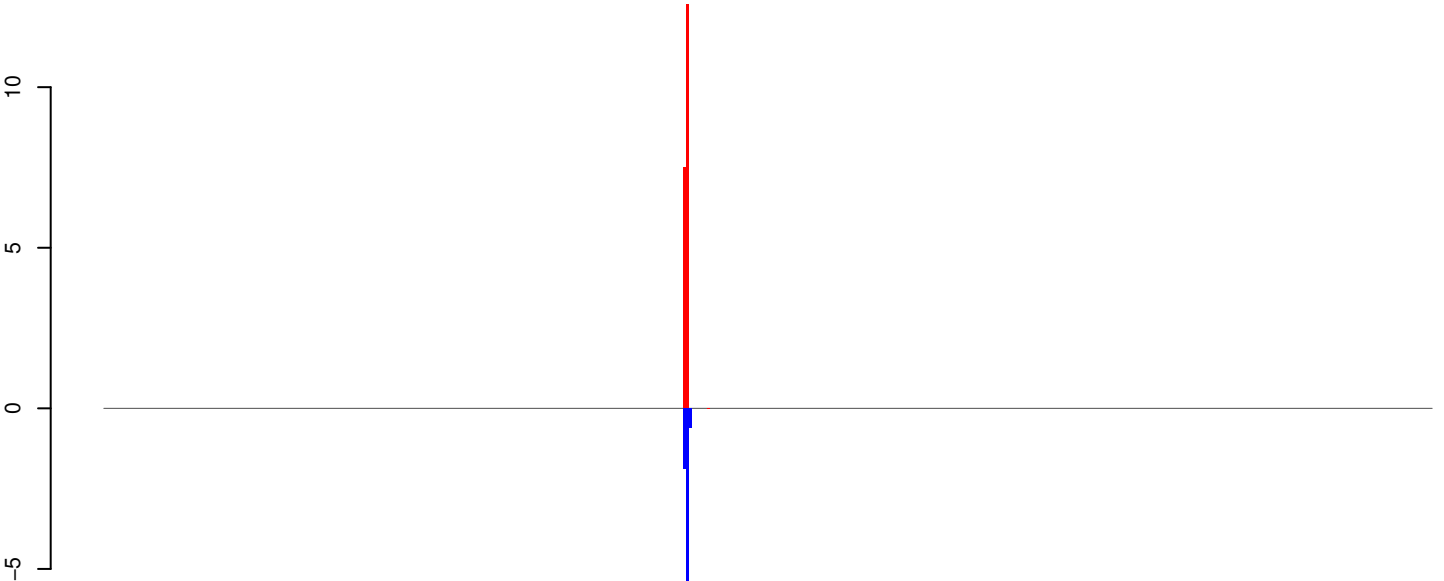
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



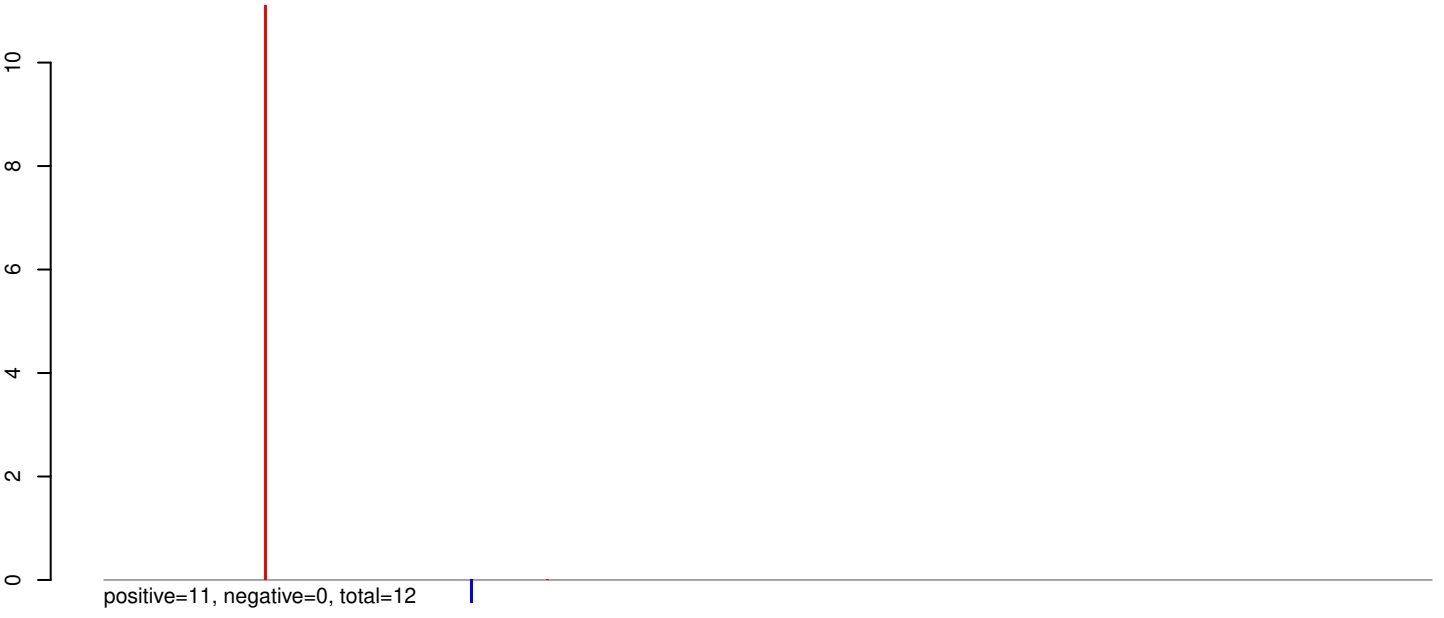
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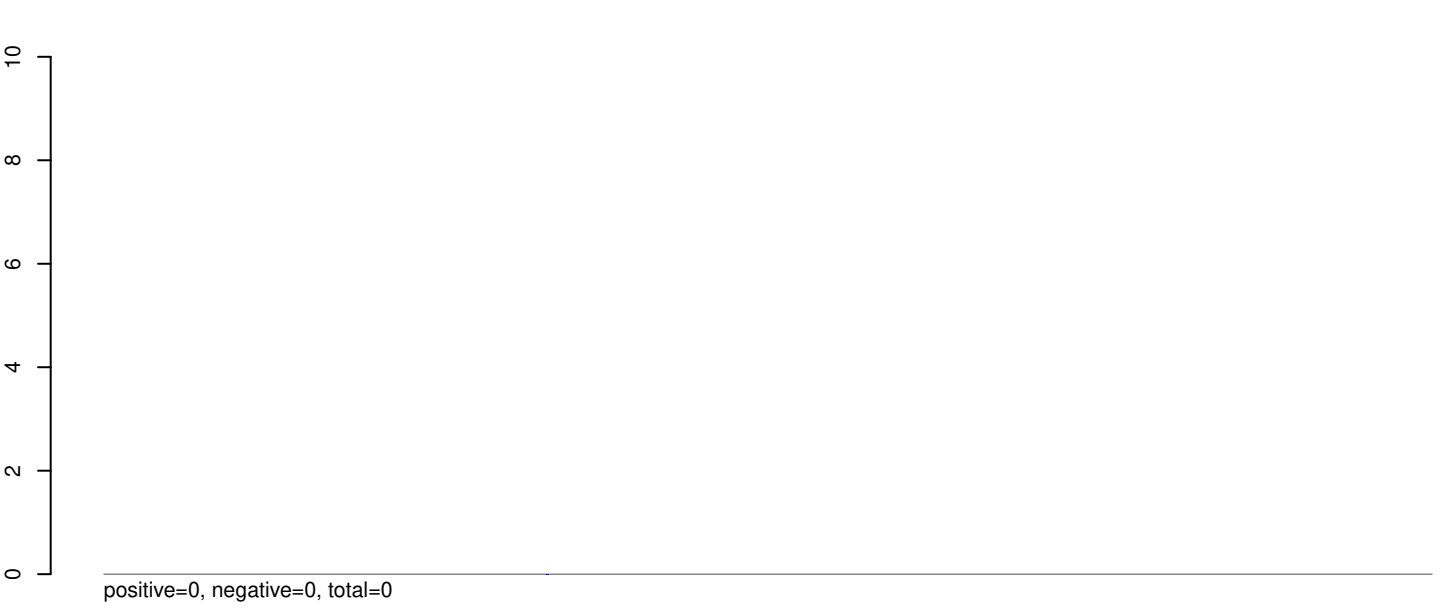
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0 2000 4000 6000 8000 10000 12000

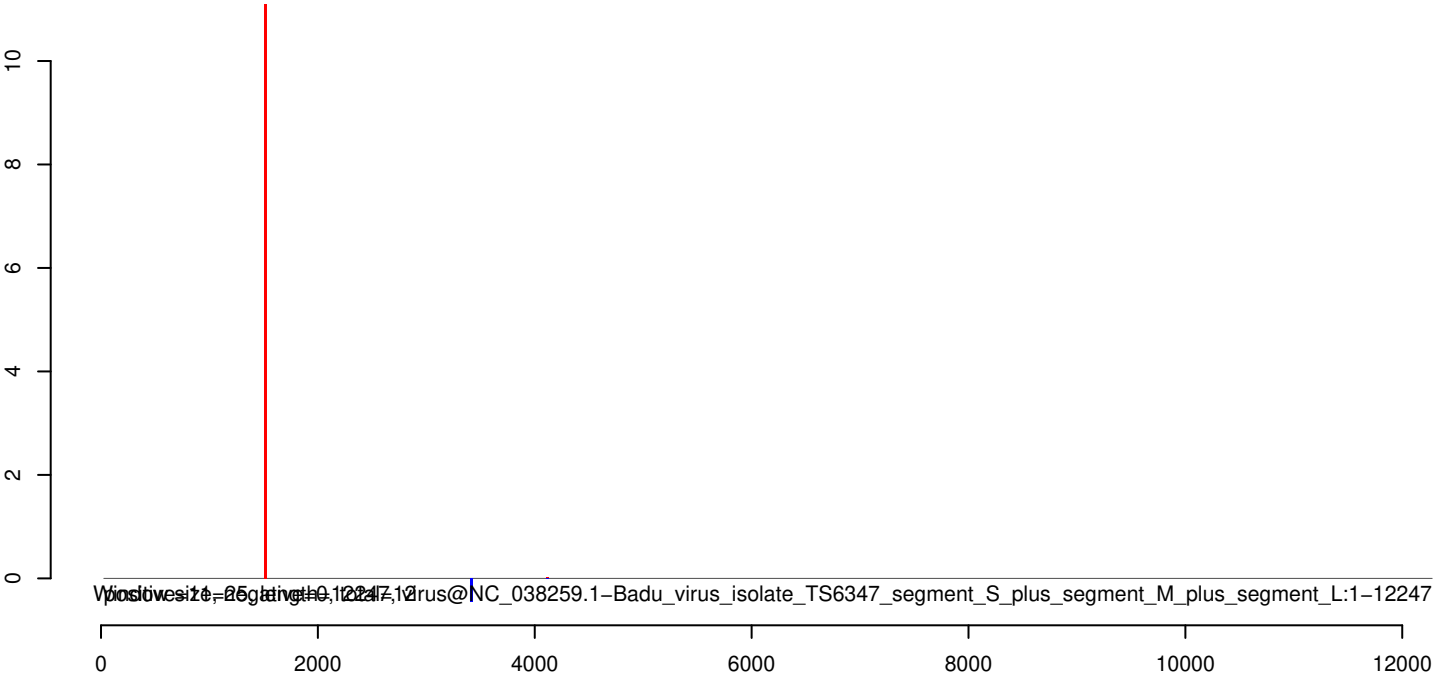
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



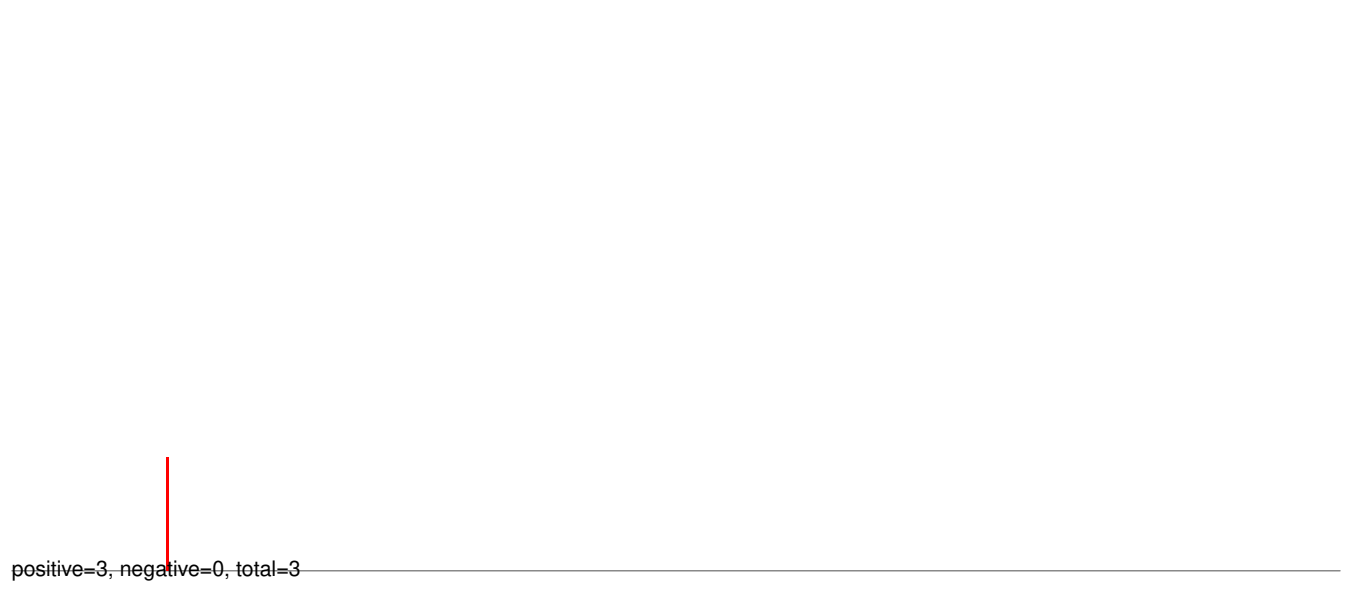
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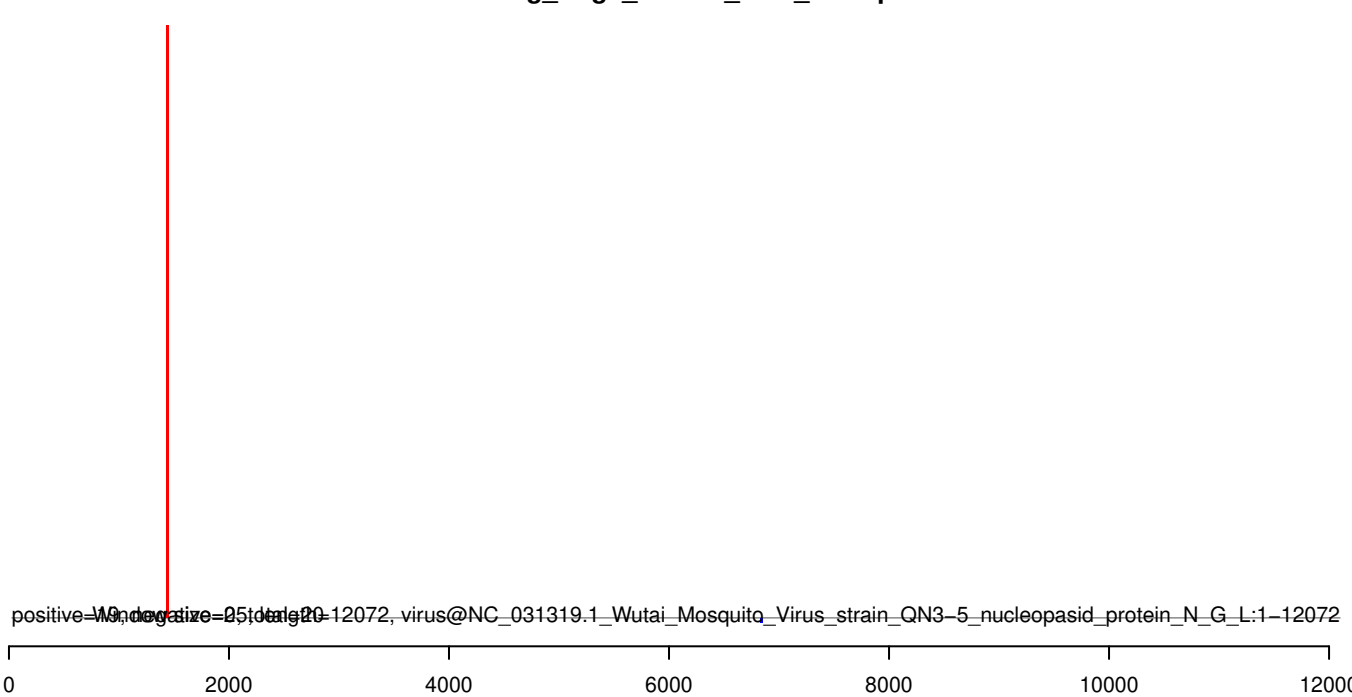
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep

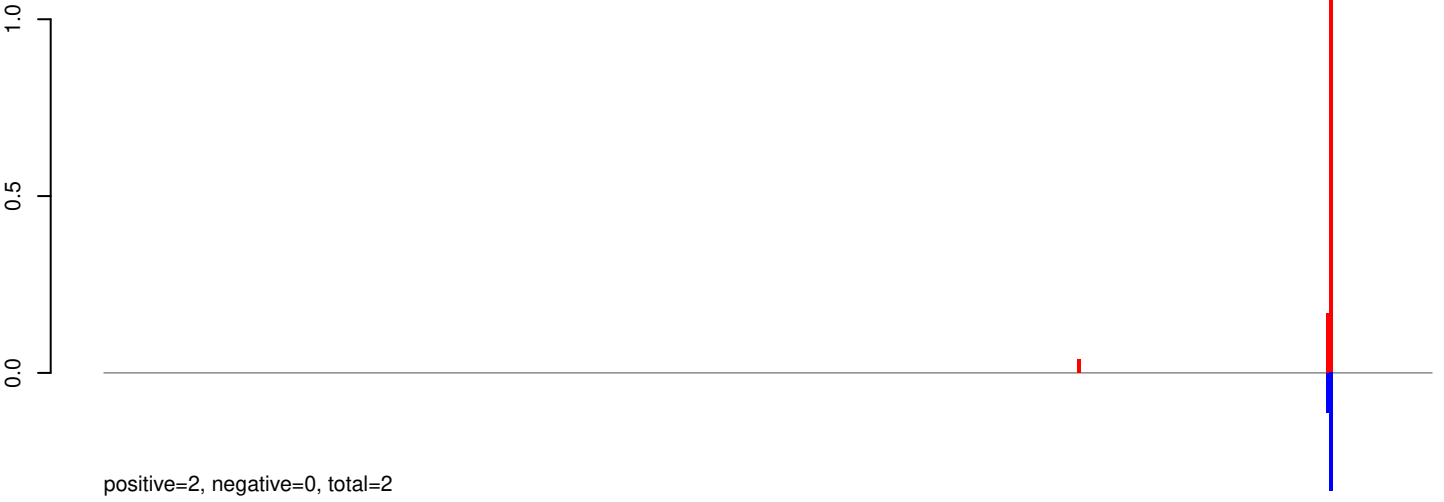


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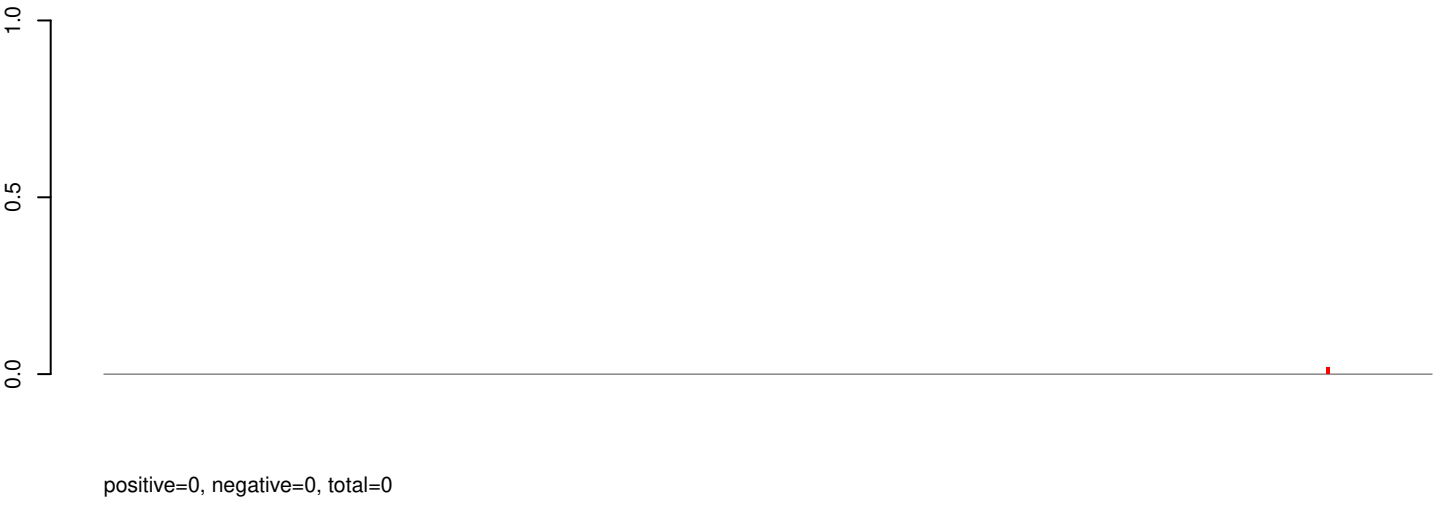


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1-12072, virus@NC_031319.1_Wutai_Mosquito_Virus_strain_QN3-5_nucleopasid_protein_N_G_L:1-12072

AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



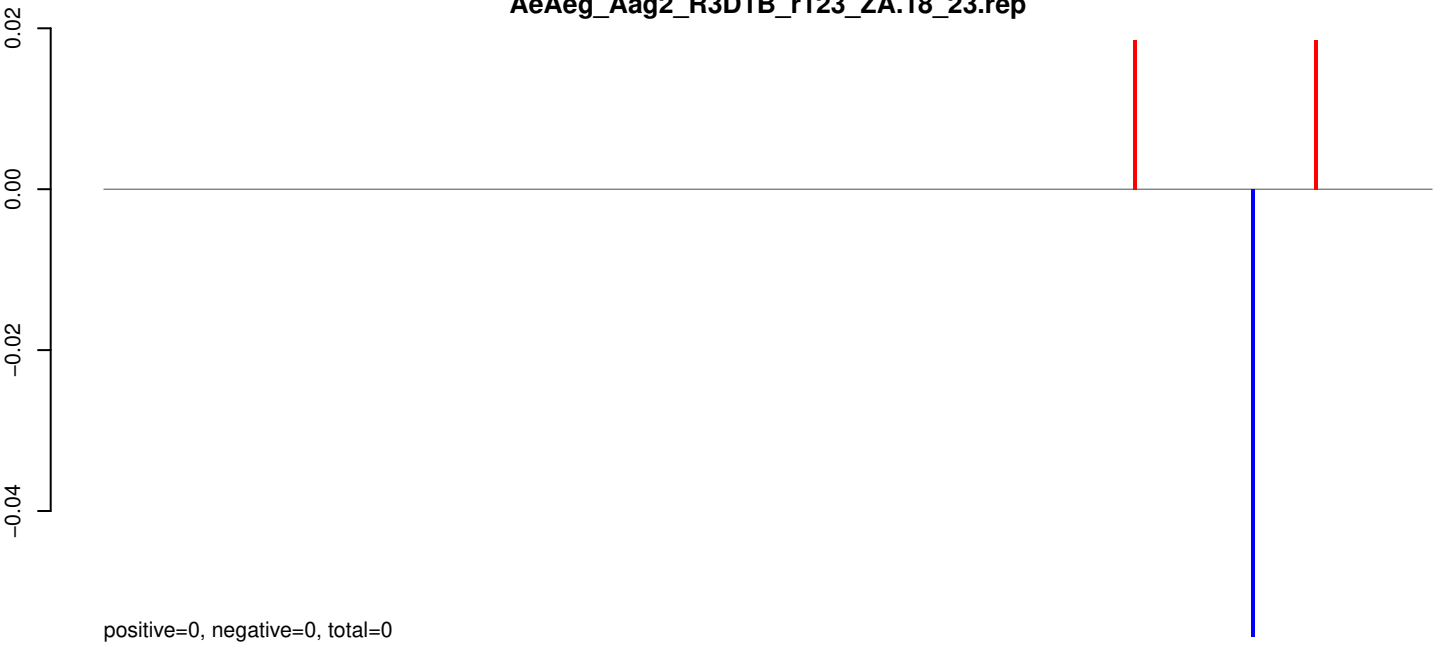
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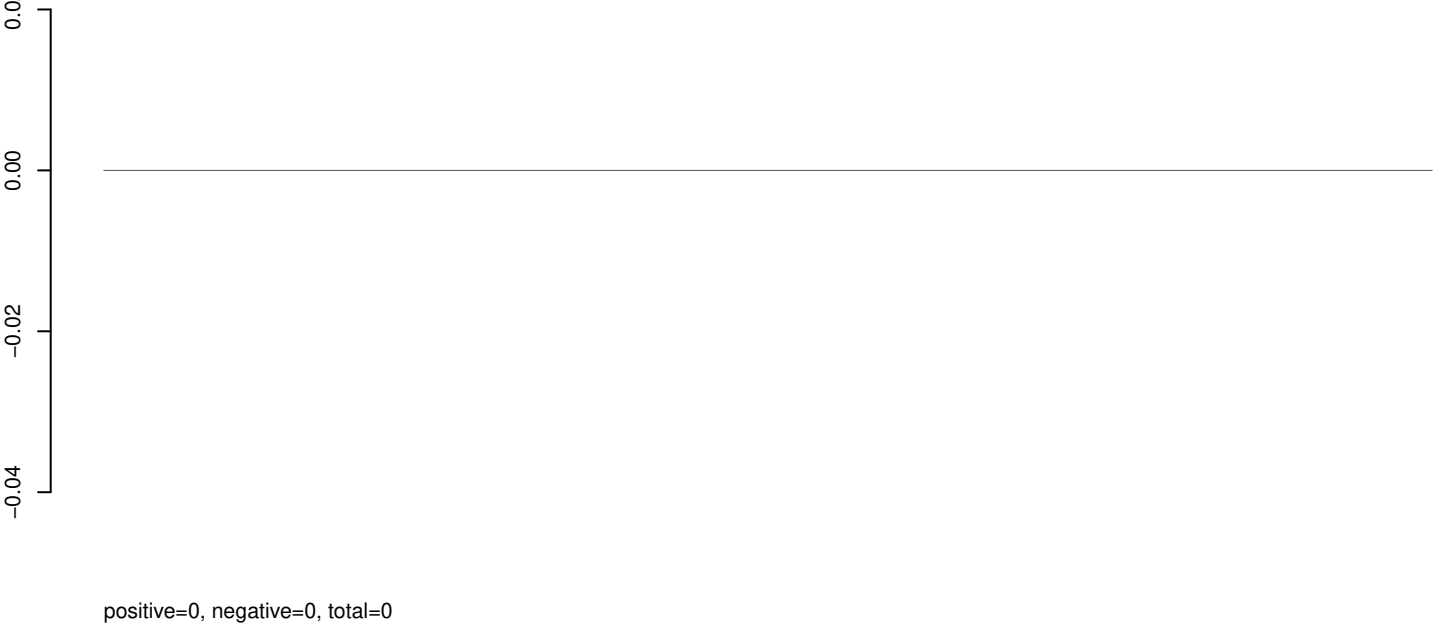
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0 2000 4000 6000 8000 10000

AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



AeAeg_Aag2_R3D1B_r123_ZA.rep



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

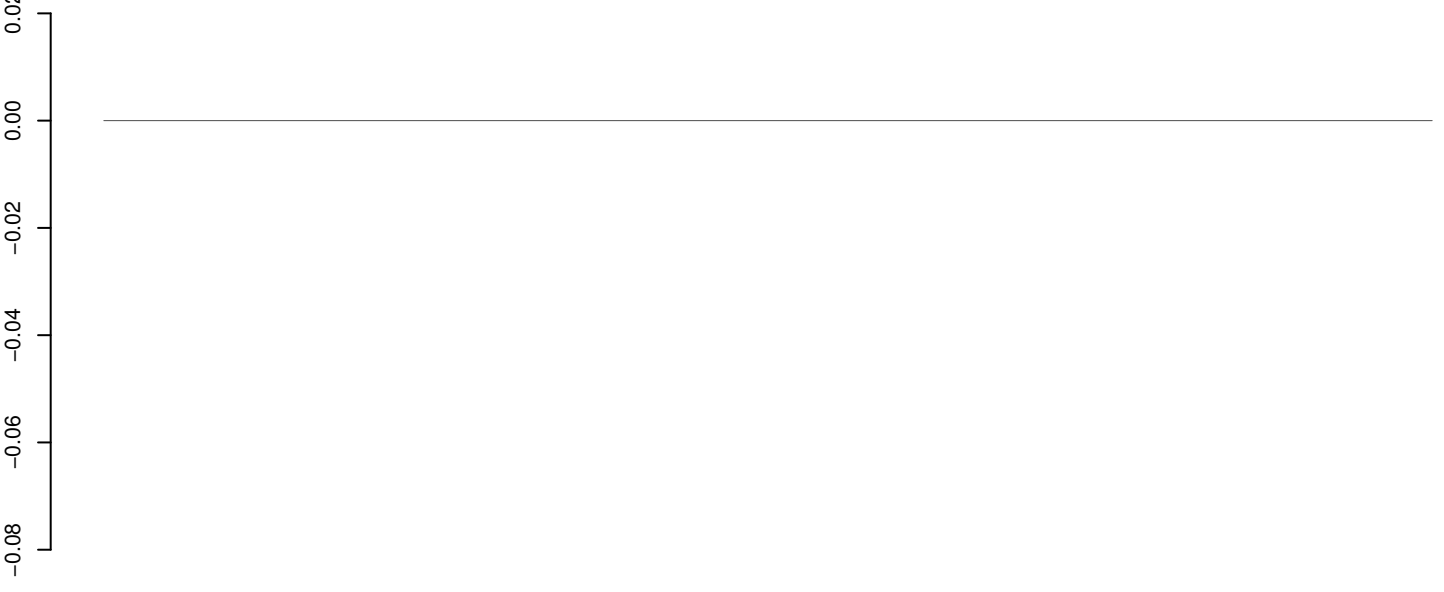
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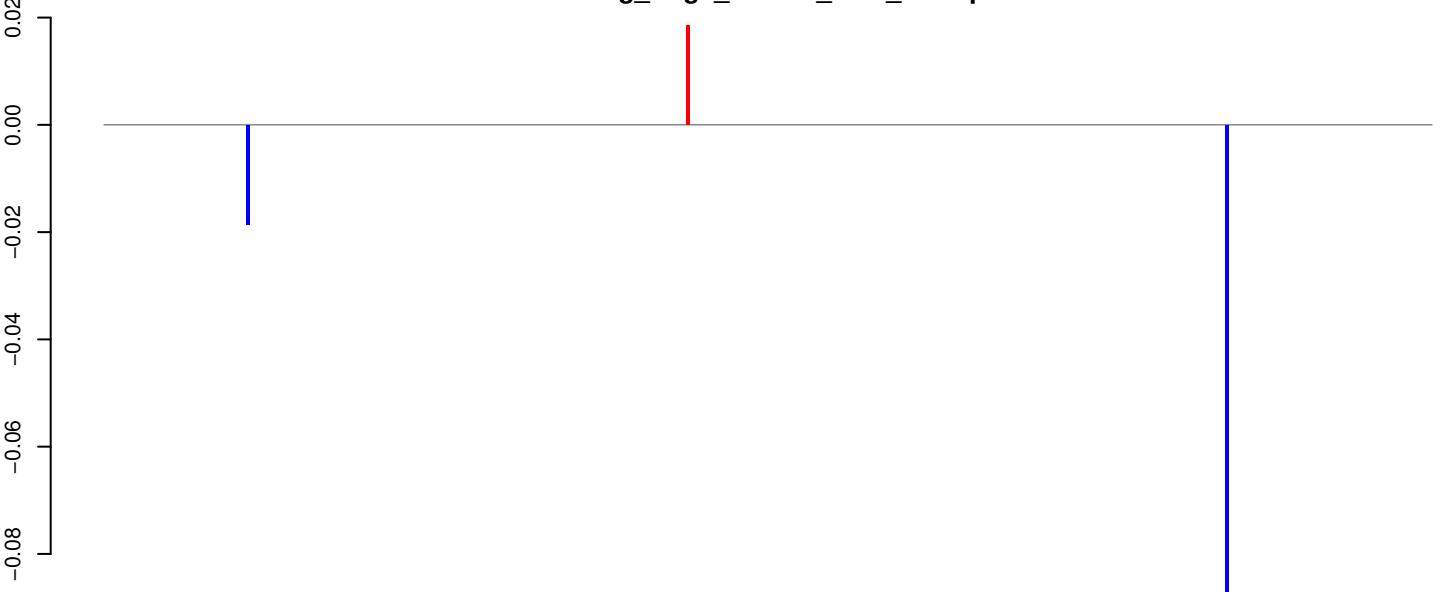
positive=0, negative=0, total=0

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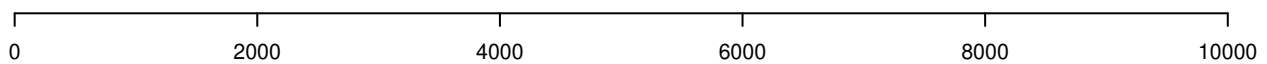
positive=0, negative=0, total=0

AeAeg_Aag2_R3D1B_r123_ZA.rep

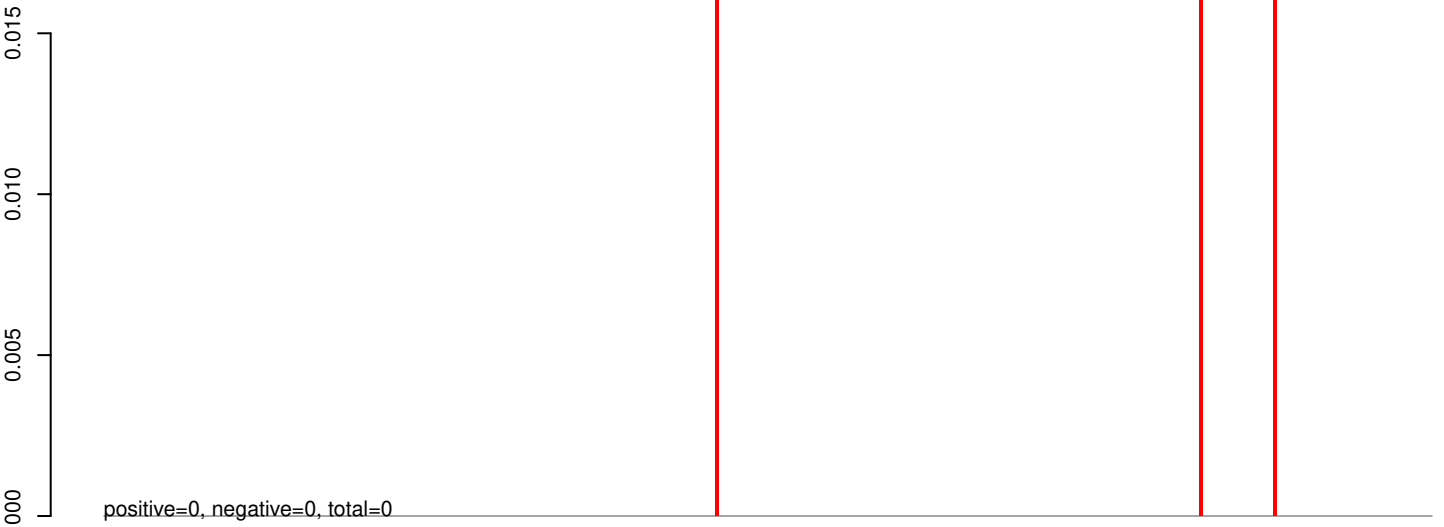


positive=0, negative=0, total=0

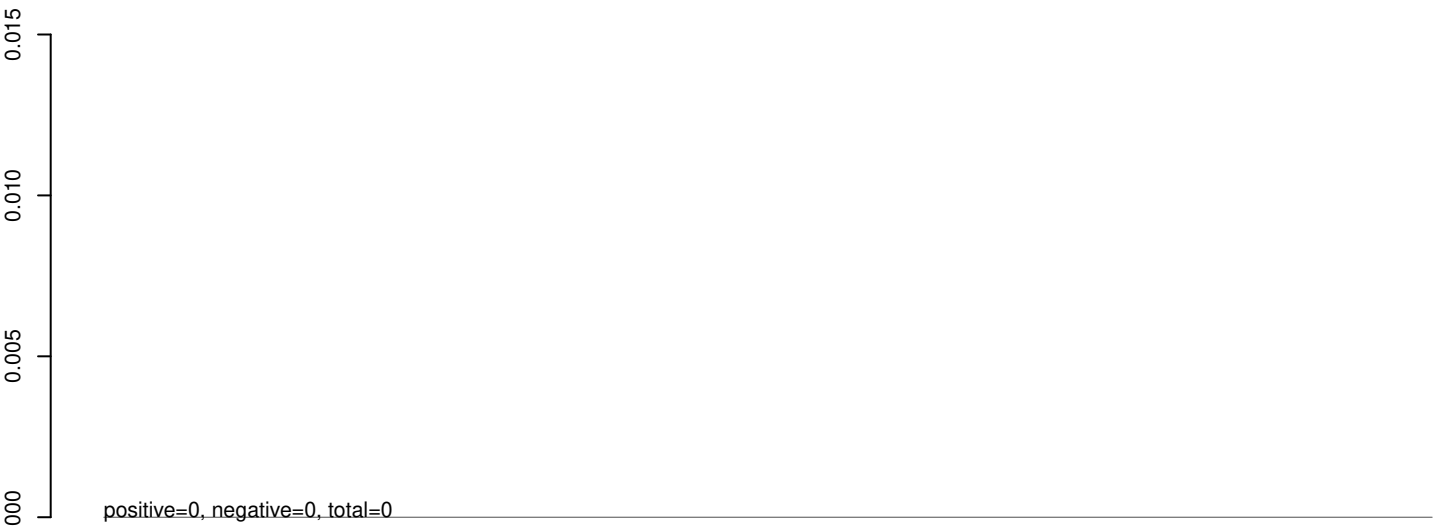
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AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep

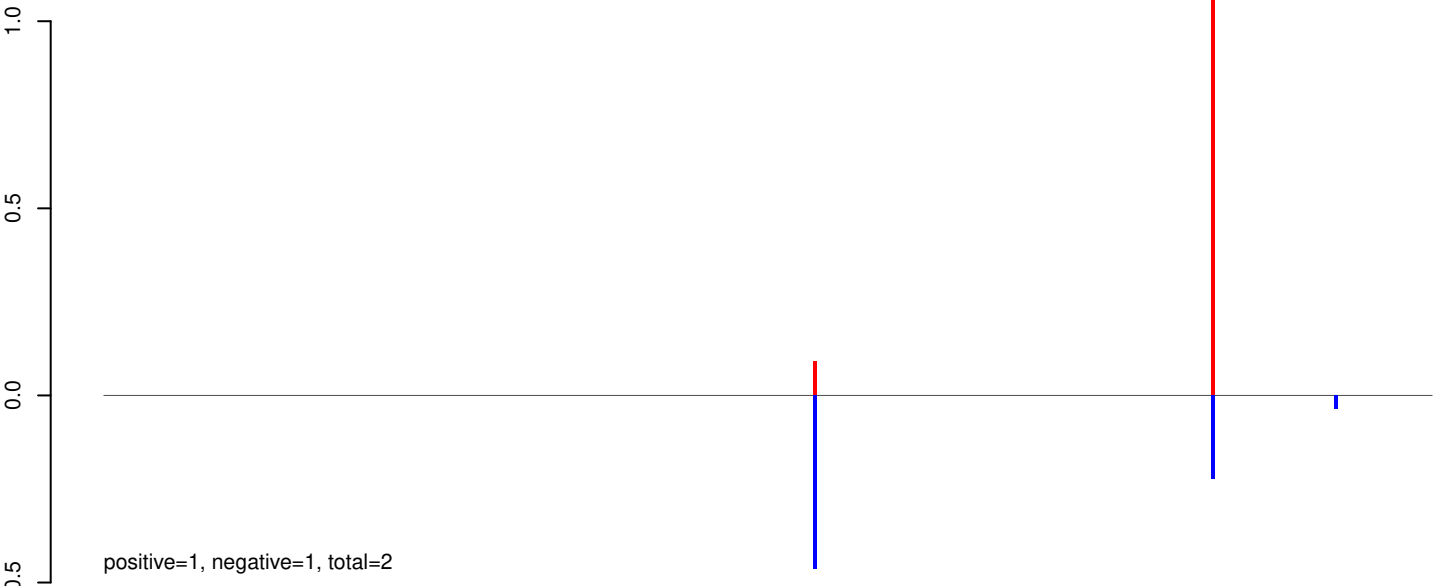


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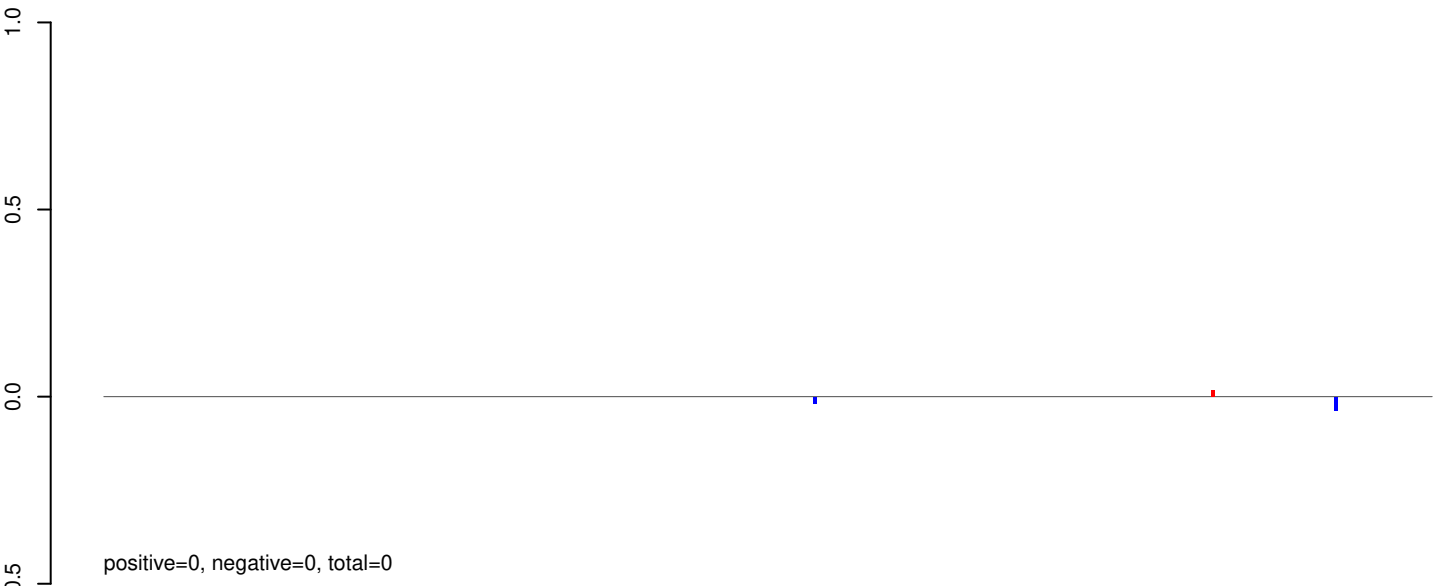


0 2000 4000 6000 8000 10000

AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep

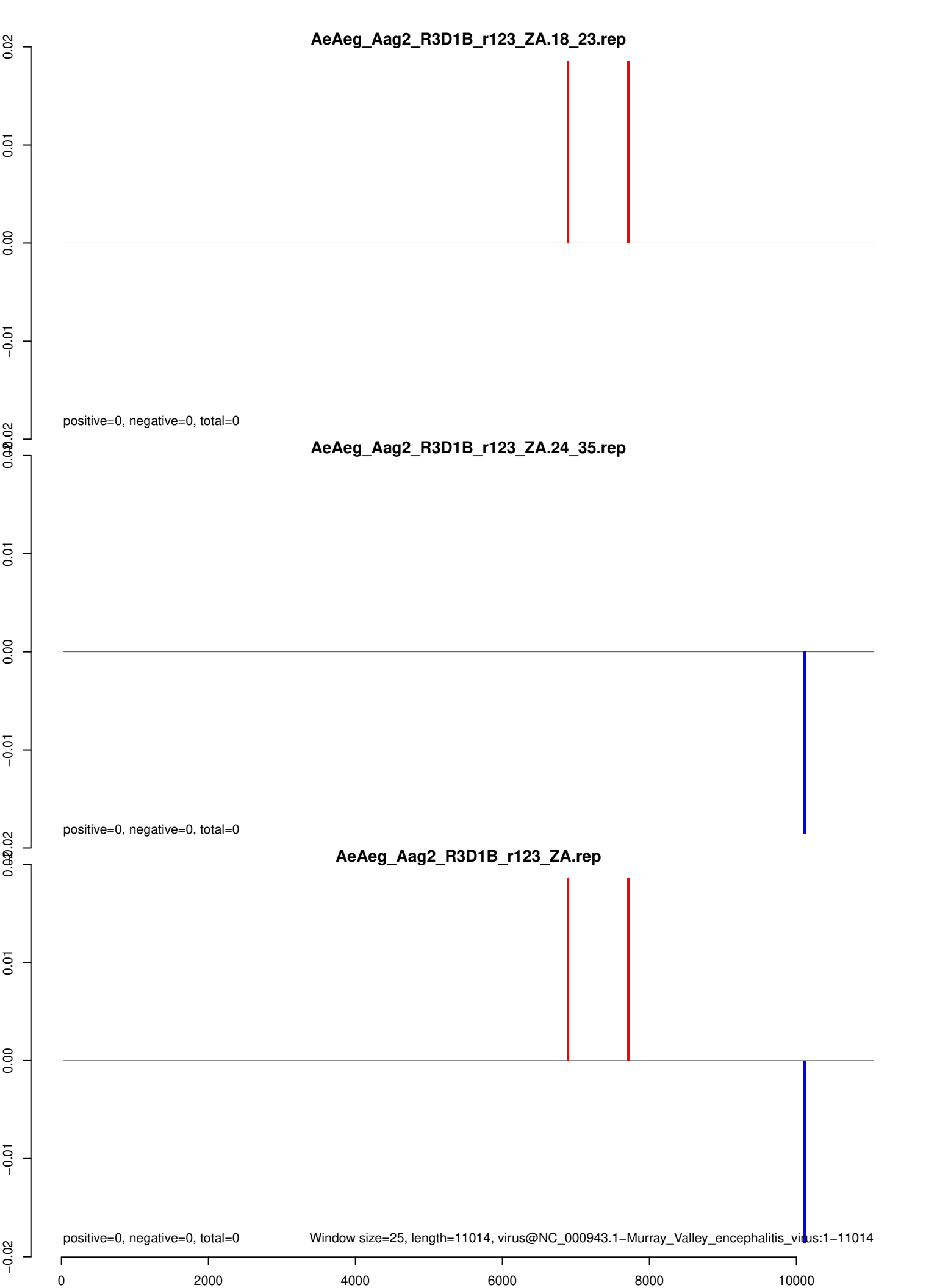


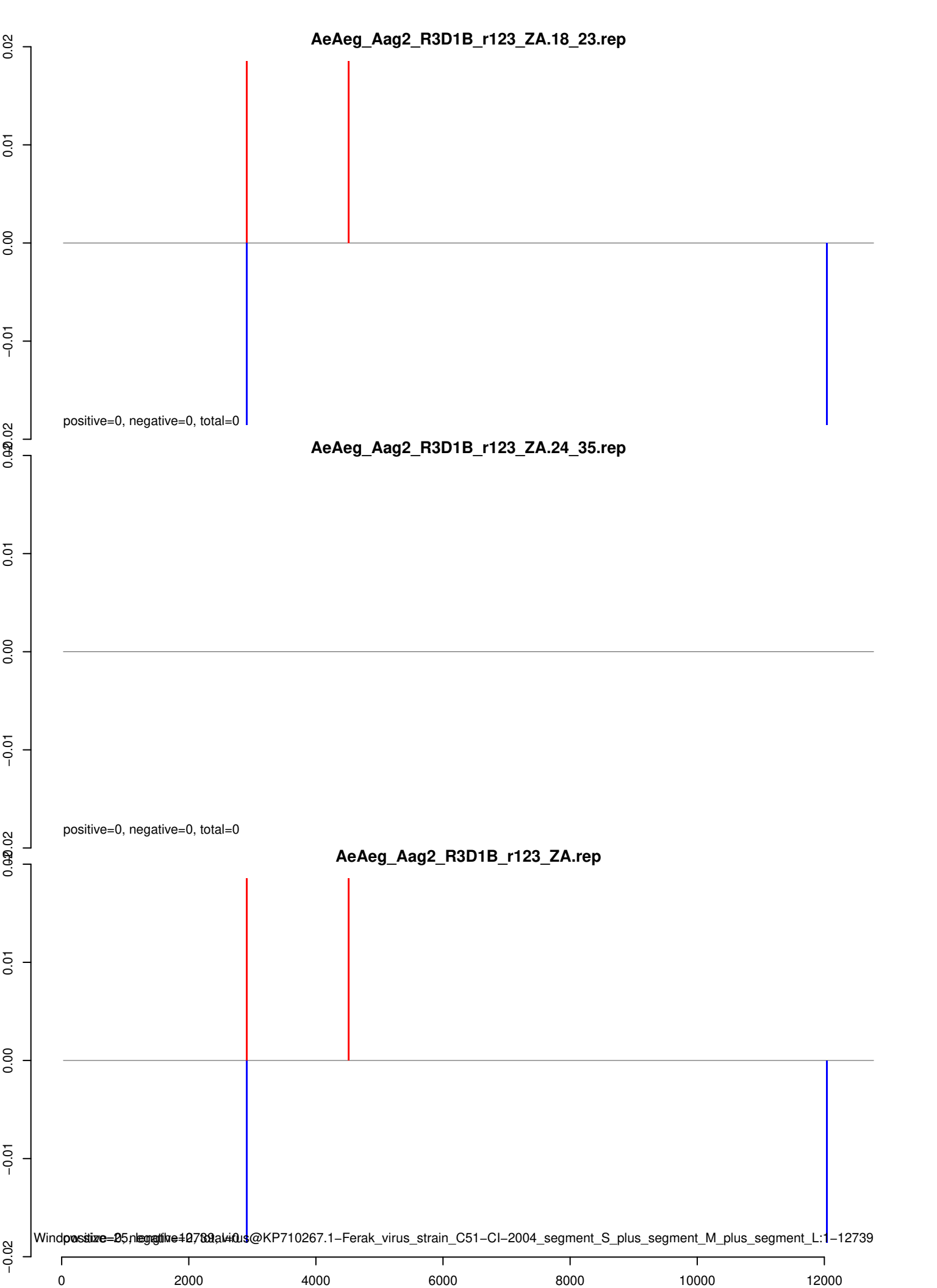
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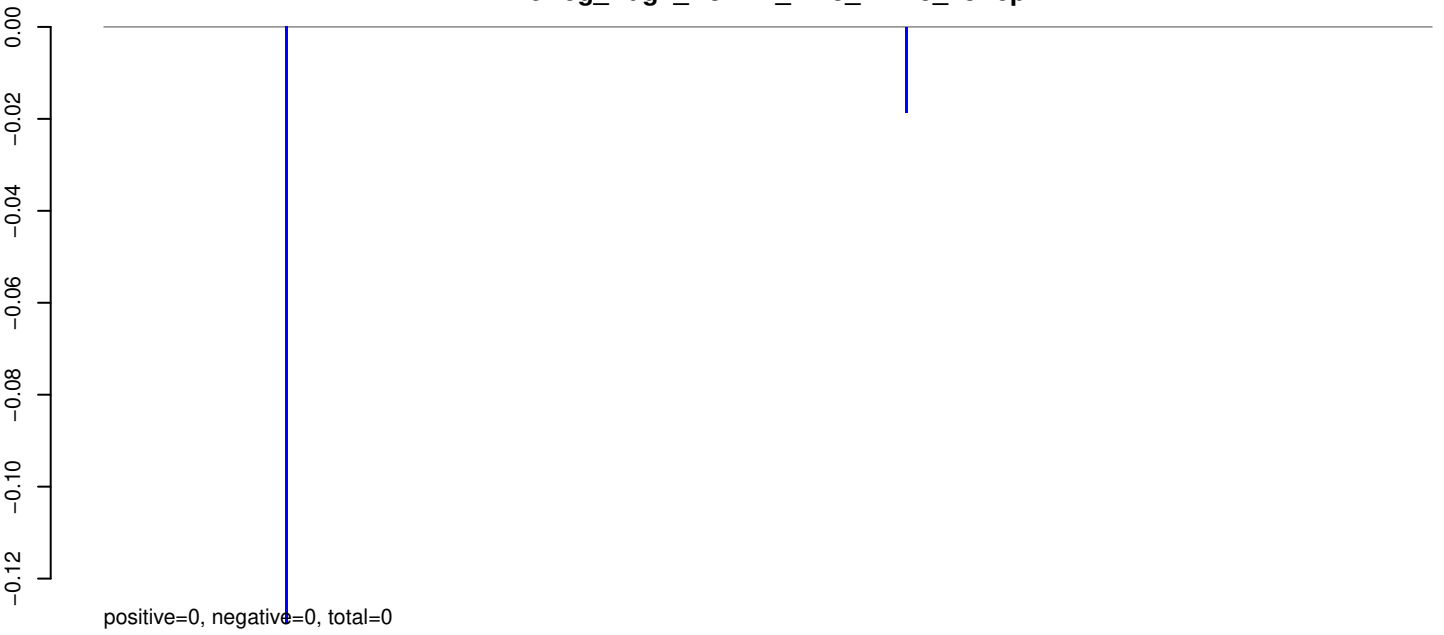
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0 2000 4000 6000 8000 10000

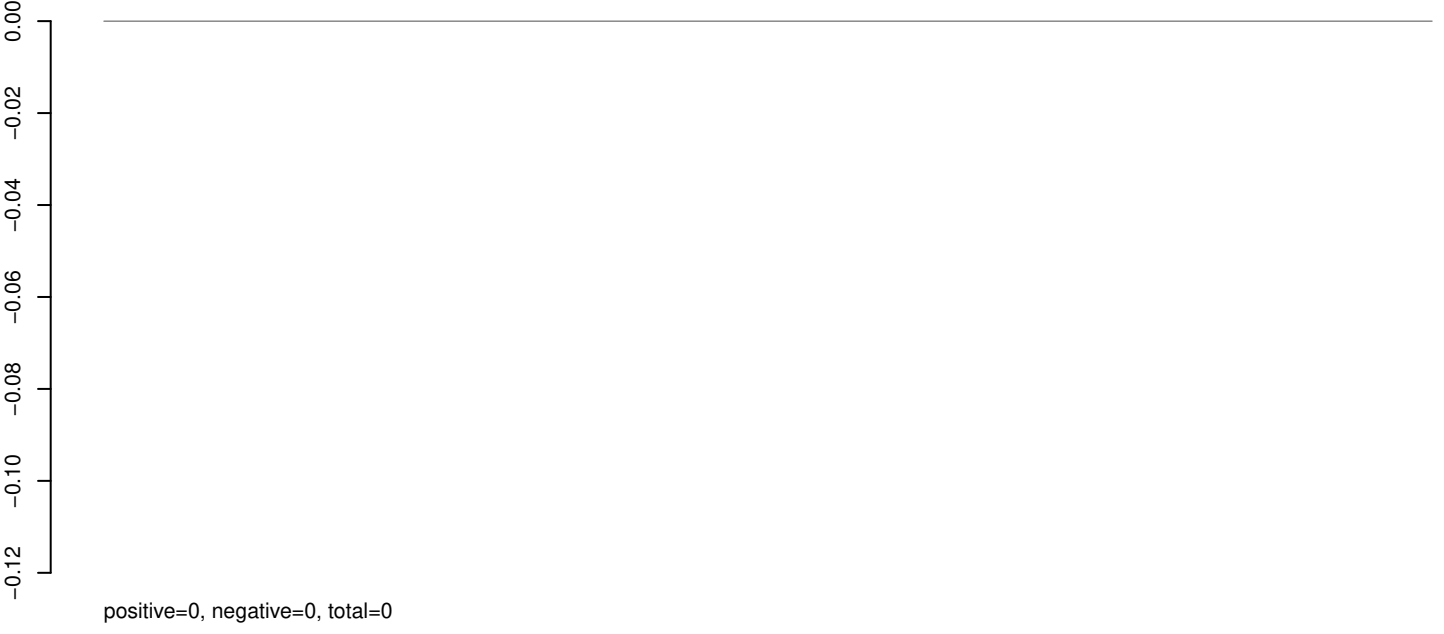




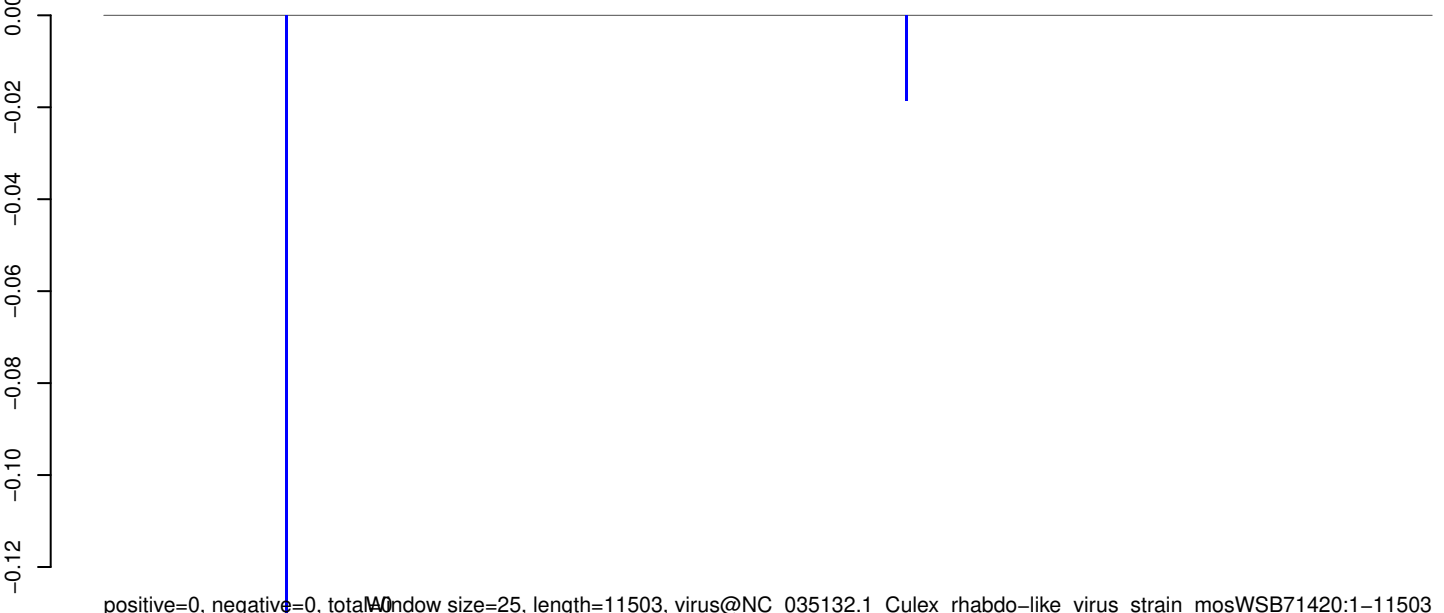
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



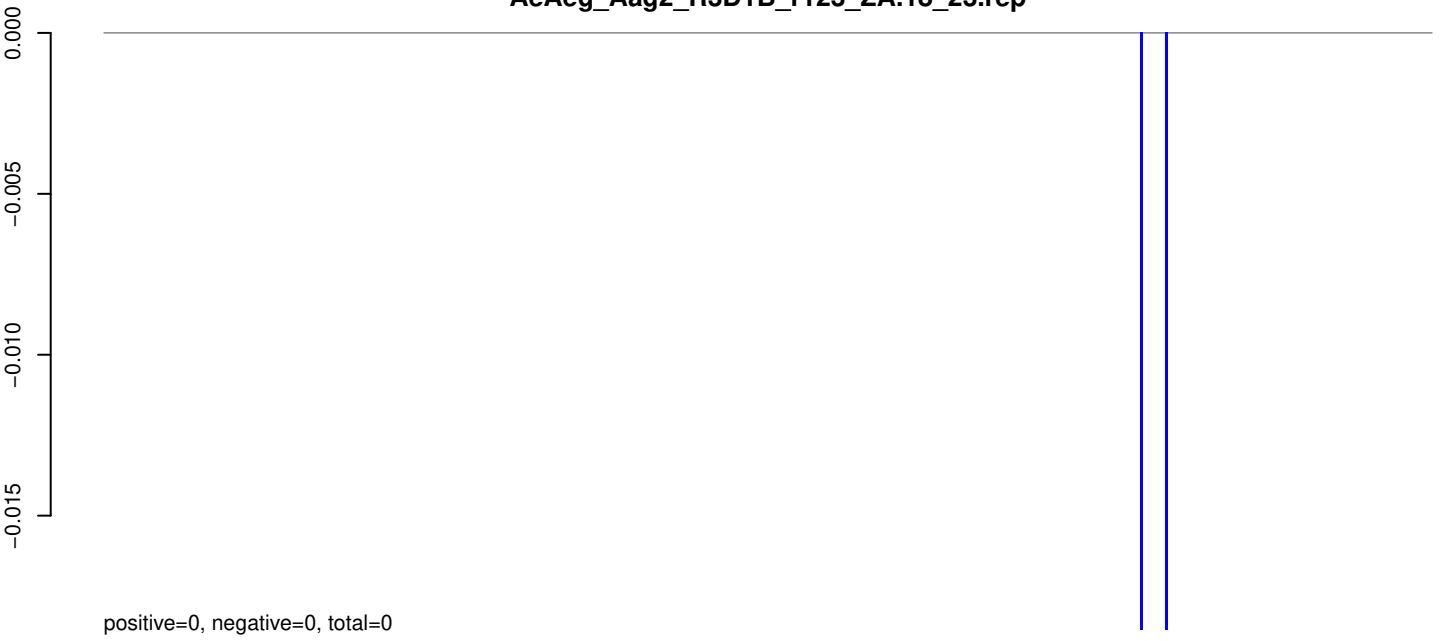
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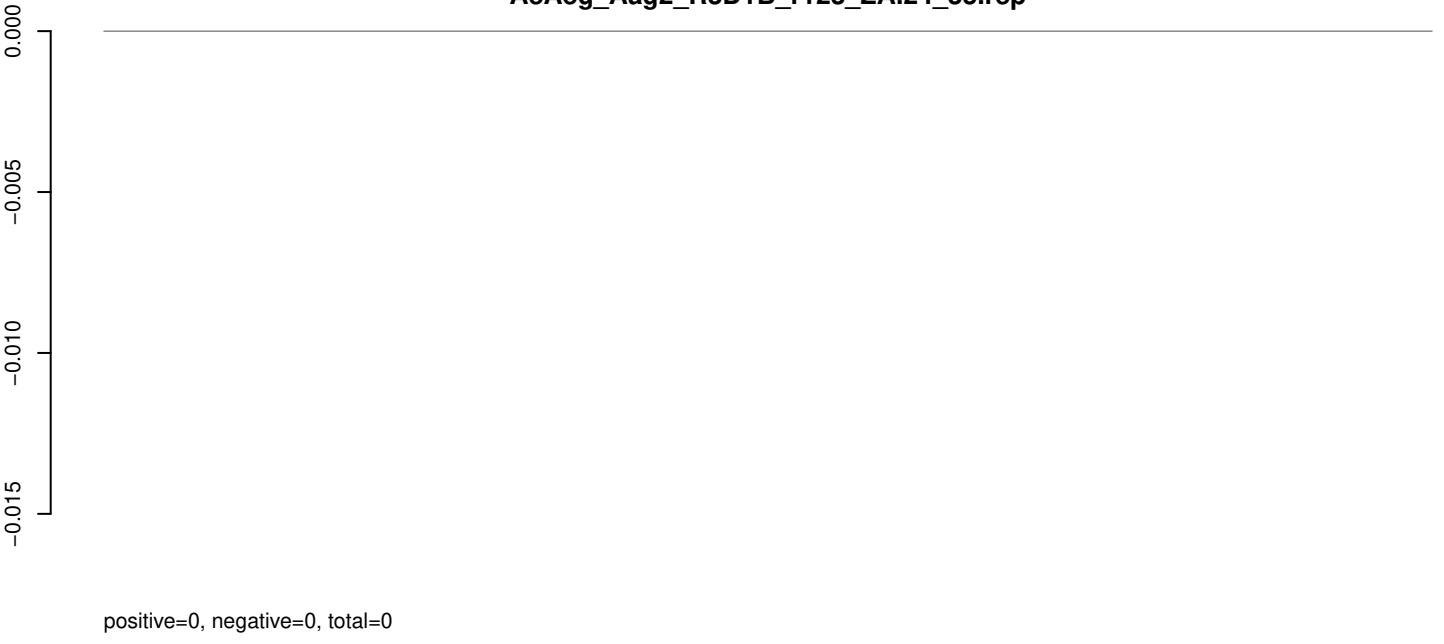
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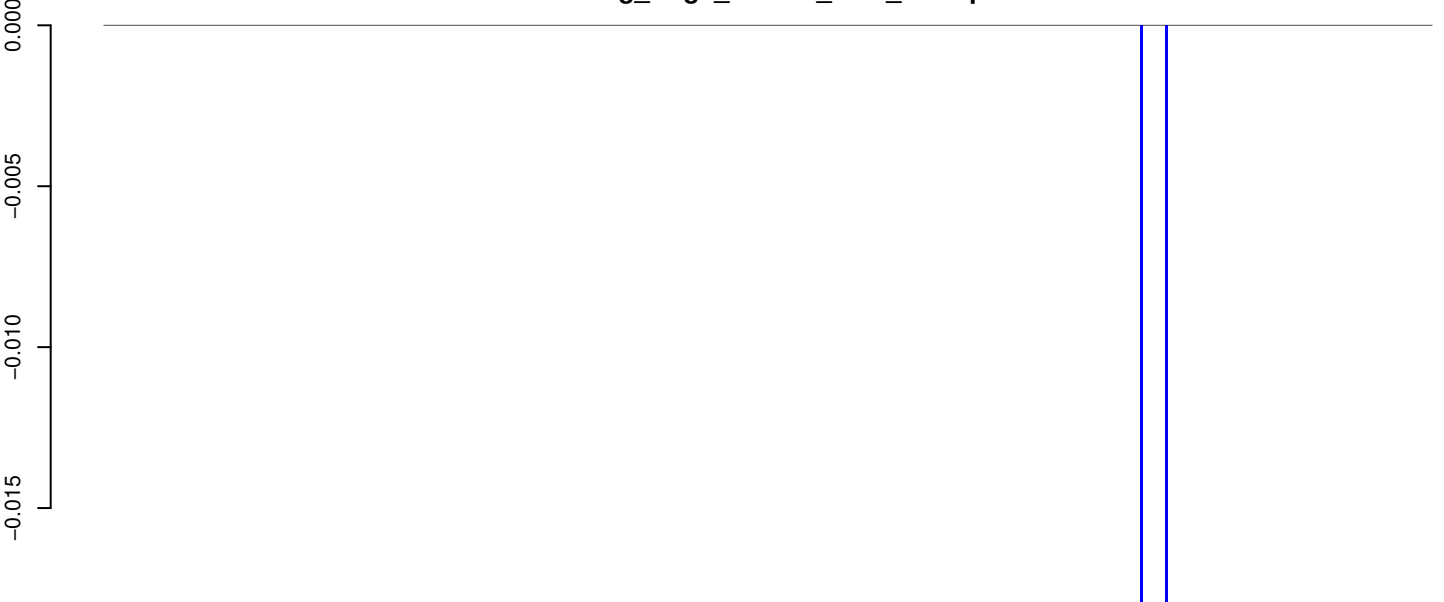
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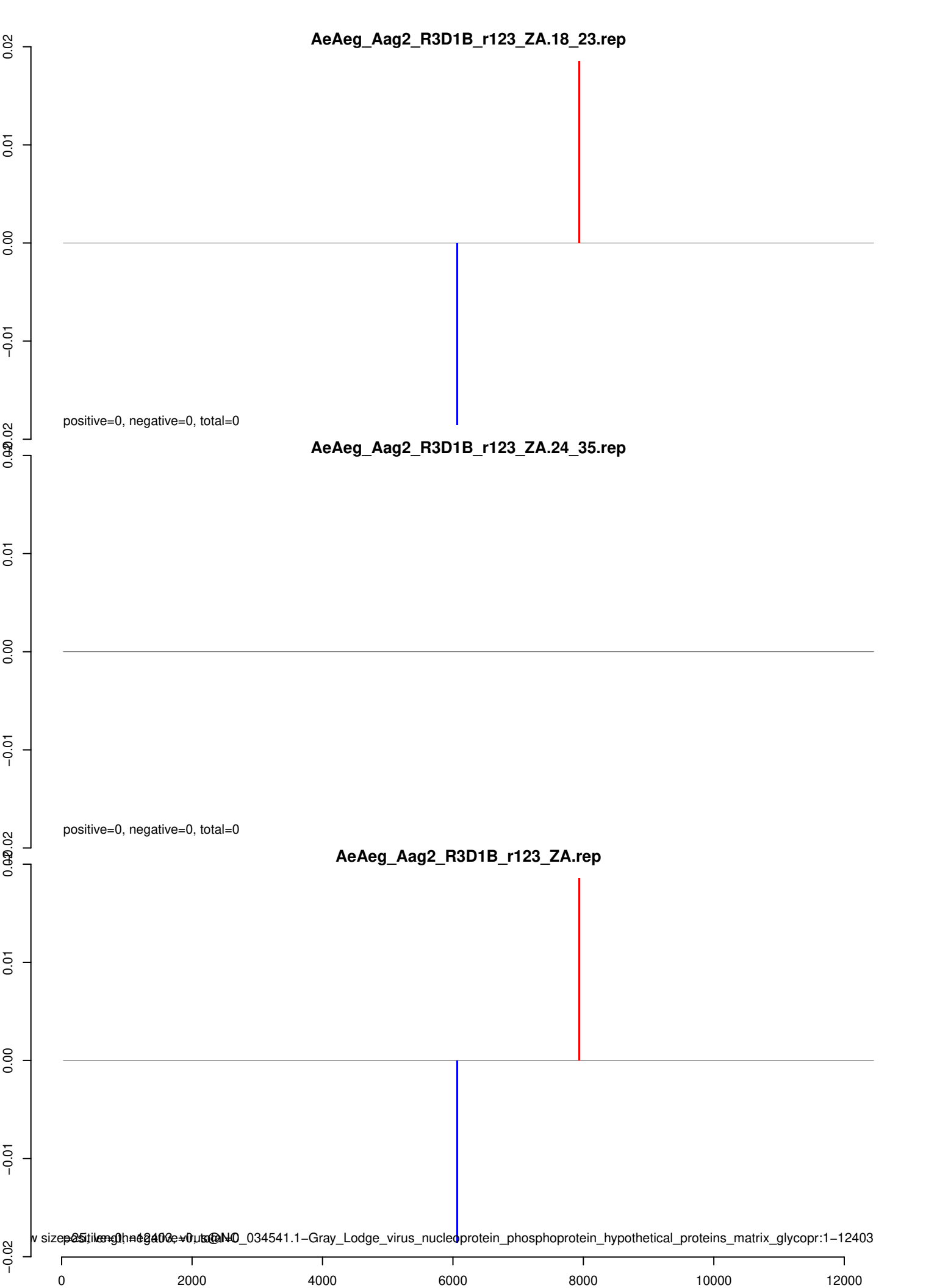


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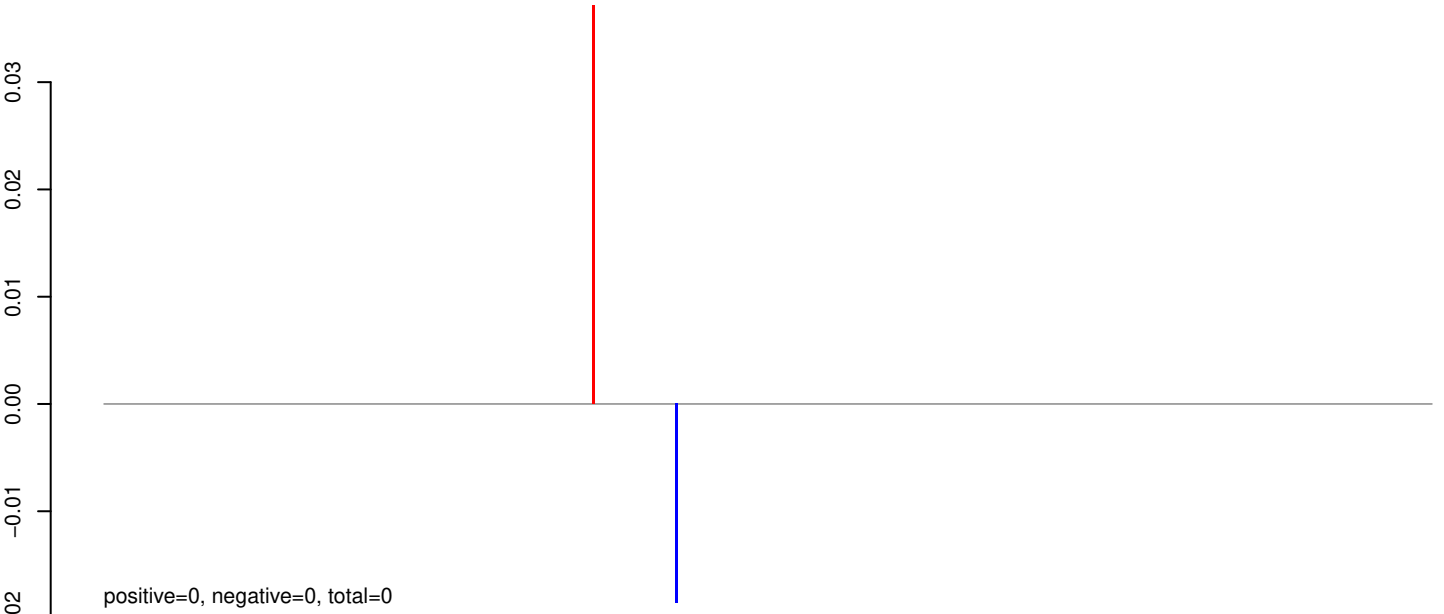


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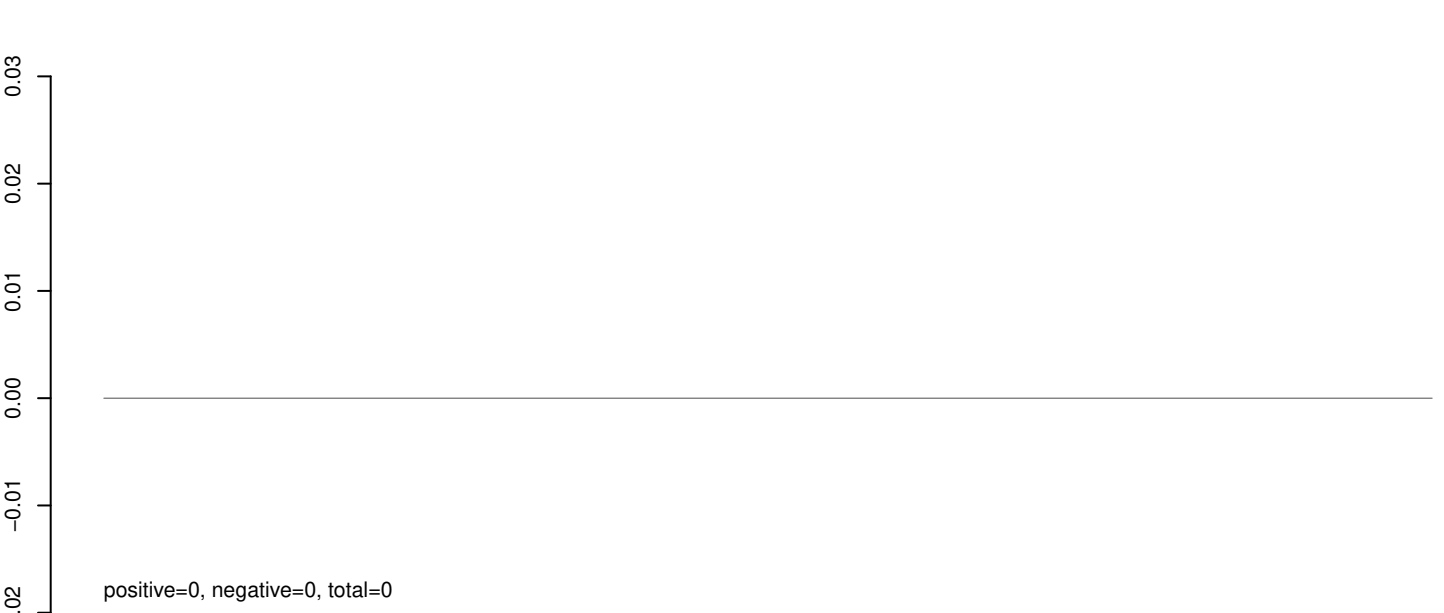




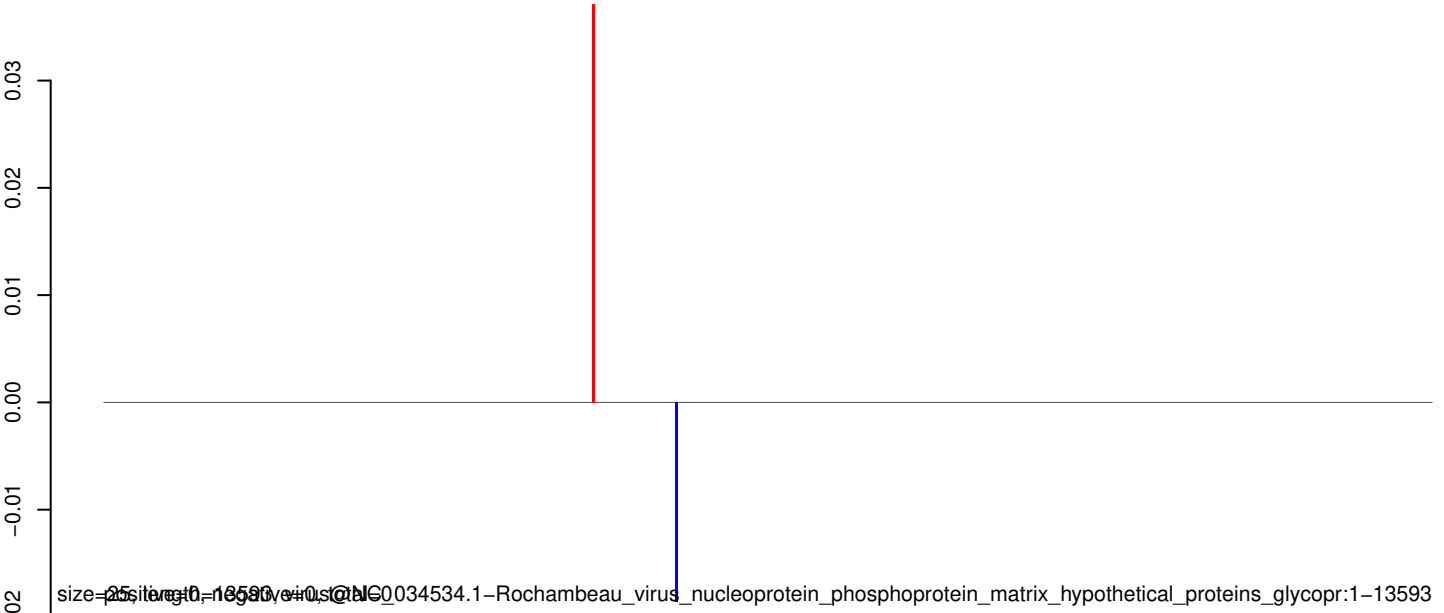
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep

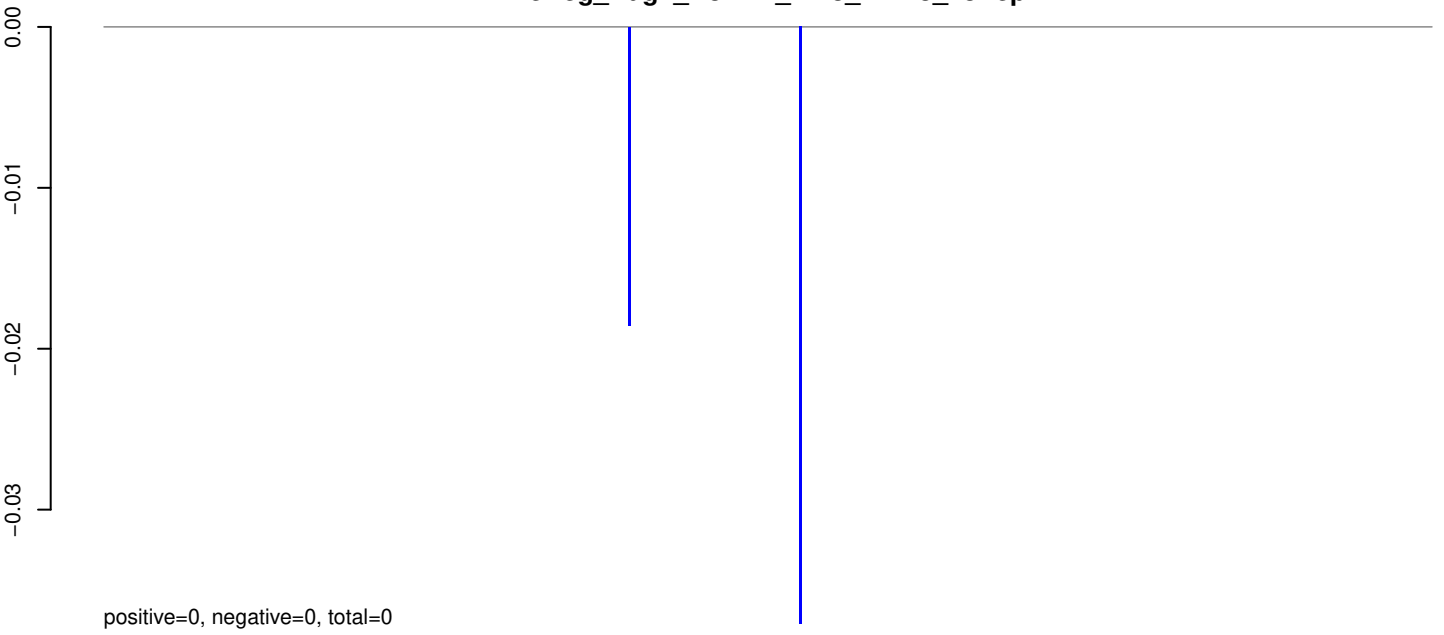


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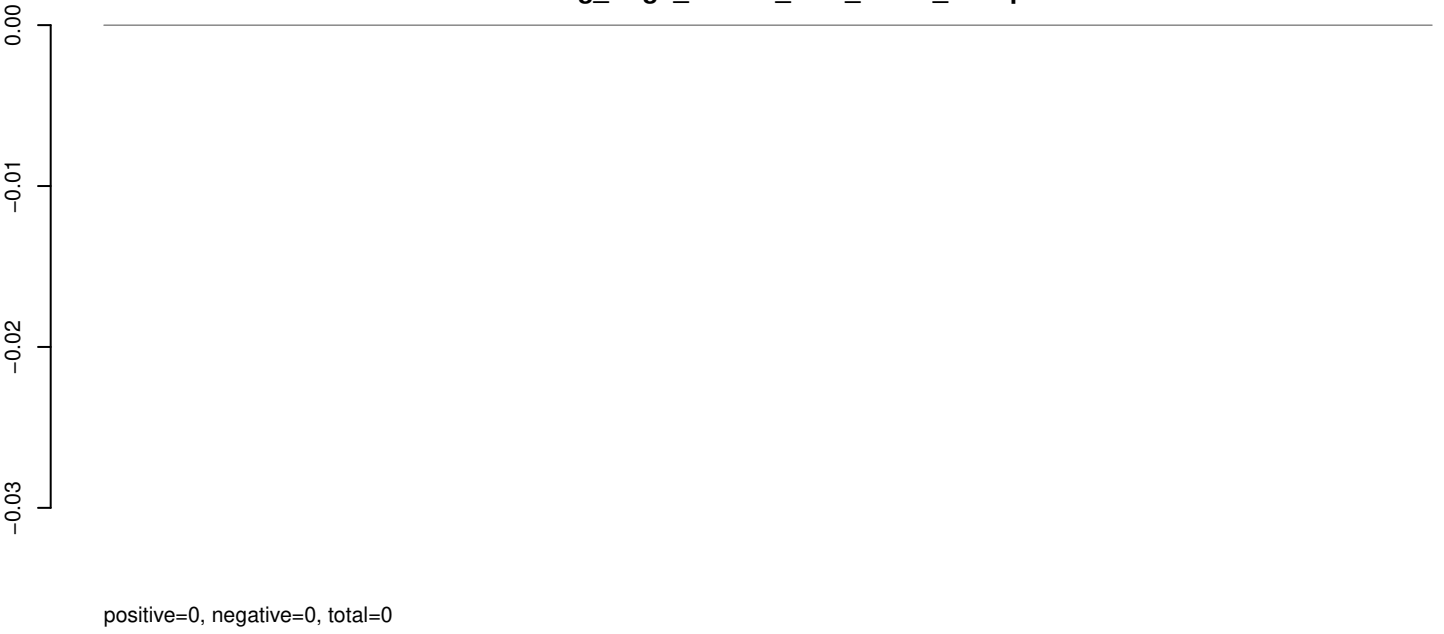


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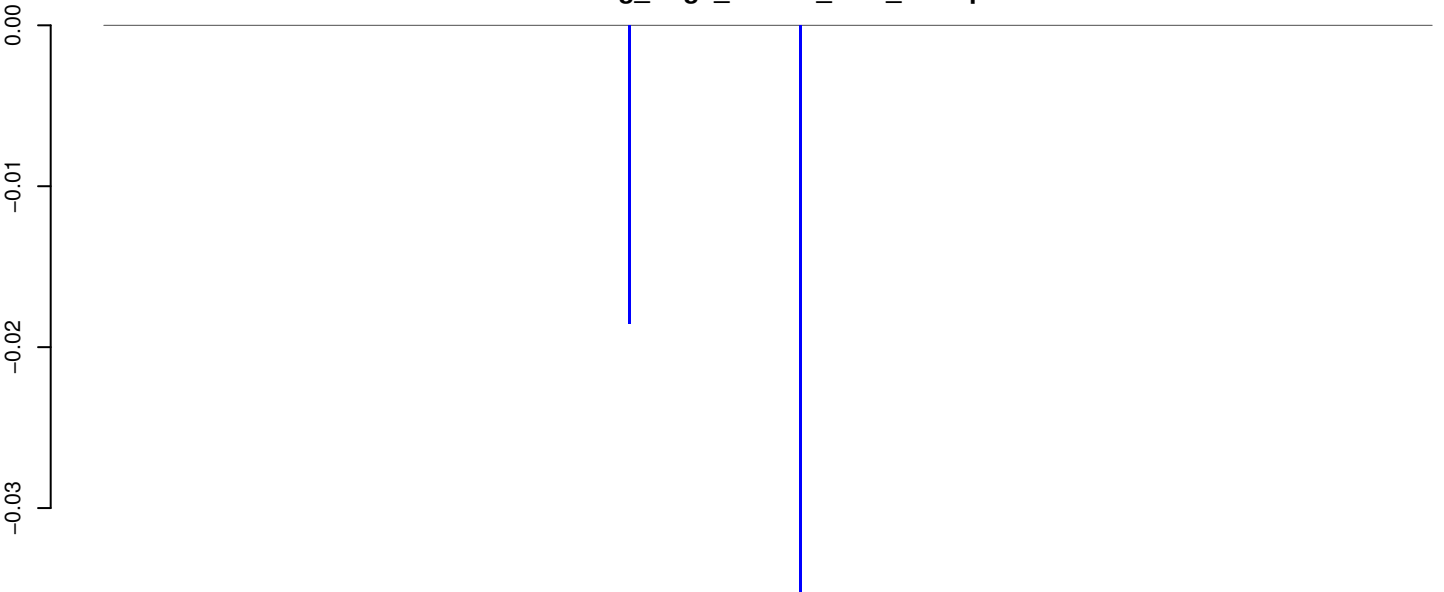
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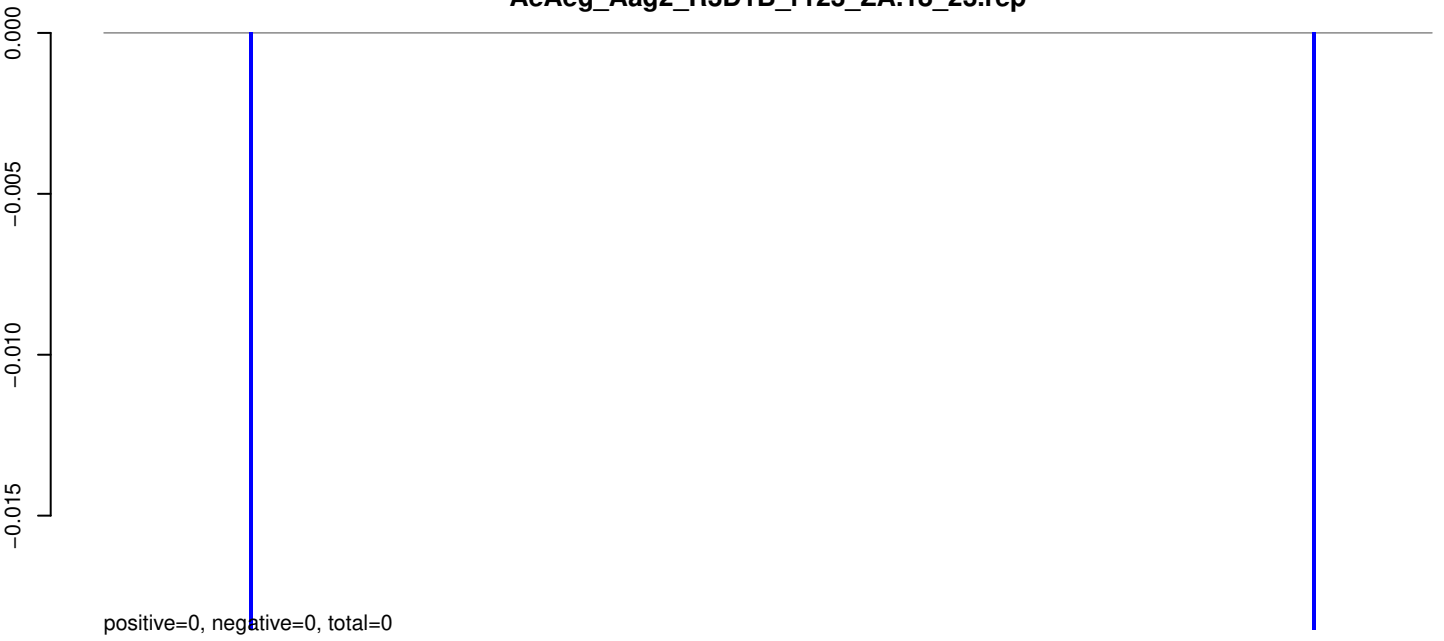
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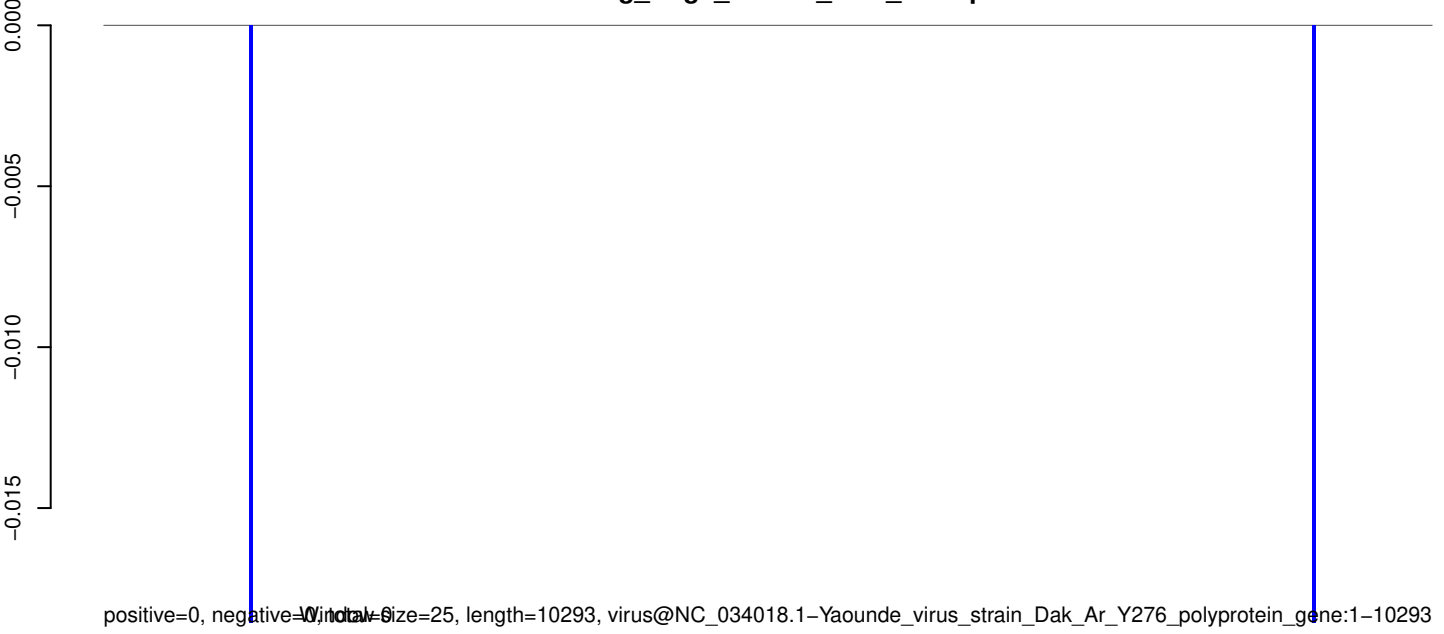
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



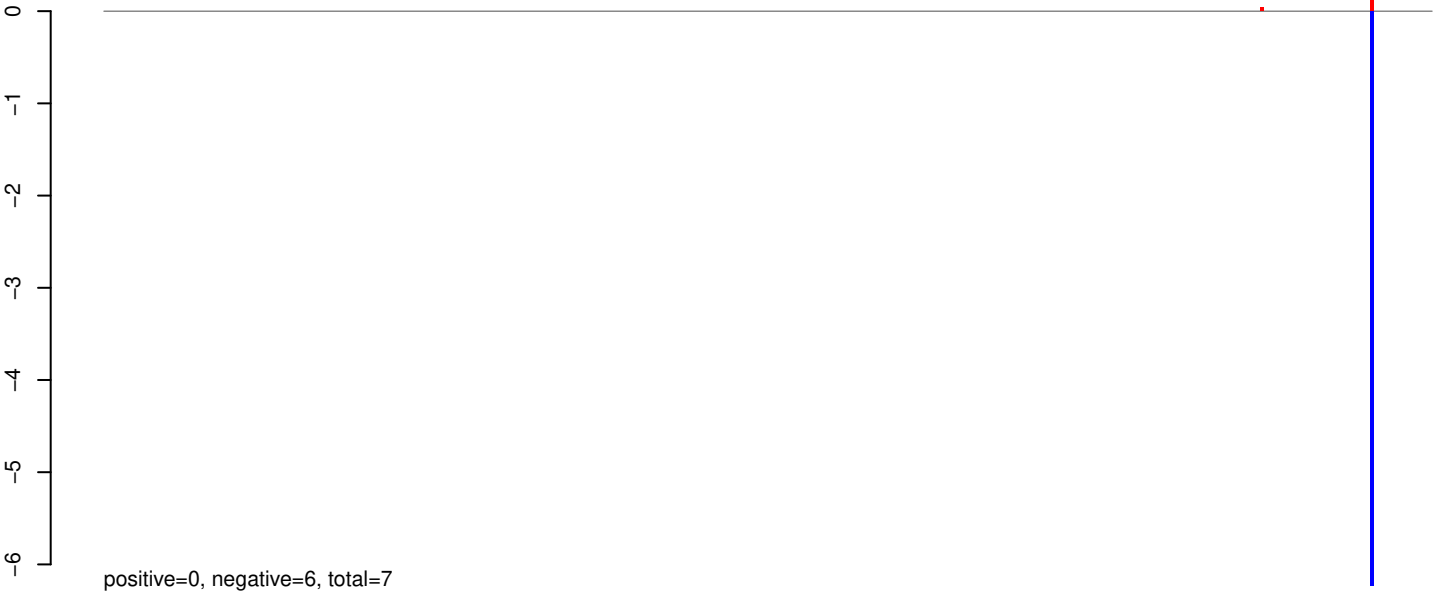
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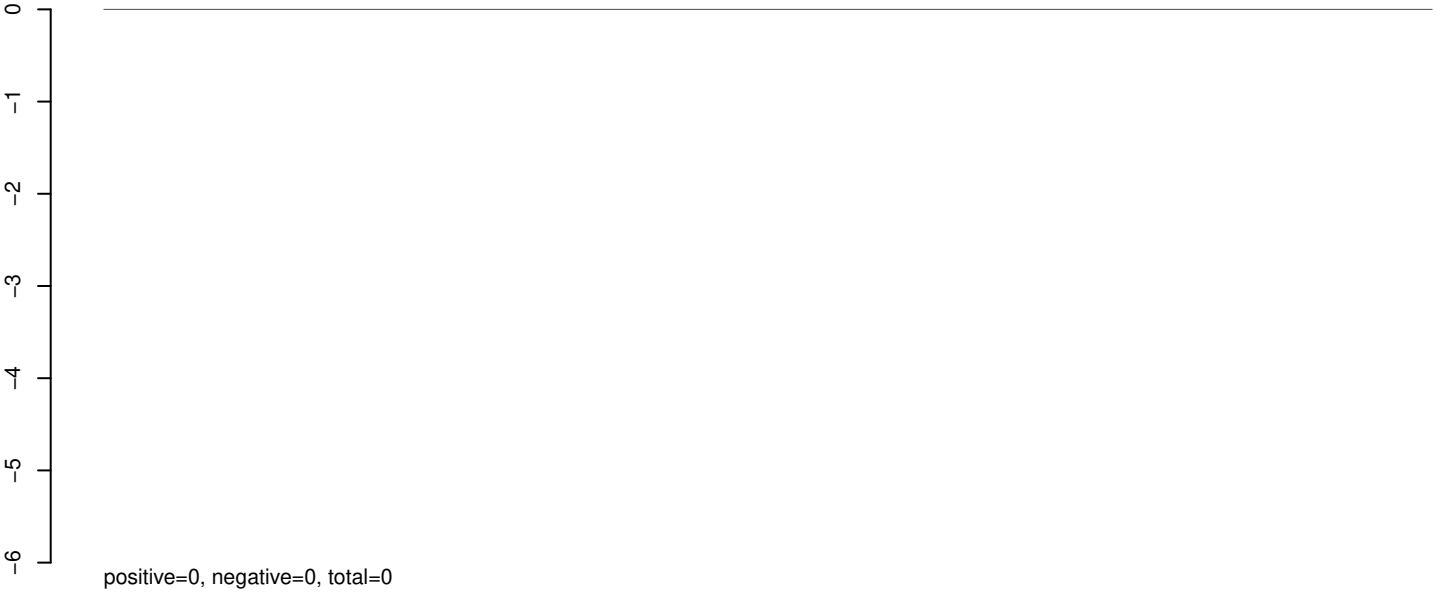
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



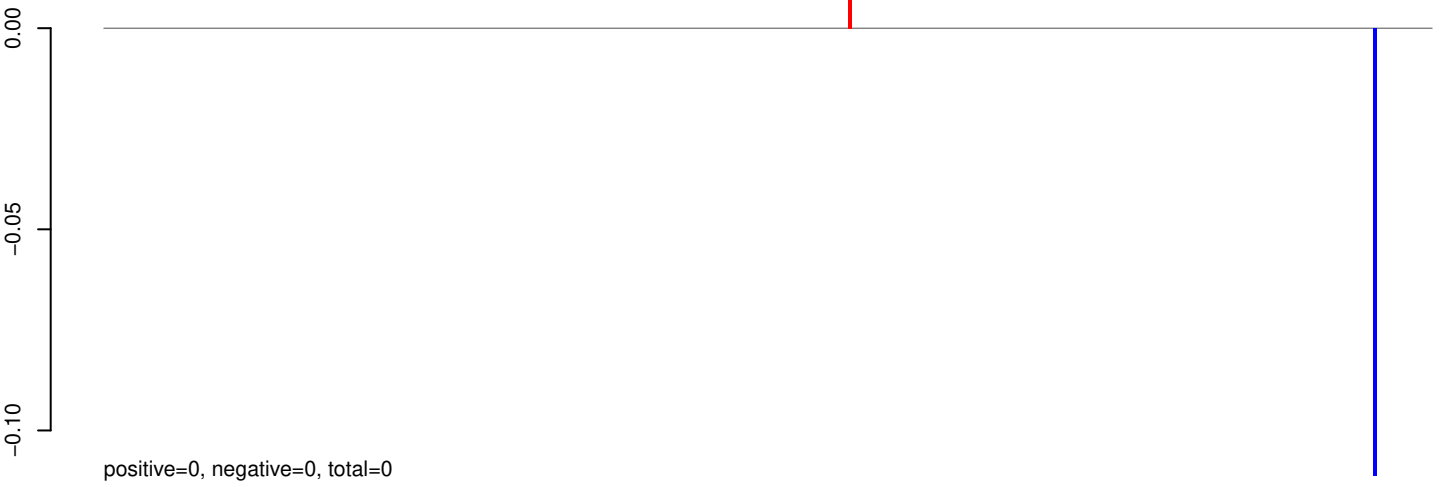
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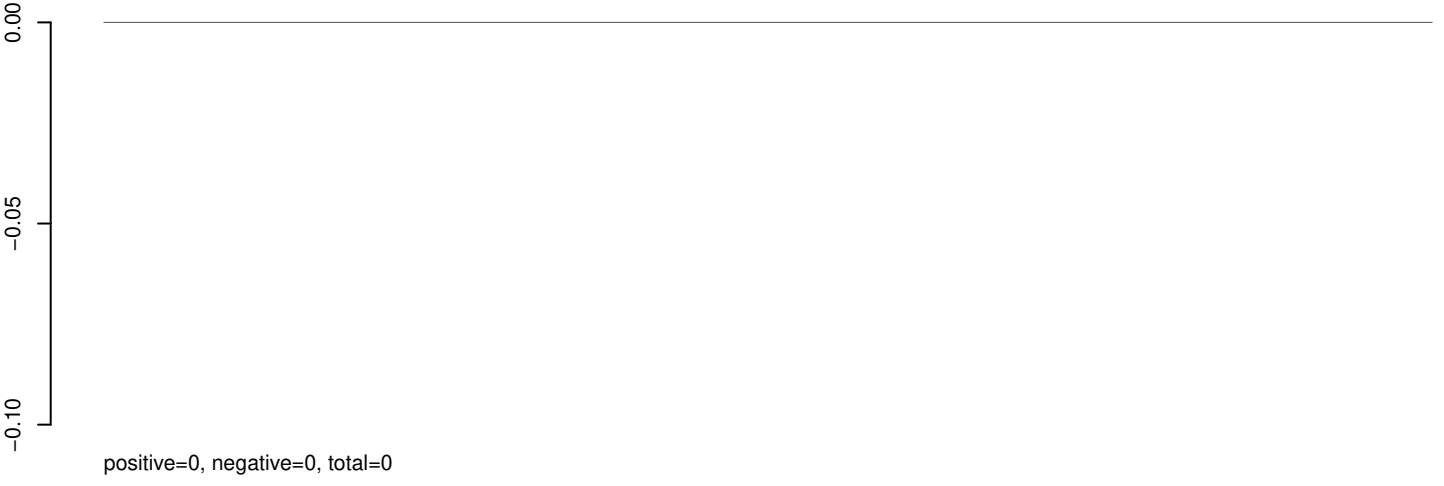
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0 2000 4000 6000 8000 10000

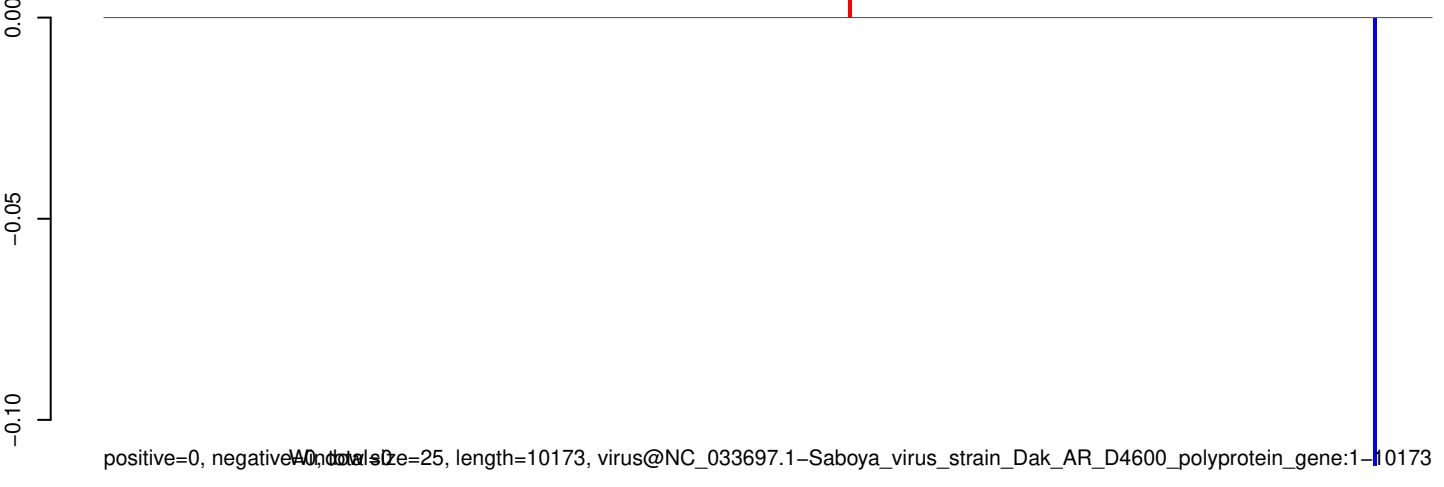
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



AeAeg_Aag2_R3D1B_r123_ZA.rep

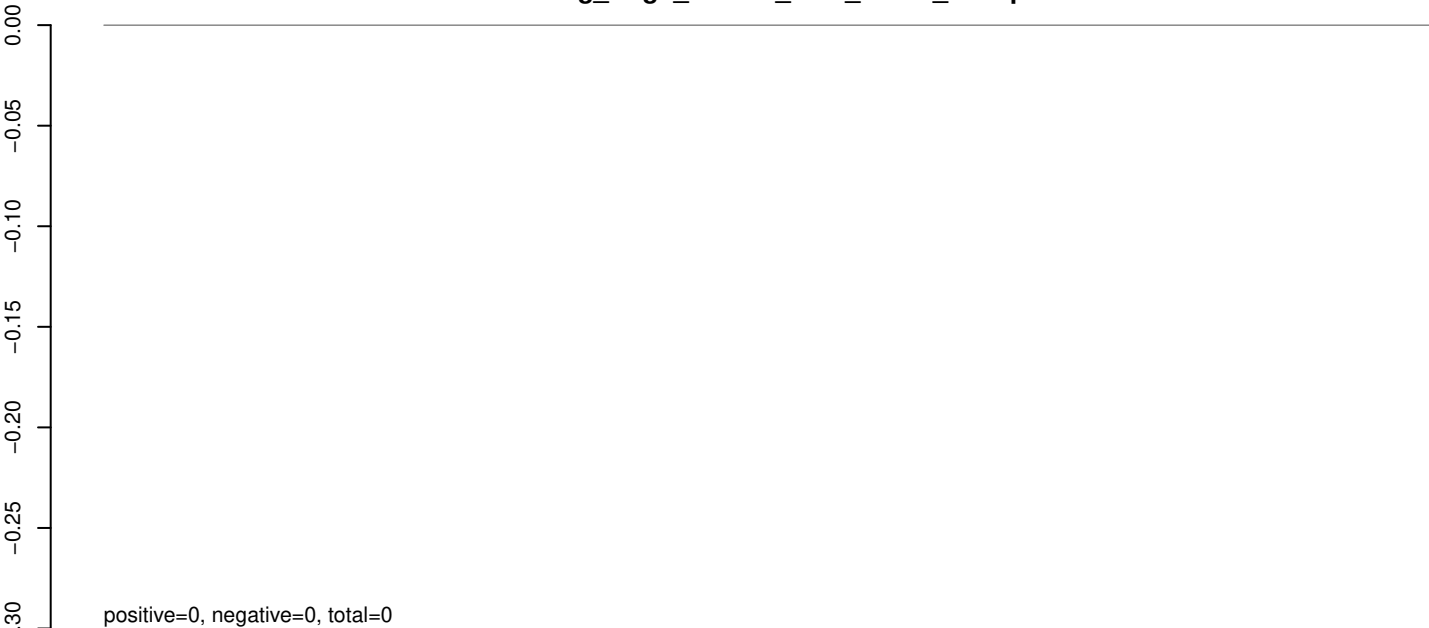


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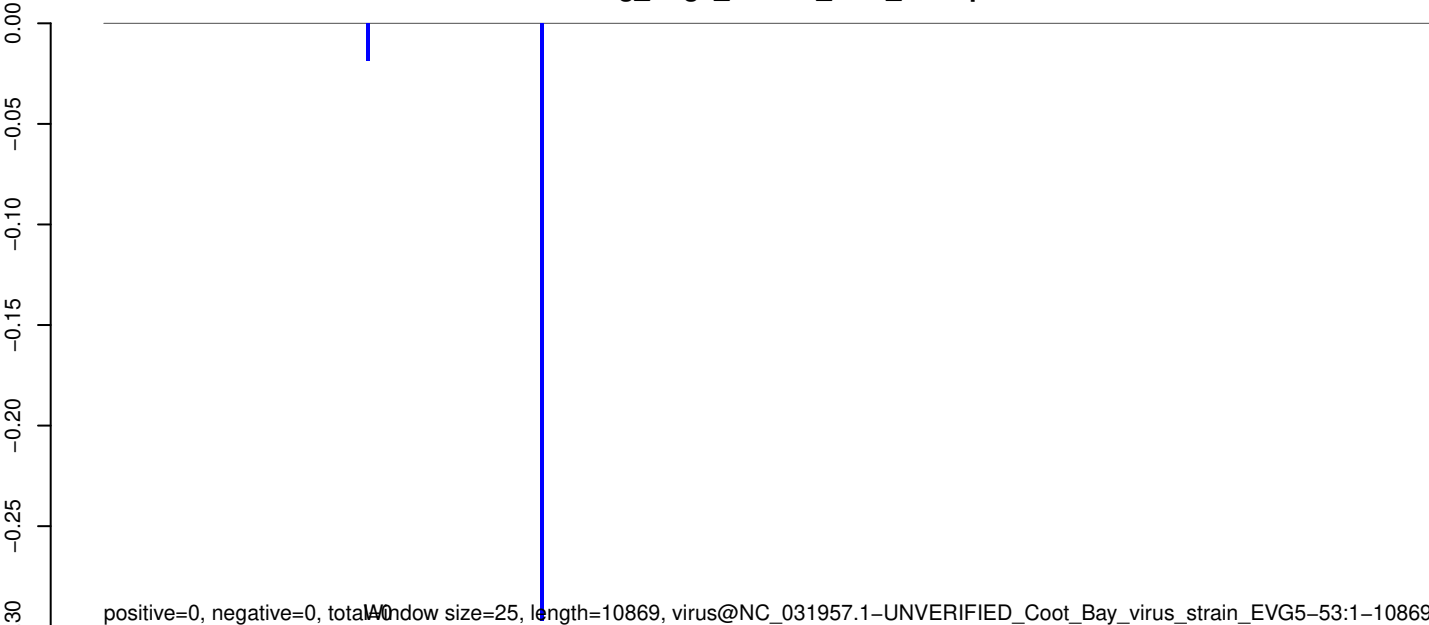
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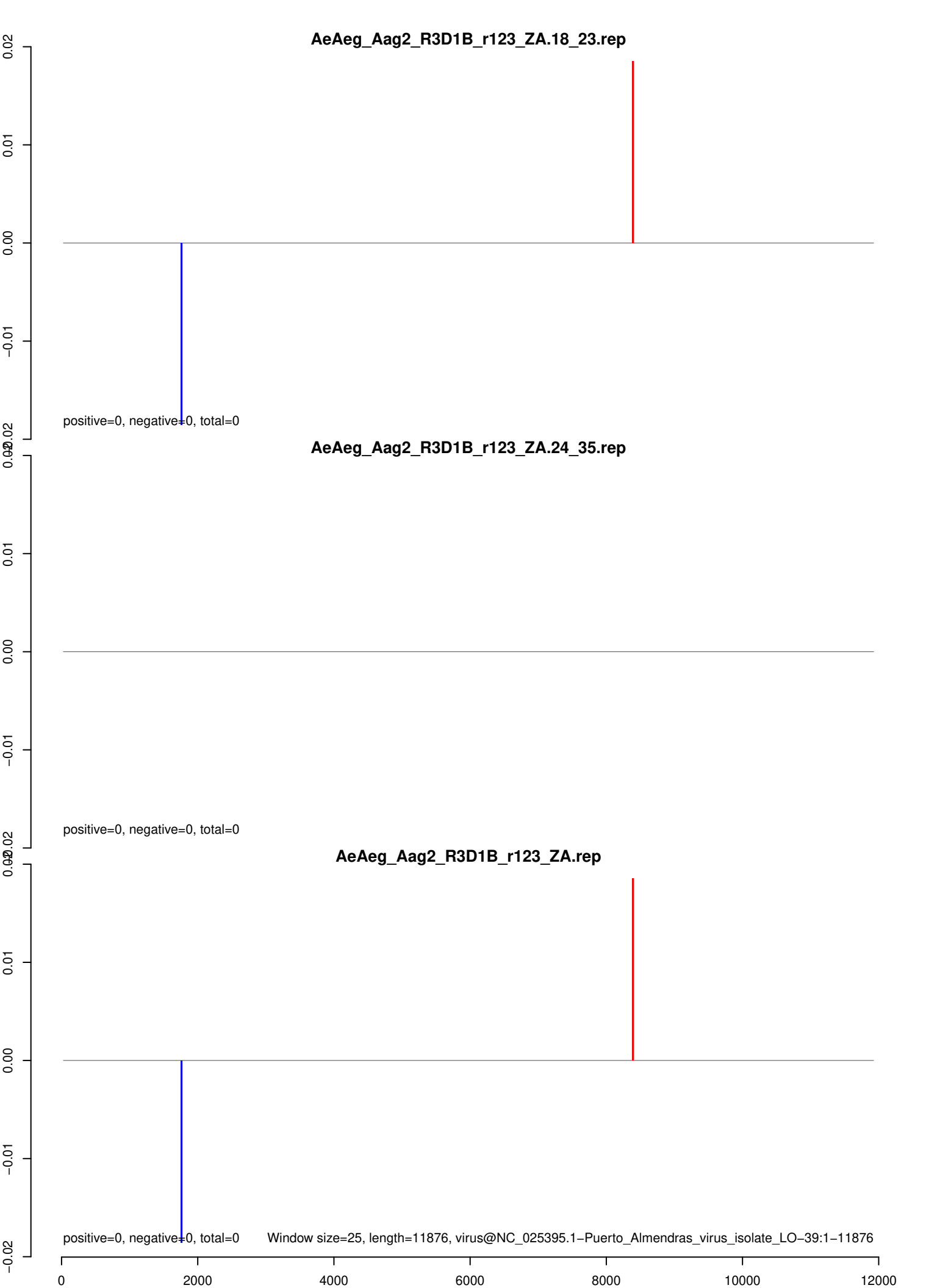


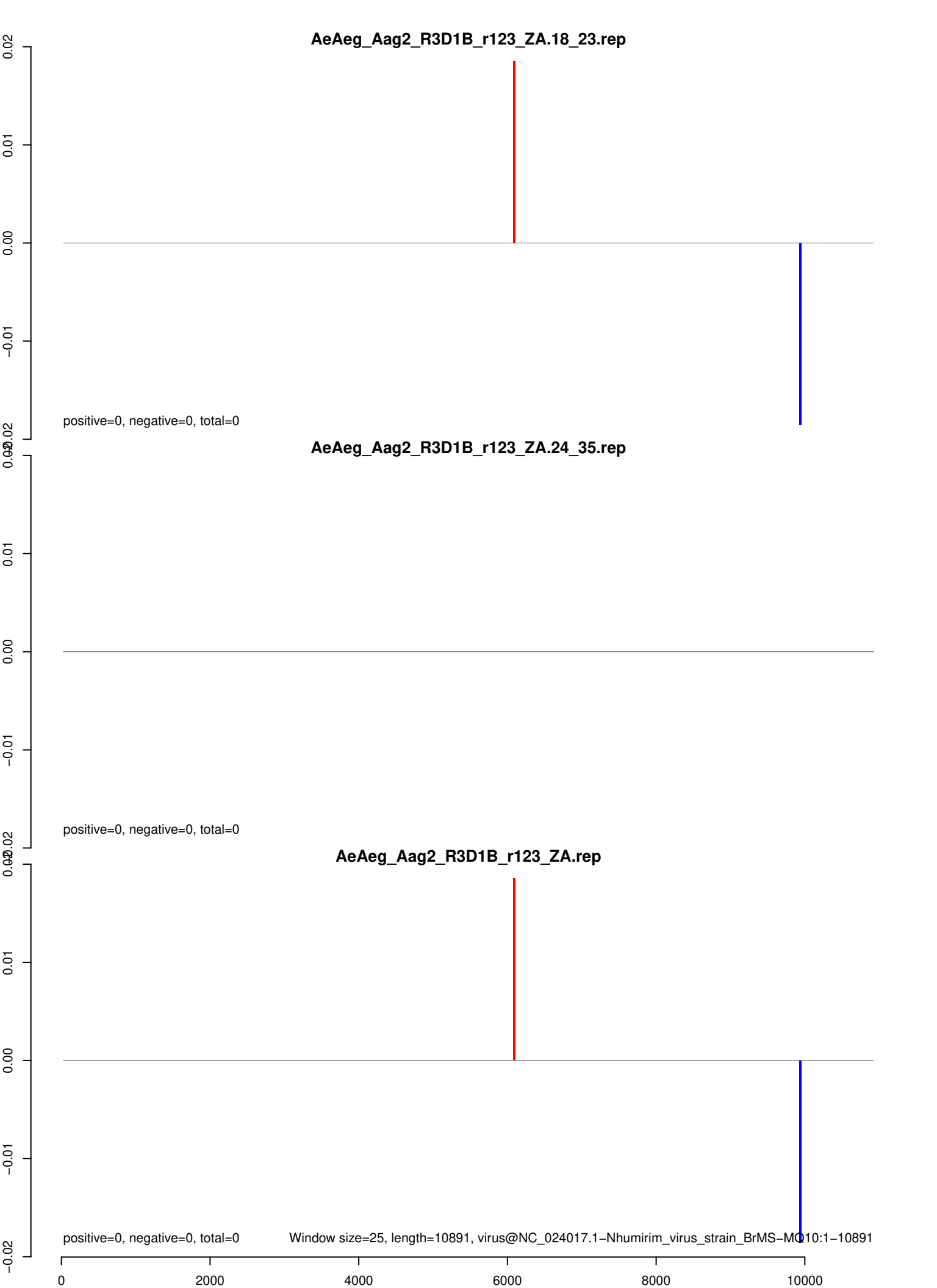
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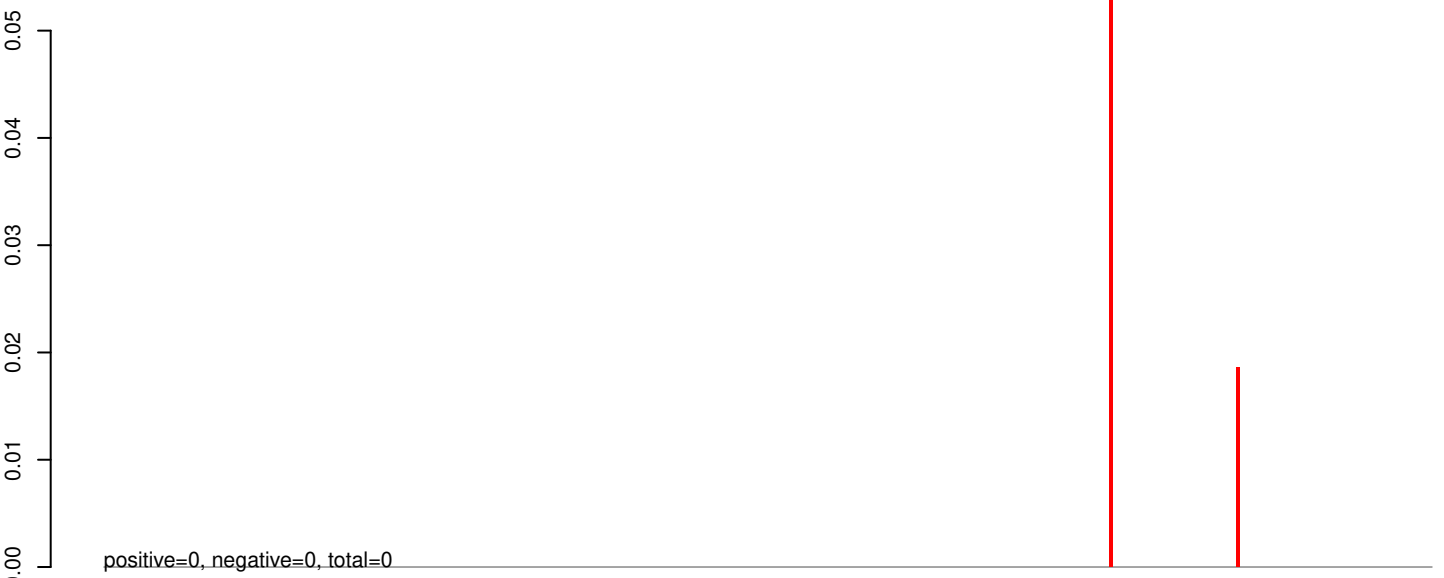
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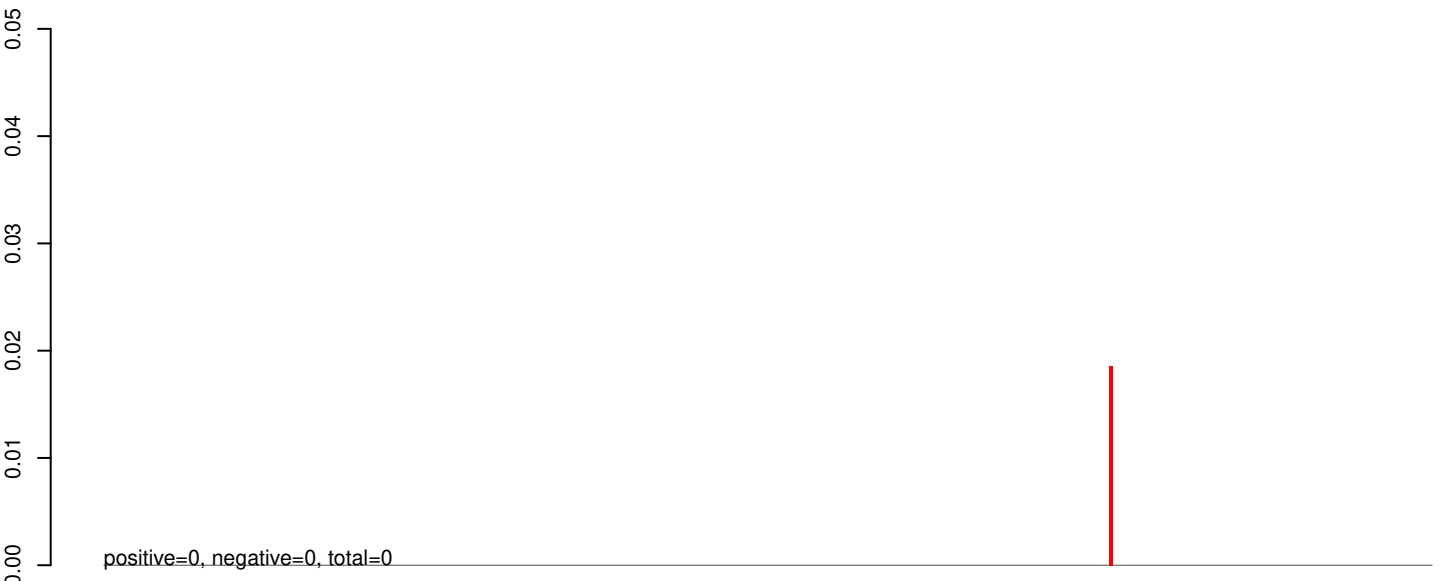




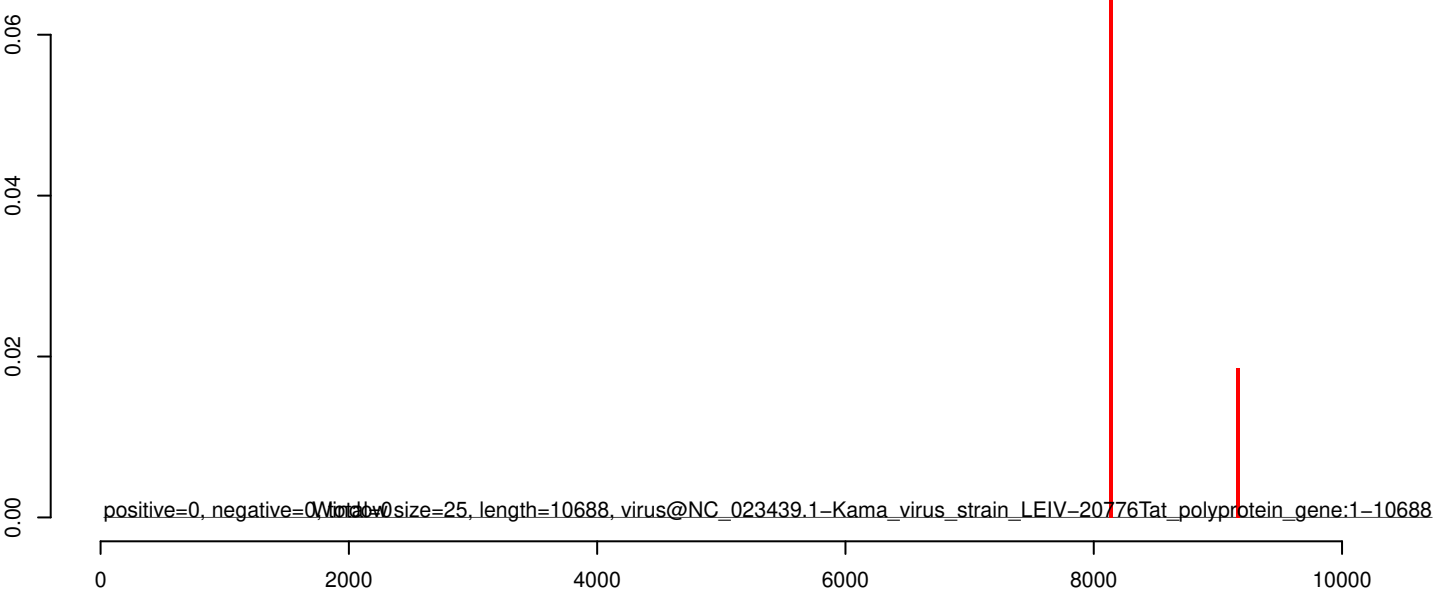
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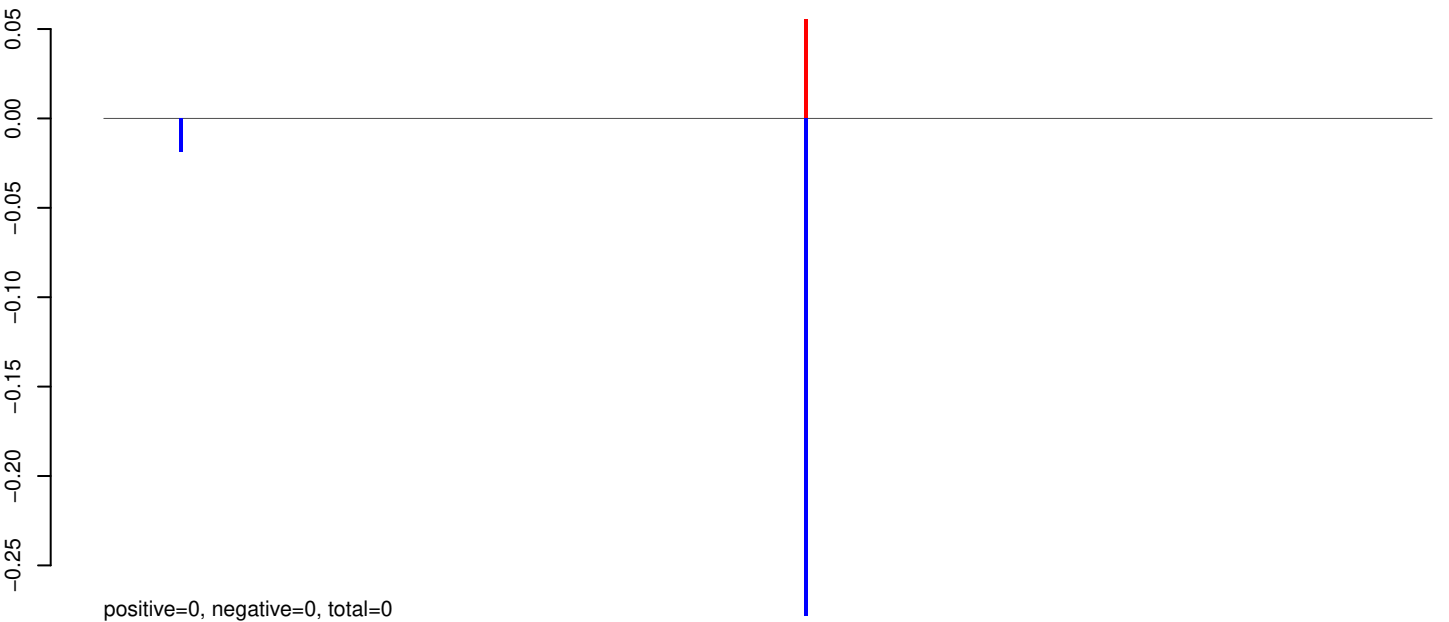
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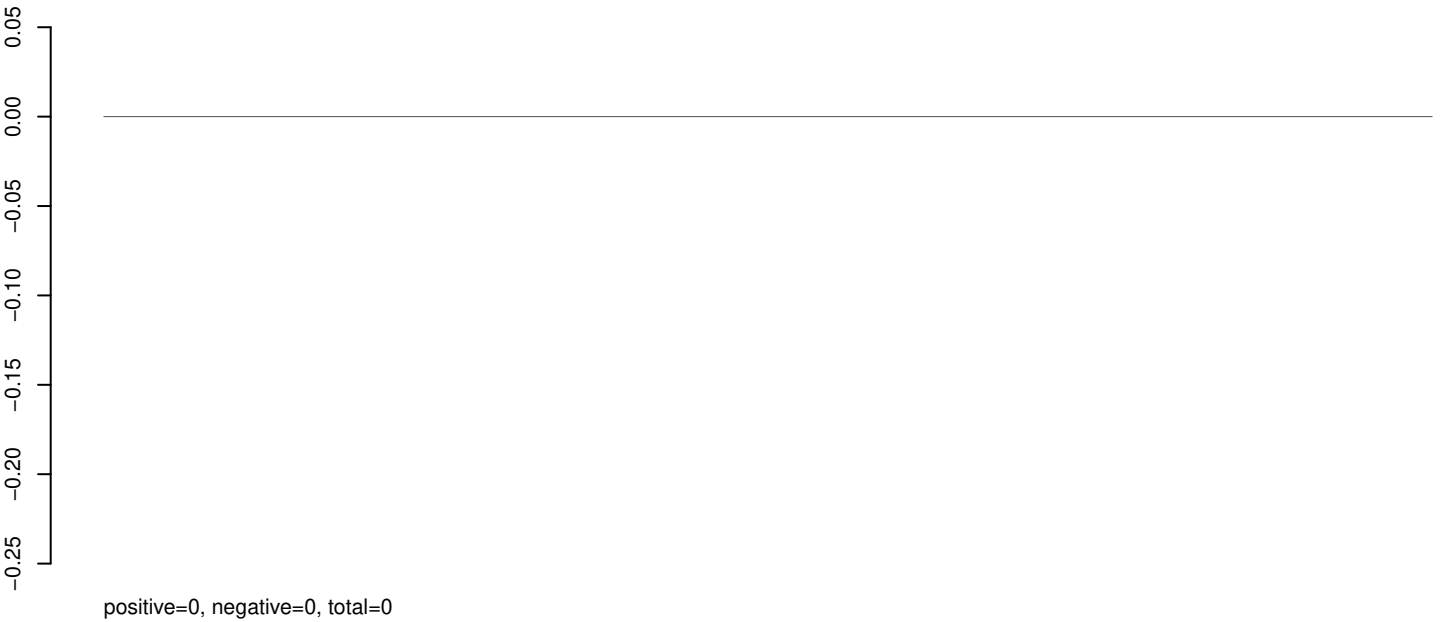
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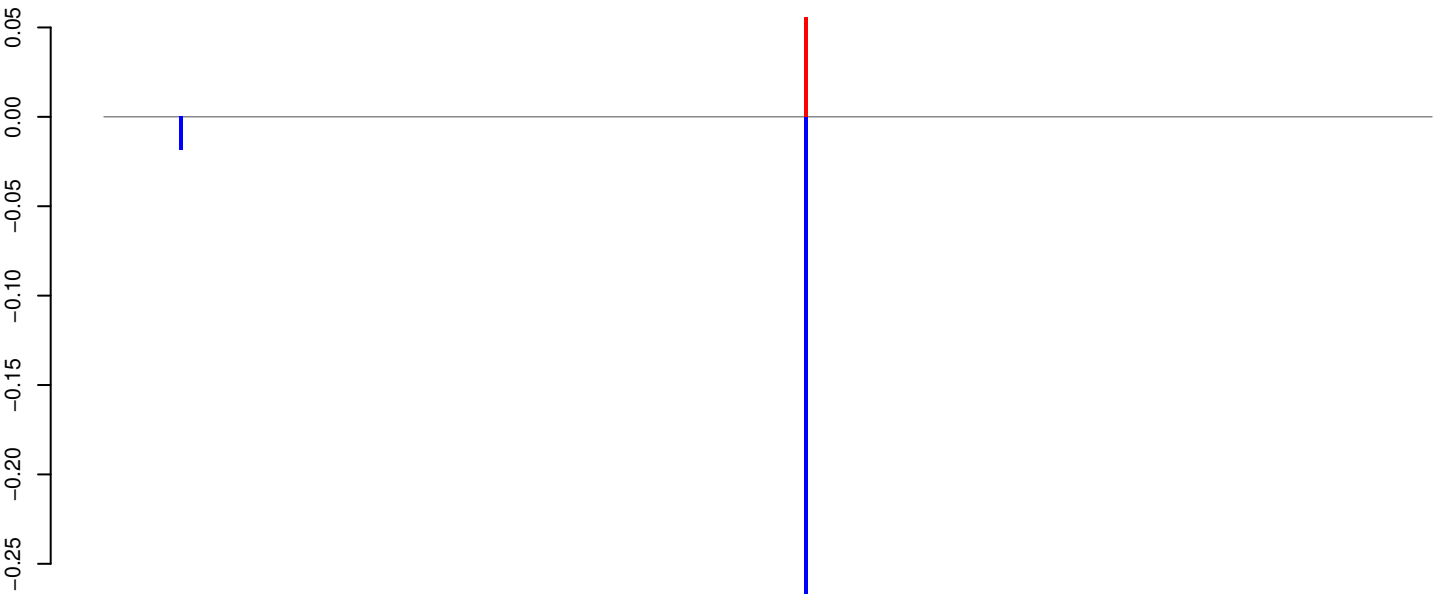
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



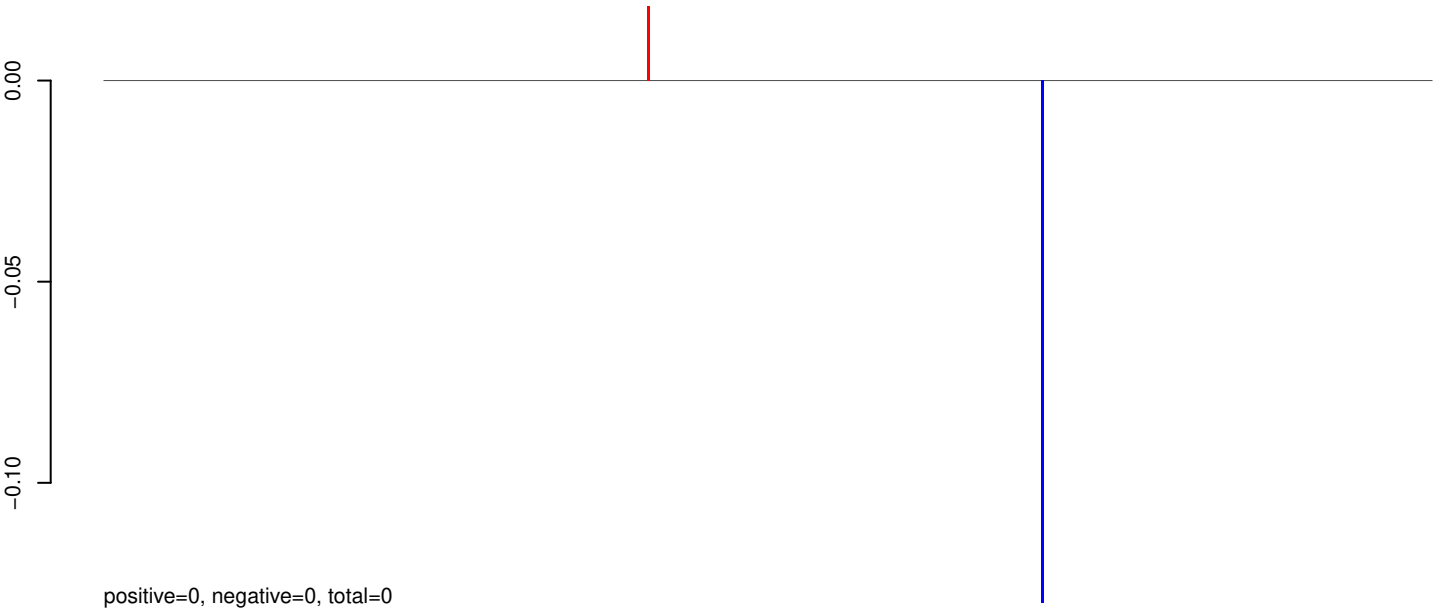
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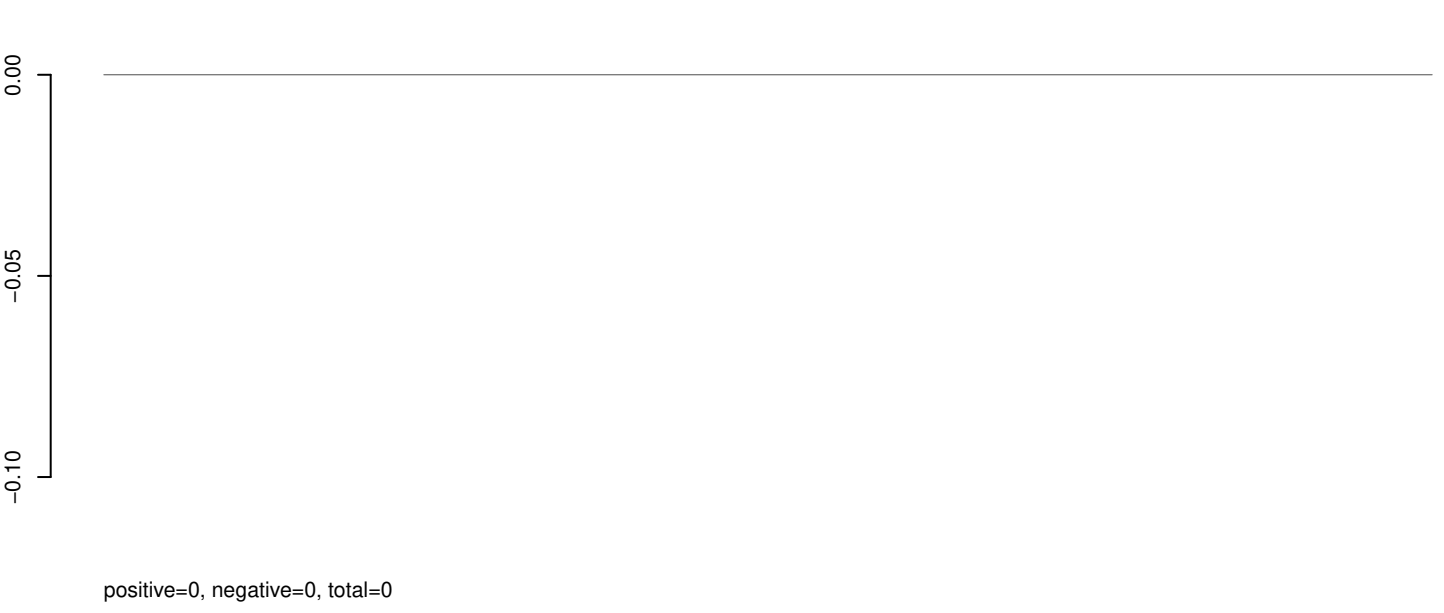
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0 2000 4000 6000 8000 10000

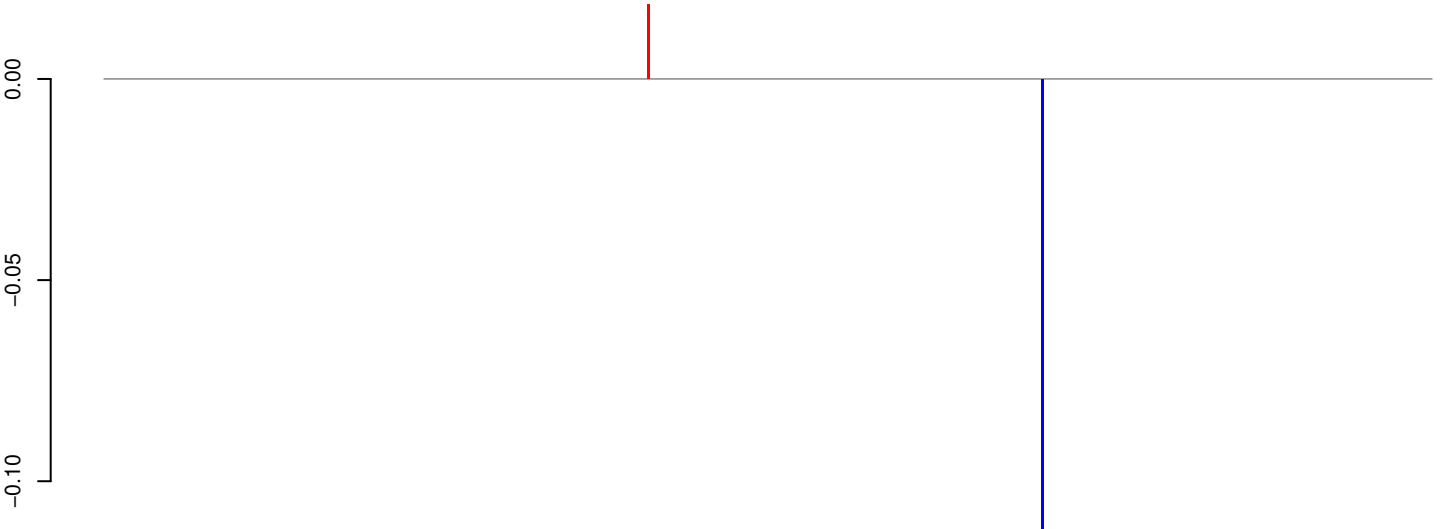
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



AeAeg_Aag2_R3D1B_r123_ZA.rep



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

0 2000 4000 6000 8000 10000 12000

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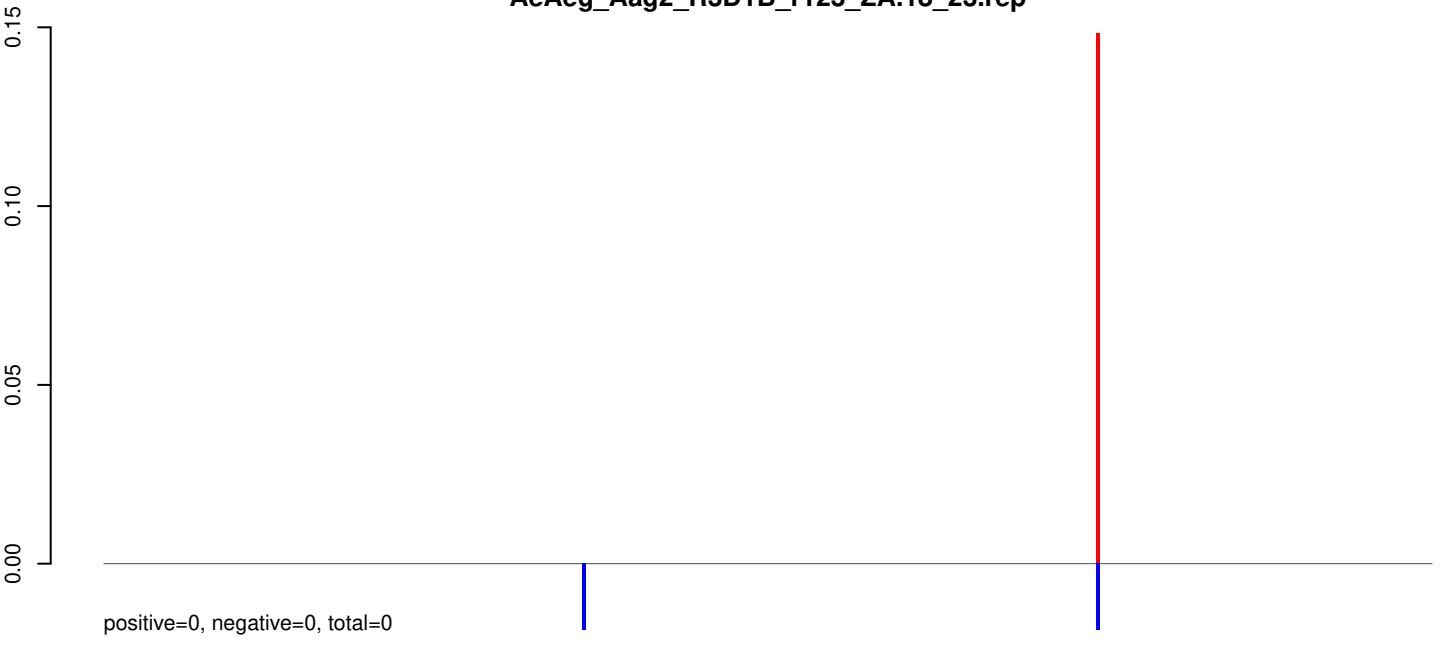
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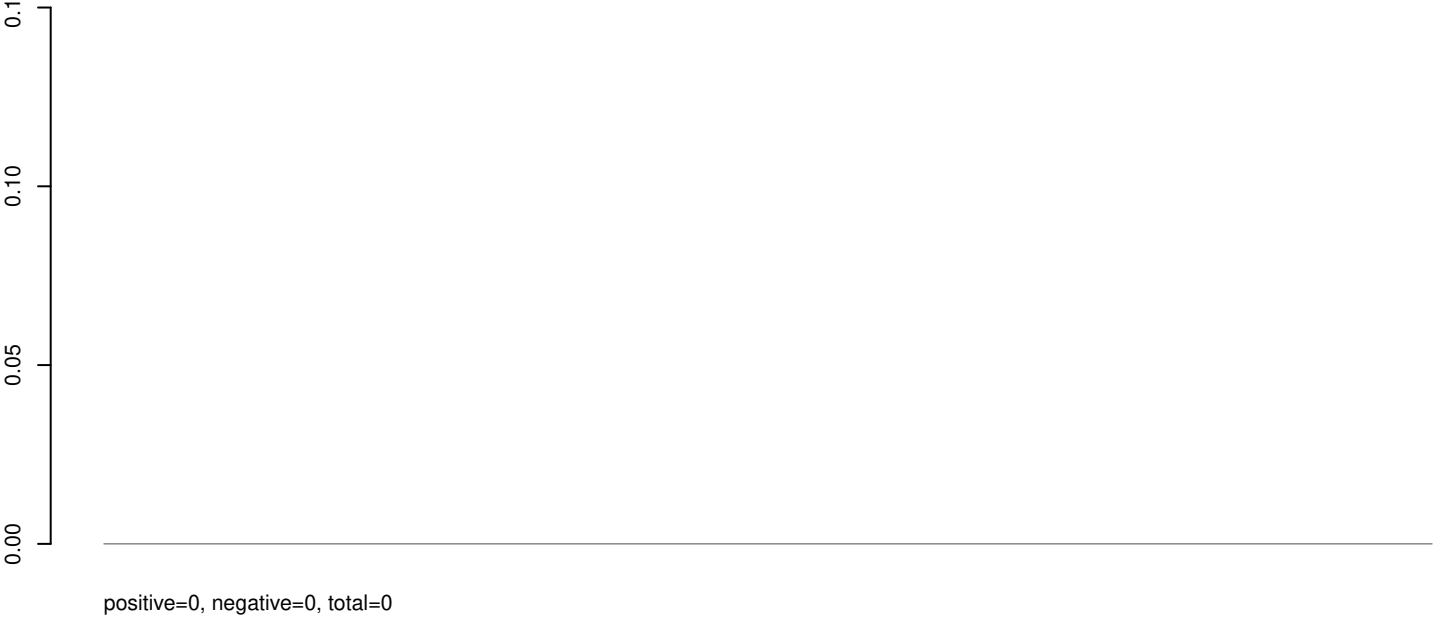
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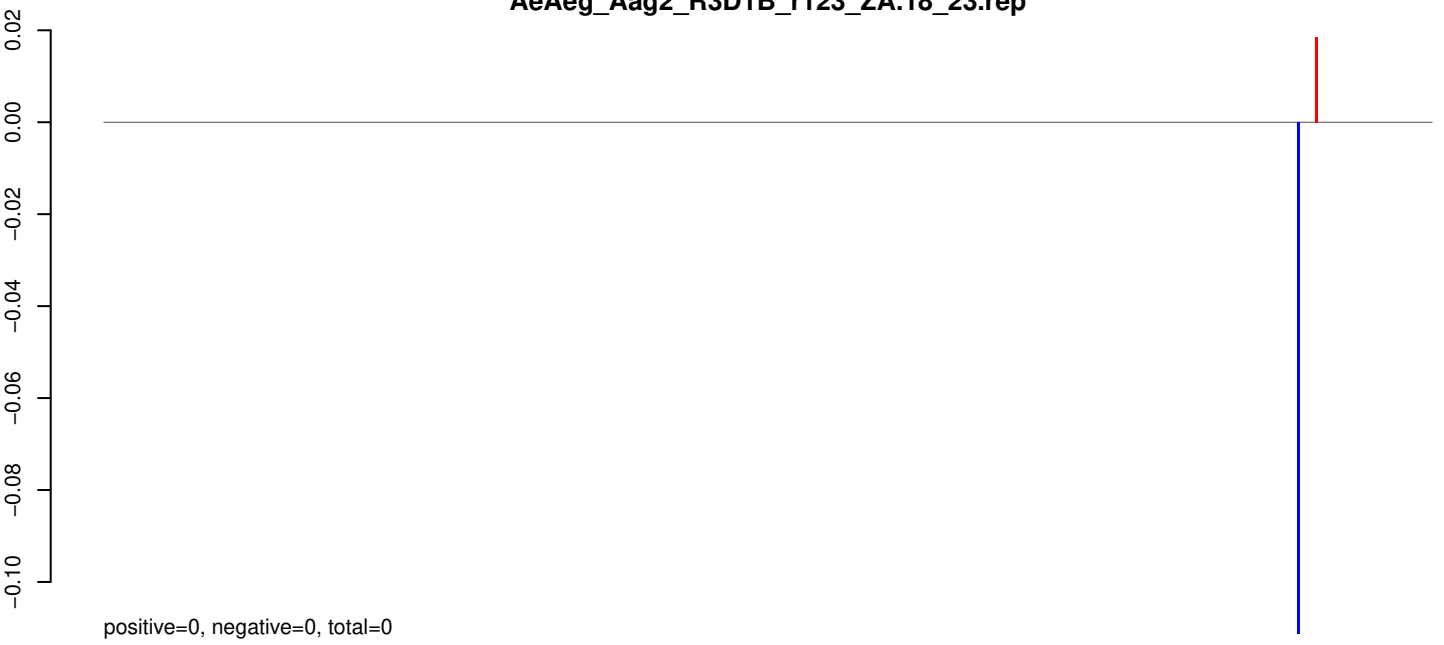
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AeAeg_Aag2_R3D1B_r123_ZA.rep

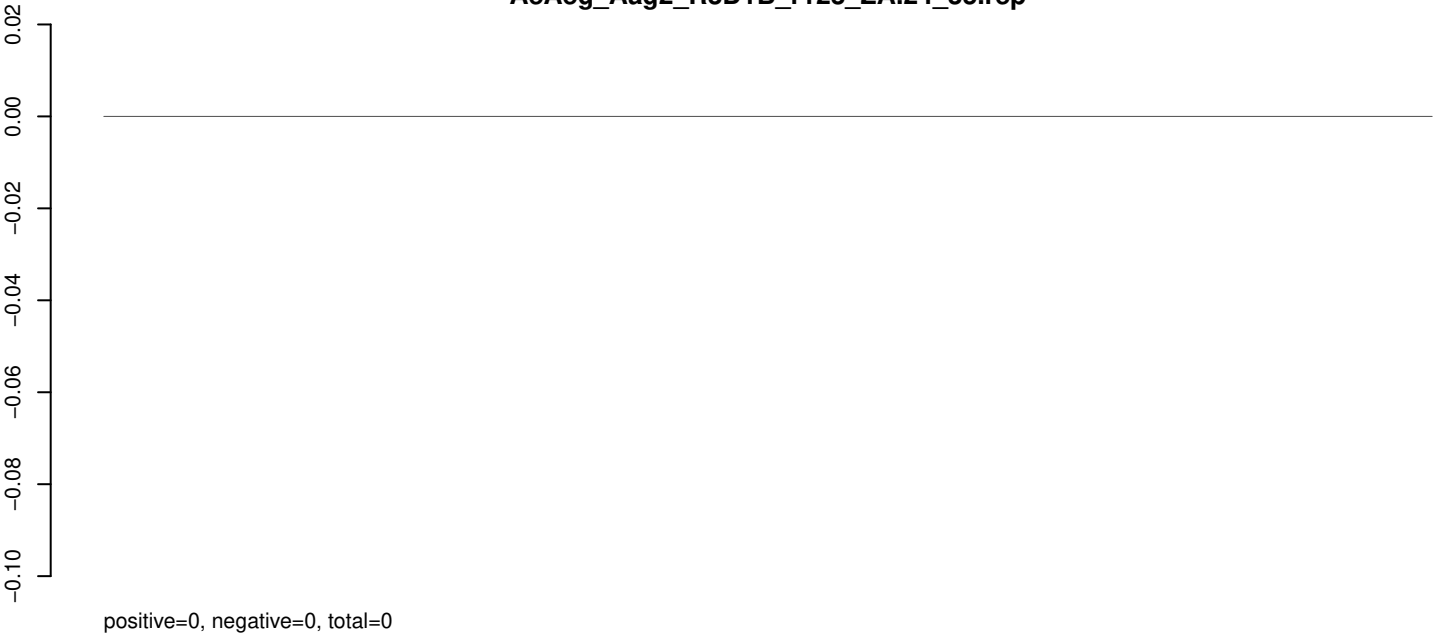


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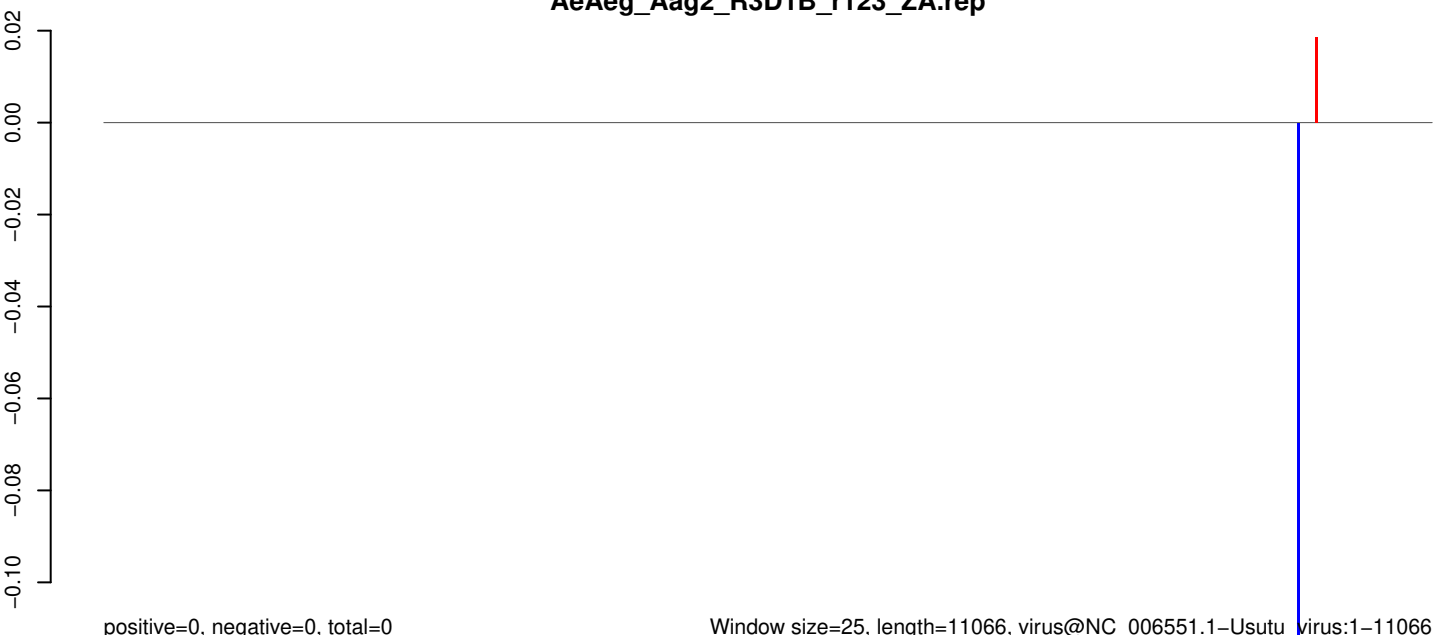
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



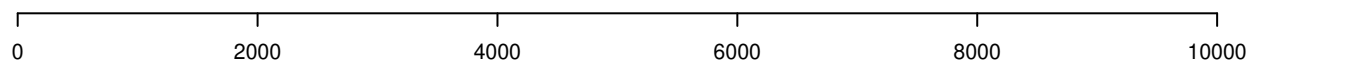
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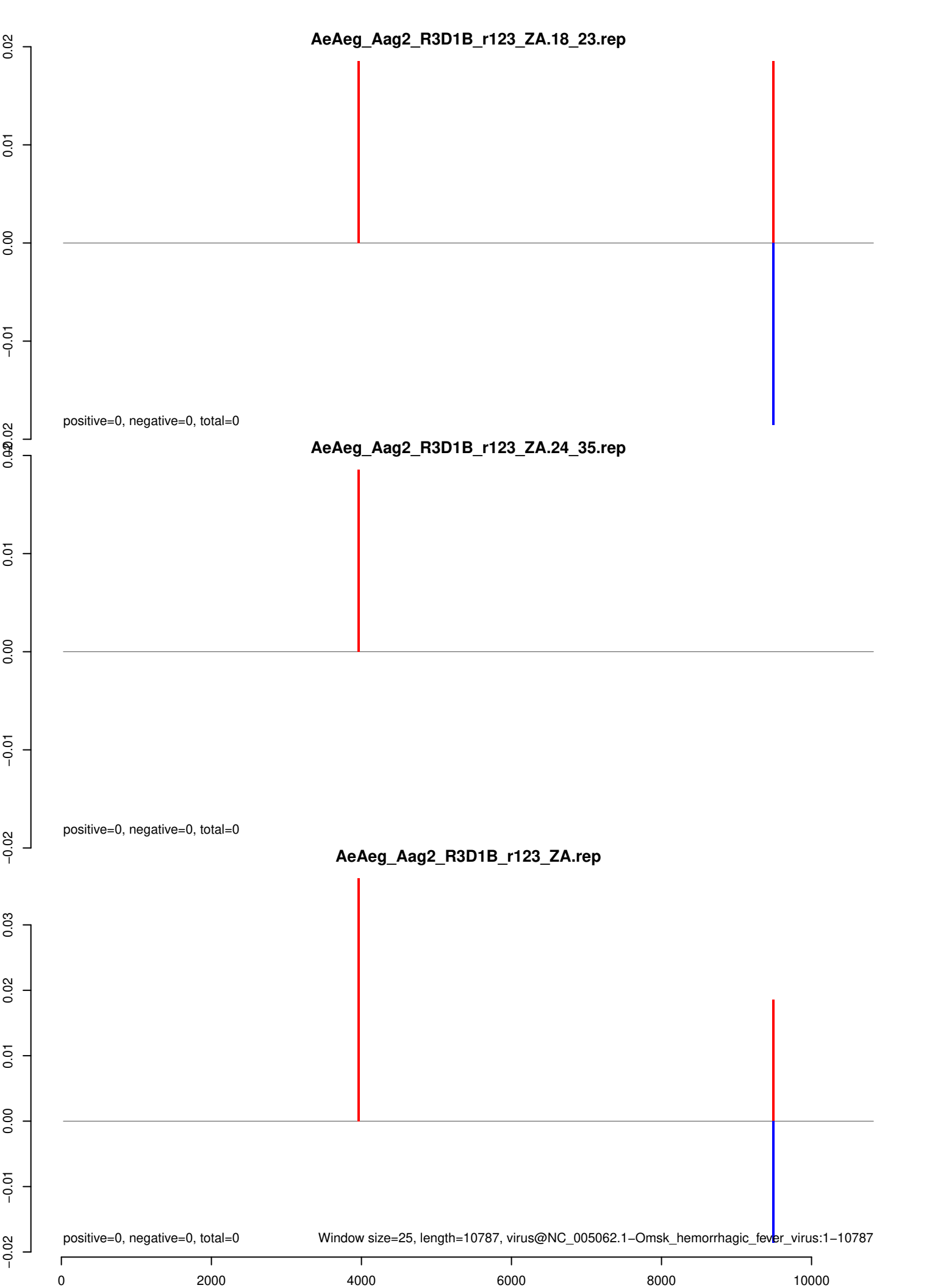
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positive=0, negative=0, total=0

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

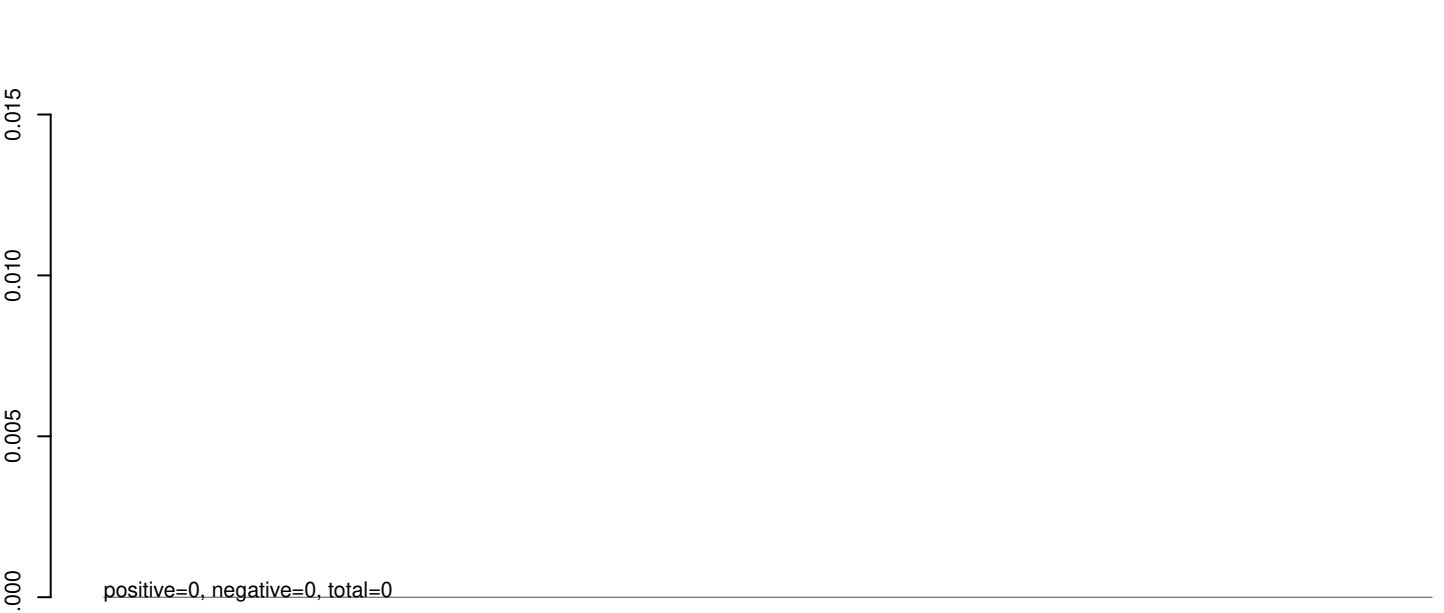




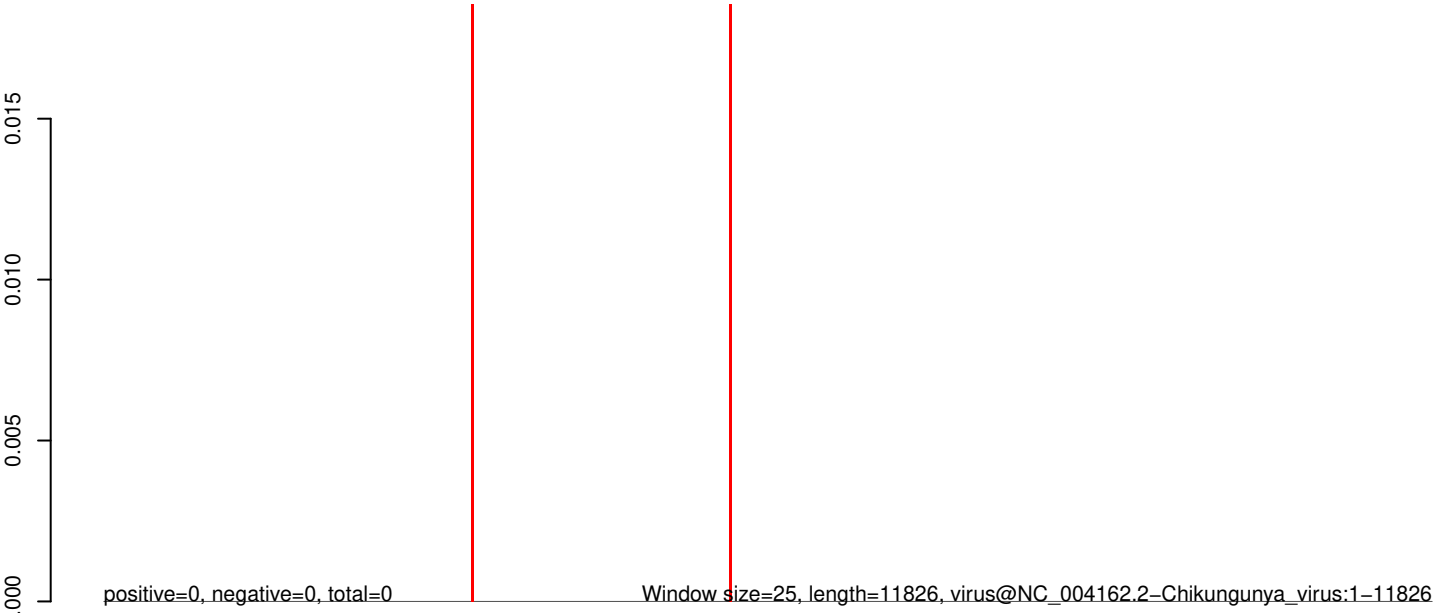
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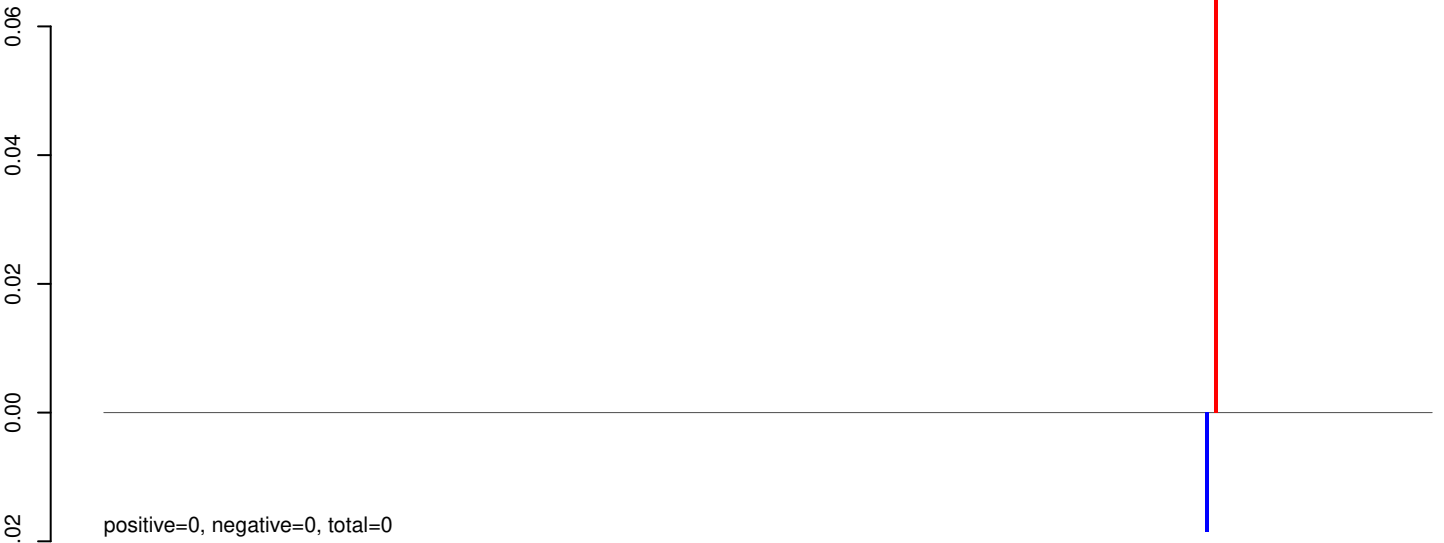


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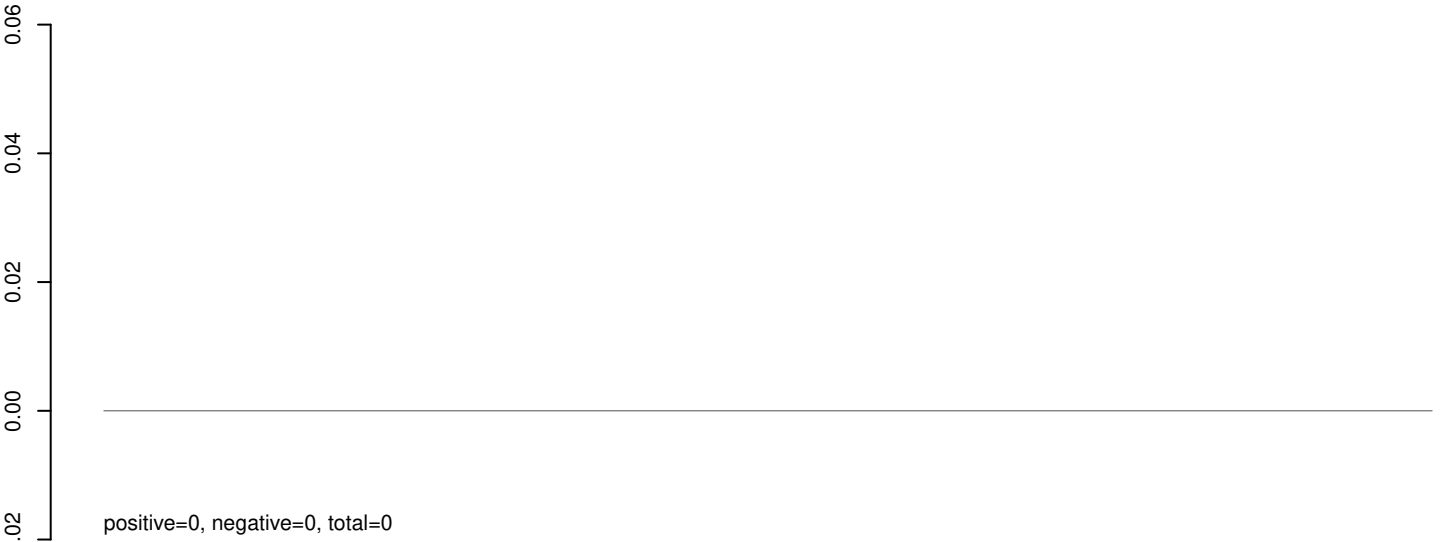


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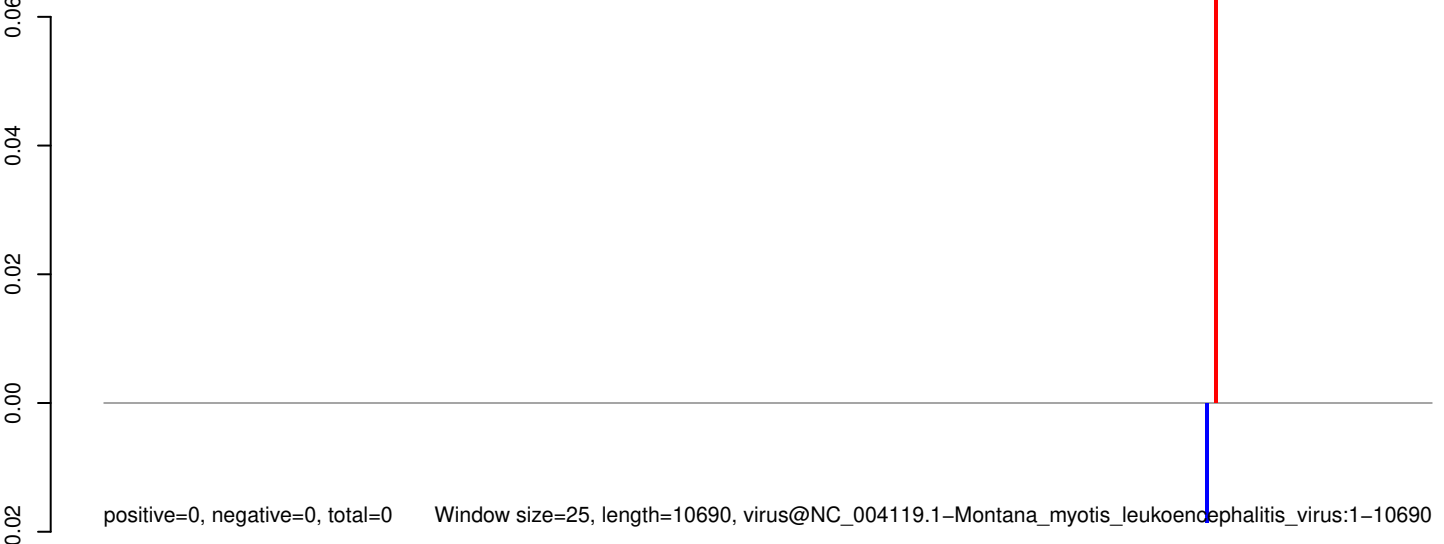
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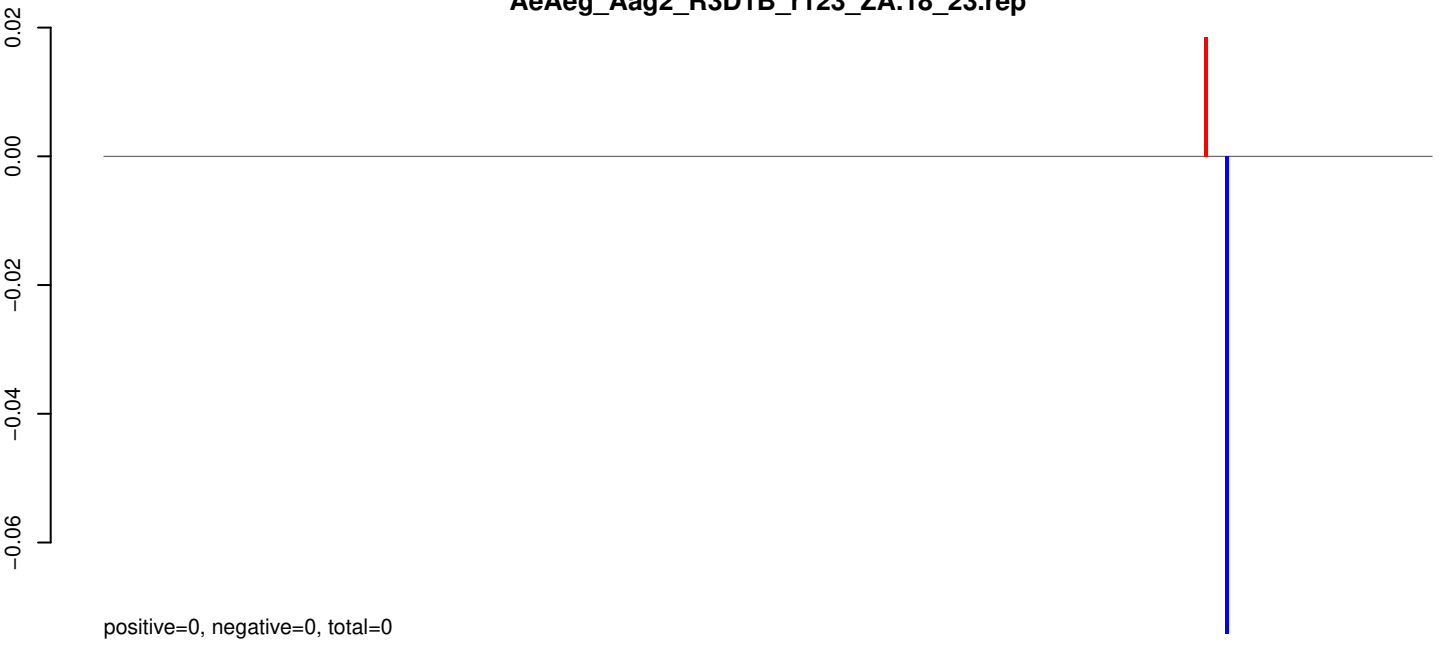


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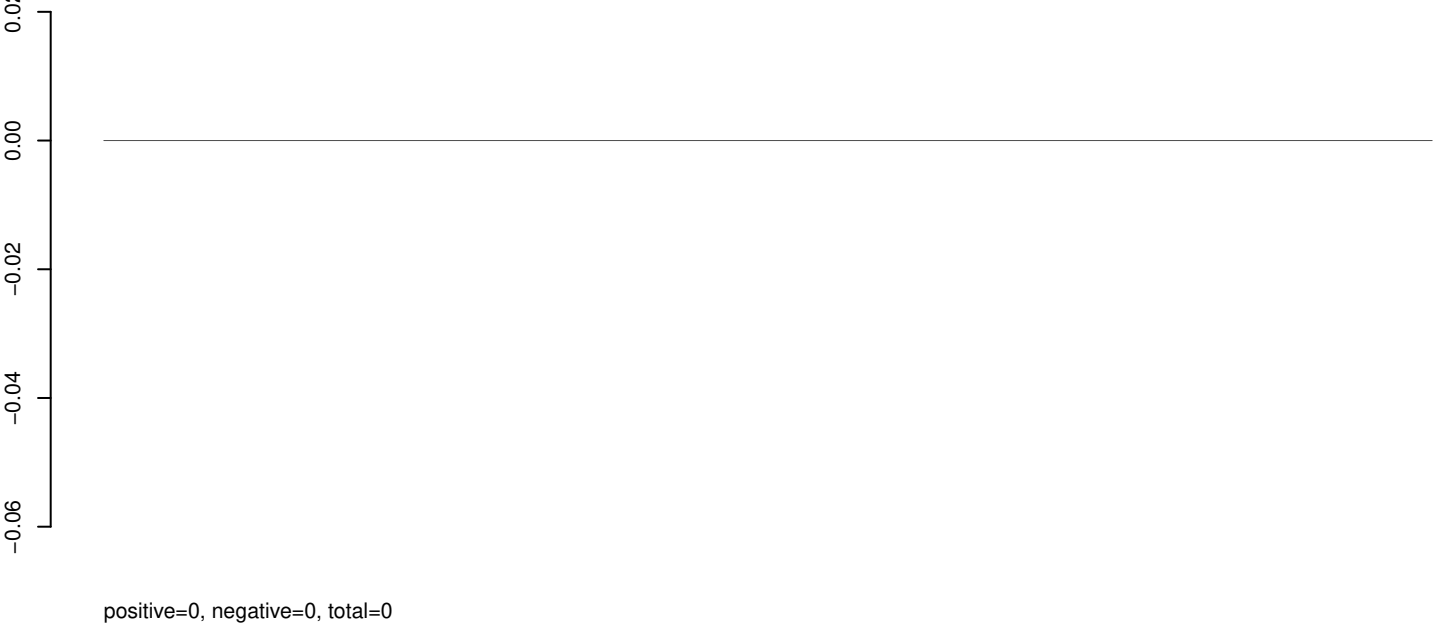


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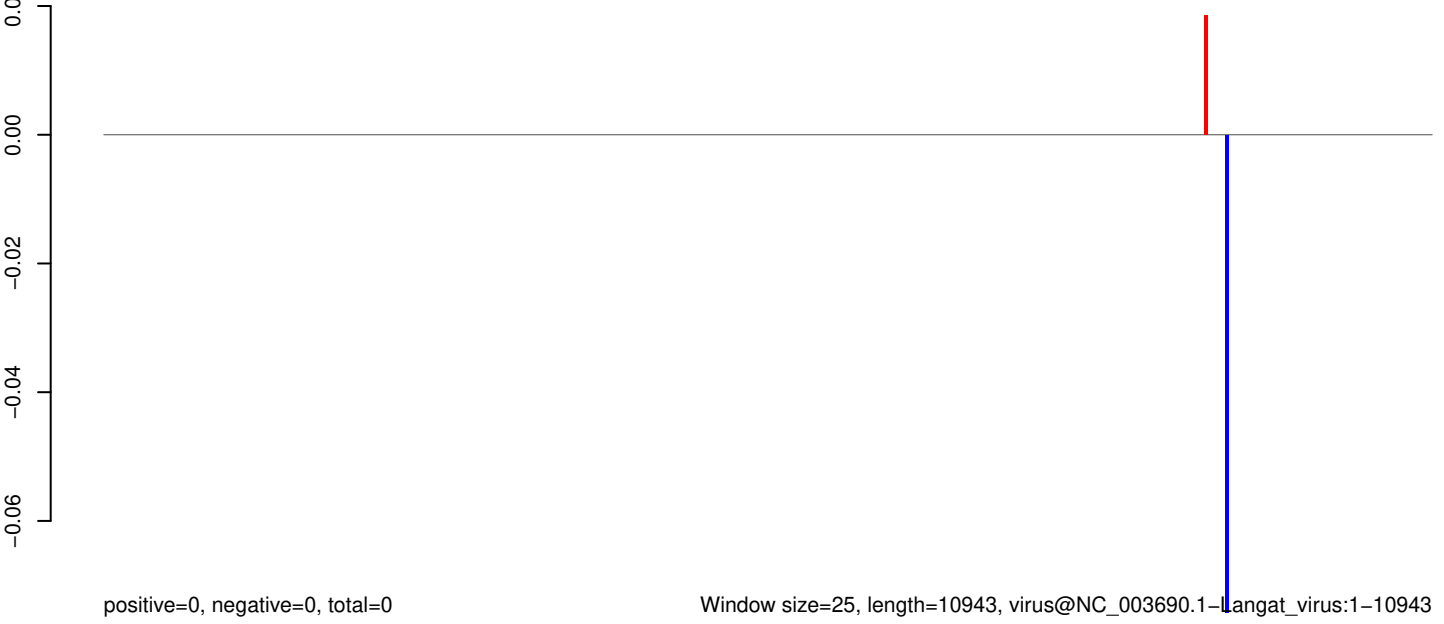
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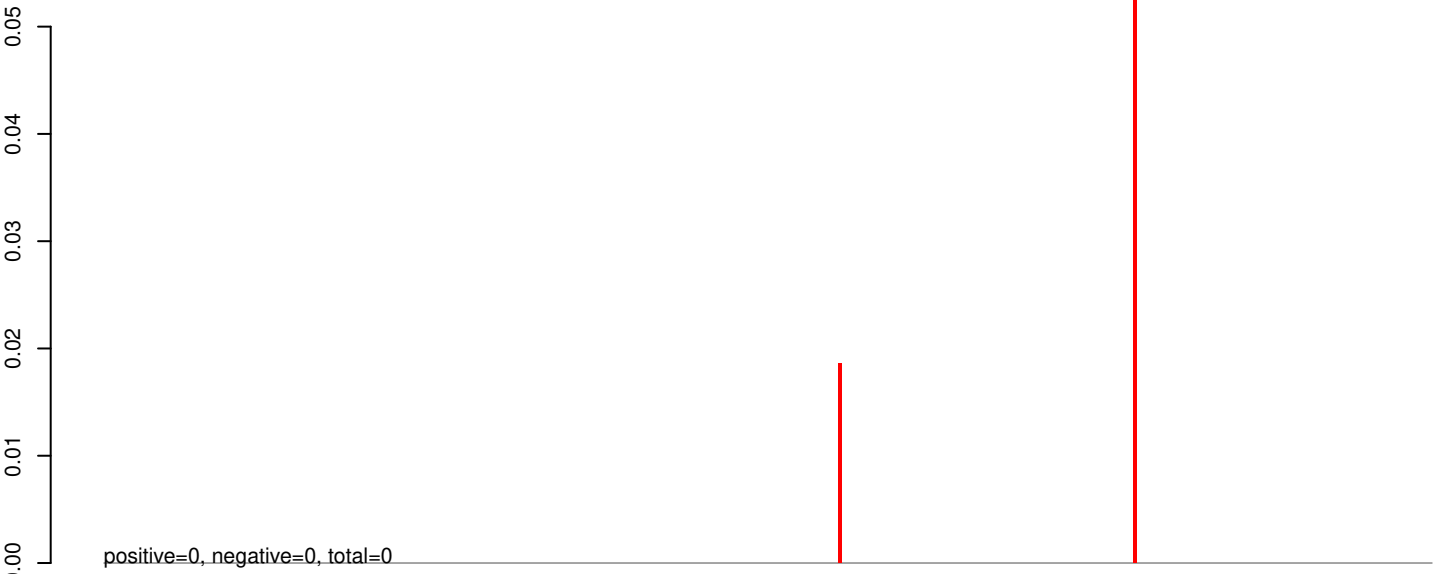
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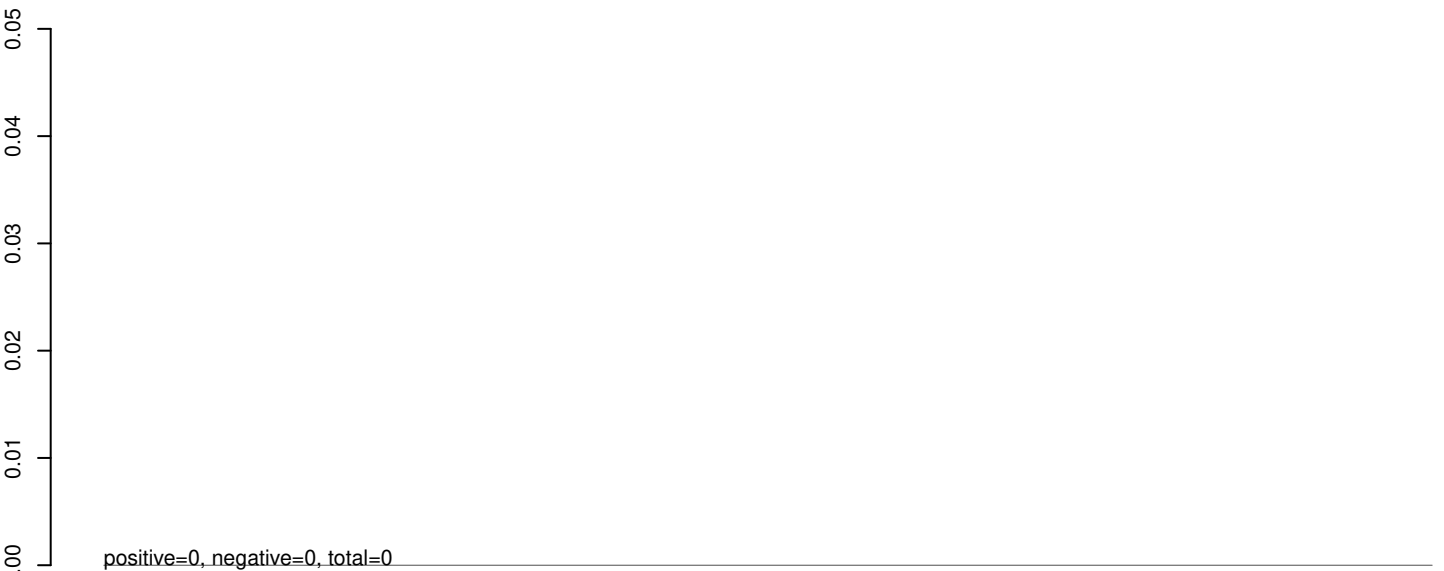
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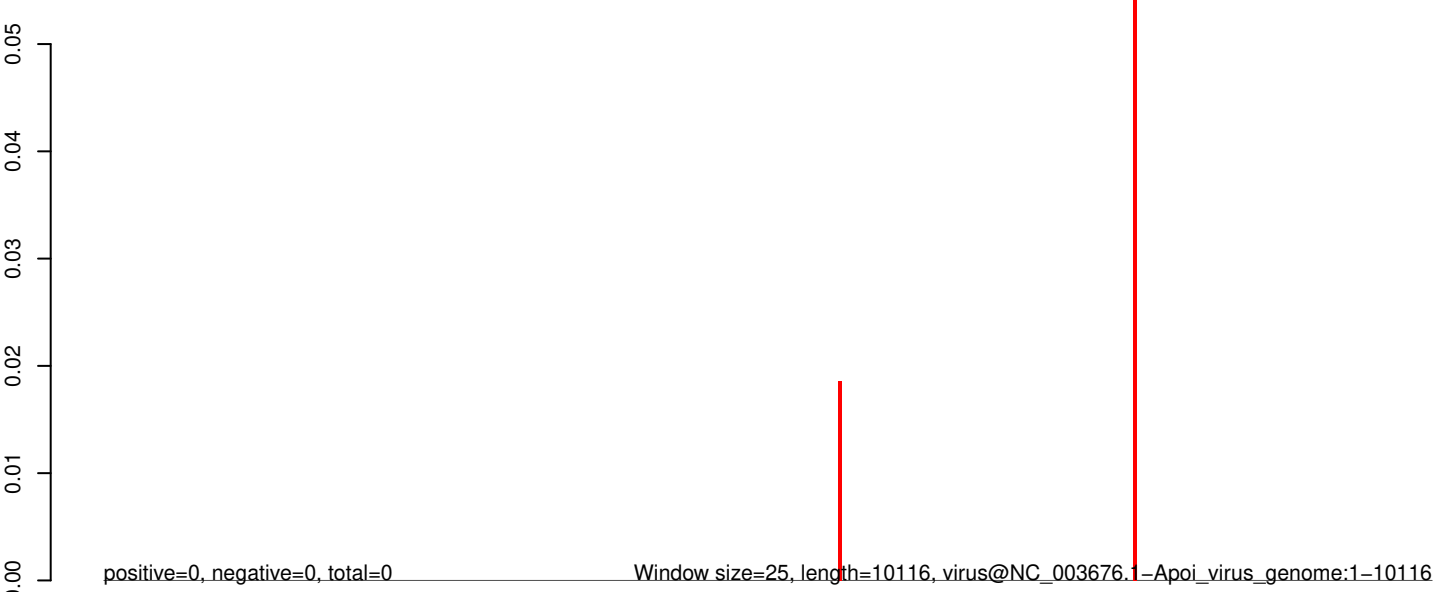
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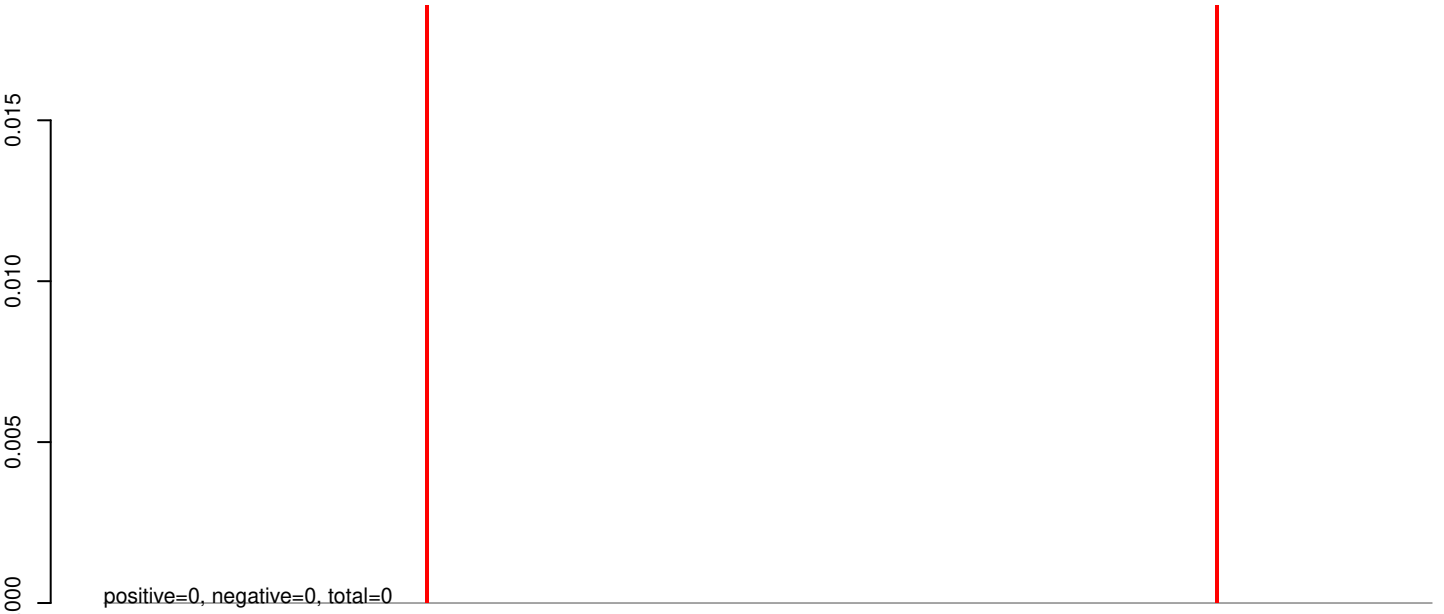
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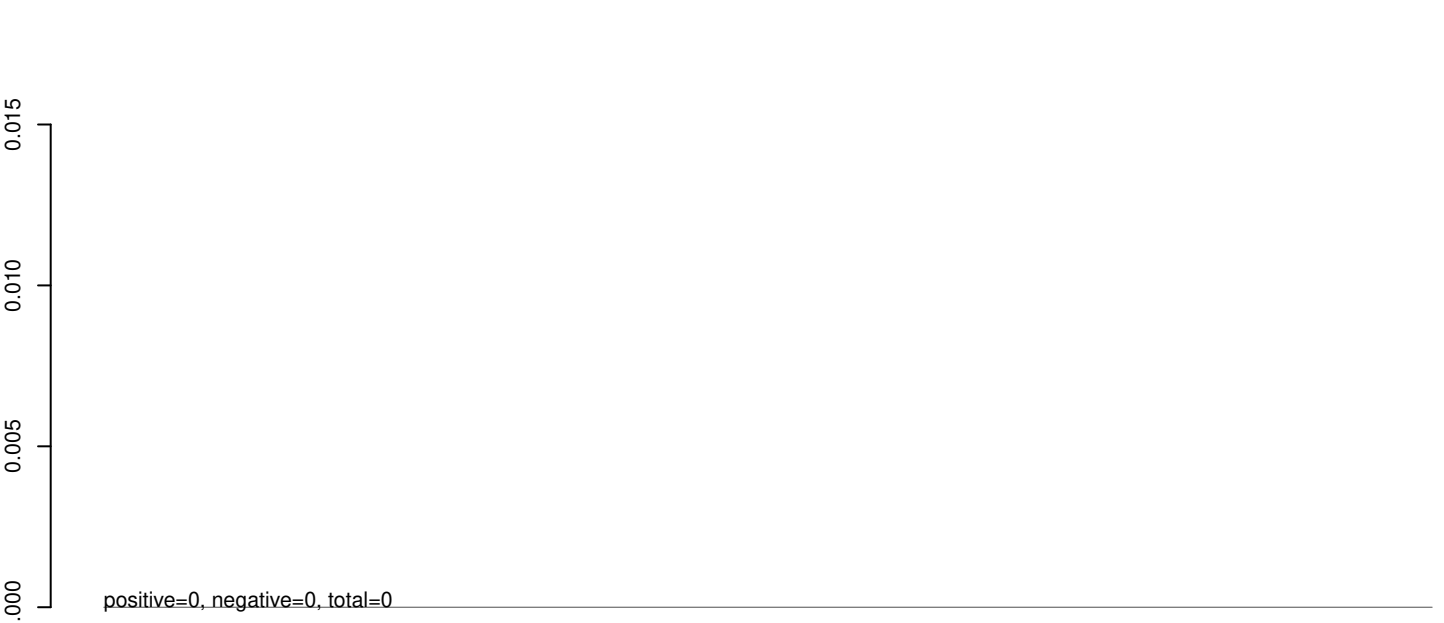
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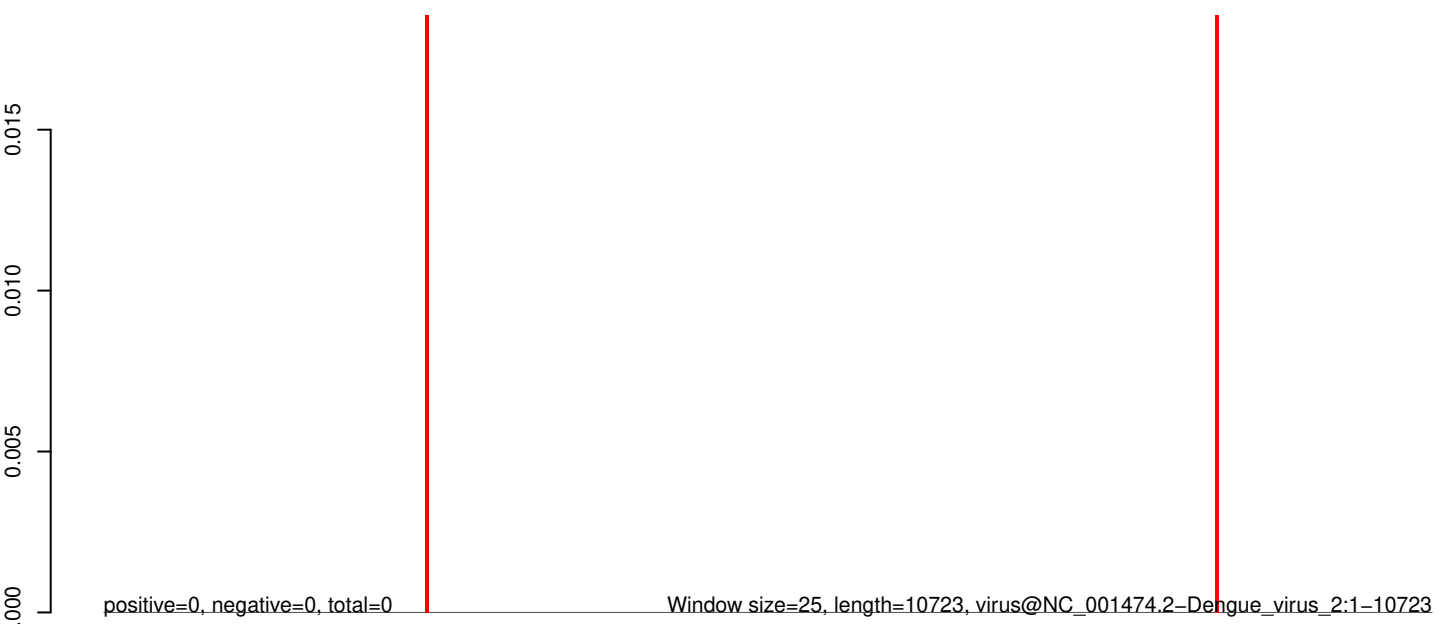
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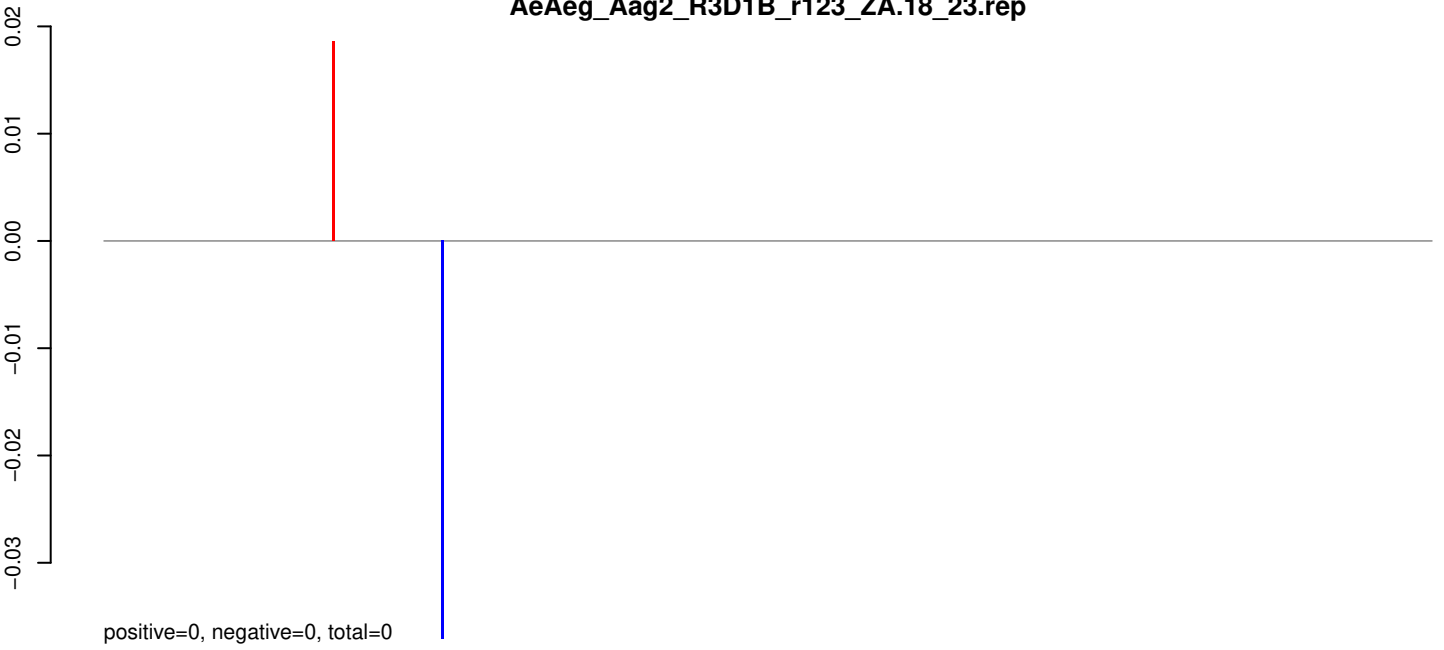


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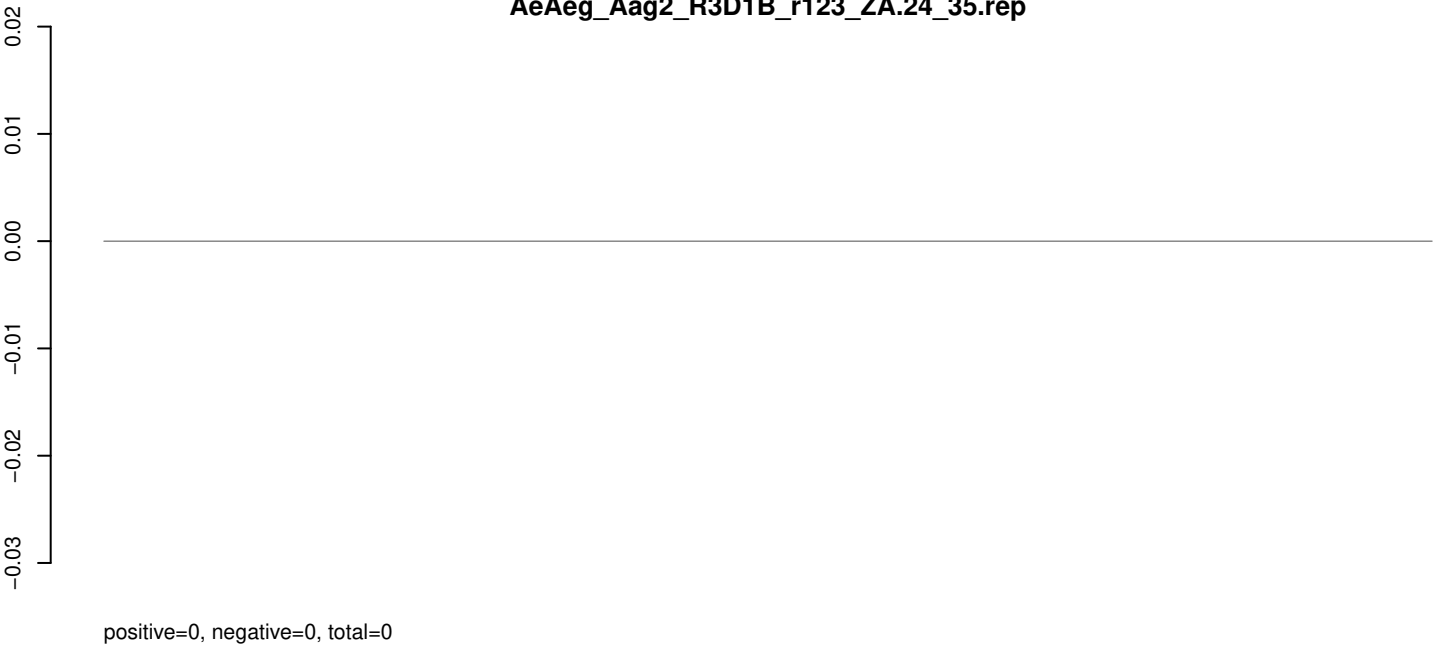


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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



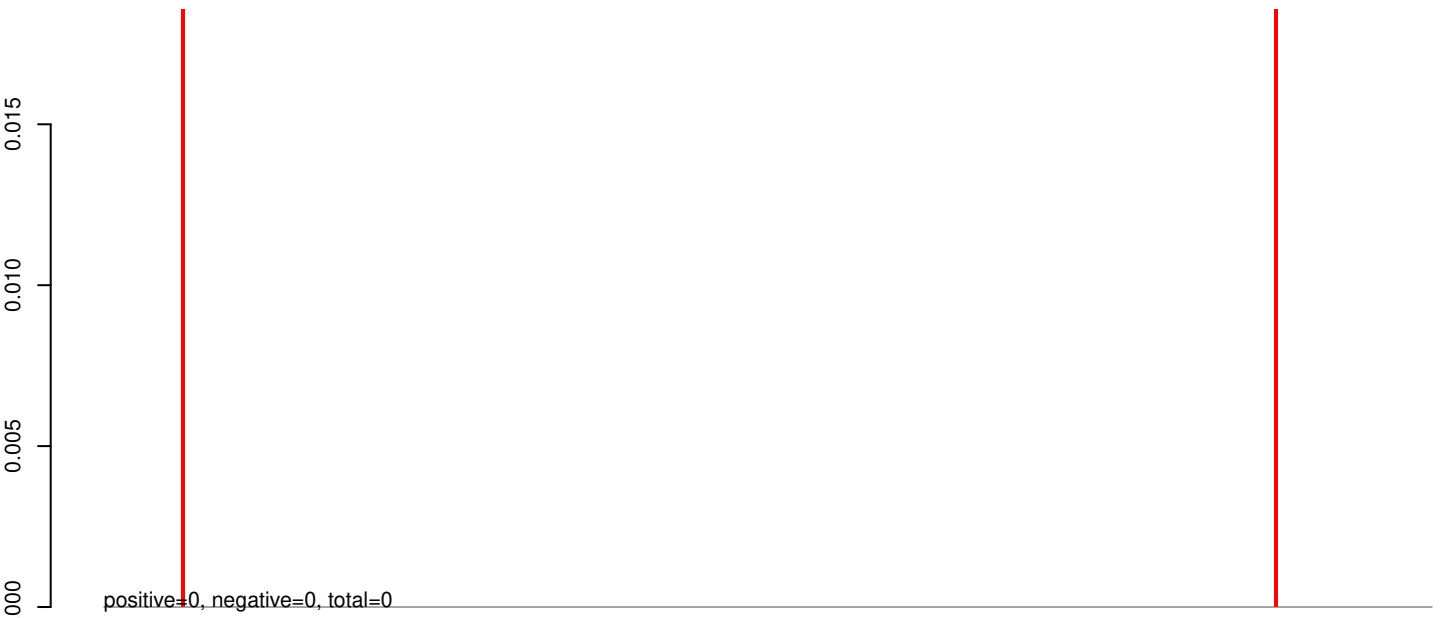
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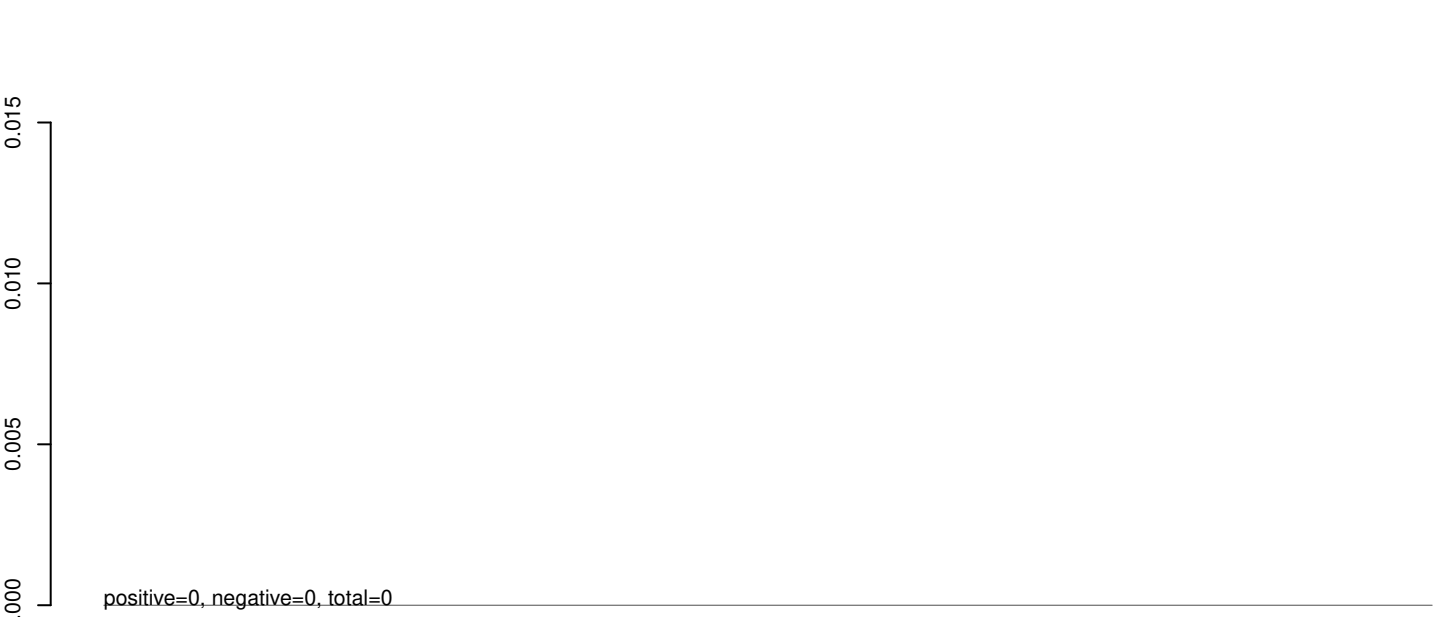
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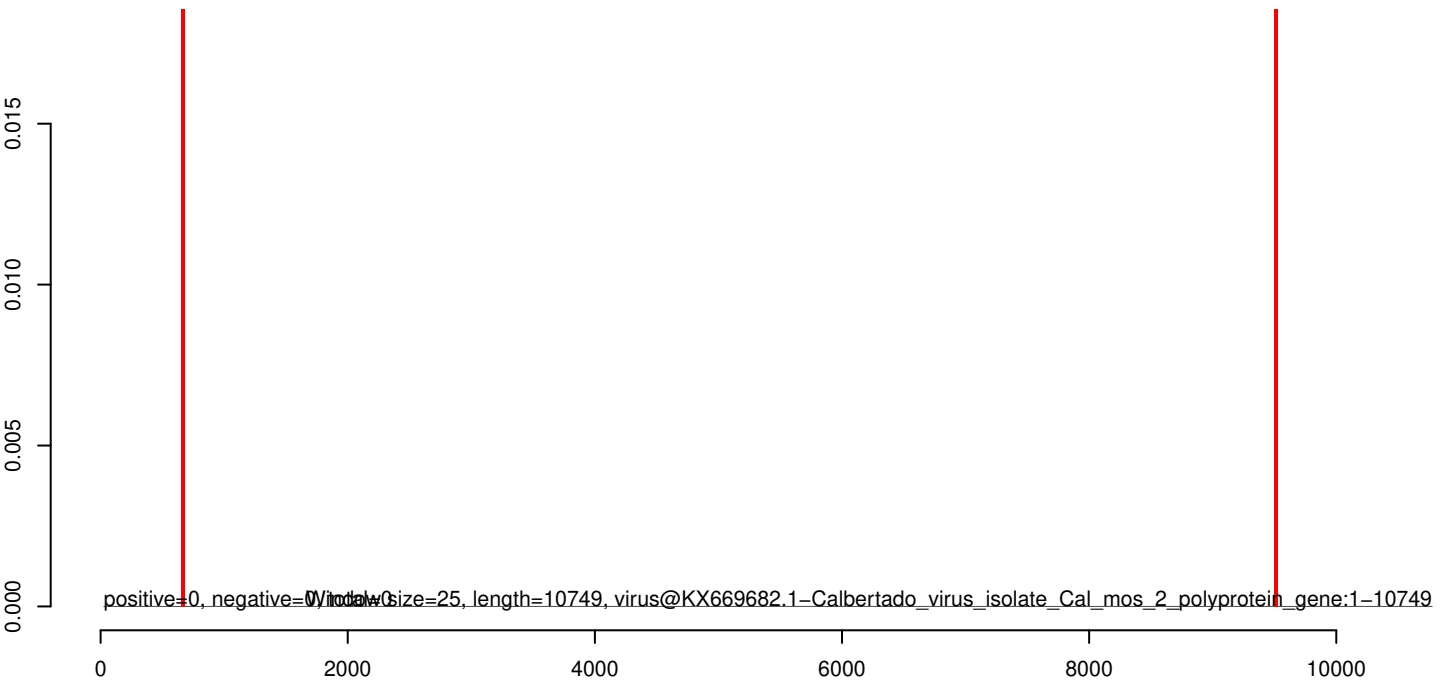
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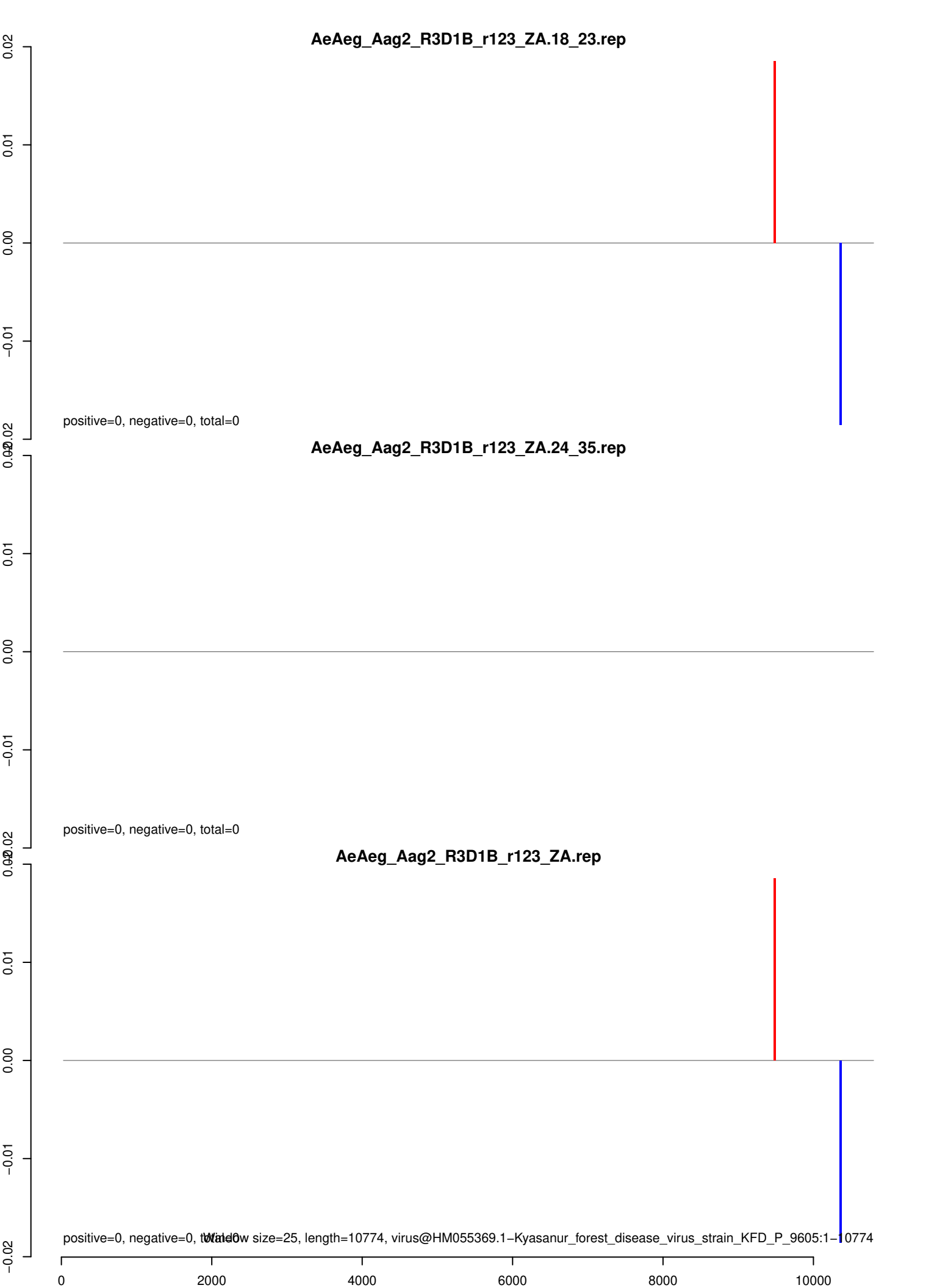


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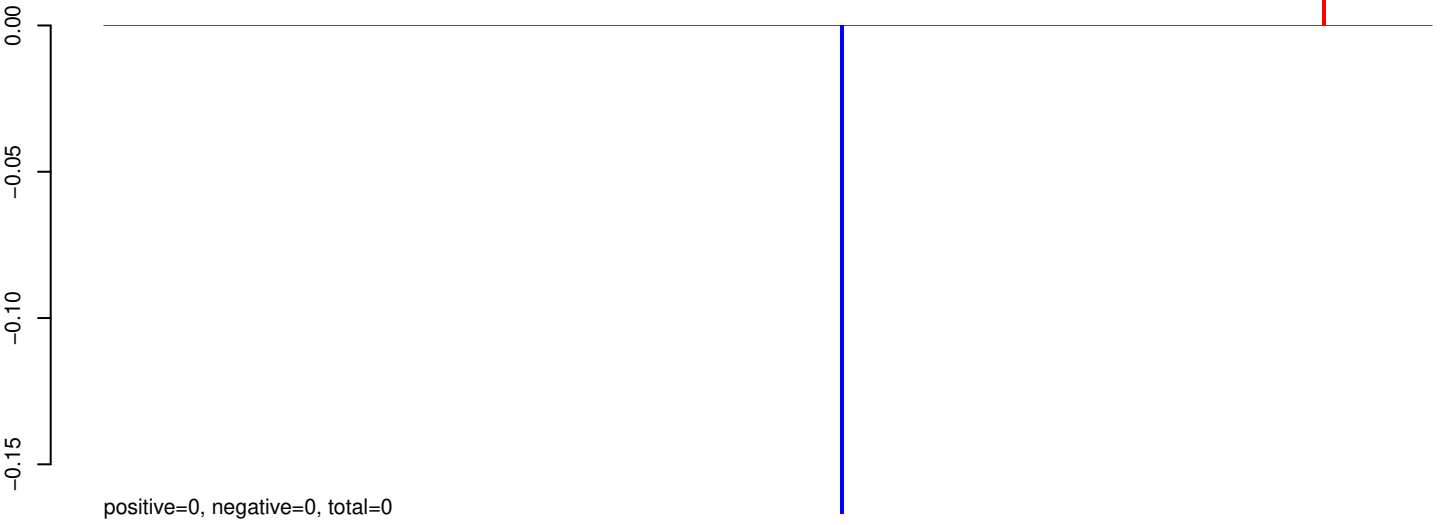


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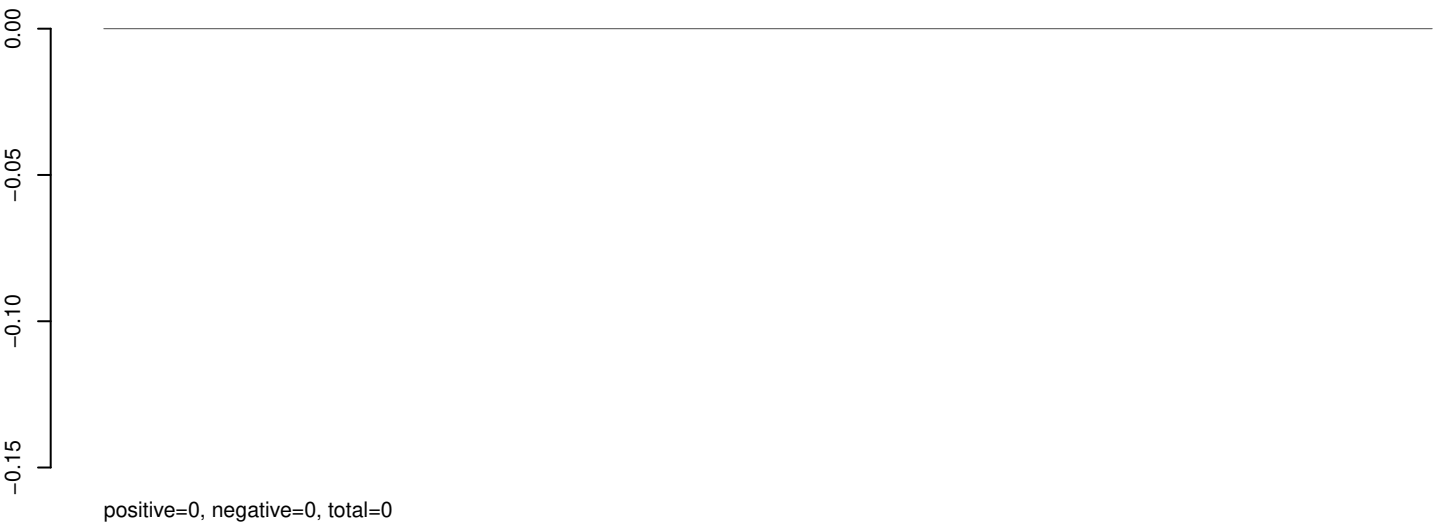




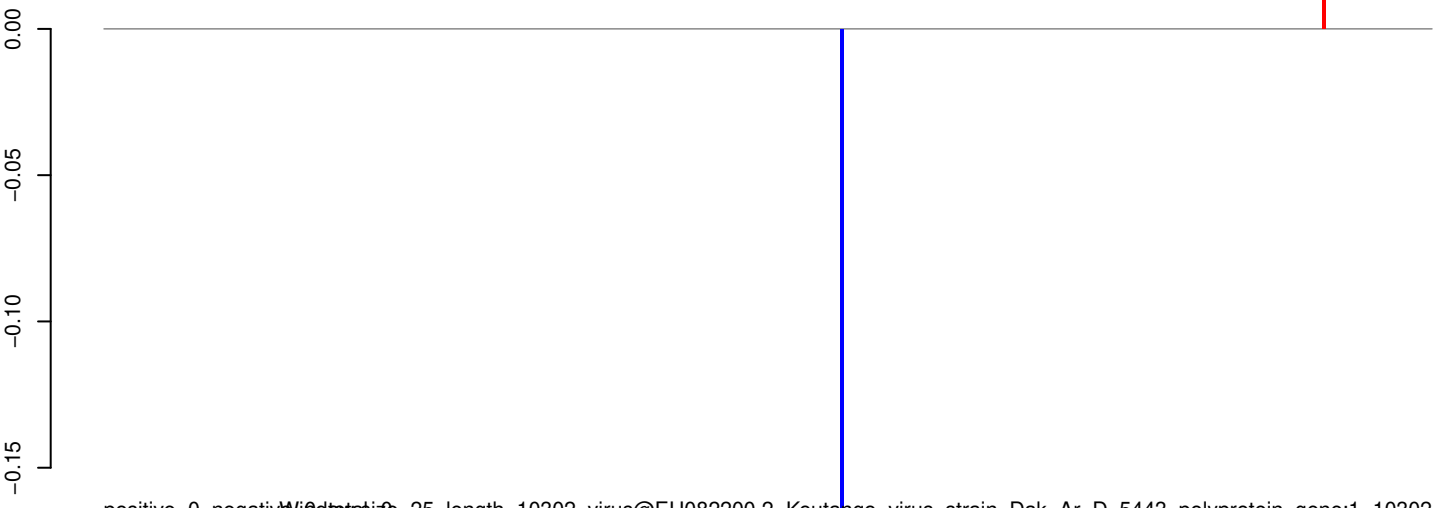
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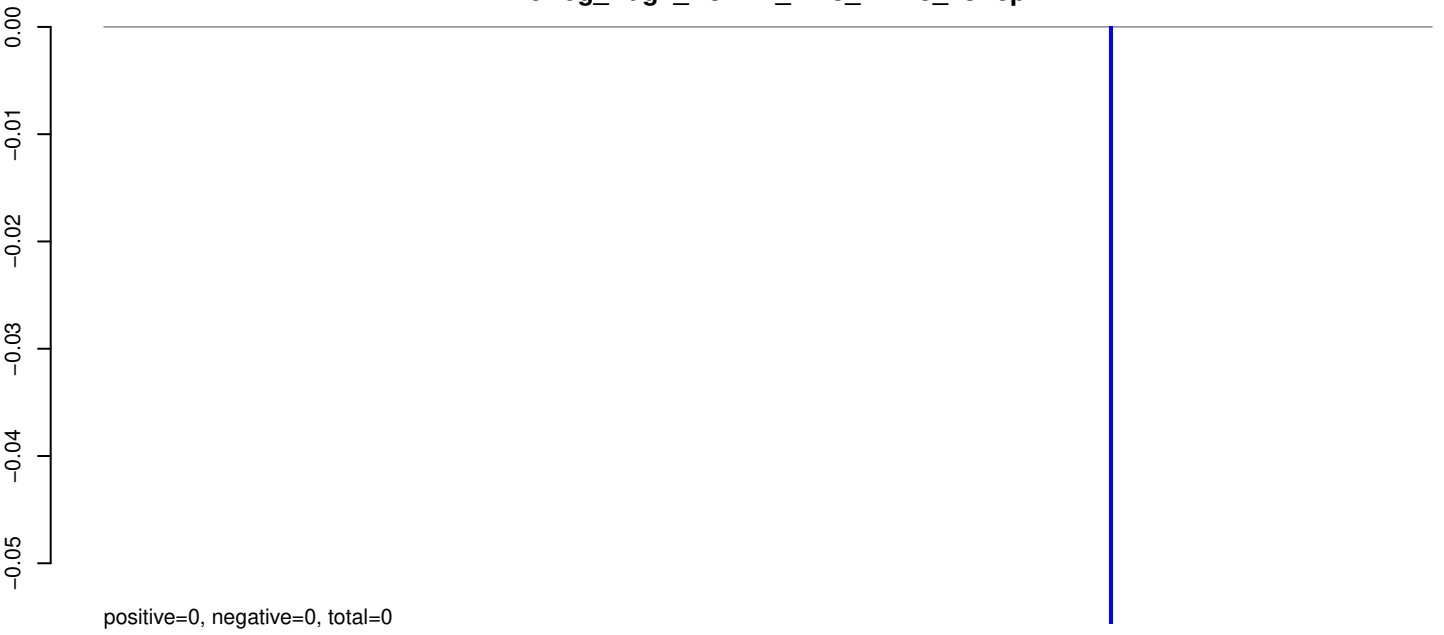
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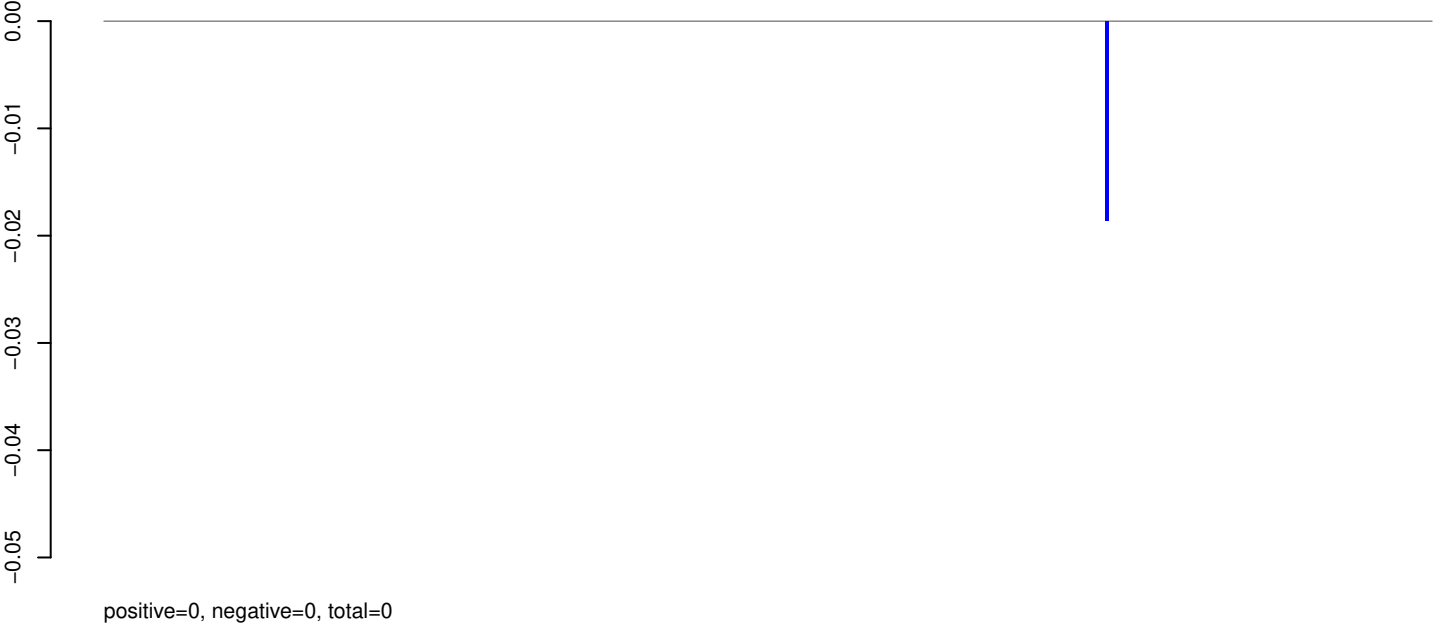
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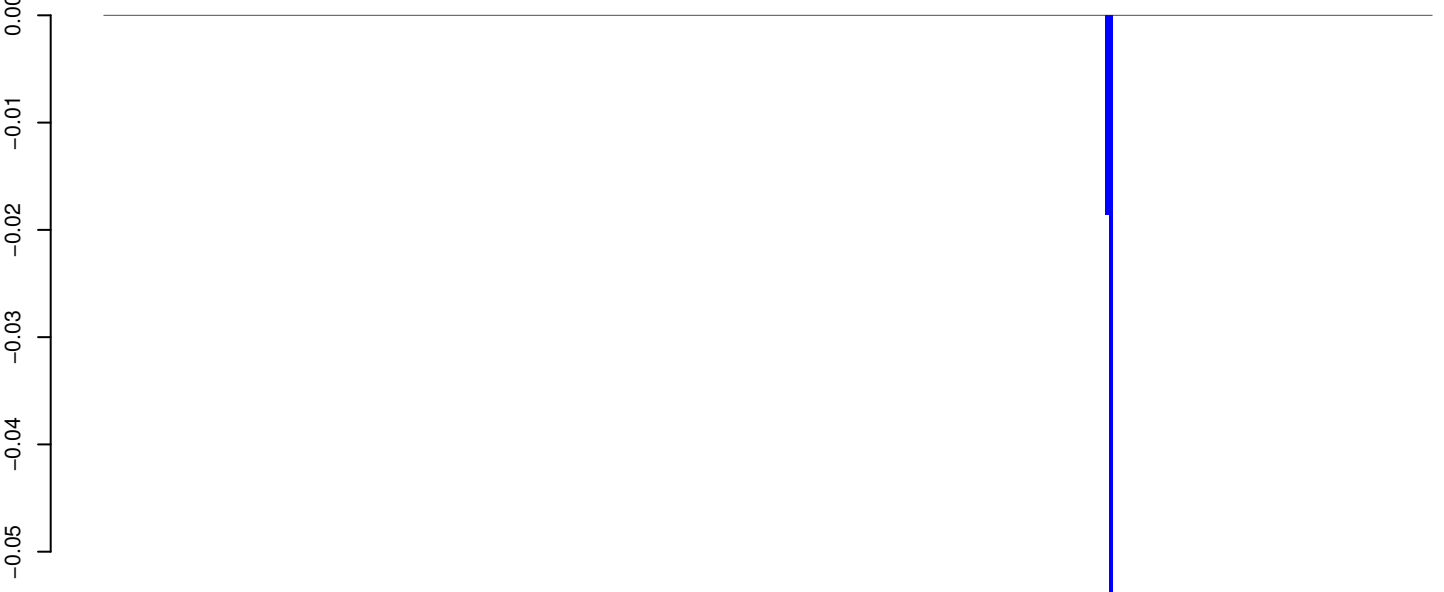
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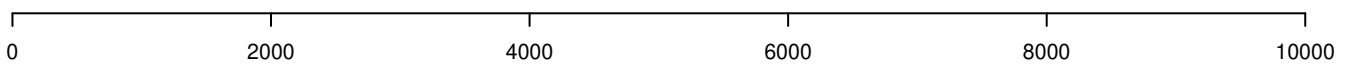
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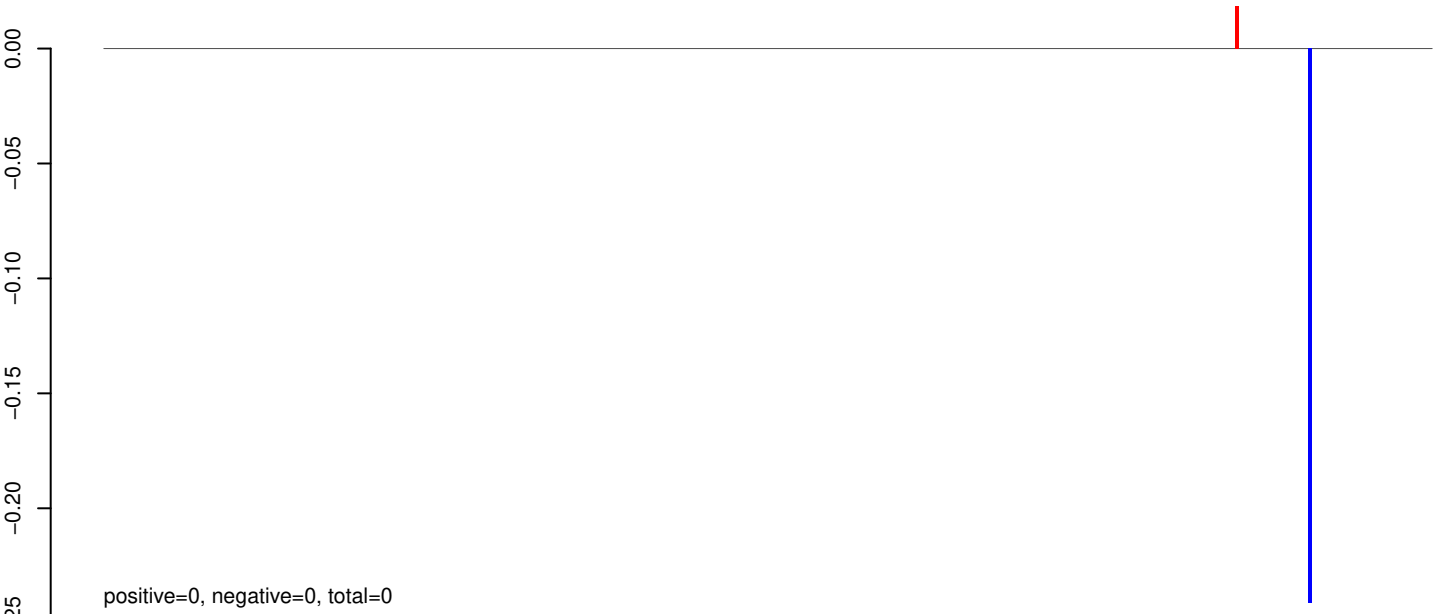
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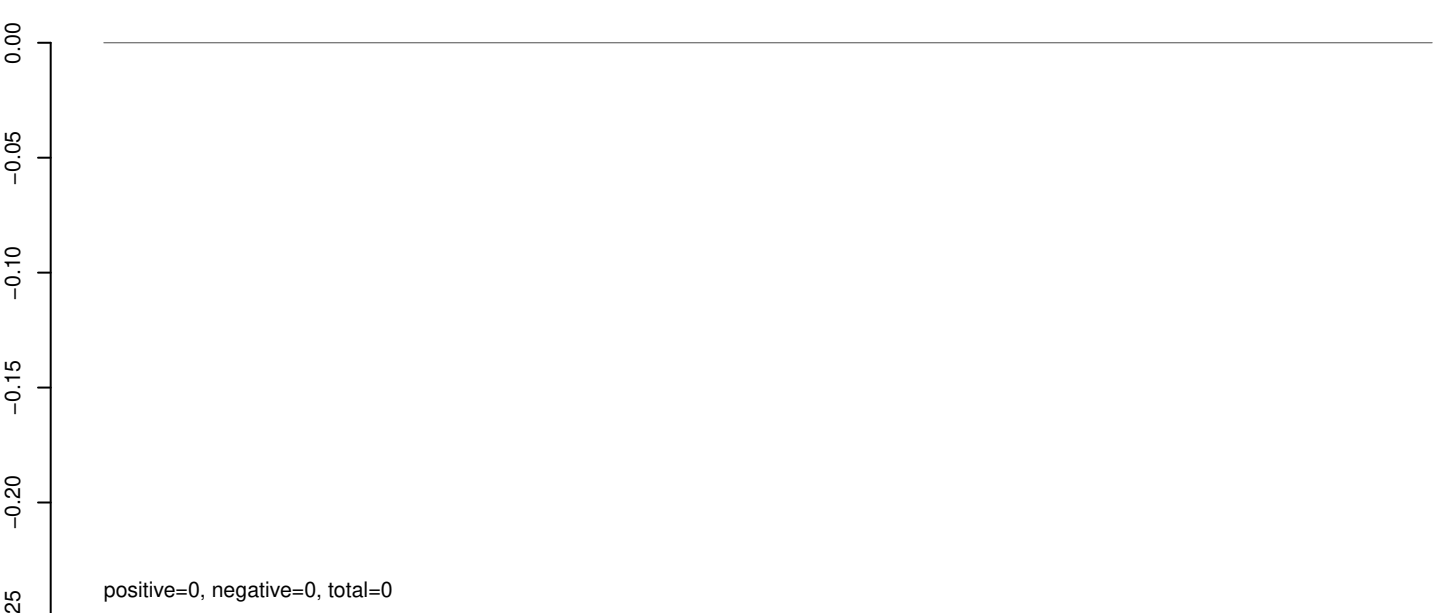
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AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



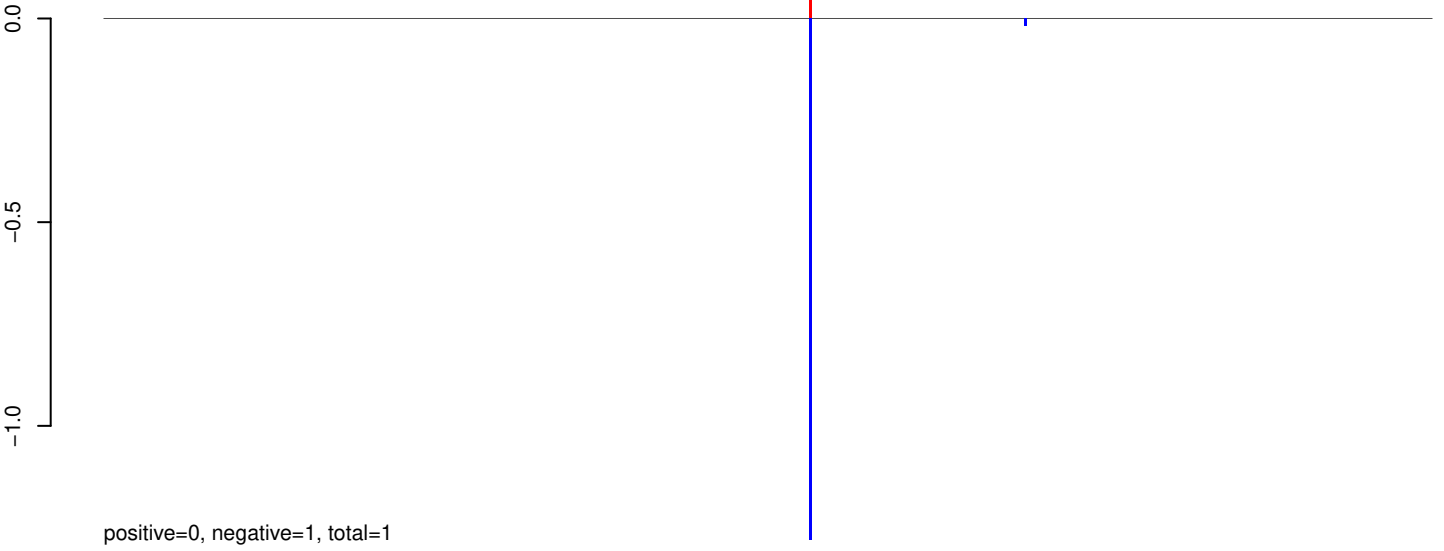
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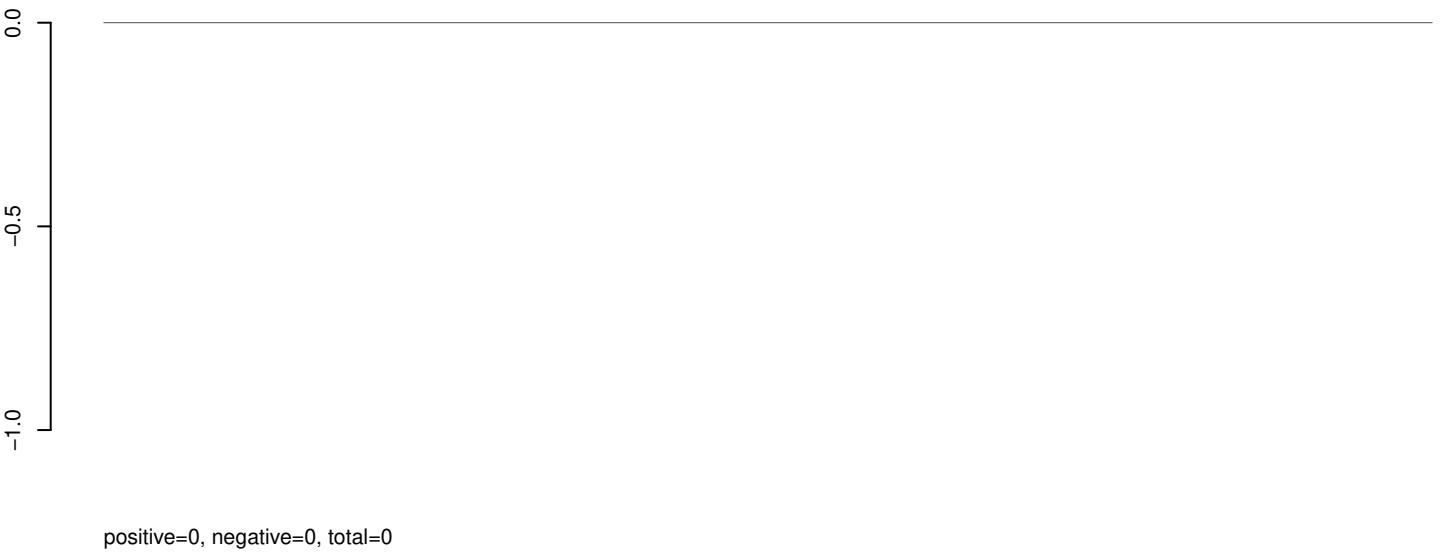
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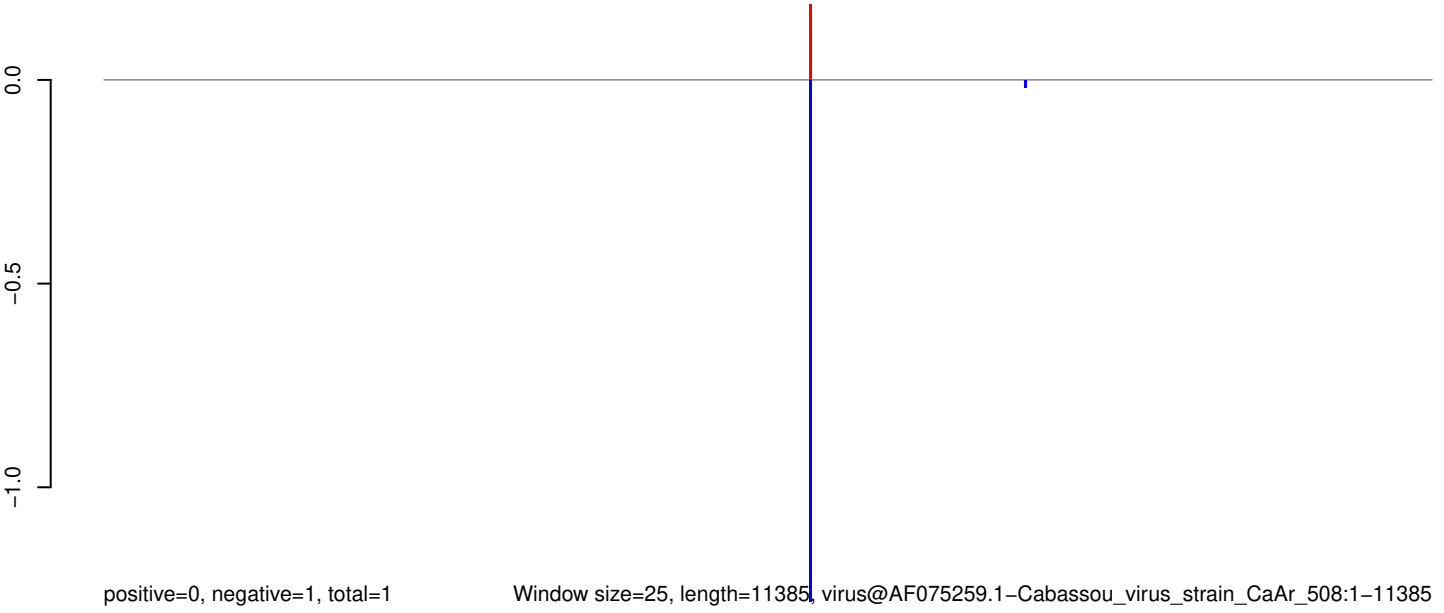
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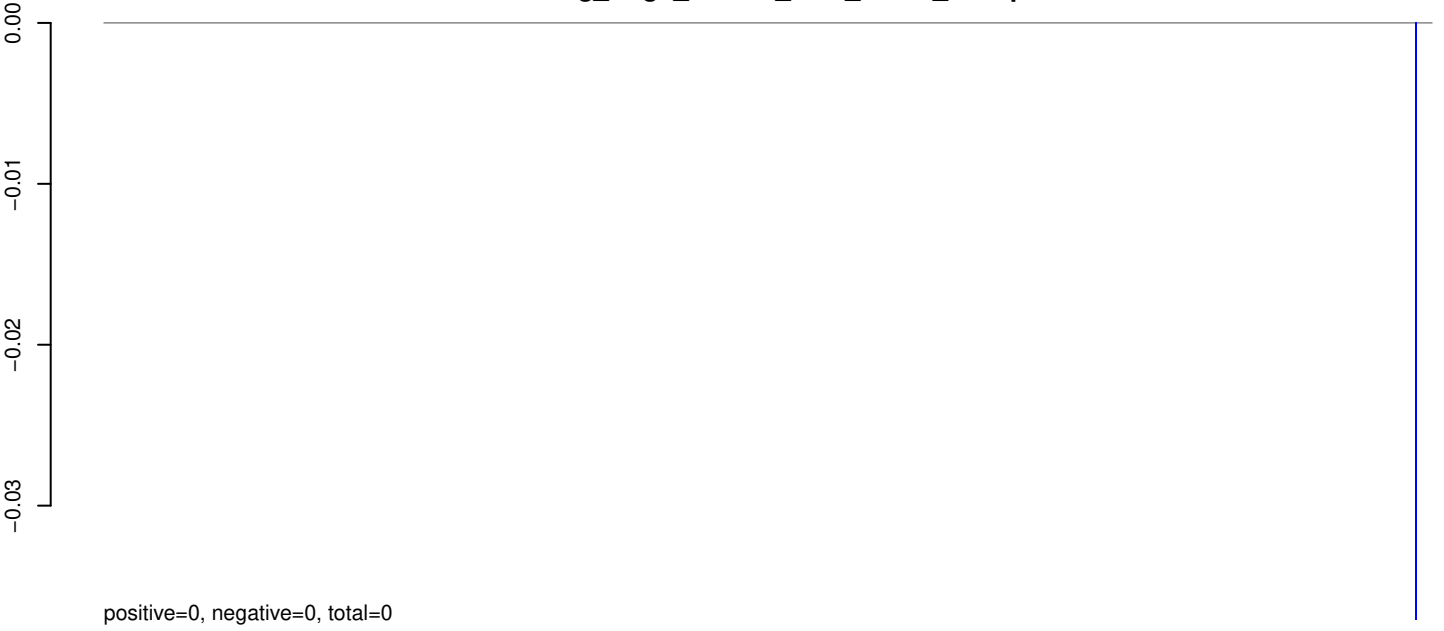
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AeAeg_Aag2_R3D1B_r123_ZA.rep

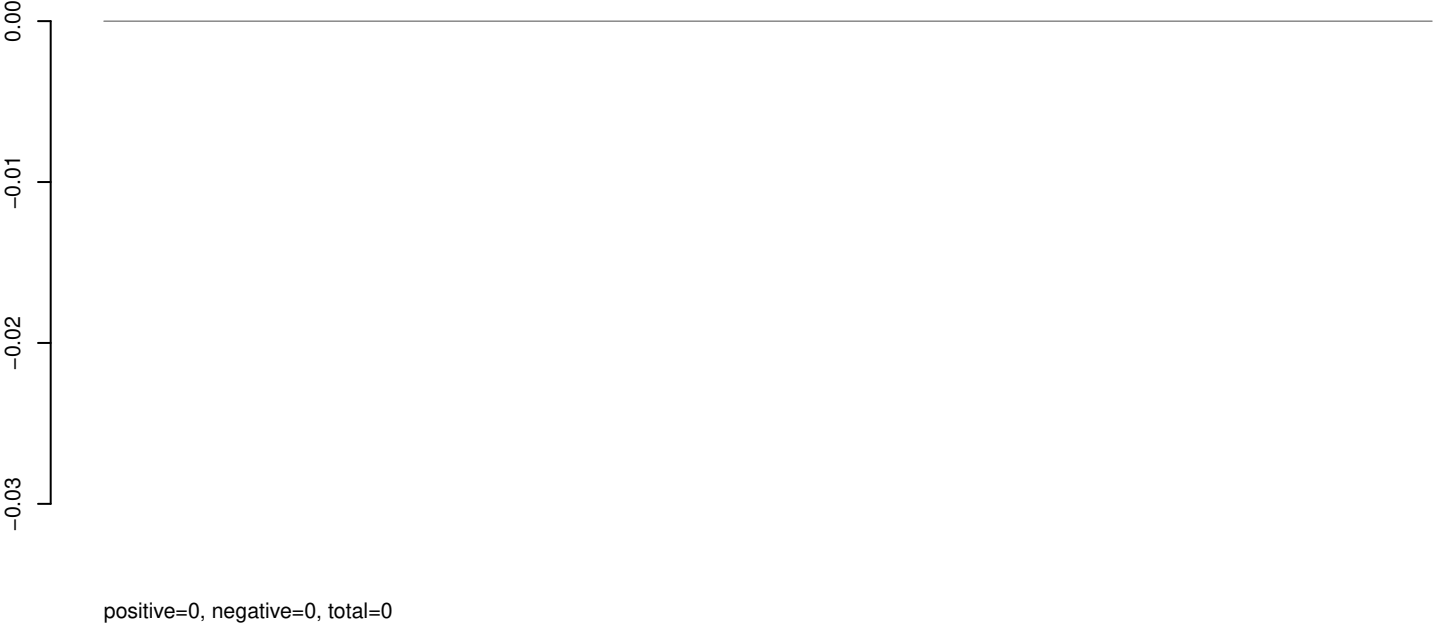


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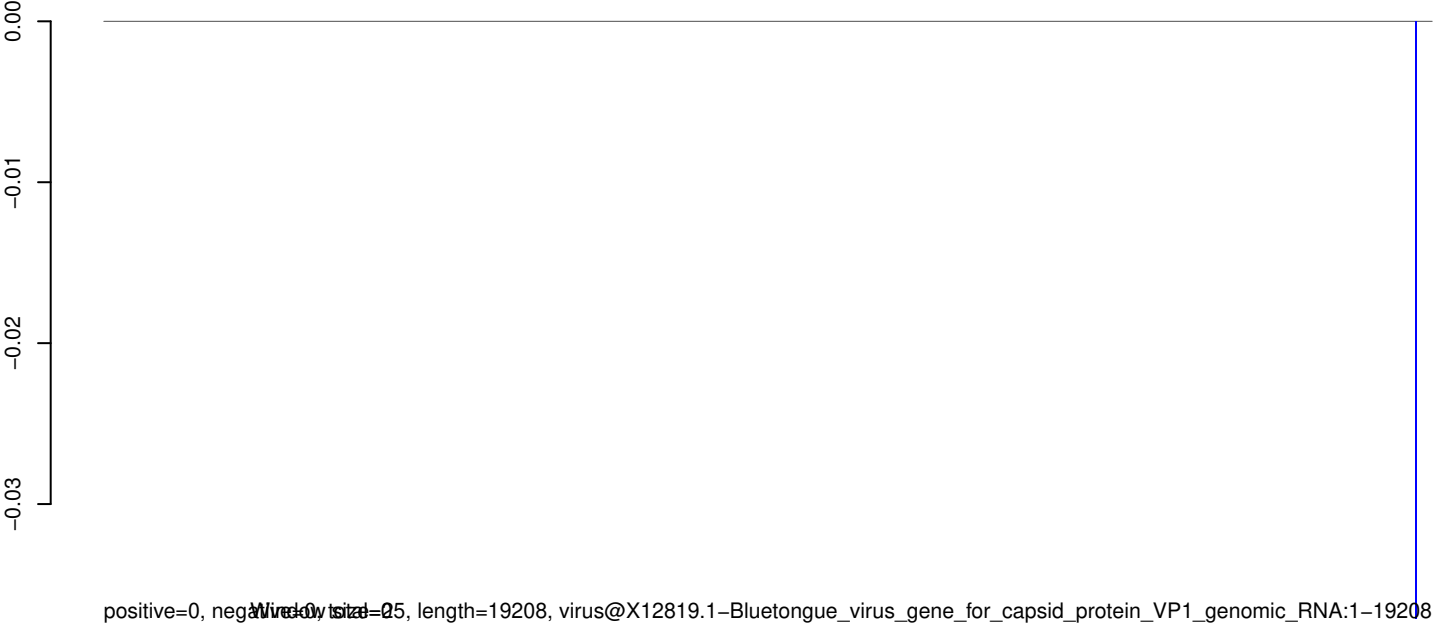
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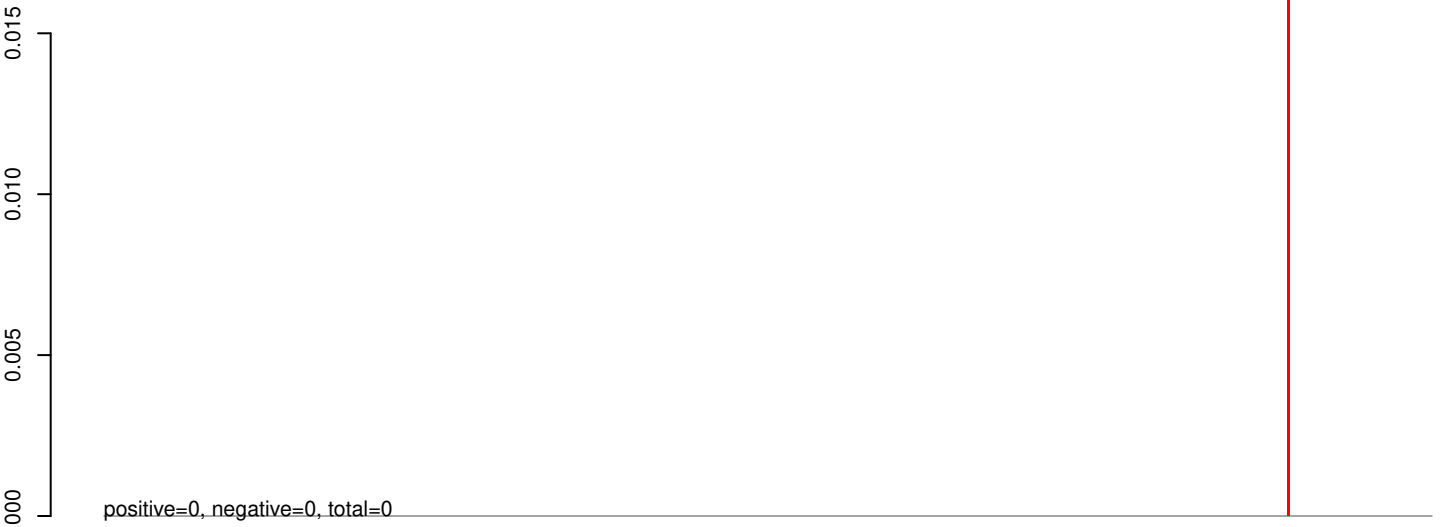
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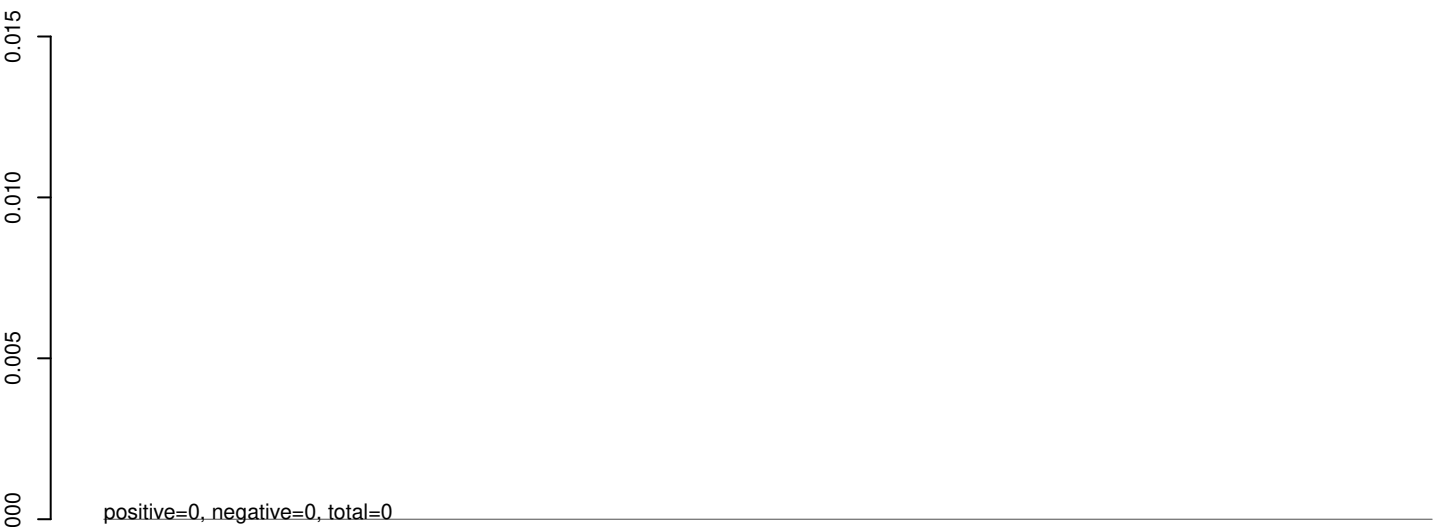
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0 5000 10000 15000 20000

AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



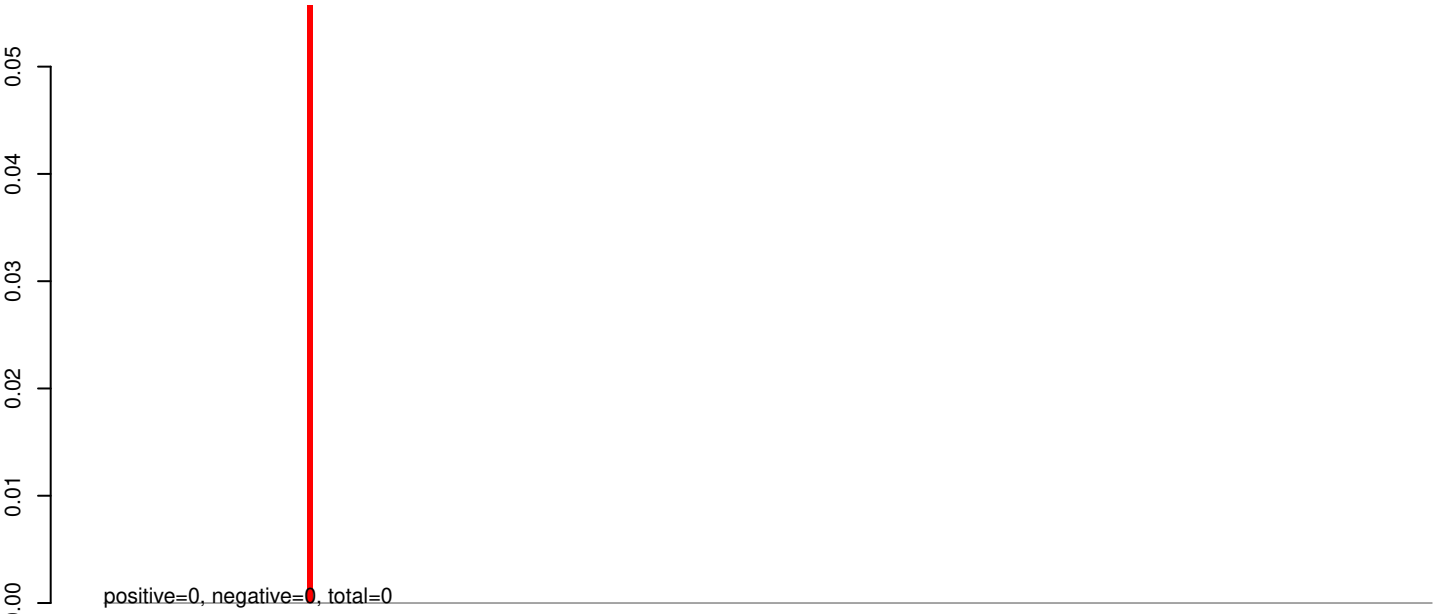
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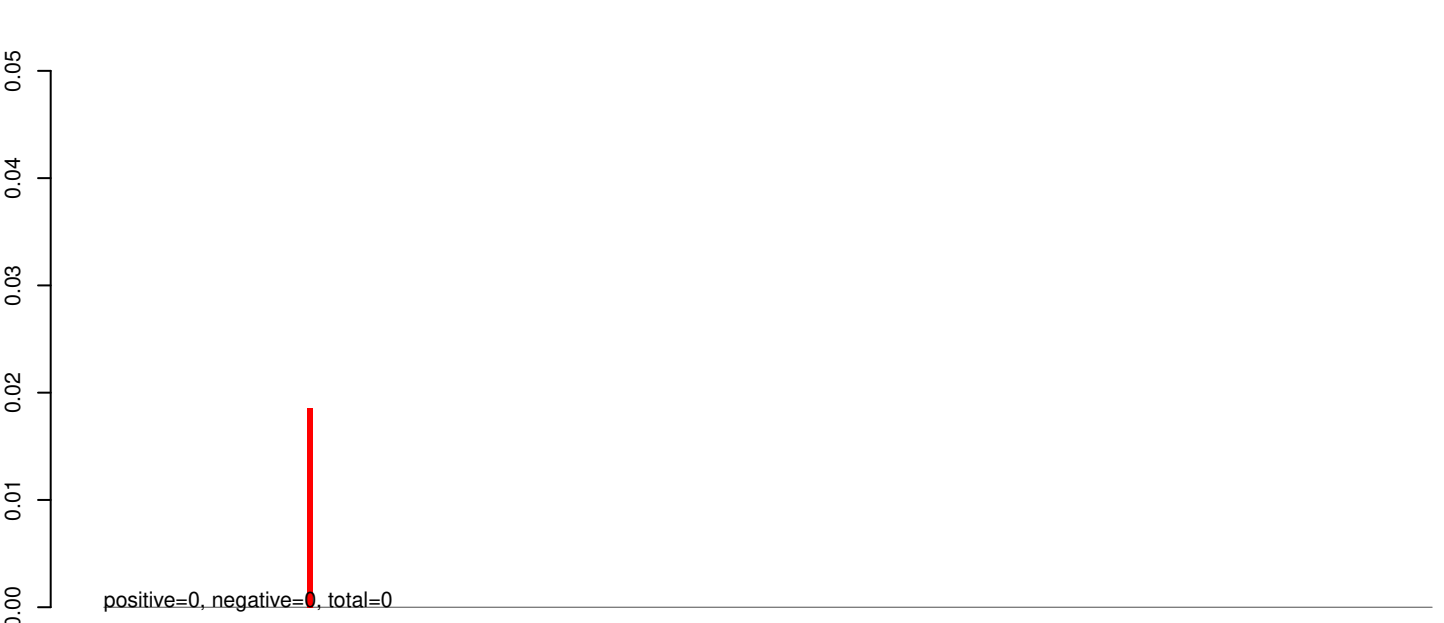
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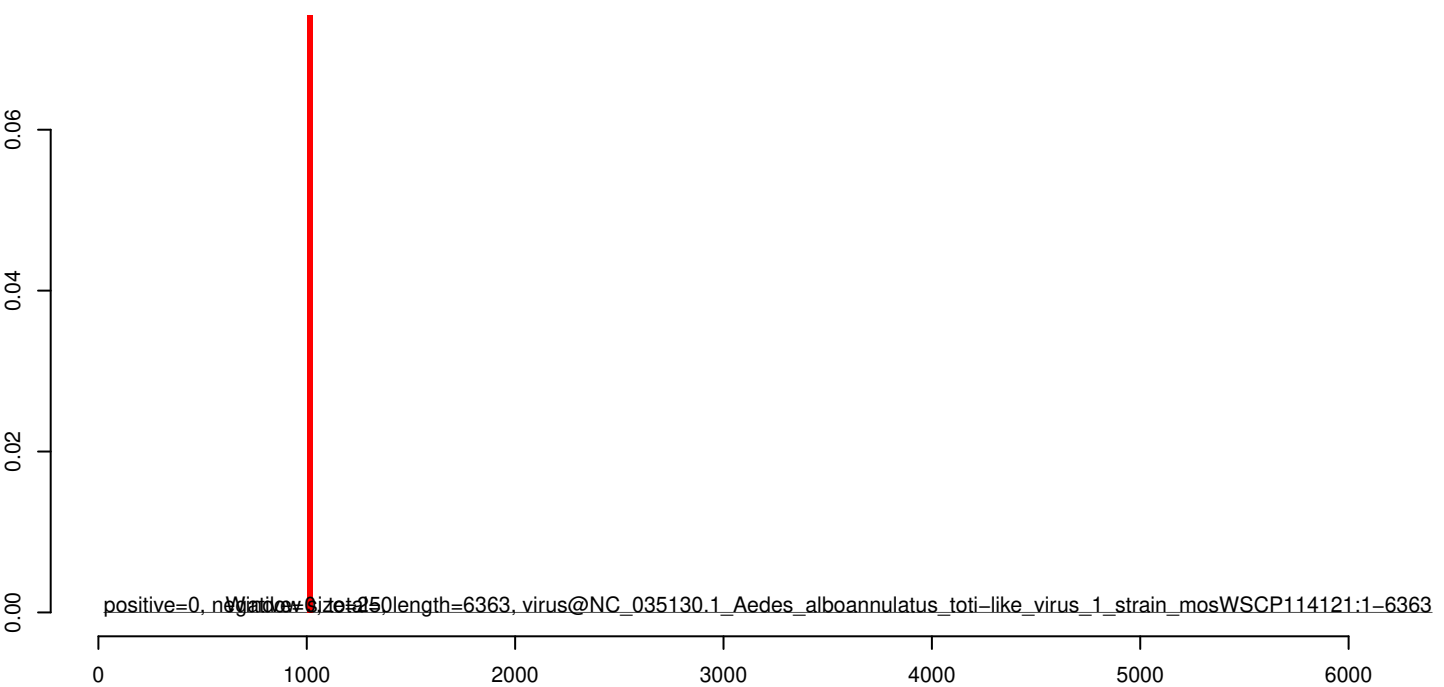
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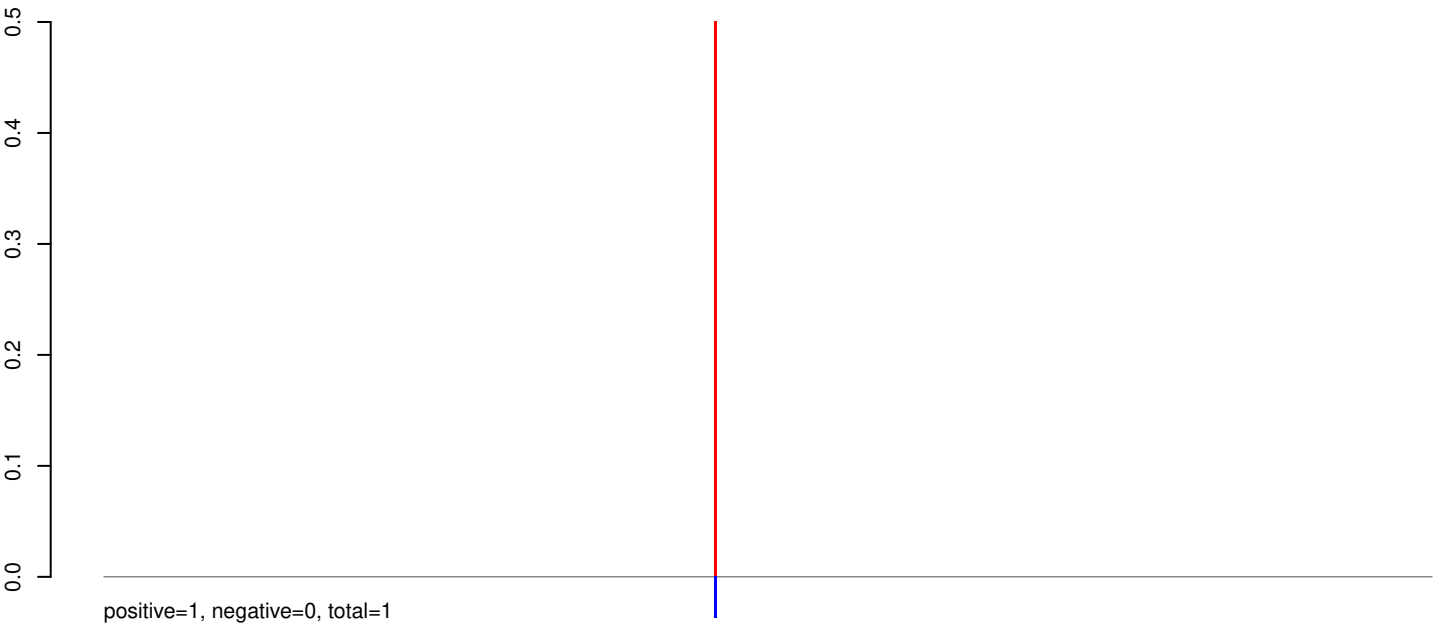
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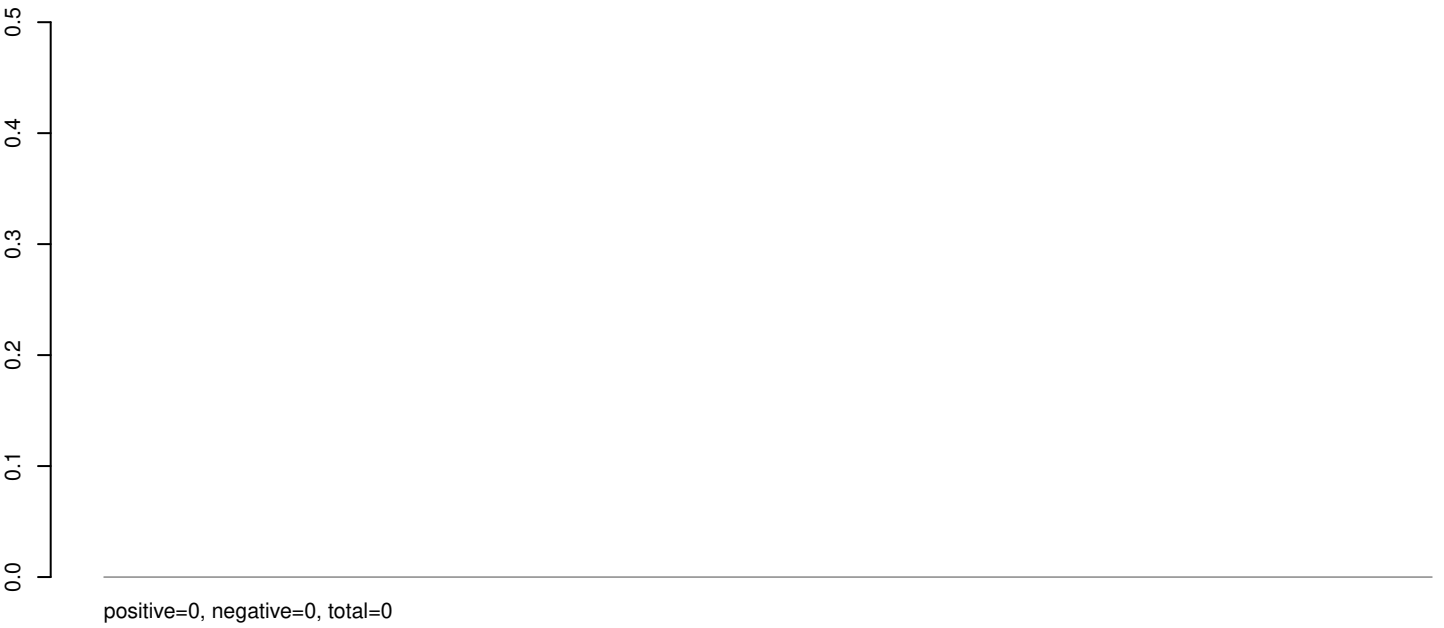
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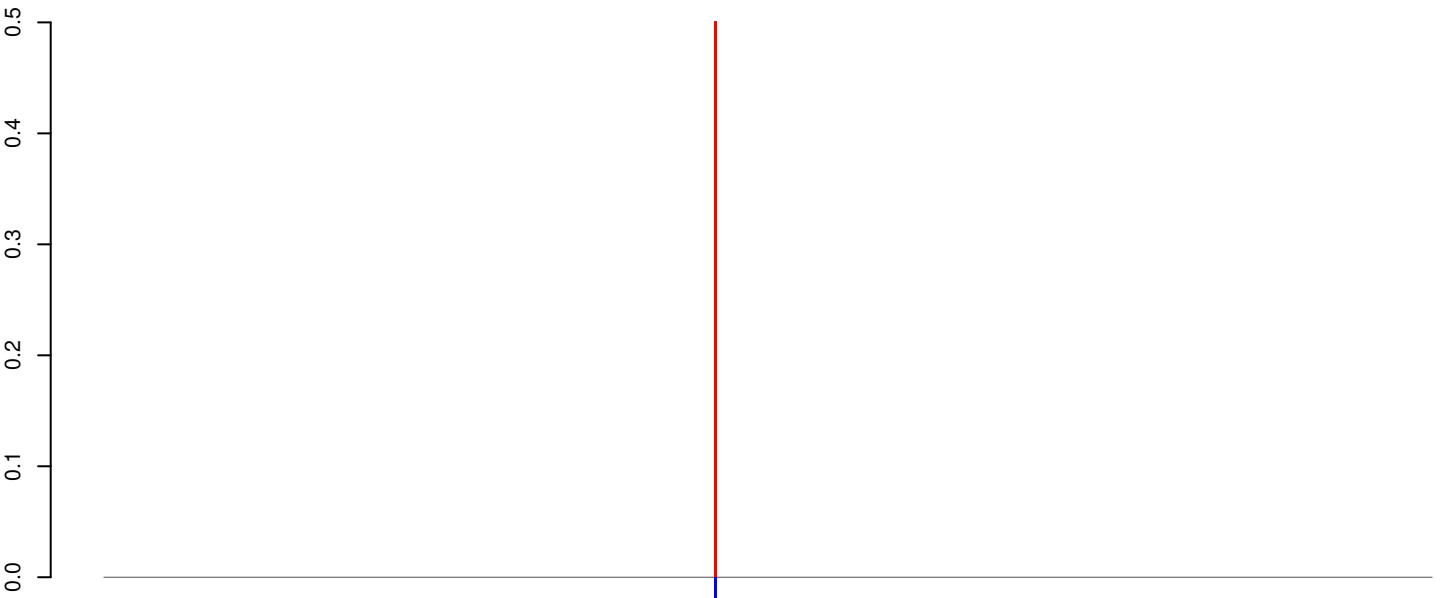
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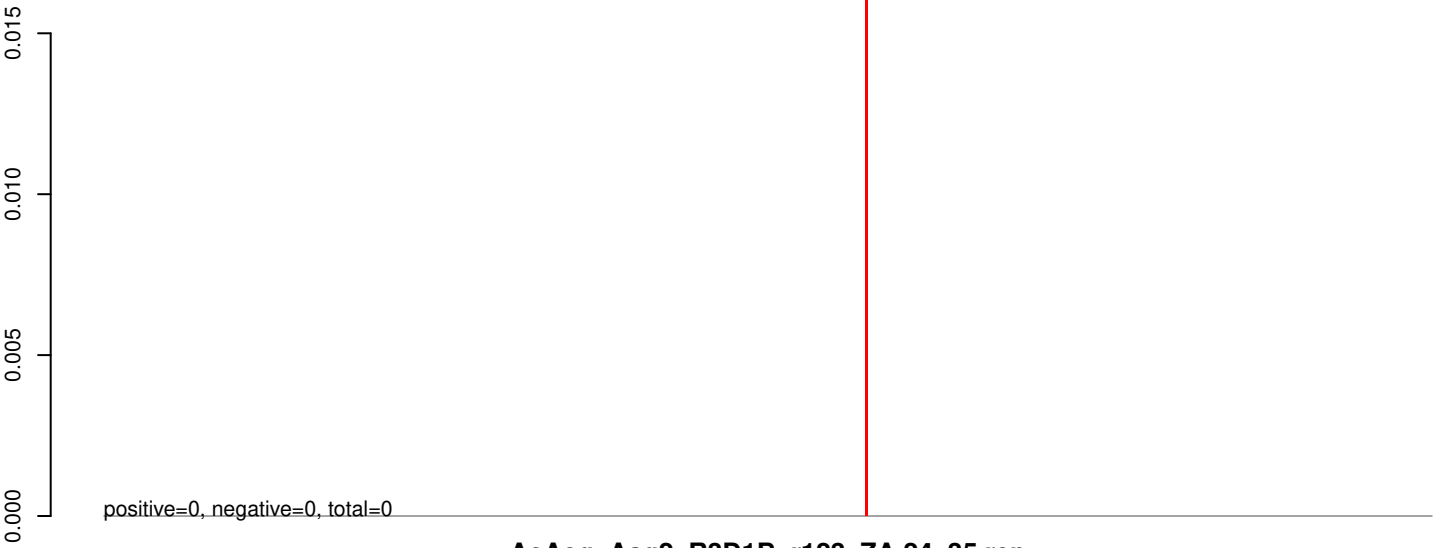
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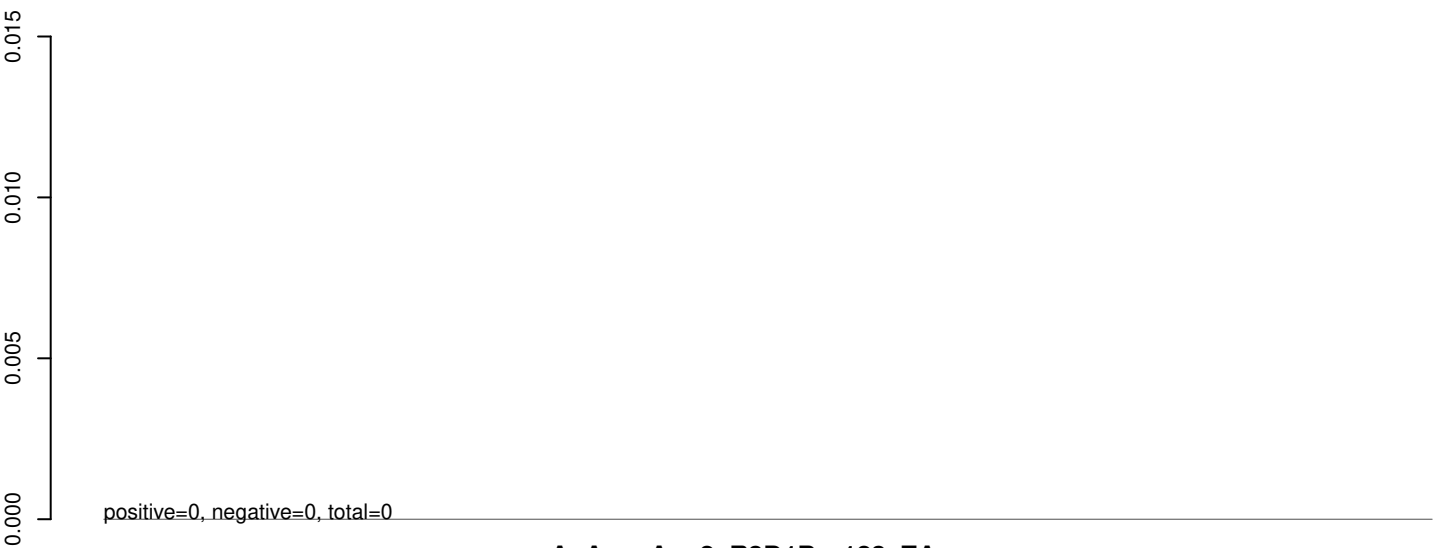
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0 2000 4000 6000 8000 10000

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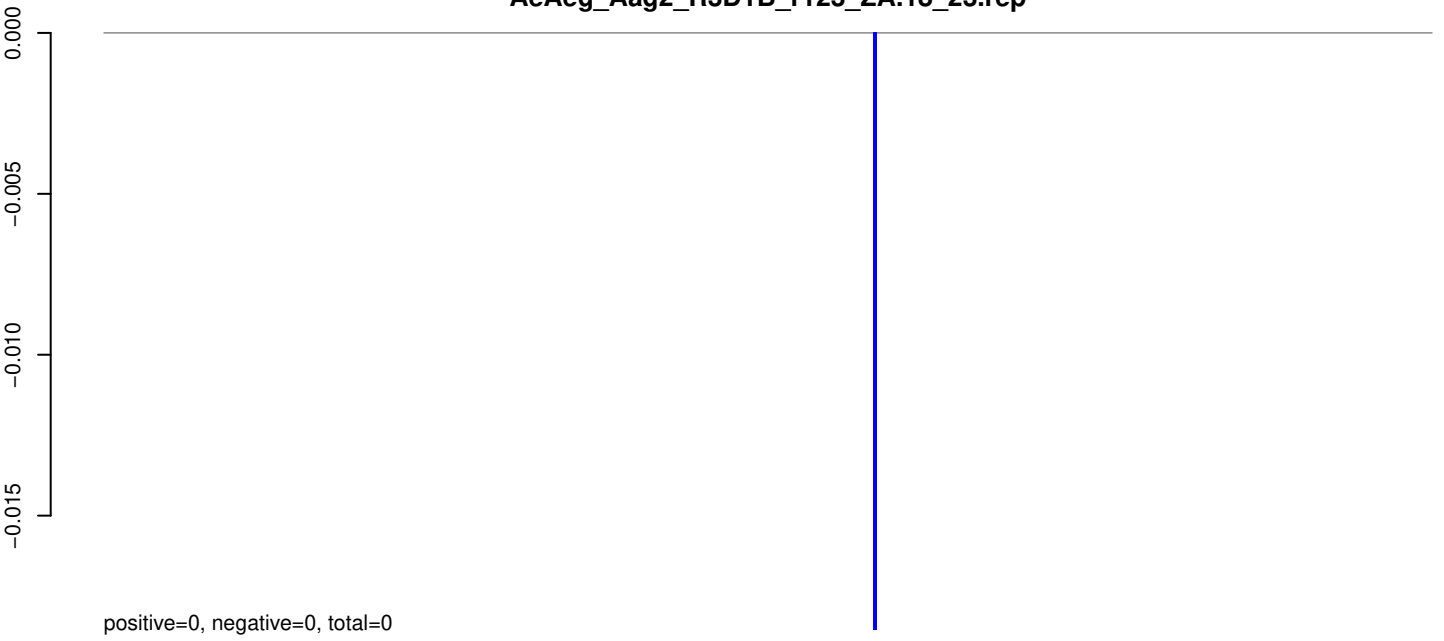
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AeAeg_Aag2_R3D1B_r123_ZA.rep



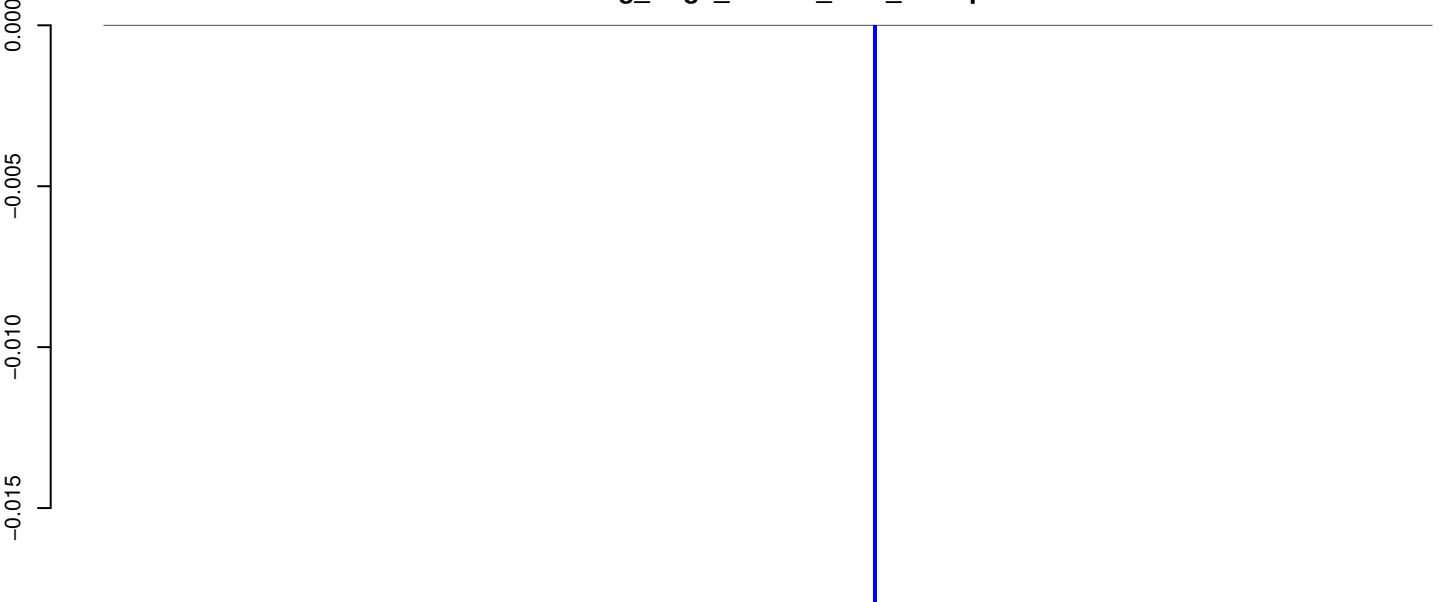
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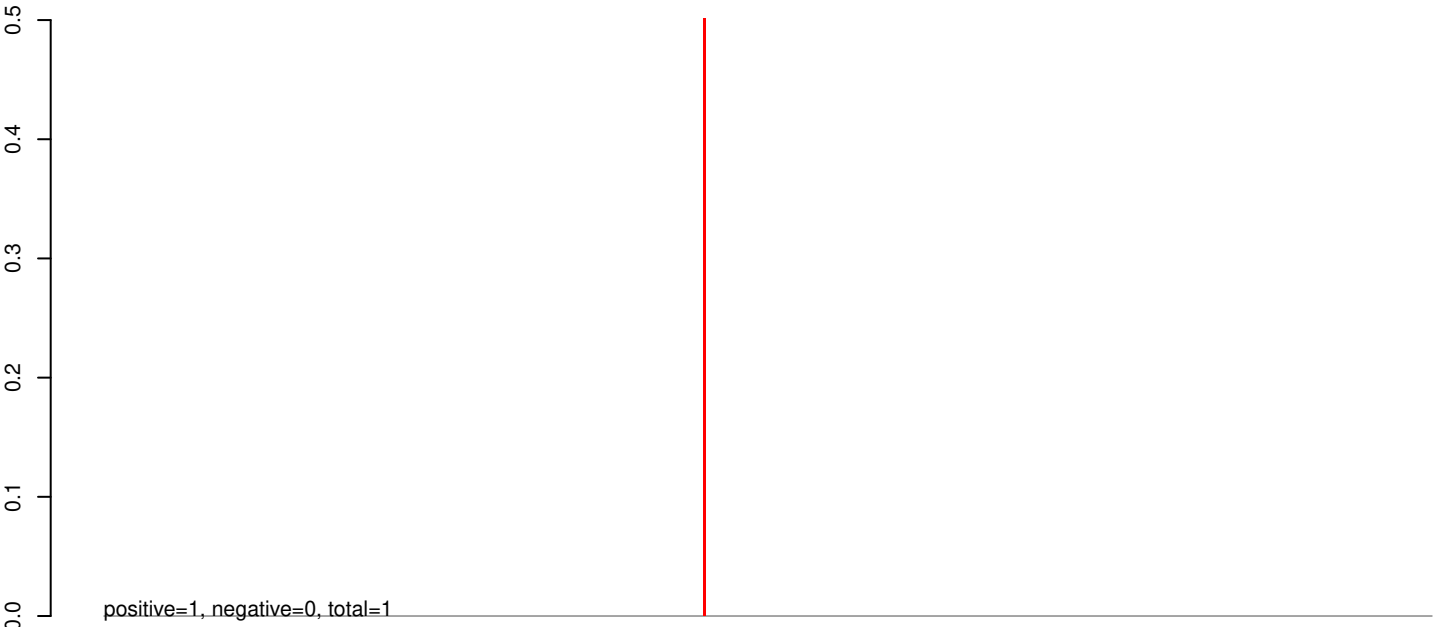
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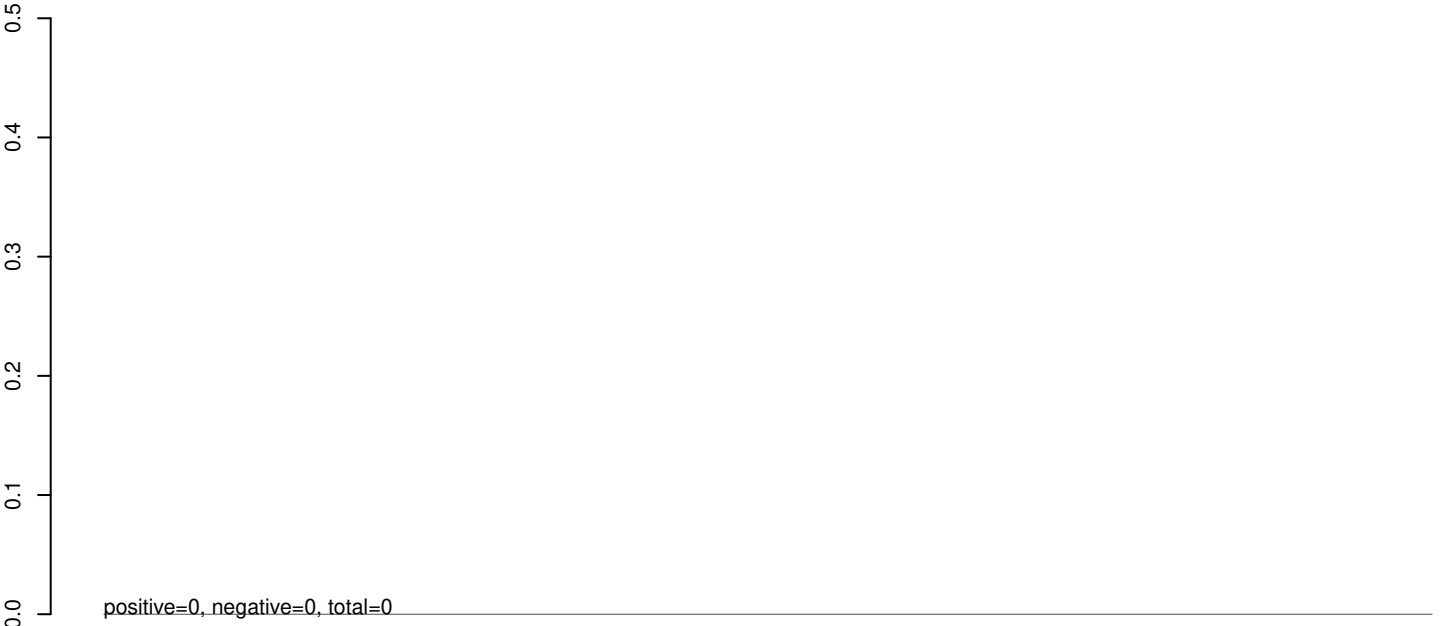
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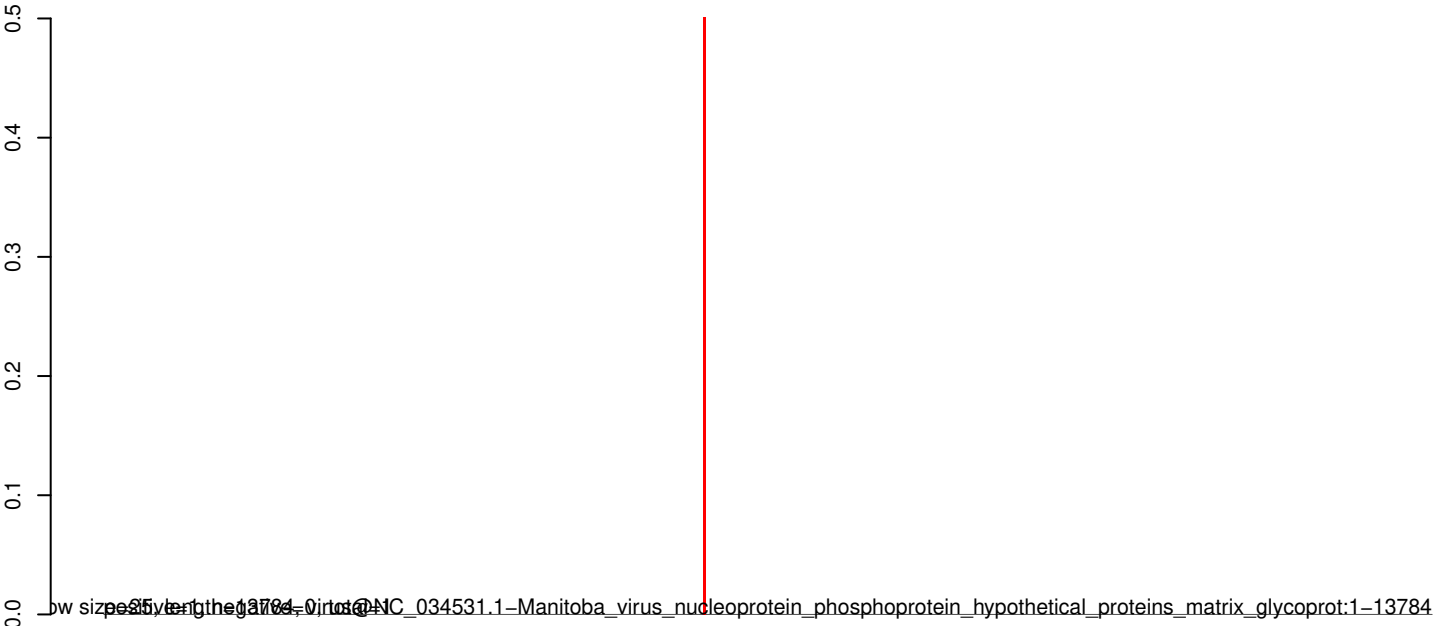
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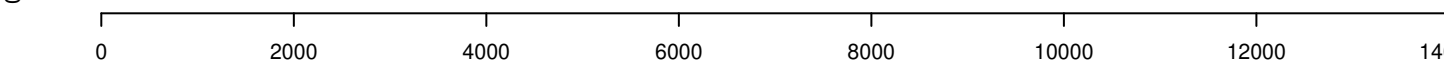
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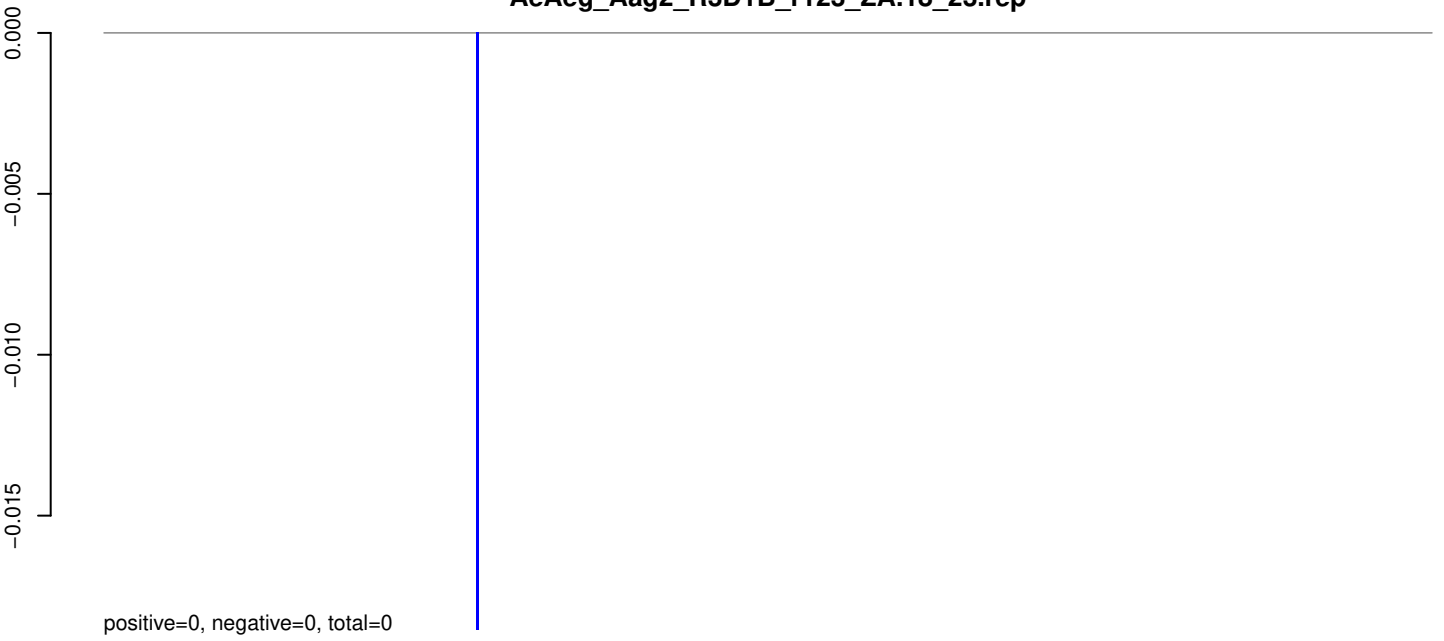
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bow size 25, length 13784, virus NC_034531.1-Manitoba_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycoprot:1-13784



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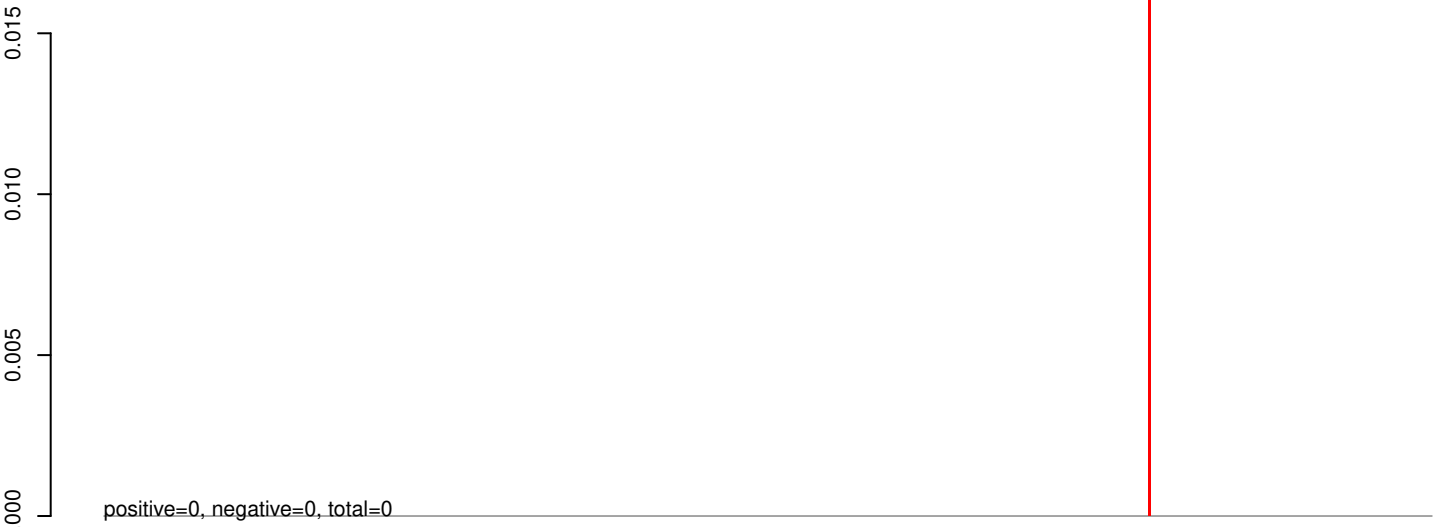
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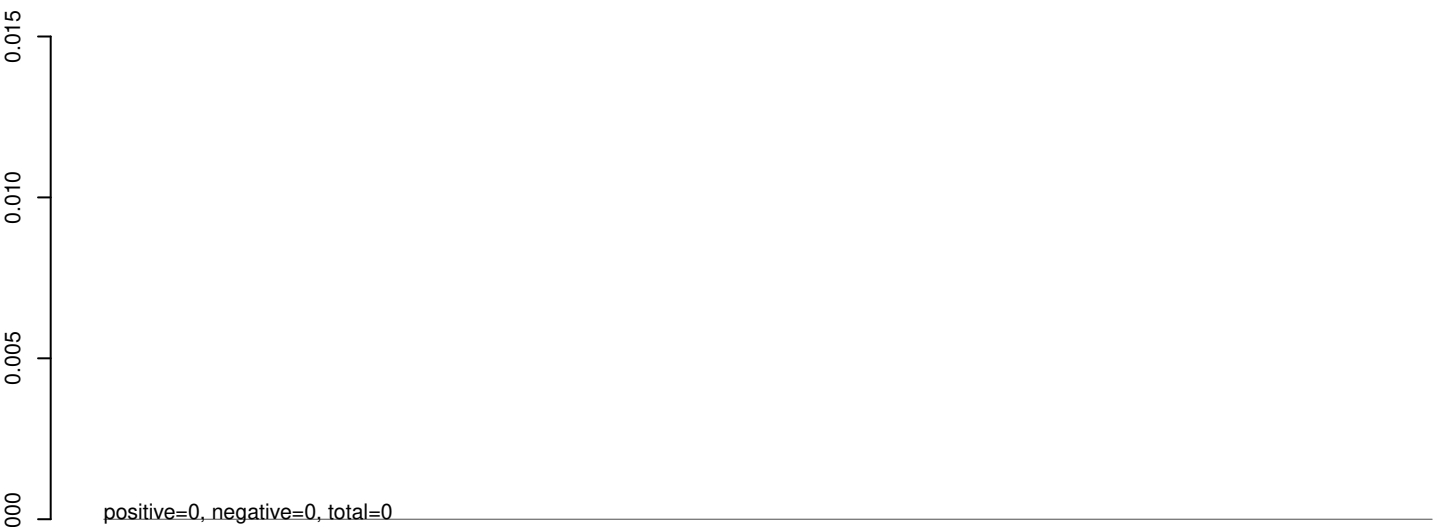
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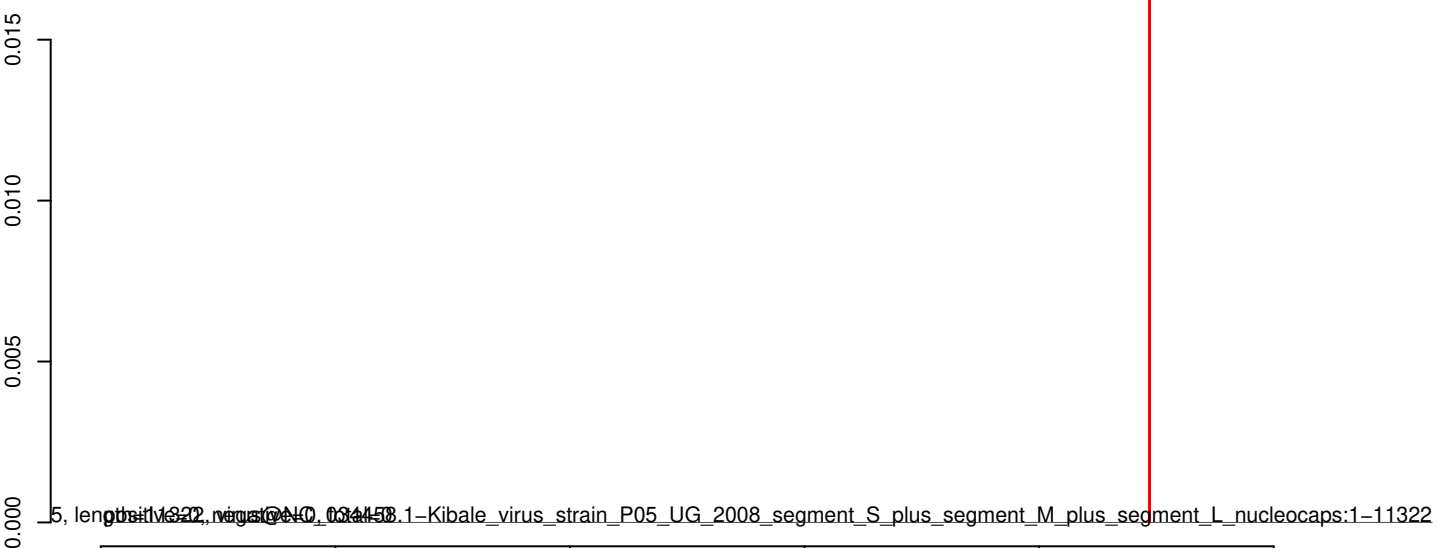
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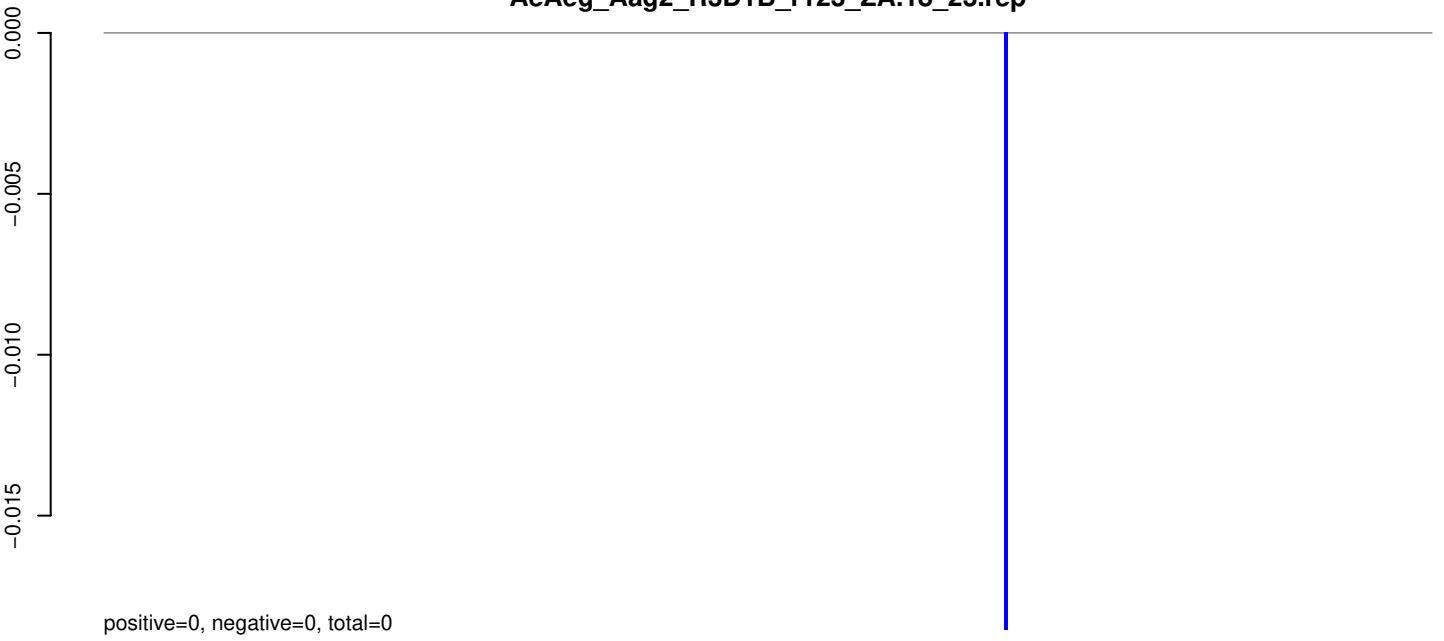


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0 2000 4000 6000 8000 10000

AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



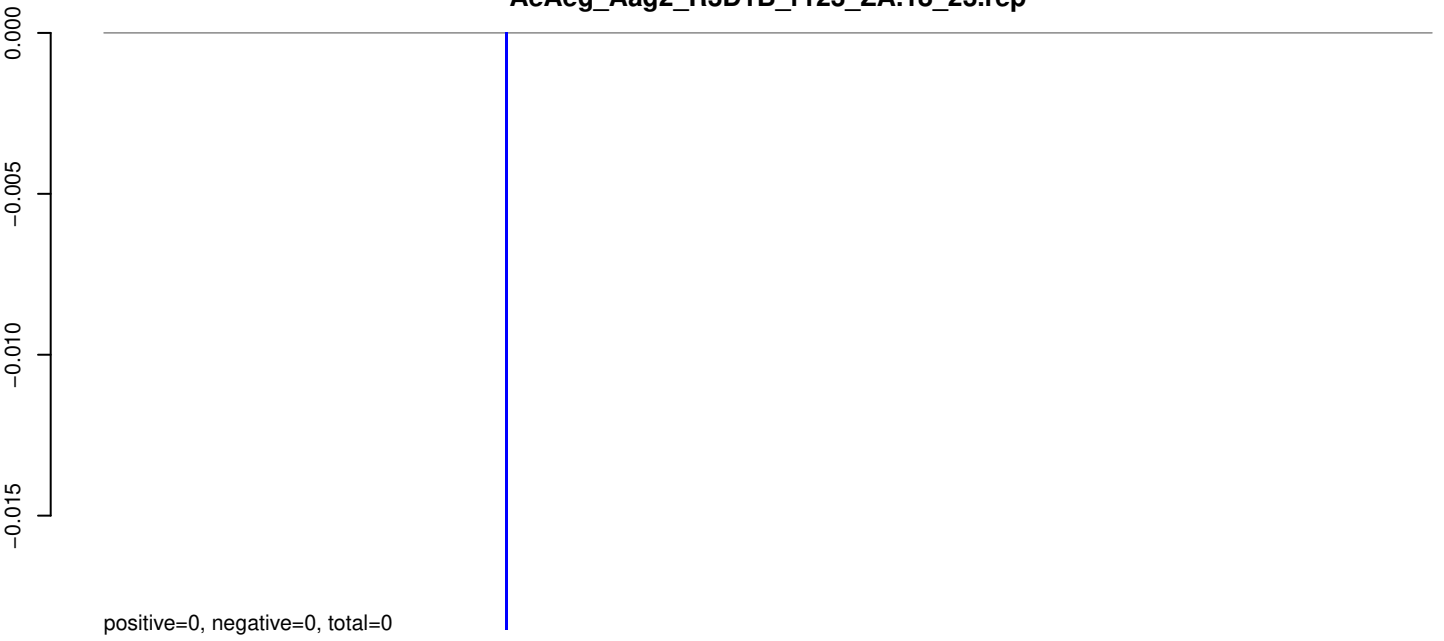
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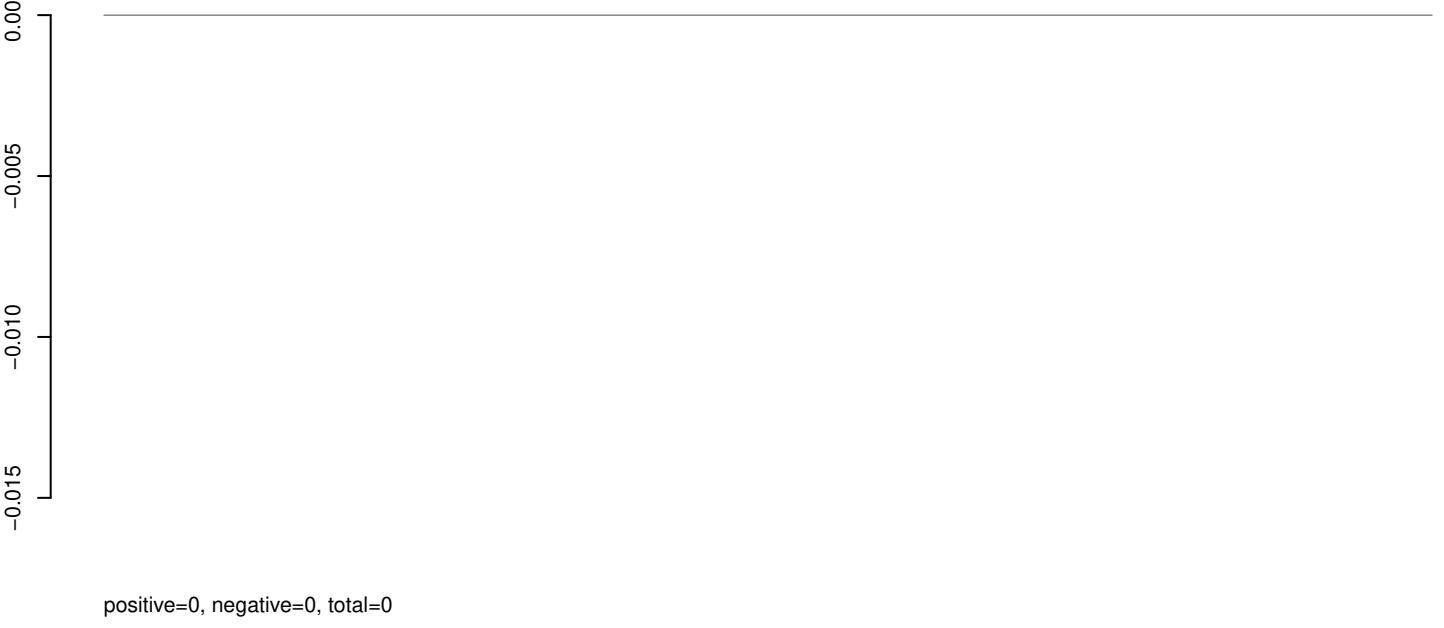
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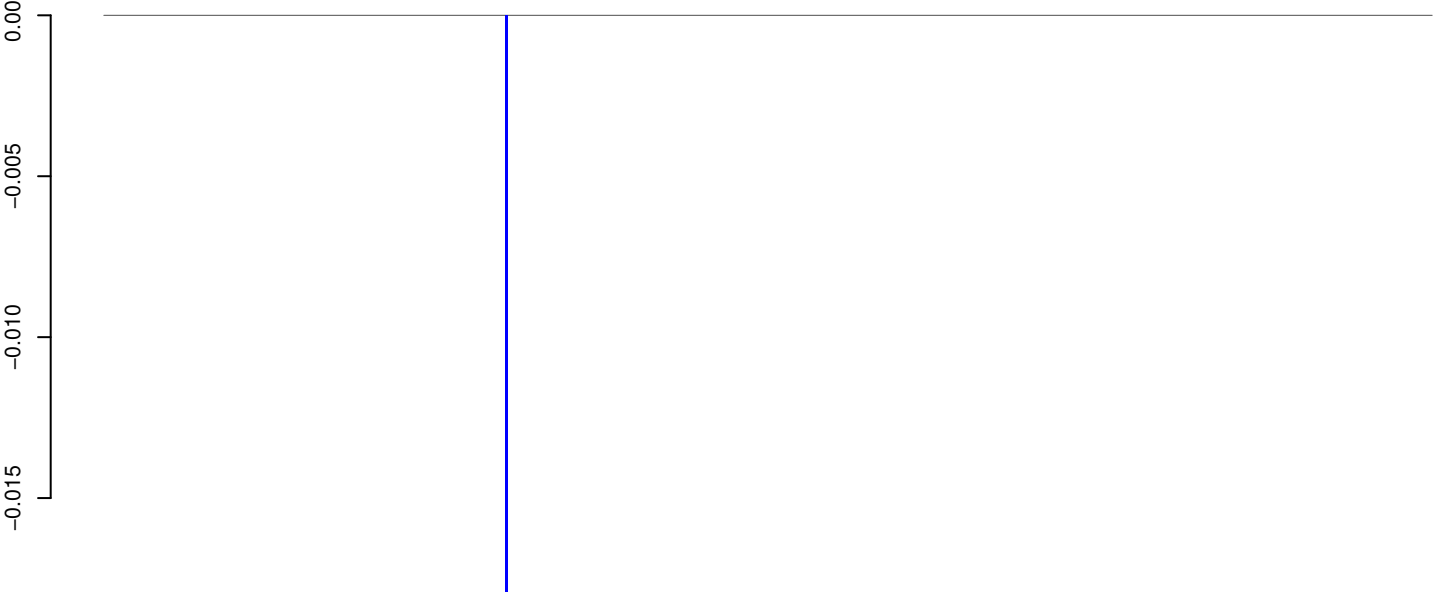
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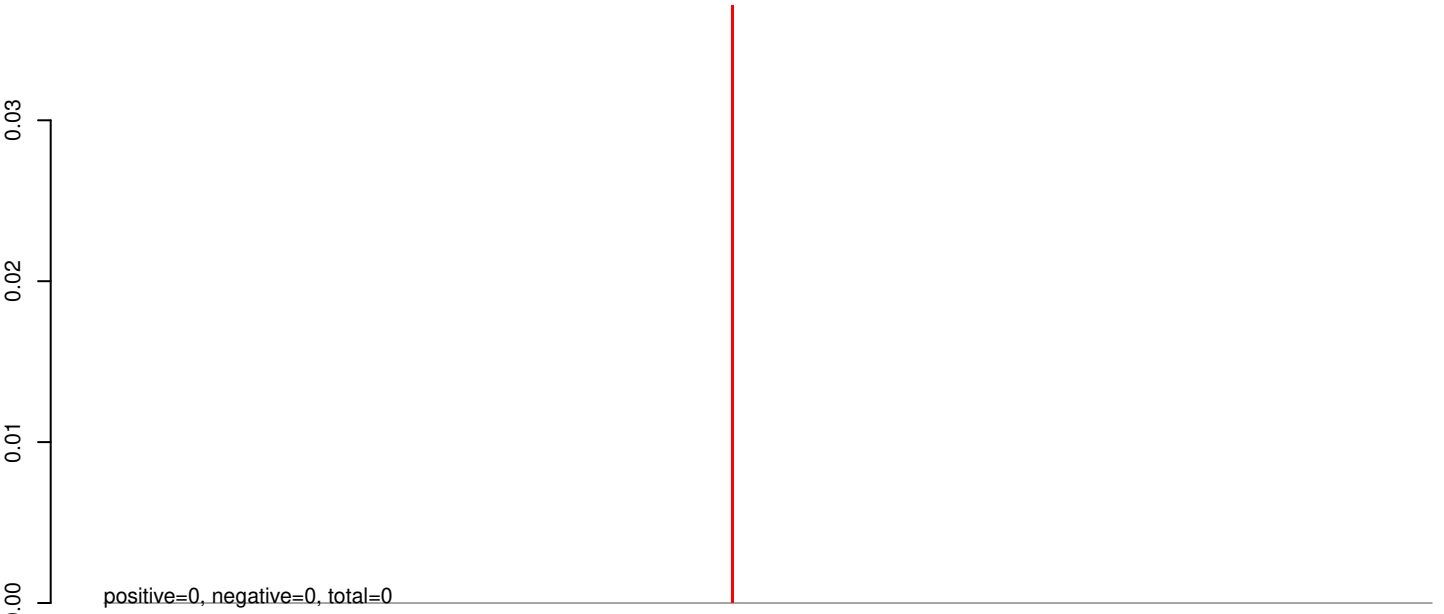
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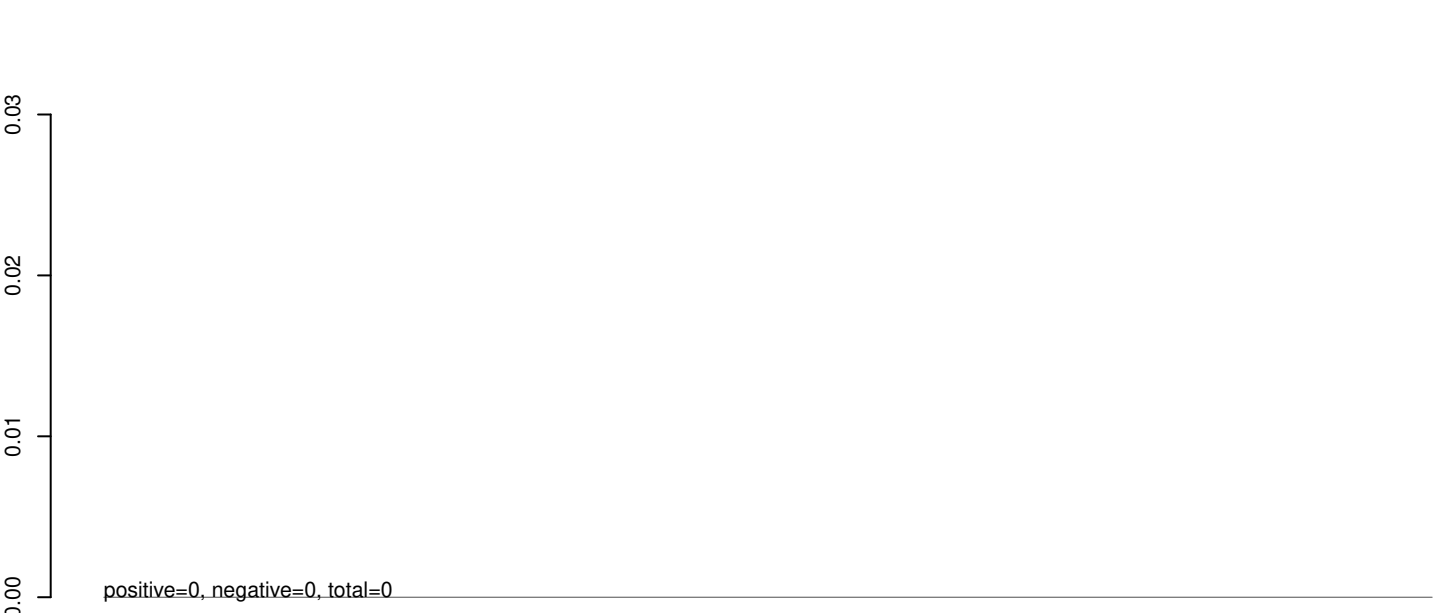
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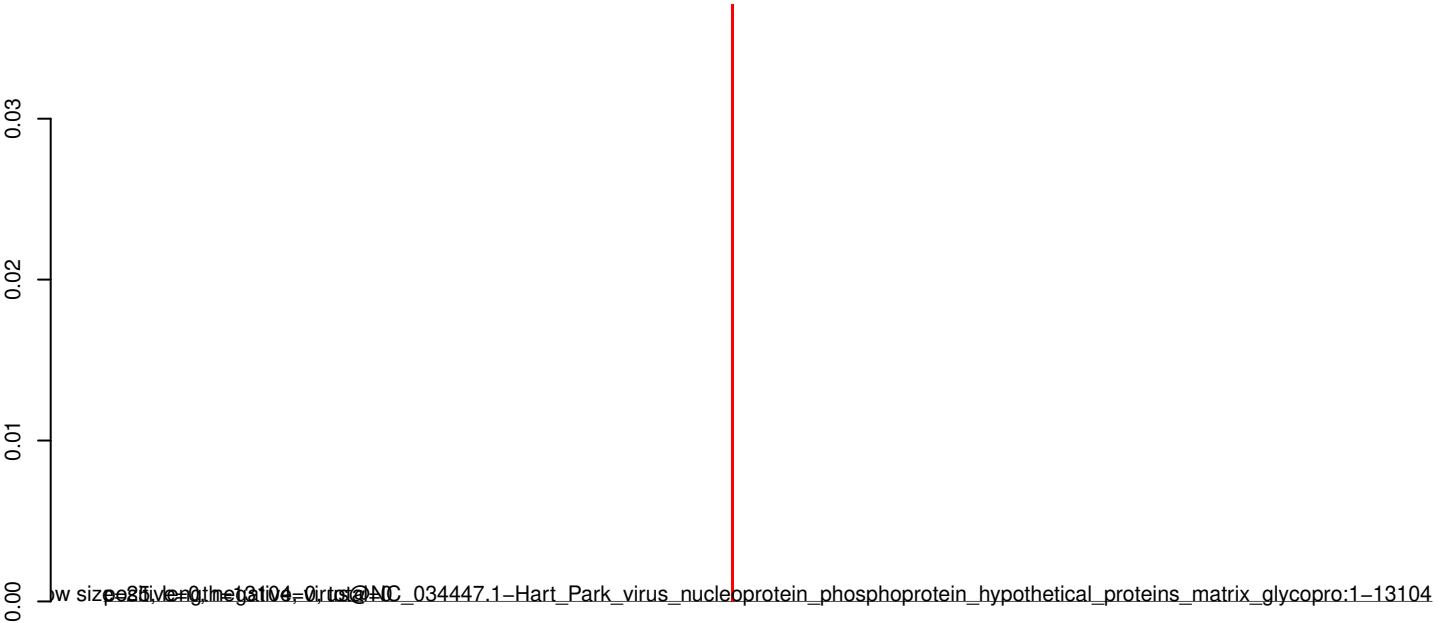
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



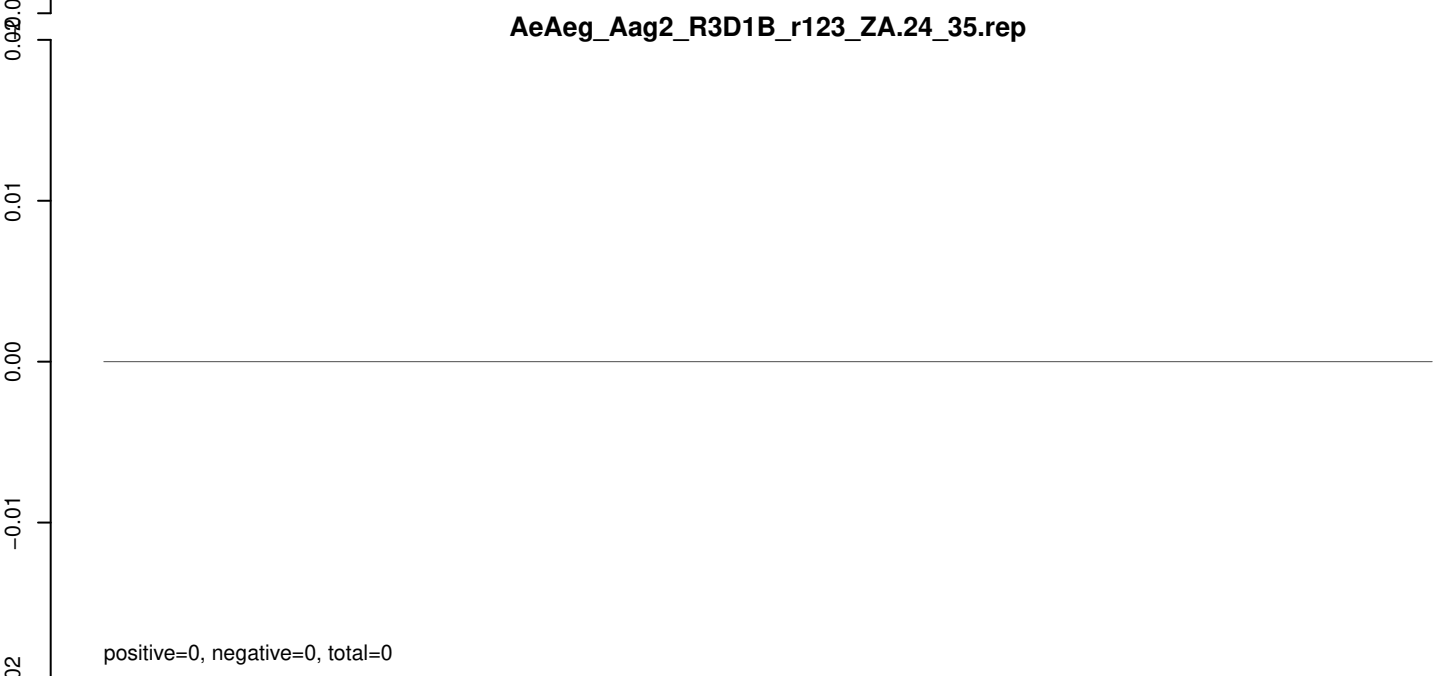
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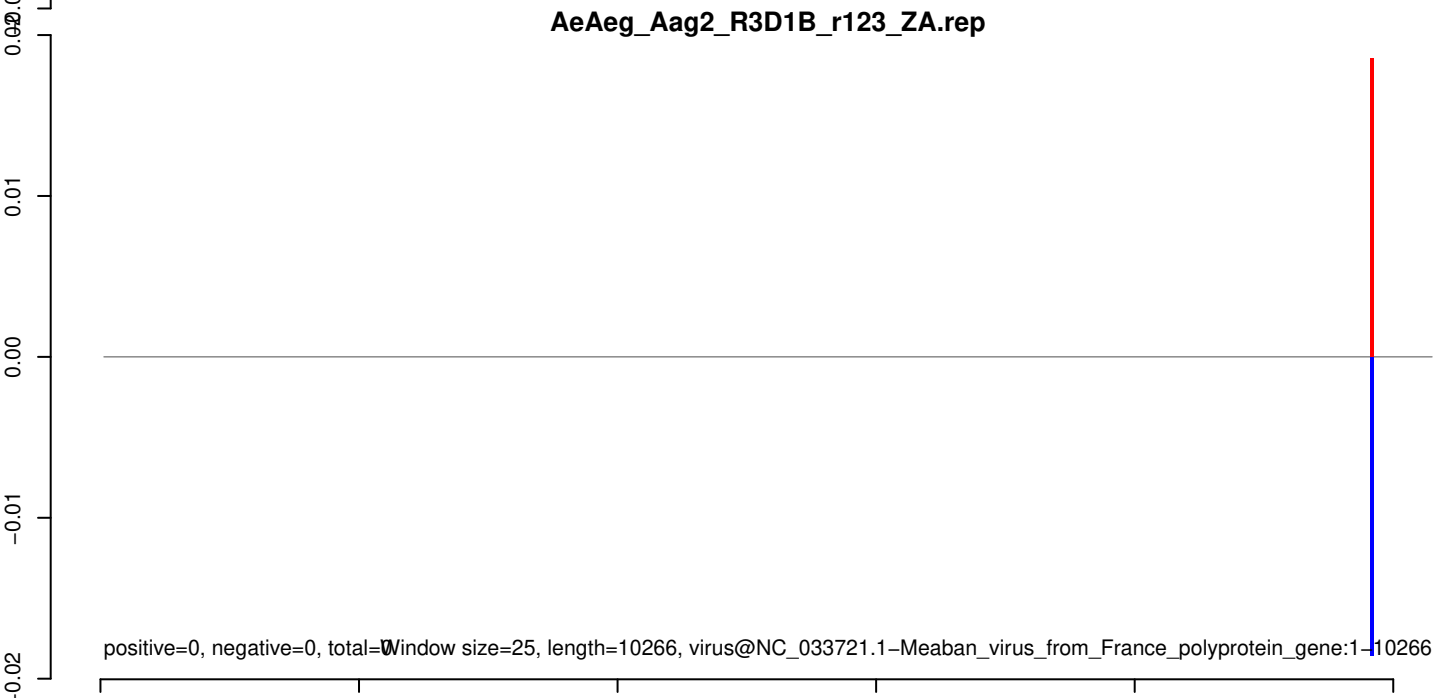
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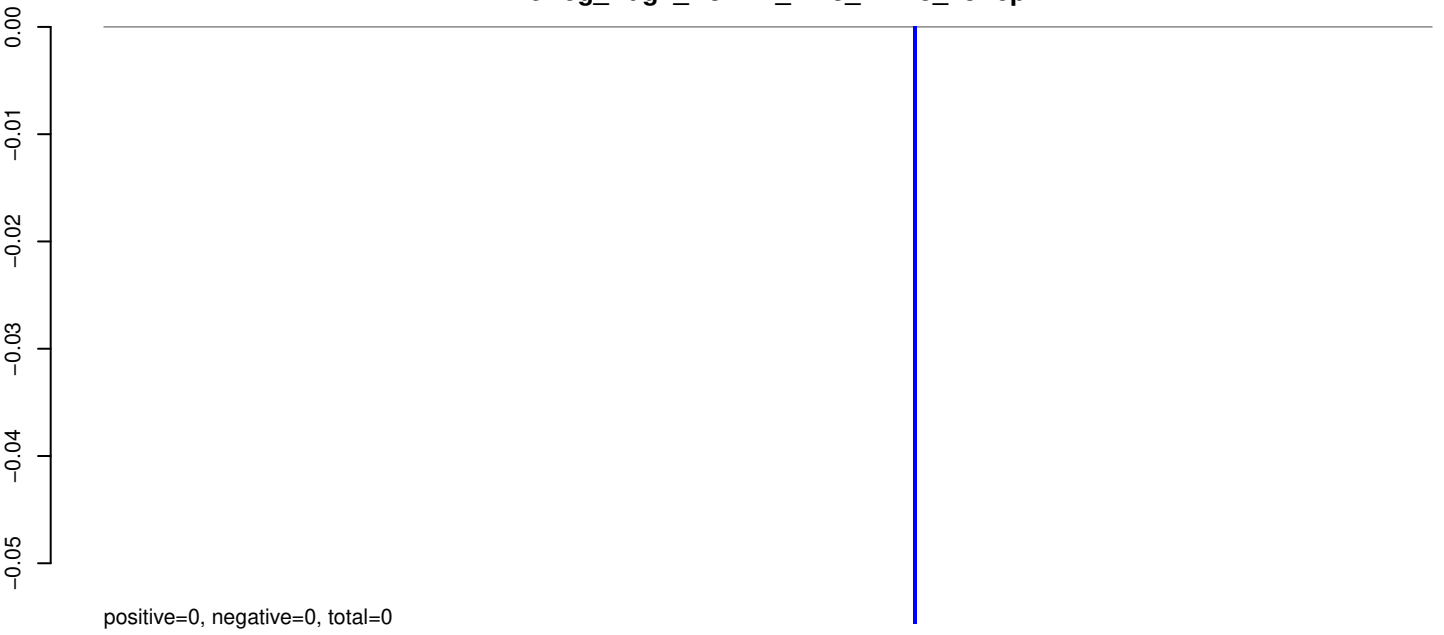
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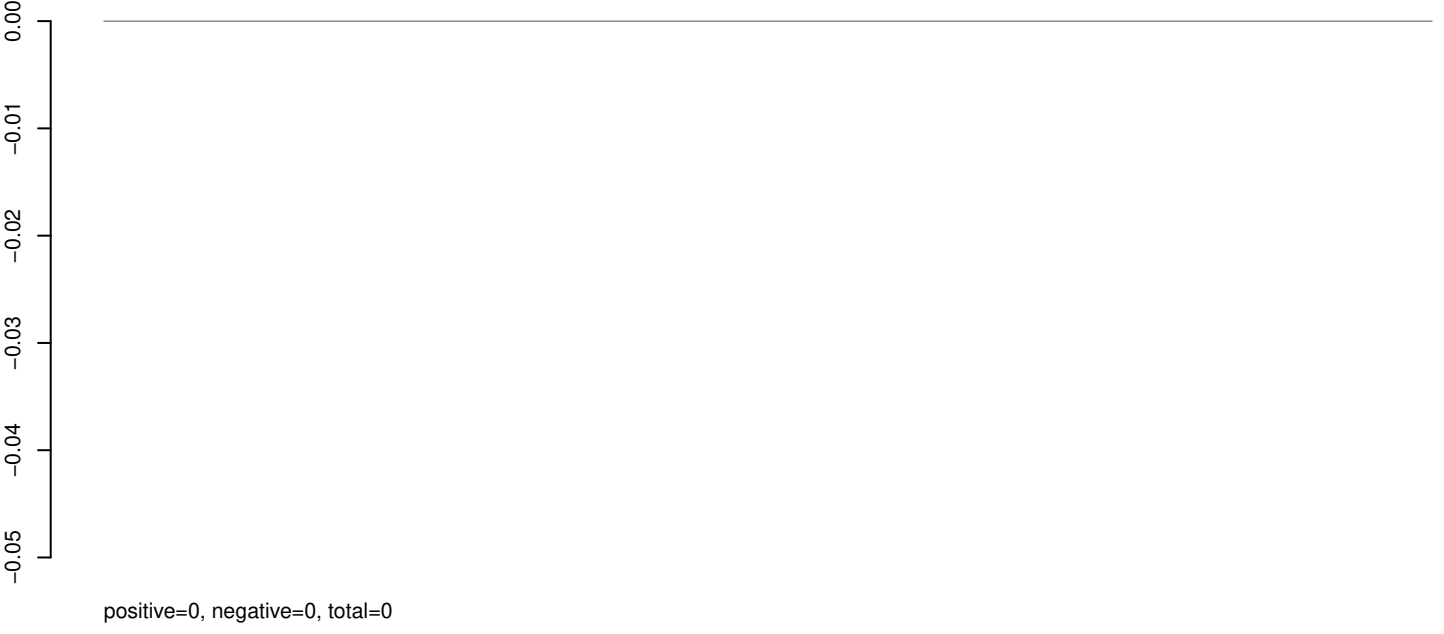
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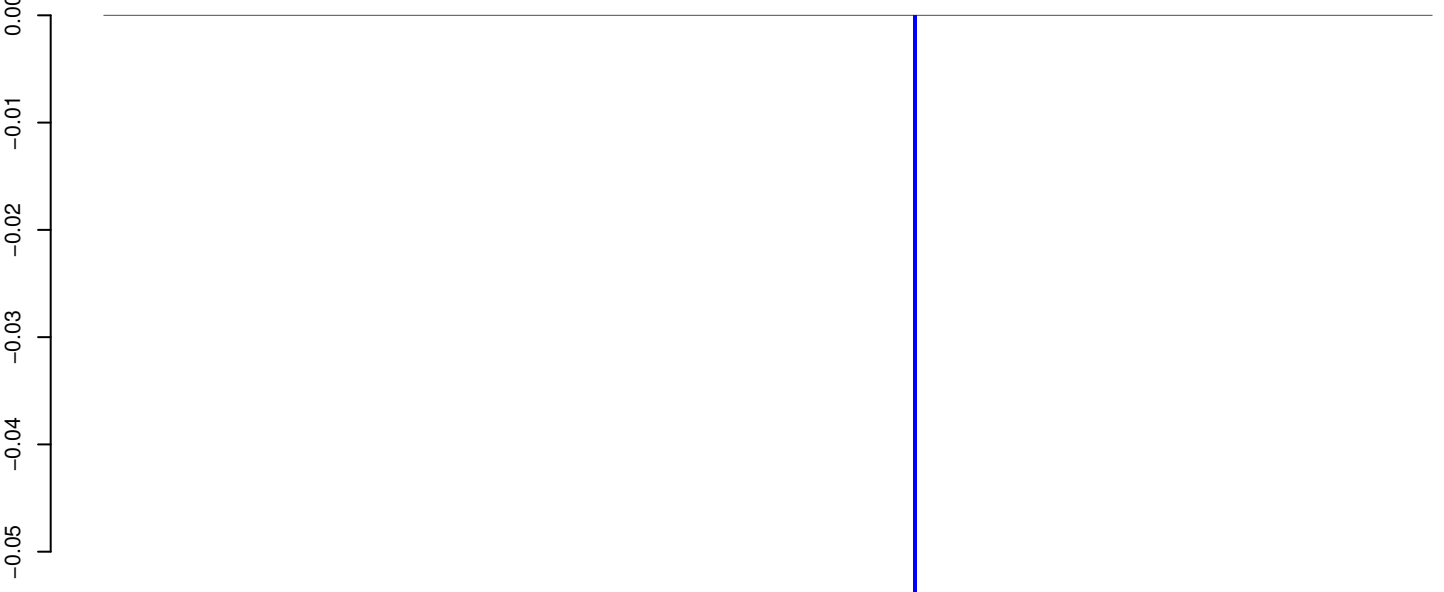
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



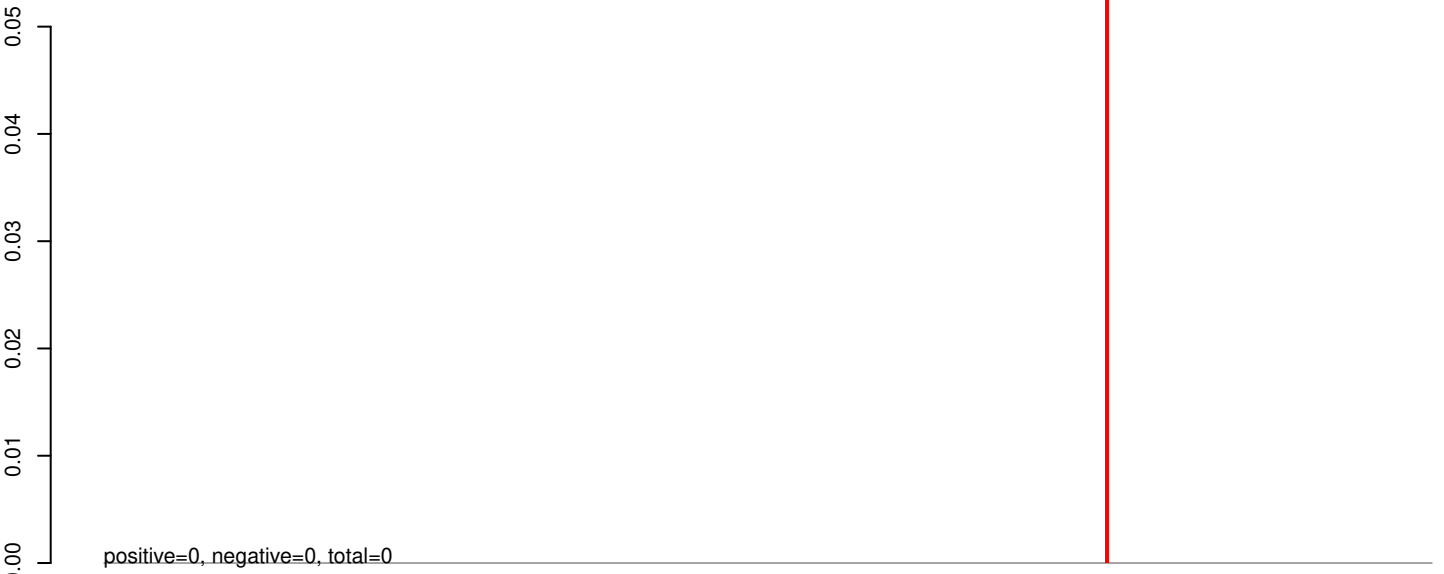
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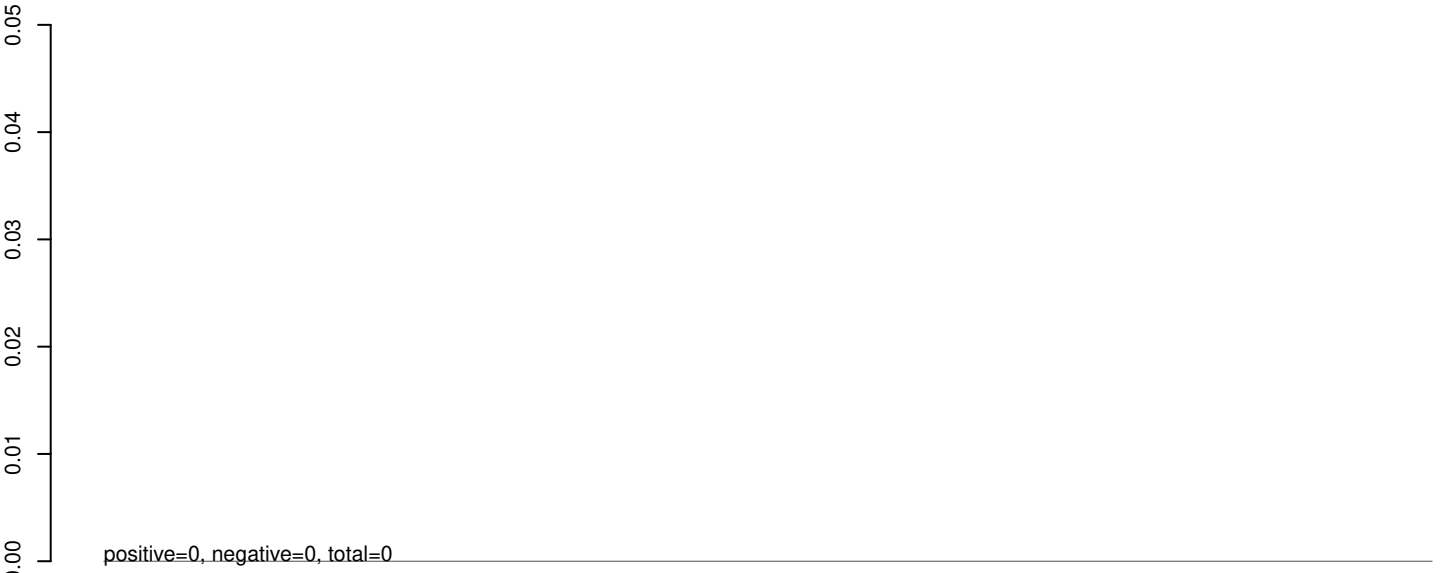
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AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep

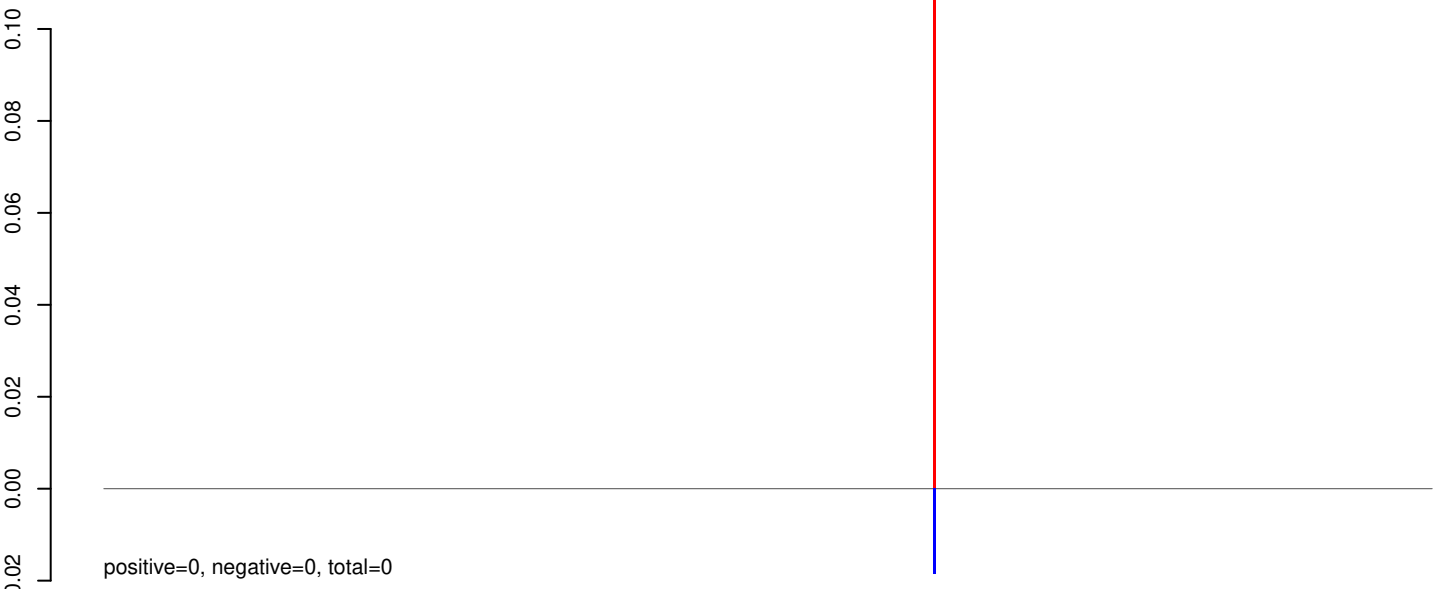


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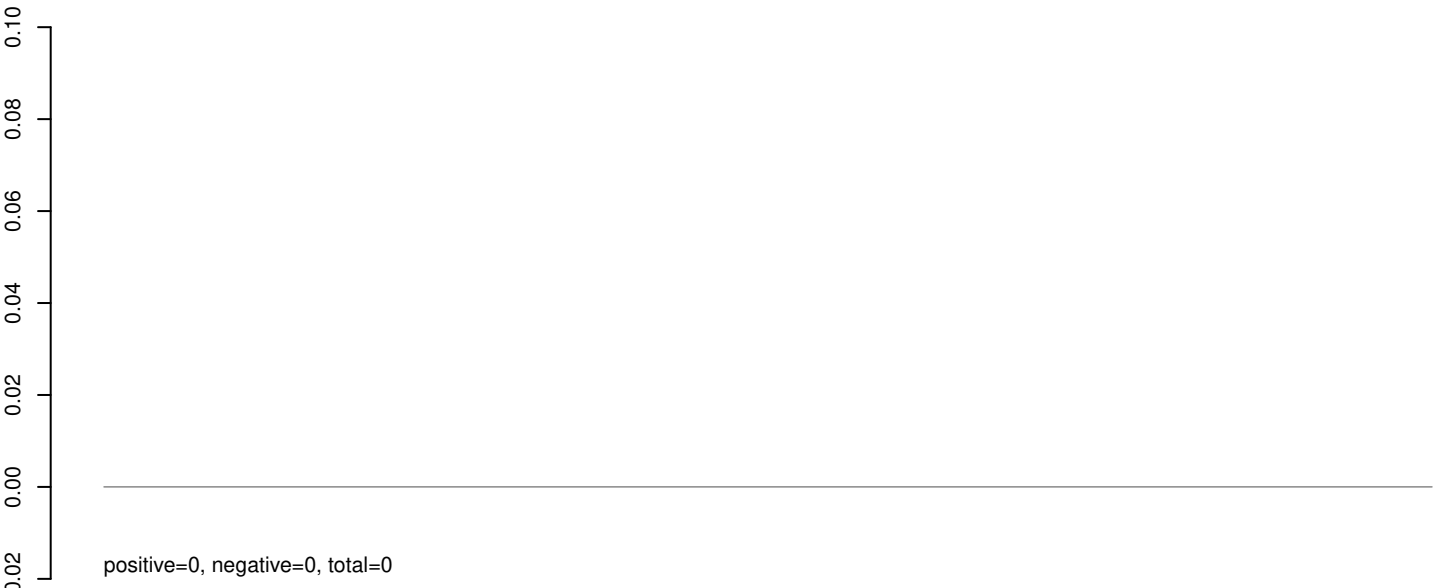


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AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



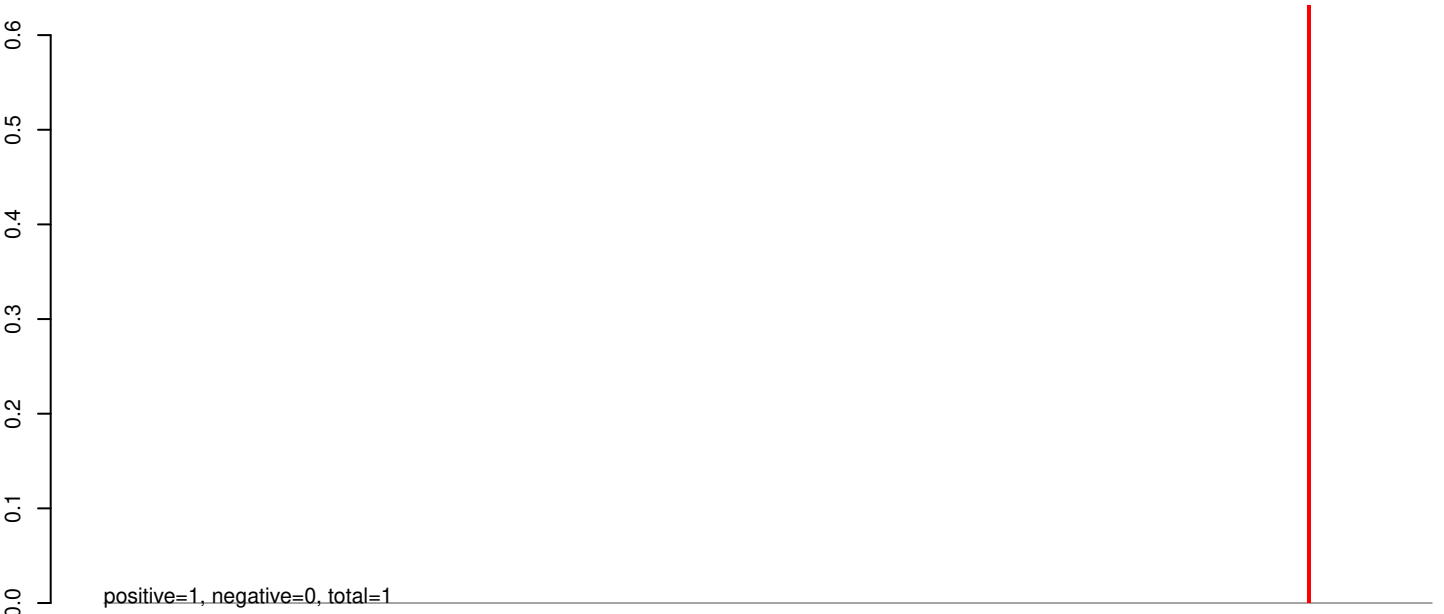
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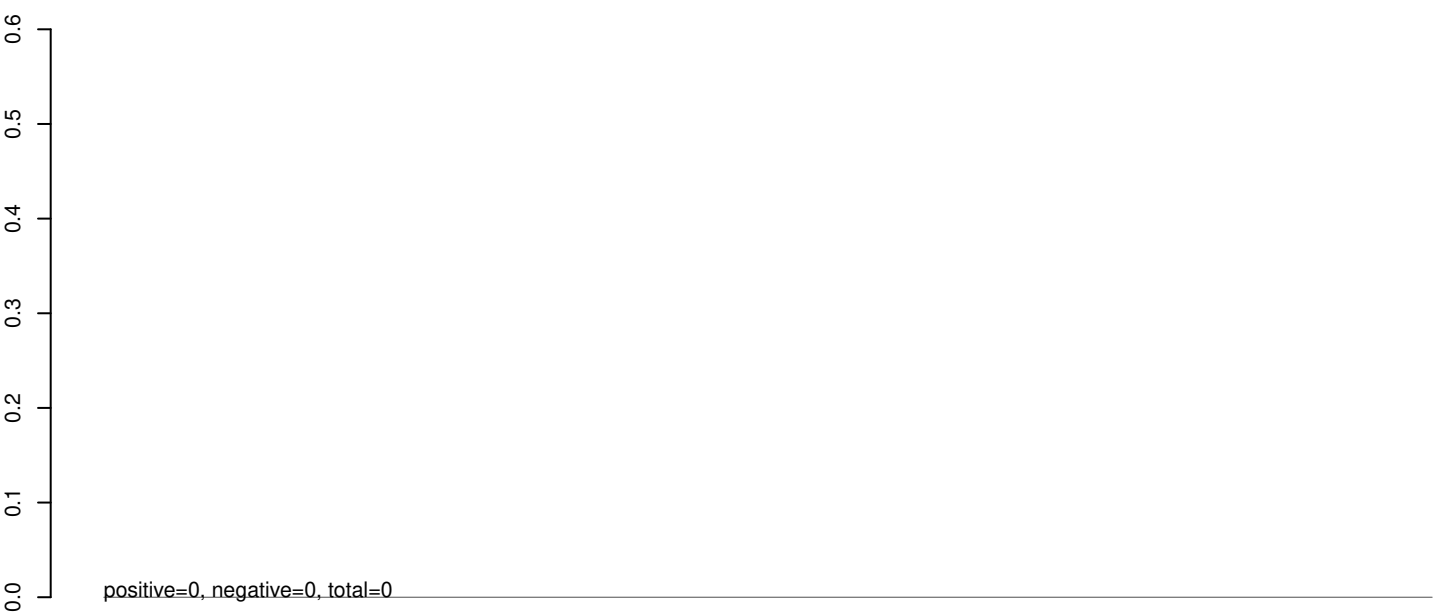
Xincheng_Mosquito_Virus_strain_XC1-6_ORF1_ORF1_ORF2_ORF2_glycoprotein_G_and_RNA-de:1-12774

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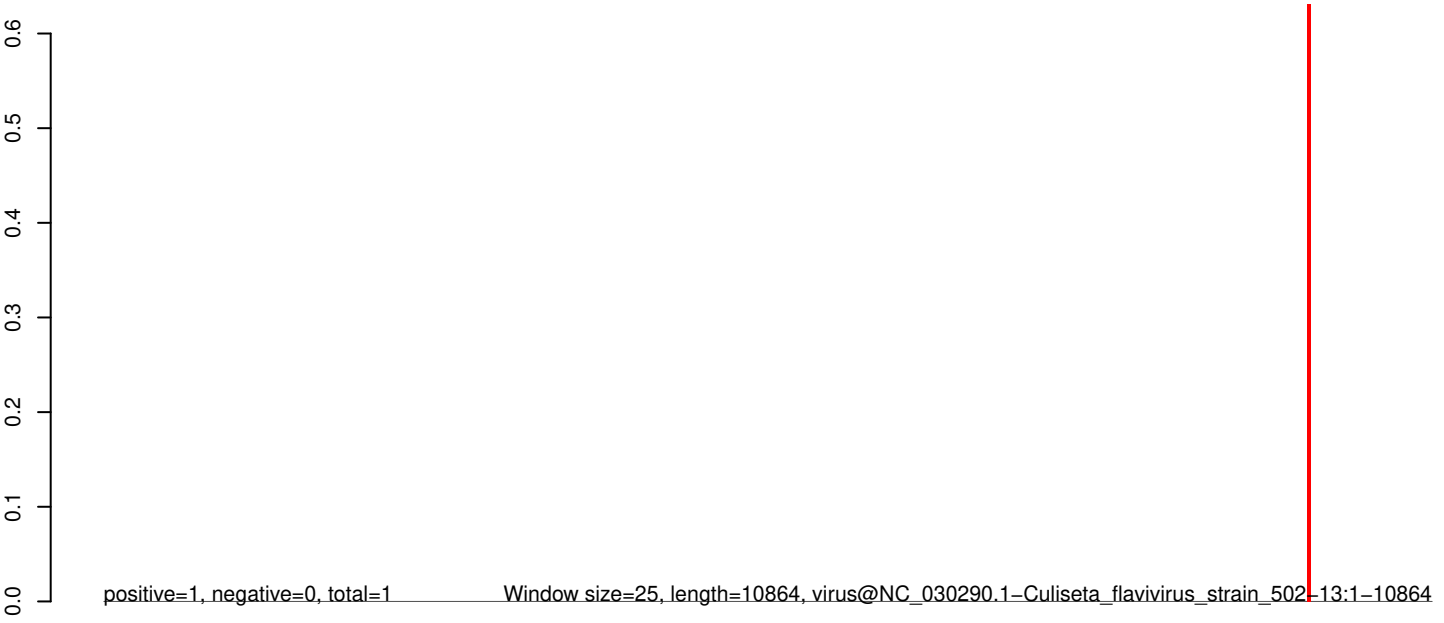
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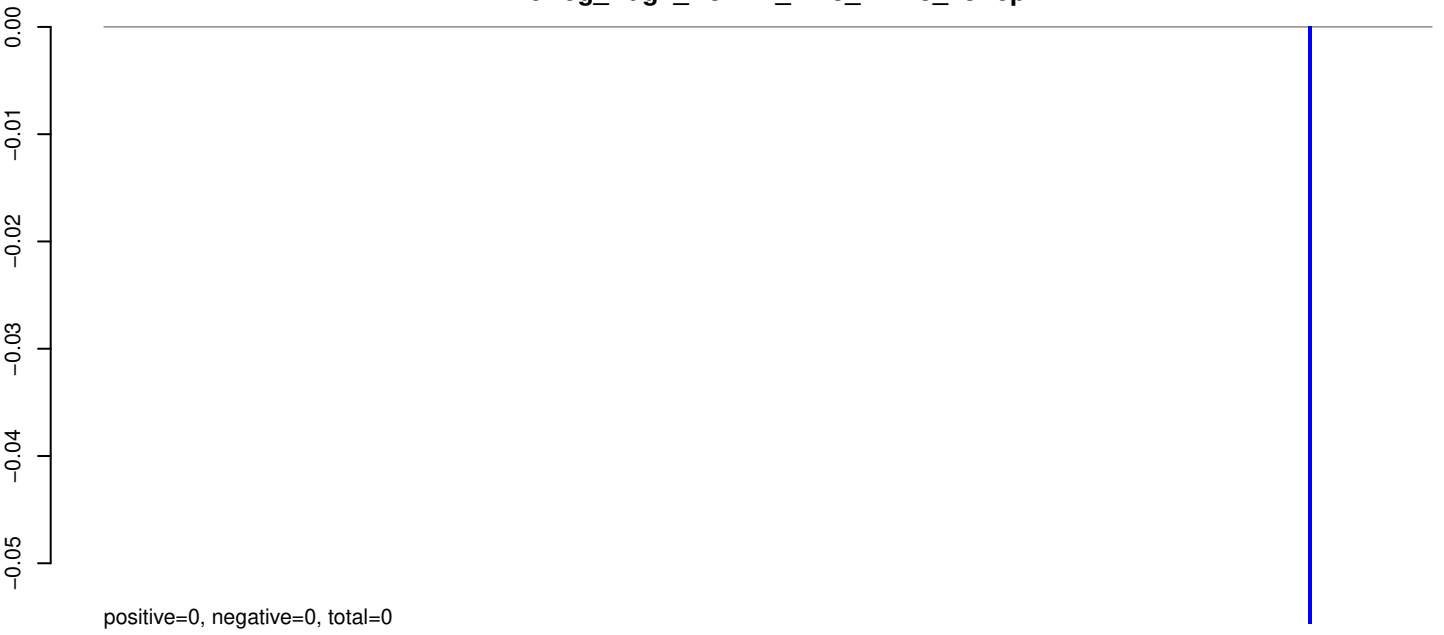


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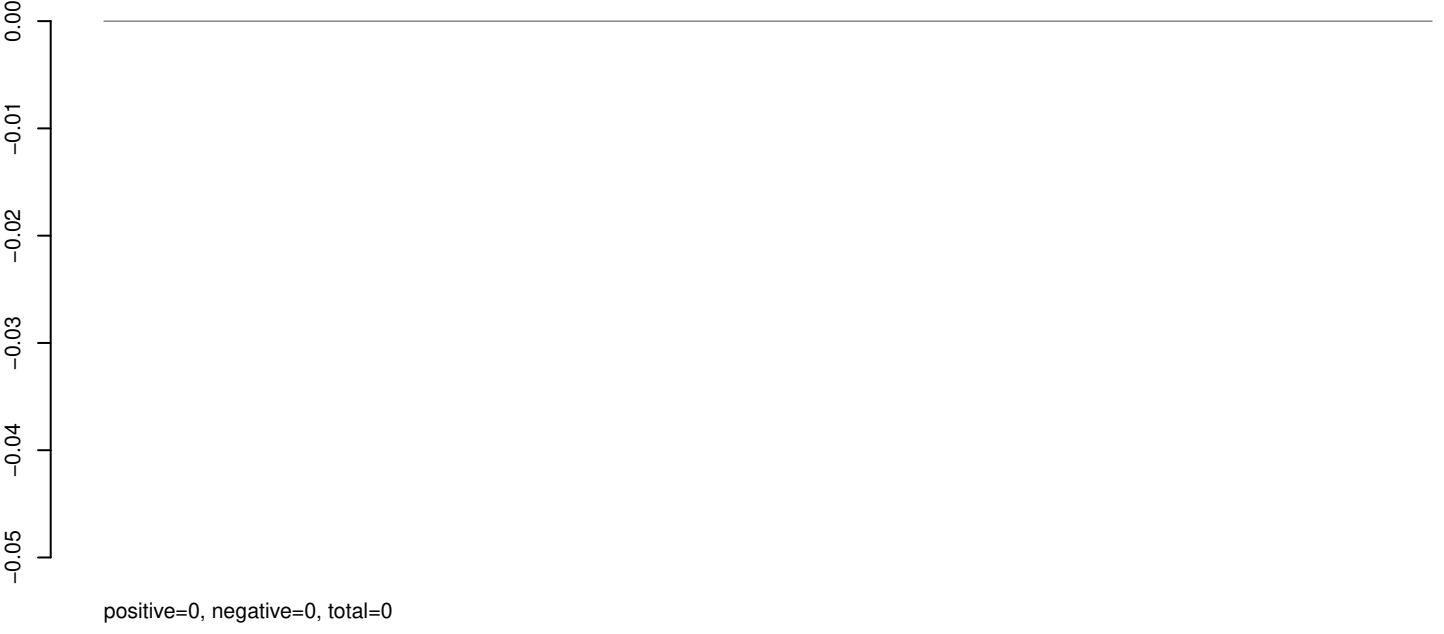
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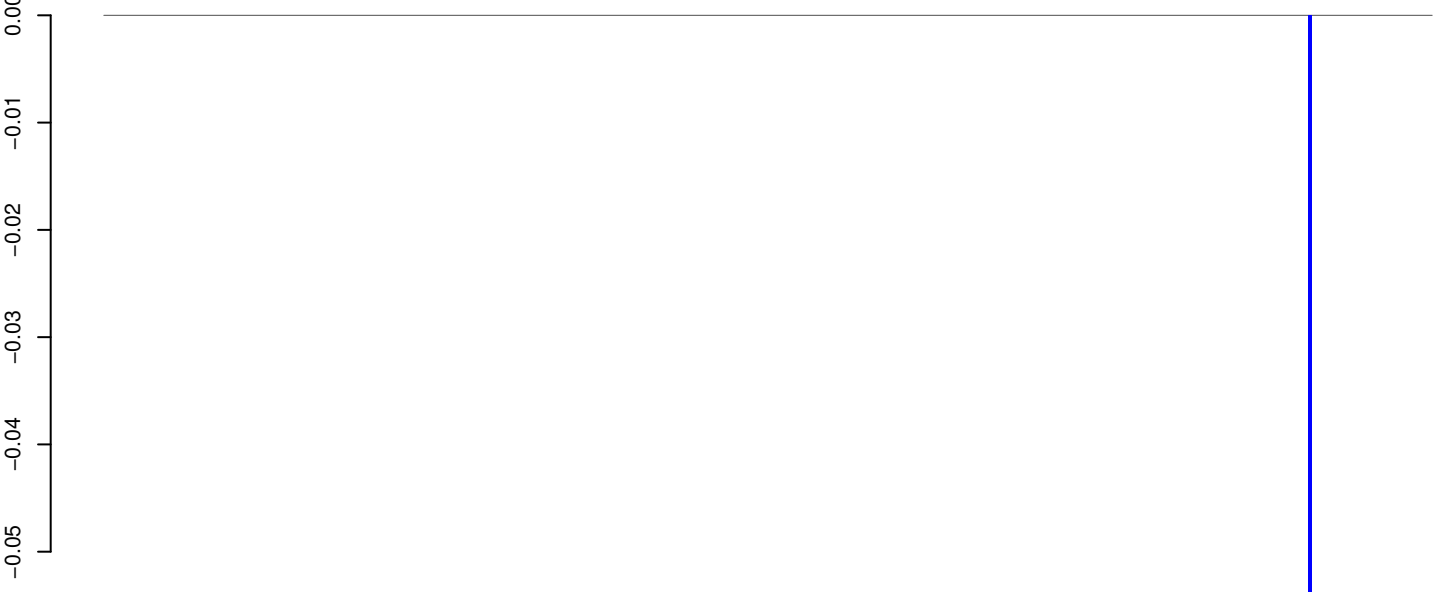
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AeAeg_Aag2_R3D1B_r123_ZA.24_35.rep



positive=0, negative=0, total=0

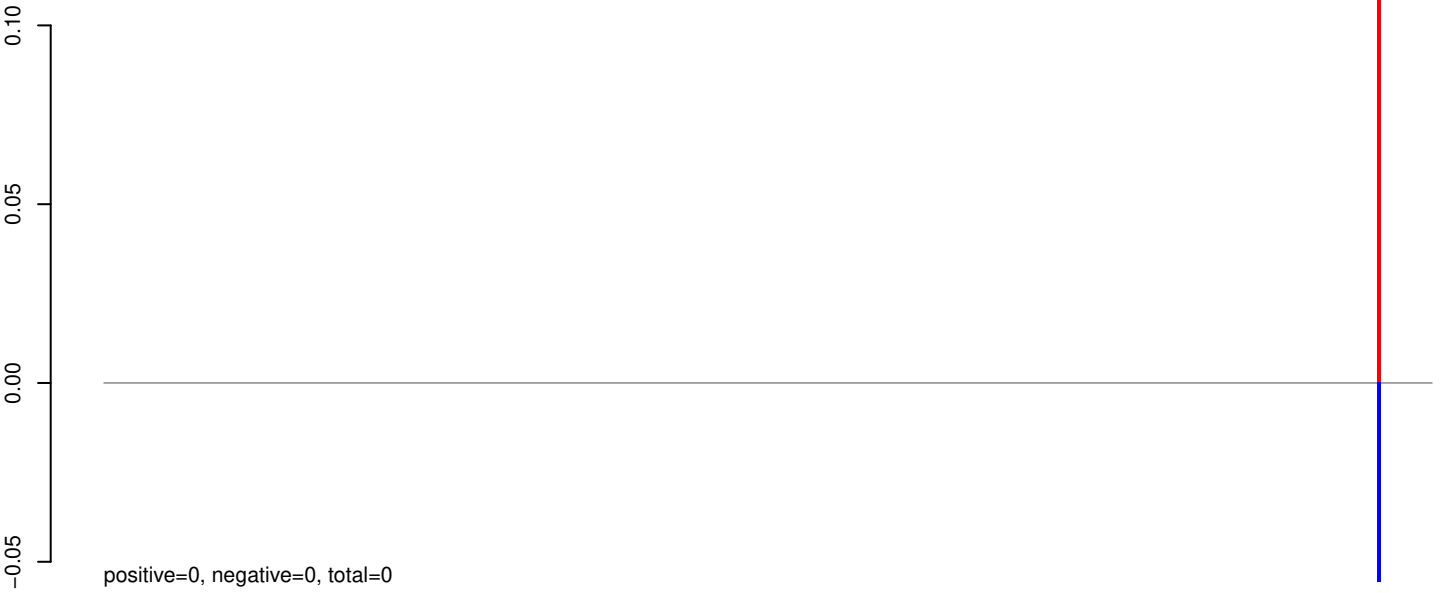
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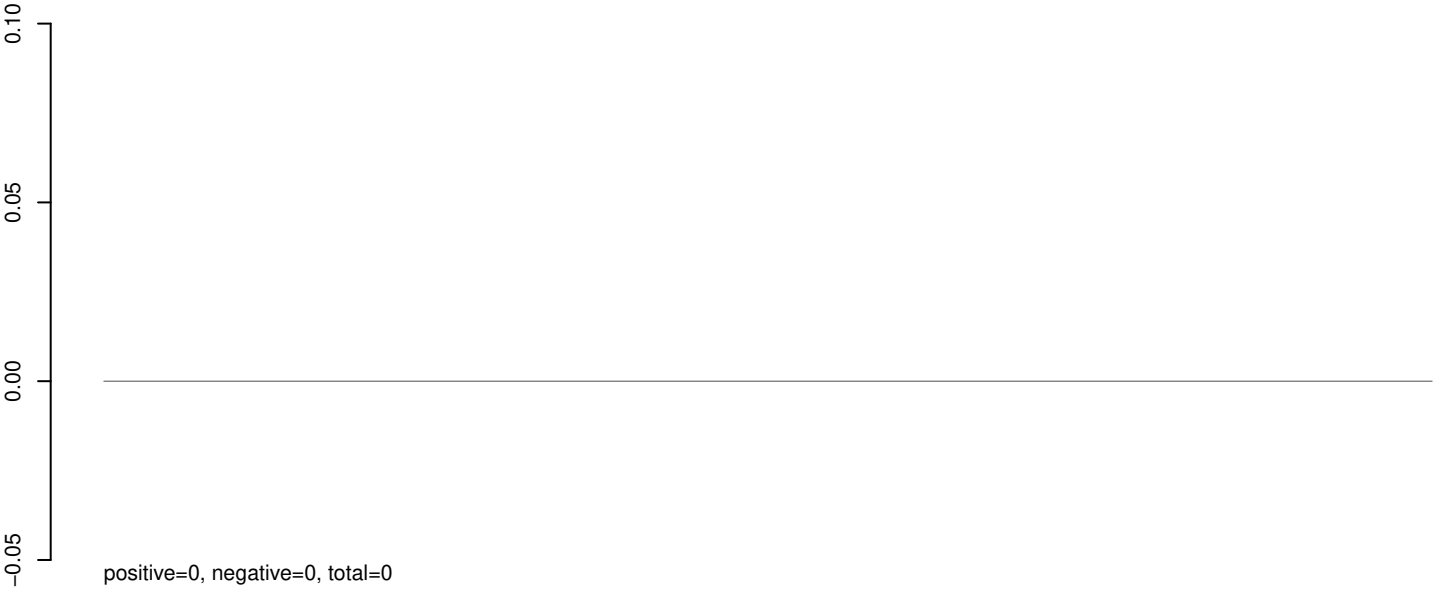
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0 2000 4000 6000 8000 10000

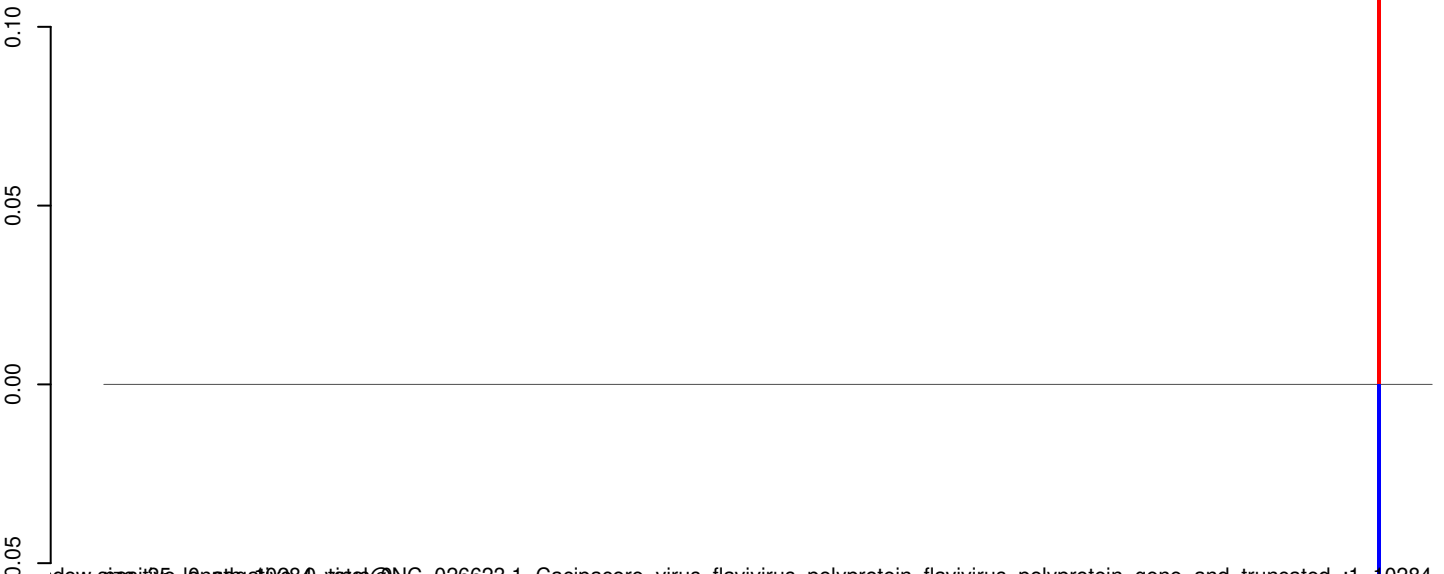
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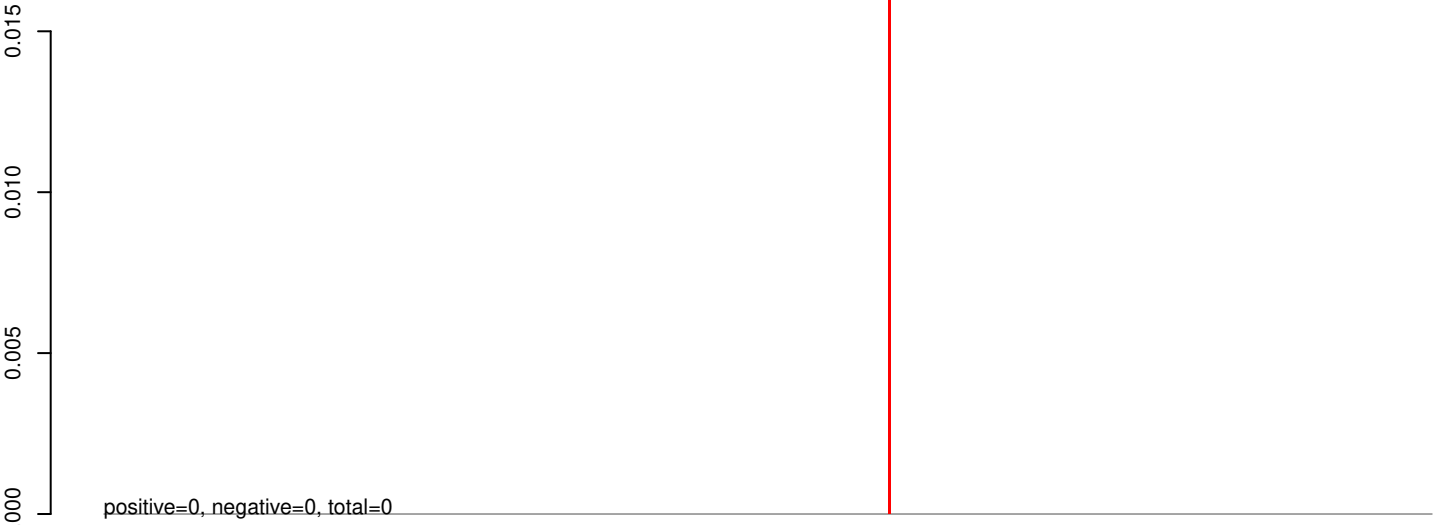
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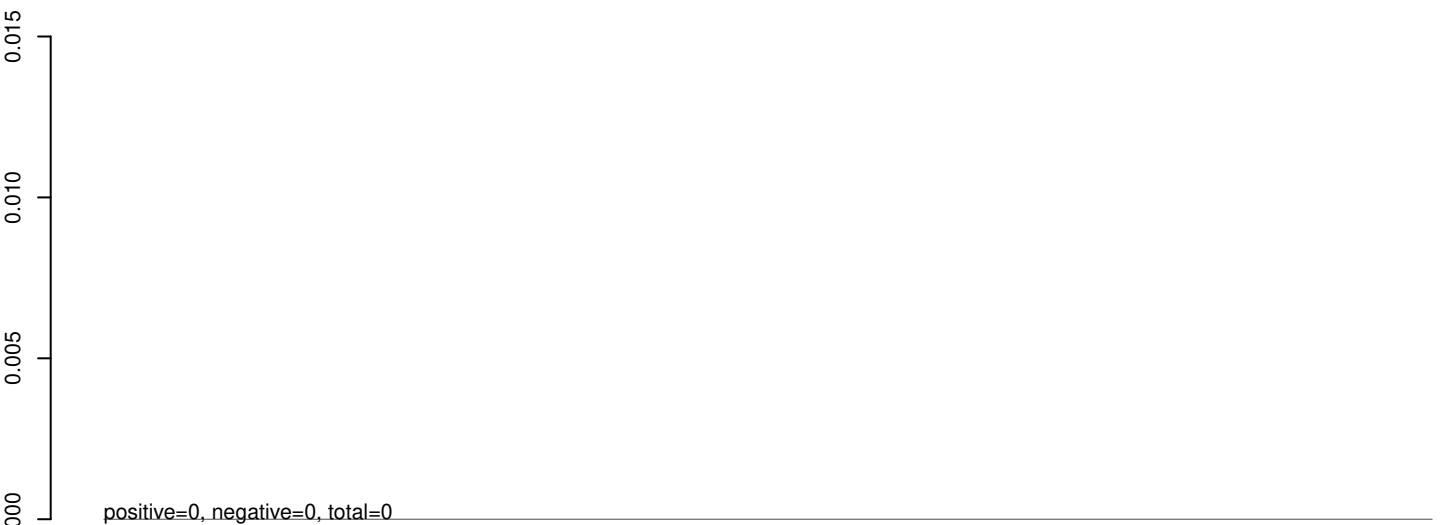
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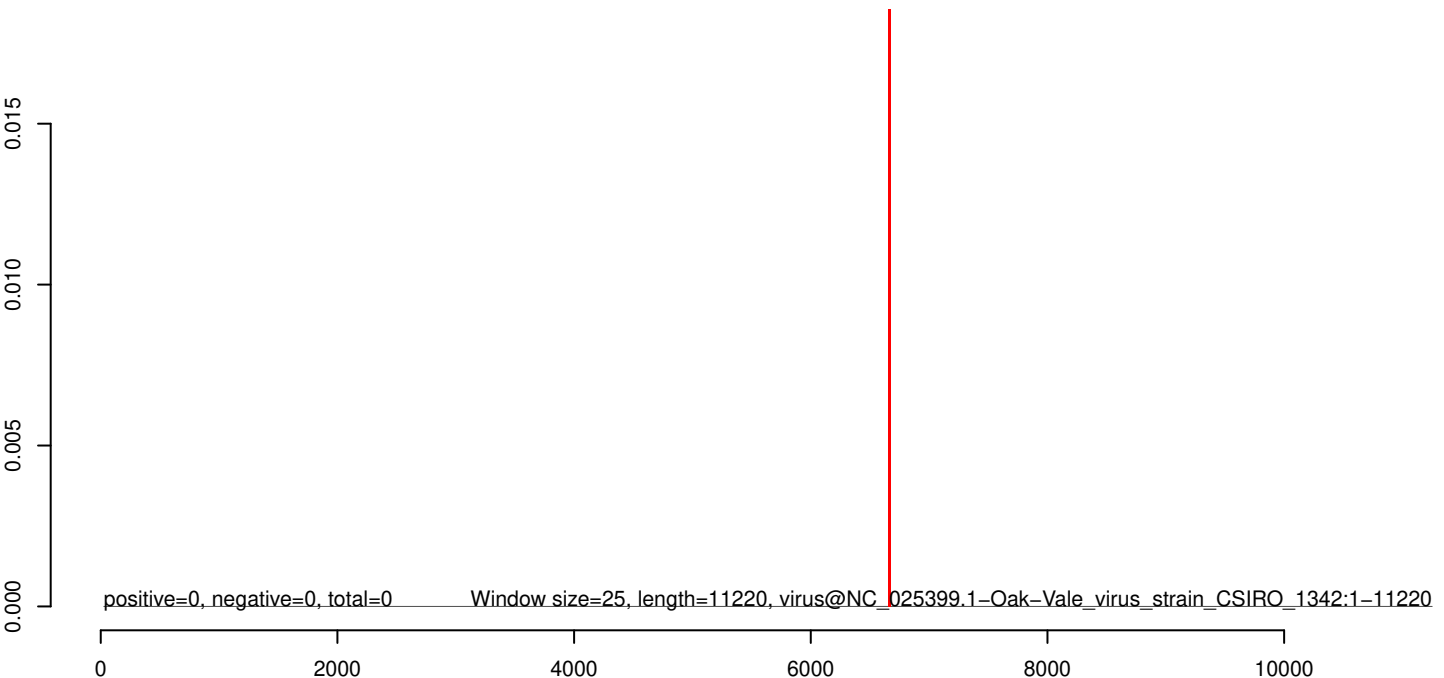
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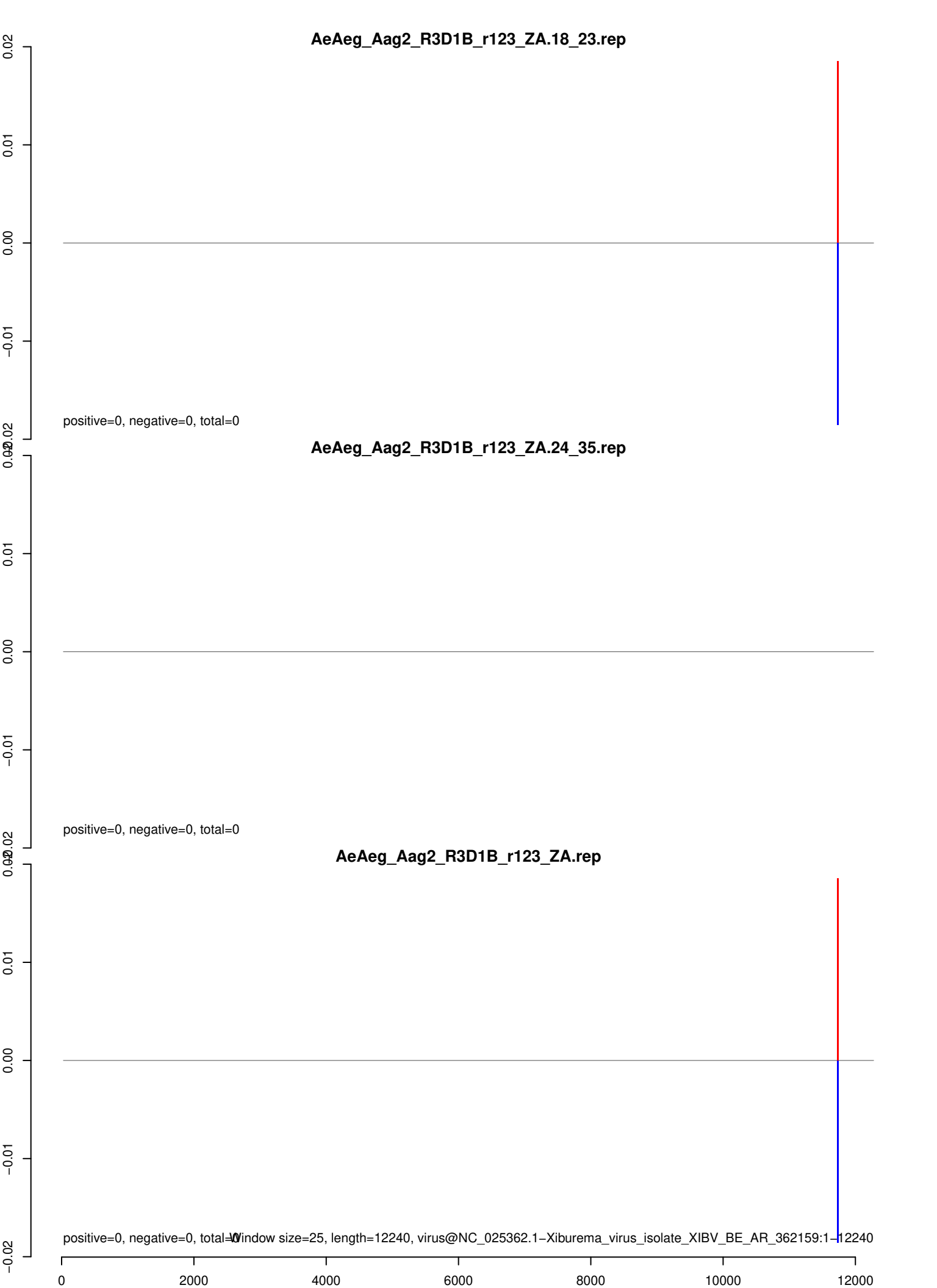


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AeAeg_Aag2_R3D1B_r123_ZA.rep

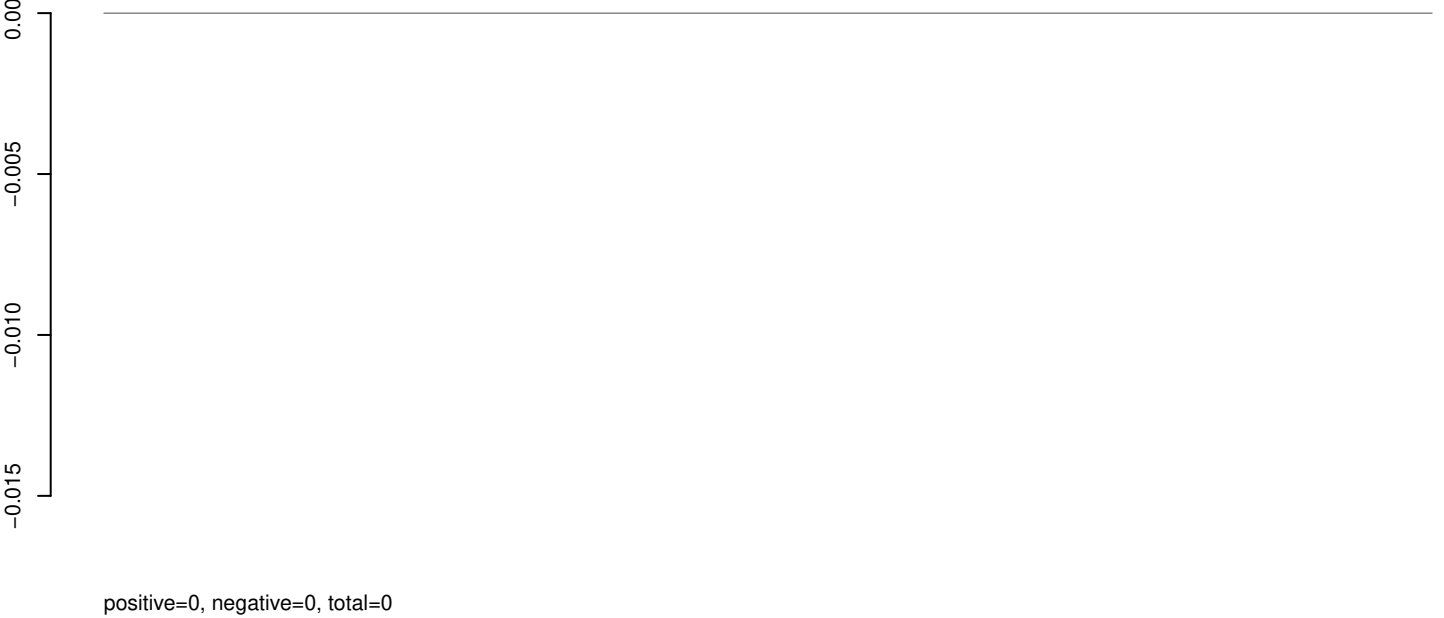




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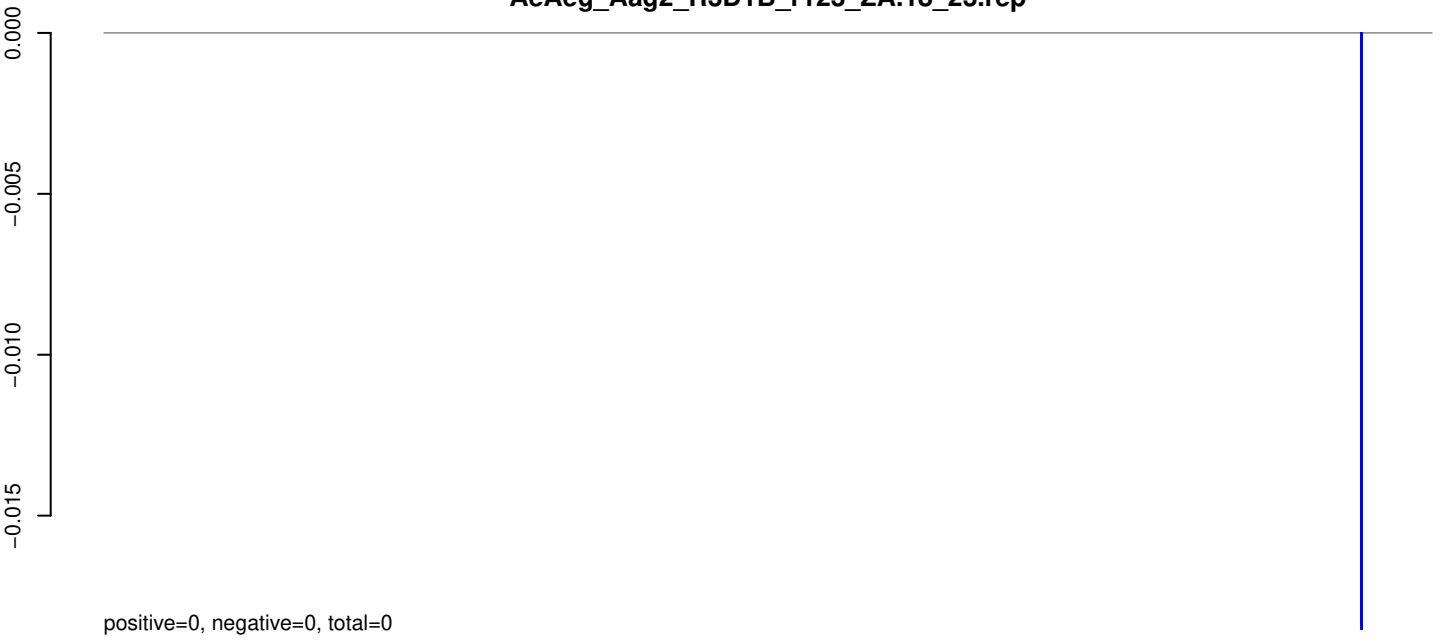
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W400 low size=25, length=10365, virus@NC_024805.1-Ilomantsi_virus_isolate_M0724_polyprotein_gene:1-10365

0 2000 4000 6000 8000 10000

AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



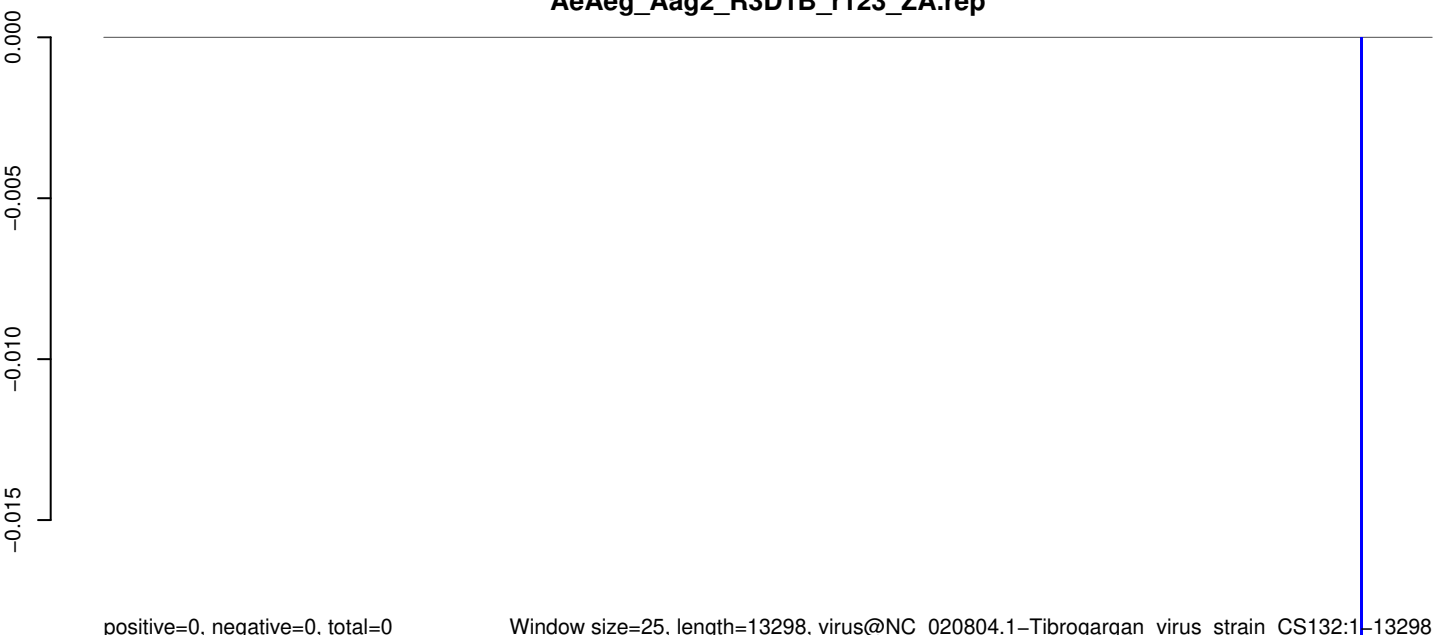
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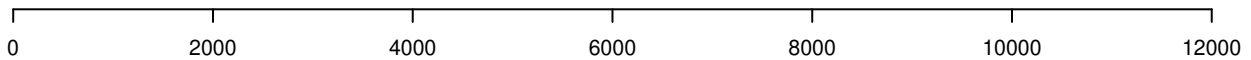
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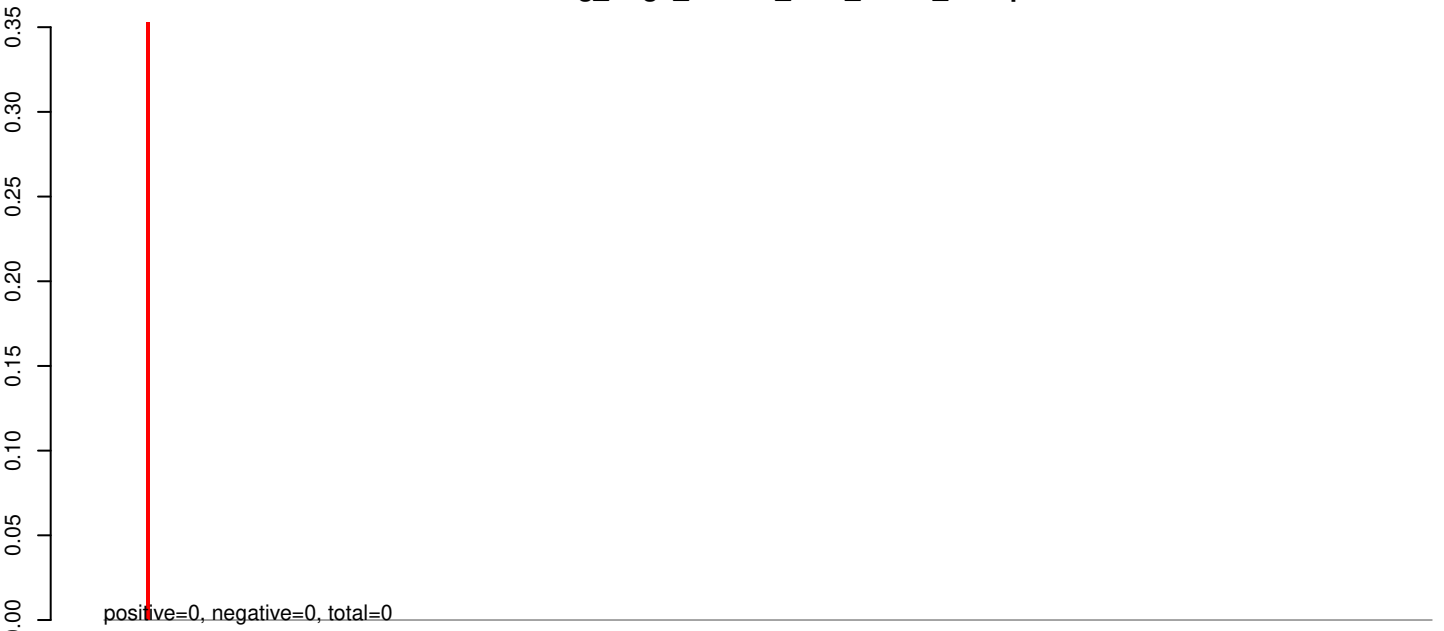


positive=0, negative=0, total=0

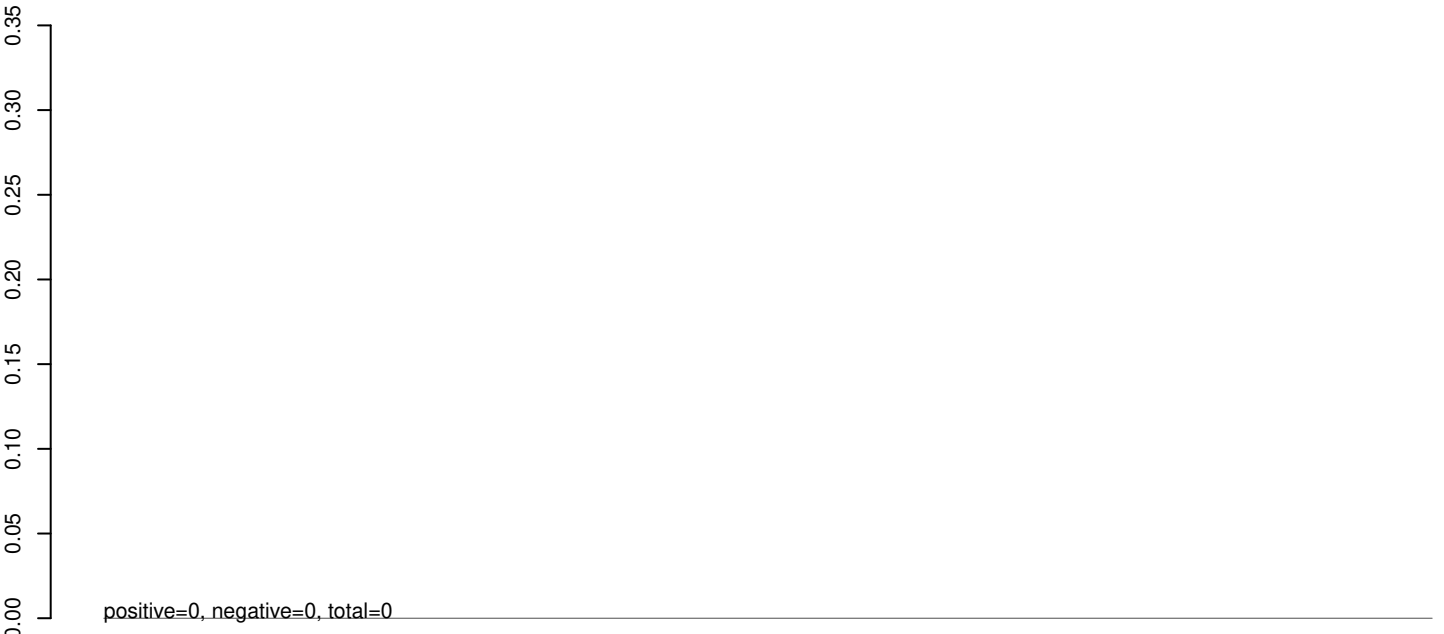
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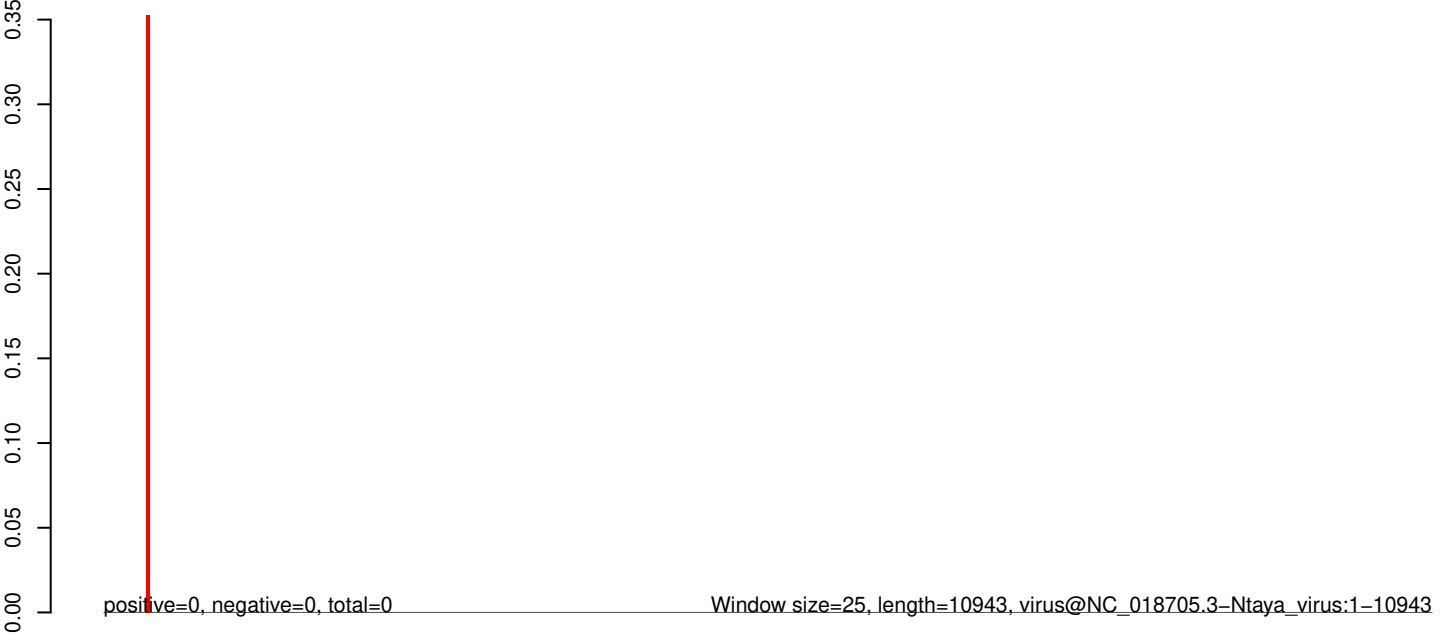
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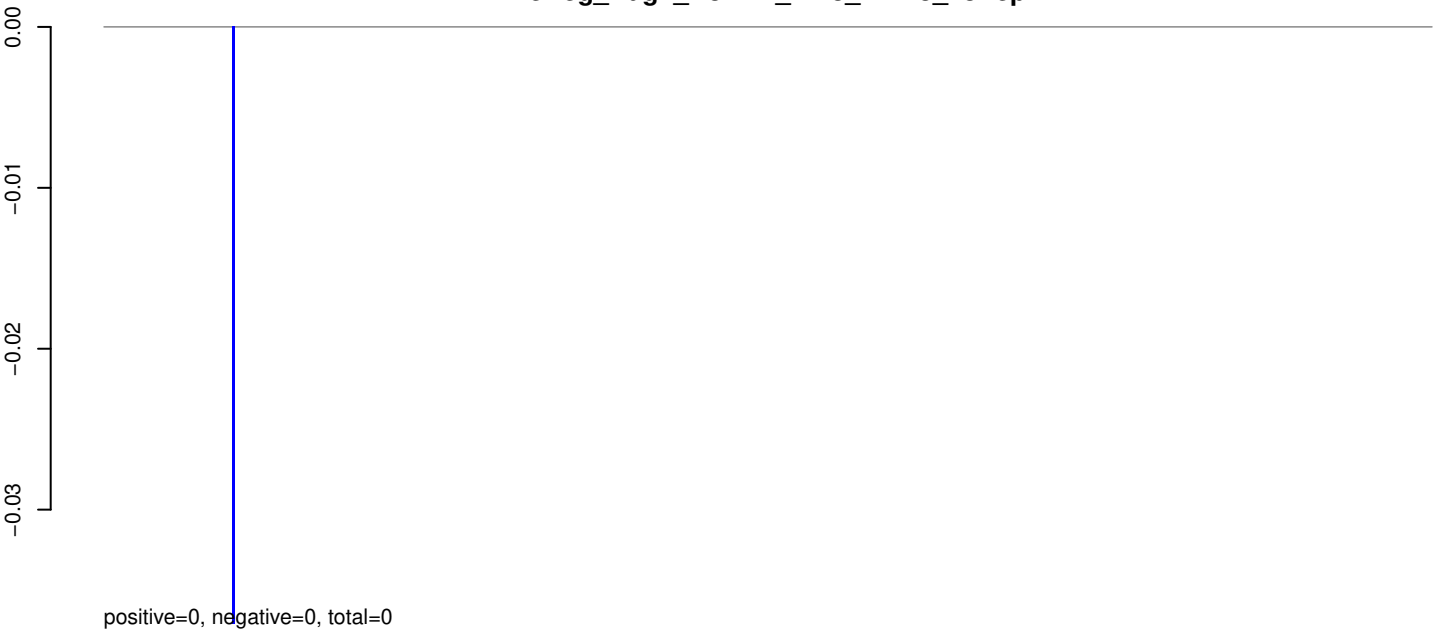


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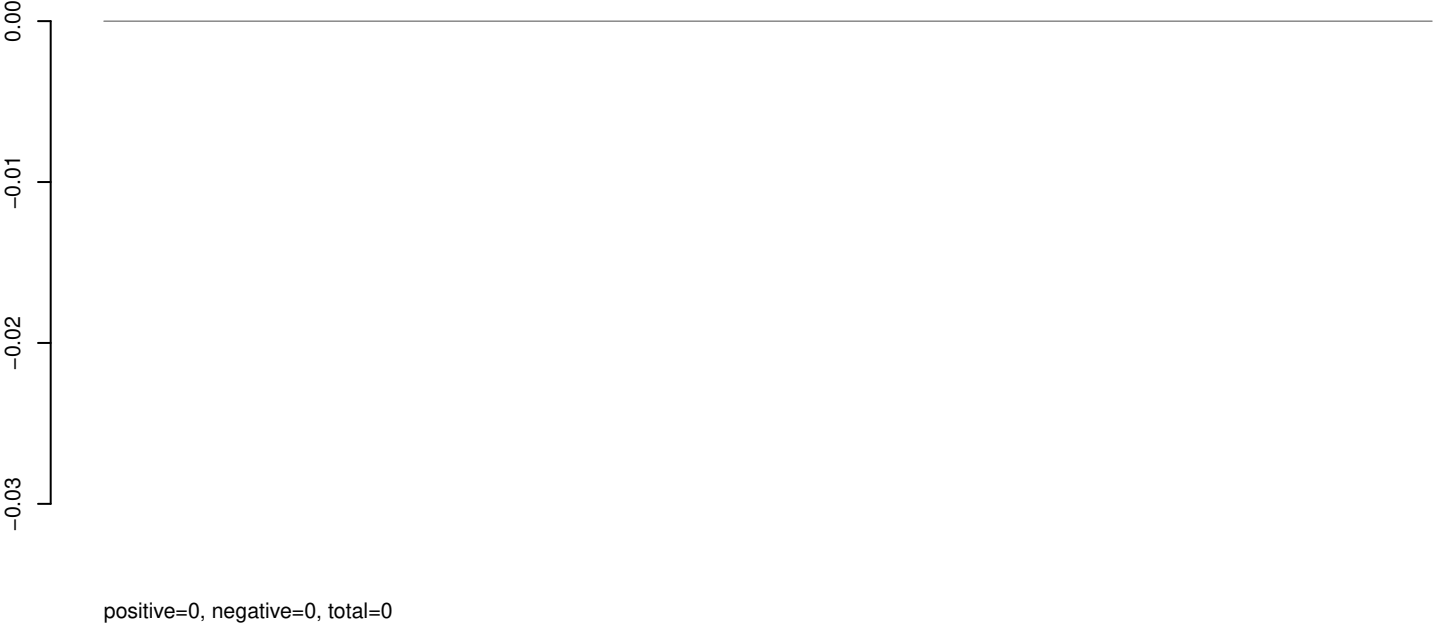


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AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep



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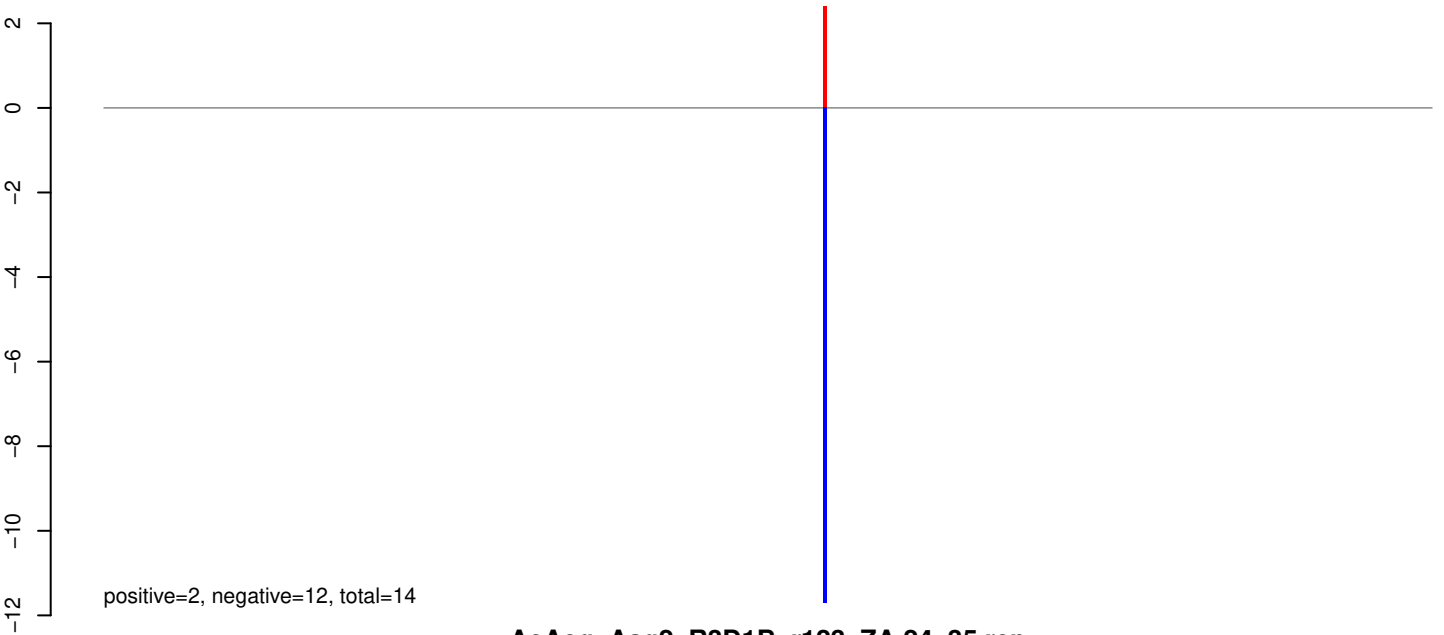
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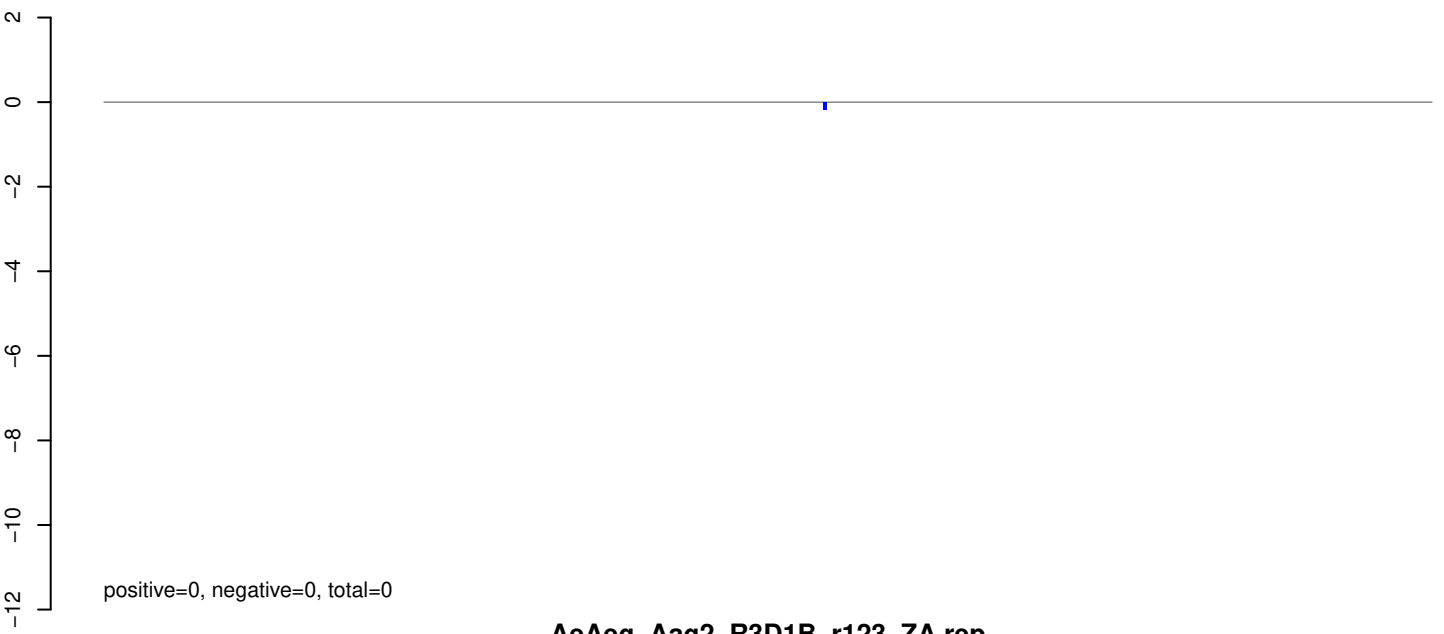
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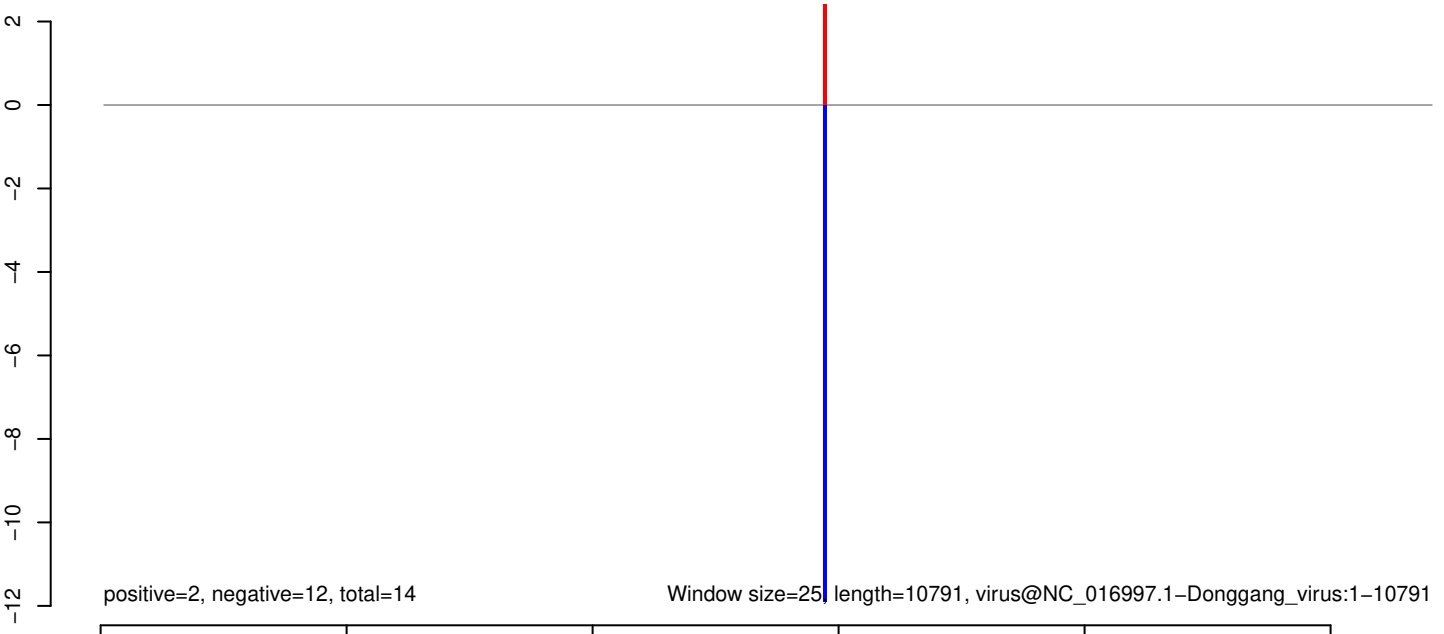
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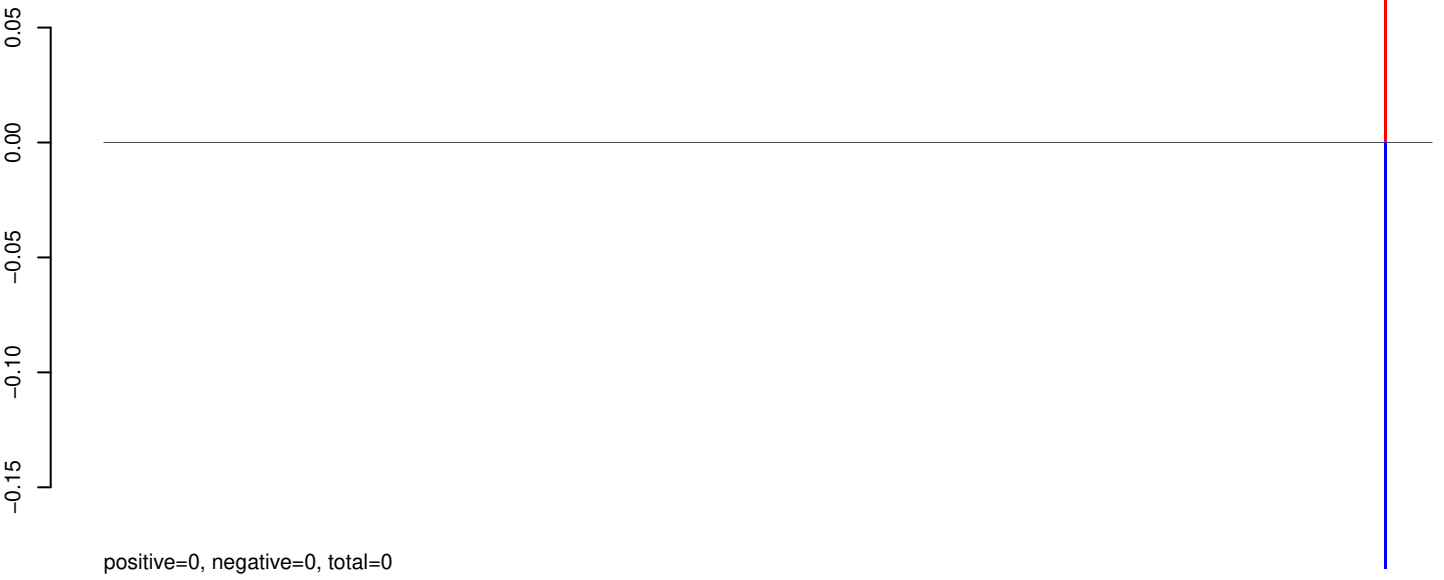


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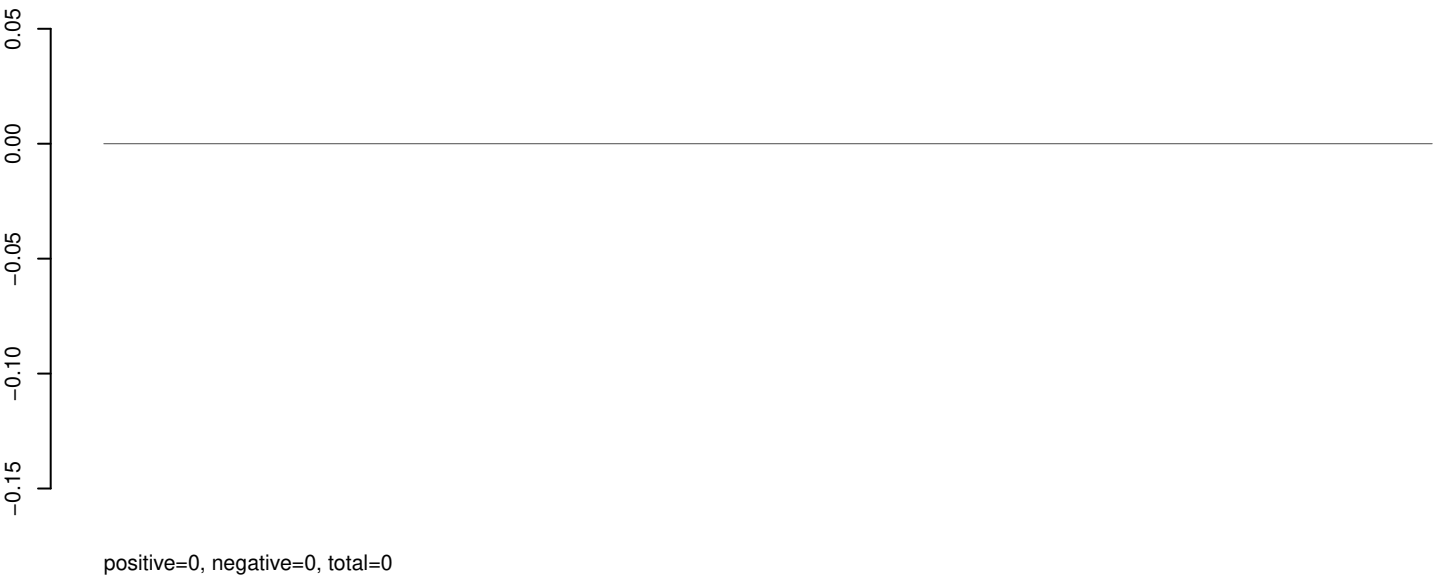
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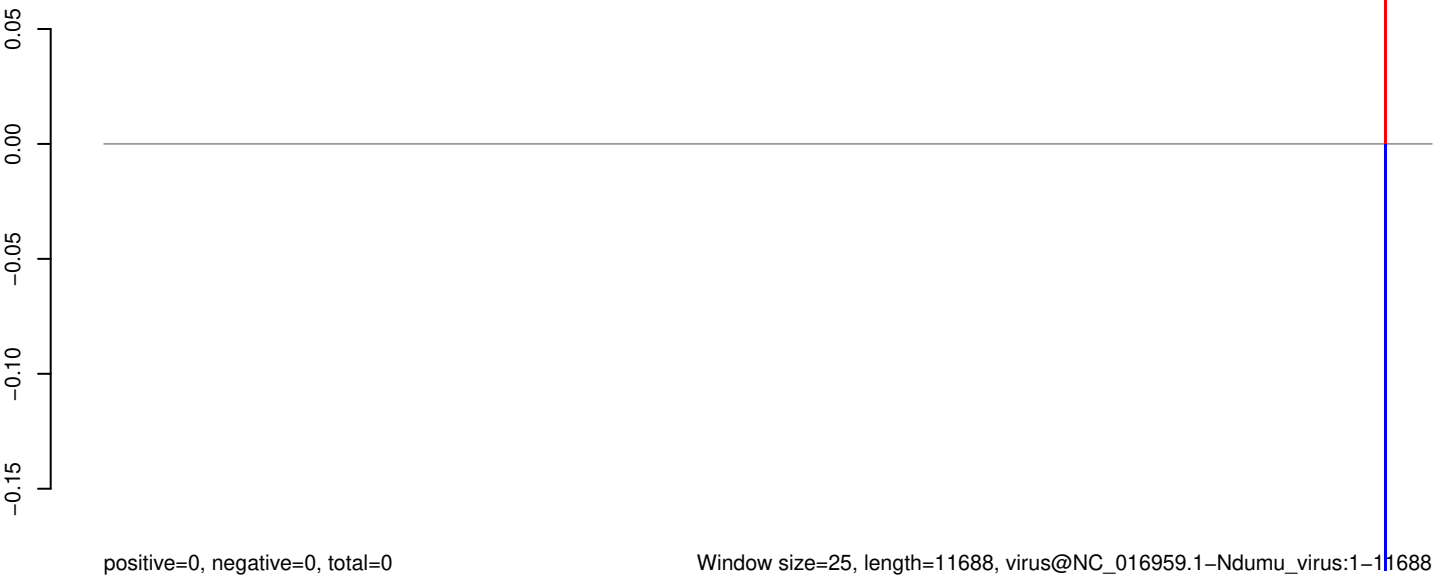
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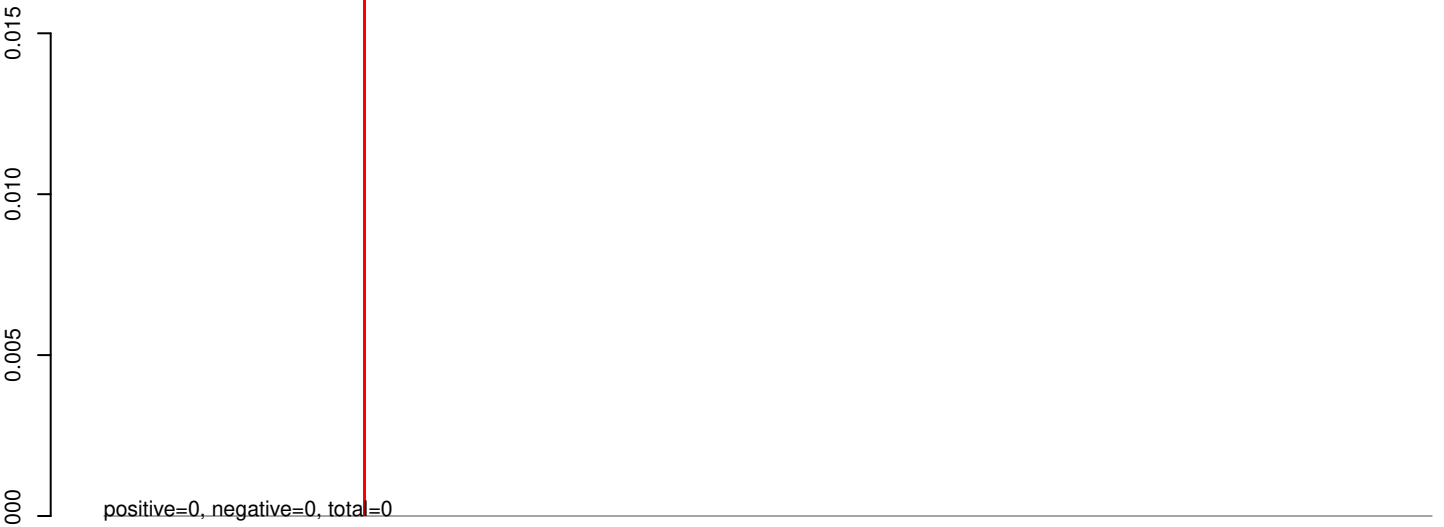
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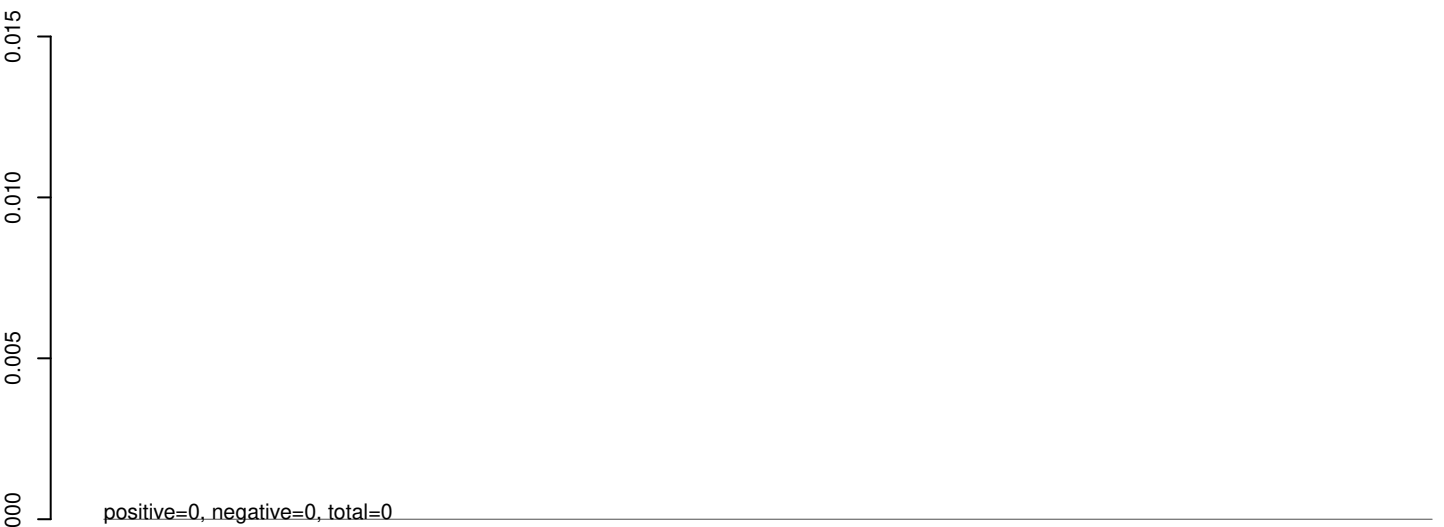
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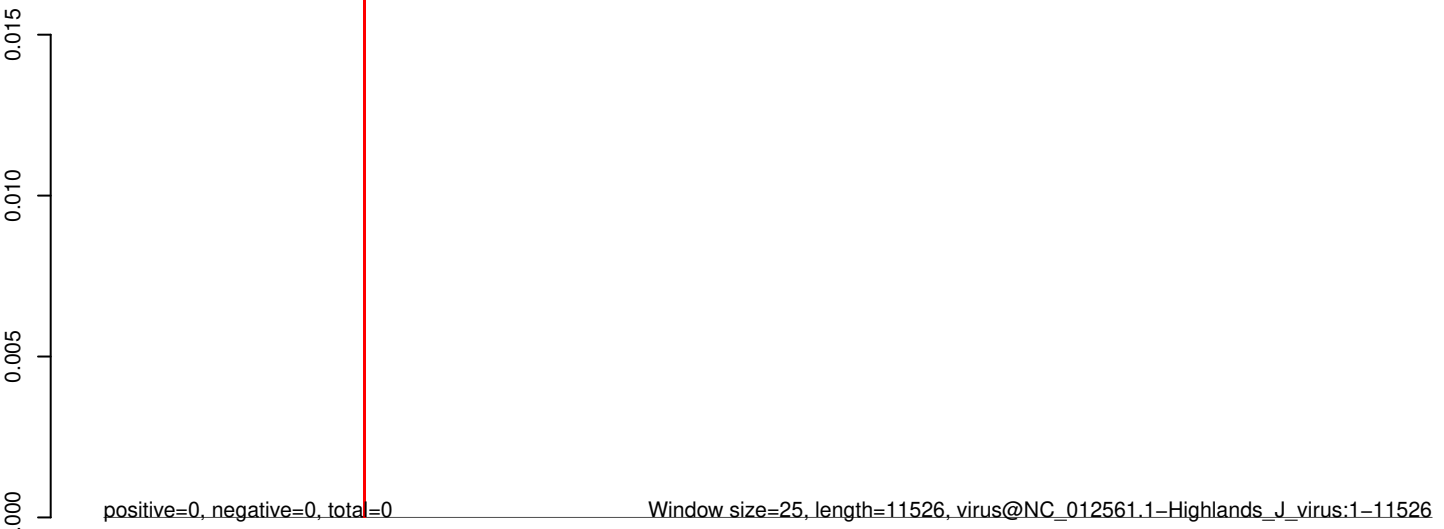
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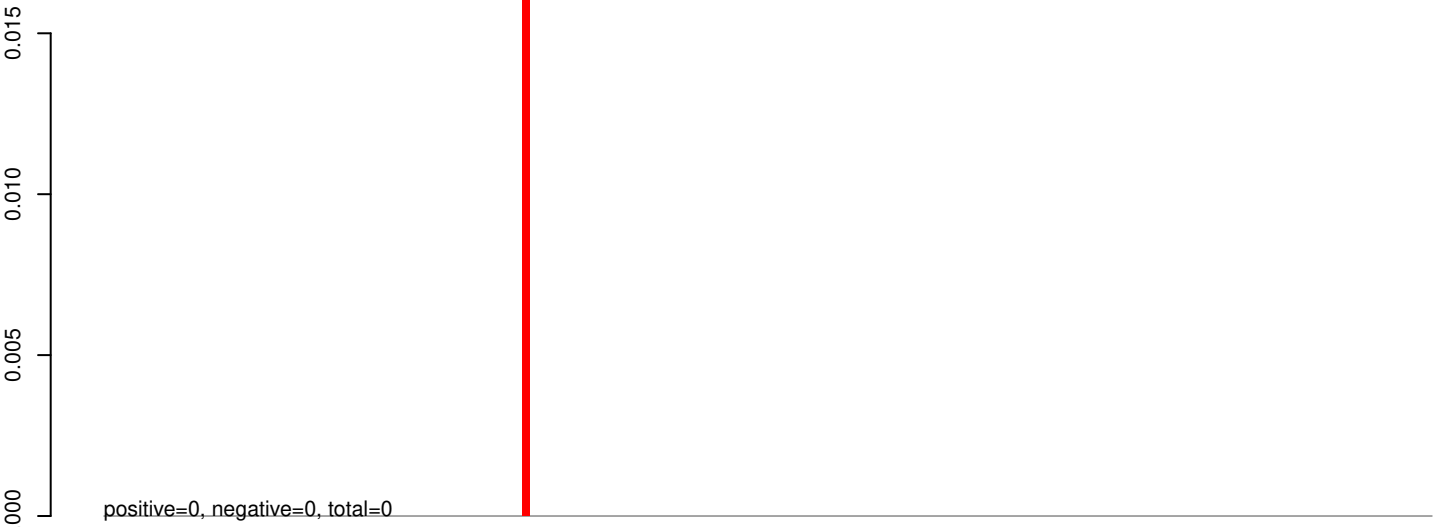
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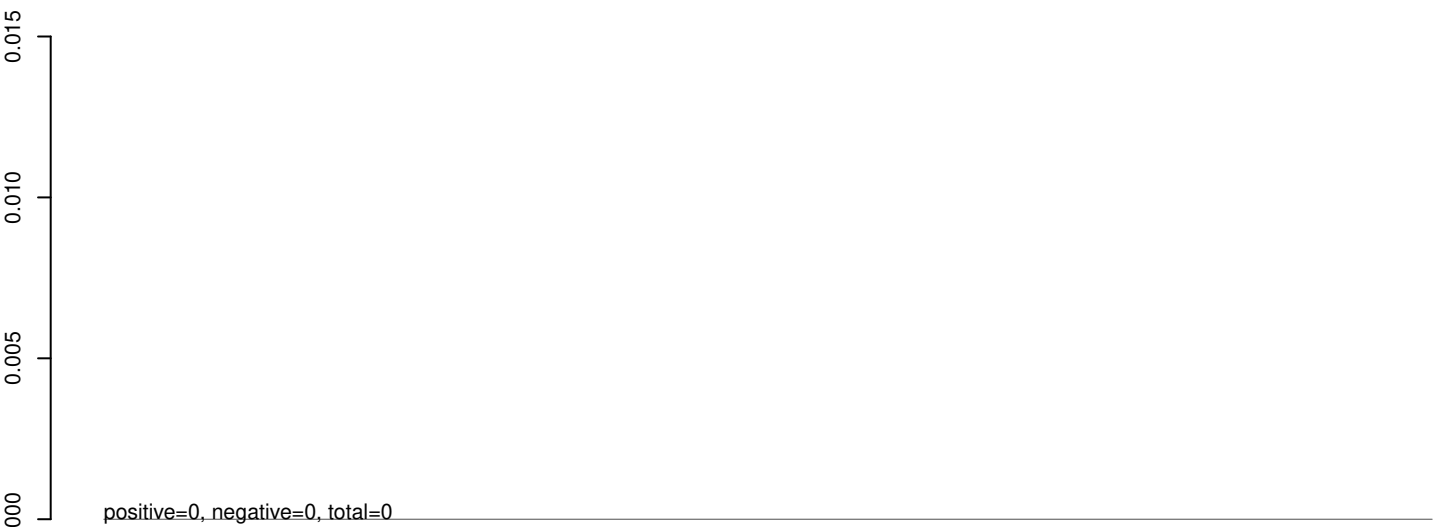
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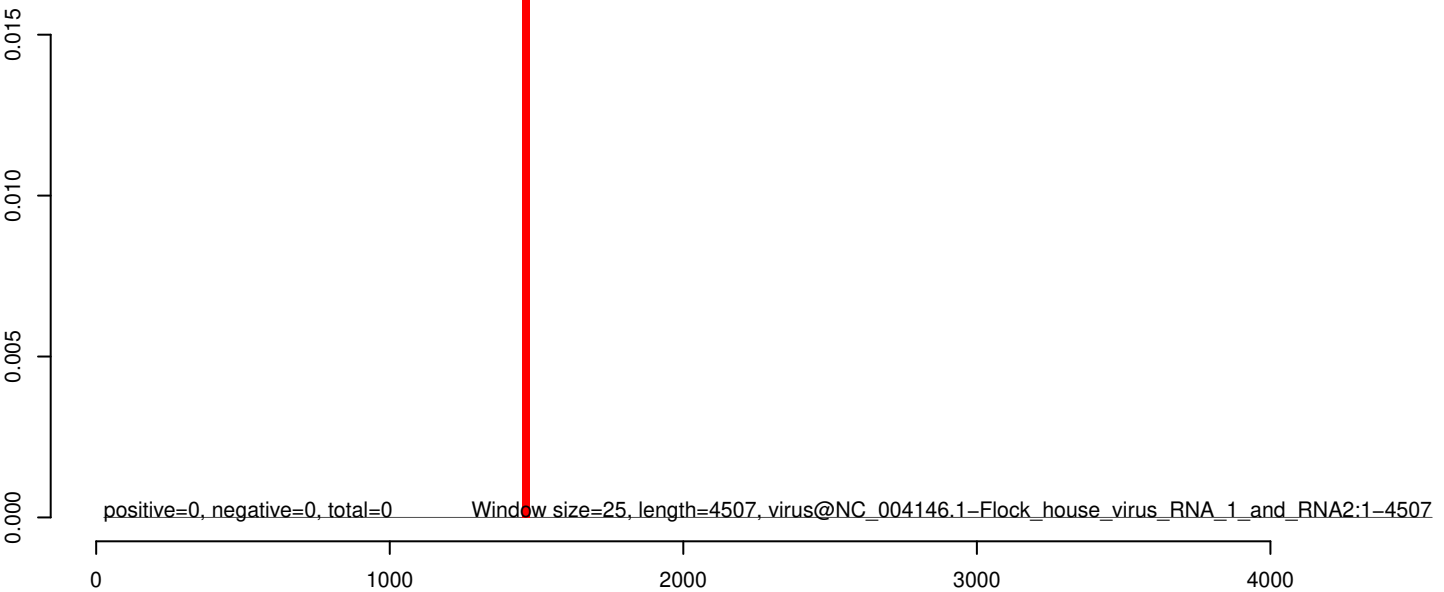
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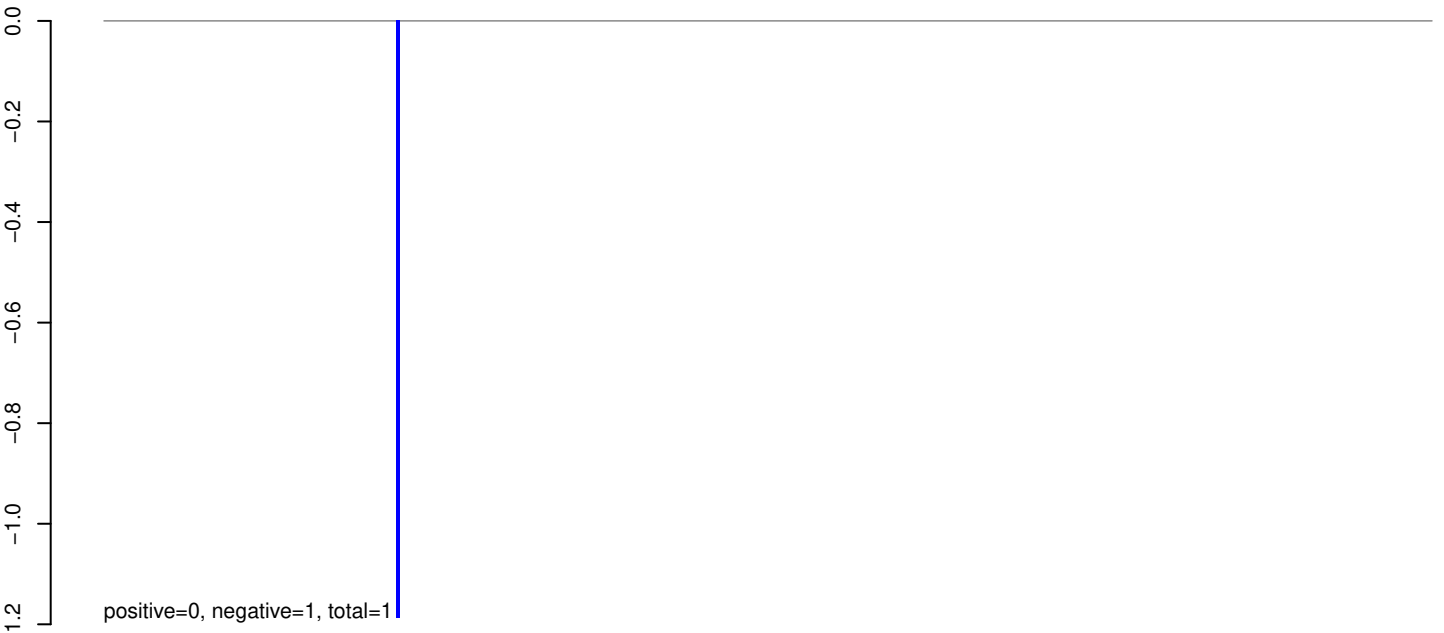
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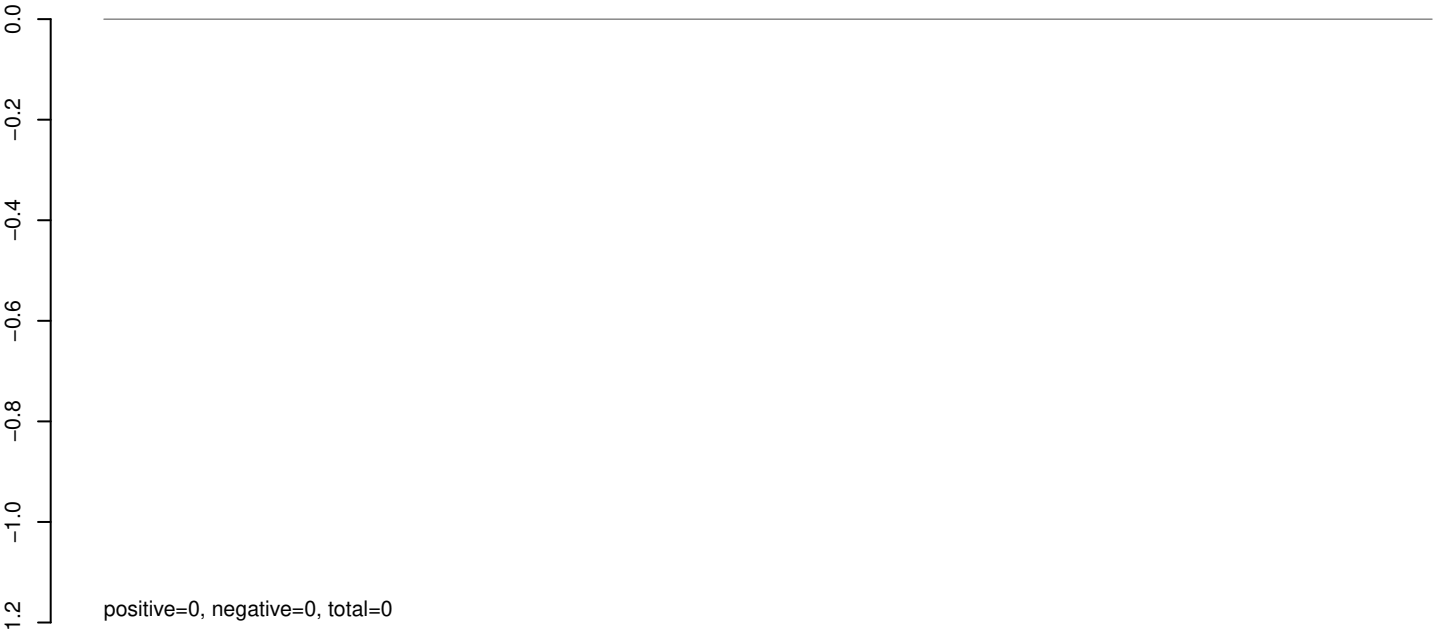
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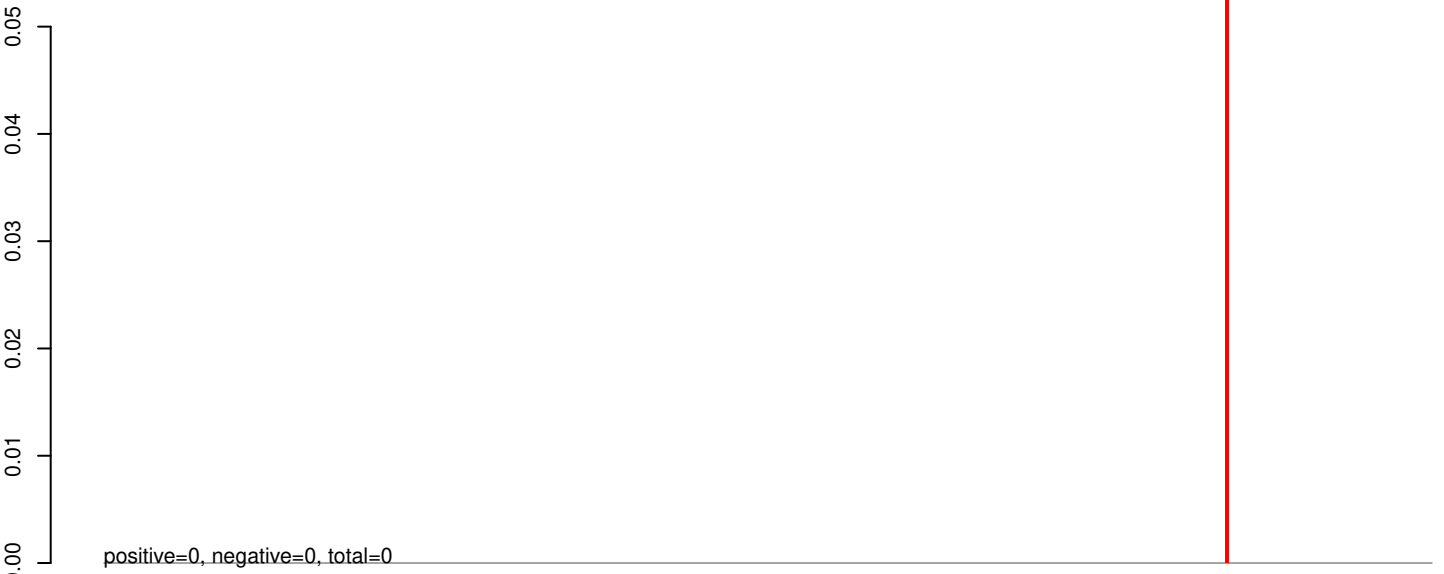


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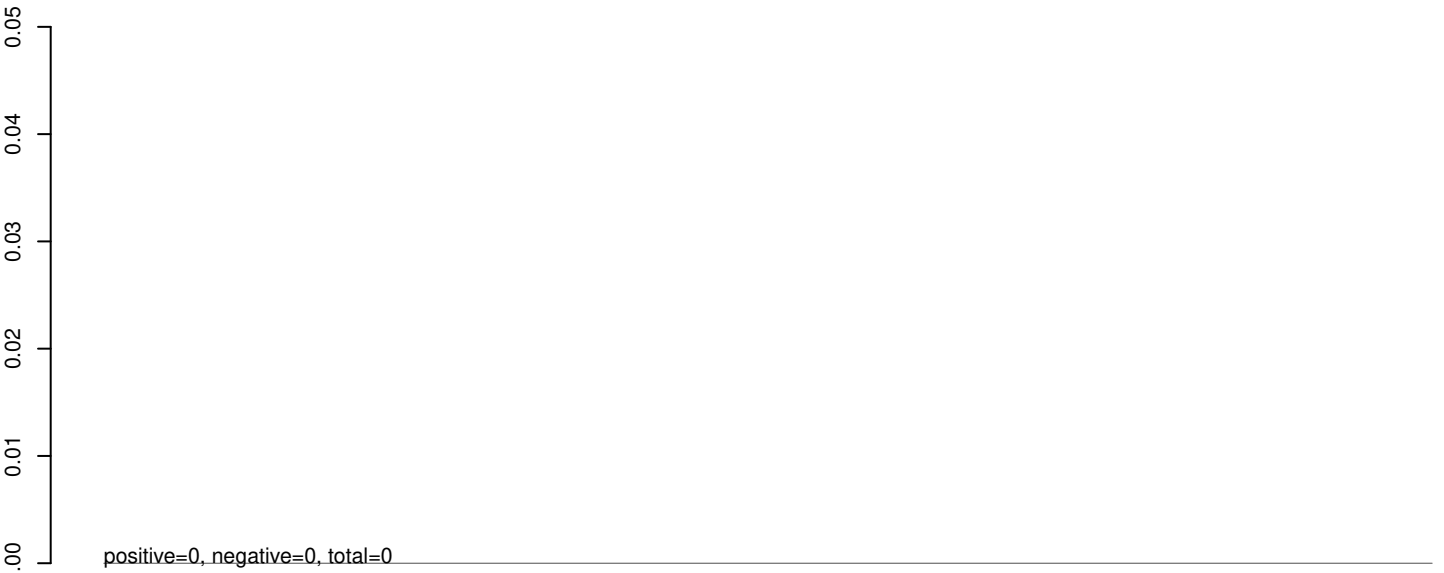




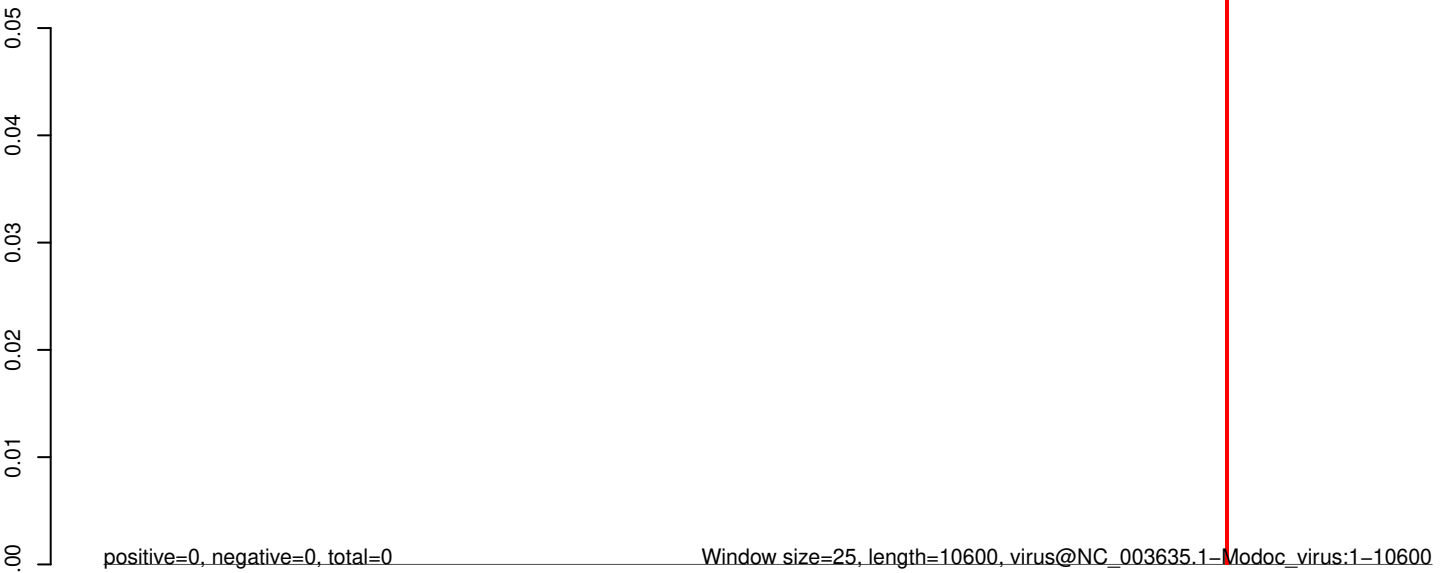
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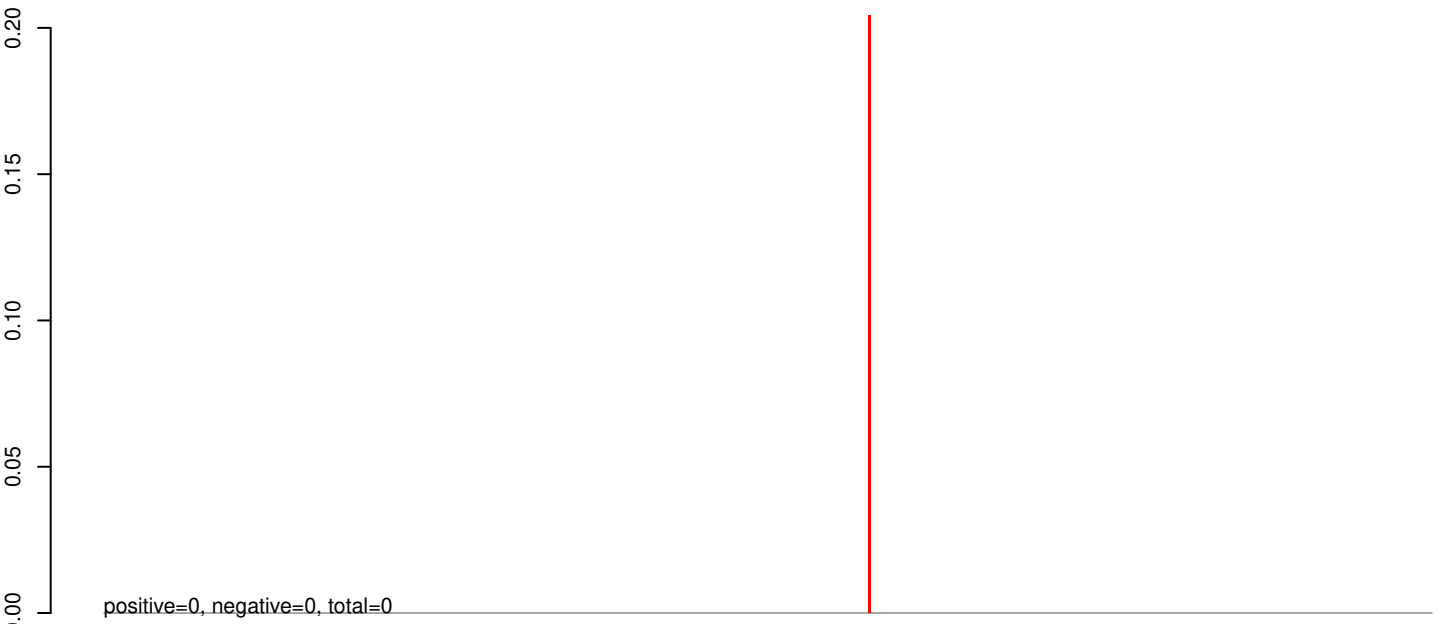
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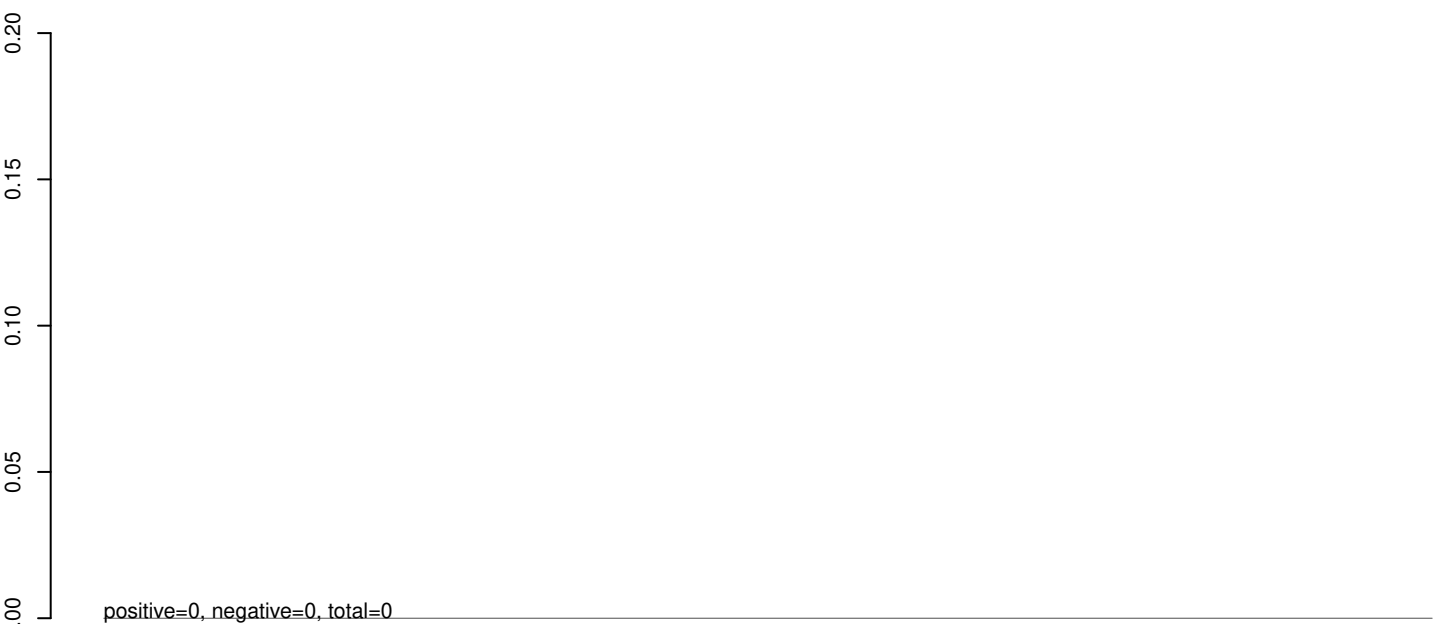
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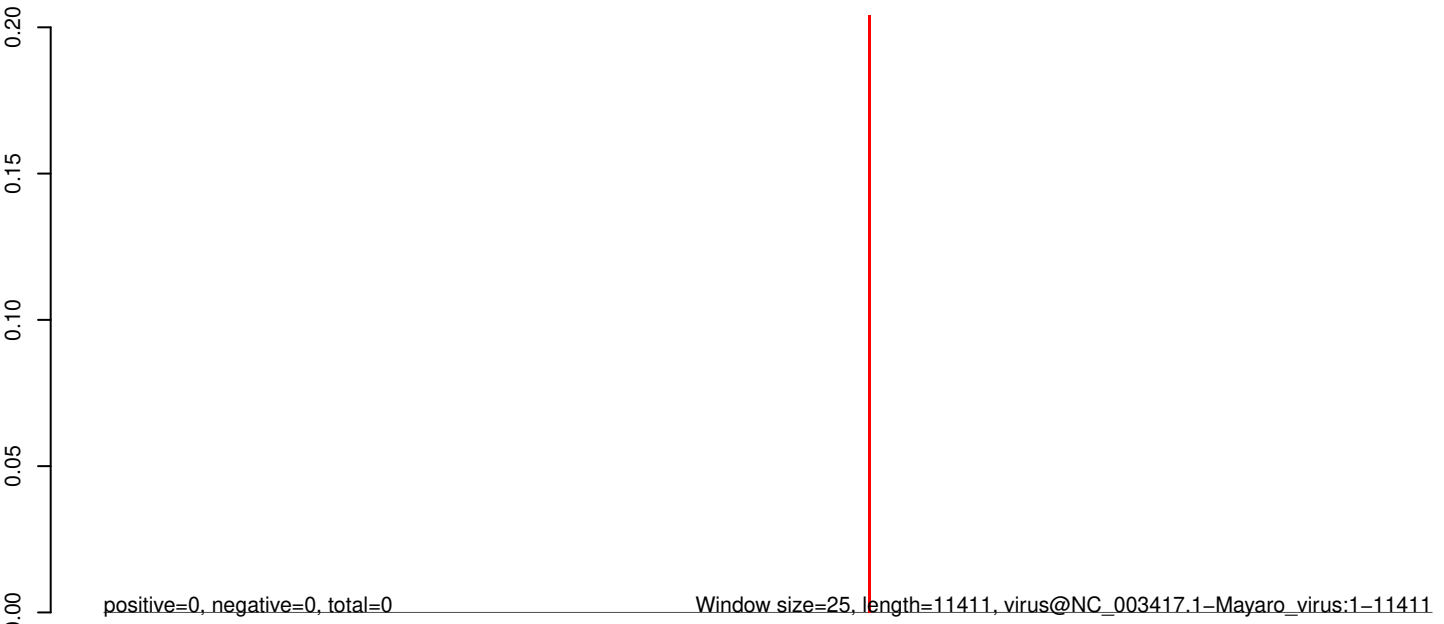
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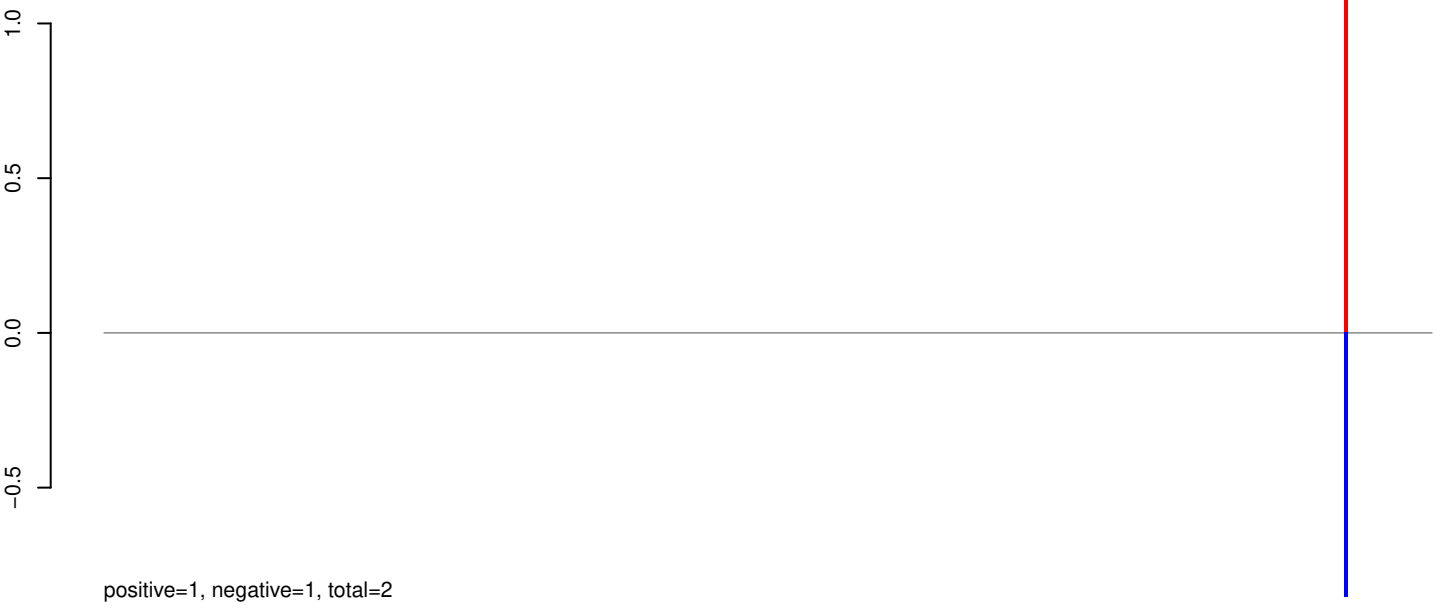


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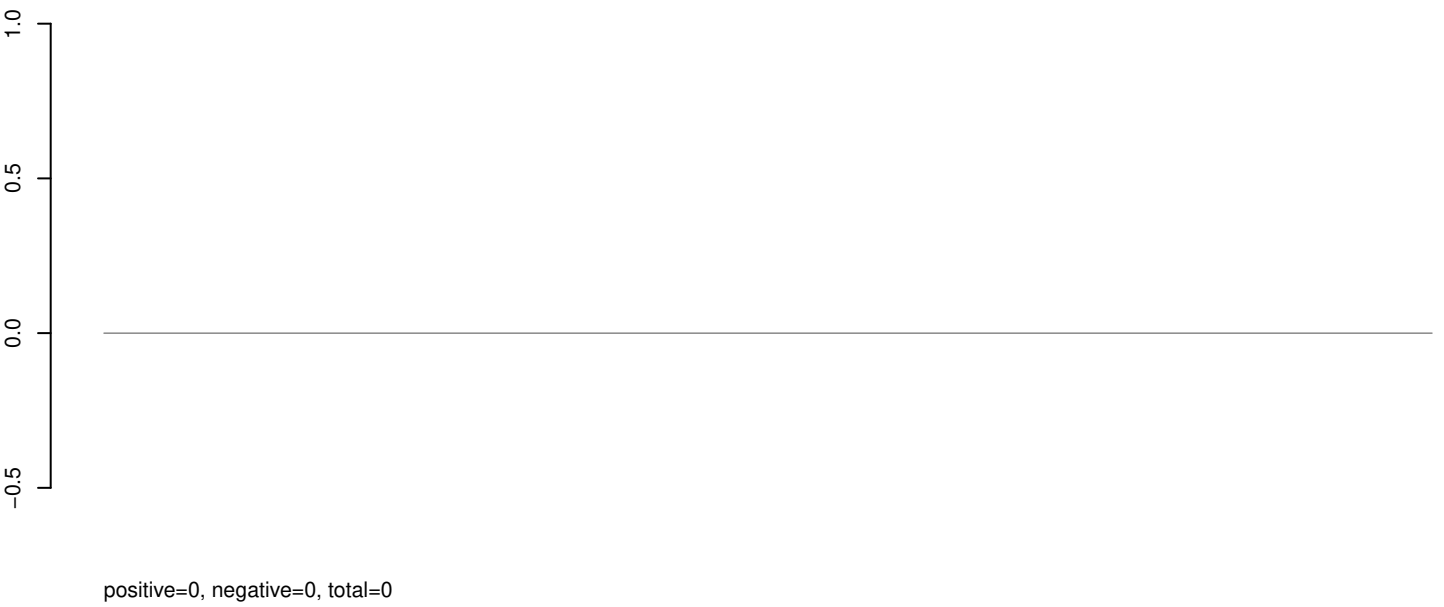


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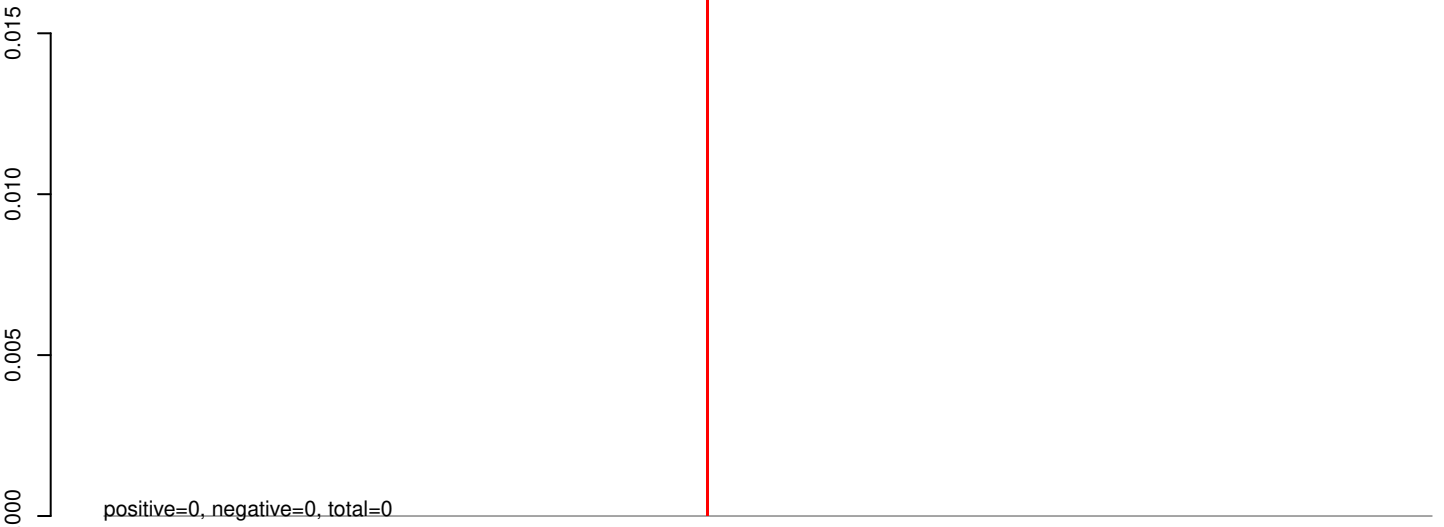
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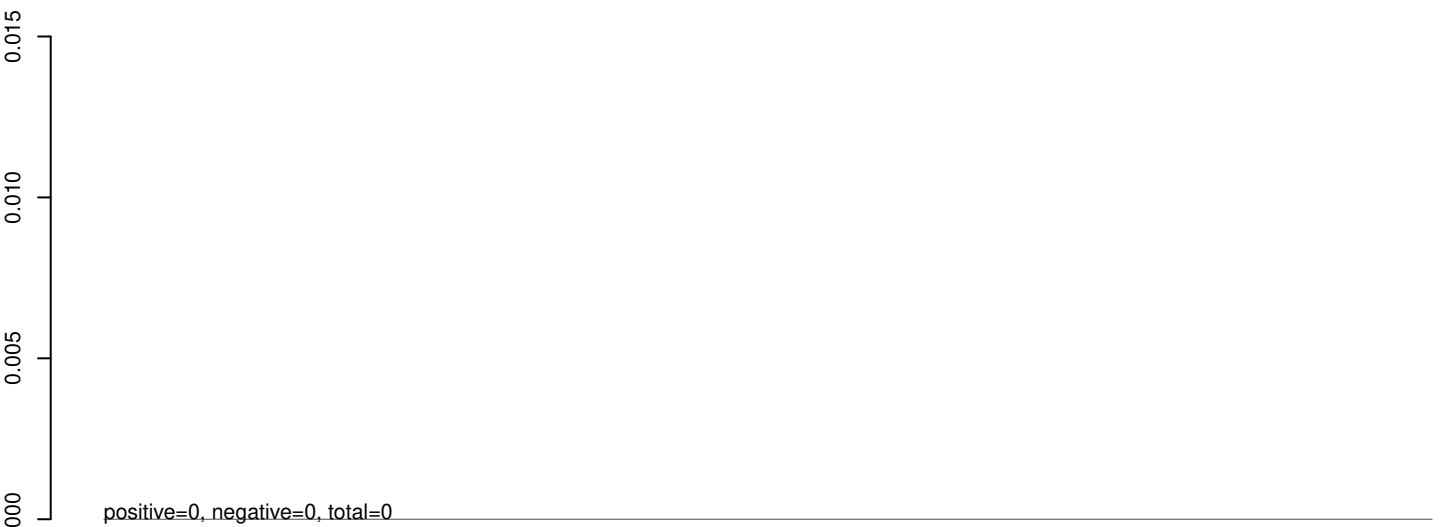
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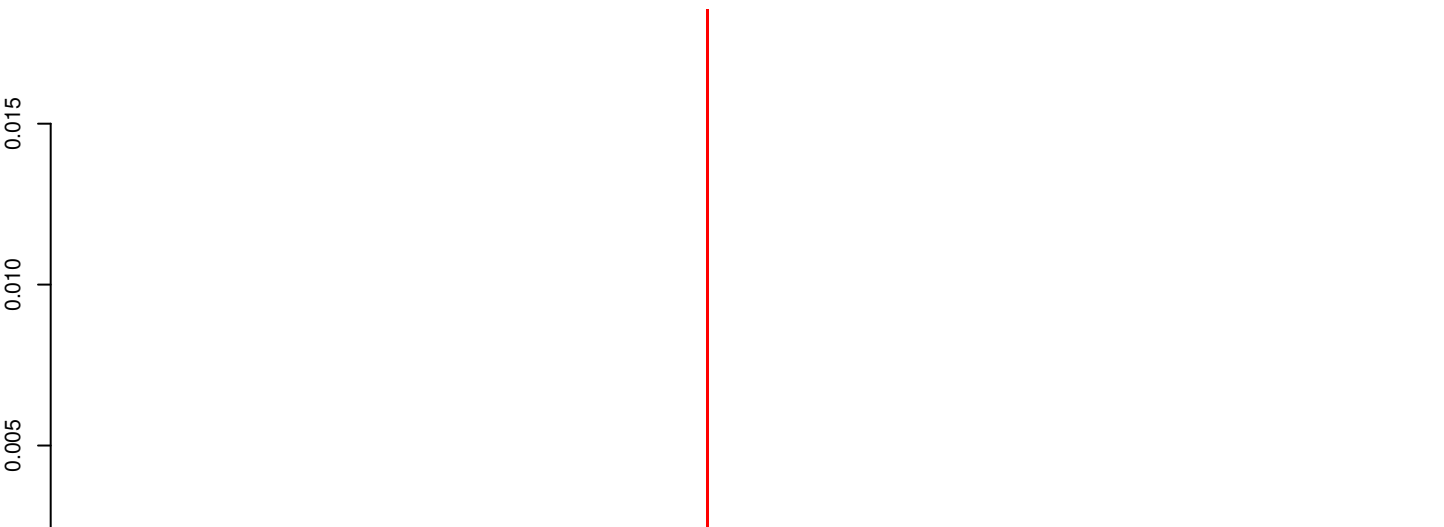
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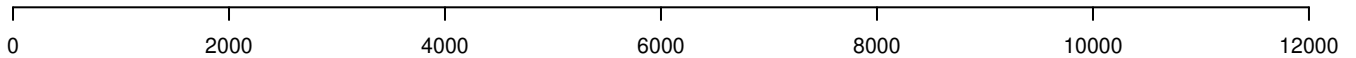
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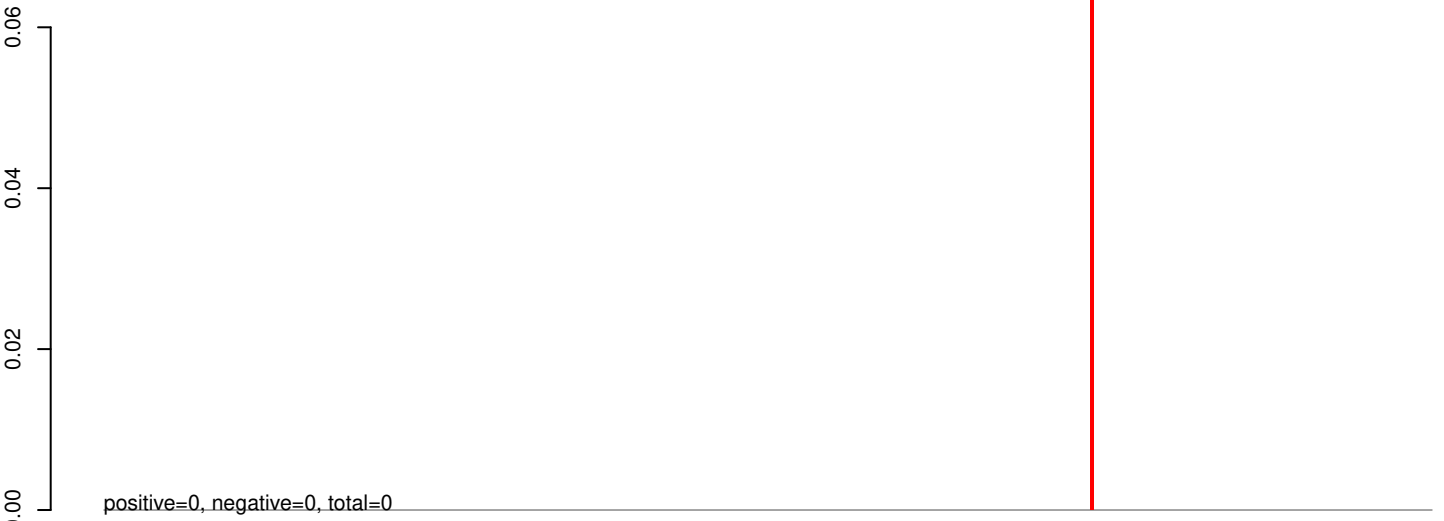
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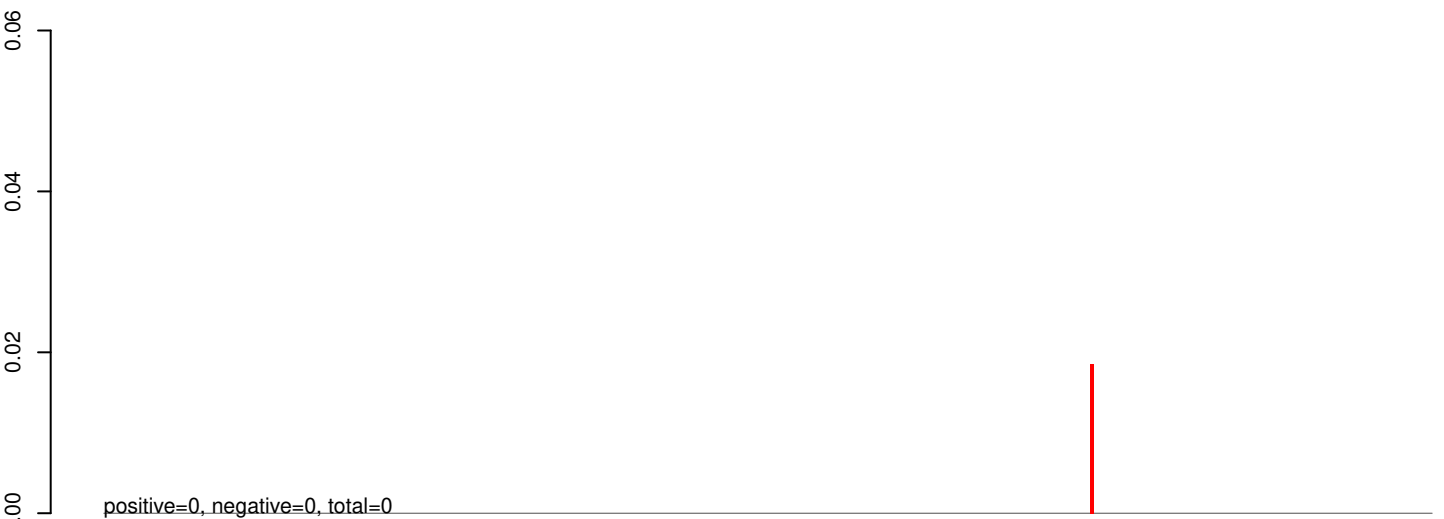
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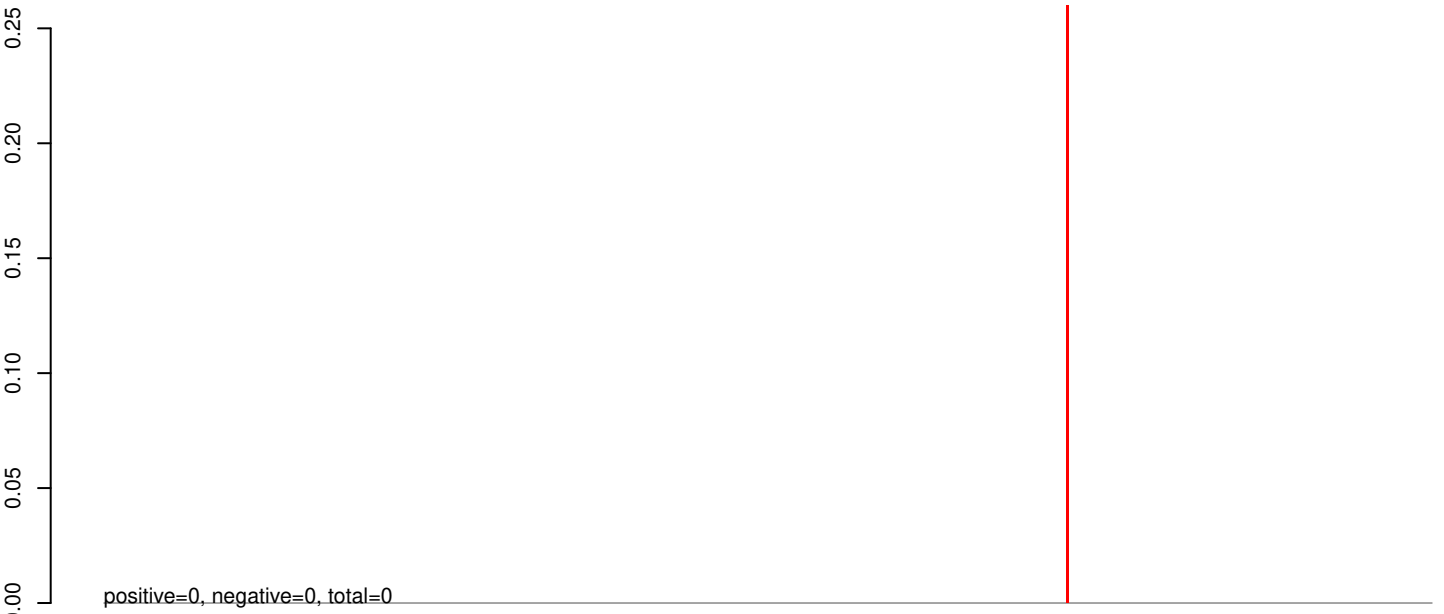


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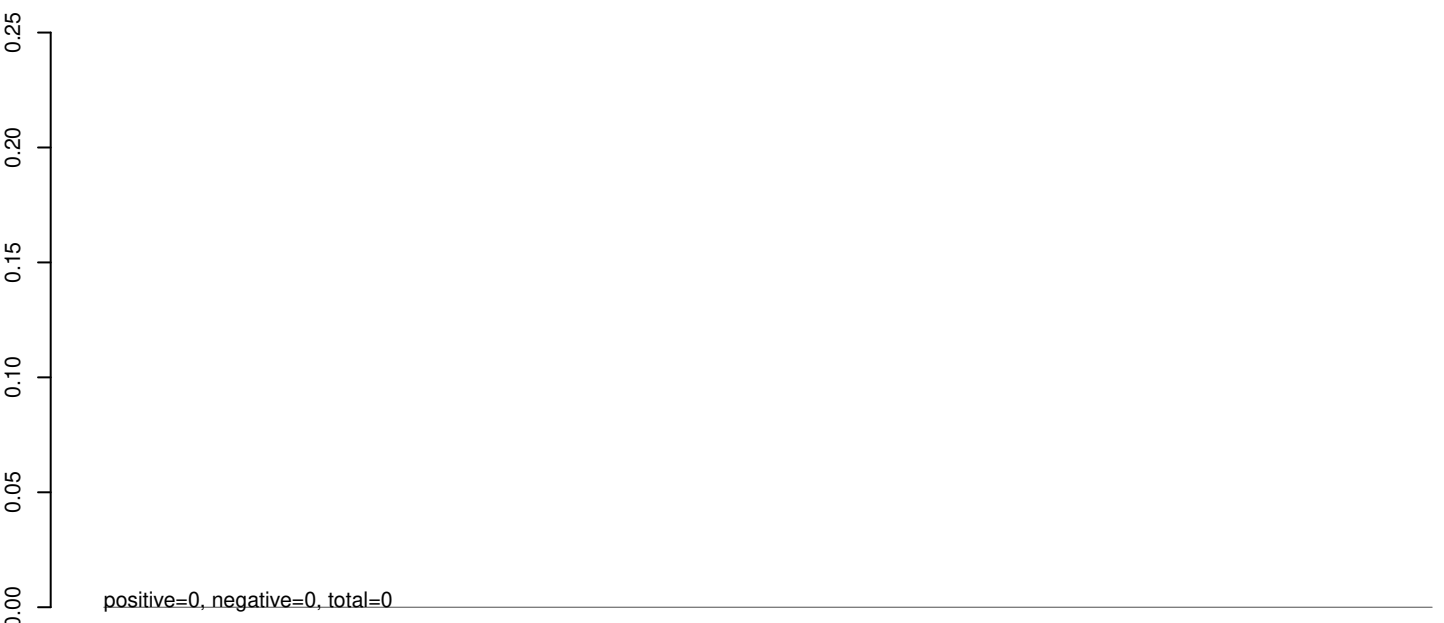


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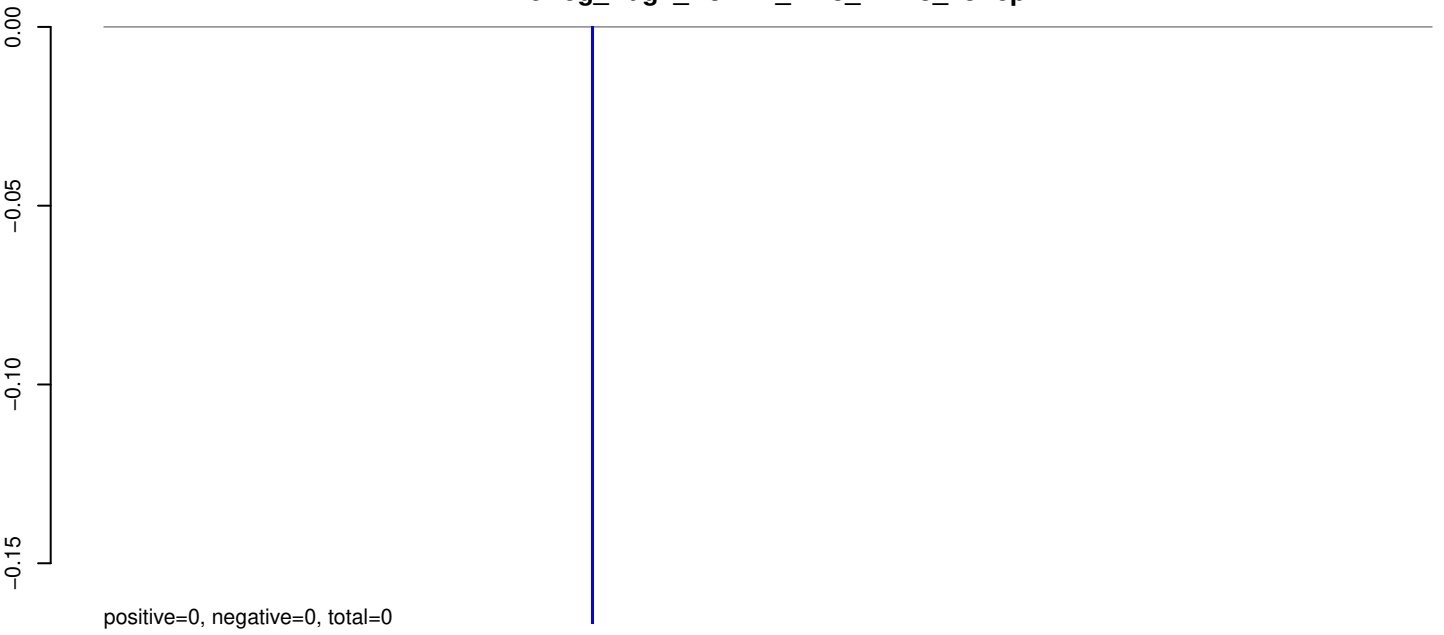
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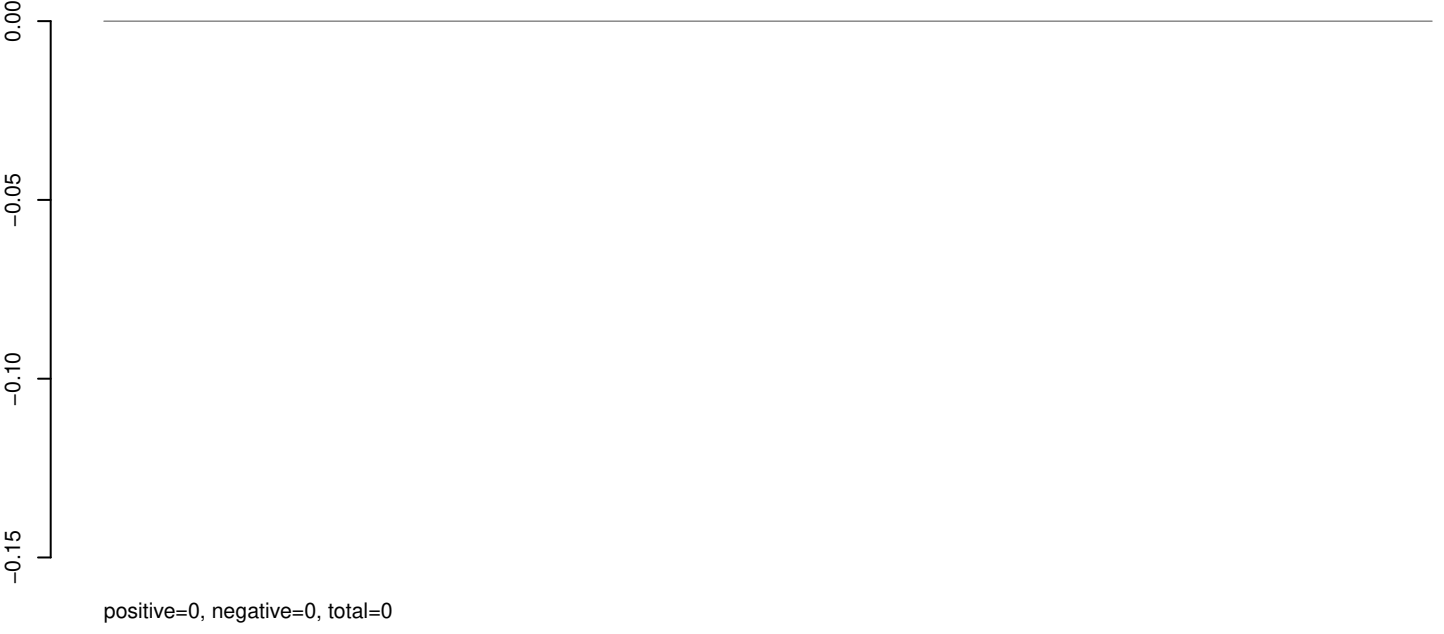
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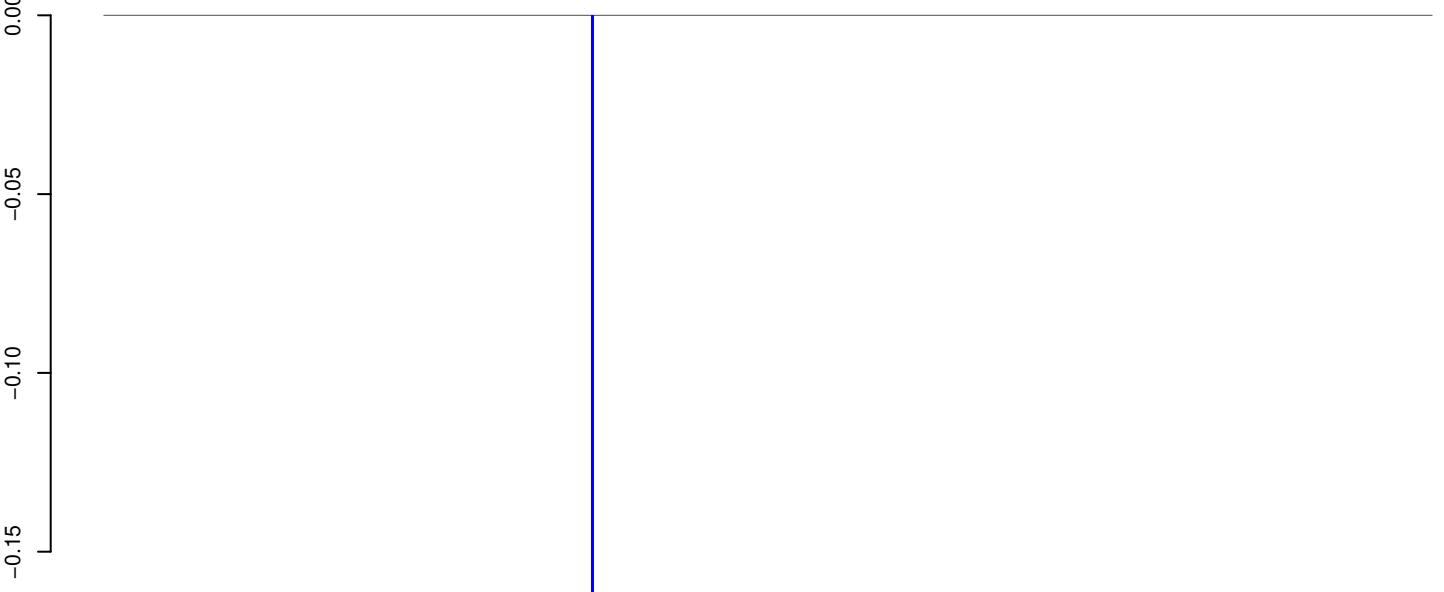
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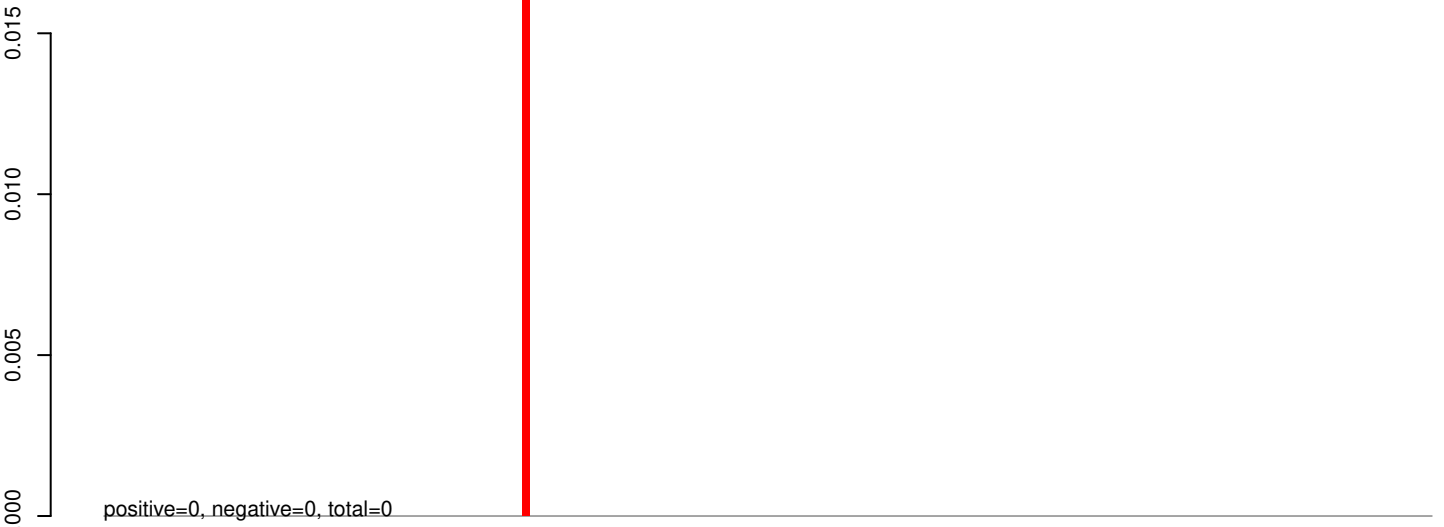
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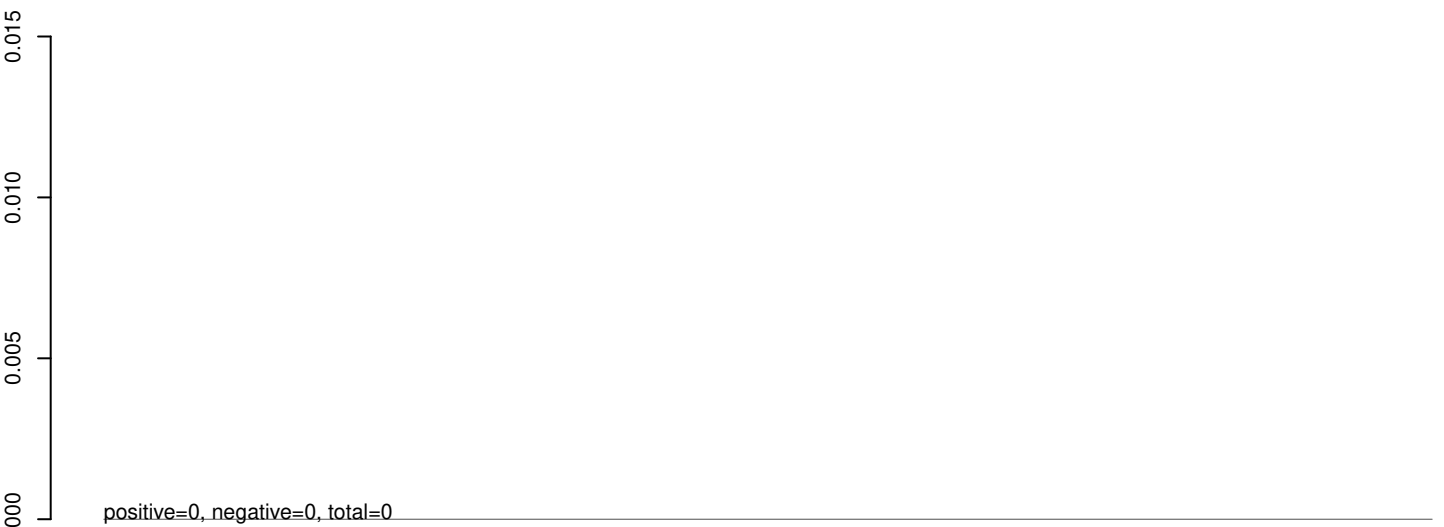
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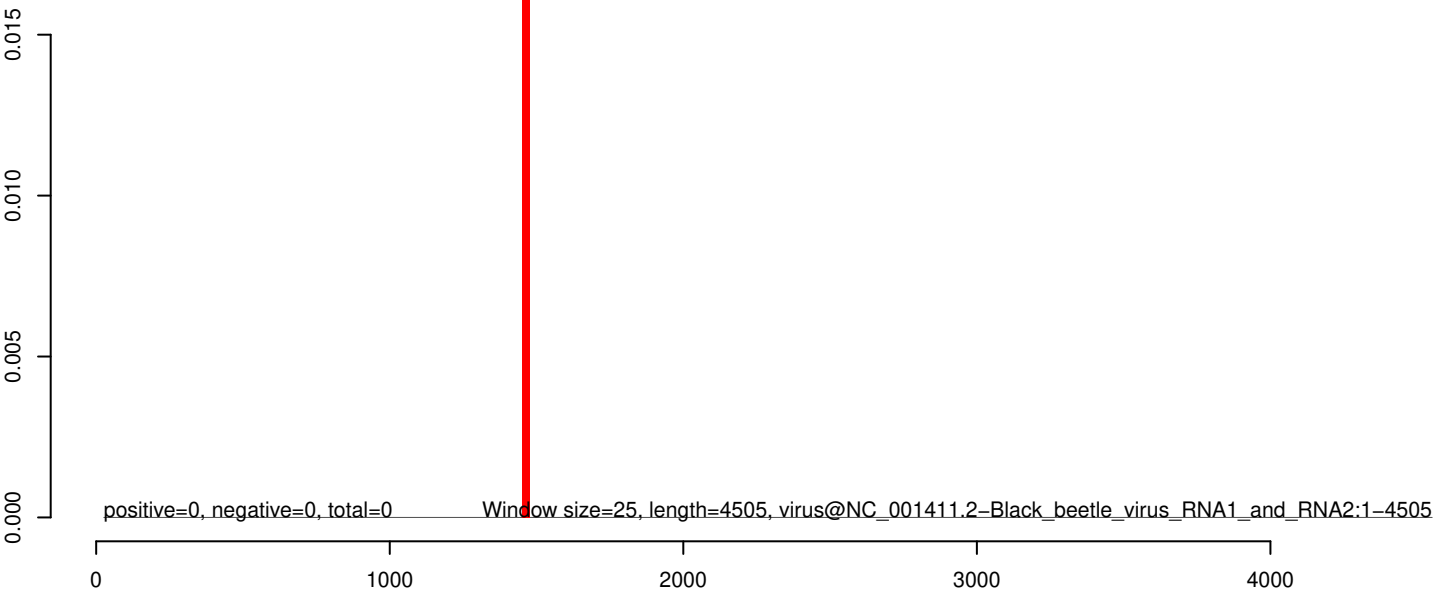
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AeAeg_Aag2_R3D1B_r123_ZA.rep



AeAeg_Aag2_R3D1B_r123_ZA.18_23.rep

0.000
-0.005
-0.010
-0.015

positive=0, negative=0, total=0

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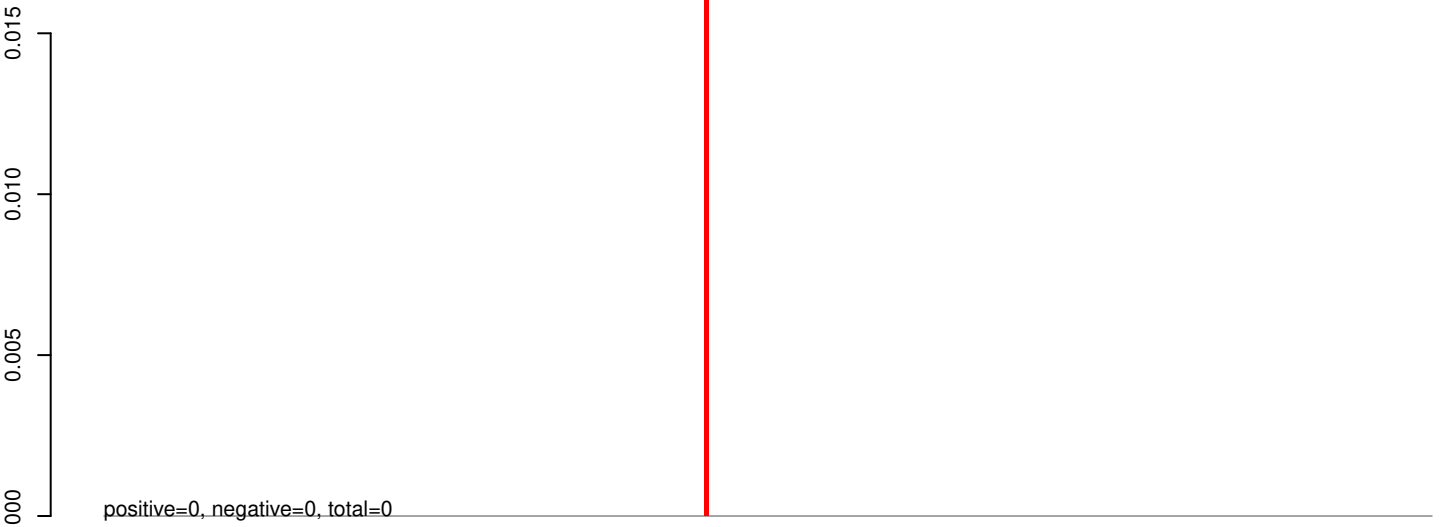
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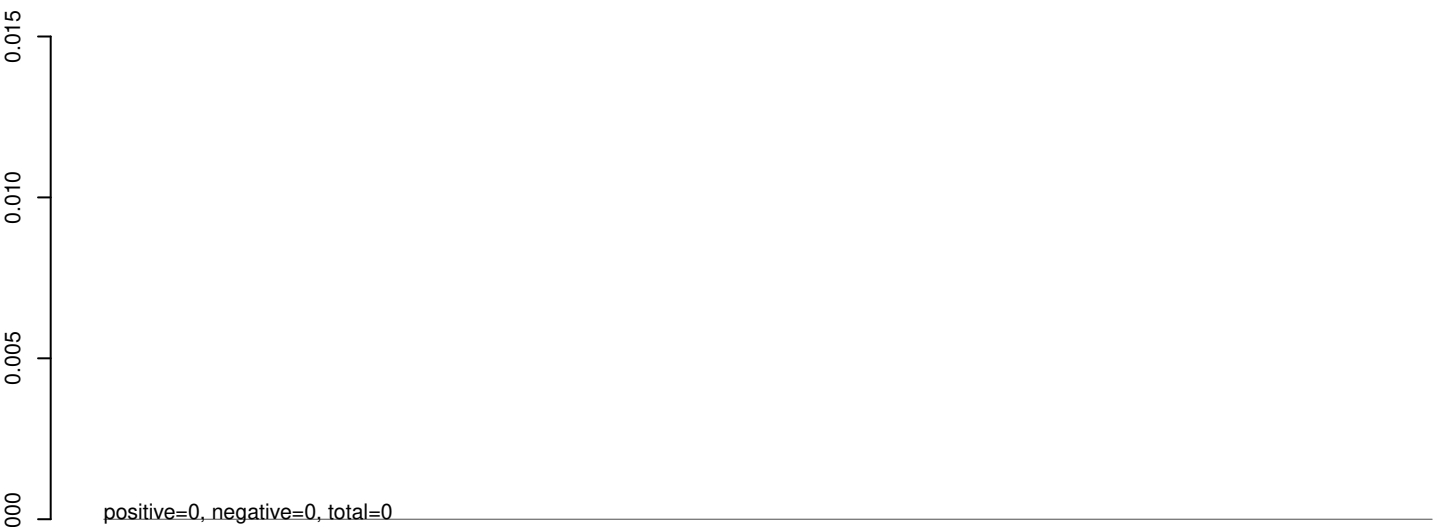
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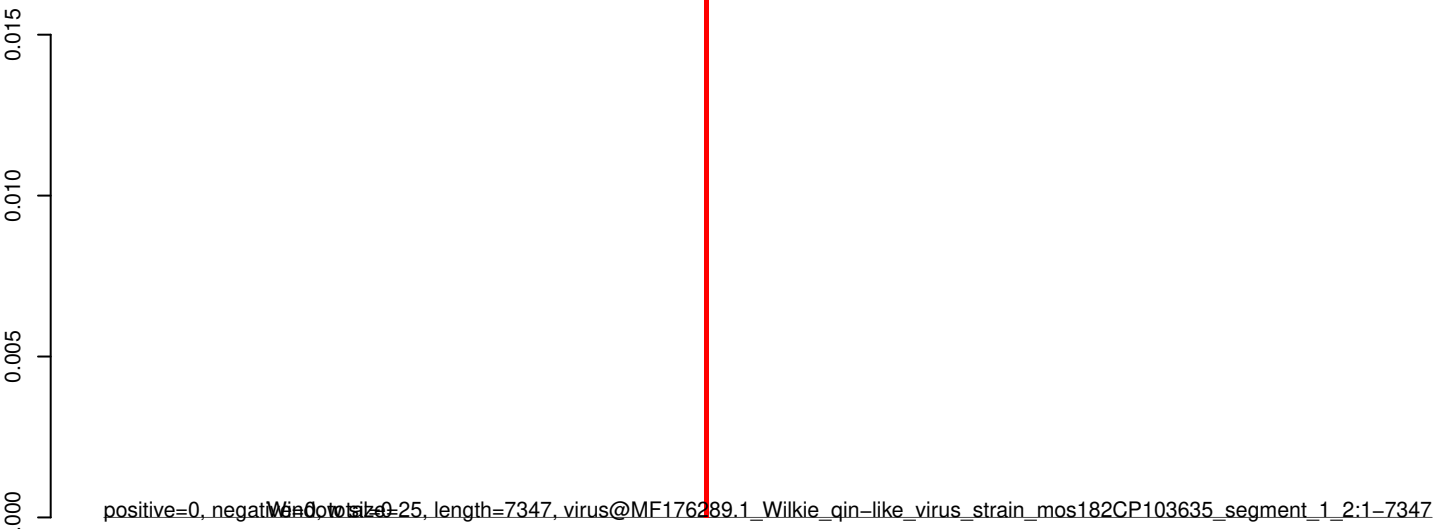
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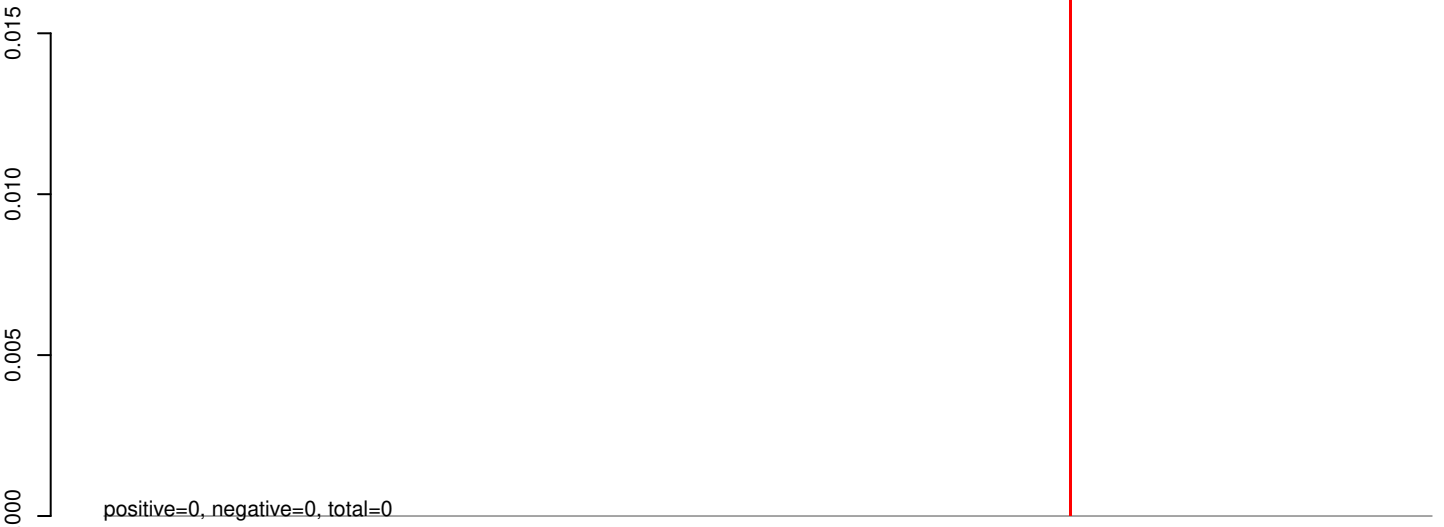
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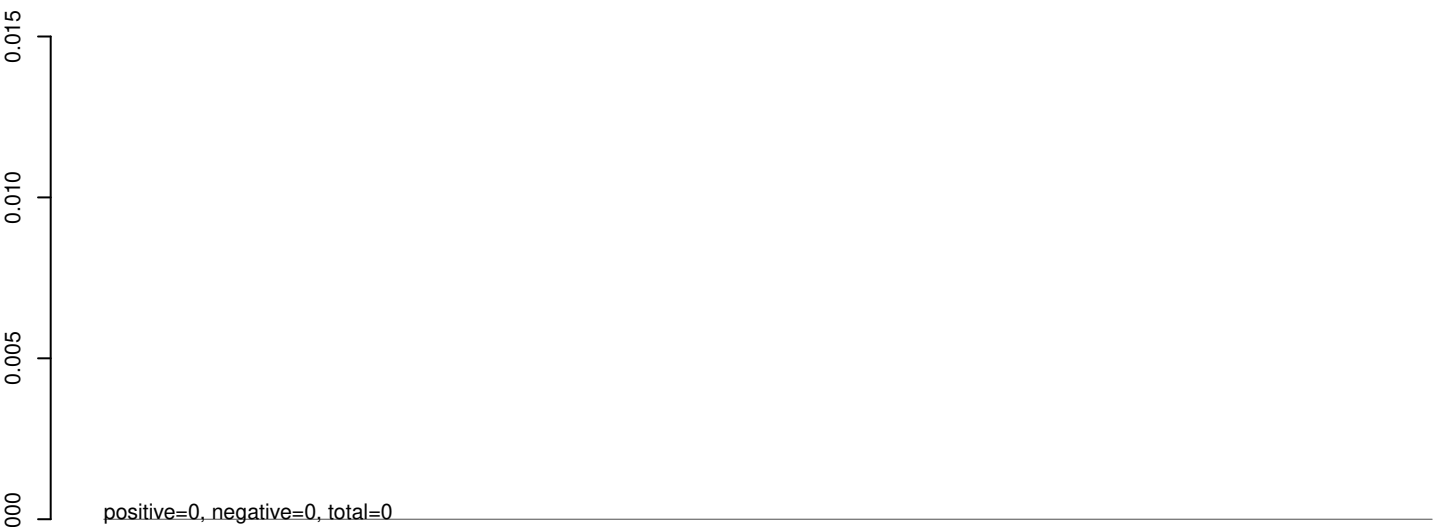
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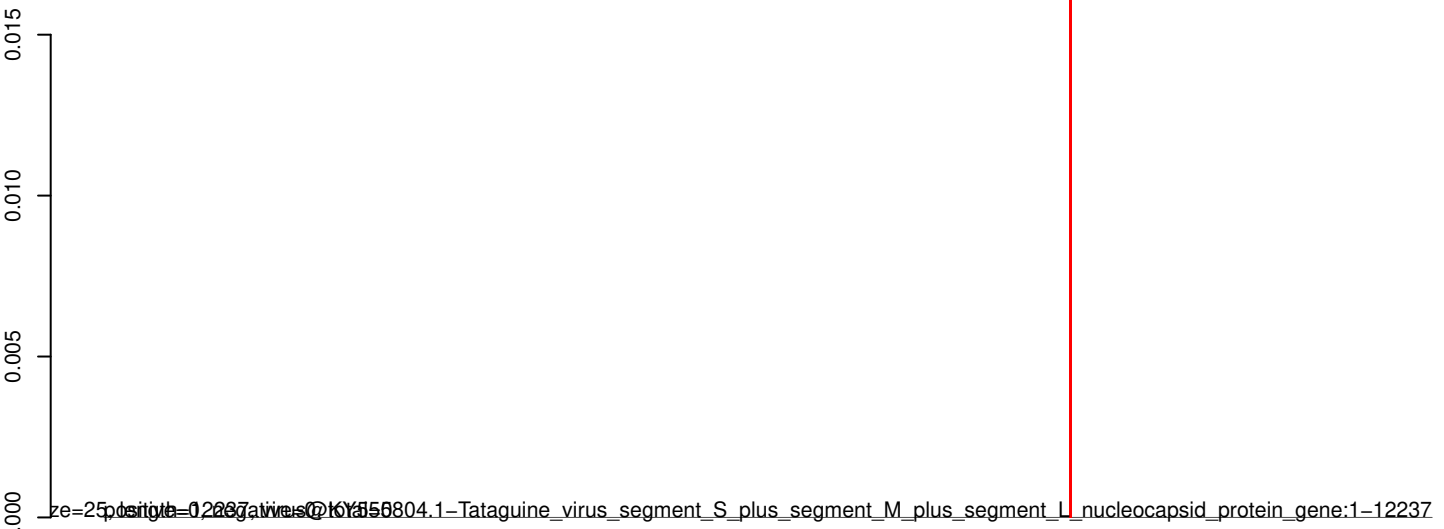
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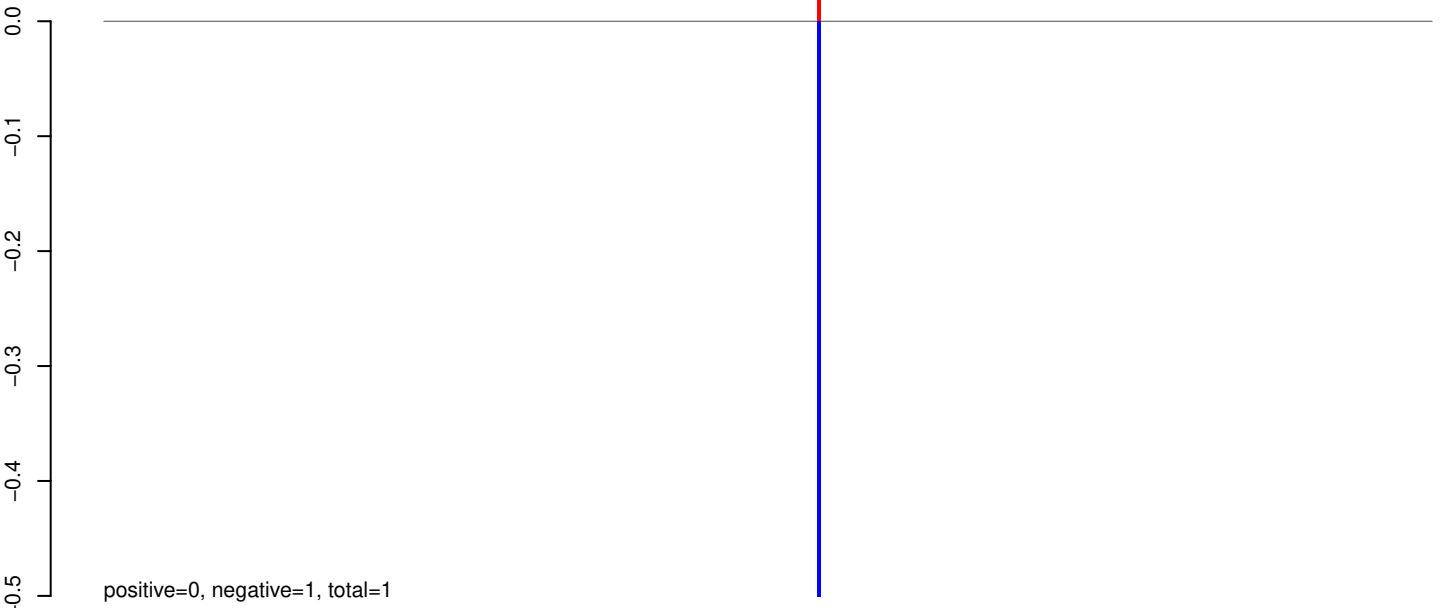
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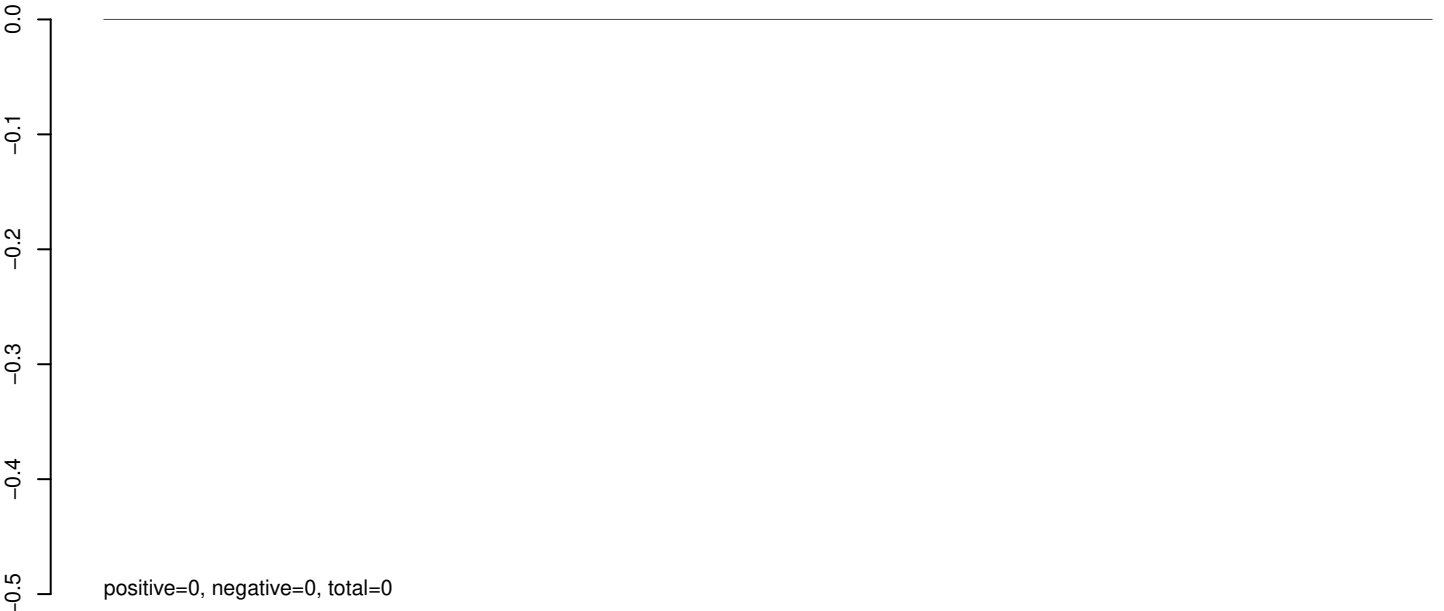
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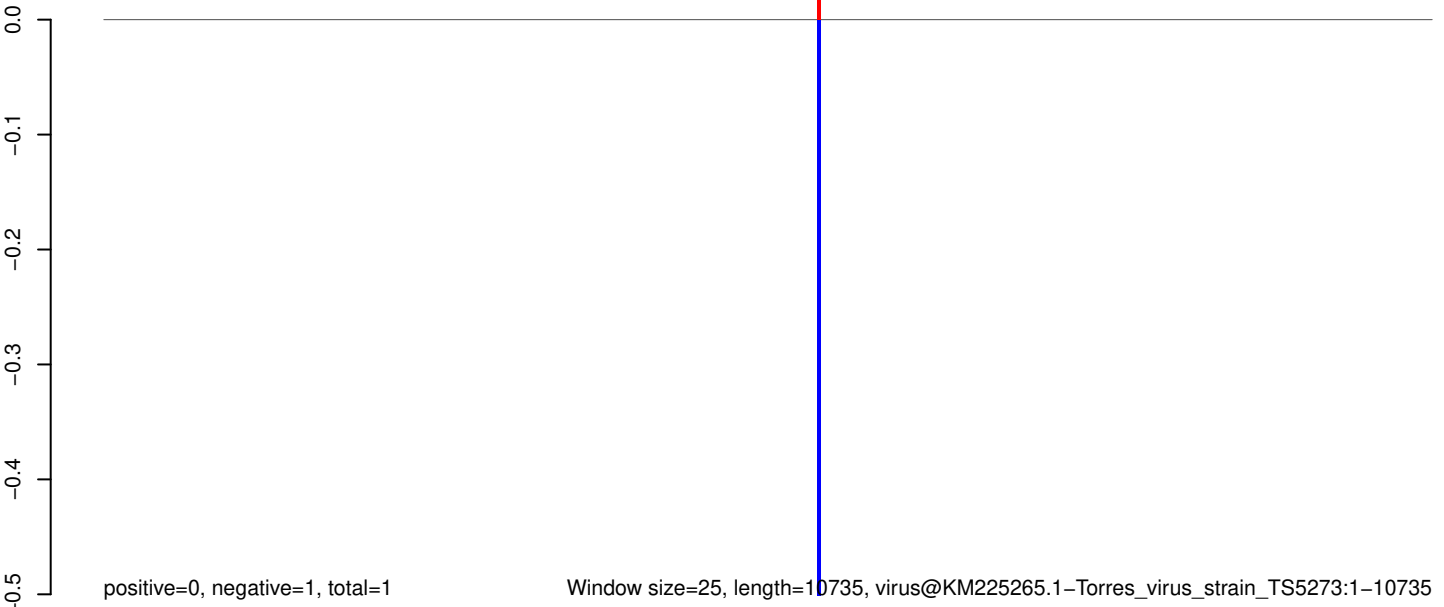
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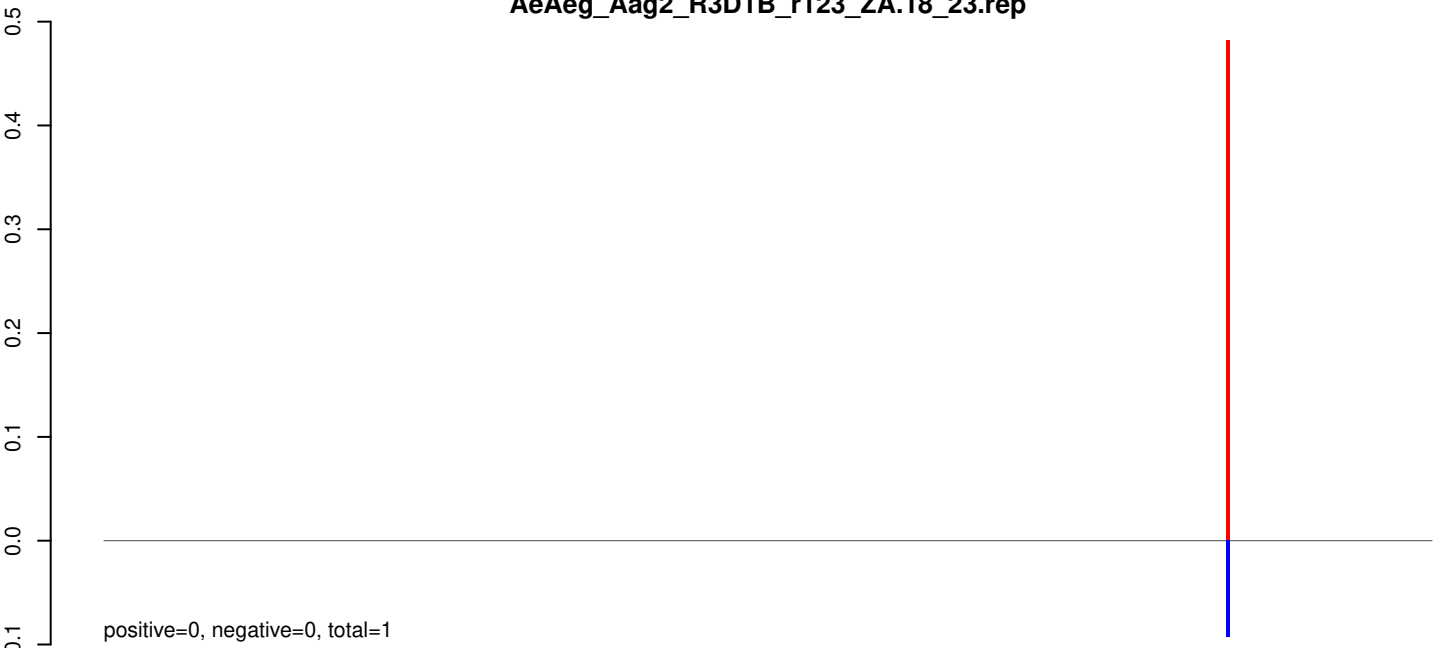
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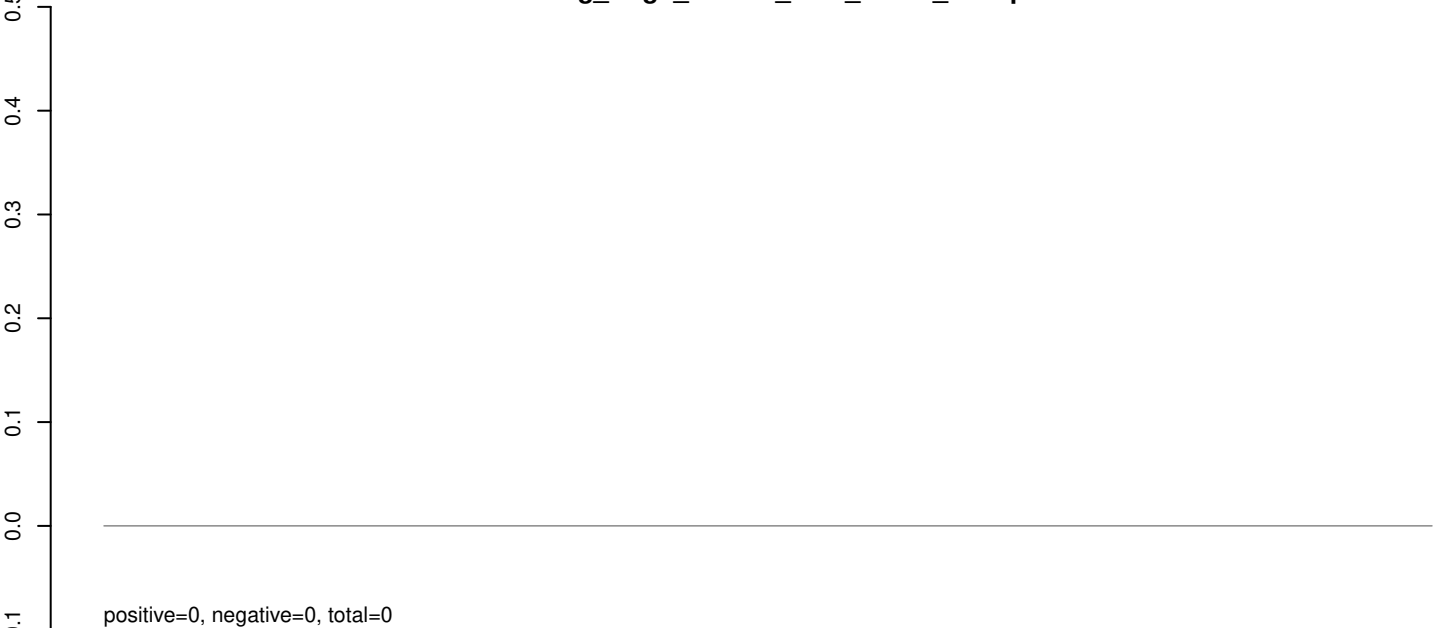
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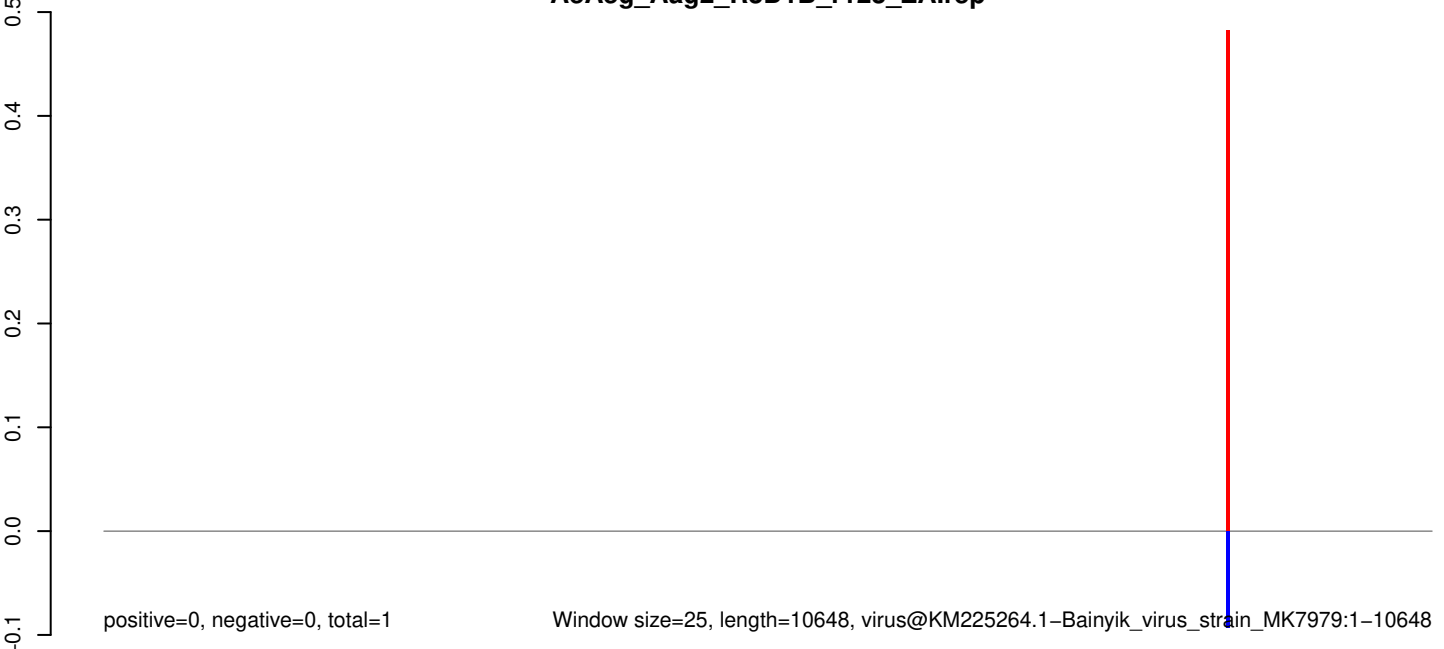
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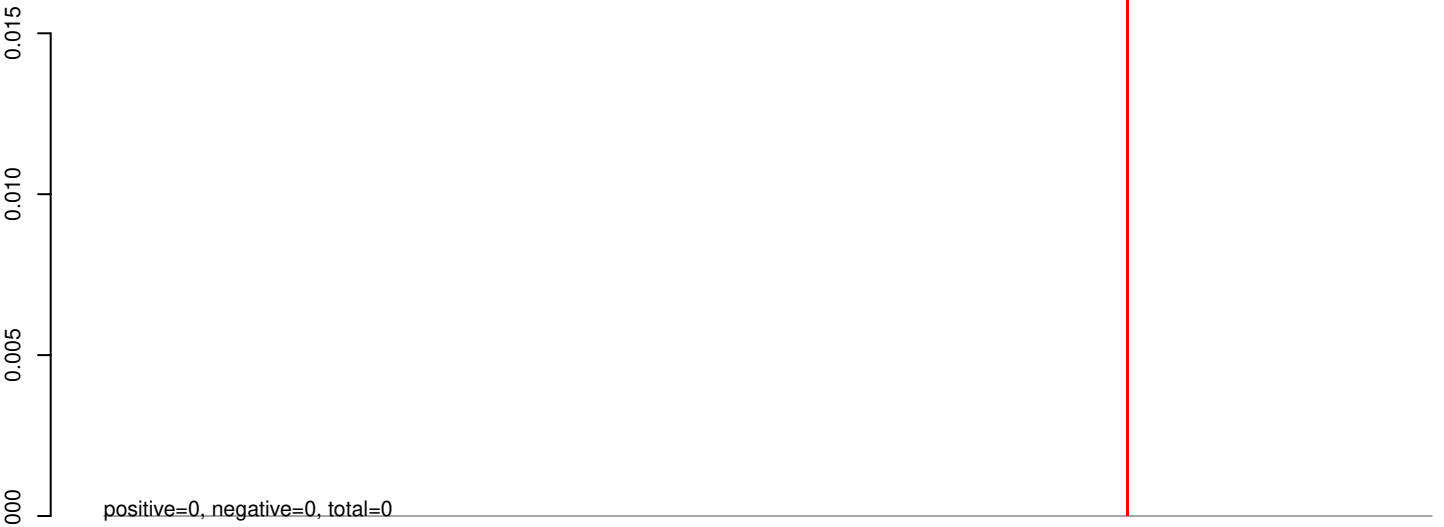
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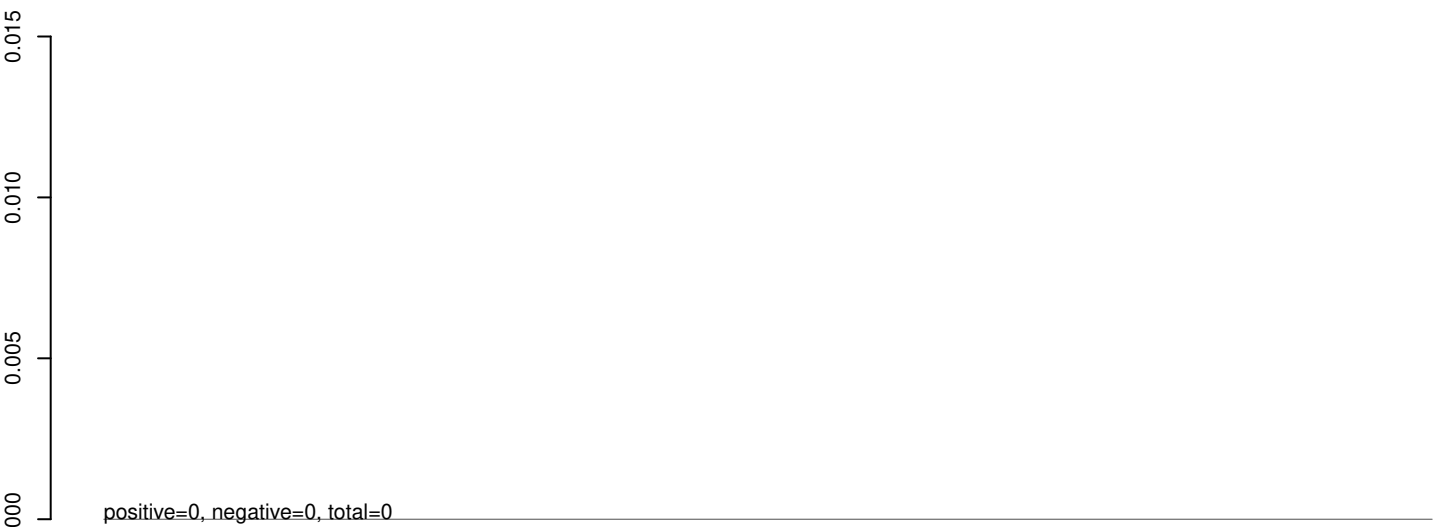
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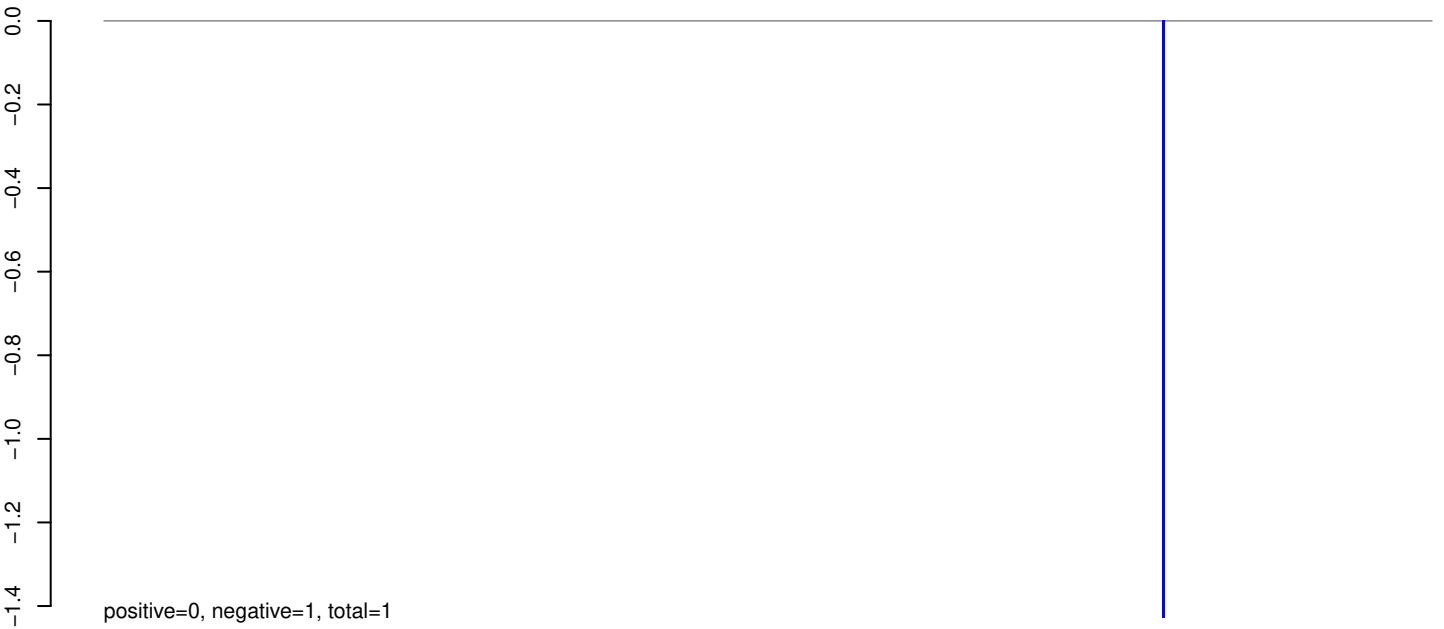
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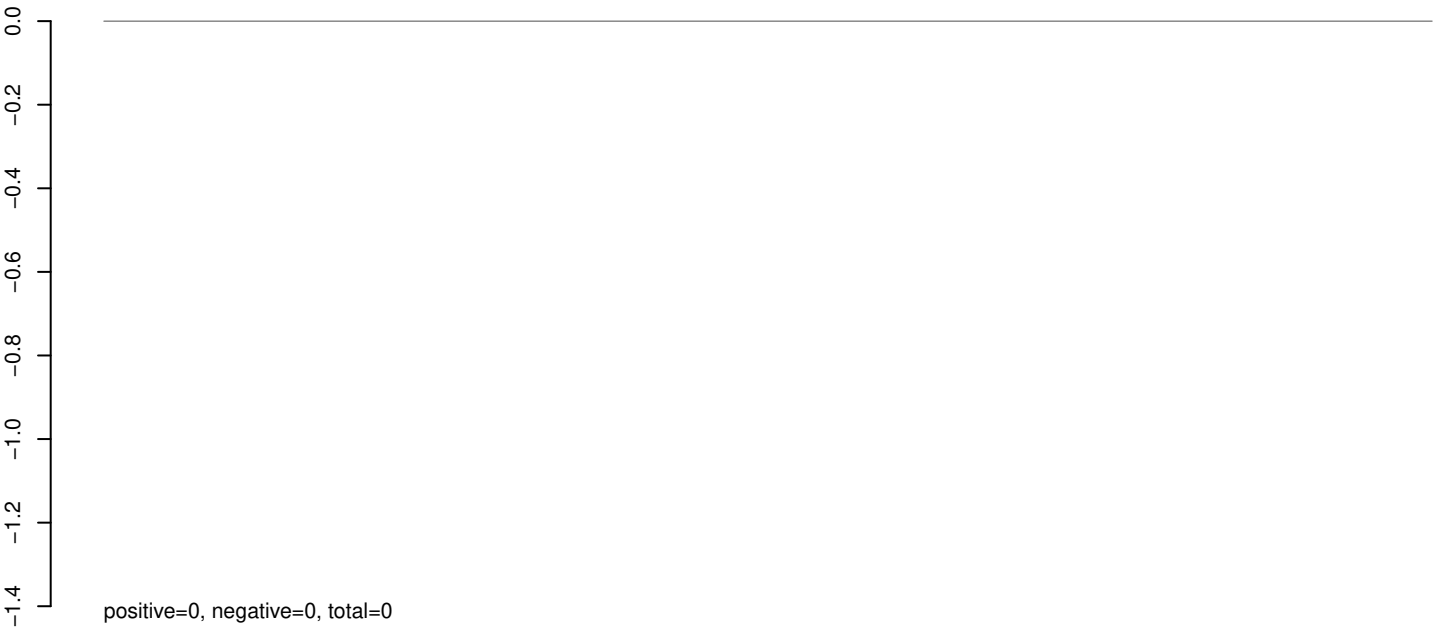
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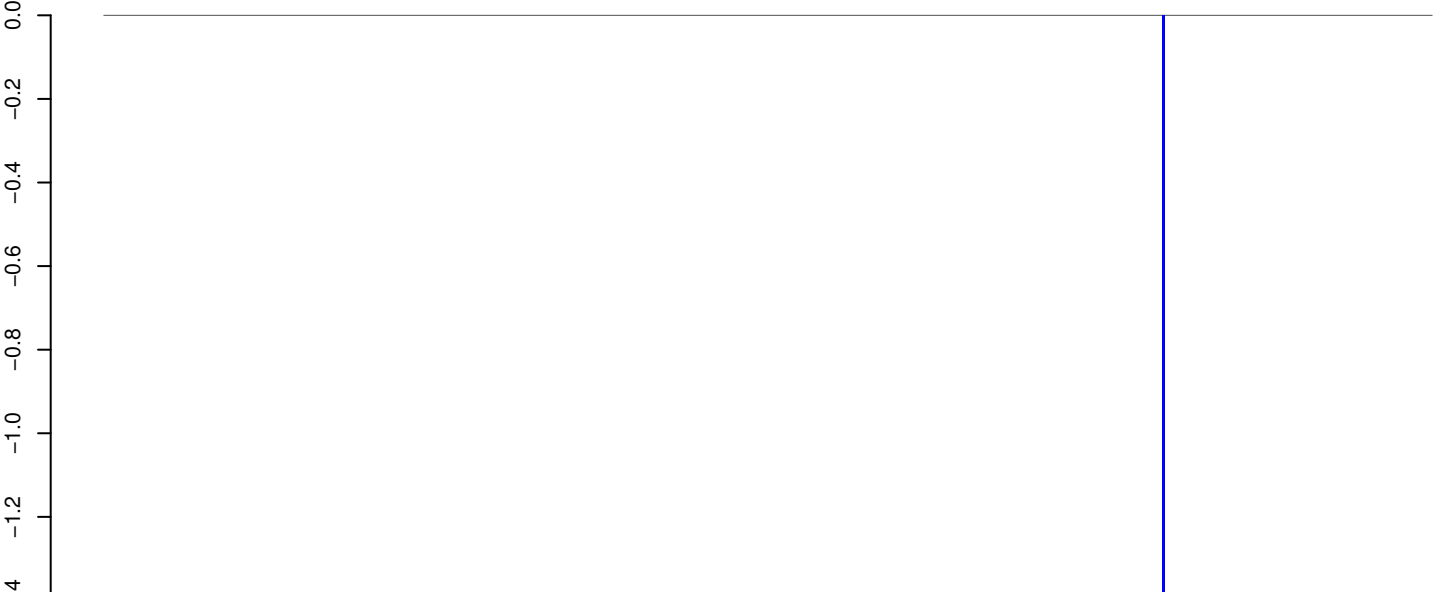
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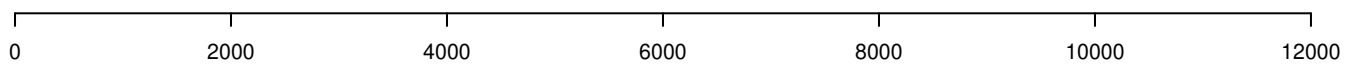
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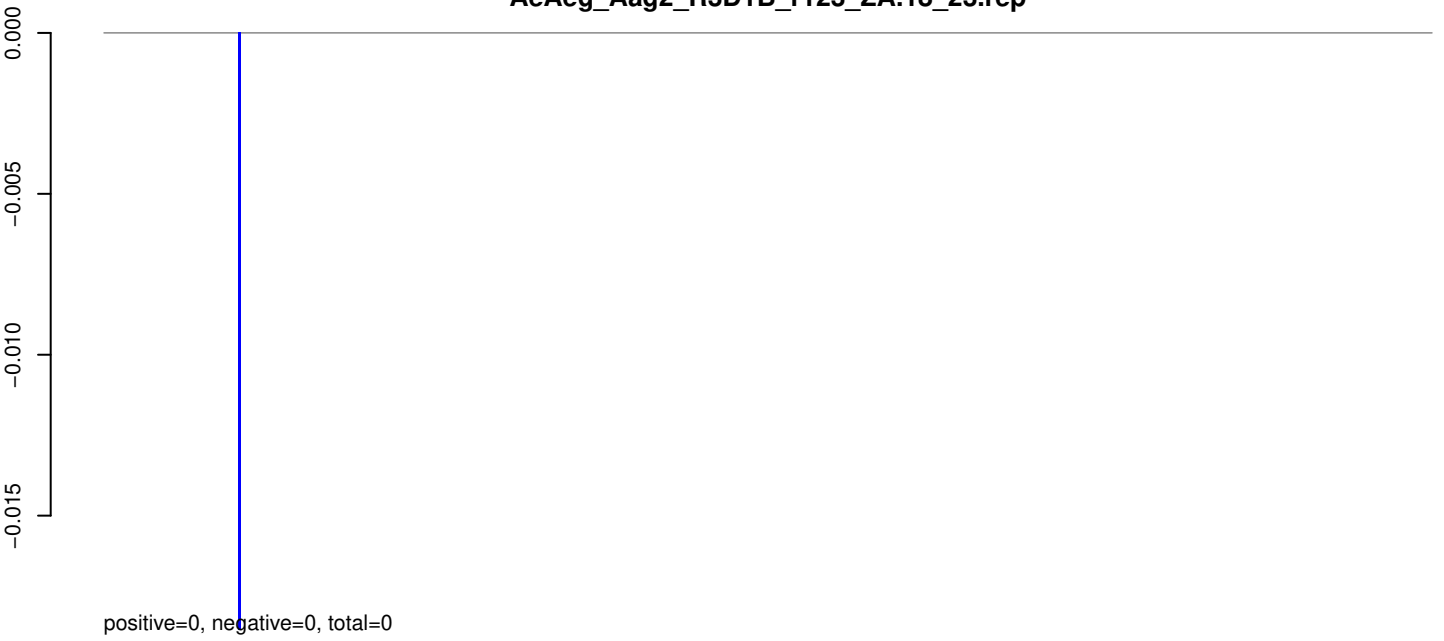
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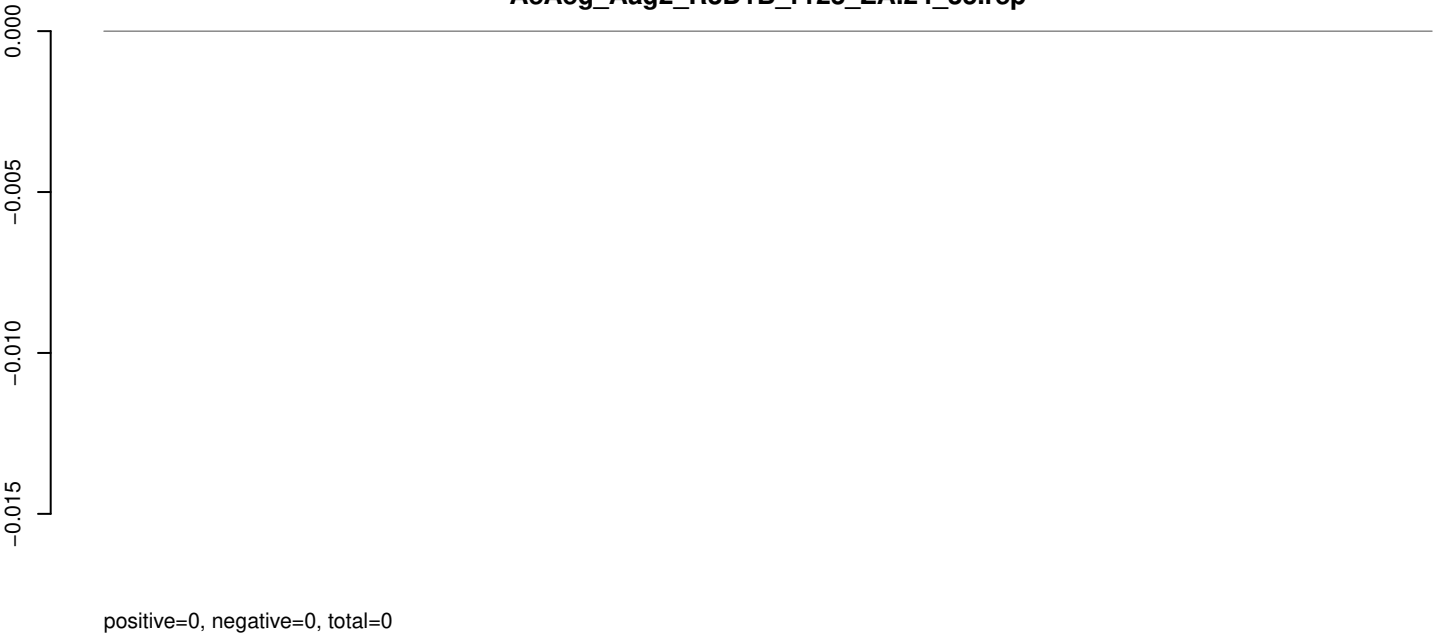
Winnipeg, Manitoba, Canada, 2015-2017, Ilesha_virus@KC608151.1-Ilesha_virus_strain_ILESHA_8e_segment_S_plus_segment_M_plus_segment_L:1-12297



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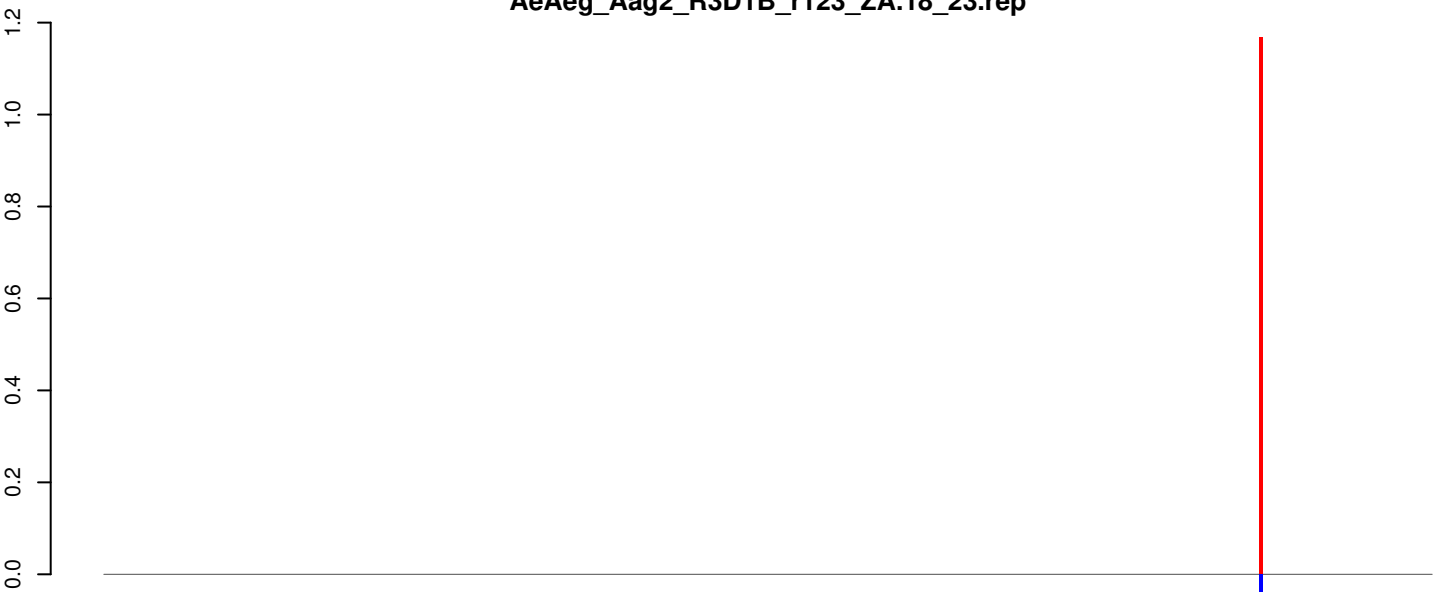
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downloaded from <https://www.ncbi.nlm.nih.gov/trace/viral/12052/> on 12/05/2024. HM147991.1-Trocaravirus non-structural polyprotein gene partial cds and structural polyprotein:1-12052

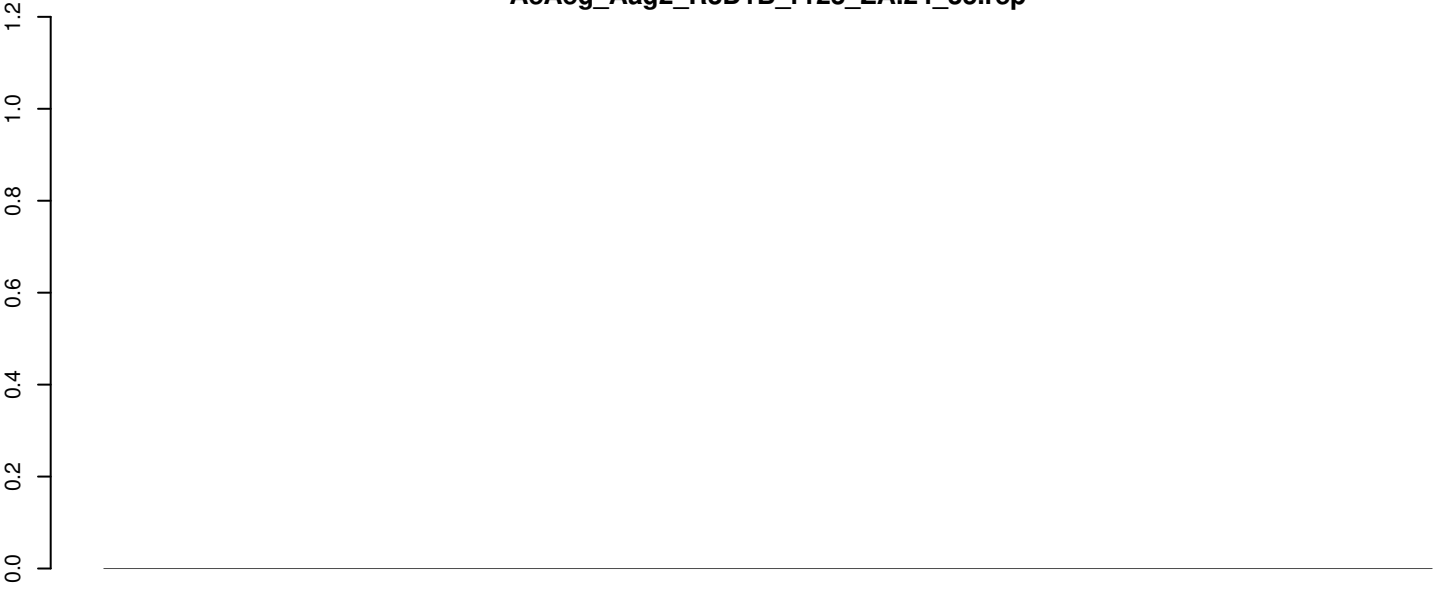
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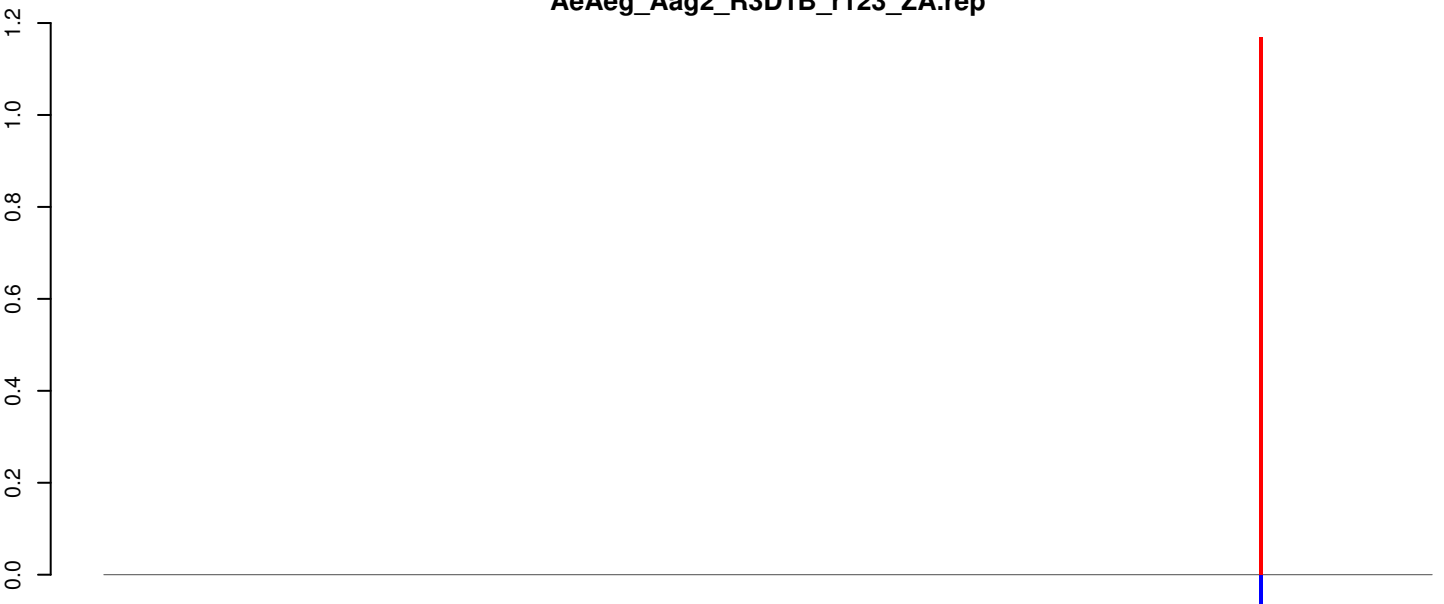
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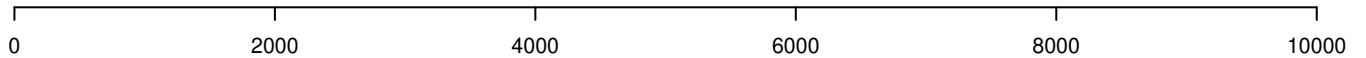


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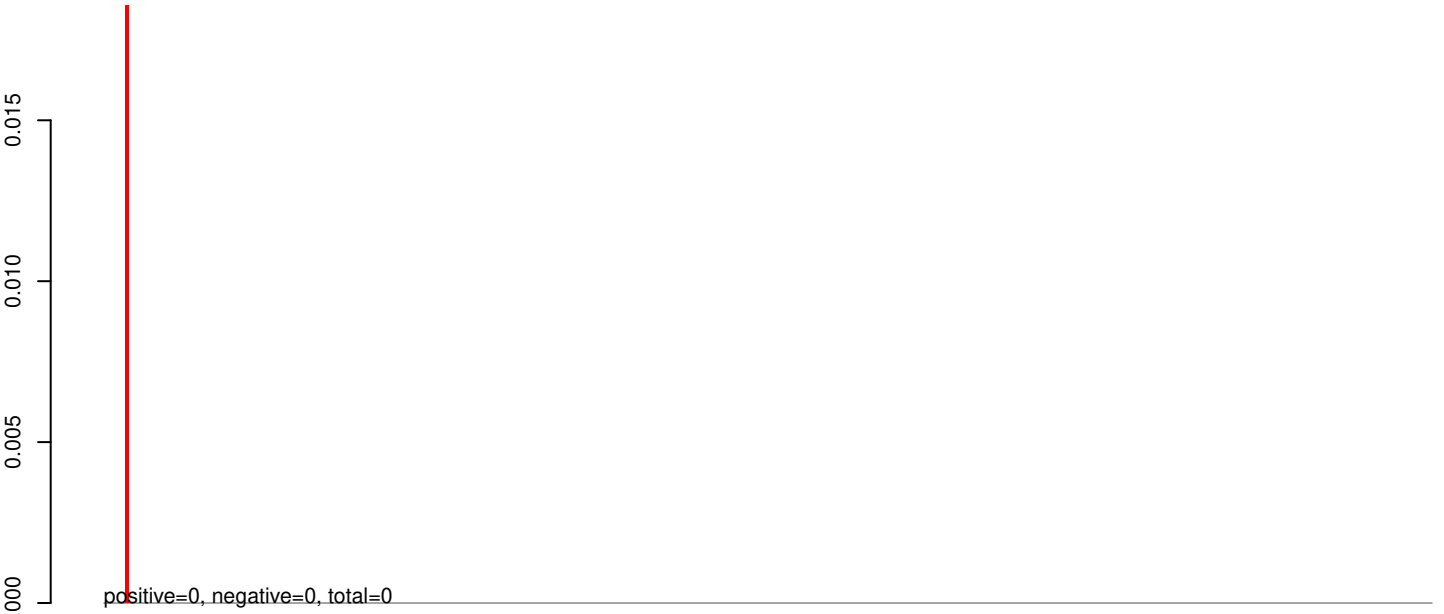
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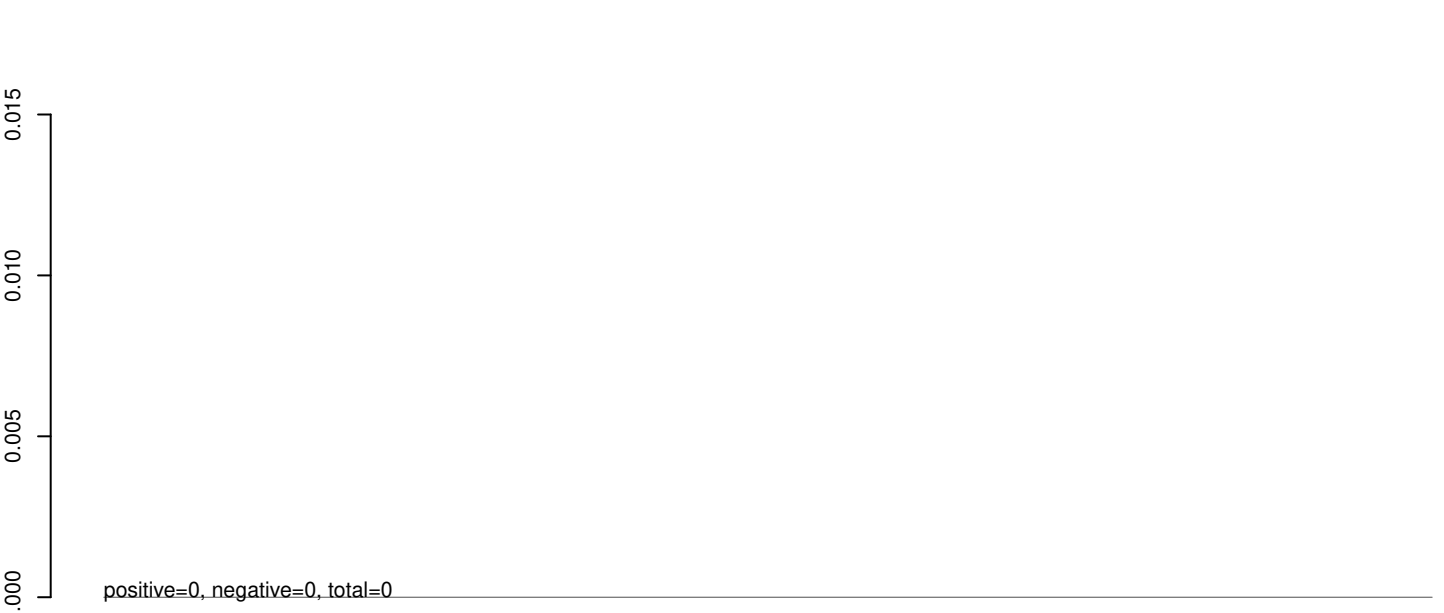
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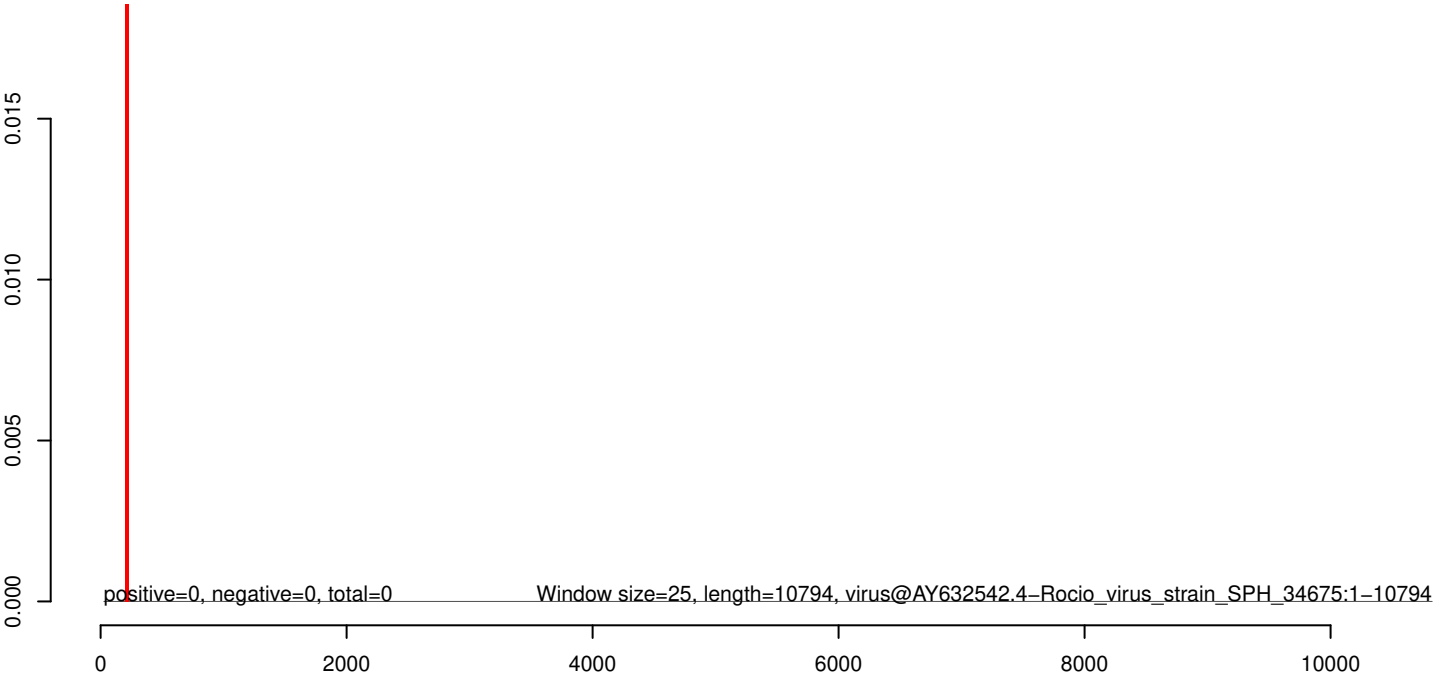
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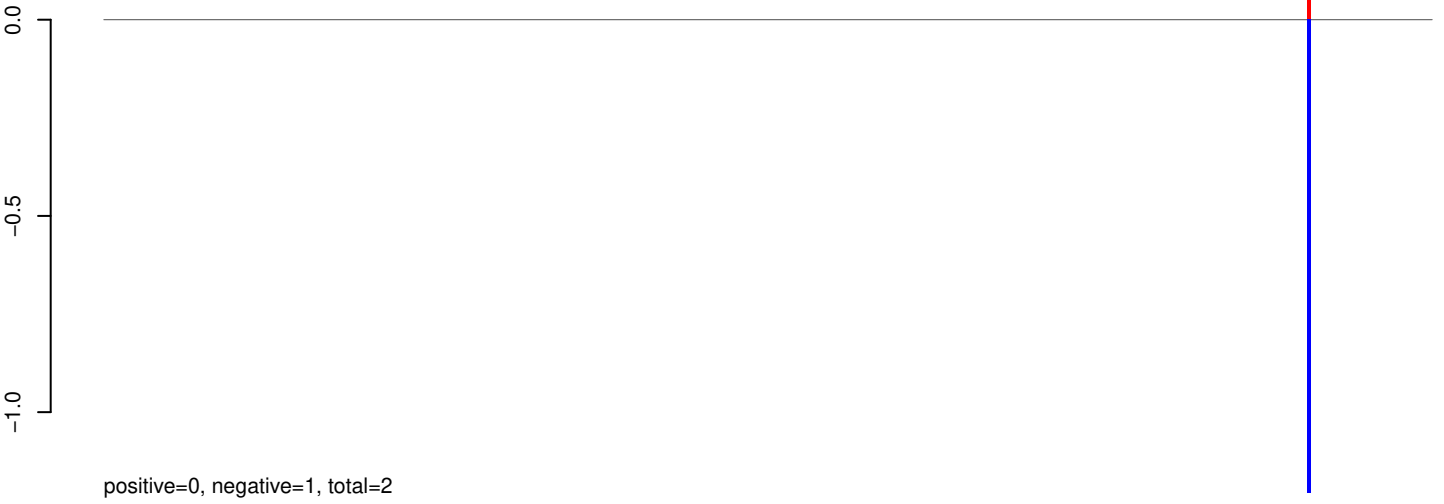
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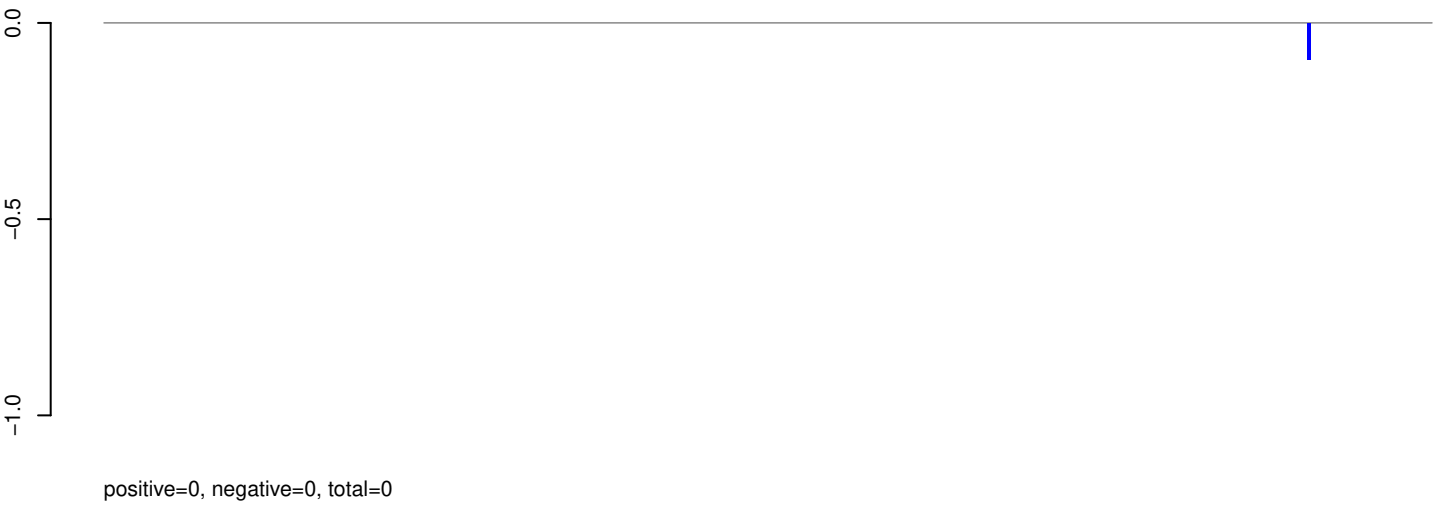
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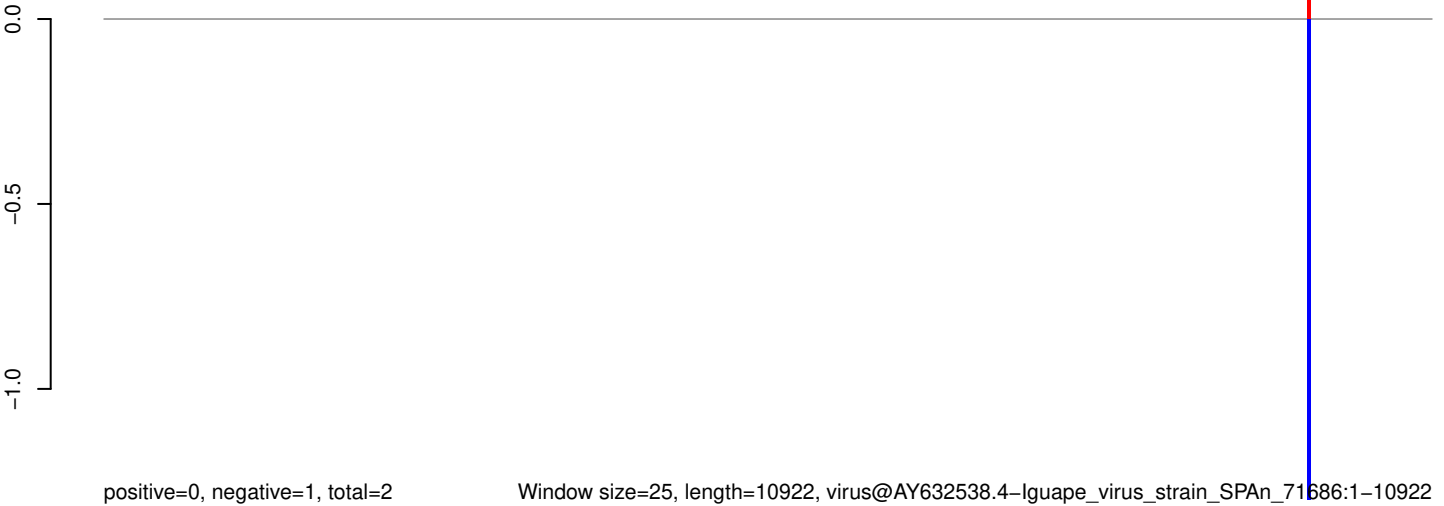
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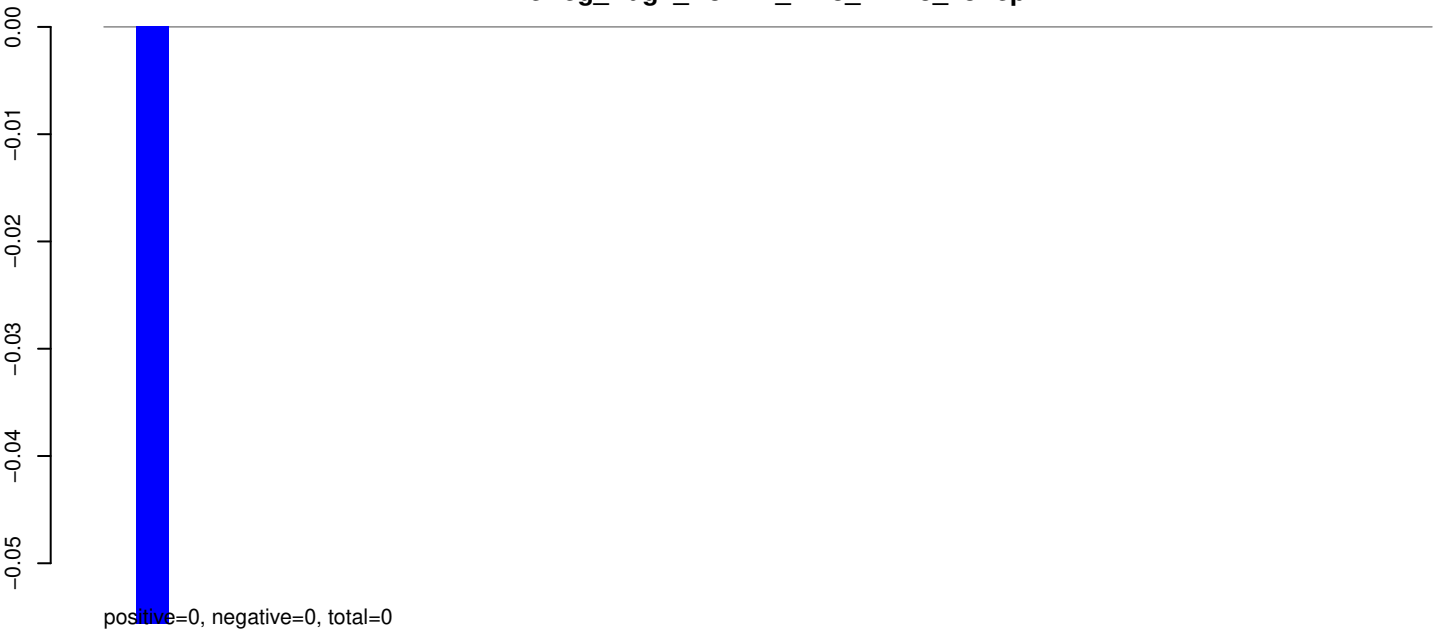
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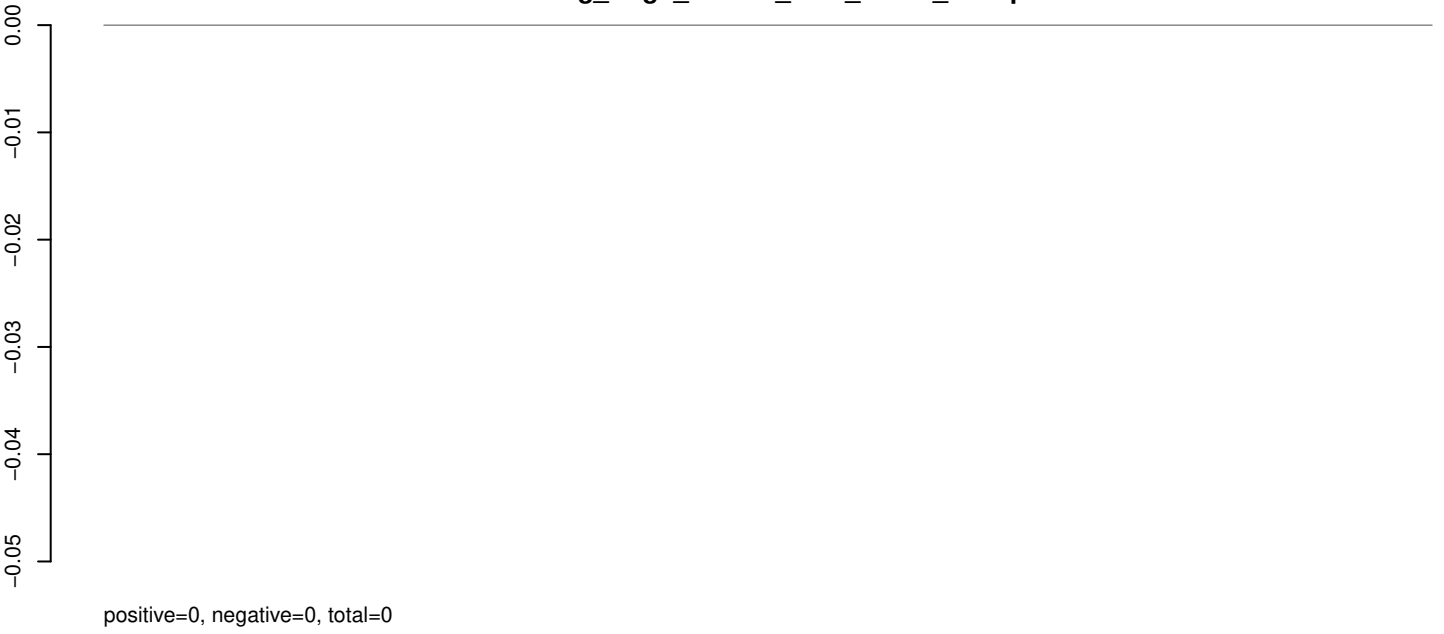
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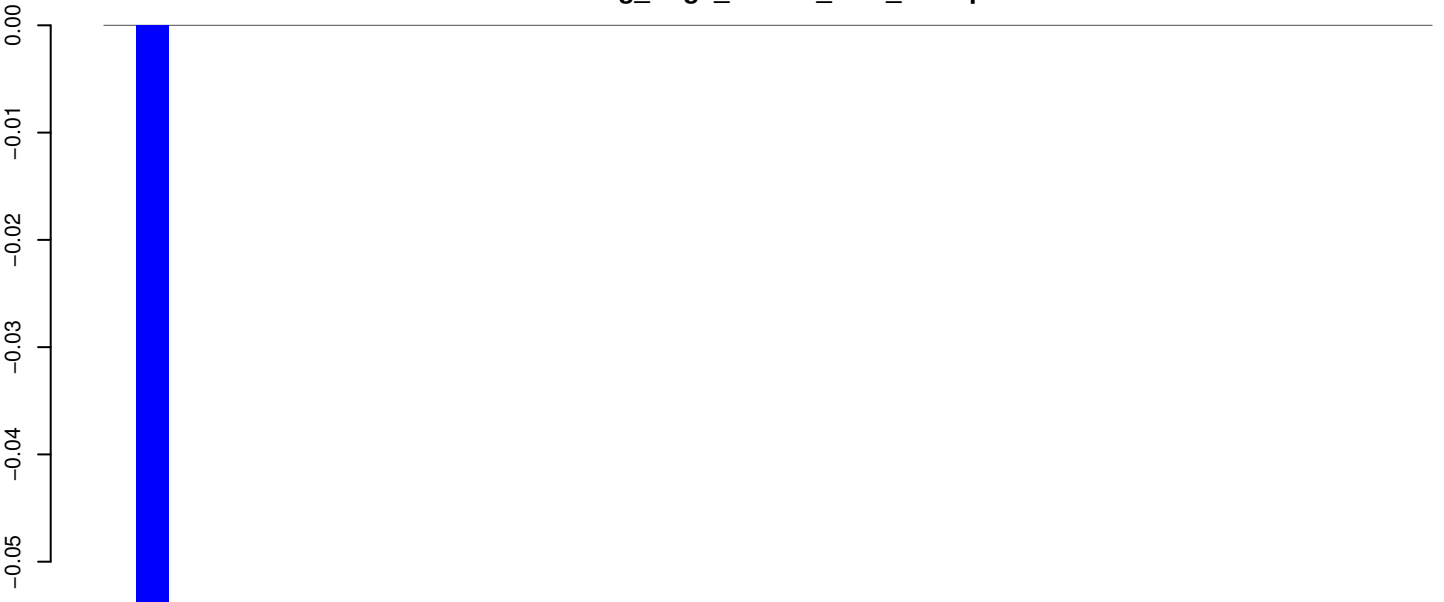
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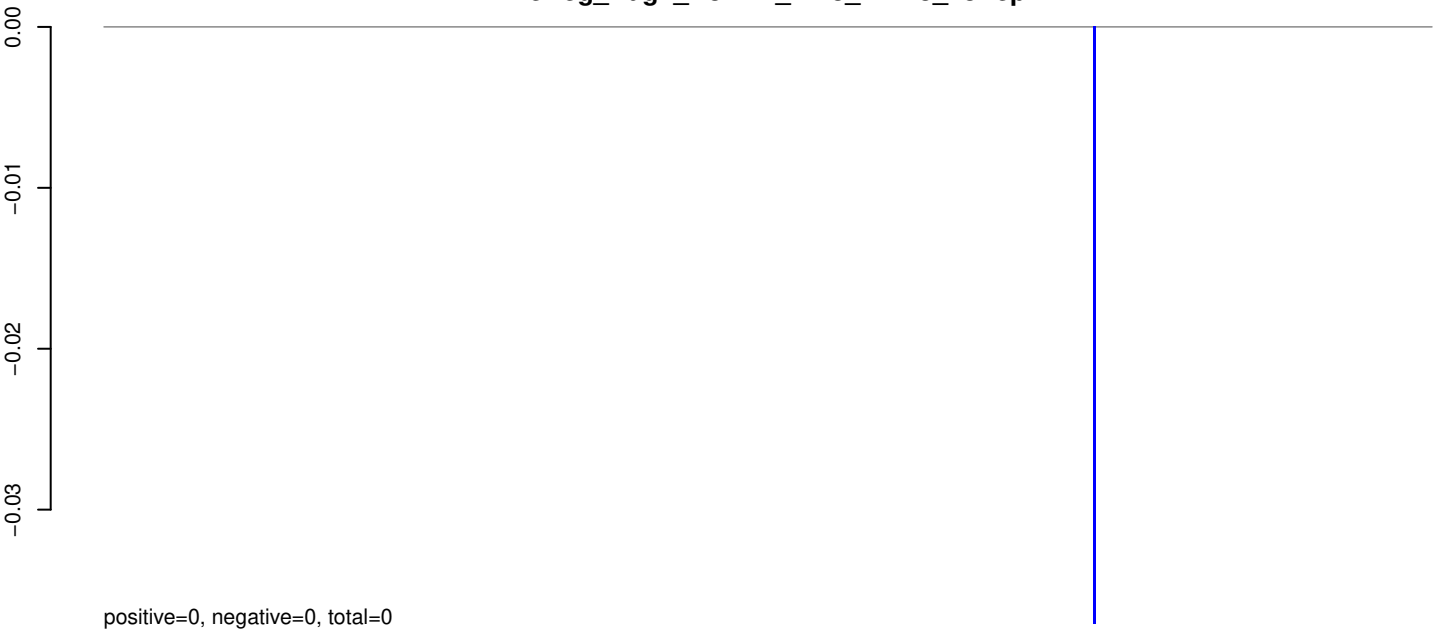
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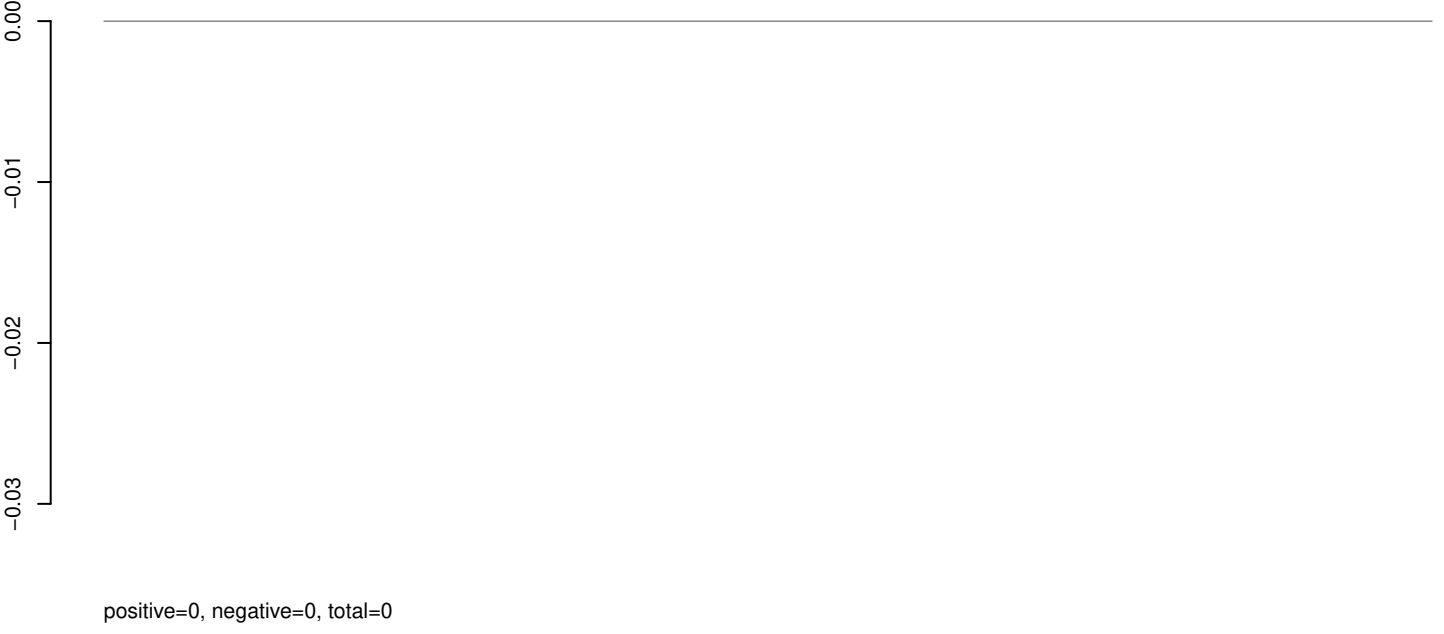


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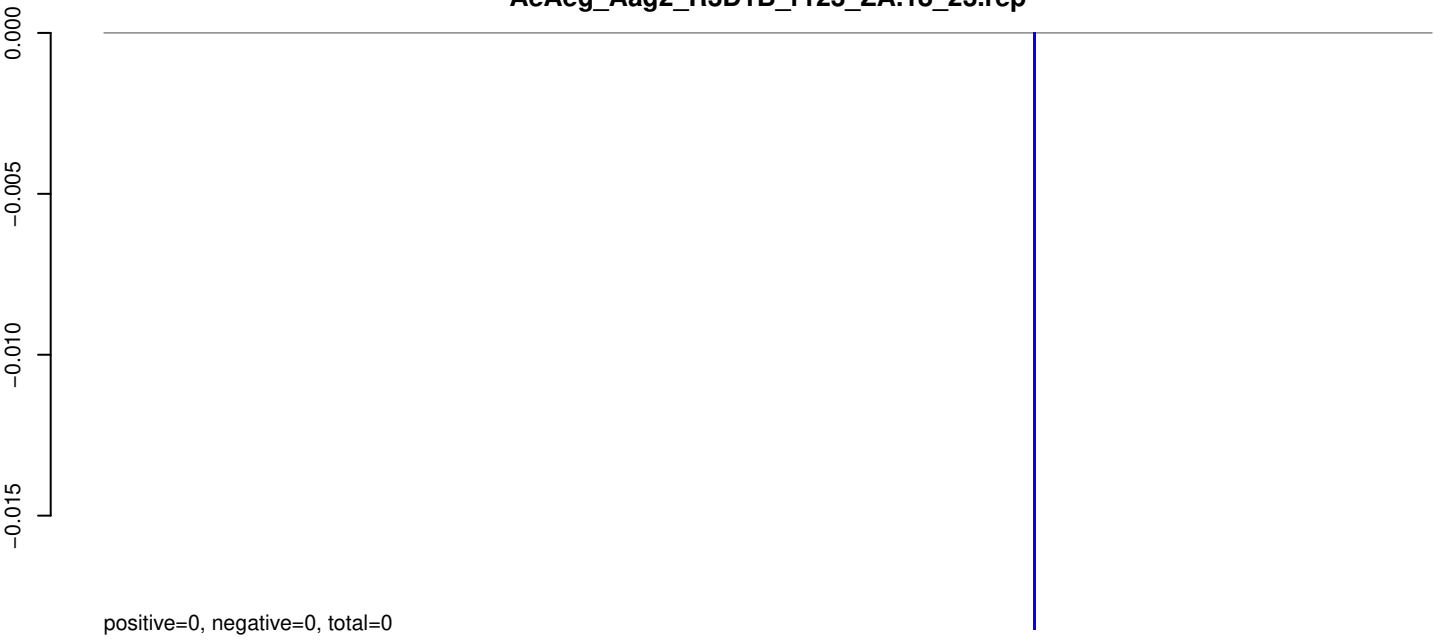


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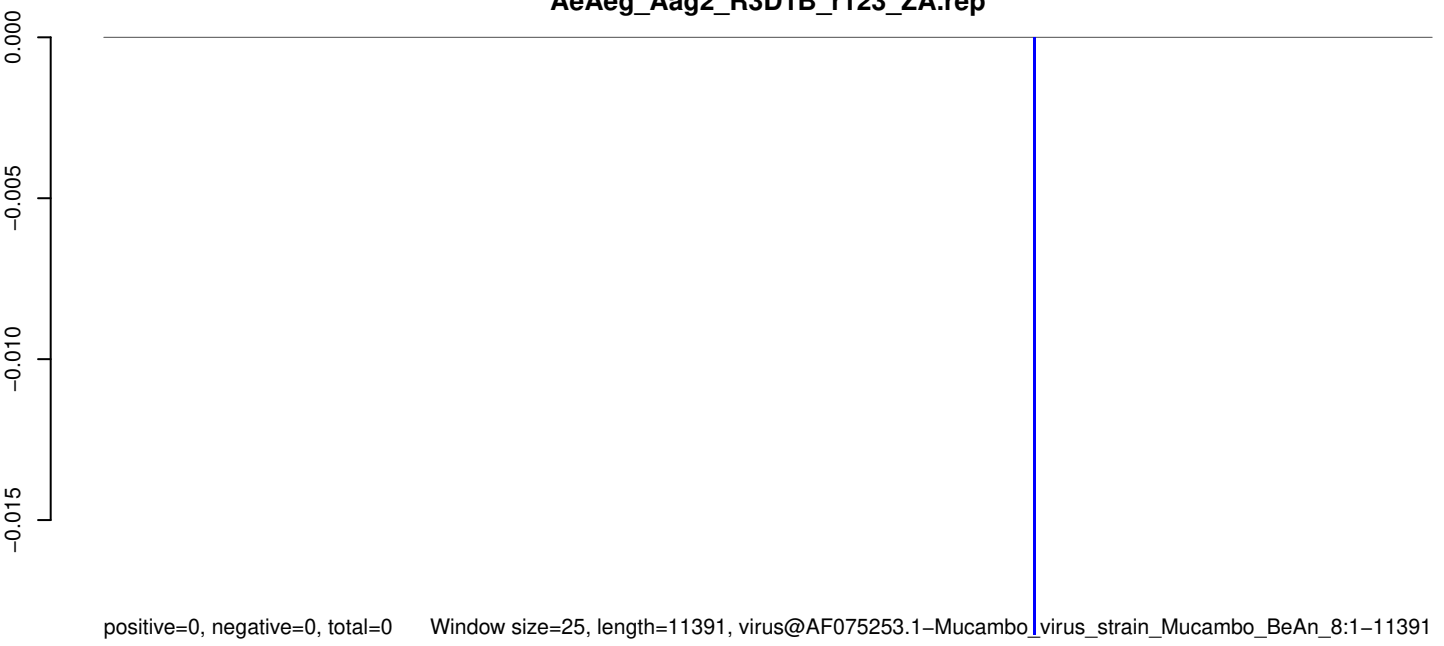
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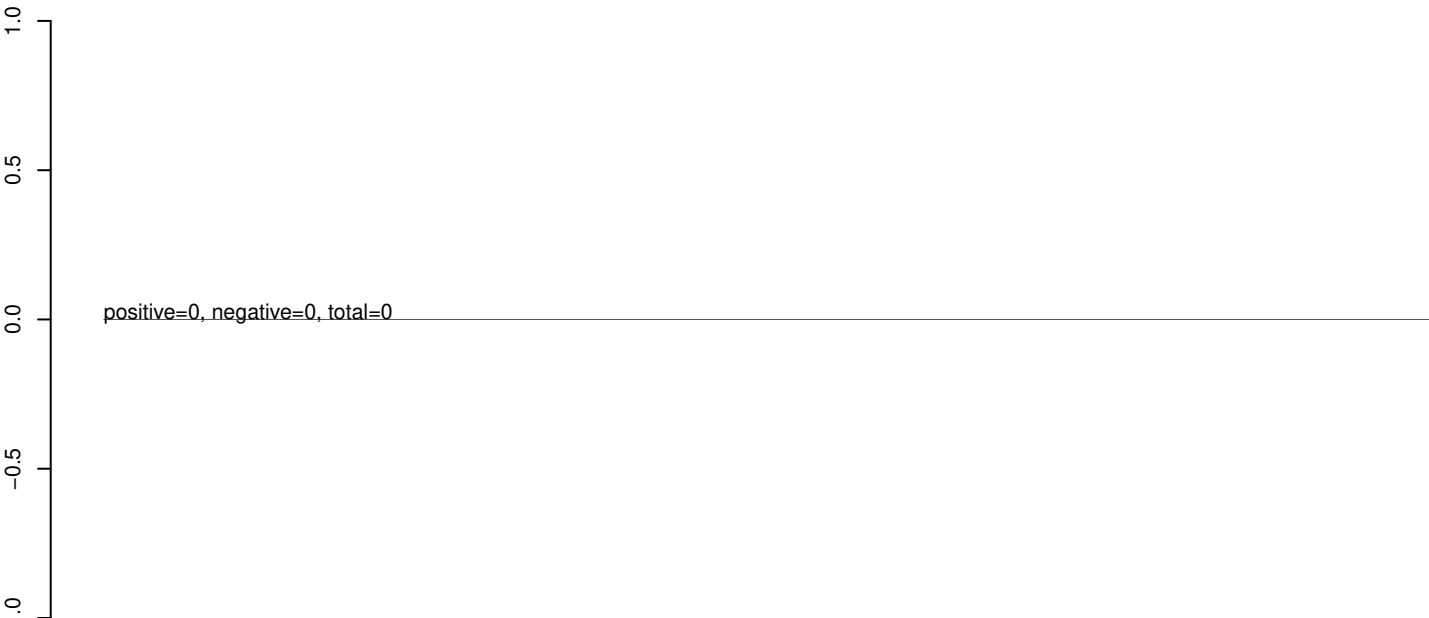
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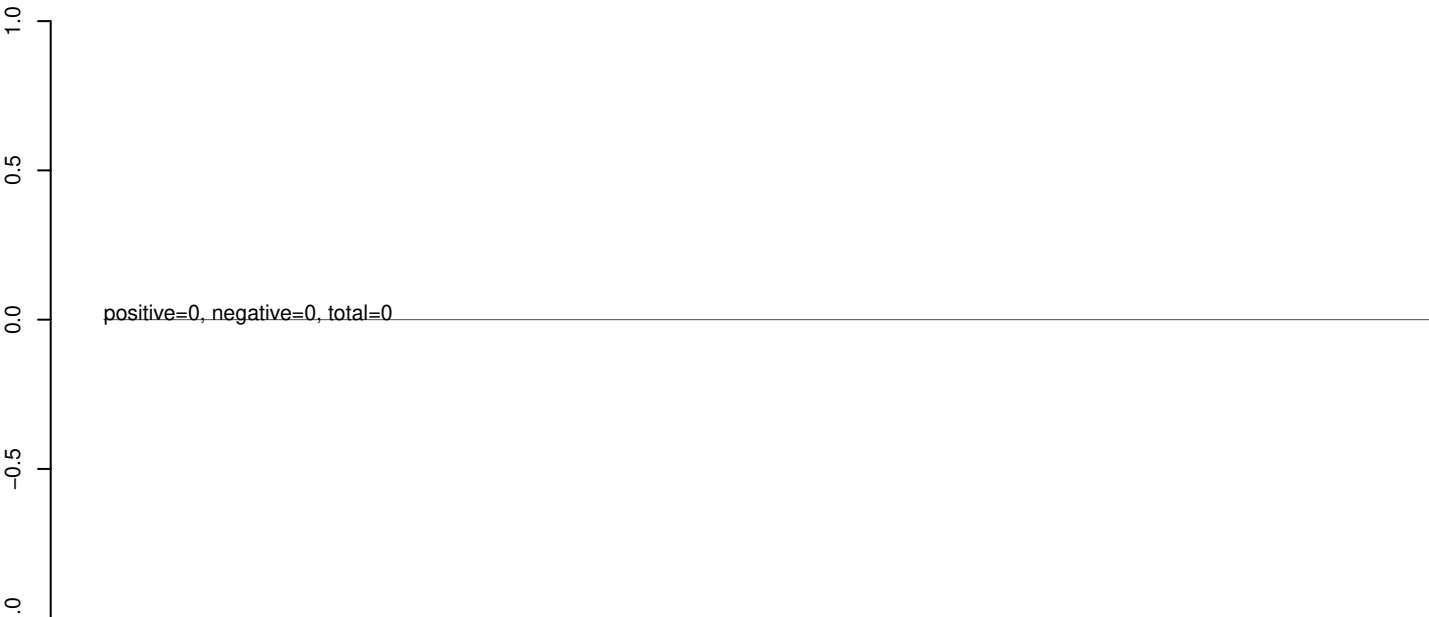
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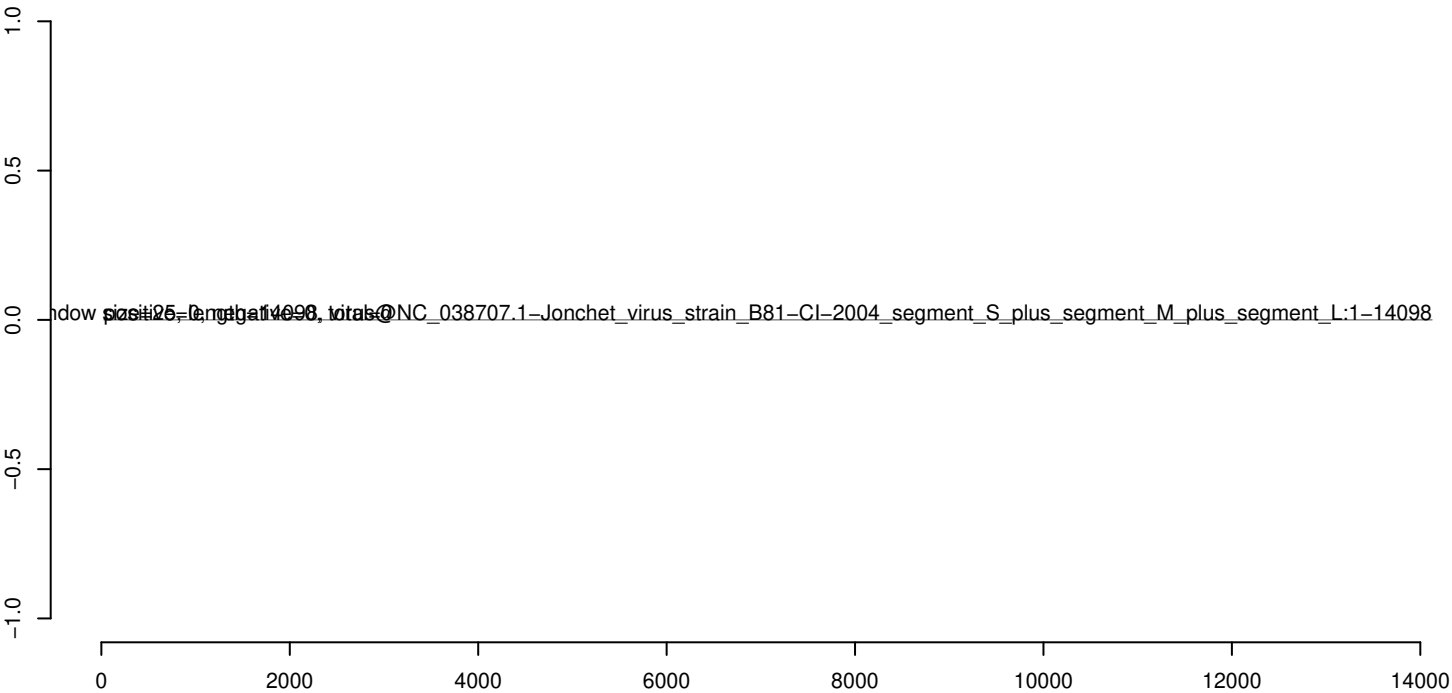
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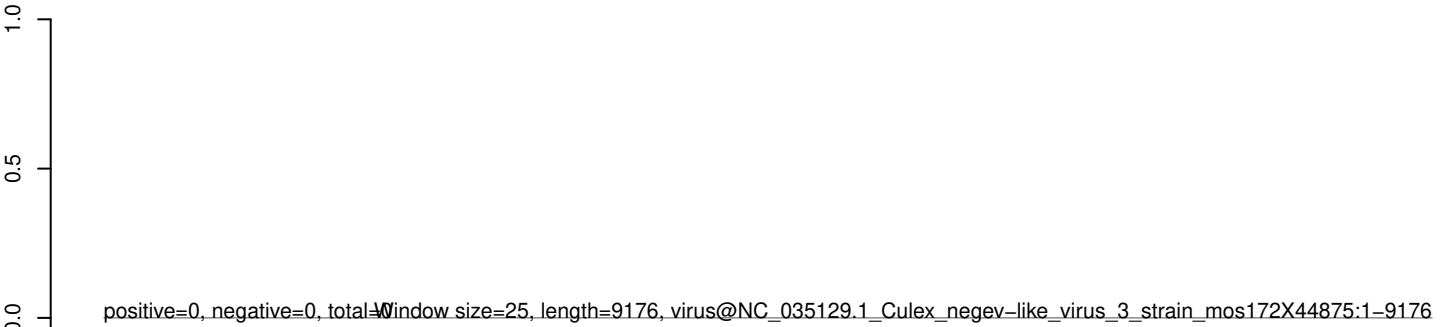
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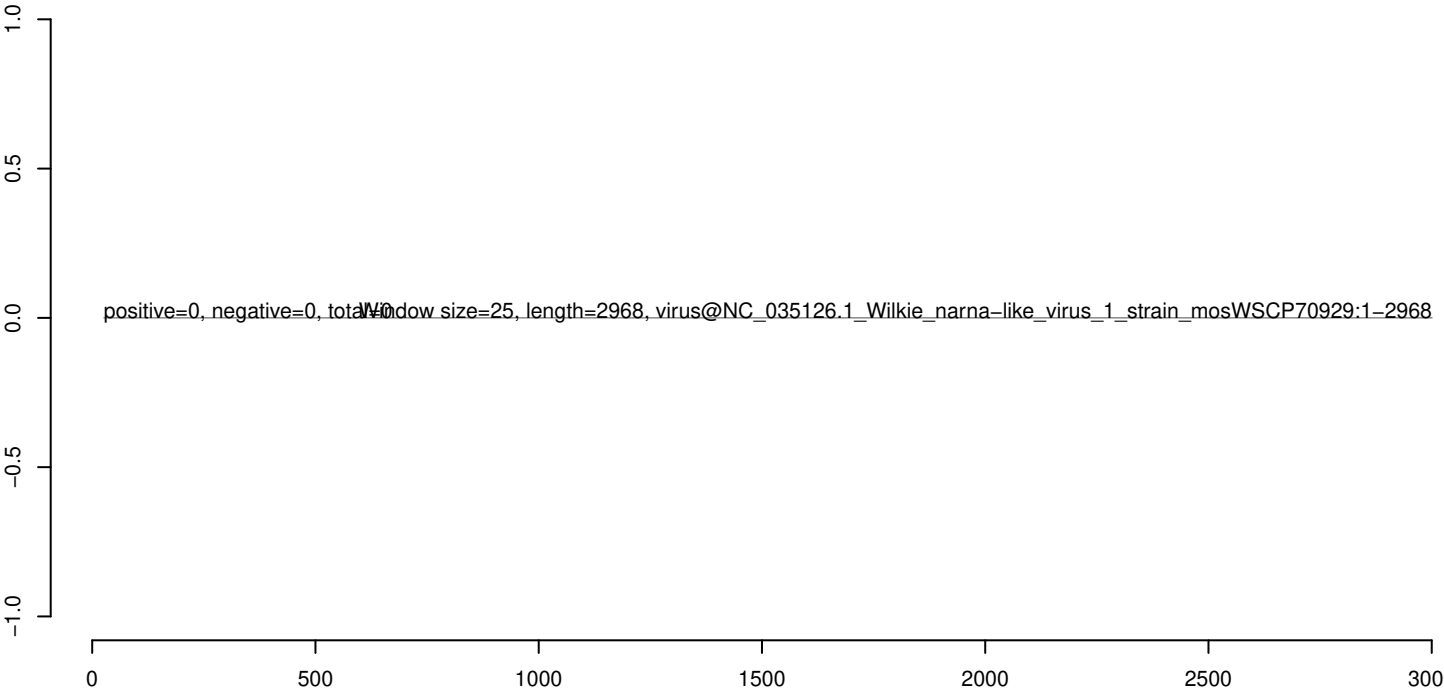
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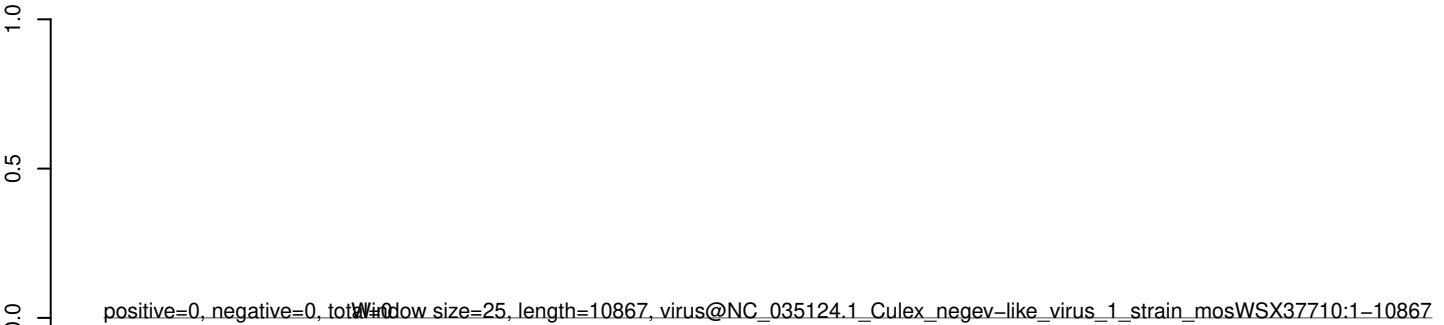
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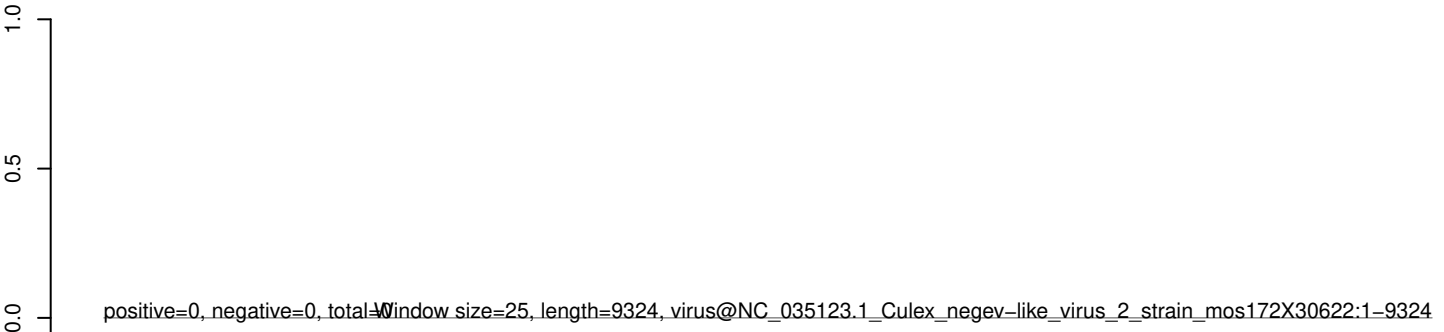
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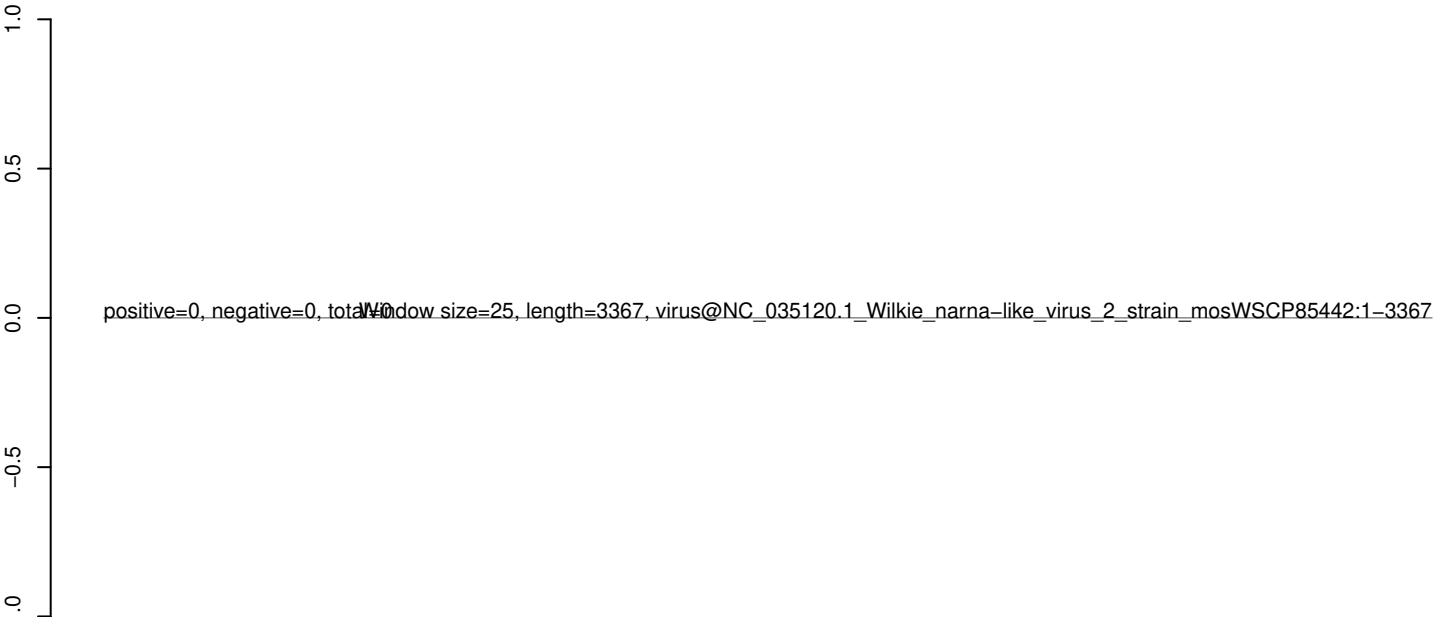
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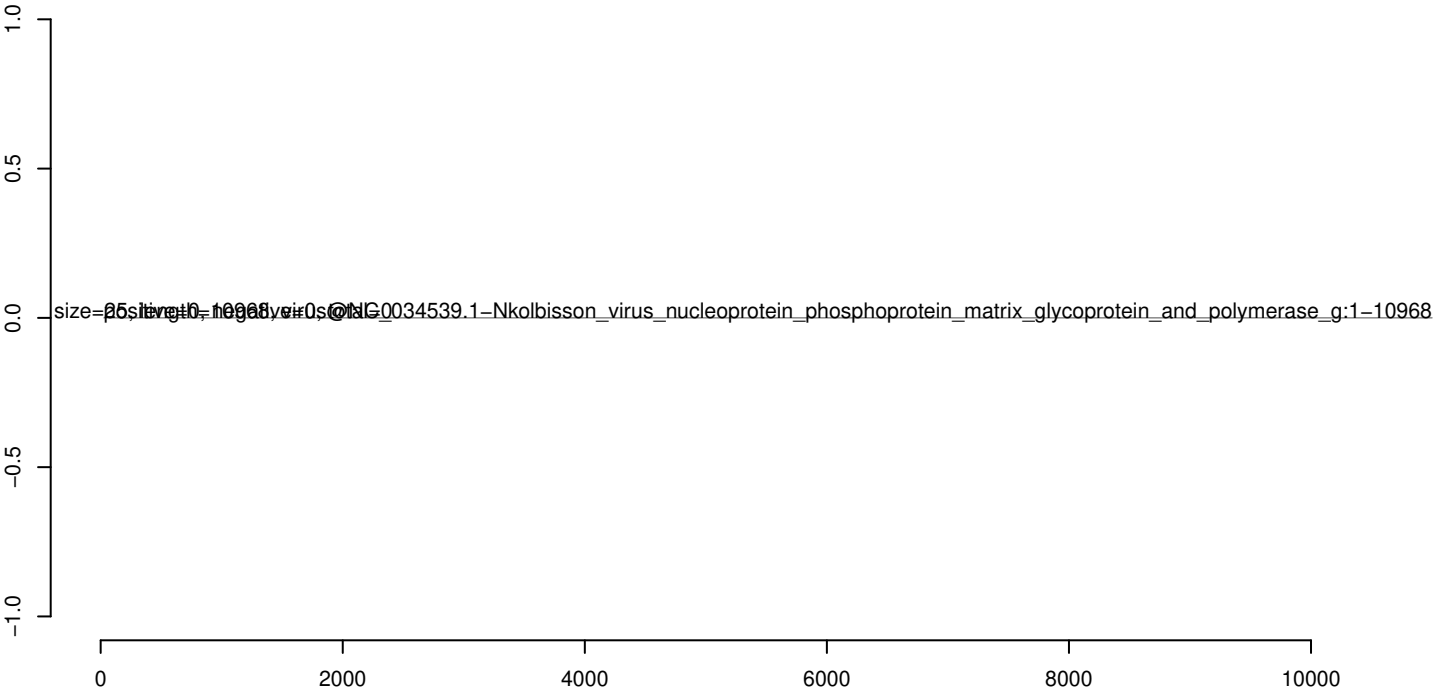
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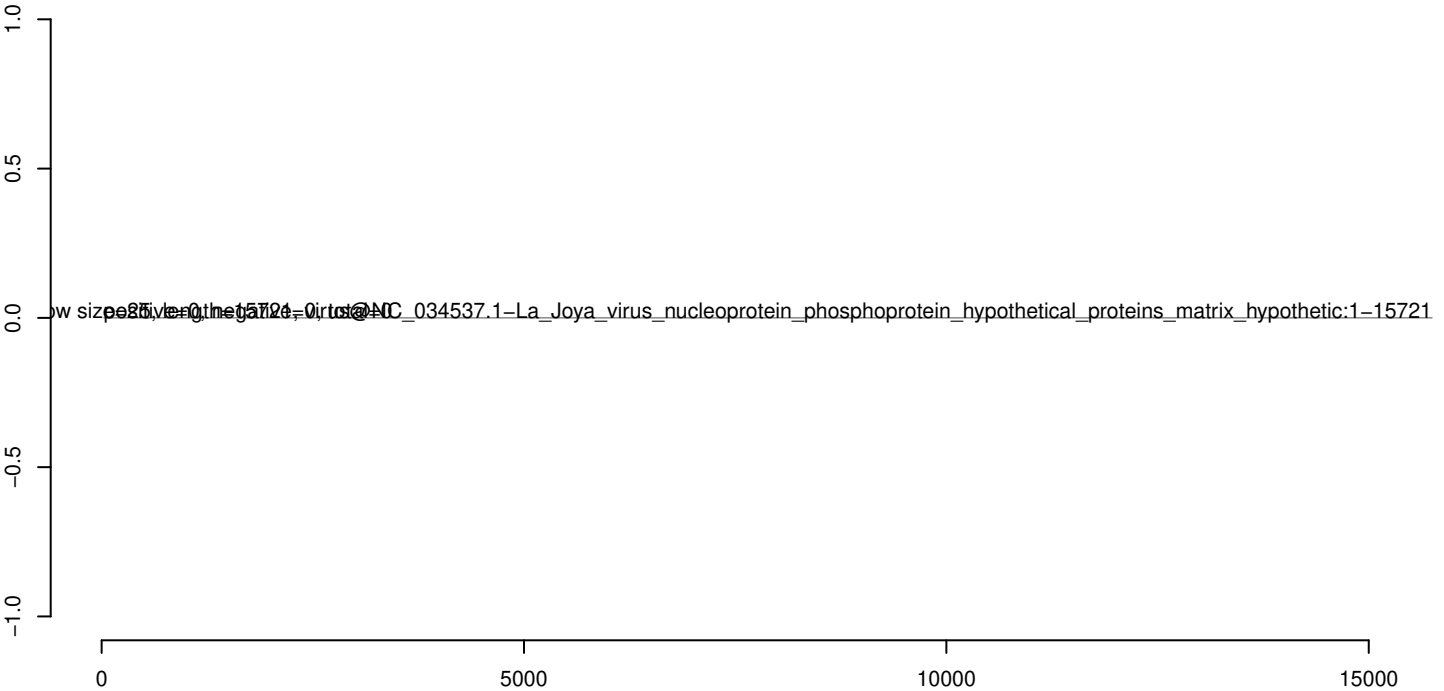
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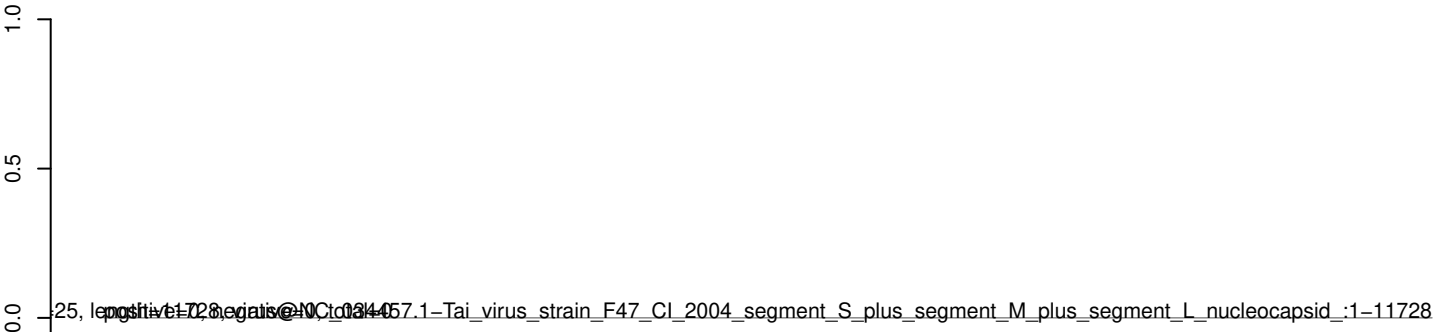
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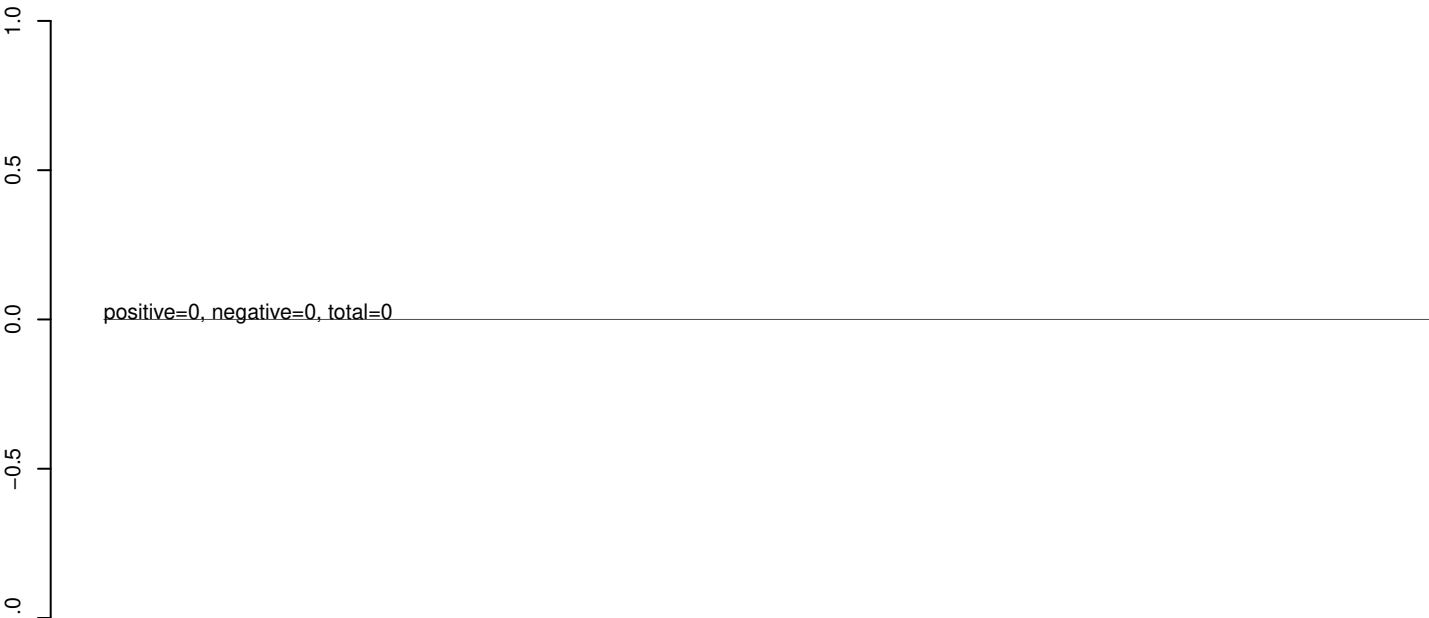


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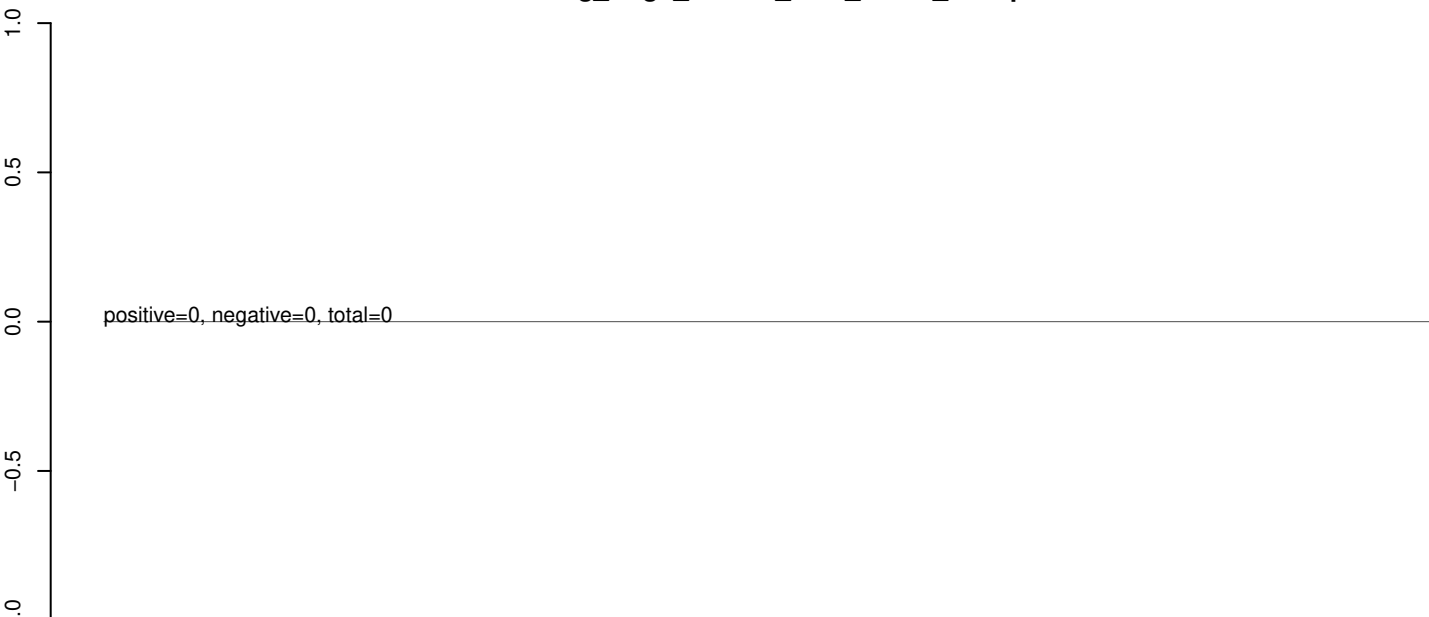


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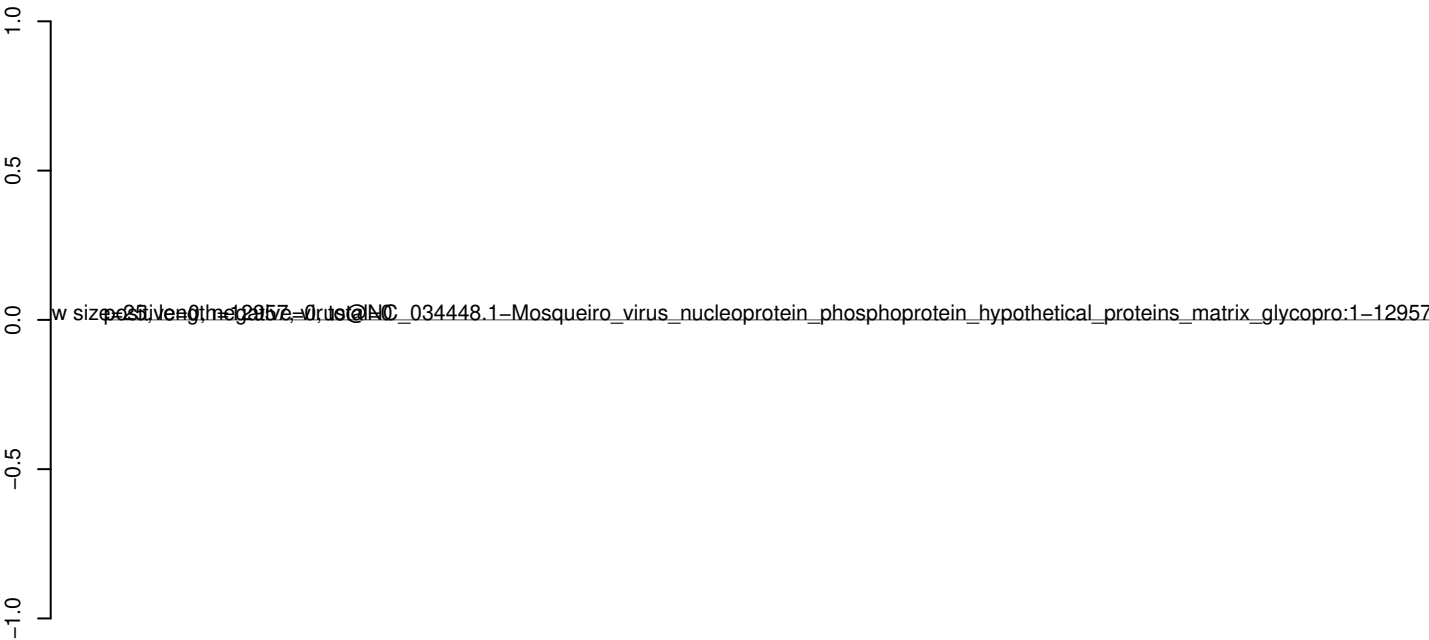
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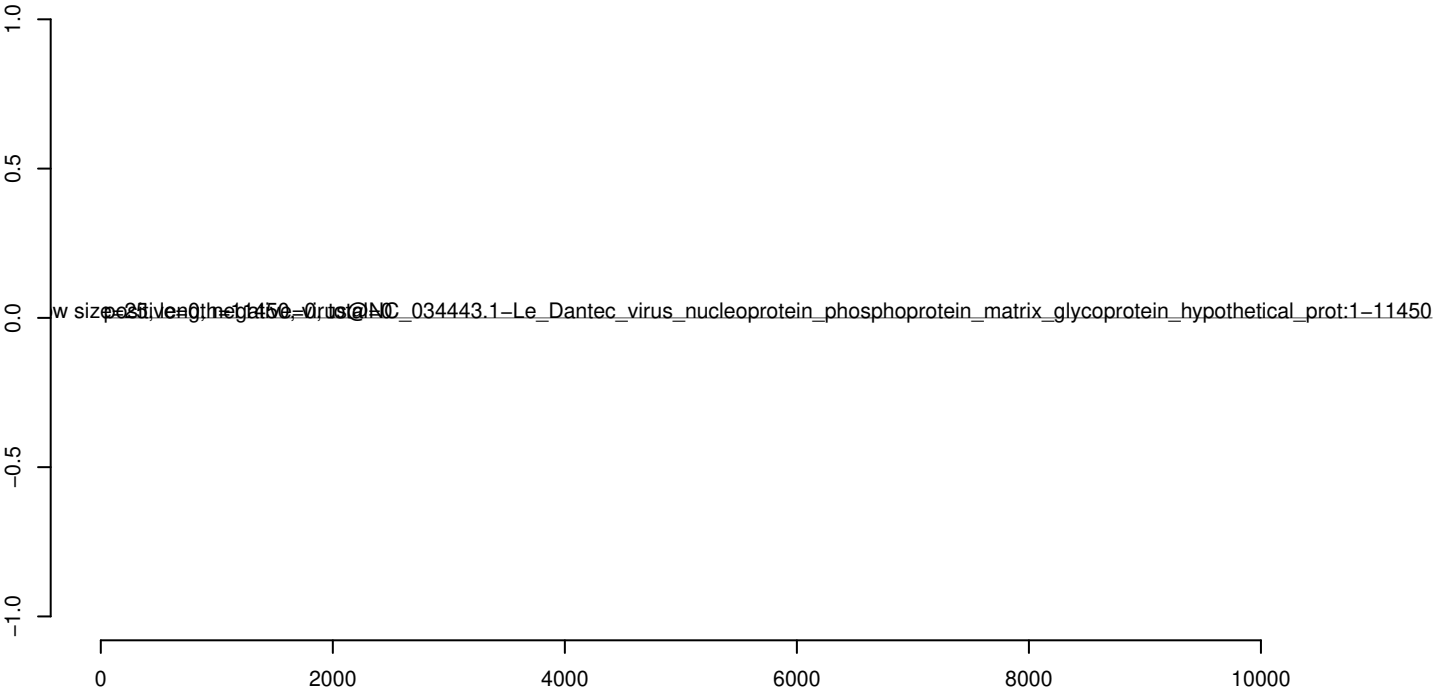
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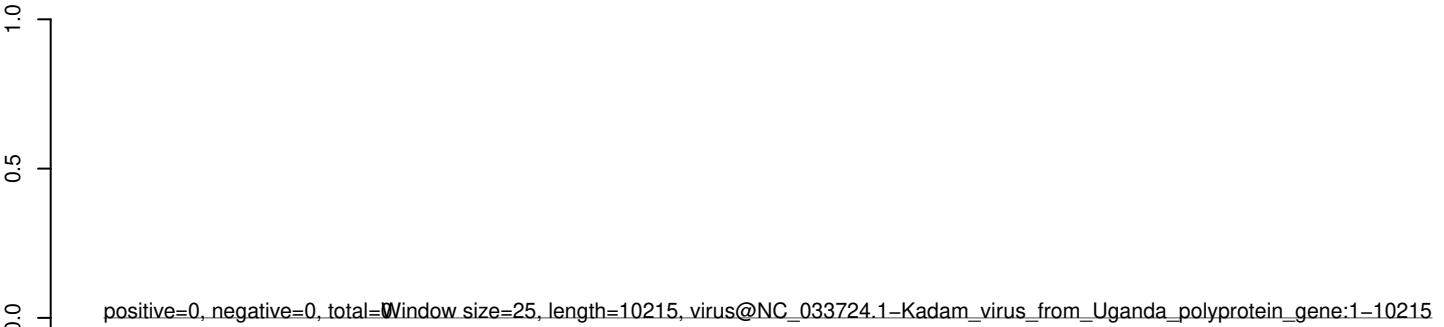
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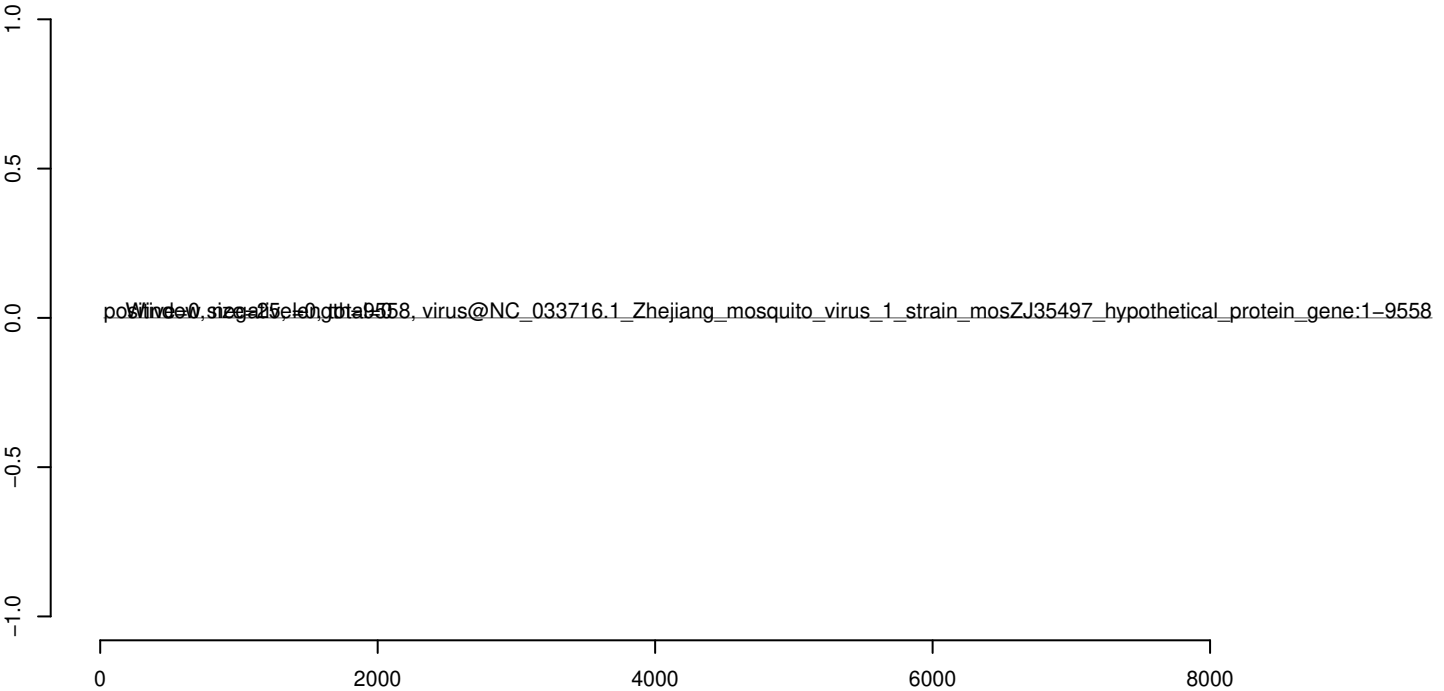
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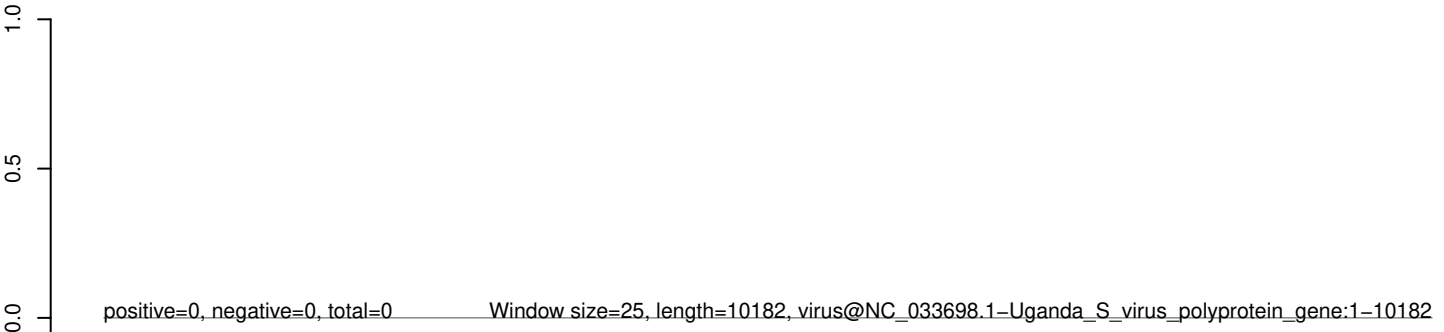
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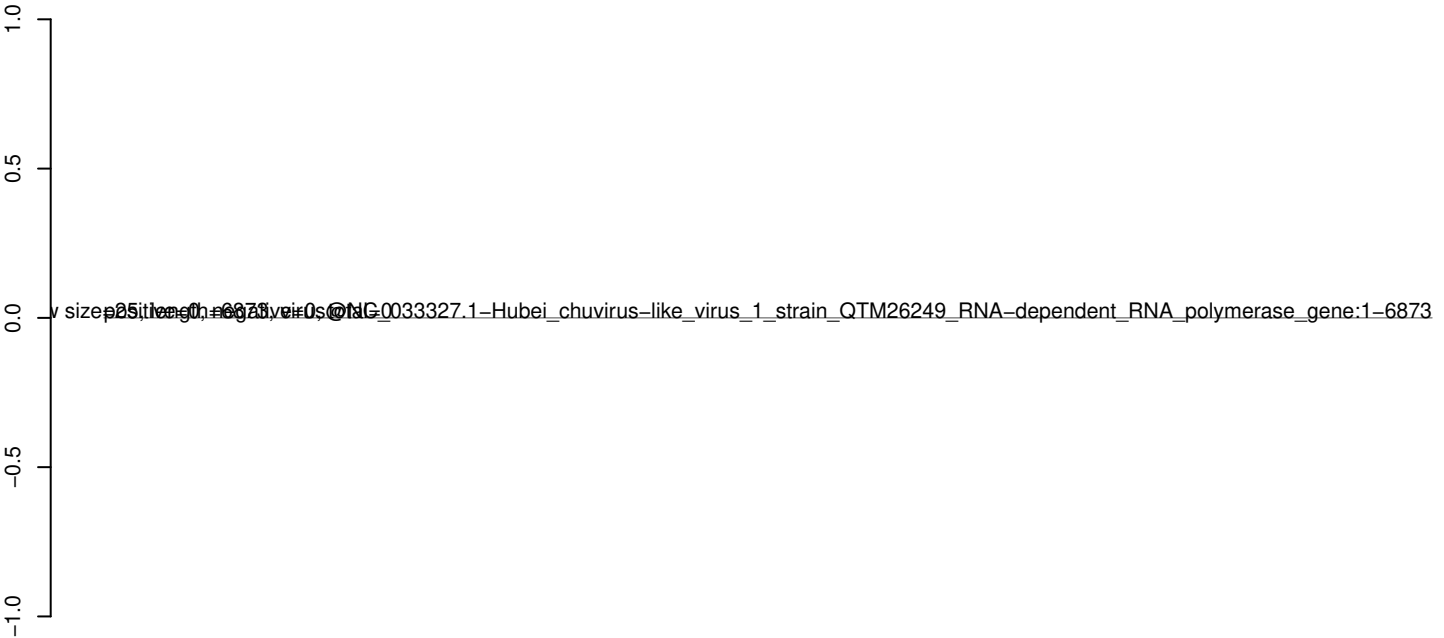
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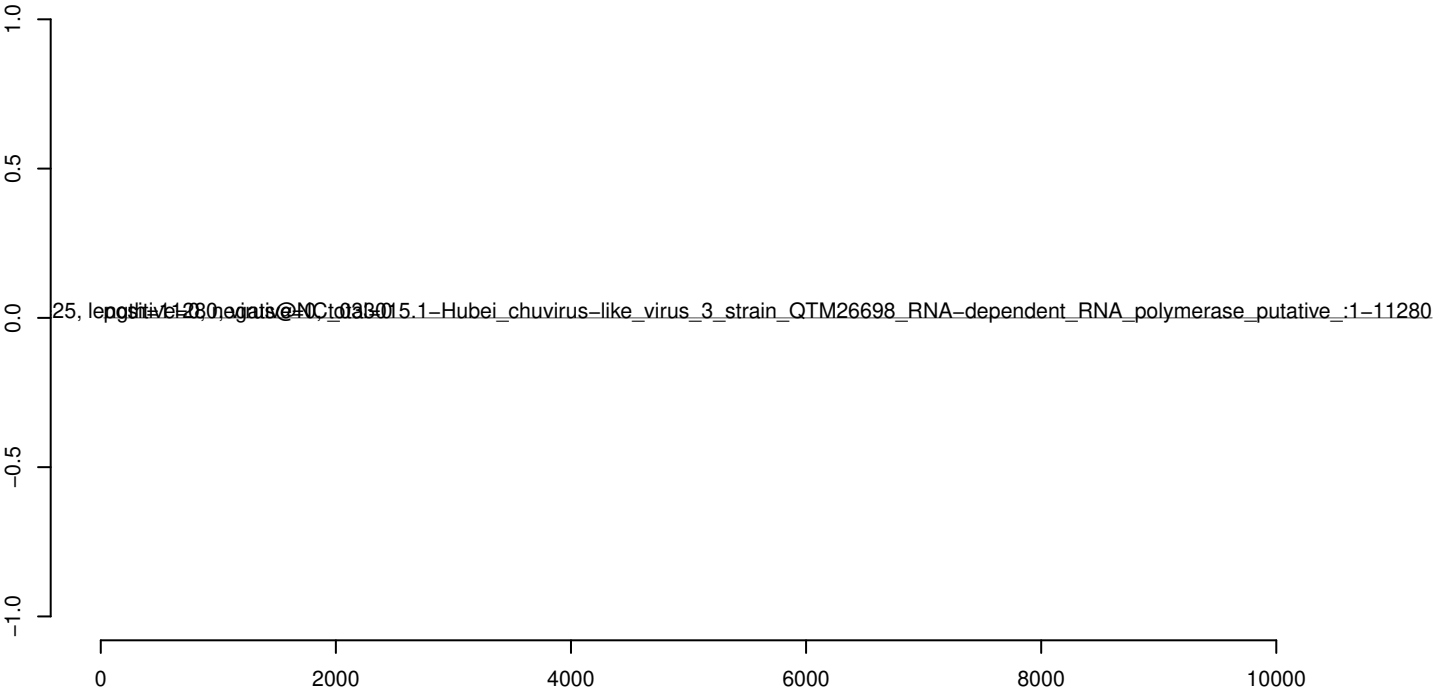
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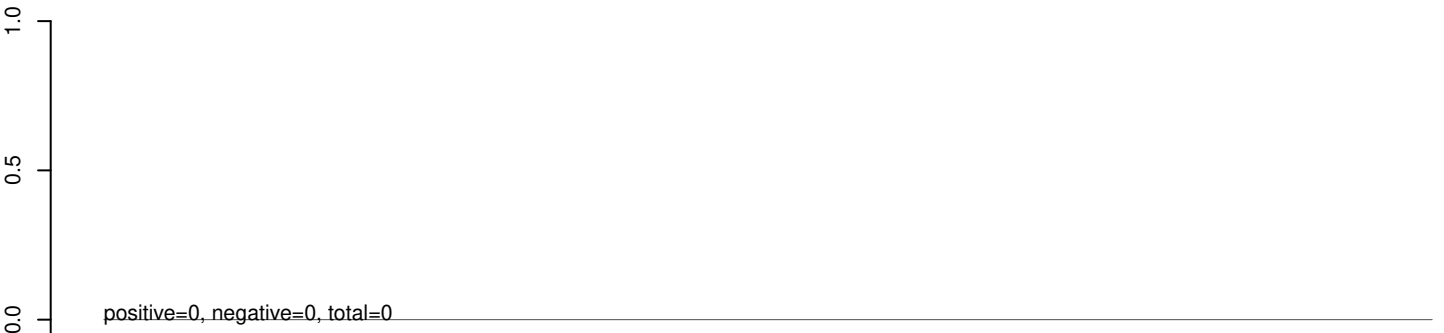
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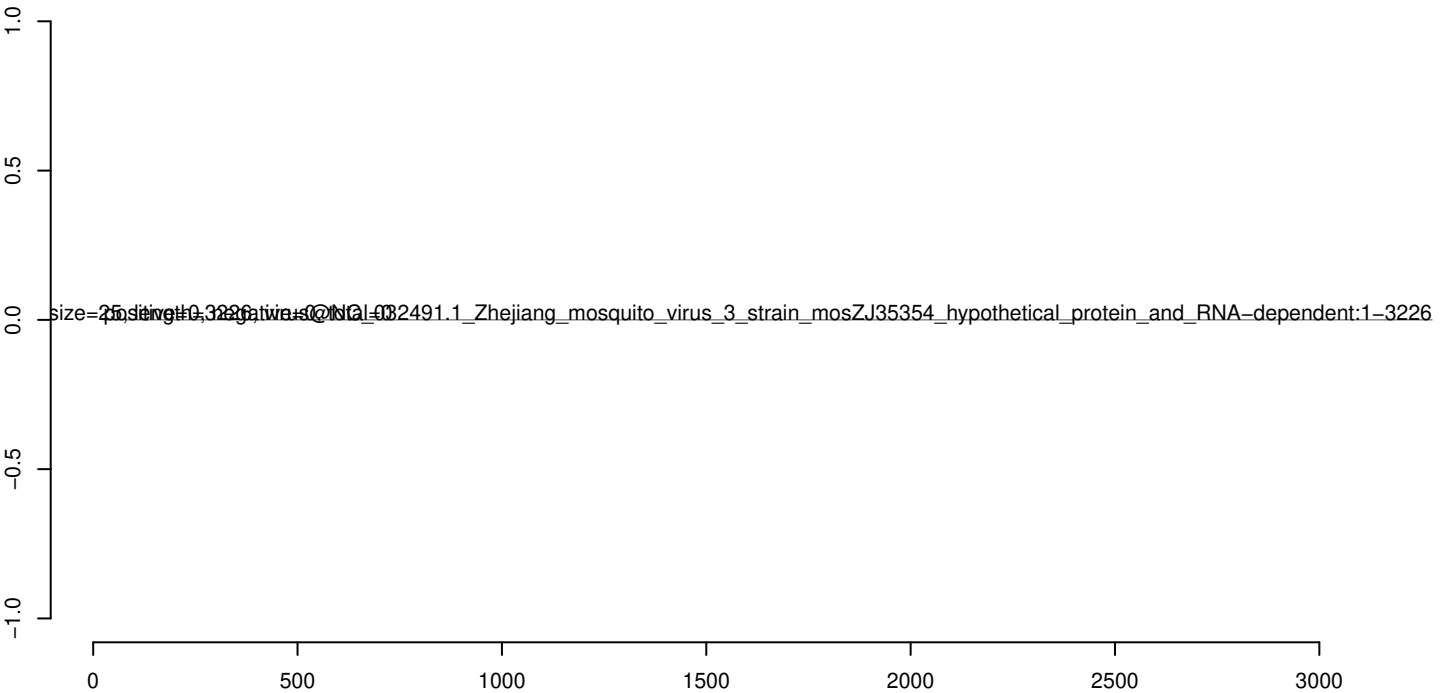
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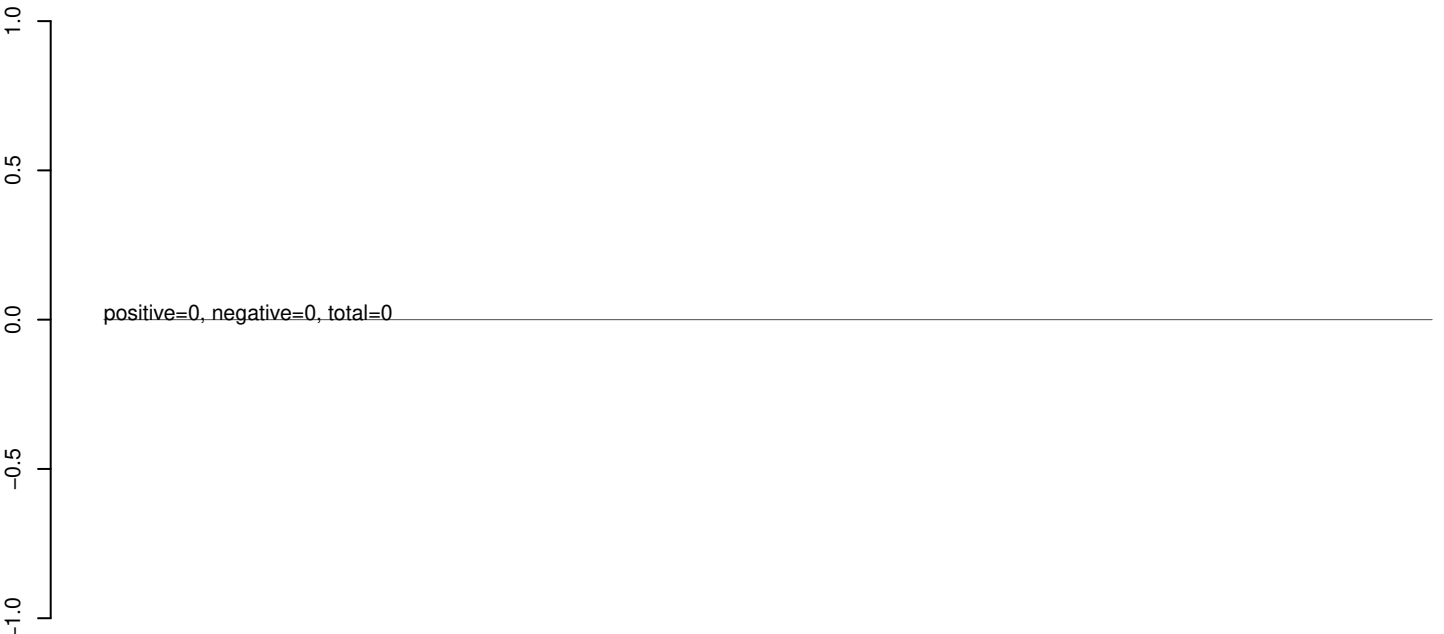
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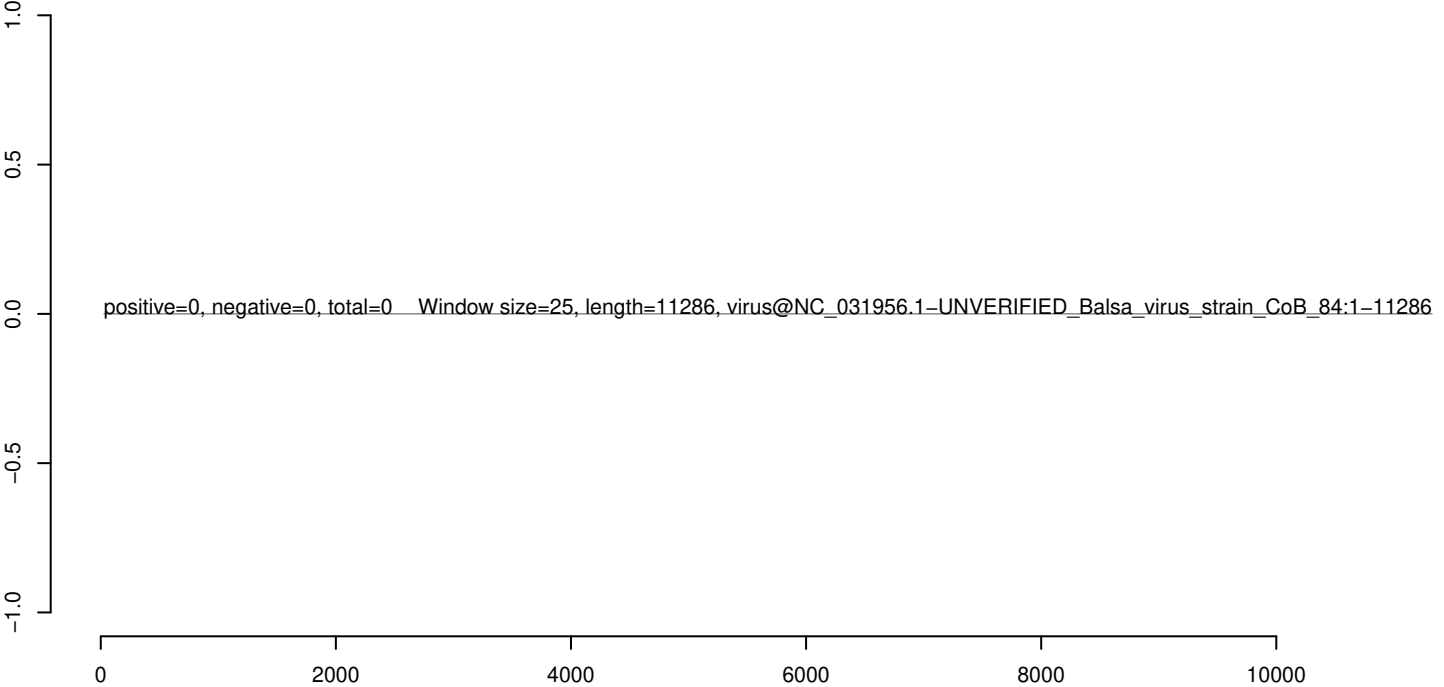
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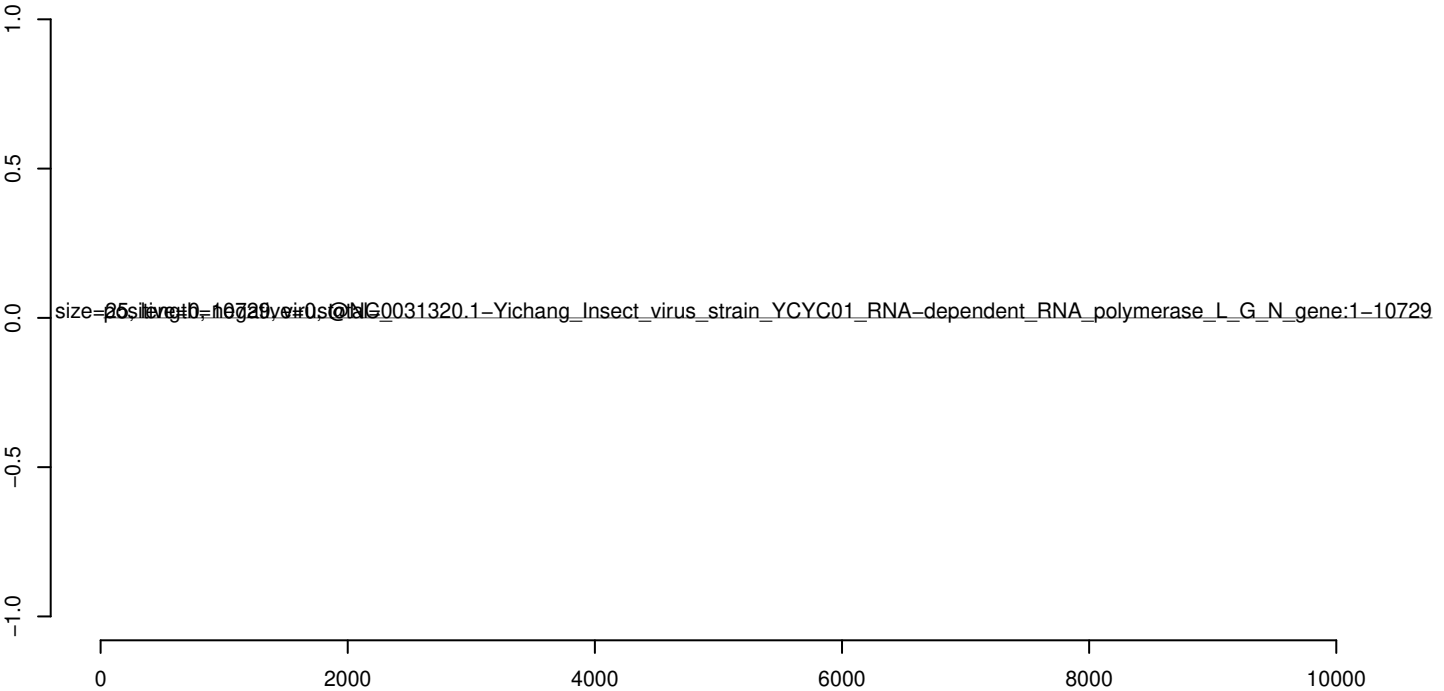
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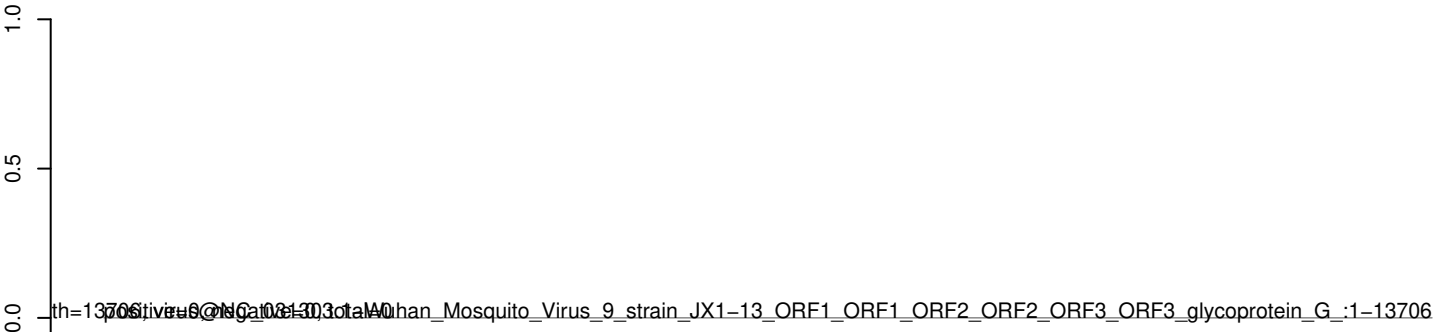
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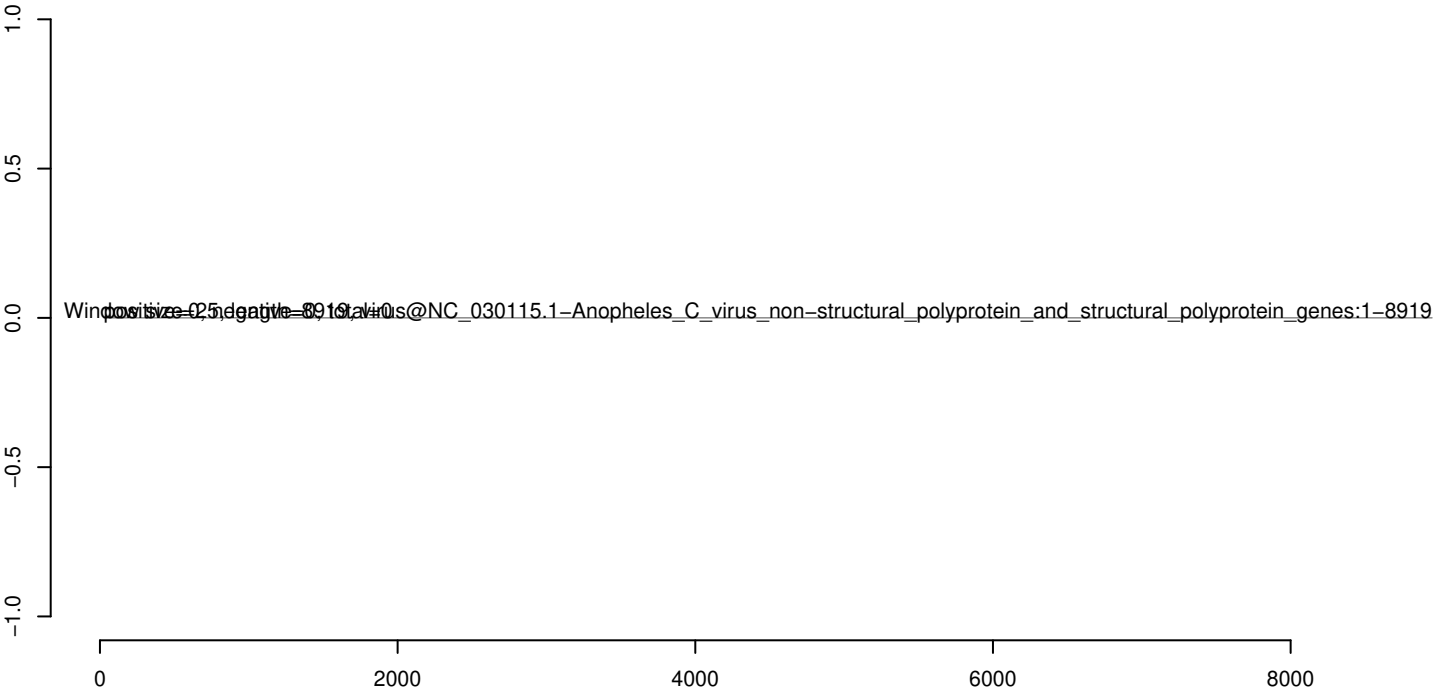
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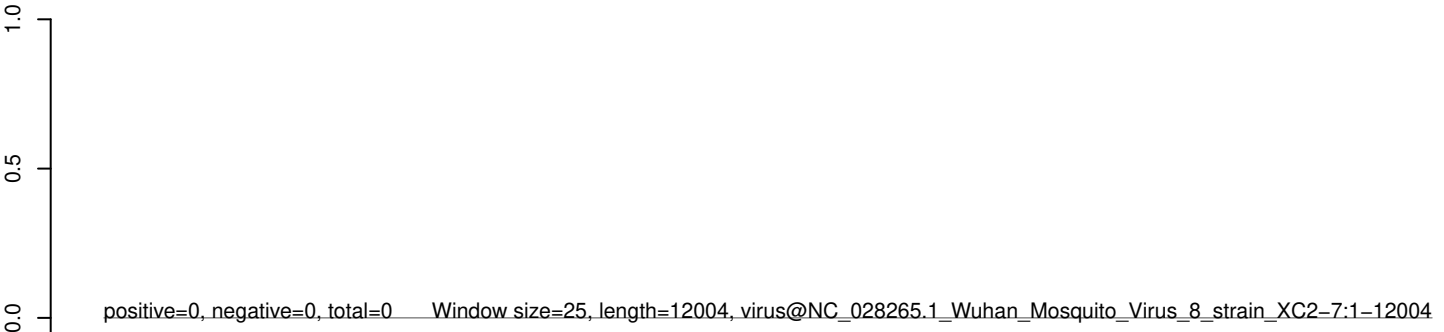
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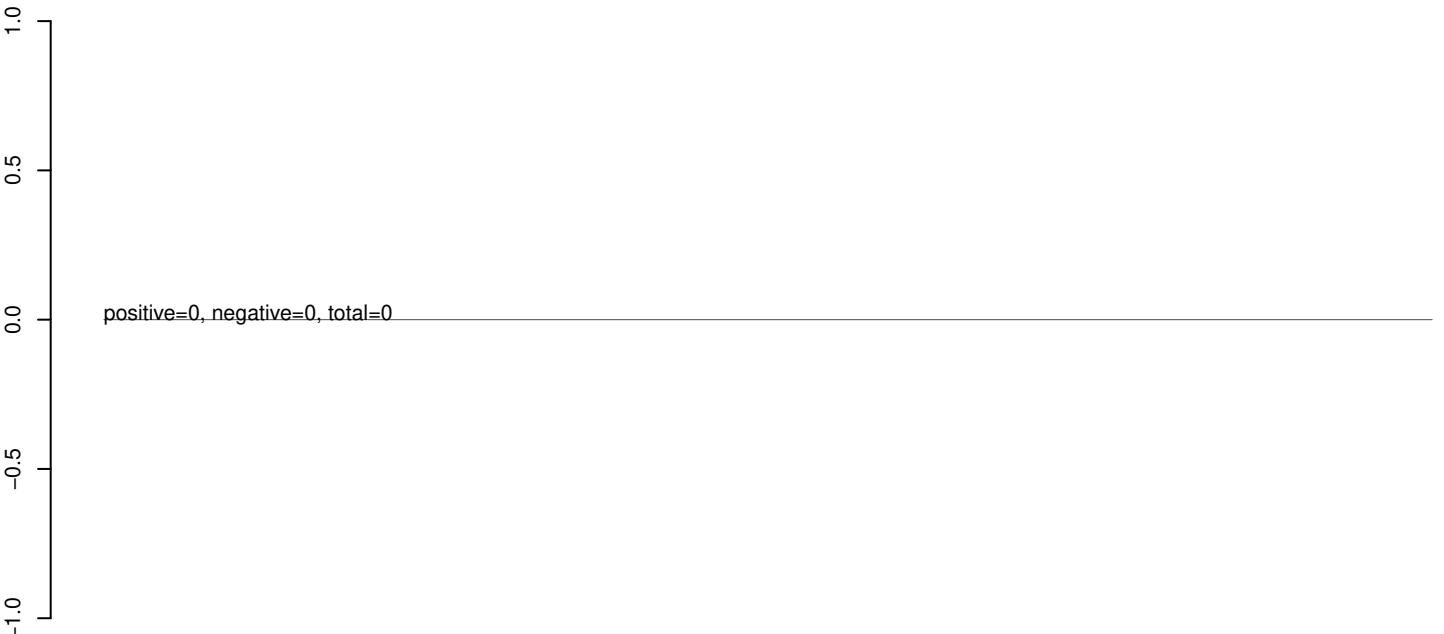


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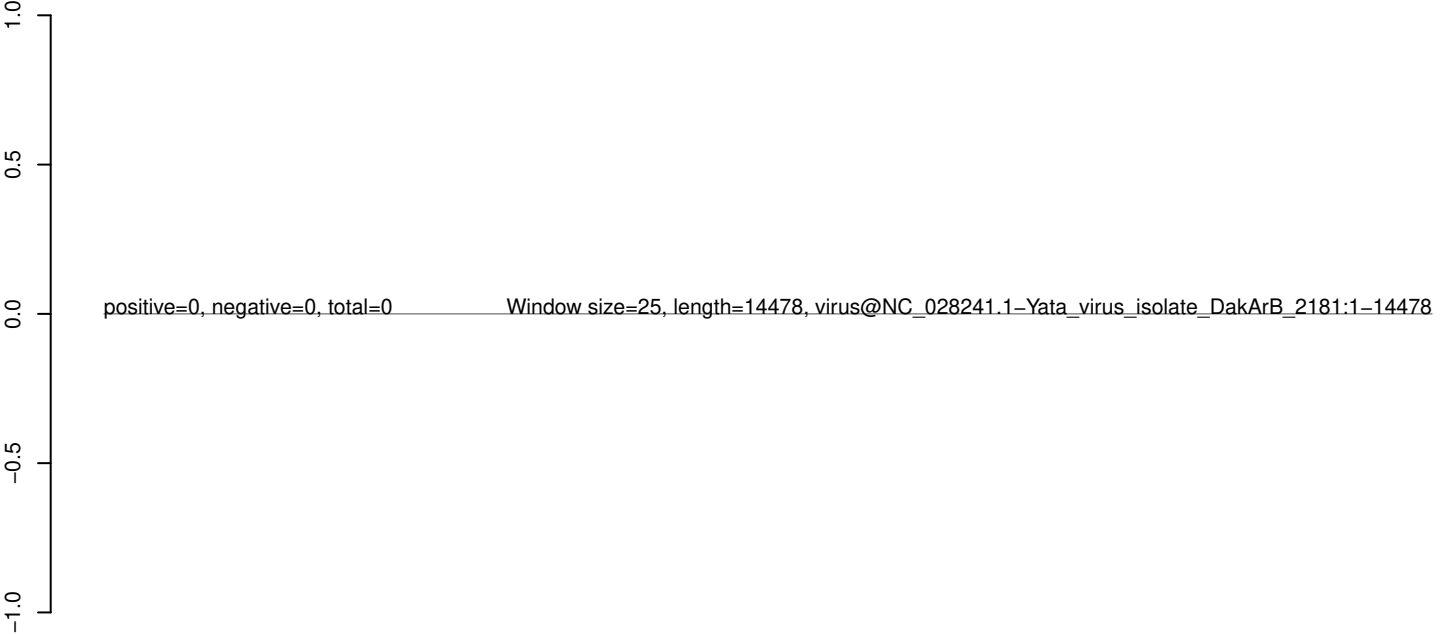
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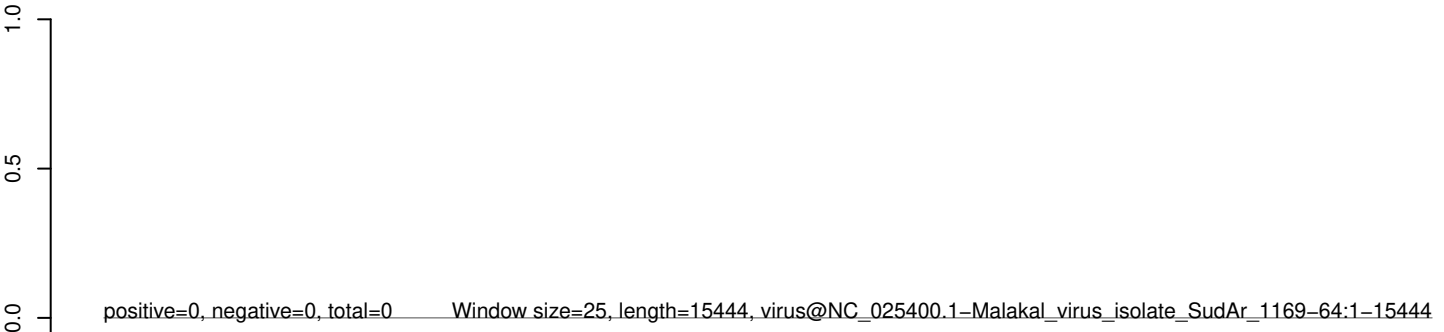
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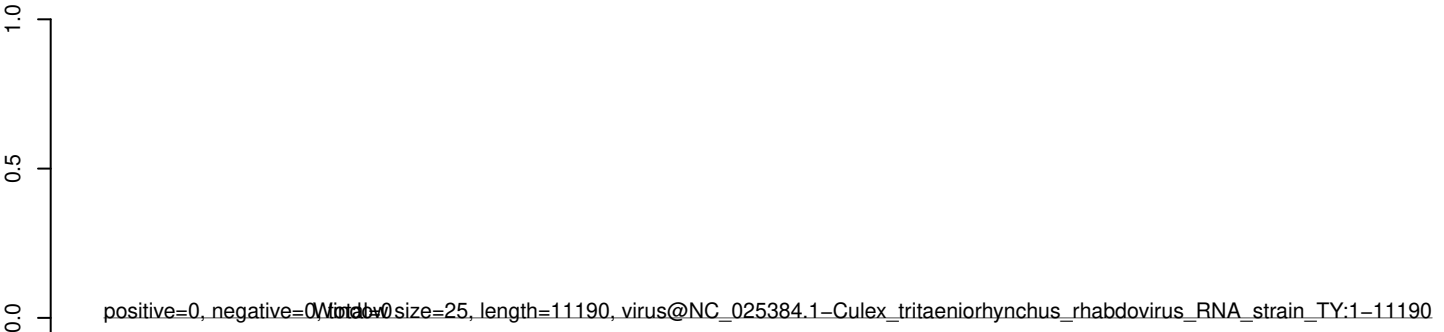
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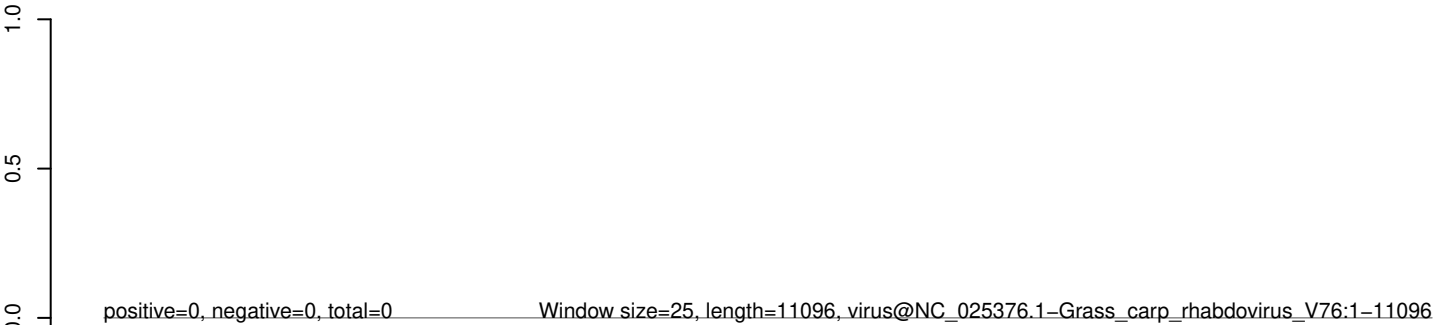
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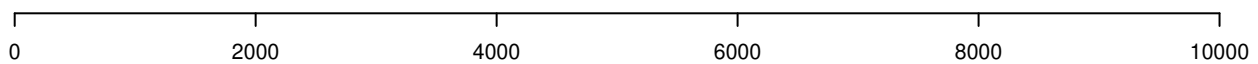
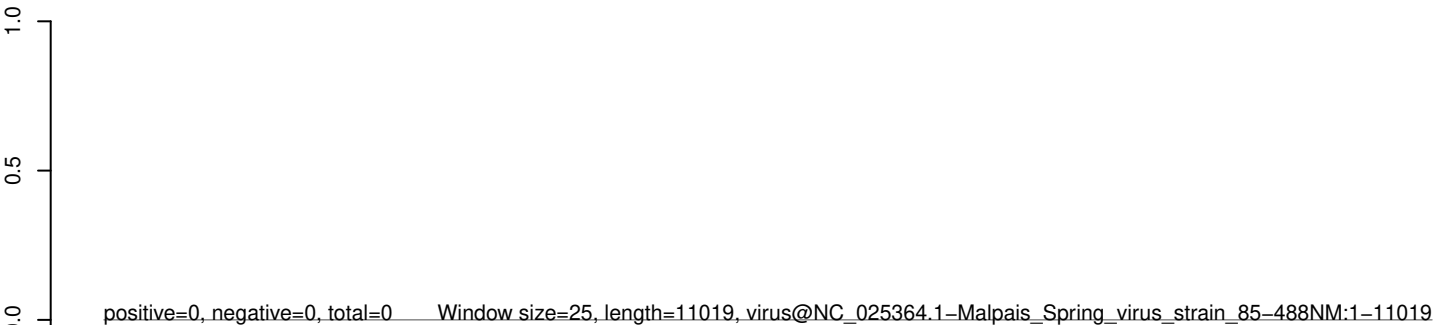
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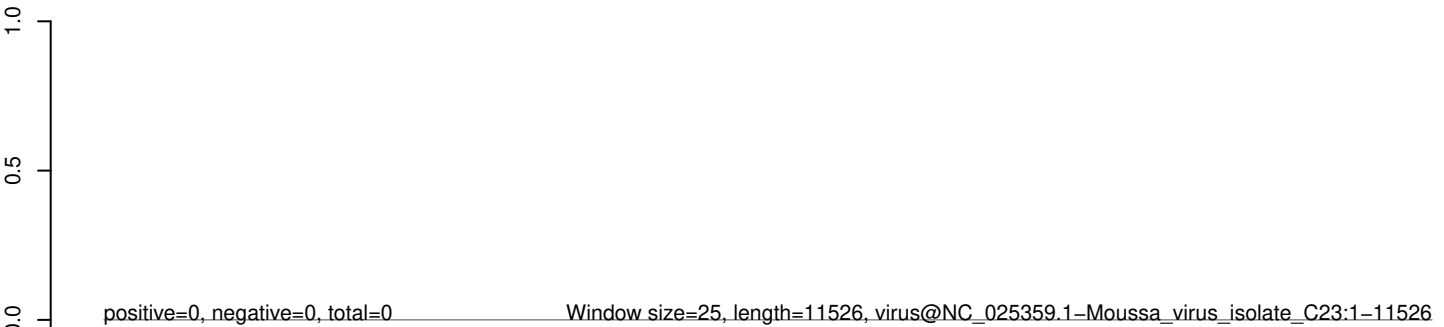
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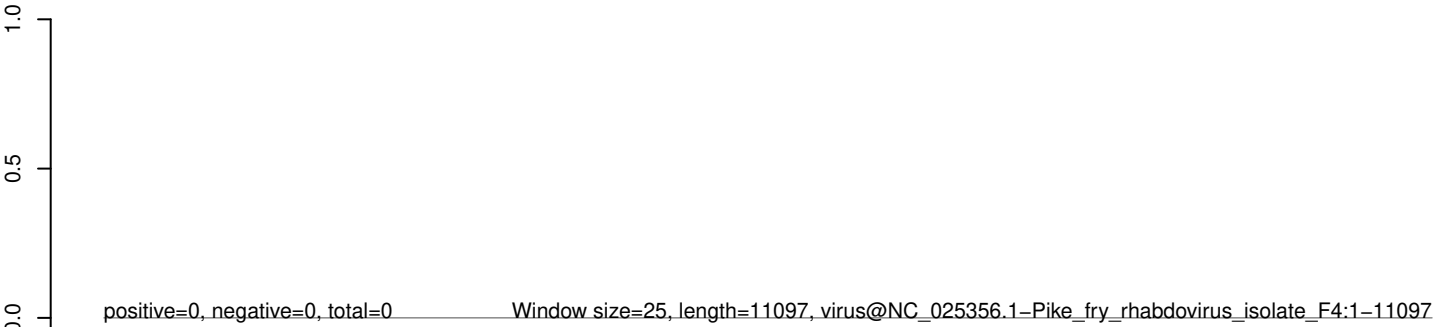
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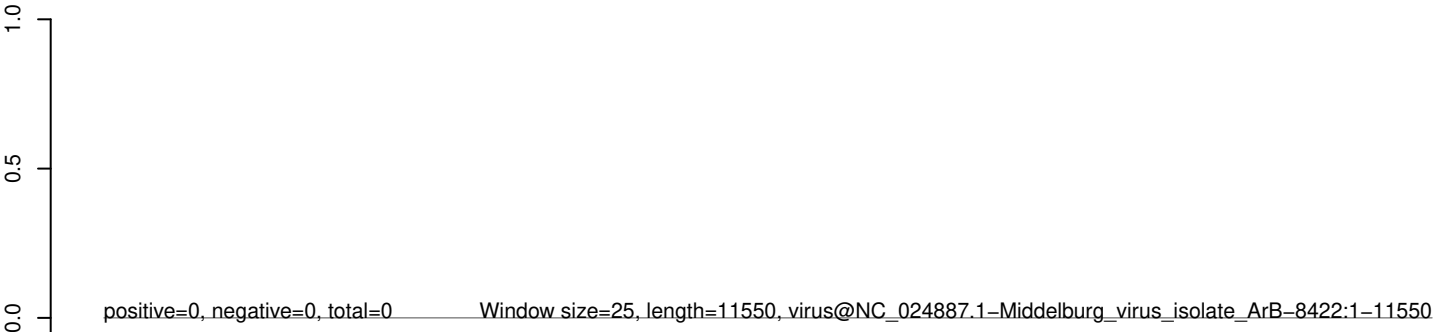
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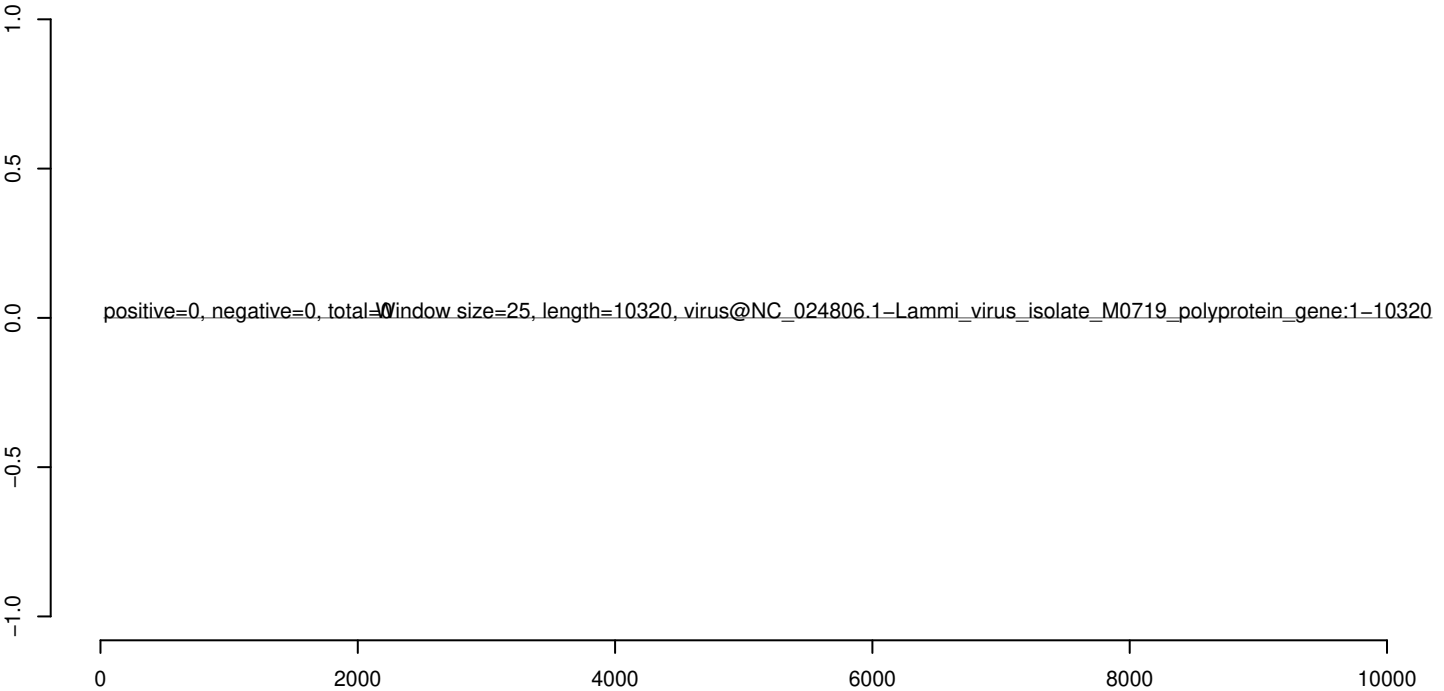
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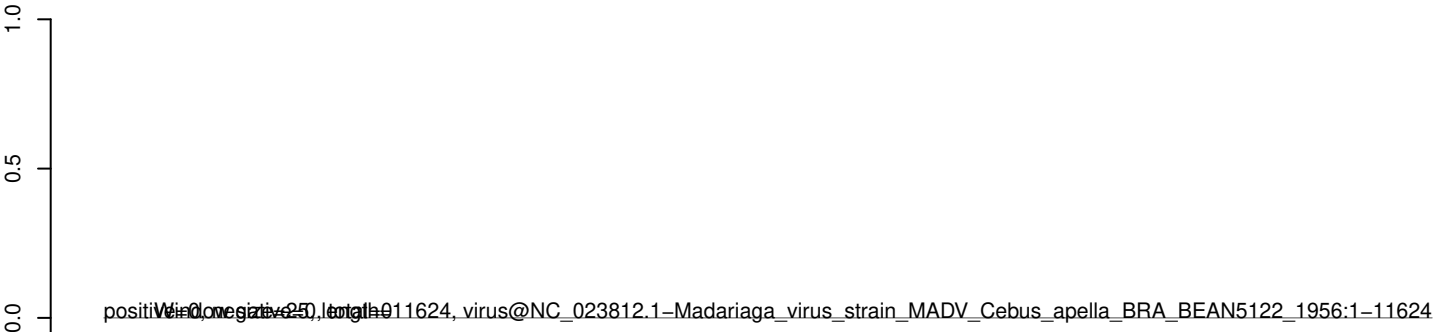
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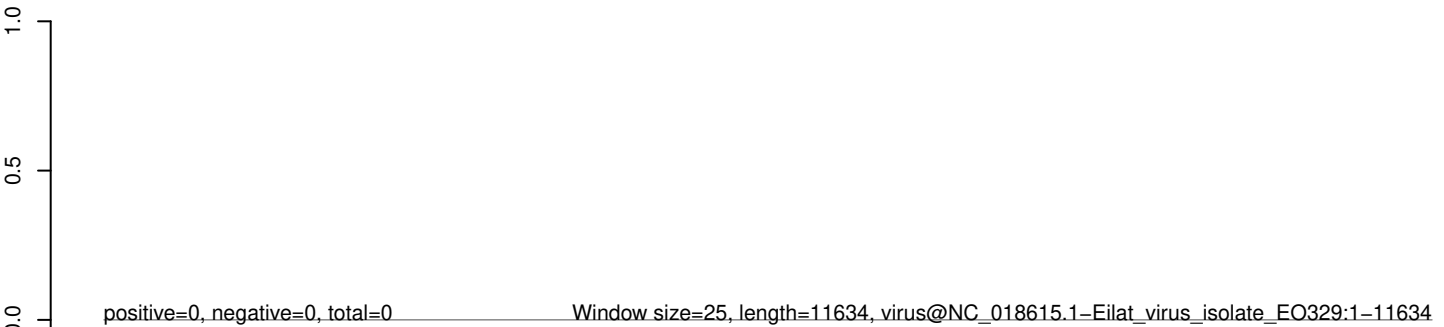
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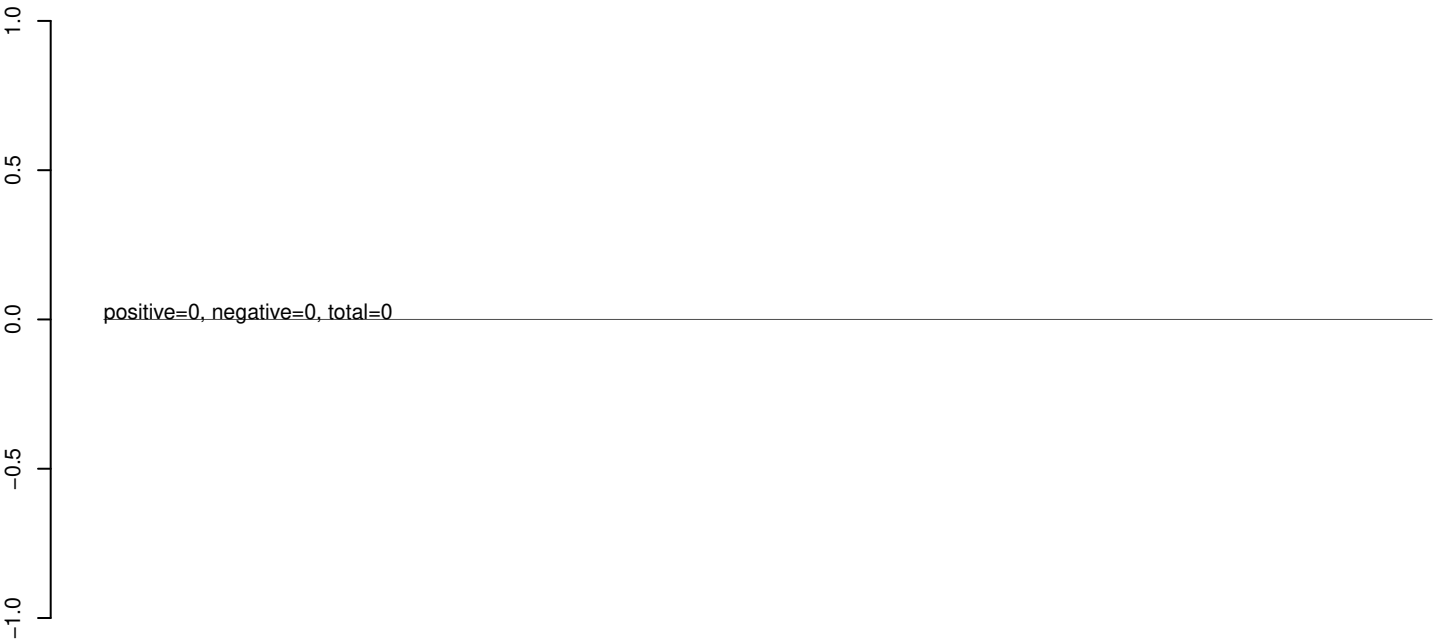


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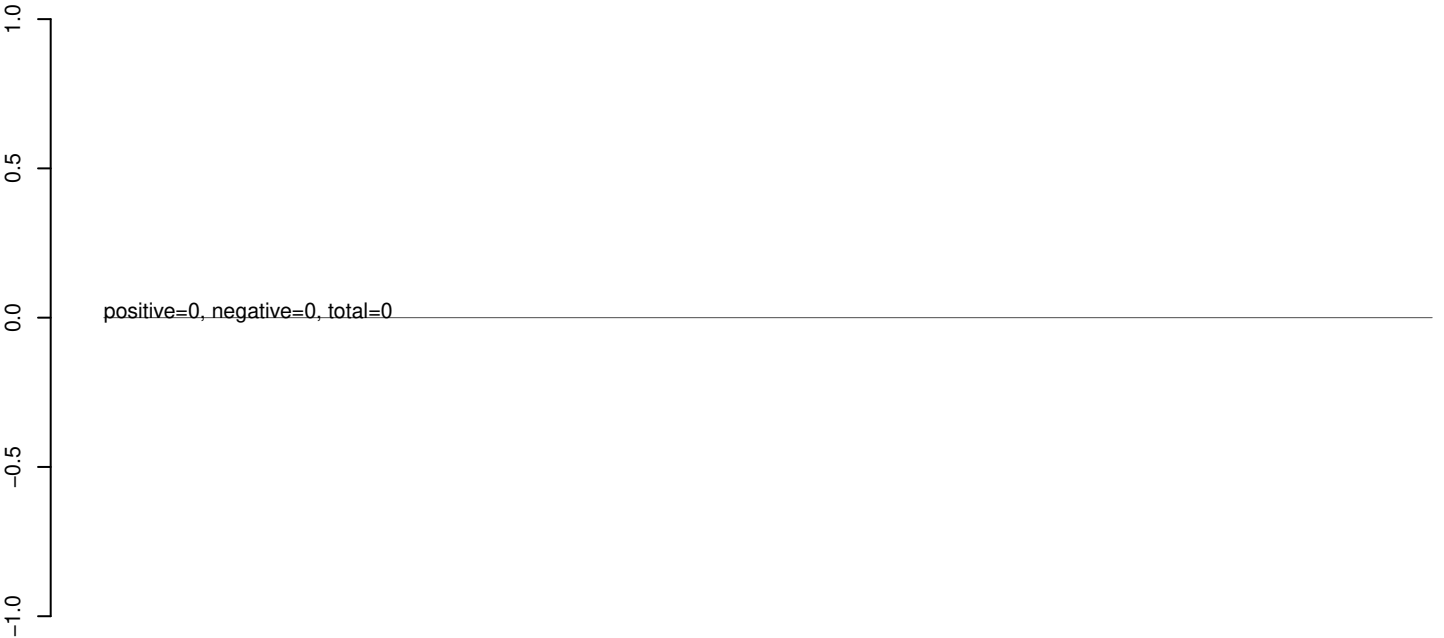


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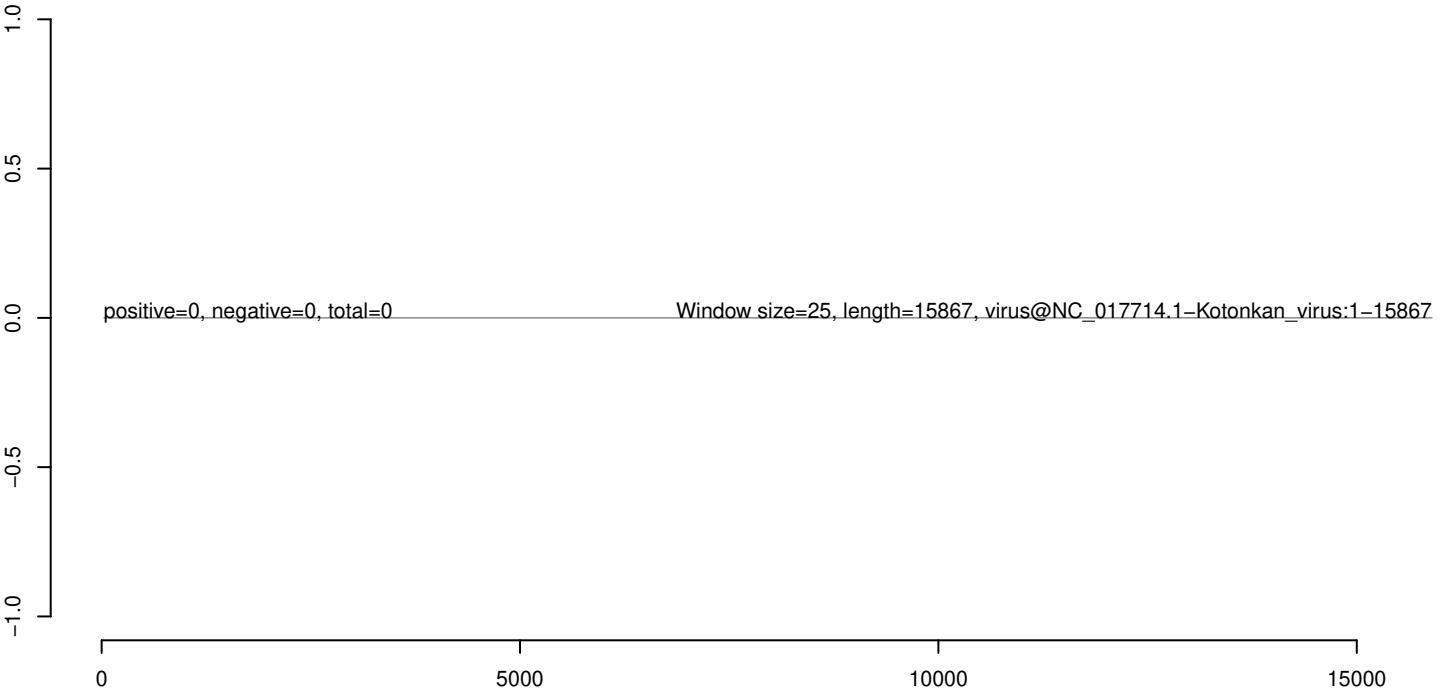
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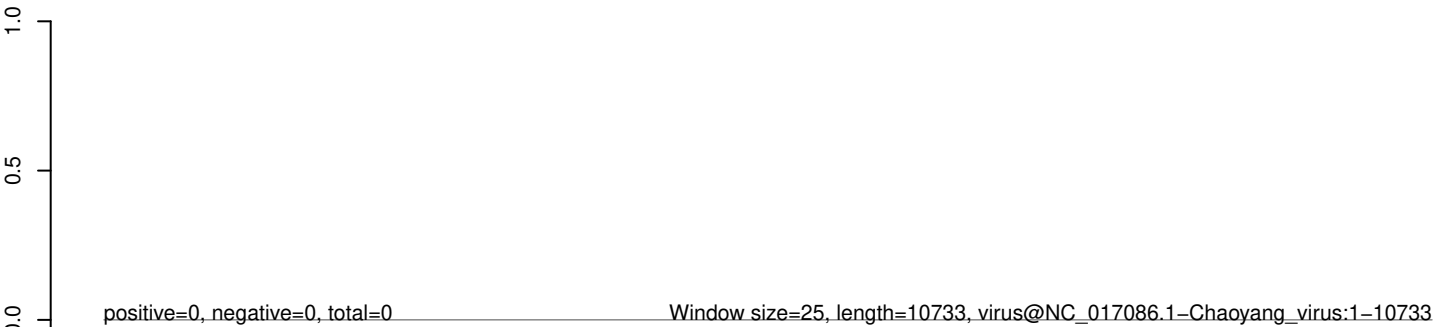
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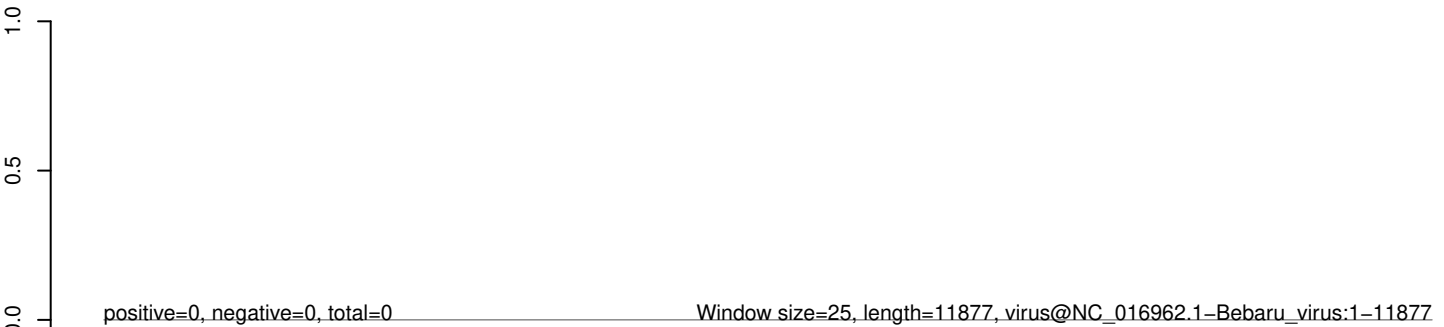
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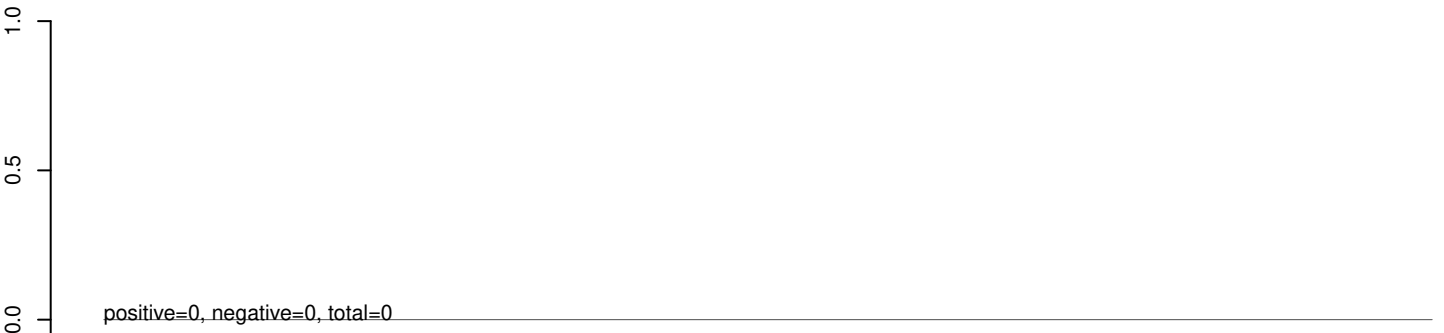
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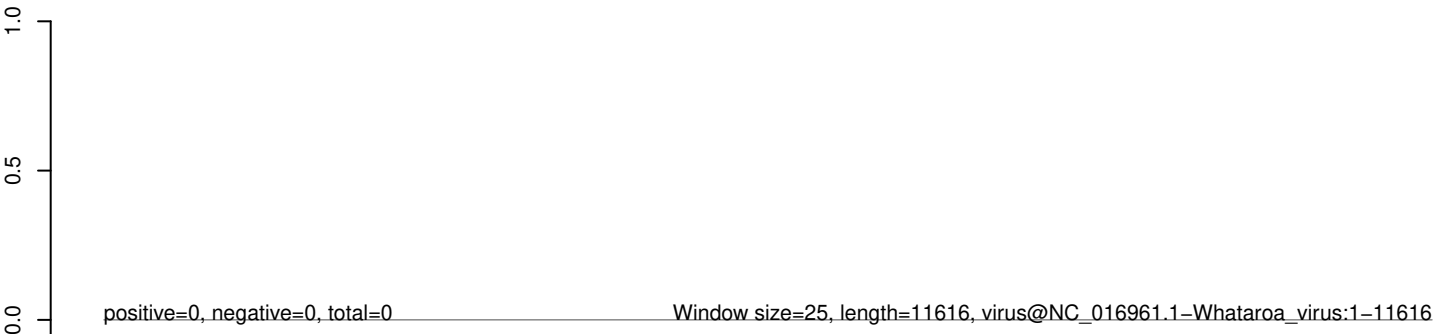
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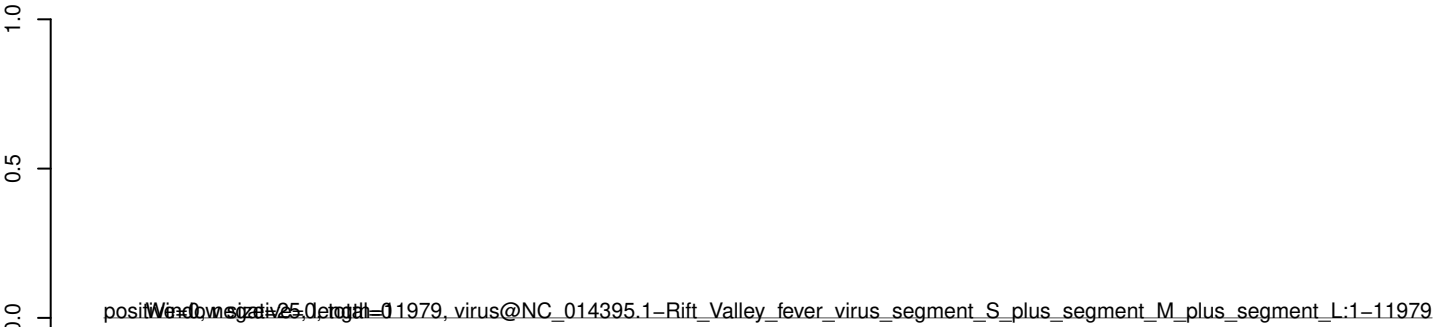
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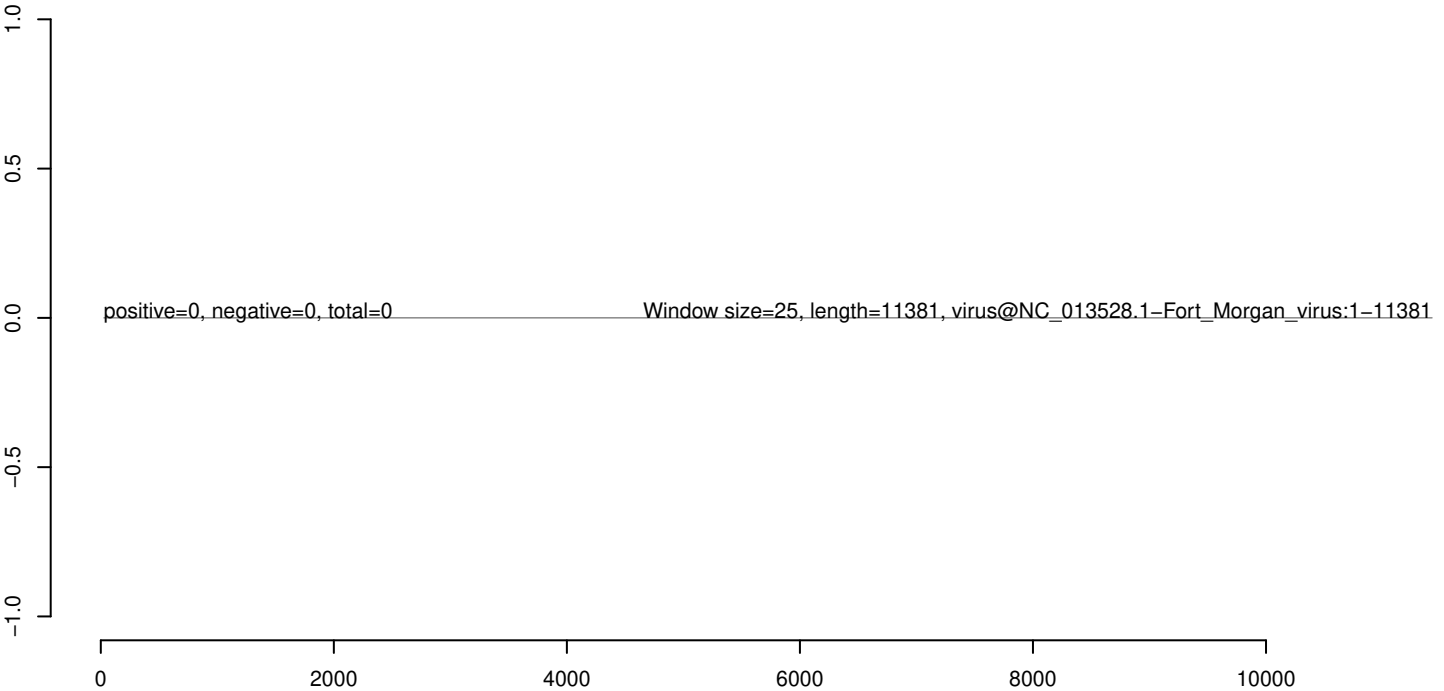
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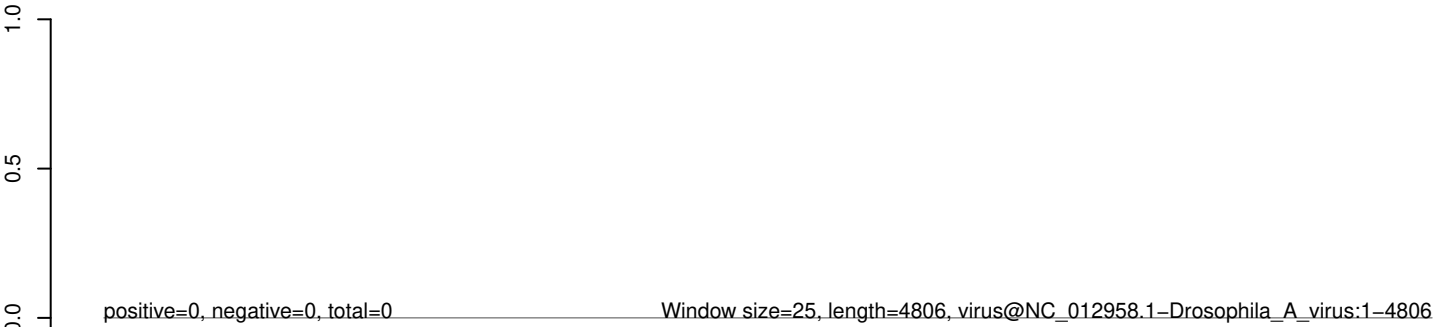
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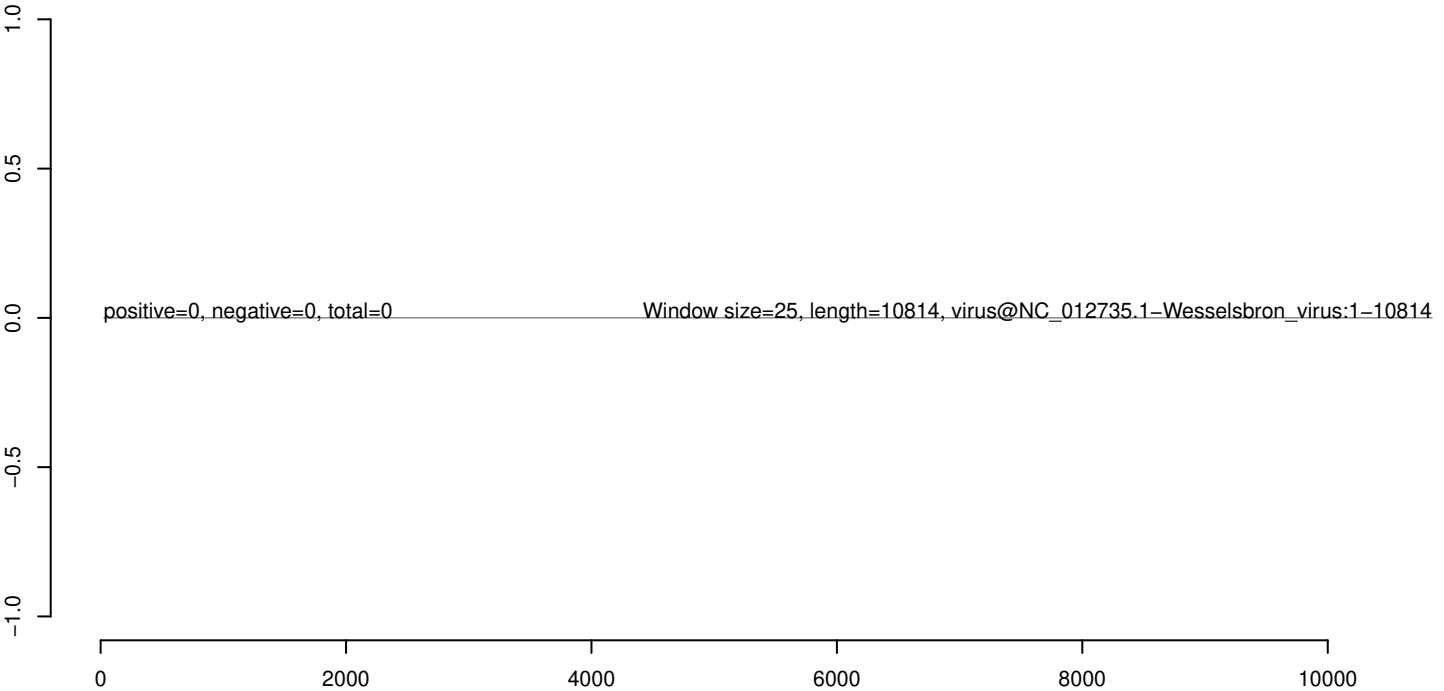
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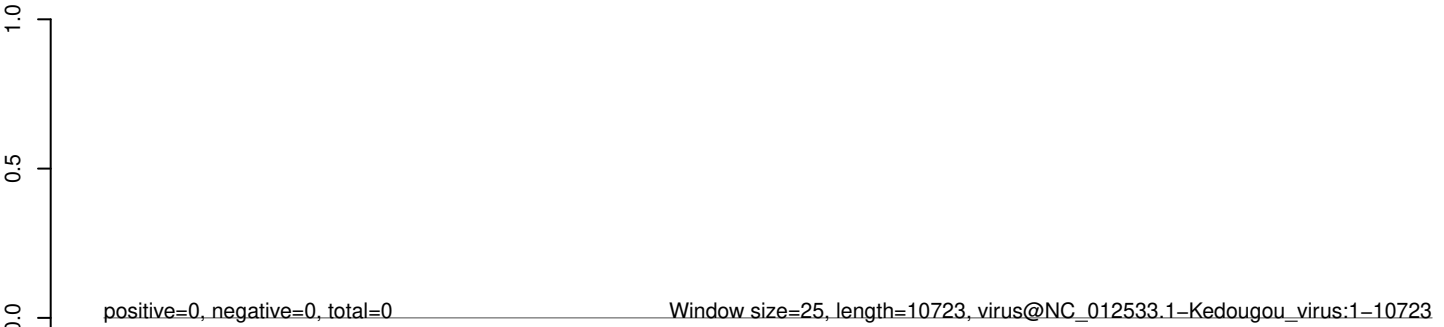
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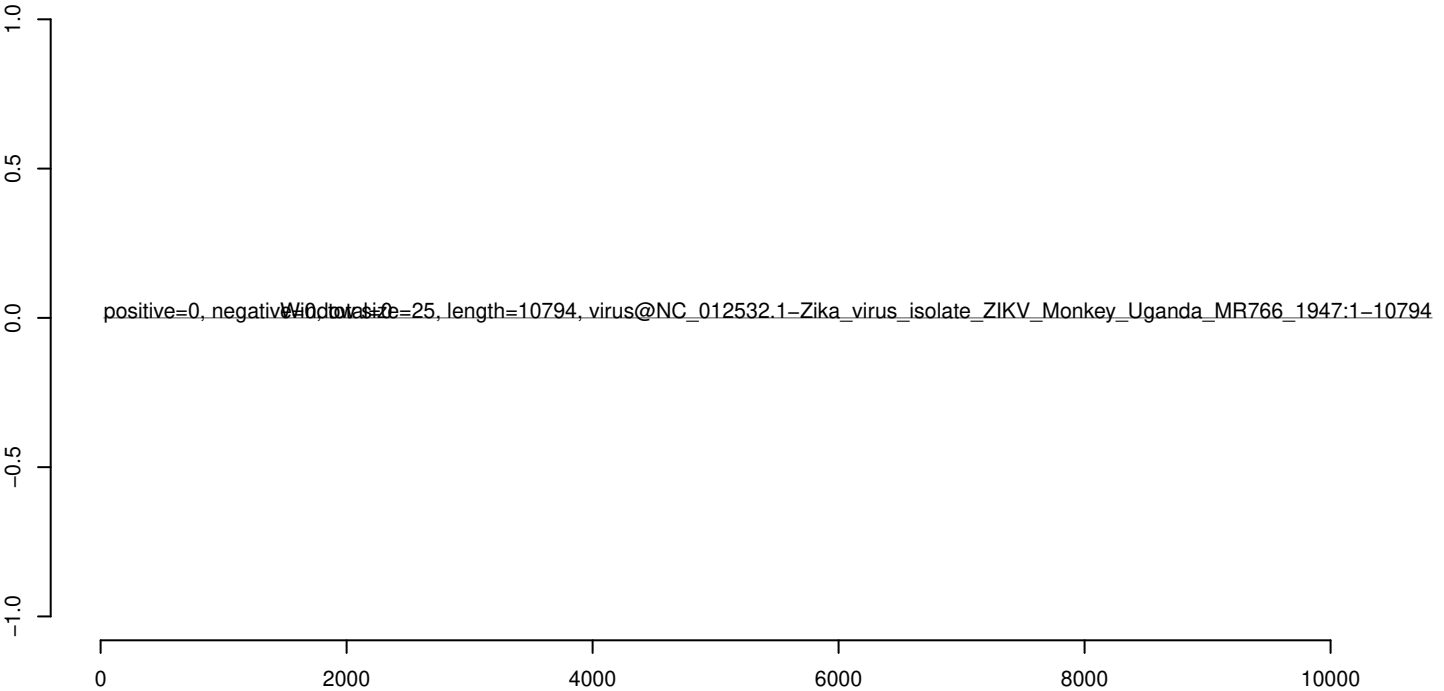
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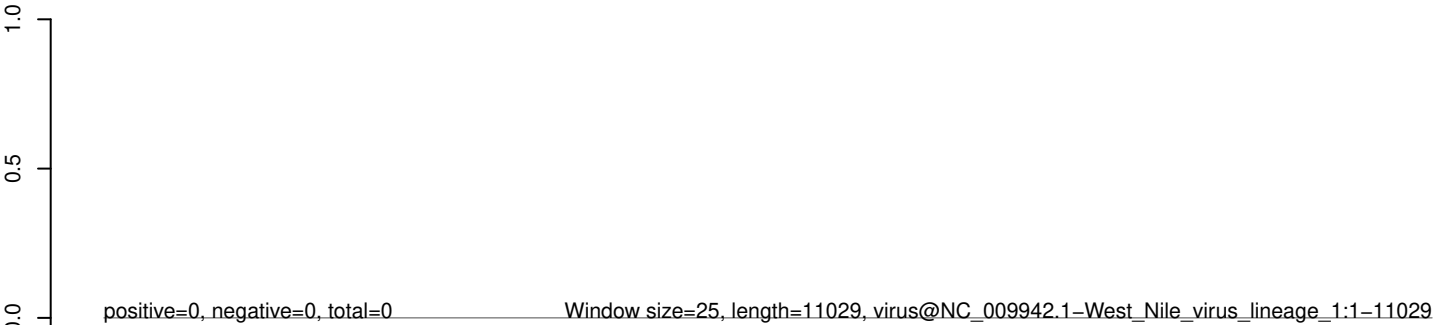
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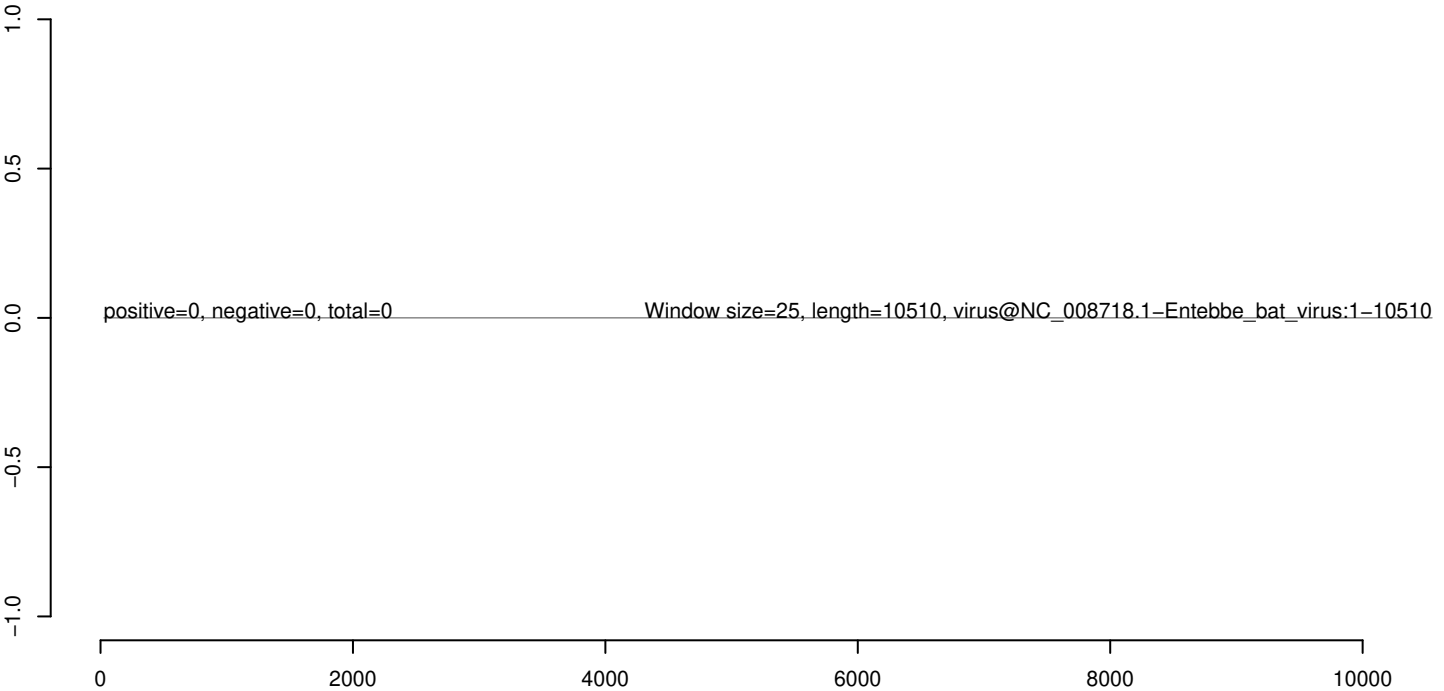
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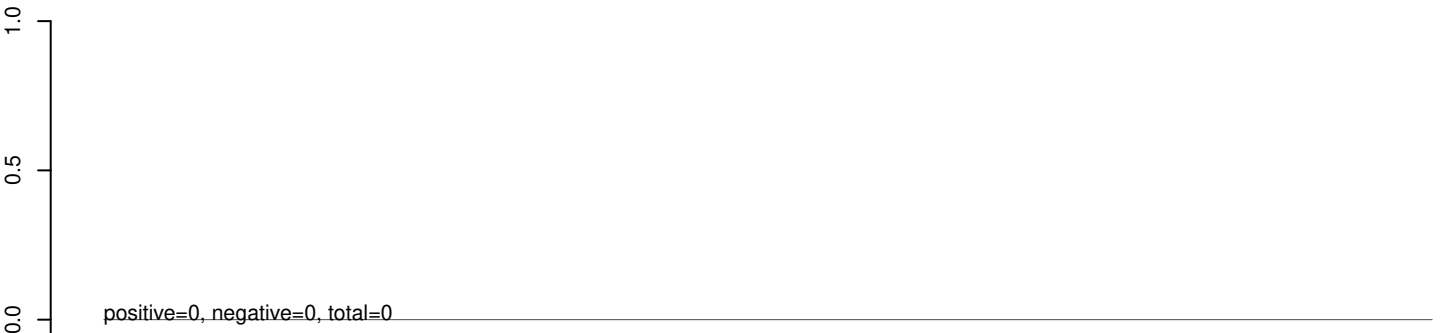
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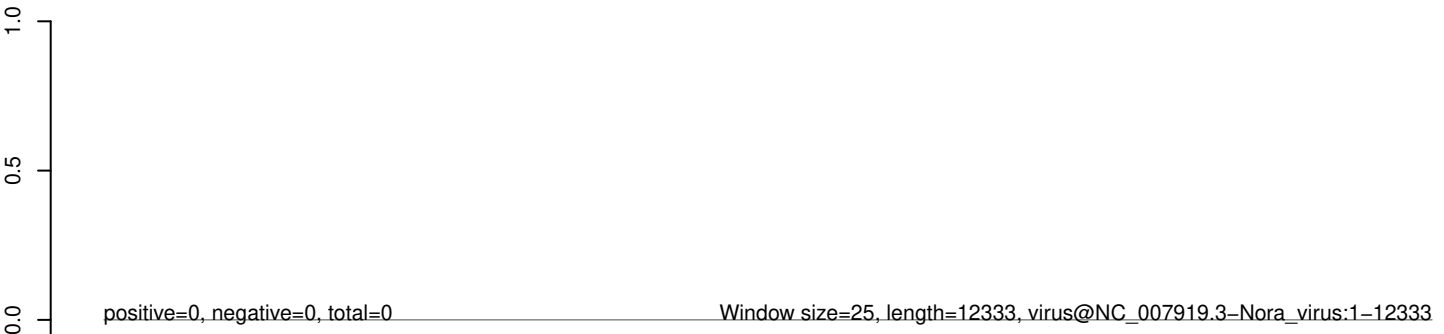
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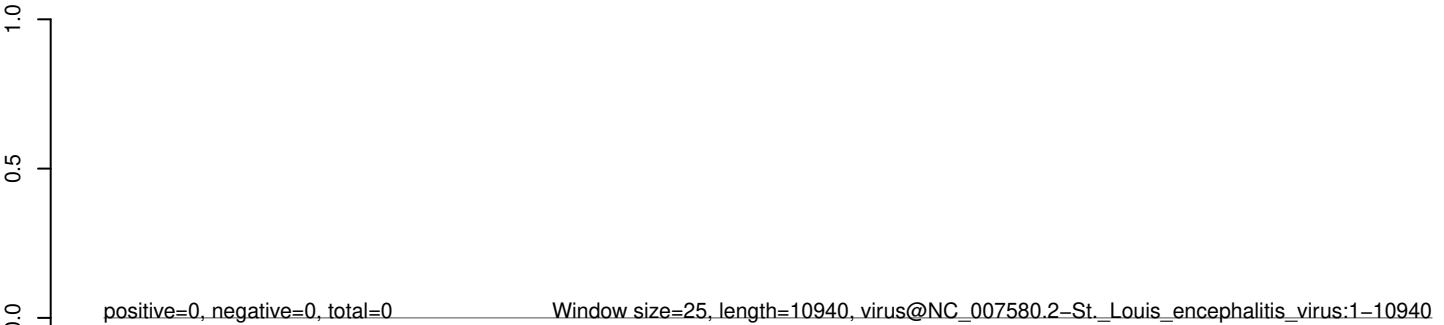
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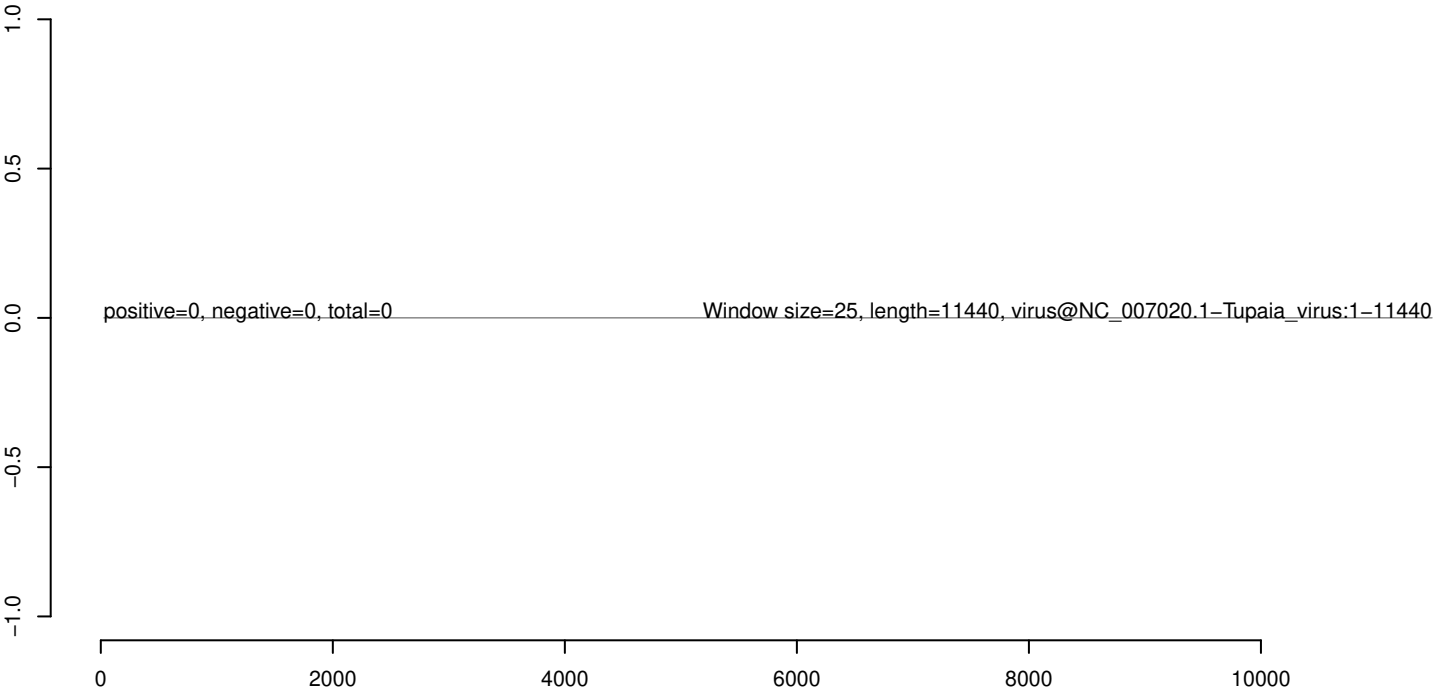
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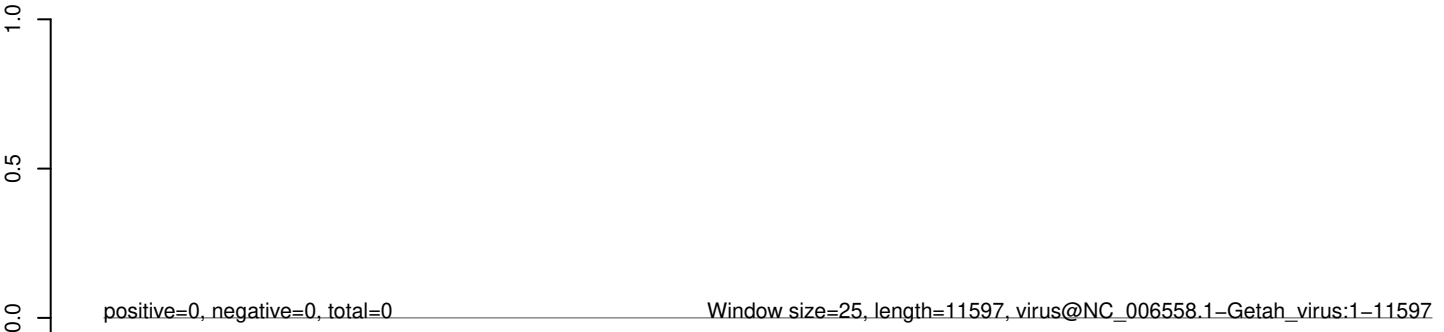
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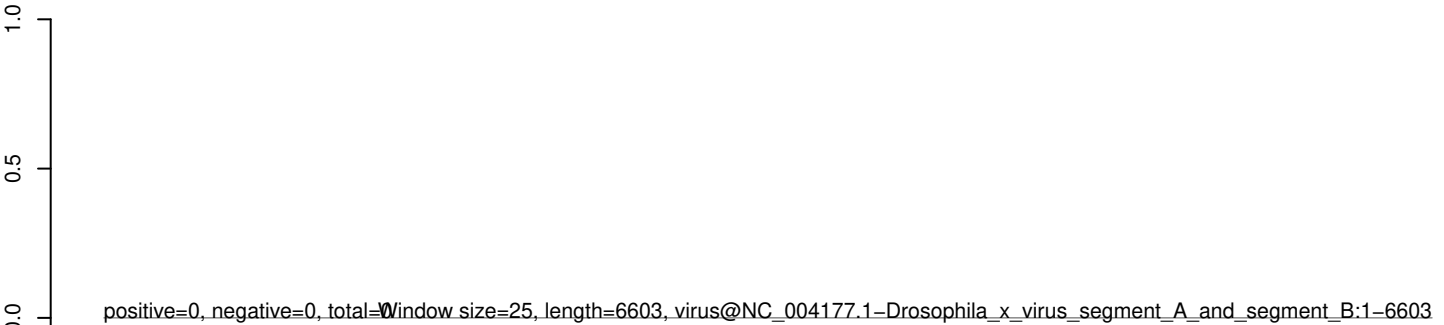
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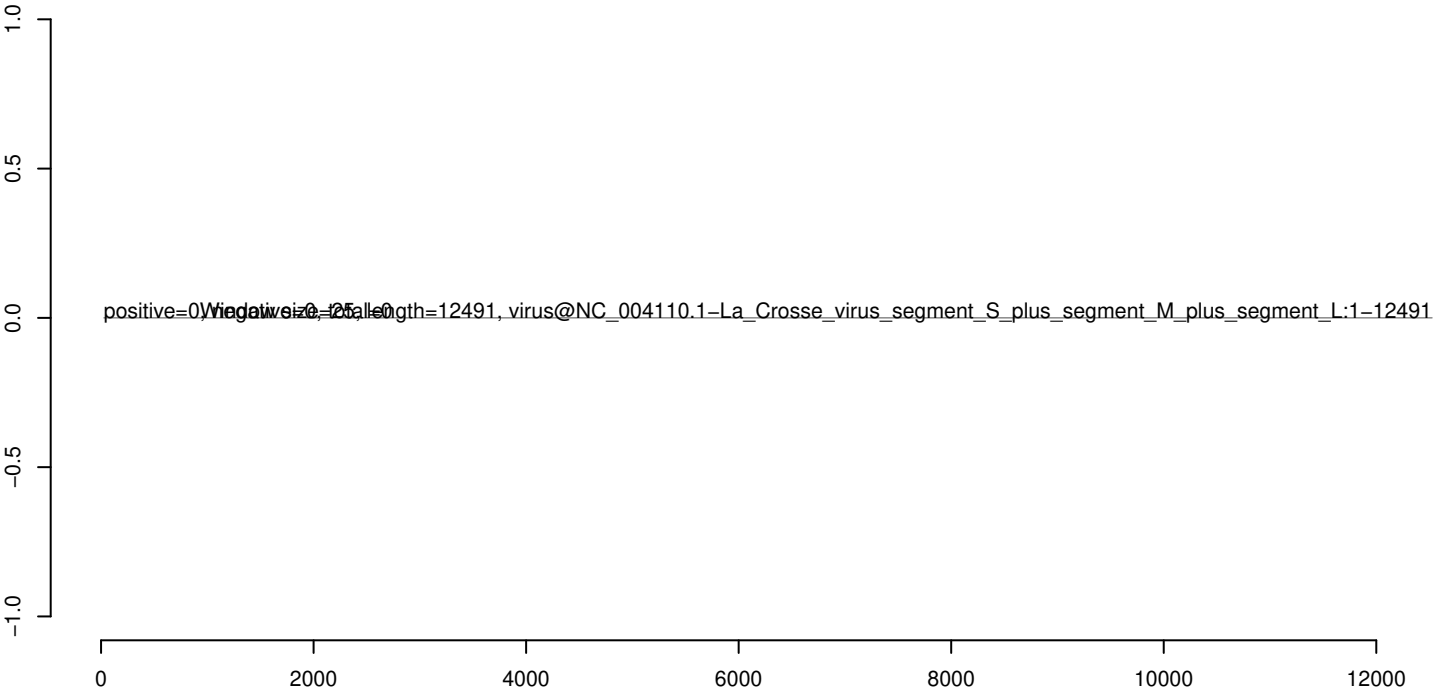
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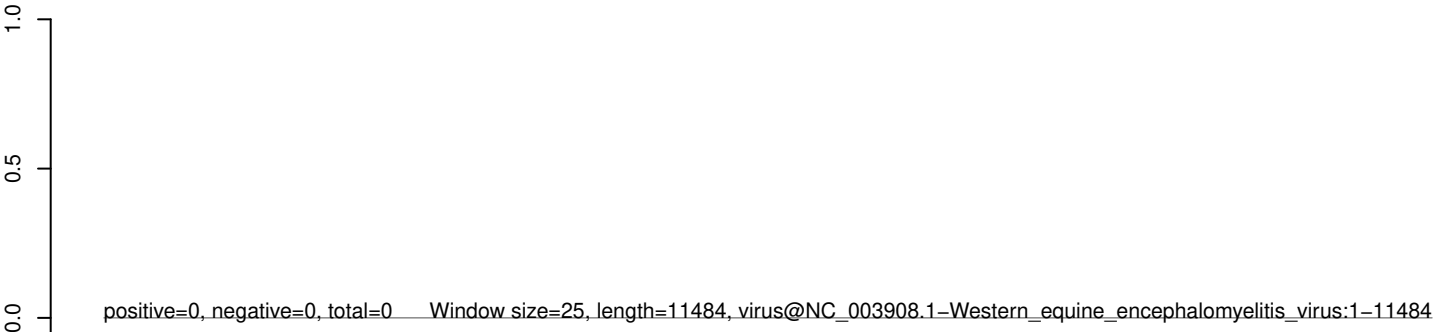
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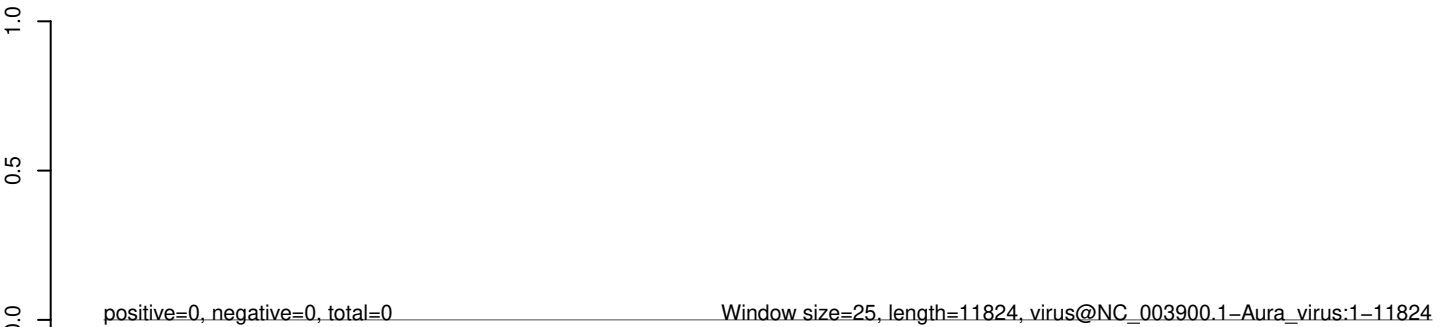
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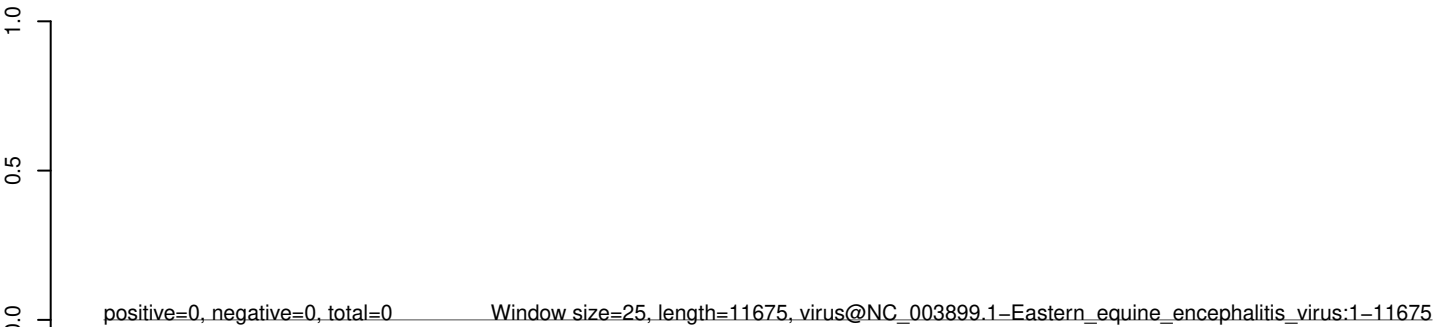
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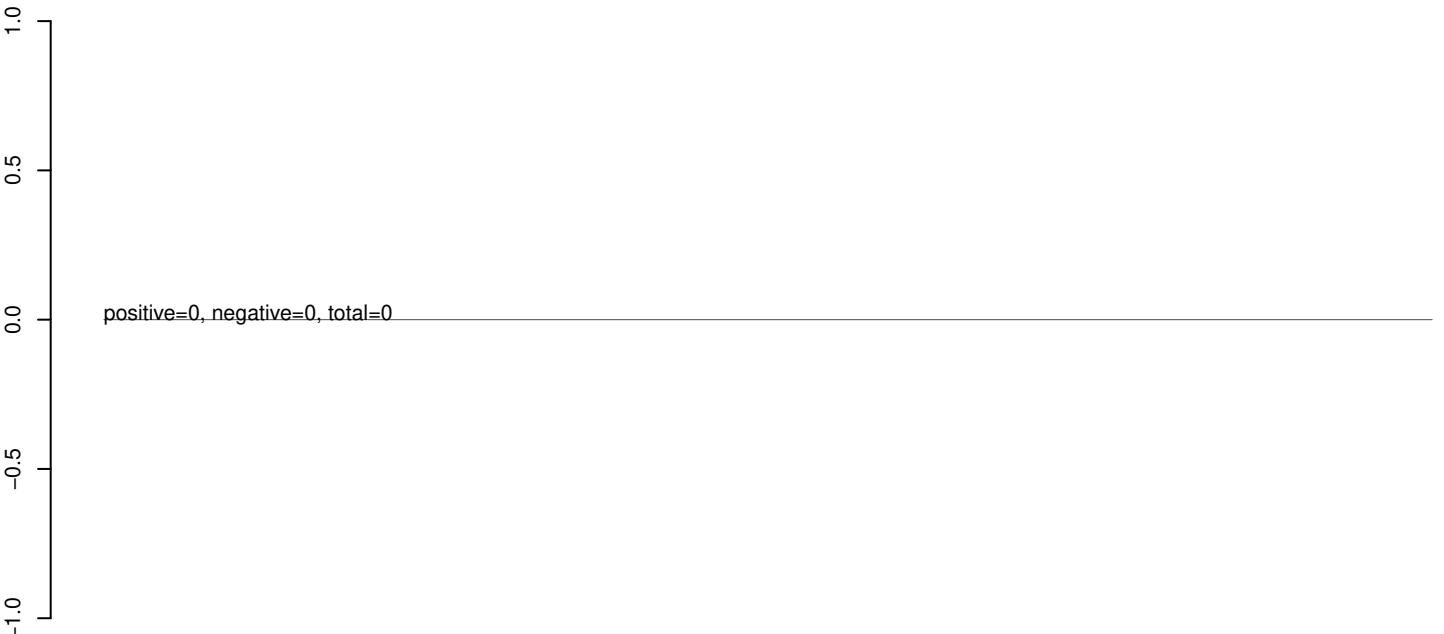


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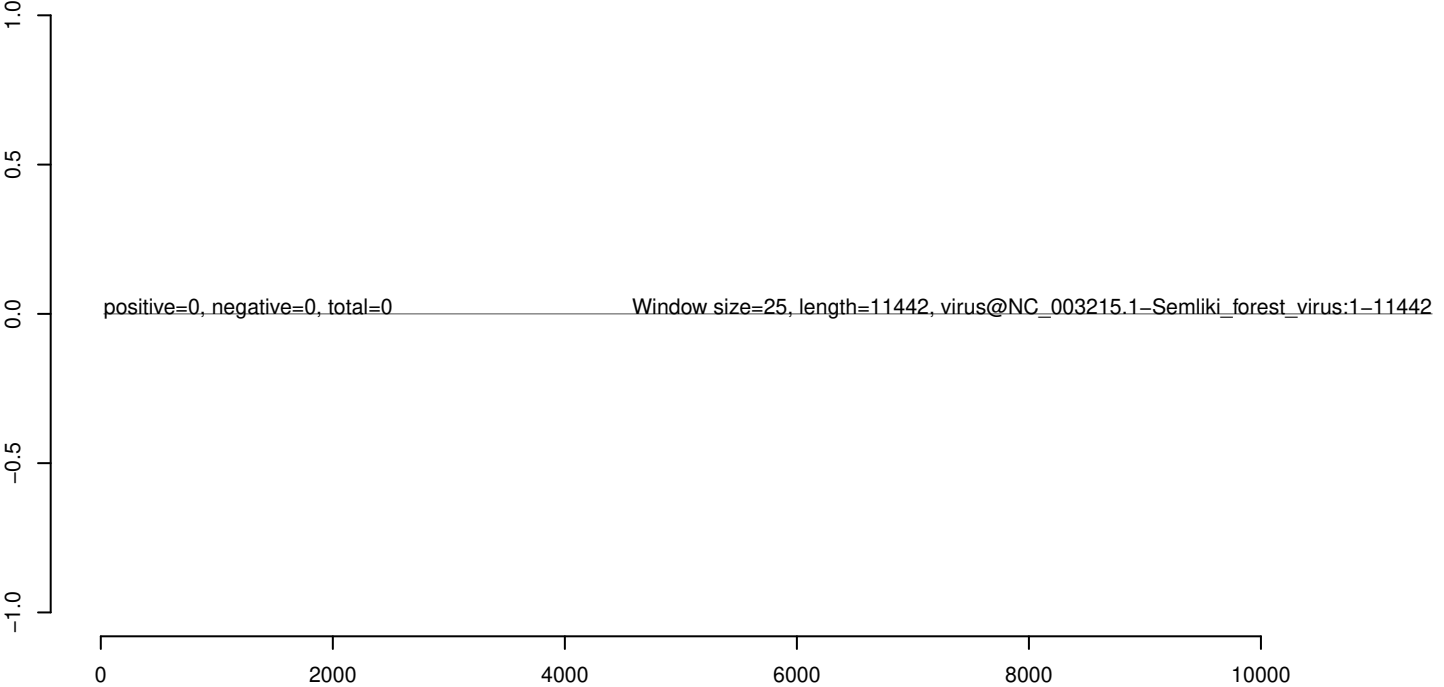
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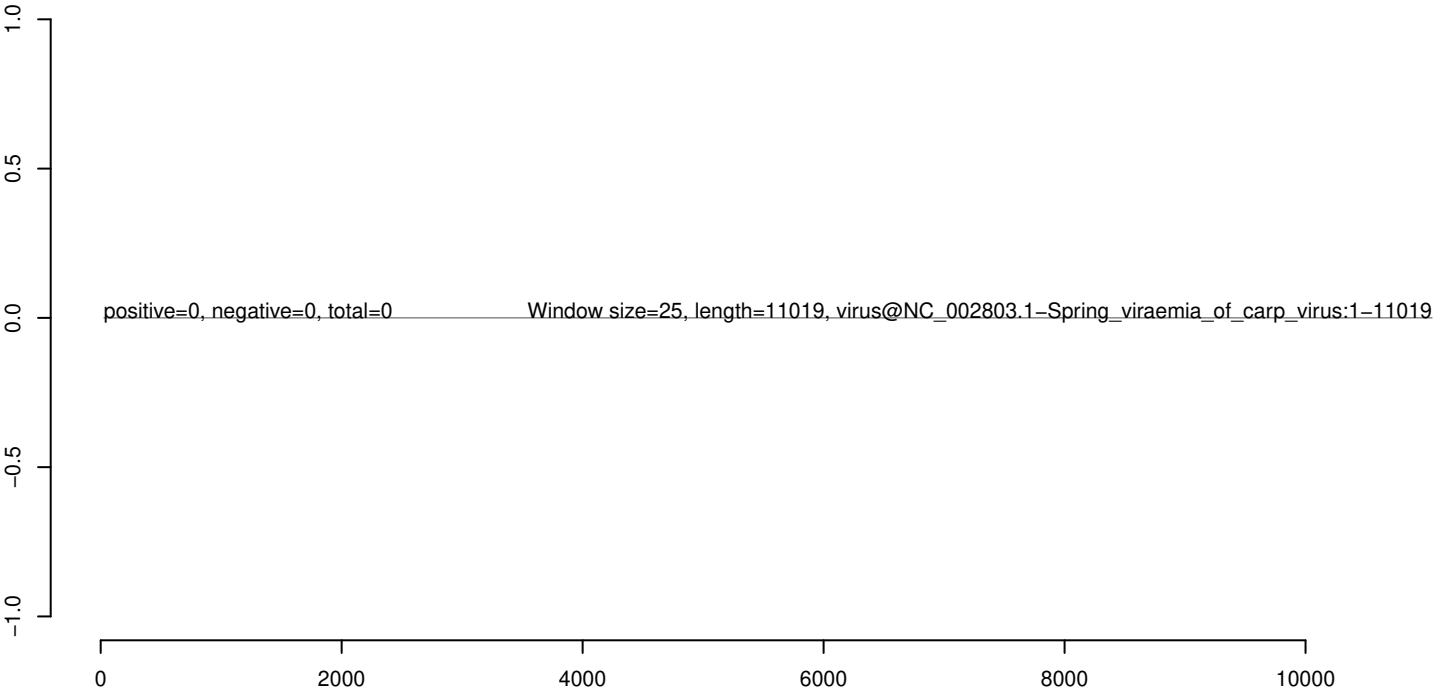
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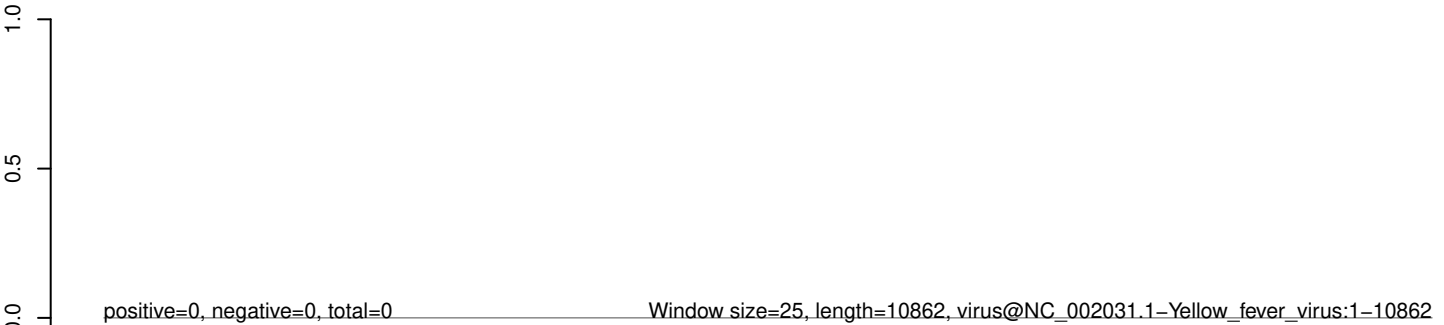
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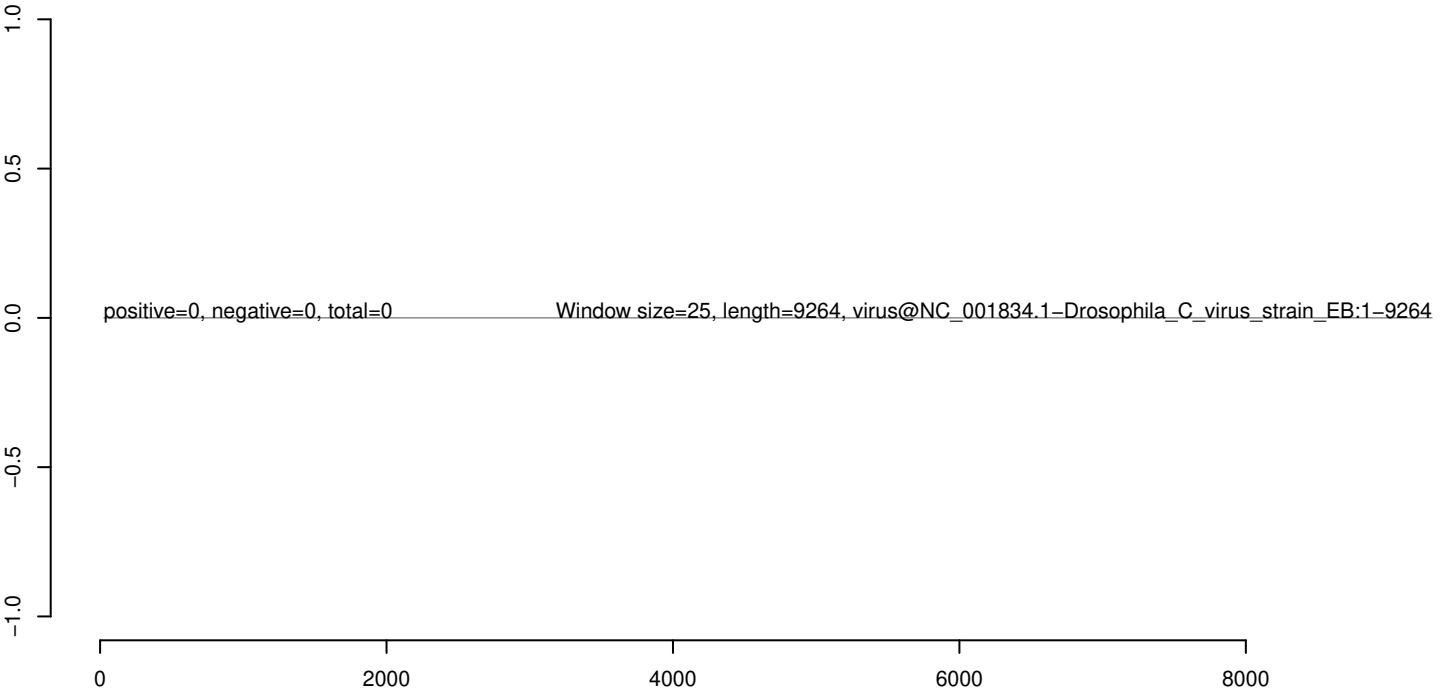
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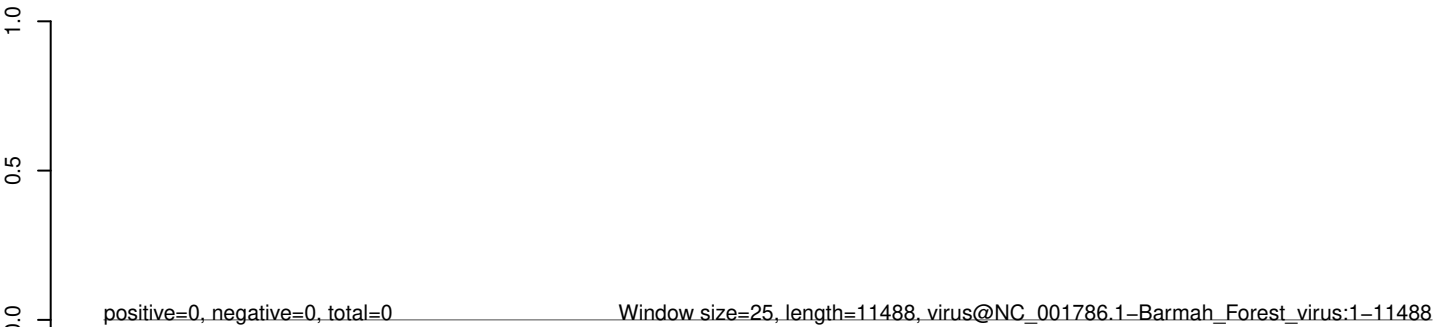
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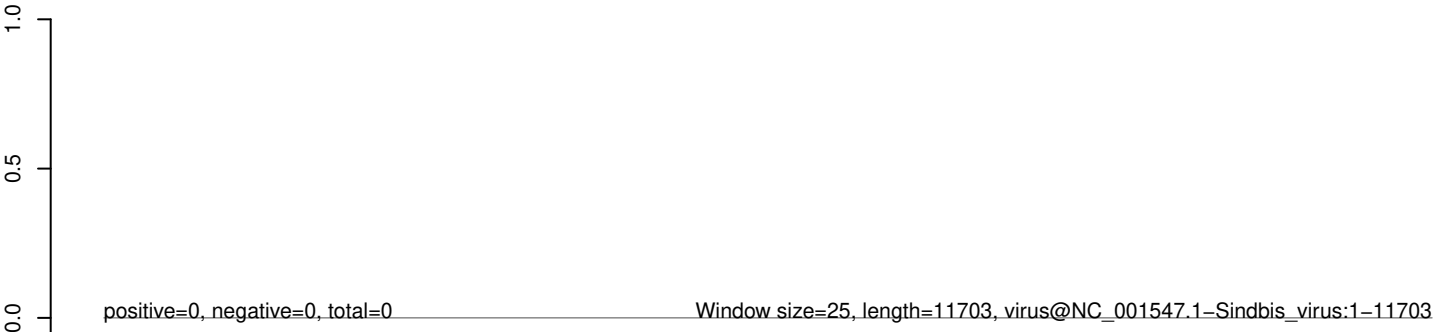
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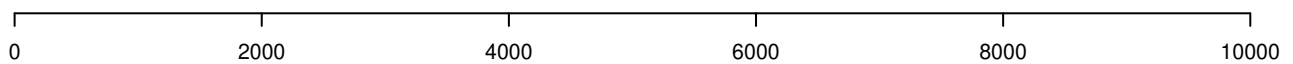
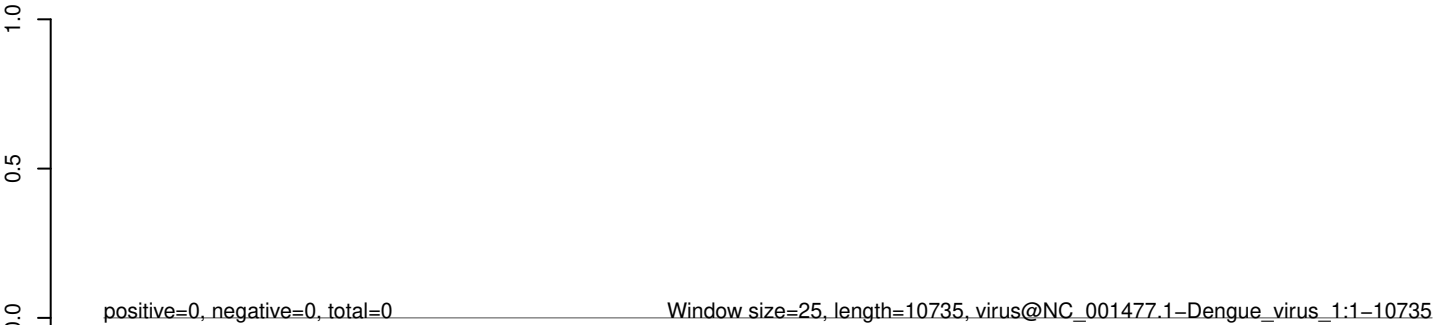
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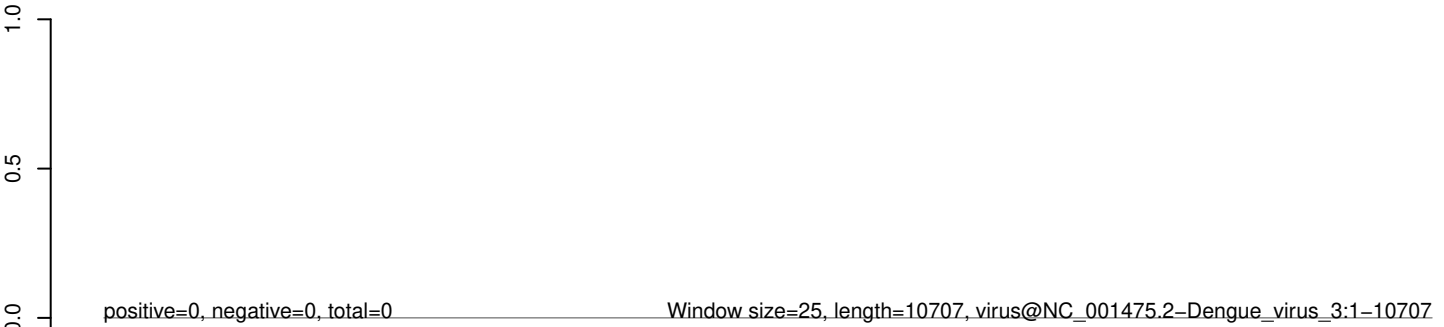
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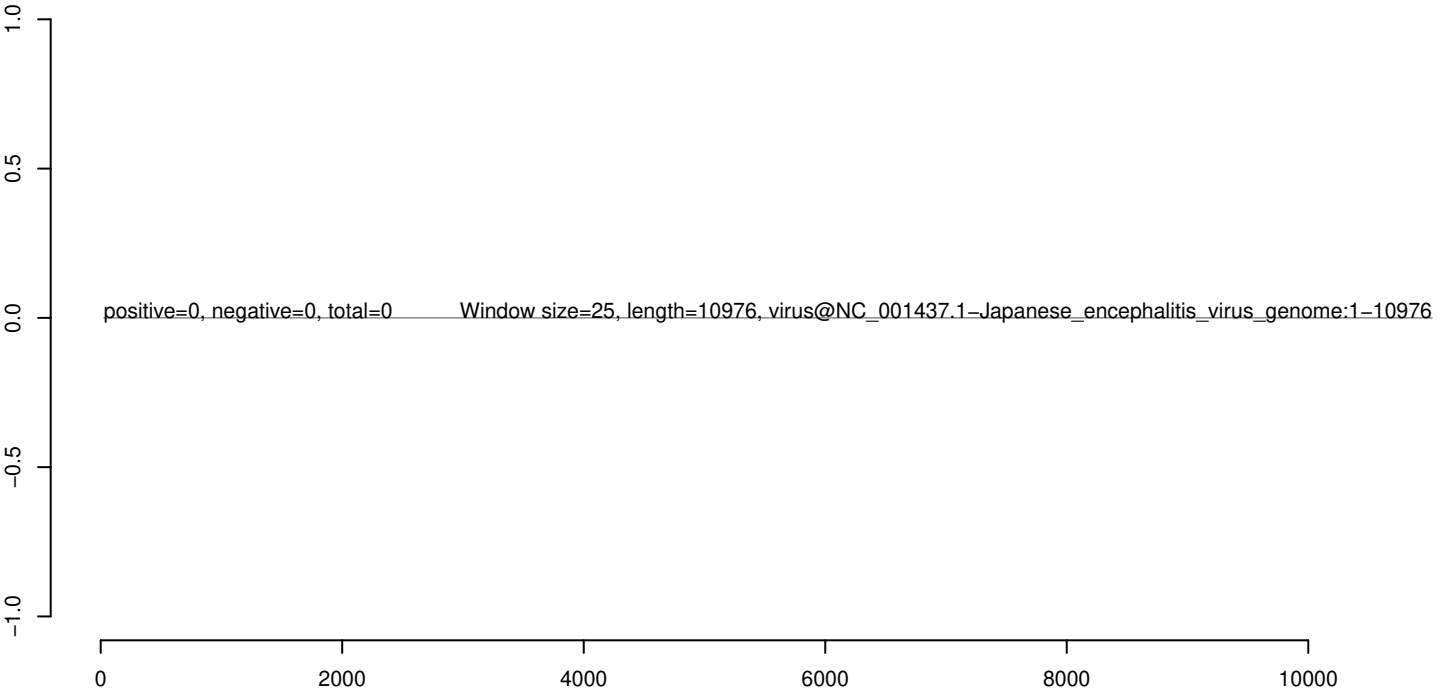
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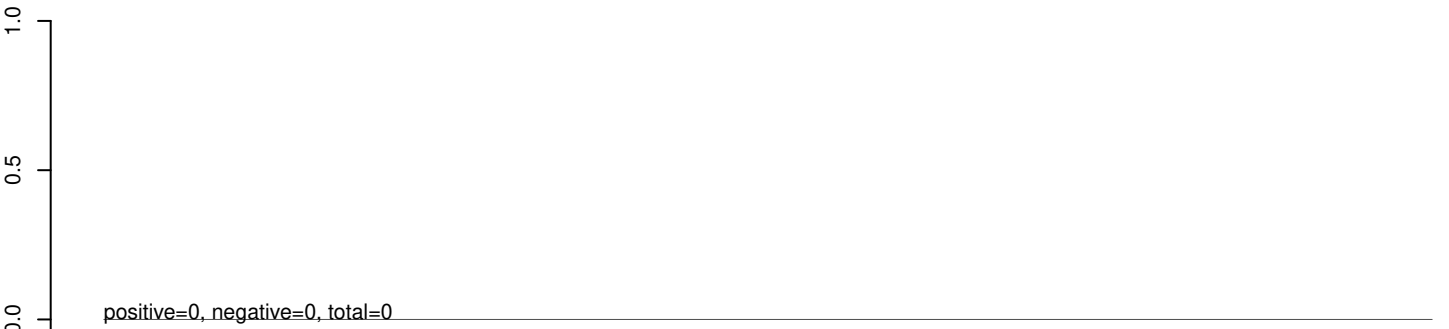
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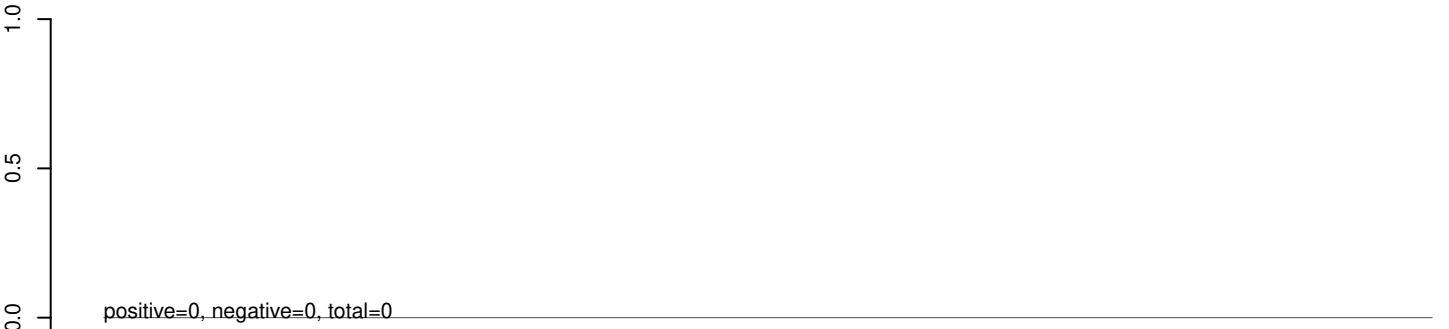
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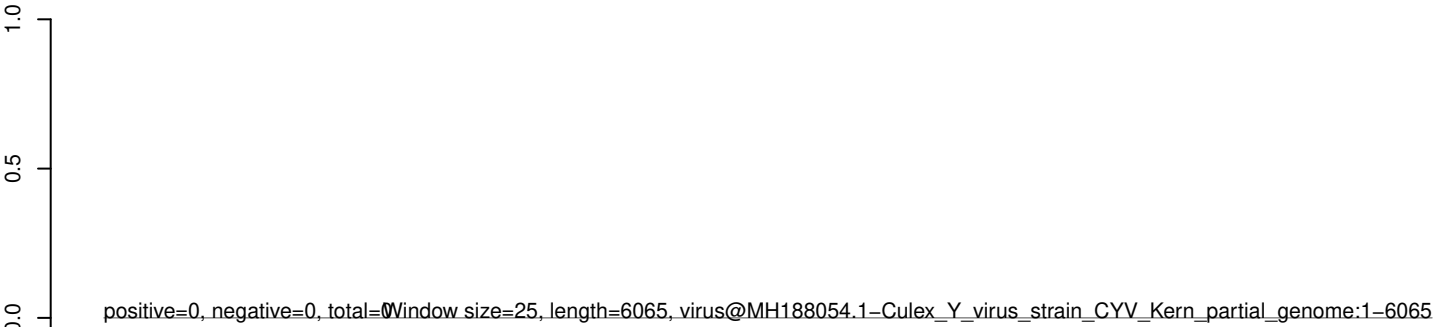
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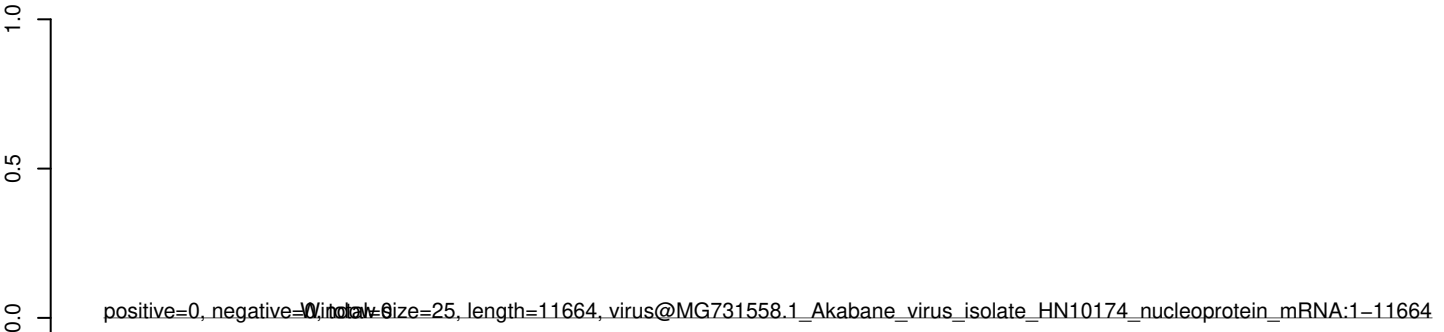
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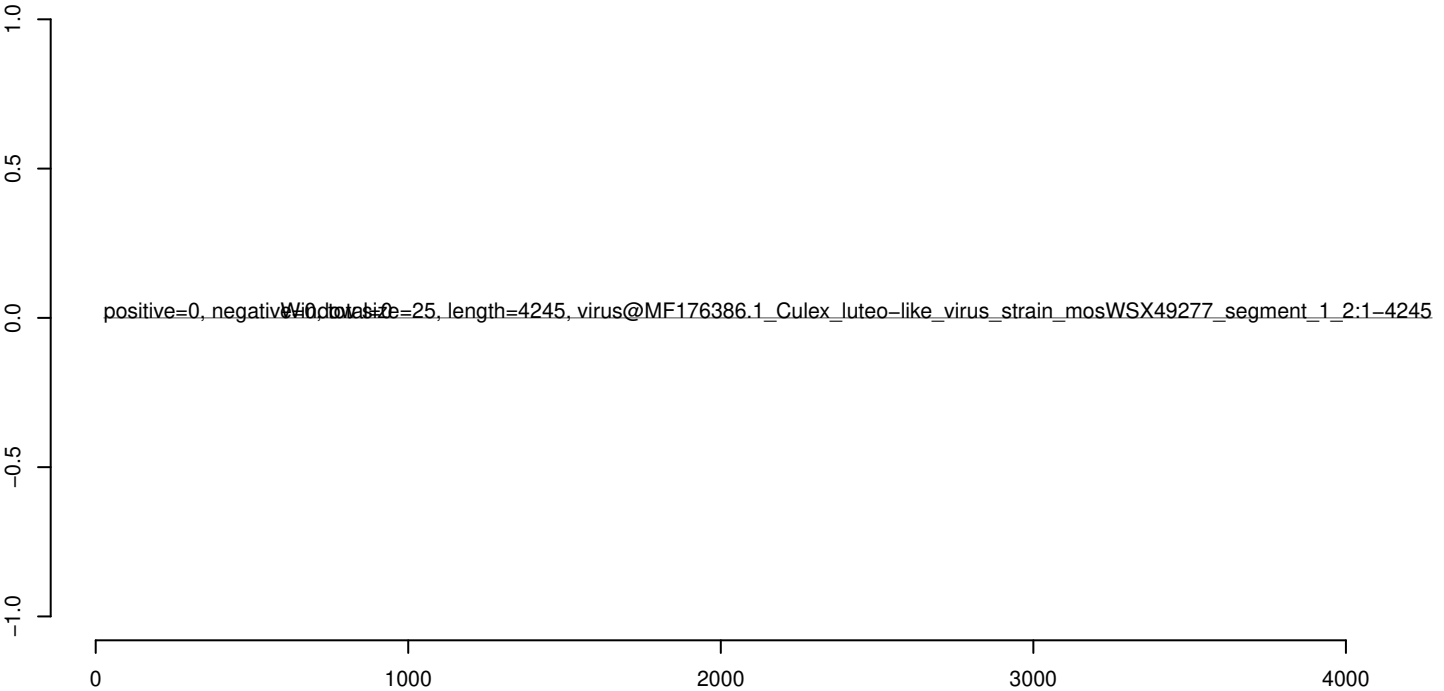
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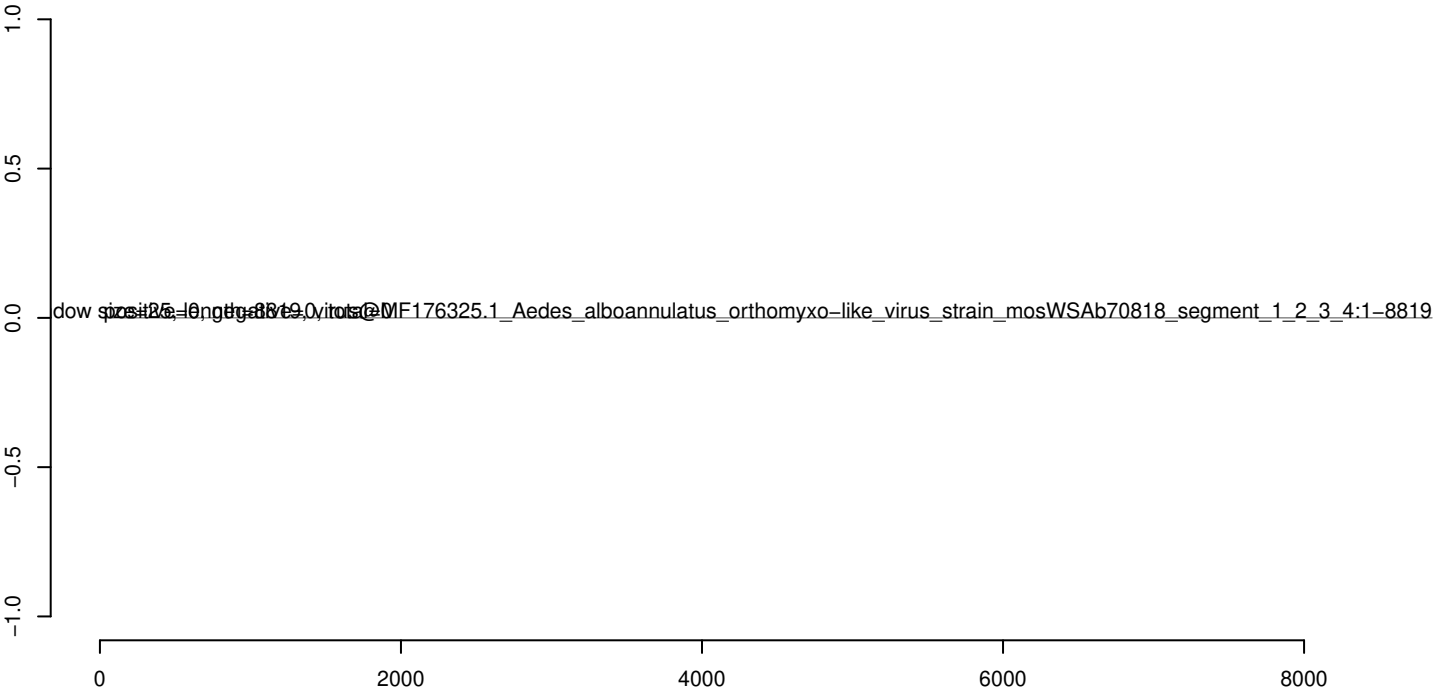
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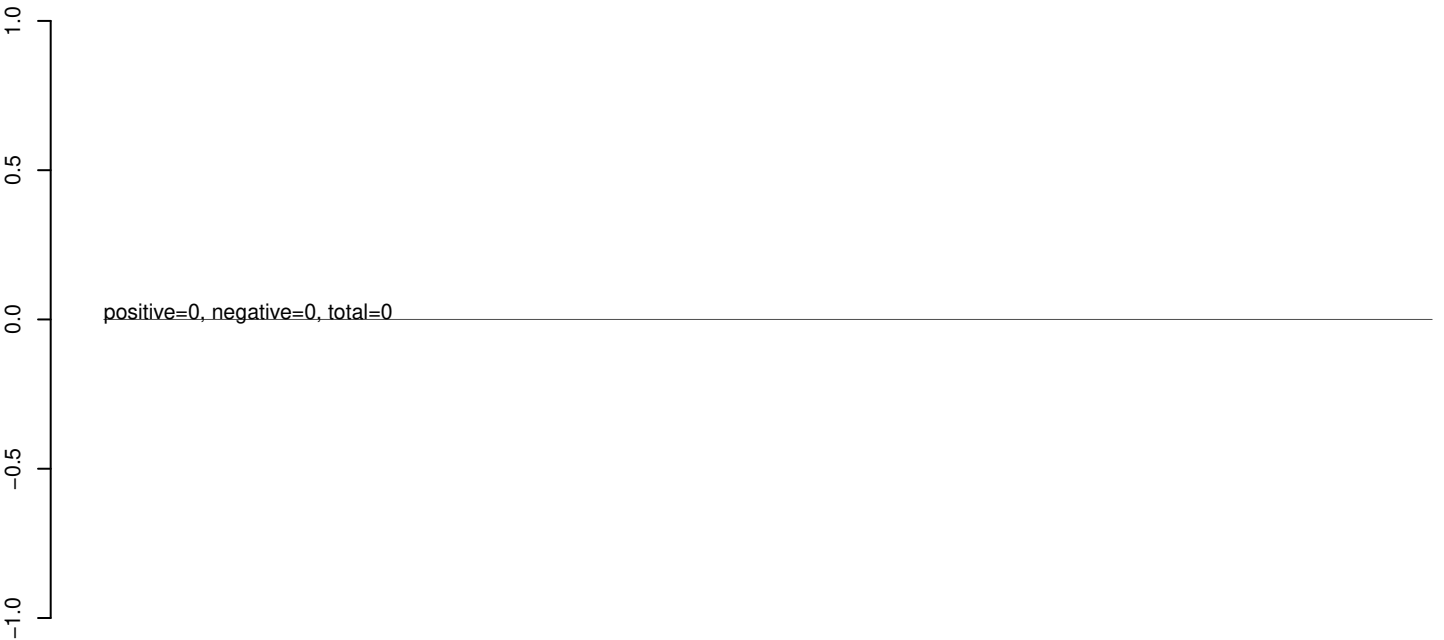
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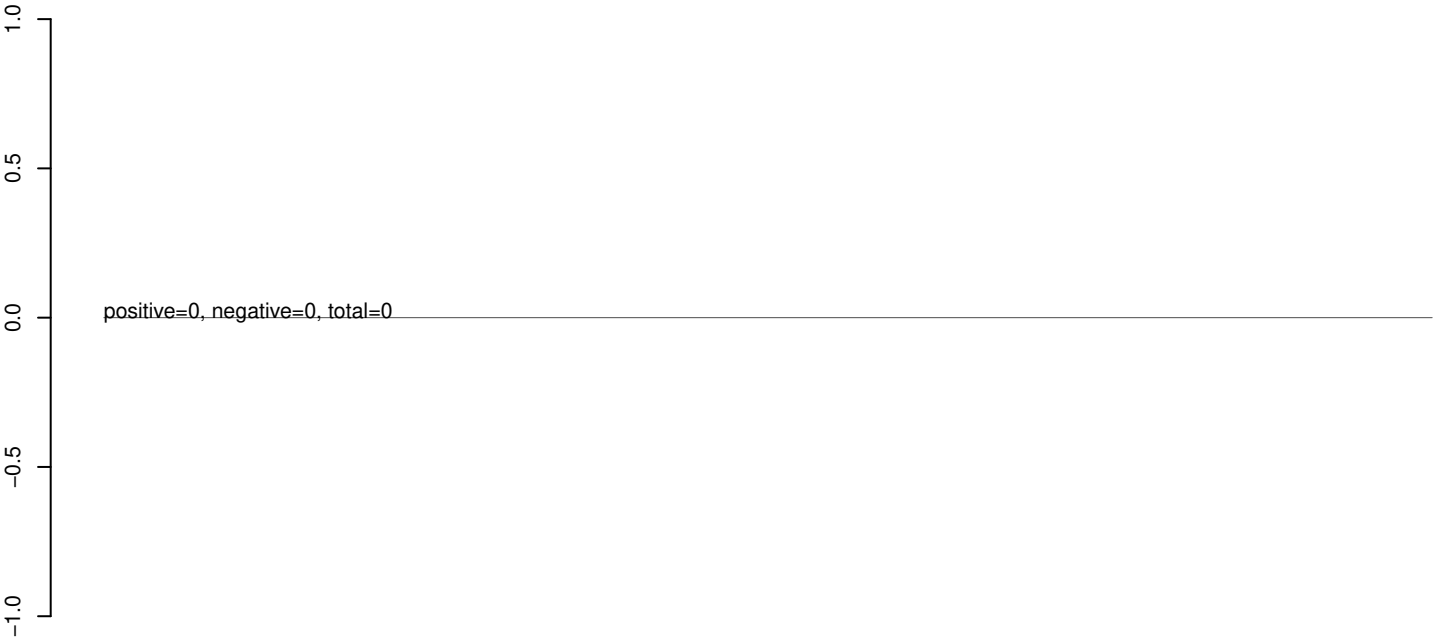
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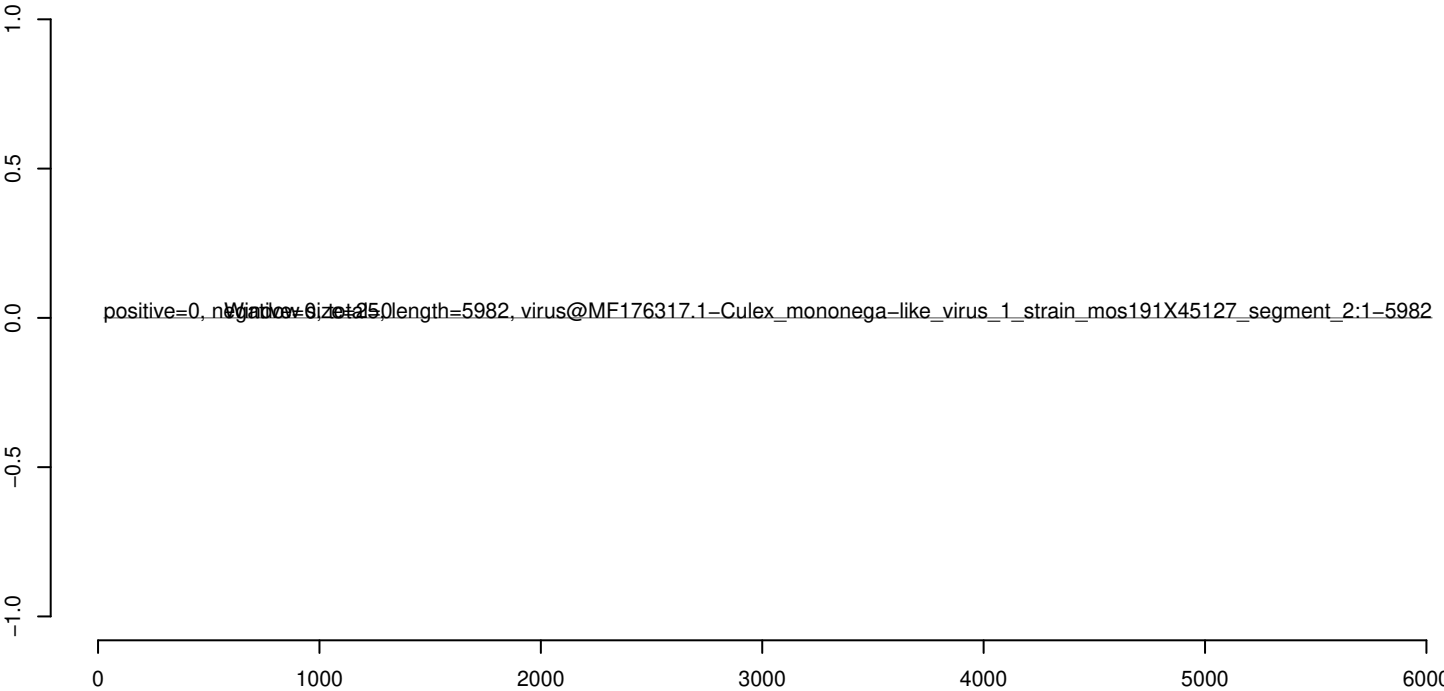
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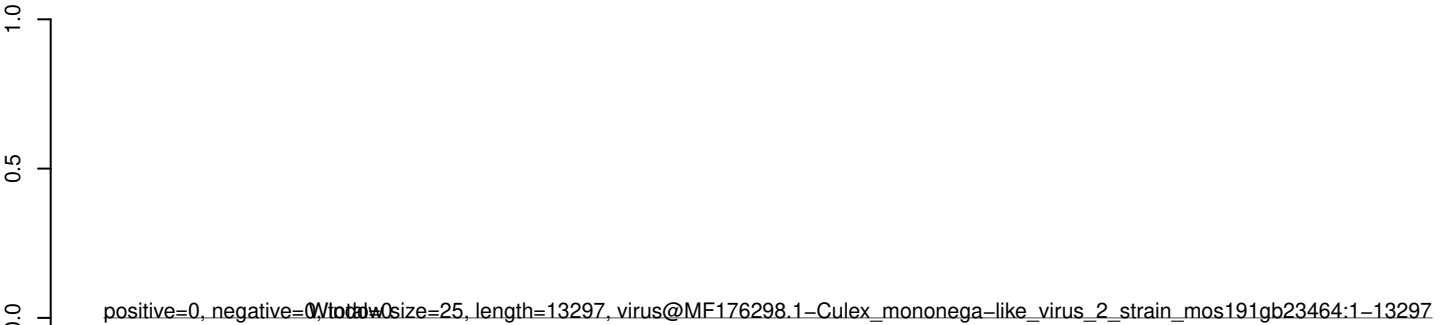
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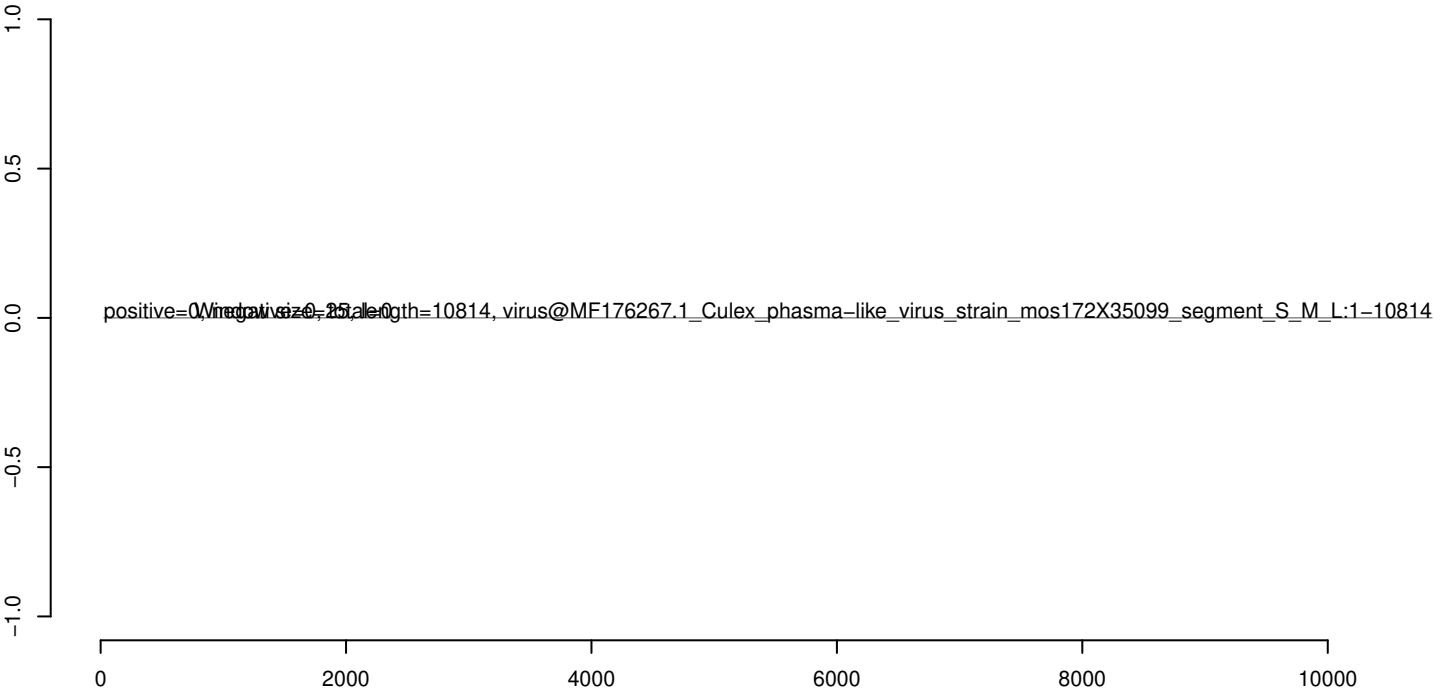
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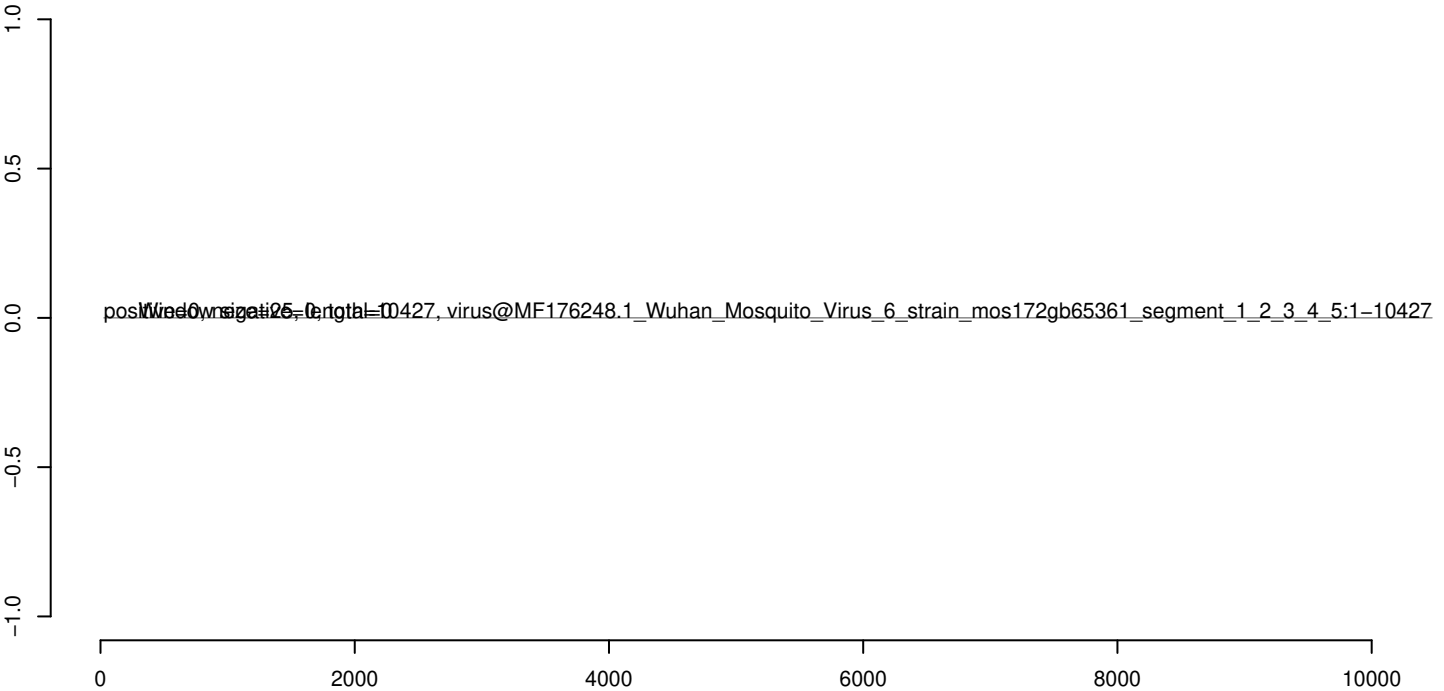
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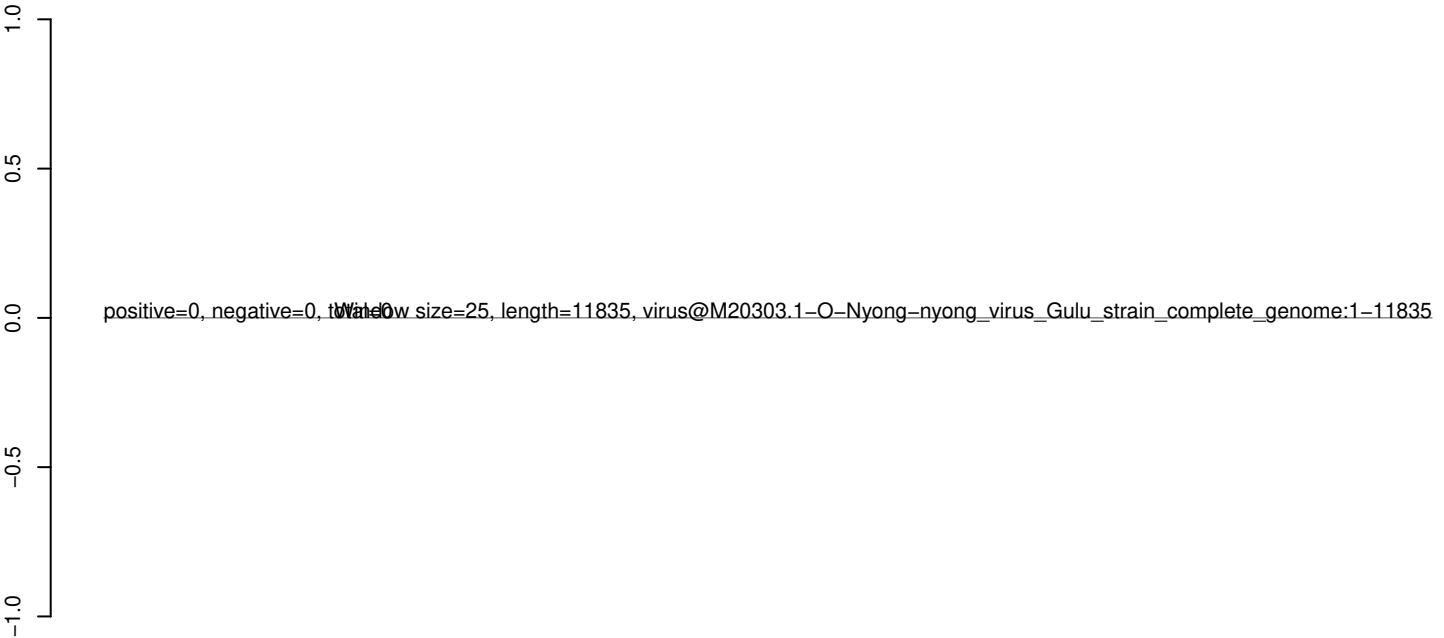
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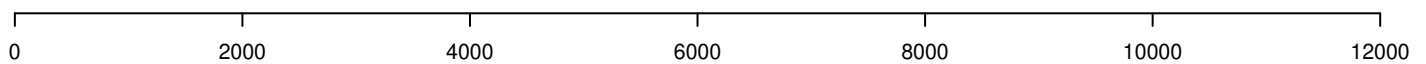
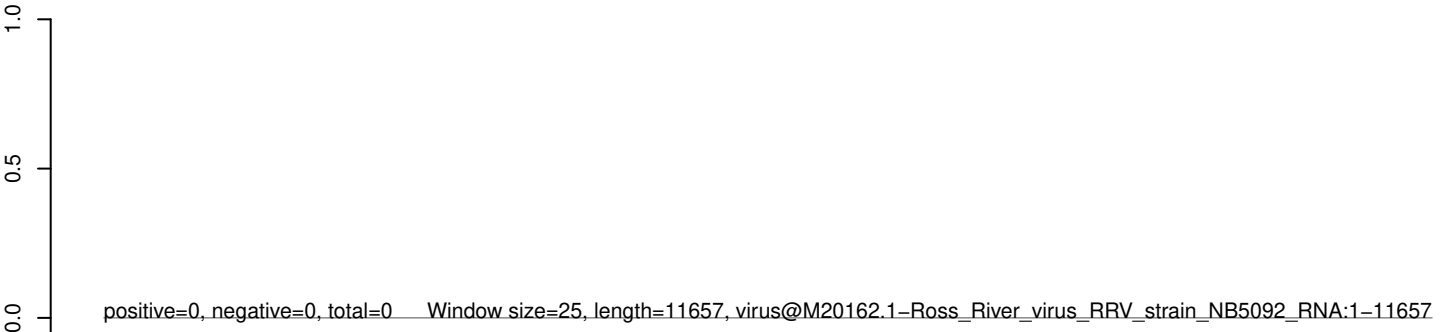
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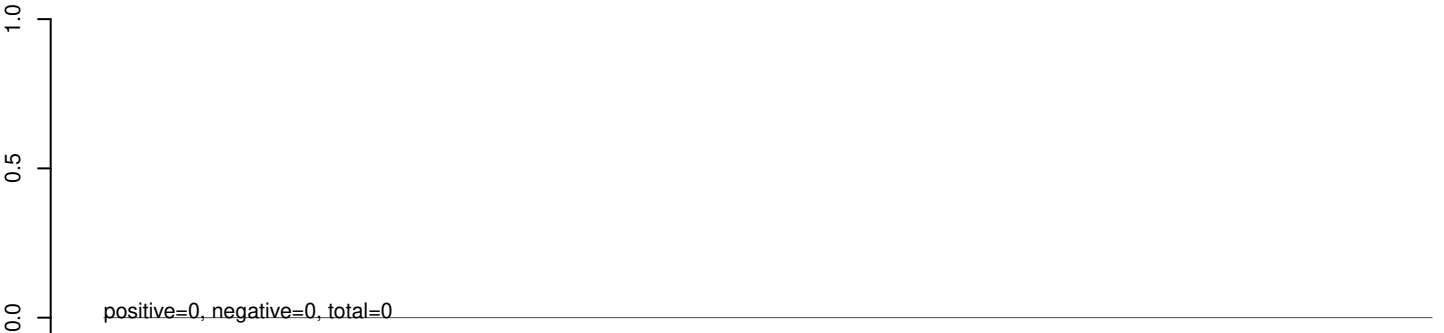
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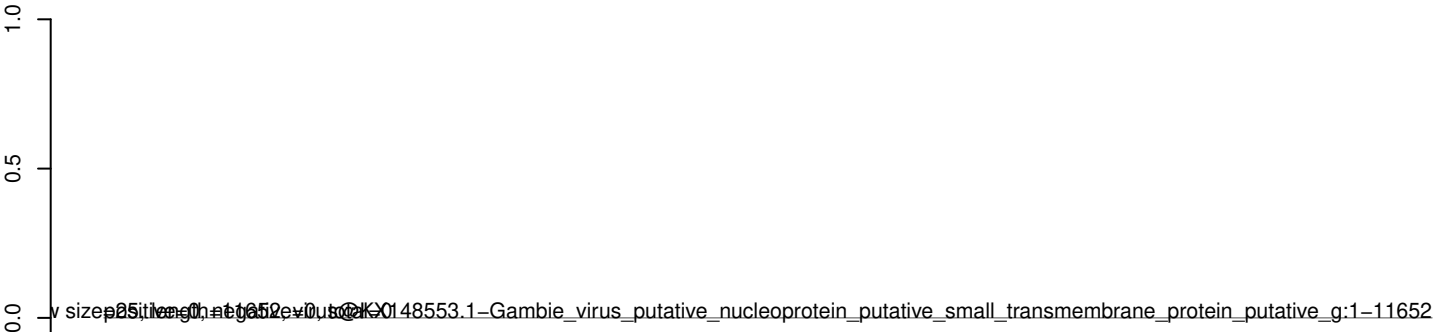
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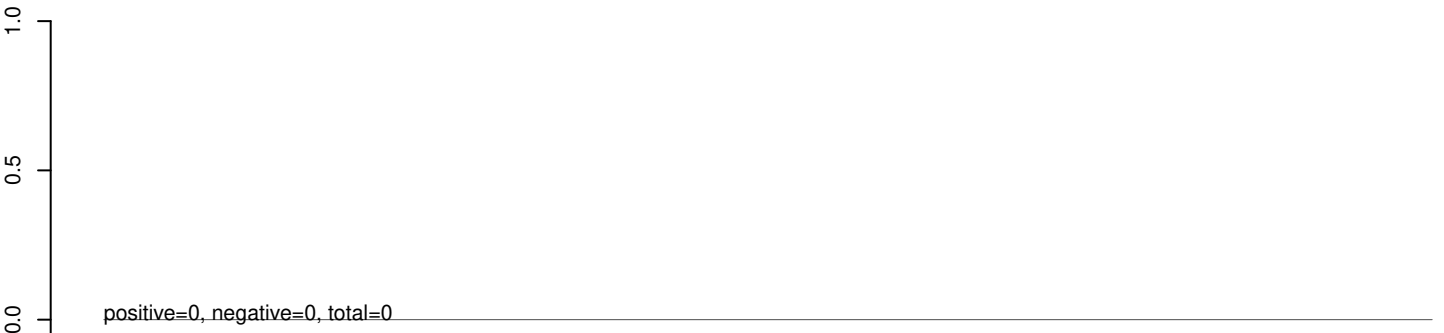
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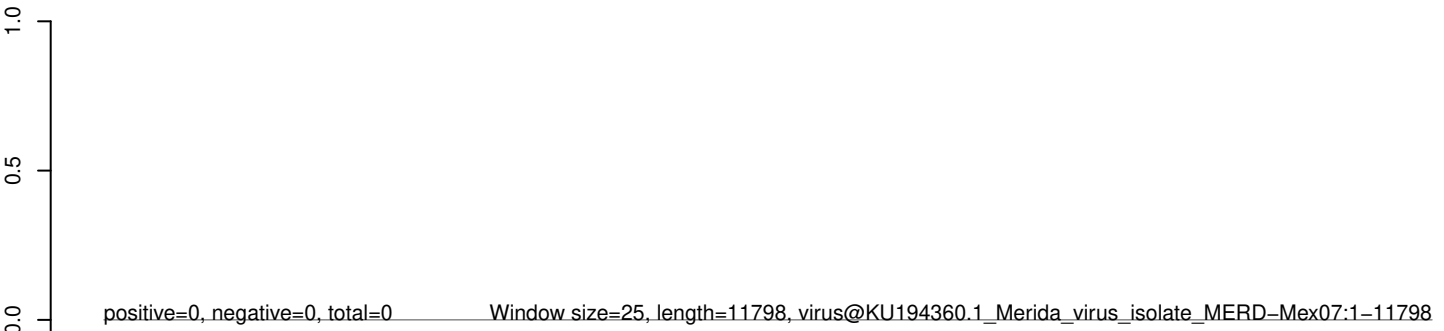
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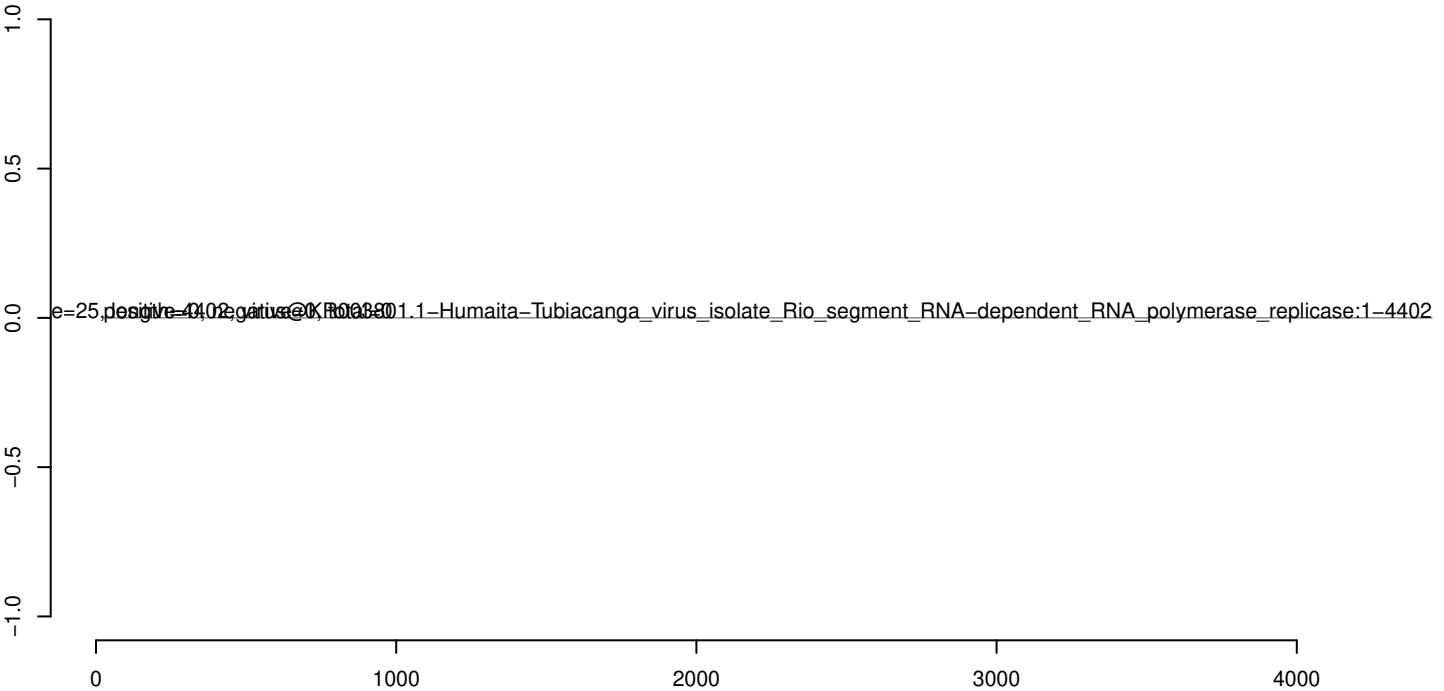
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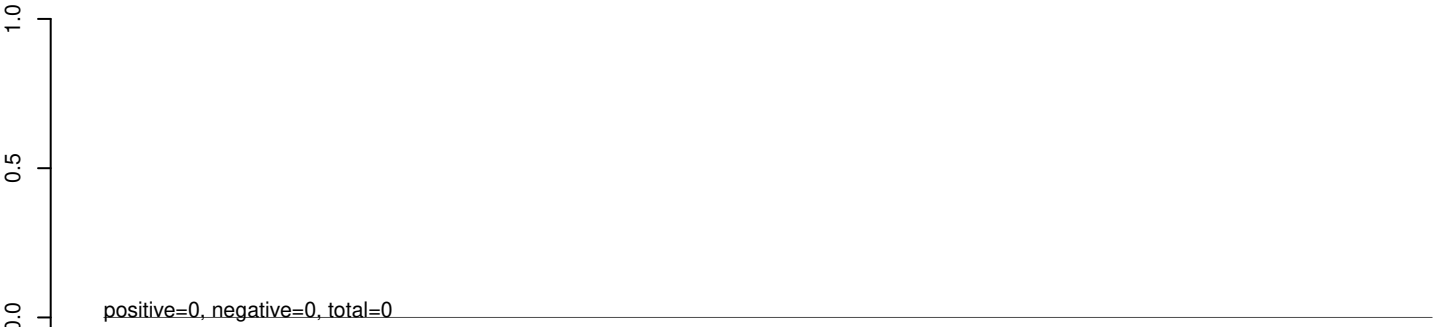
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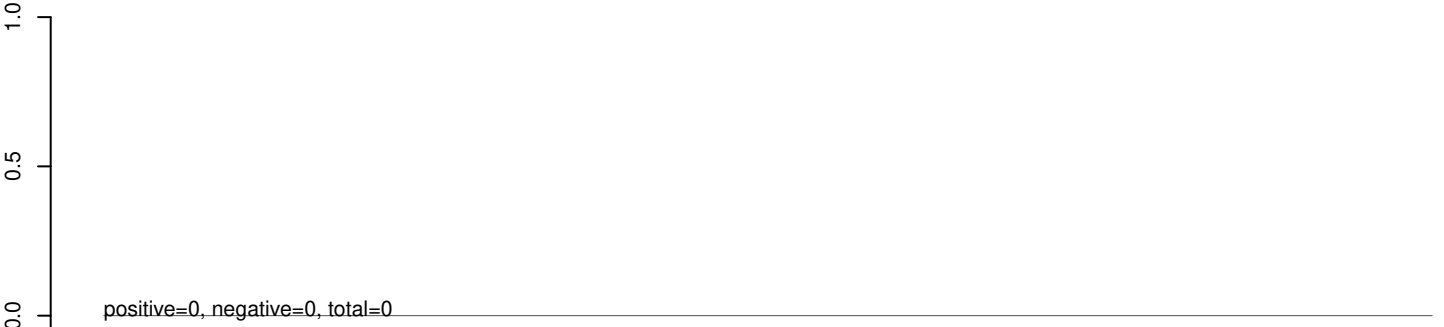
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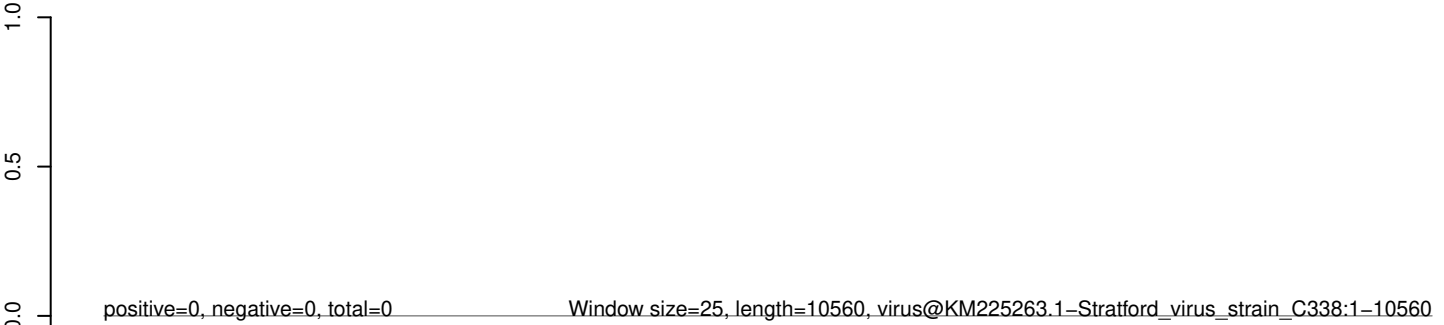
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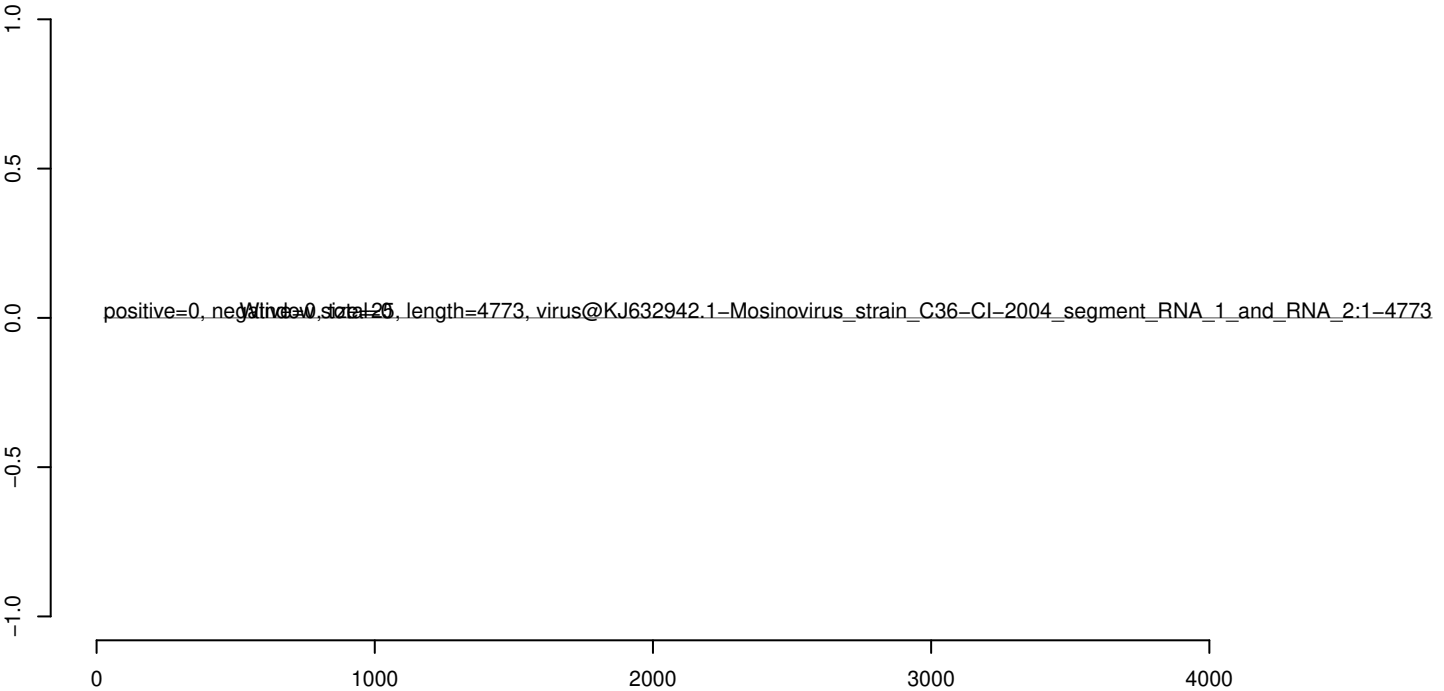
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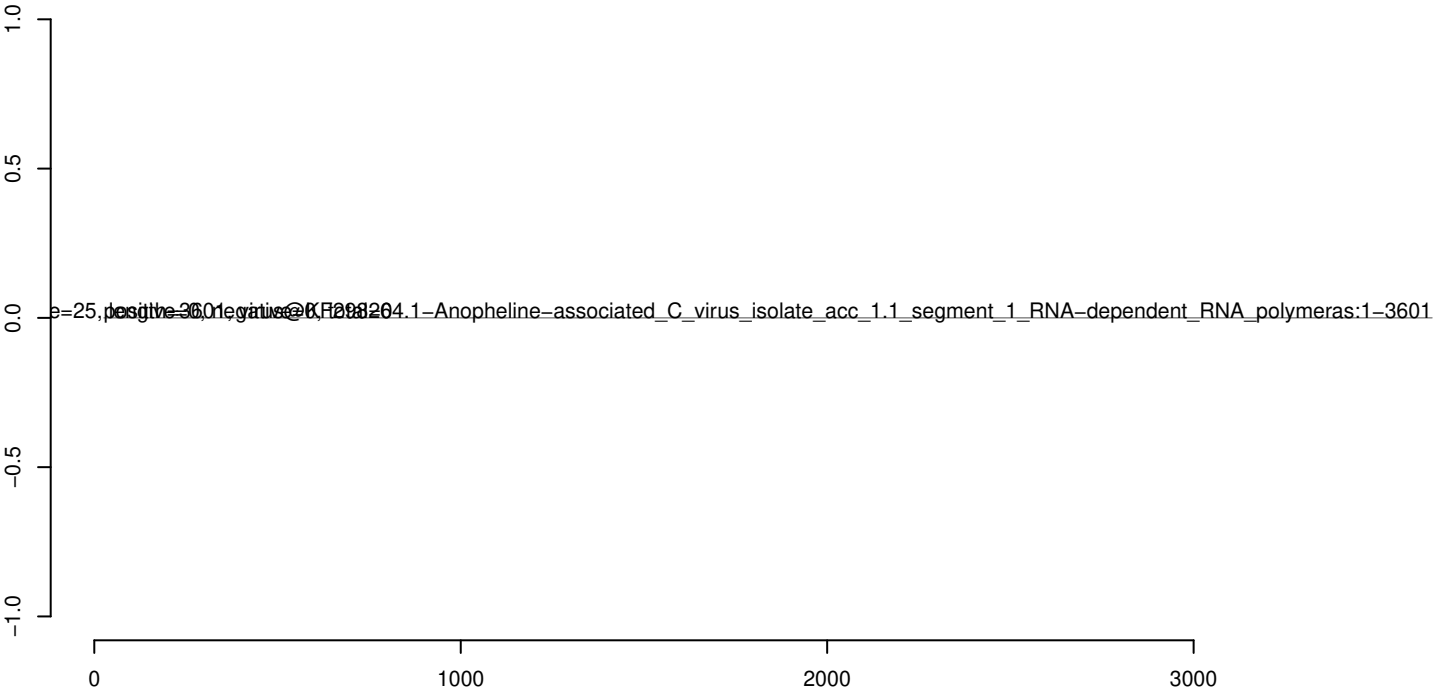
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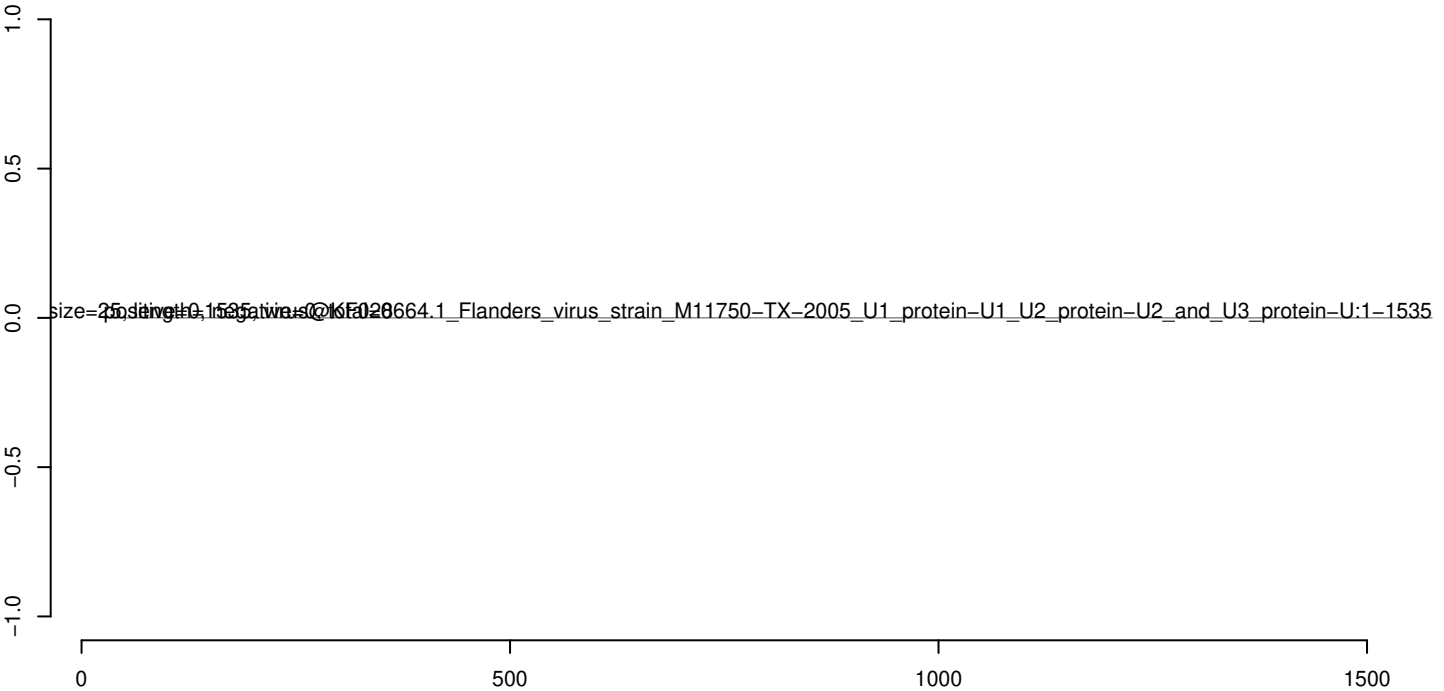
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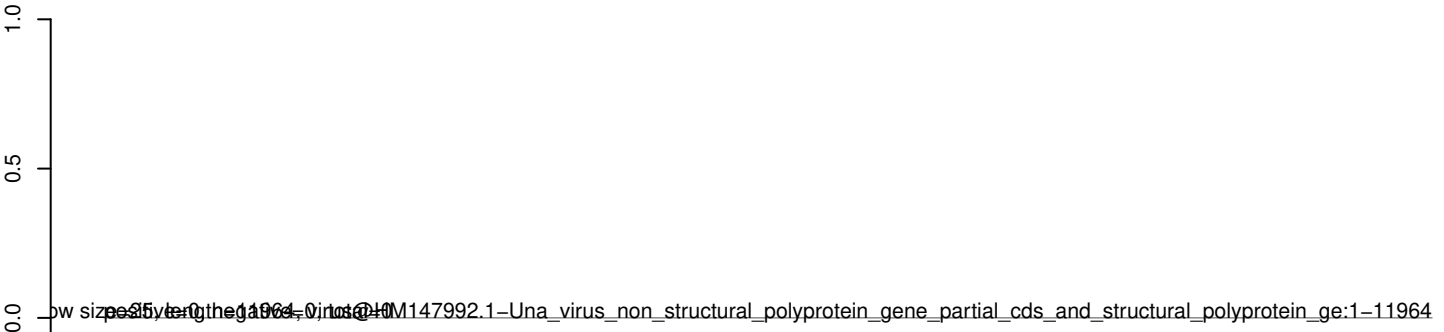
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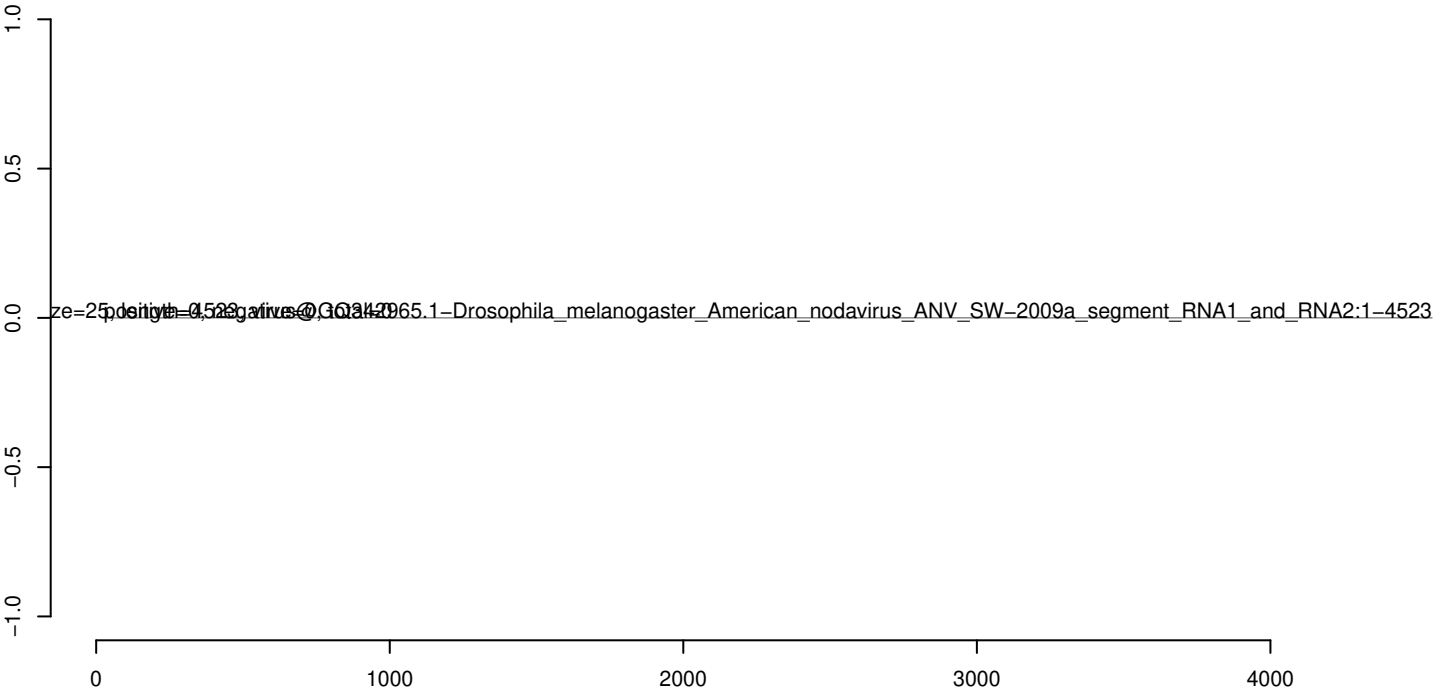
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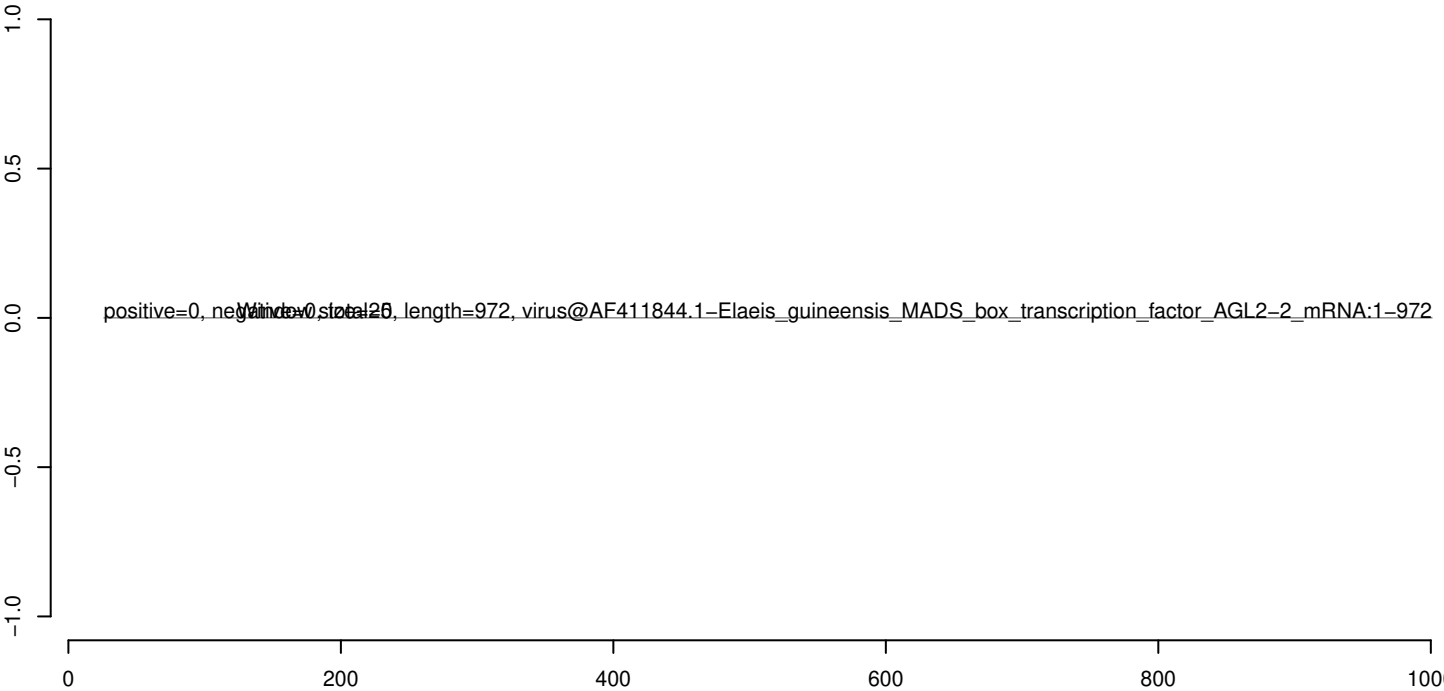
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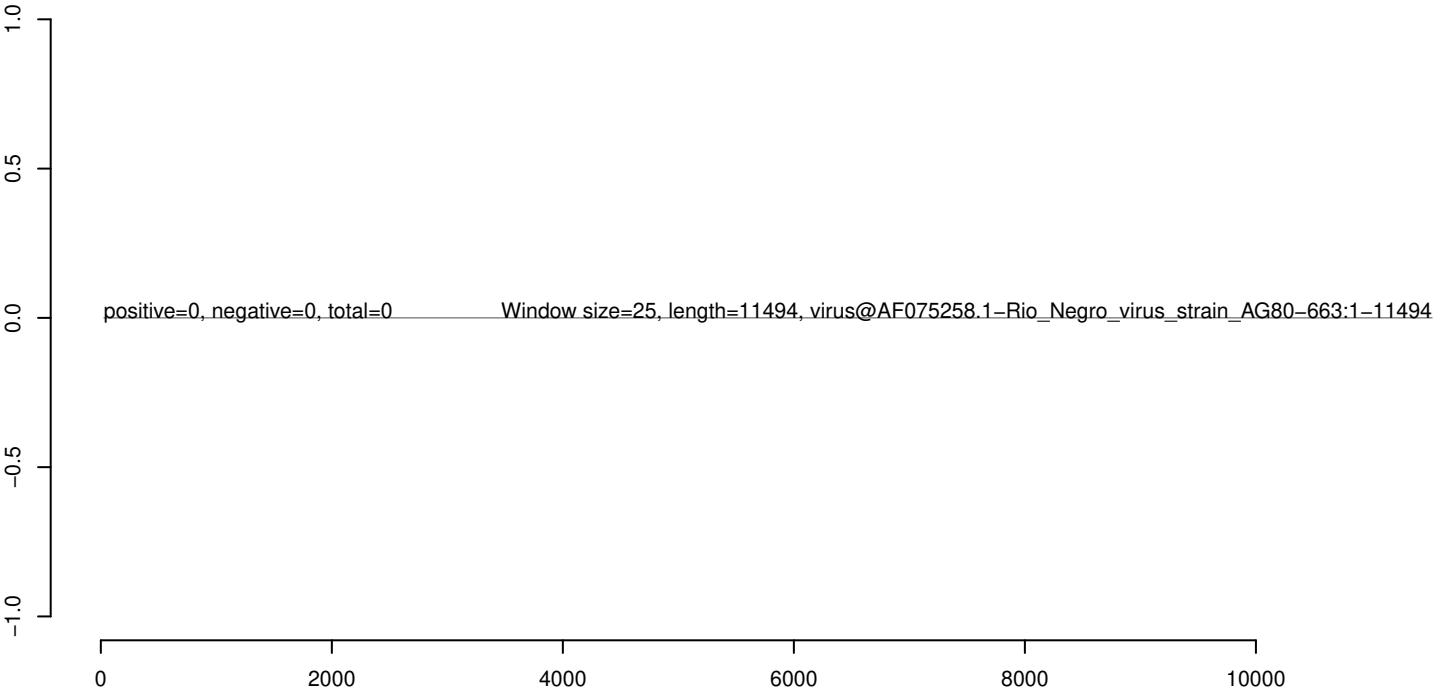
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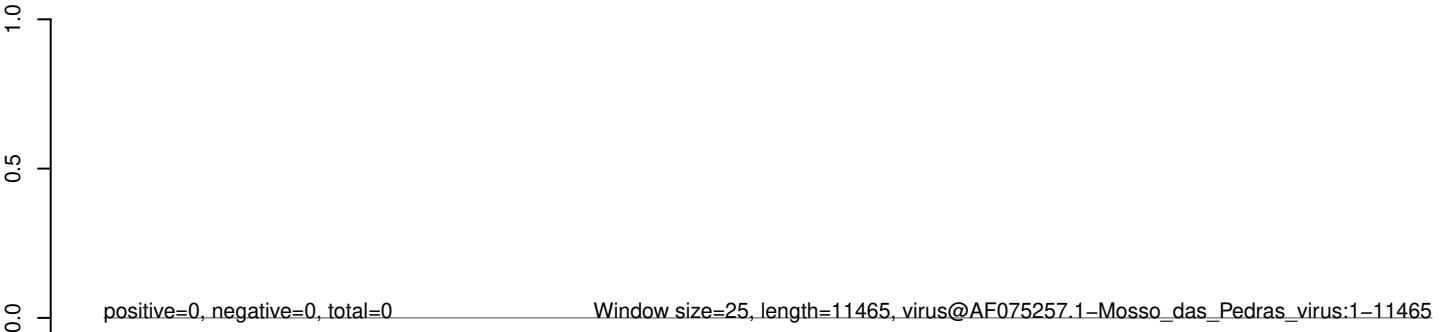
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