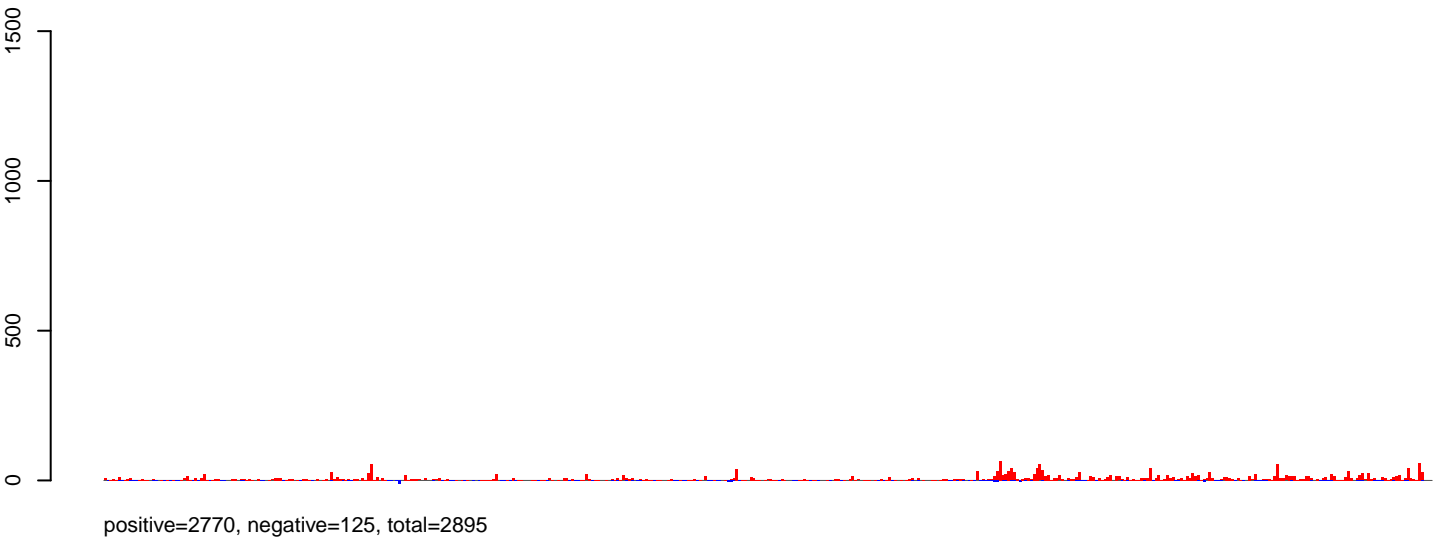
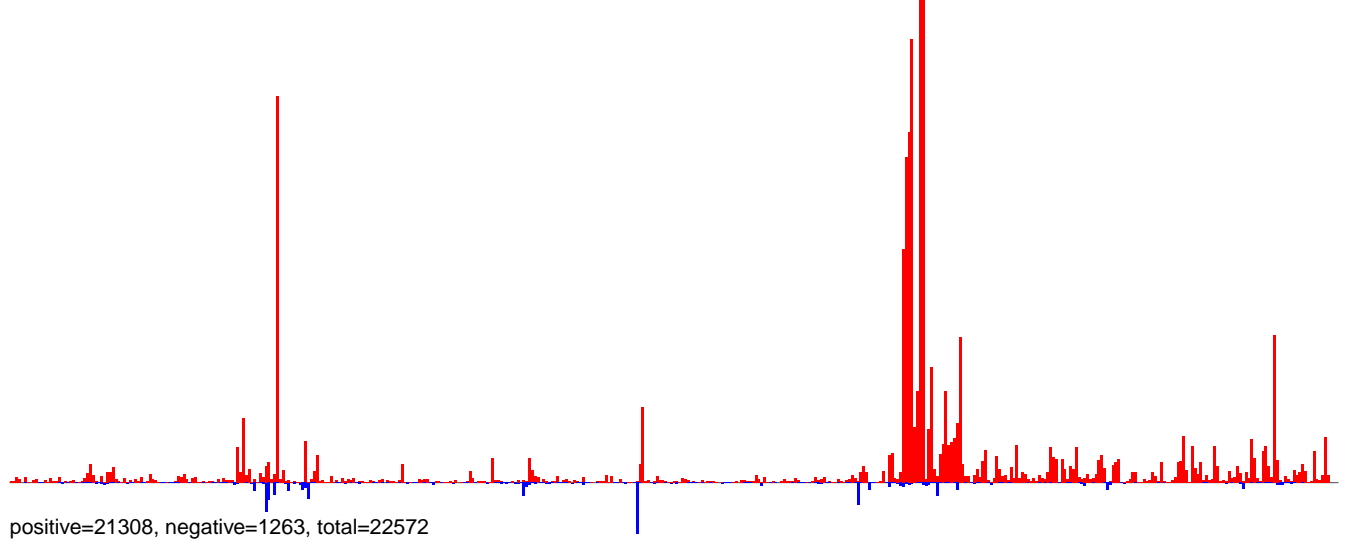


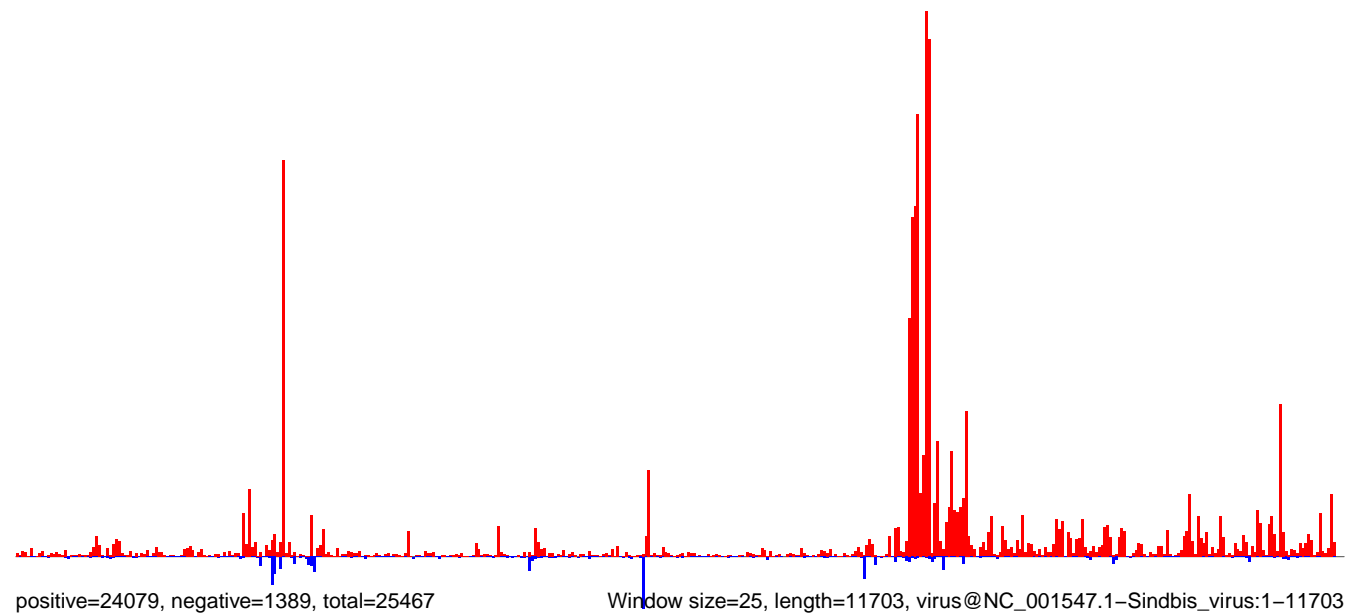
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



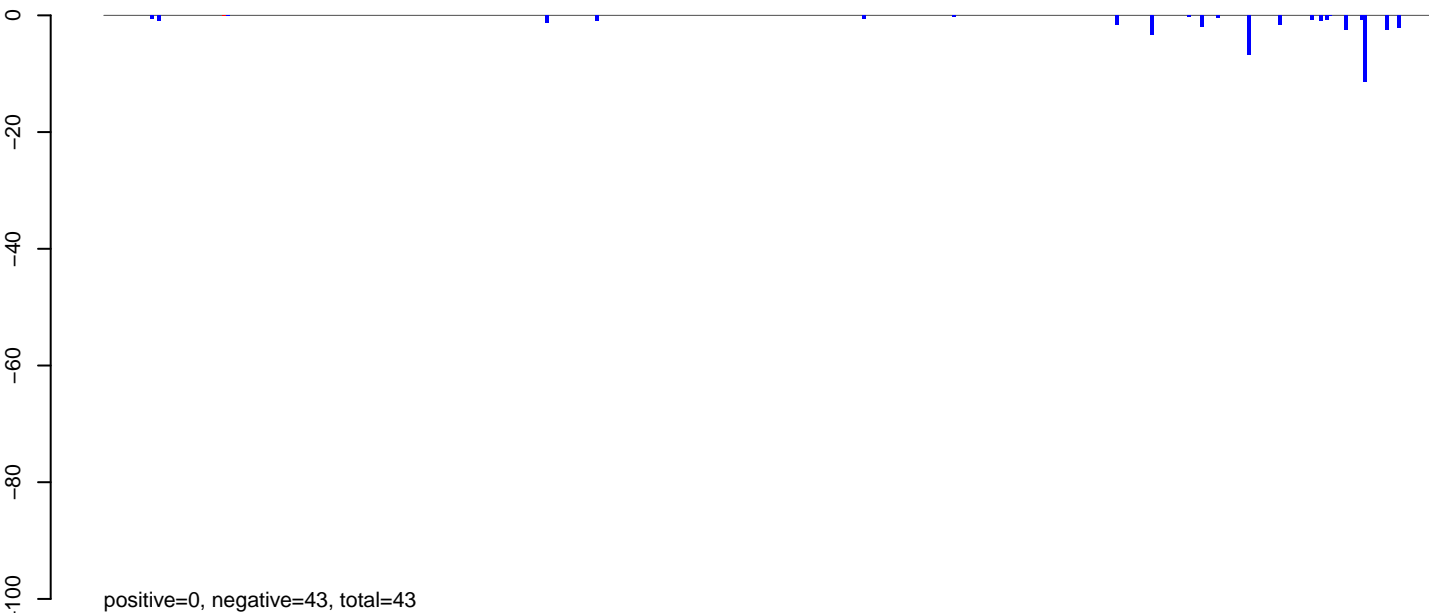
AeAlbo_C636_SINV_GFP_betaE.24_35.rep



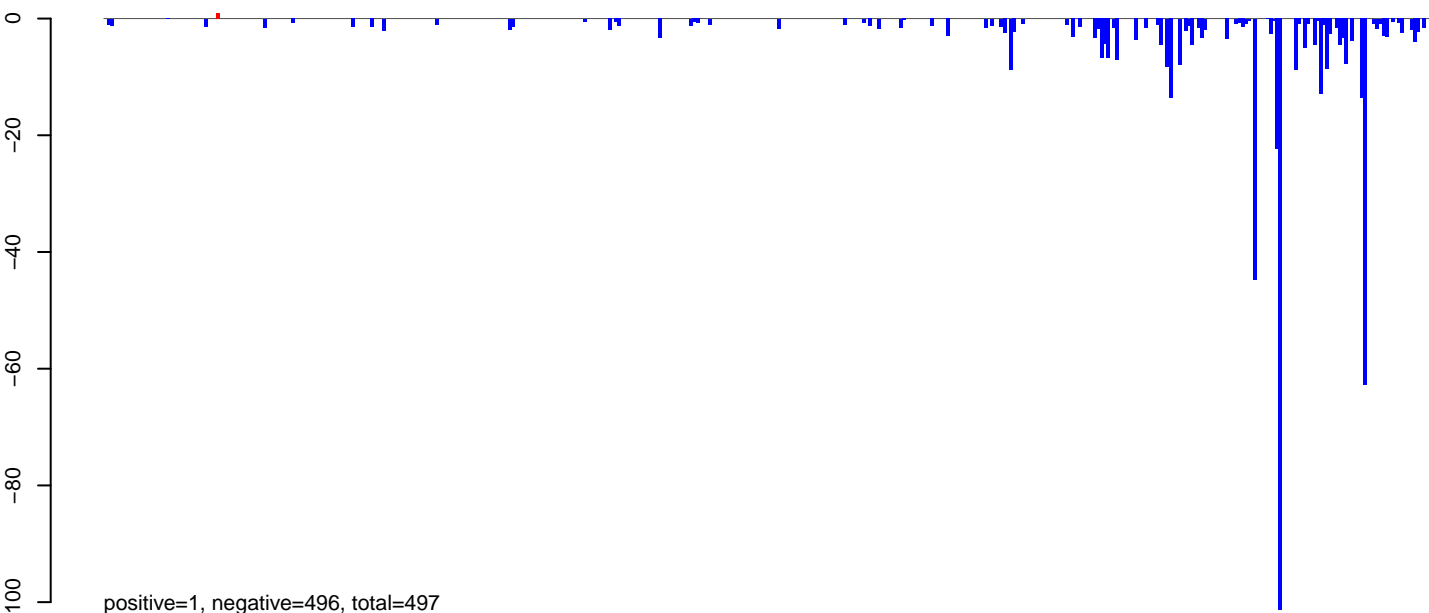
AeAlbo_C636_SINV_GFP_betaE.rep



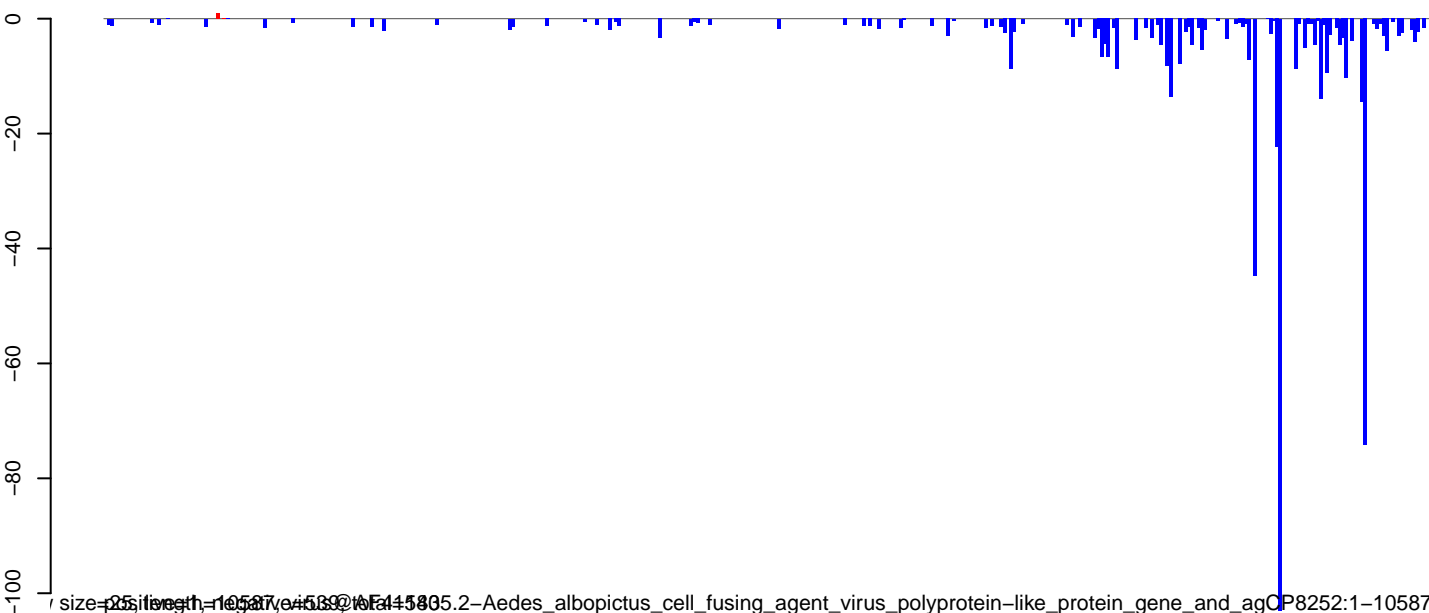
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep

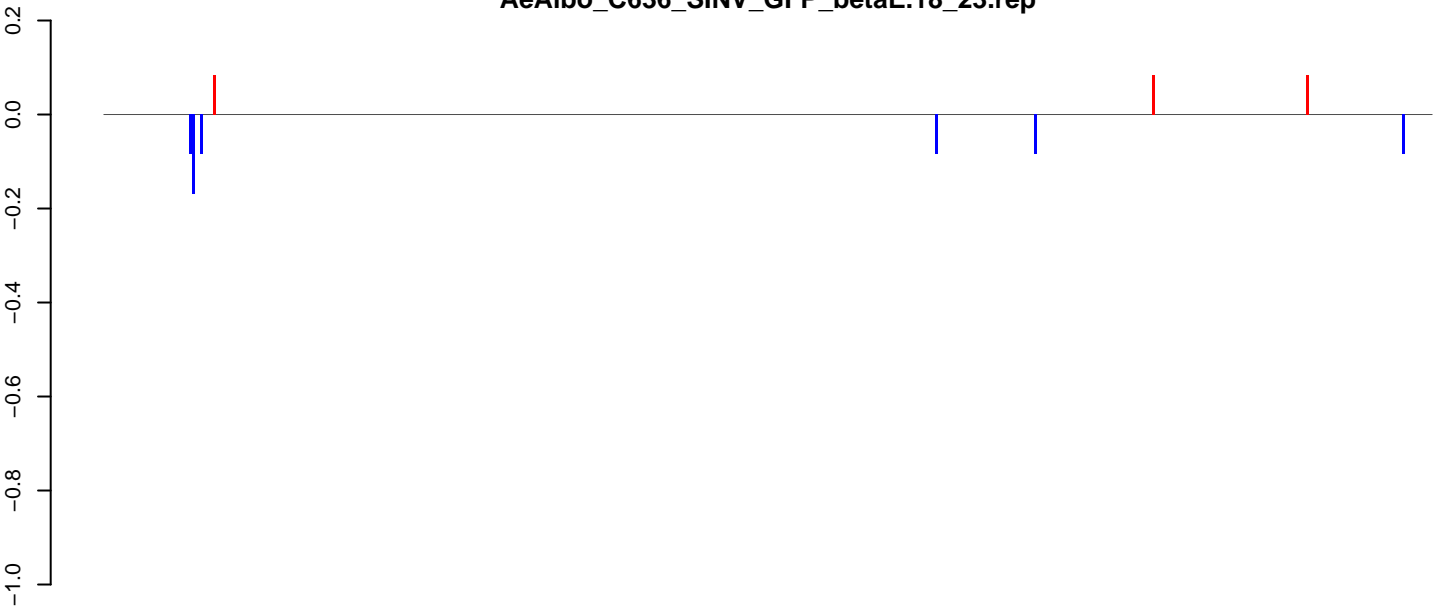


AeAlbo_C636_SINV_GFP_betaE.rep



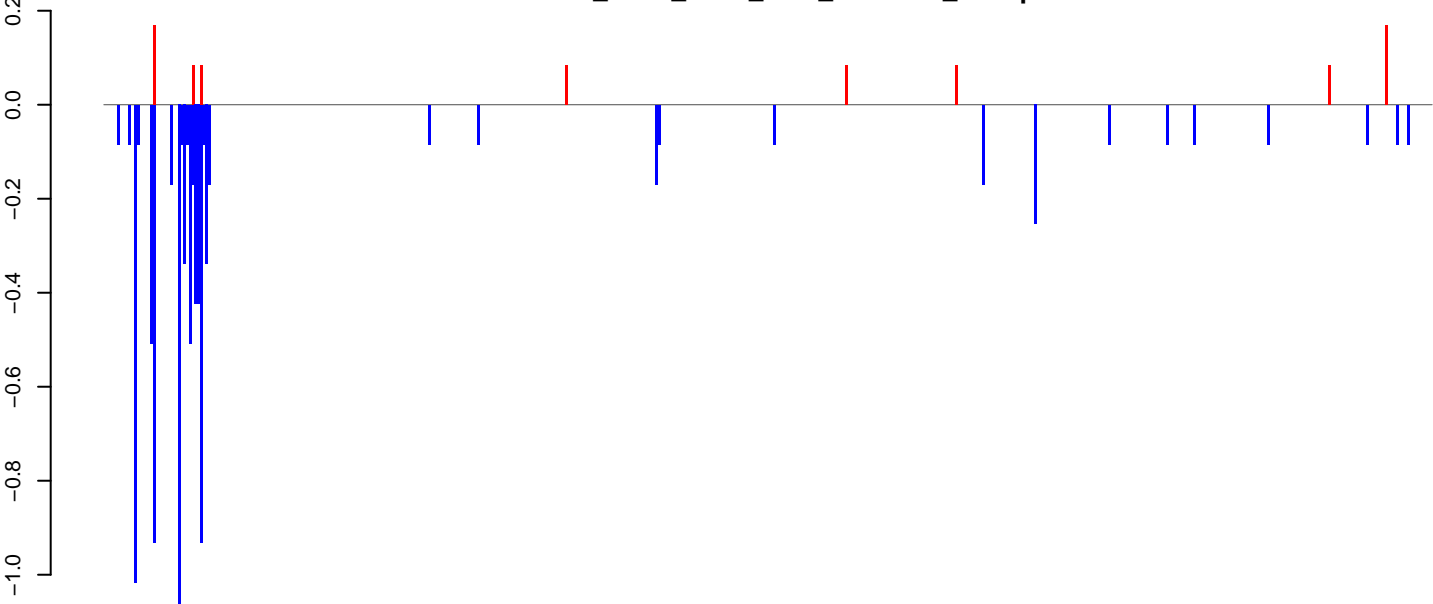
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AeAlbo_C636_SINV_GFP_betaE.18_23.rep



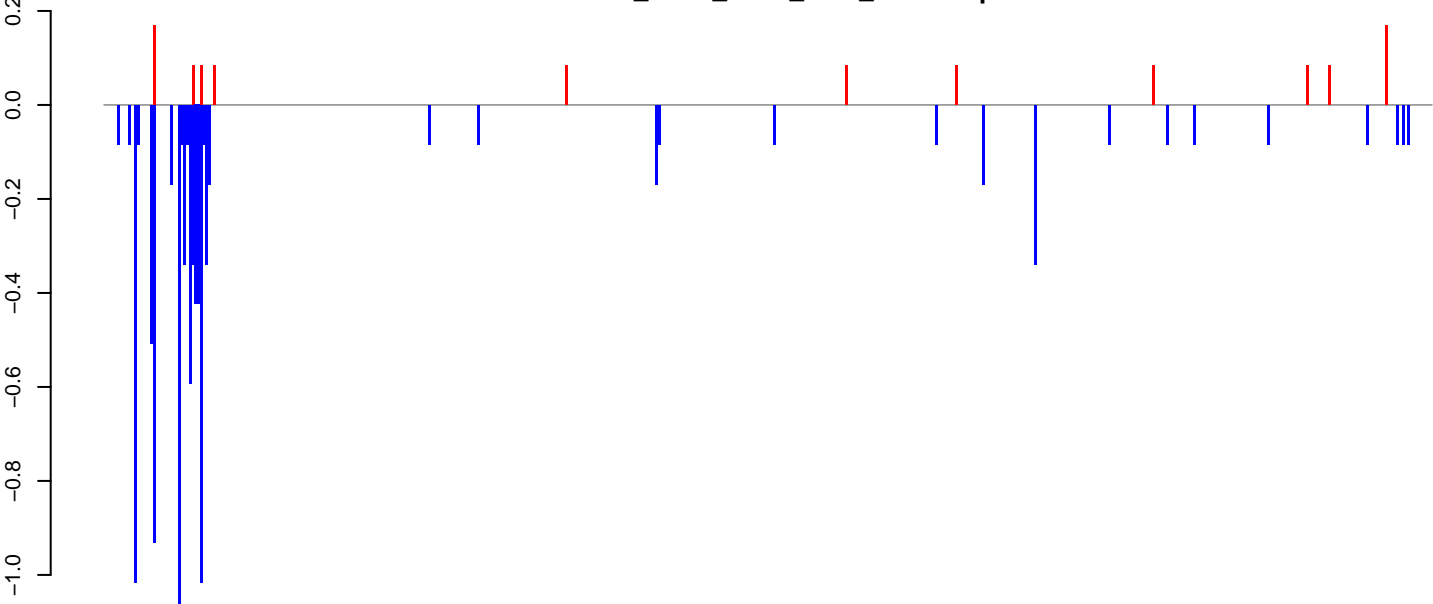
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AeAlbo_C636_SINV_GFP_betaE.24_35.rep



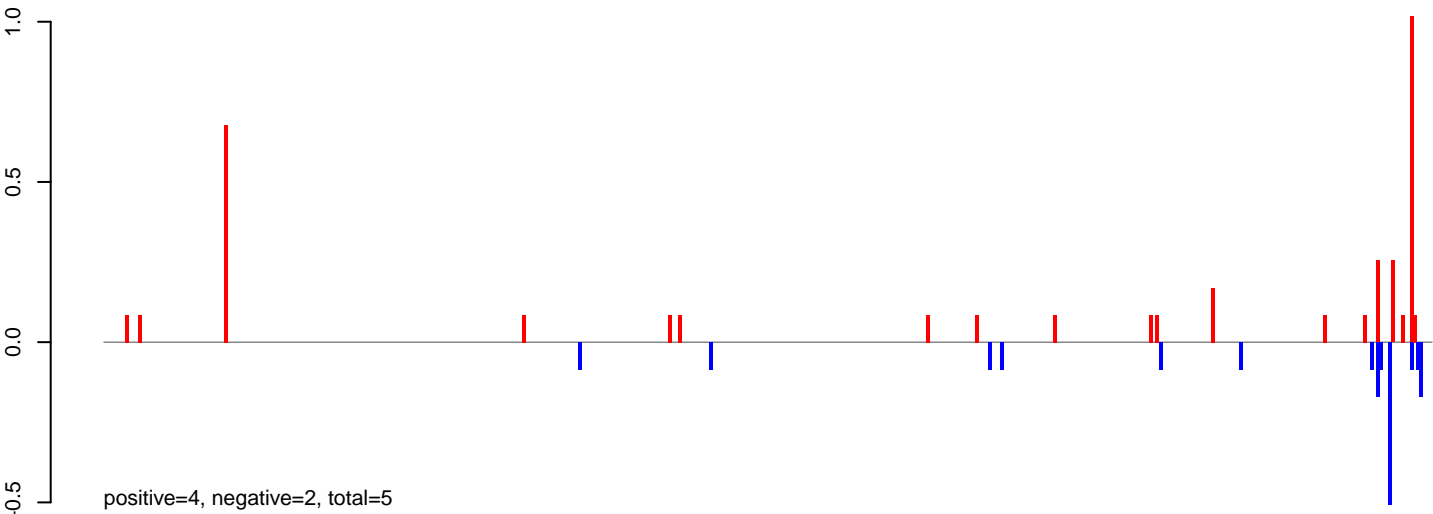
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AeAlbo_C636_SINV_GFP_betaE.rep

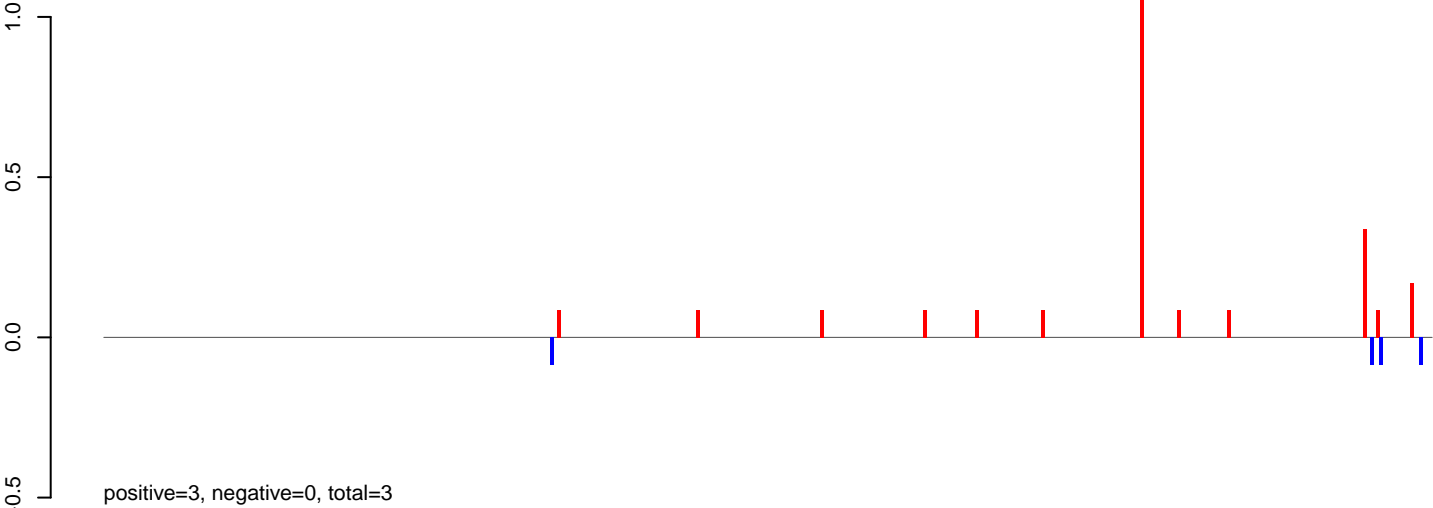


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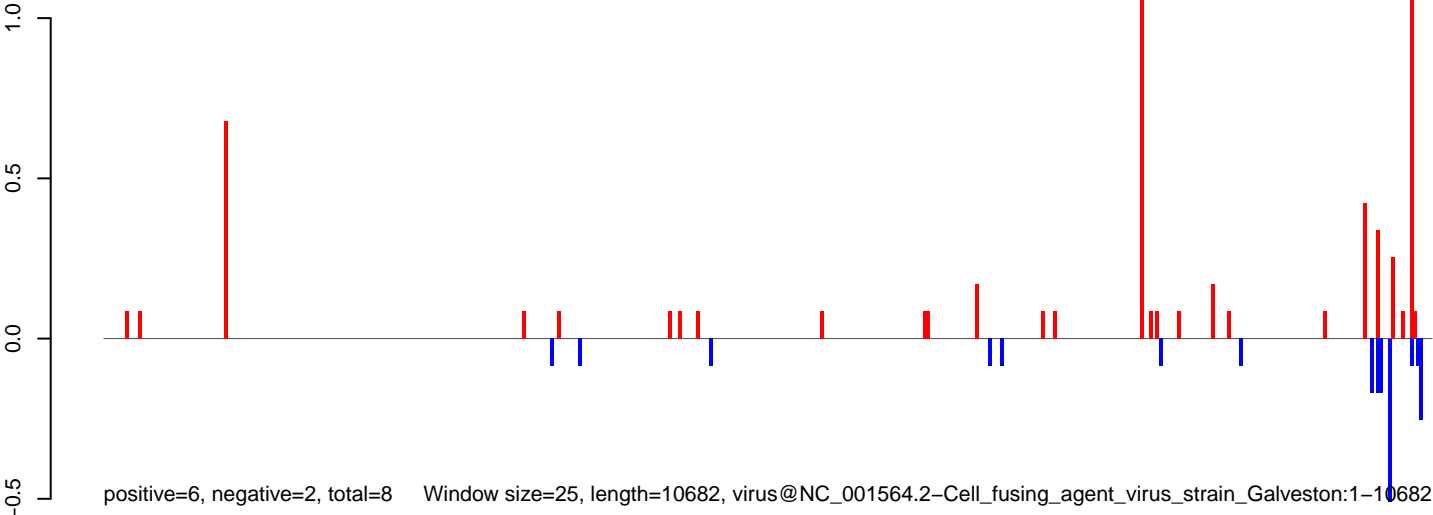
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep

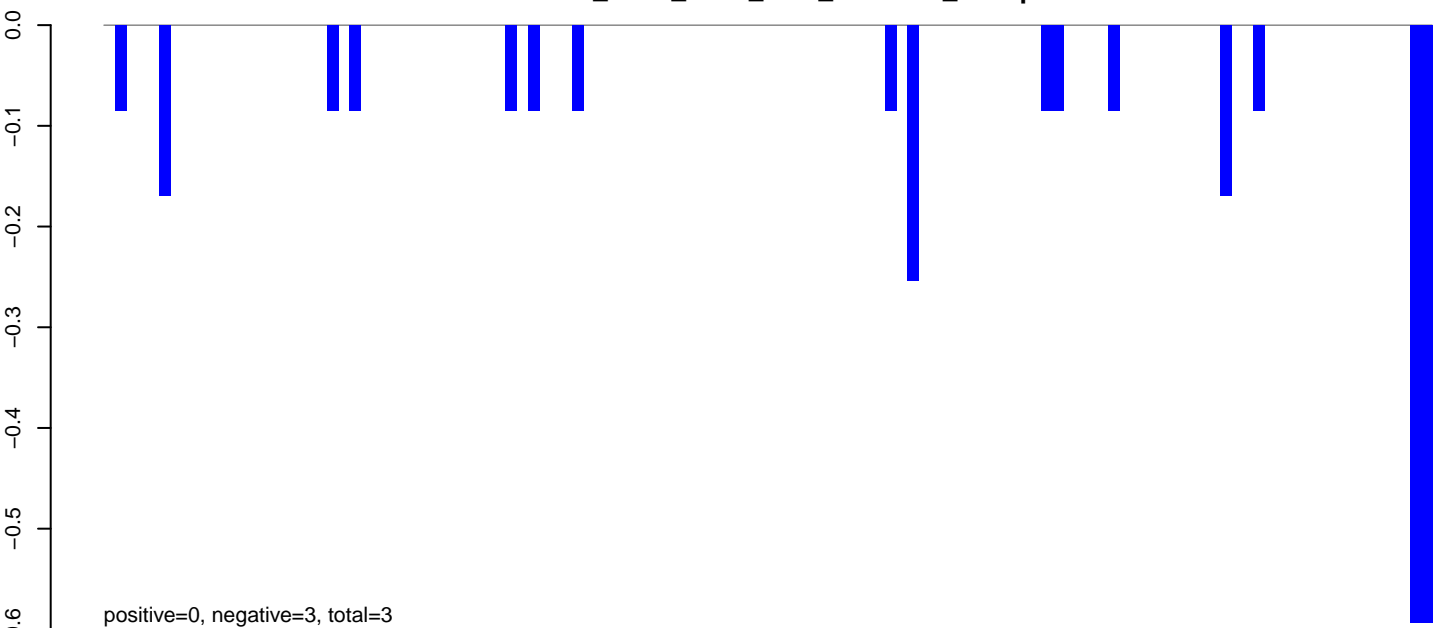


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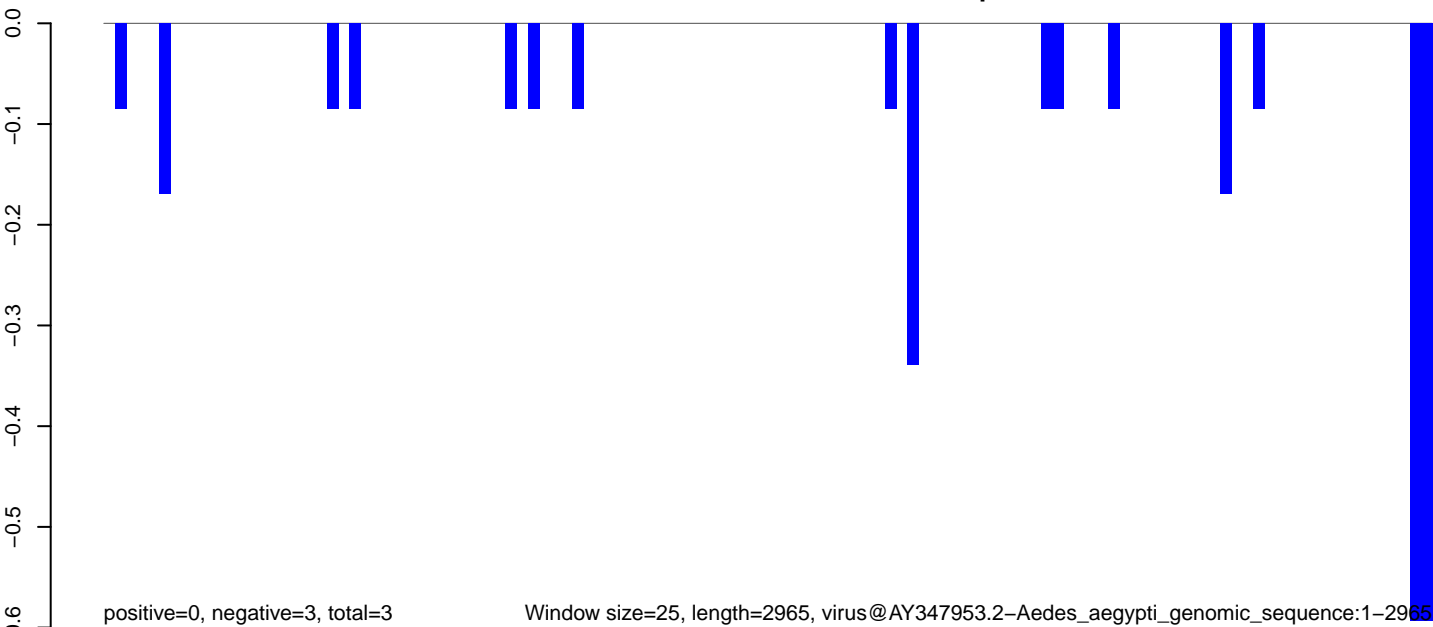
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep

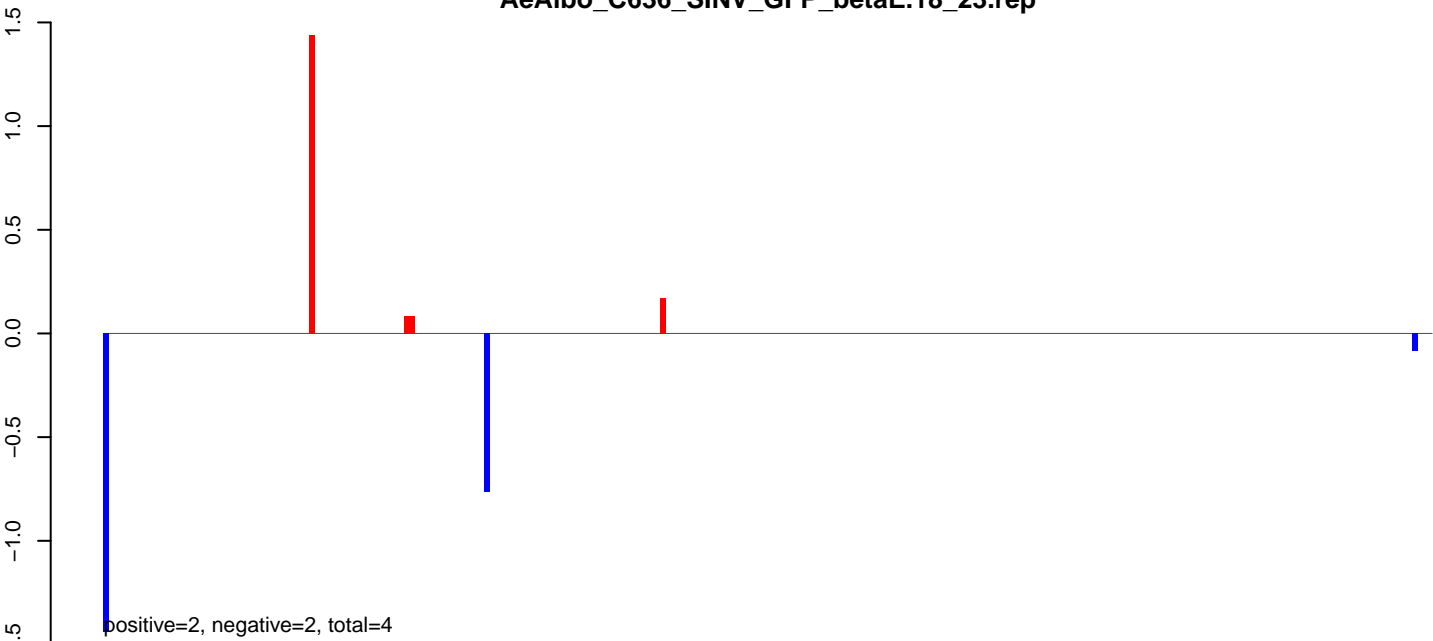


AeAlbo_C636_SINV_GFP_betaE.rep

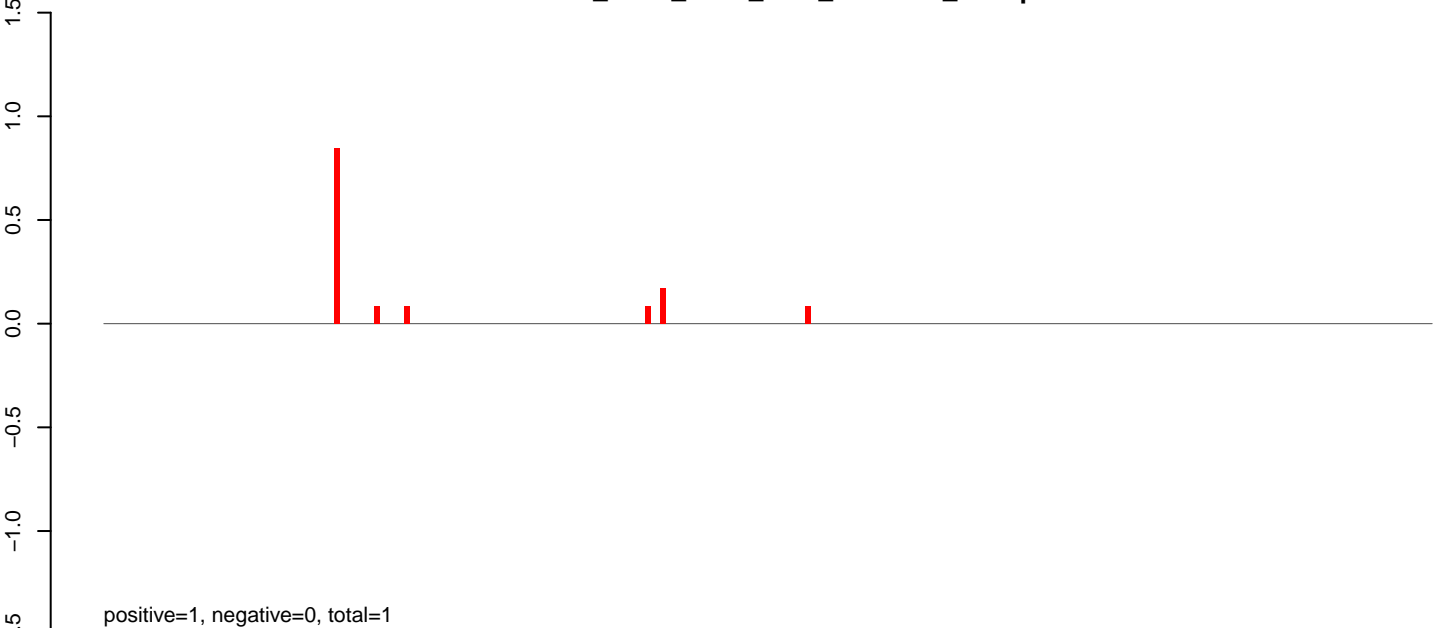


Window size=25, length=2965, virus@AY347953.2-Aedes_aegypti_genomic_sequence:1-2965

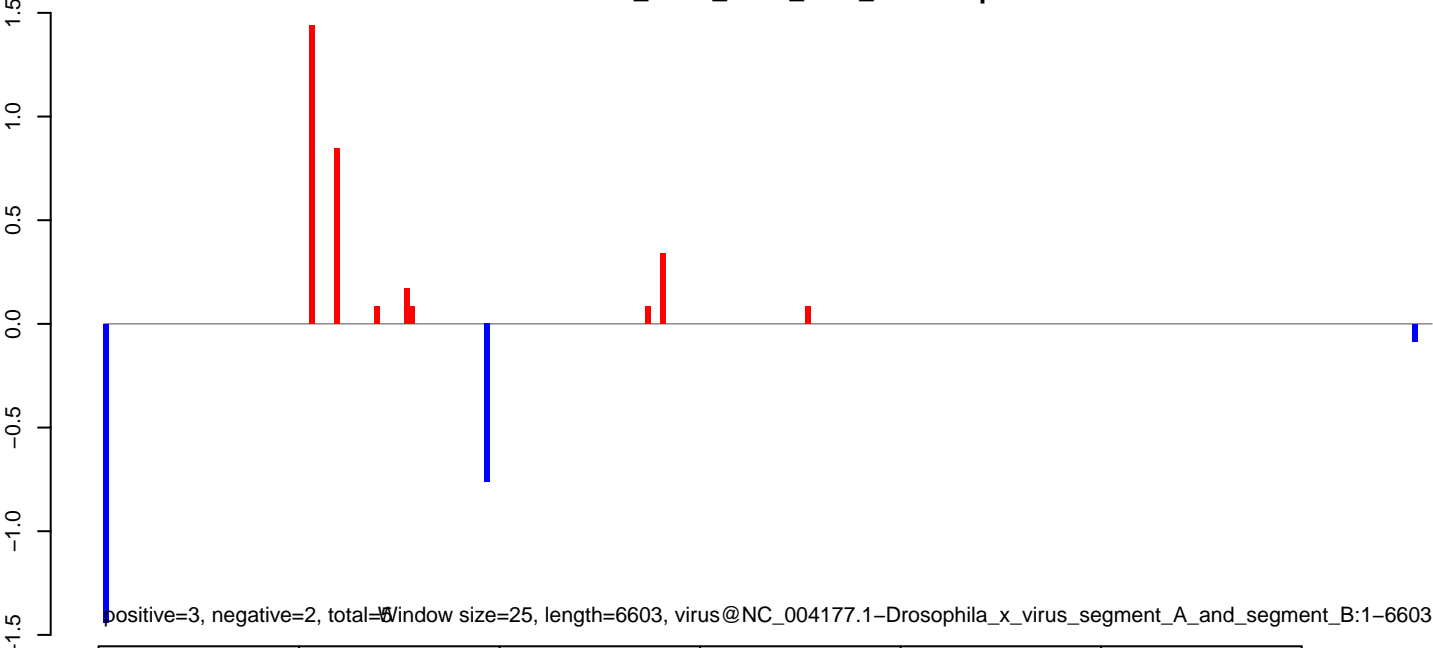
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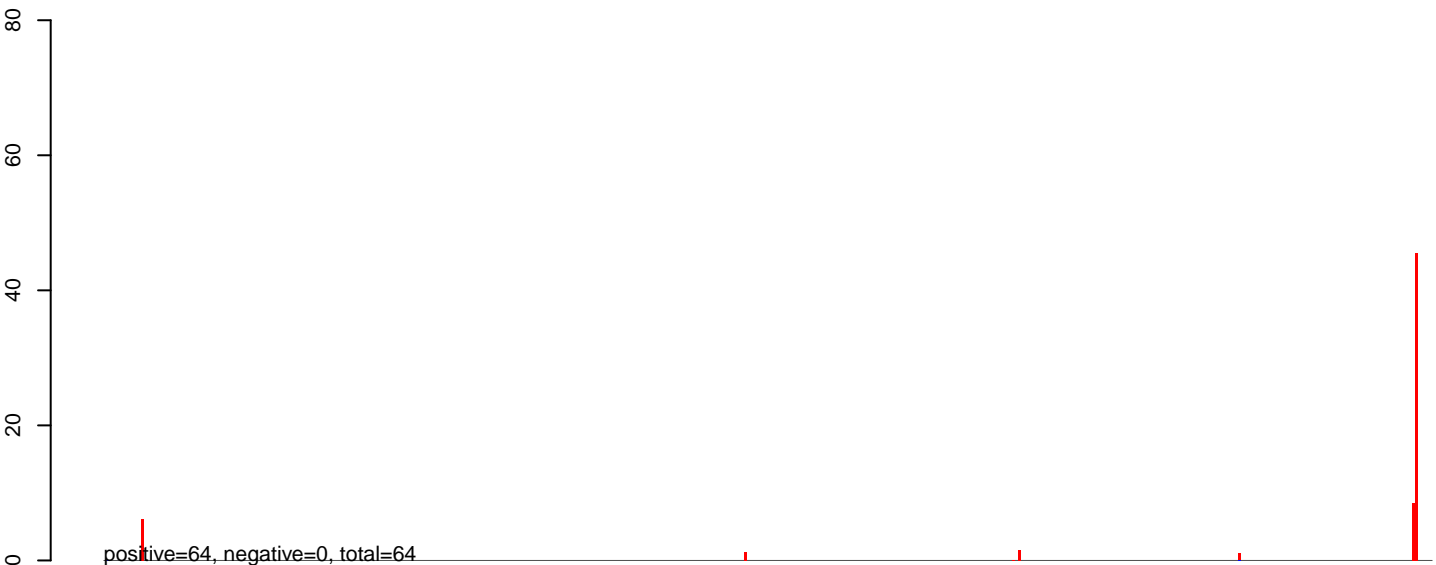
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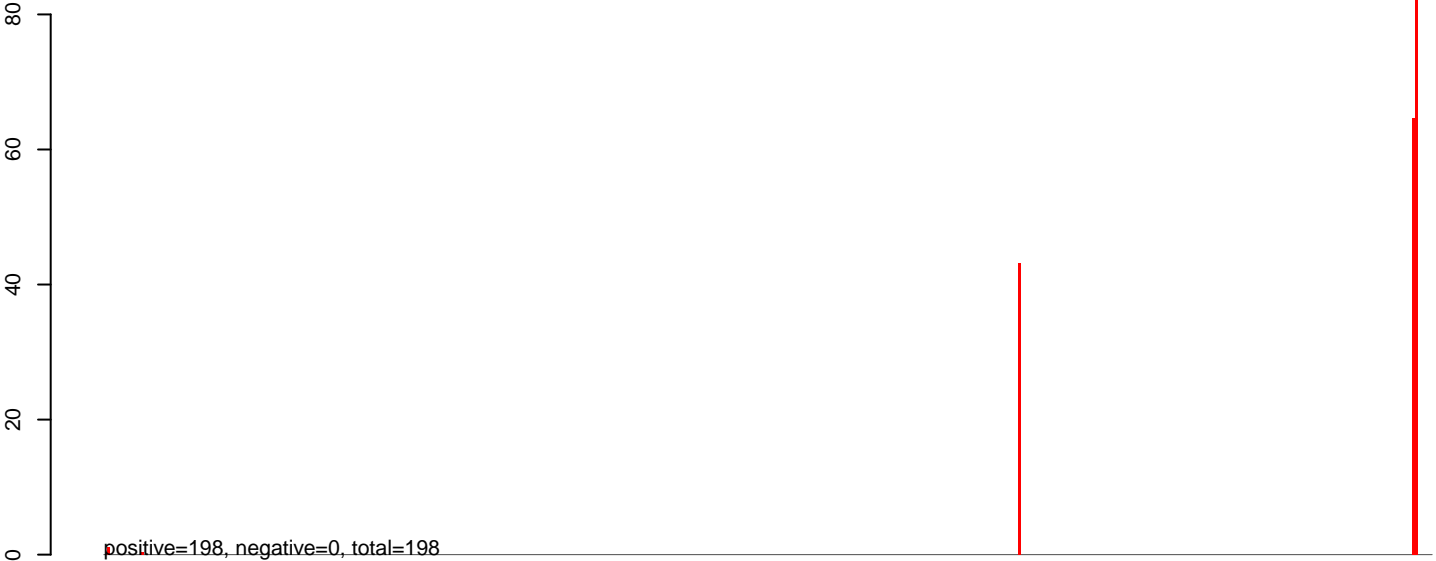
AeAlbo_C636_SINV_GFP_betaE.rep



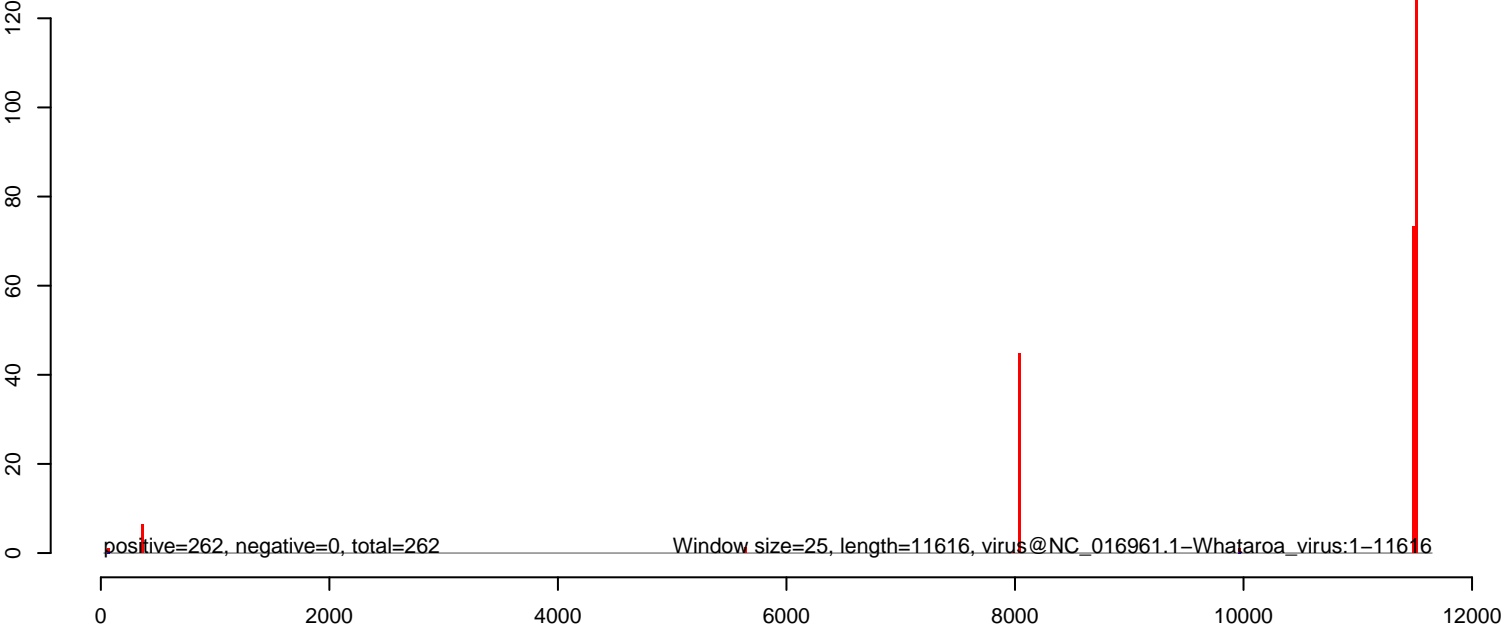
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



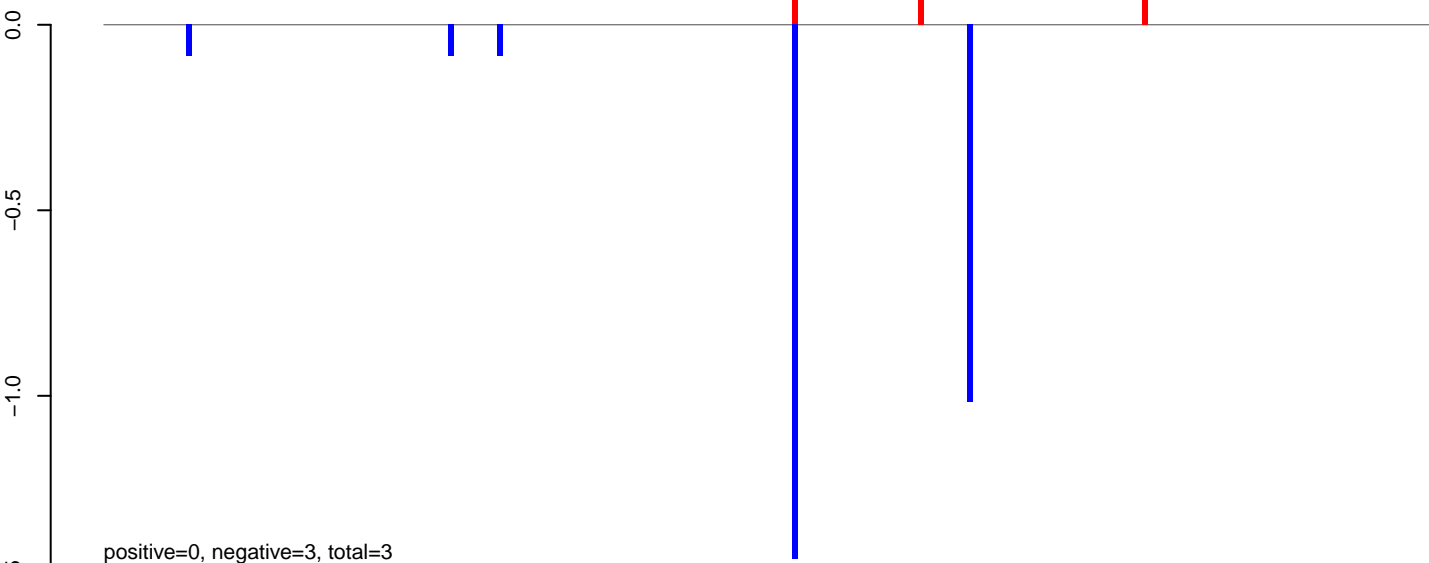
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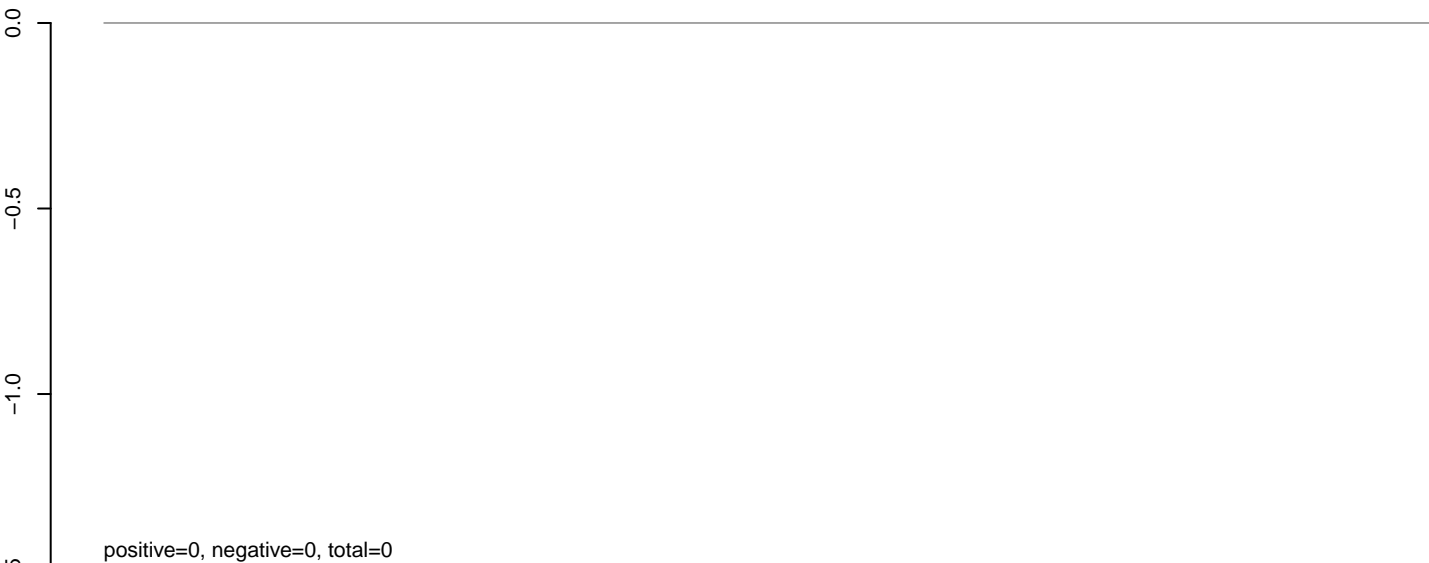
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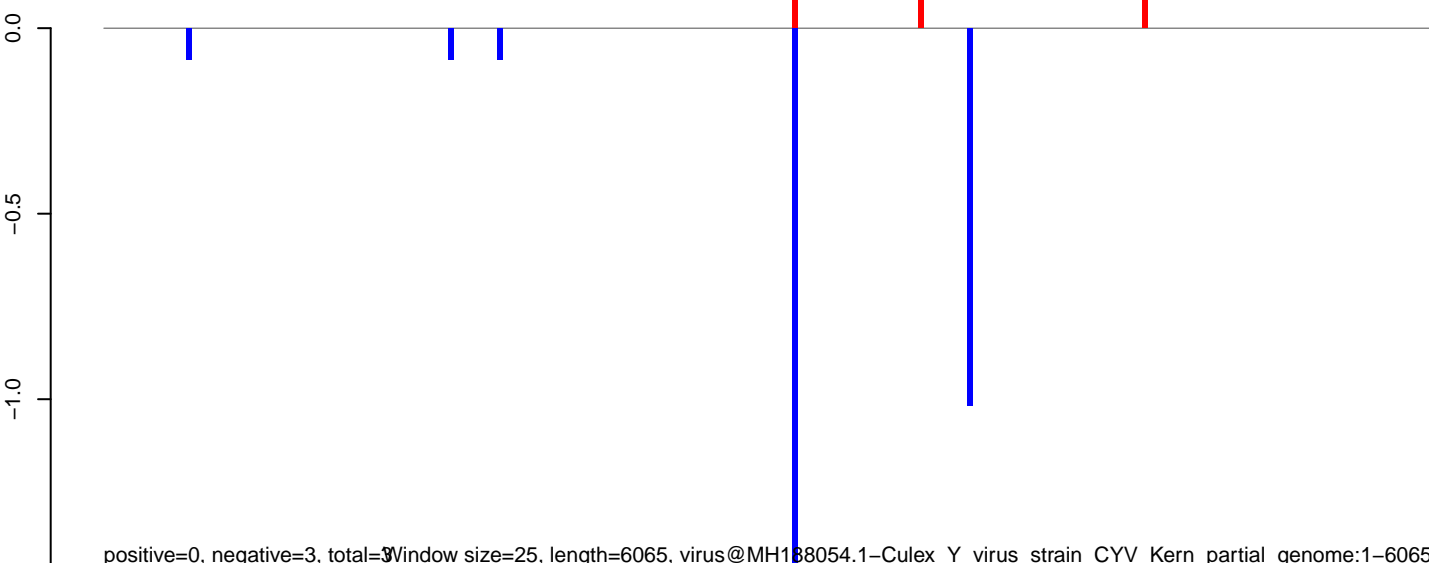
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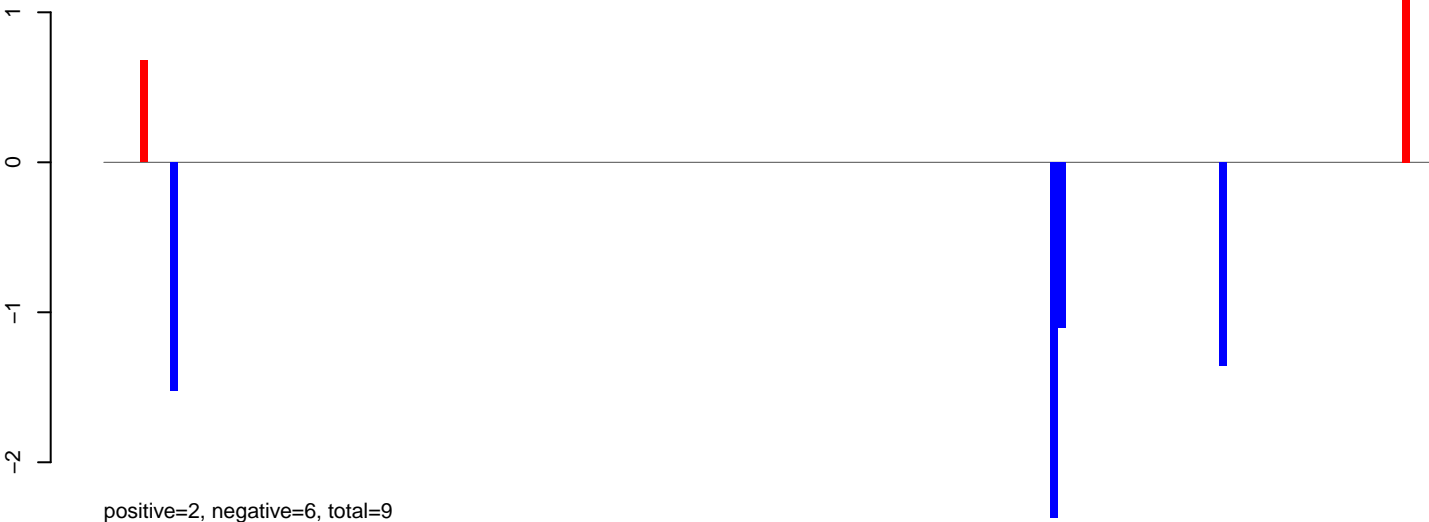
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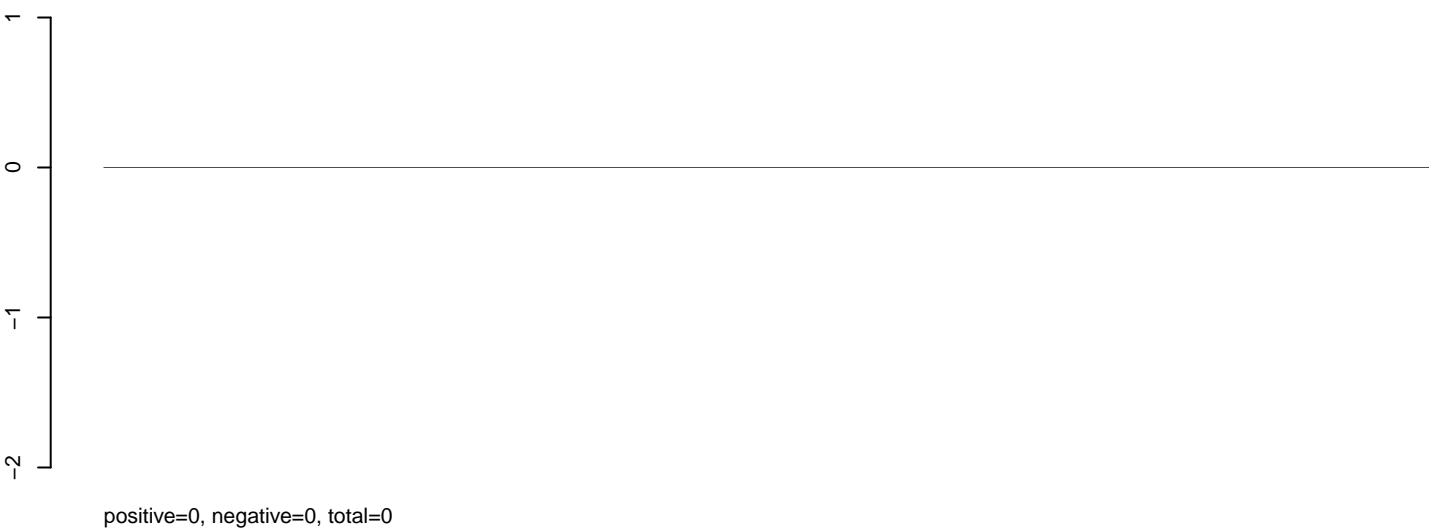
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AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



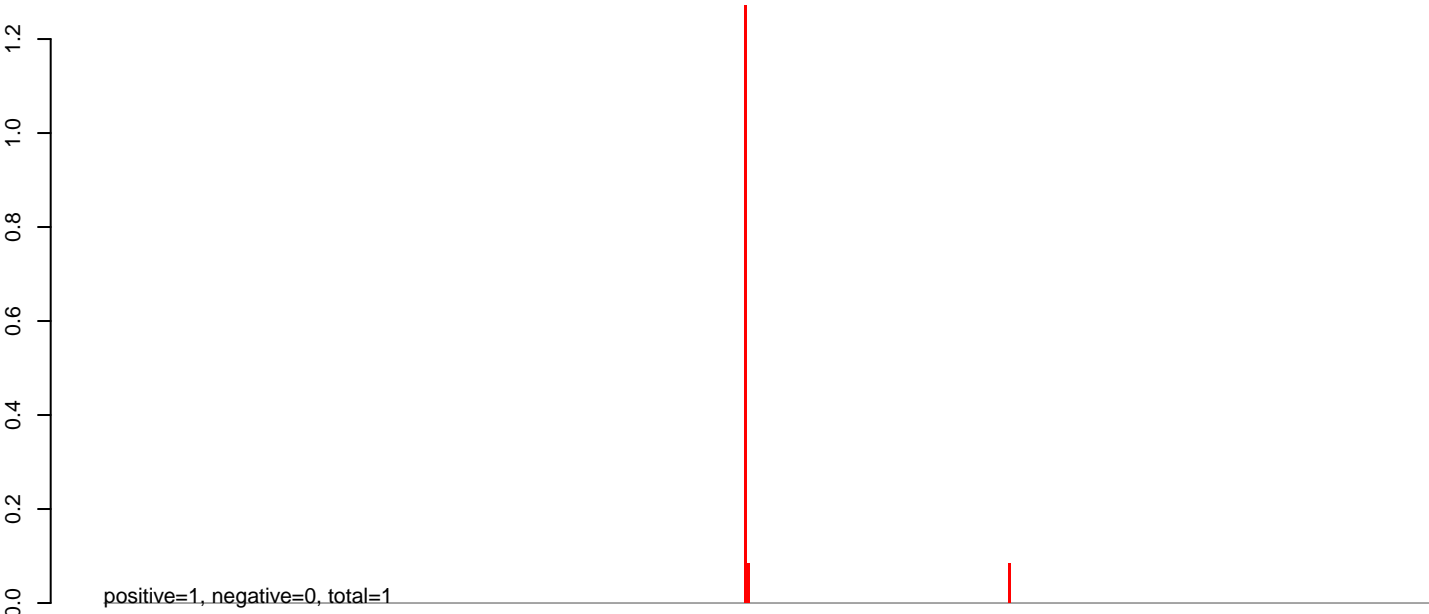
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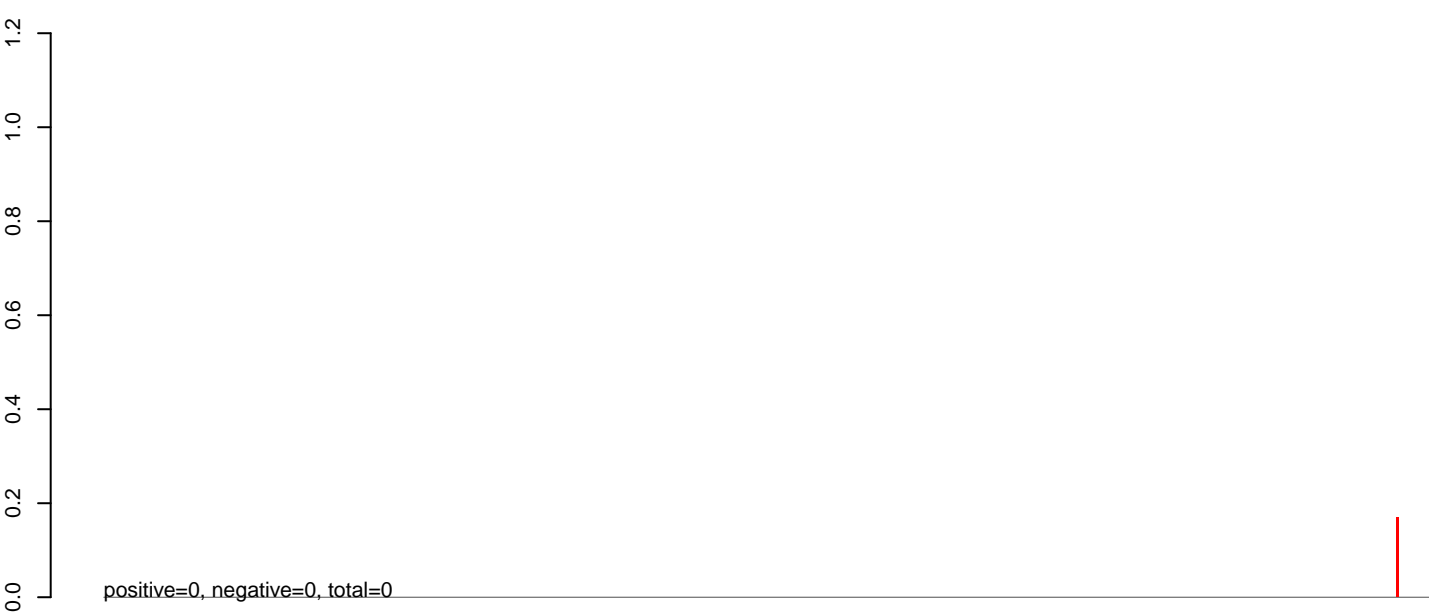
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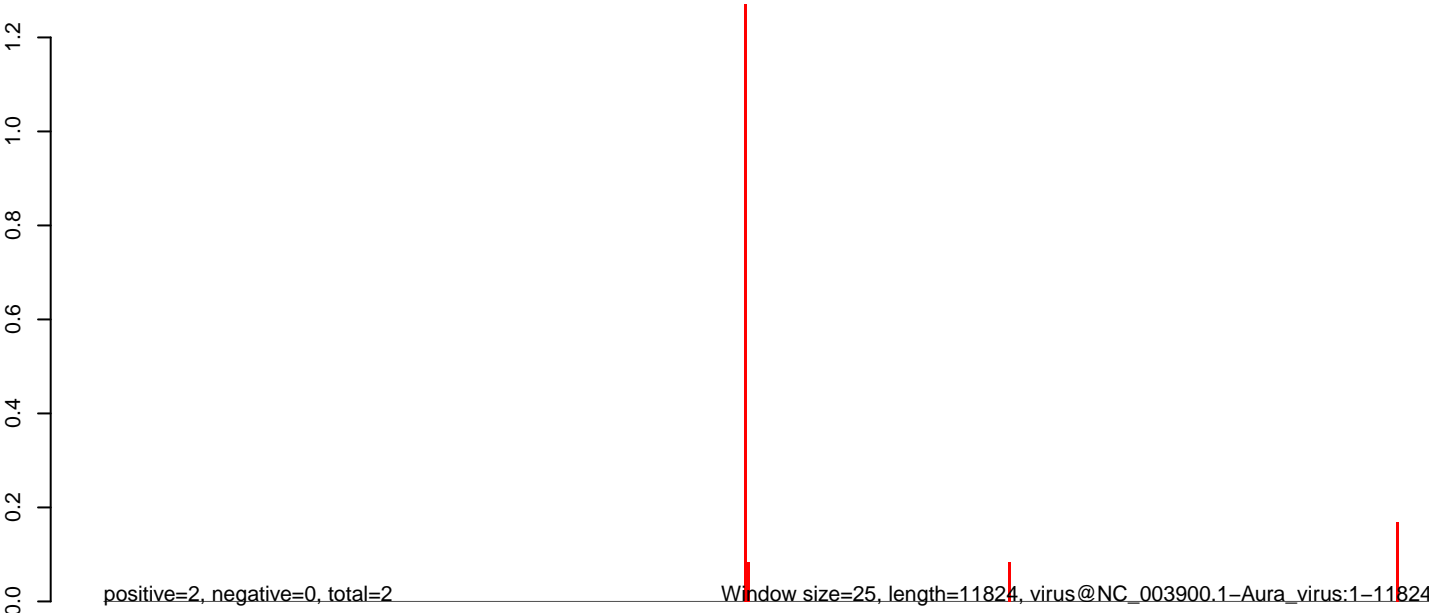
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AeAlbo_C636_SINV_GFP_betaE.24_35.rep



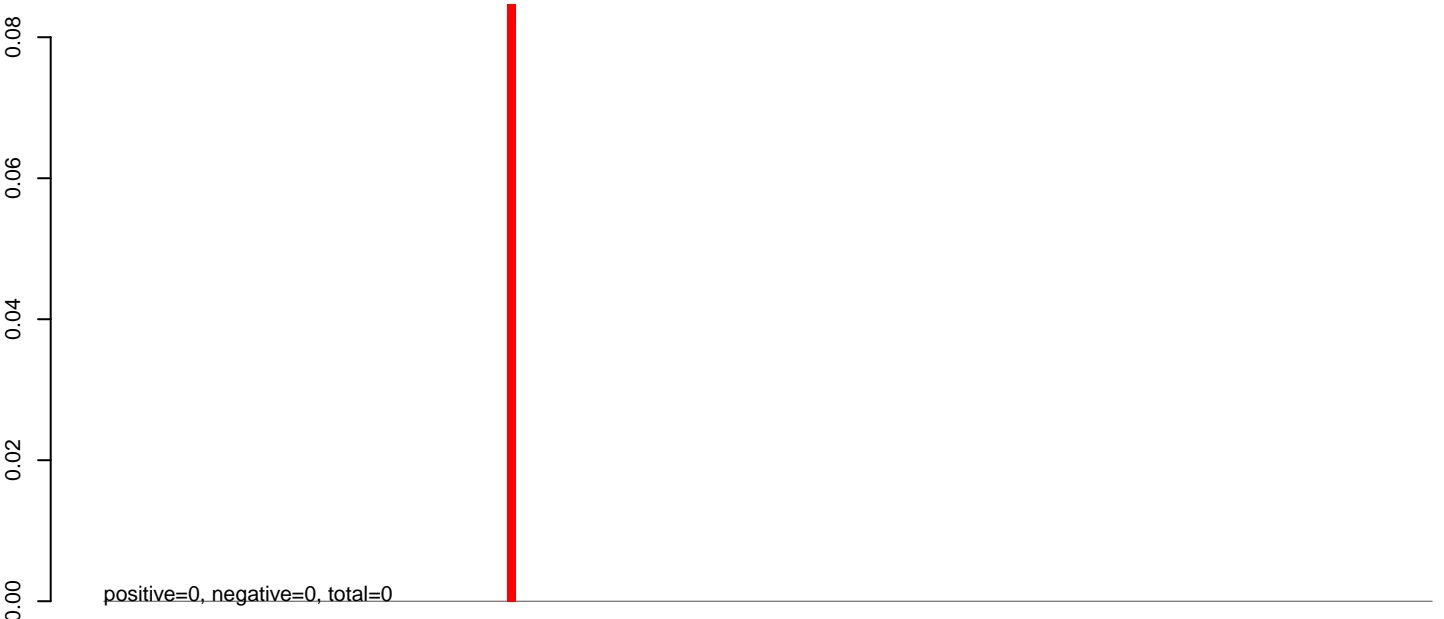
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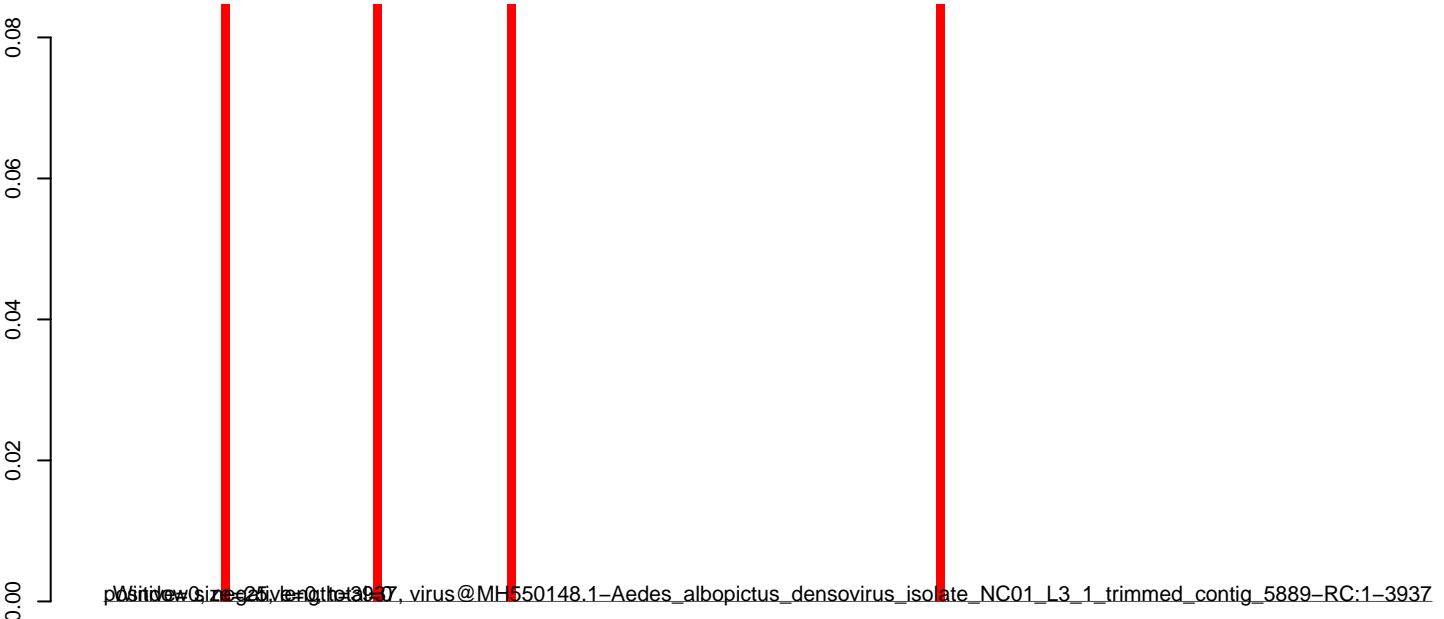
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep

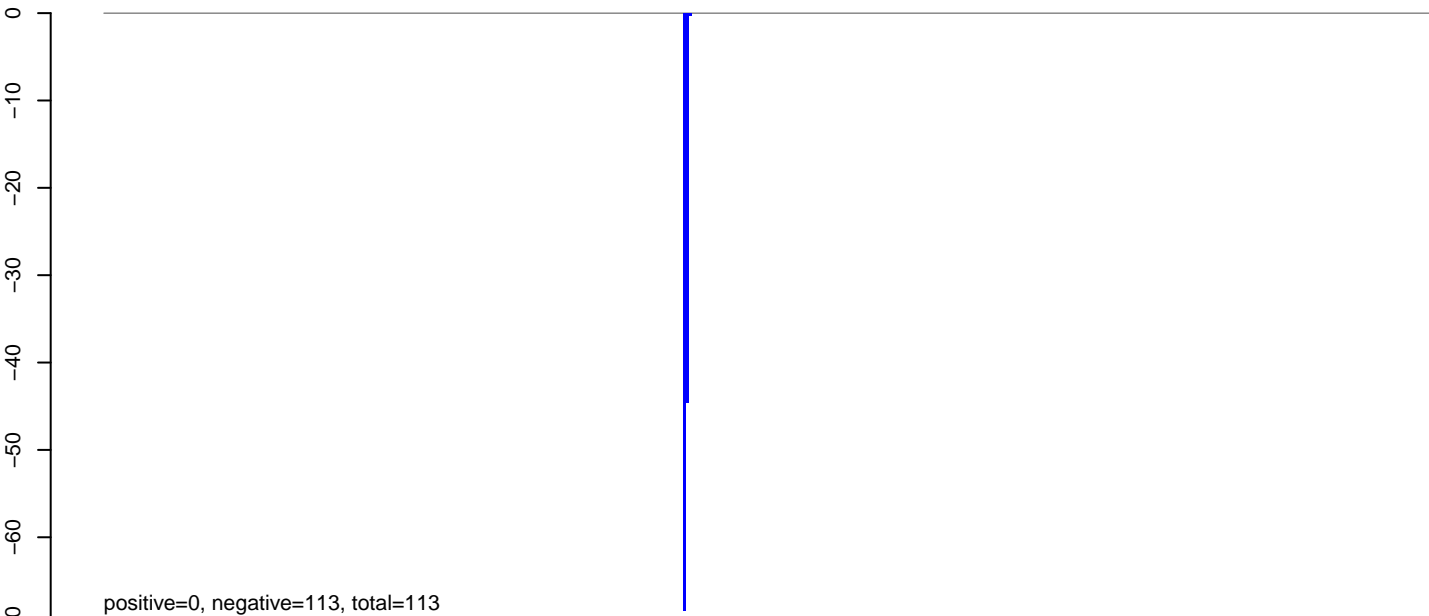


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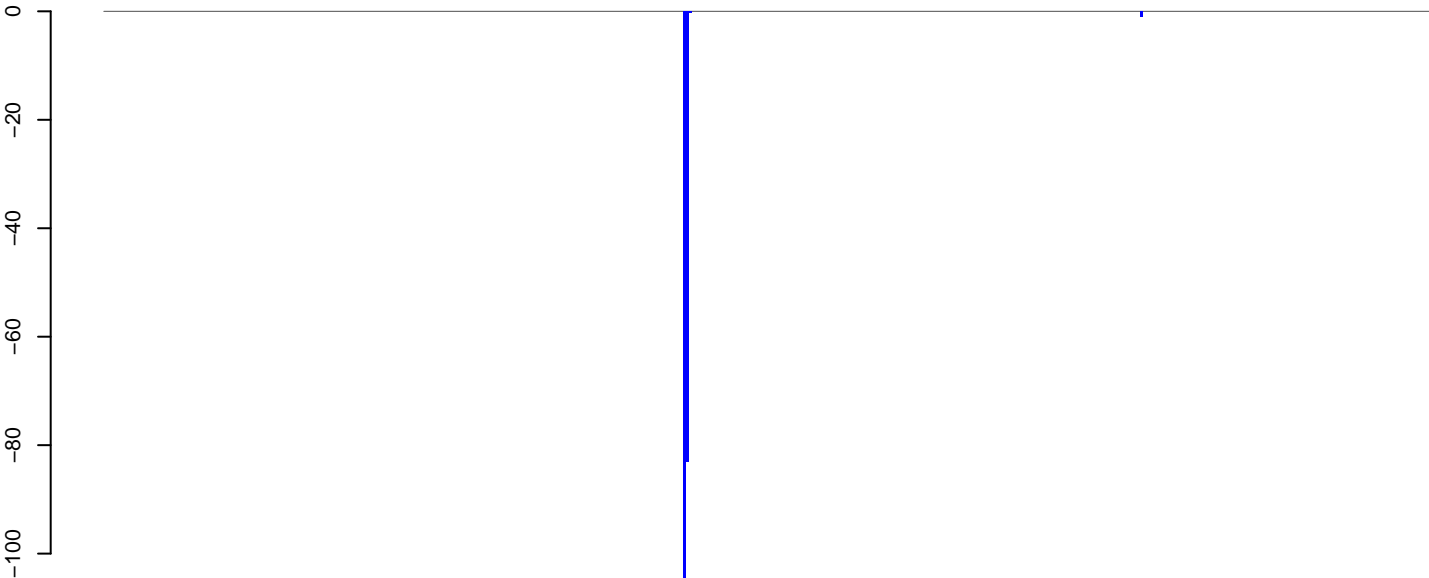
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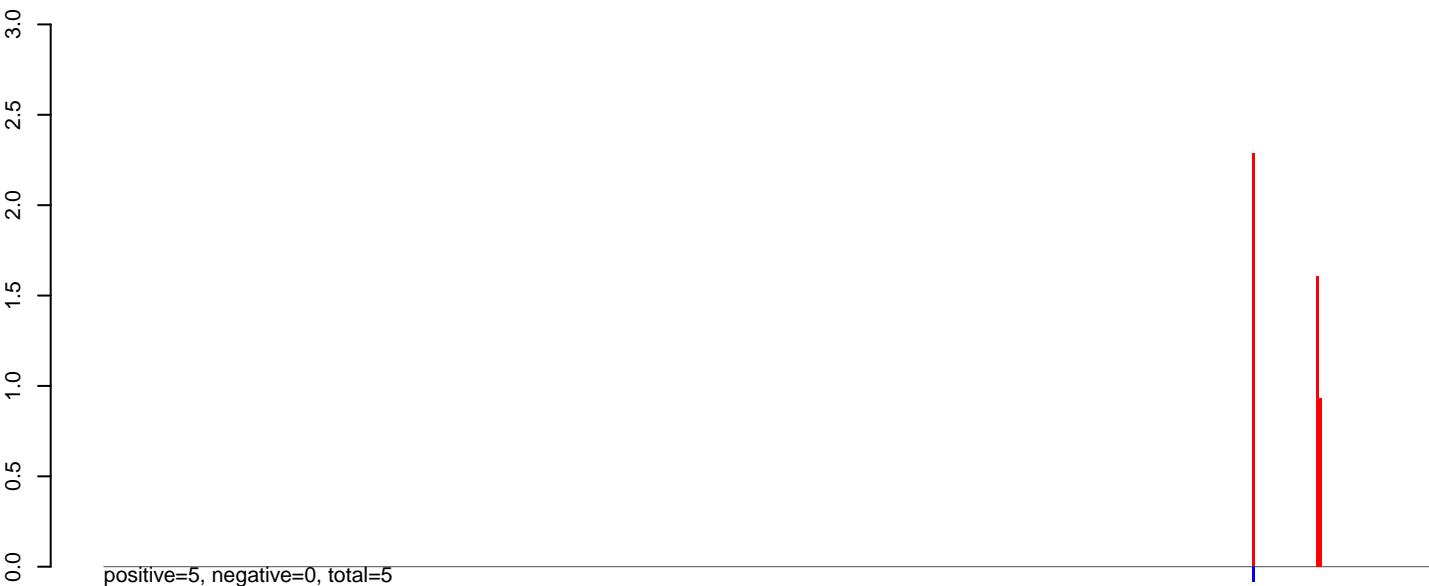
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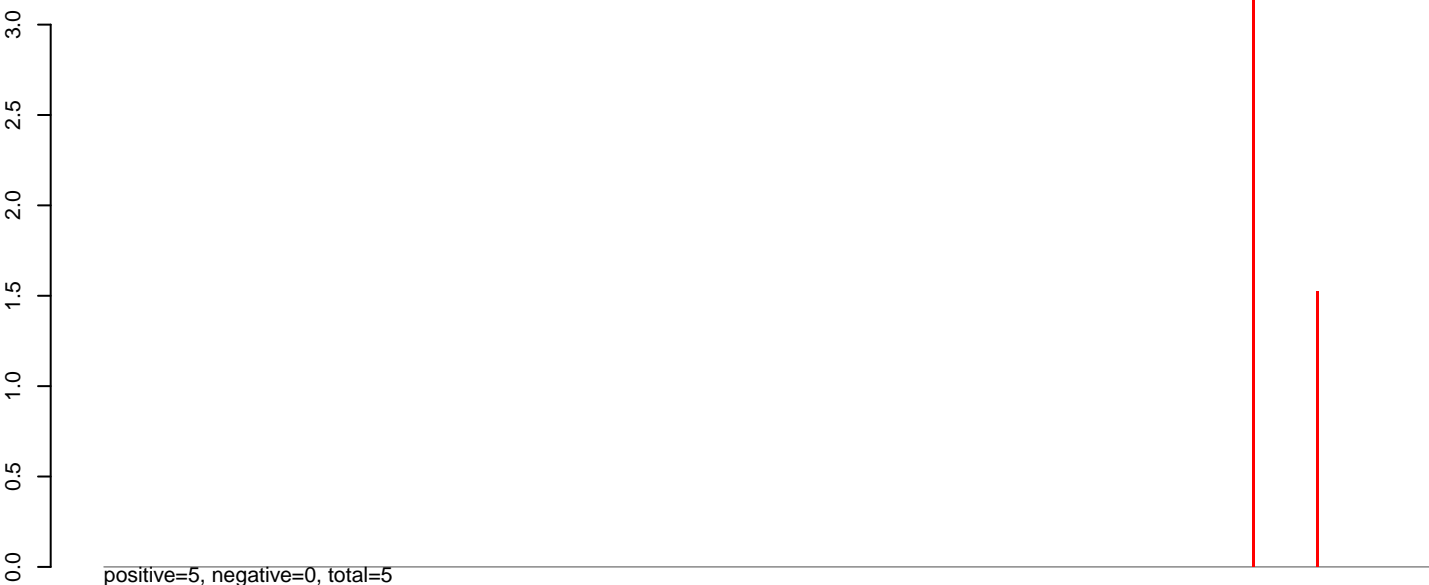
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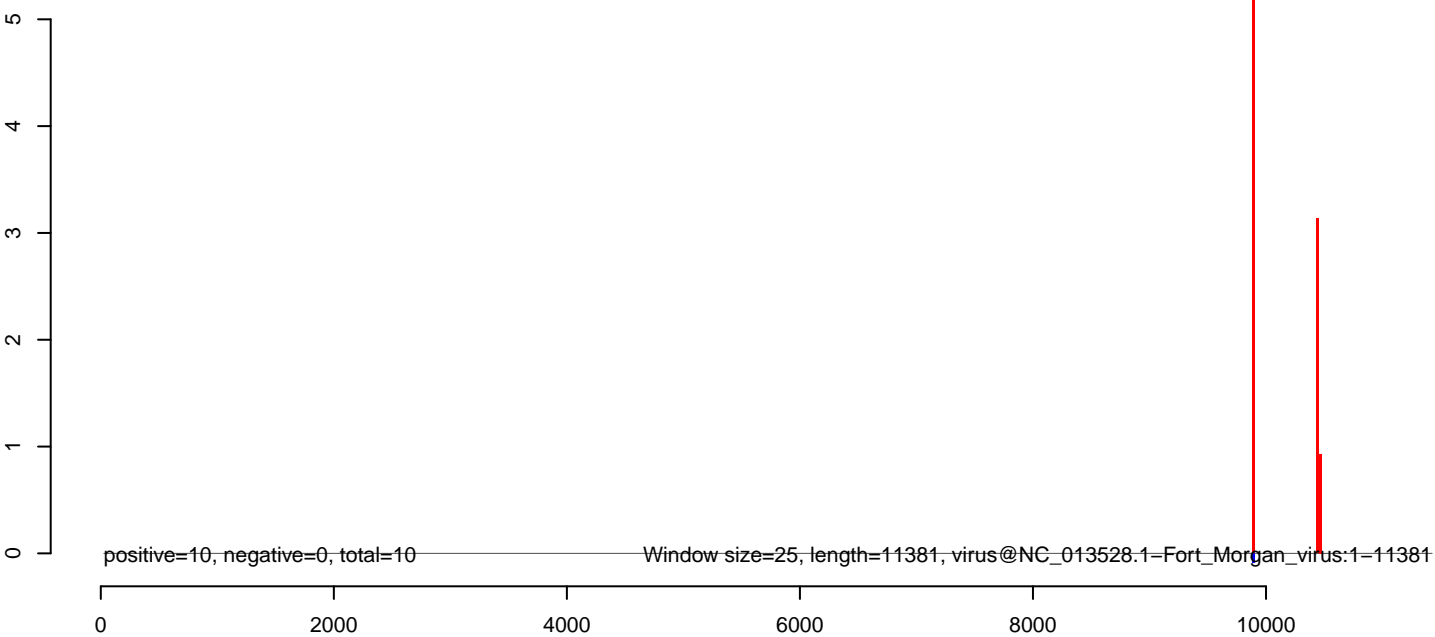
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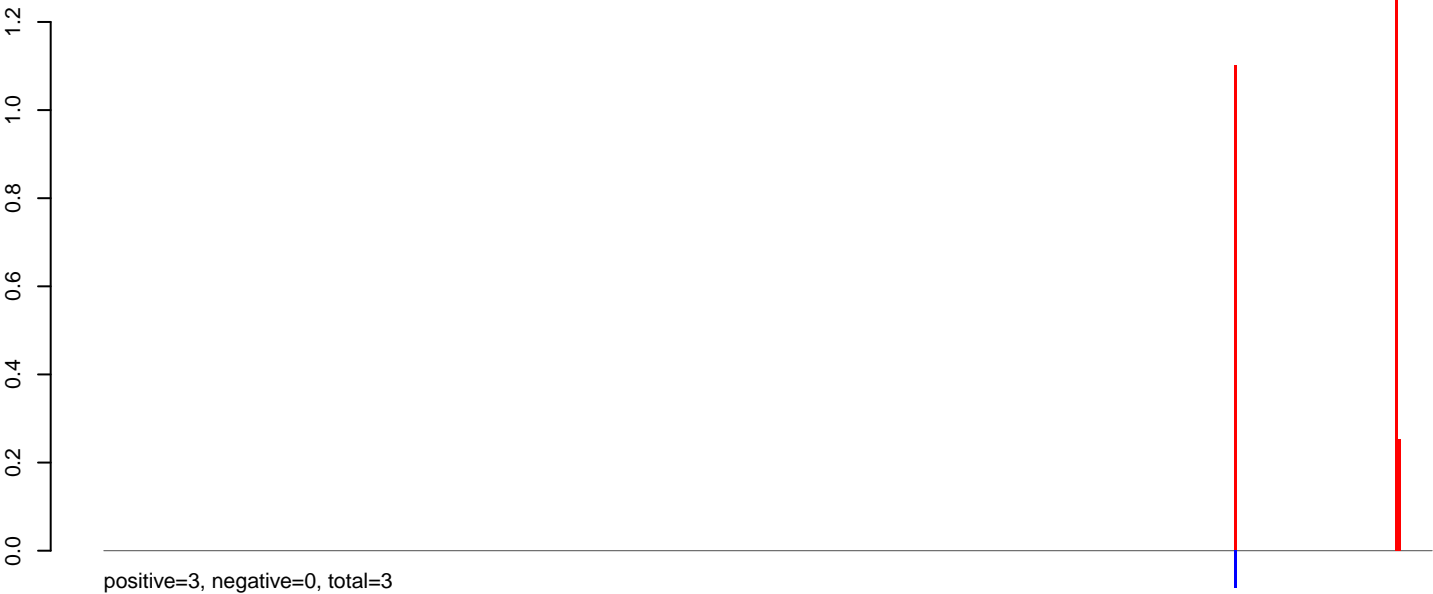
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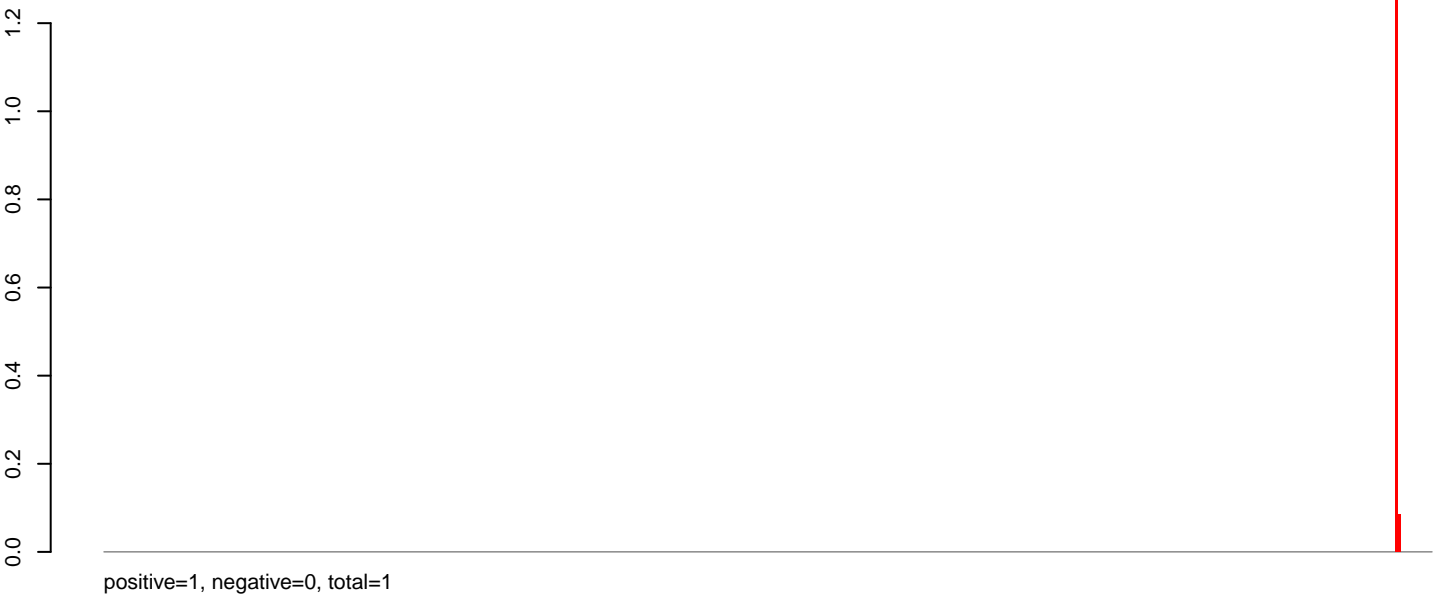
AeAlbo_C636_SINV_GFP_betaE.rep



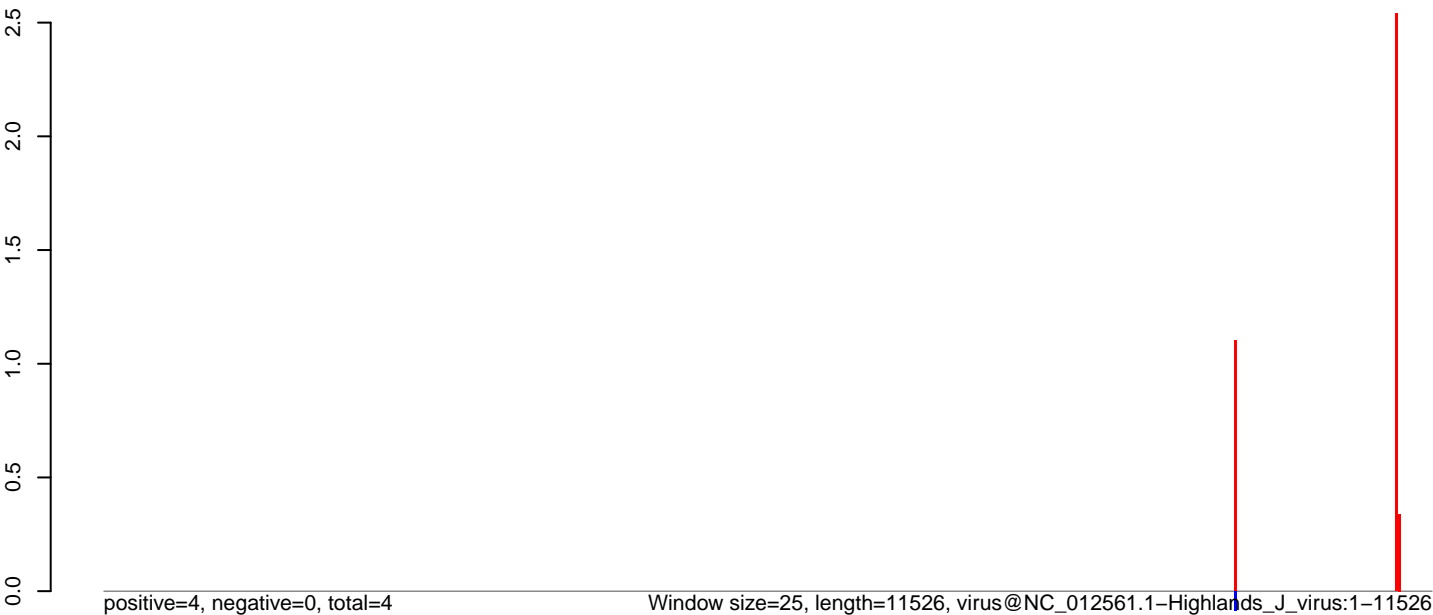
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



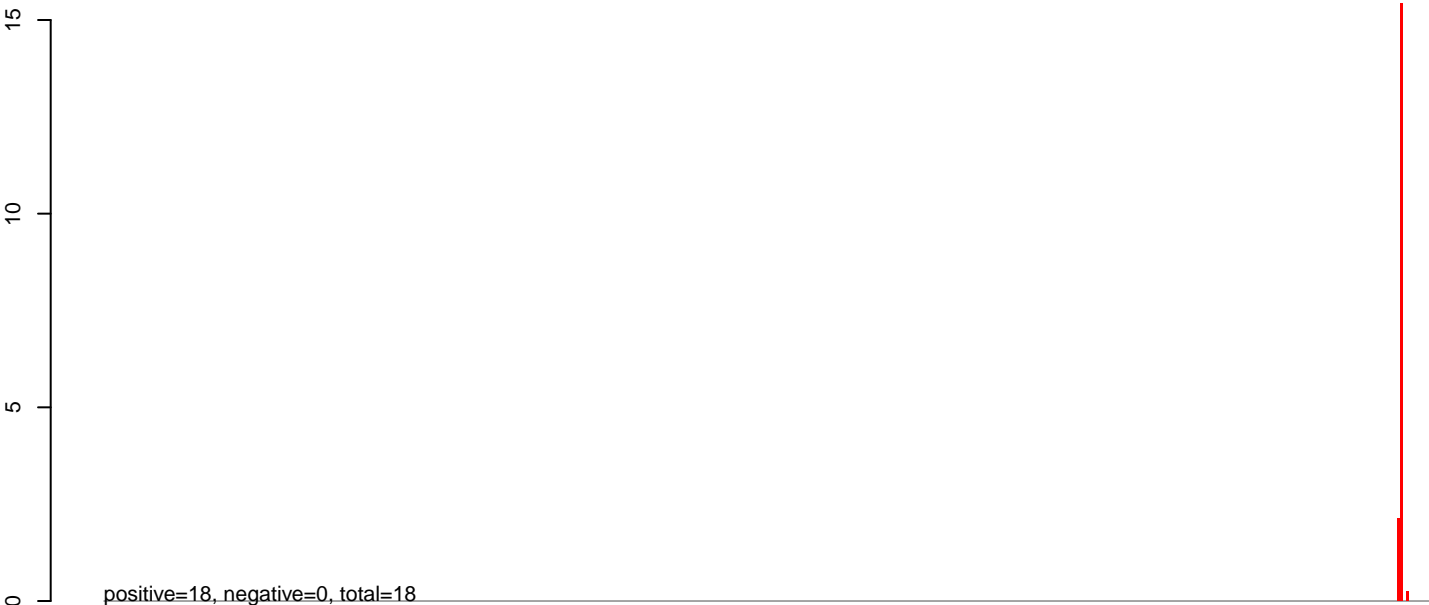
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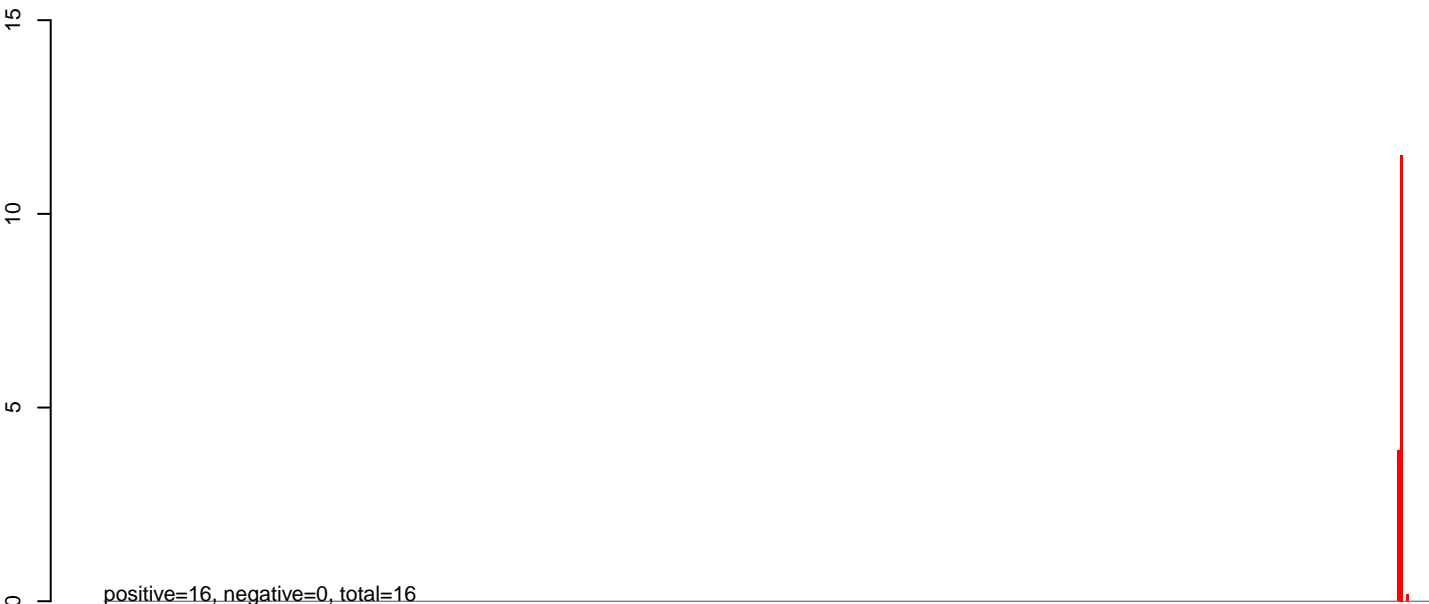
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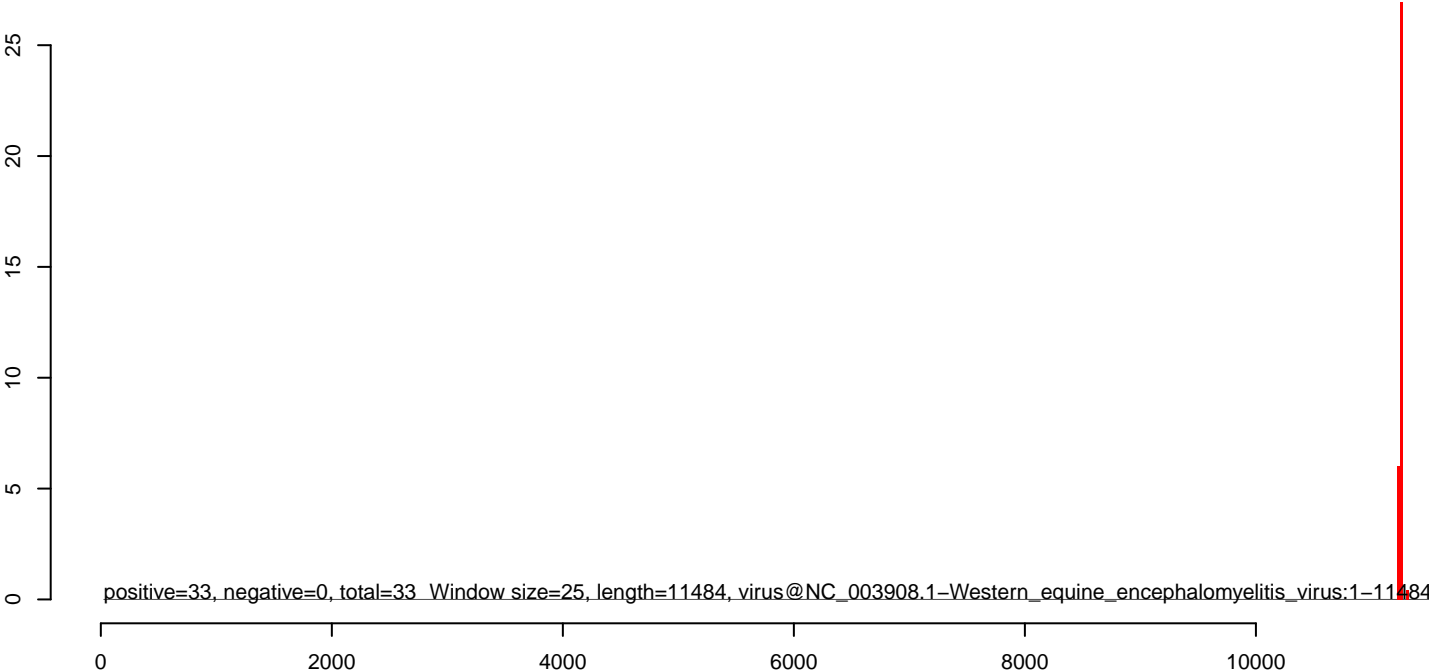
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AeAlbo_C636_SINV_GFP_betaE.24_35.rep



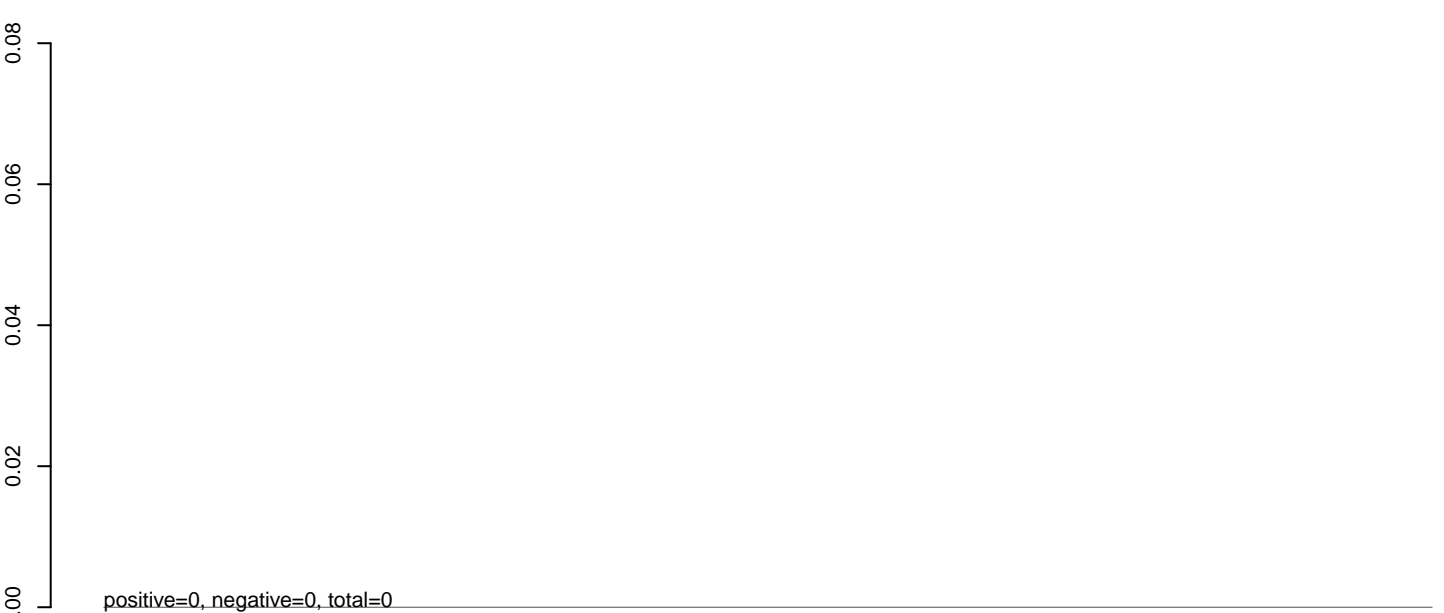
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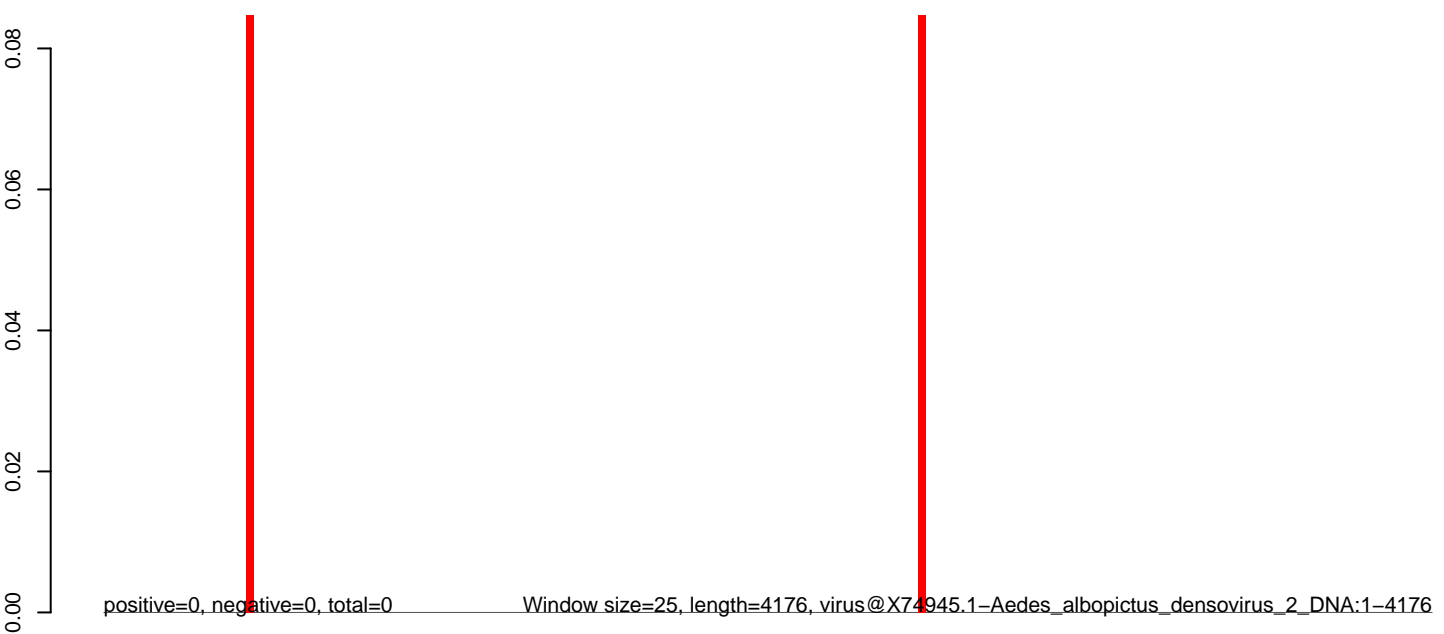
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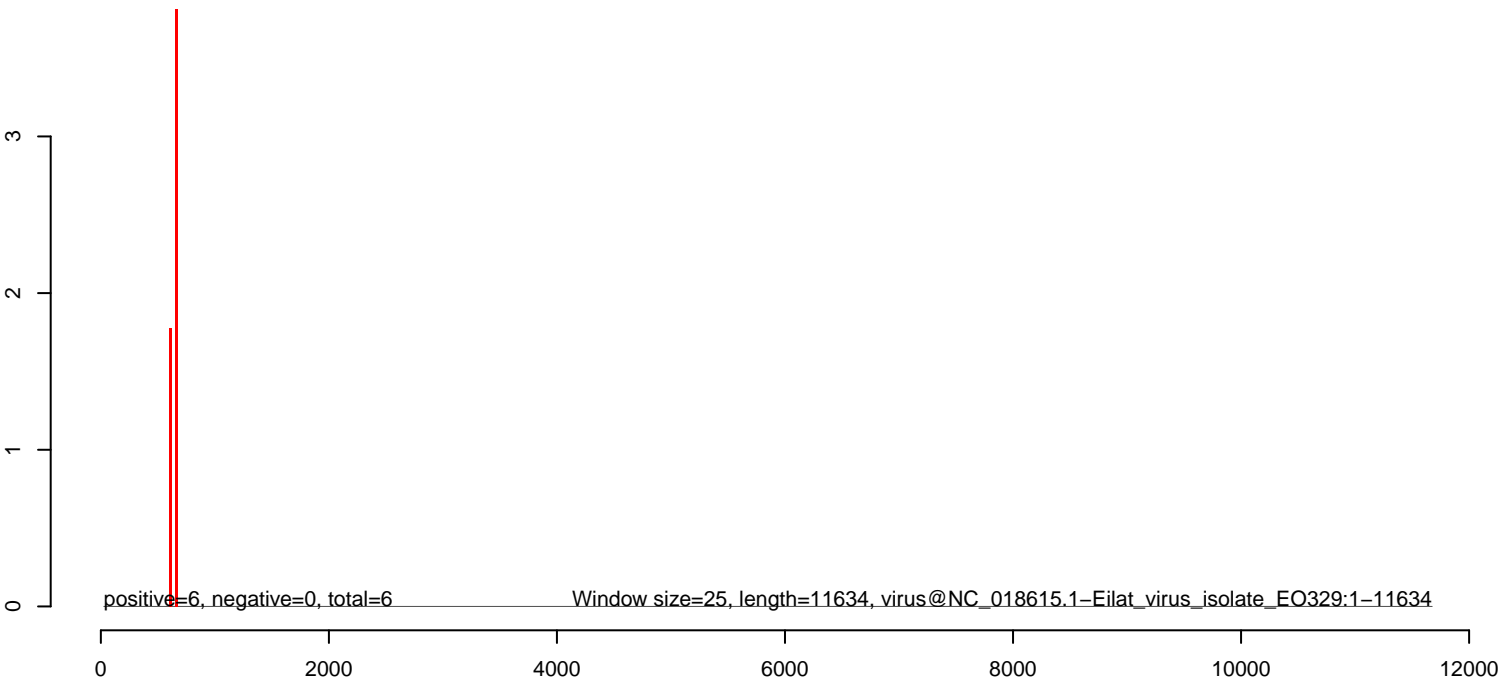
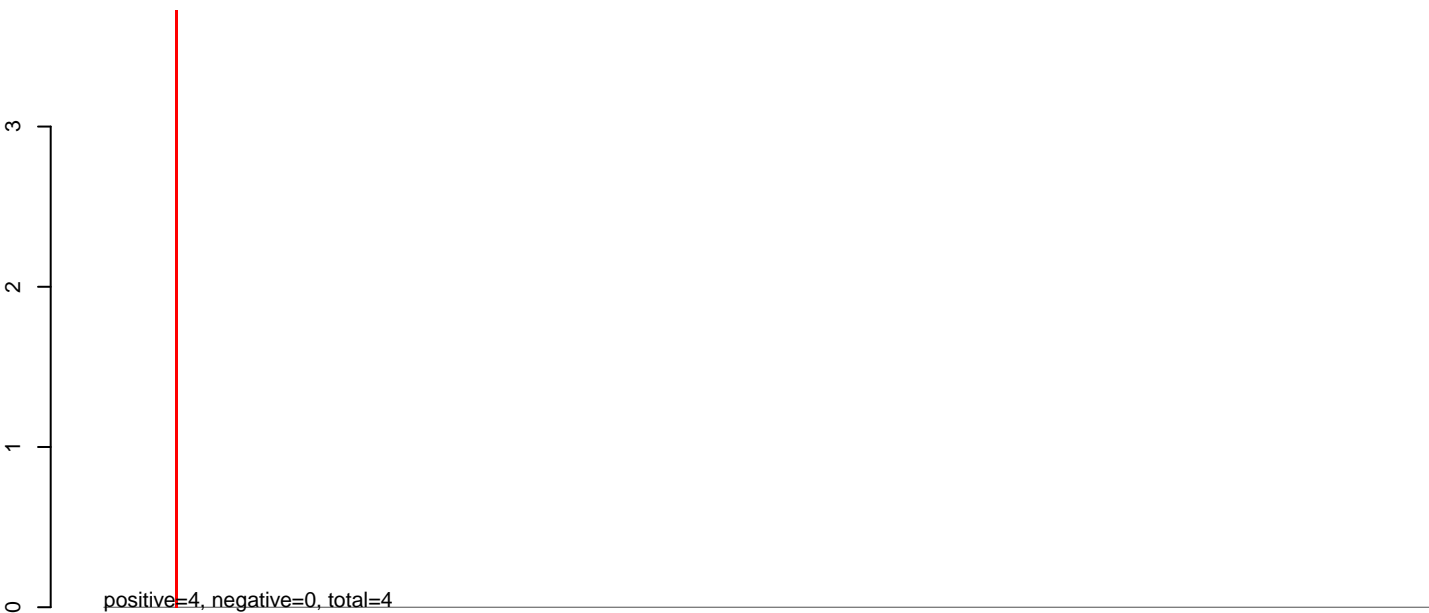
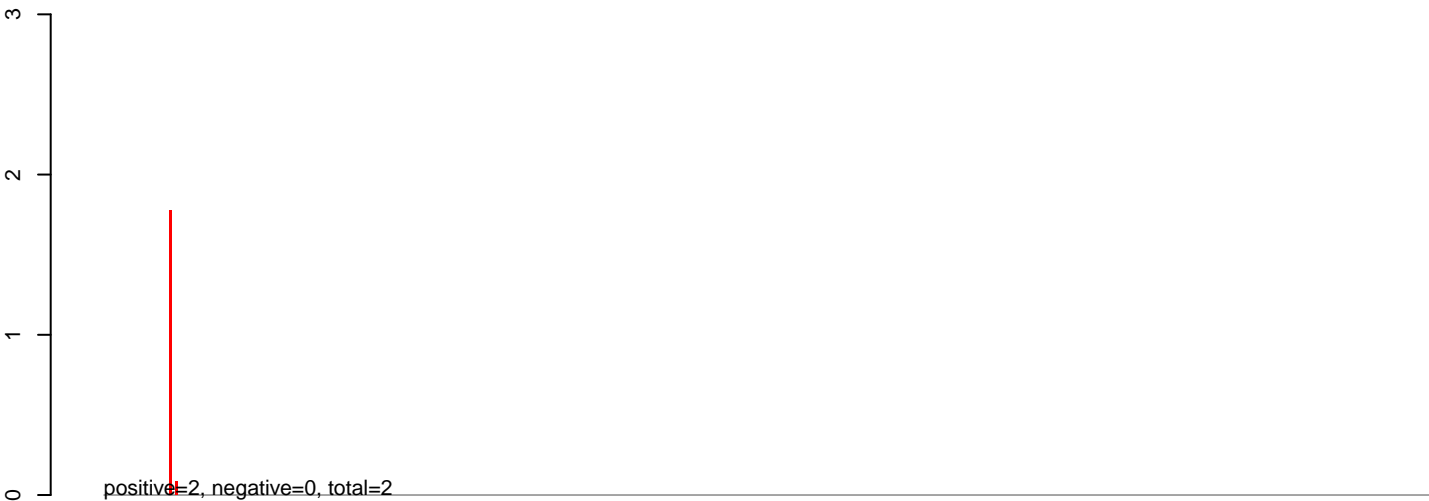


AeAlbo_C636_SINV_GFP_betaE.24_35.rep

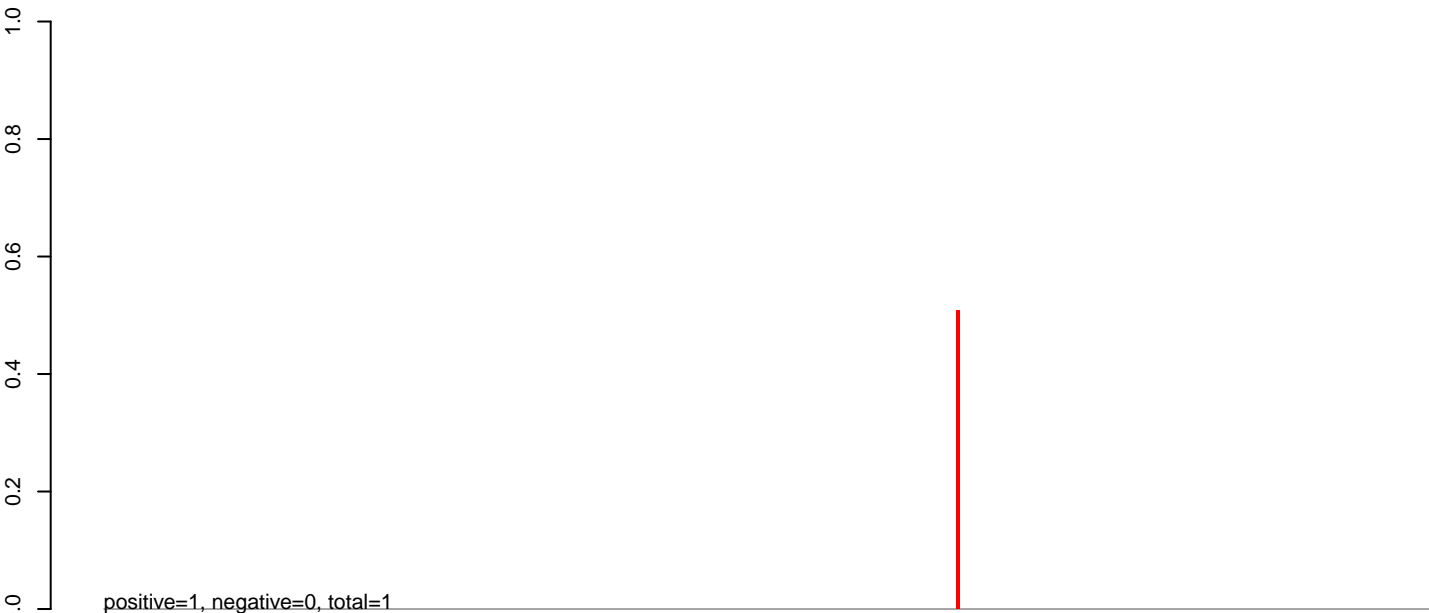


AeAlbo_C636_SINV_GFP_betaE.rep

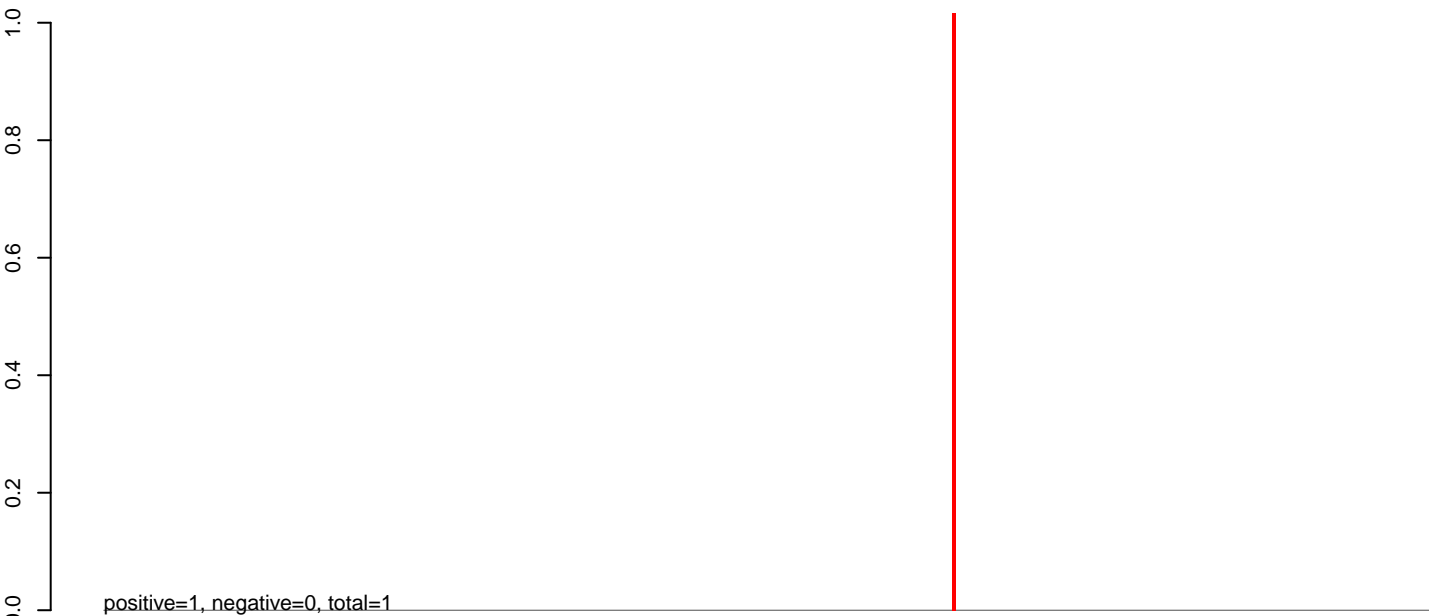




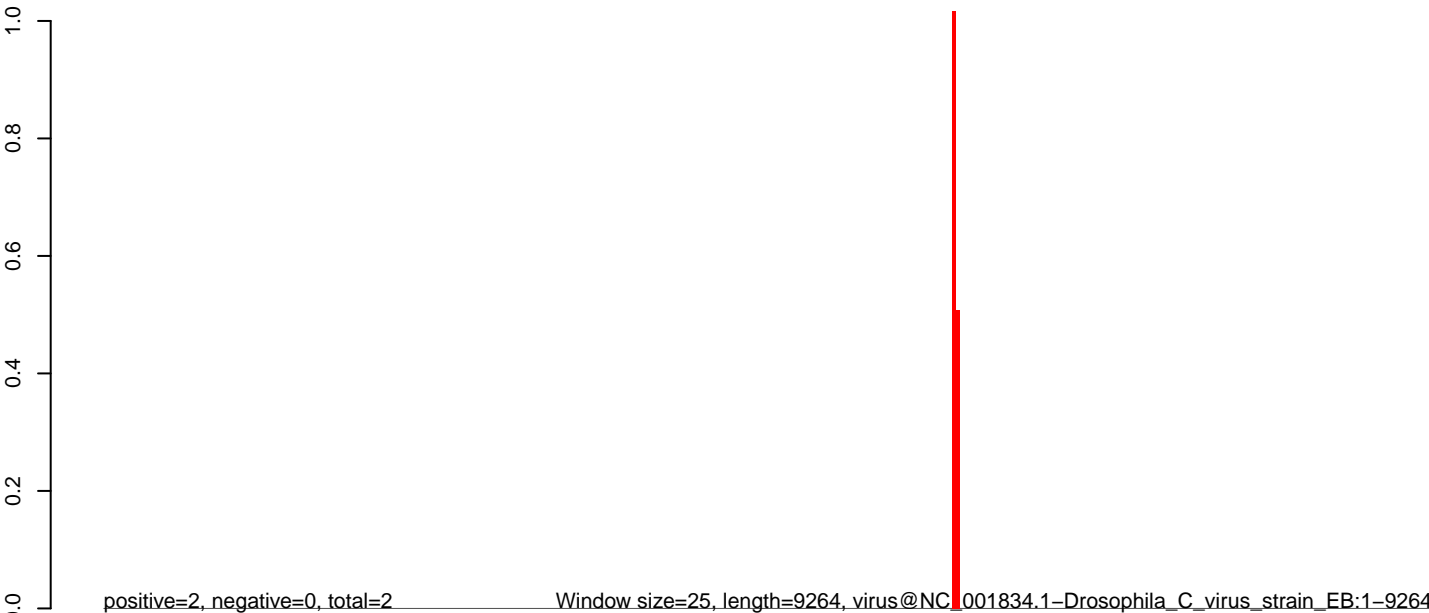
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



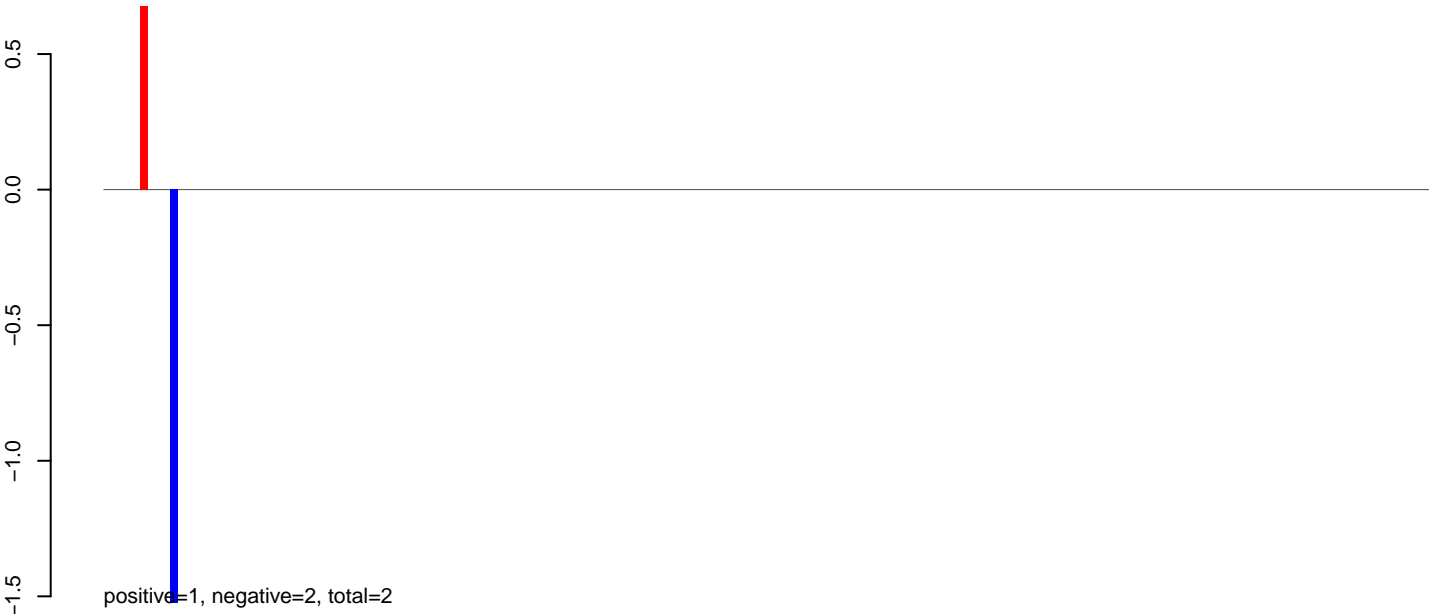
AeAlbo_C636_SINV_GFP_betaE.rep



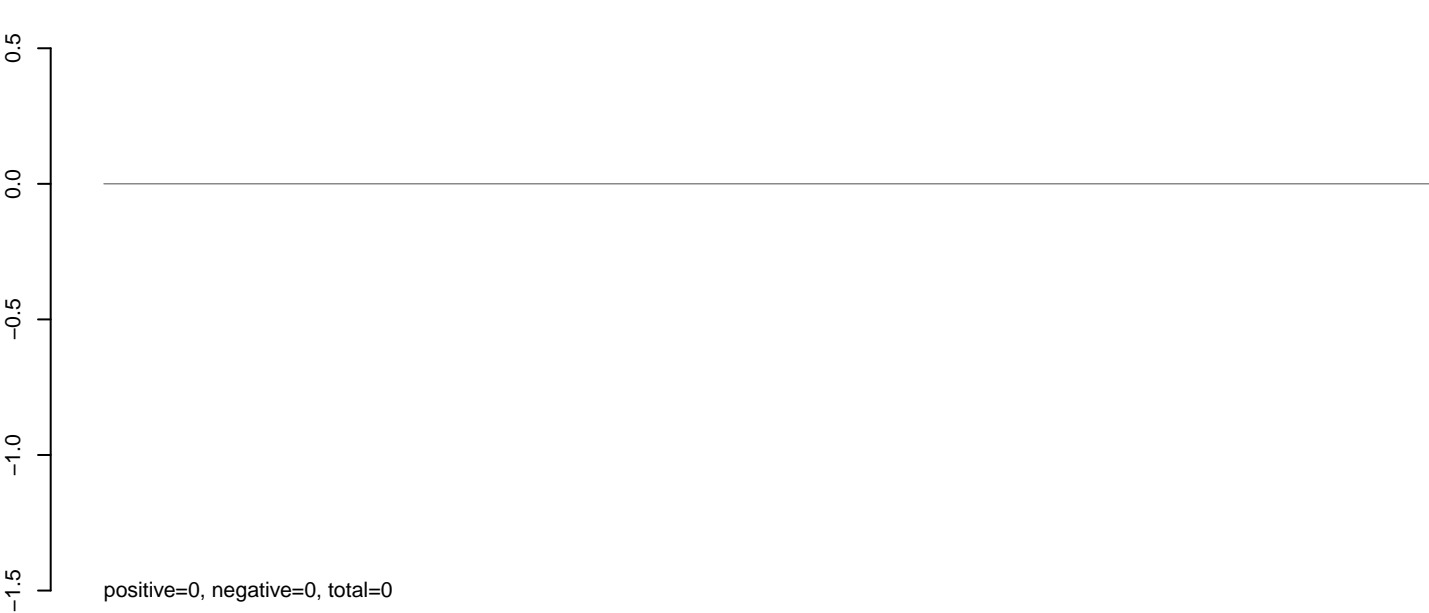
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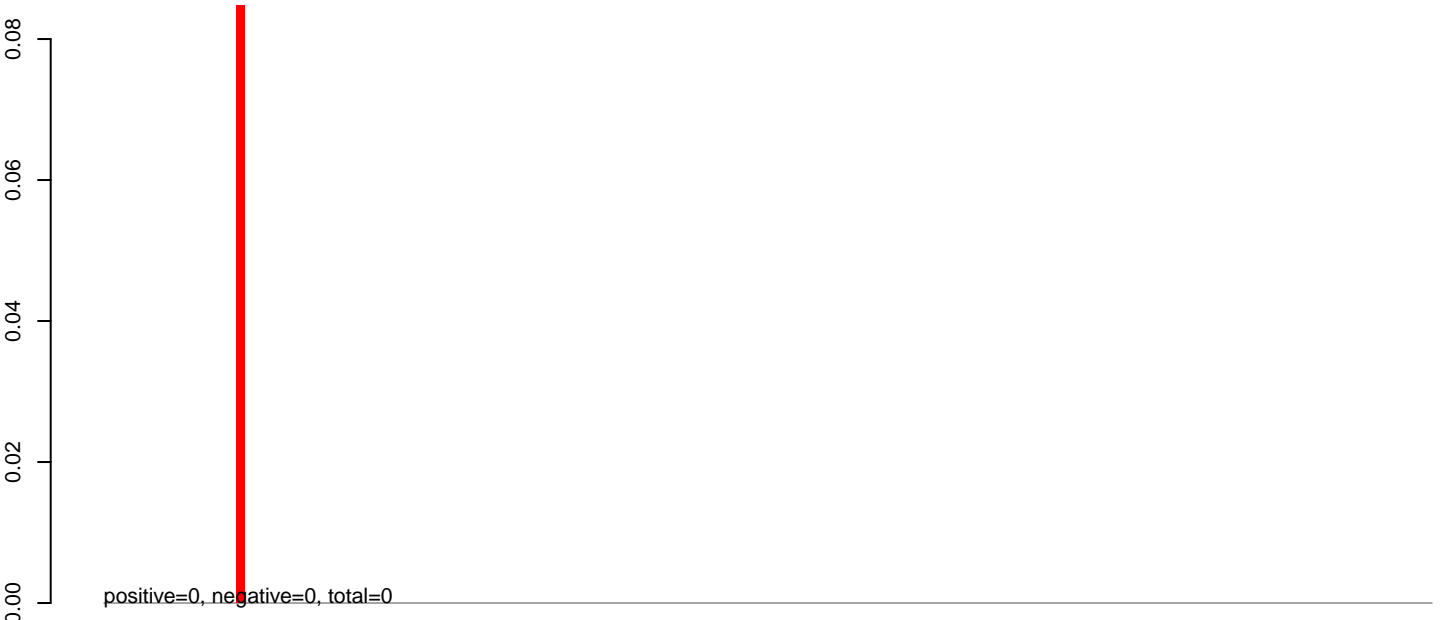
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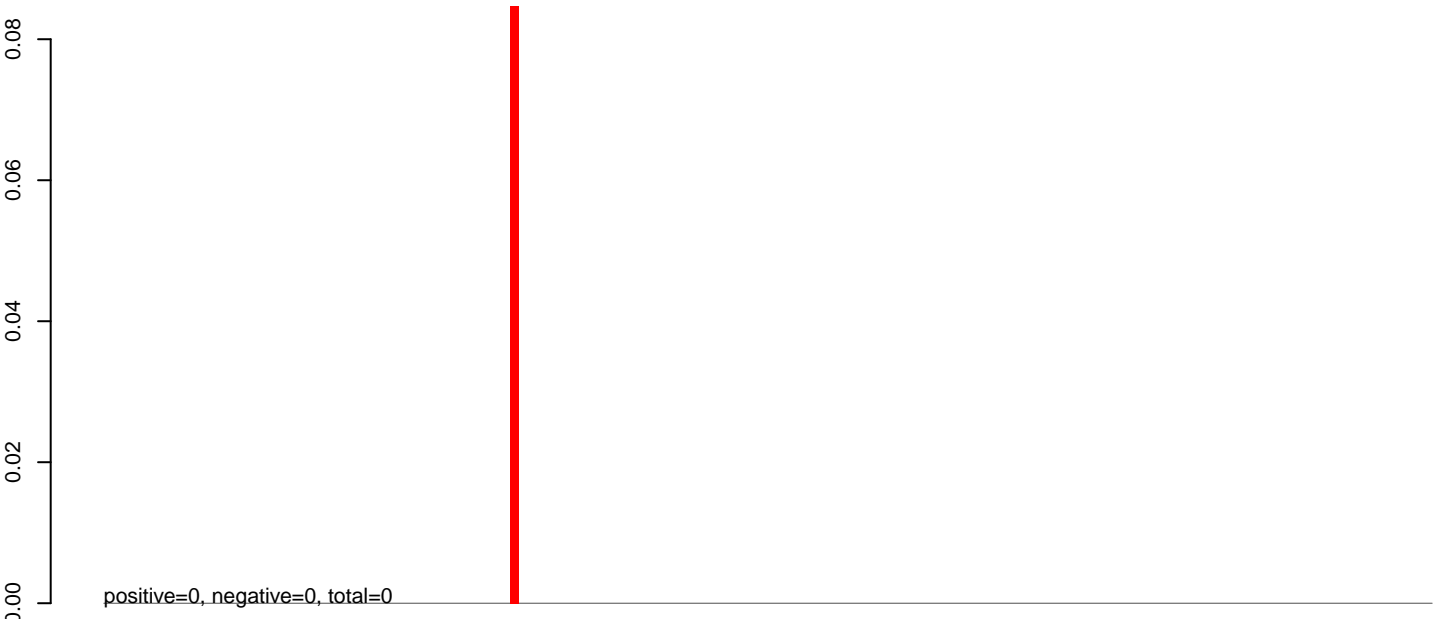
AeAlbo_C636_SINV_GFP_betaE.rep



AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep

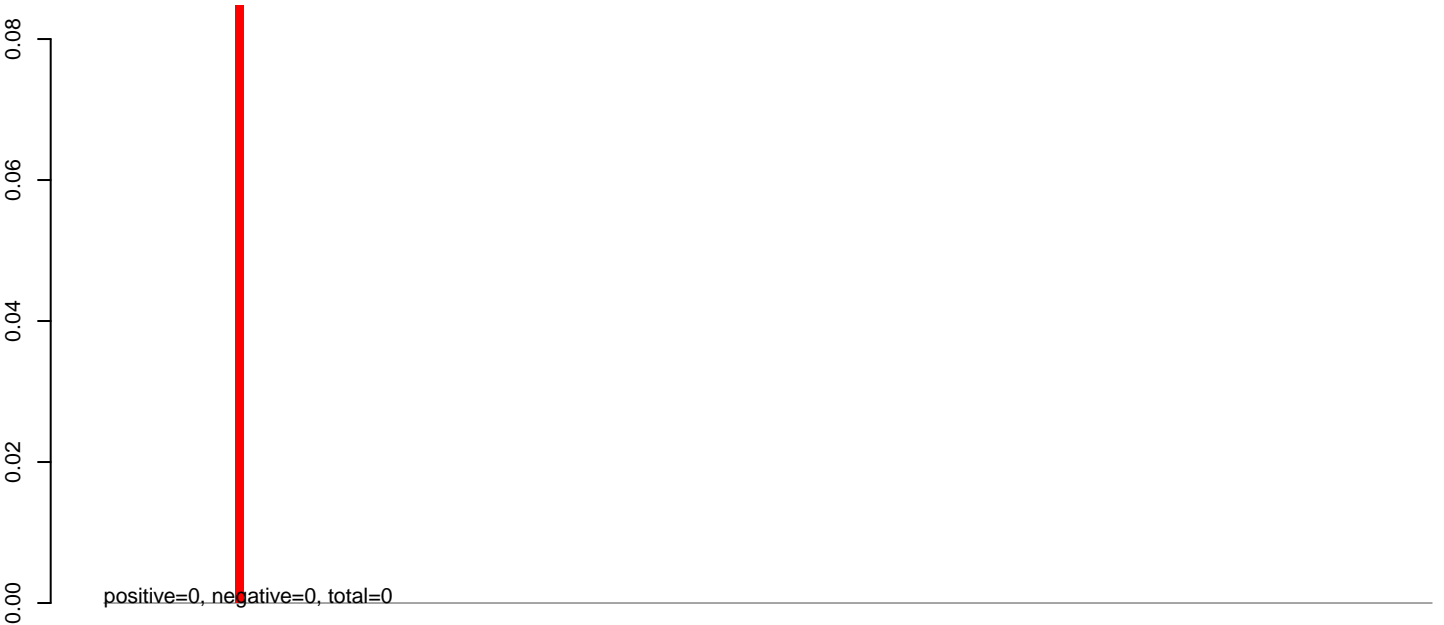


AeAlbo_C636_SINV_GFP_betaE.rep

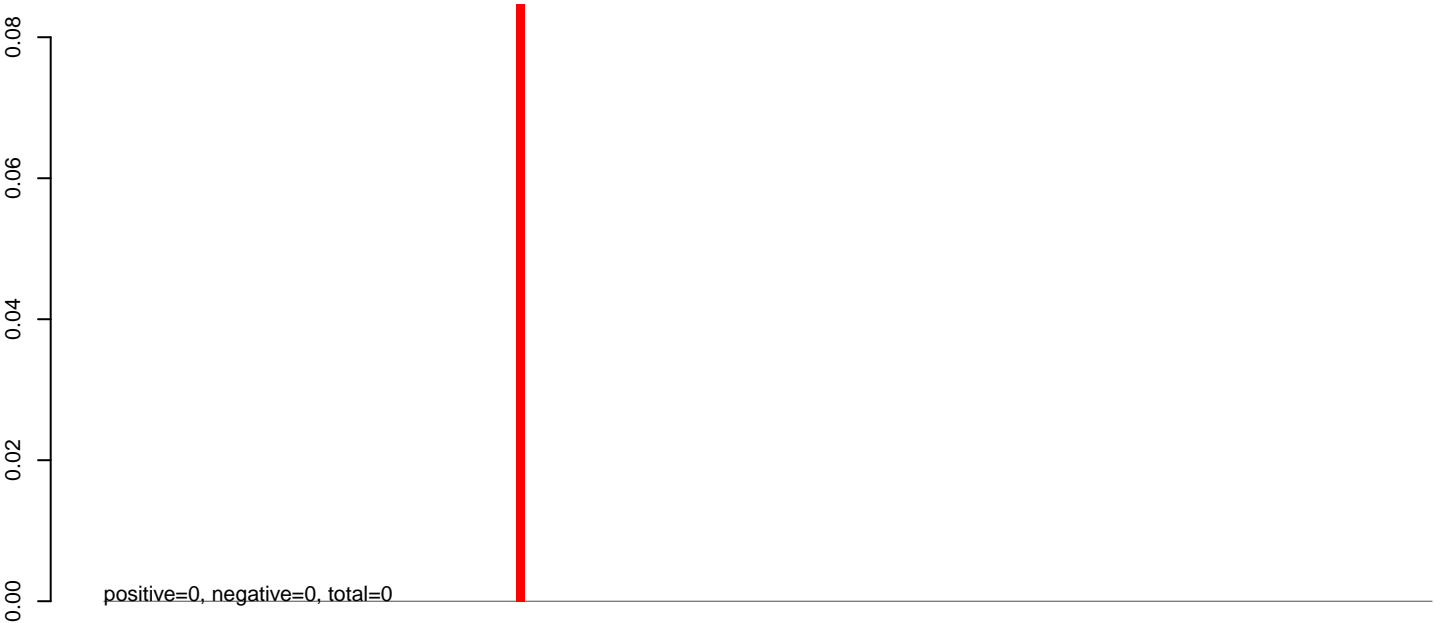


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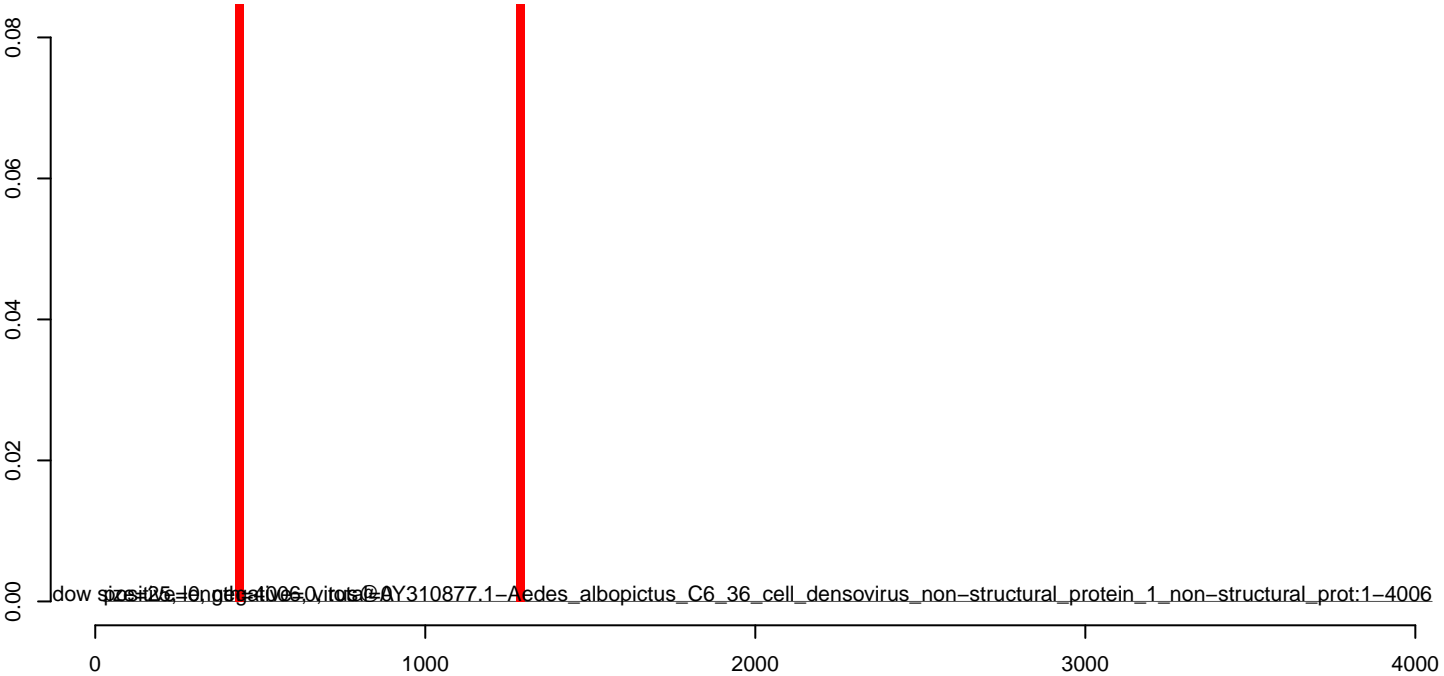
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



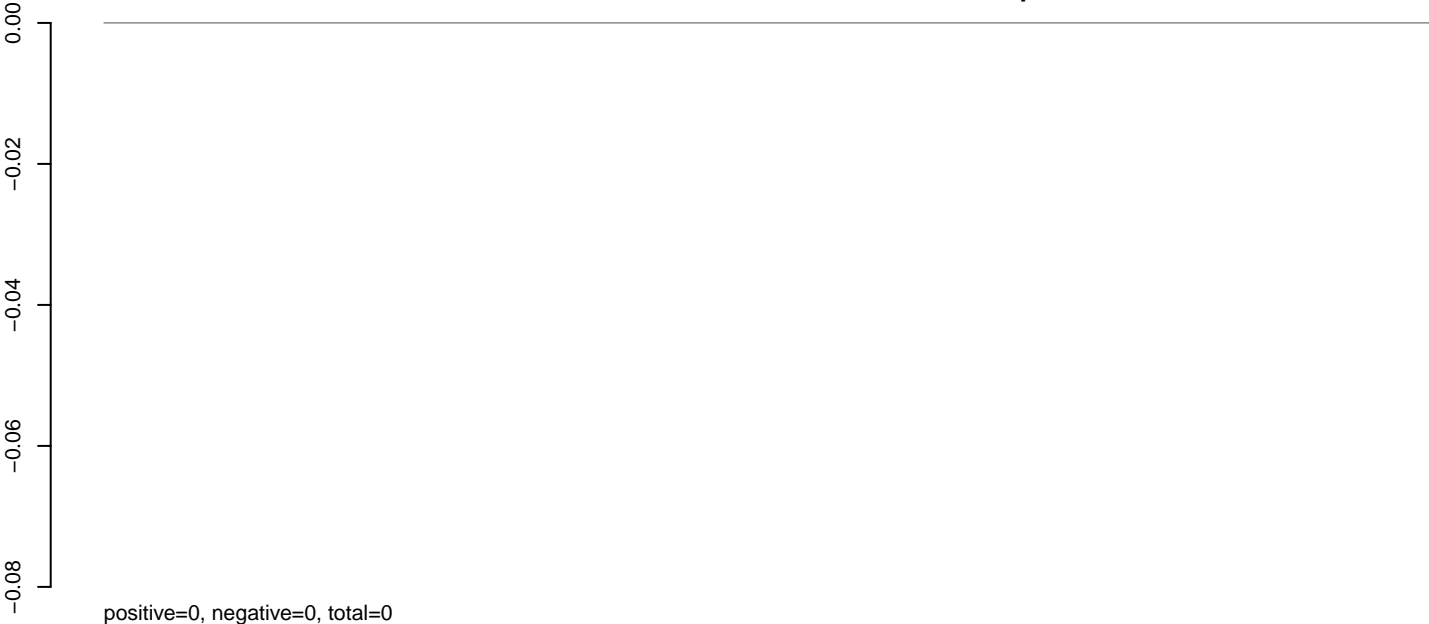
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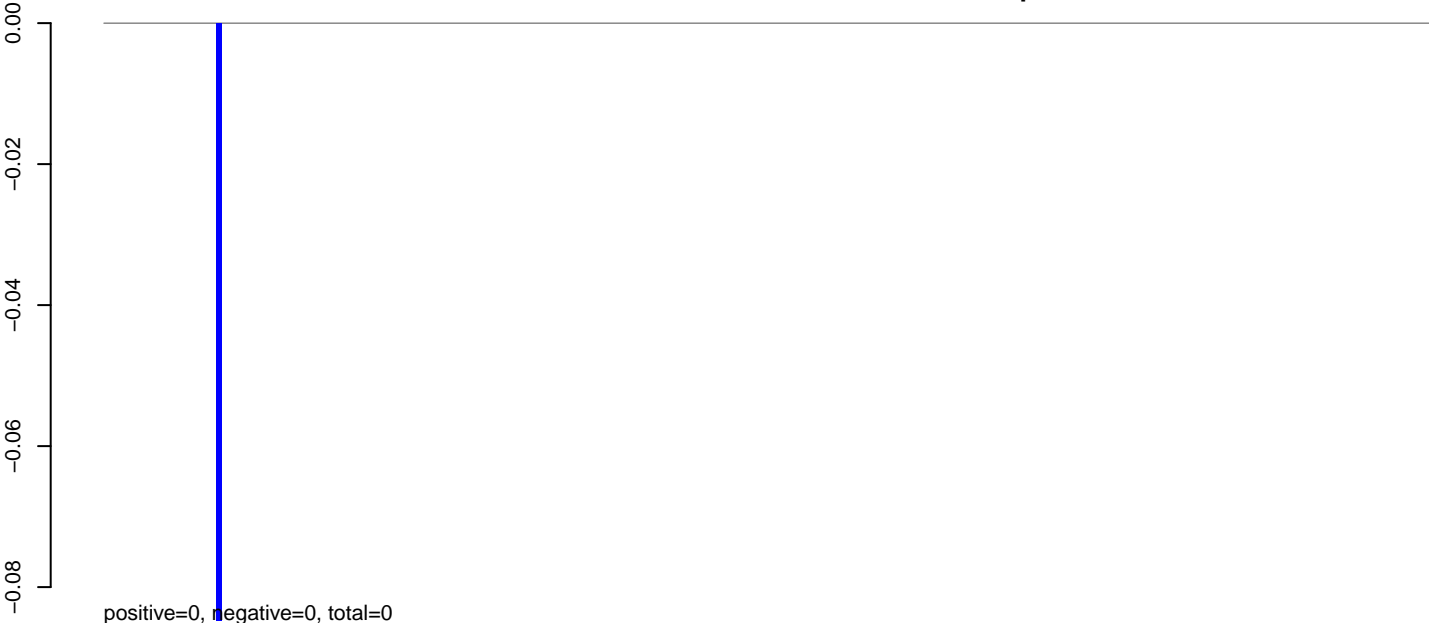
AeAlbo_C636_SINV_GFP_betaE.rep



AeAlbo_C636_SINV_GFP_betaE.18_23.rep



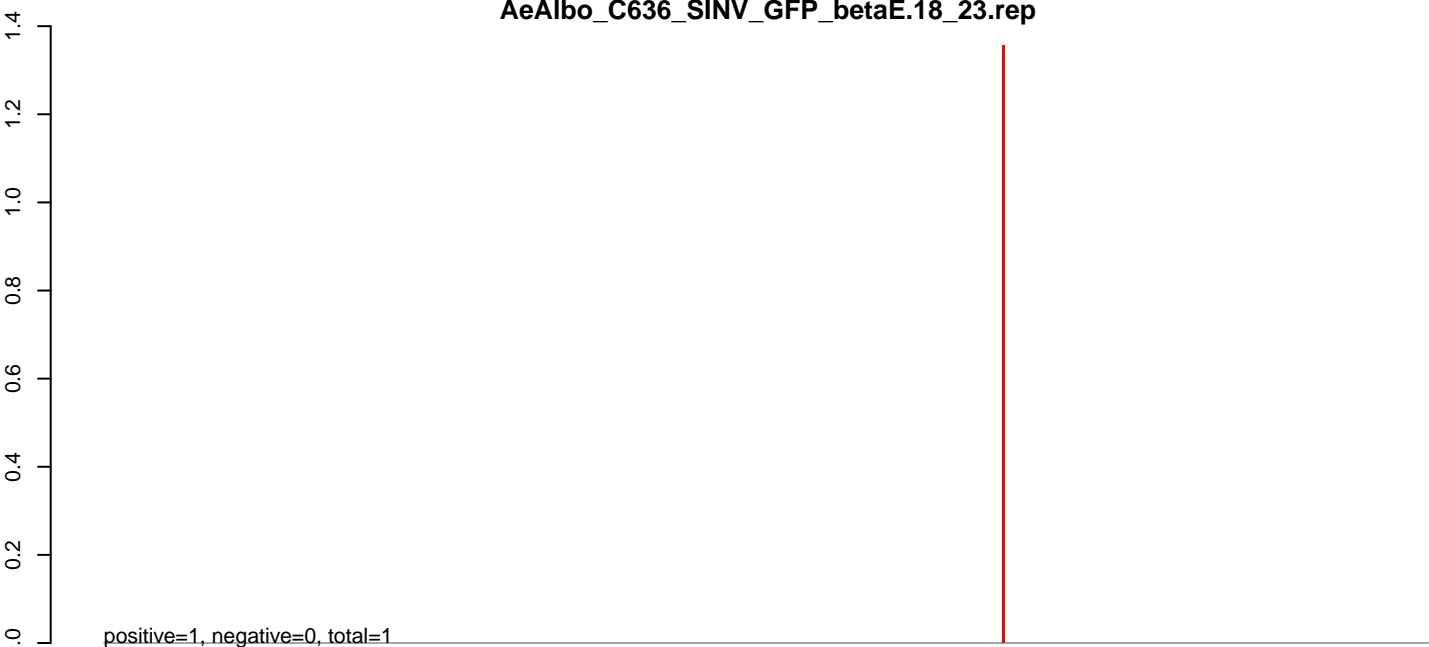
AeAlbo_C636_SINV_GFP_betaE.24_35.rep



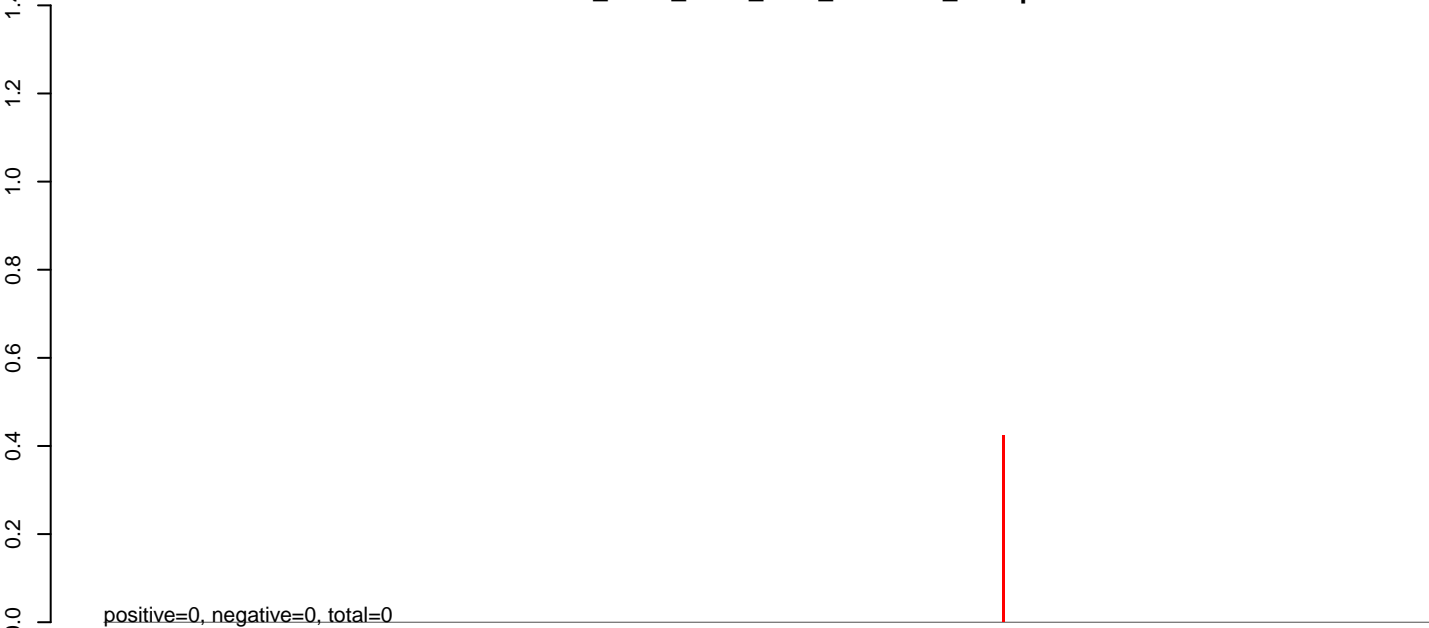
AeAlbo_C636_SINV_GFP_betaE.rep



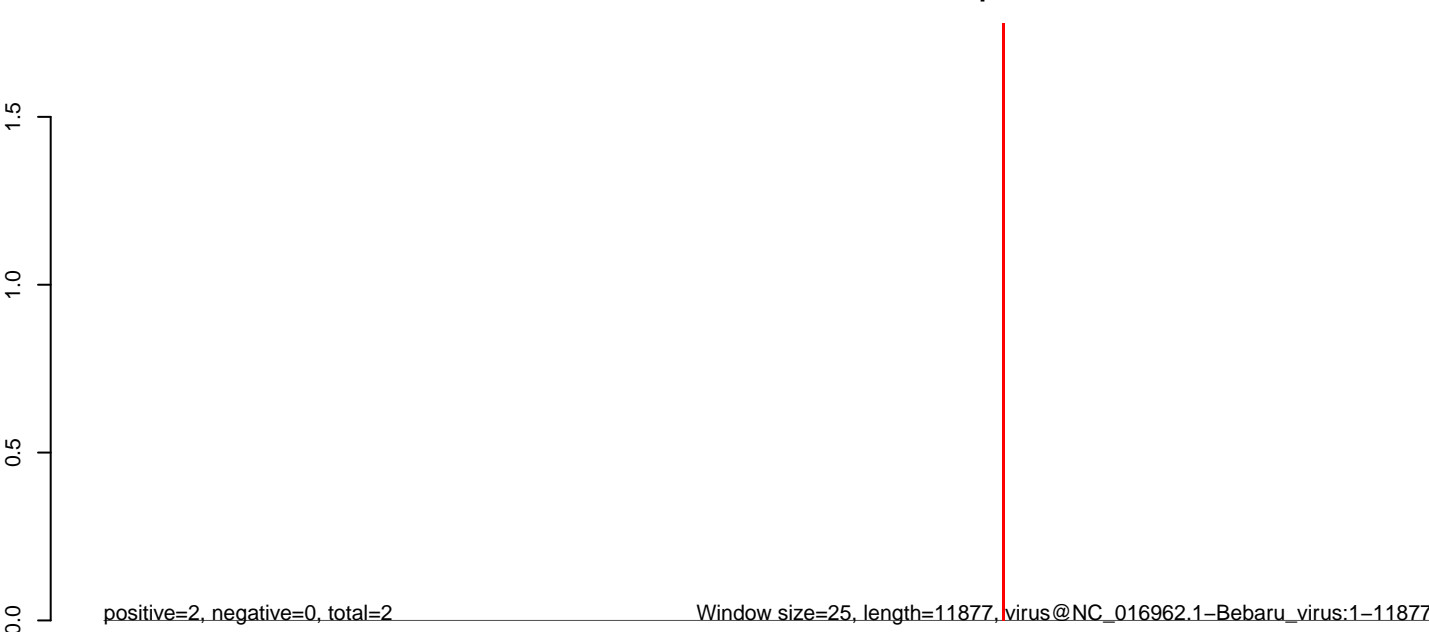
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AeAlbo_C636_SINV_GFP_betaE.24_35.rep



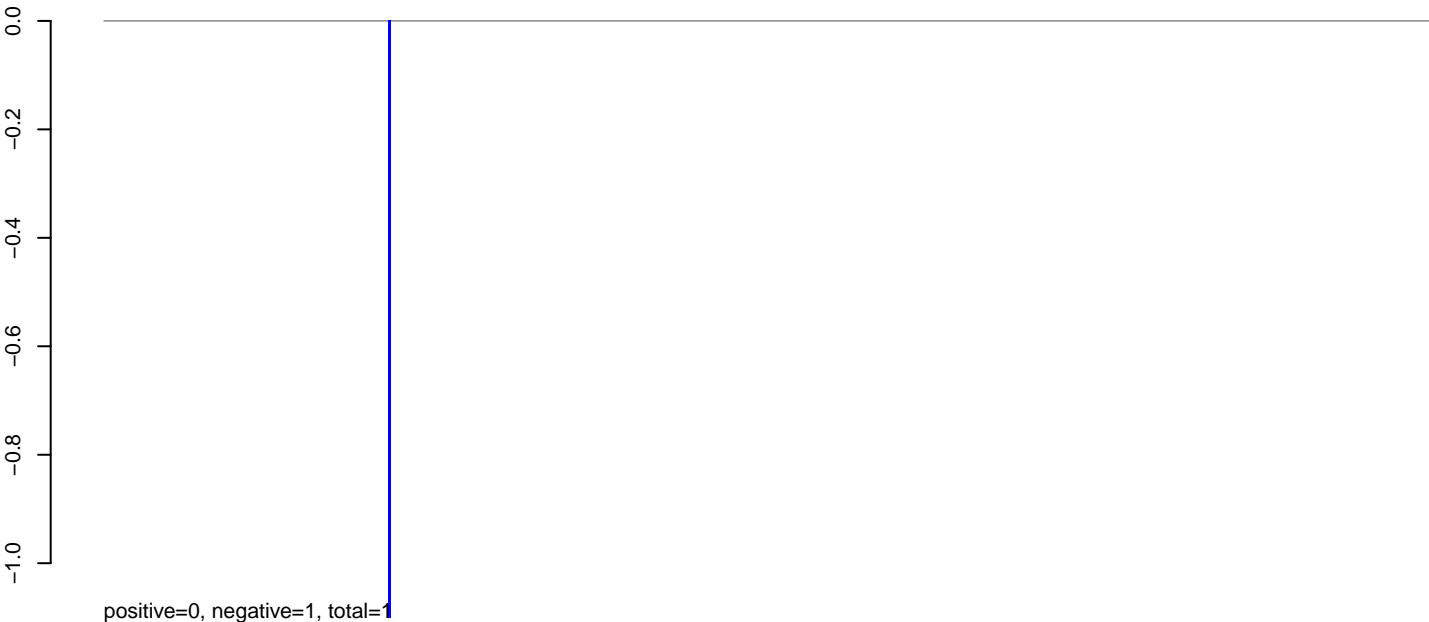
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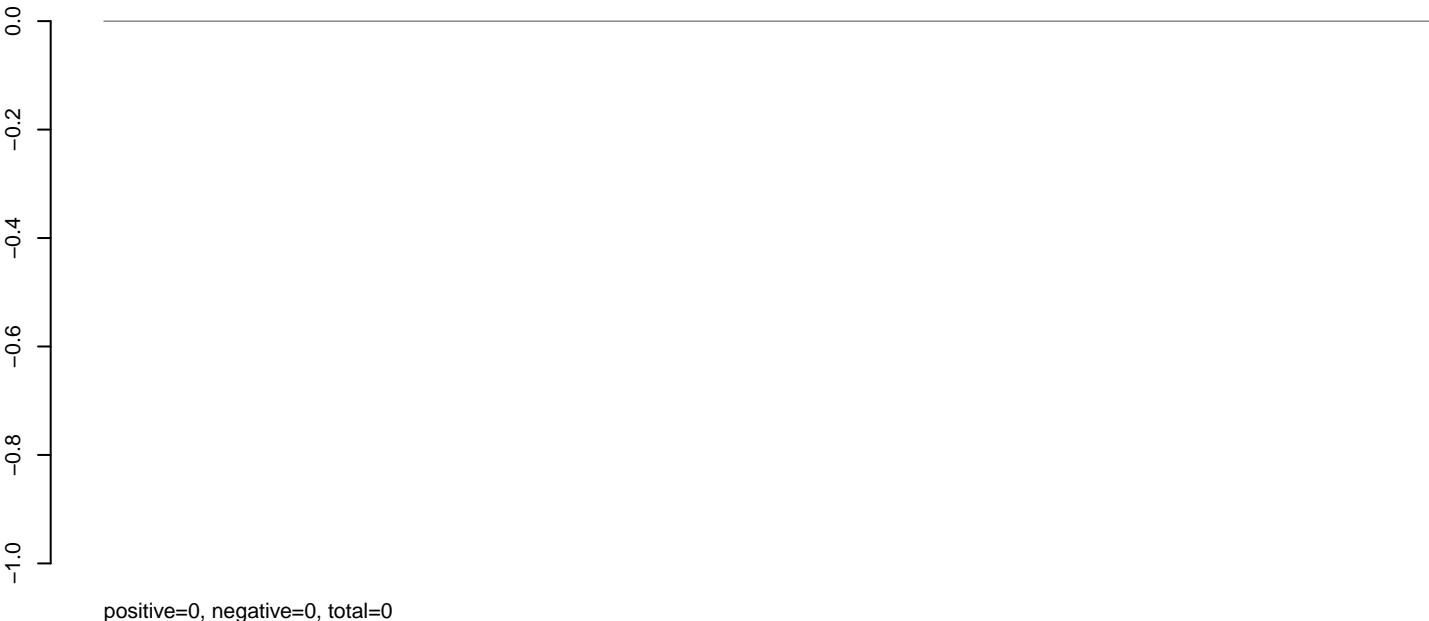
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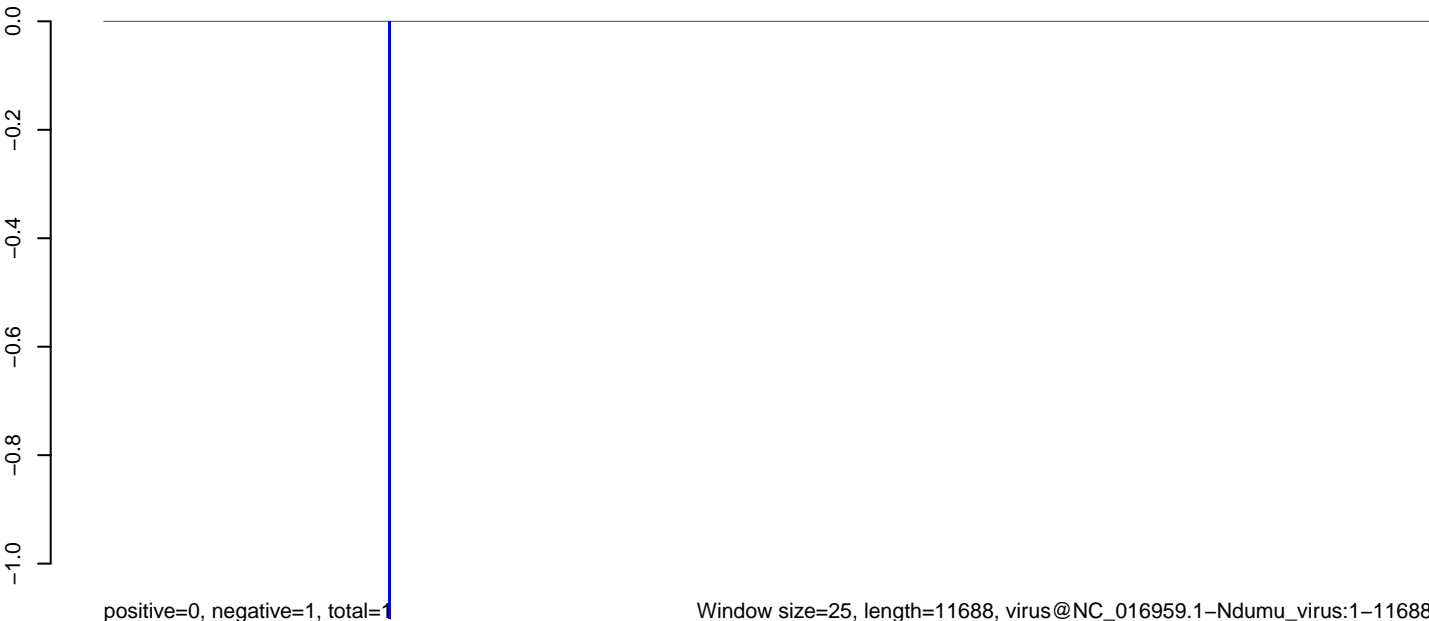
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



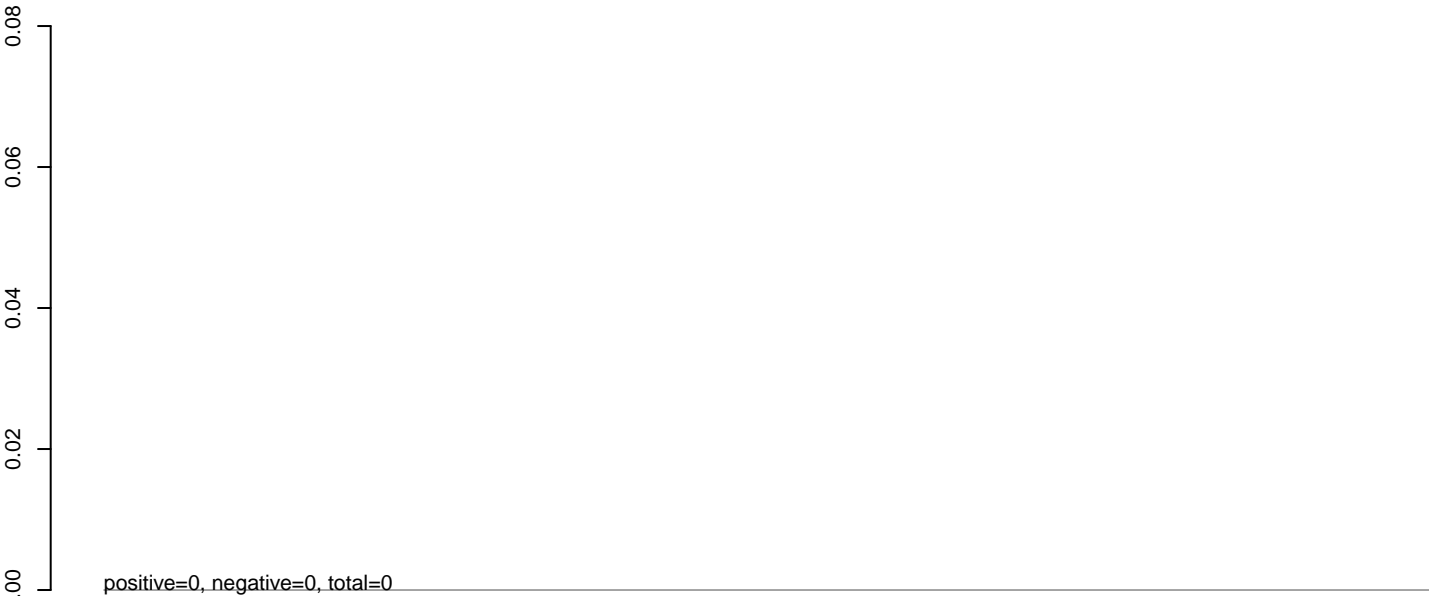
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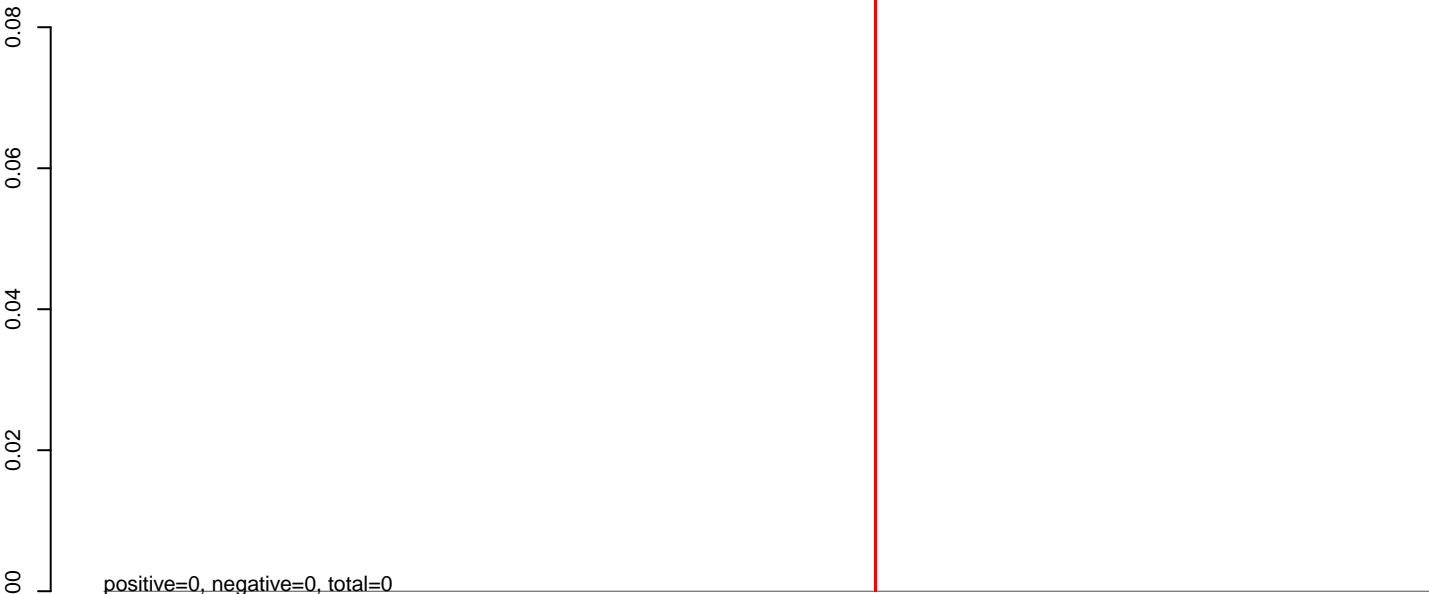
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0 2000 4000 6000 8000 10000 12000

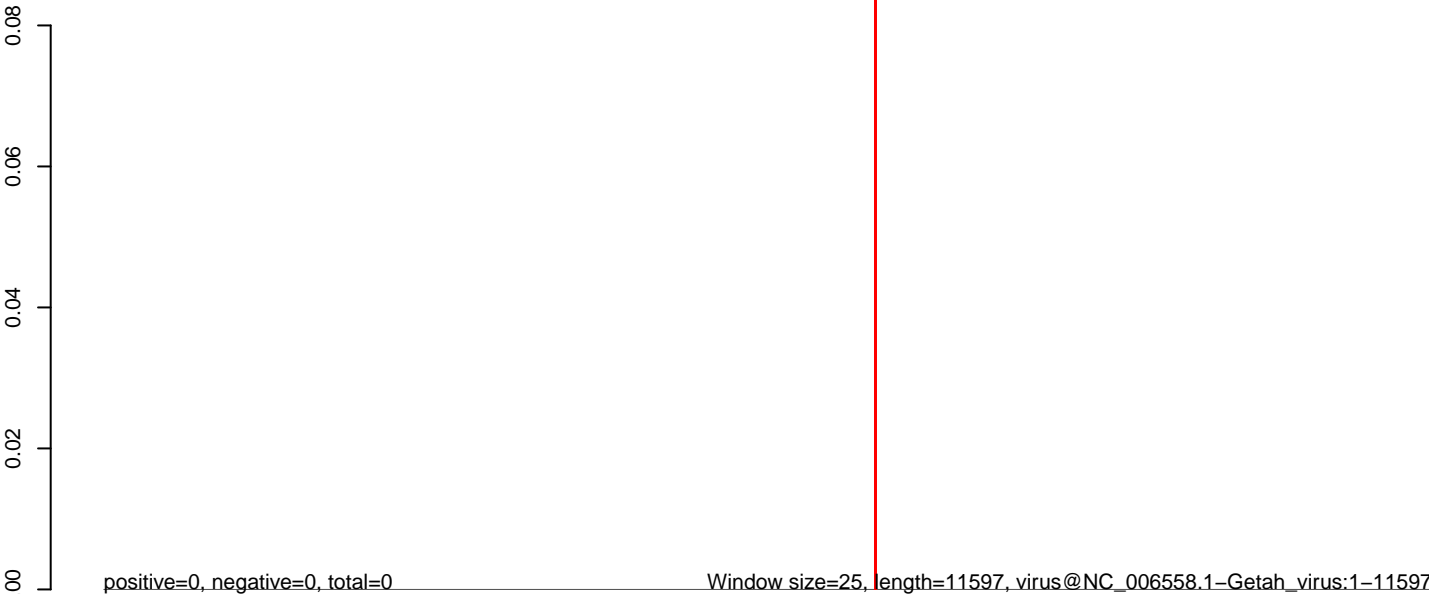
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep

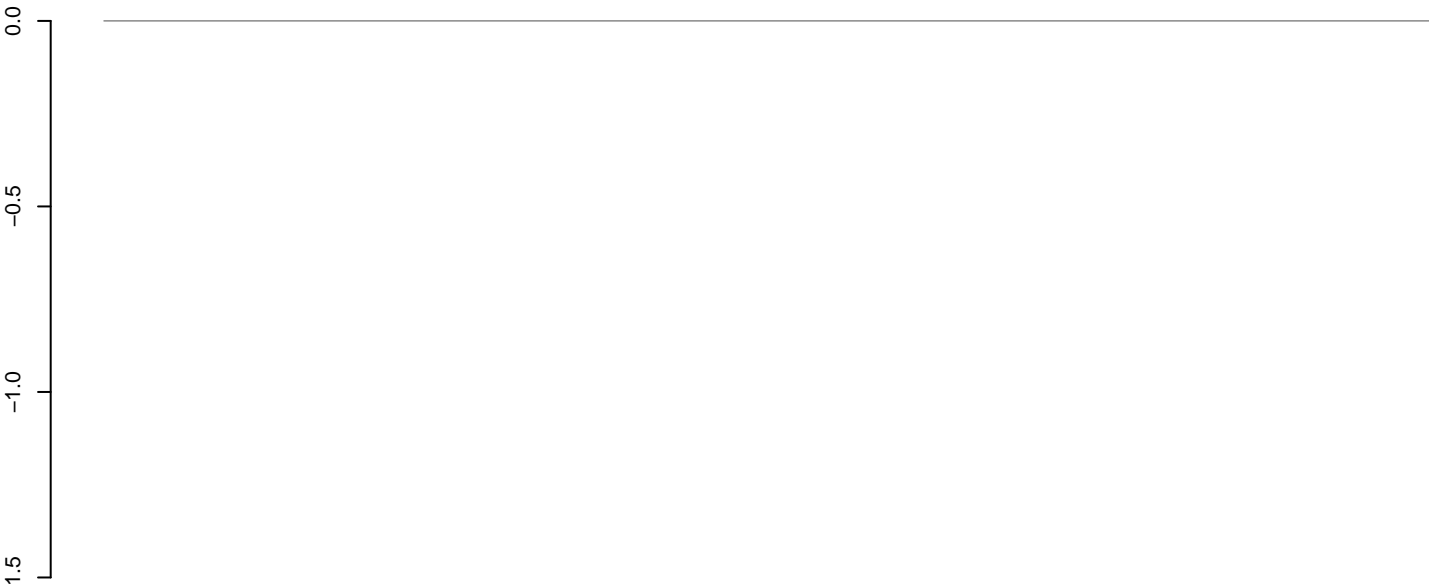


AeAlbo_C636_SINV_GFP_betaE.rep



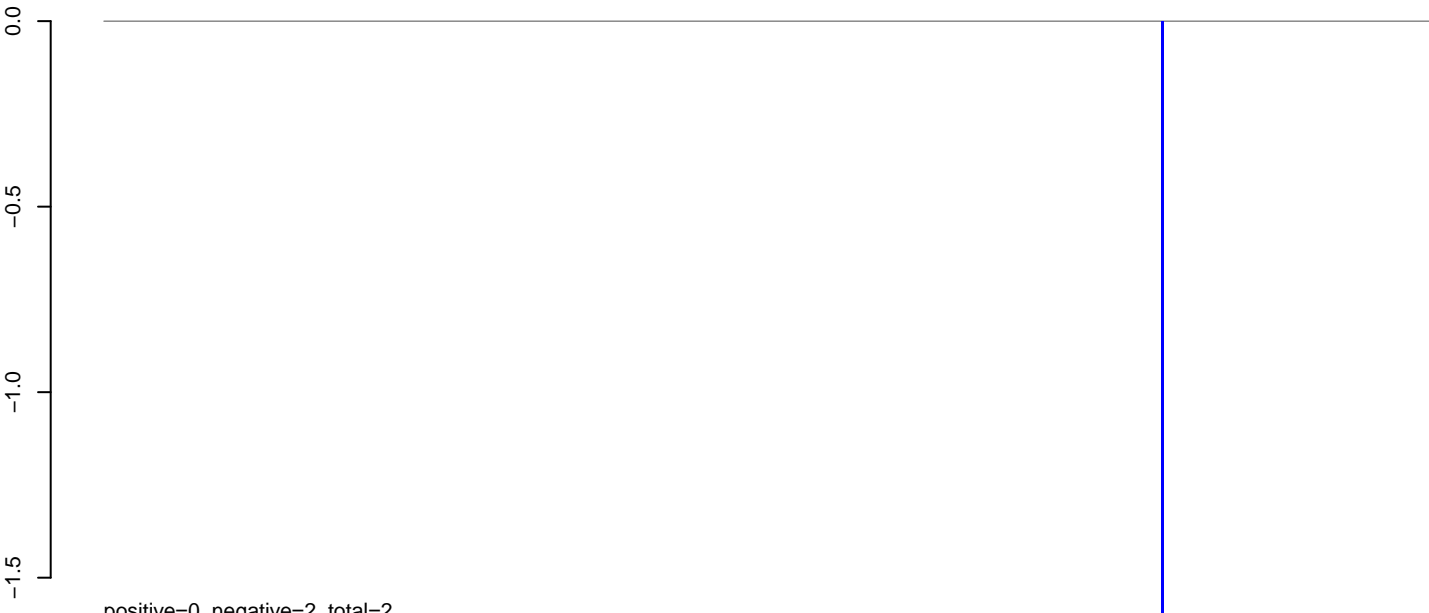
0 2000 4000 6000 8000 10000 12000

AeAlbo_C636_SINV_GFP_betaE.18_23.rep



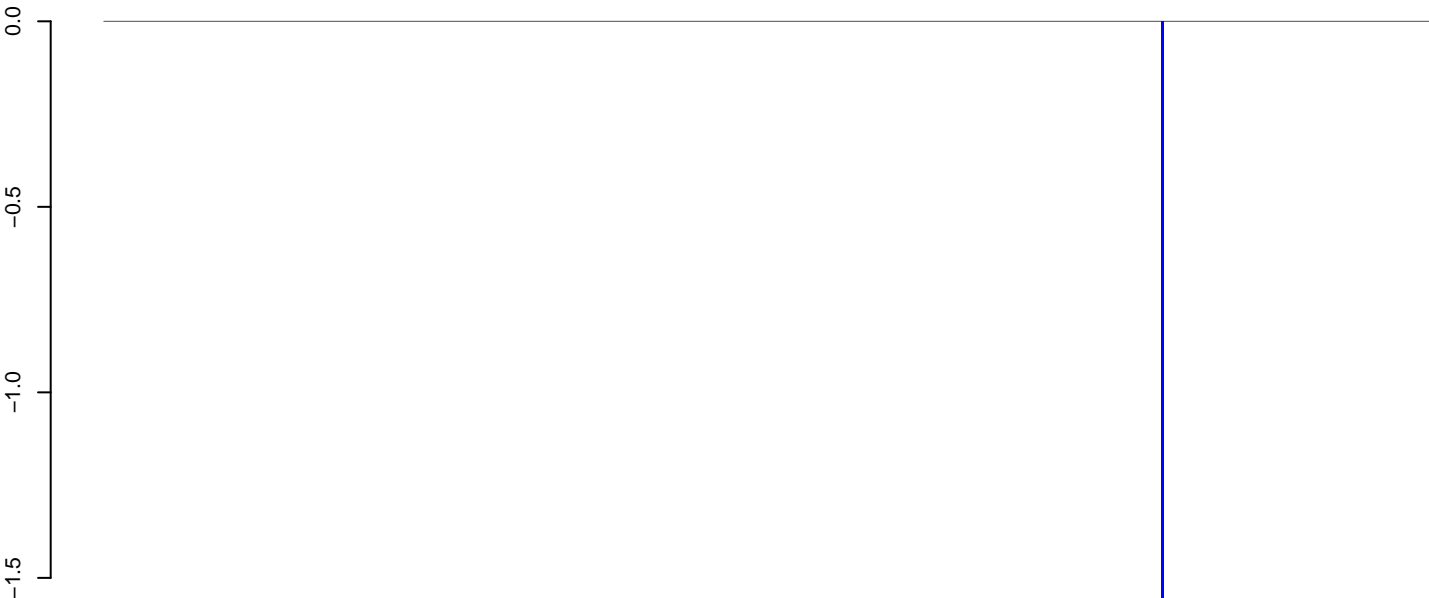
positive=0, negative=0, total=0

AeAlbo_C636_SINV_GFP_betaE.24_35.rep



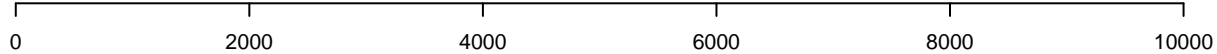
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AeAlbo_C636_SINV_GFP_betaE.rep

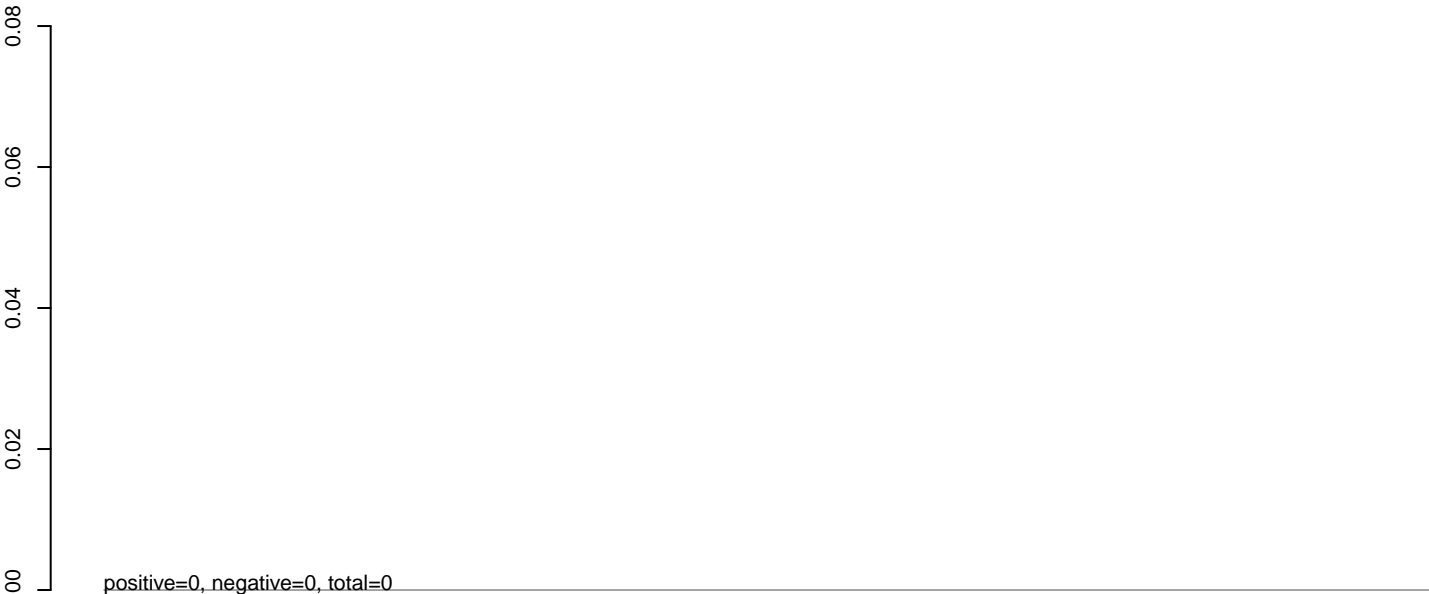


positive=0, negative=2, total=2

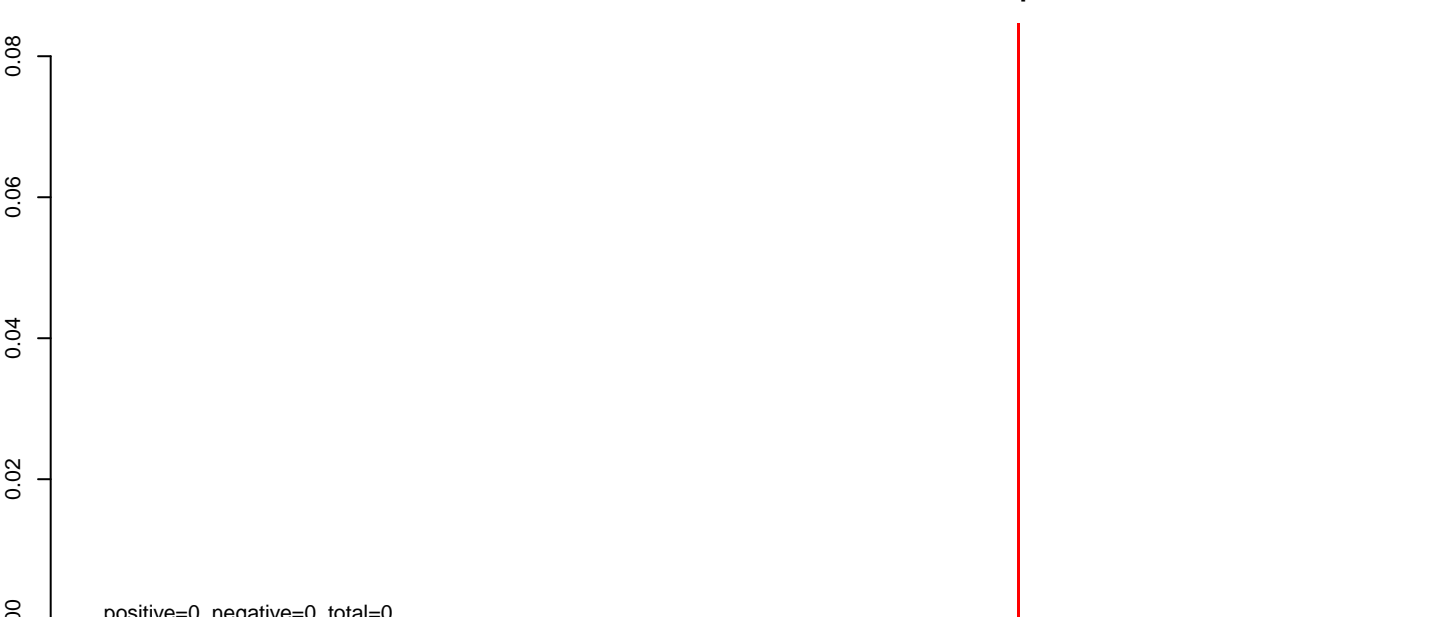
Window size=25, length=11375, virus@NC_005064.1-Kamiti_River_virus:1-11375



AeAlbo_C636_SINV_GFP_betaE.18_23.rep



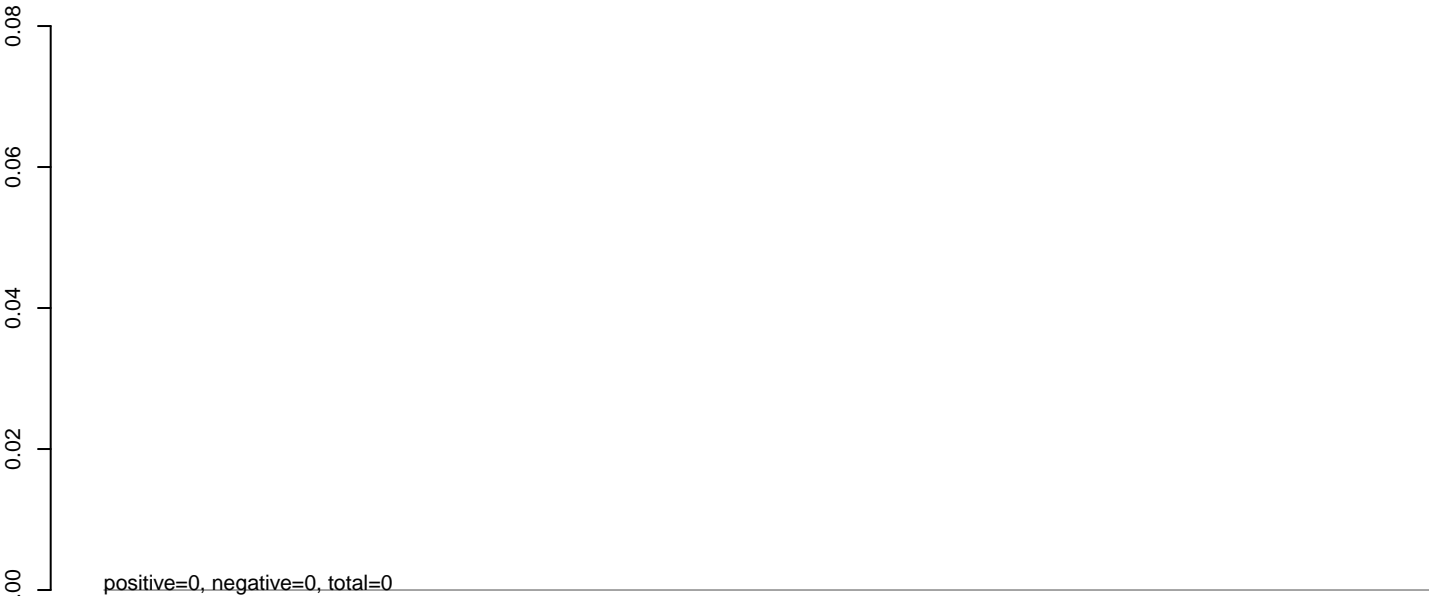
AeAlbo_C636_SINV_GFP_betaE.24_35.rep



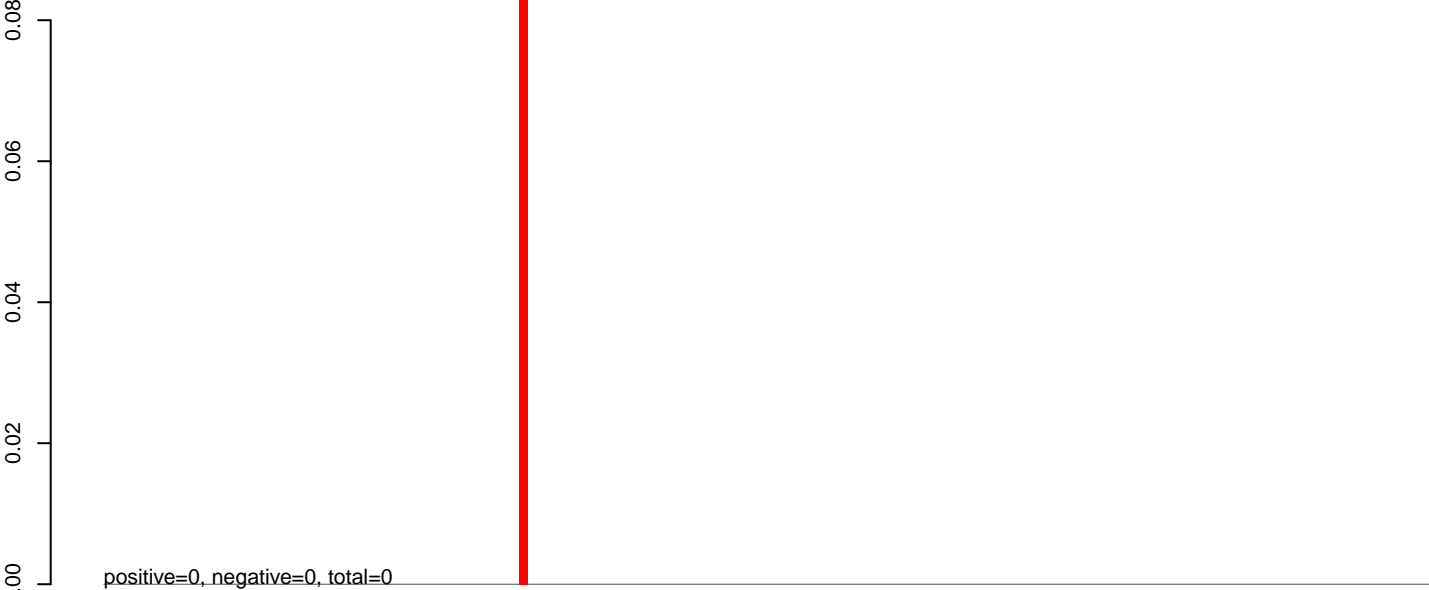
AeAlbo_C636_SINV_GFP_betaE.rep



AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep

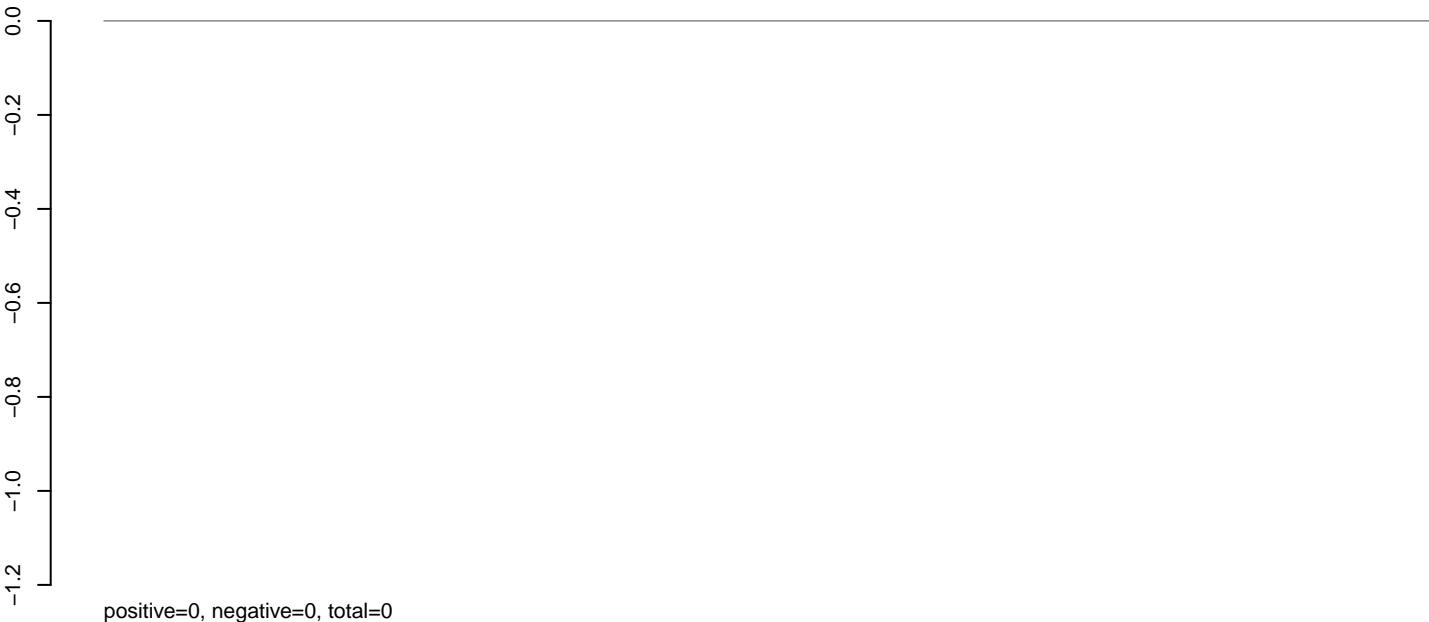


AeAlbo_C636_SINV_GFP_betaE.rep

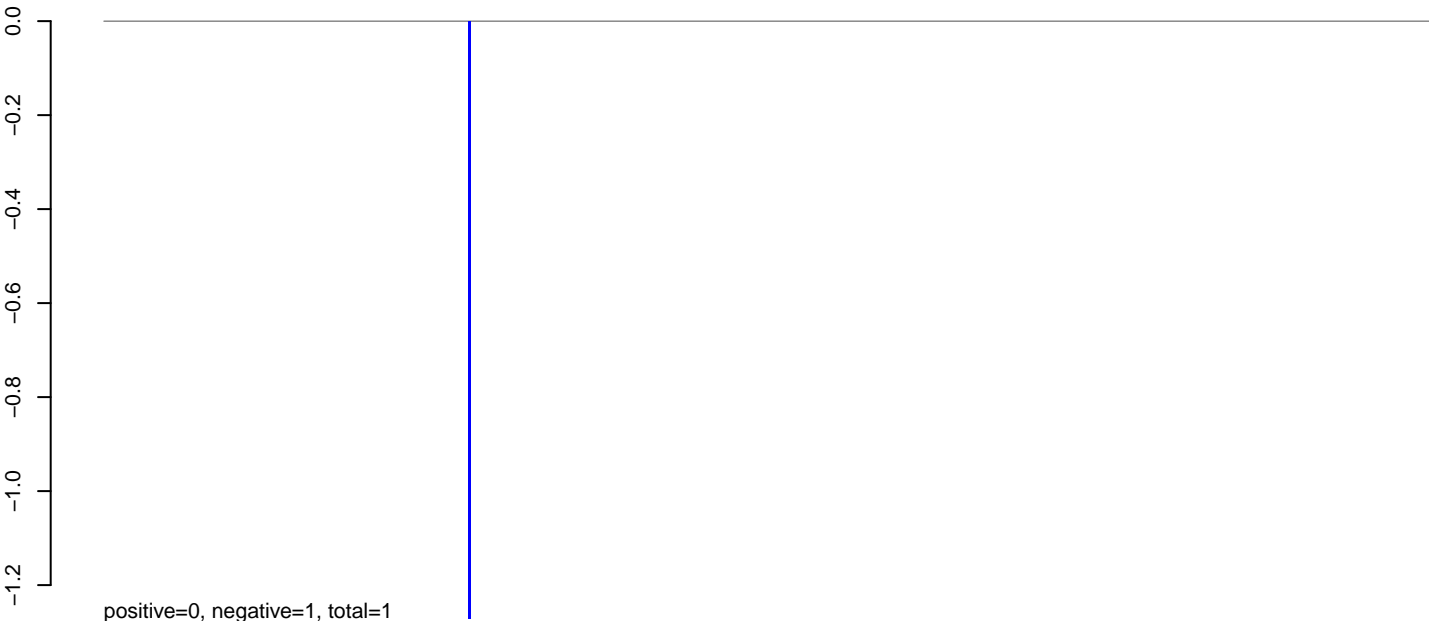


0 1000 2000 3000 4000

AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



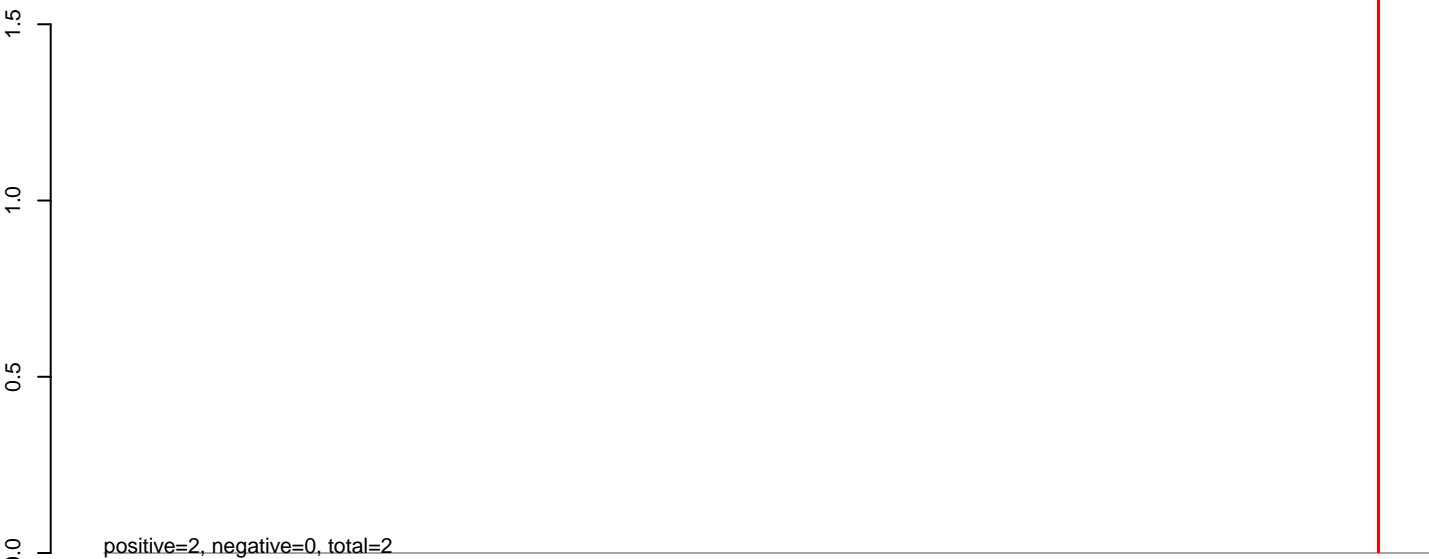
AeAlbo_C636_SINV_GFP_betaE.rep



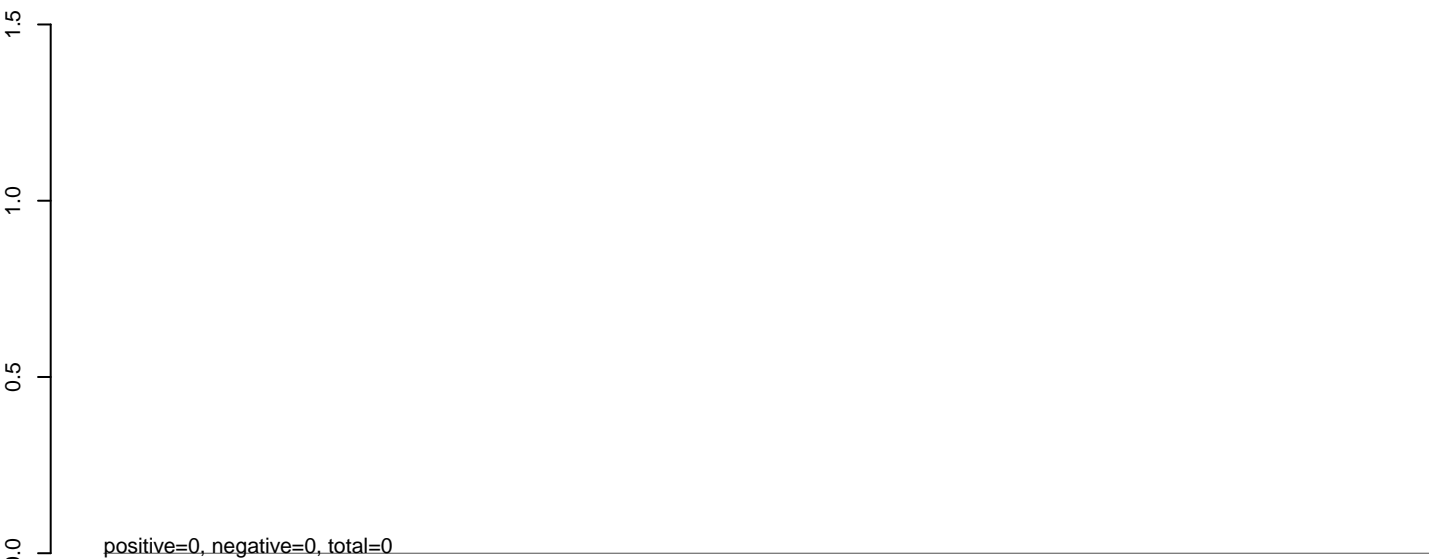
v size=25, length=11652, virus=AK1148553.1-Gambie_virus_putative_nucleoprotein_putative_small_transmembrane_protein_putative_g:1-11652

0 2000 4000 6000 8000 10000 12000

AeAlbo_C636_SINV_GFP_betaE.18_23.rep



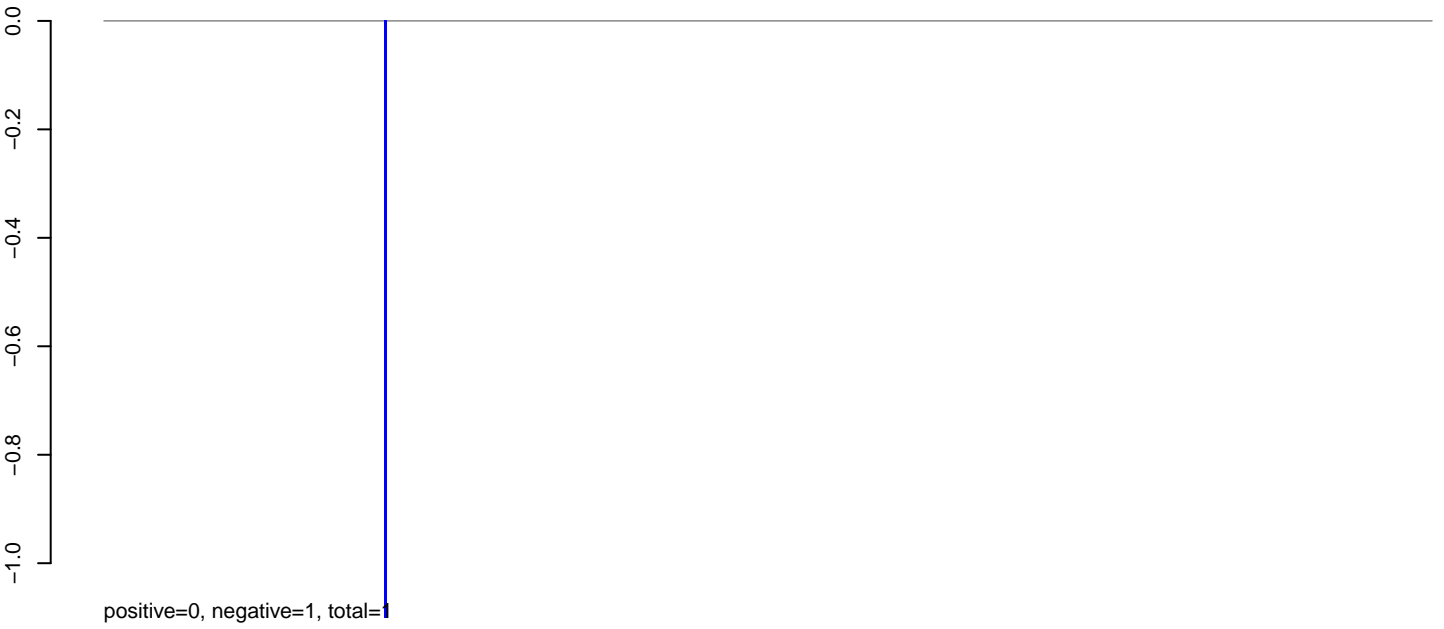
AeAlbo_C636_SINV_GFP_betaE.24_35.rep



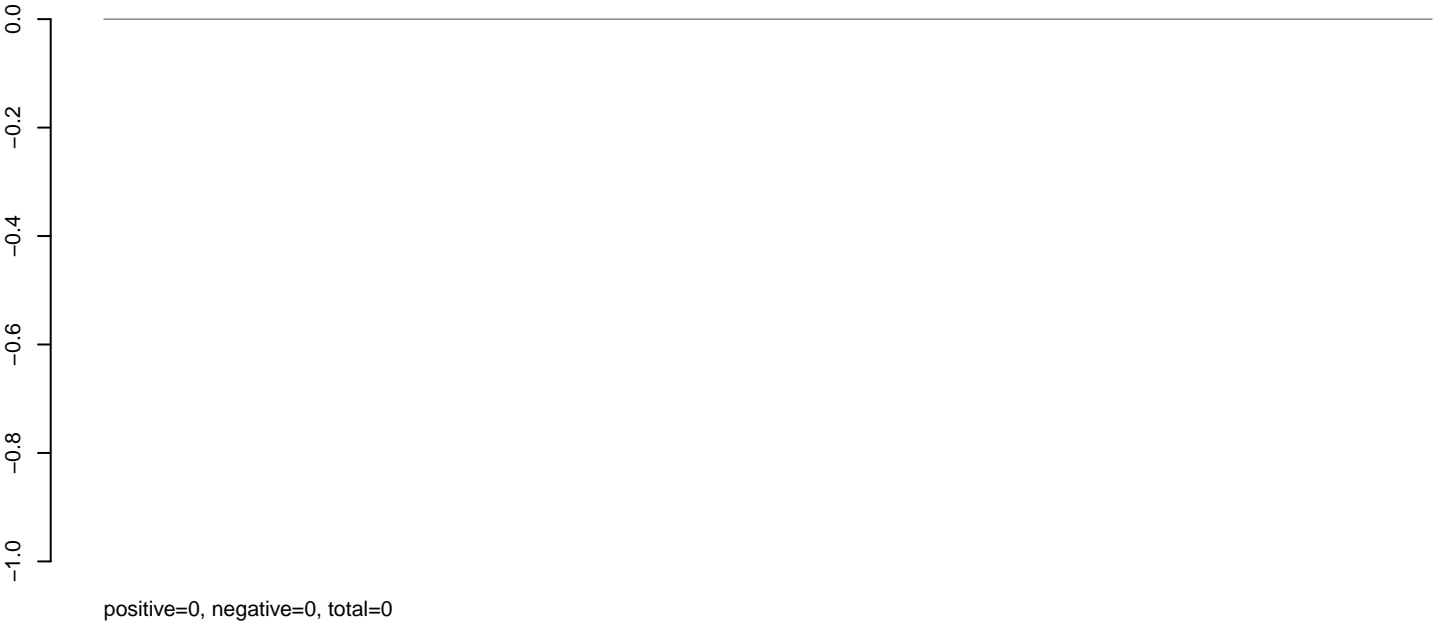
AeAlbo_C636_SINV_GFP_betaE.rep



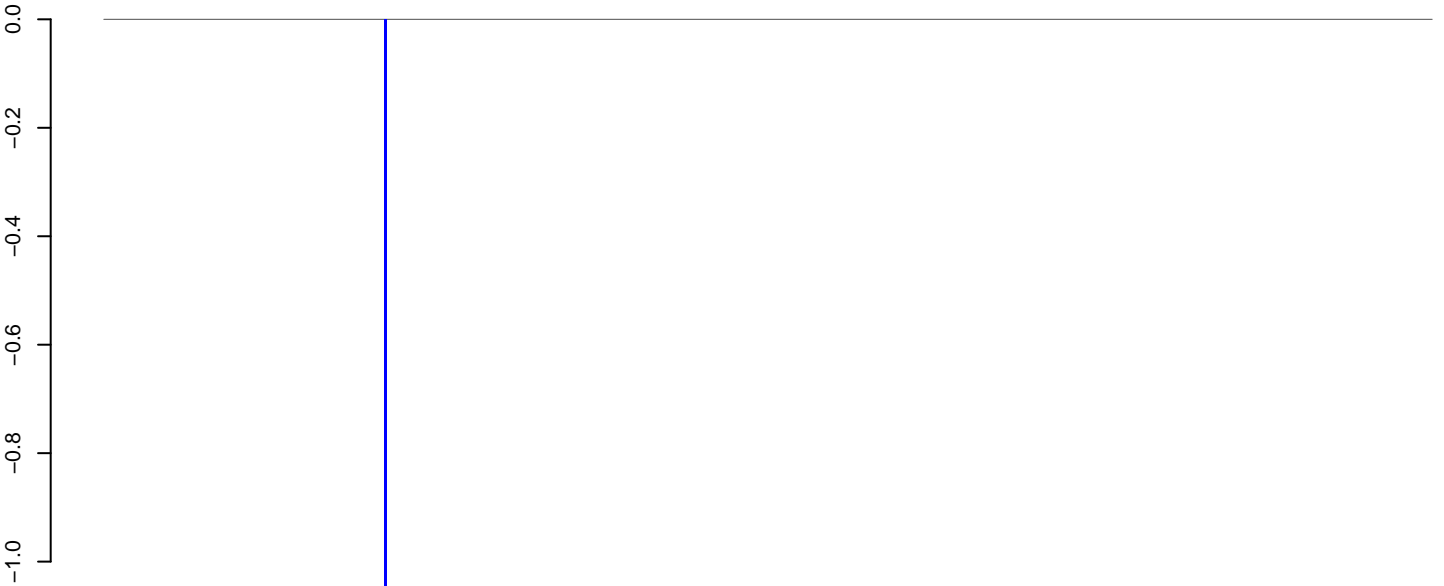
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



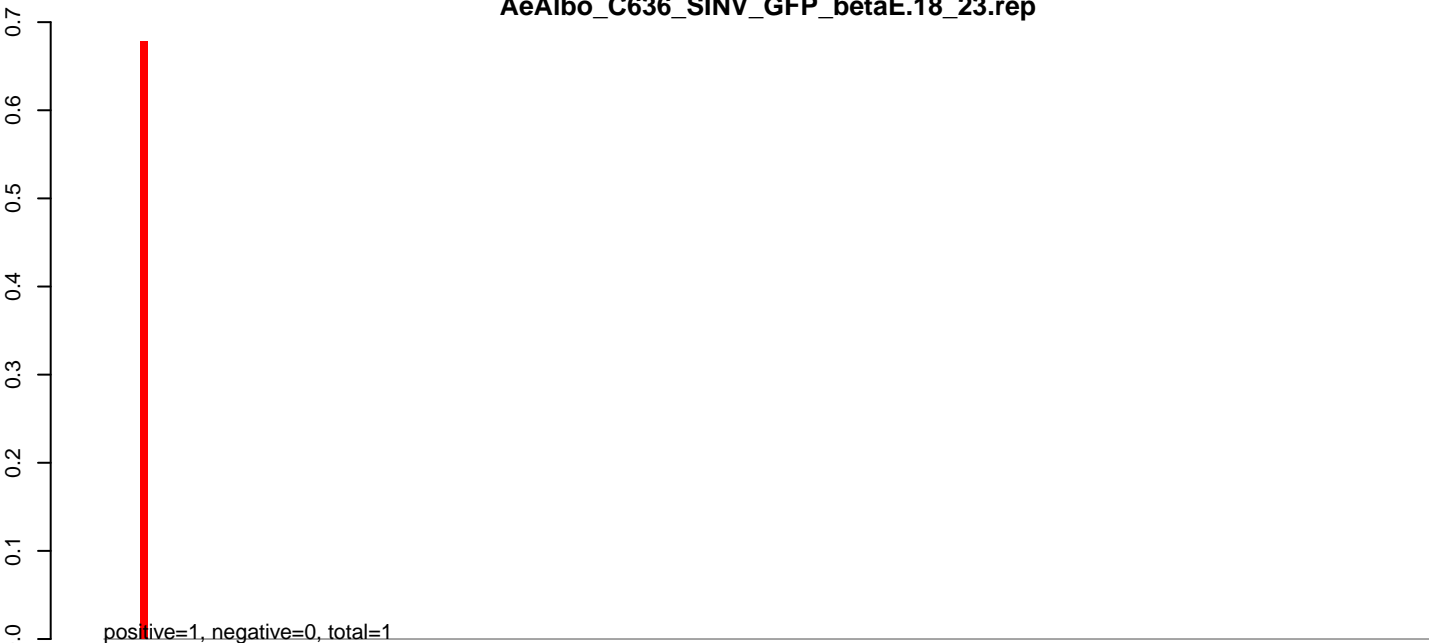
AeAlbo_C636_SINV_GFP_betaE.24_35.rep



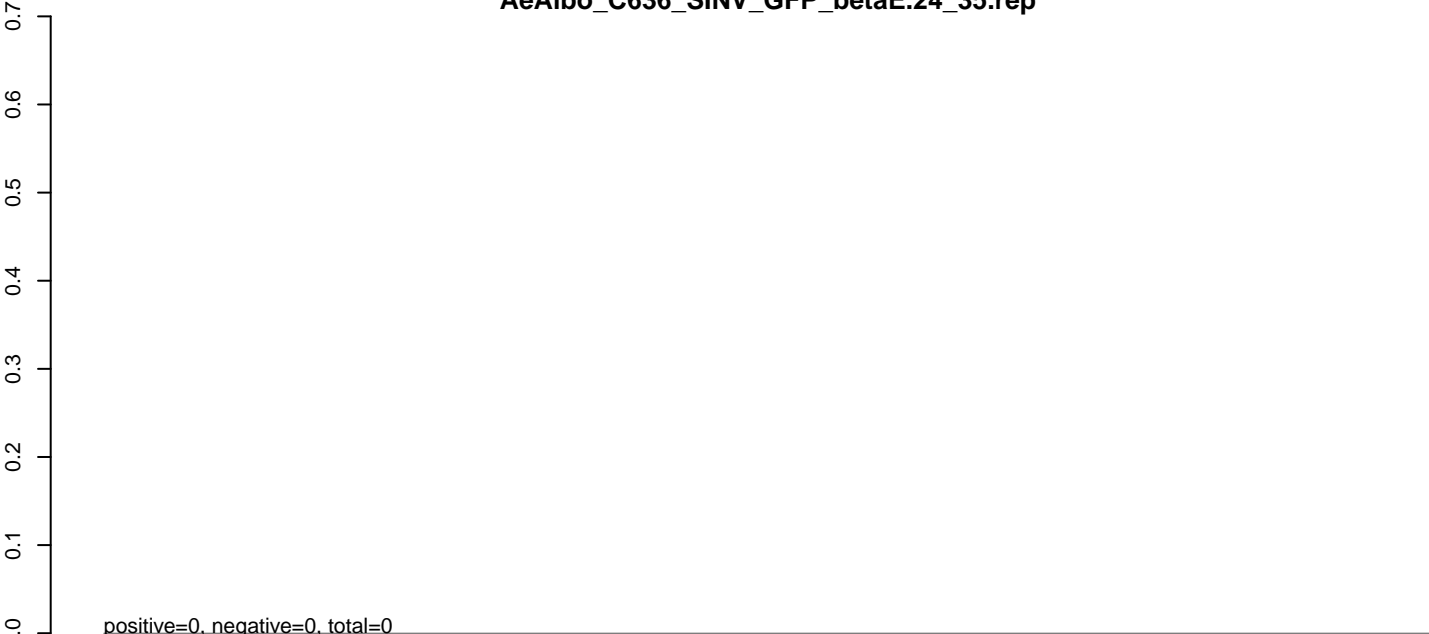
AeAlbo_C636_SINV_GFP_betaE.rep



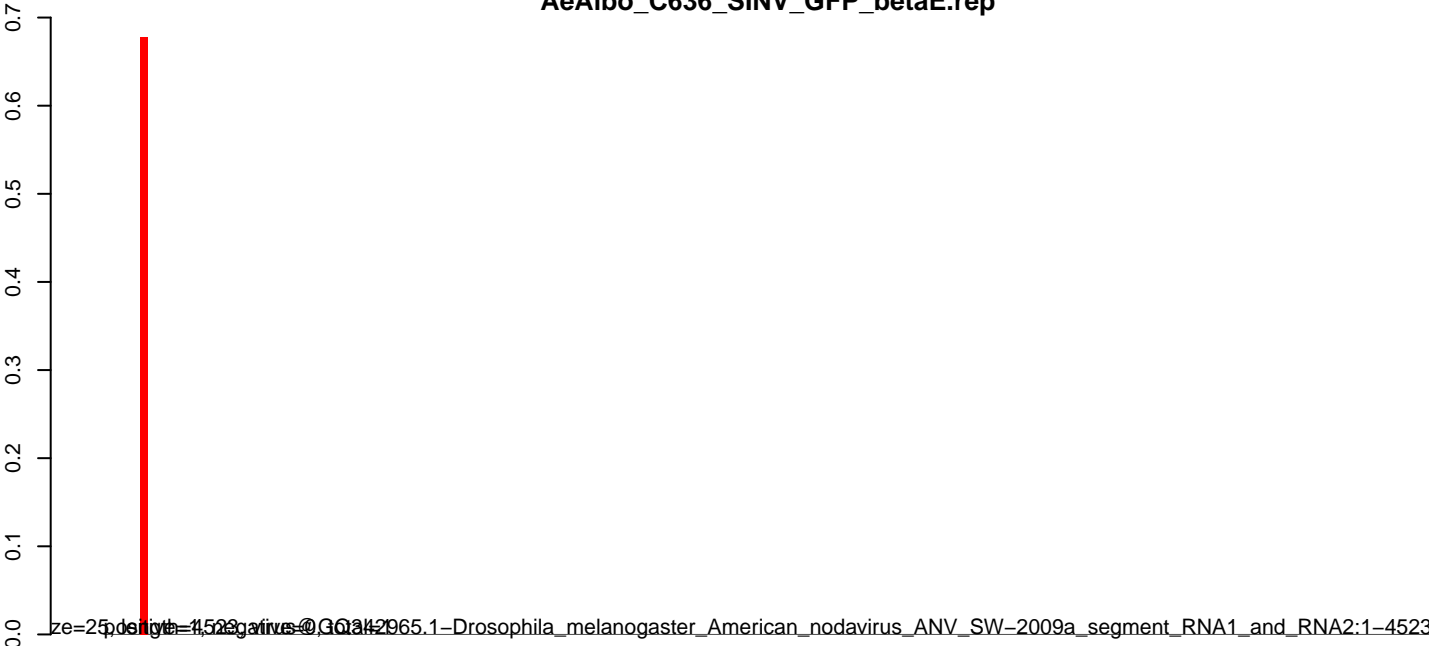
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



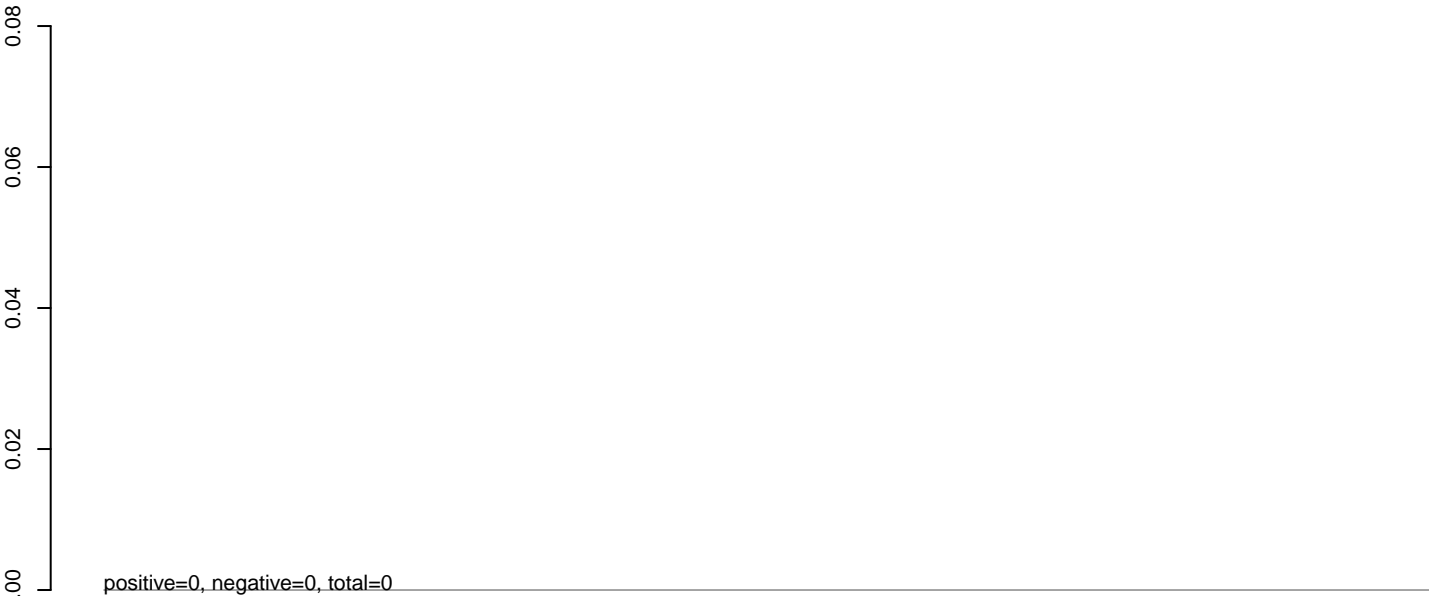
AeAlbo_C636_SINV_GFP_betaE.24_35.rep



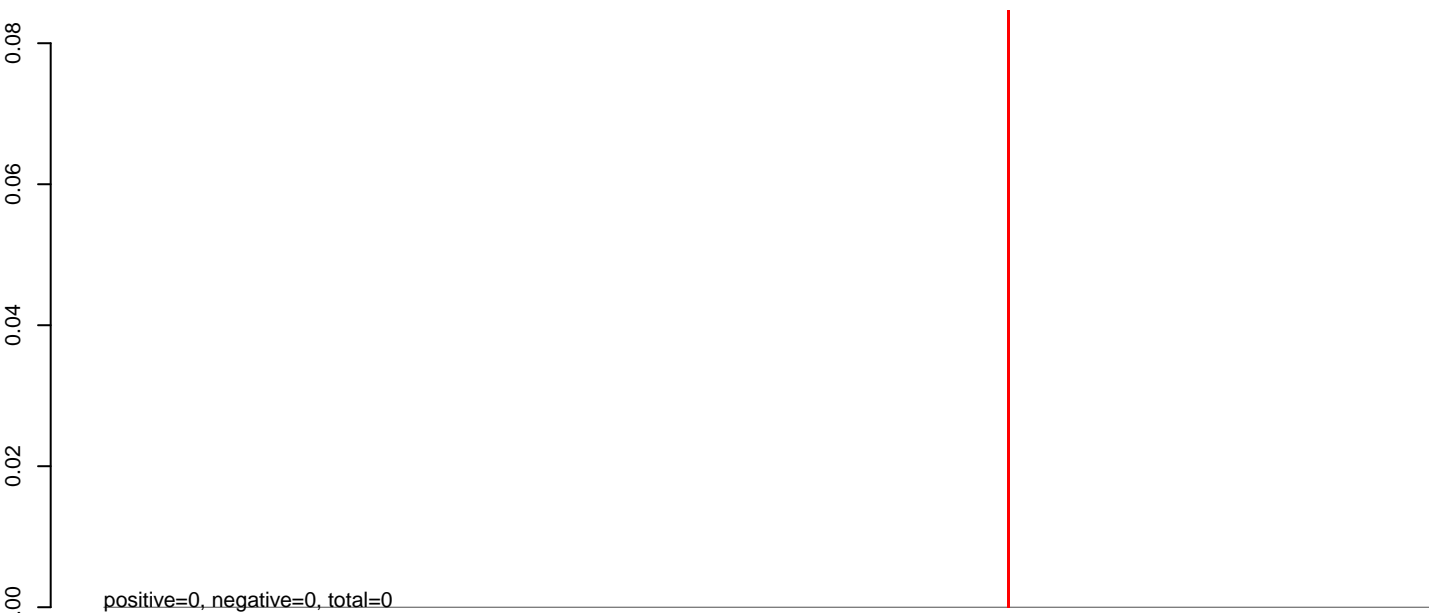
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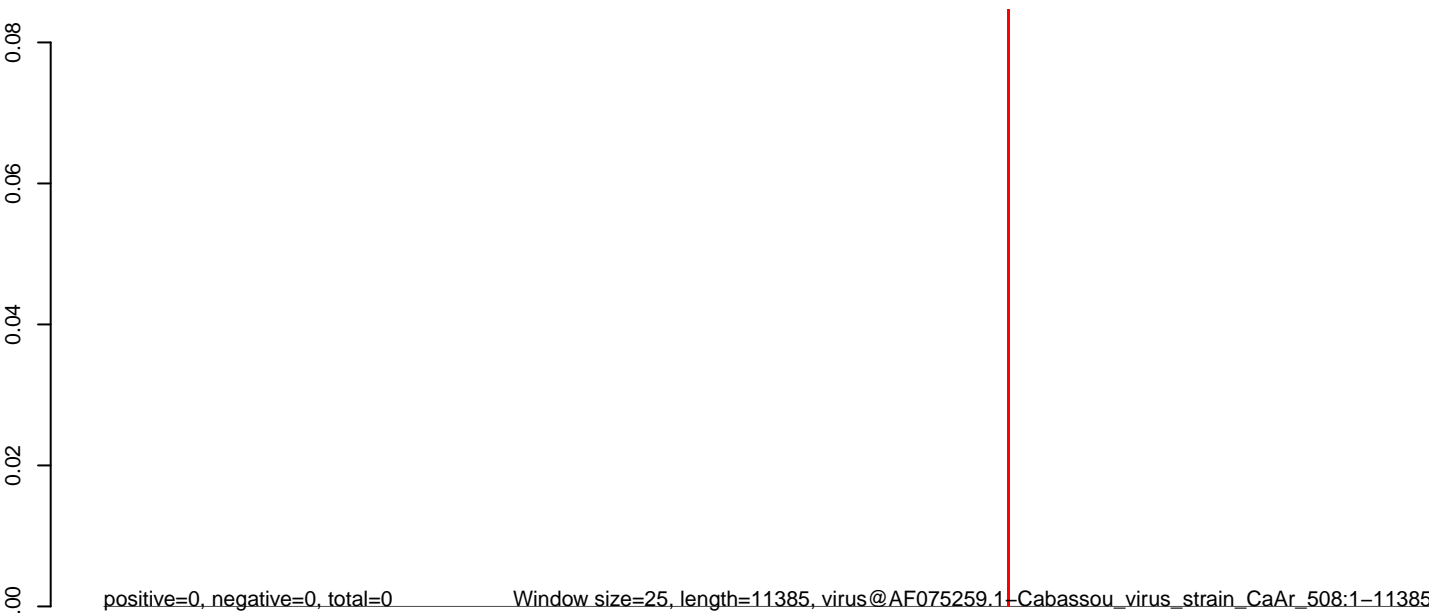
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



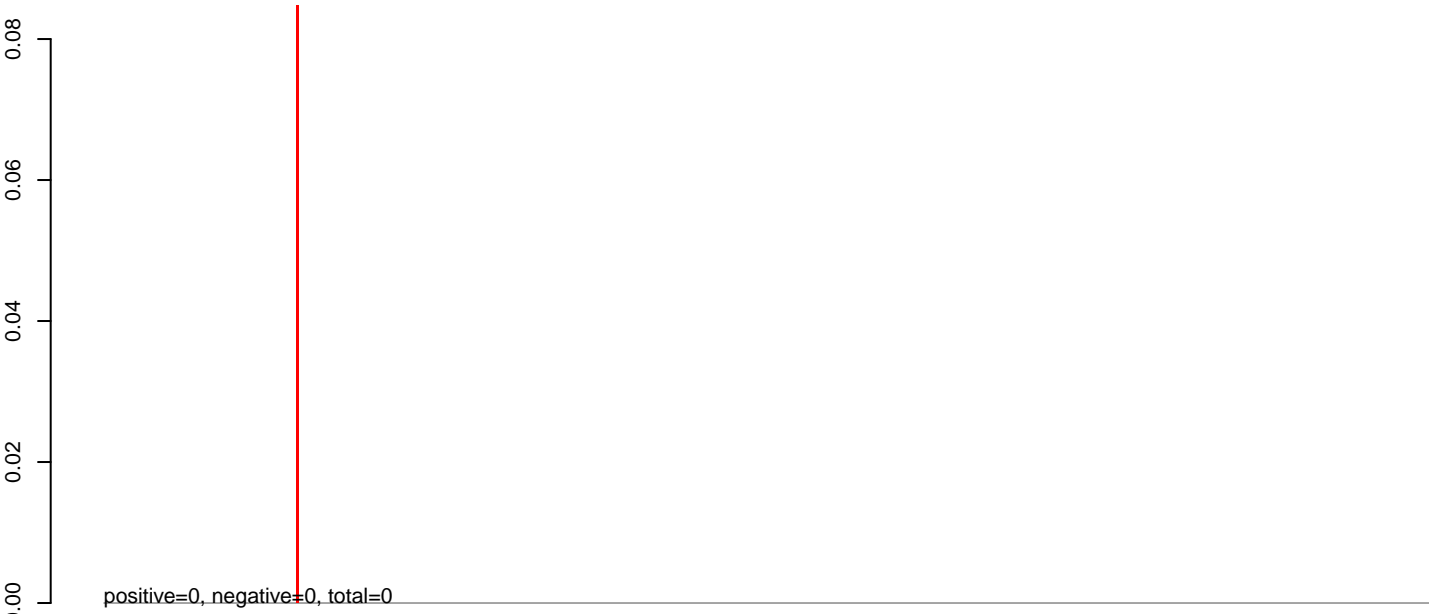
AeAlbo_C636_SINV_GFP_betaE.rep



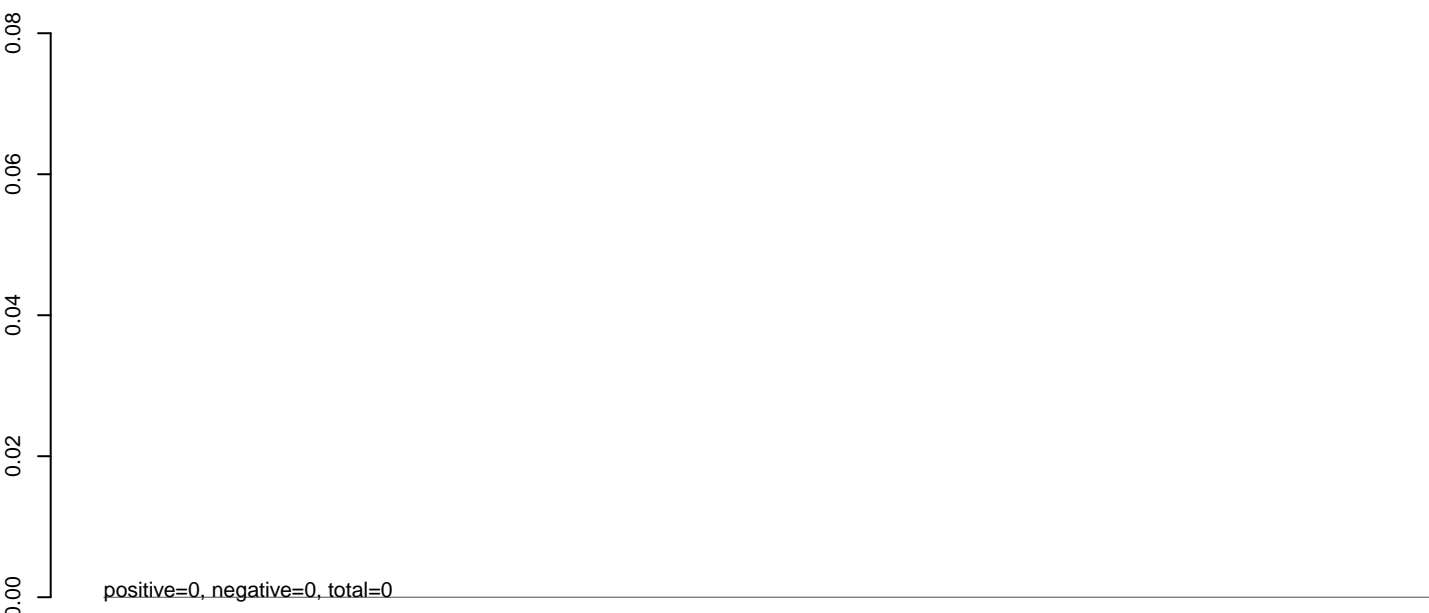
Window size=25, length=11385, virus@AF075259.1-Cabassou_virus_strain_CaAr_508:1-11385

0 2000 4000 6000 8000 10000

AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



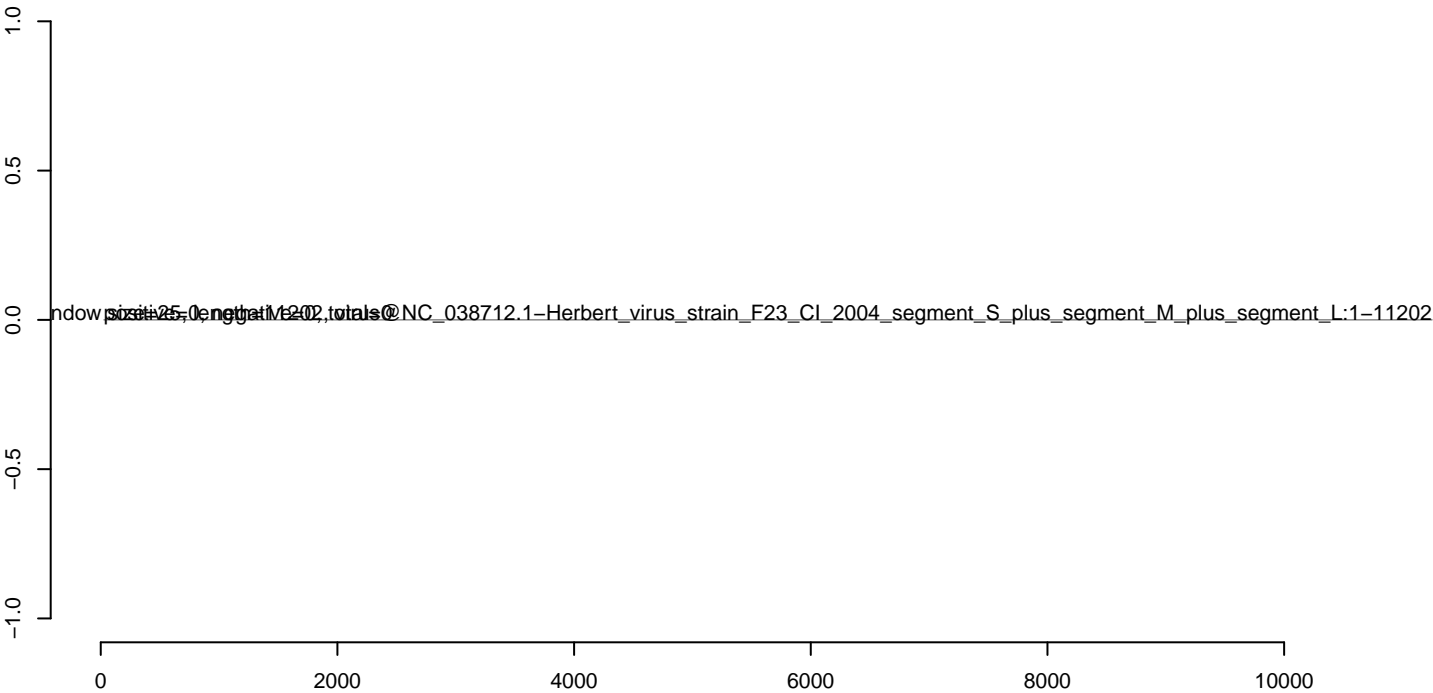
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



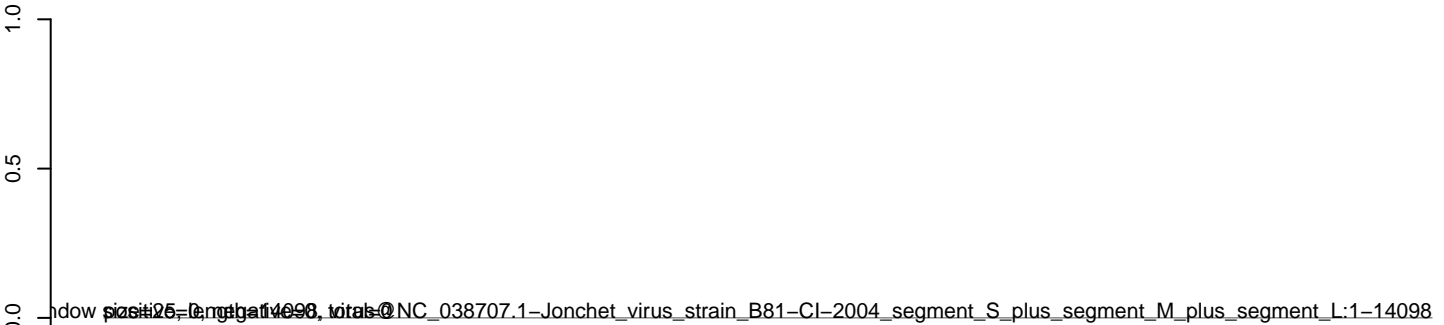
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



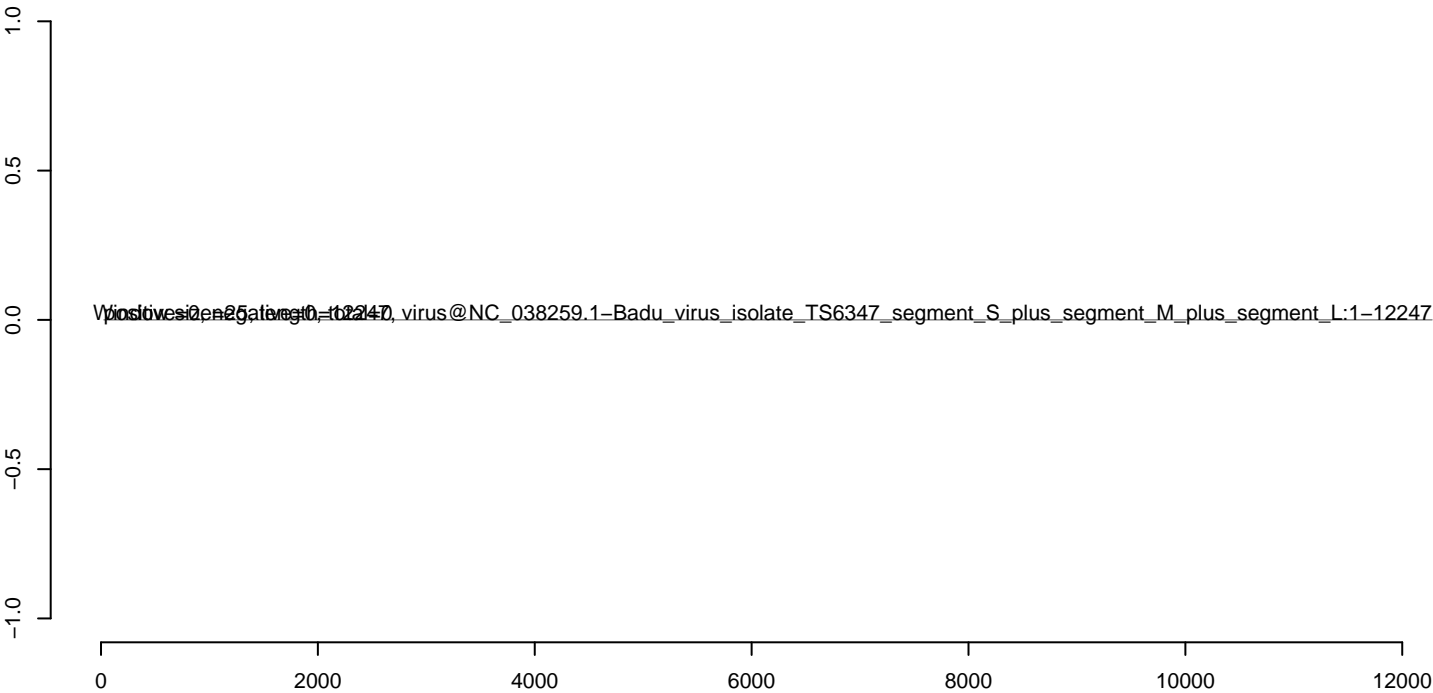
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AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



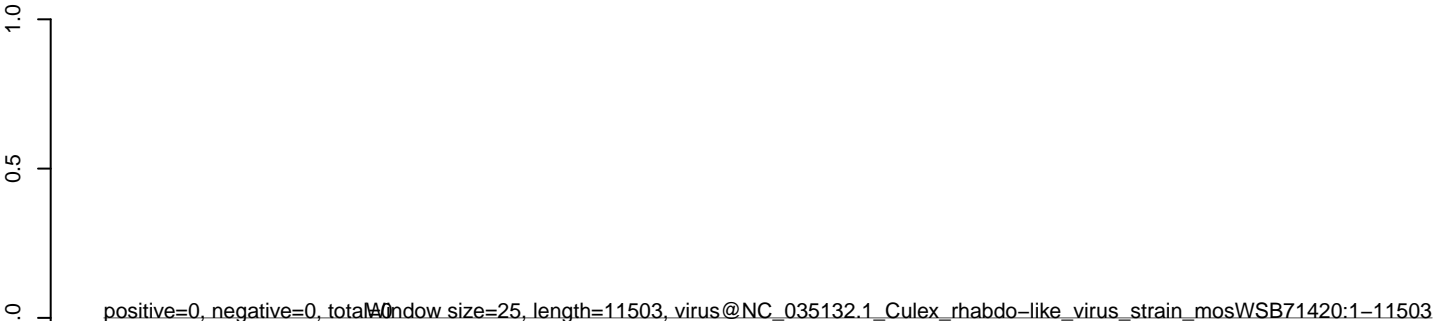
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



0 2000 4000 6000 8000 10000 12000

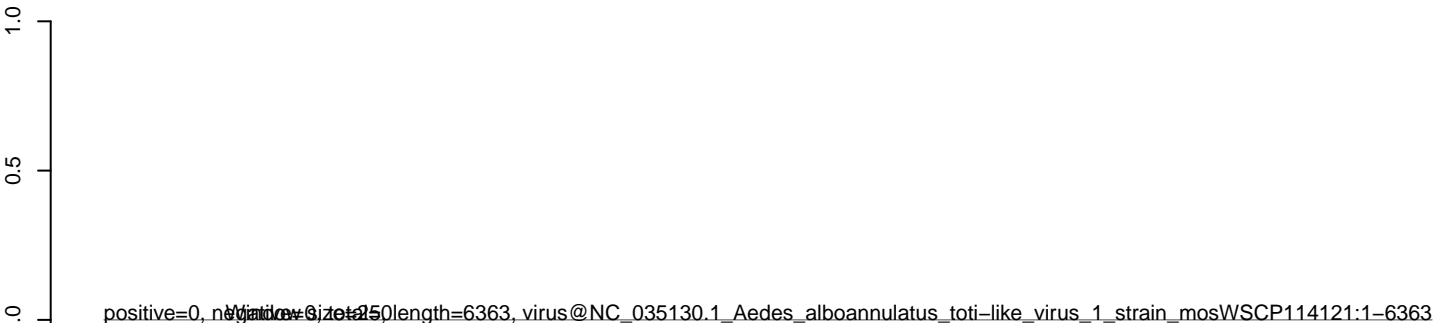
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



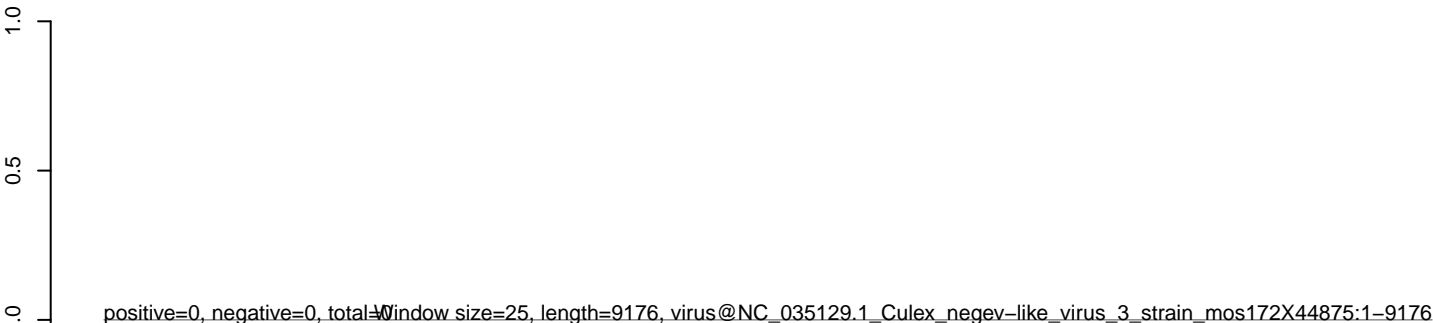
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



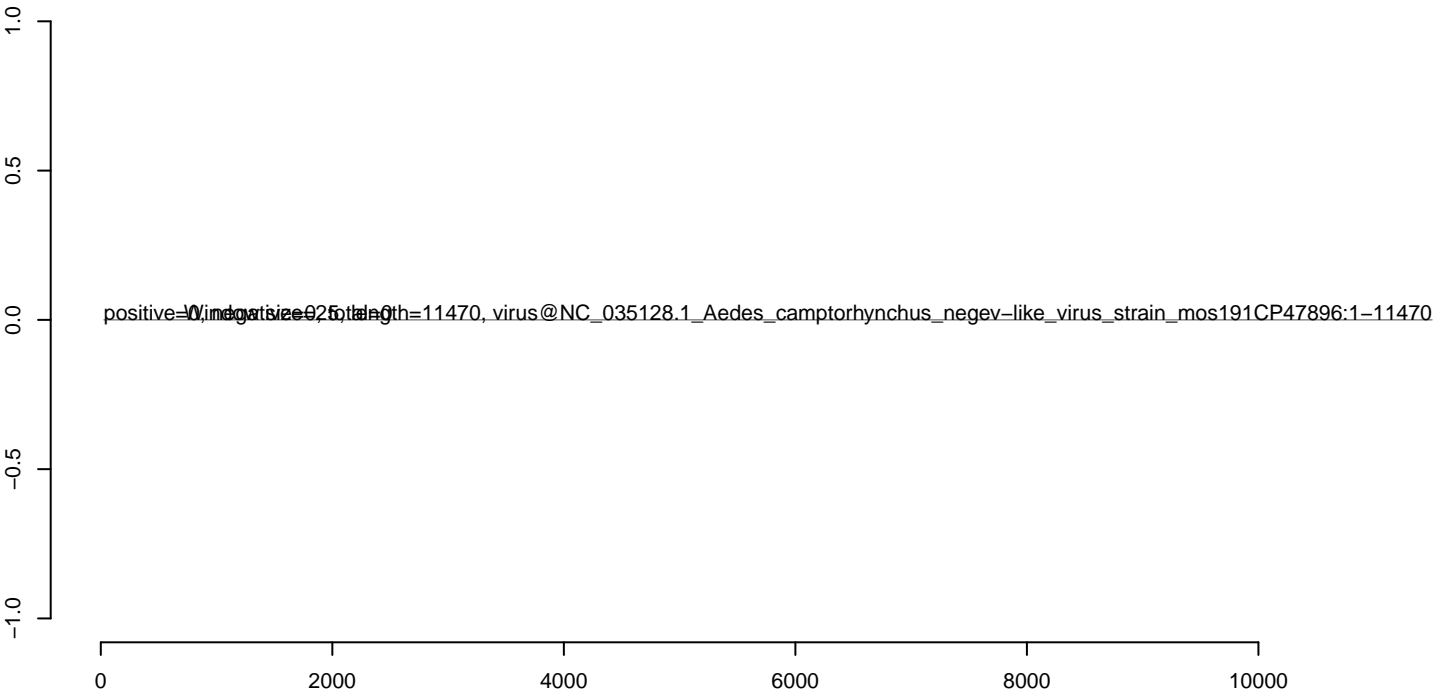
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



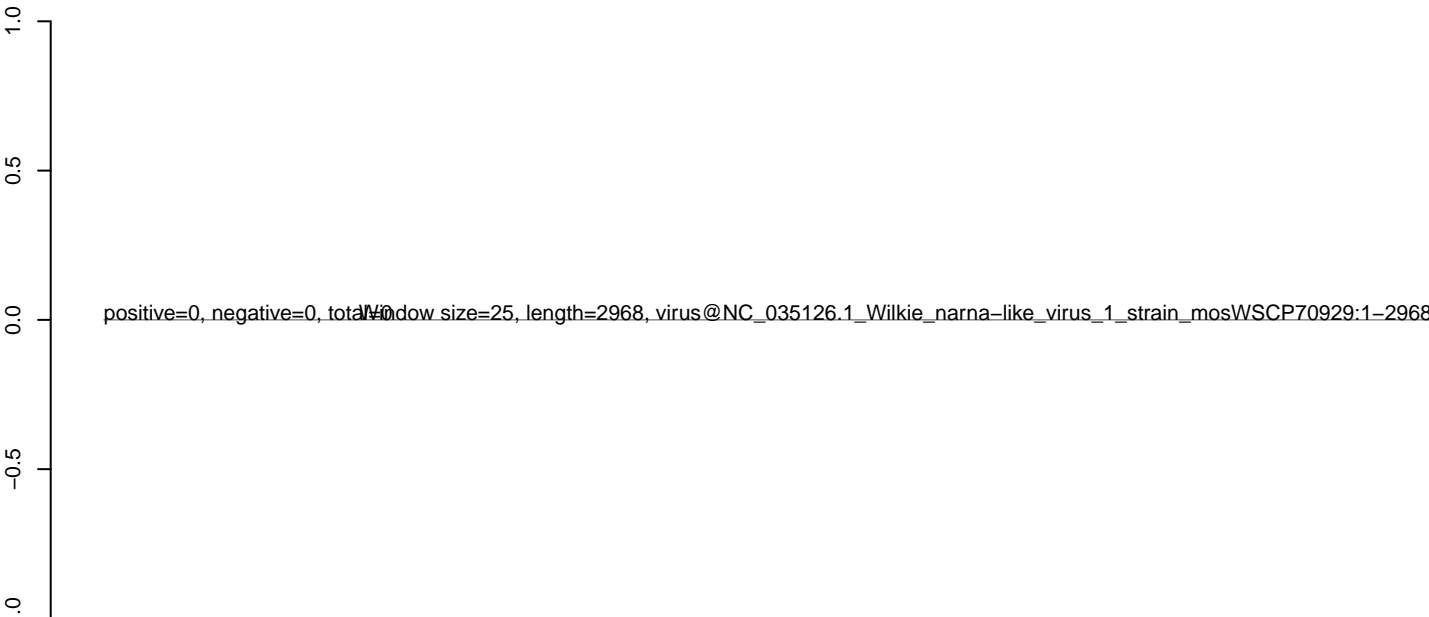
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



0 500 1000 1500 2000 2500 3000

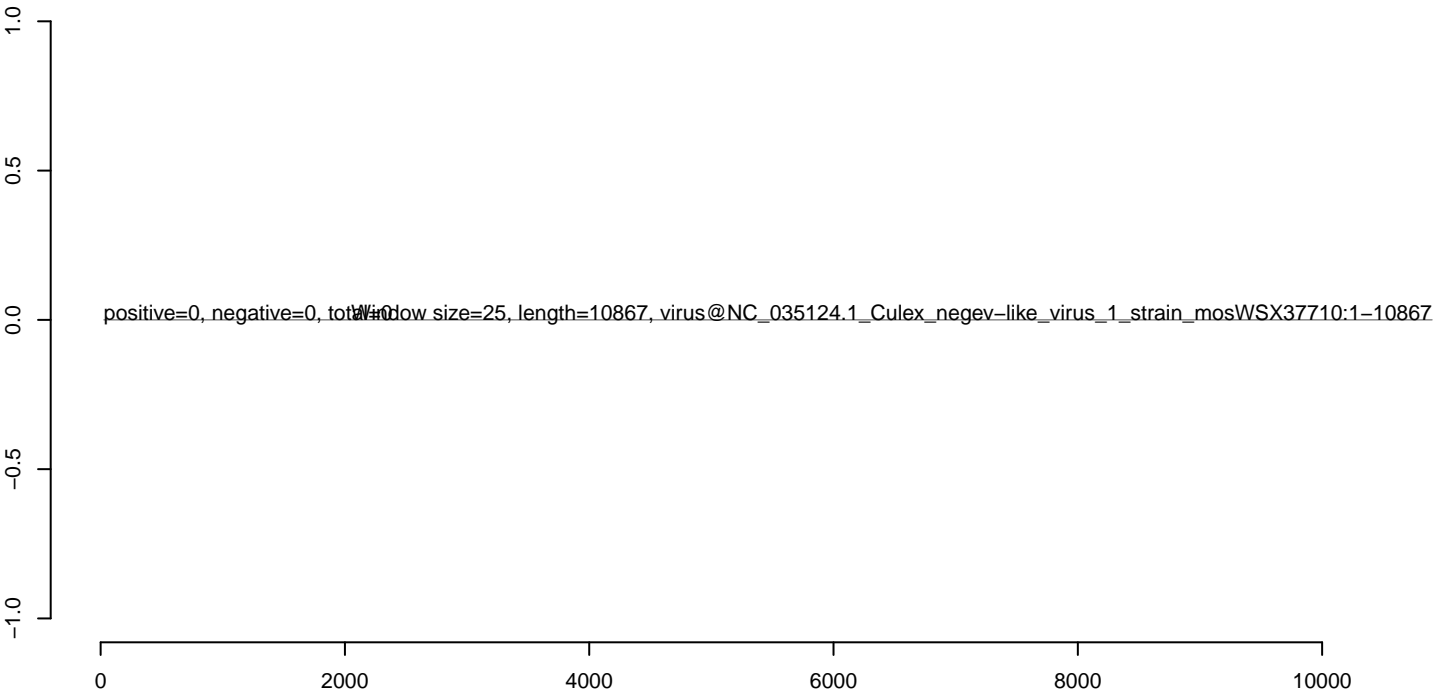
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



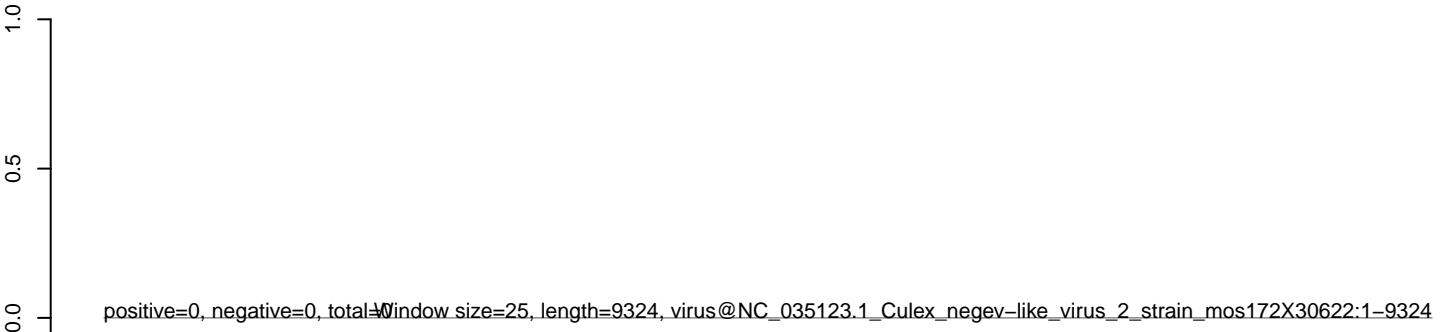
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



0 2000 4000 6000 8000

AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



0 500 1000 1500 2000 2500 3000

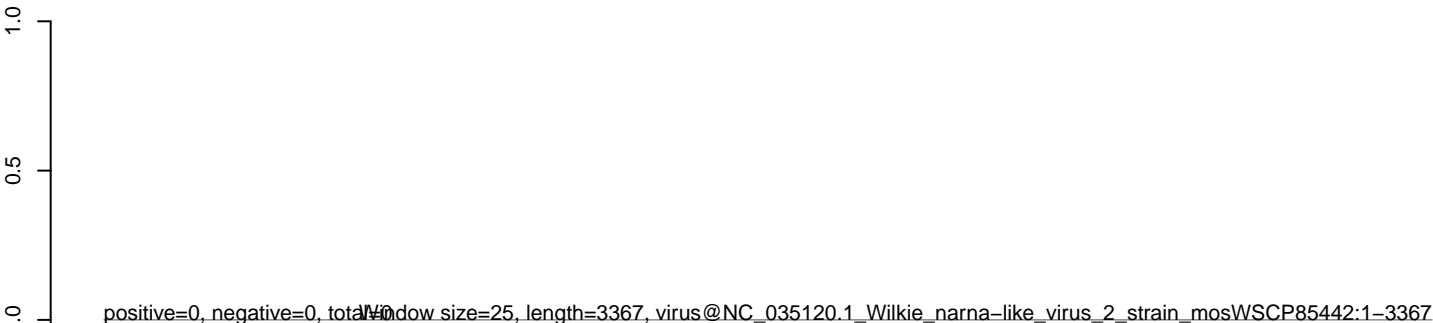
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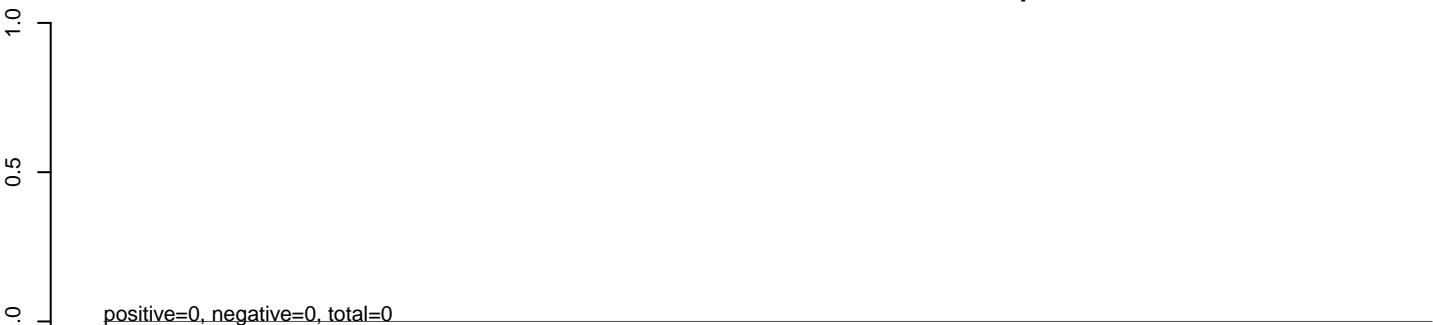
AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



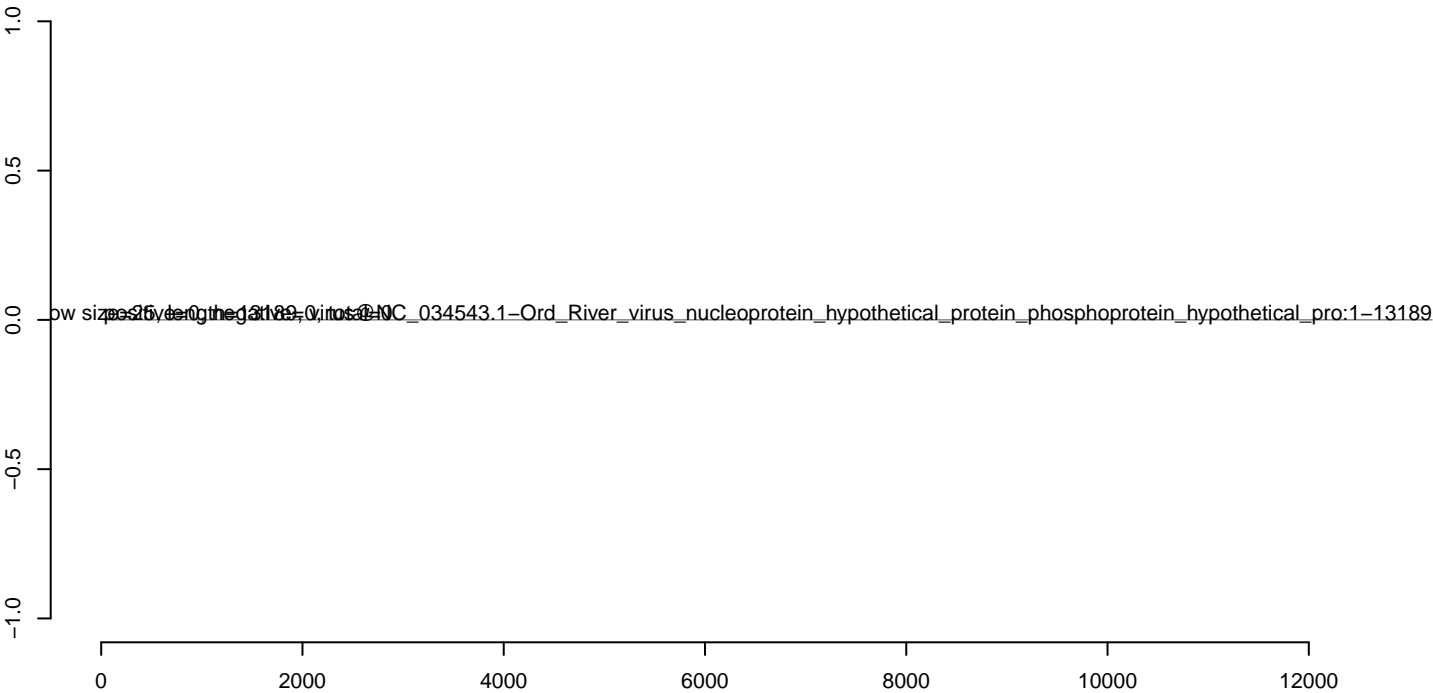
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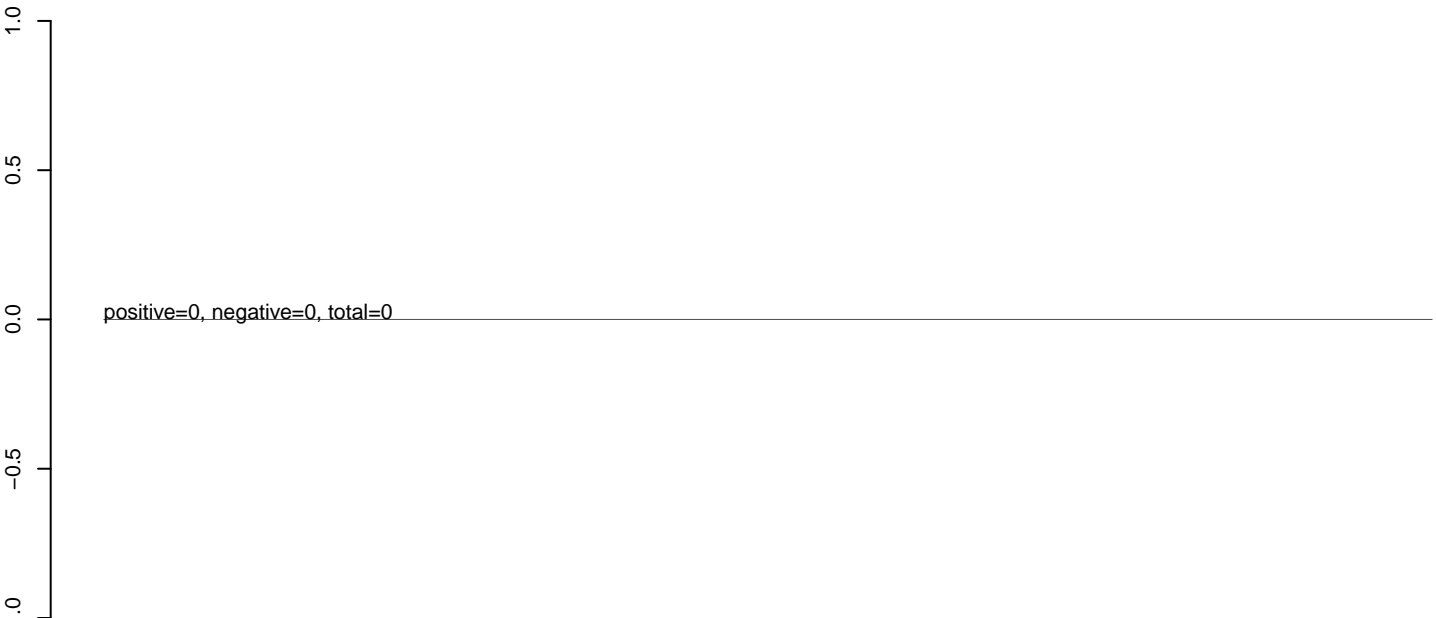
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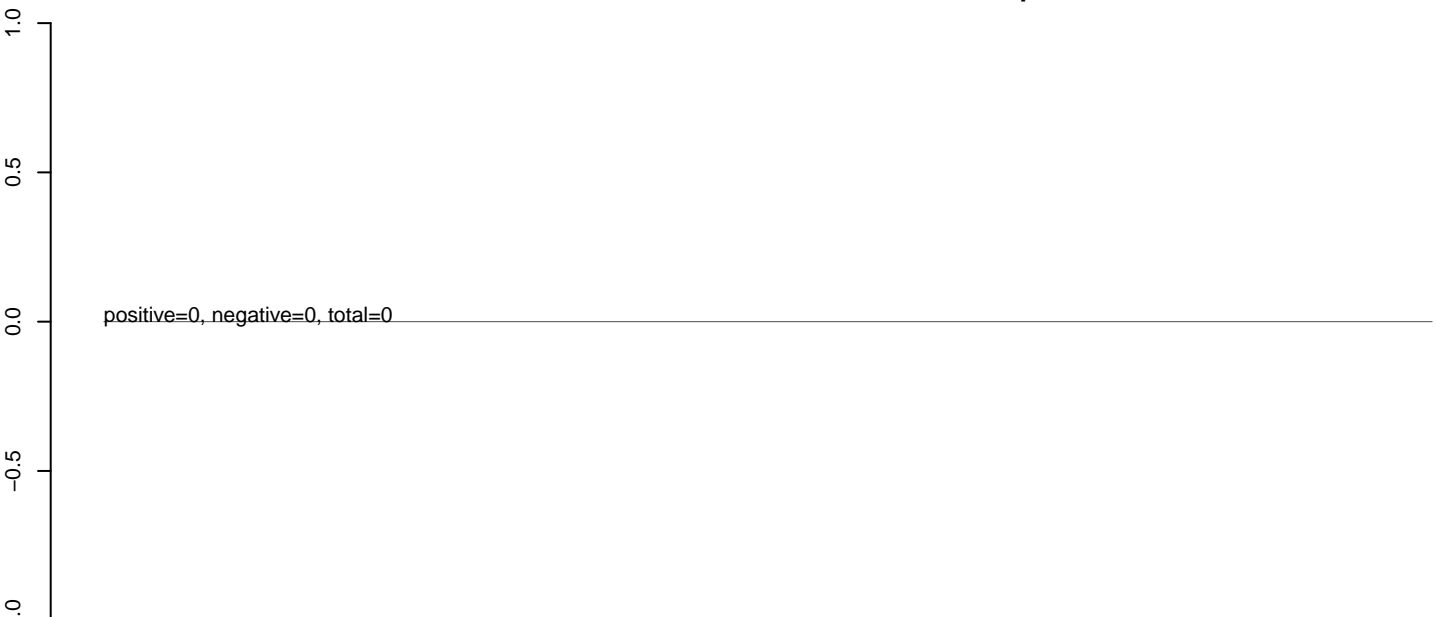
AeAlbo_C636_SINV_GFP_betaE.rep



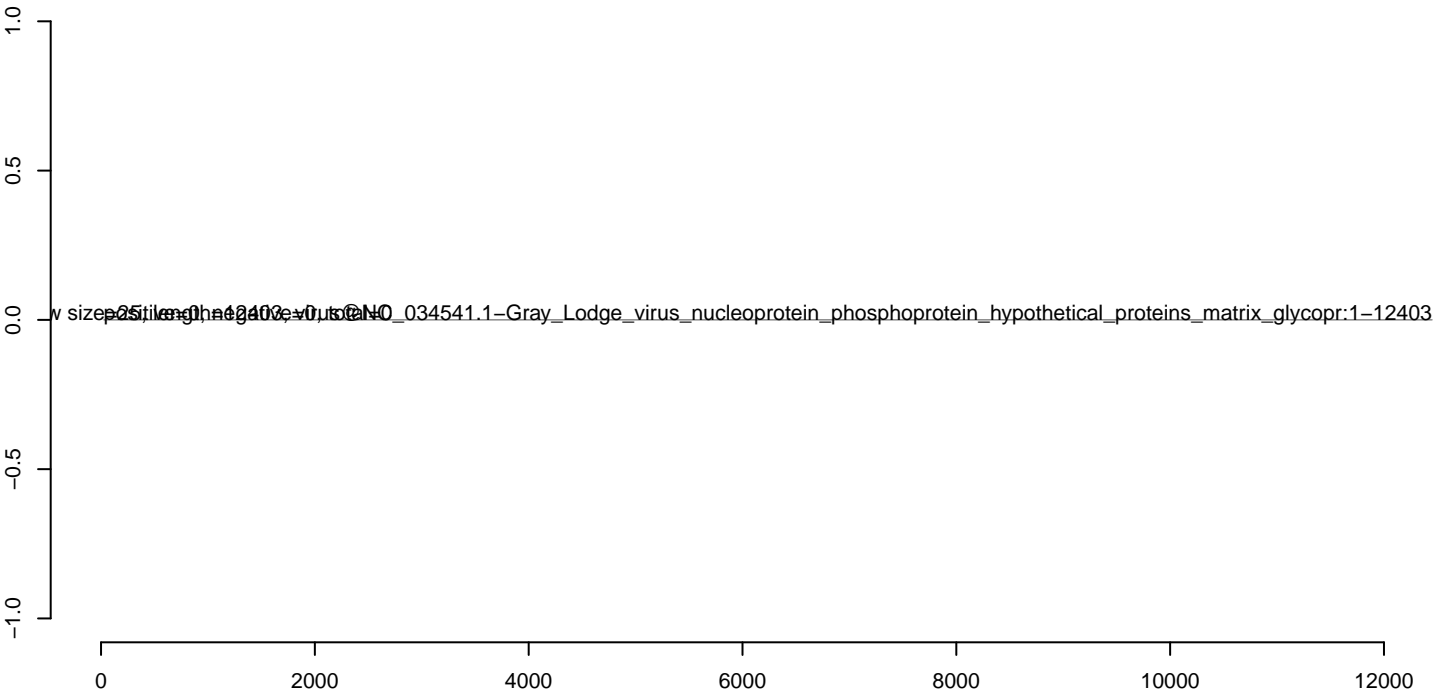
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



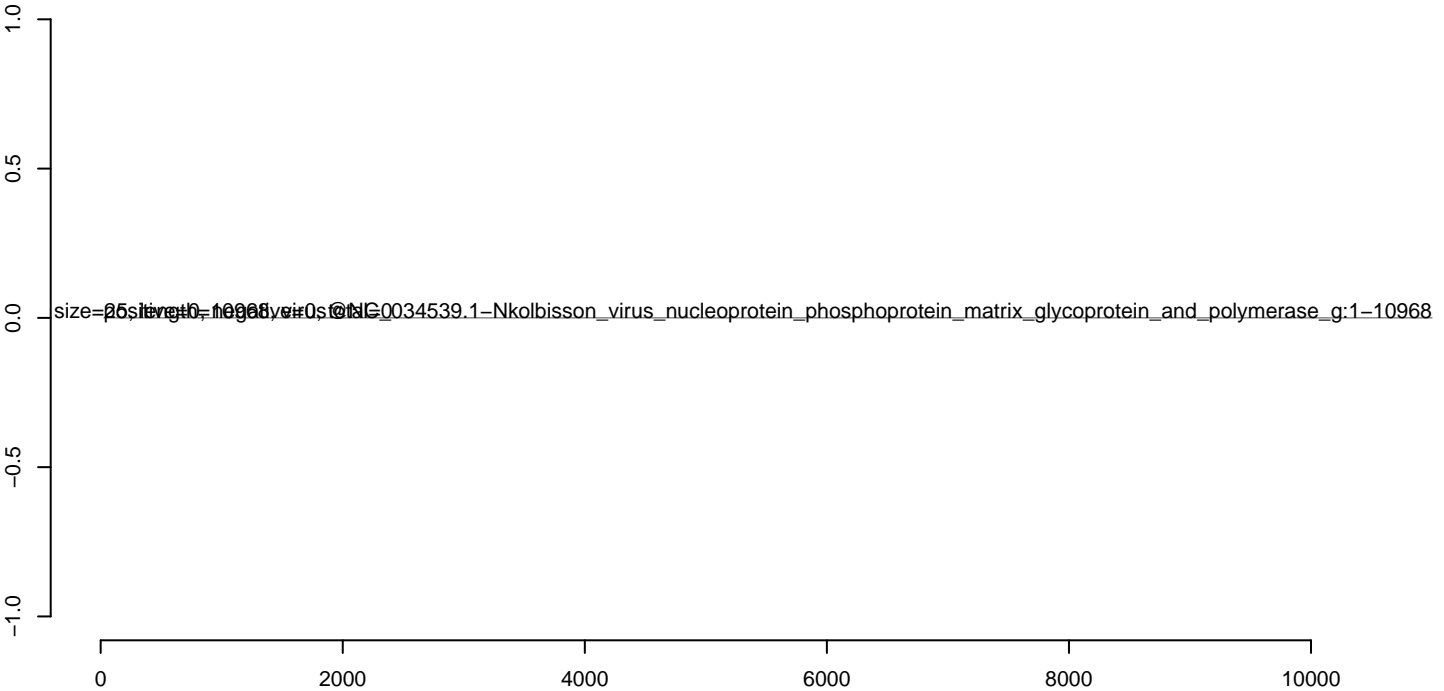
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



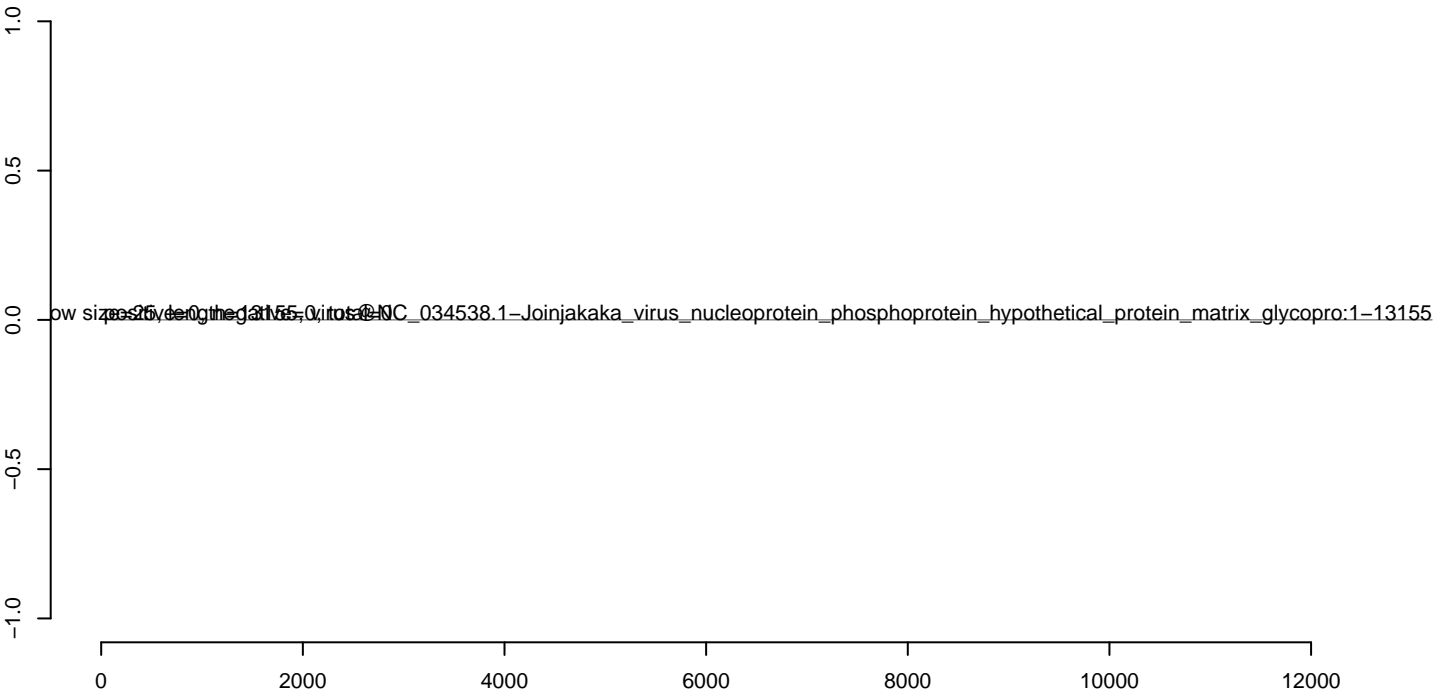
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



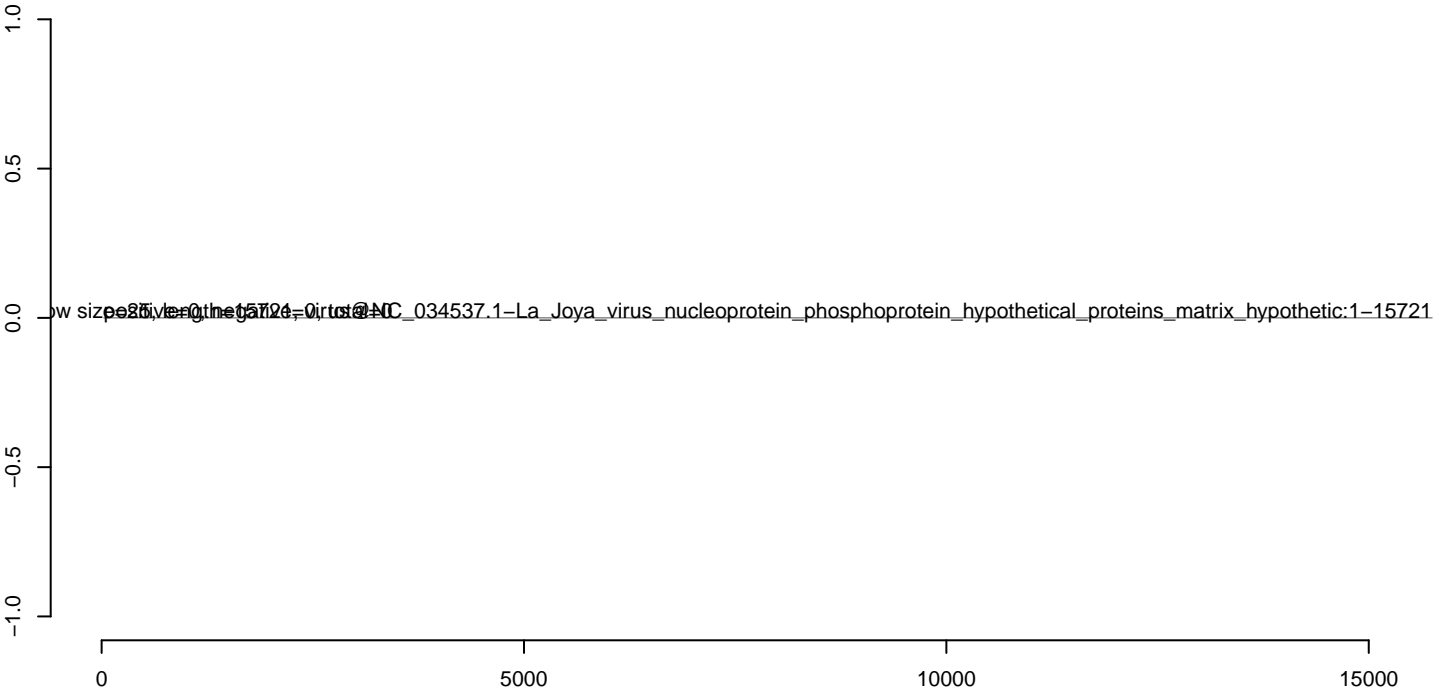
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



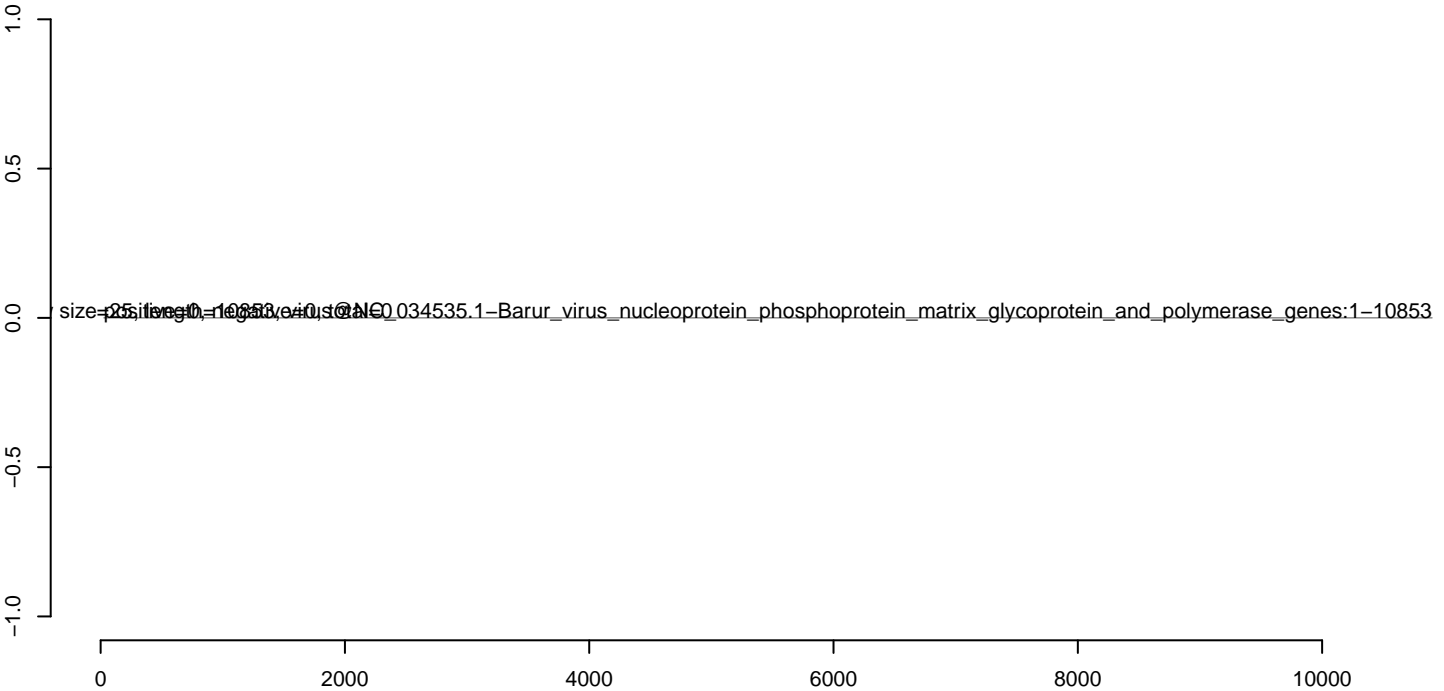
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AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



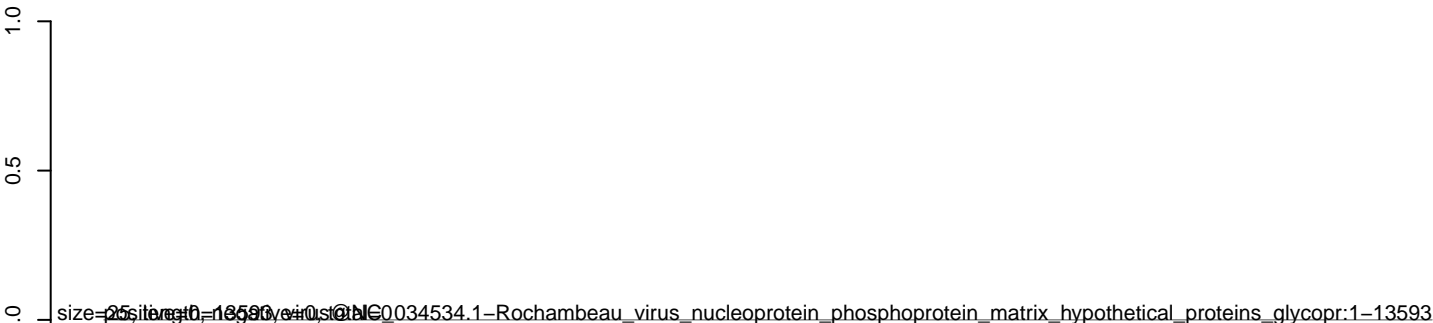
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



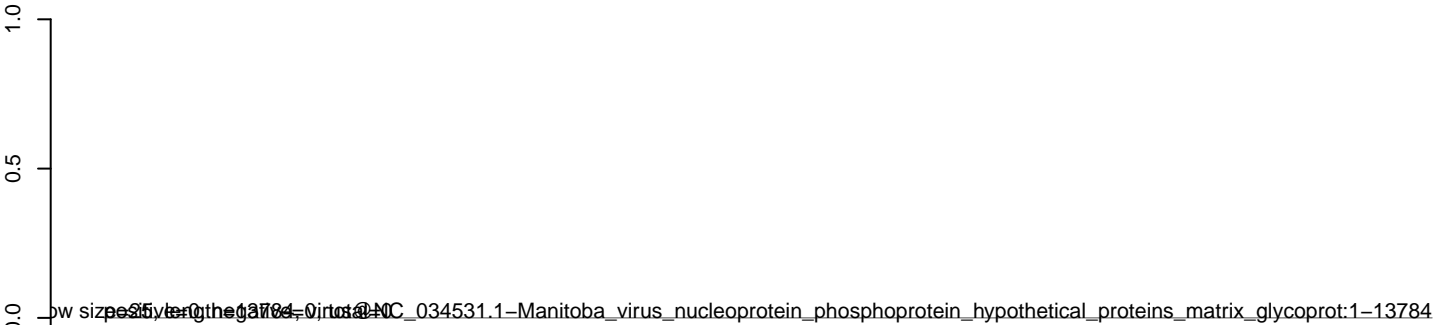
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



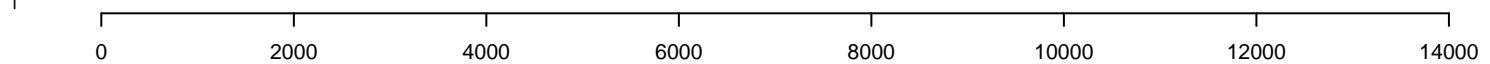
AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



low size of the genome of the virus. NC_034531.1-Manitoba_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycoprot:1-13784



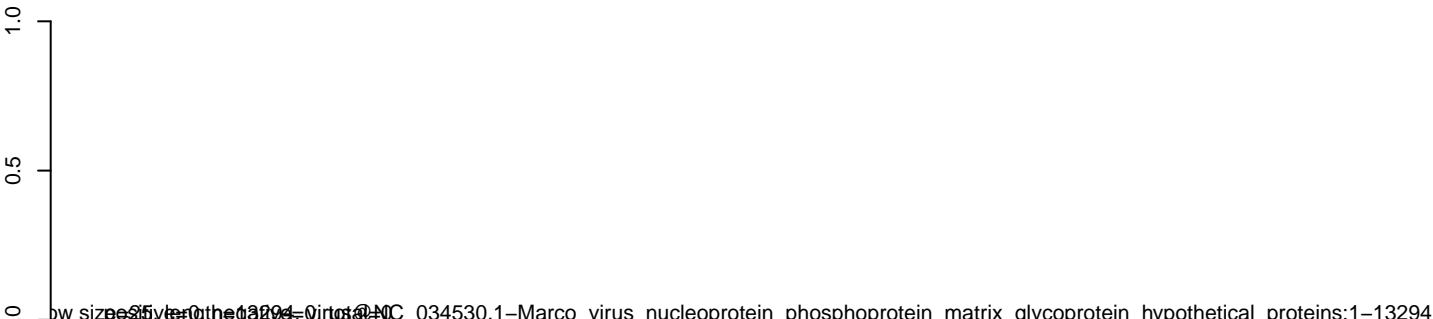
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



low size of the genome. The genome of the virus is approximately 12.5 kb. The genome contains a single open reading frame (ORF) that encodes a polyprotein of approximately 13.294 amino acids. The polyprotein is cleaved into several proteins, including the nucleoprotein, phosphoprotein, matrix protein, glycoprotein, and several hypothetical proteins. The genome also contains several non-coding regions, including the 5' and 3' untranslated regions (UTRs) and the intergenic region (IGR).

0 2000 4000 6000 8000 10000 12000

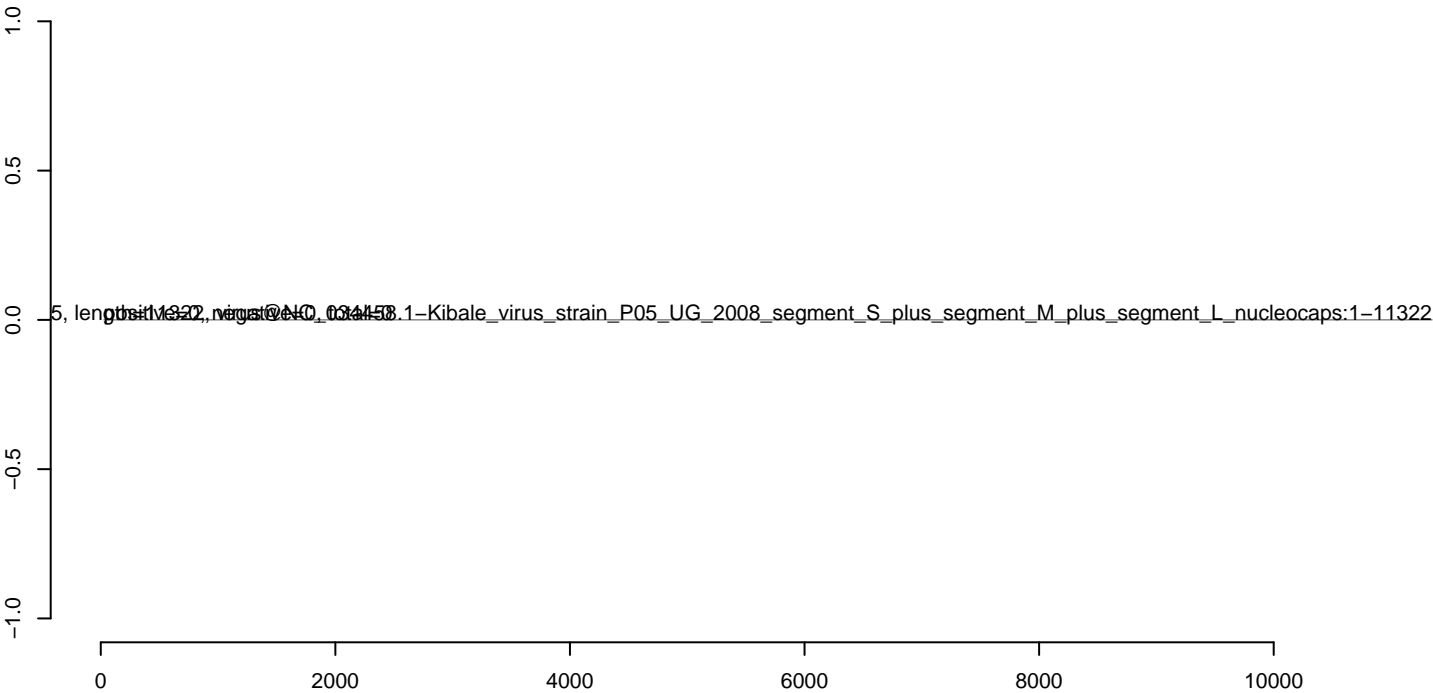
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



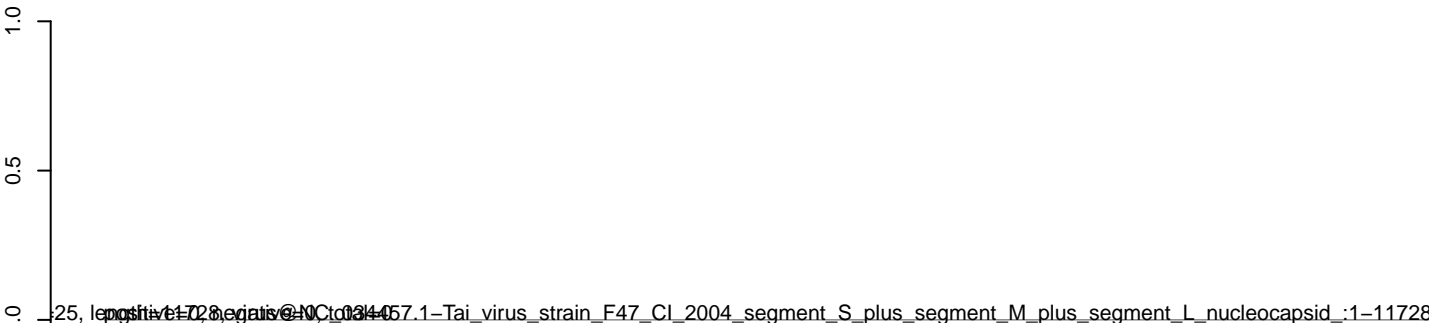
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



0 2000 4000 6000 8000 10000 12000

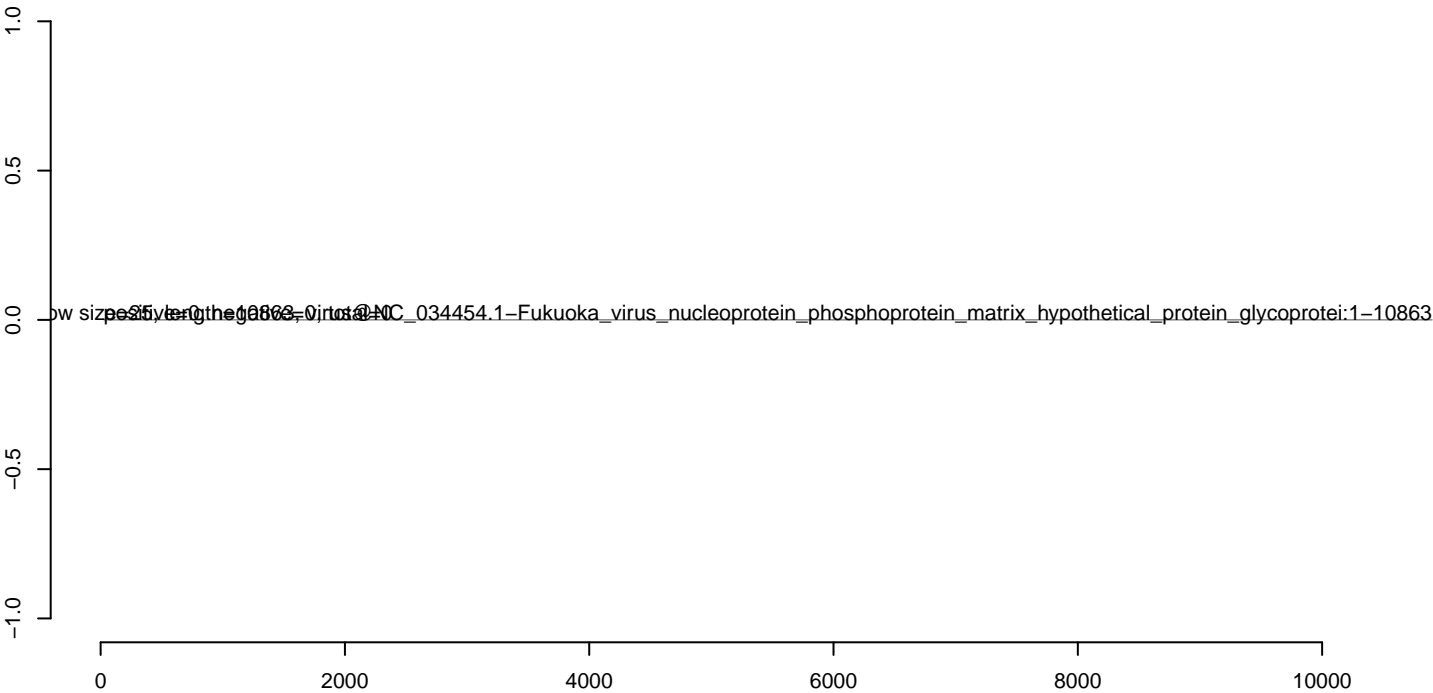
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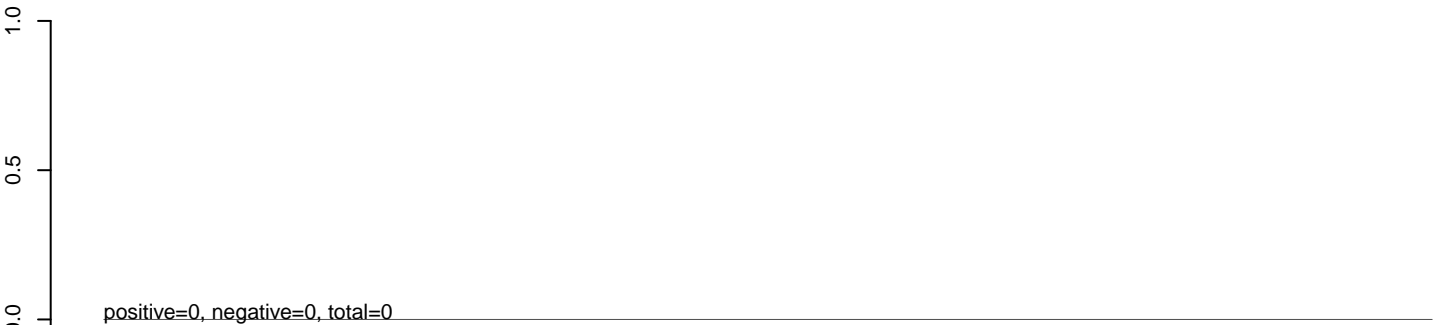
AeAlbo_C636_SINV_GFP_betaE.24_35.rep



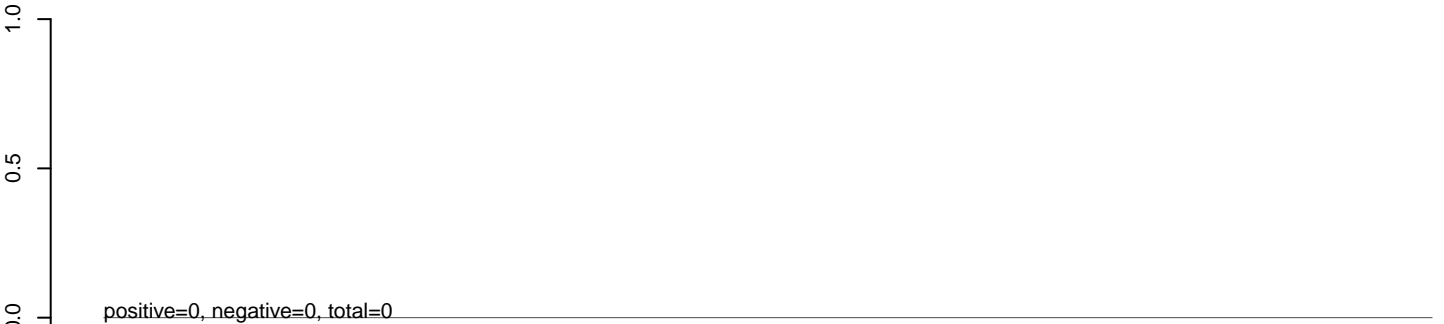
AeAlbo_C636_SINV_GFP_betaE.rep



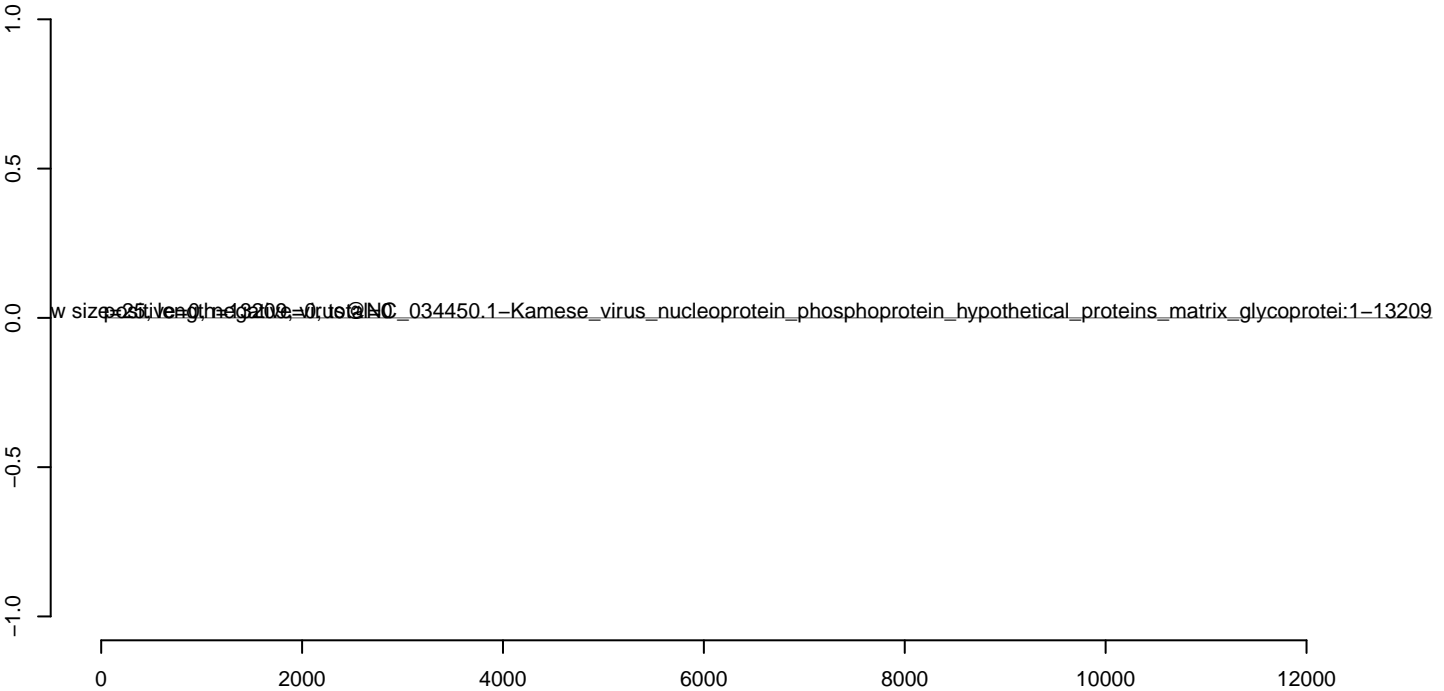
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



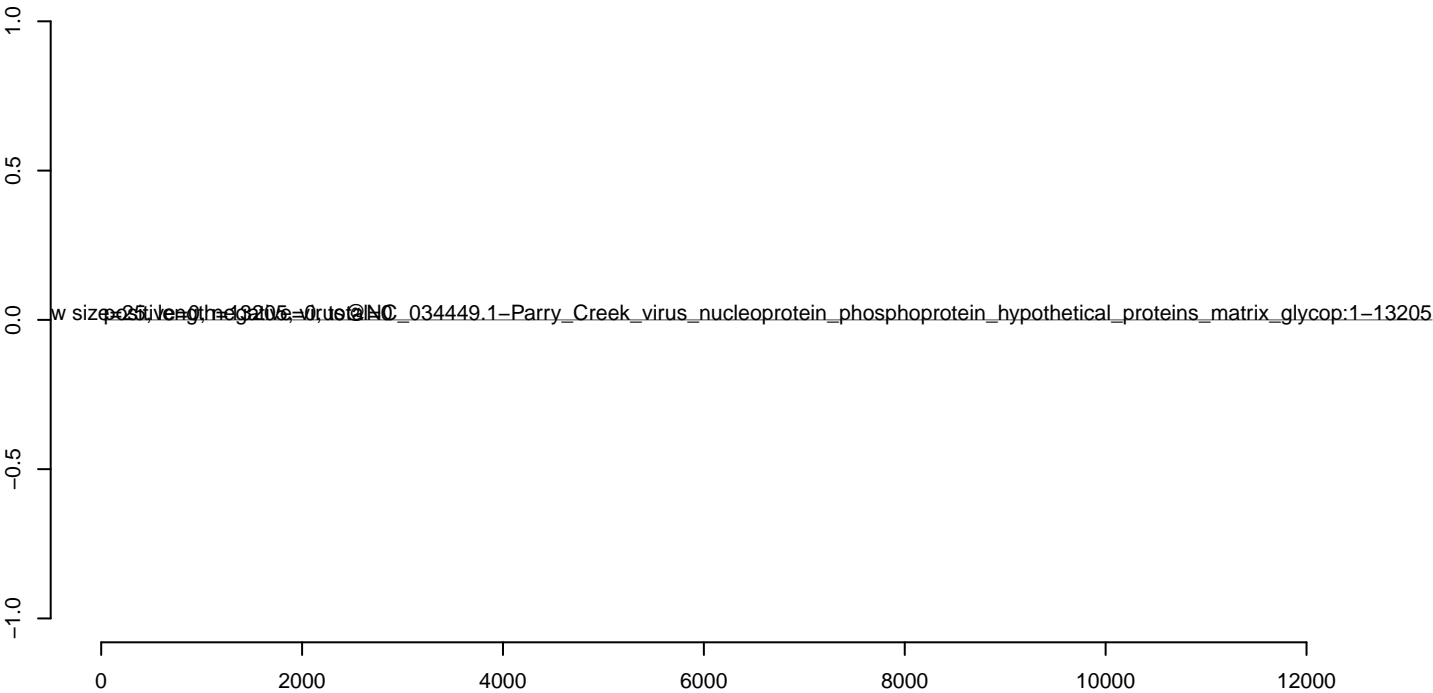
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



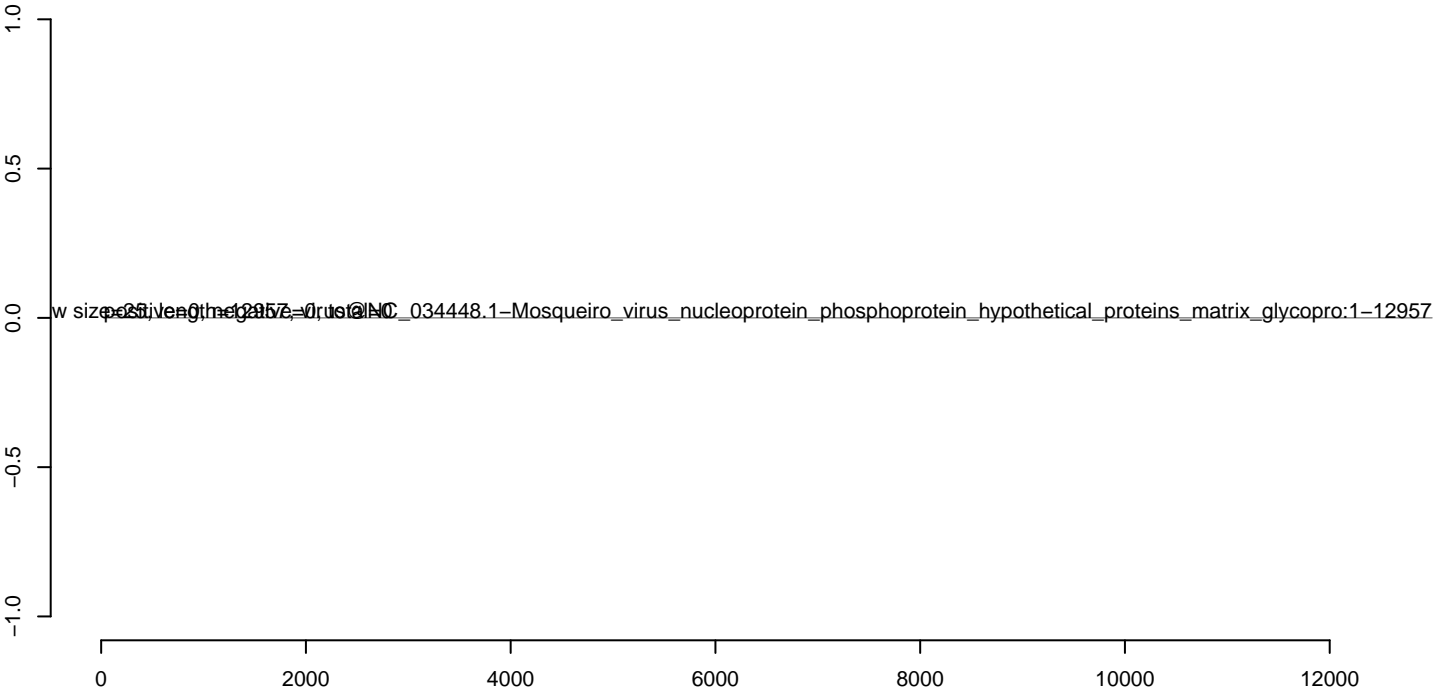
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



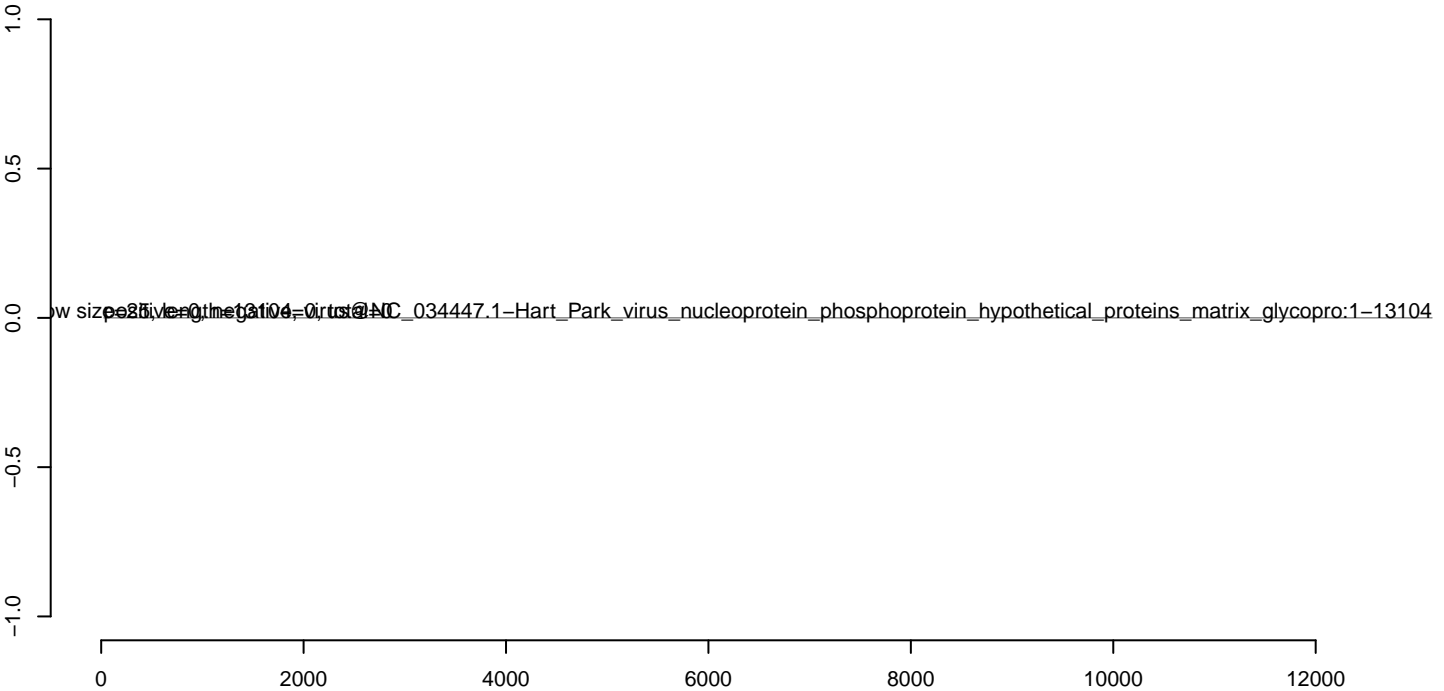
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



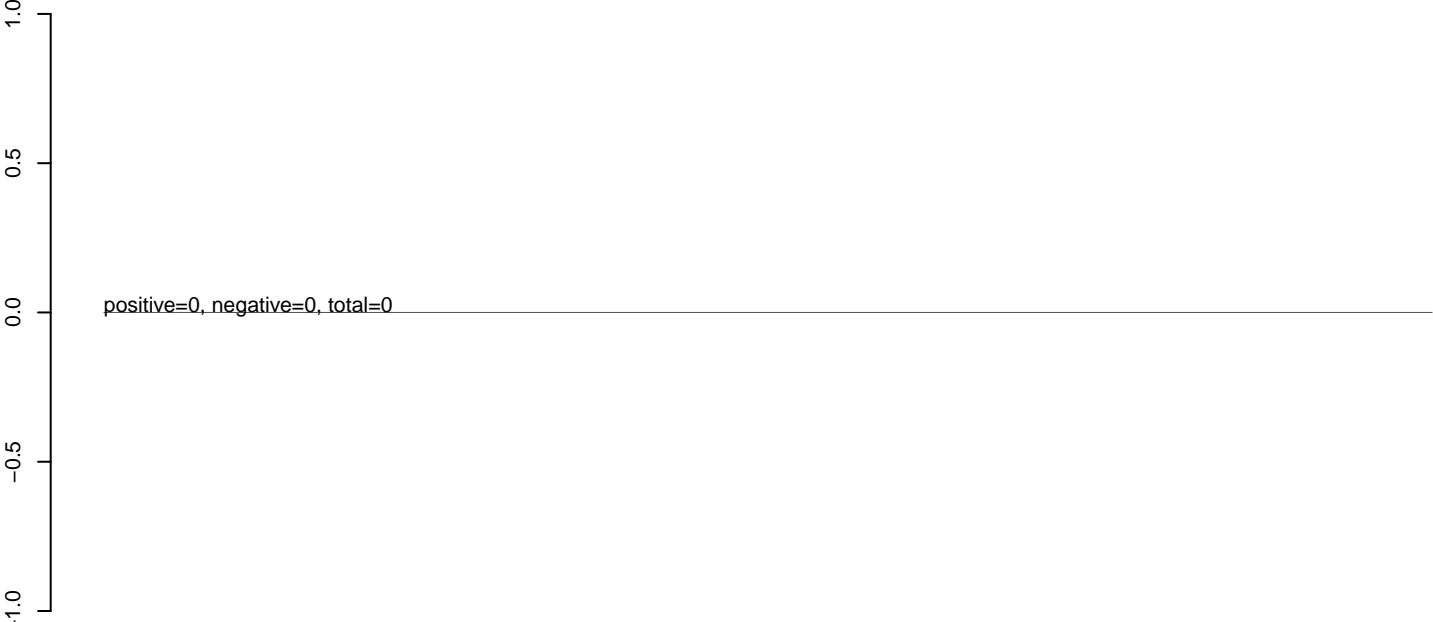
AeAlbo_C636_SINV_GFP_betaE.24_35.rep



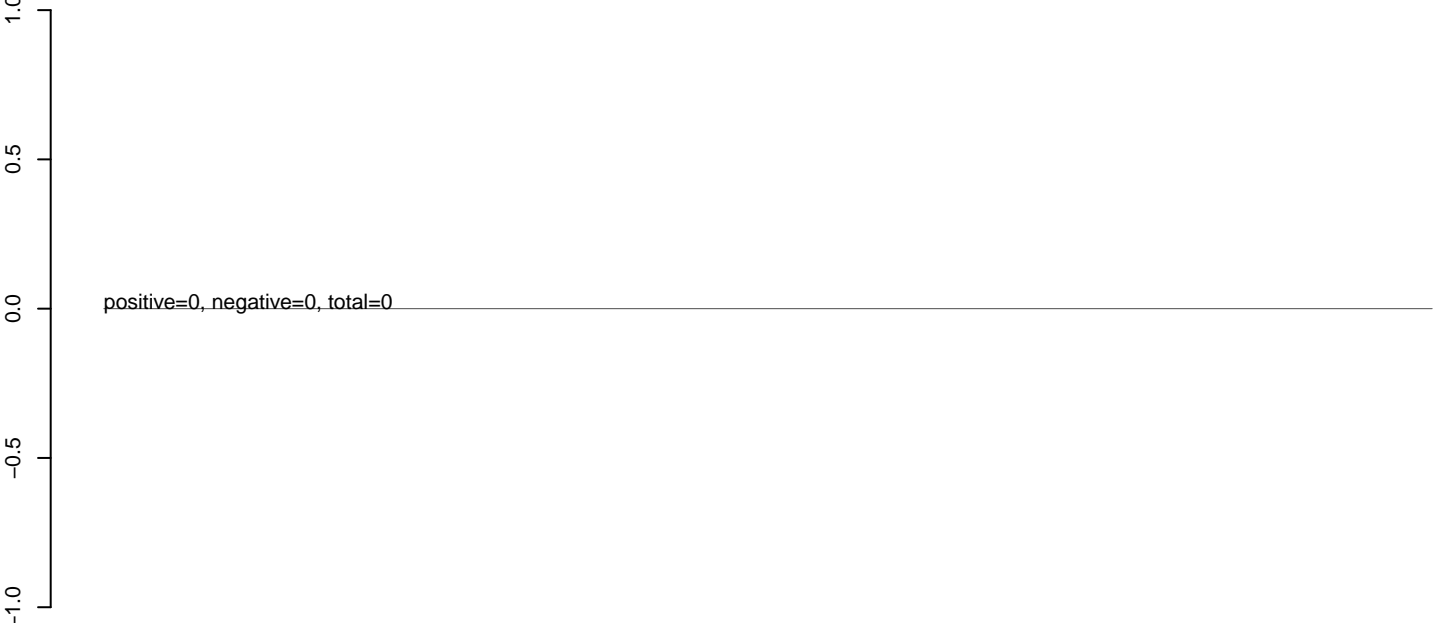
AeAlbo_C636_SINV_GFP_betaE.rep



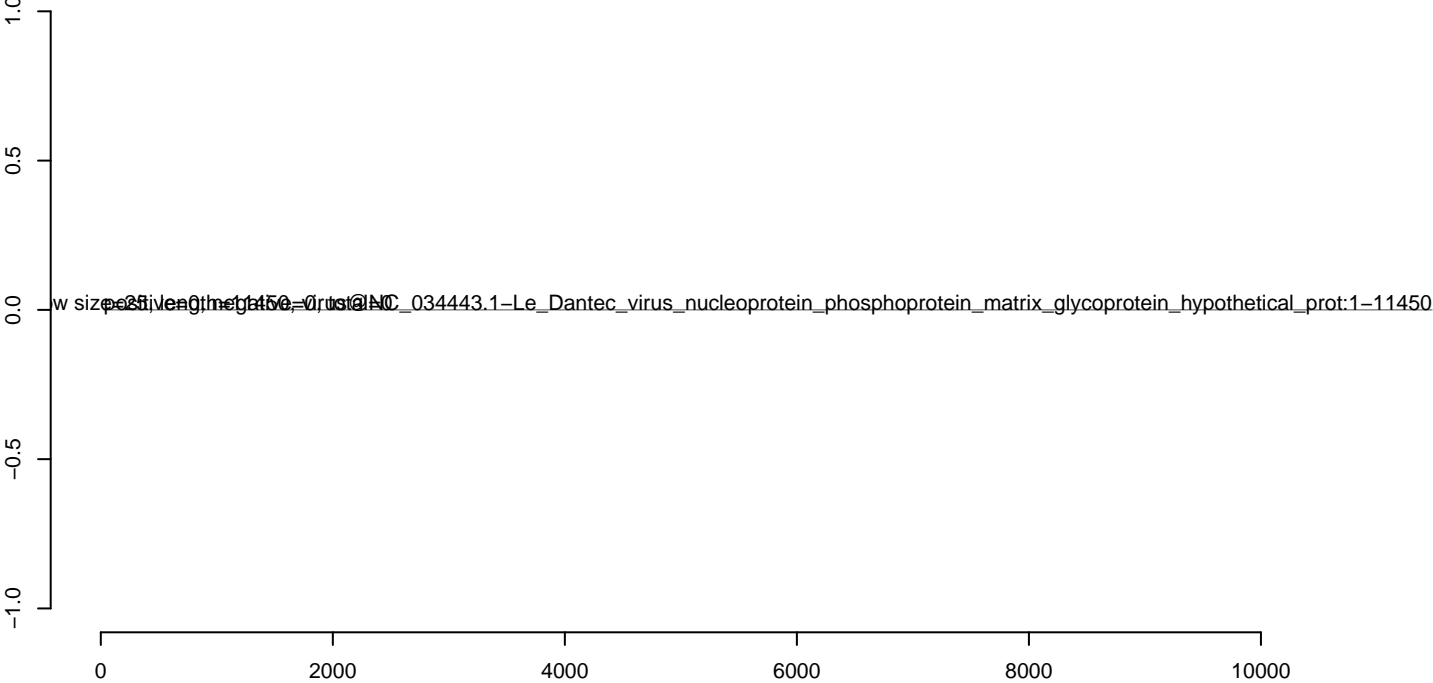
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



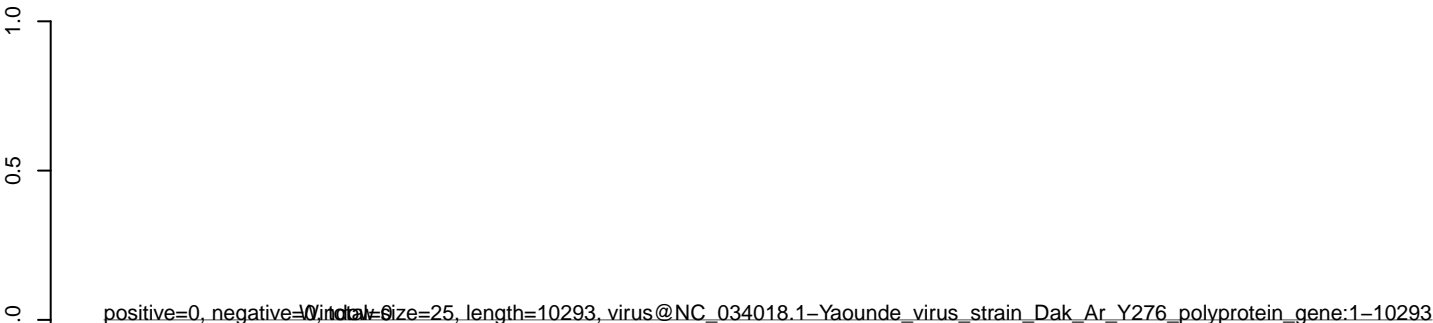
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



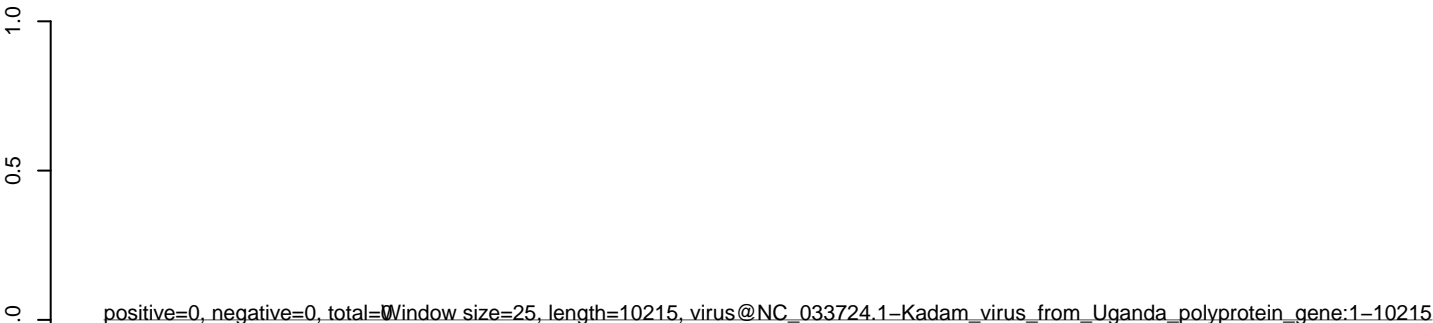
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



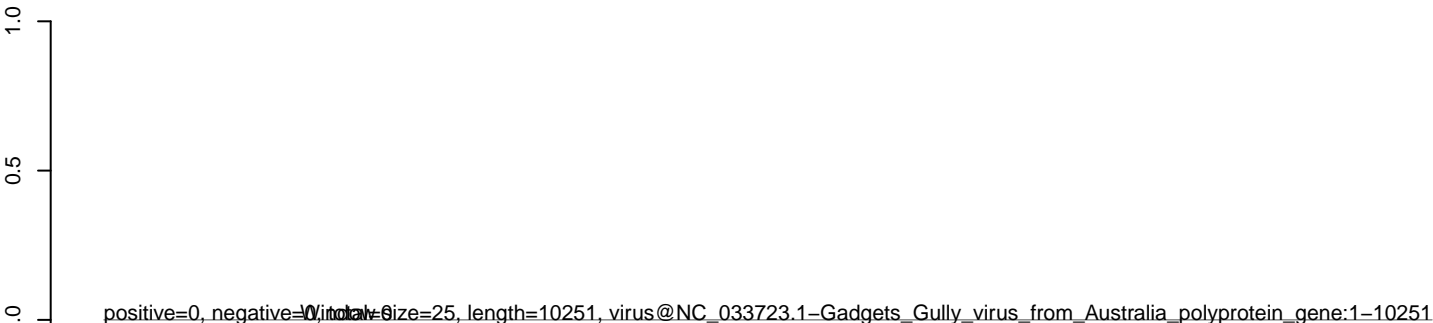
AeAlbo_C636_SINV_GFP_betaE.18_23.rep



AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



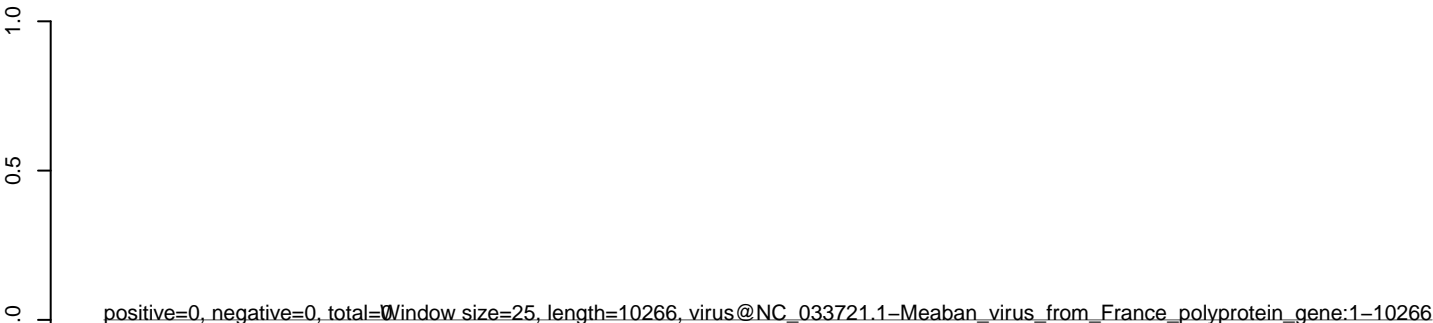
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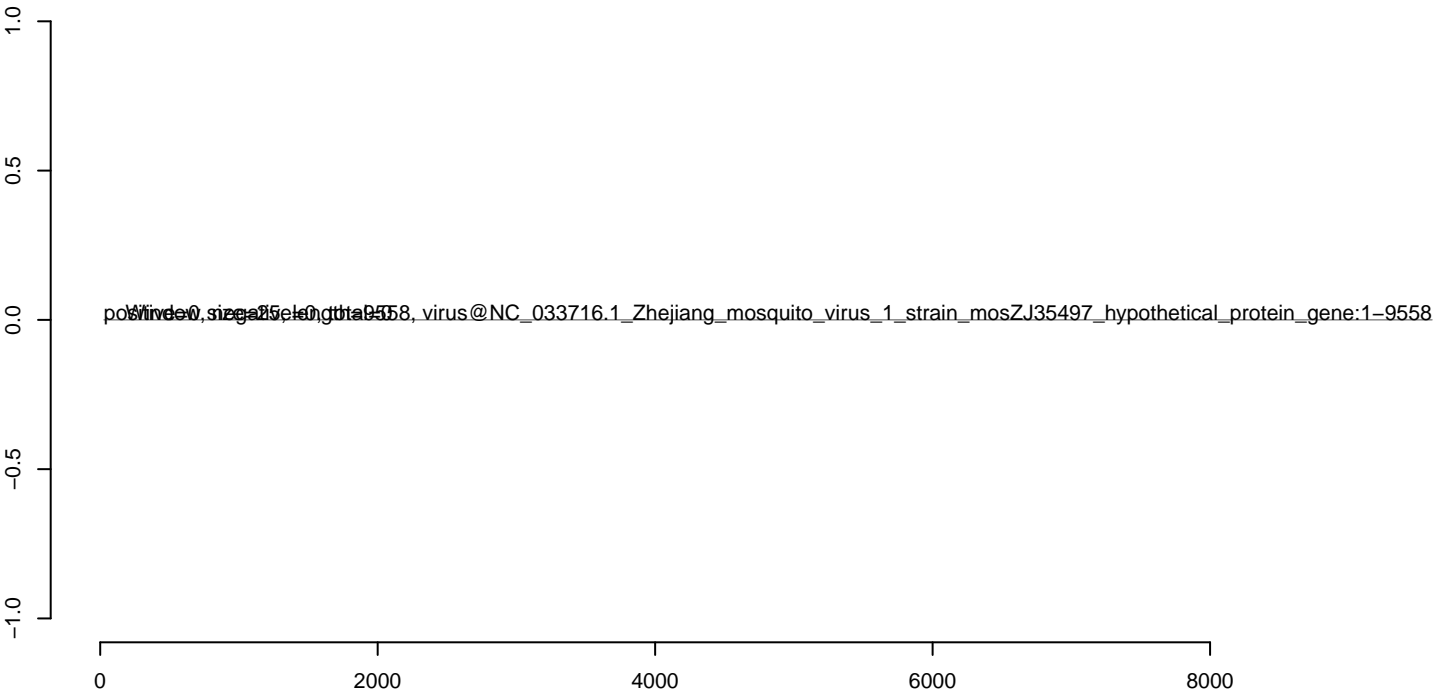
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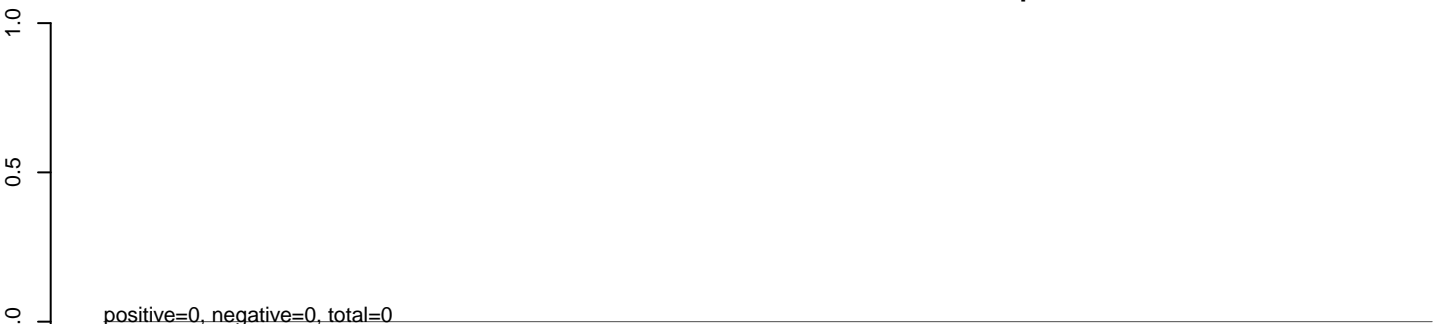
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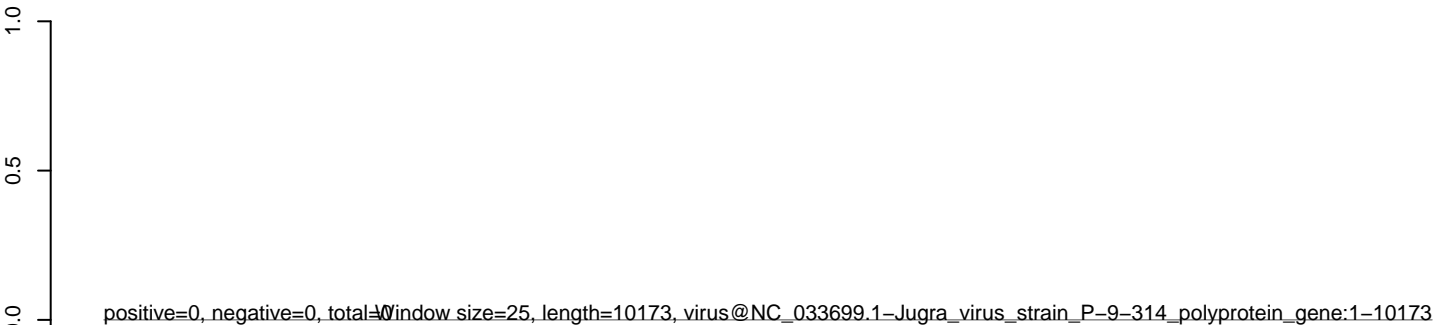
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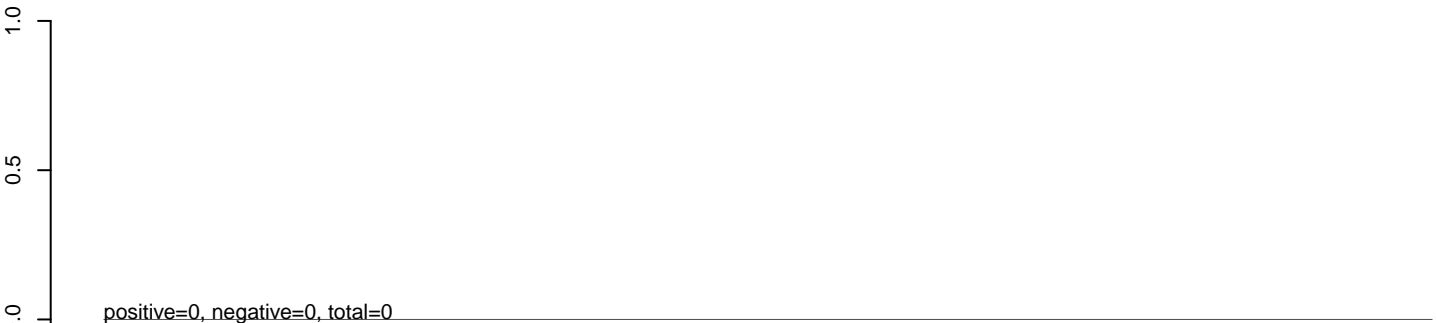
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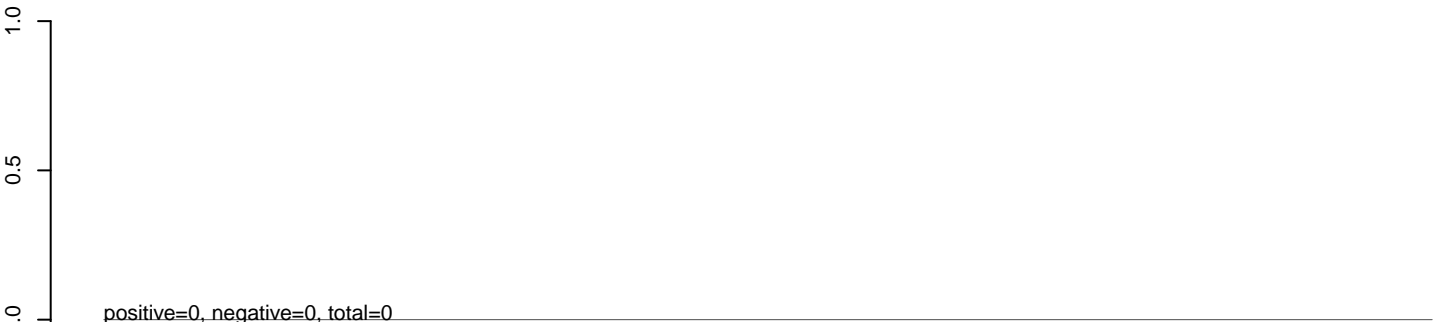
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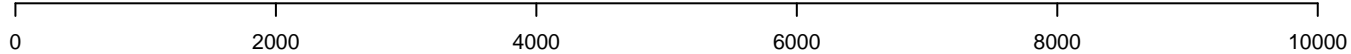
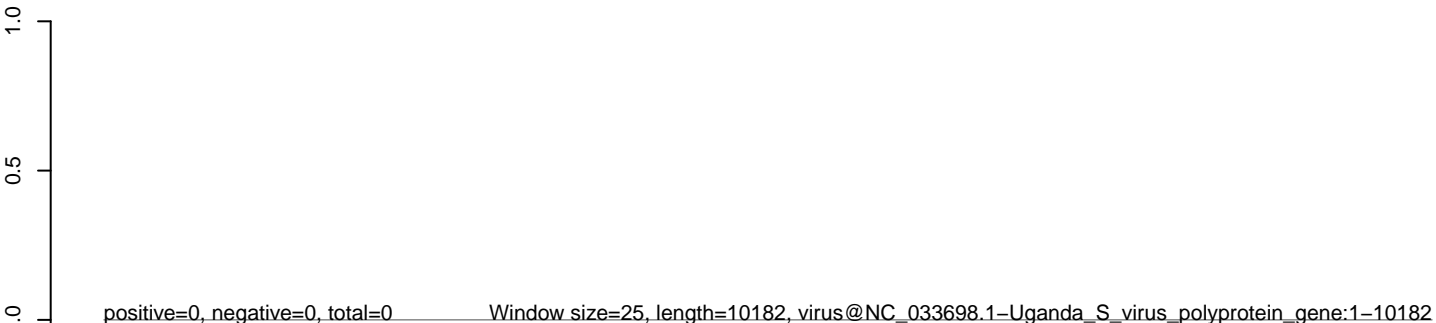
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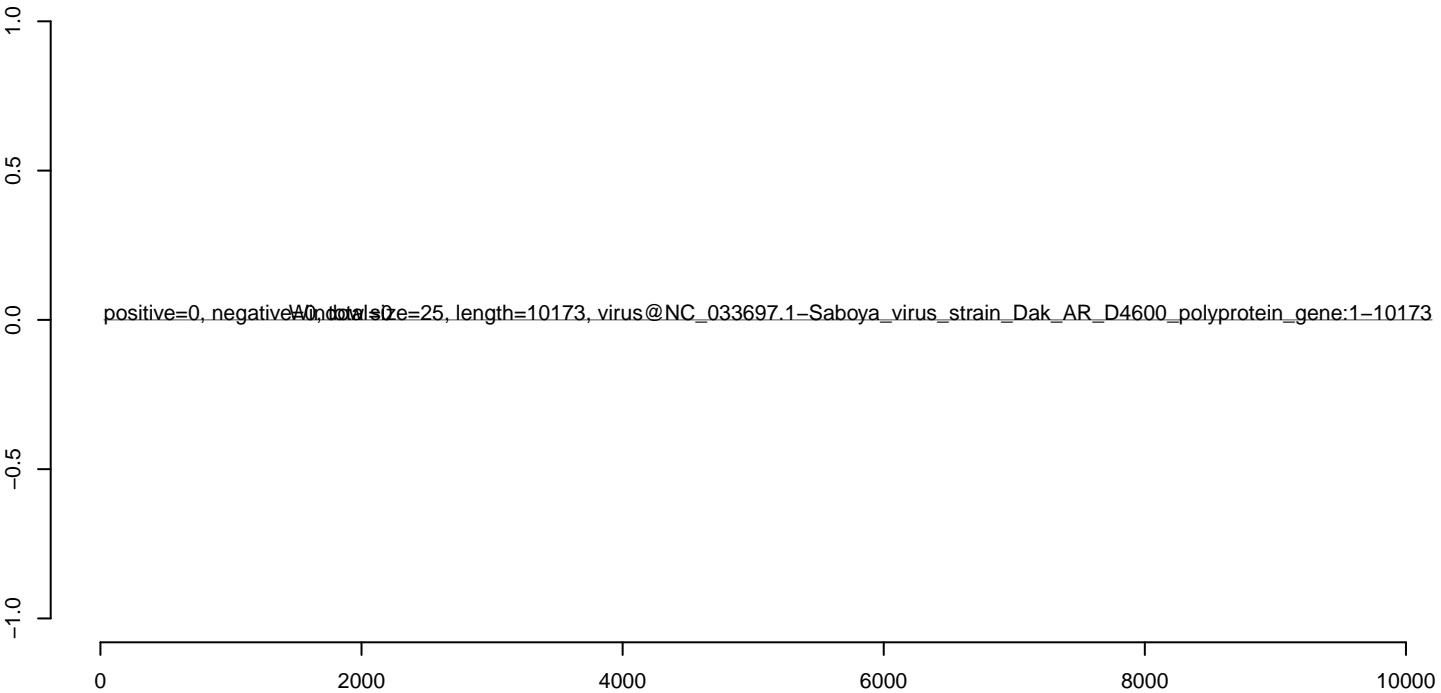
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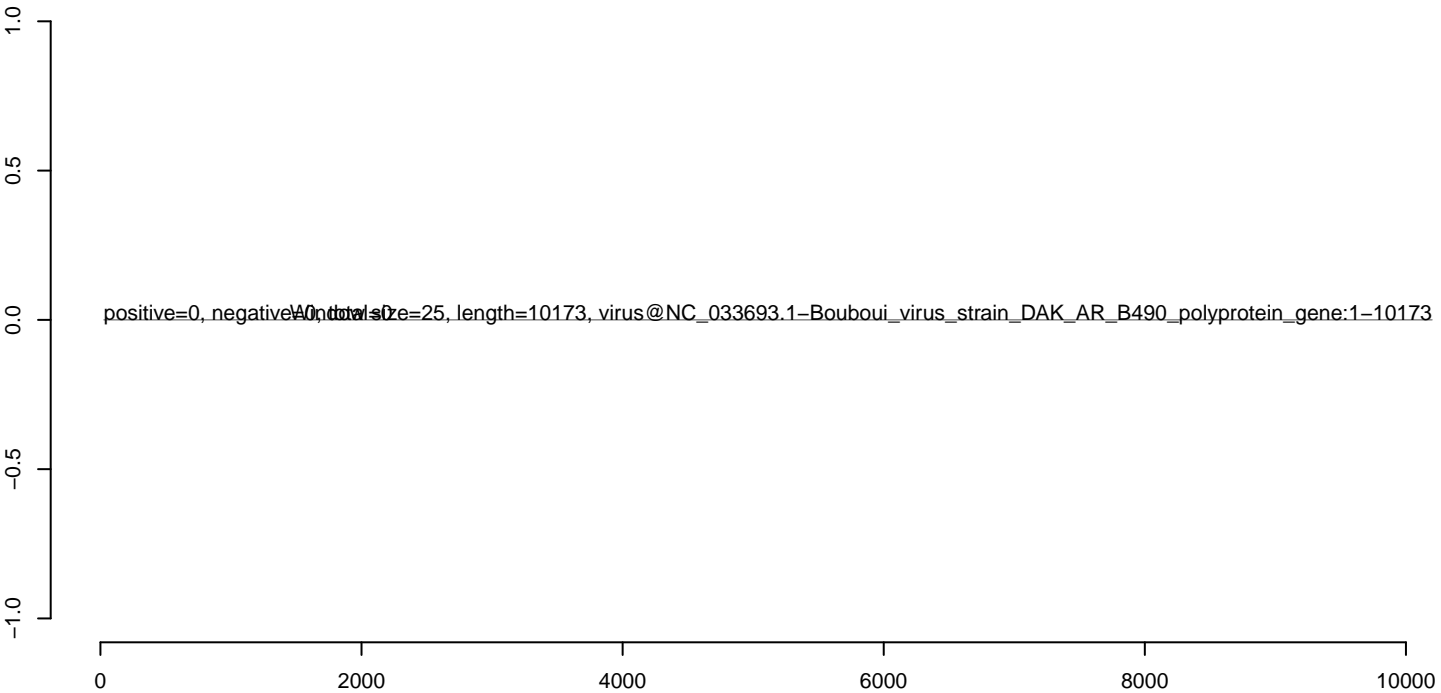
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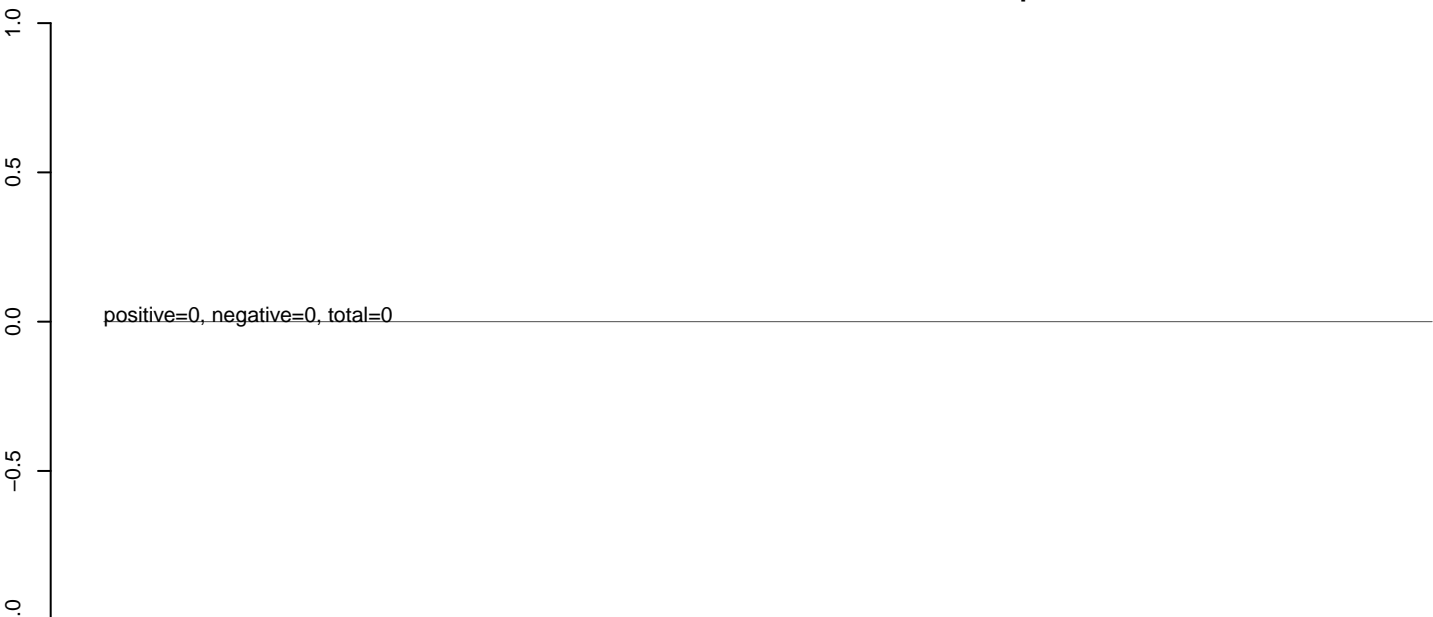
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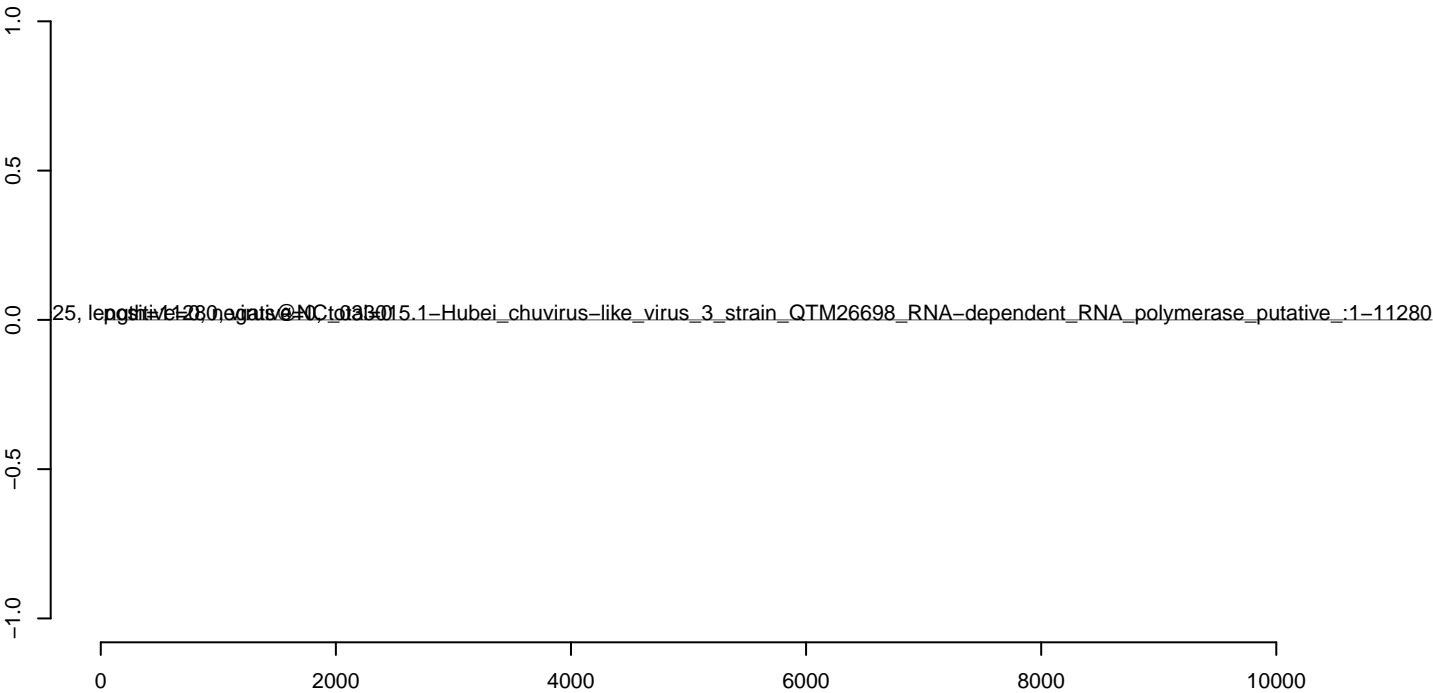
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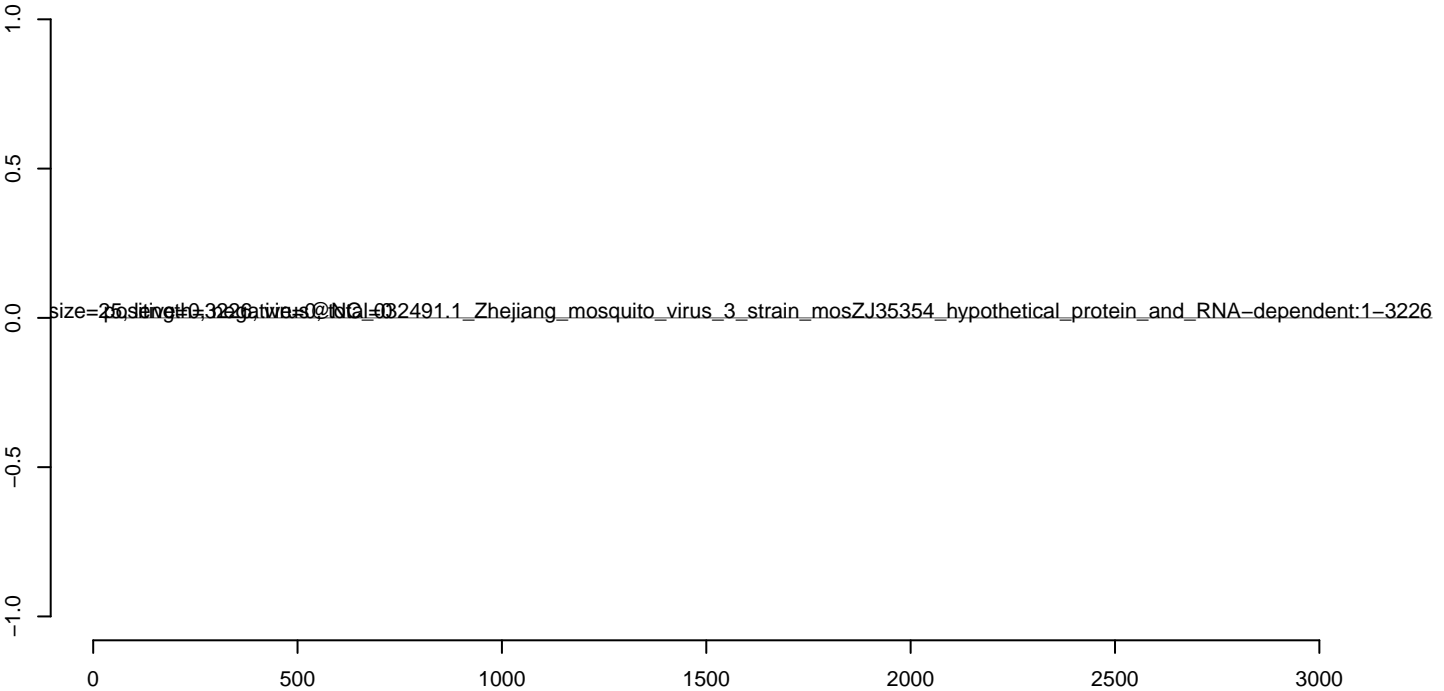
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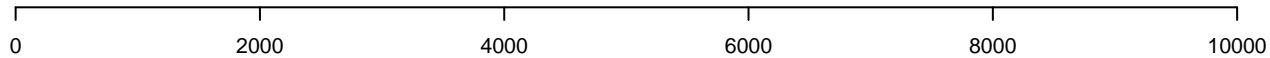
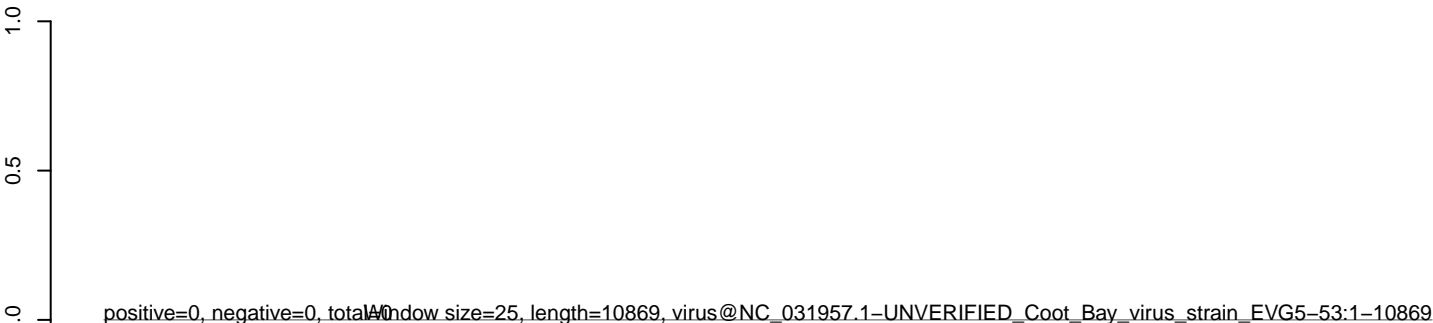
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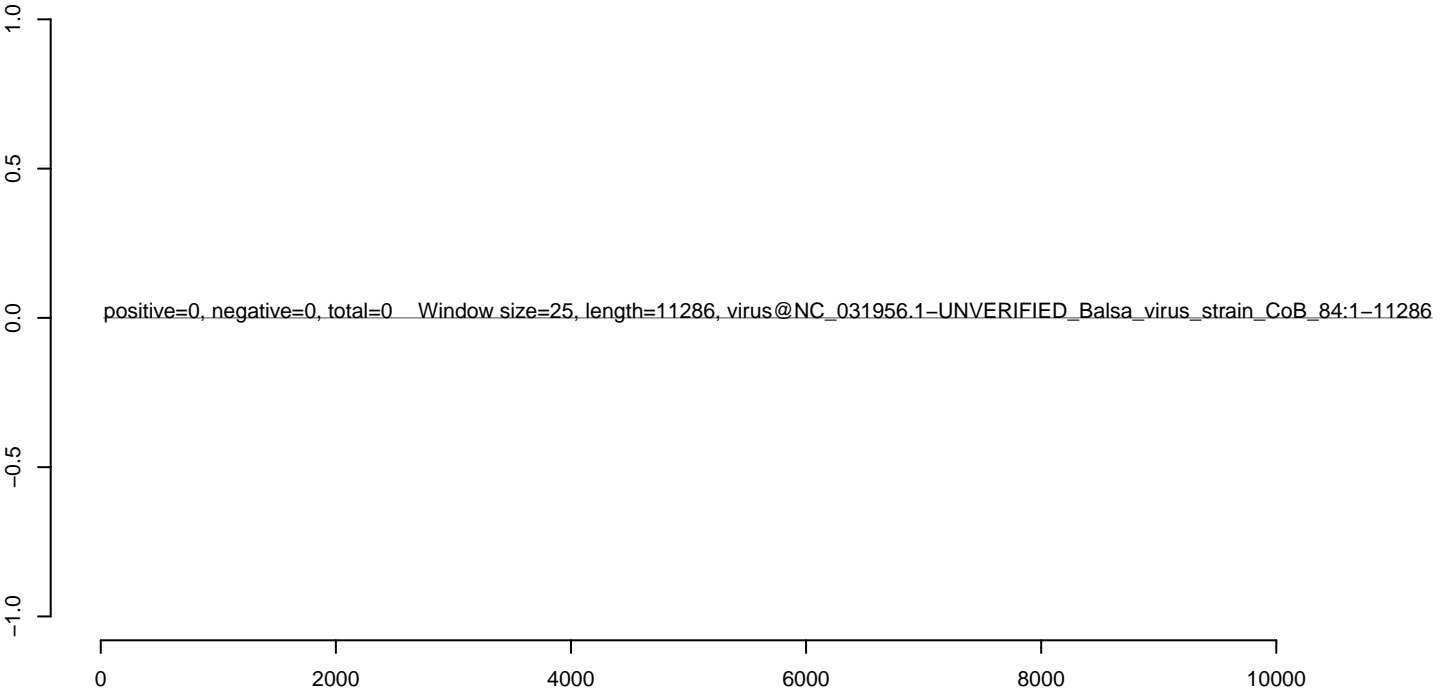
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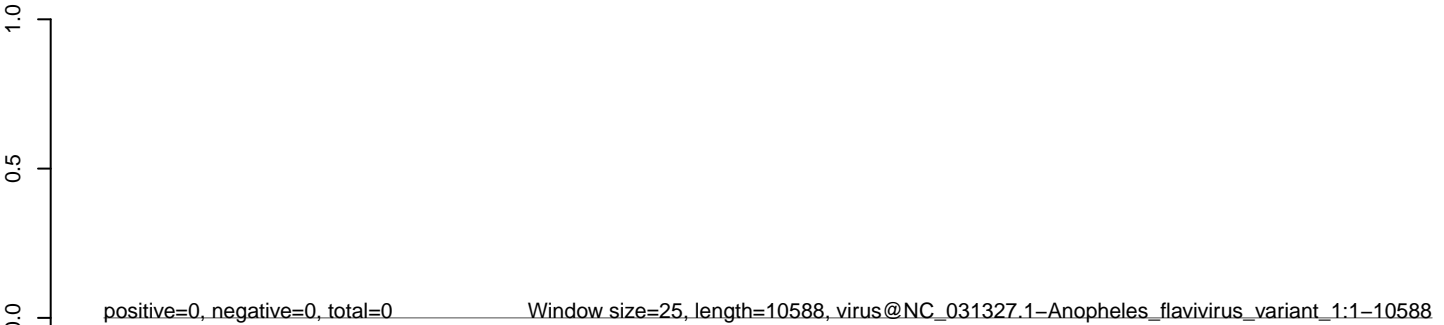
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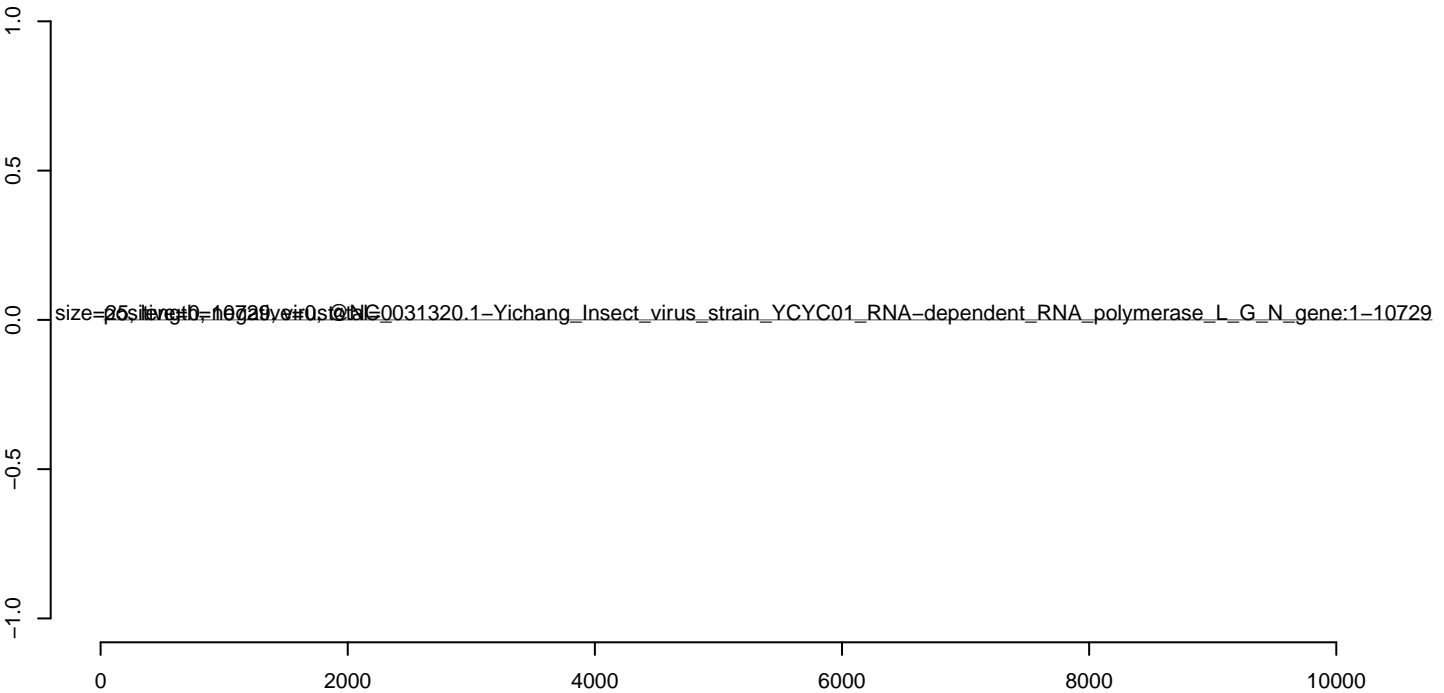
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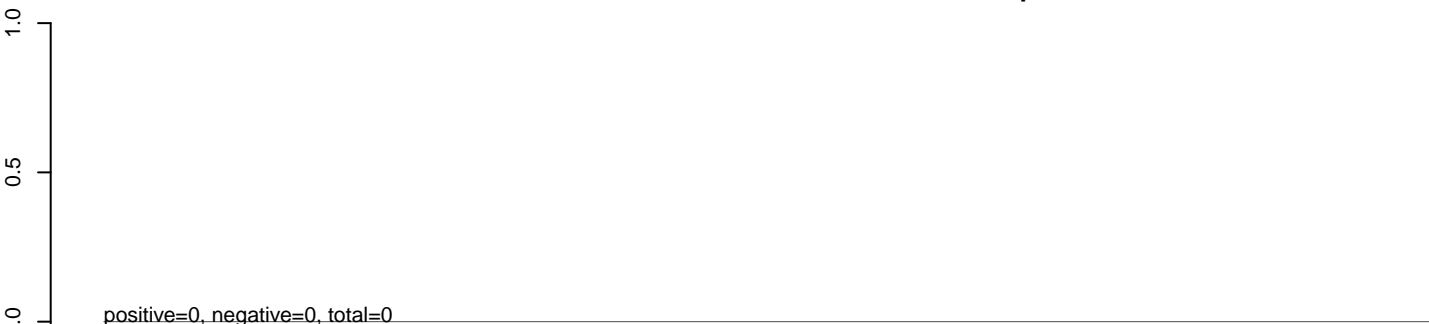
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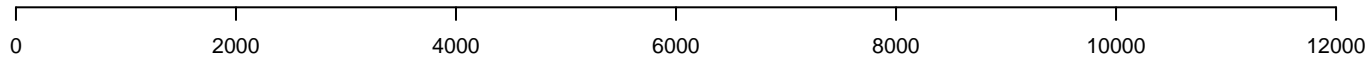
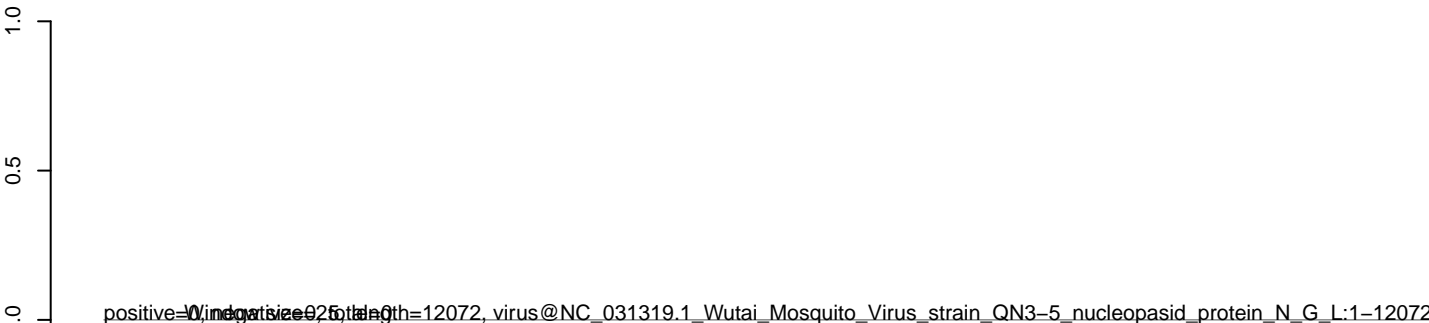
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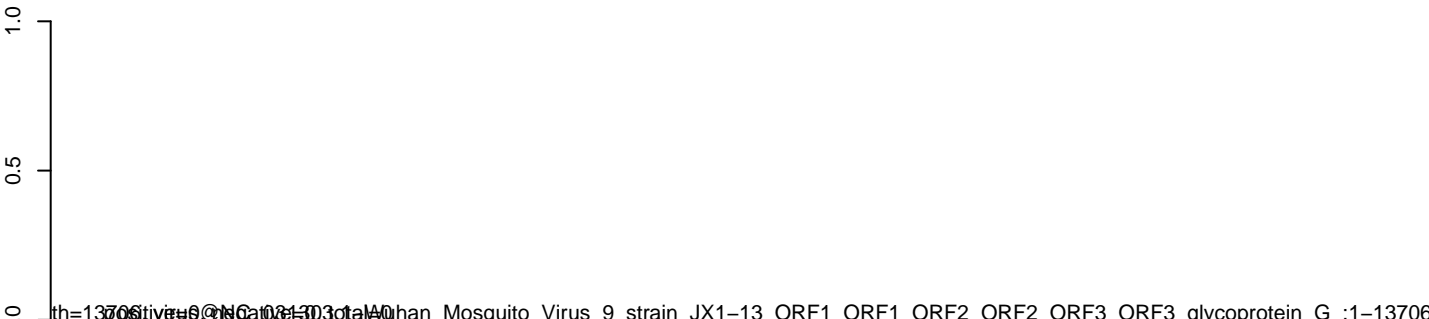
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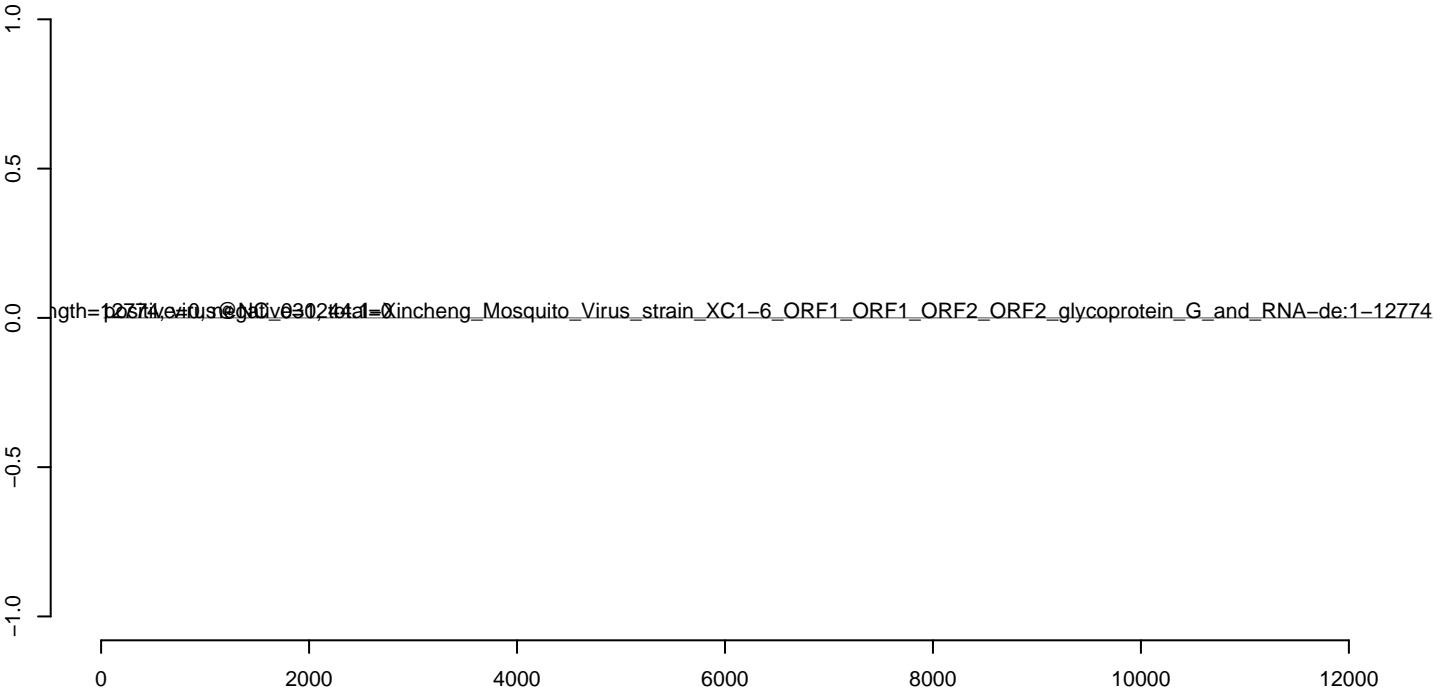
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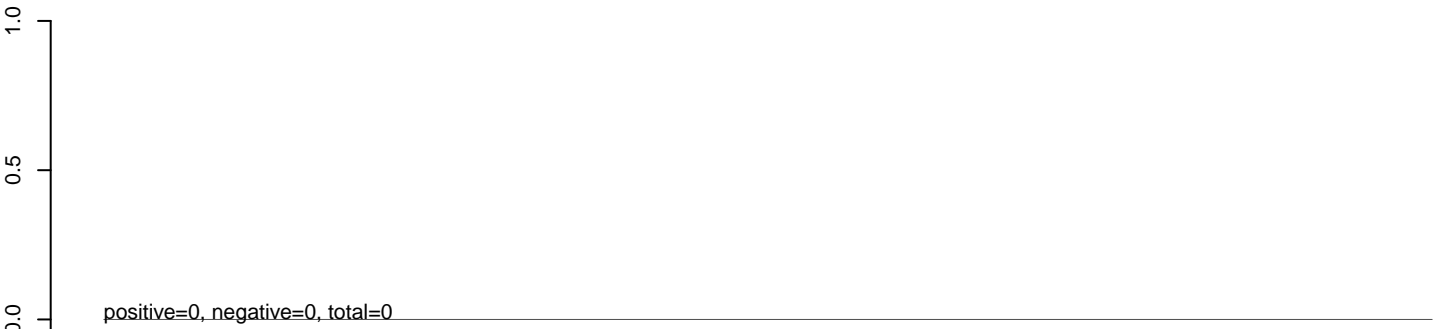
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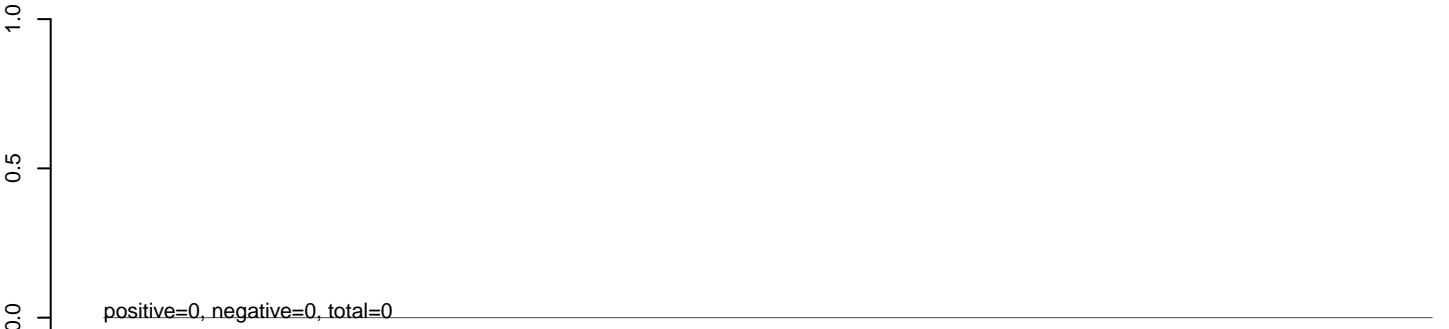
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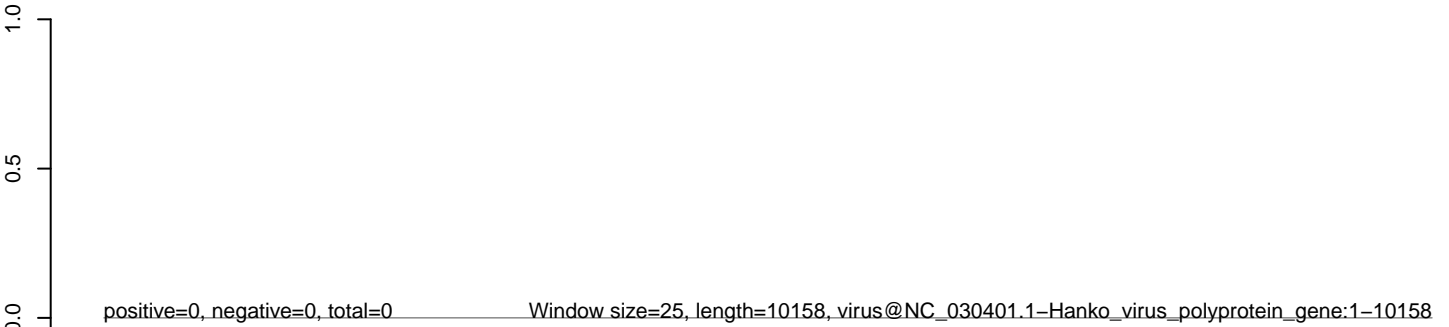
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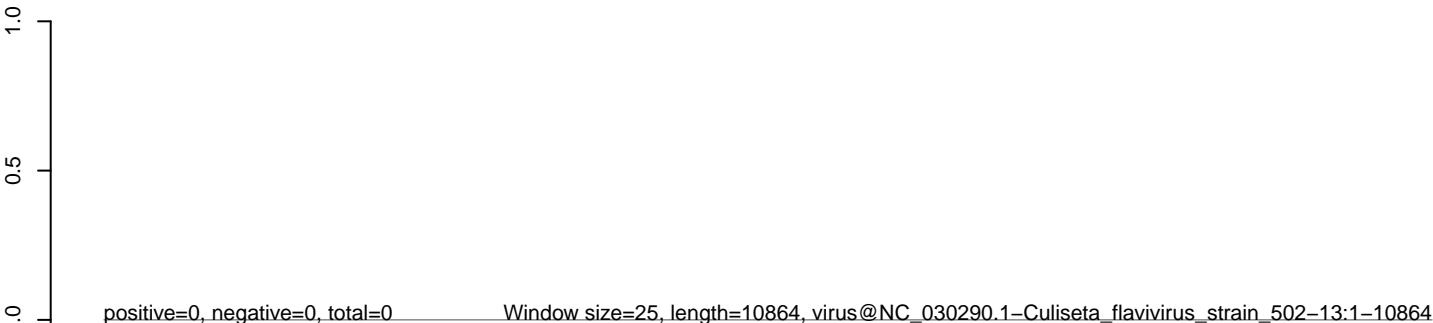
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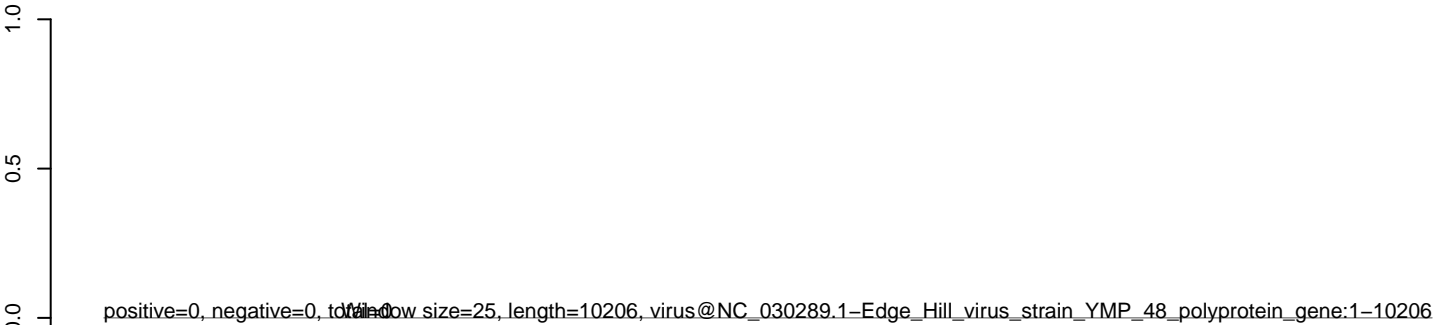
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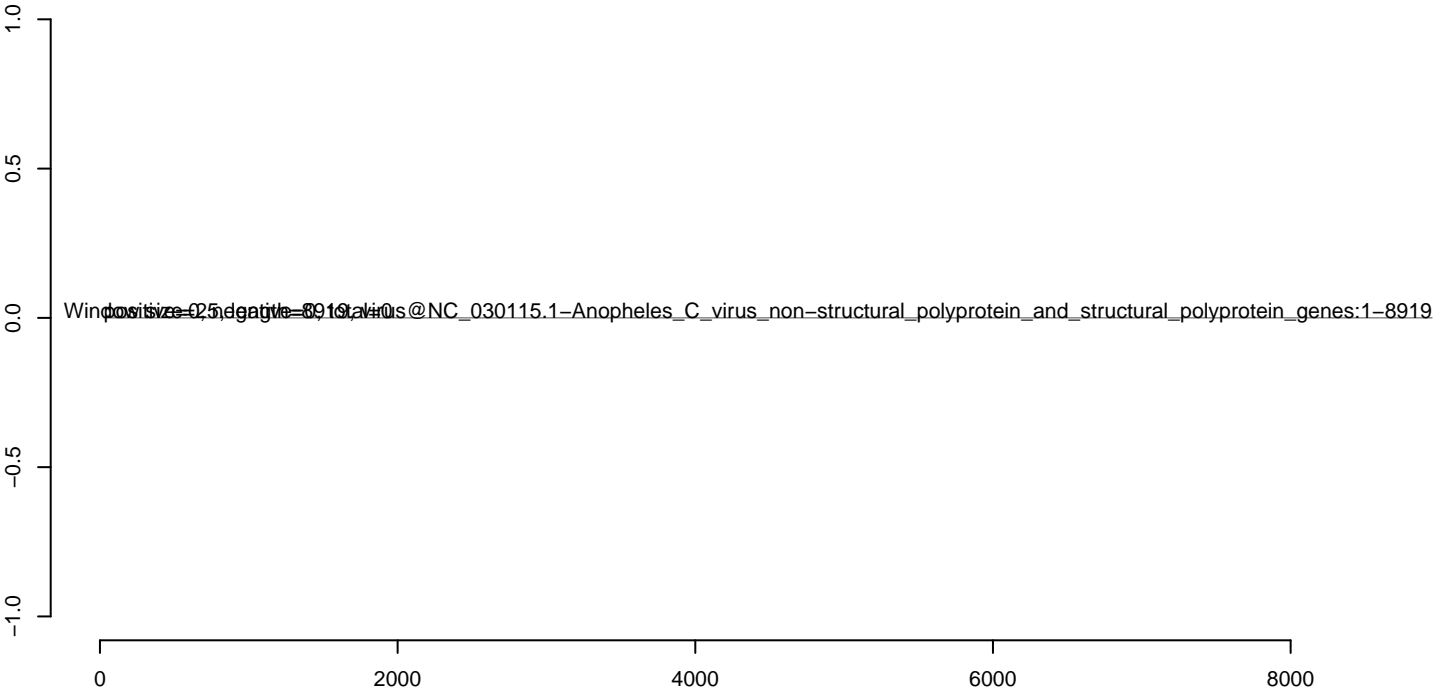
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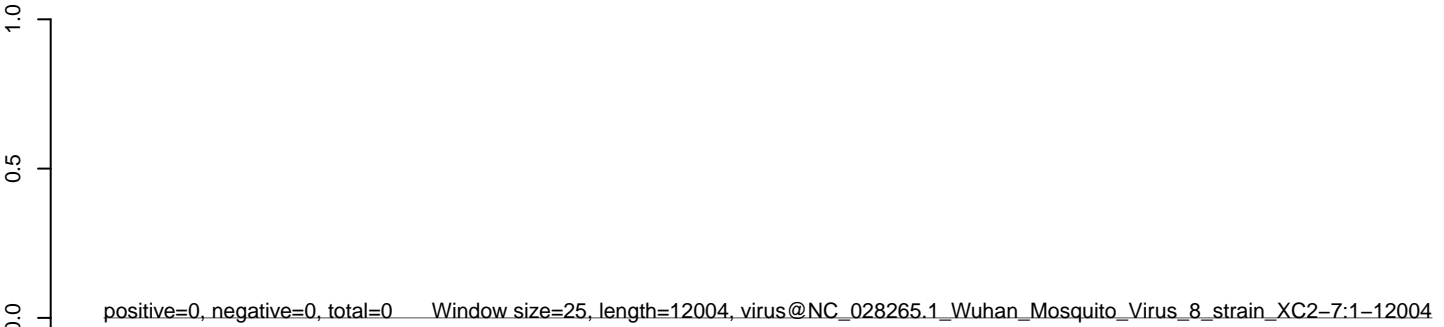
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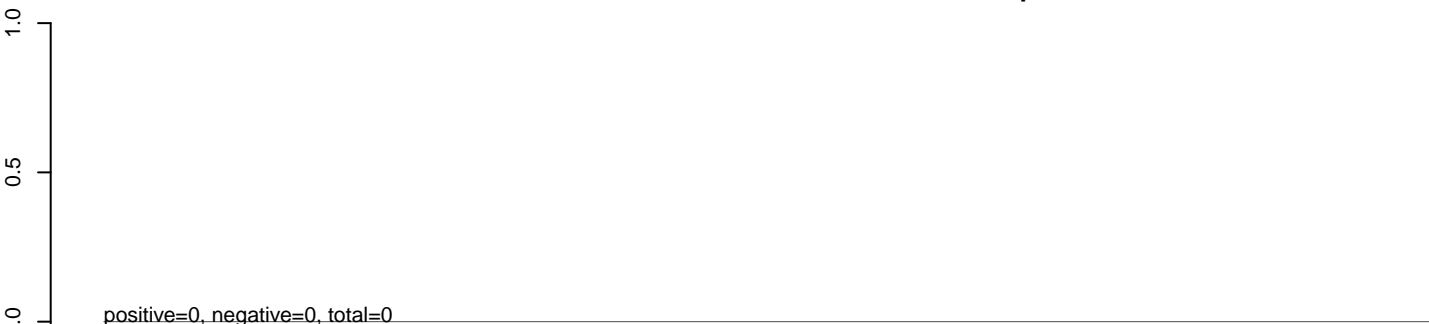
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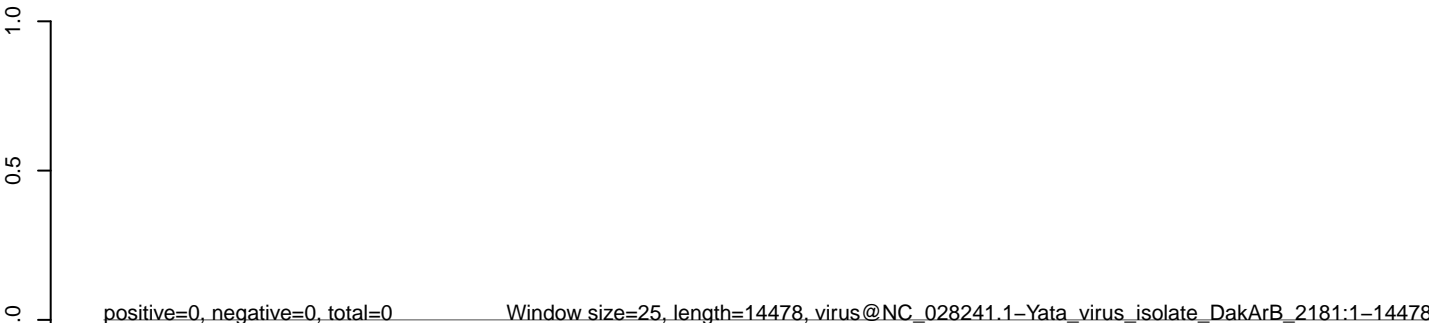
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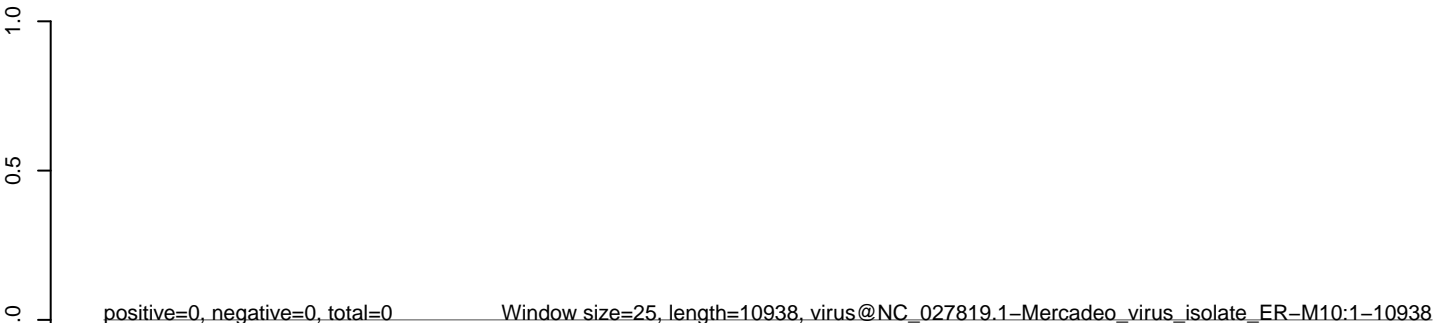
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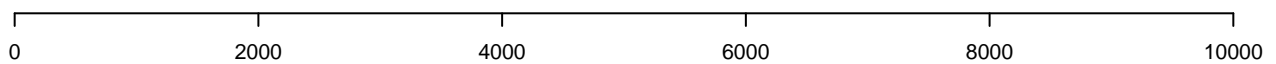
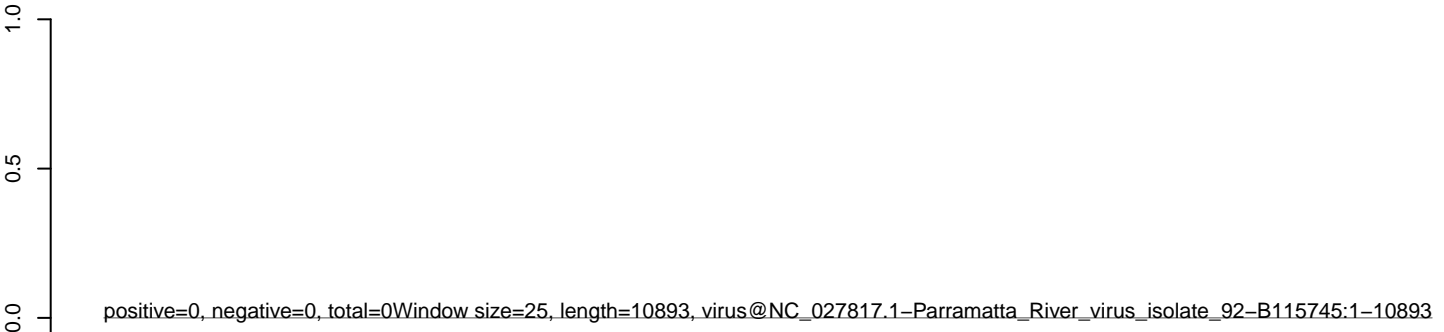
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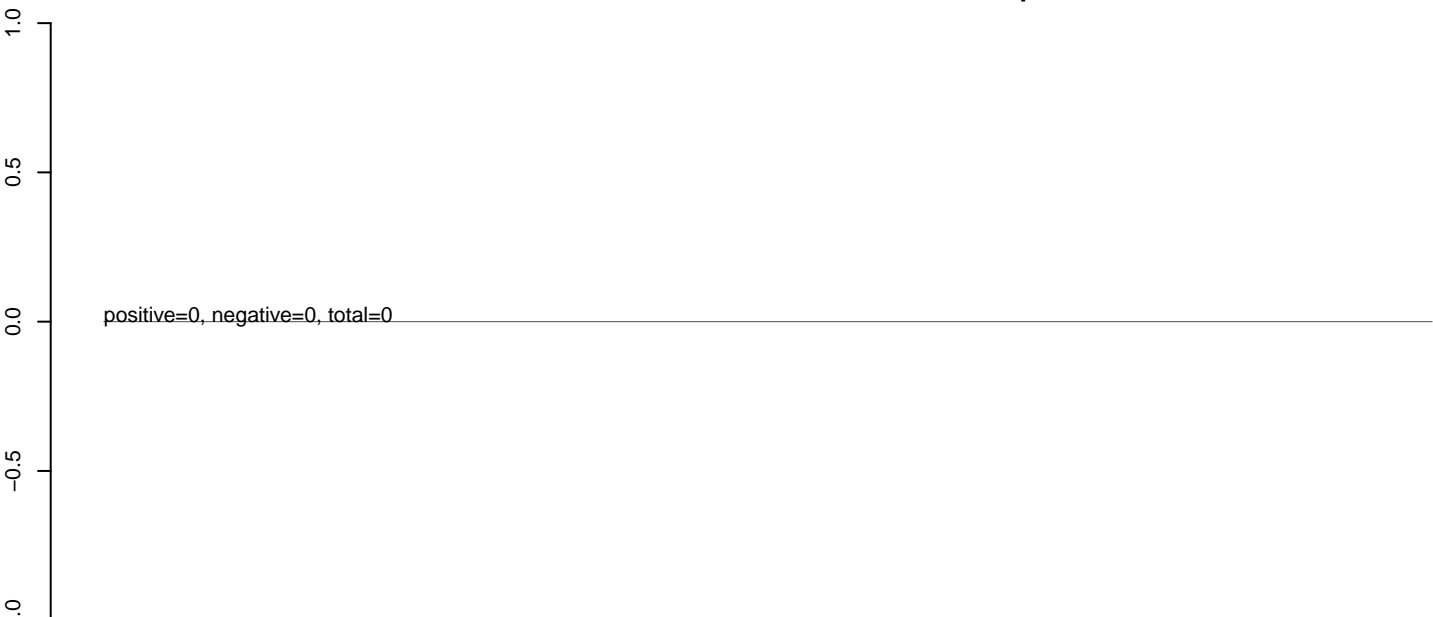
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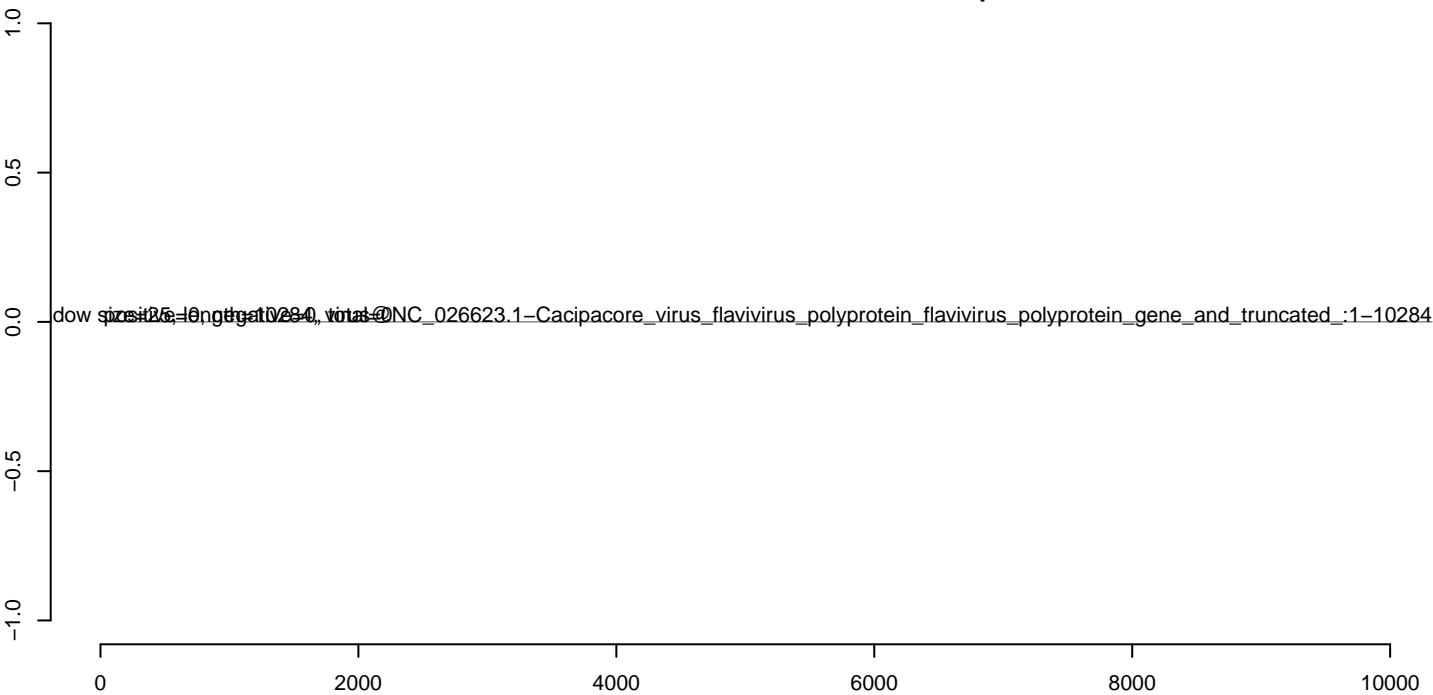
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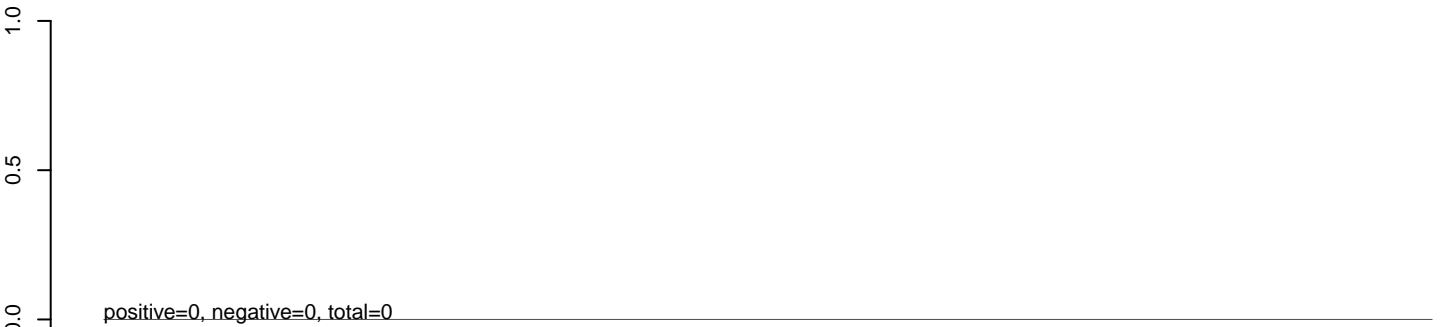
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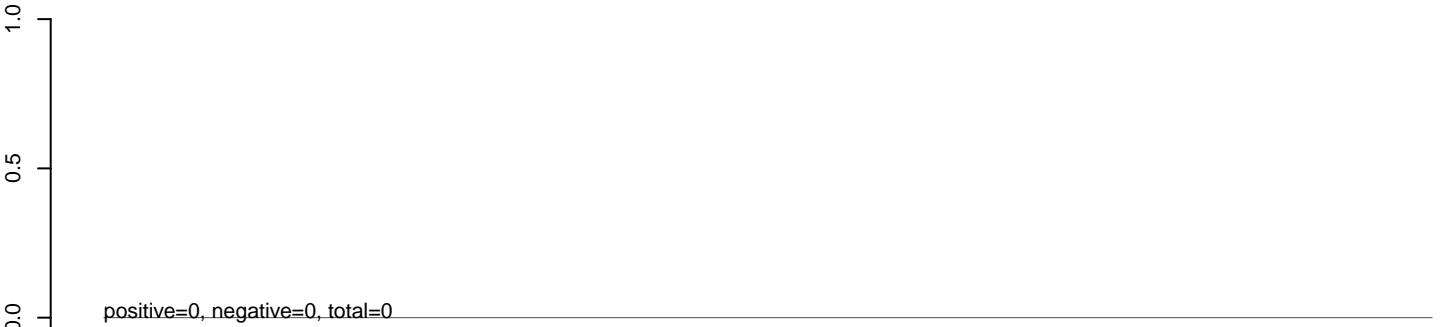
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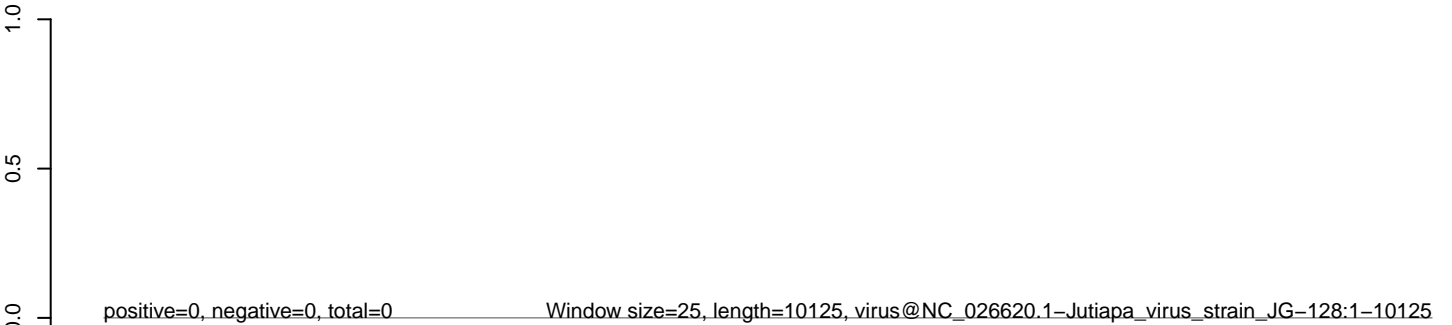
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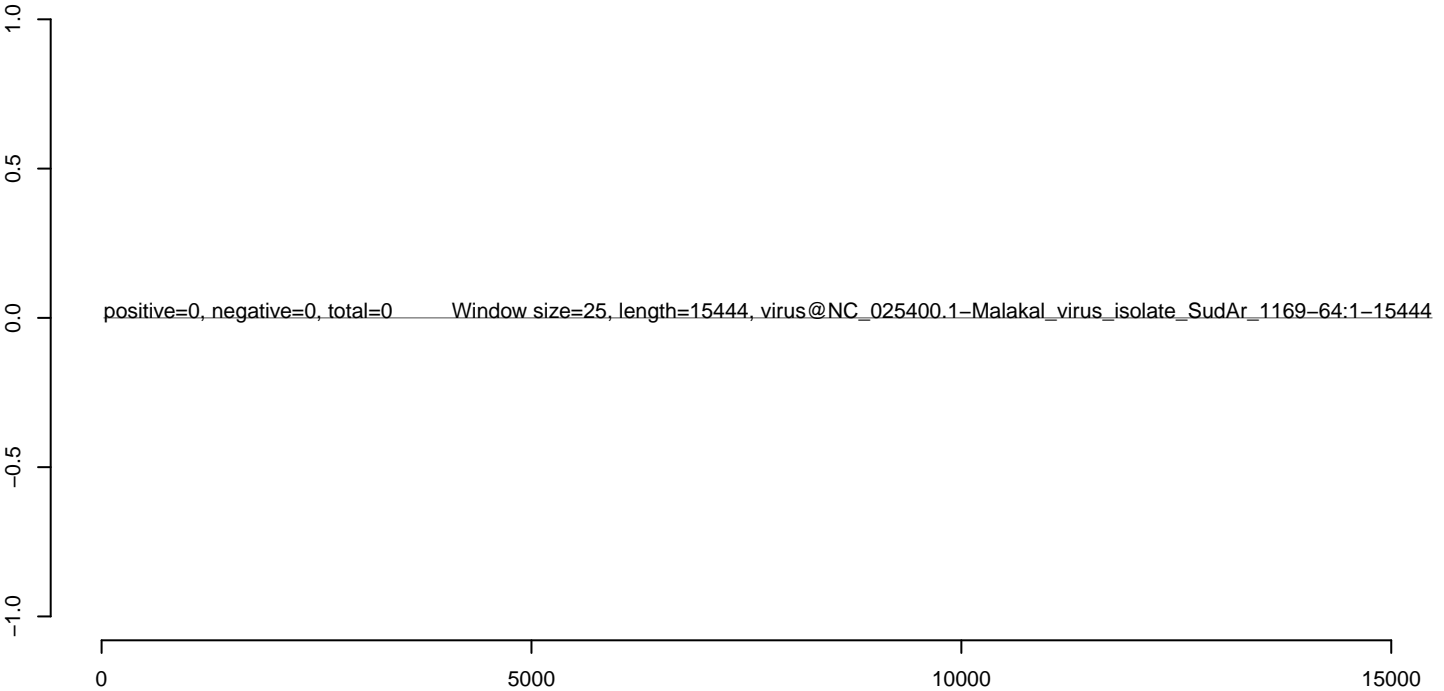
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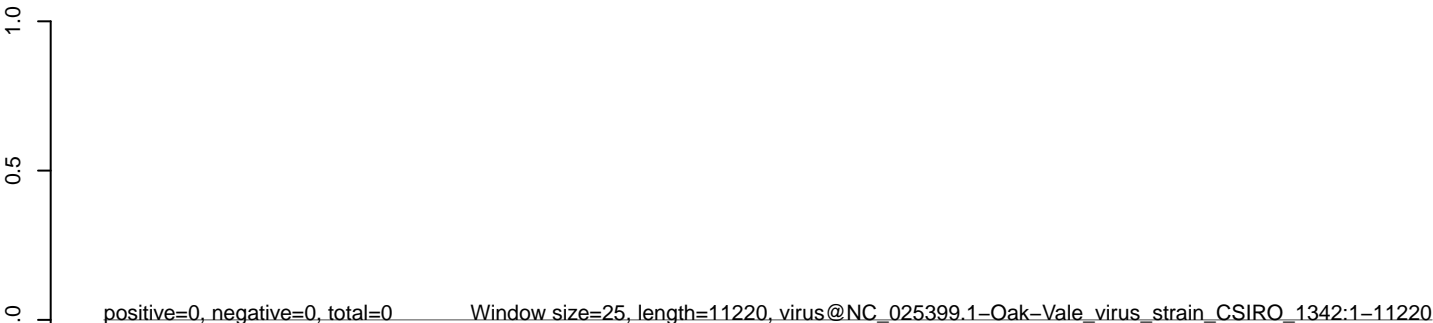
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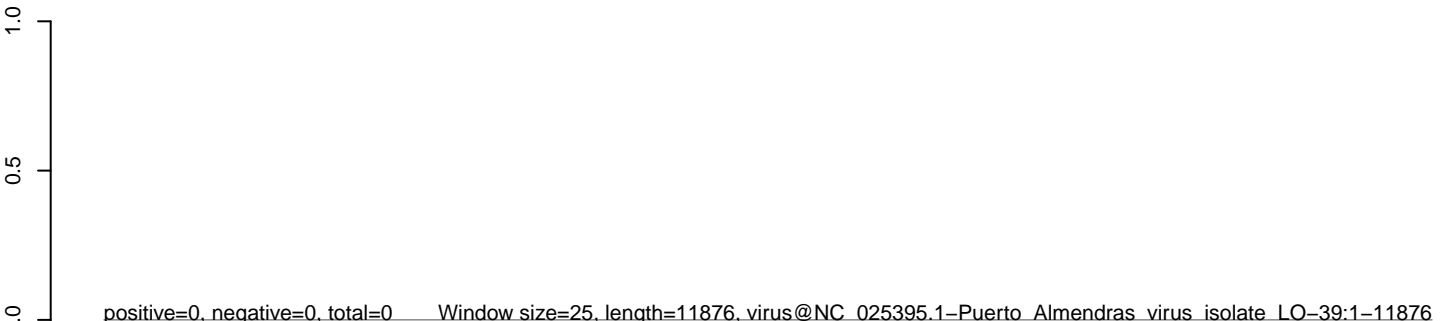
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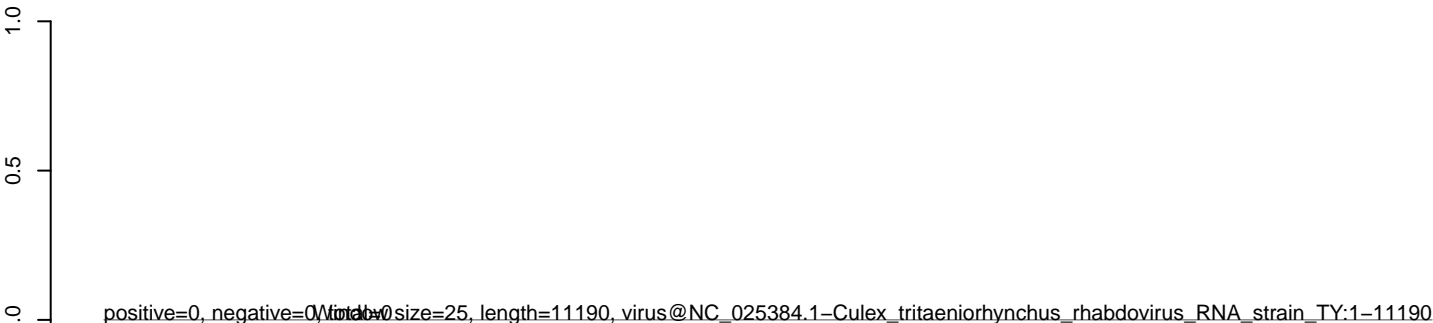
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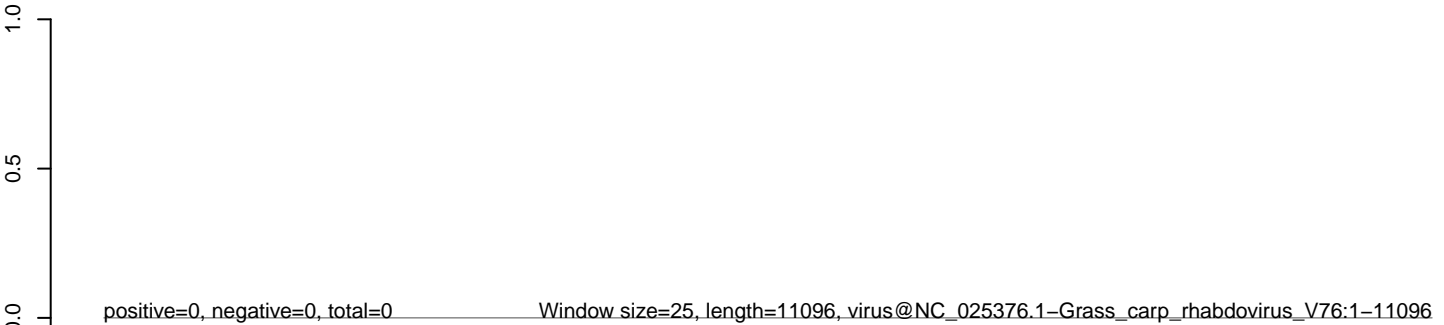
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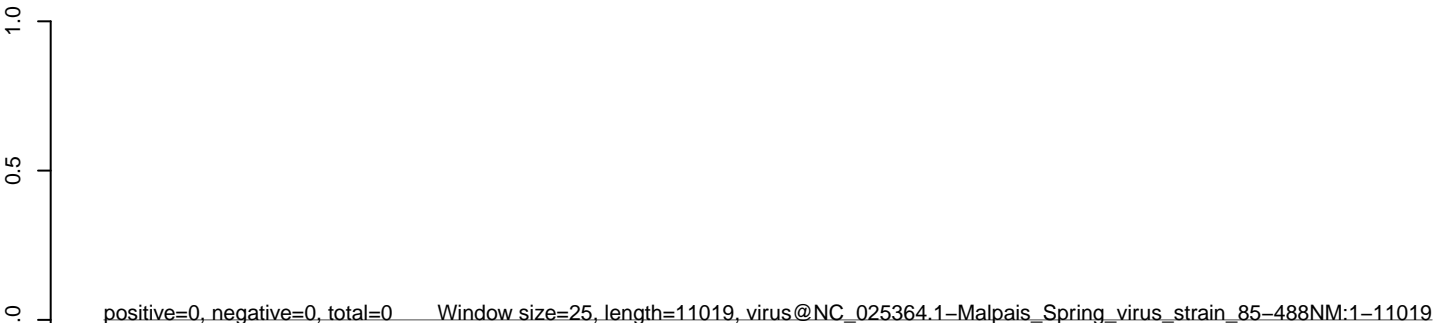
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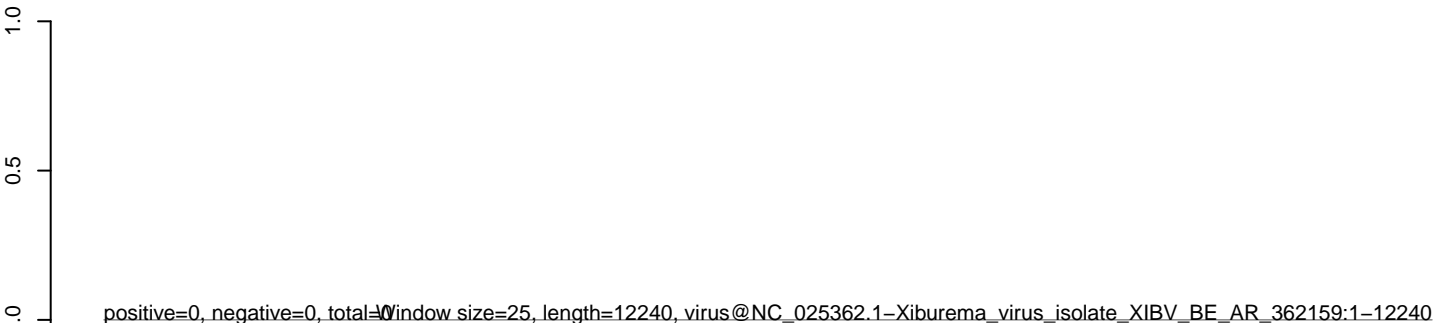
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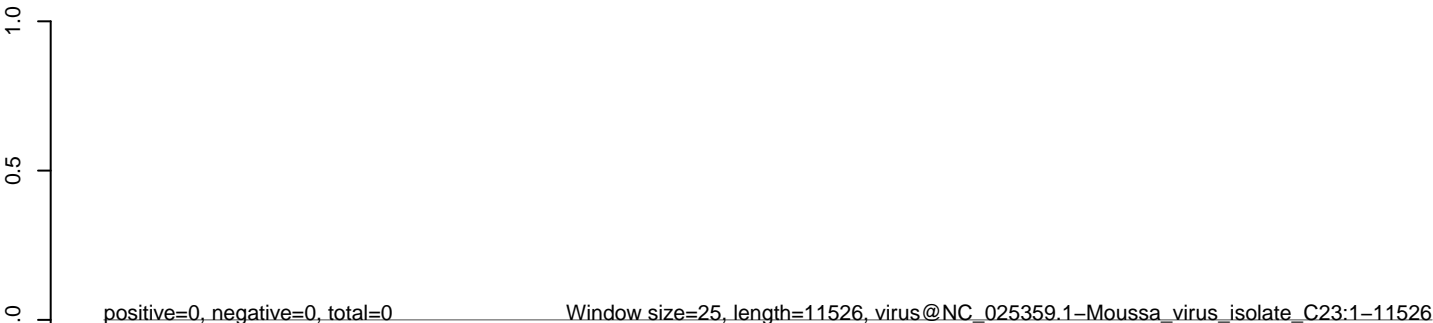
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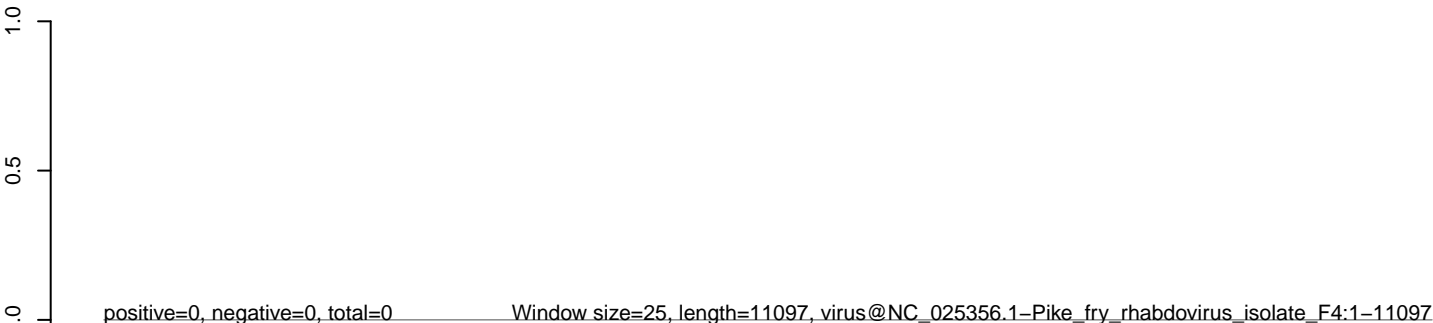
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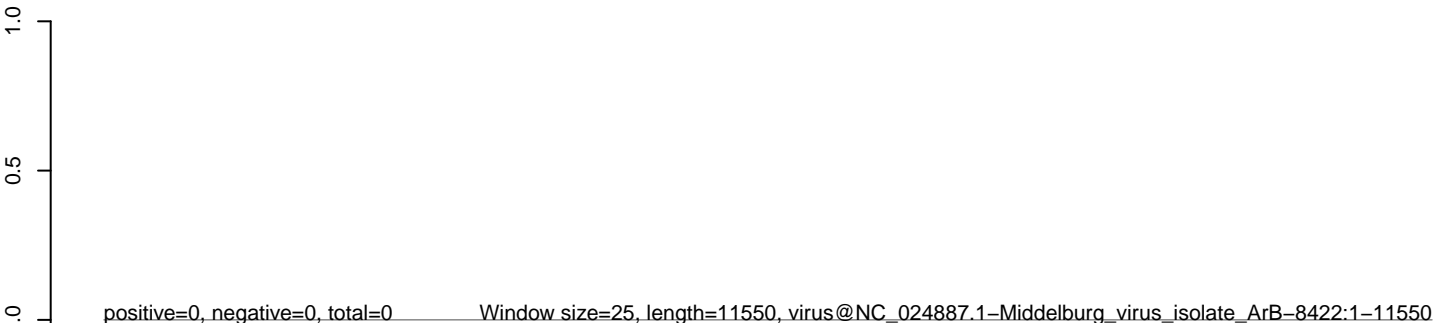
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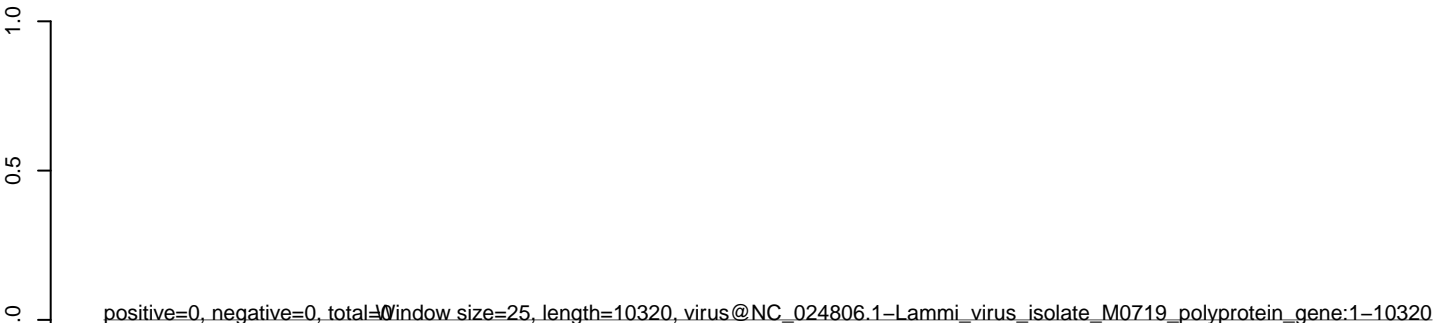
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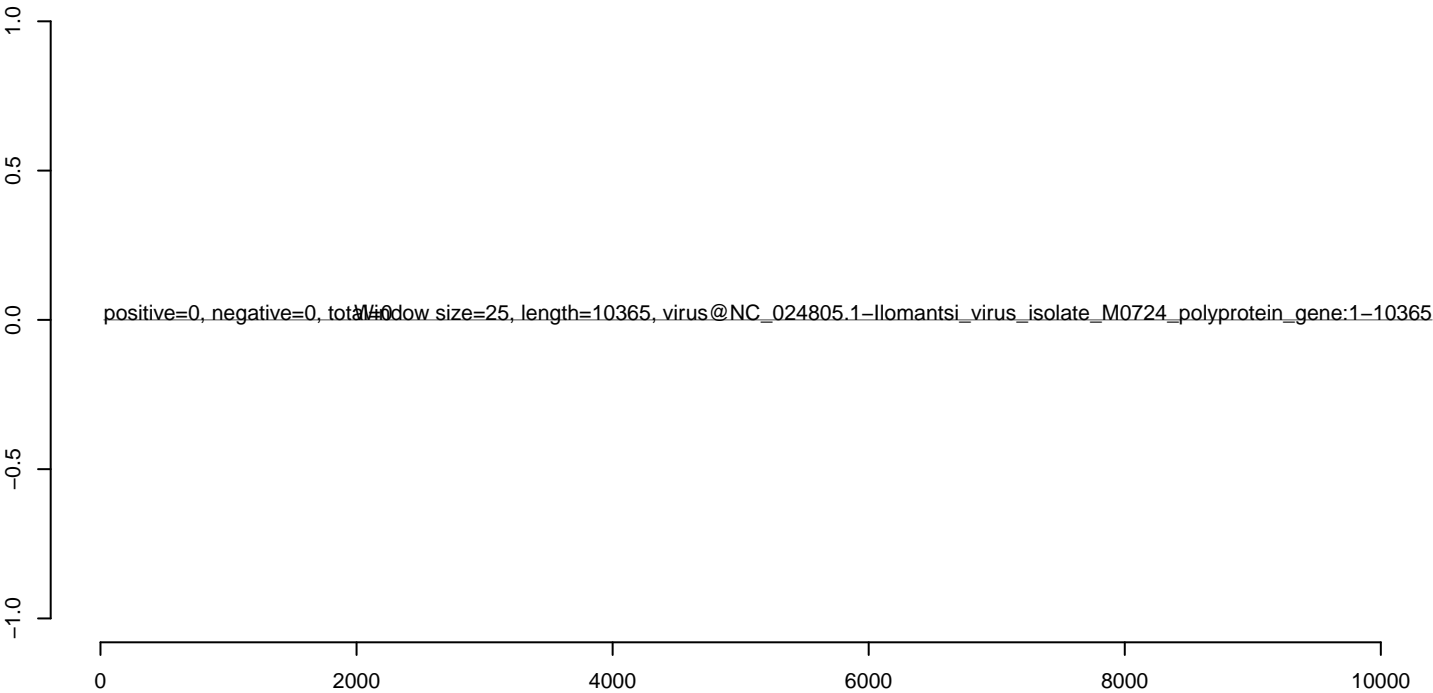
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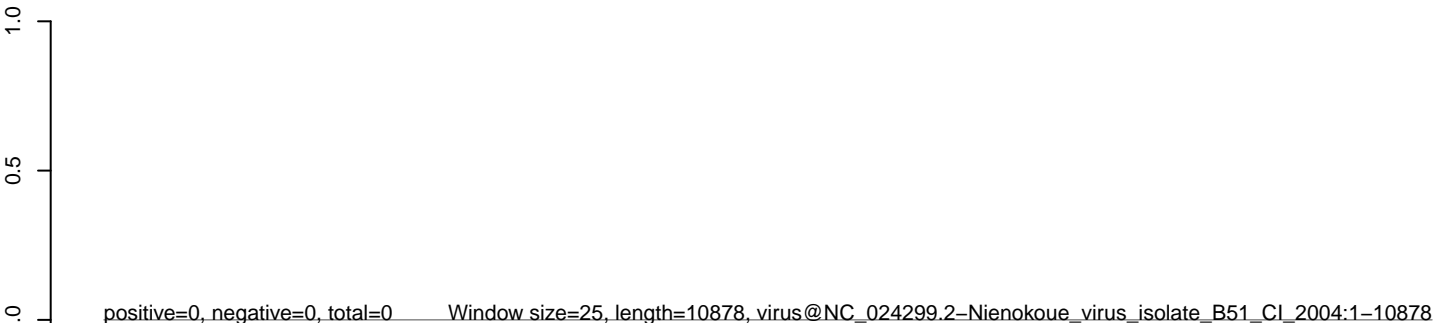
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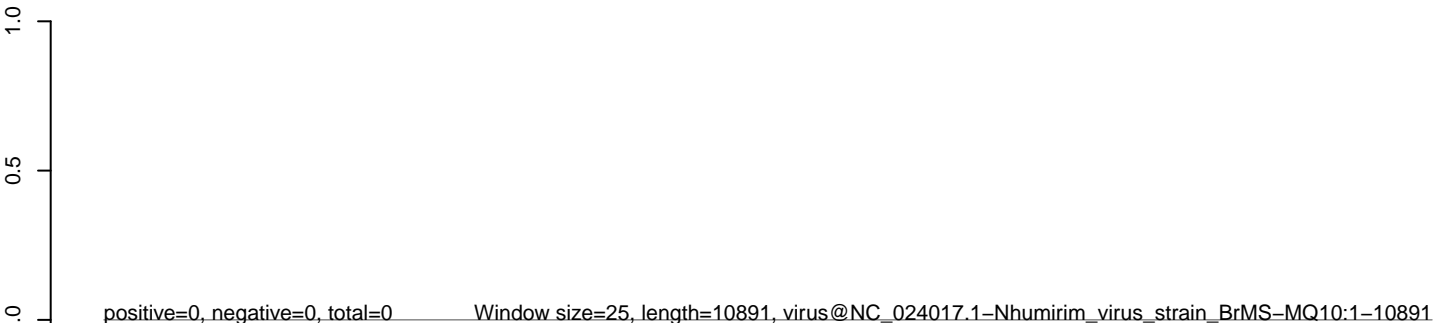
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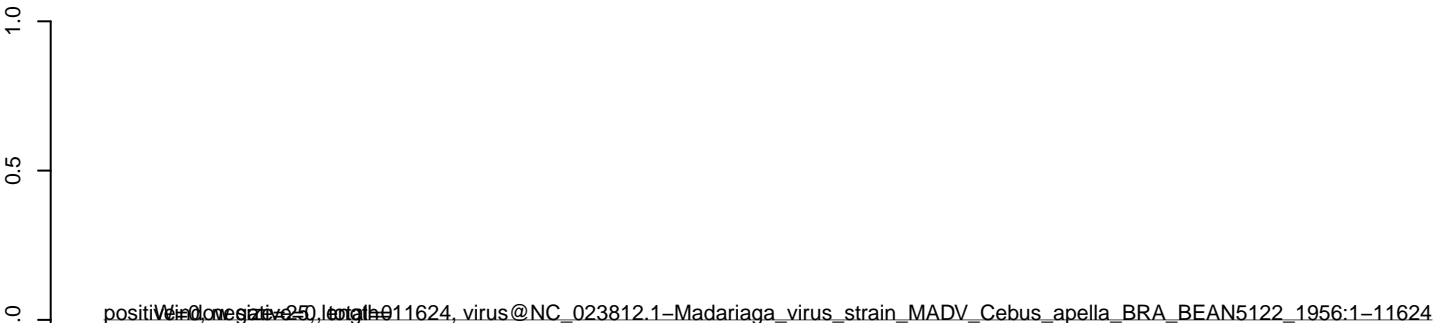
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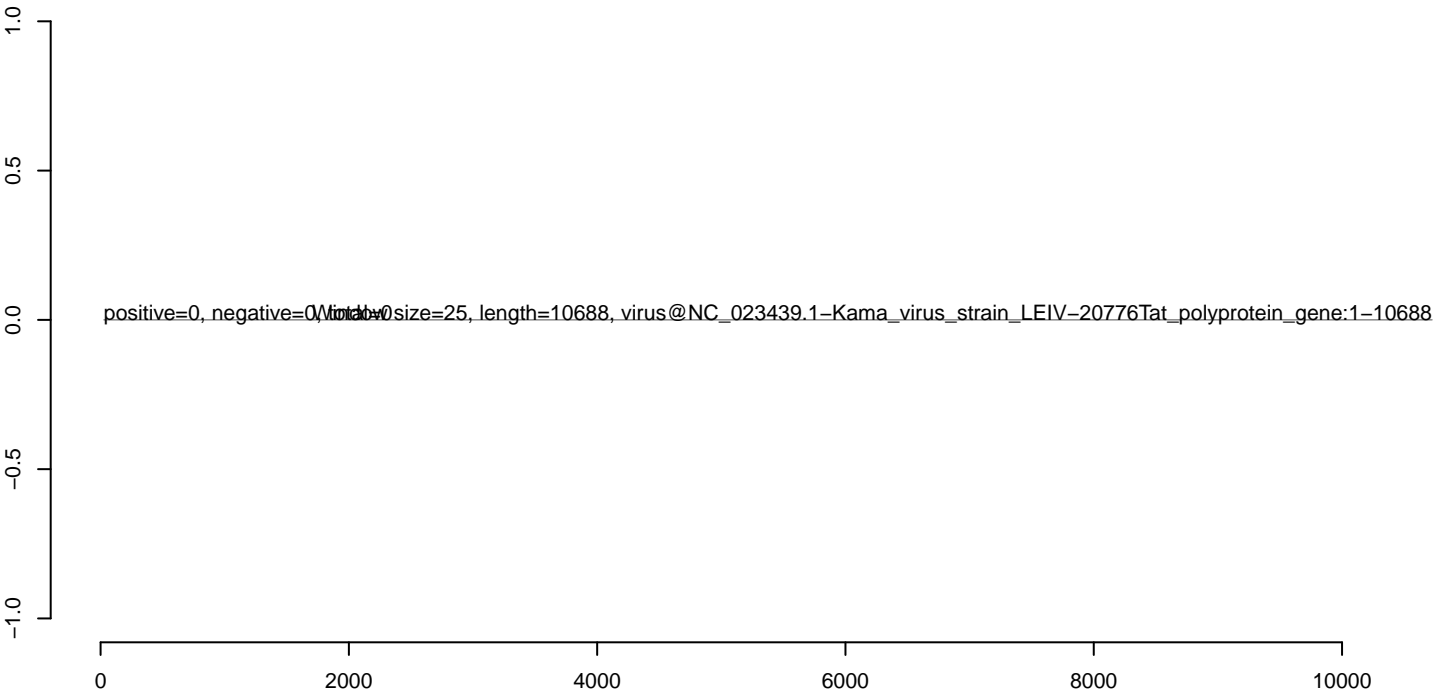
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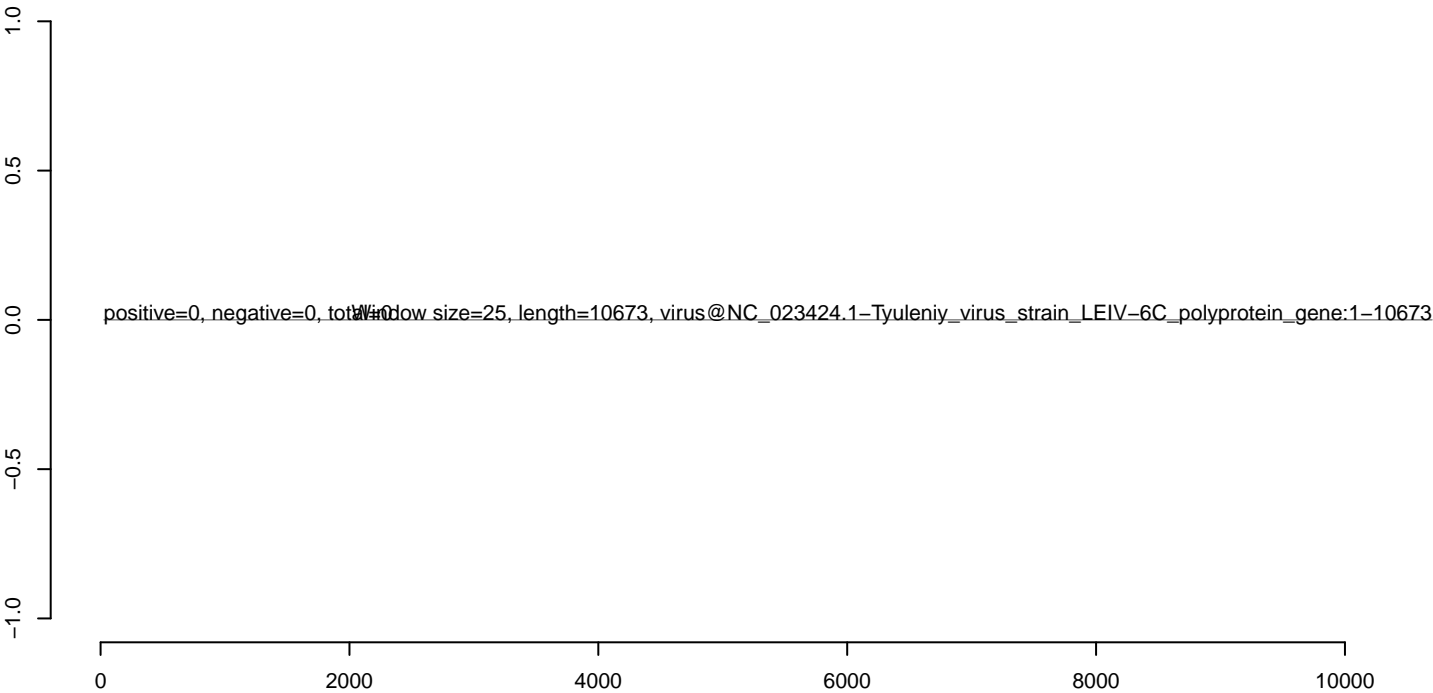
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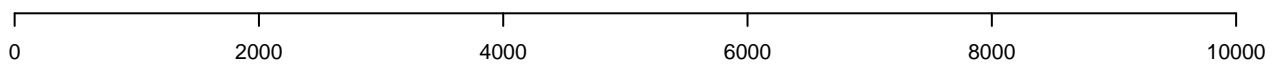
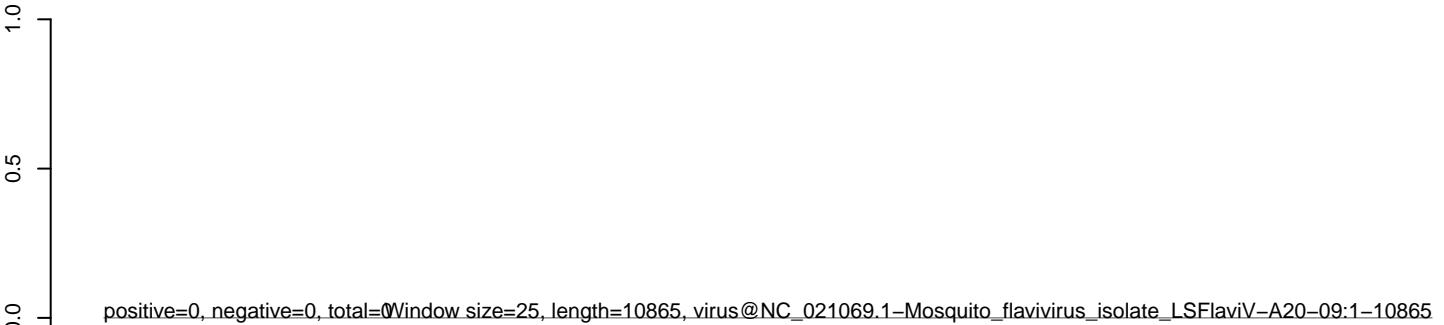
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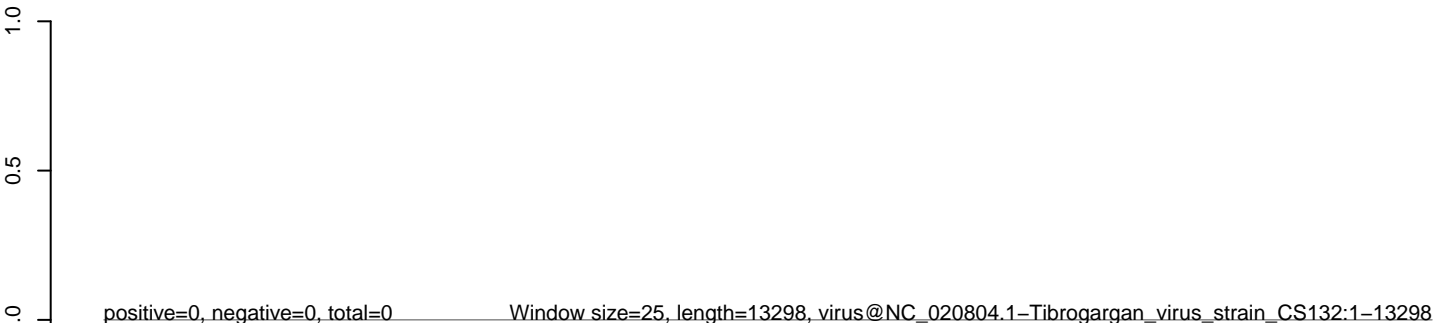
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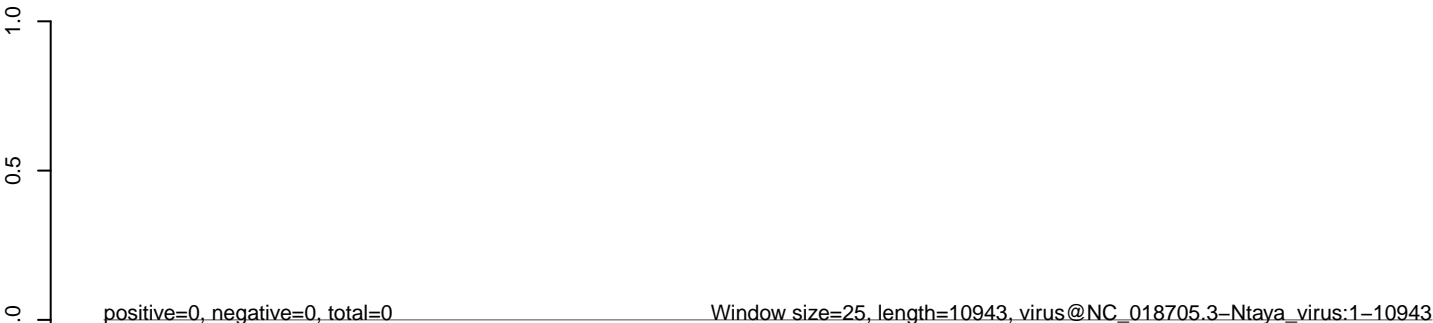
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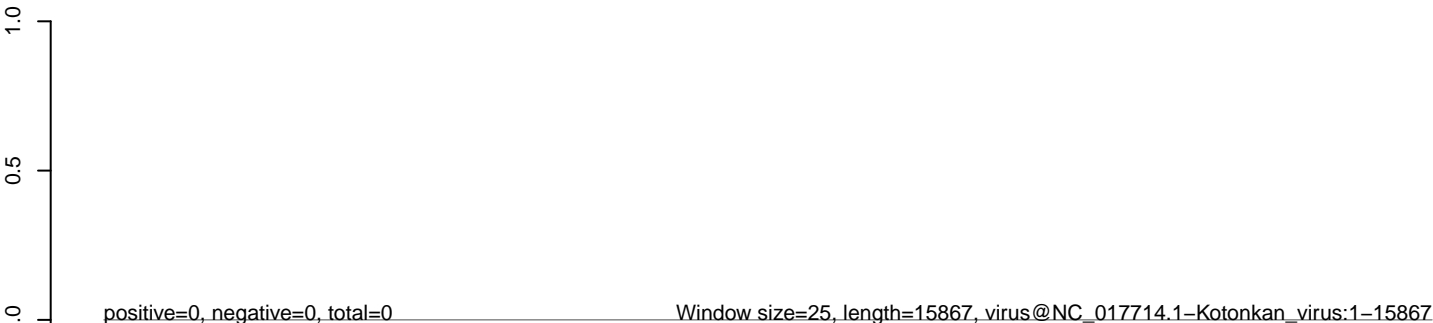
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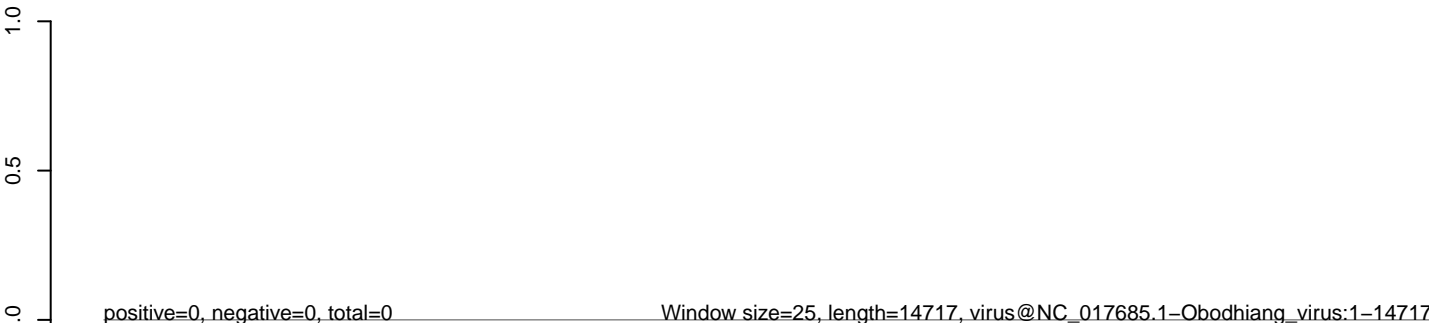
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AeAlbo_C636_SINV_GFP_betaE.rep



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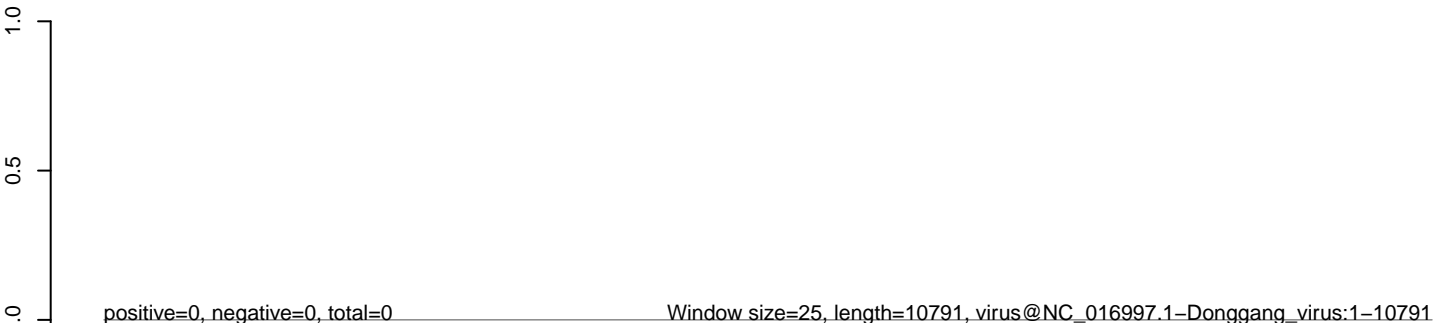
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AeAlbo_C636_SINV_GFP_betaE.rep



0 2000 4000 6000 8000 10000

AeAlbo_C636_SINV_GFP_betaE.18_23.rep



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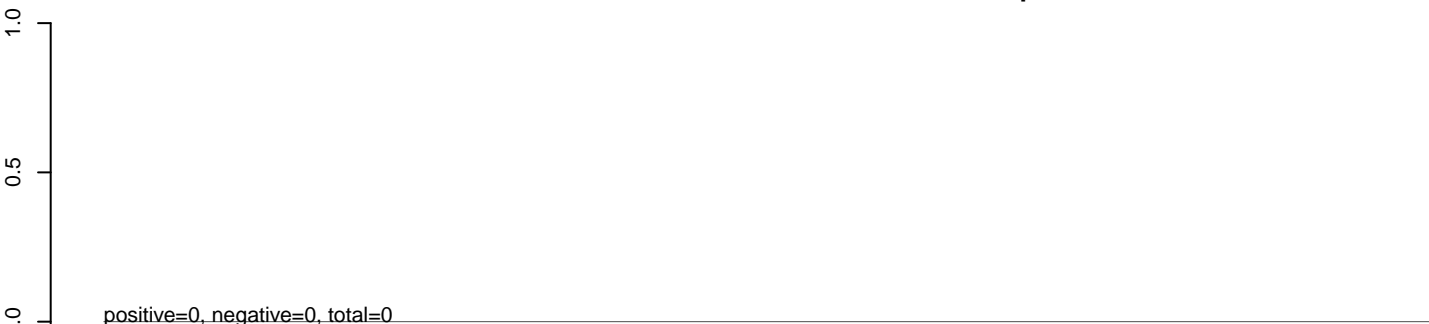


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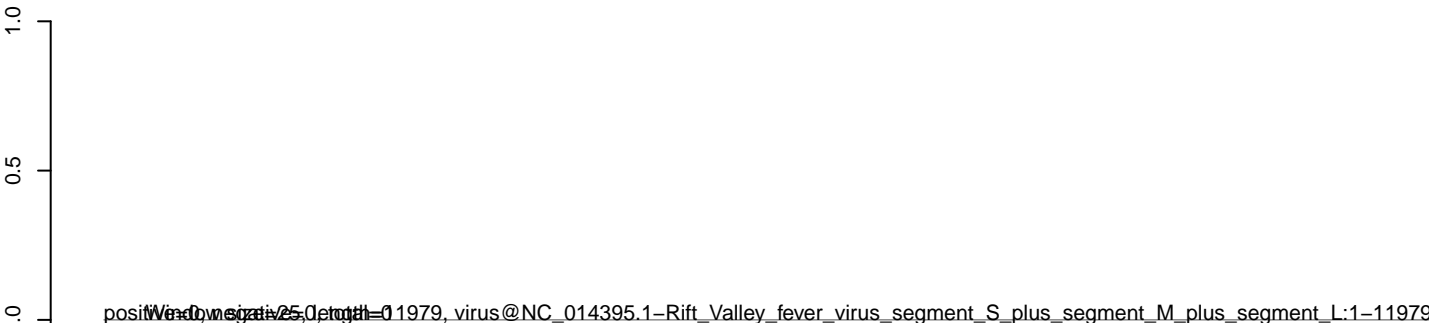
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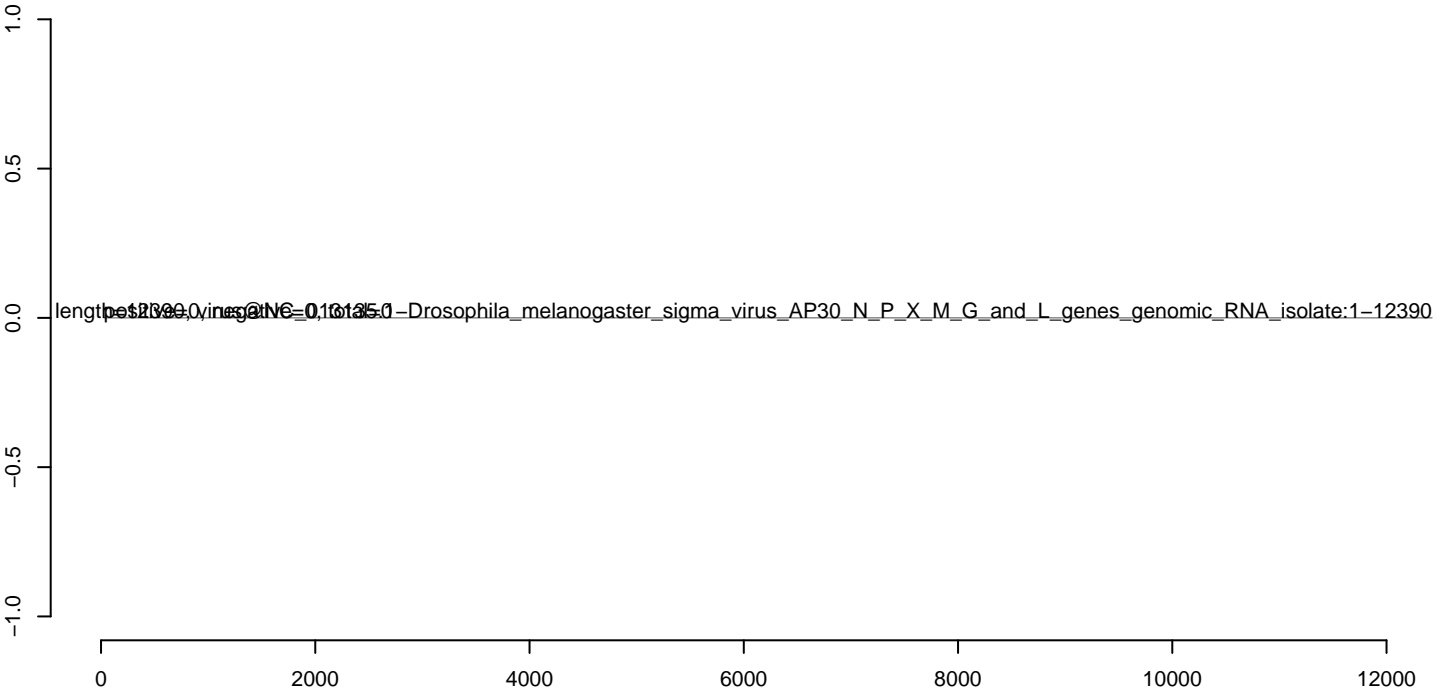
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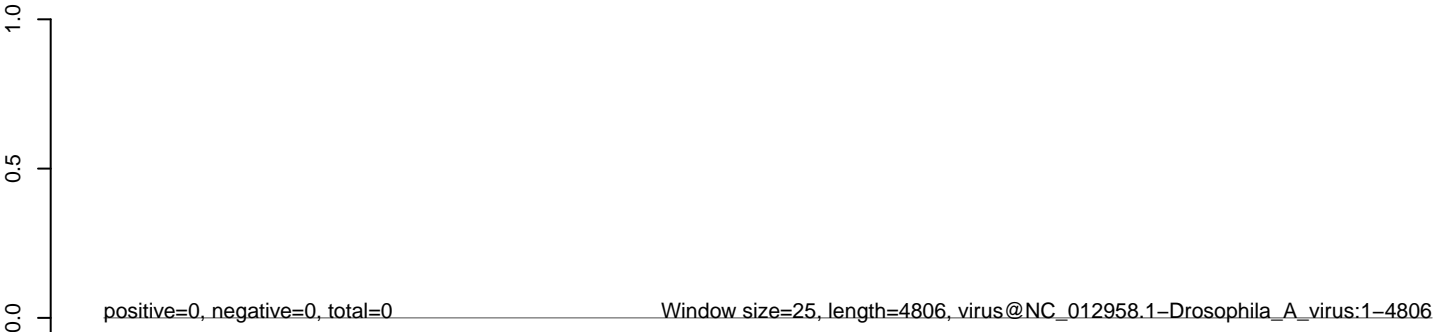
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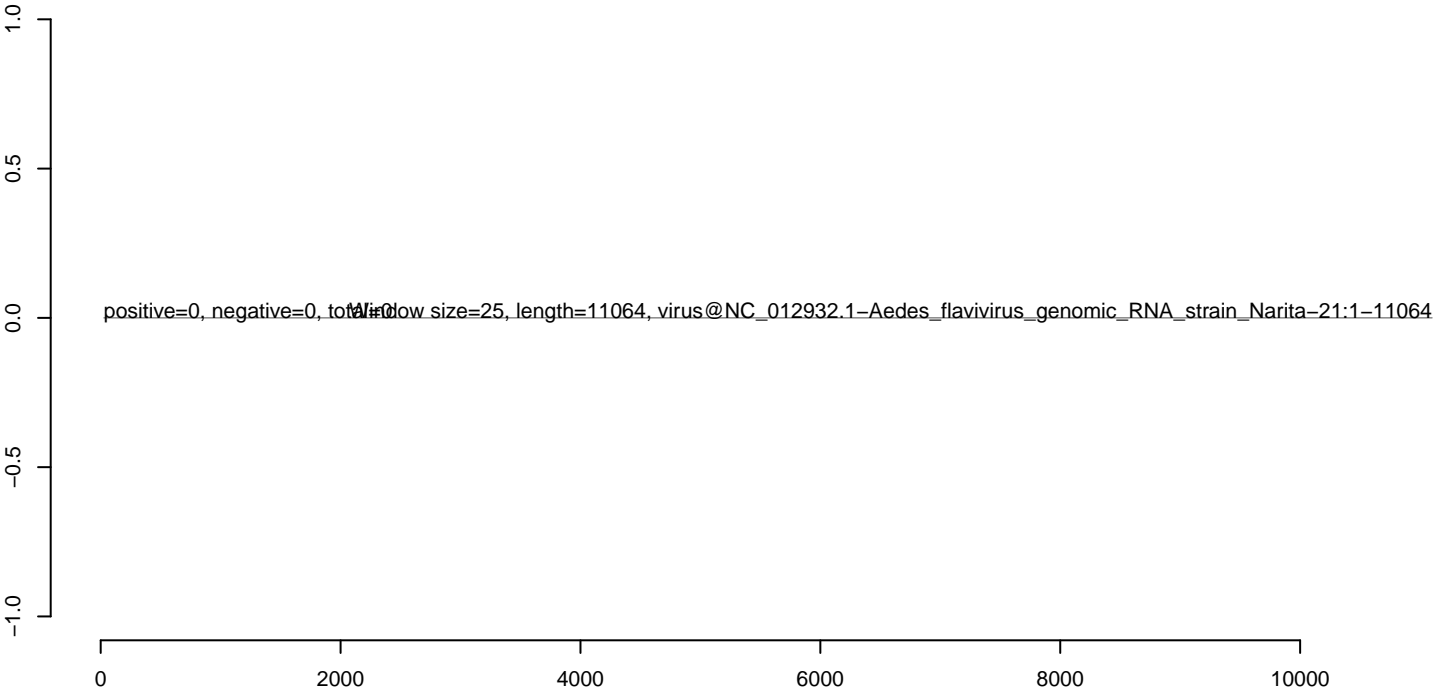
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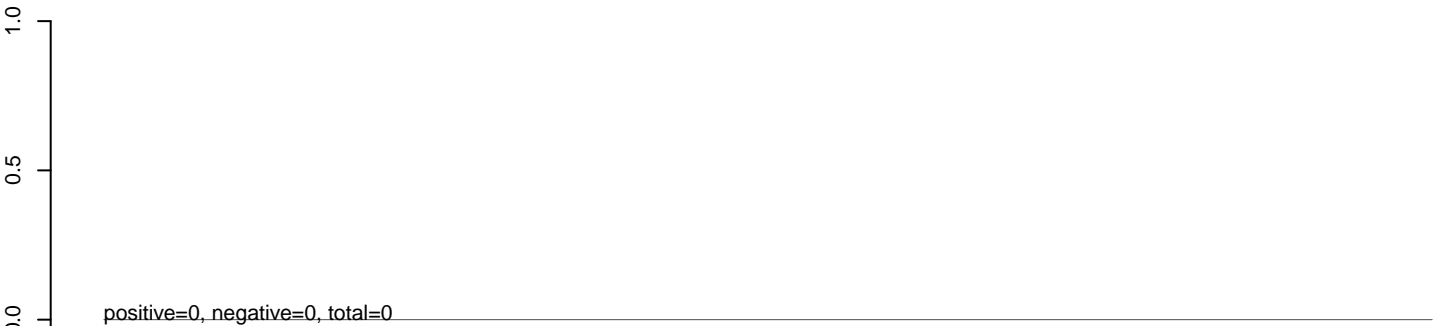
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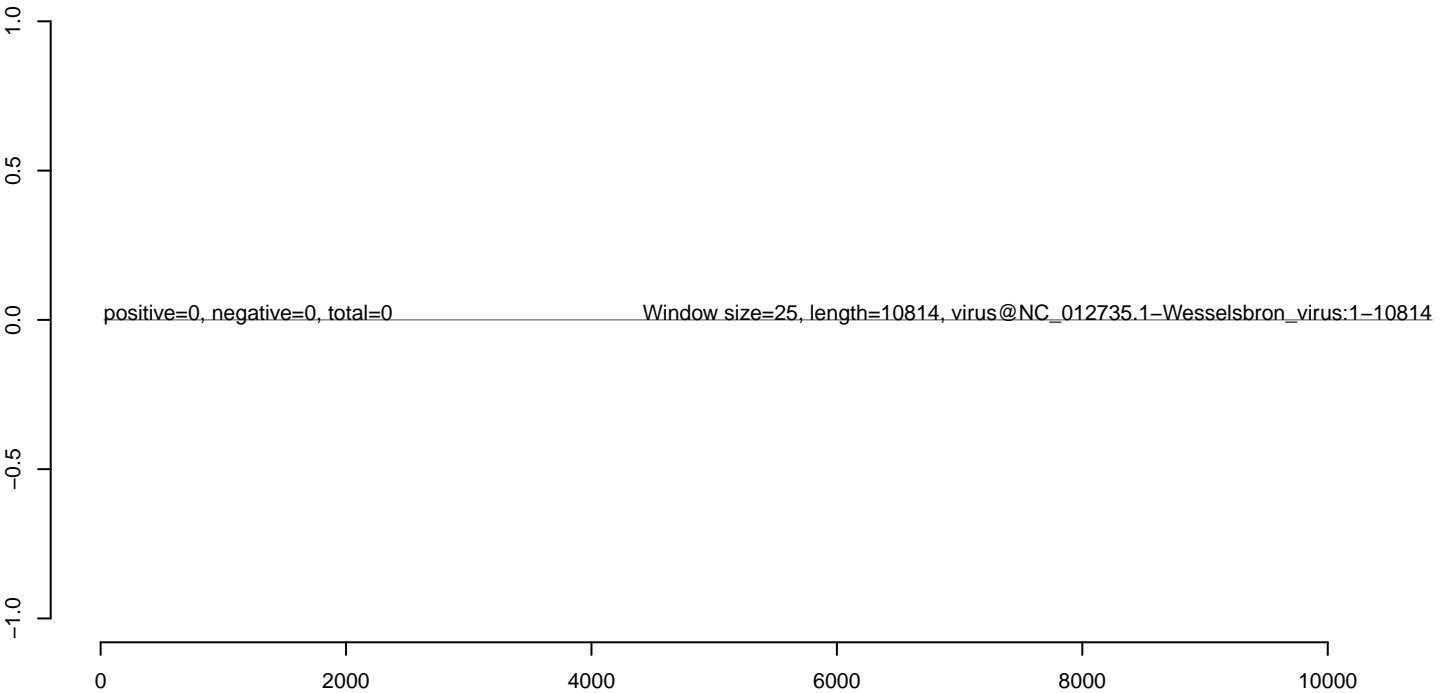
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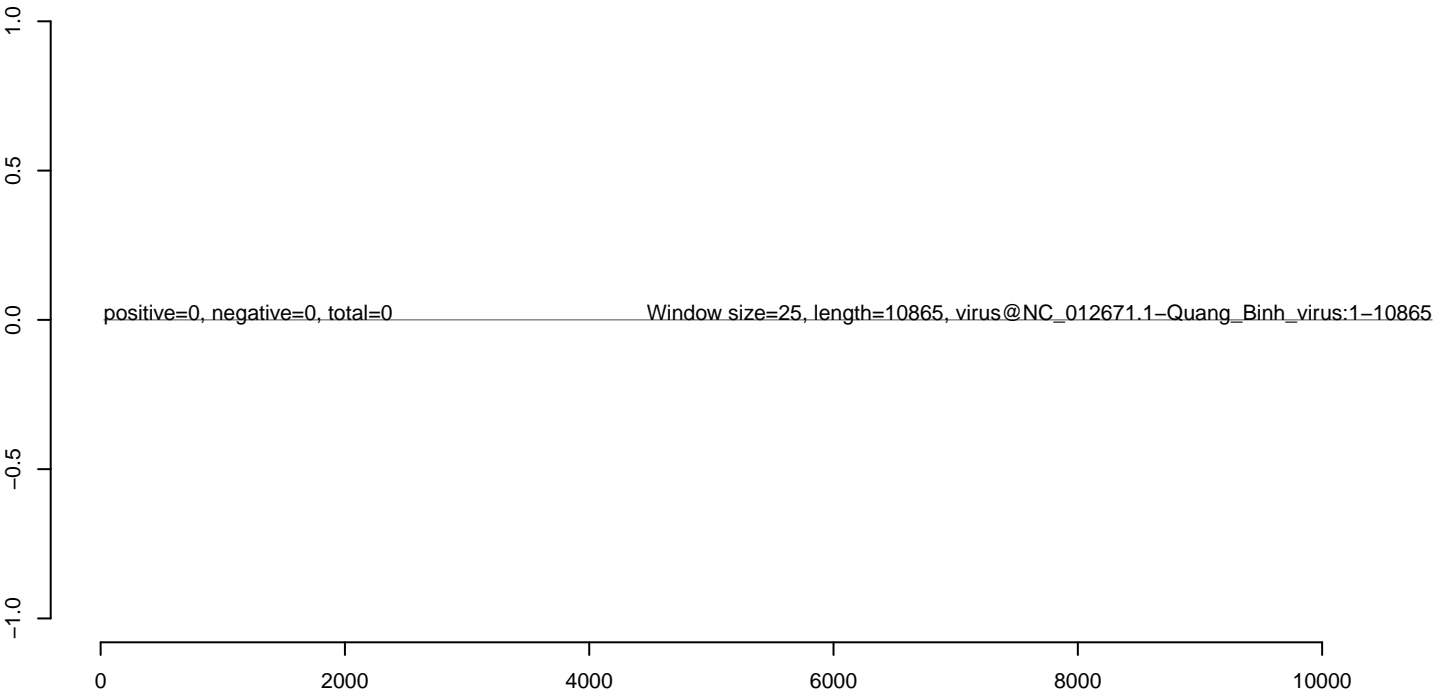
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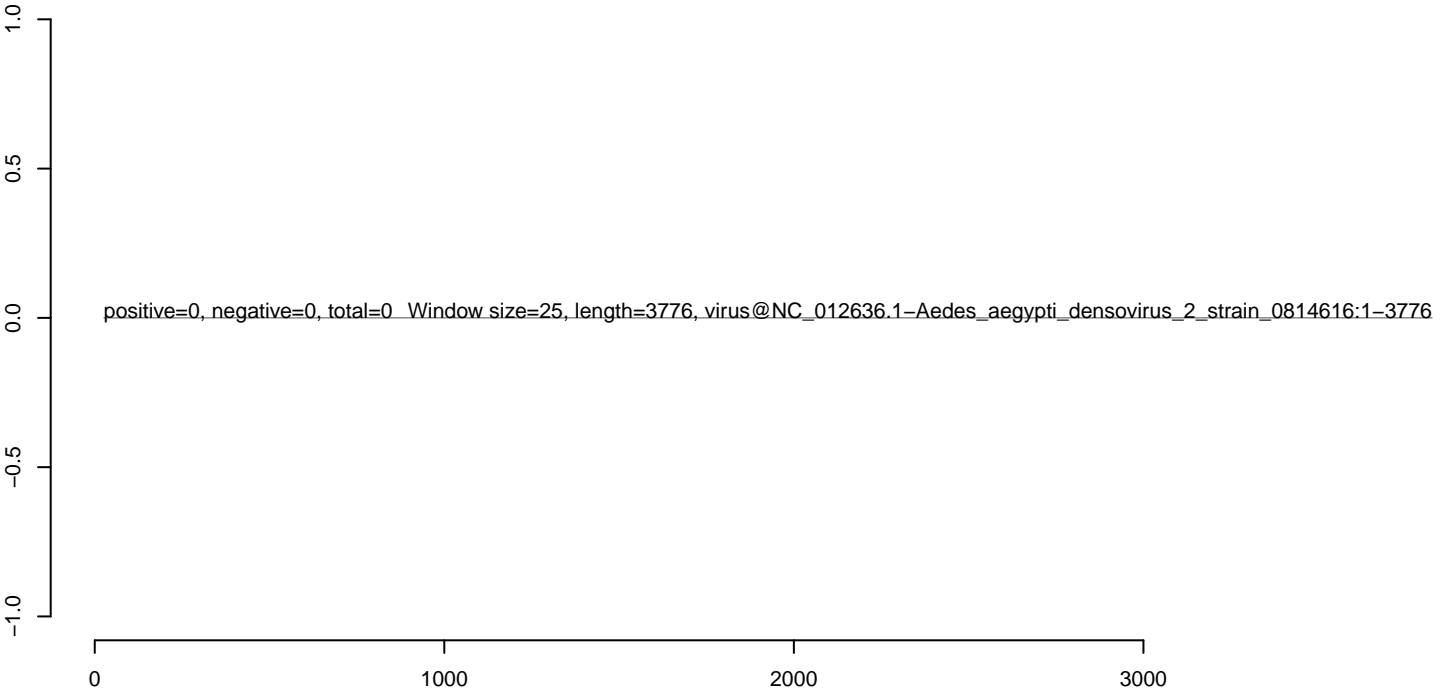
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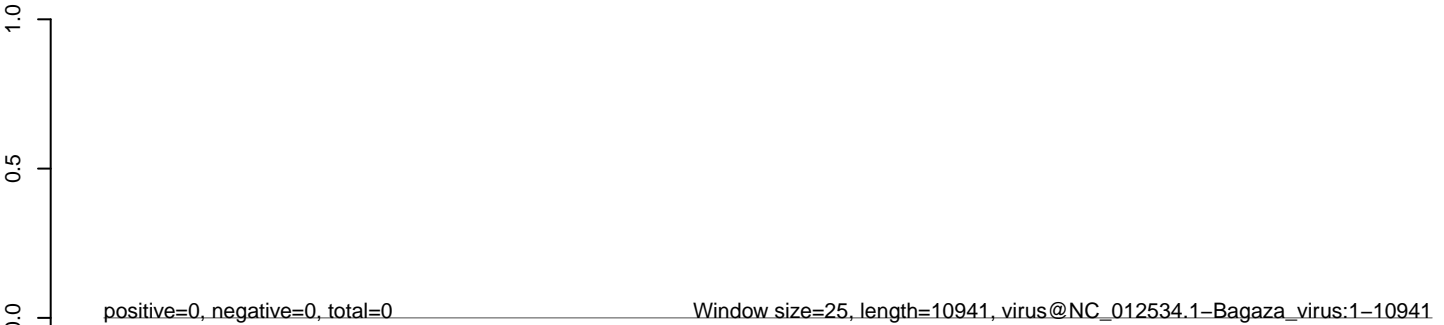
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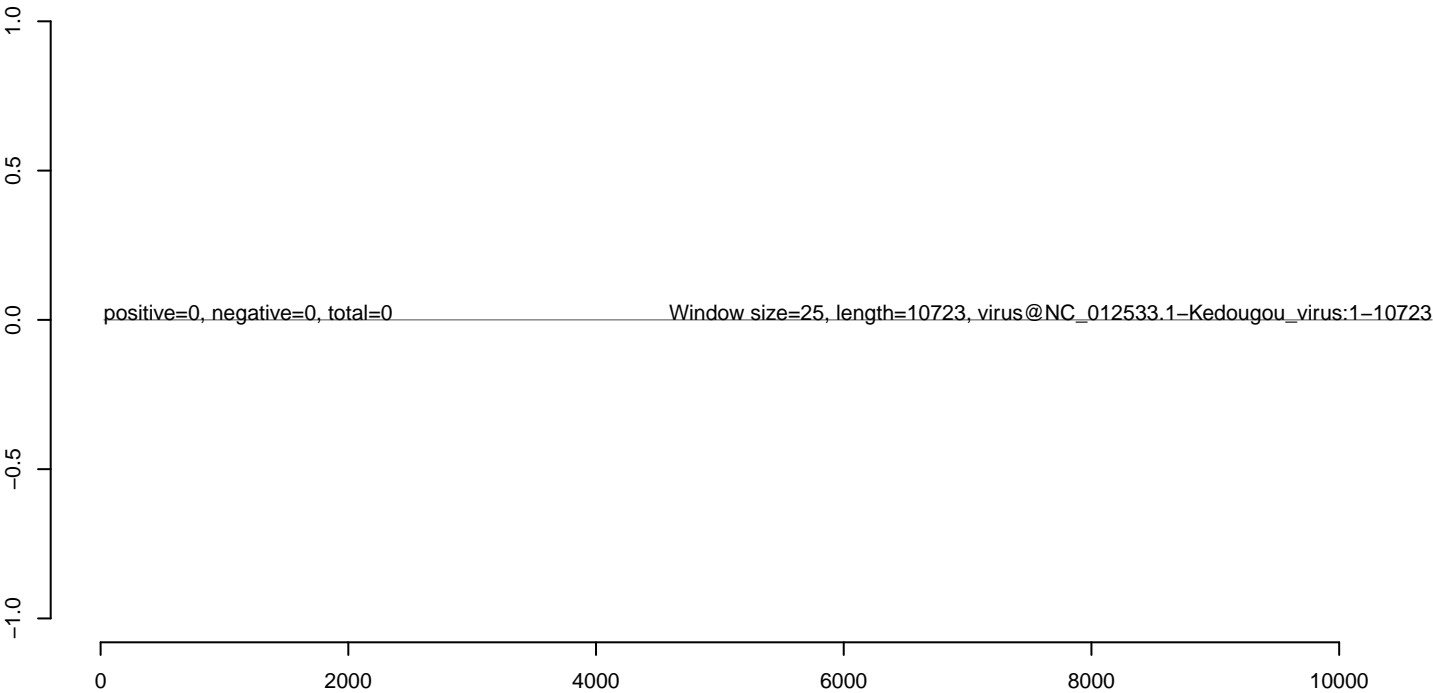
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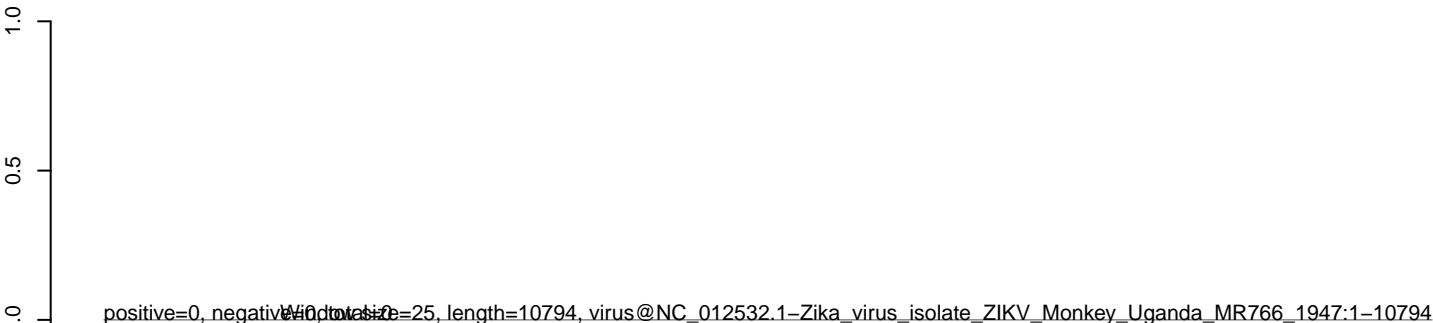
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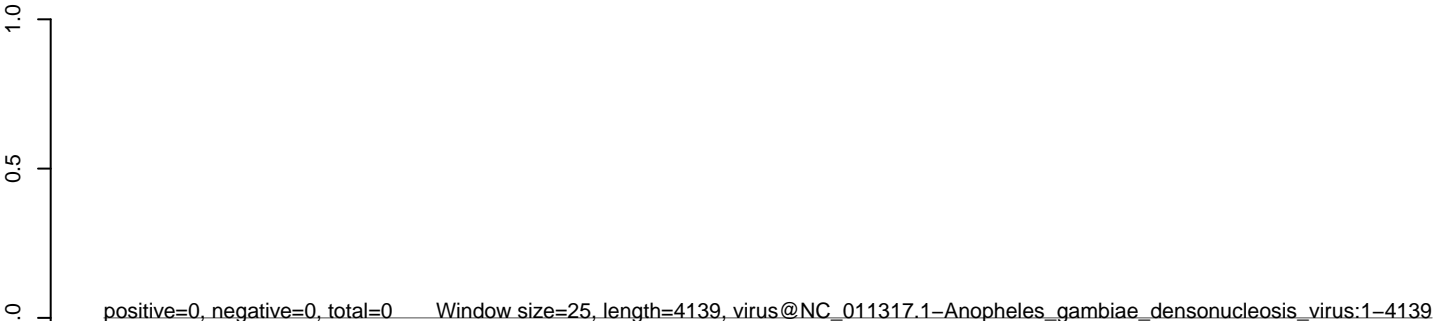
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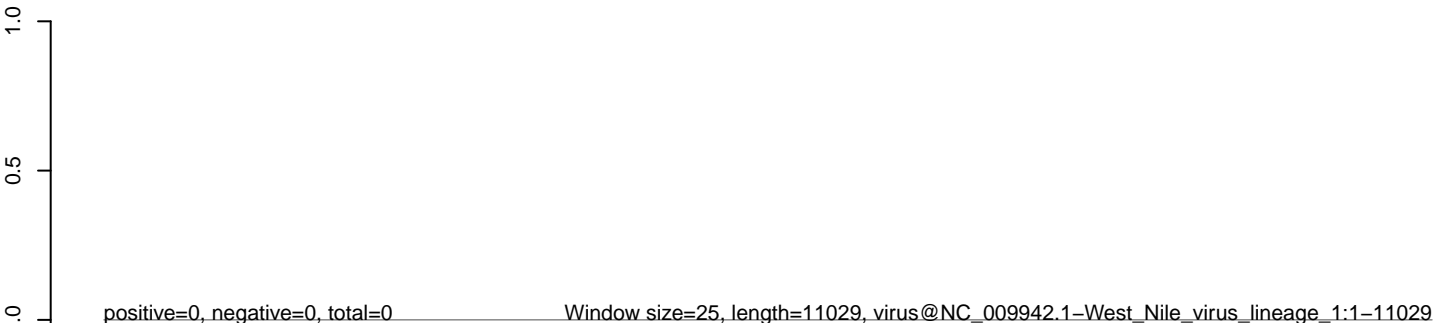
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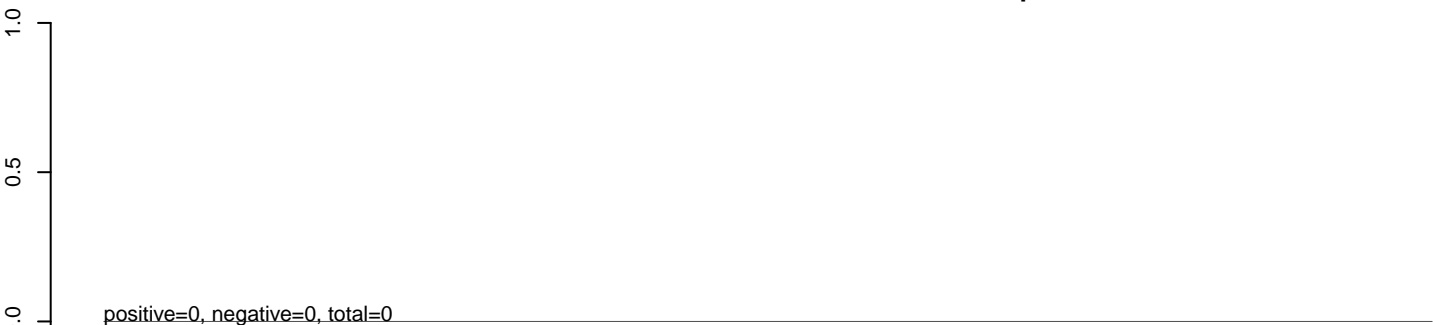


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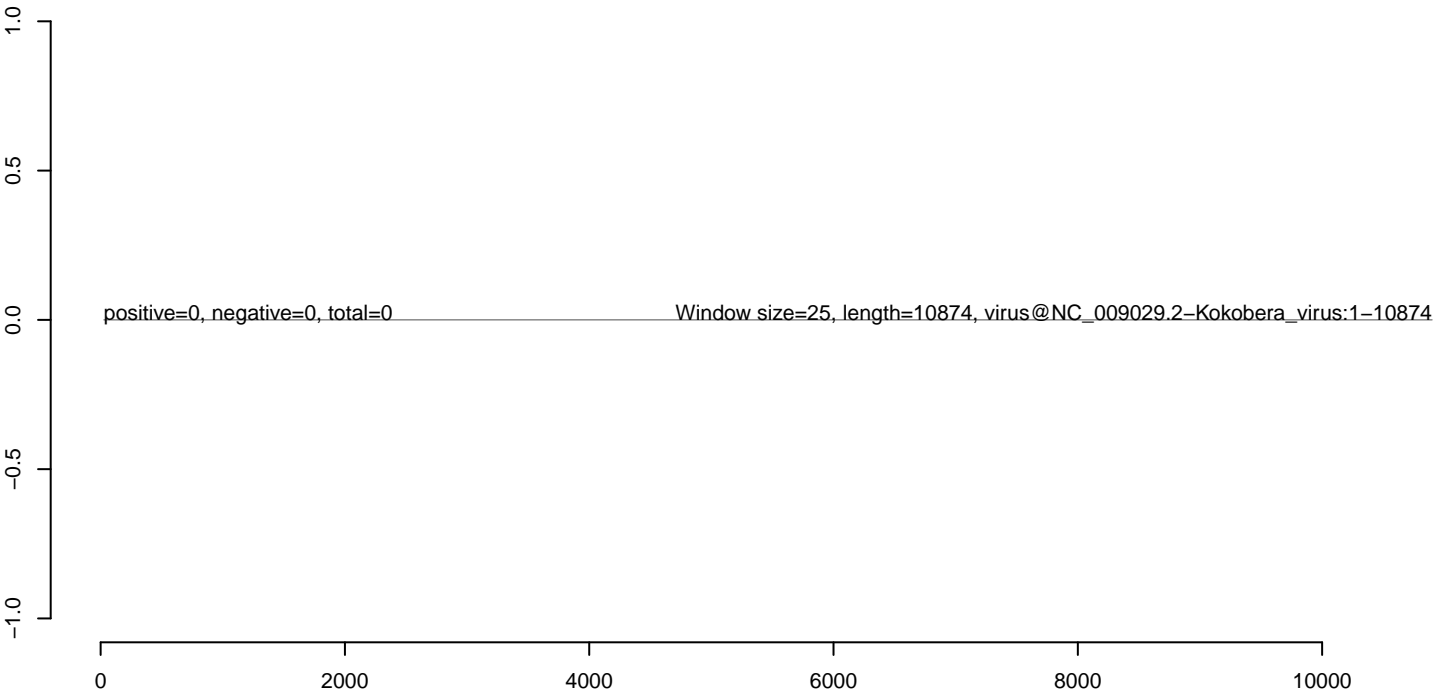
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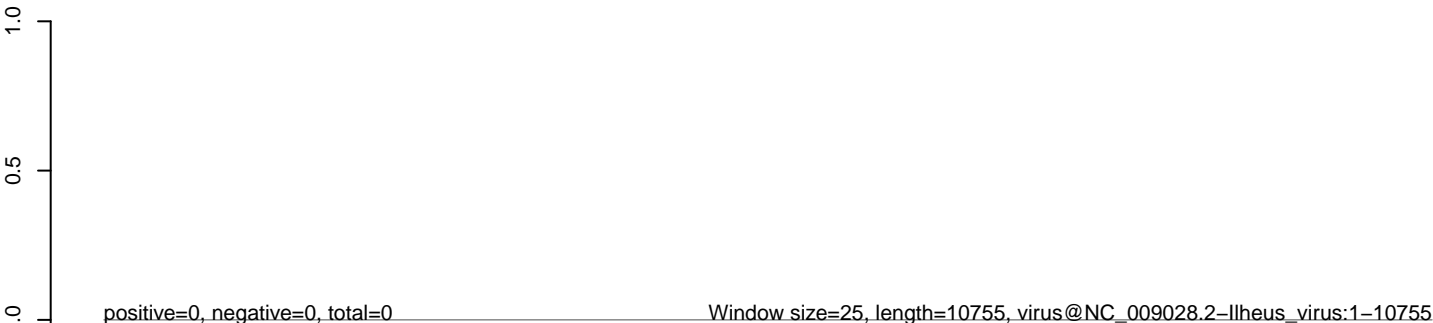
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AeAlbo_C636_SINV_GFP_betaE.rep



0 2000 4000 6000 8000 10000

AeAlbo_C636_SINV_GFP_betaE.18_23.rep



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AeAlbo_C636_SINV_GFP_betaE.rep



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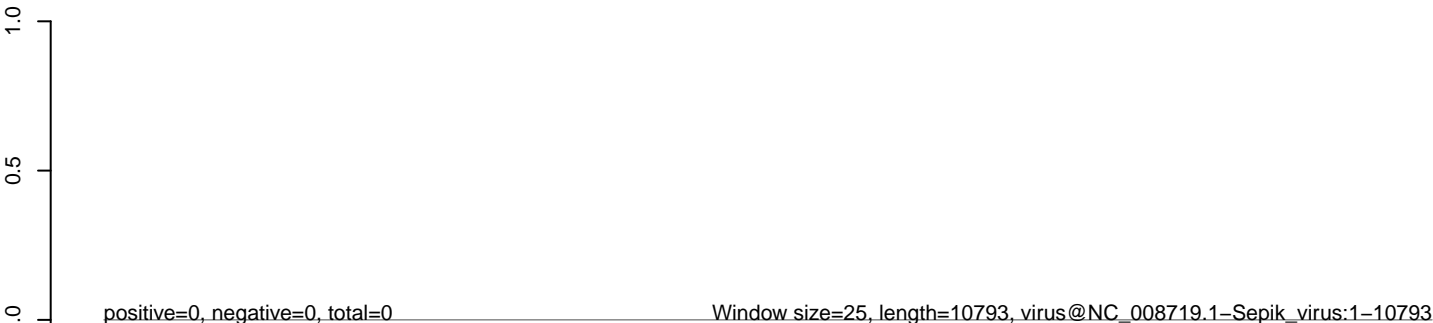
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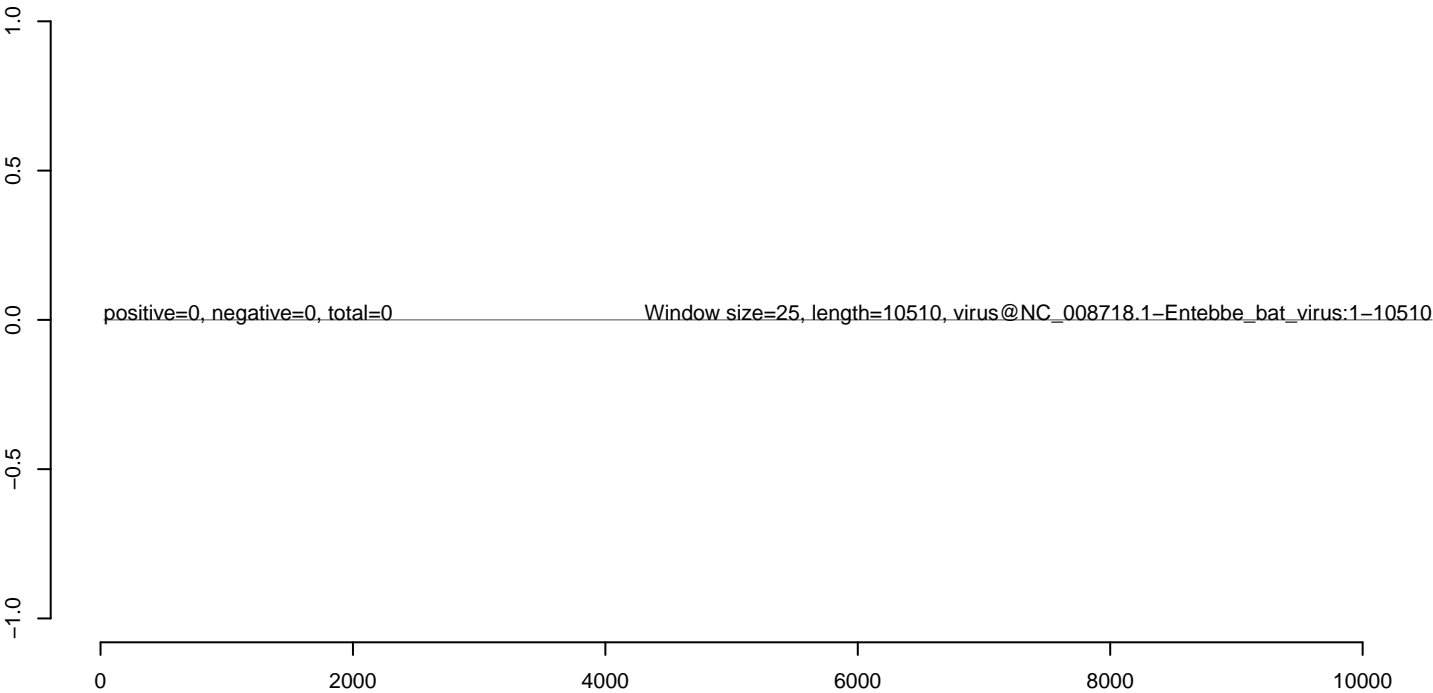
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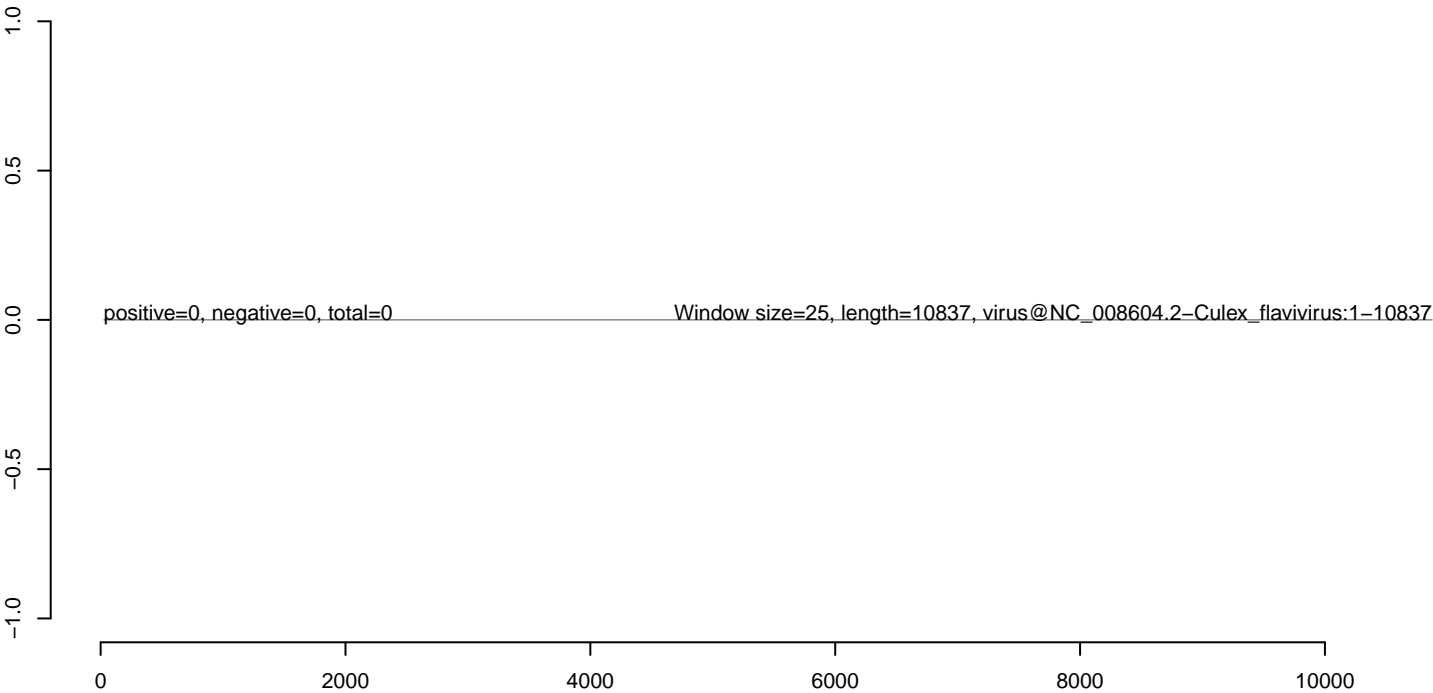
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AeAlbo_C636_SINV_GFP_betaE.rep



AeAlbo_C636_SINV_GFP_betaE.18_23.rep



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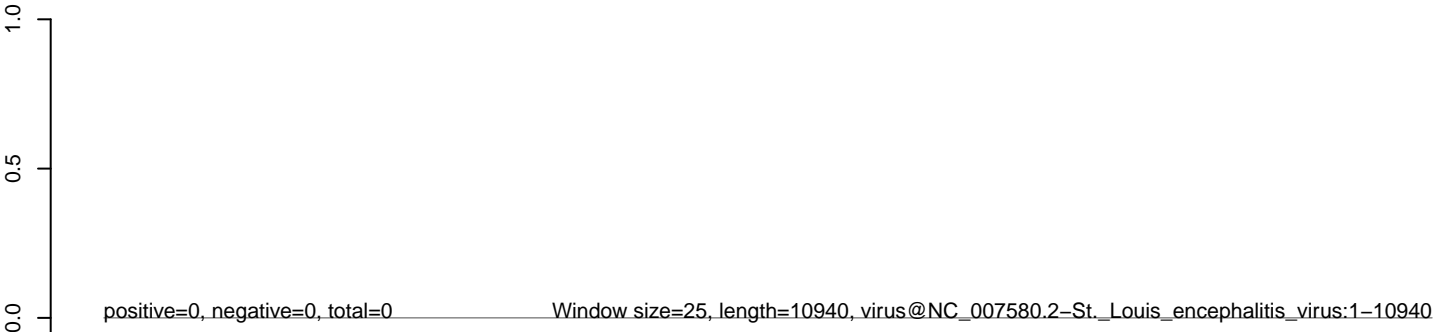
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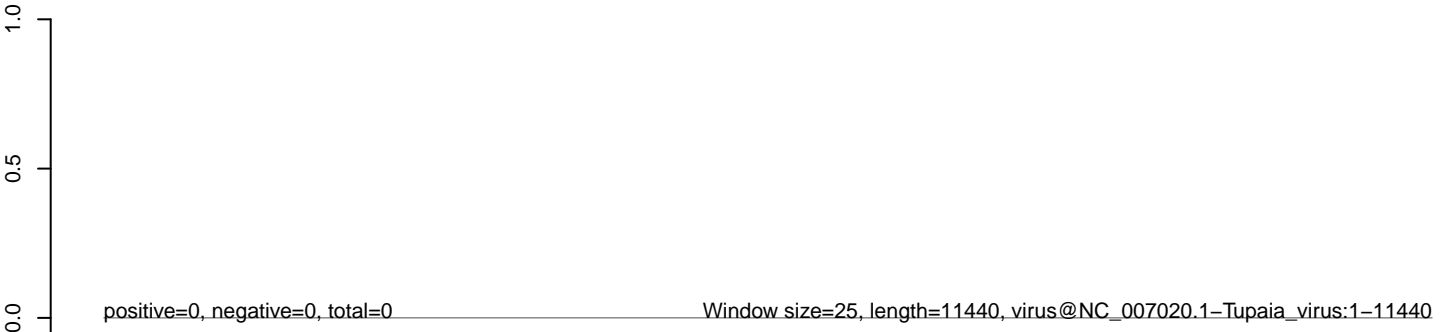
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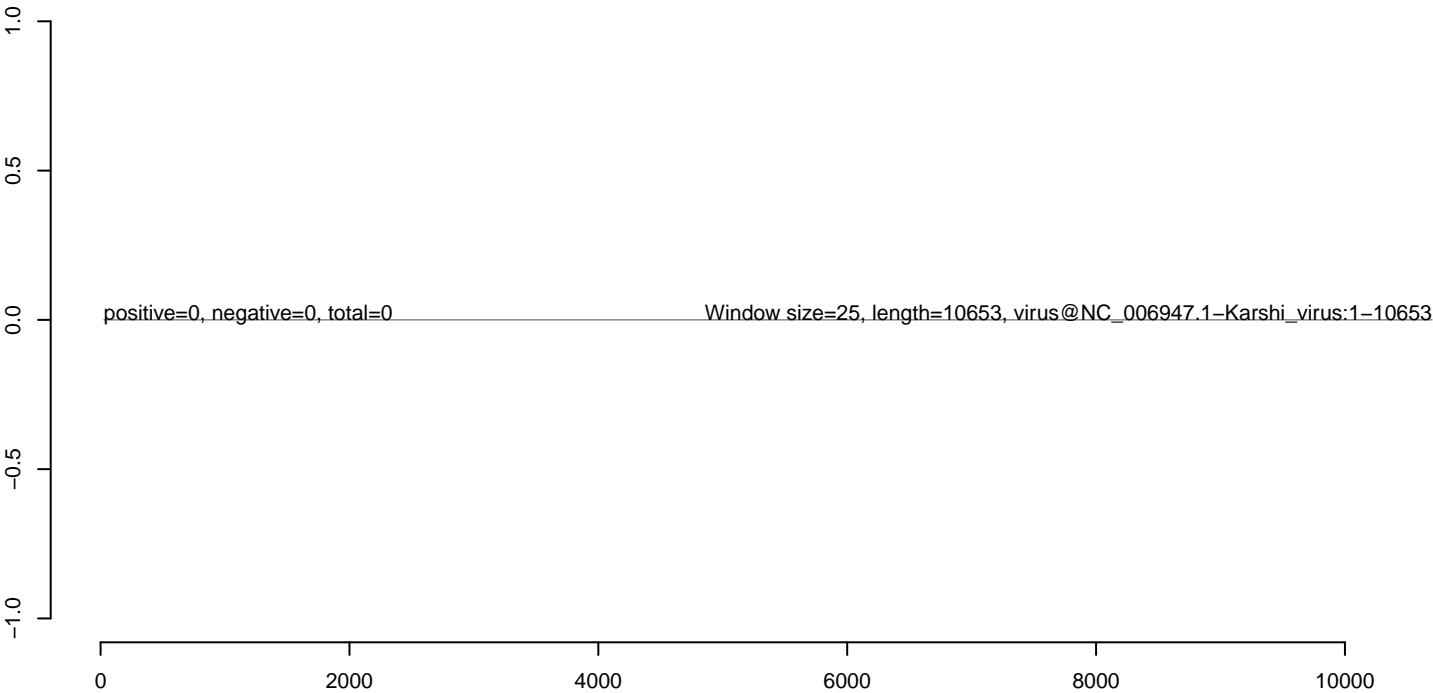
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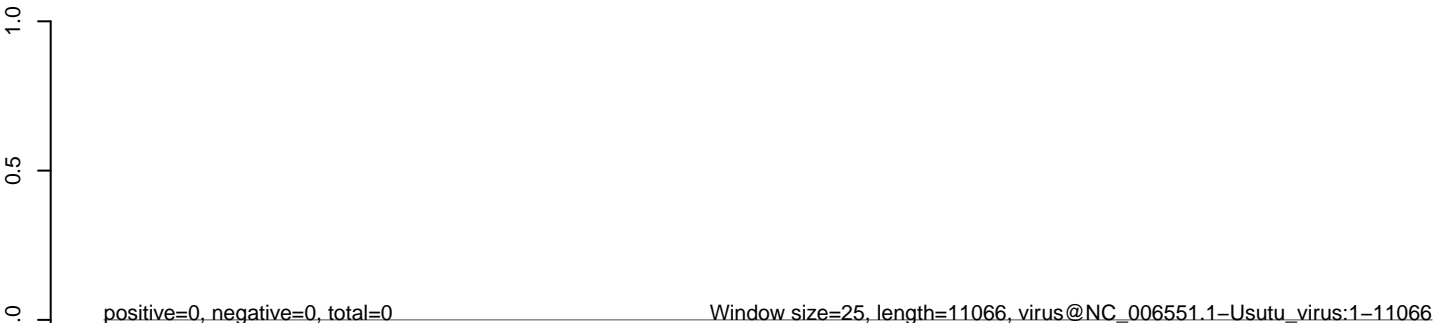
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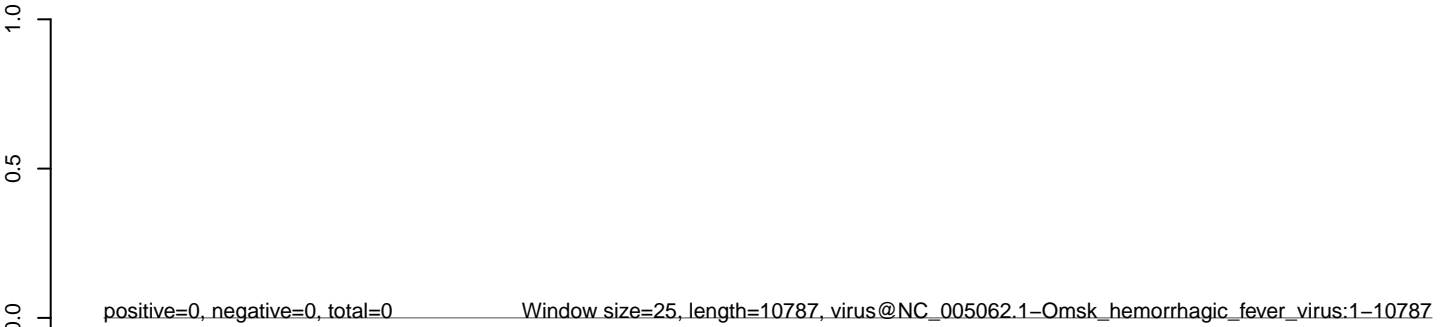
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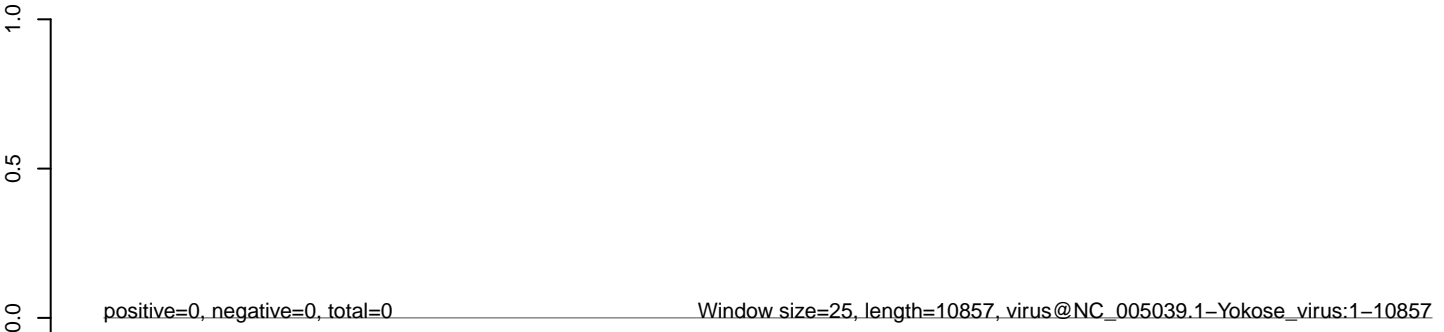
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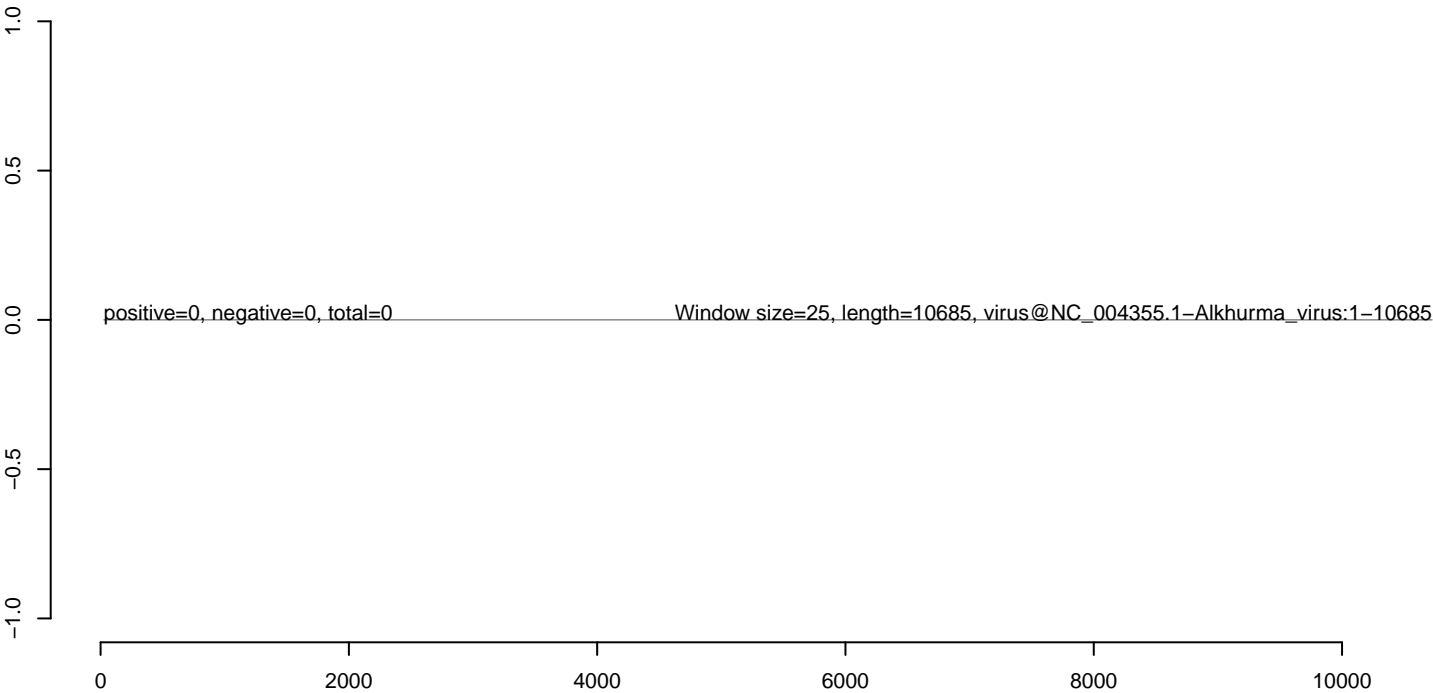
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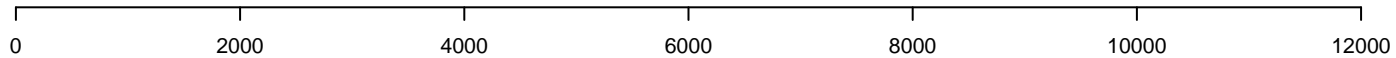
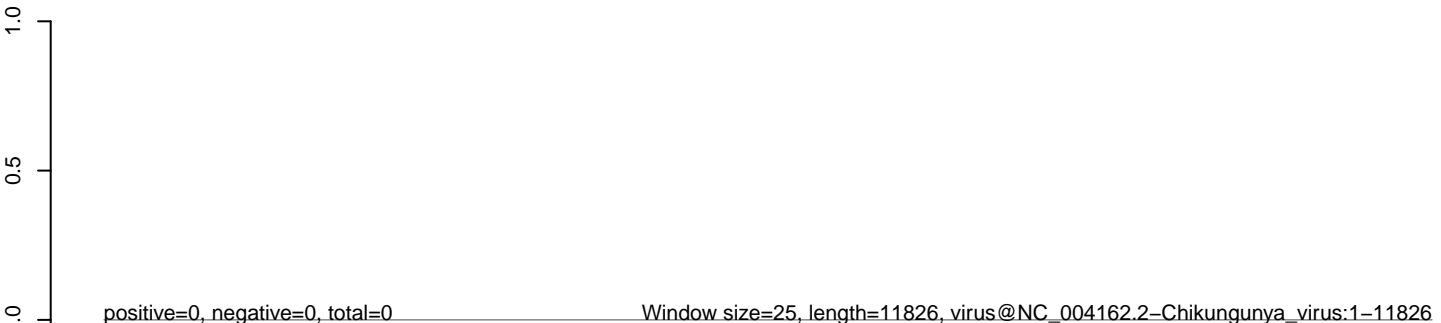
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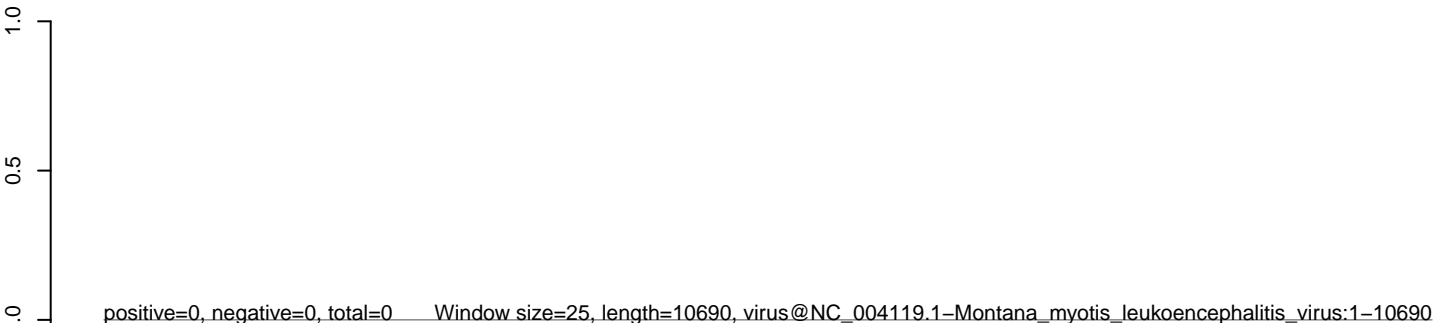
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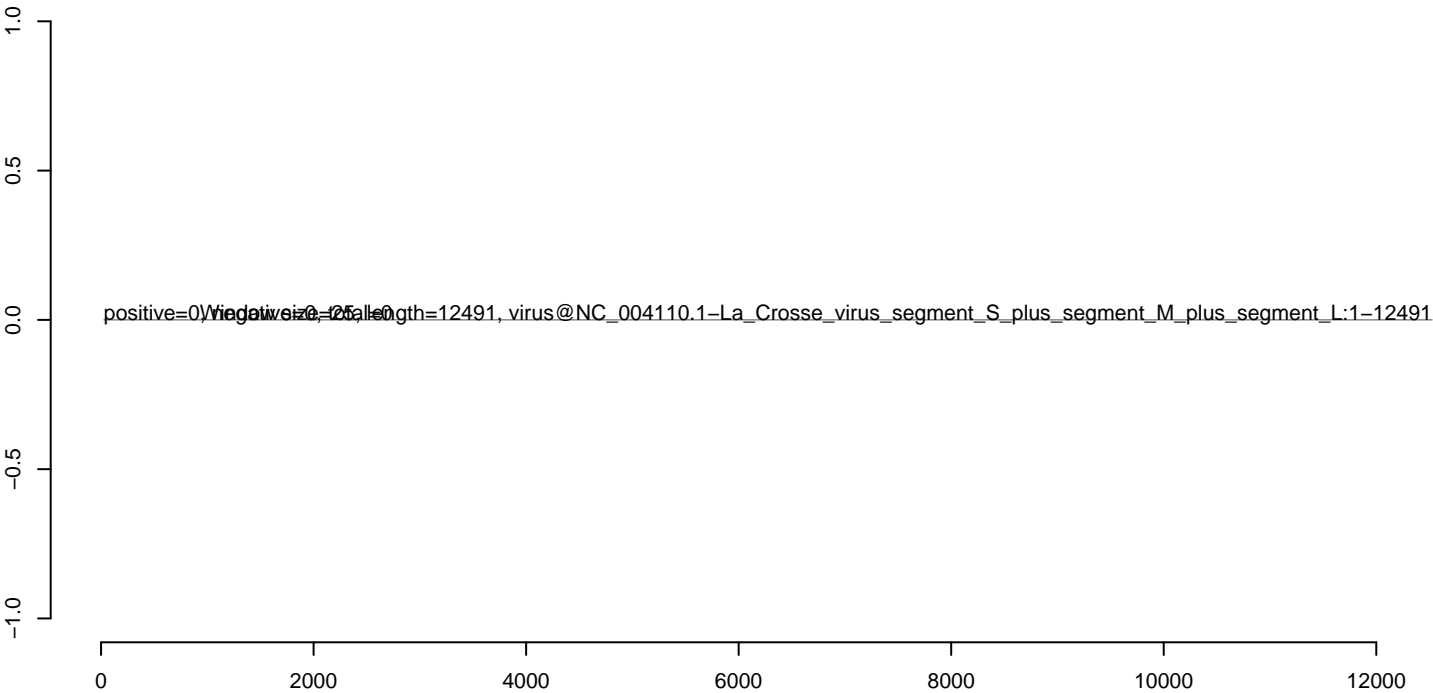
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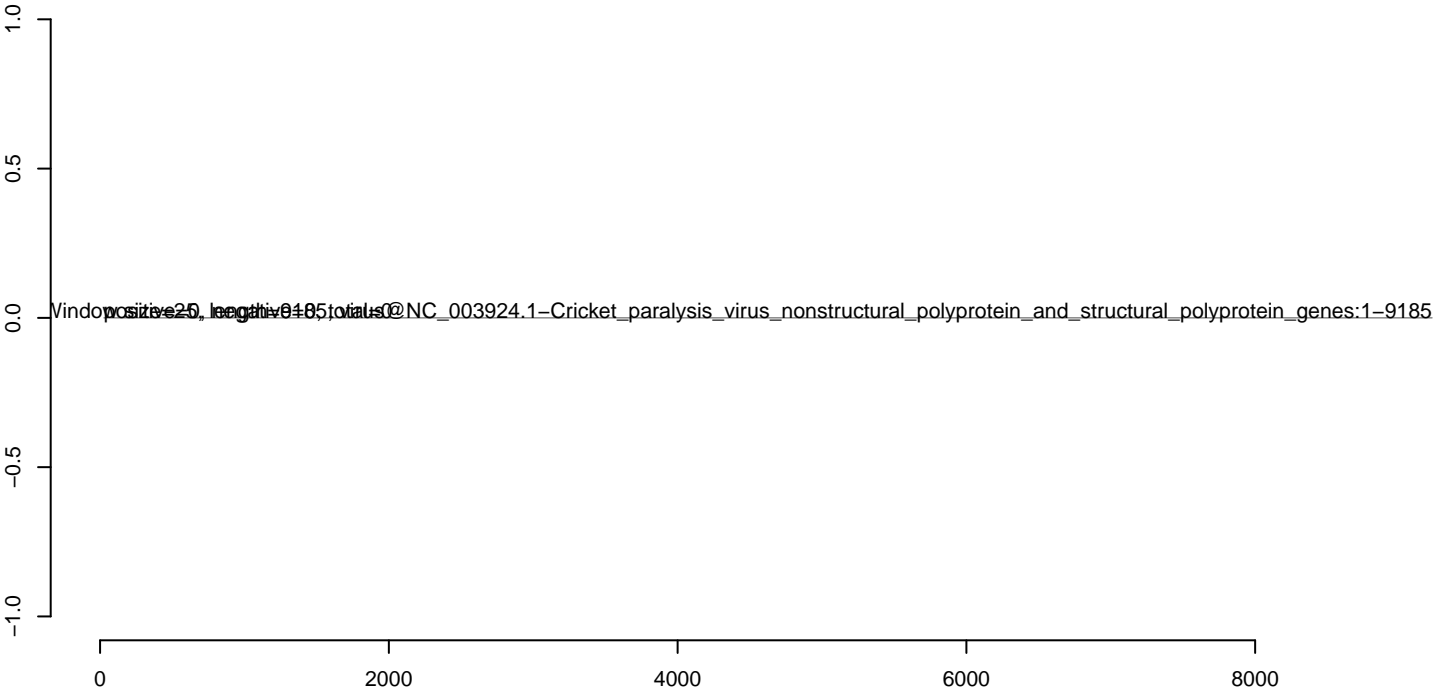
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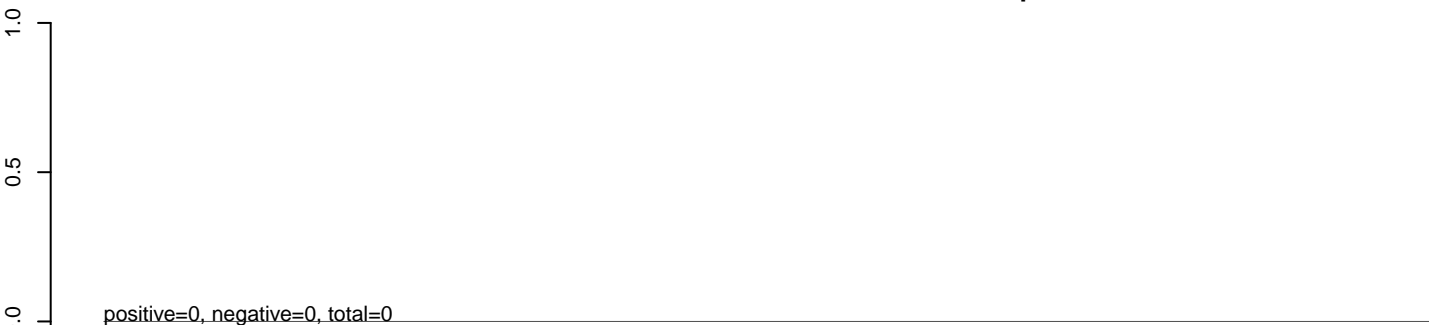
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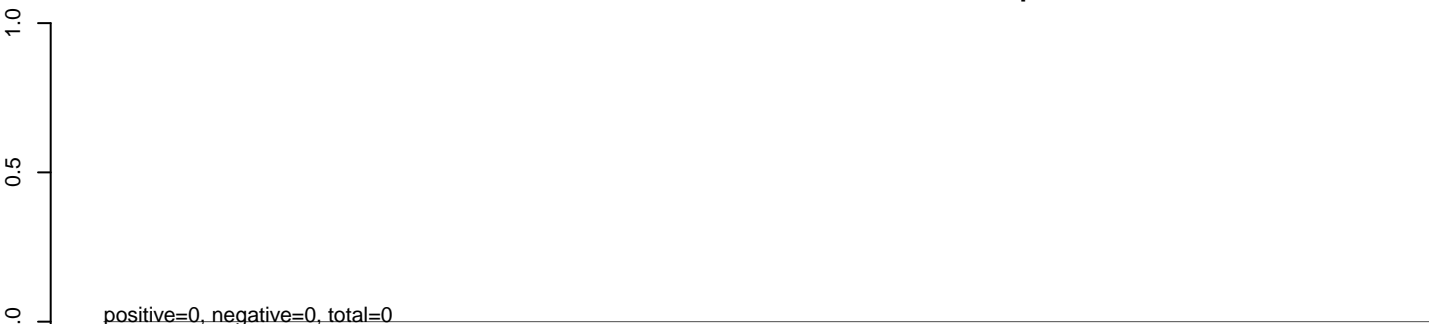
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AeAlbo_C636_SINV_GFP_betaE.rep



0 2000 4000 6000 8000 10000 12000

AeAlbo_C636_SINV_GFP_betaE.18_23.rep



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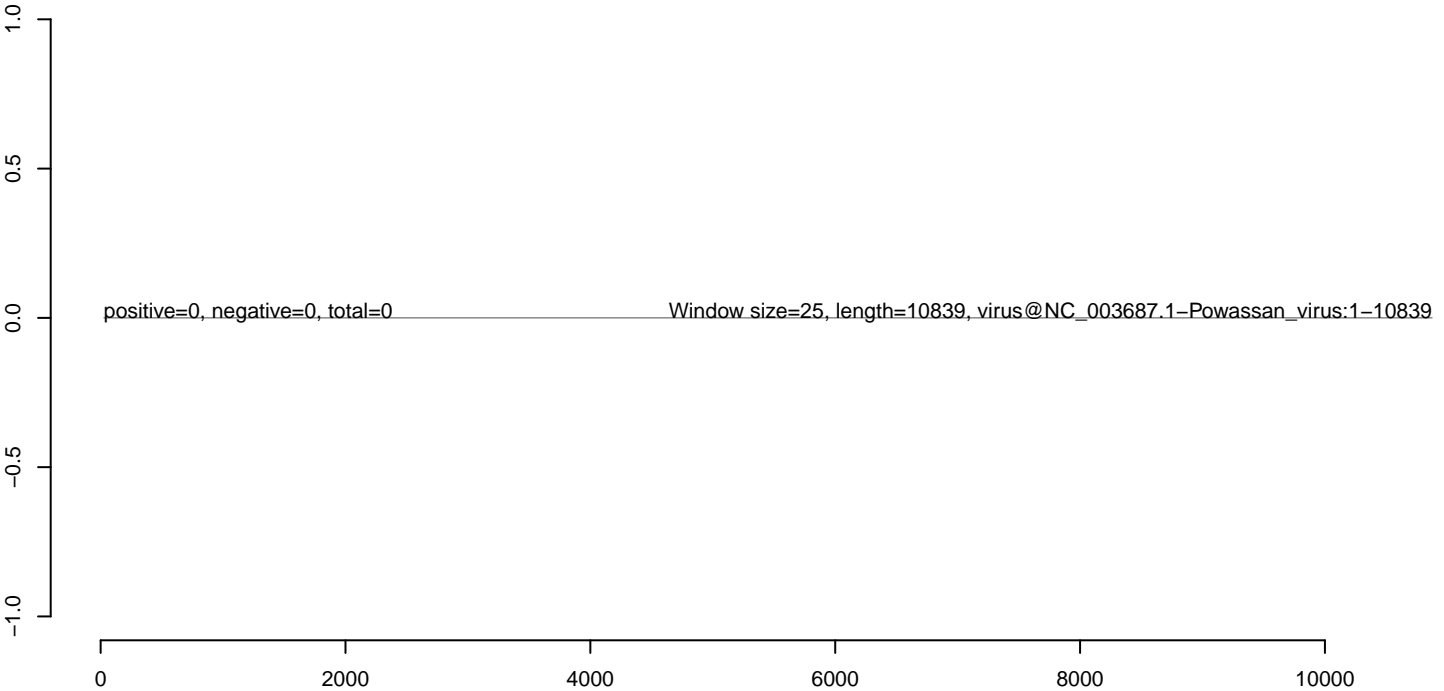
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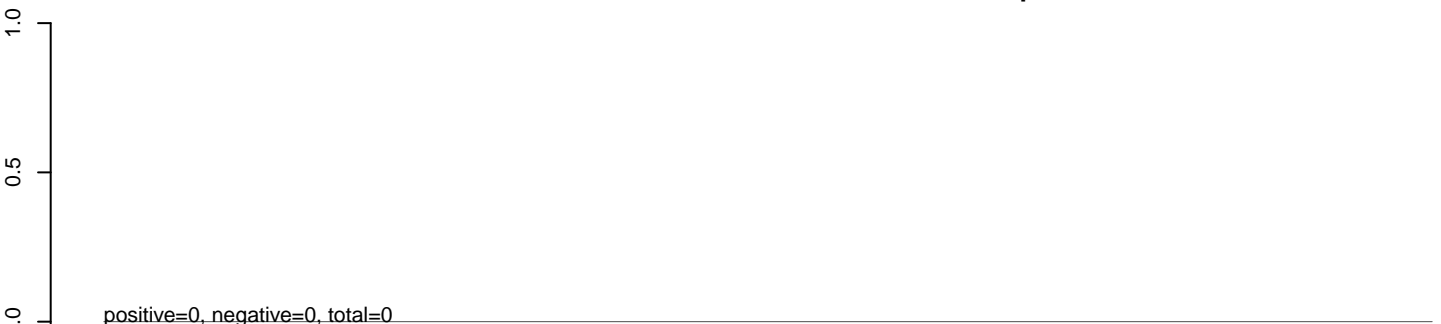
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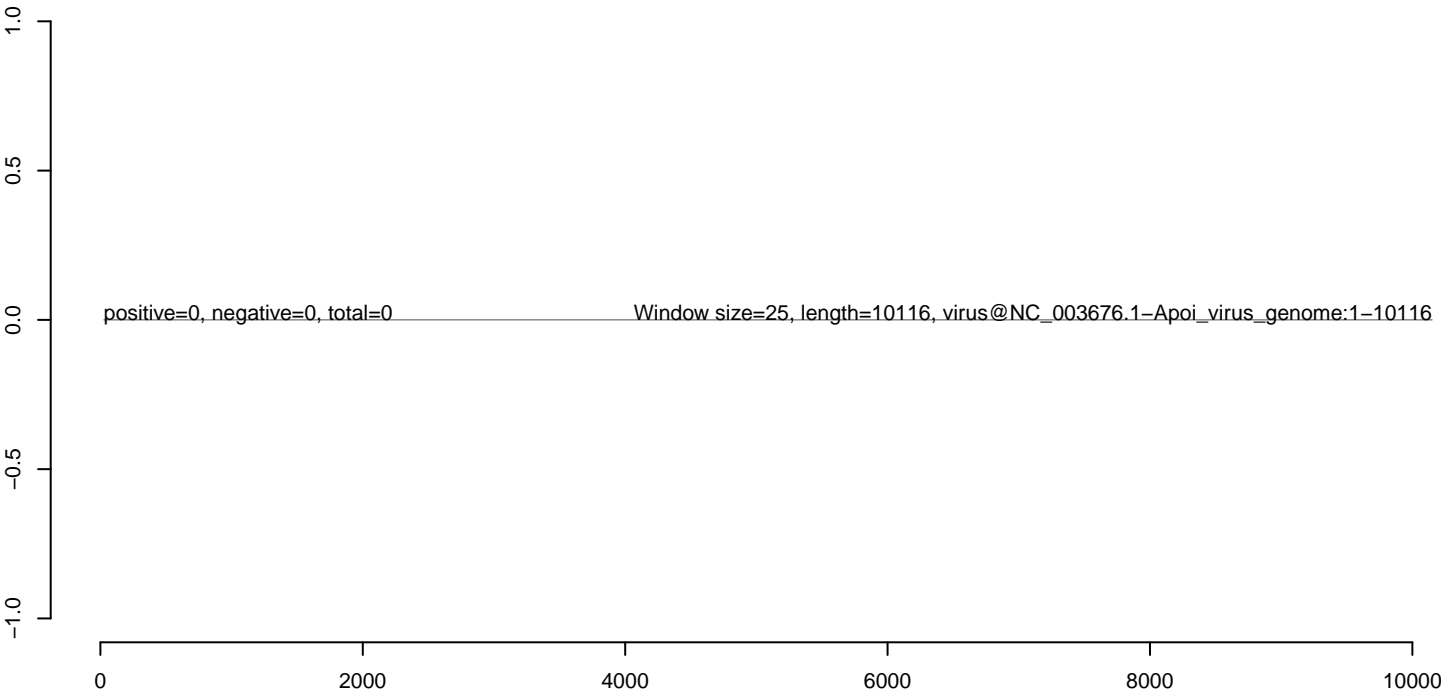
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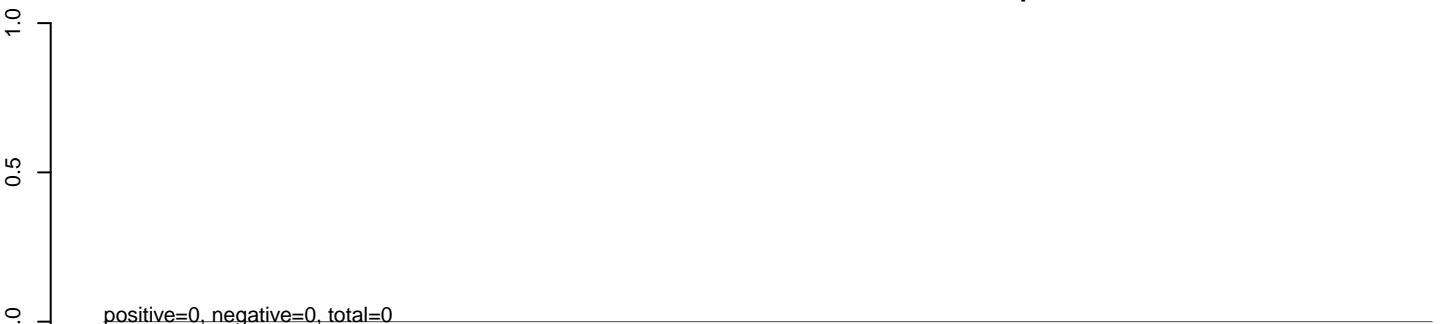
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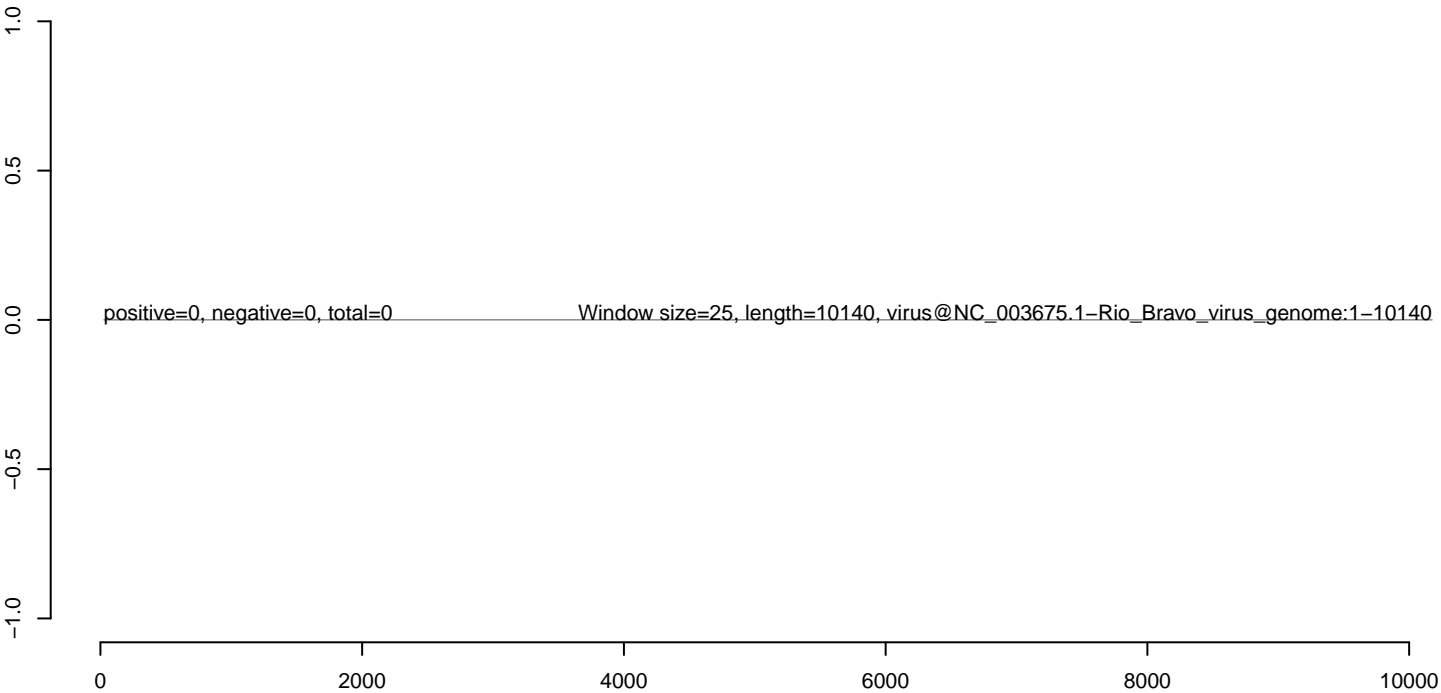
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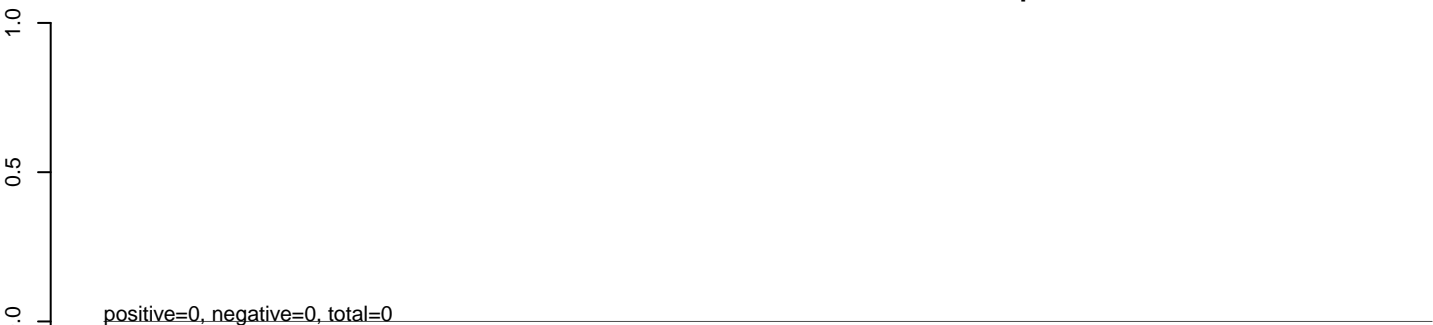
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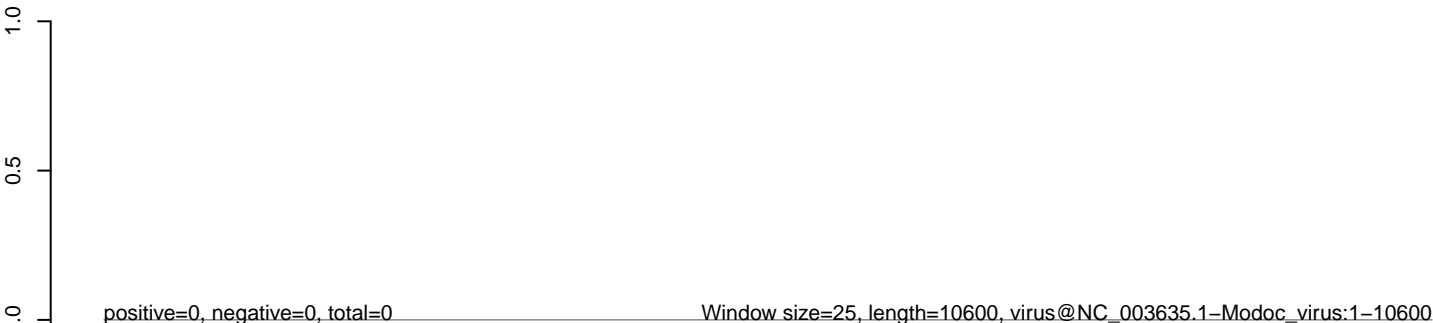
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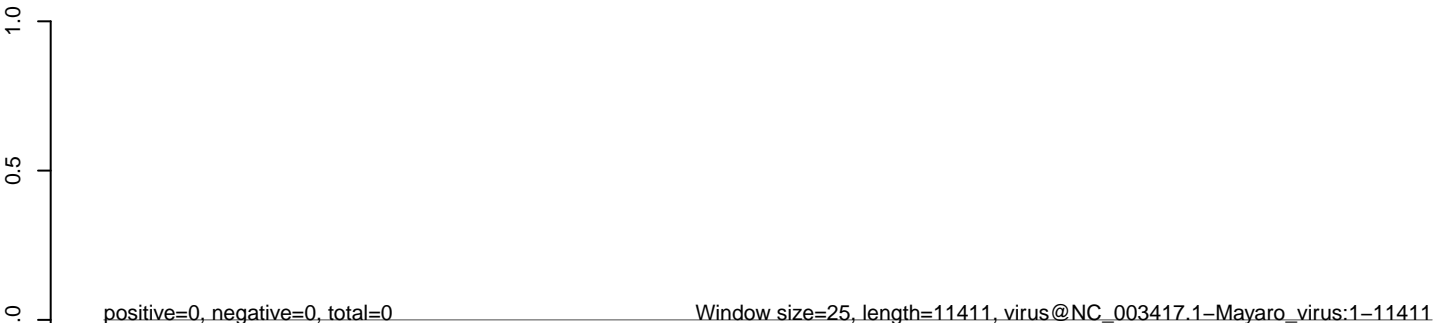
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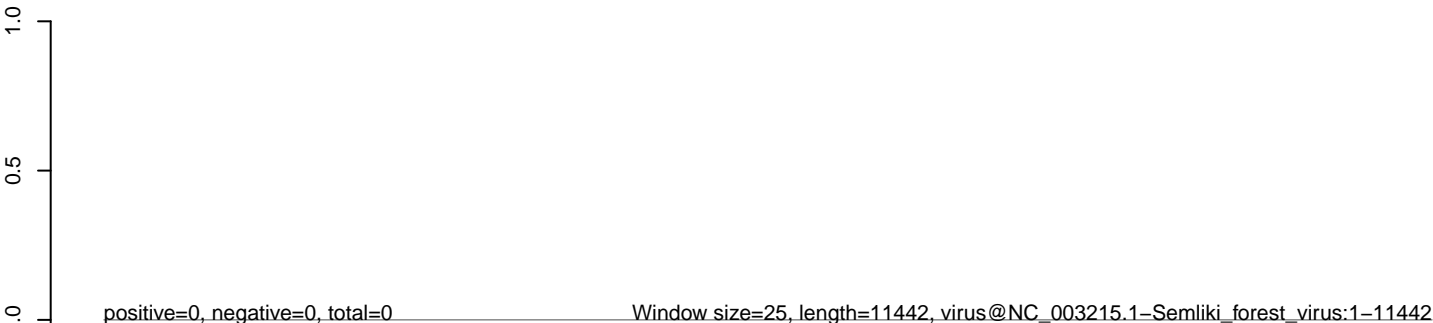
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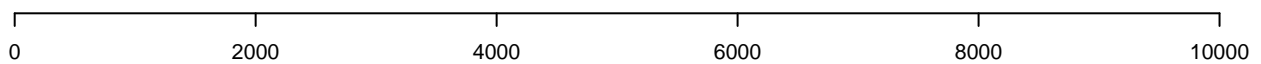
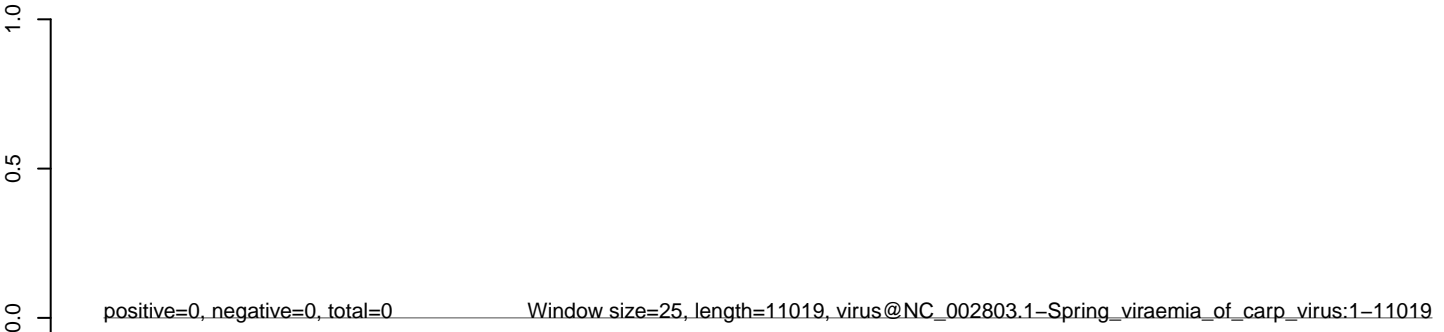
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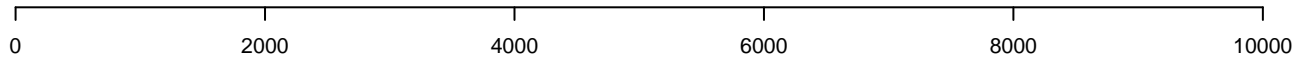
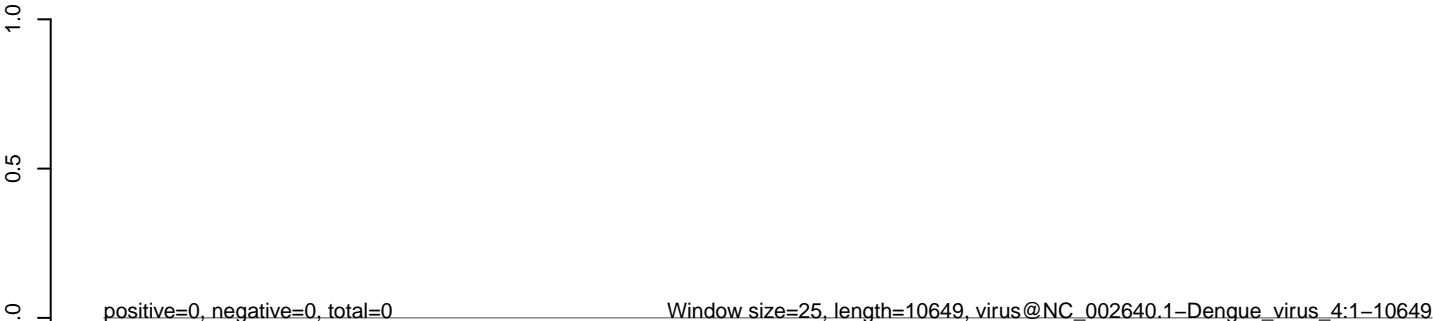
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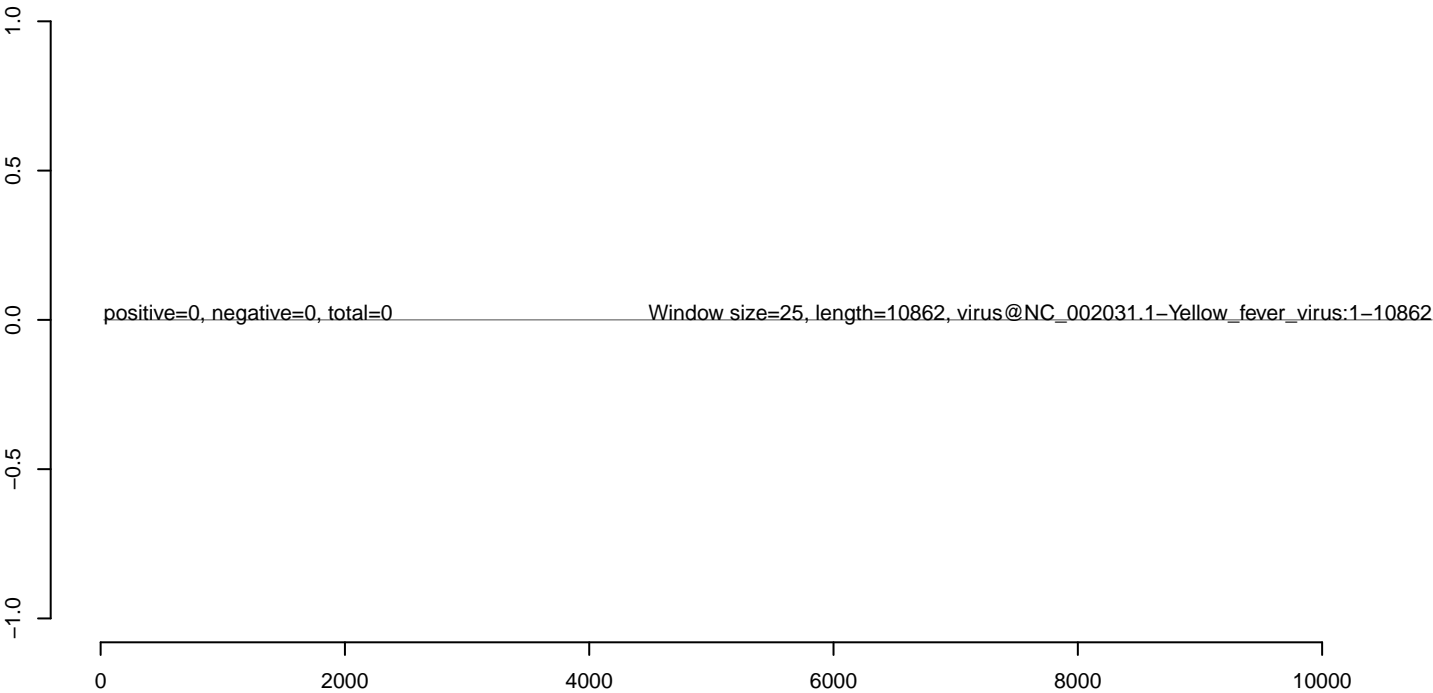
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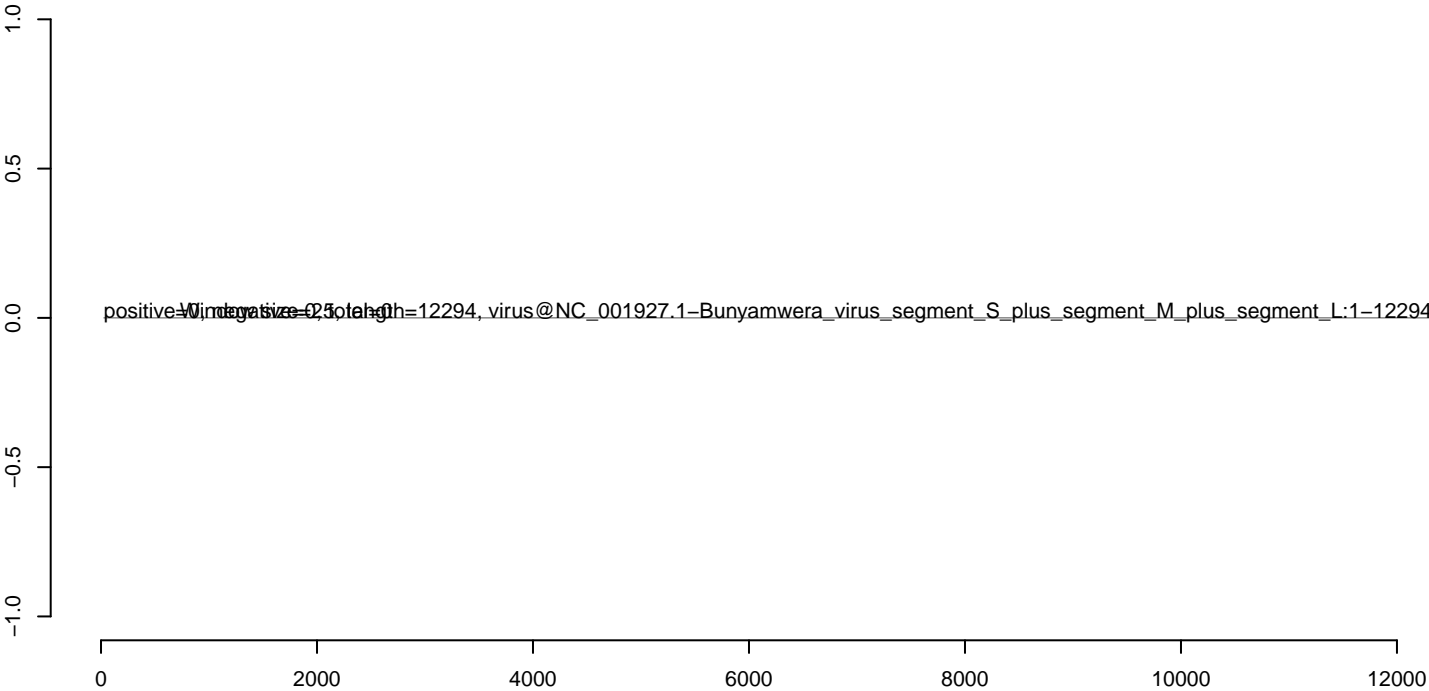
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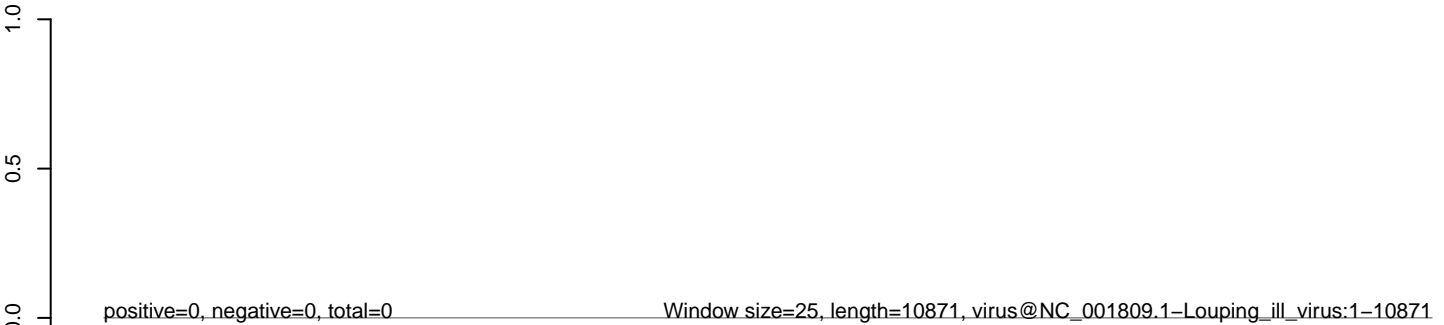
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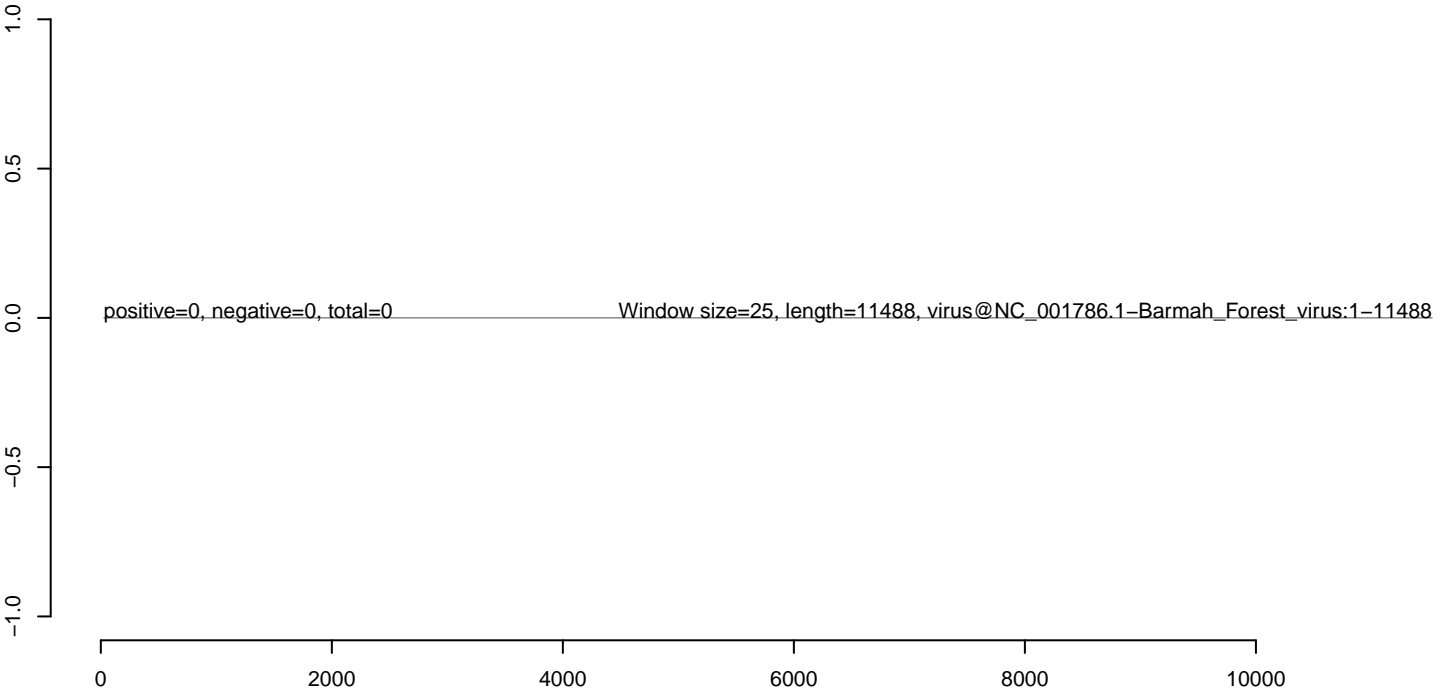
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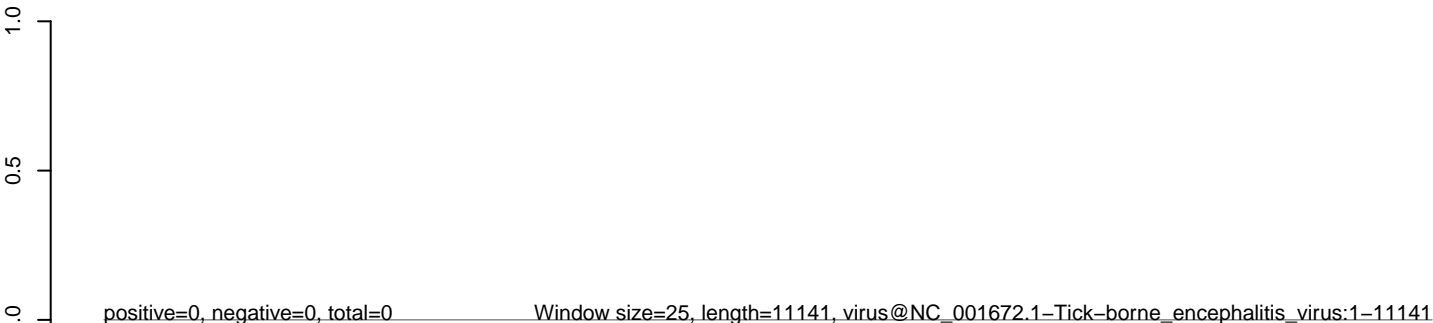
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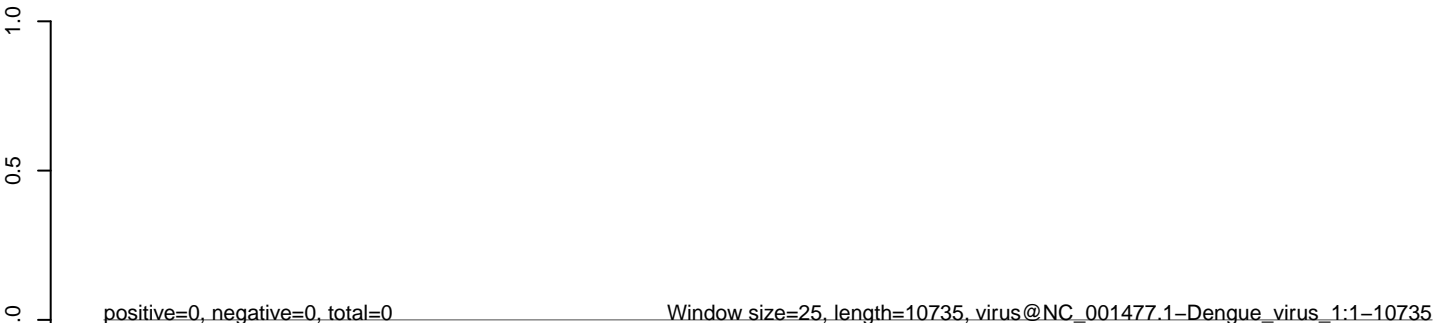
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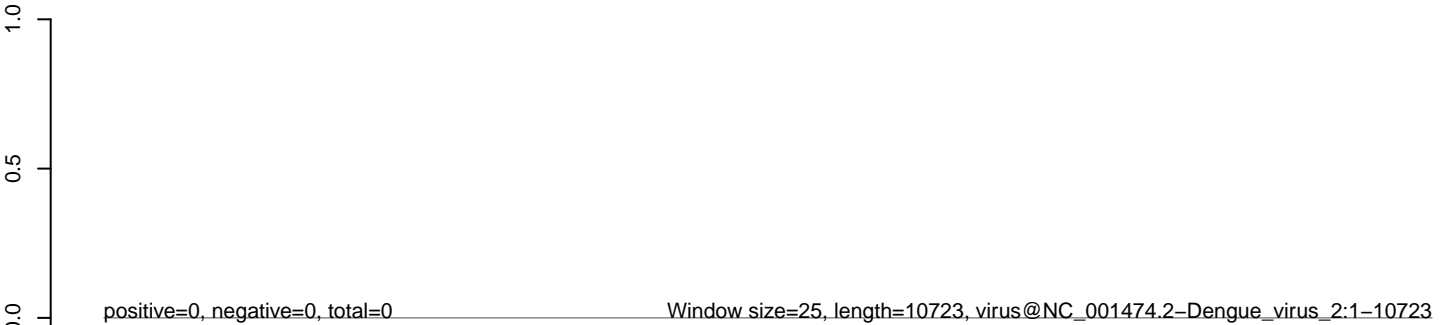
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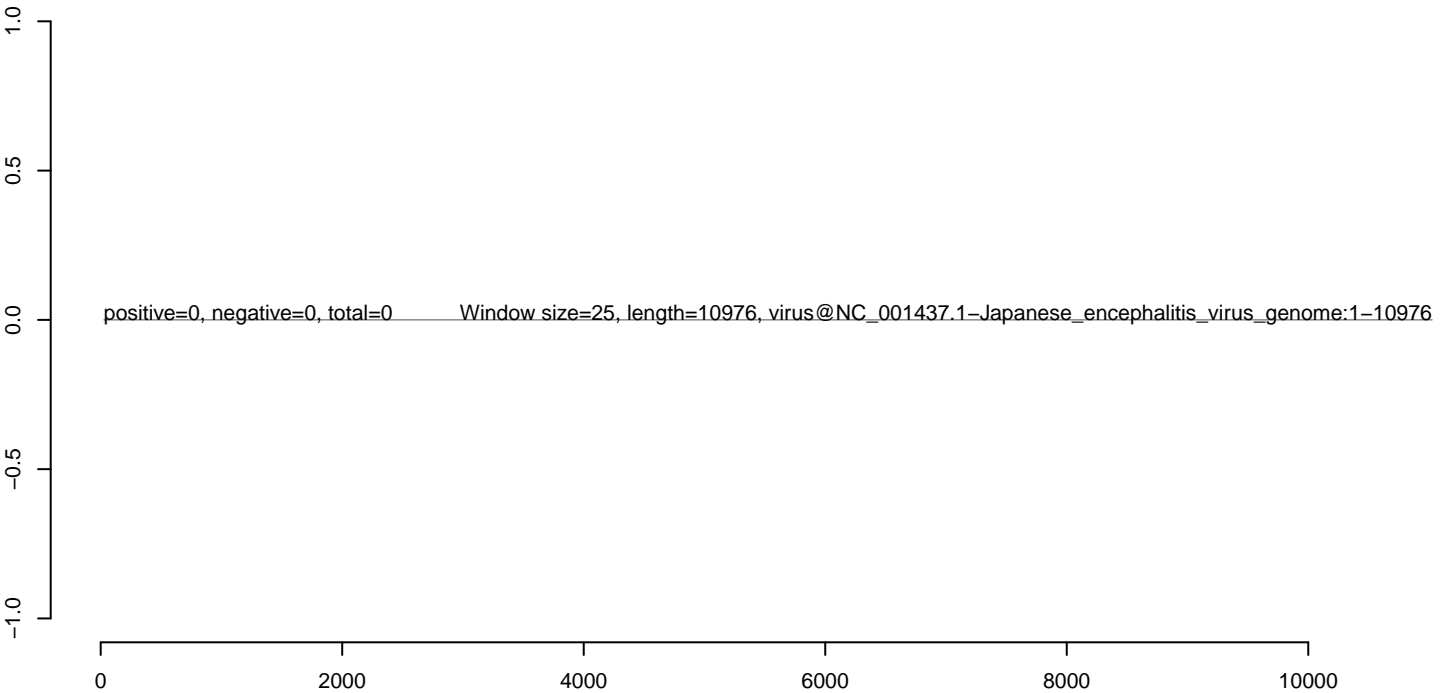
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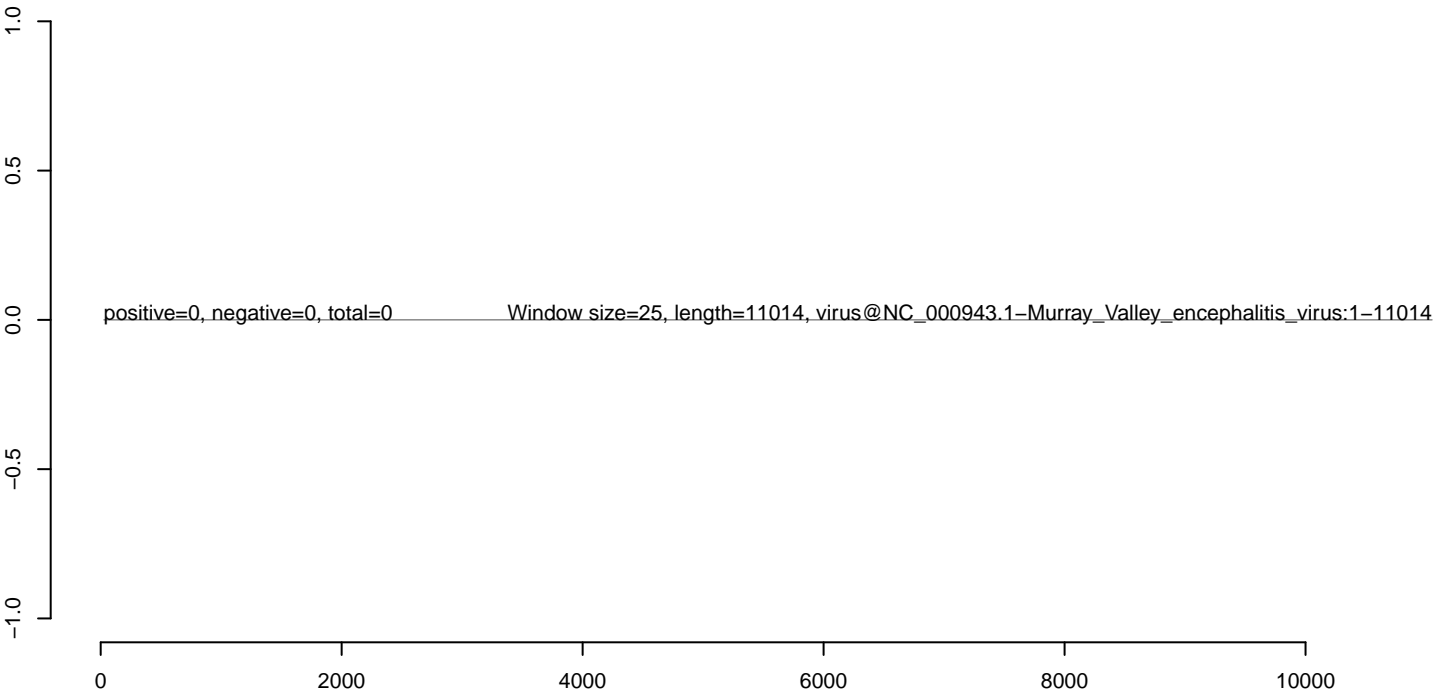
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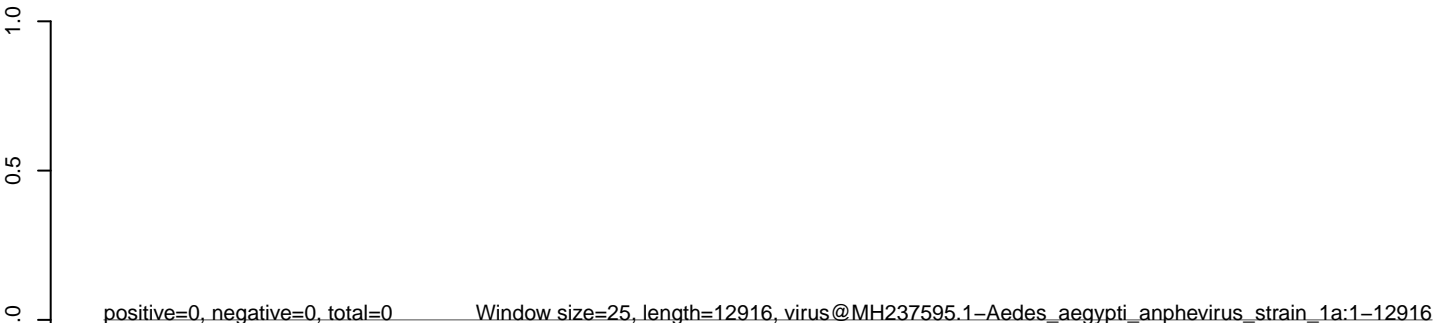
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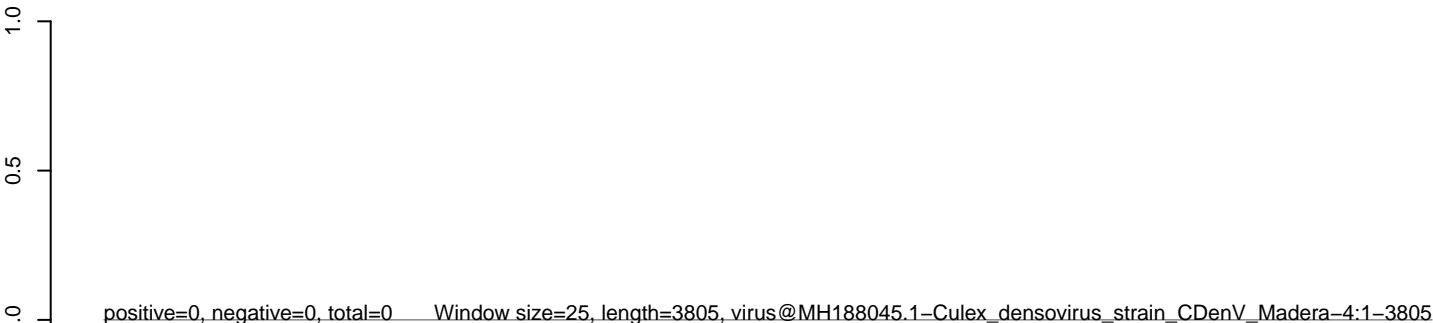
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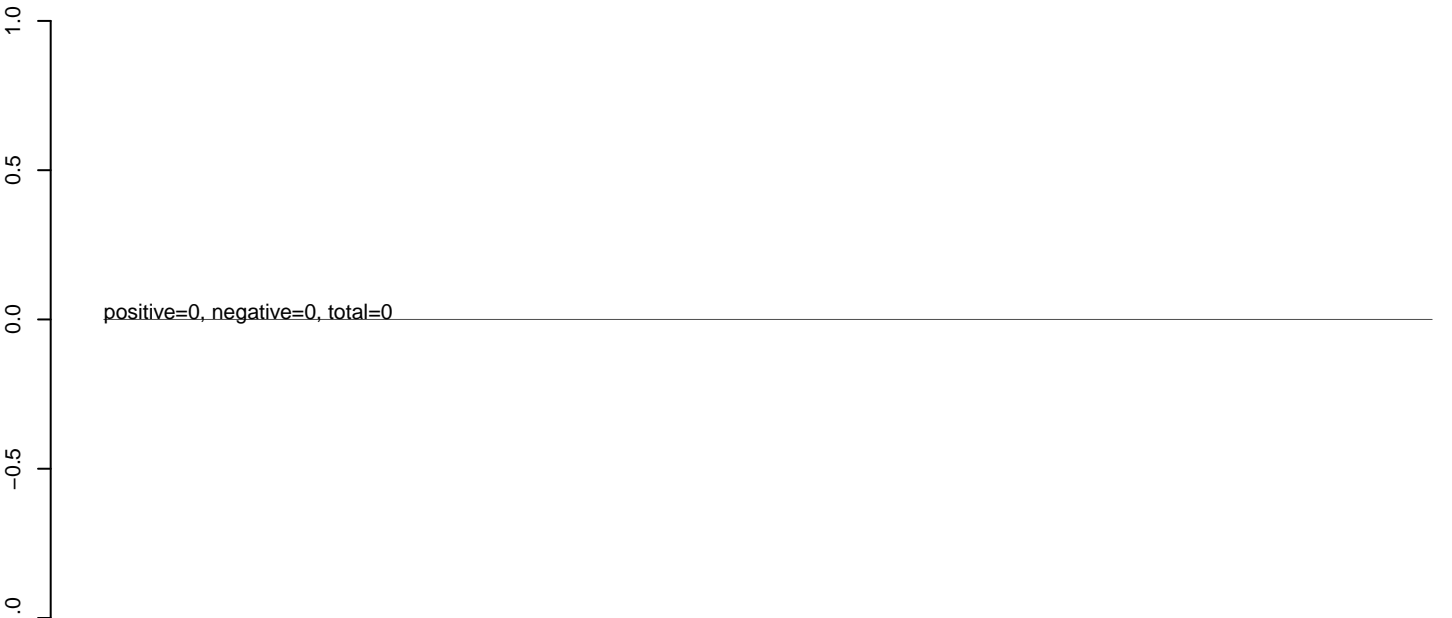
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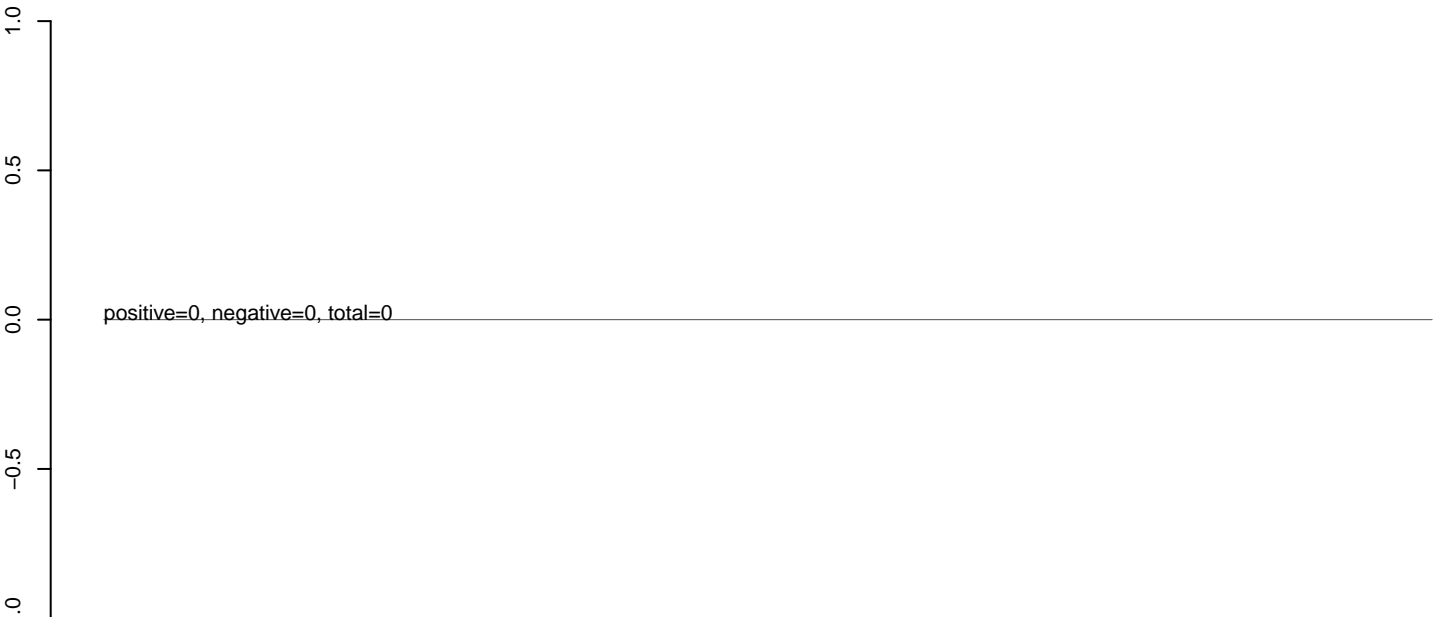
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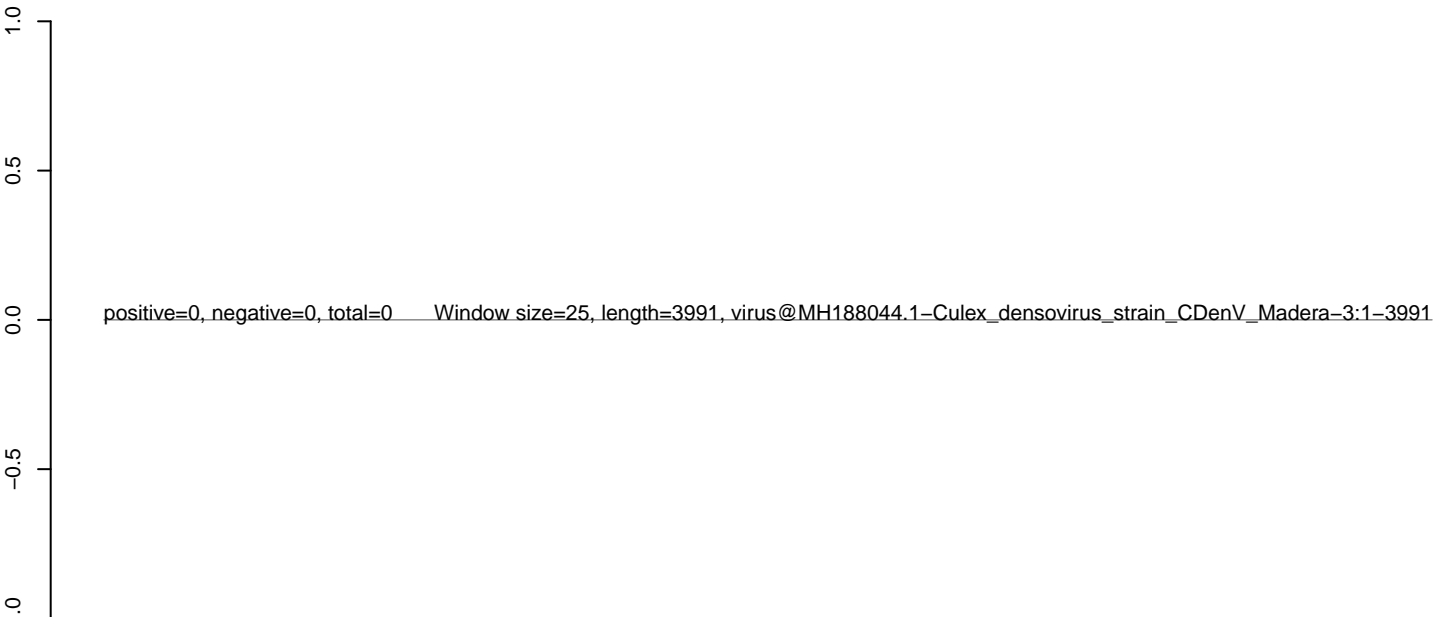
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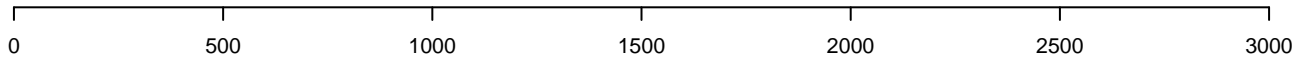
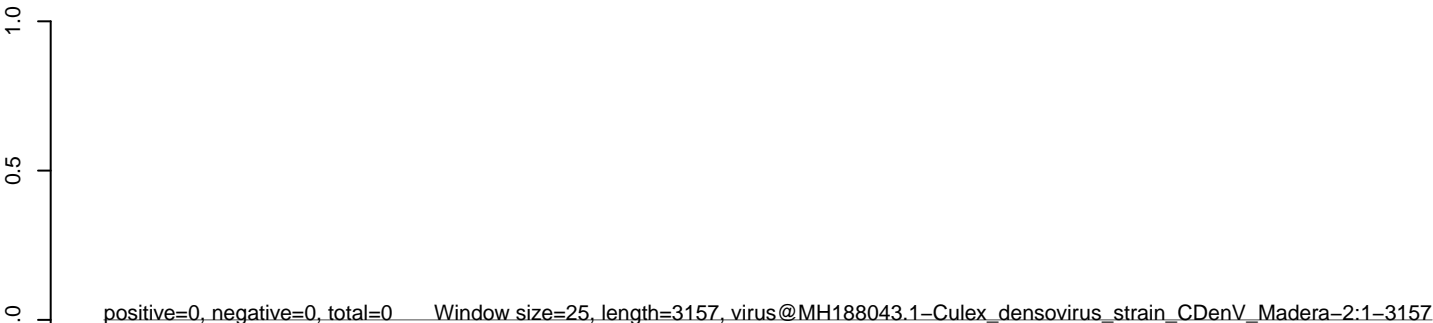
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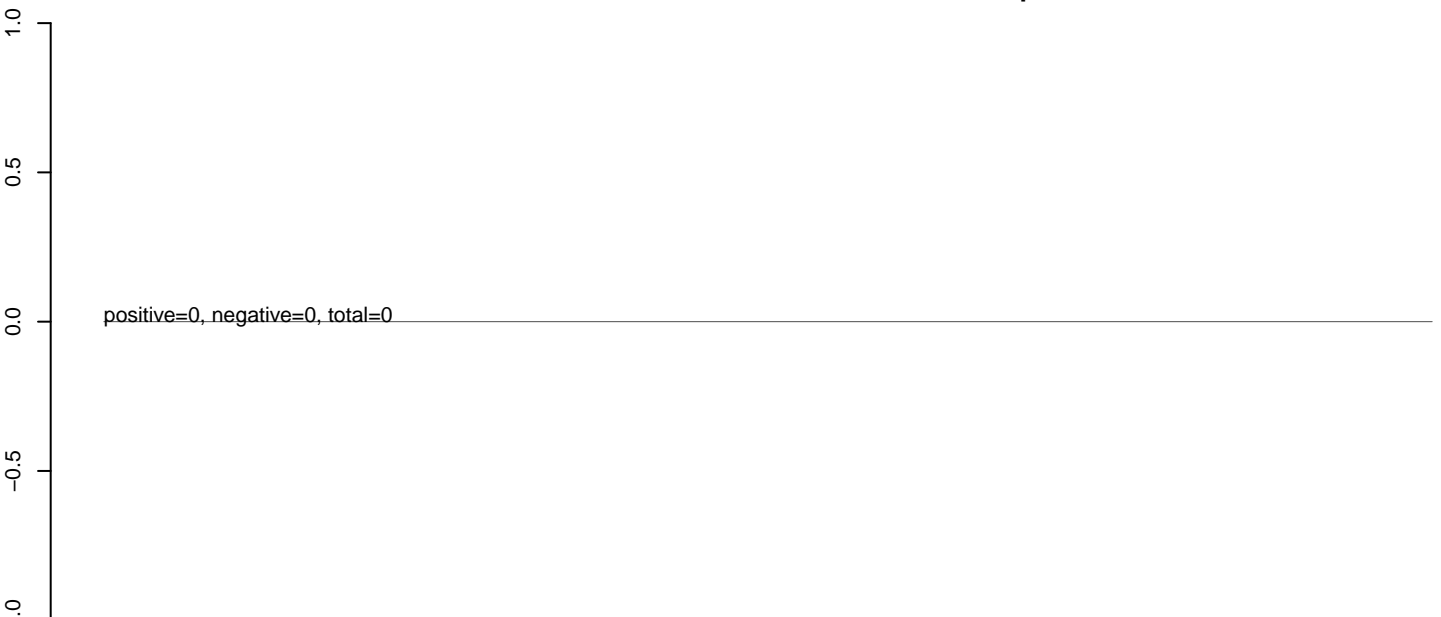
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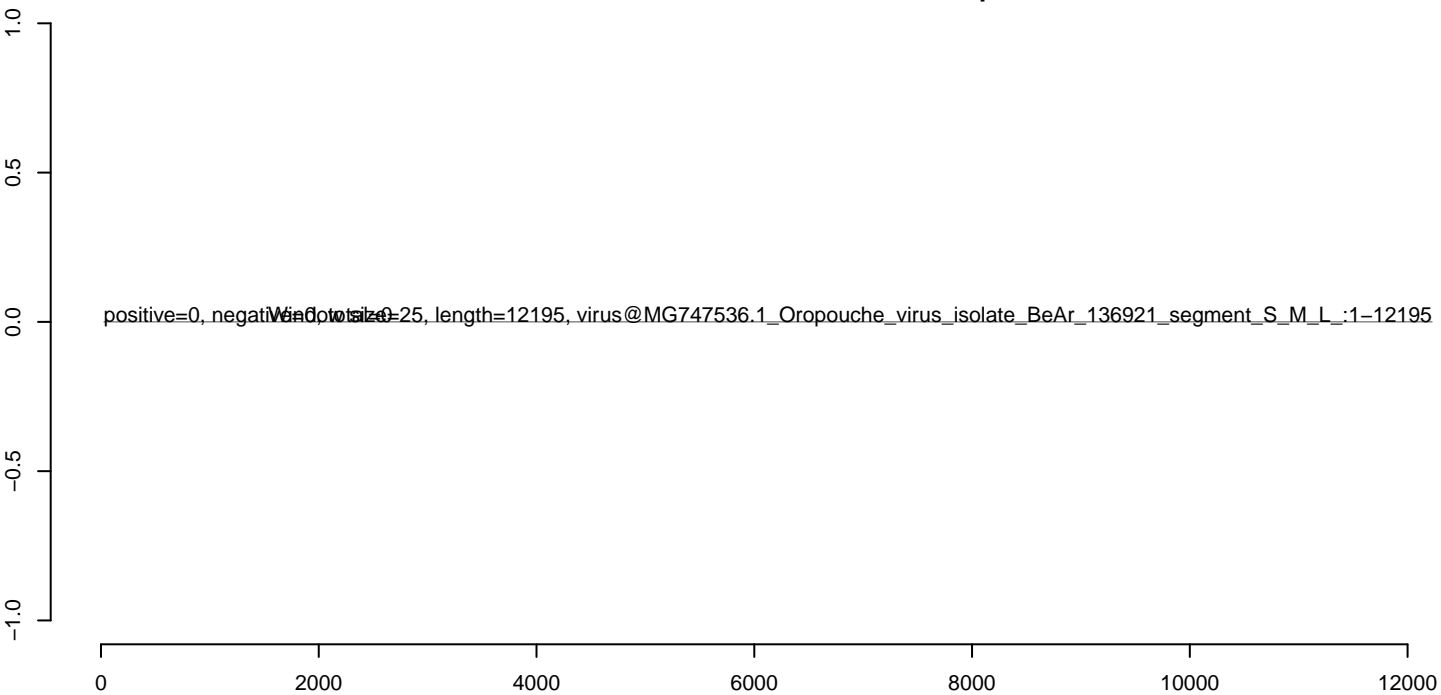
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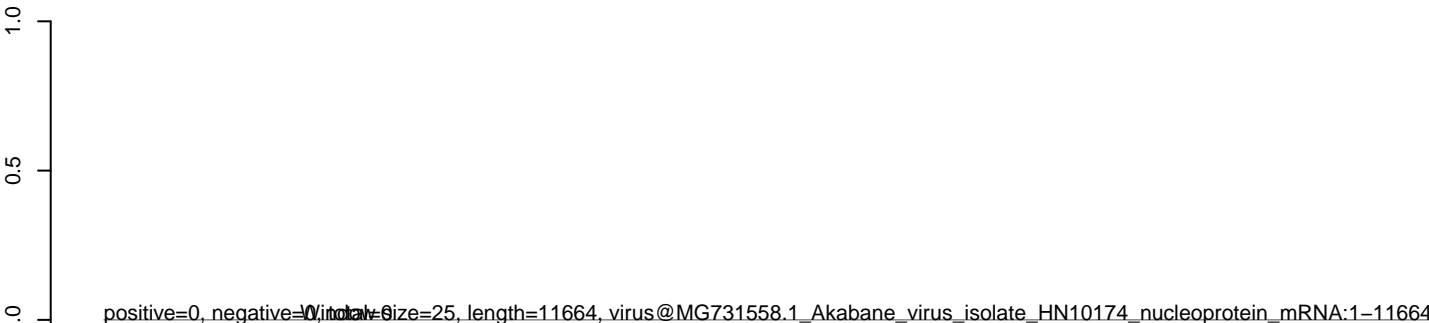
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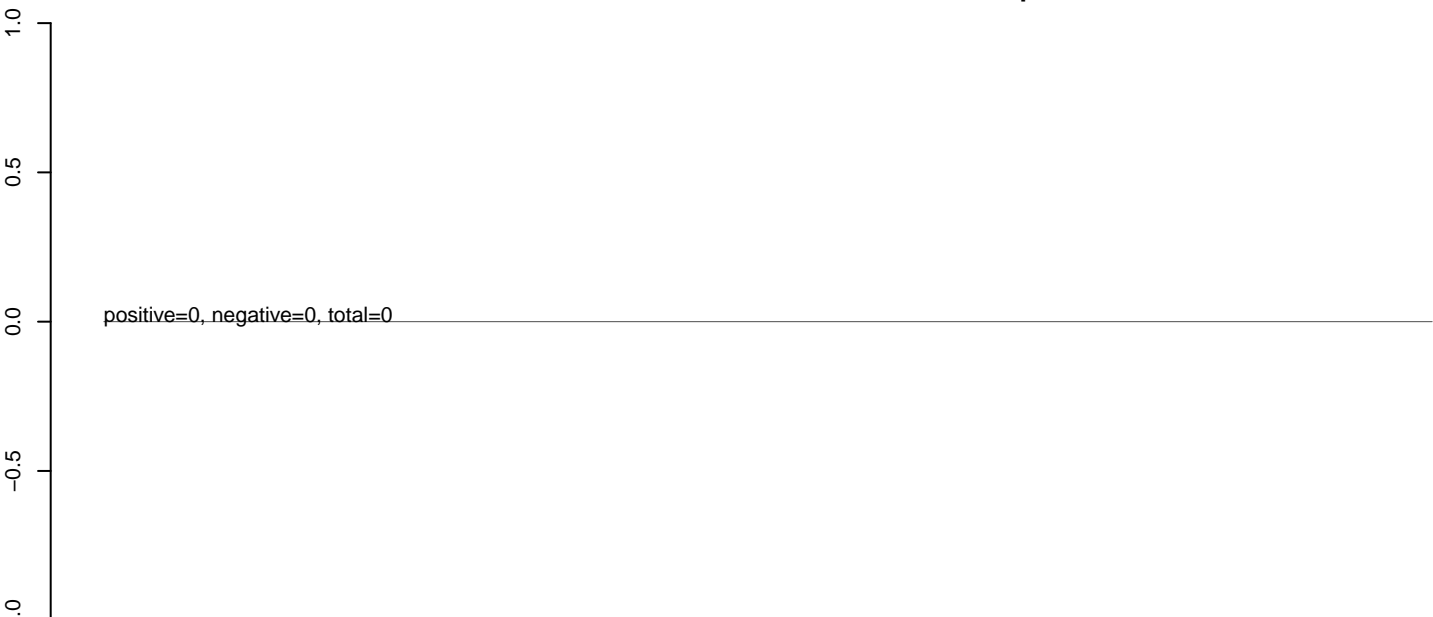


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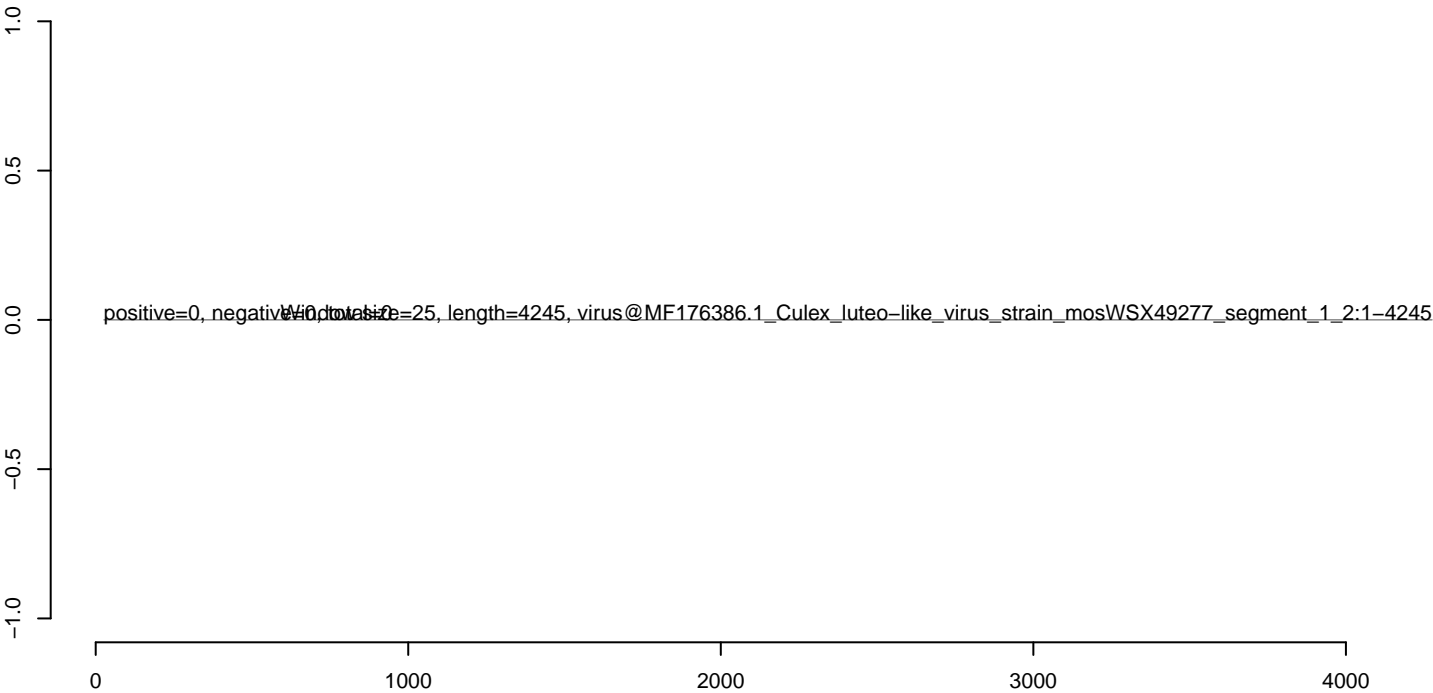
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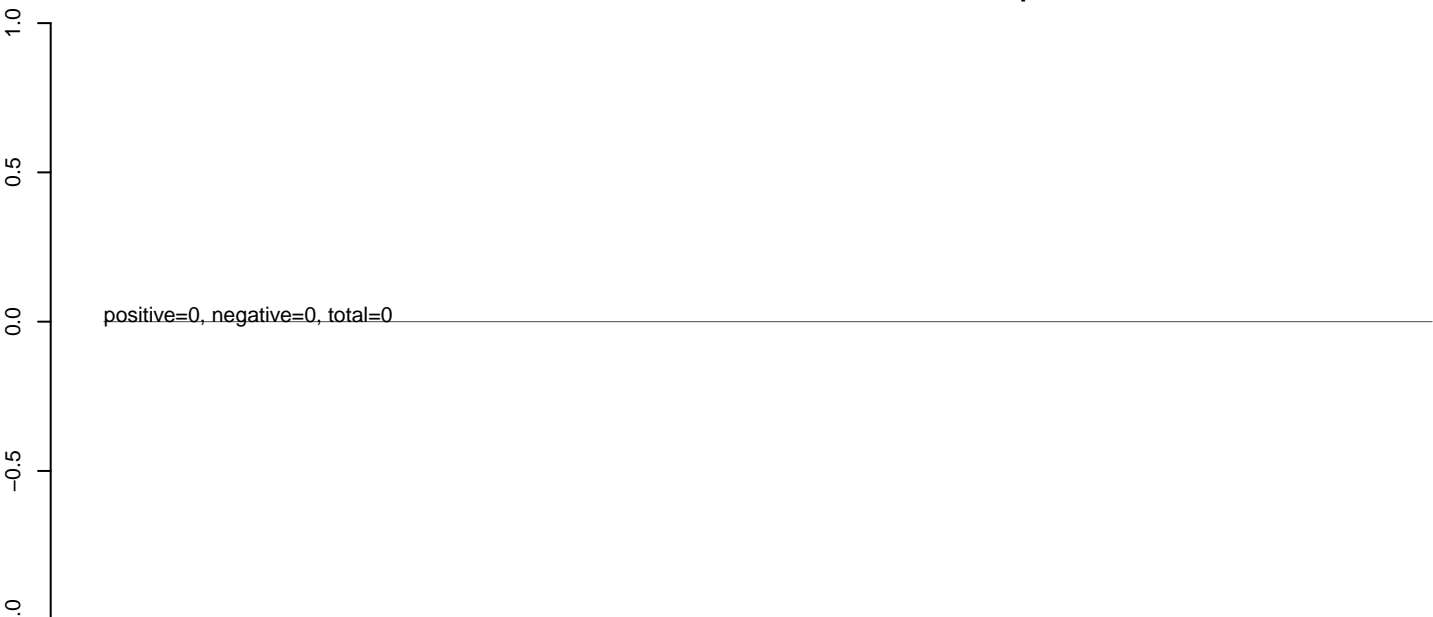
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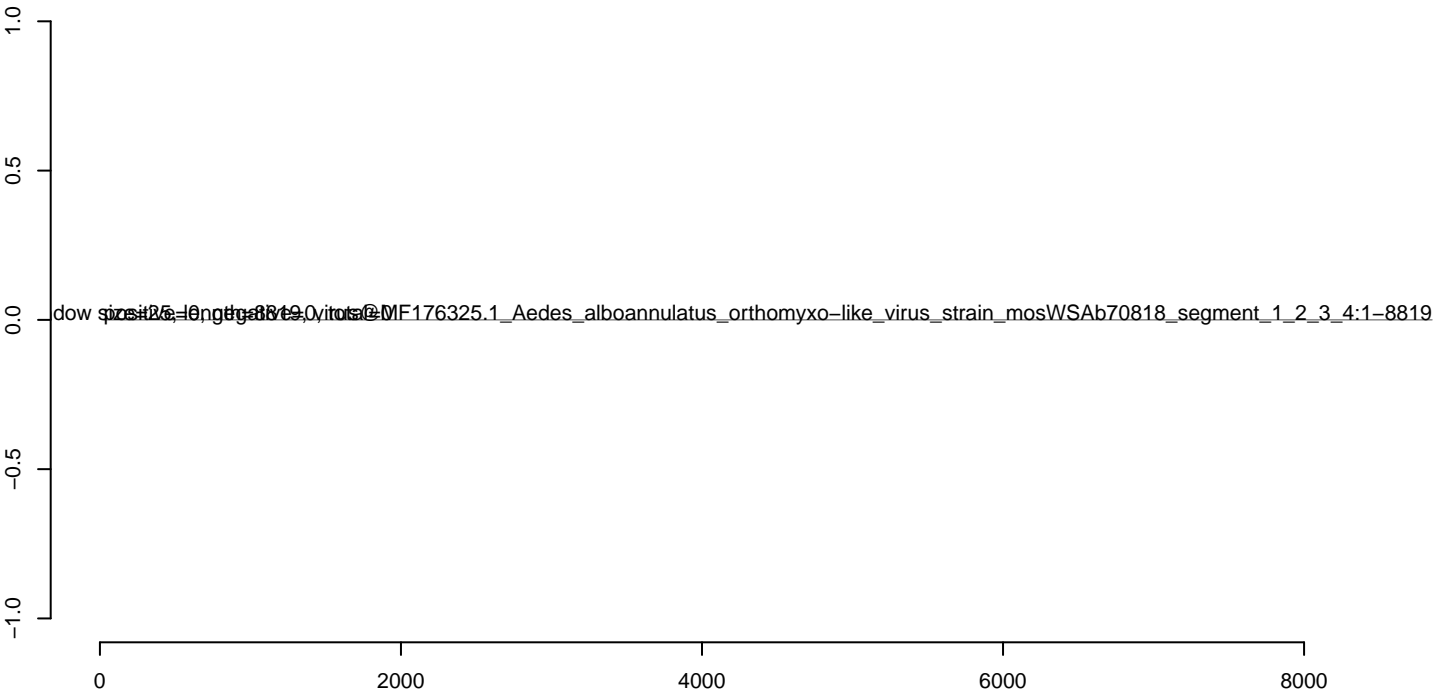
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AeAlbo_C636_SINV_GFP_betaE.24_35.rep



AeAlbo_C636_SINV_GFP_betaE.rep



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AeAlbo_C636_SINV_GFP_betaE.rep



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AeAlbo_C636_SINV_GFP_betaE.18_23.rep



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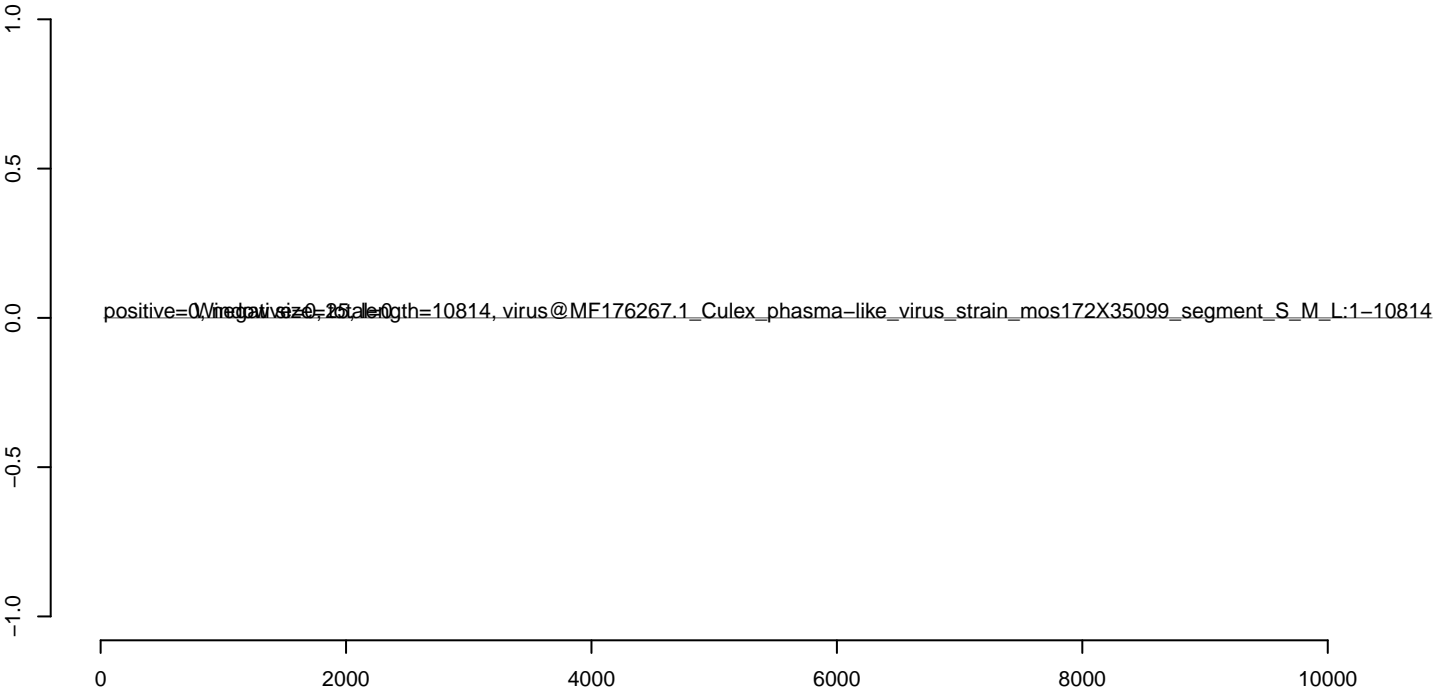
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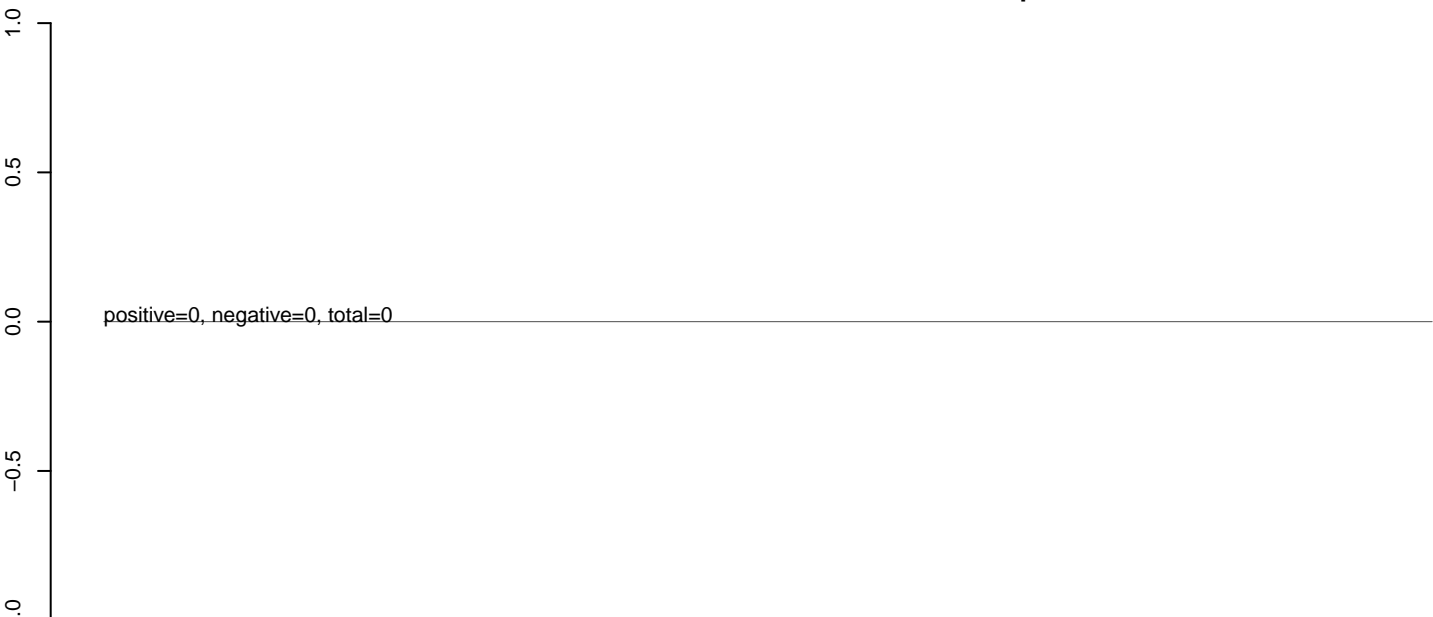
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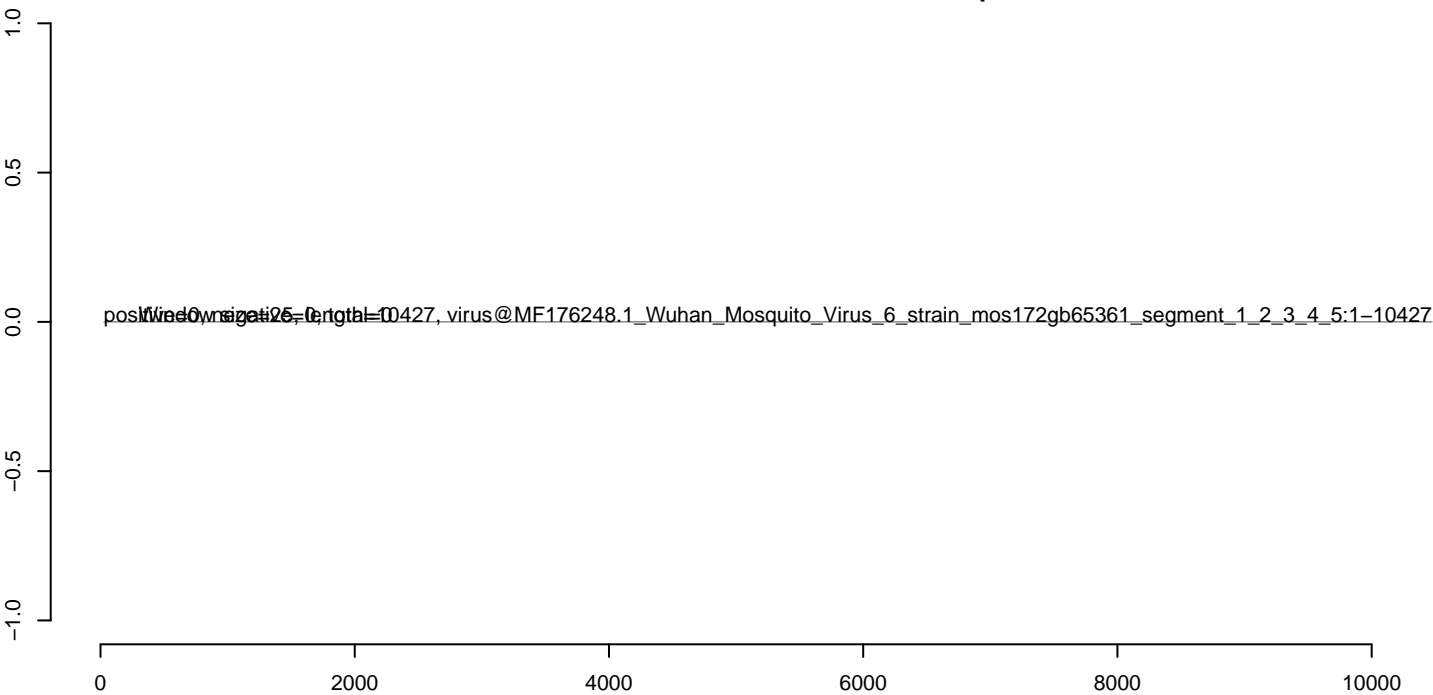
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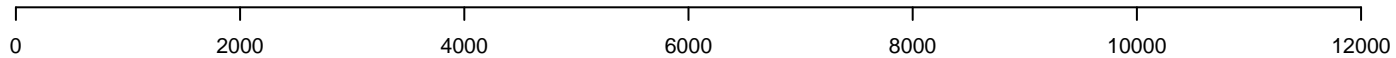
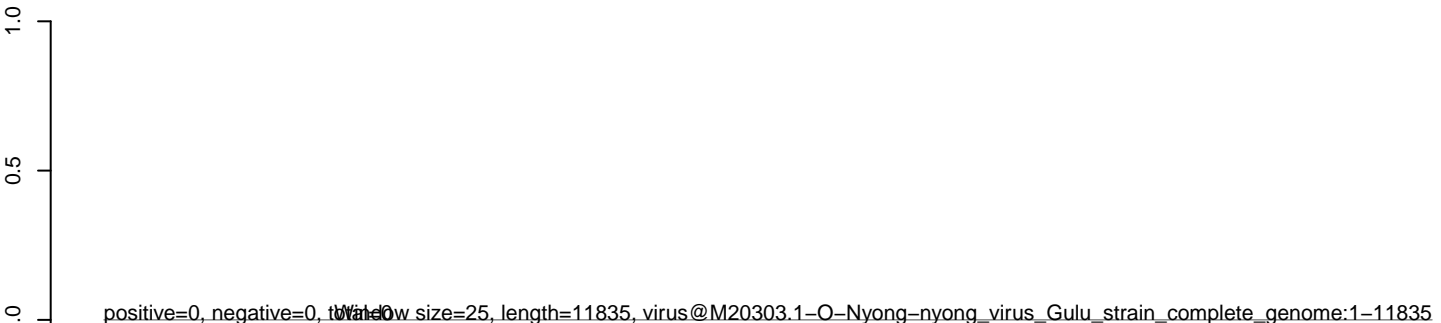
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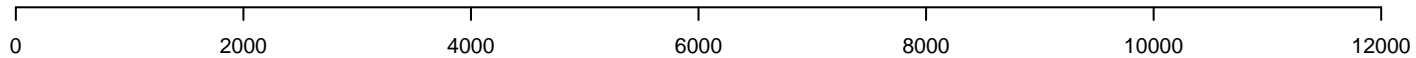
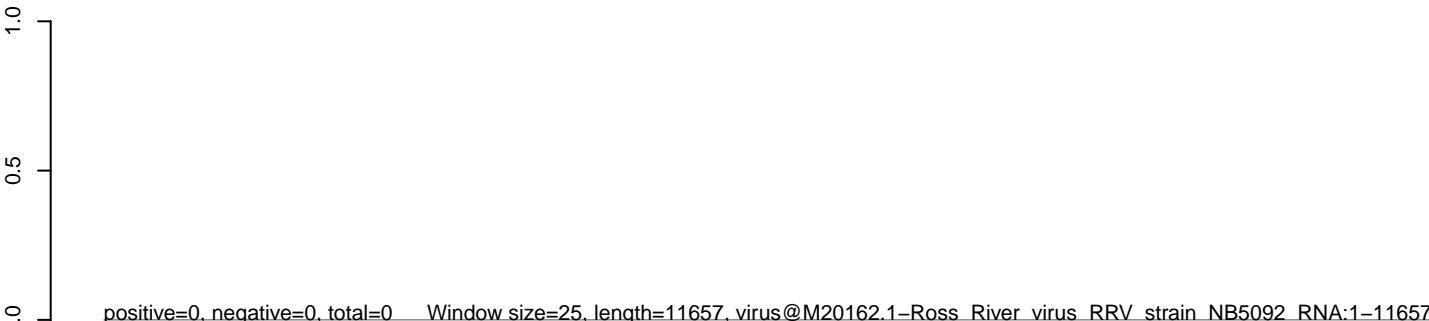
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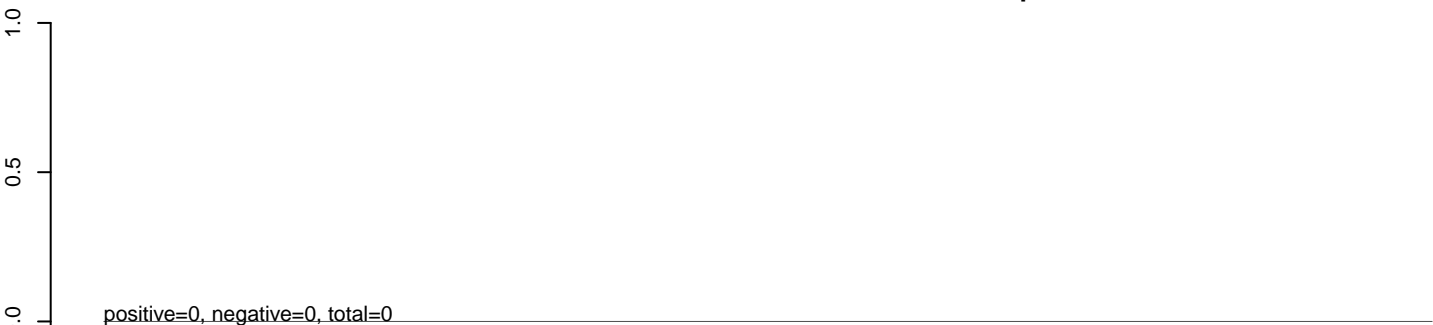
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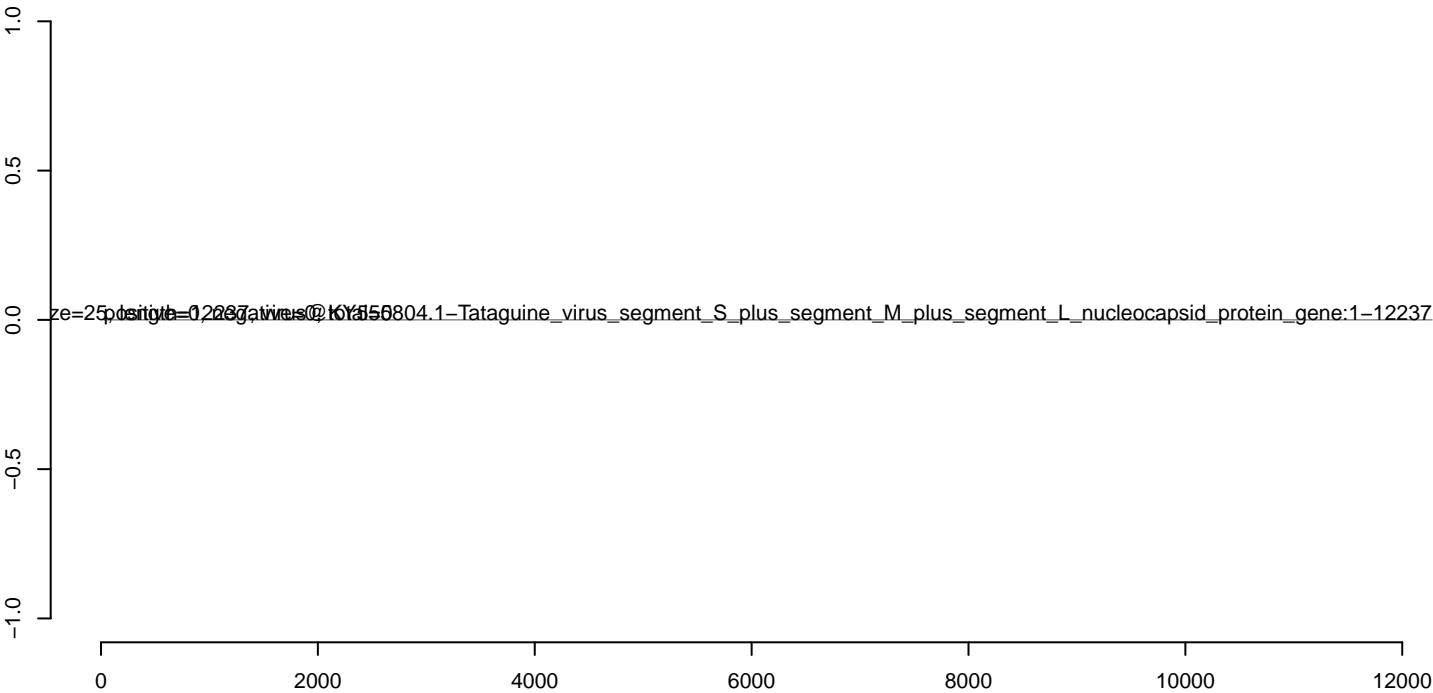
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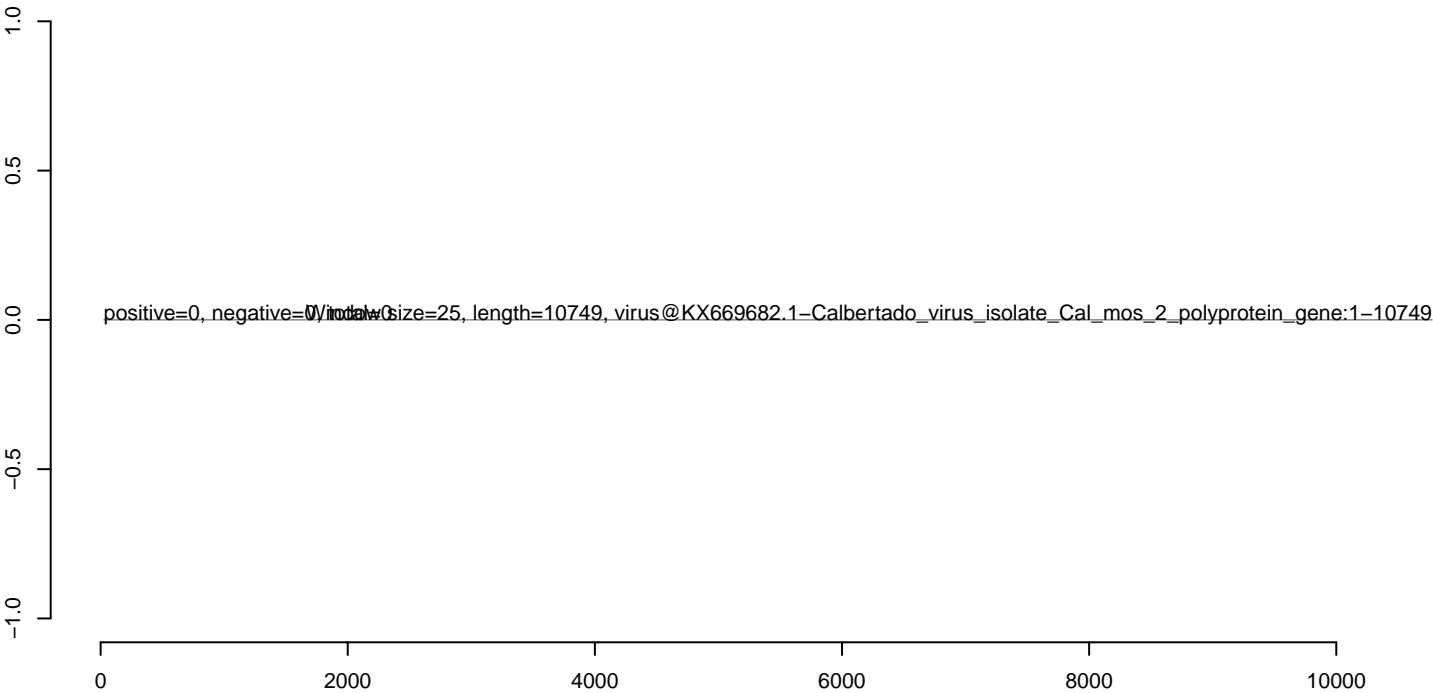
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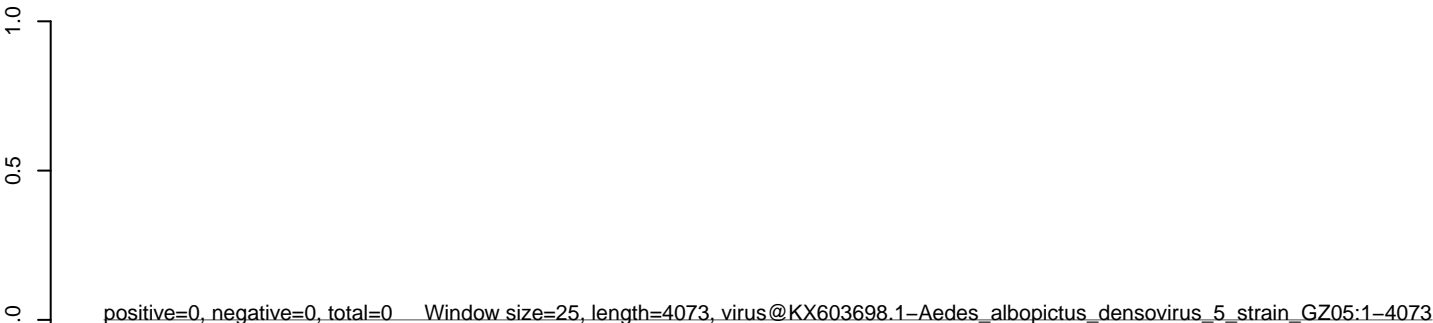
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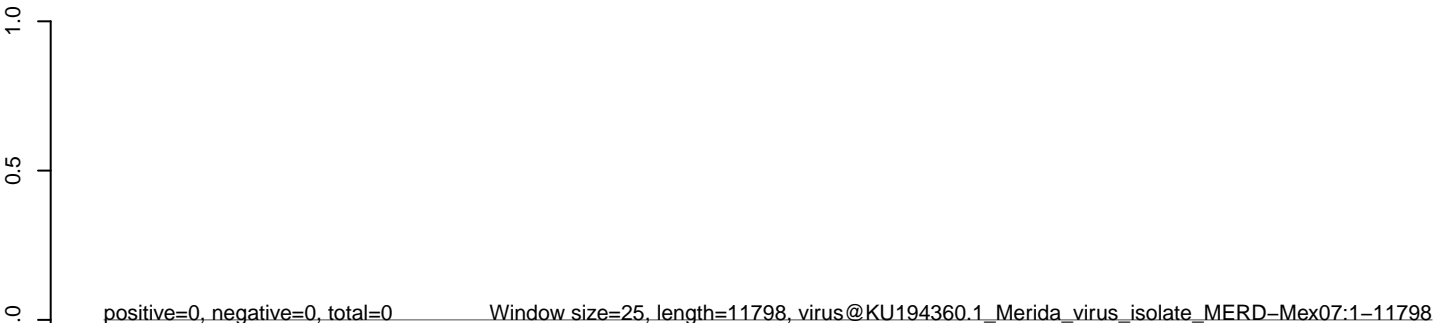
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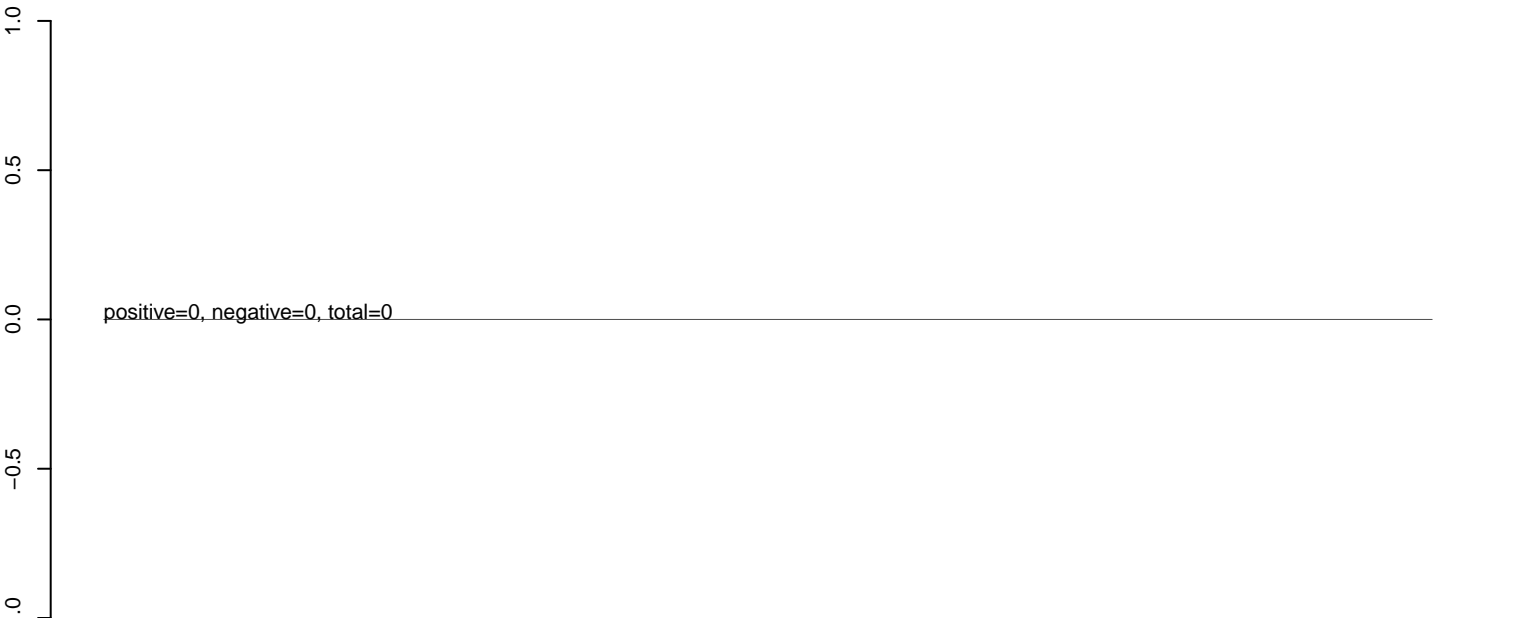


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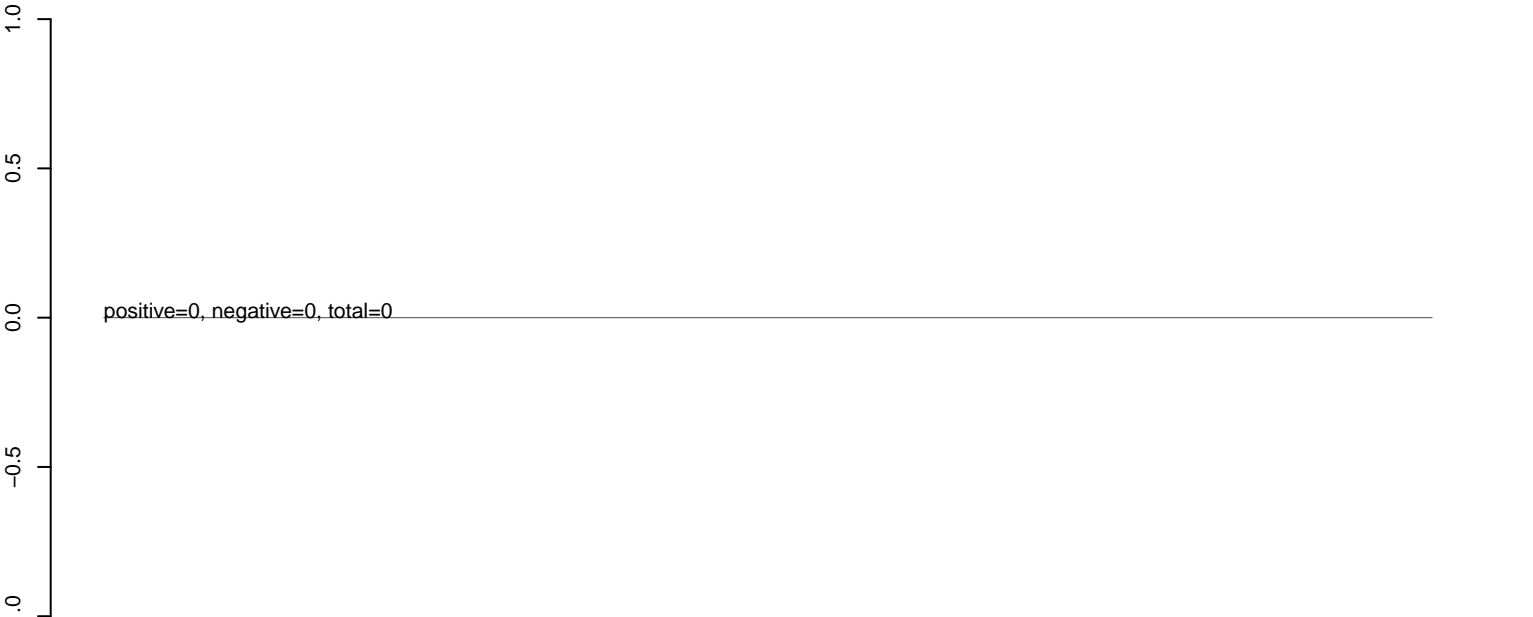


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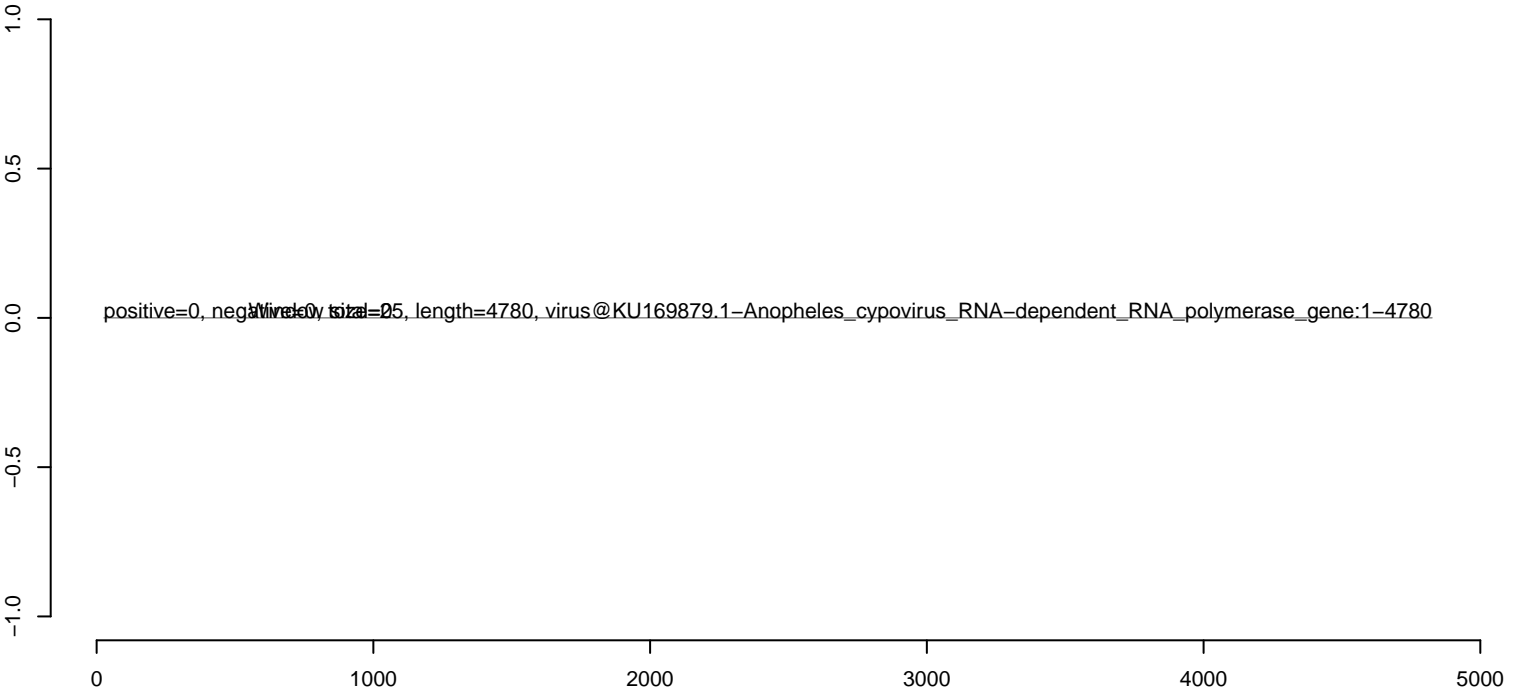
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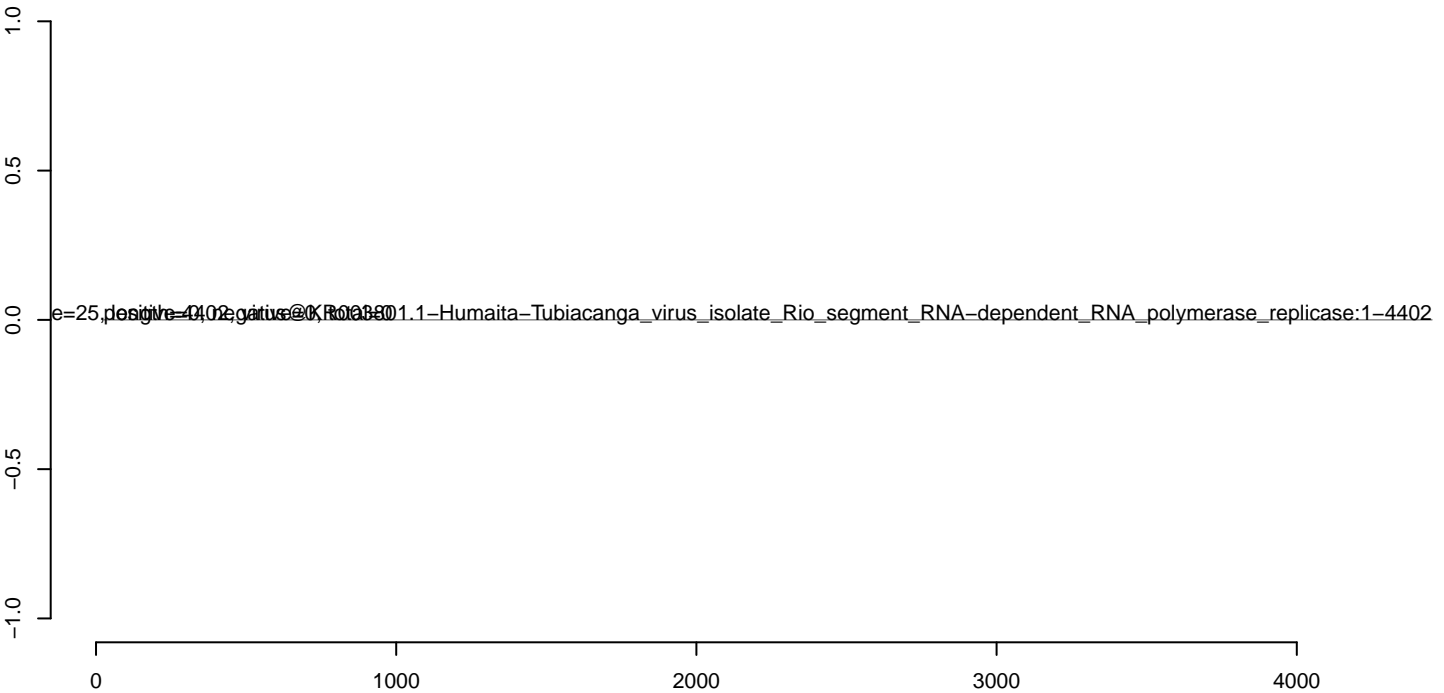
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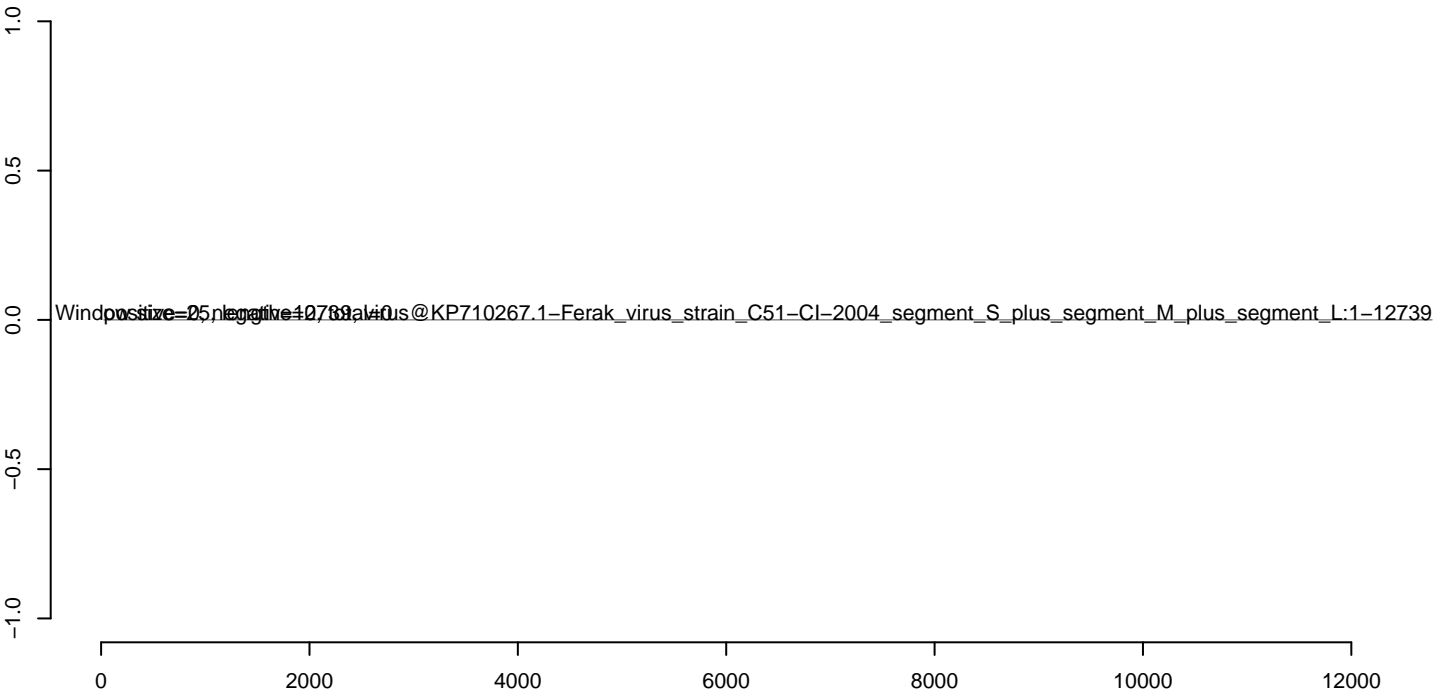
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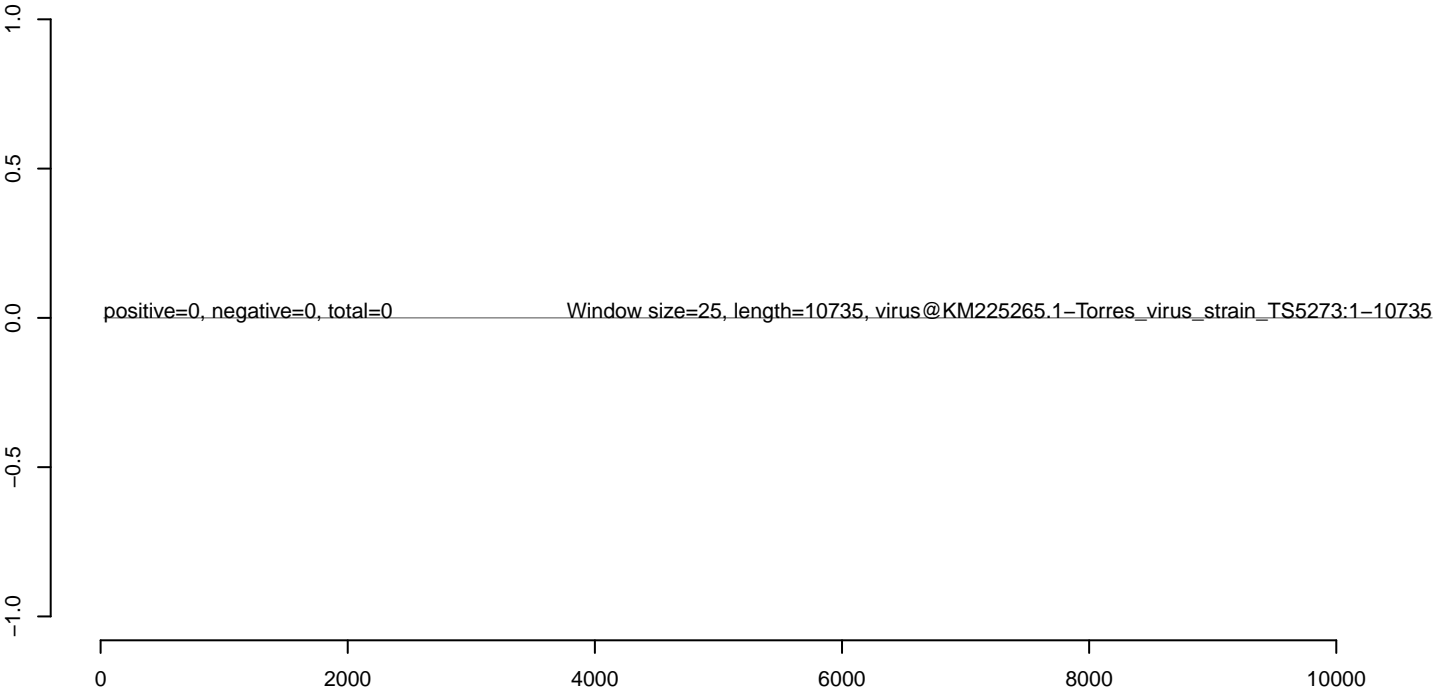
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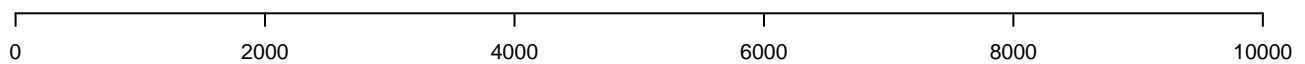
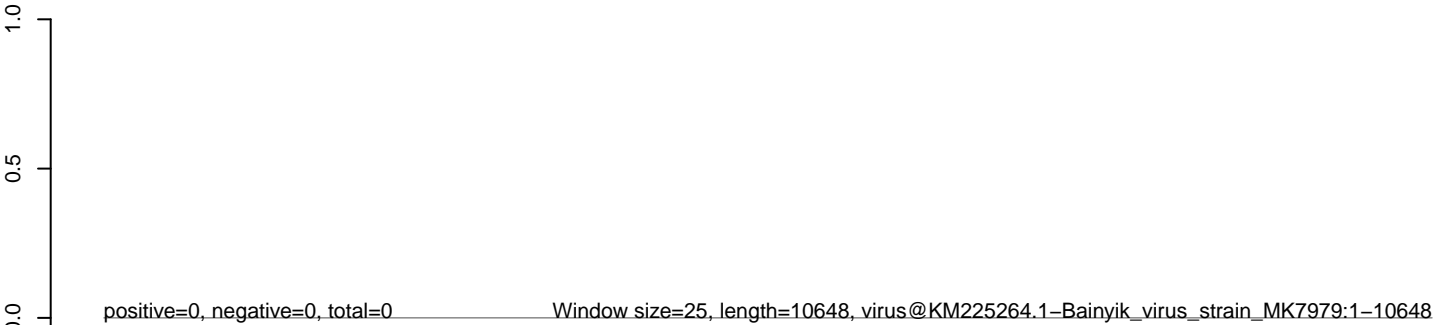
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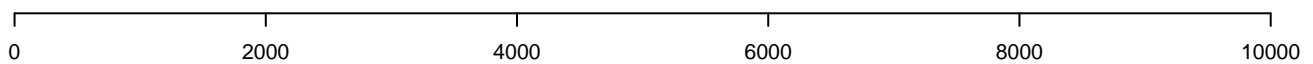
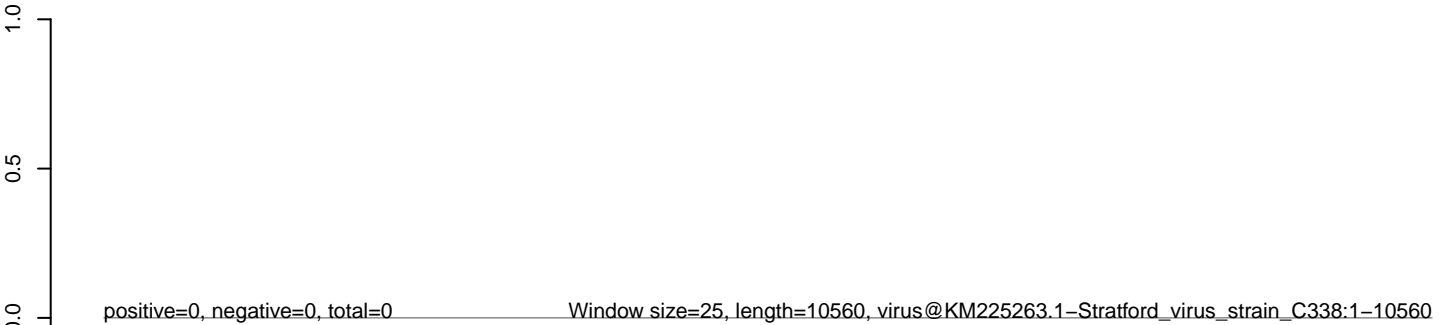
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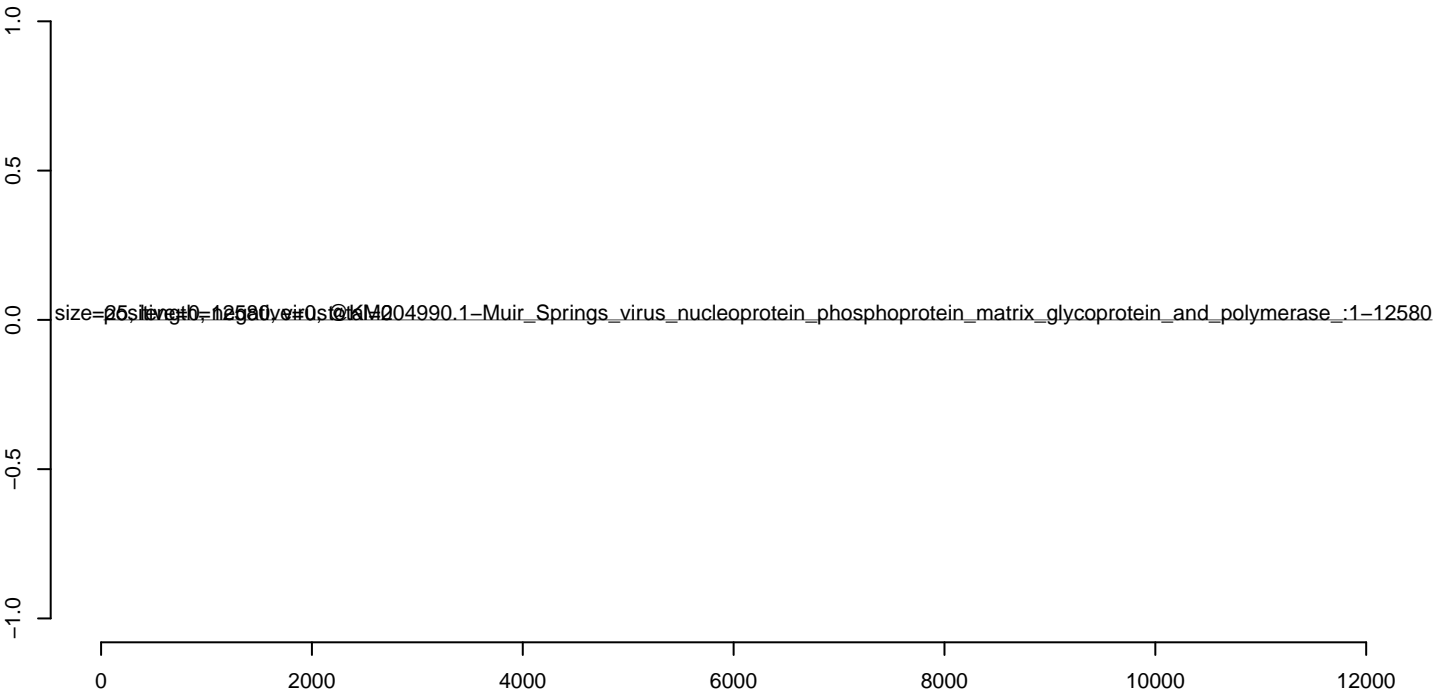
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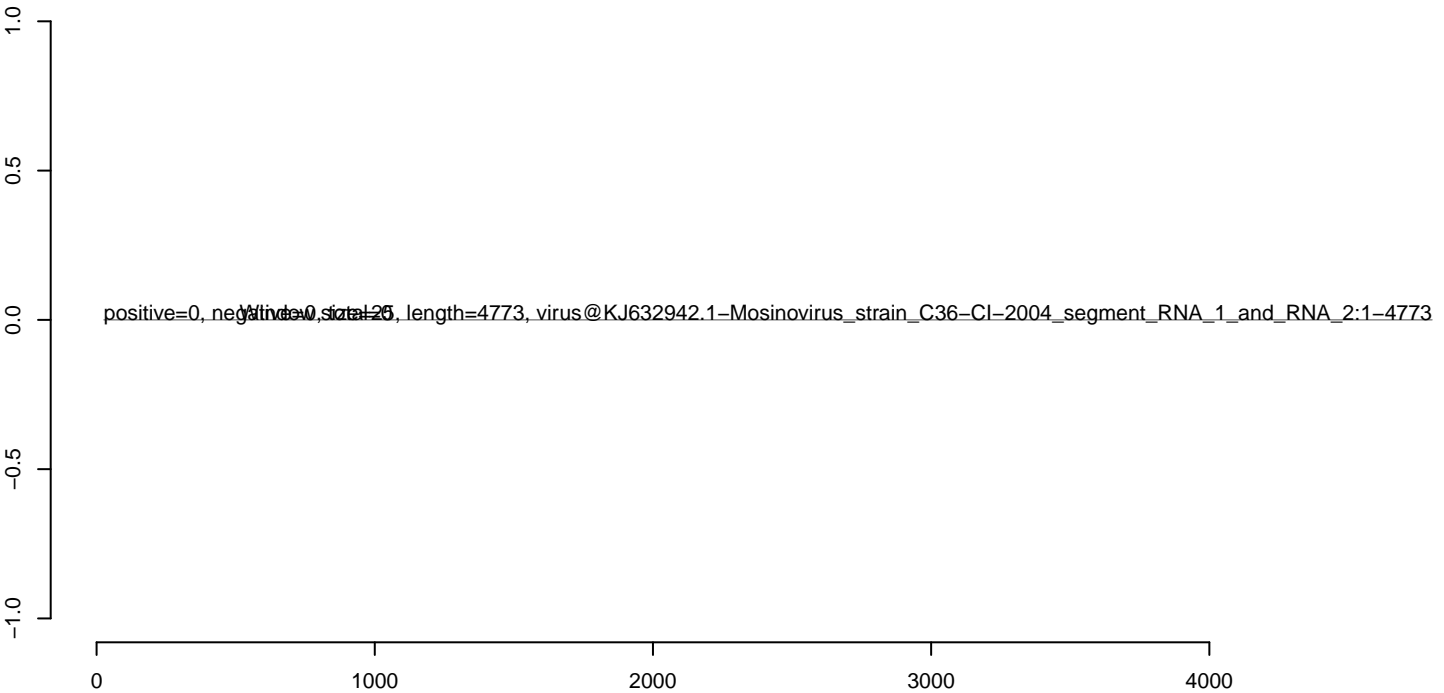
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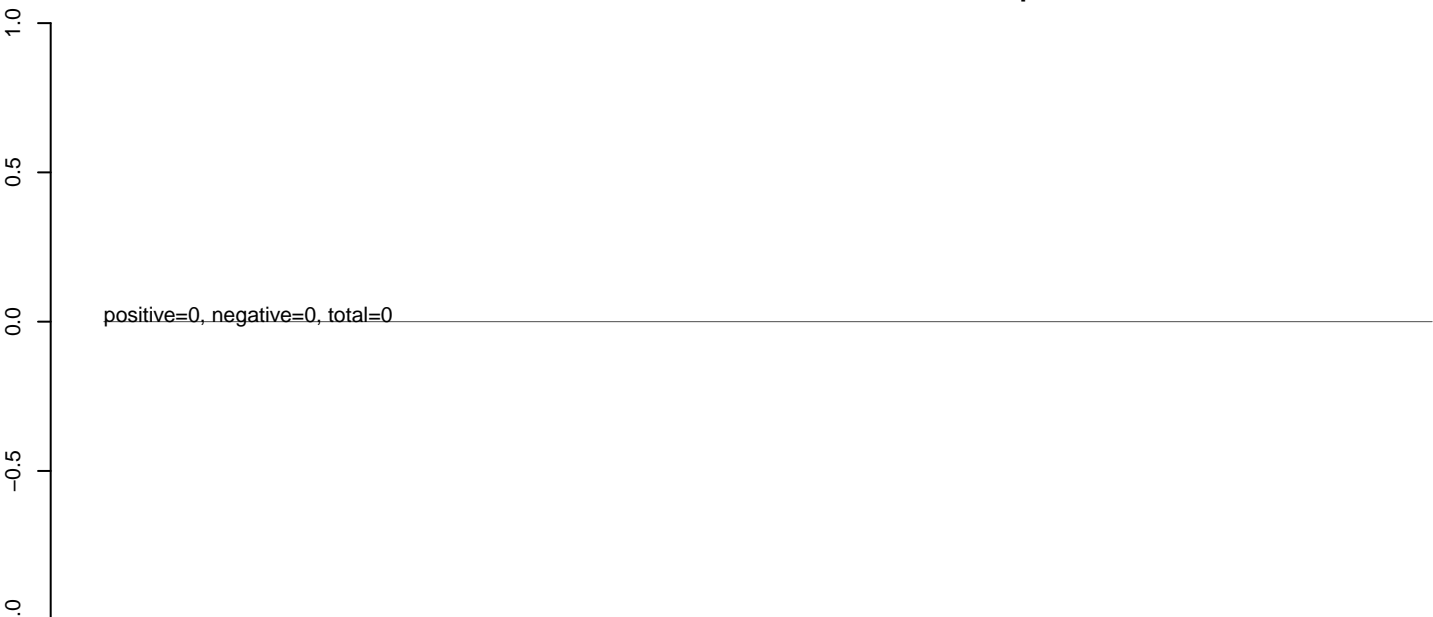
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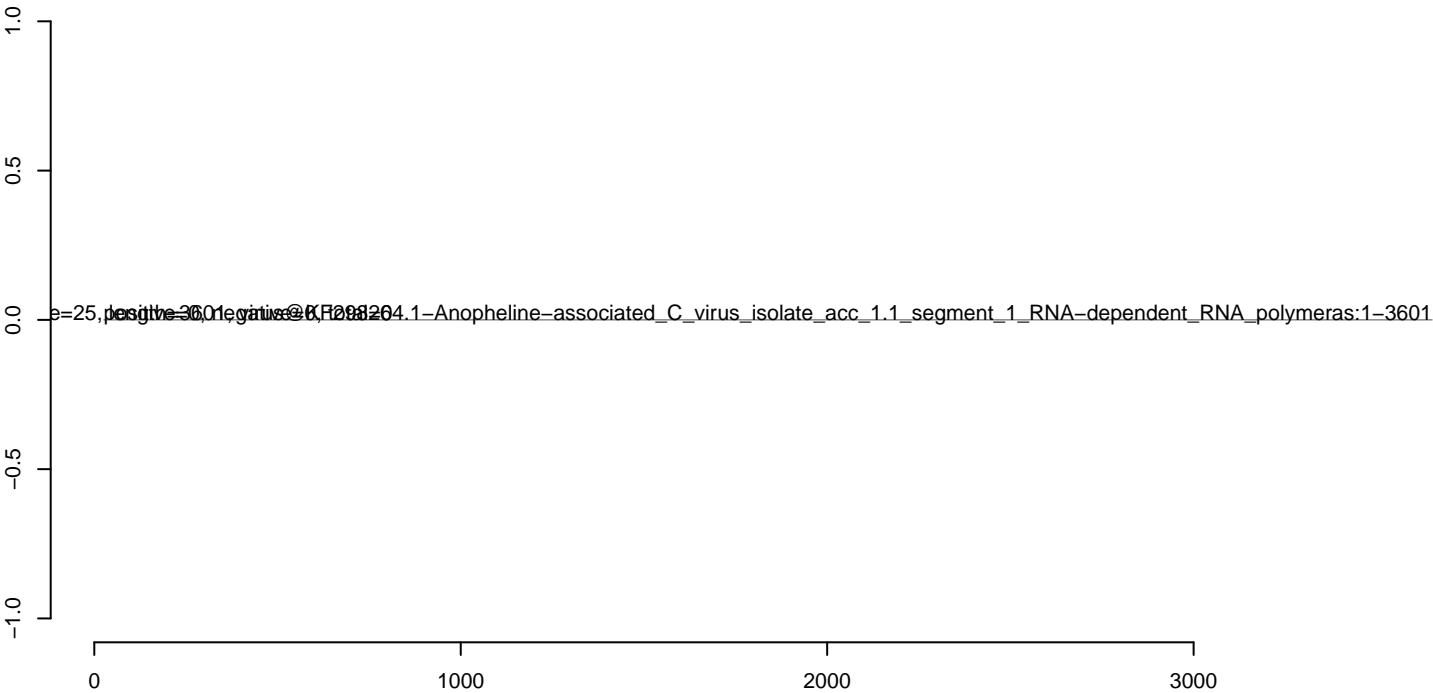
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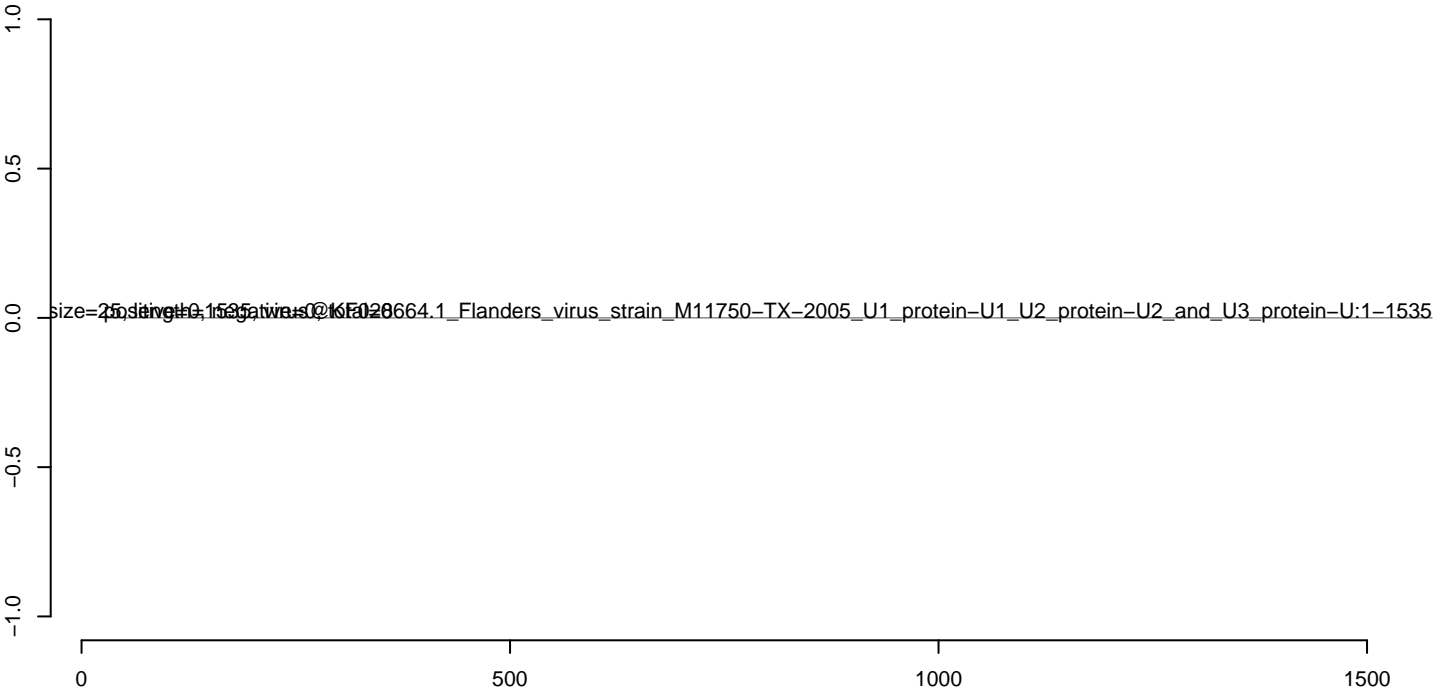
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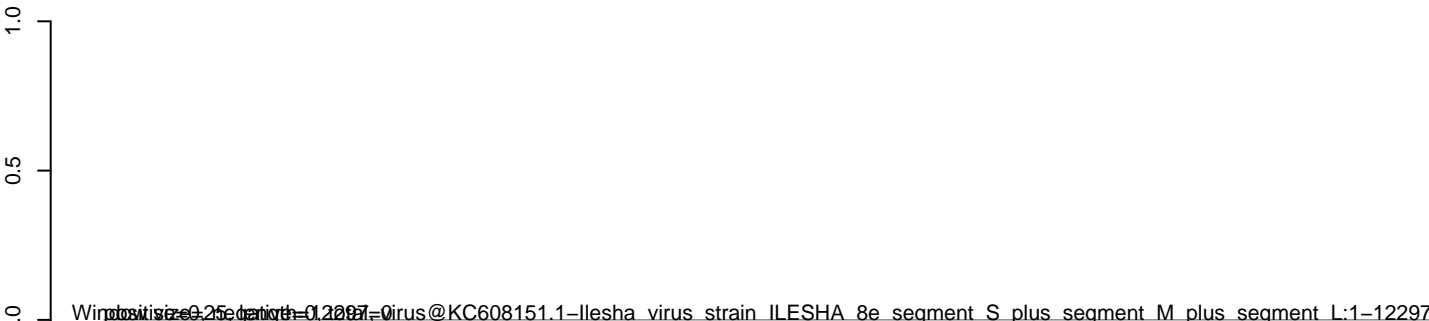
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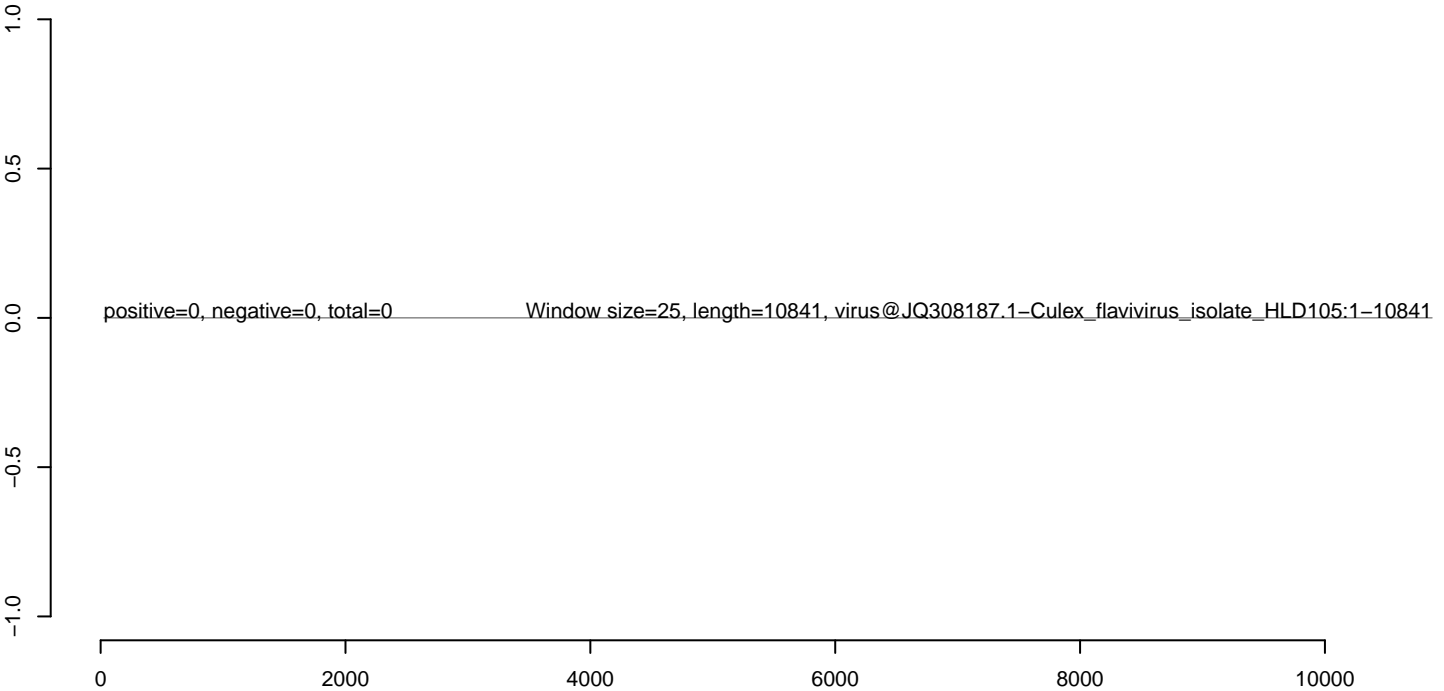
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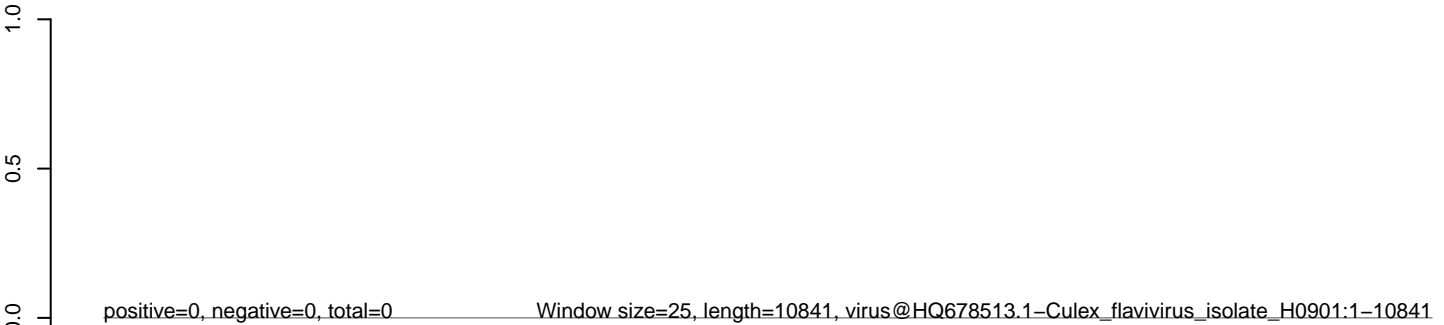
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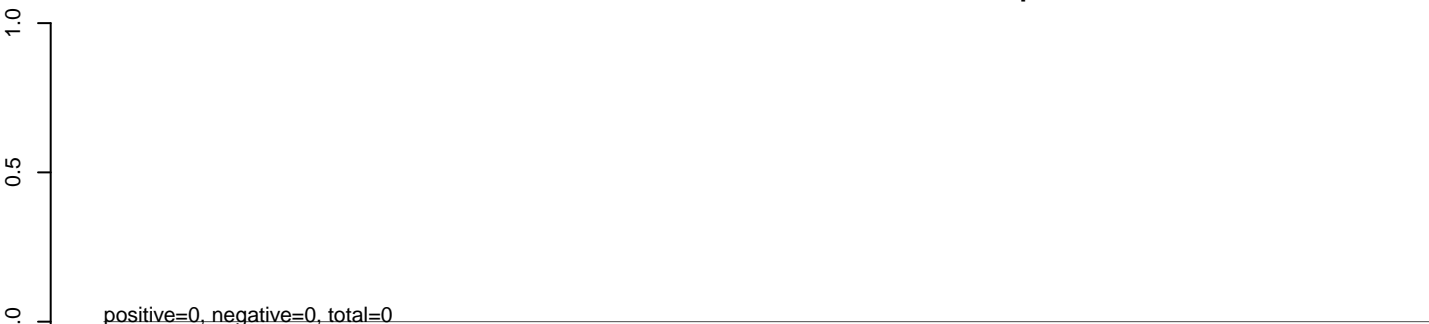


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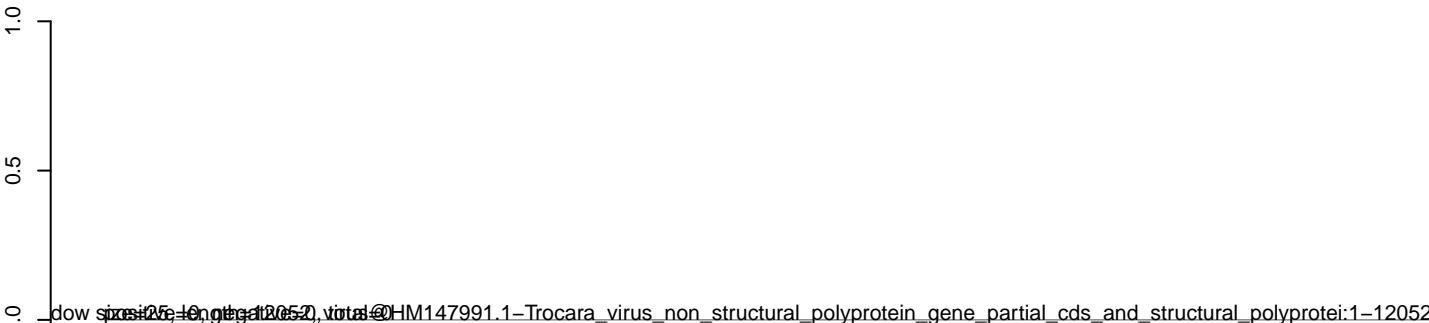
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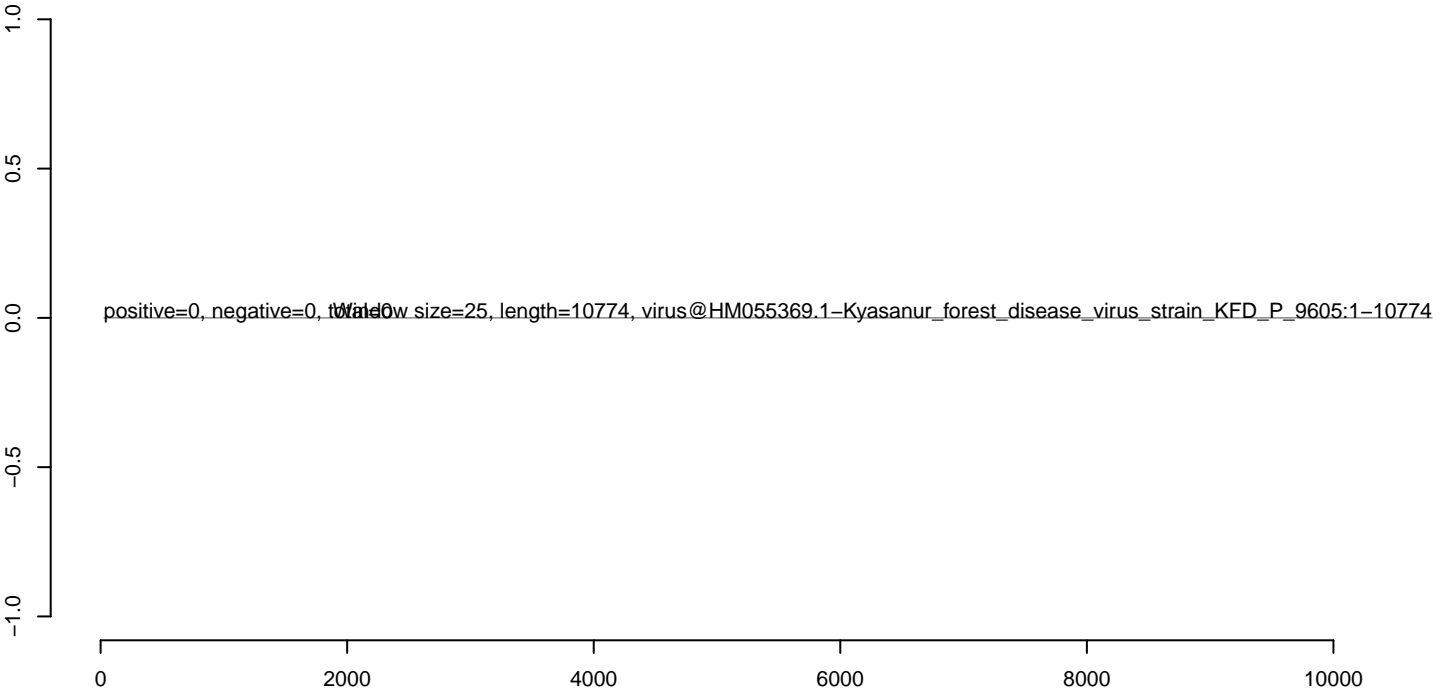
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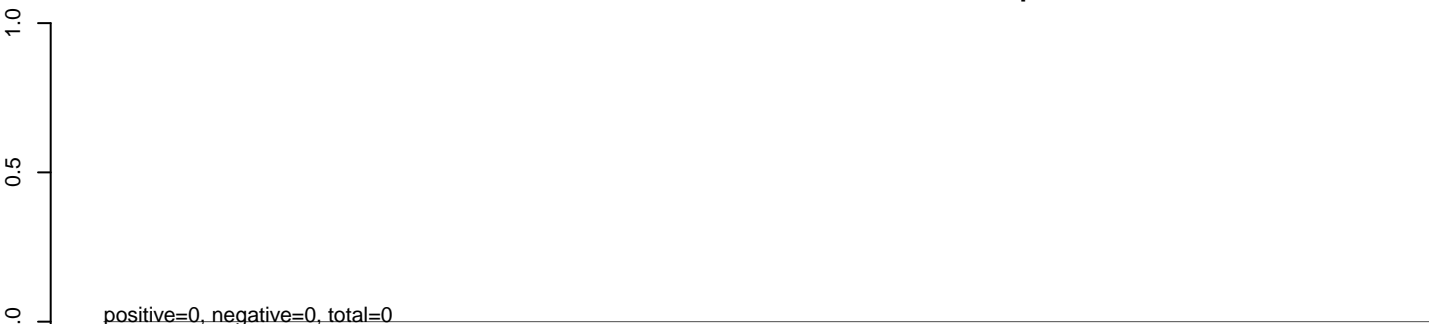
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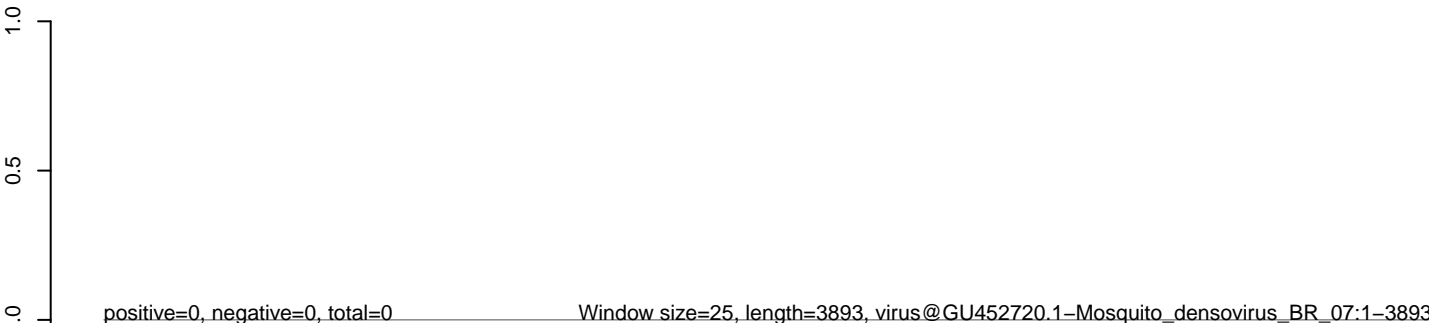
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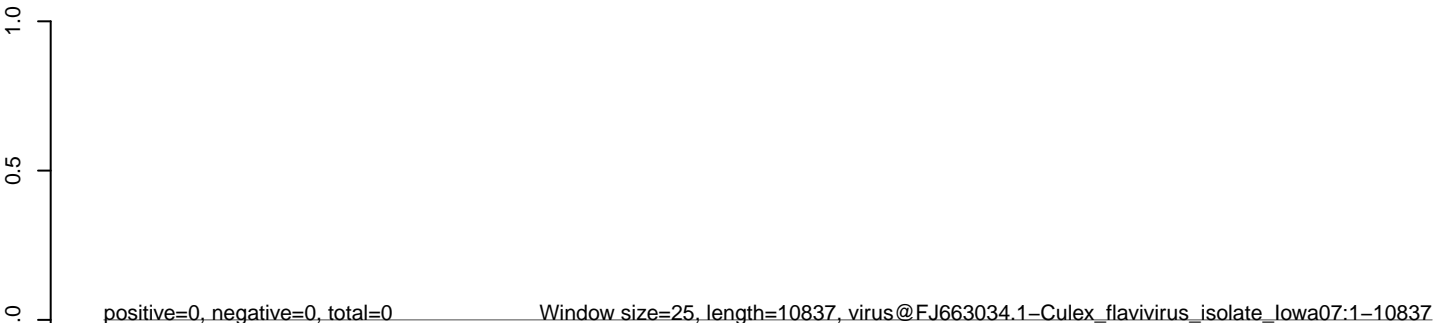
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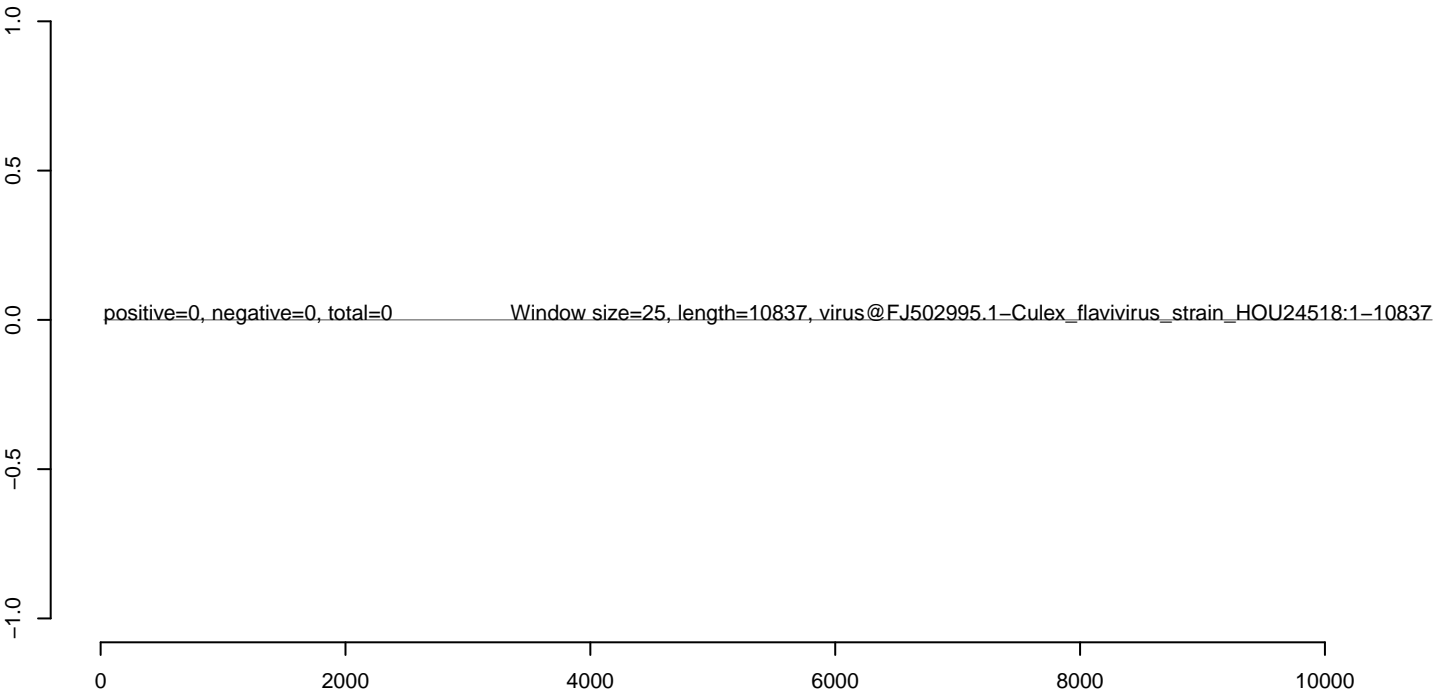
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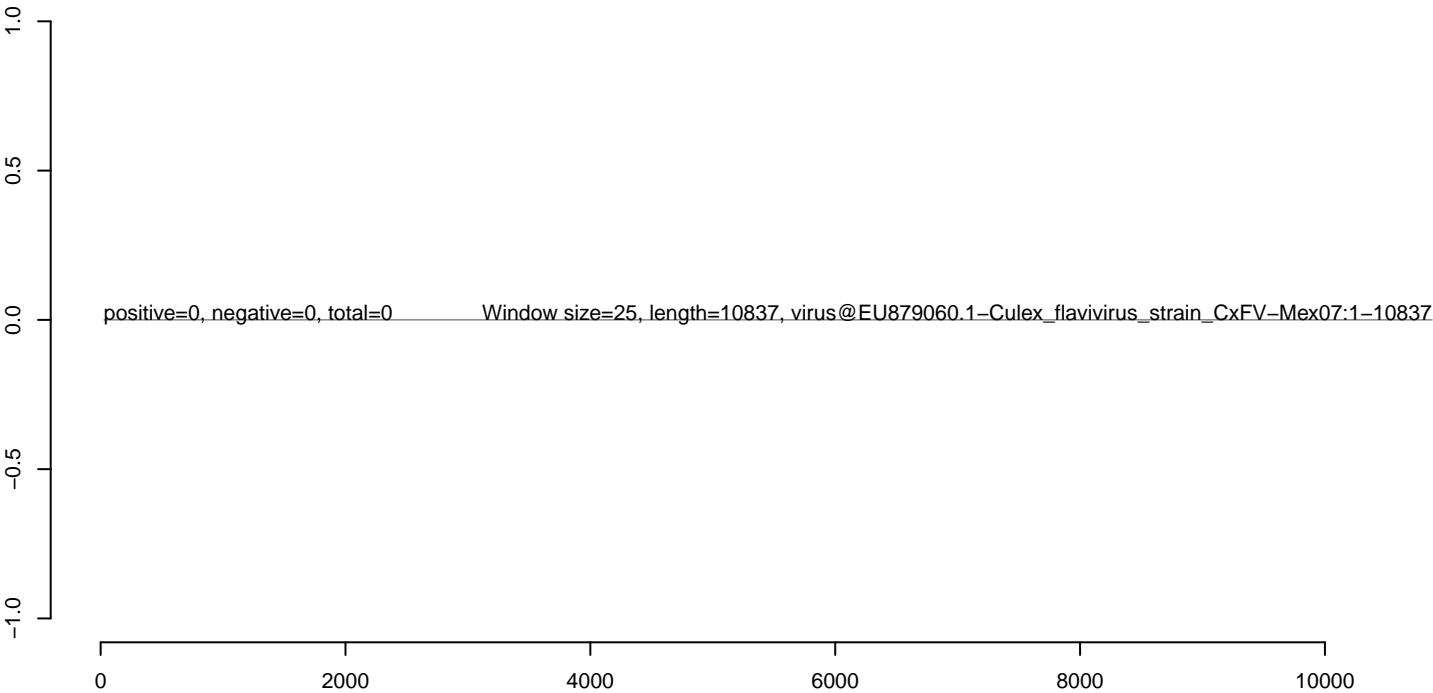
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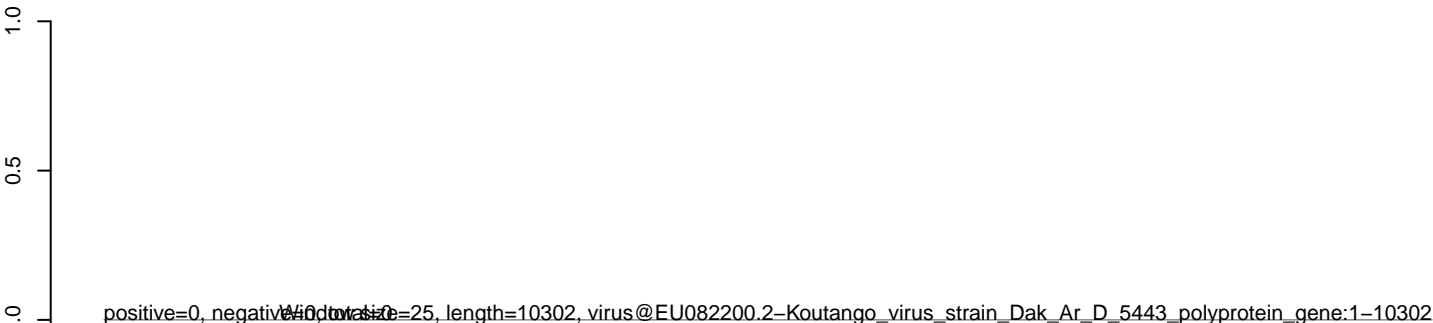
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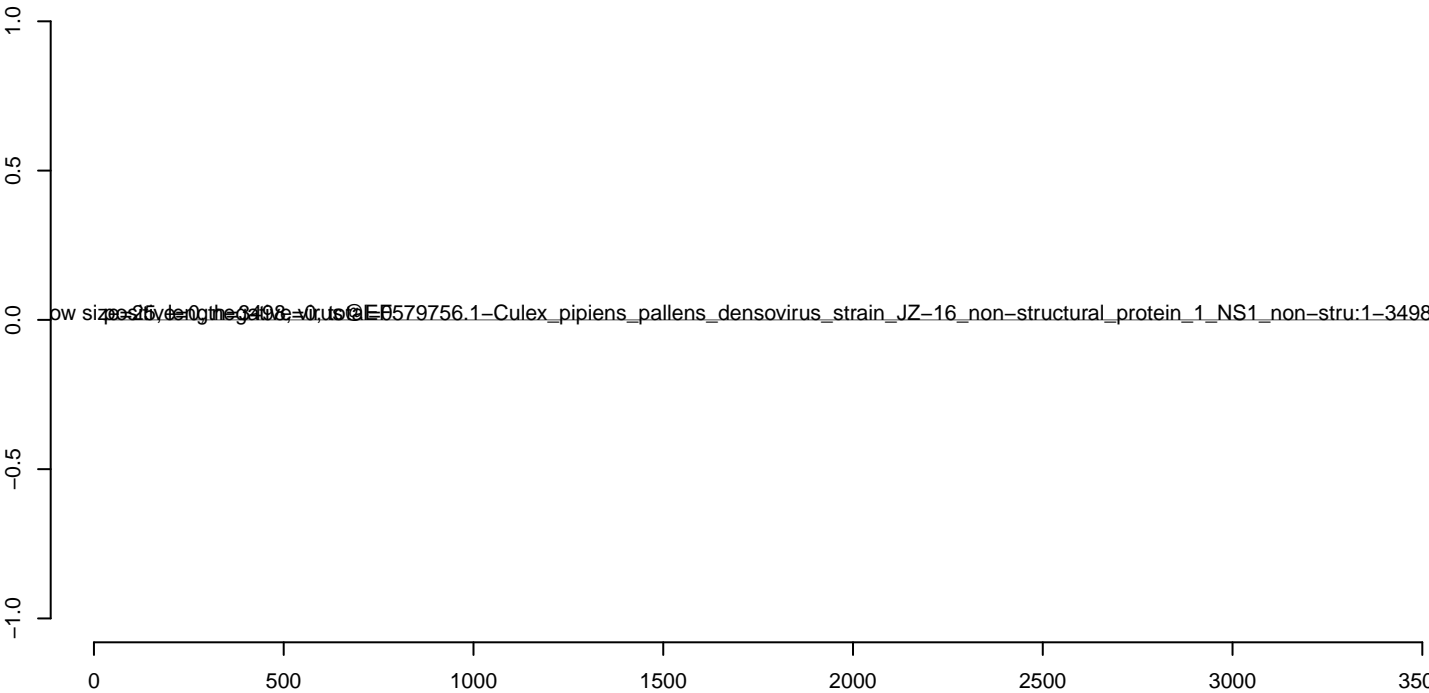
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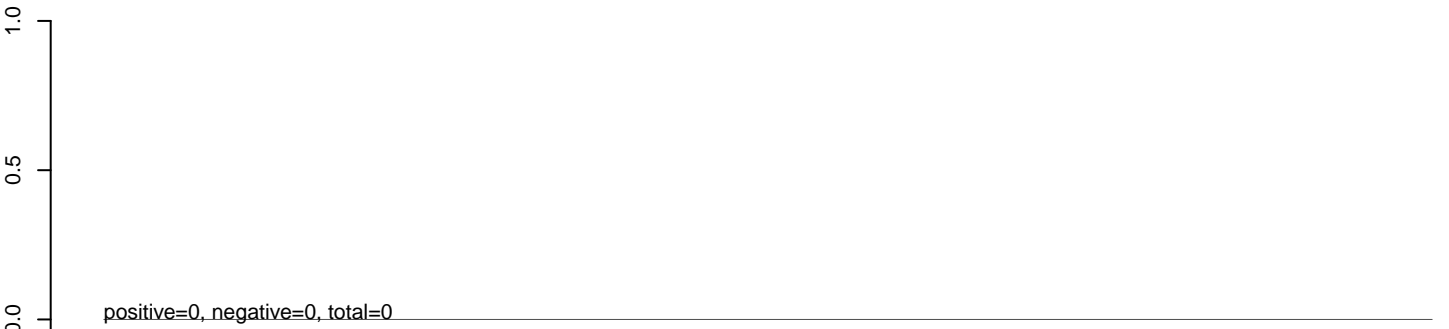
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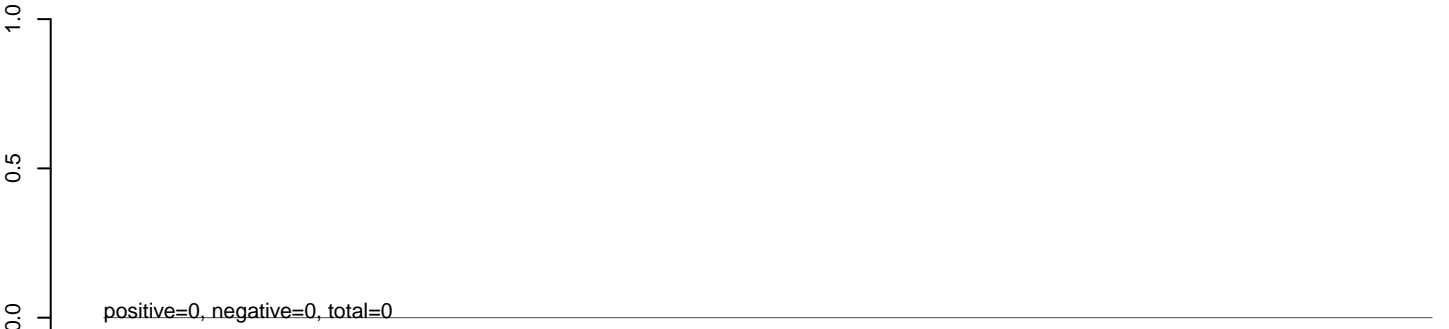
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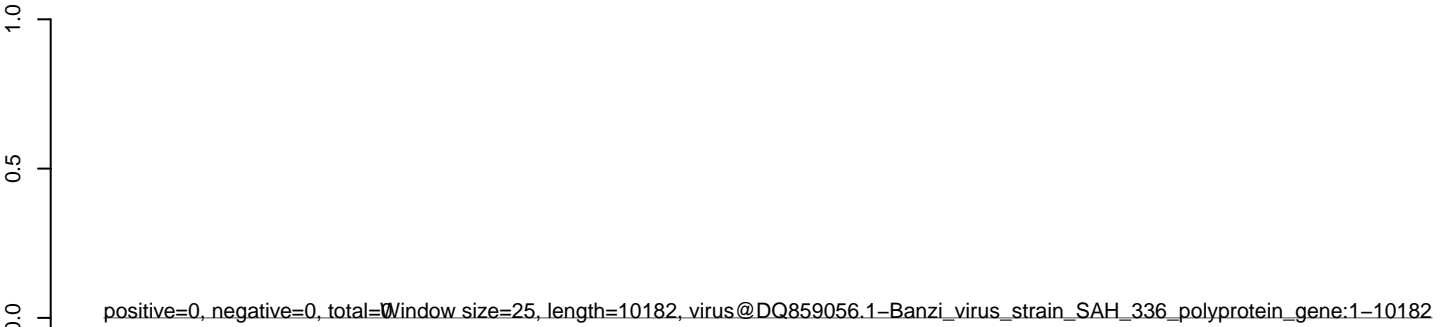
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AeAlbo_C636_SINV_GFP_betaE.rep



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AeAlbo_C636_SINV_GFP_betaE.18_23.rep



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AeAlbo_C636_SINV_GFP_betaE.rep



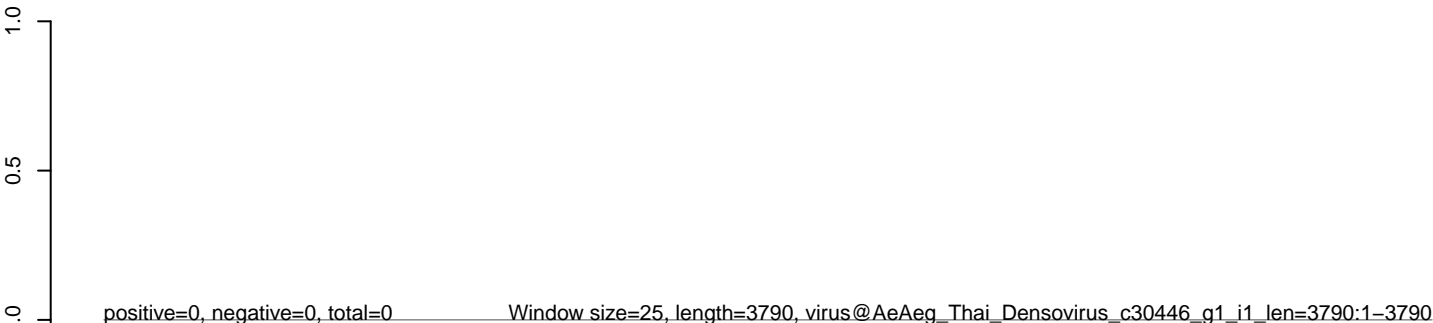
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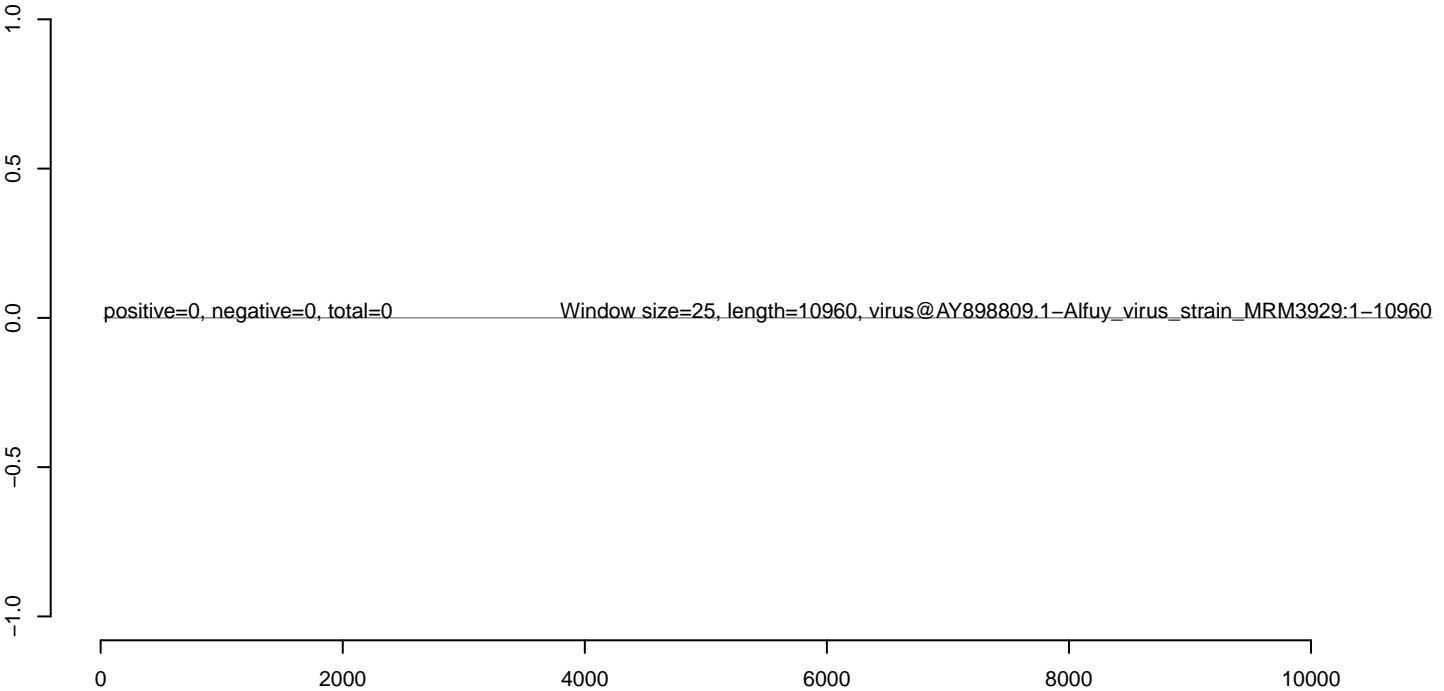
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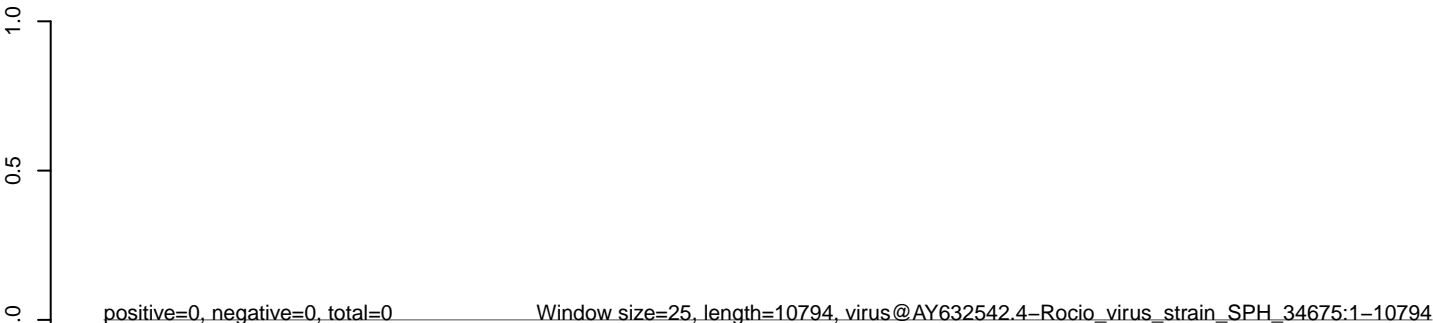
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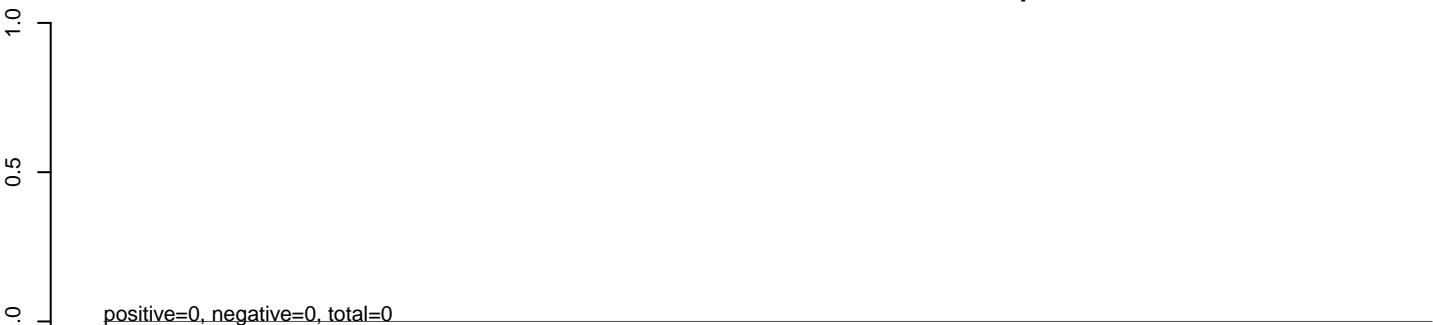


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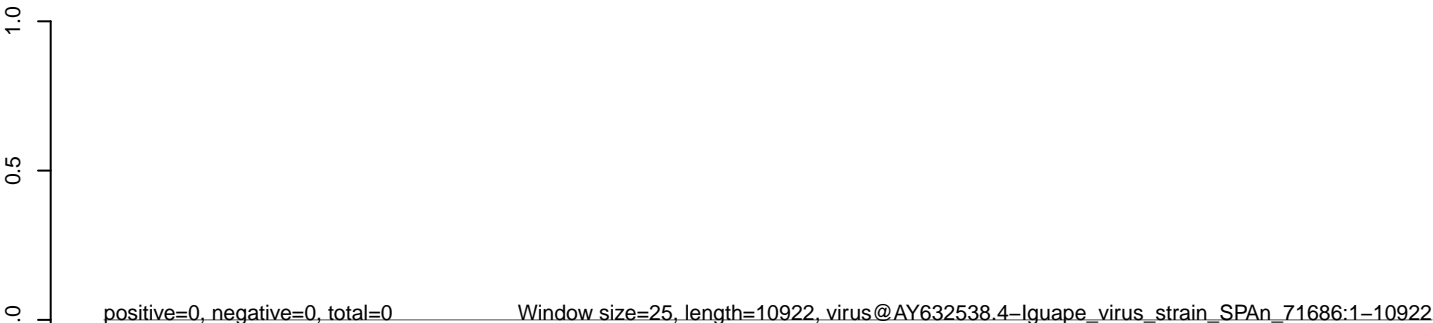
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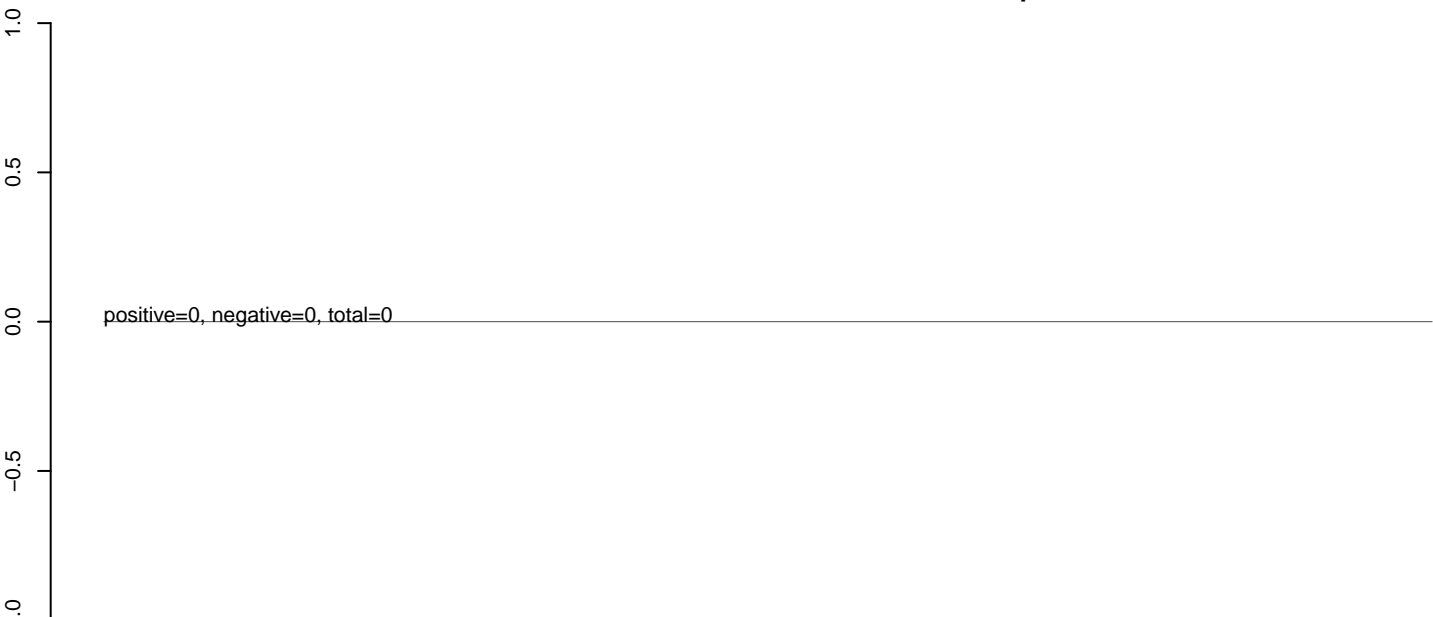


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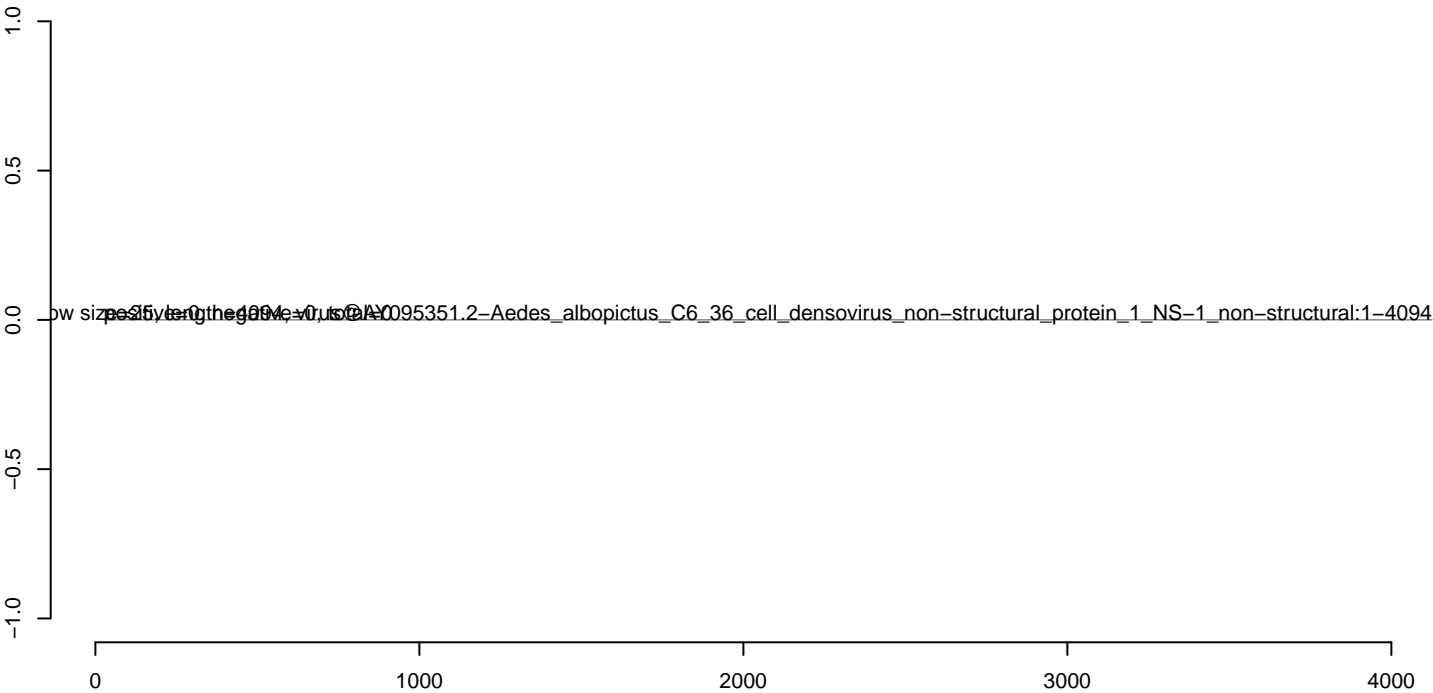
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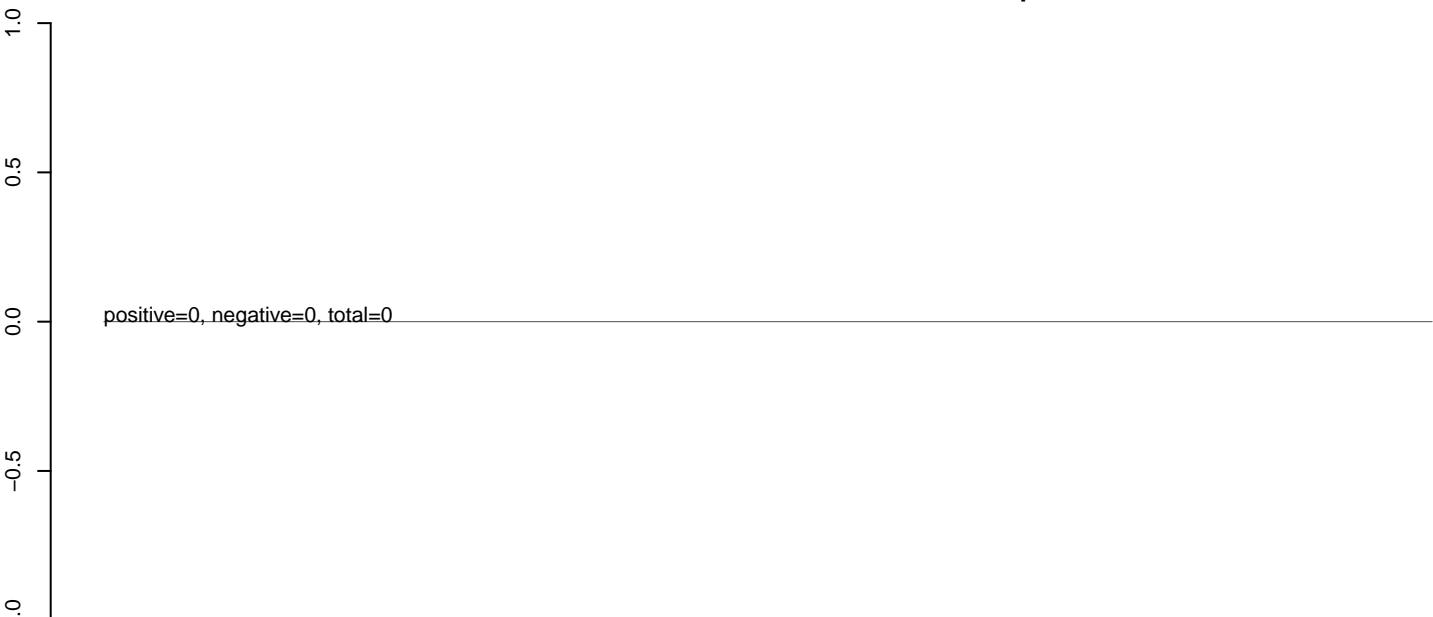
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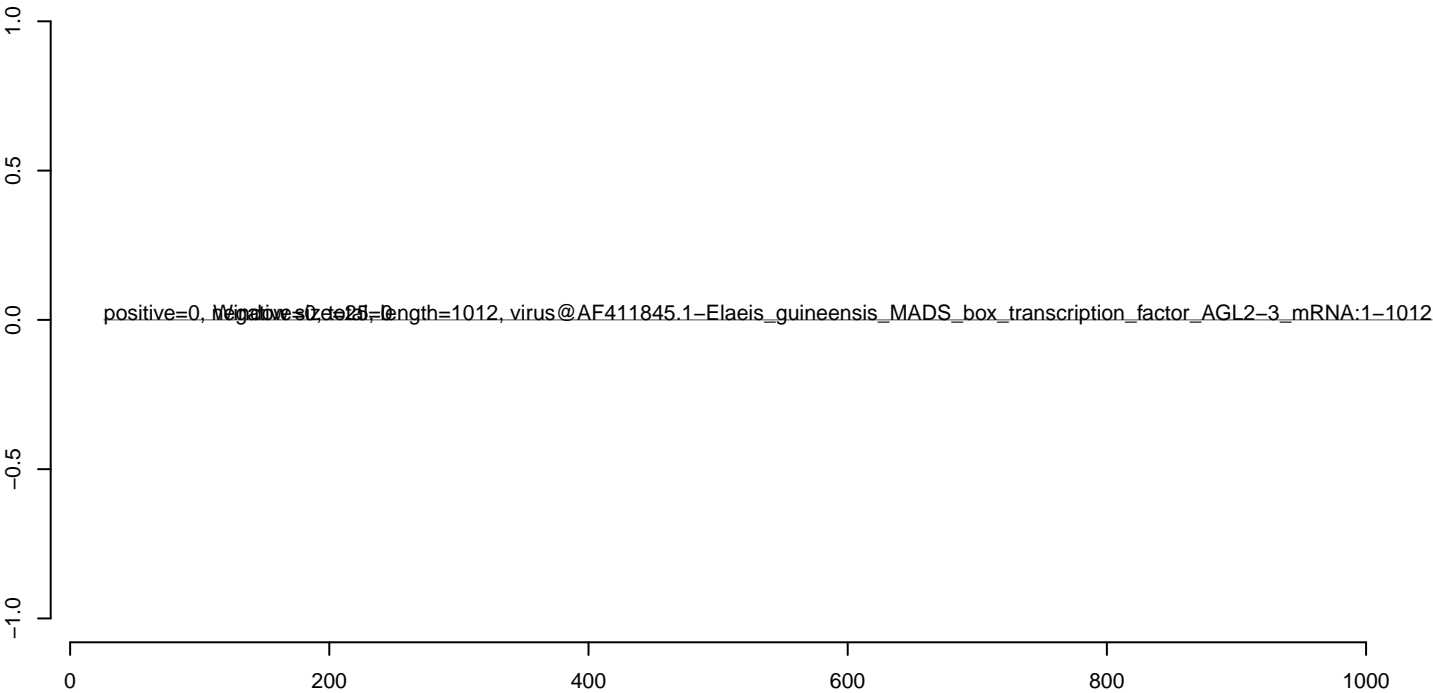
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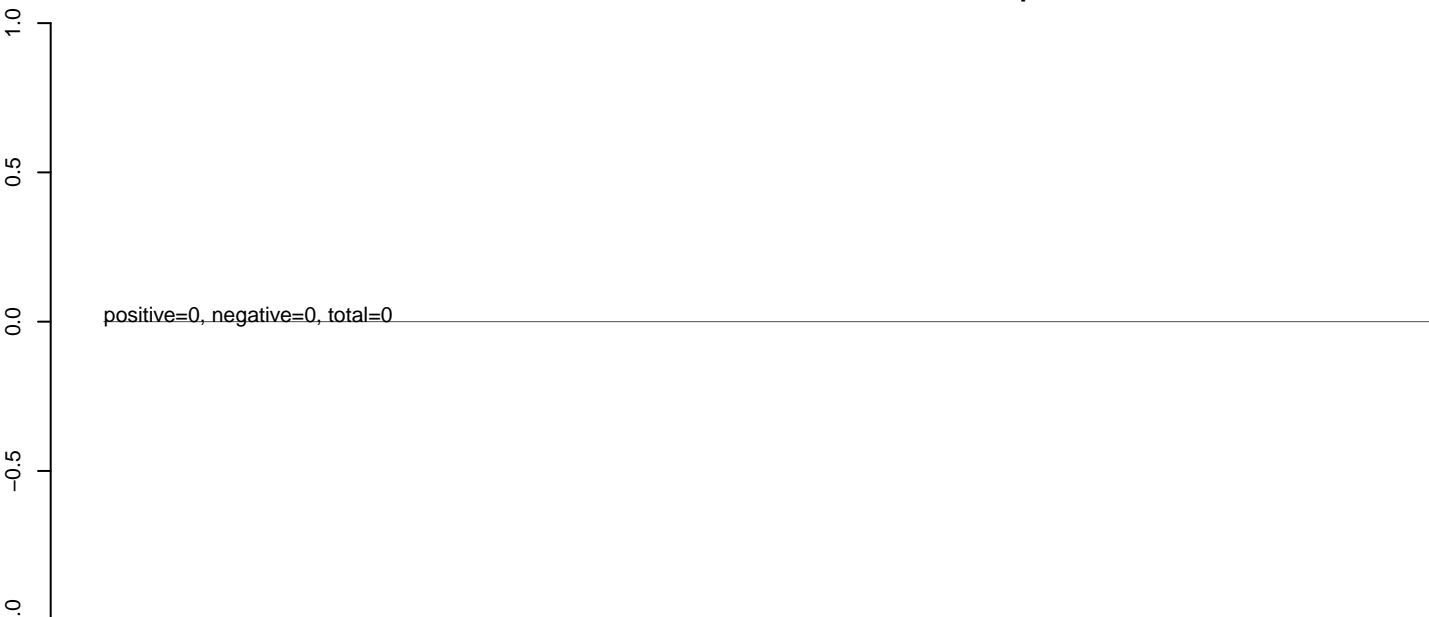
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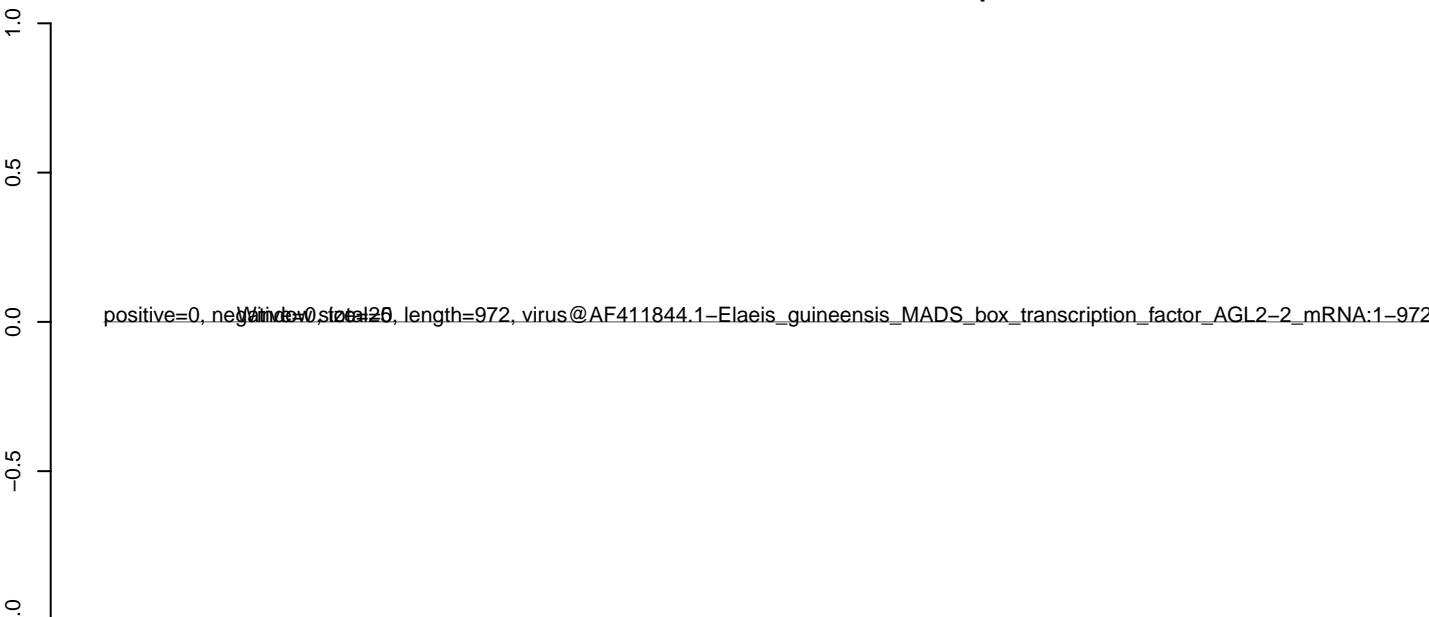
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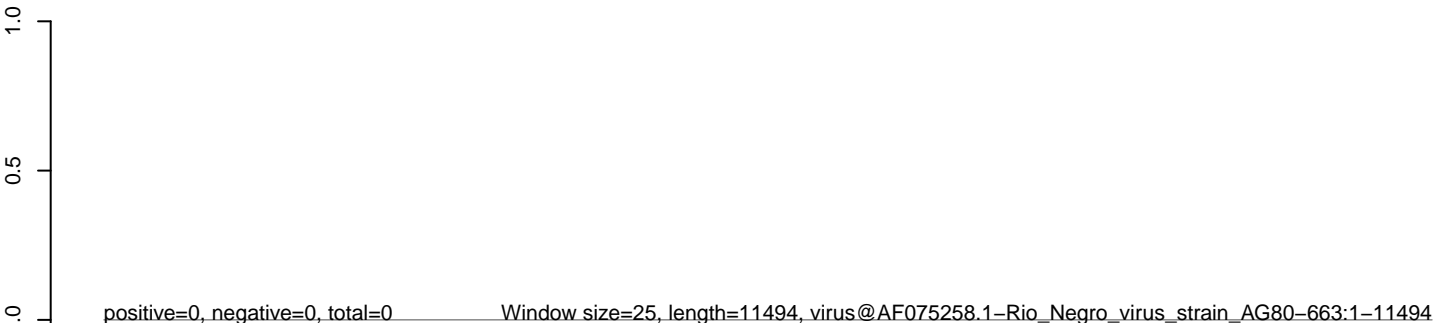
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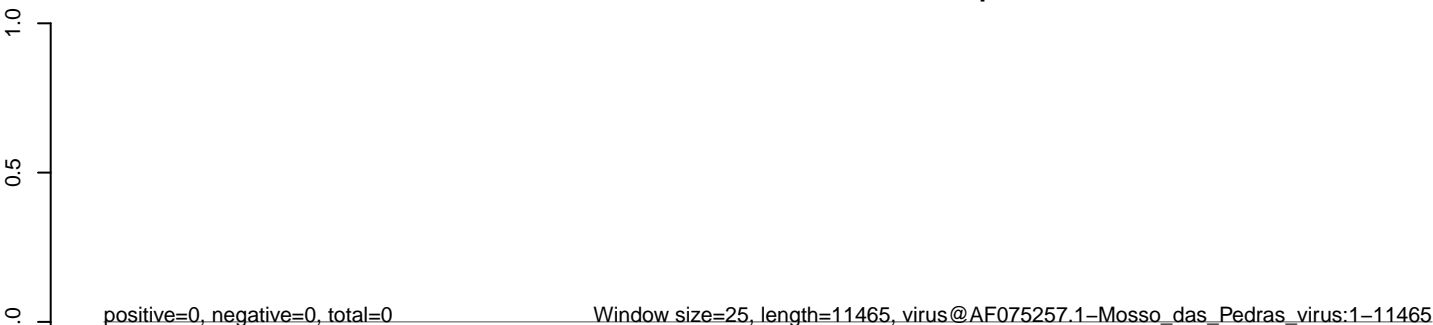
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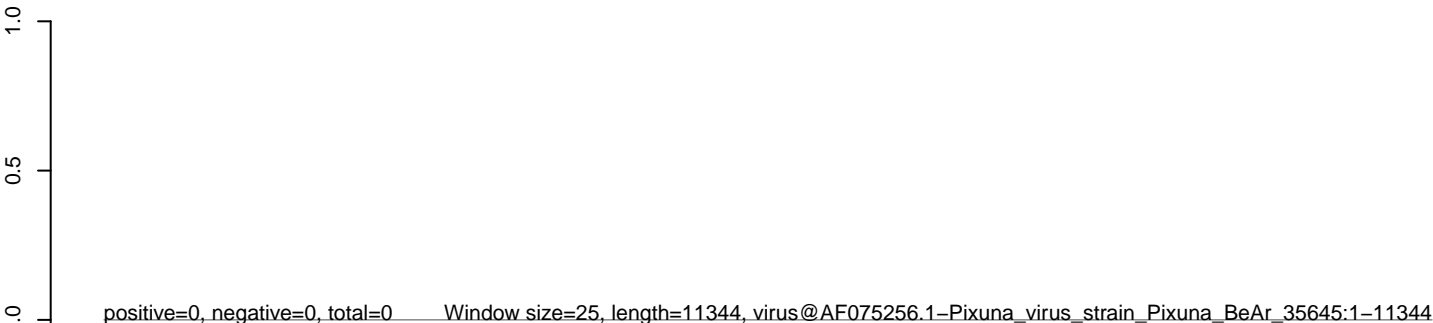
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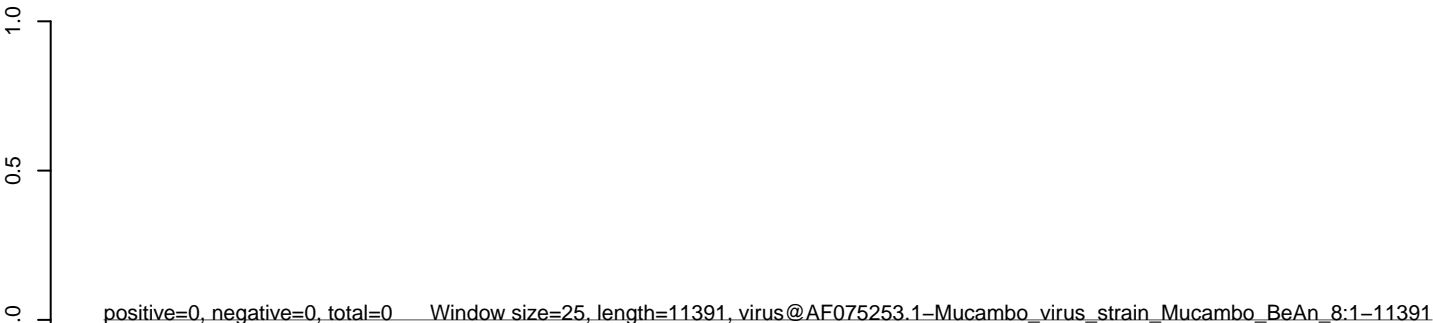
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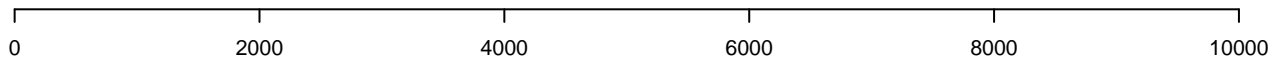
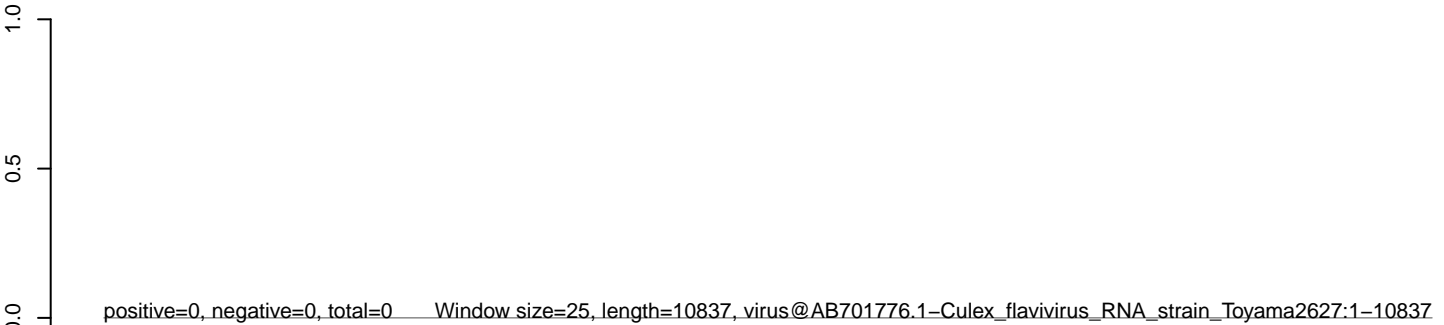
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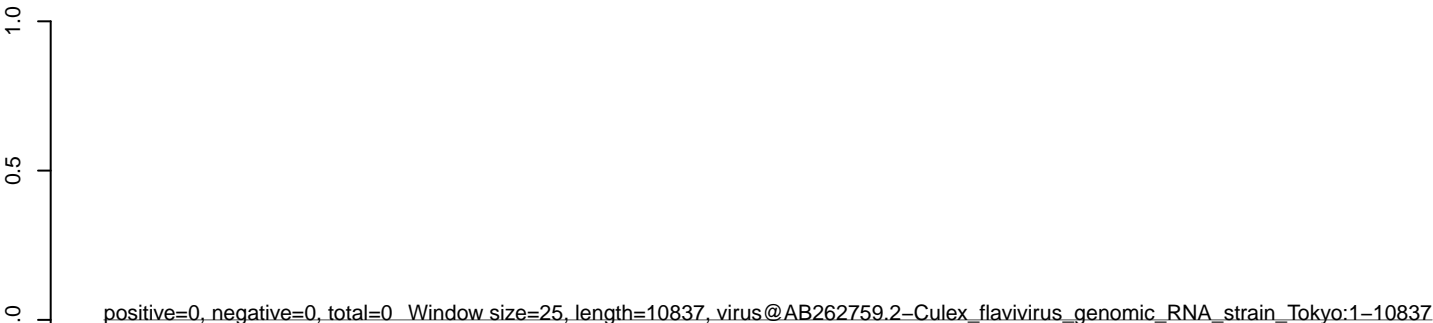
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