

AeAeg_Aag2_NL.M3.18_23.rep



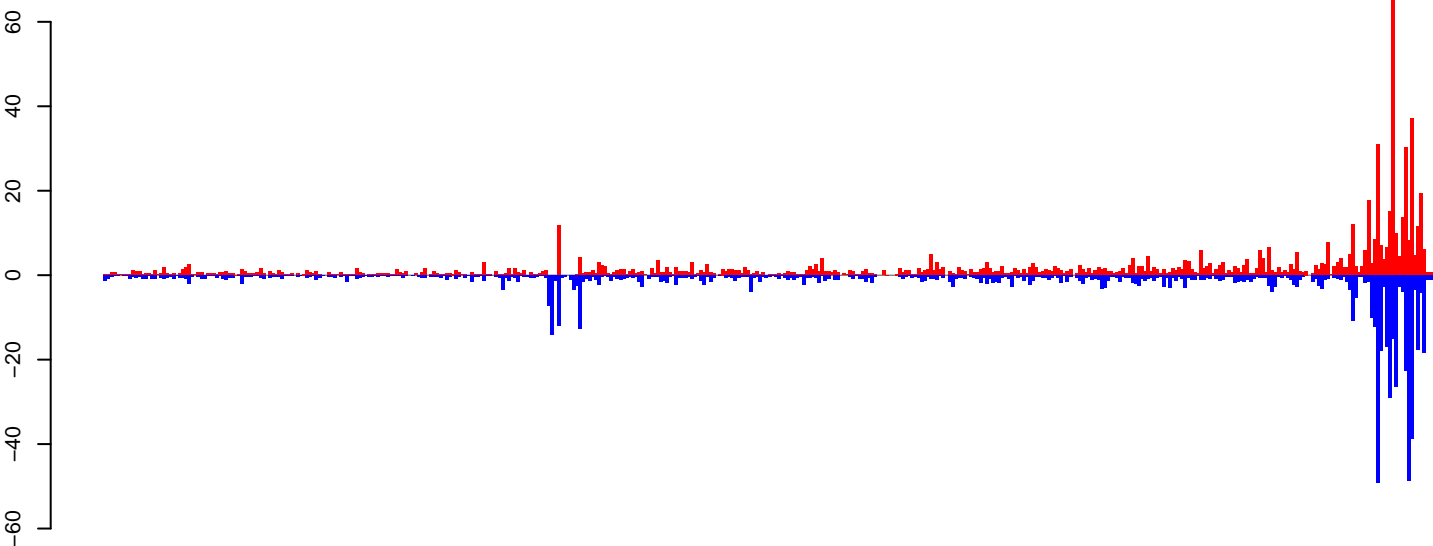
AeAeg_Aag2_NL.M3.24_35.rep



AeAeg_Aag2_NL.M3.rep

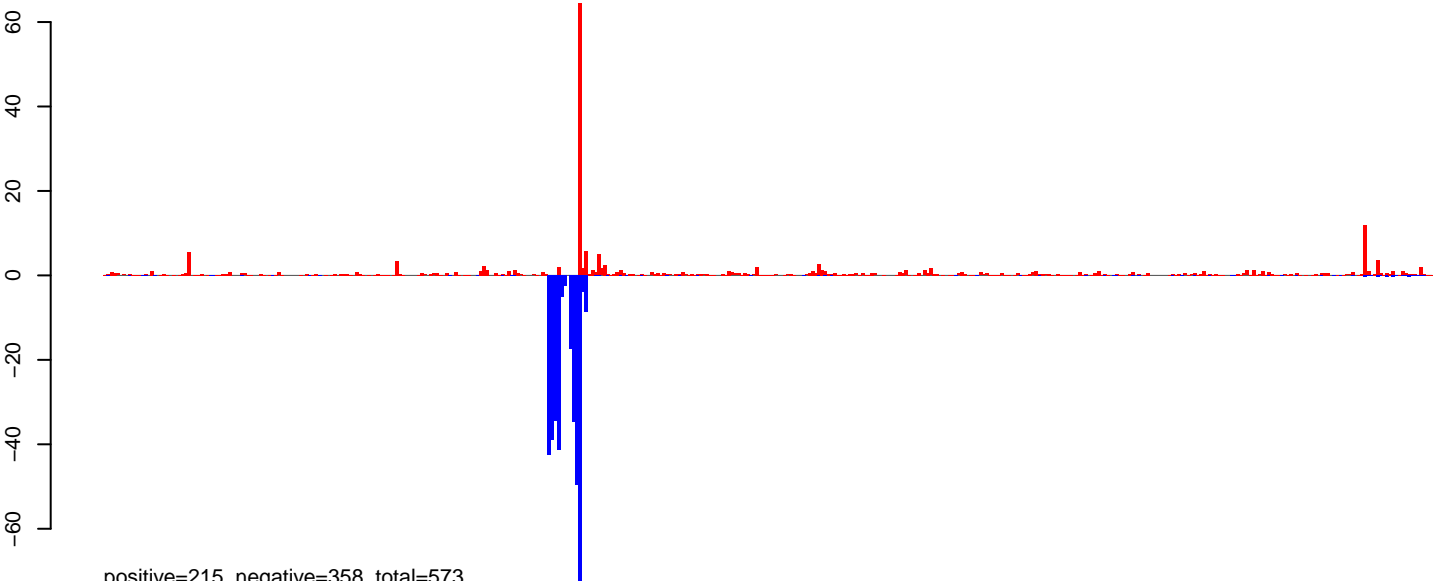


AeAeg_Aag2_NL.M3.18_23.rep



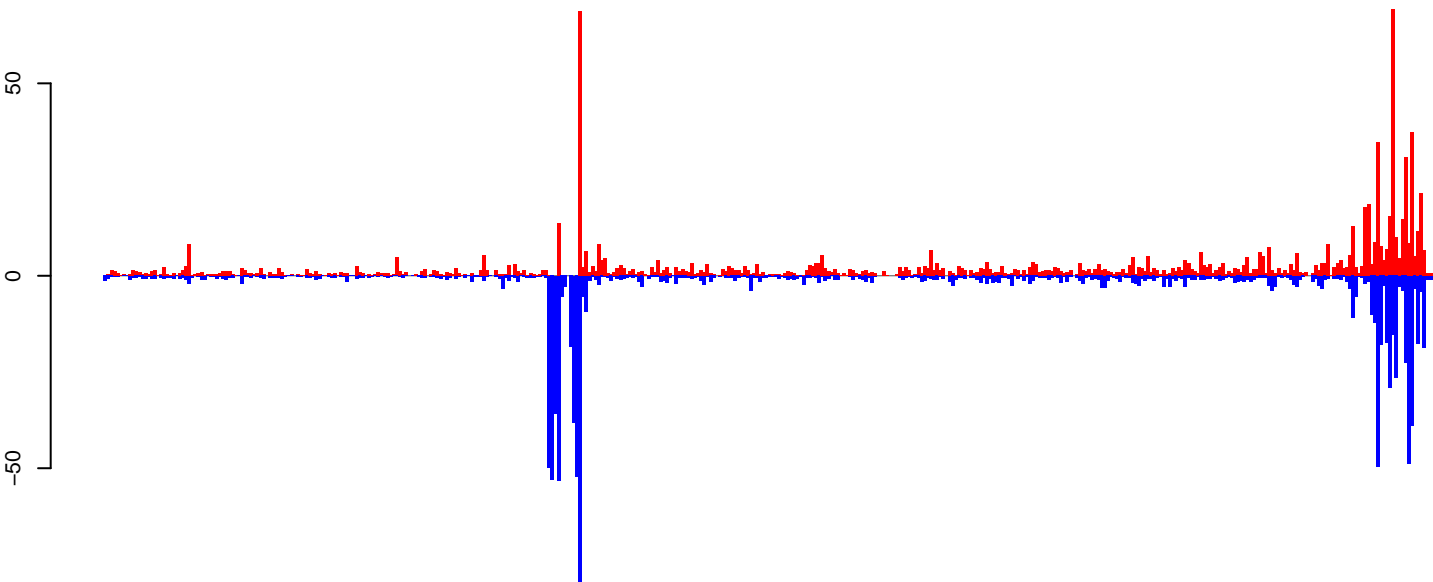
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AeAeg_Aag2_NL.M3.24_35.rep

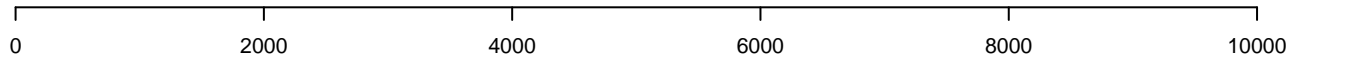


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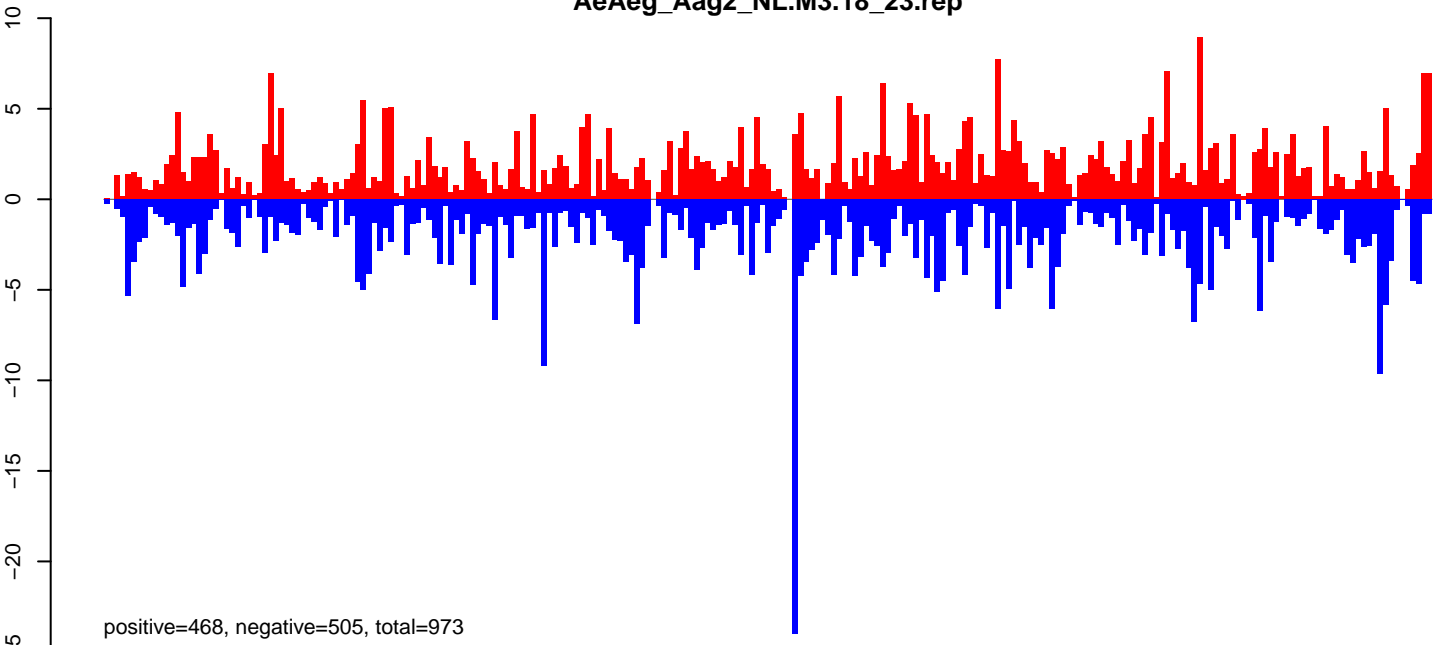
AeAeg_Aag2_NL.M3.rep



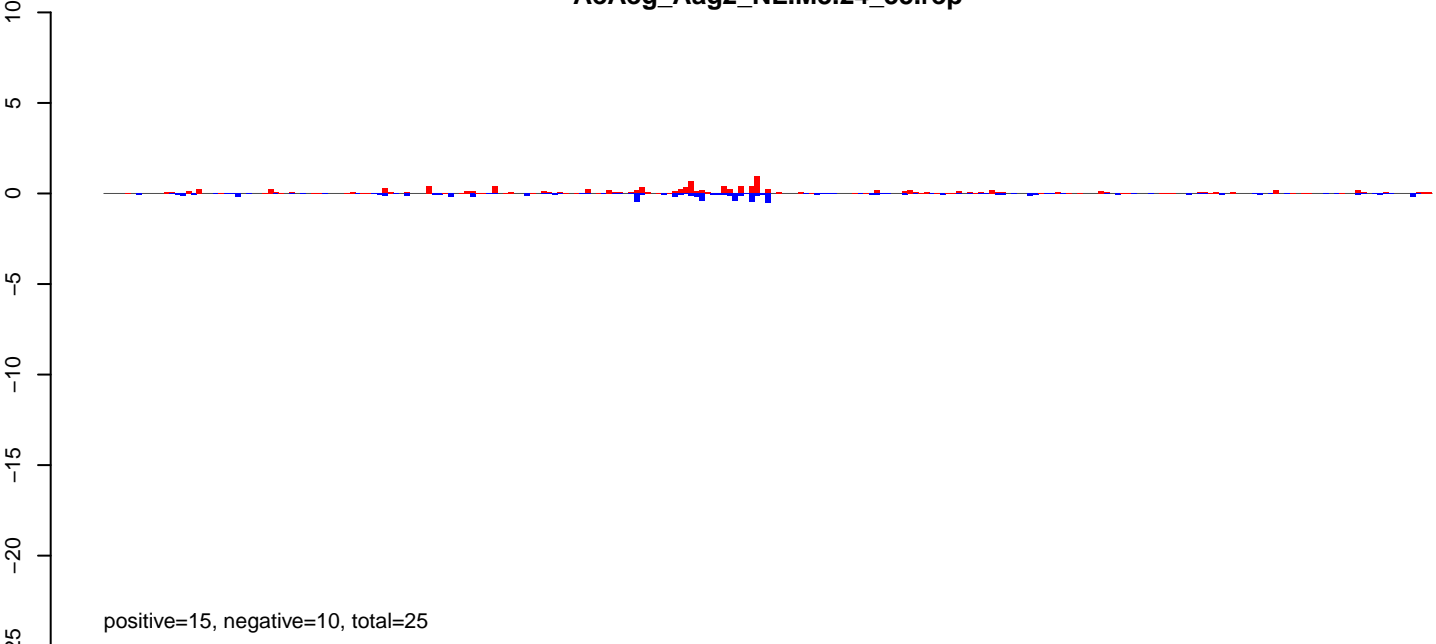
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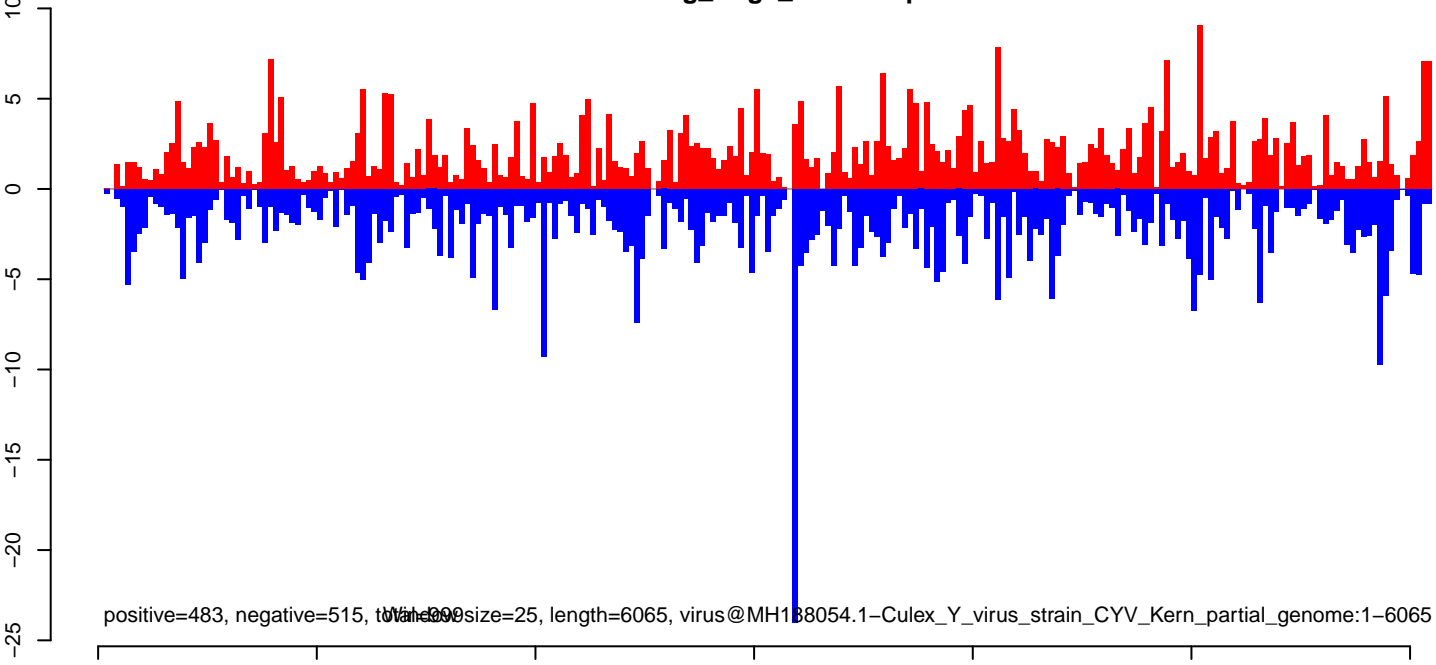
AeAeg_Aag2_NL.M3.18_23.rep



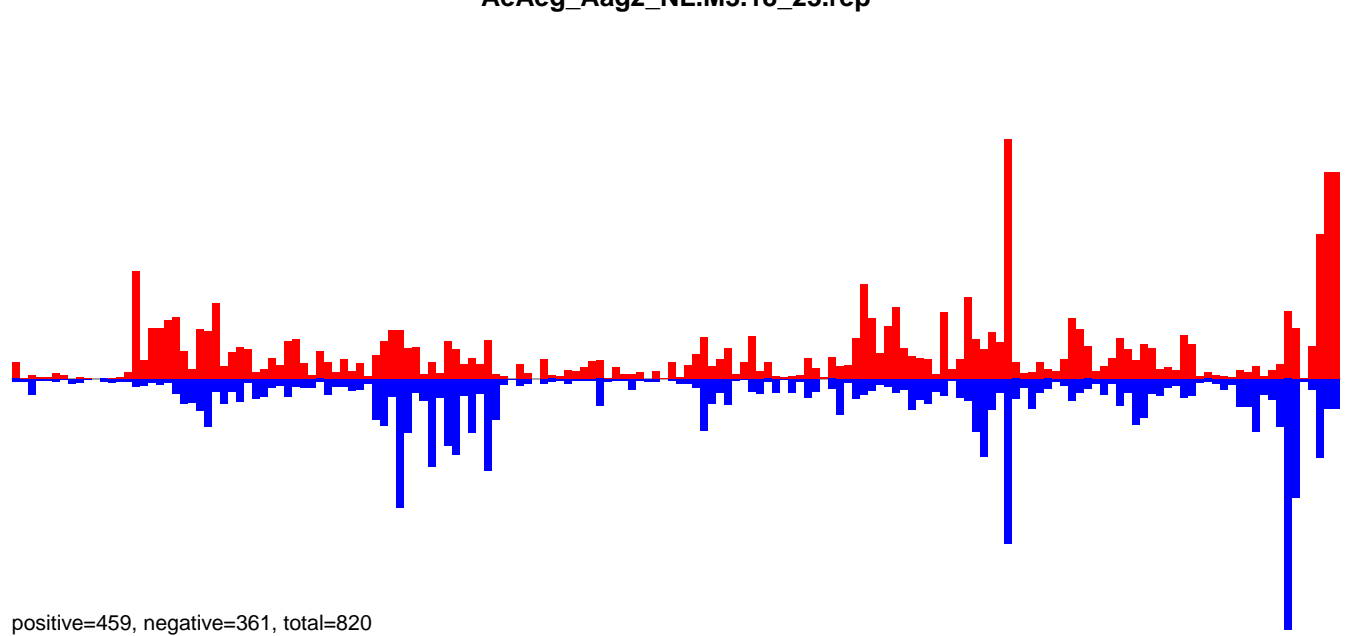
AeAeg_Aag2_NL.M3.24_35.rep



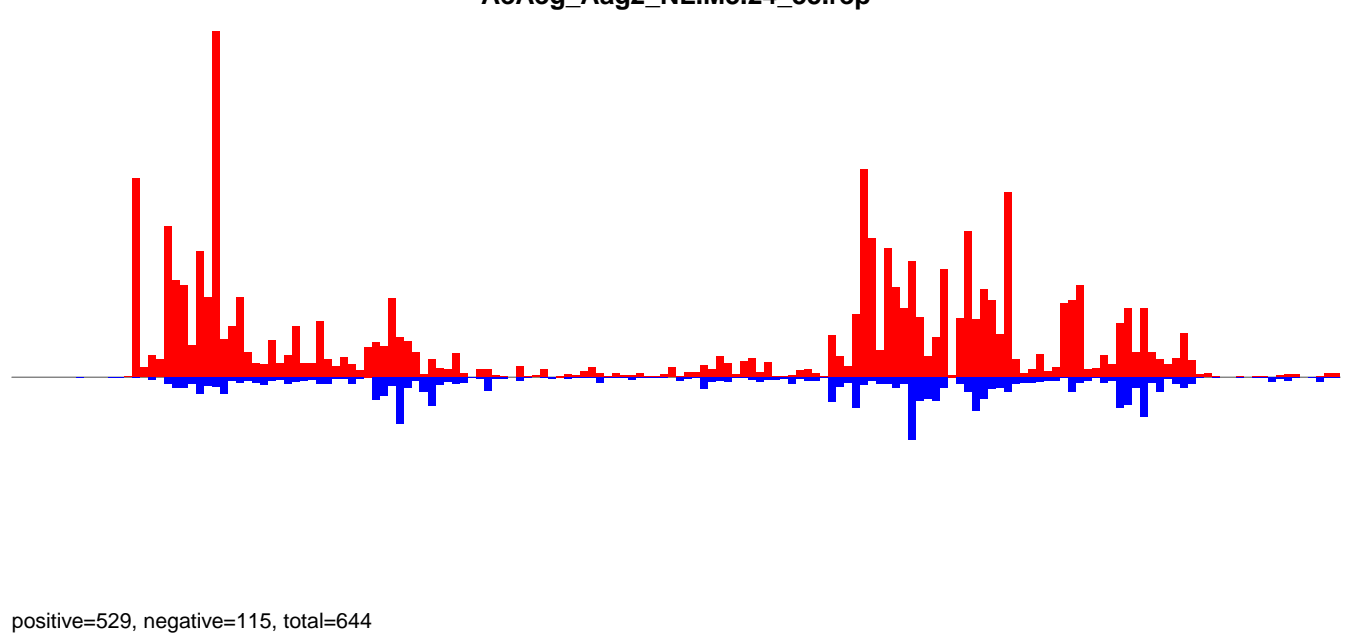
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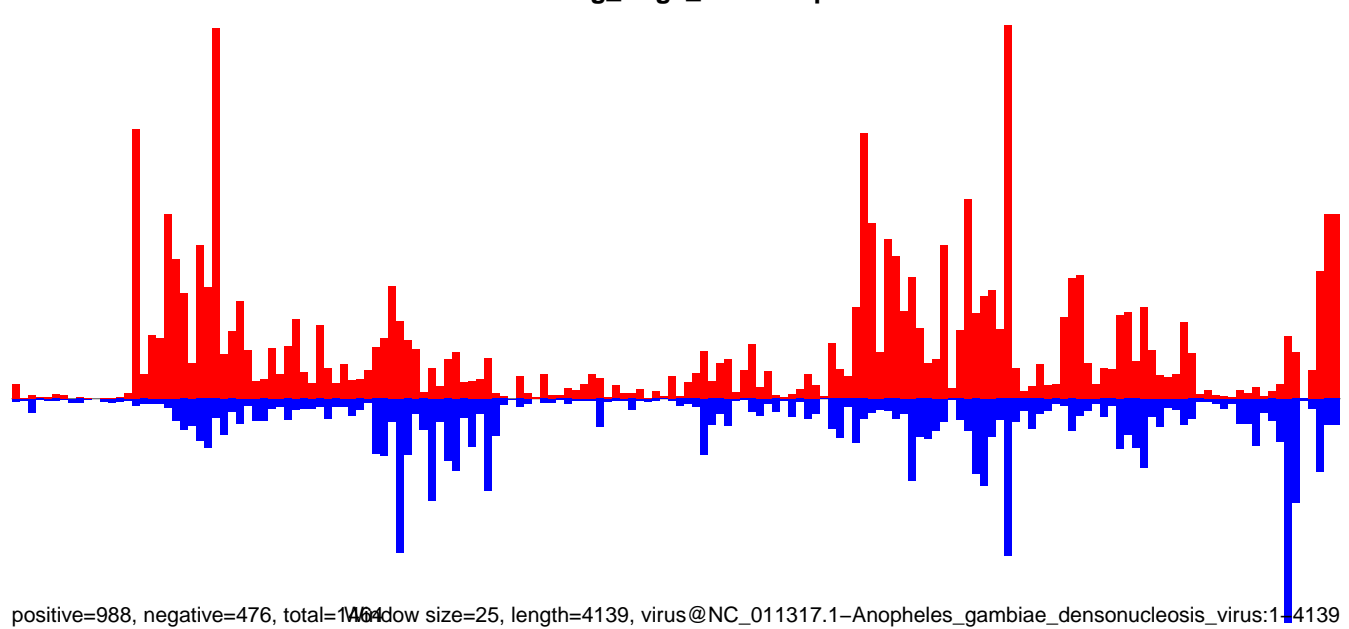
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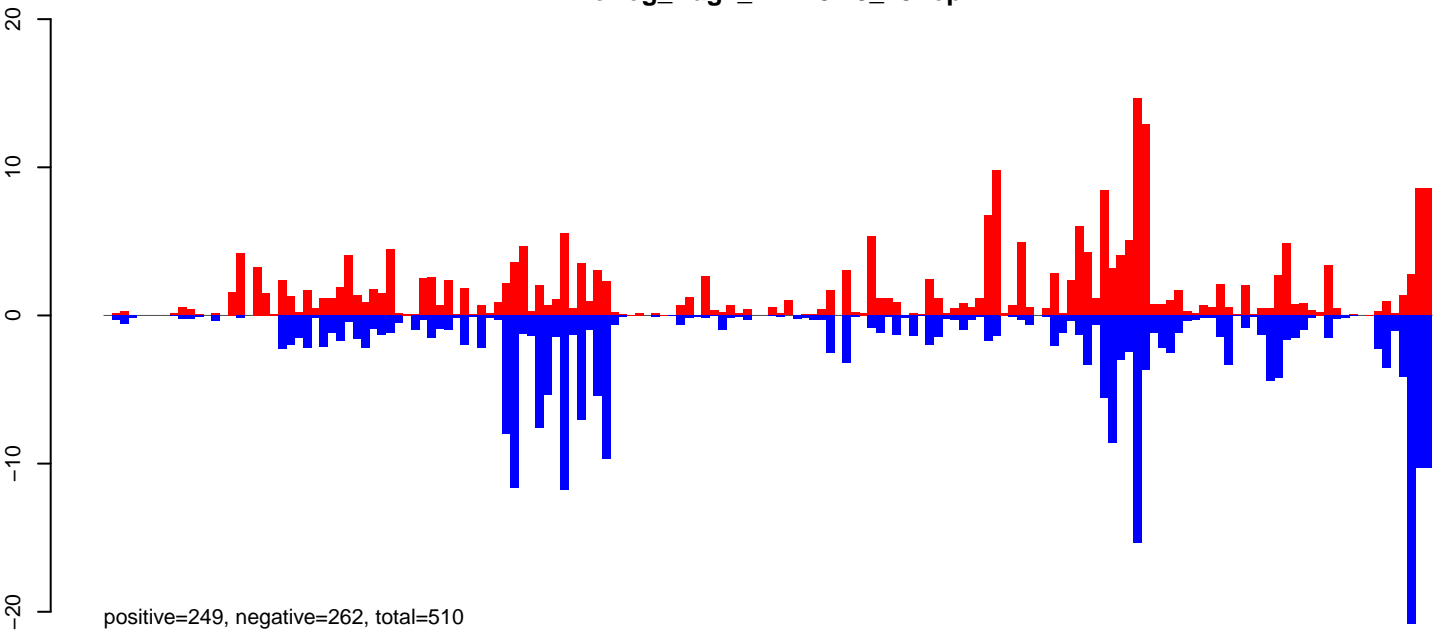
AeAeg_Aag2_NL.M3.24_35.rep



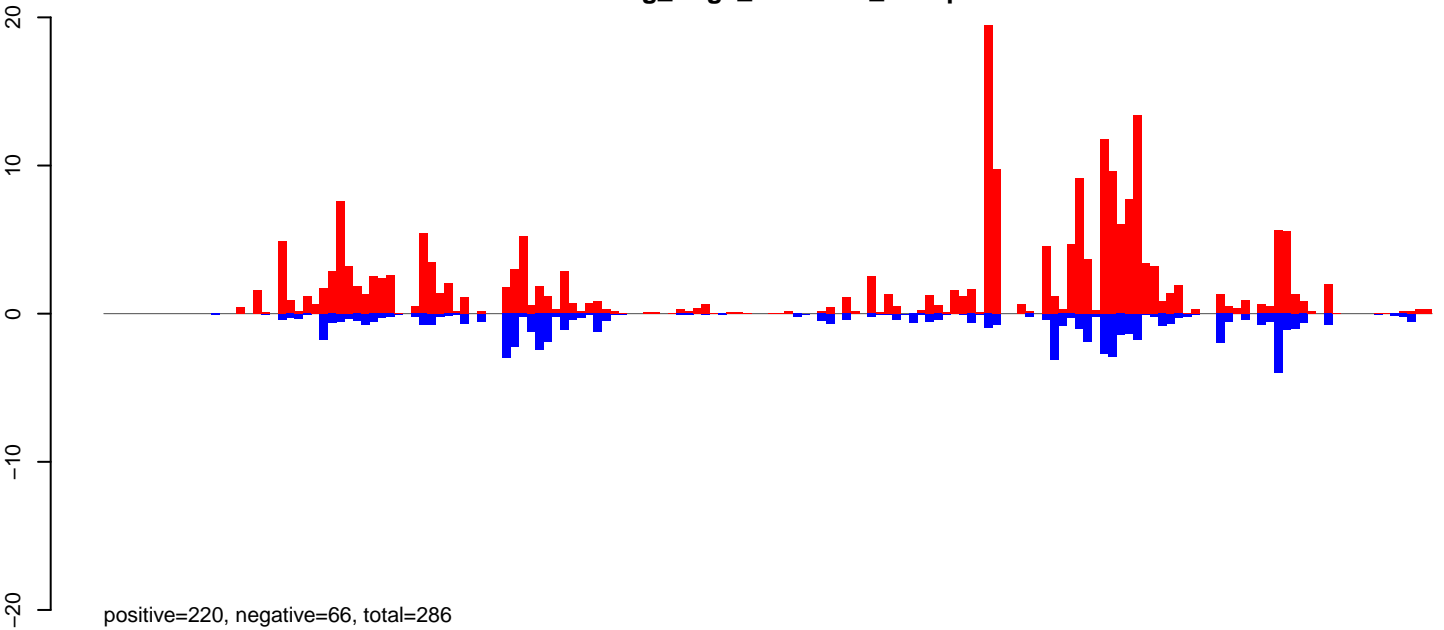
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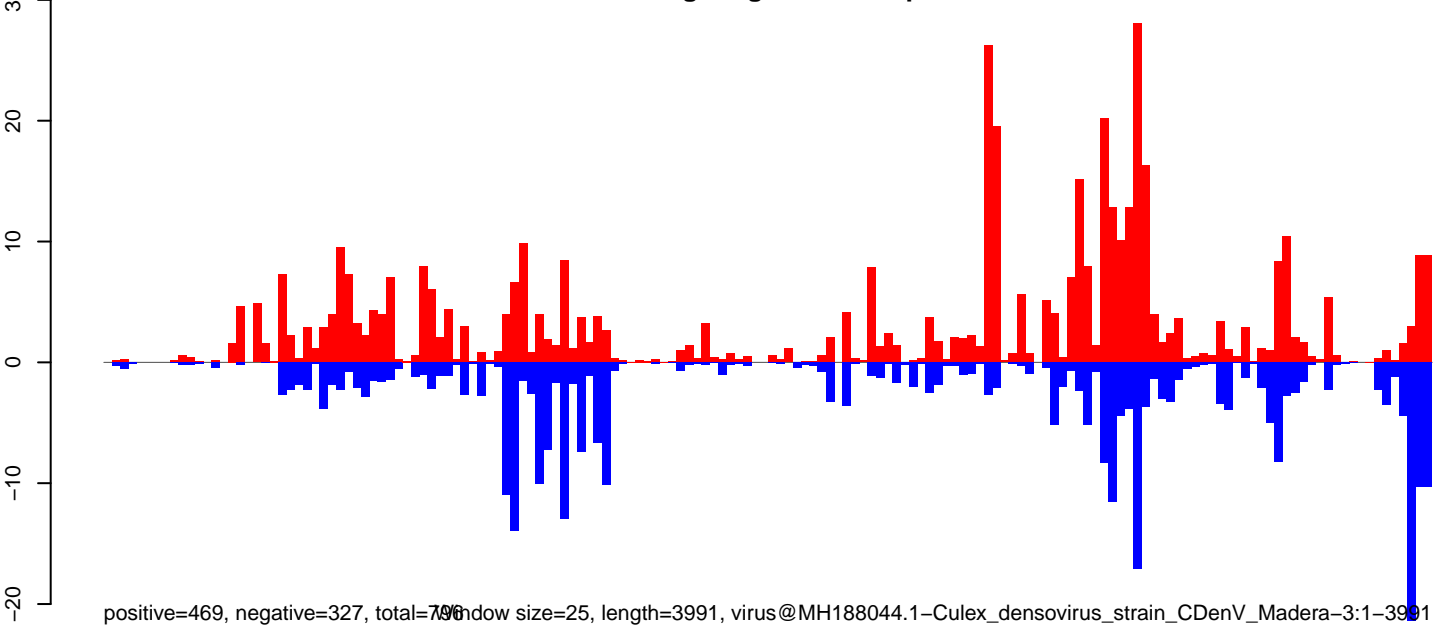
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep



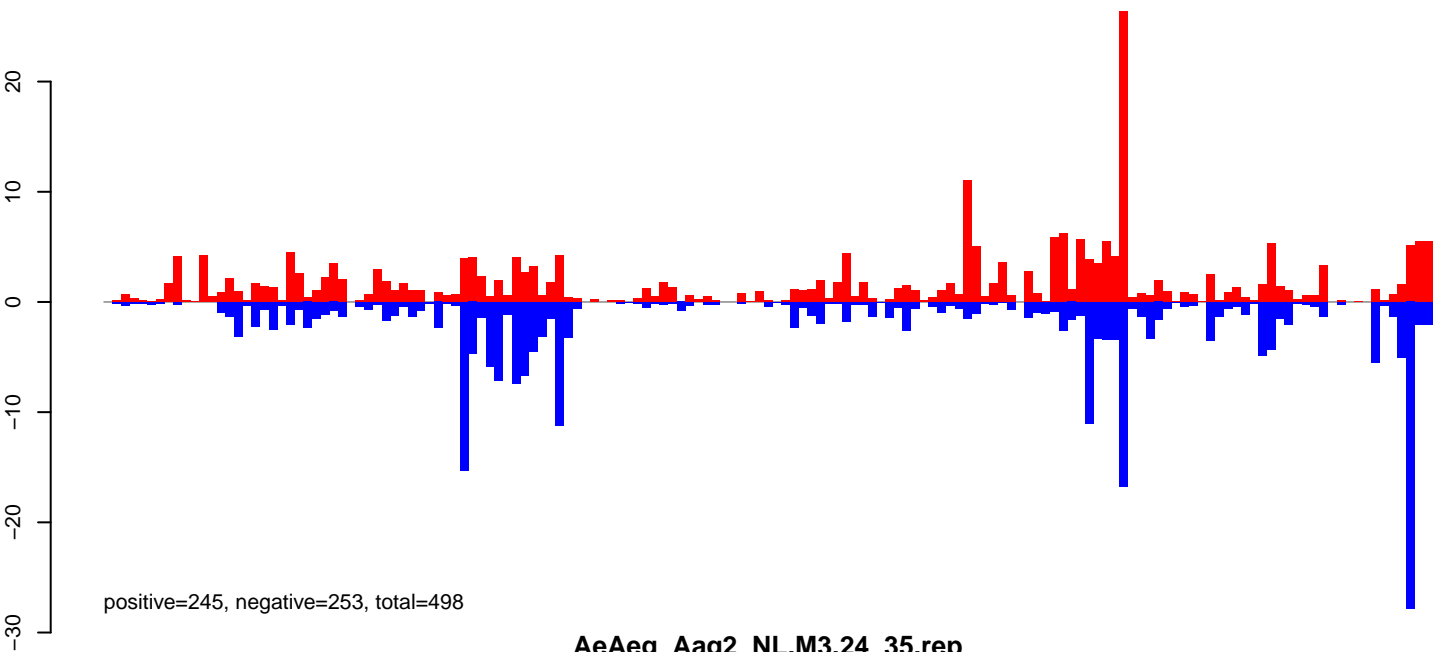
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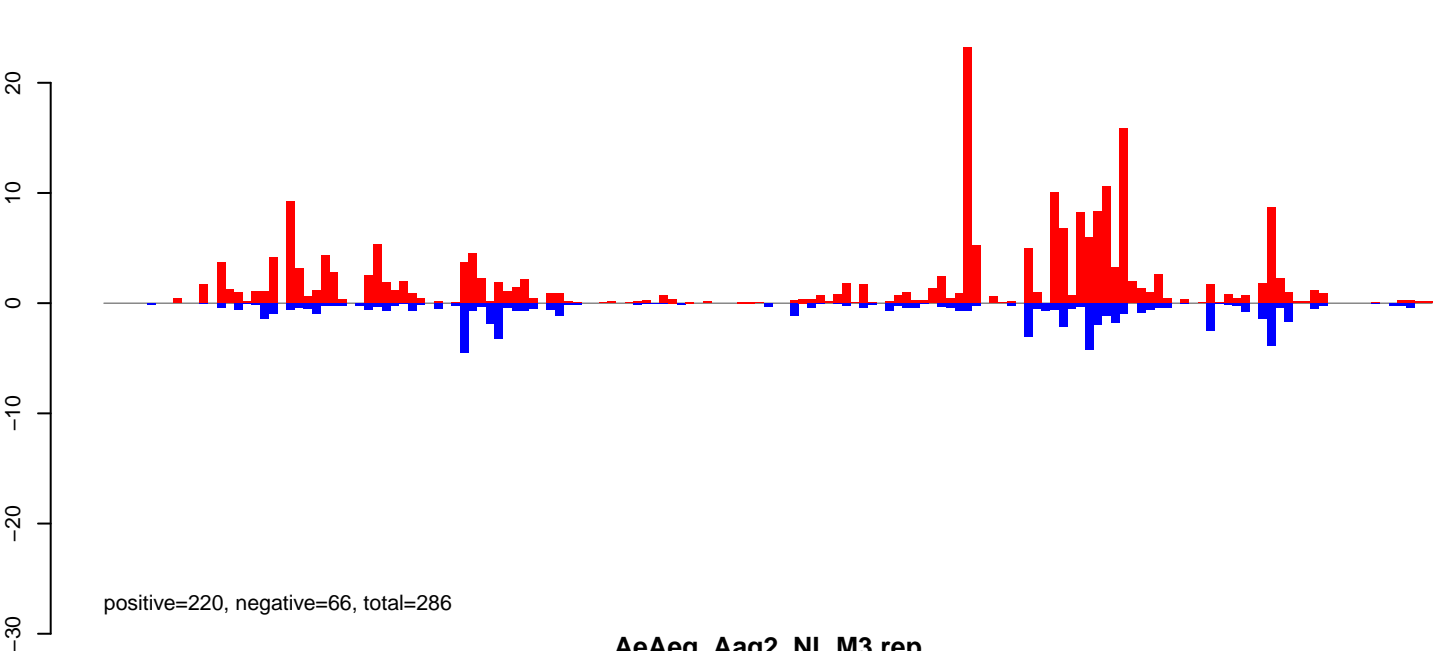
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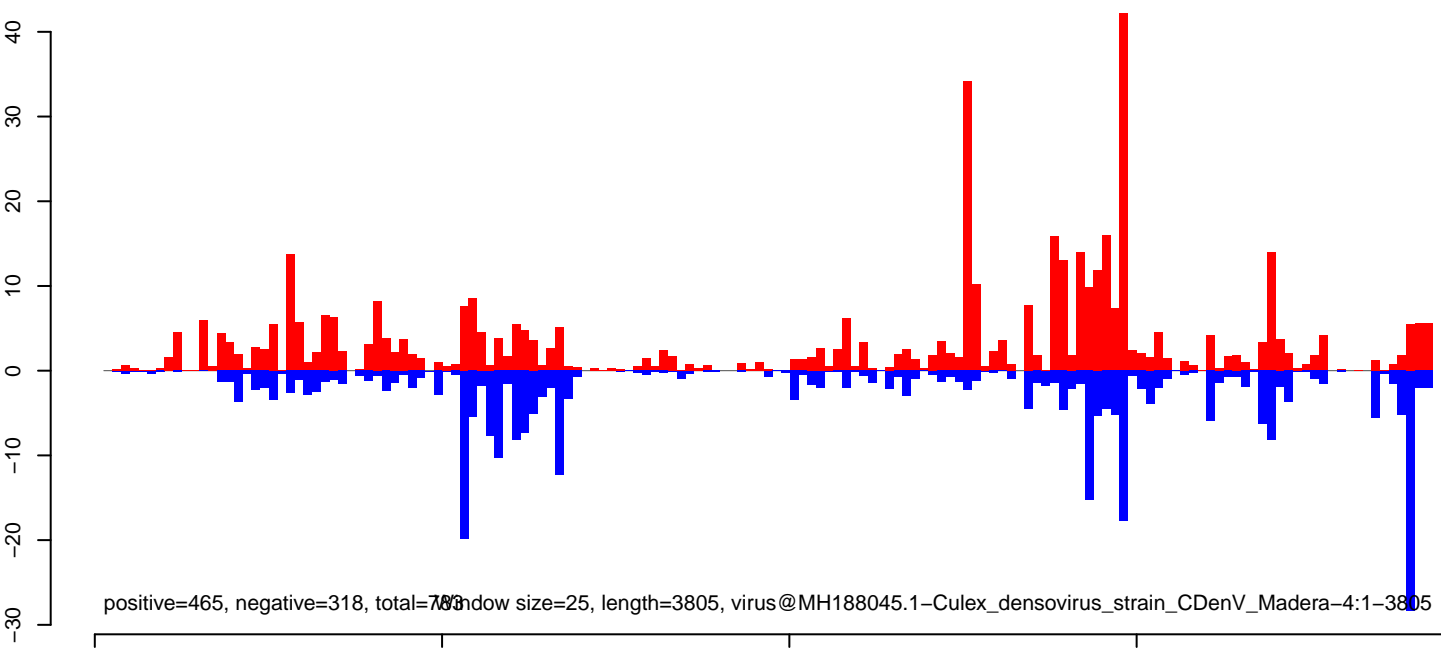
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep

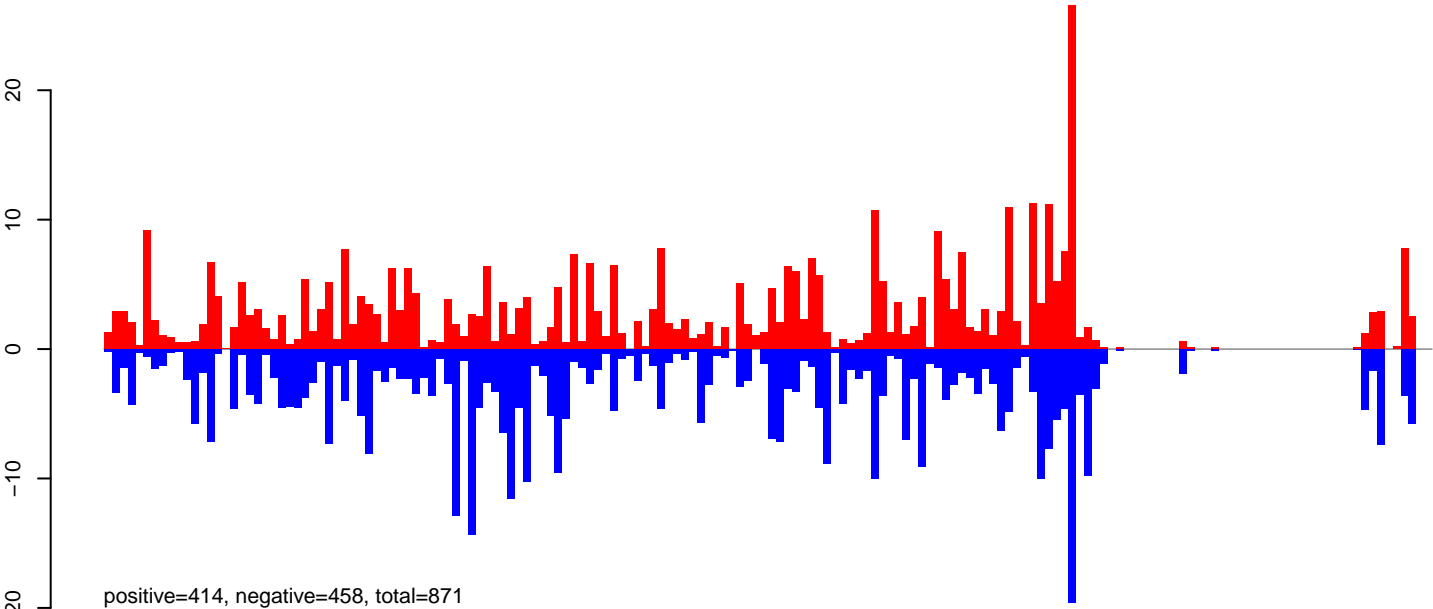


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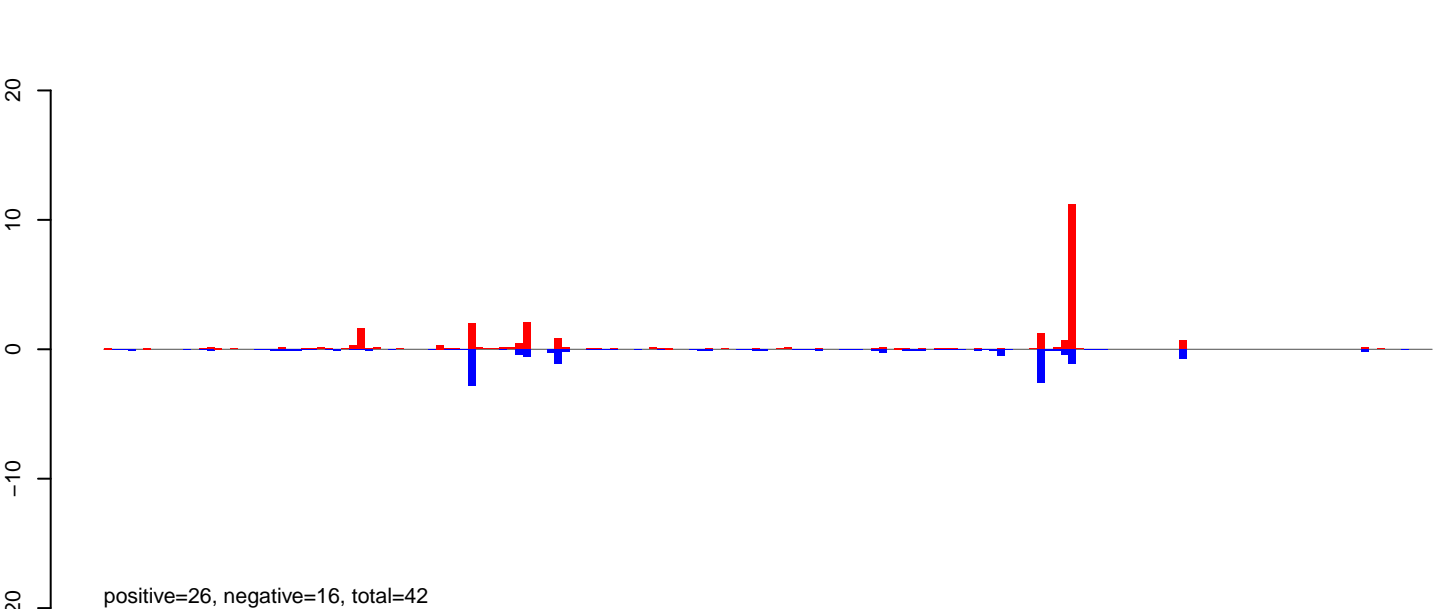


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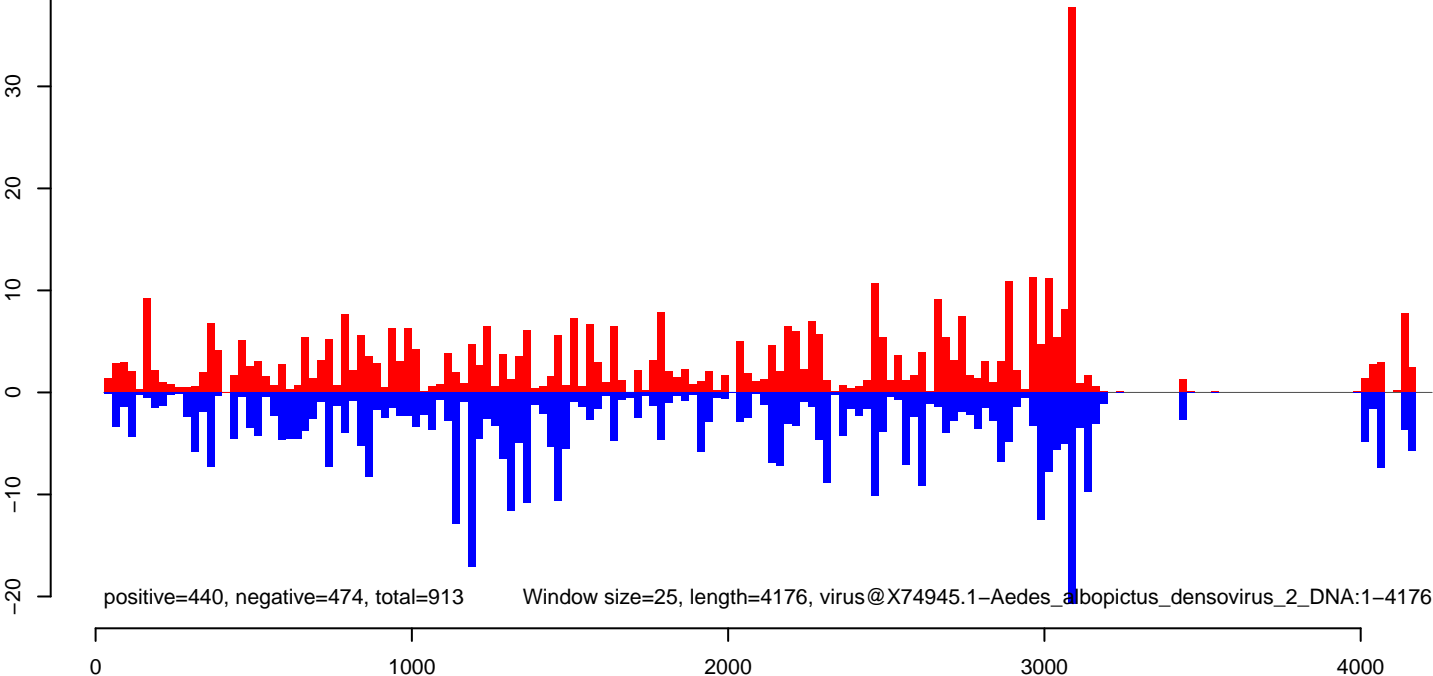
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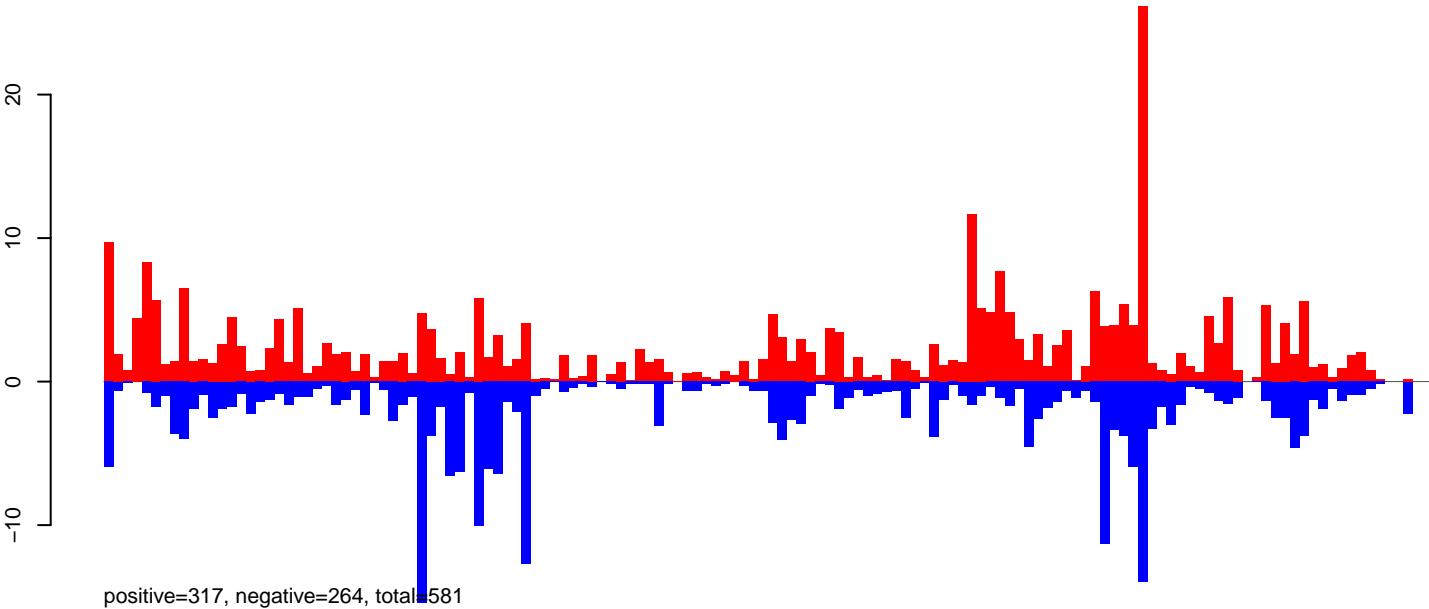
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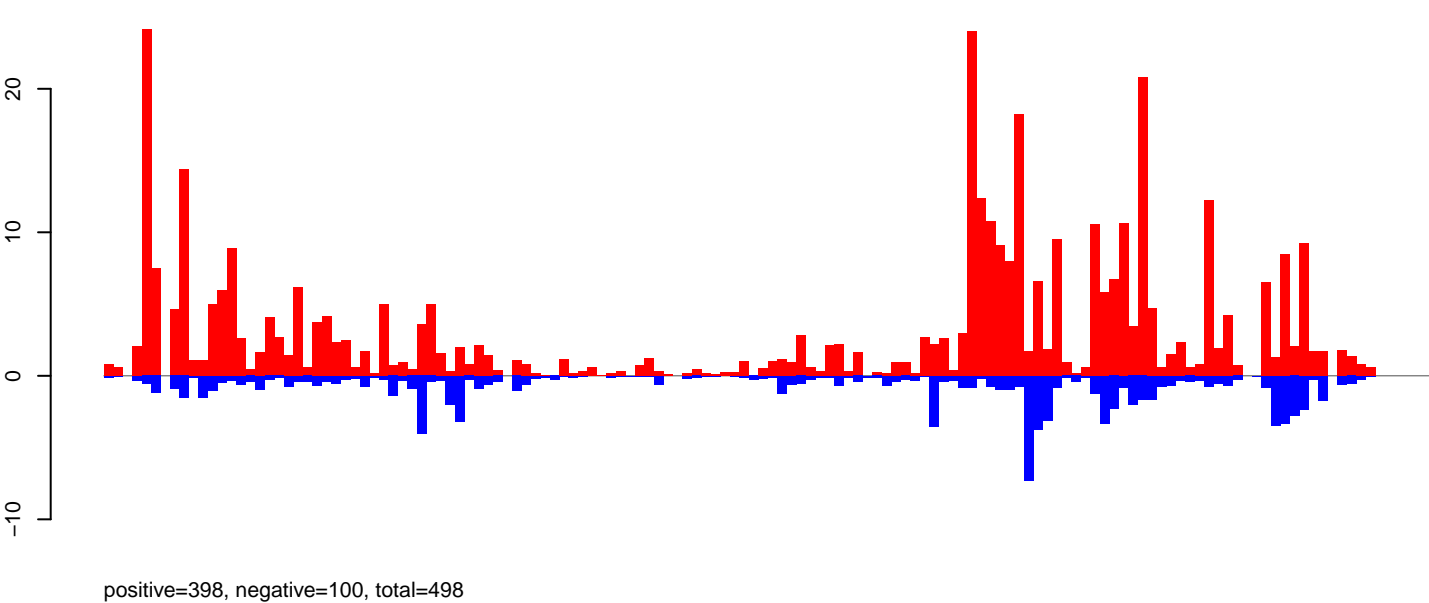
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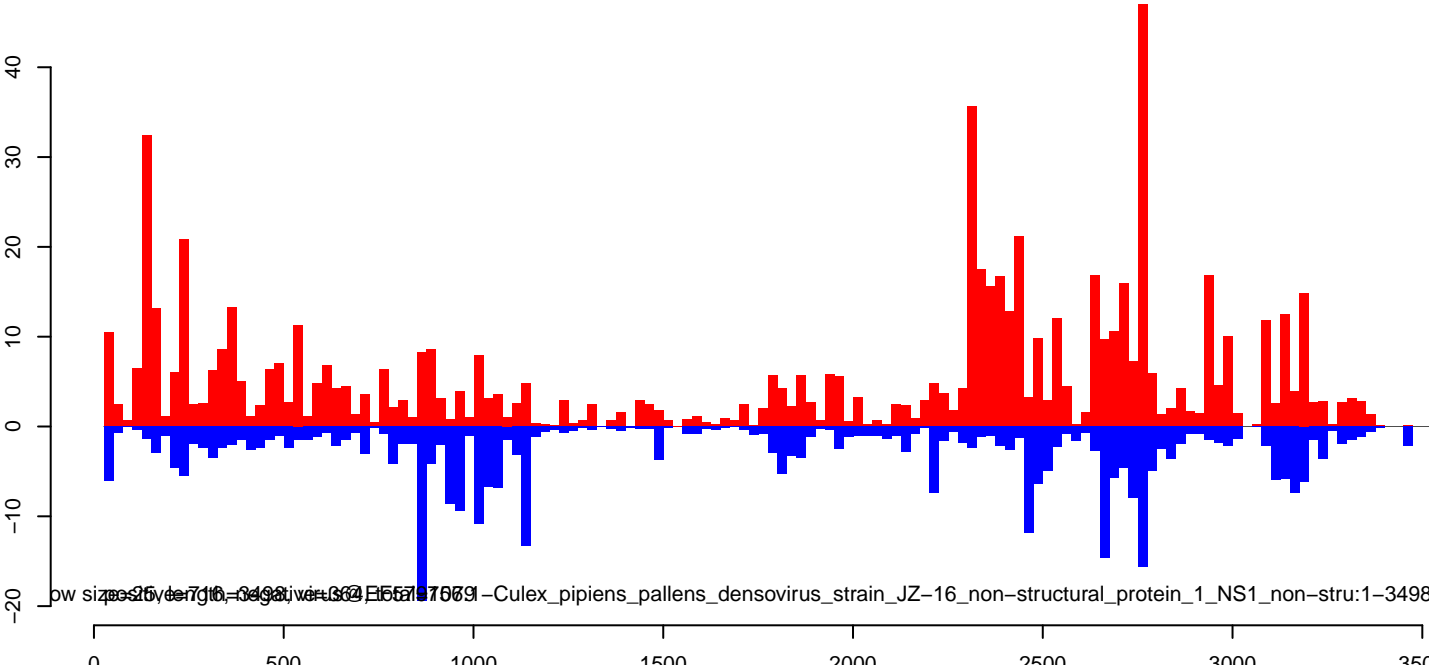
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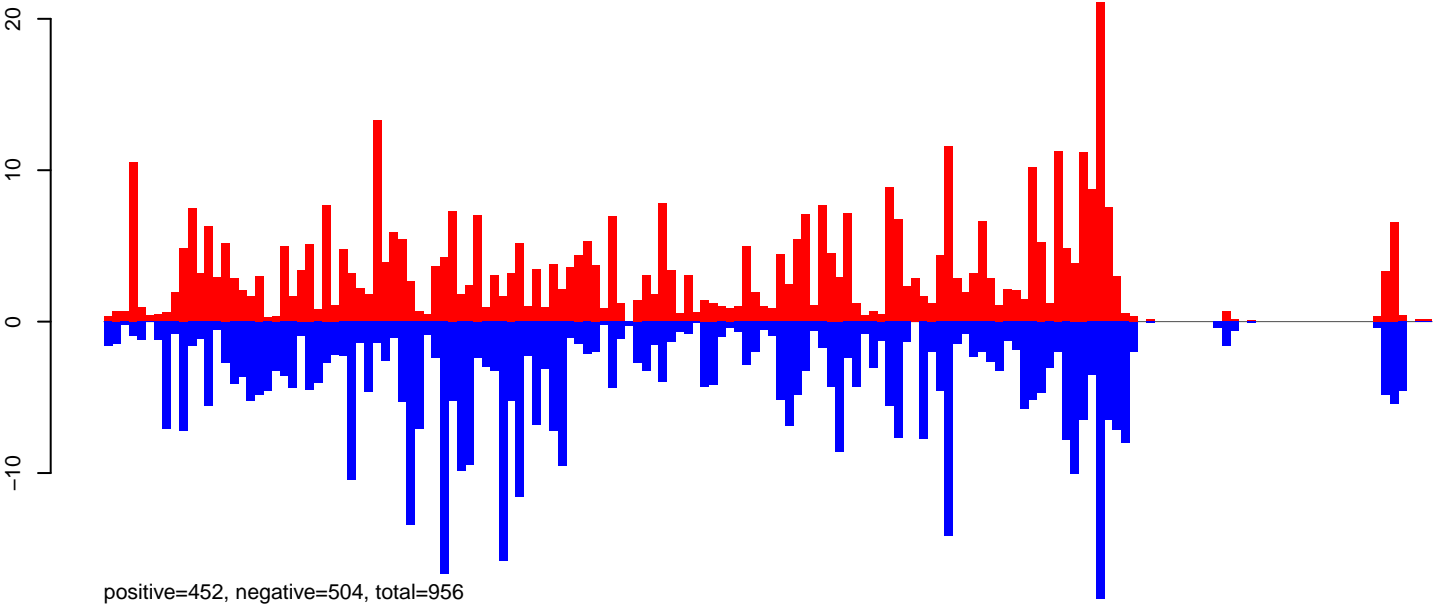
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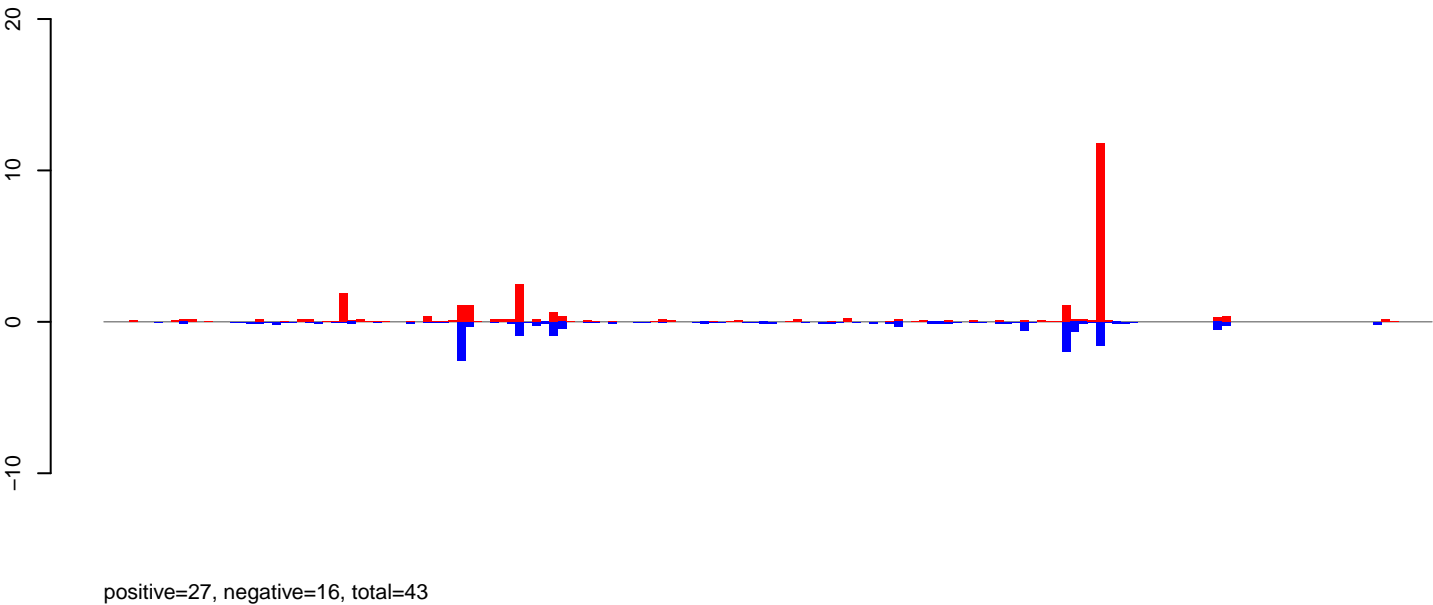
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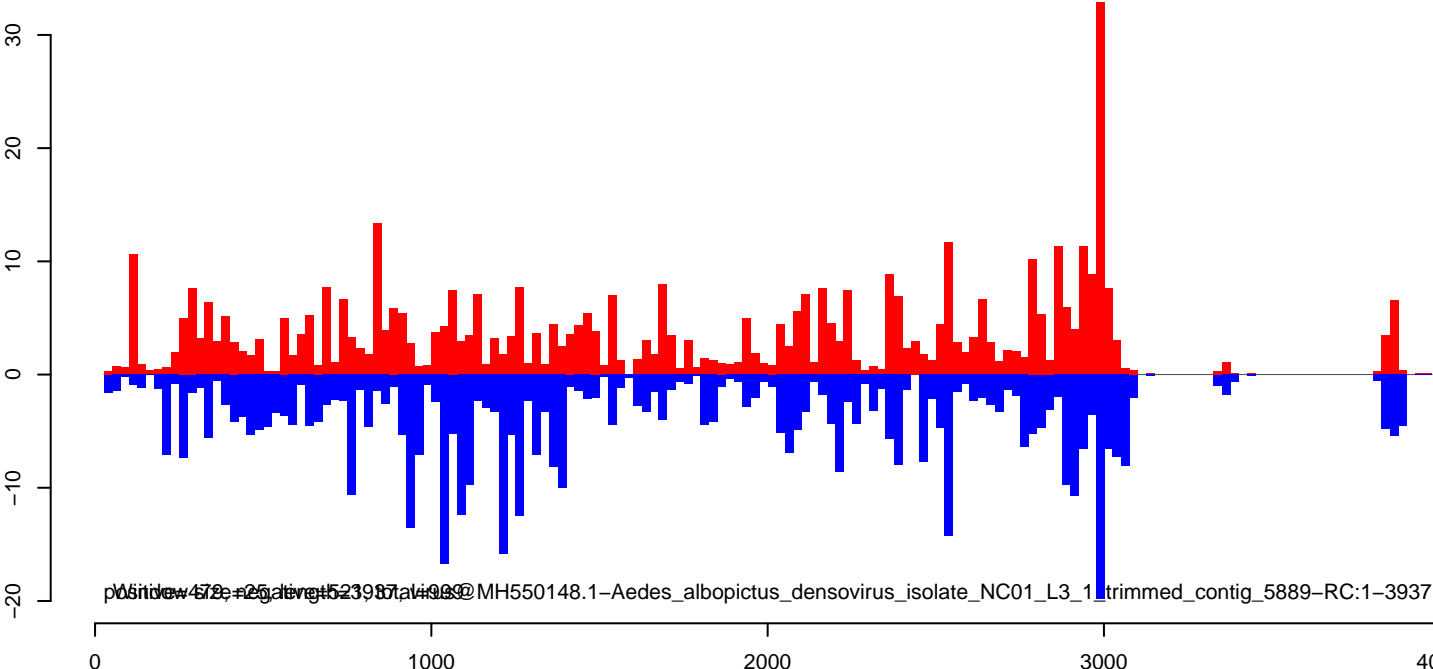
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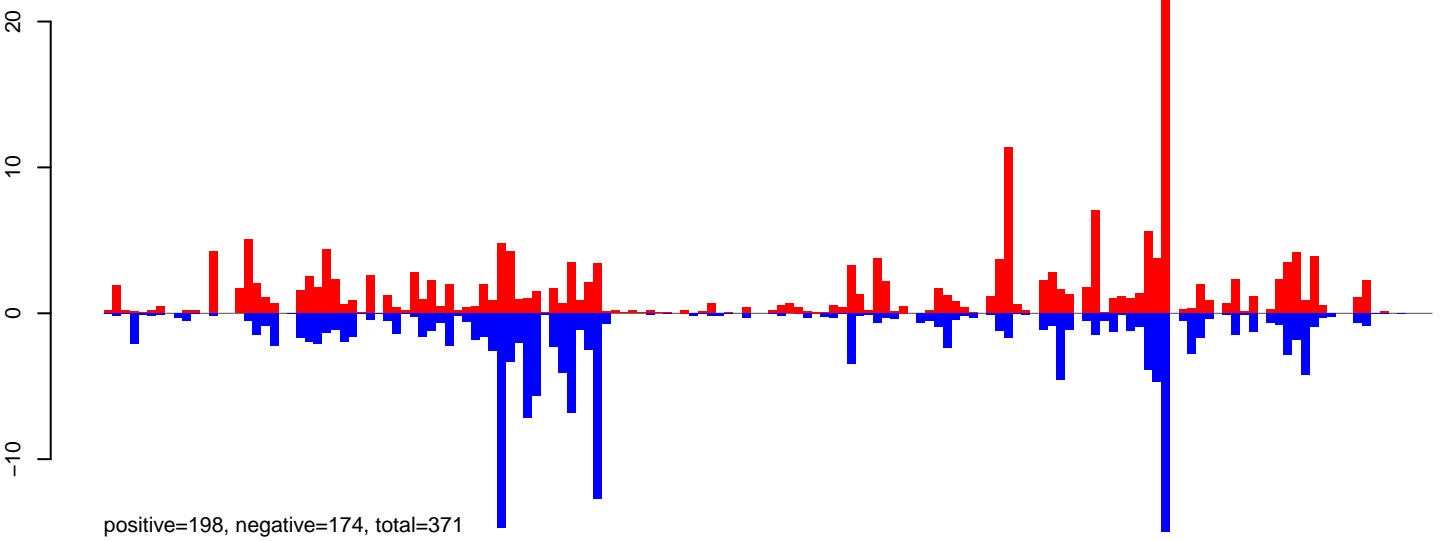
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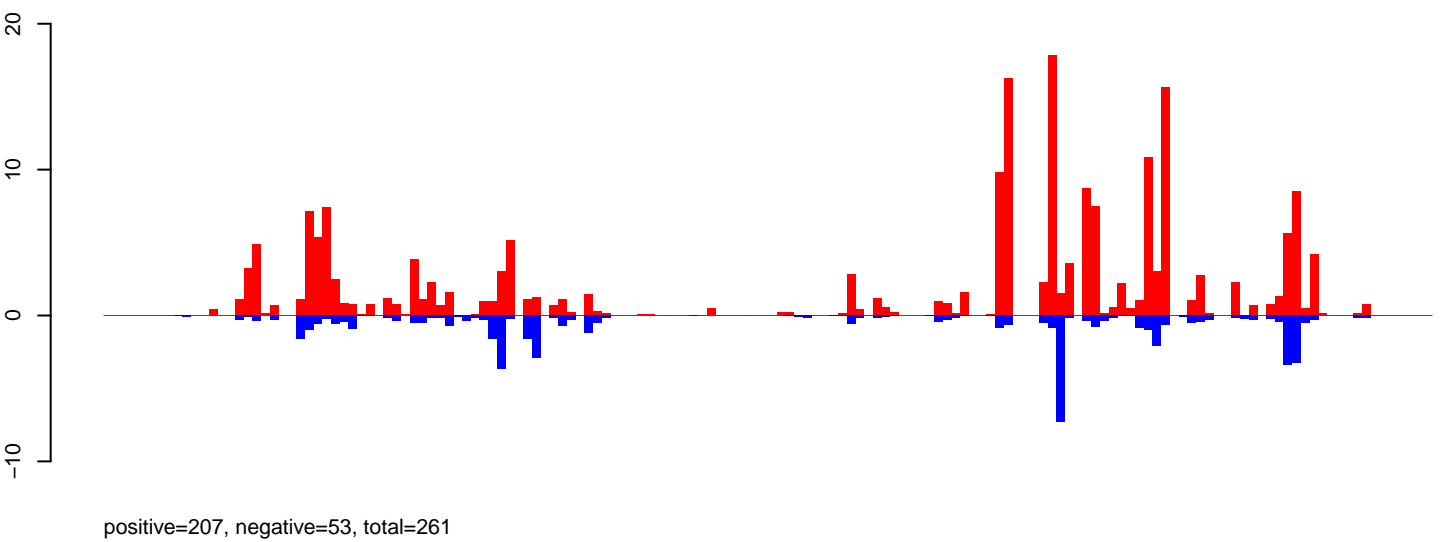
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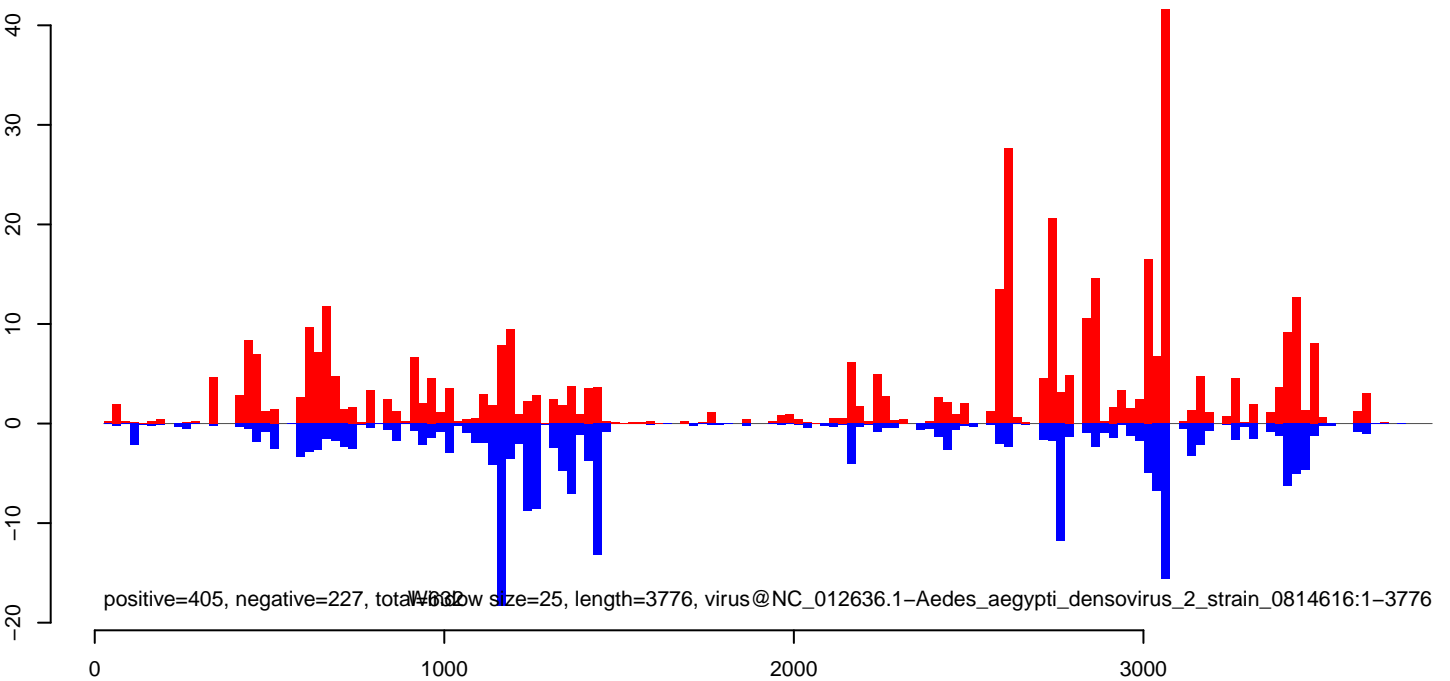
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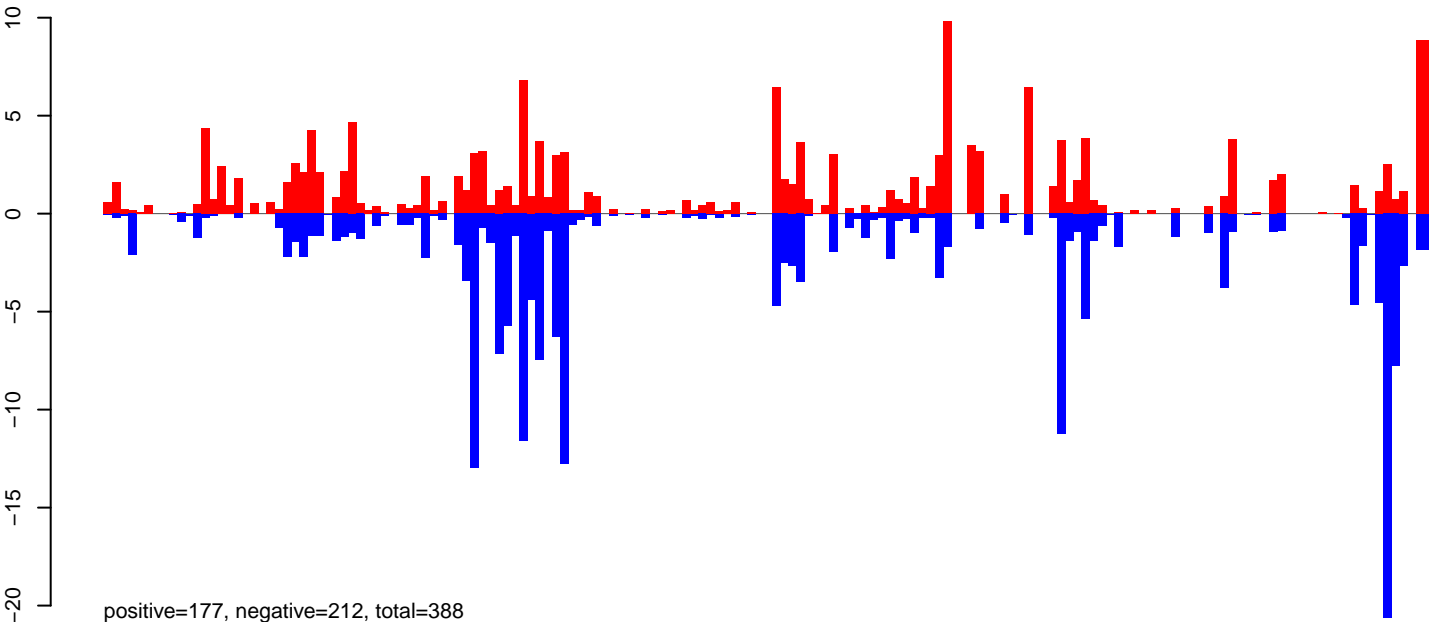
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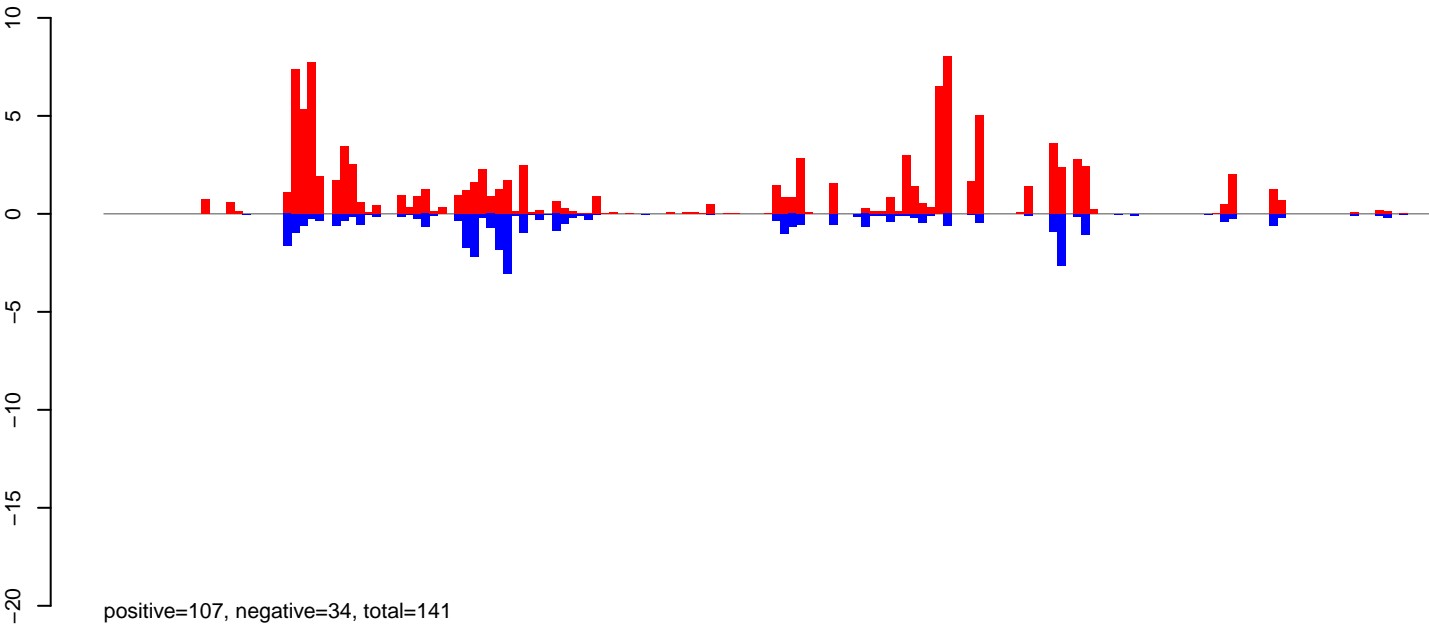
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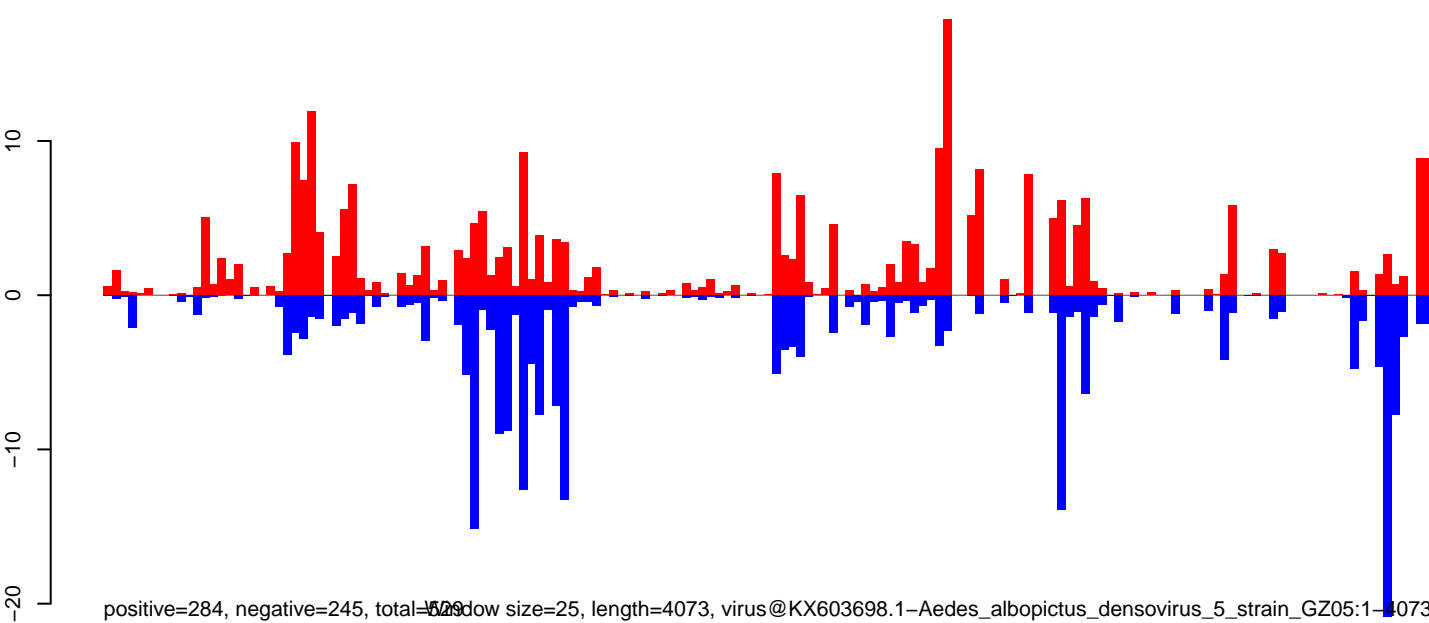
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AeAeg_Aag2_NL.M3.24_35.rep

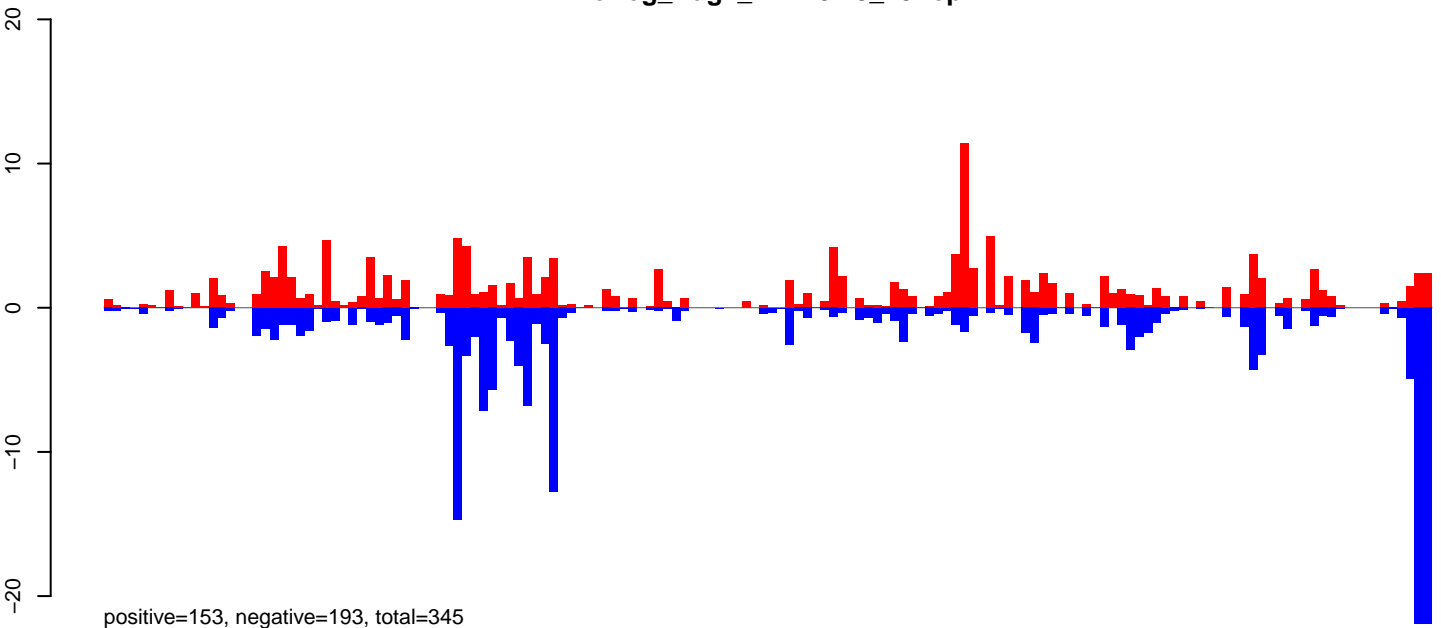


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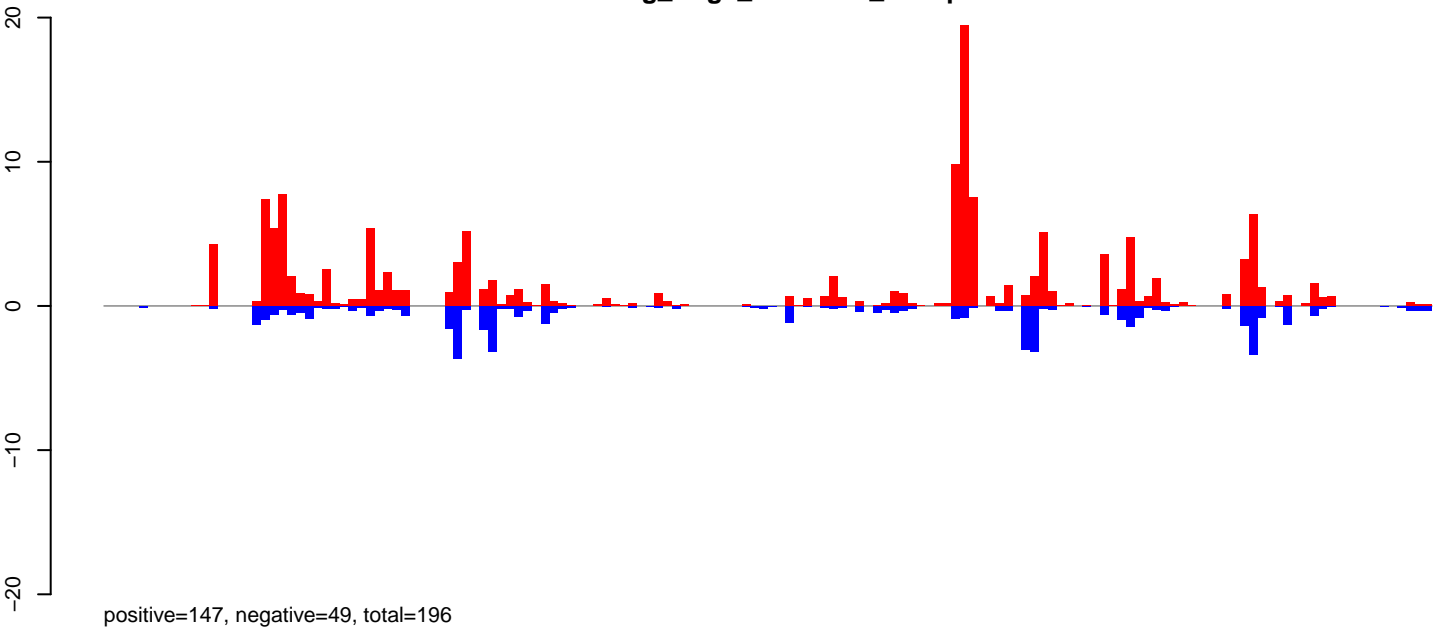


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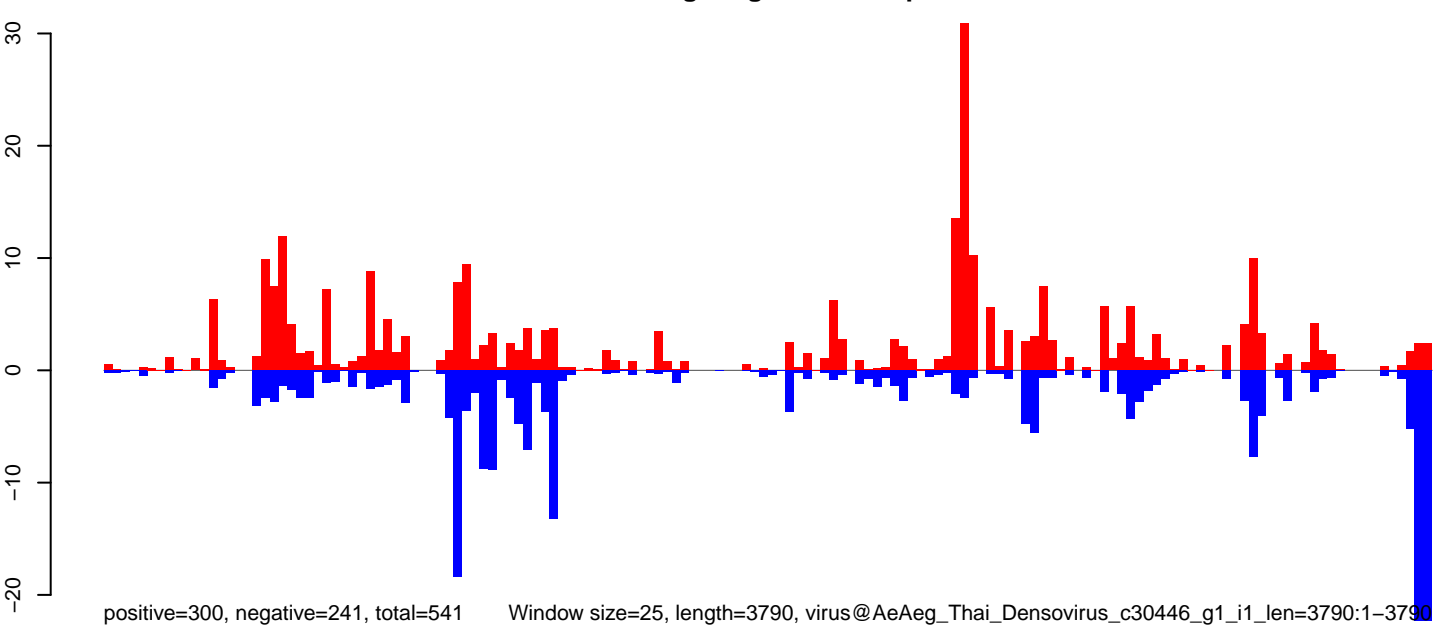
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AeAeg_Aag2_NL.M3.24_35.rep



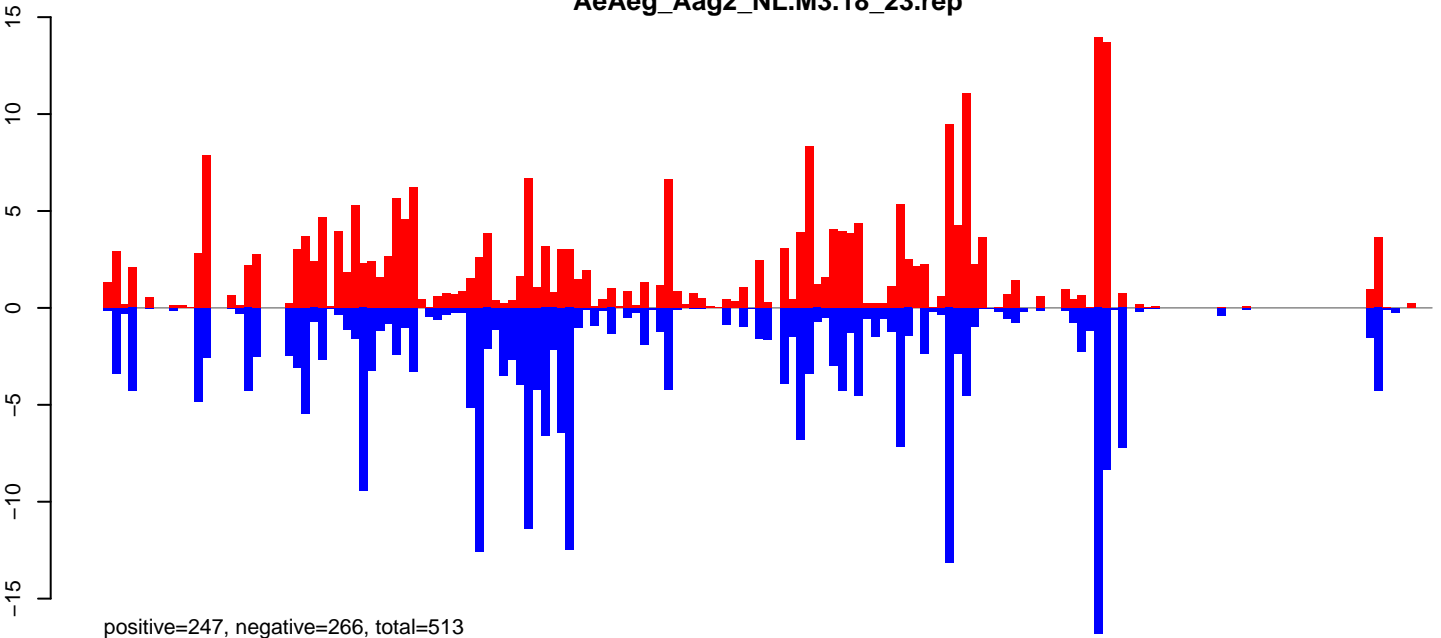
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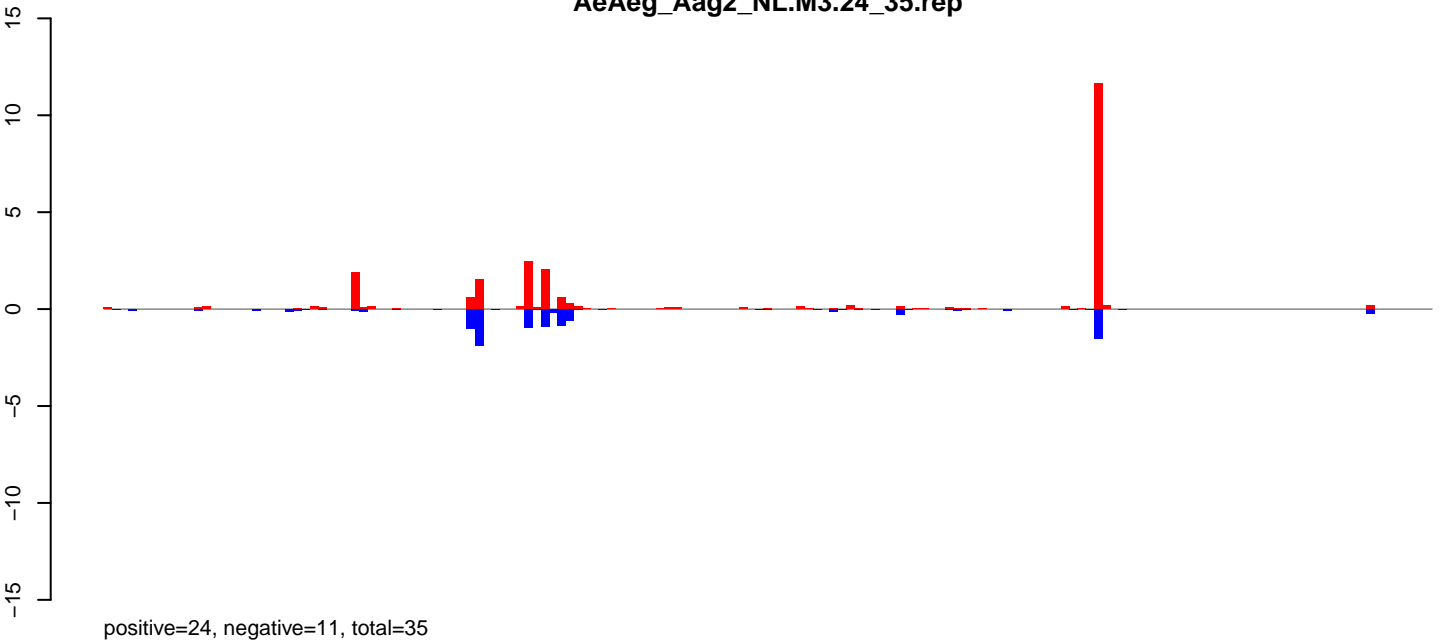
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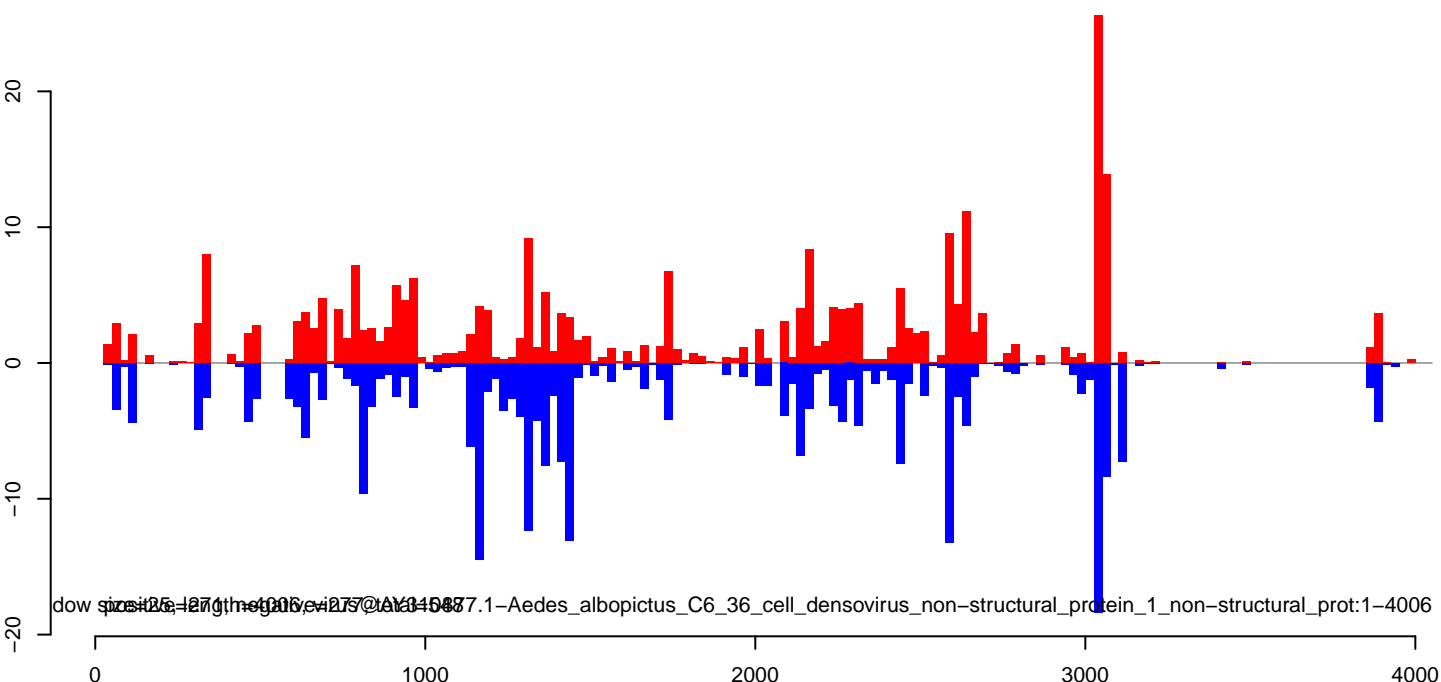
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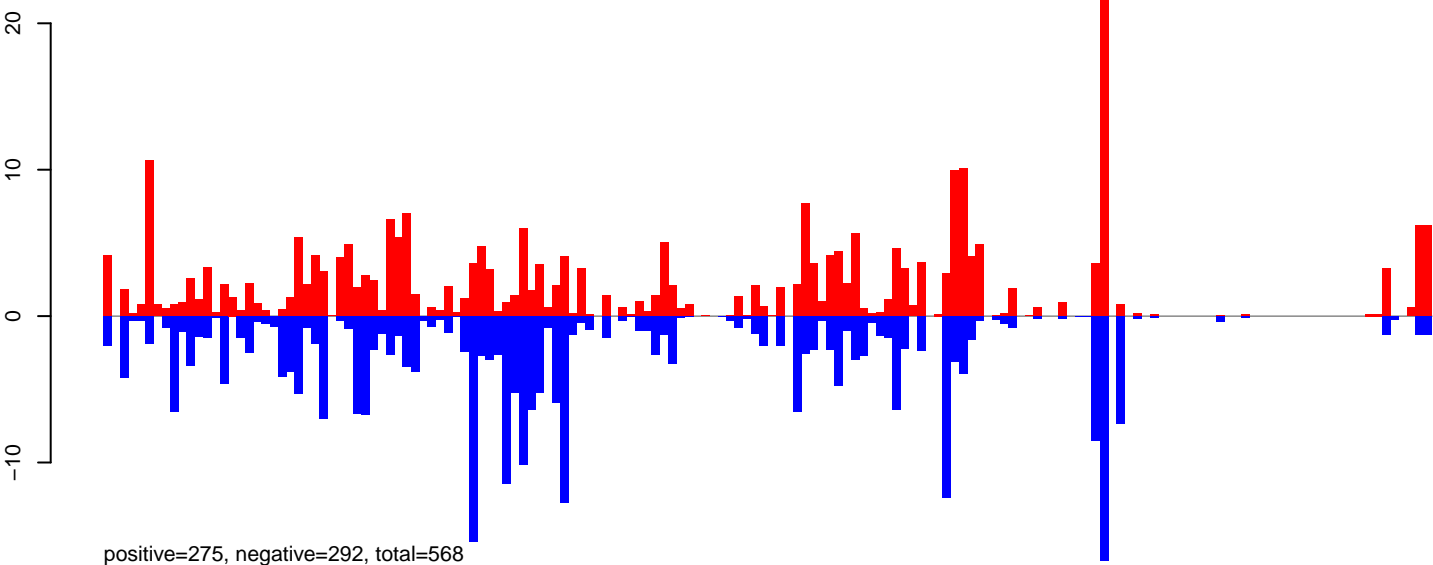
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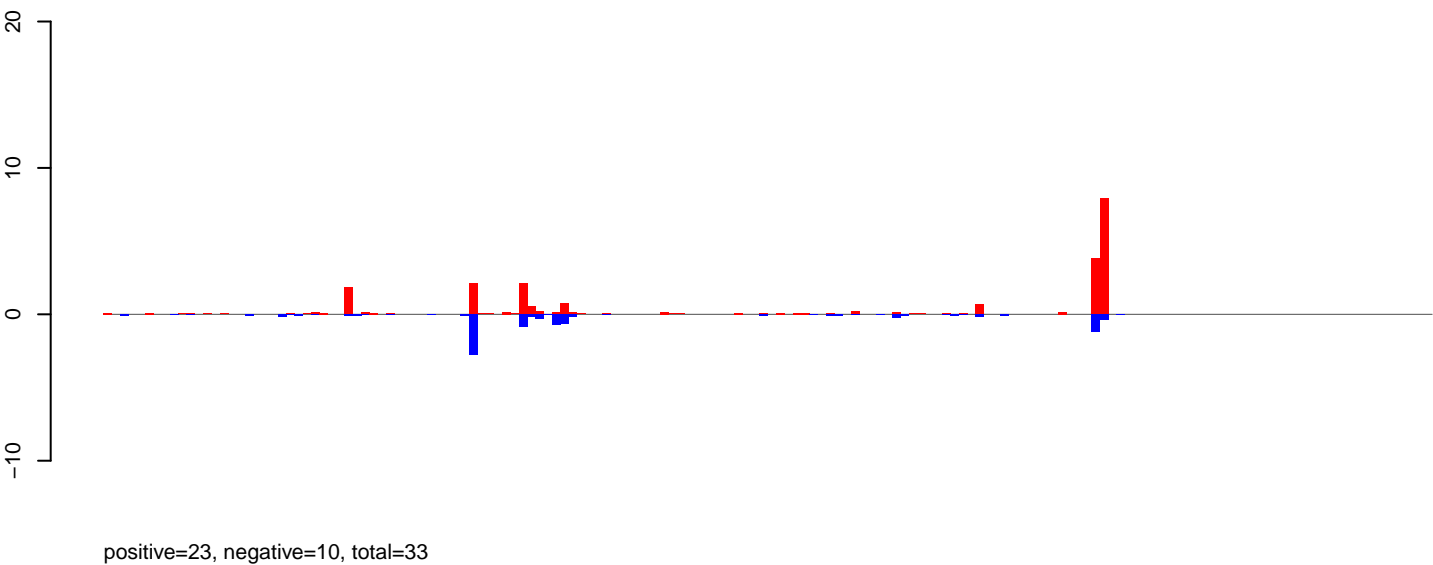
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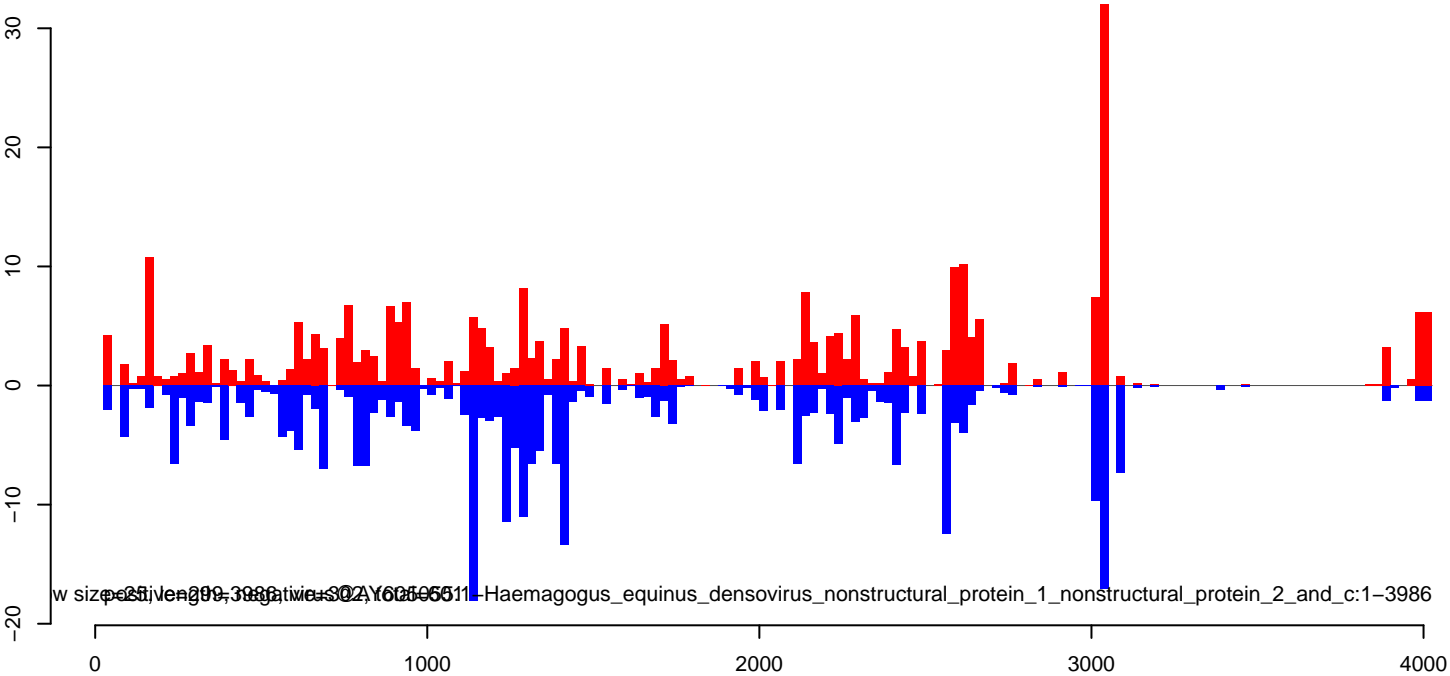
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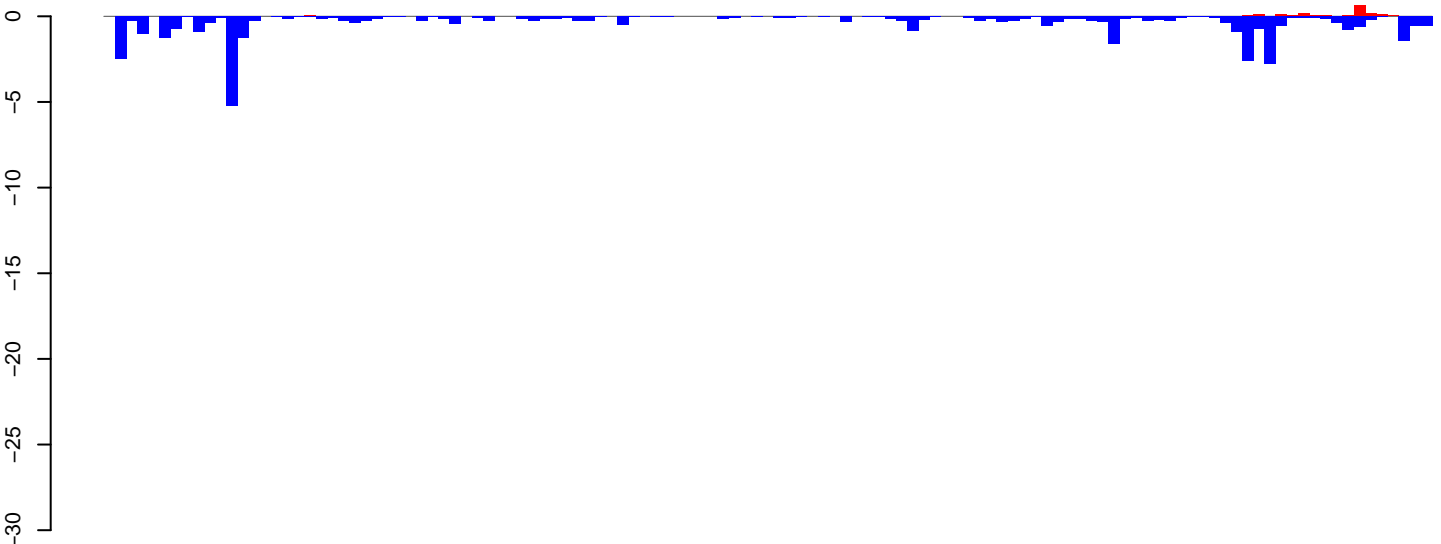
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AeAeg_Aag2_NL.M3.rep

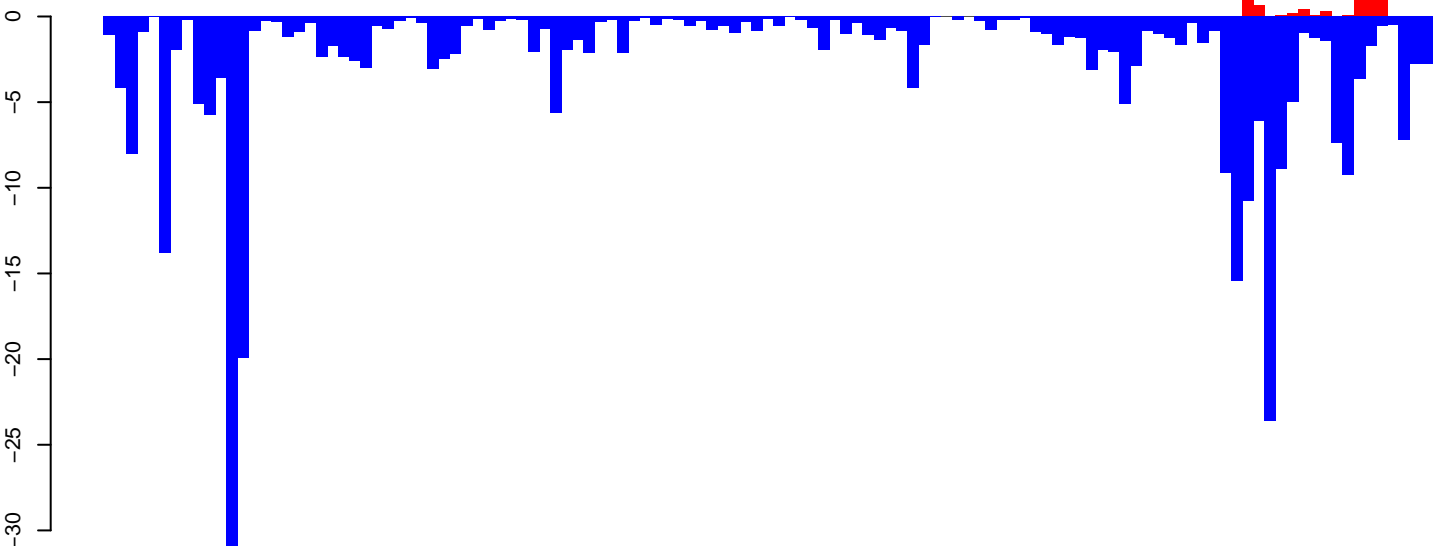


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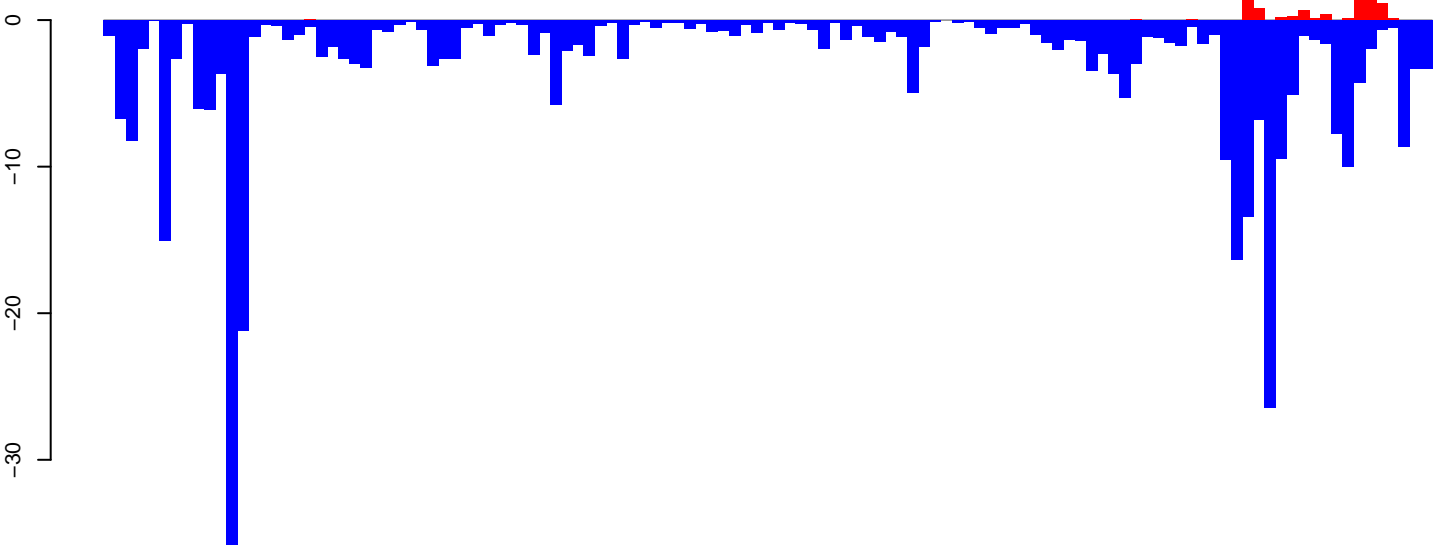
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AeAeg_Aag2_NL.M3.24_35.rep



positive=8, negative=311, total=319

AeAeg_Aag2_NL.M3.rep

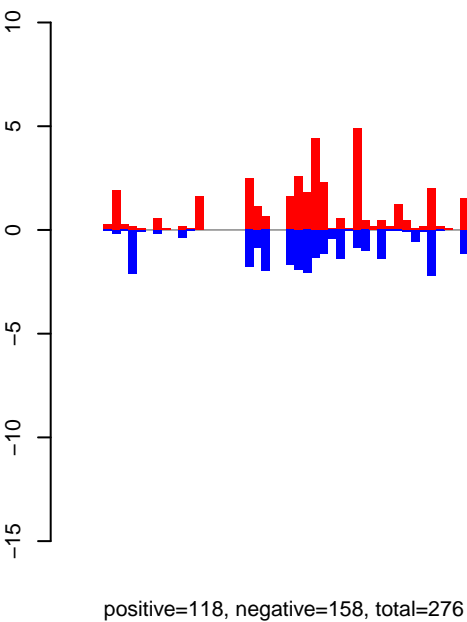


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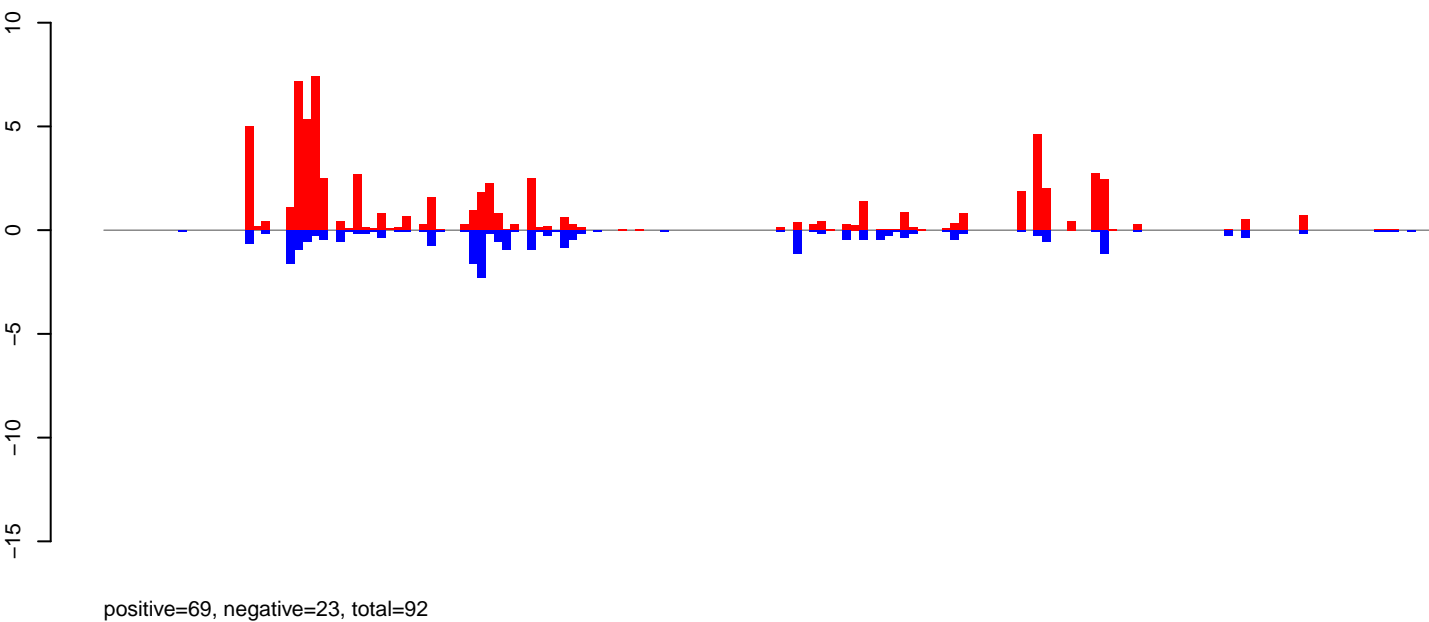
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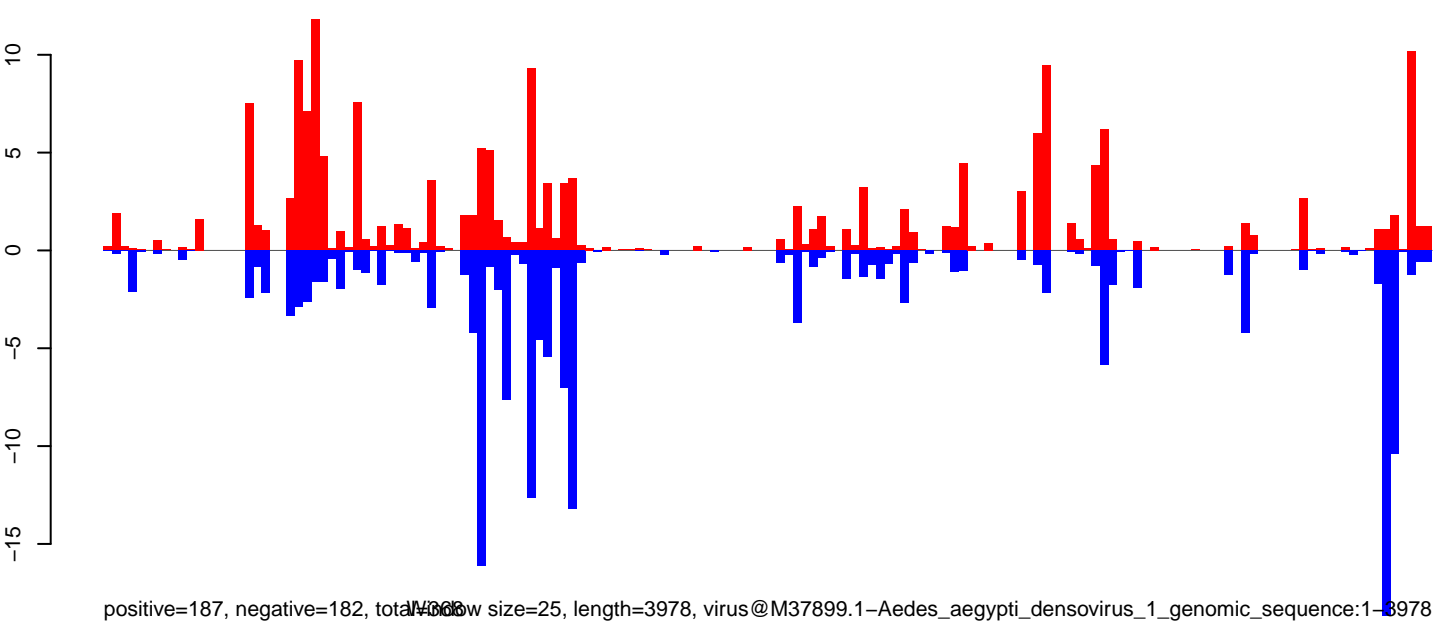
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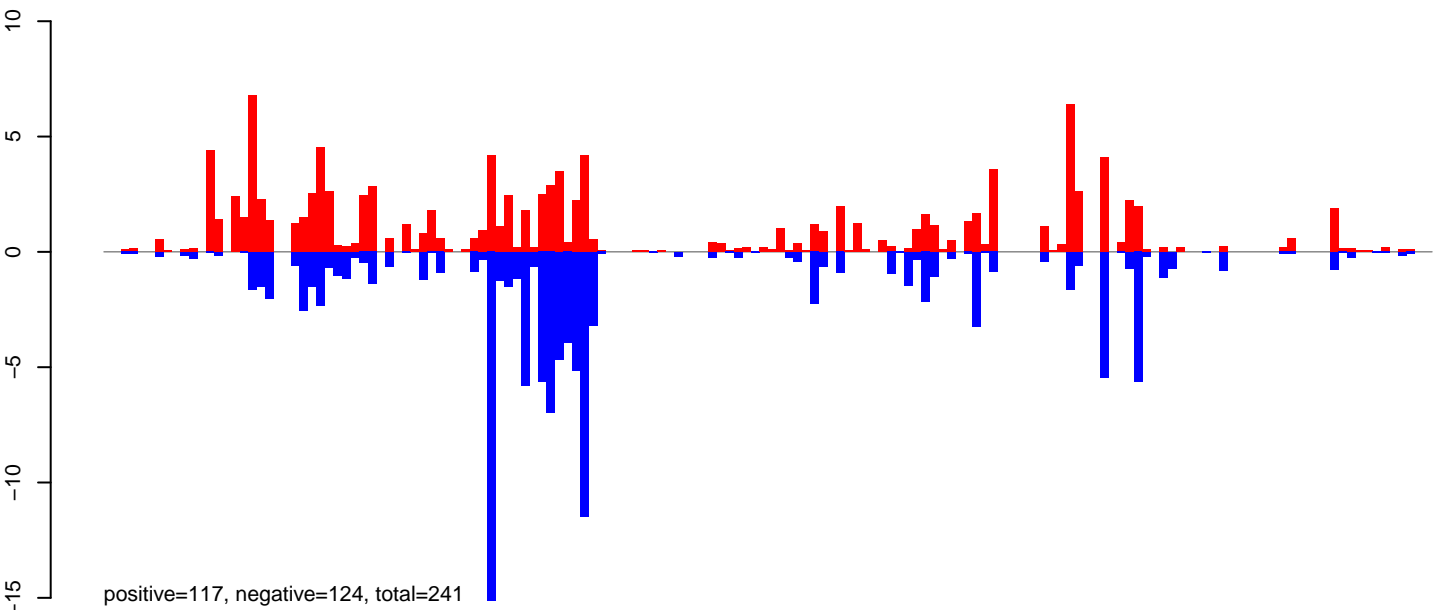
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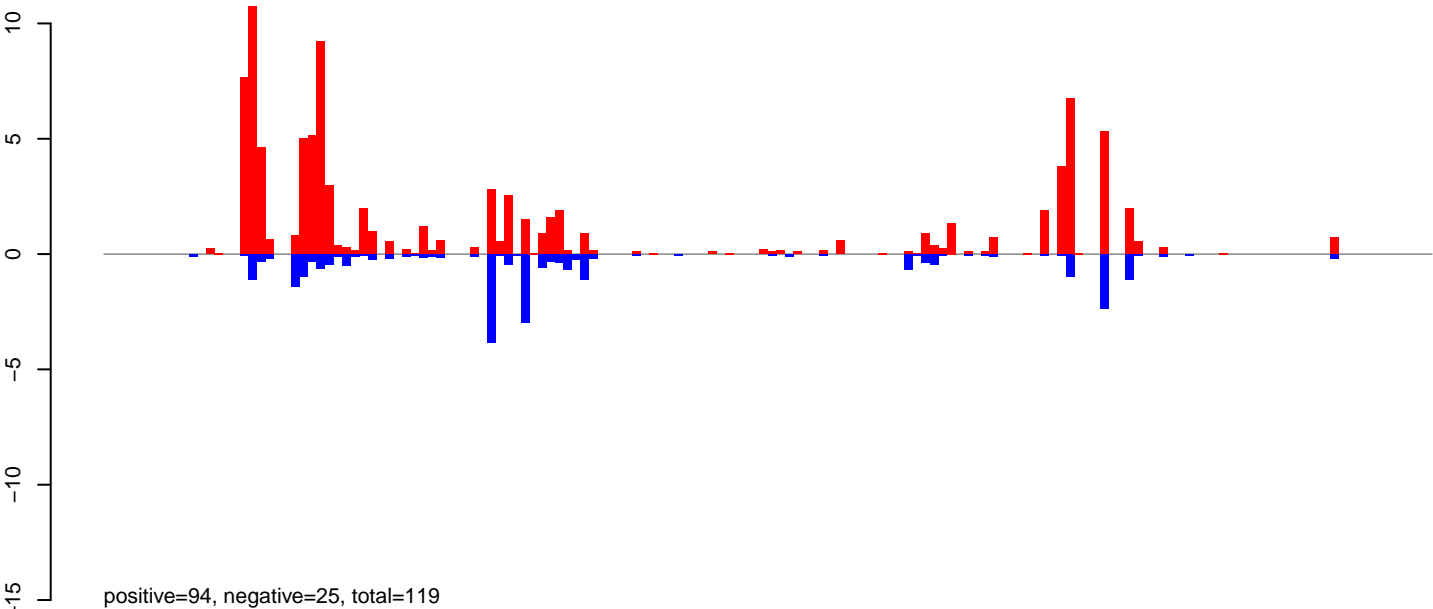
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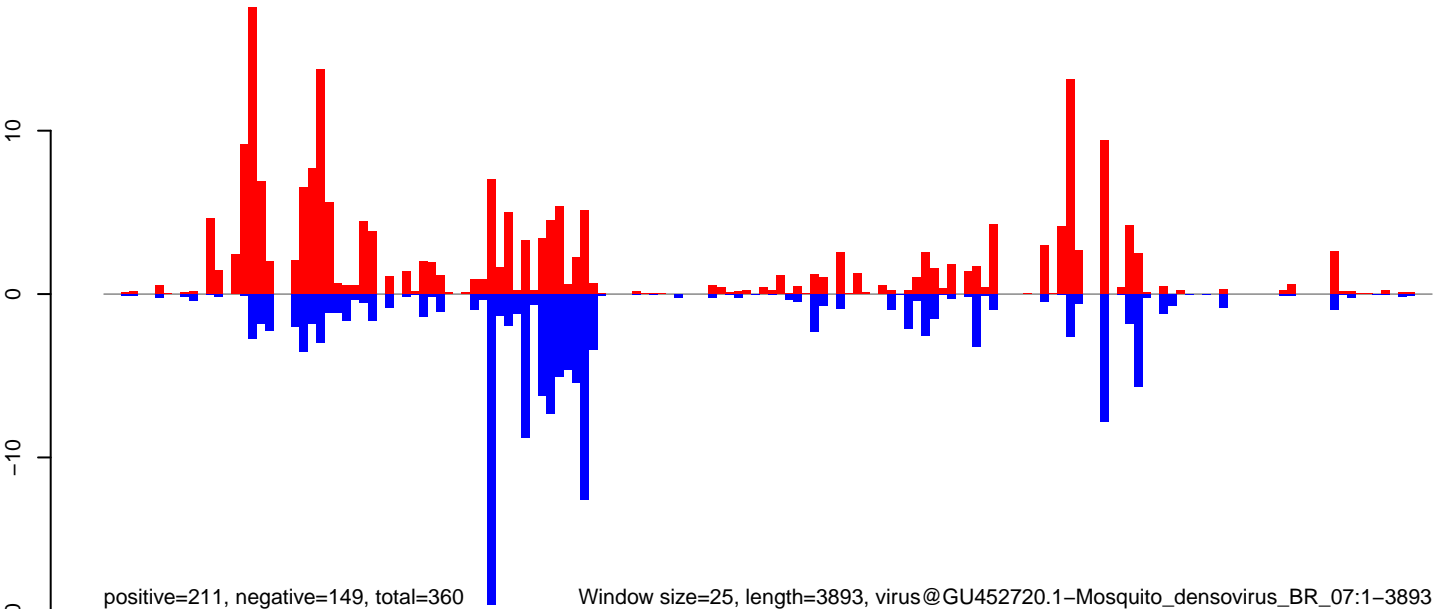
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AeAeg_Aag2_NL.M3.24_35.rep



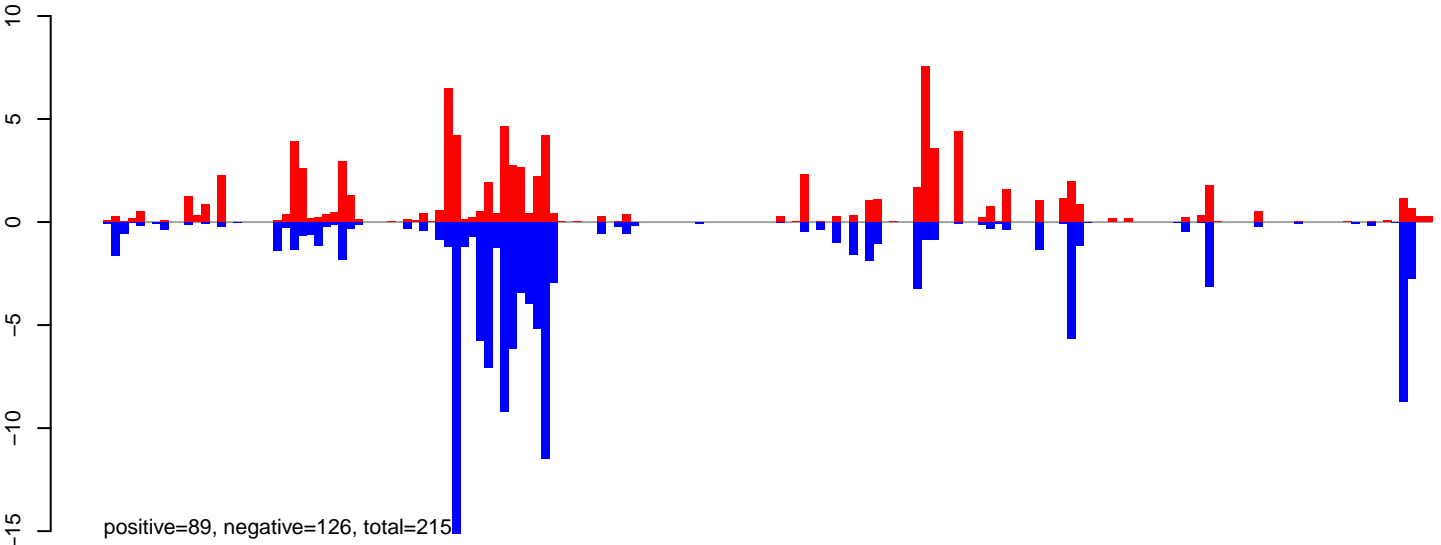
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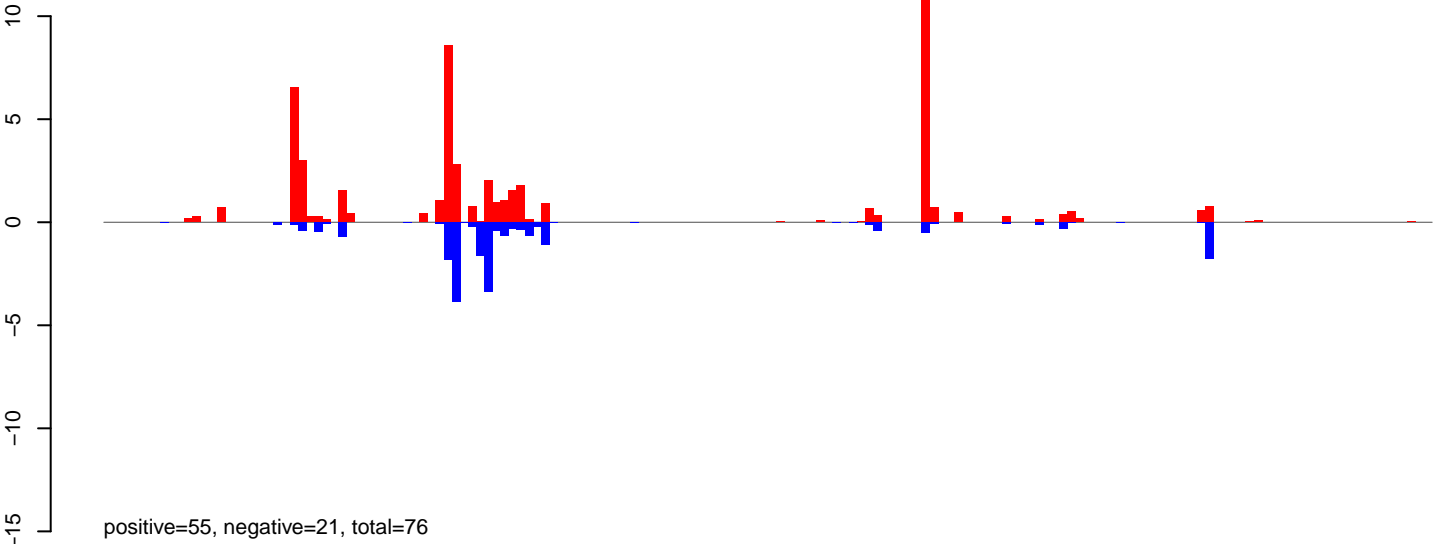
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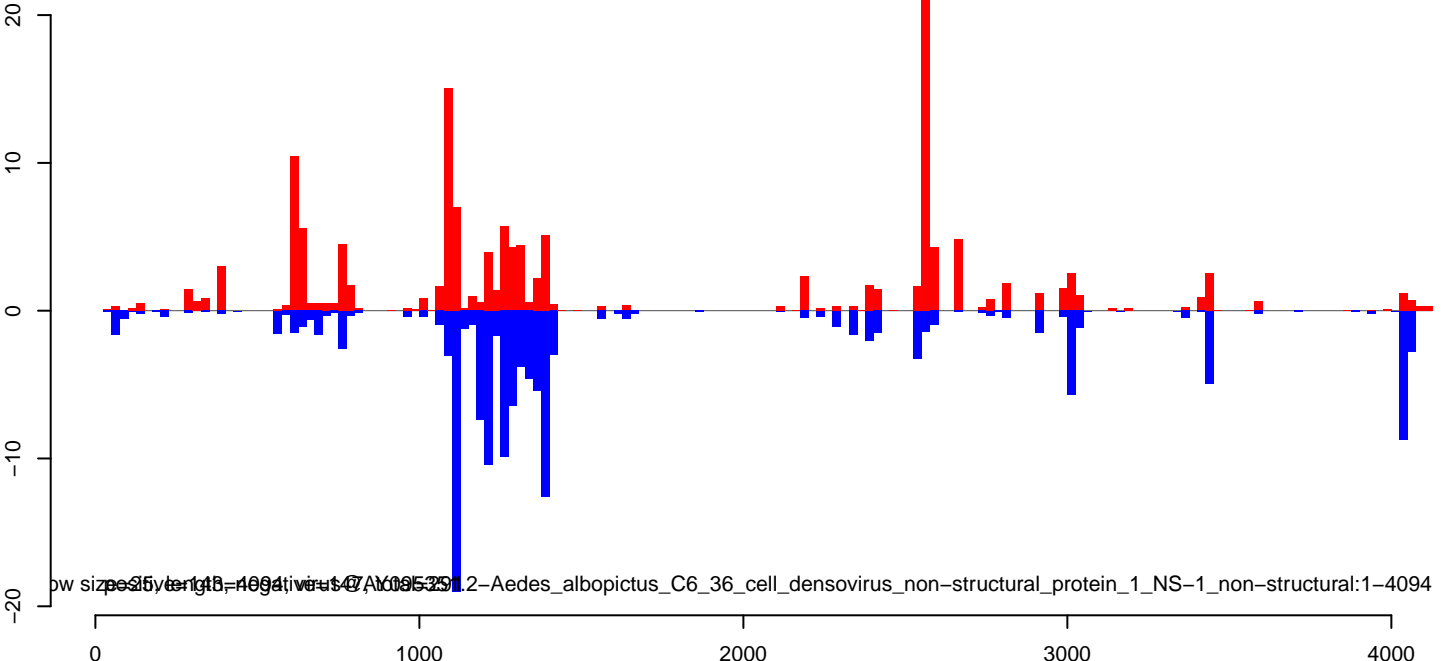
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AeAeg_Aag2_NL.M3.24_35.rep

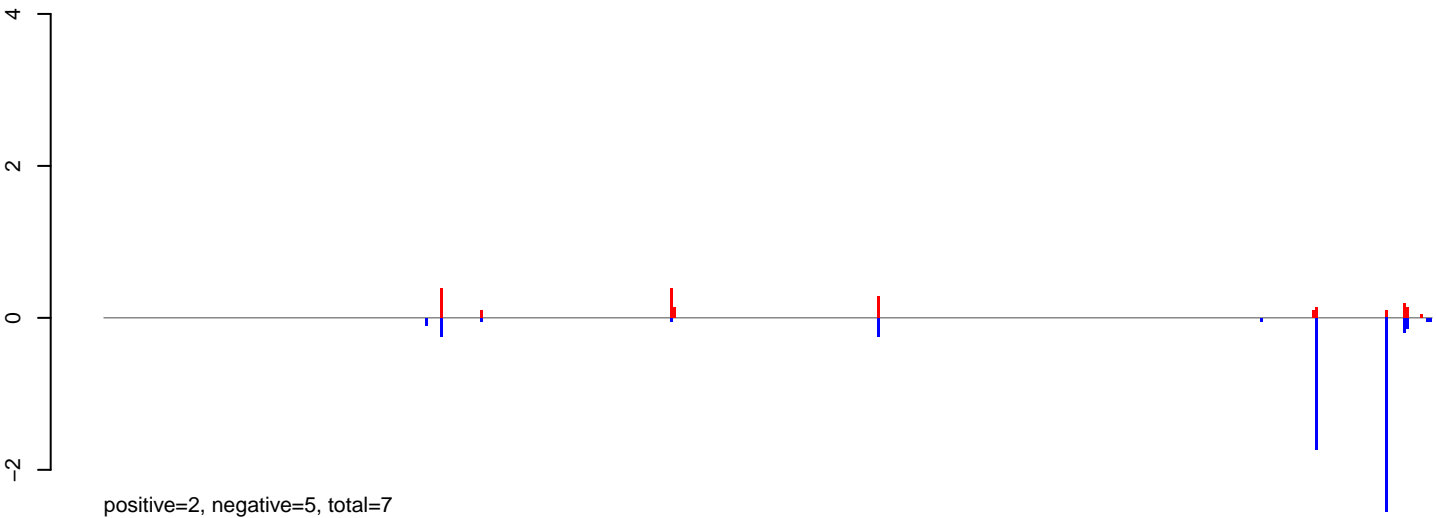


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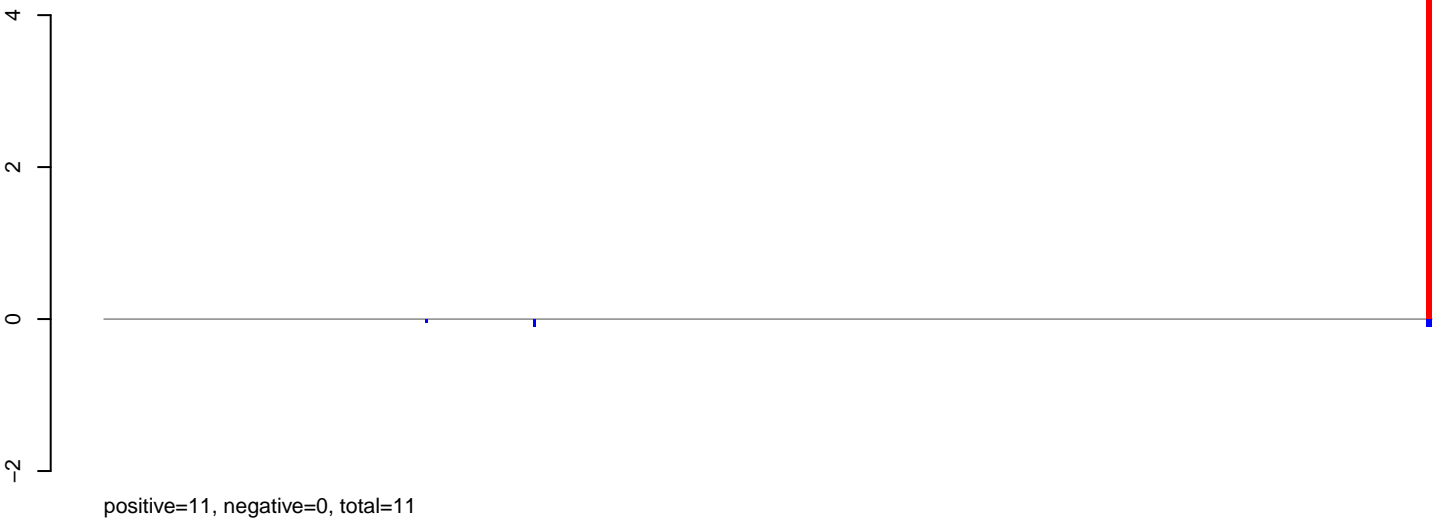


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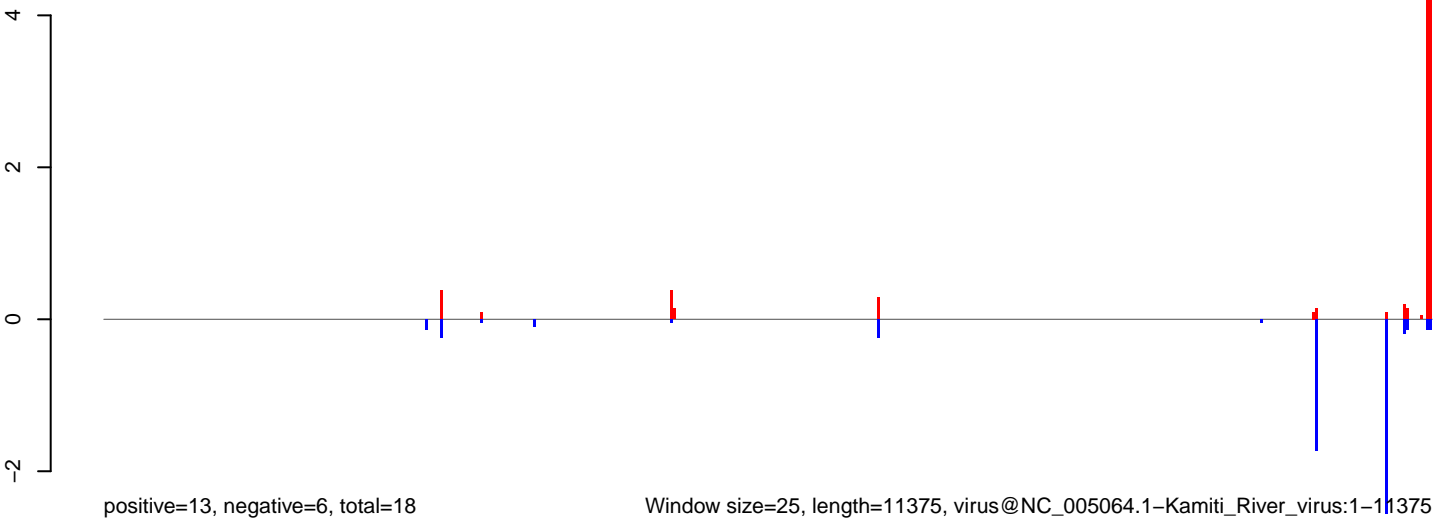
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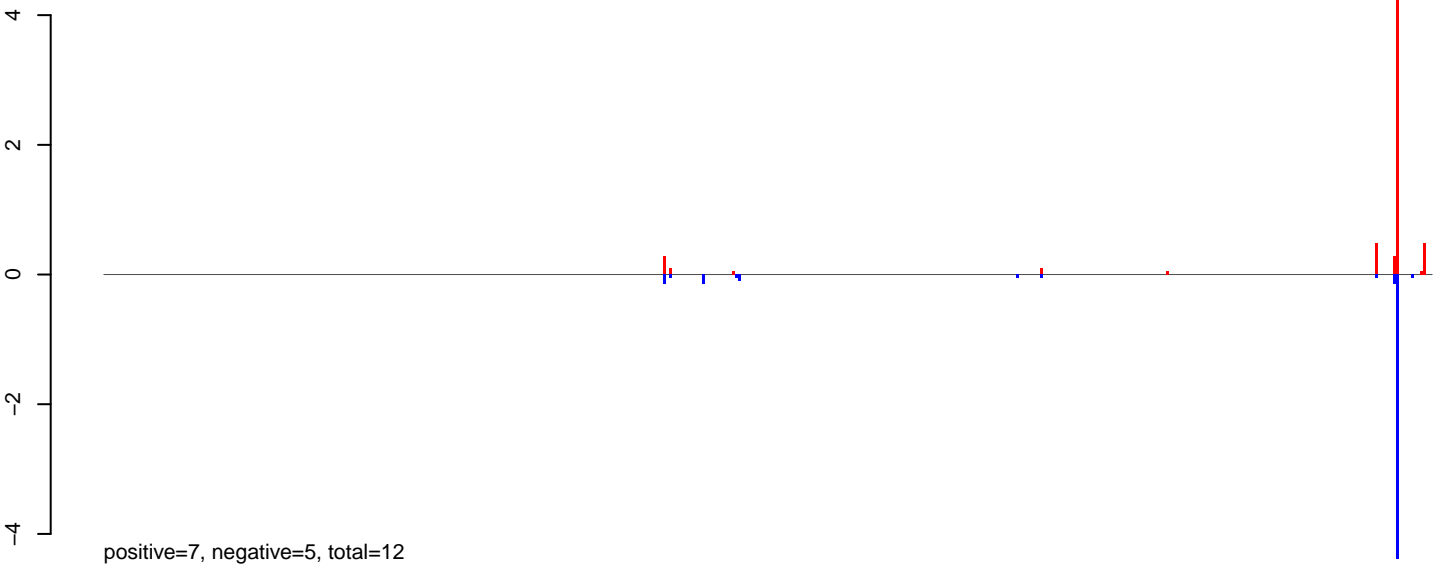
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AeAeg_Aag2_NL.M3.rep



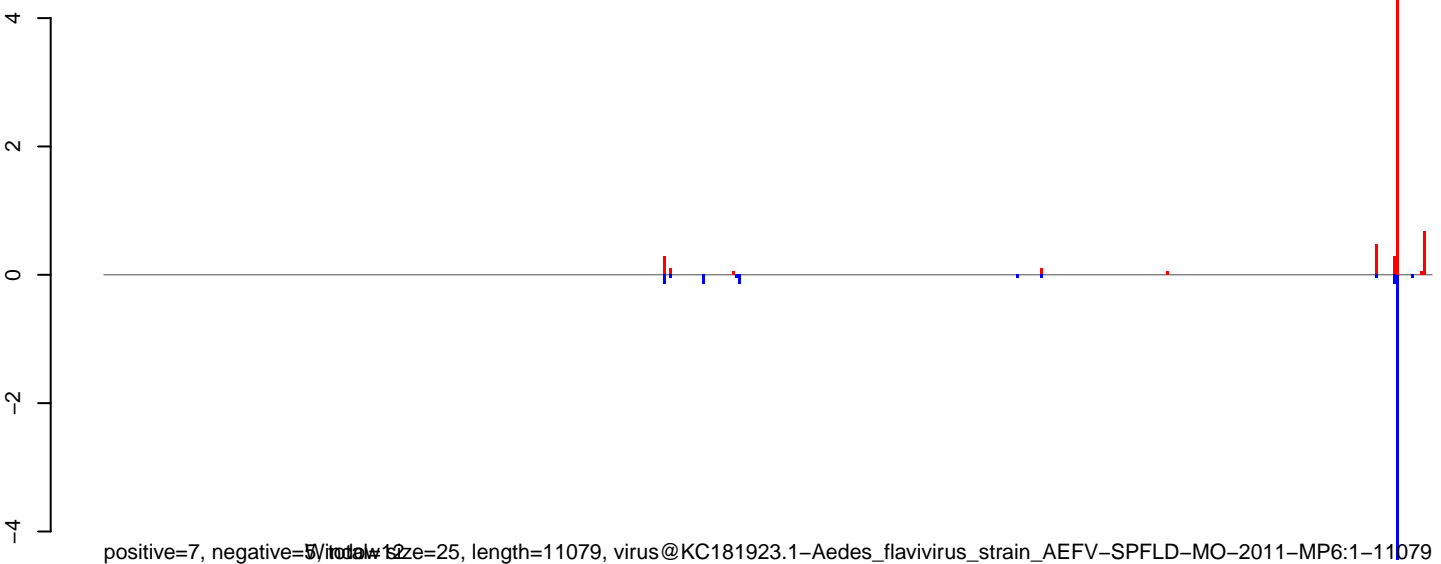
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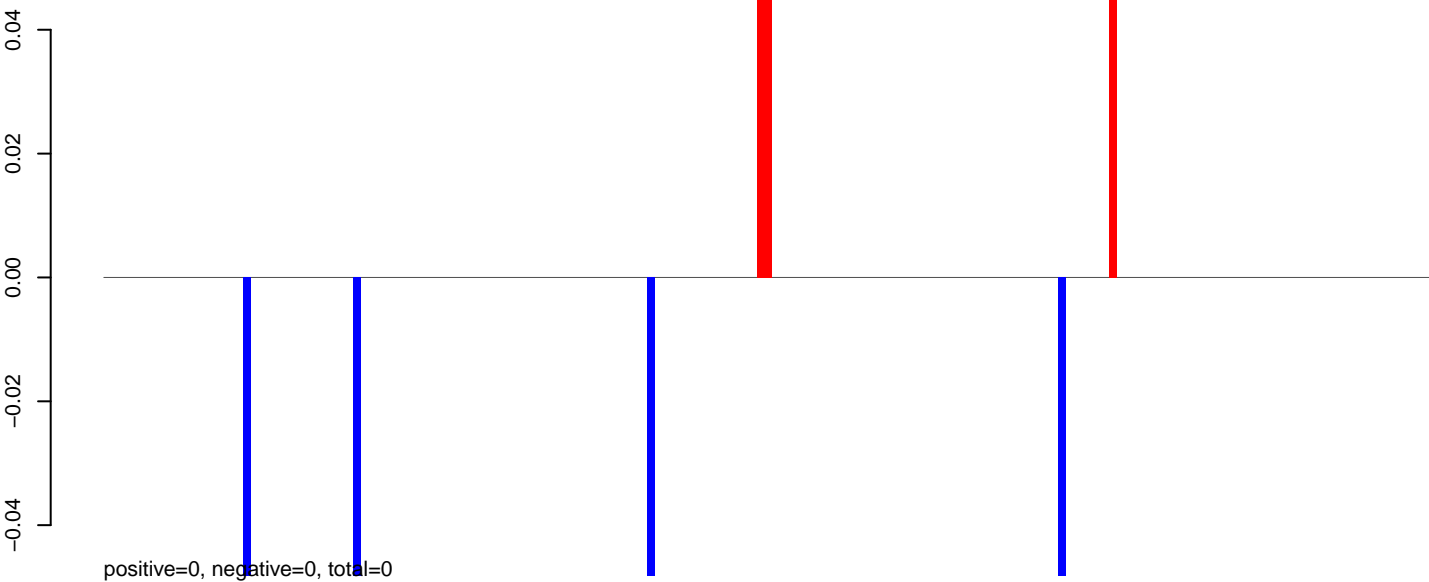
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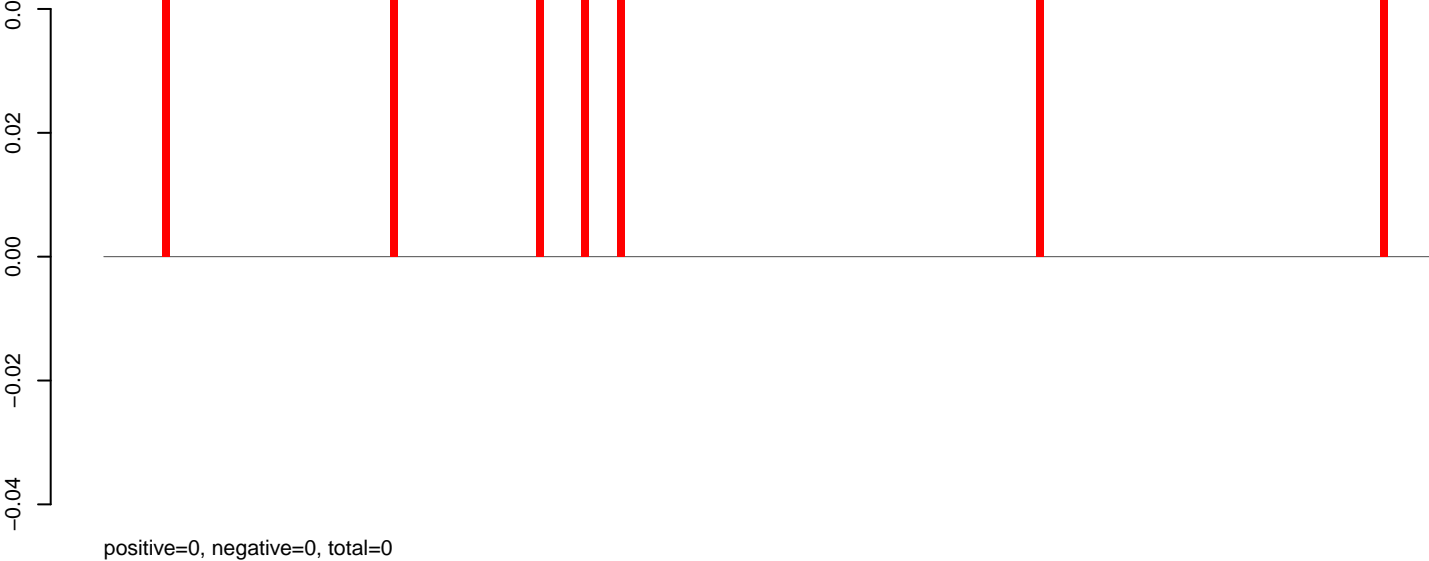
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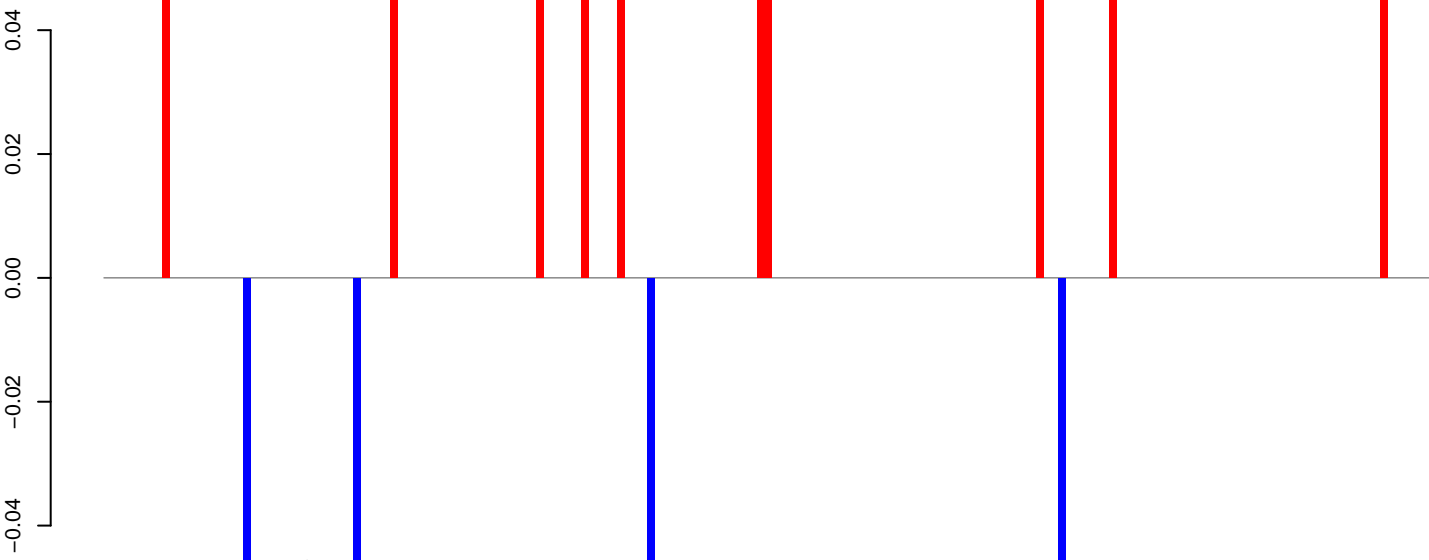
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep



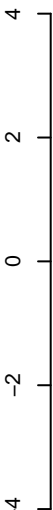
AeAeg_Aag2_NL.M3.rep



ze=25 positive=4, negative=3, total=1-Drosophila_melanogaster_American_nodavirus_ANV_SW-2009a_segment_RNA1_and_RNA2:1-4523

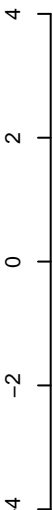
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AeAeg_Aag2_NL.M3.18_23.rep



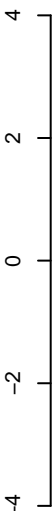
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AeAeg_Aag2_NL.M3.24_35.rep



positive=0, negative=0, total=1

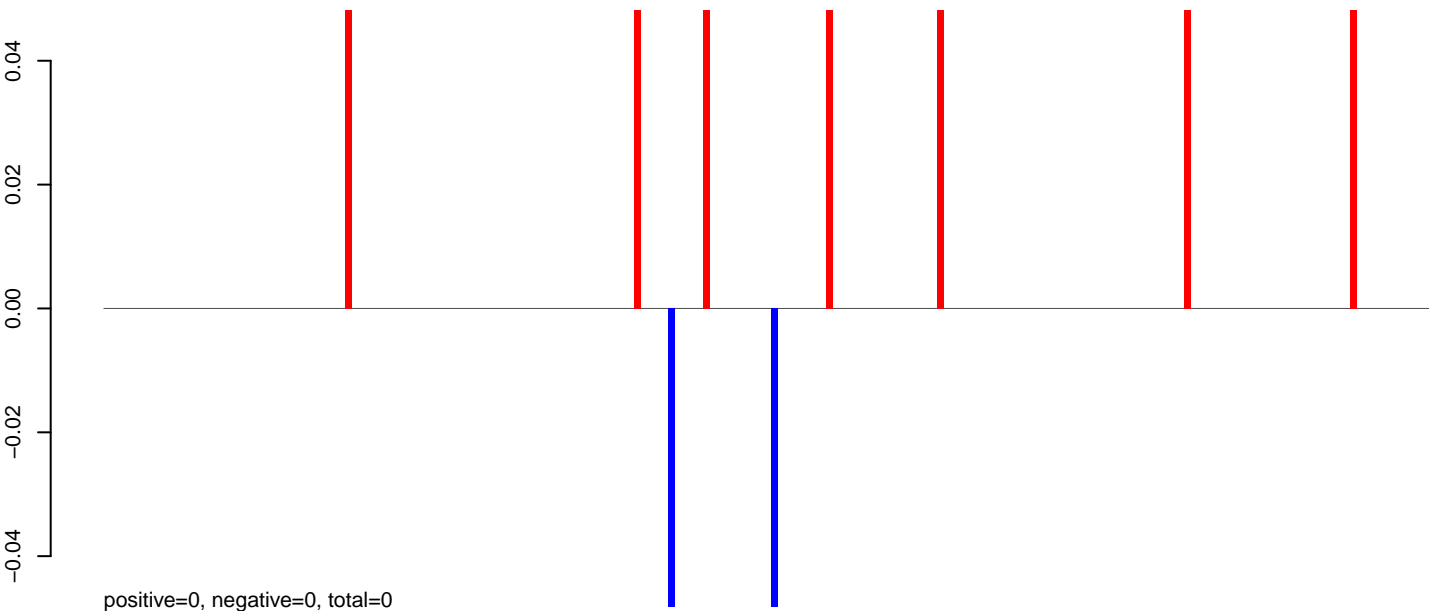
AeAeg_Aag2_NL.M3.rep



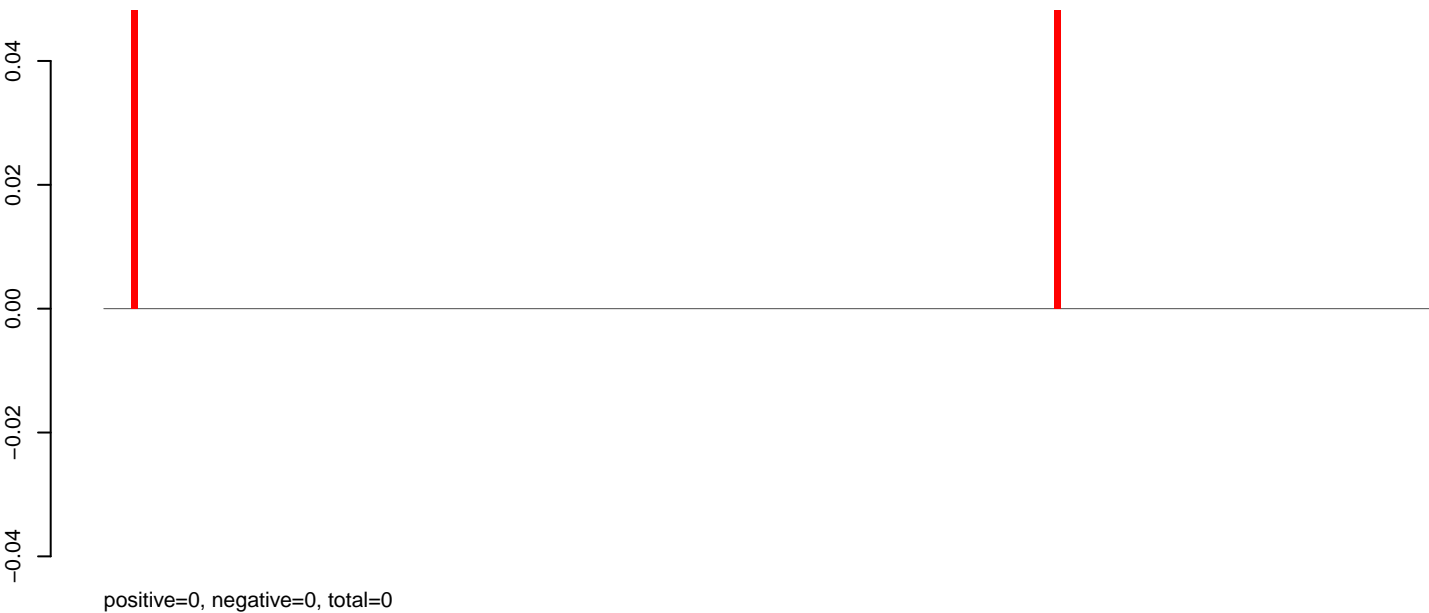
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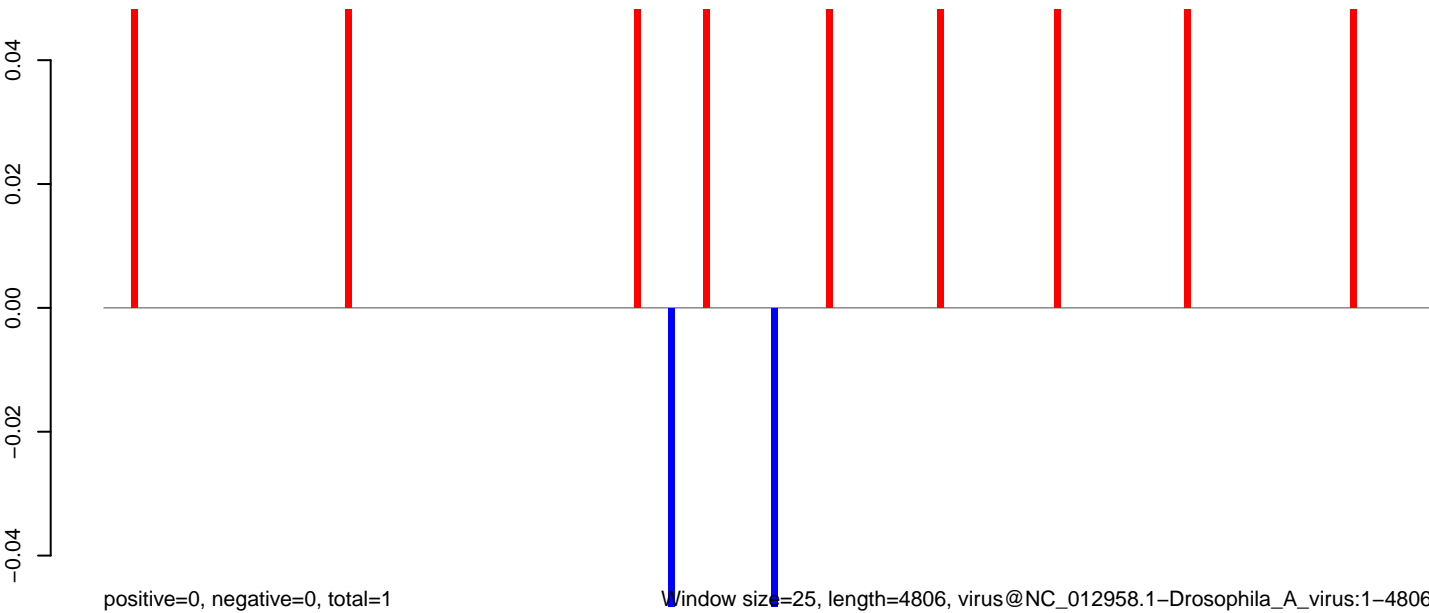
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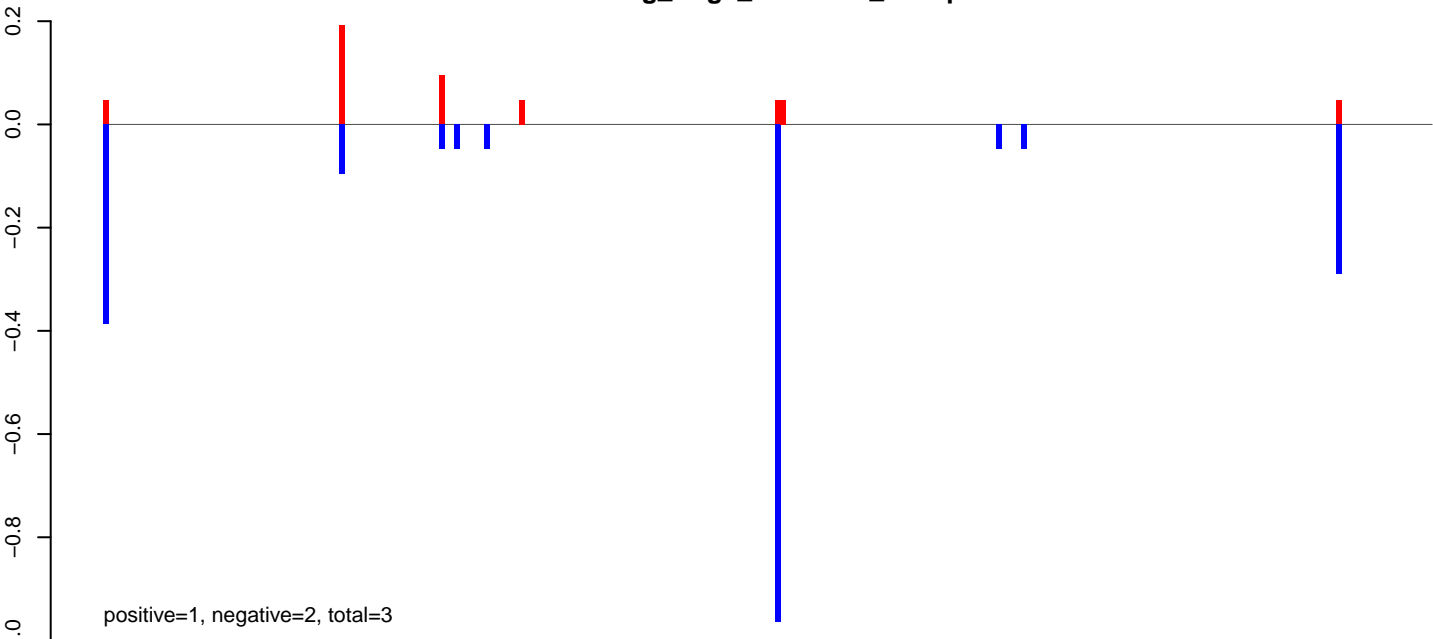
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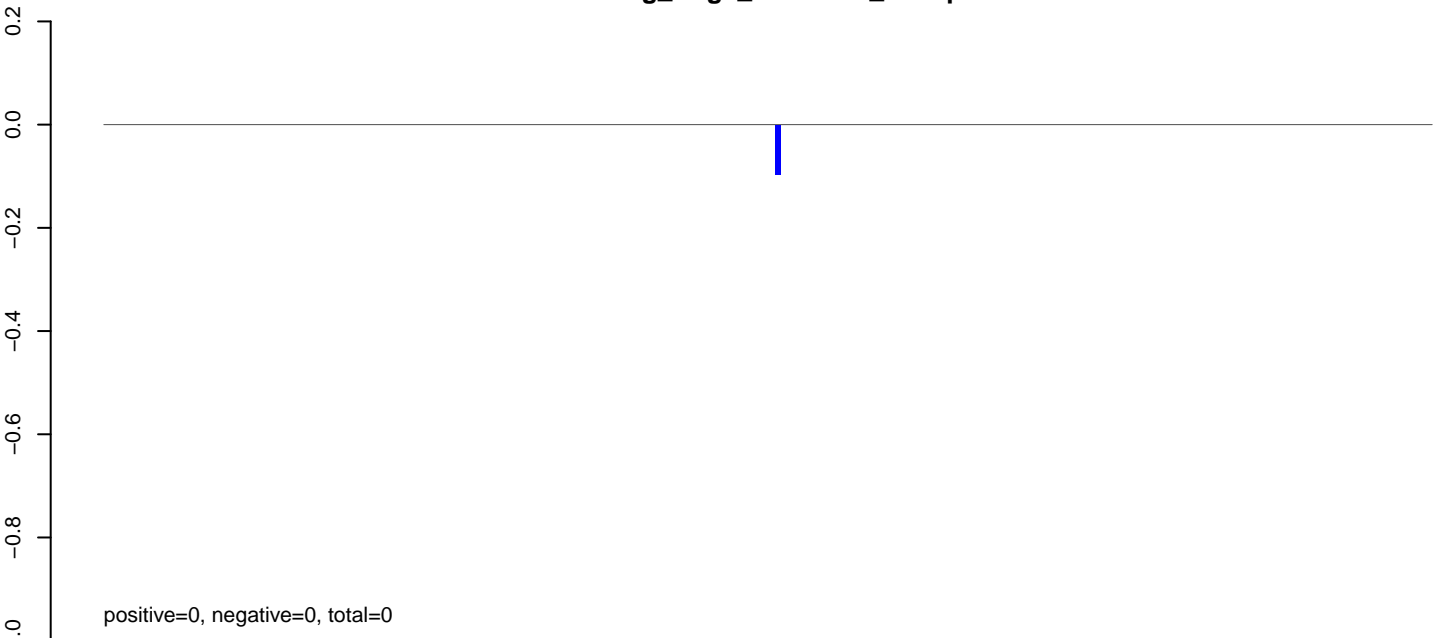
AeAeg_Aag2_NL.M3.rep



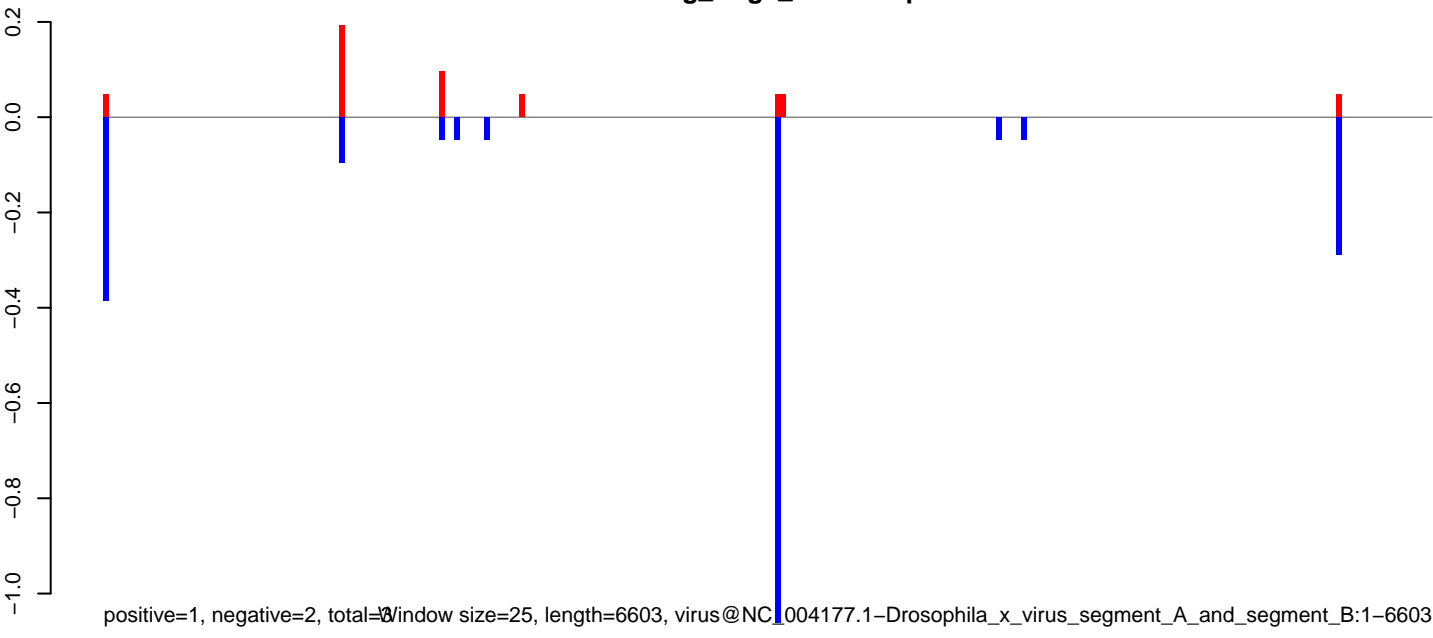
AeAeg_Aag2_NL.M3.18_23.rep



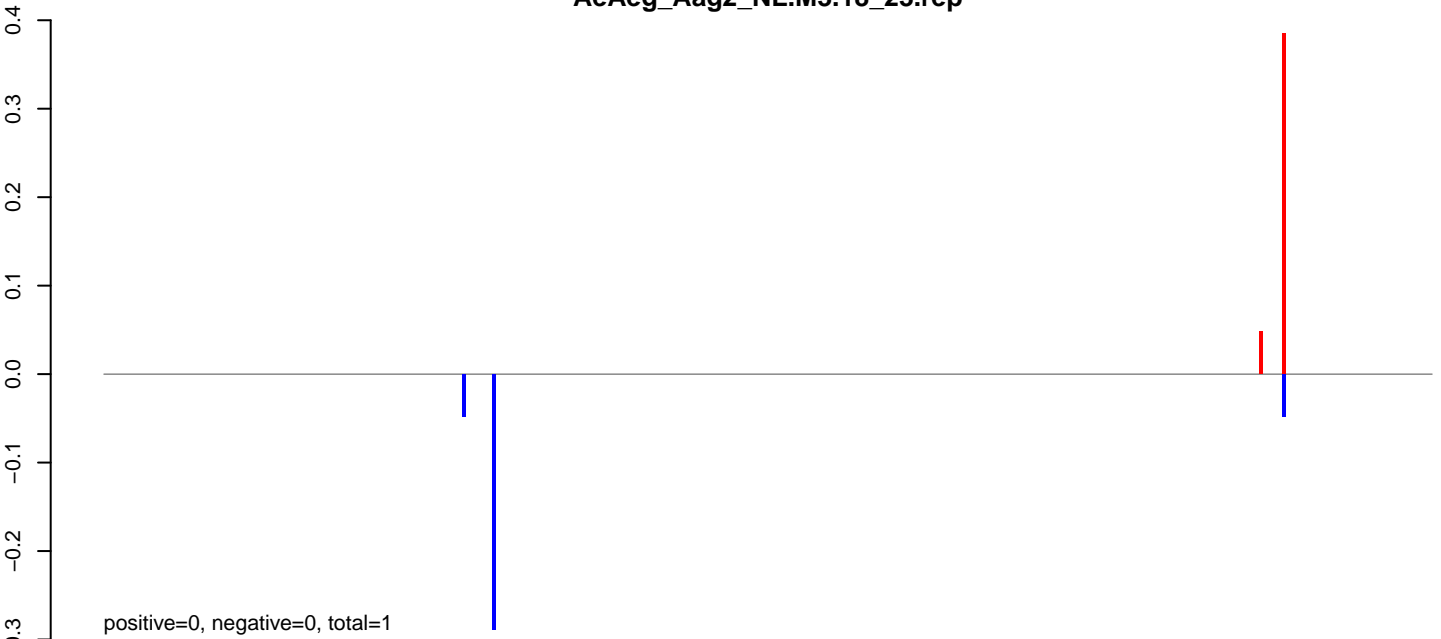
AeAeg_Aag2_NL.M3.24_35.rep



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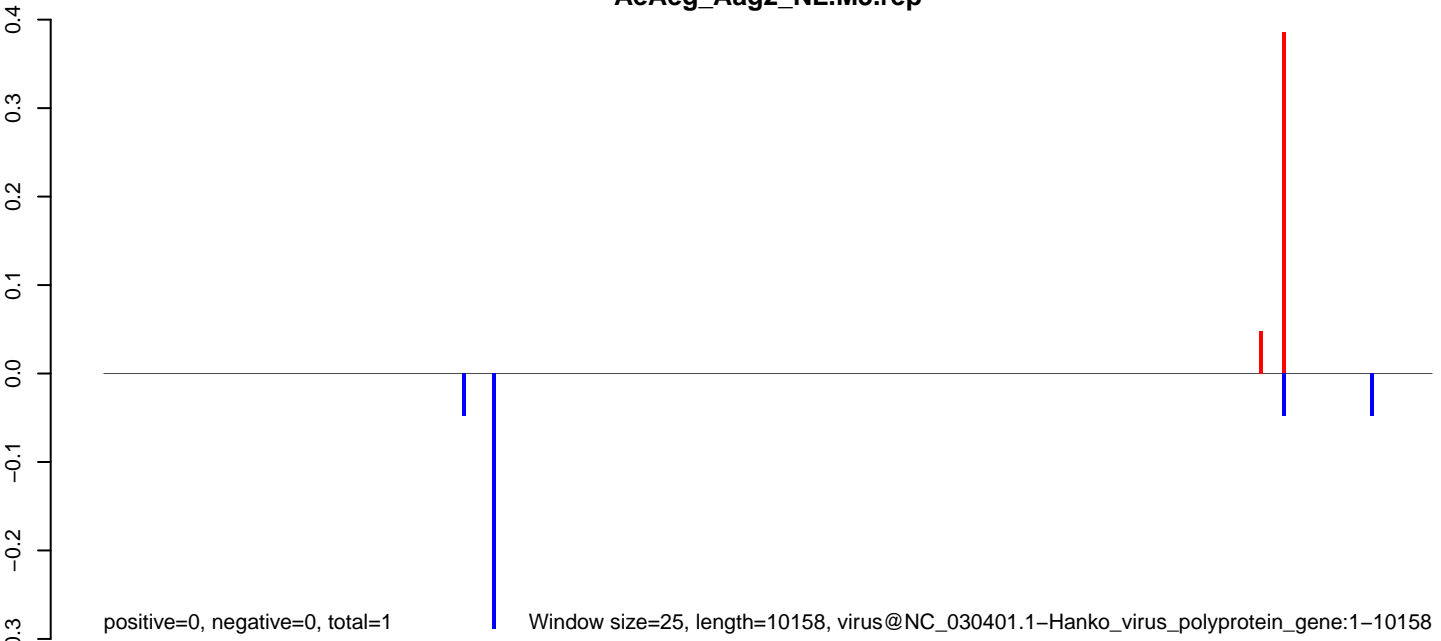
AeAeg_Aag2_NL.M3.18_23.rep



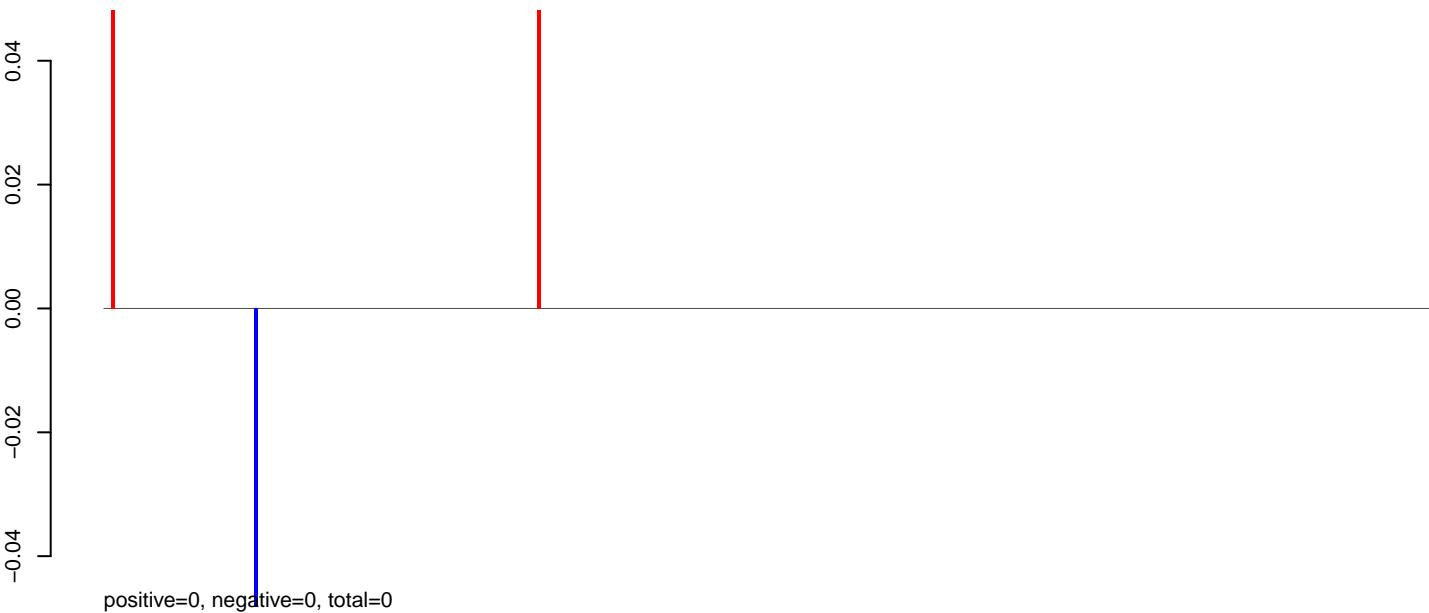
AeAeg_Aag2_NL.M3.24_35.rep



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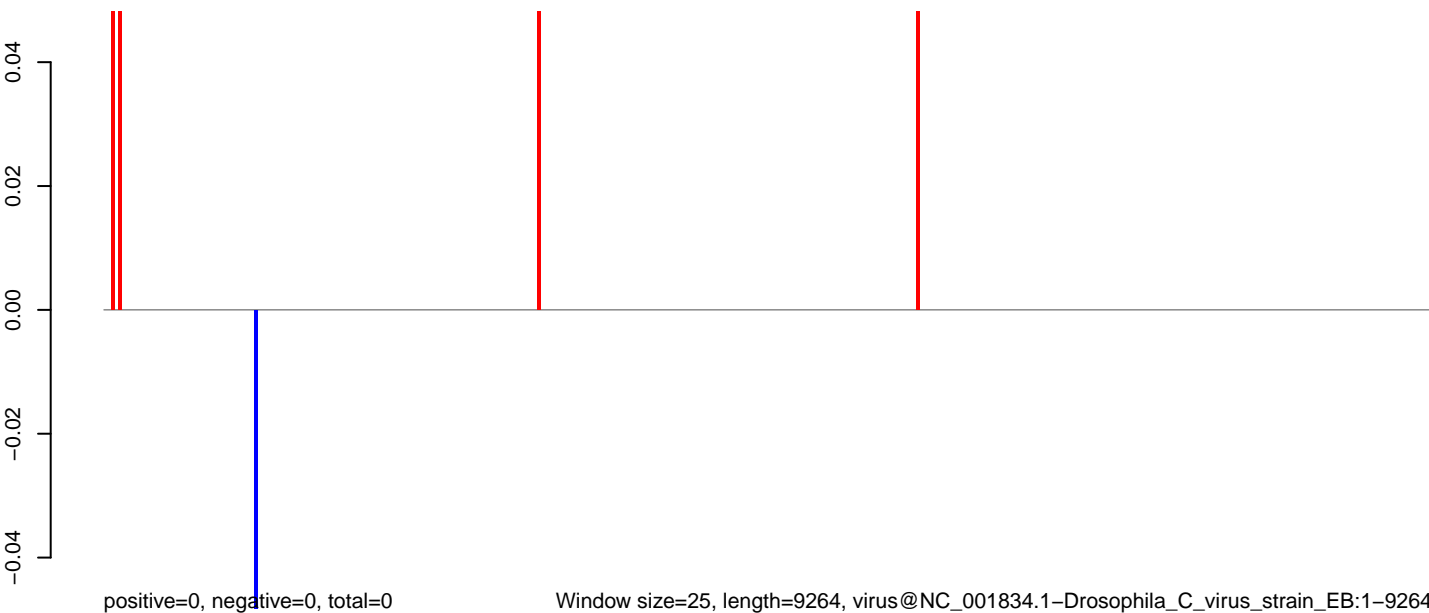
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep



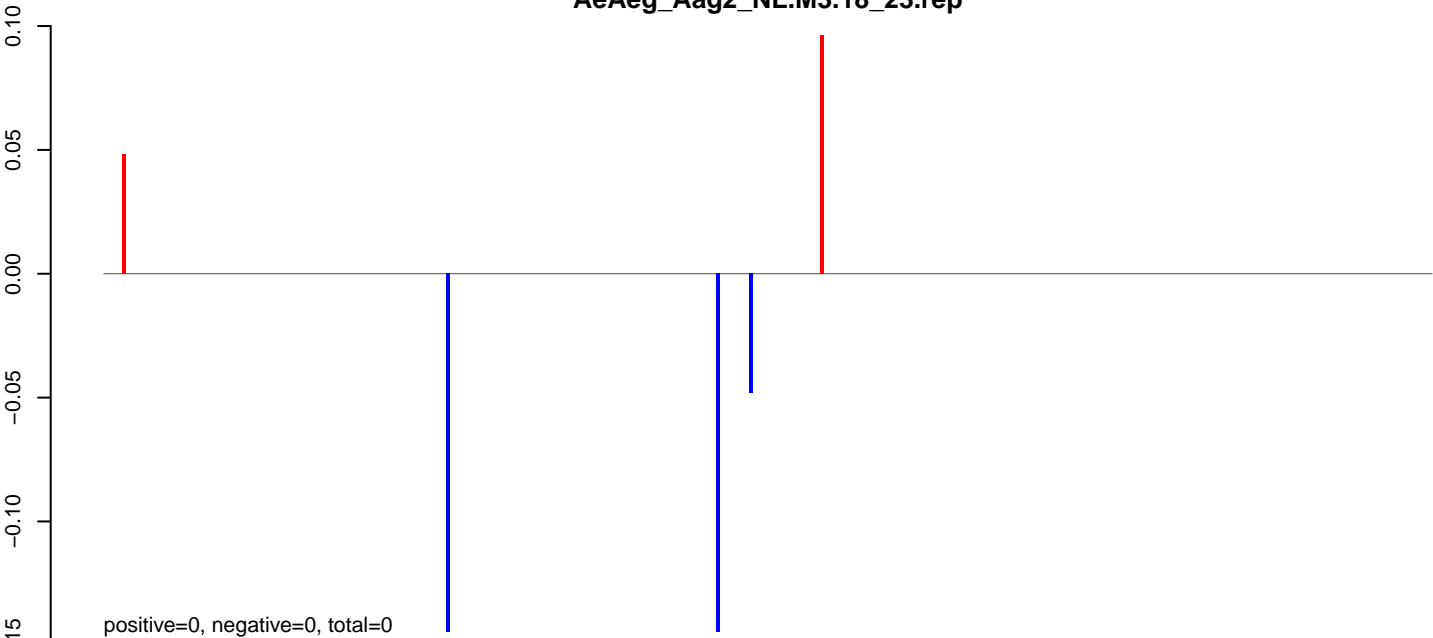
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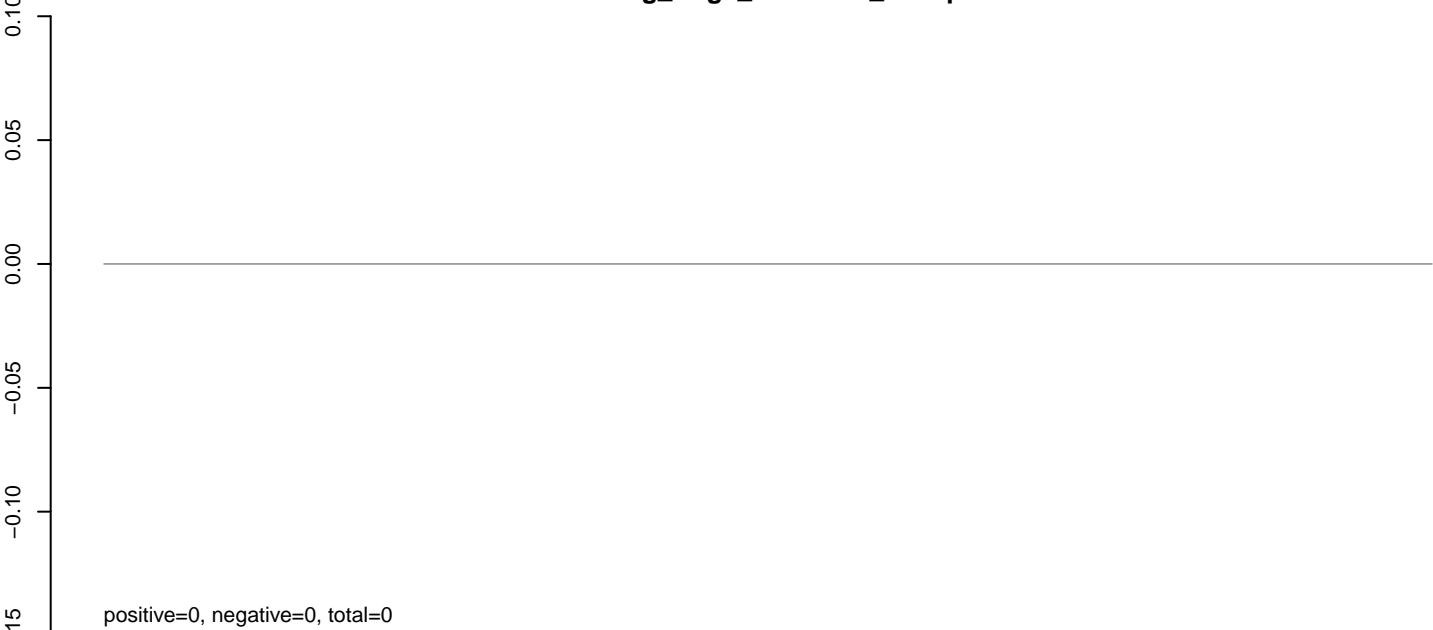
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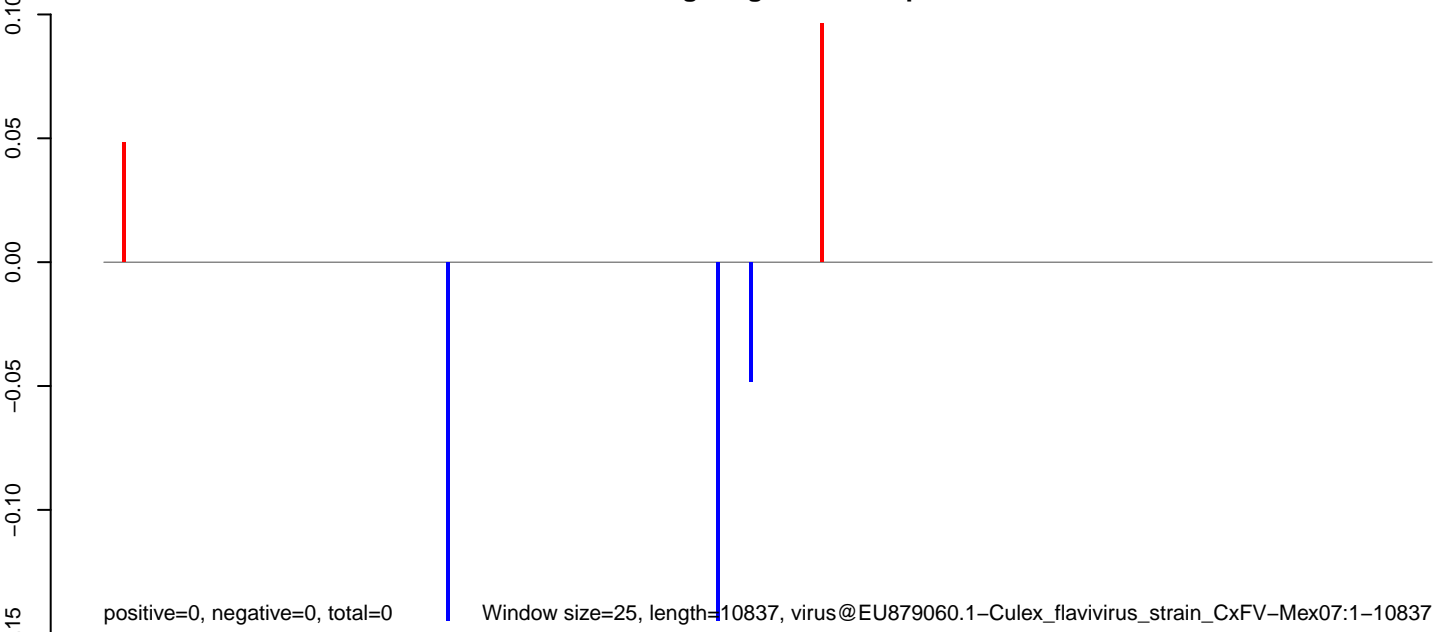
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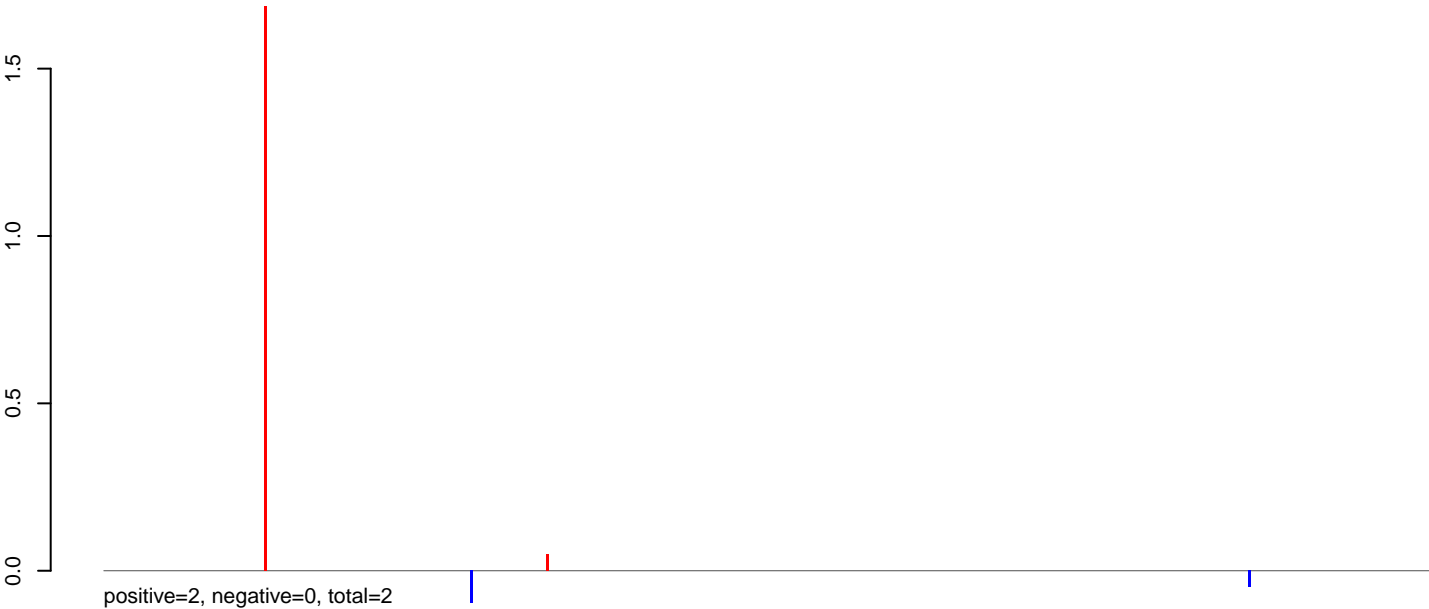
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AeAeg_Aag2_NL.M3.rep



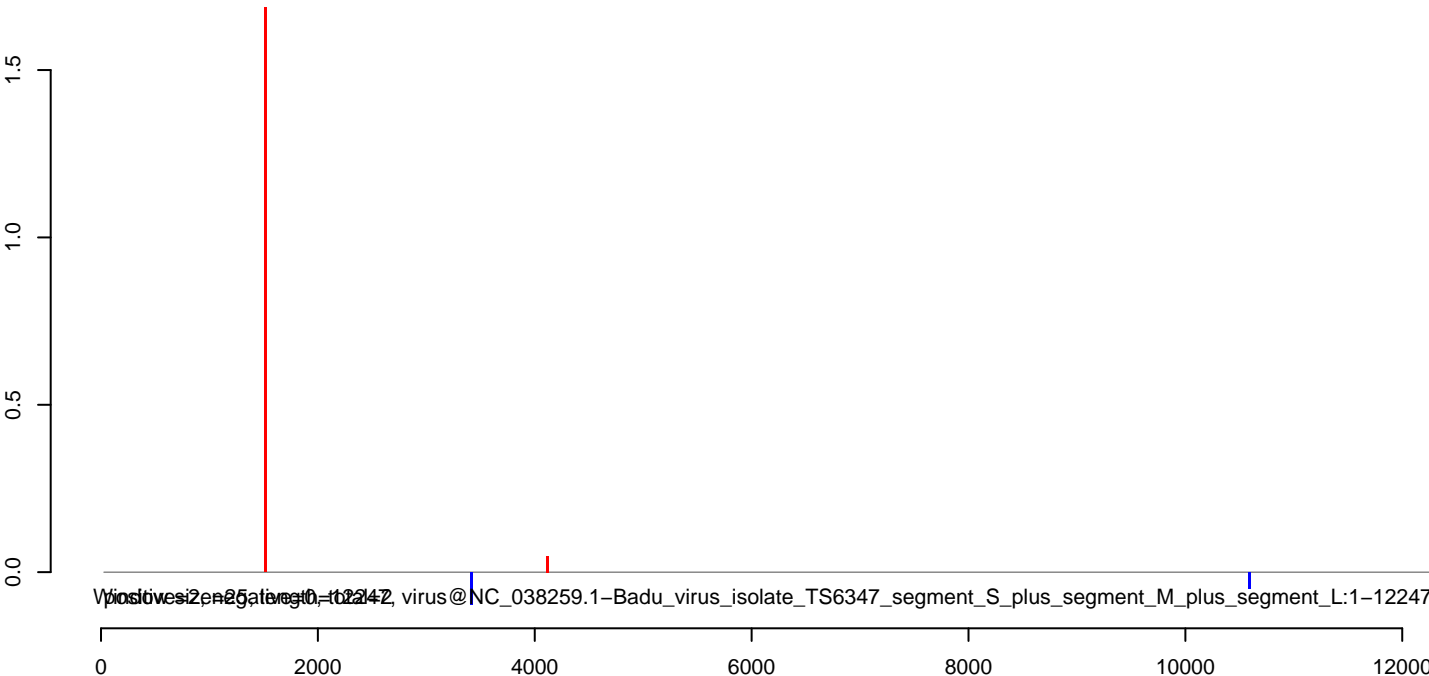
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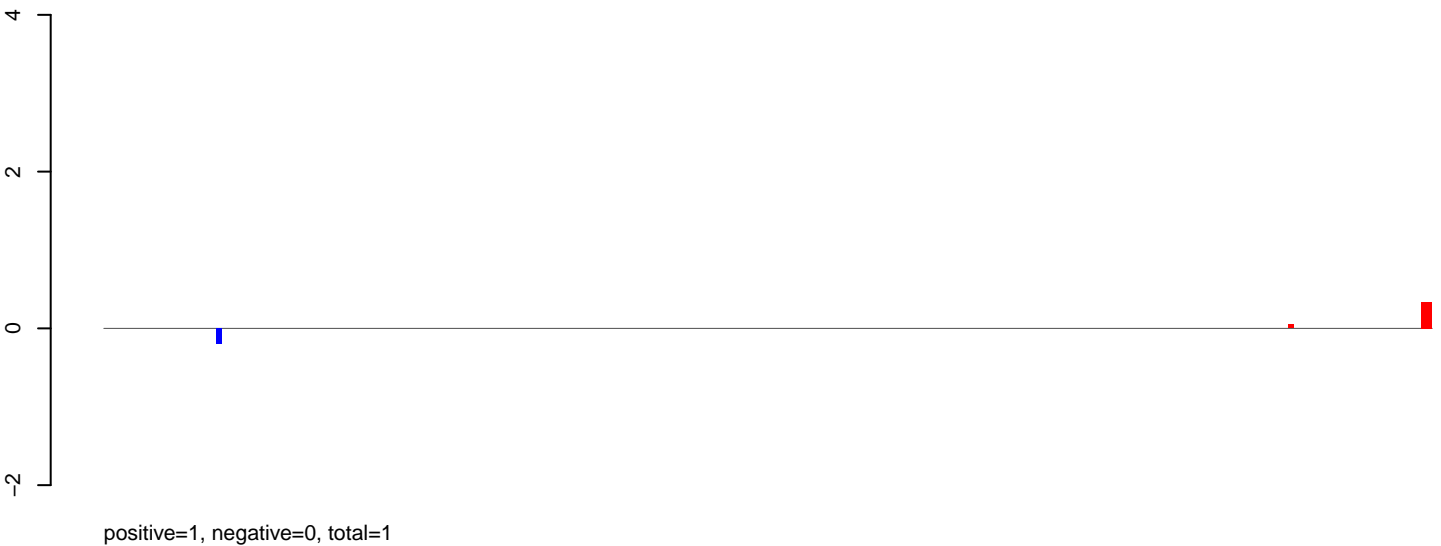
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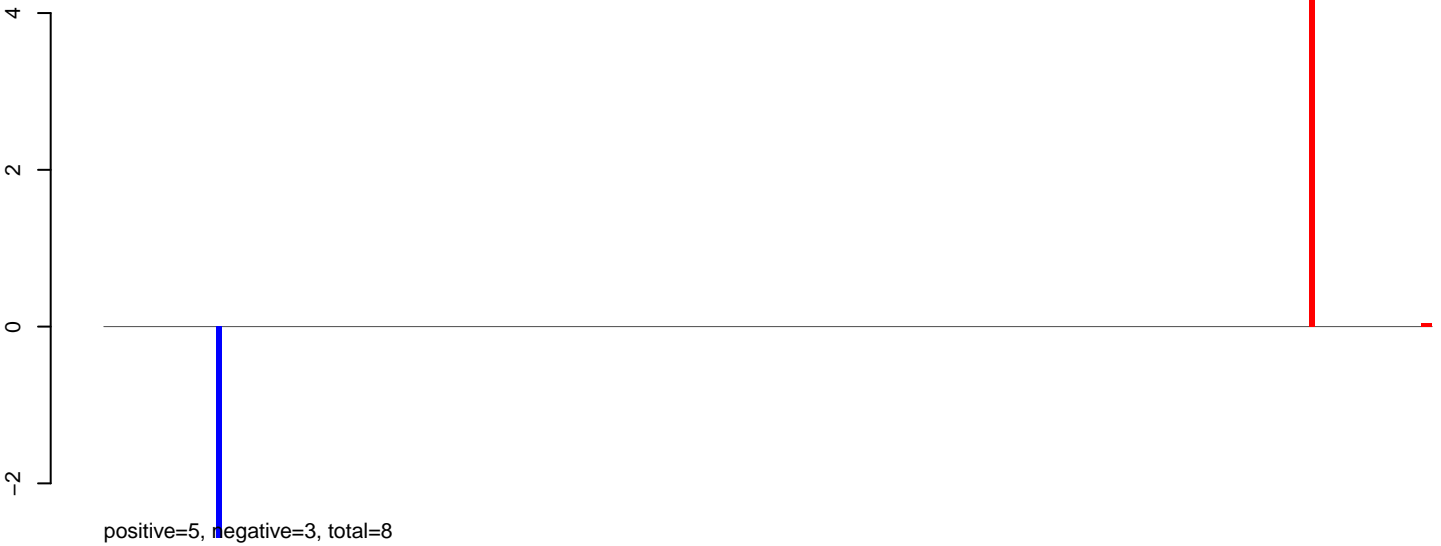
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AeAeg_Aag2_NL.M3.18_23.rep



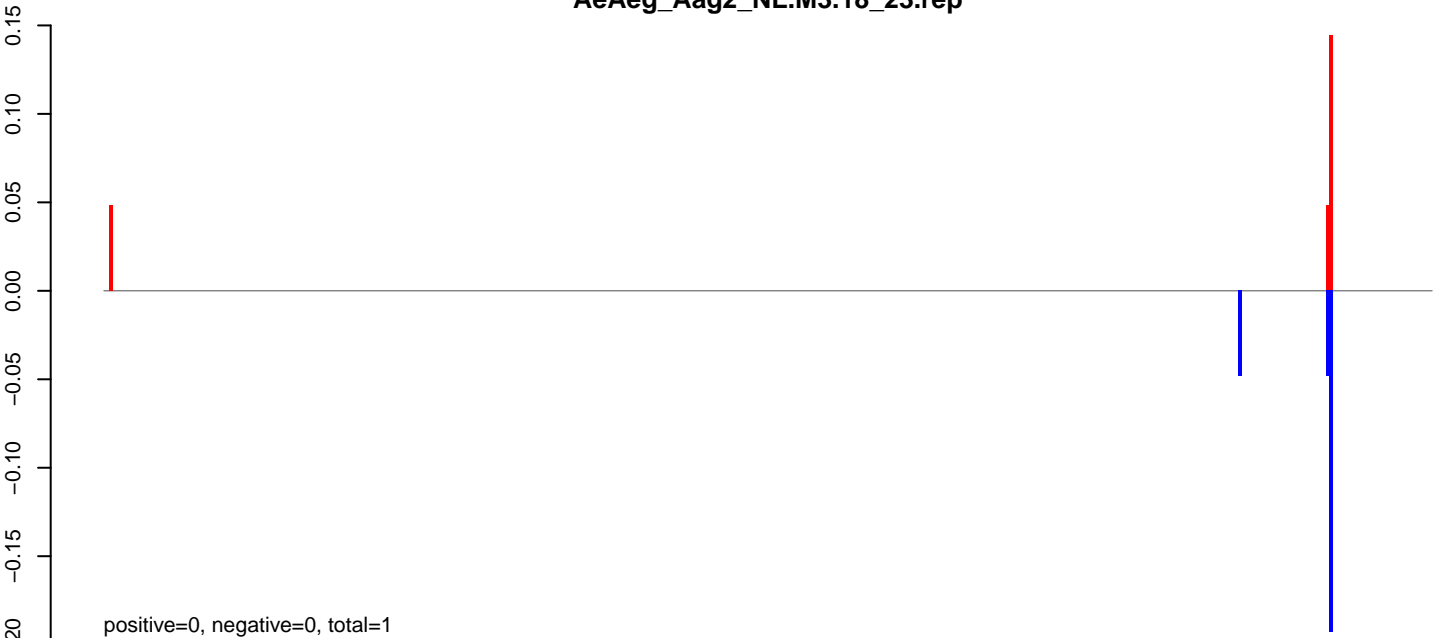
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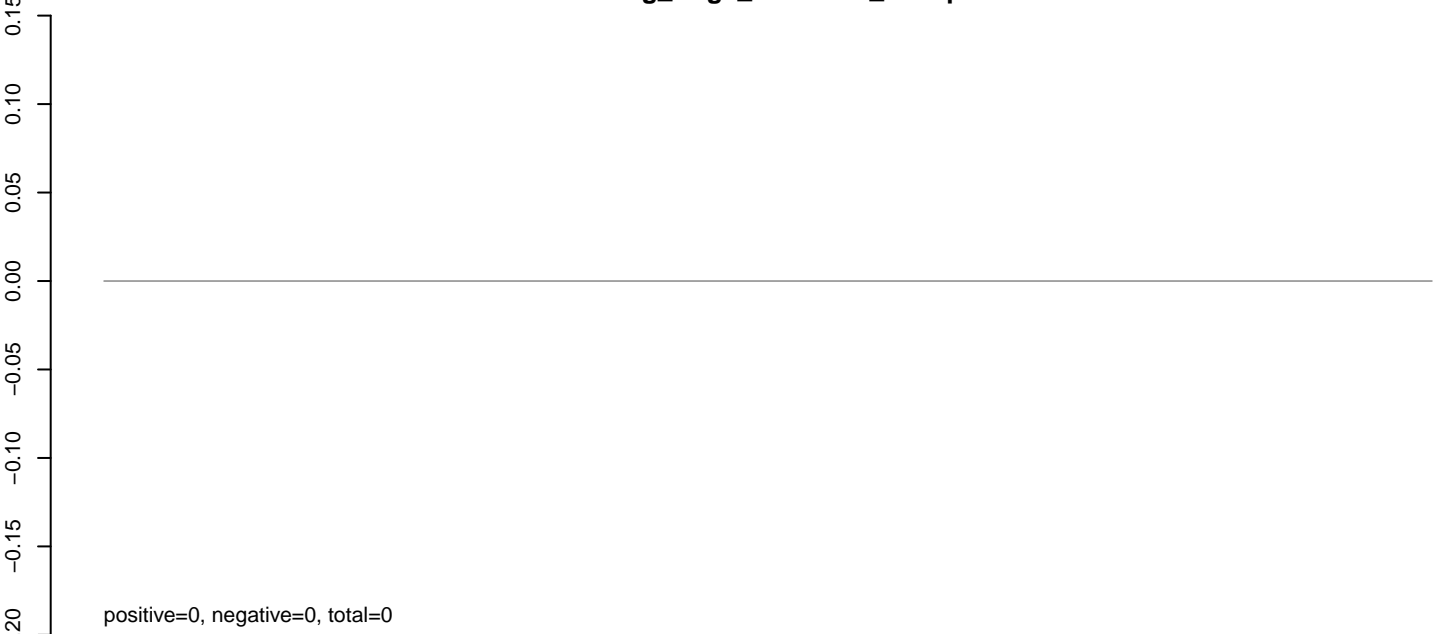
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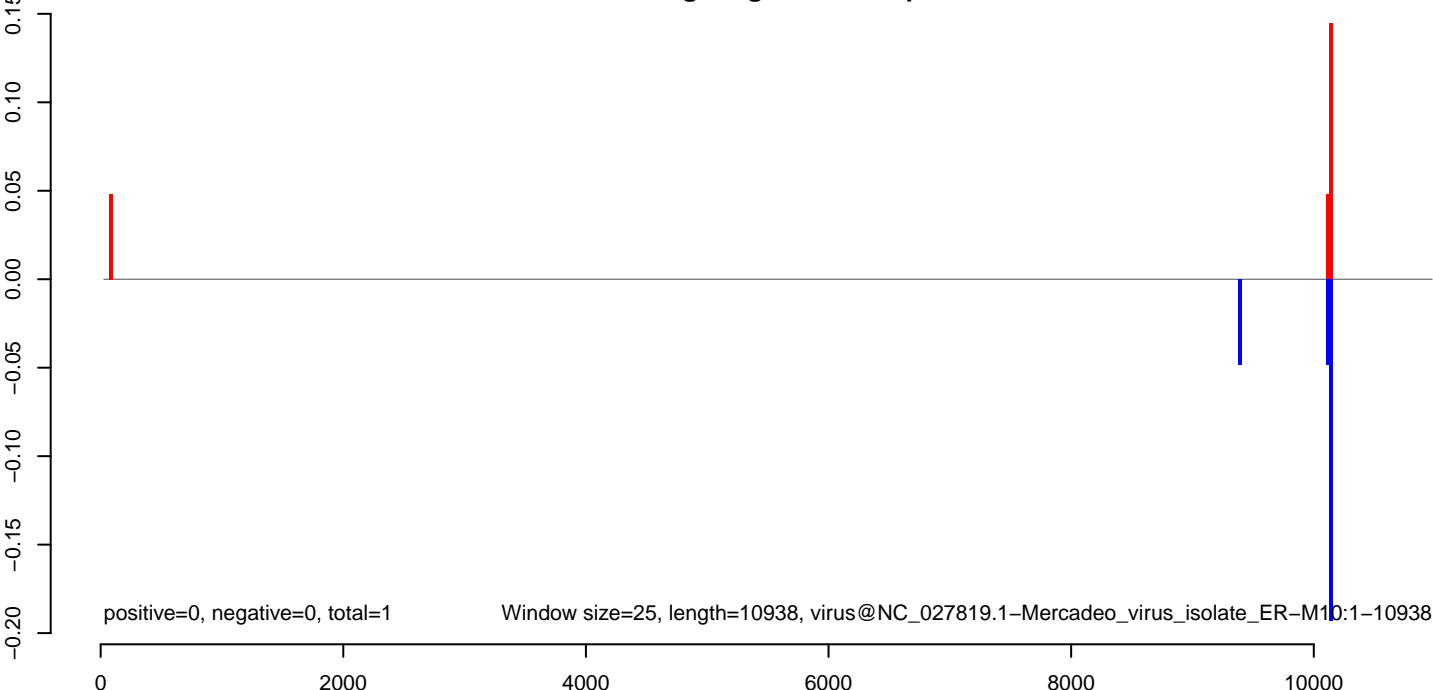
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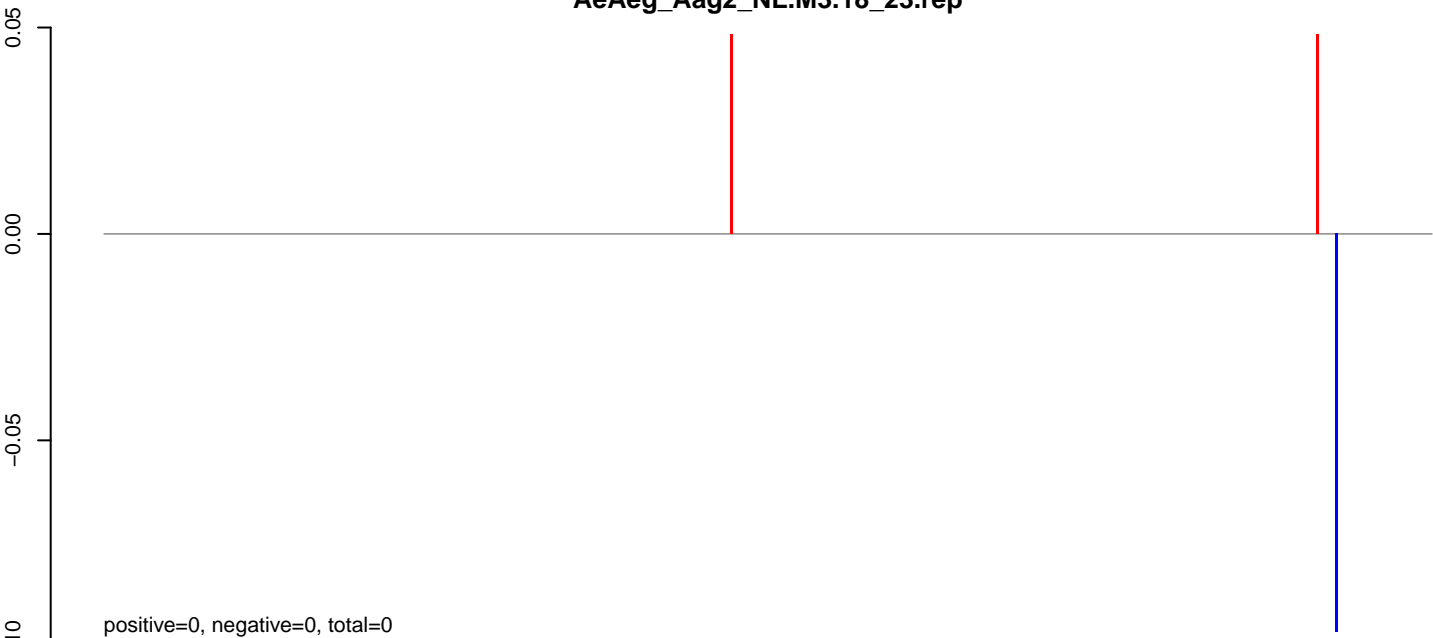
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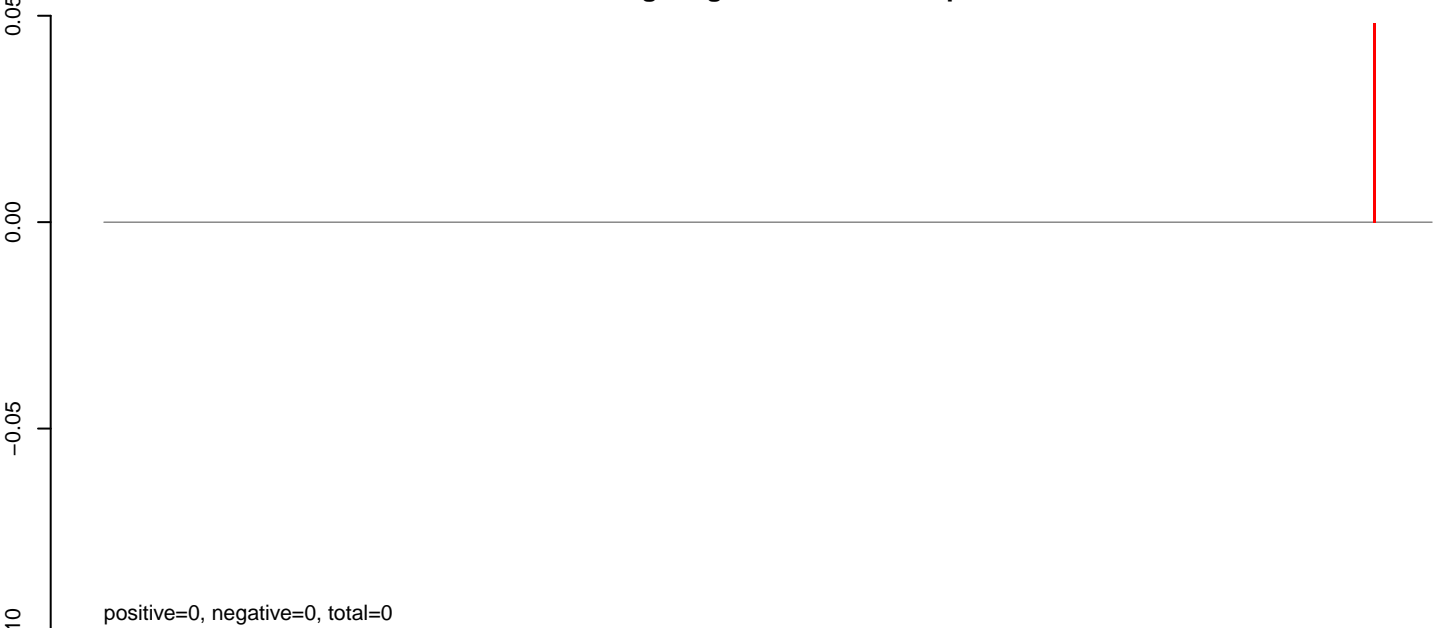
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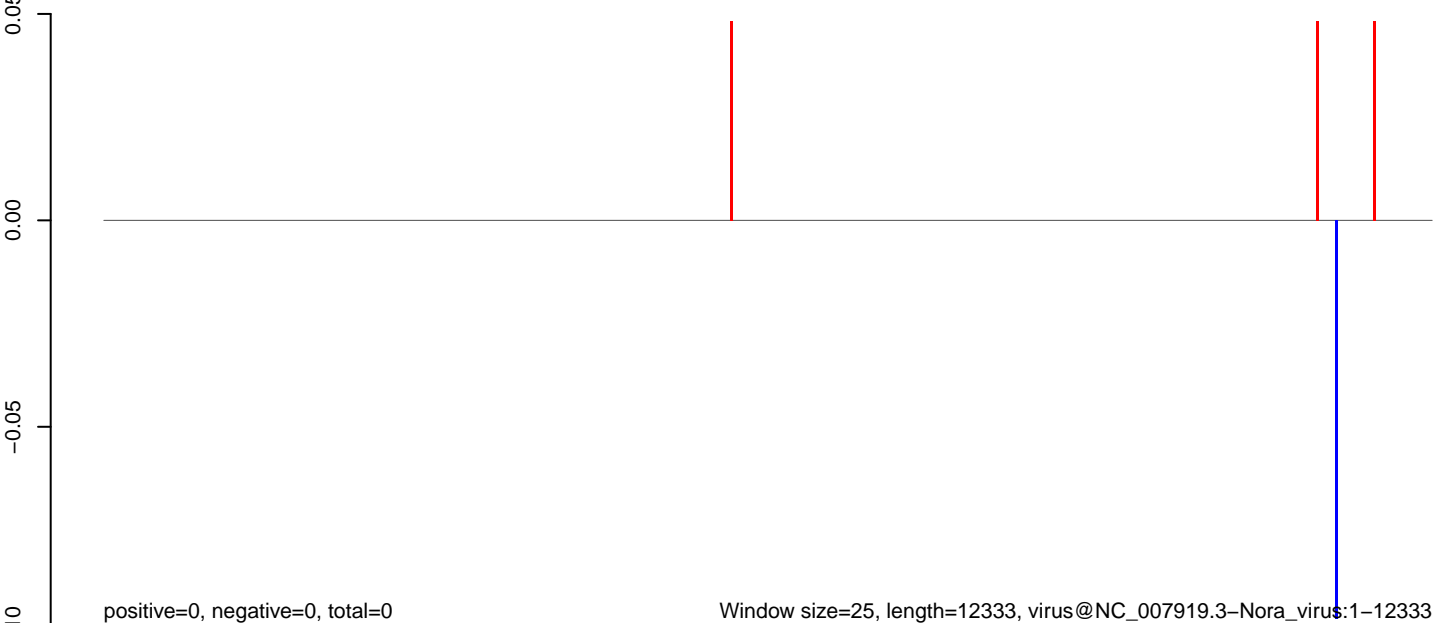
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AeAeg_Aag2_NL.M3.24_35.rep

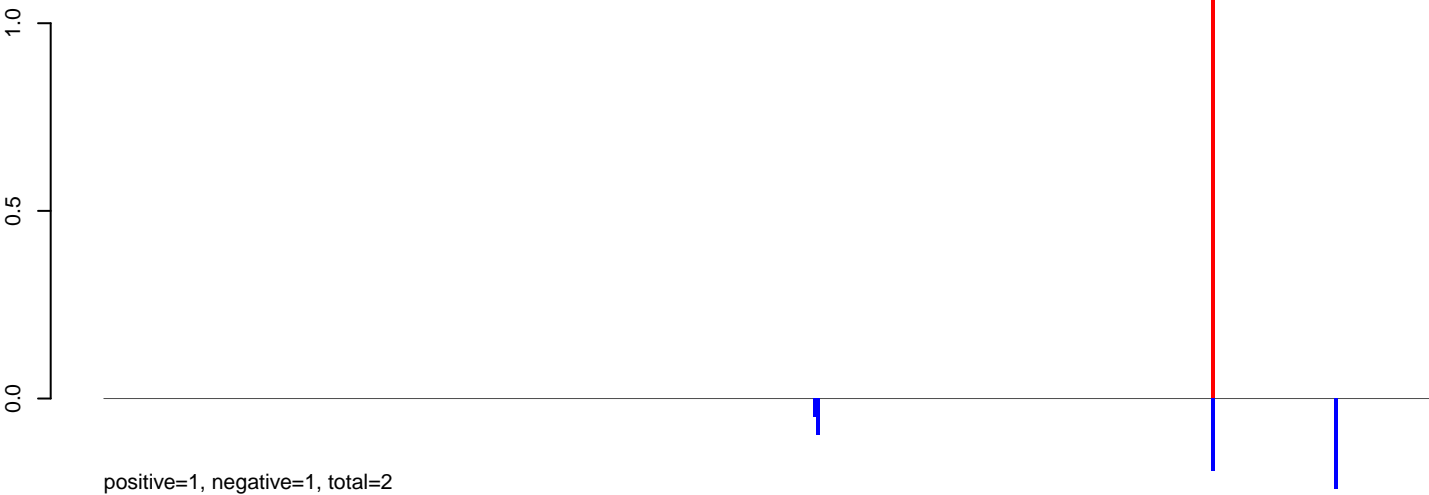


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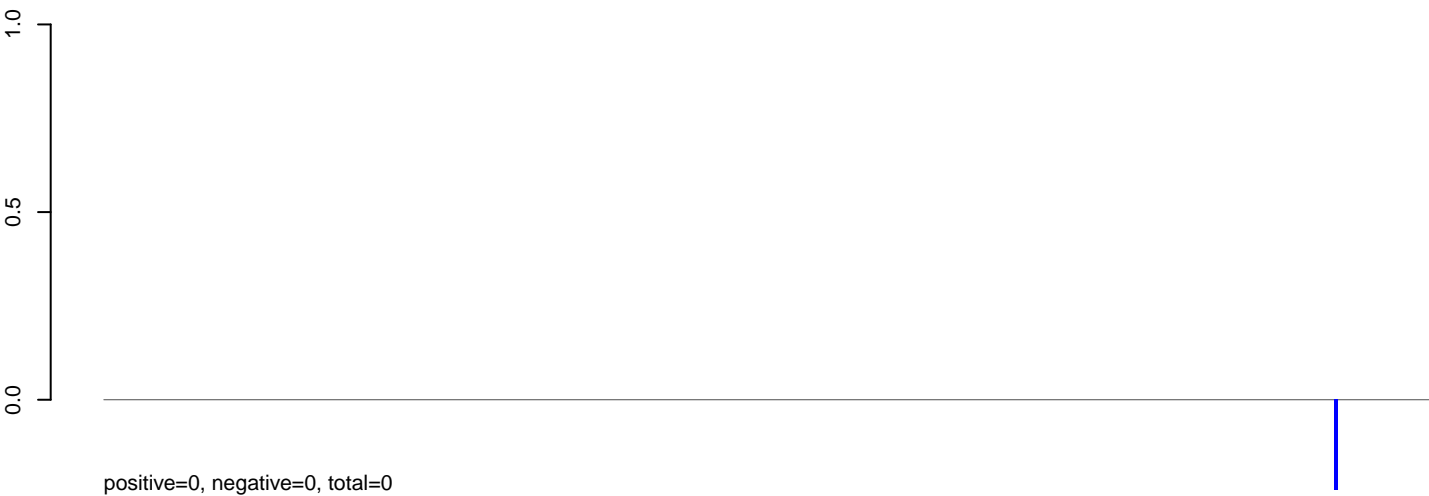


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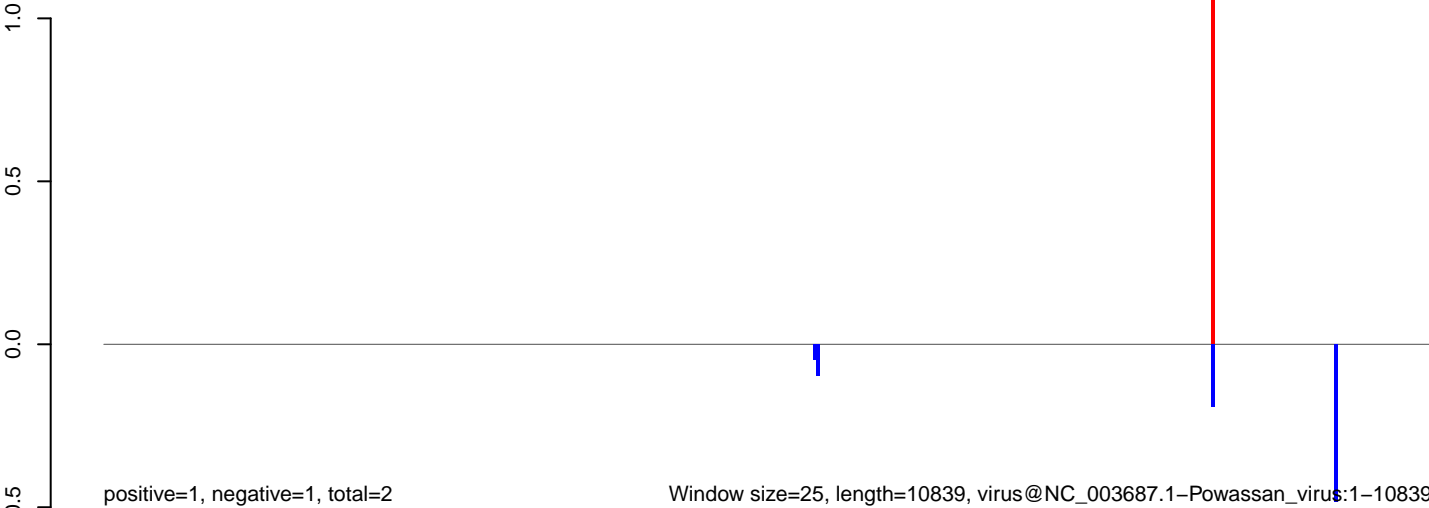
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AeAeg_Aag2_NL.M3.24_35.rep



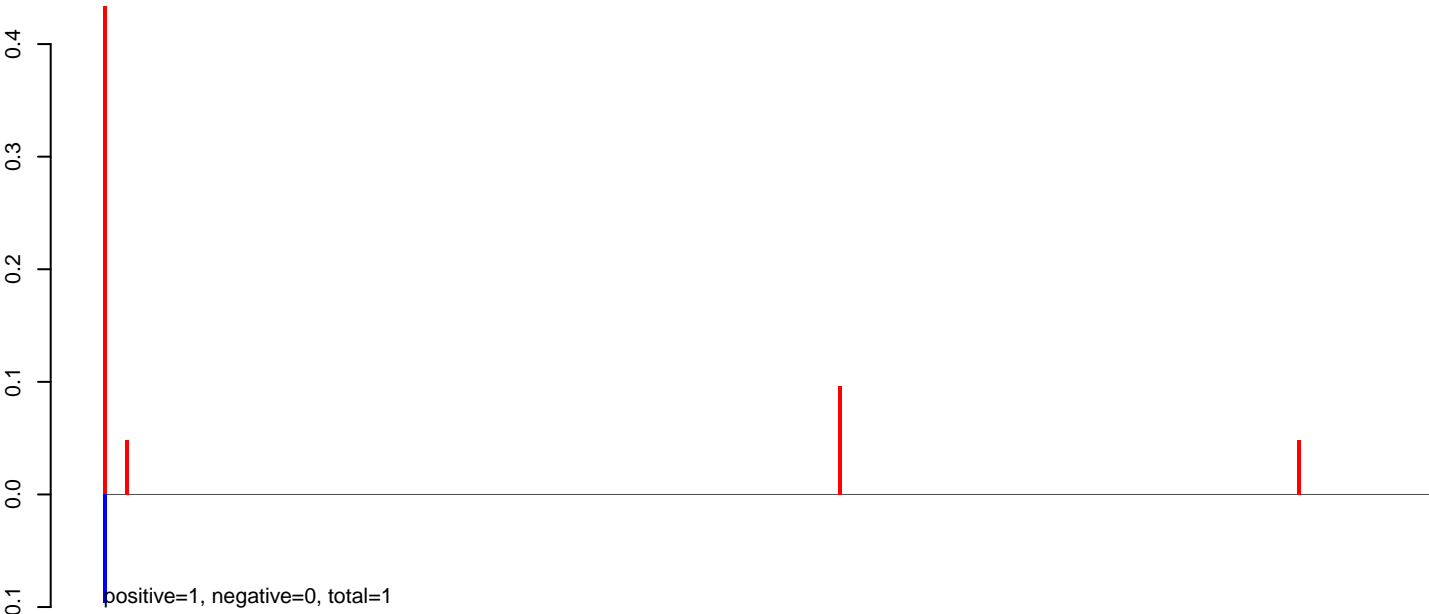
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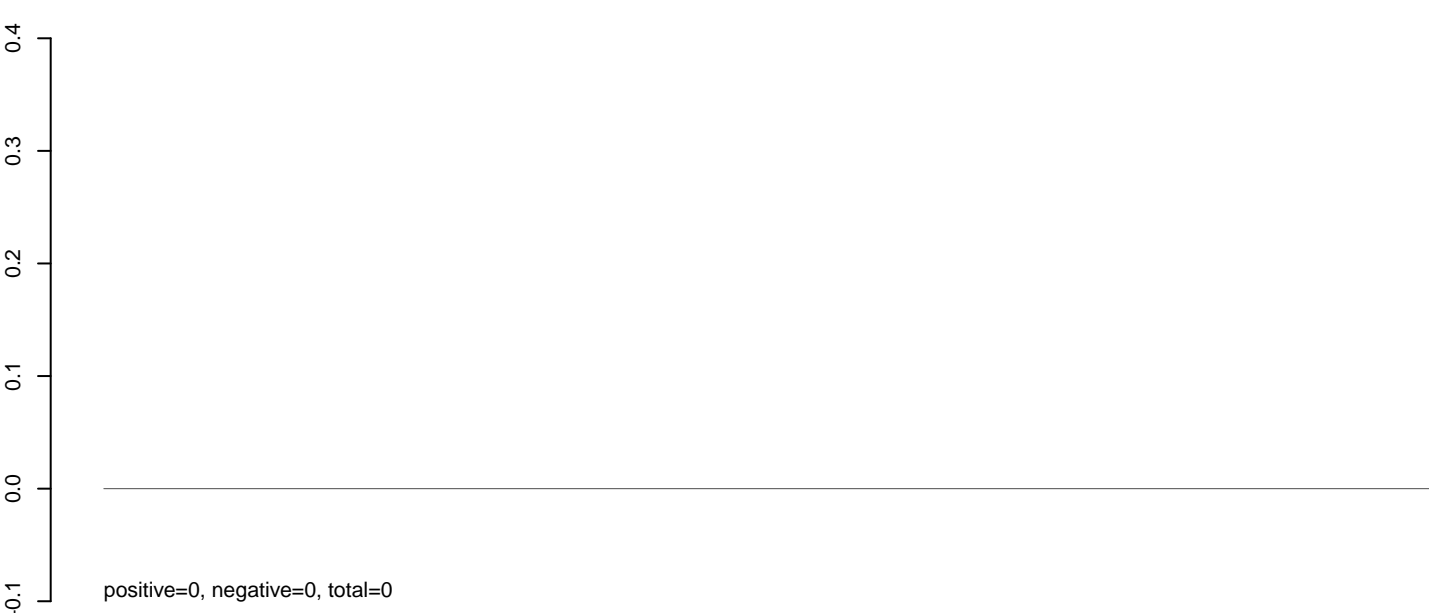
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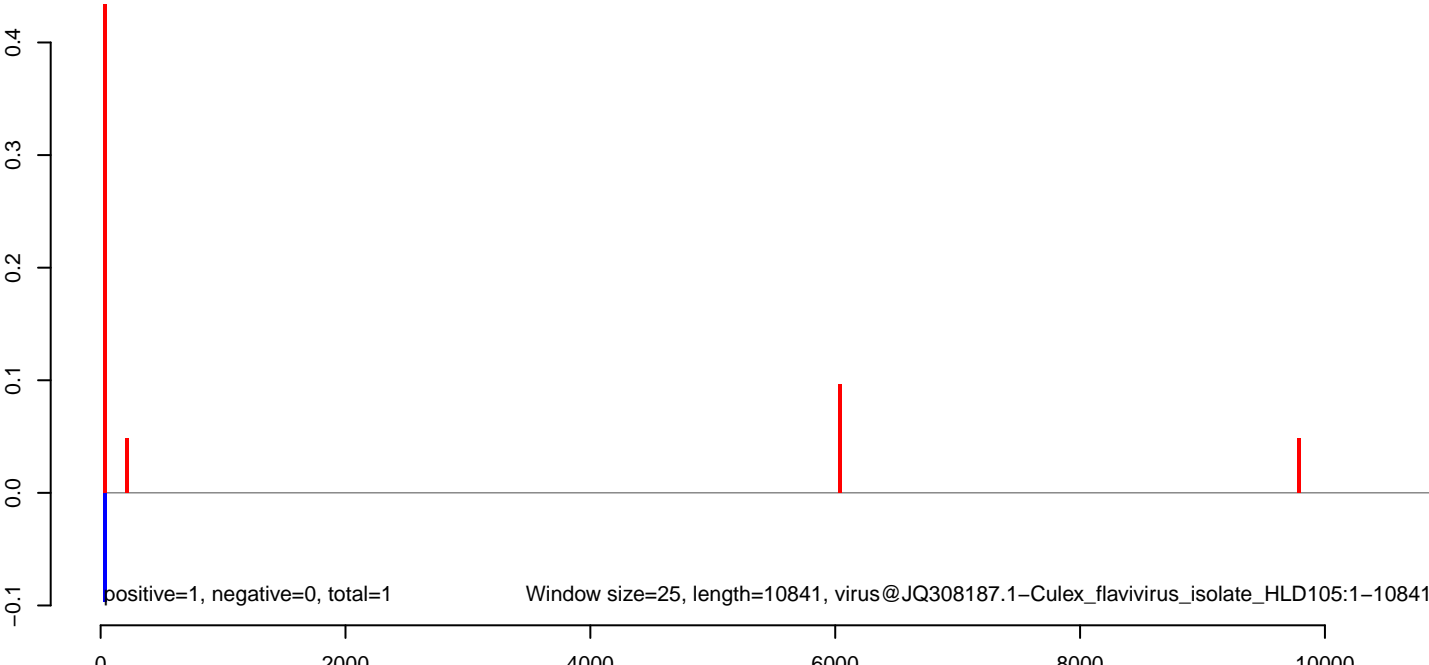
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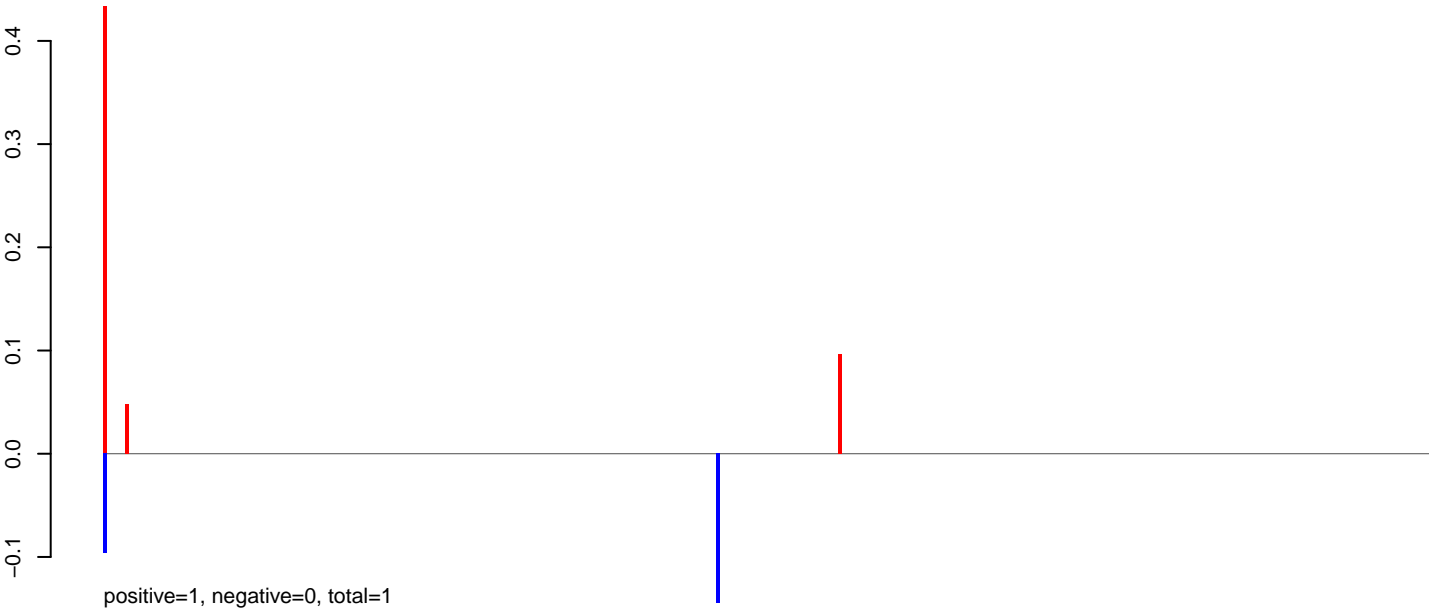
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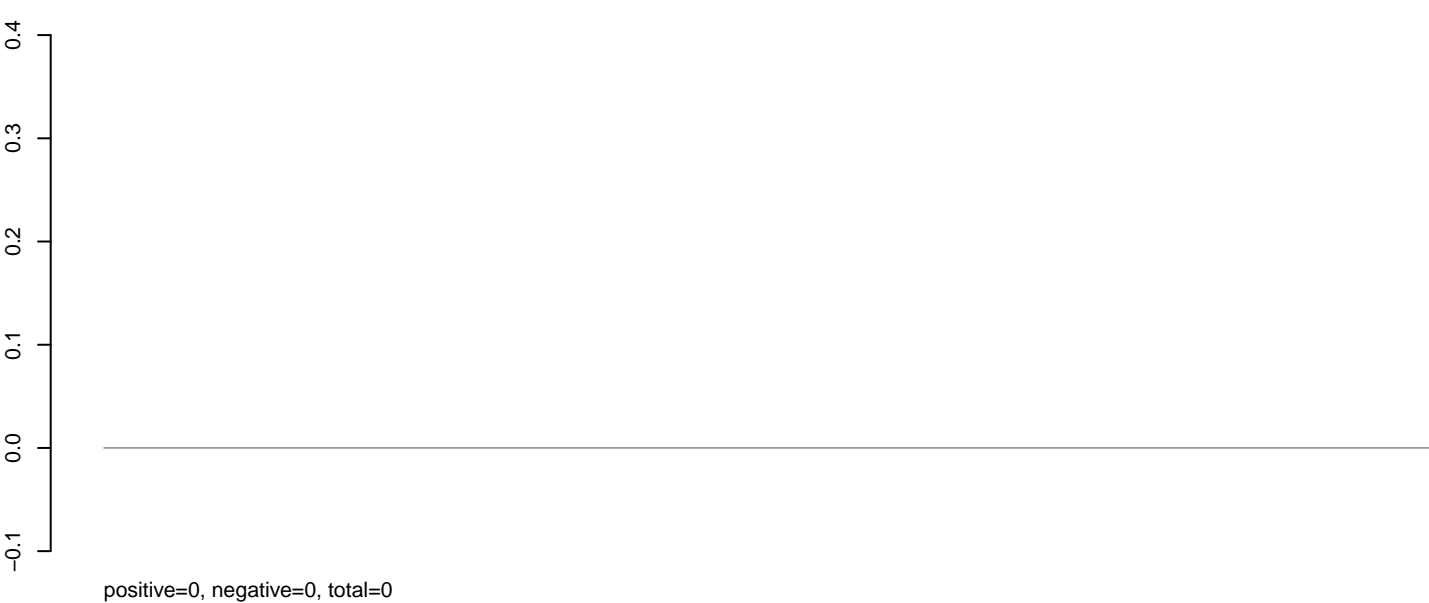
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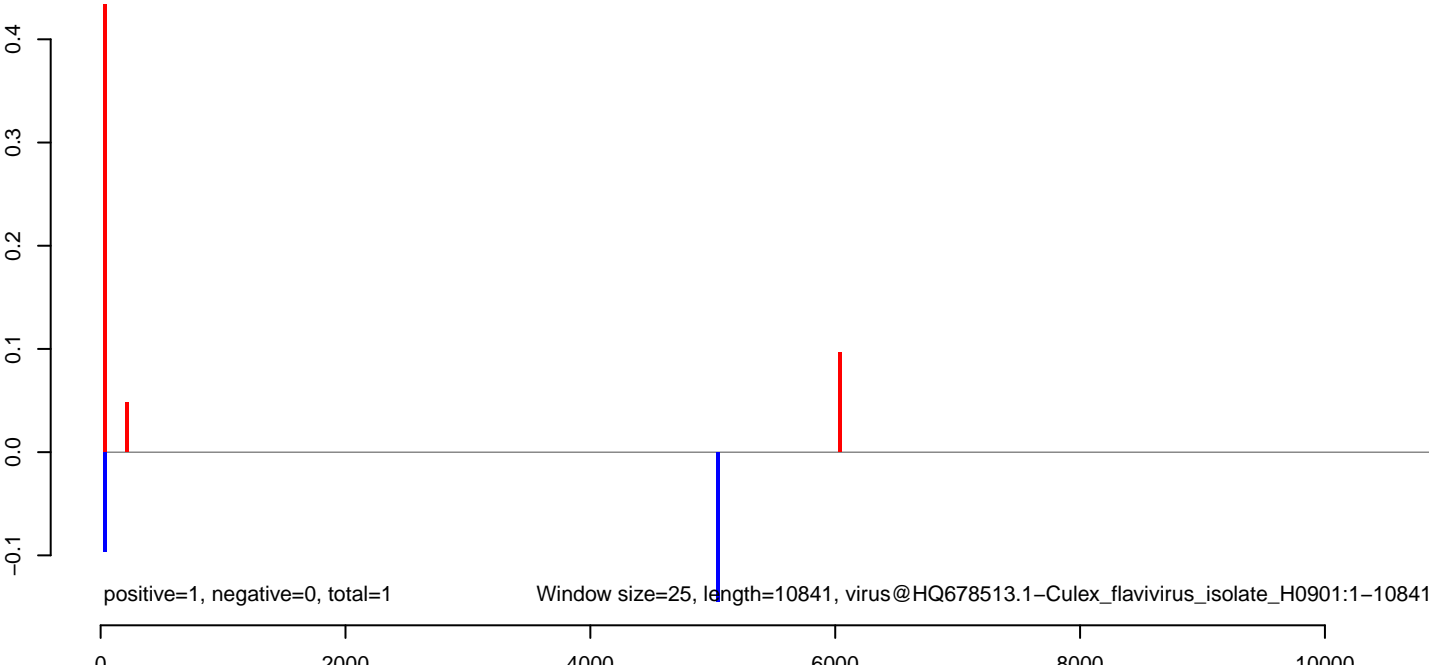
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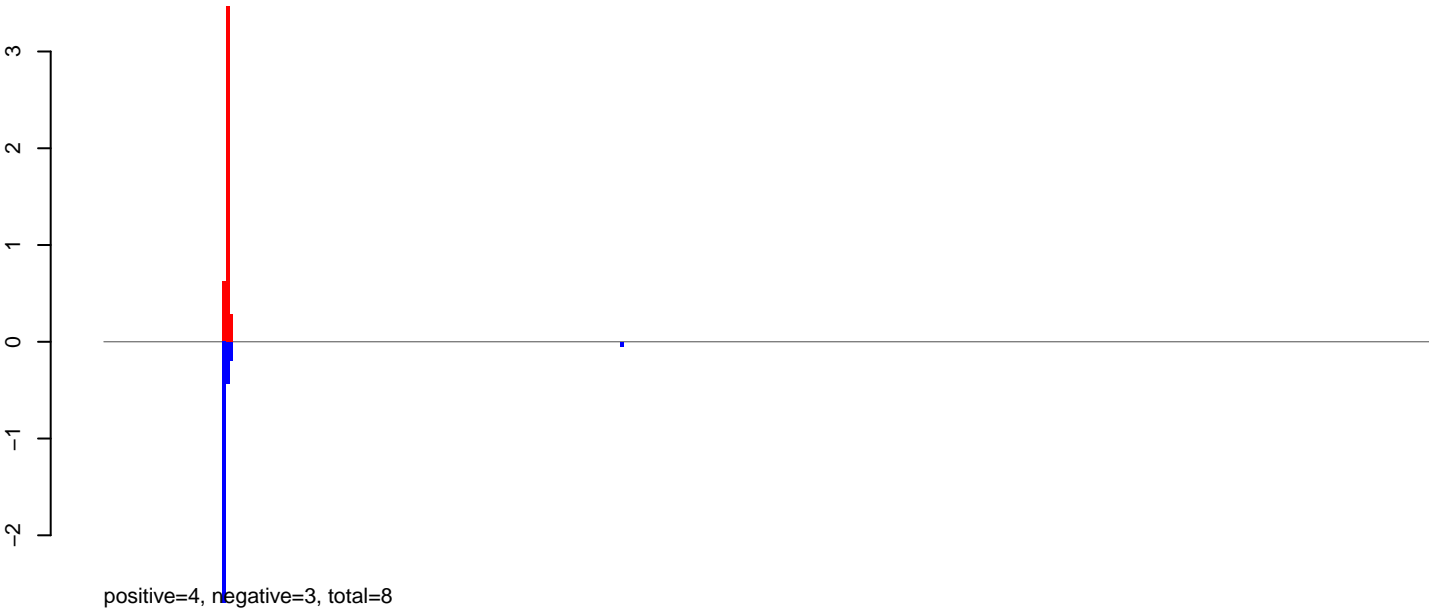
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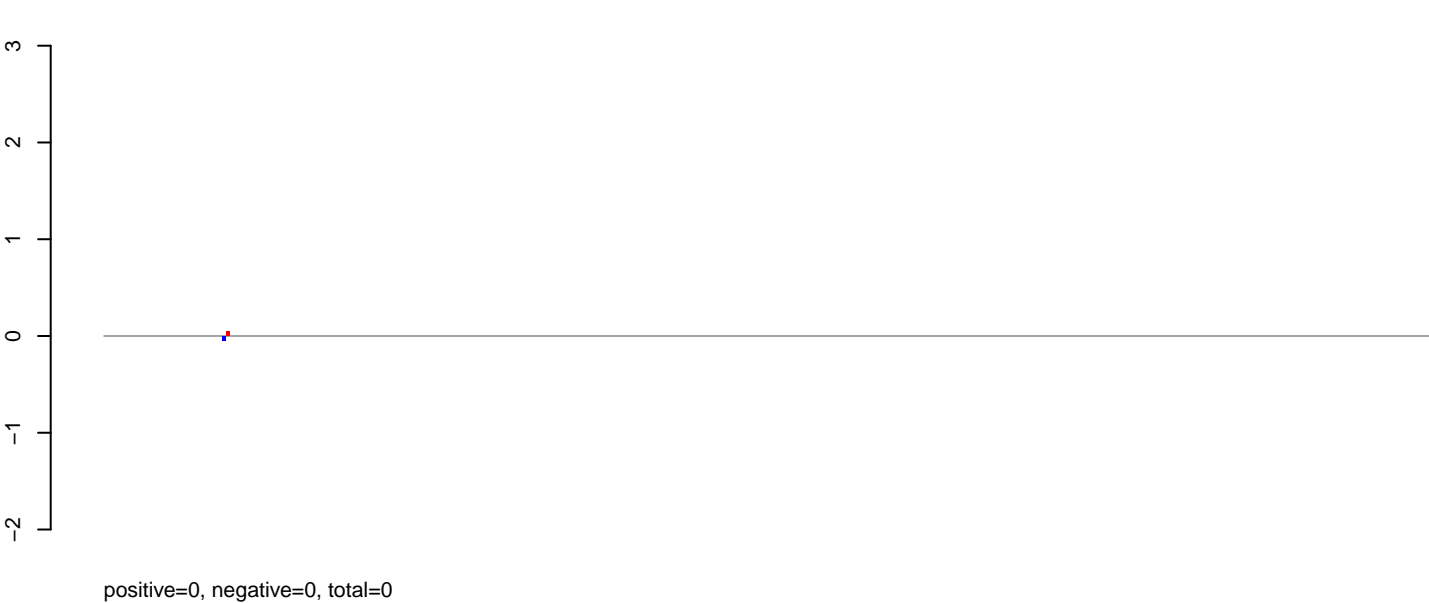
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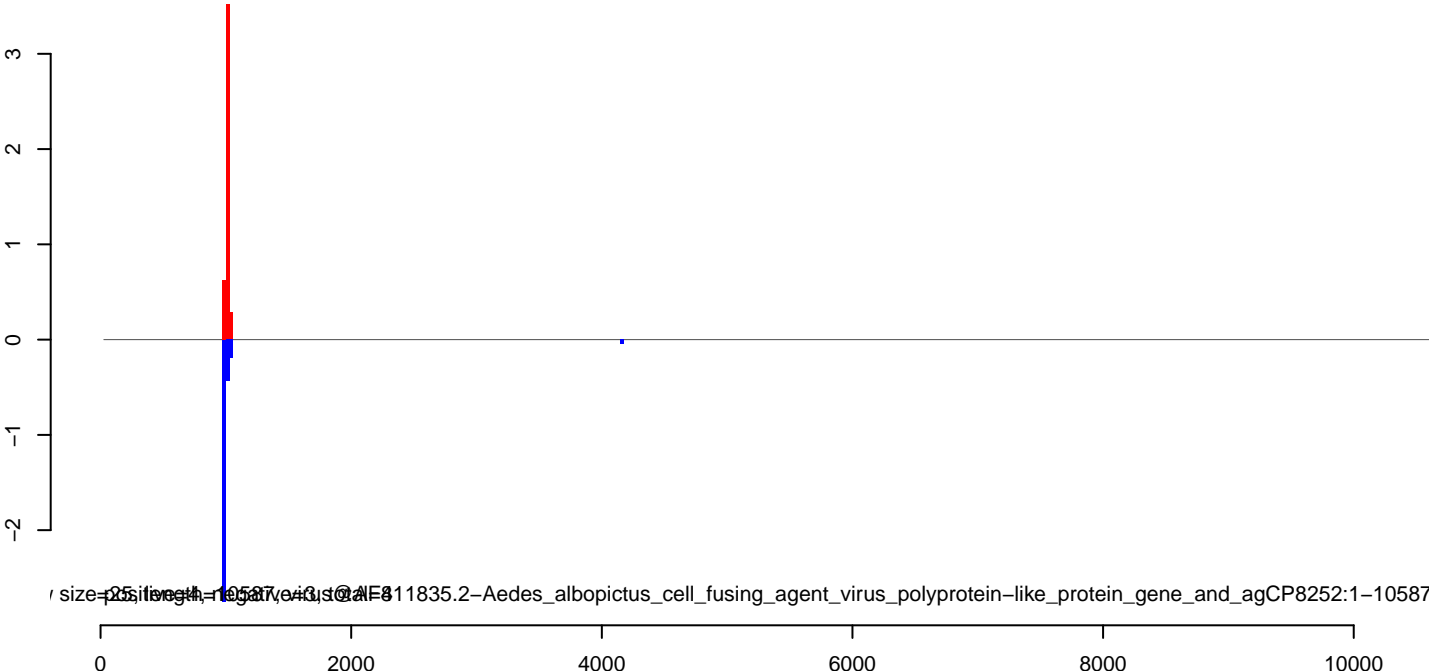
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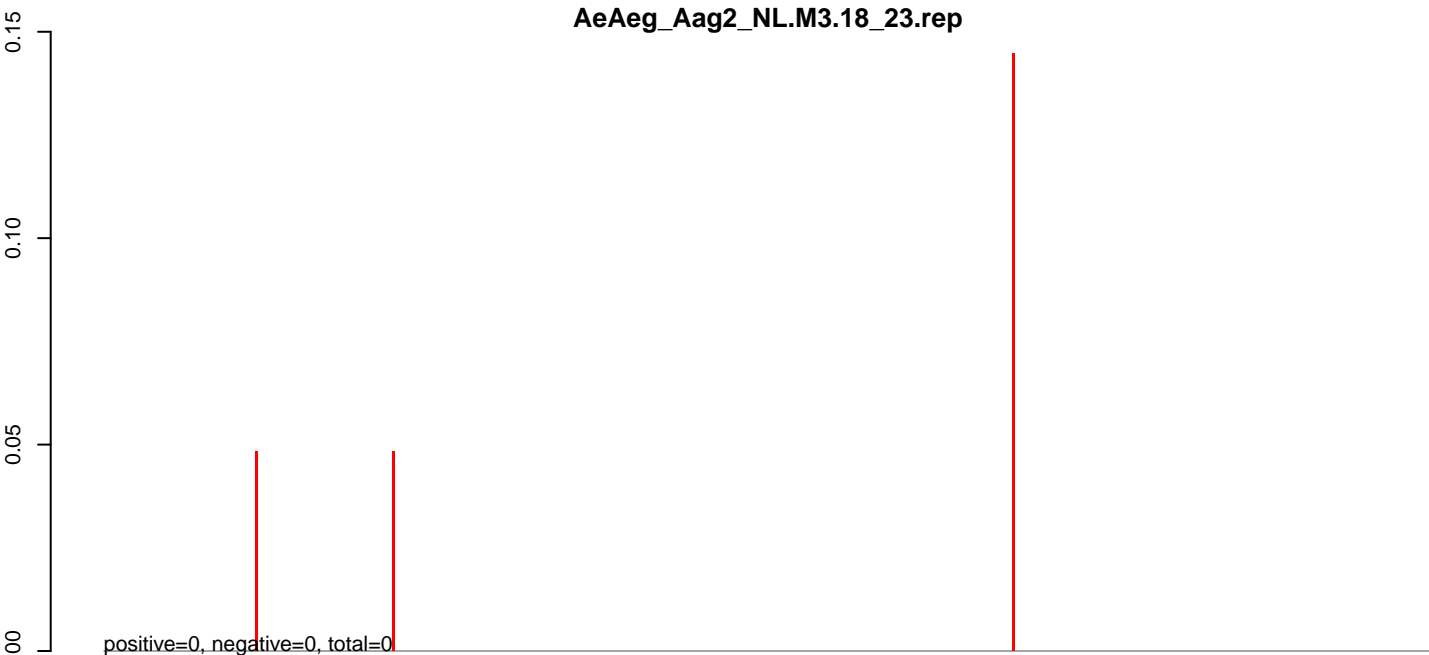
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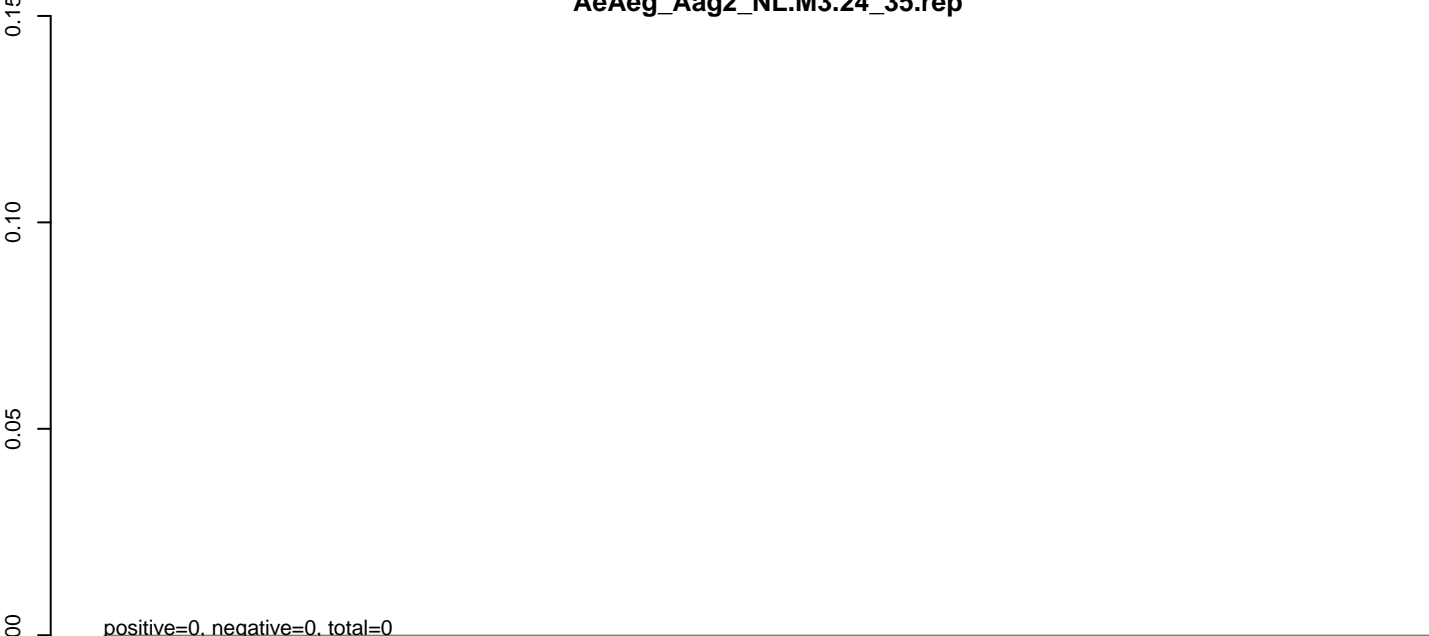
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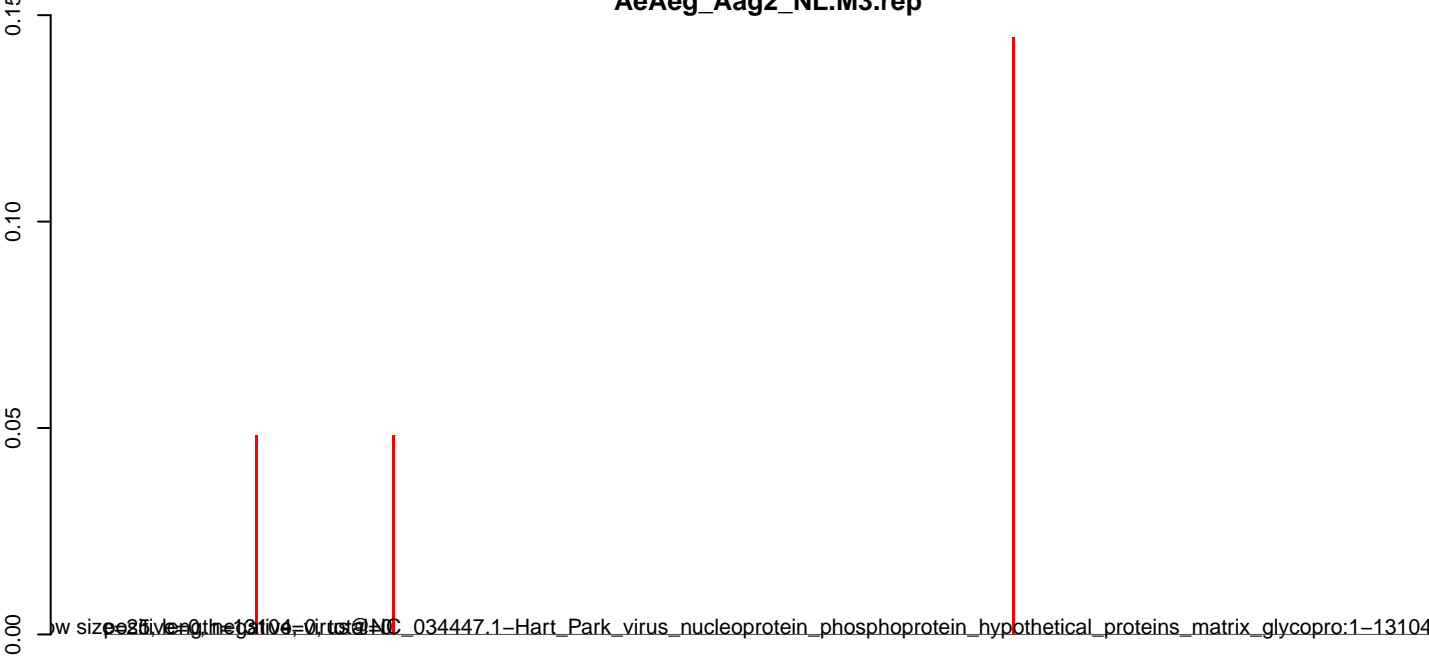
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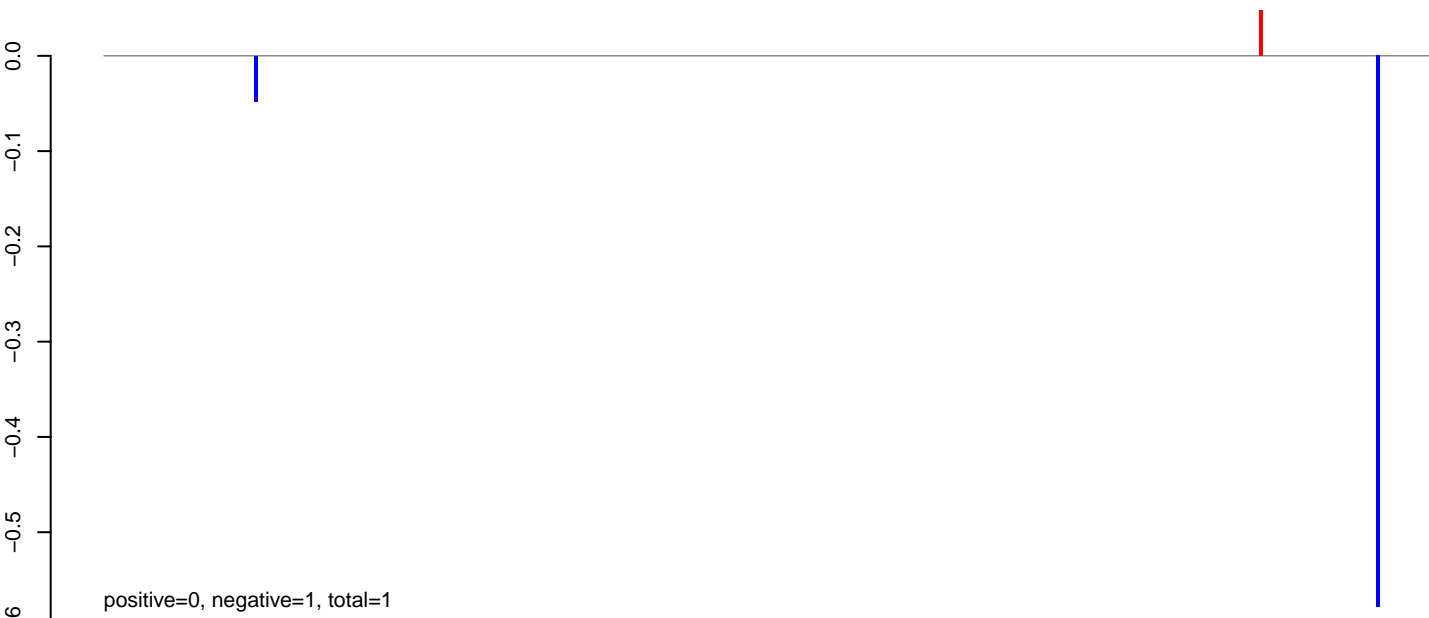


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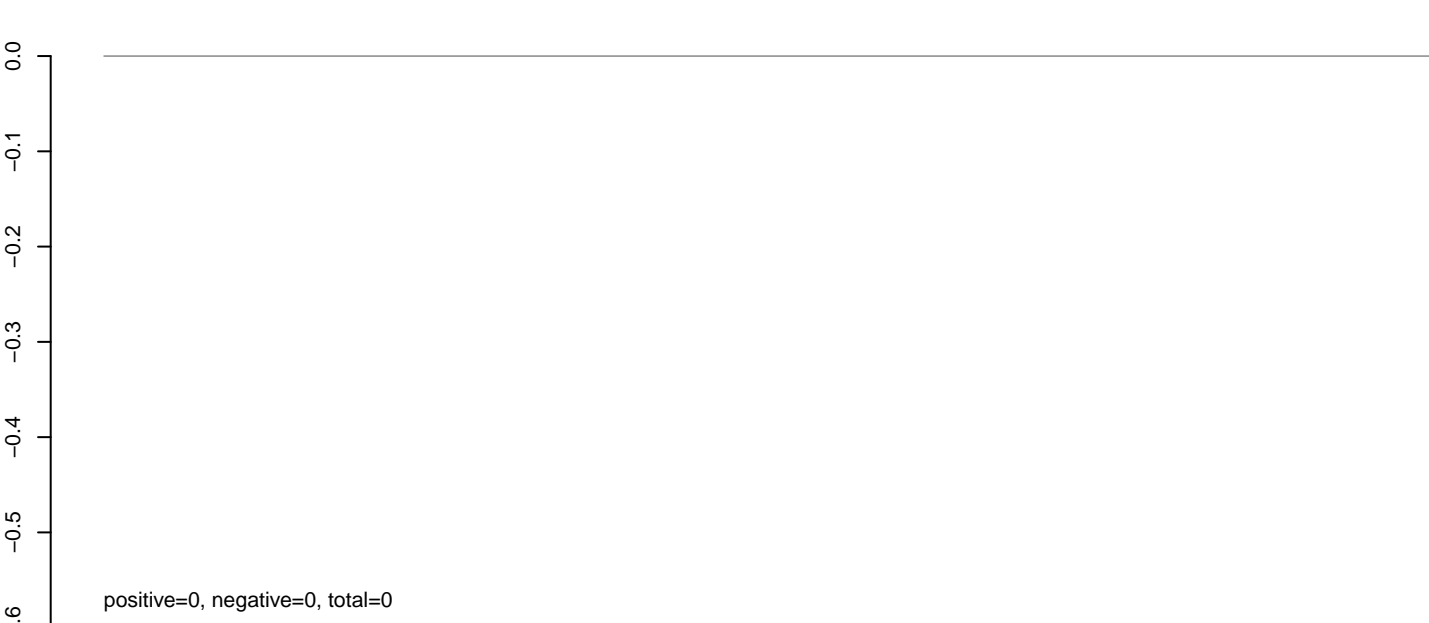


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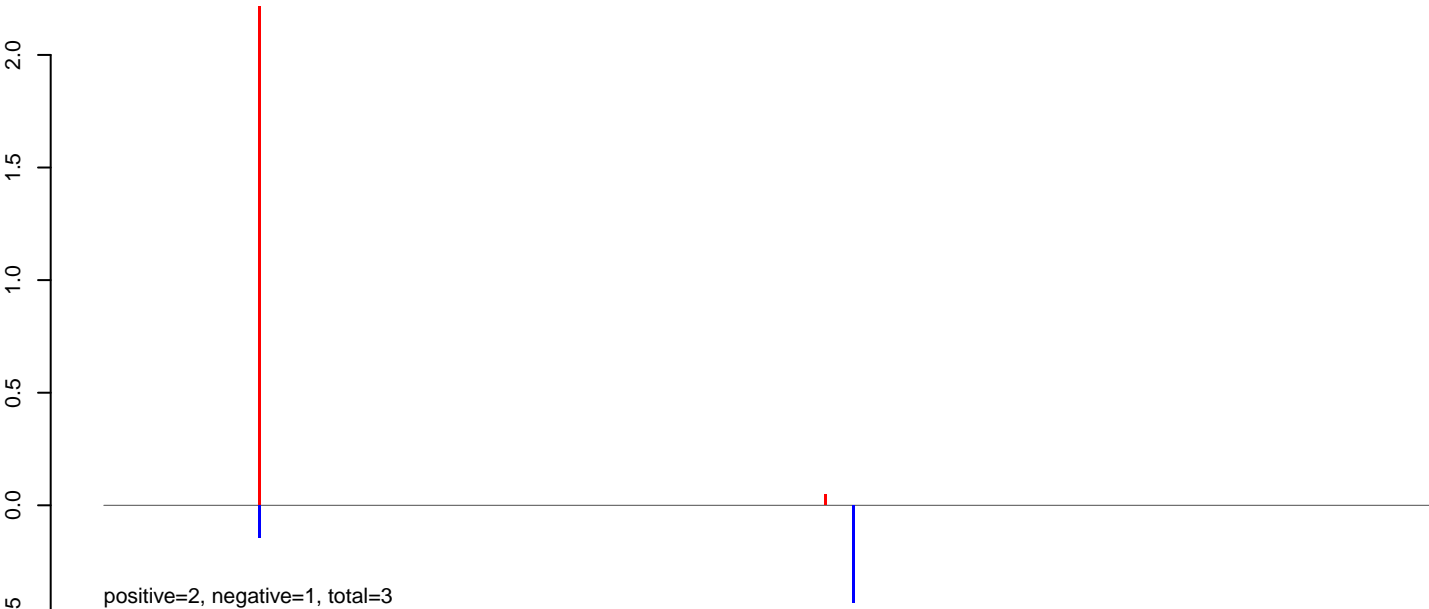
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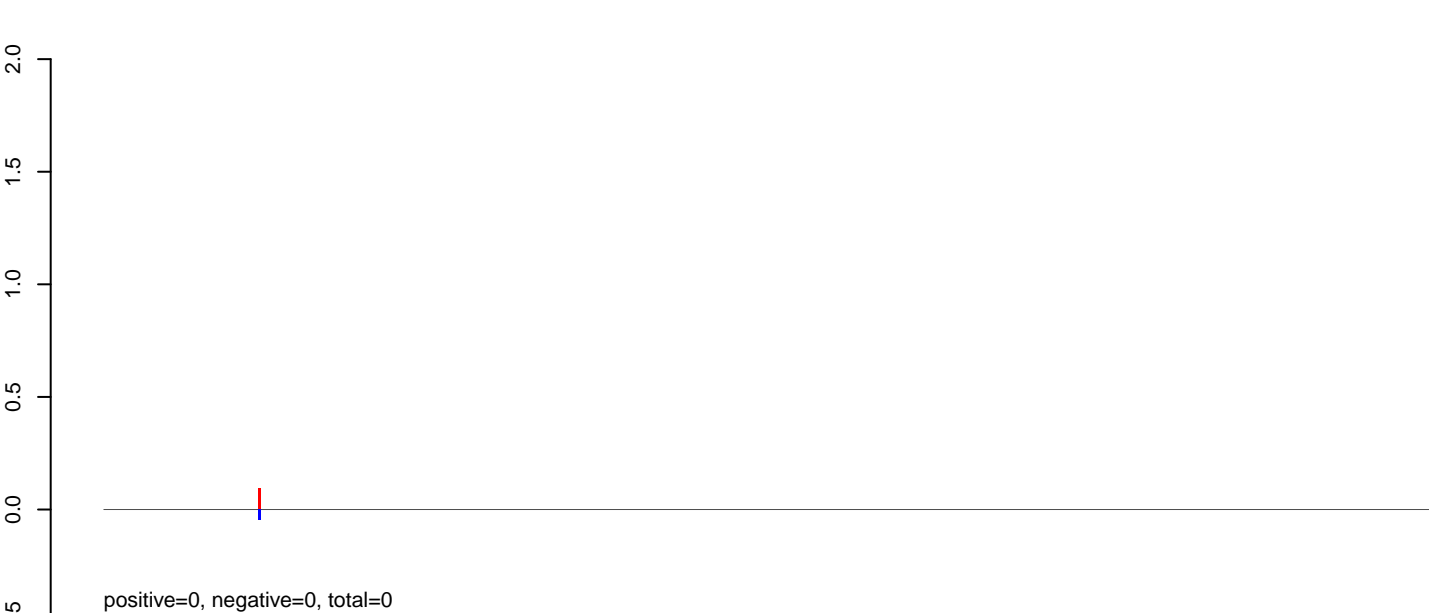
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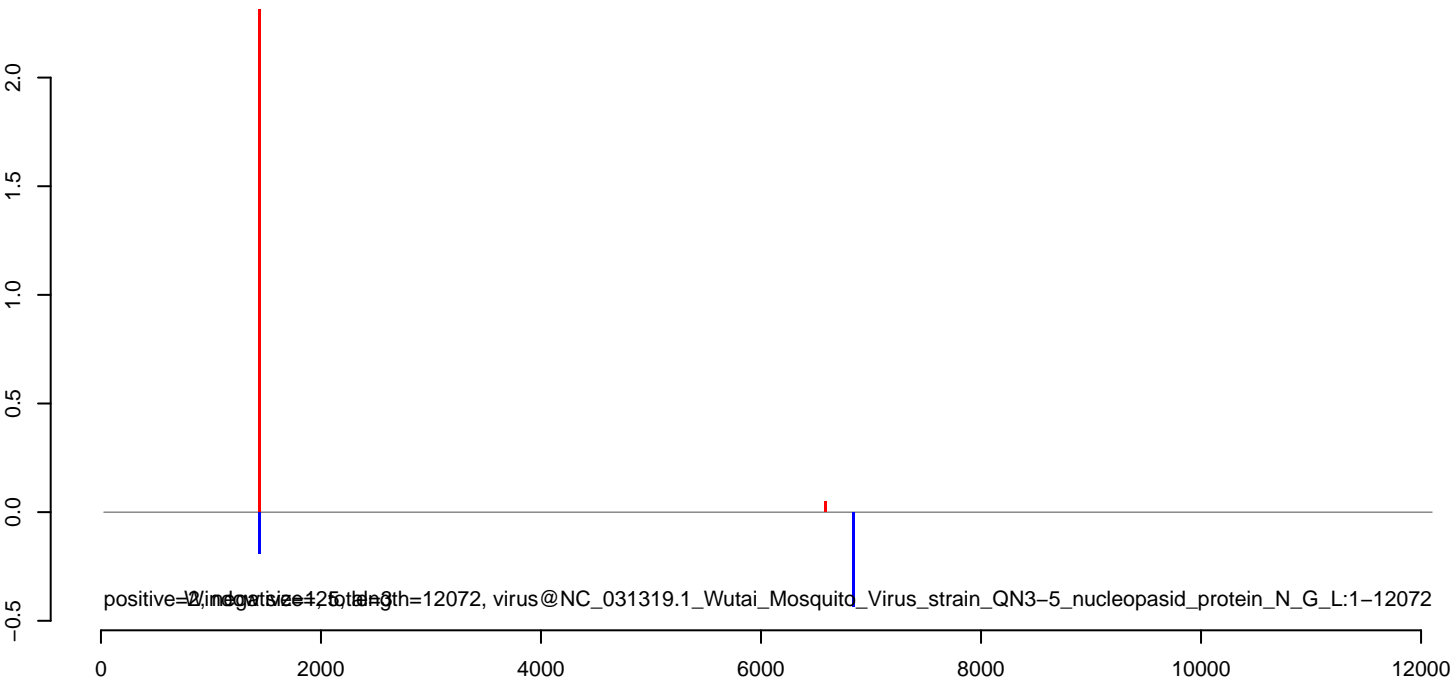
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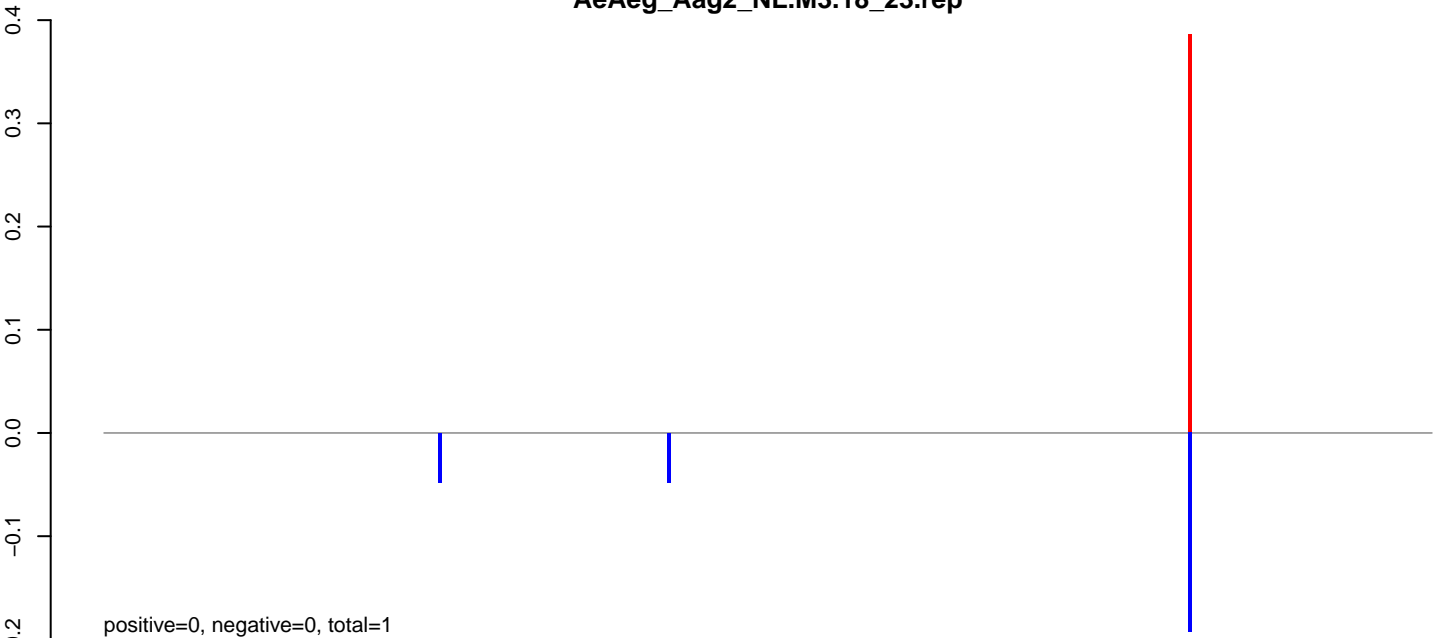
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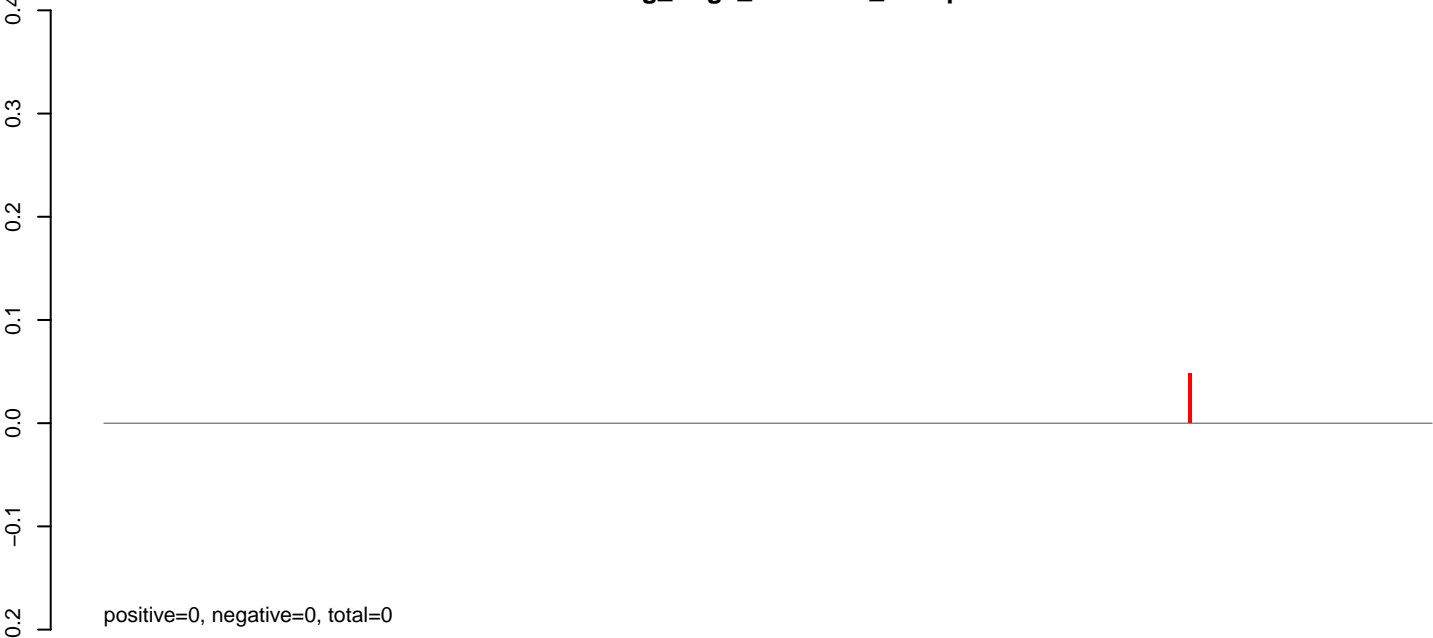
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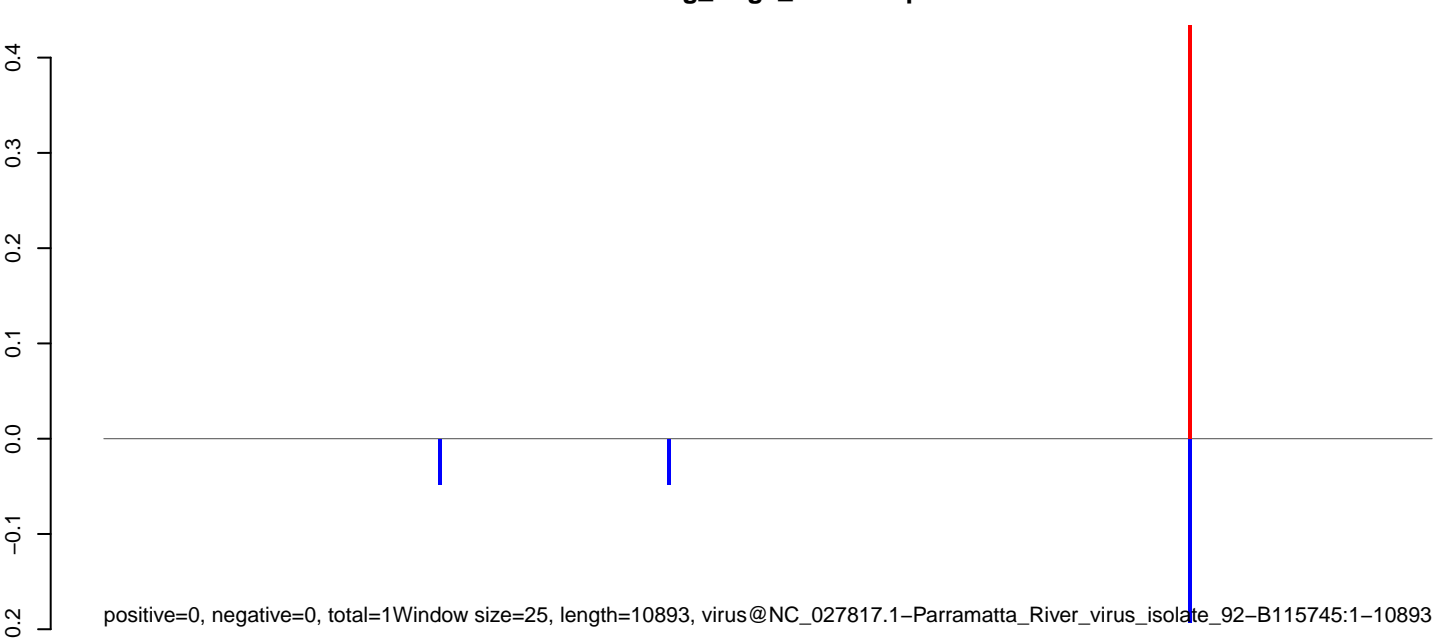
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AeAeg_Aag2_NL.M3.24_35.rep



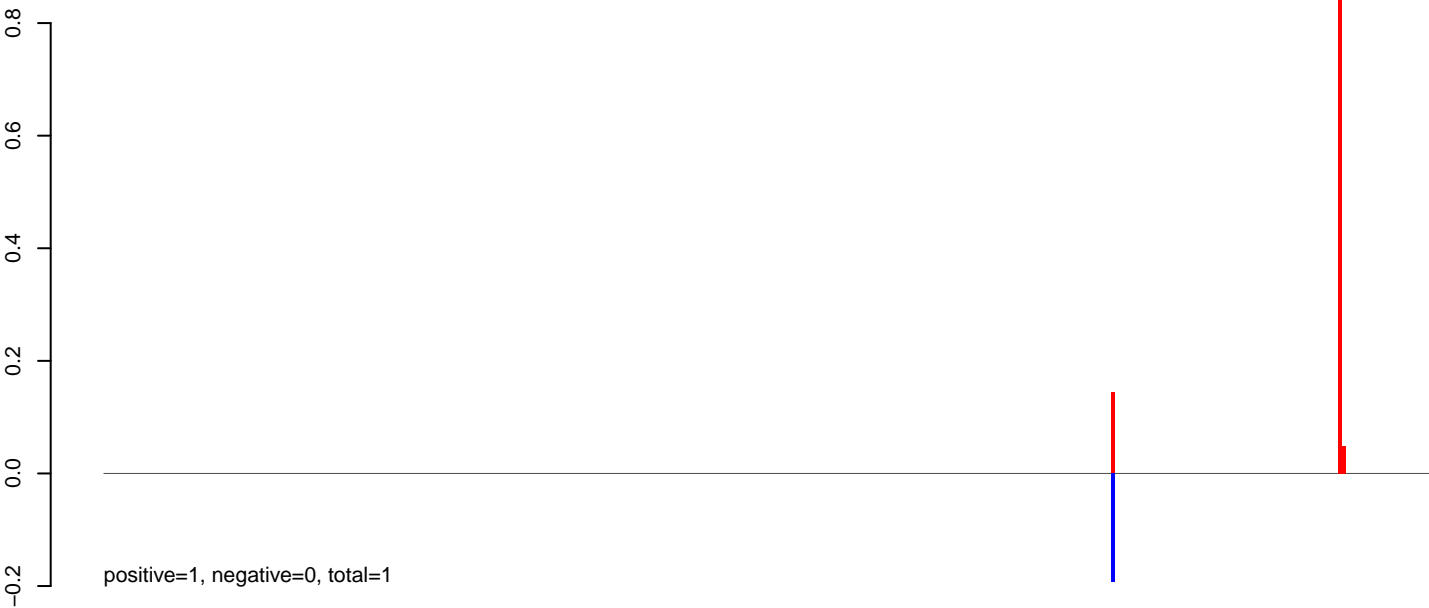
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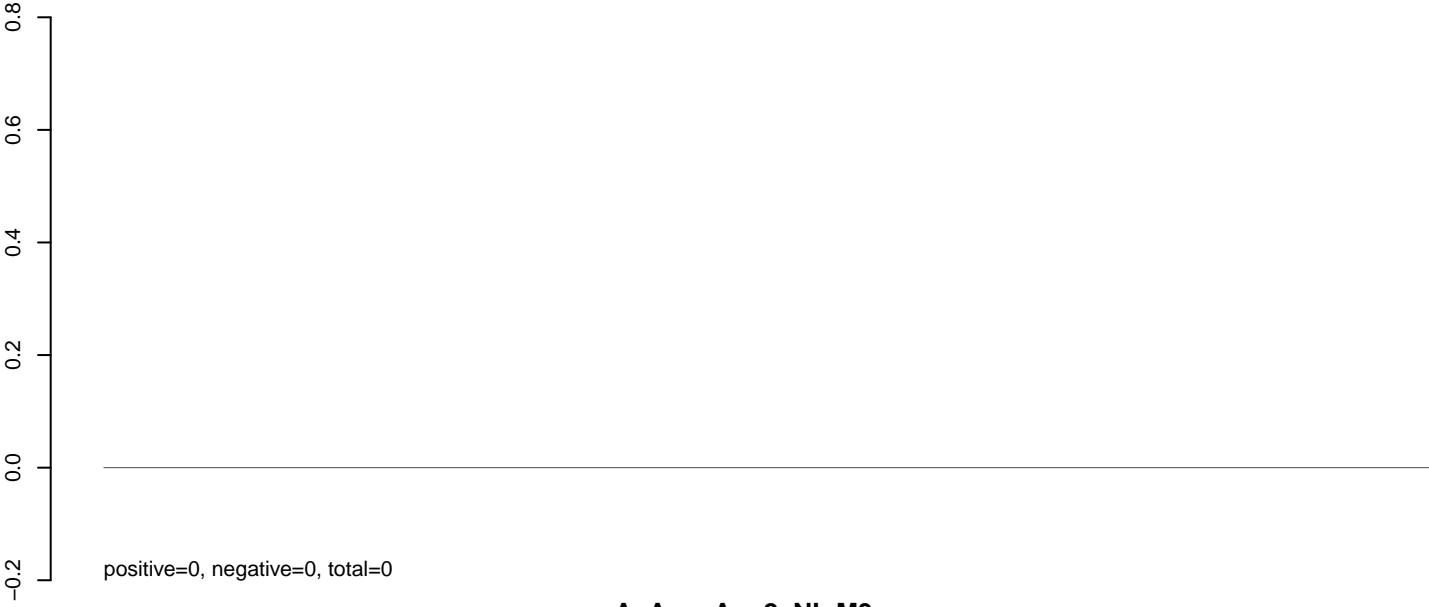
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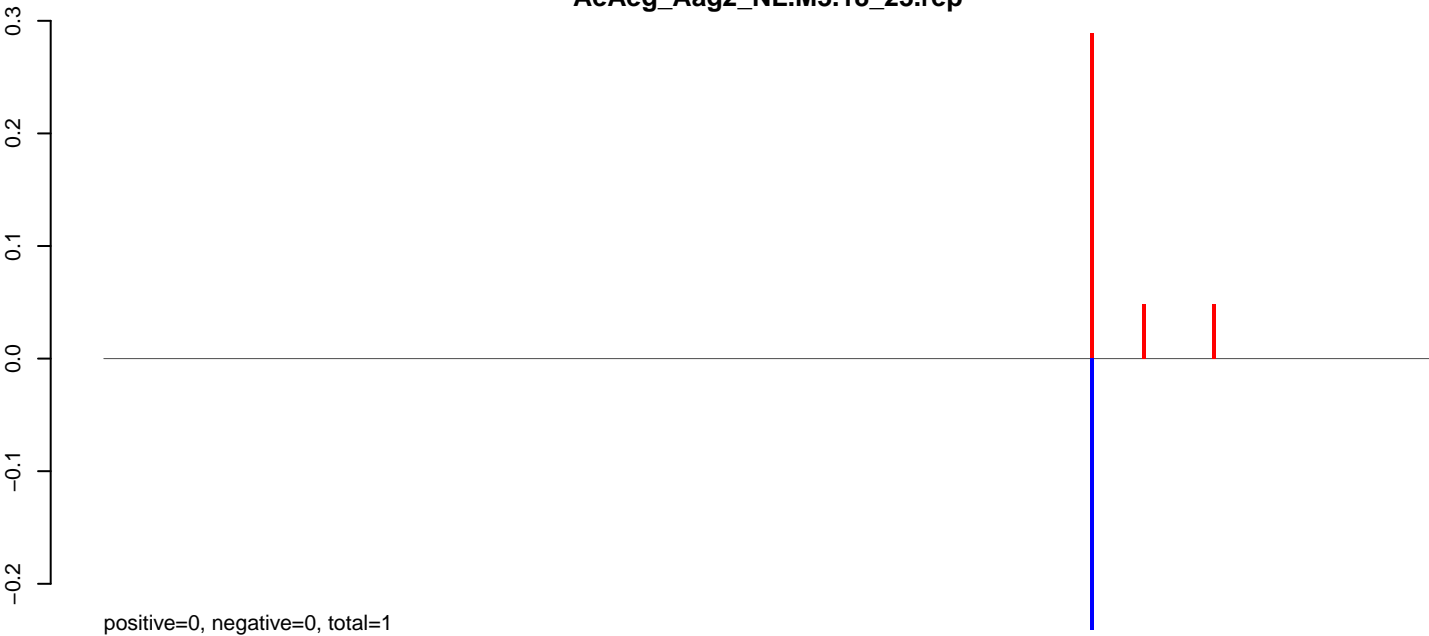
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AeAeg_Aag2_NL.M3.rep



AeAeg_Aag2_NL.M3.18_23.rep



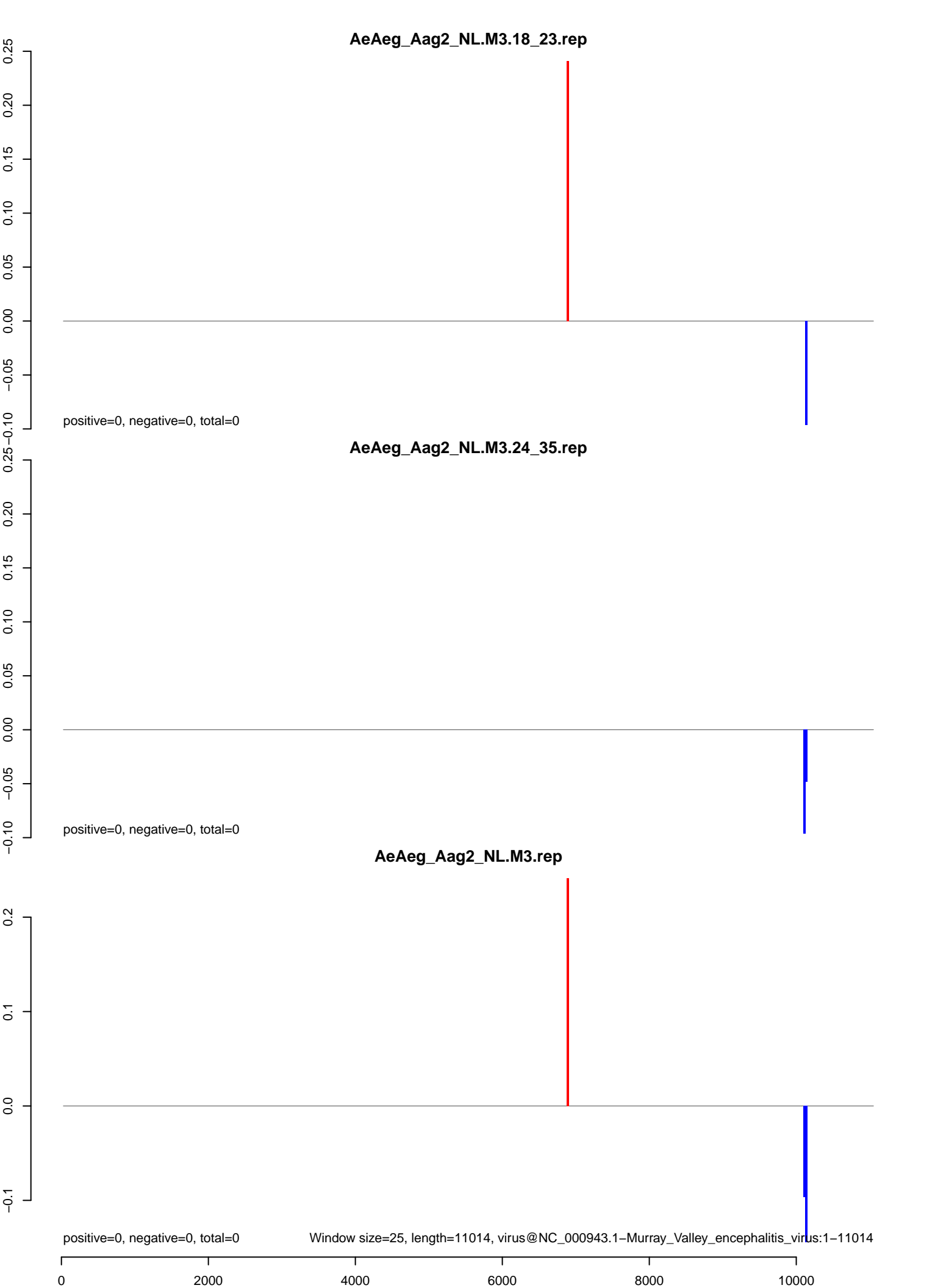
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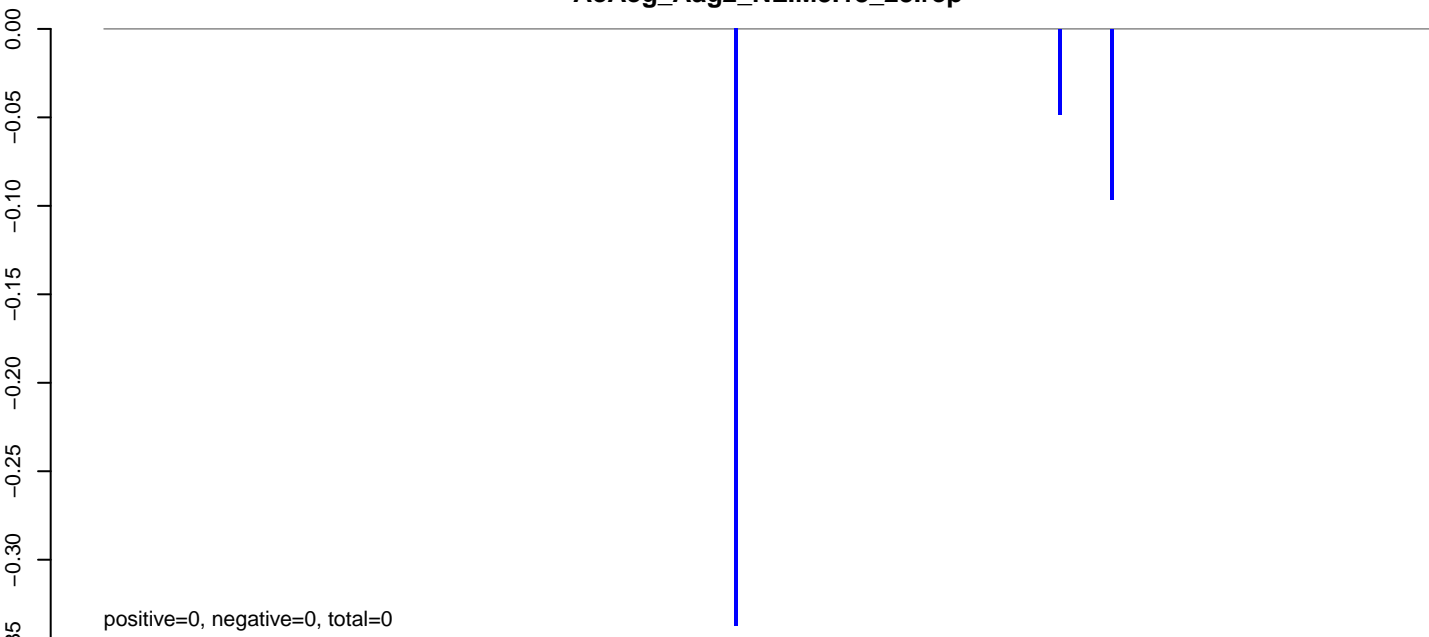
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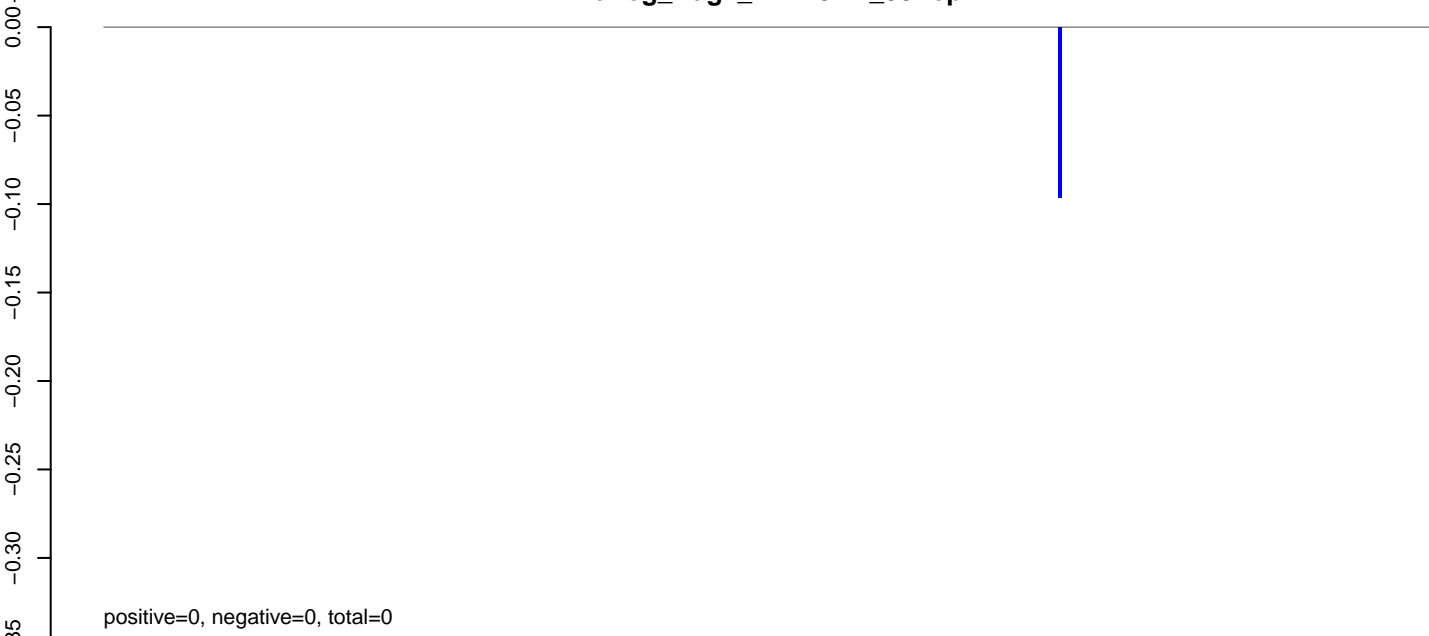
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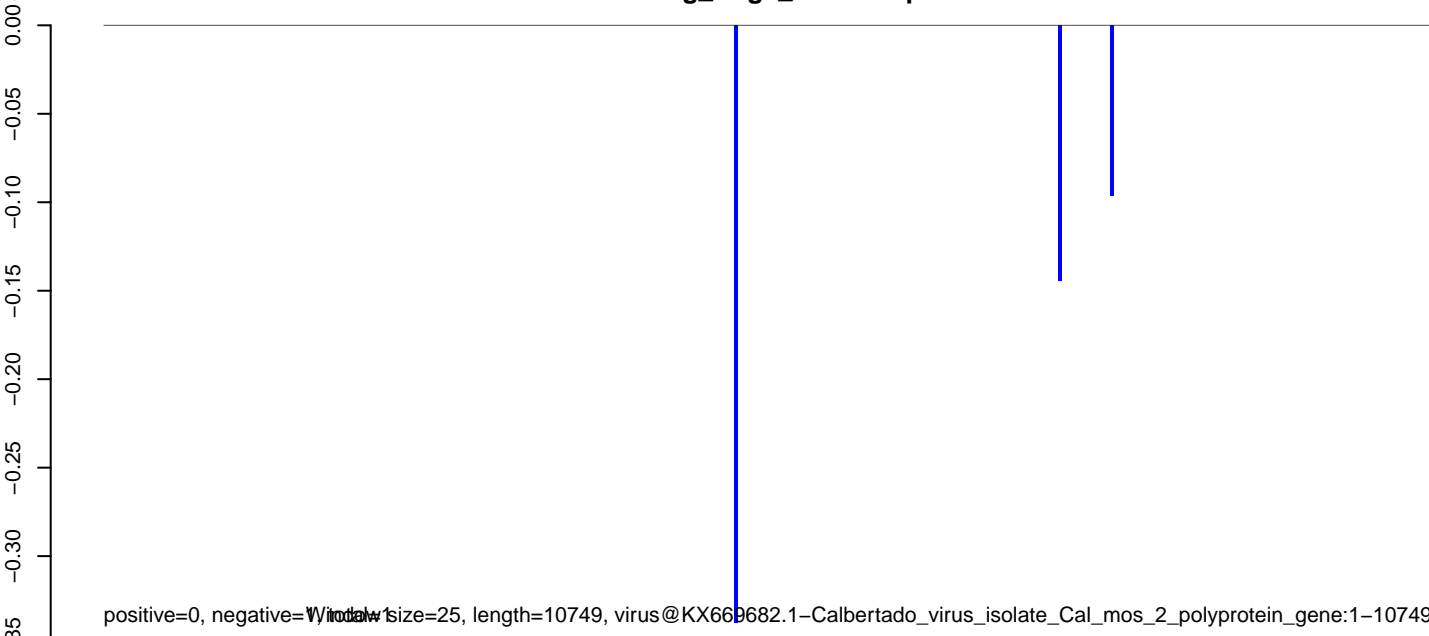
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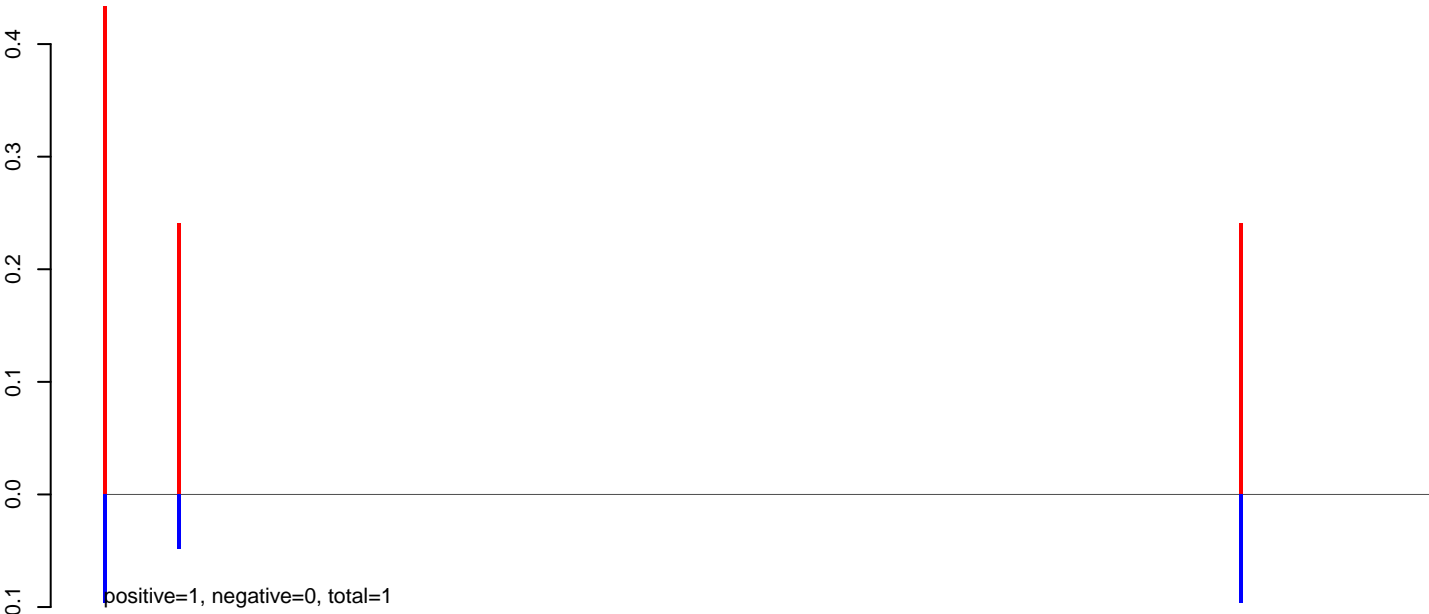
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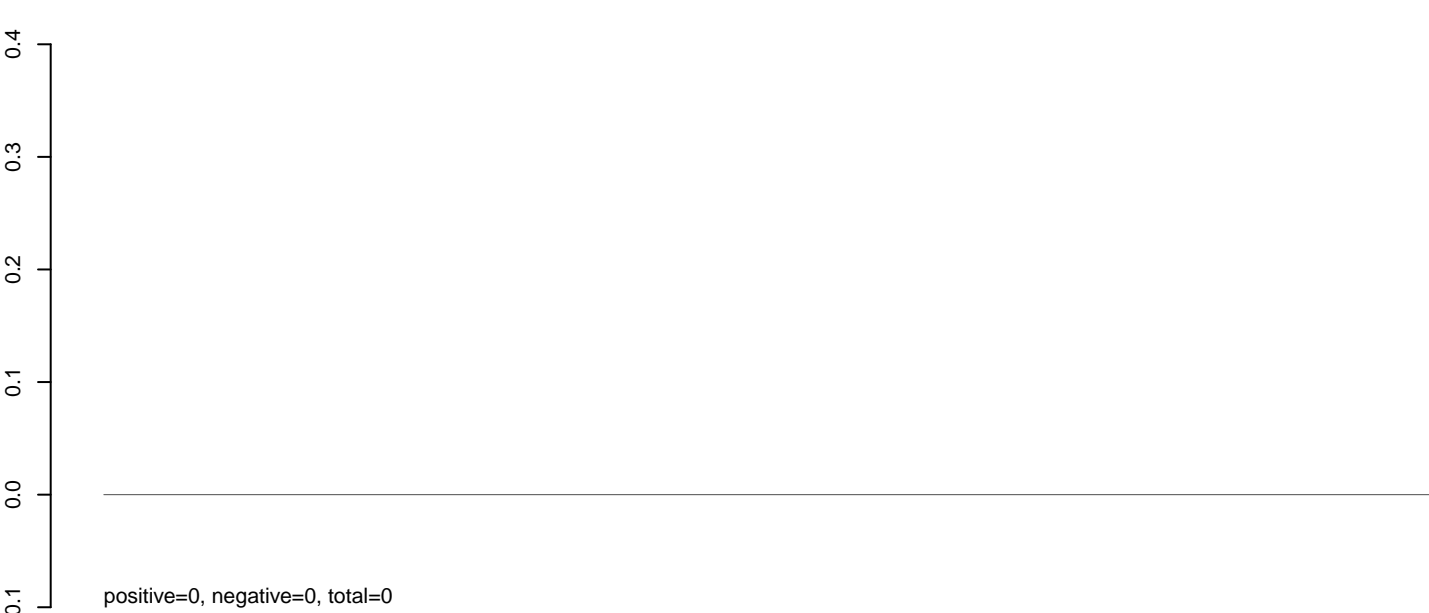
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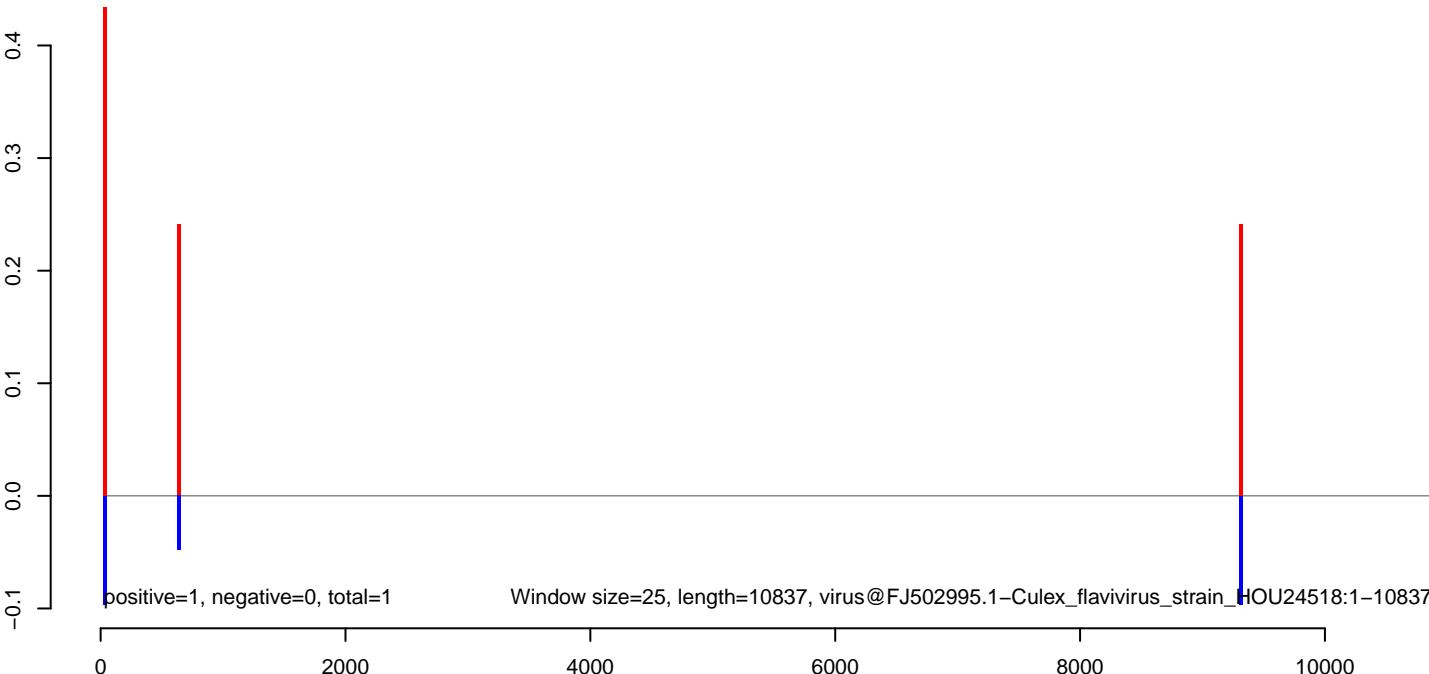
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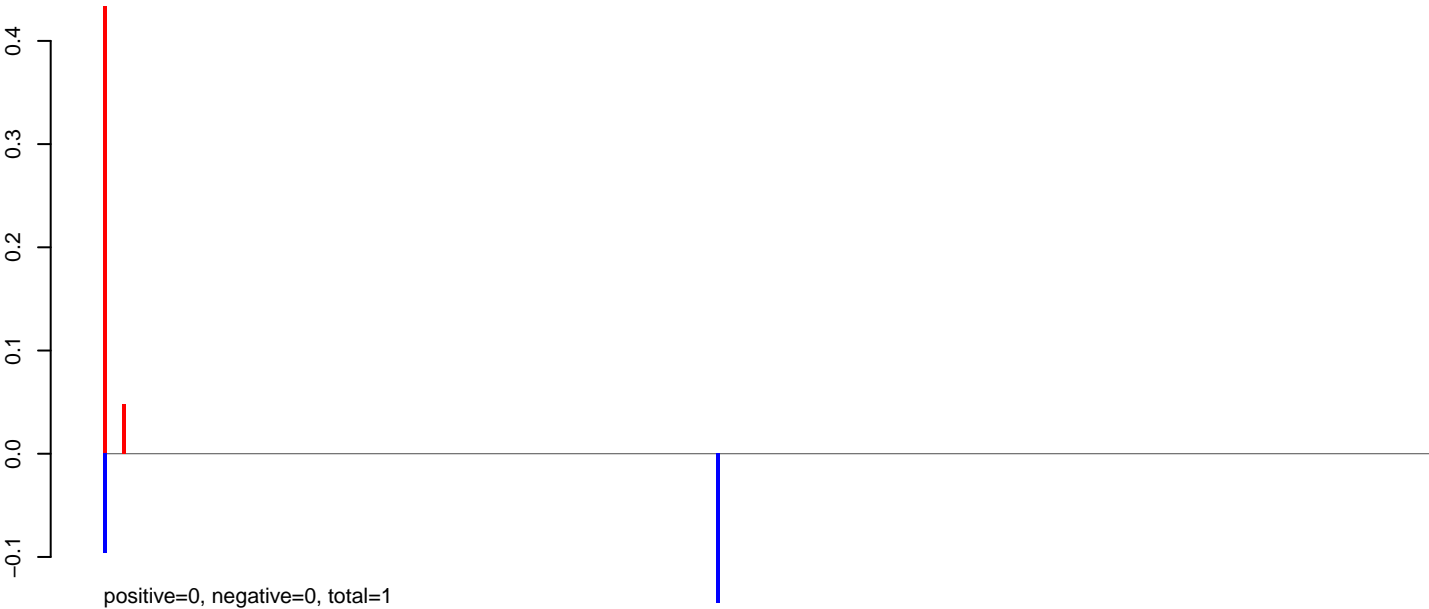
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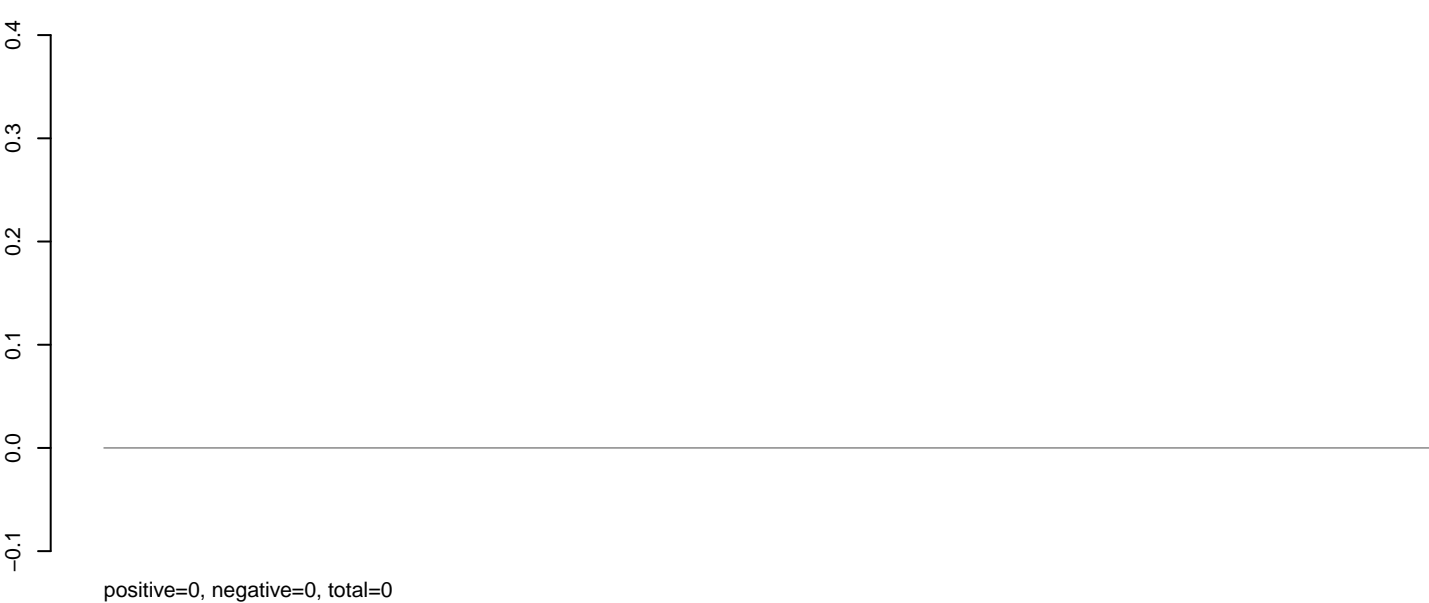
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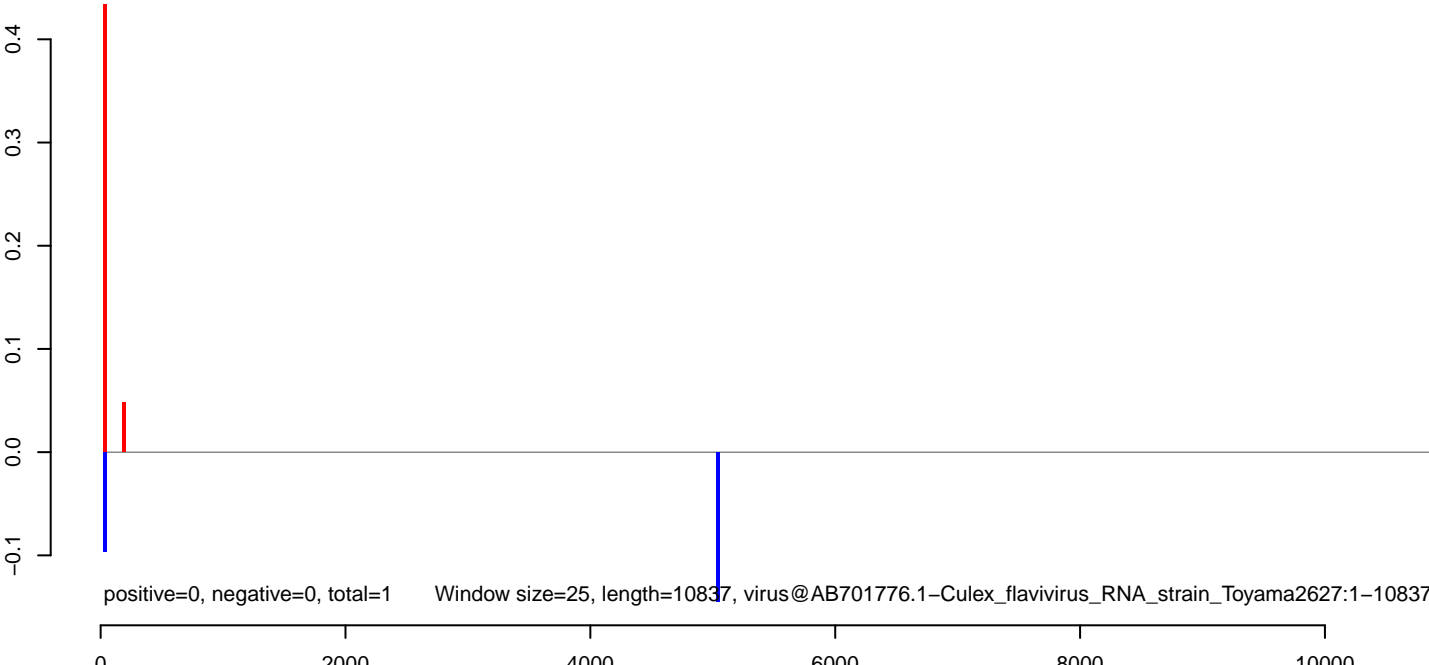
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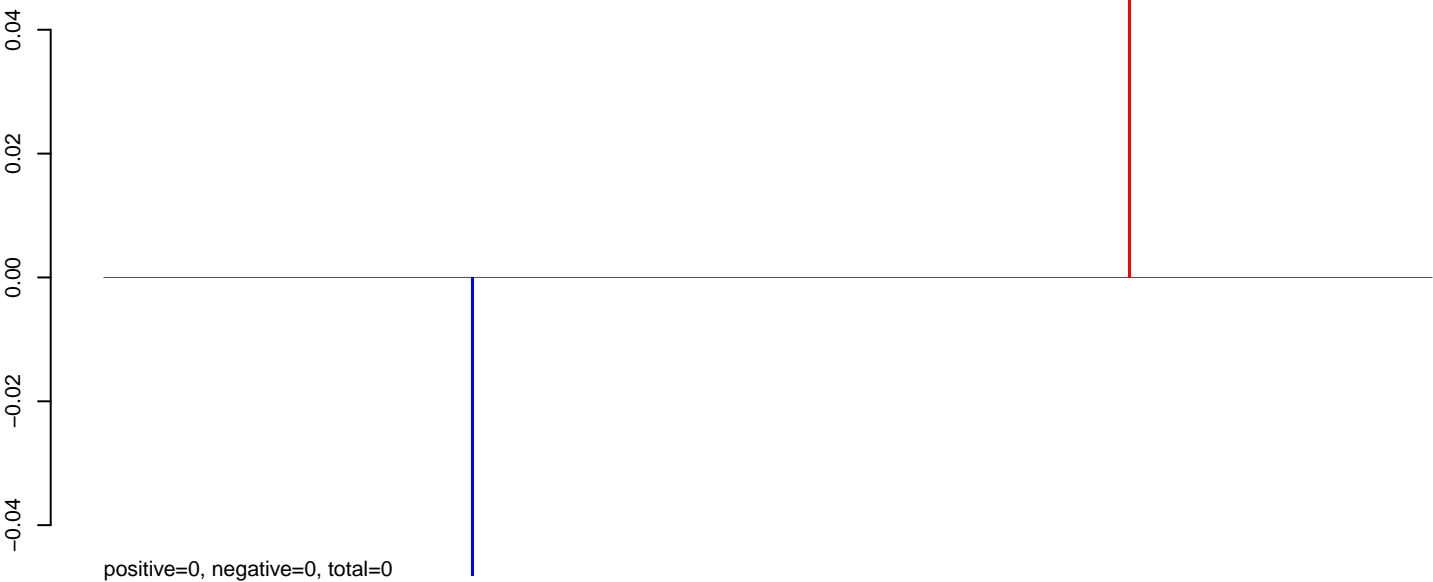
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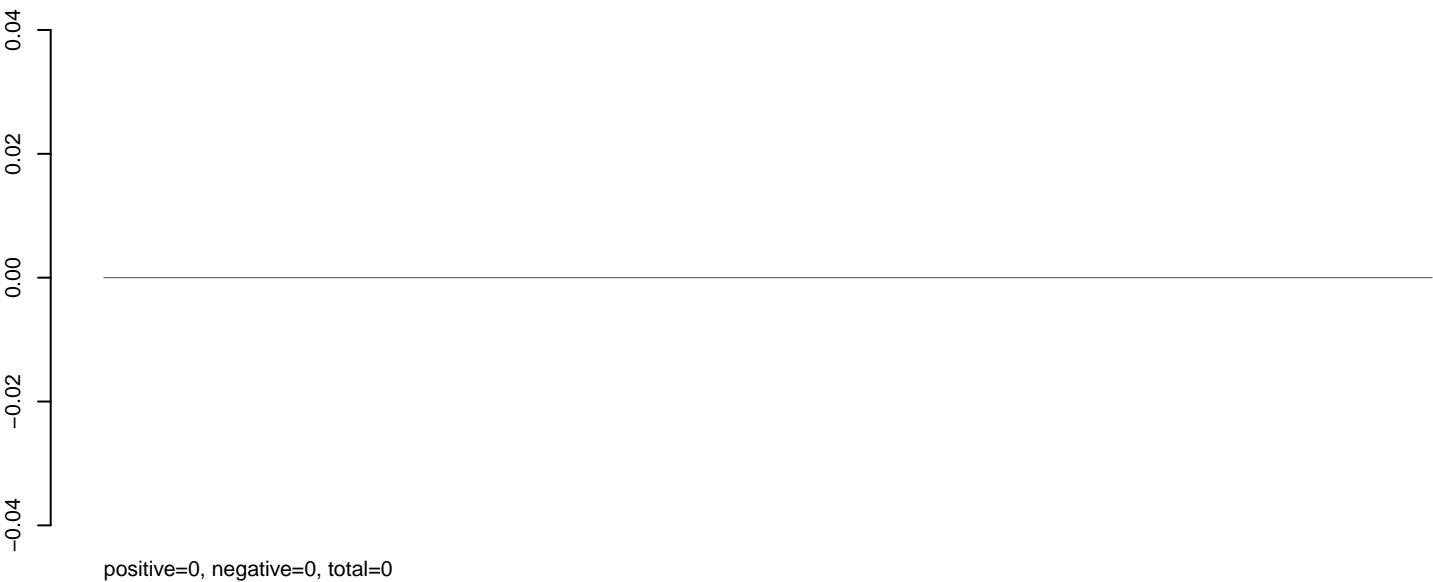
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AeAeg_Aag2_NL.M3.24_35.rep

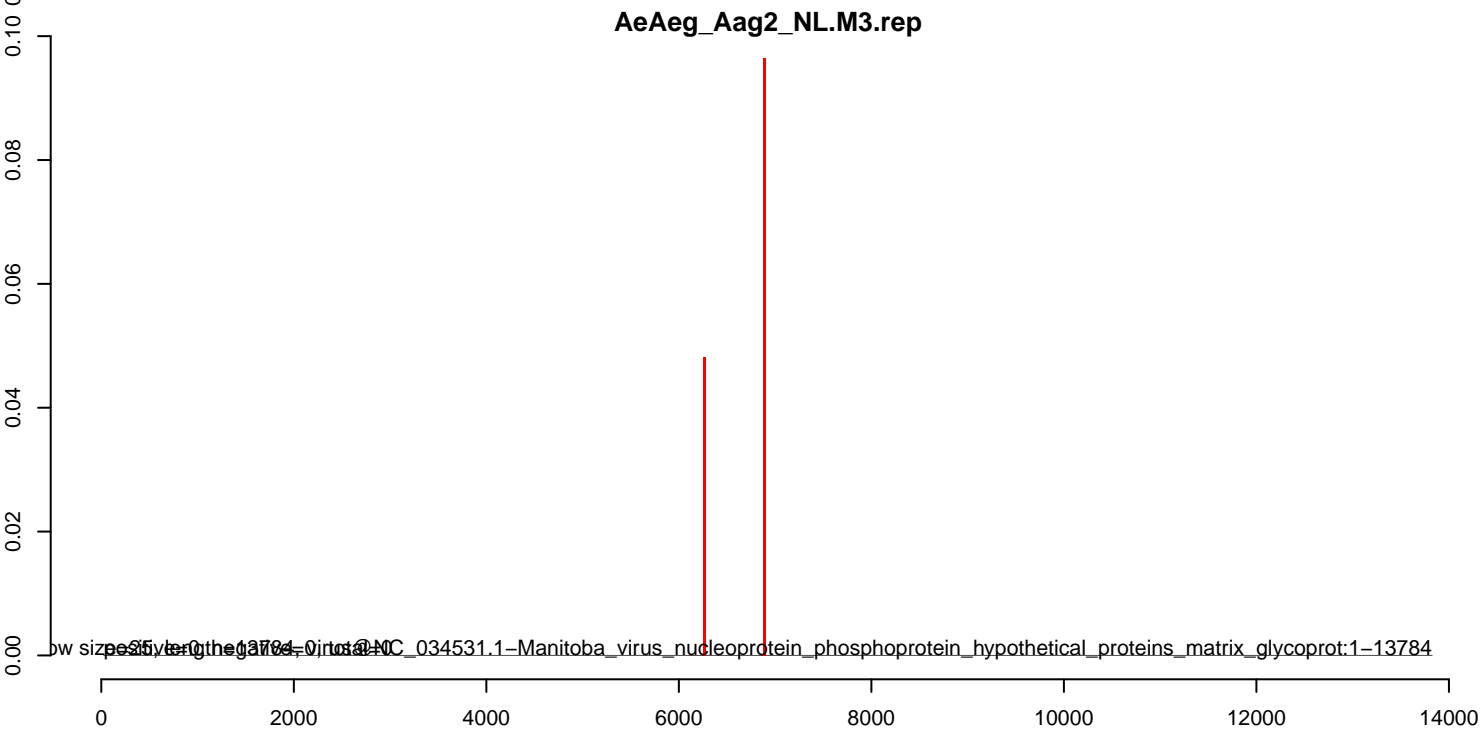
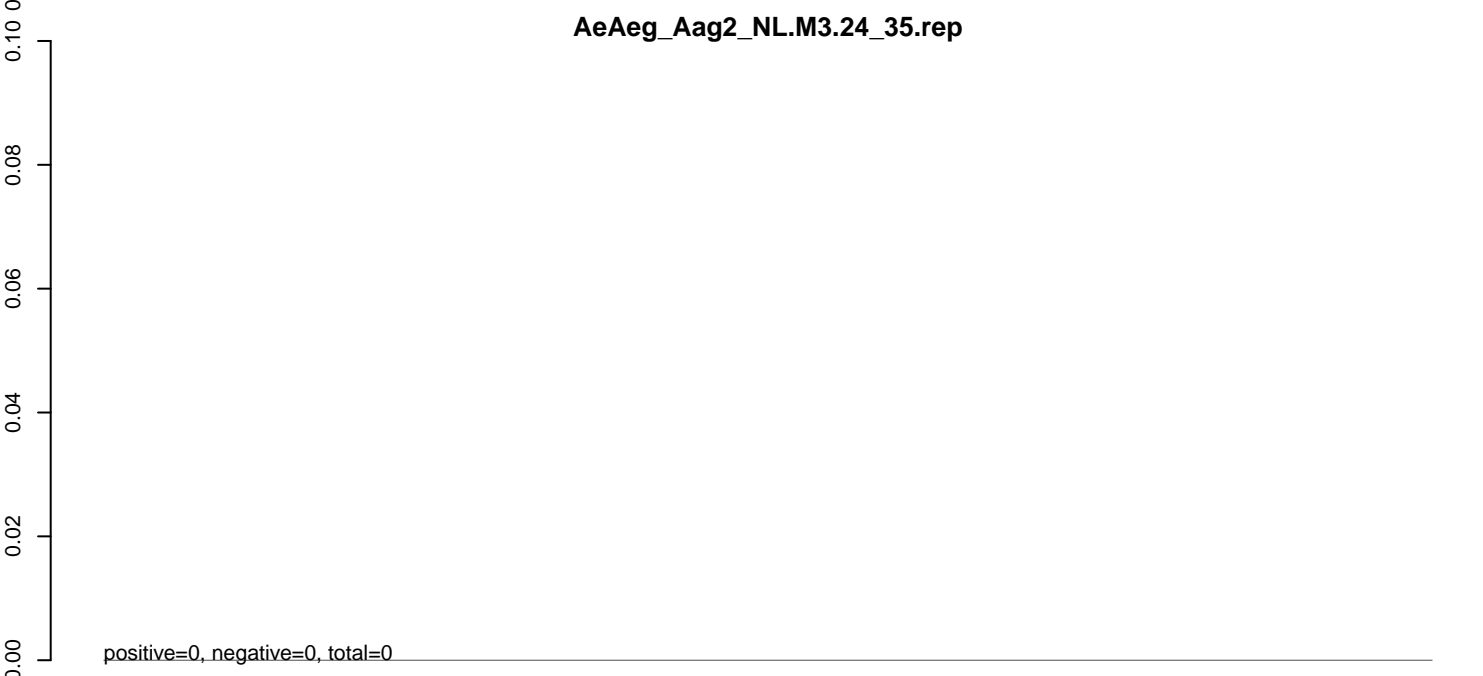
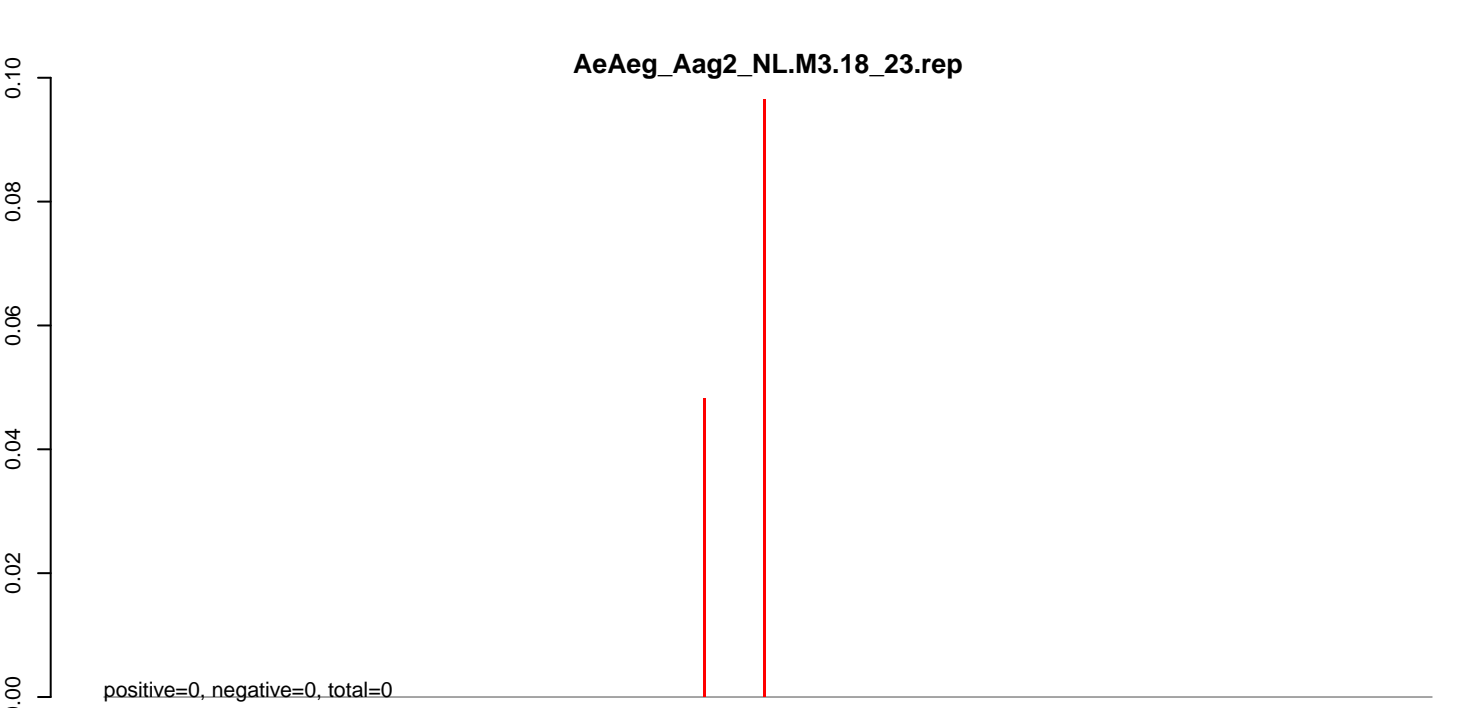


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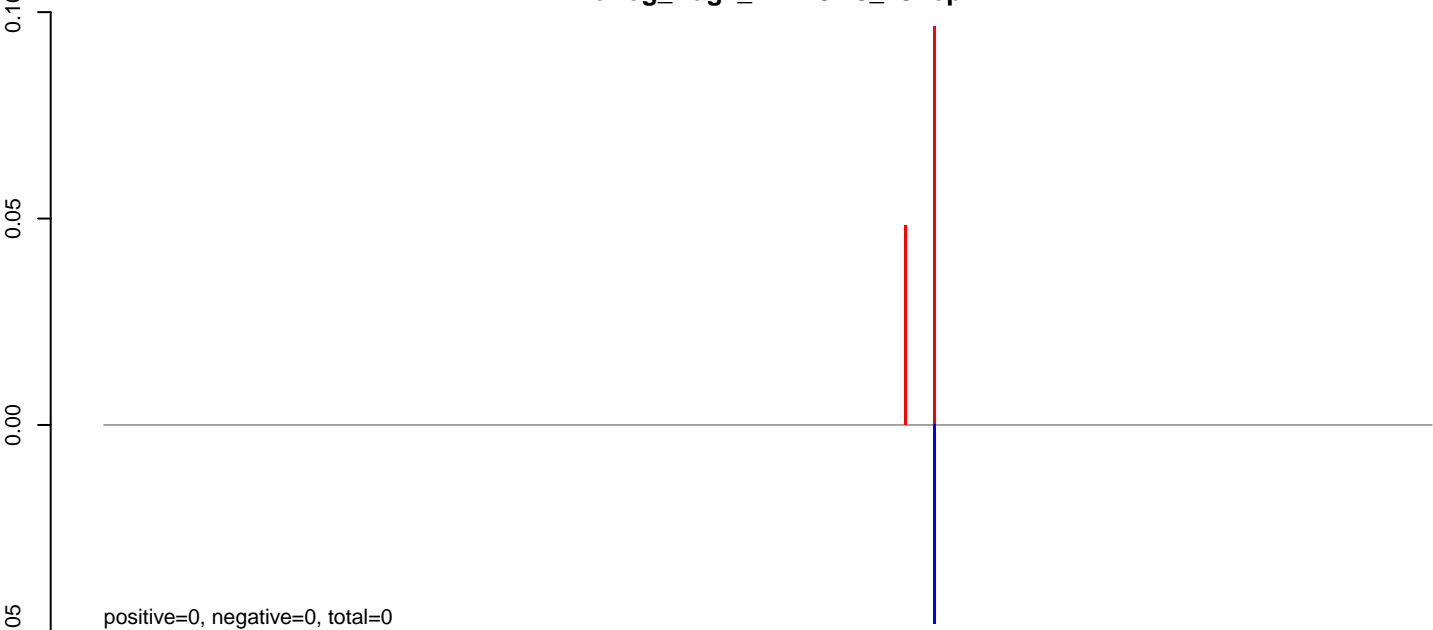
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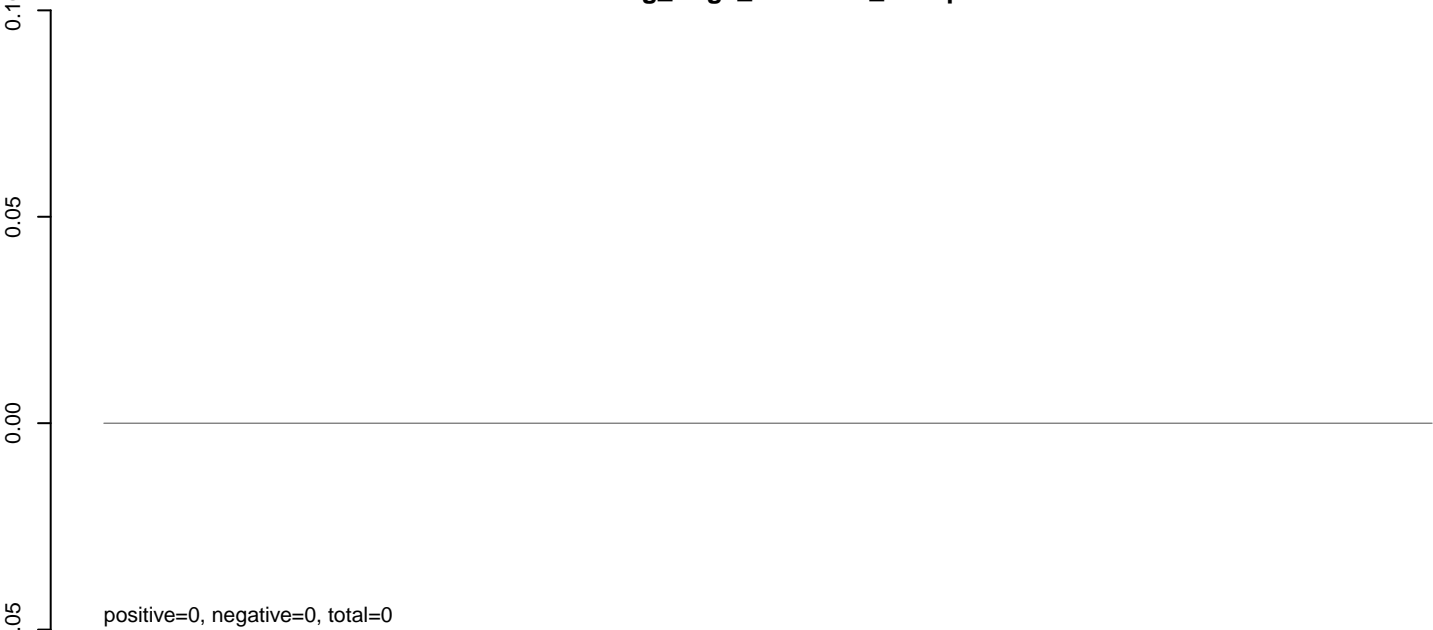


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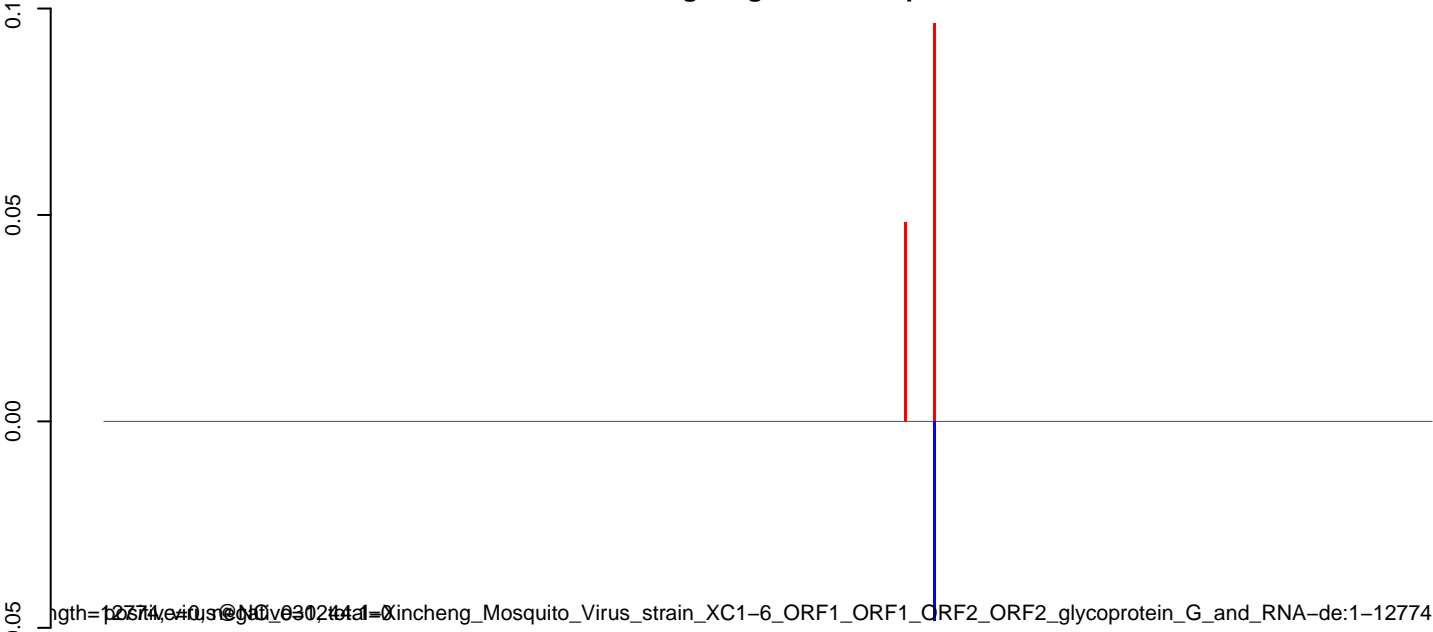
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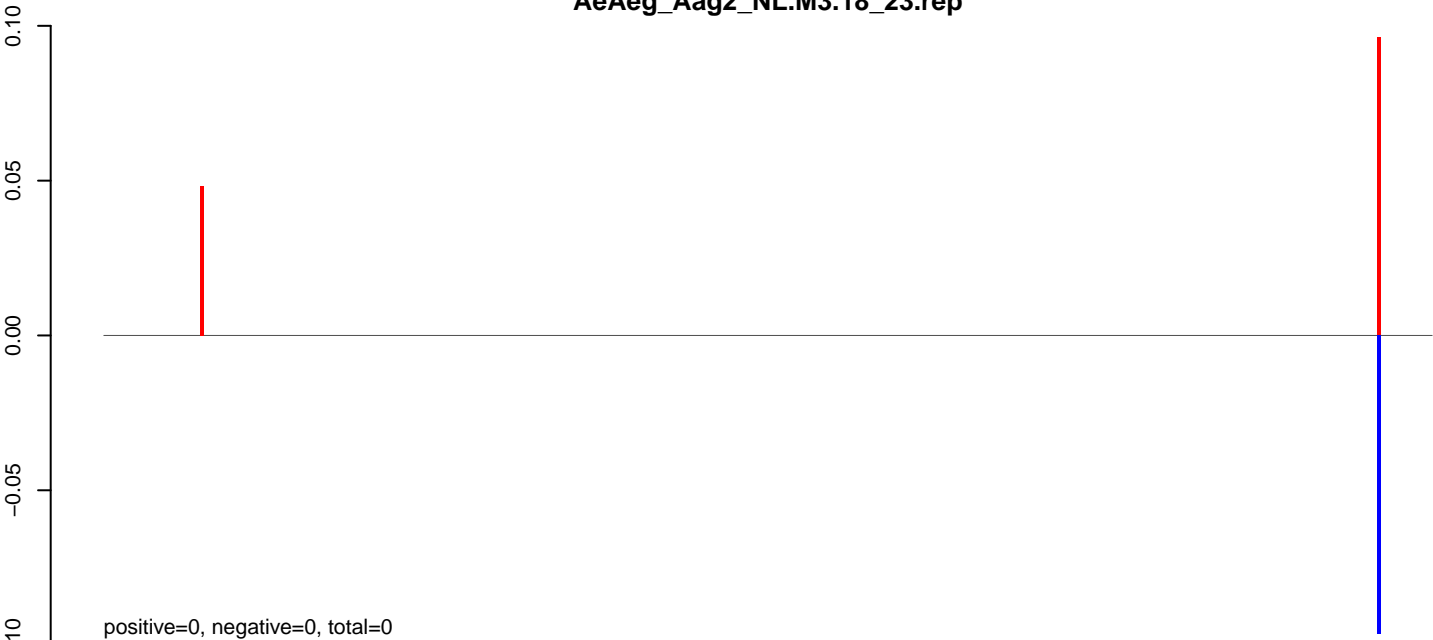
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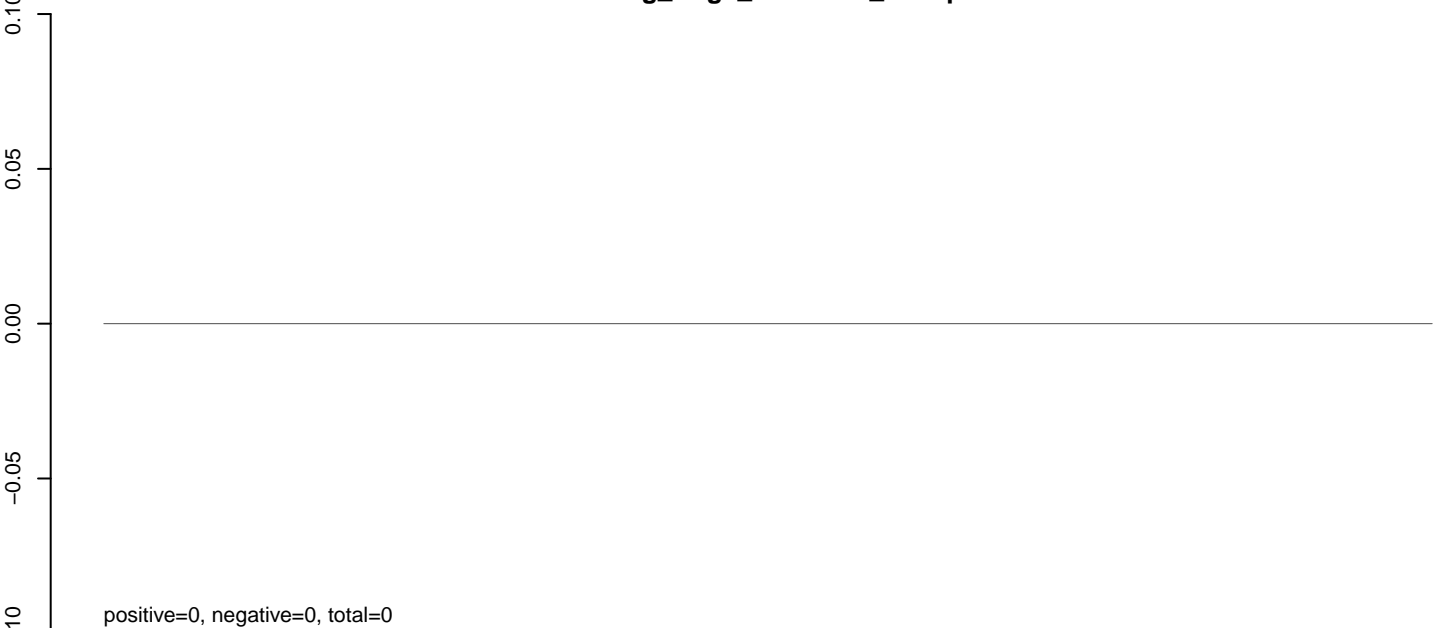
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AeAeg_Aag2_NL.M3.18_23.rep



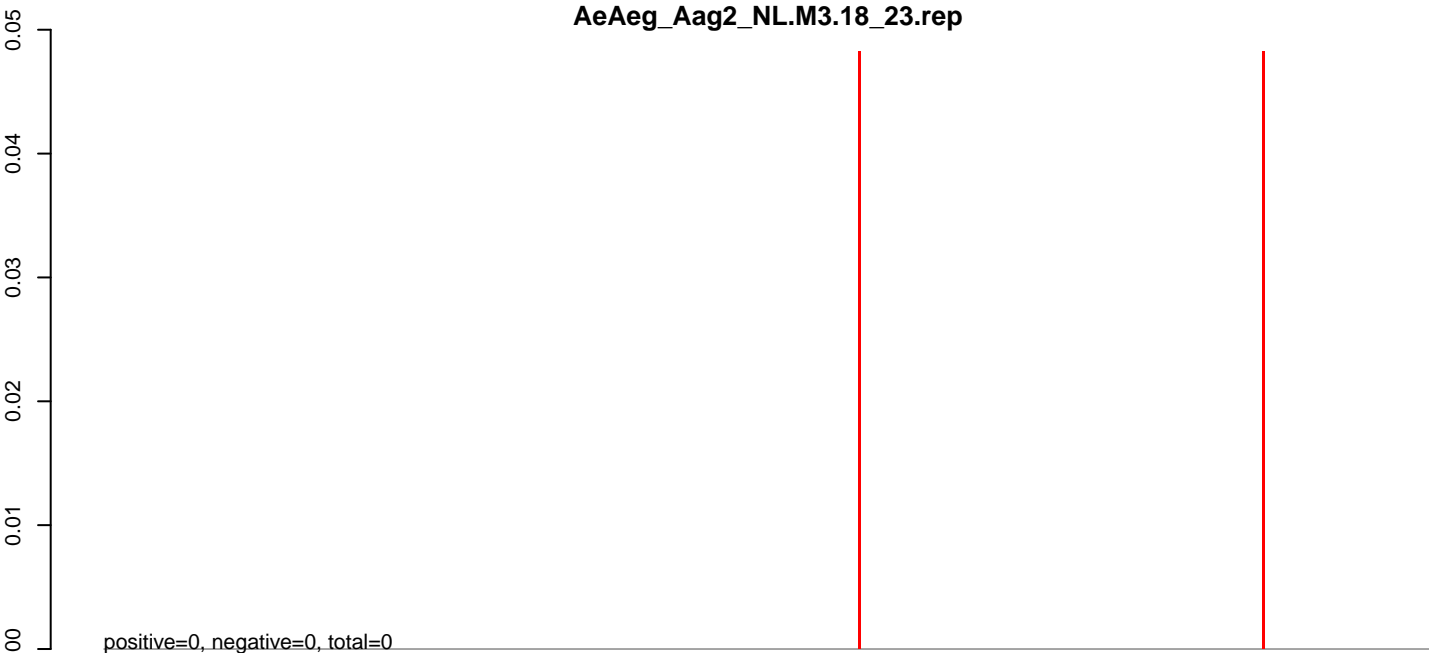
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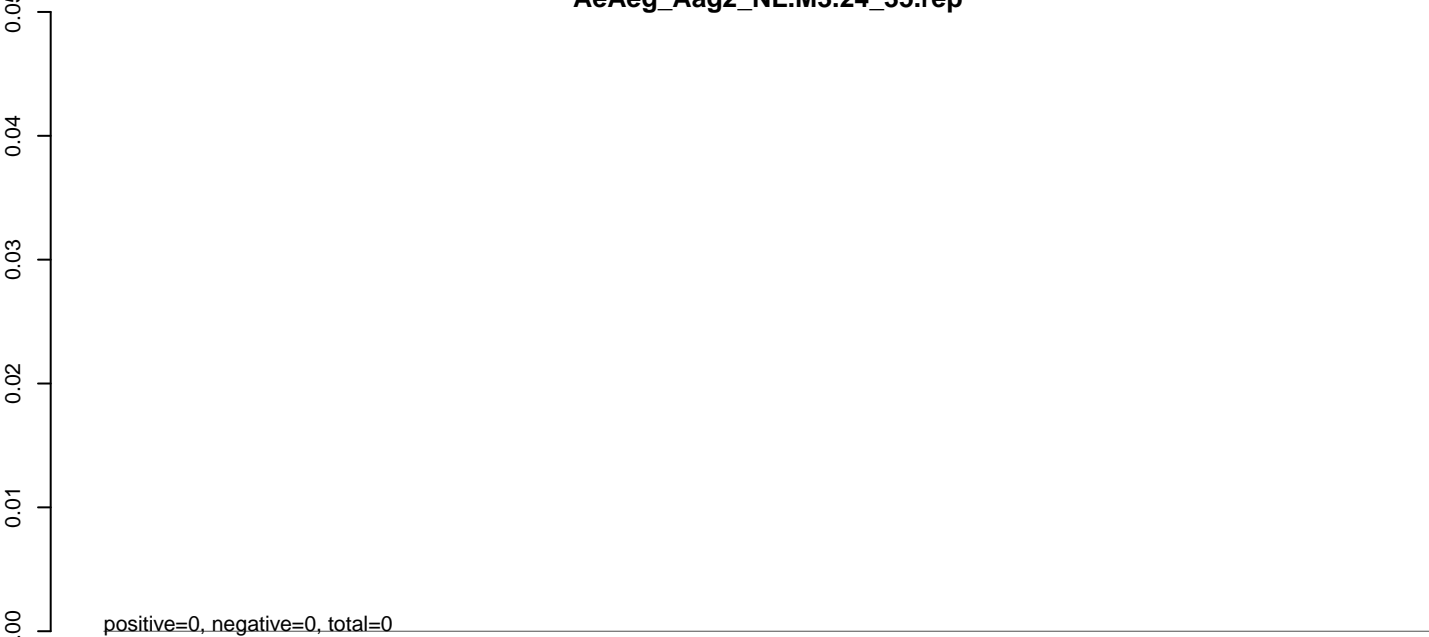
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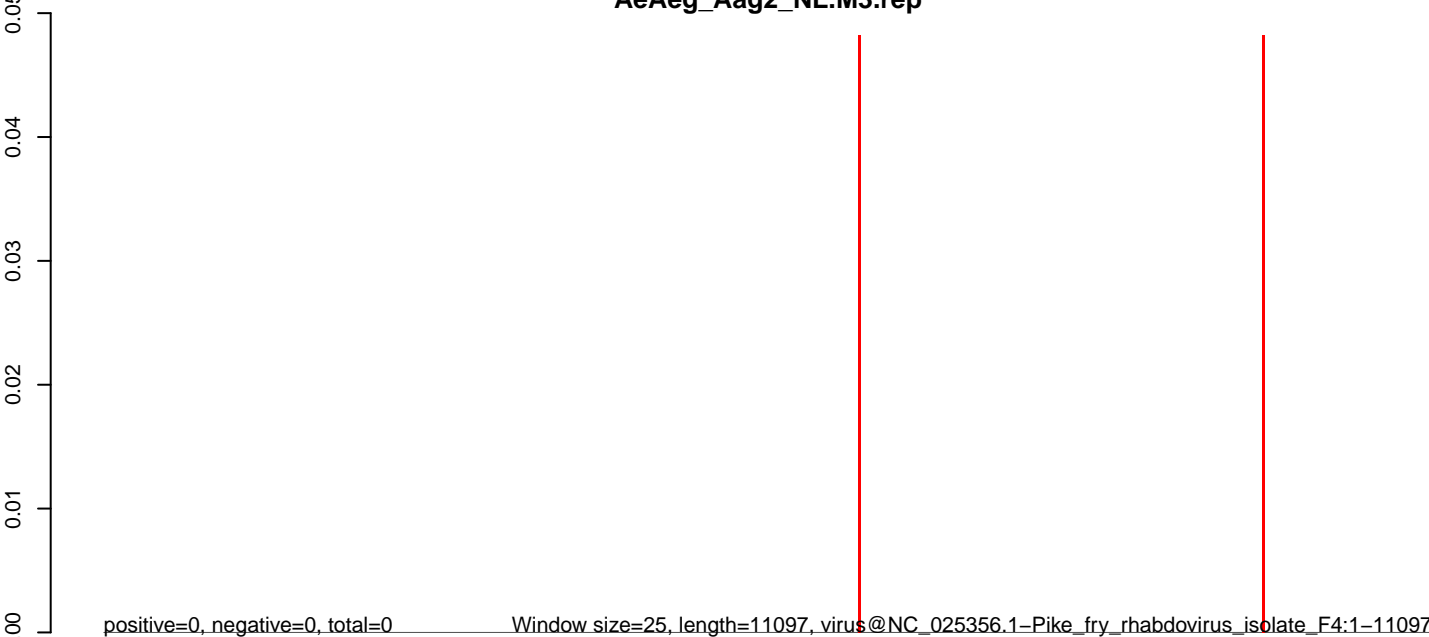
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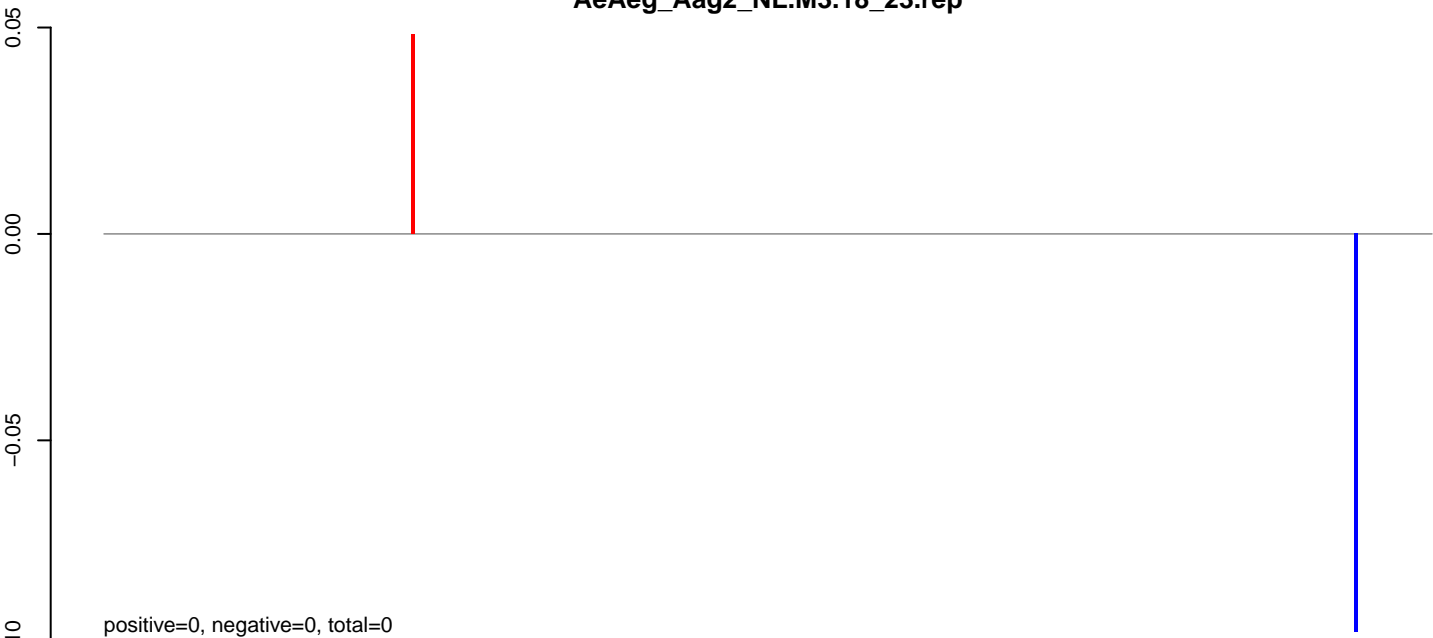


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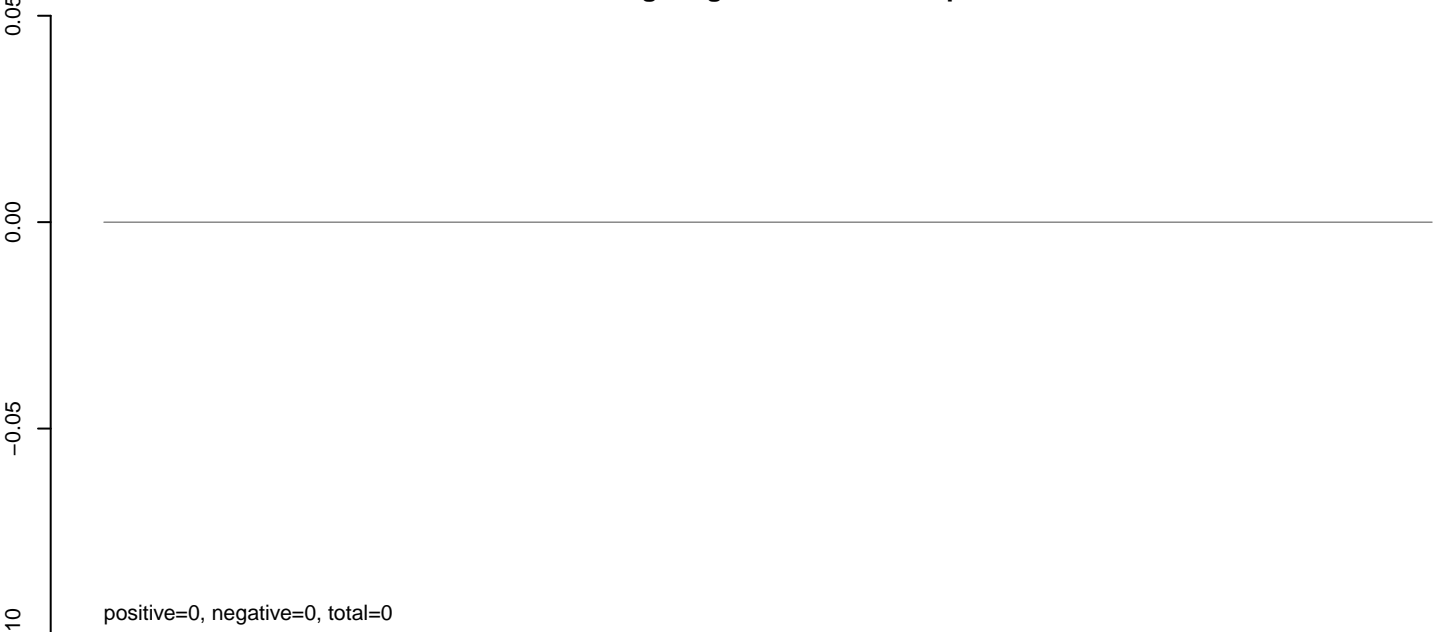


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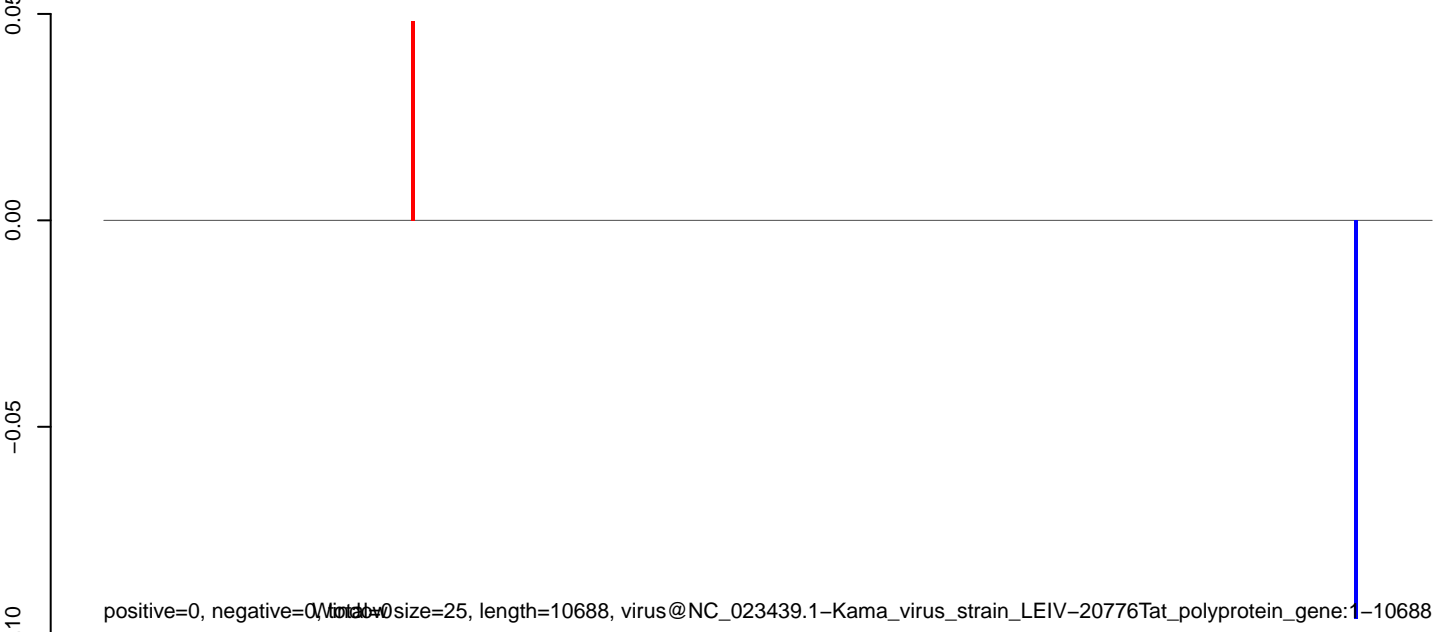
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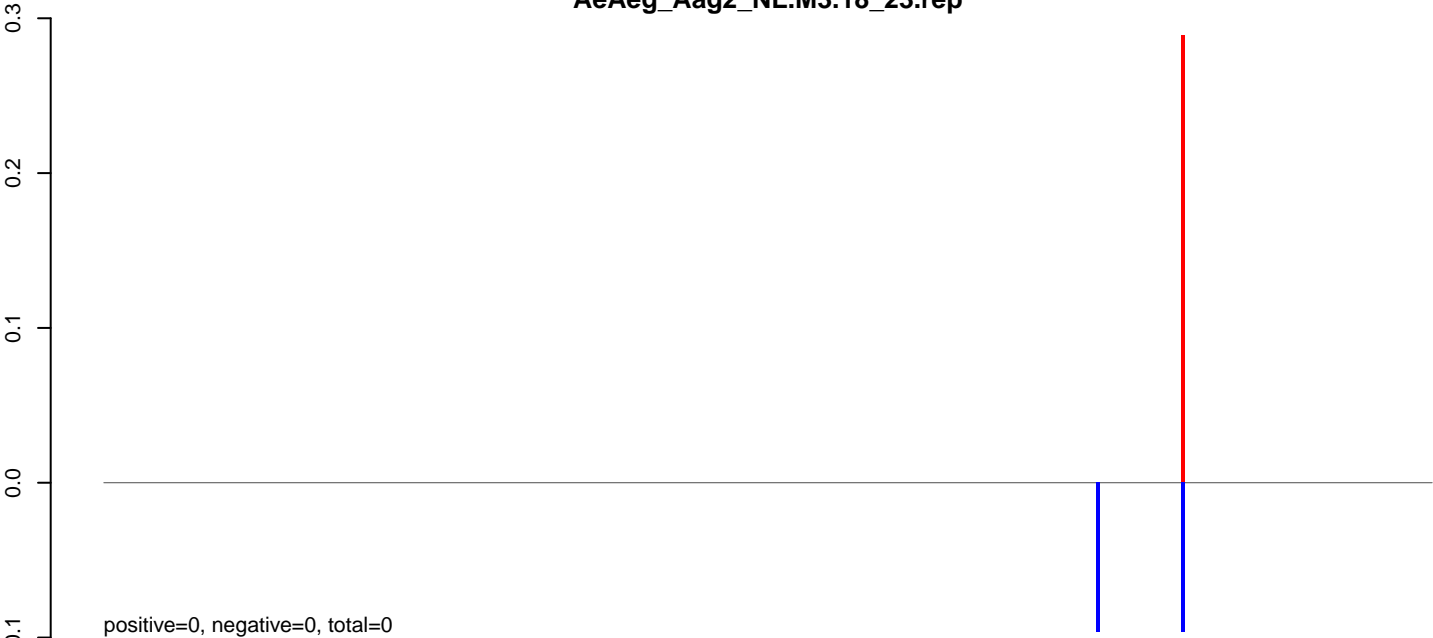
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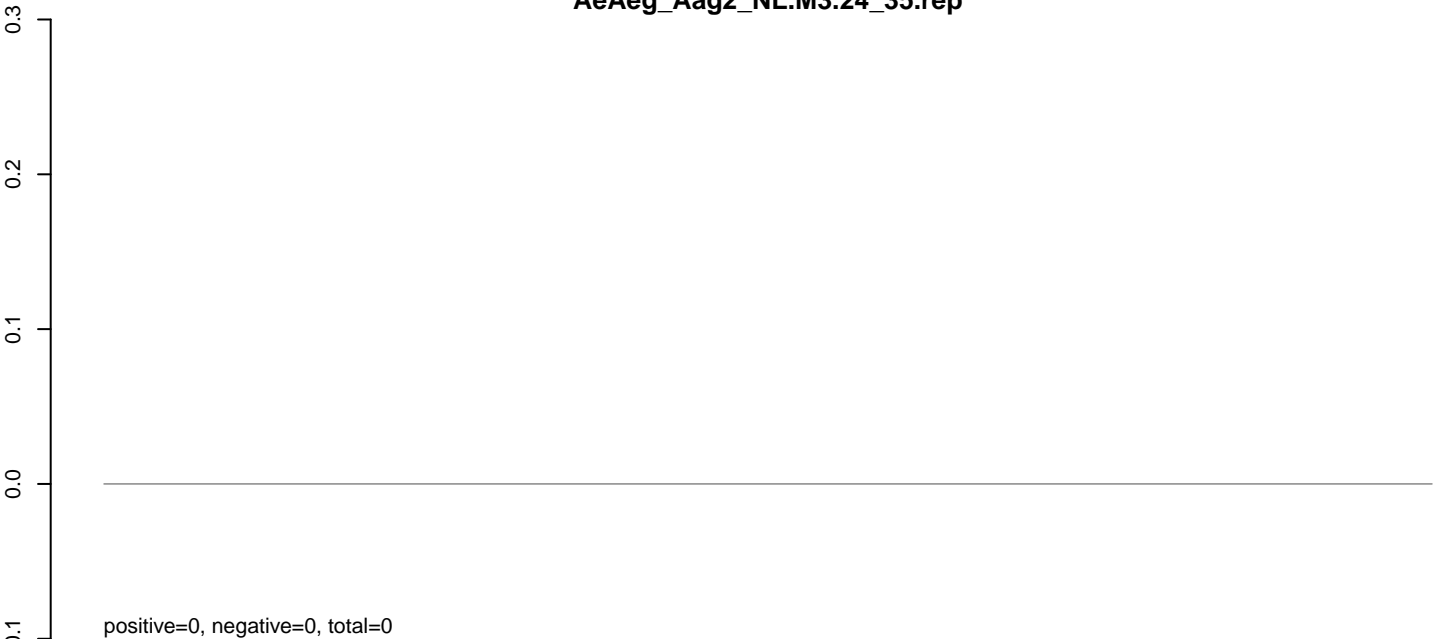
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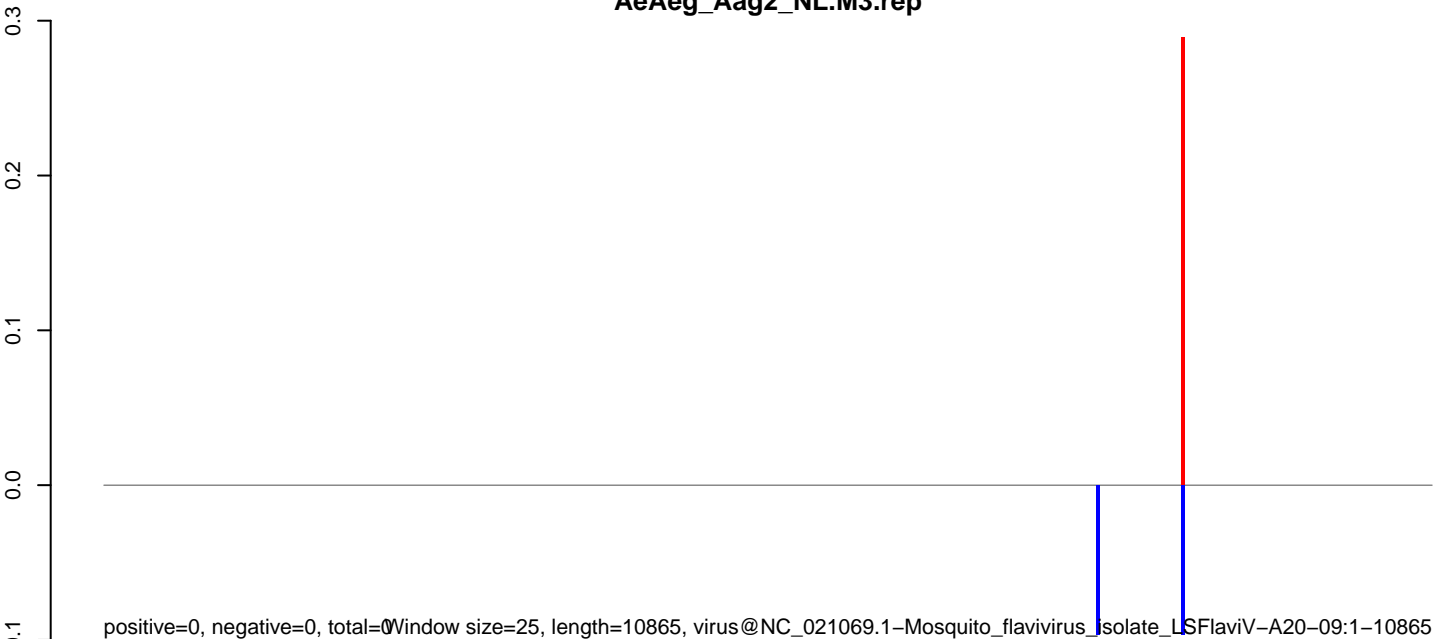
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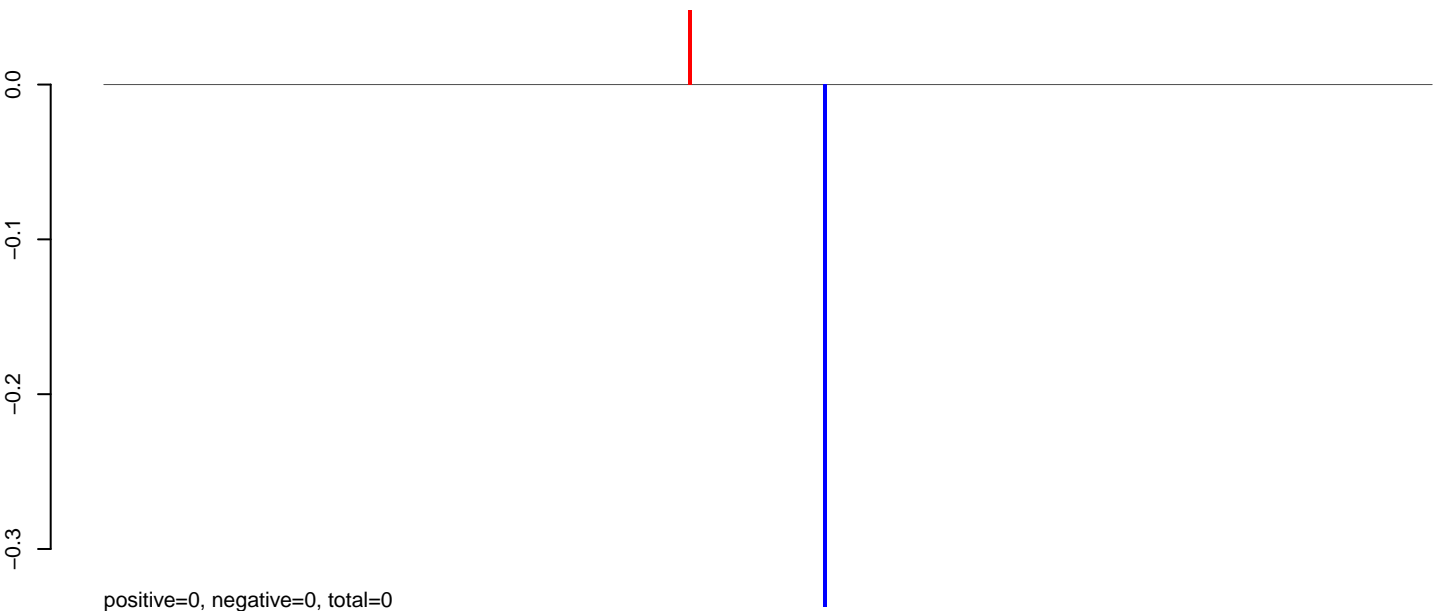
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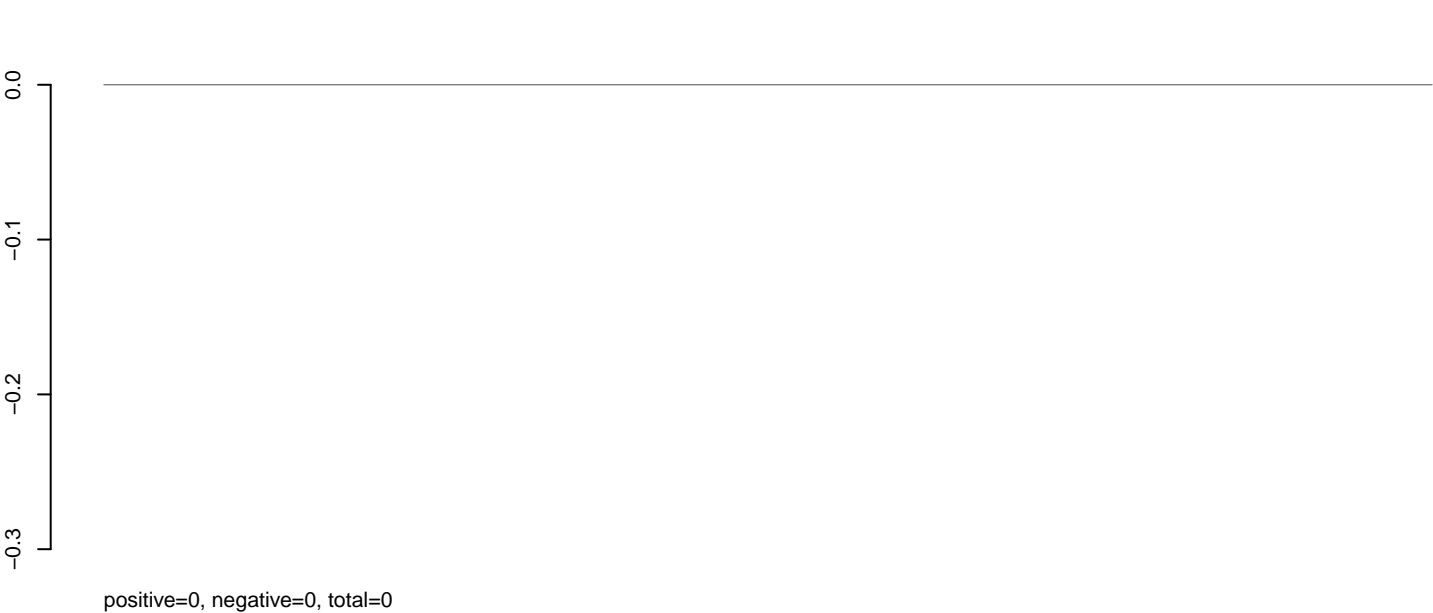
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AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep

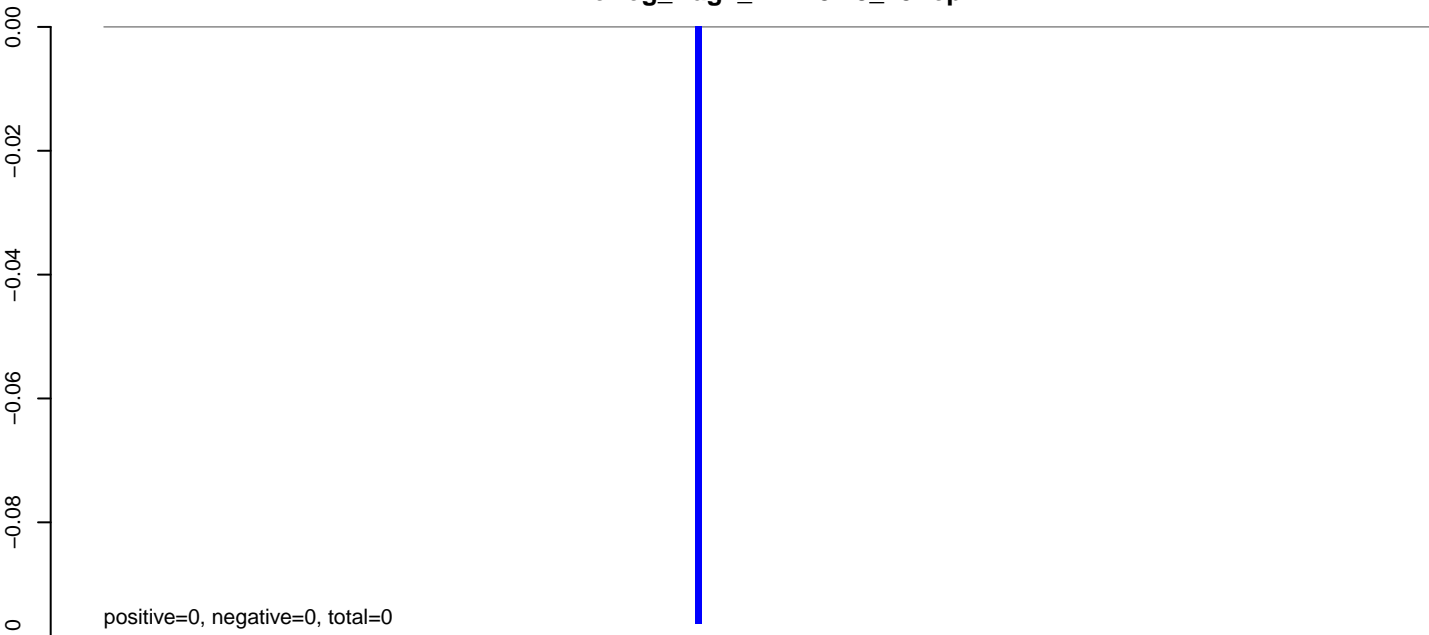


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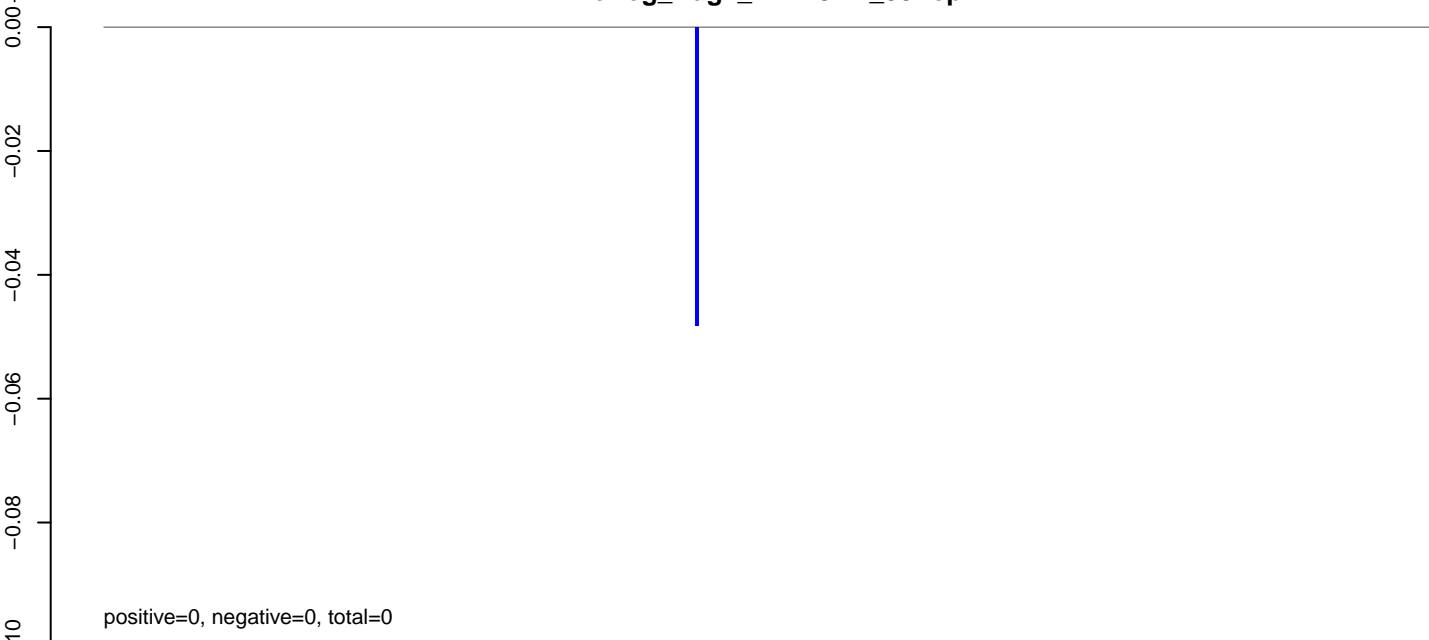


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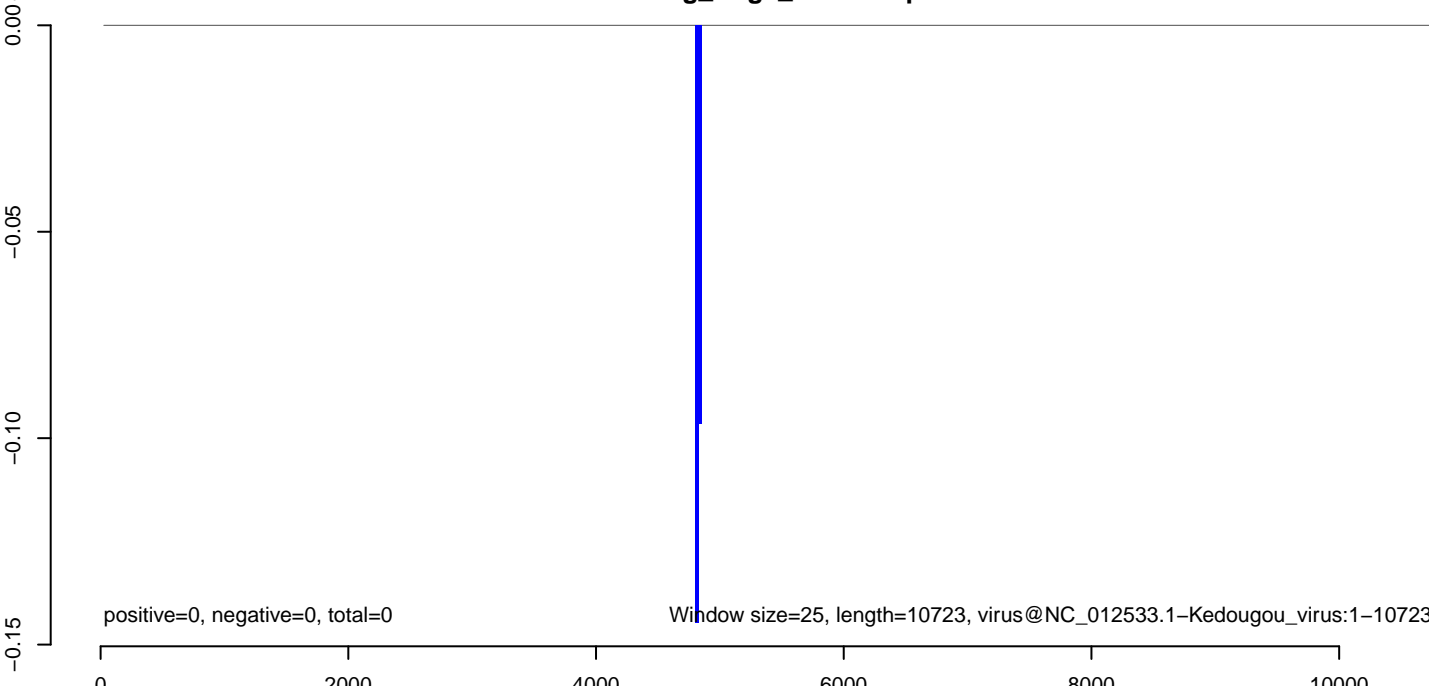
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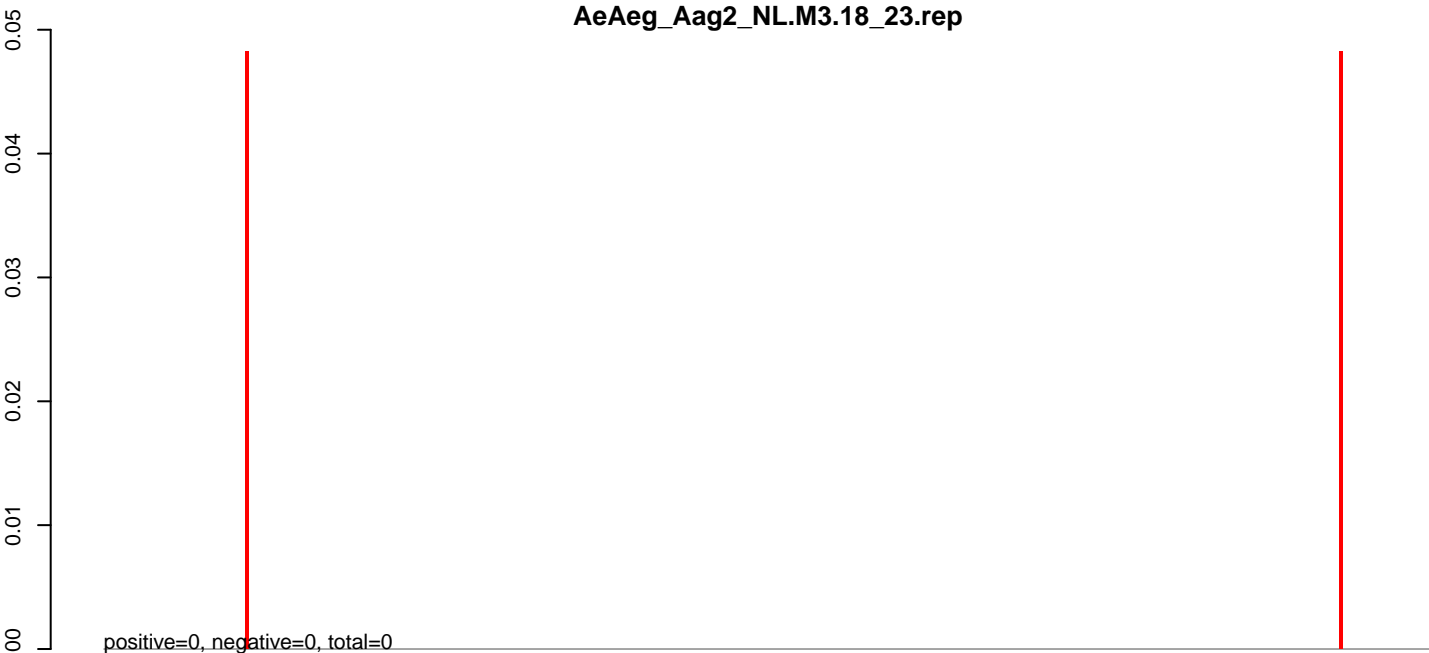
AeAeg_Aag2_NL.M3.24_35.rep



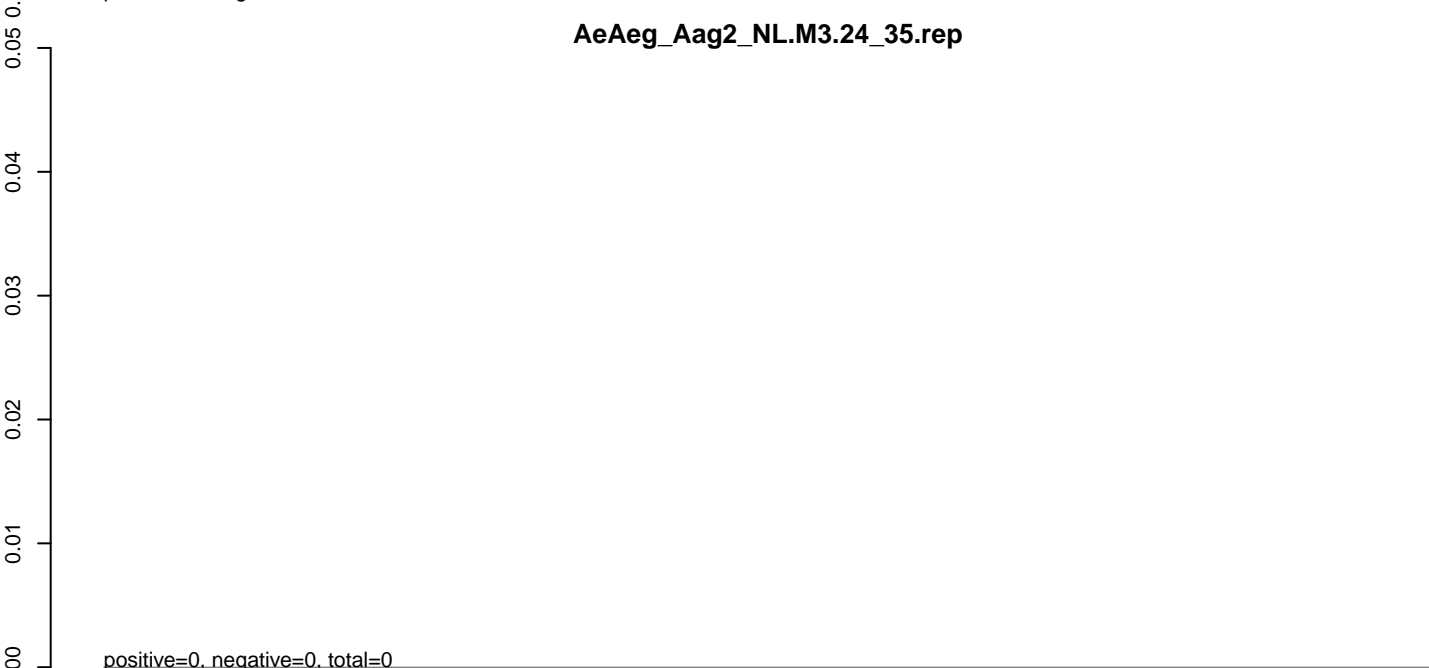
AeAeg_Aag2_NL.M3.rep



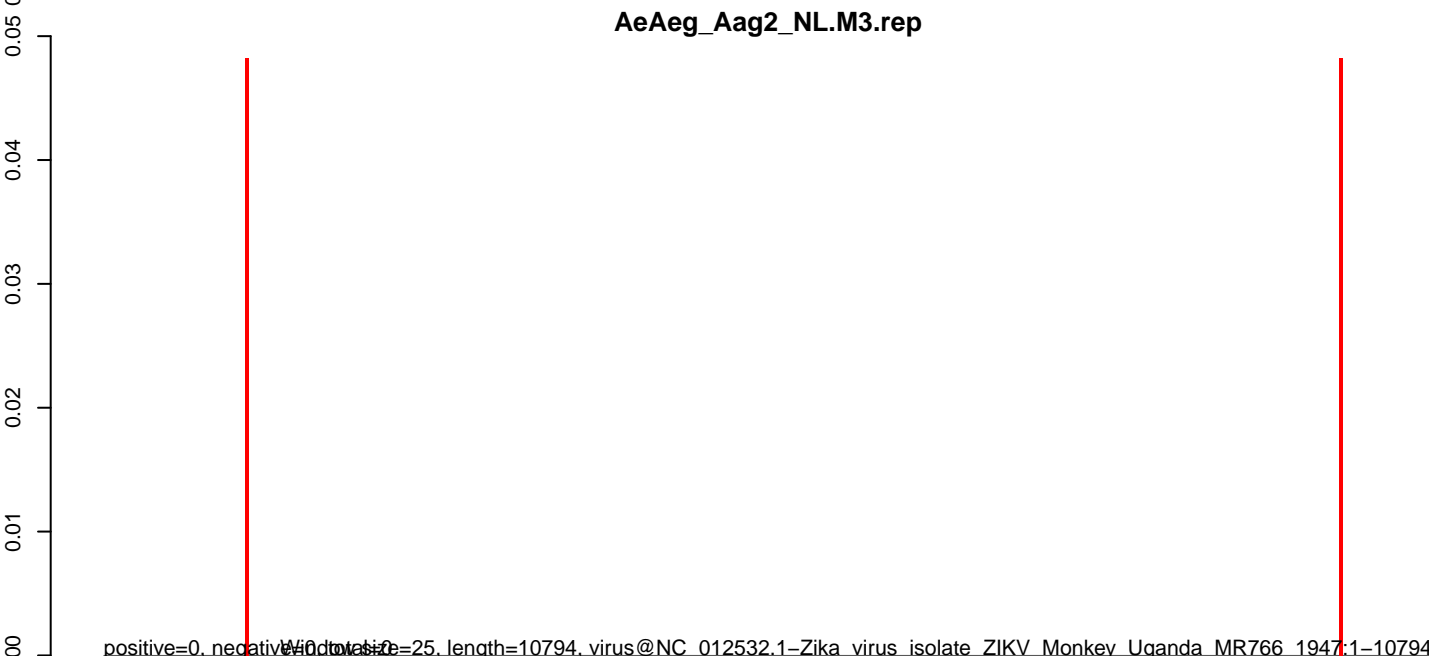
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep

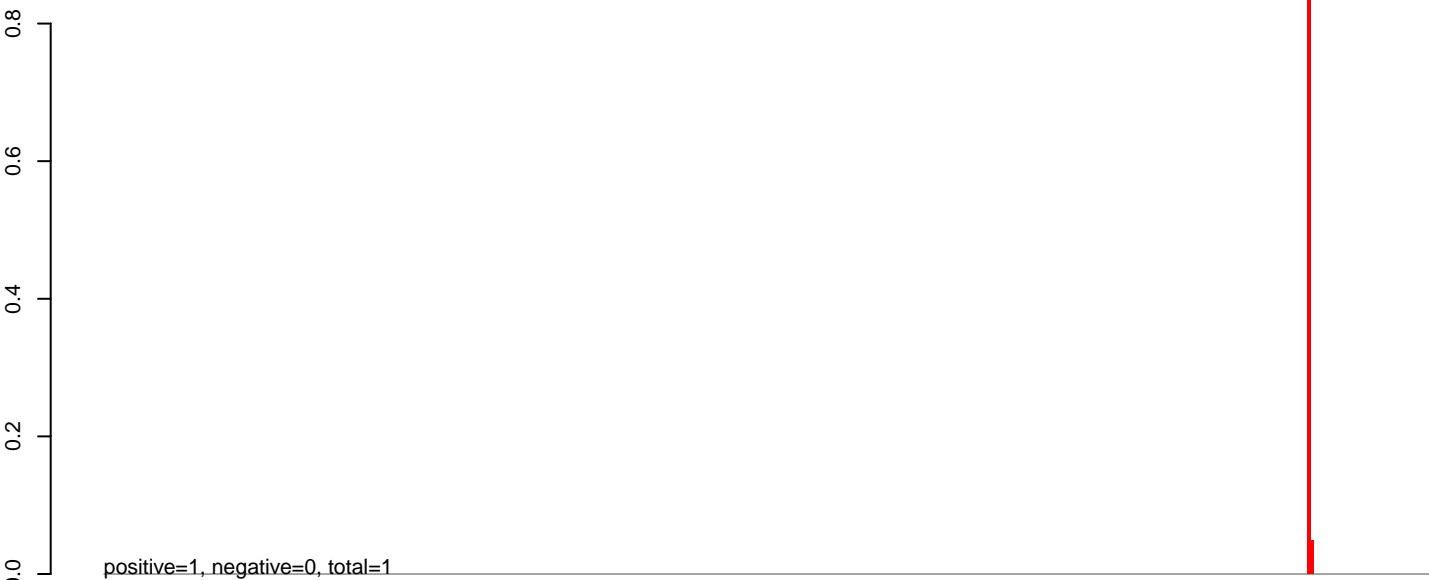


AeAeg_Aag2_NL.M3.rep

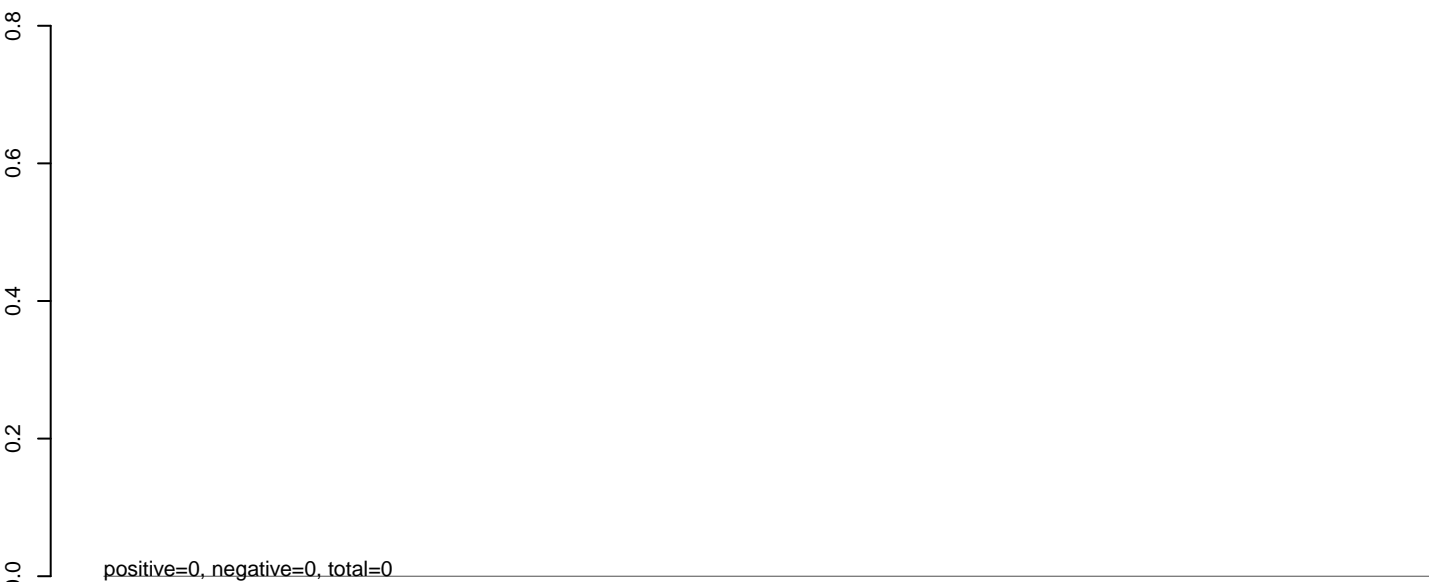


0 2000 4000 6000 8000 10000

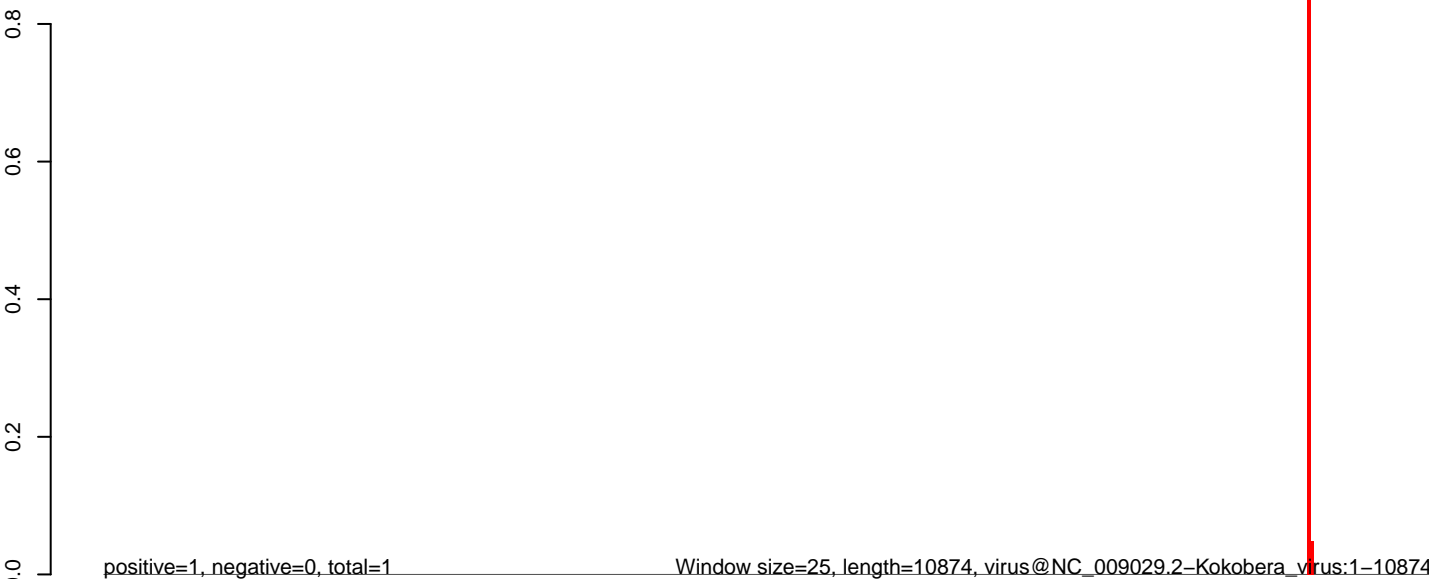
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep

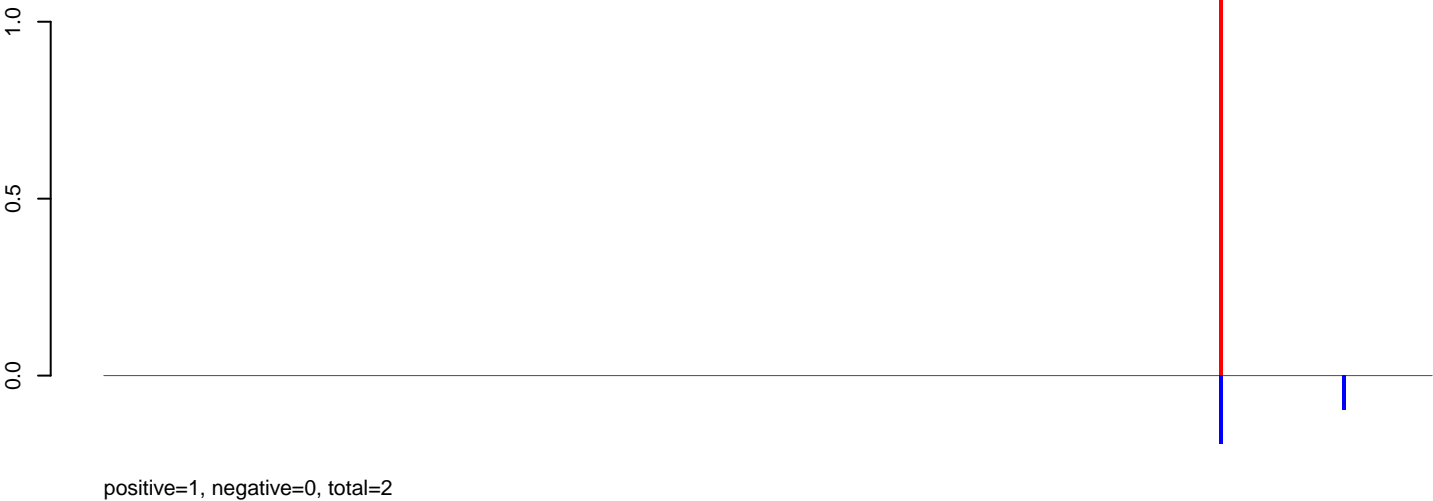


AeAeg_Aag2_NL.M3.rep

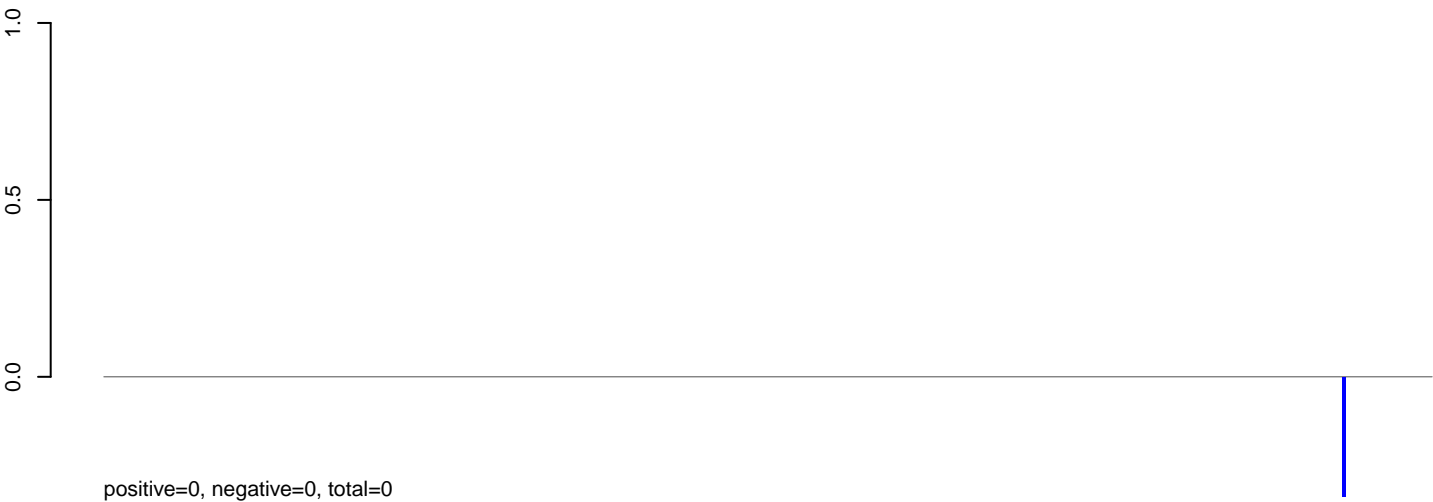


0 2000 4000 6000 8000 10000

AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep

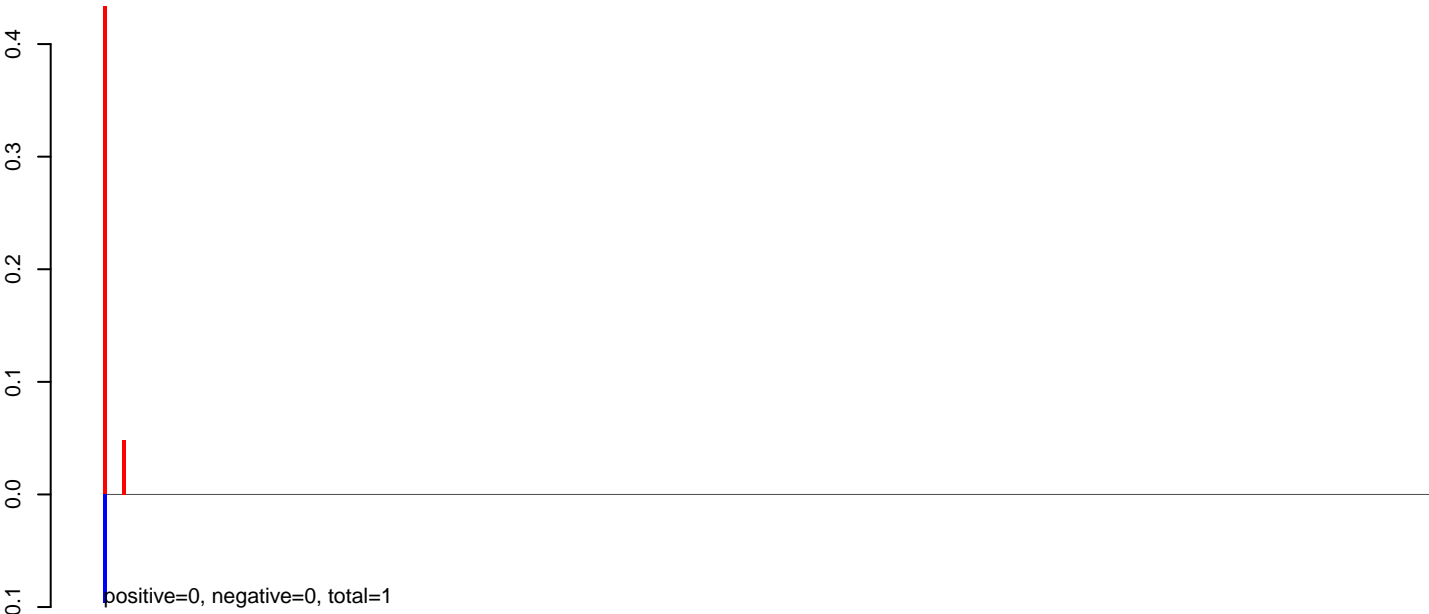


AeAeg_Aag2_NL.M3.rep

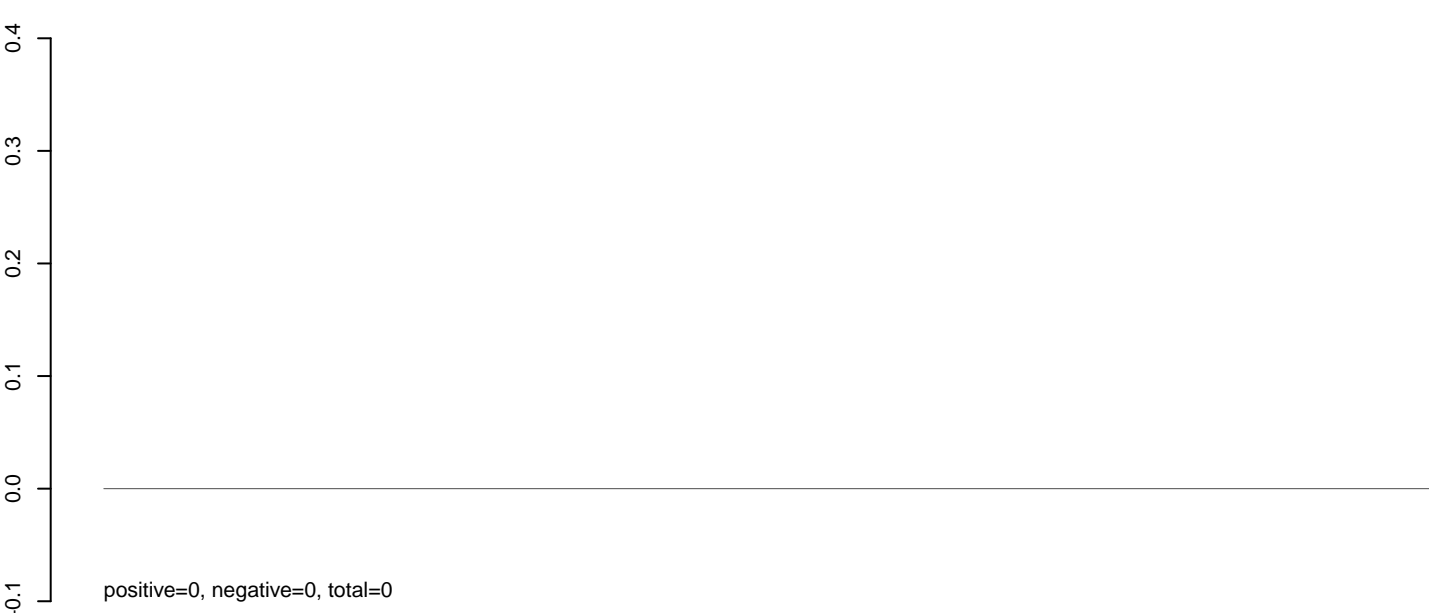


Window size=25, length=10755, virus@NC_009028.2-Ilheus_virus1-10755

AeAeg_Aag2_NL.M3.18_23.rep



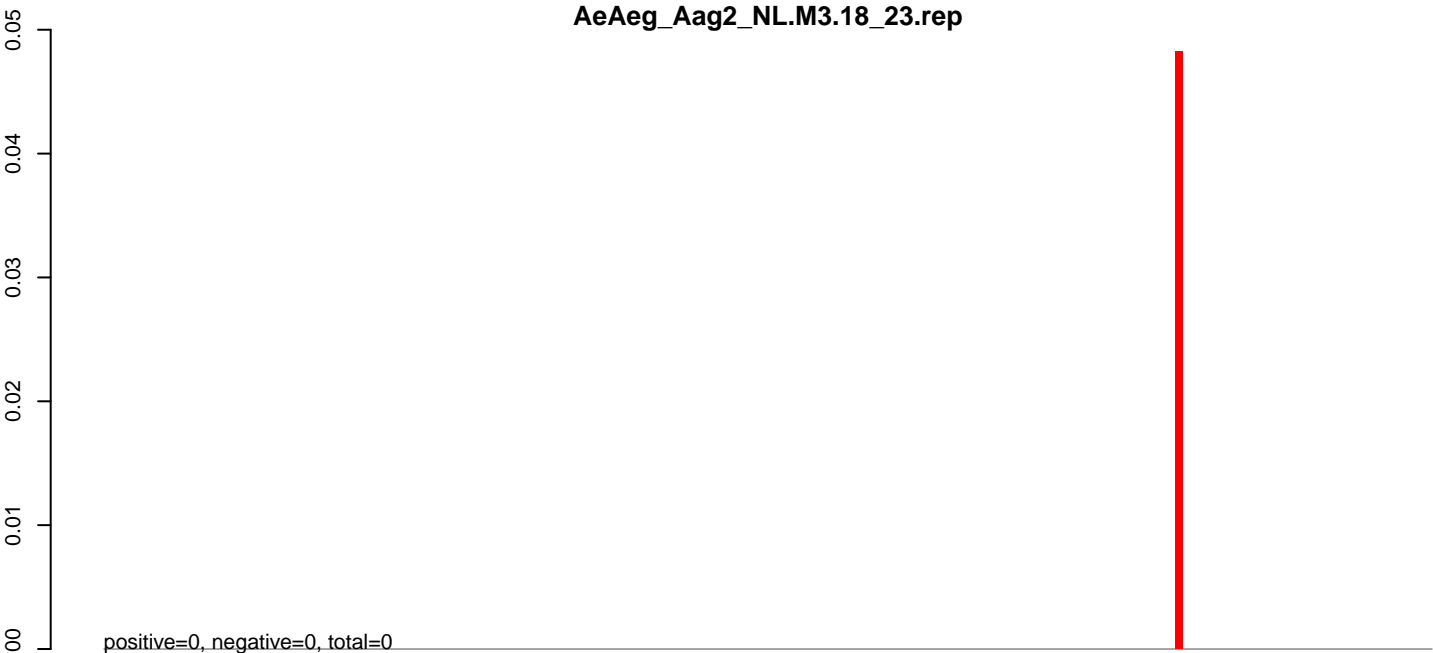
AeAeg_Aag2_NL.M3.24_35.rep



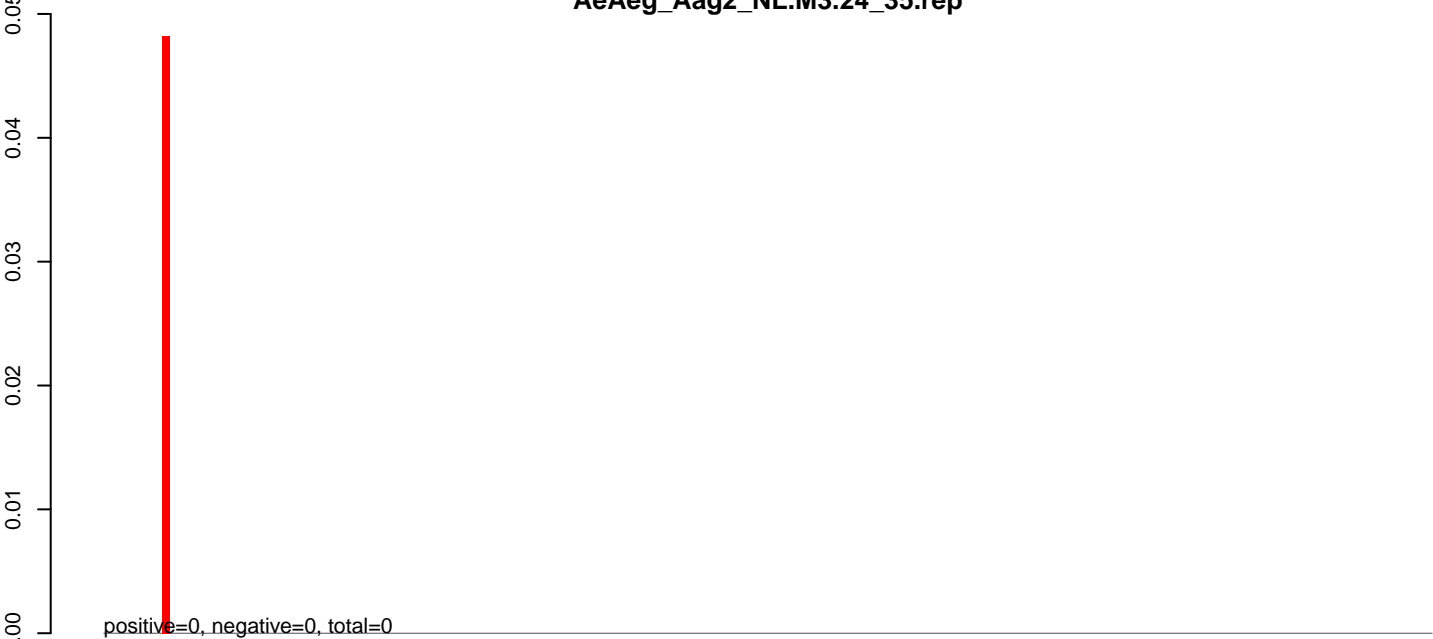
AeAeg_Aag2_NL.M3.rep



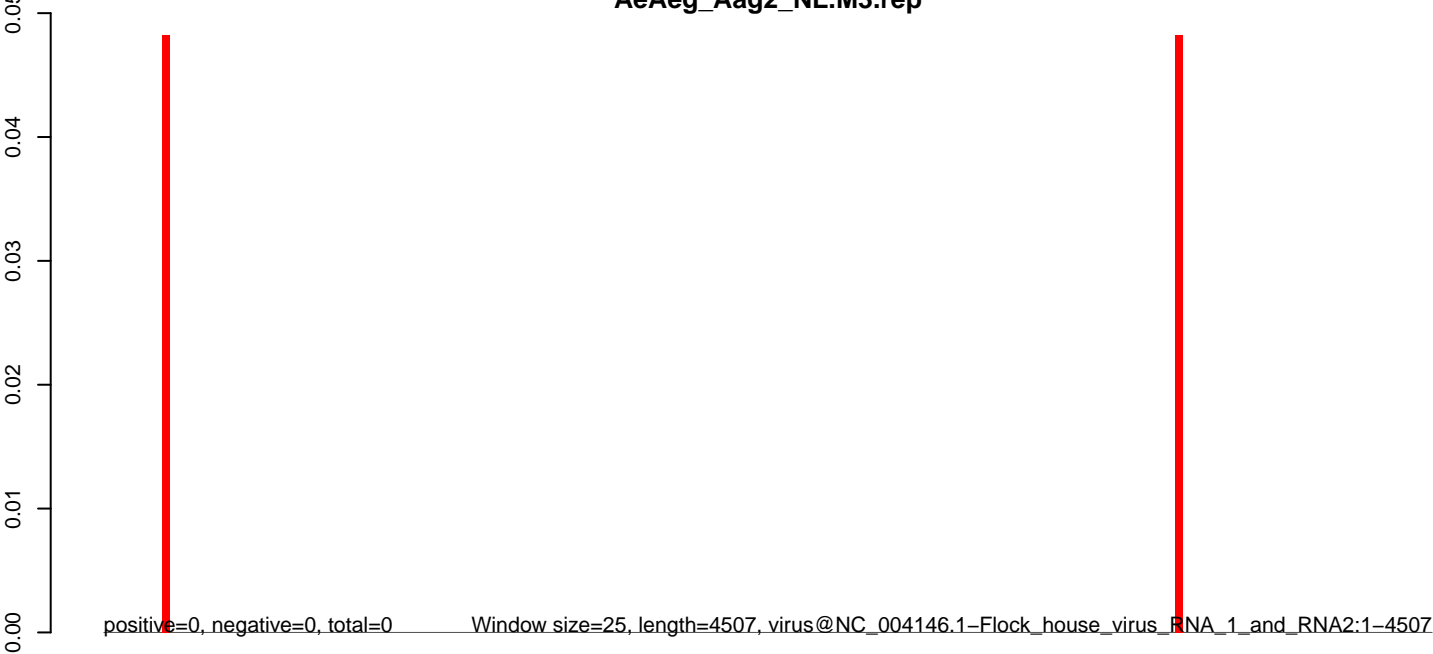
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep



AeAeg_Aag2_NL.M3.rep



AeAeg_Aag2_NL.M3.18_23.rep

positive=1, negative=0, total=1

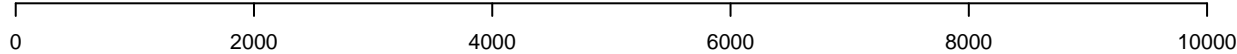
AeAeg_Aag2_NL.M3.24_35.rep

positive=0, negative=0, total=0

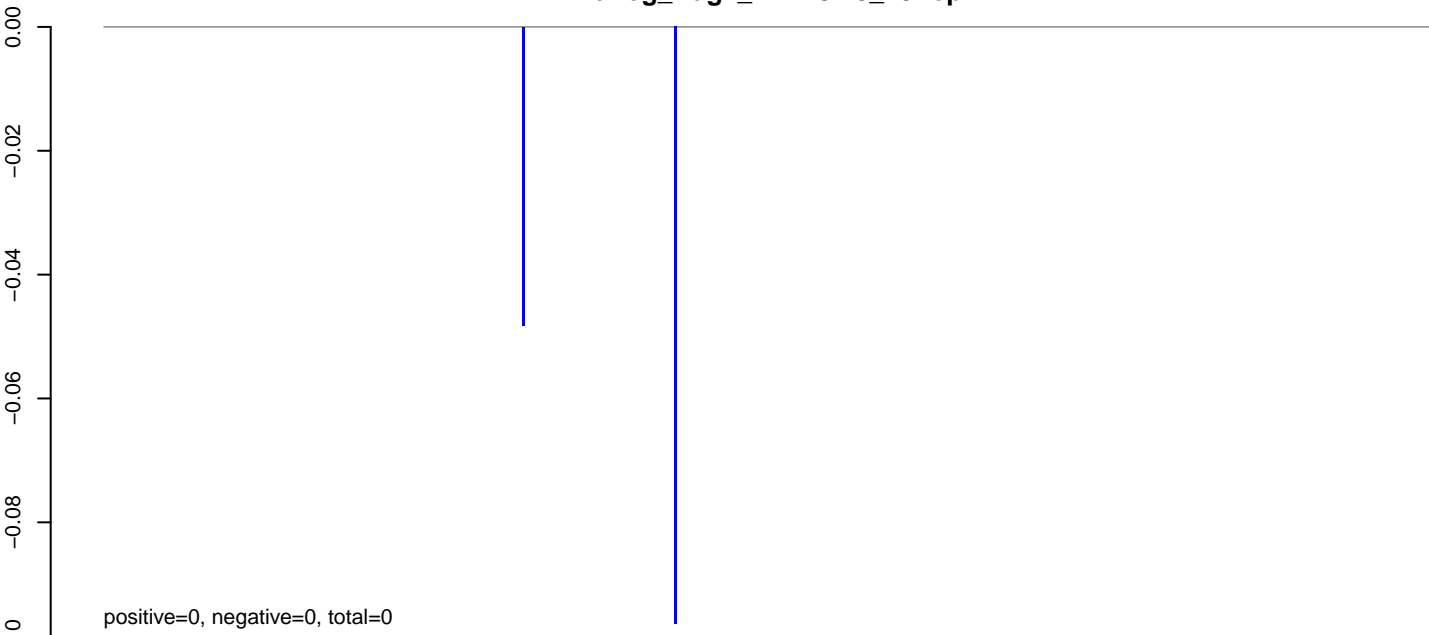
AeAeg_Aag2_NL.M3.rep

positive=1, negative=0, total=1

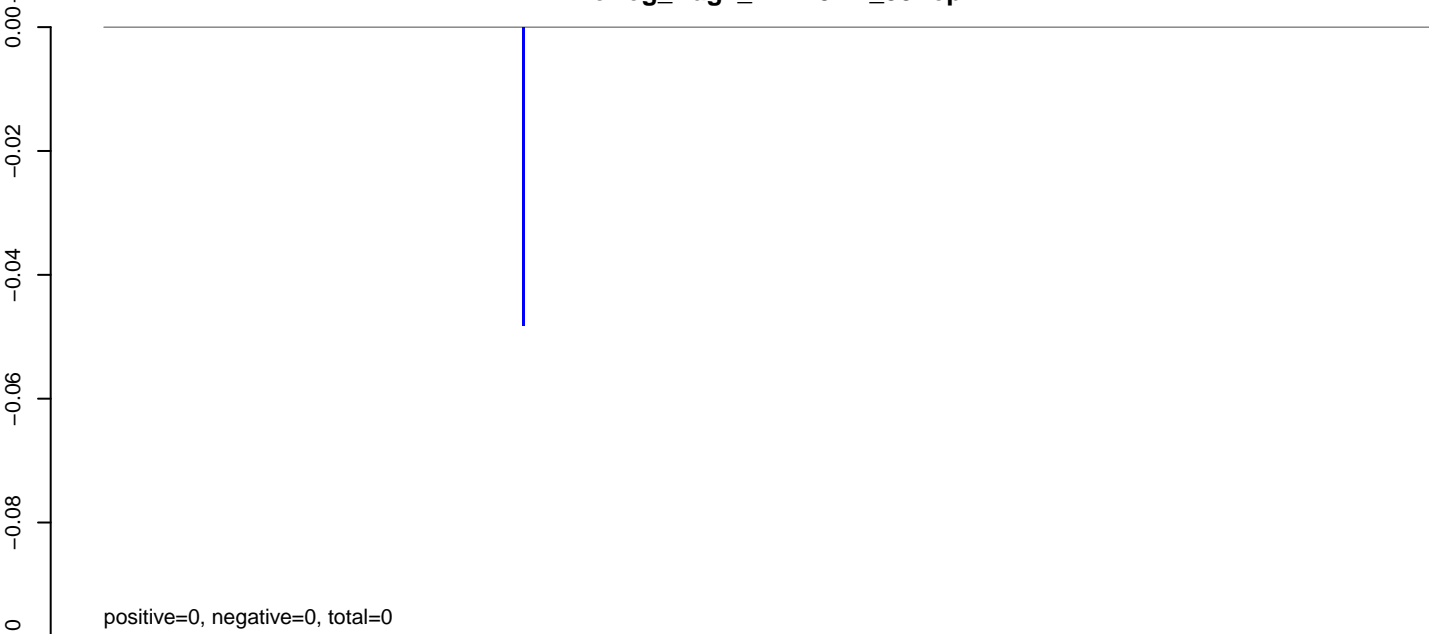
Window size=25, length=11141, virus@NC_001672.1-Tick-borne_encephalitis_virus:1-11141



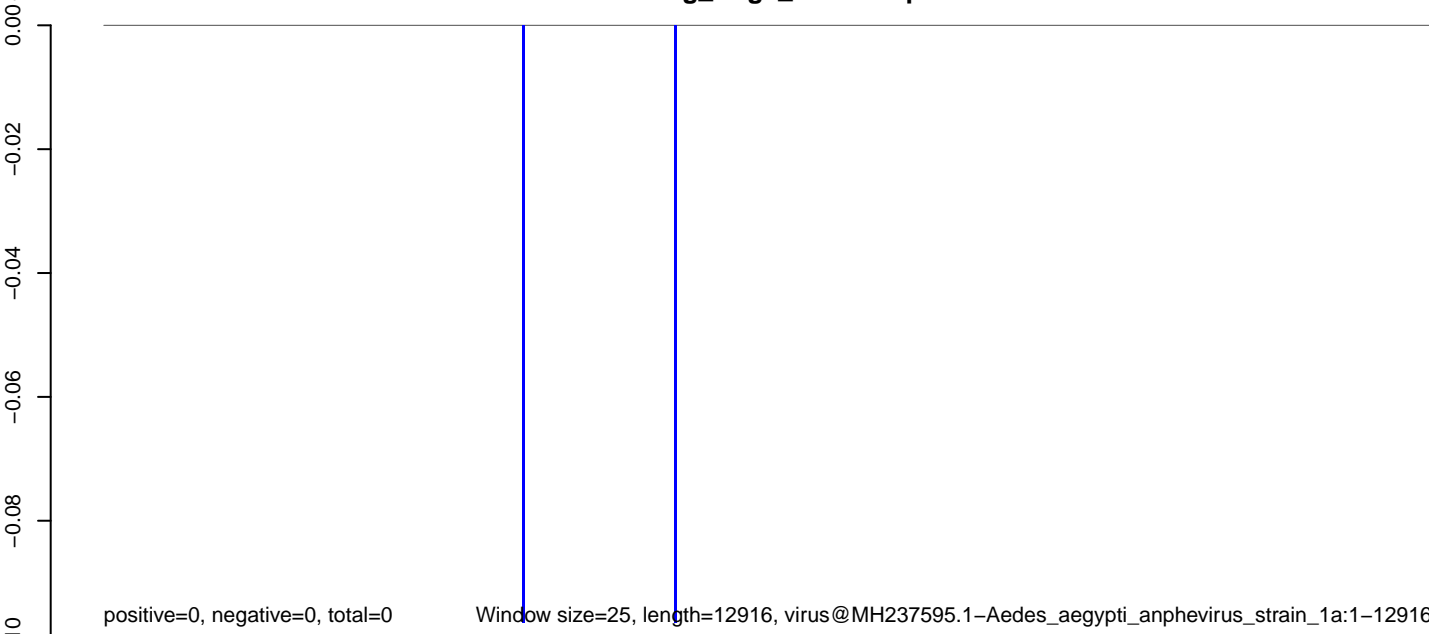
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep

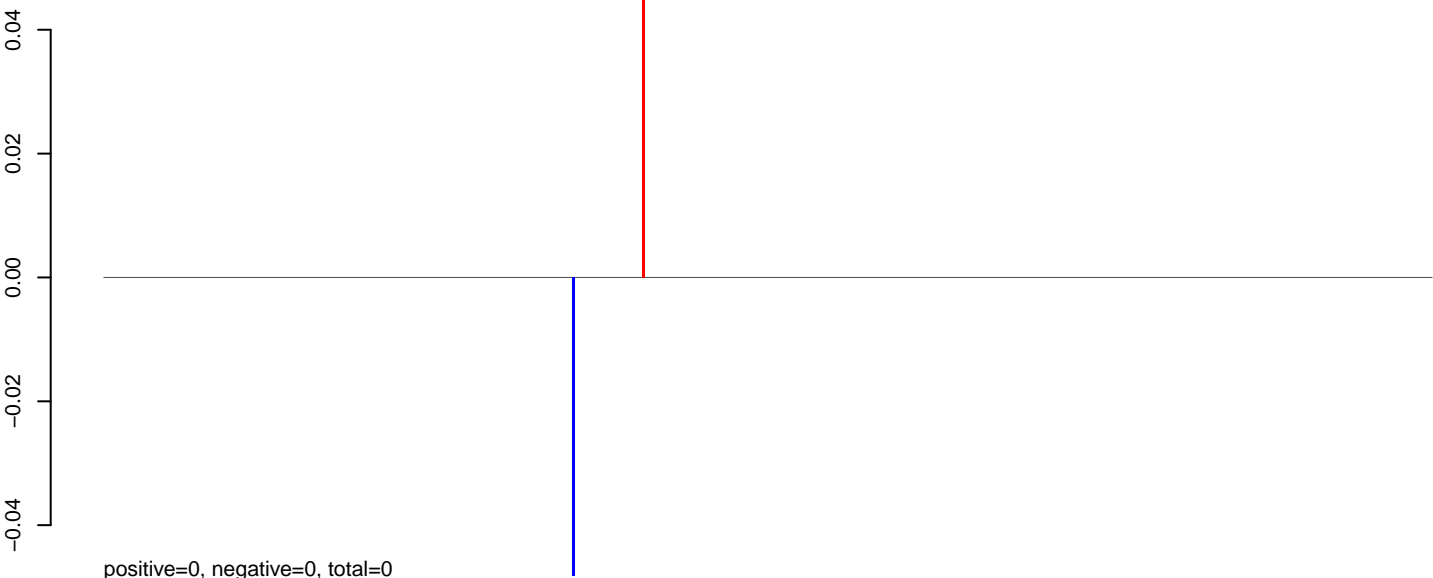


AeAeg_Aag2_NL.M3.rep

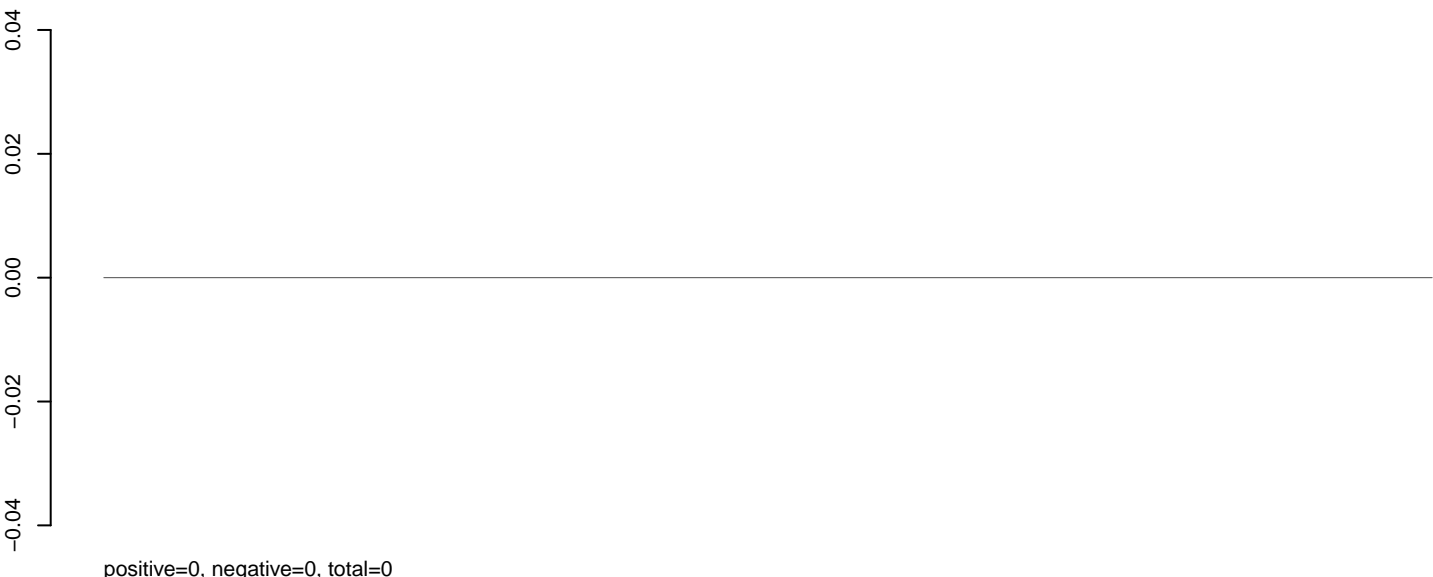


0 2000 4000 6000 8000 10000 12000

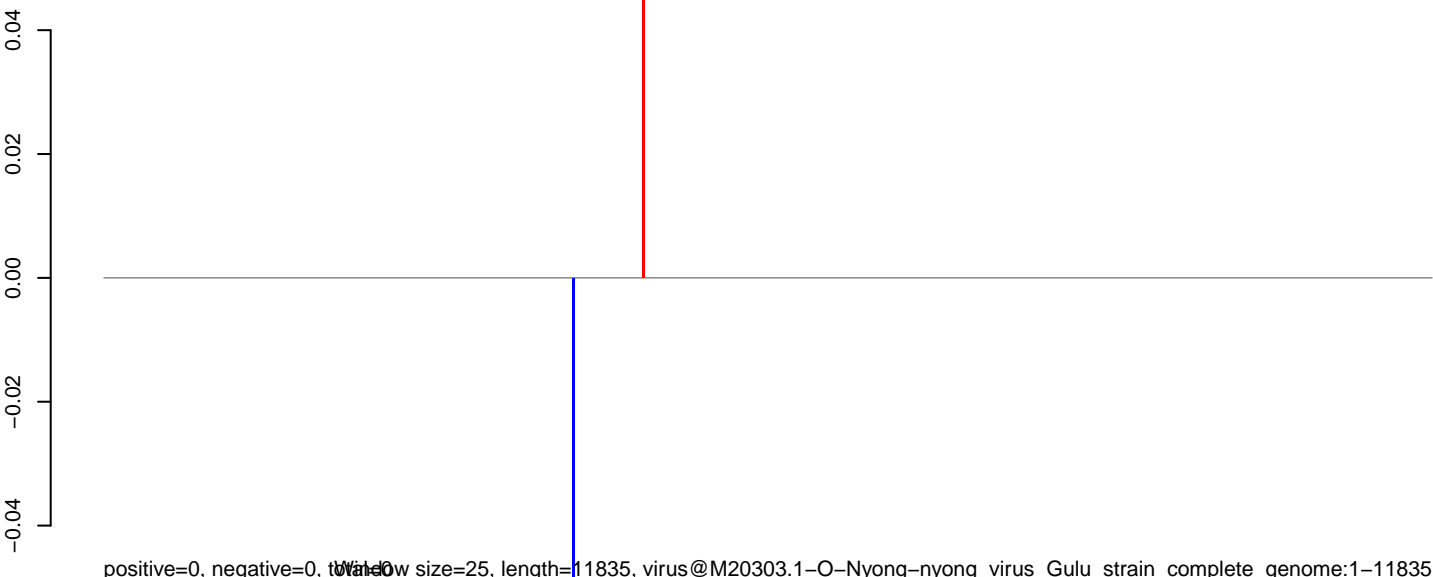
AeAeg_Aag2_NL.M3.18_23.rep



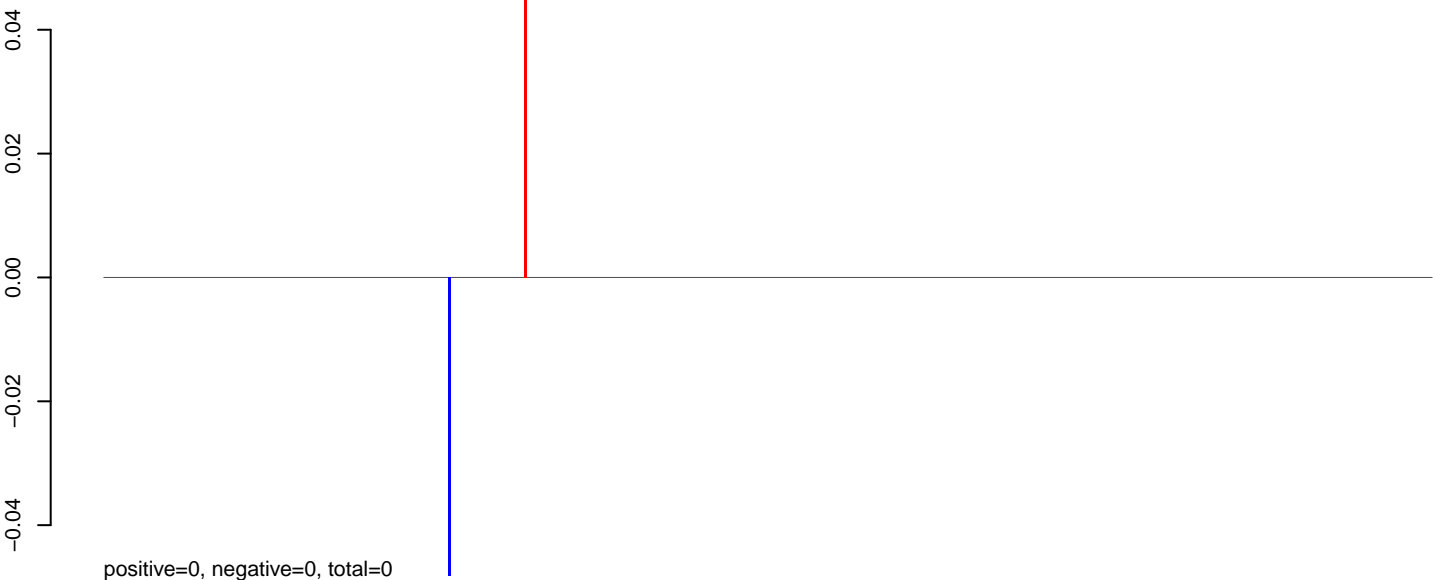
AeAeg_Aag2_NL.M3.24_35.rep



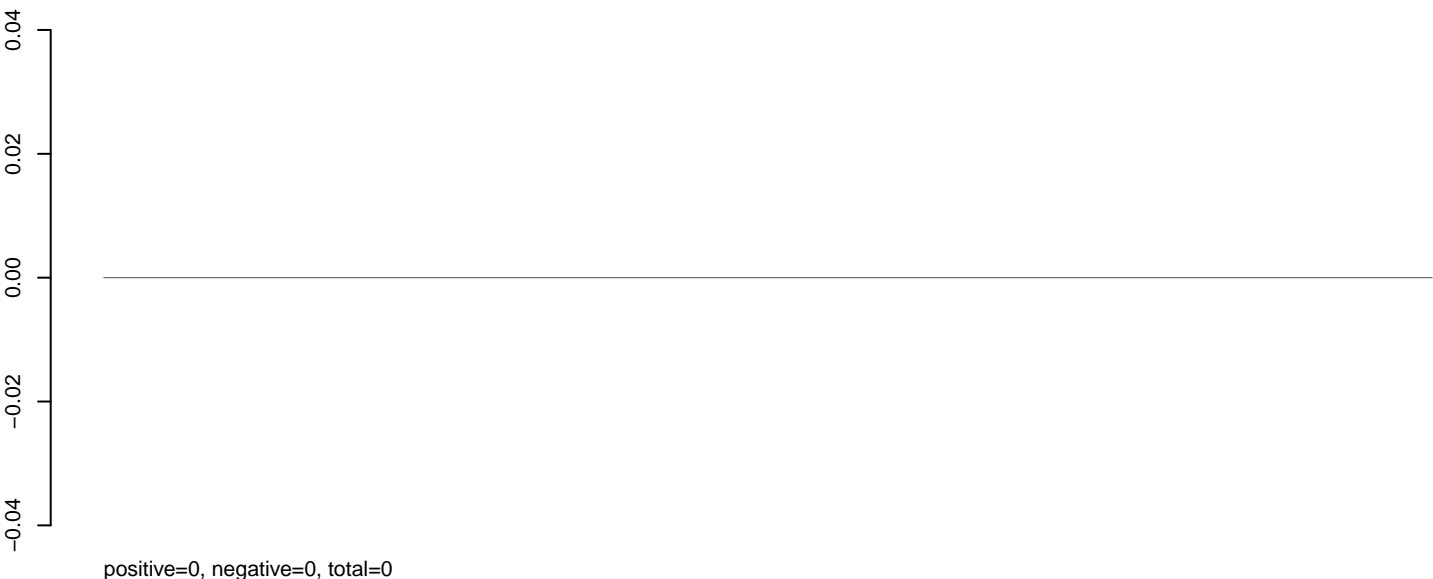
AeAeg_Aag2_NL.M3.rep



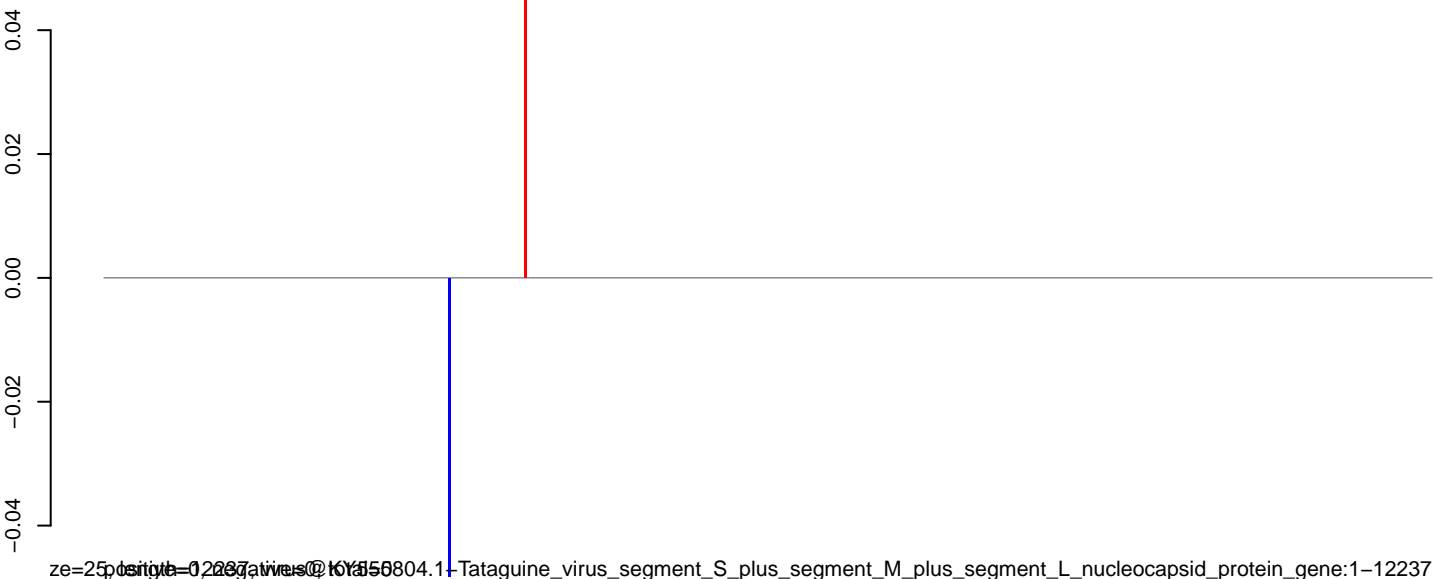
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep



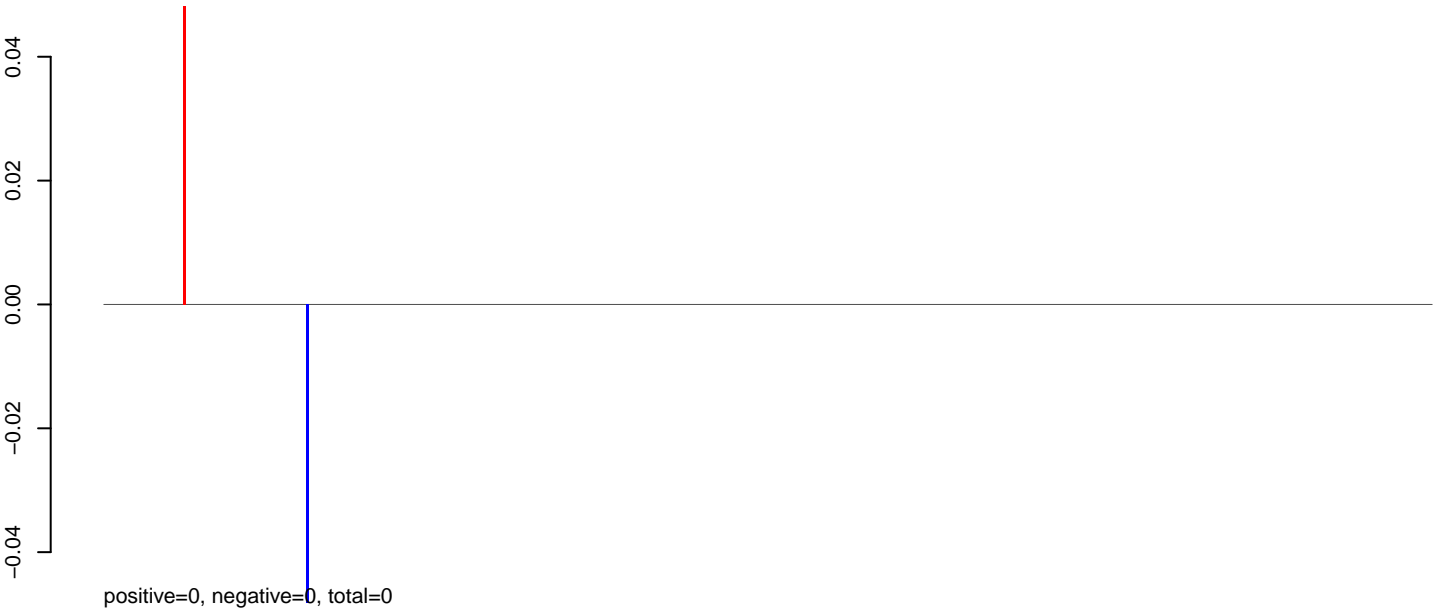
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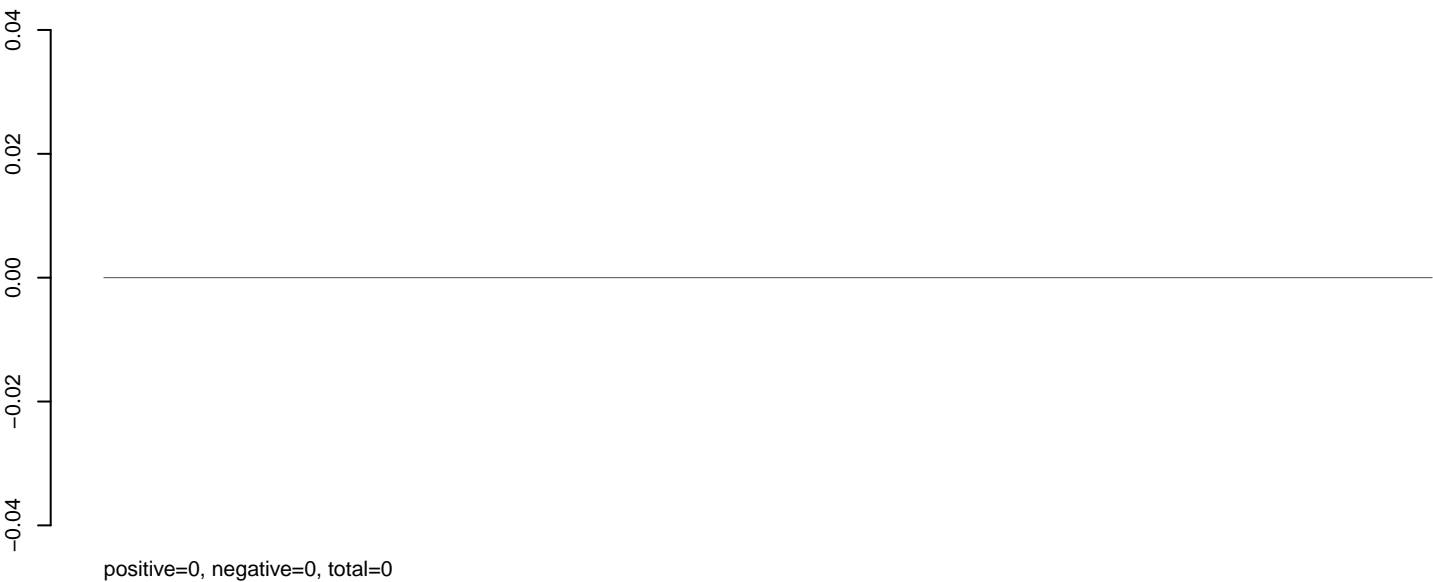
ze=25 path=12237atives@16156804.1-Tataguine_virus_segment_S_plus_segment_M_plus_segment_L_nucleocapsid_protein_gene:1-12237

0 2000 4000 6000 8000 10000 12000

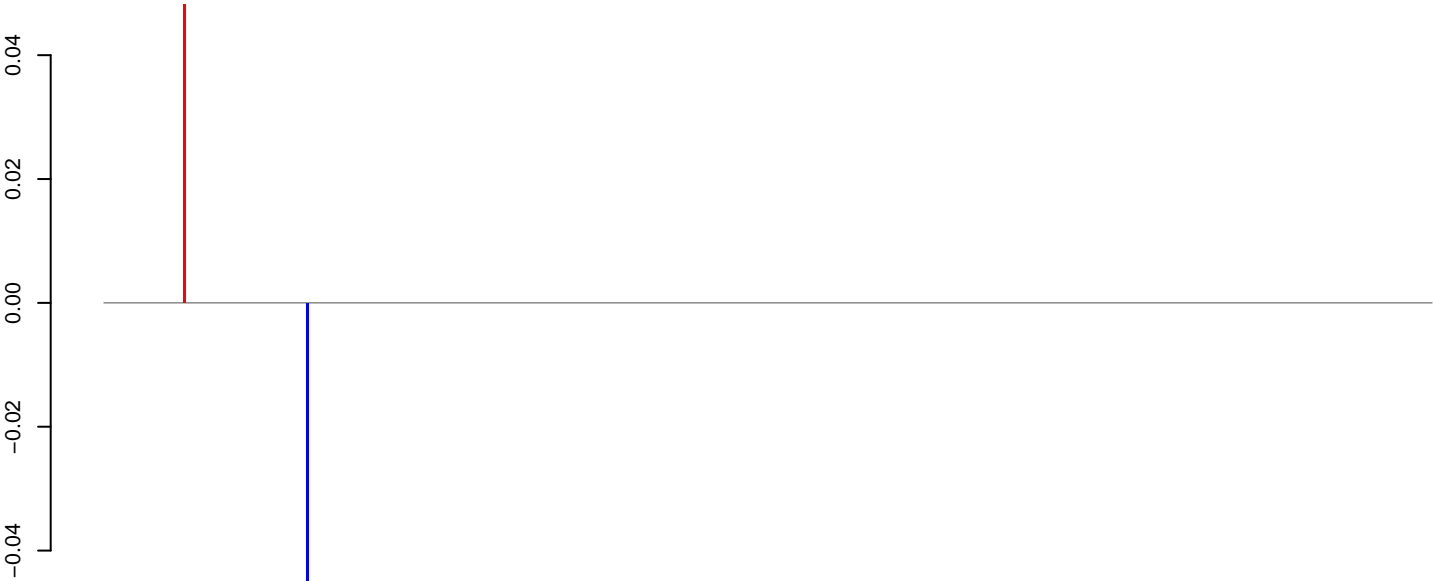
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep



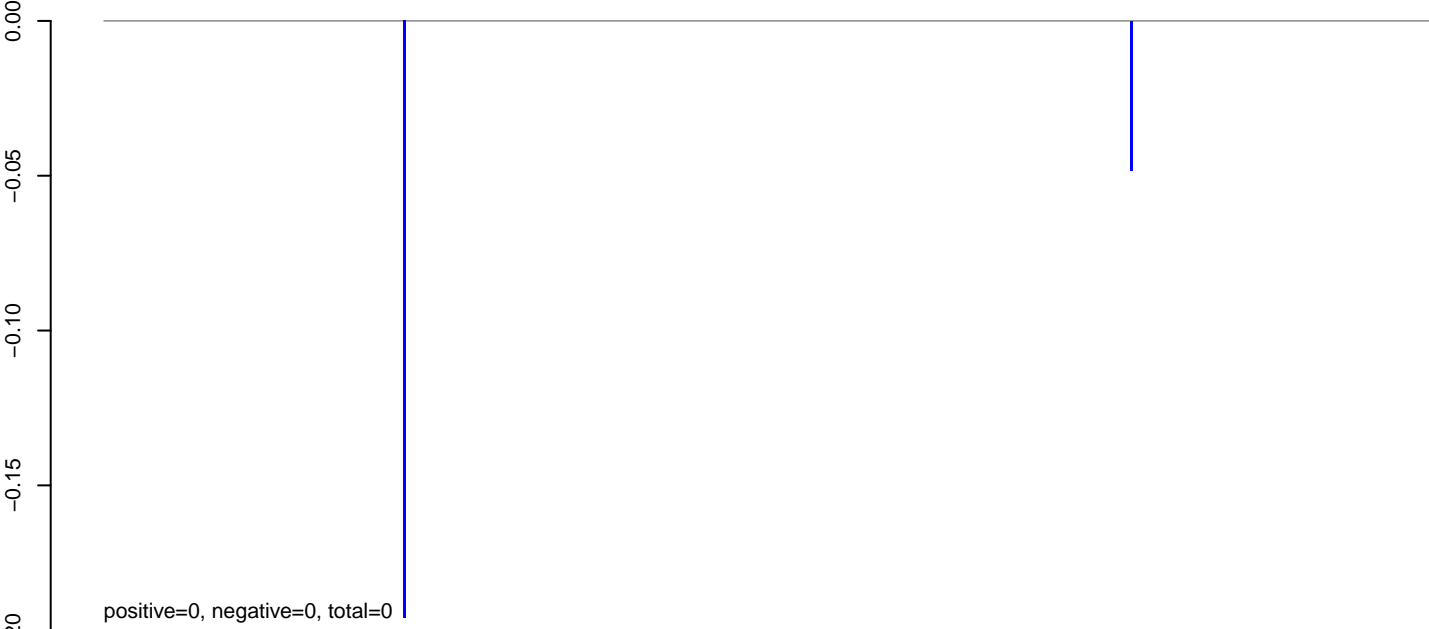
AeAeg_Aag2_NL.M3.rep



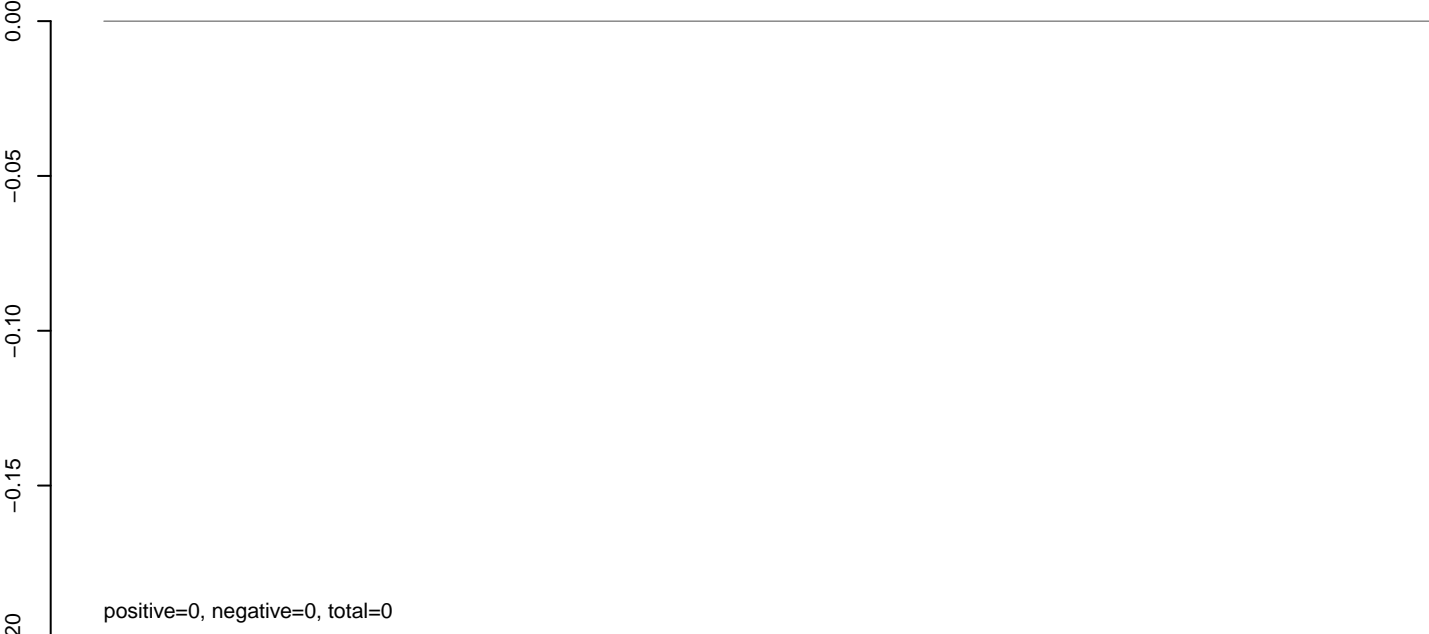
v size=25, time=0, net gain=0, total=0, DUSK=0, 148553.1-Gambie_virus_putative_nucleoprotein_putative_small_transmembrane_protein_putative_g:1-11652

0 2000 4000 6000 8000 10000 12000

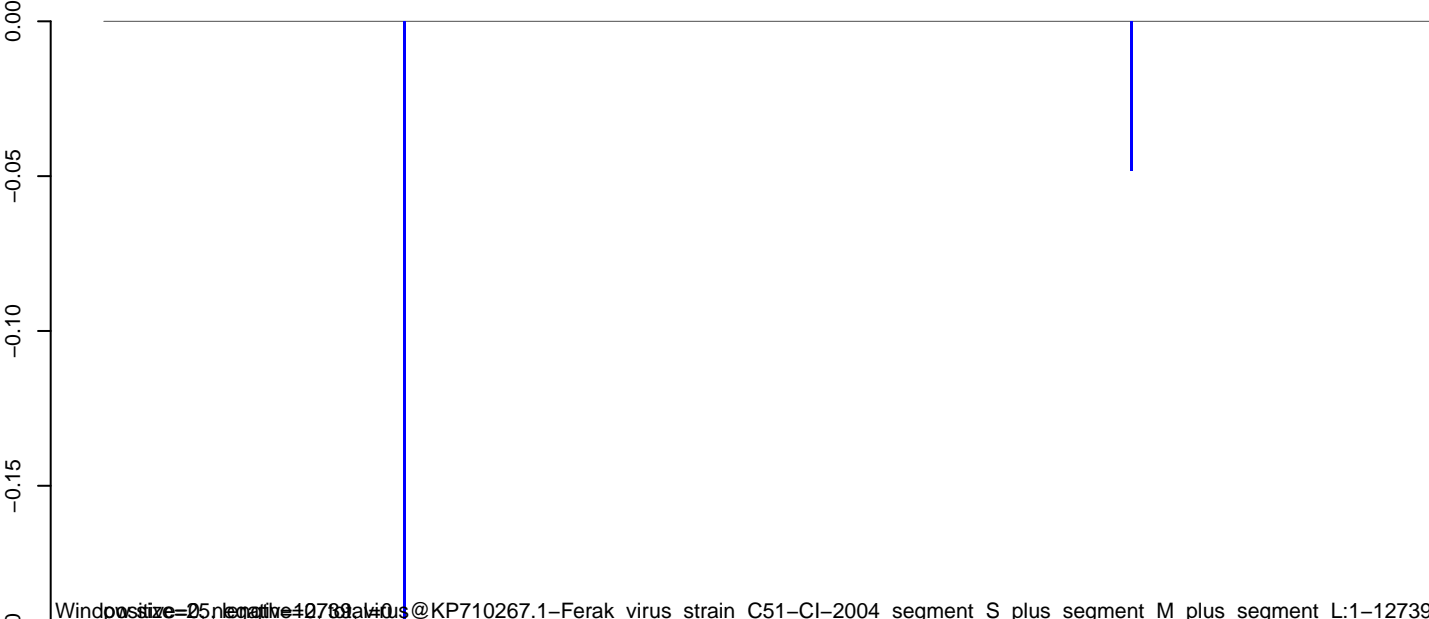
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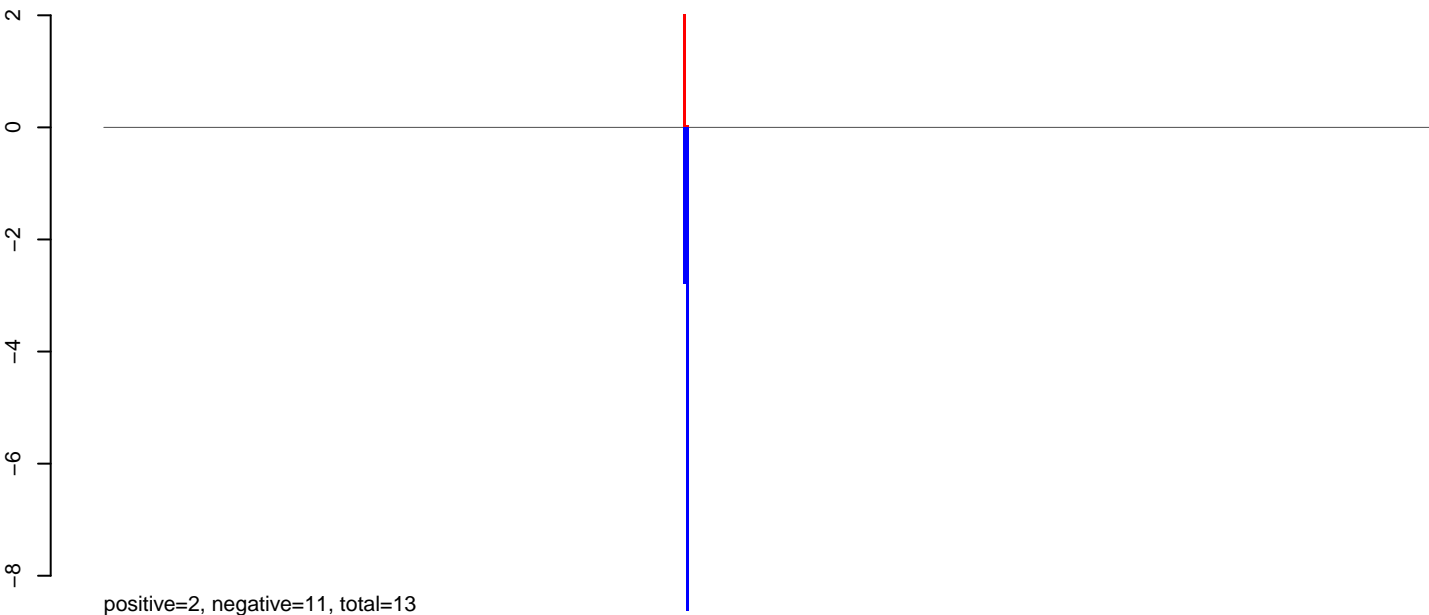
AeAeg_Aag2_NL.M3.24_35.rep



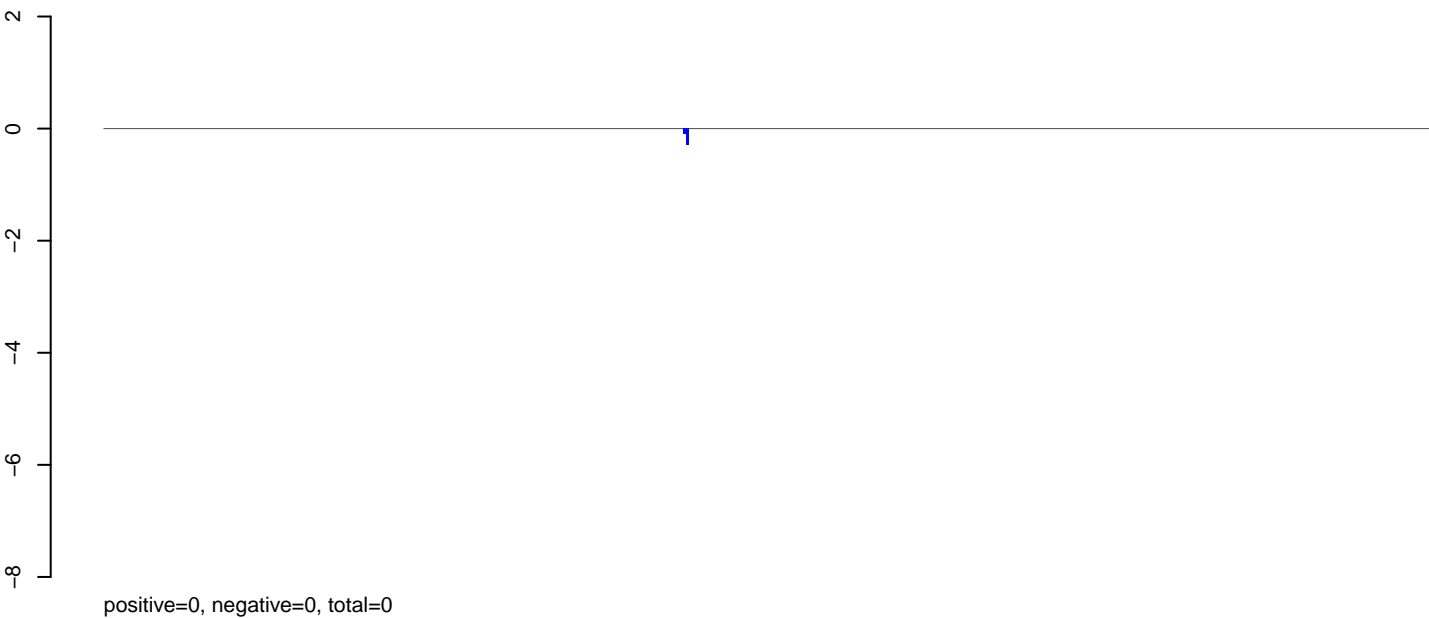
AeAeg_Aag2_NL.M3.rep



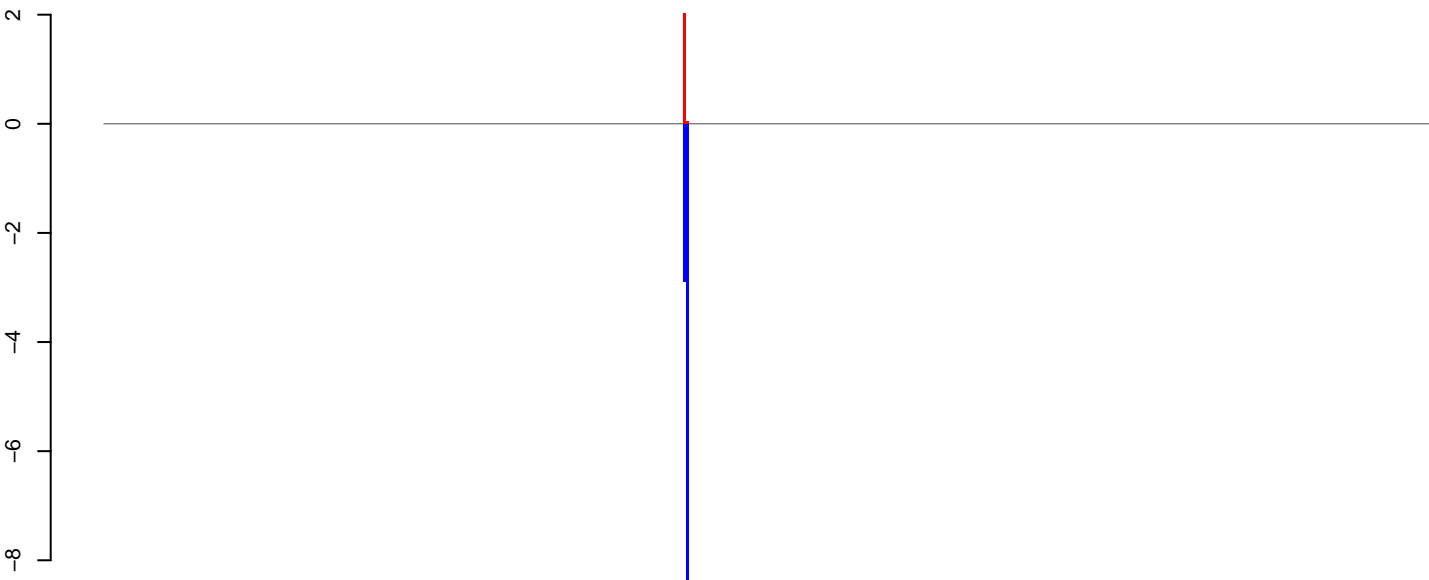
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep



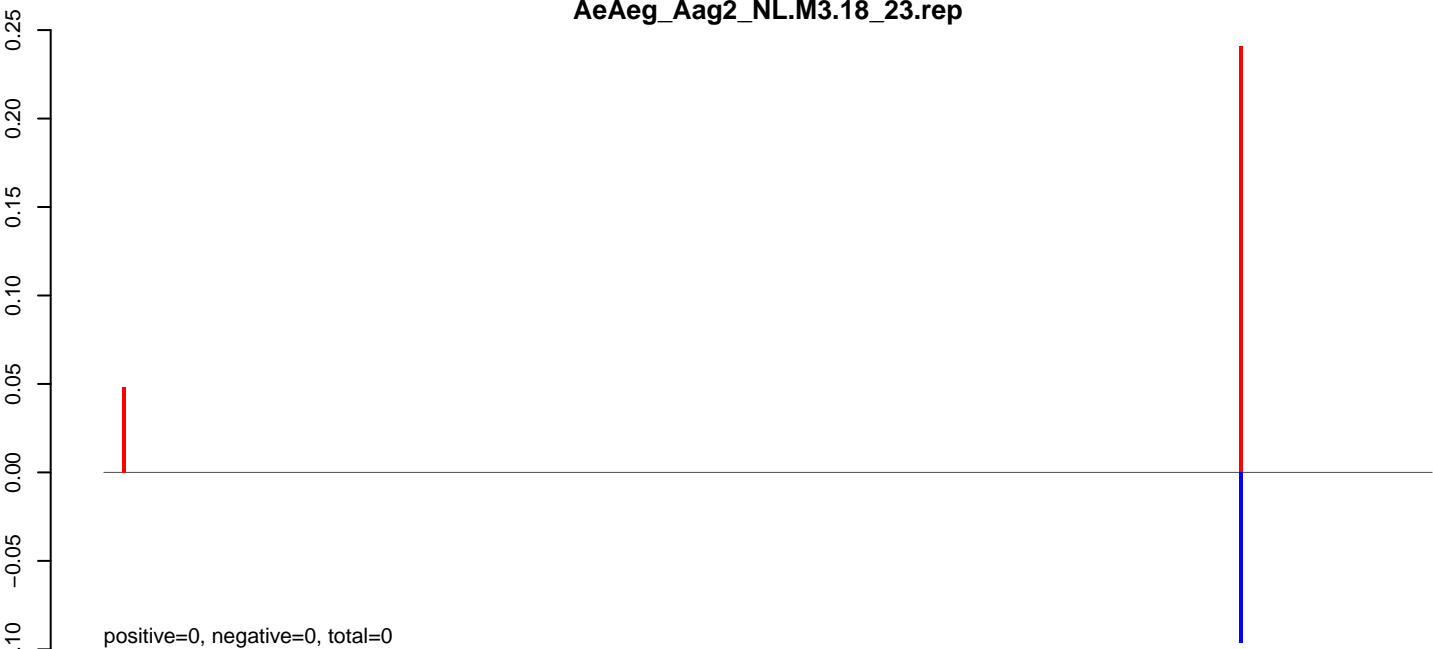
AeAeg_Aag2_NL.M3.rep



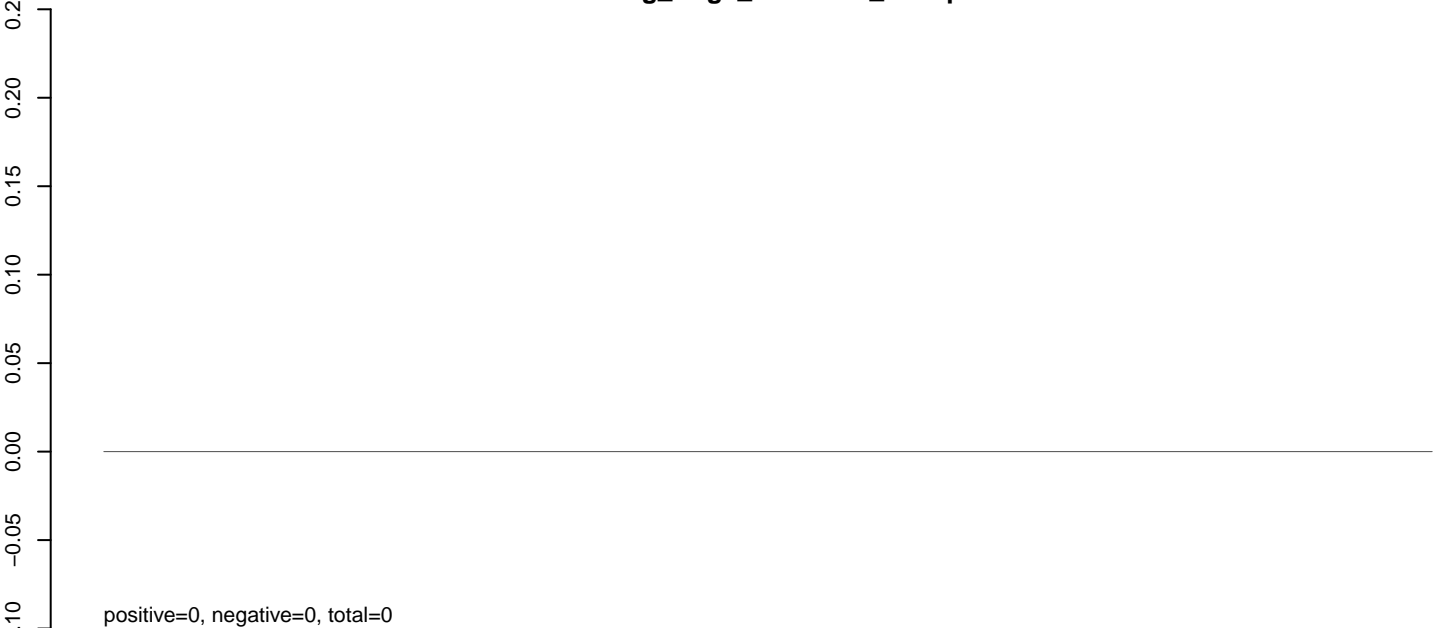
Wad2, size=25, length=2369, virus@KJ716848.1-Ngari_virus_strain_Adrar_segment_S_plus_segment_M_plus_segment_L:1-12369

0 2000 4000 6000 8000 10000 12000

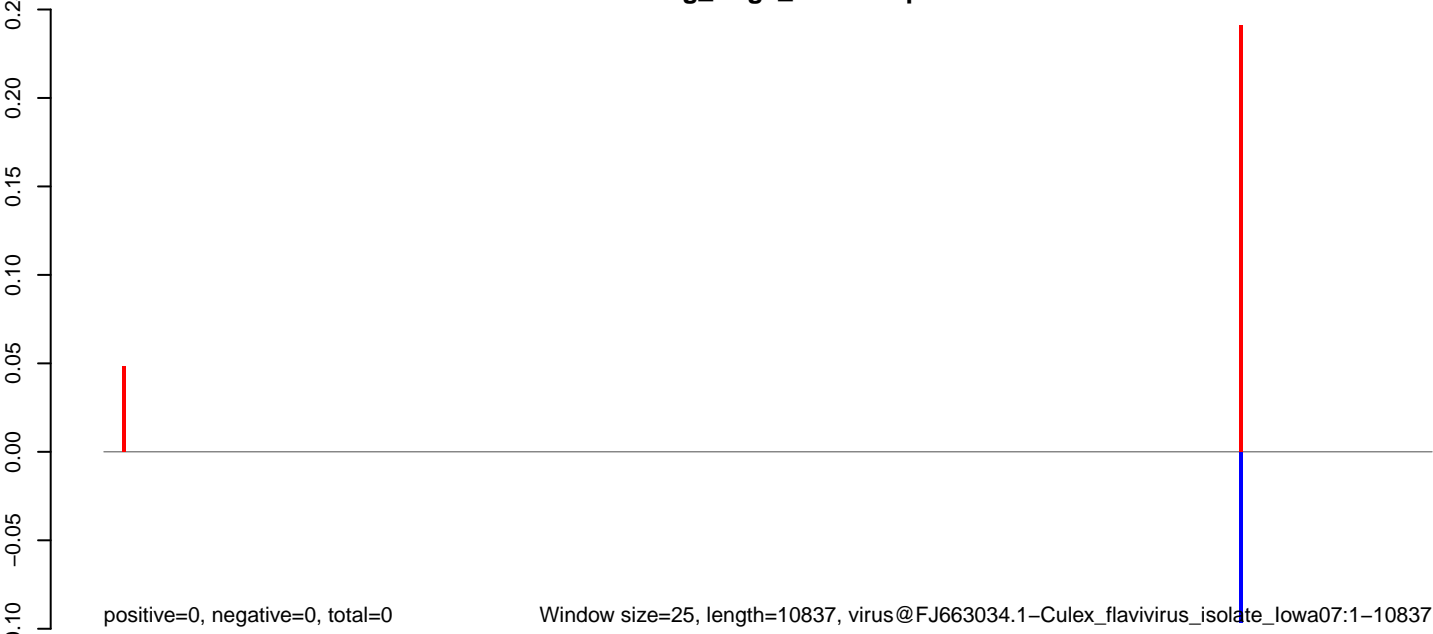
AeAeg_Aag2_NL.M3.18_23.rep

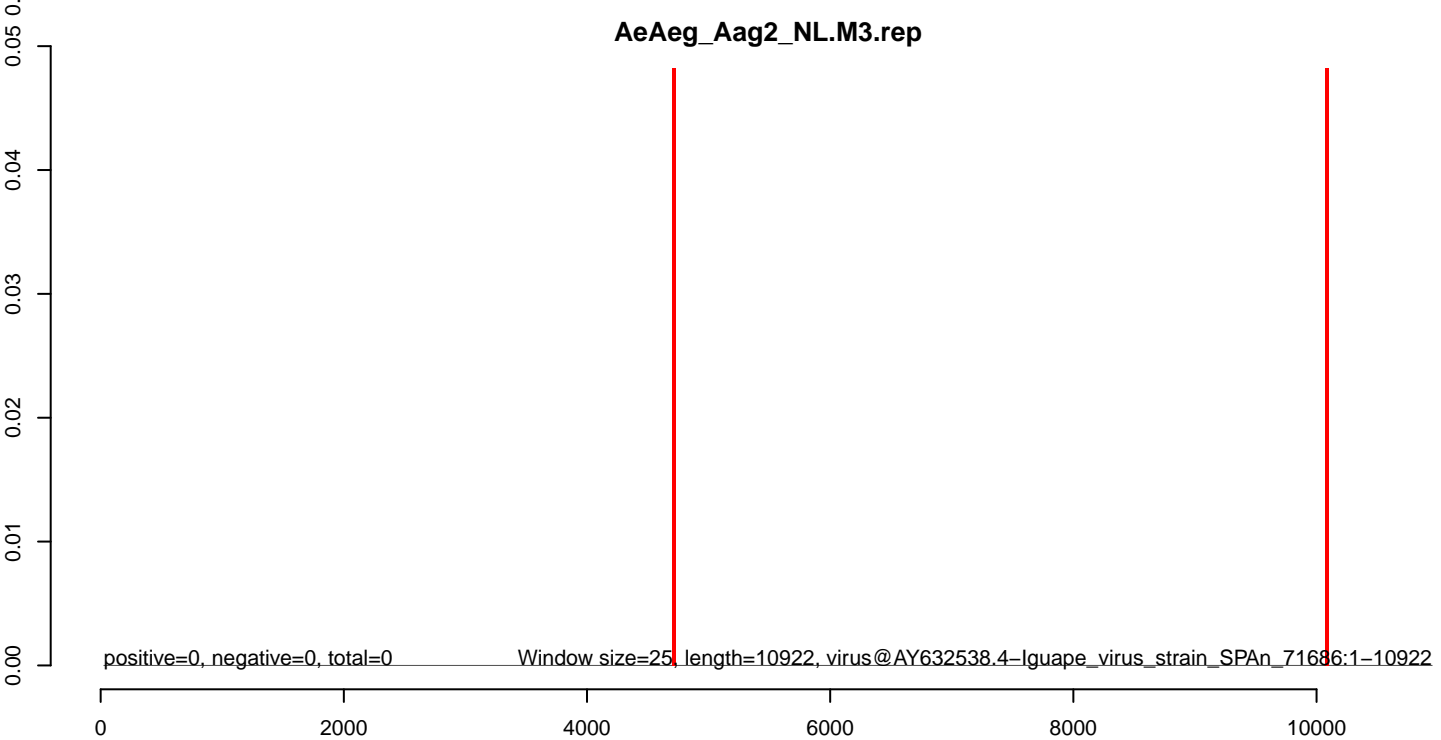
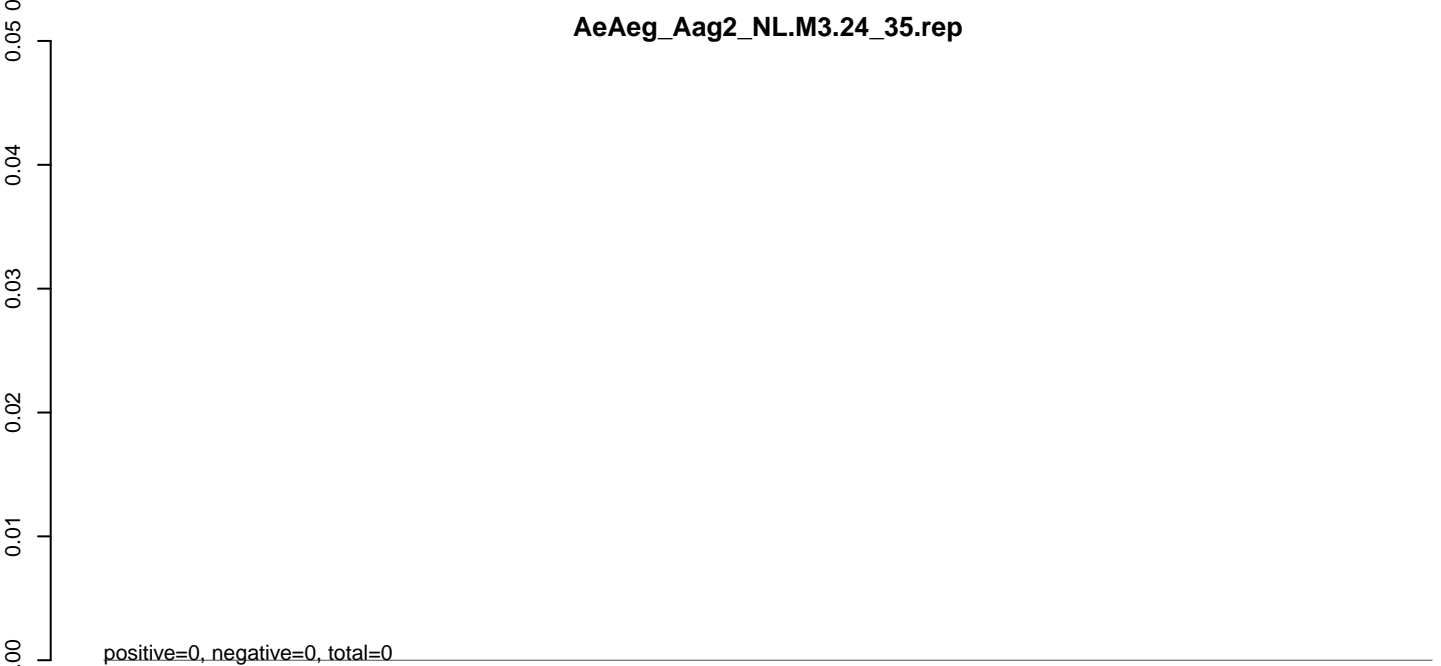
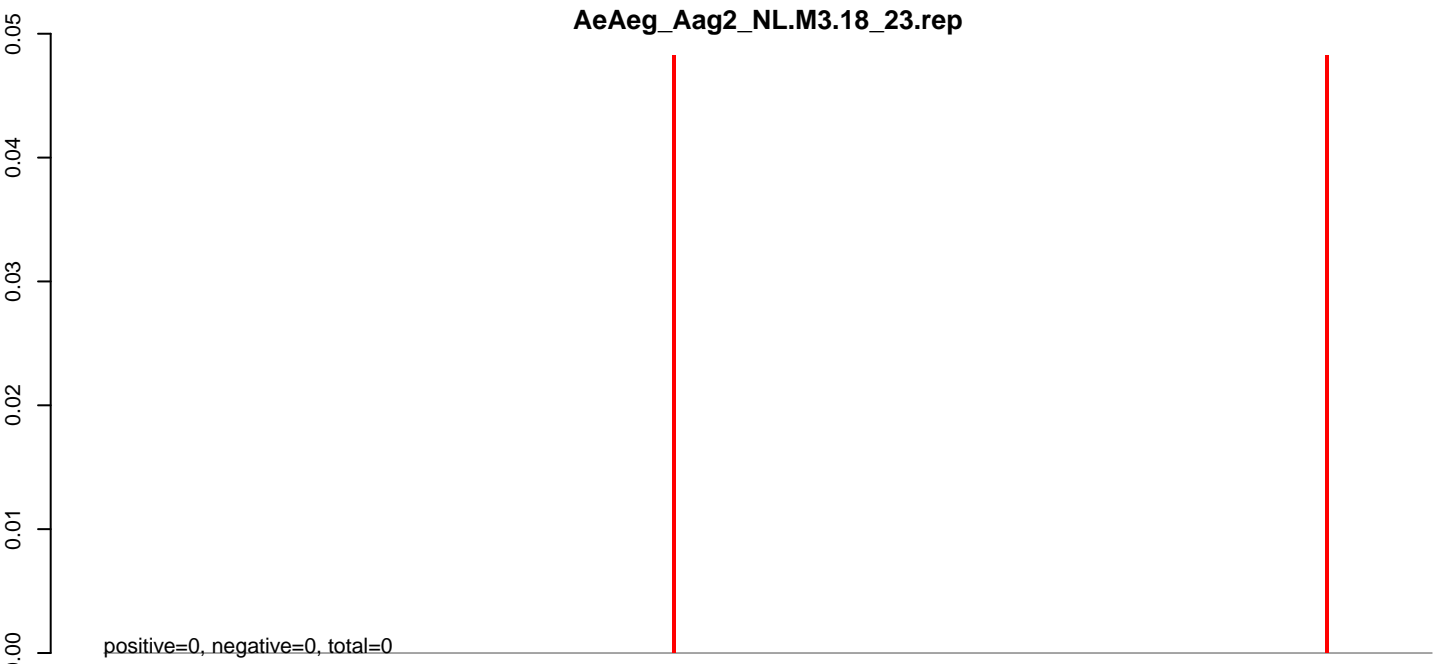


AeAeg_Aag2_NL.M3.24_35.rep

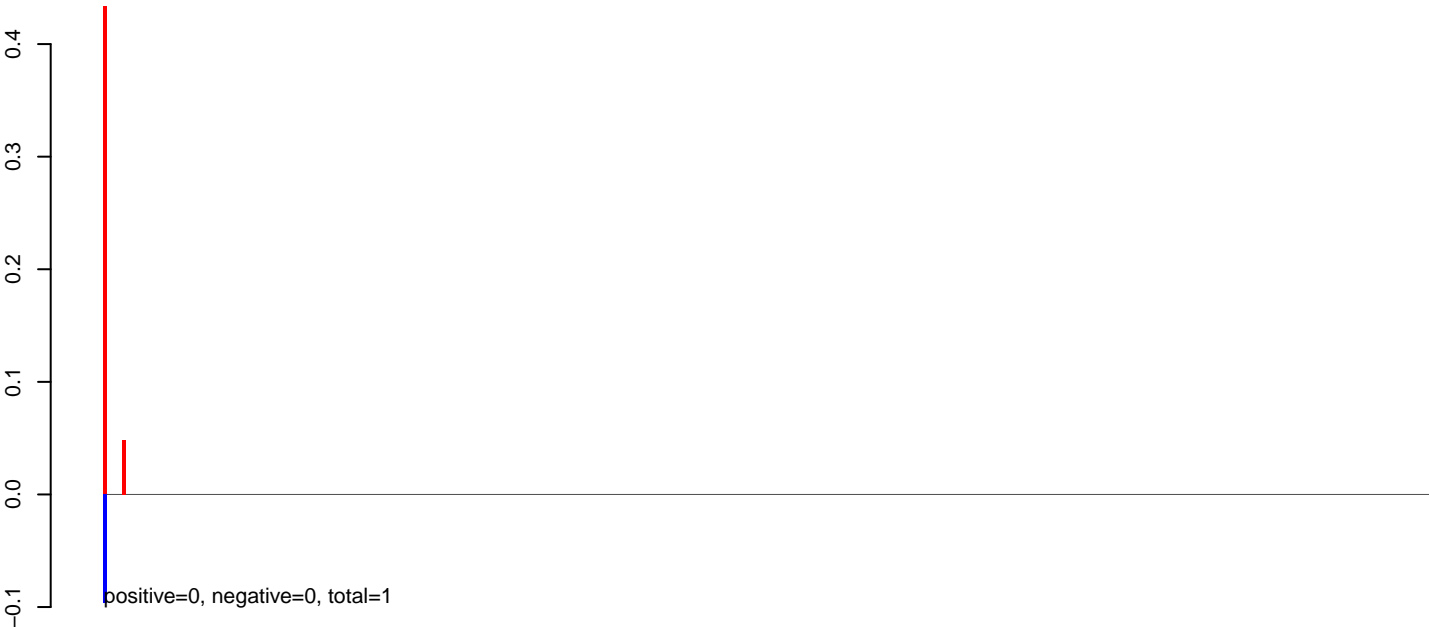


AeAeg_Aag2_NL.M3.rep

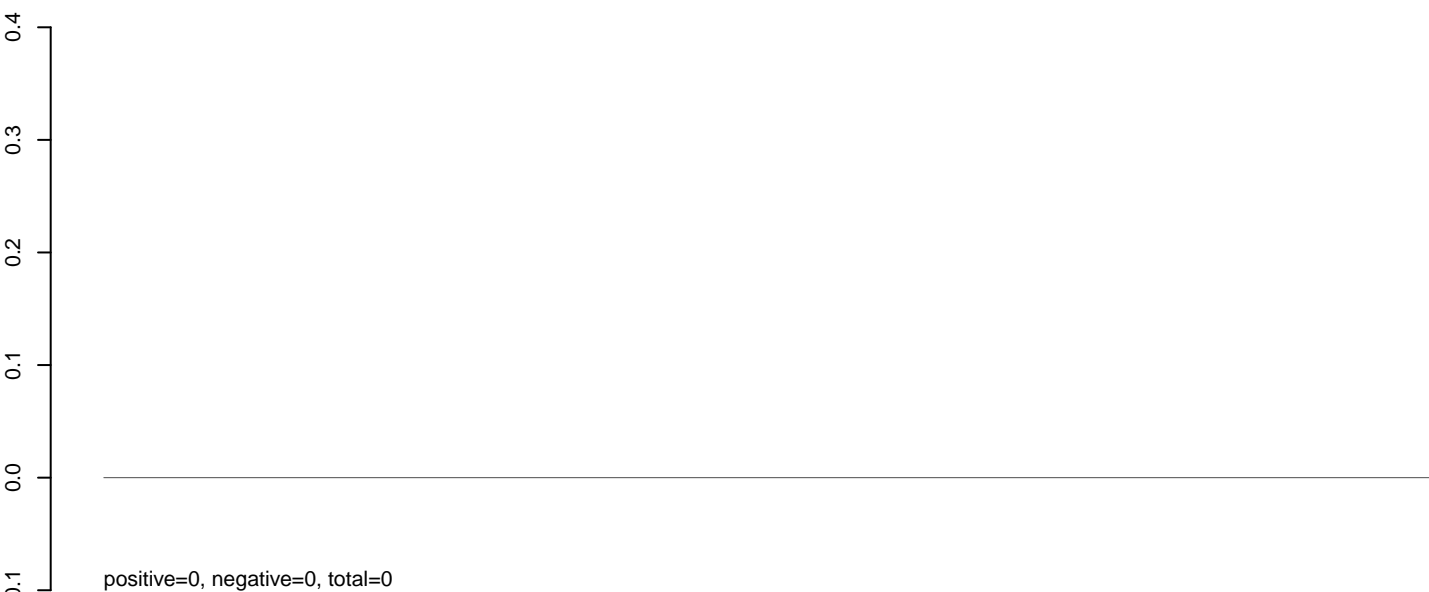




AeAeg_Aag2_NL.M3.18_23.rep



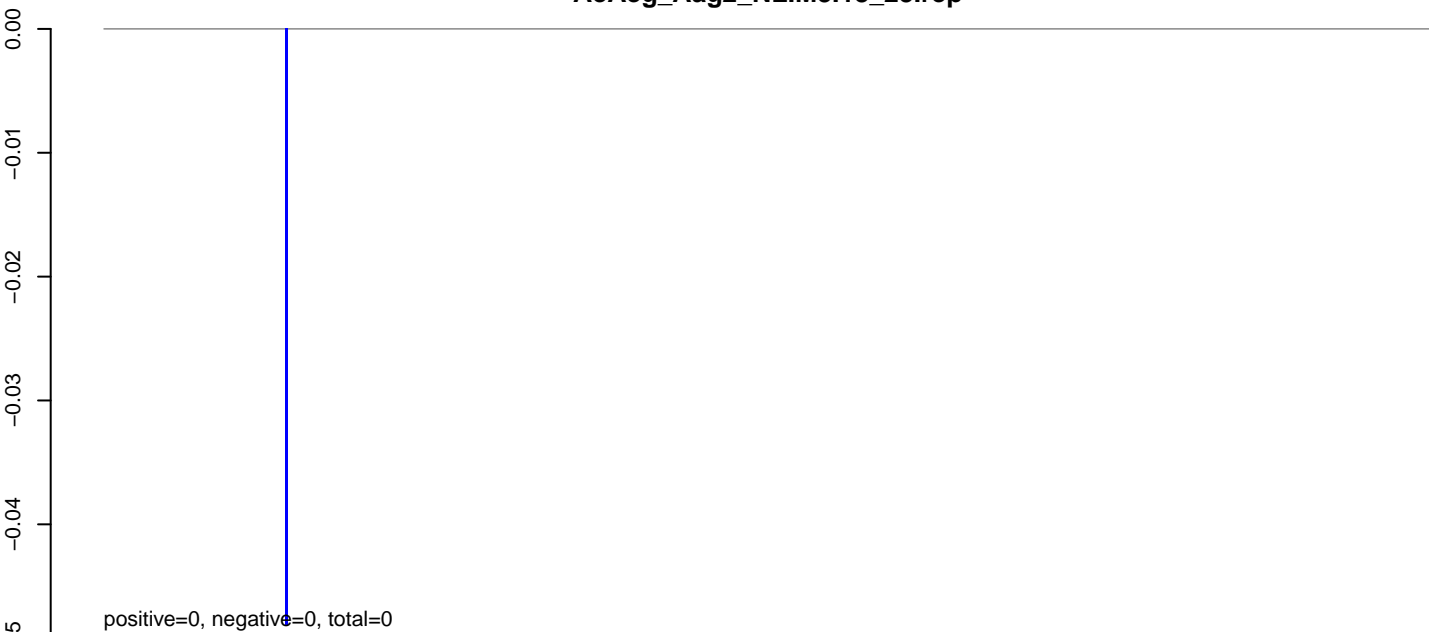
AeAeg_Aag2_NL.M3.24_35.rep



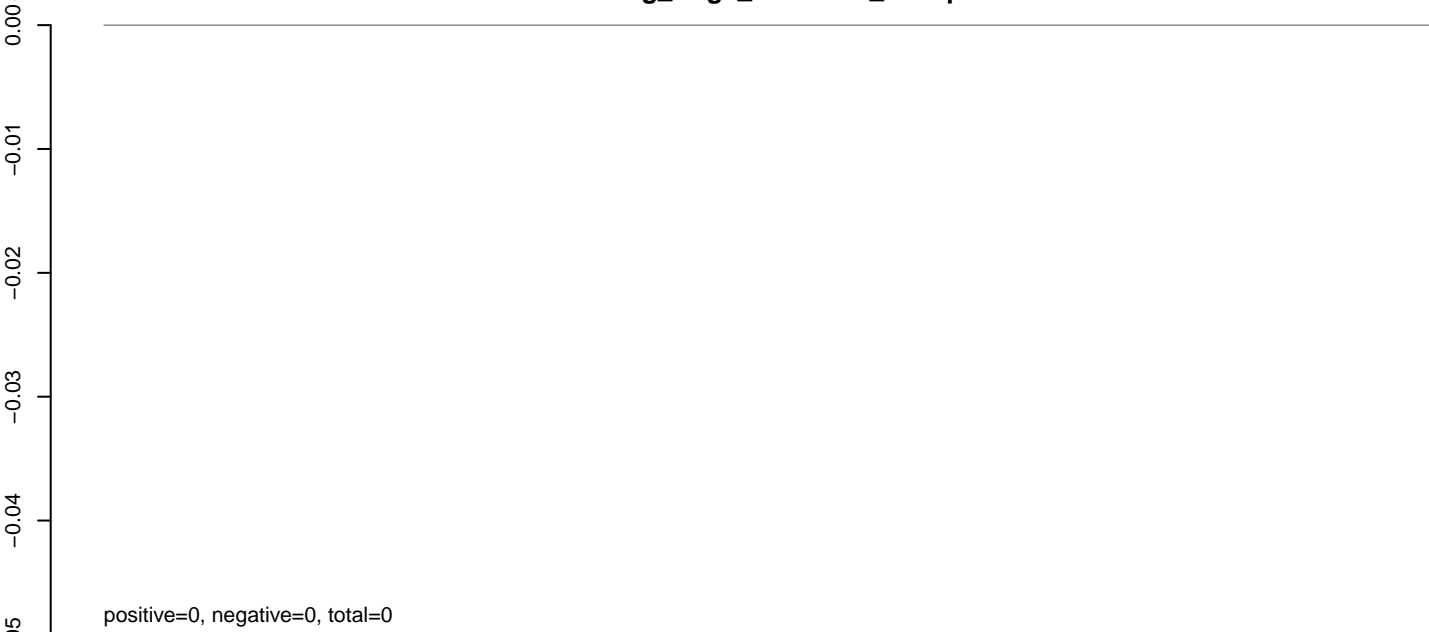
AeAeg_Aag2_NL.M3.rep



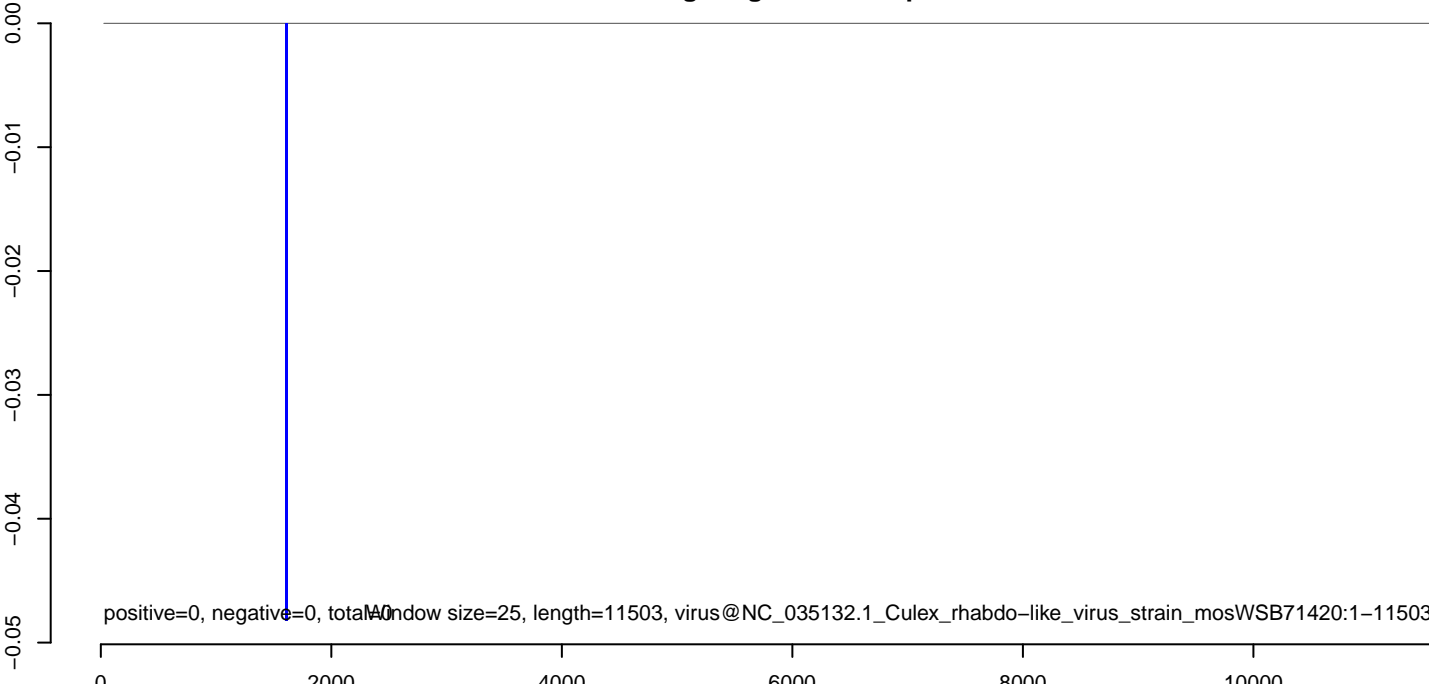
AeAeg_Aag2_NL.M3.18_23.rep

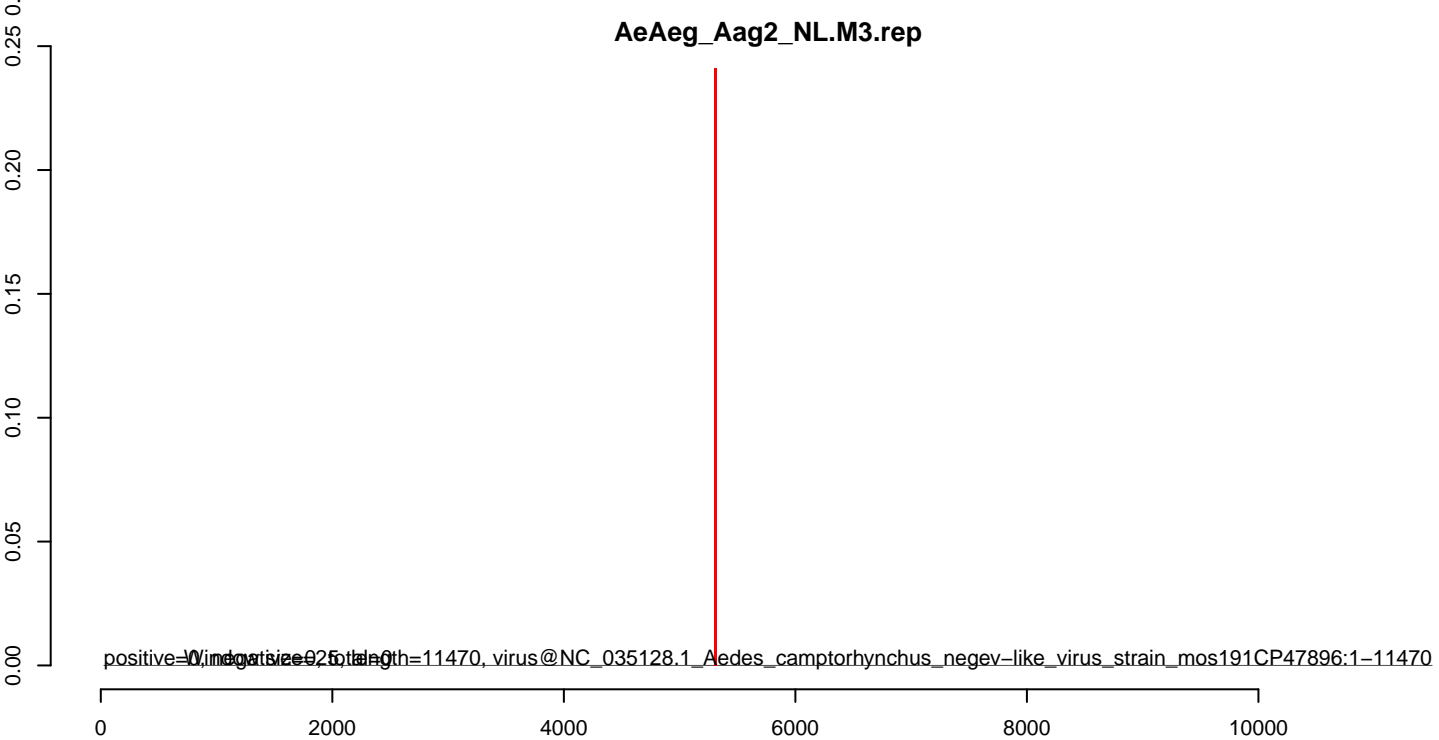
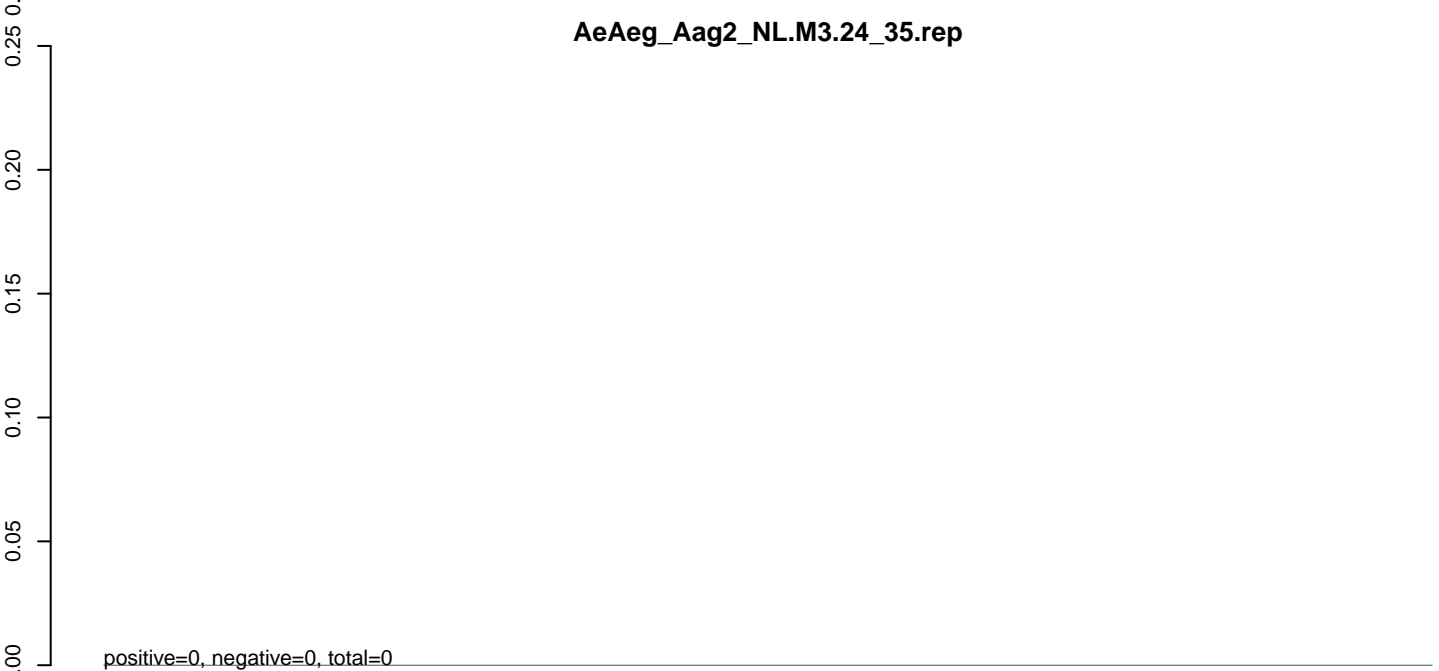
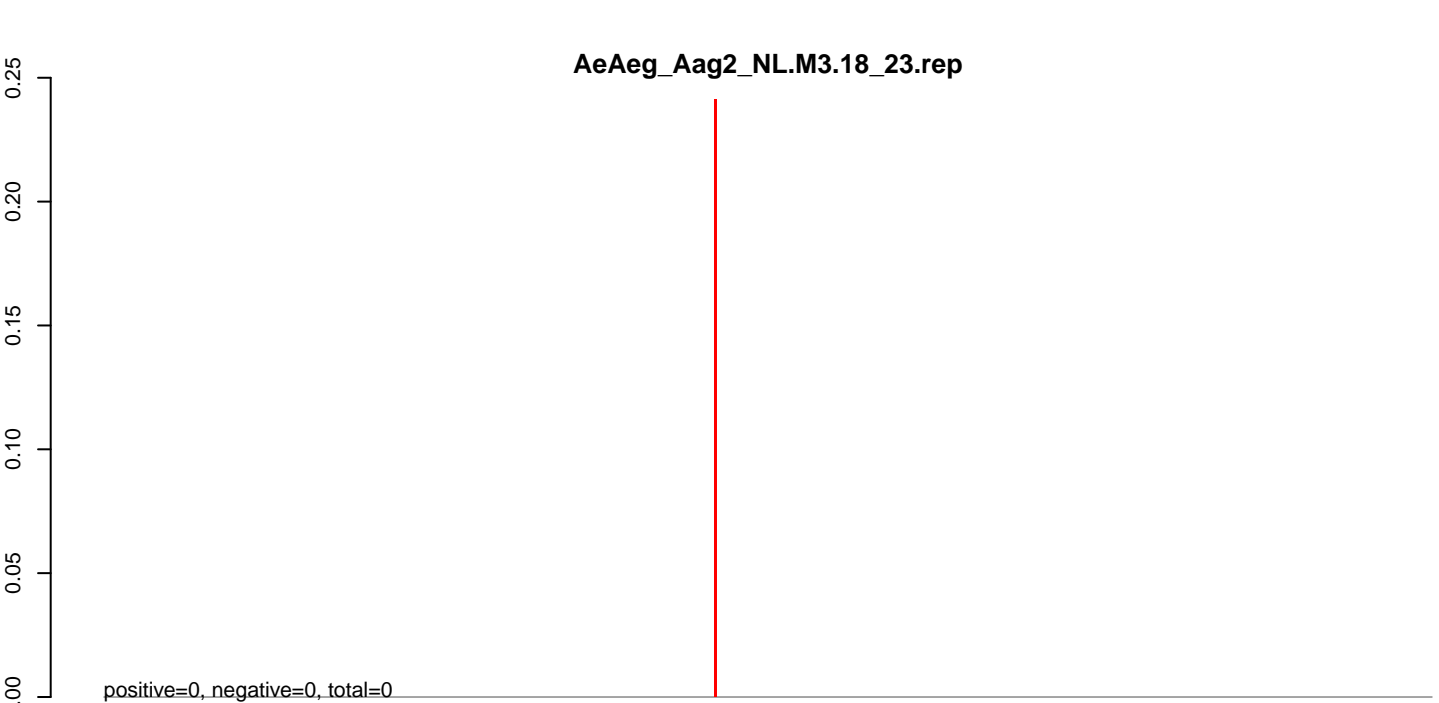


AeAeg_Aag2_NL.M3.24_35.rep

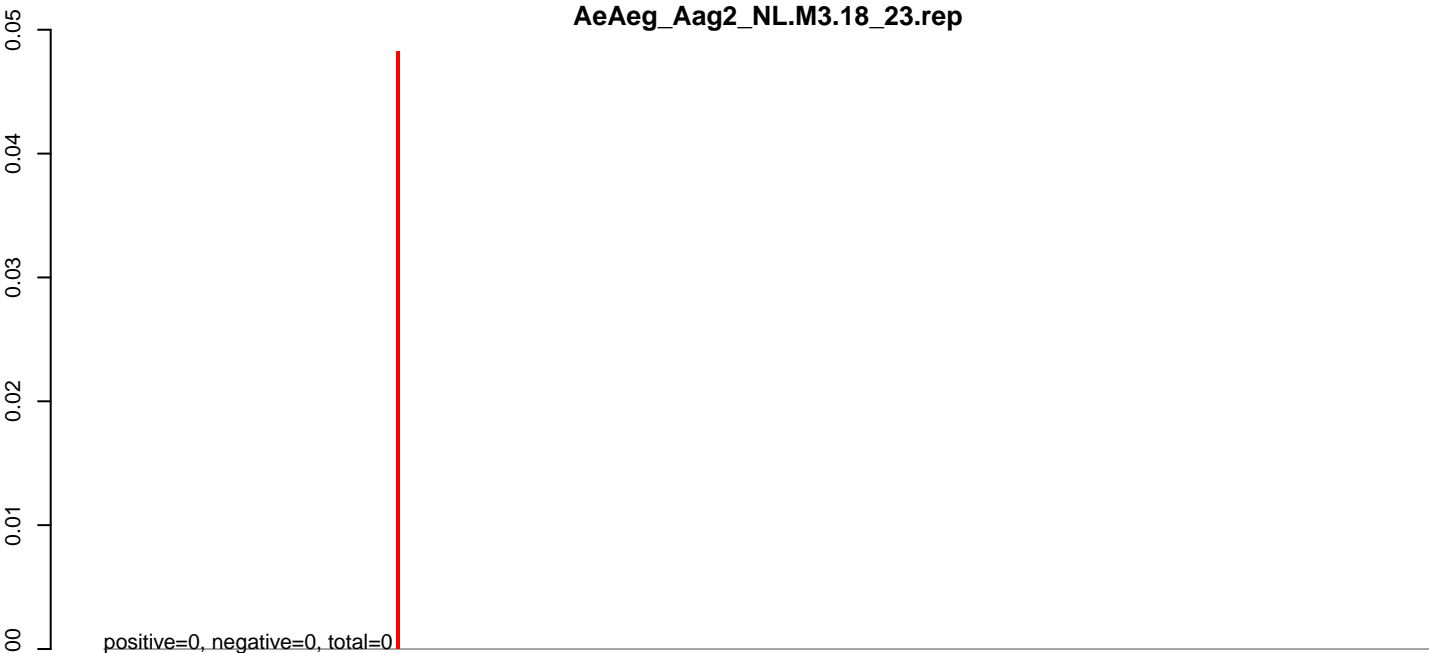


AeAeg_Aag2_NL.M3.rep





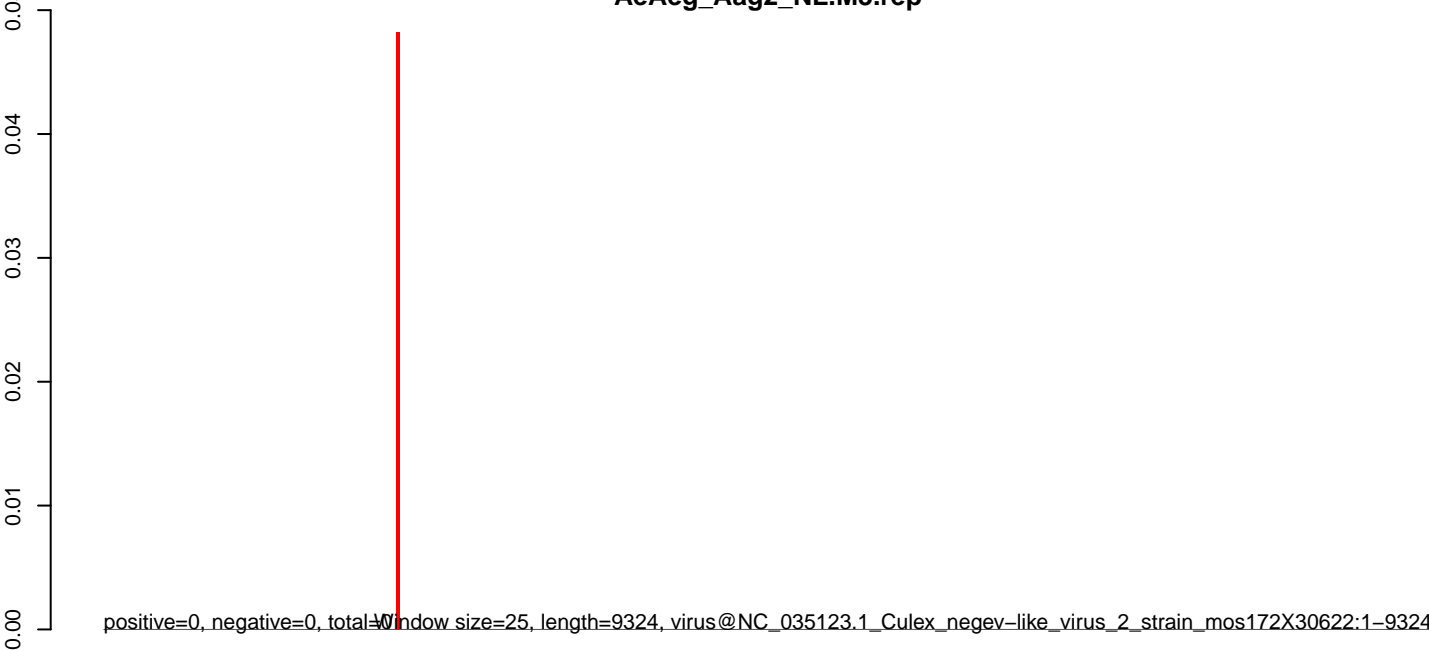
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep



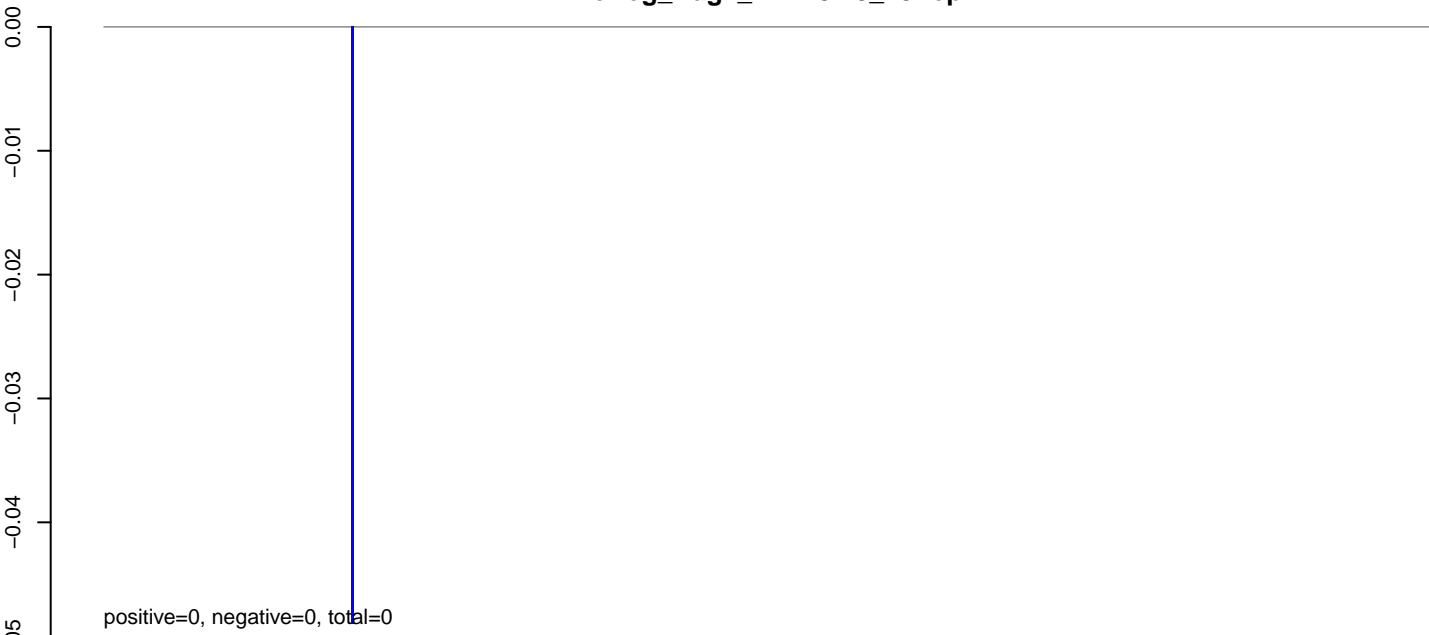
AeAeg_Aag2_NL.M3.rep



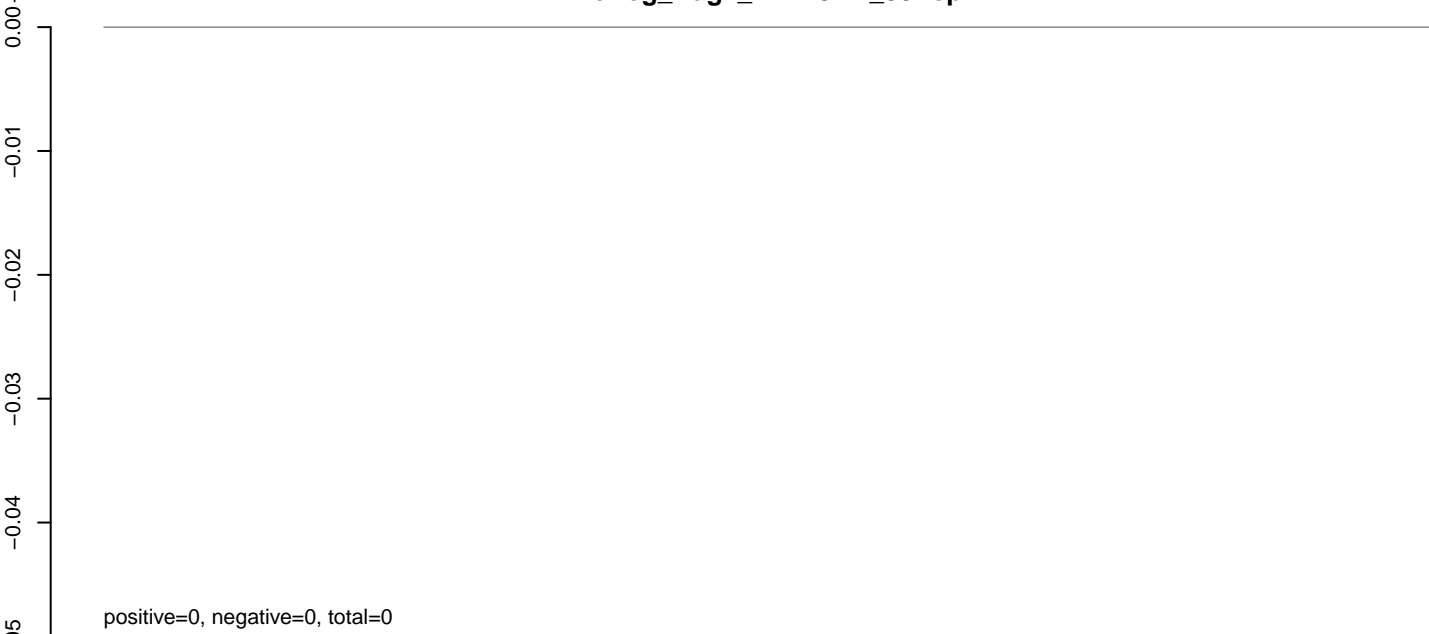
Window size=25, length=9324, virus@NC_035123.1_Culex_negev-like_virus_2_strain_mos172X30622:1-9324

0 2000 4000 6000 8000

AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep



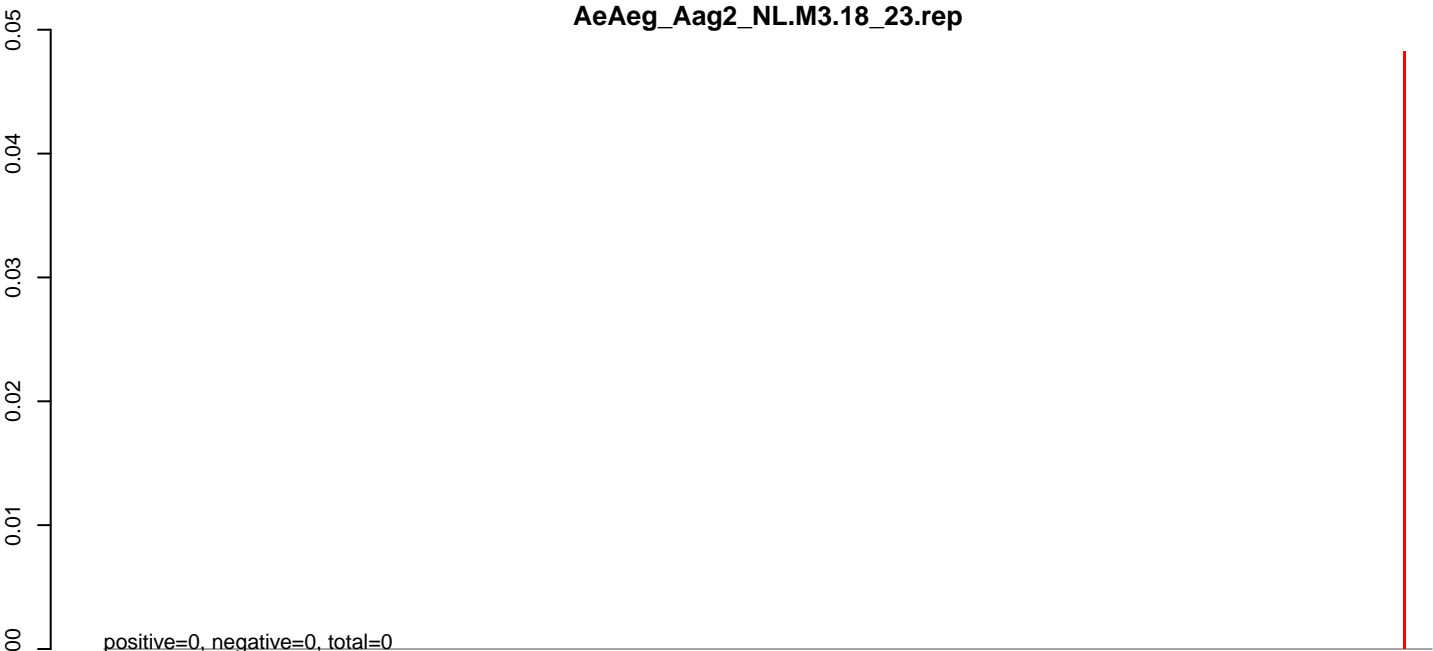
AeAeg_Aag2_NL.M3.rep



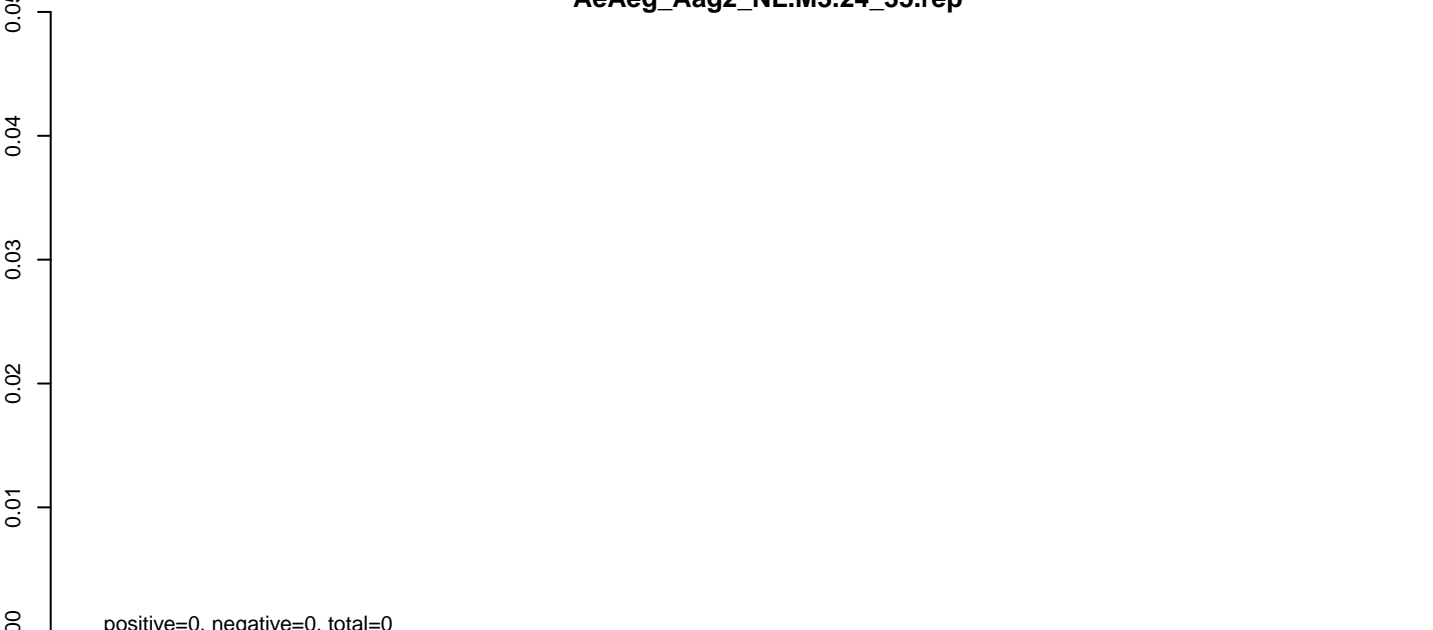
NC_034538.1-Joinjakaka_virus_nucleoprotein_phosphoprotein_hypothetical_protein_matrix_glycopro:1-13155

0 2000 4000 6000 8000 10000 12000

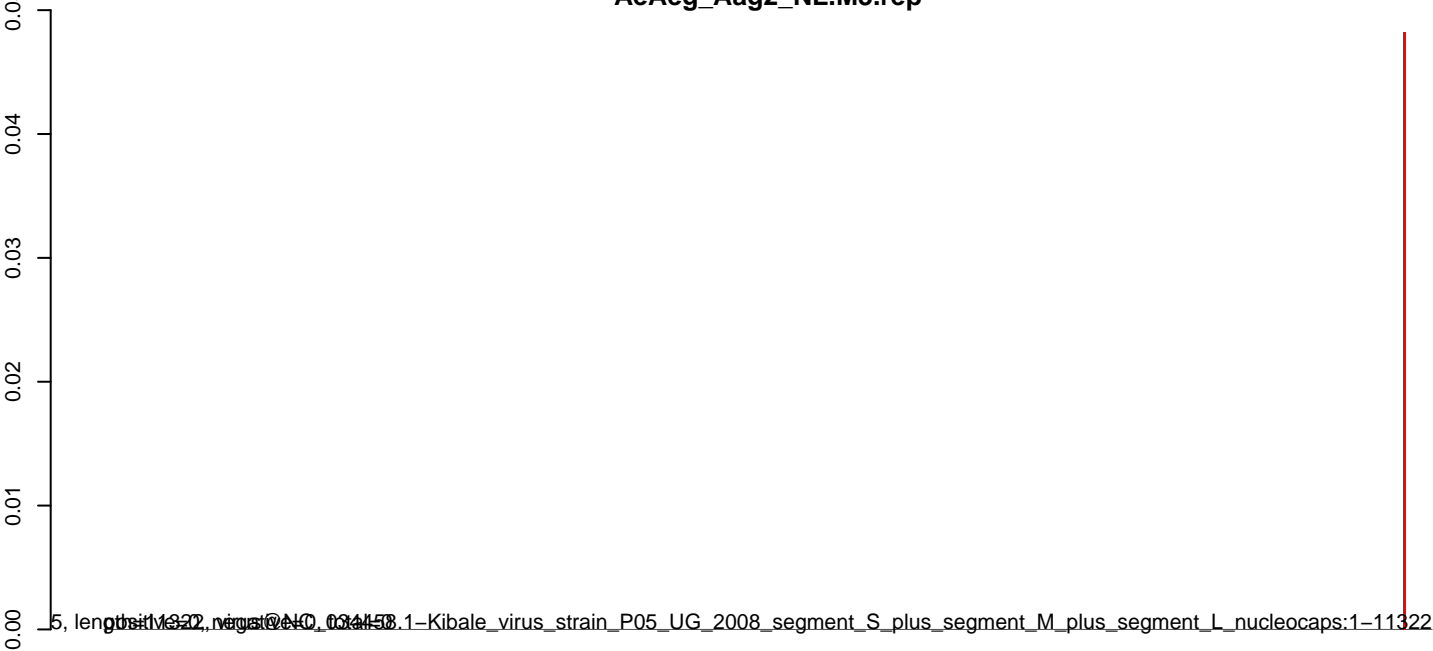
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep

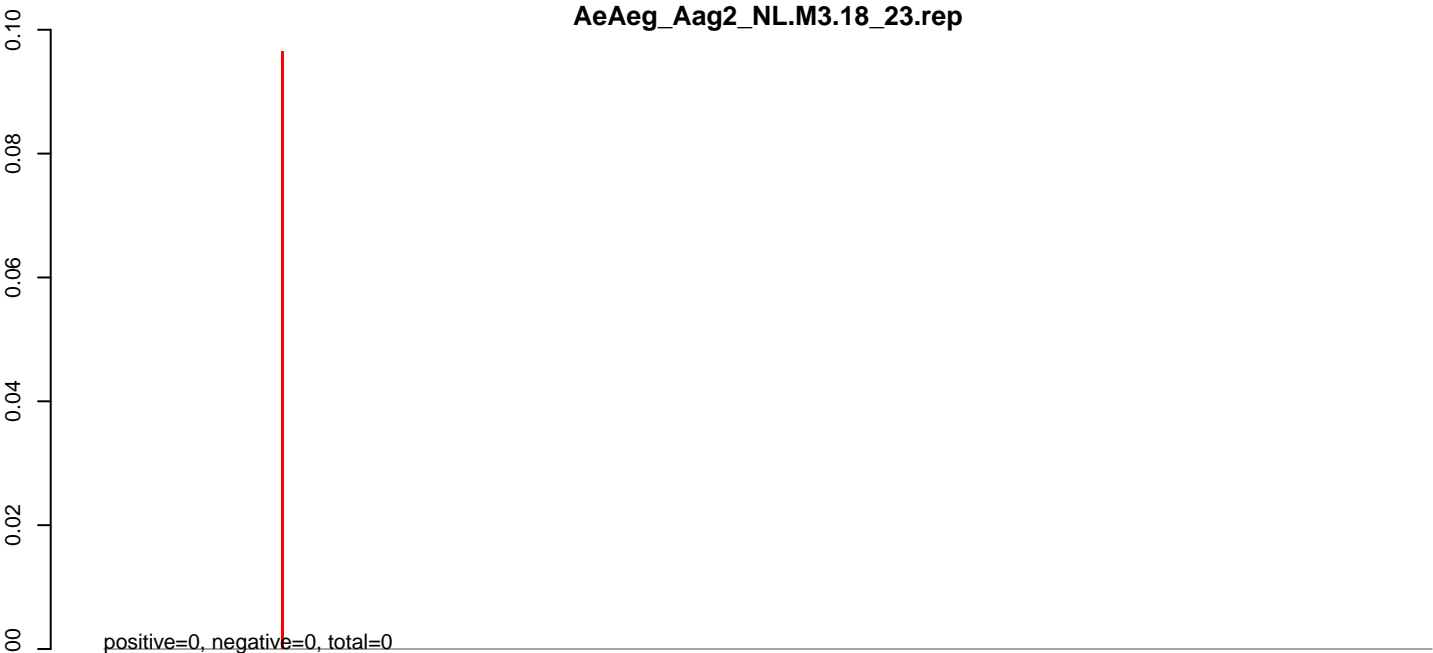


AeAeg_Aag2_NL.M3.rep

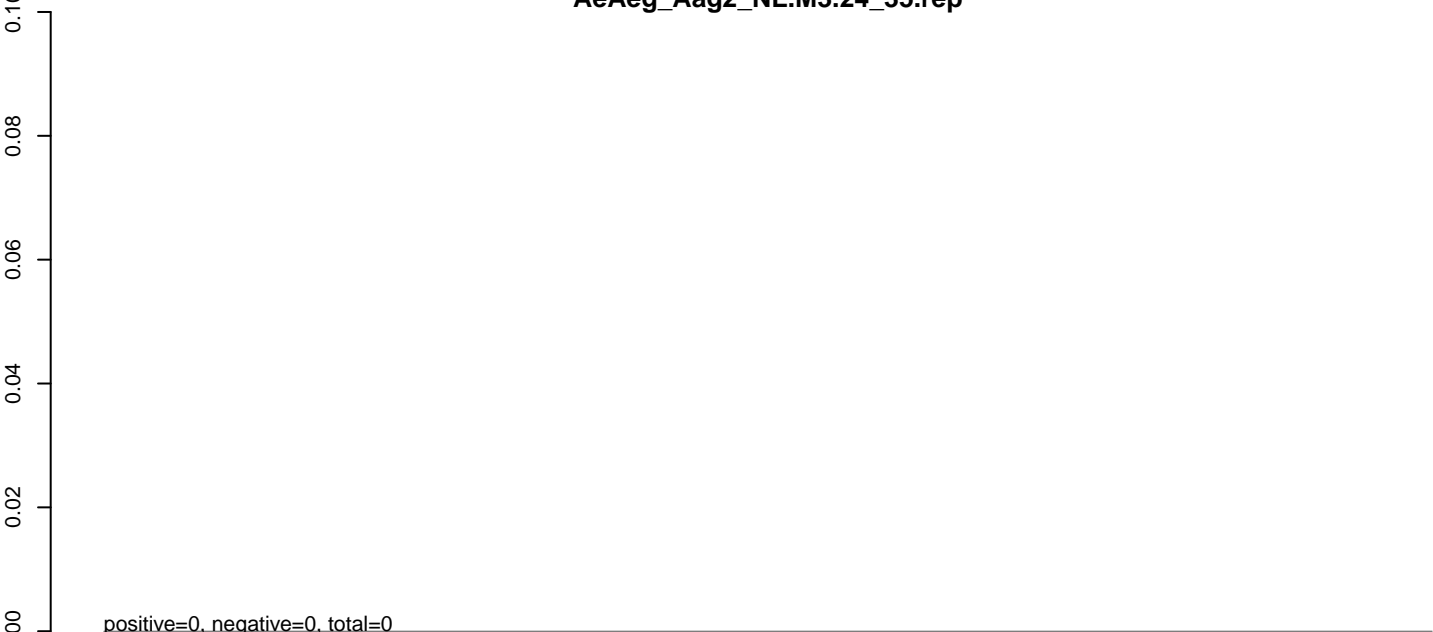


0 2000 4000 6000 8000 10000

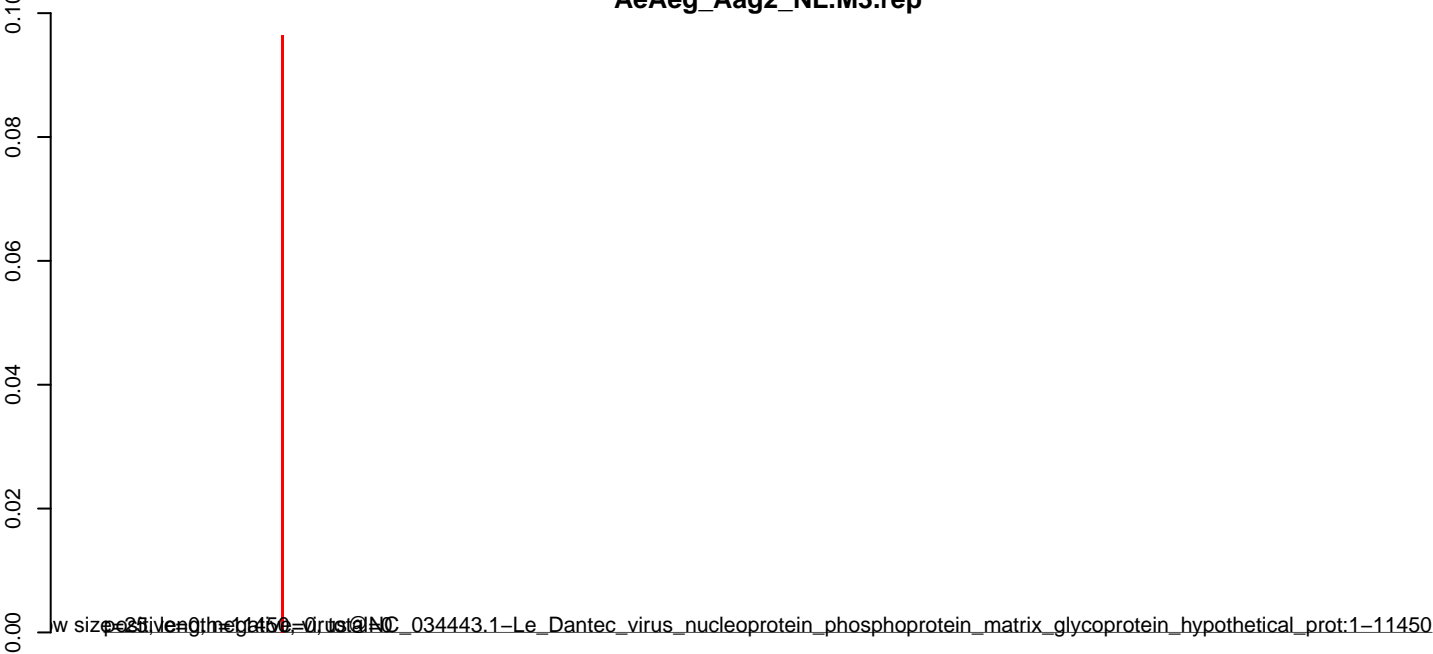
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep



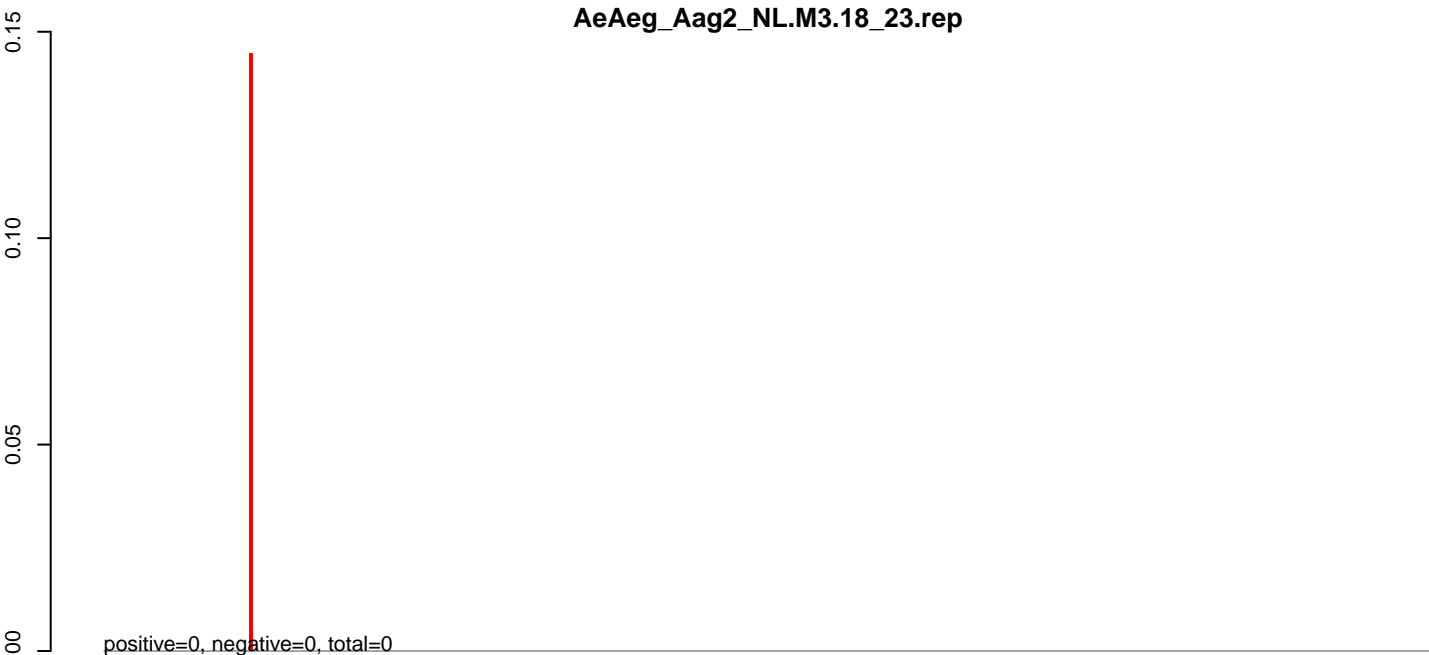
AeAeg_Aag2_NL.M3.rep



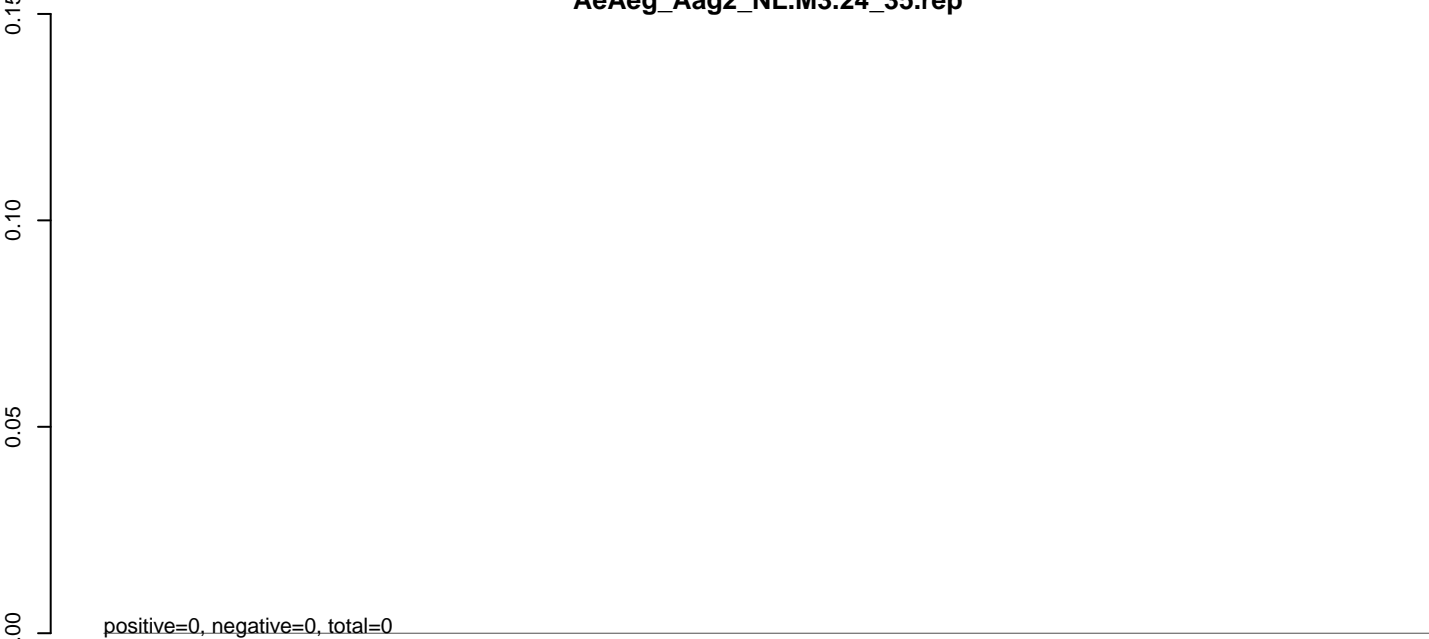
sw size=251 length=1456 @NC_034443.1-Le_Dantec_virus_nucleoprotein_phosphoprotein_matrix_glycoprotein_hypothetical_prot:1-11450

0 2000 4000 6000 8000 10000

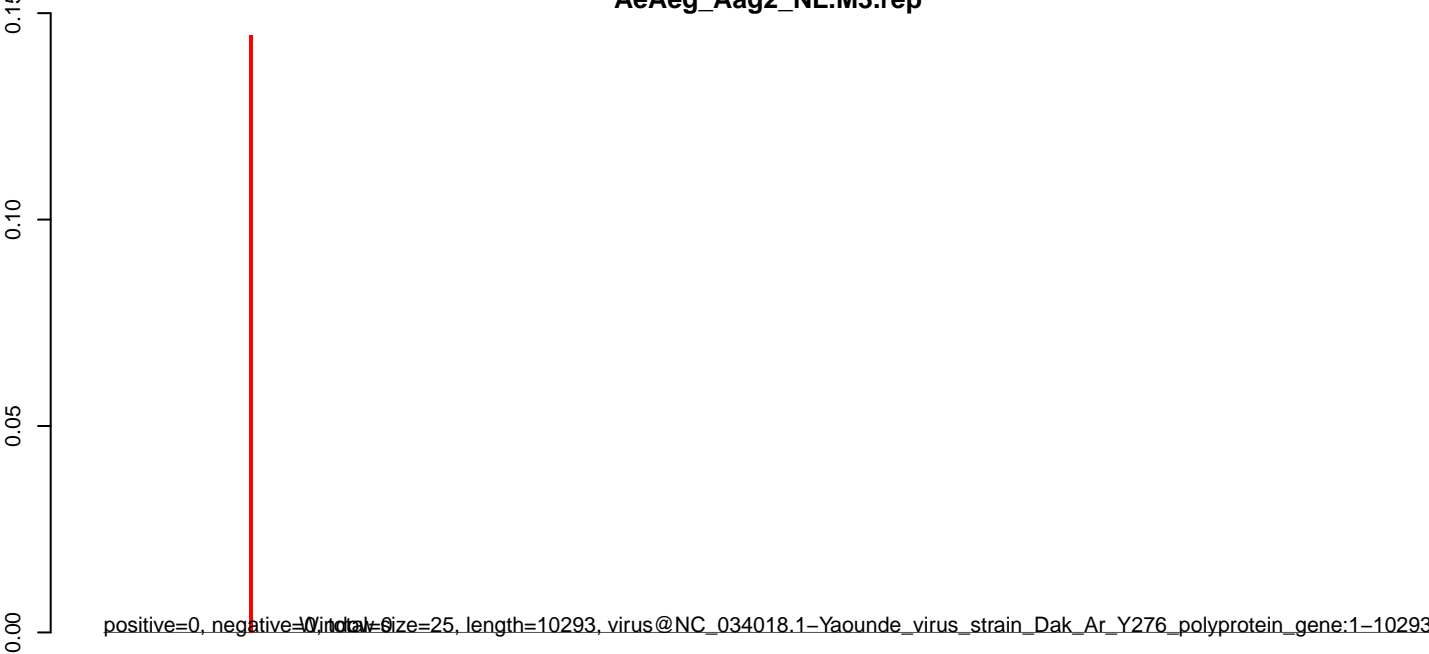
AeAeg_Aag2_NL.M3.18_23.rep



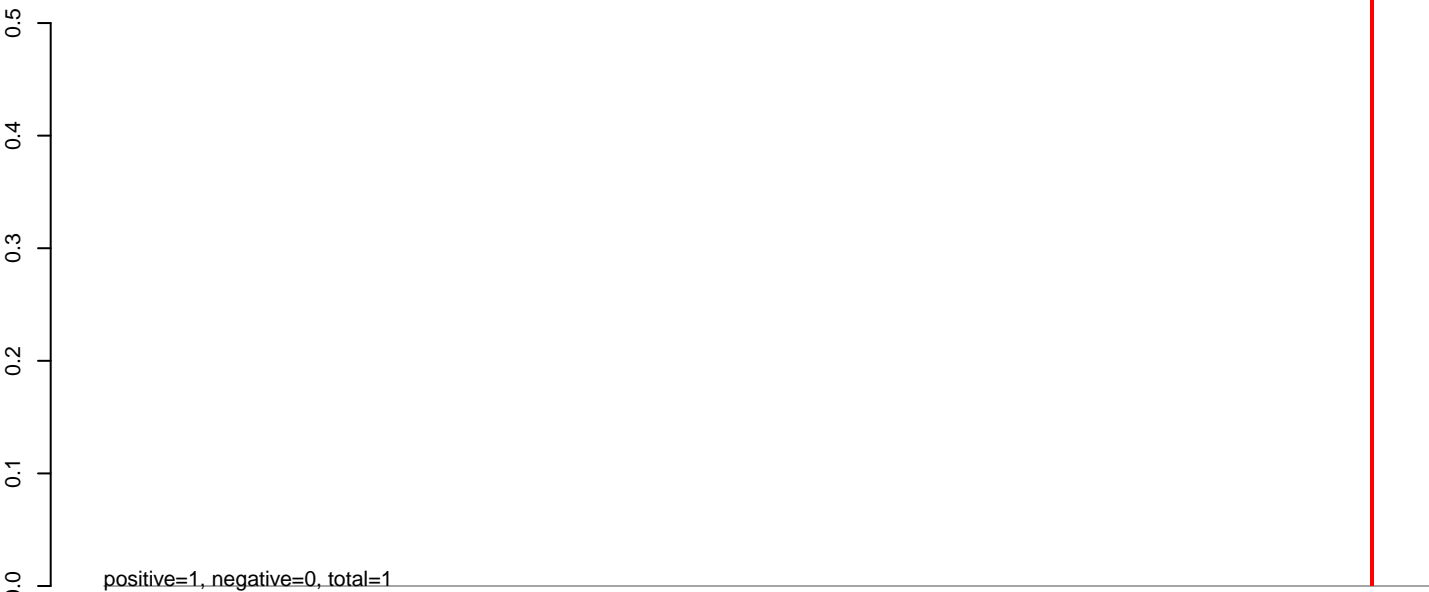
AeAeg_Aag2_NL.M3.24_35.rep



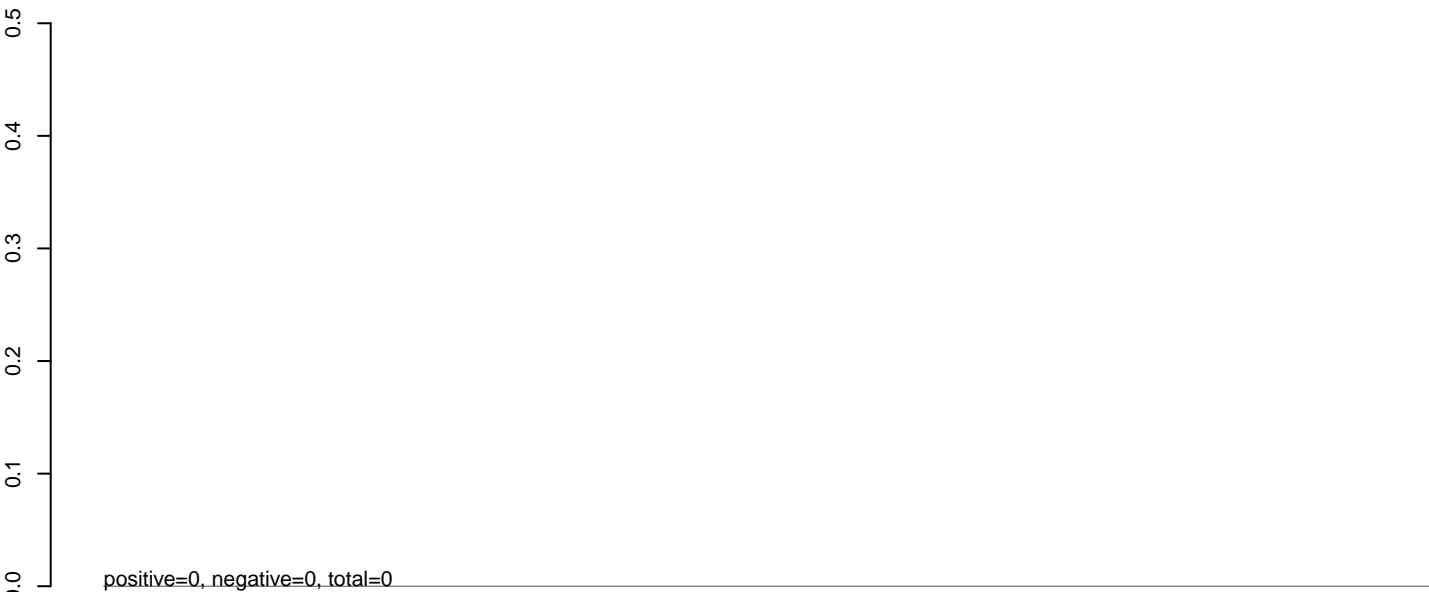
AeAeg_Aag2_NL.M3.rep



AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep

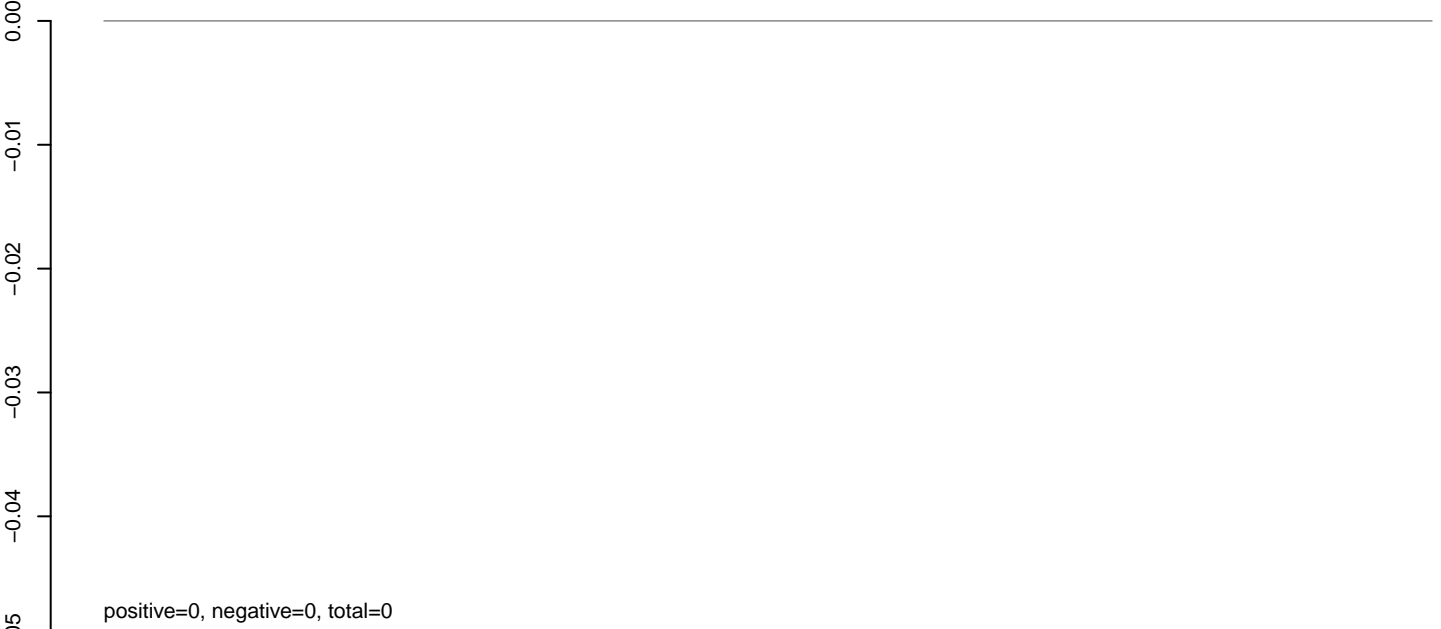


AeAeg_Aag2_NL.M3.rep

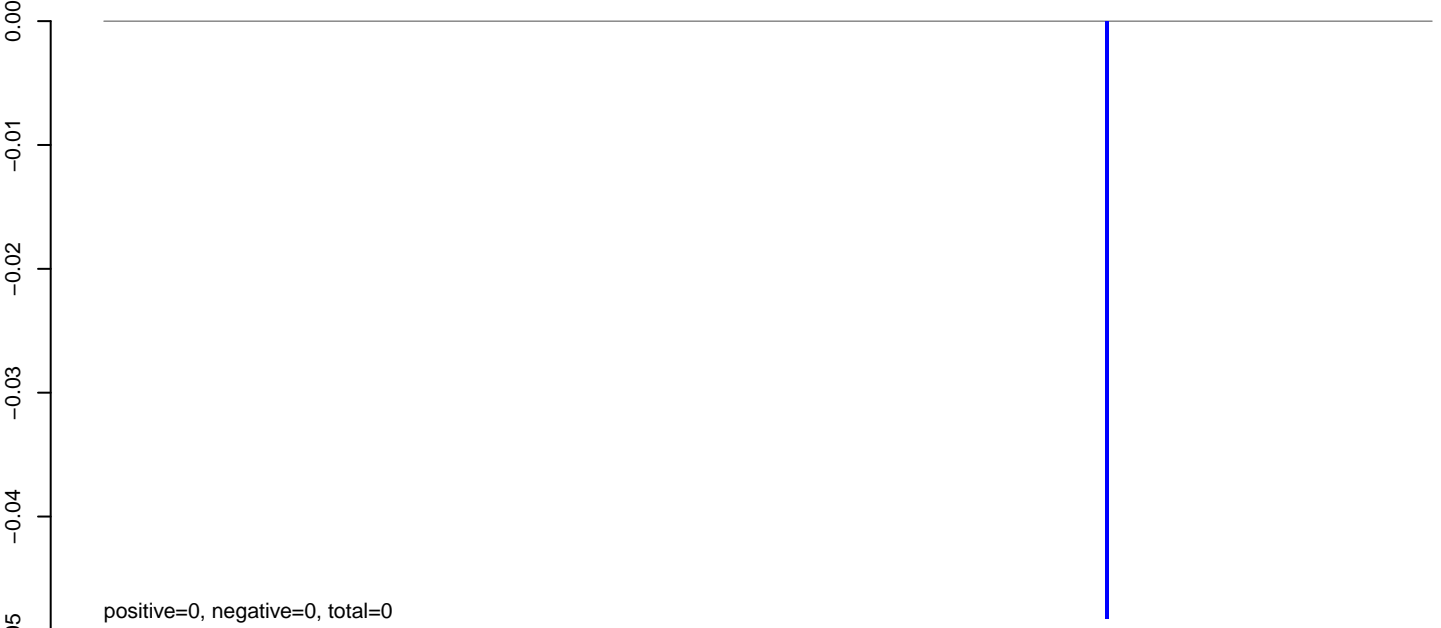


0 2000 4000 6000 8000 10000

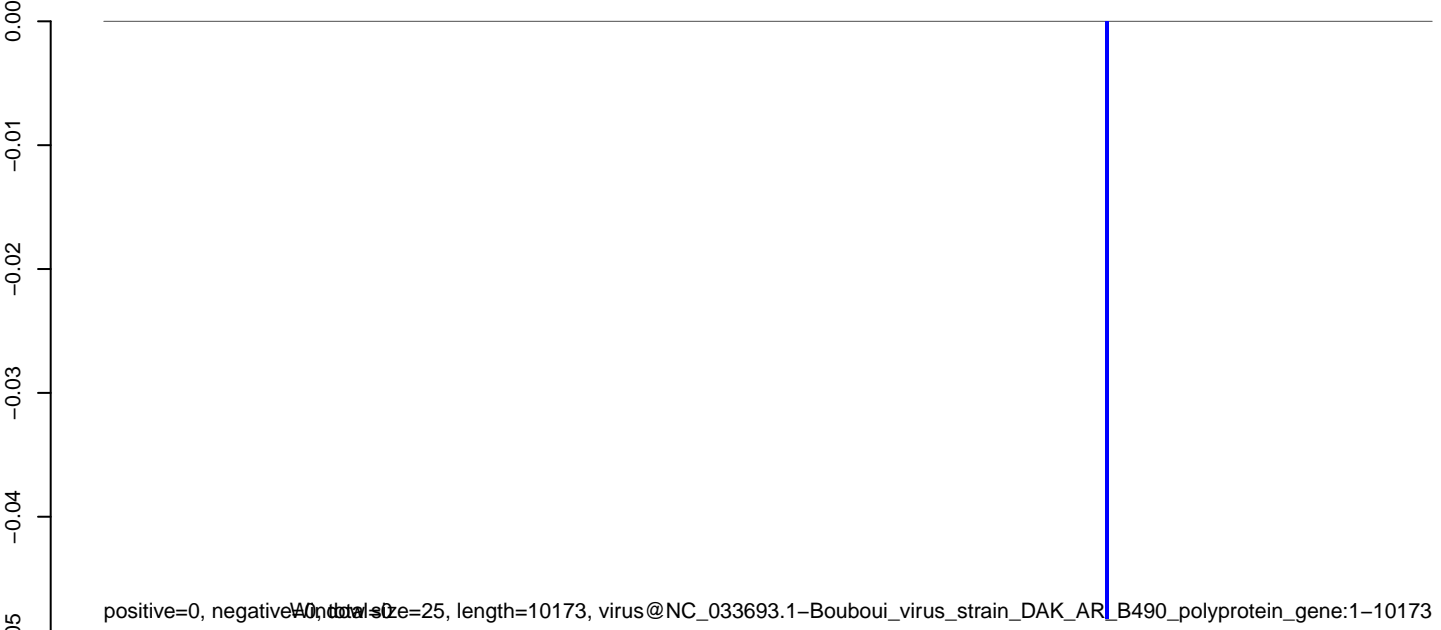
AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep



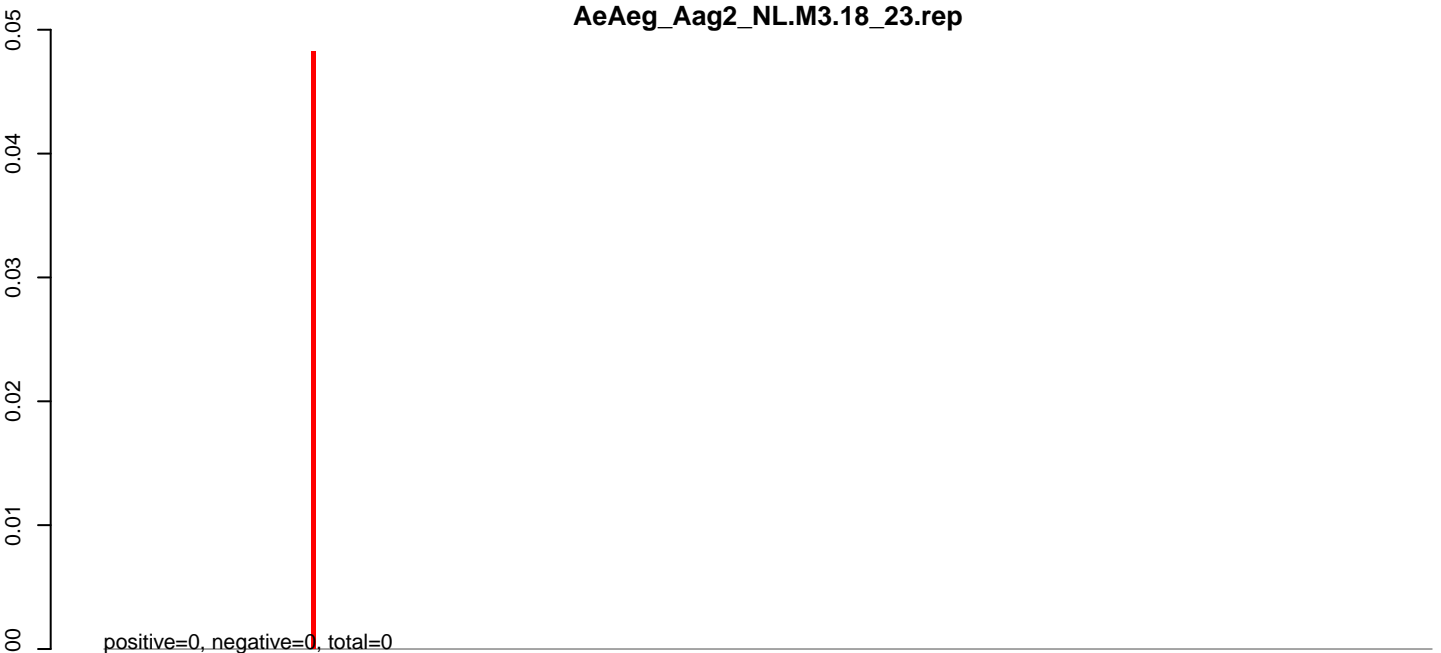
AeAeg_Aag2_NL.M3.rep



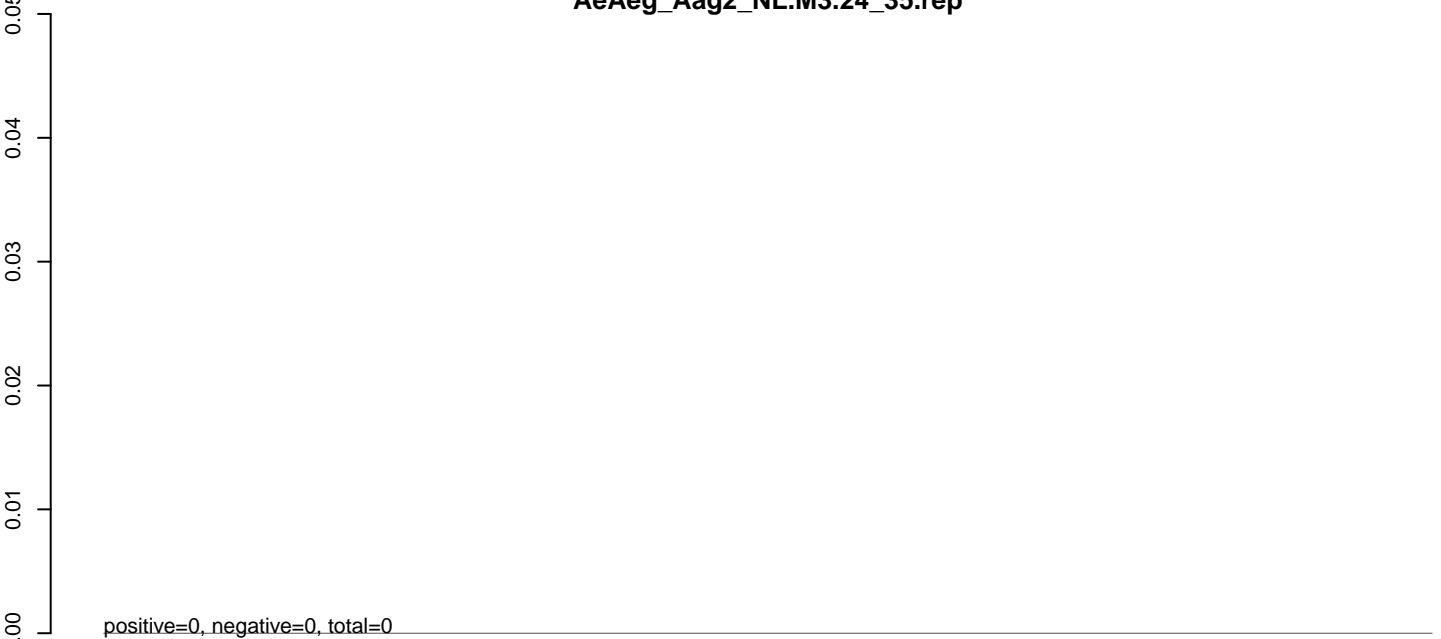
positive=0, negative=0, total=0, n=25, length=10173, virus@NC_033693.1-Bouboui_virus_strain_DAK_AR_B490_polyprotein_gene:1-10173

0 2000 4000 6000 8000 10000

AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep



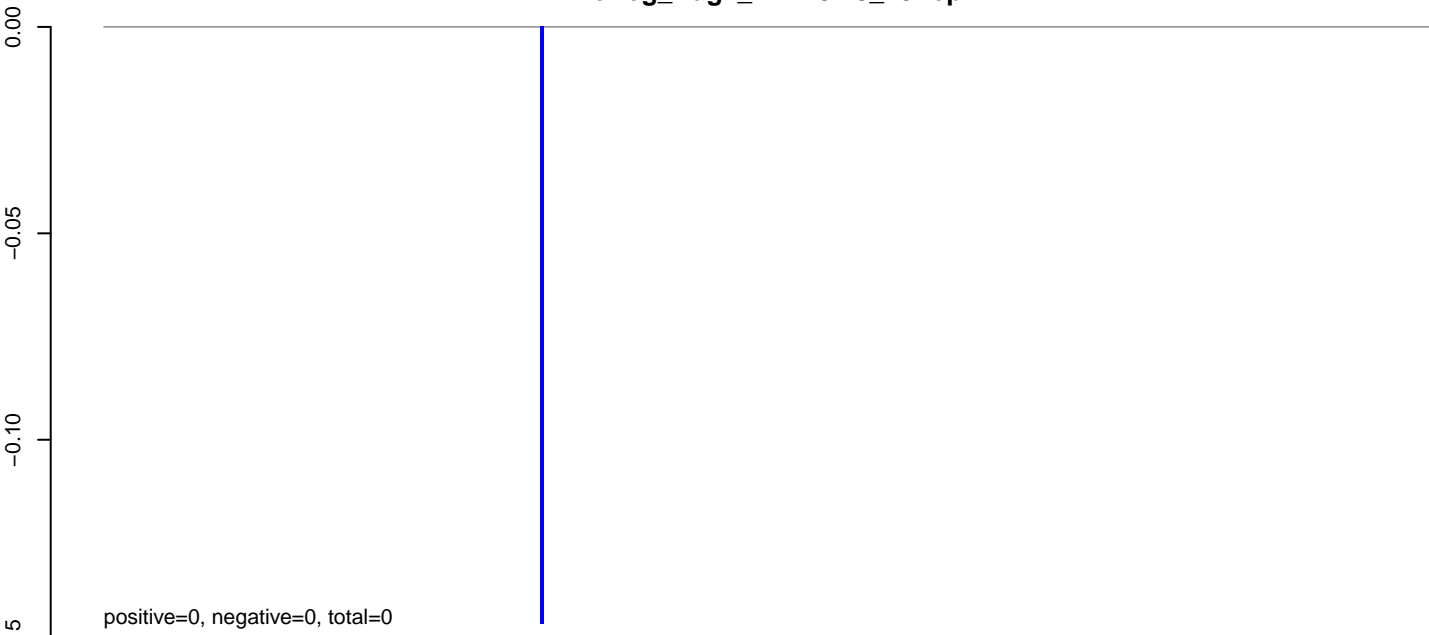
AeAeg_Aag2_NL.M3.rep



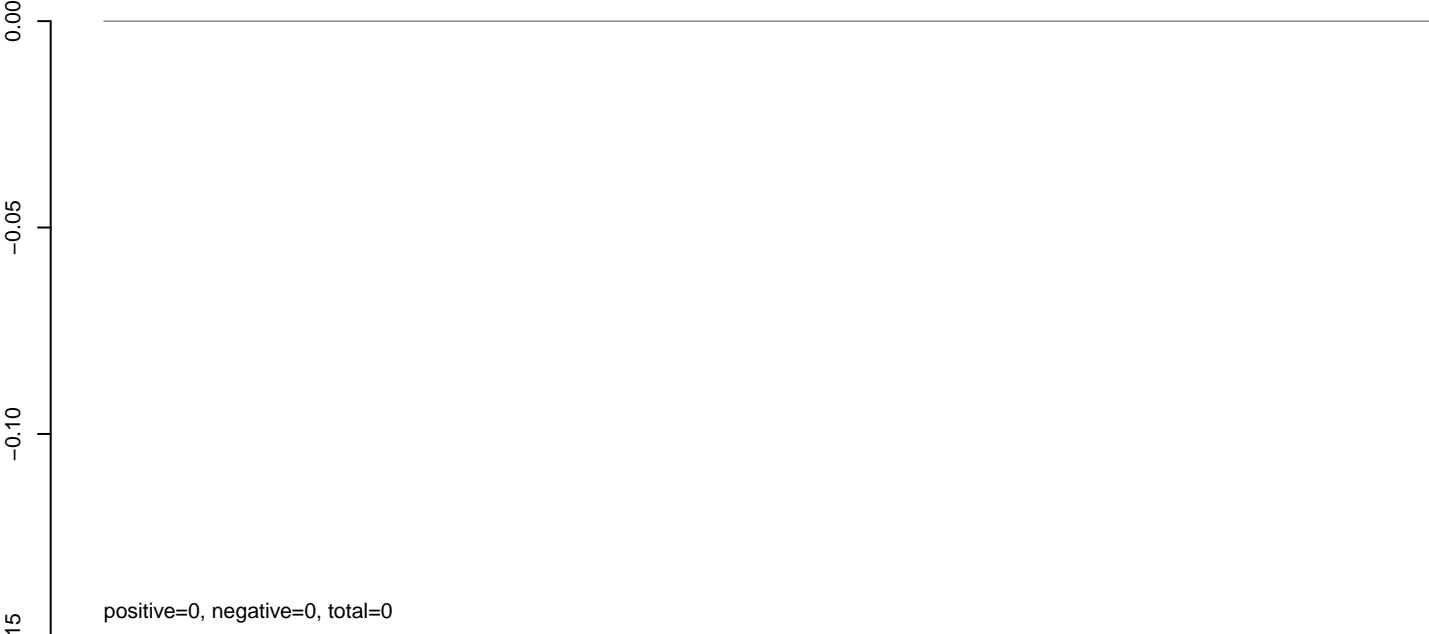
v size=251 length=6675 virus @ NC_033327.1-Hubei_chuvirus-like_virus_1_strain_QTM26249_RNA-dependent_RNA_polymerase_gene:1-6873

0 1000 2000 3000 4000 5000 6000 7000

AeAeg_Aag2_NL.M3.18_23.rep



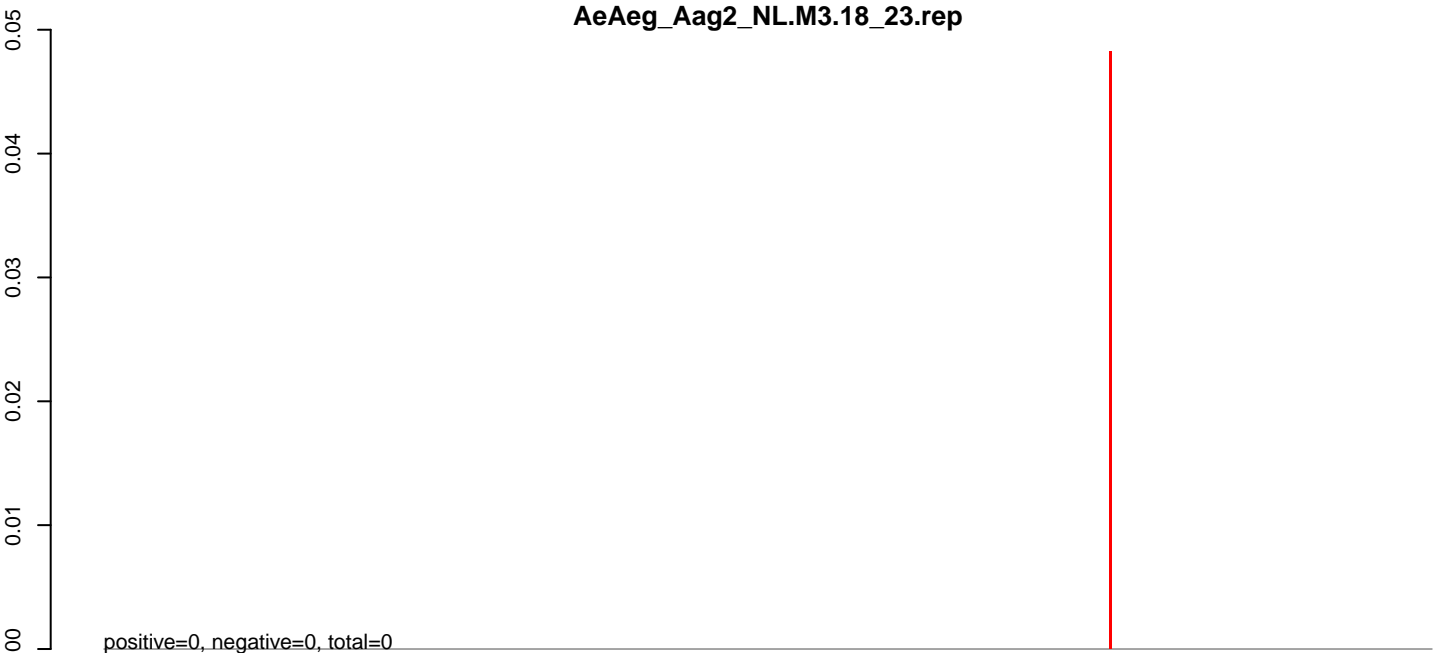
AeAeg_Aag2_NL.M3.24_35.rep



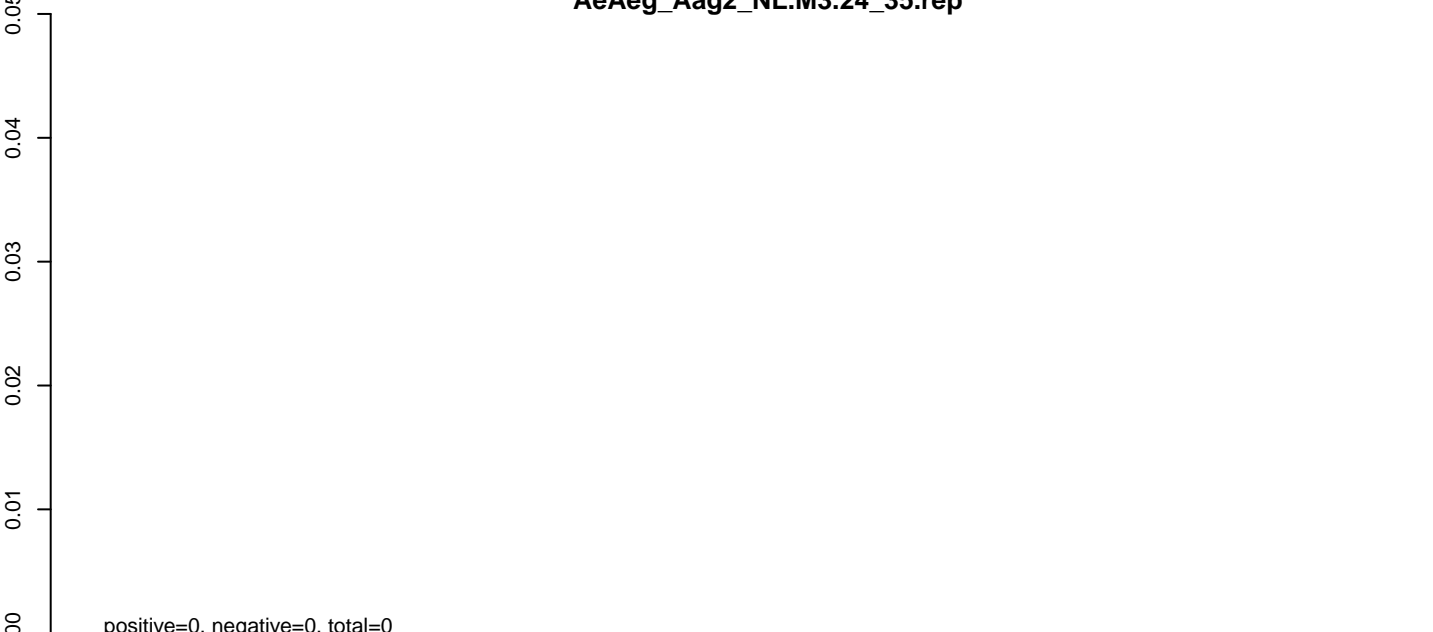
AeAeg_Aag2_NL.M3.rep



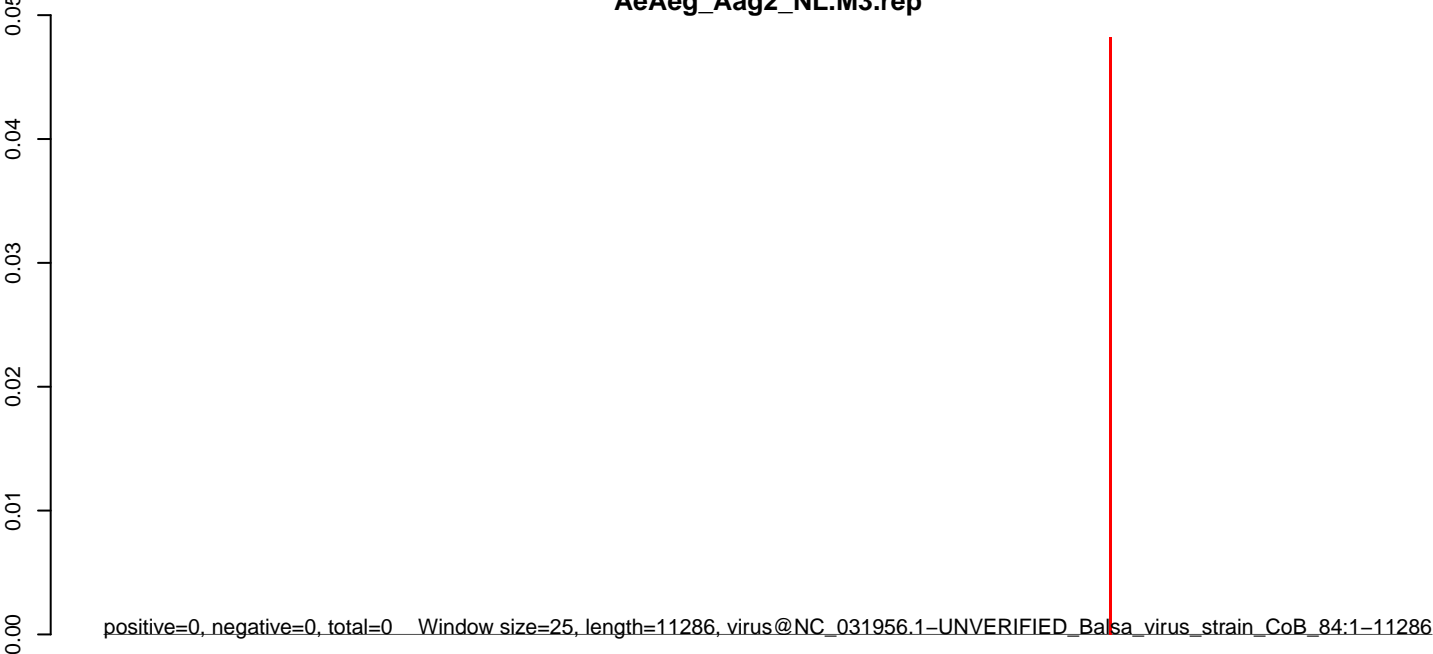
AeAeg_Aag2_NL.M3.18_23.rep



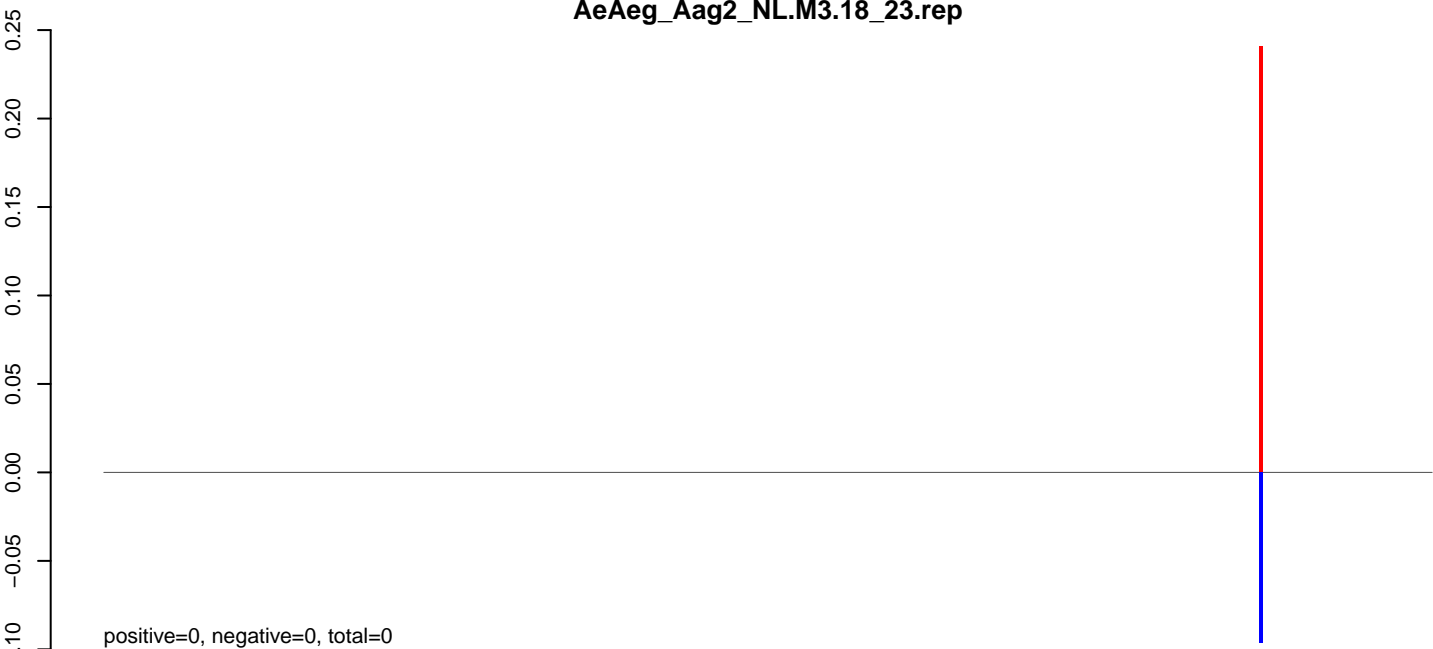
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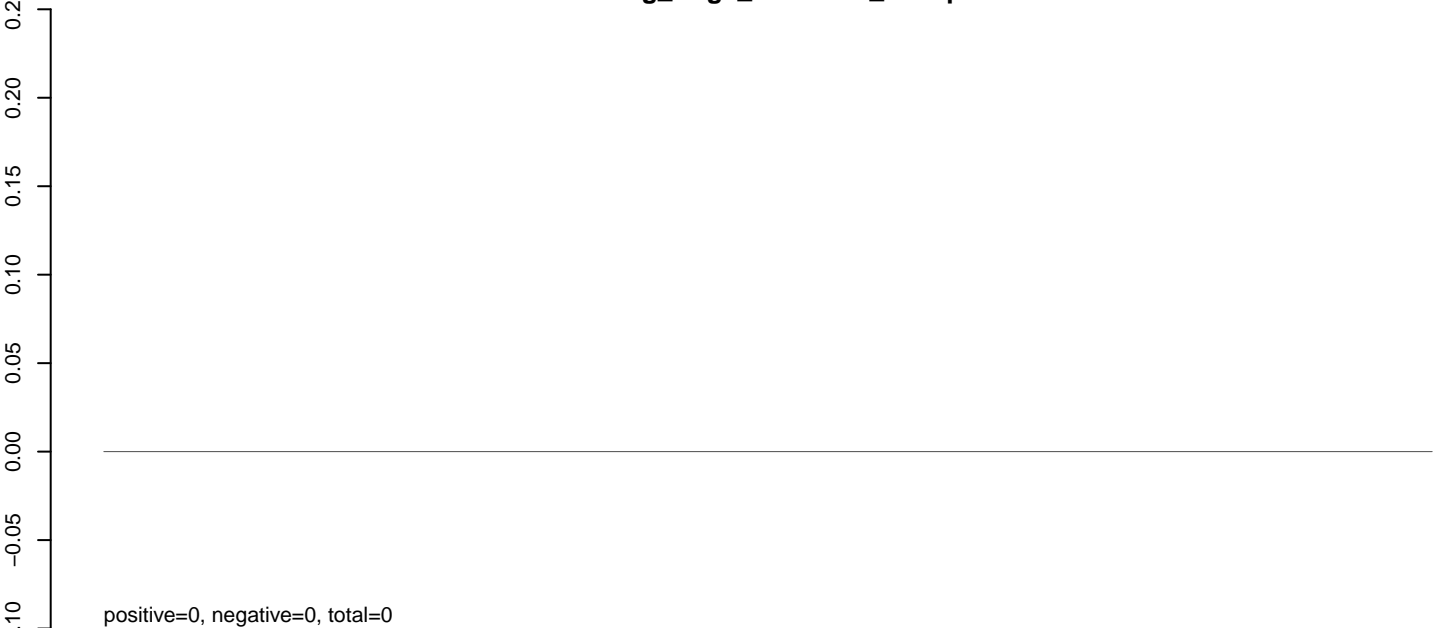
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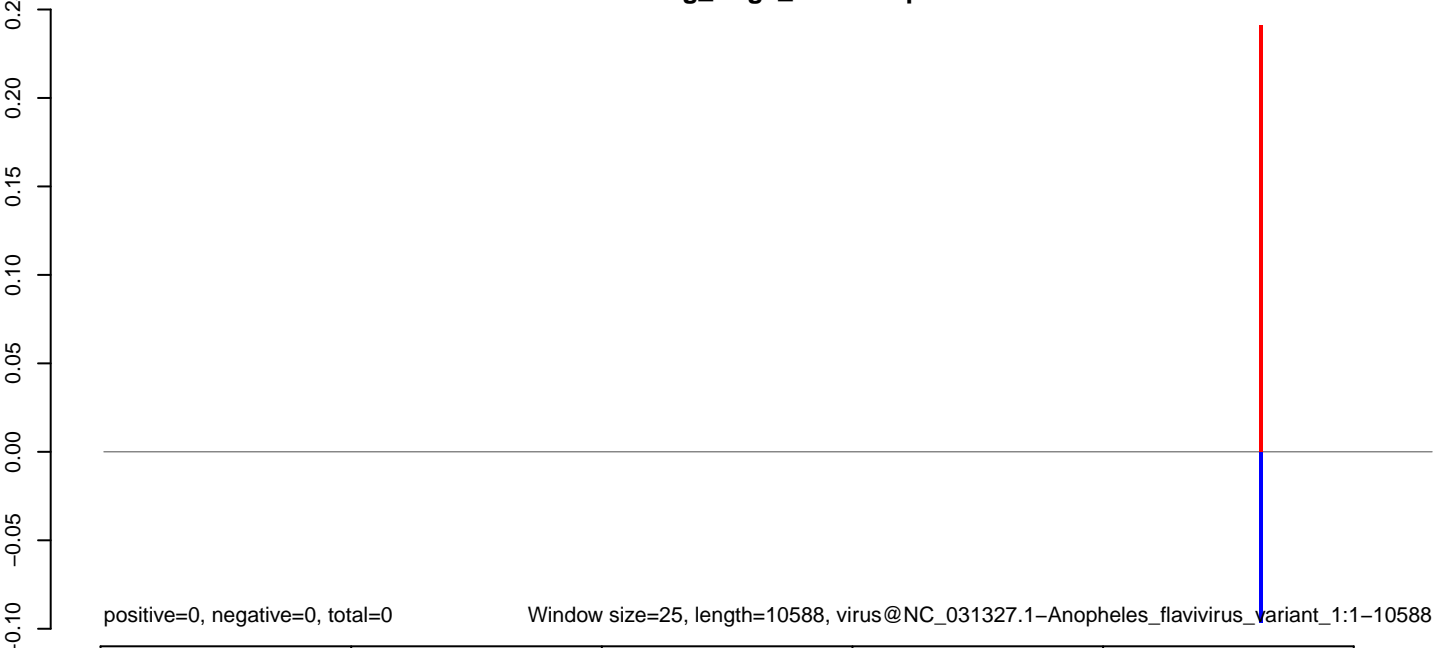
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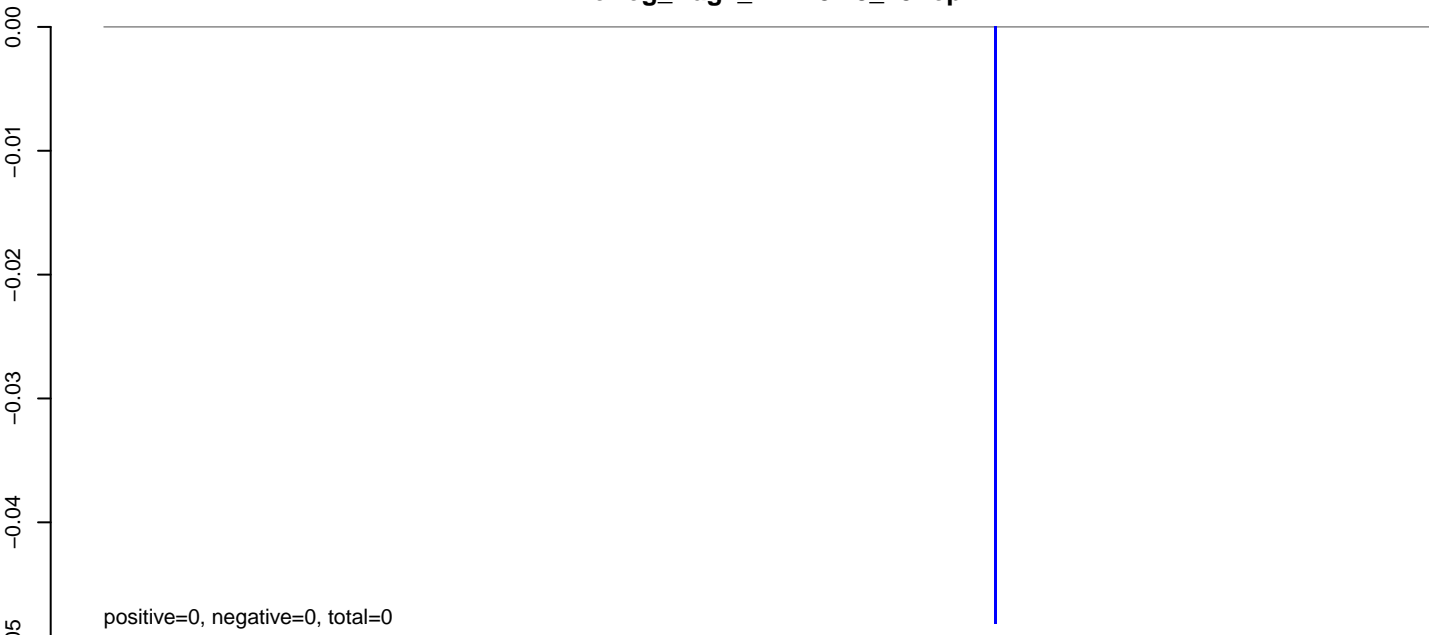
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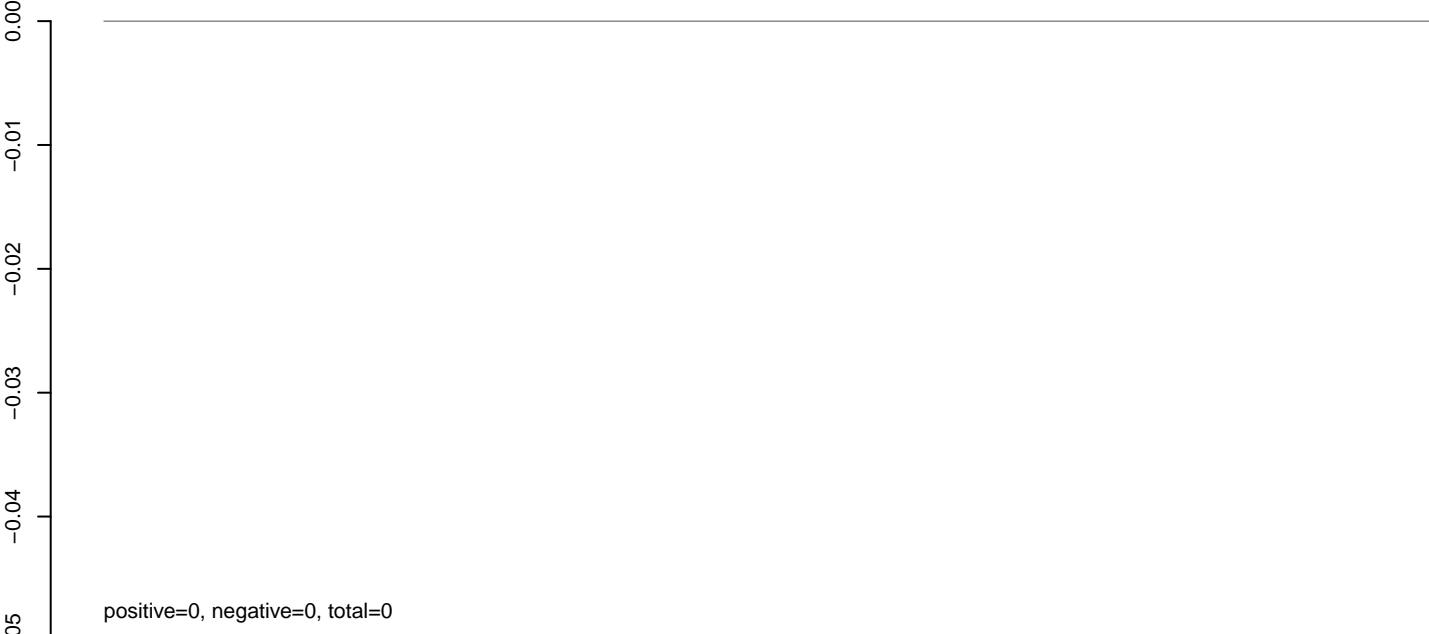
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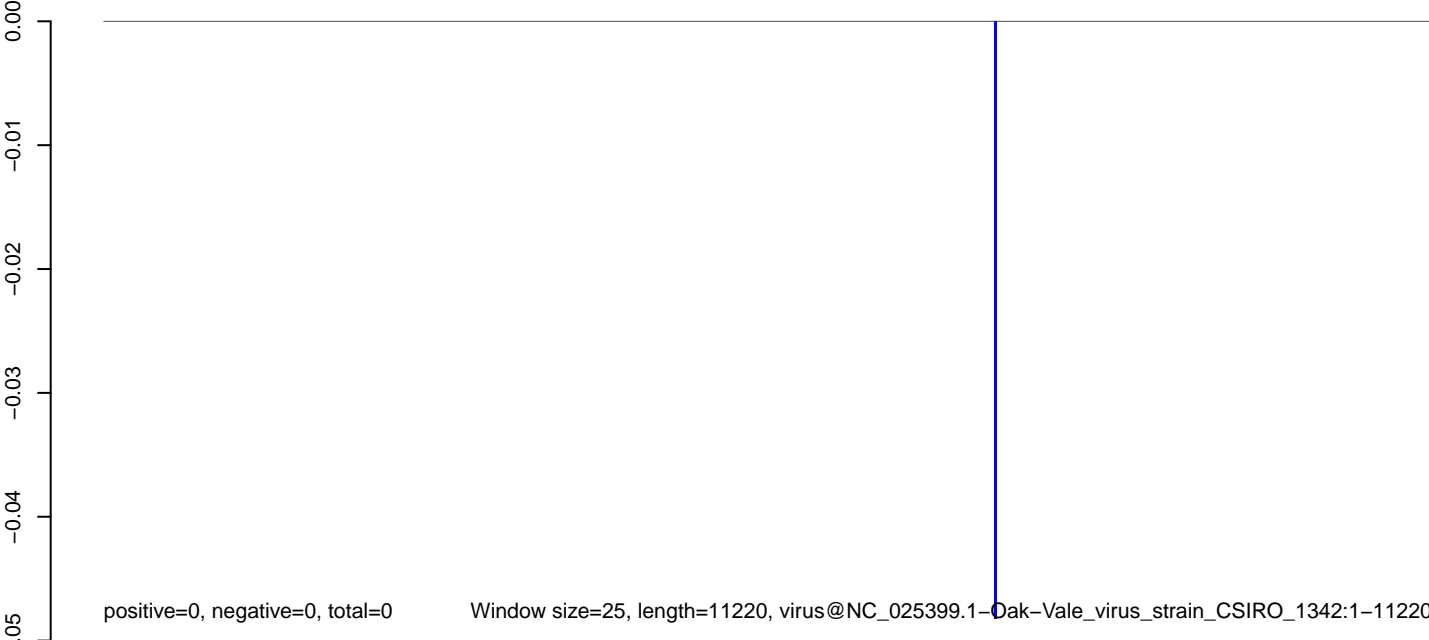
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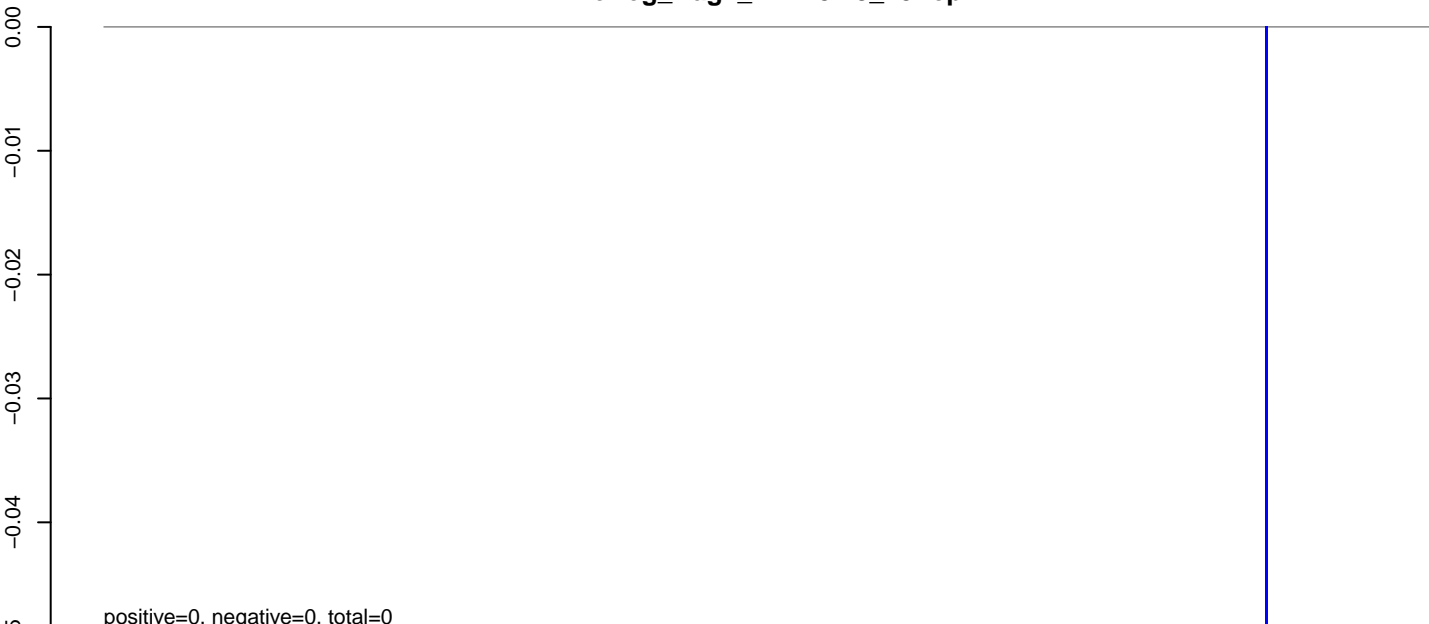
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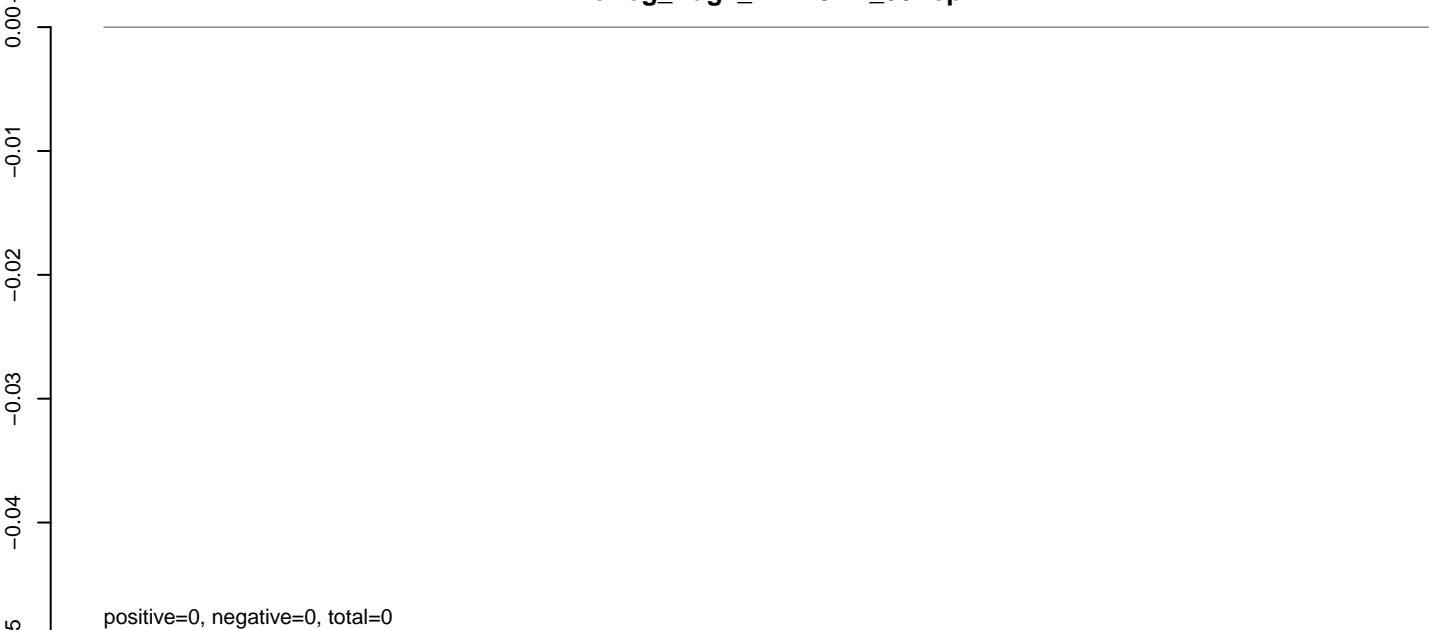
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AeAeg_Aag2_NL.M3.18_23.rep



AeAeg_Aag2_NL.M3.24_35.rep



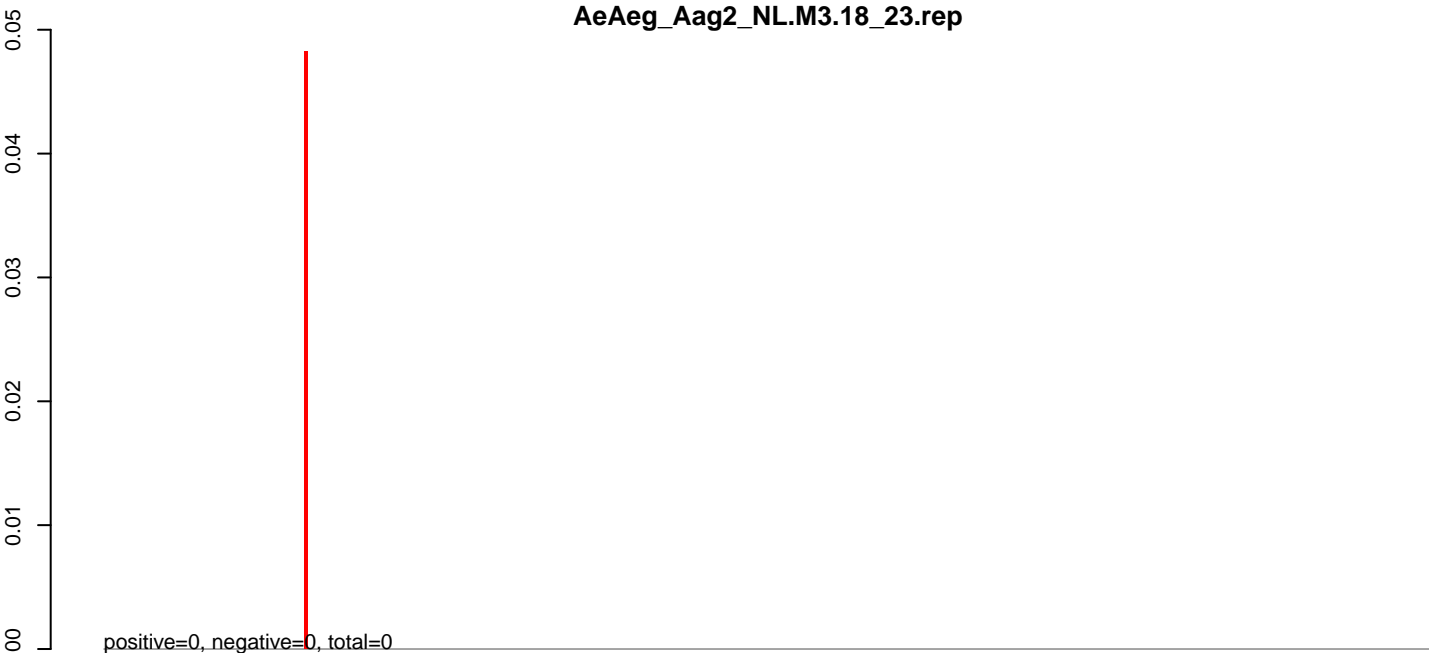
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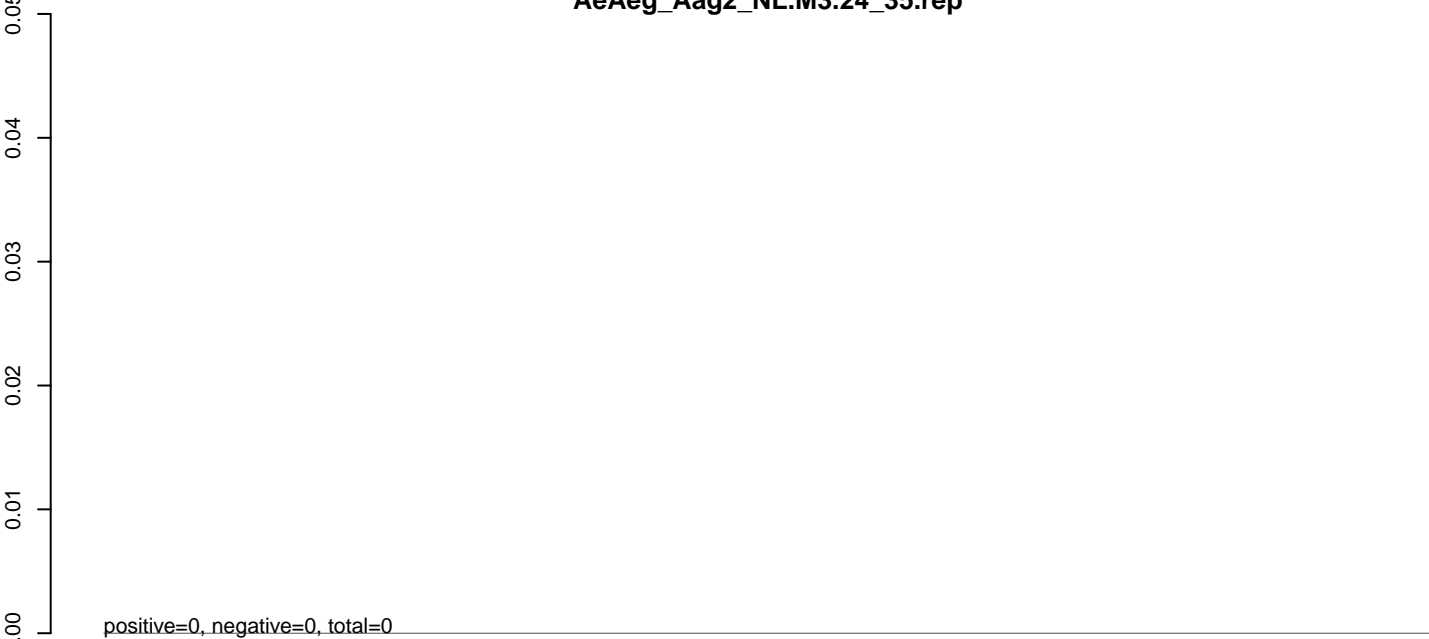
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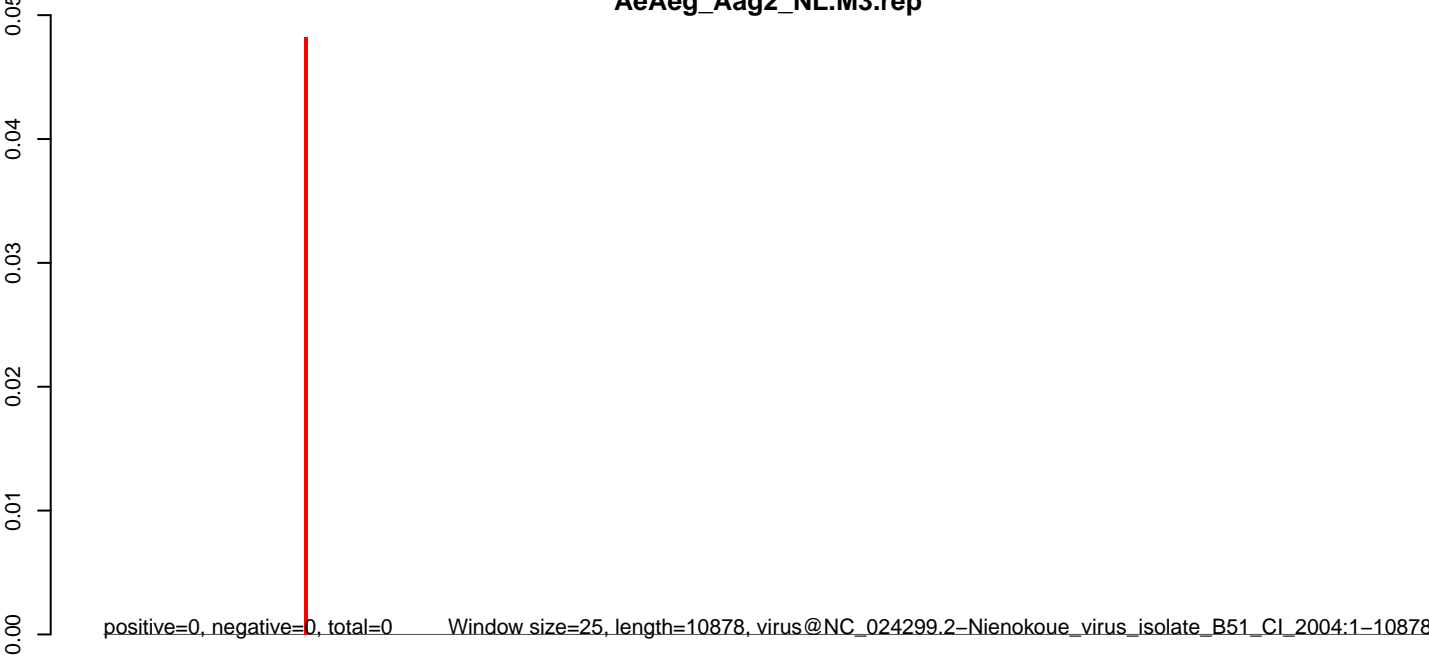
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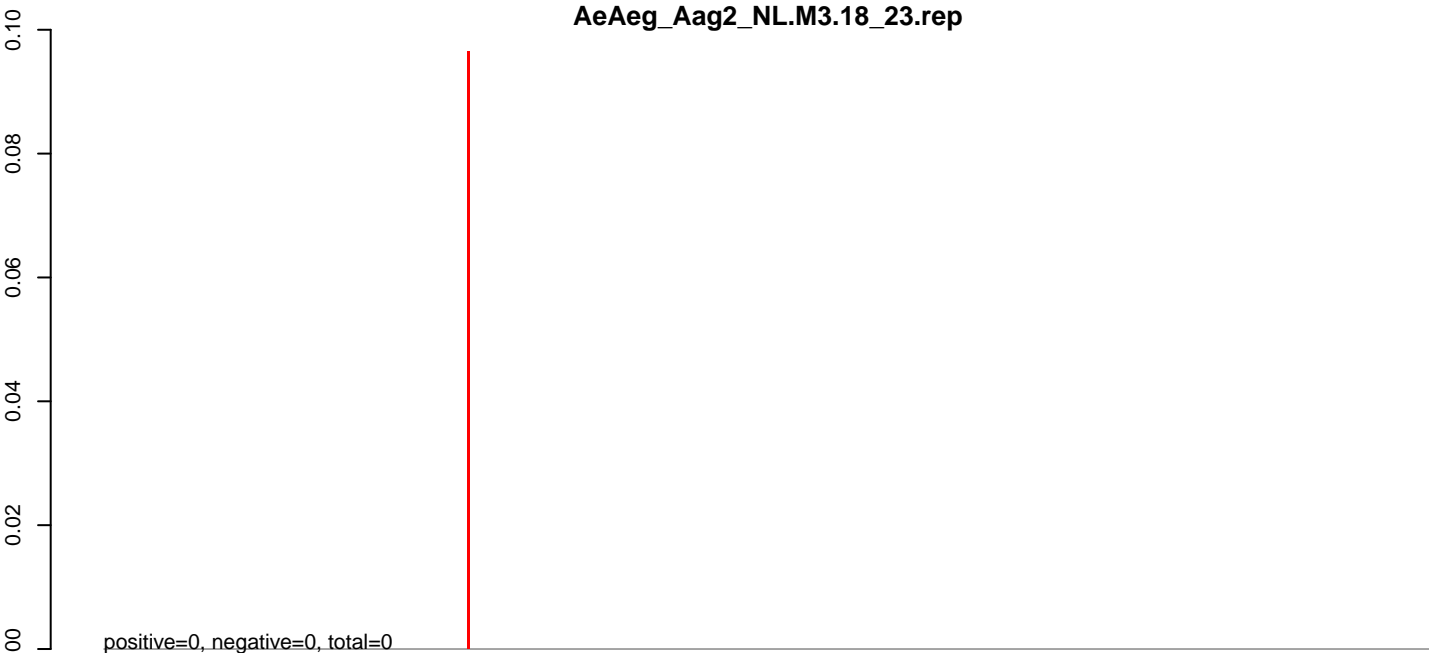
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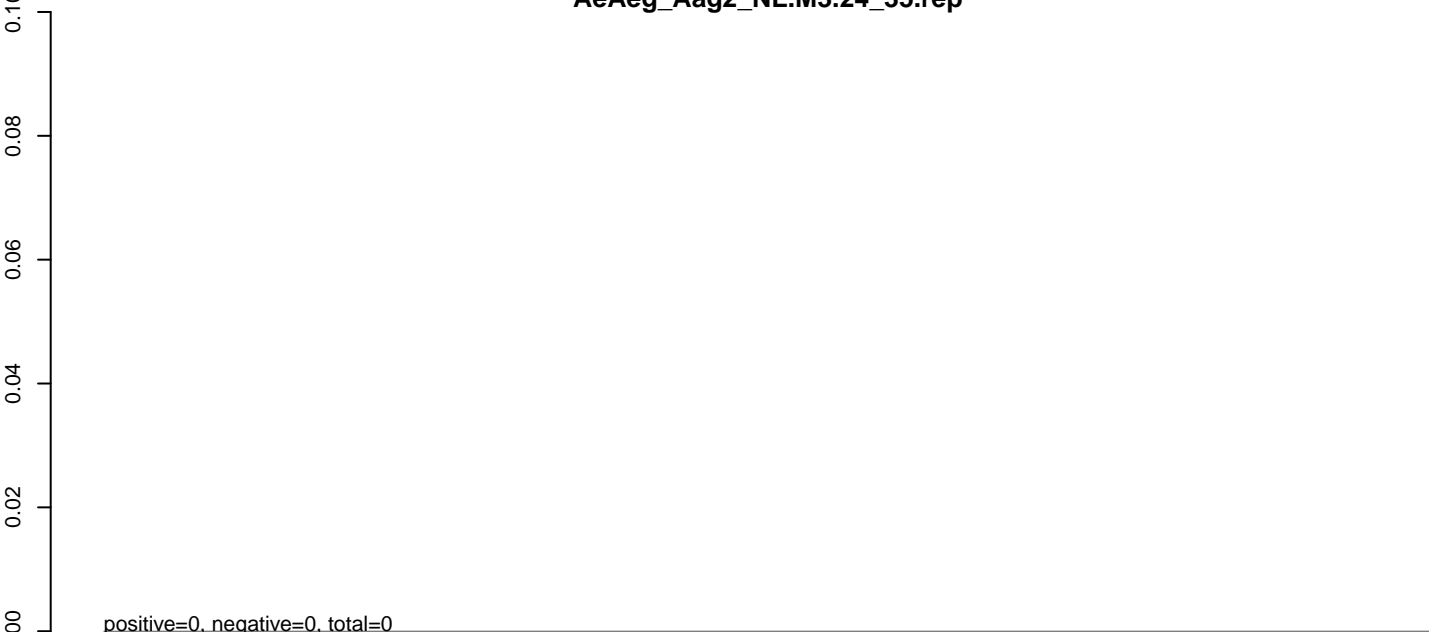
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AeAeg_Aag2_NL.M3.18_23.rep



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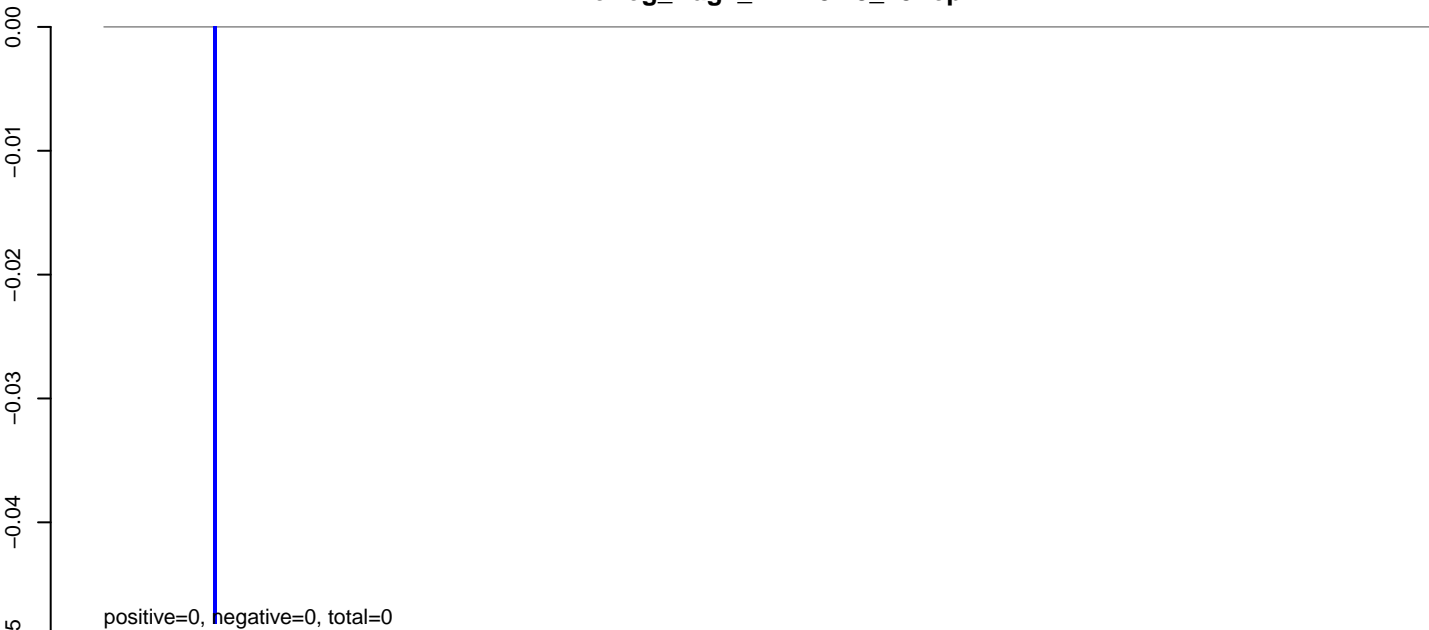
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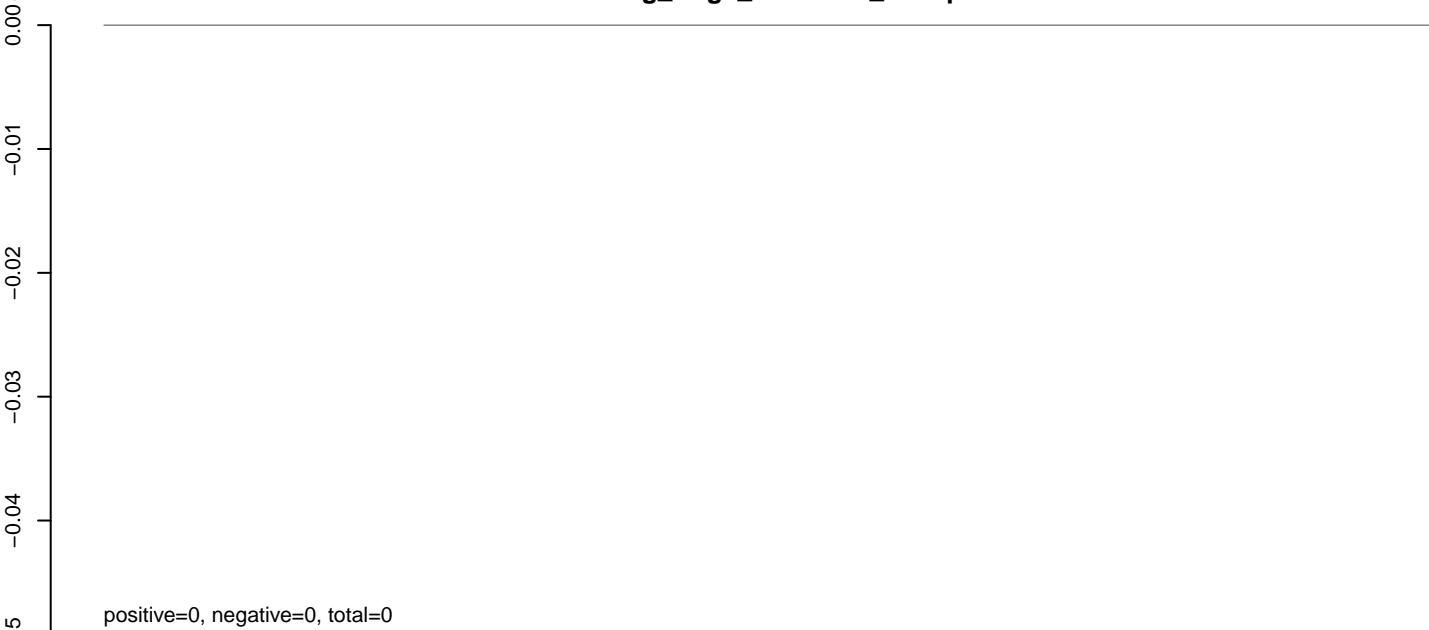
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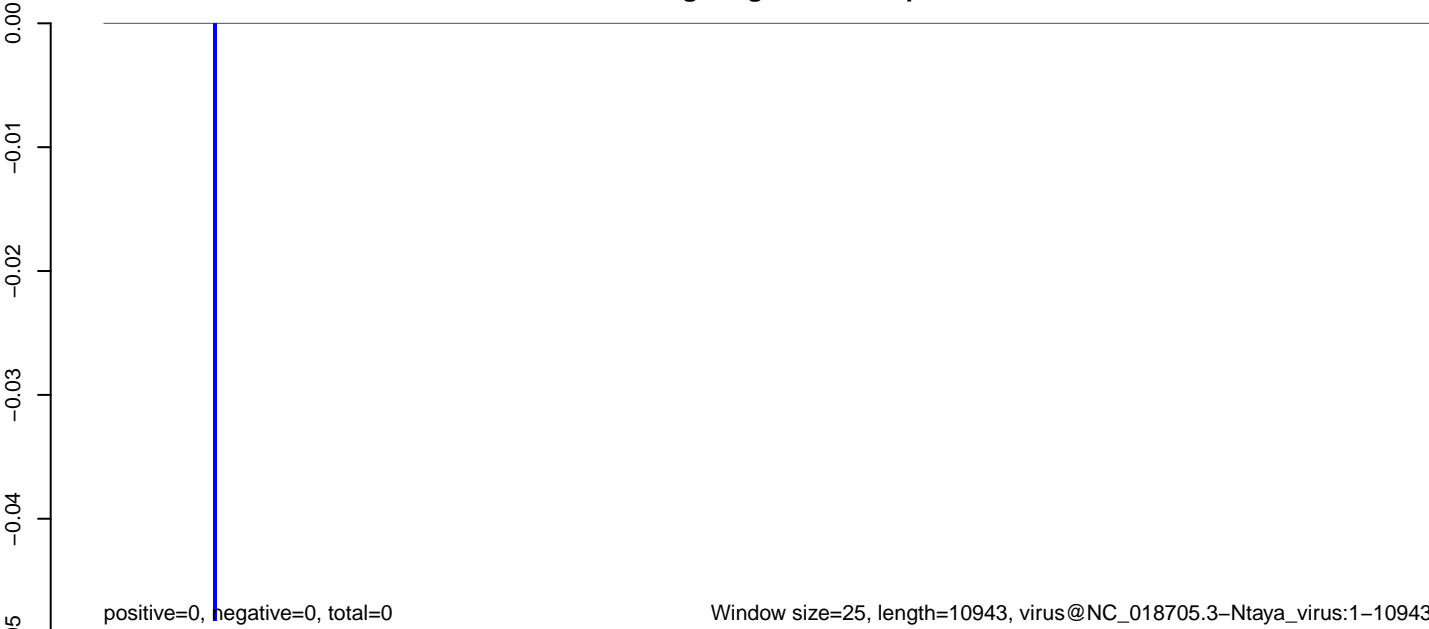
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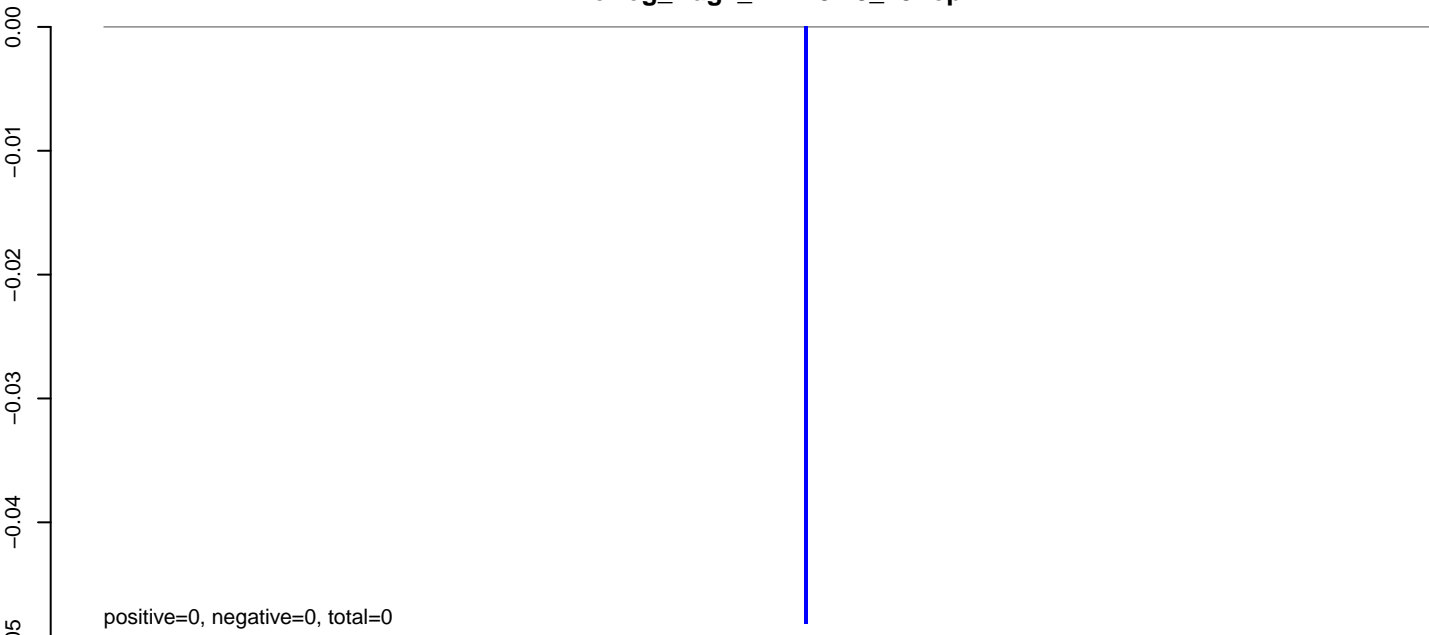
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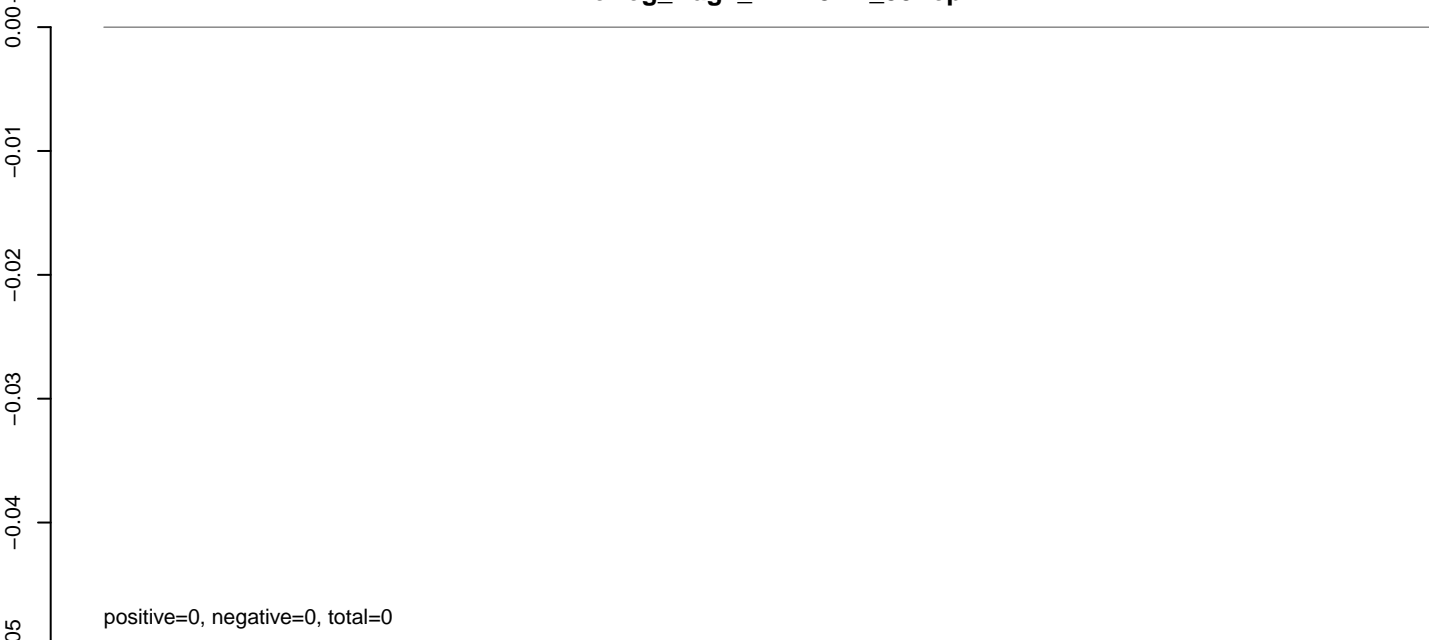
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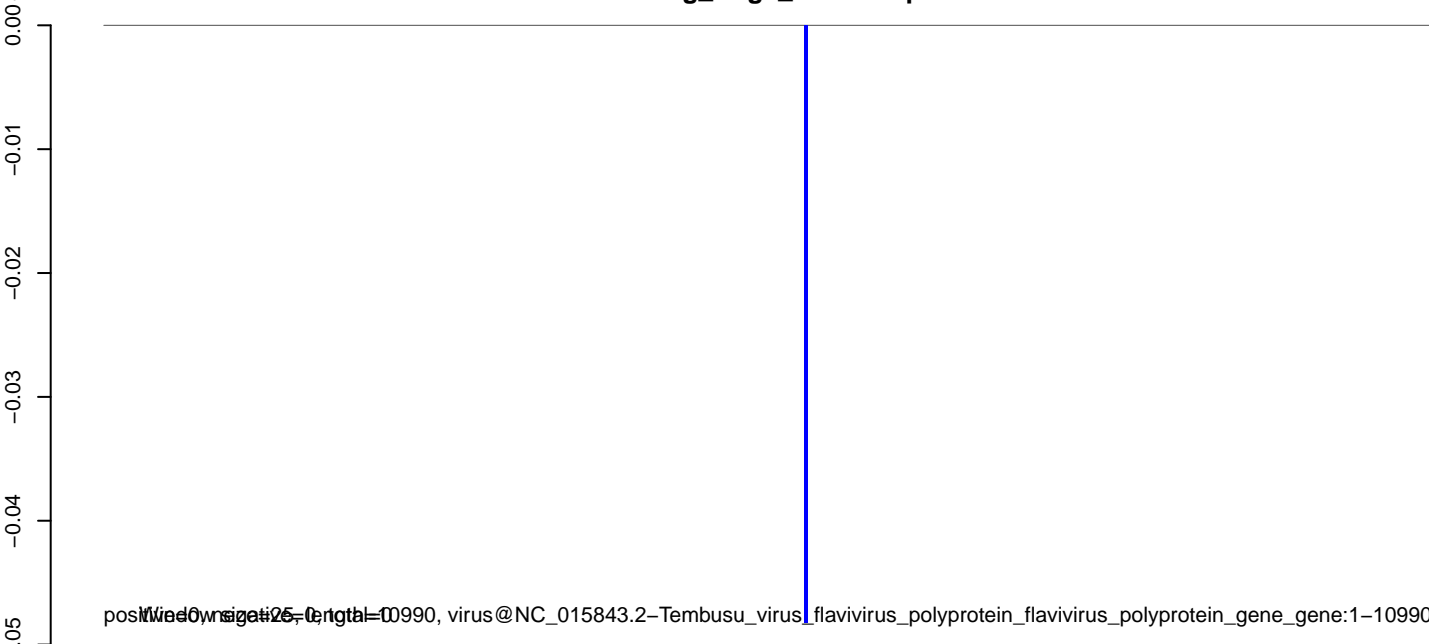
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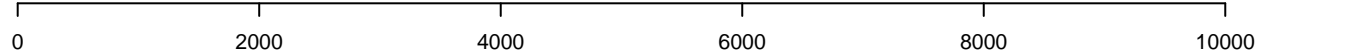
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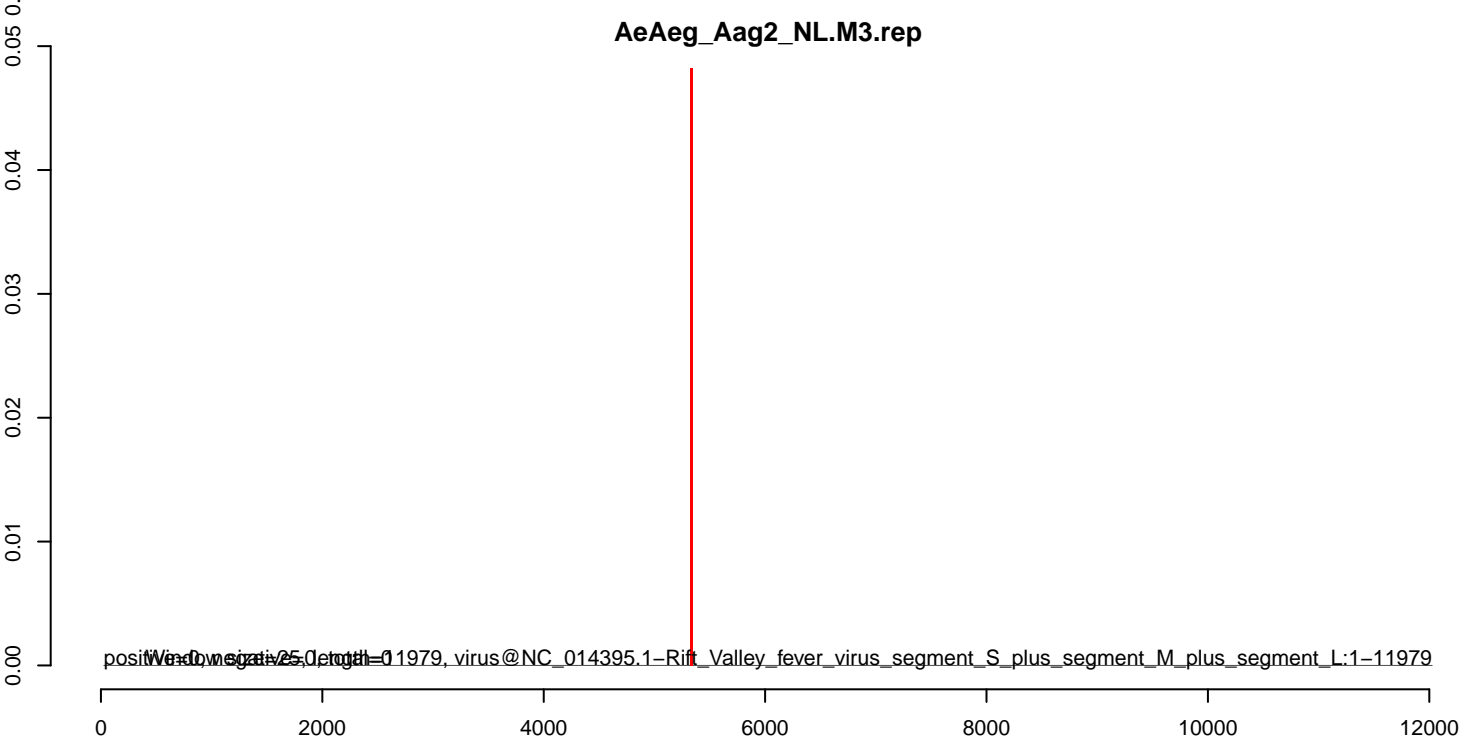
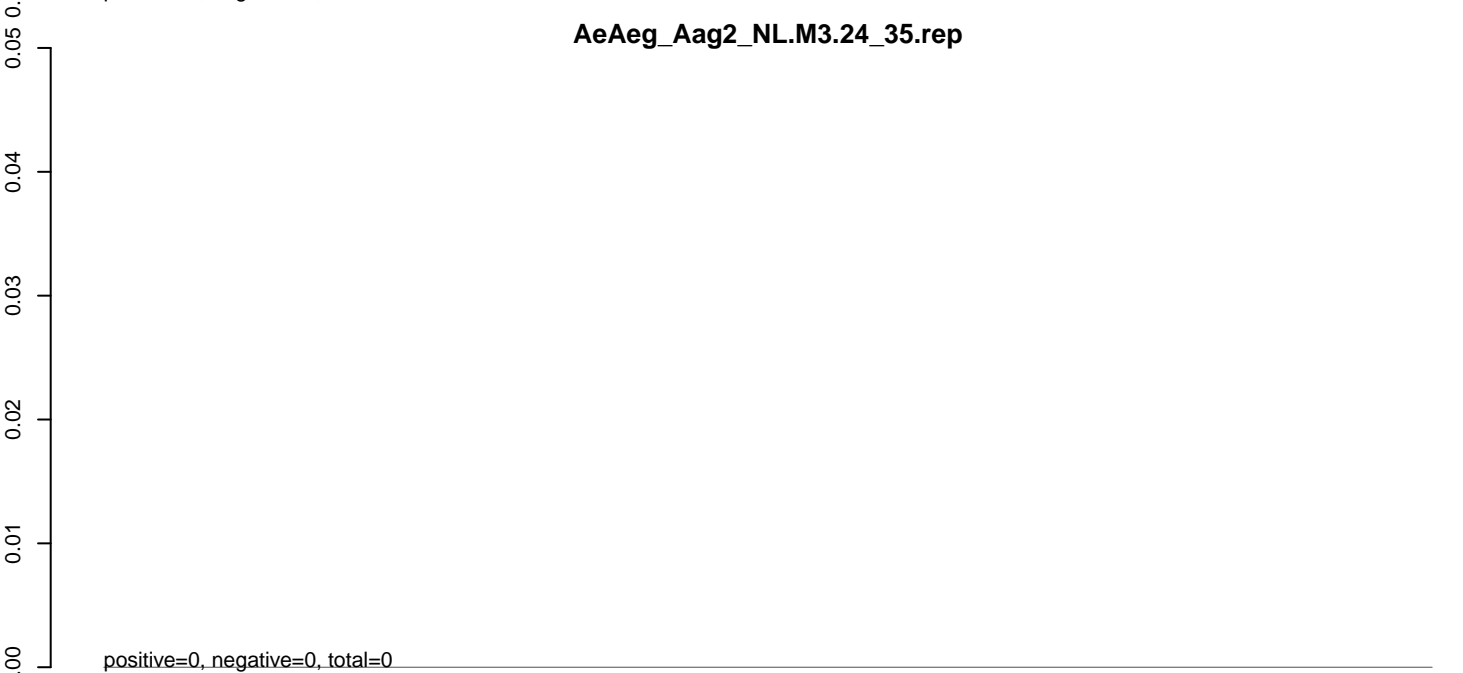
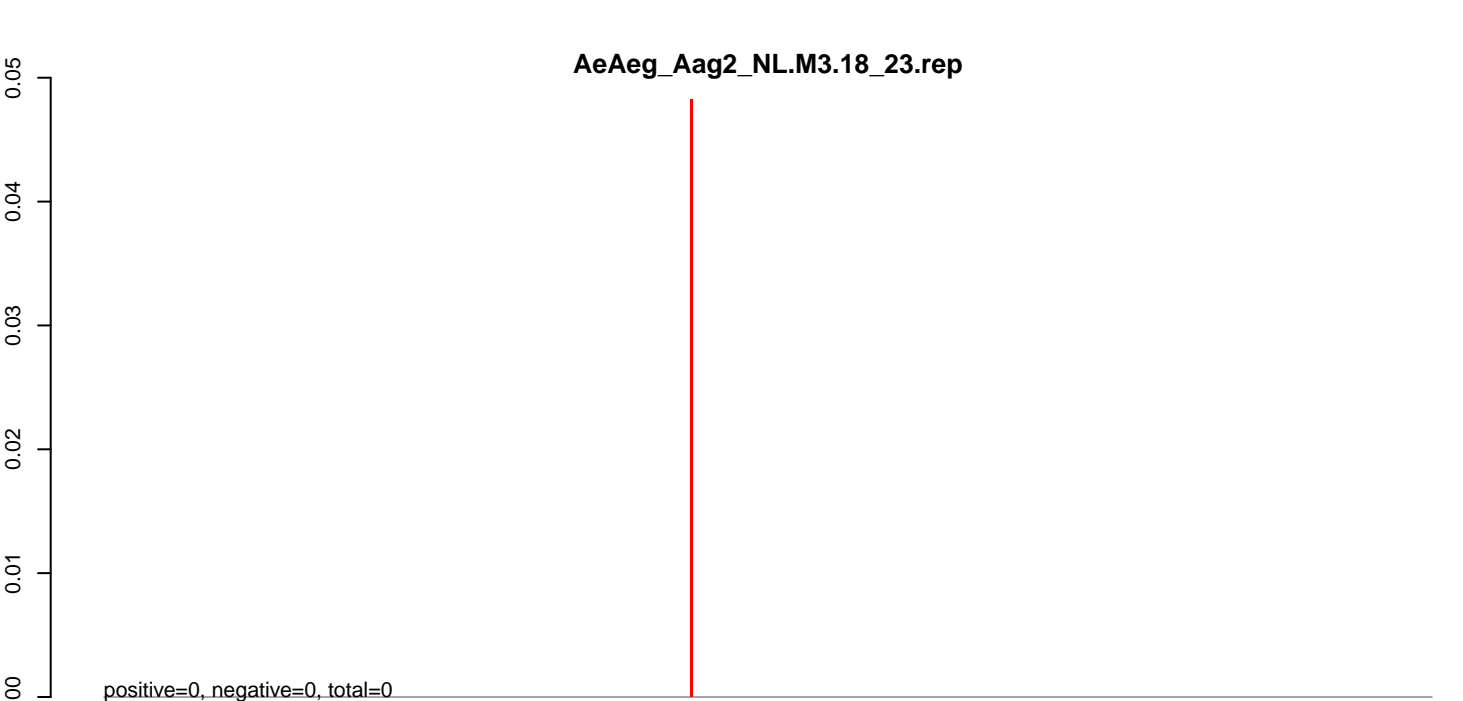


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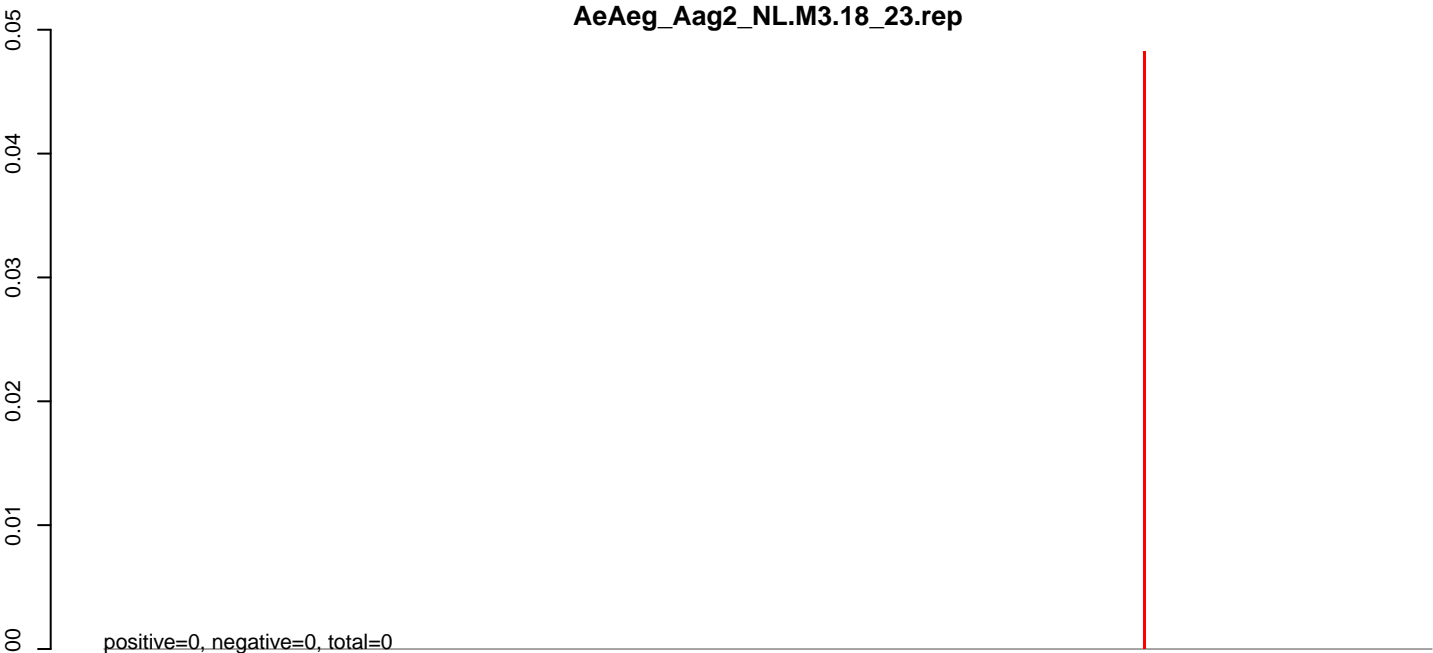


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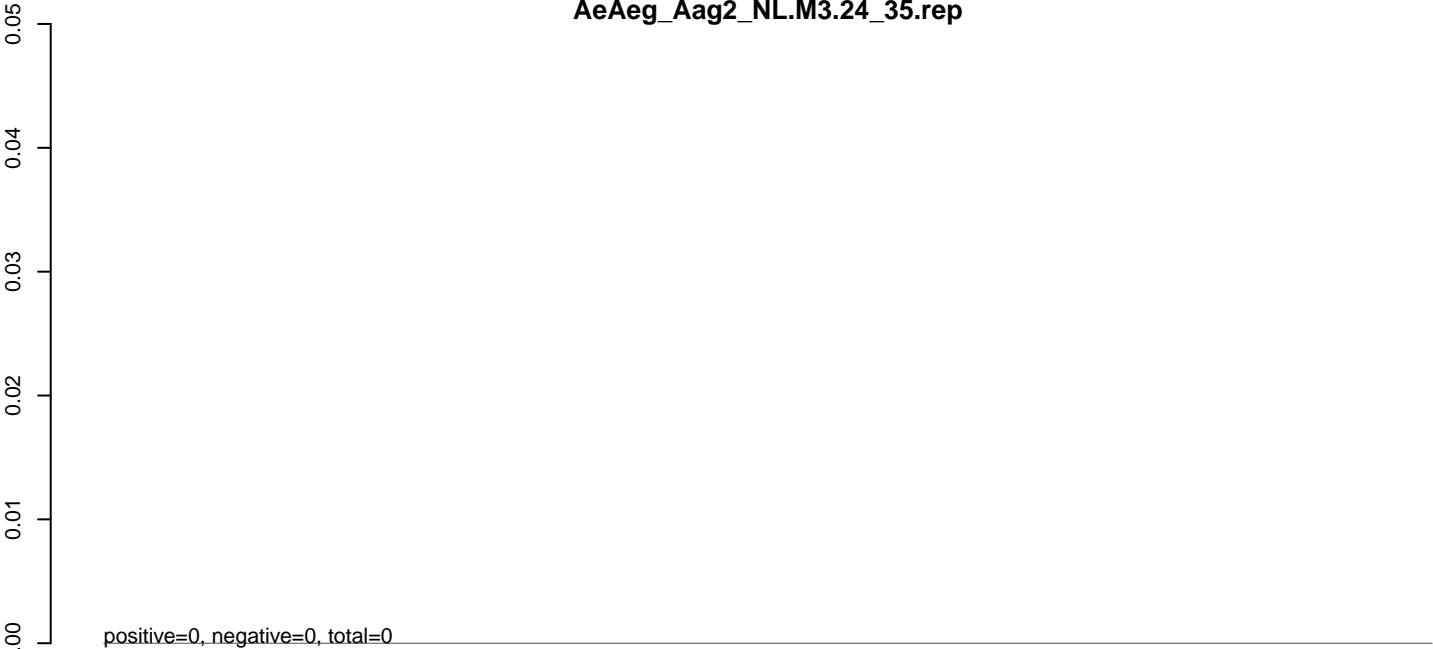




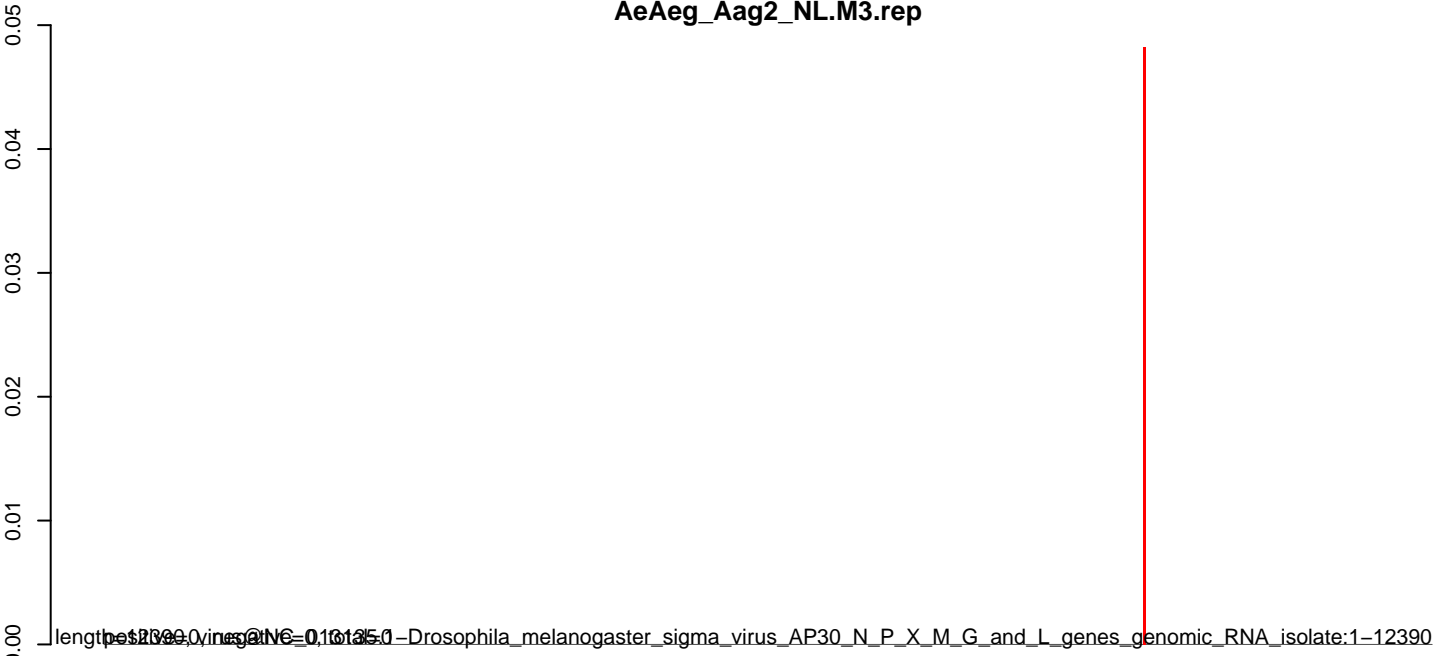
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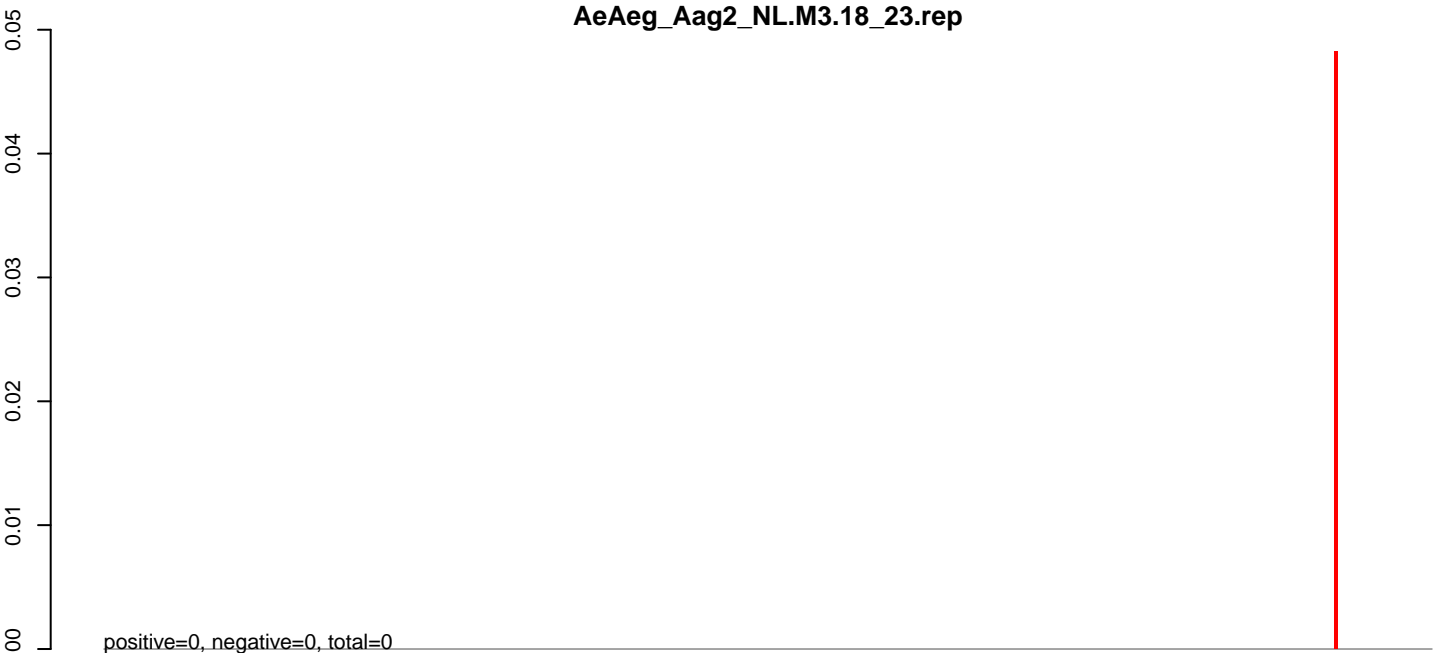
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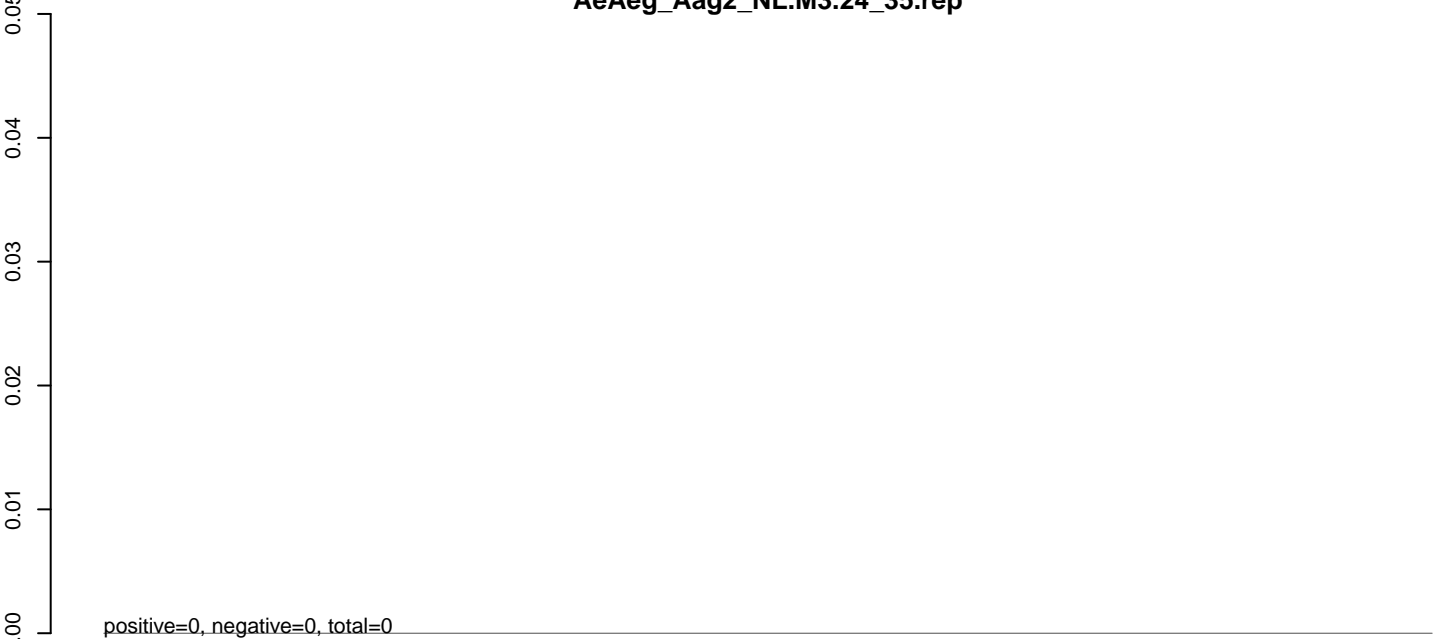
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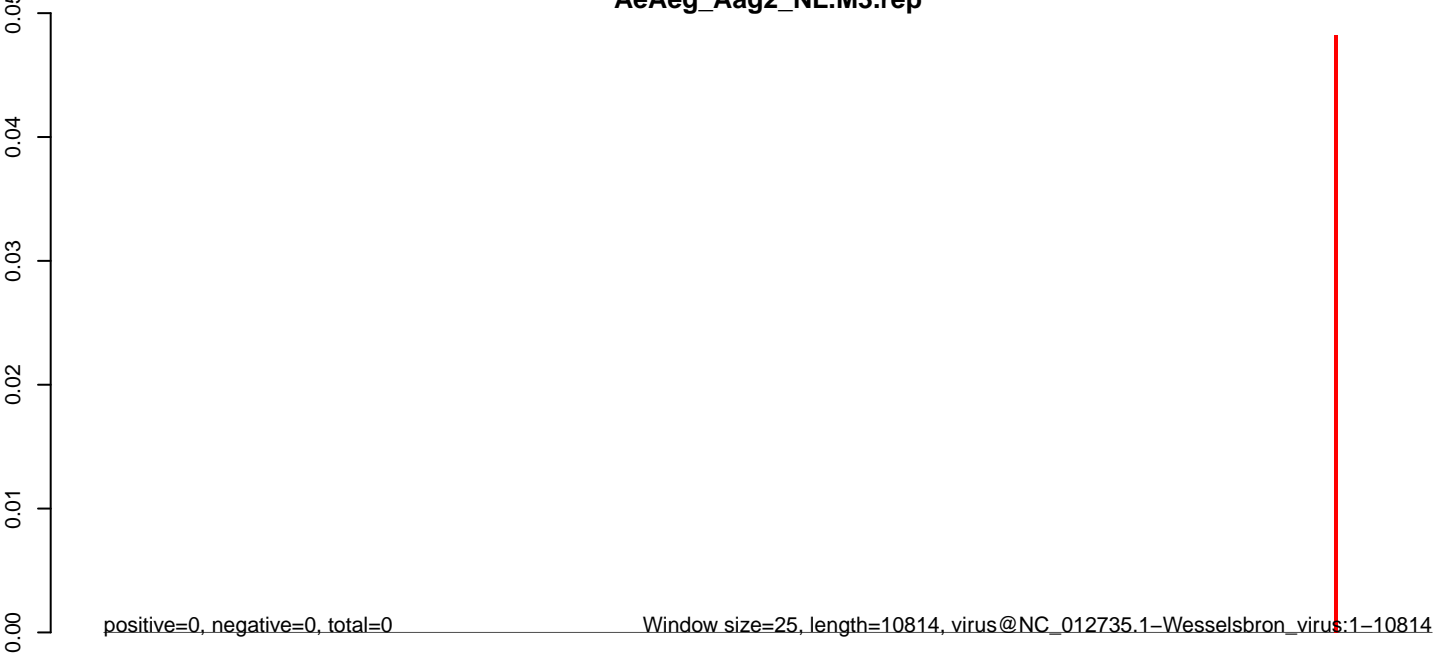
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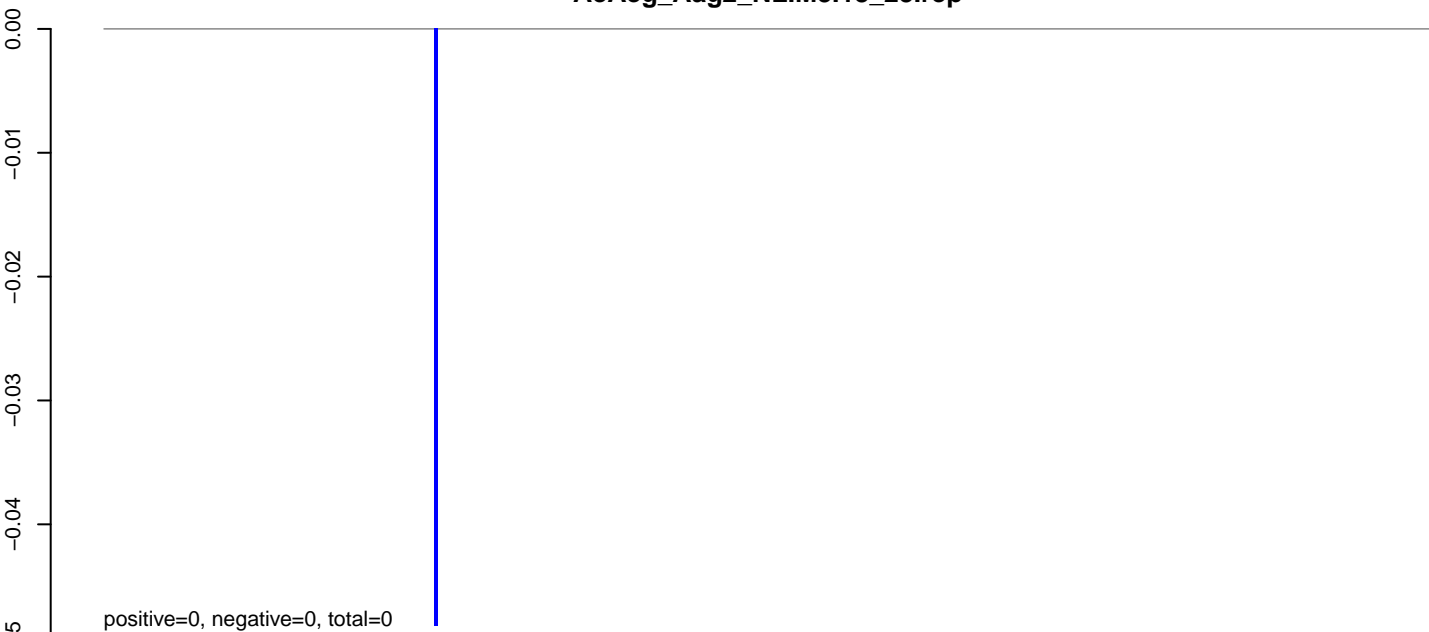


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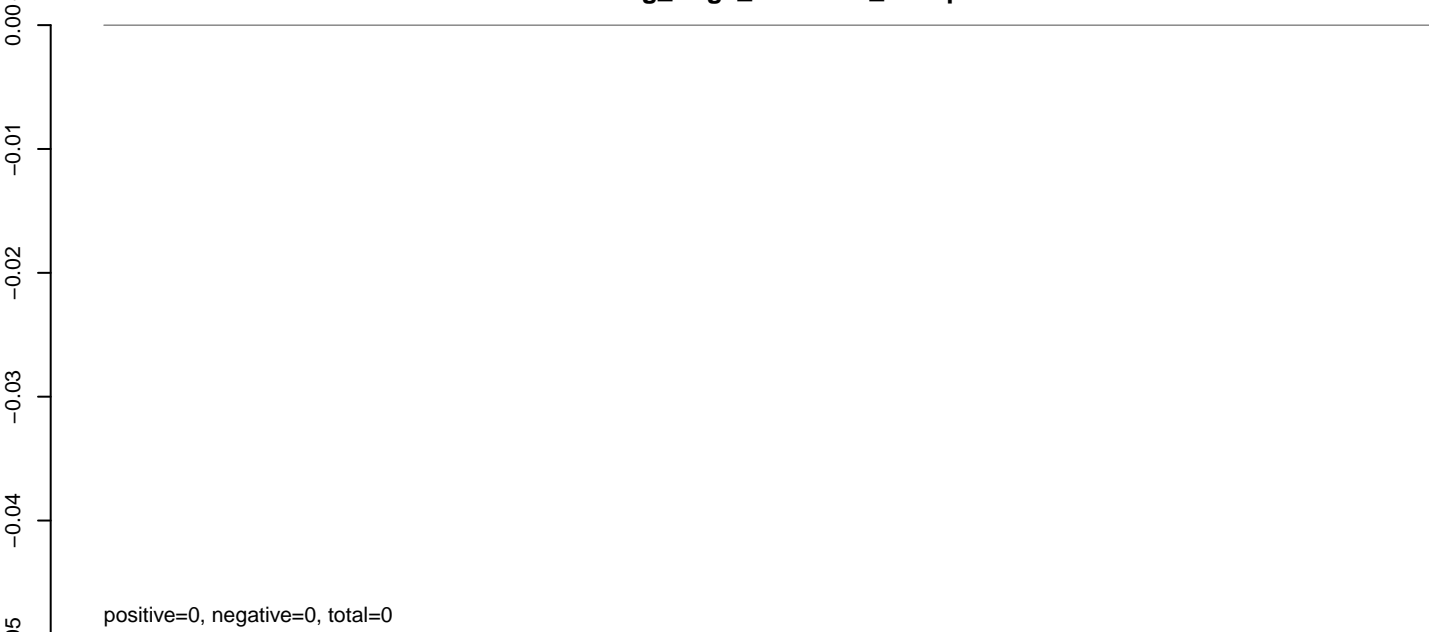


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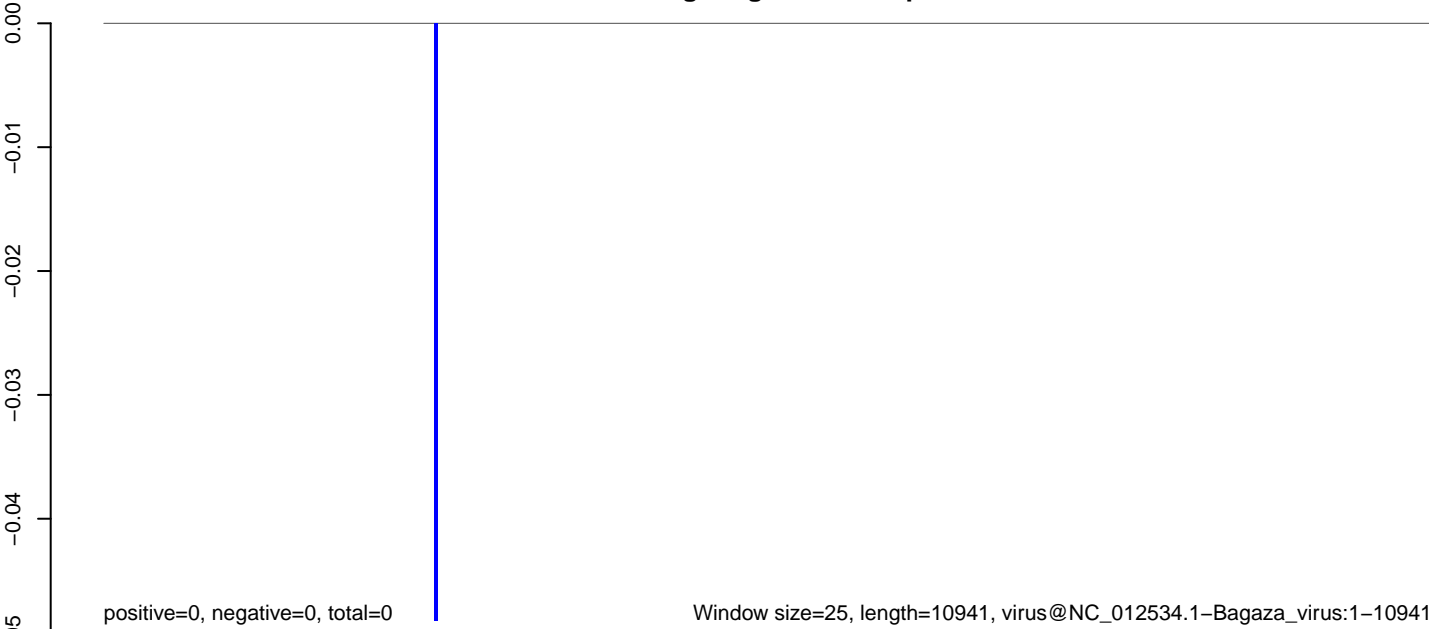
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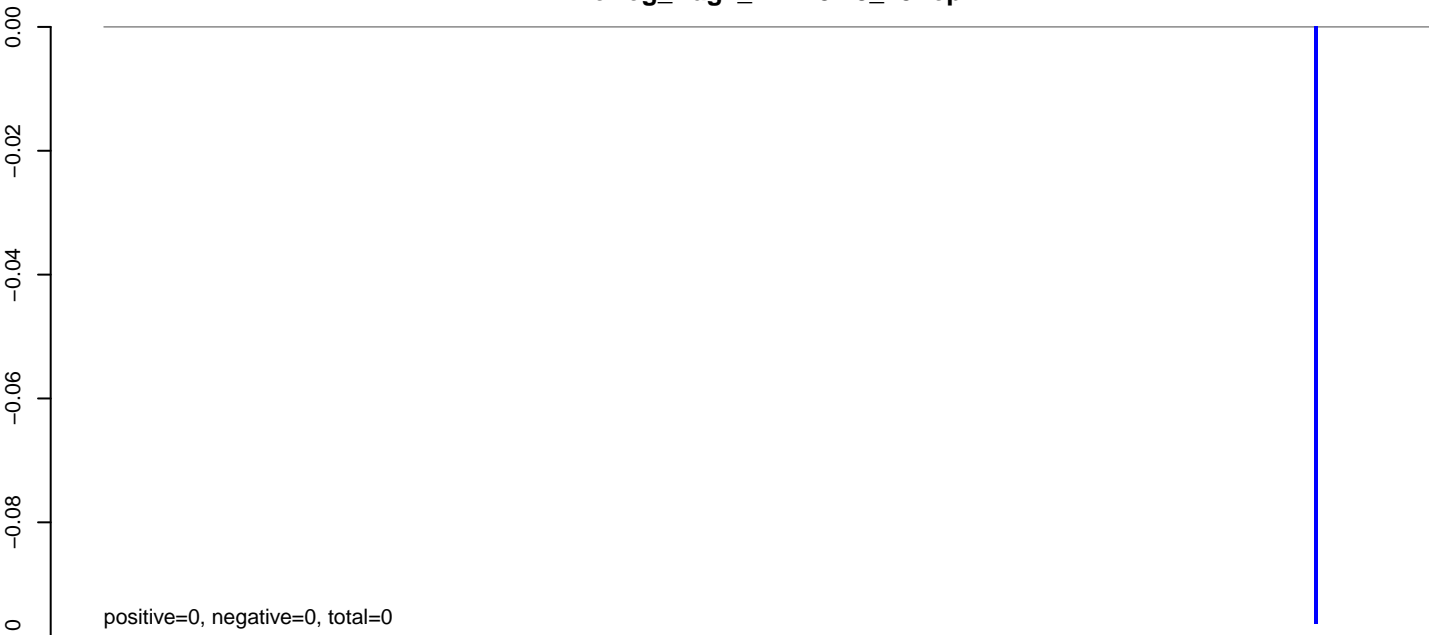
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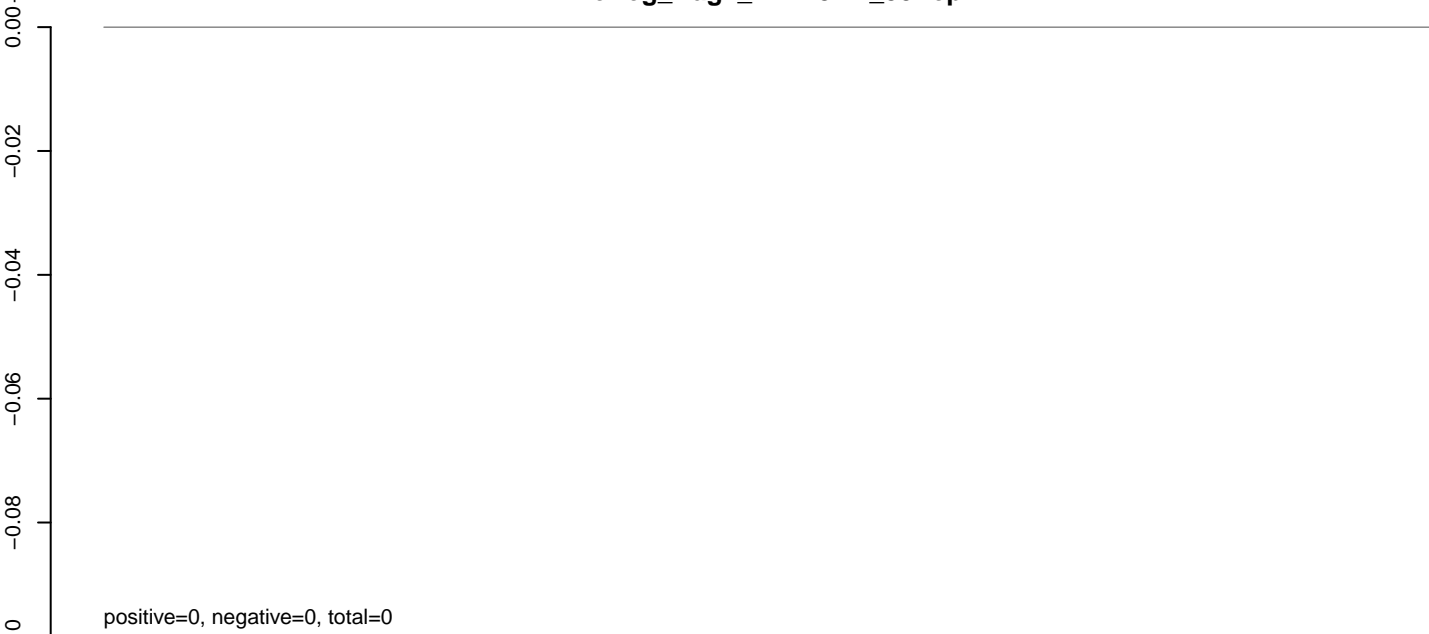
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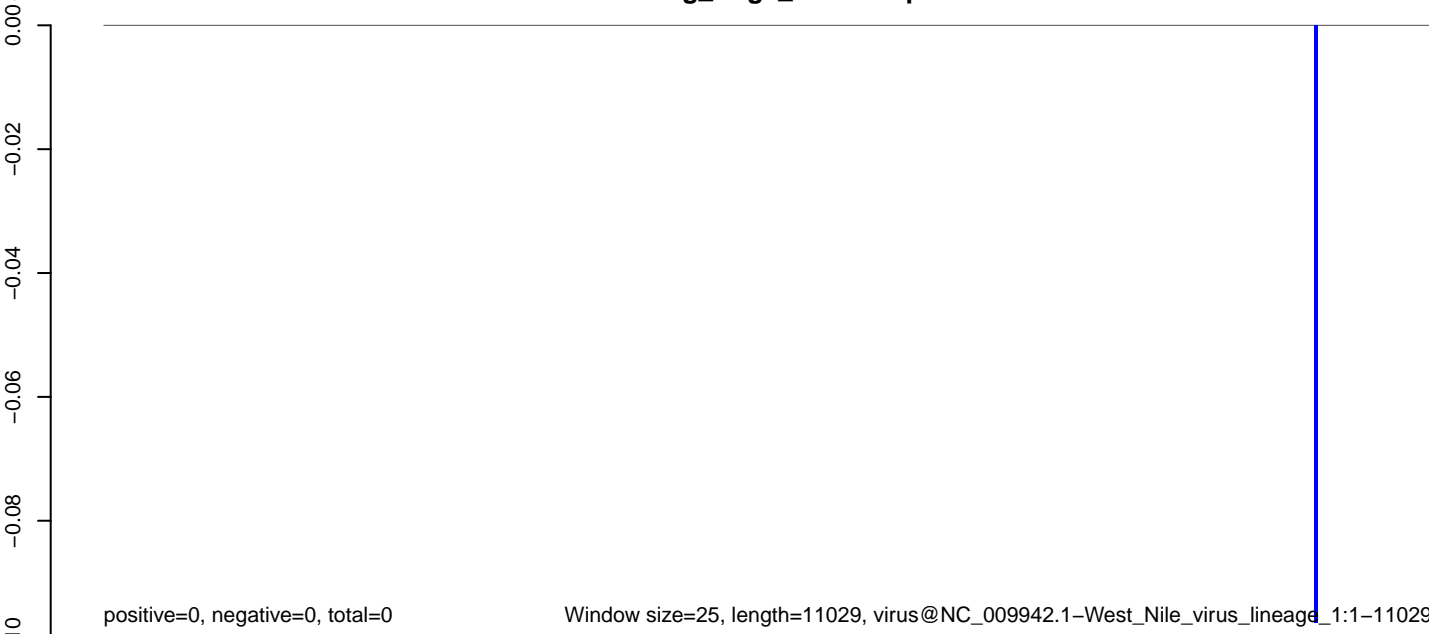
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AeAeg_Aag2_NL.M3.rep



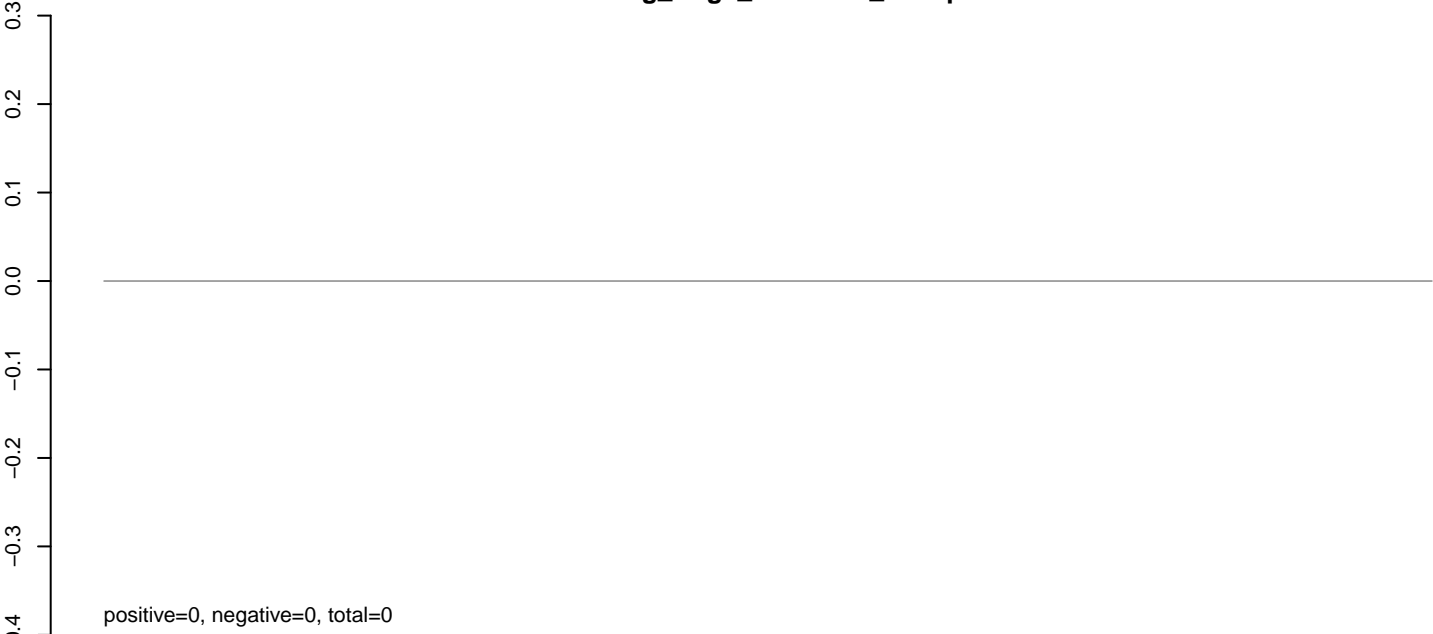
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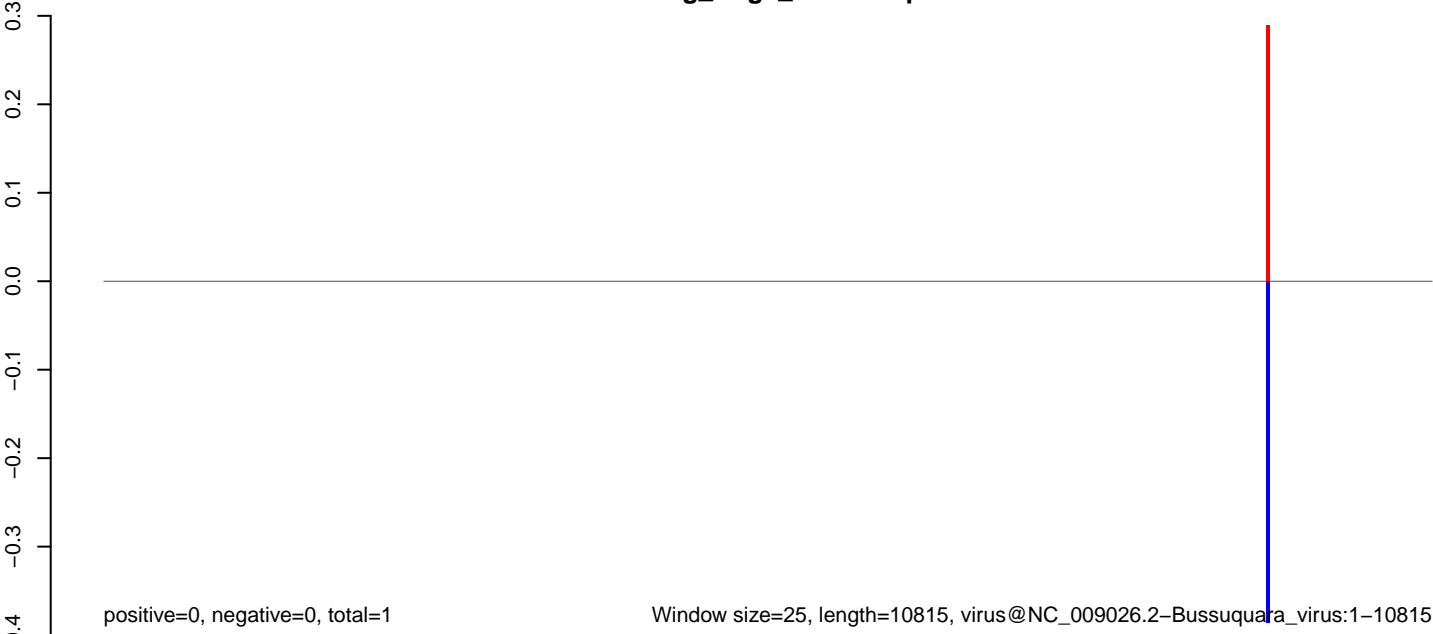
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positive=0, negative=0, total=0

AeAeg_Aag2_NL.M3.rep



positive=0, negative=0, total=1

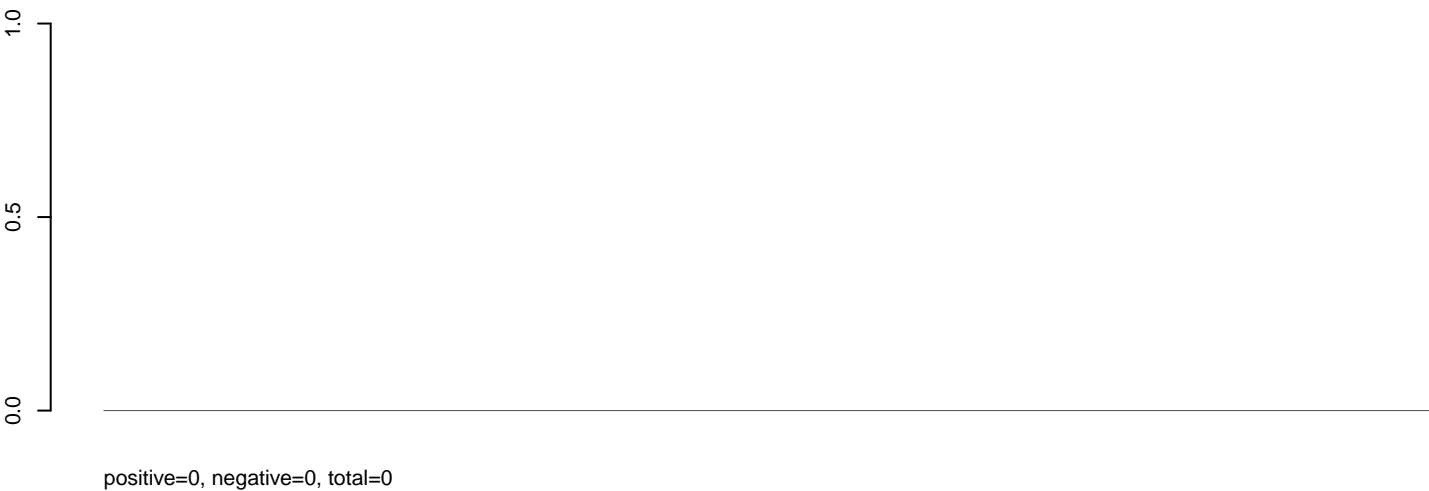
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AeAeg_Aag2_NL.M3.24_35.rep



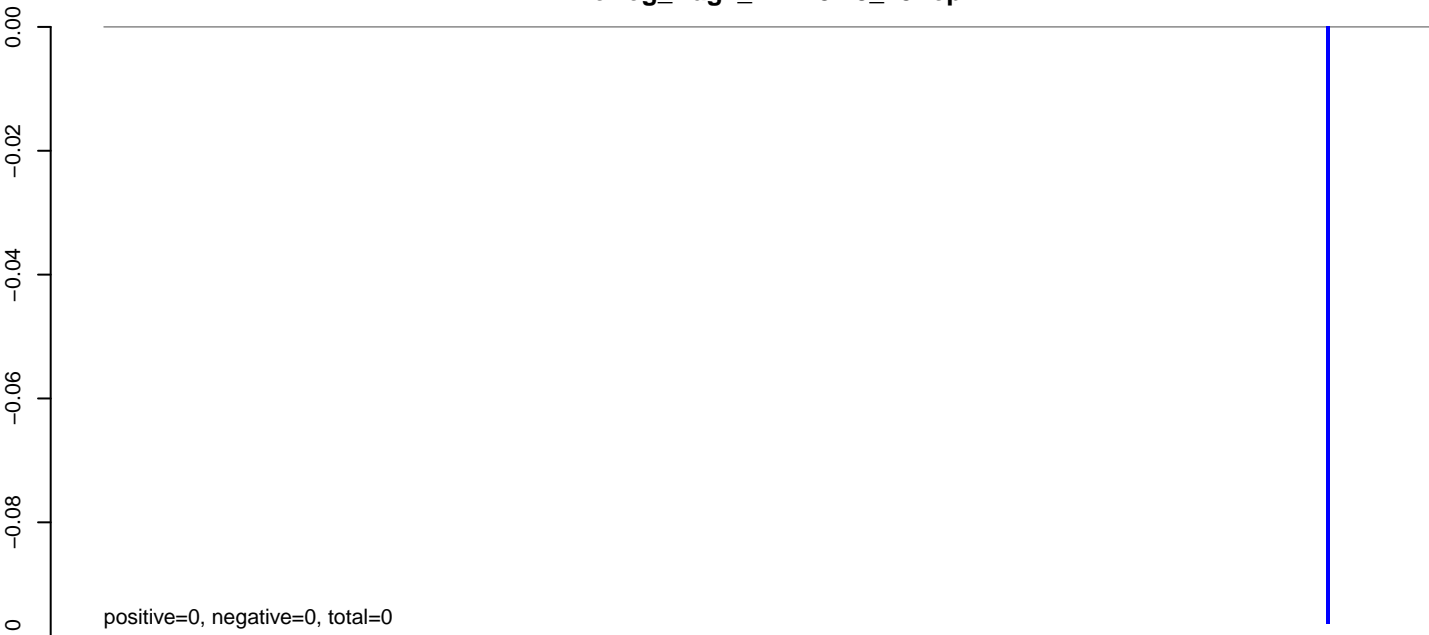
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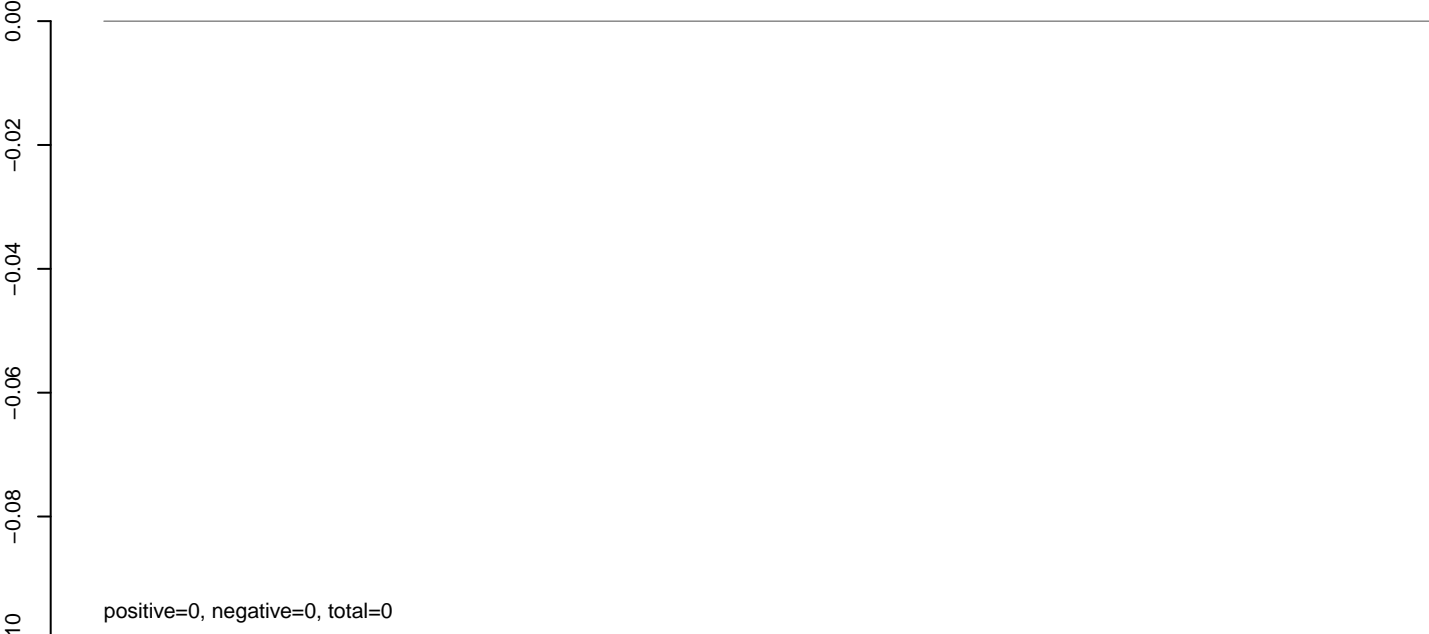
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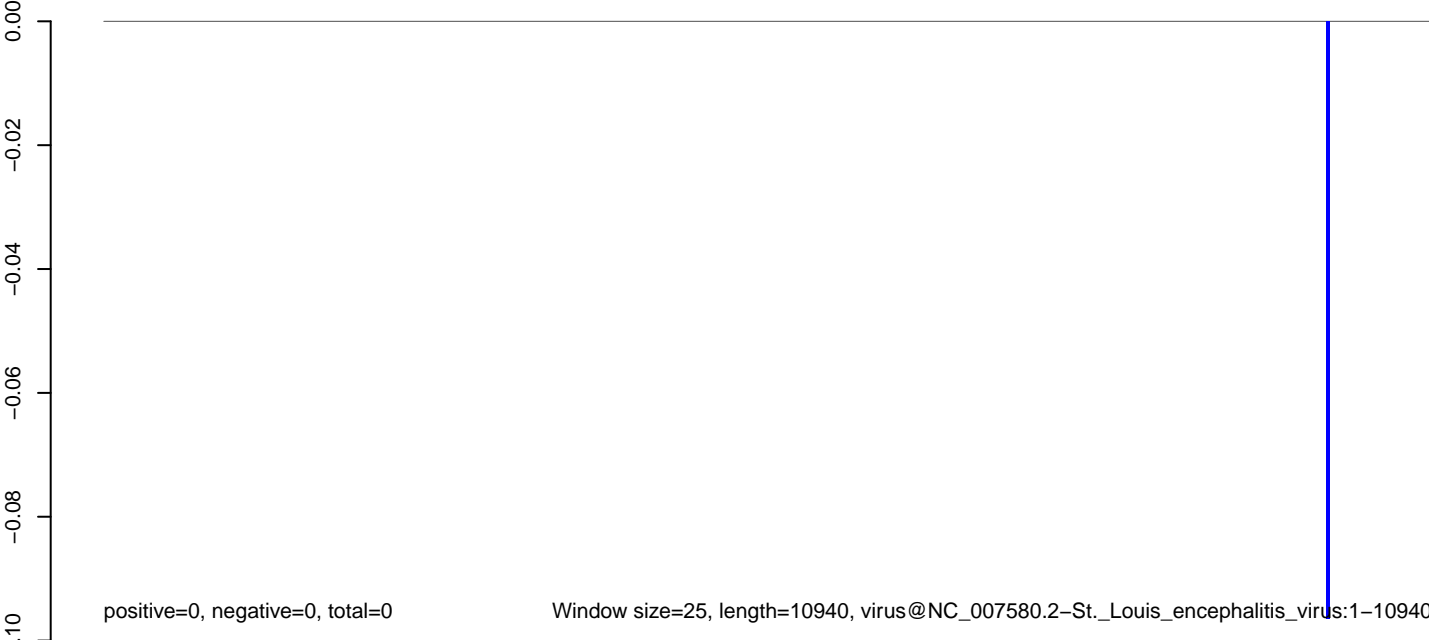
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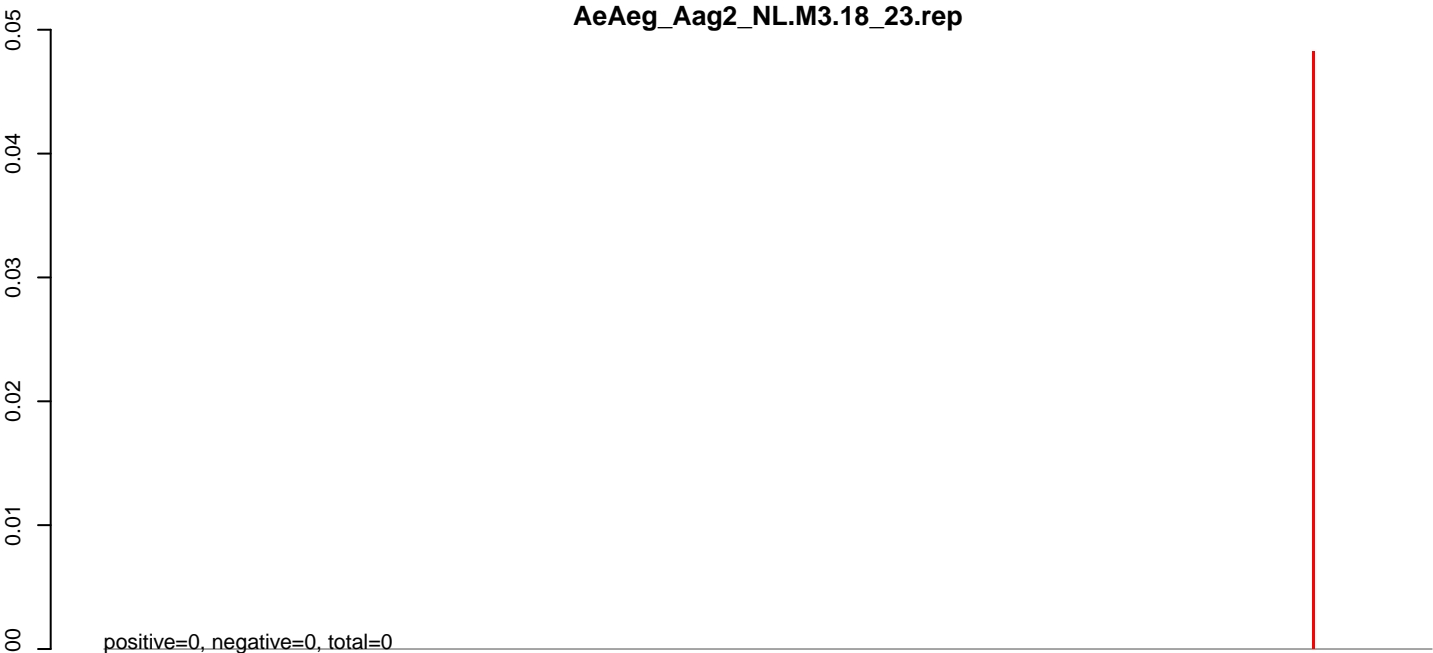
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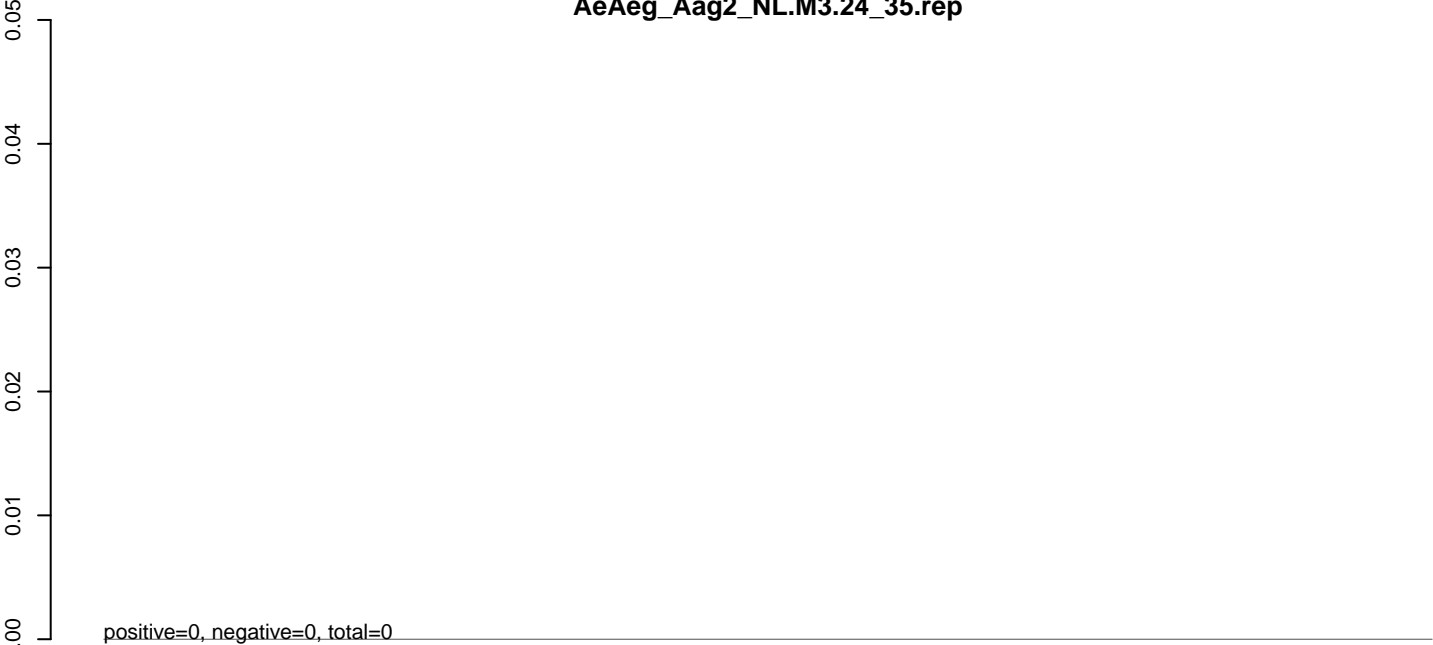
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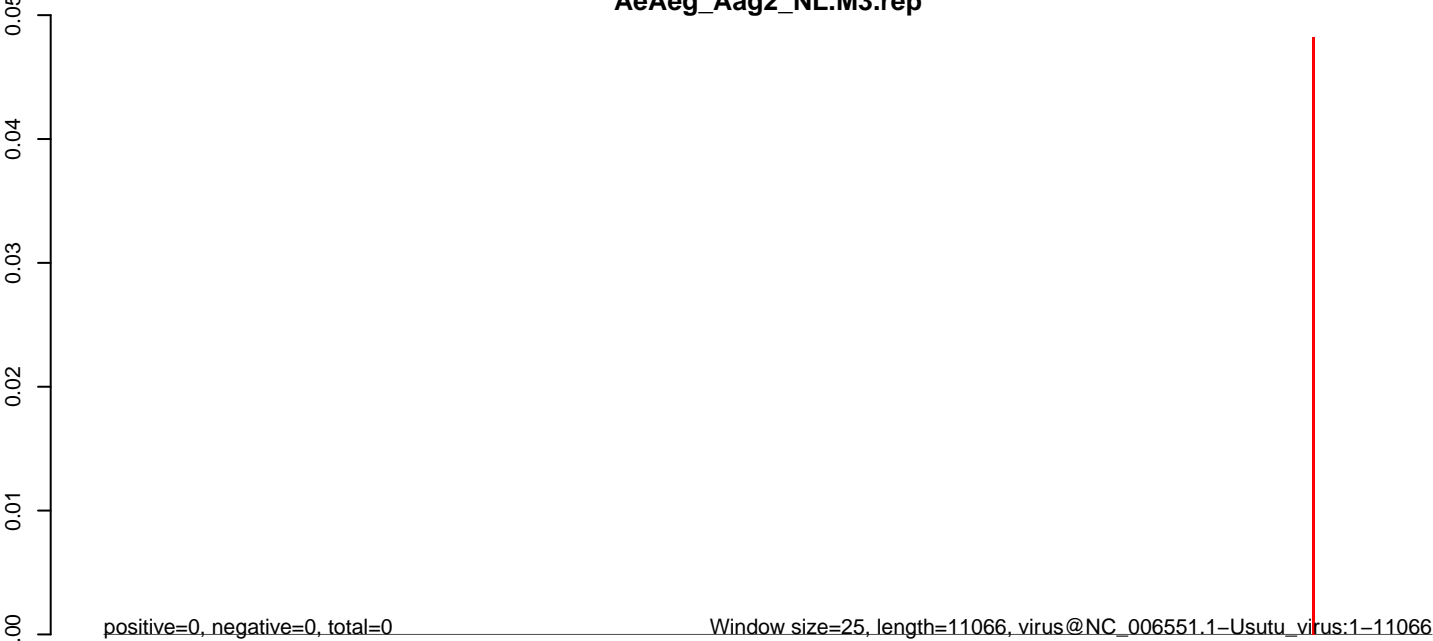
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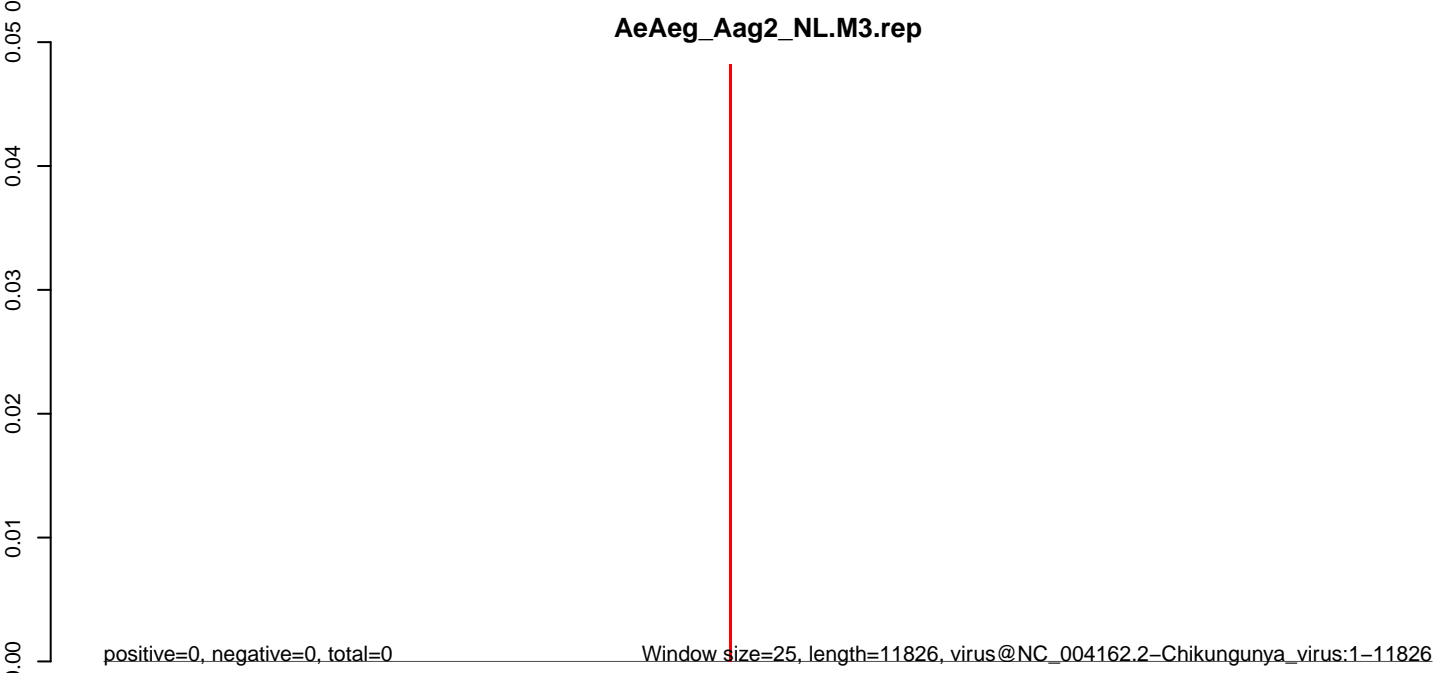
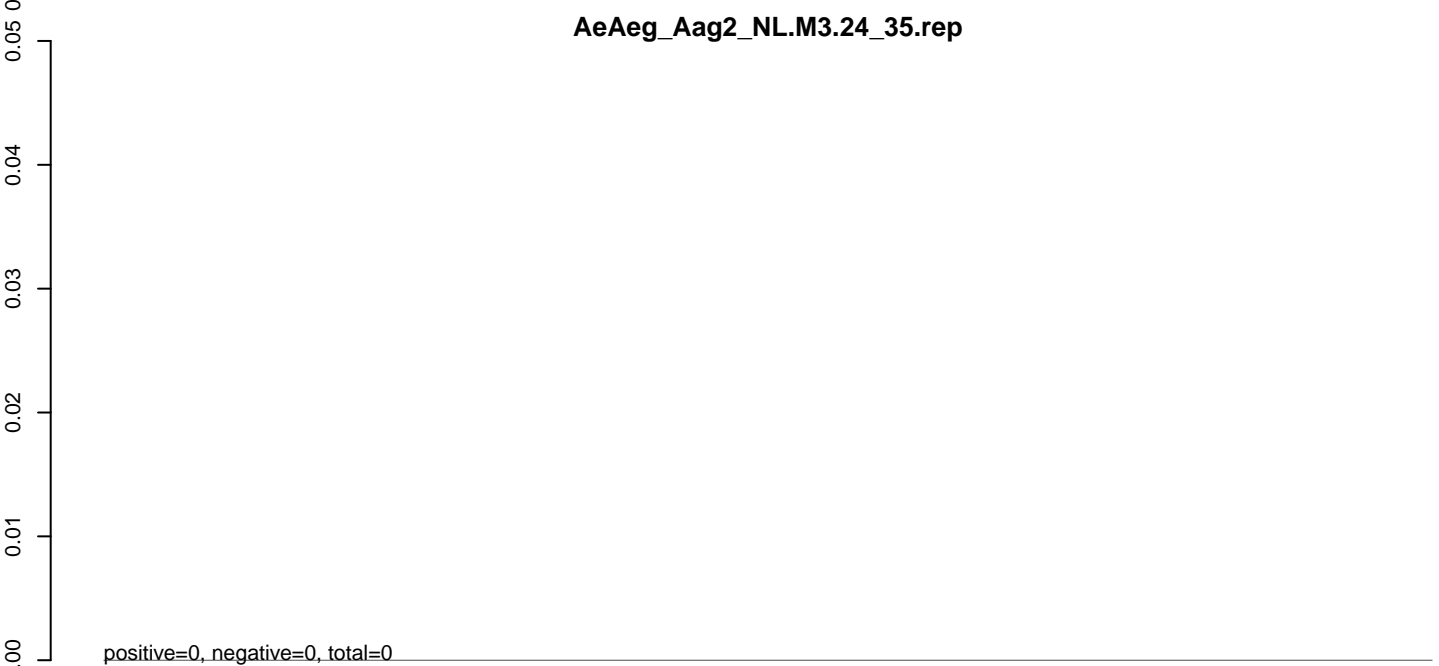
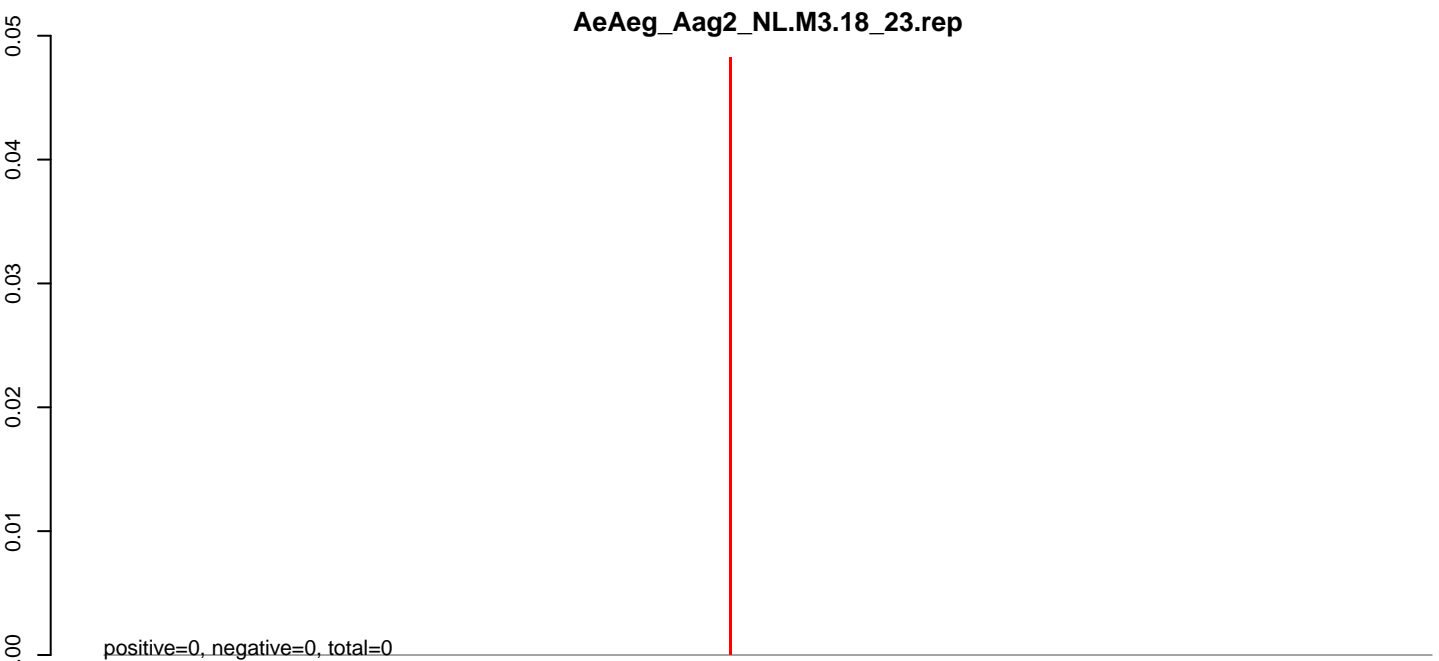
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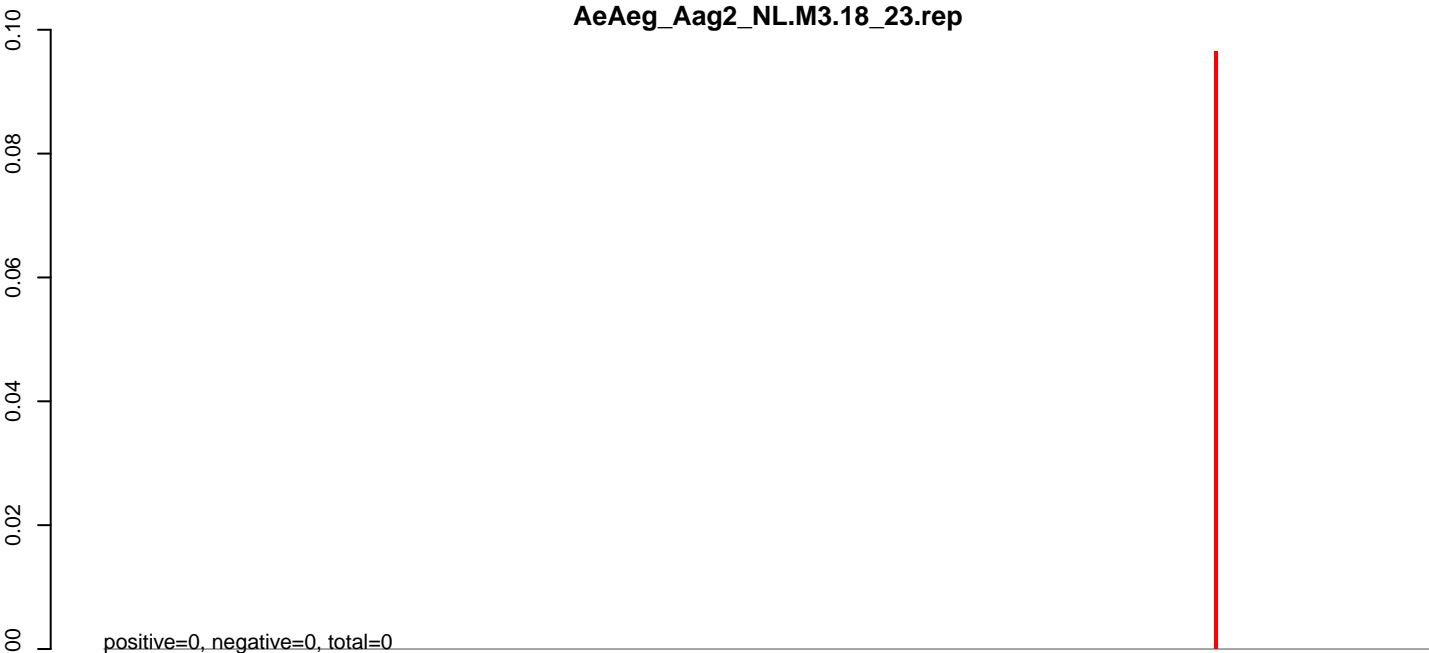


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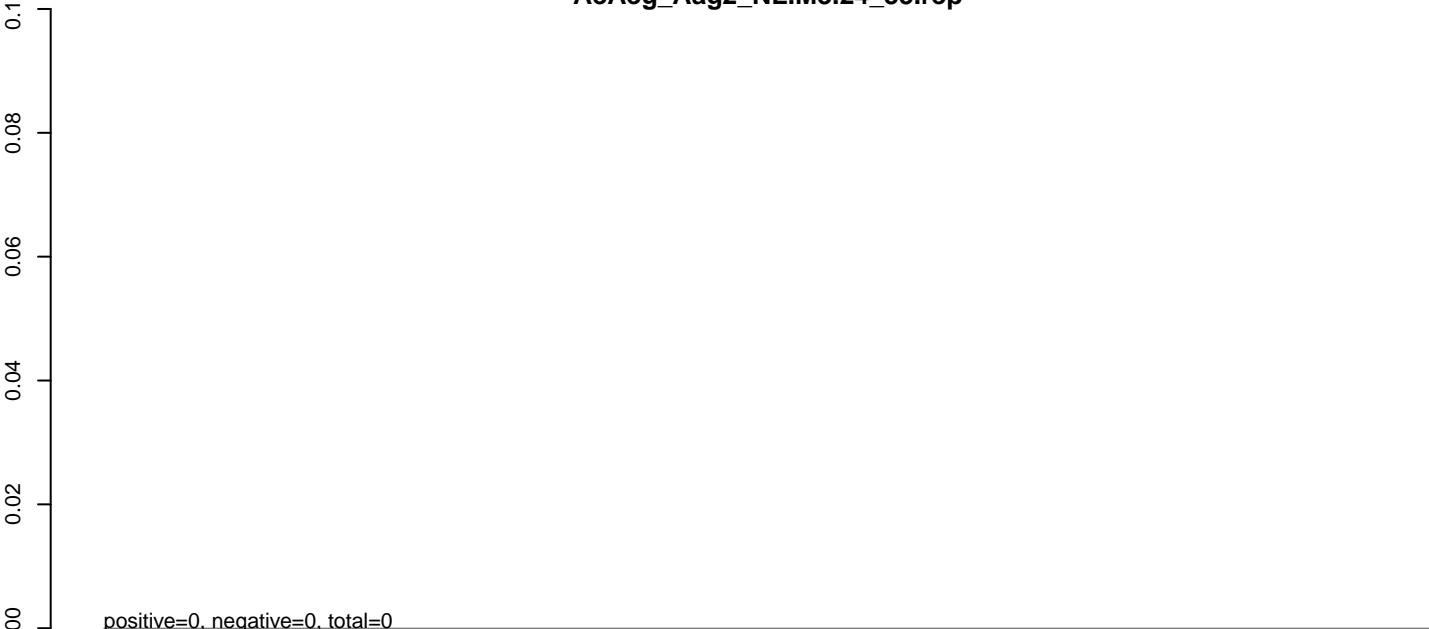


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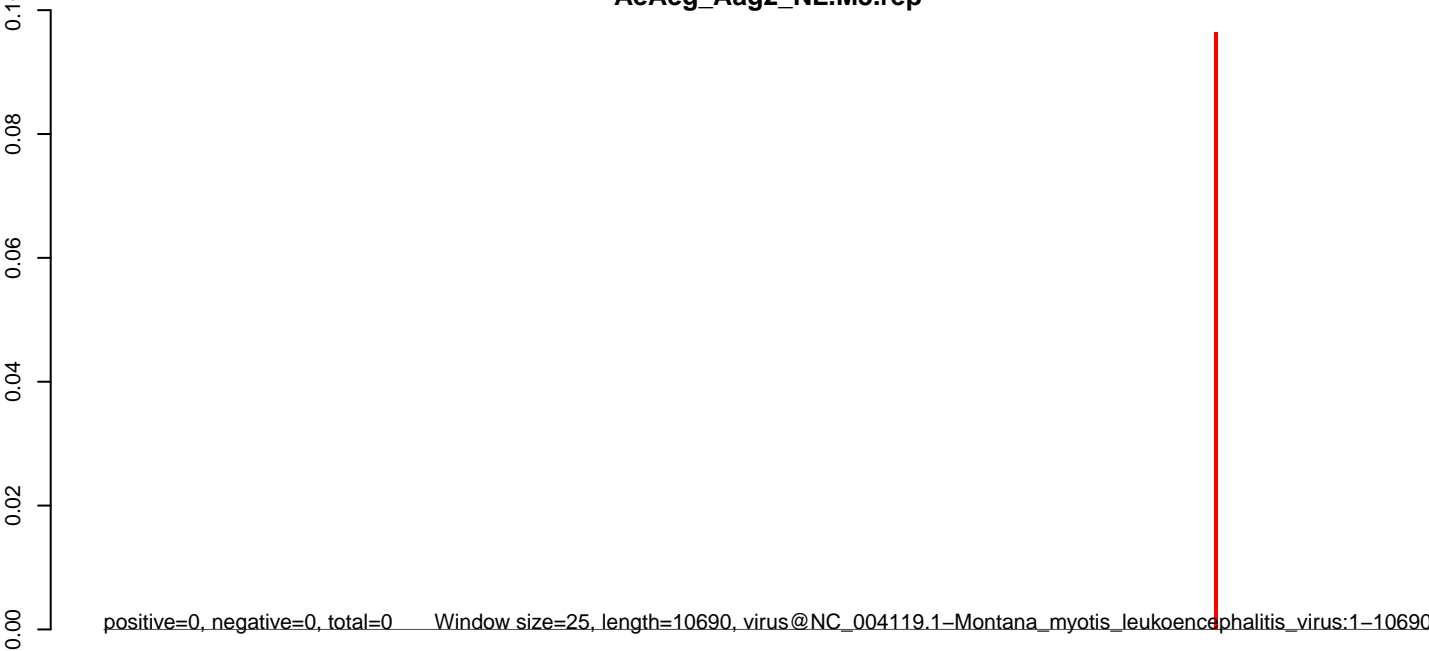
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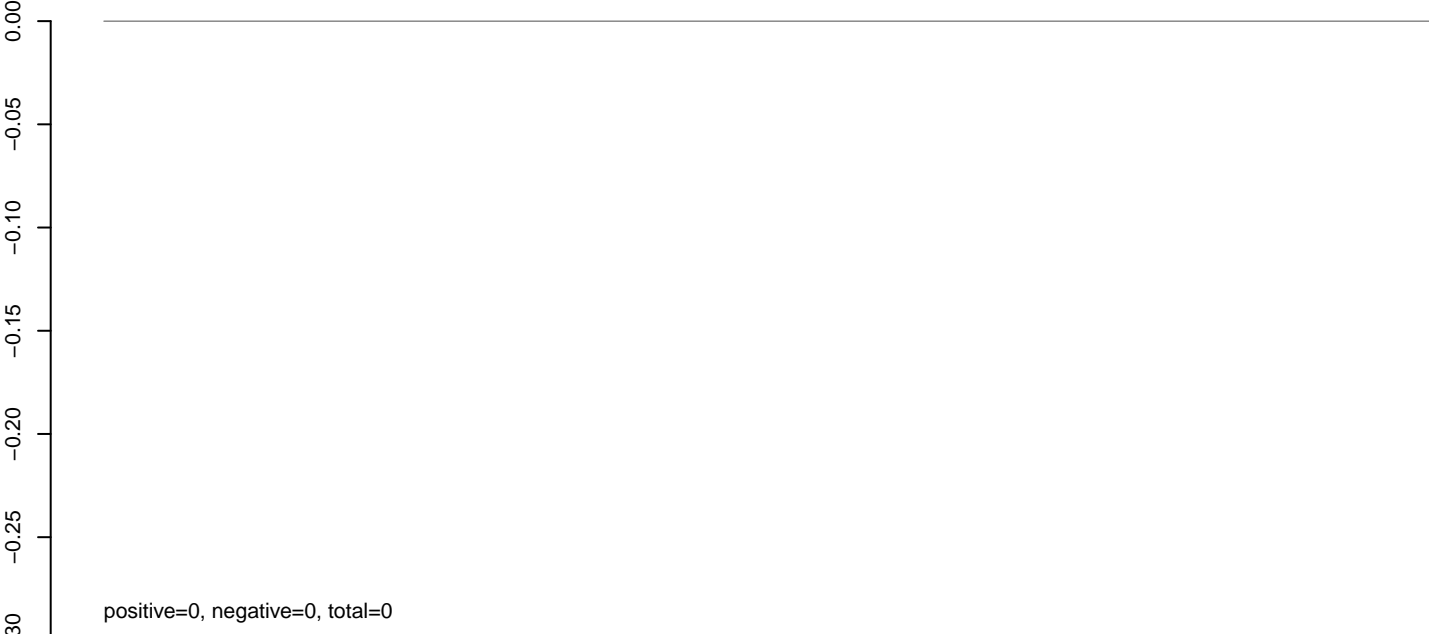


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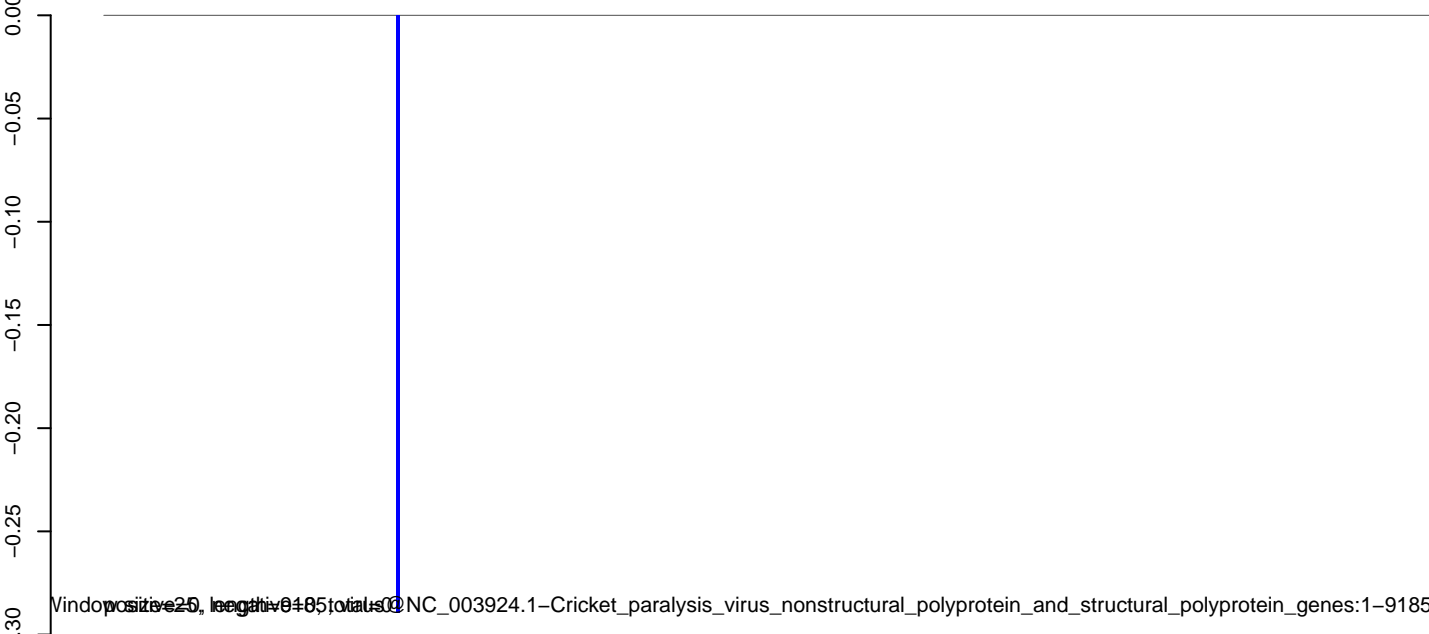
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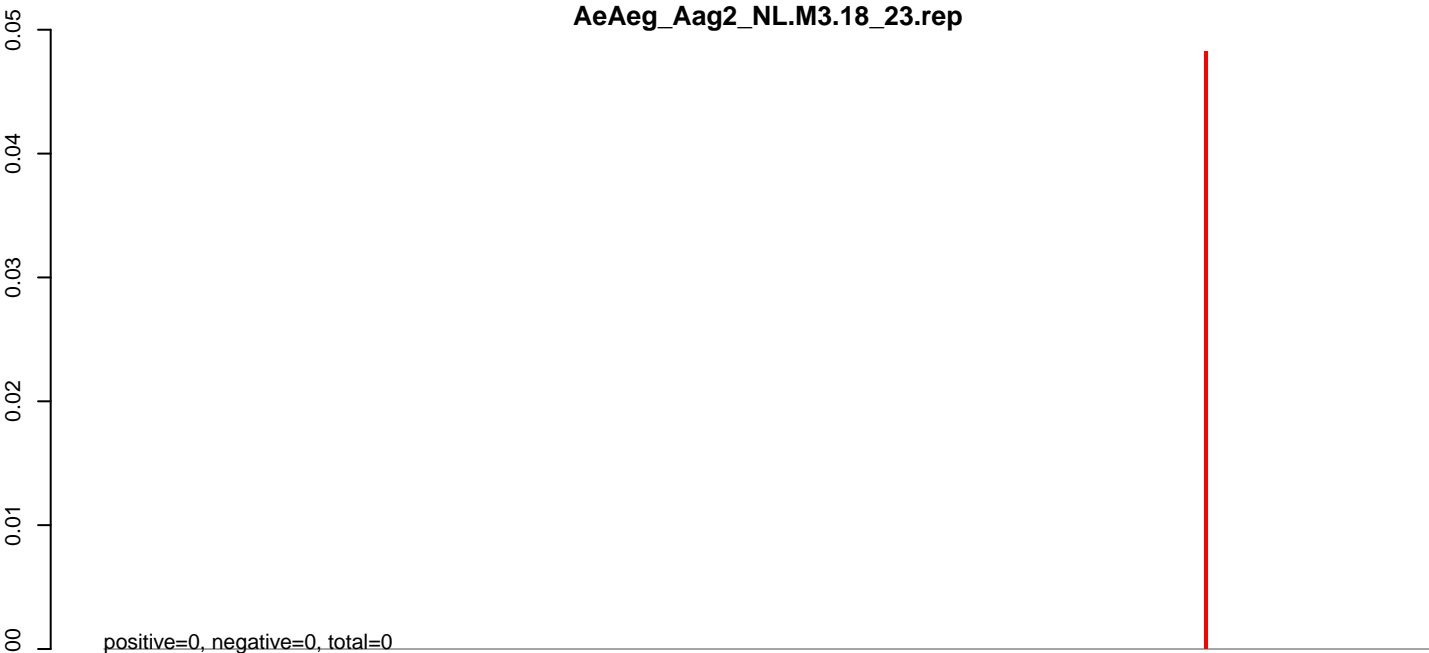
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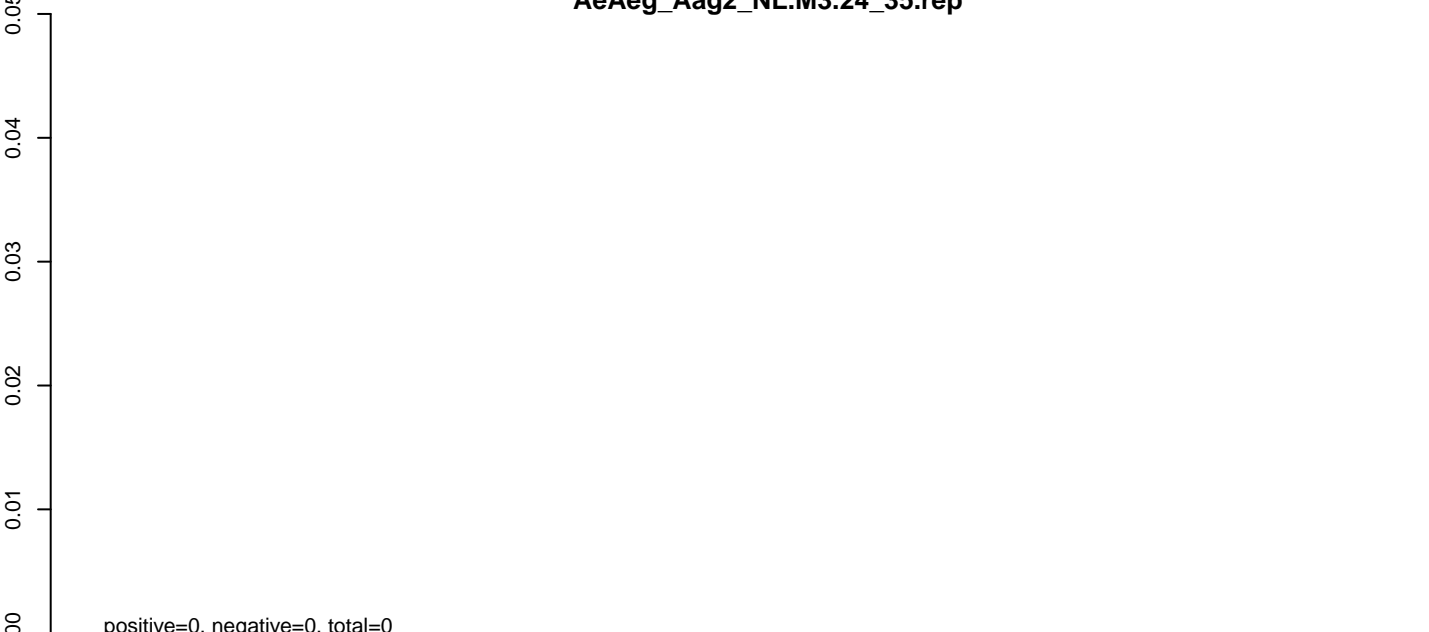
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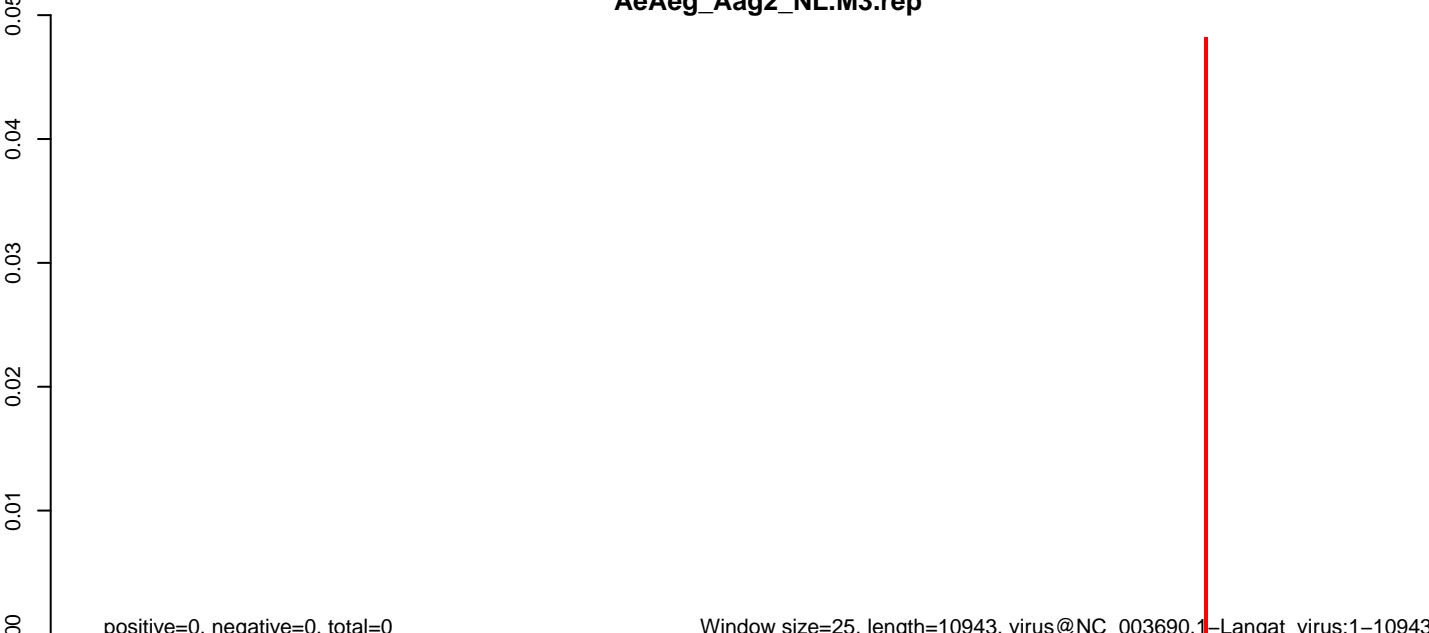
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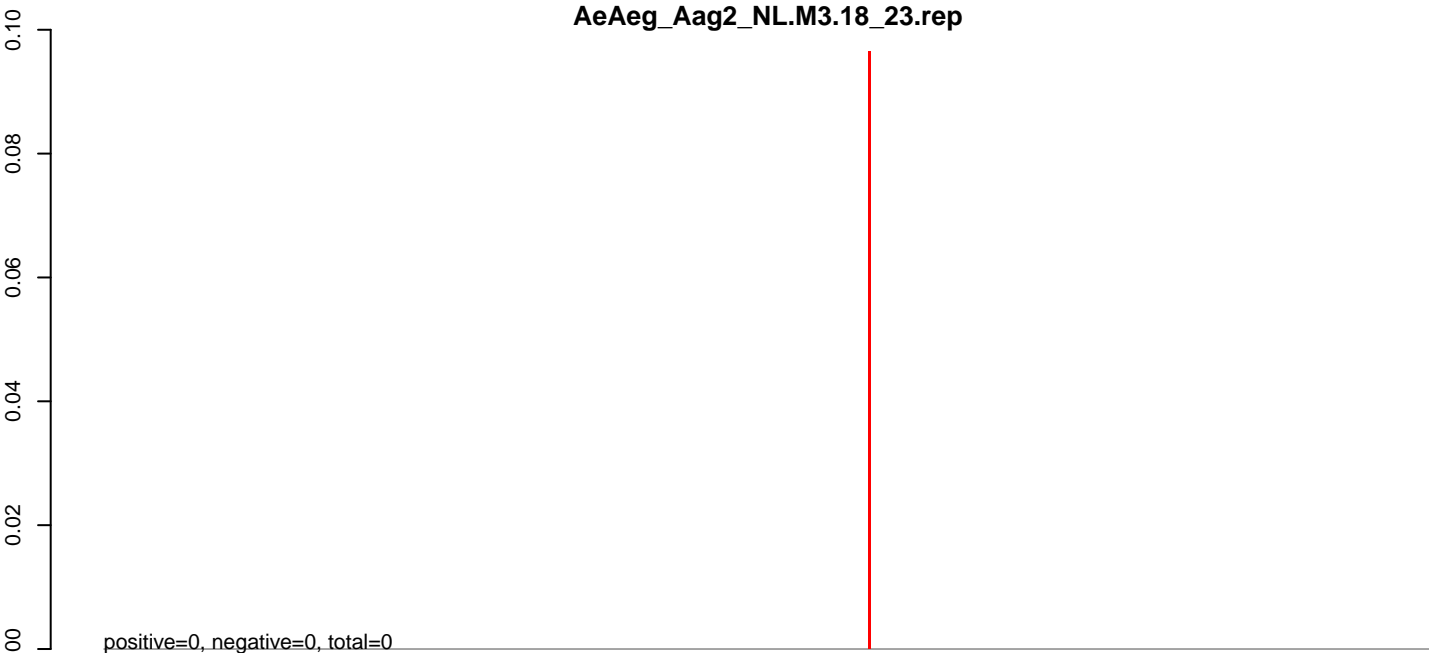


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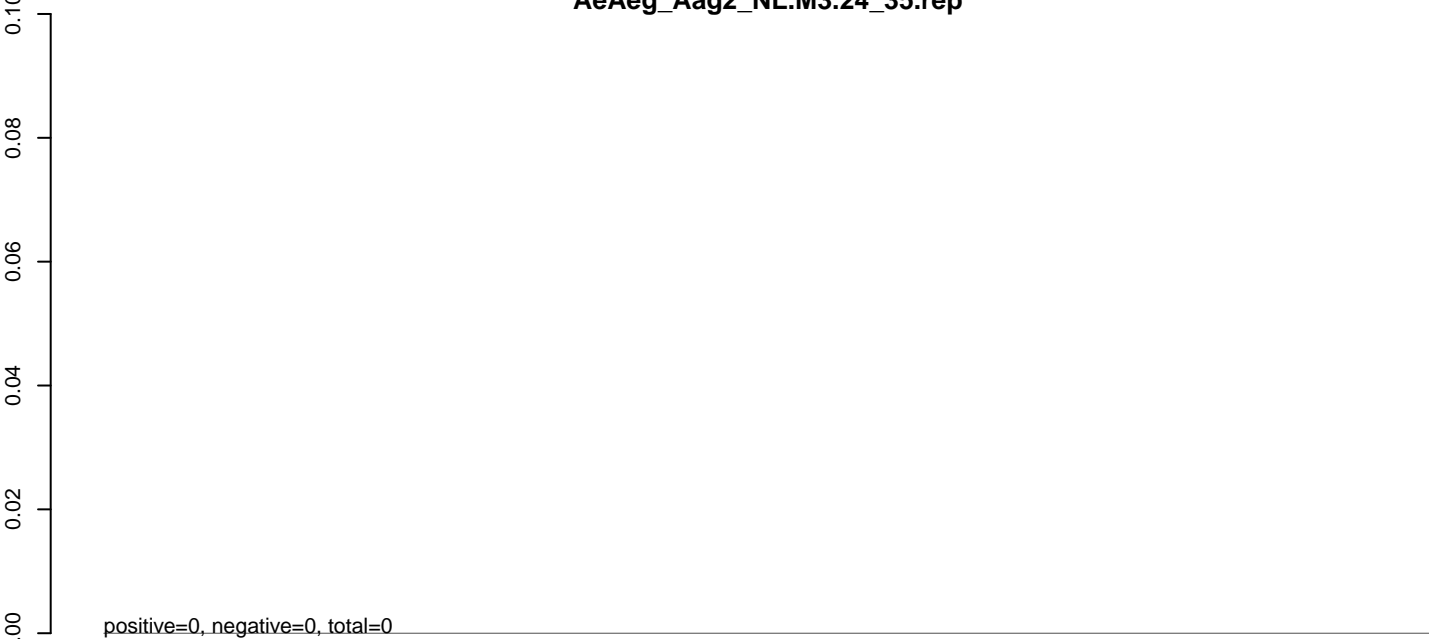


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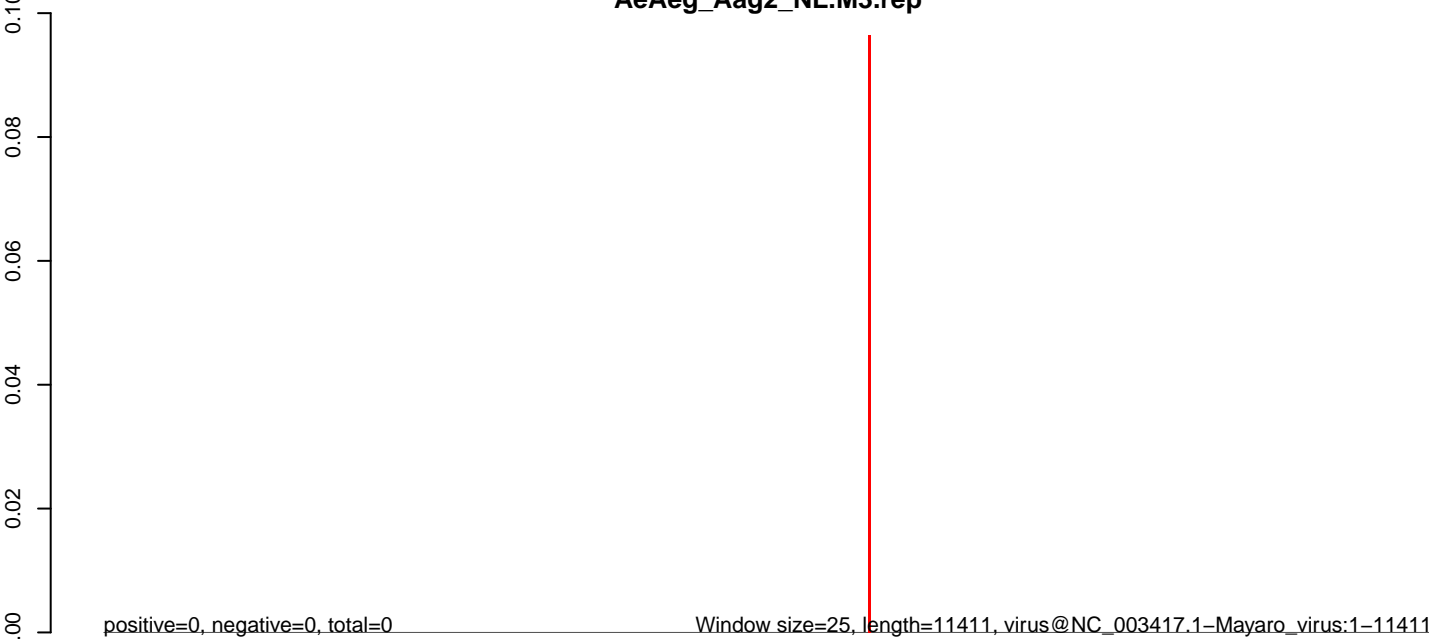
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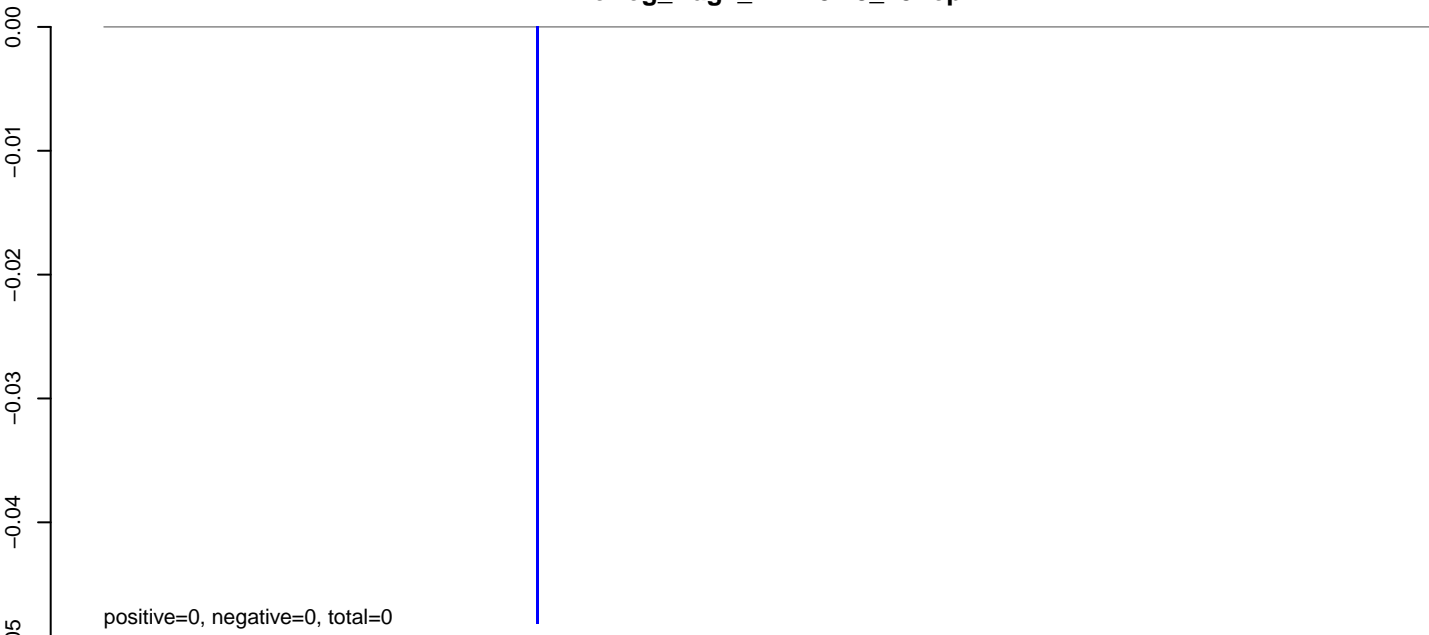


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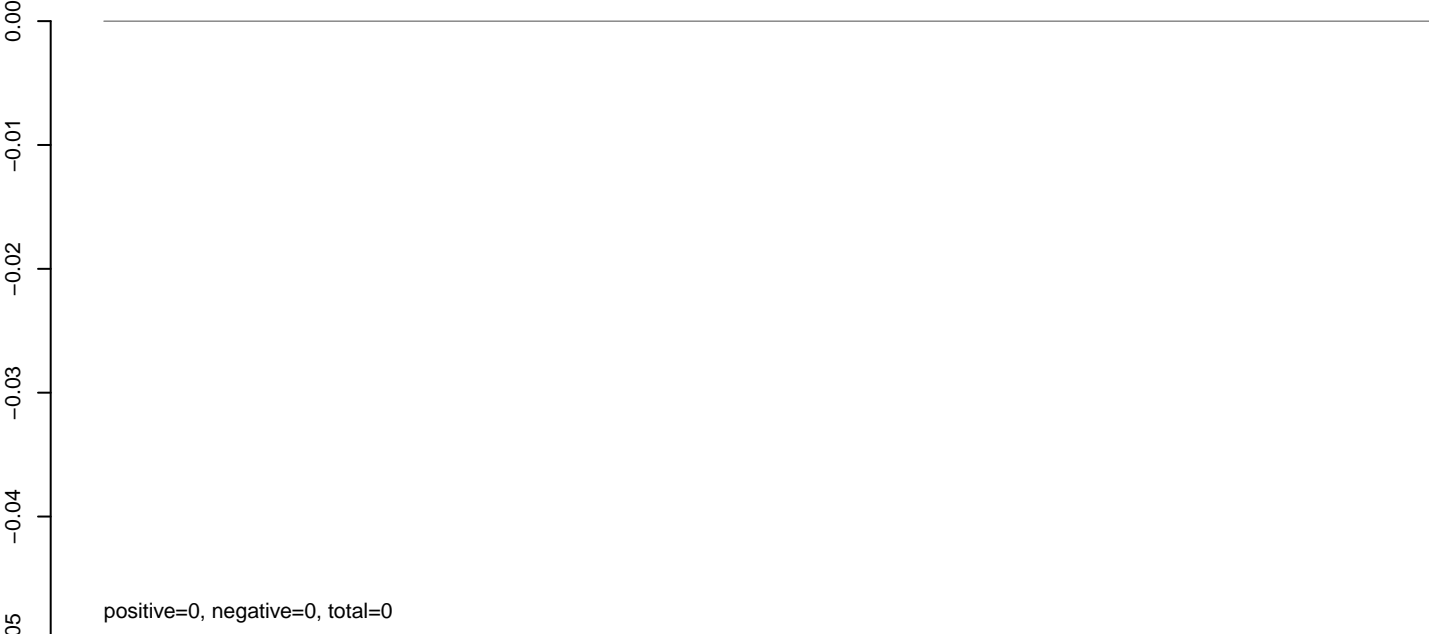


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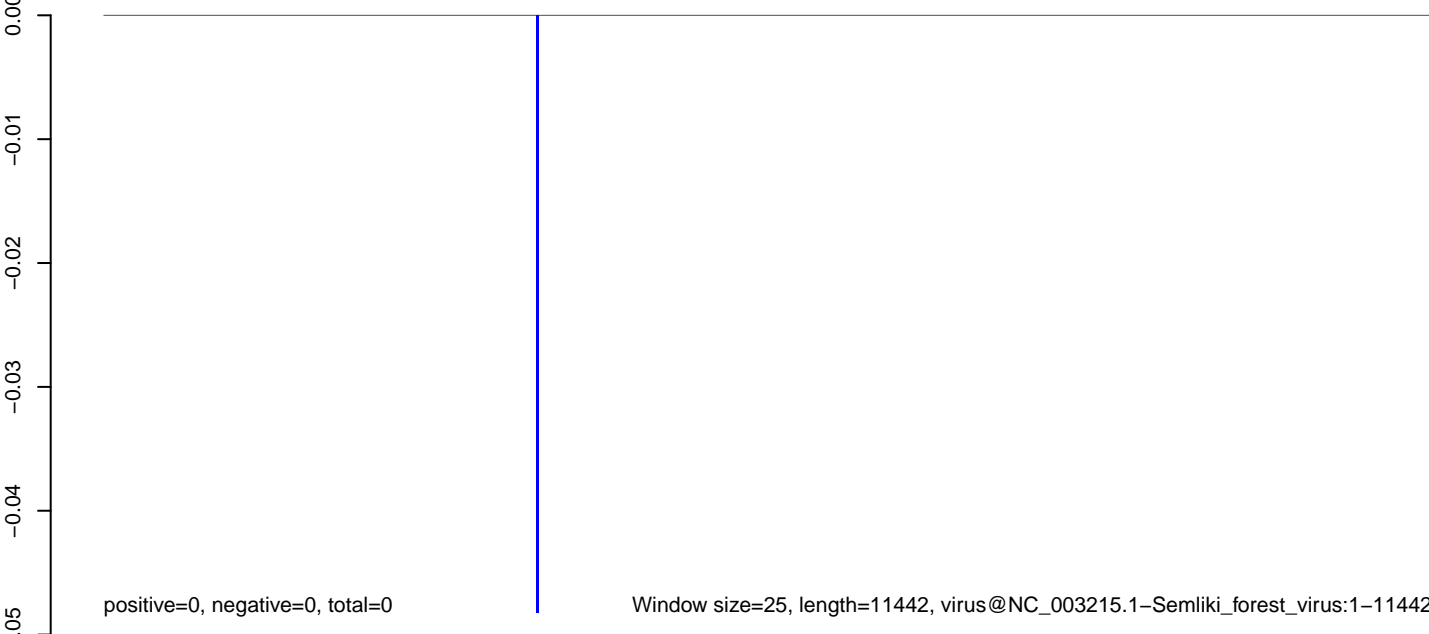
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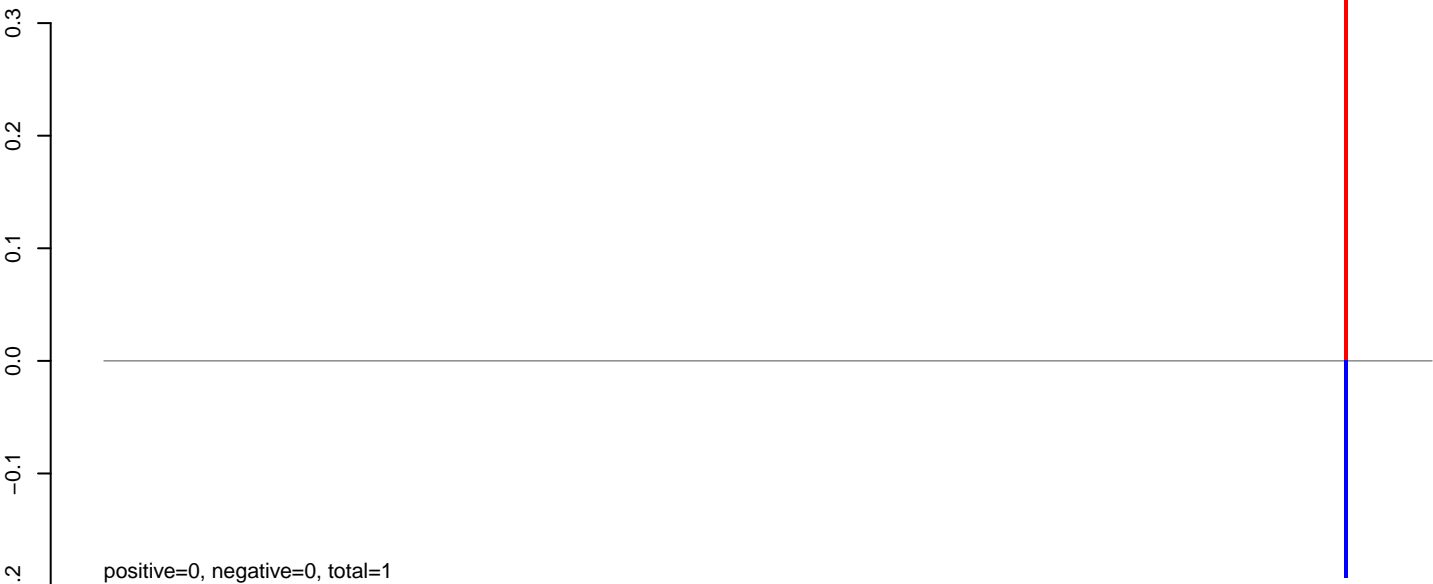
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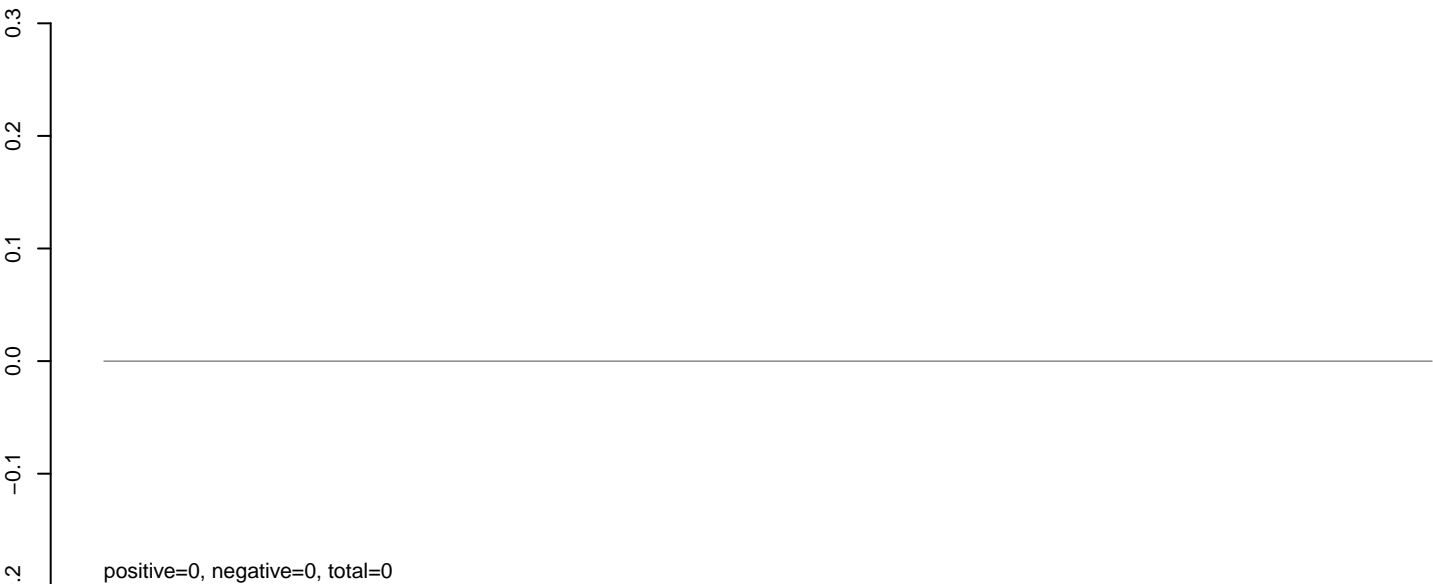
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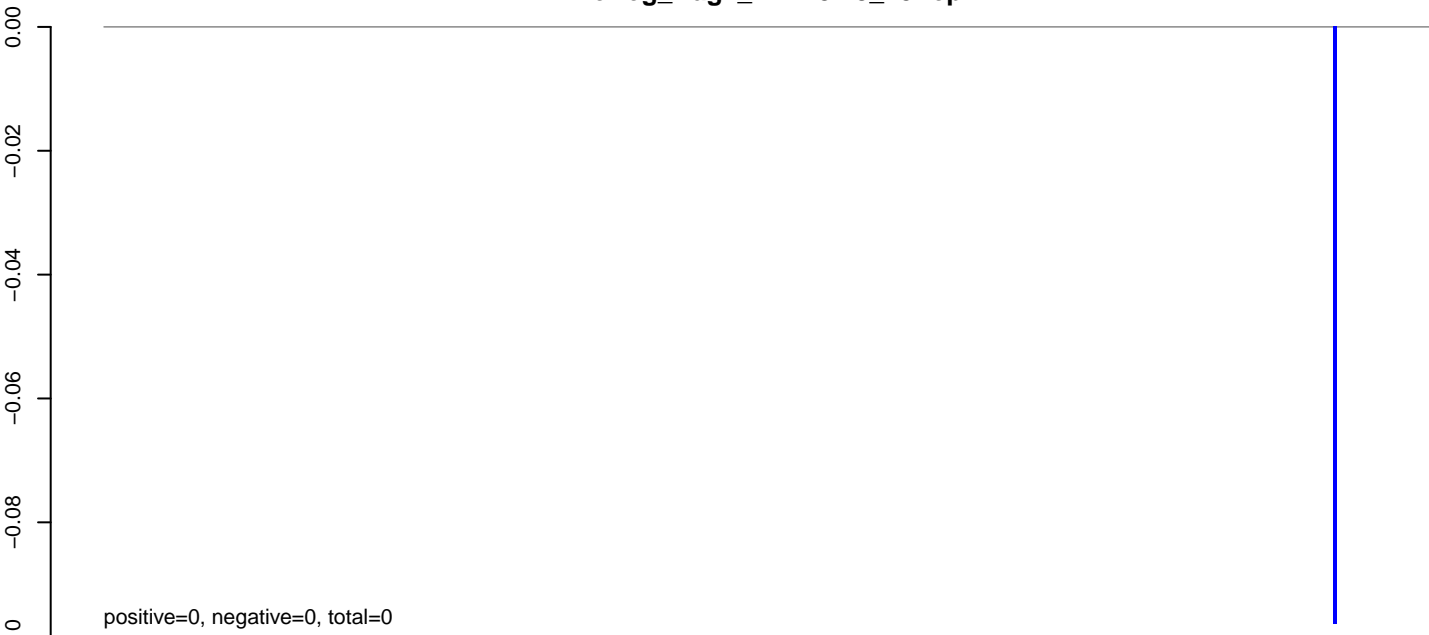
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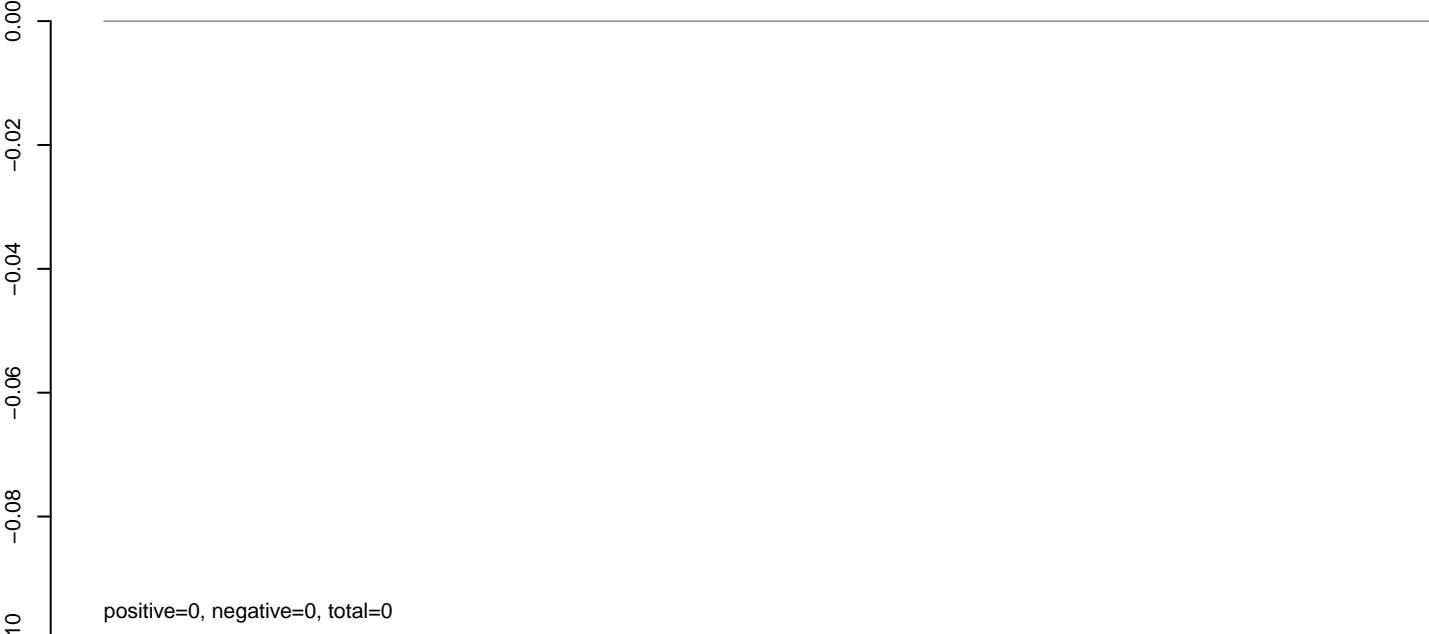
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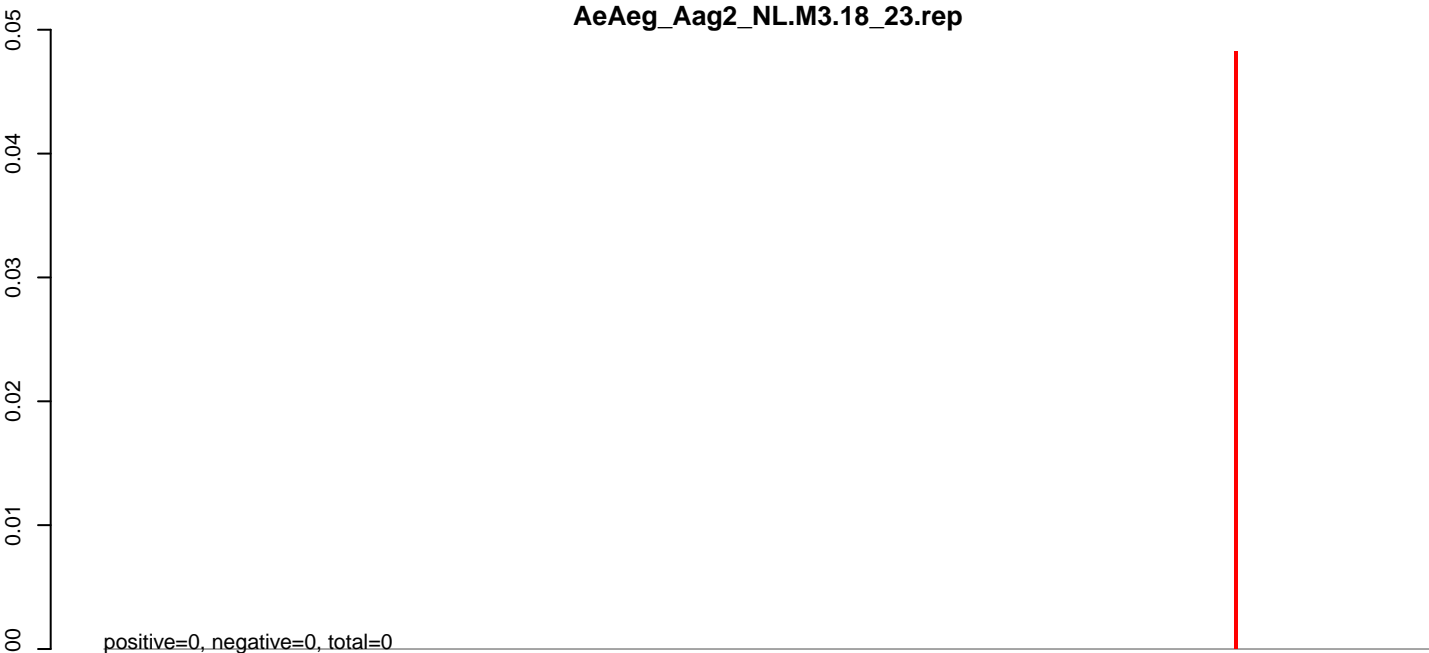
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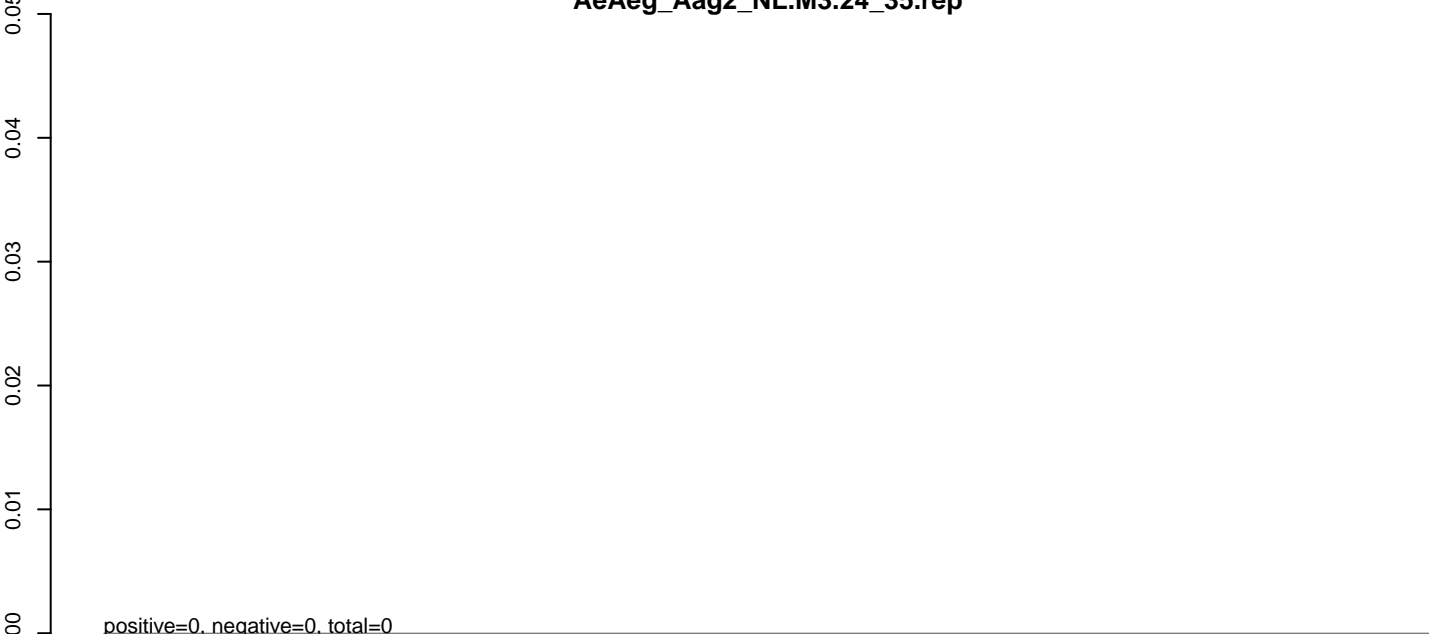
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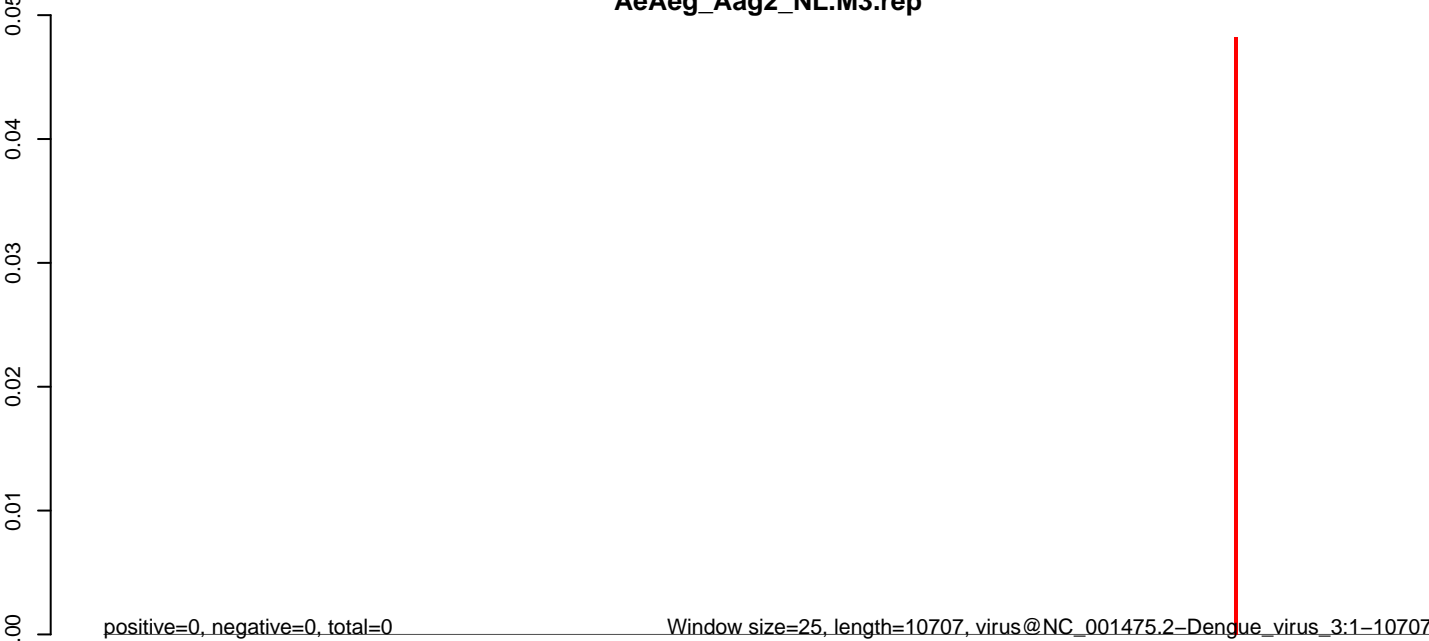
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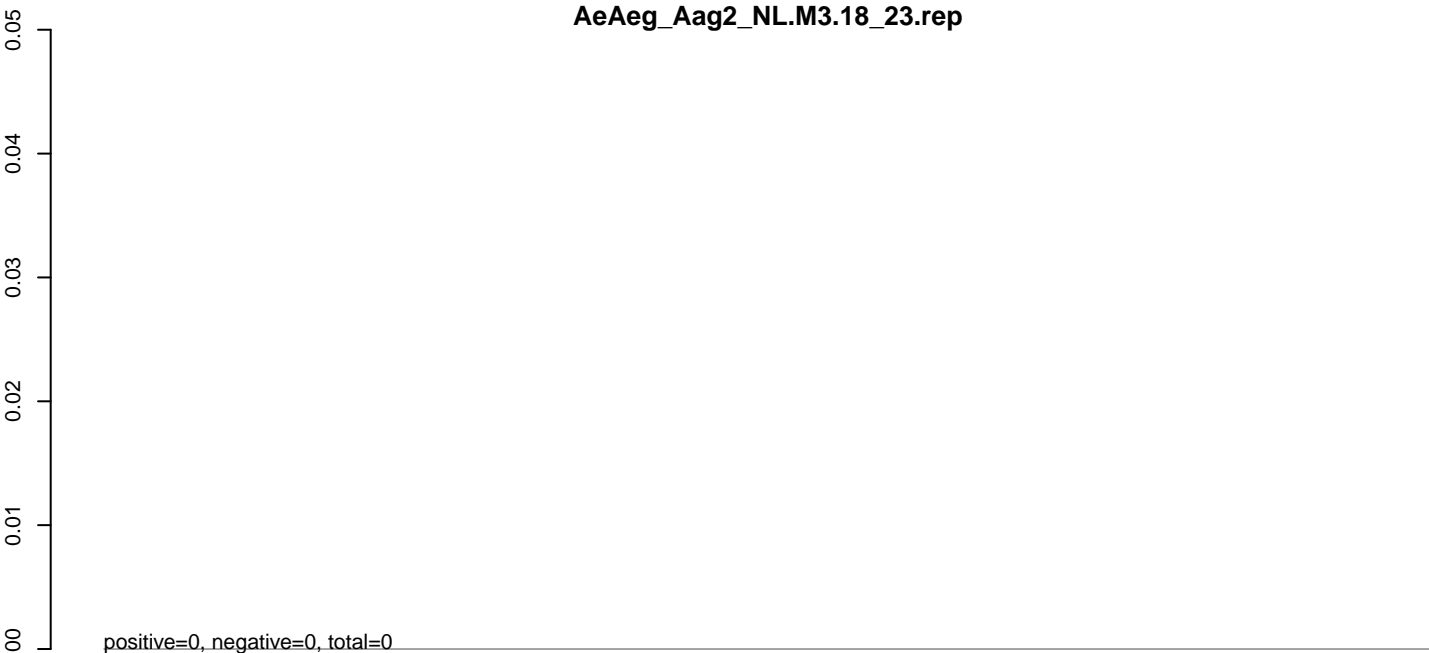


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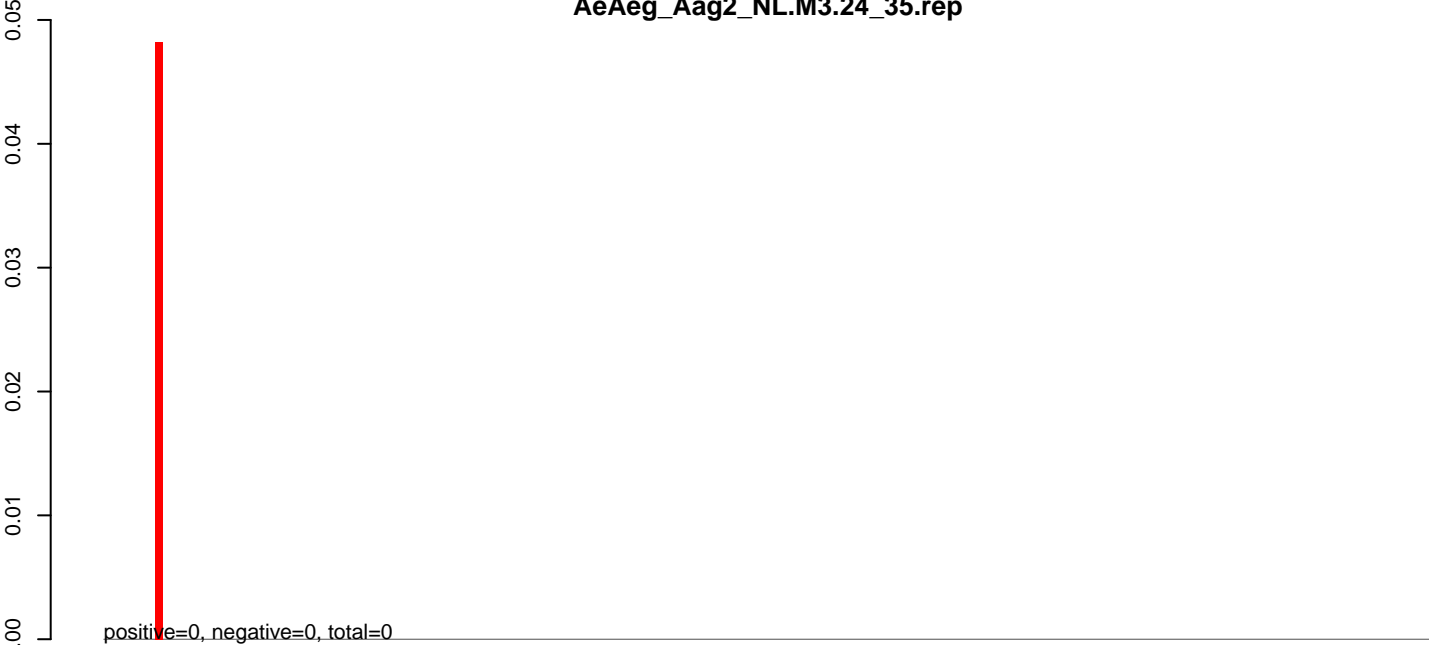


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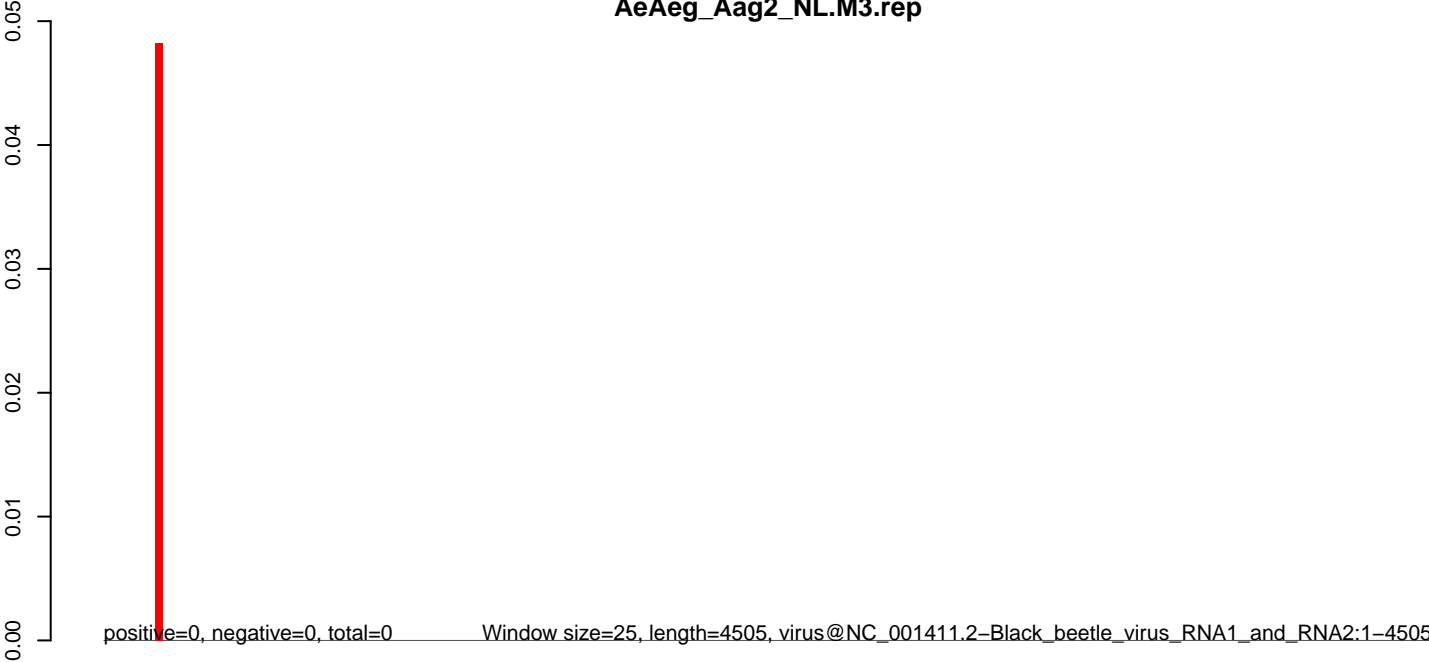
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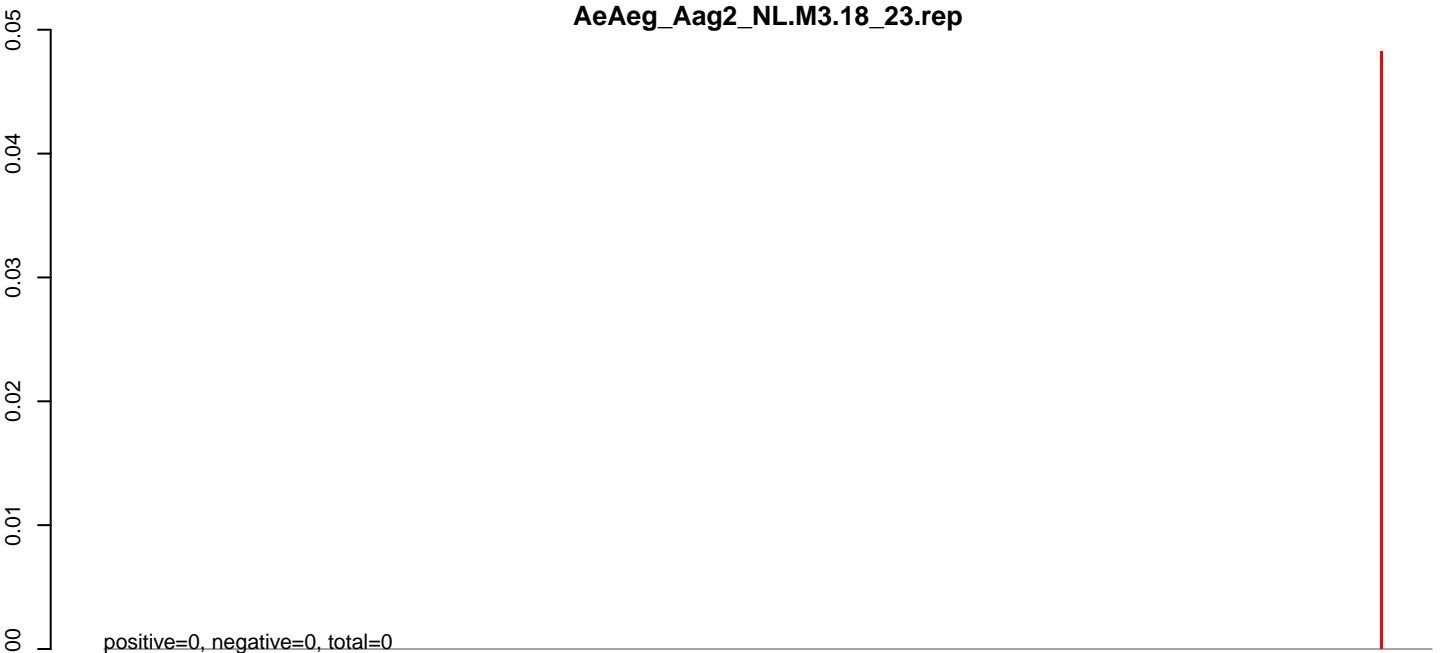
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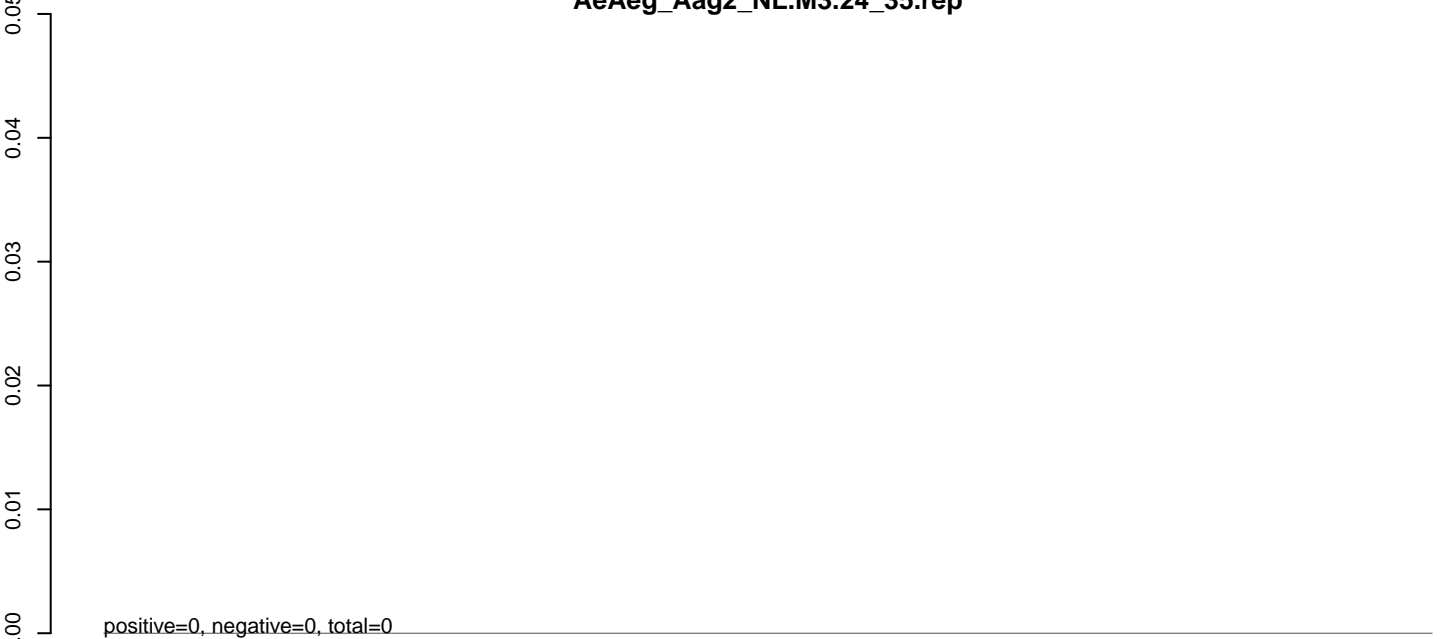
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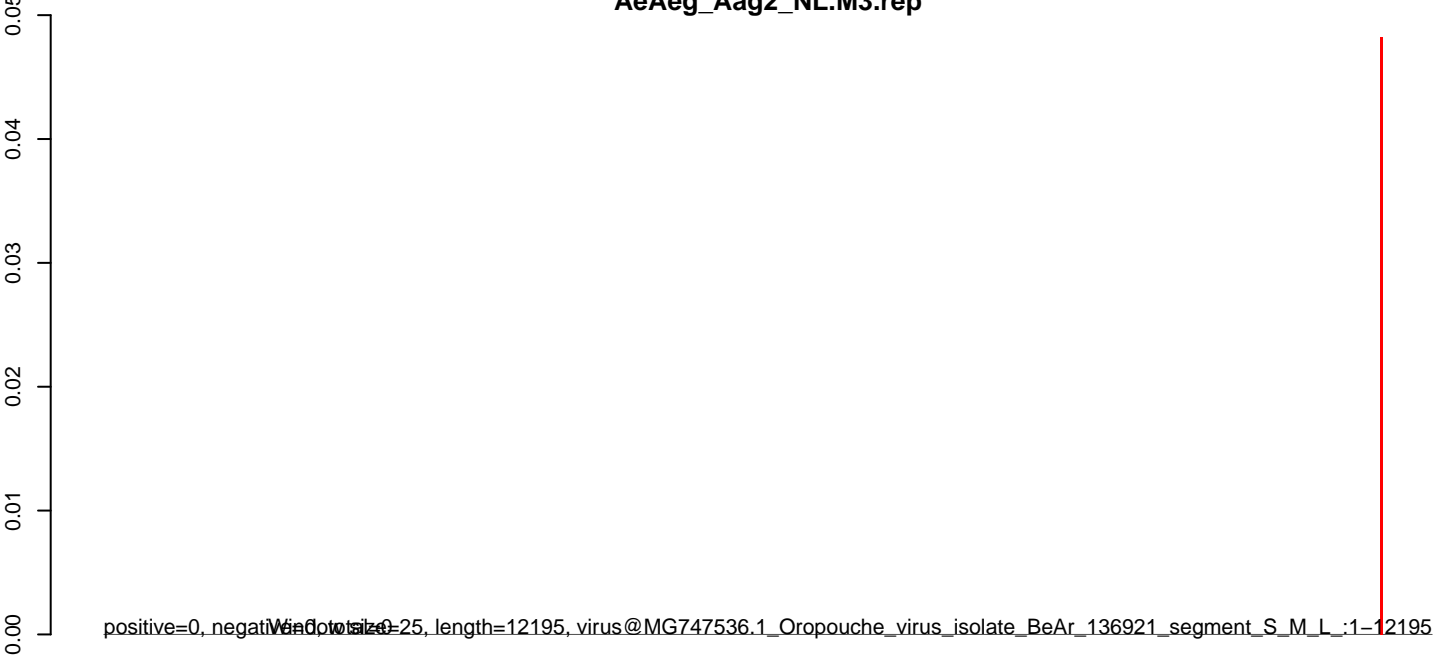
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AeAeg_Aag2_NL.M3.24_35.rep



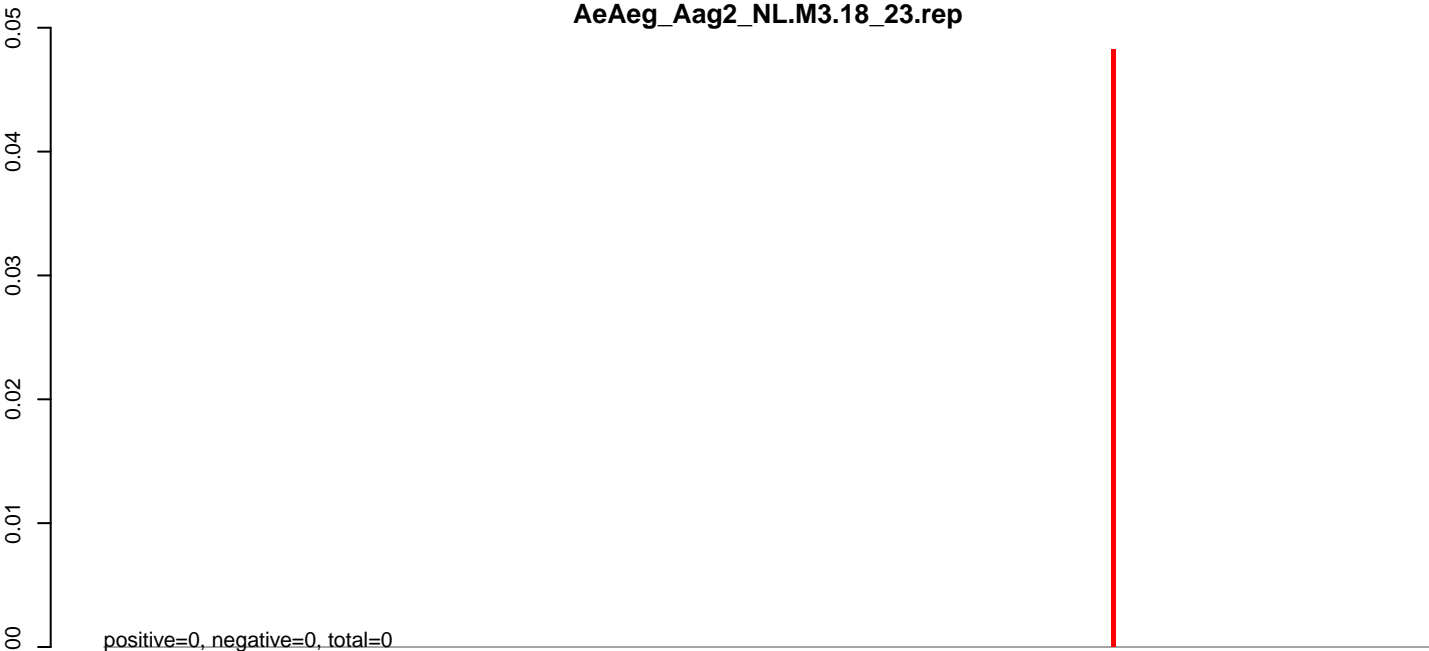
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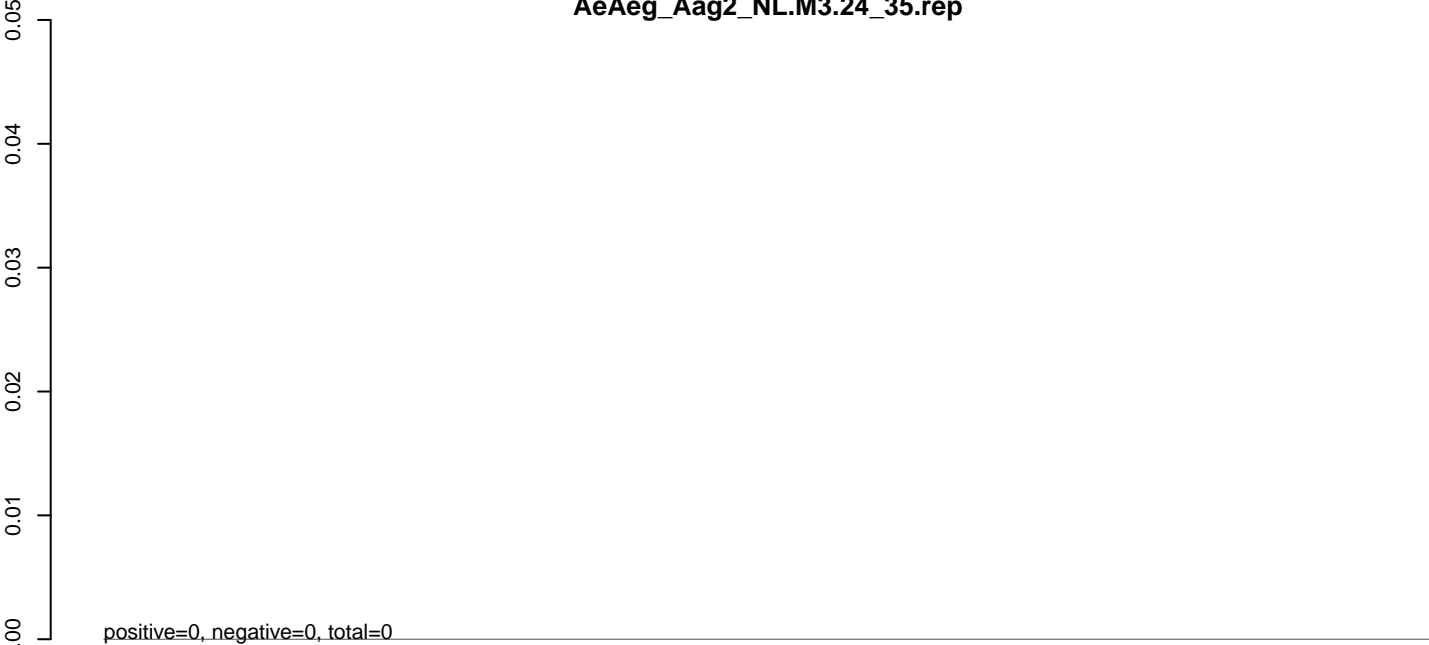
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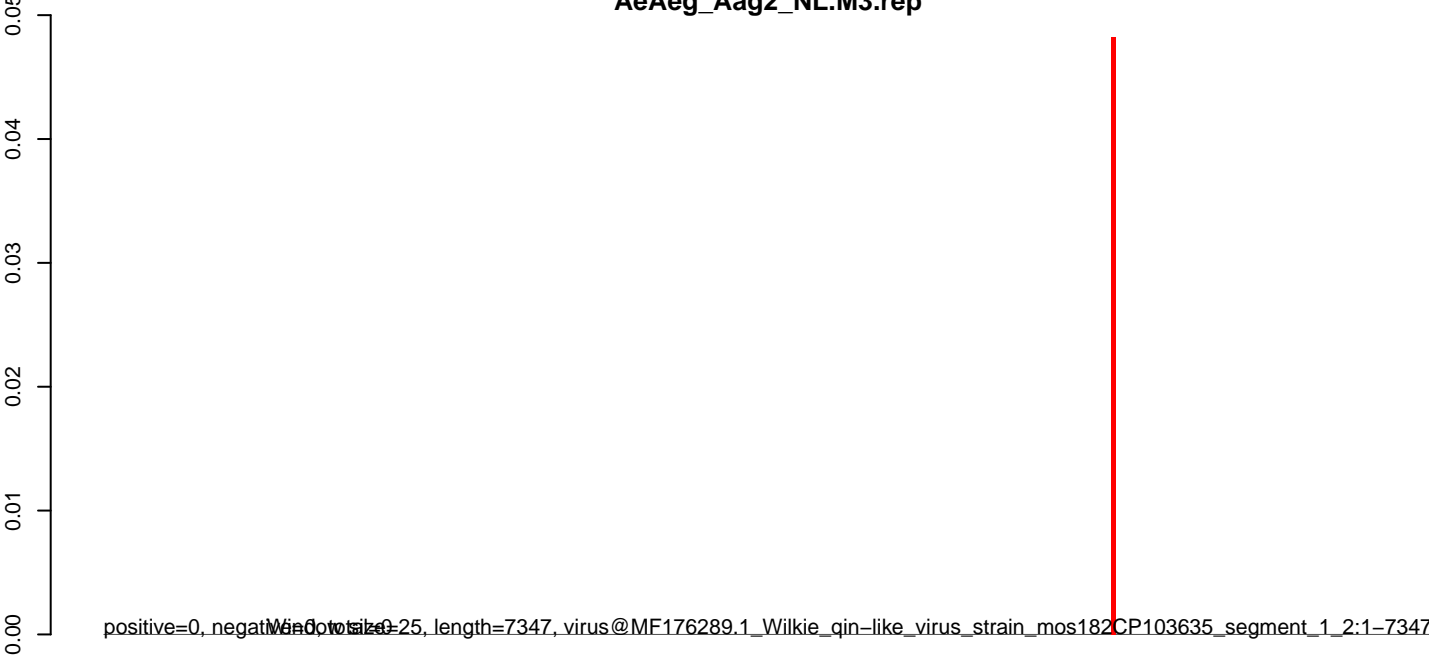
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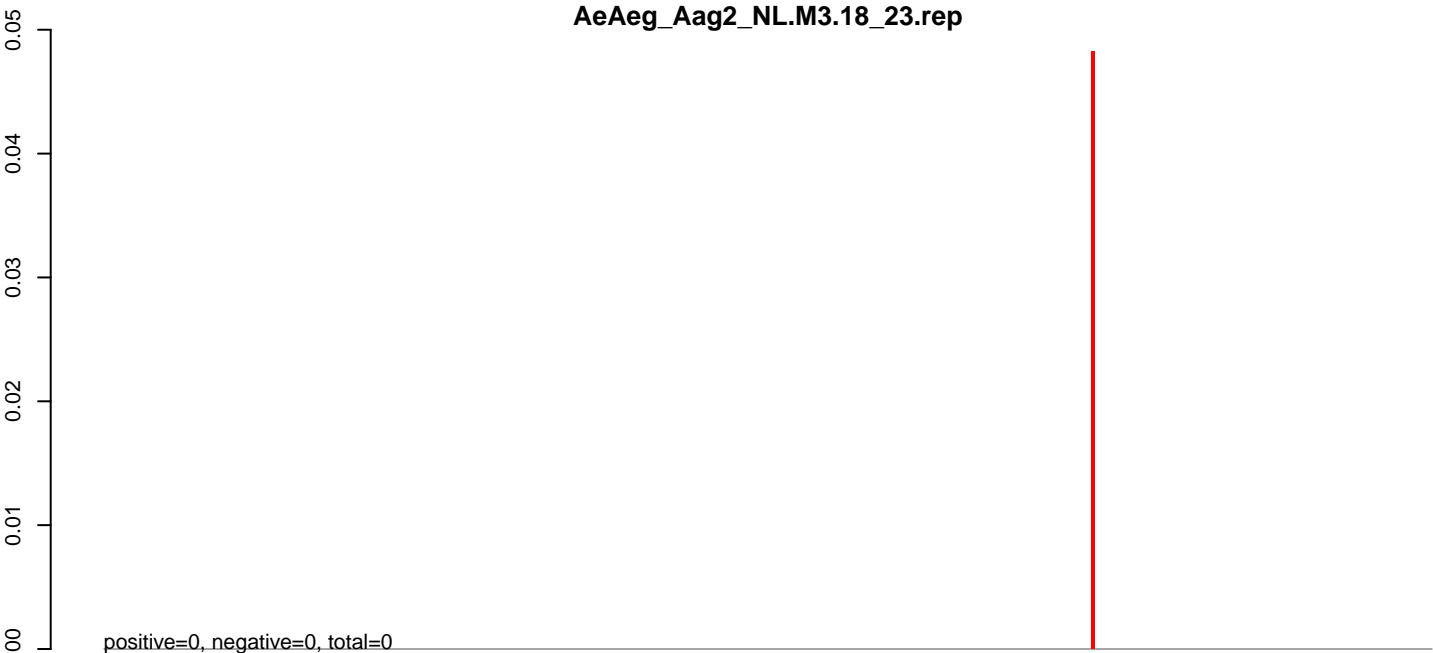
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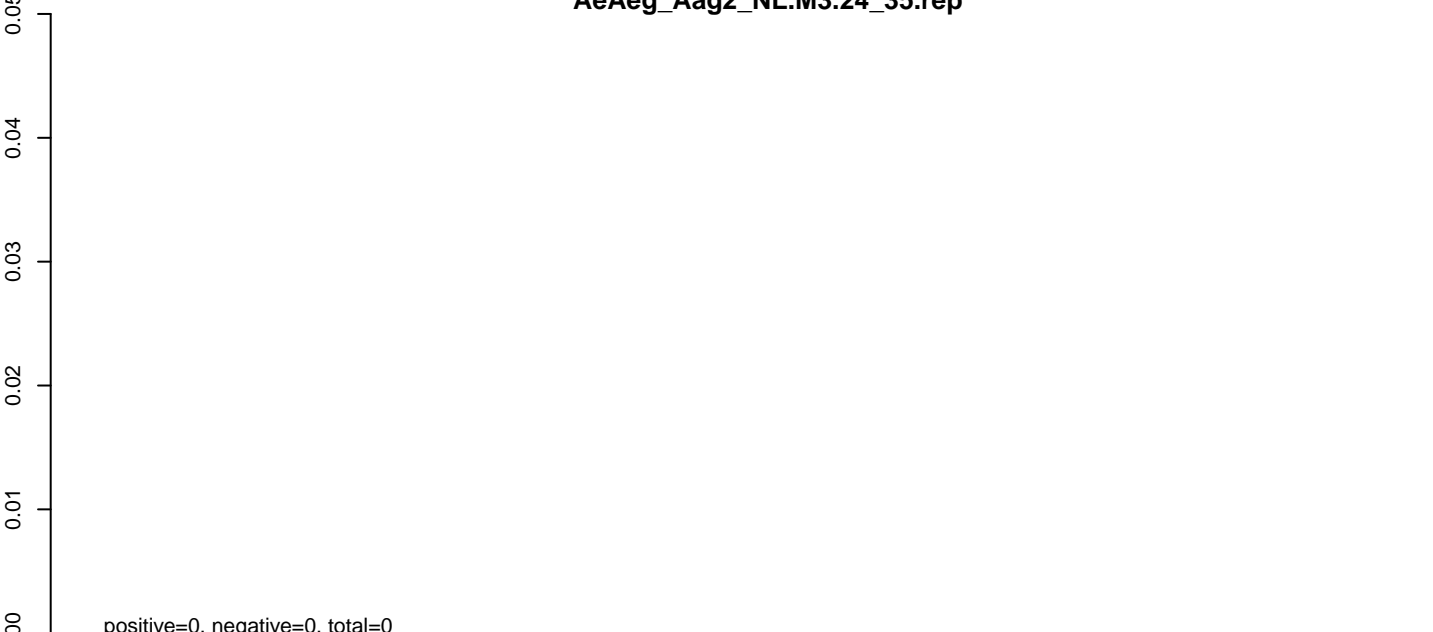
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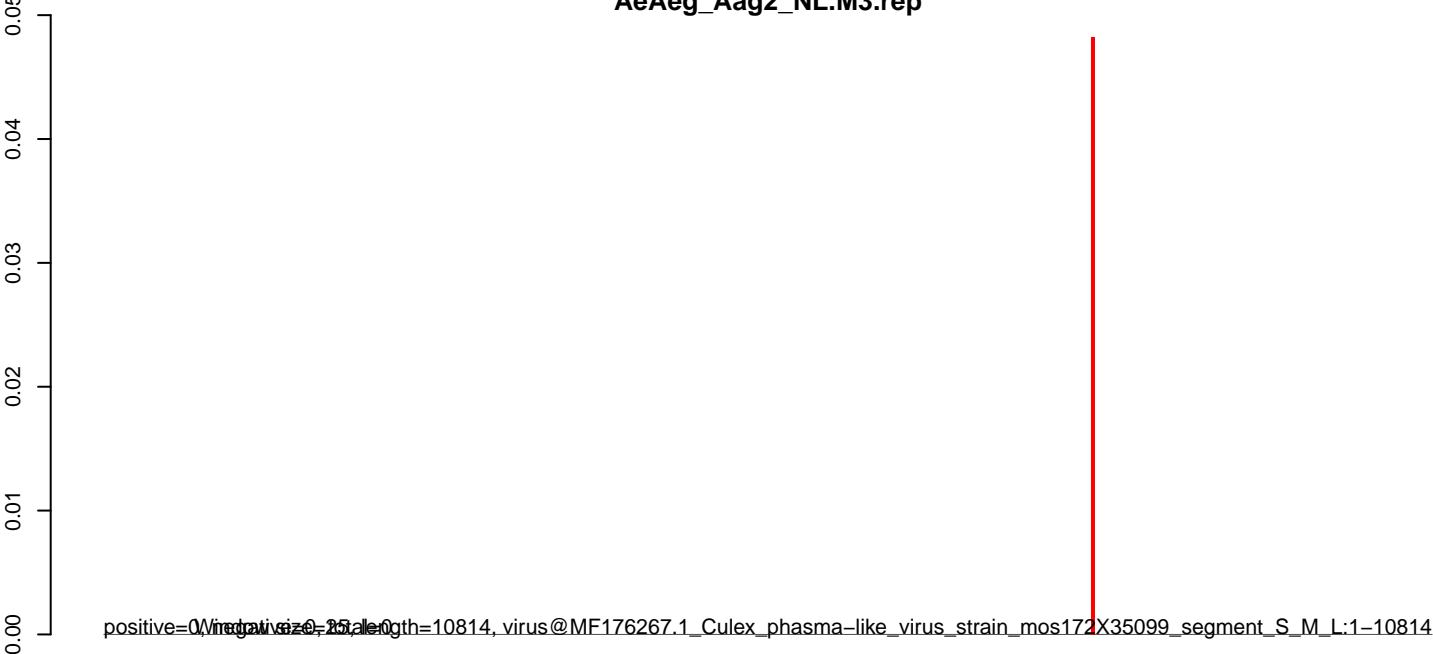
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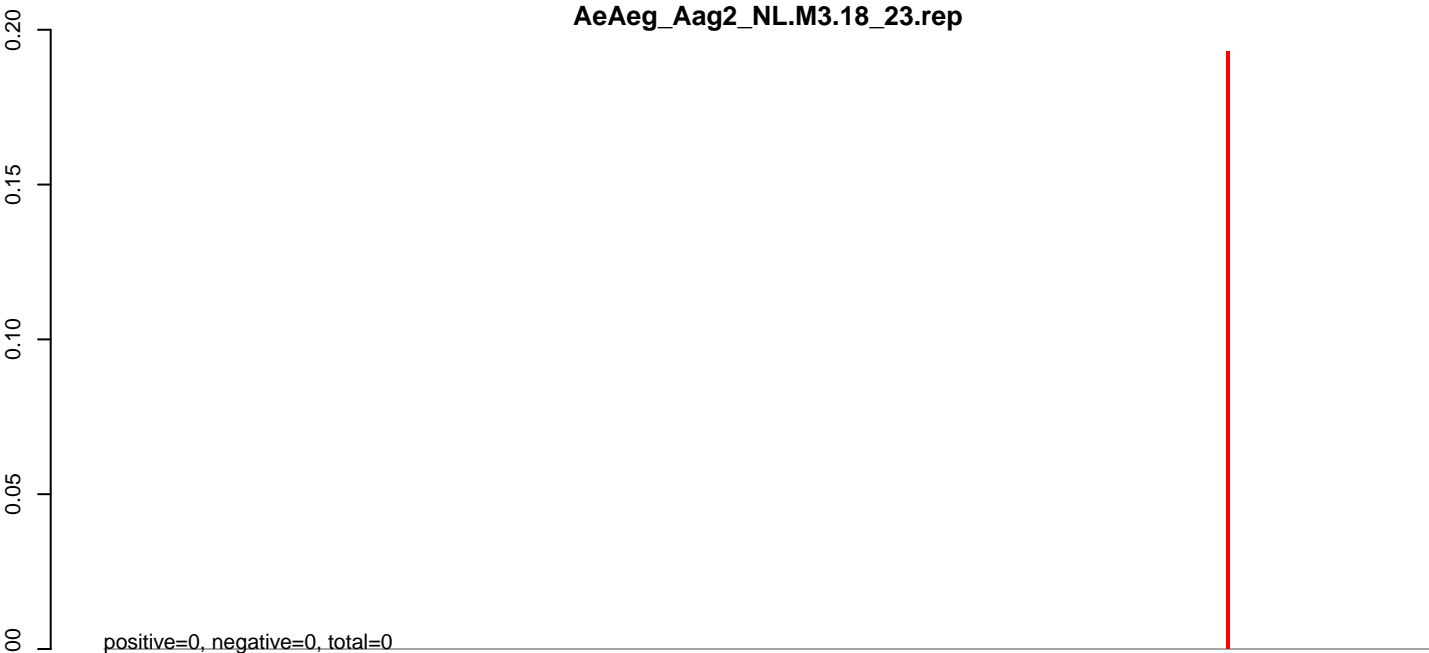


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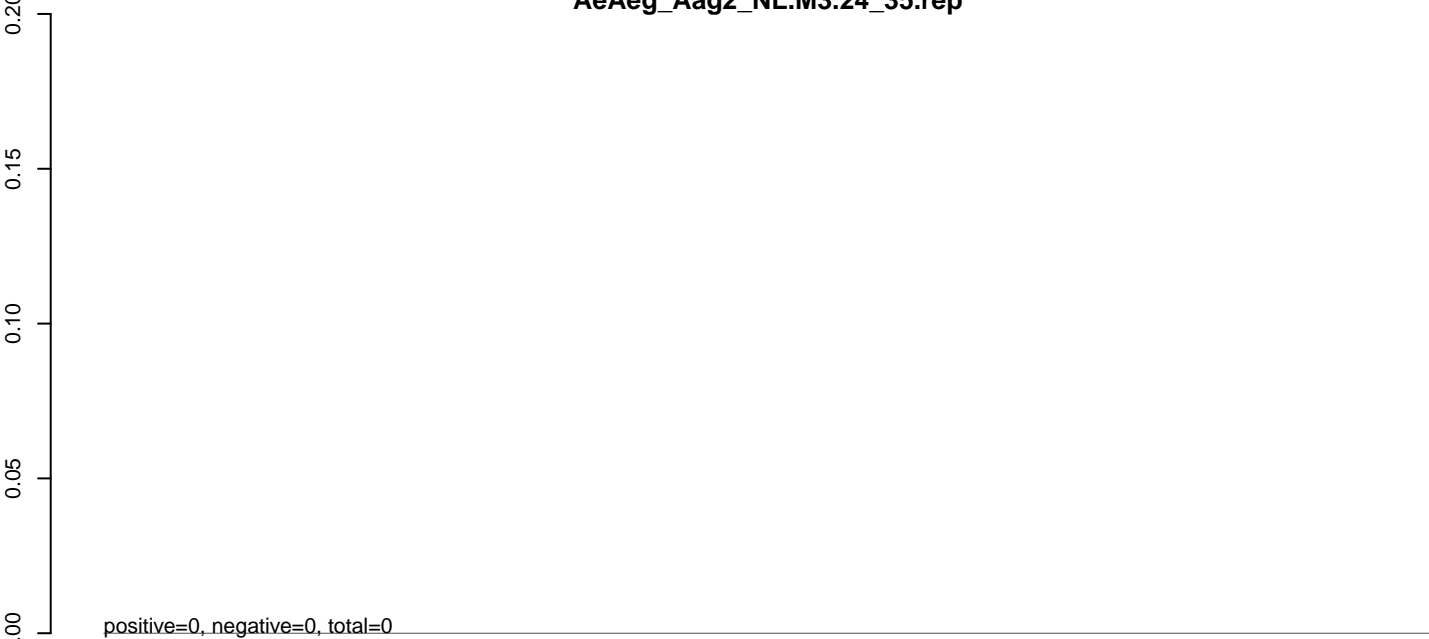


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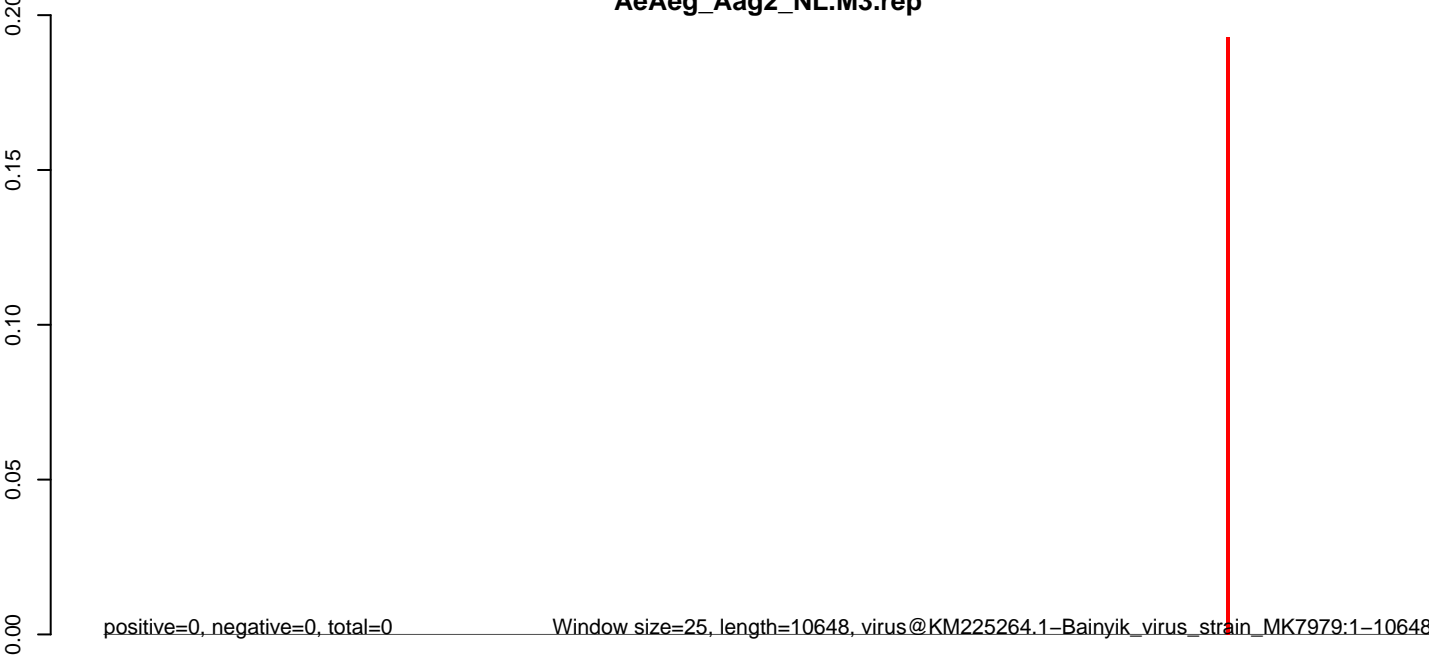
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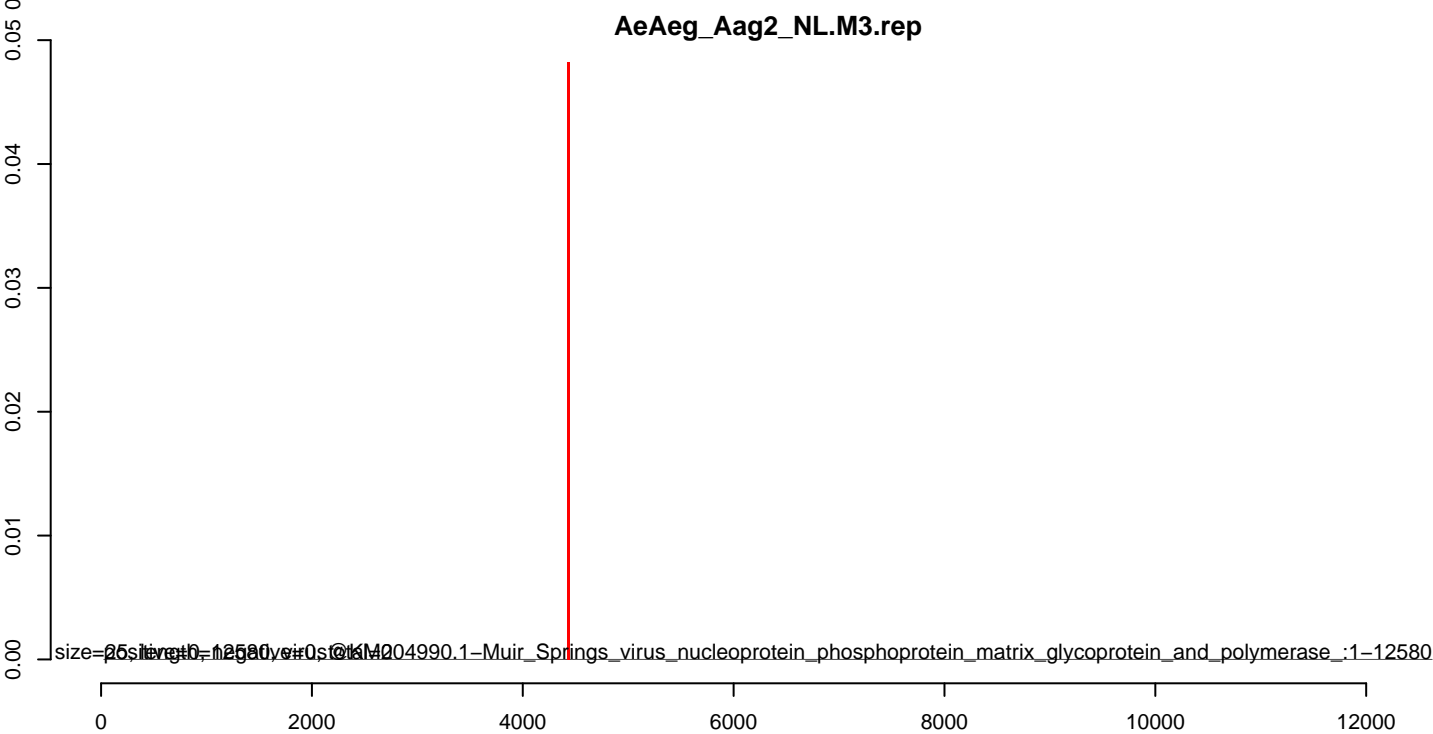
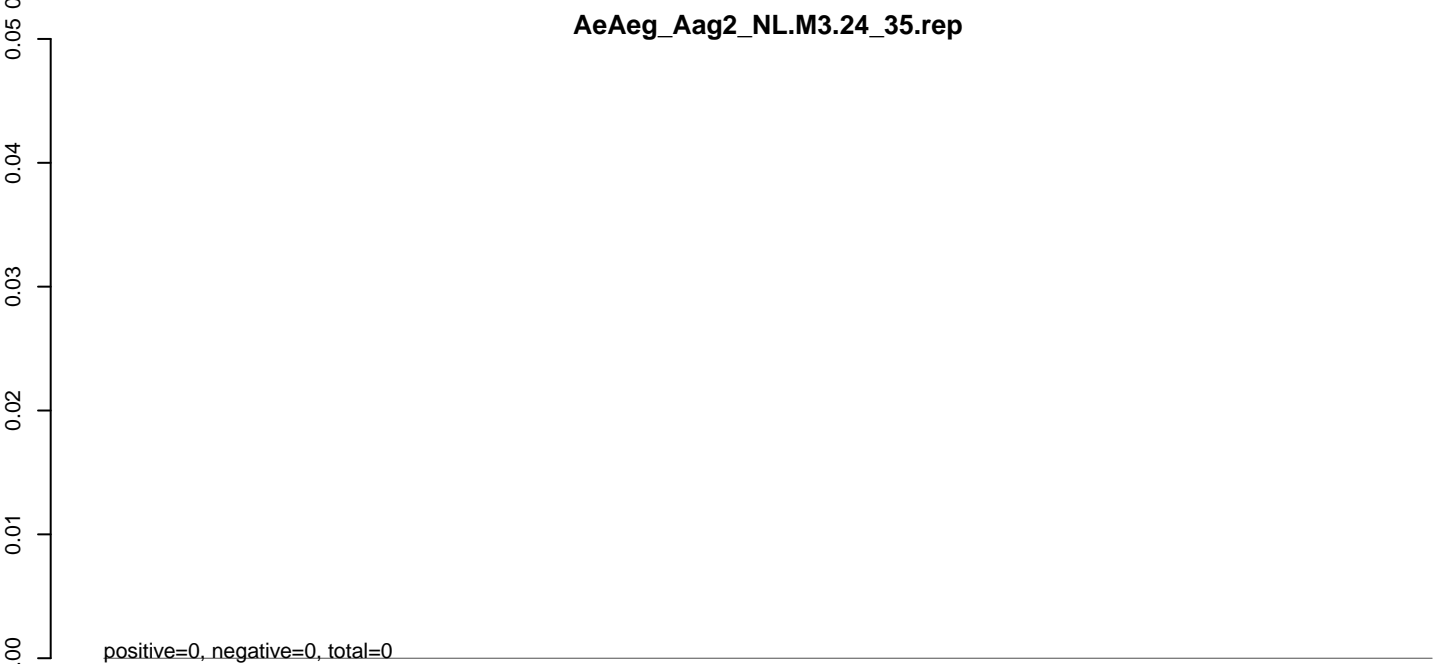
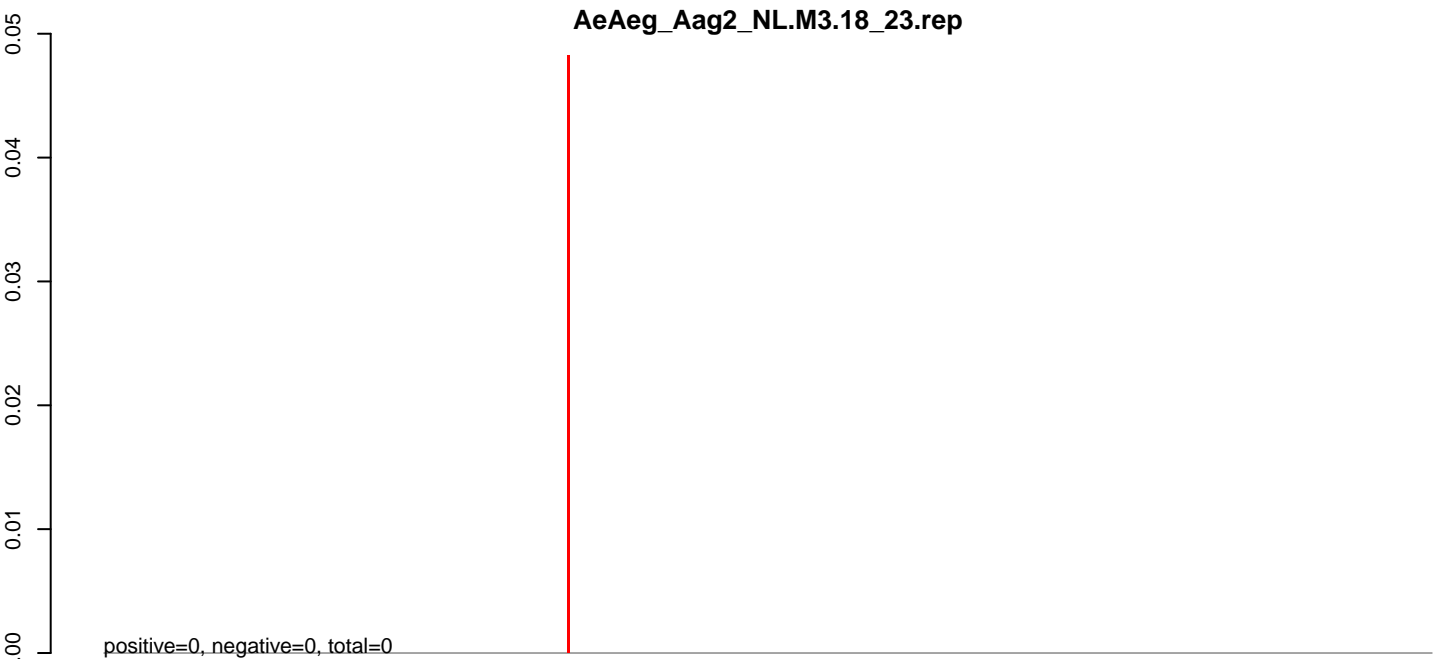
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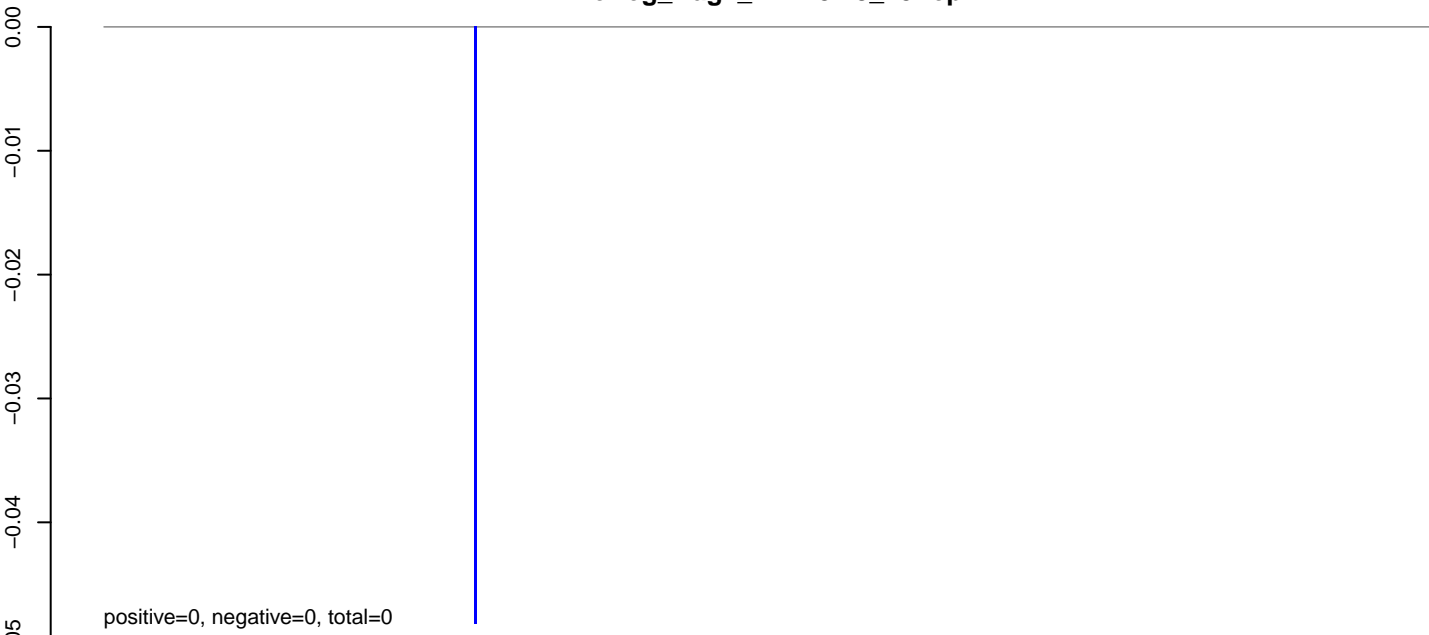
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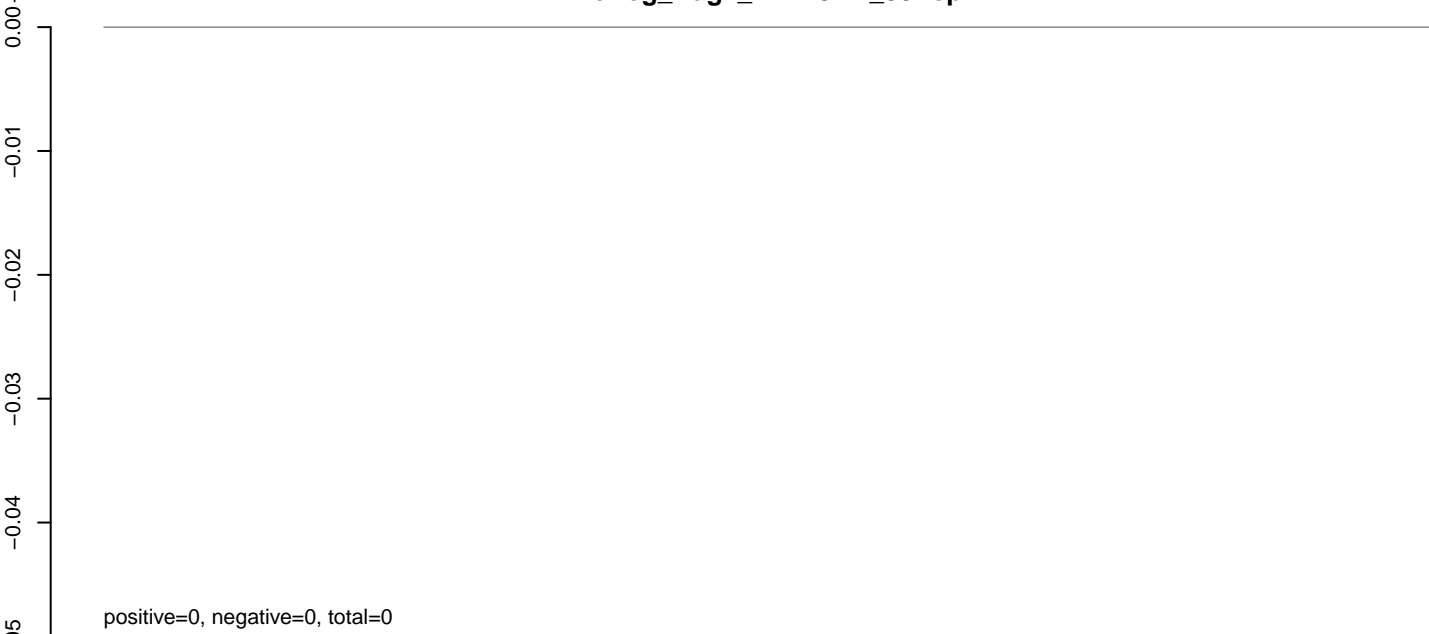
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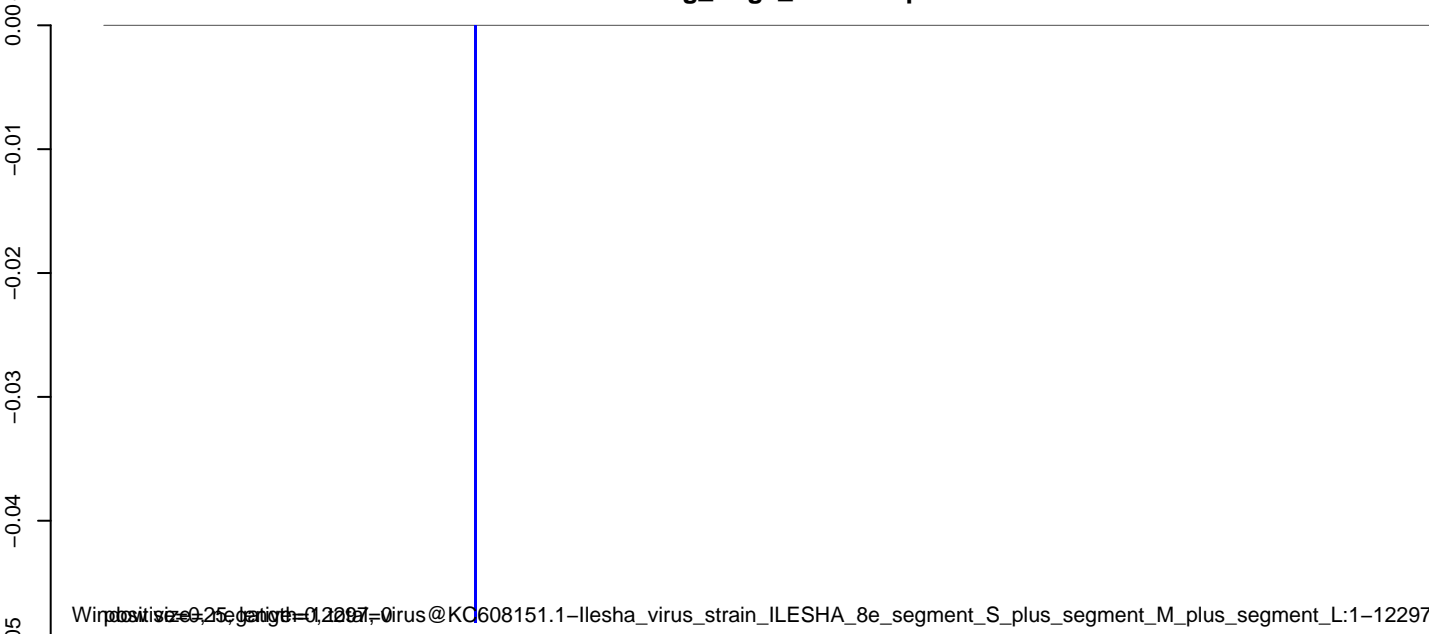
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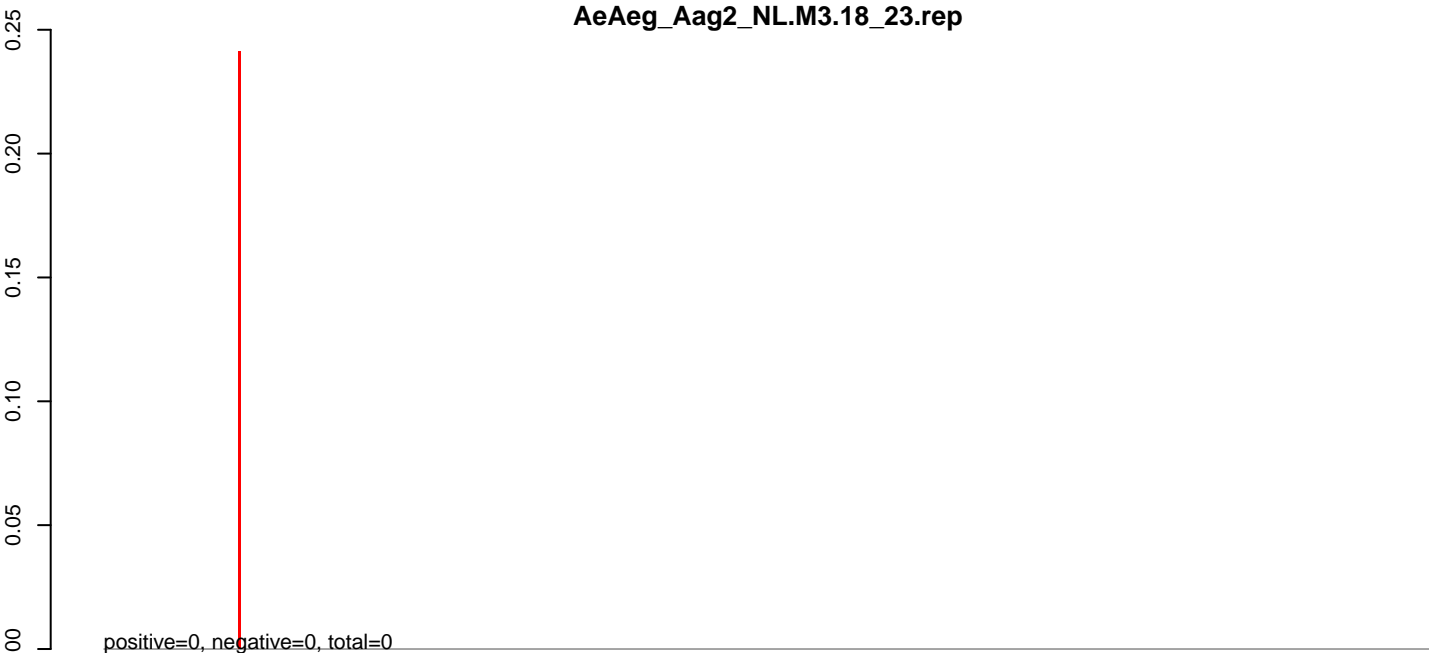
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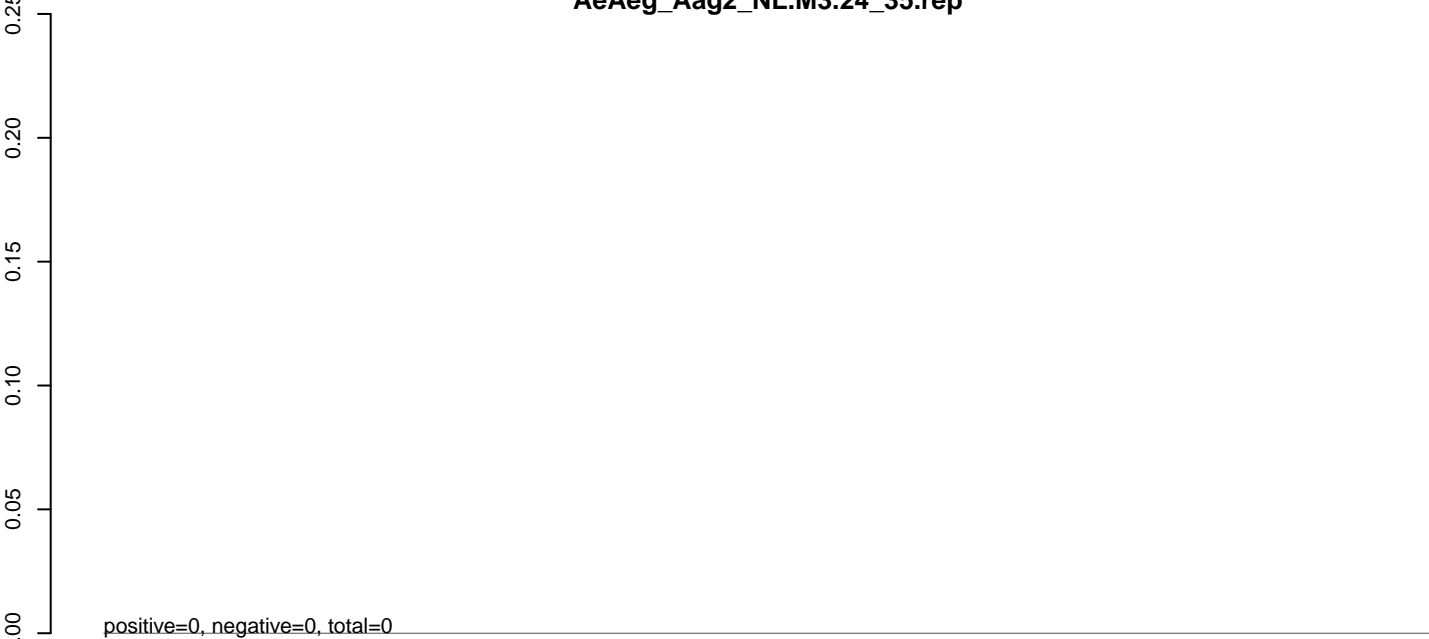
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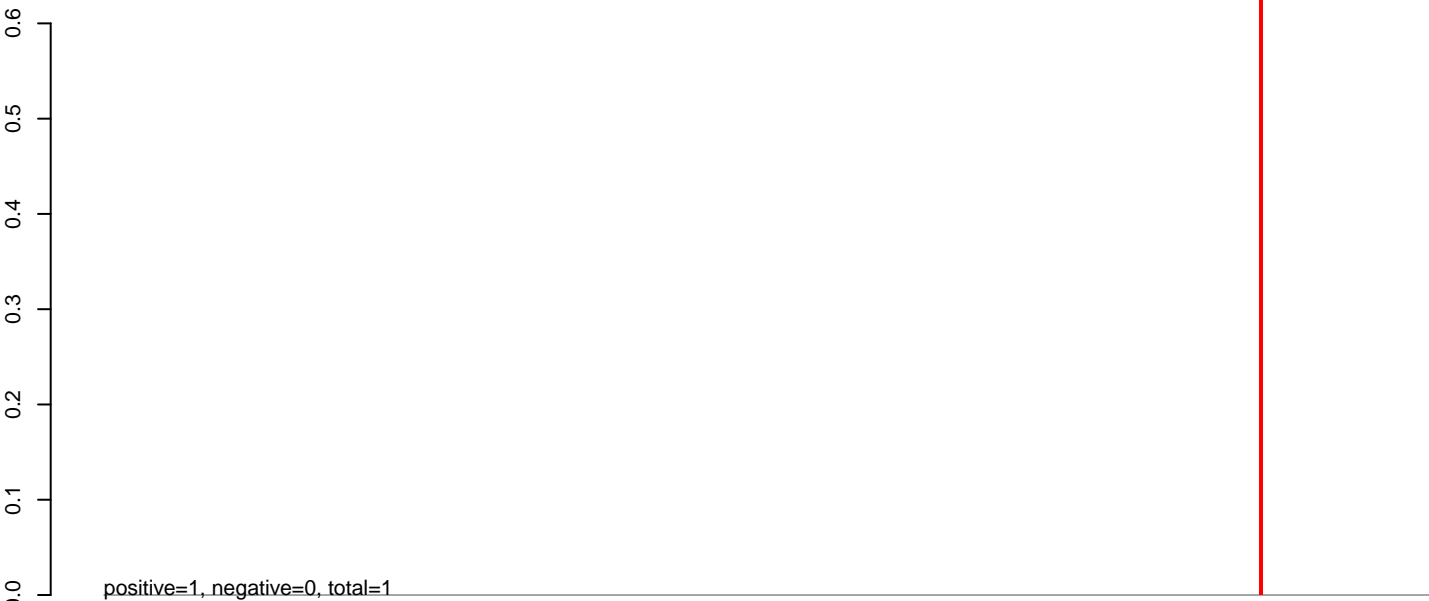
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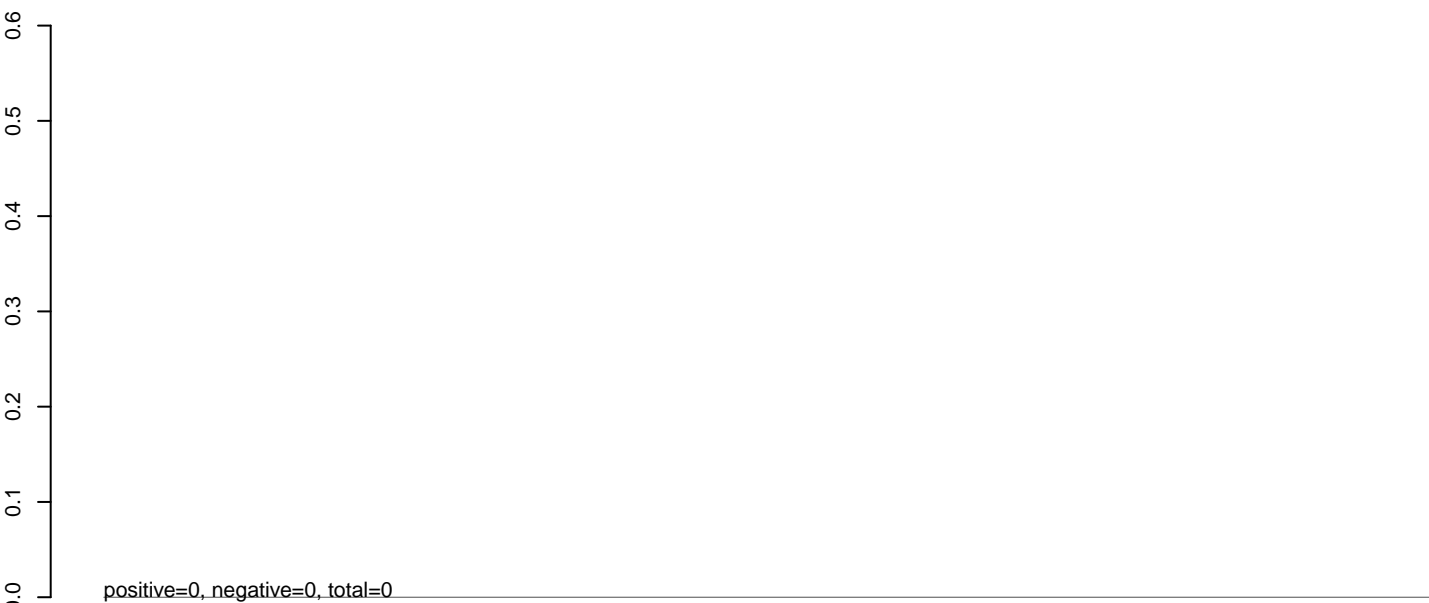
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0 2000 4000 6000 8000 10000 12000

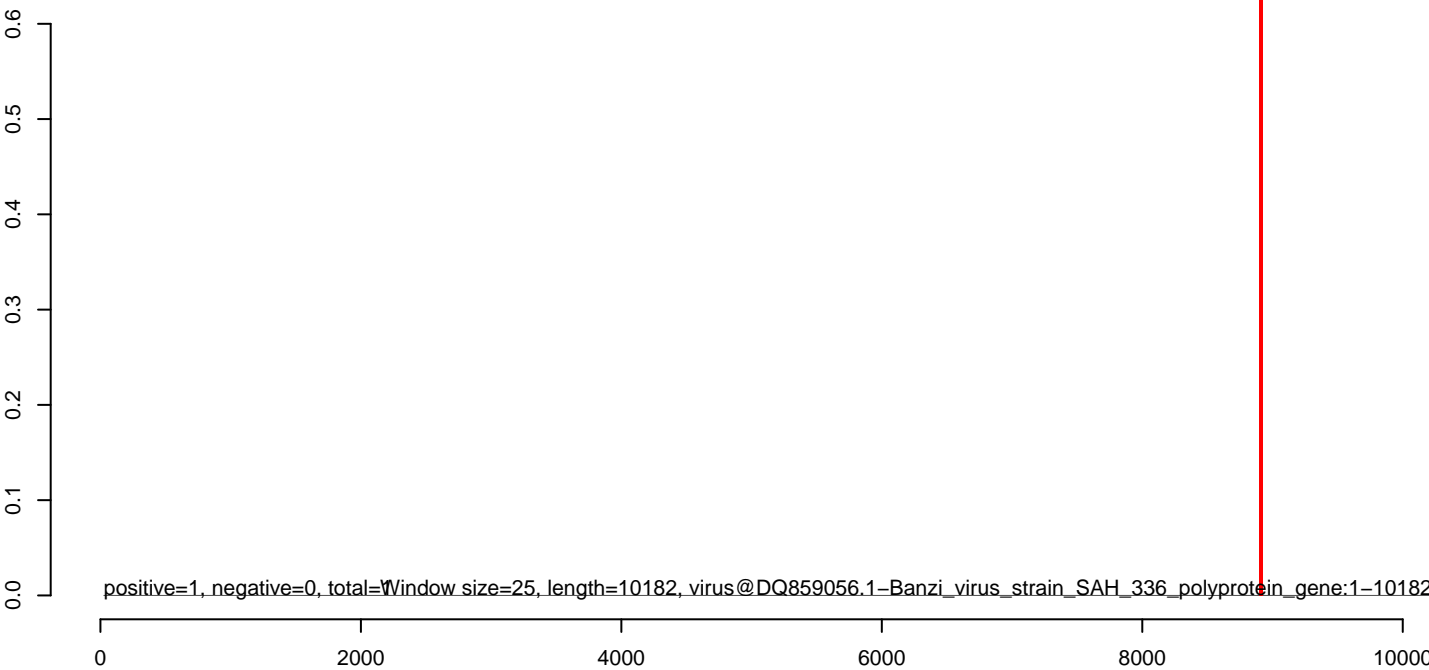
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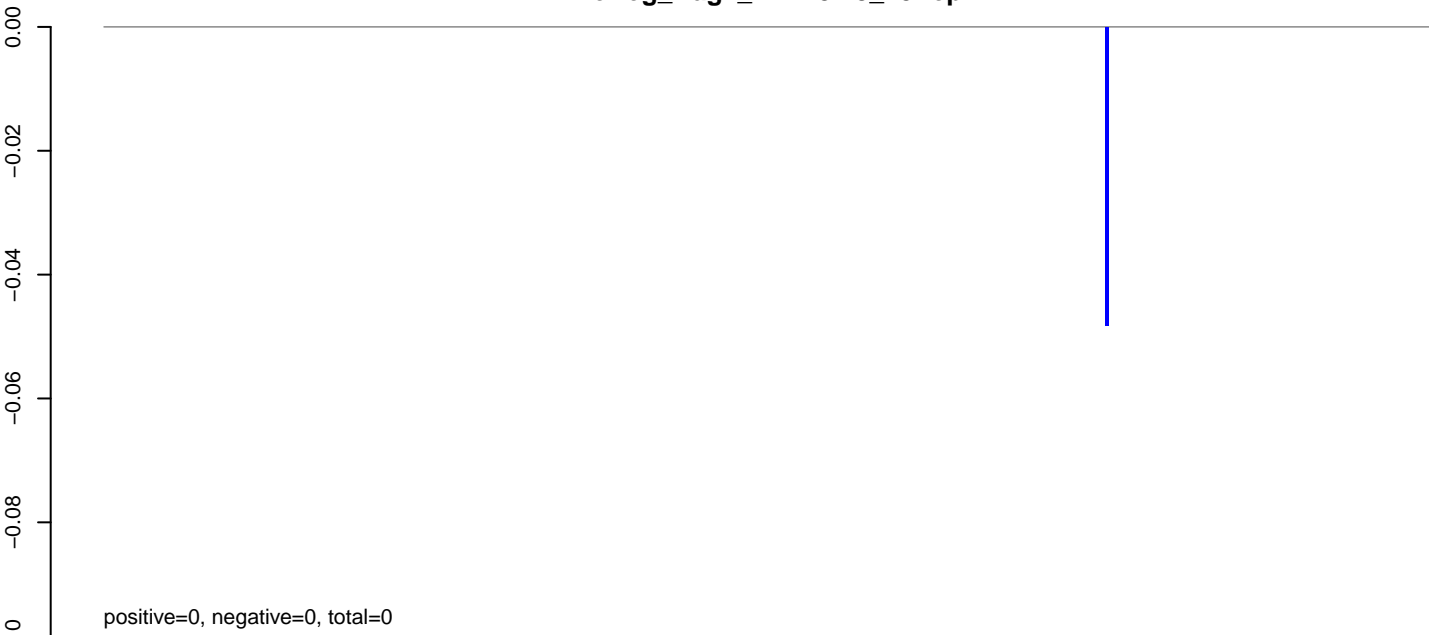
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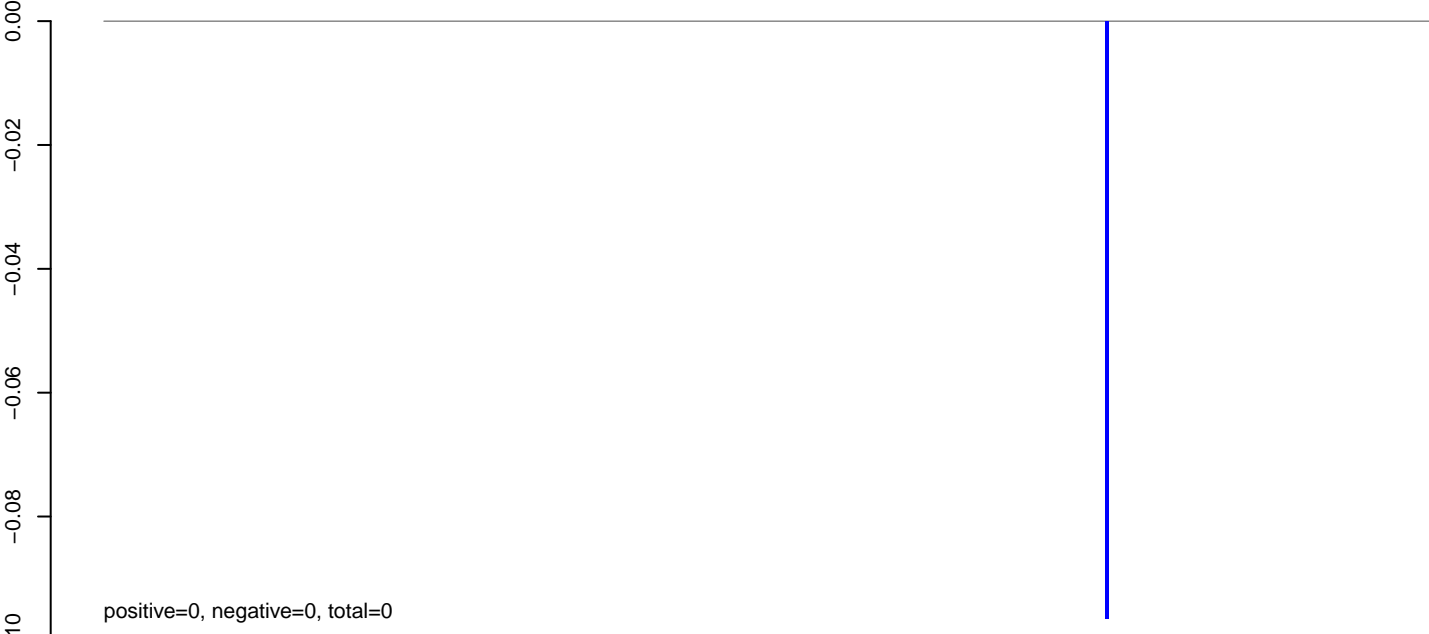
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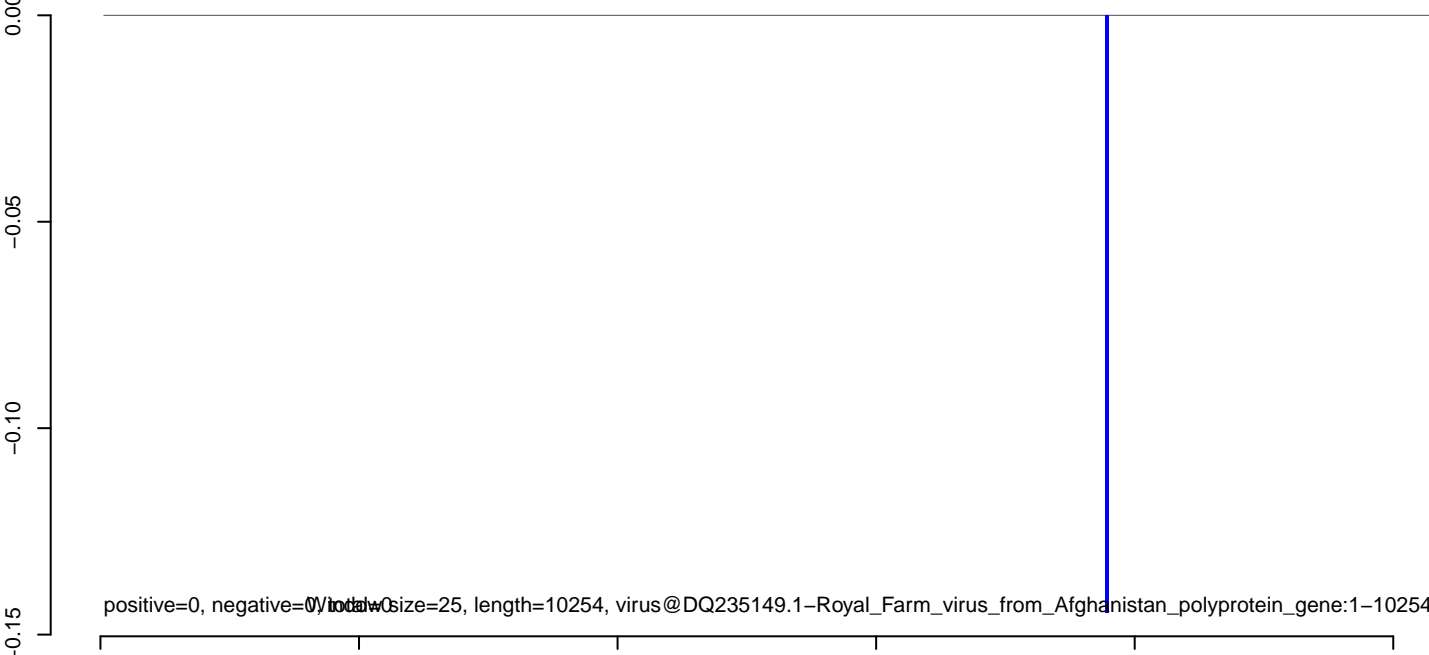
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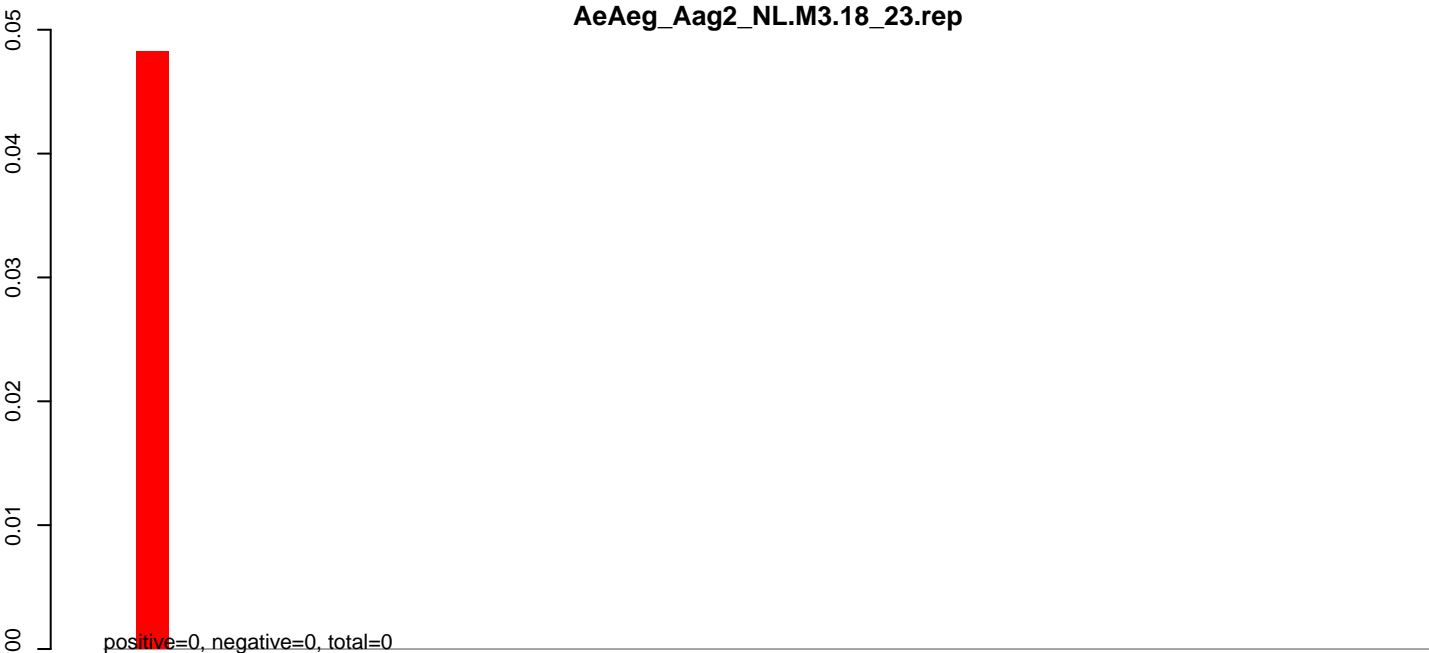
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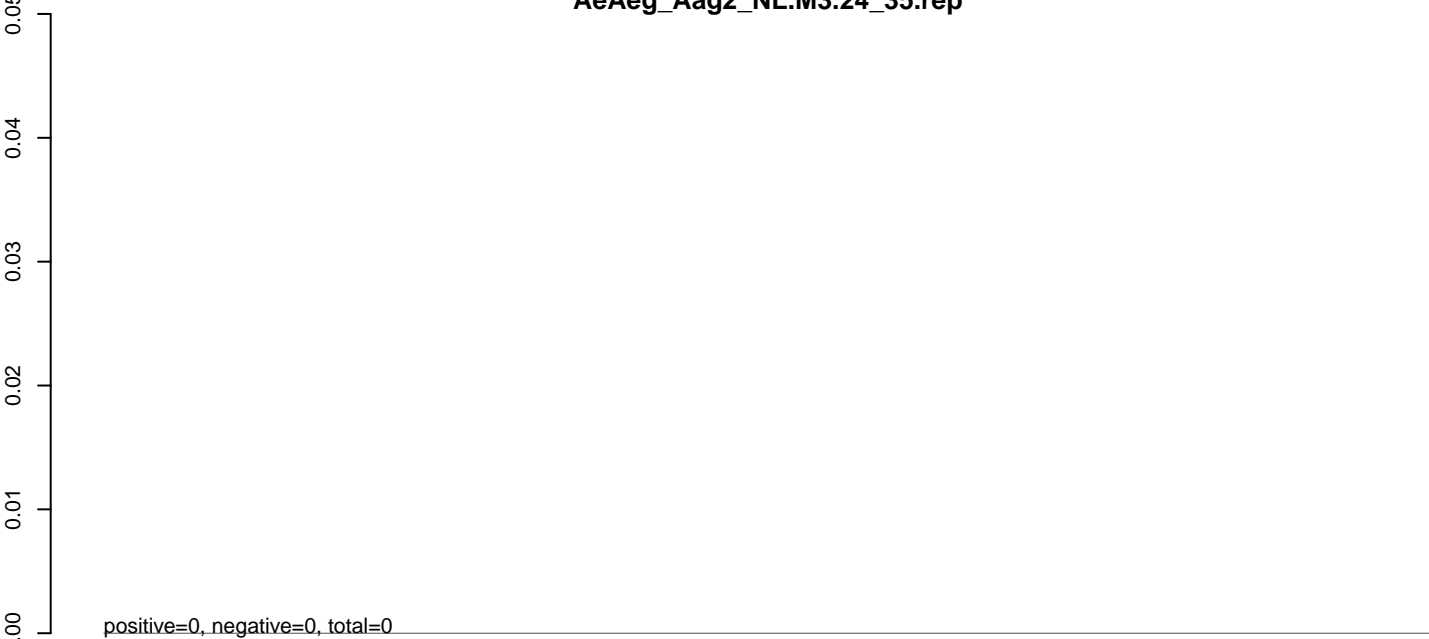
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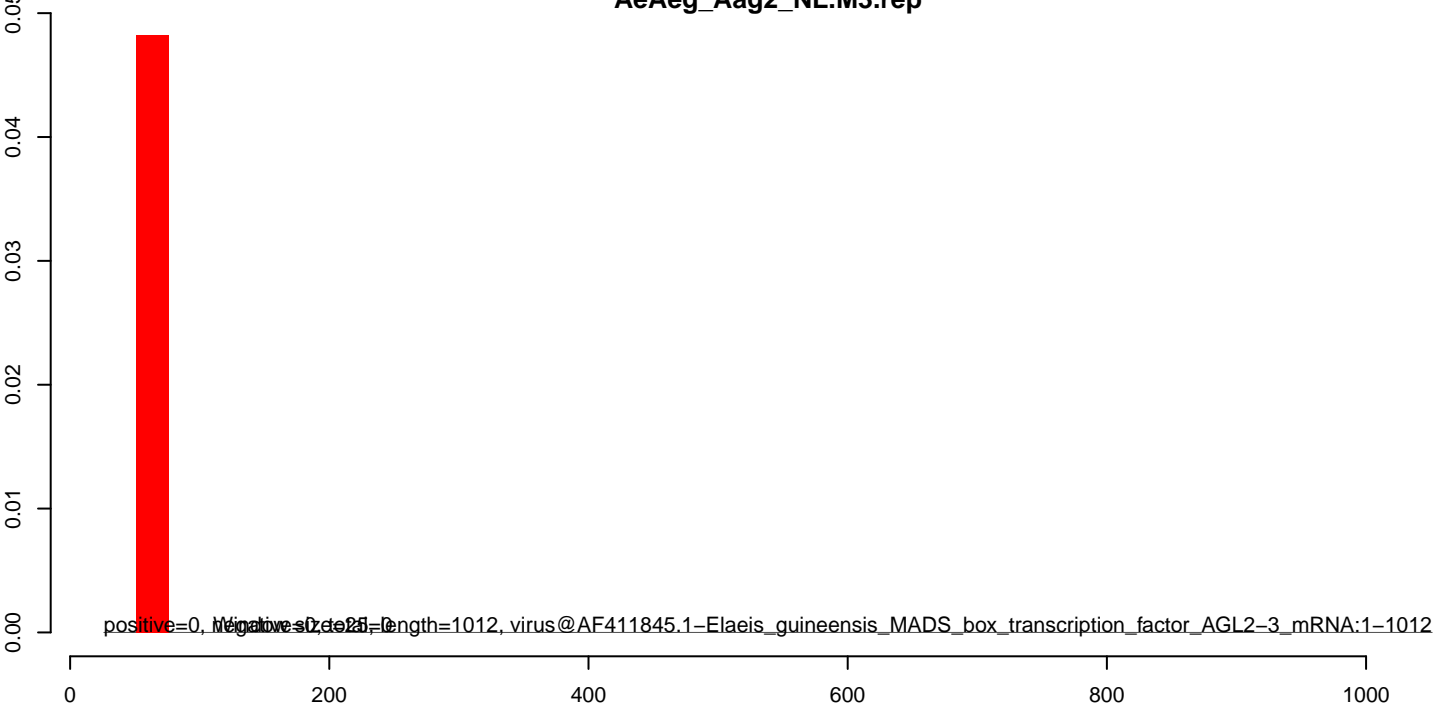
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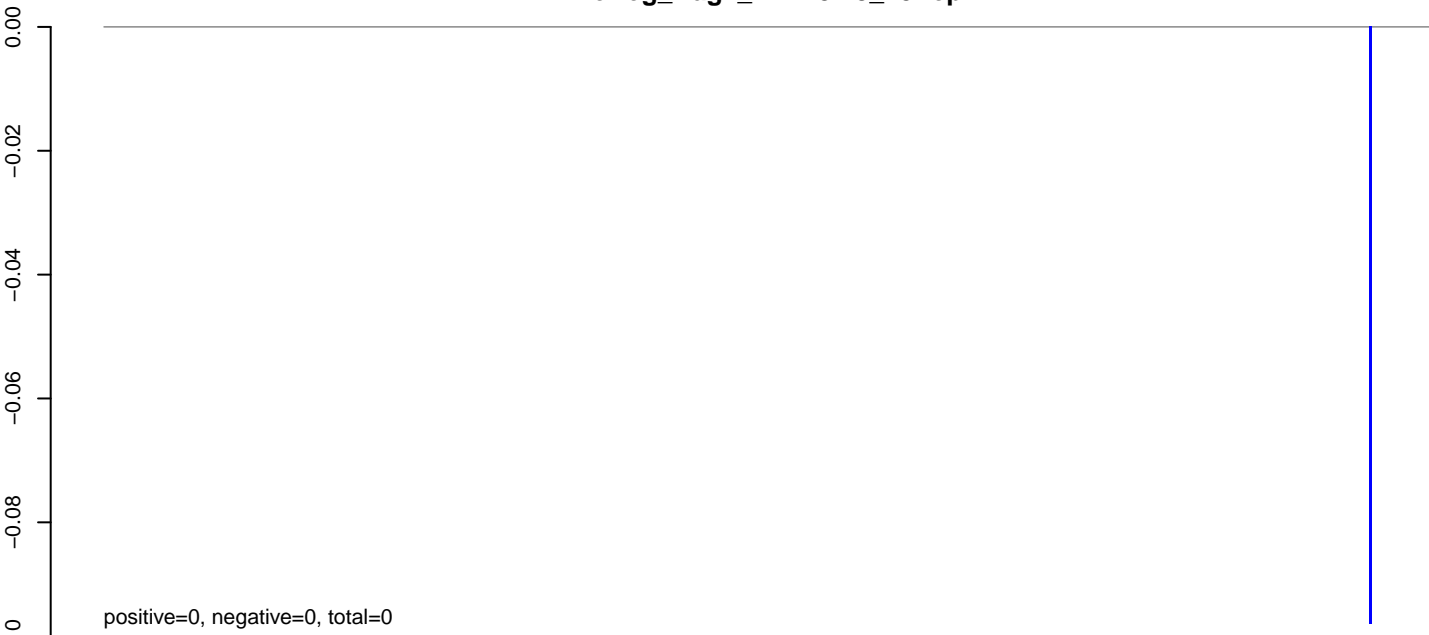
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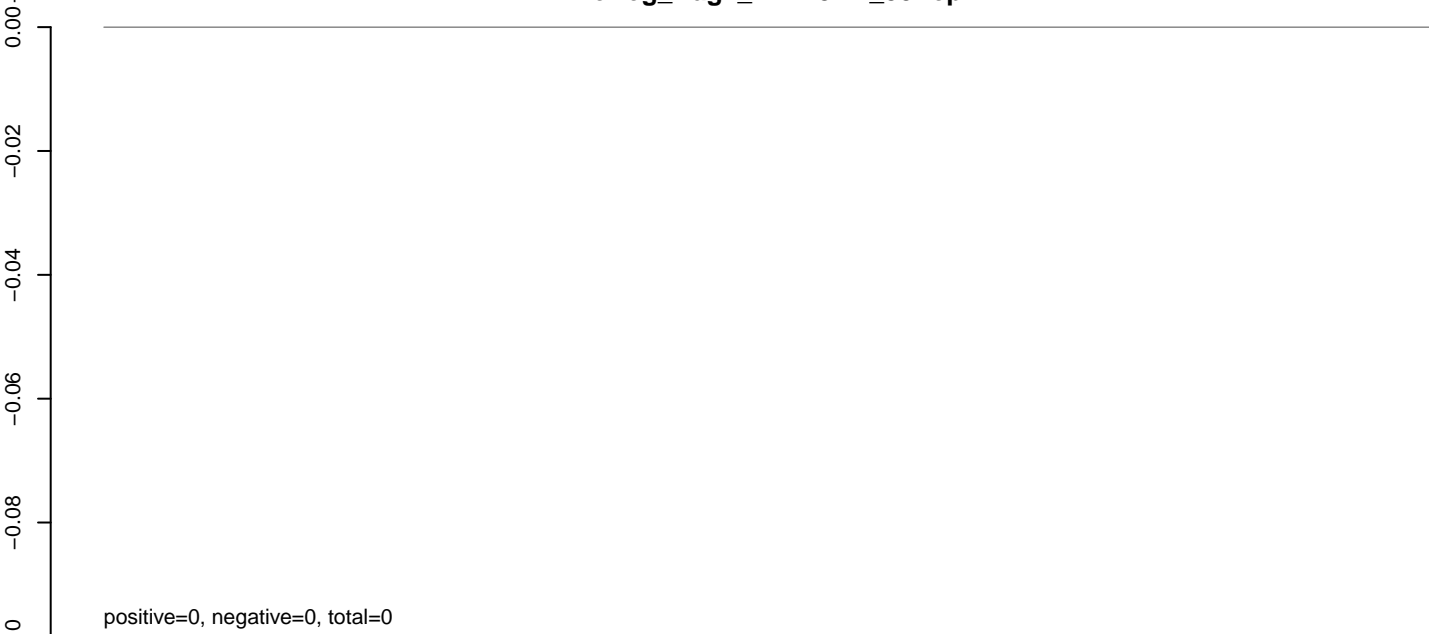
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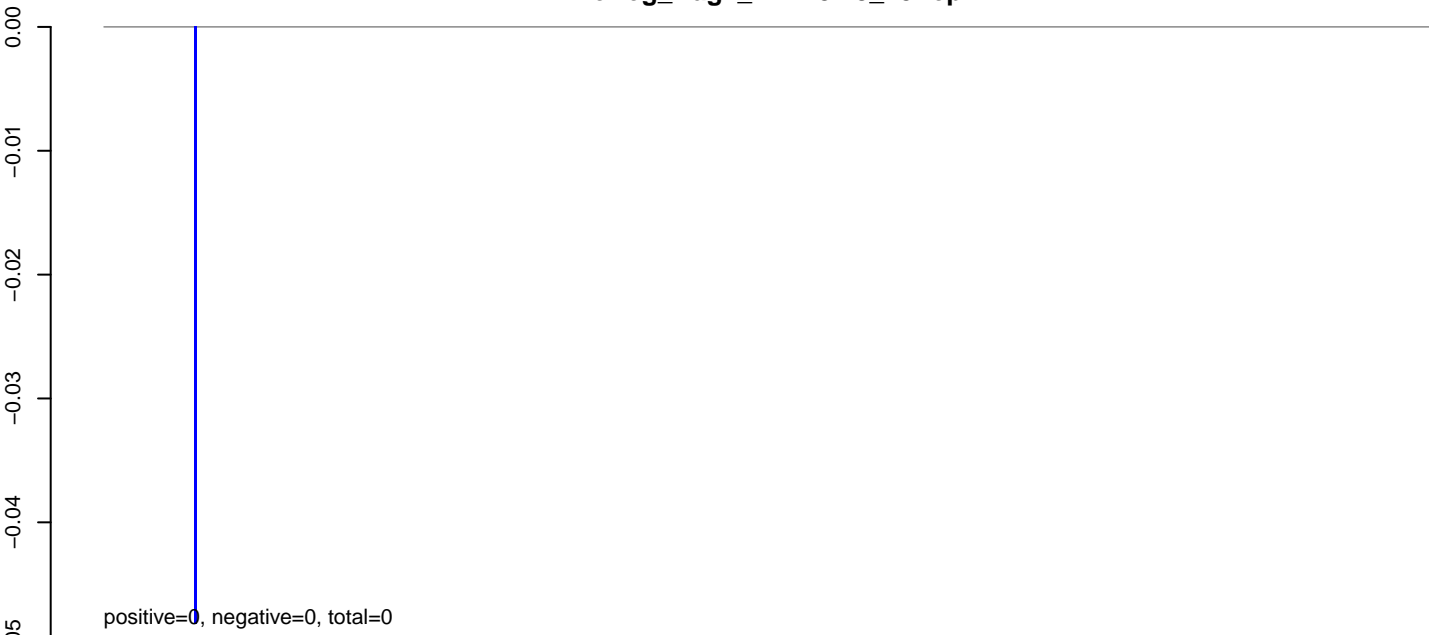
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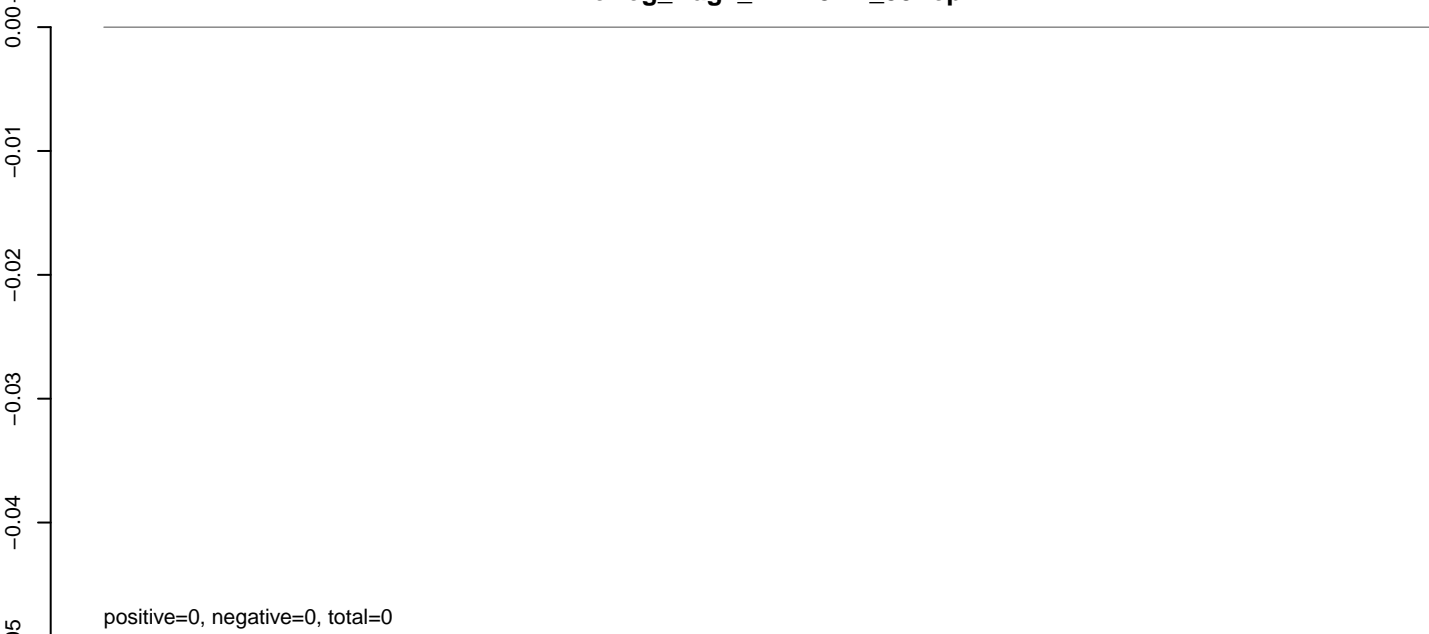
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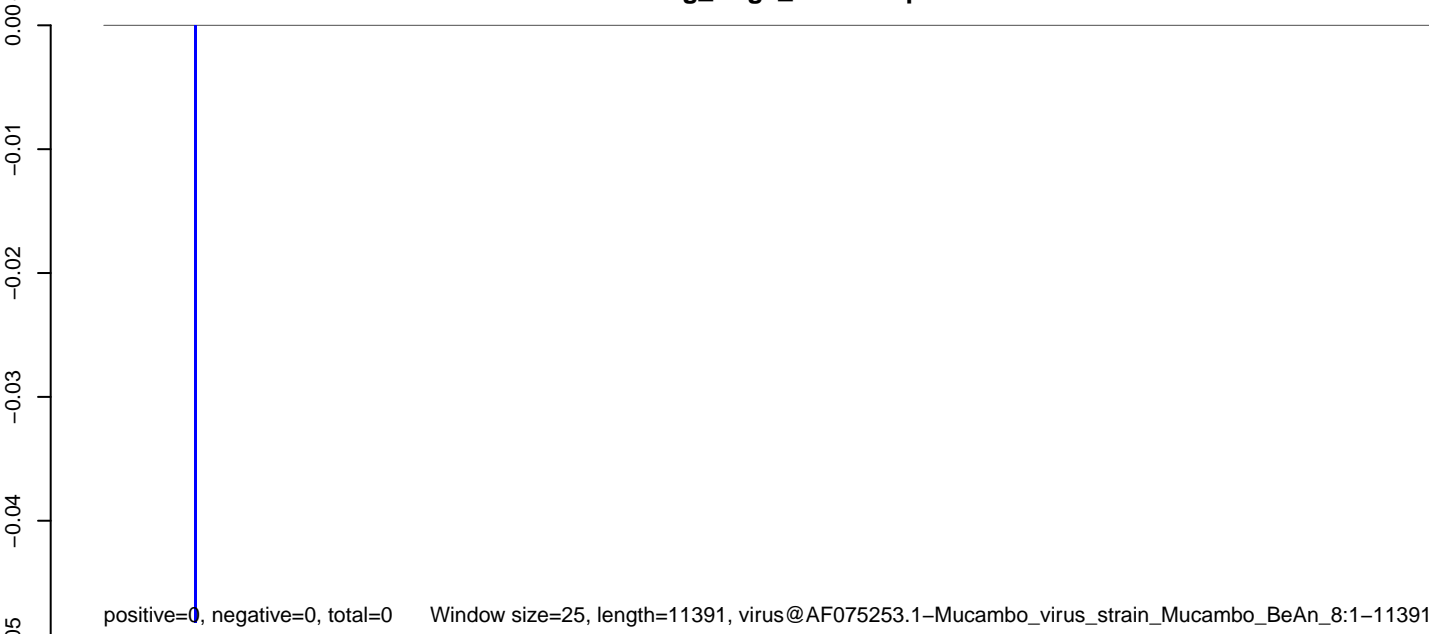
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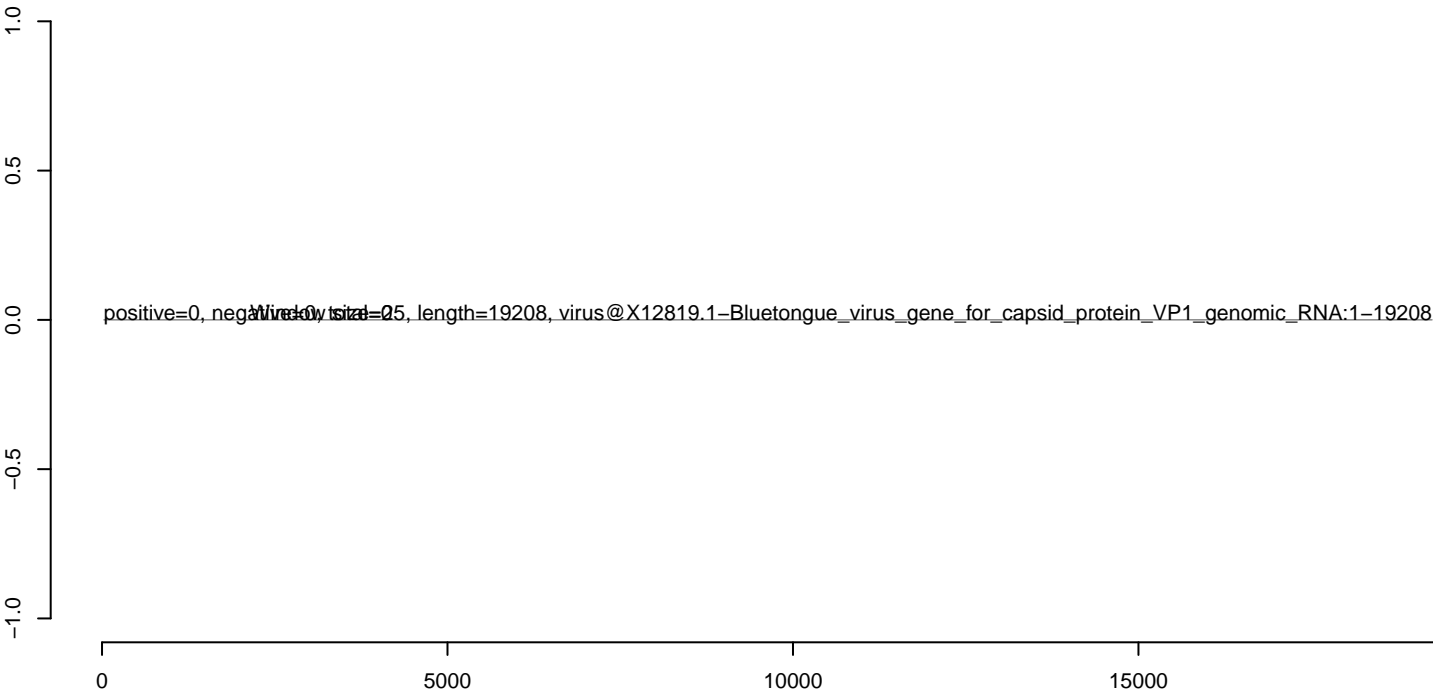
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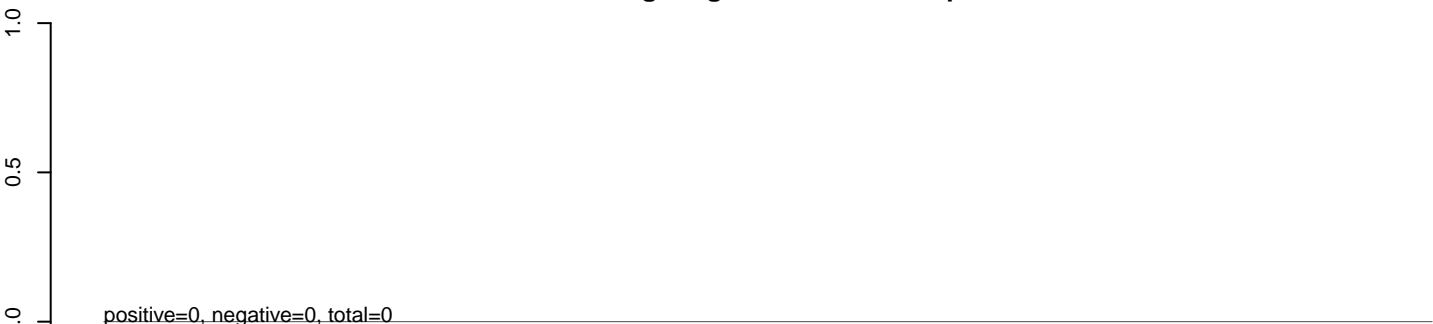
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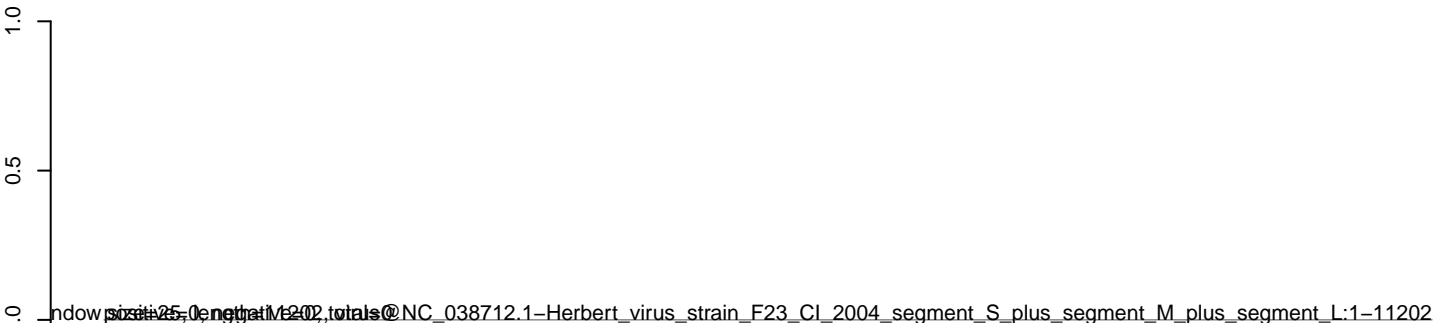
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AeAeg_Aag2_NL.M3.rep



0 2000 4000 6000 8000 10000

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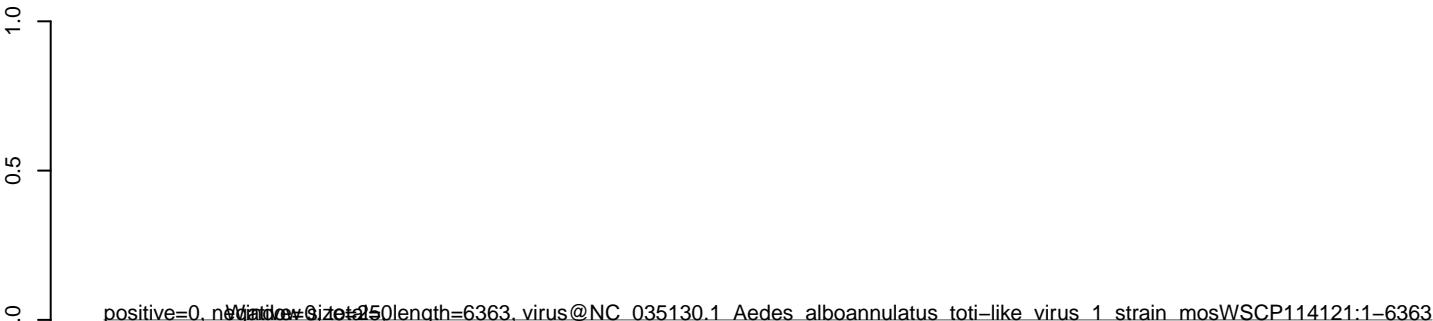
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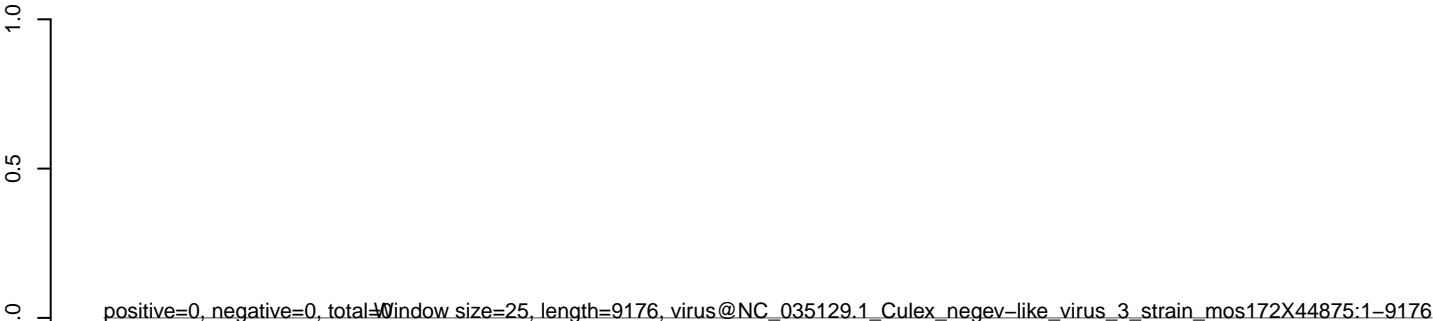
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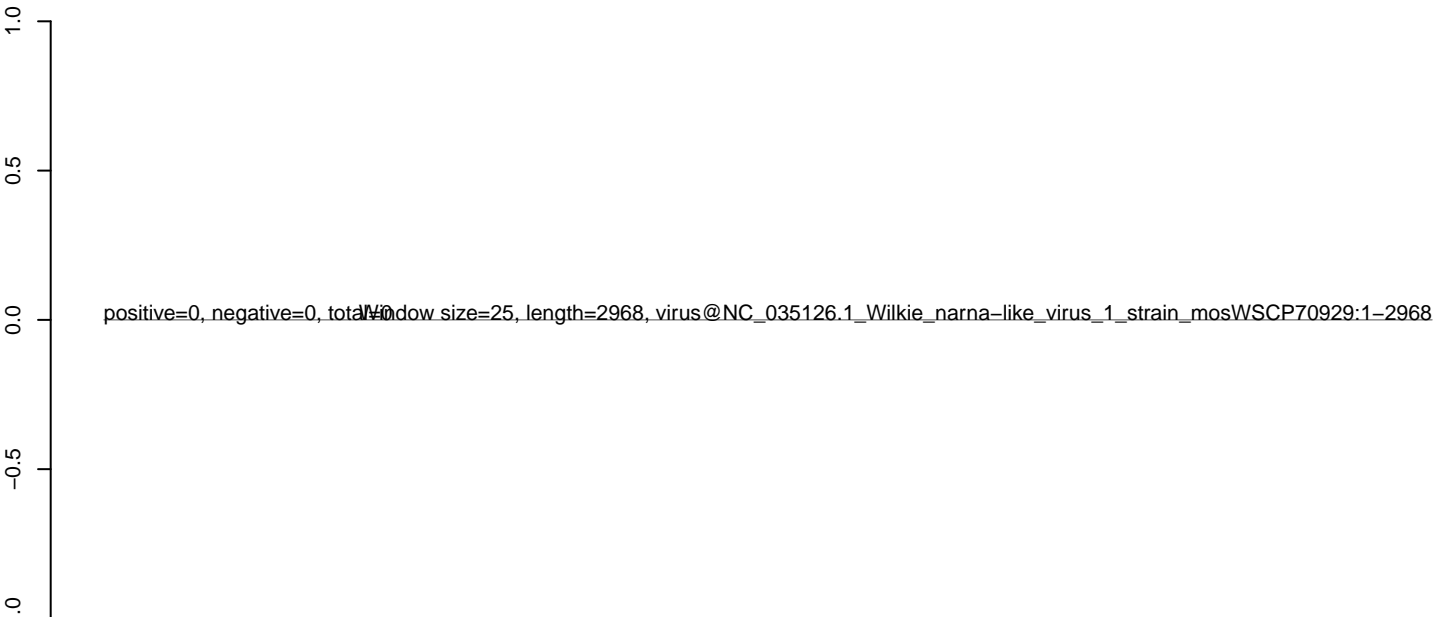
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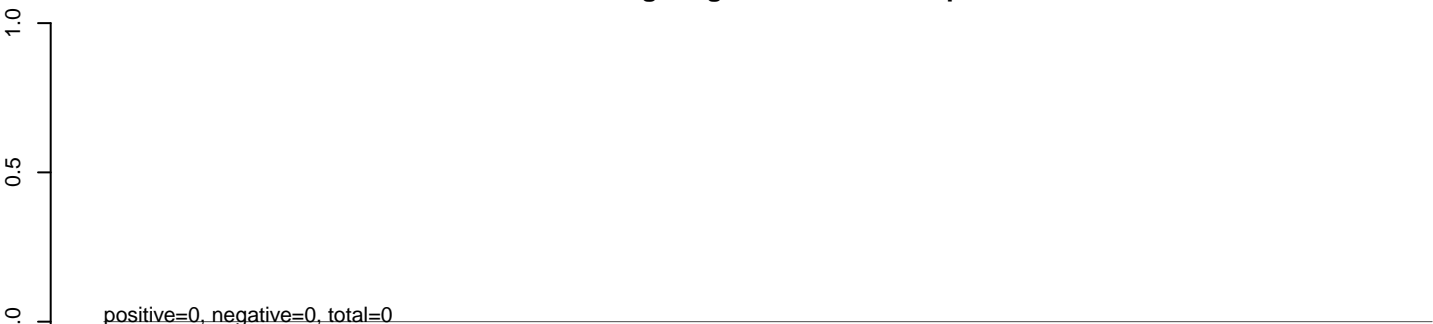
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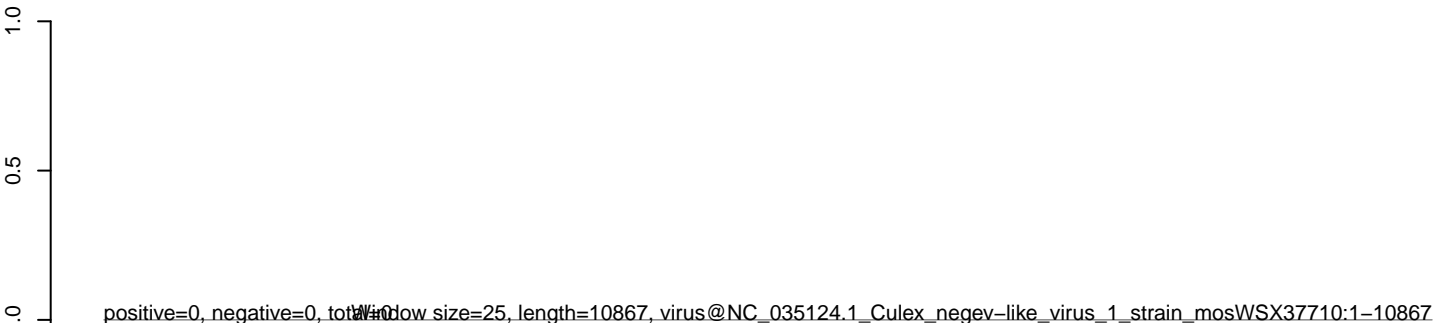
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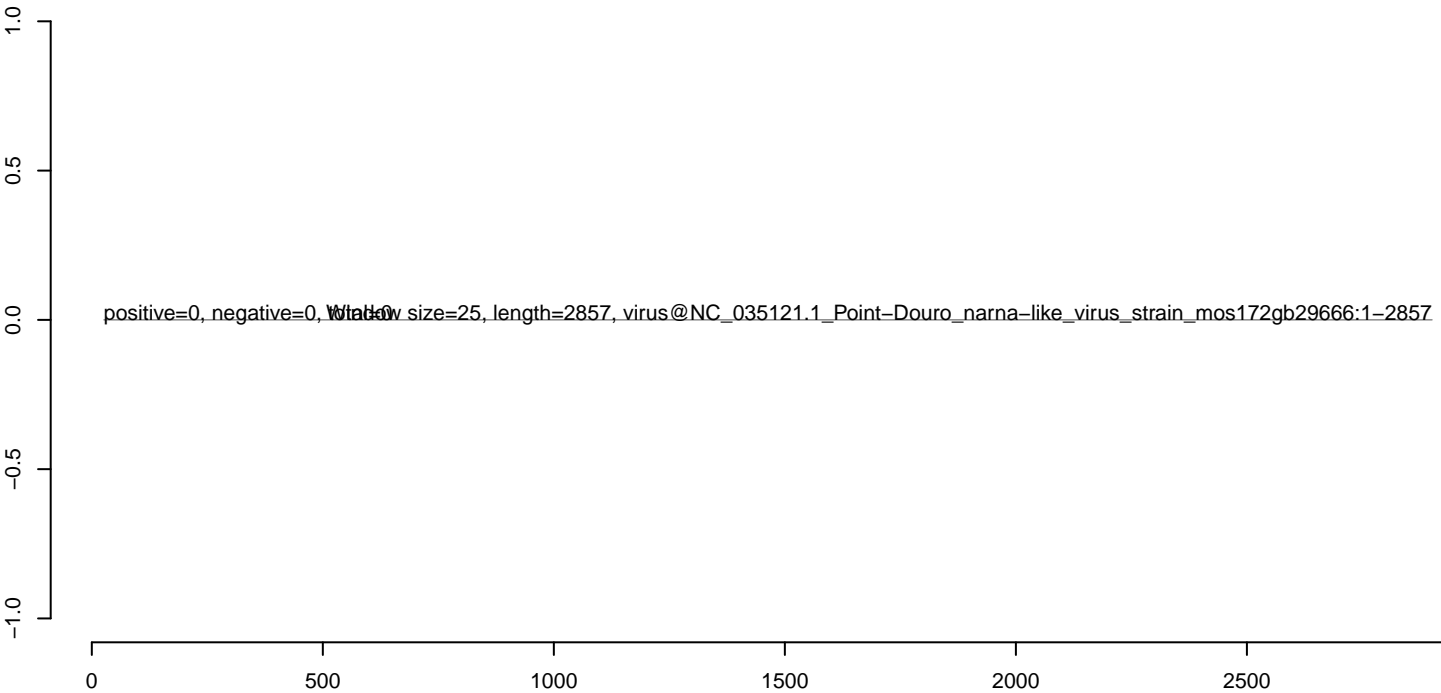
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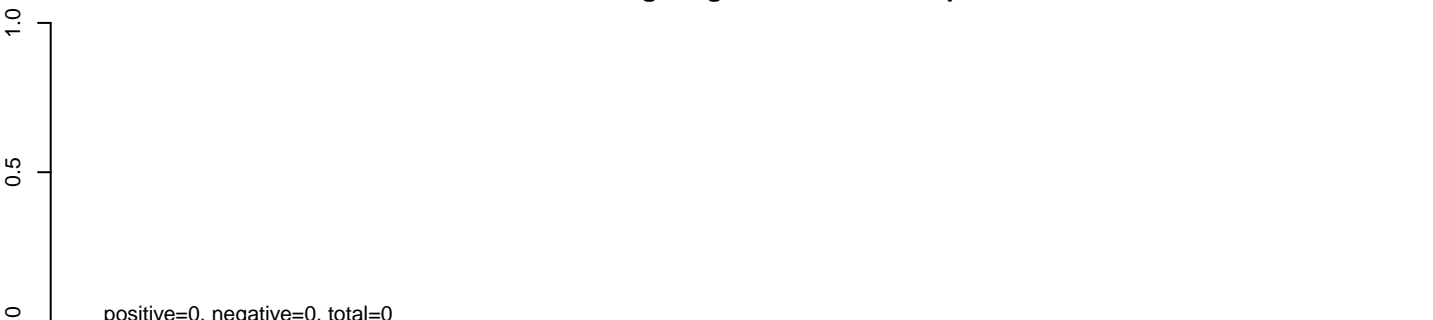
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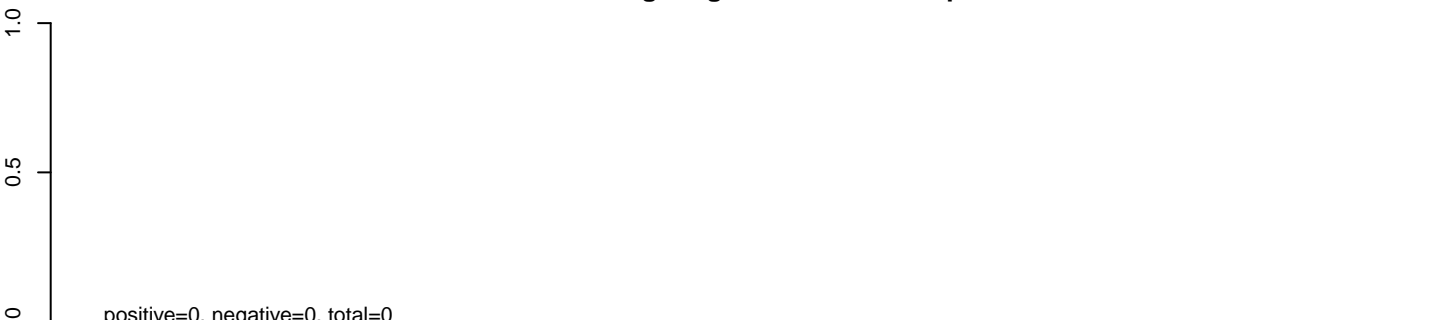
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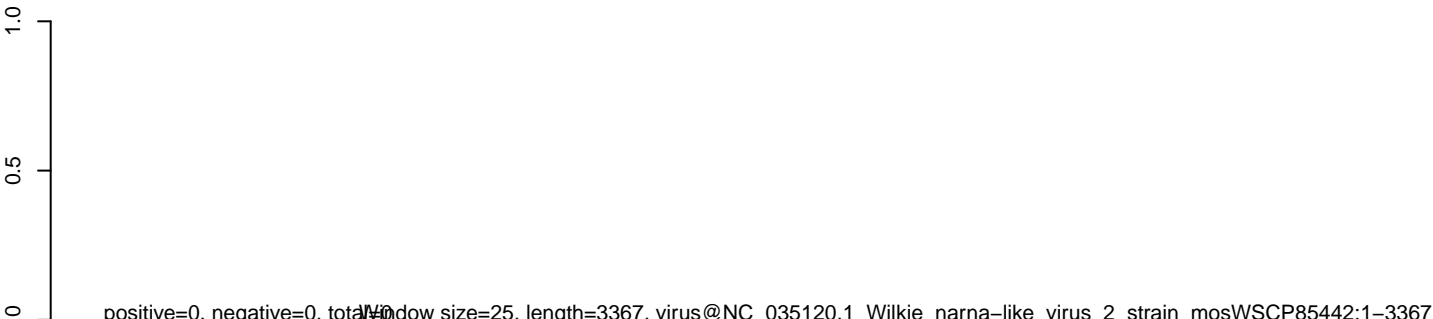
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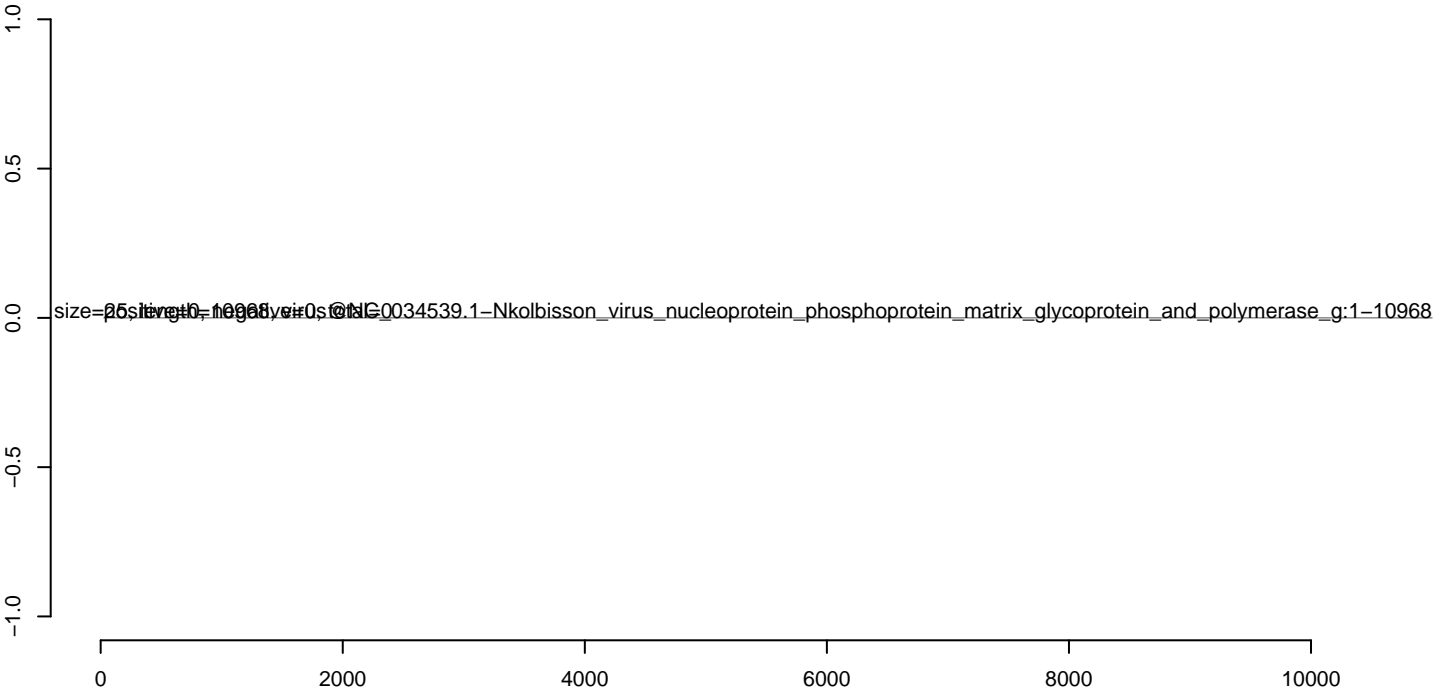
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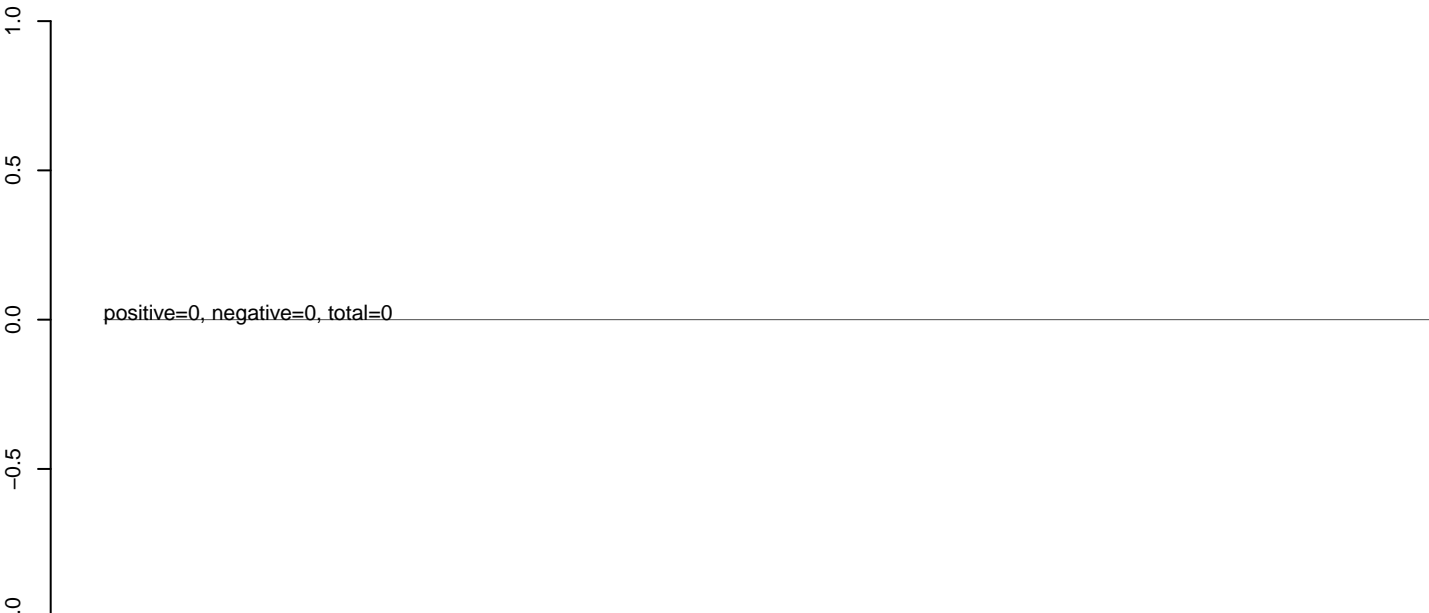
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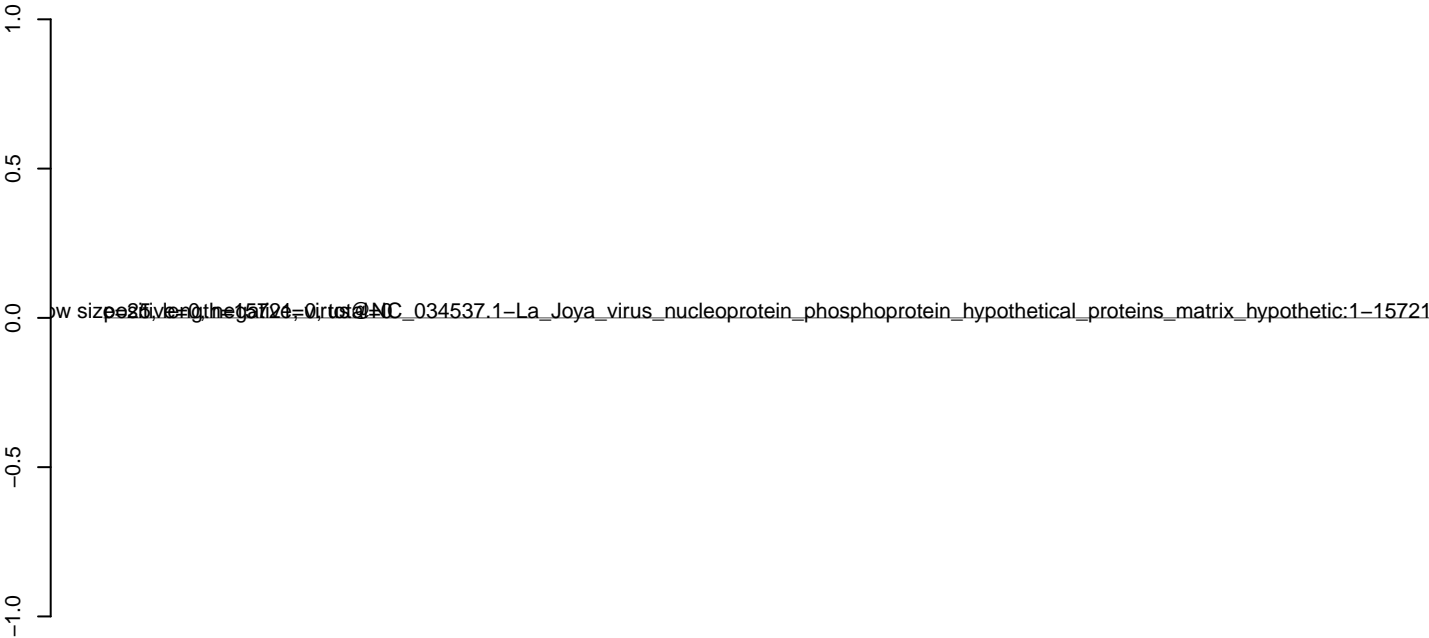
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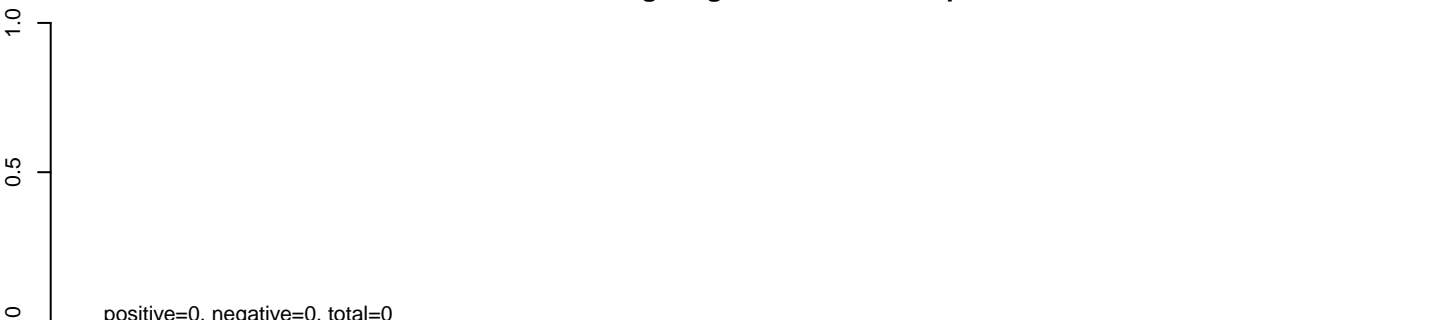
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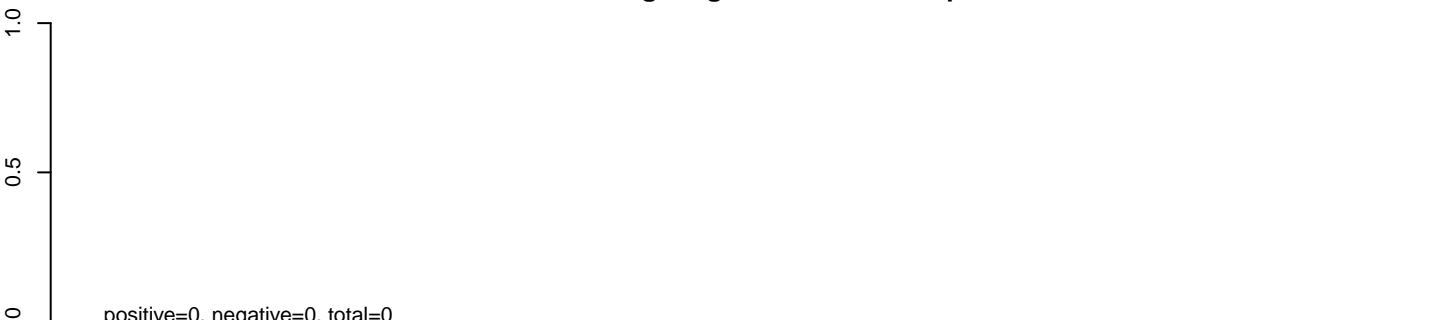
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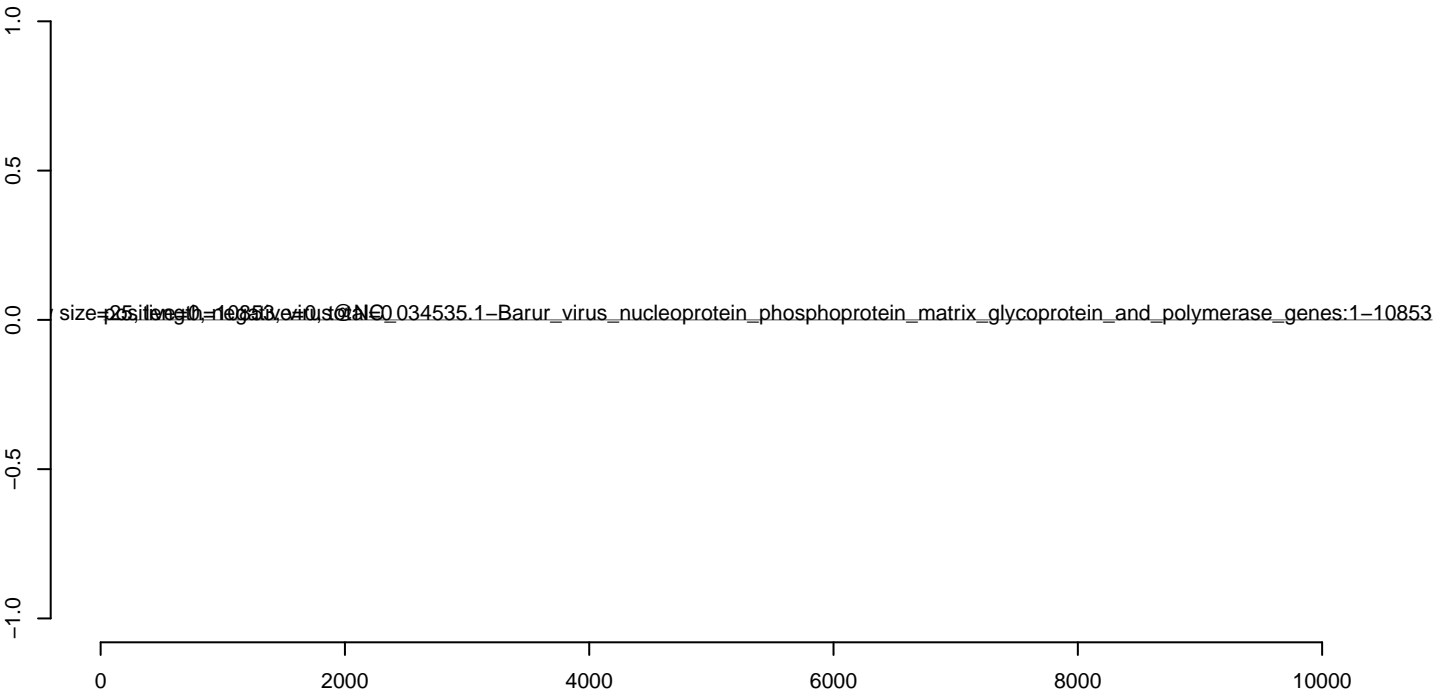
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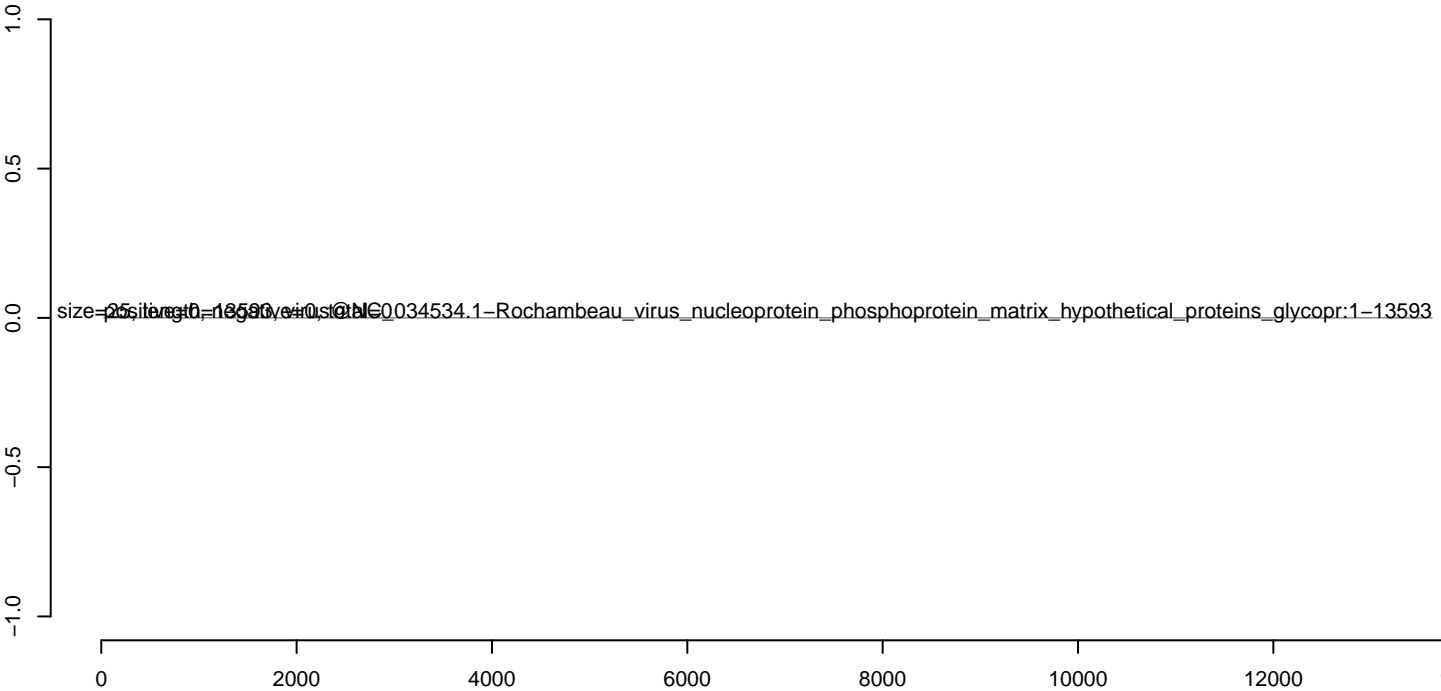
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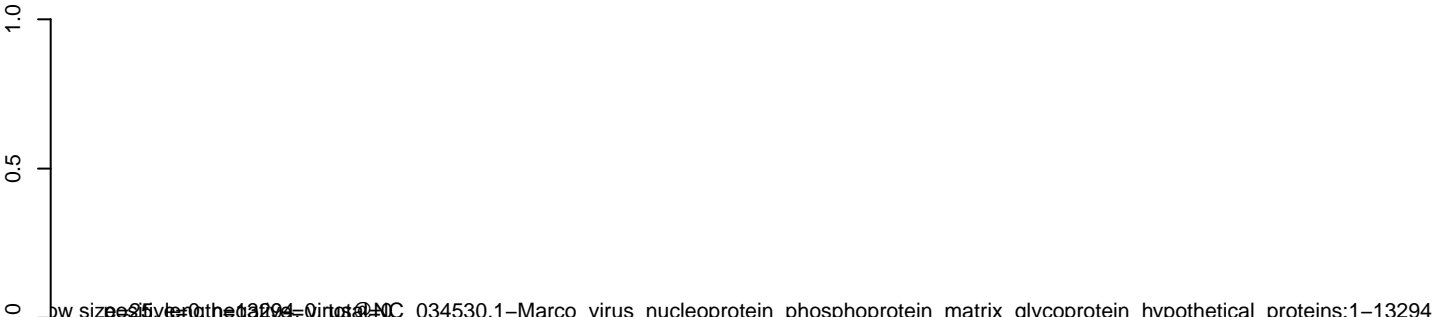
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AeAeg_Aag2_NL.M3.24_35.rep



AeAeg_Aag2_NL.M3.rep



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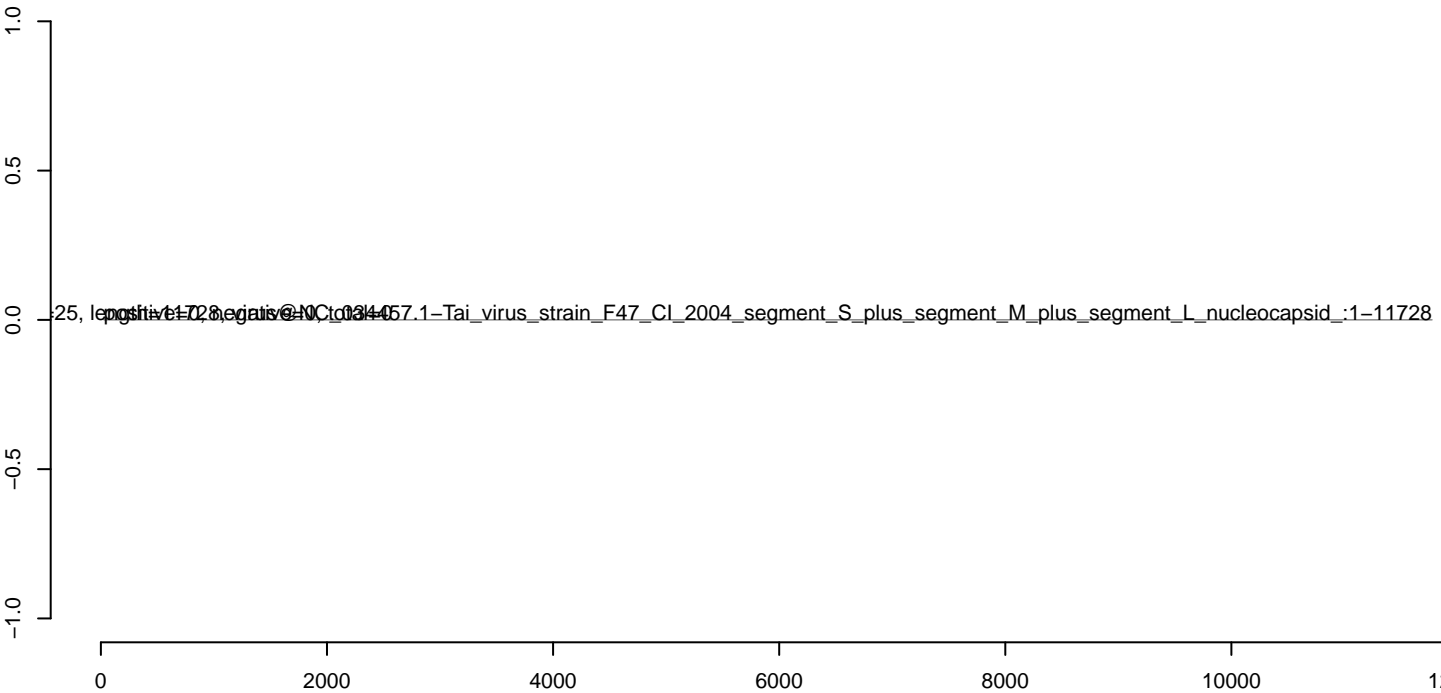
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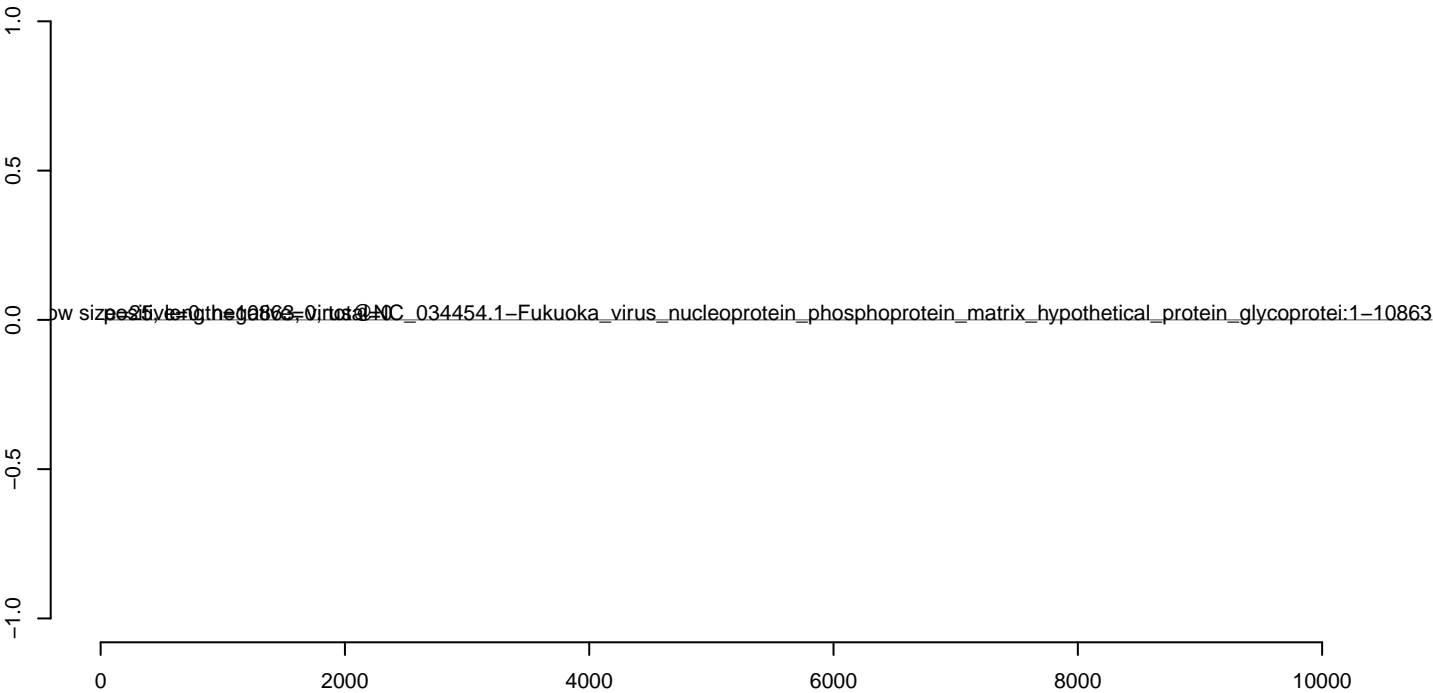
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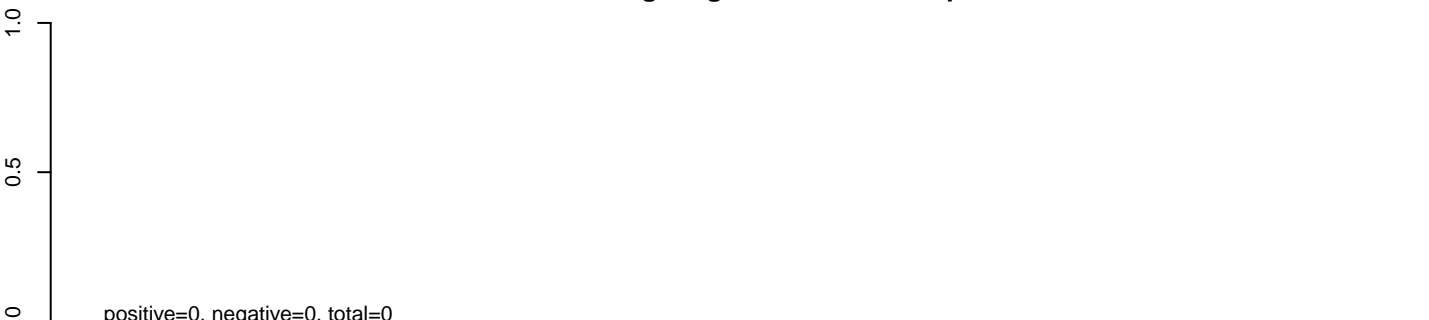
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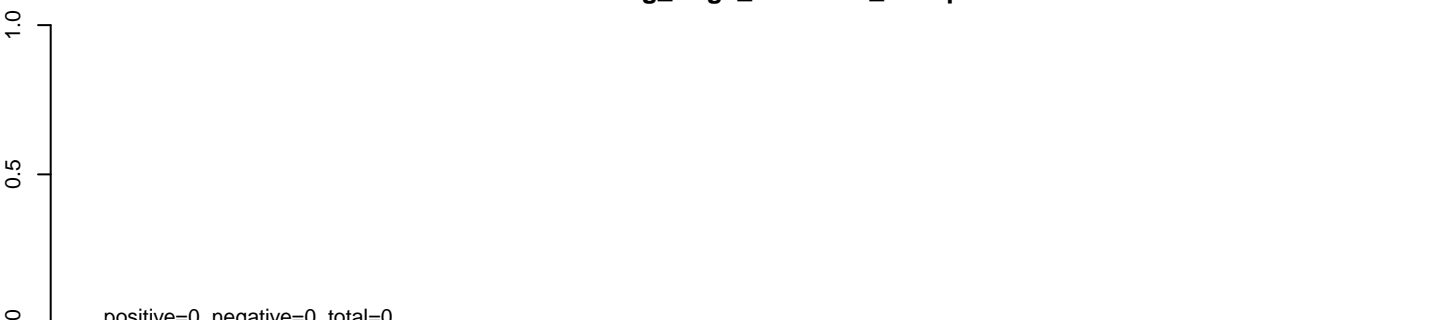
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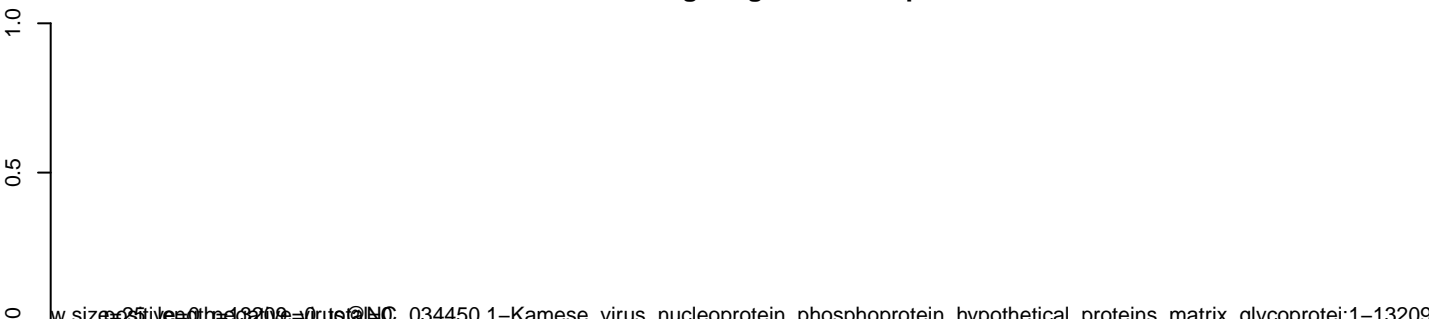
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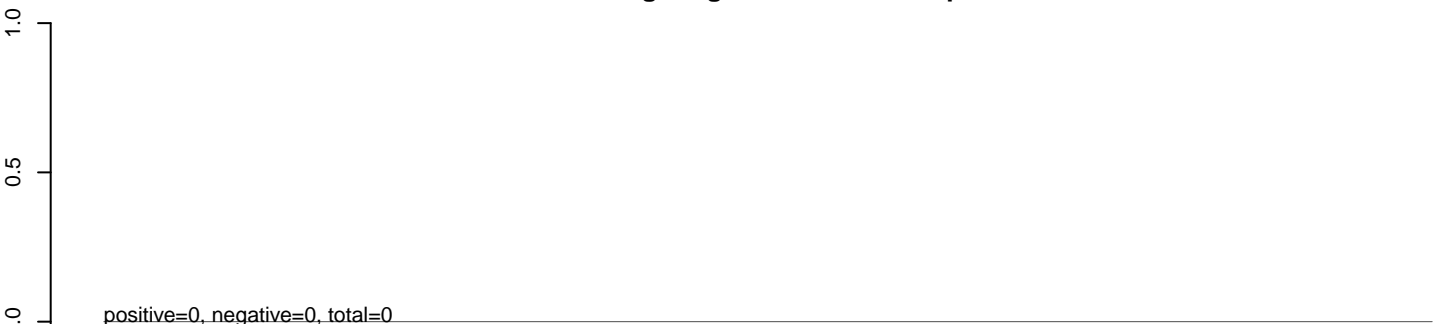
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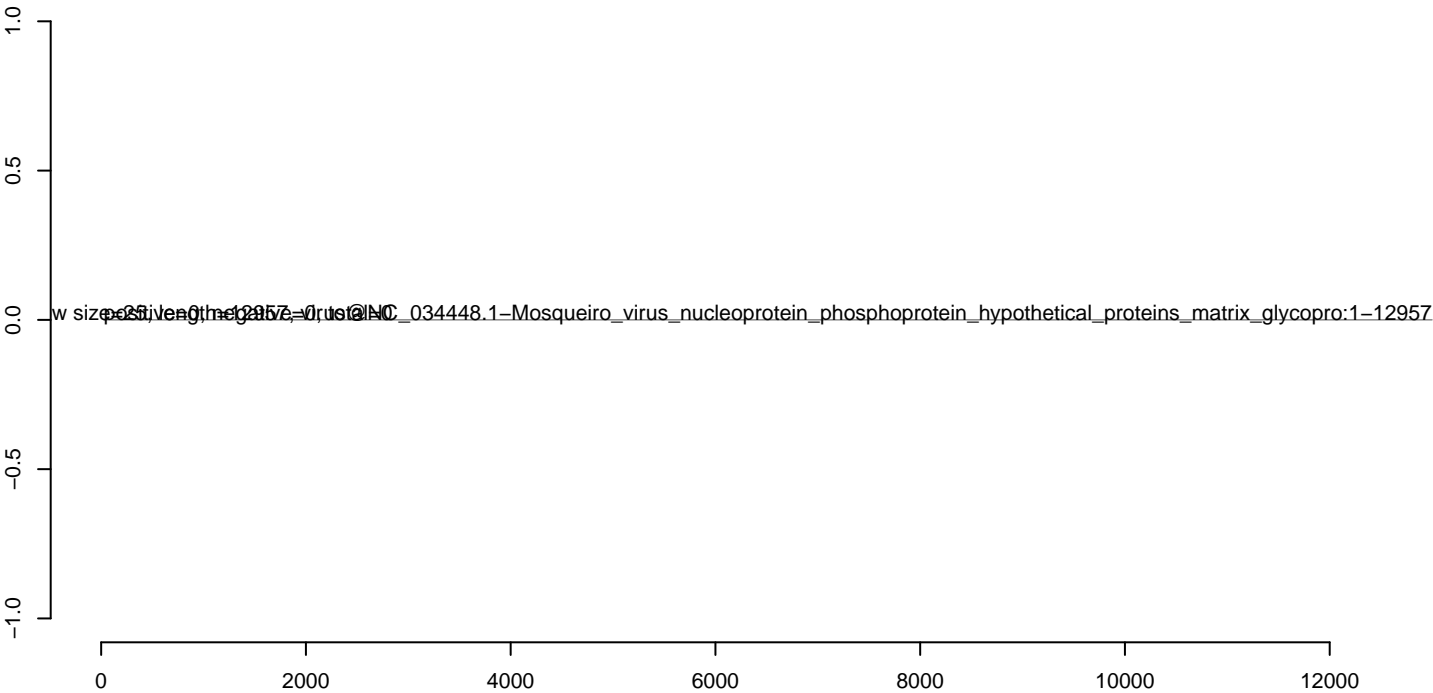
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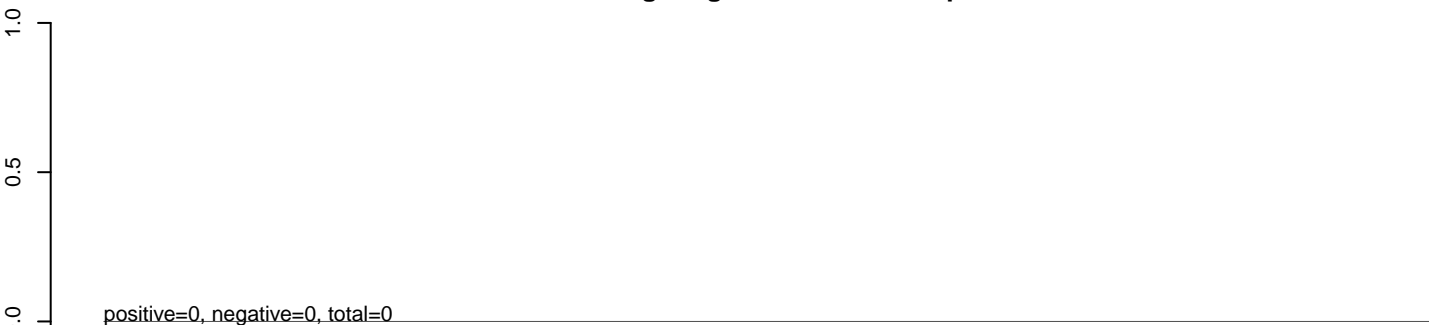
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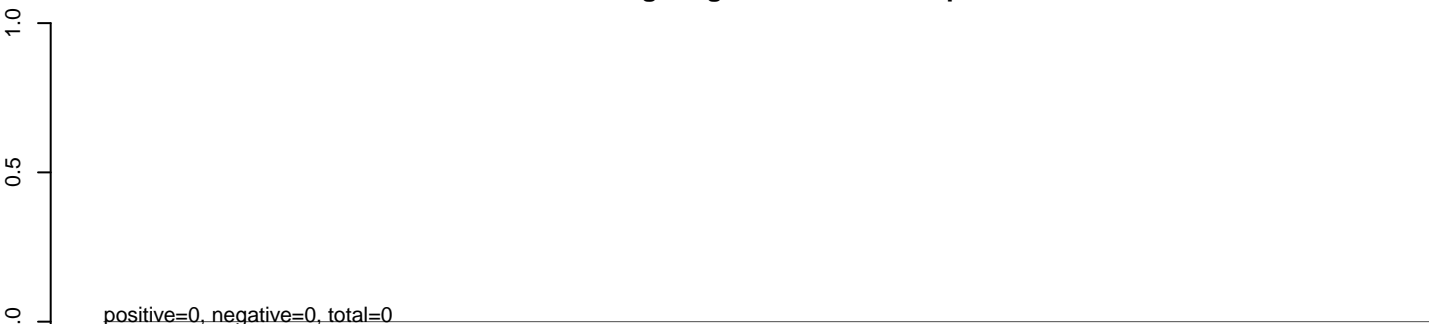
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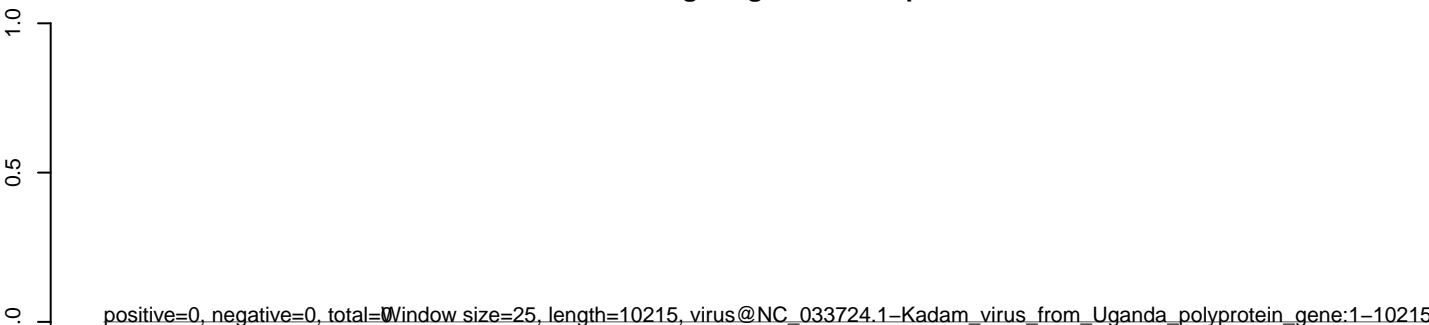
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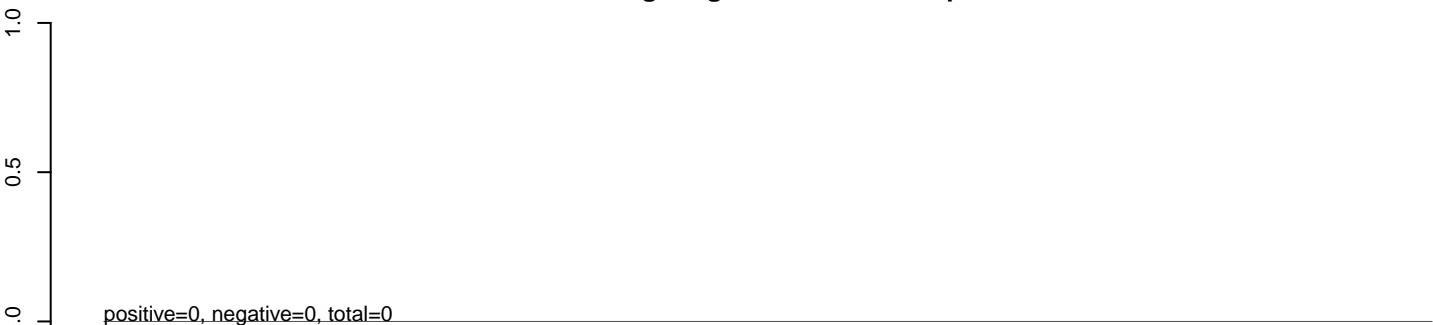


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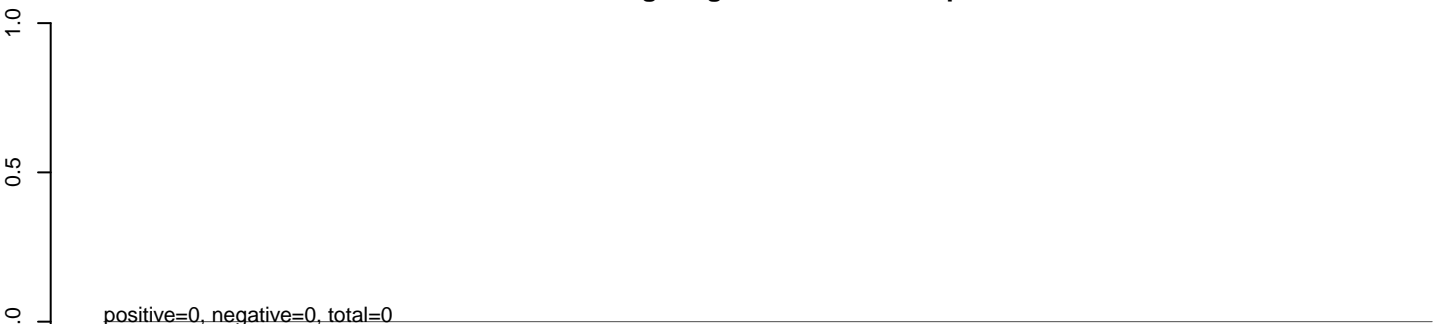


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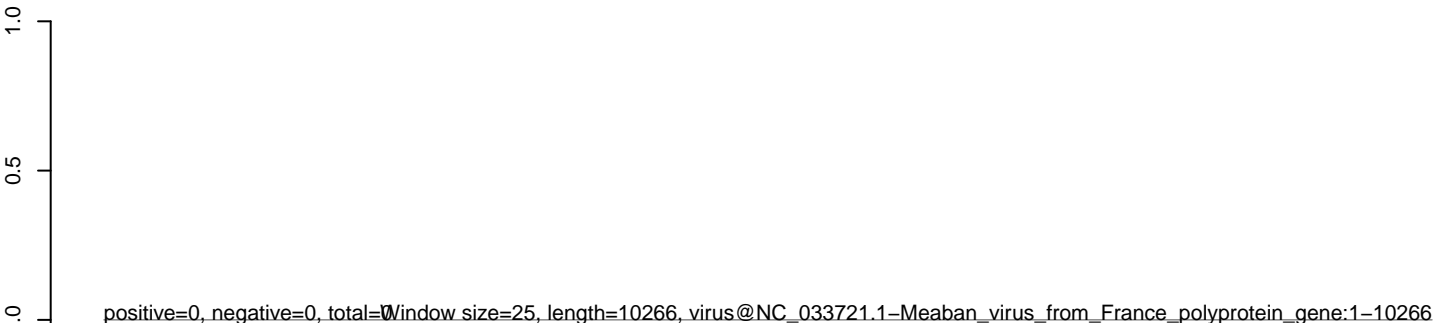
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AeAeg_Aag2_NL.M3.rep



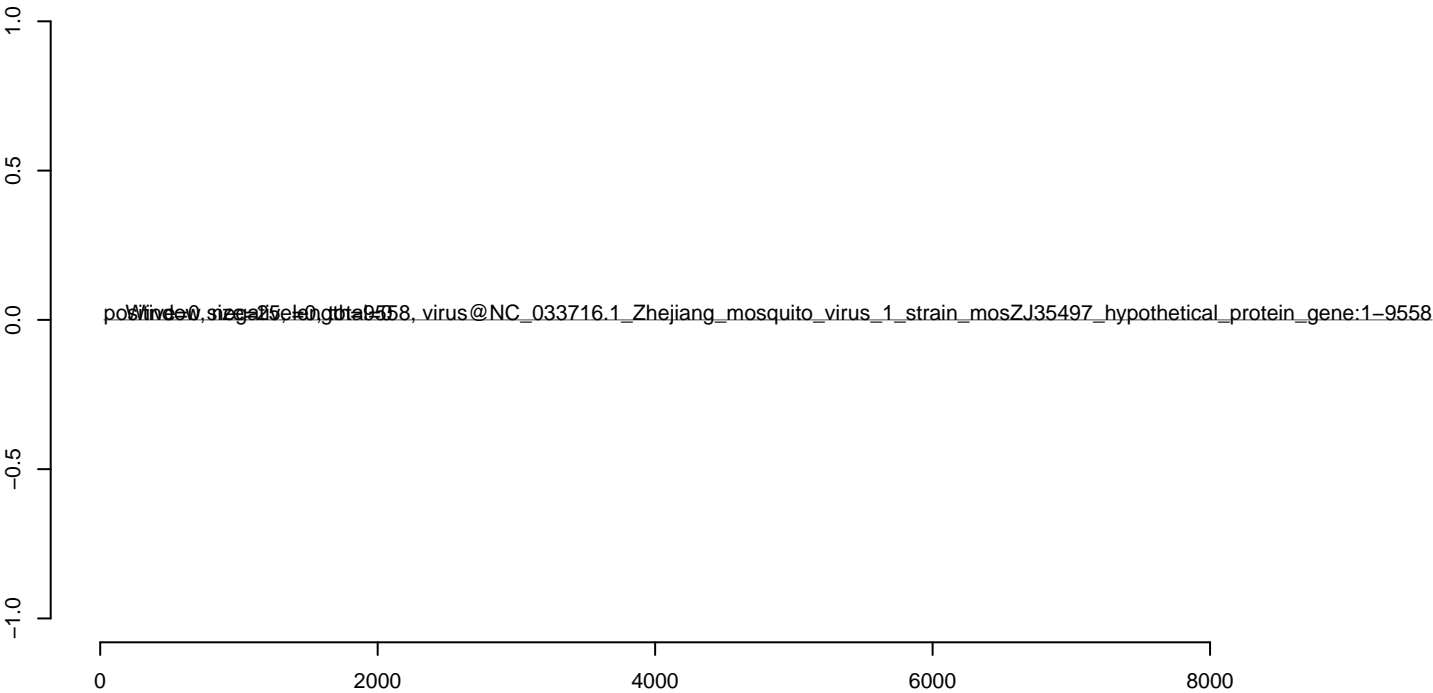
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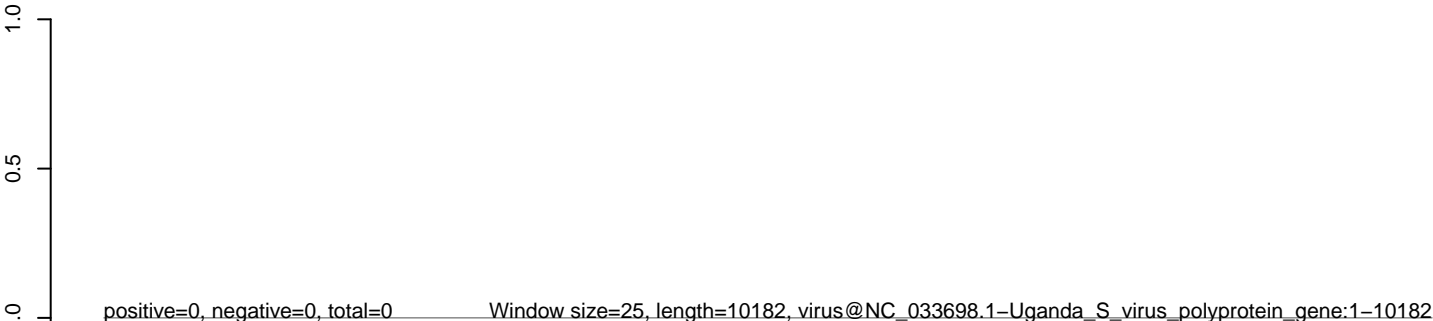
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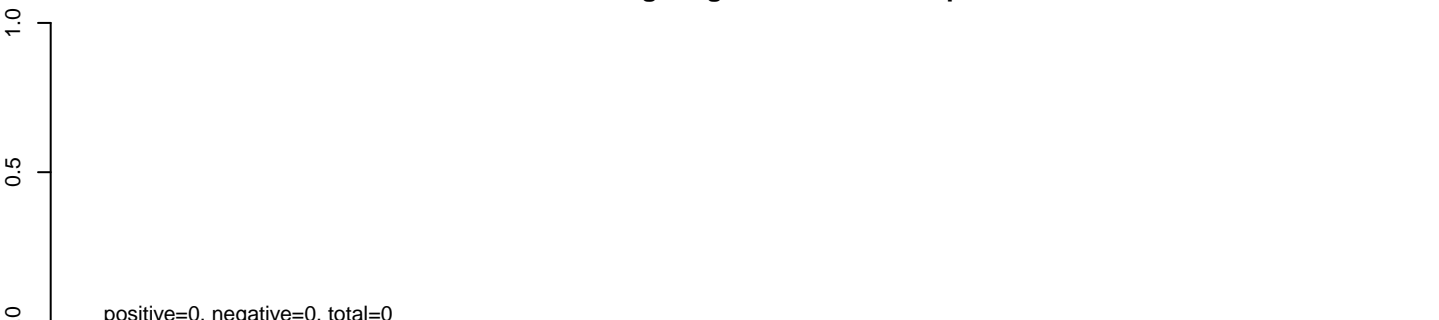
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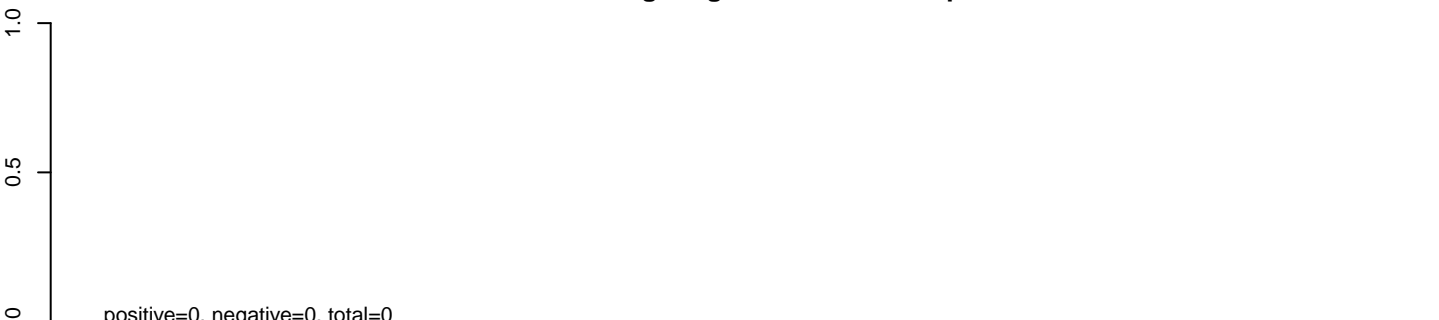
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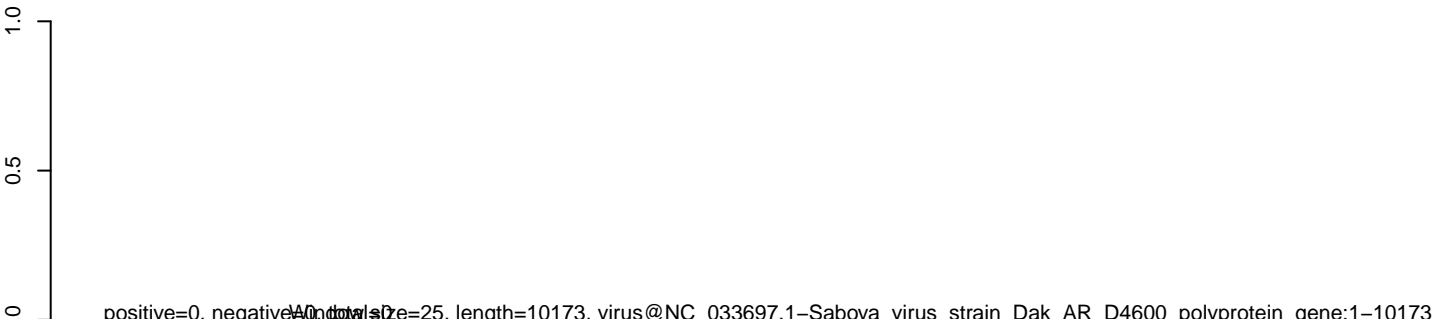
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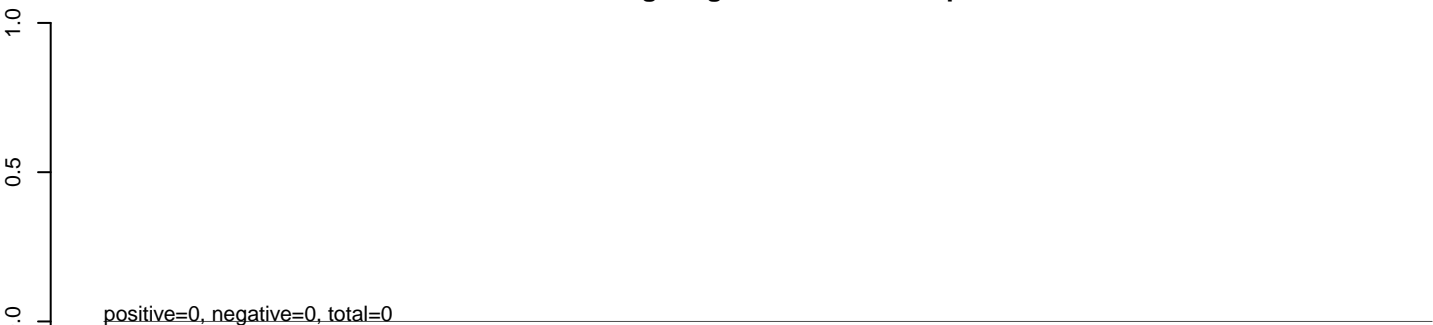
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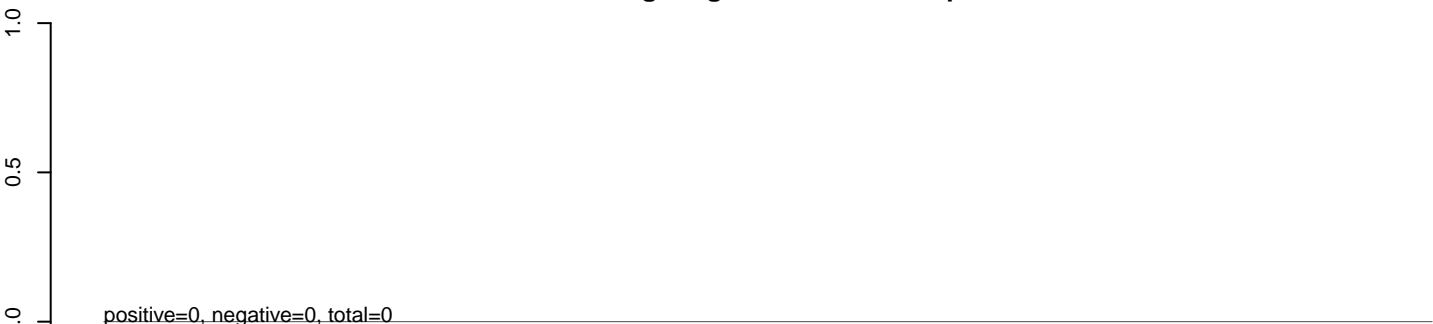
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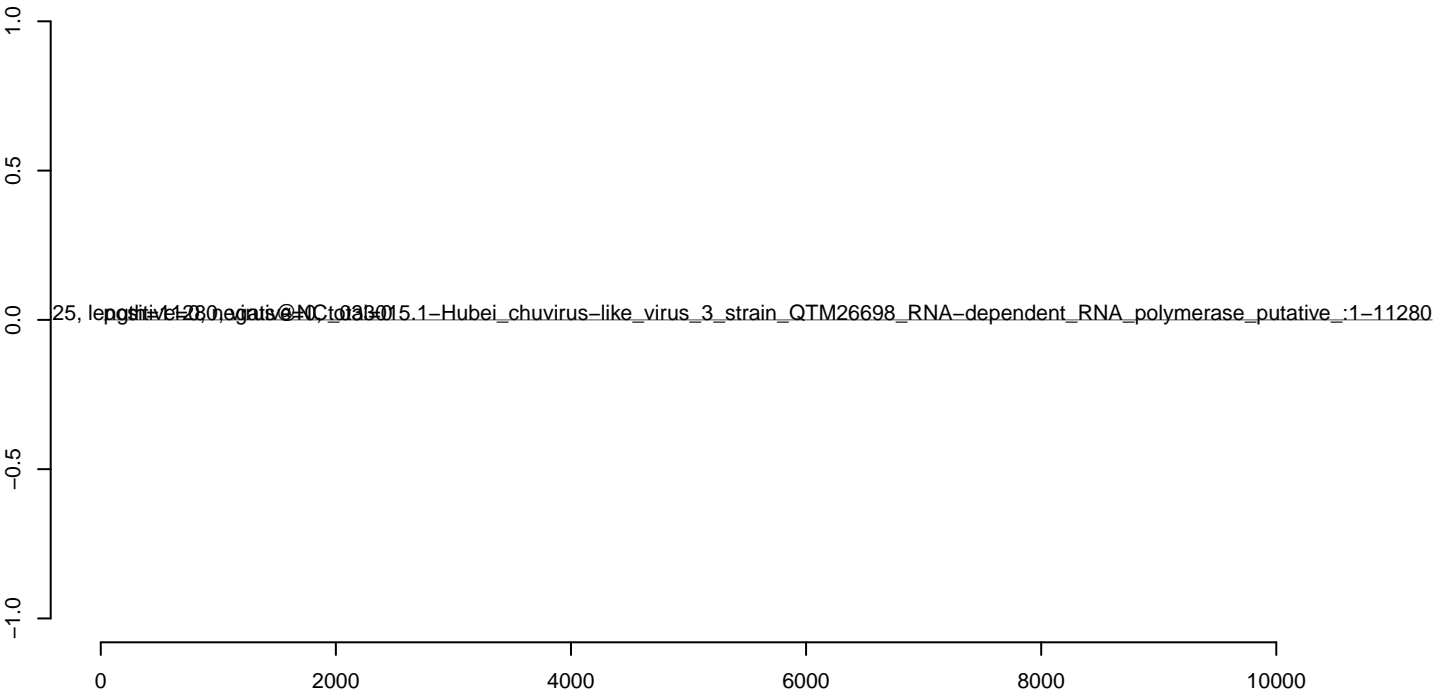
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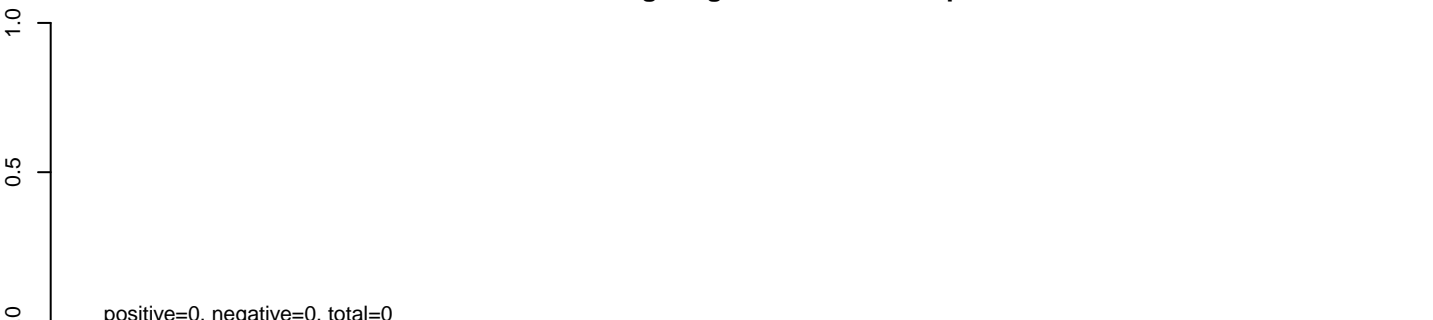
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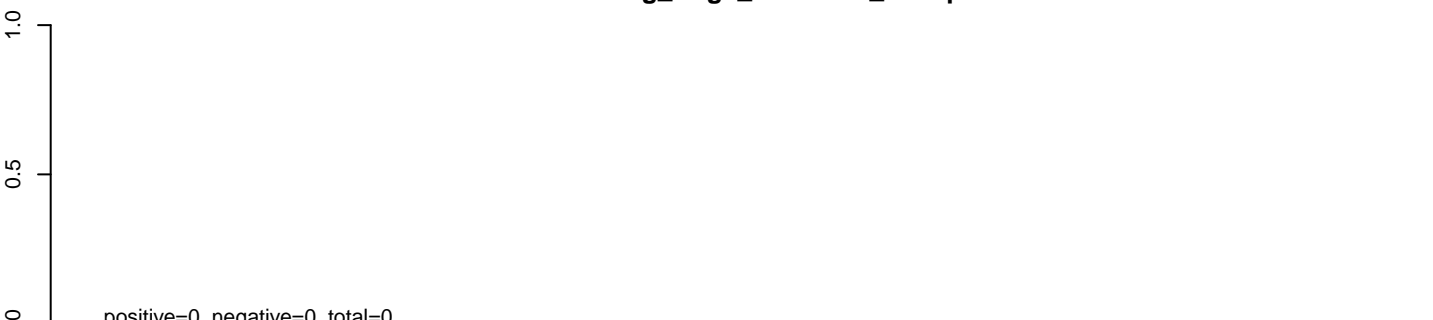
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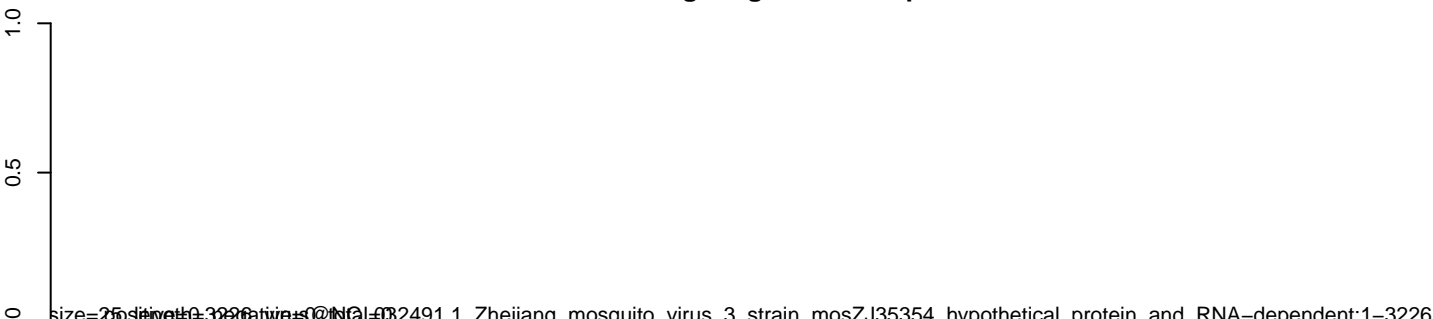
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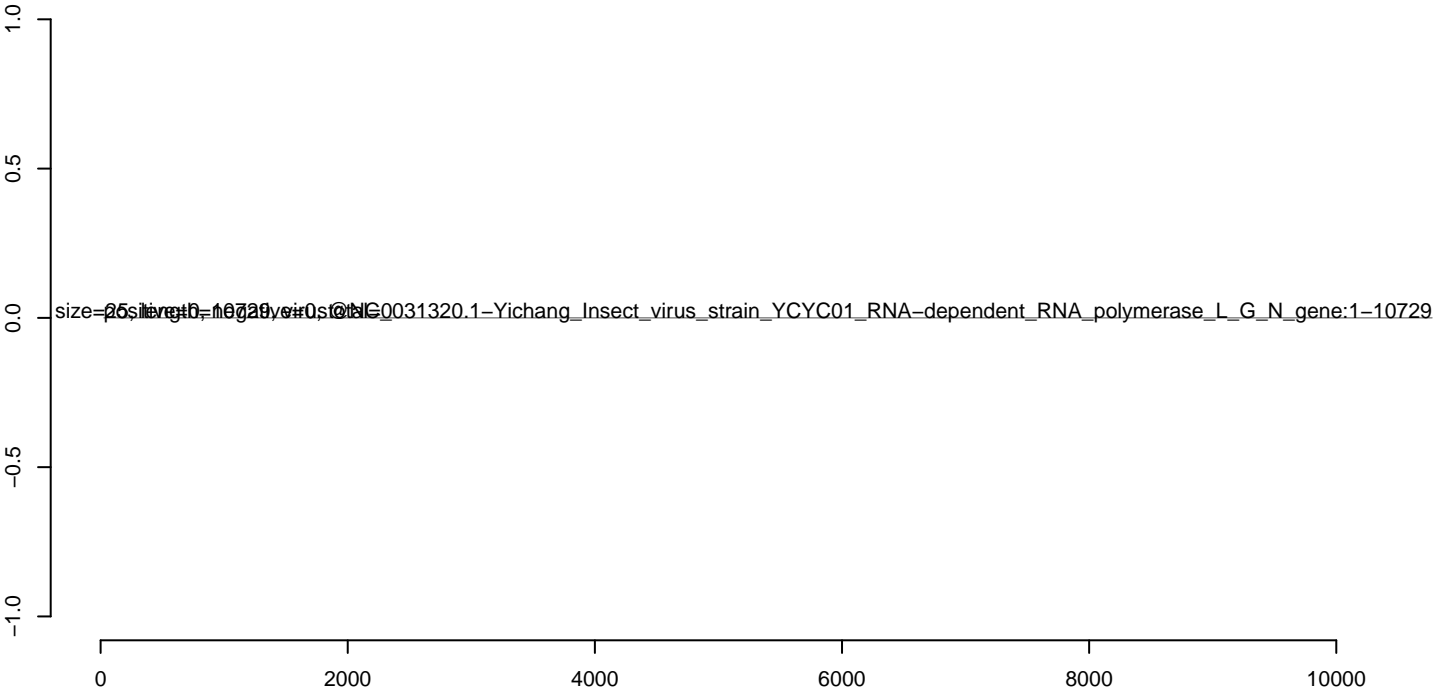
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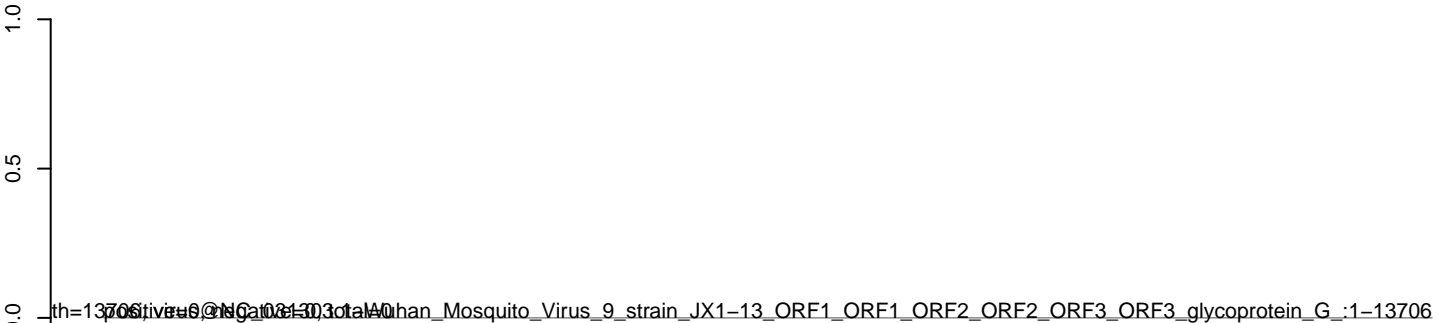
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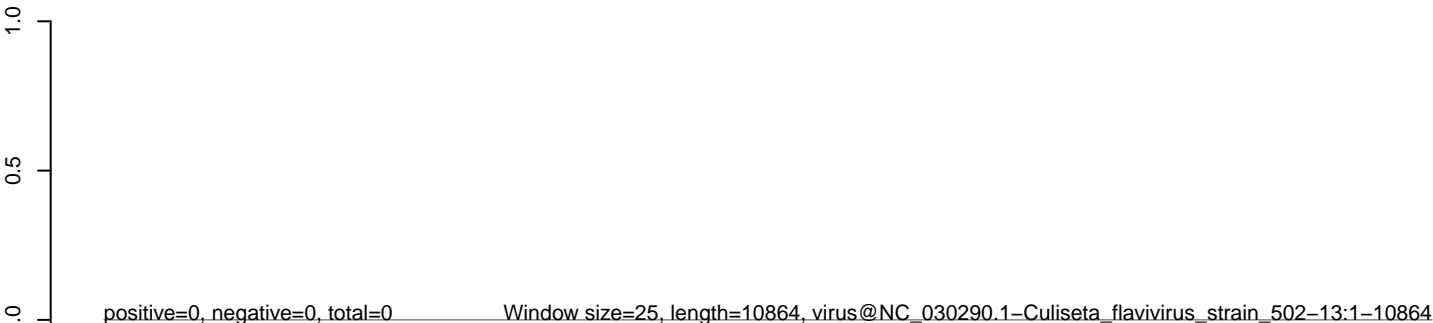
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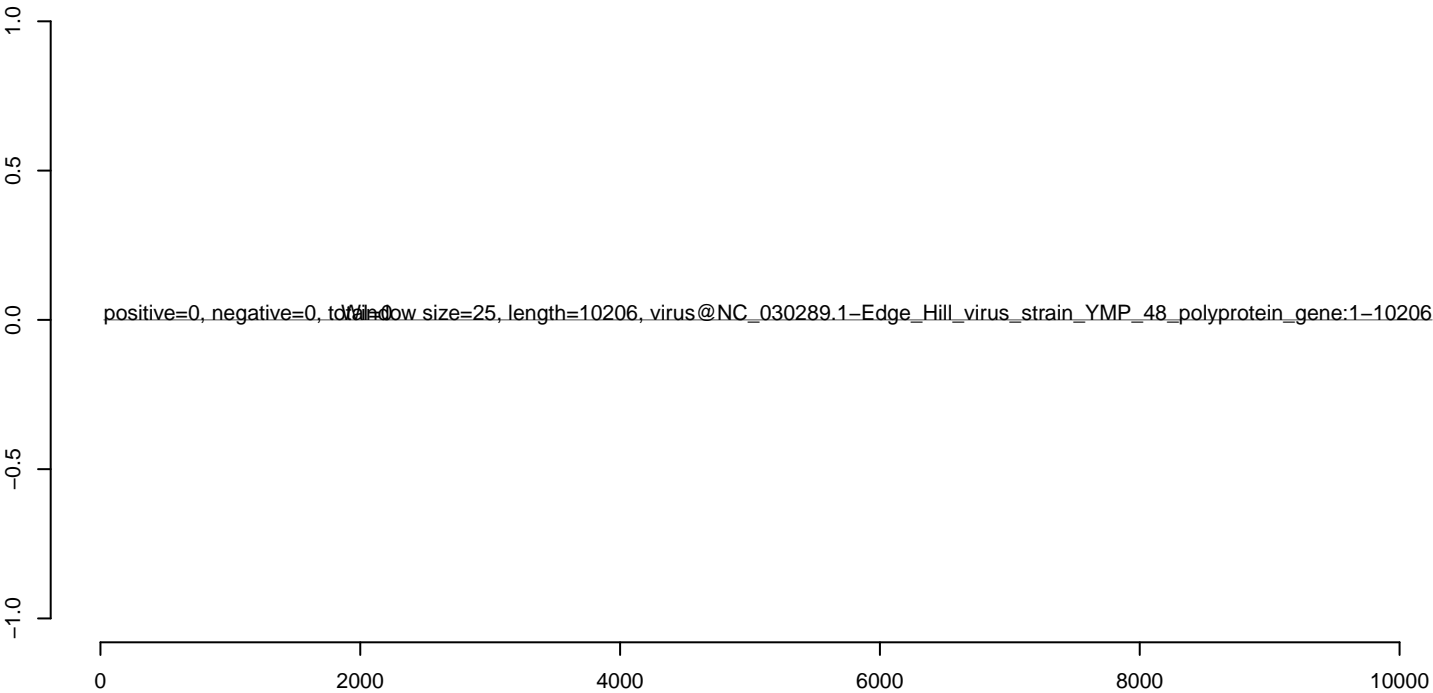
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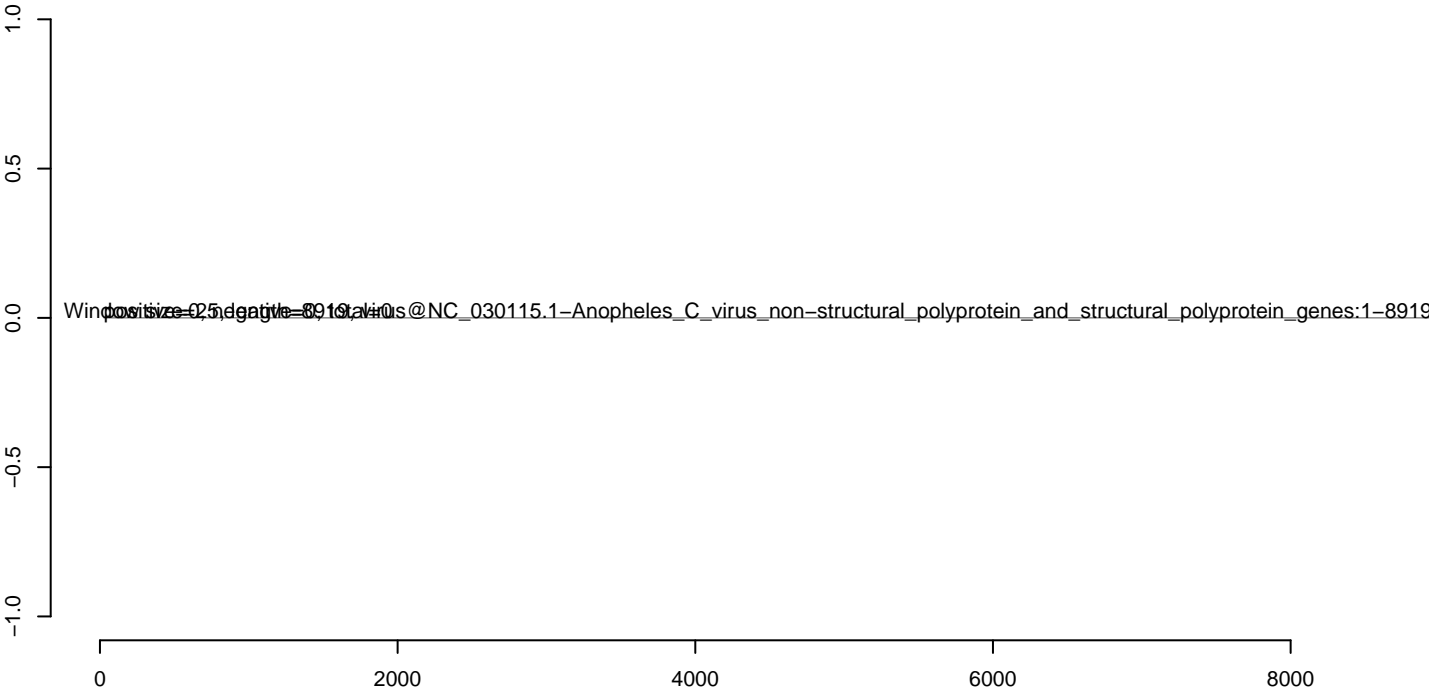
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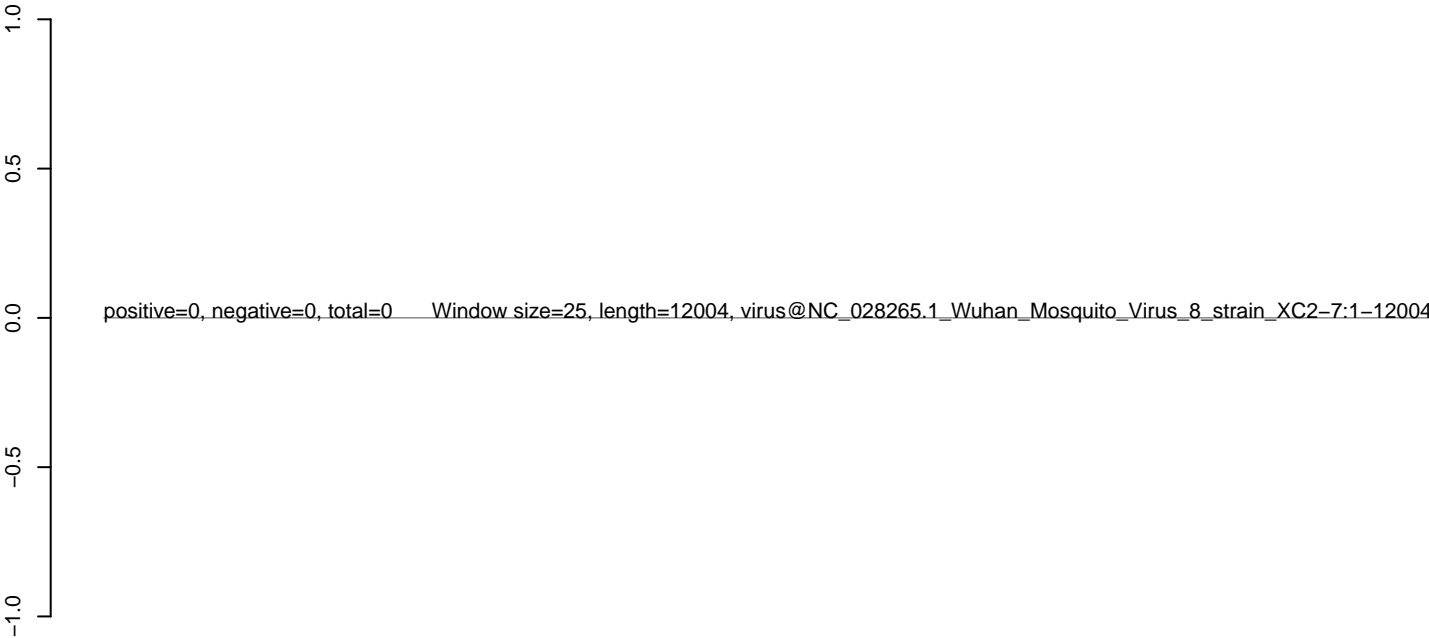
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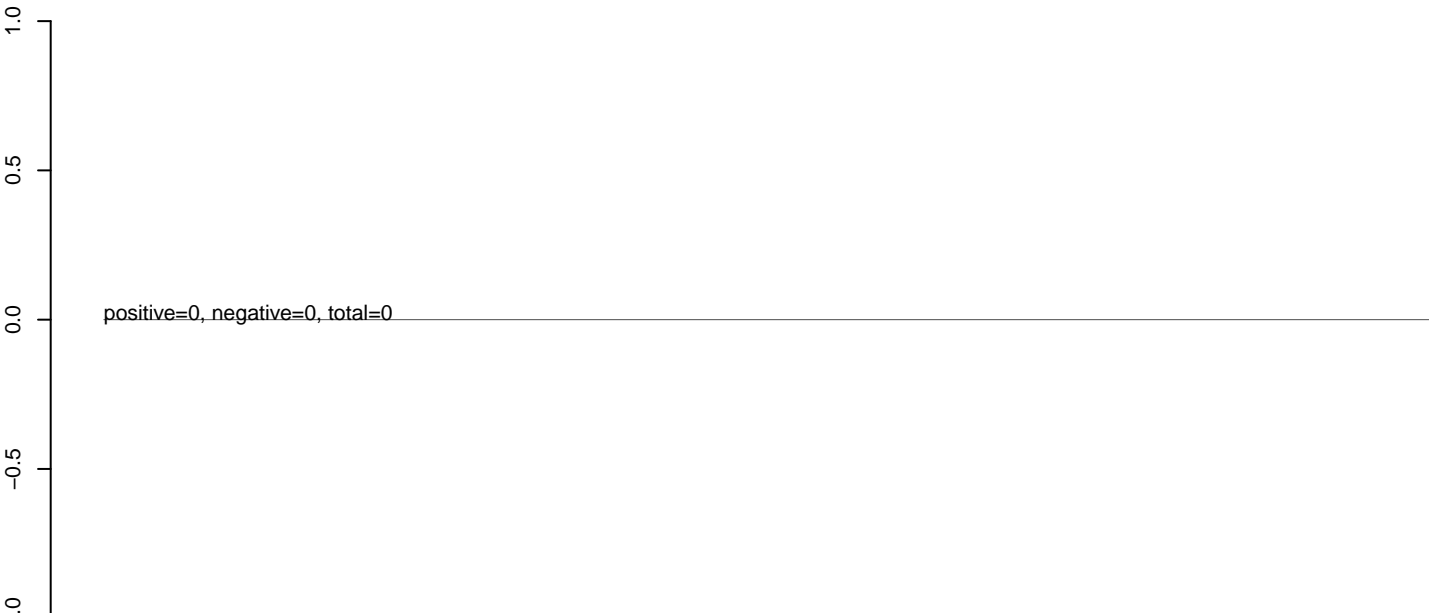


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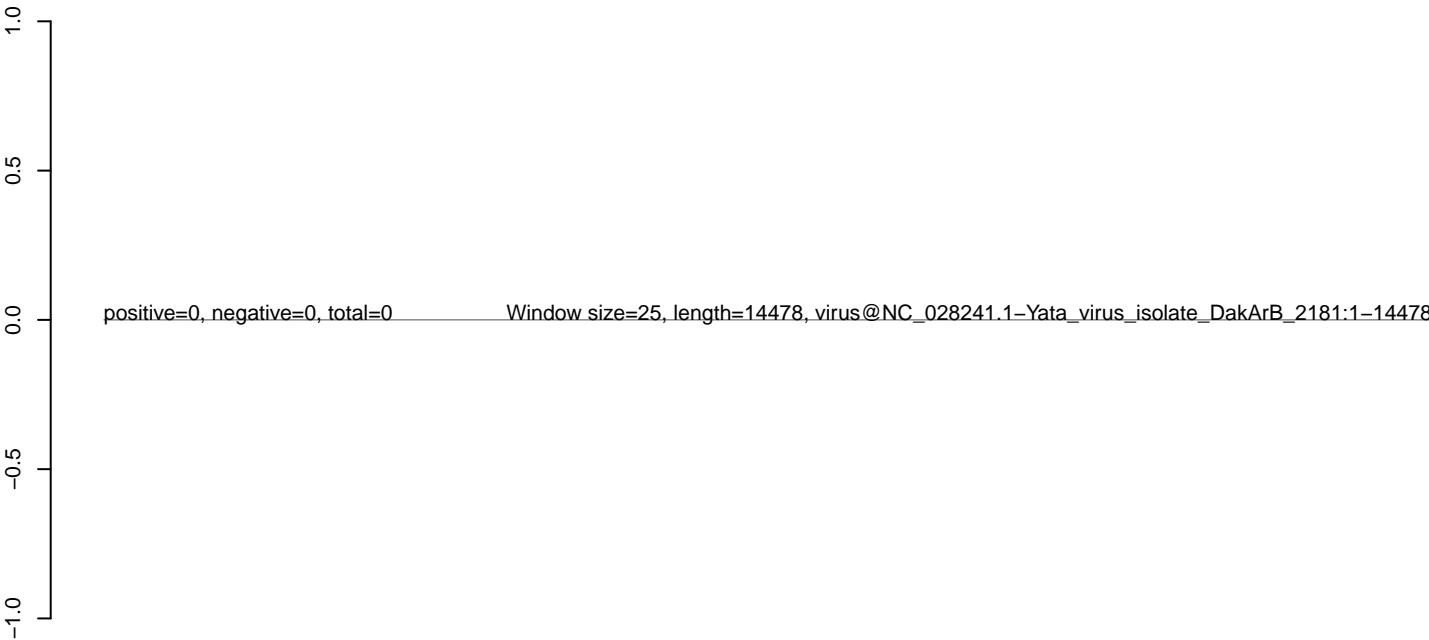
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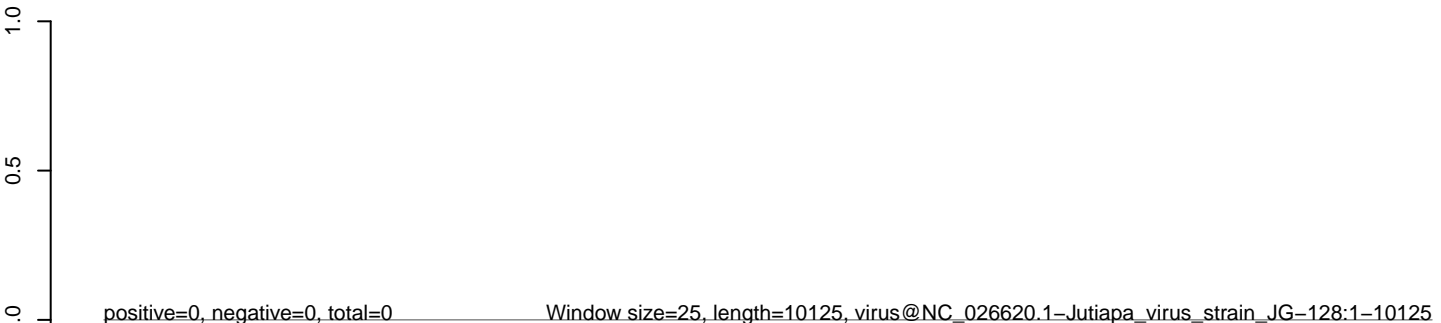
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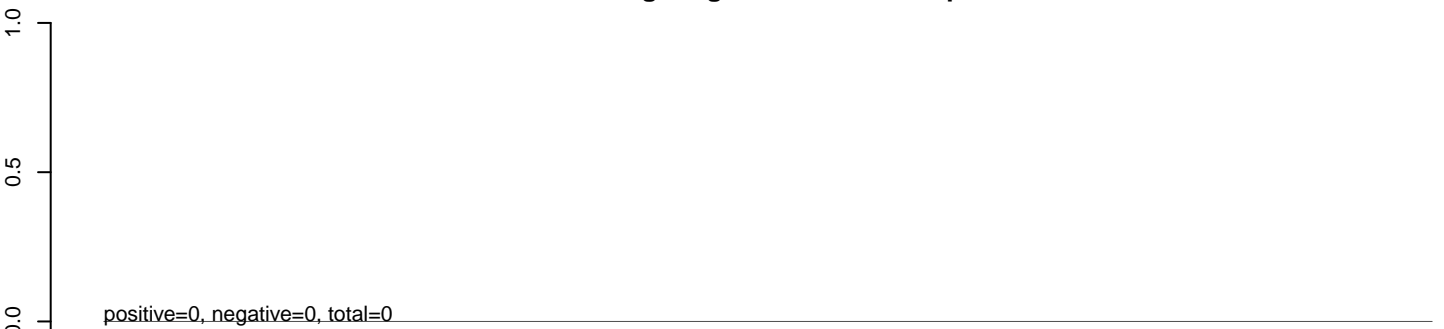
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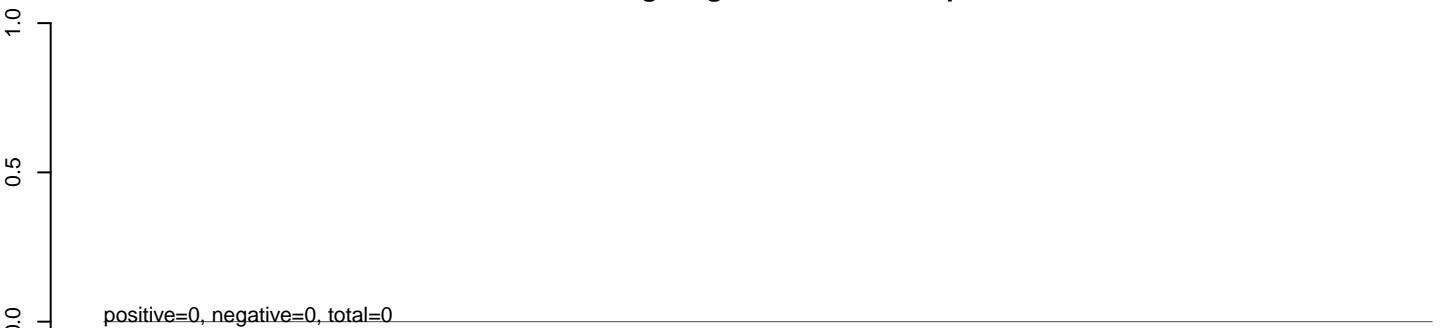
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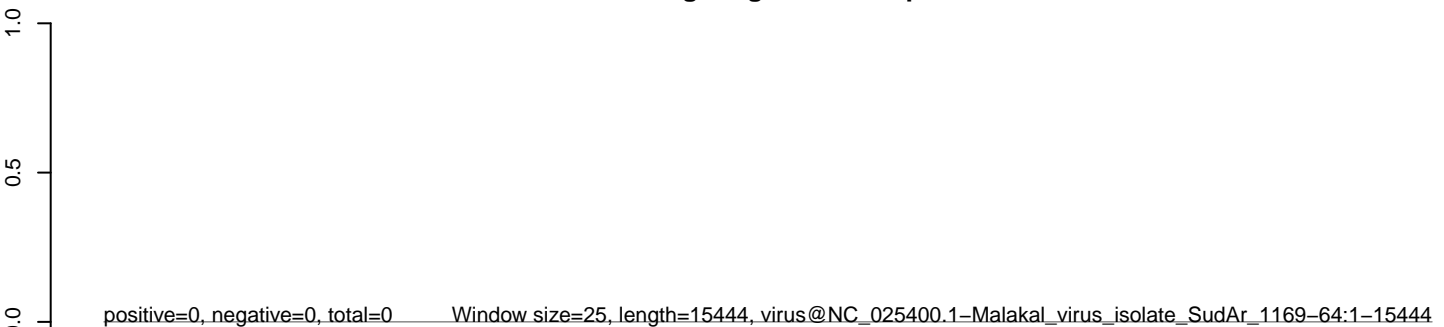
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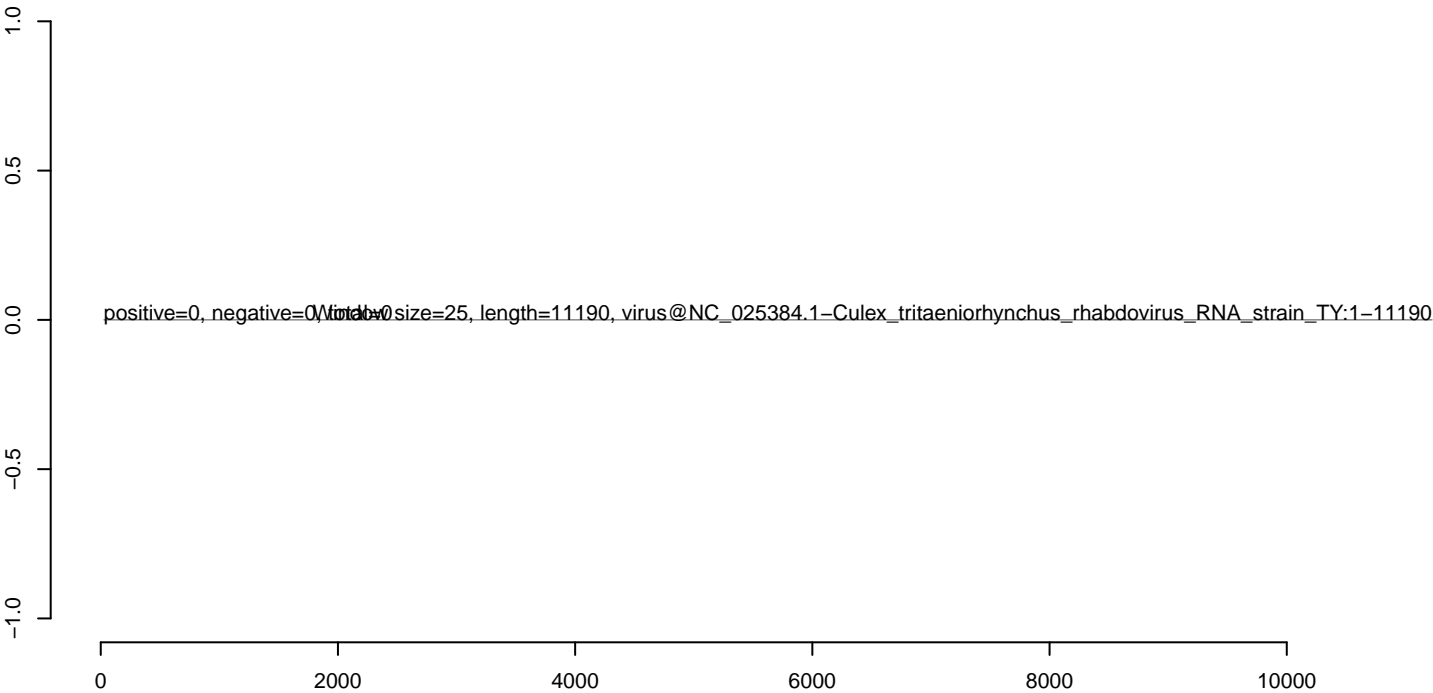
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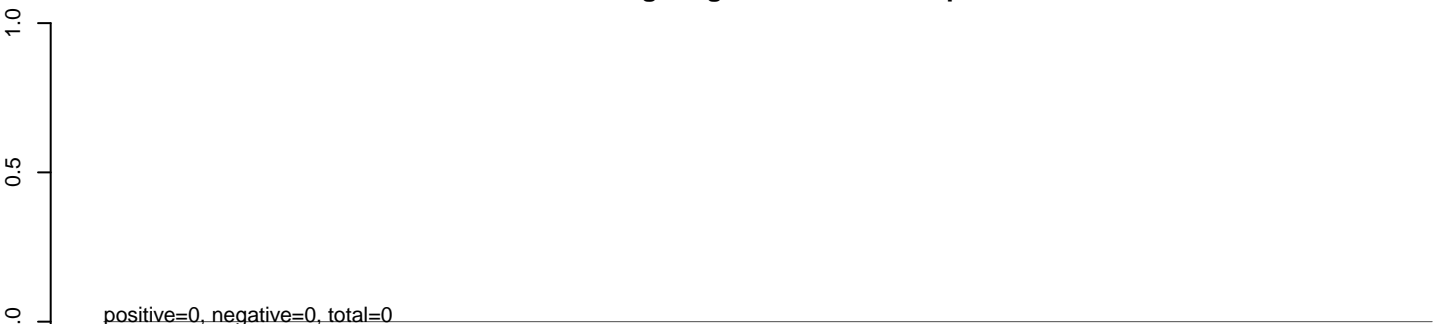
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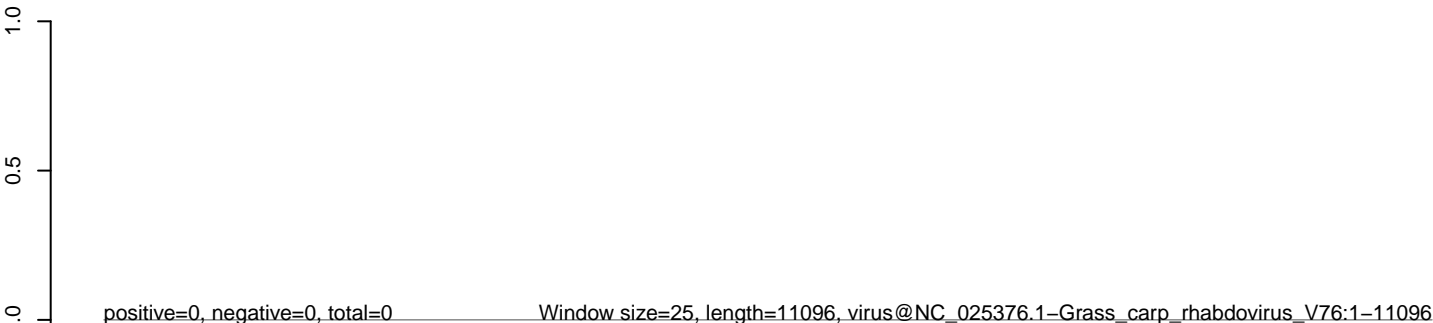
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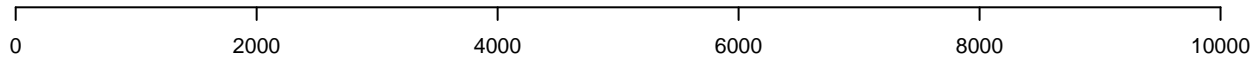
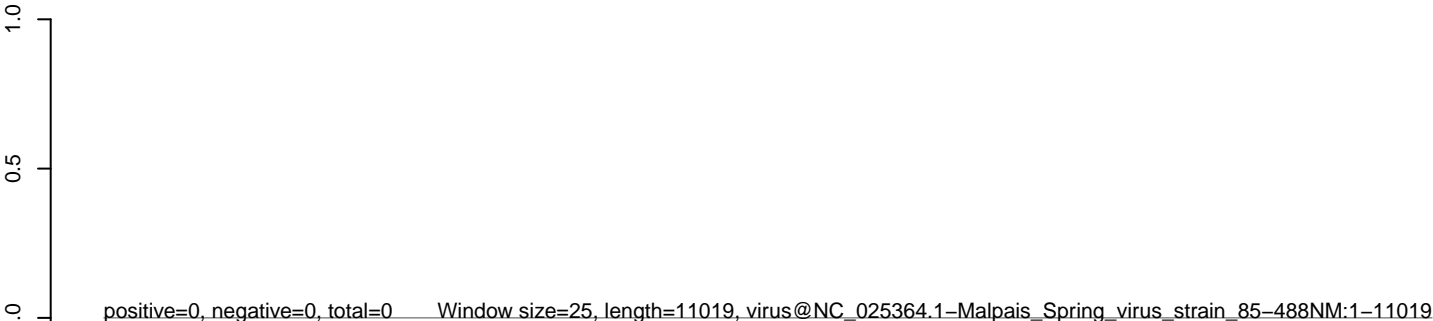
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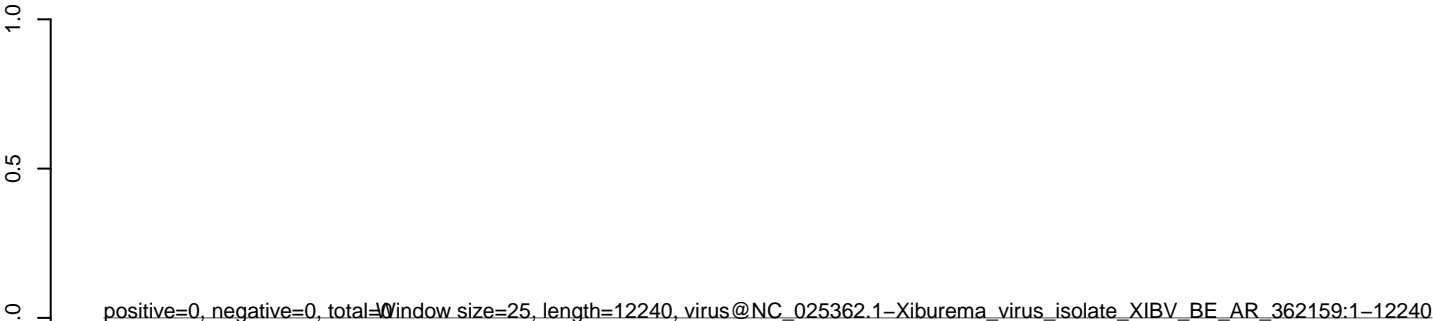
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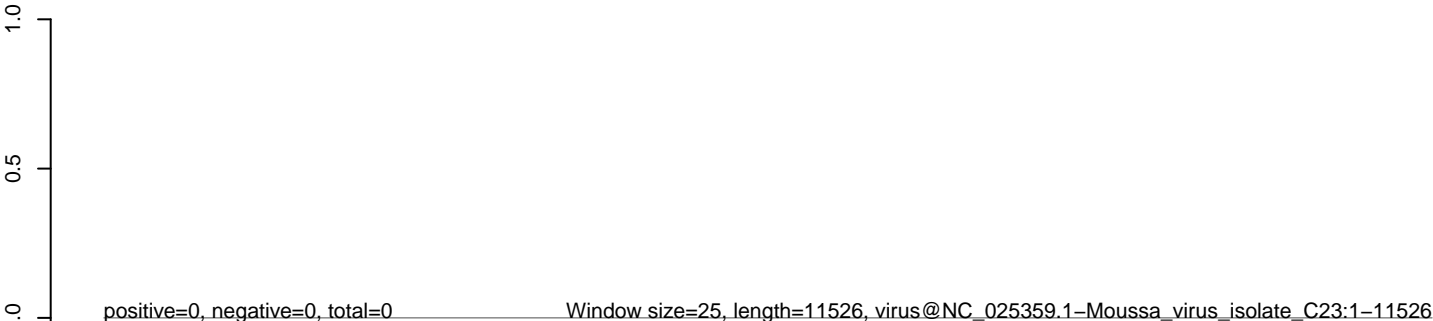
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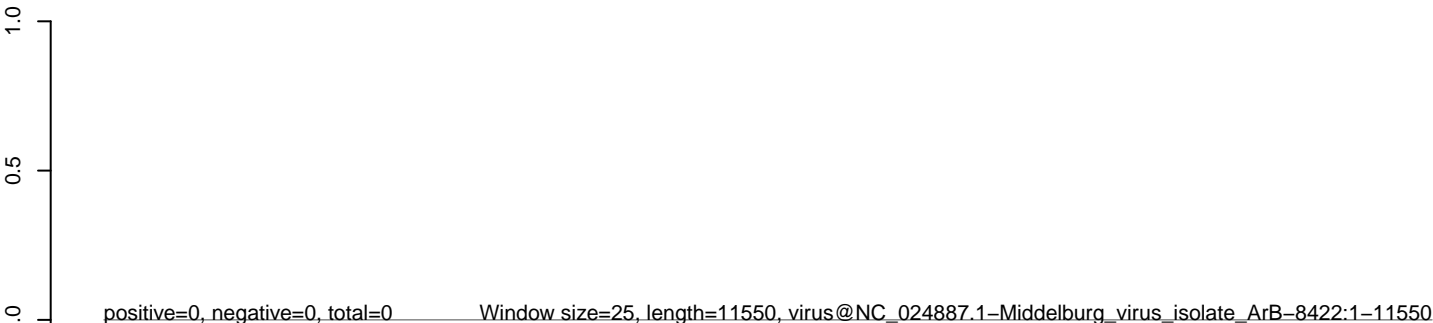
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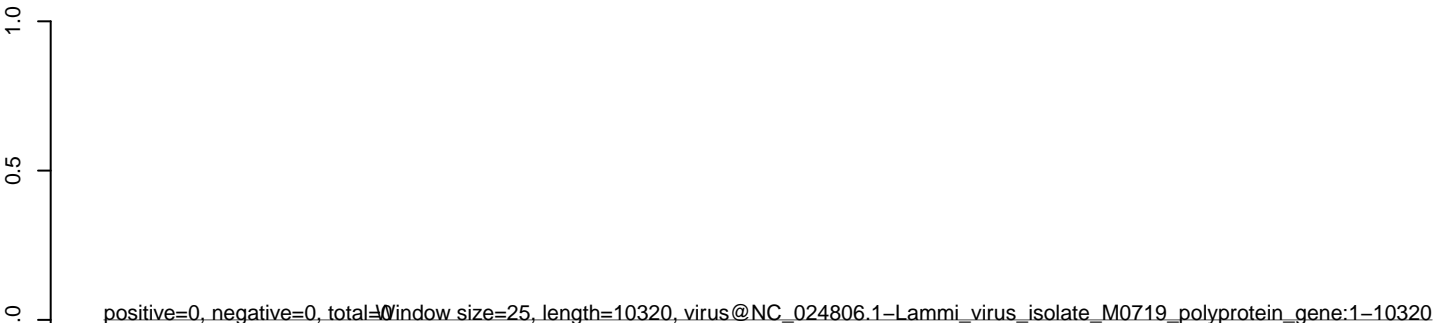
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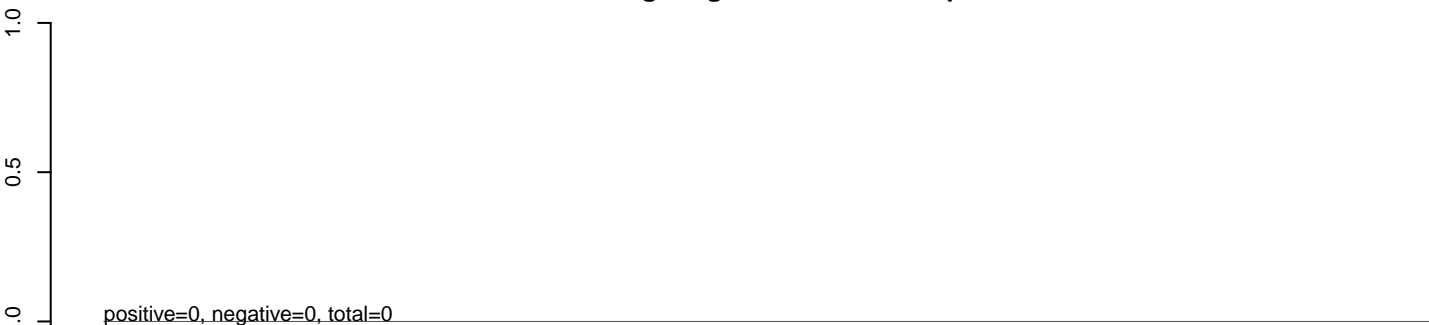
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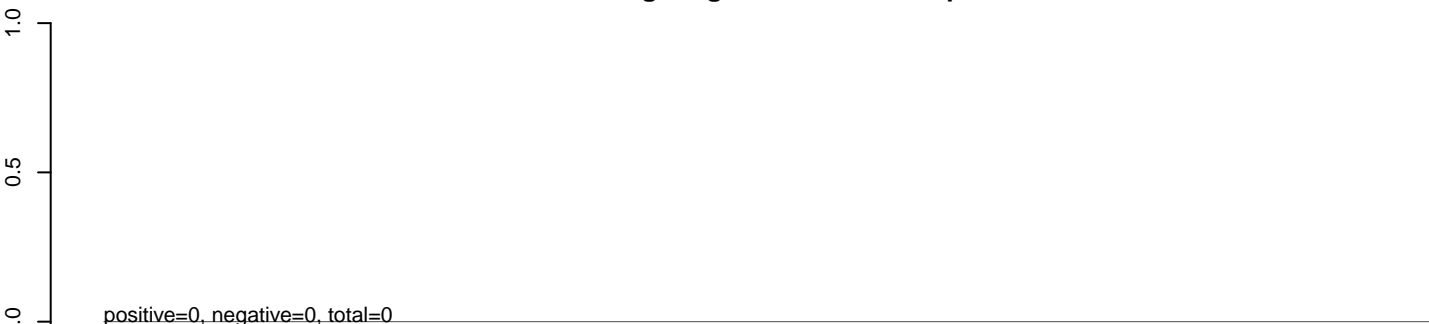
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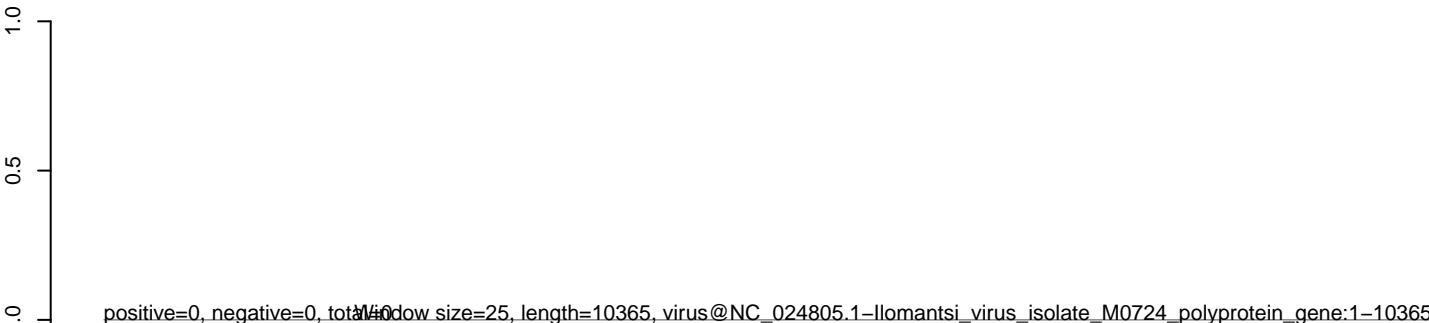
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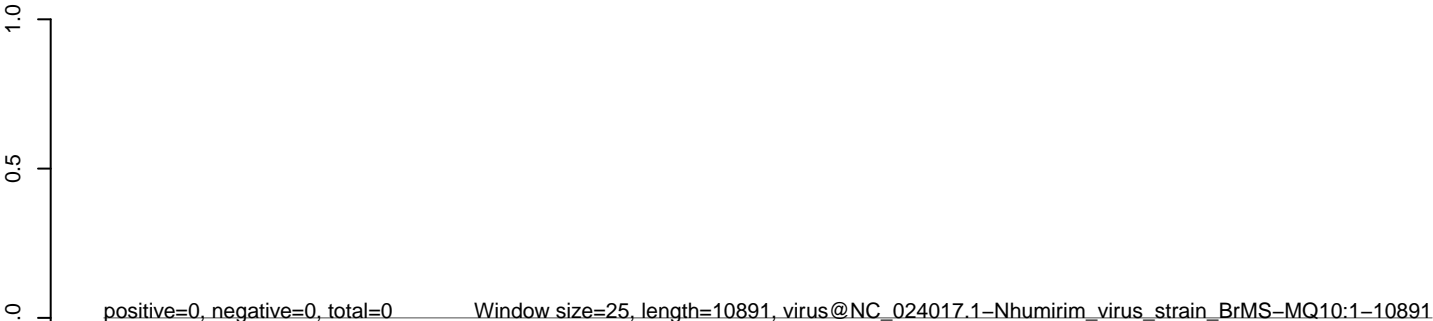
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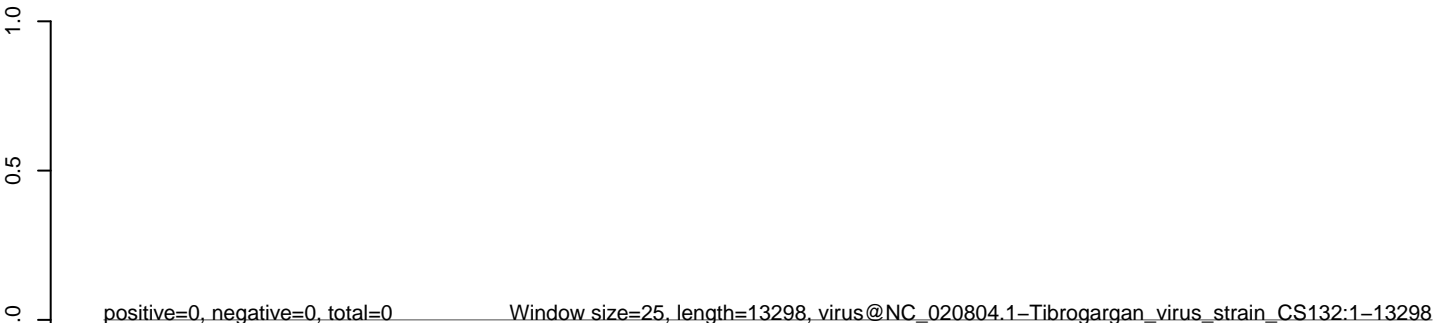
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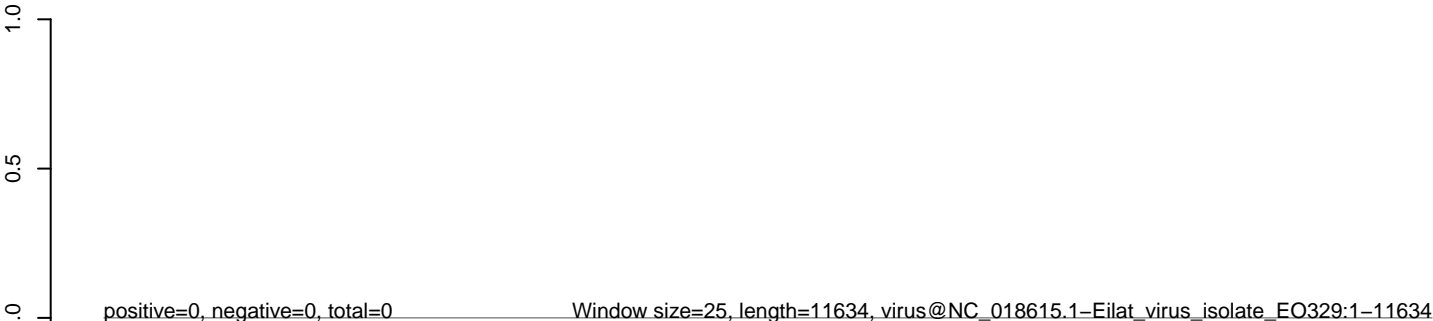
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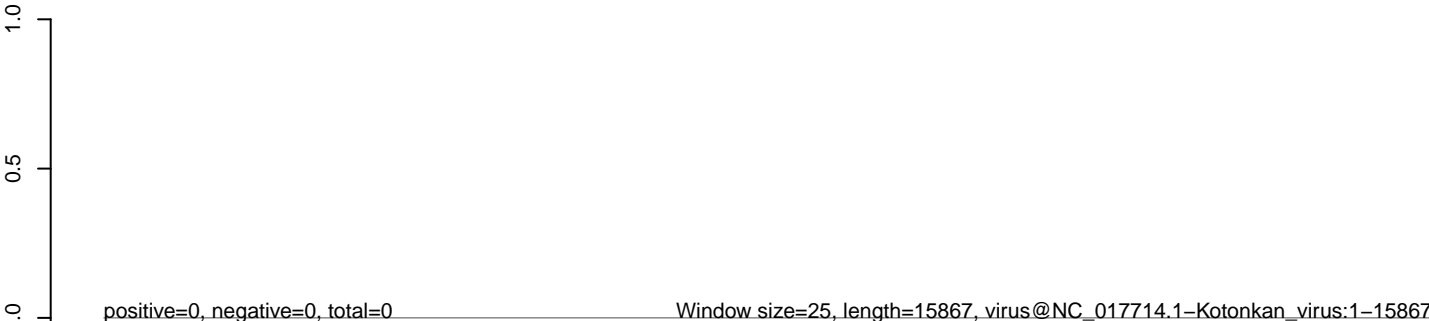
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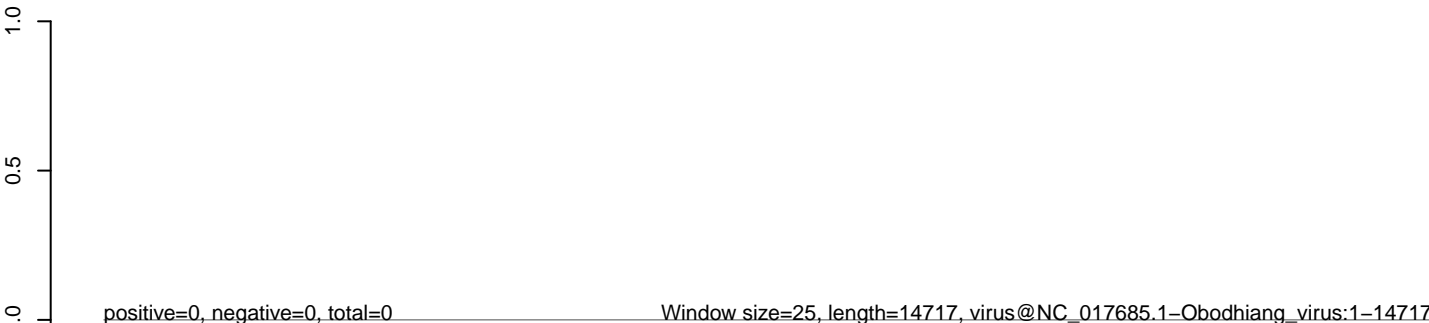
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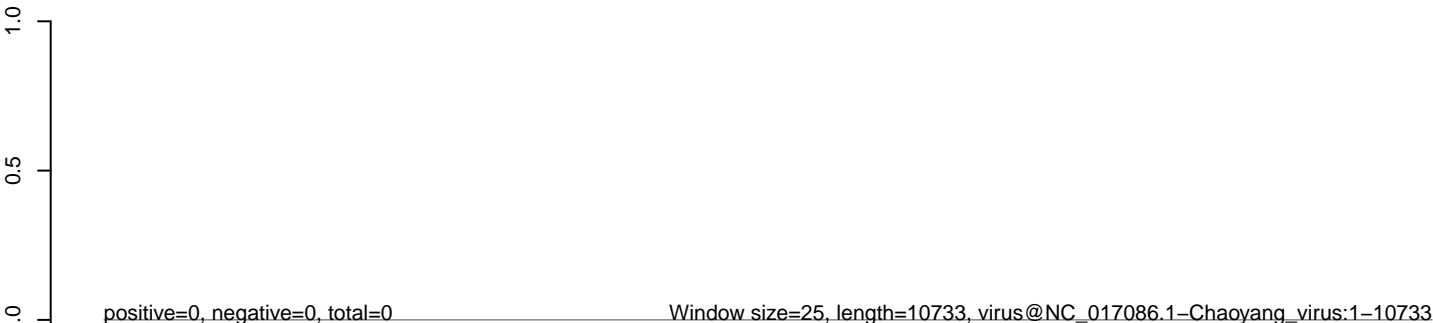
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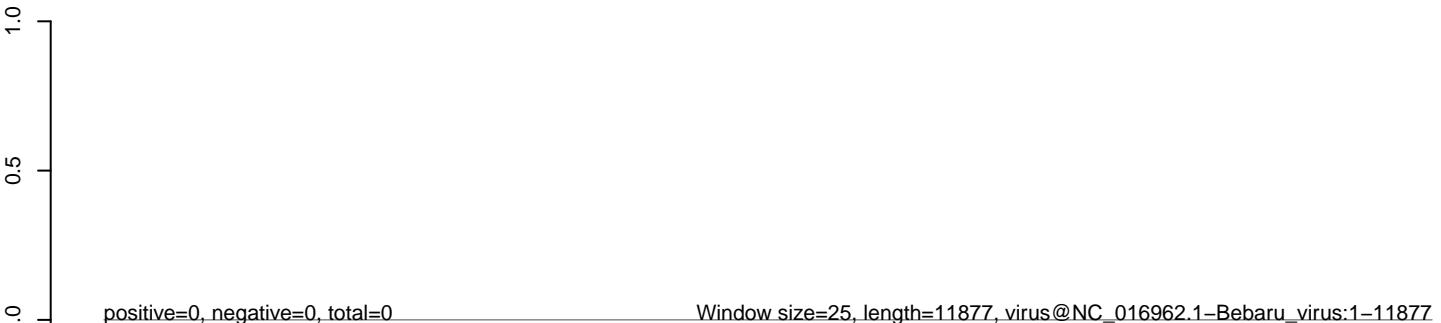
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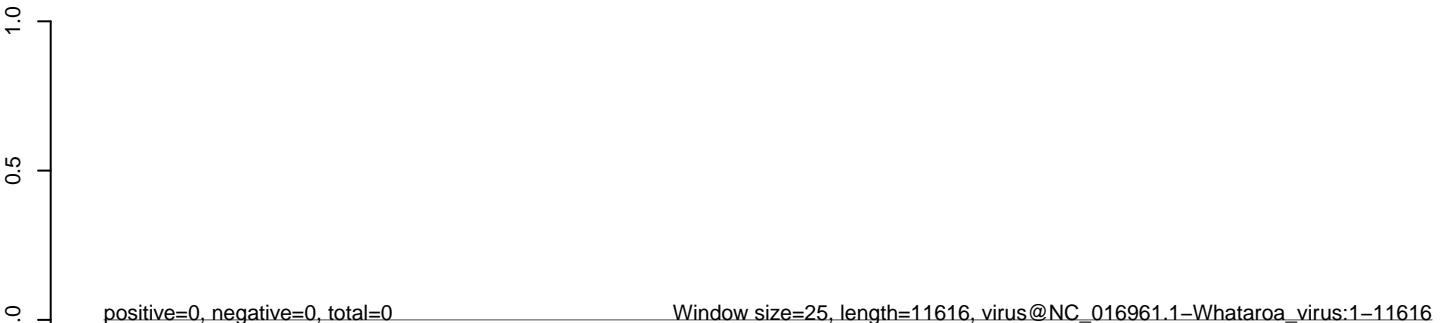
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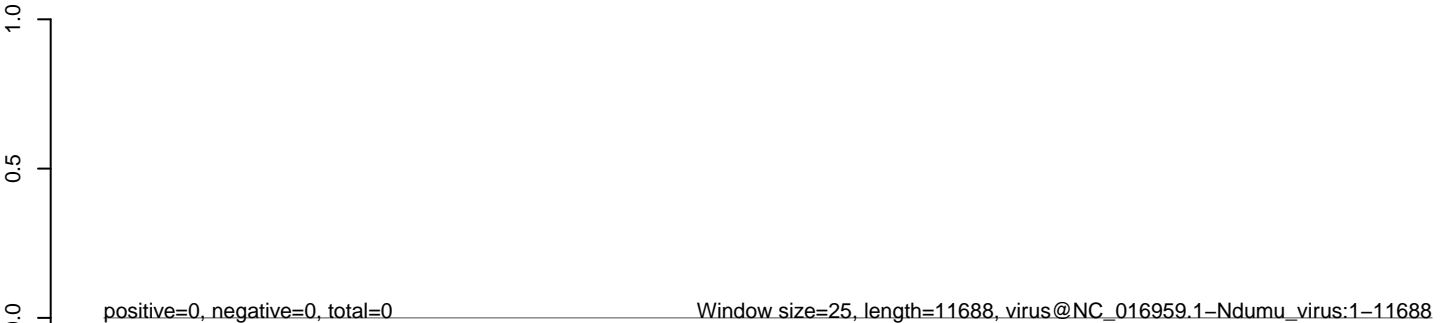
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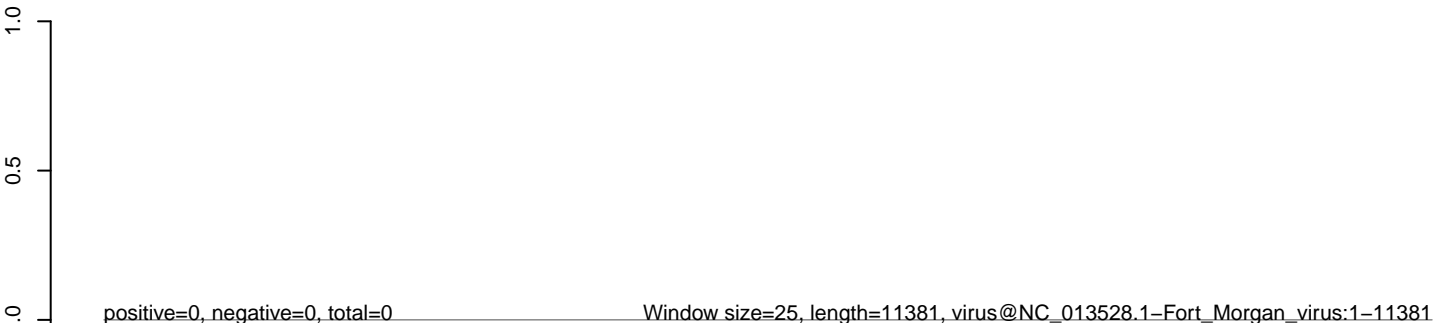
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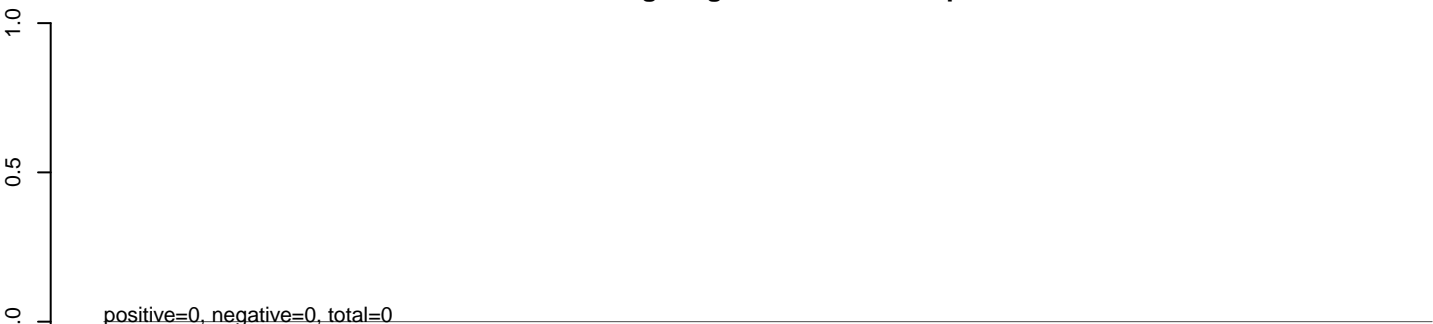


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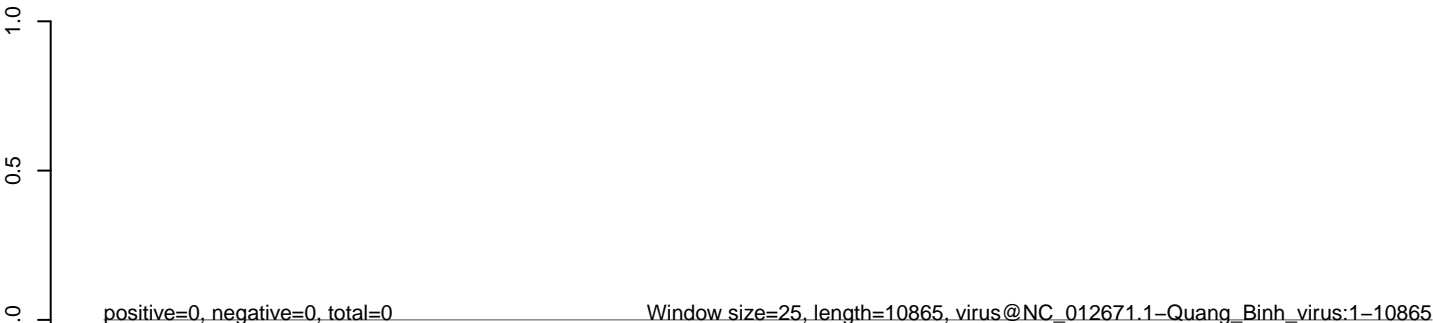
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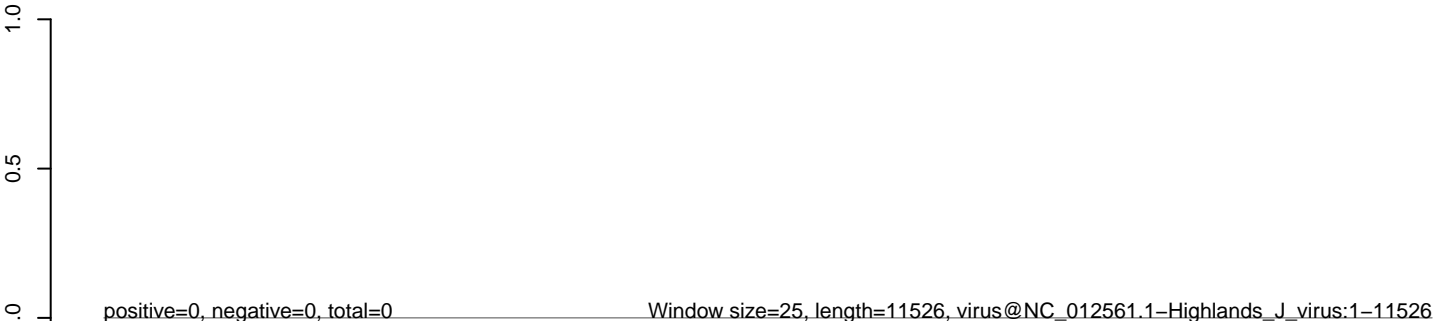
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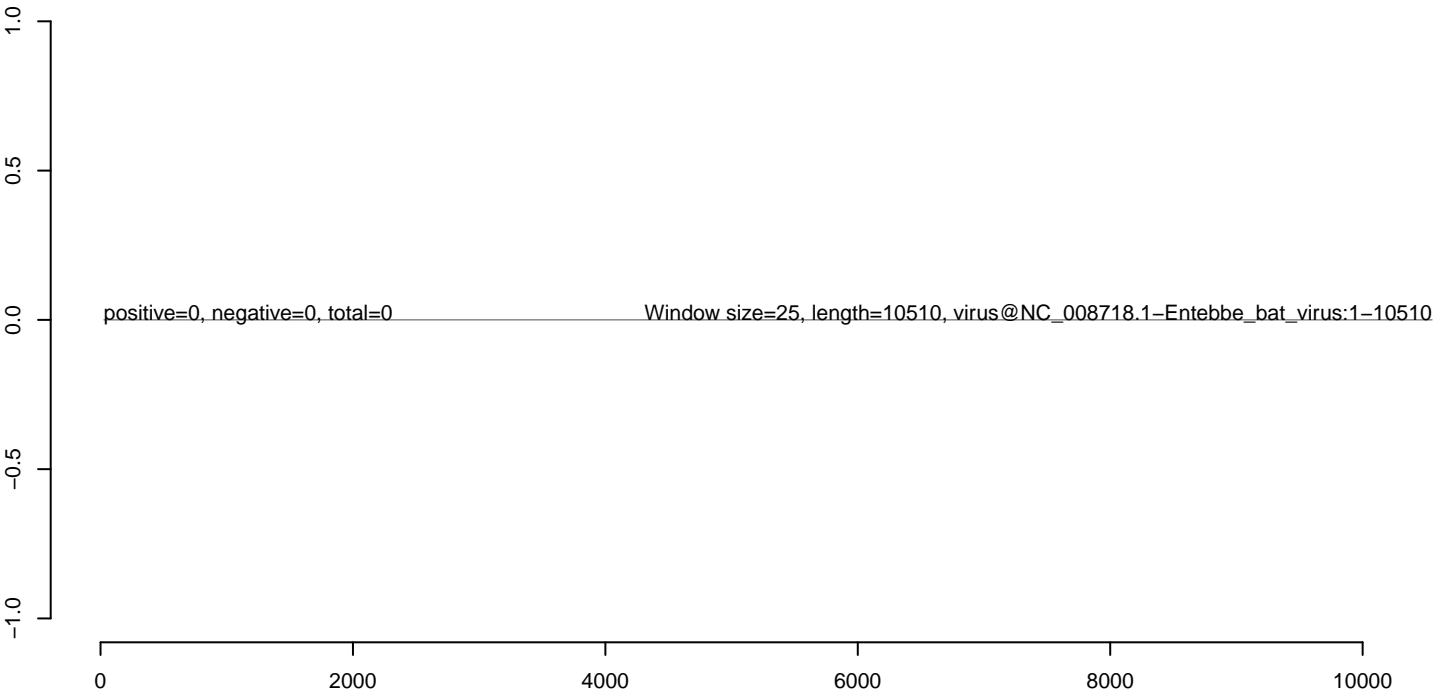
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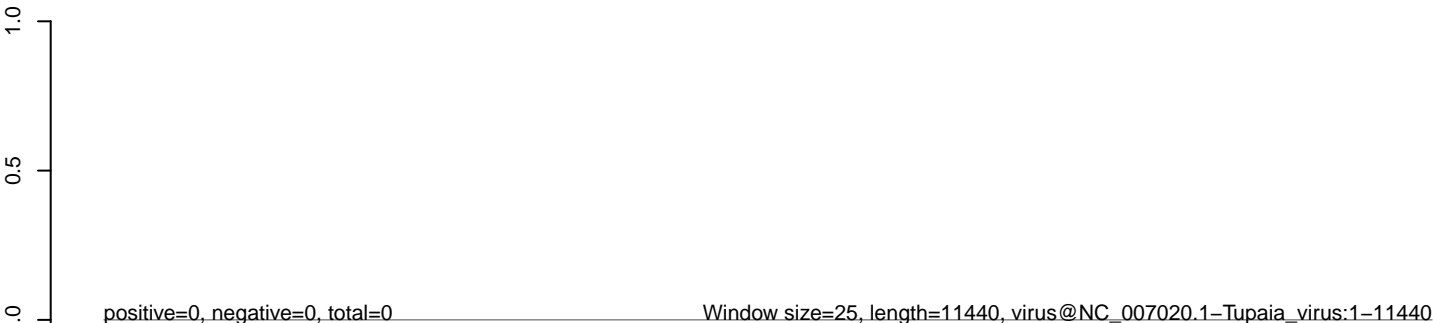
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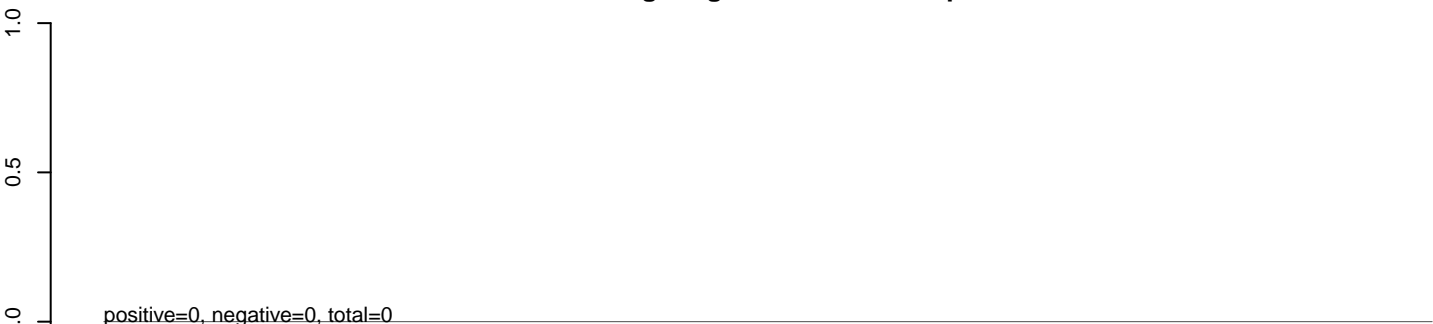


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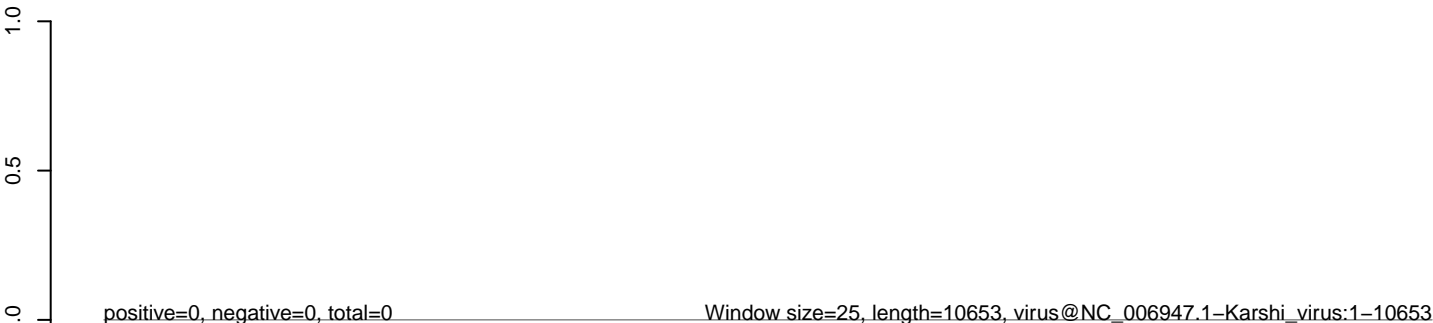
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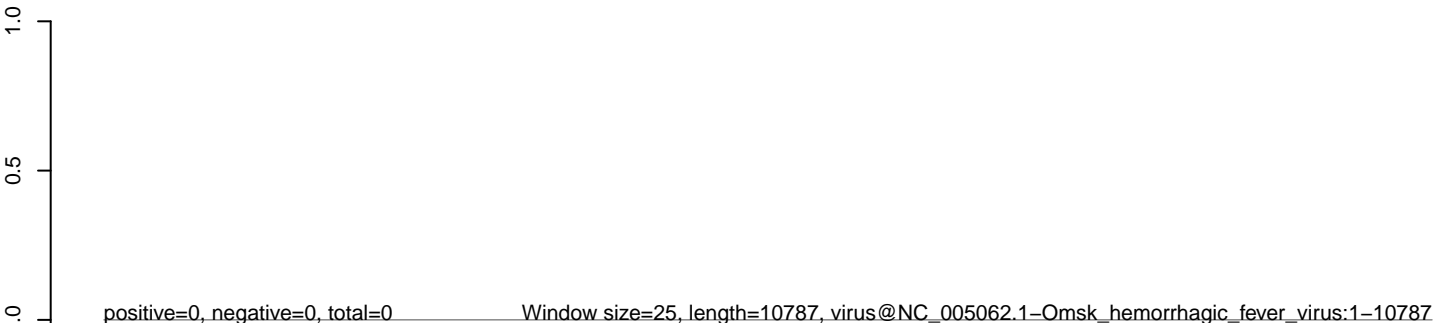
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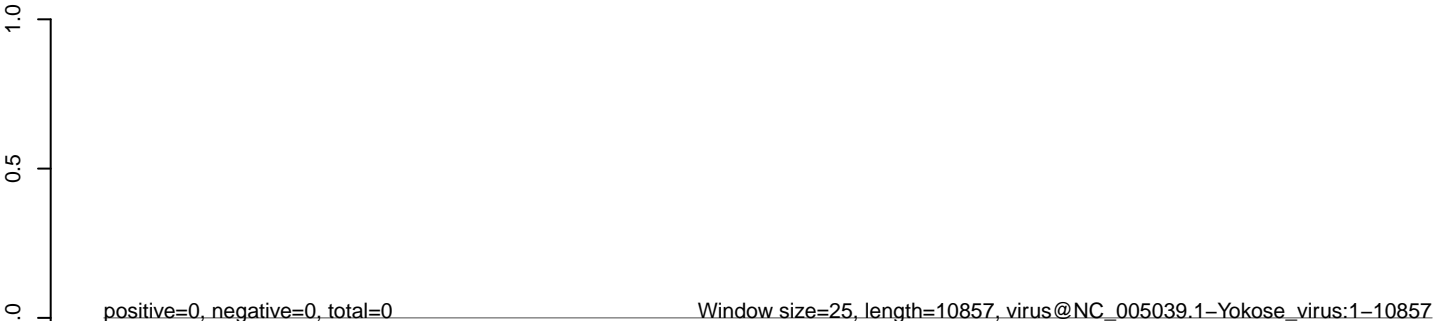
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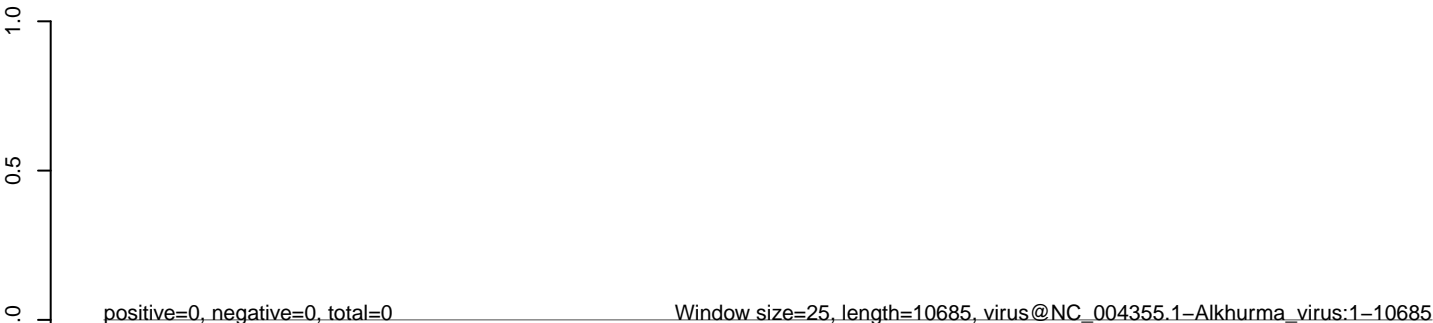
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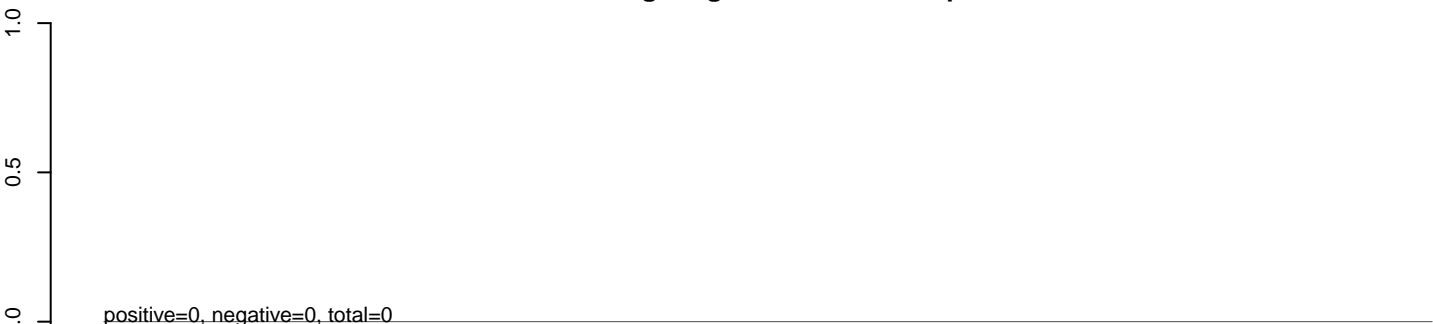


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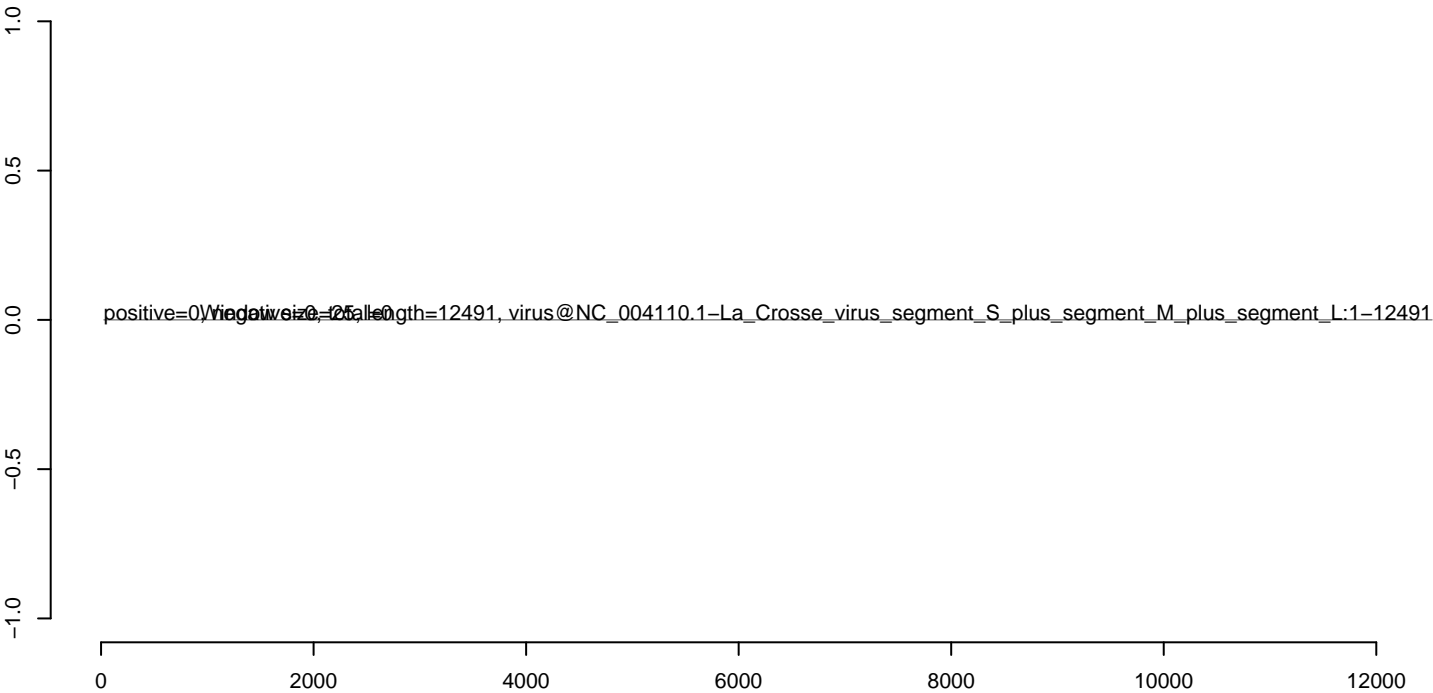
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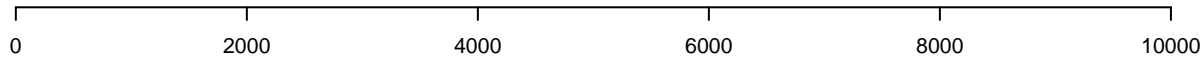
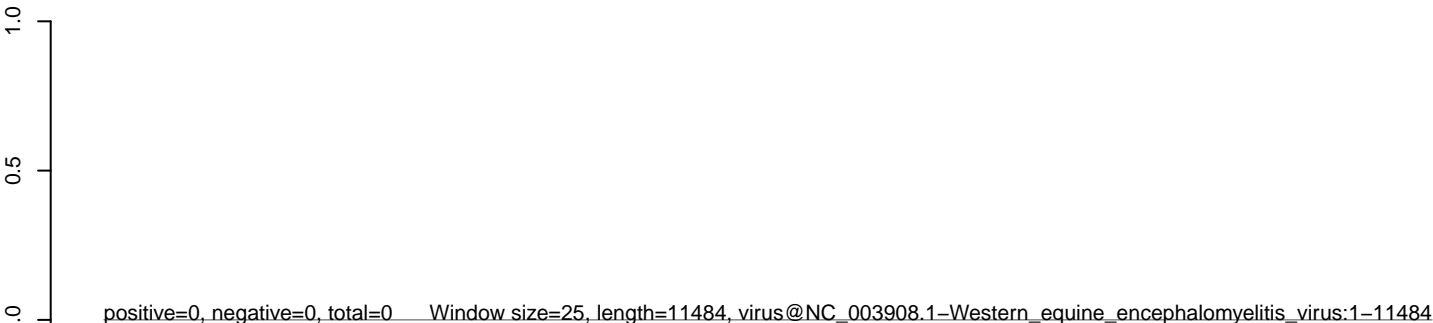
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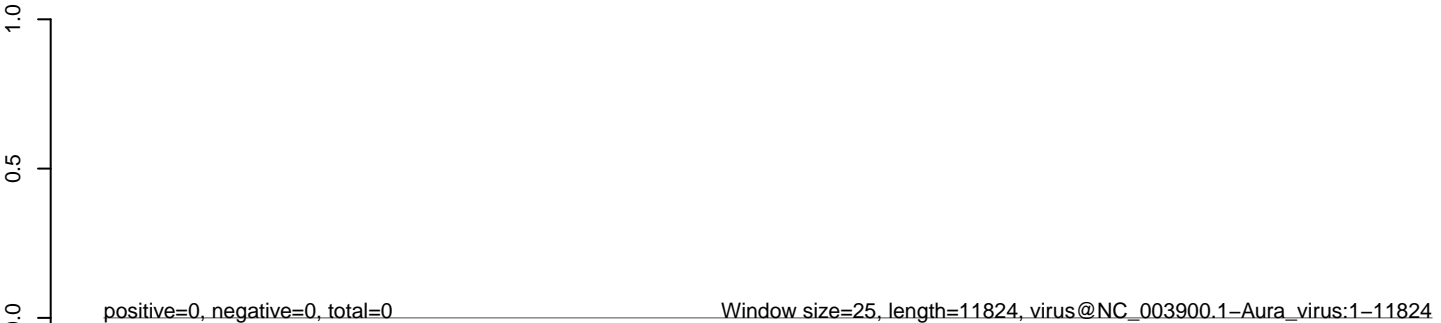
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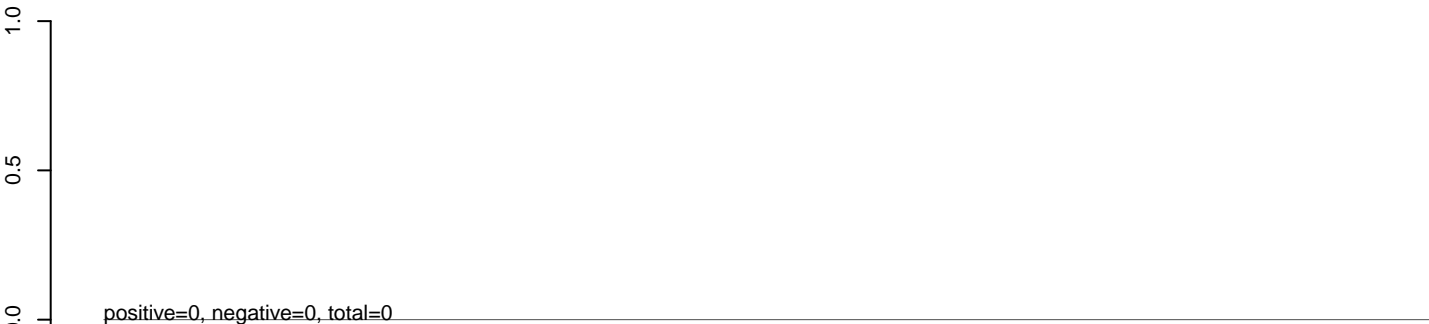


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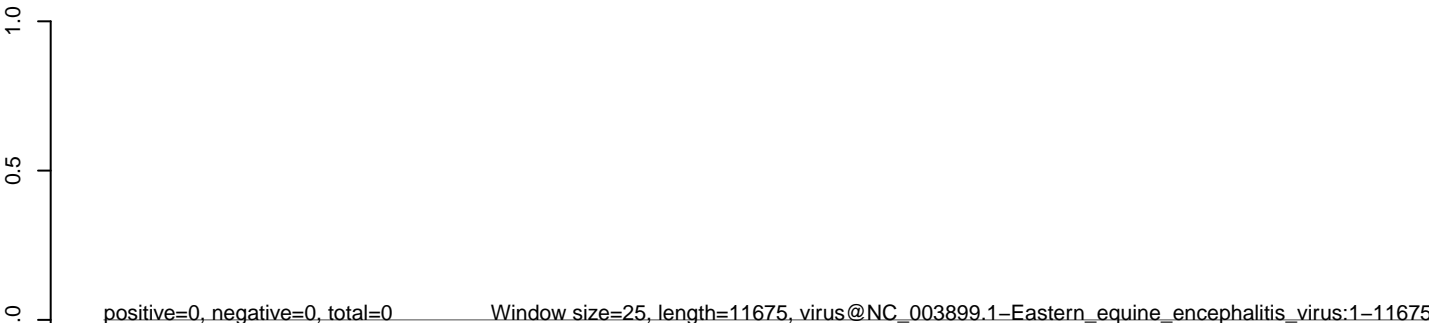
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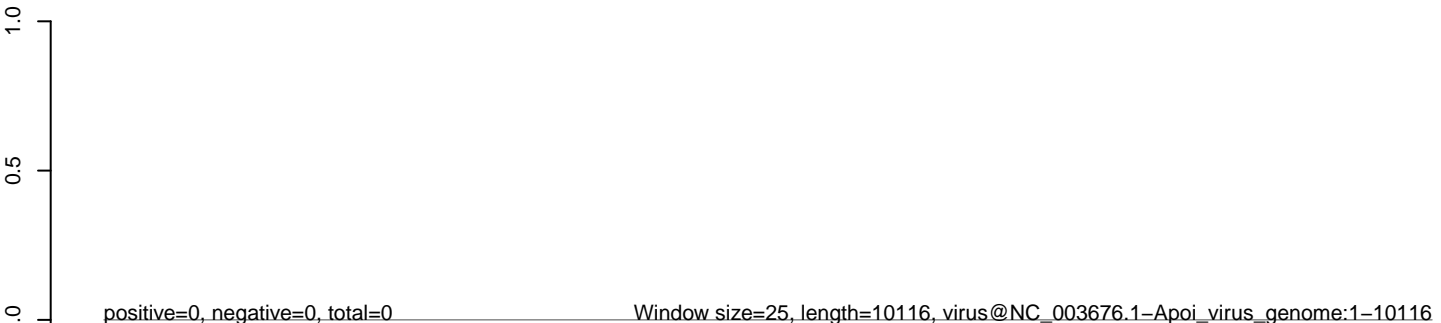
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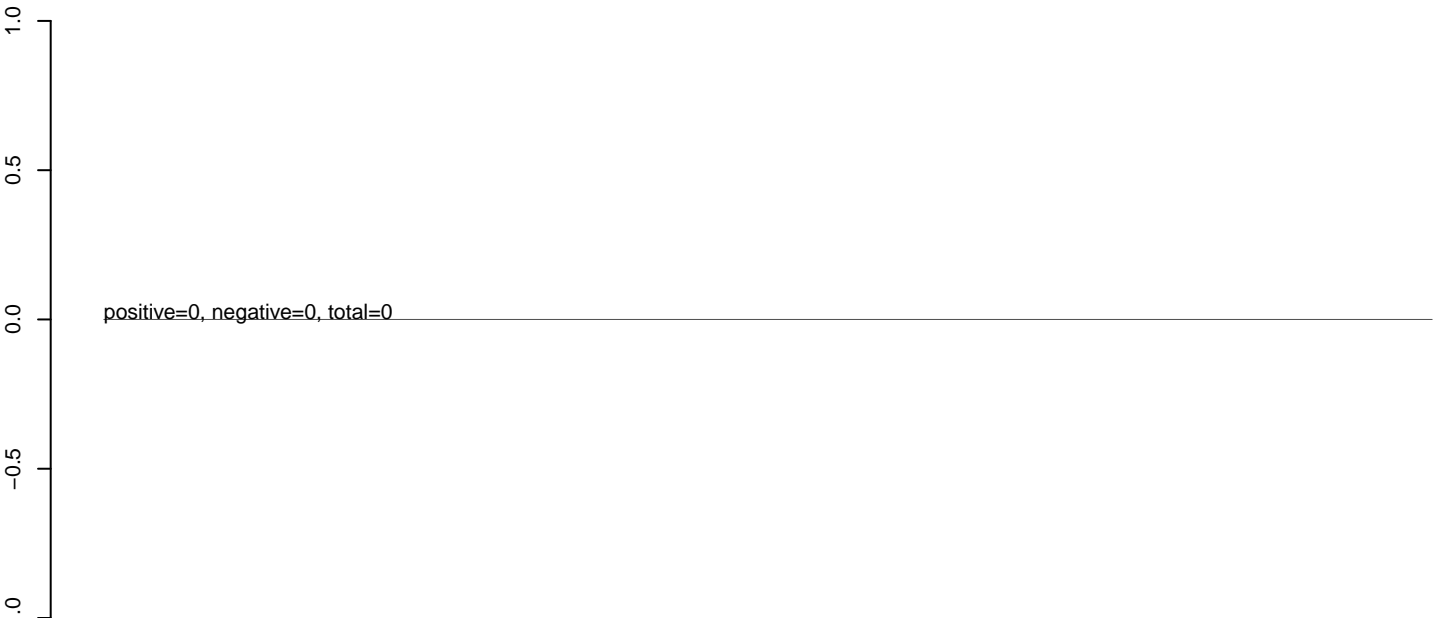
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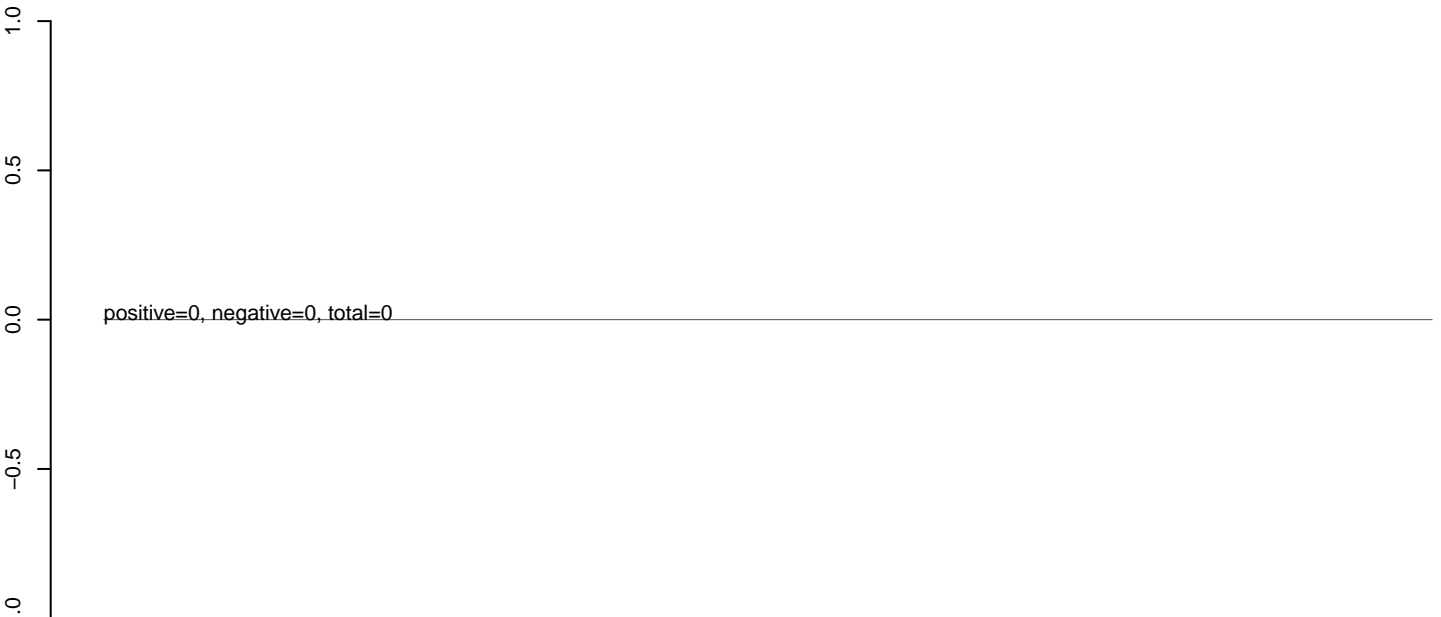
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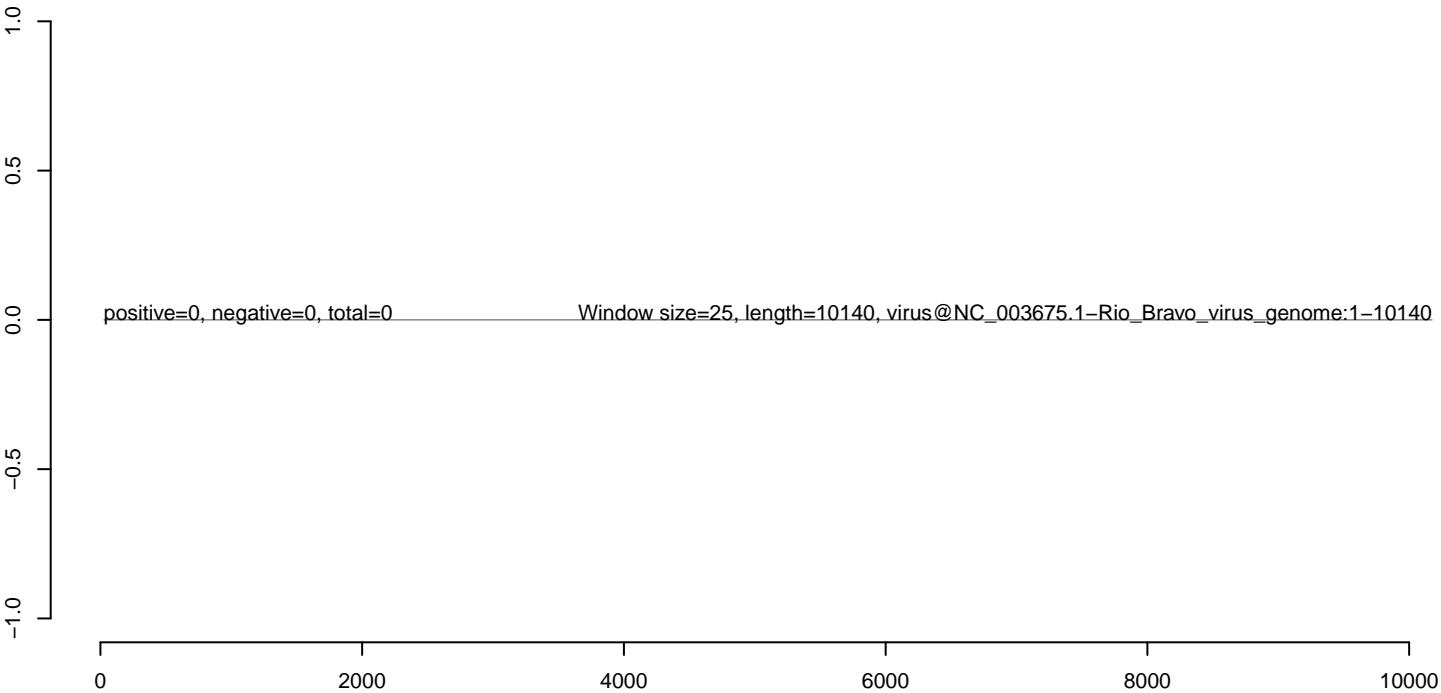
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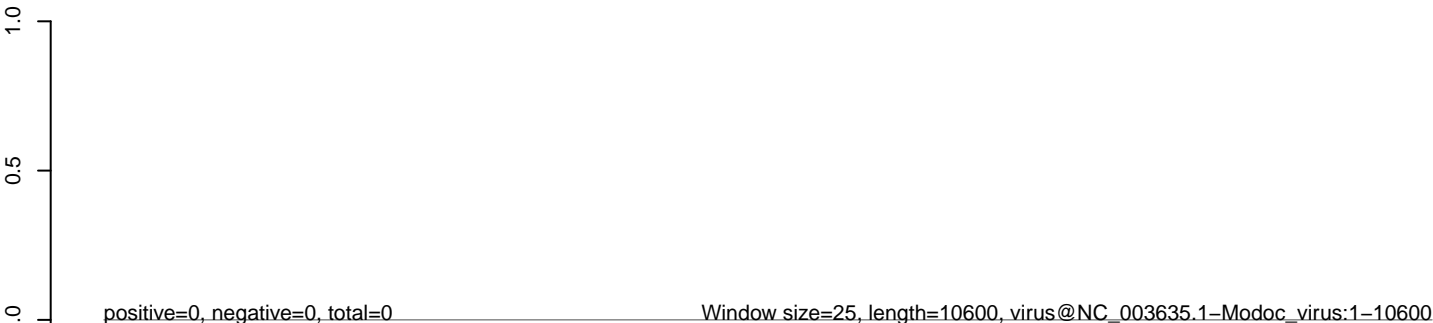
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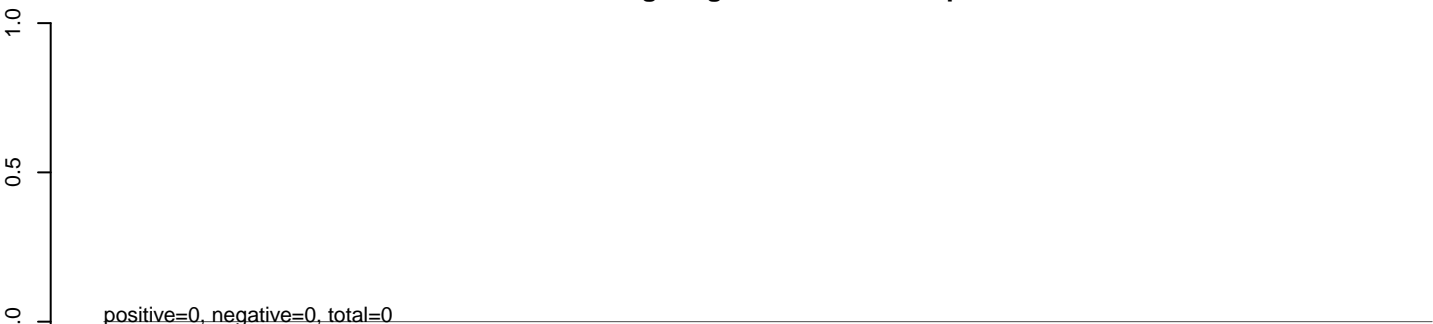


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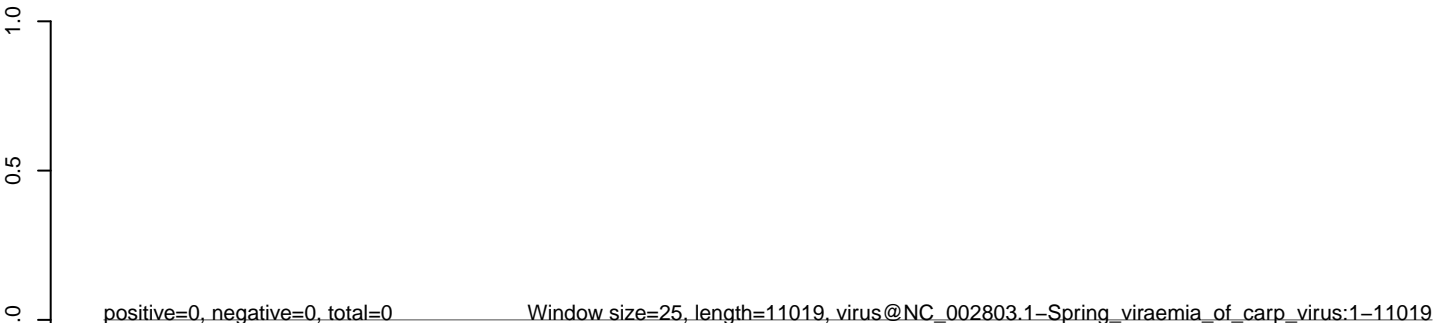
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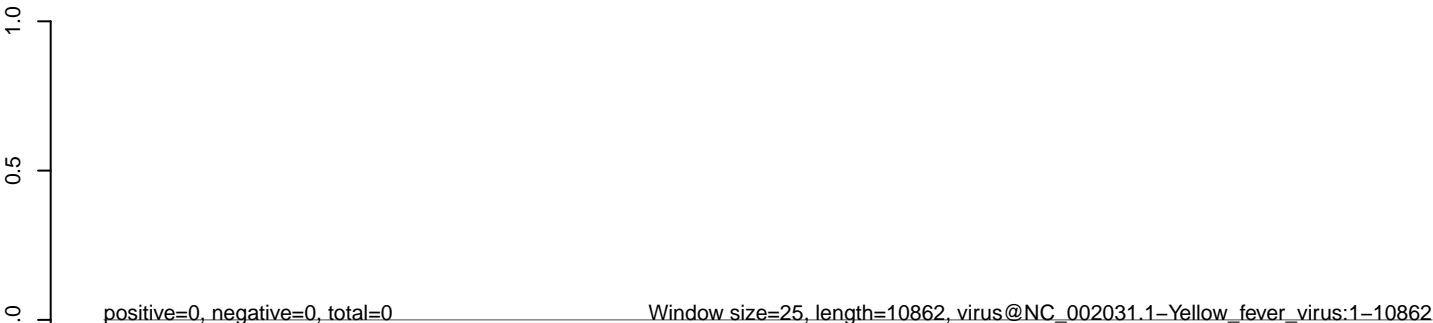
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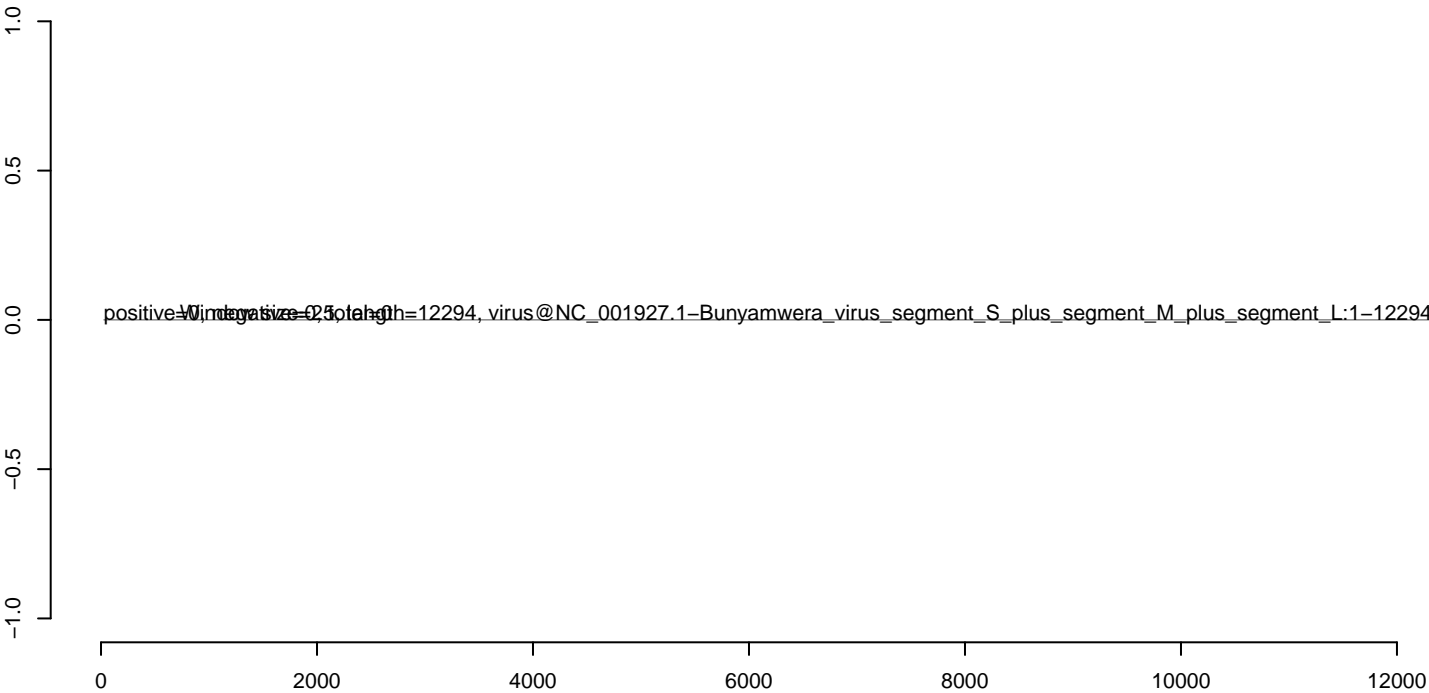
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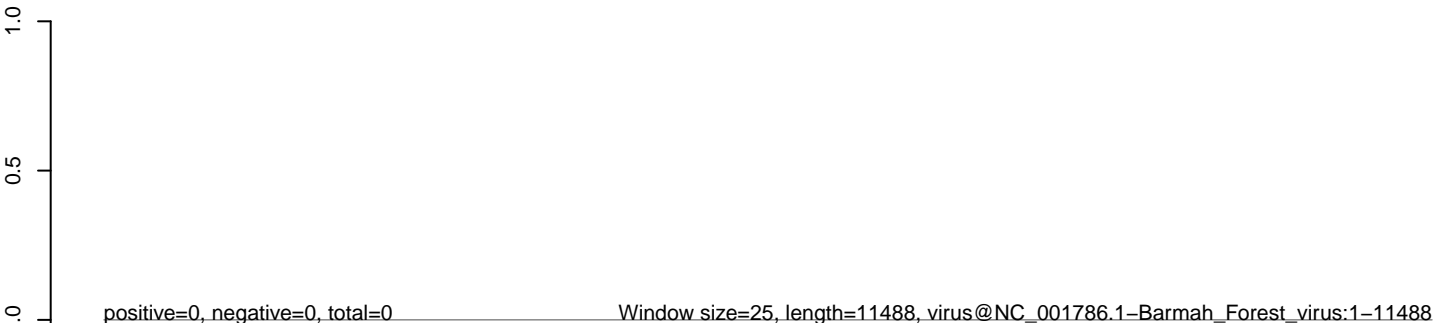
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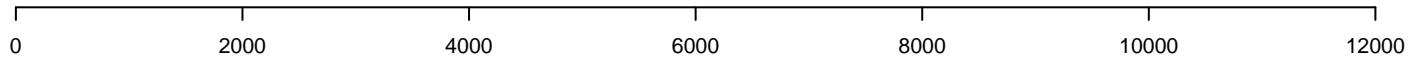
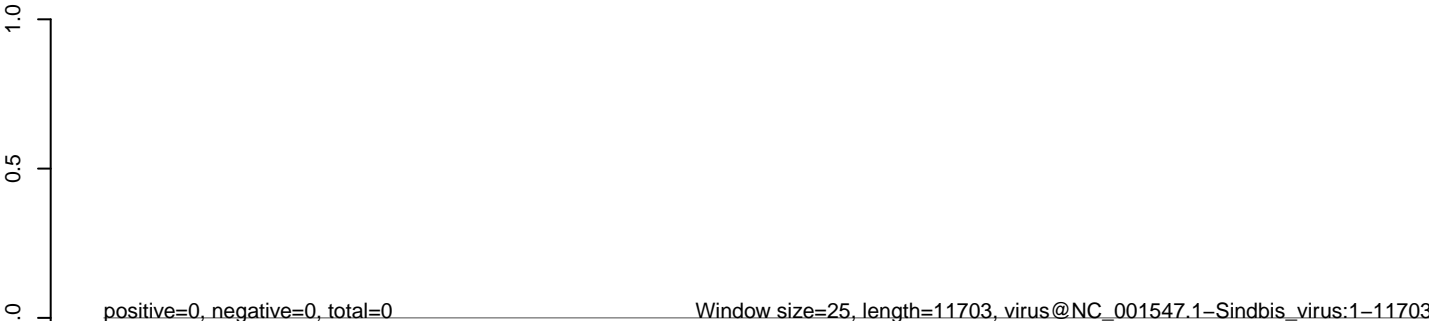
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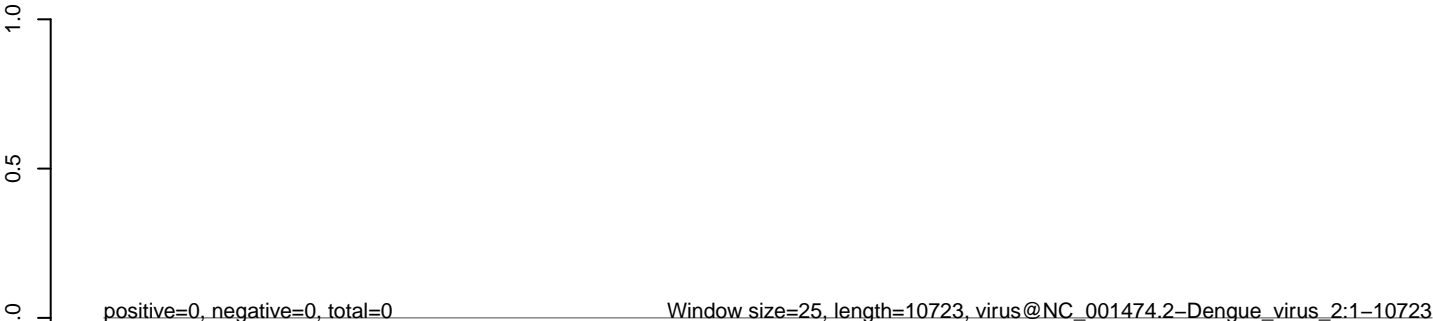
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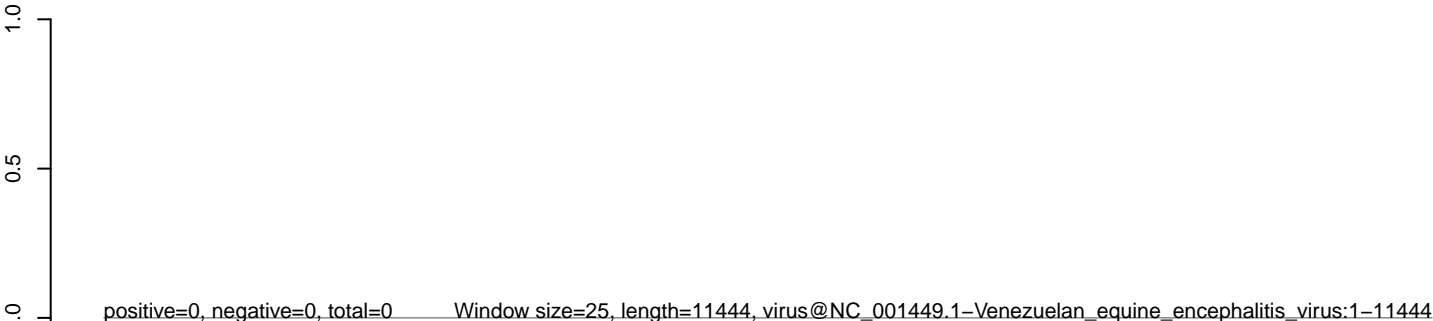
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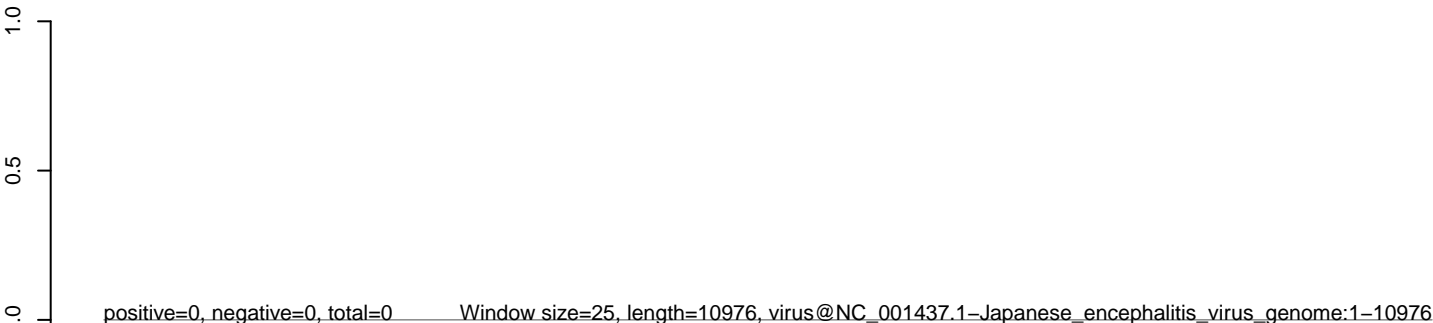
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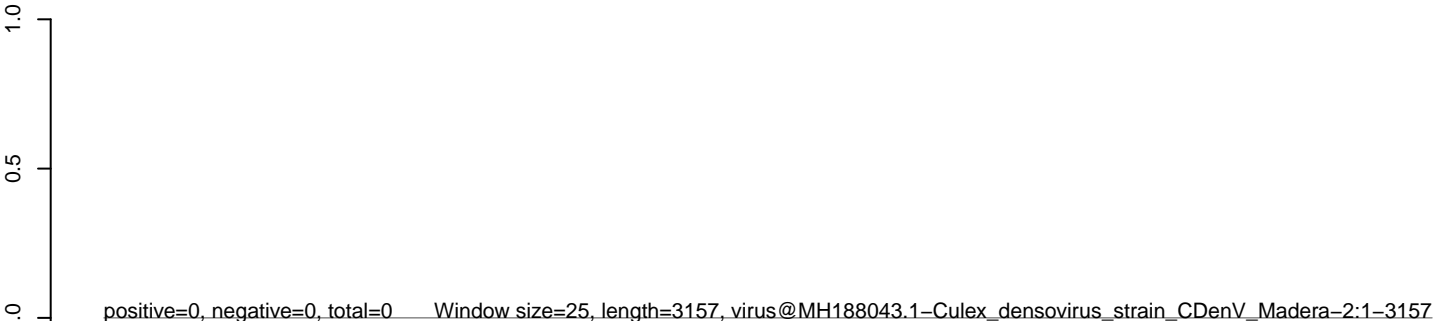
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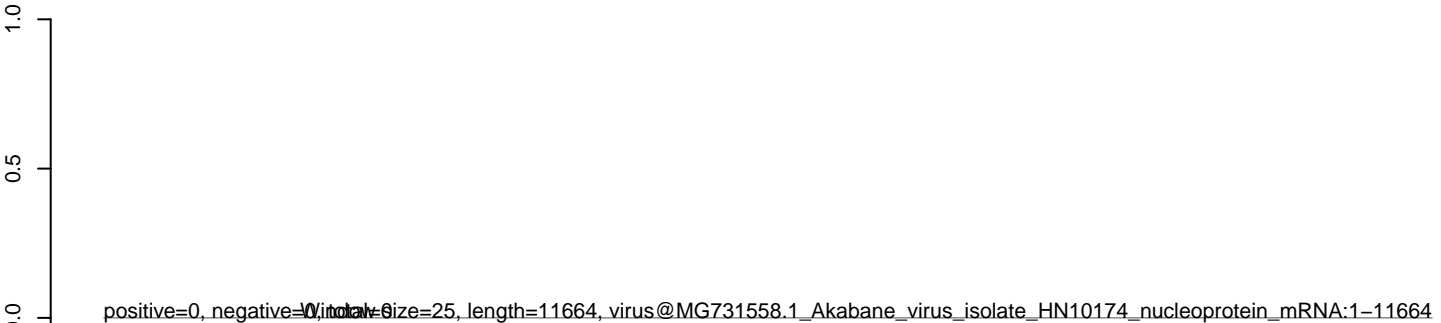
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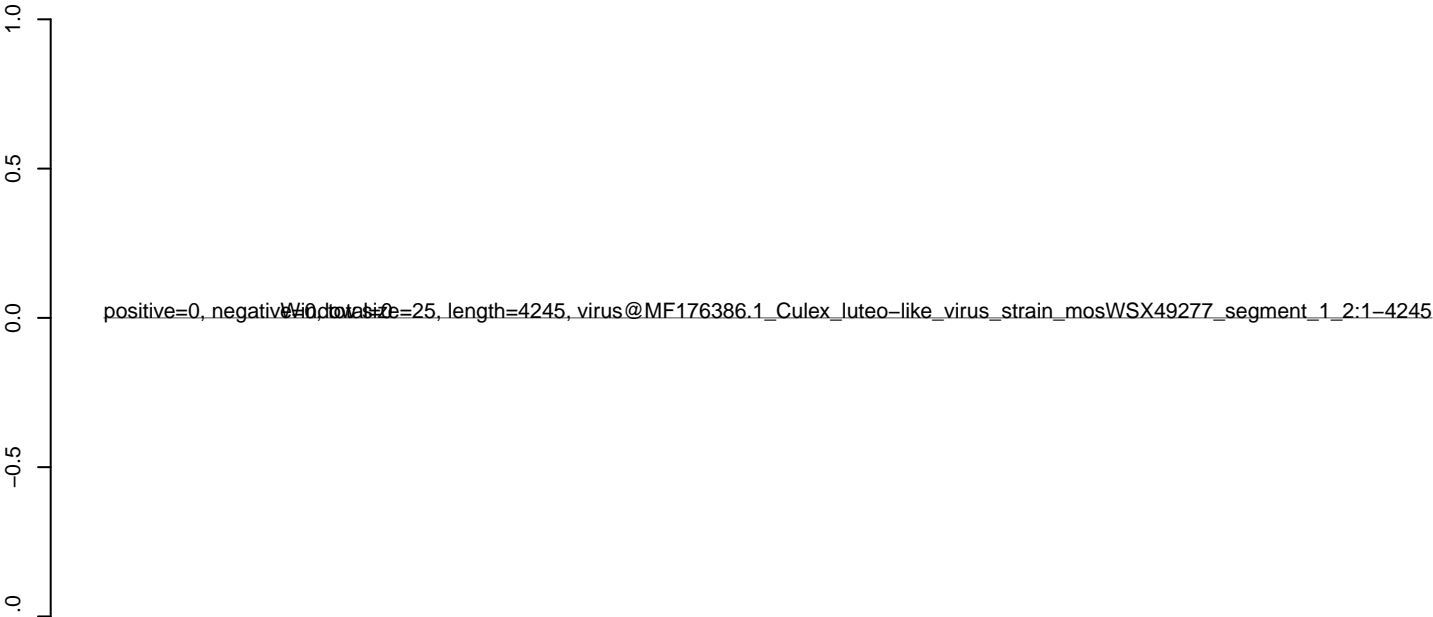
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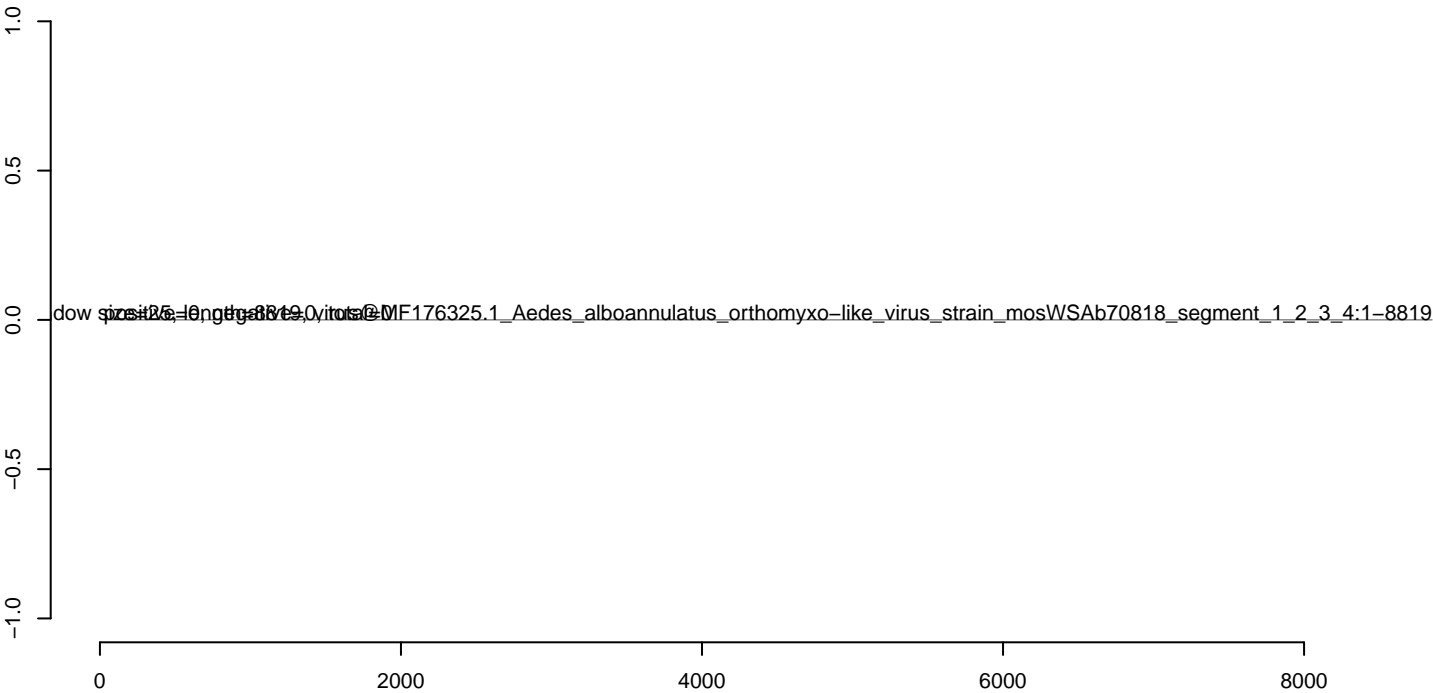
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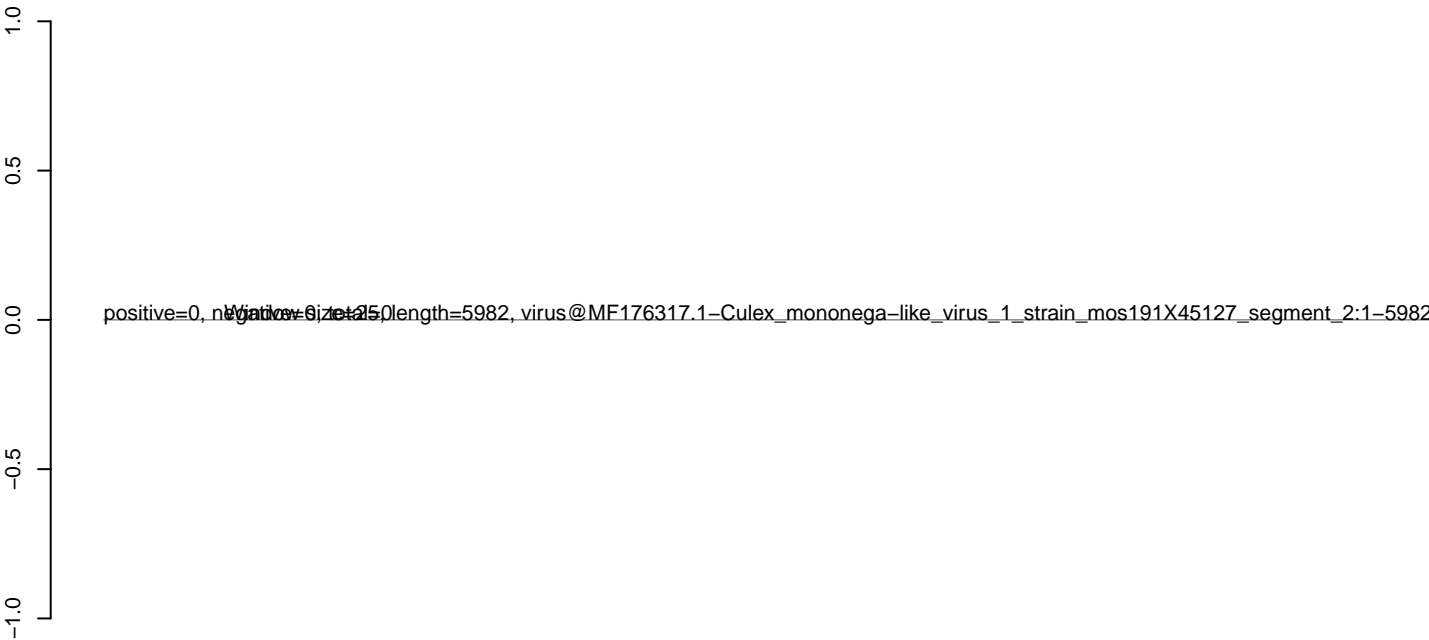
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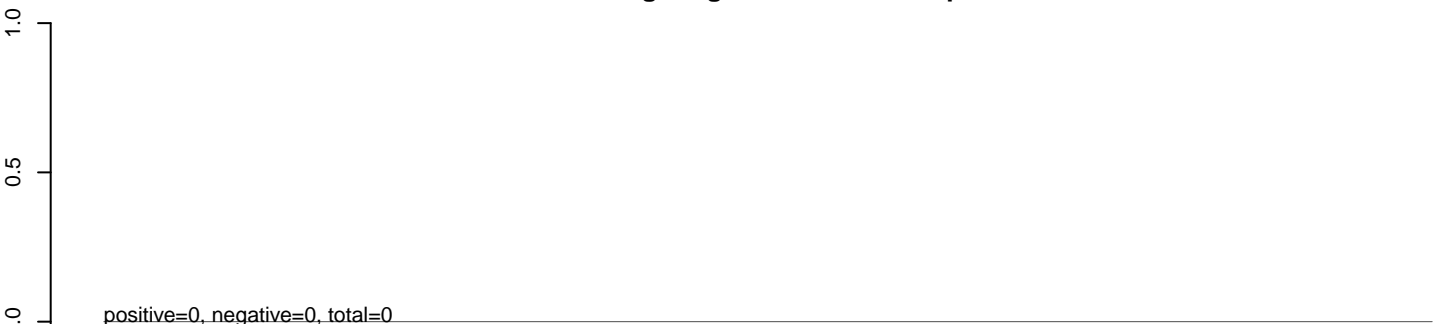


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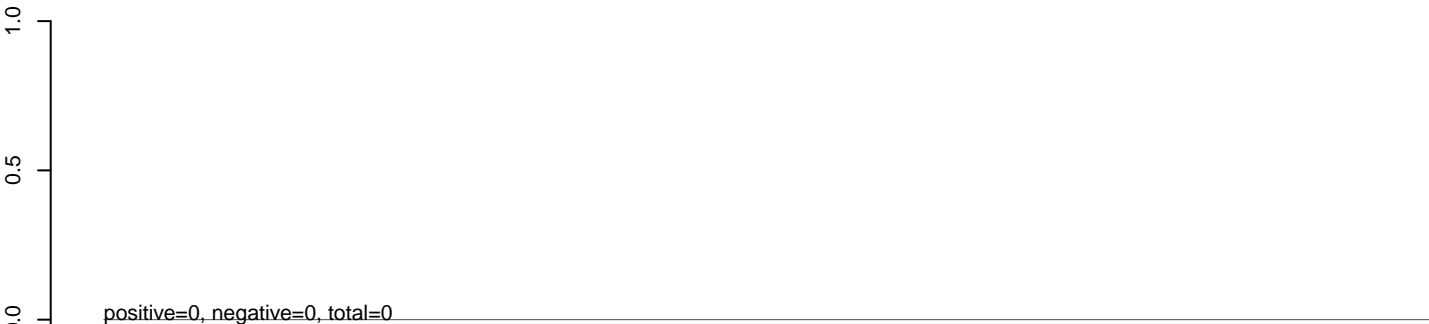


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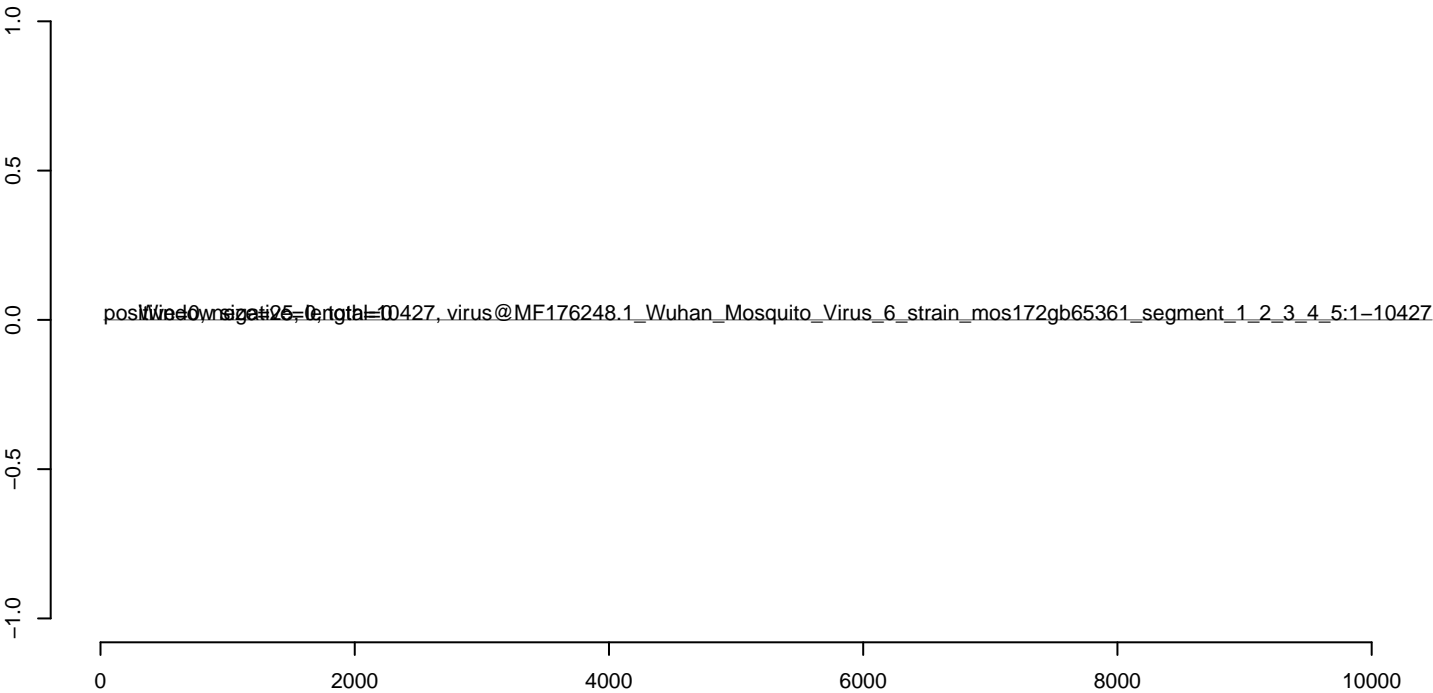
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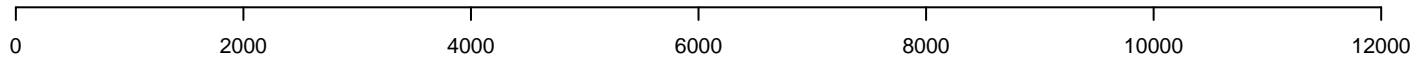
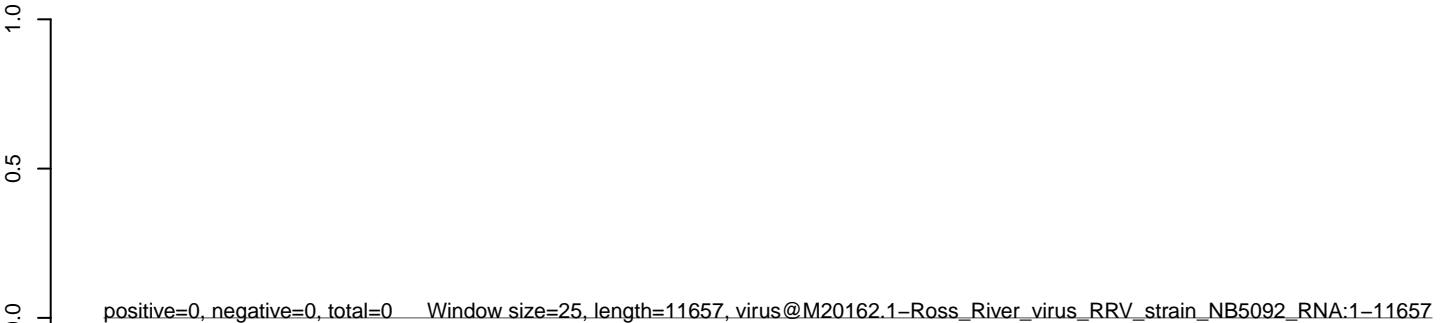
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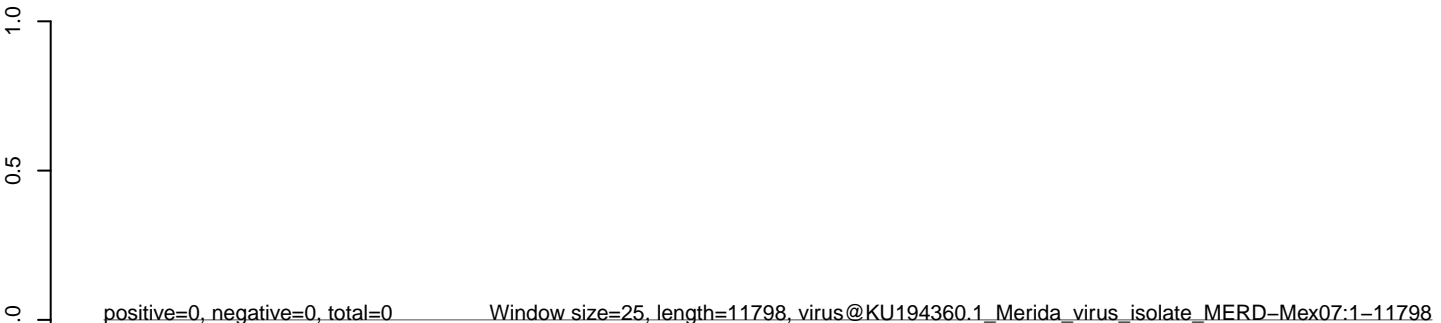
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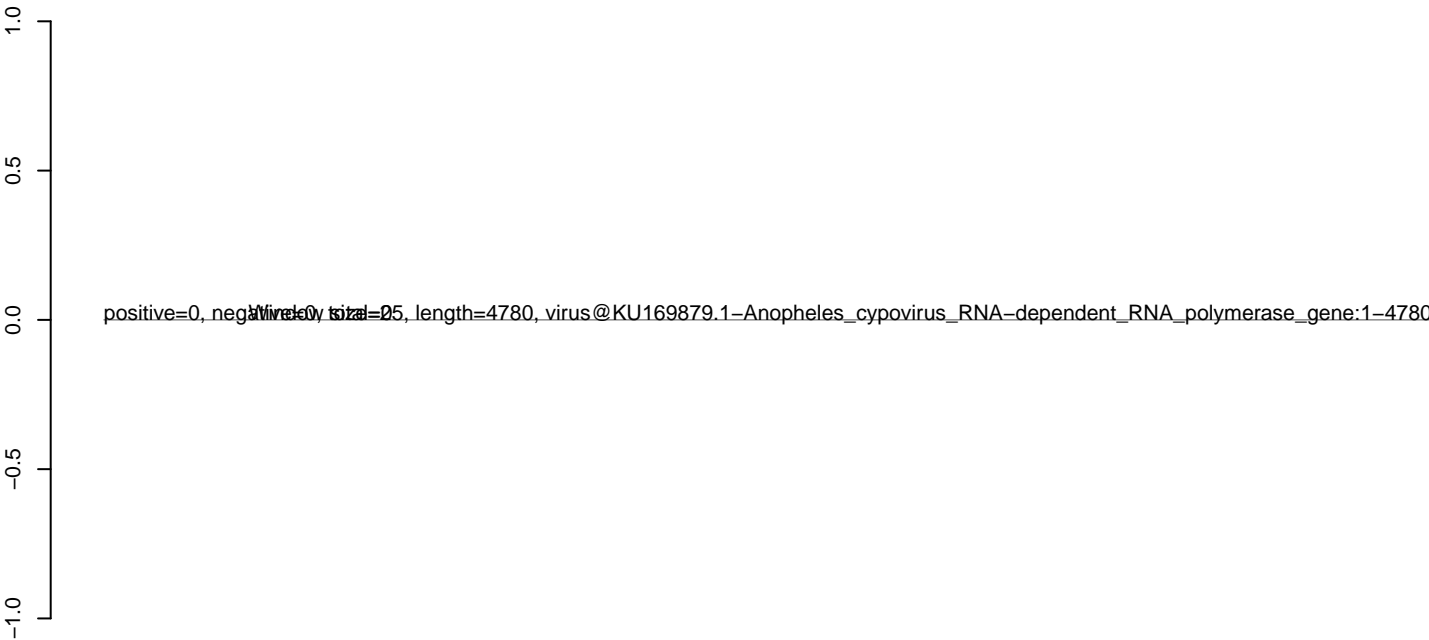
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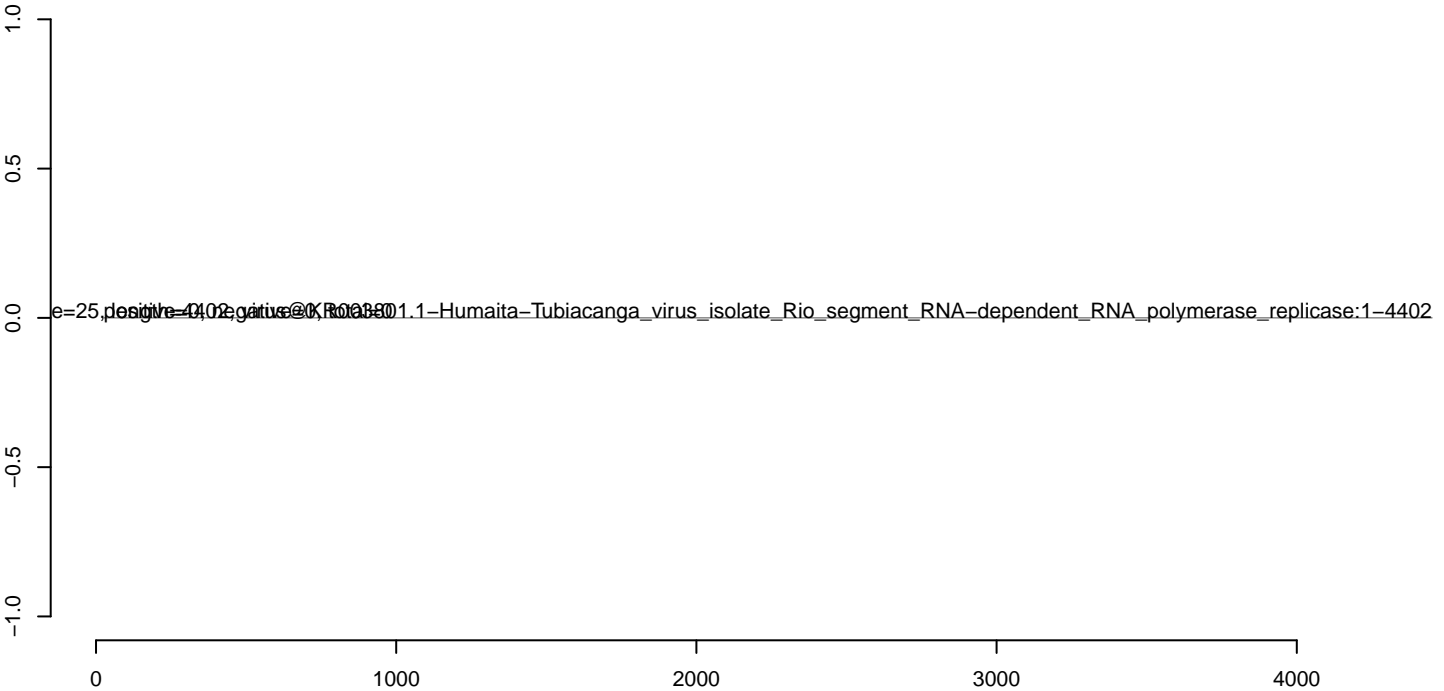
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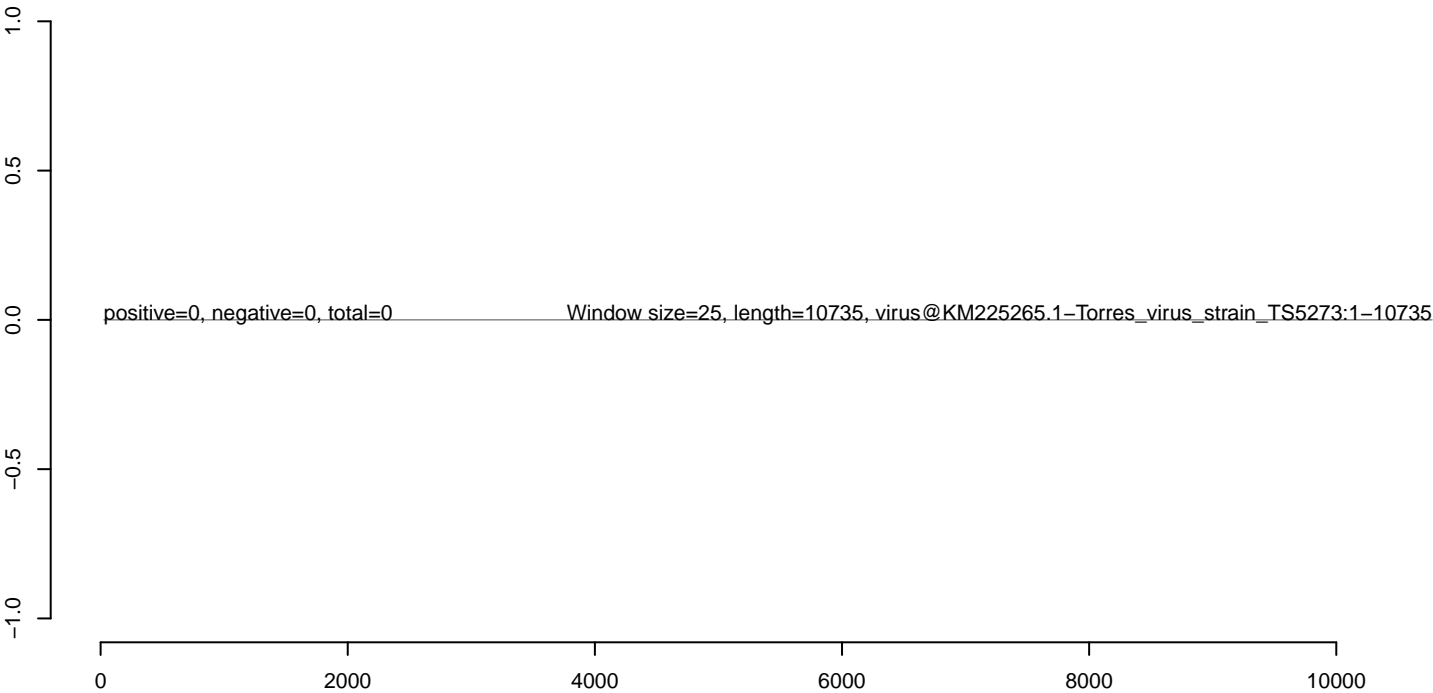
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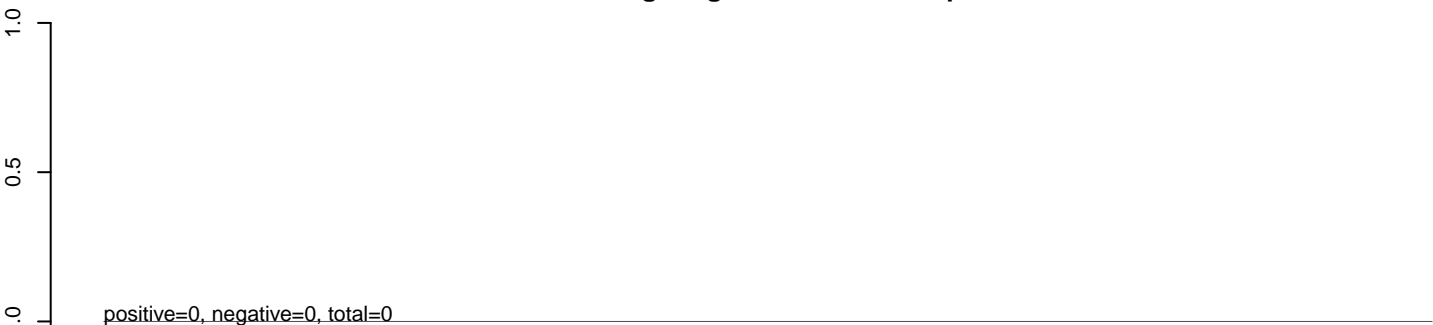
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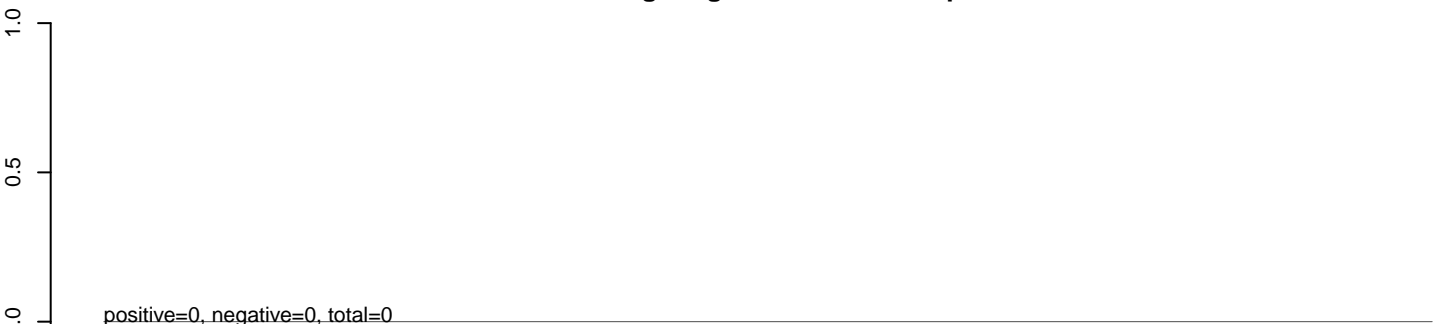
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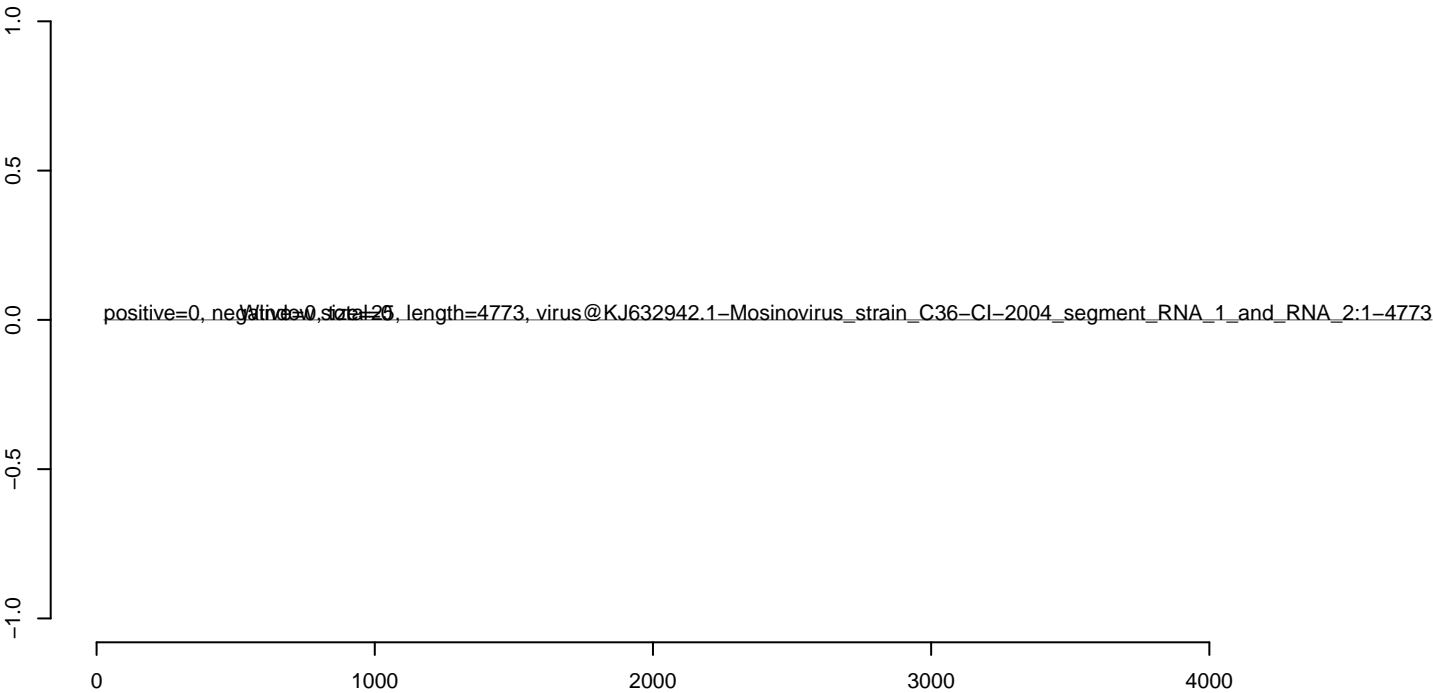
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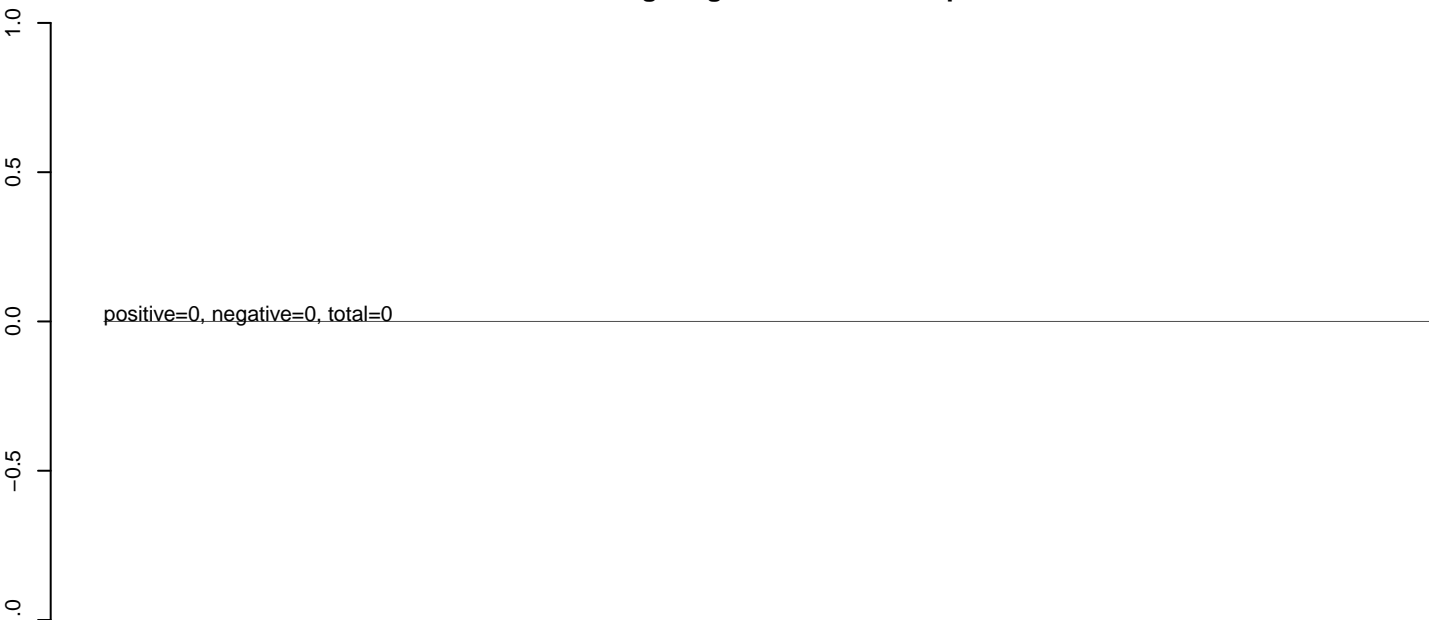
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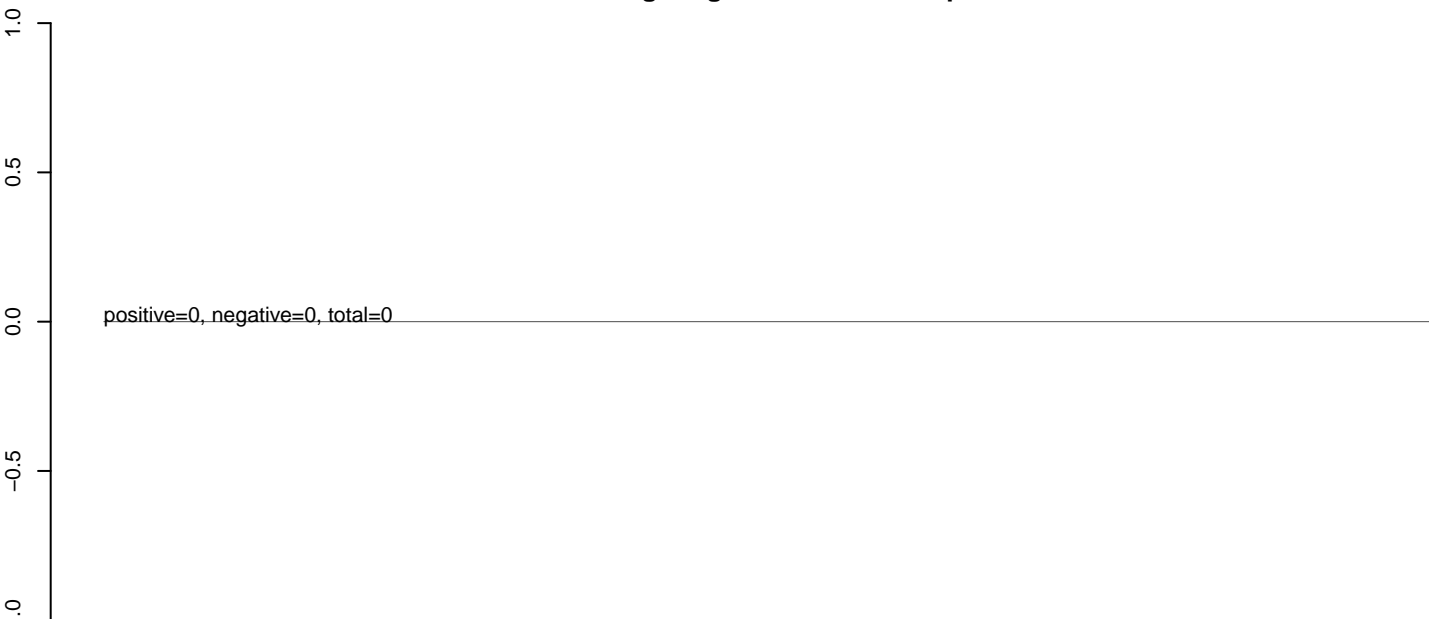
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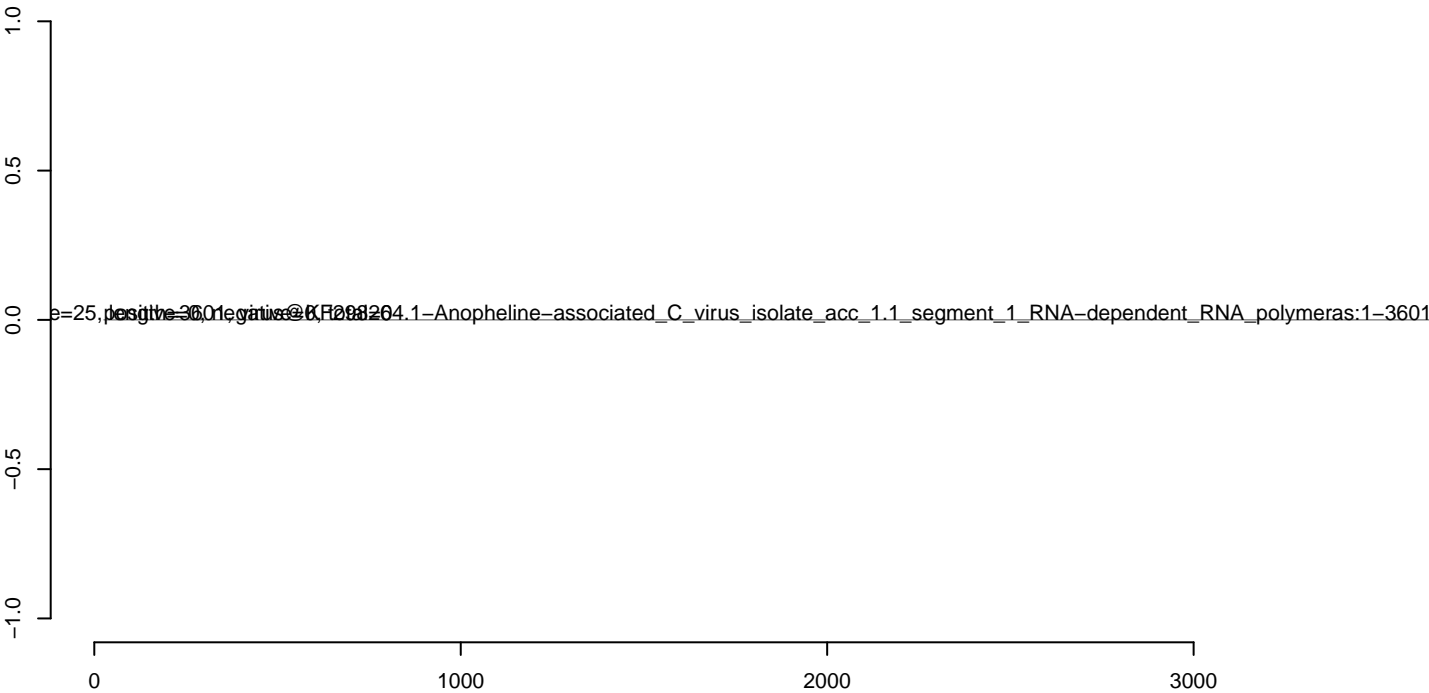
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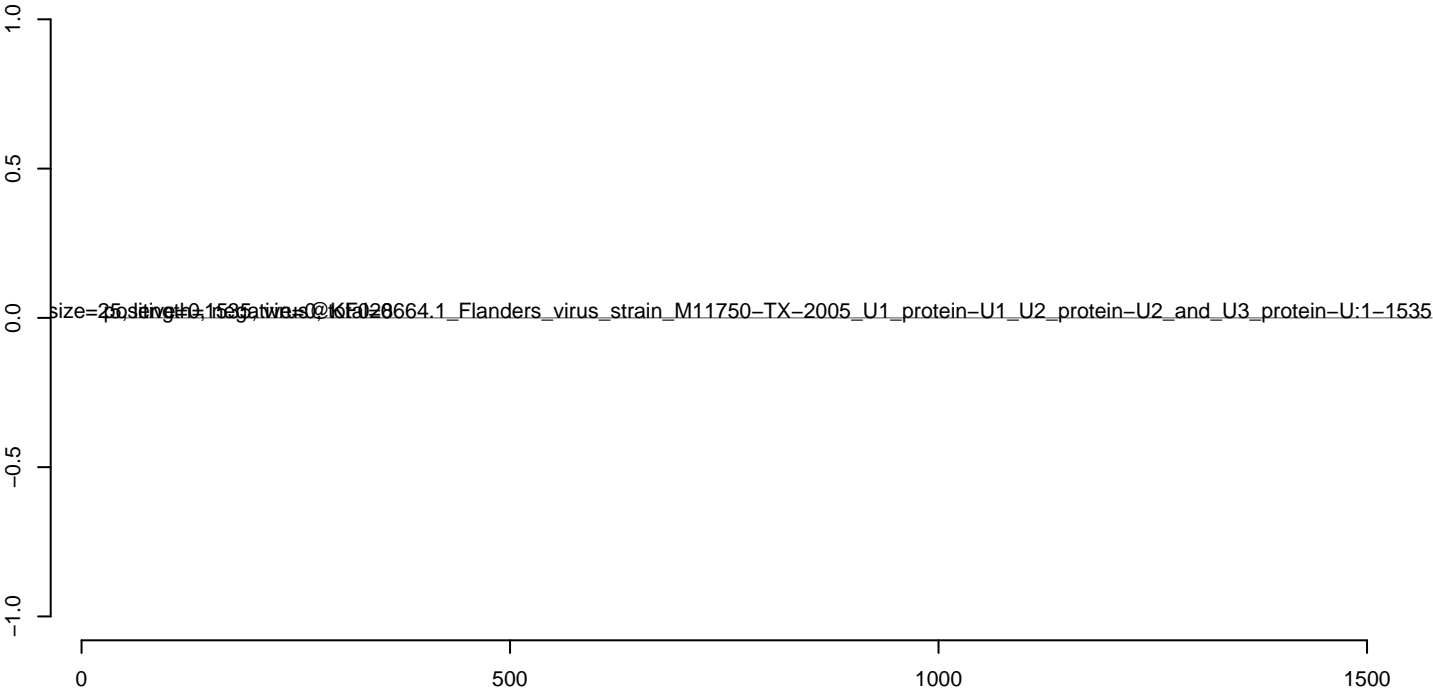
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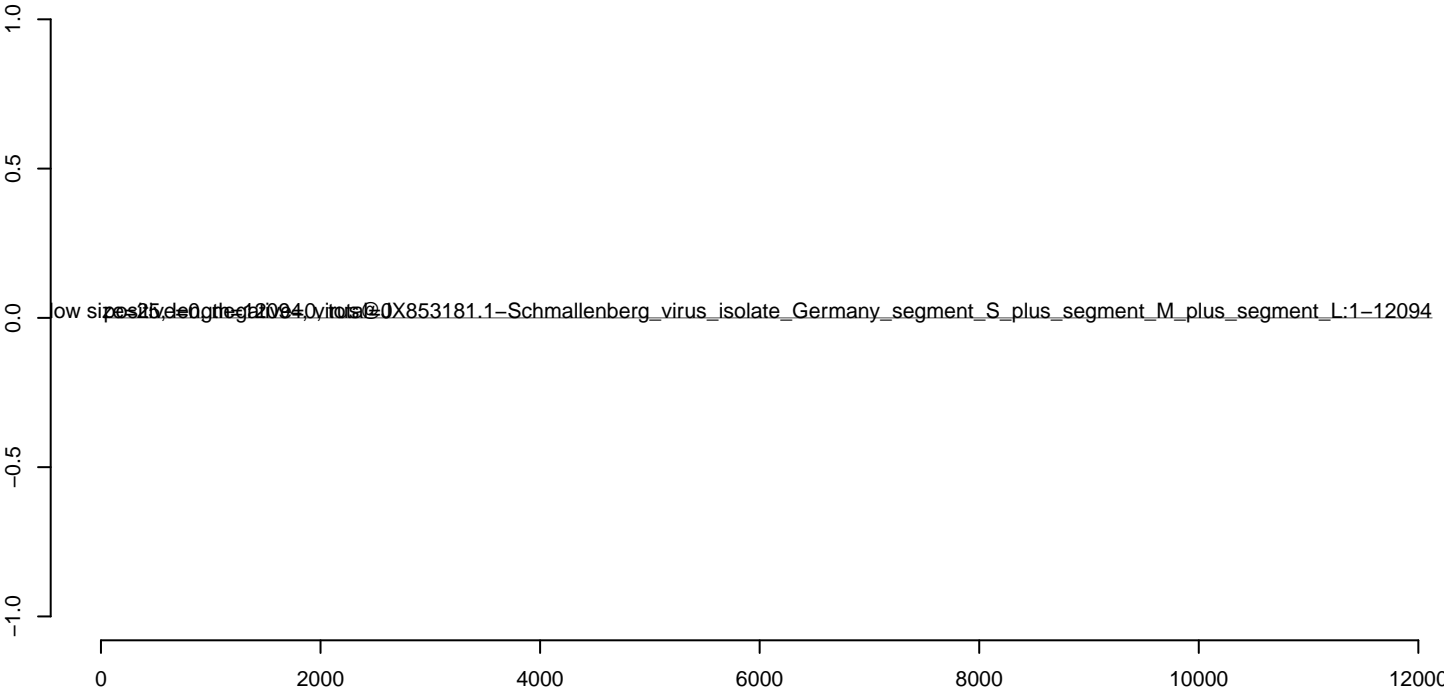
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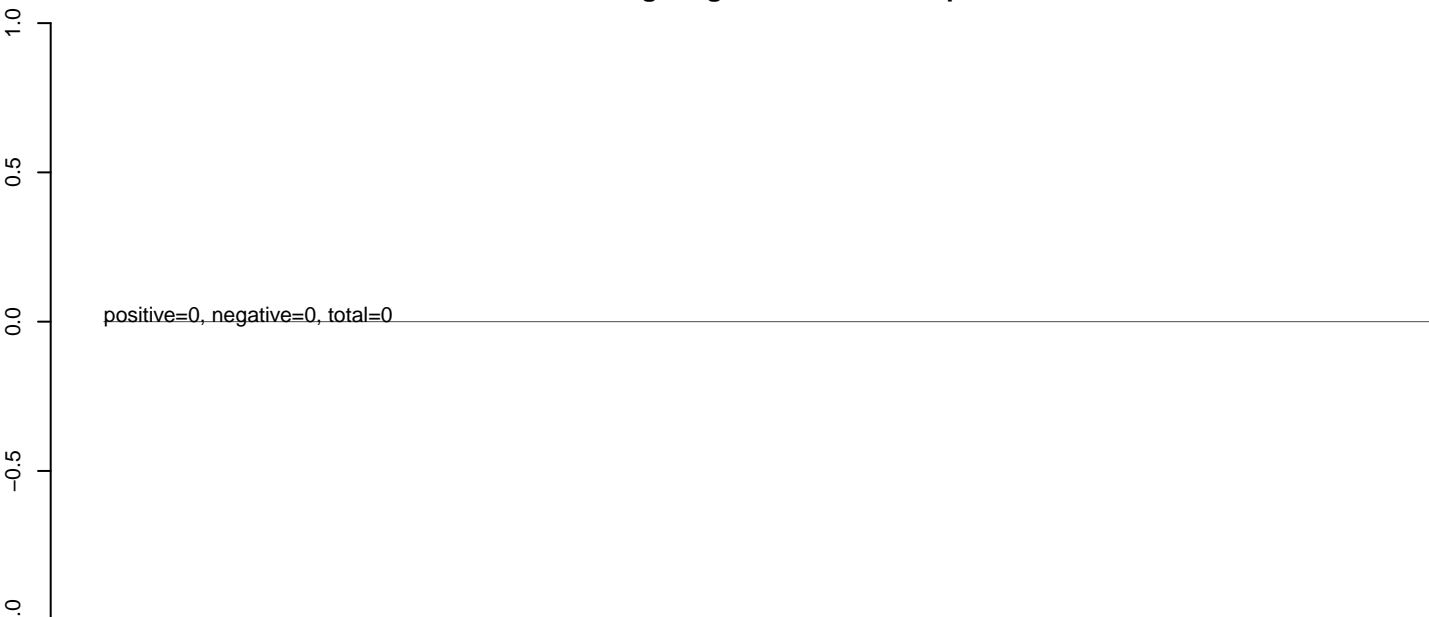
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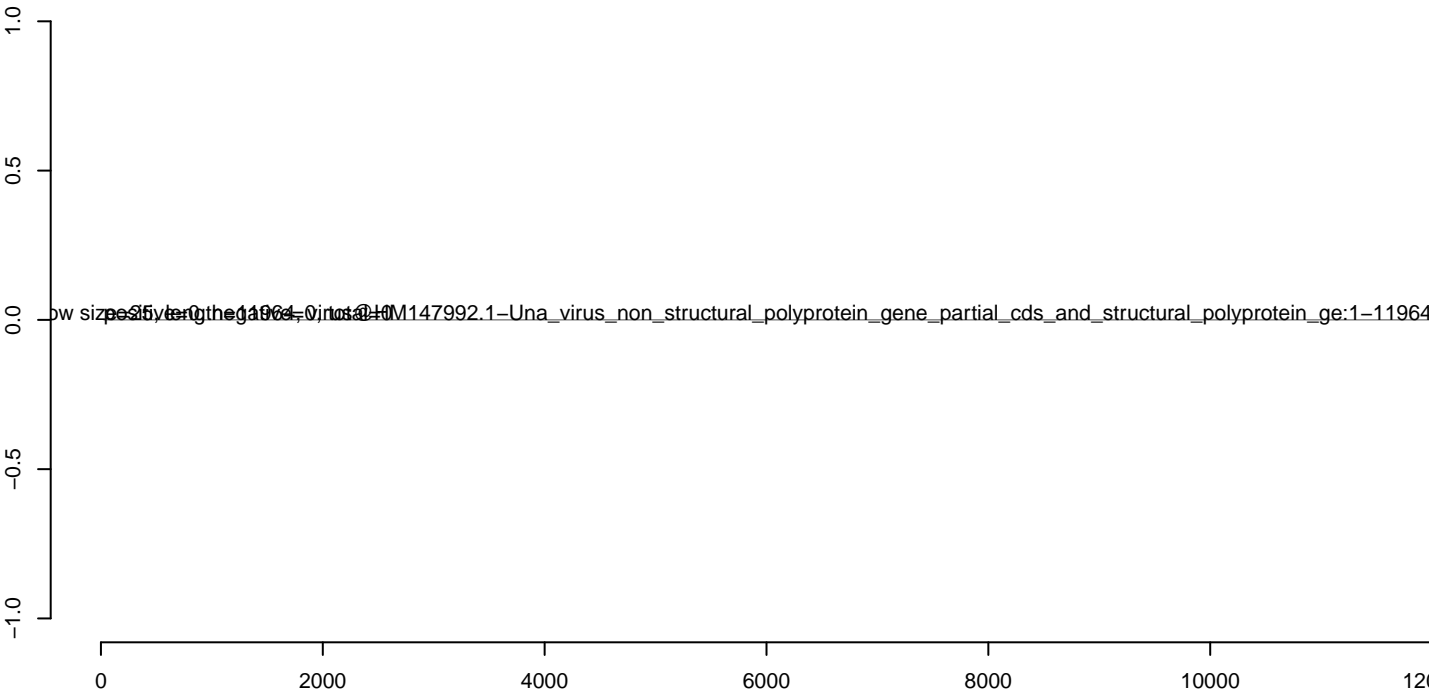
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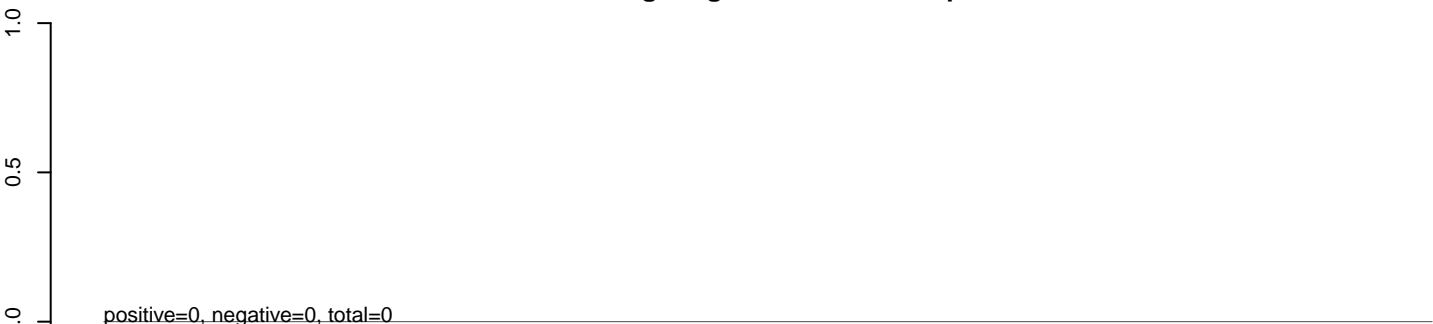
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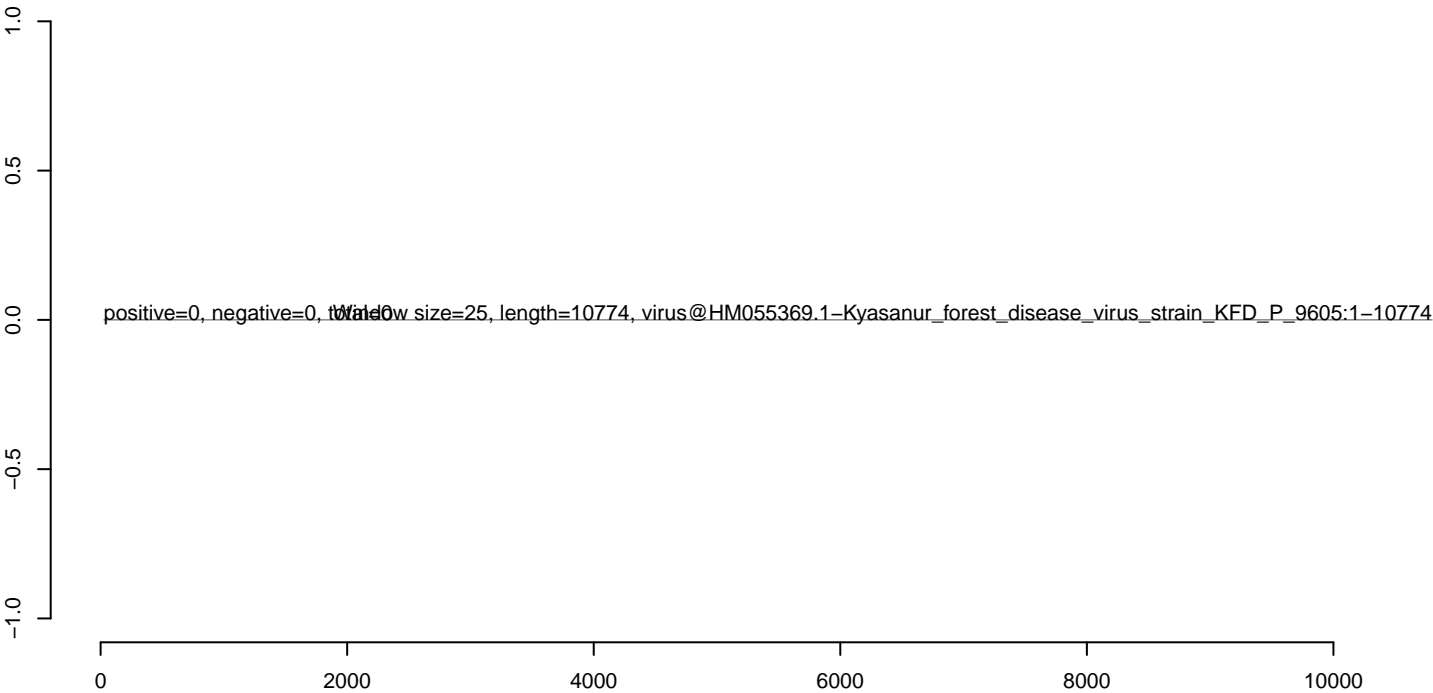
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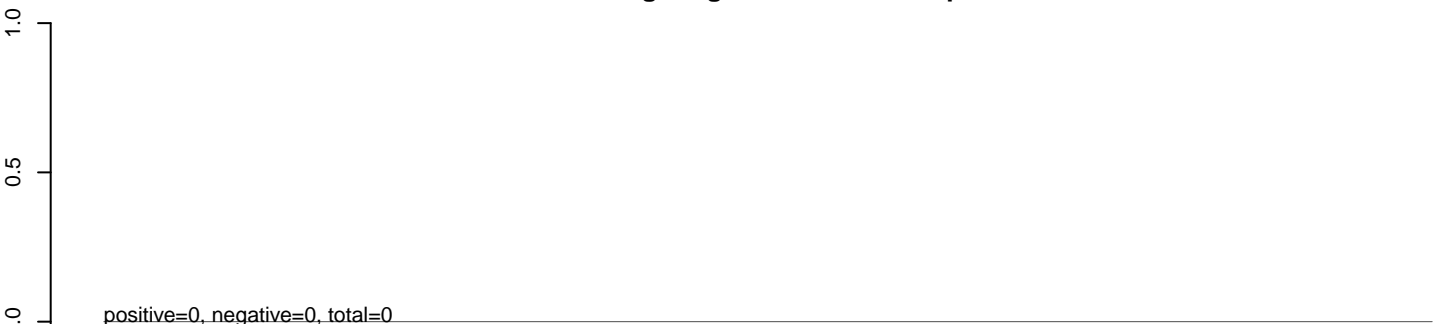


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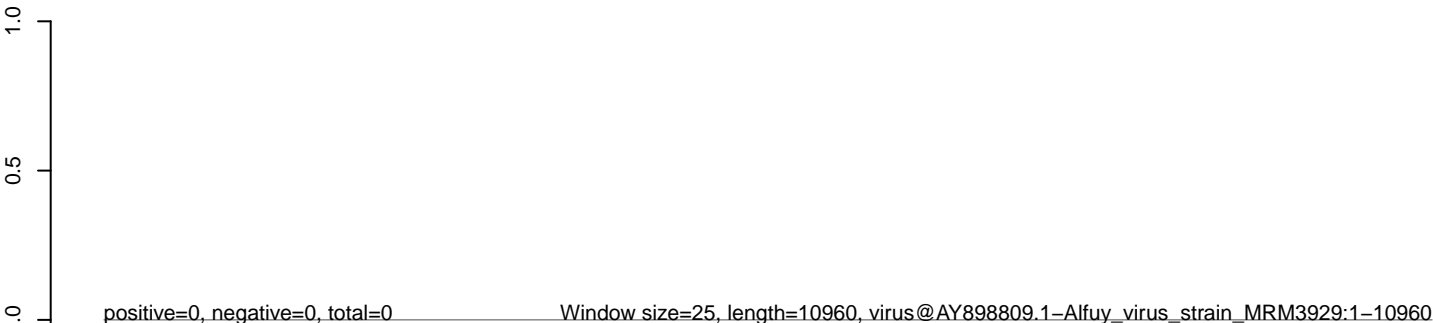
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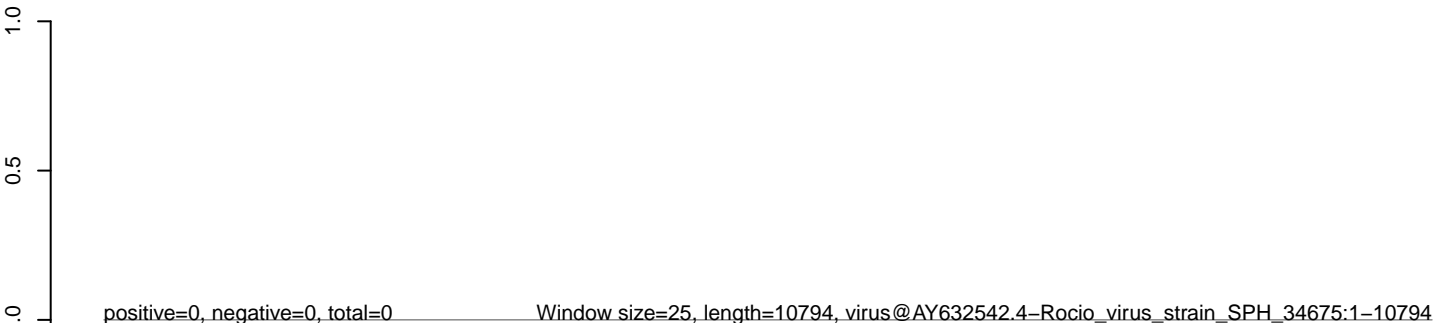
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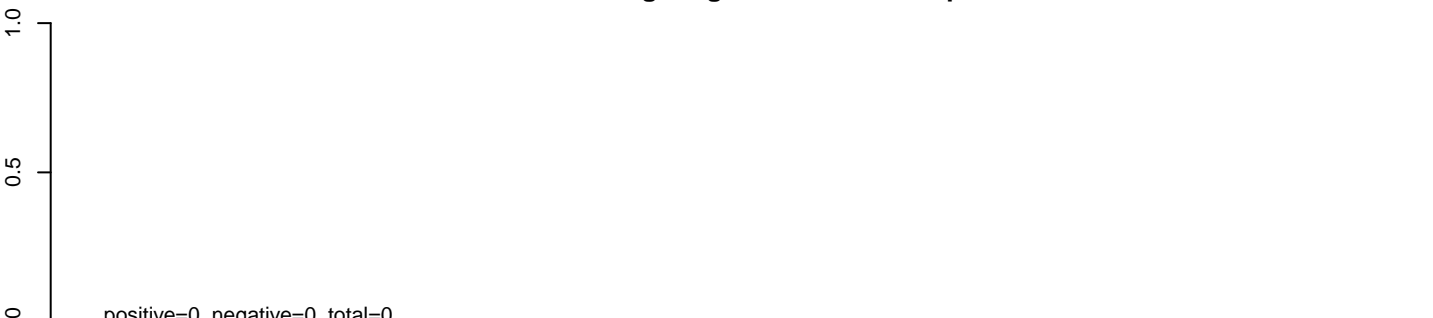
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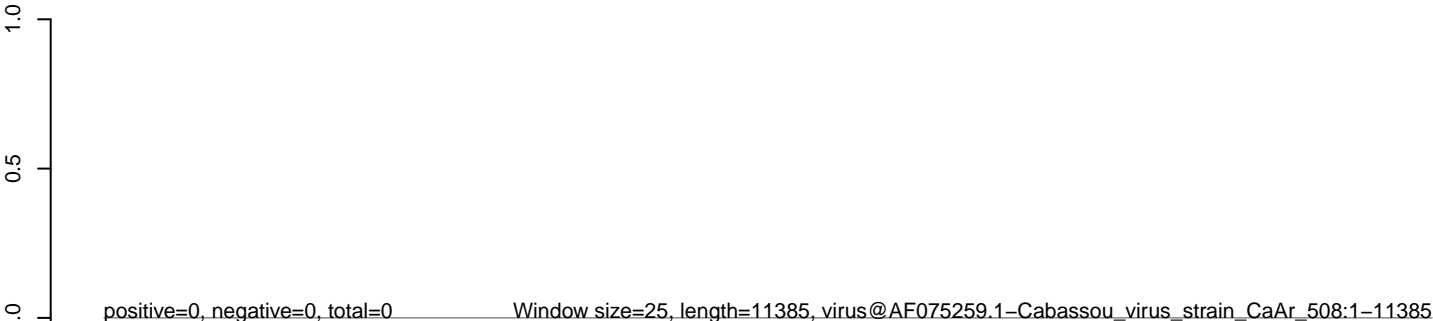
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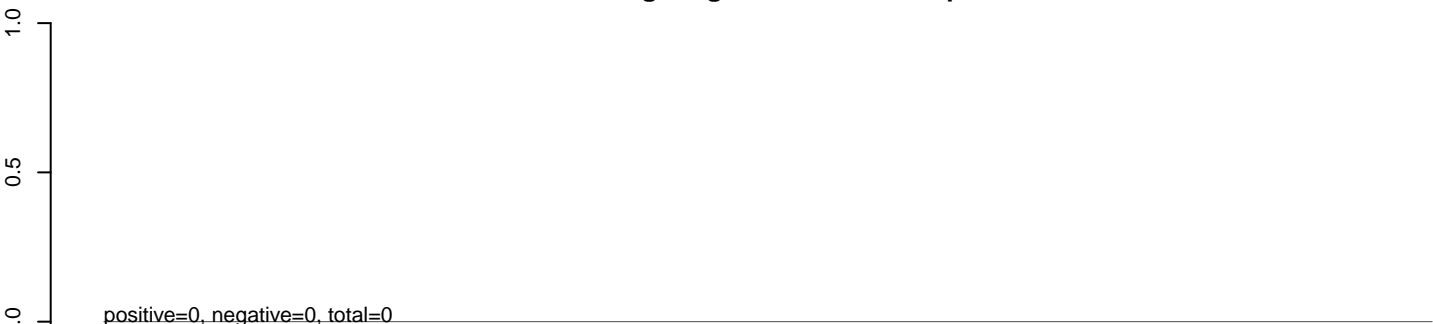


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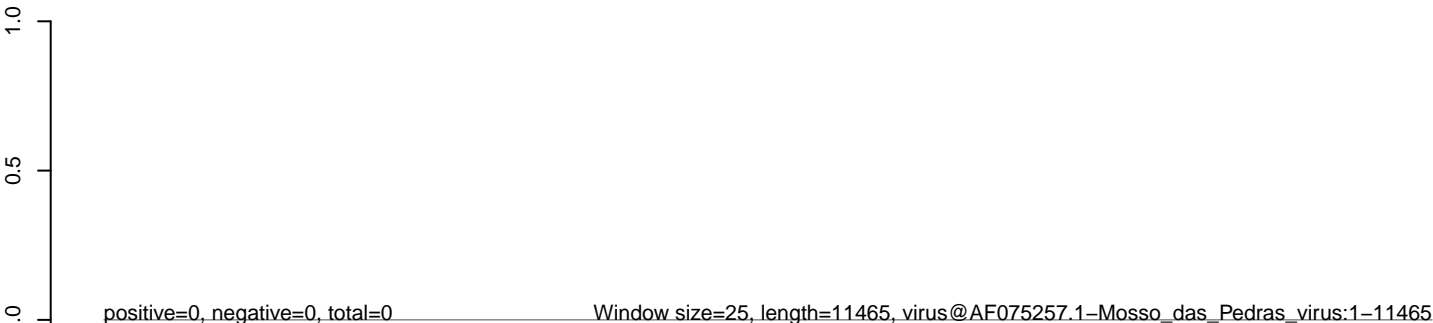
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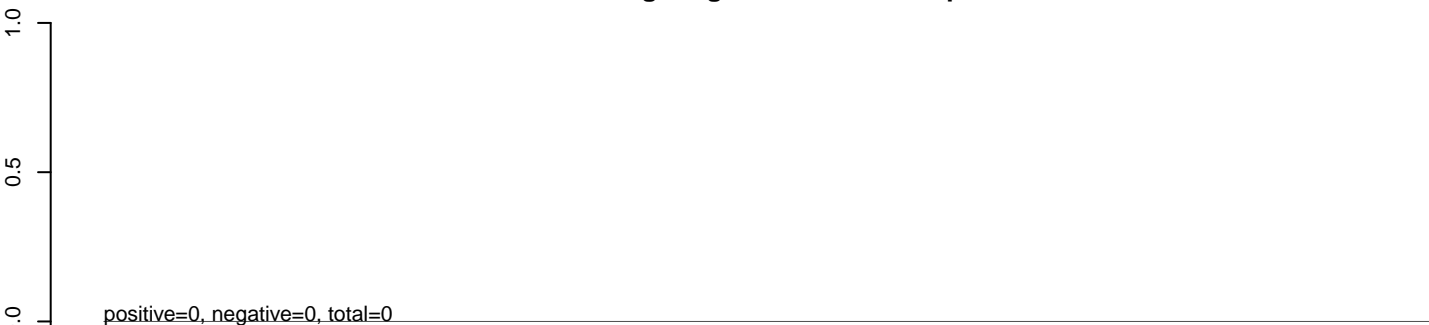
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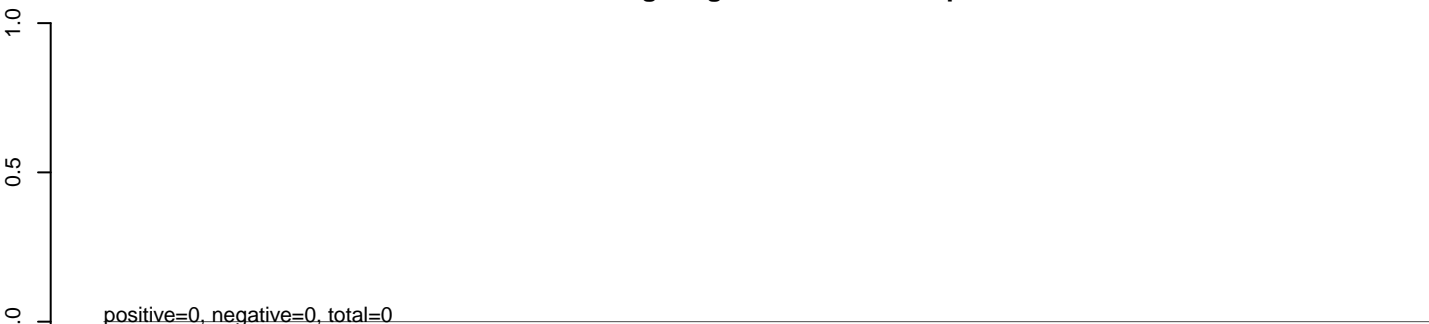
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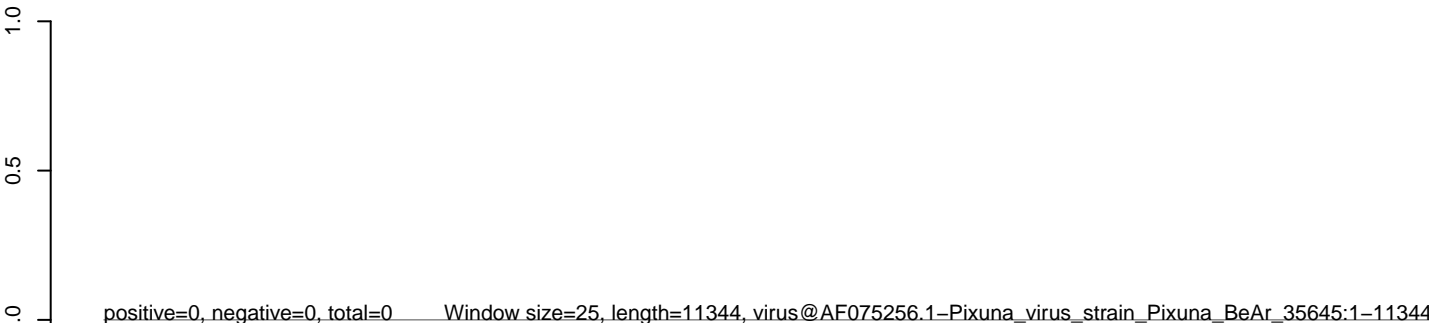
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