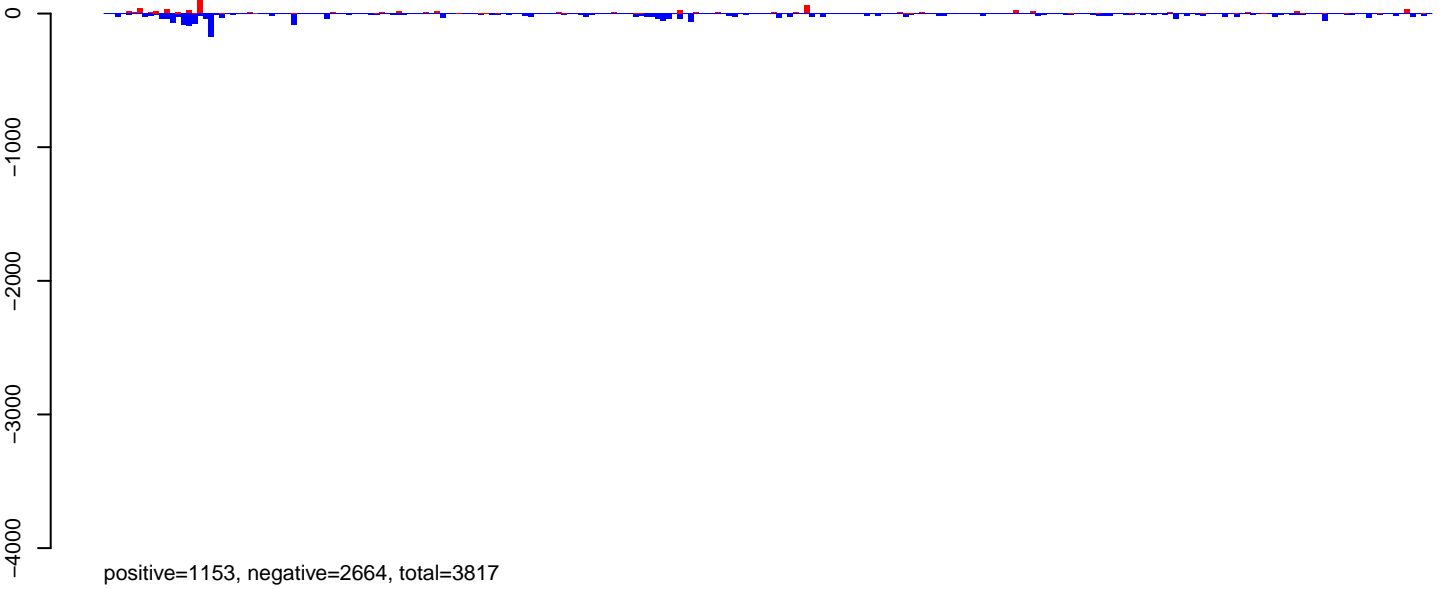
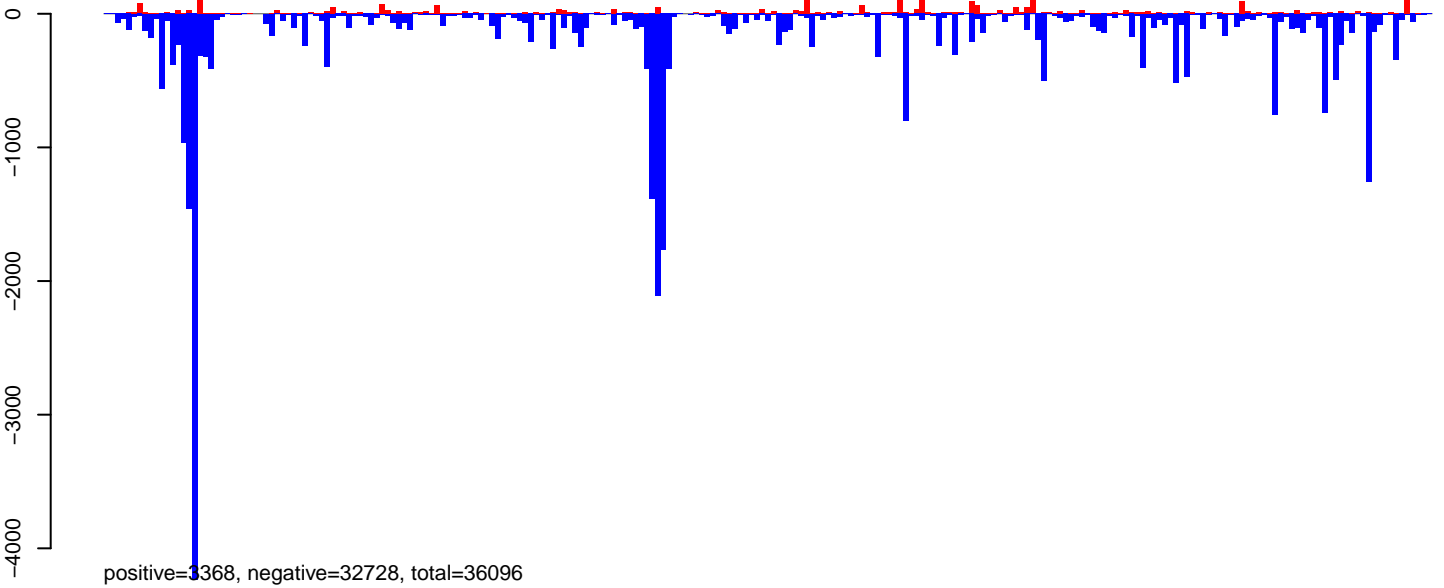


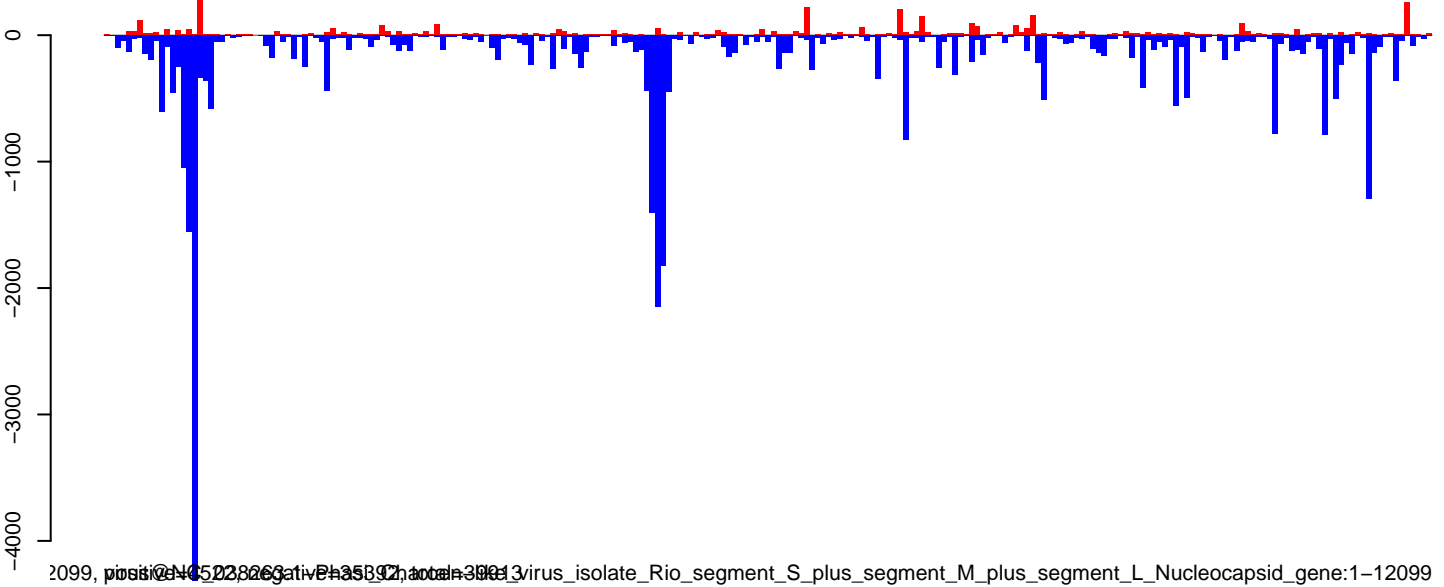
AeAeg_Aag2cells_CHIKV.18_23.rep



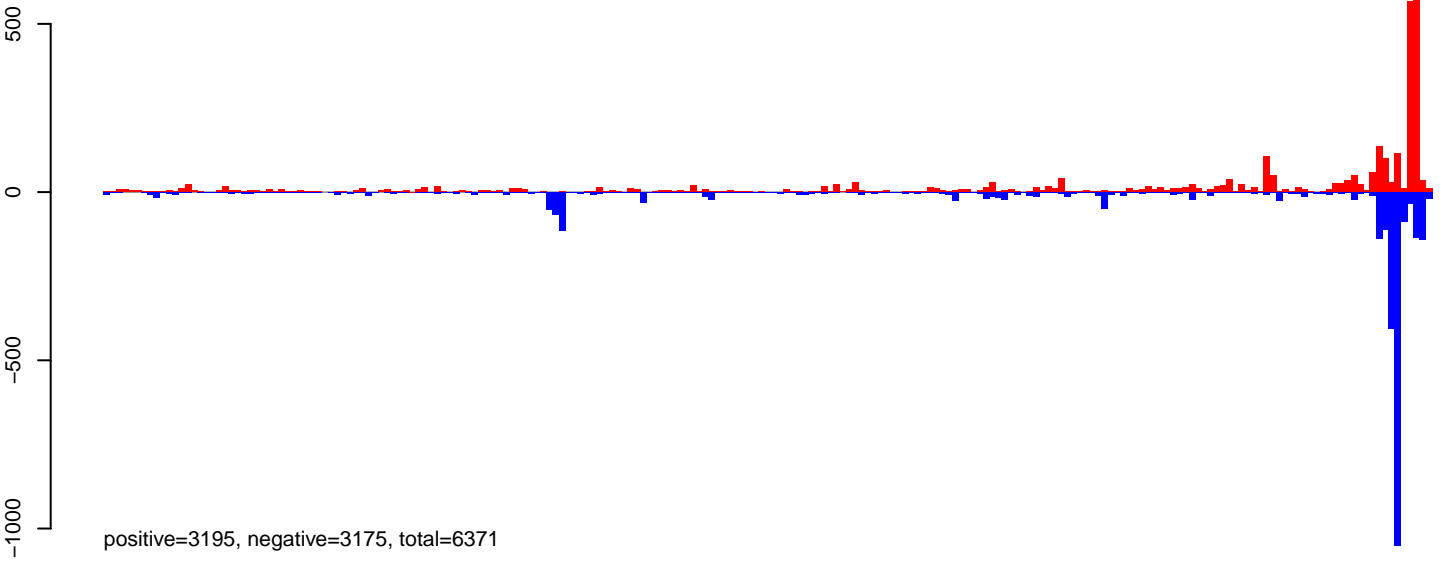
AeAeg_Aag2cells_CHIKV.24_35.rep



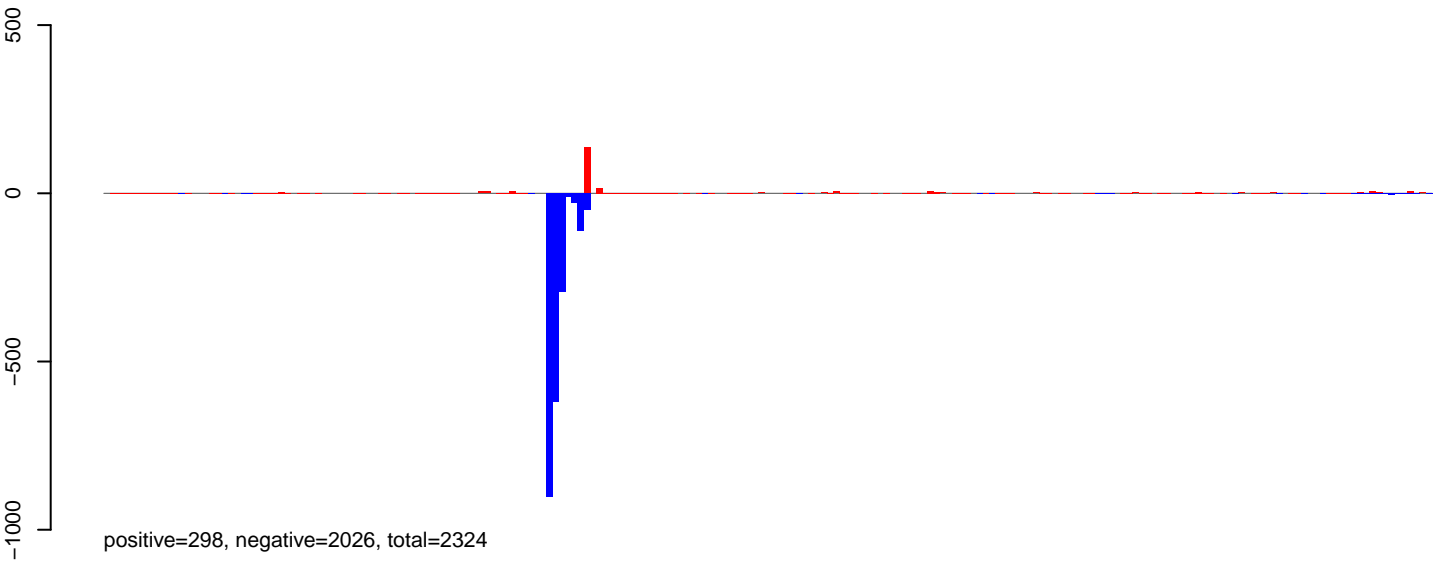
AeAeg_Aag2cells_CHIKV.rep



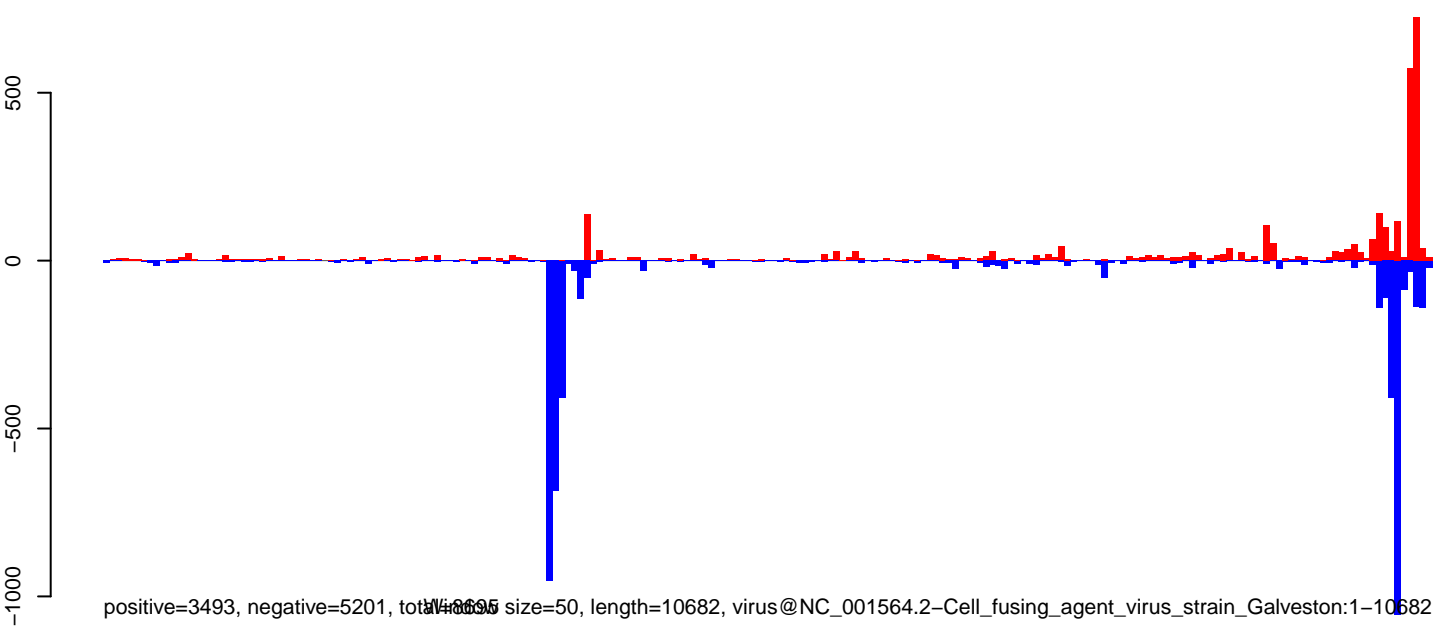
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

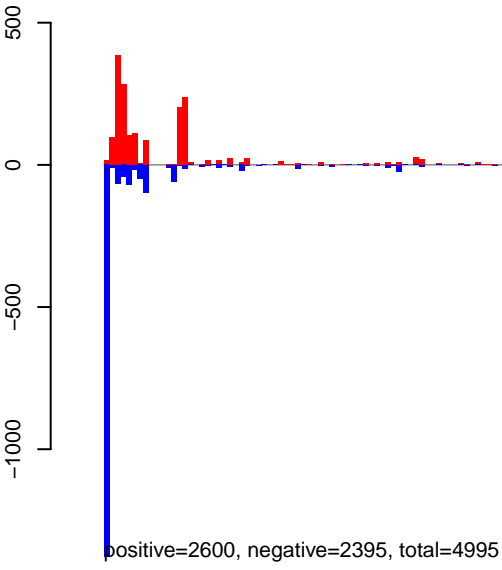


AeAeg_Aag2cells_CHIKV.rep

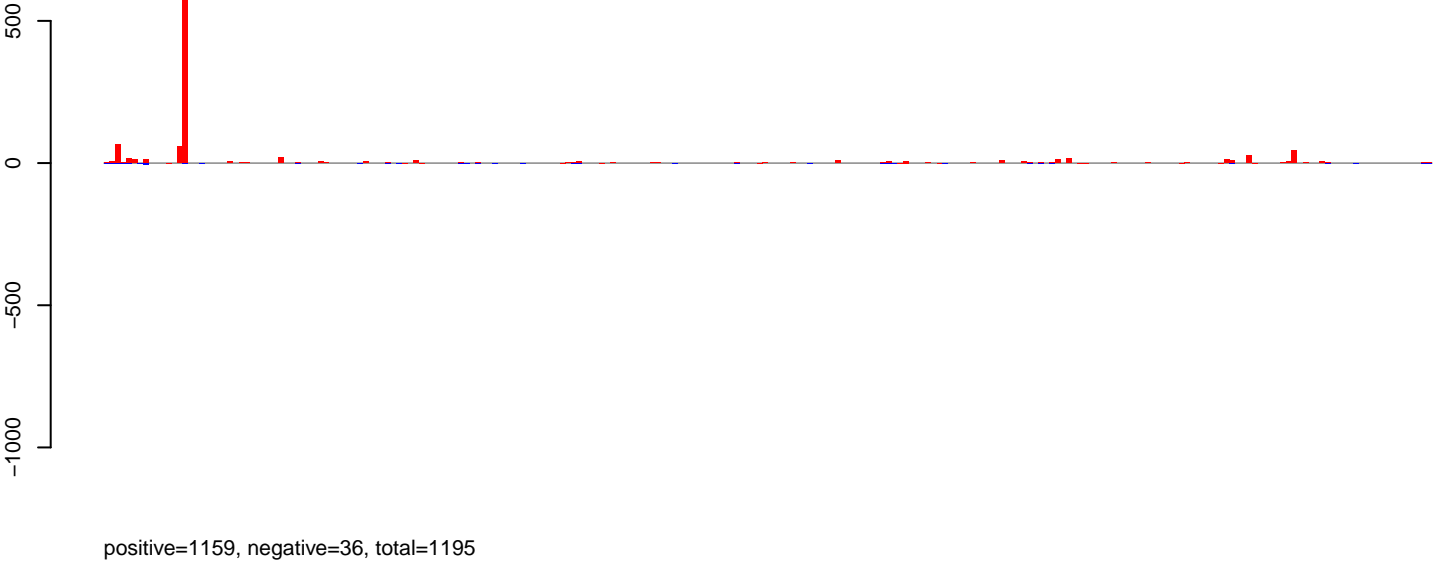


0 2000 4000 6000 8000 10000

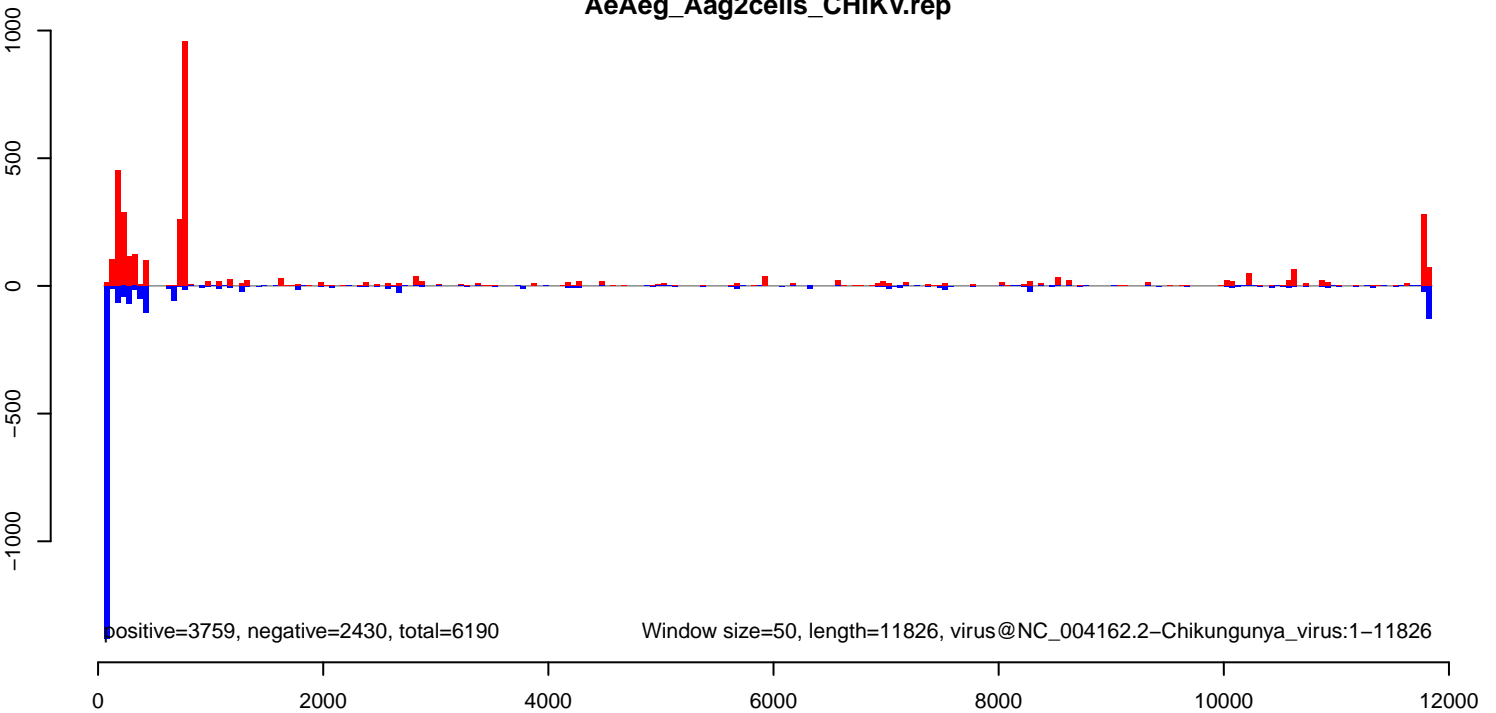
AeAeg_Aag2cells_CHIKV.18_23.rep



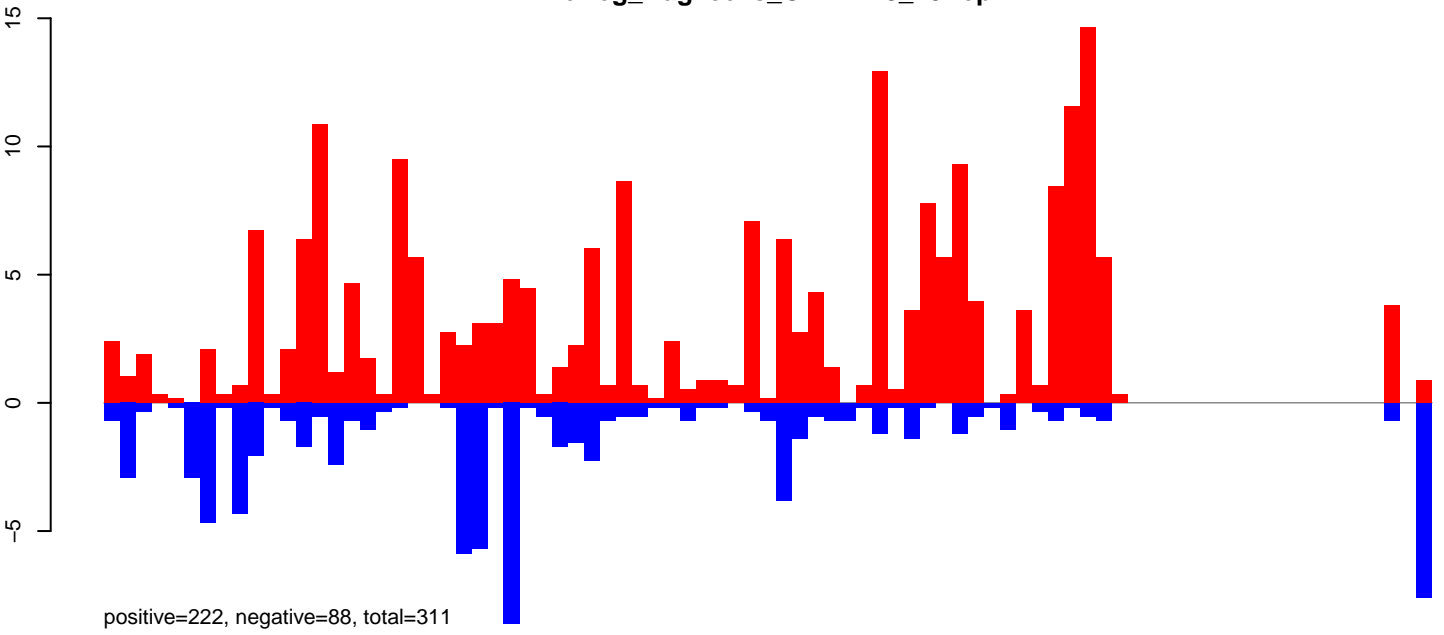
AeAeg_Aag2cells_CHIKV.24_35.rep



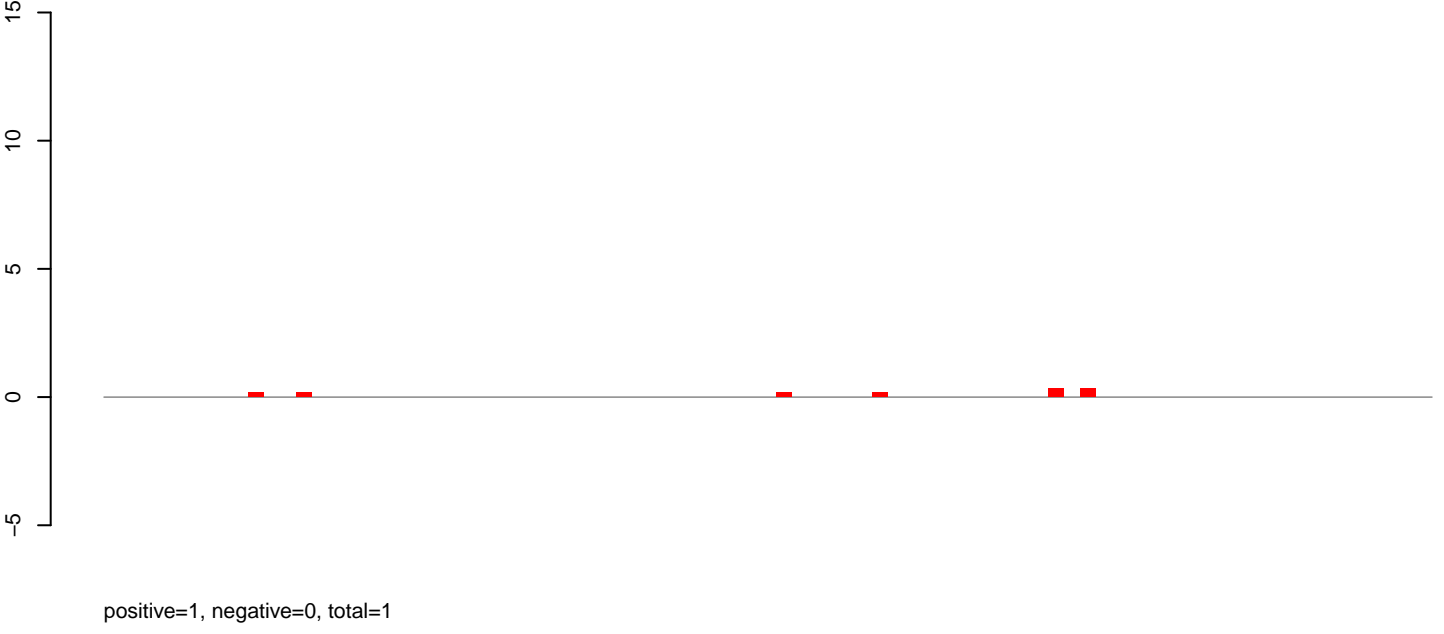
AeAeg_Aag2cells_CHIKV.rep



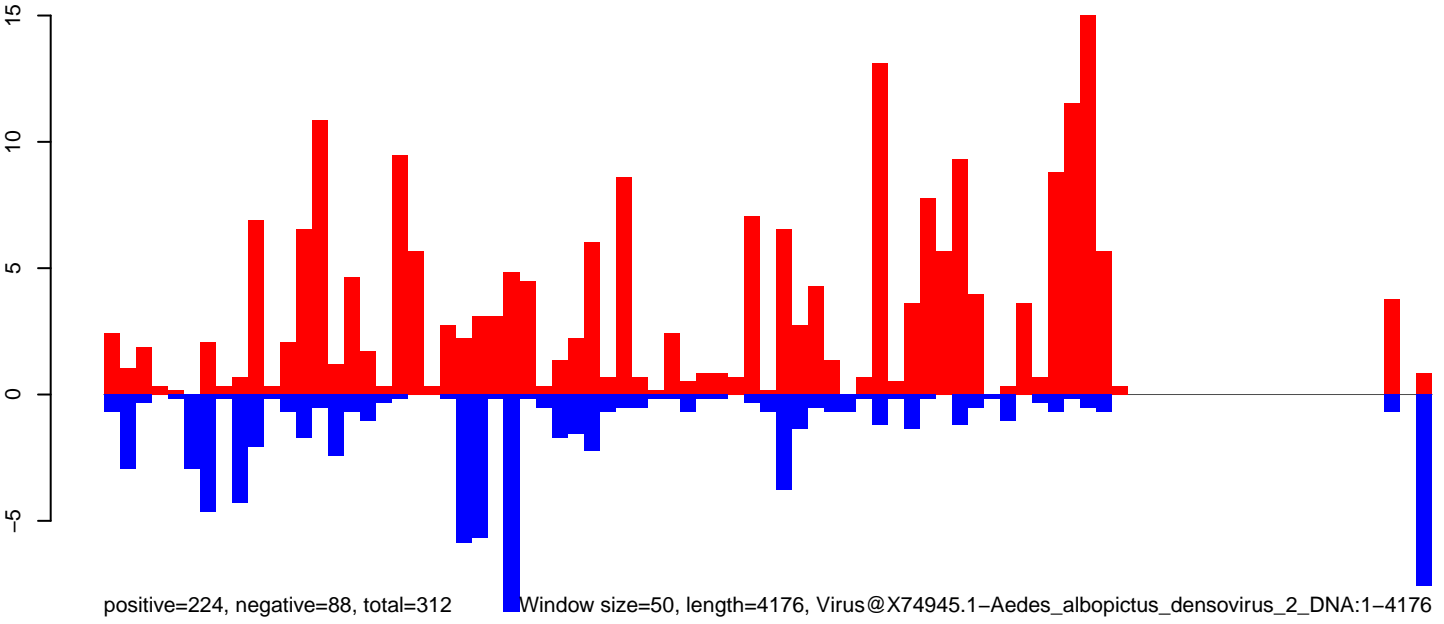
AeAeg_Aag2cells_CHIKV.18_23.rep



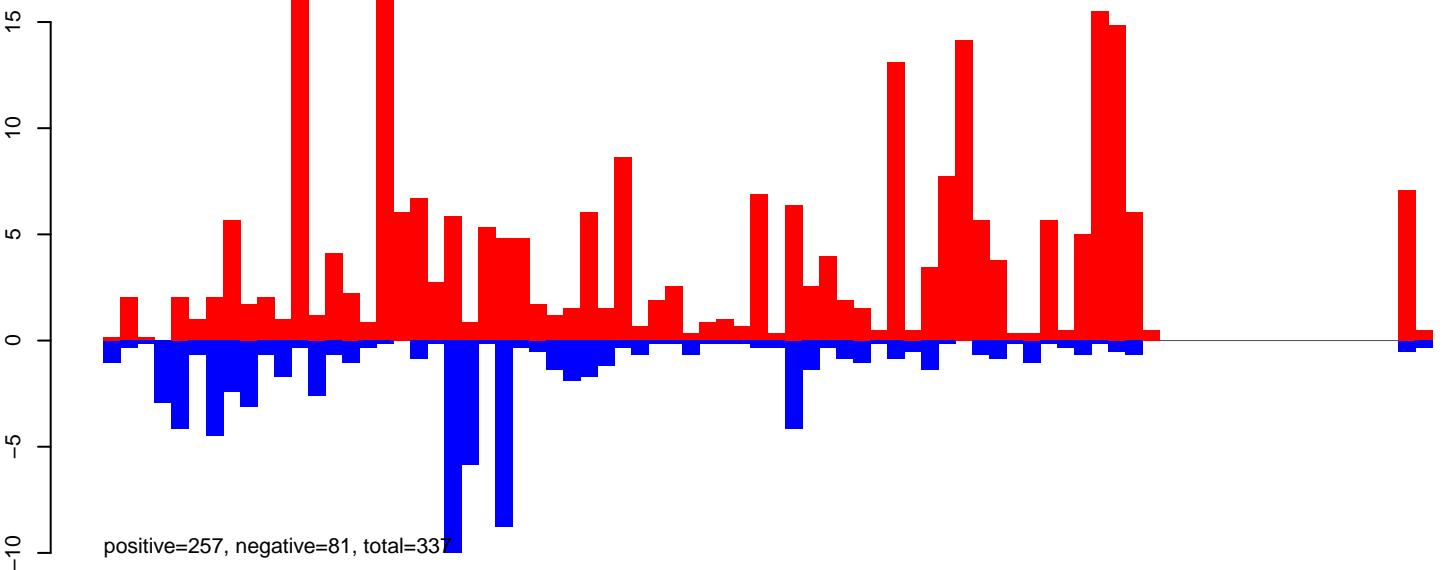
AeAeg_Aag2cells_CHIKV.24_35.rep



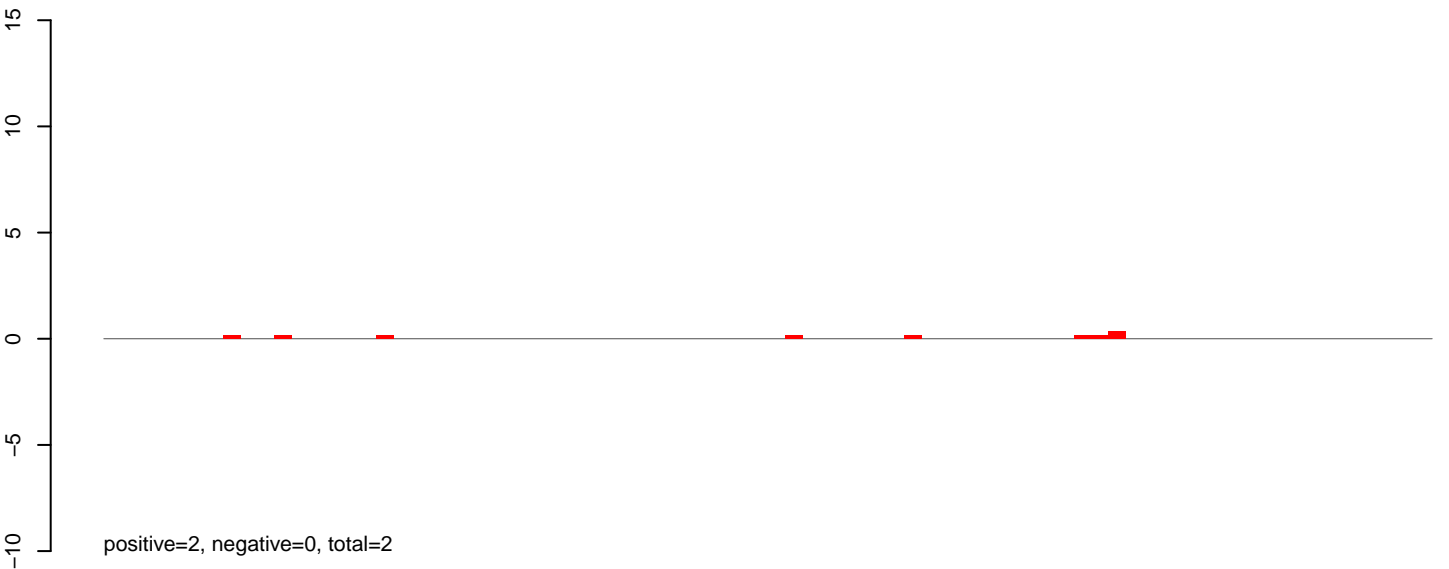
AeAeg_Aag2cells_CHIKV.rep



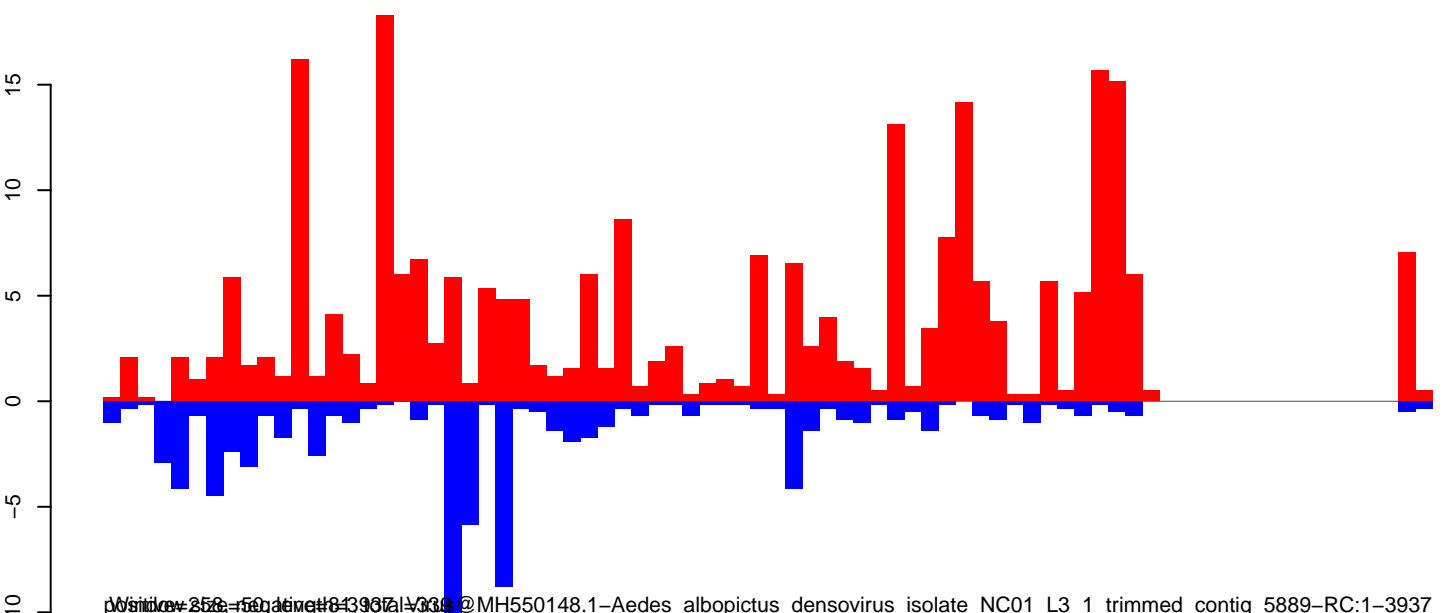
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



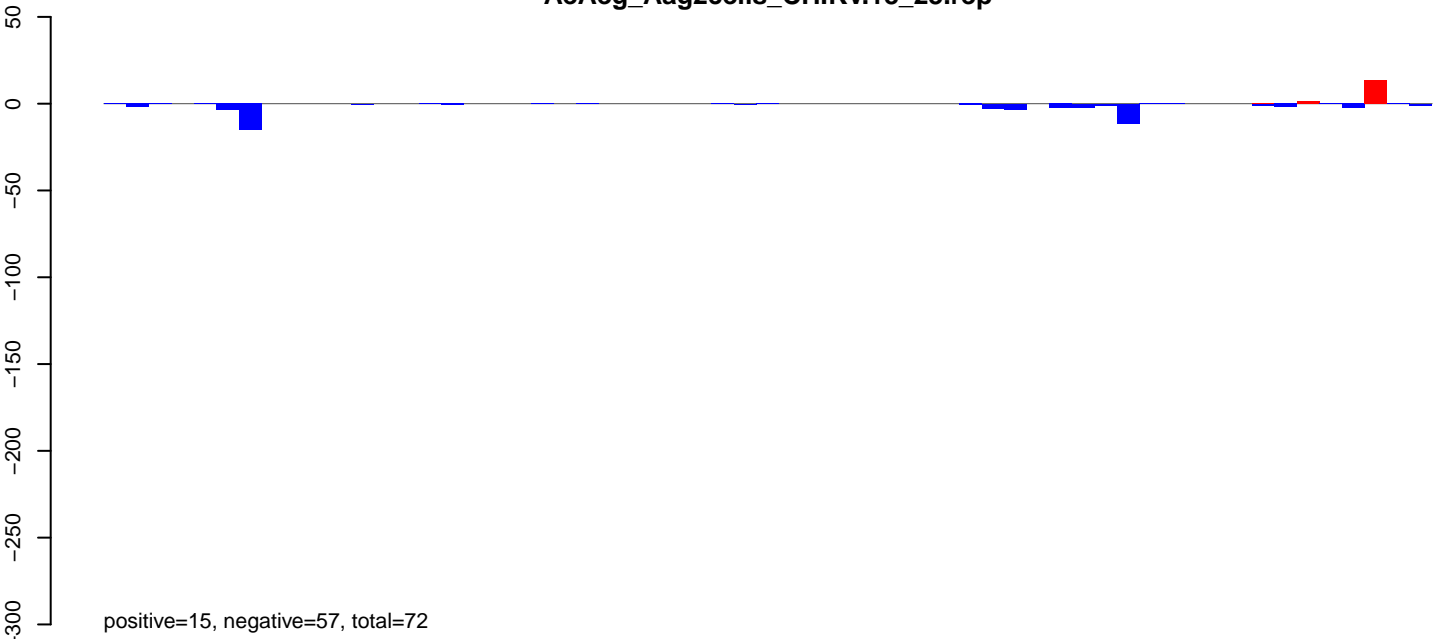
AeAeg_Aag2cells_CHIKV.rep



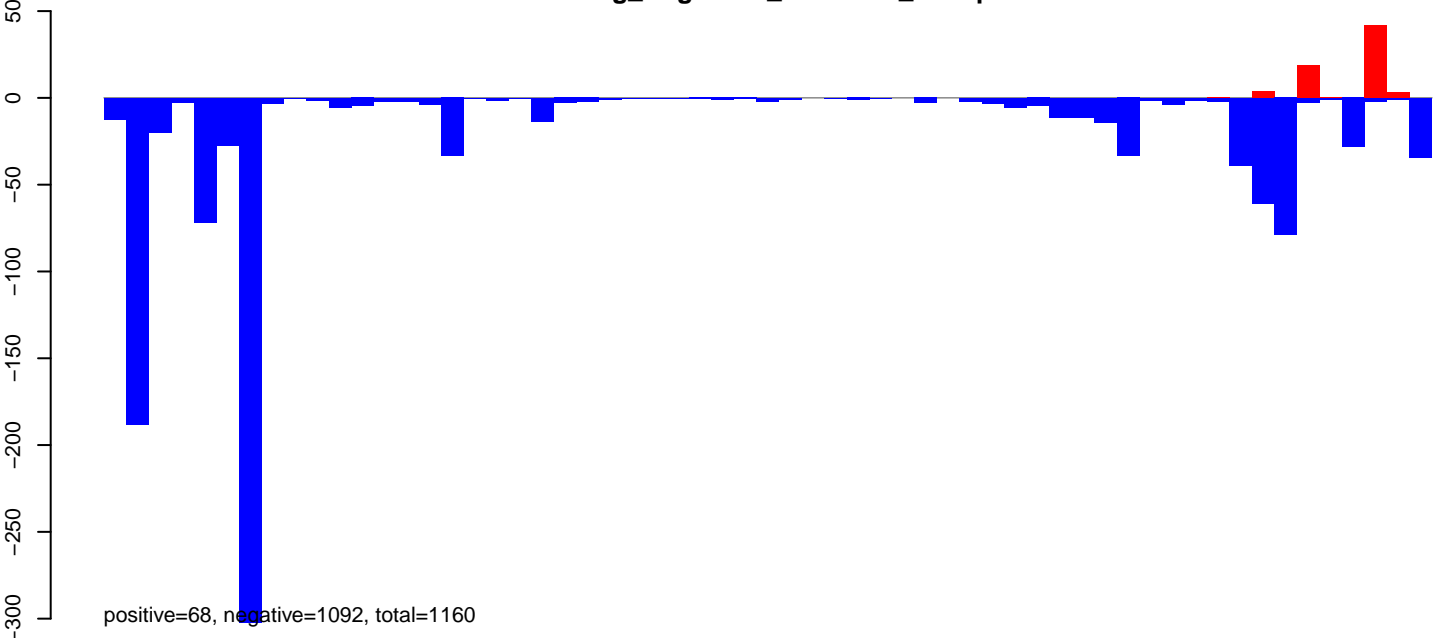
Window 358, starting at 3967, ending at 4336 @MH550148.1-Aedes_albopictus_densovirus_isolate_NC01_L3_1_trimmed_contig_5889-RC:1-3937

0 1000 2000 3000 4000

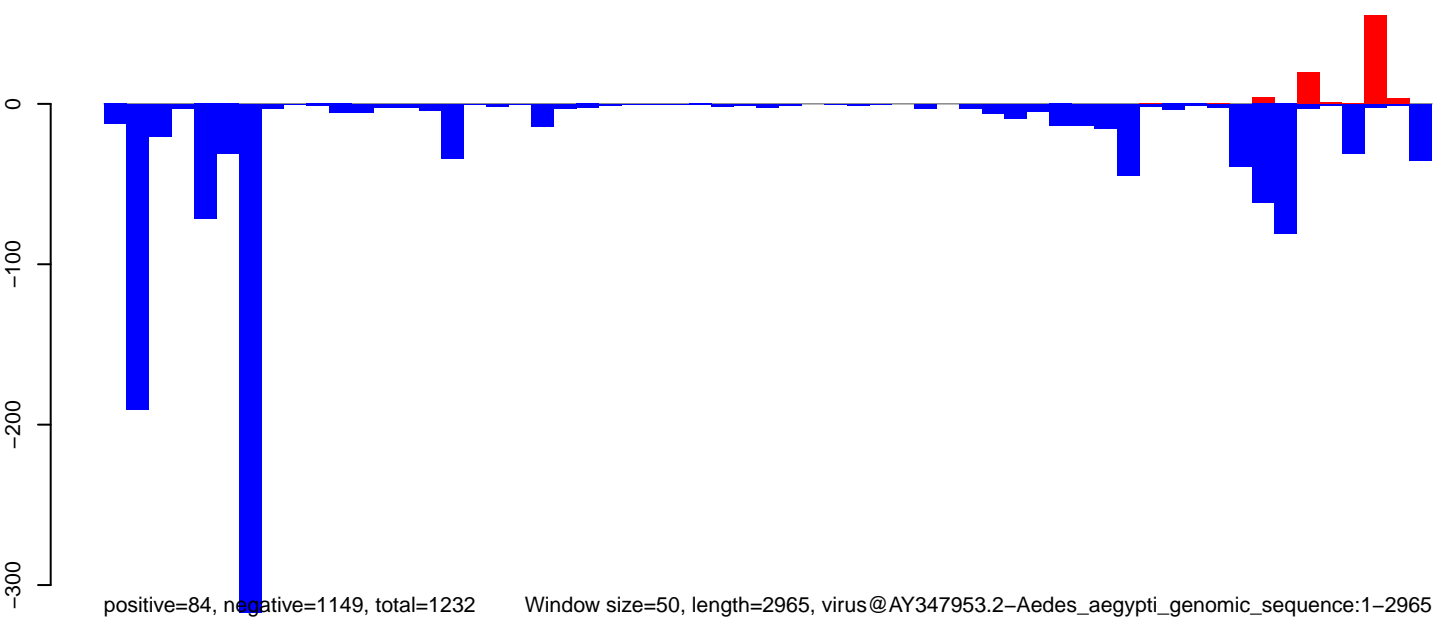
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



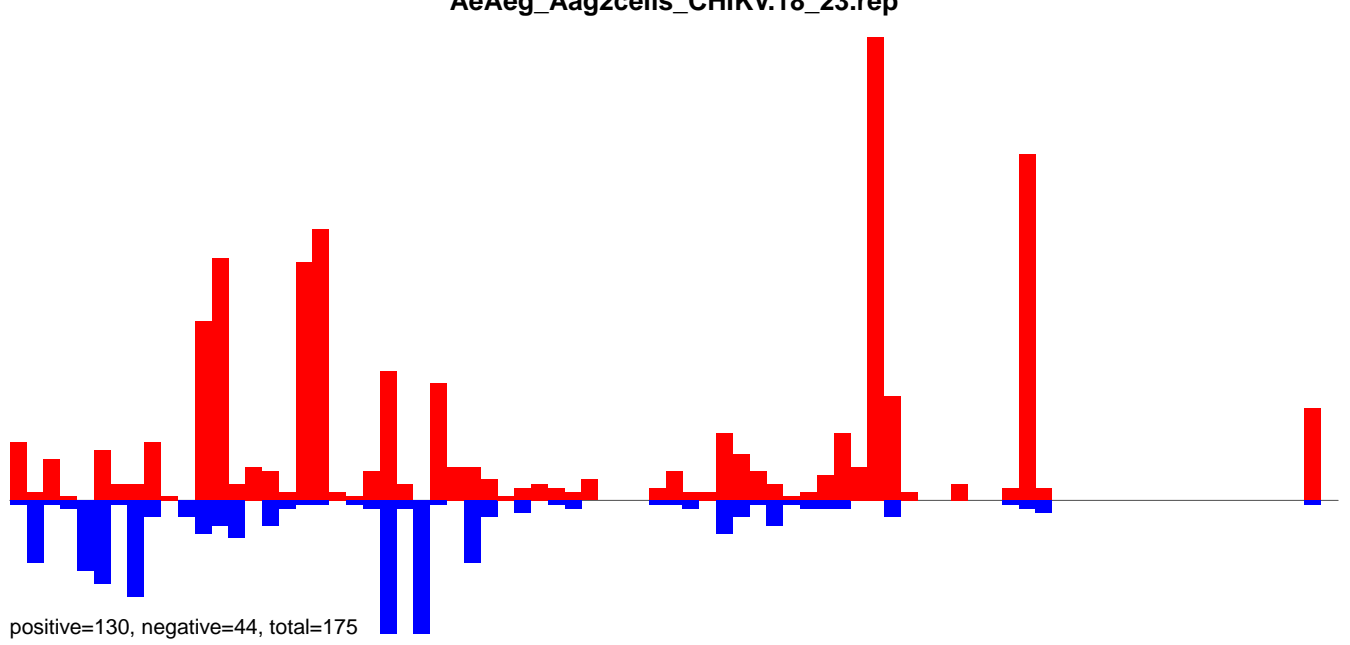
AeAeg_Aag2cells_CHIKV.rep



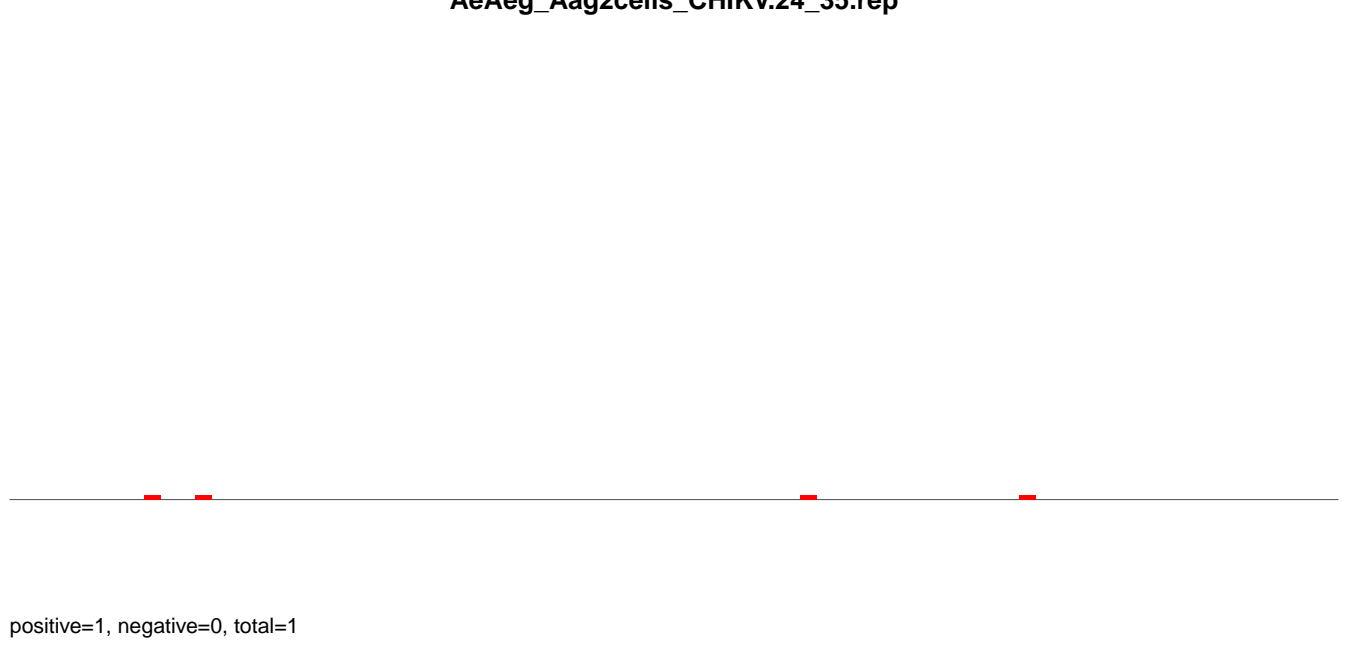
Window size=50, length=2965, virus@AY347953.2-Aedes_aegypti_genomic_sequence:1-2965

0 500 1000 1500 2000 2500 3000

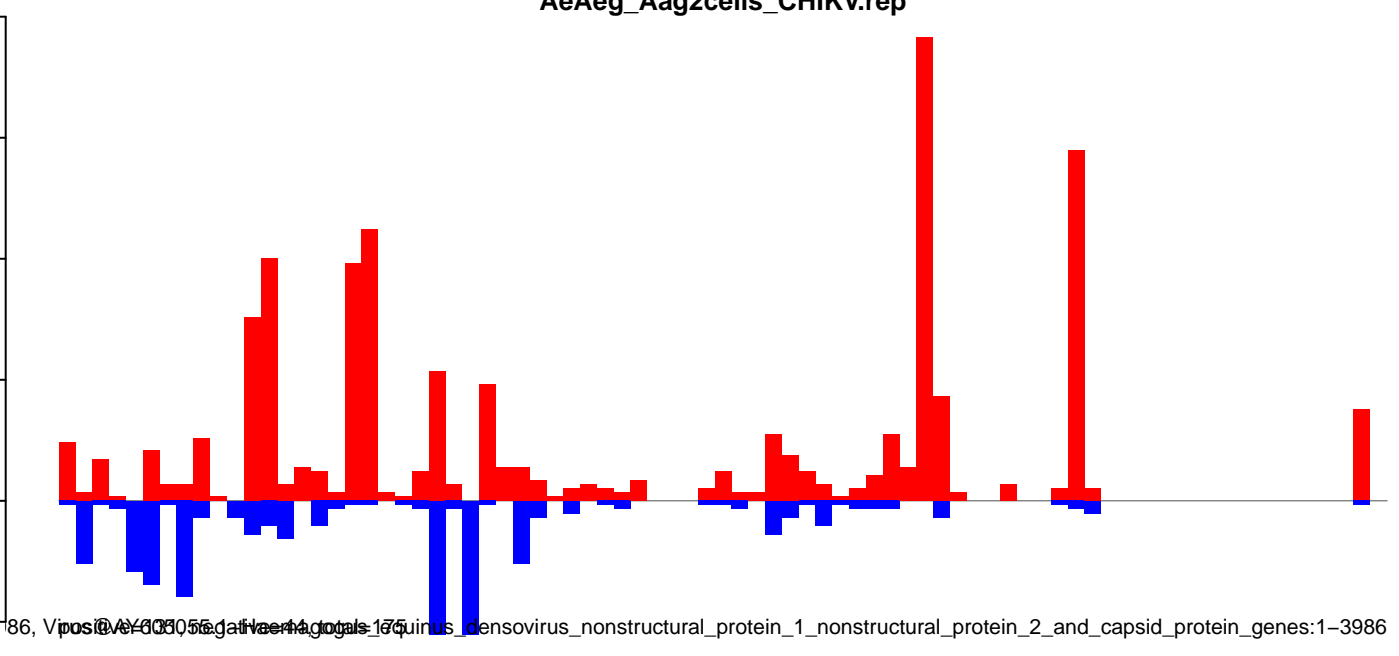
AeAeg_Aag2cells_CHIKV.18_23.rep



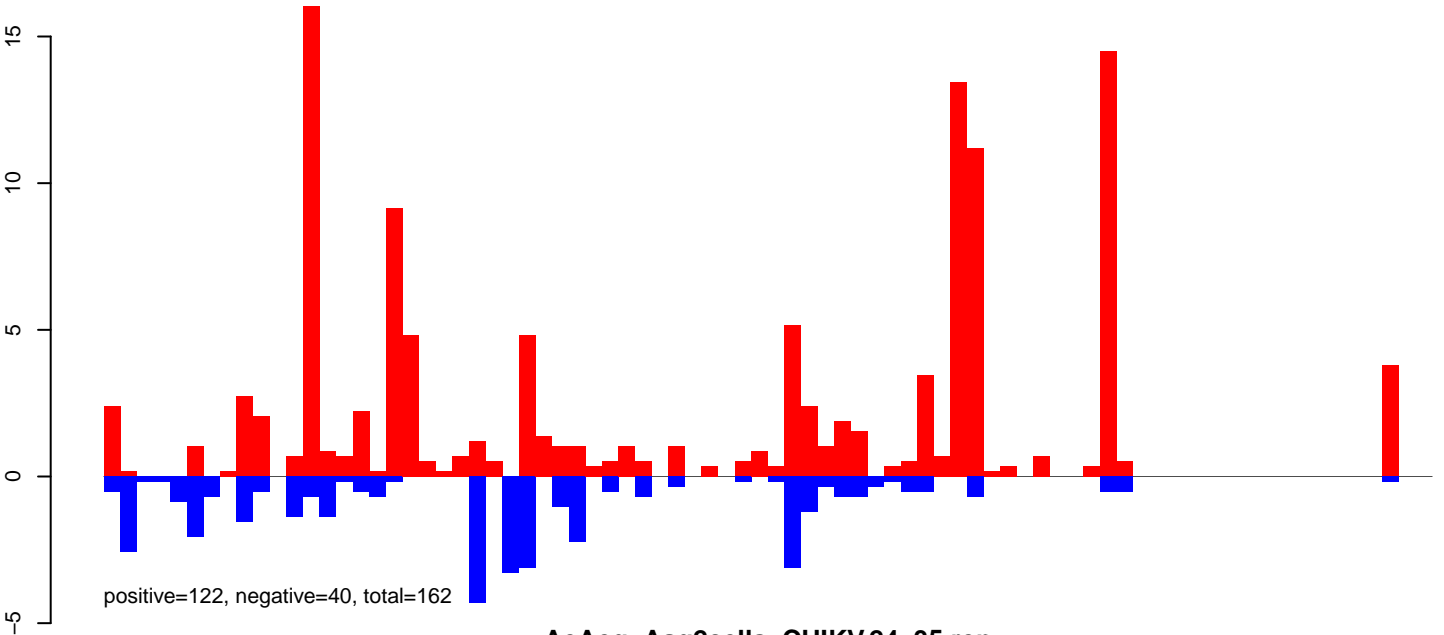
AeAeg_Aag2cells_CHIKV.24_35.rep



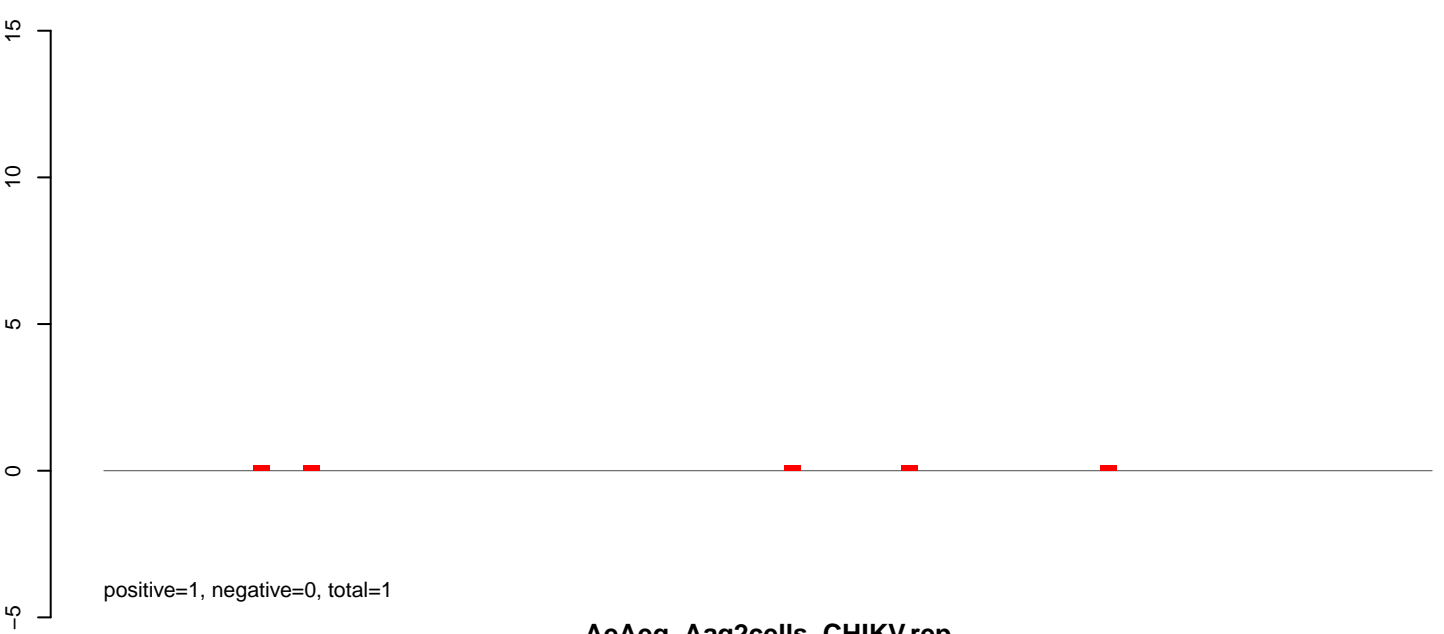
AeAeg_Aag2cells_CHIKV.rep



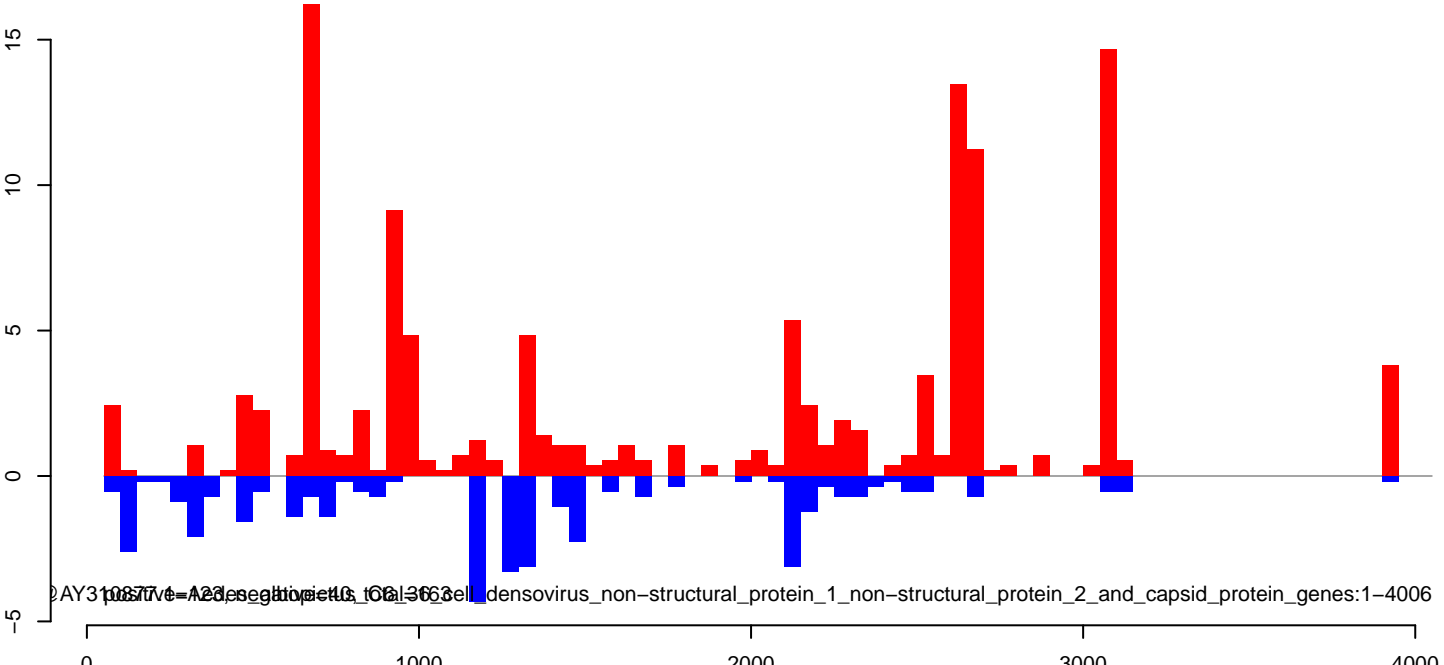
AeAeg_Aag2cells_CHIKV.18_23.rep



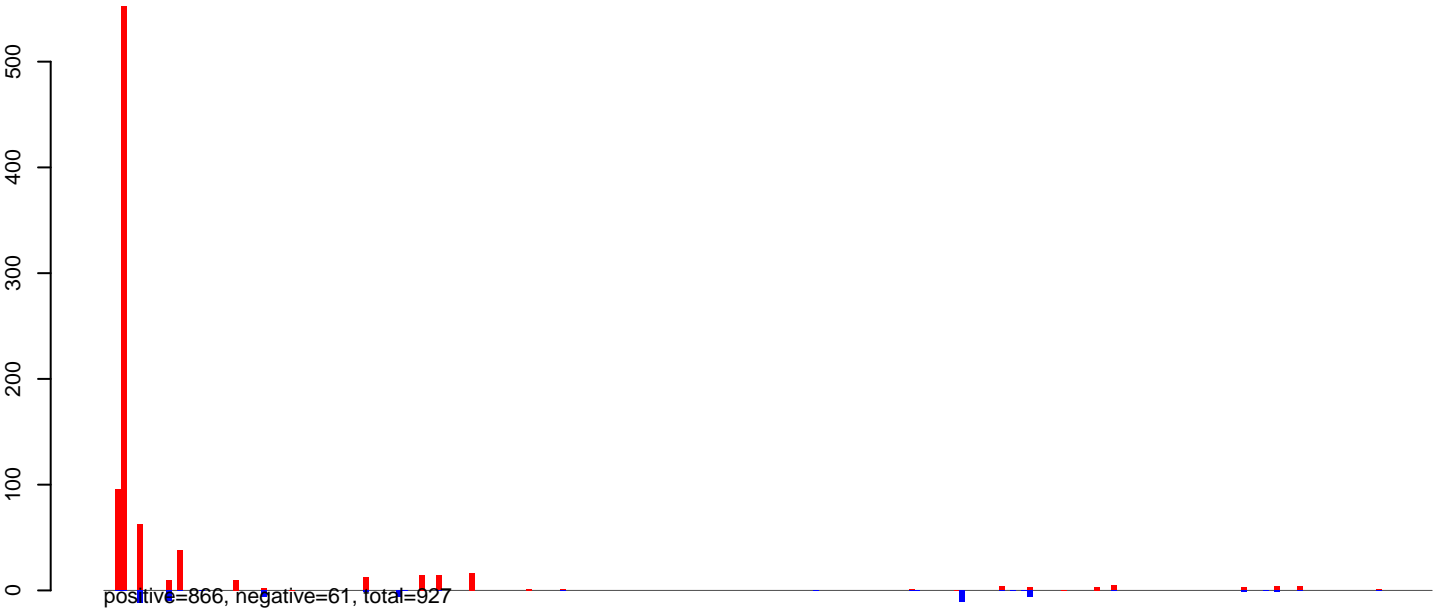
AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep



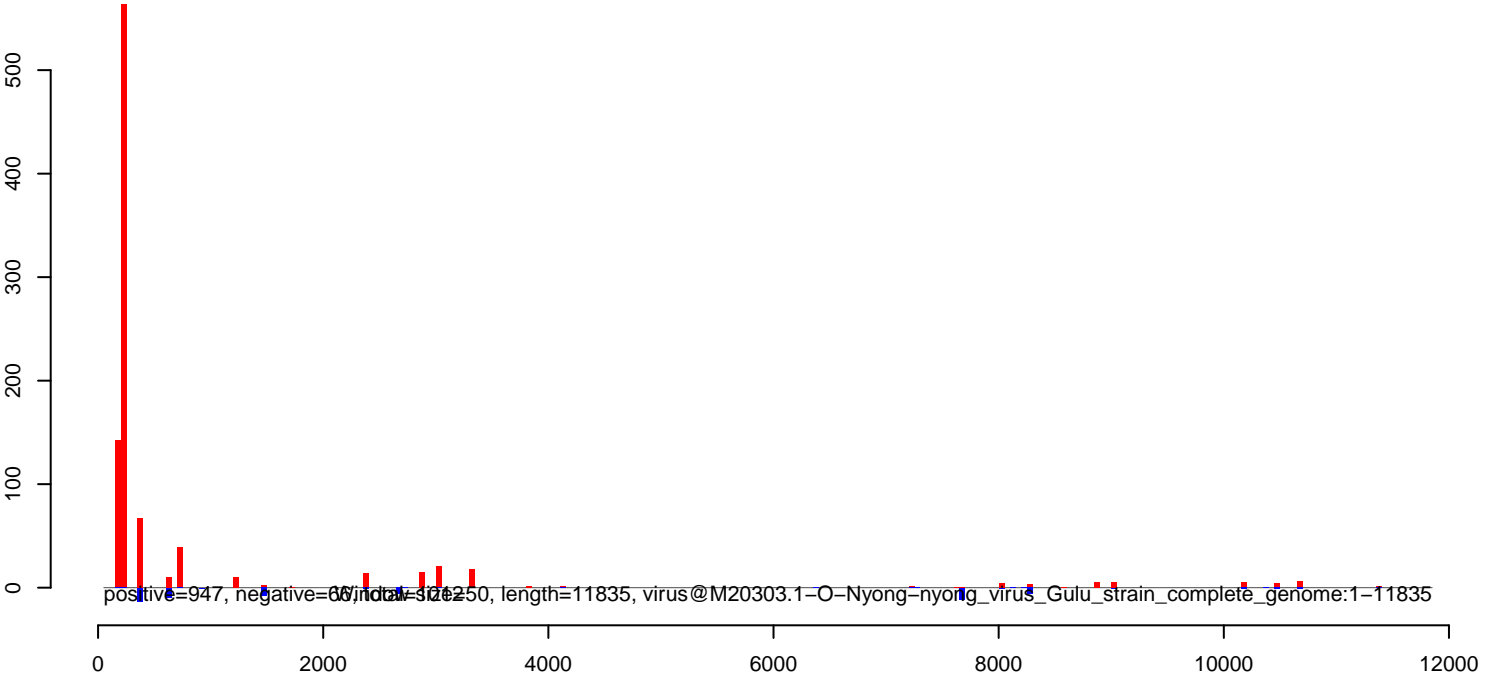
AeAeg_Aag2cells_CHIKV.18_23.rep



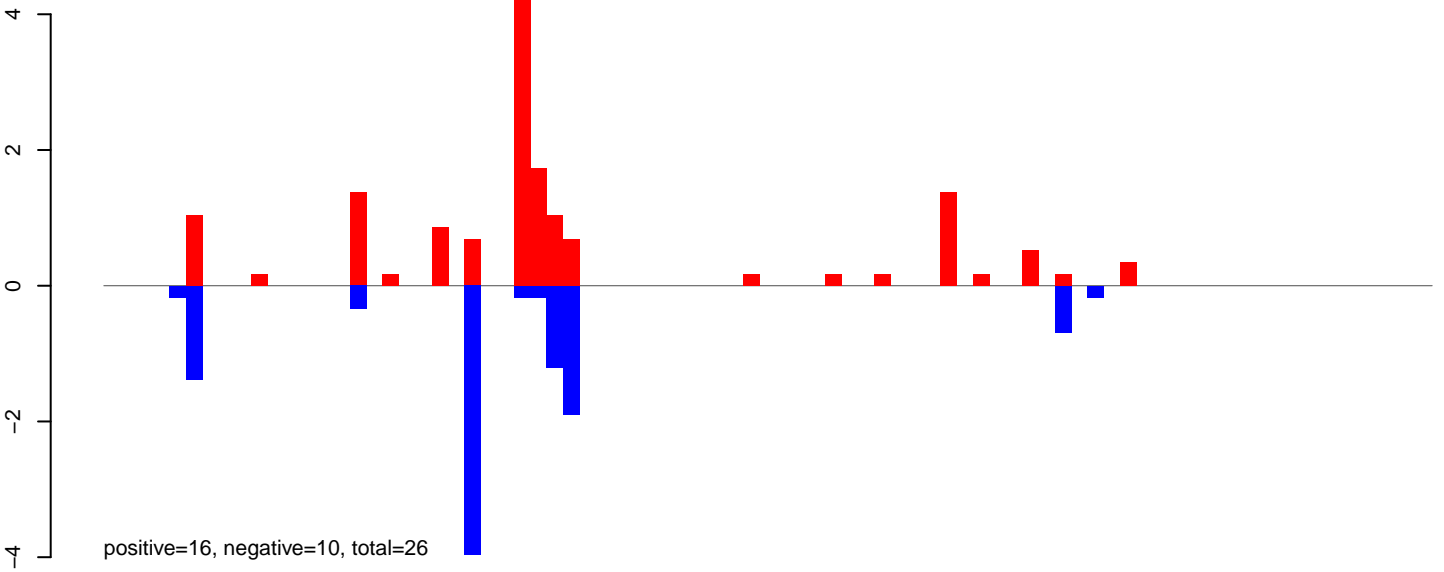
AeAeg_Aag2cells_CHIKV.24_35.rep



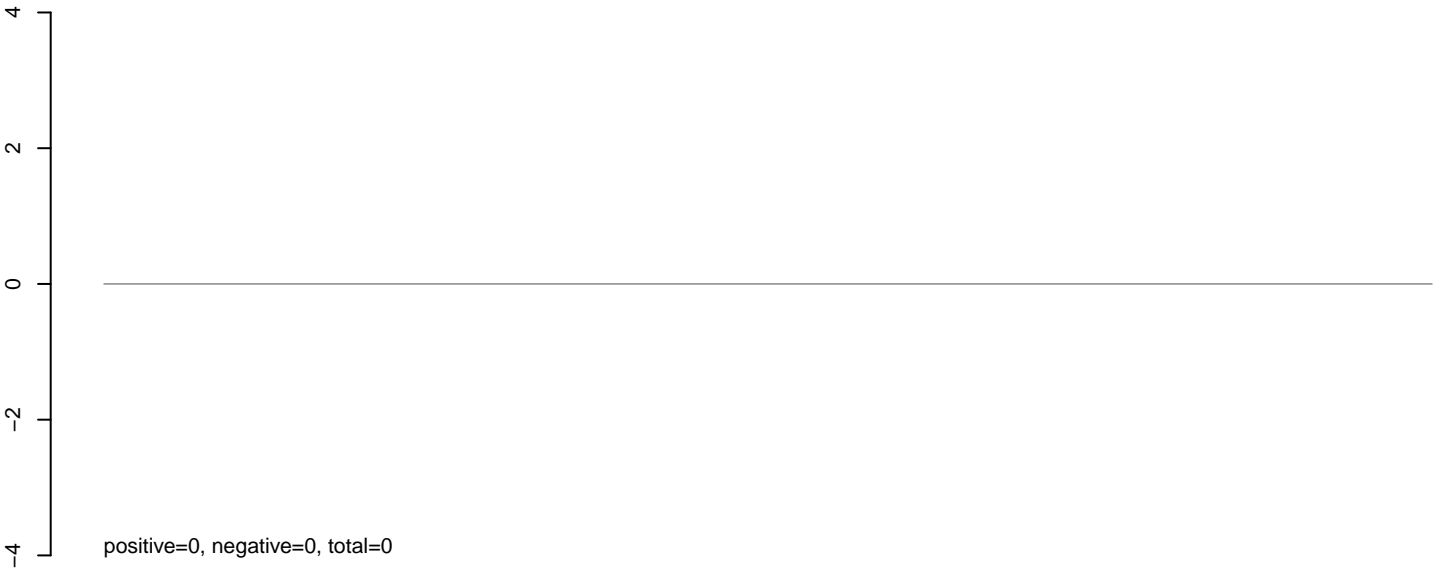
AeAeg_Aag2cells_CHIKV.rep



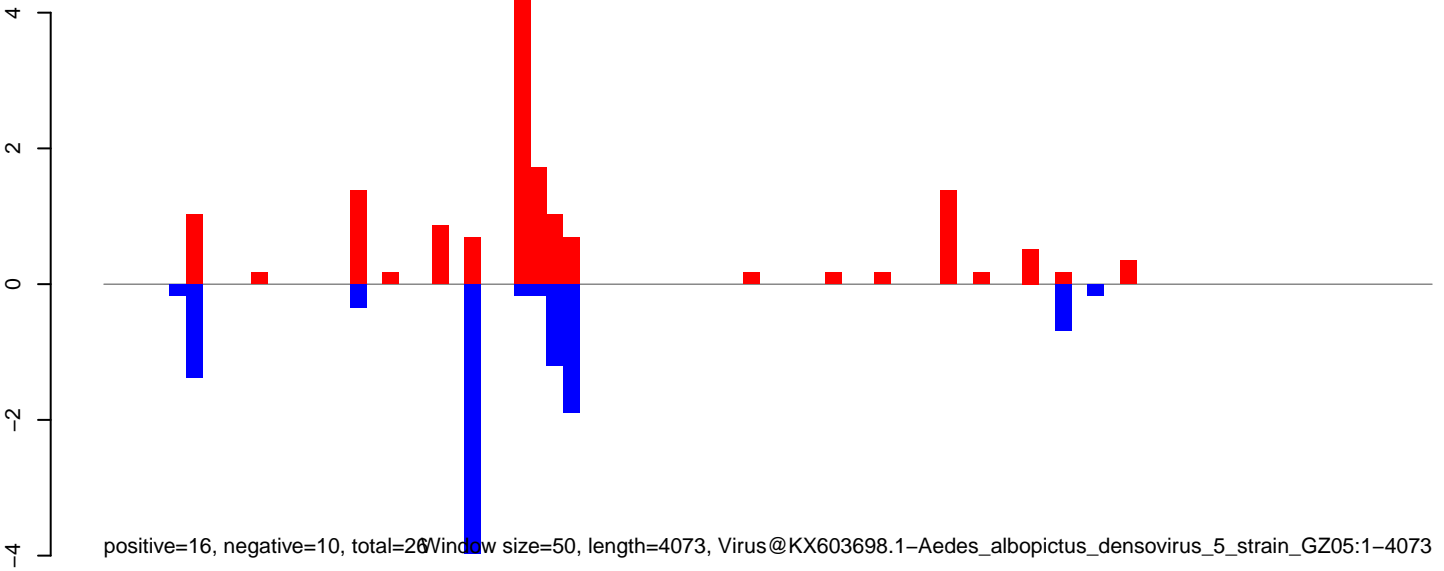
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

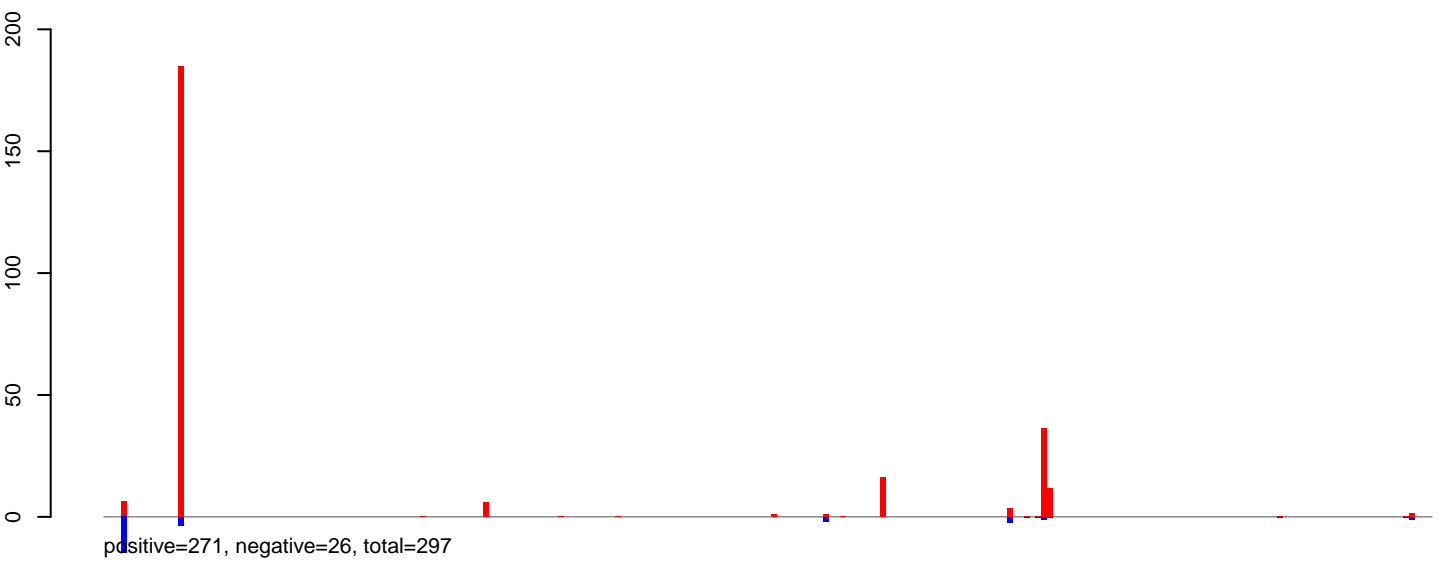


AeAeg_Aag2cells_CHIKV.rep

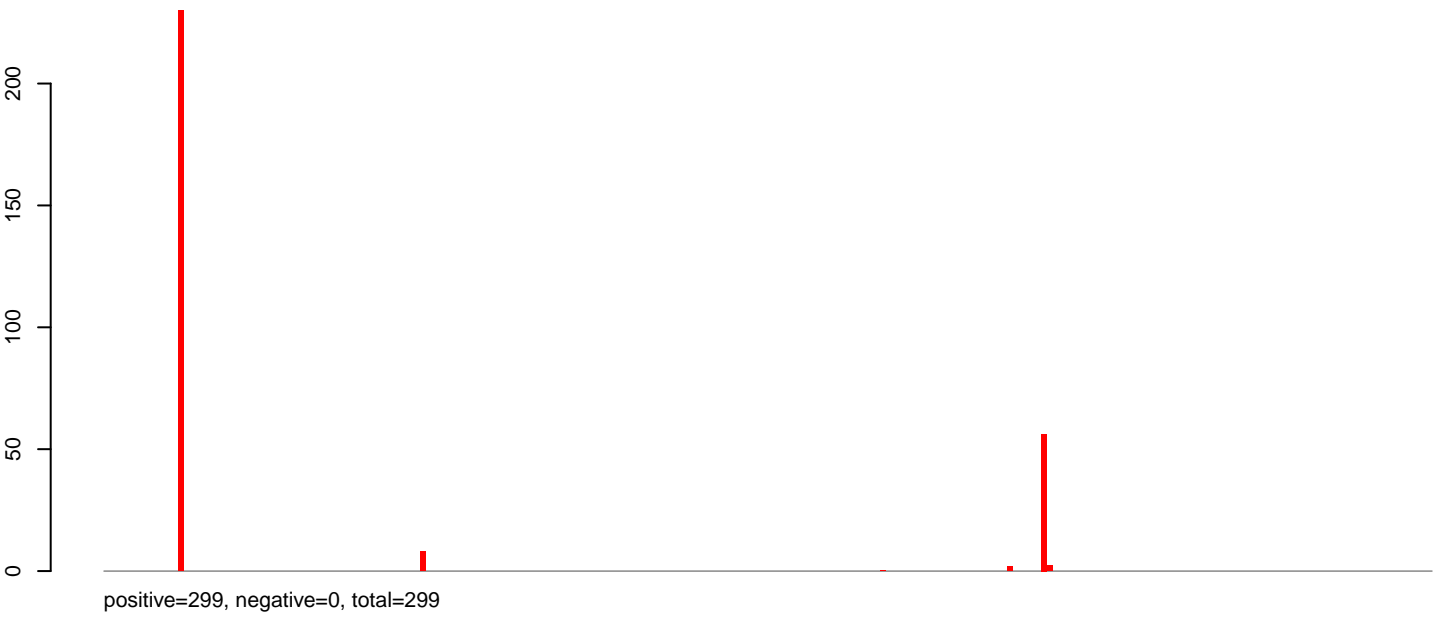


Window size=50, length=4073, Virus@KX603698.1-Aedes albopictus_densovirus_5_strain_GZ05:1-4073

AeAeg_Aag2cells_CHIKV.18_23.rep



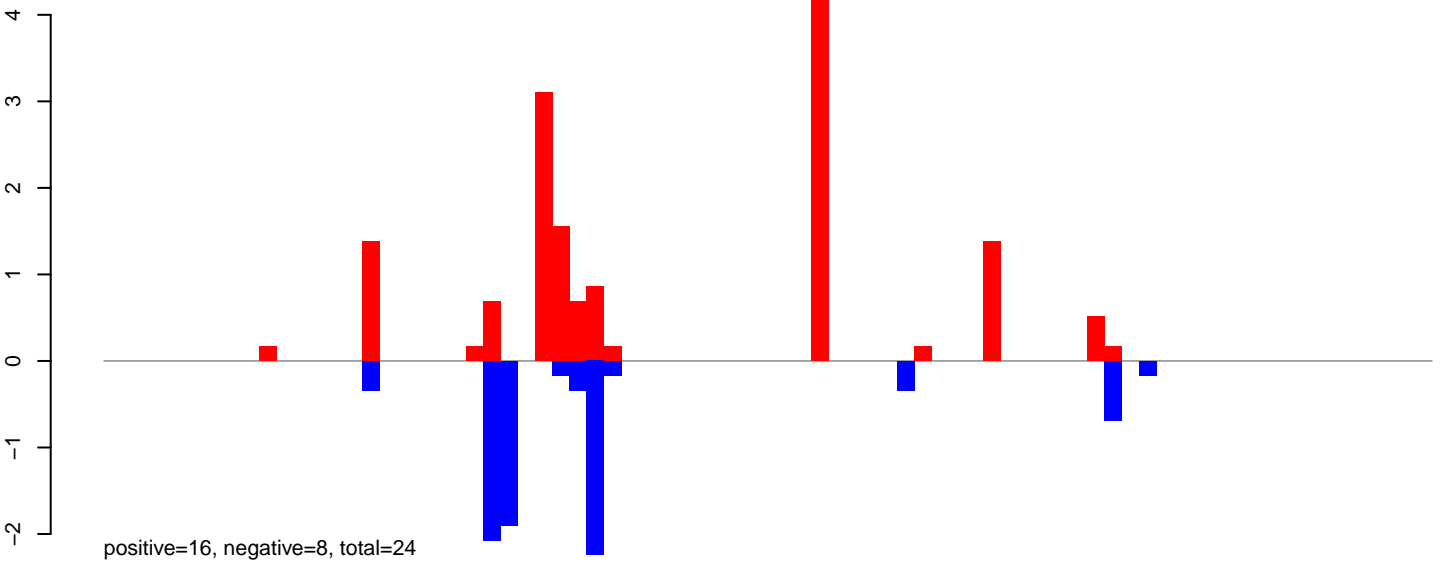
AeAeg_Aag2cells_CHIKV.24_35.rep



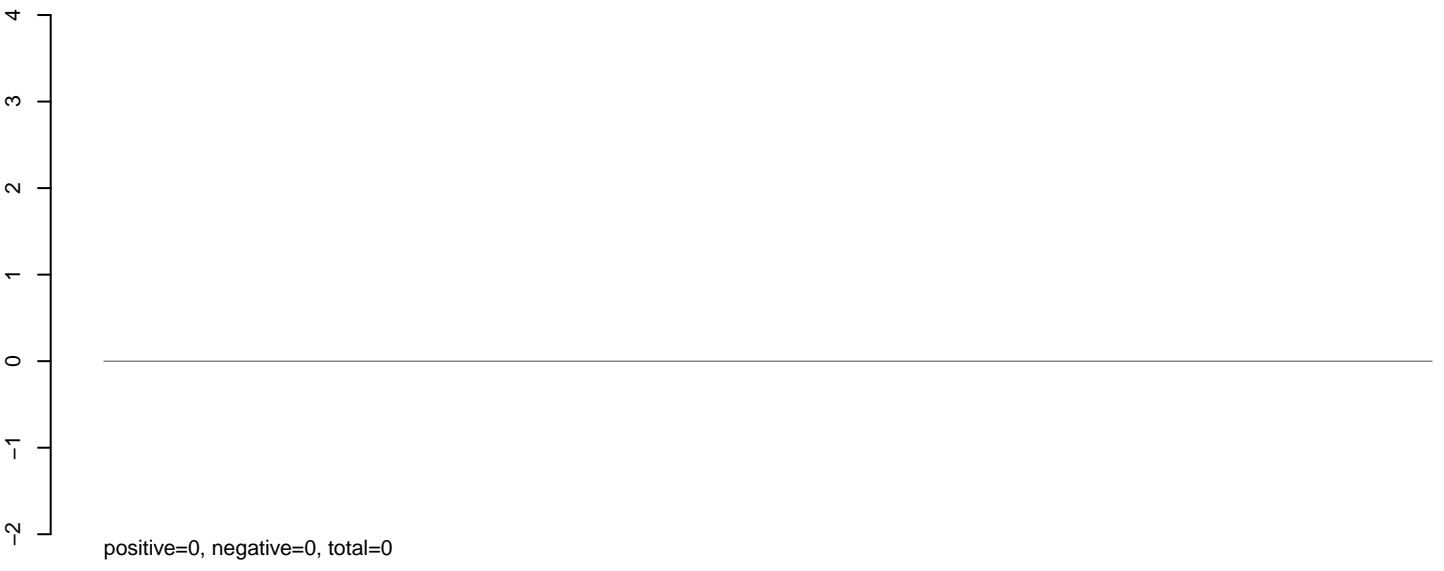
AeAeg_Aag2cells_CHIKV.rep



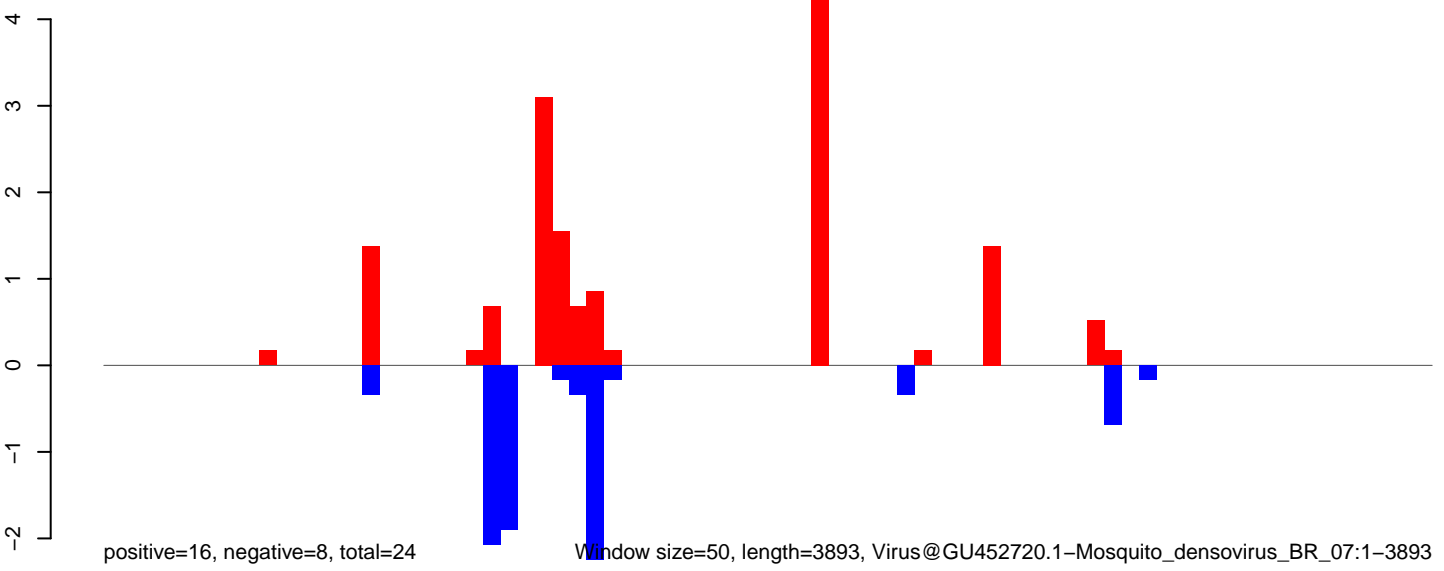
AeAeg_Aag2cells_CHIKV.18_23.rep



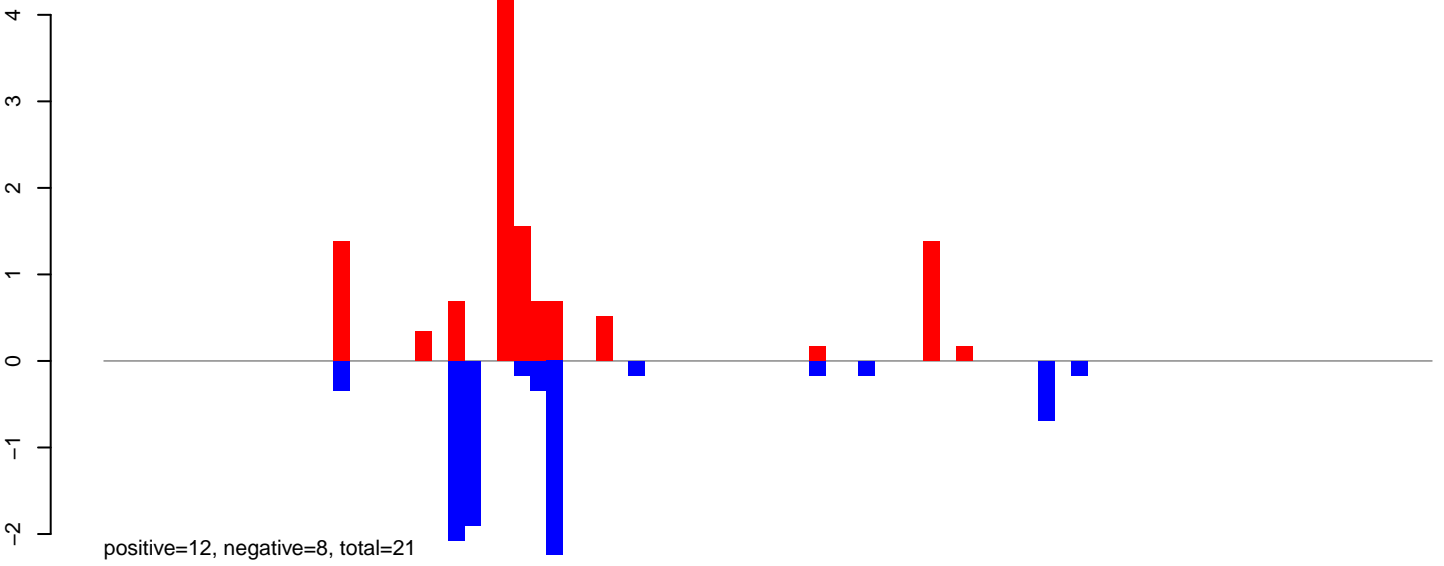
AeAeg_Aag2cells_CHIKV.24_35.rep



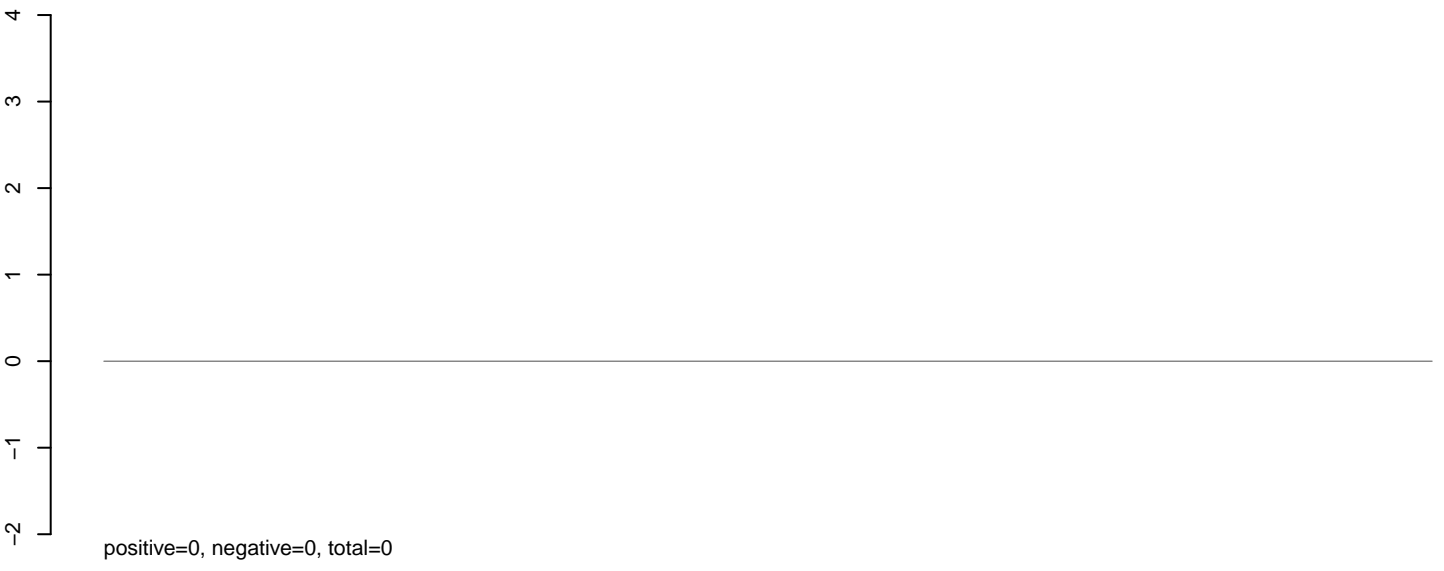
AeAeg_Aag2cells_CHIKV.rep



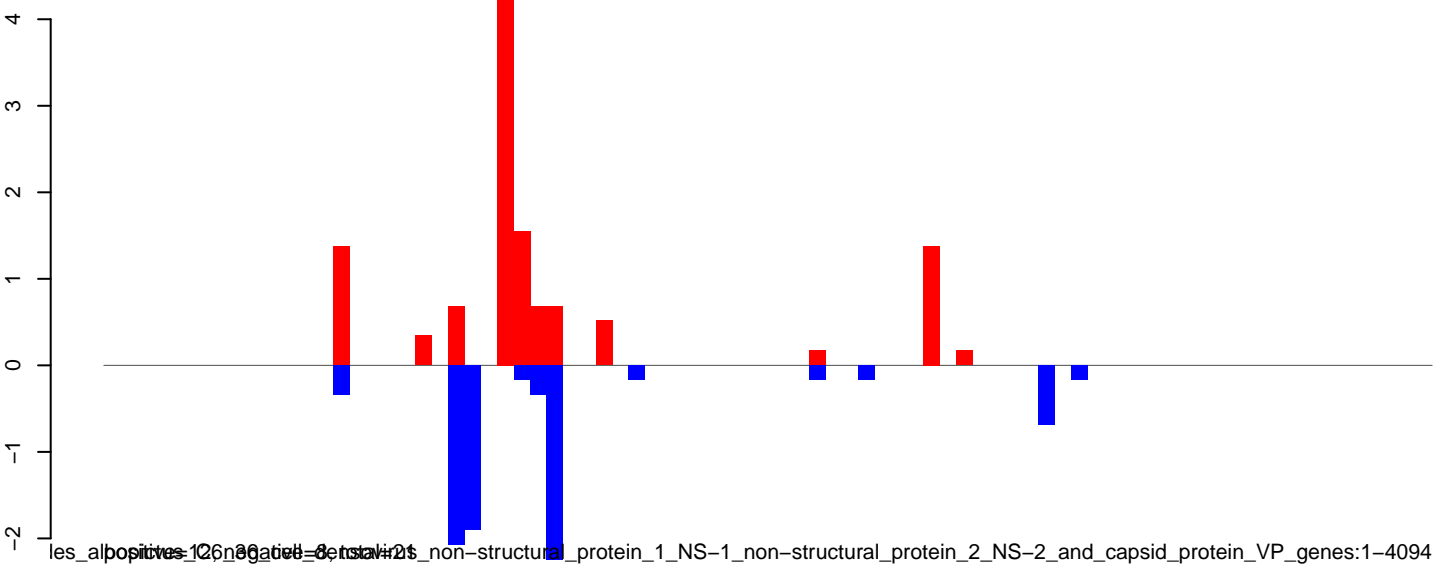
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



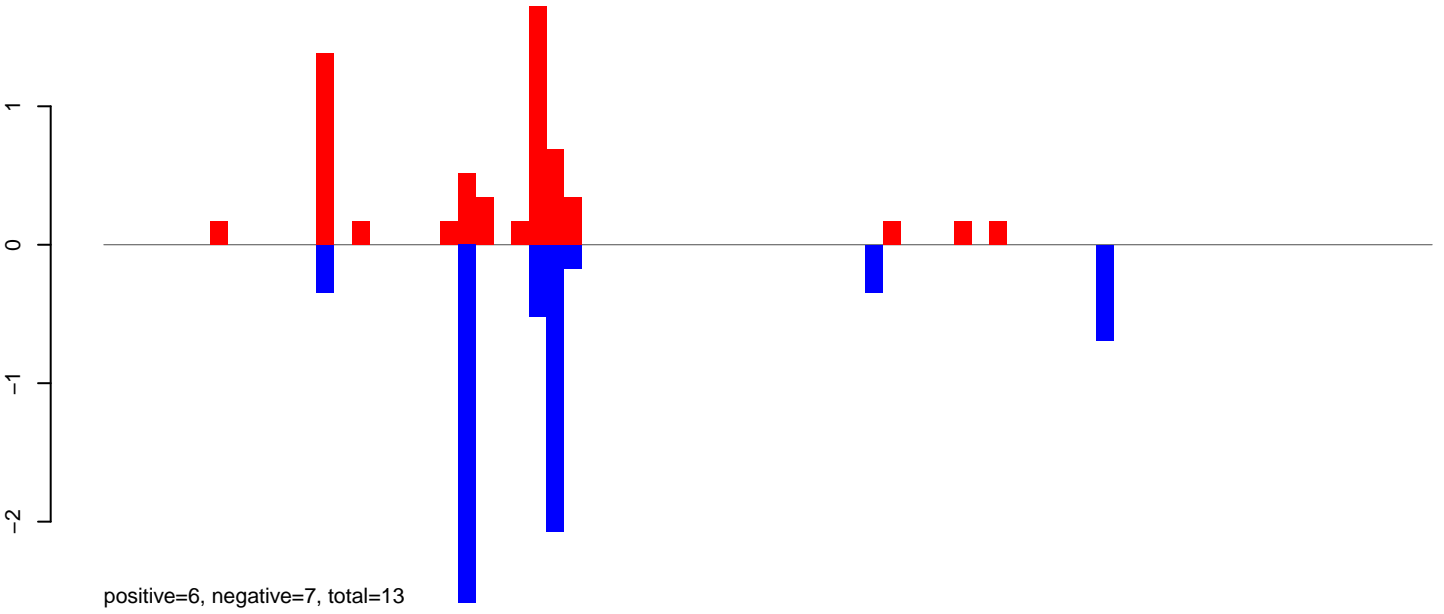
AeAeg_Aag2cells_CHIKV.rep



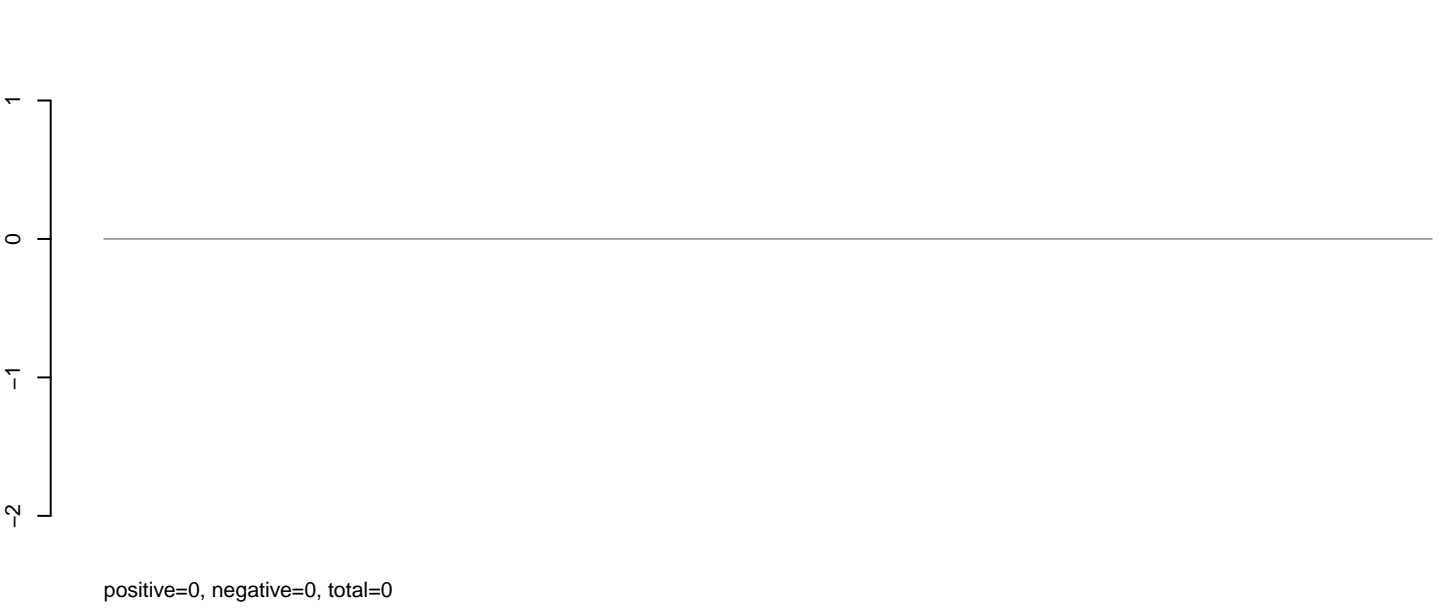
les_alphavirus_126_nucleocapsid_protein_1_NS-1_non-structural_protein_2_NS-2_and_capsid_protein_VP_genes:1-4094

0 1000 2000 3000 4000

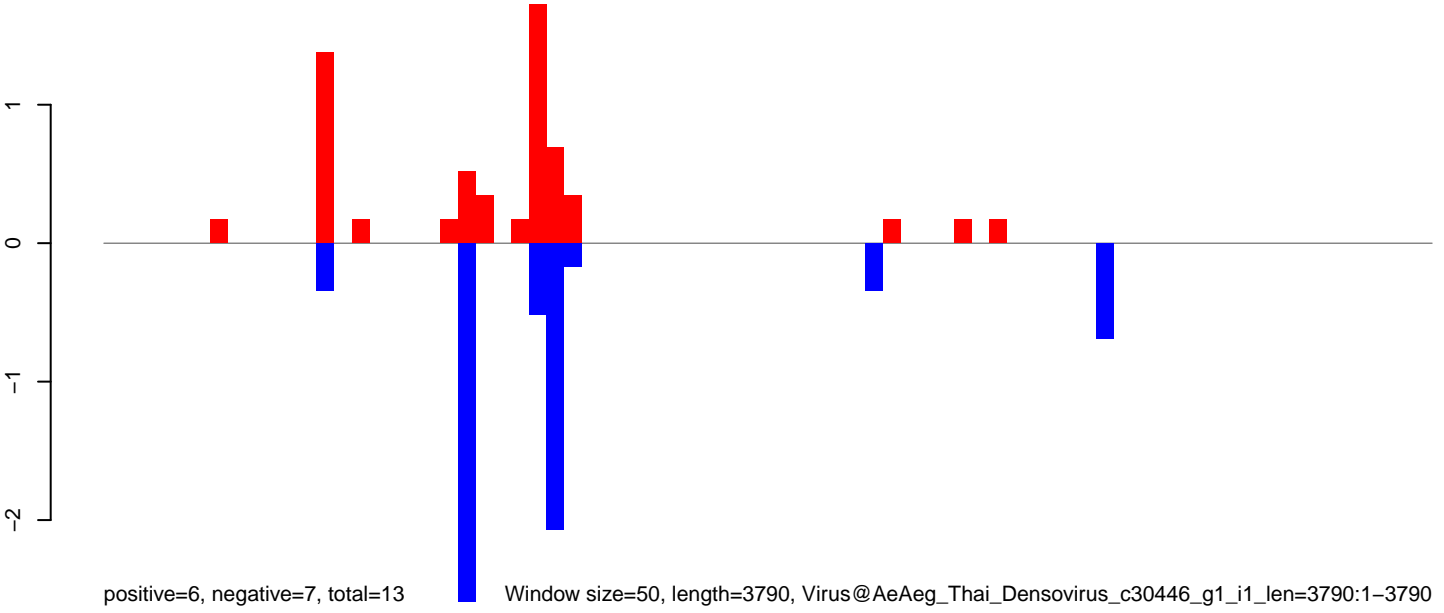
AeAeg_Aag2cells_CHIKV.18_23.rep



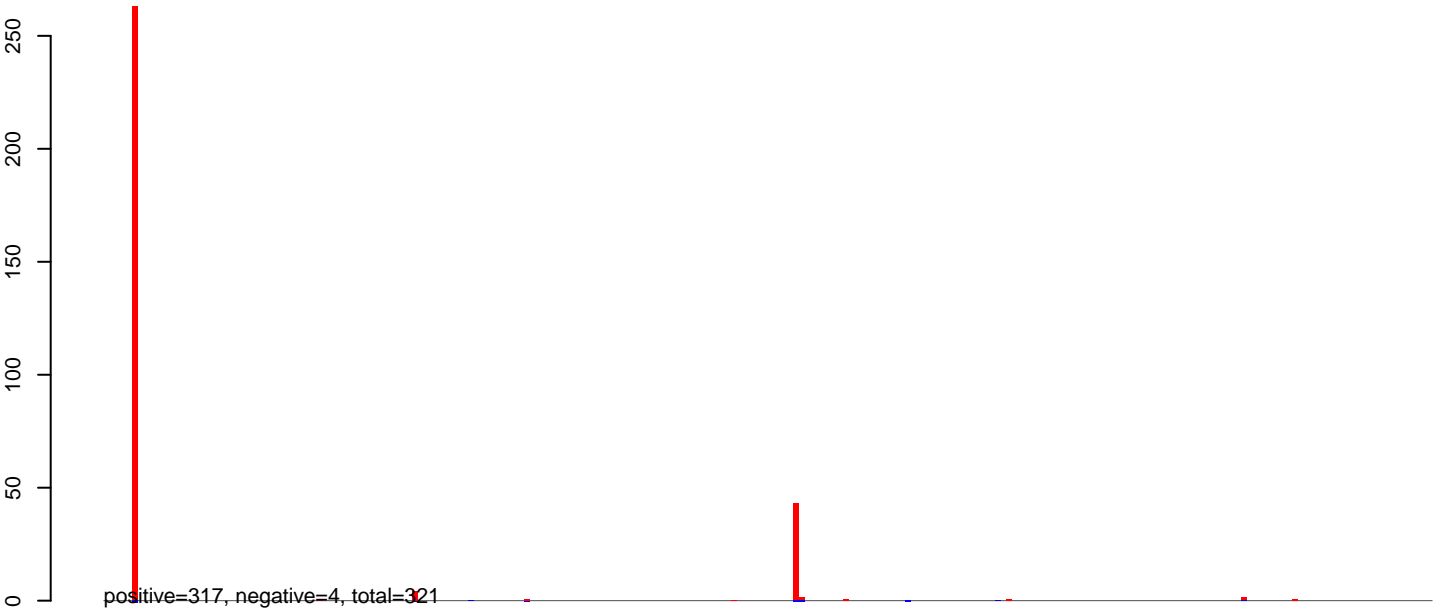
AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep



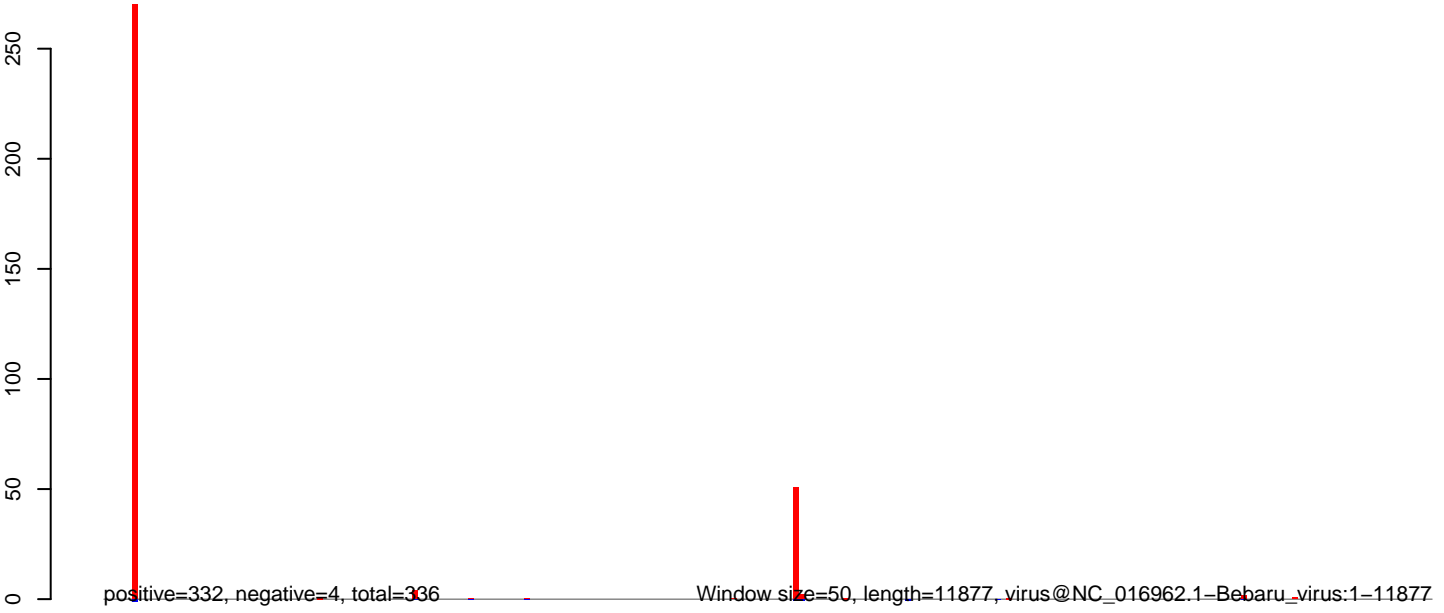
AeAeg_Aag2cells_CHIKV.18_23.rep



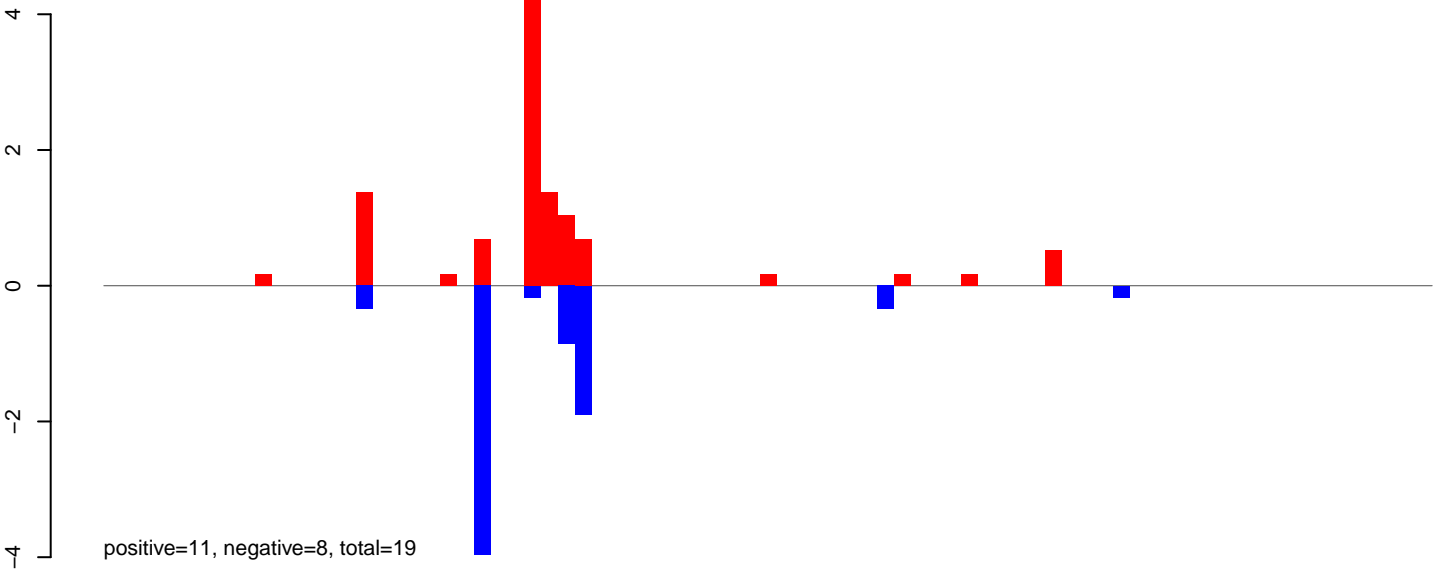
AeAeg_Aag2cells_CHIKV.24_35.rep



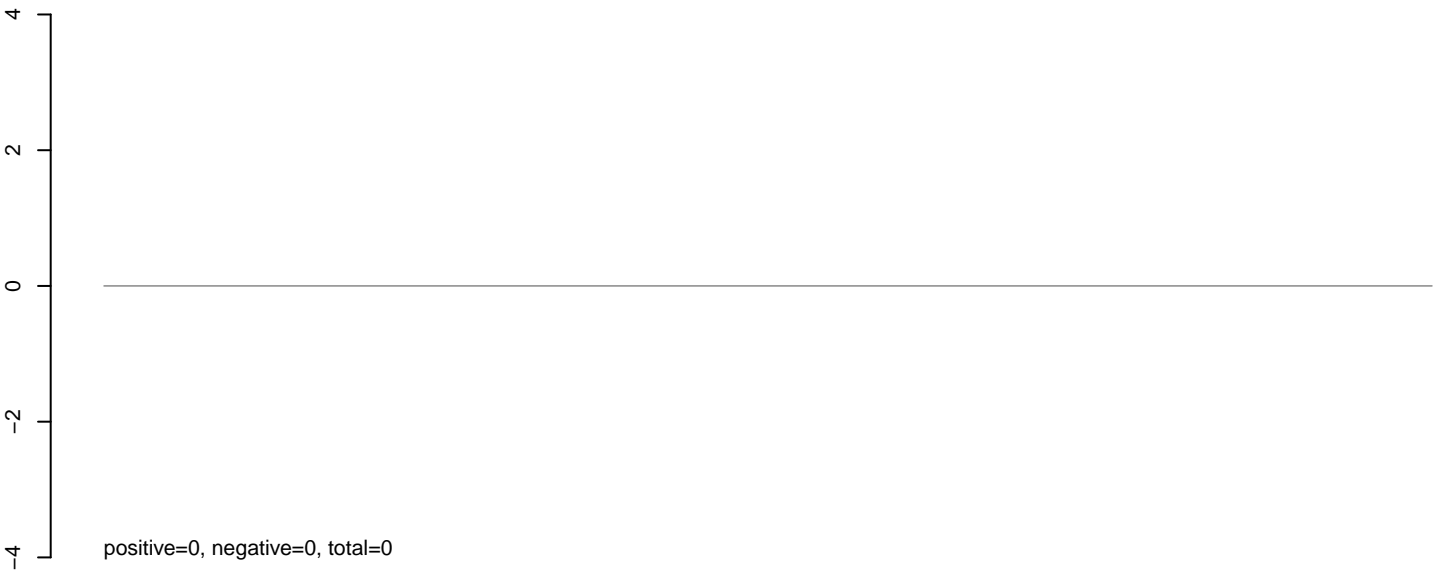
AeAeg_Aag2cells_CHIKV.rep



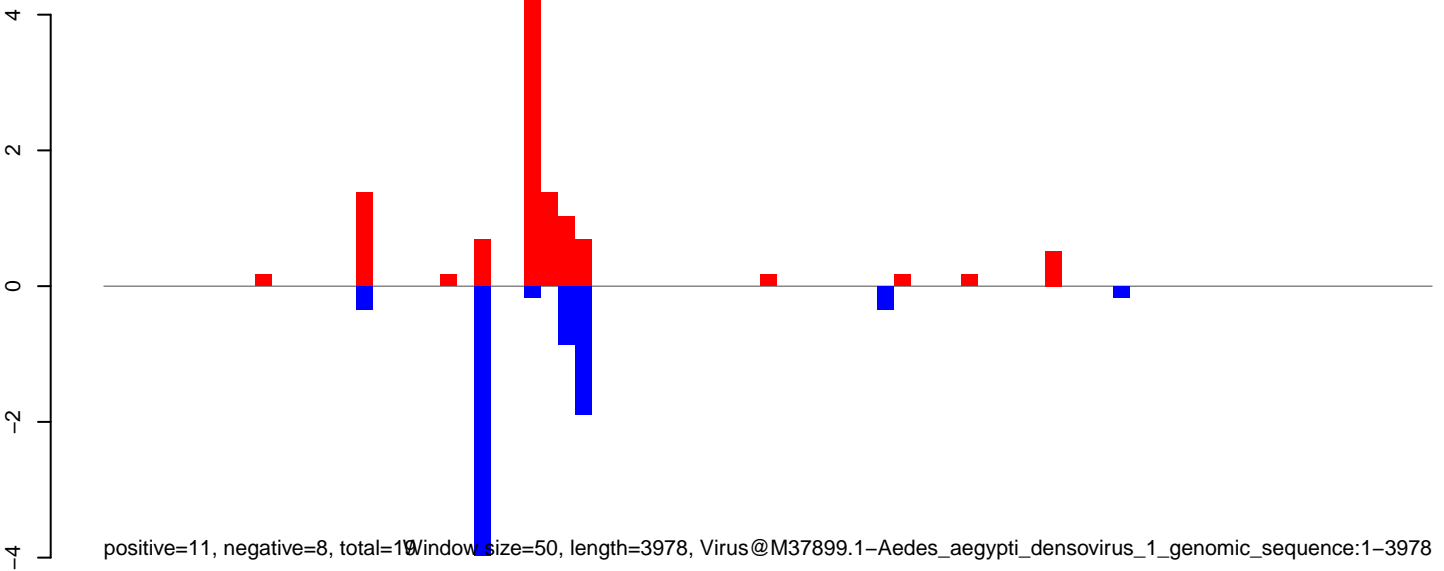
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

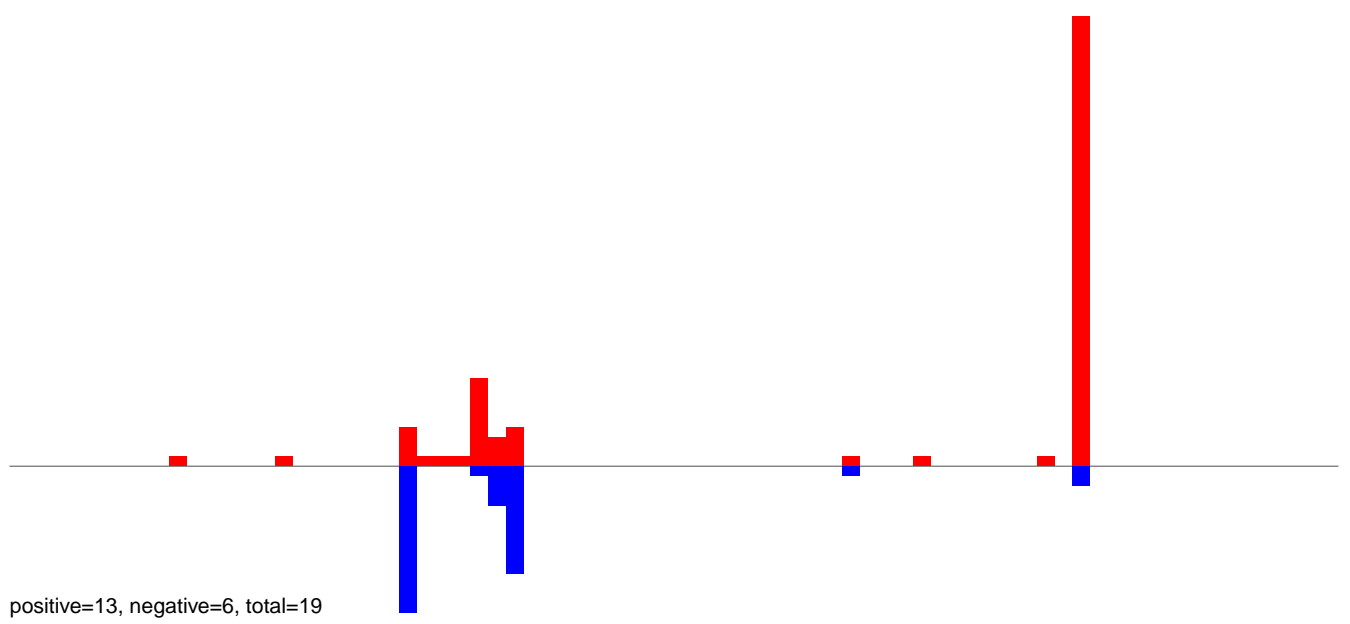


AeAeg_Aag2cells_CHIKV.rep

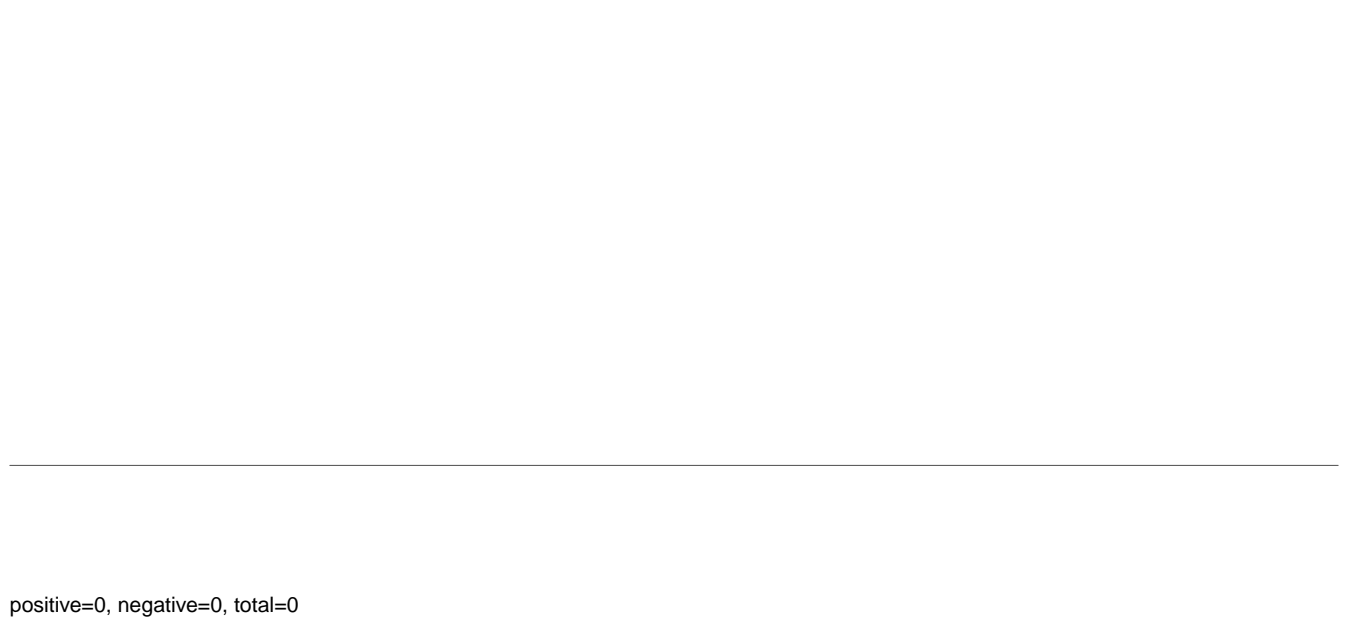


Window size=50, length=3978, Virus@M37899.1-Aedes_aegypti_densovirus_1_genomic_sequence:1-3978

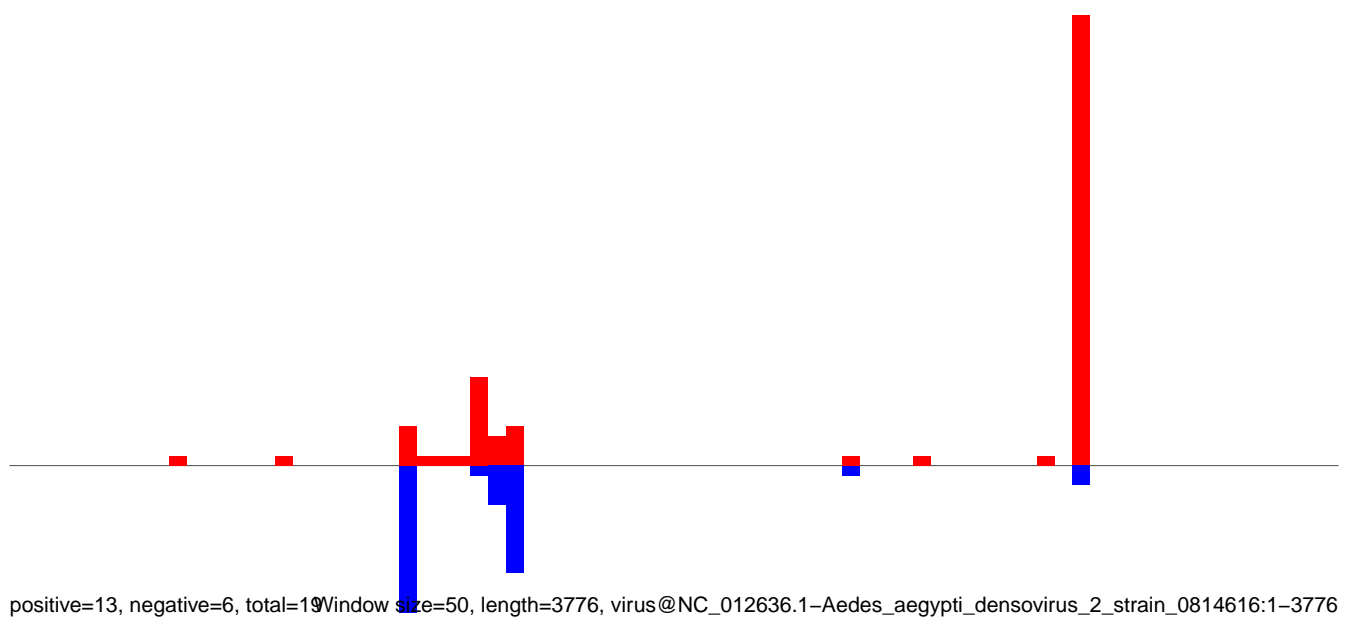
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

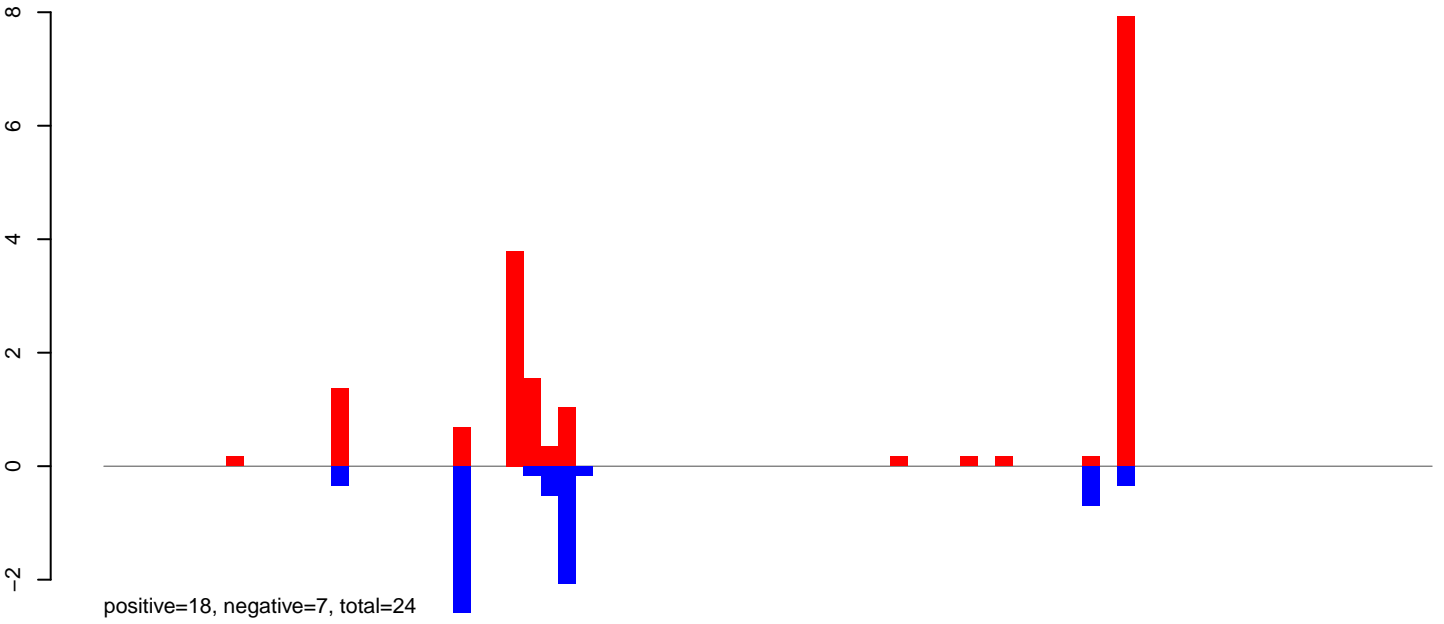


AeAeg_Aag2cells_CHIKV.rep



Window size=50, length=3776, virus@NC_012636.1-Aedes_aegypti_densovirus_2_strain_0814616:1-3776

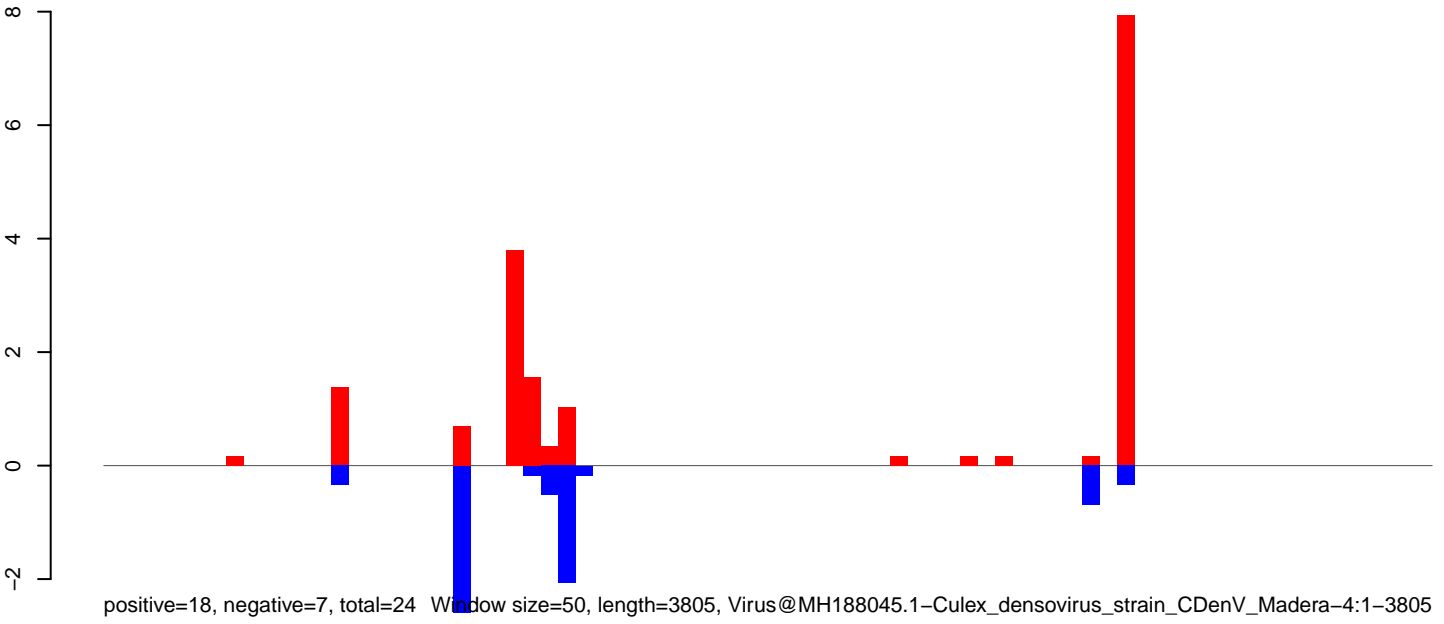
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



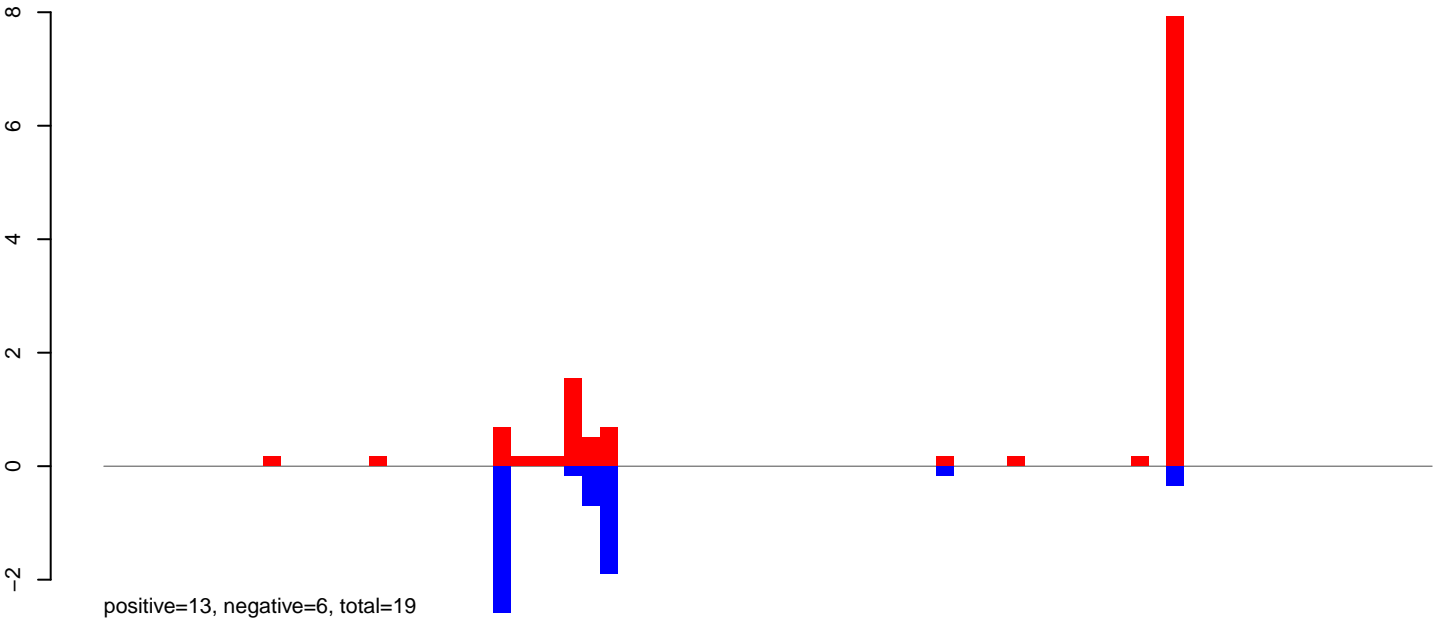
AeAeg_Aag2cells_CHIKV.rep



Window size=50, length=3805, Virus@MH188045.1-Culex_densovirus_strain_CDenV_Madera-4:1-3805

0 1000 2000 3000 4000

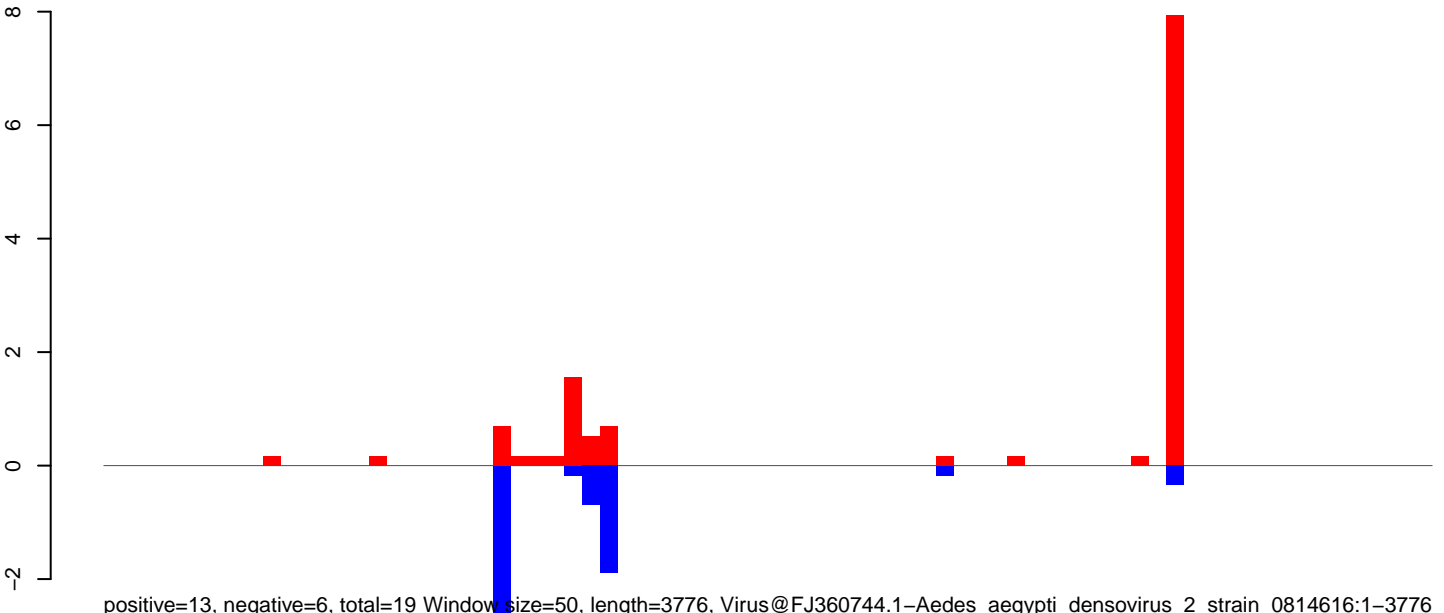
AeAeg_Aag2cells_CHIKV.18_23.rep



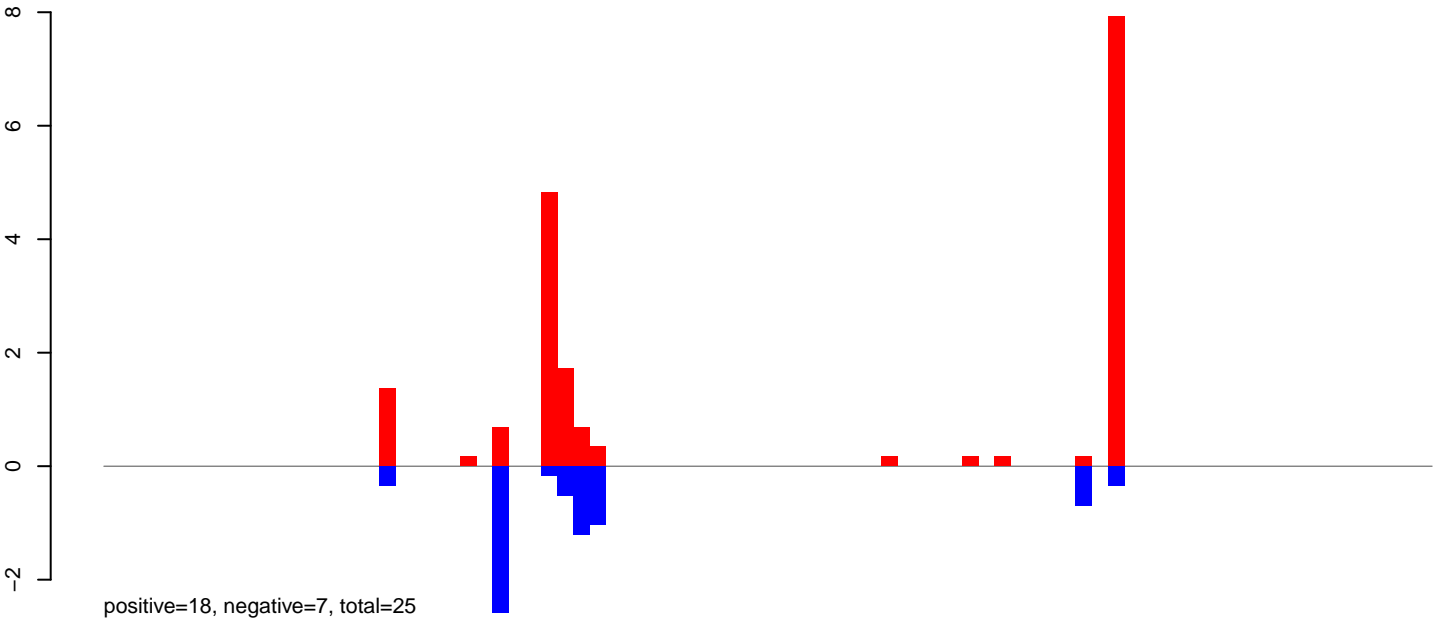
AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep



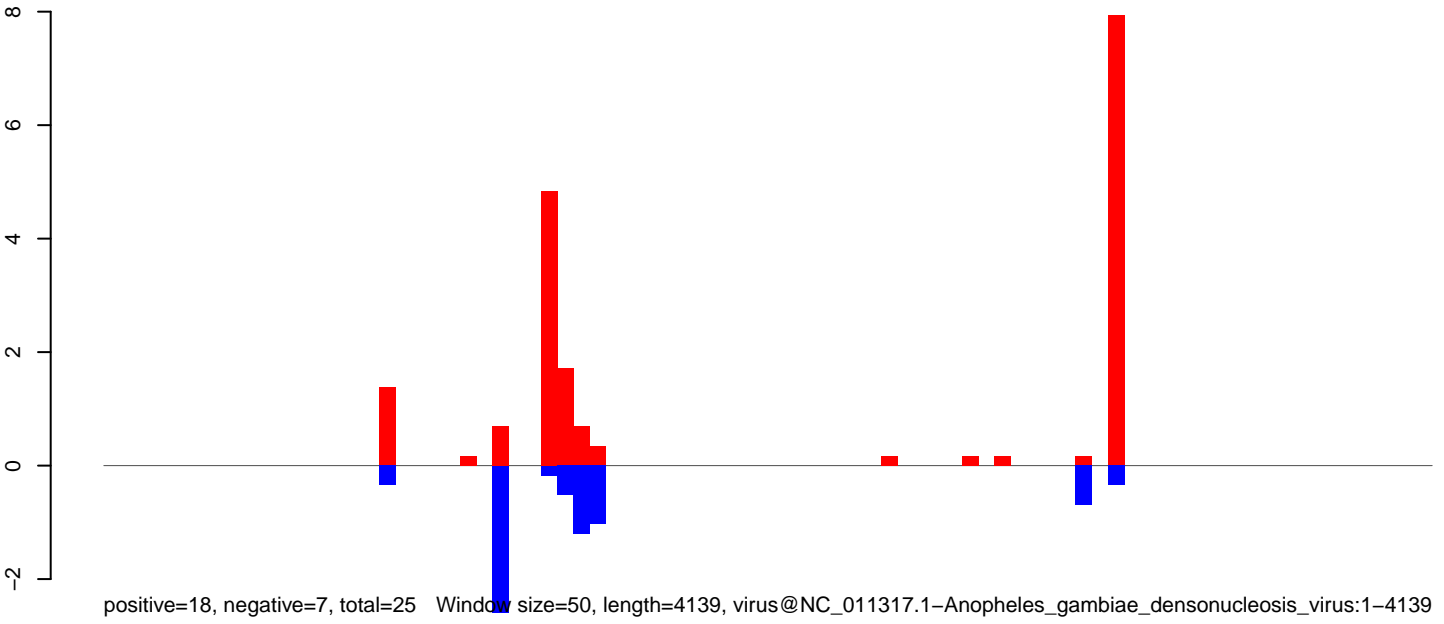
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



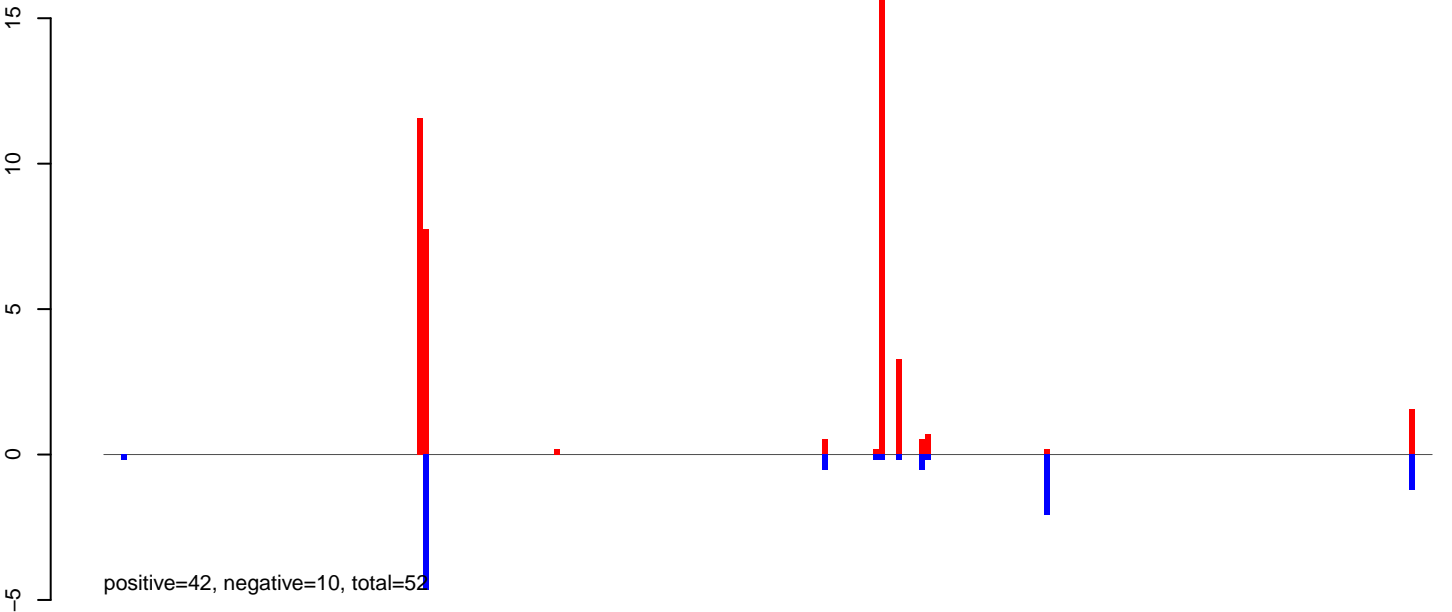
AeAeg_Aag2cells_CHIKV.rep



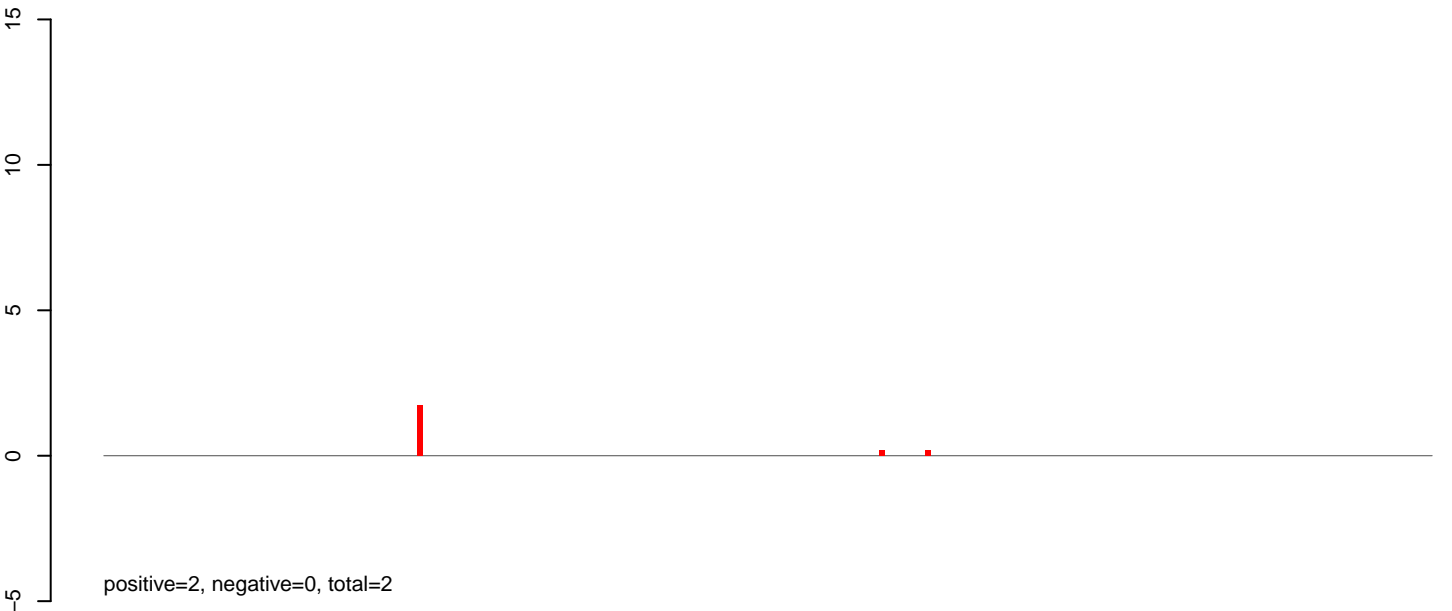
Window size=50, length=4139, virus@NC_011317.1-Anopheles_gambiae_densonucleosis_virus:1-4139

0 1000 2000 3000 4000

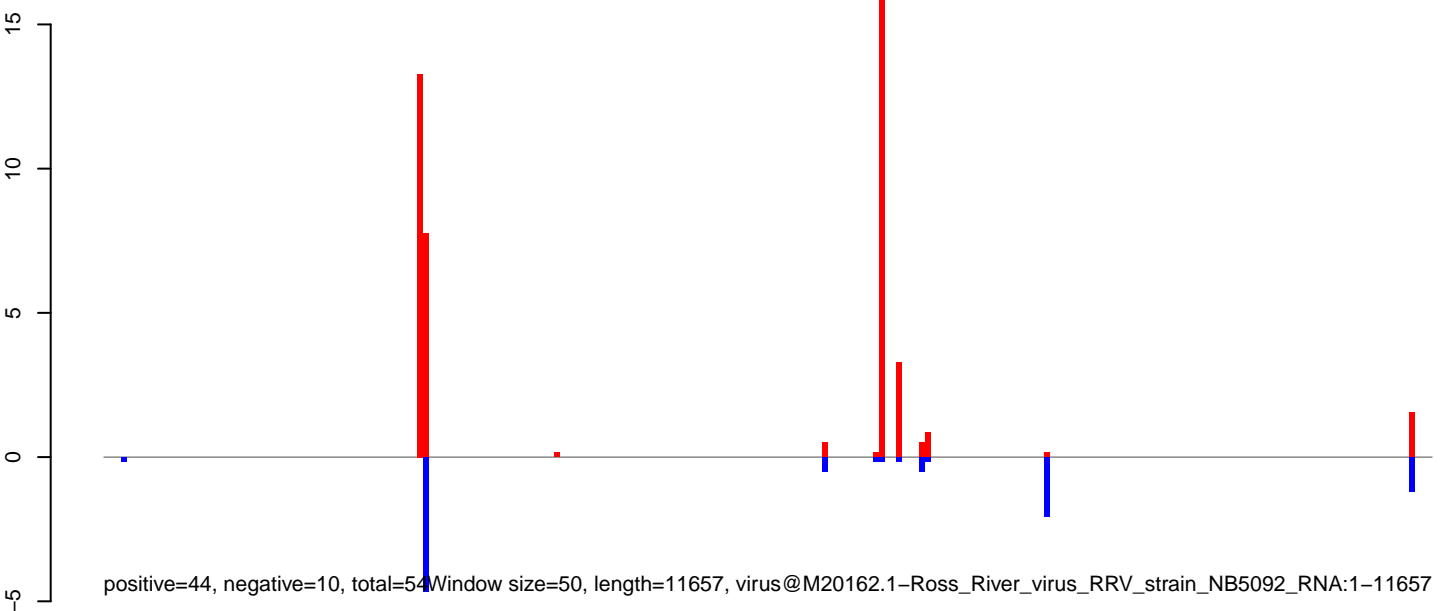
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

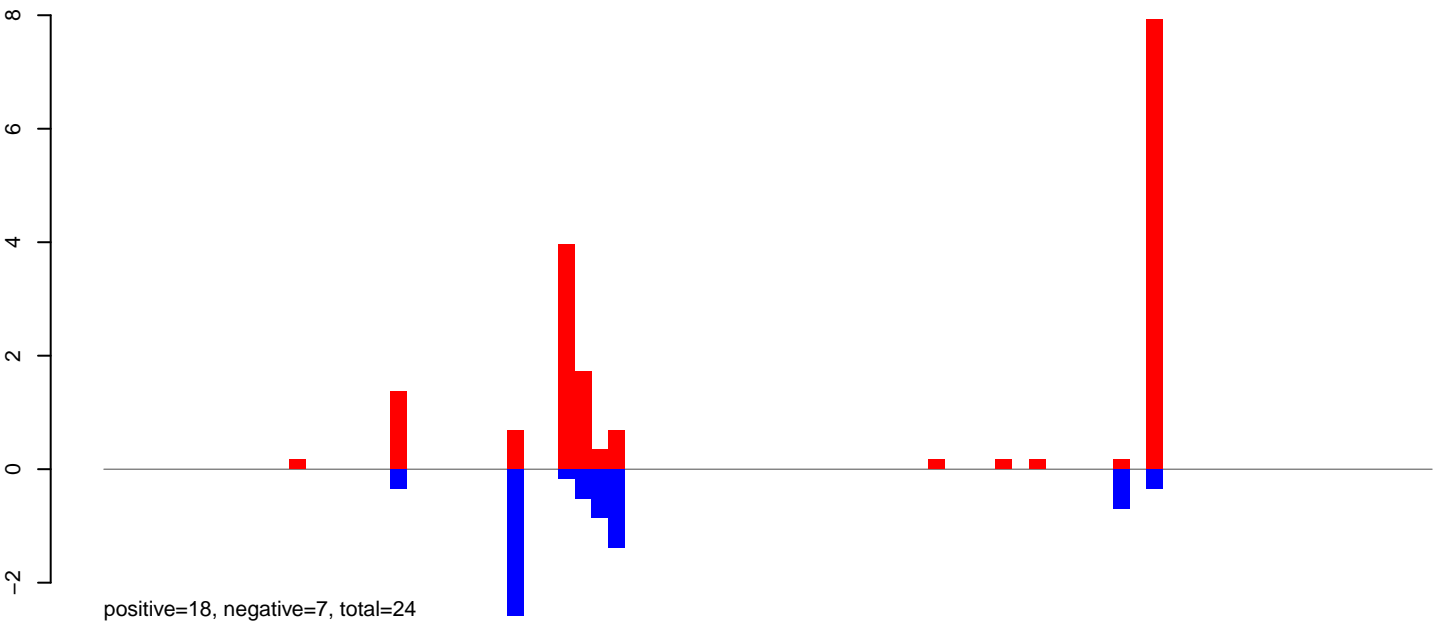


AeAeg_Aag2cells_CHIKV.rep



Window size=50, length=11657, virus@M20162.1-Ross_River_virus_RRV_strain_NB5092_RNA:1-11657

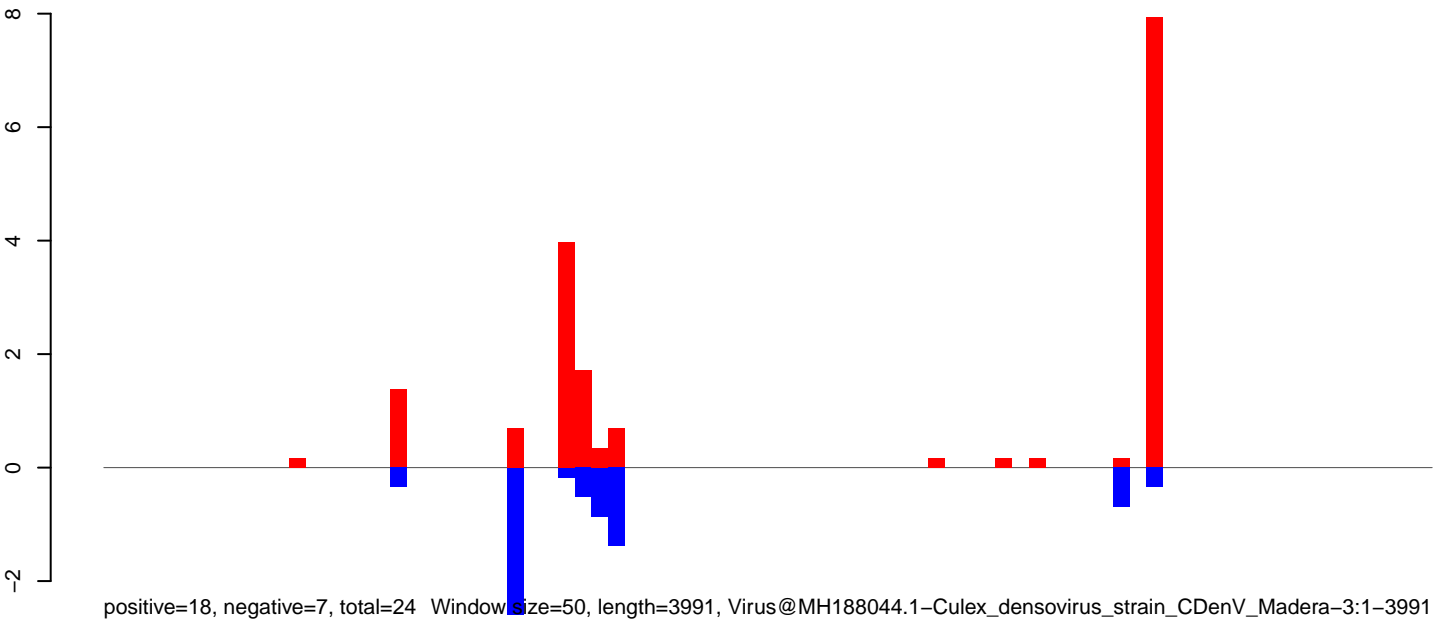
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

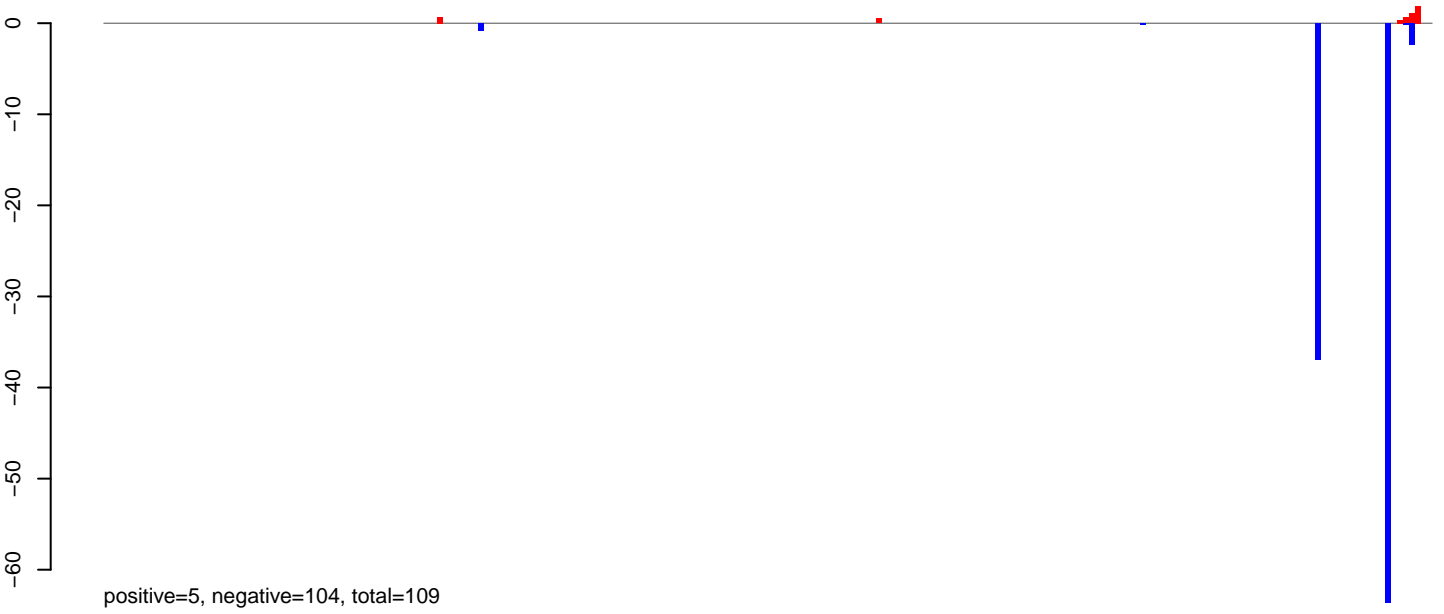


AeAeg_Aag2cells_CHIKV.rep

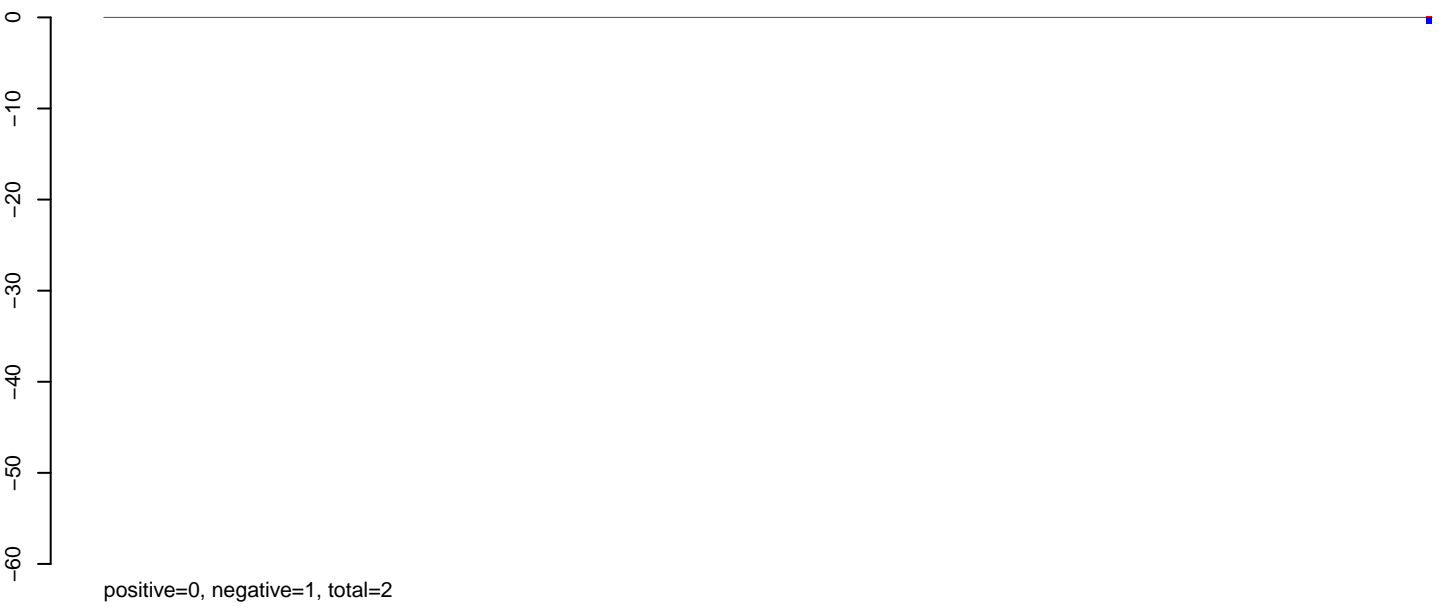


Window Size=50, length=3991, Virus@MH188044.1-Culex_densovirus_strain_CDenV_Madera-3:1-3991

AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



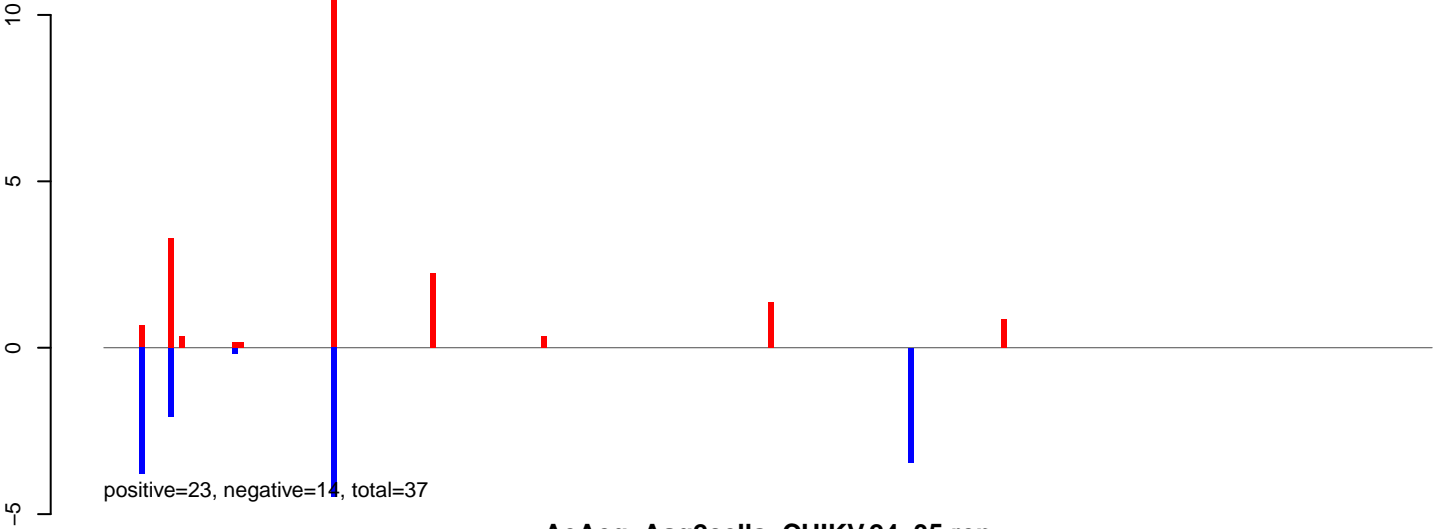
AeAeg_Aag2cells_CHIKV.rep



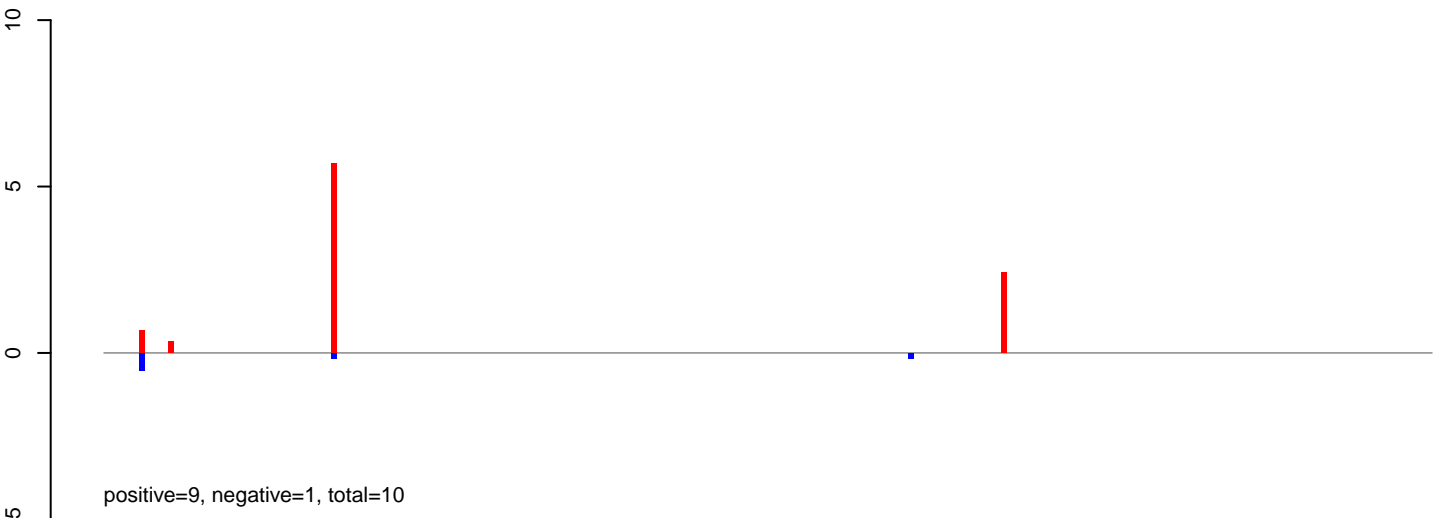
Window size=50, length=11375, virus@NC_005064.1-Kamiti_River_virus:1-11375

0 2000 4000 6000 8000 10000

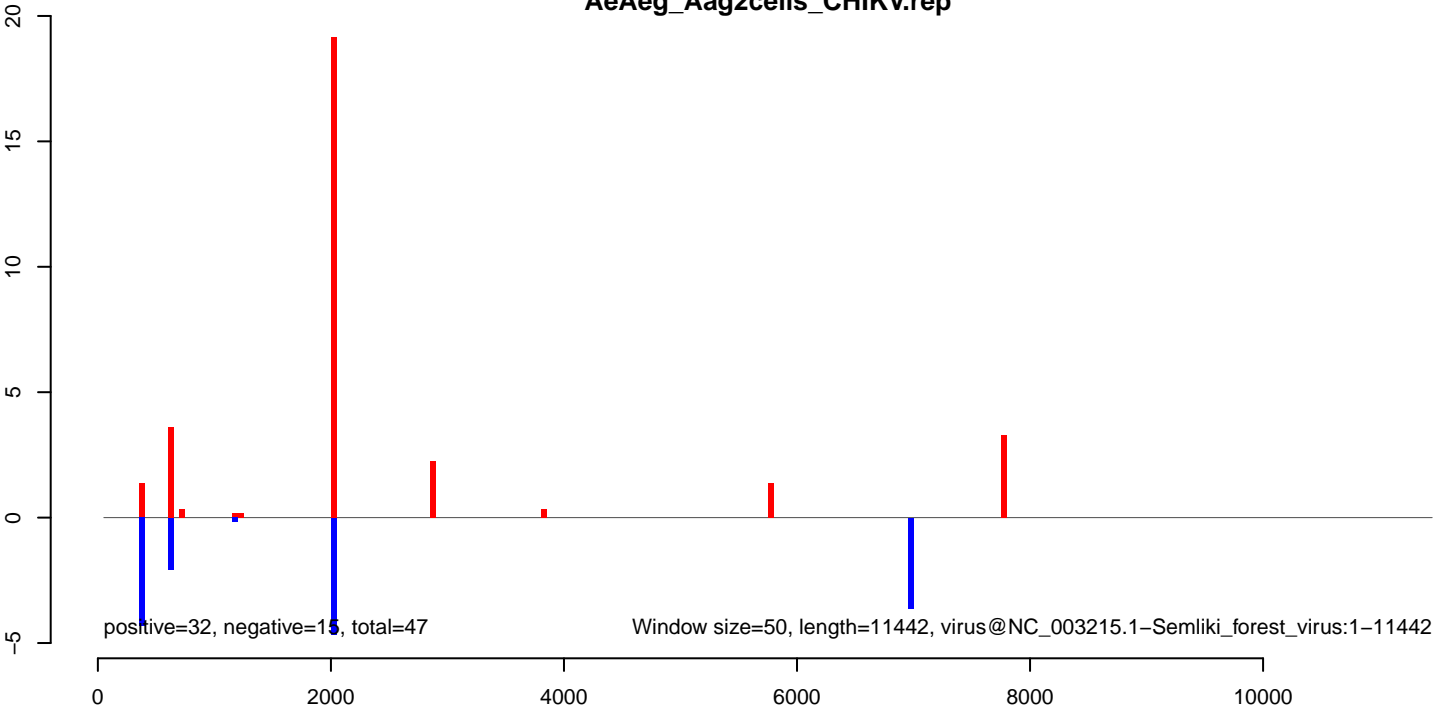
AeAeg_Aag2cells_CHIKV.18_23.rep



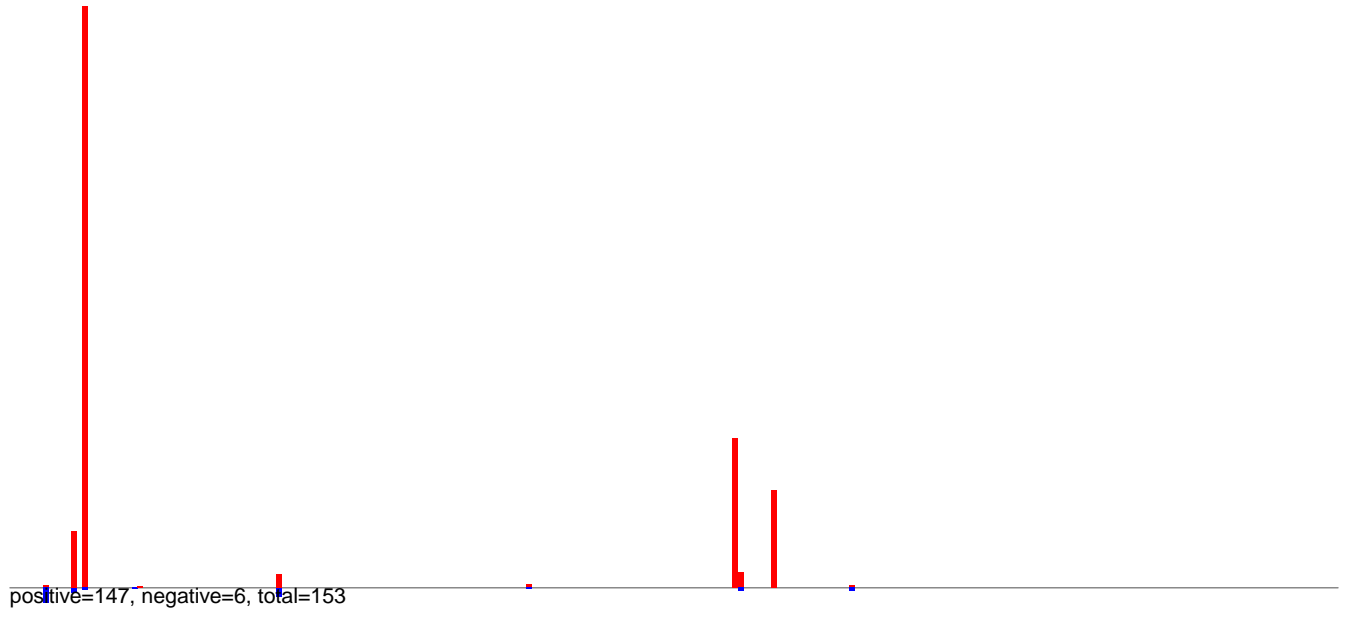
AeAeg_Aag2cells_CHIKV.24_35.rep



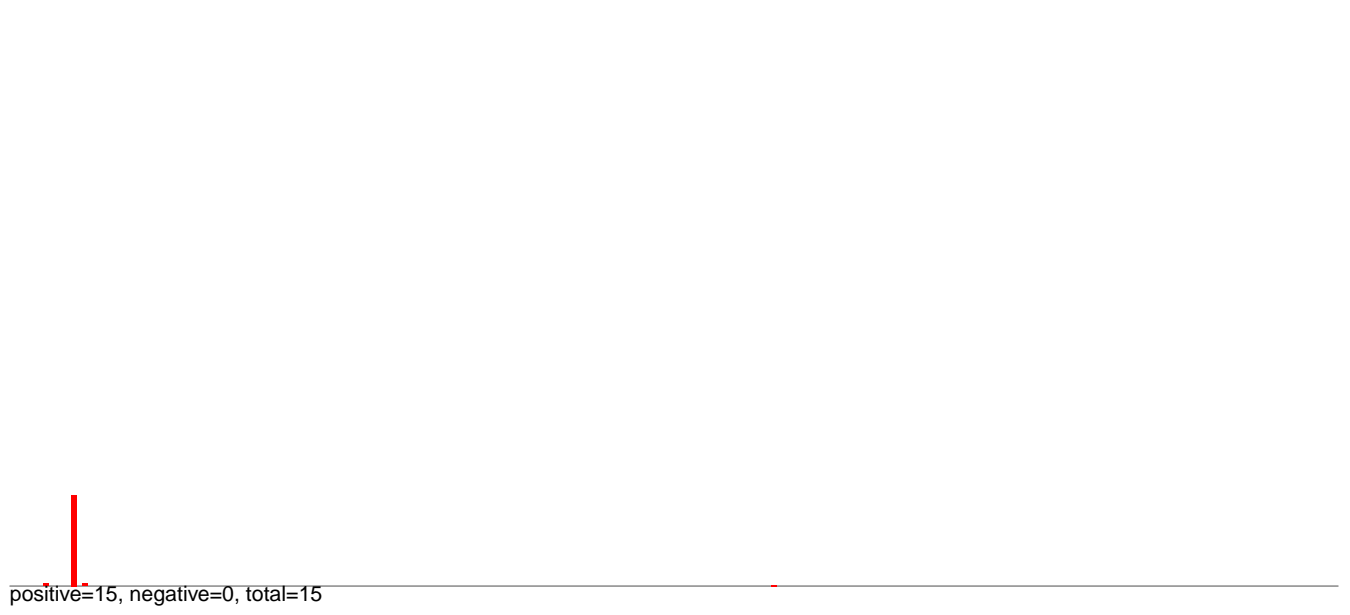
AeAeg_Aag2cells_CHIKV.rep



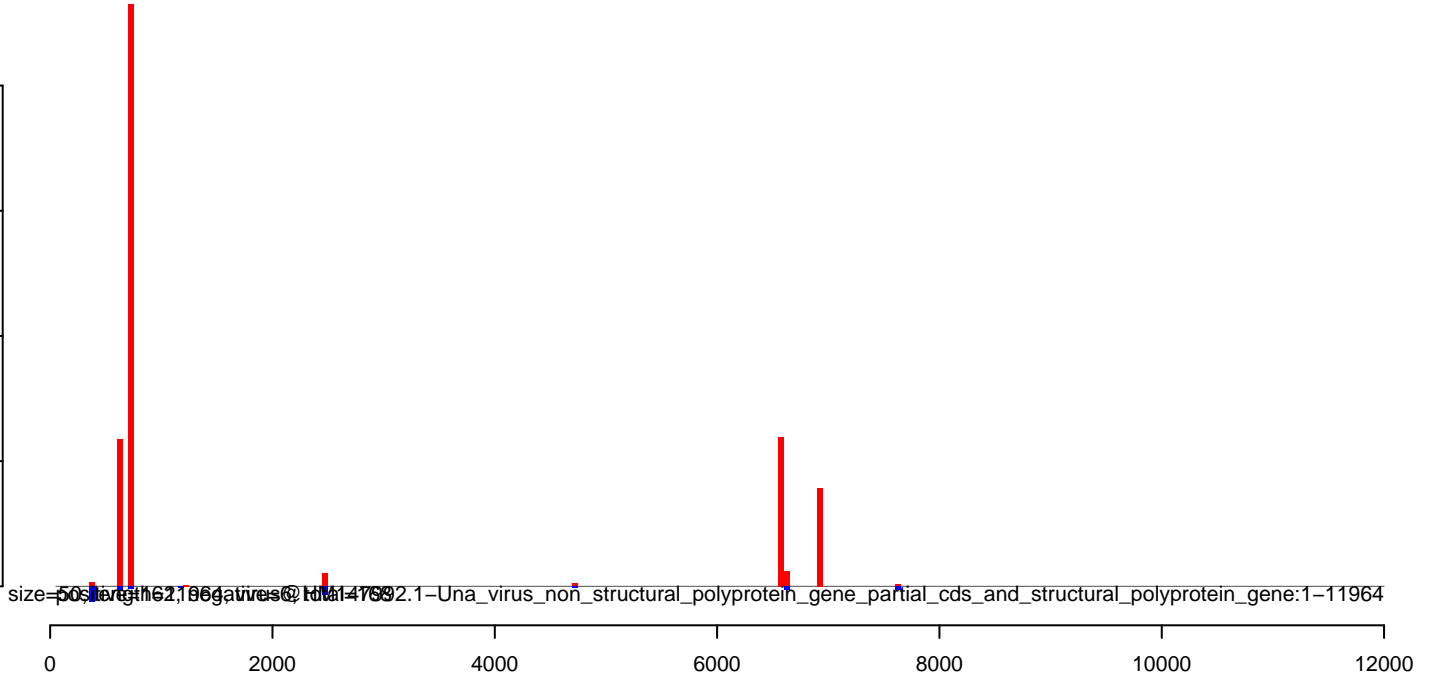
AeAeg_Aag2cells_CHIKV.18_23.rep



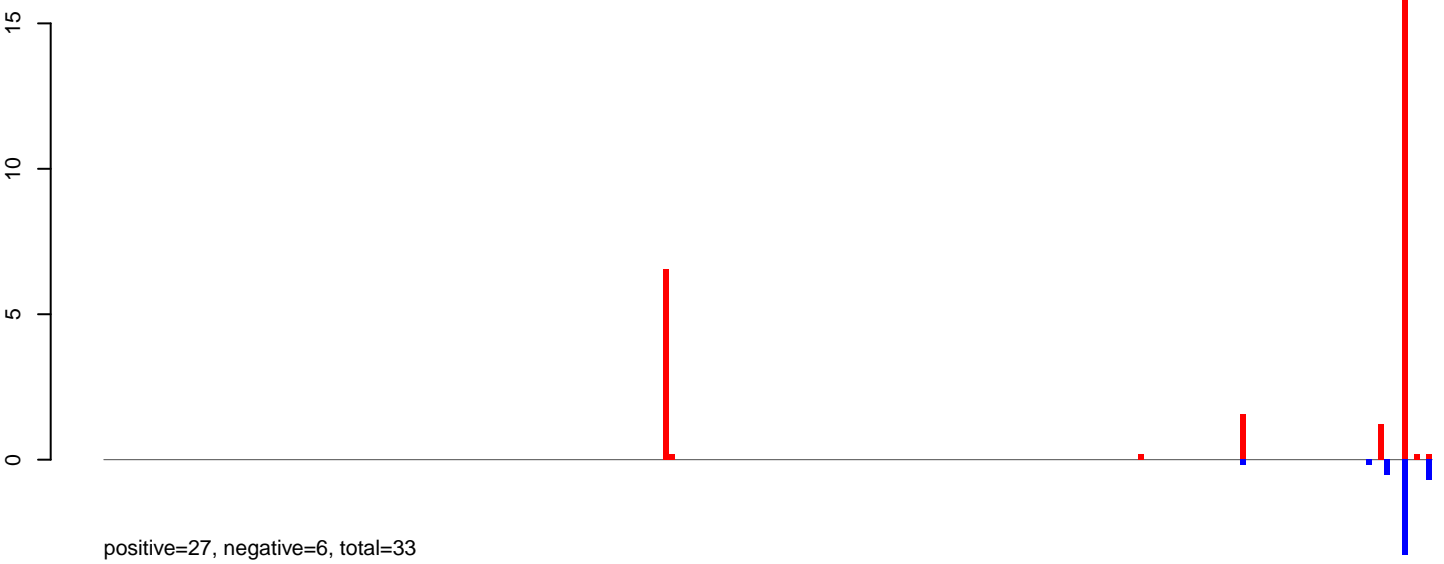
AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep



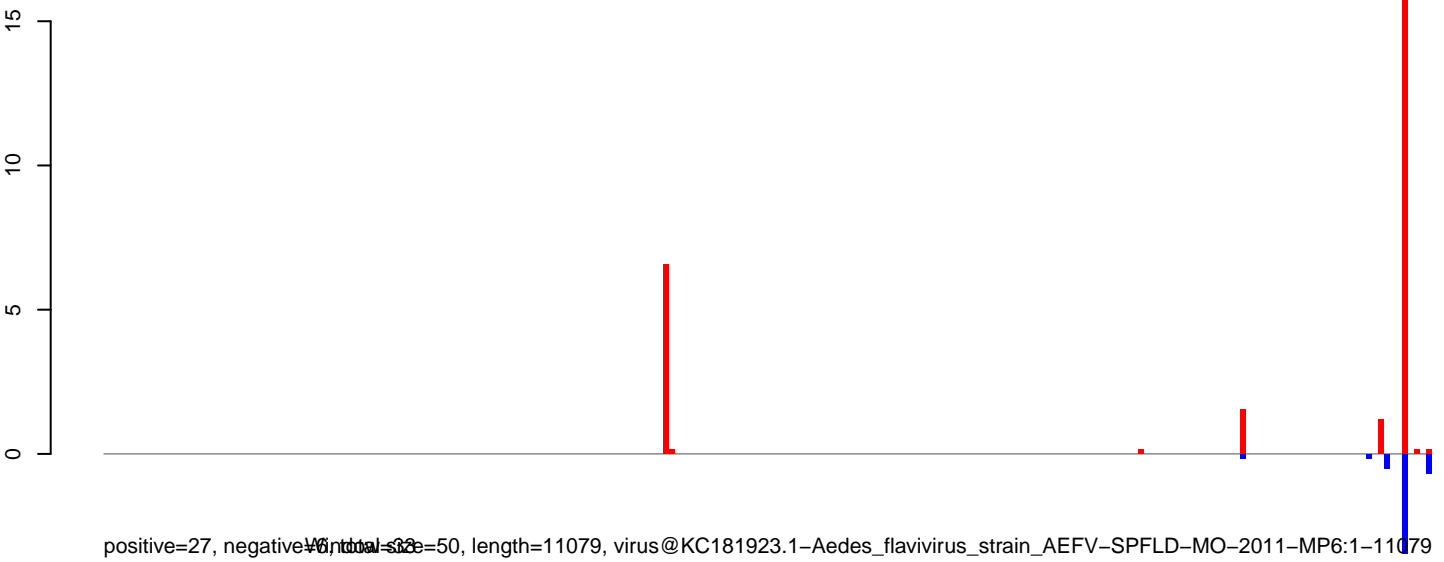
AeAeg_Aag2cells_CHIKV.18_23.rep



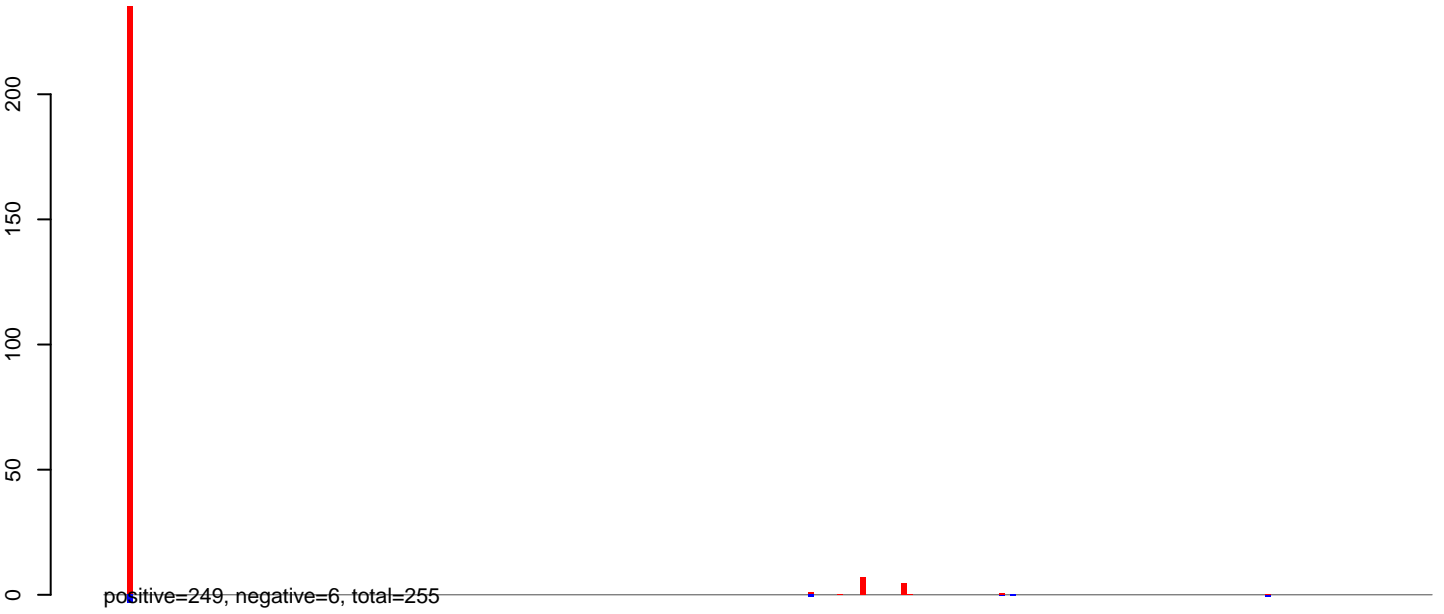
AeAeg_Aag2cells_CHIKV.24_35.rep



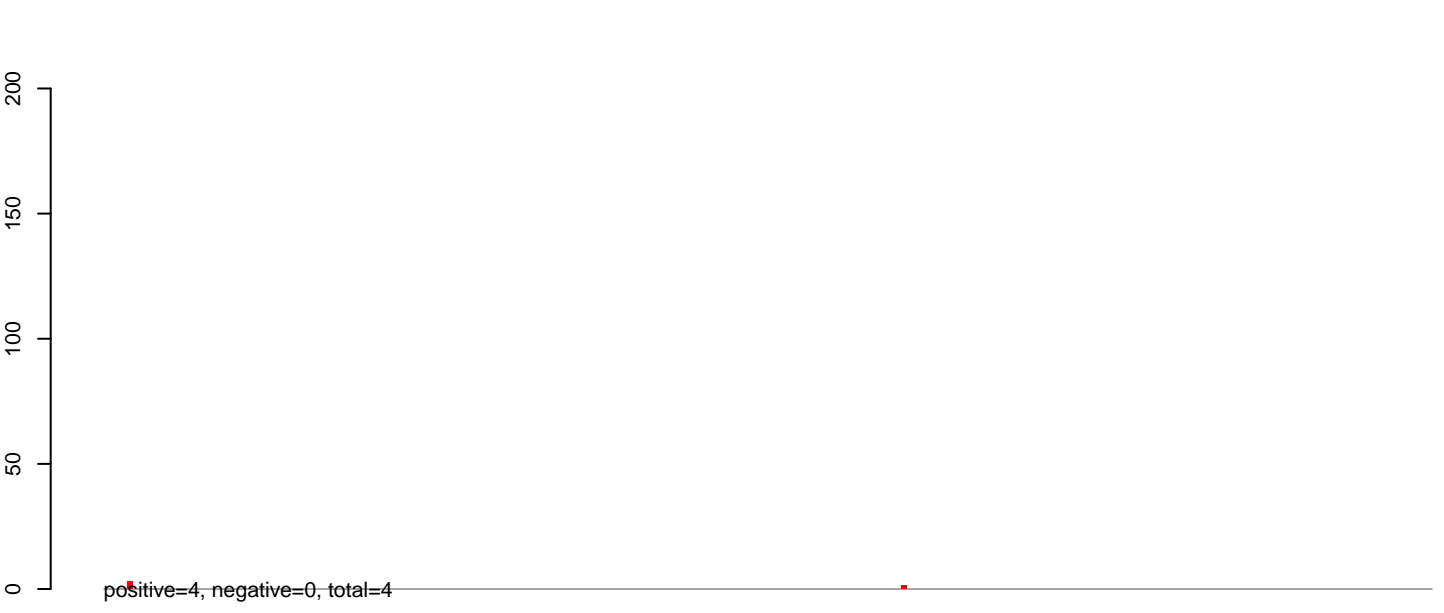
AeAeg_Aag2cells_CHIKV.rep



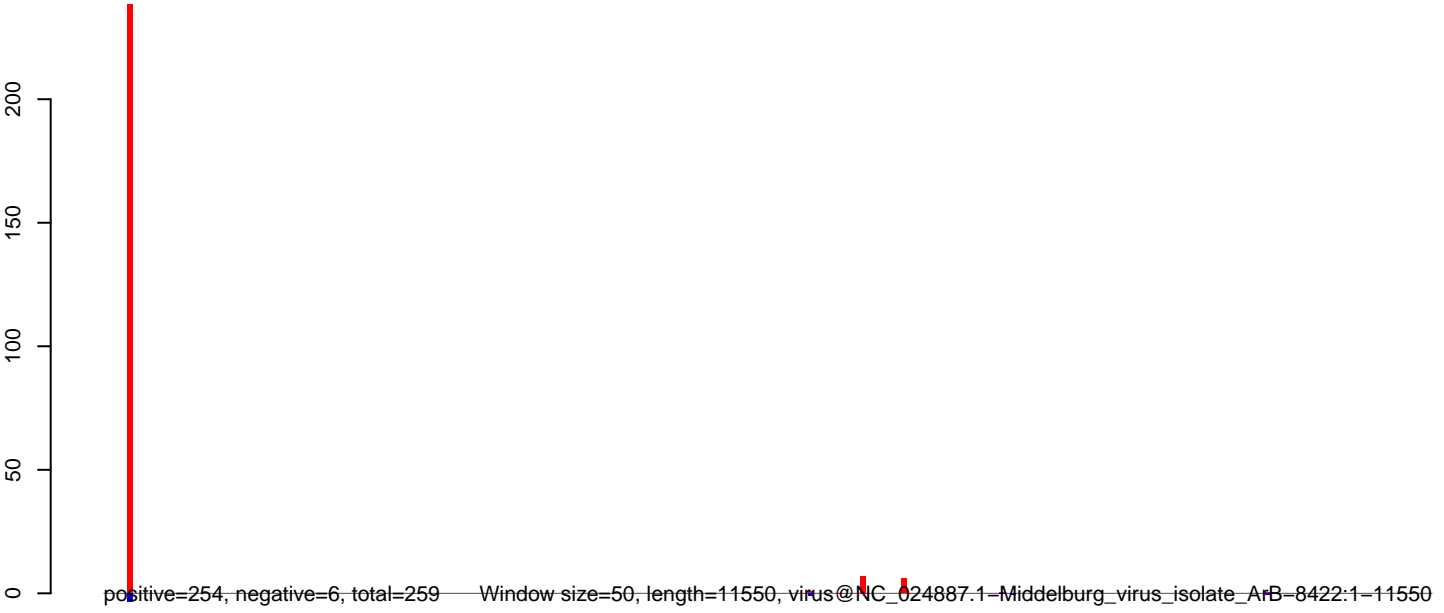
AeAeg_Aag2cells_CHIKV.18_23.rep



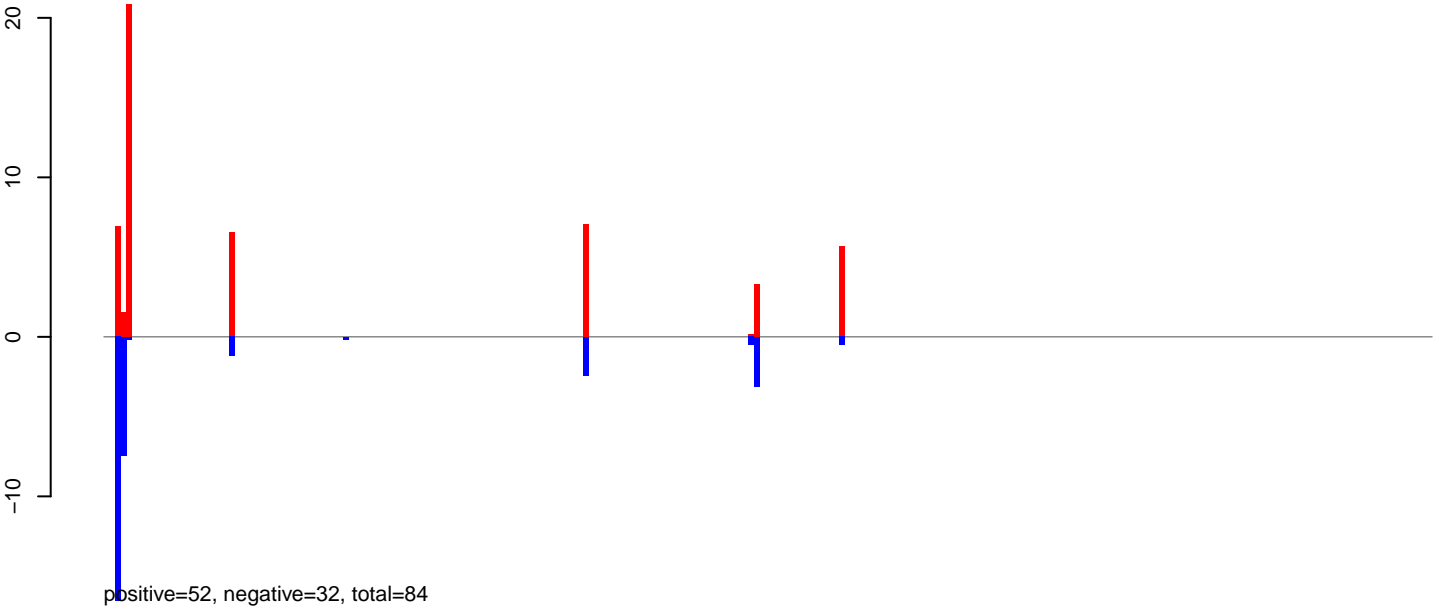
AeAeg_Aag2cells_CHIKV.24_35.rep



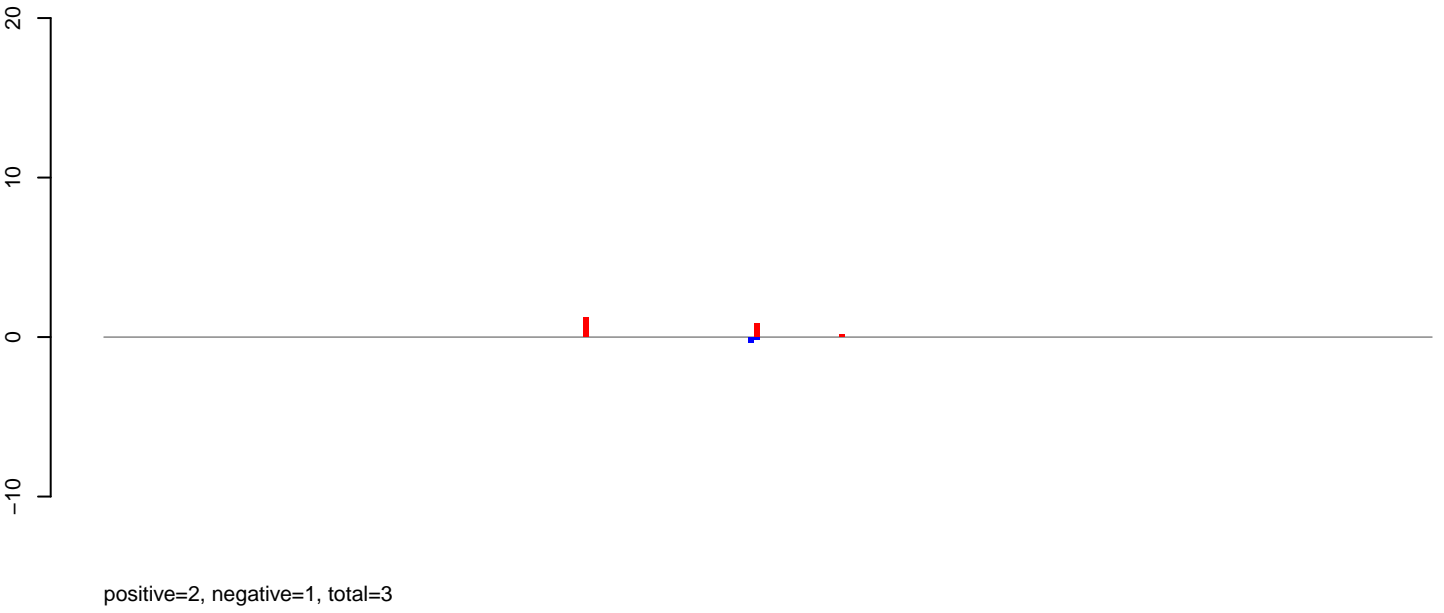
AeAeg_Aag2cells_CHIKV.rep



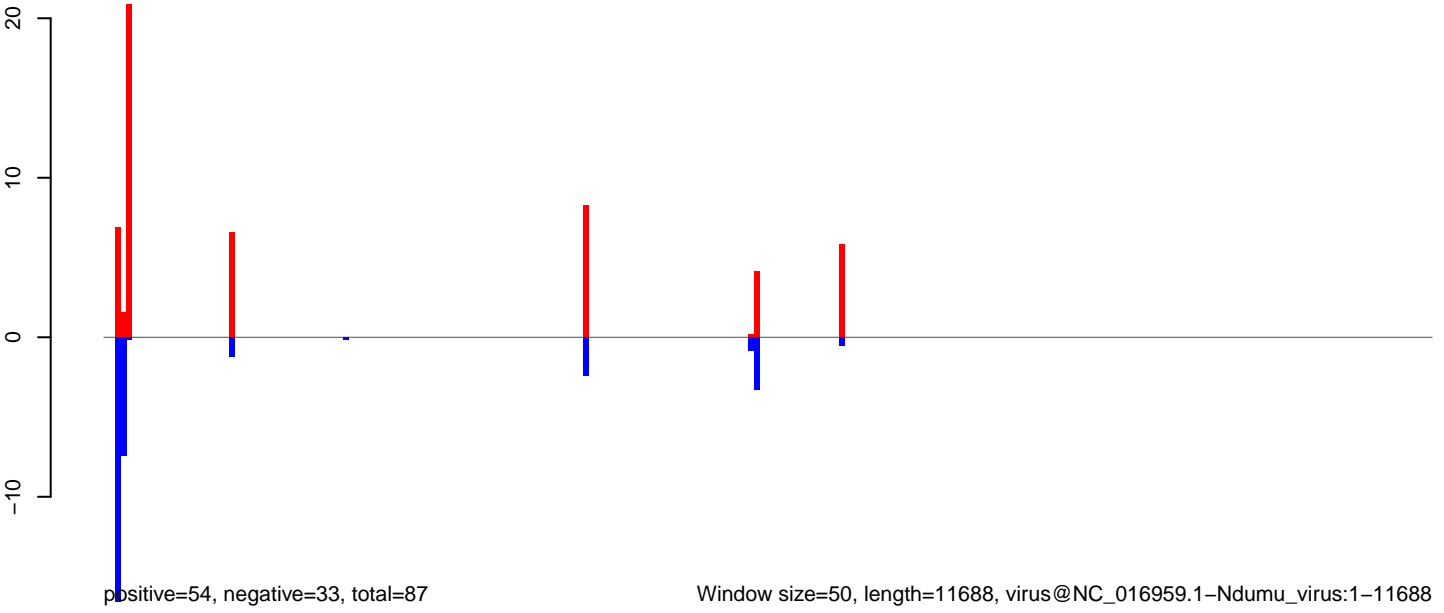
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep



Window size=50, length=11688, virus@NC_016959.1-Ndumu_virus:1-11688

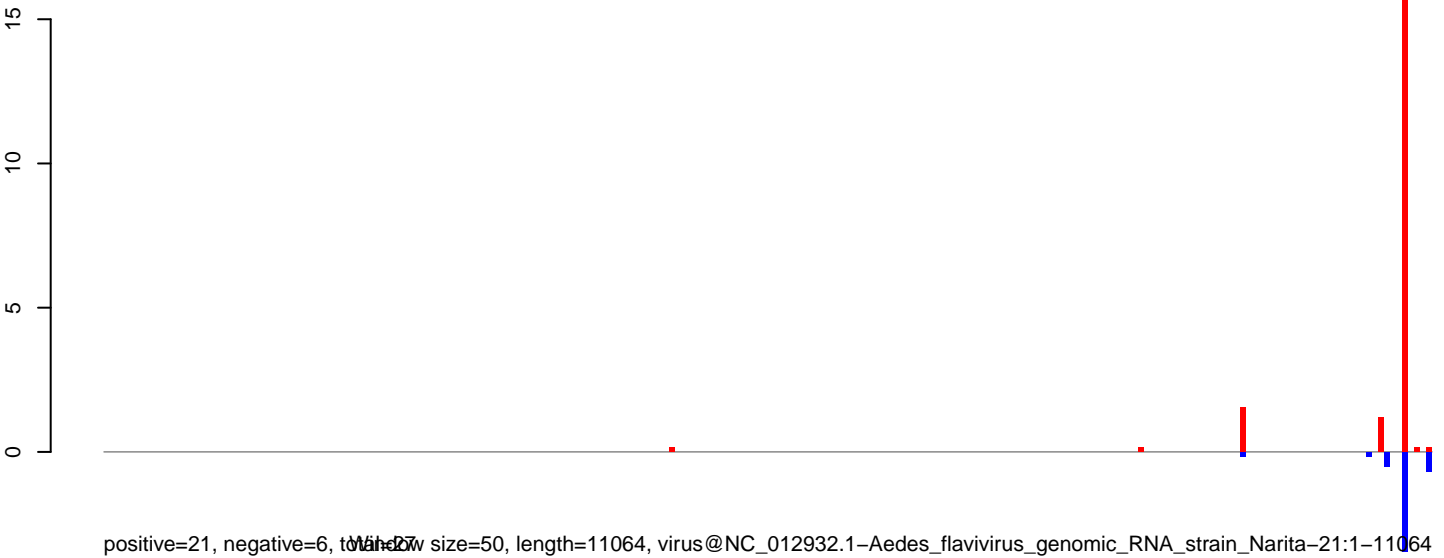
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep



Window size=50, length=11064, virus@NC_012932.1-Aedes_flavivirus_genomic_RNA_strain_Narita-21:1-11064

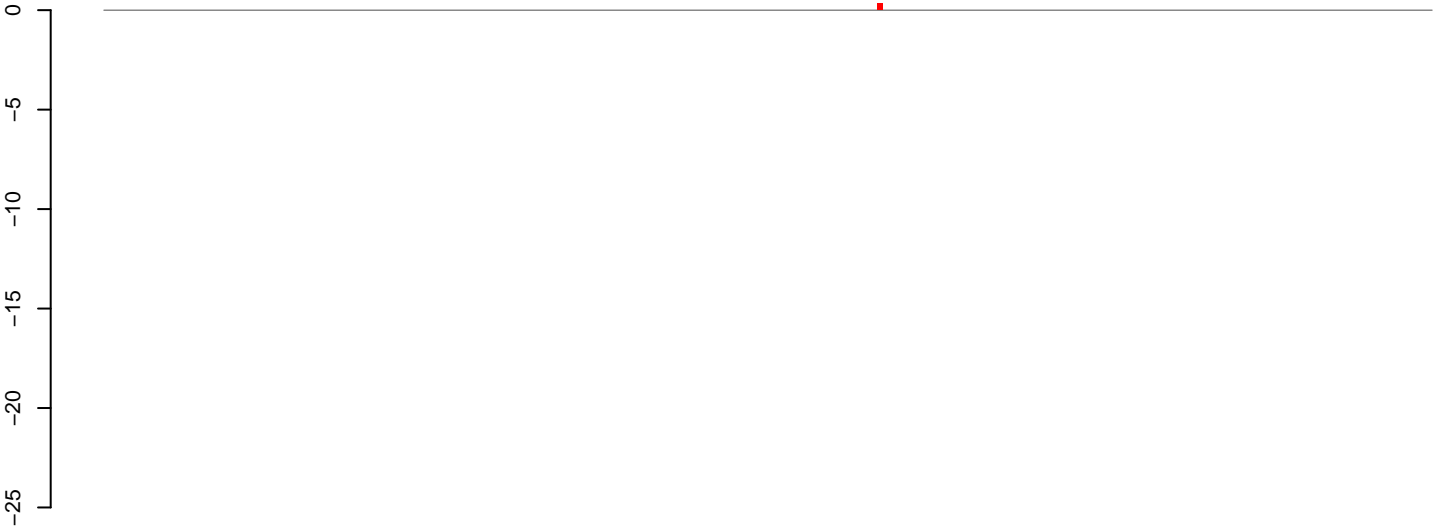
0 2000 4000 6000 8000 10000

AeAeg_Aag2cells_CHIKV.18_23.rep



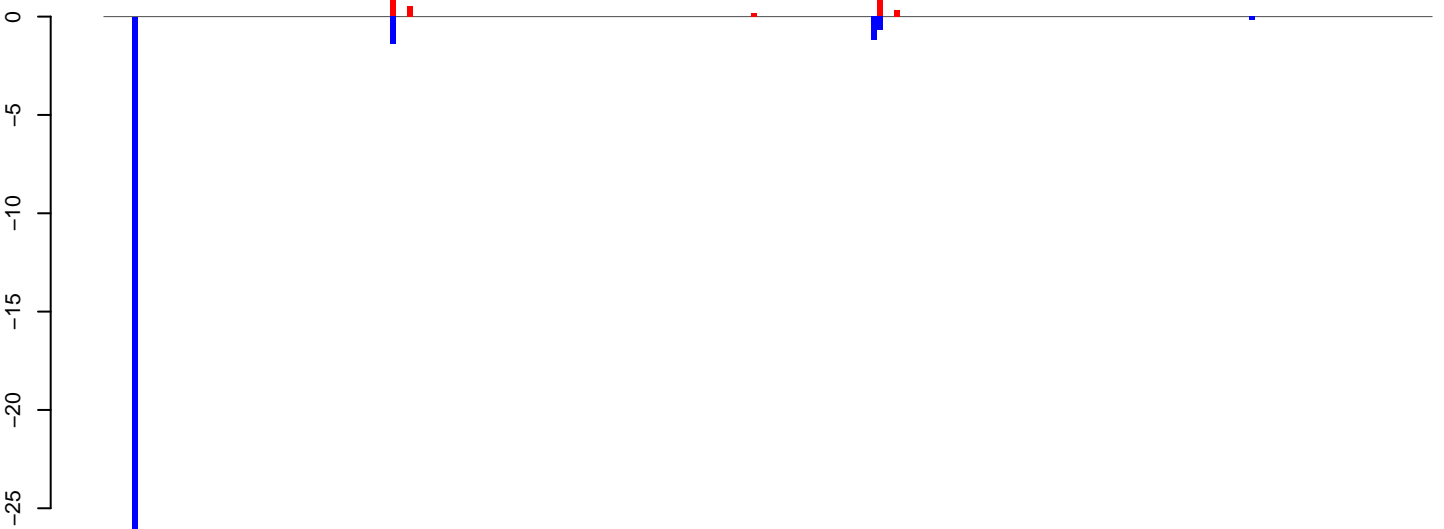
positive=6, negative=31, total=37

AeAeg_Aag2cells_CHIKV.24_35.rep



positive=0, negative=0, total=0

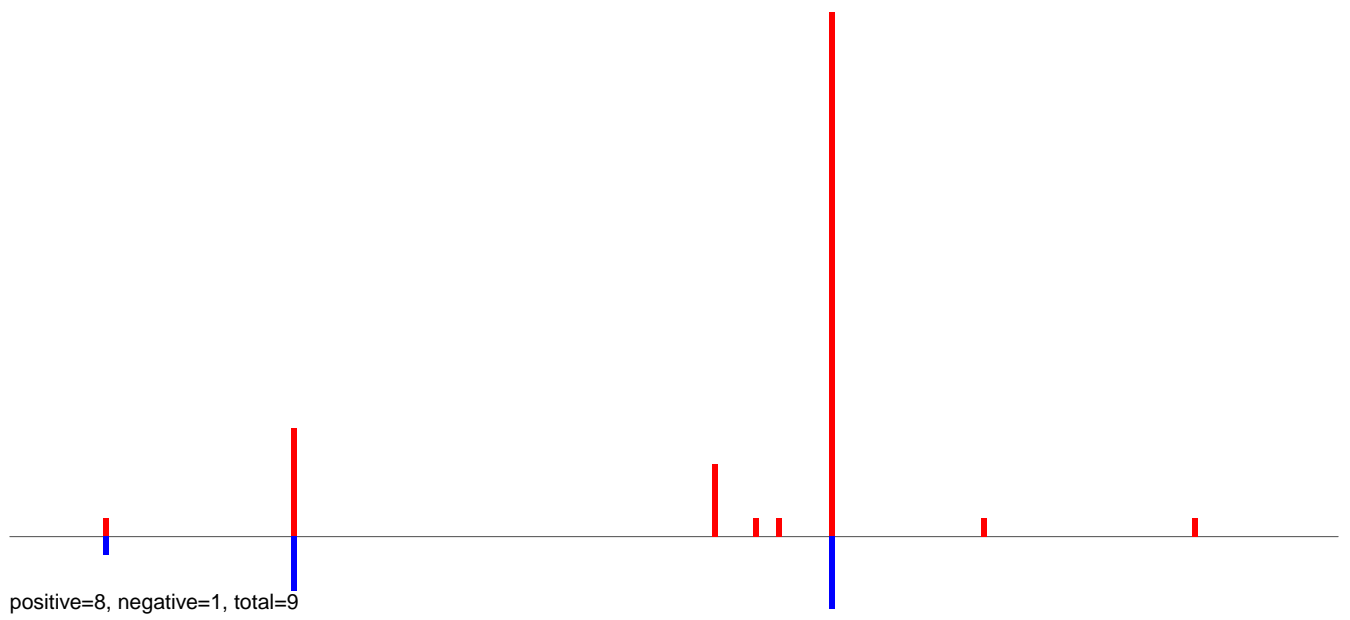
AeAeg_Aag2cells_CHIKV.rep



positive=6, negative=31, total=37

Window size=50, length=11634, virus@NC_018615.1-Eilat_virus_isolate_EO329:1-11634

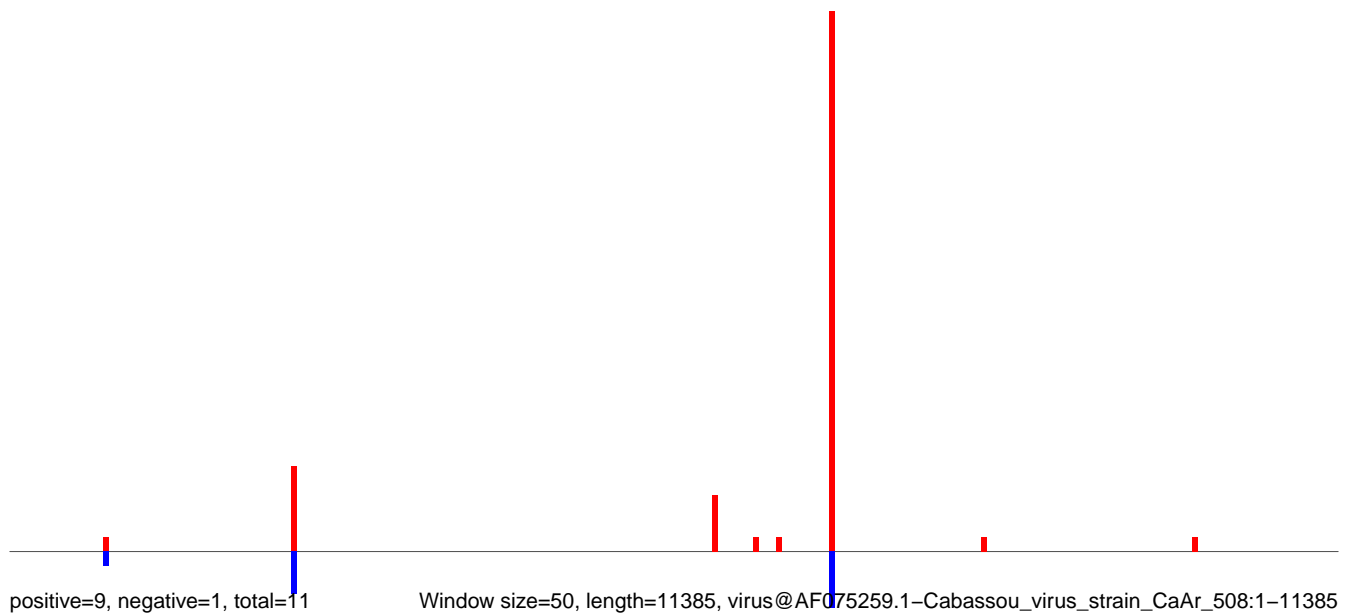
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



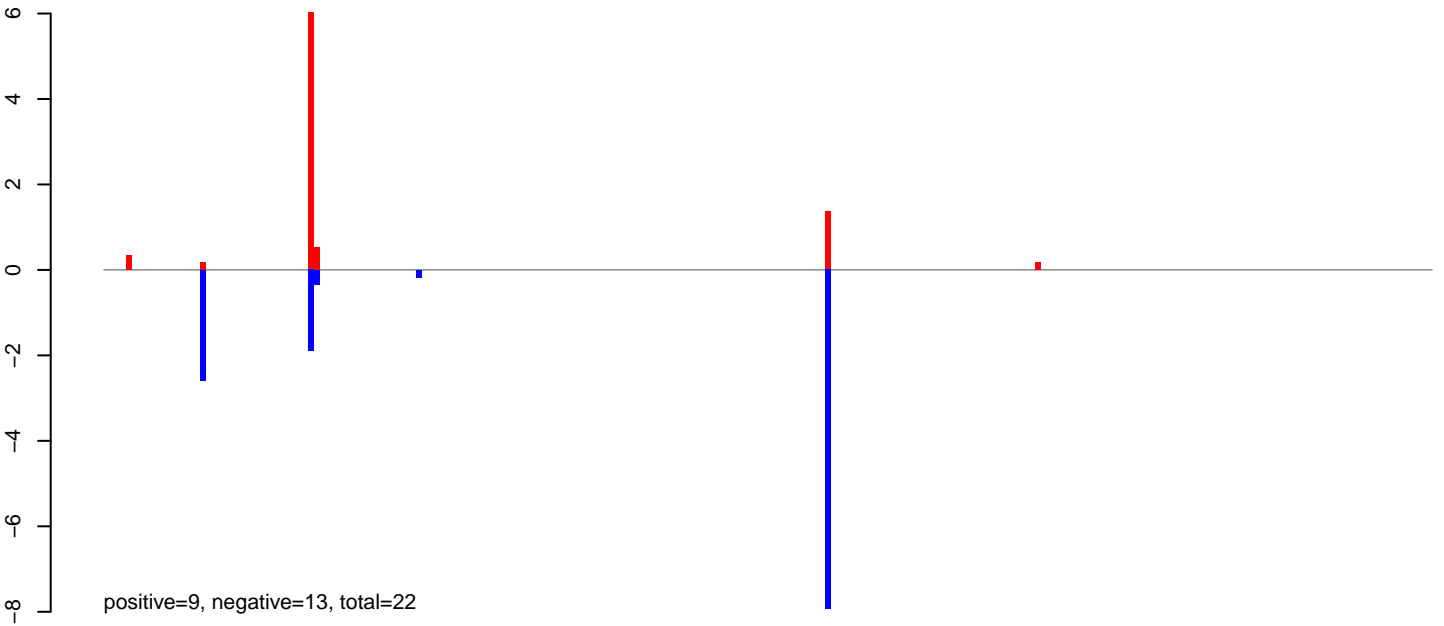
AeAeg_Aag2cells_CHIKV.rep



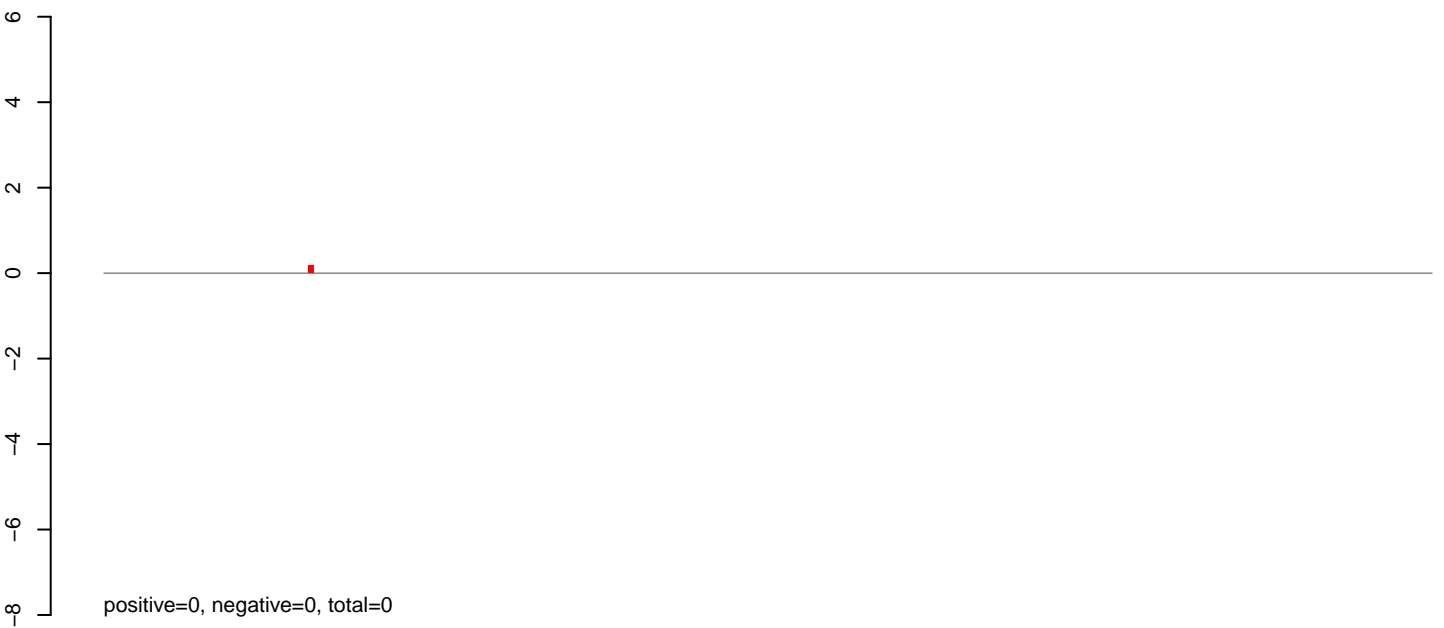
Window size=50, length=11385, virus@AF075259.1-Cabassou_virus_strain_CaAr_508:1-11385

0 2000 4000 6000 8000 10000

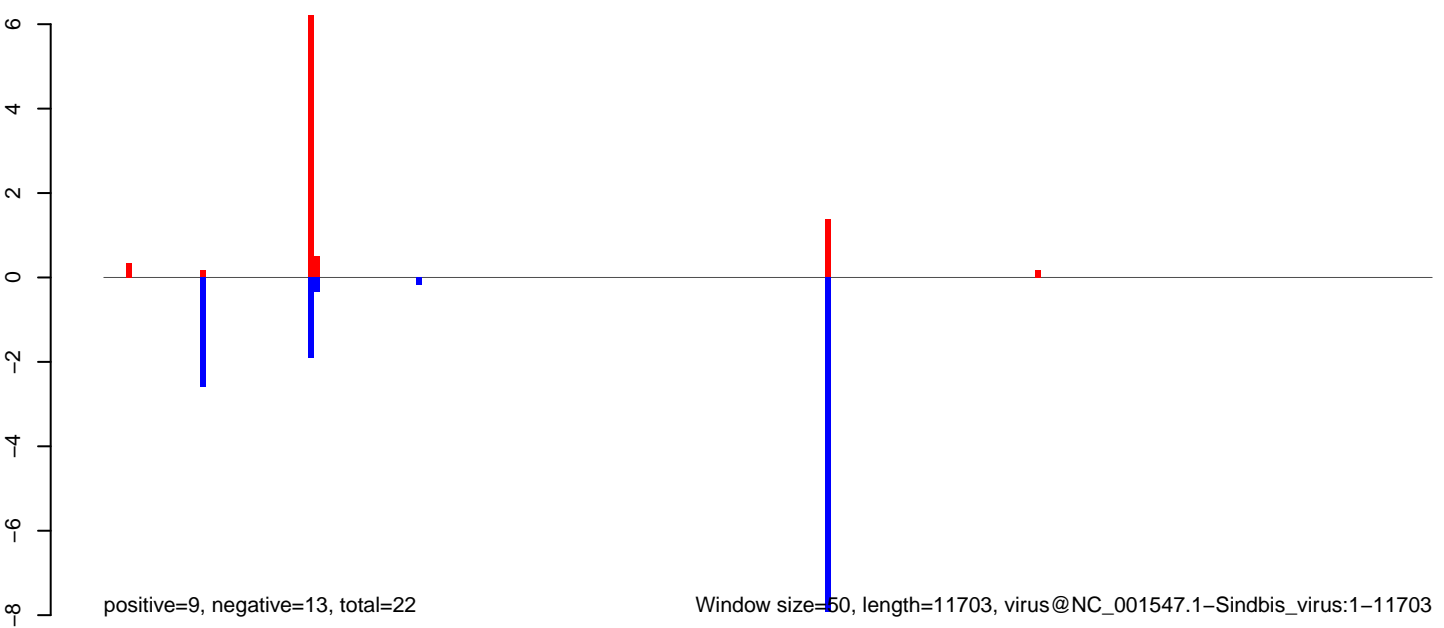
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep



0 2000 4000 6000 8000 10000 12000

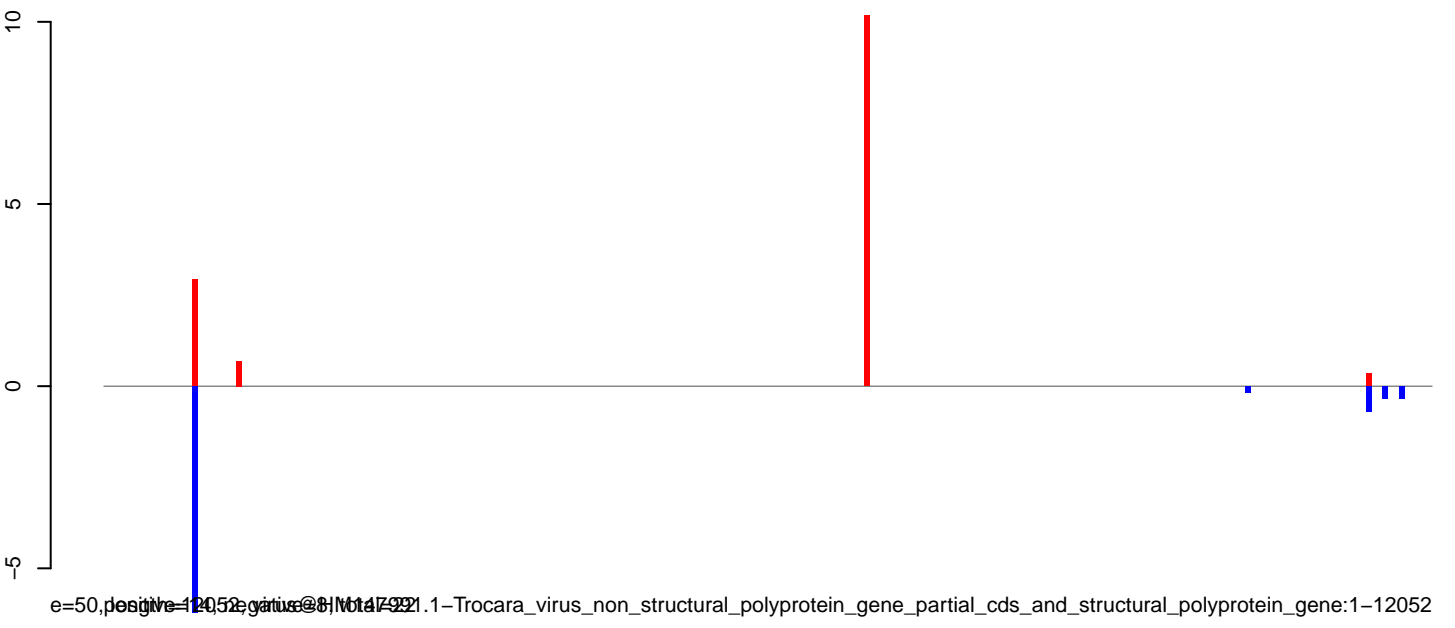
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep



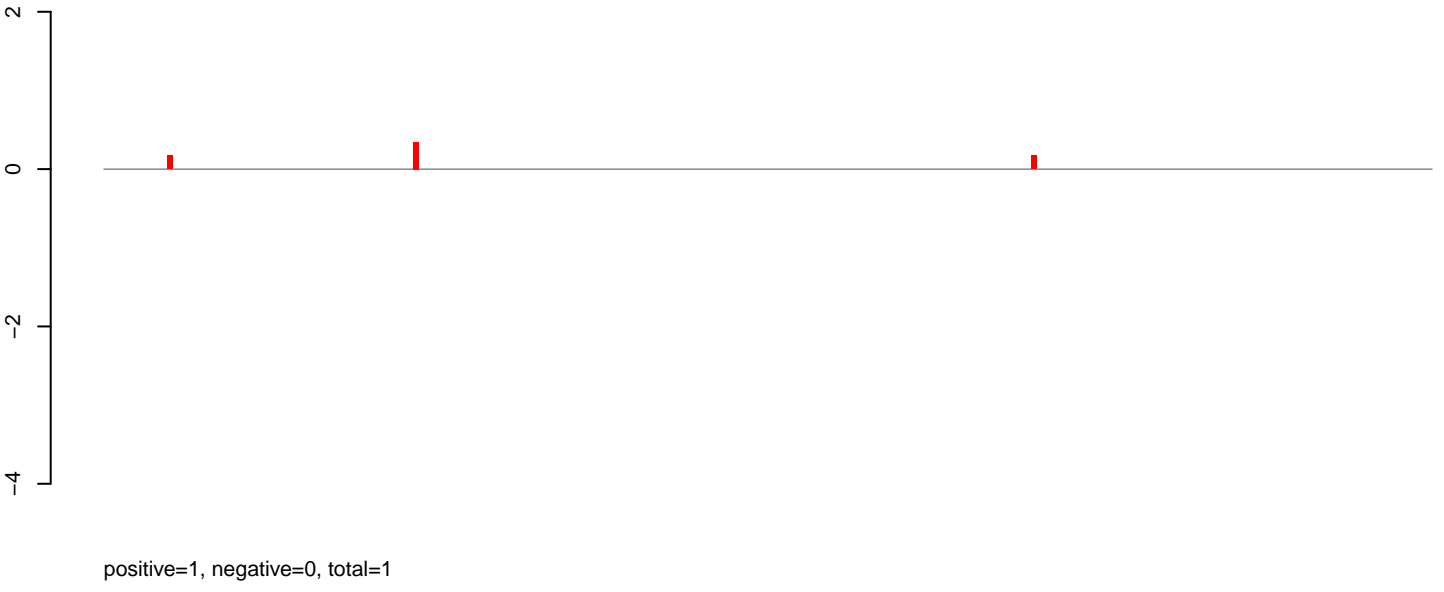
e=50, positive=12052, negative=8, total=12060
1-Trocaro_virus_non_structural_polyprotein_gene_partial_cds_and_structural_polyprotein_gene:1-12052

0 2000 4000 6000 8000 10000 12000

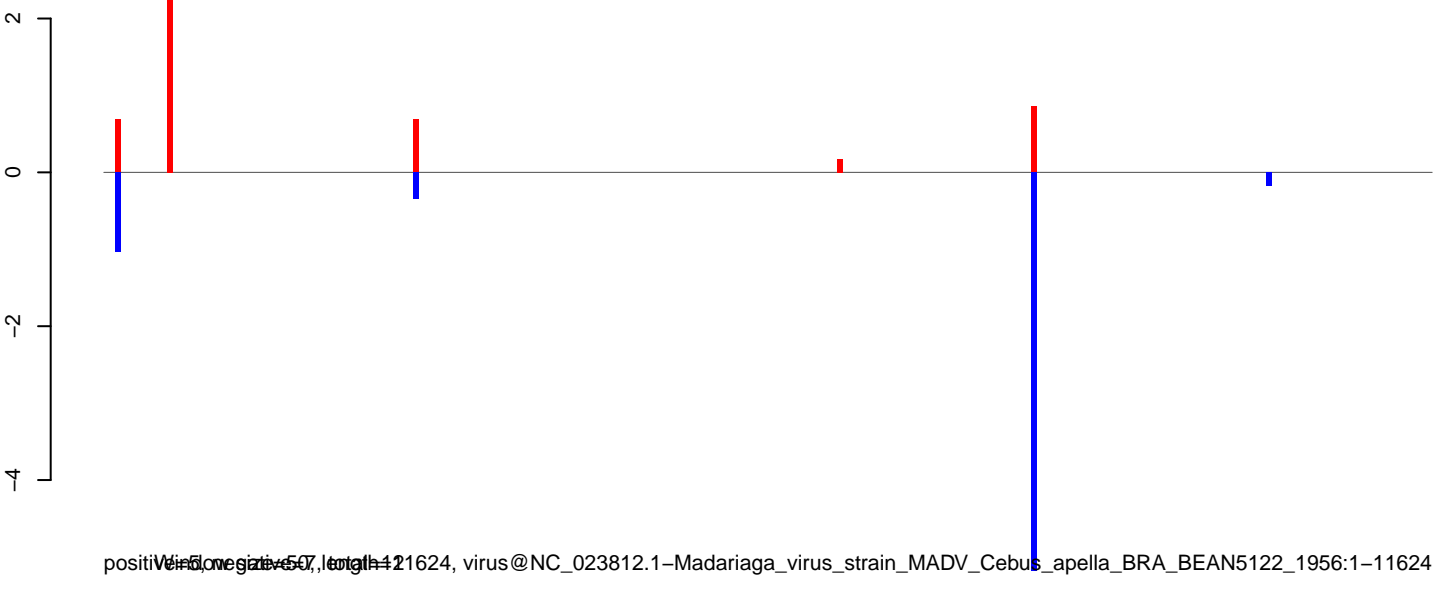
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

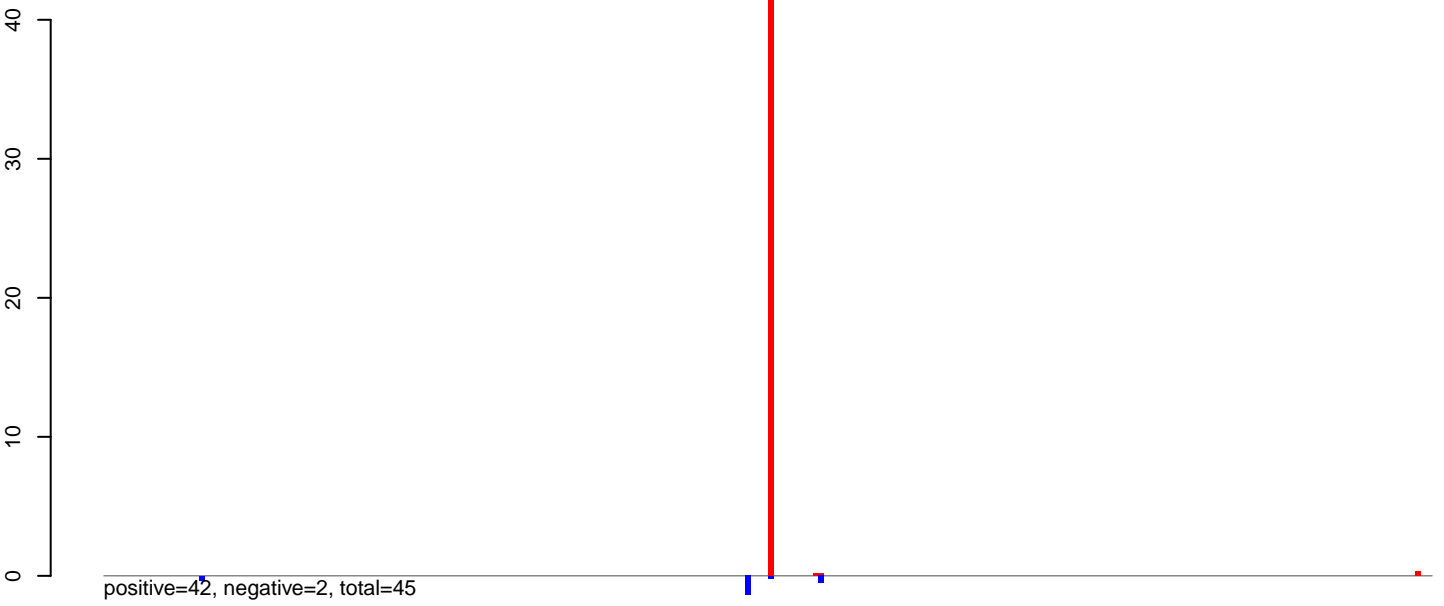


AeAeg_Aag2cells_CHIKV.rep

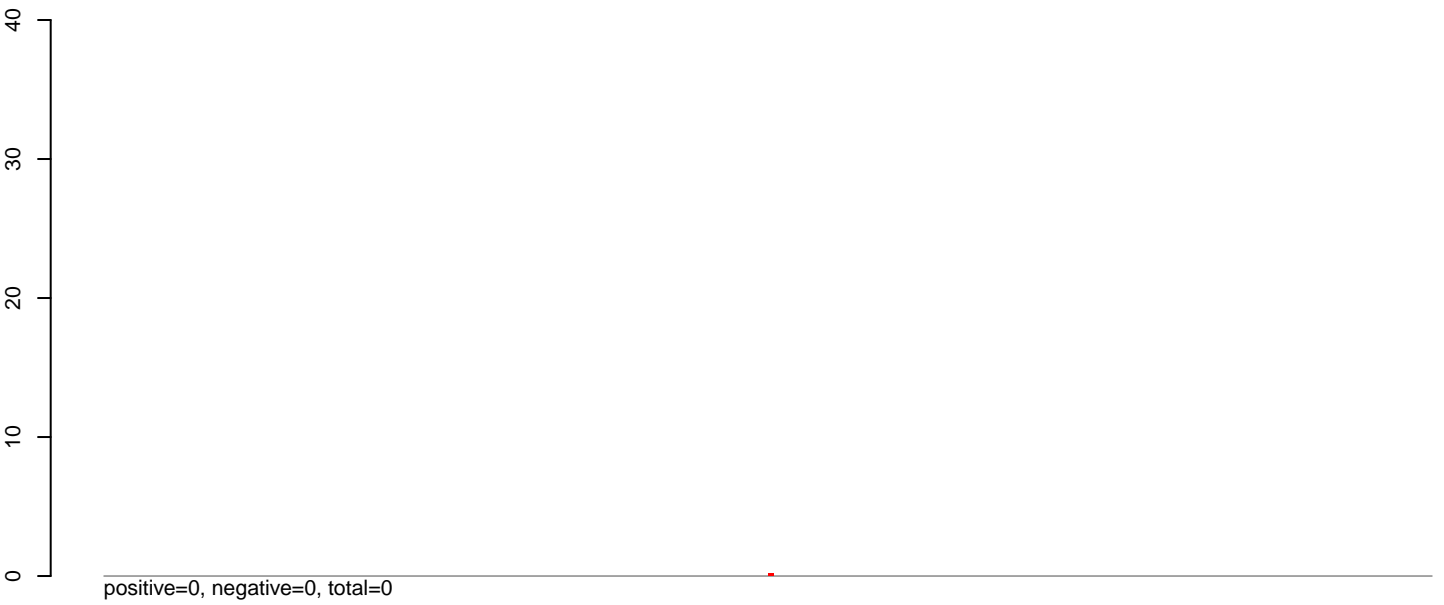


View on genome browser: [https://www.ncbi.nlm.nih.gov/genomes/Genomes.org/Home?CID=NC_023812.1-Madariaga_virus_strain_MADV_Cebus_apella_BRA_BEAN5122_1956:1-11624](#)

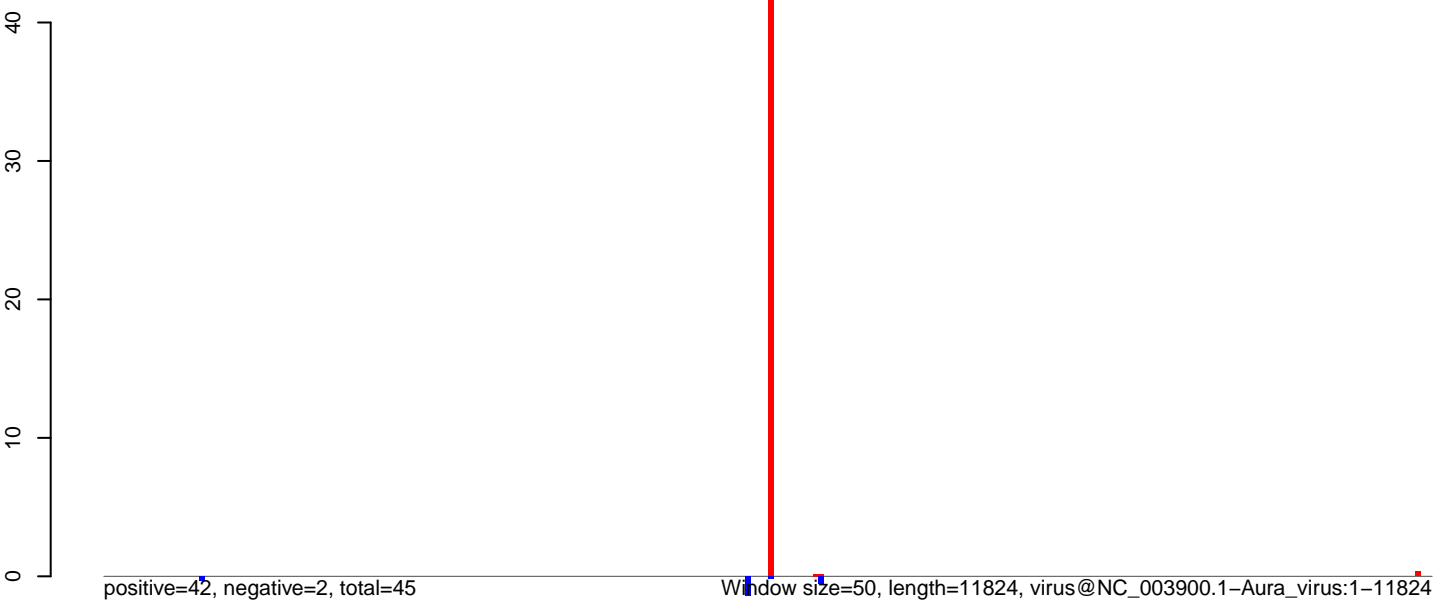
AeAeg_Aag2cells_CHIKV.18_23.rep



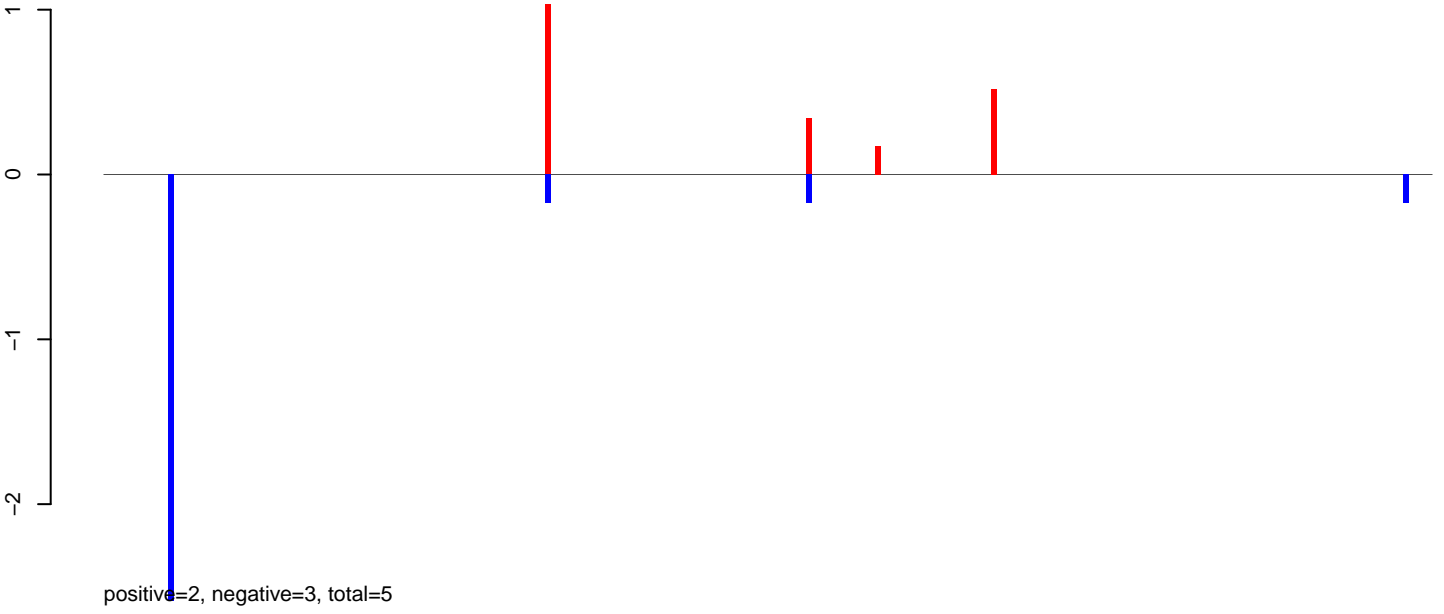
AeAeg_Aag2cells_CHIKV.24_35.rep



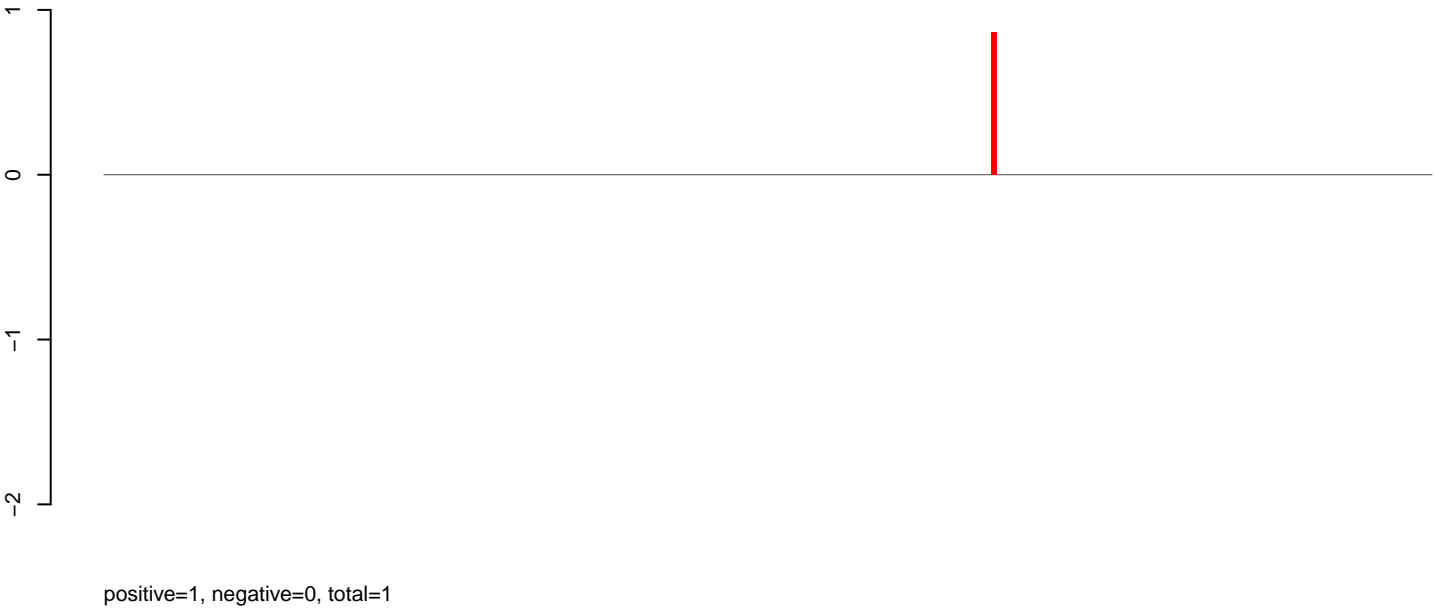
AeAeg_Aag2cells_CHIKV.rep



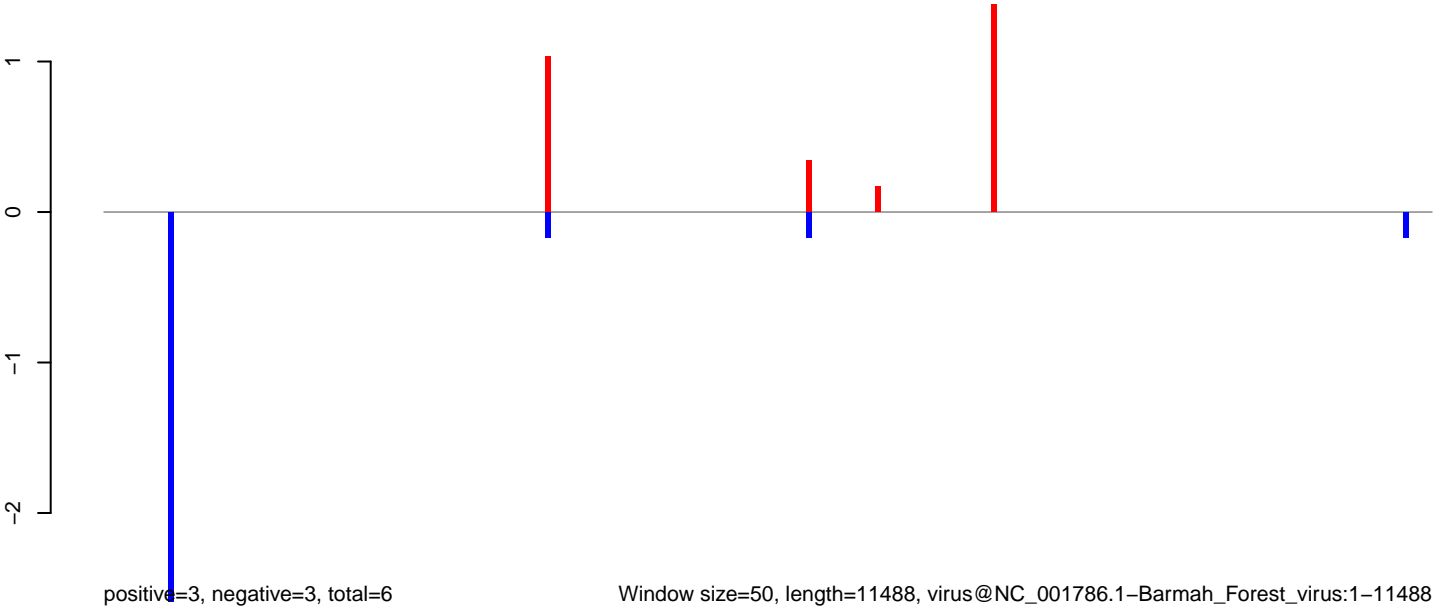
AeAeg_Aag2cells_CHIKV.18_23.rep



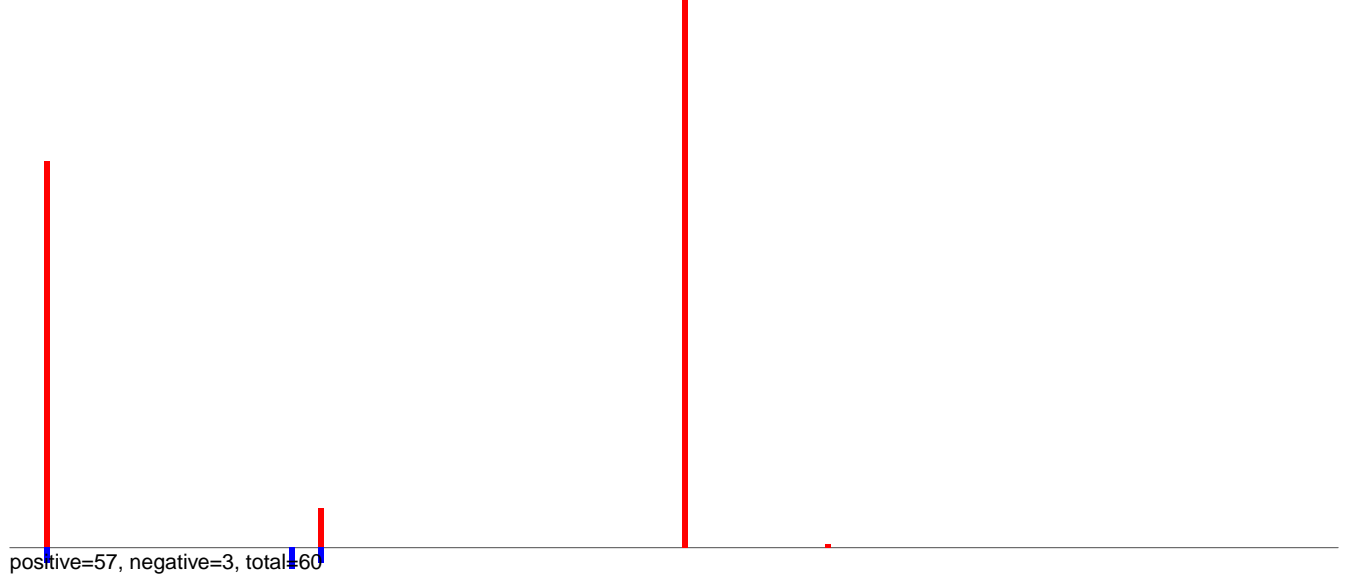
AeAeg_Aag2cells_CHIKV.24_35.rep



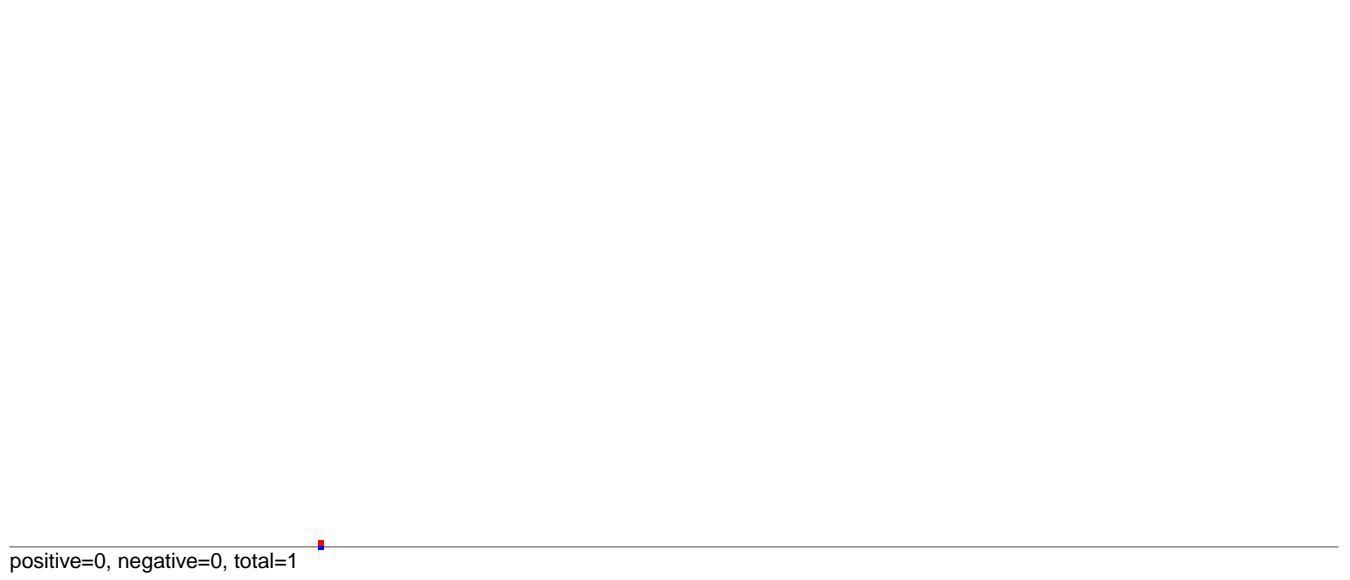
AeAeg_Aag2cells_CHIKV.rep



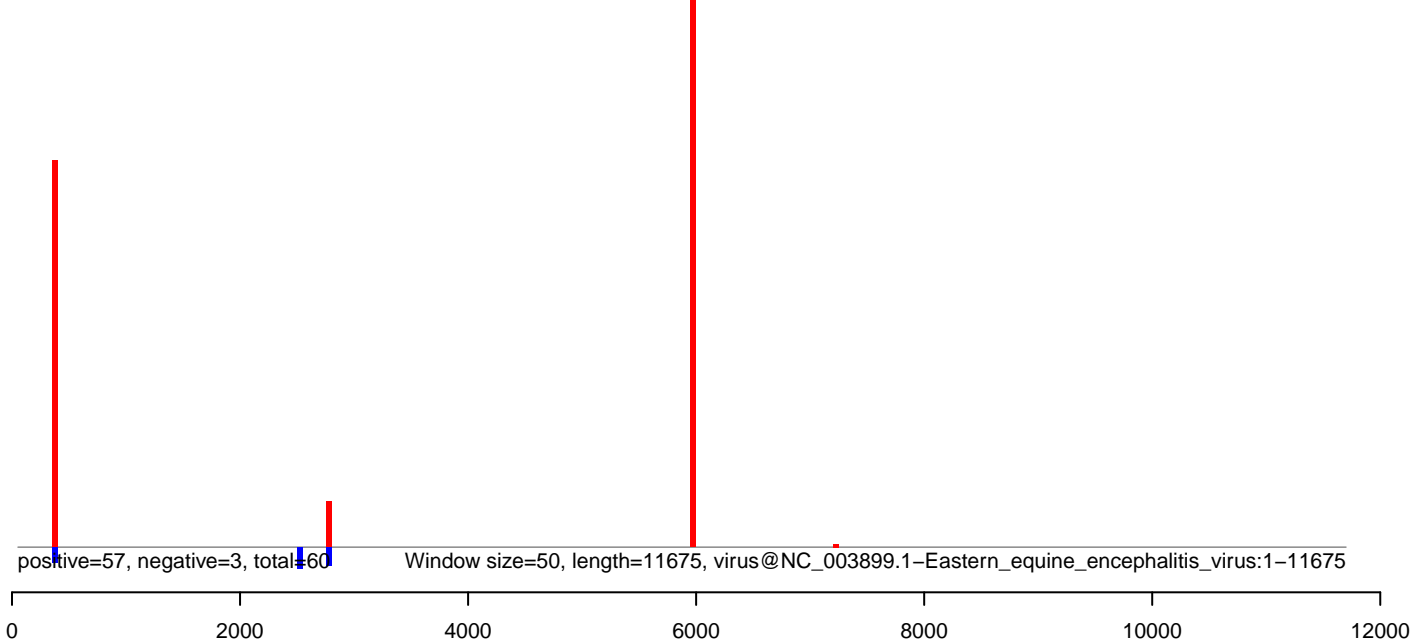
AeAeg_Aag2cells_CHIKV.18_23.rep



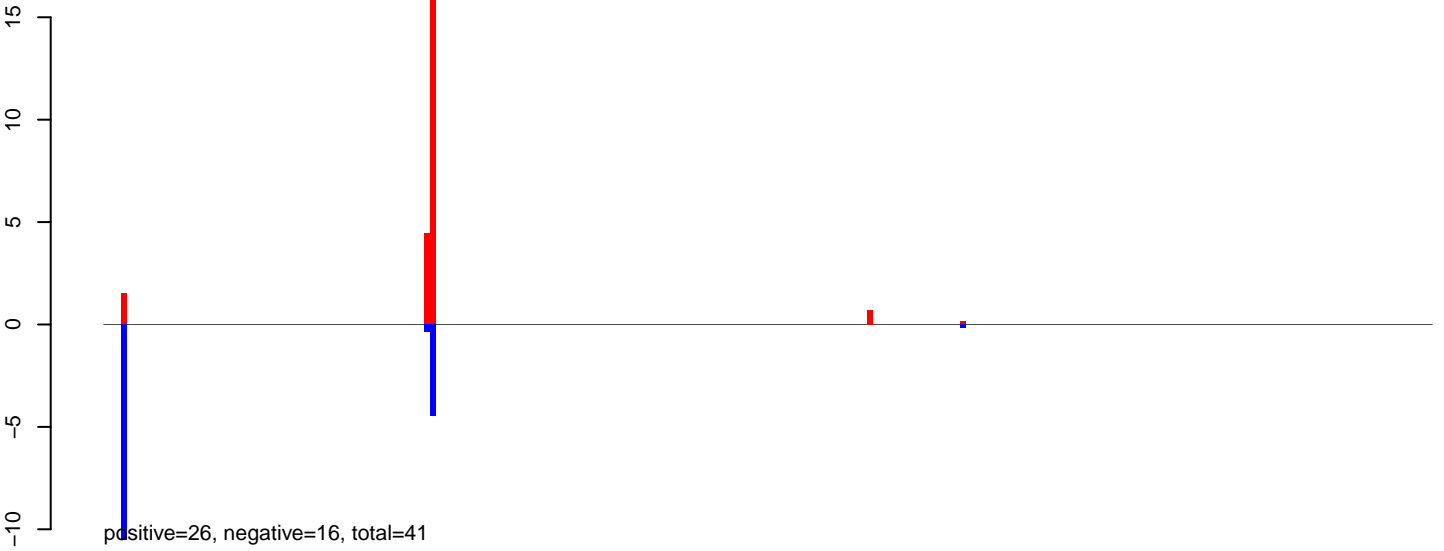
AeAeg_Aag2cells_CHIKV.24_35.rep



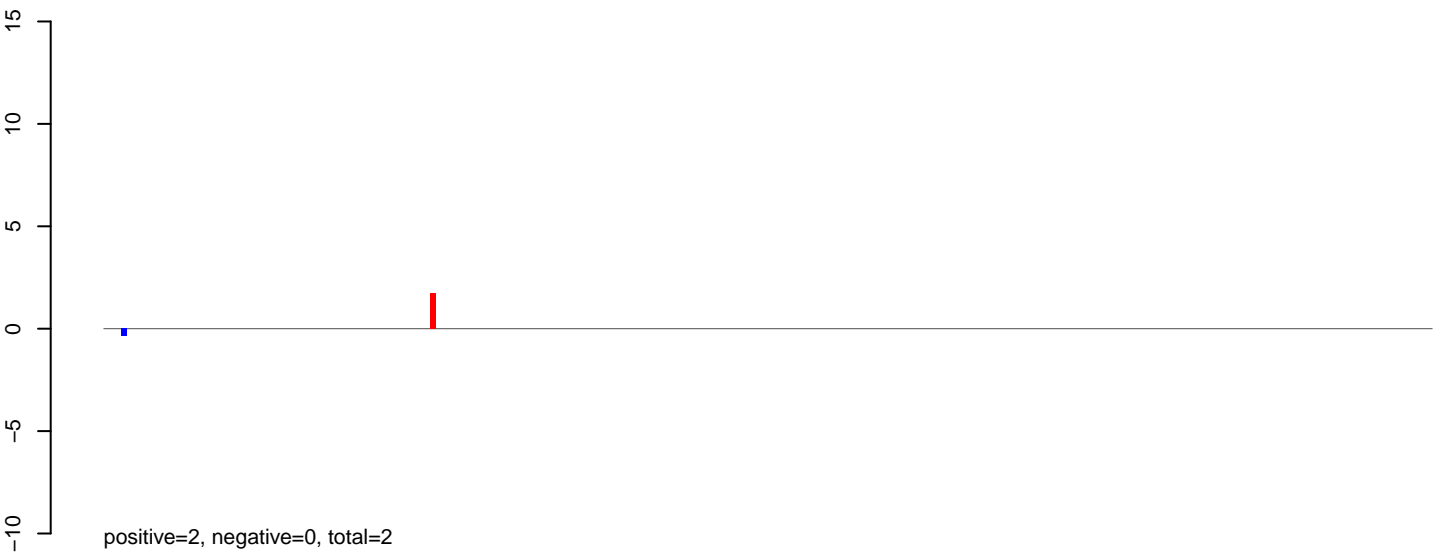
AeAeg_Aag2cells_CHIKV.rep



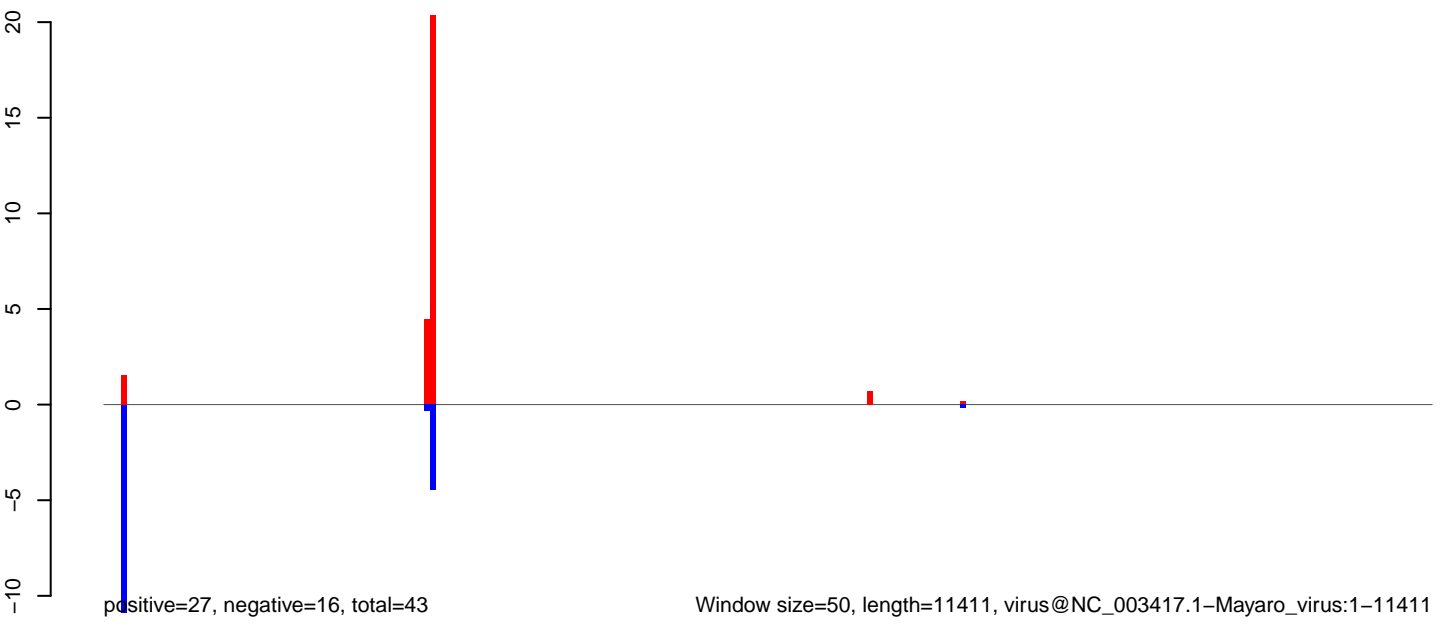
AeAeg_Aag2cells_CHIKV.18_23.rep



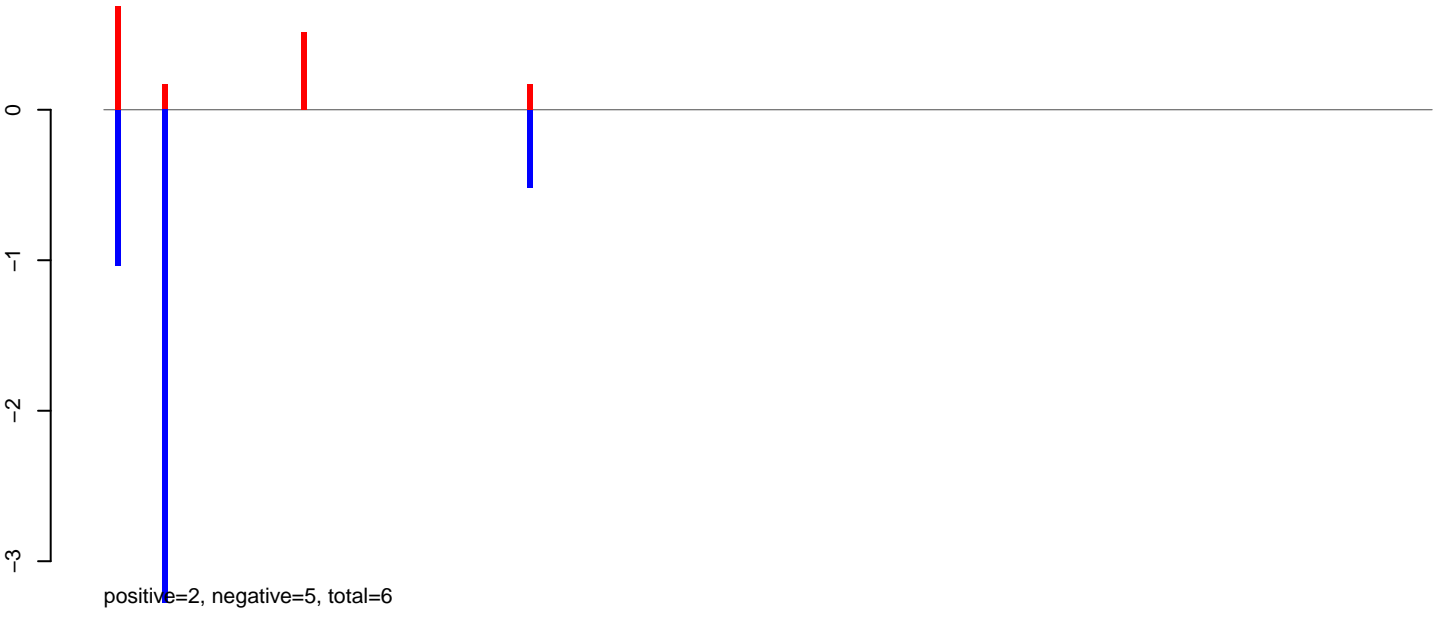
AeAeg_Aag2cells_CHIKV.24_35.rep



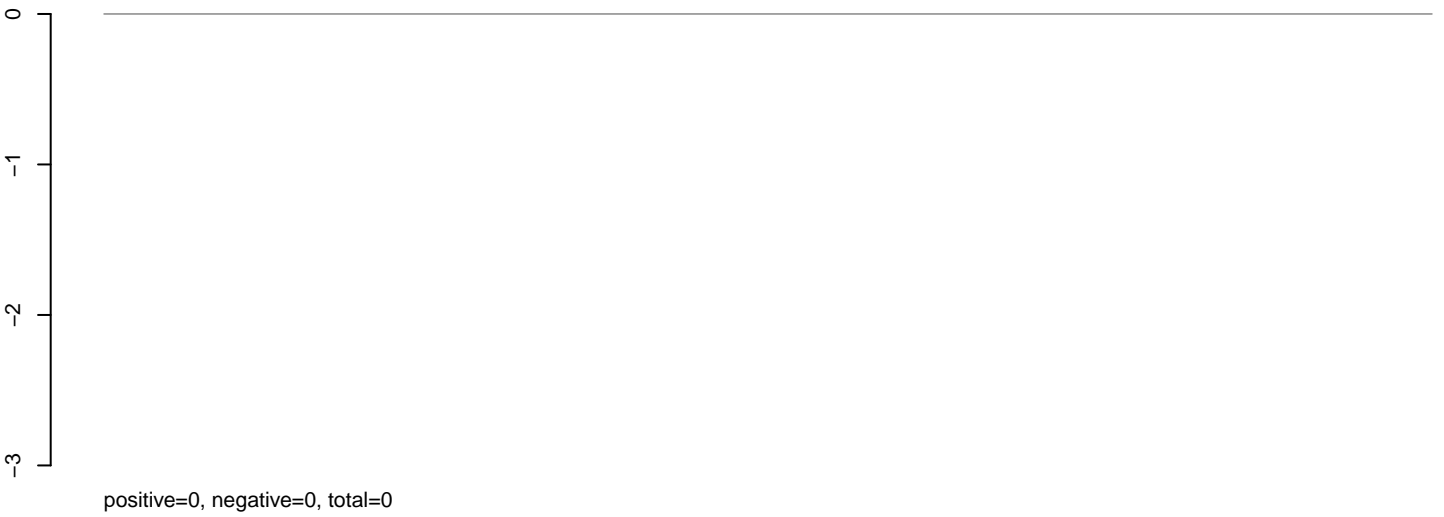
AeAeg_Aag2cells_CHIKV.rep



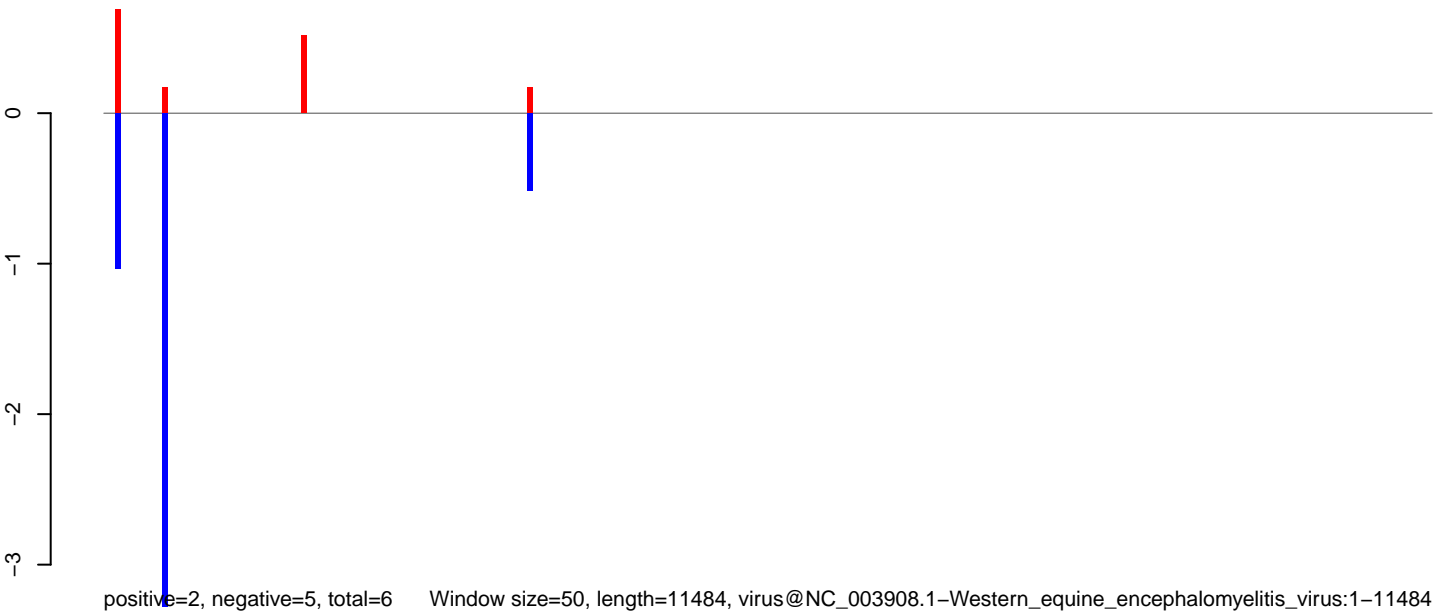
AeAeg_Aag2cells_CHIKV.18_23.rep



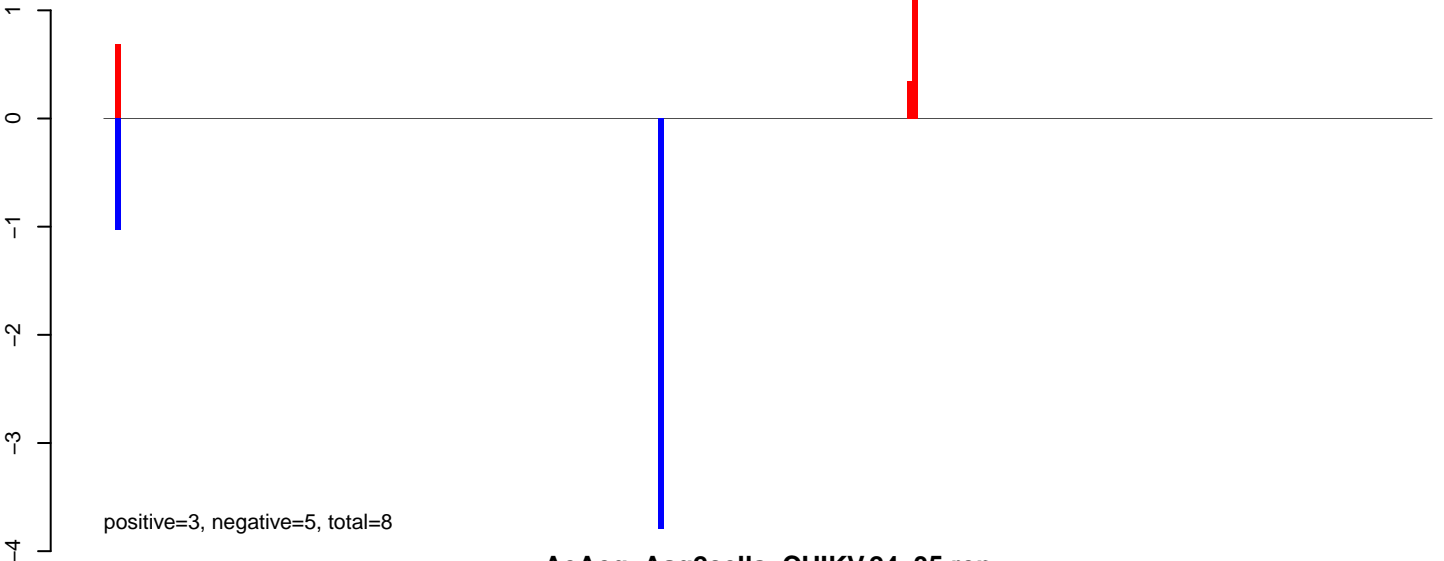
AeAeg_Aag2cells_CHIKV.24_35.rep



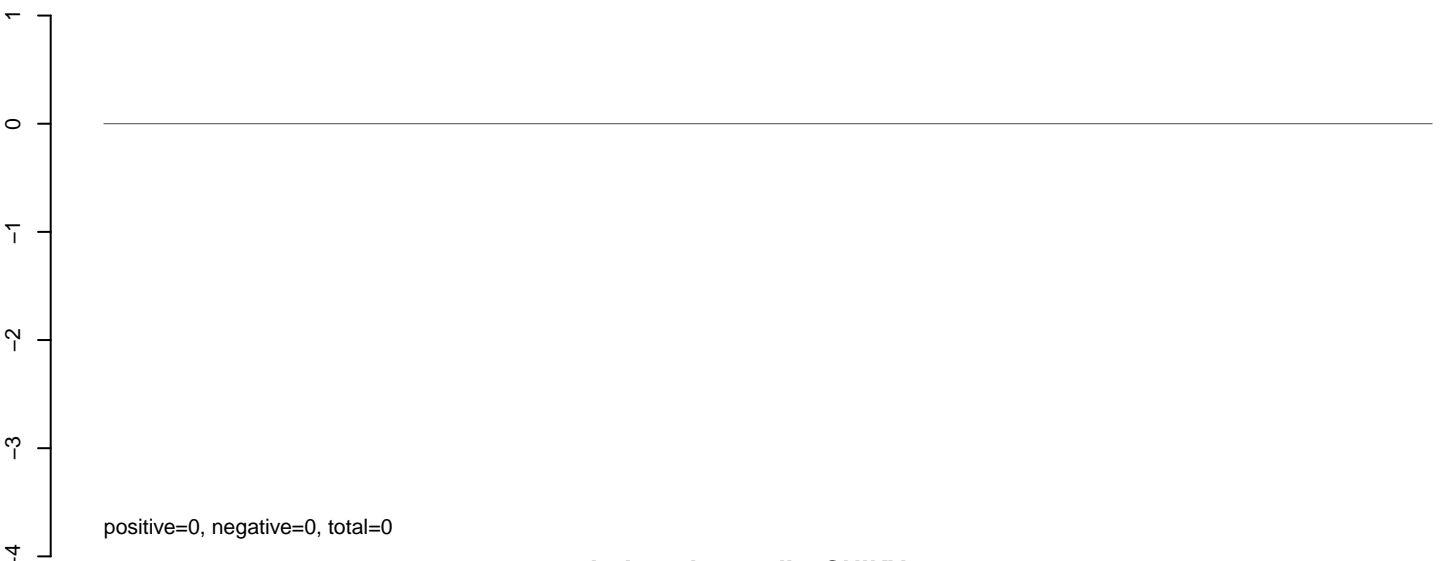
AeAeg_Aag2cells_CHIKV.rep



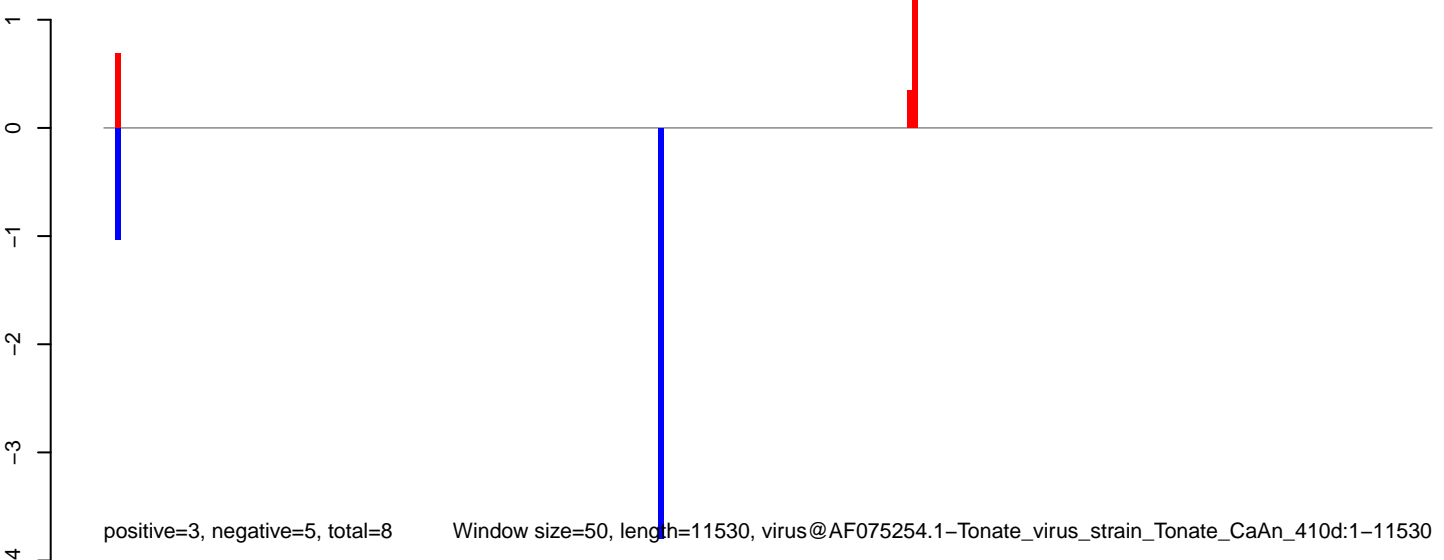
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

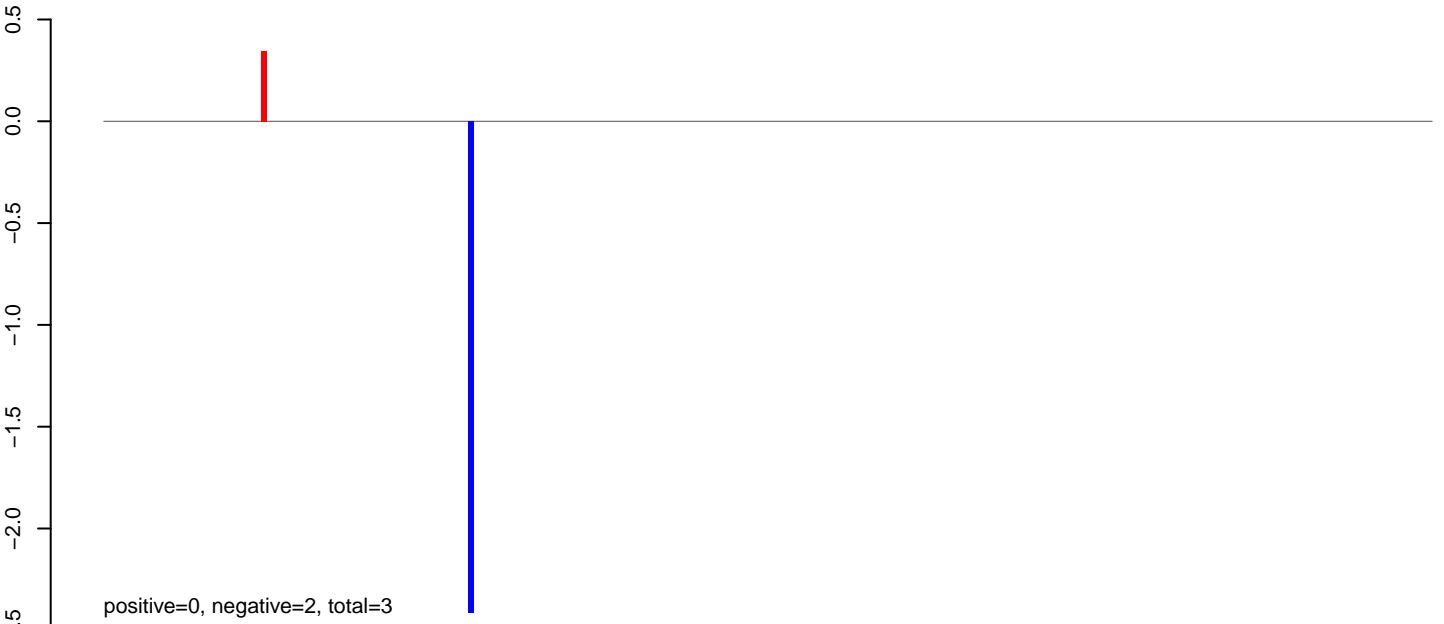


AeAeg_Aag2cells_CHIKV.rep

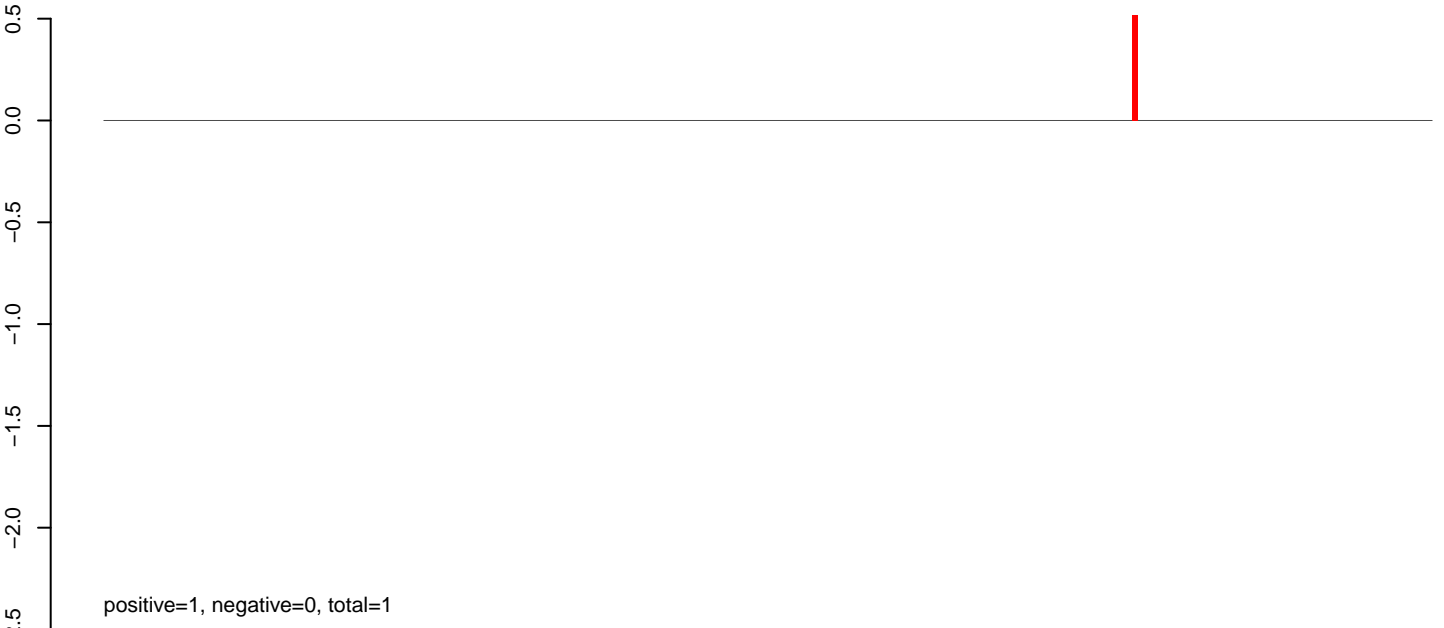


Window size=50, length=11530, virus@AF075254.1-Tonate_virus_strain_Tonate_CaAn_410d:1-11530

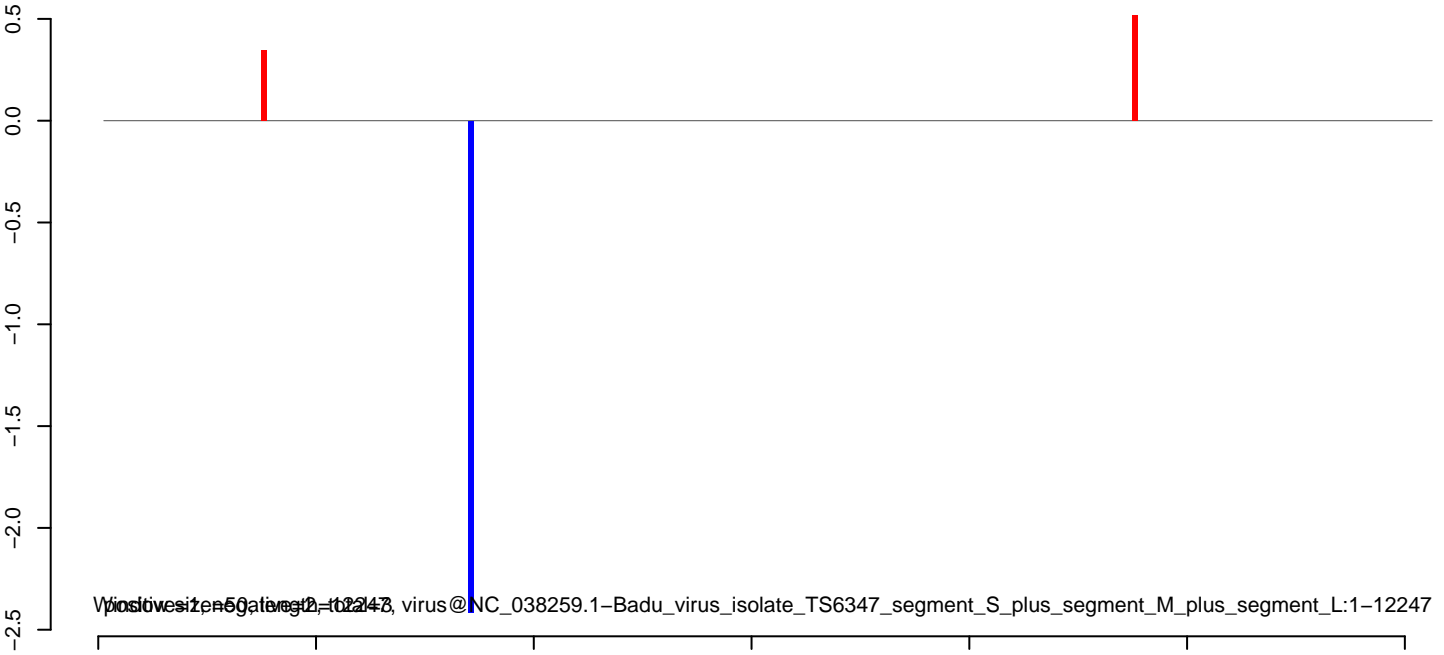
AeAeg_Aag2cells_CHIKV.18_23.rep



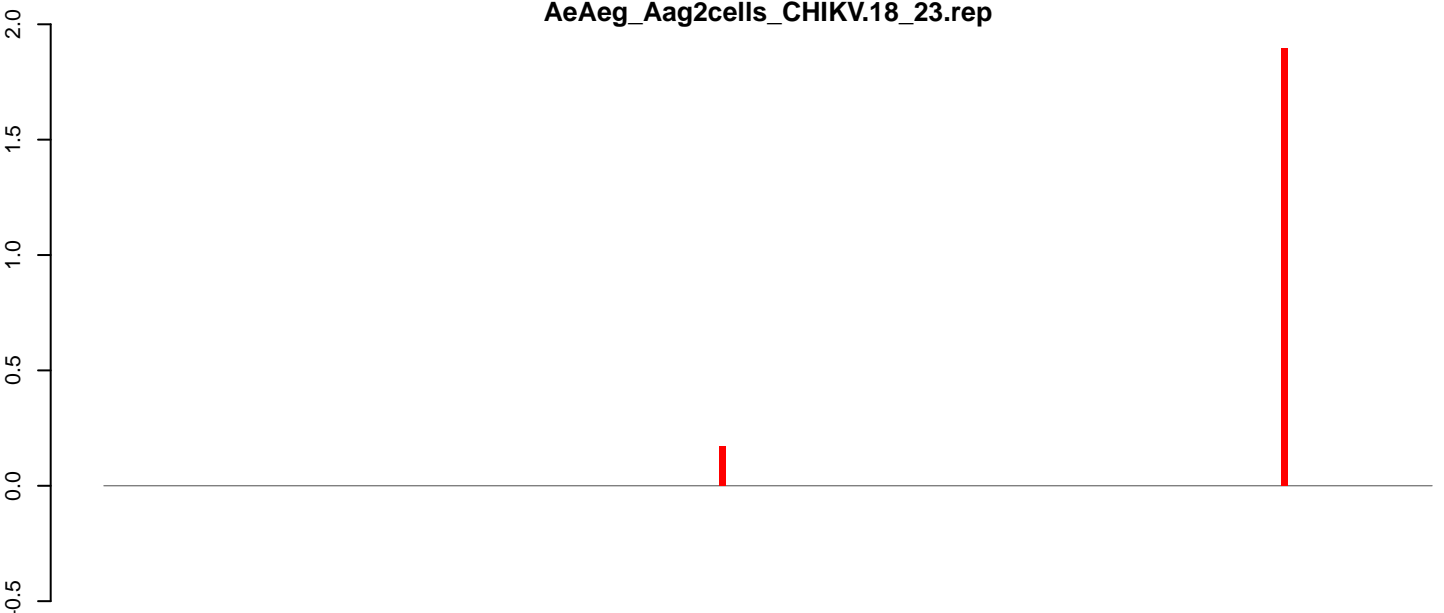
AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep

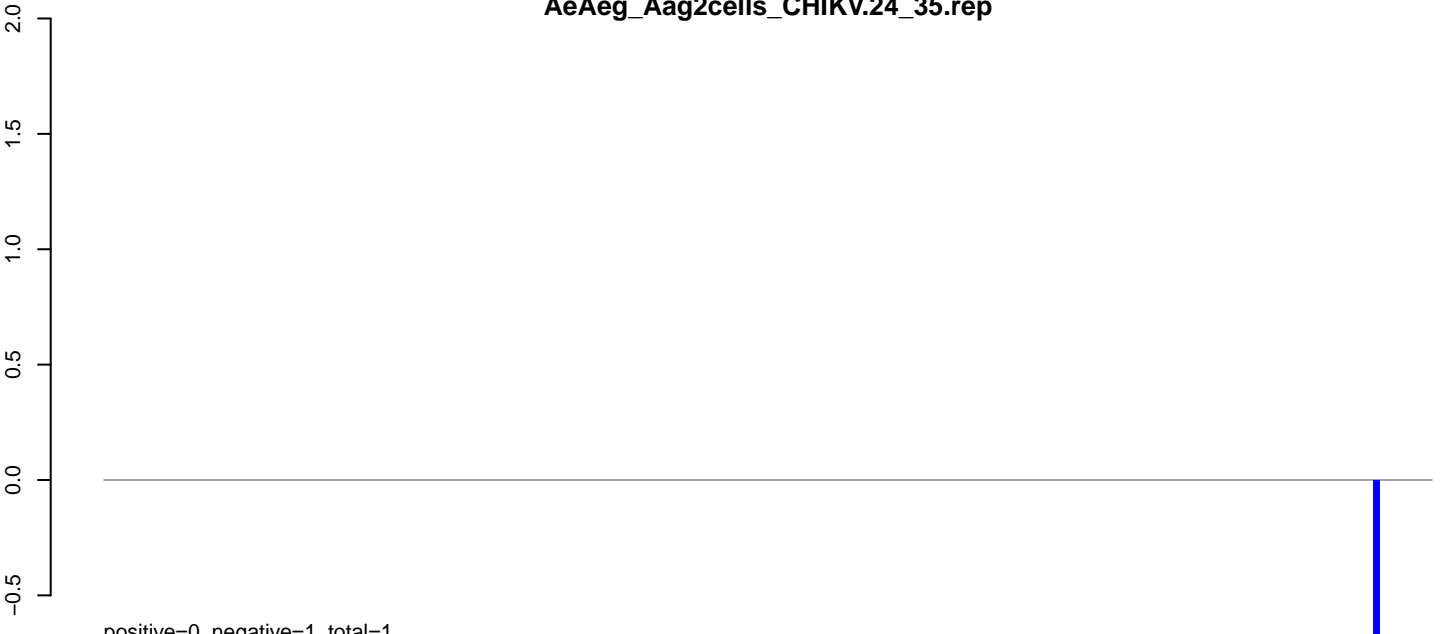


AeAeg_Aag2cells_CHIKV.18_23.rep



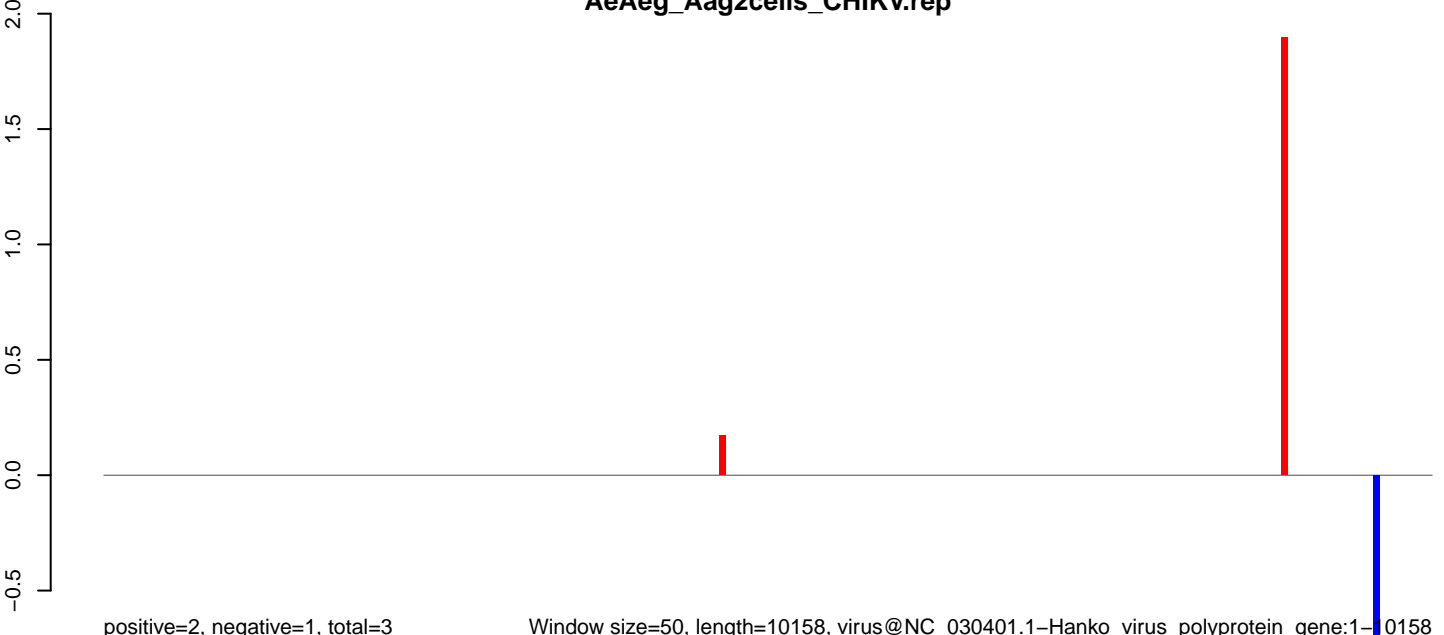
positive=2, negative=0, total=2

AeAeg_Aag2cells_CHIKV.24_35.rep



positive=0, negative=1, total=1

AeAeg_Aag2cells_CHIKV.rep



positive=2, negative=1, total=3

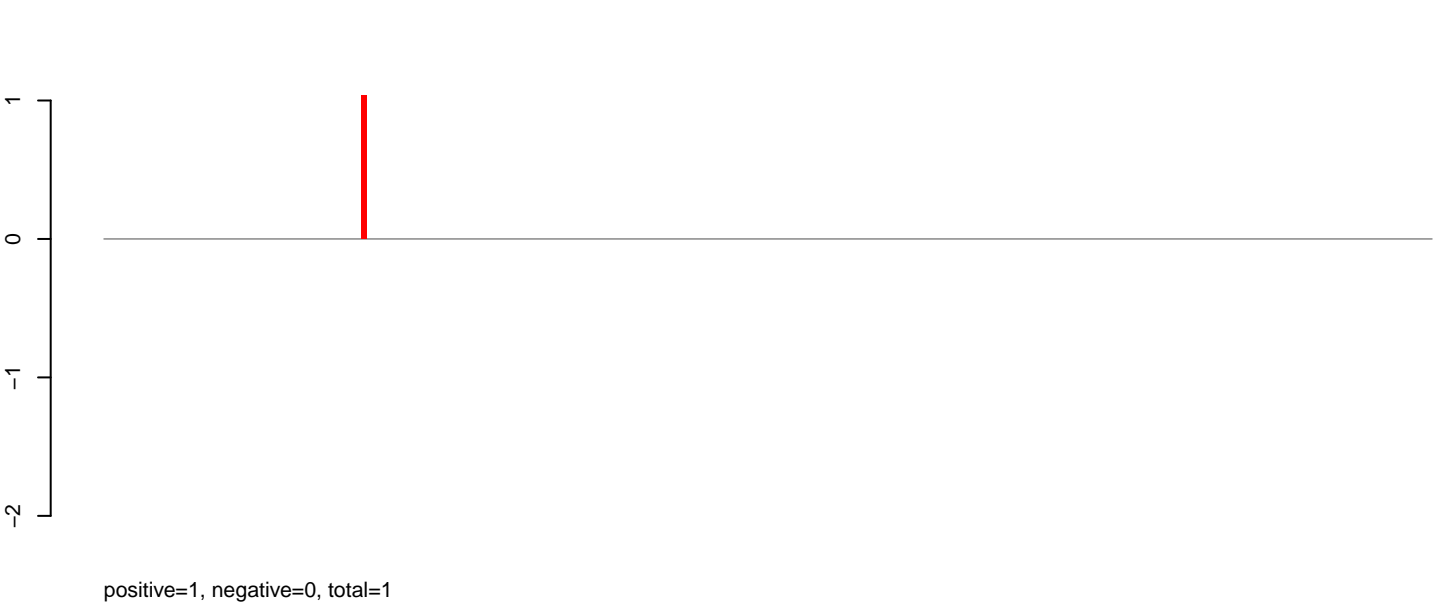
Window size=50, length=10158, virus@NC_030401.1-Hanko_virus_polyprotein_gene:1-10158

0 2000 4000 6000 8000 10000

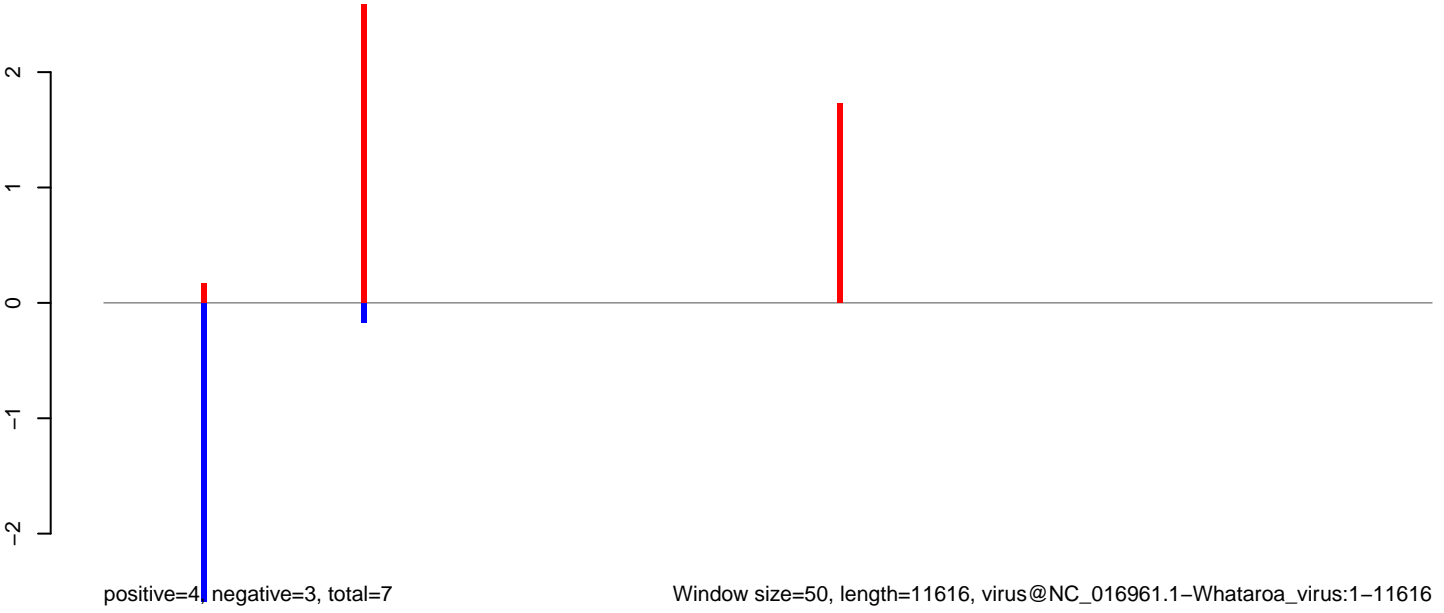
AeAeg_Aag2cells_CHIKV.18_23.rep



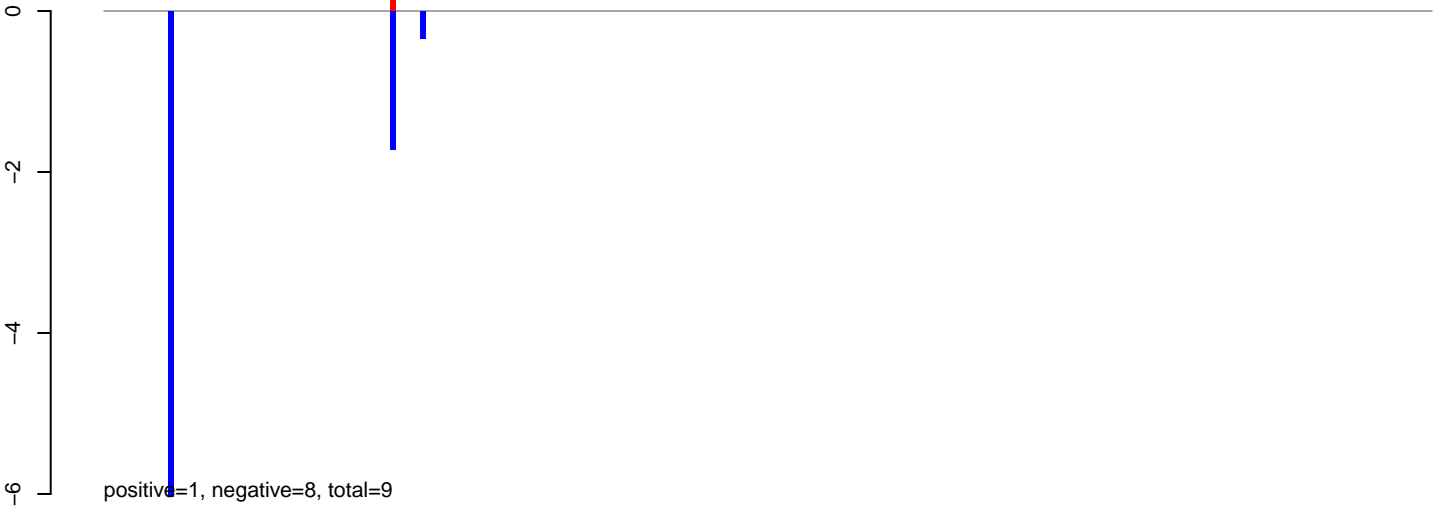
AeAeg_Aag2cells_CHIKV.24_35.rep



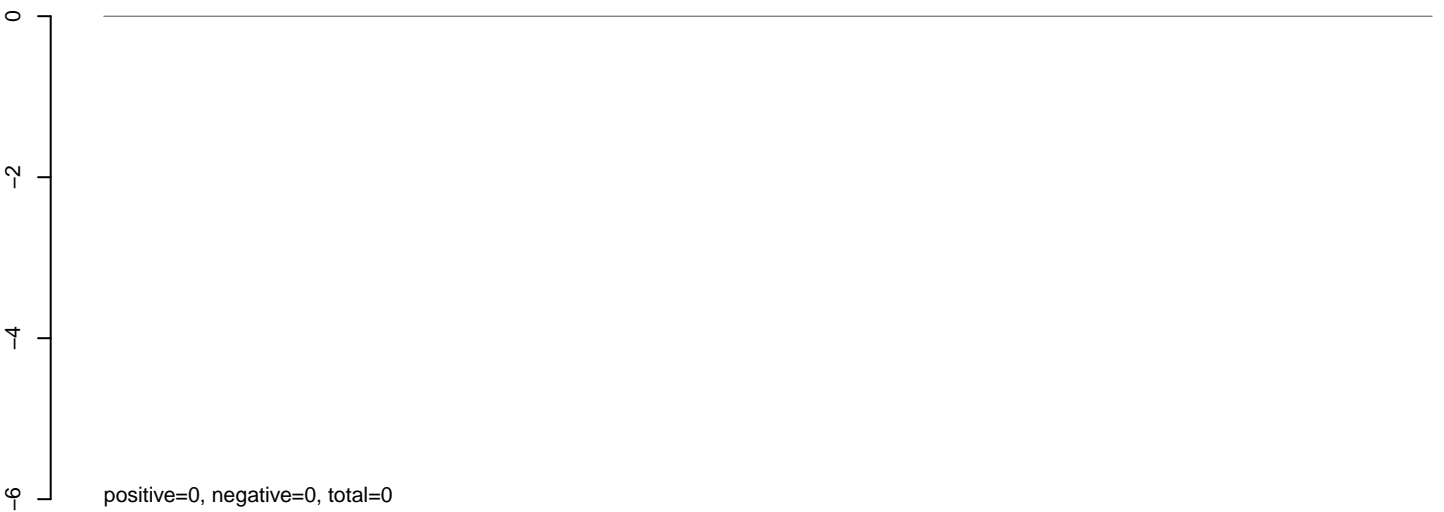
AeAeg_Aag2cells_CHIKV.rep



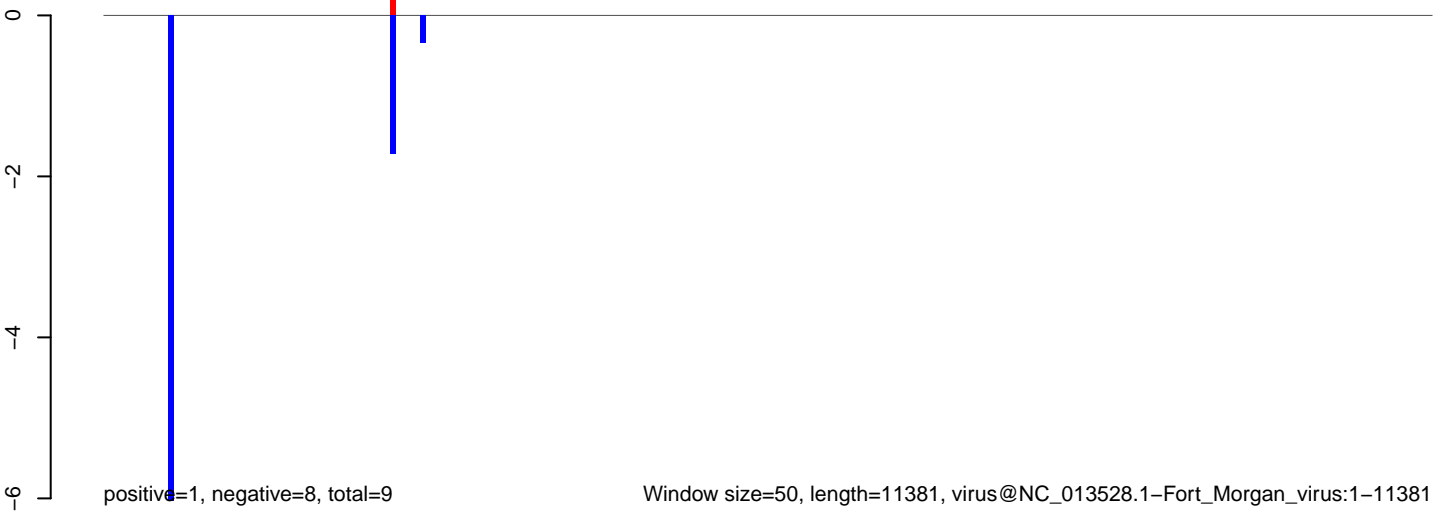
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



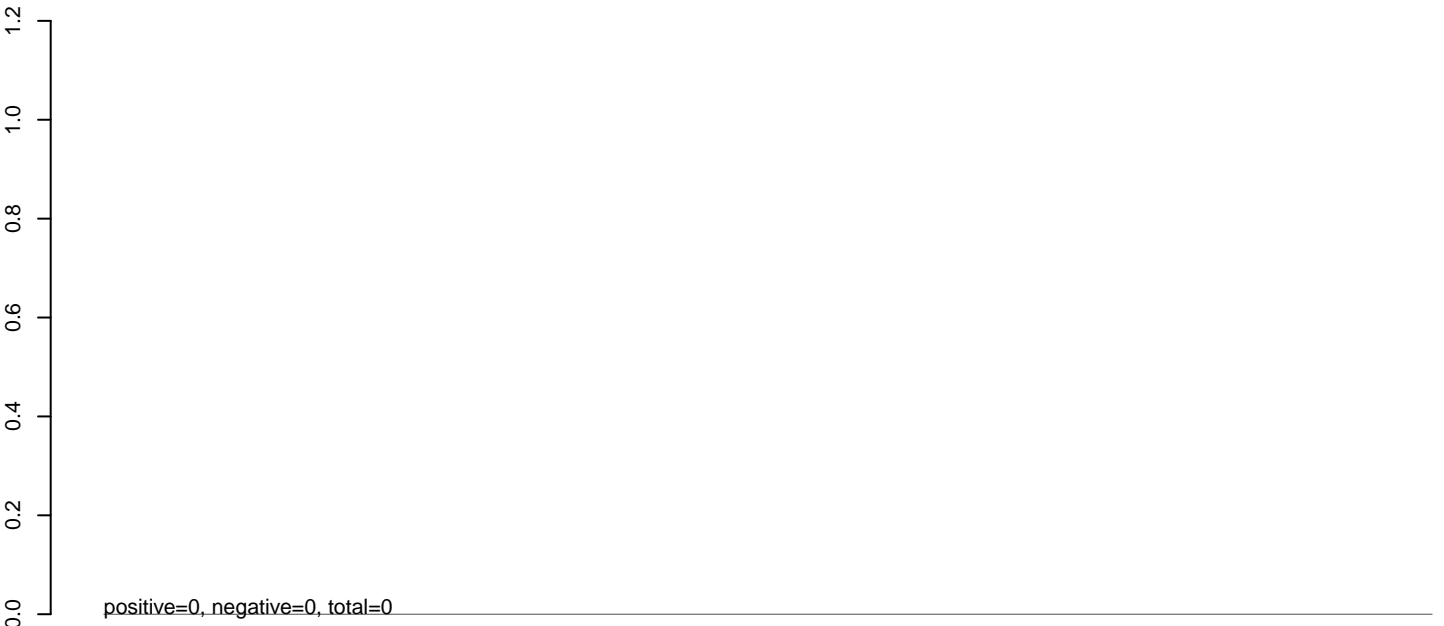
AeAeg_Aag2cells_CHIKV.rep



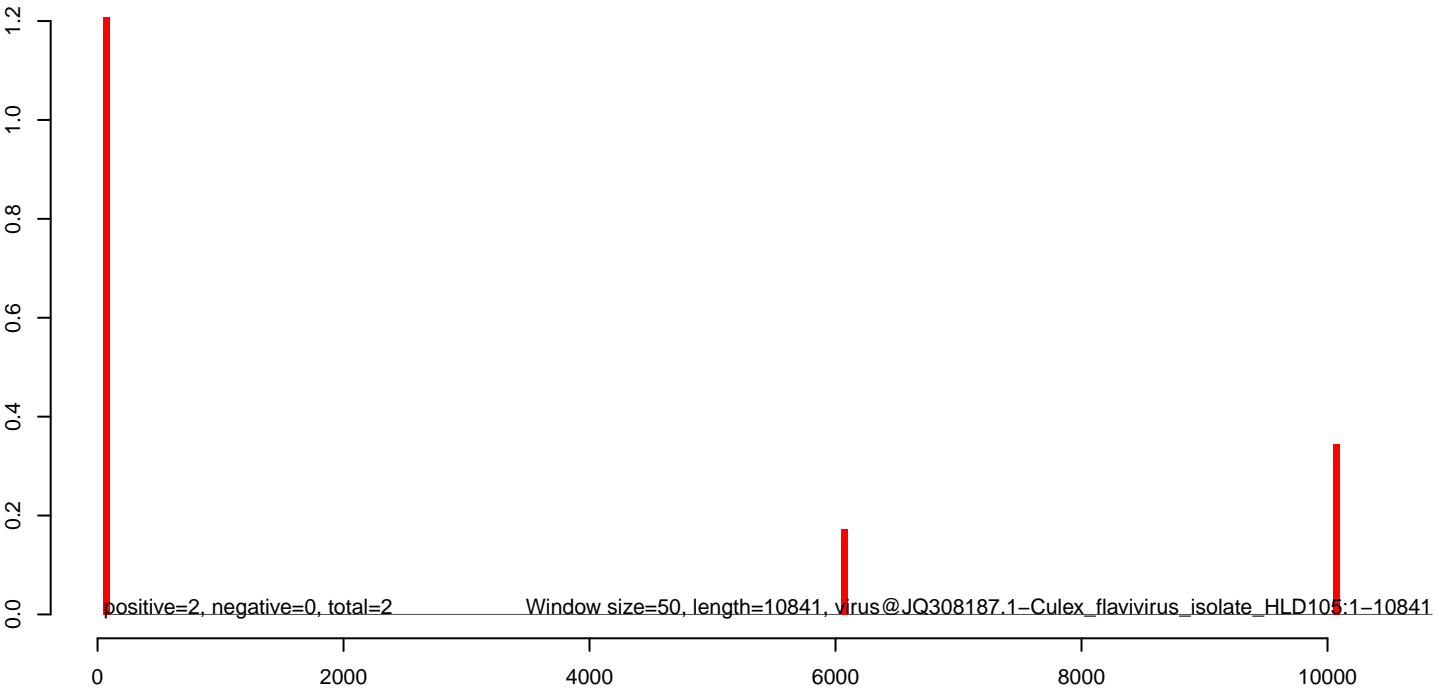
AeAeg_Aag2cells_CHIKV.18_23.rep



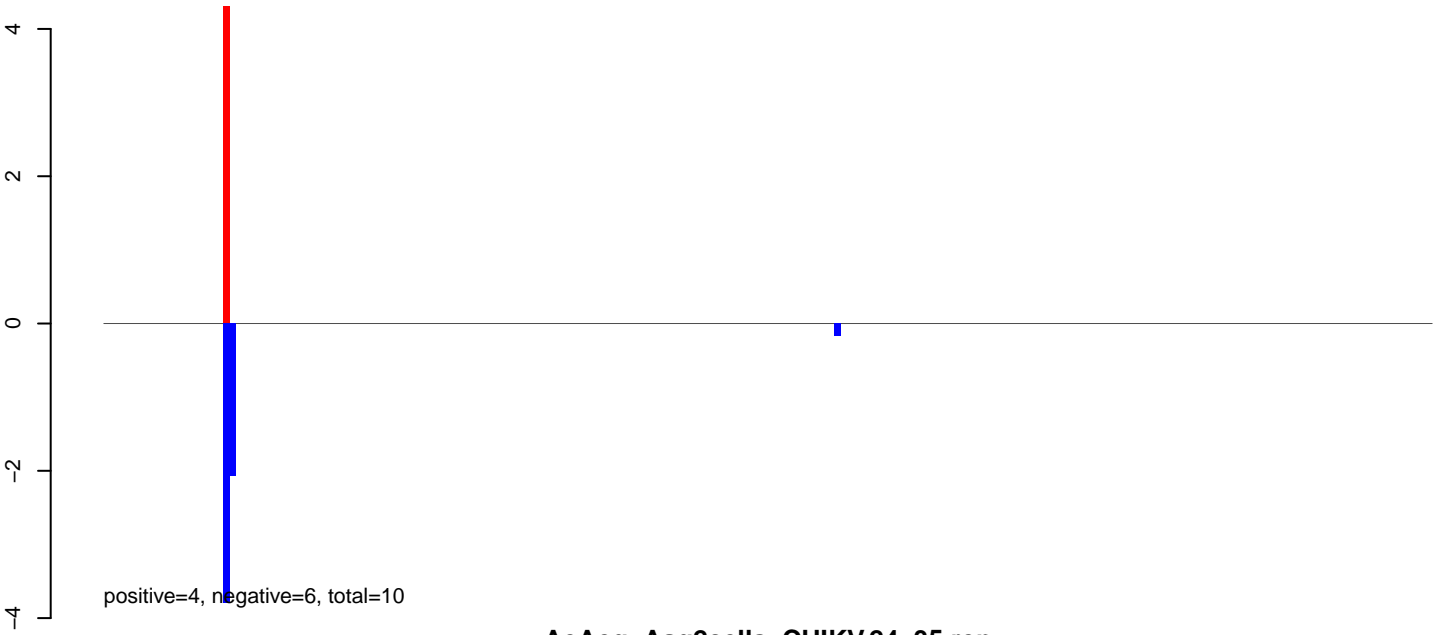
AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep



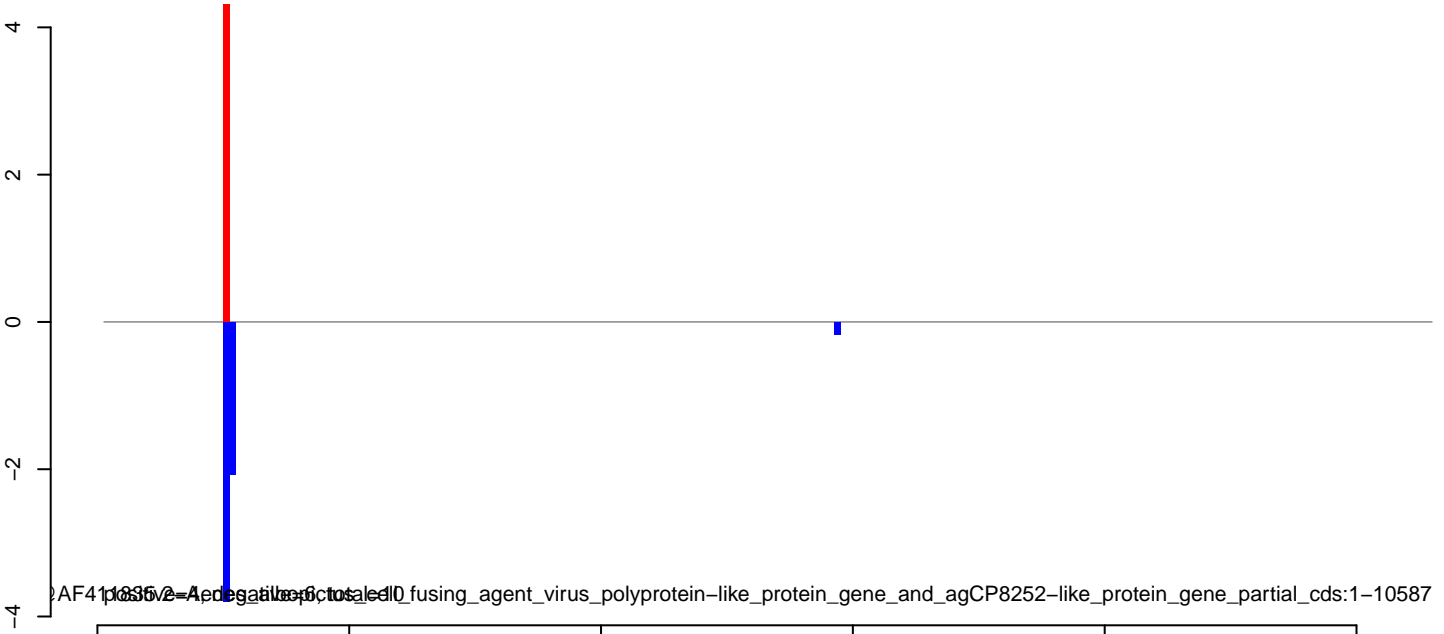
AeAeg_Aag2cells_CHIKV.18_23.rep



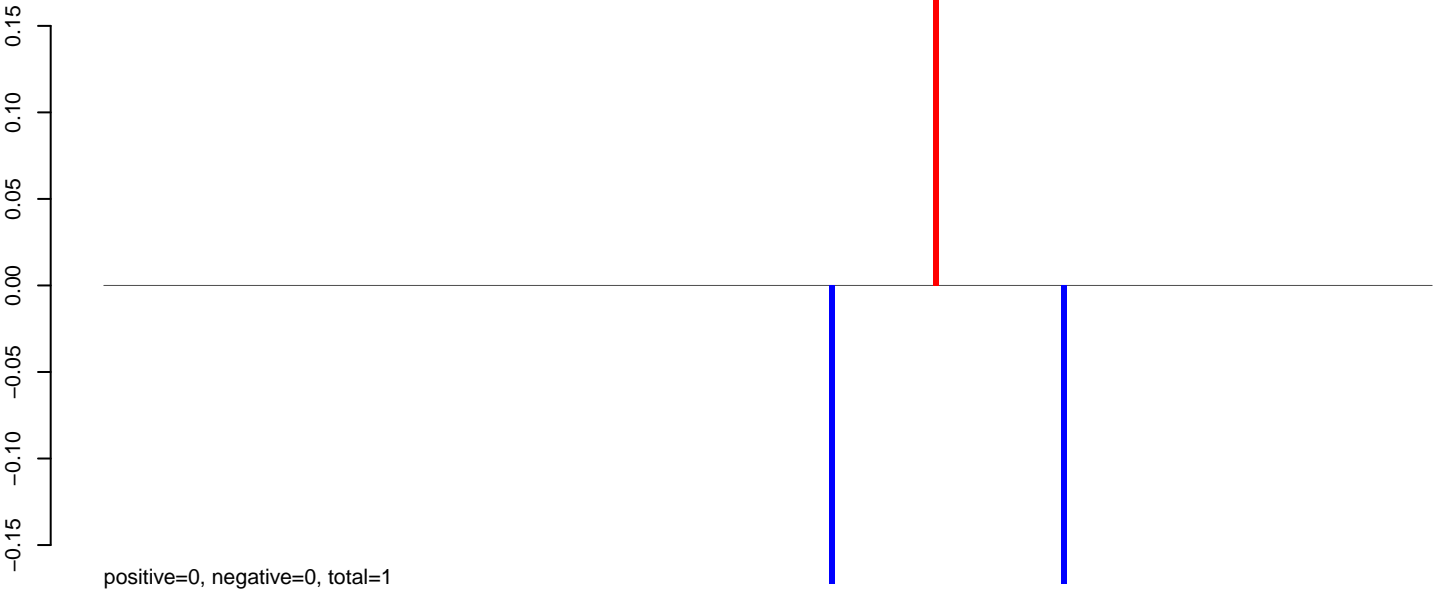
AeAeg_Aag2cells_CHIKV.24_35.rep



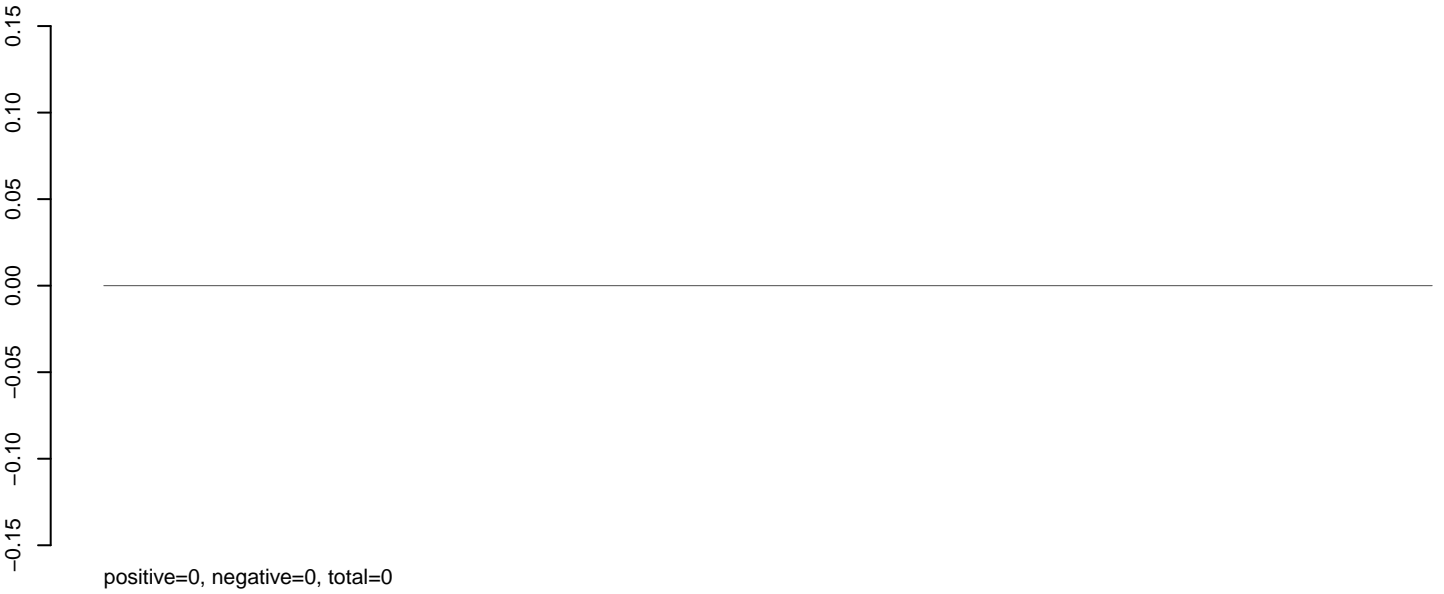
AeAeg_Aag2cells_CHIKV.rep



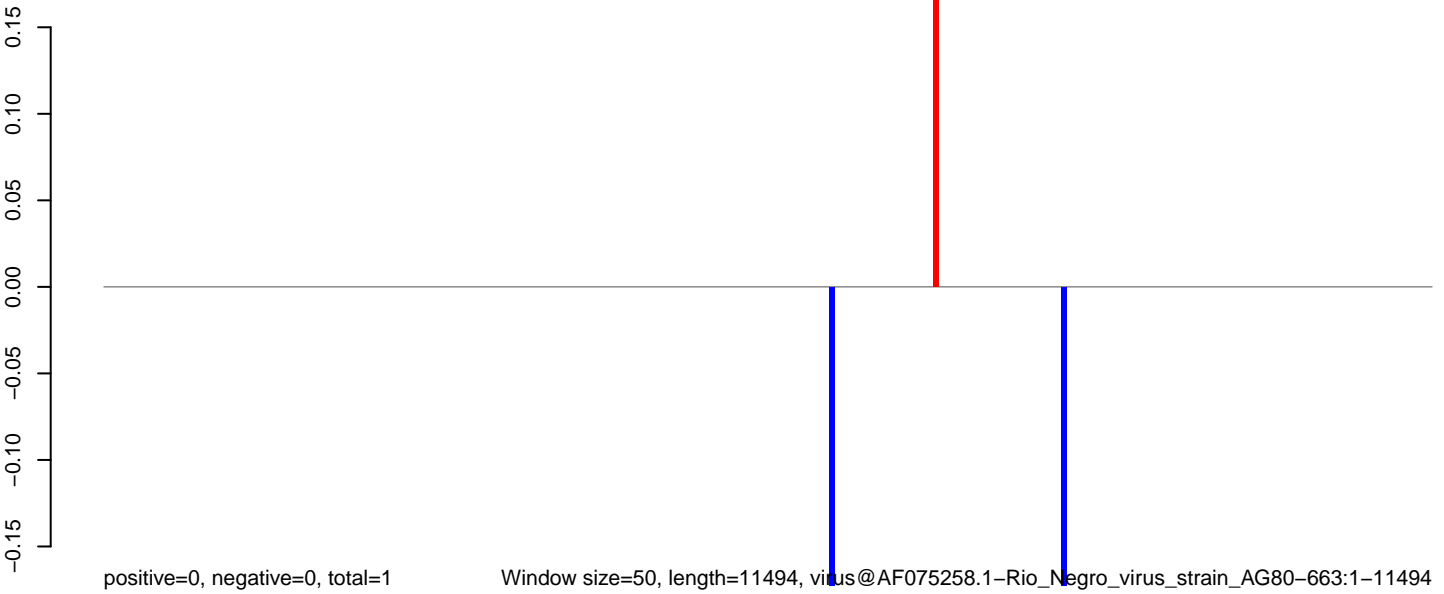
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

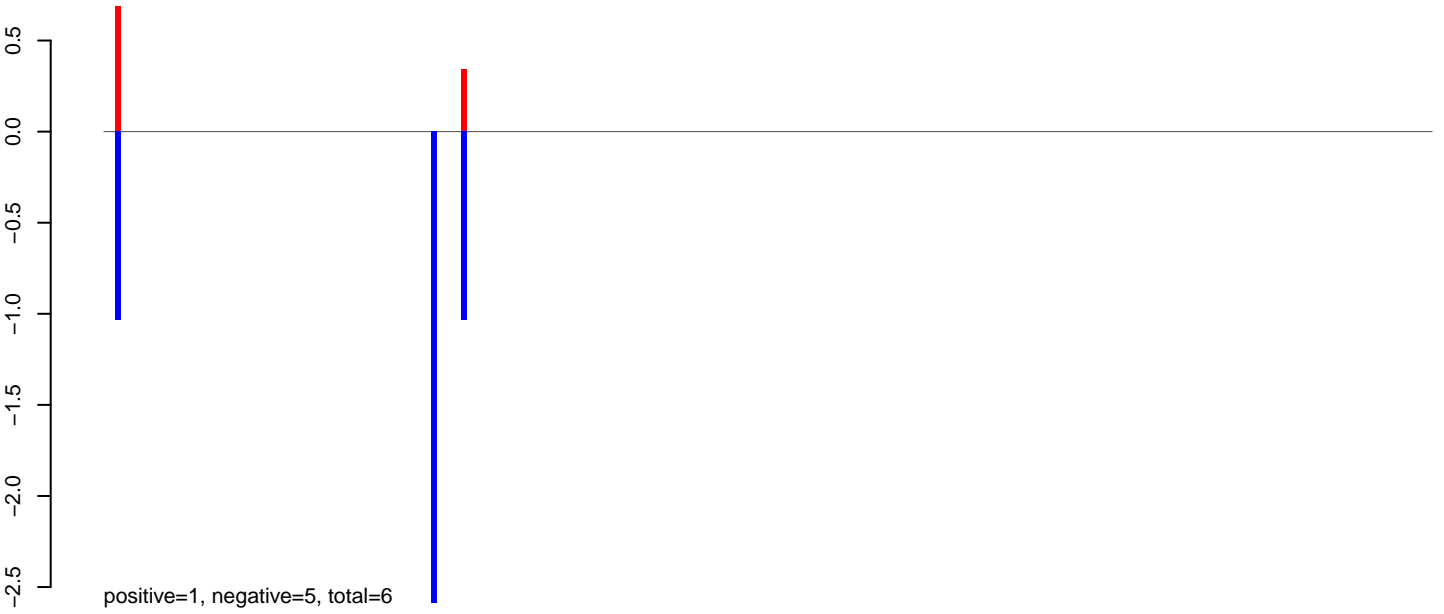


AeAeg_Aag2cells_CHIKV.rep



0 2000 4000 6000 8000 10000

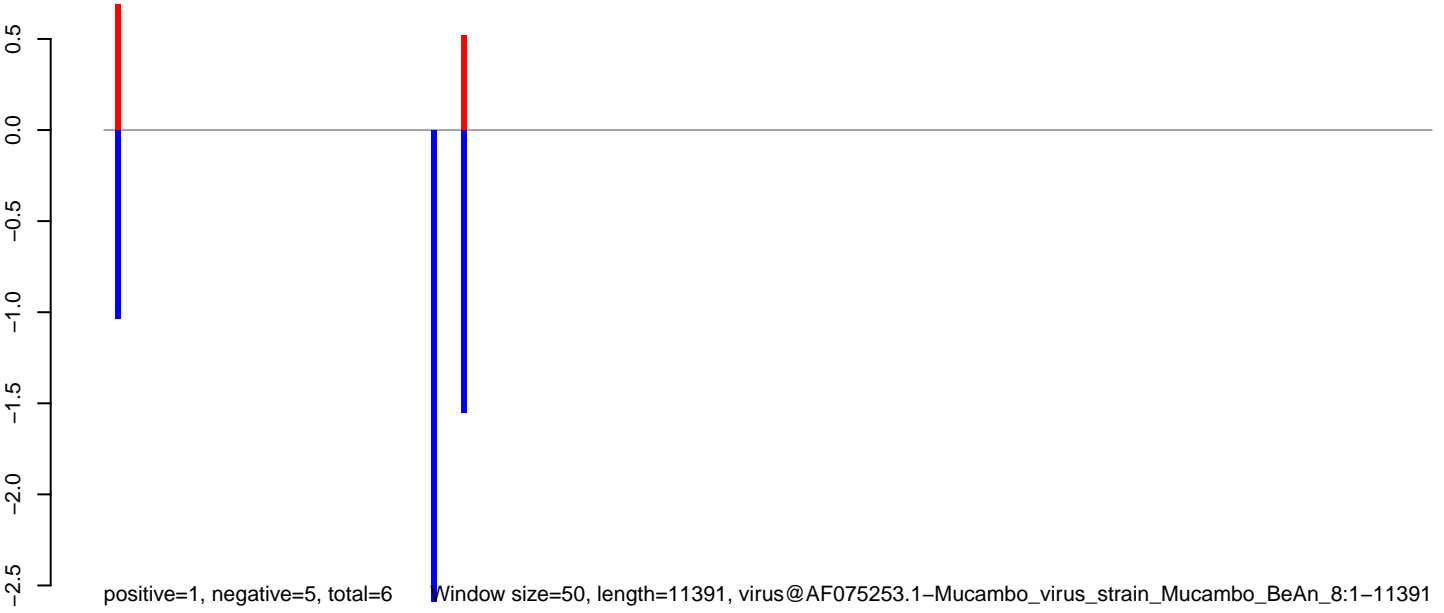
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



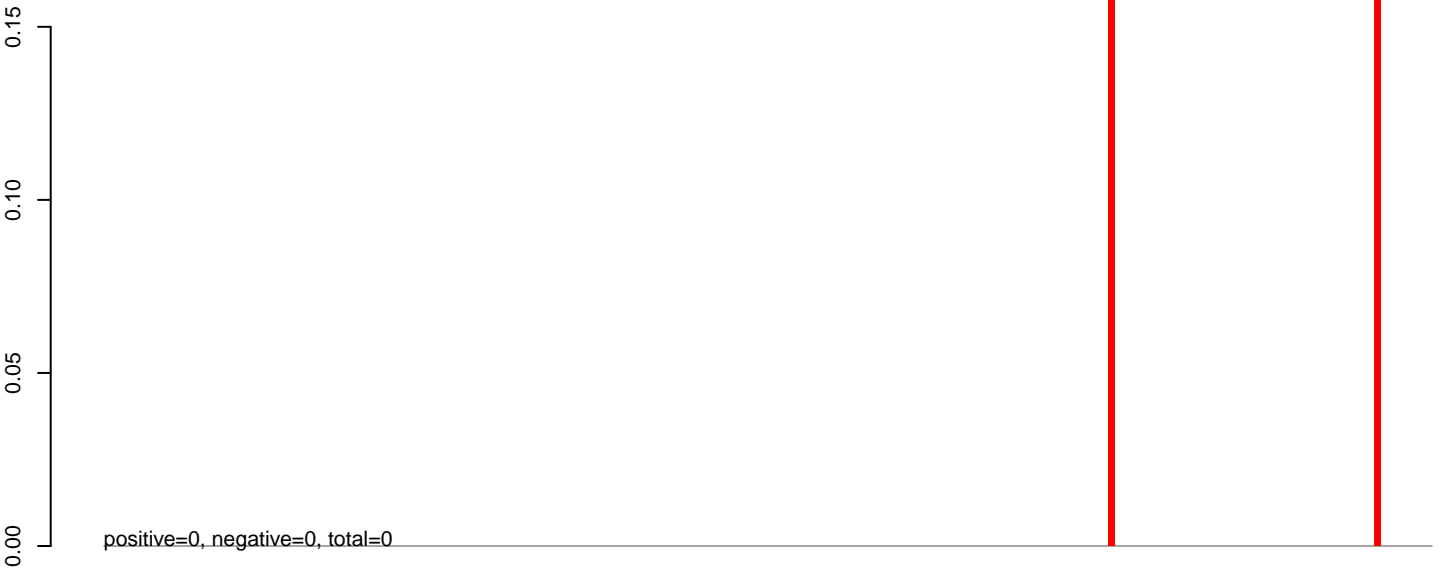
AeAeg_Aag2cells_CHIKV.rep



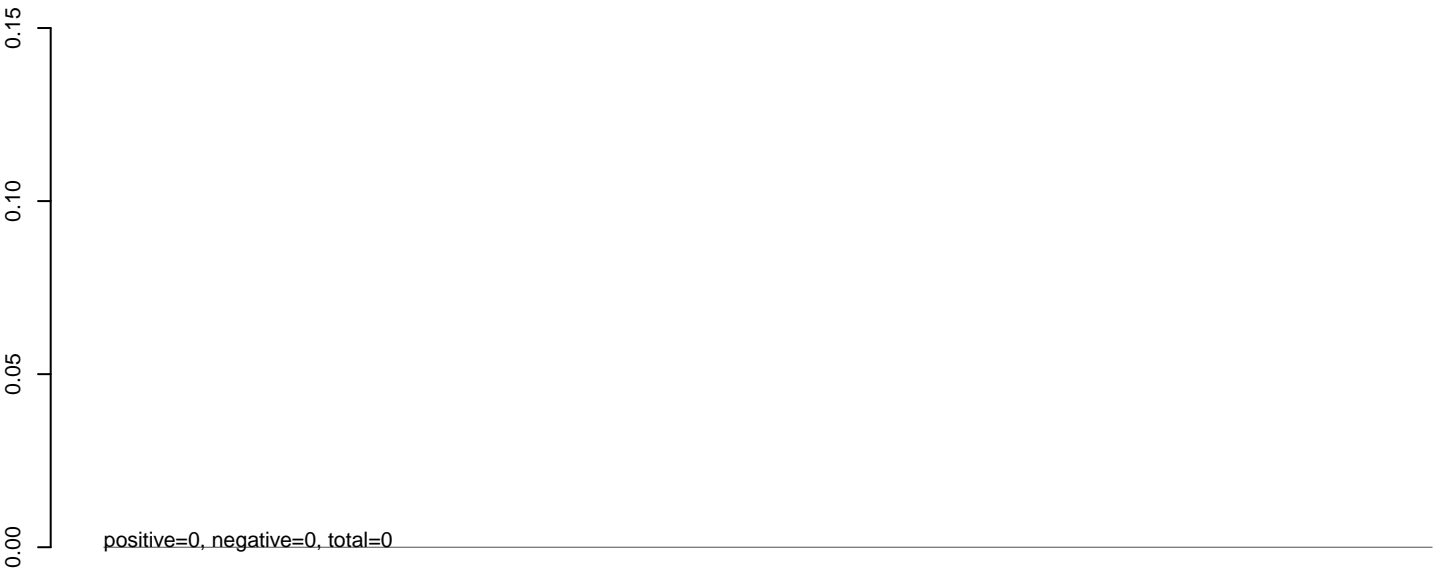
Window size=50, length=11391, virus@AF075253.1-Mucambo_virus_strain_Mucambo_BeAn_8:1-11391

0 2000 4000 6000 8000 10000

AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

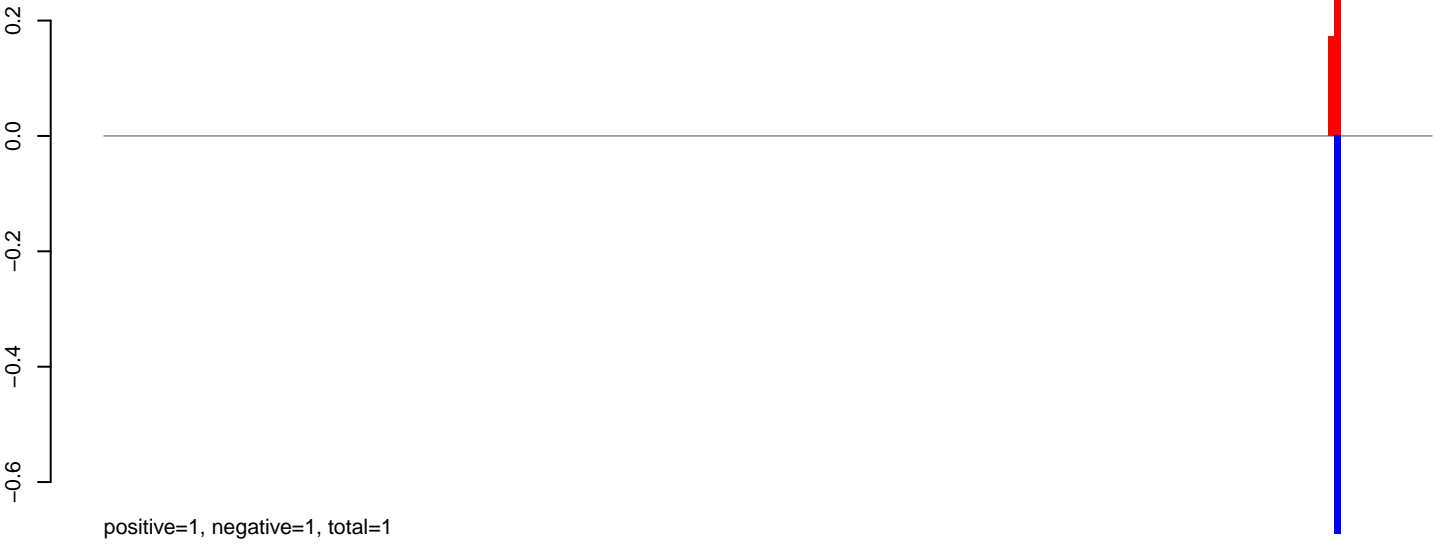


AeAeg_Aag2cells_CHIKV.rep

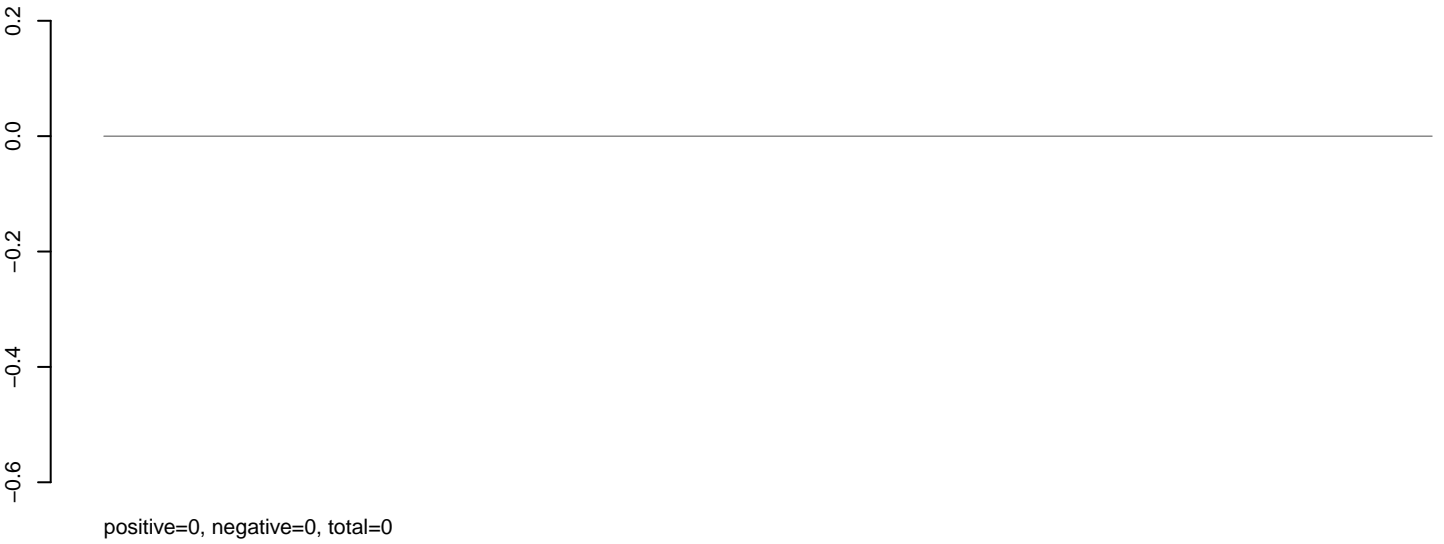


0 2000 4000 6000 8000 10000

AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



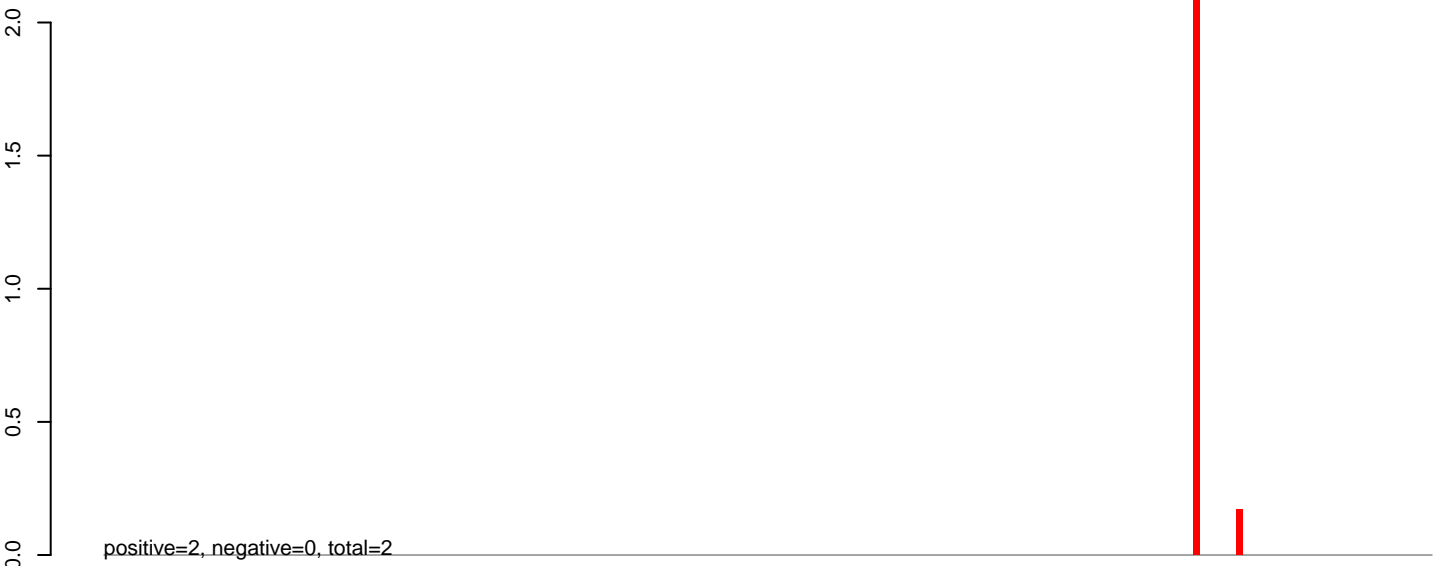
AeAeg_Aag2cells_CHIKV.rep



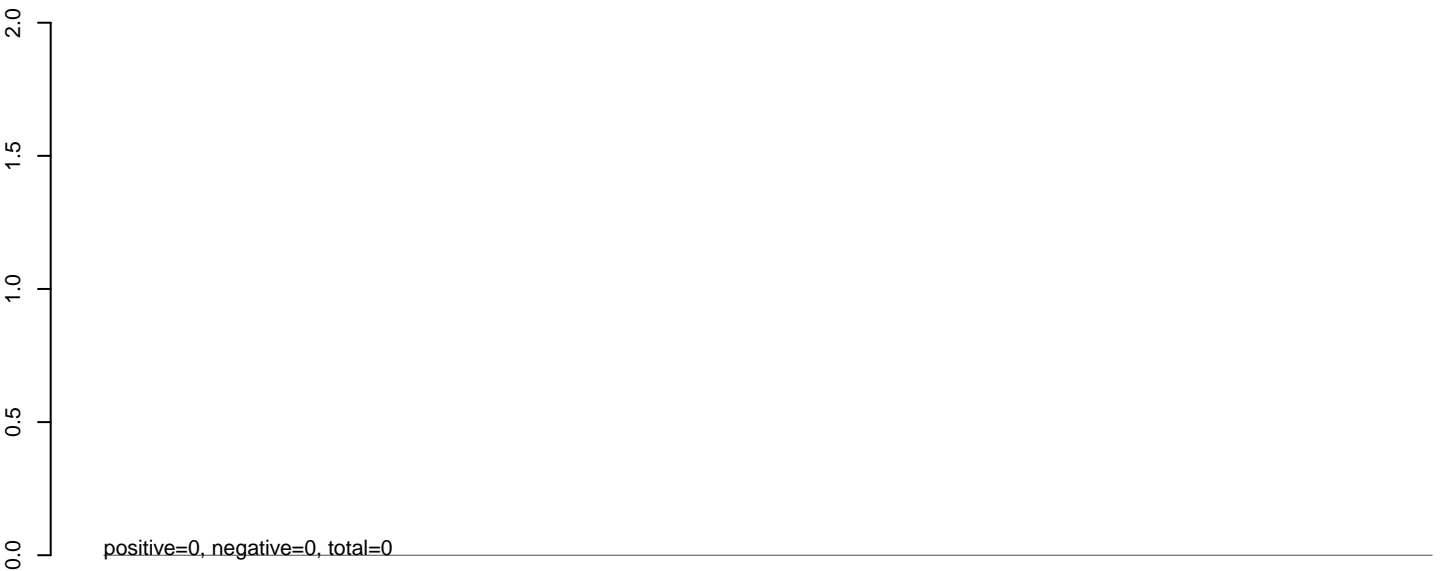
Window size=50, length=10938, virus@NC_027819.1-Mercadeo_virus_isolate_ER-M10:1-10938

0 2000 4000 6000 8000 10000

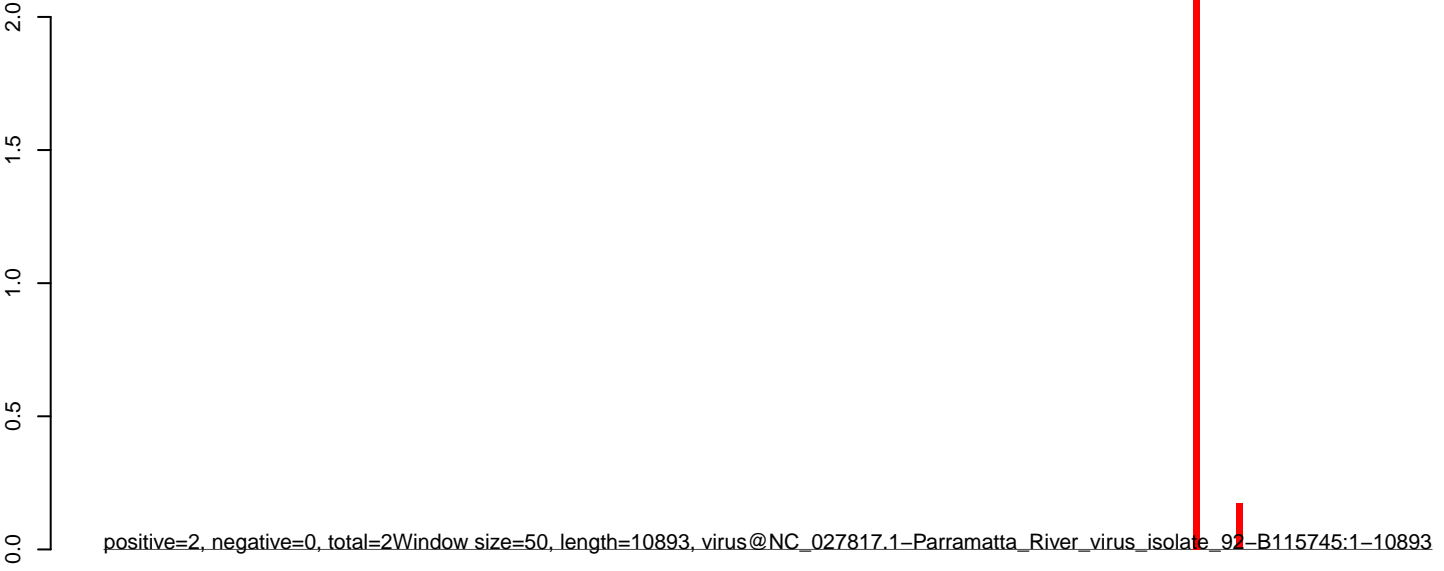
AeAeg_Aag2cells_CHIKV.18_23.rep



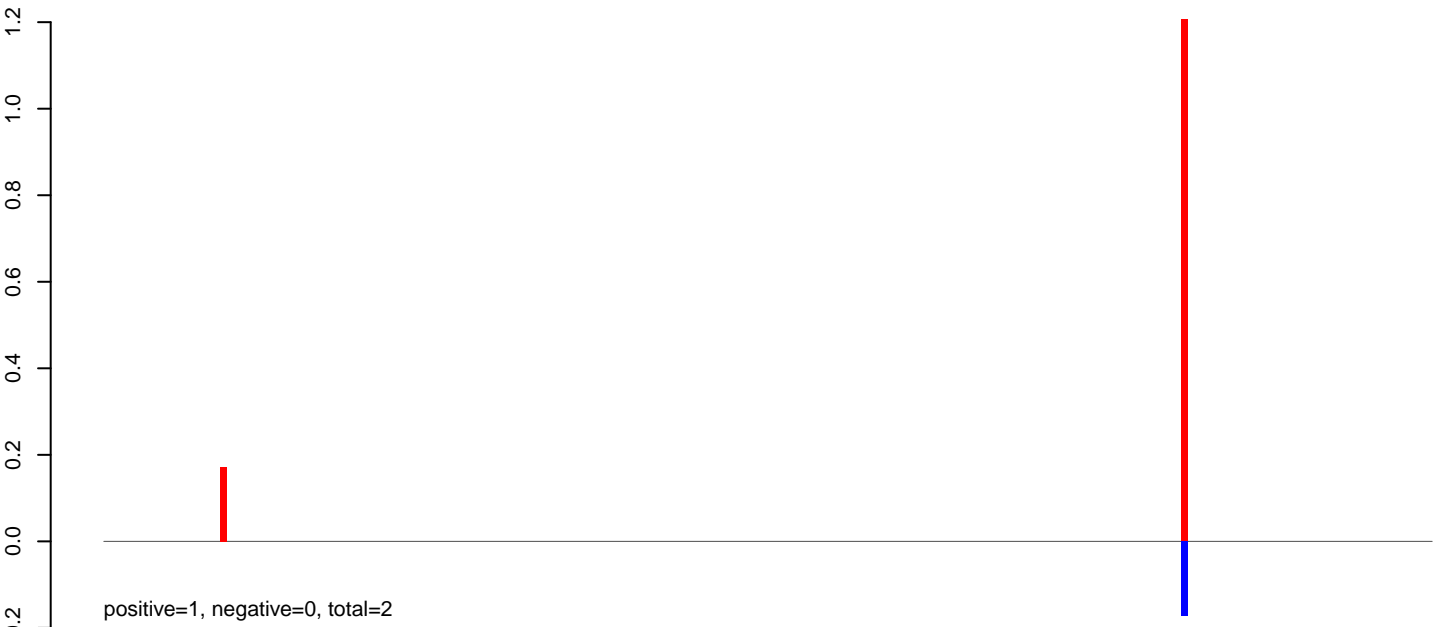
AeAeg_Aag2cells_CHIKV.24_35.rep



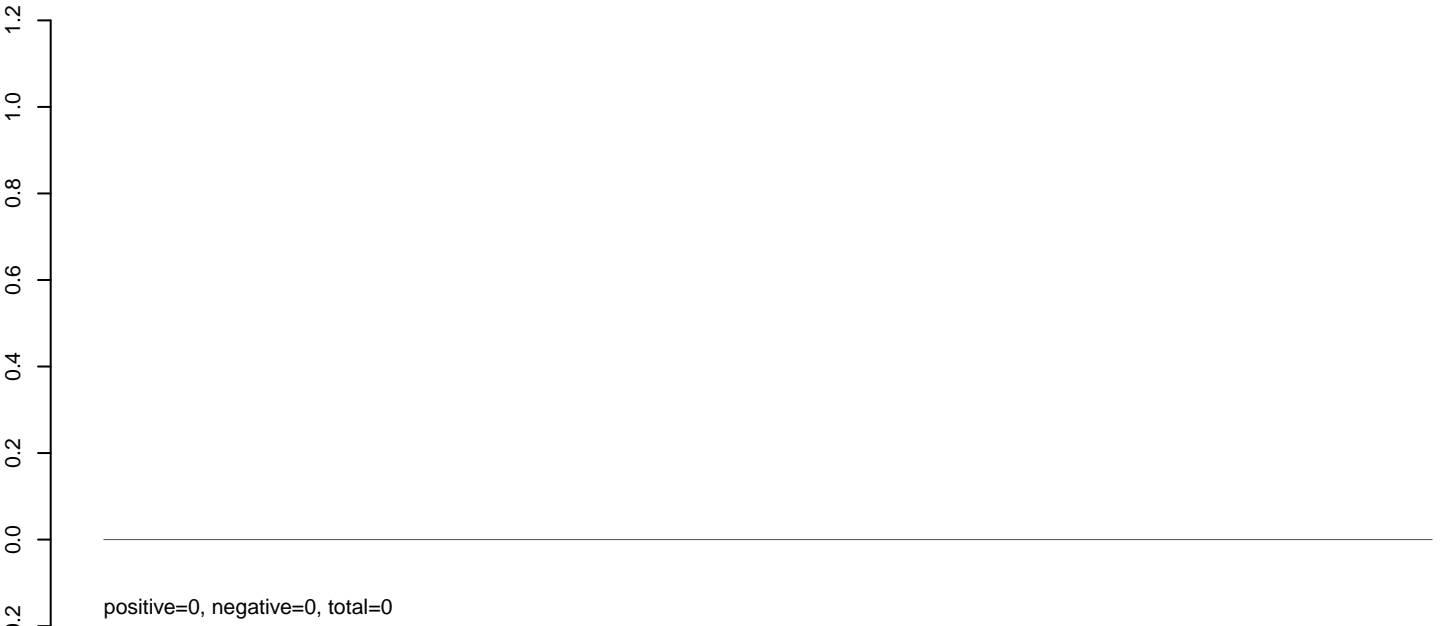
AeAeg_Aag2cells_CHIKV.rep



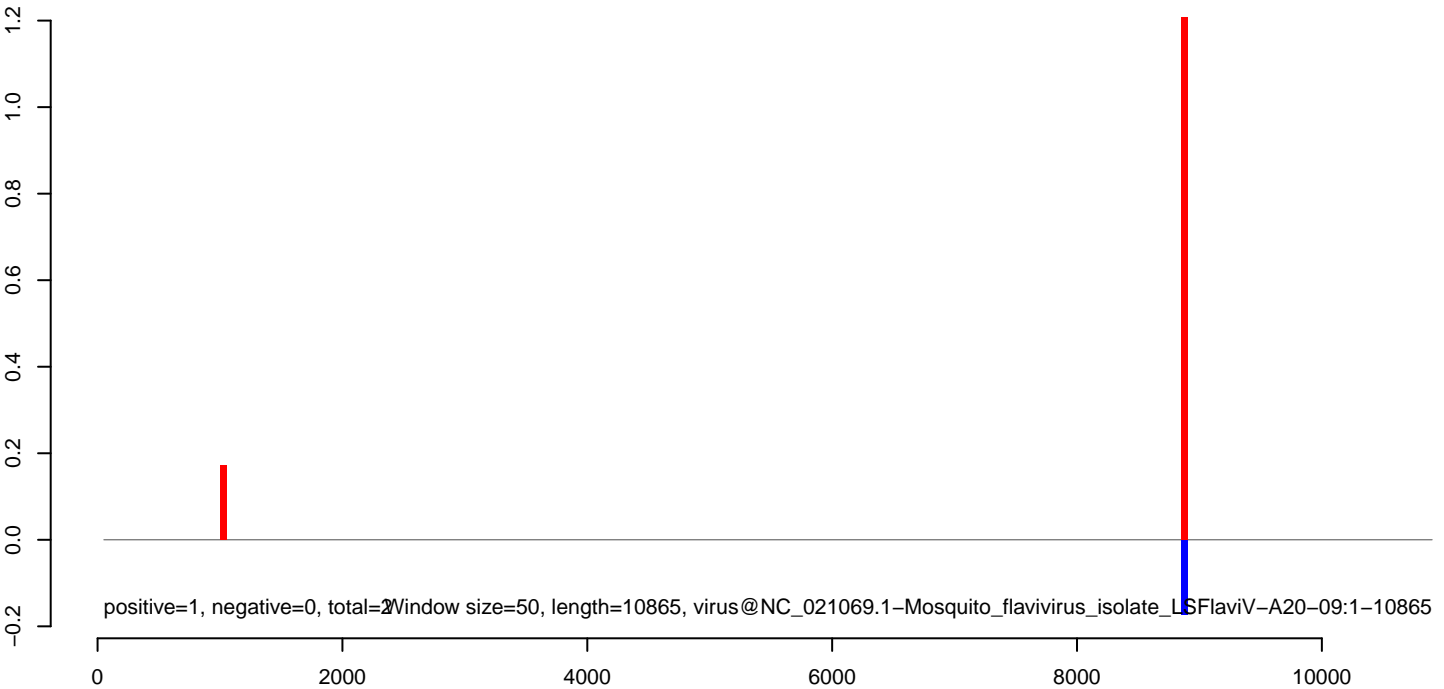
AeAeg_Aag2cells_CHIKV.18_23.rep



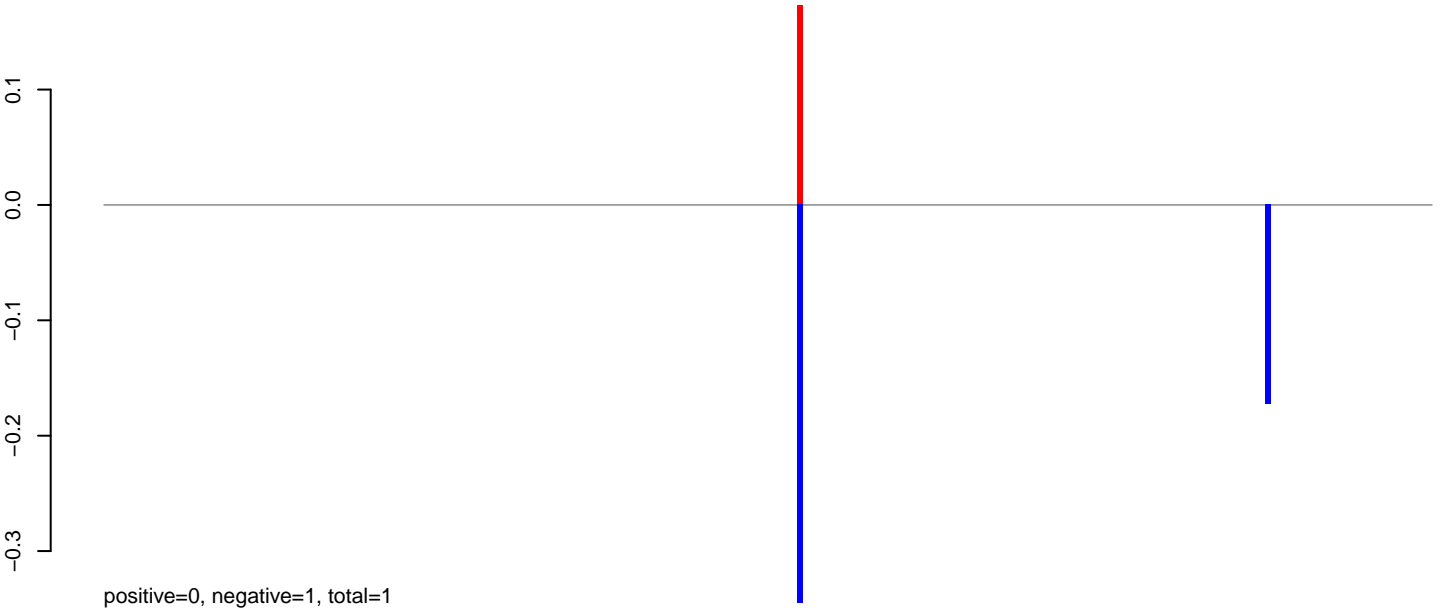
AeAeg_Aag2cells_CHIKV.24_35.rep



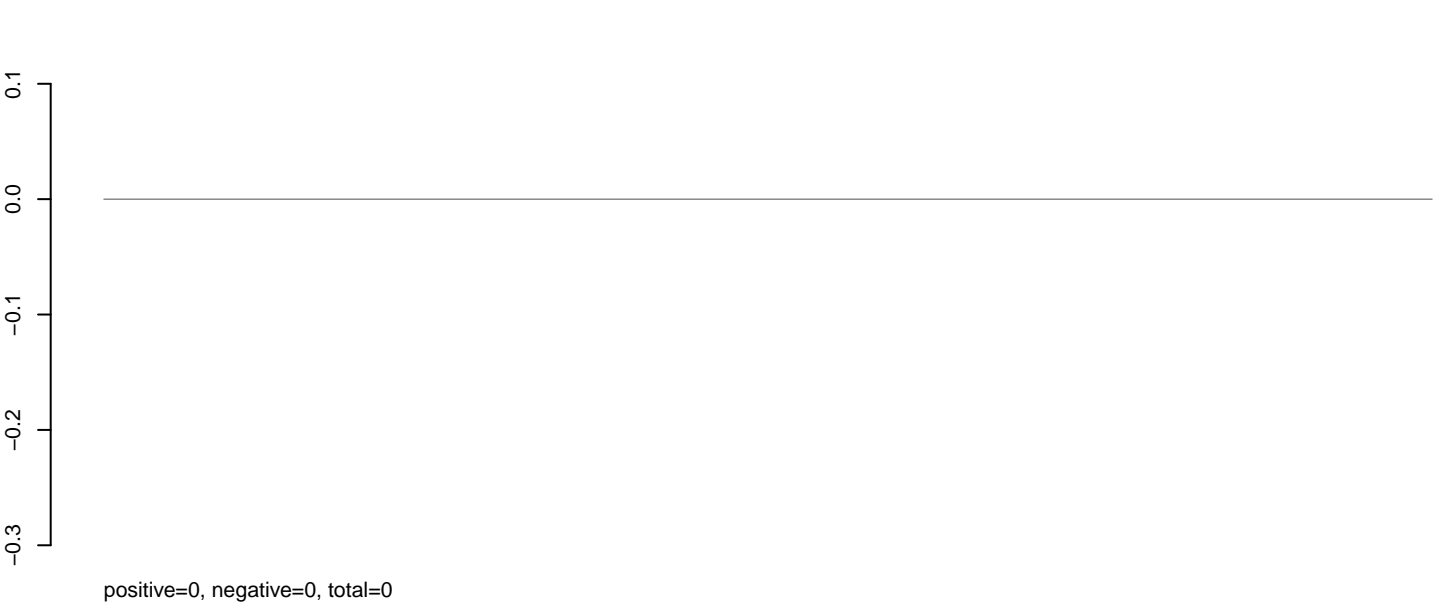
AeAeg_Aag2cells_CHIKV.rep



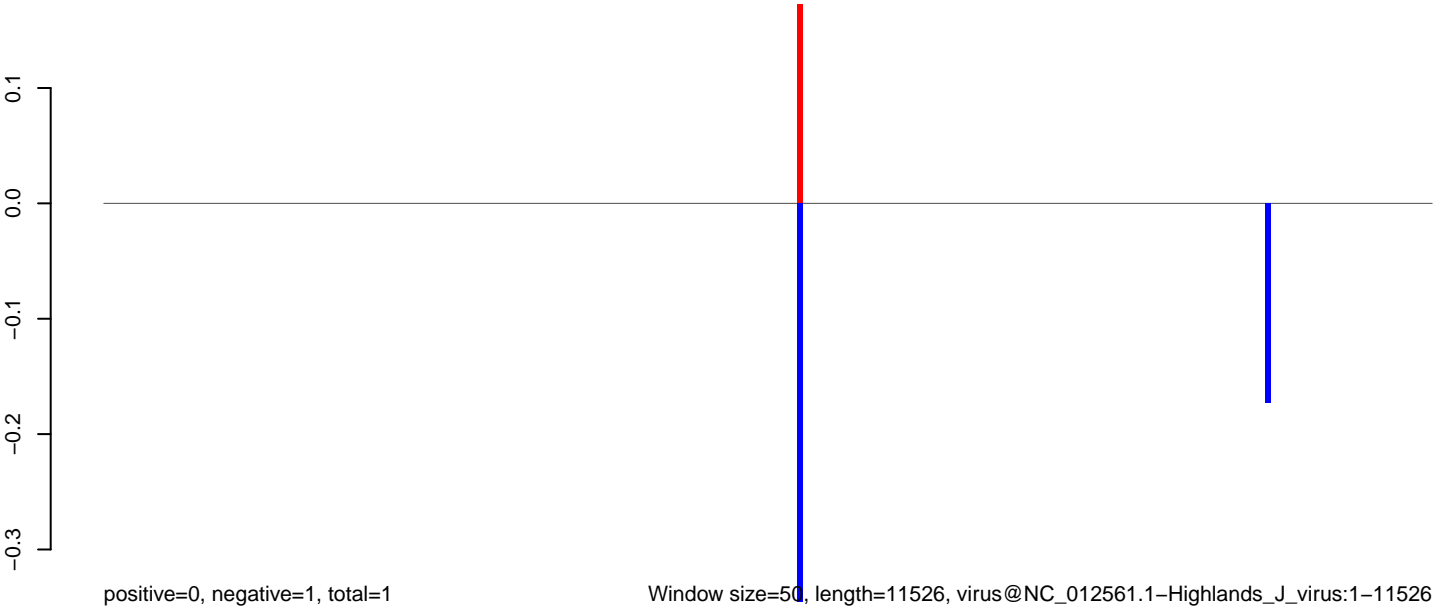
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



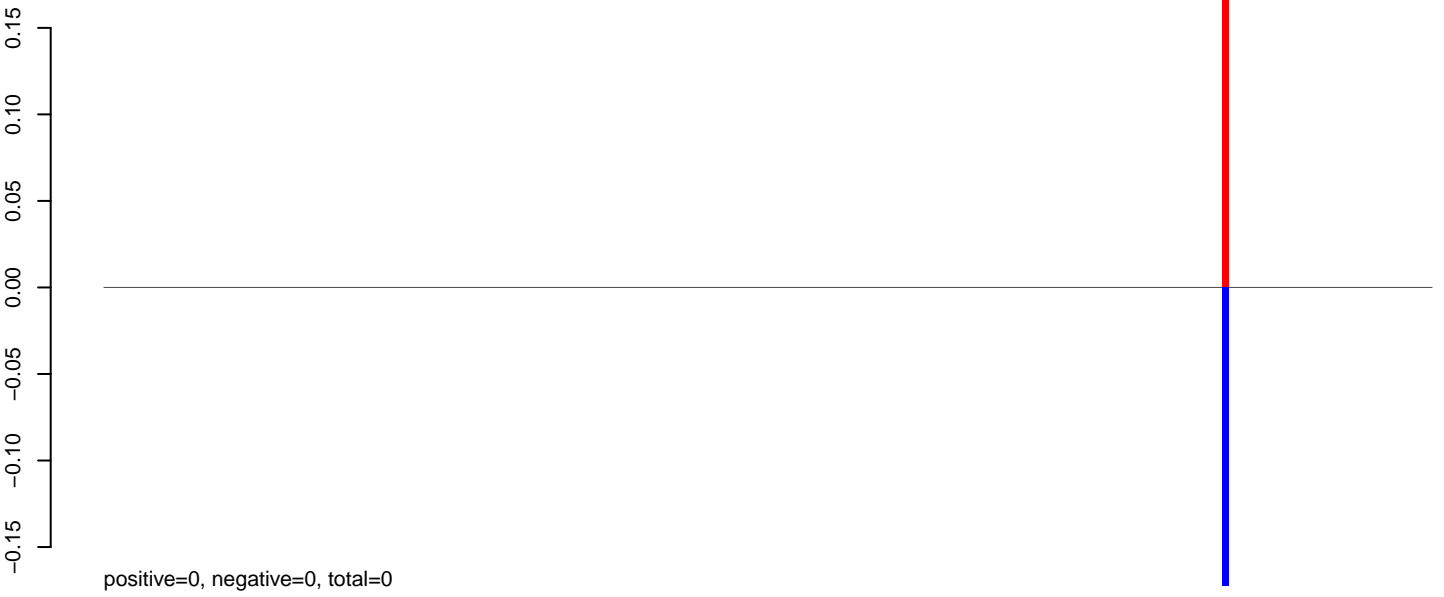
AeAeg_Aag2cells_CHIKV.rep



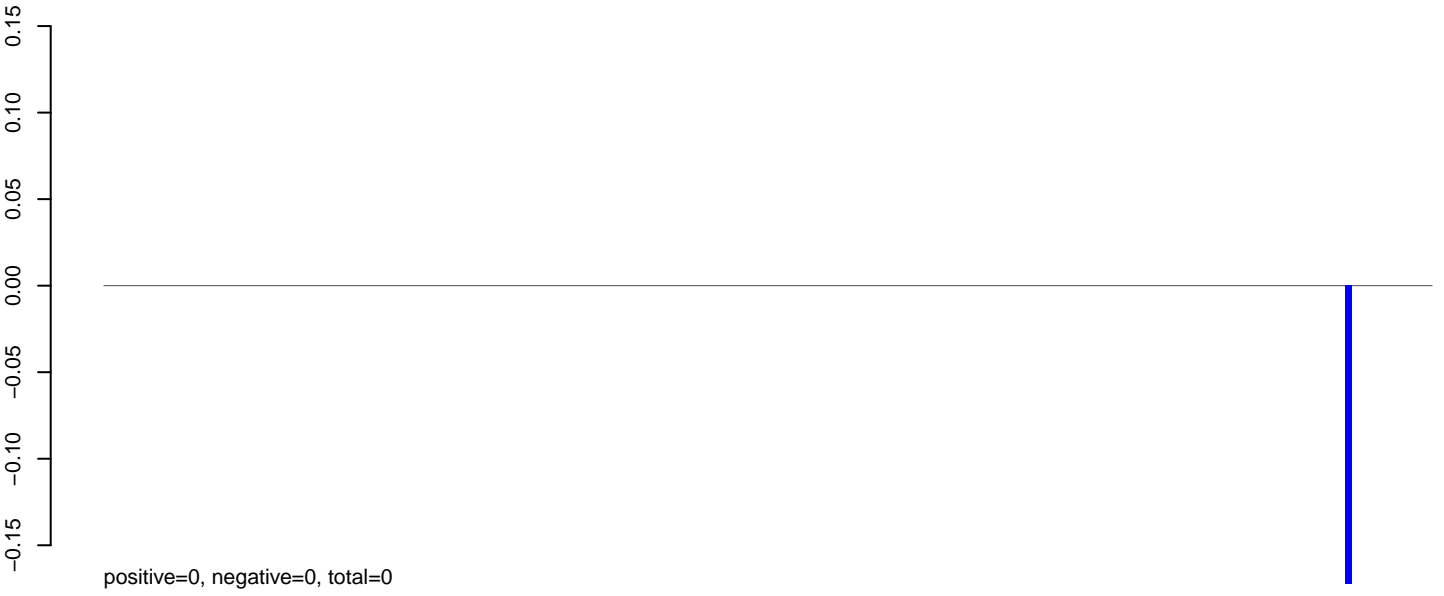
Window size=50, length=11526, virus@NC_012561.1-Highlands_J_virus:1-11526

0 2000 4000 6000 8000 10000 12000

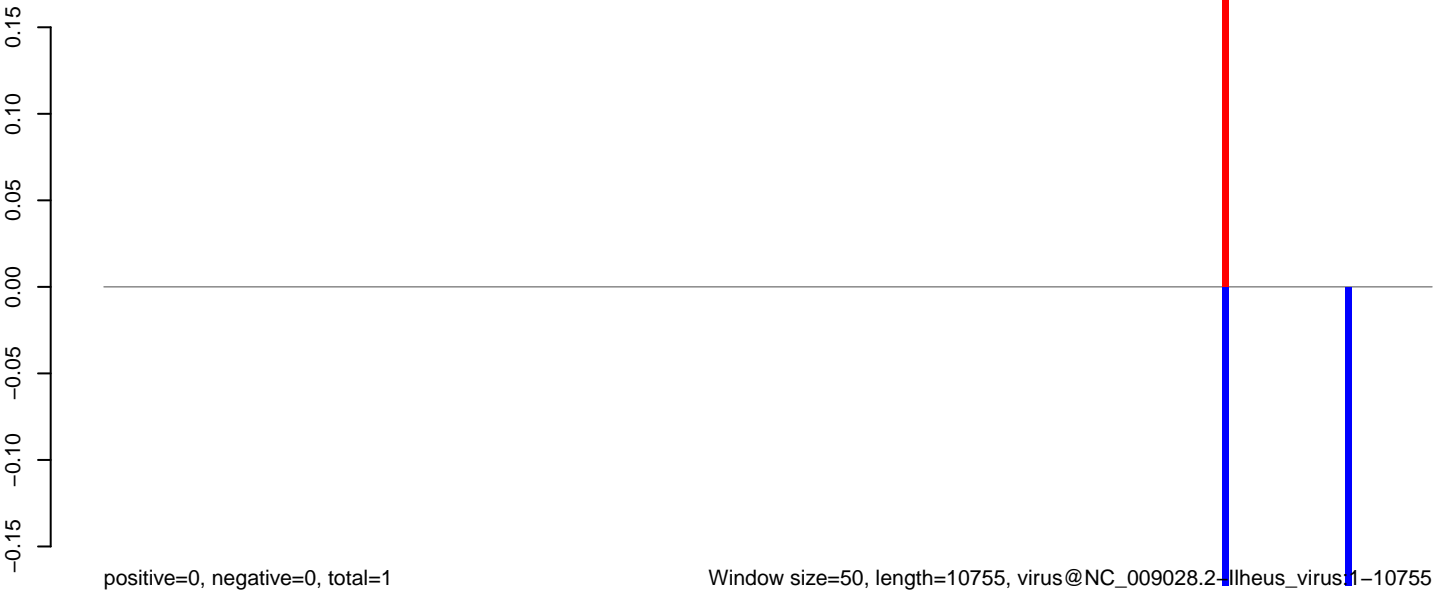
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



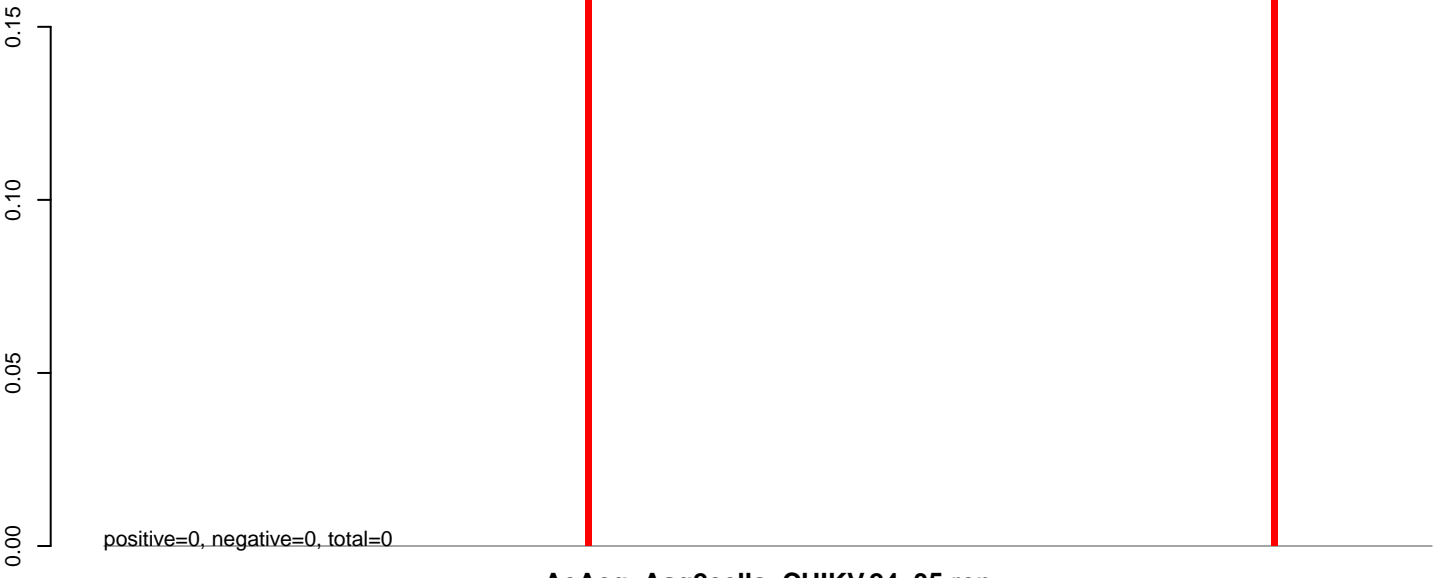
AeAeg_Aag2cells_CHIKV.rep



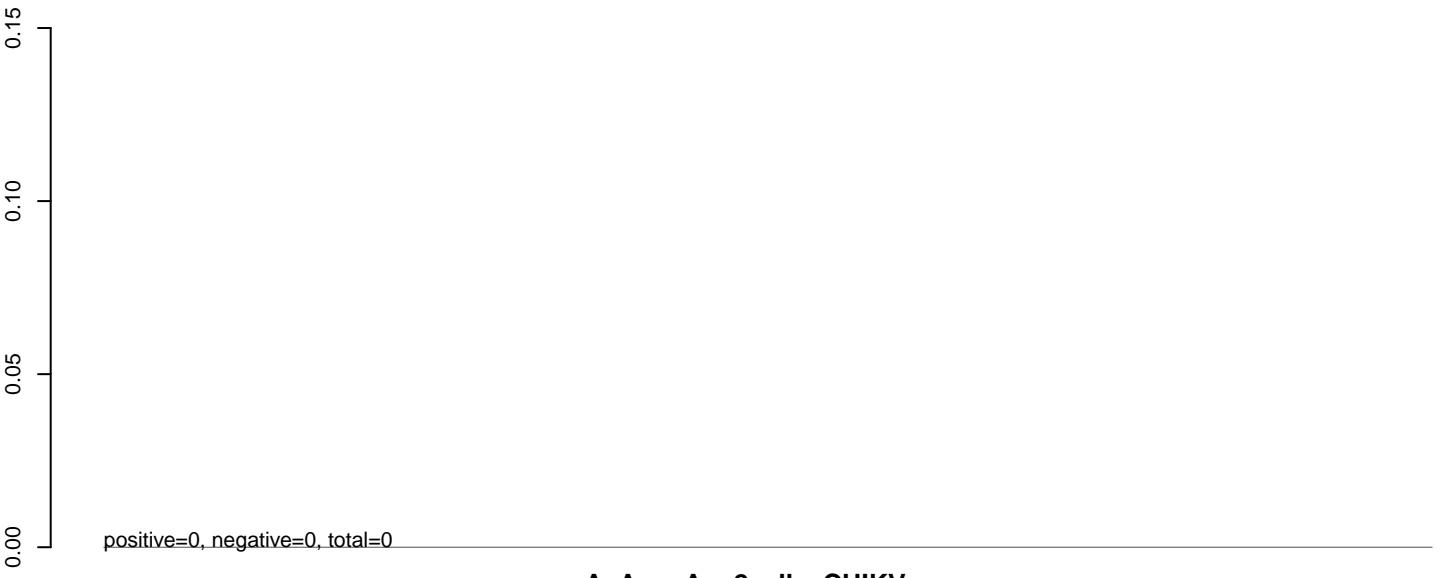
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0 2000 4000 6000 8000 10000

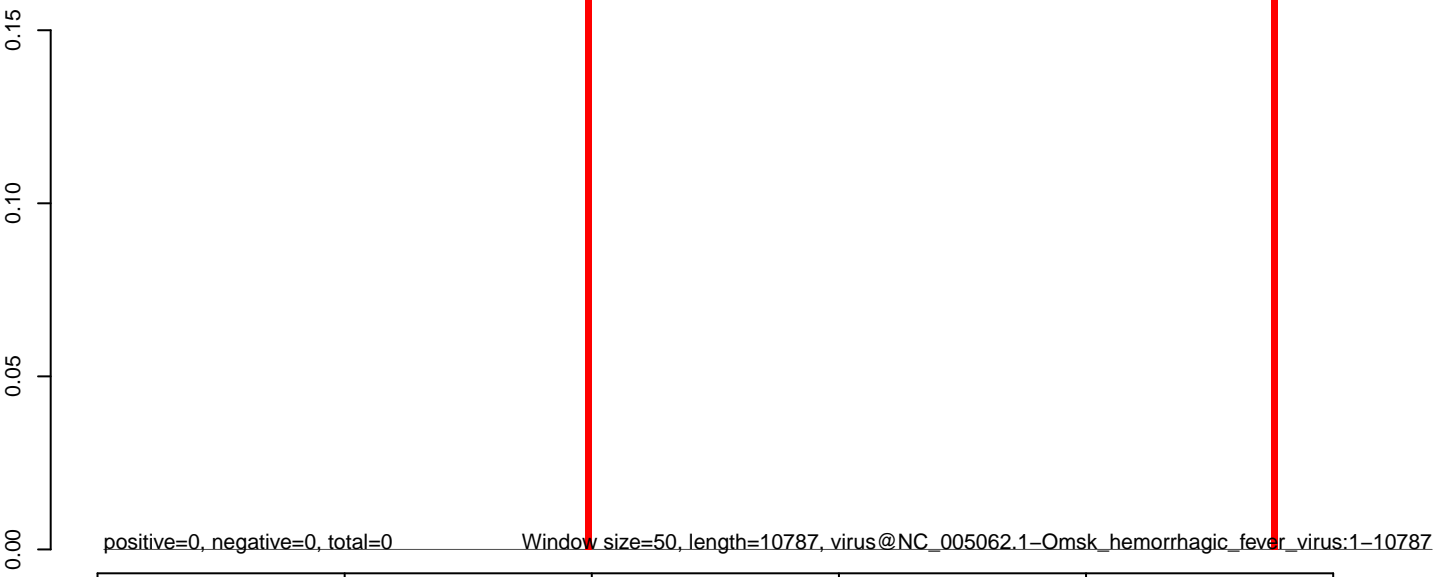
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

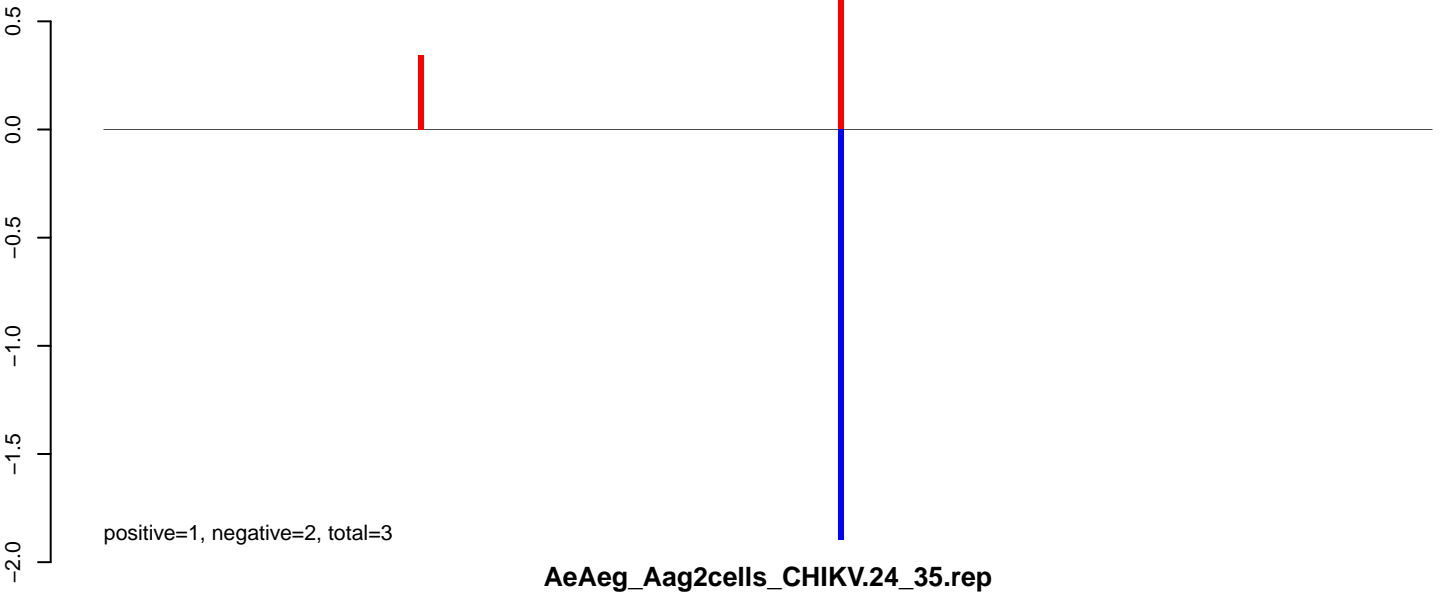


AeAeg_Aag2cells_CHIKV.rep

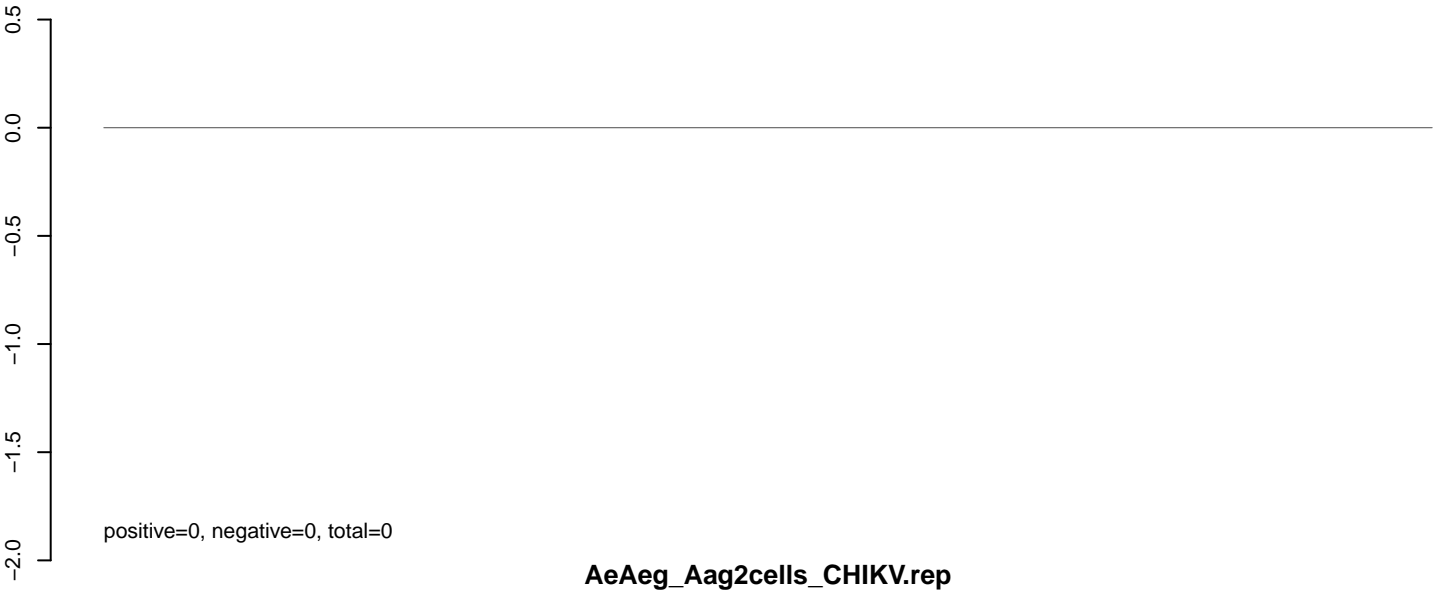


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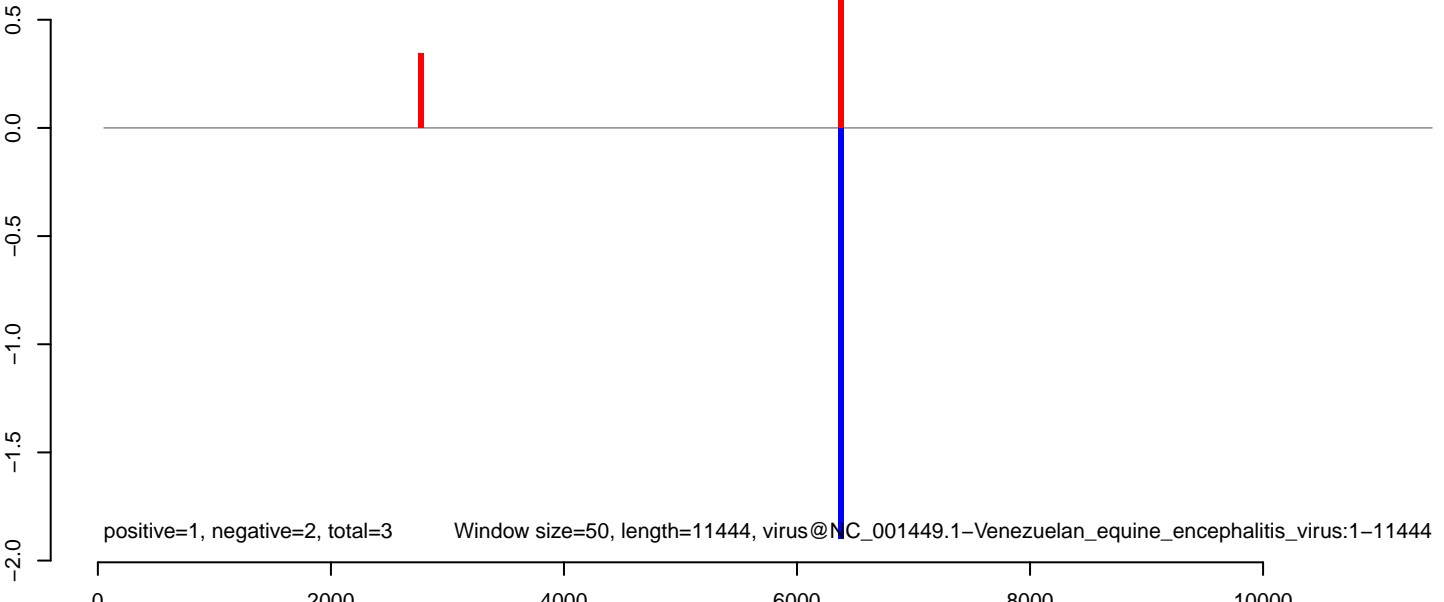
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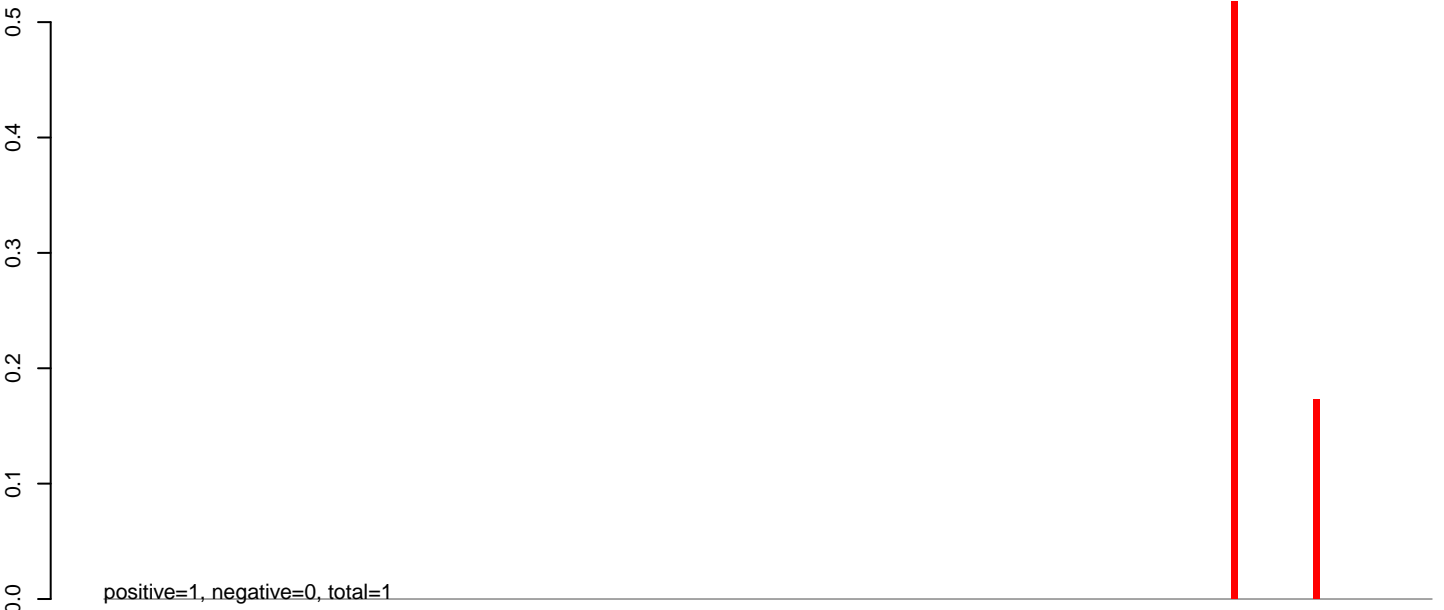
AeAeg_Aag2cells_CHIKV.24_35.rep



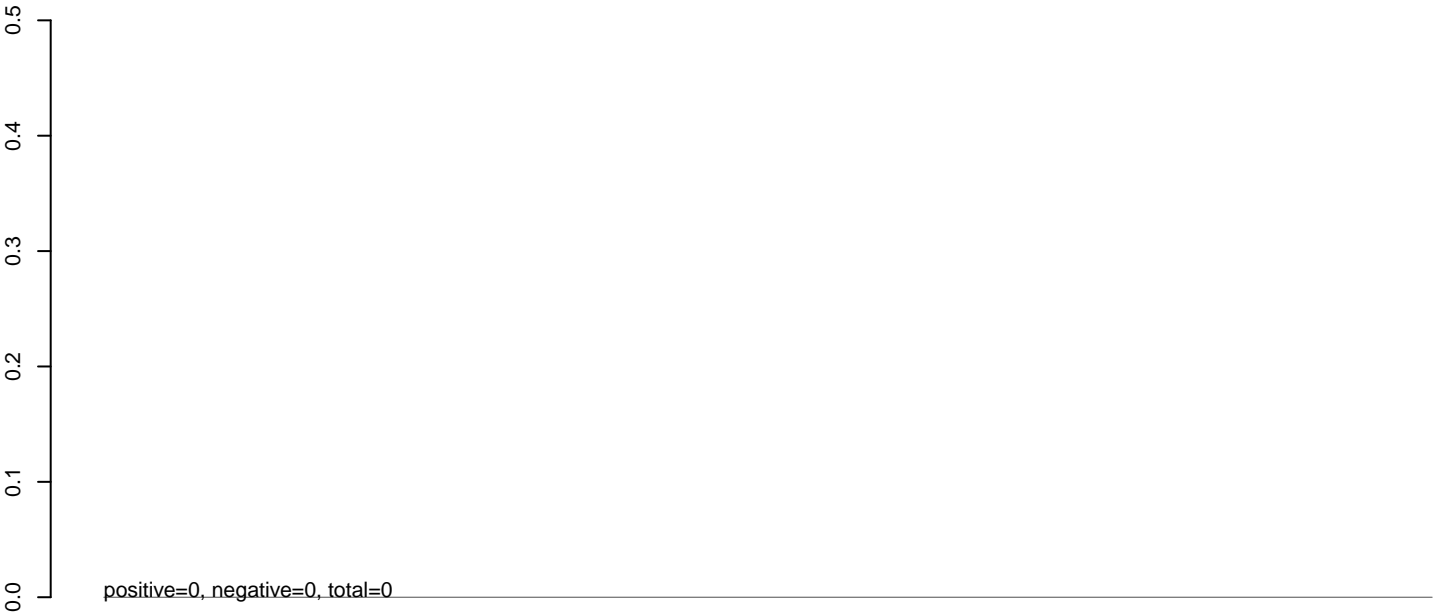
AeAeg_Aag2cells_CHIKV.rep



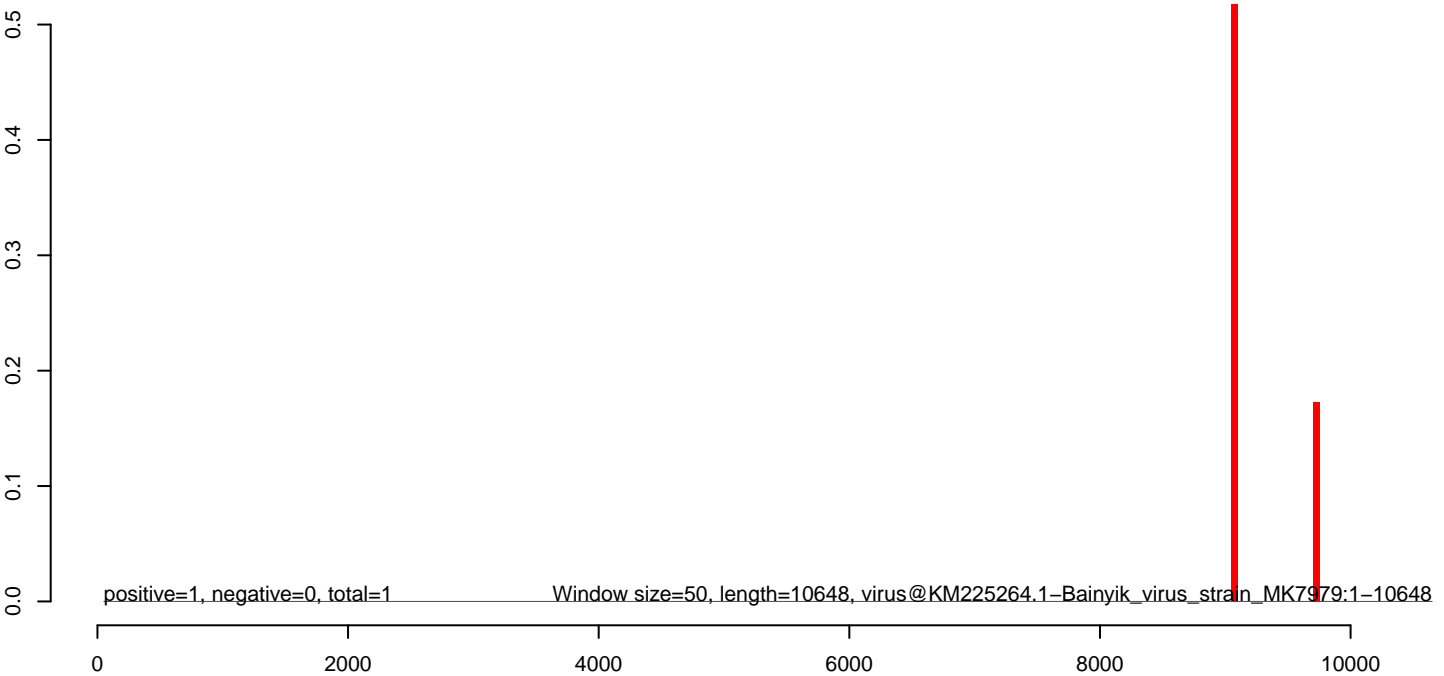
AeAeg_Aag2cells_CHIKV.18_23.rep



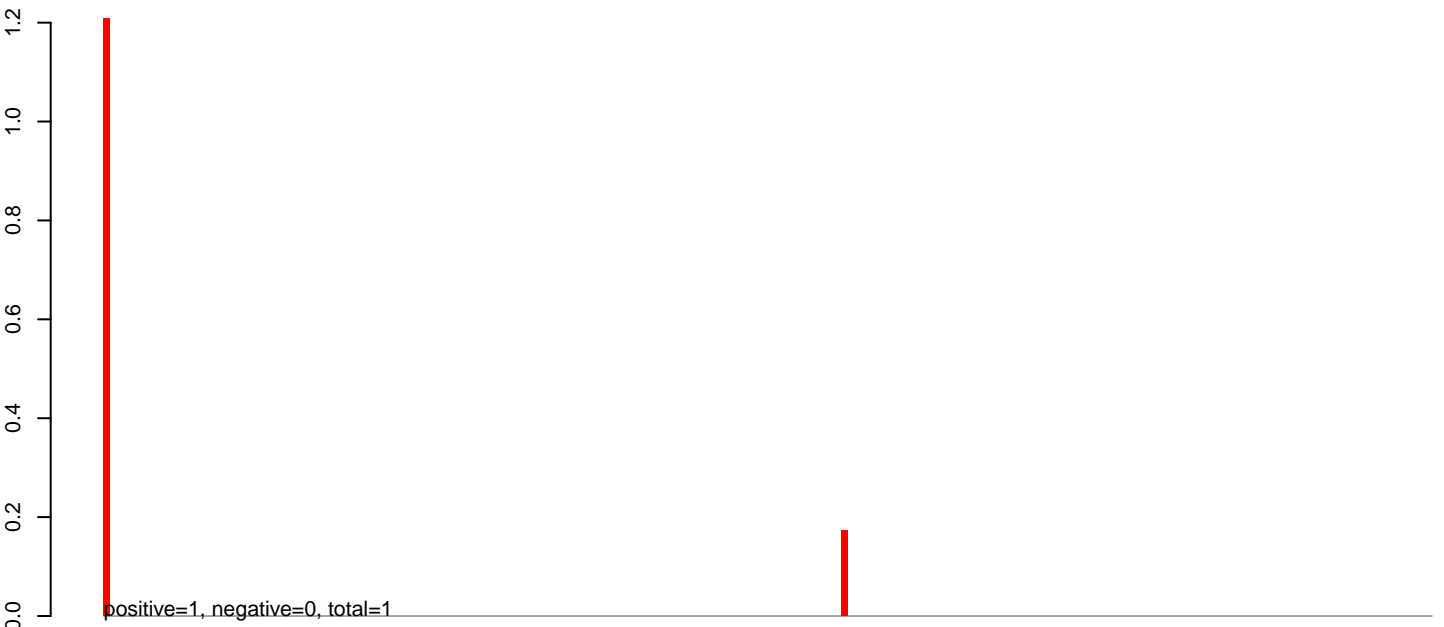
AeAeg_Aag2cells_CHIKV.24_35.rep



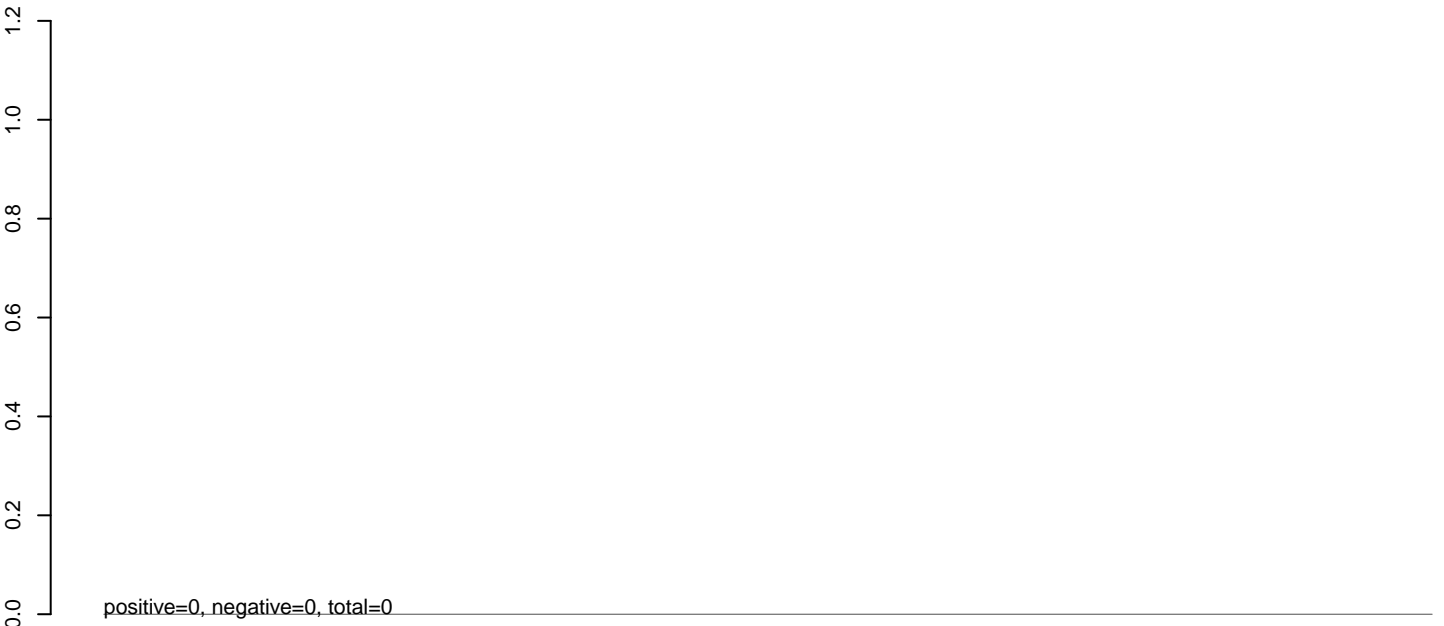
AeAeg_Aag2cells_CHIKV.rep



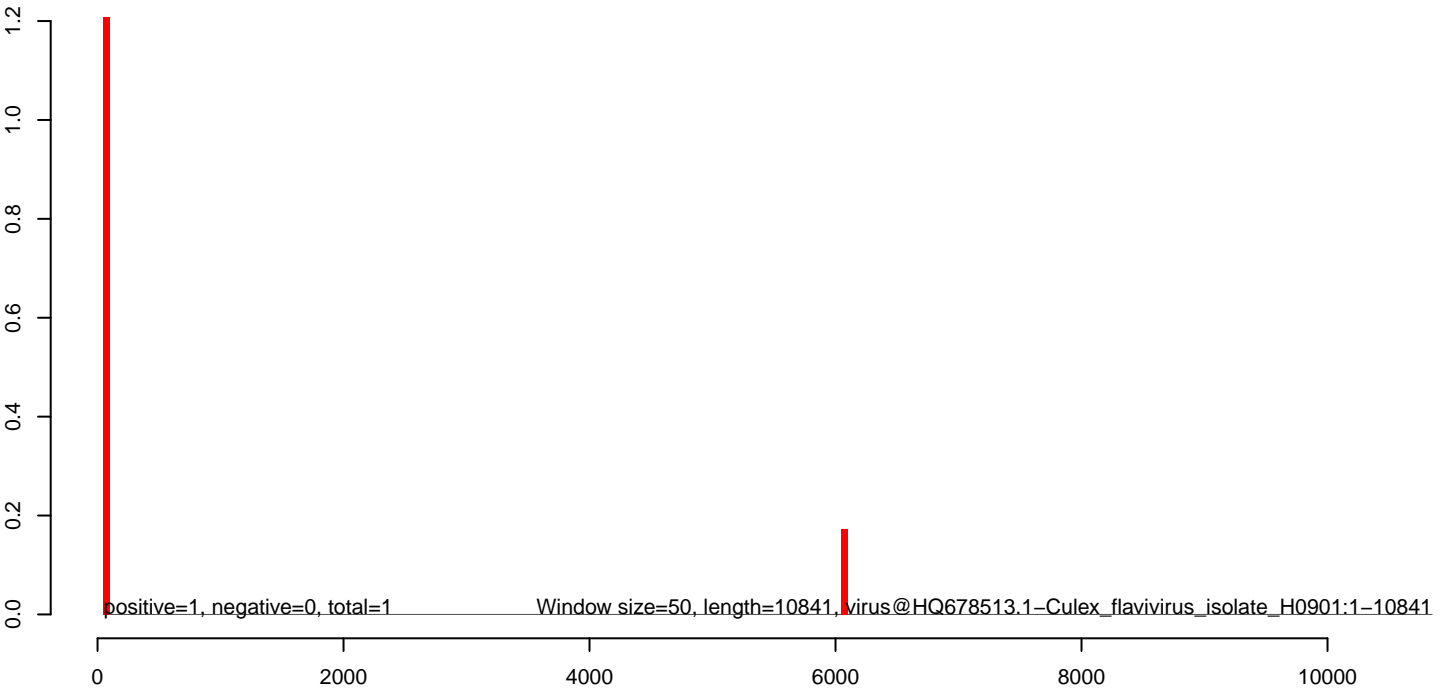
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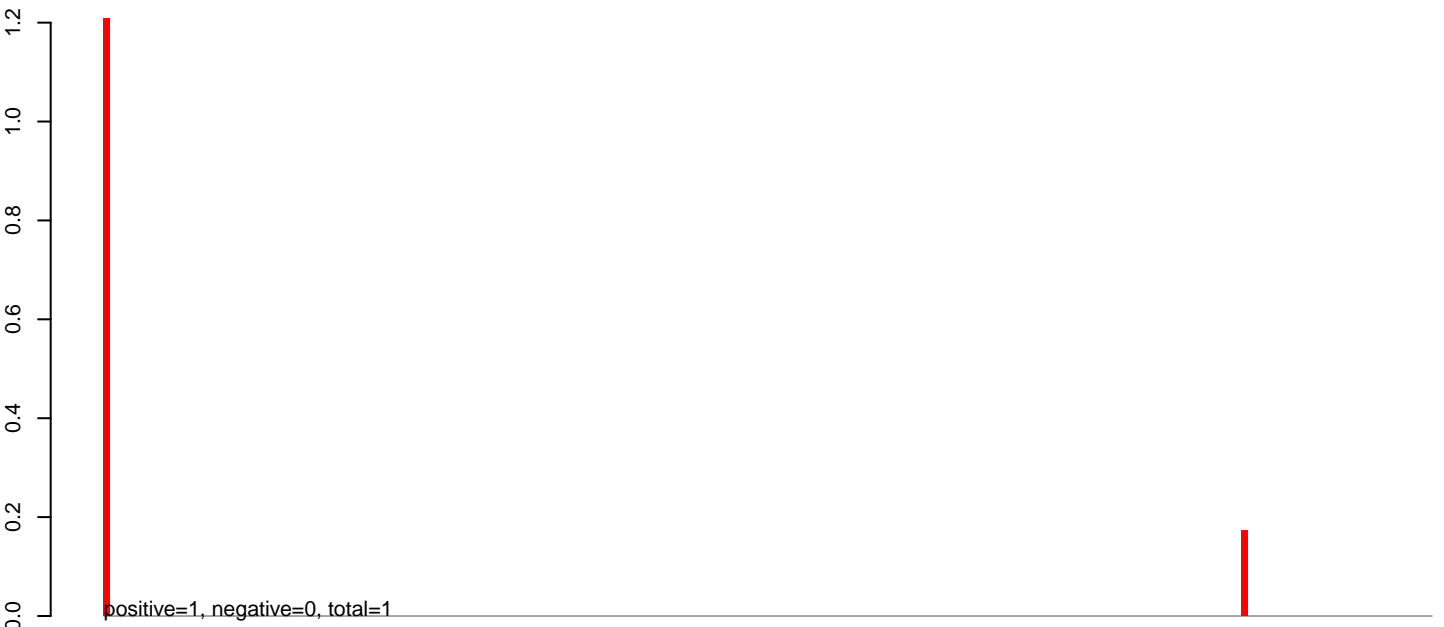
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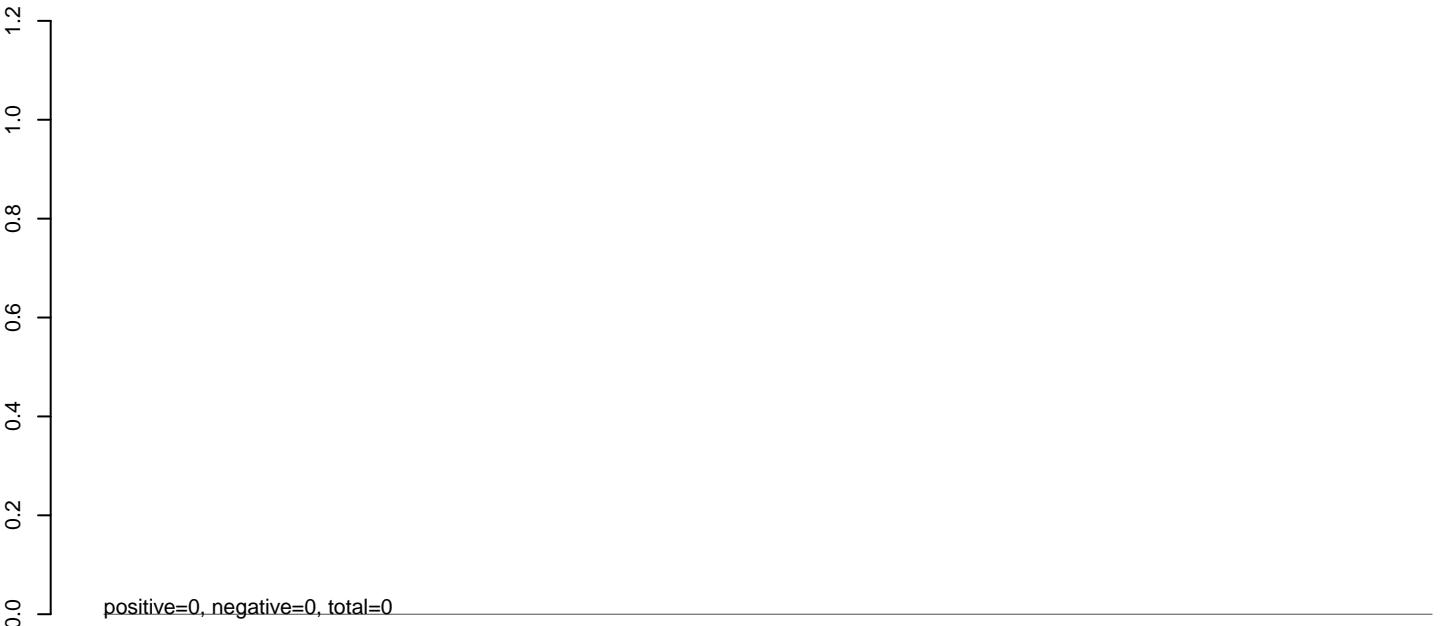
AeAeg_Aag2cells_CHIKV.rep



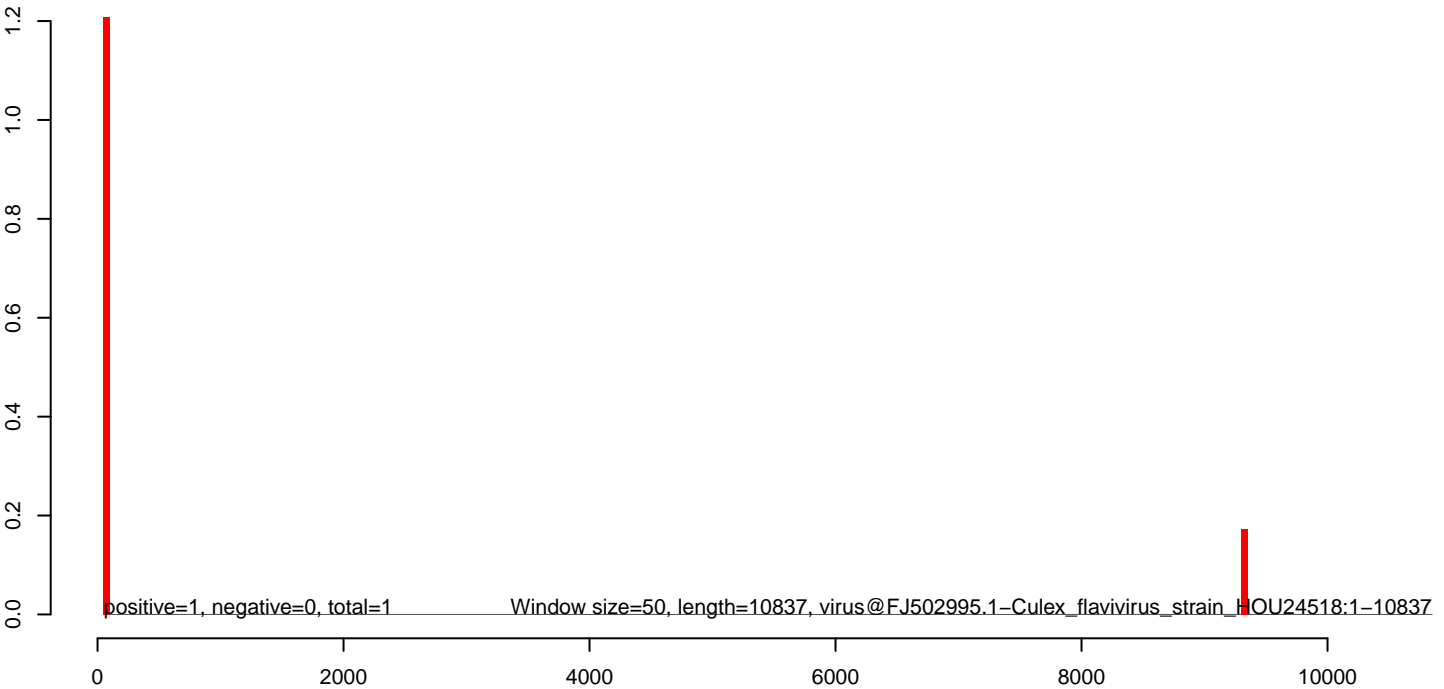
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep



AeAeg_Aag2cells_CHIKV.18_23.rep

positive=2, negative=0, total=2

AeAeg_Aag2cells_CHIKV.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2cells_CHIKV.rep

positive=2, negative=0, total=2

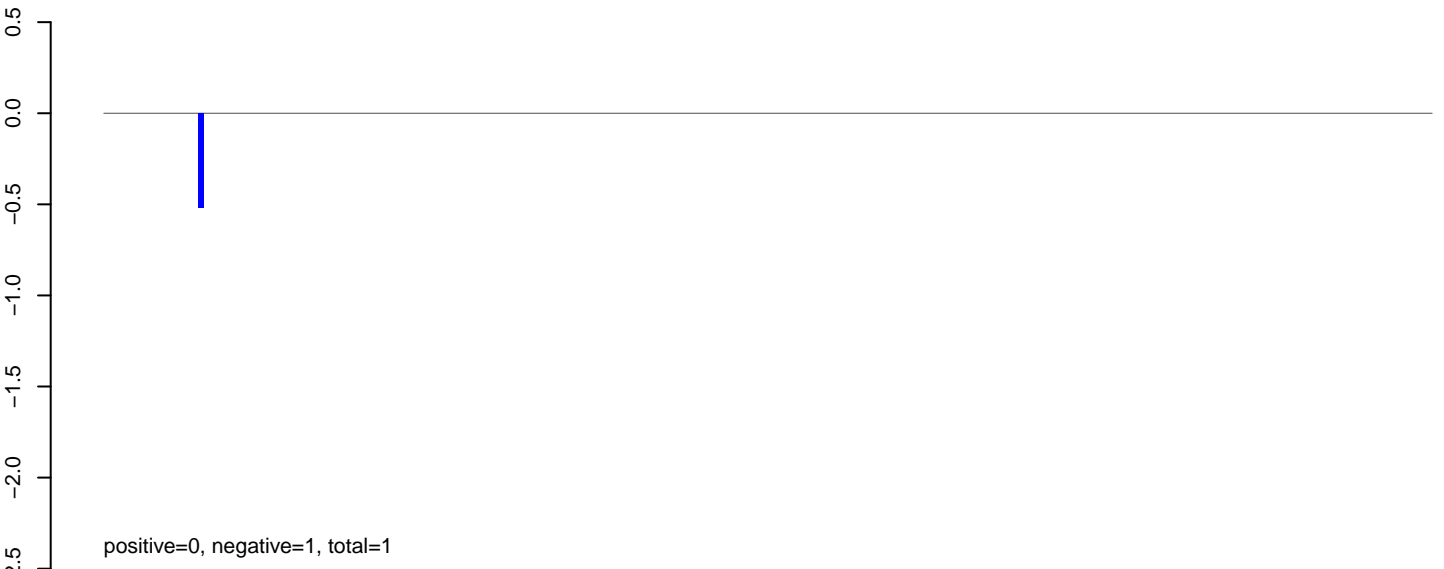
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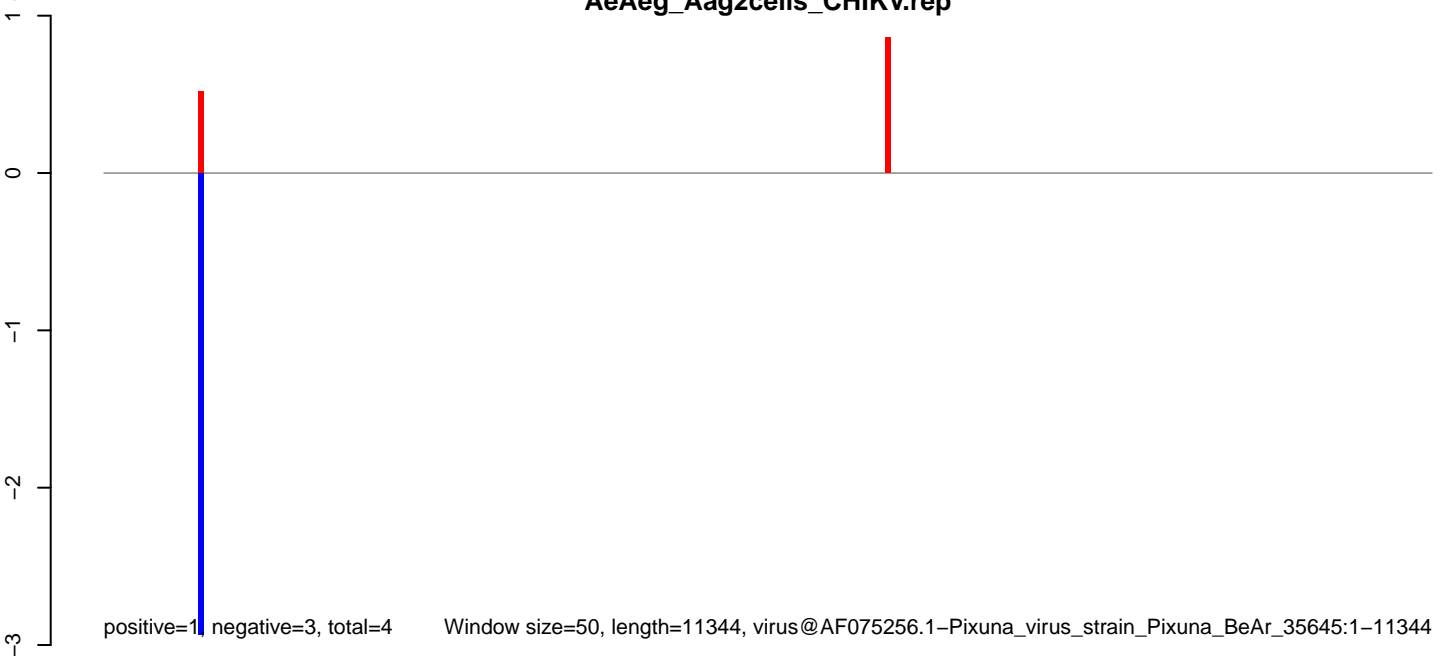
AeAeg_Aag2cells_CHIKV.18_23.rep



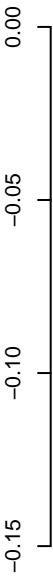
AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep

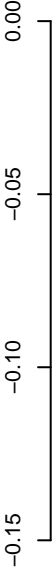


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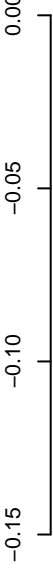
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AeAeg_Aag2cells_CHIKV.24_35.rep

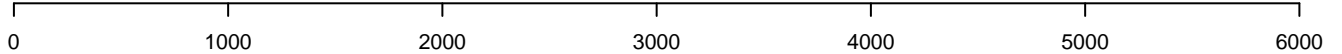


positive=0, negative=0, total=0

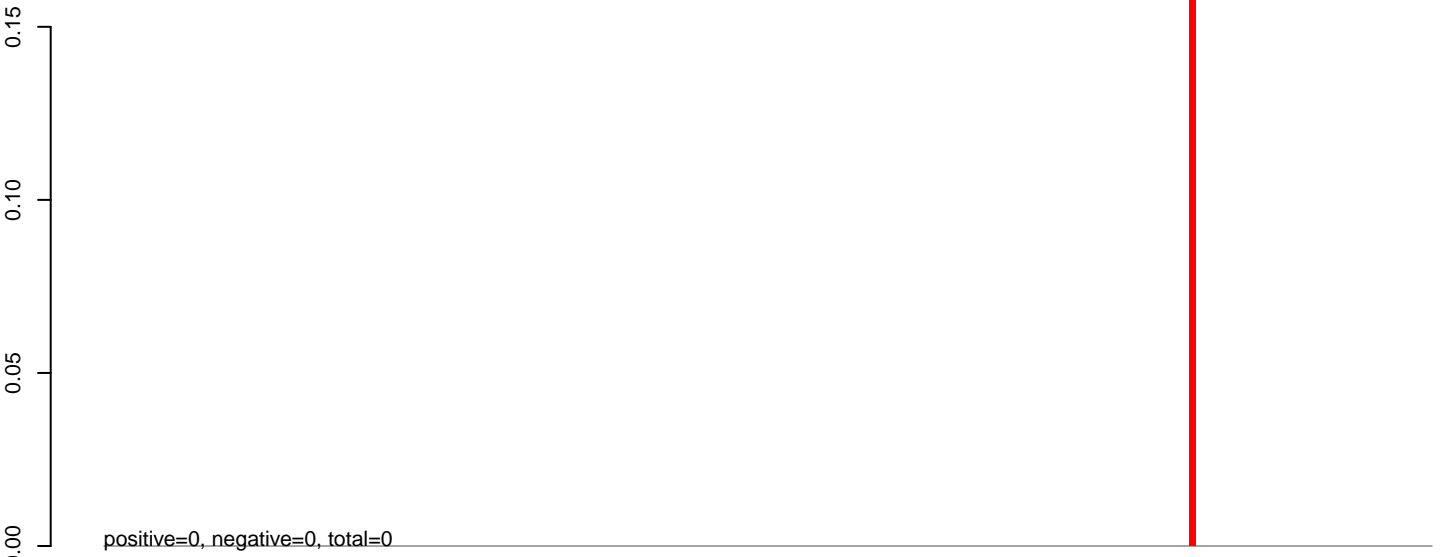
AeAeg_Aag2cells_CHIKV.rep



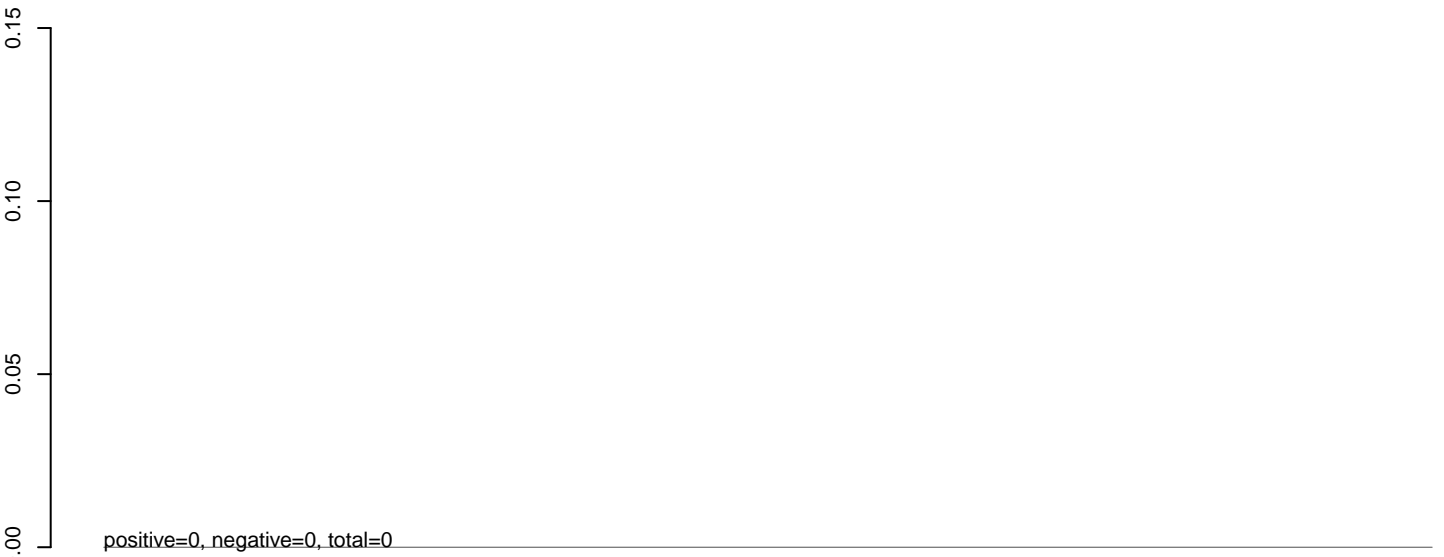
positive=0, negative=0, total=0, window size=50, length=6203, virus@NC_035674.1-Australian_Anopheles_totivirus_isolate_AATV_150840:1-6203



AeAeg_Aag2cells_CHIKV.18_23.rep



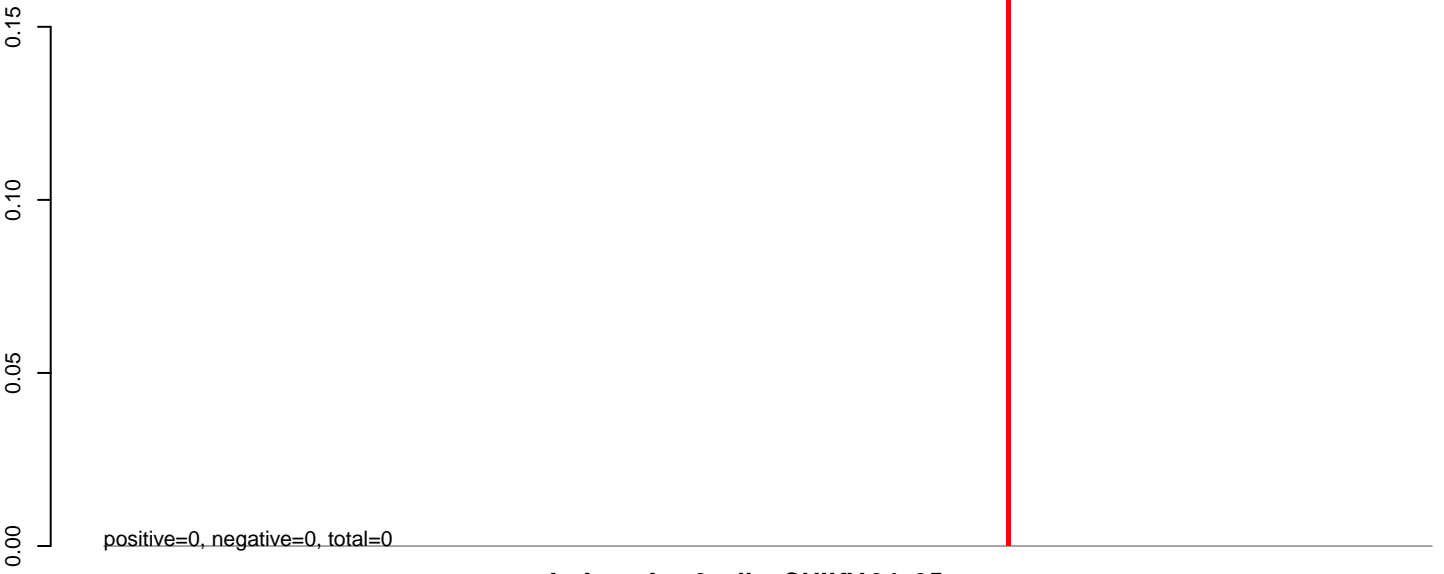
AeAeg_Aag2cells_CHIKV.24_35.rep



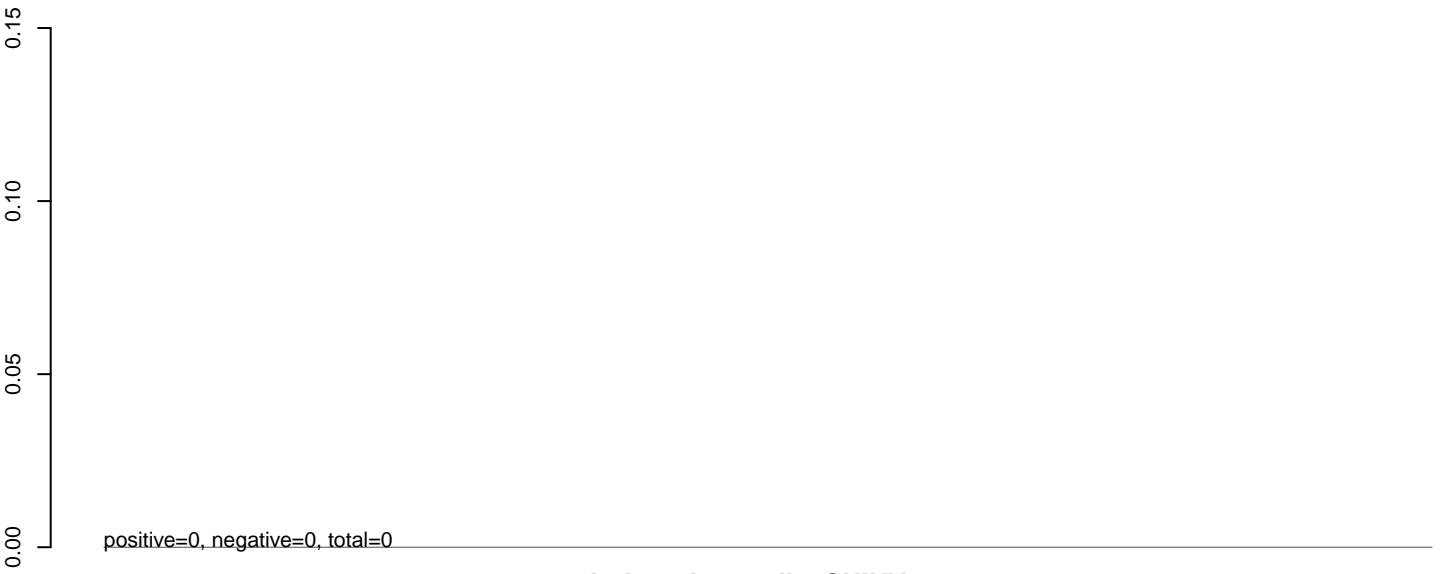
AeAeg_Aag2cells_CHIKV.rep



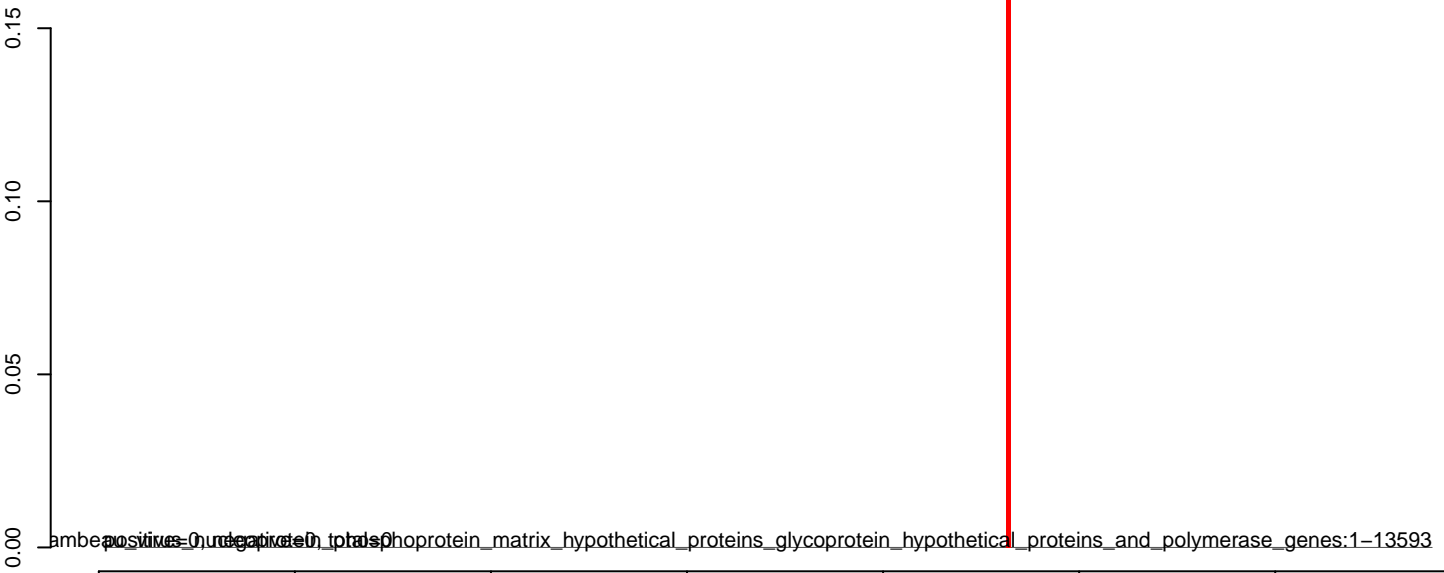
AeAeg_Aag2cells_CHIKV.18_23.rep



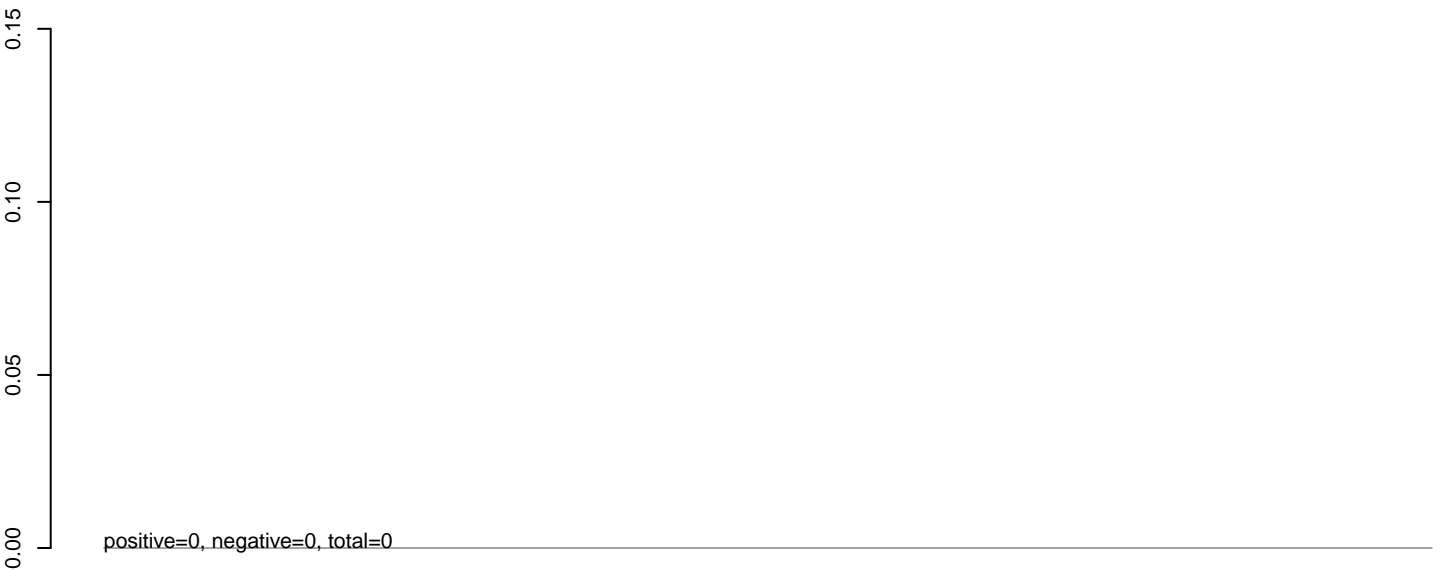
AeAeg_Aag2cells_CHIKV.24_35.rep



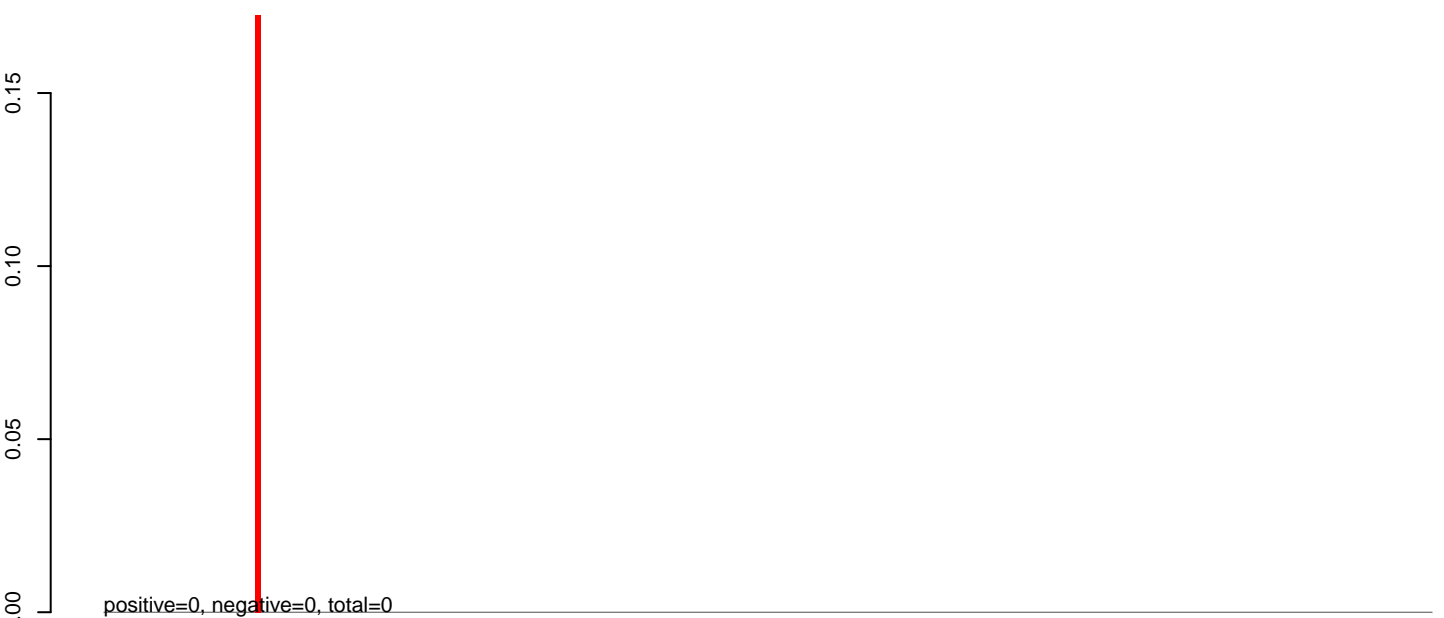
AeAeg_Aag2cells_CHIKV.rep



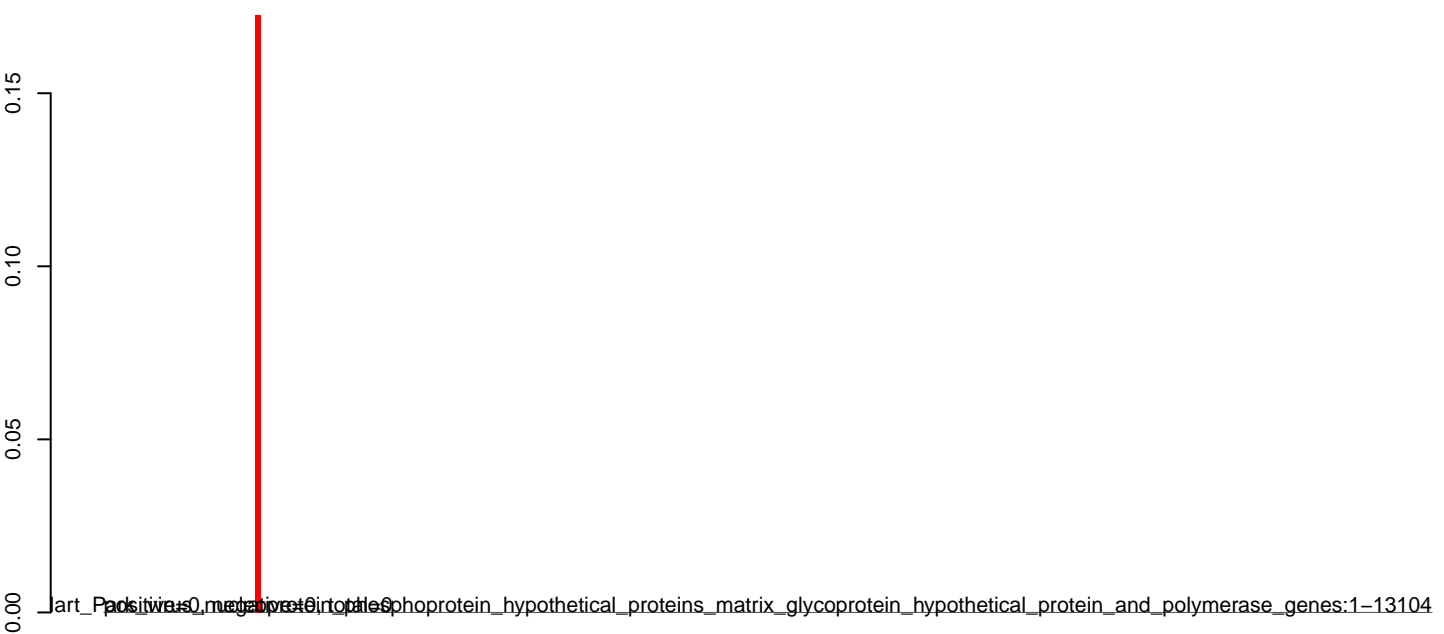
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

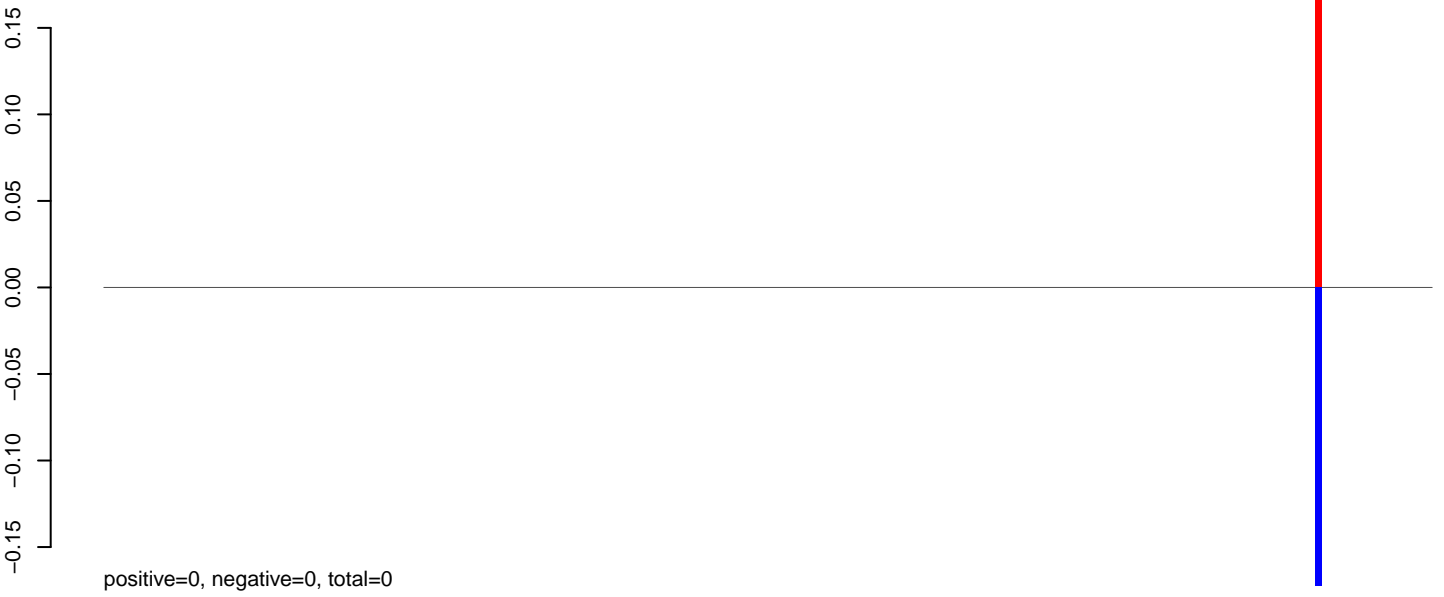


AeAeg_Aag2cells_CHIKV.rep

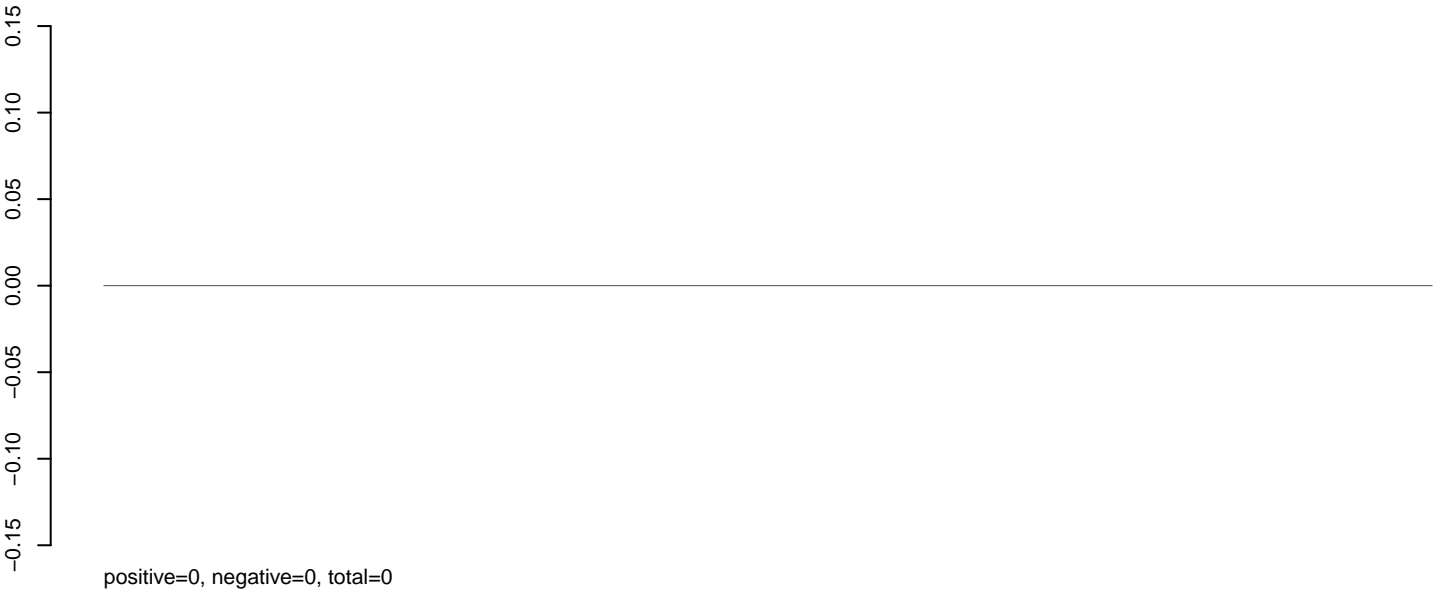


0 2000 4000 6000 8000 10000 12000
part_positive, negative, total, phosphoprotein_hypothetical_proteins_matrix_glycoprotein_hypothetical_protein_and_polymerase_genes:1-13104

AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



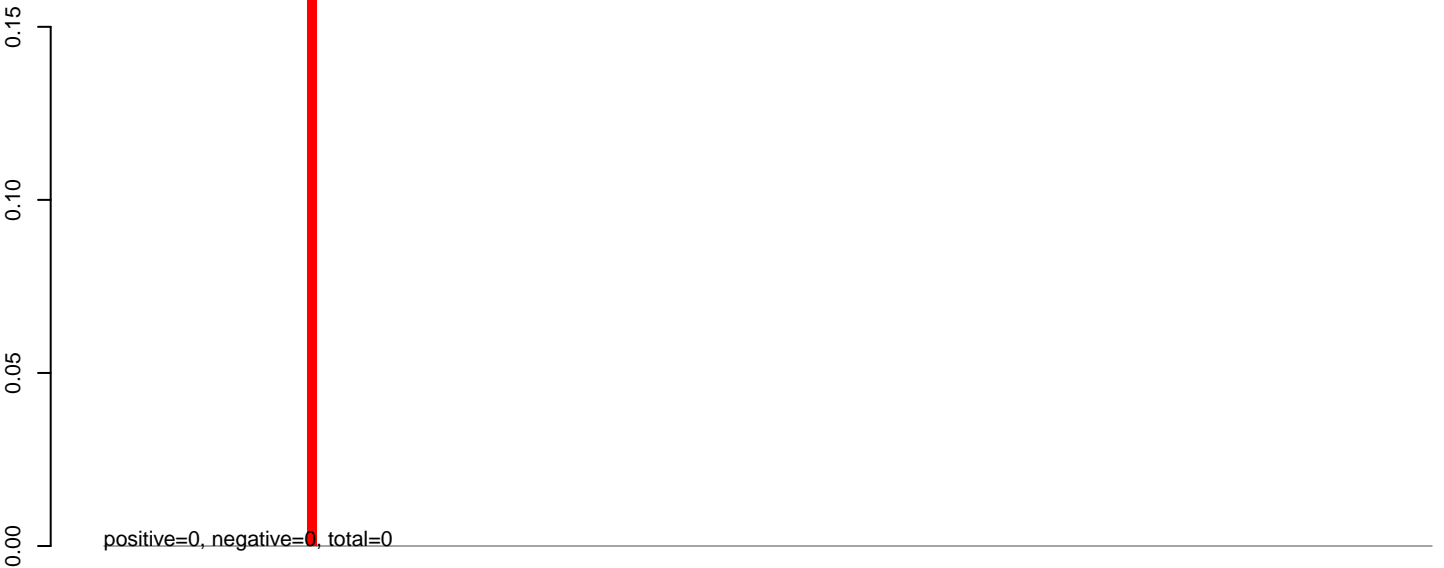
AeAeg_Aag2cells_CHIKV.rep



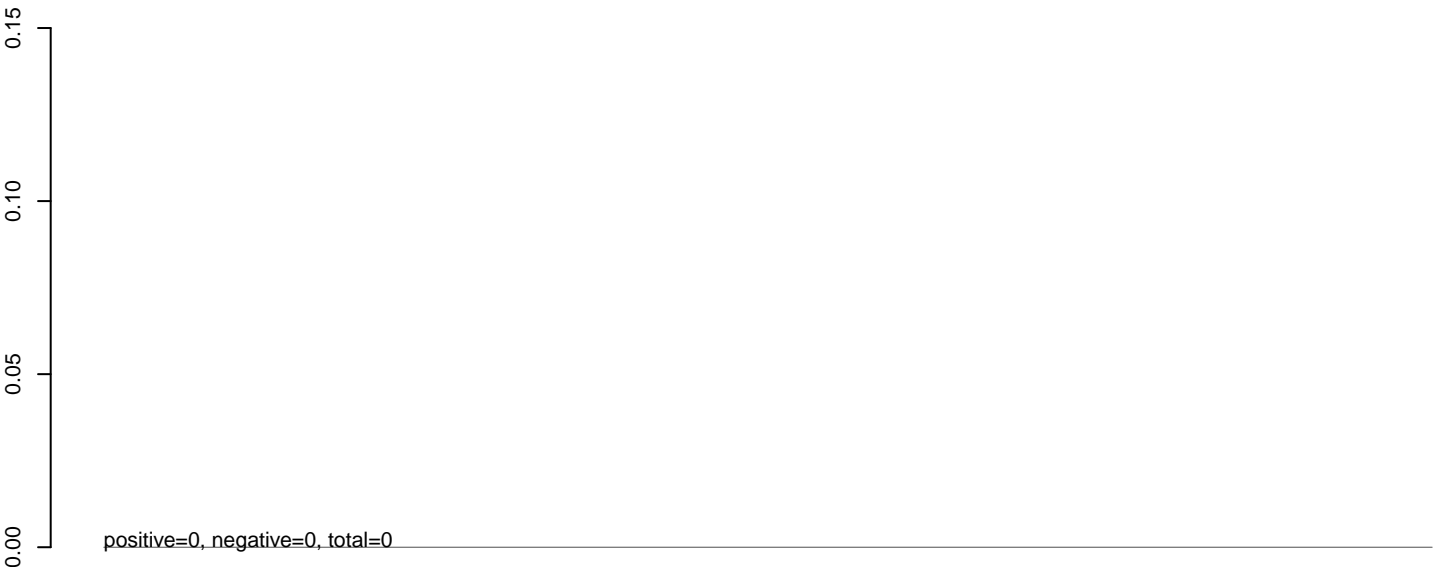
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0 2000 4000 6000 8000 10000

AeAeg_Aag2cells_CHIKV.18_23.rep



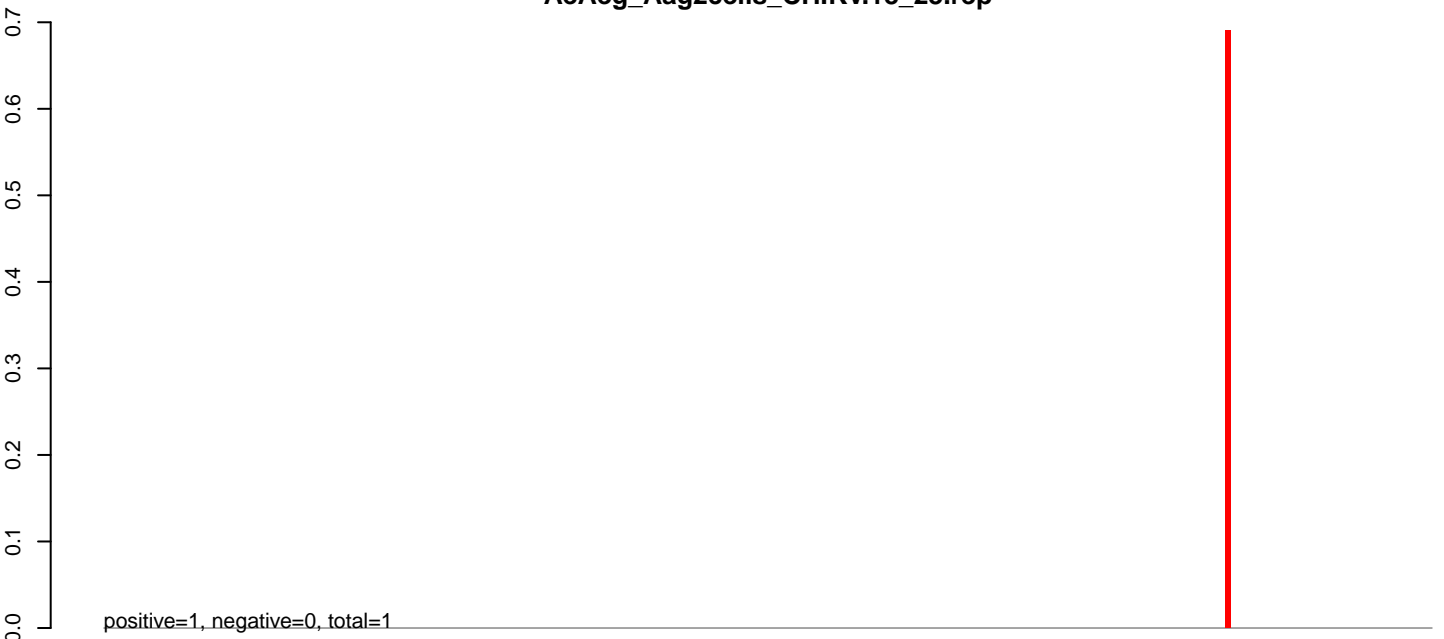
AeAeg_Aag2cells_CHIKV.24_35.rep



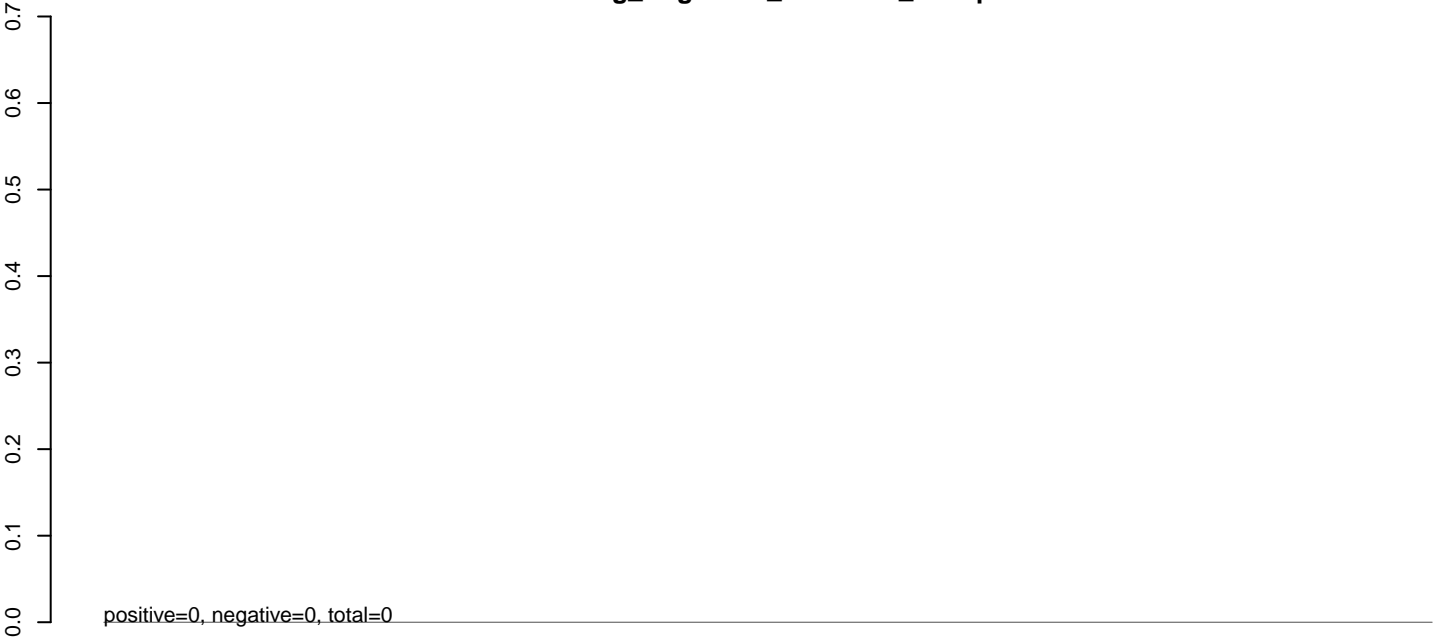
AeAeg_Aag2cells_CHIKV.rep



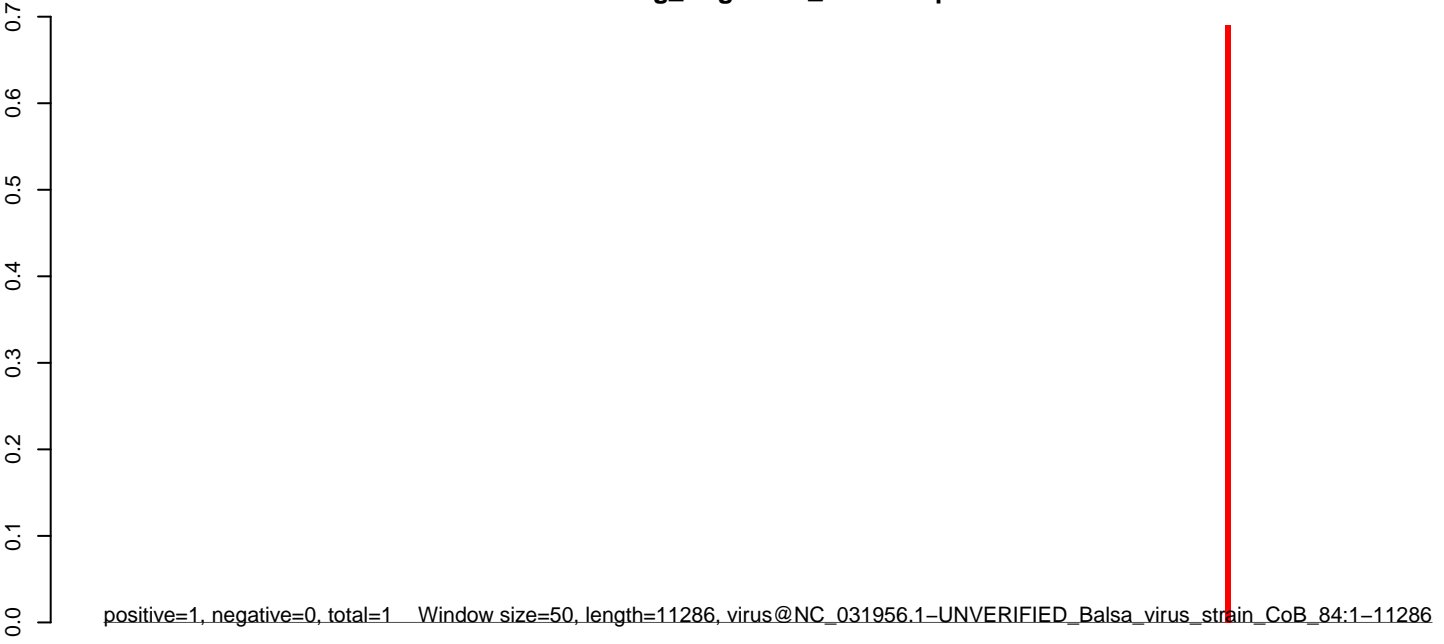
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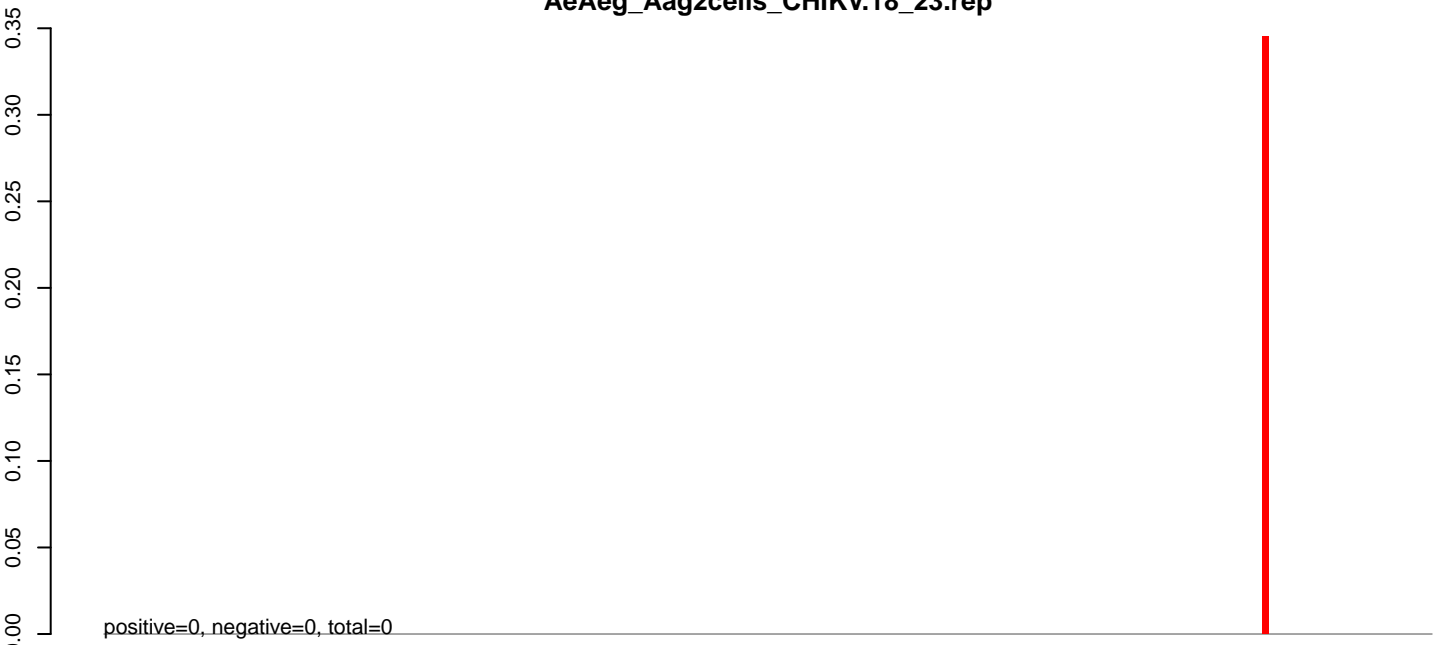
AeAeg_Aag2cells_CHIKV.24_35.rep



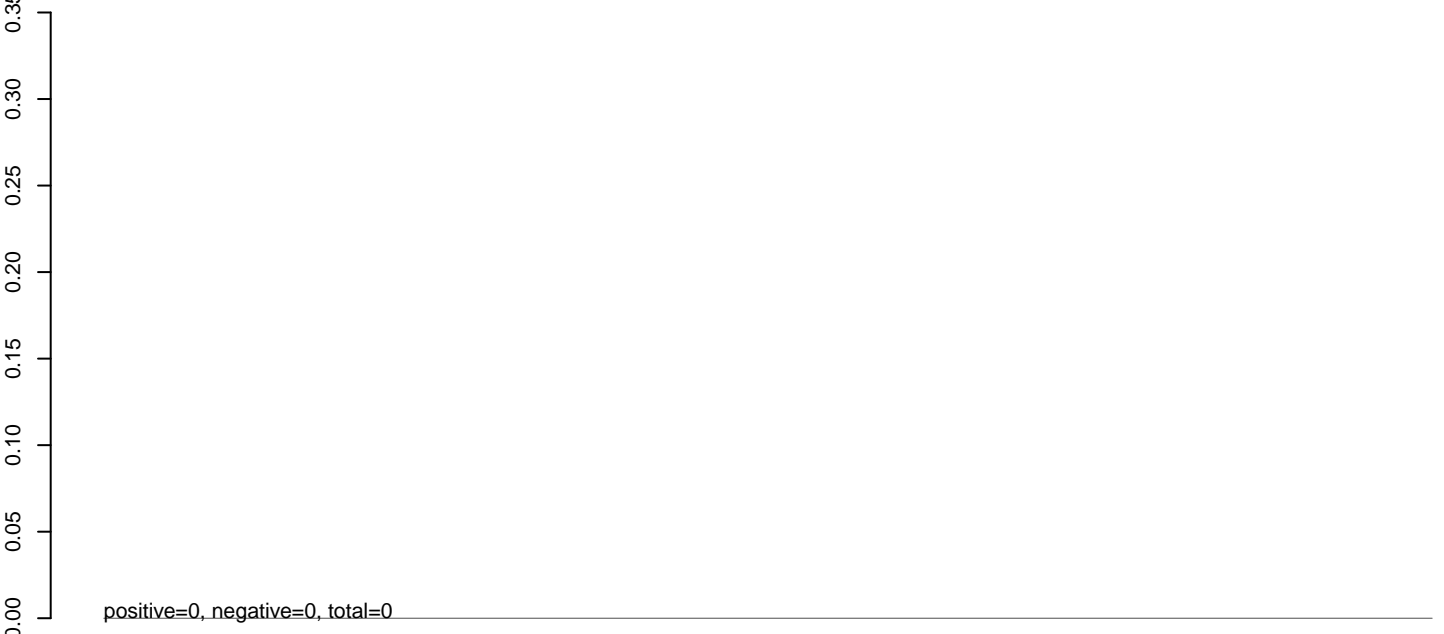
AeAeg_Aag2cells_CHIKV.rep



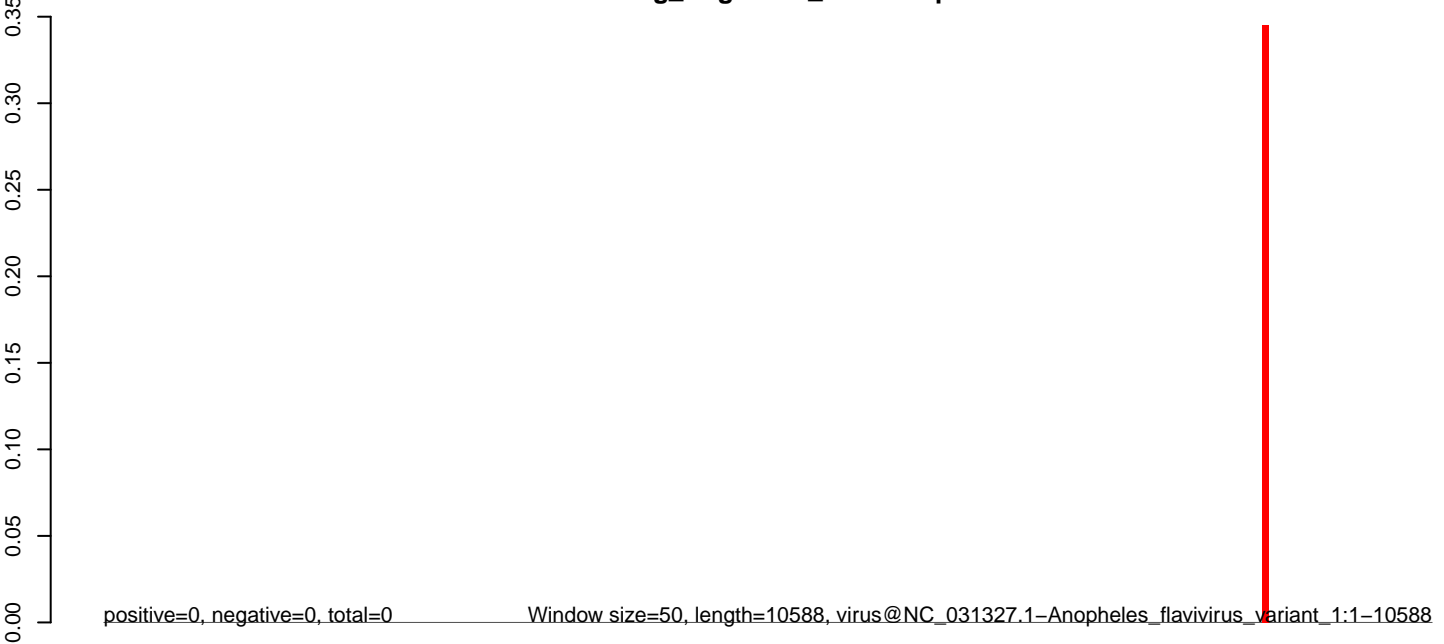
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

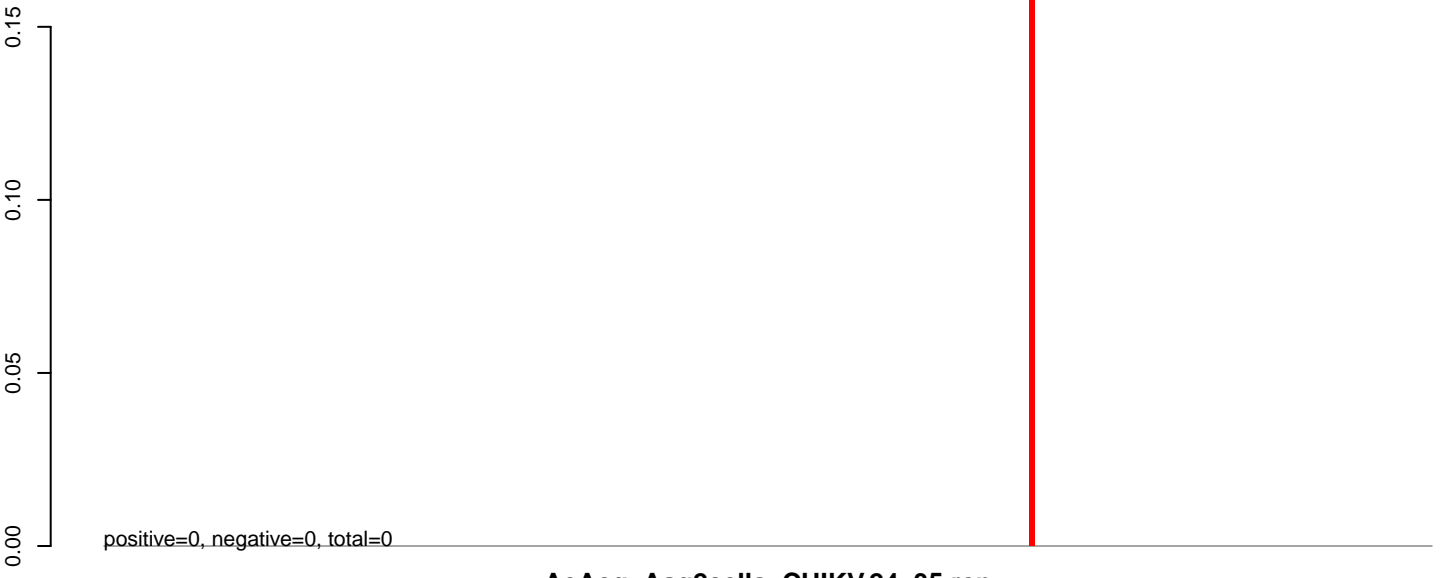


AeAeg_Aag2cells_CHIKV.rep

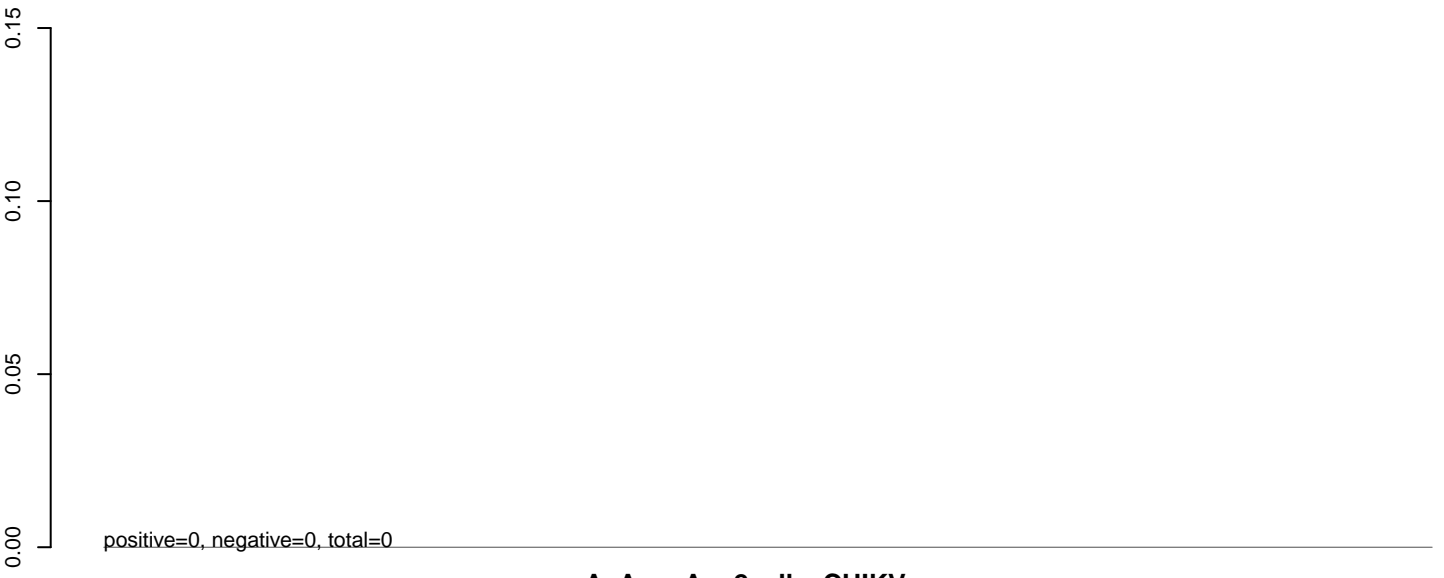


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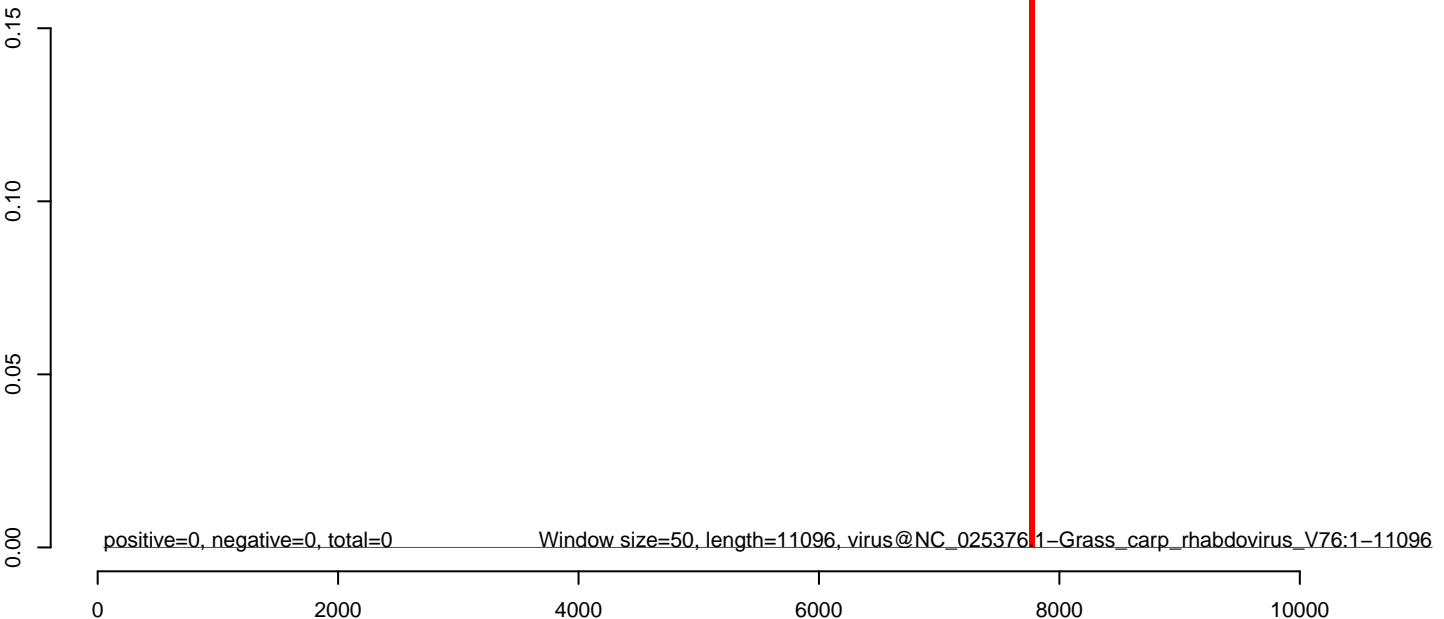
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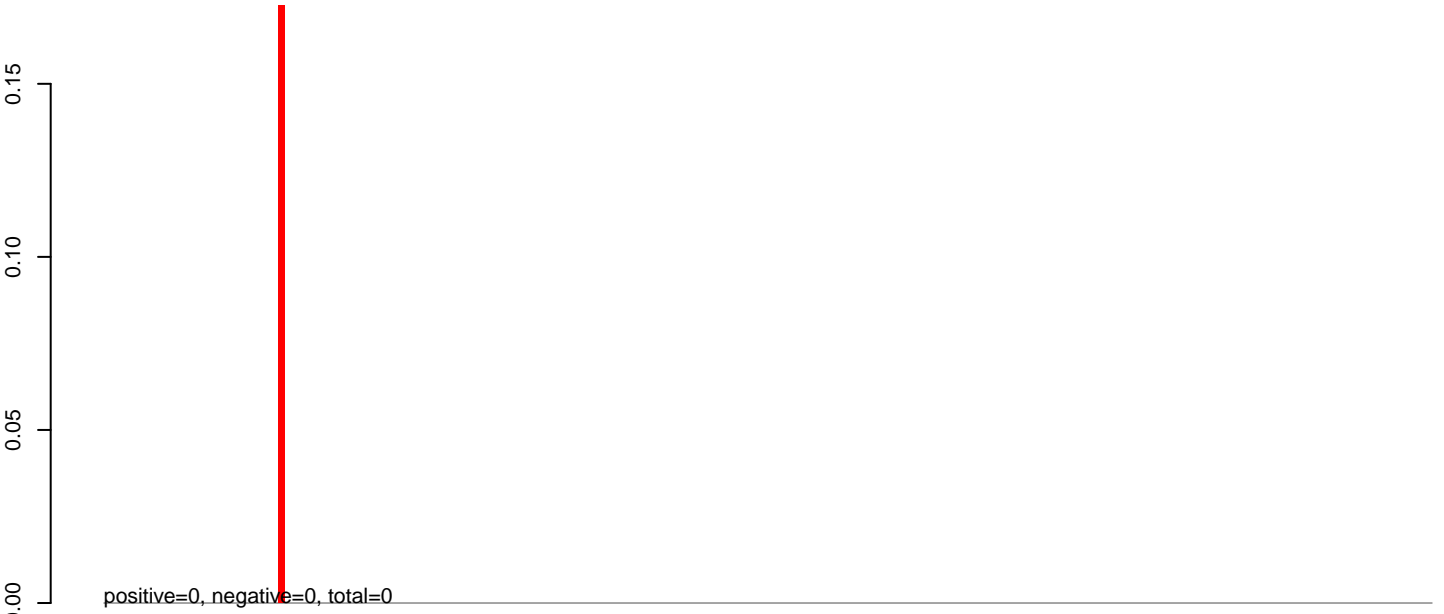
AeAeg_Aag2cells_CHIKV.24_35.rep



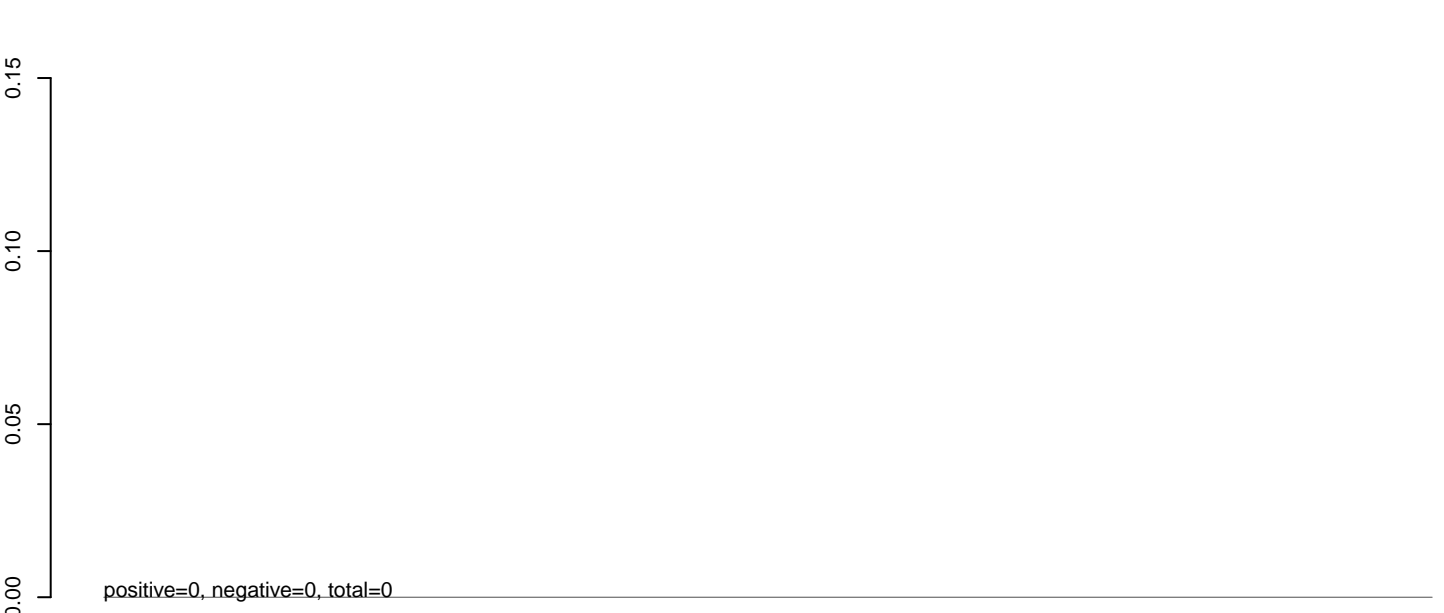
AeAeg_Aag2cells_CHIKV.rep



AeAeg_Aag2cells_CHIKV.18_23.rep



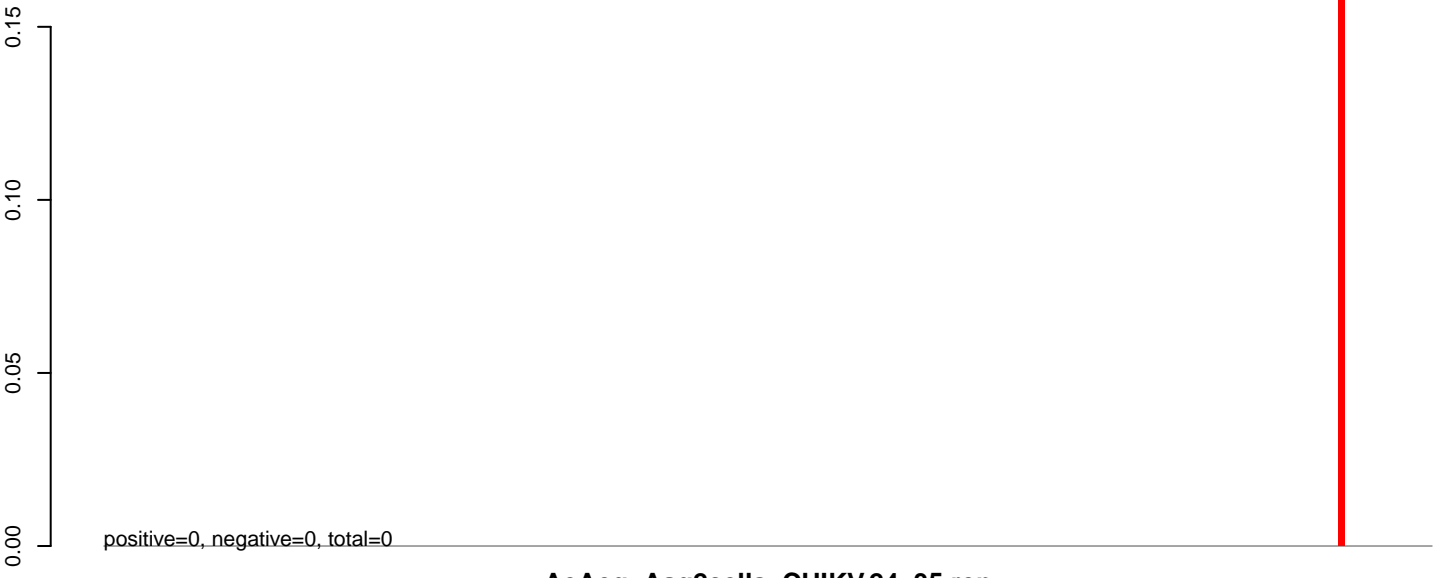
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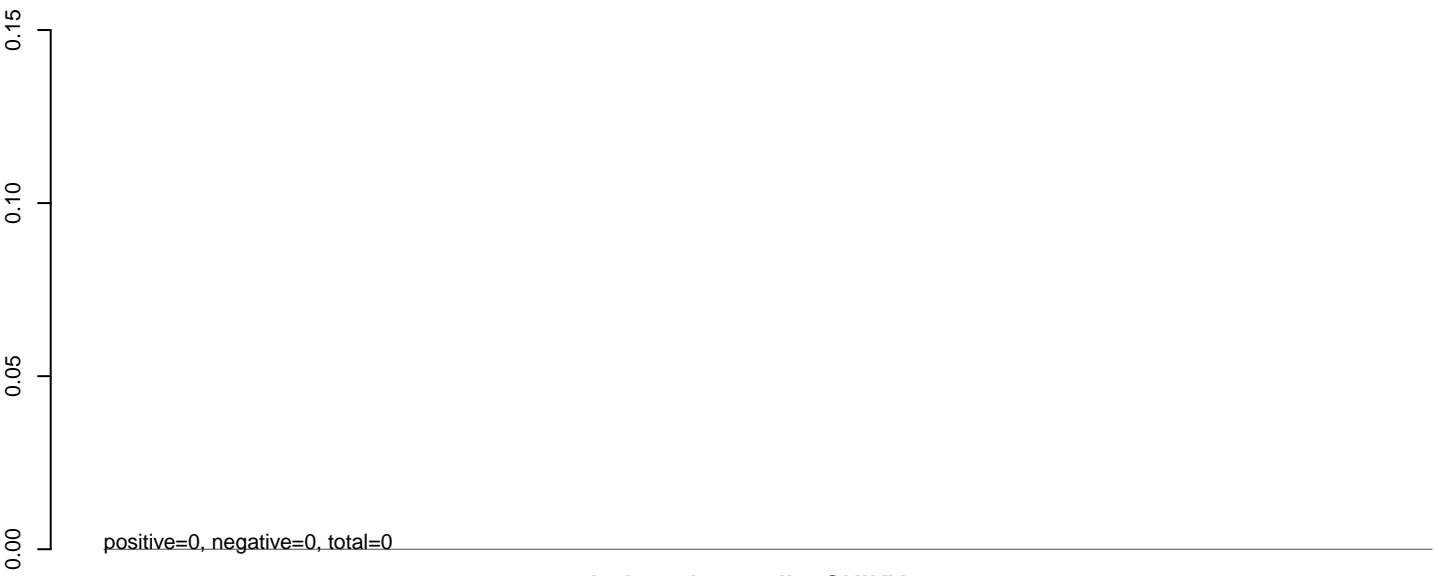
AeAeg_Aag2cells_CHIKV.rep



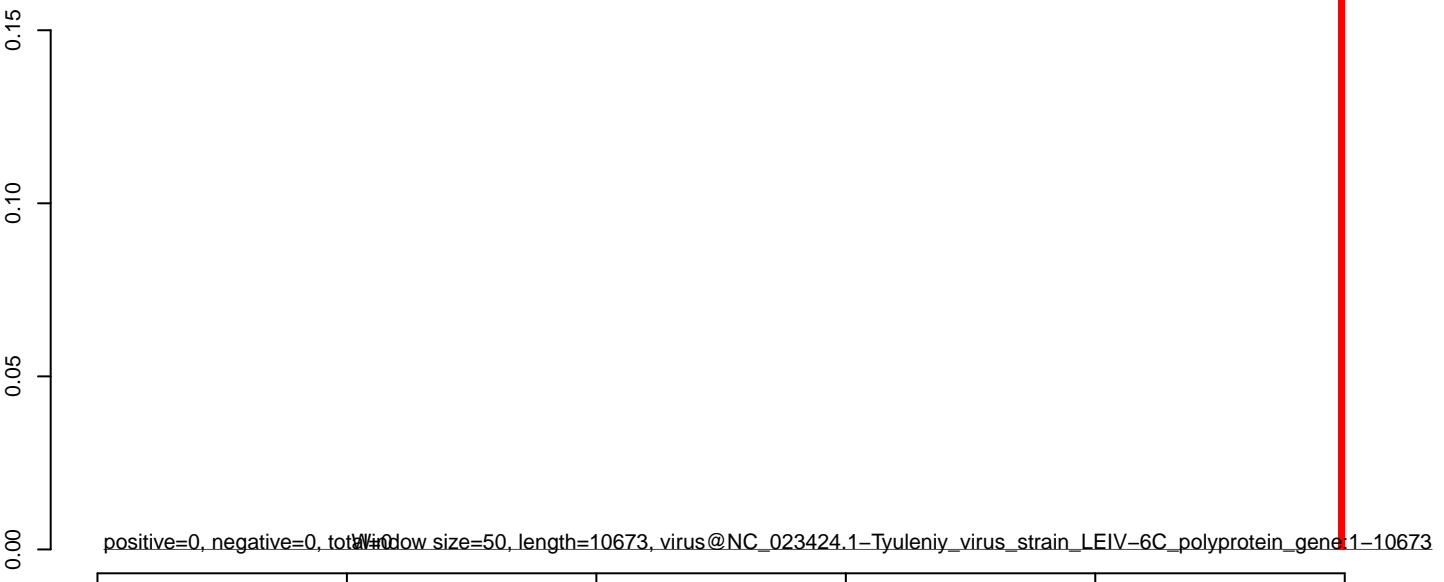
AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep



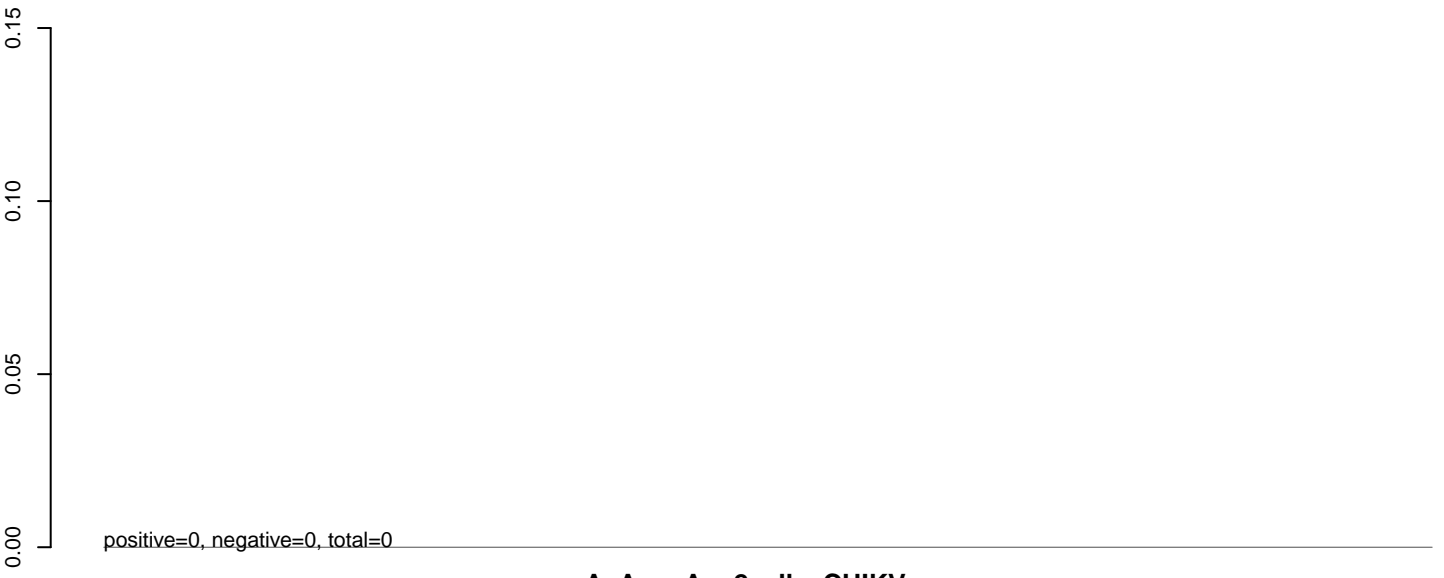
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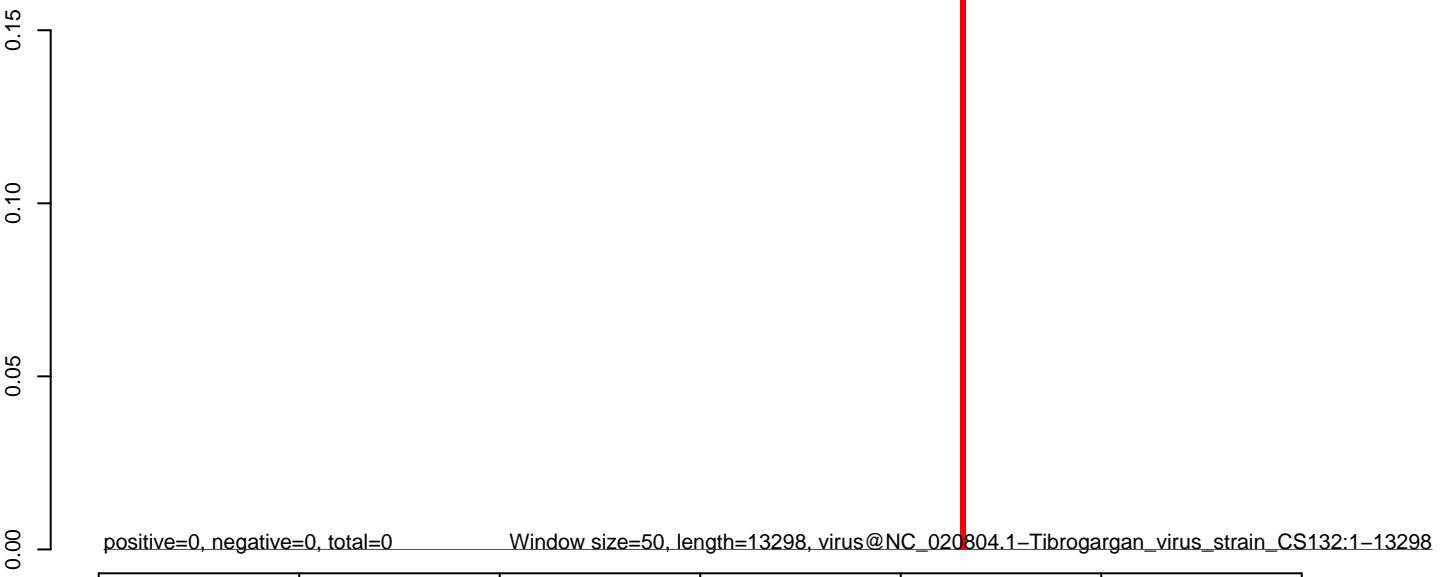
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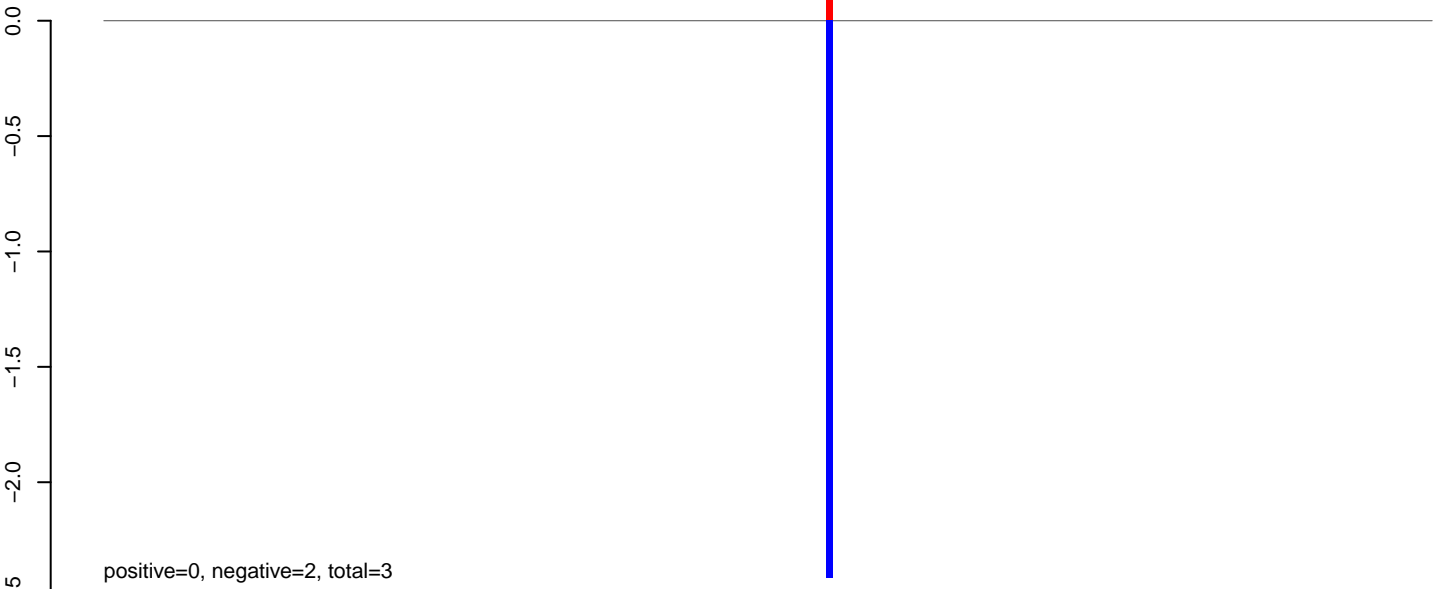
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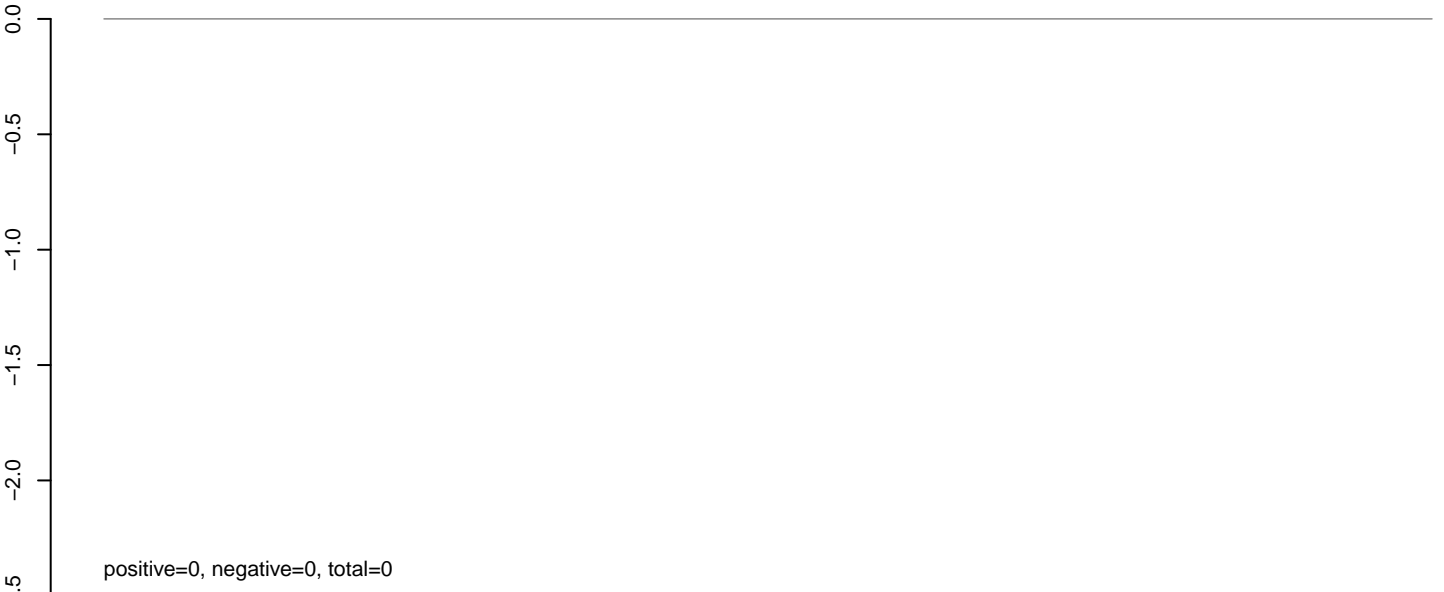
AeAeg_Aag2cells_CHIKV.rep



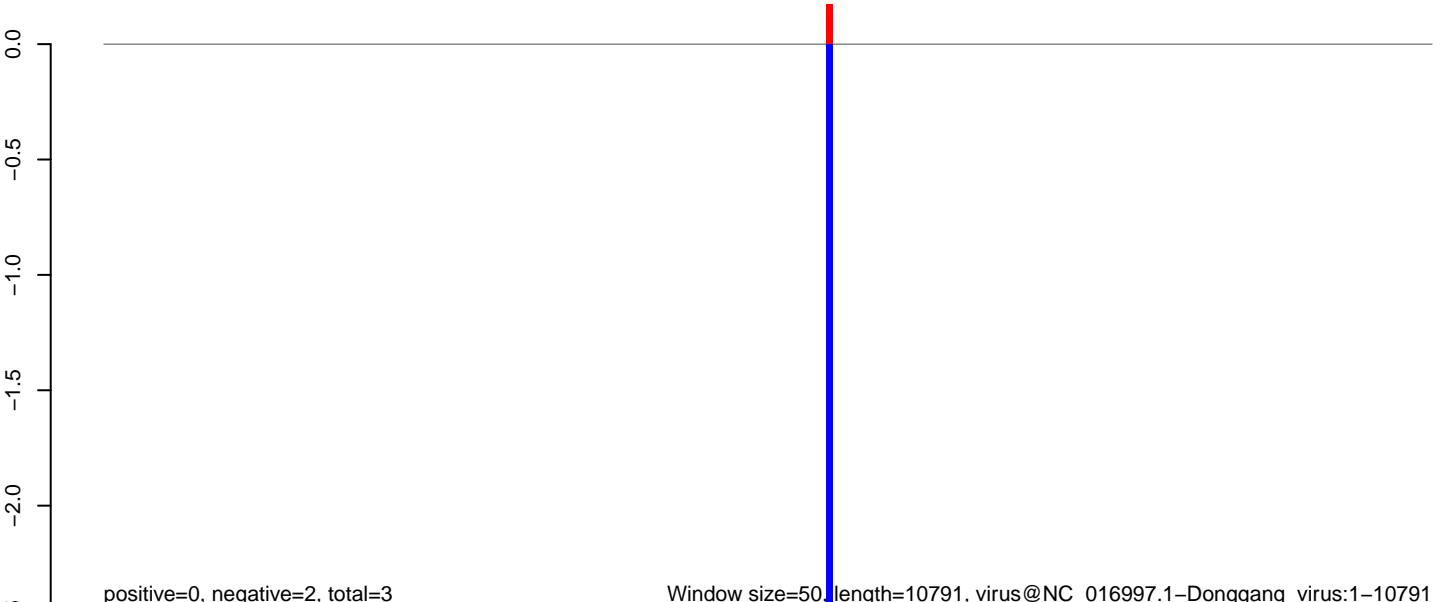
AeAeg_Aag2cells_CHIKV.18_23.rep



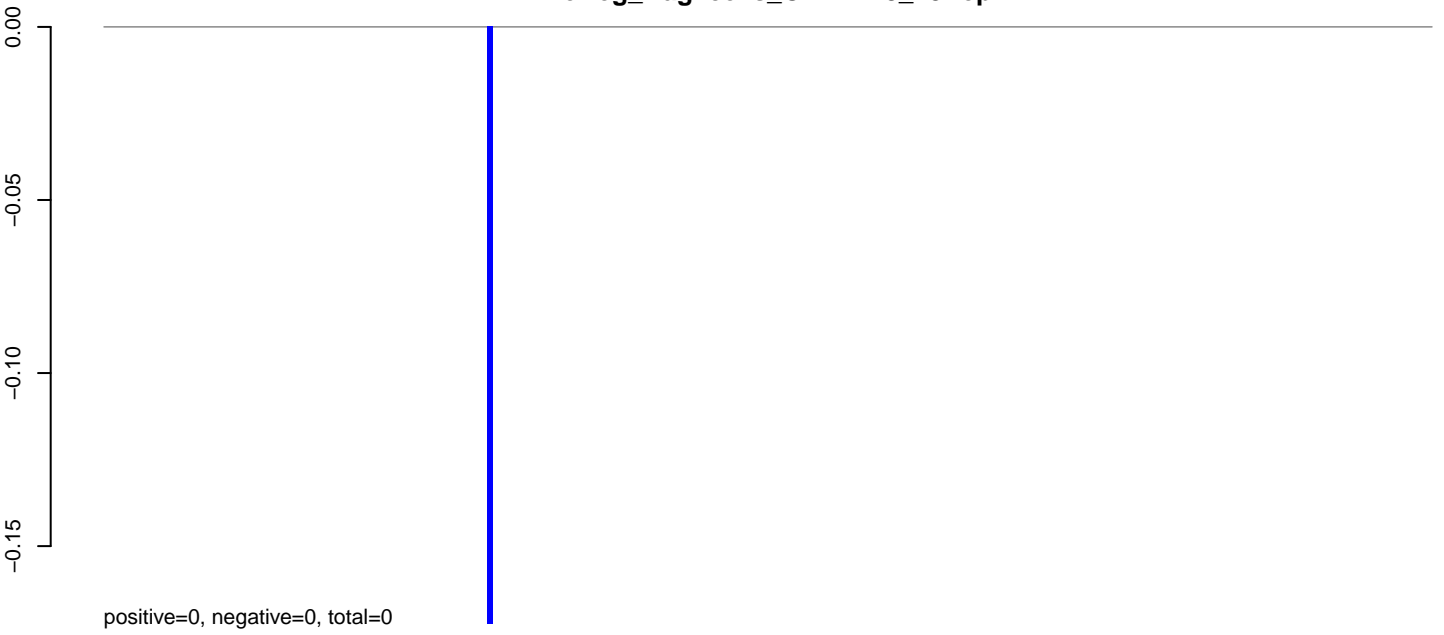
AeAeg_Aag2cells_CHIKV.24_35.rep



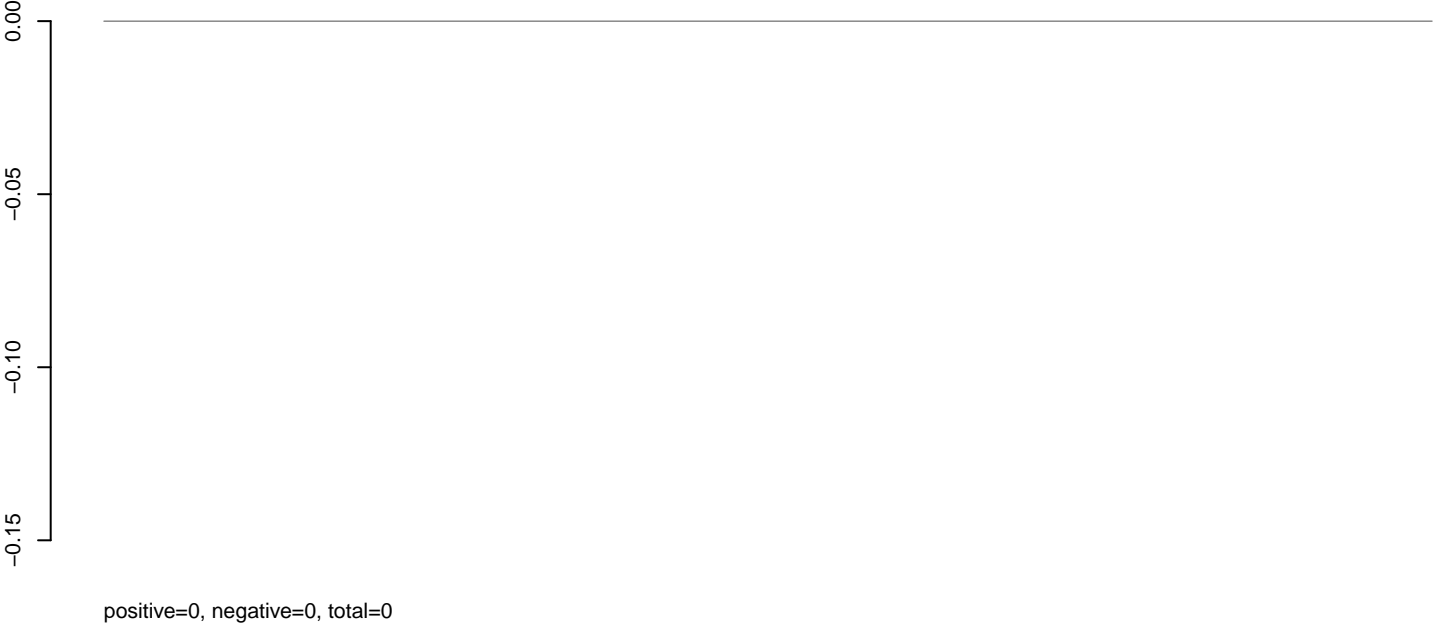
AeAeg_Aag2cells_CHIKV.rep



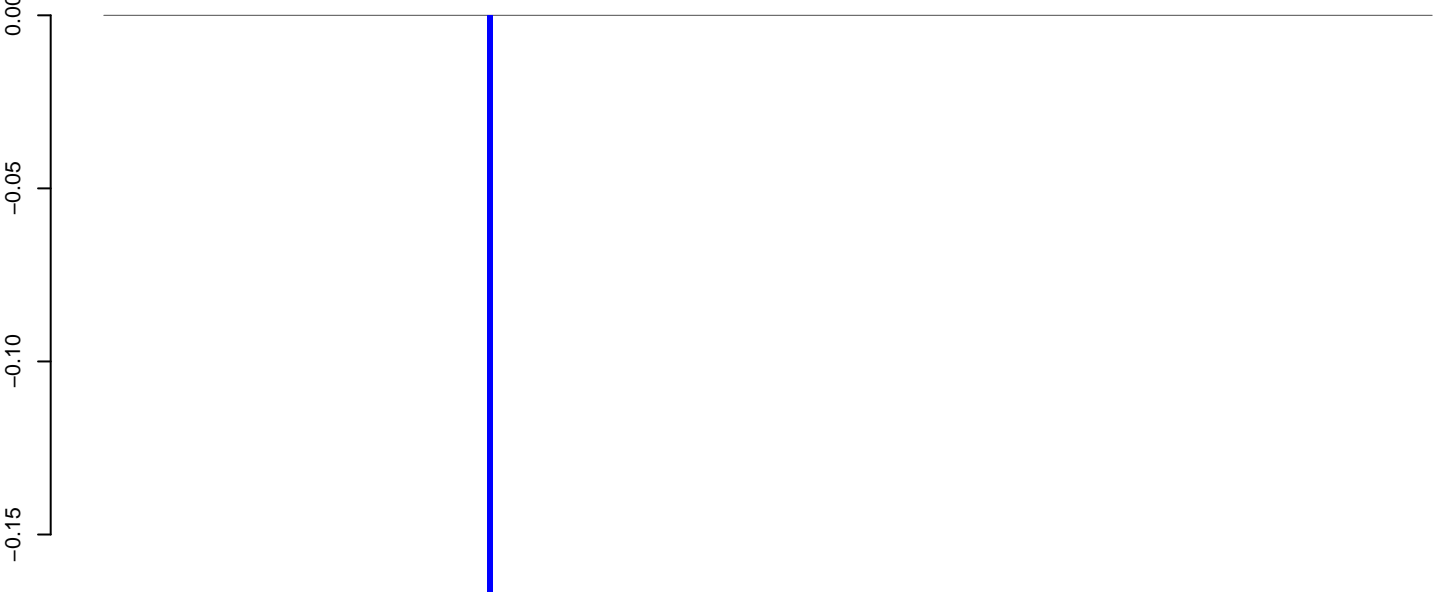
AeAeg_Aag2cells_CHIKV.18_23.rep



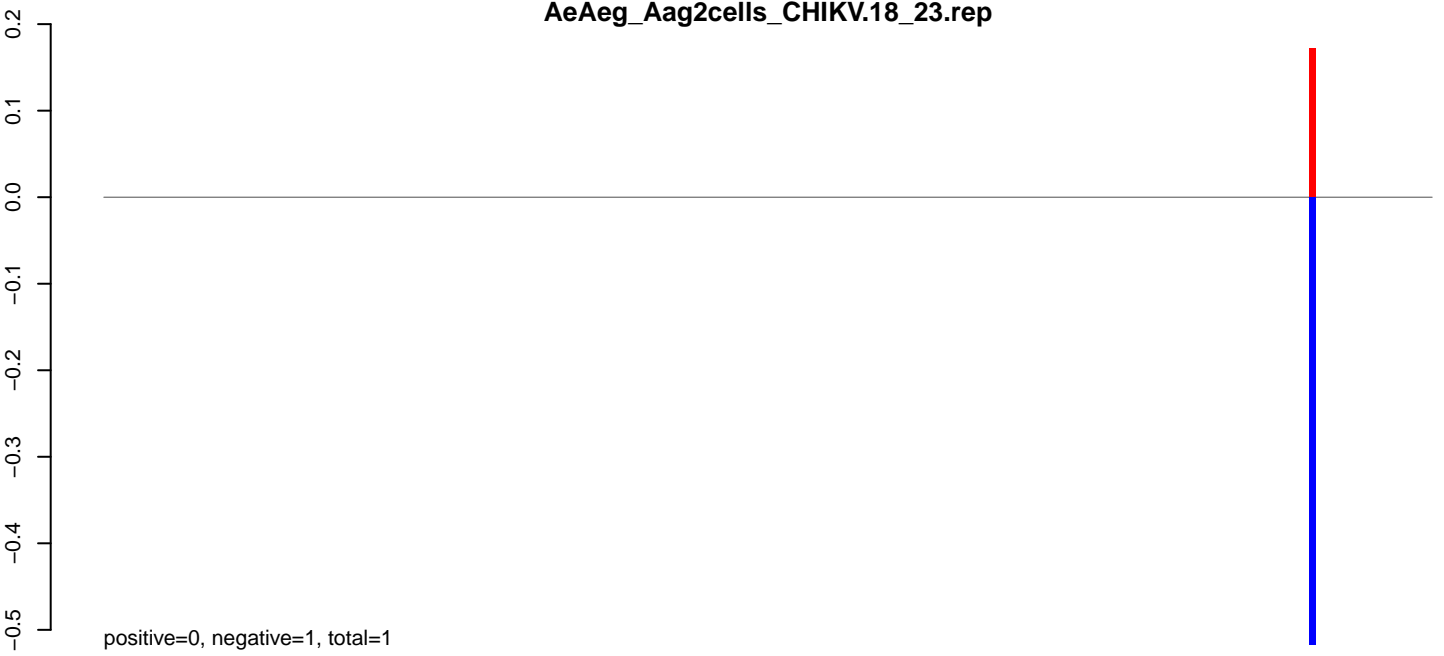
AeAeg_Aag2cells_CHIKV.24_35.rep



AeAeg_Aag2cells_CHIKV.rep

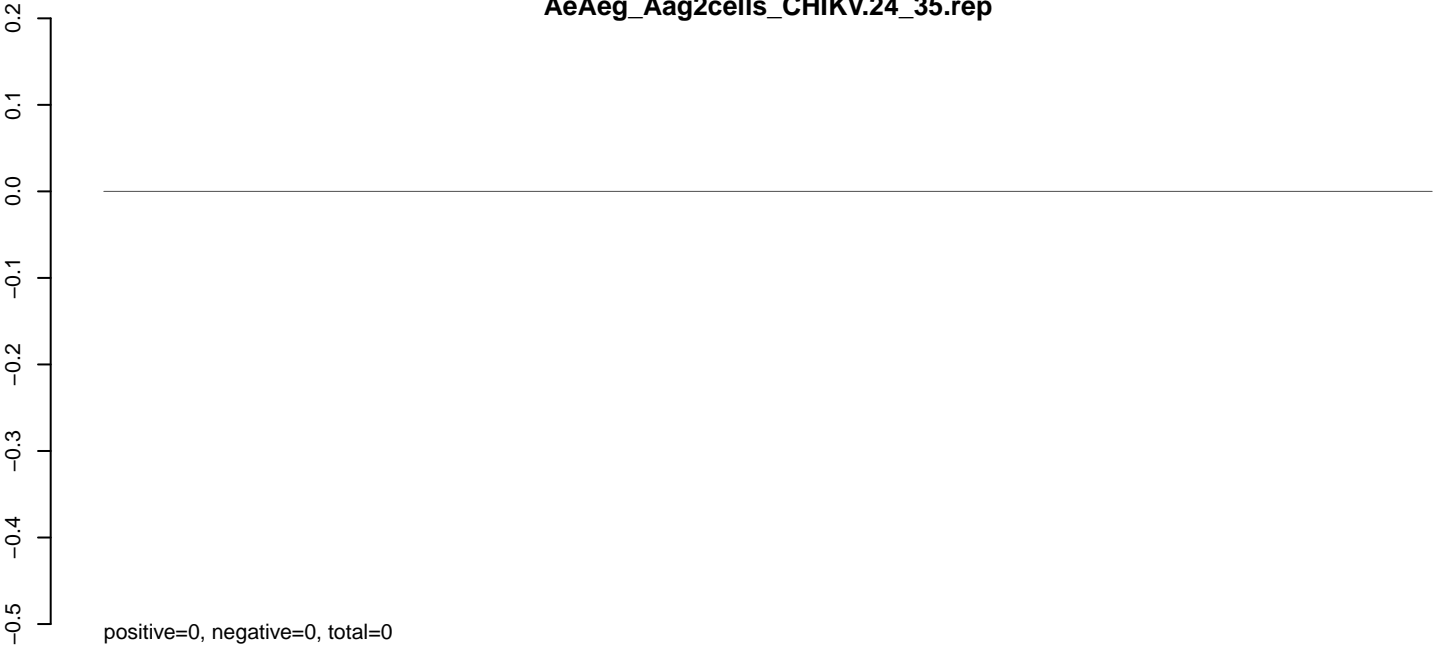


AeAeg_Aag2cells_CHIKV.18_23.rep



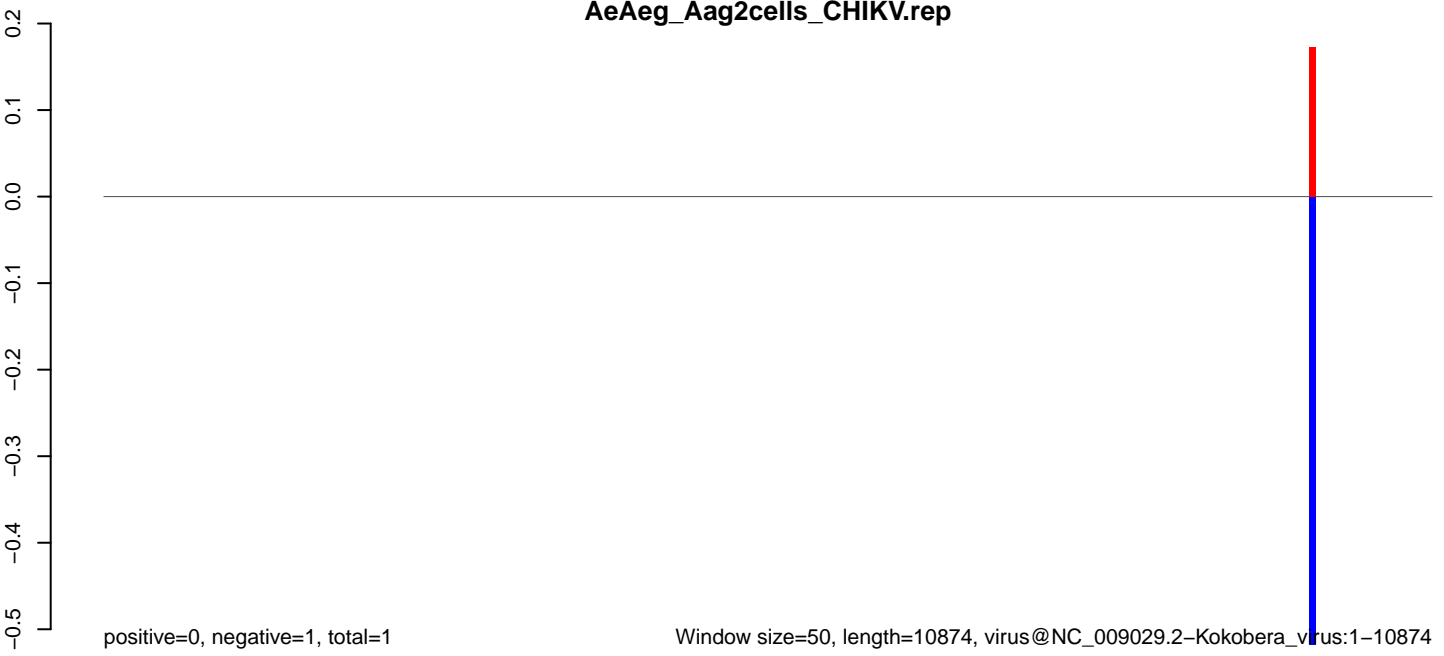
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AeAeg_Aag2cells_CHIKV.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2cells_CHIKV.rep

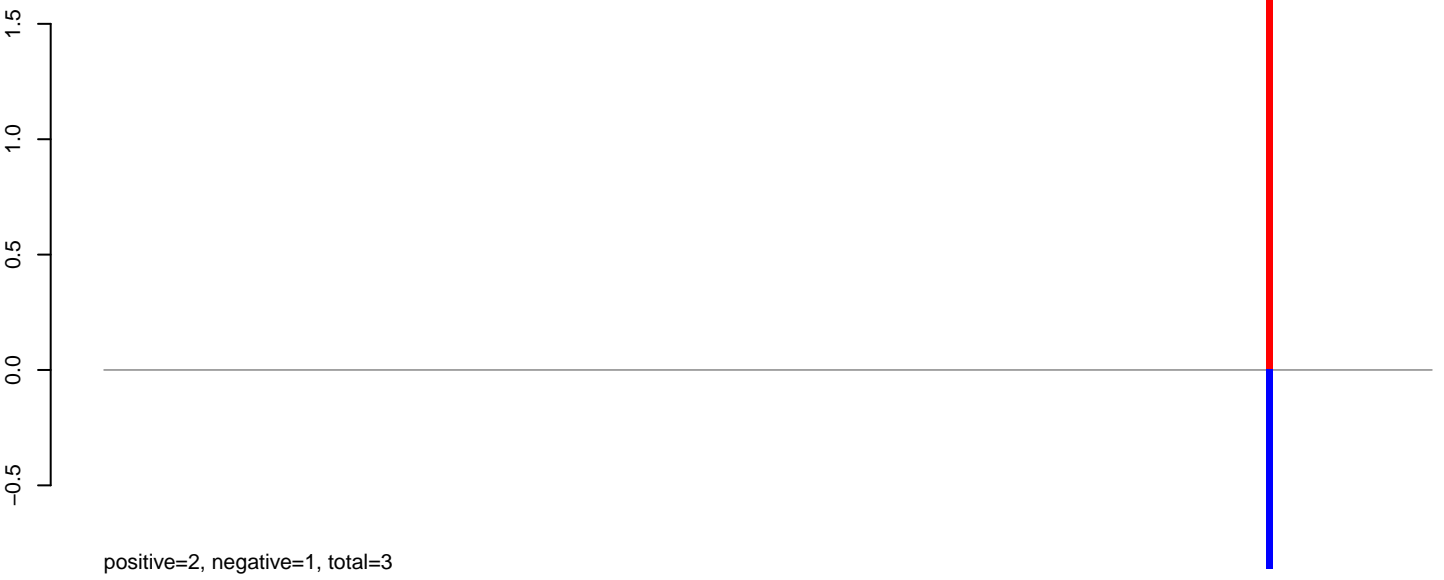


positive=0, negative=1, total=1

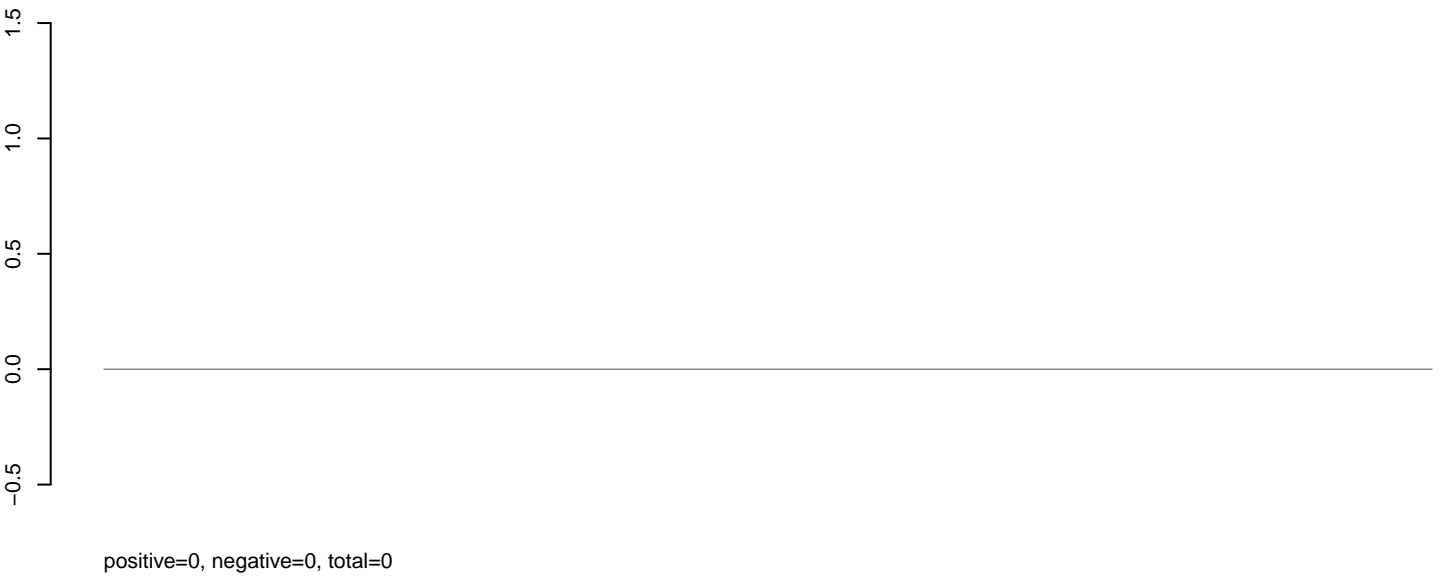
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AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep



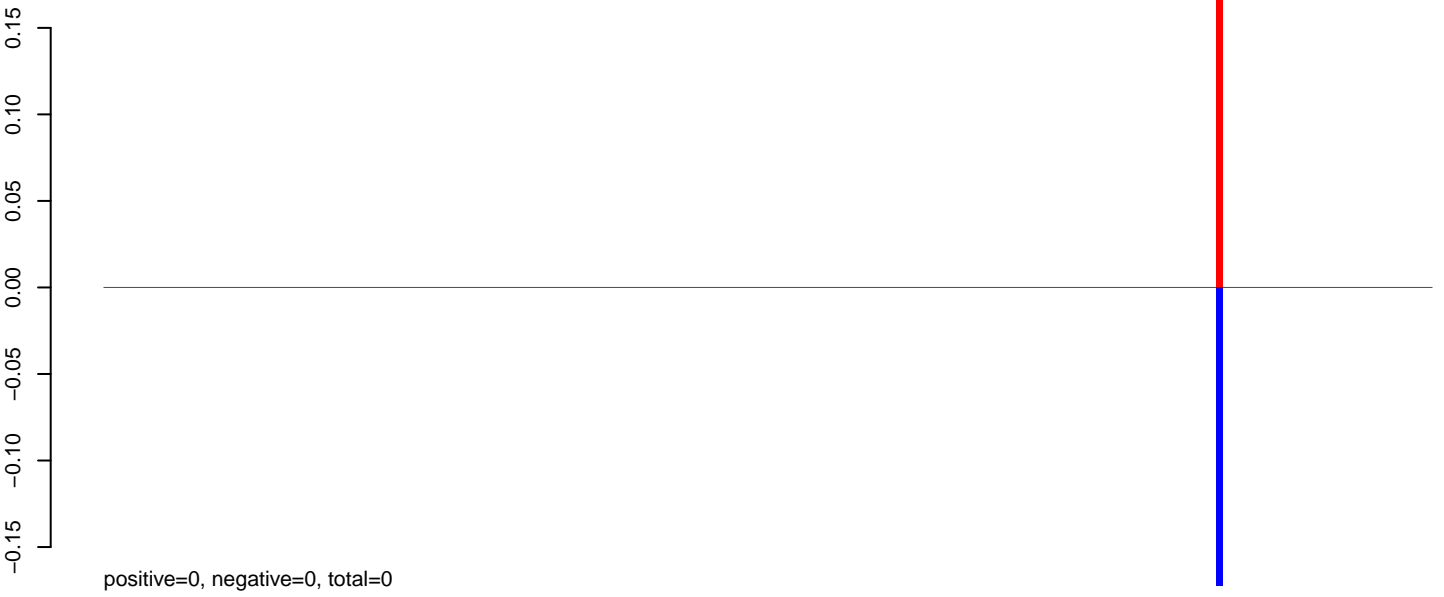
AeAeg_Aag2cells_CHIKV.rep



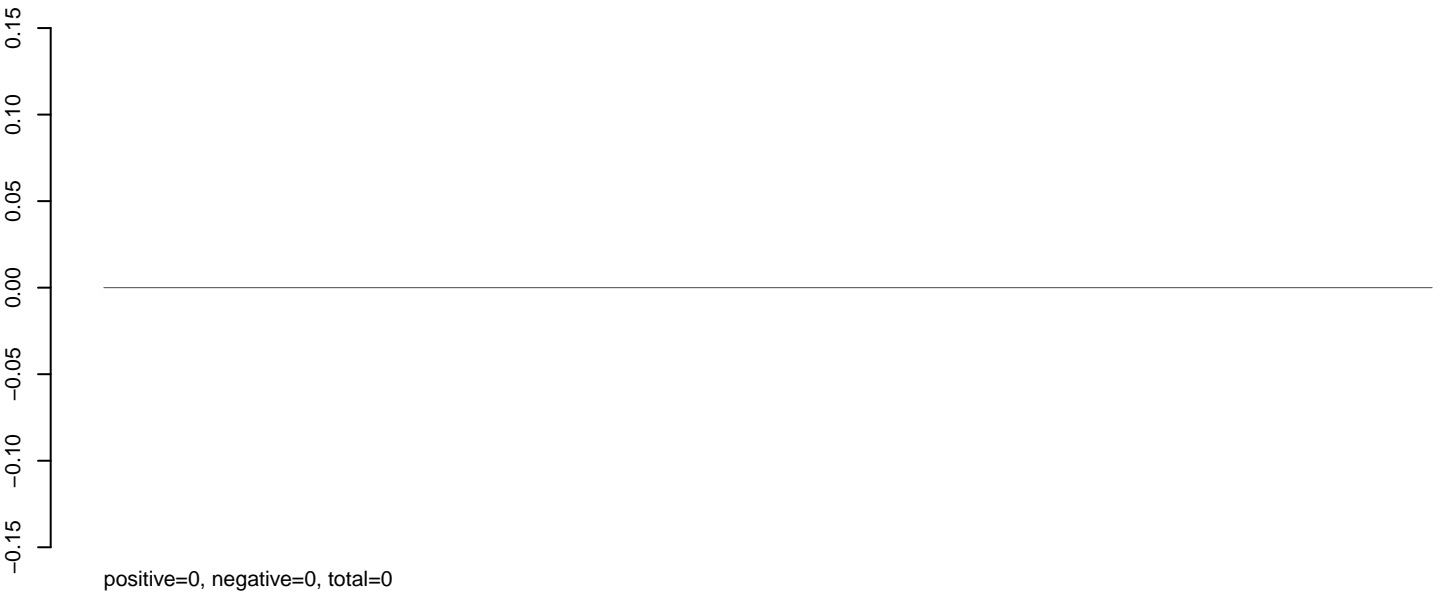
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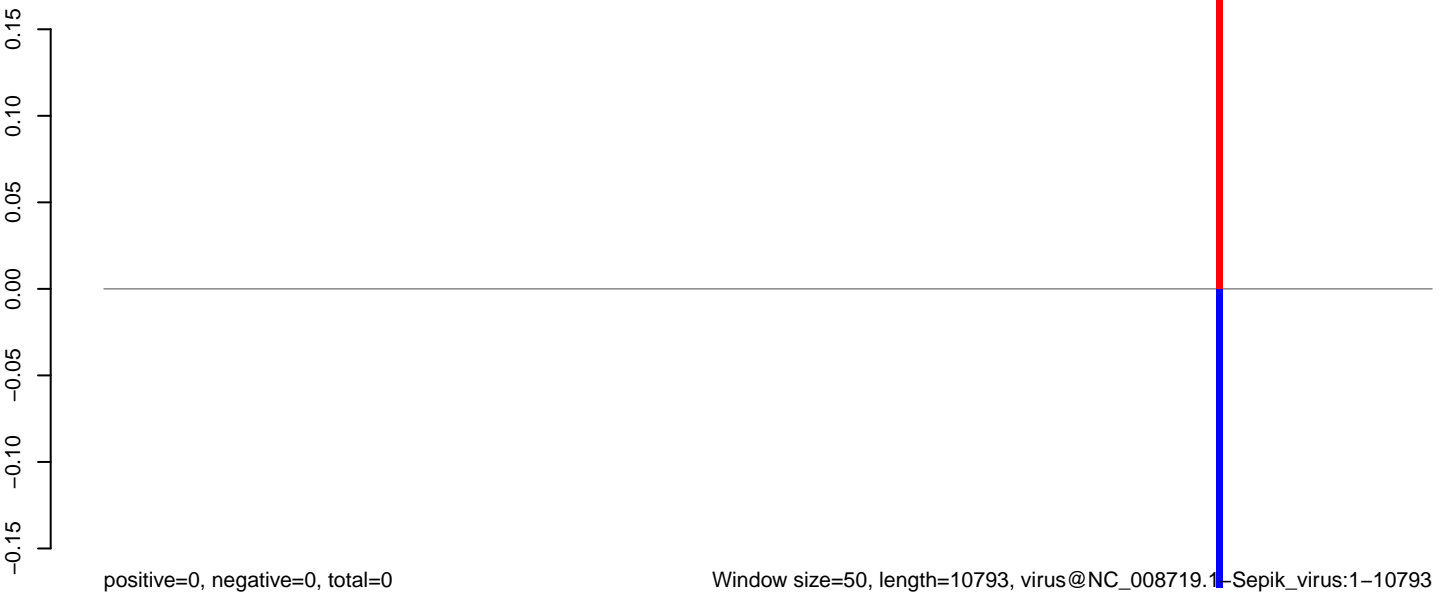
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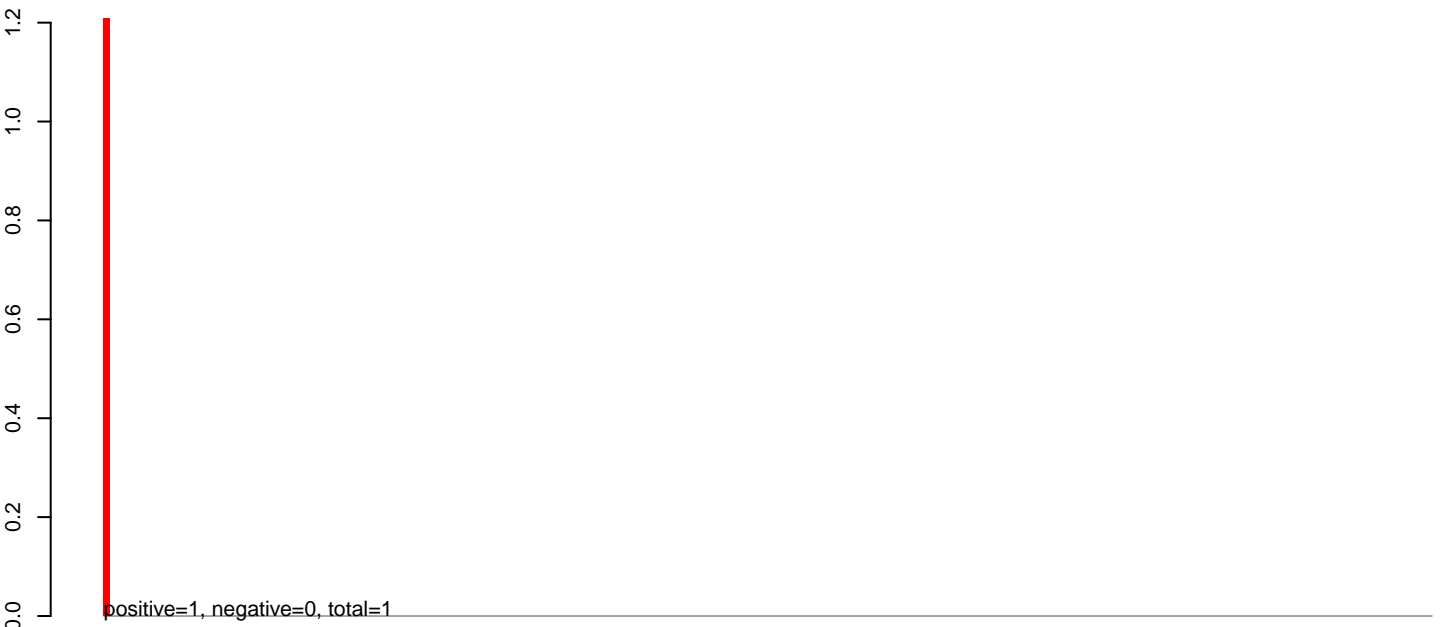
AeAeg_Aag2cells_CHIKV.24_35.rep



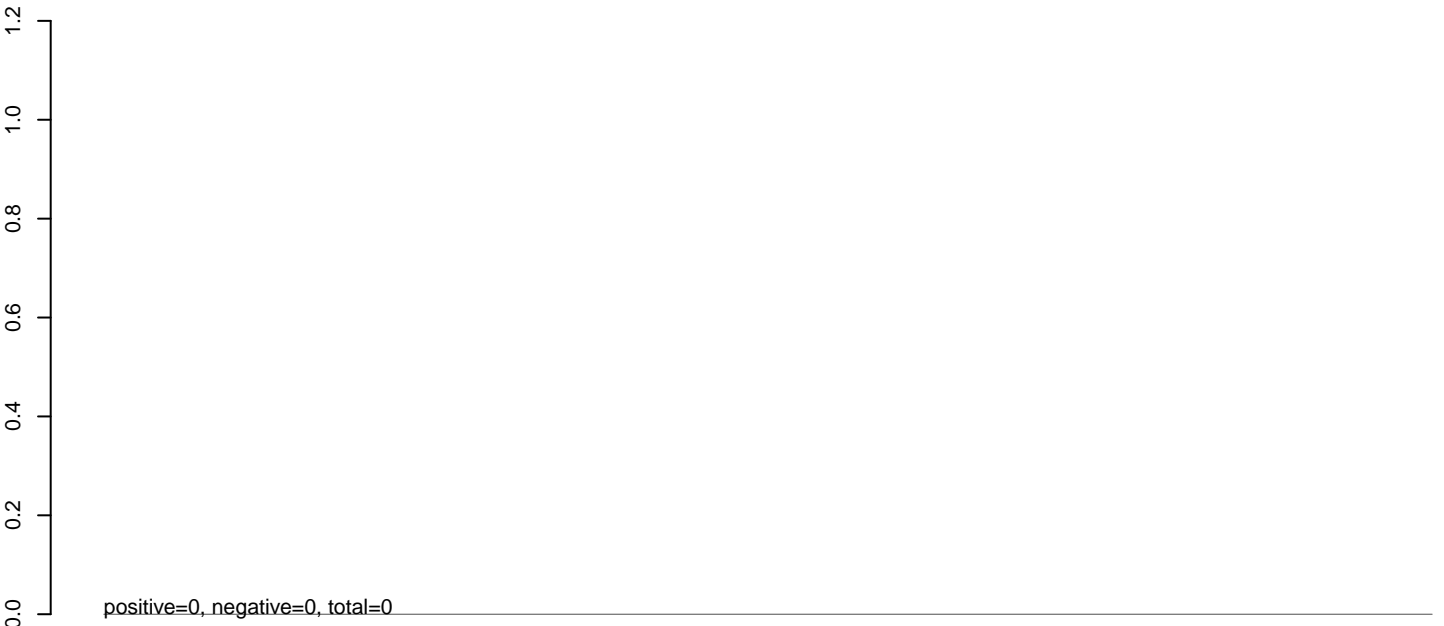
AeAeg_Aag2cells_CHIKV.rep



AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

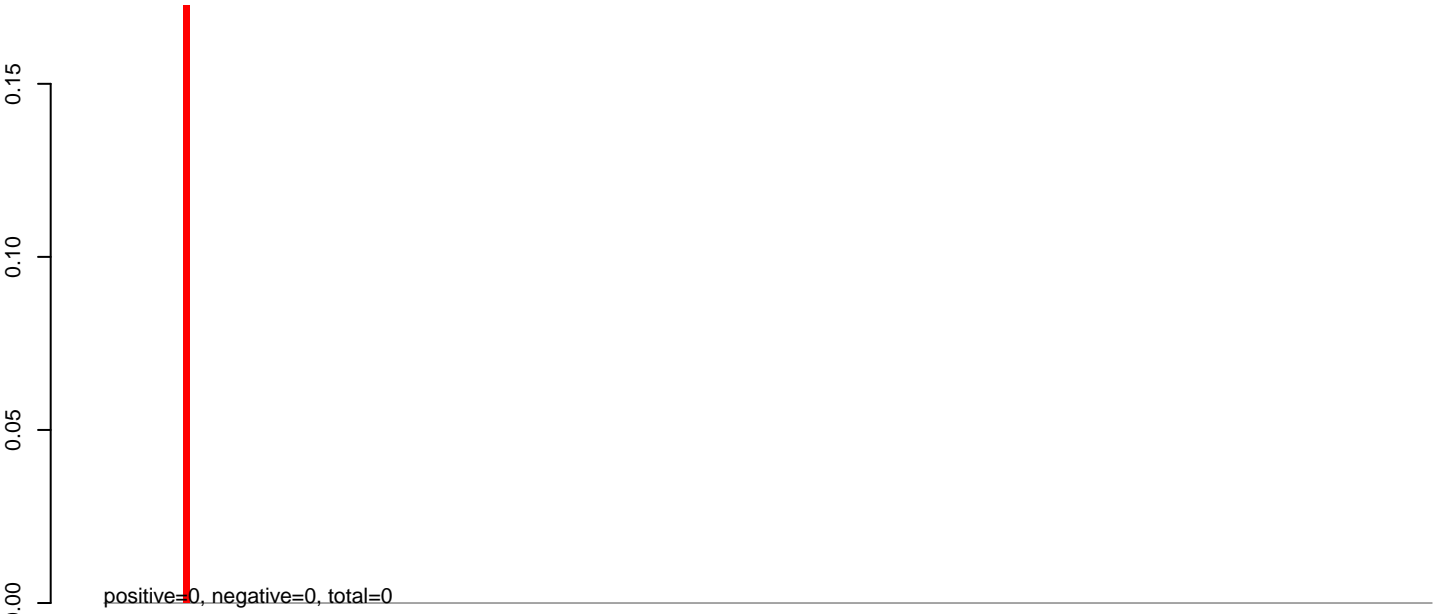


AeAeg_Aag2cells_CHIKV.rep

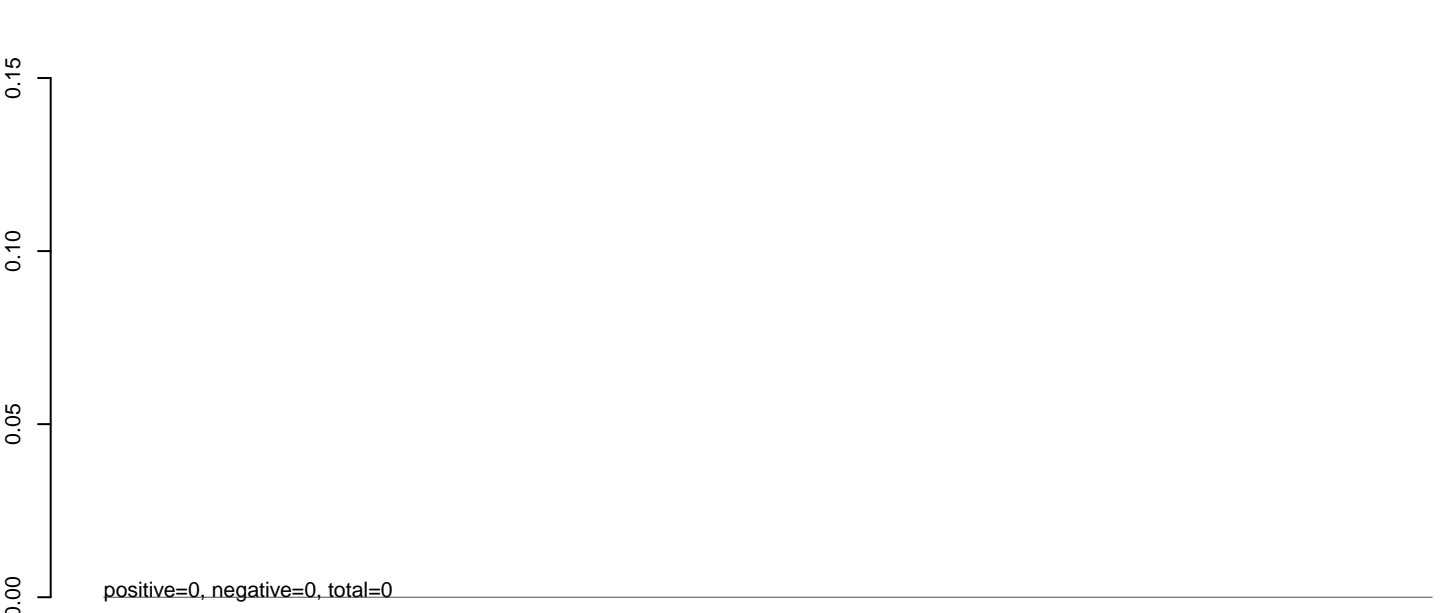


0 2000 4000 6000 8000 10000

AeAeg_Aag2cells_CHIKV.18_23.rep



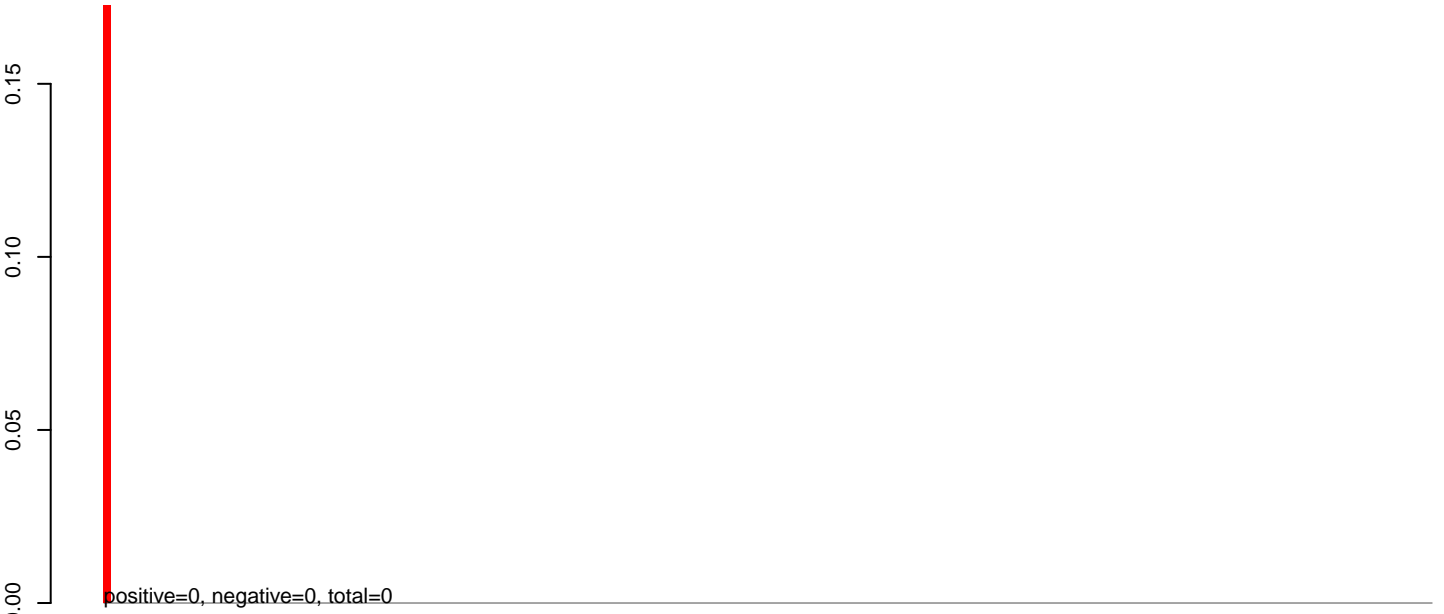
AeAeg_Aag2cells_CHIKV.24_35.rep



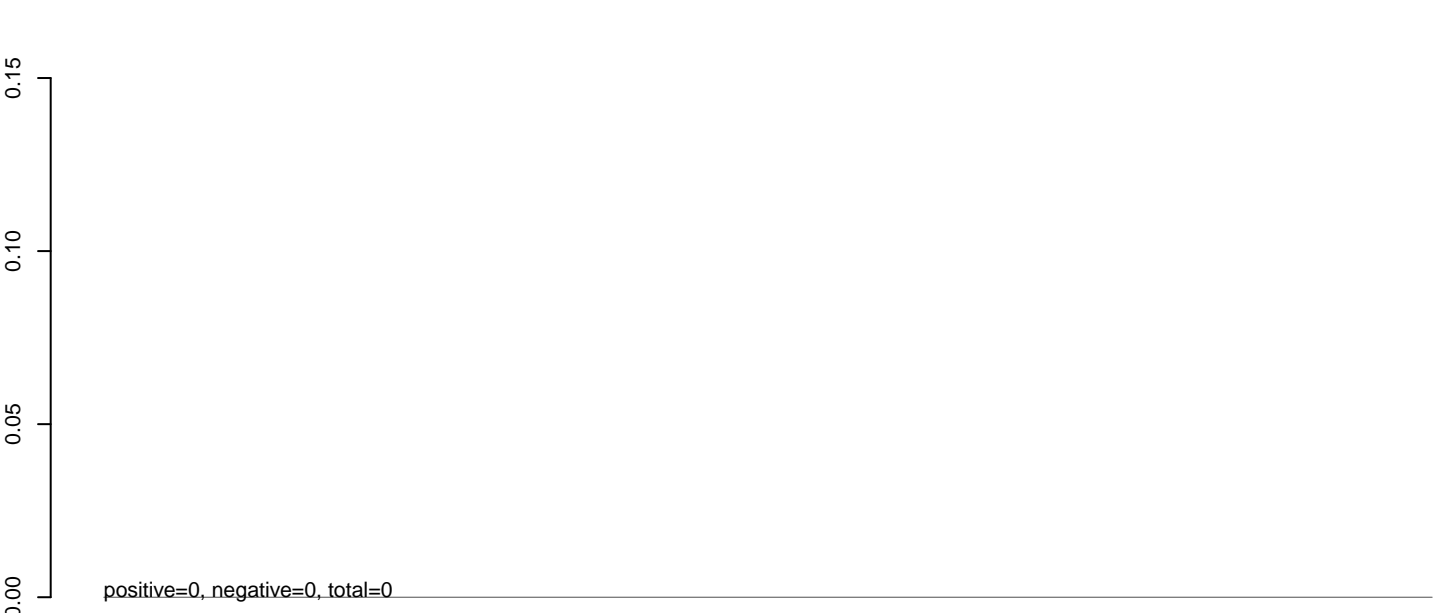
AeAeg_Aag2cells_CHIKV.rep



AeAeg_Aag2cells_CHIKV.18_23.rep



AeAeg_Aag2cells_CHIKV.24_35.rep

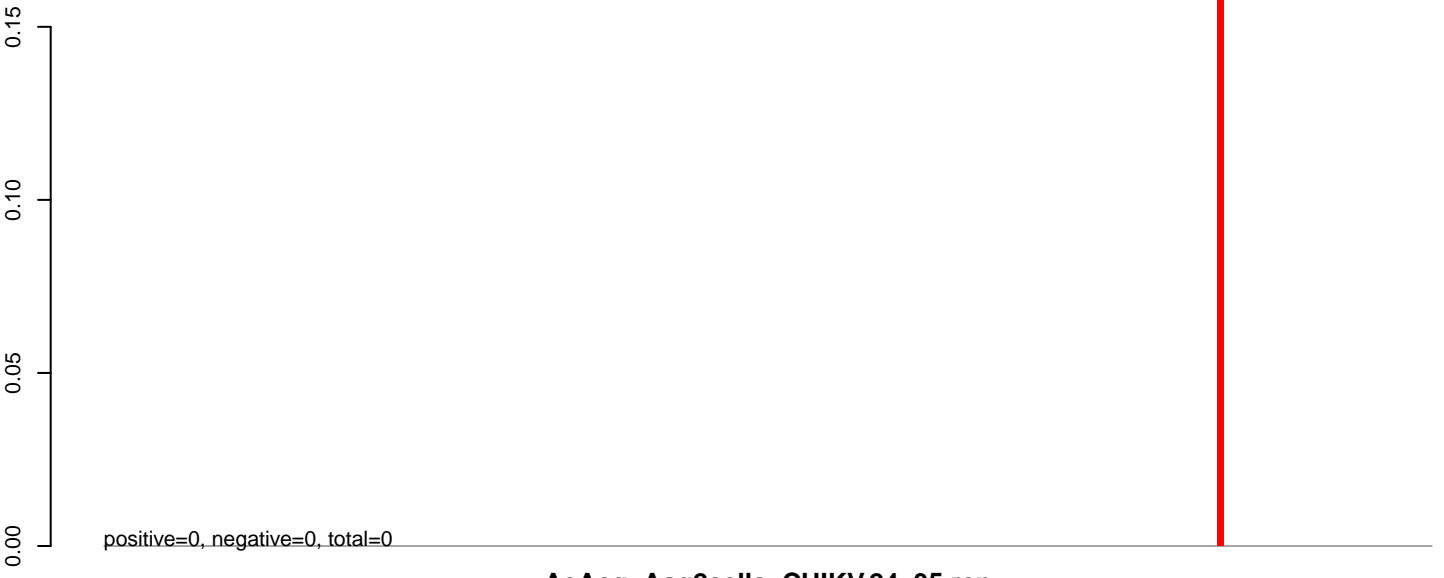


AeAeg_Aag2cells_CHIKV.rep

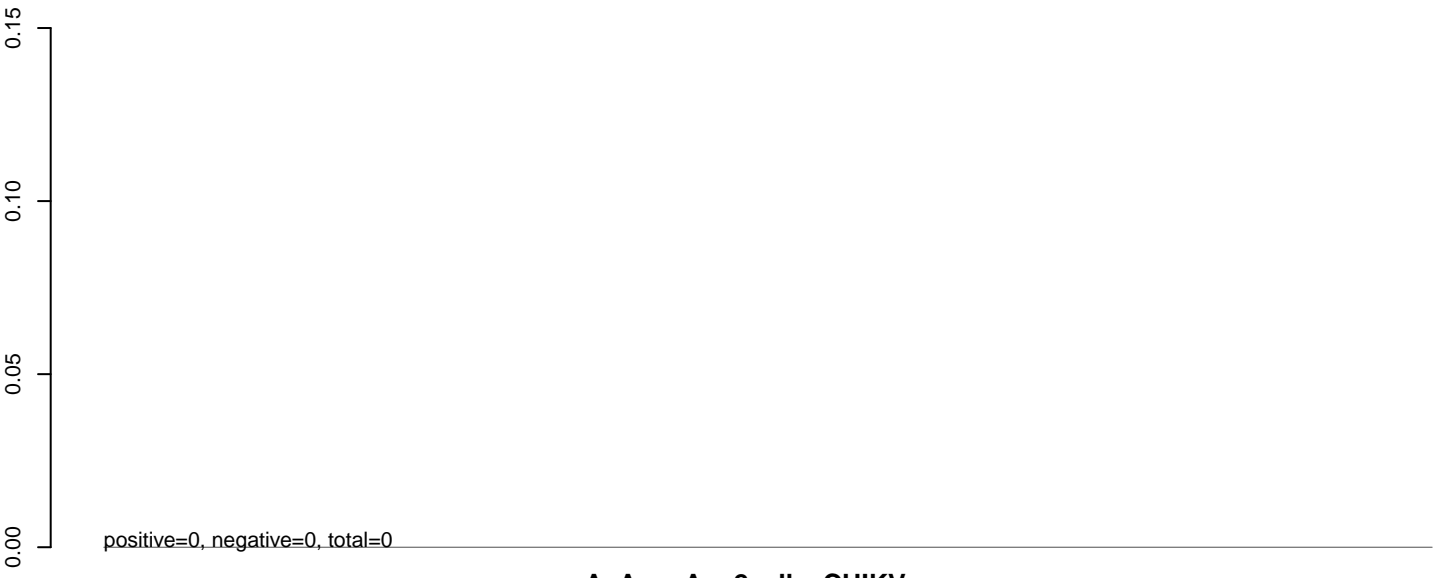


WINDOMAS NC_003924.1-Cricket paralysis virus nonstructural polyprotein and structural polyprotein genes:1-9185

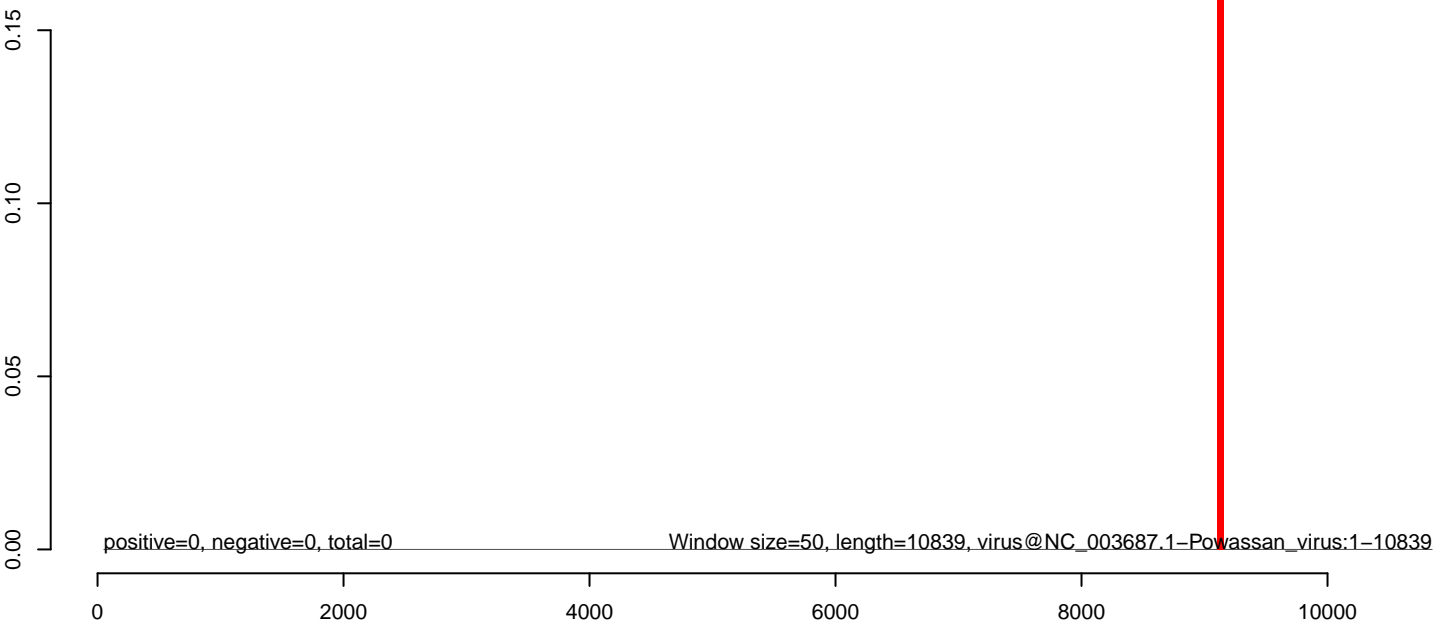
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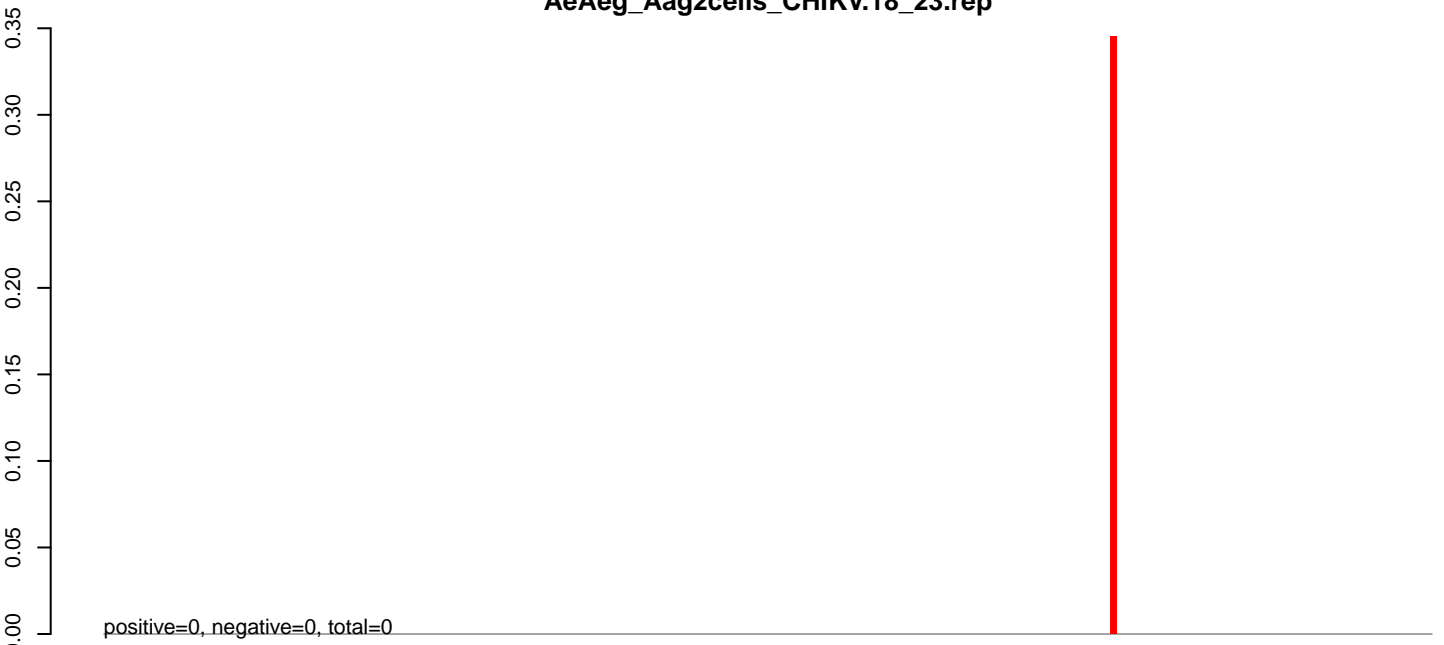
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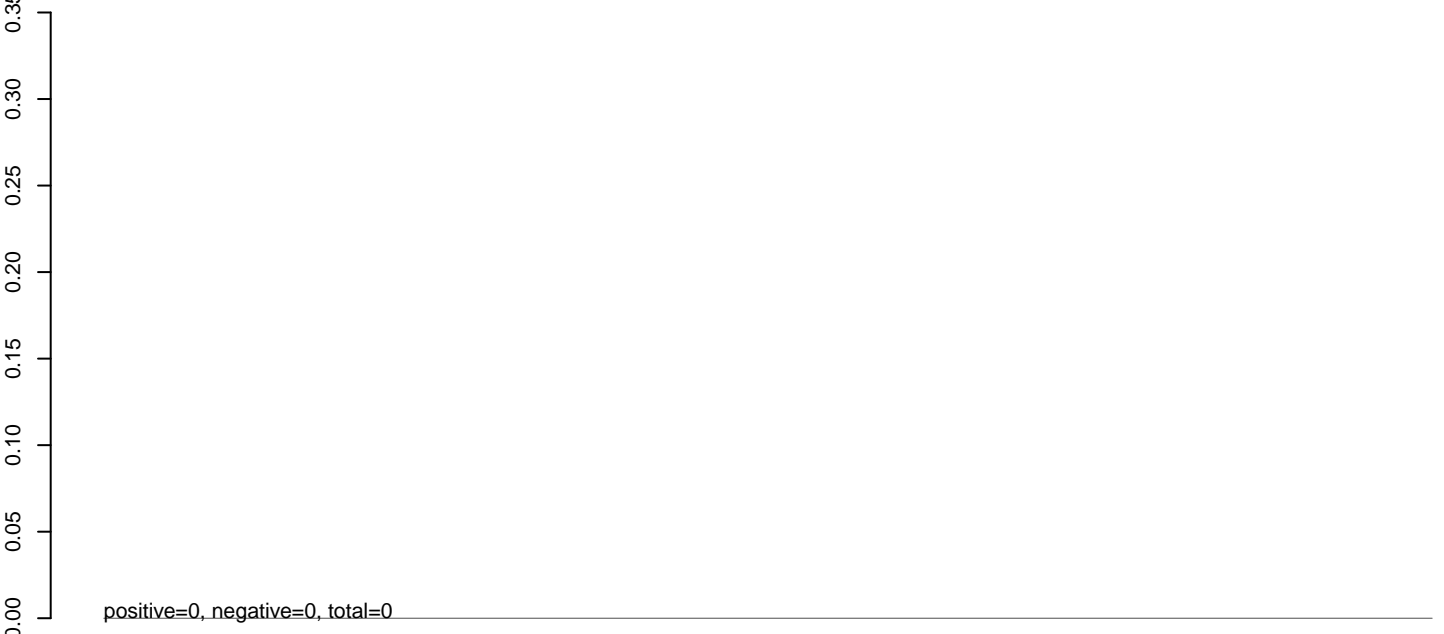
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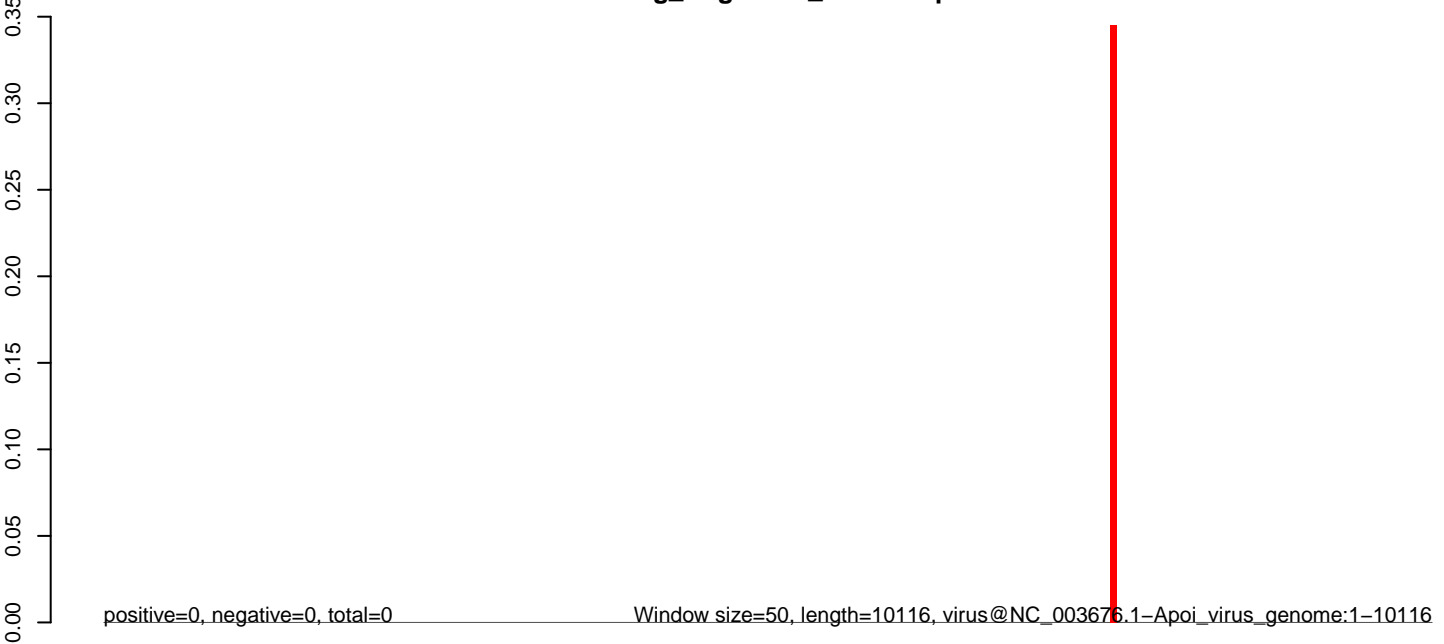
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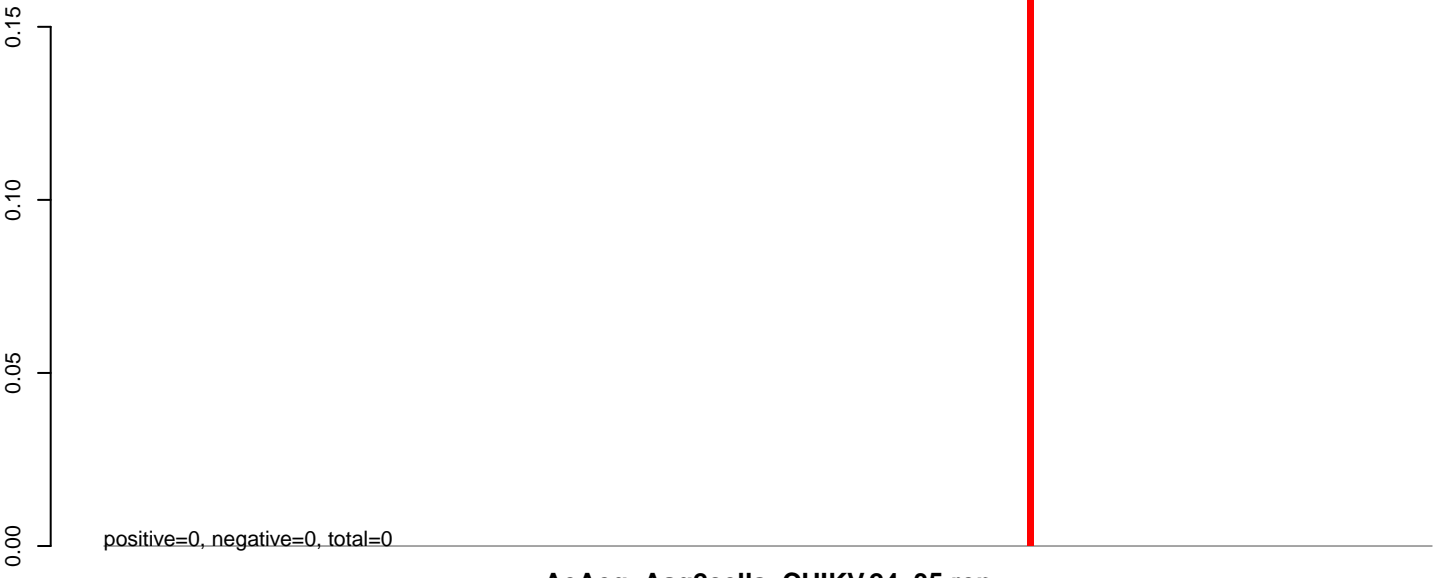


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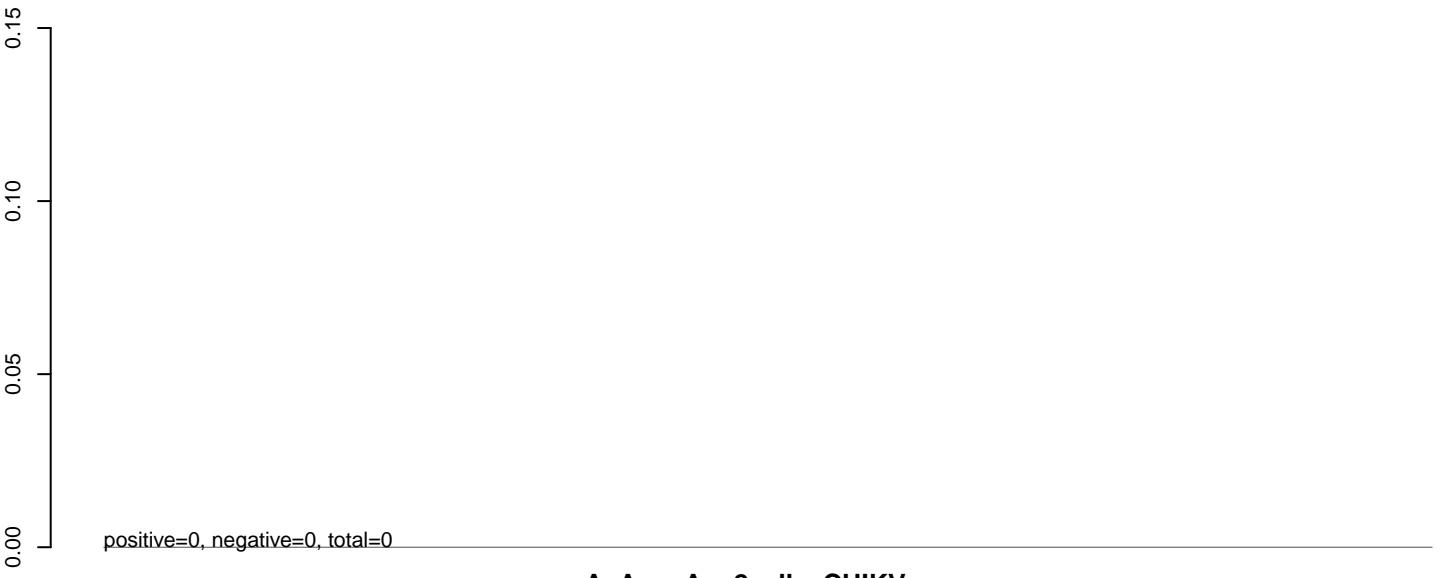


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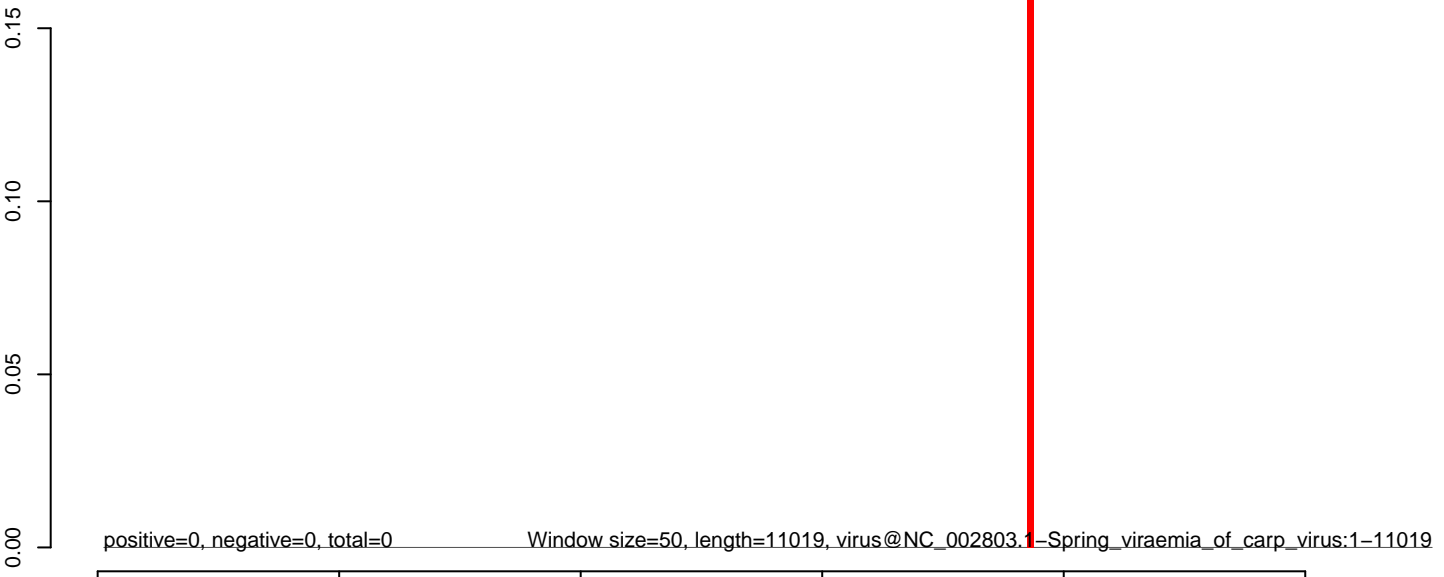
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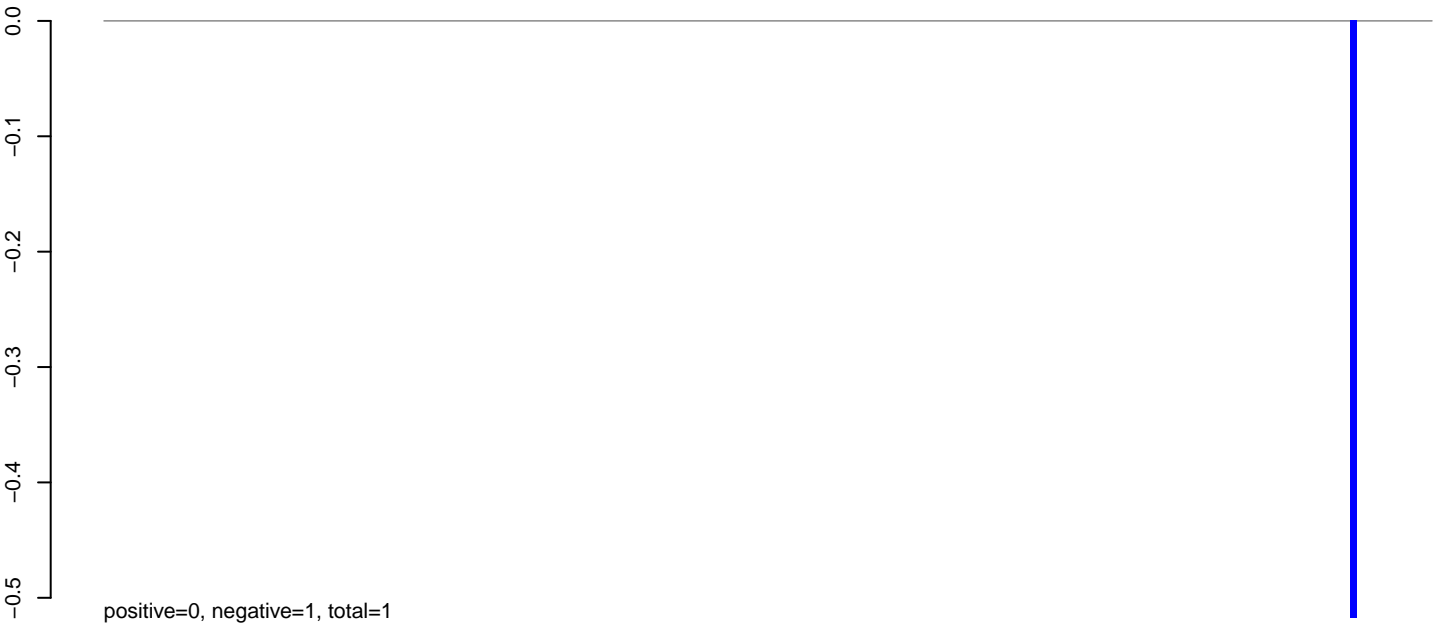
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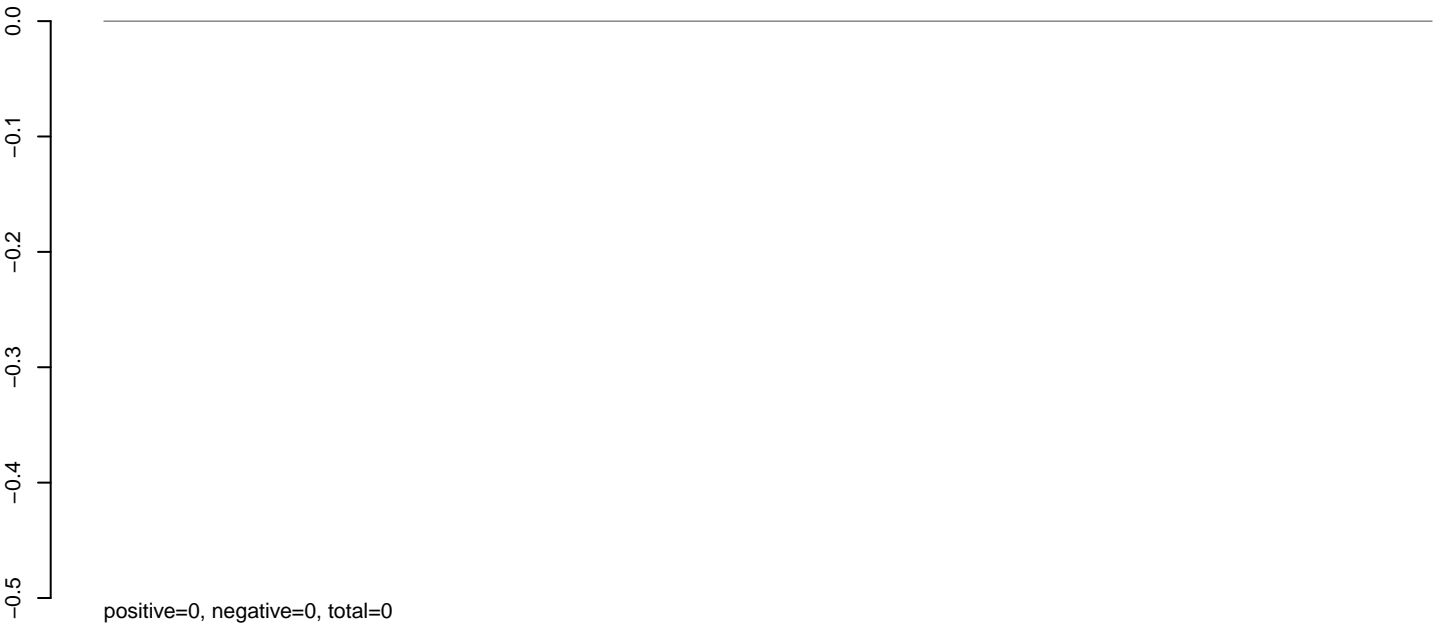
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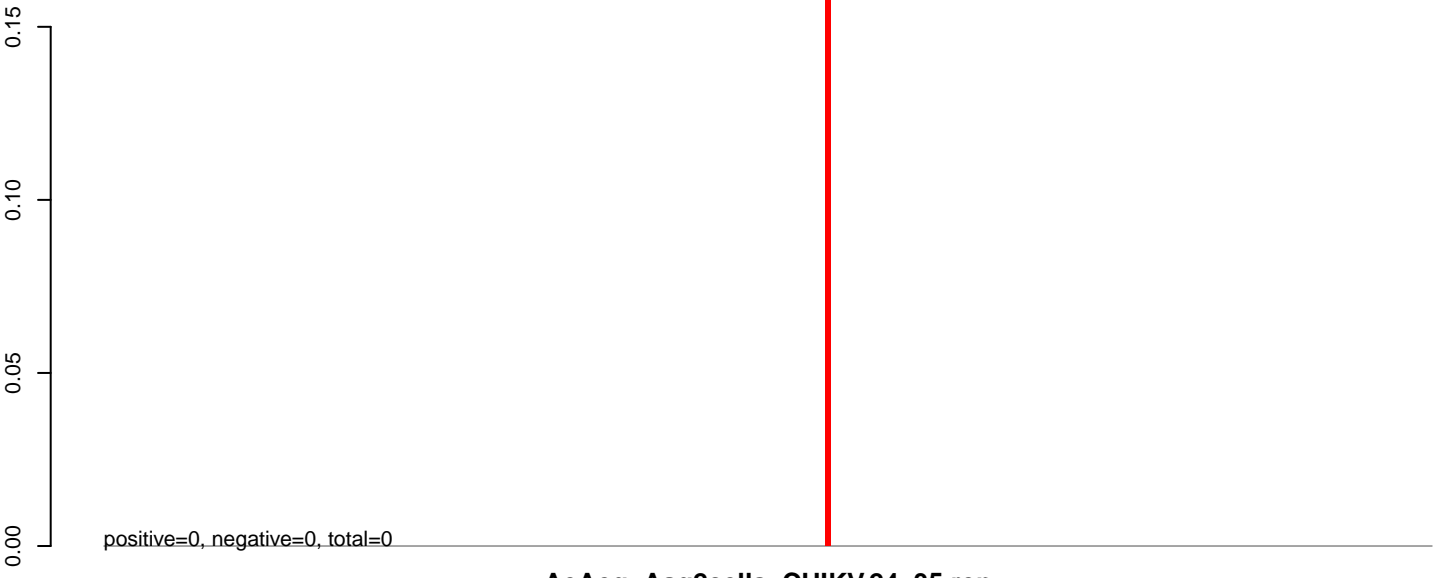
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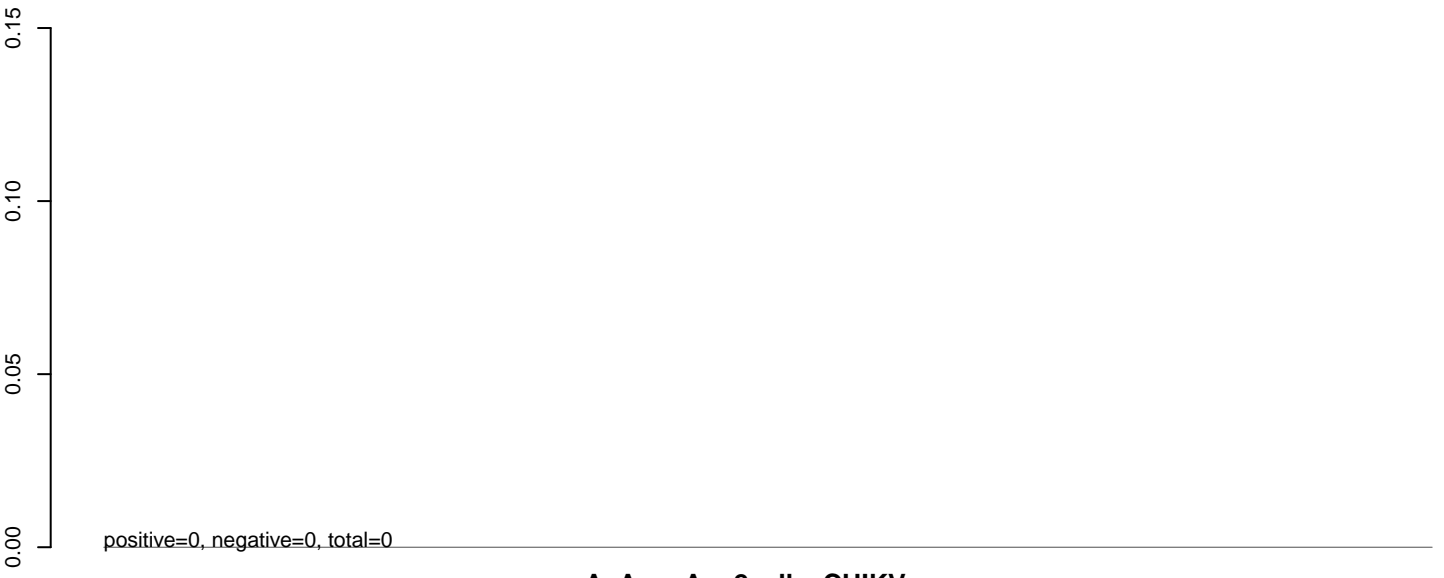
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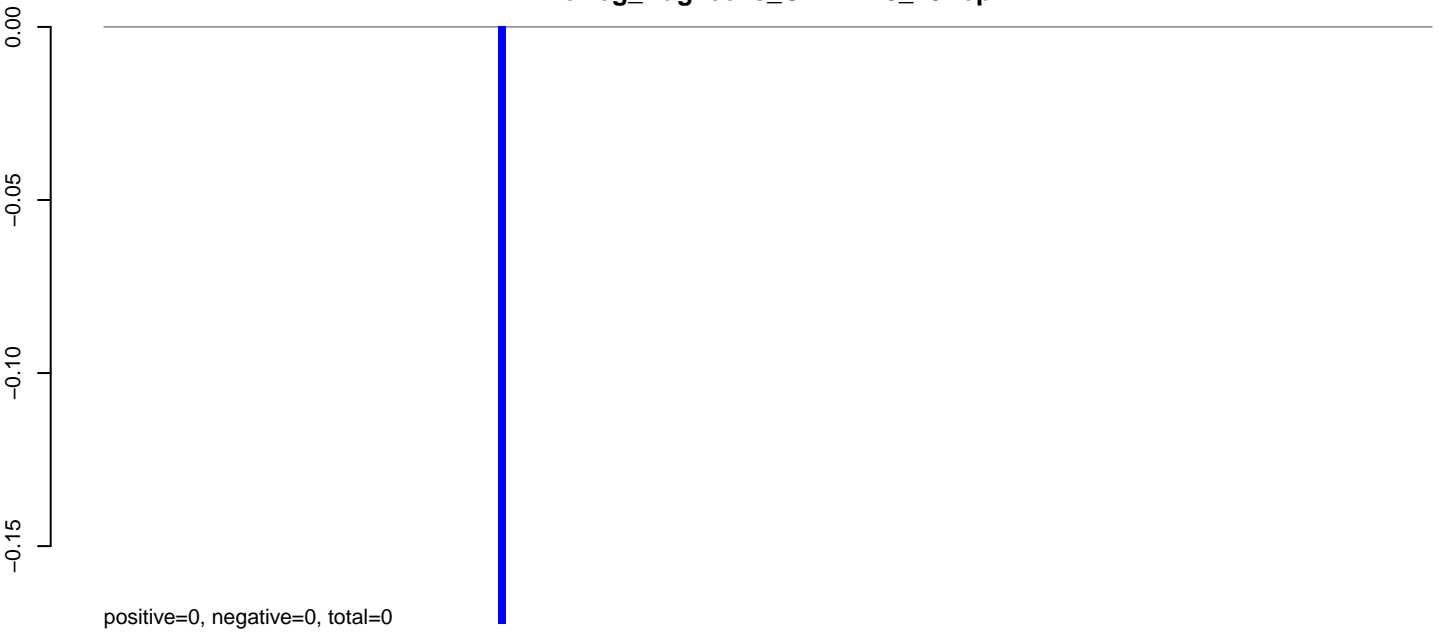
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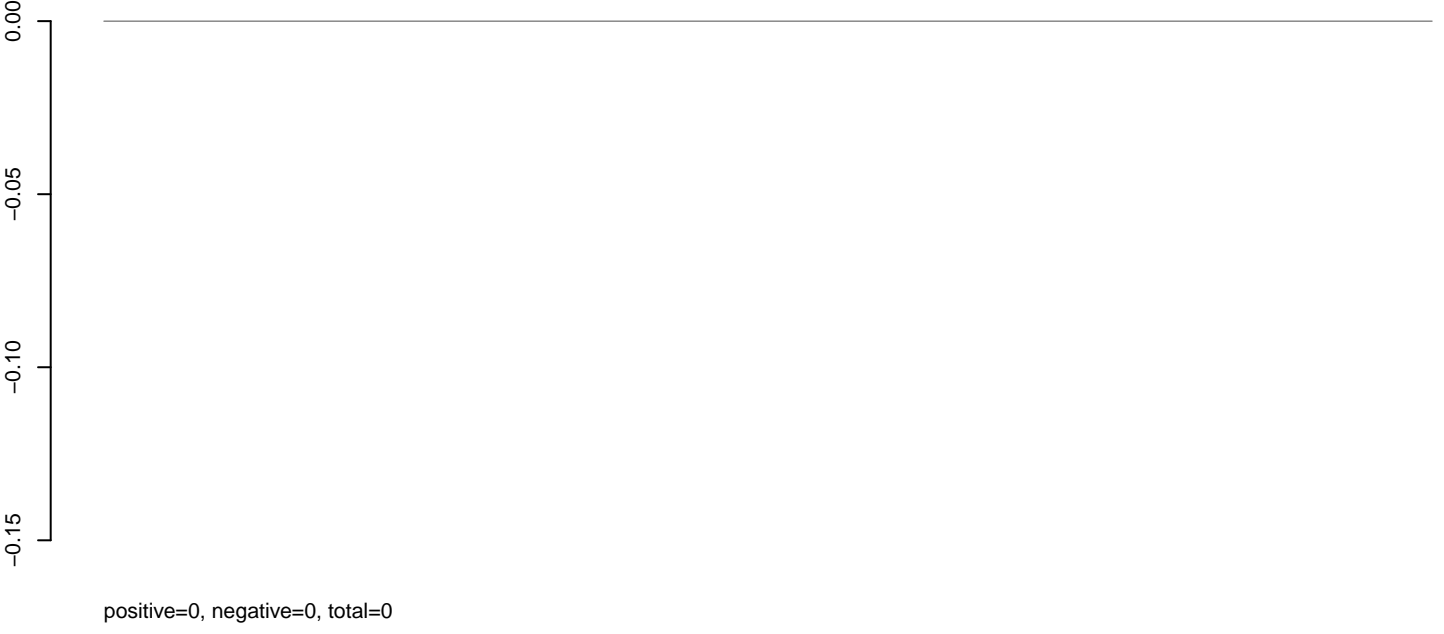
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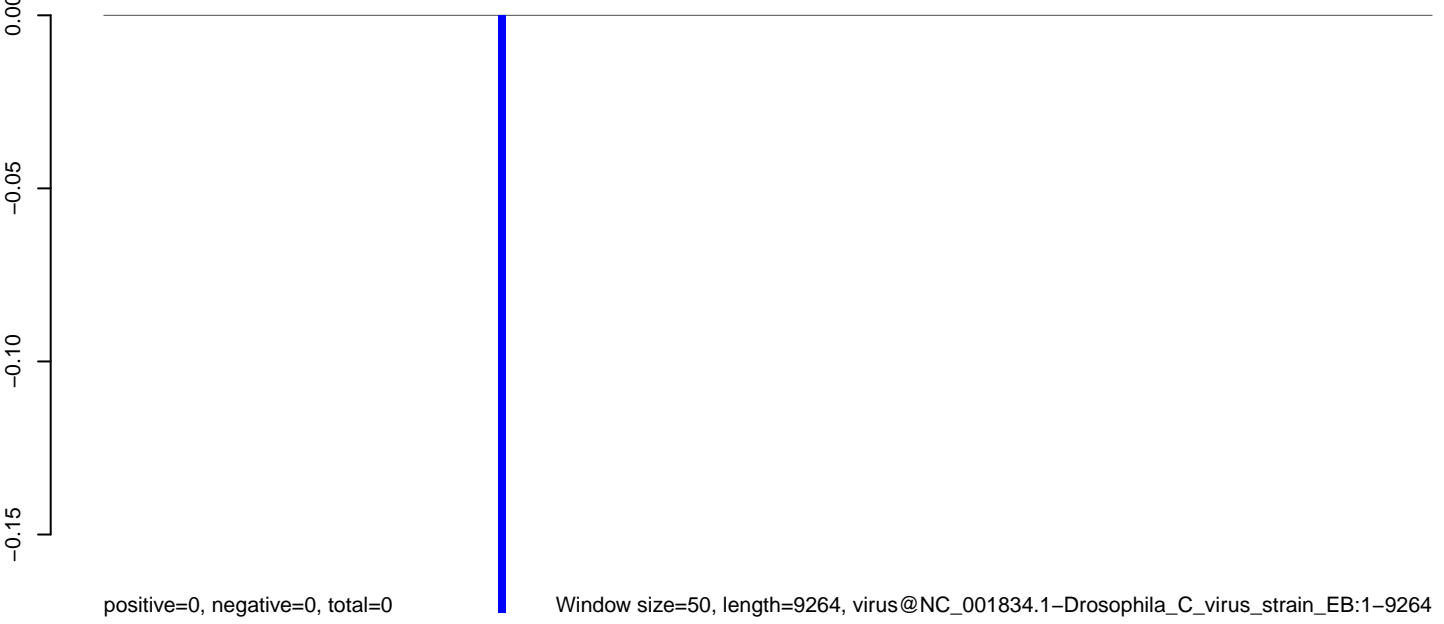
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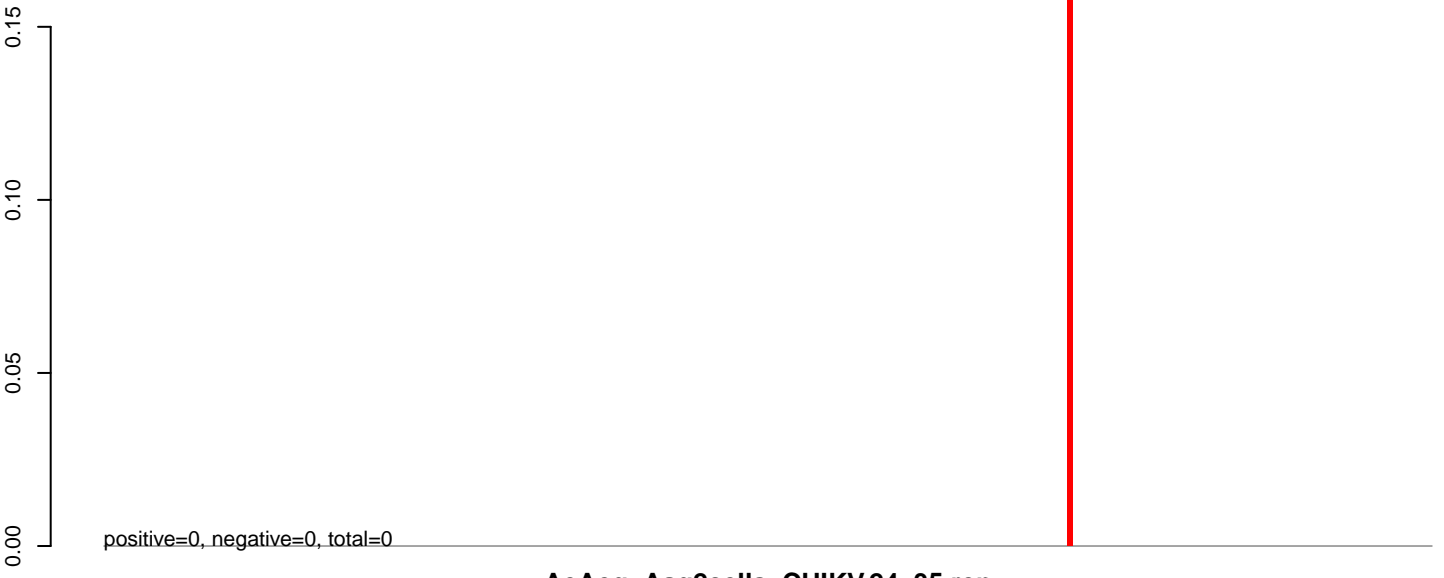
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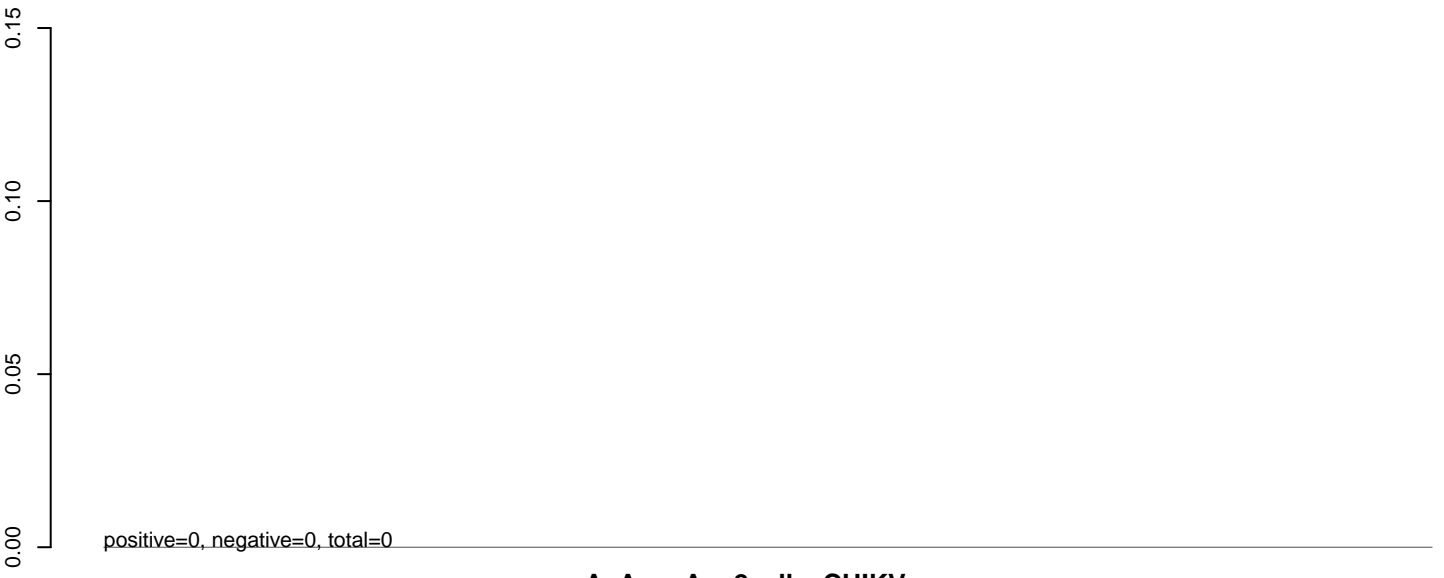
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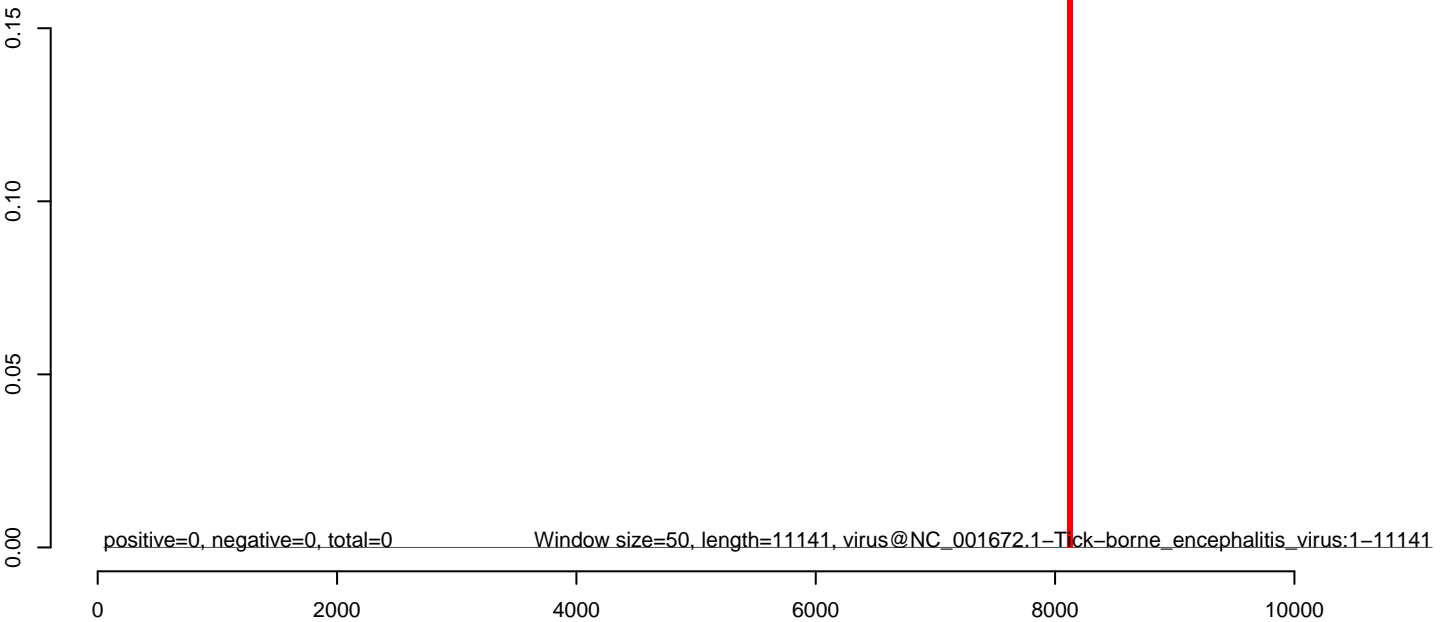
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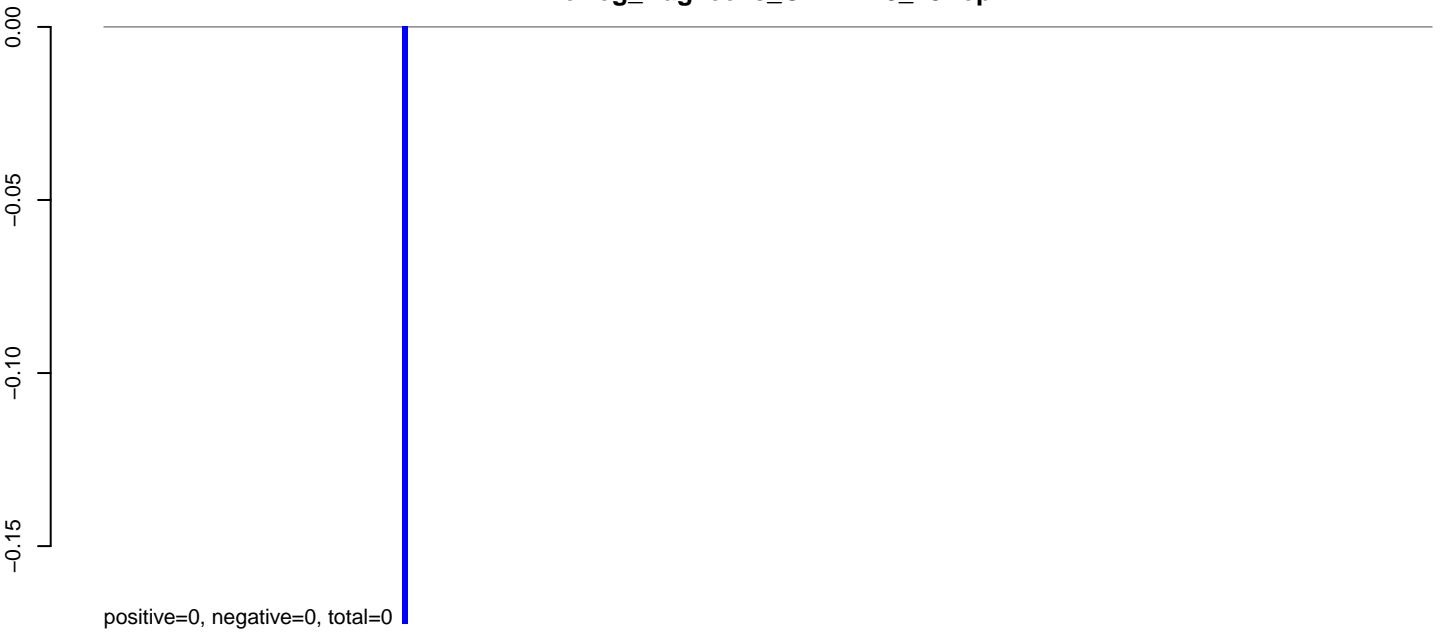
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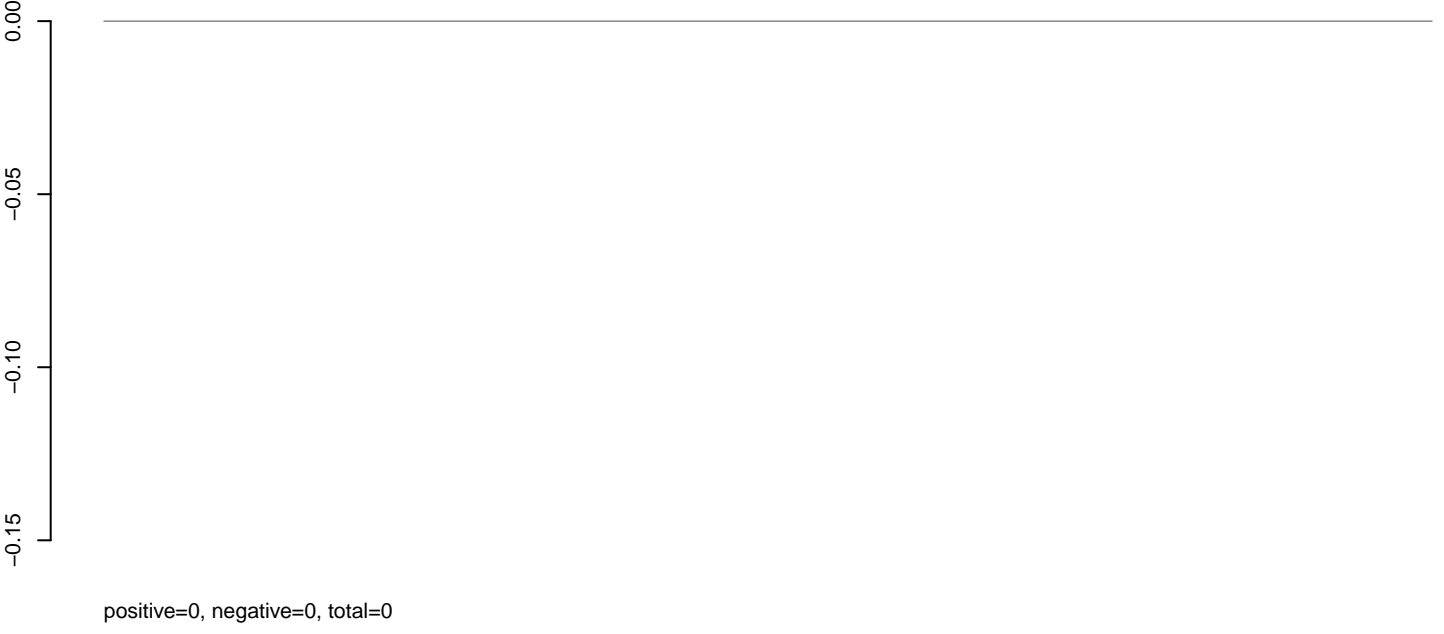
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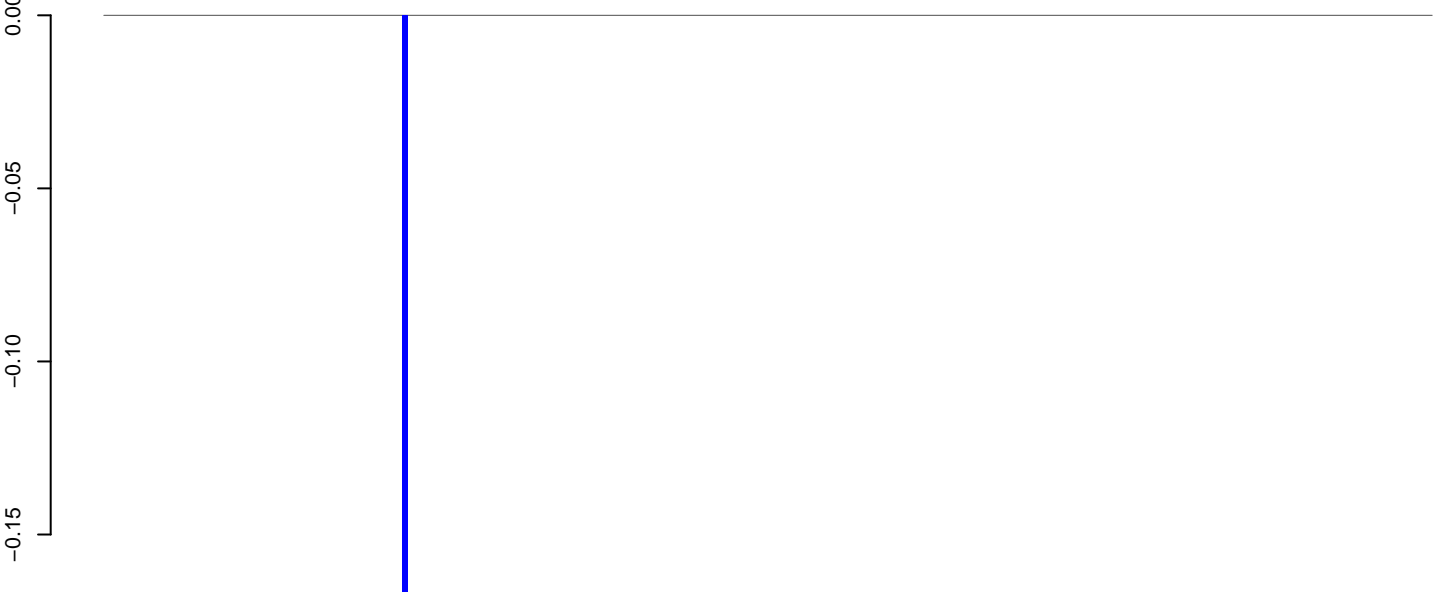
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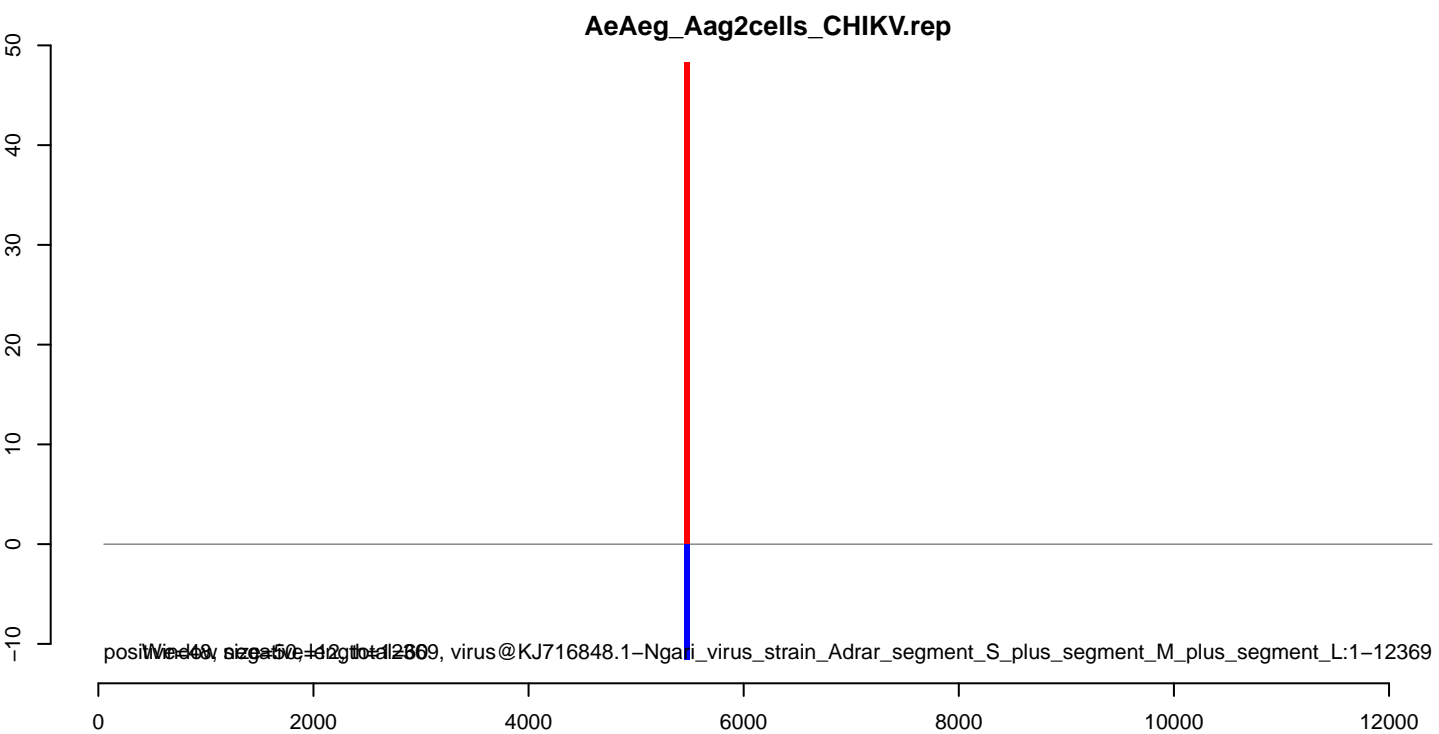
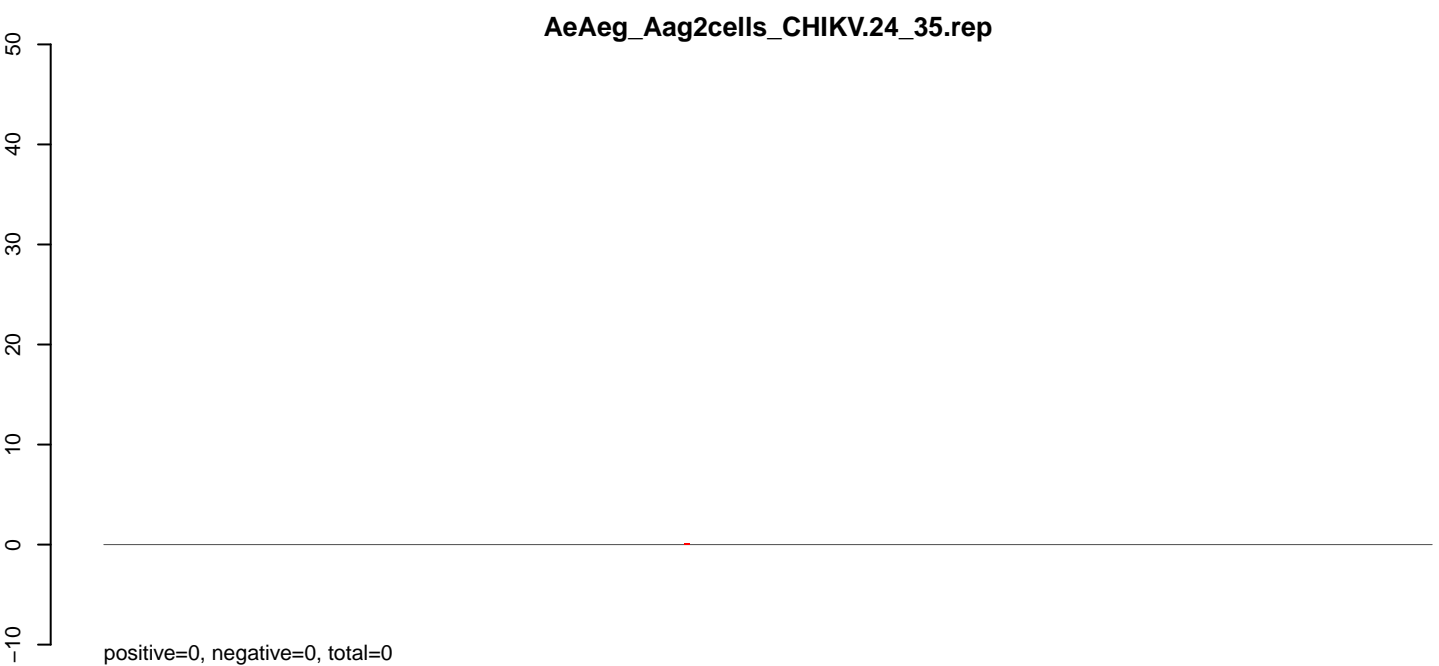
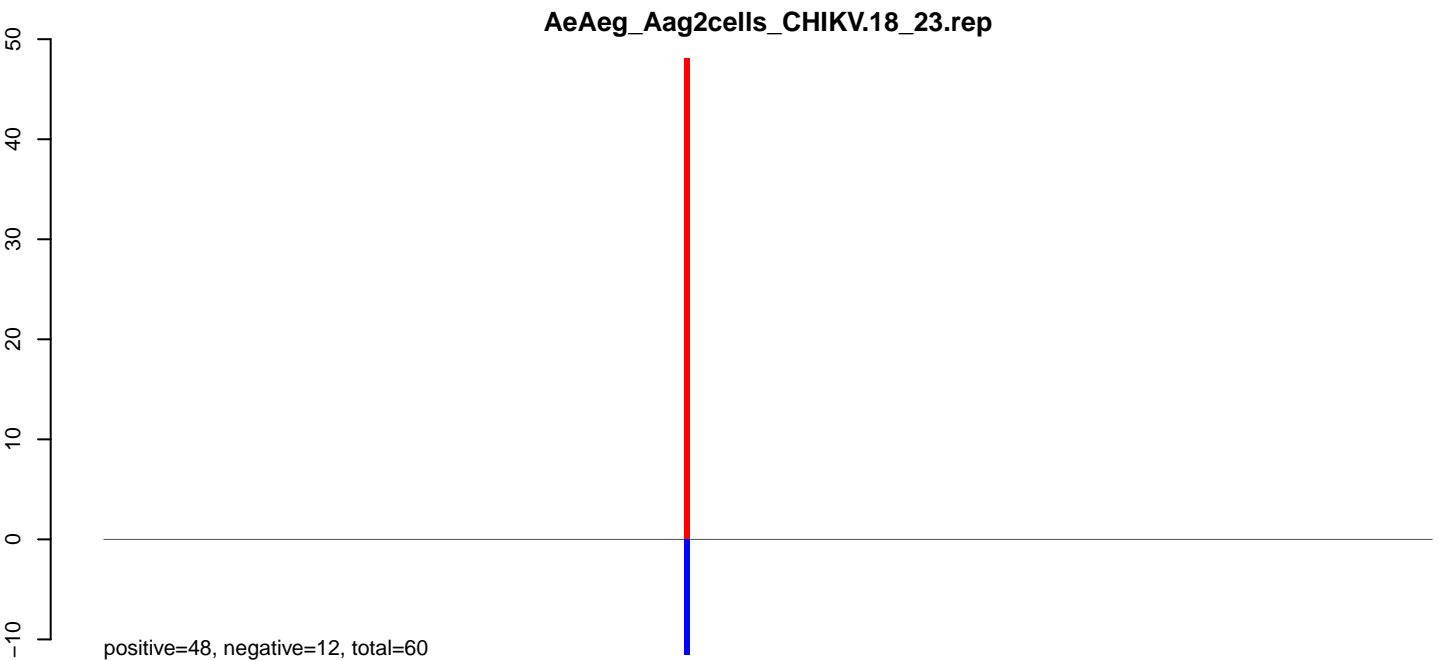


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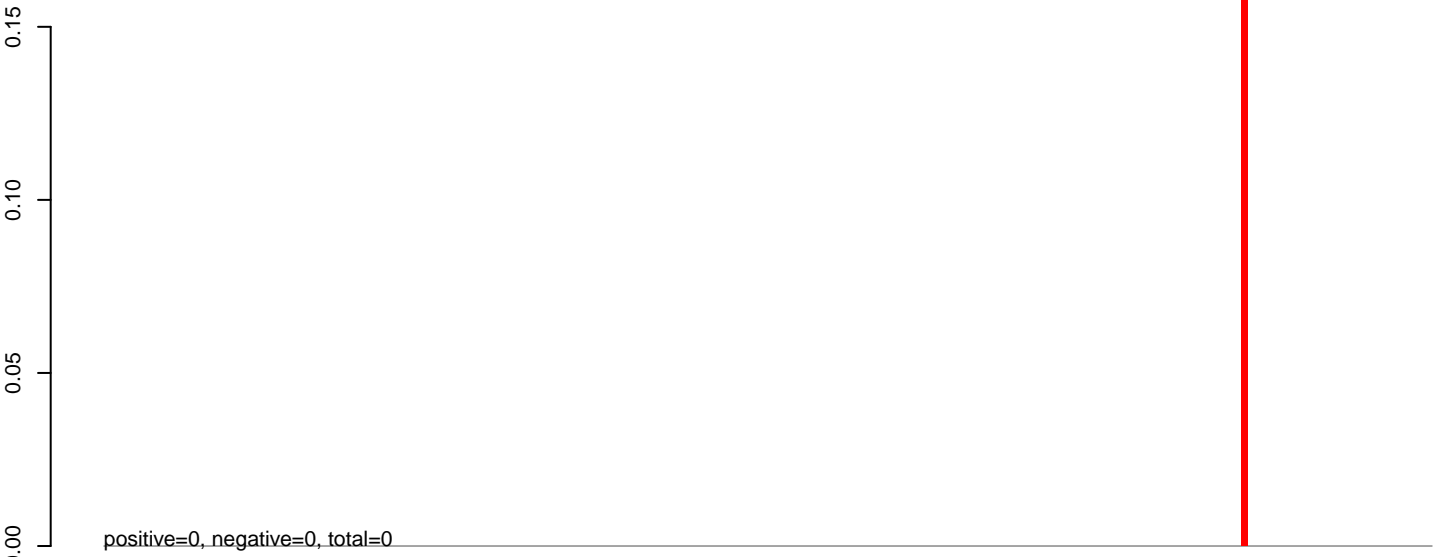


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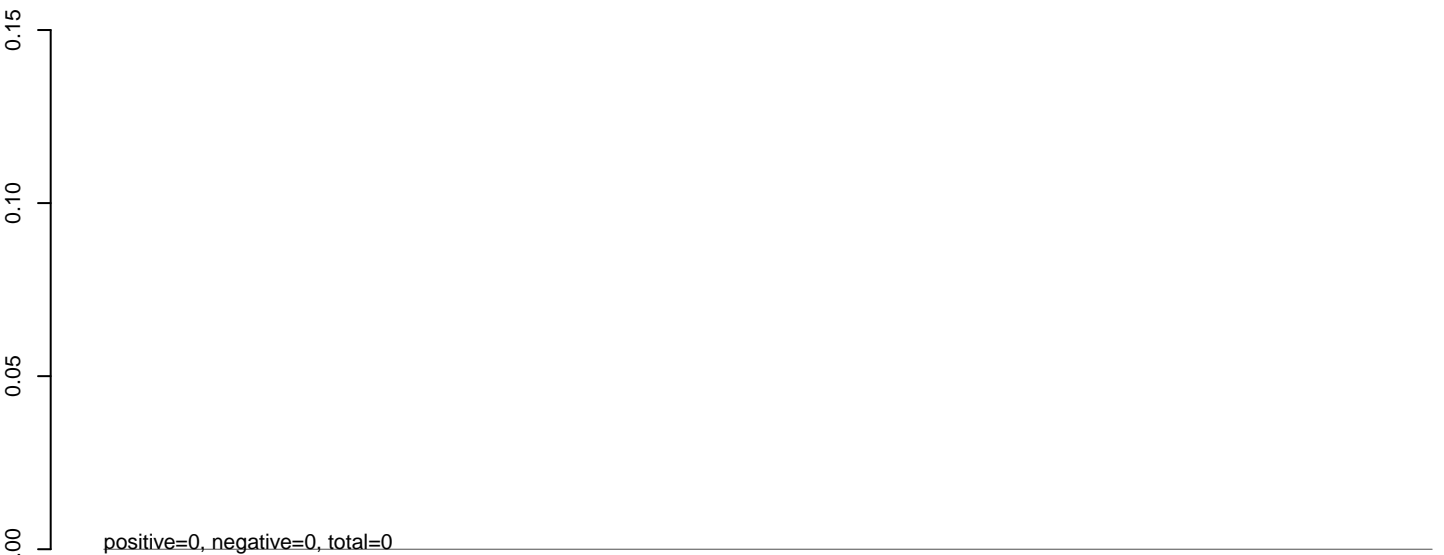




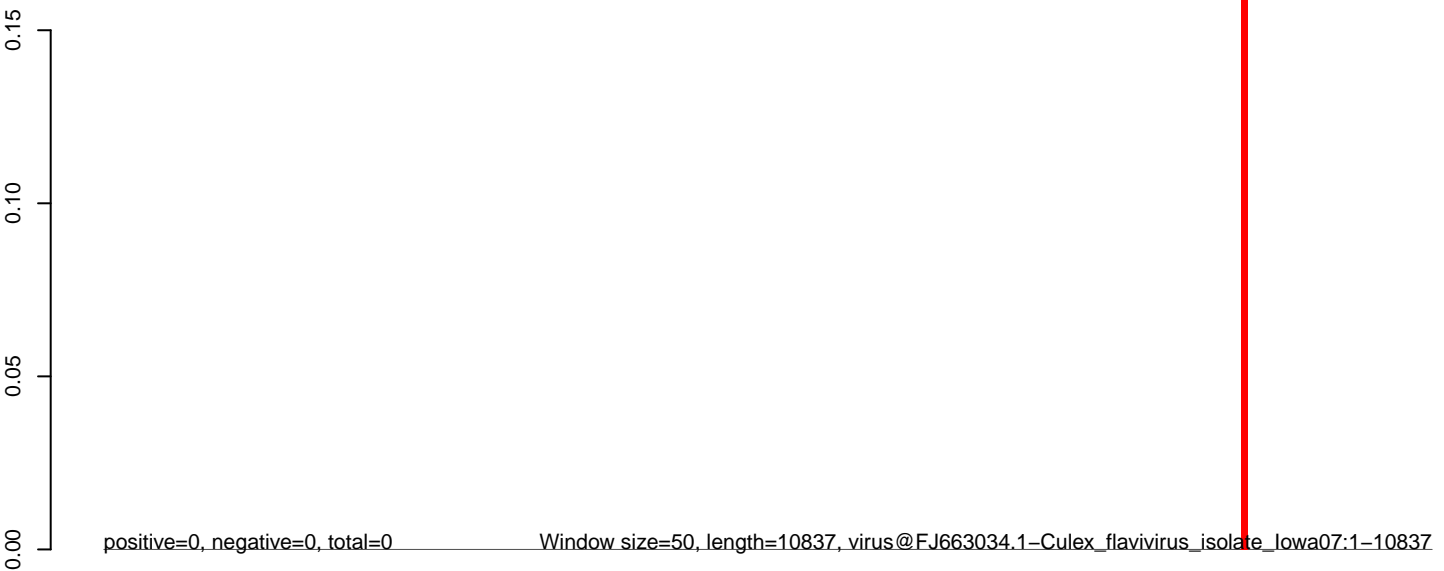
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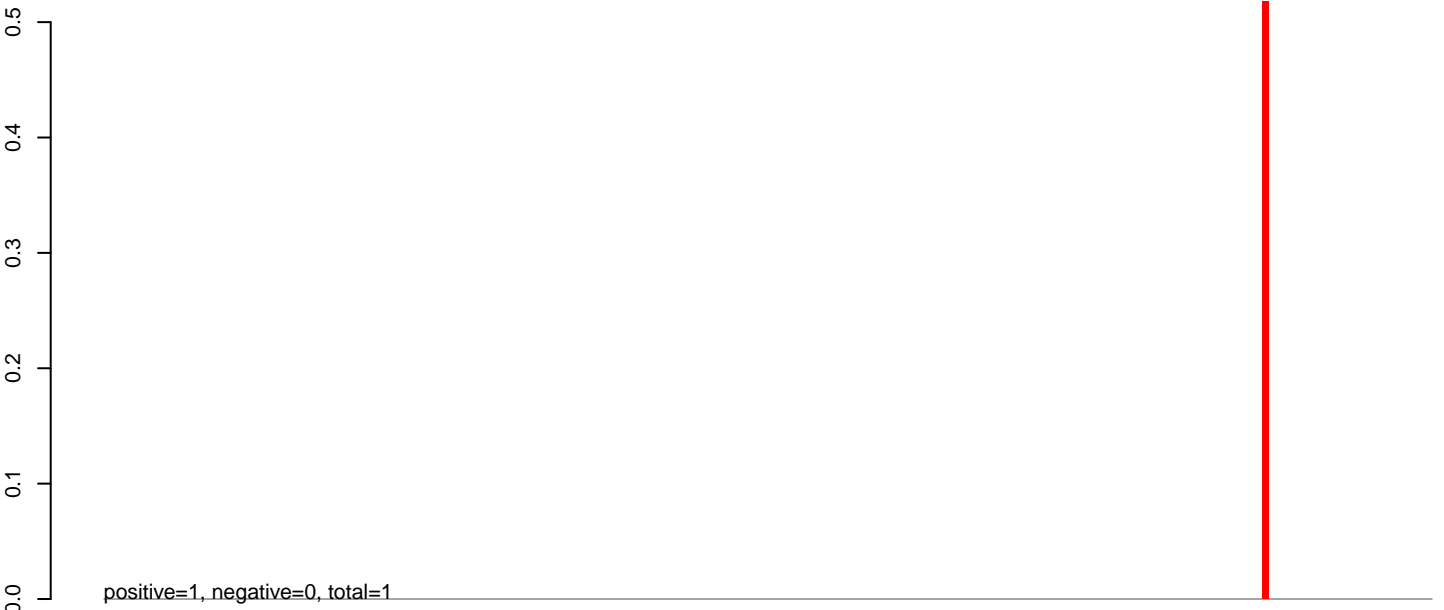


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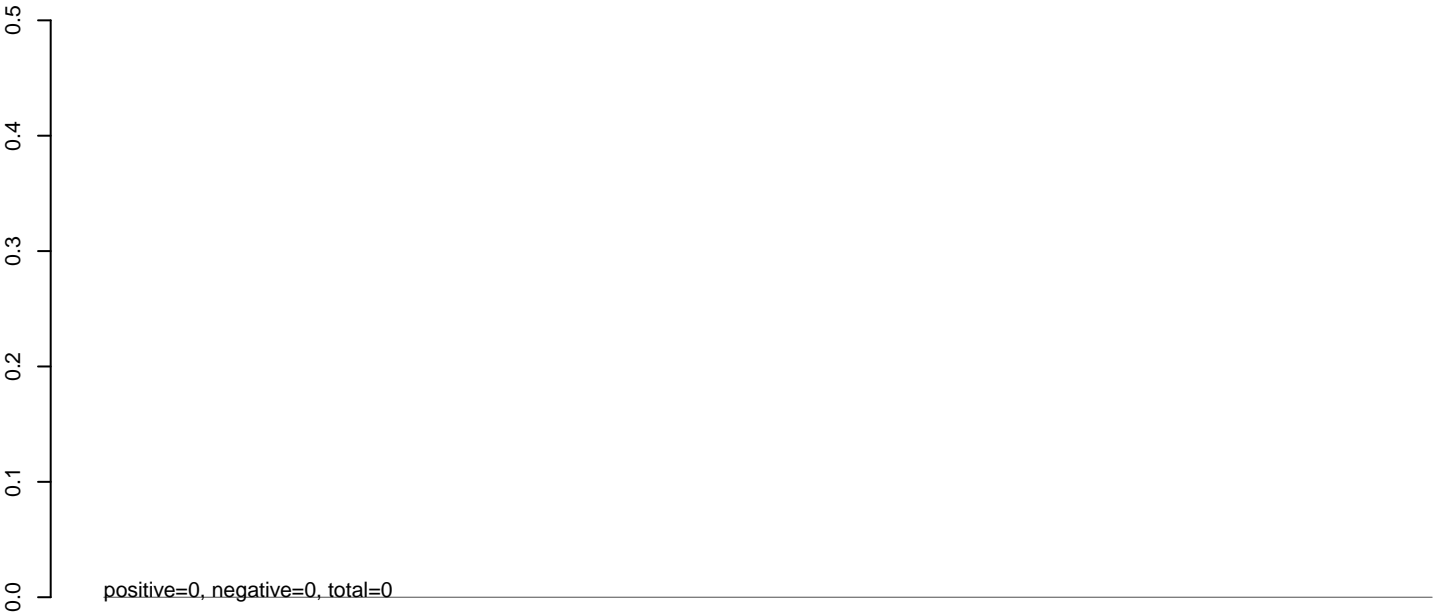


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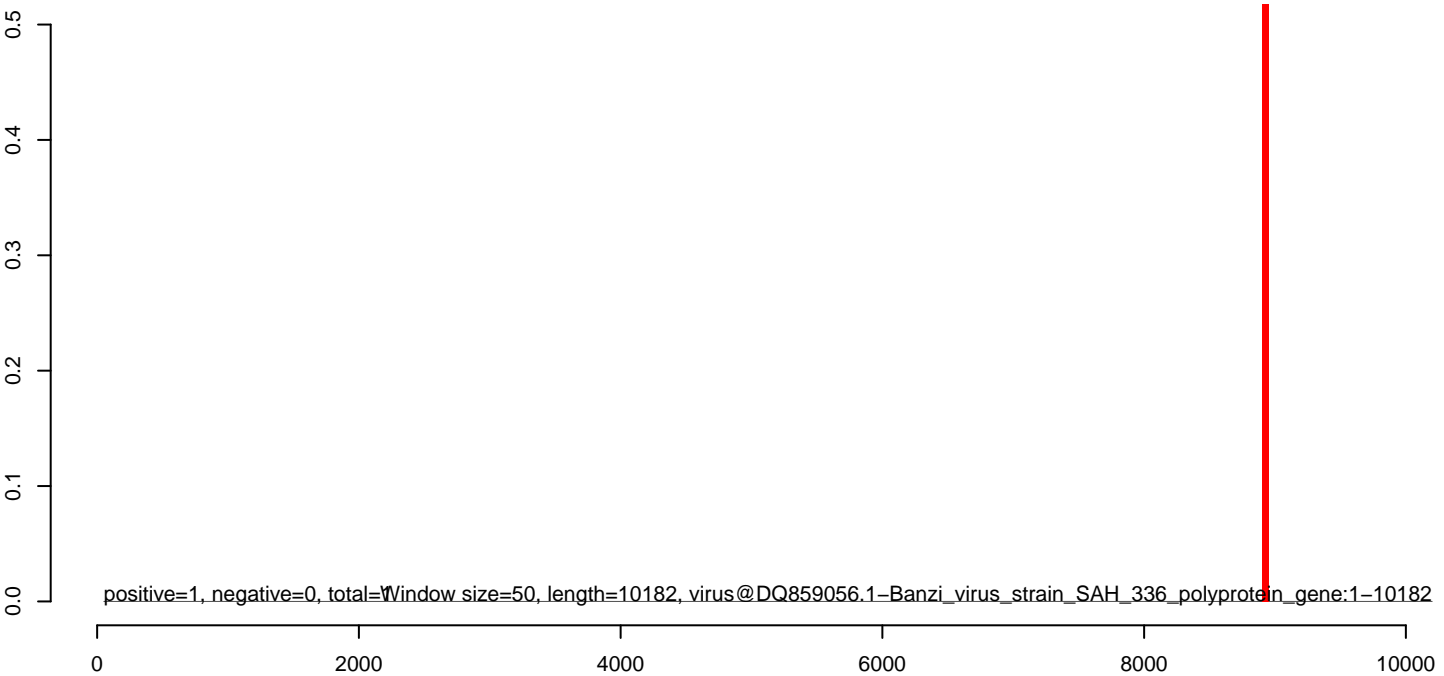
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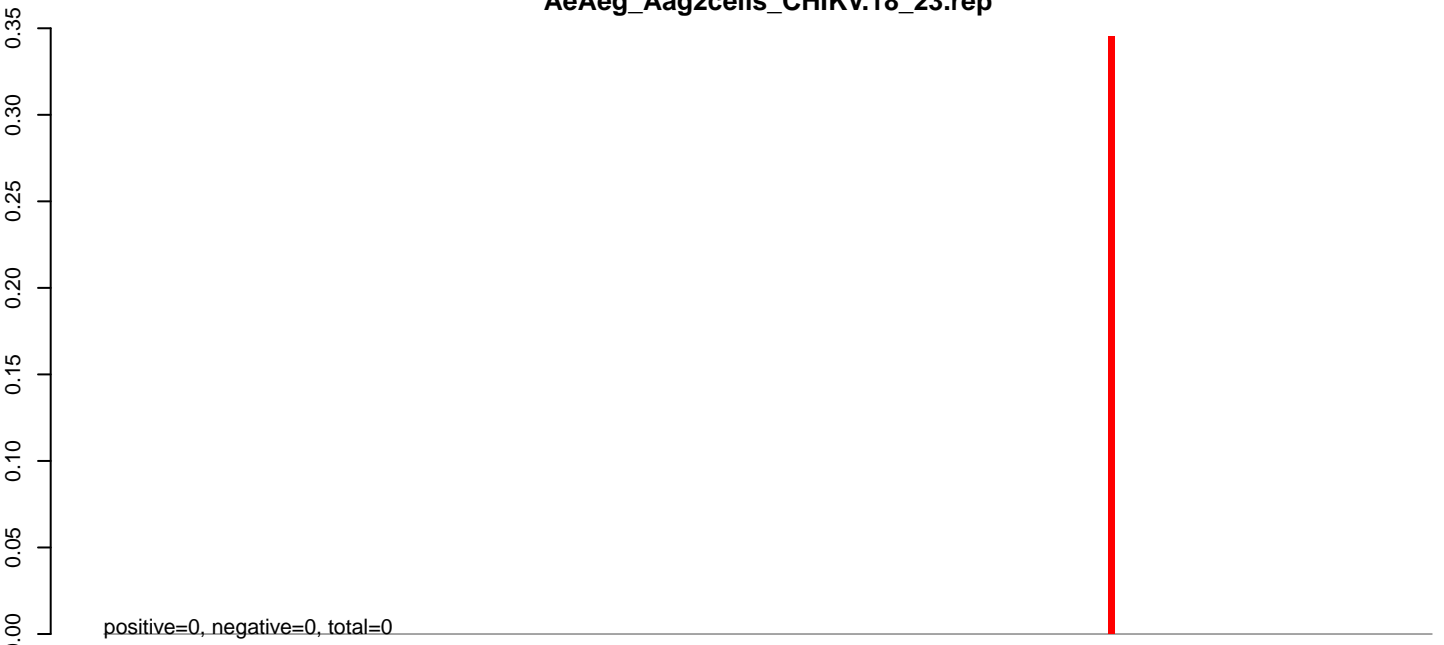
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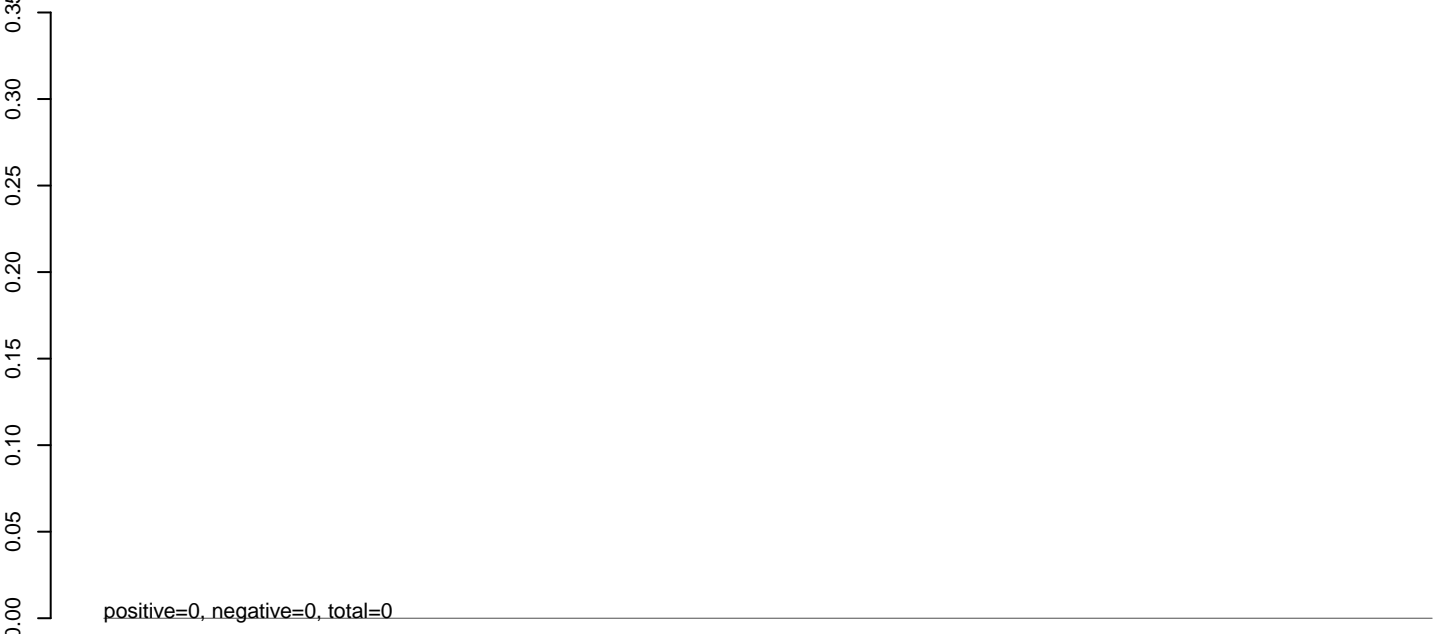
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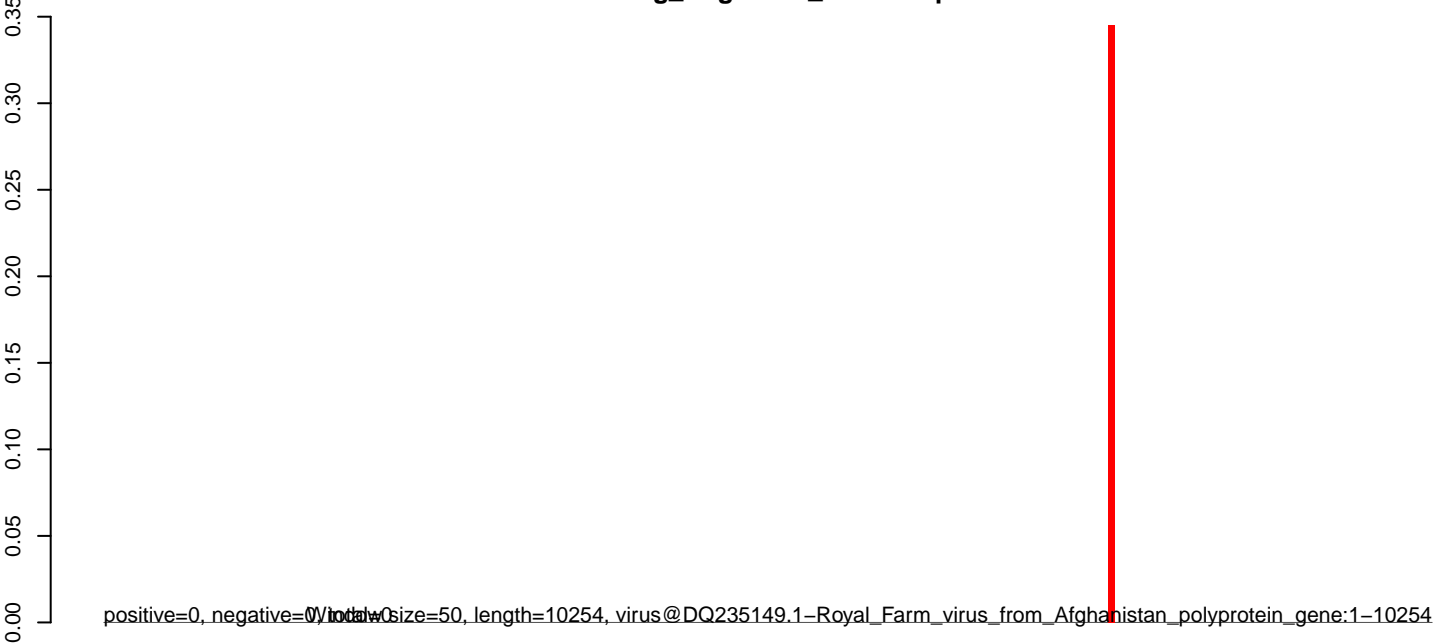
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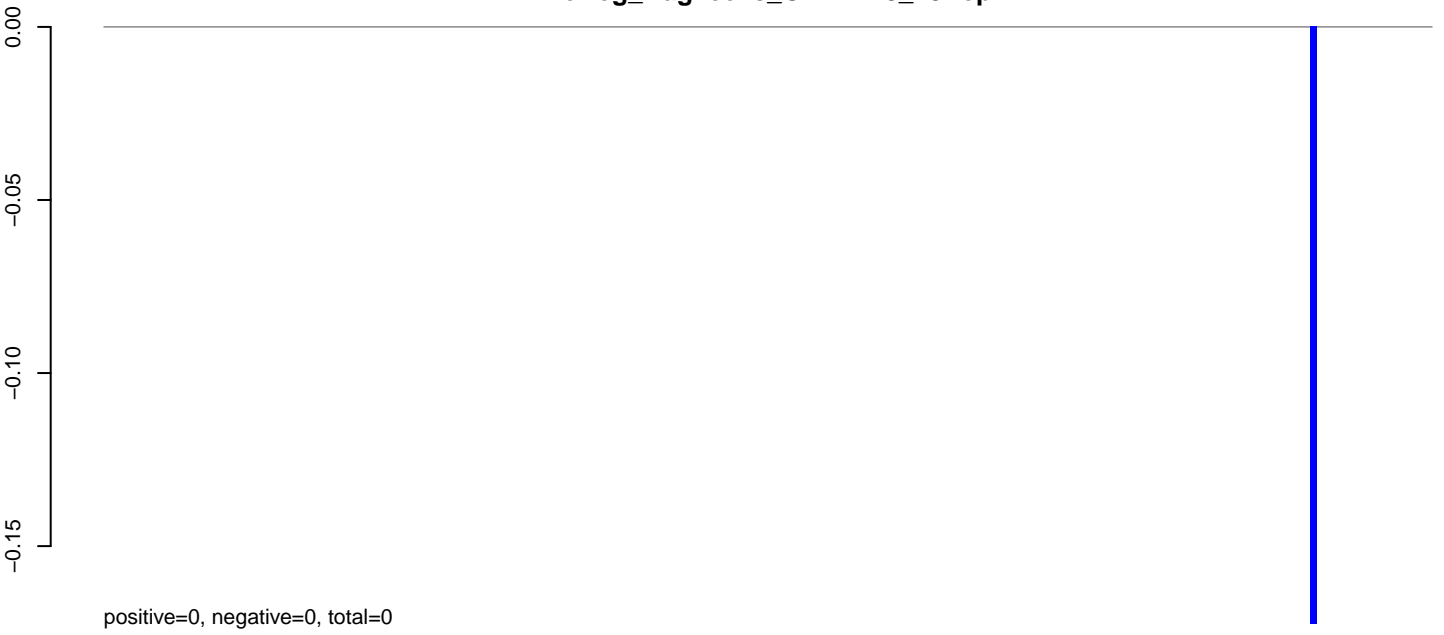
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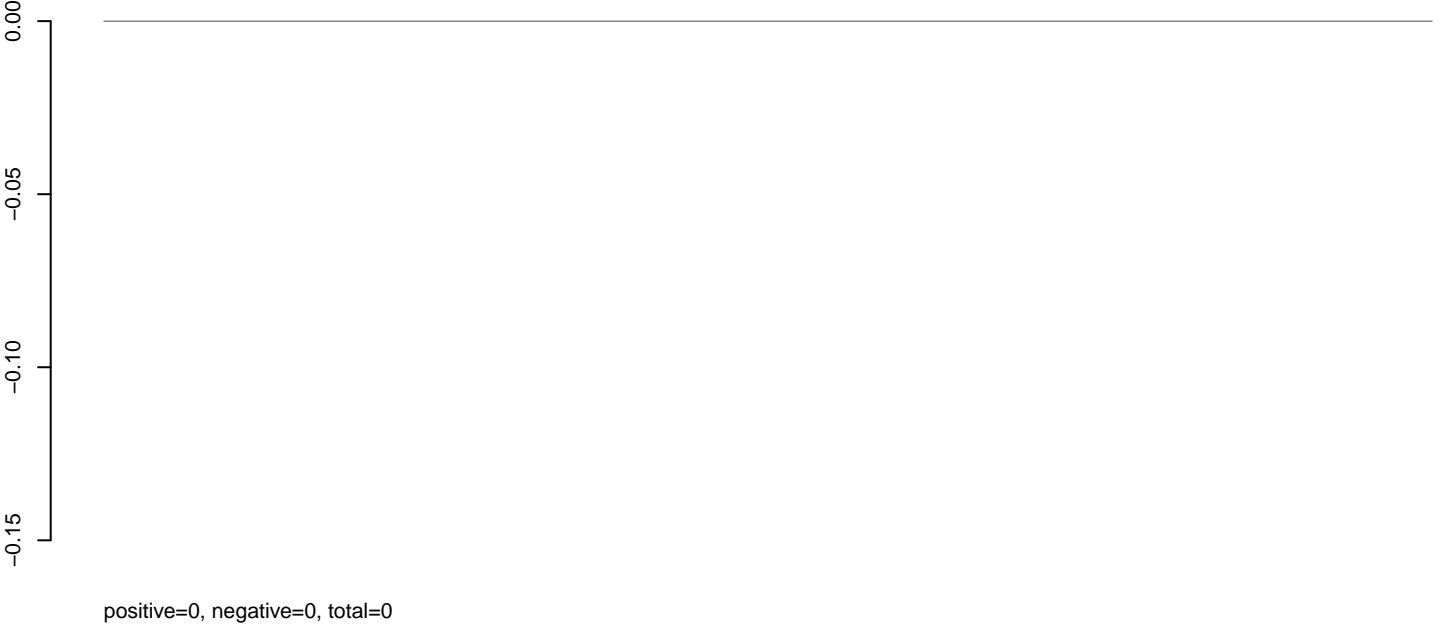
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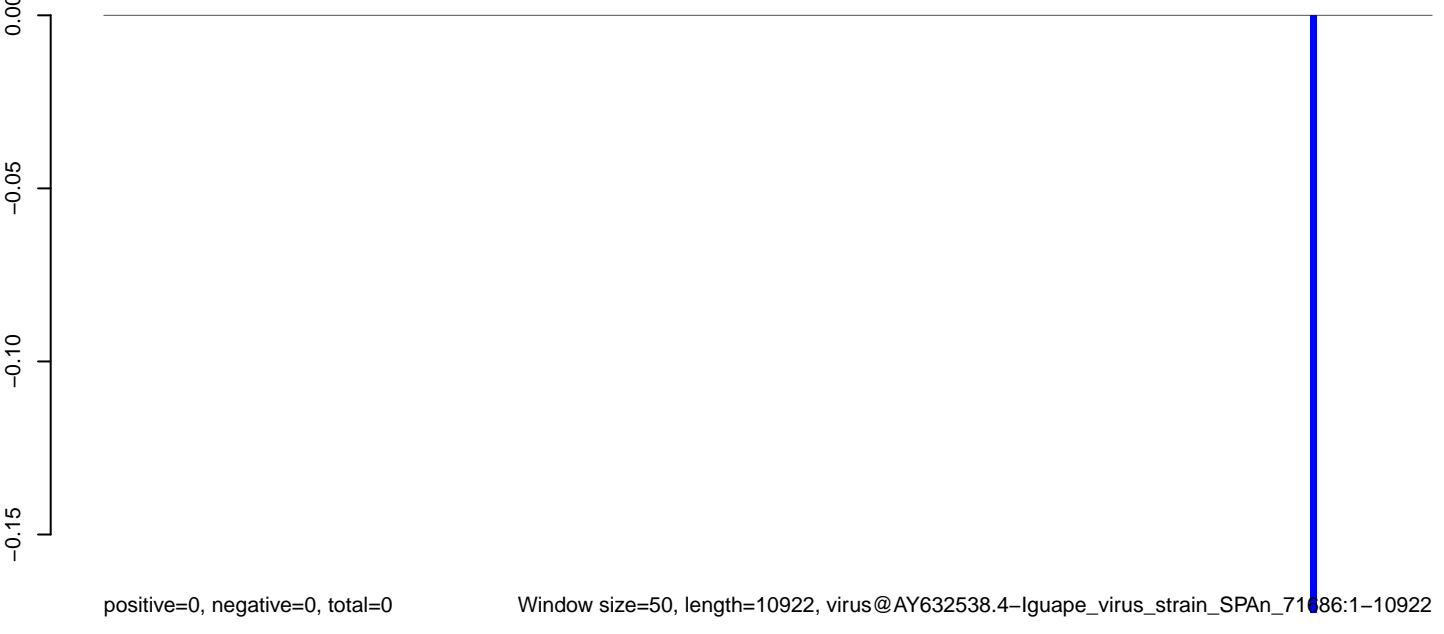
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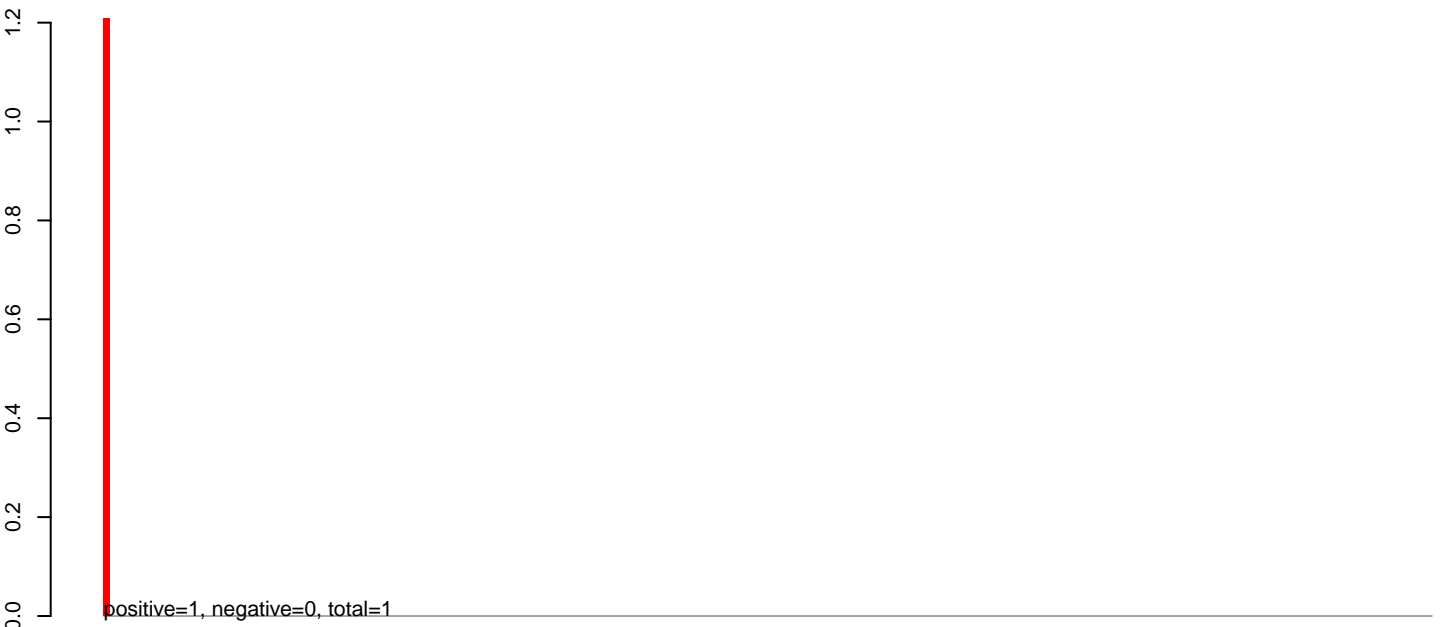
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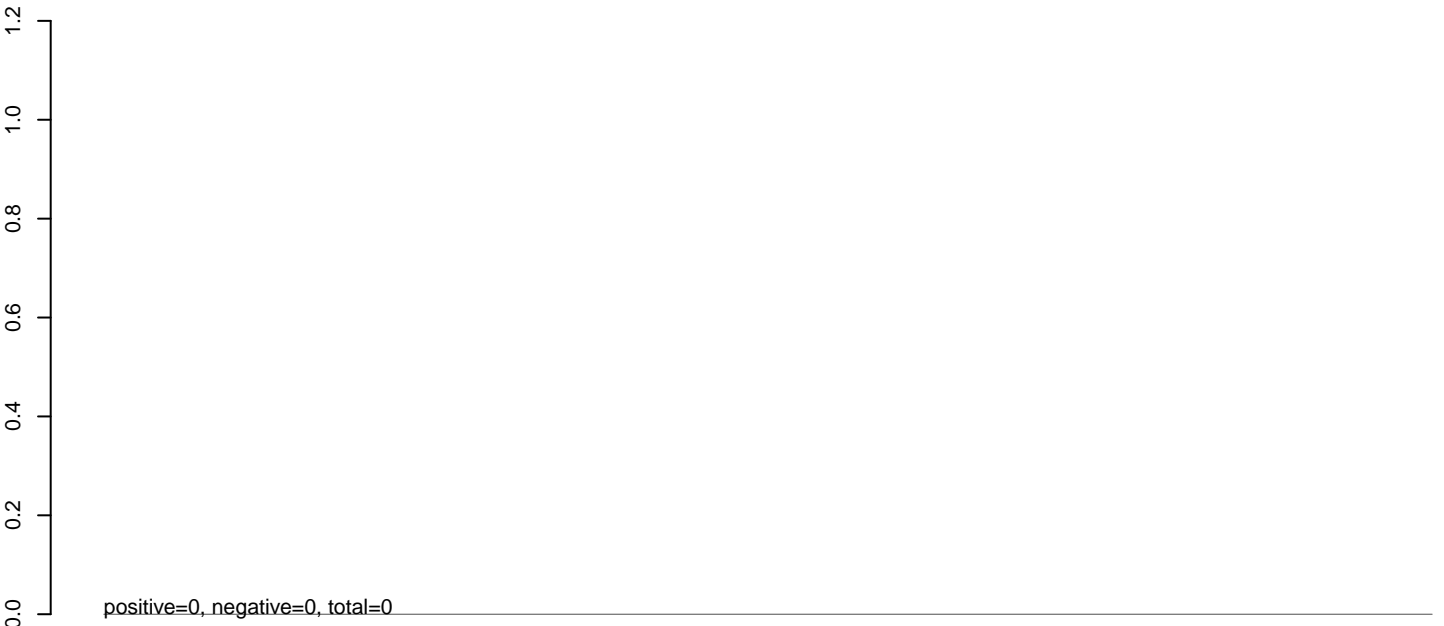
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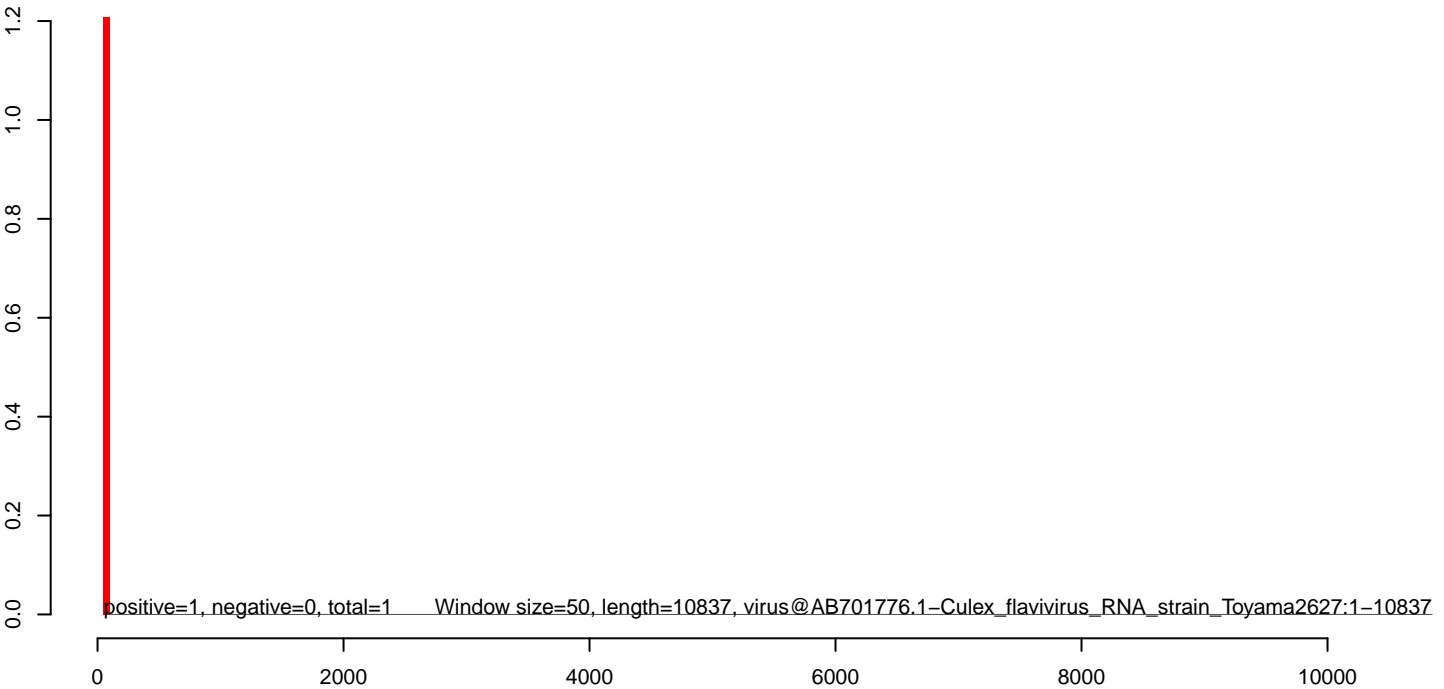
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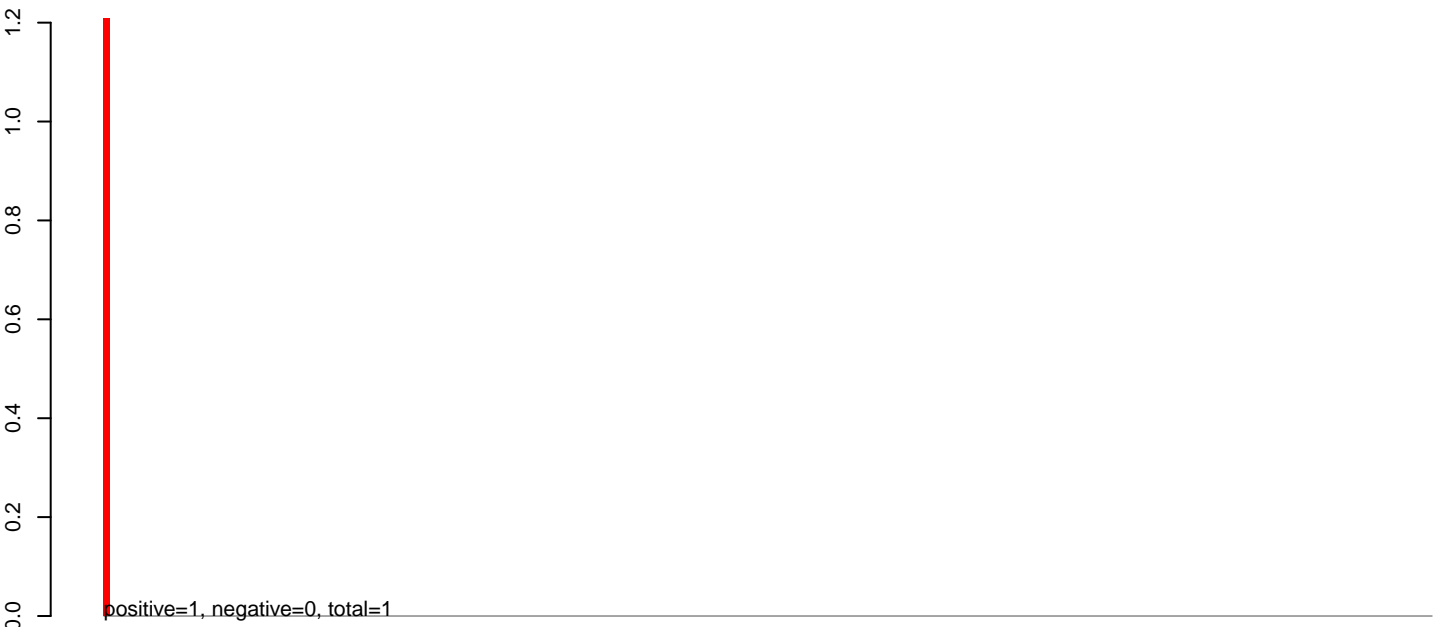
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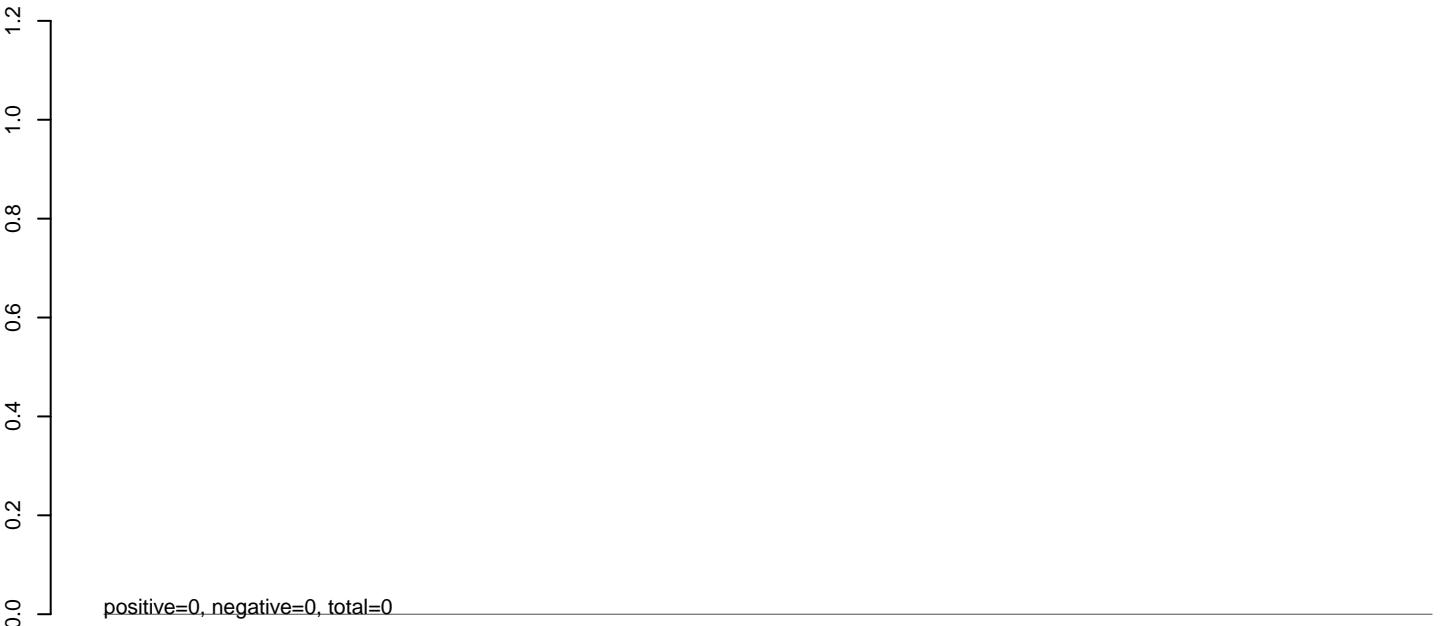
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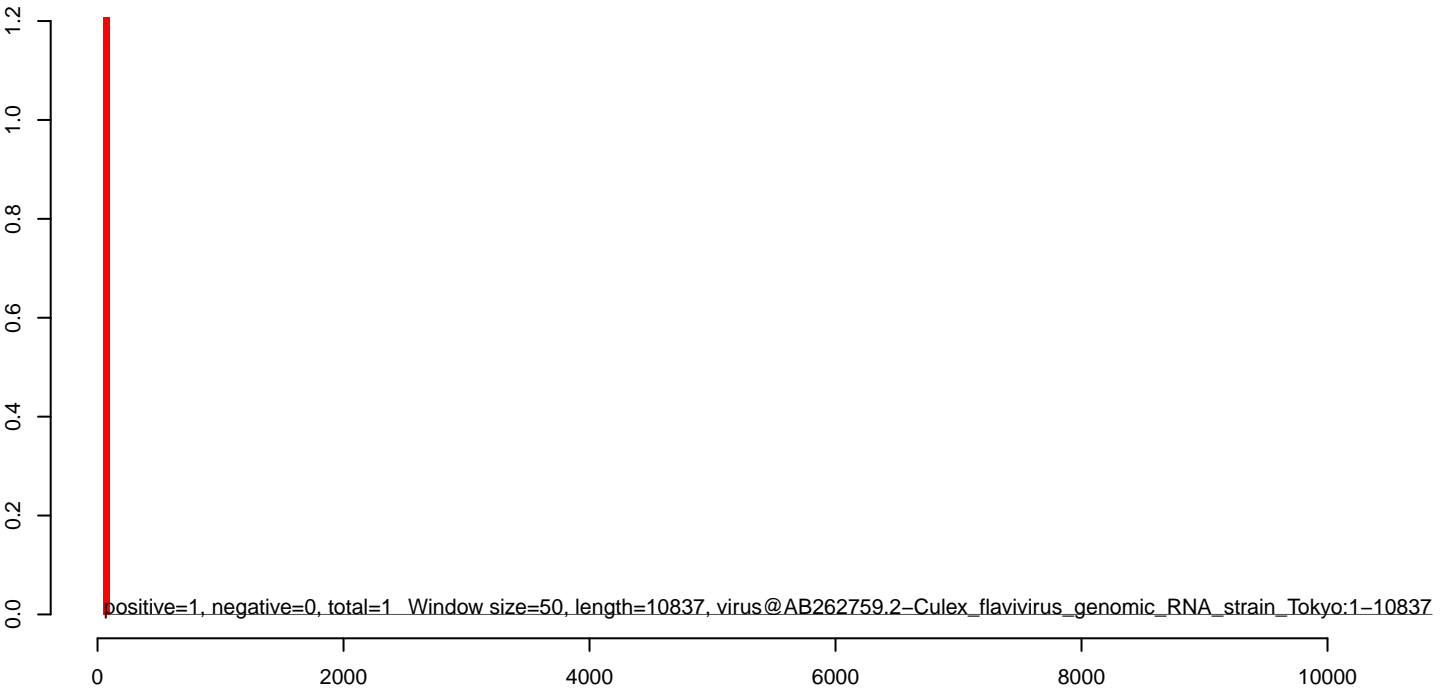
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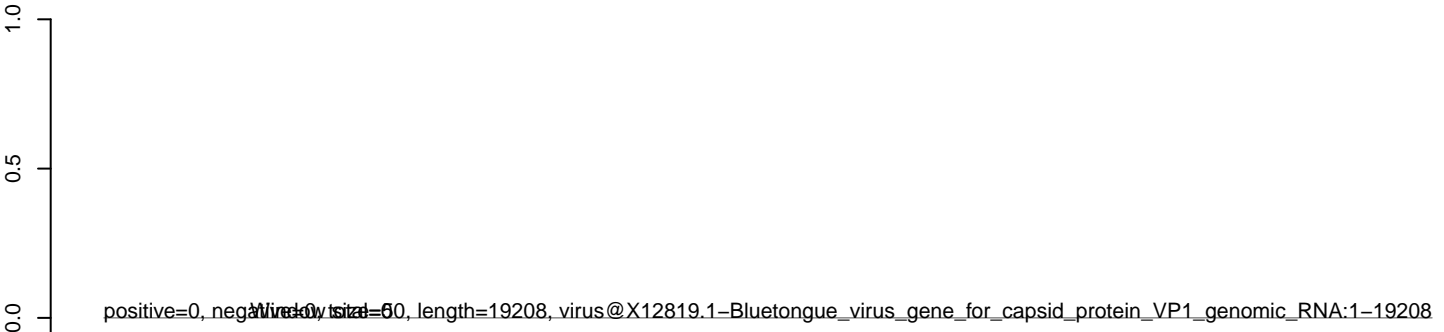
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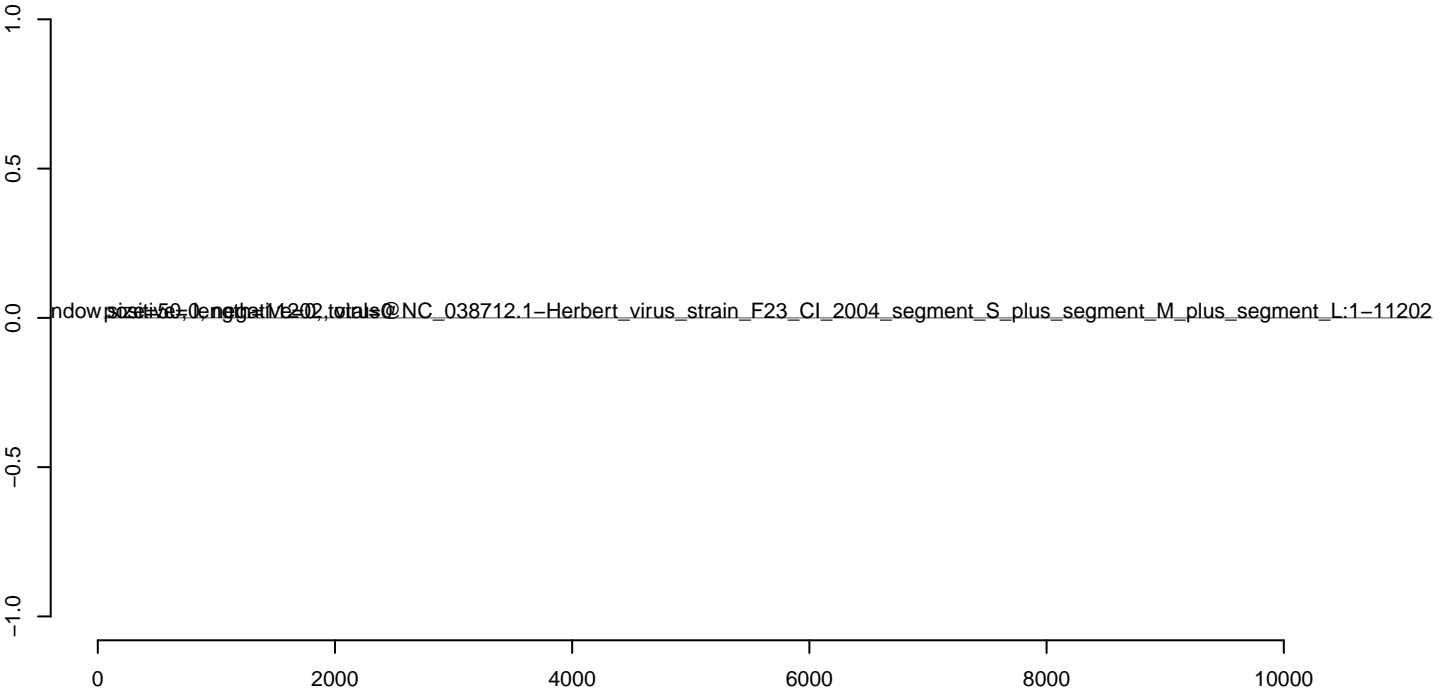
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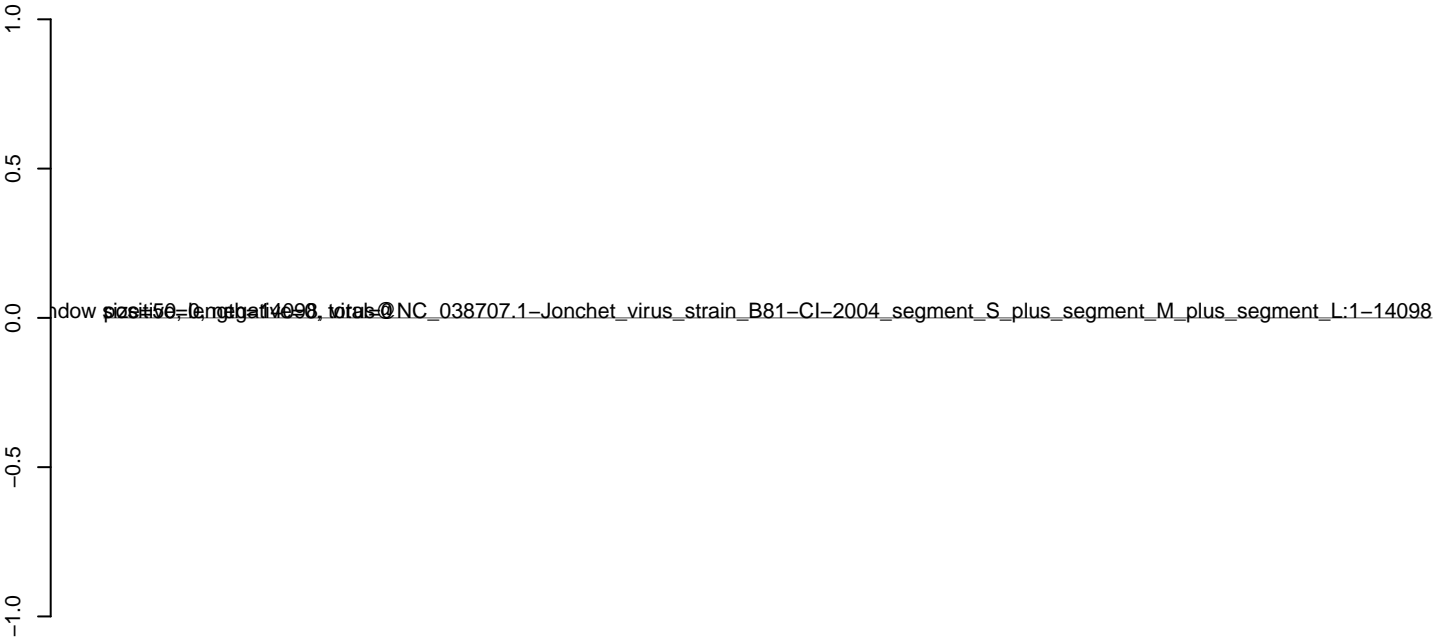
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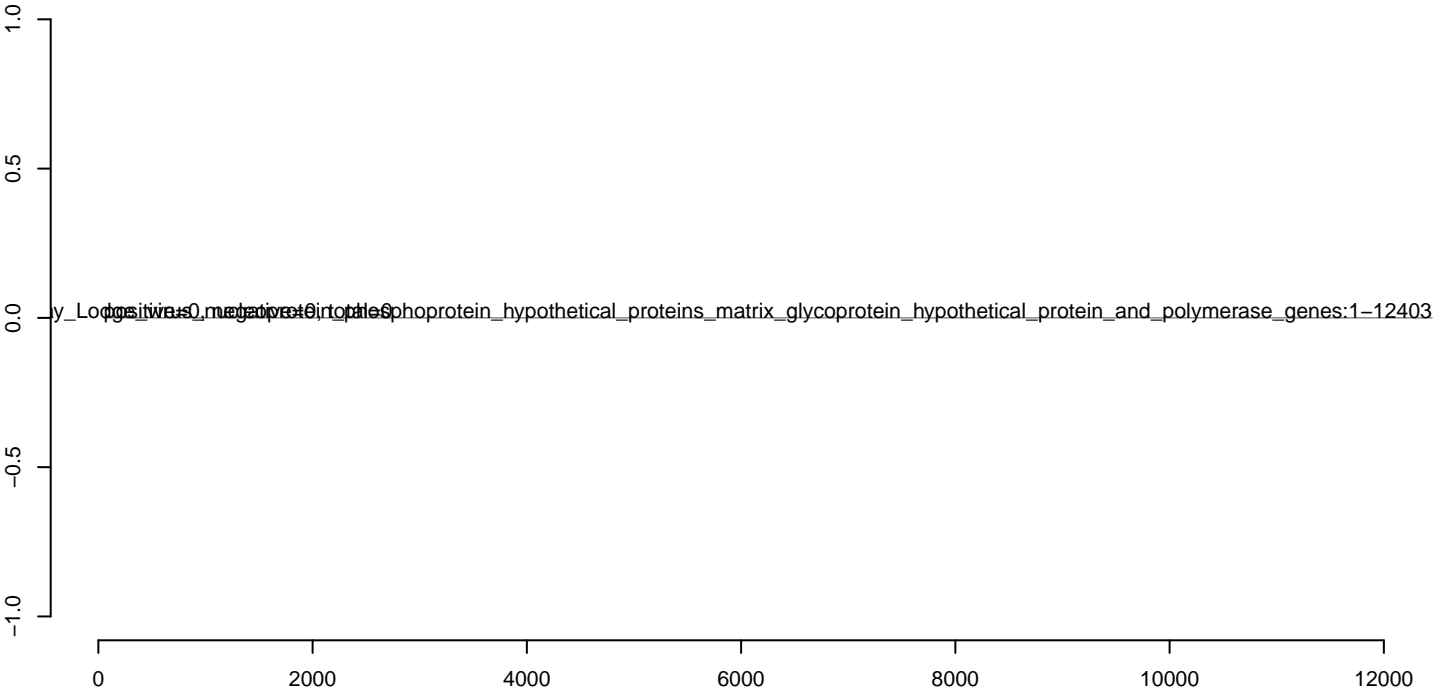
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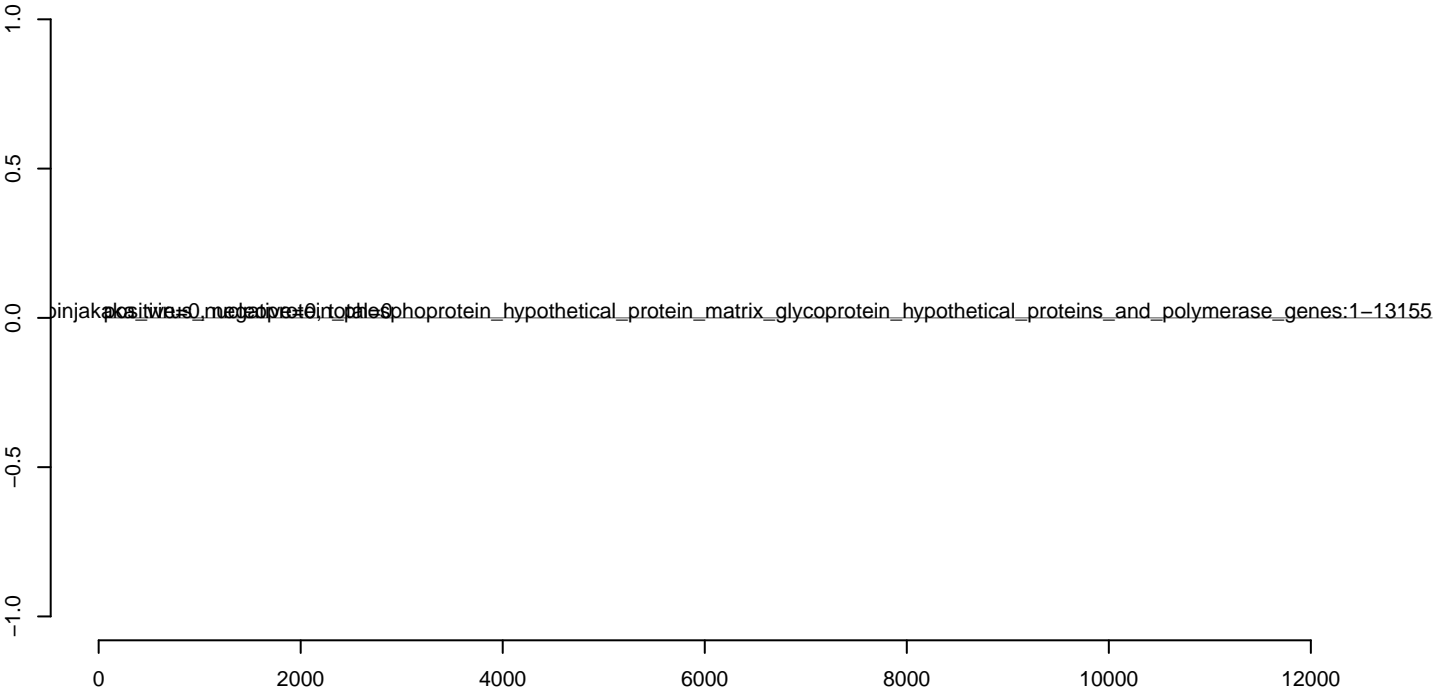
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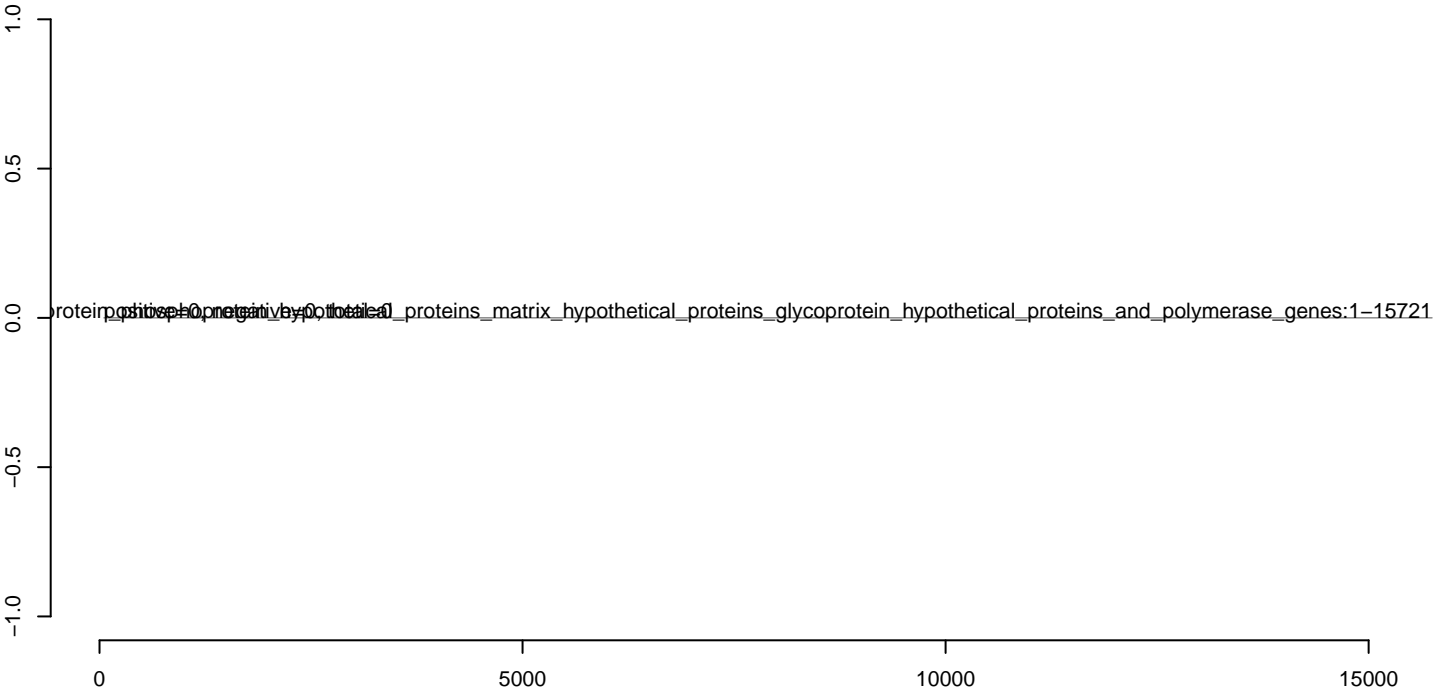
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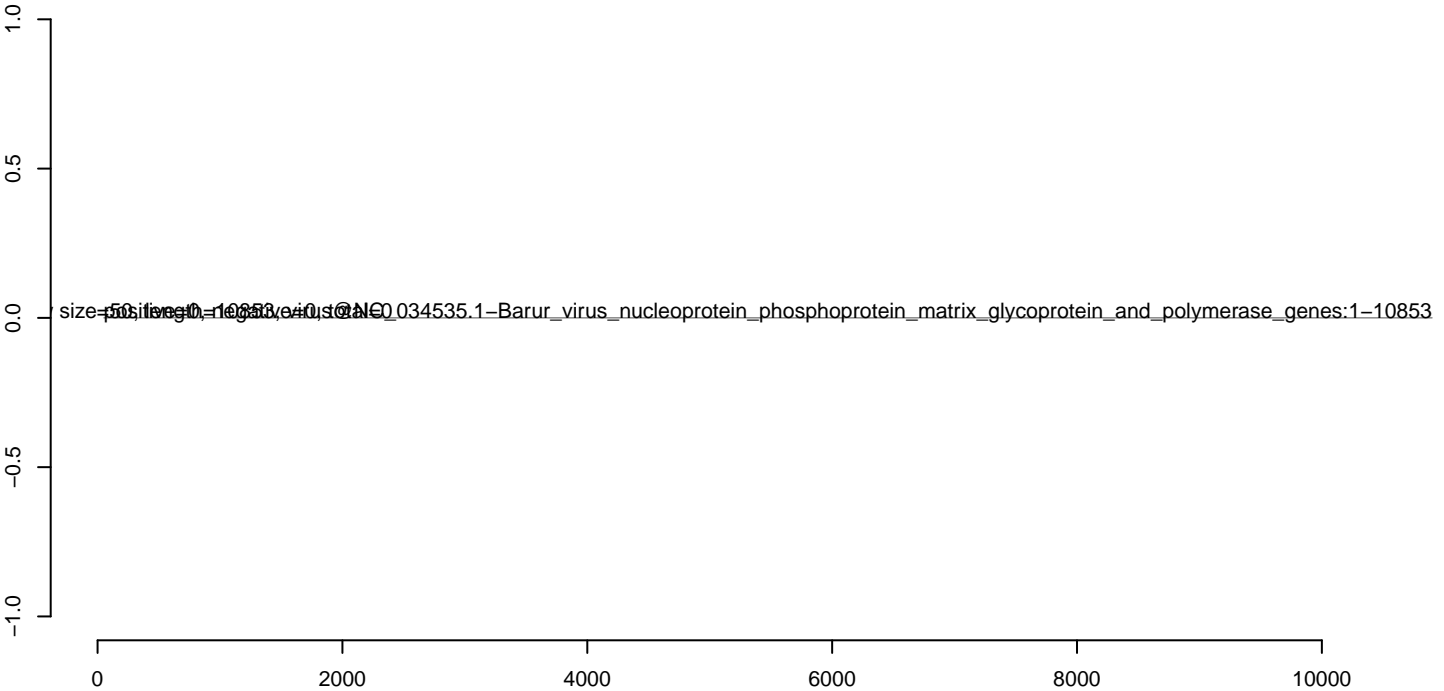
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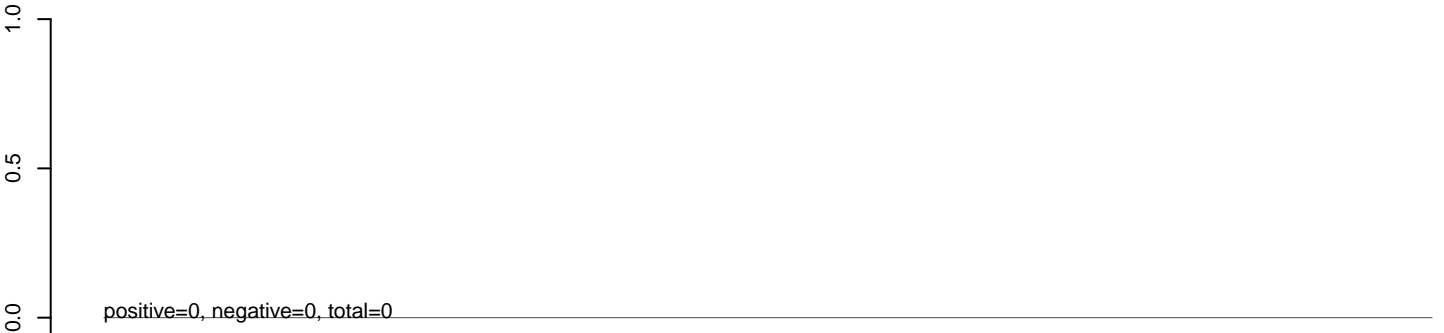
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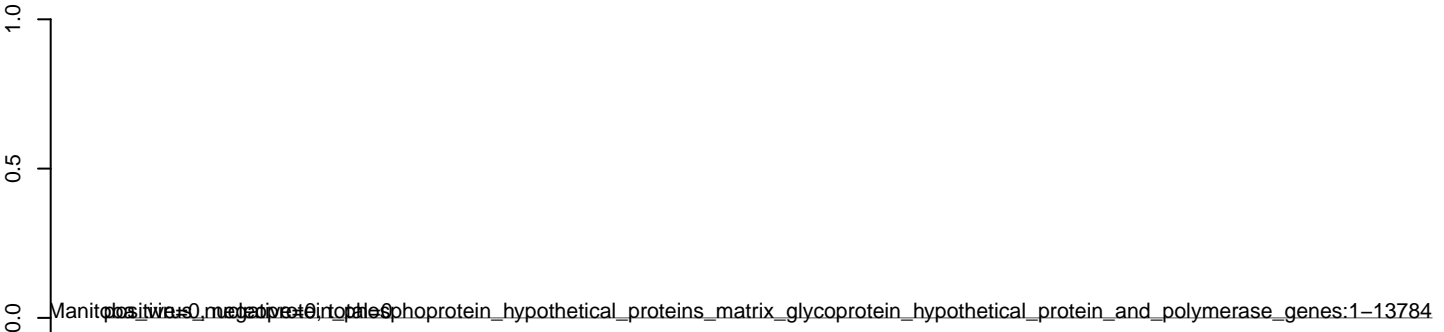
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AeAeg_Aag2cells_CHIKV.rep



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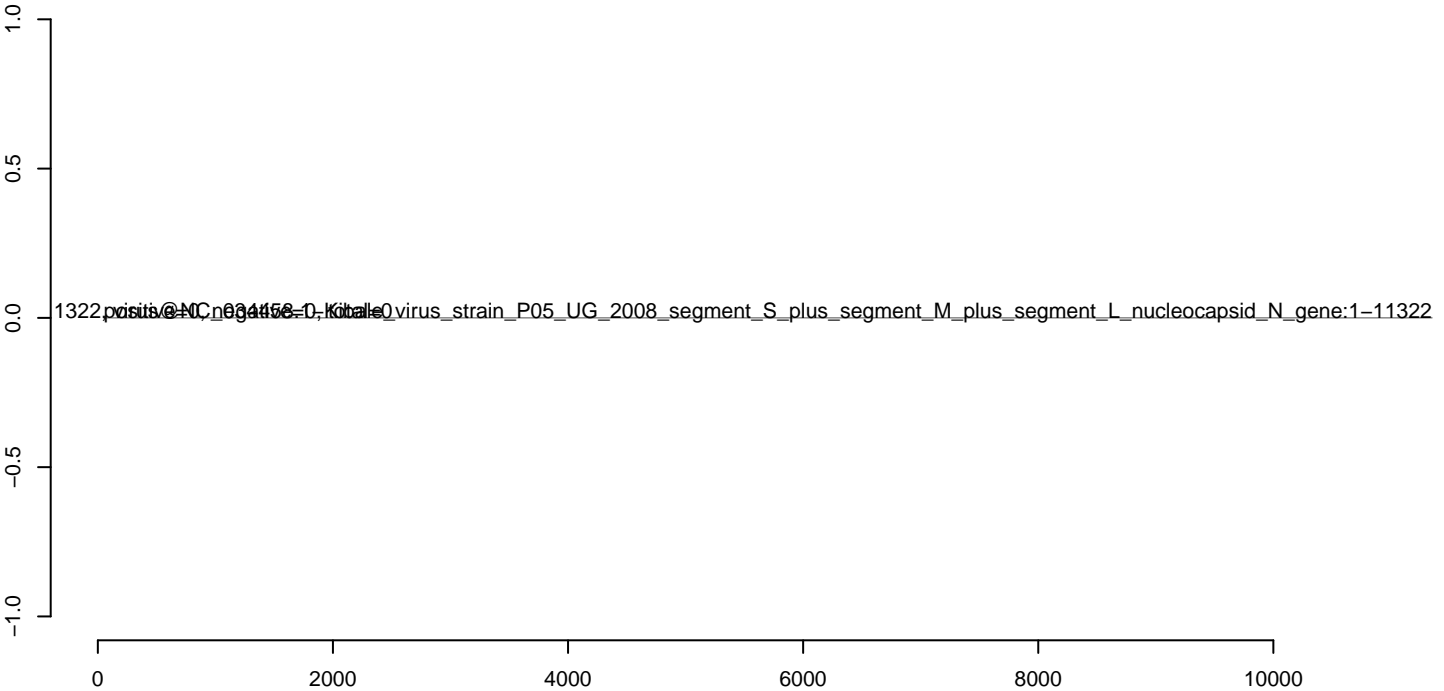
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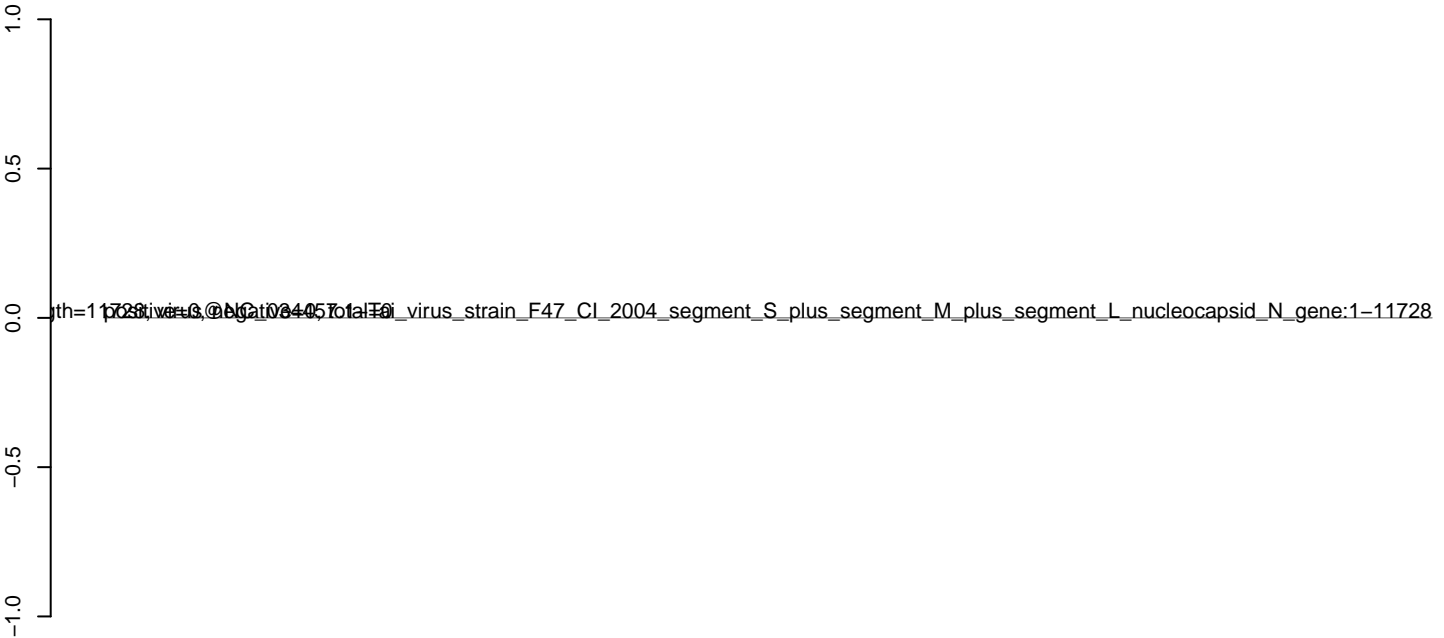
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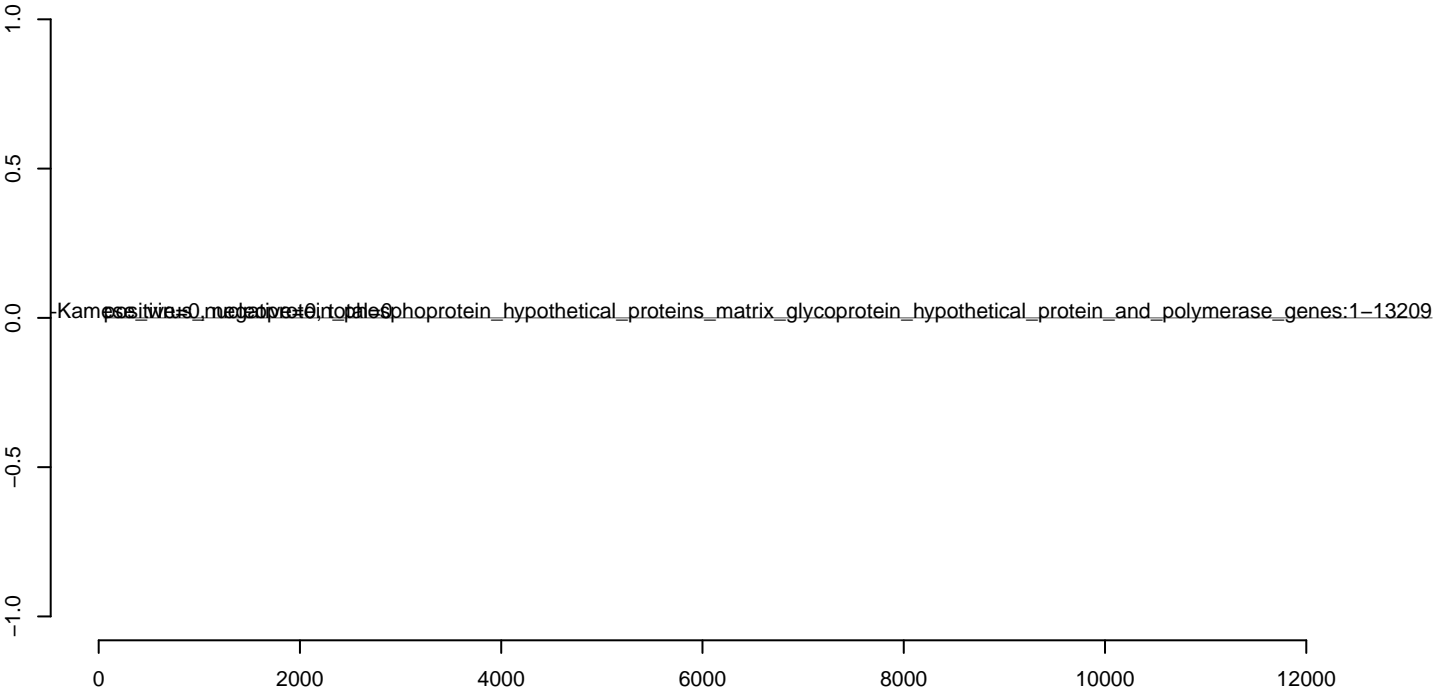
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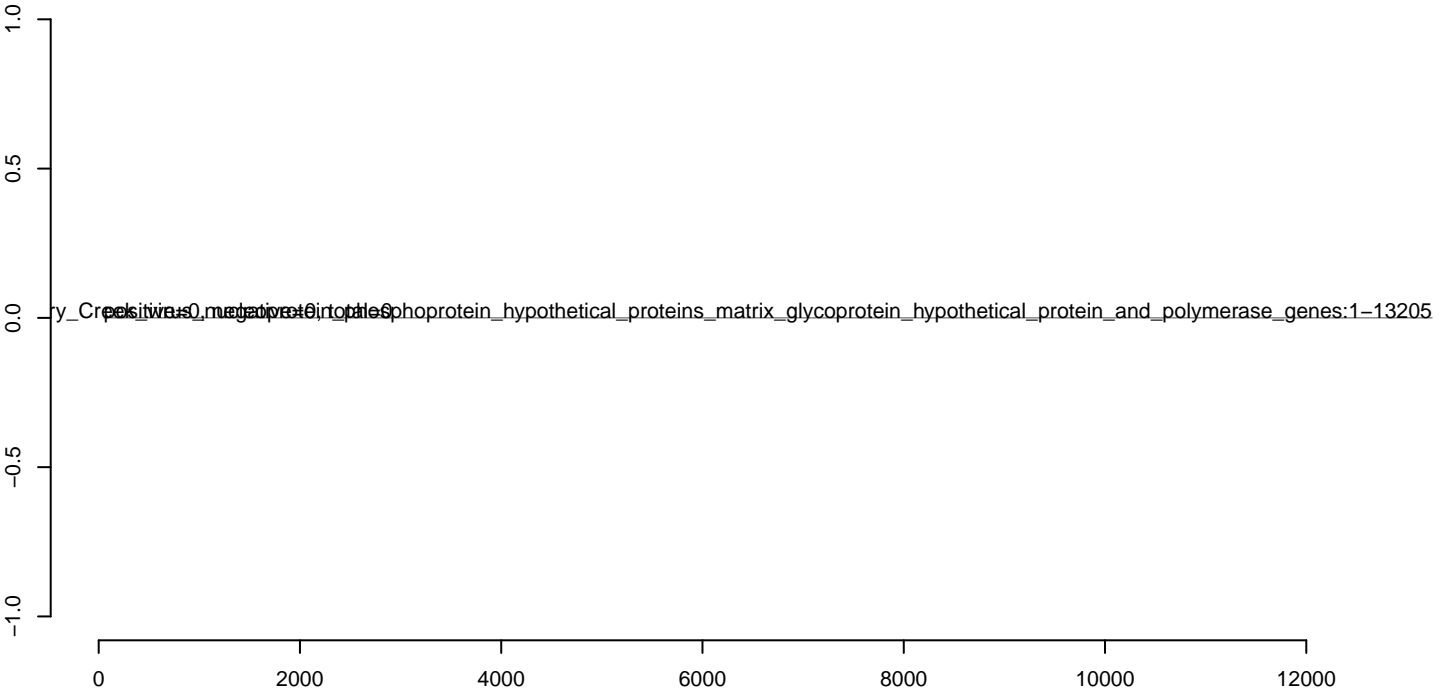
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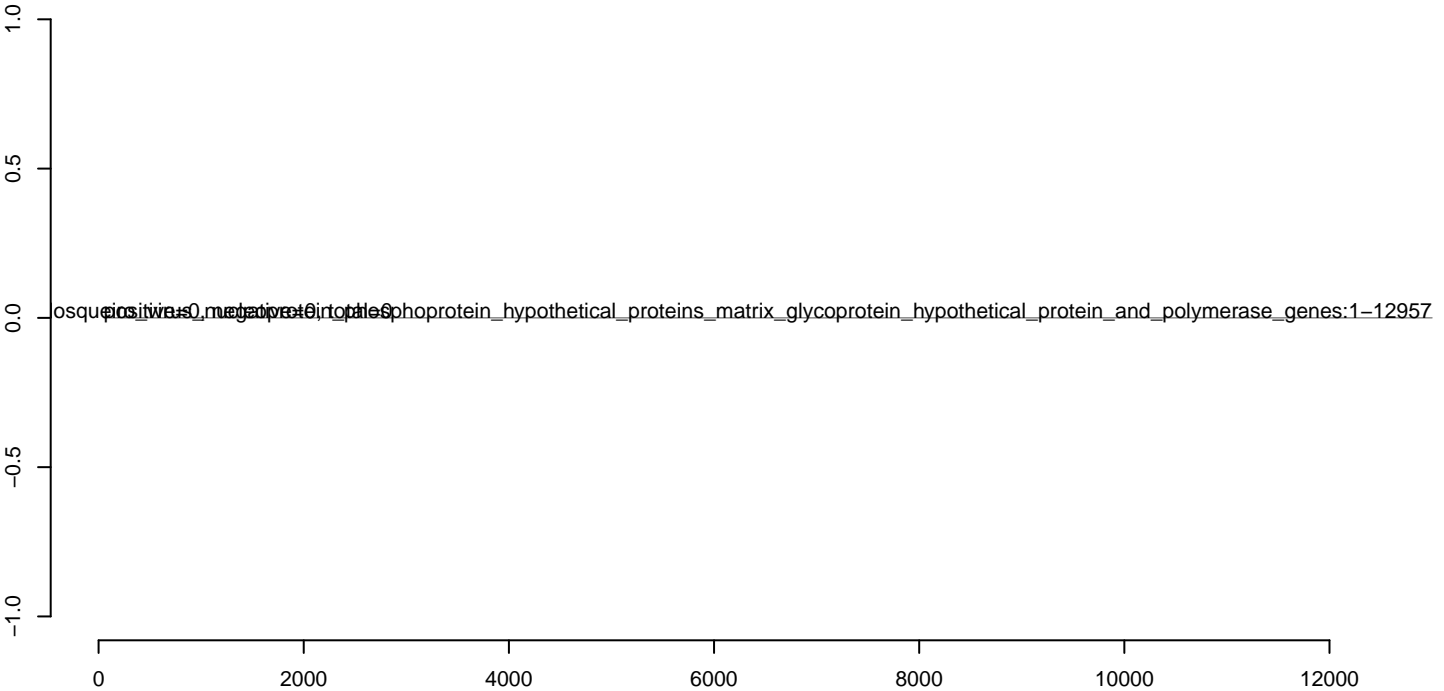
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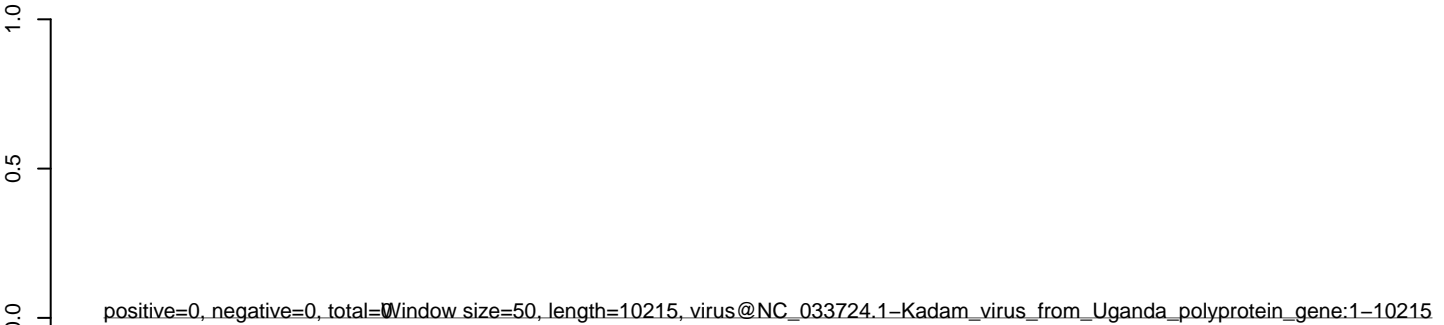
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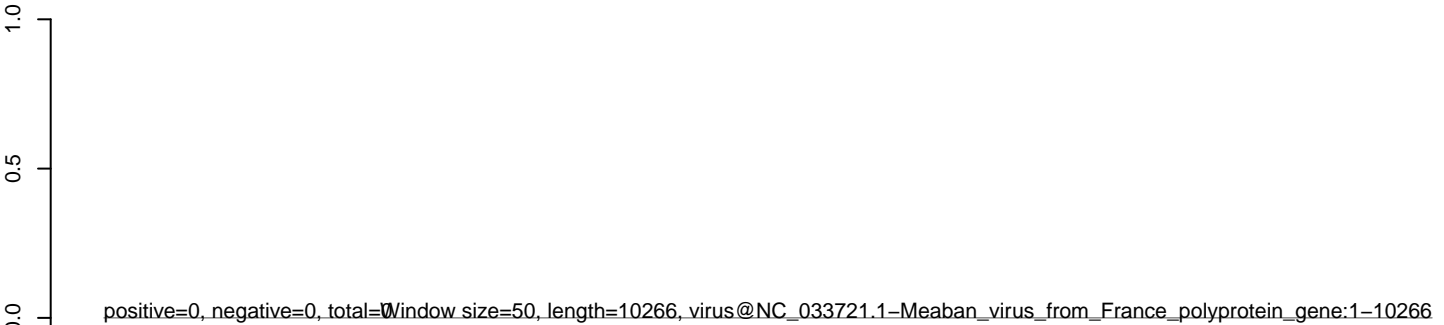
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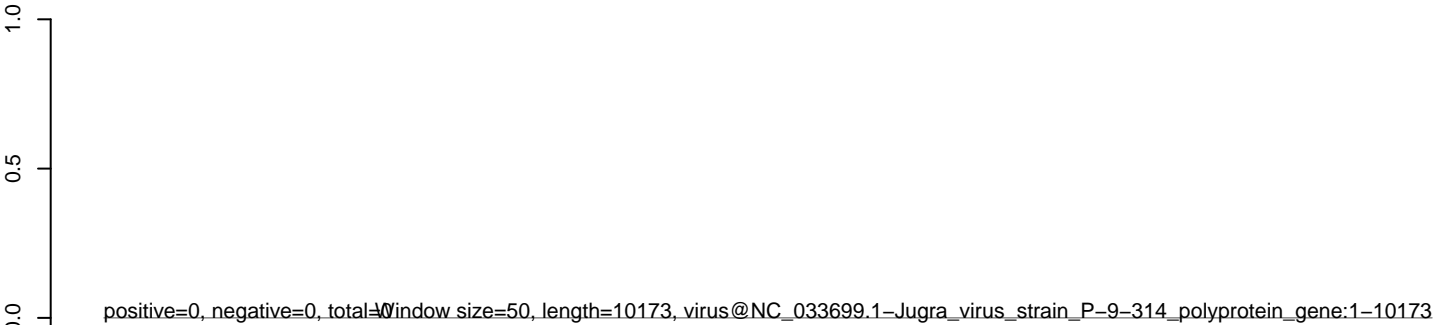
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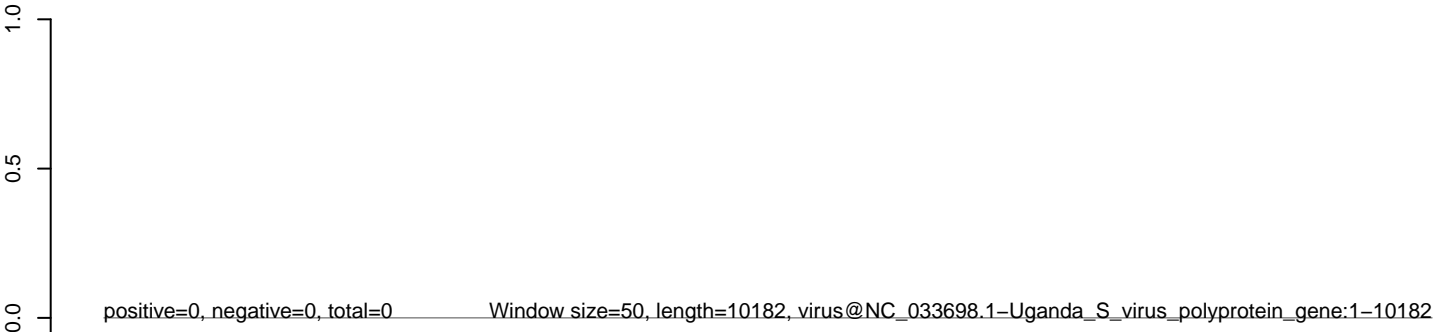
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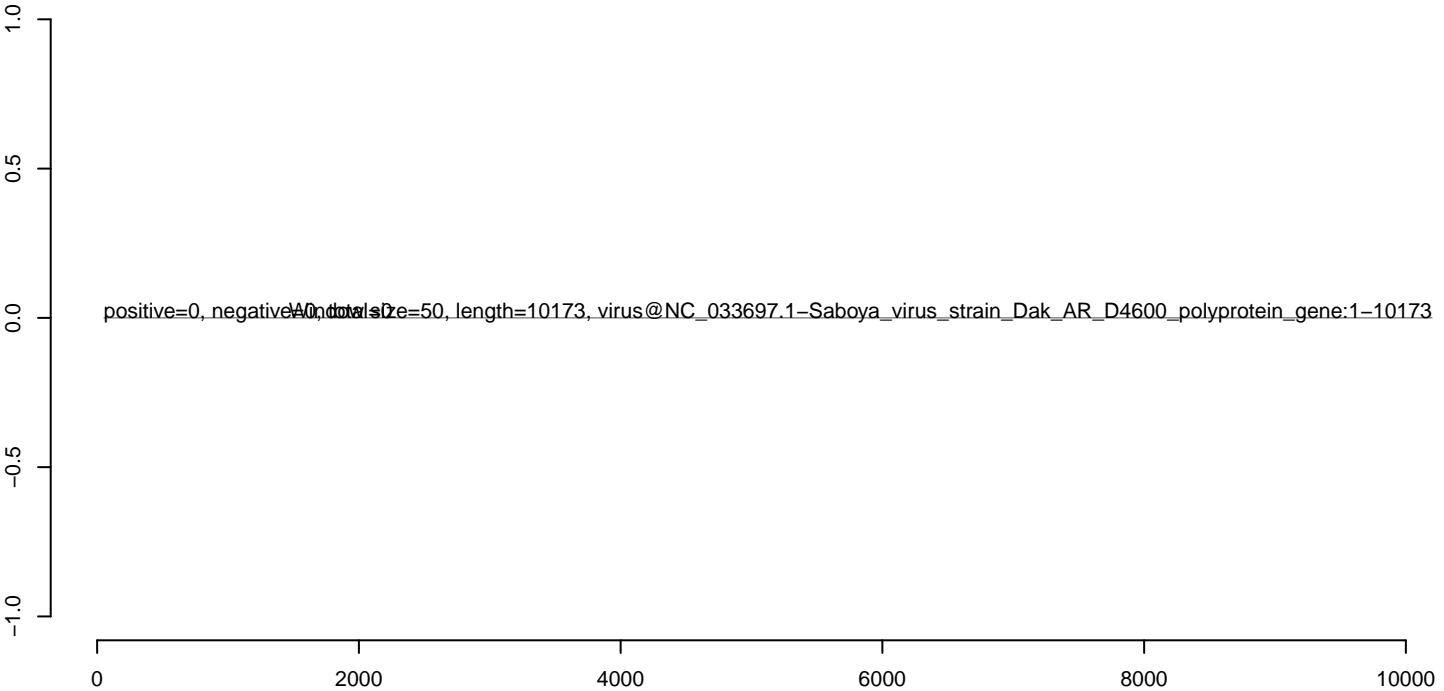
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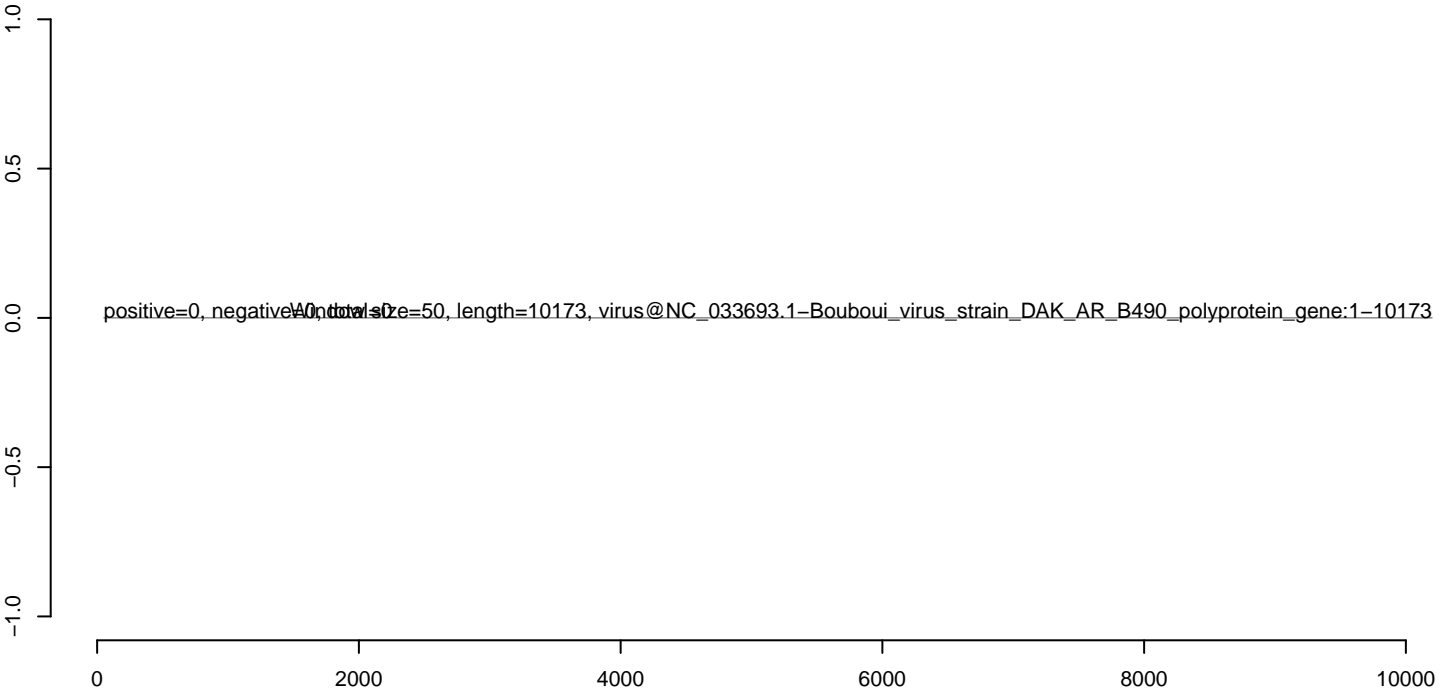
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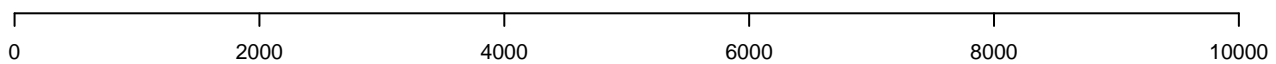
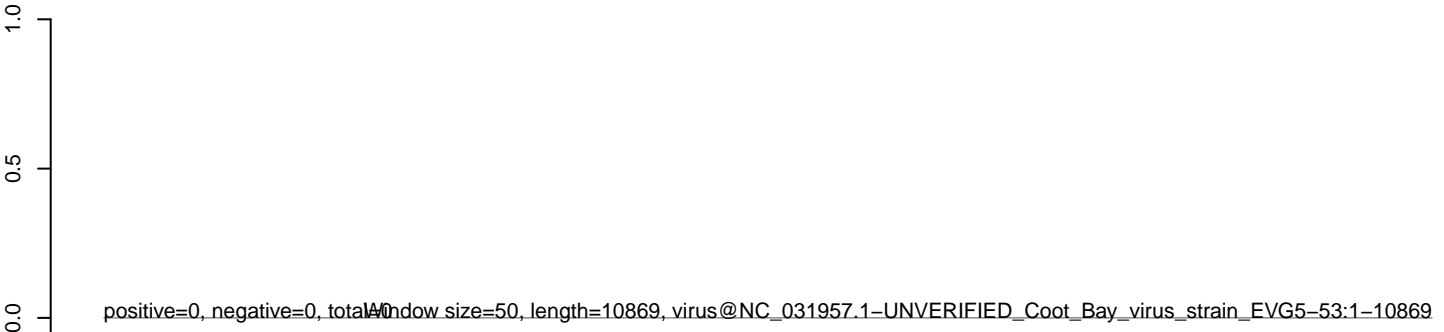
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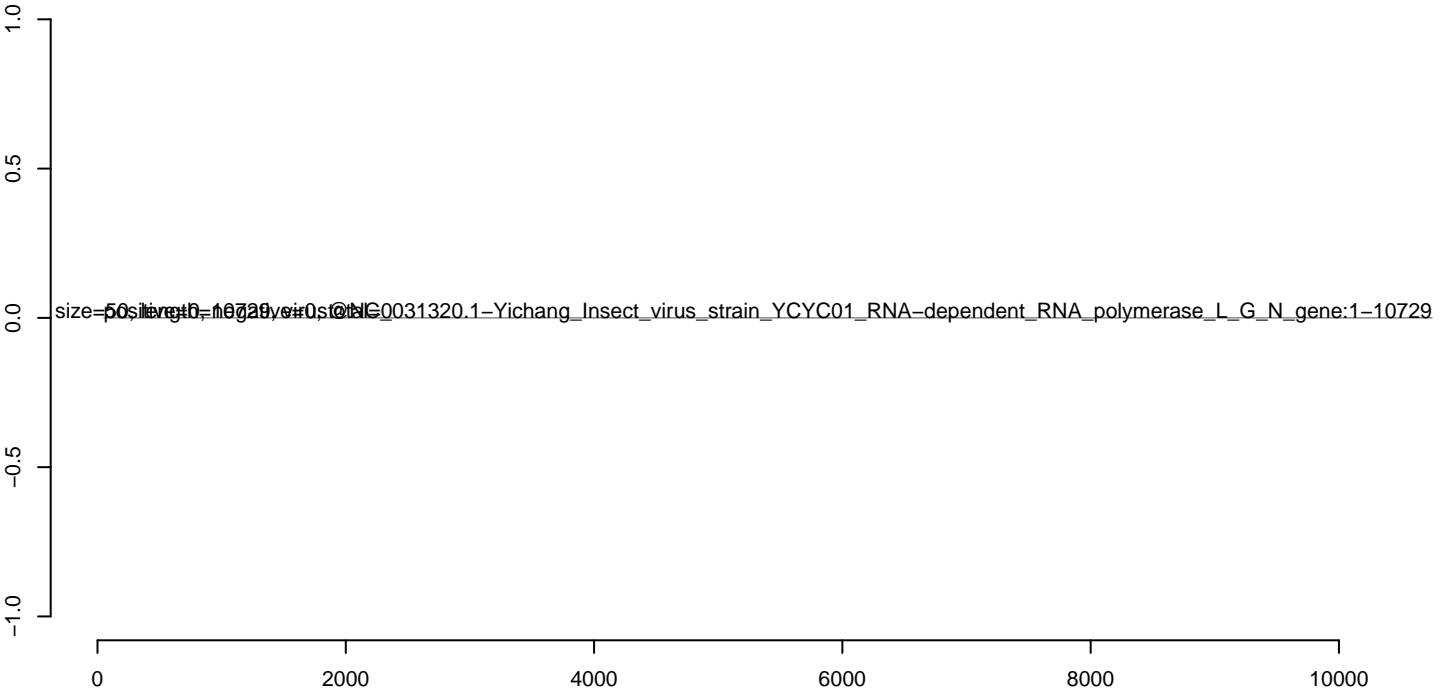
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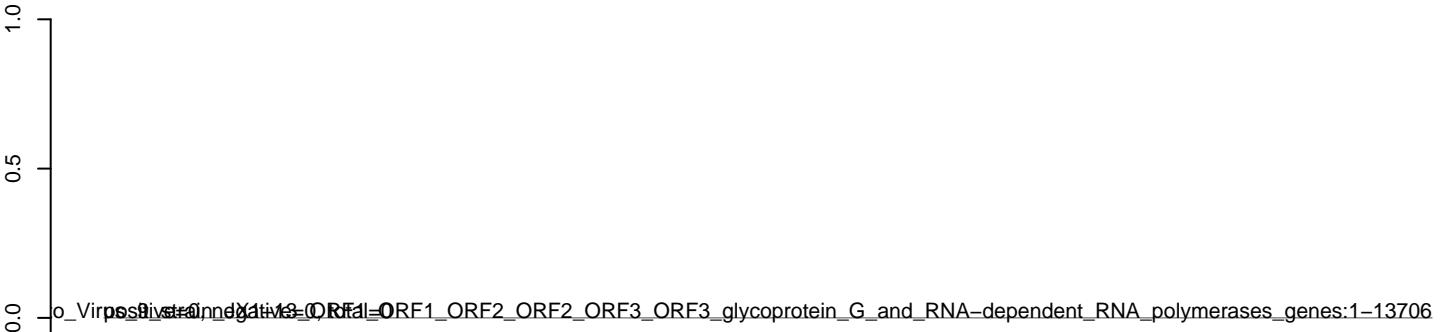
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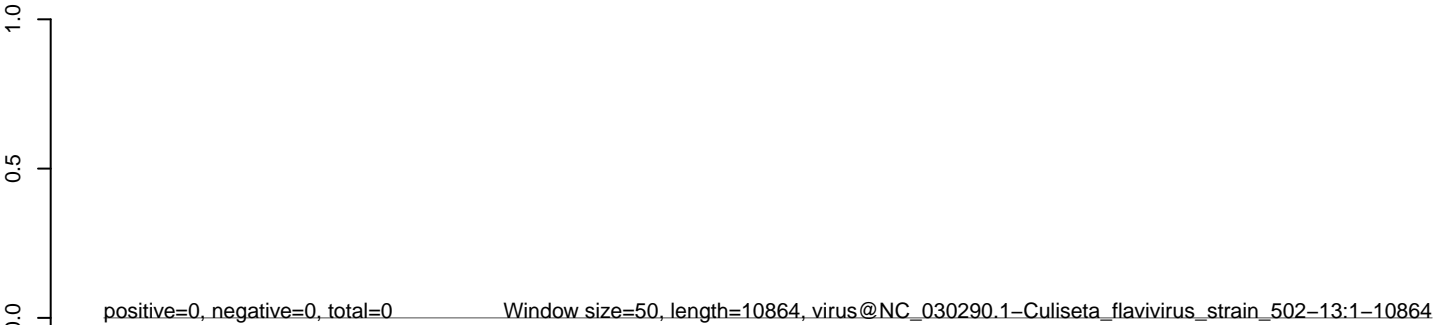
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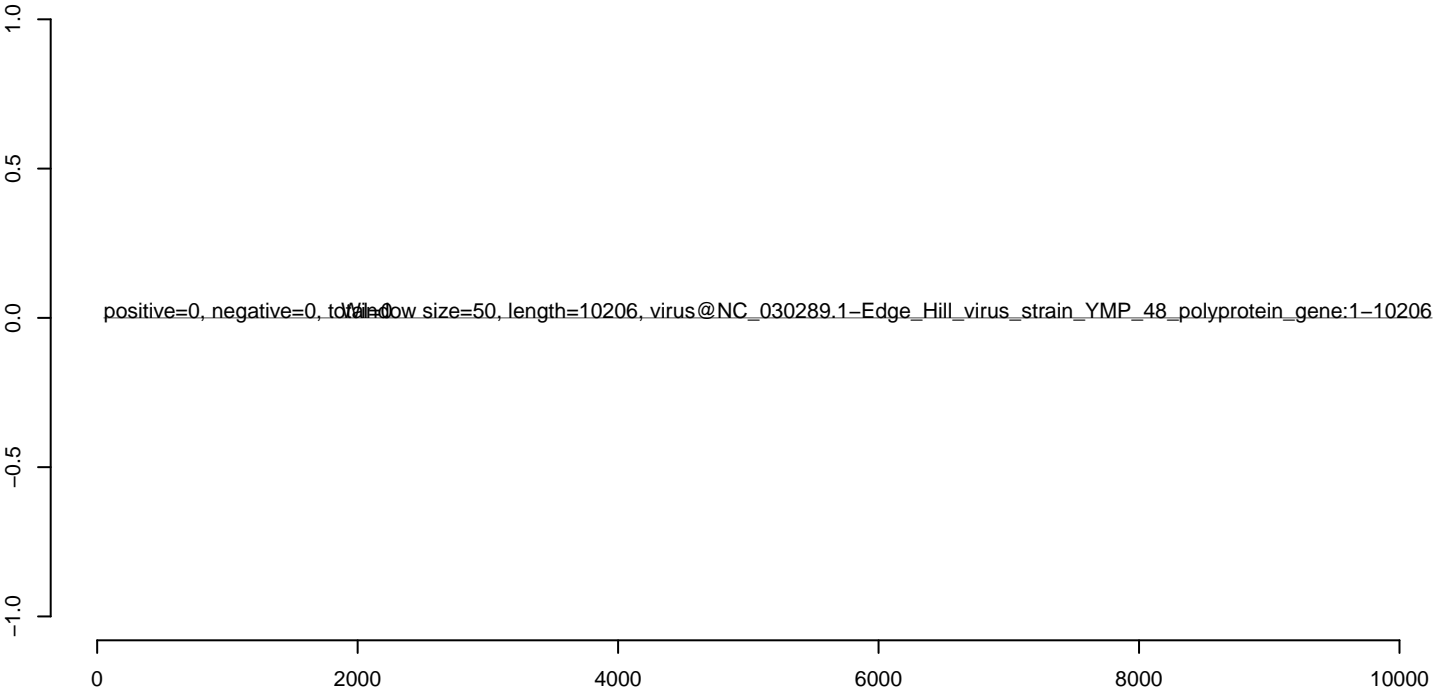
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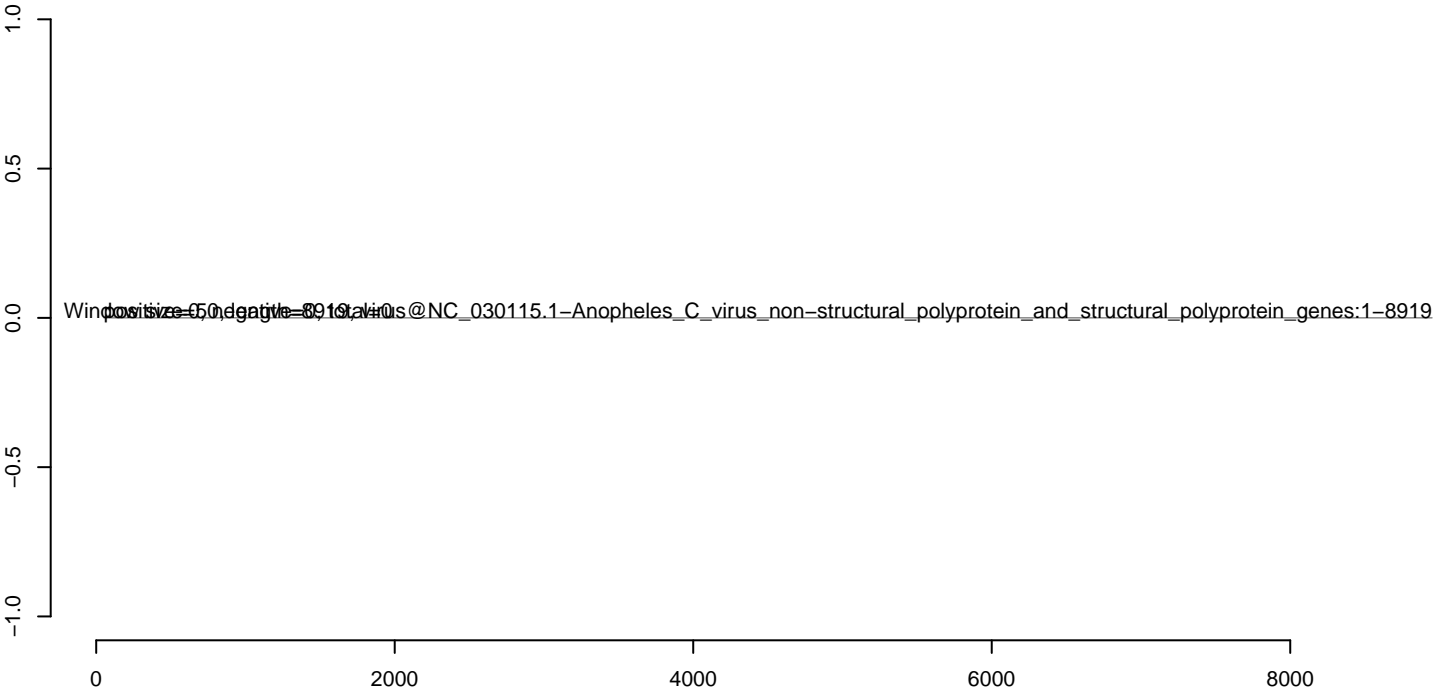
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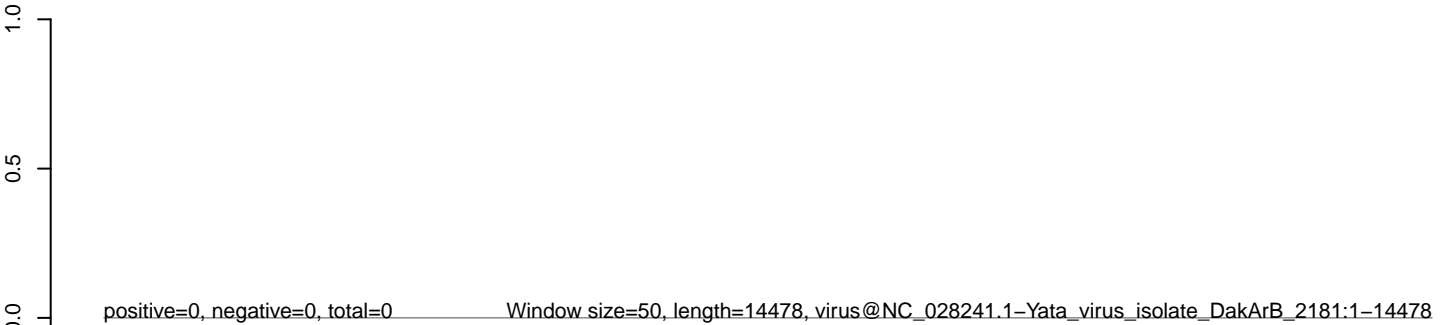
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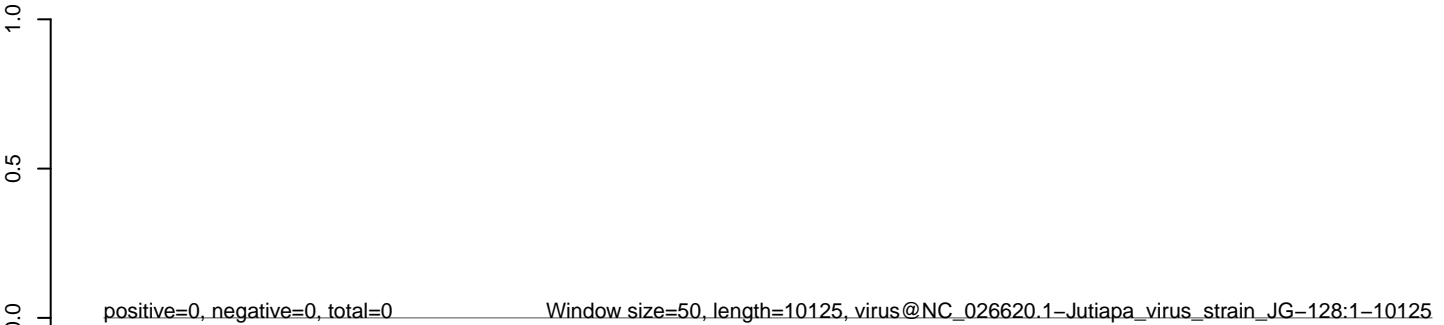
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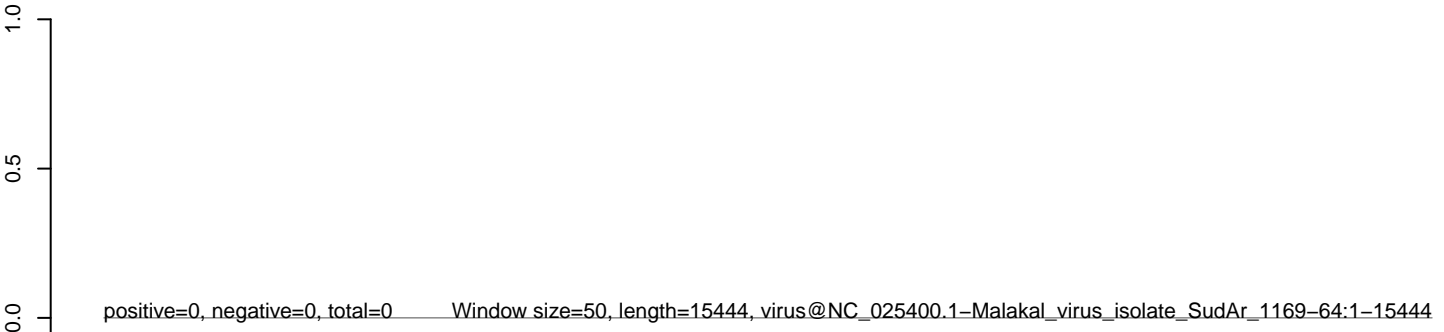
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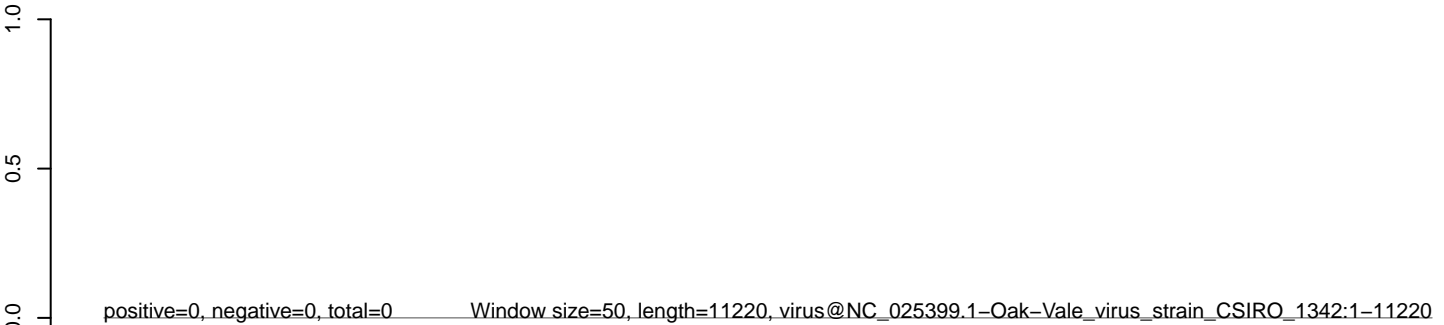
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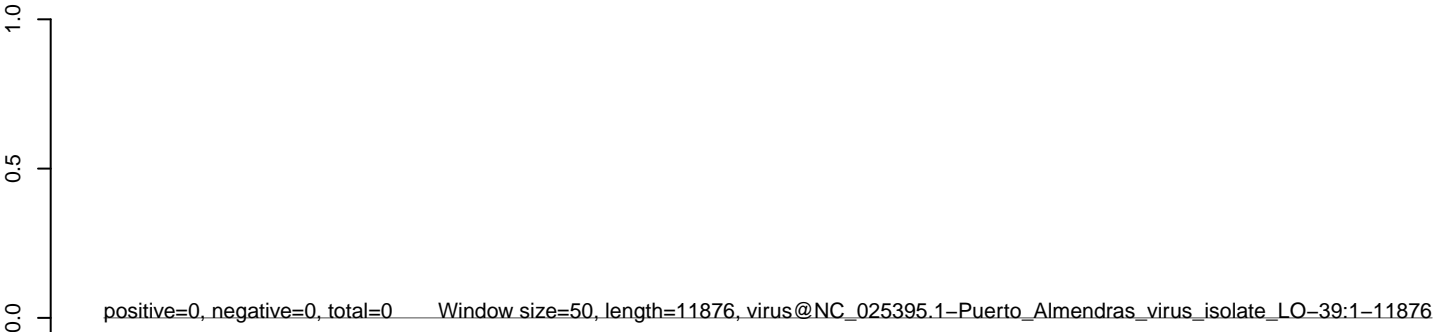
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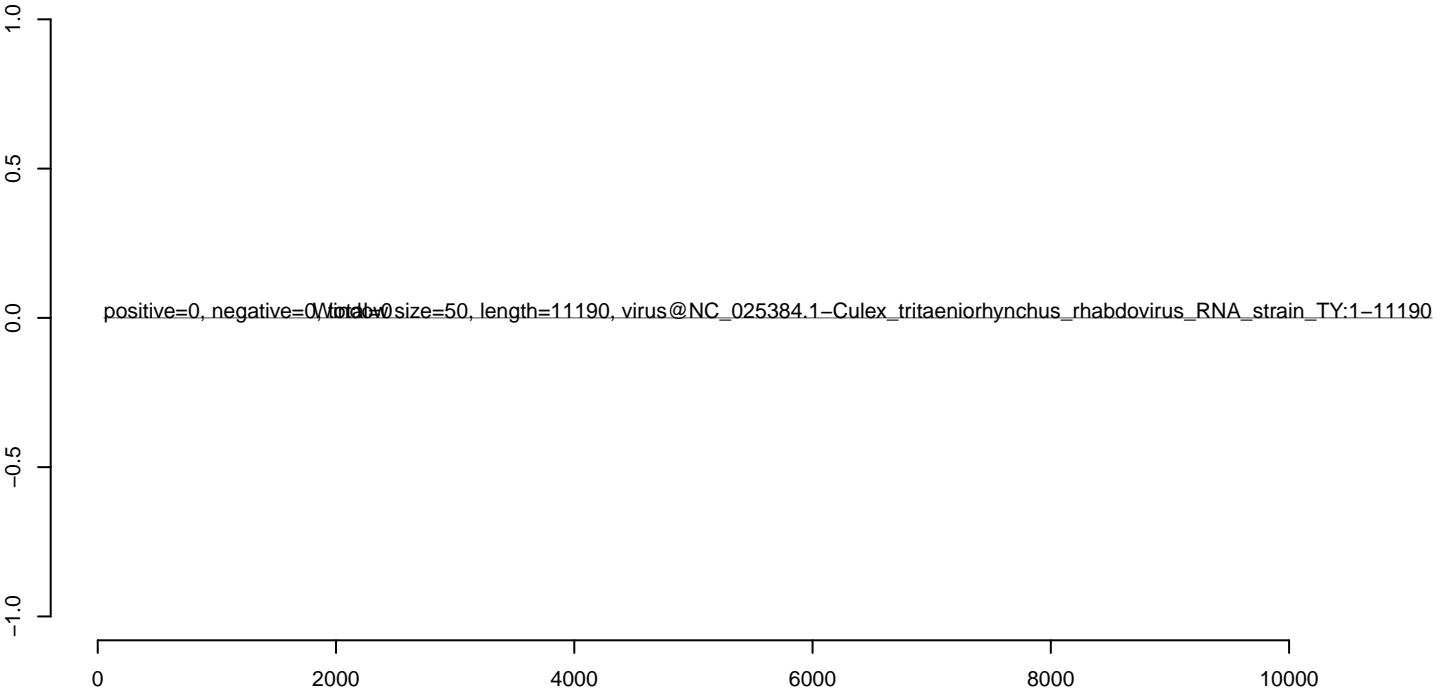
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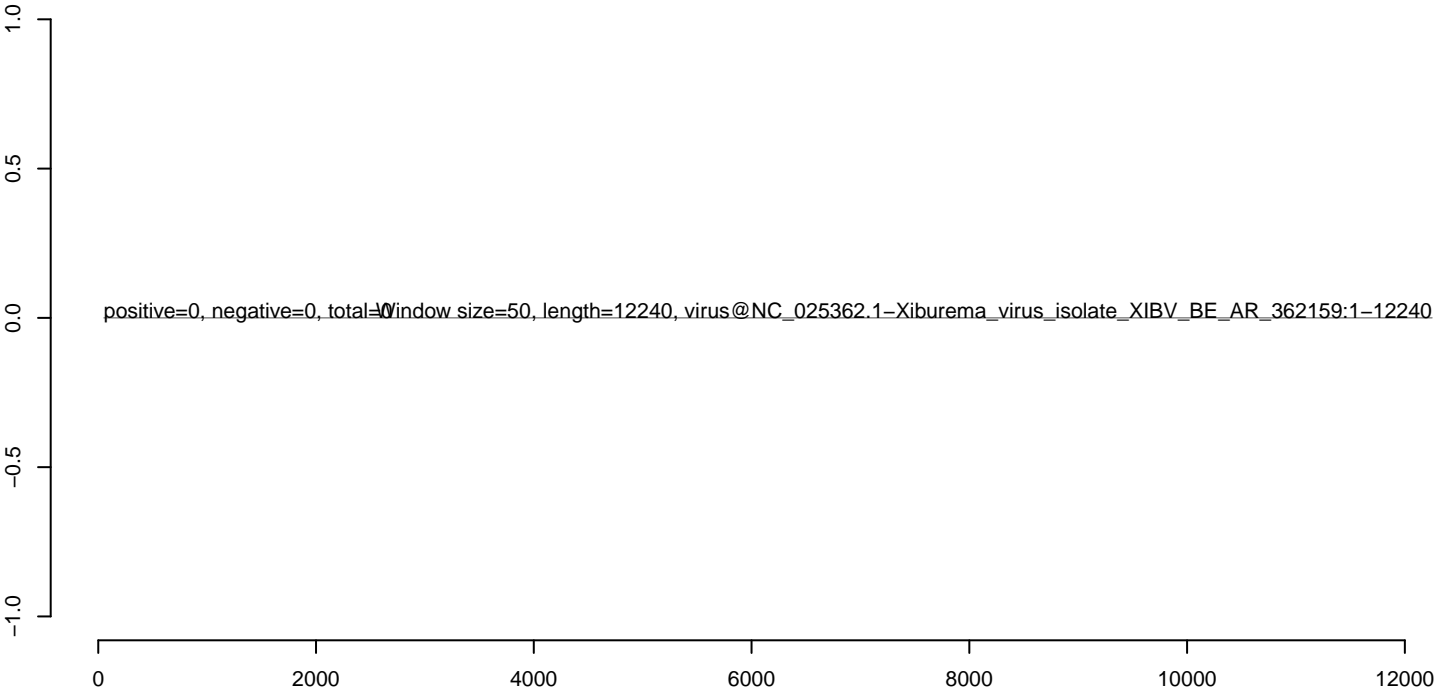
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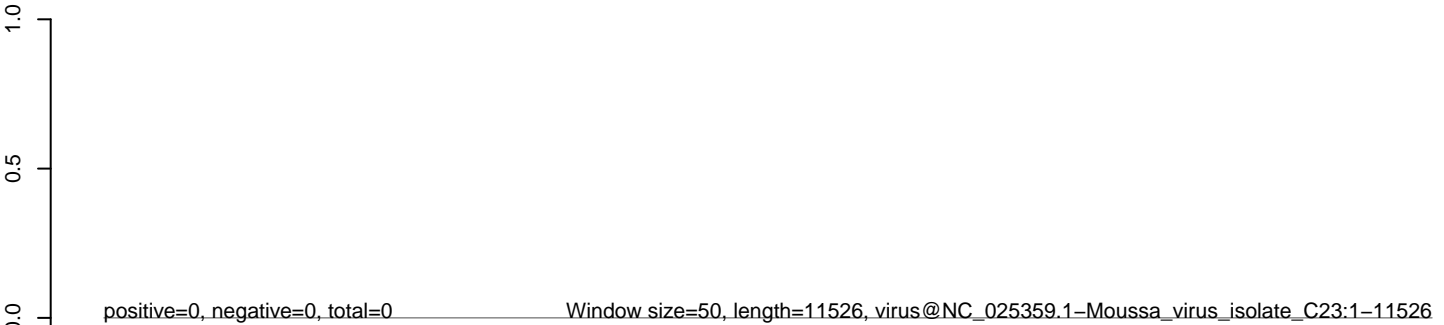
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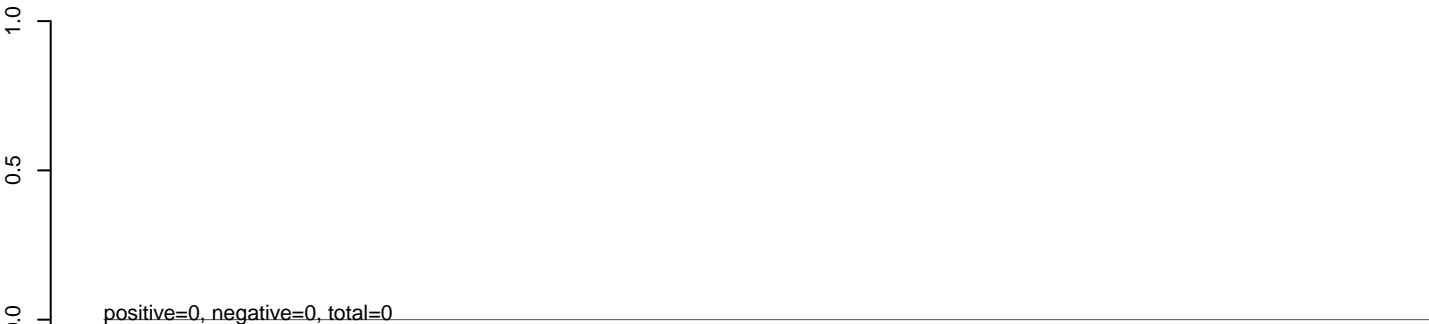


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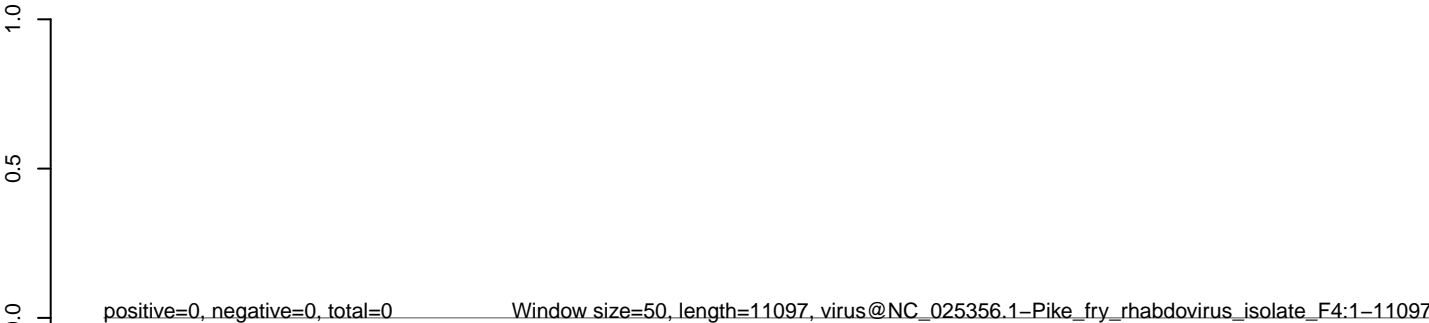
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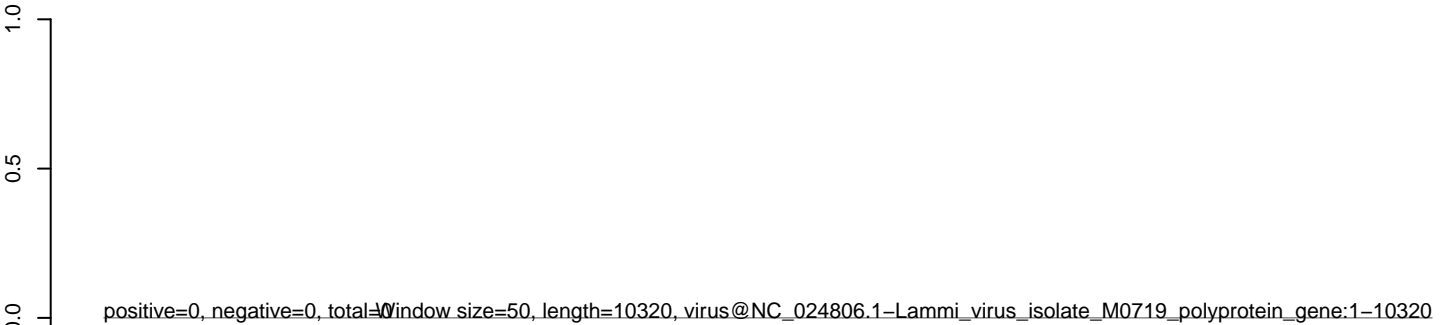
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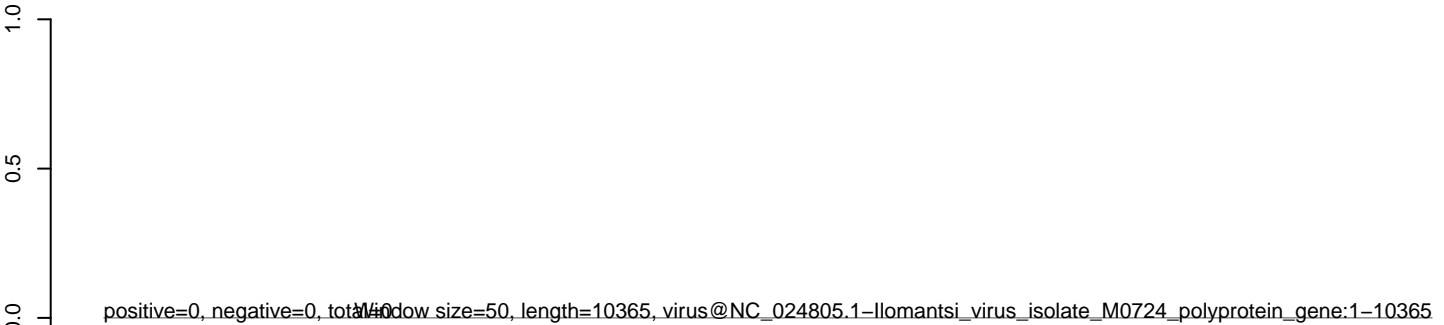
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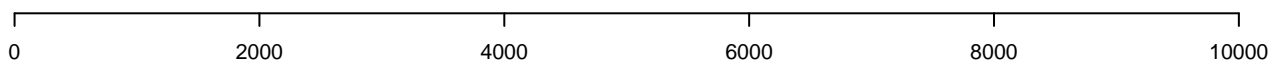
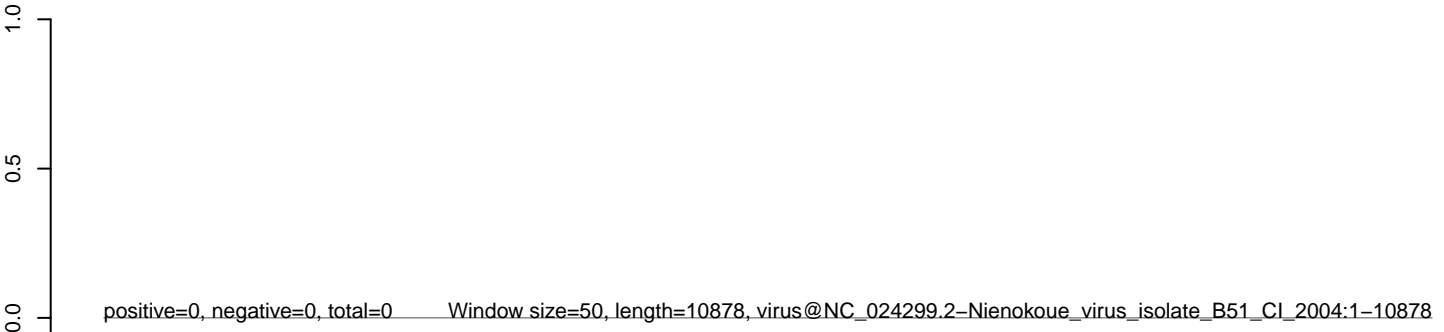
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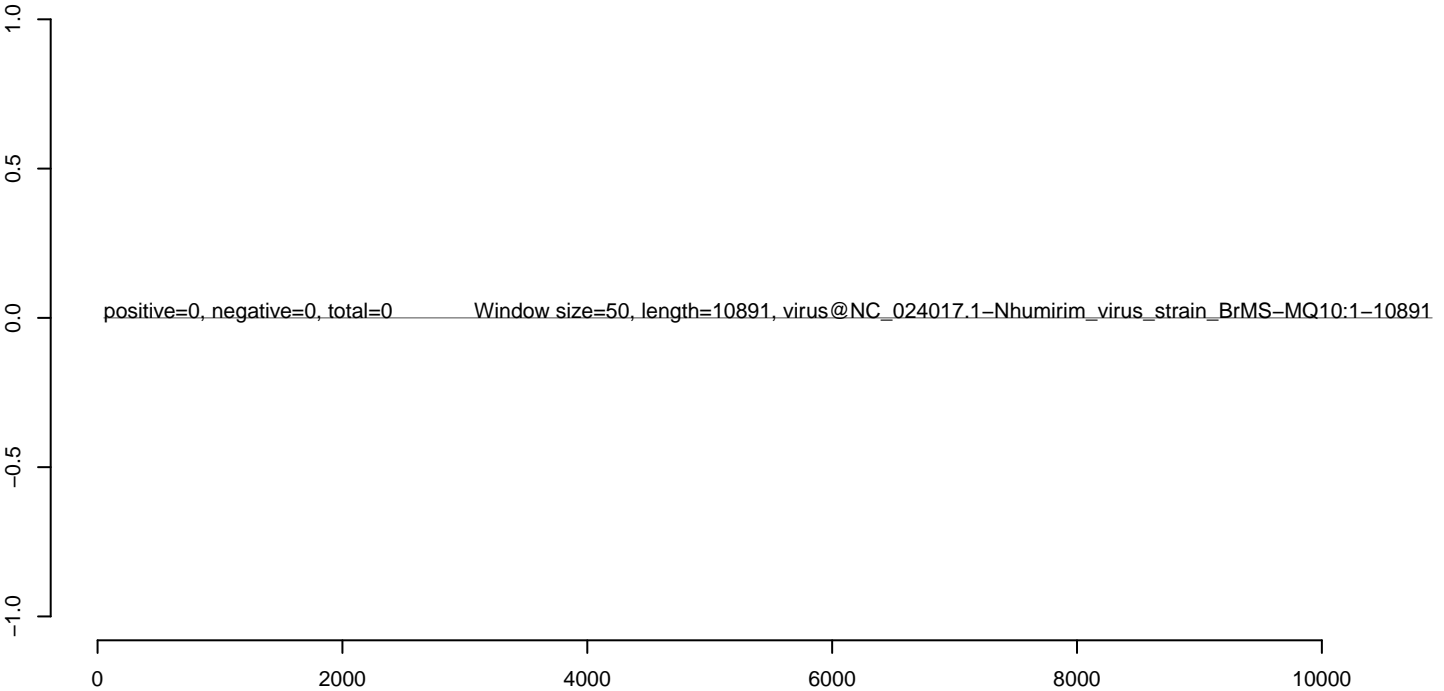
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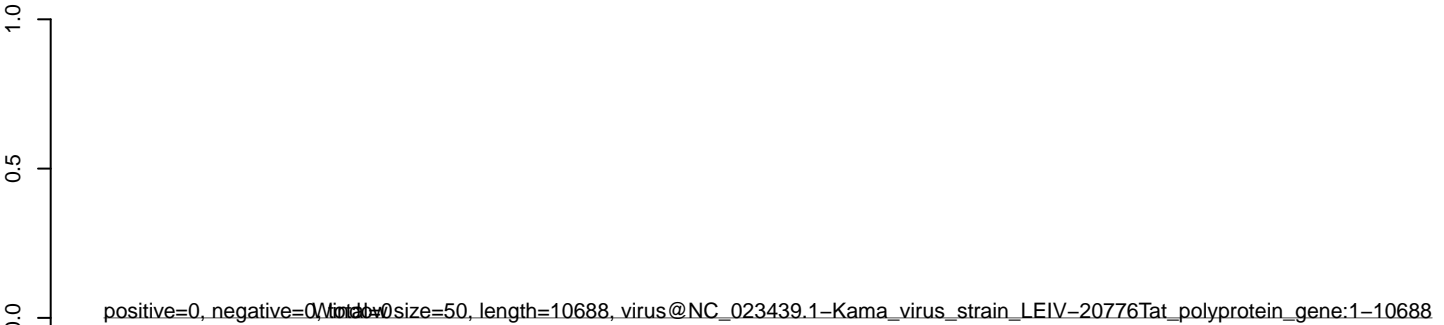
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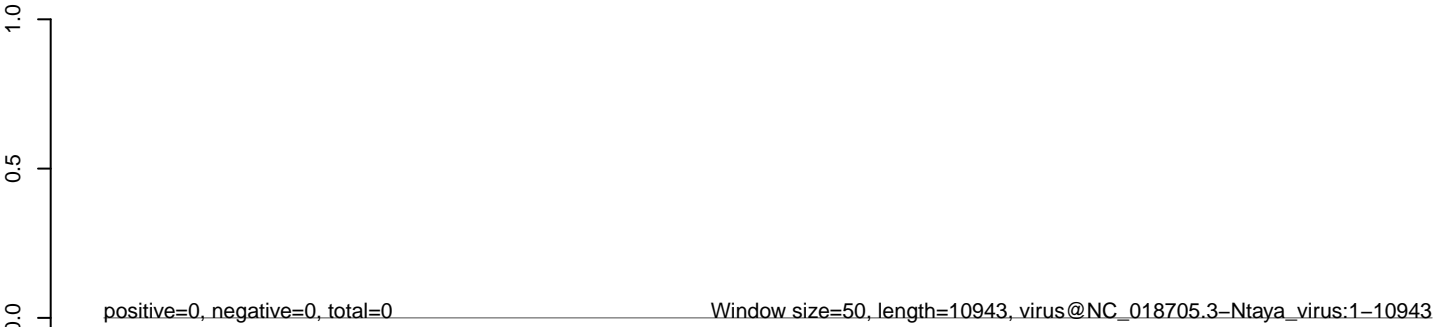
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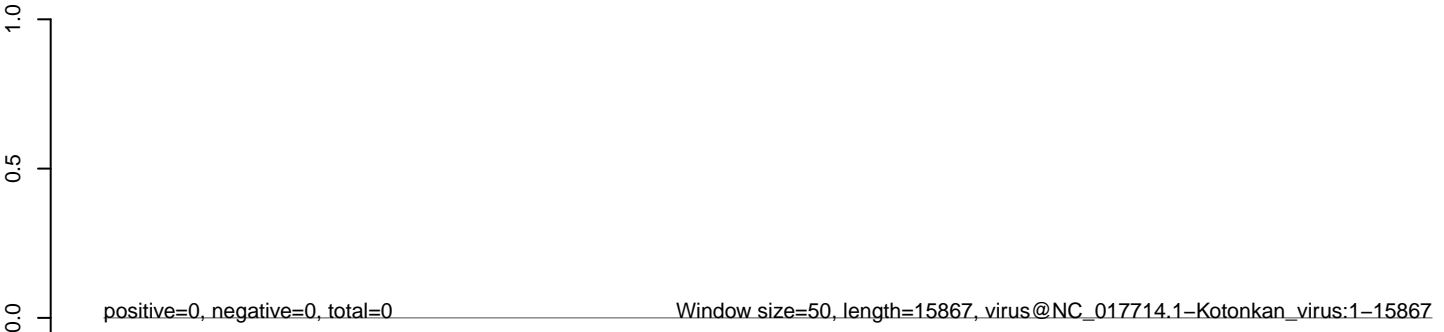
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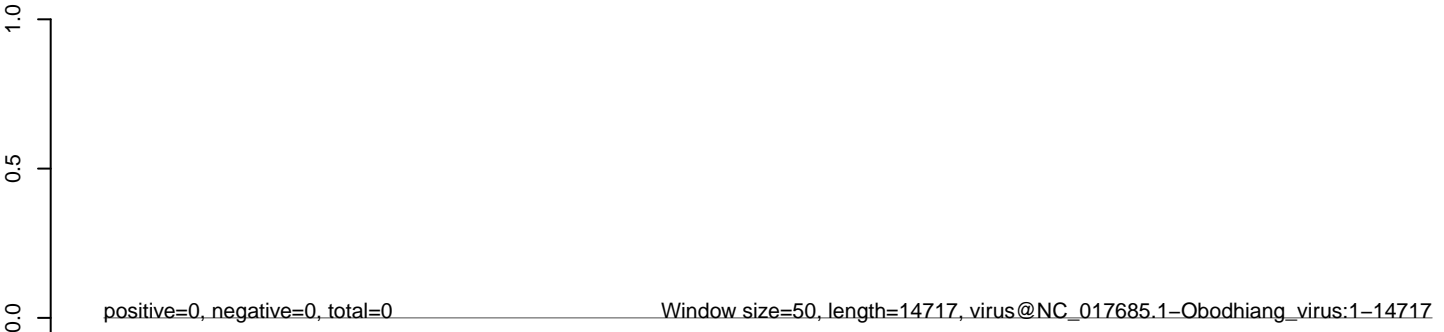
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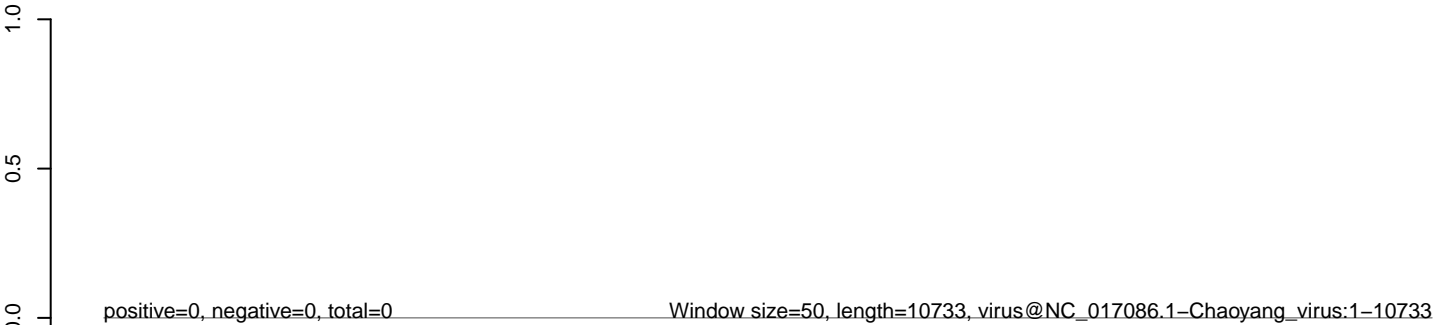
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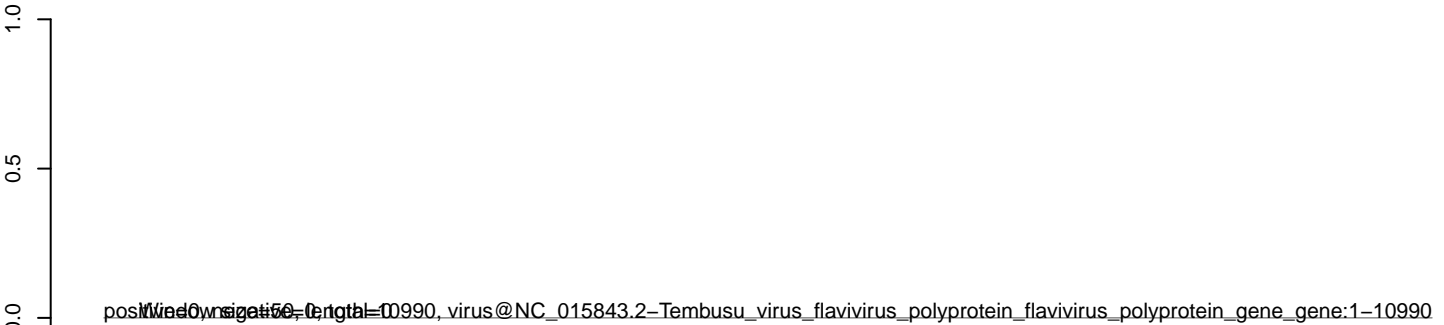
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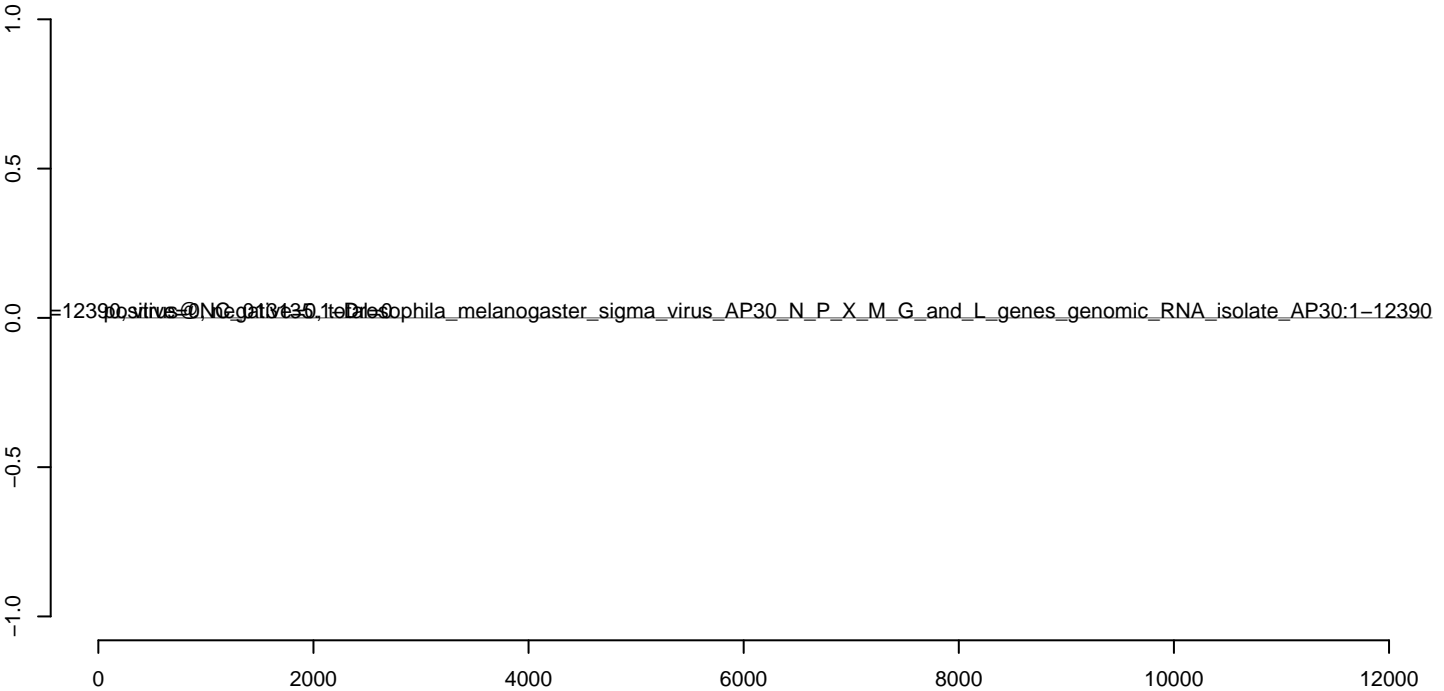
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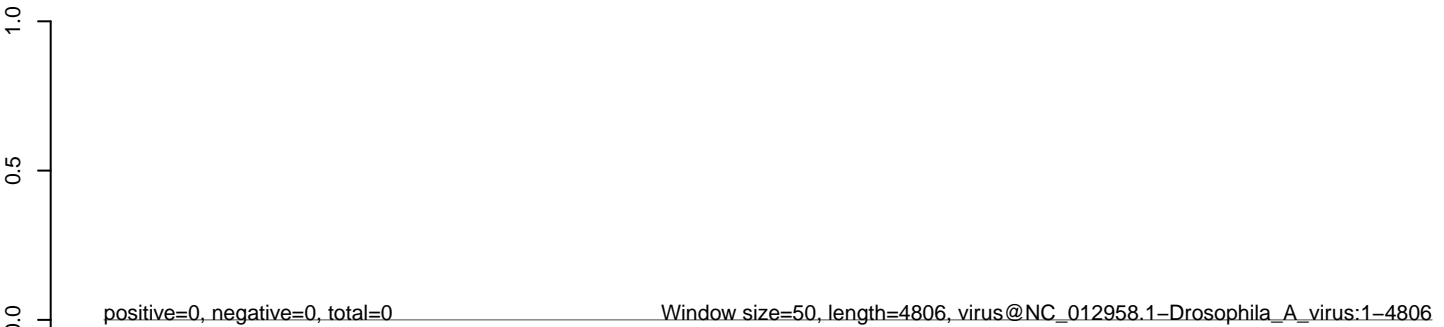
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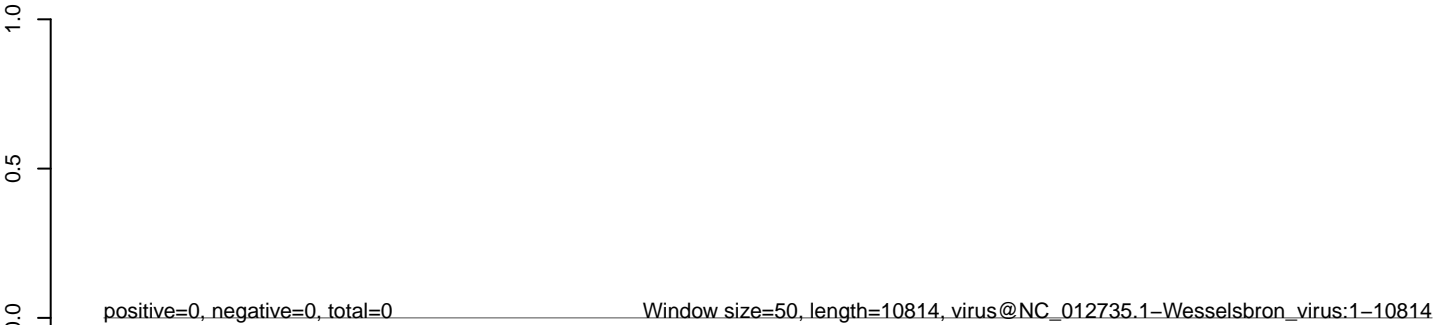
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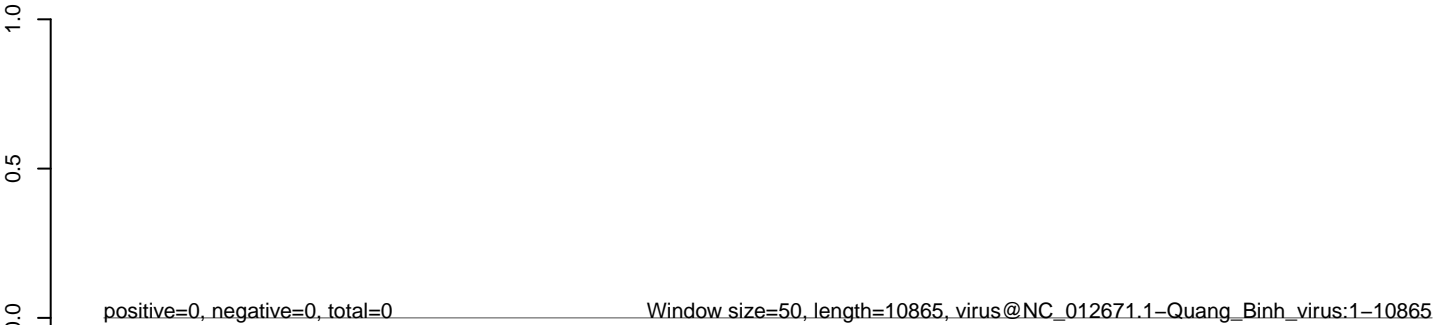
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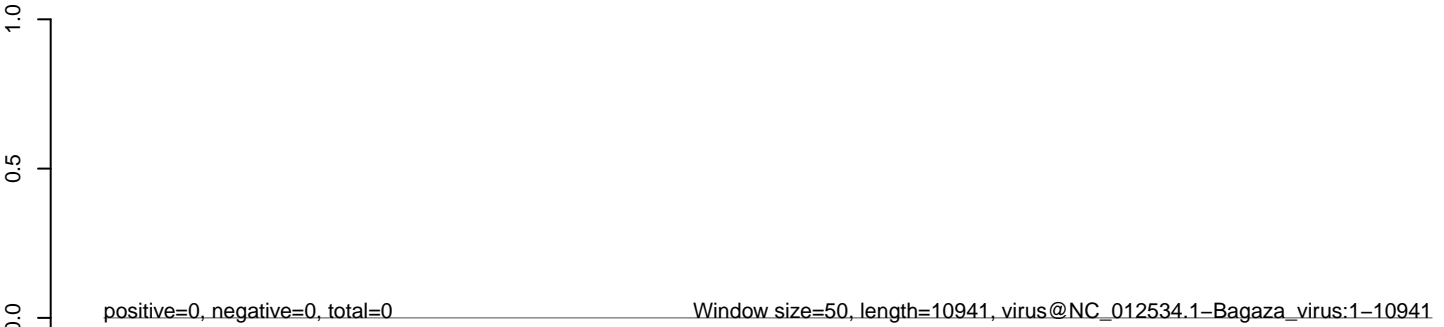
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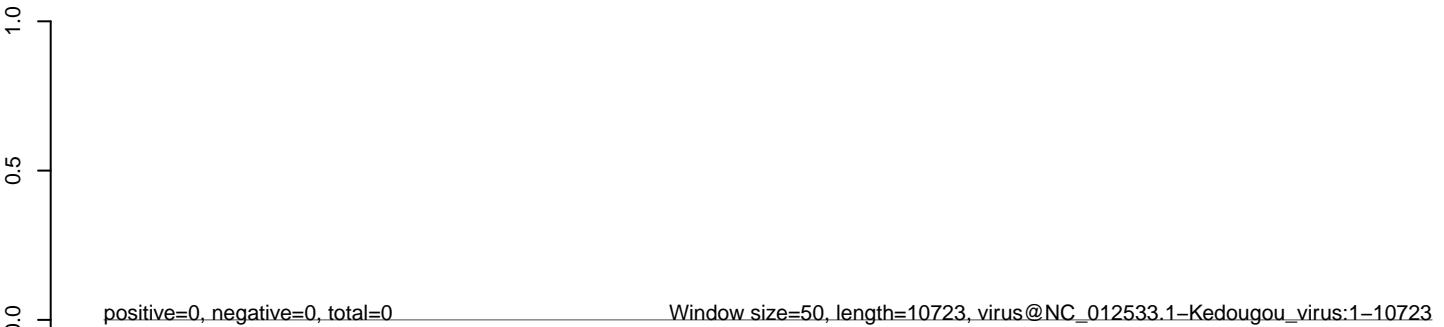
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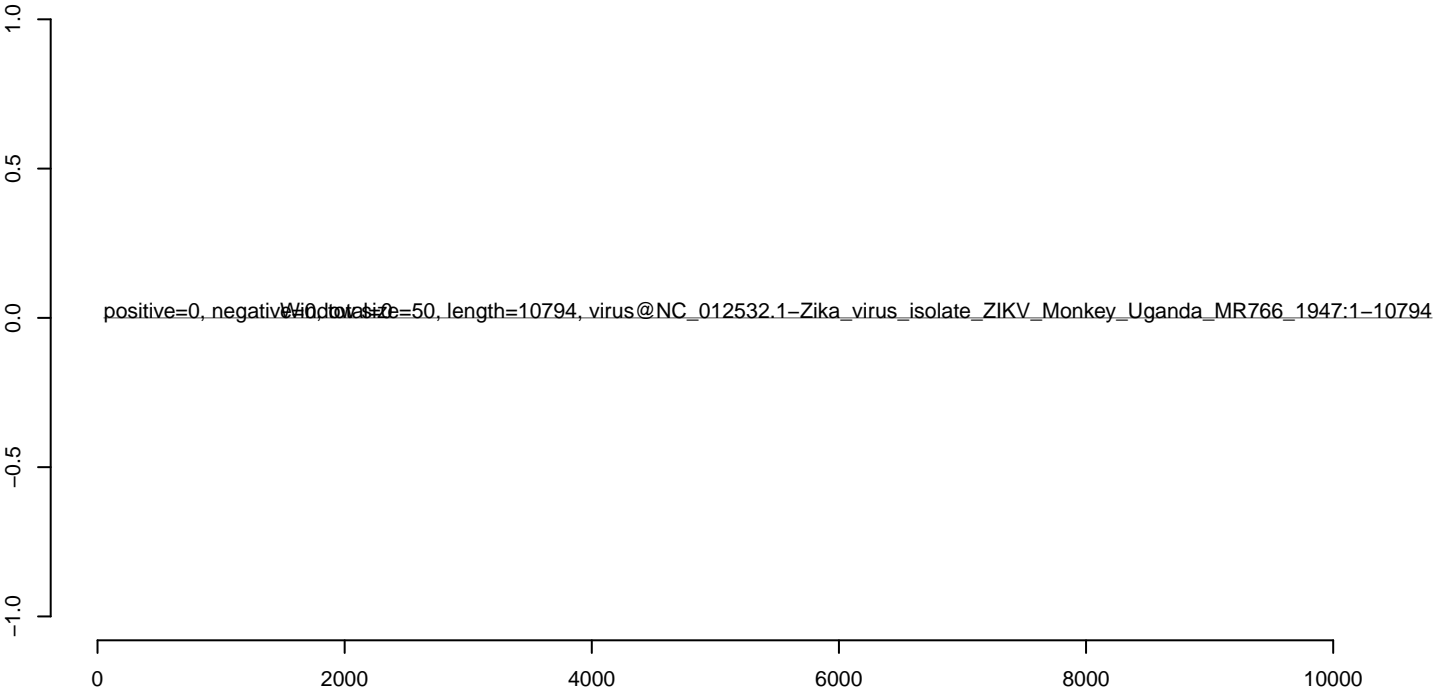
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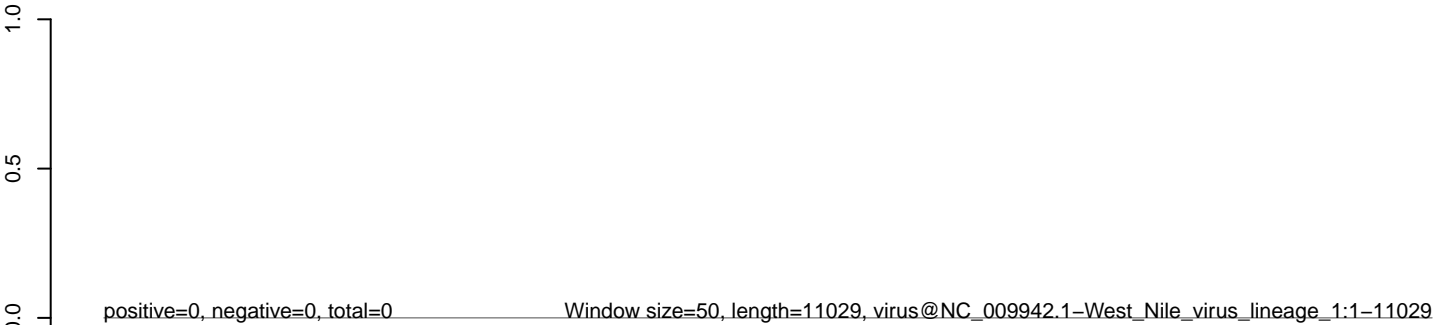
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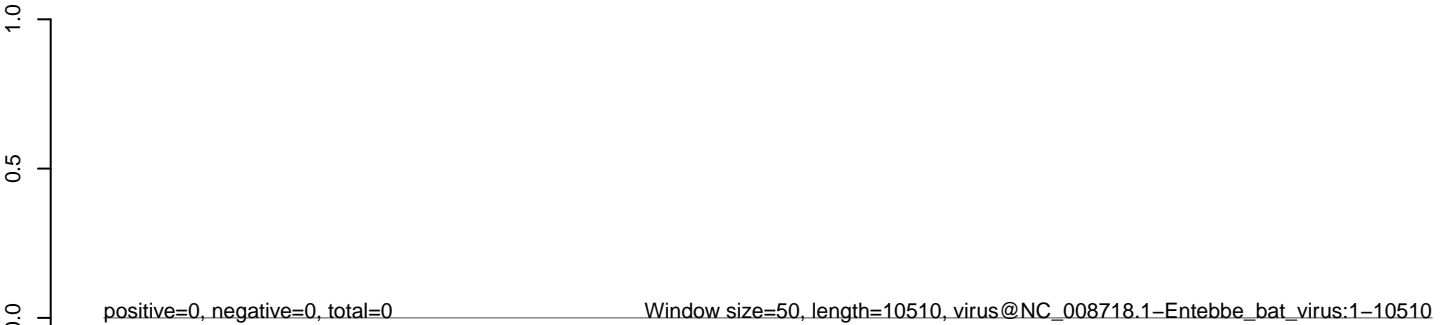
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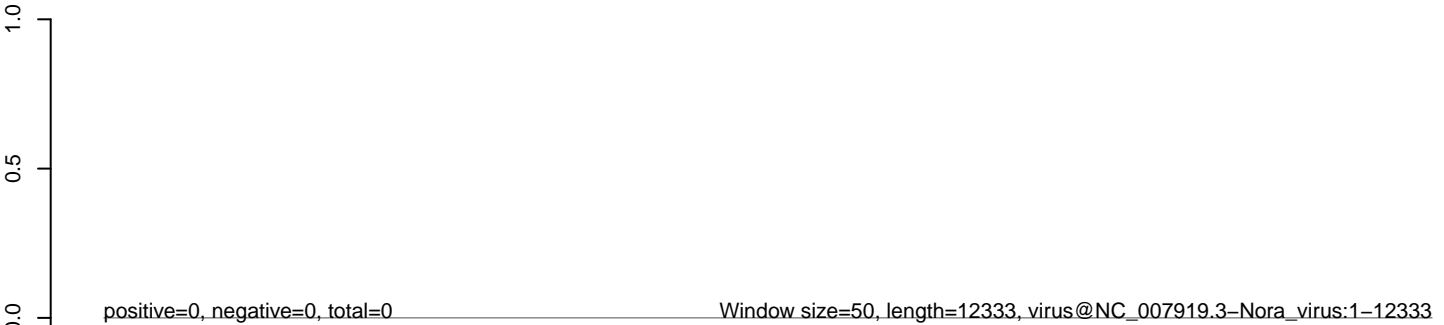
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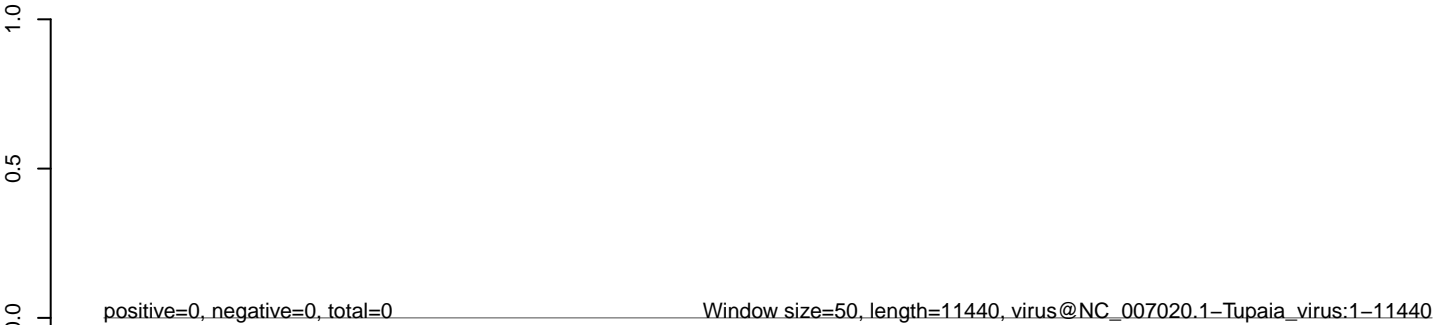
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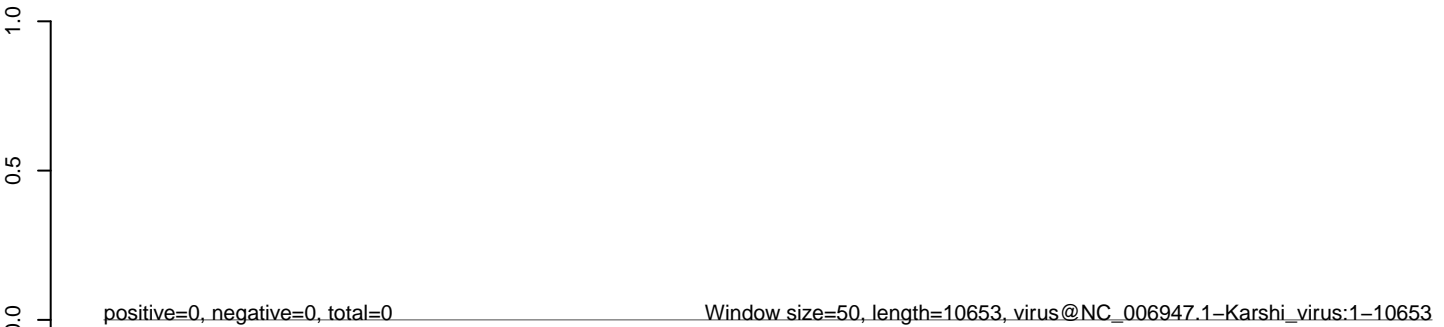
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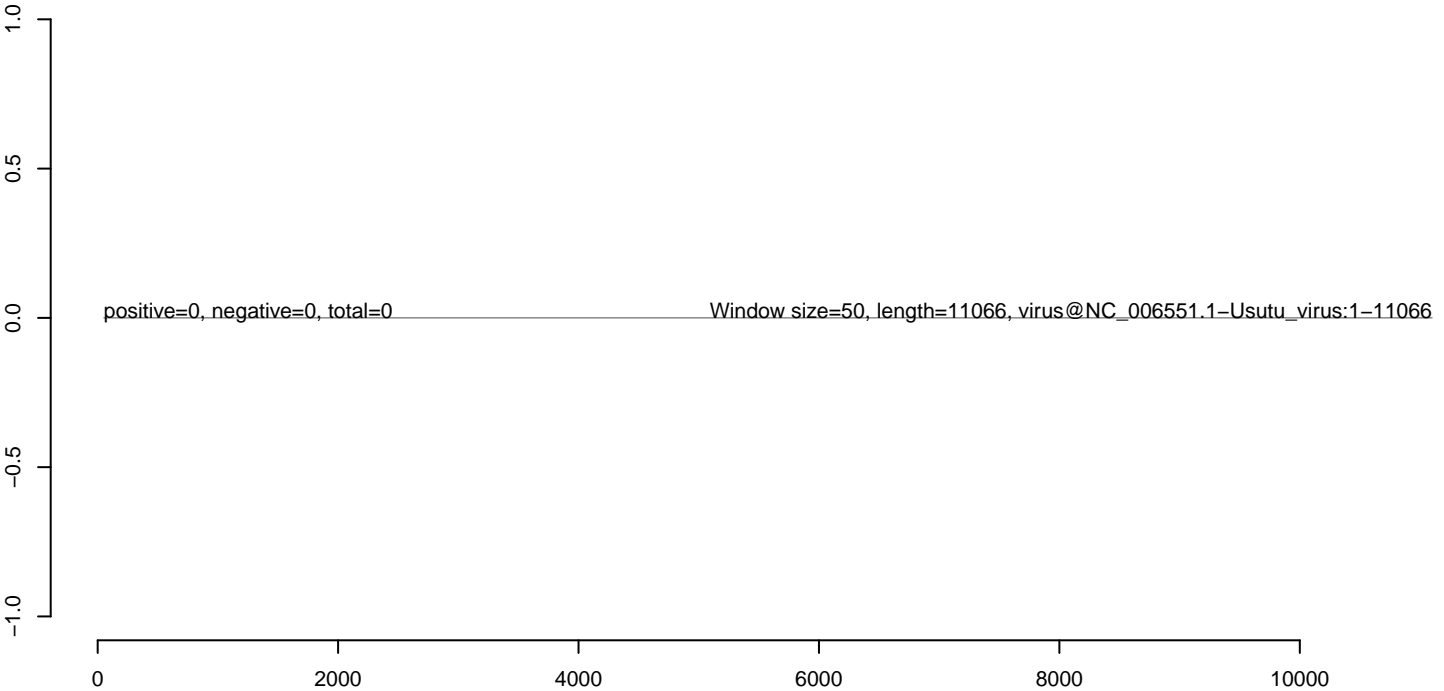
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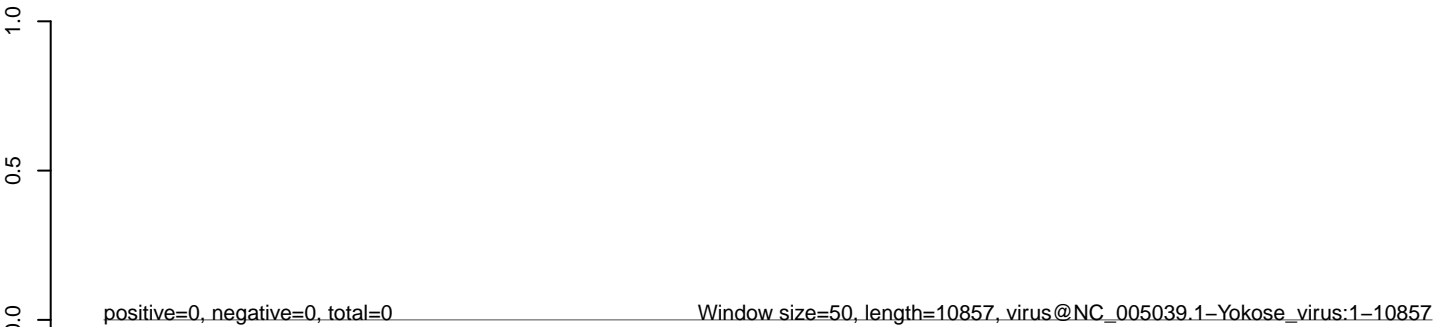
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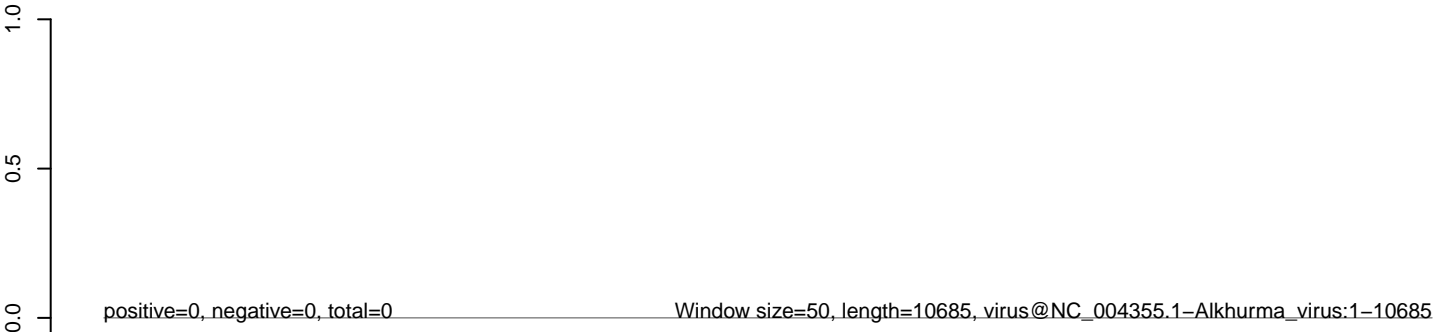
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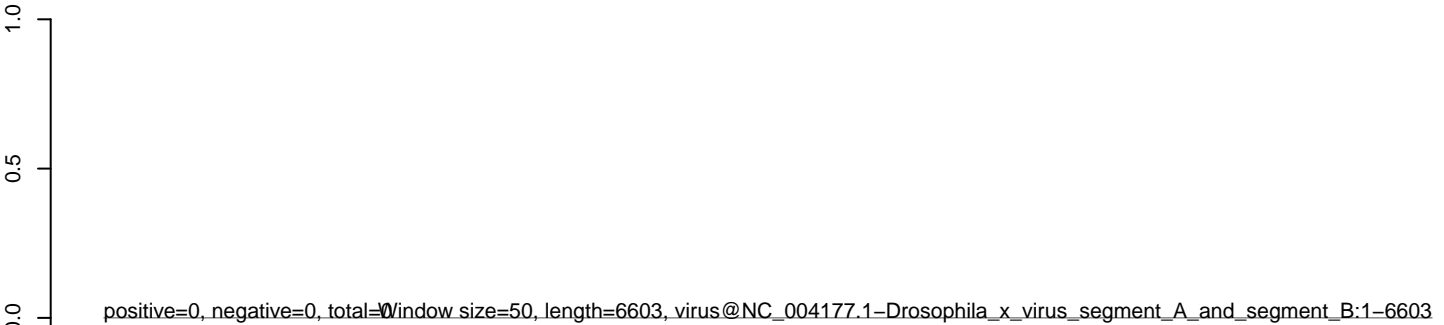
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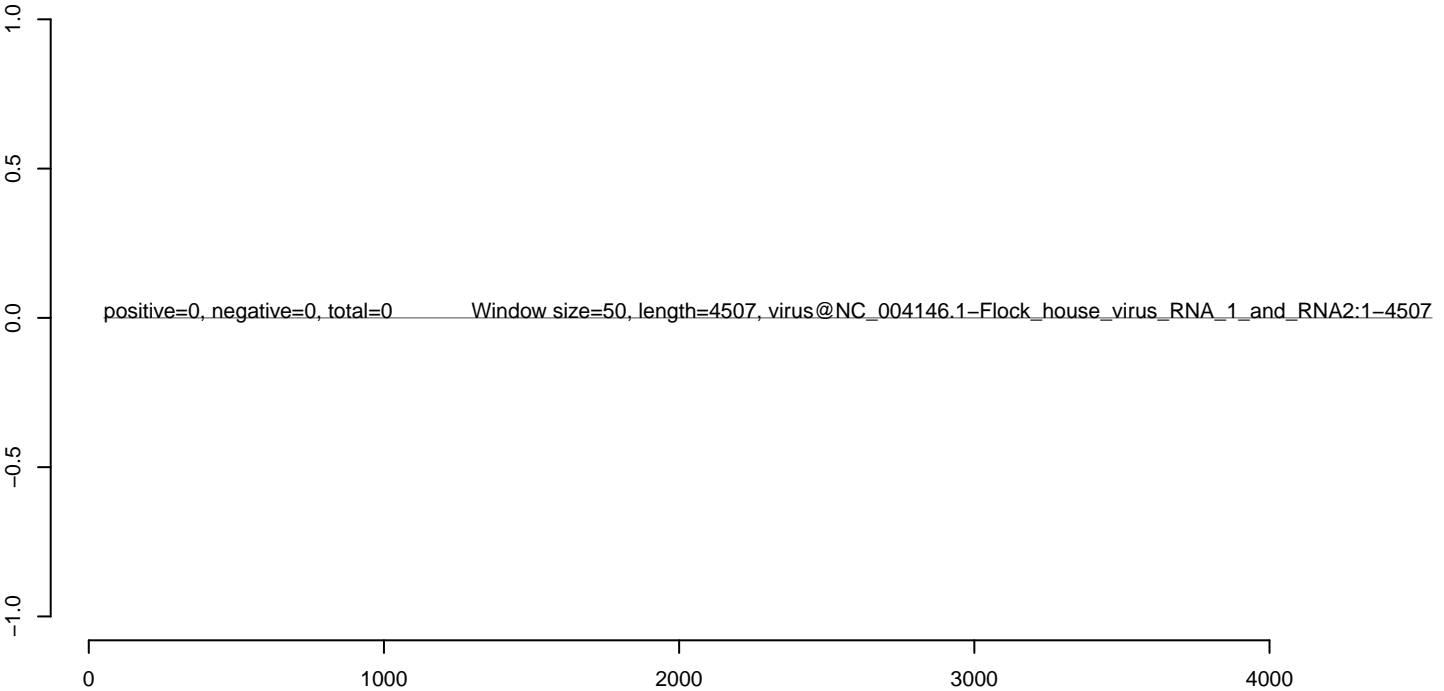
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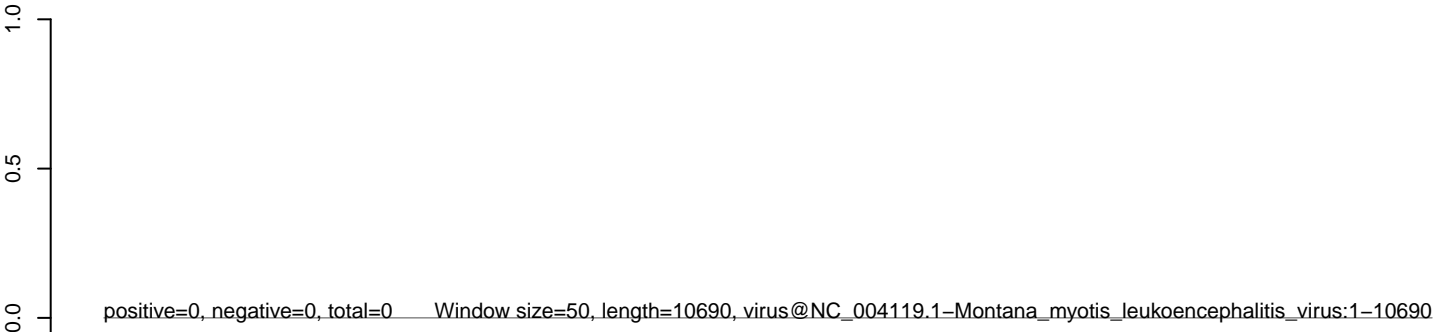
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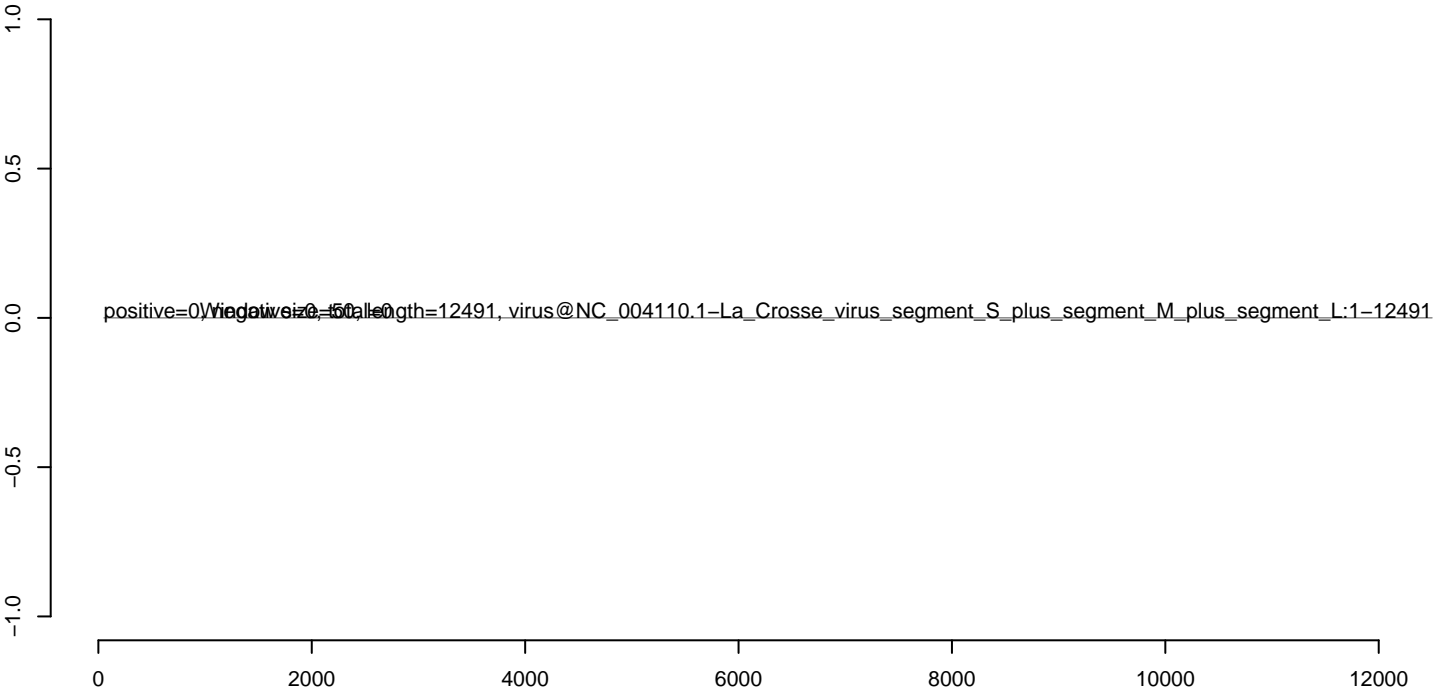
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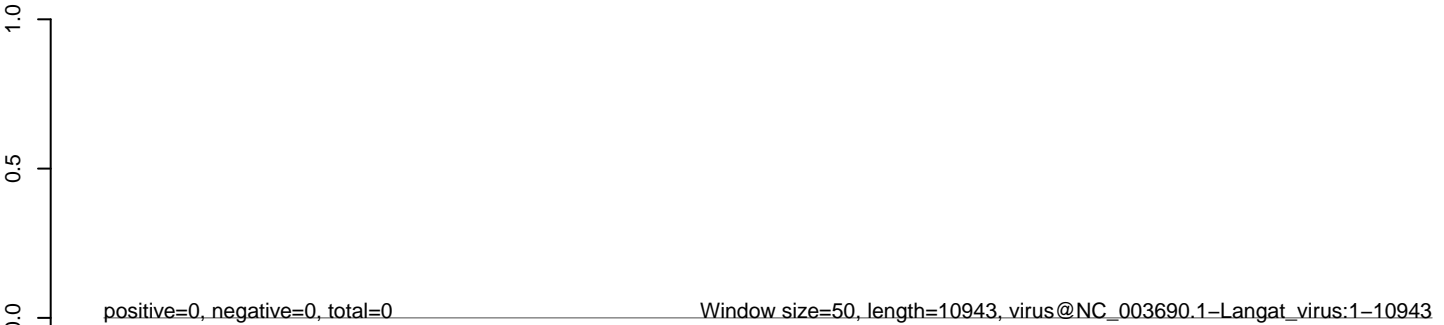
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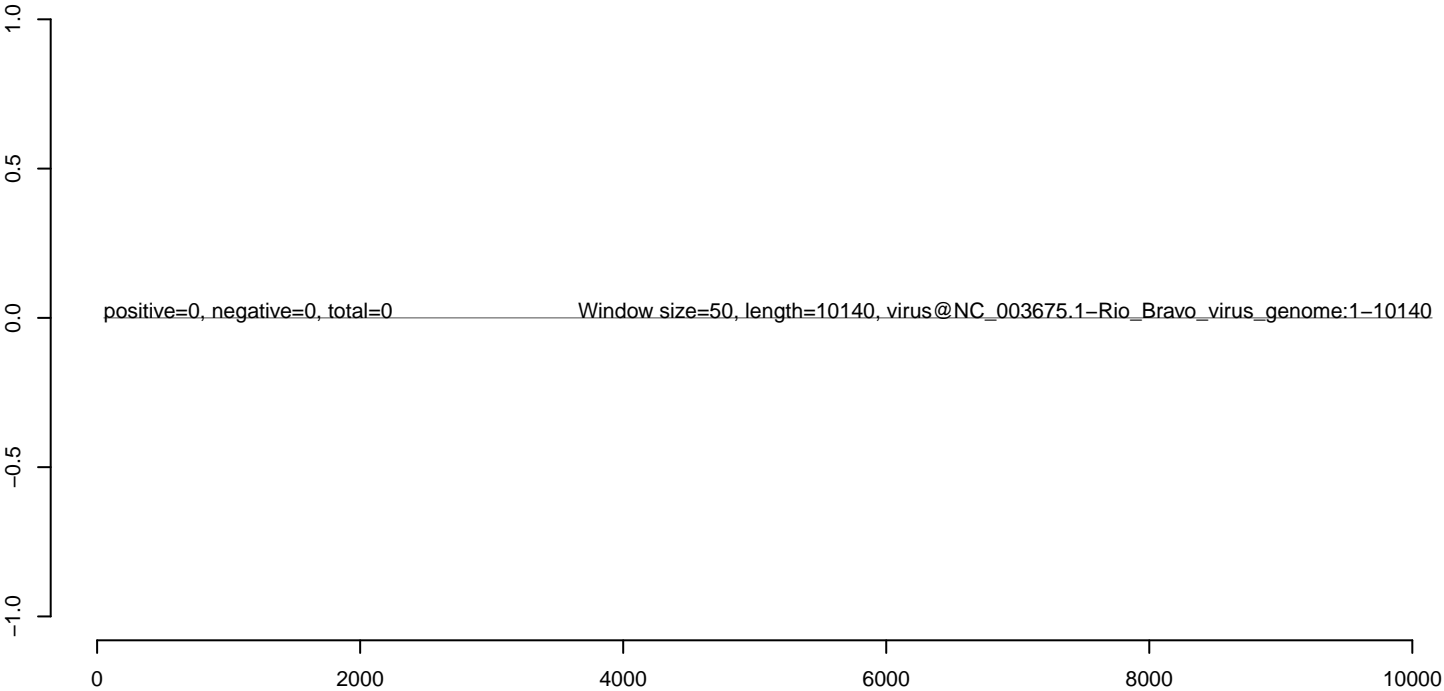
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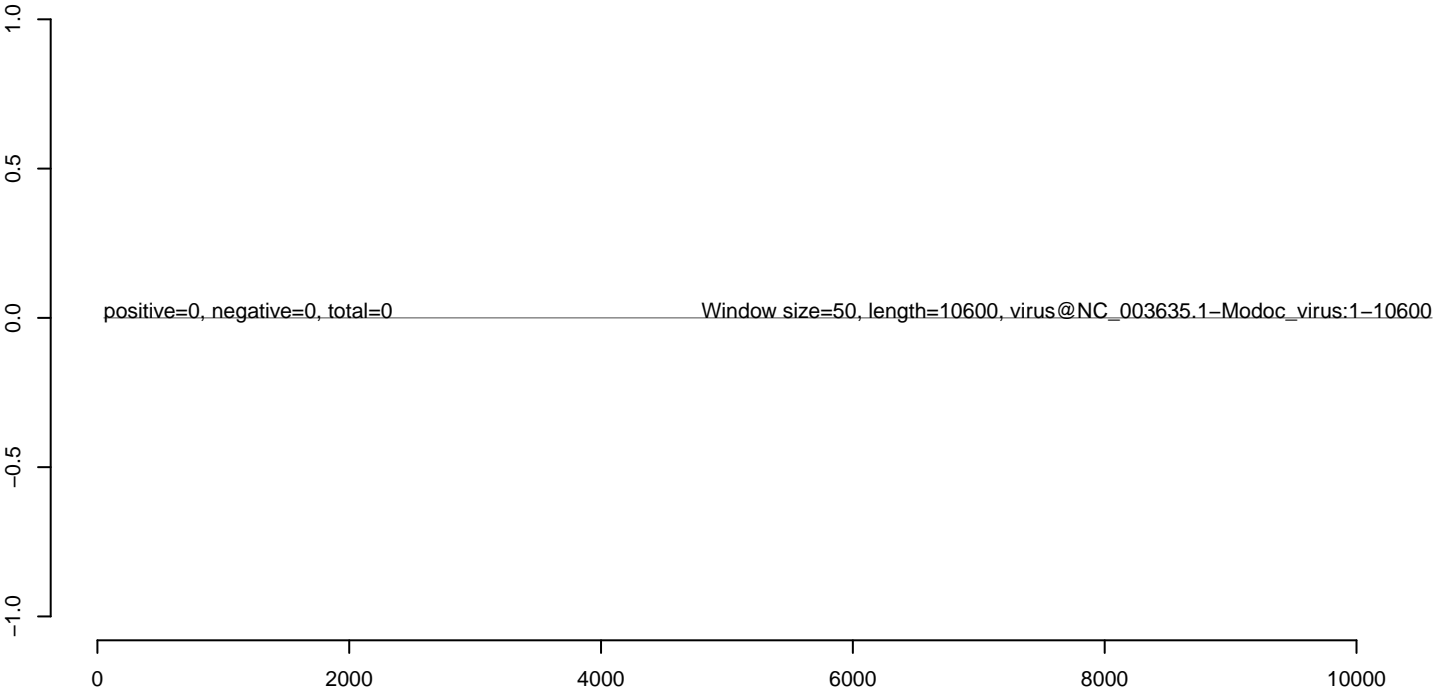
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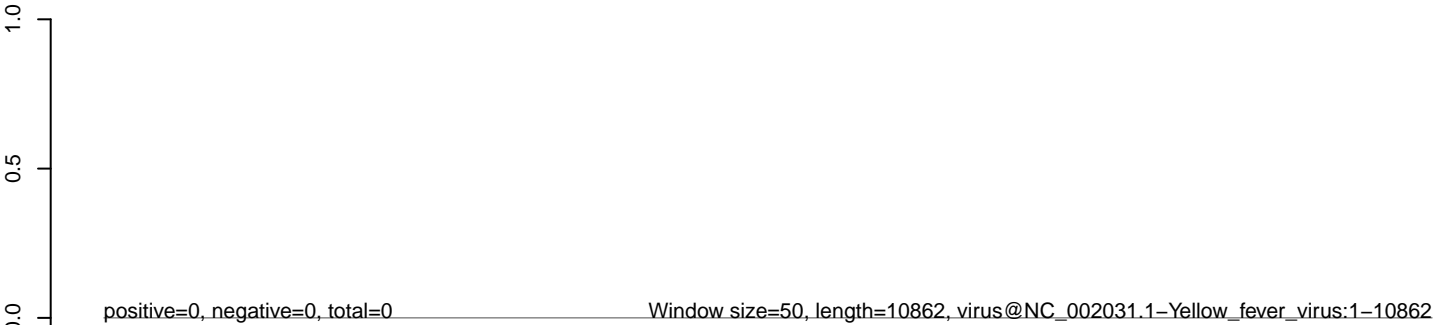
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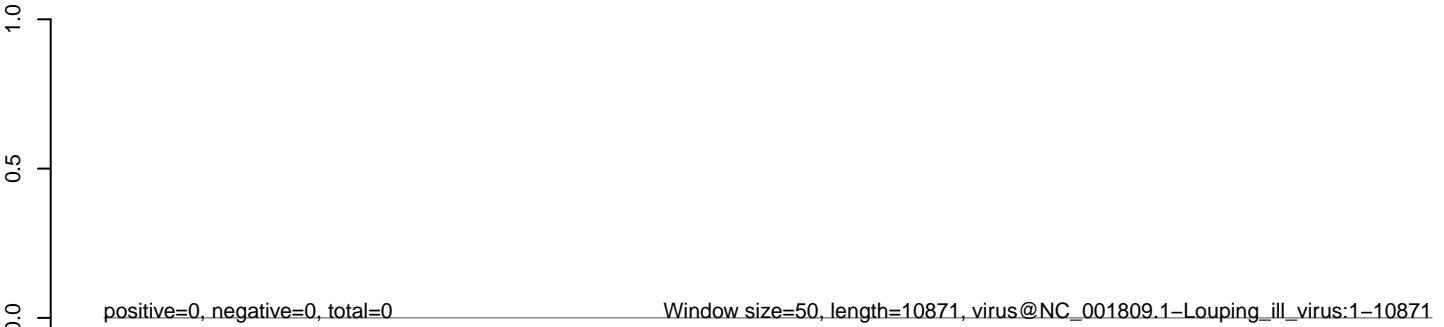
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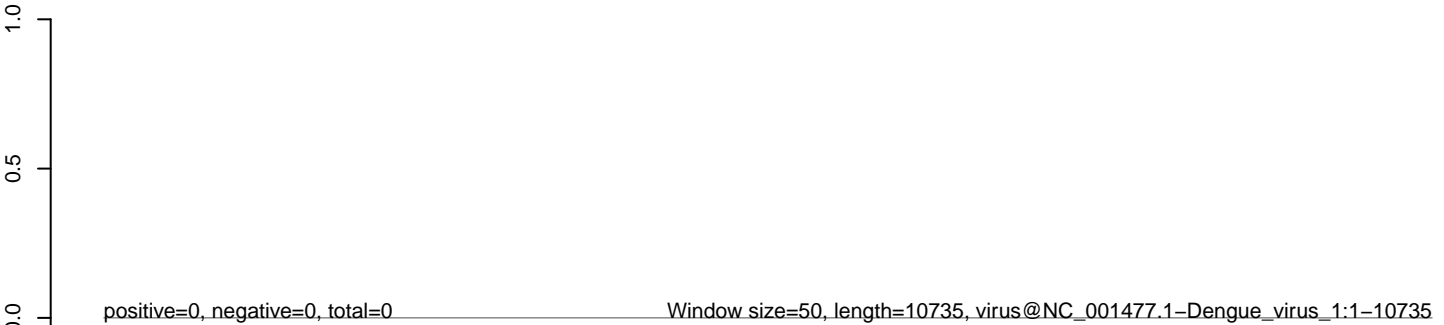
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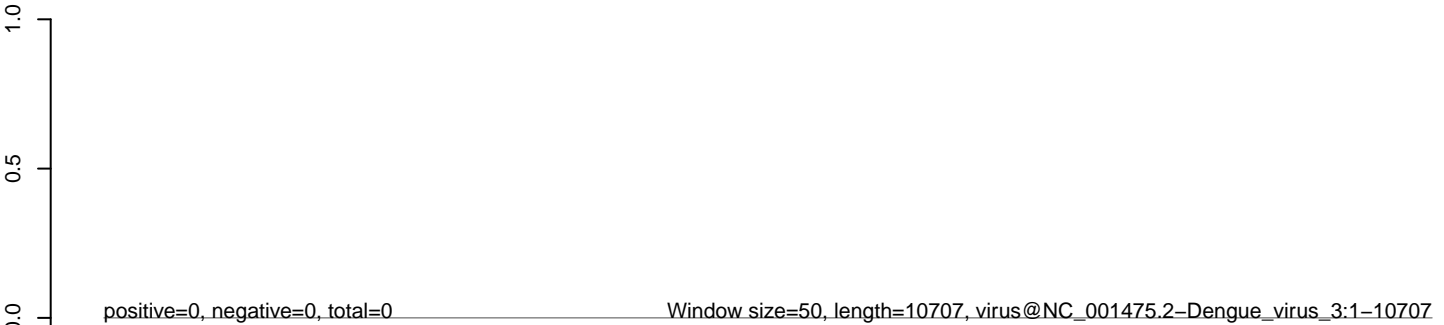
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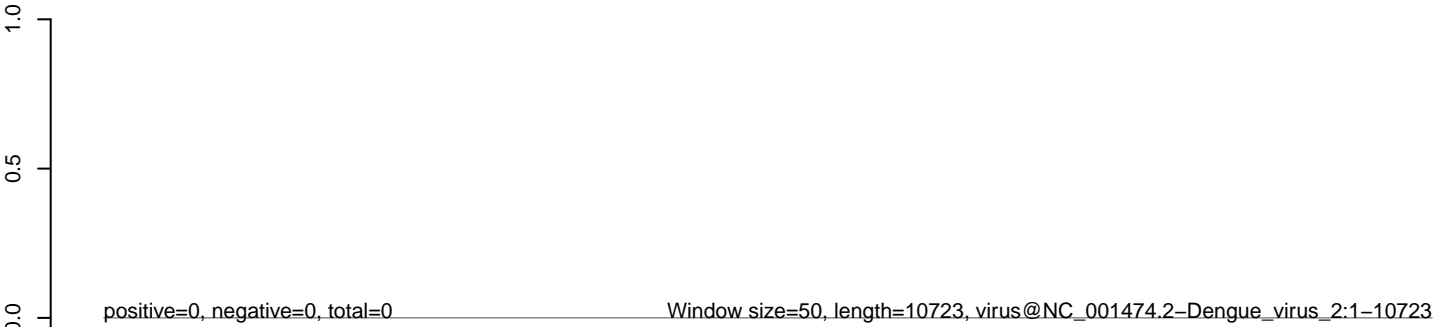
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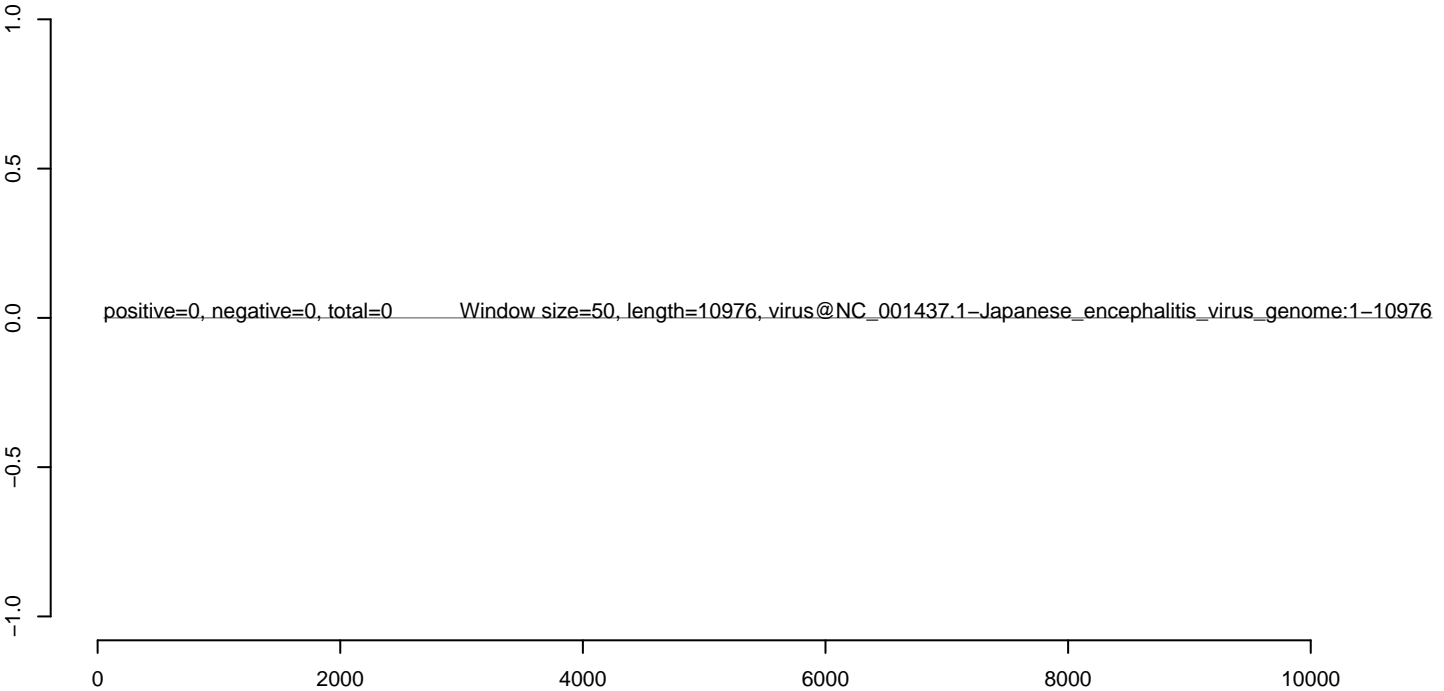
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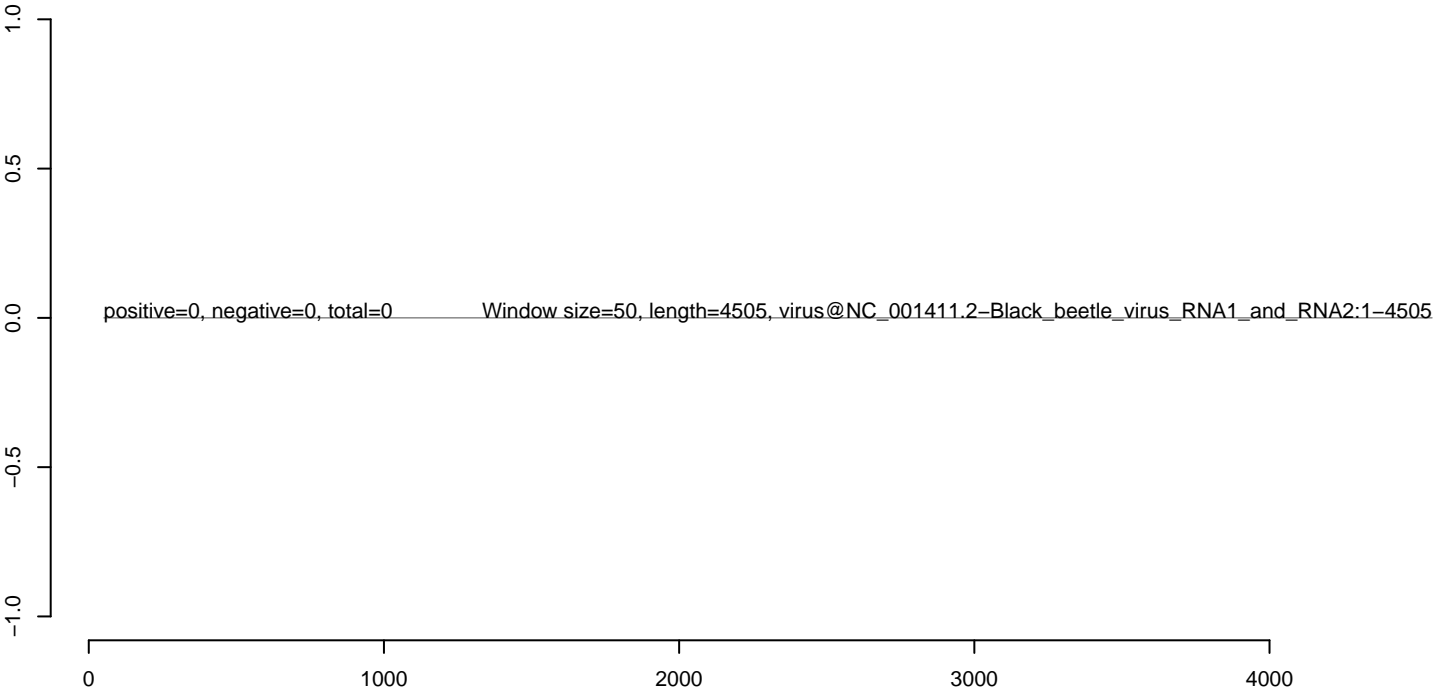
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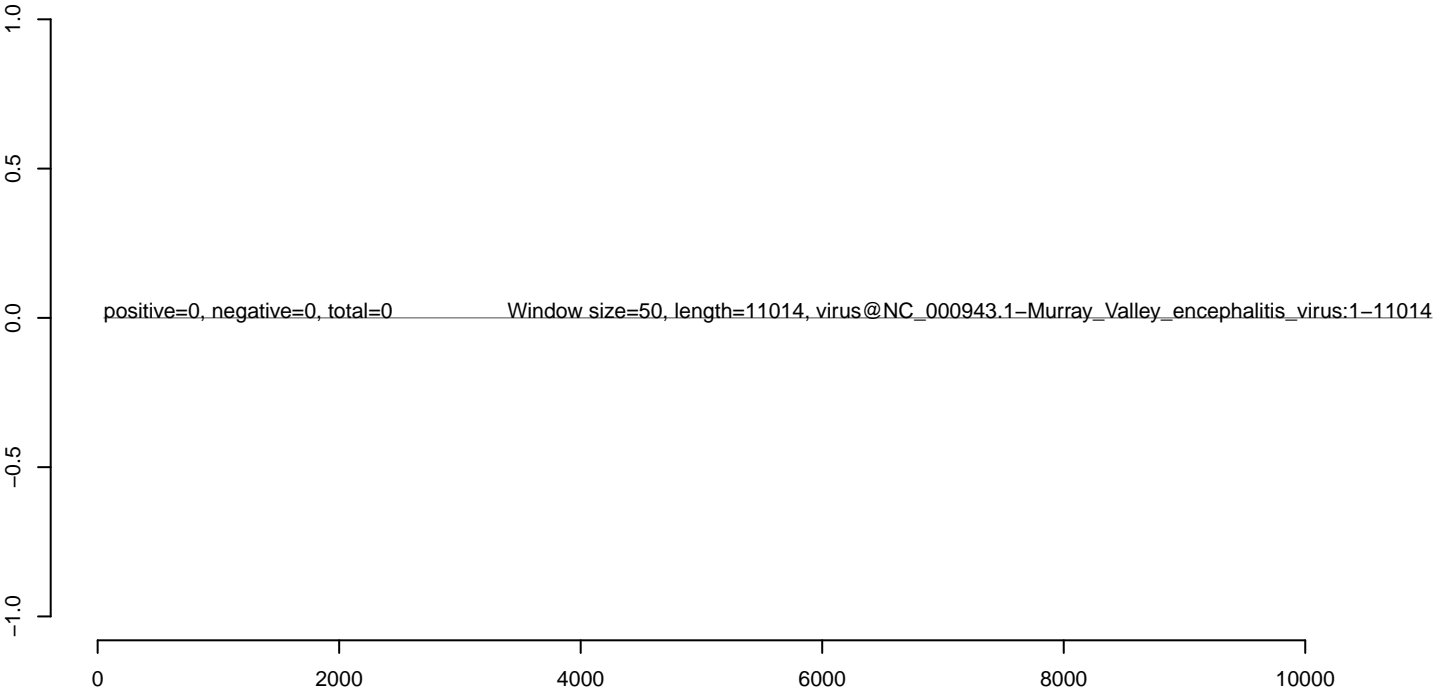
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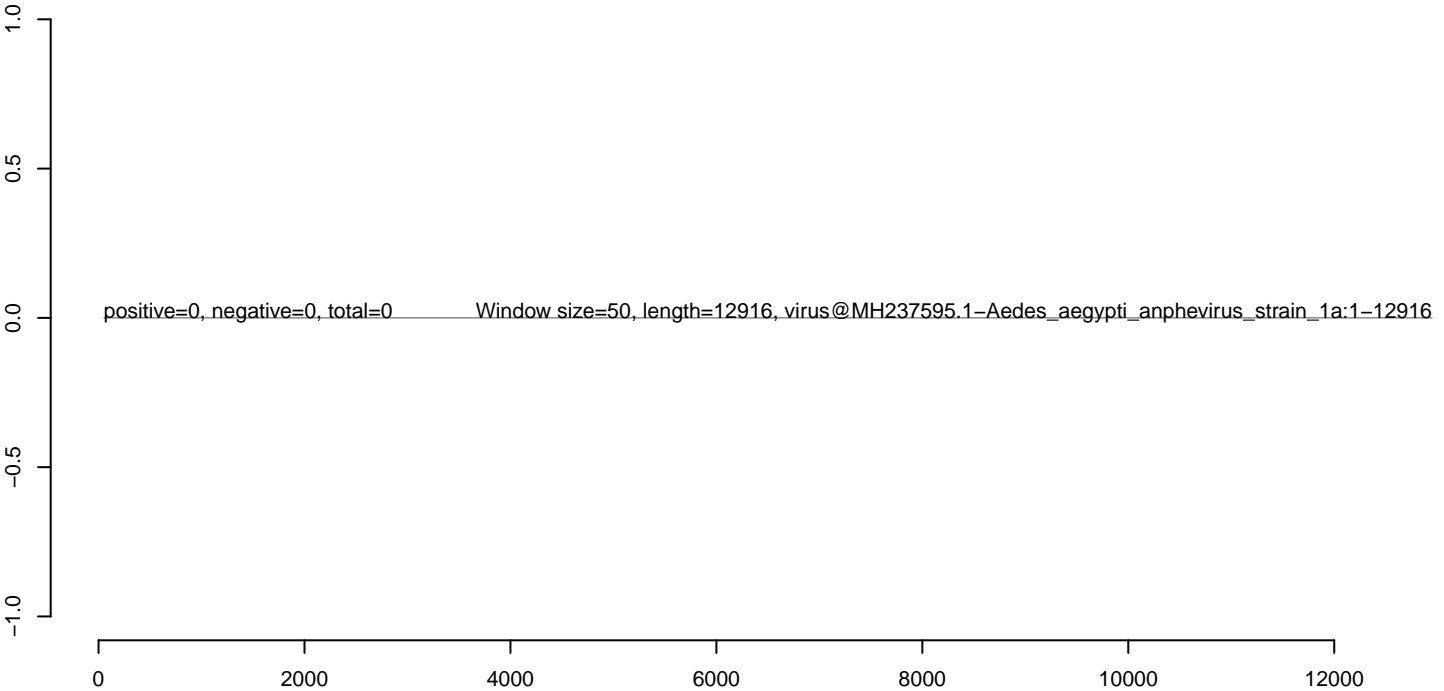
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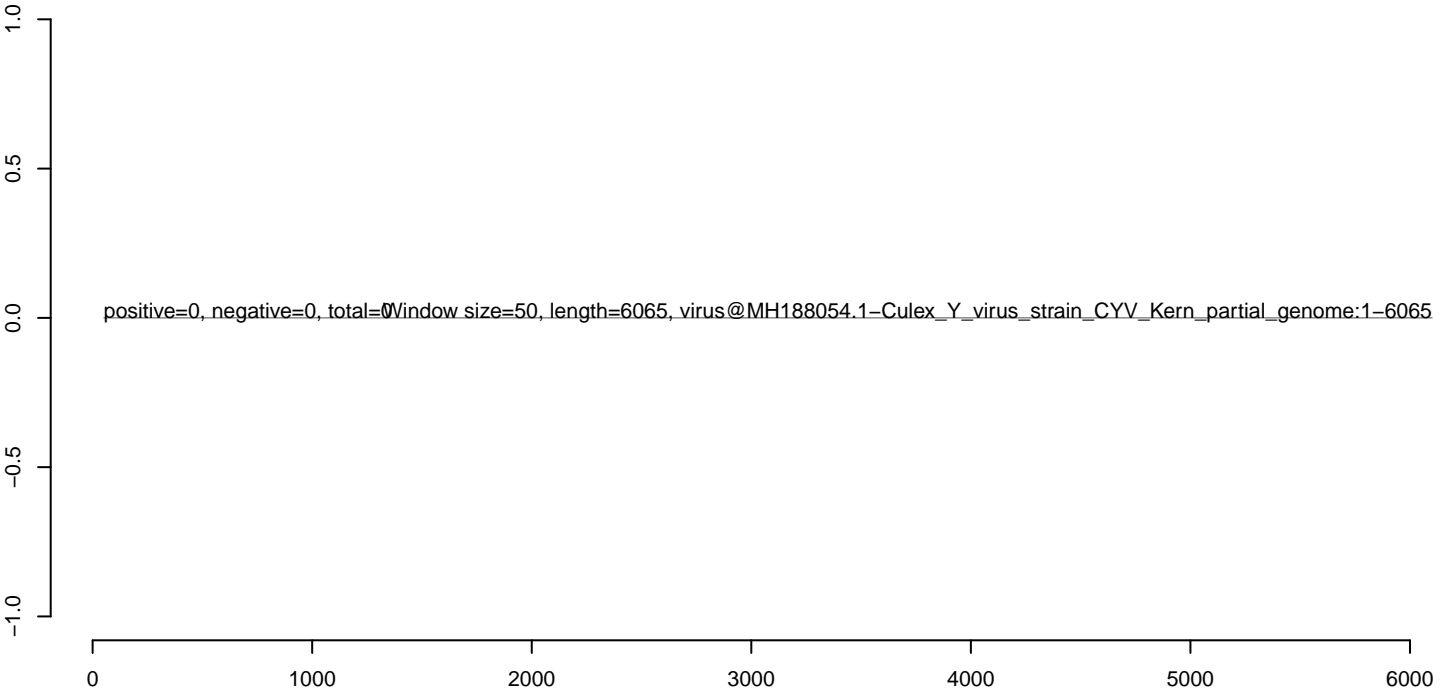
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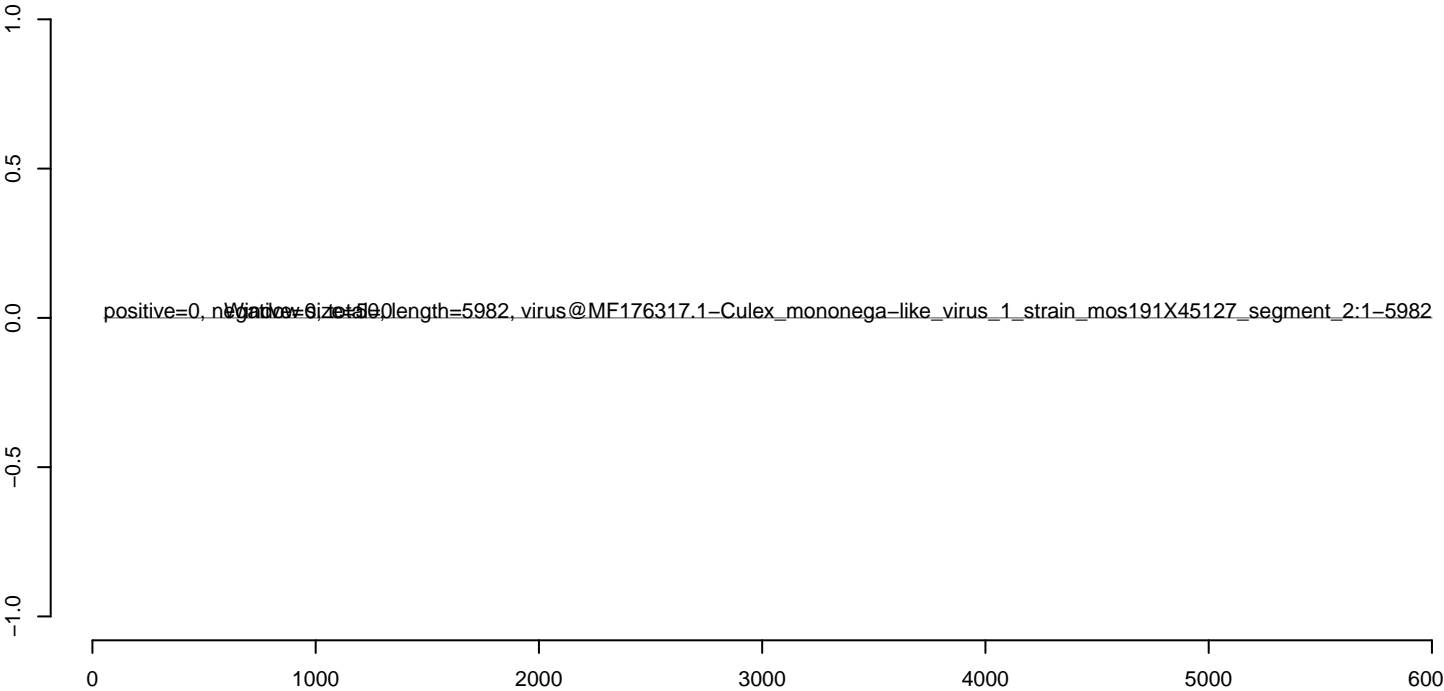
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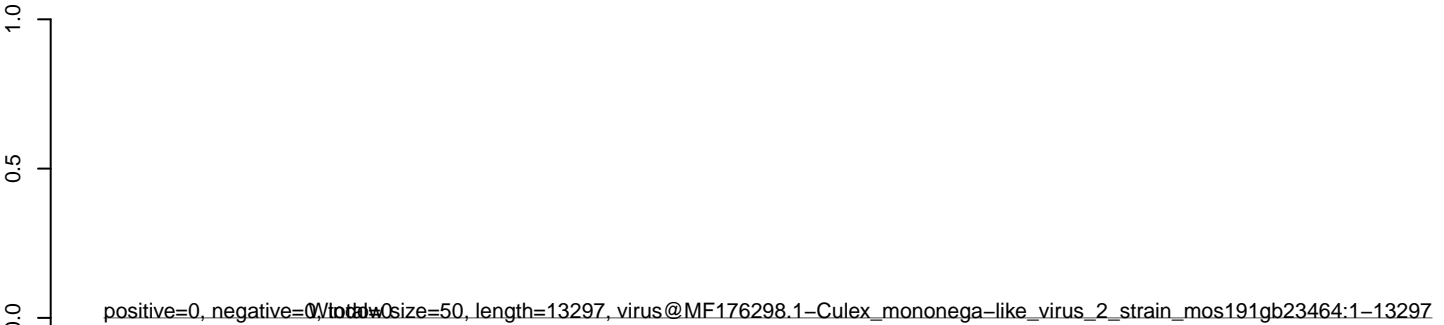
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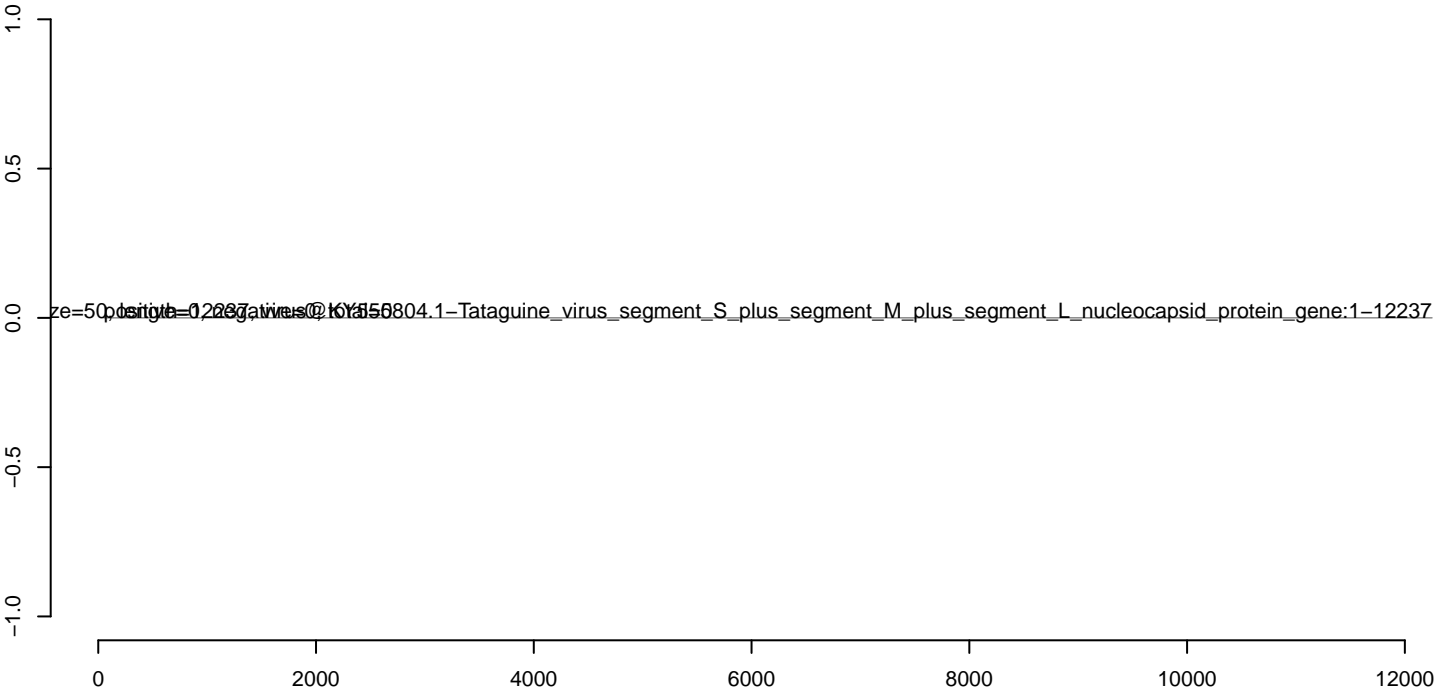
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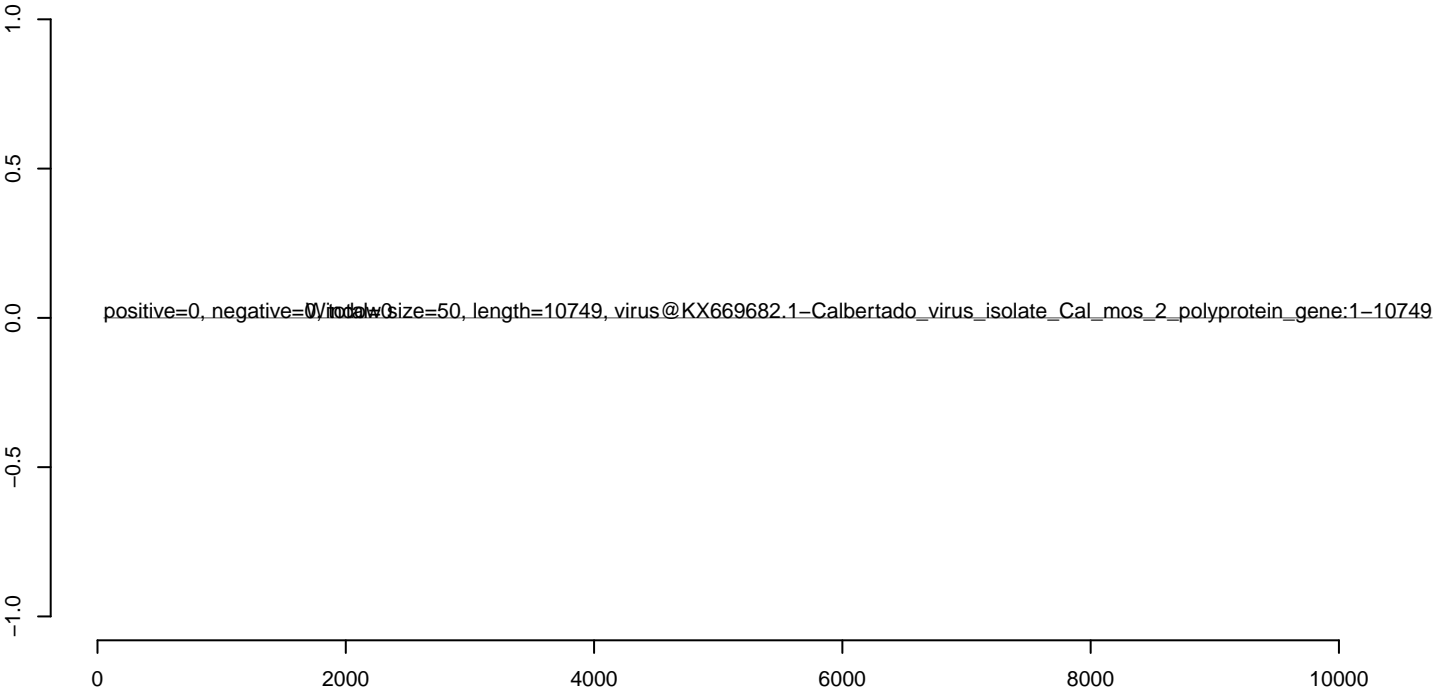
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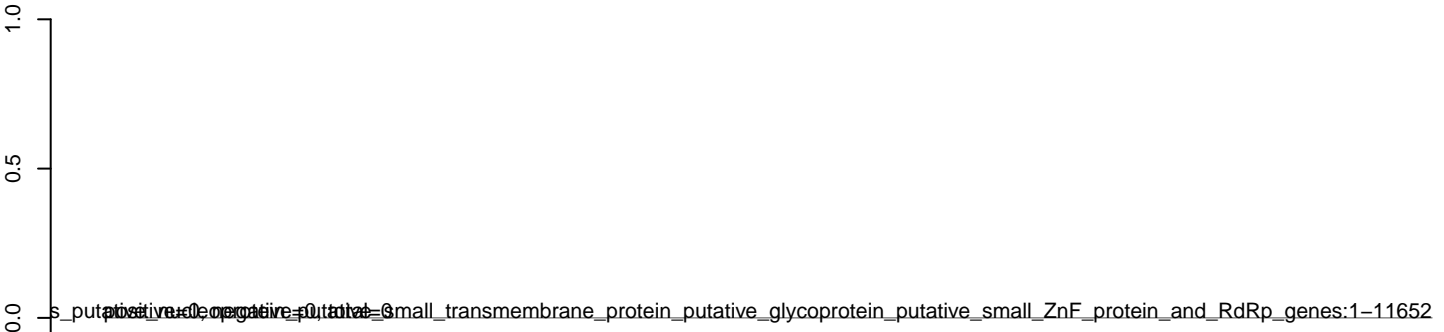
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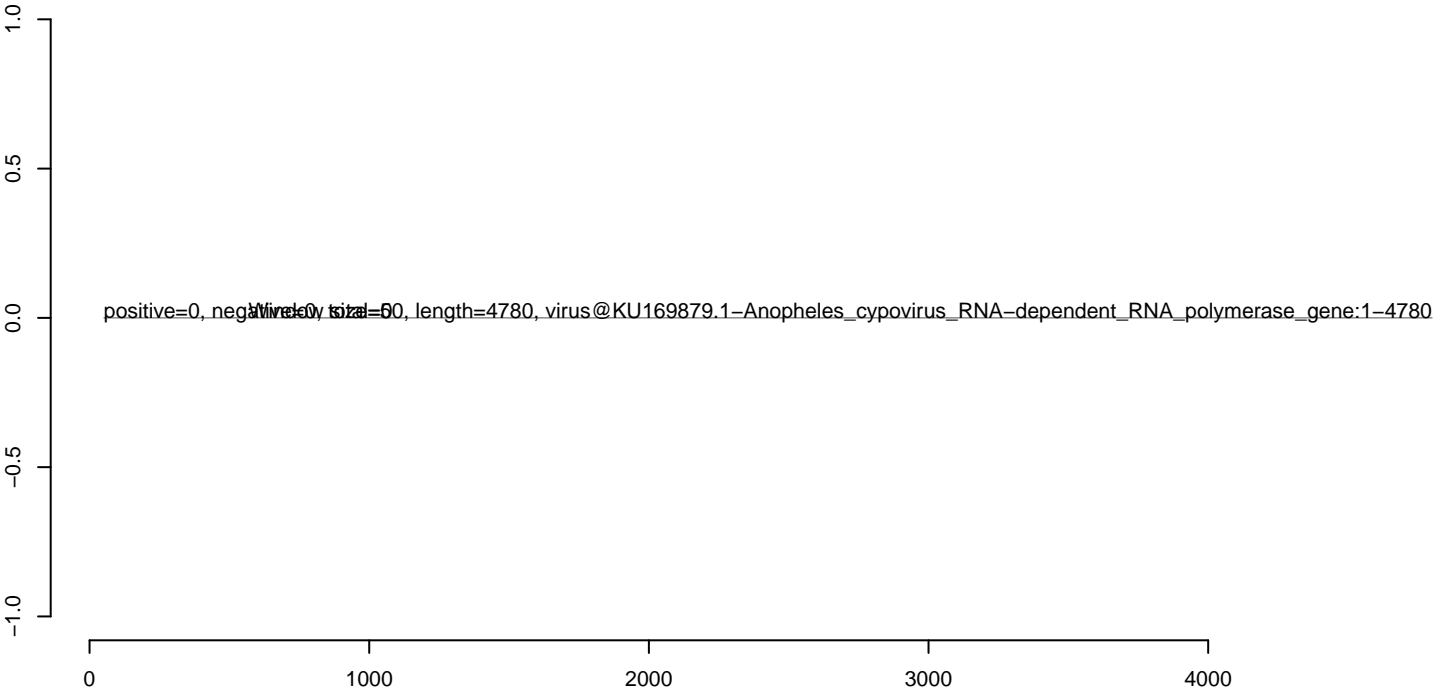
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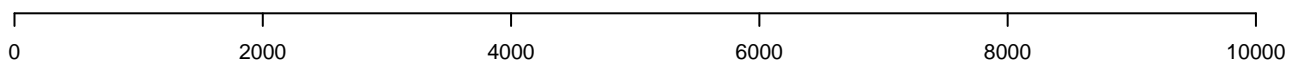
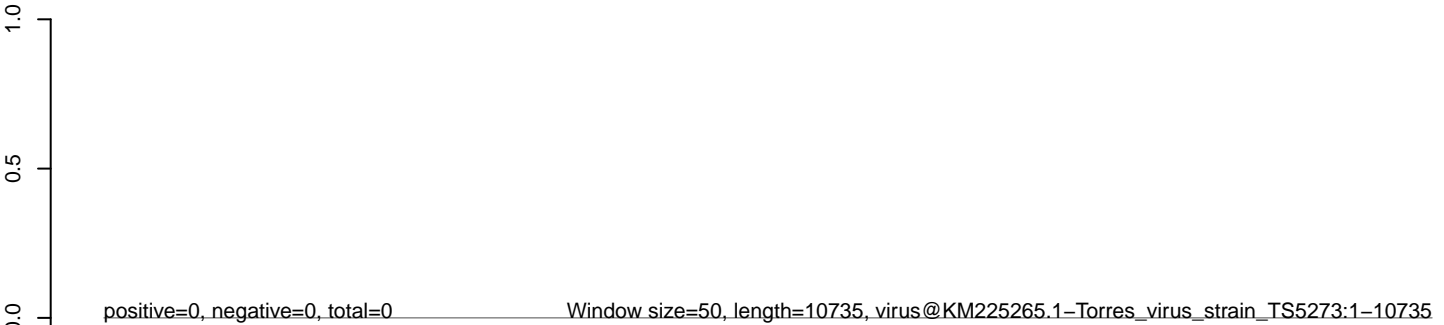
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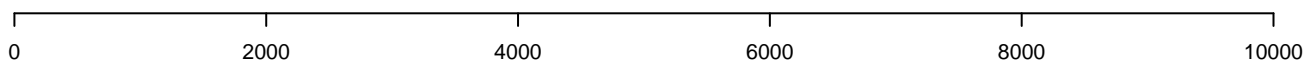
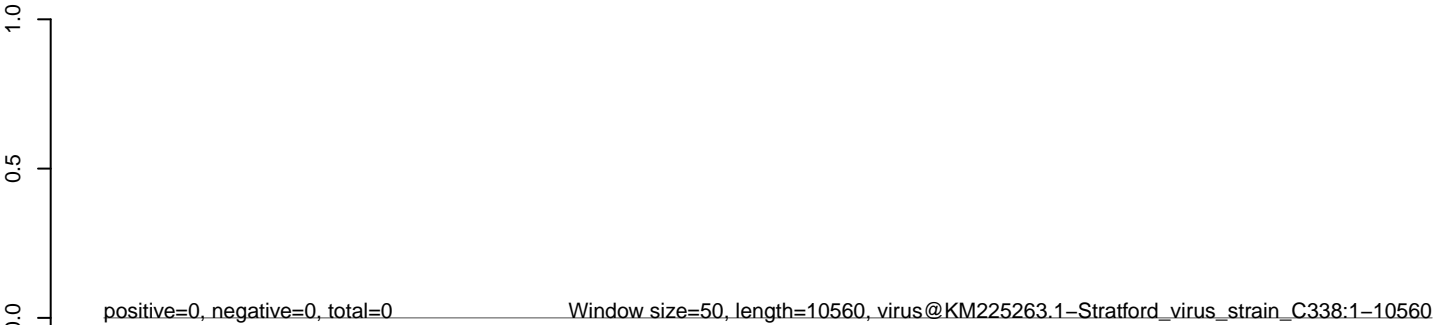
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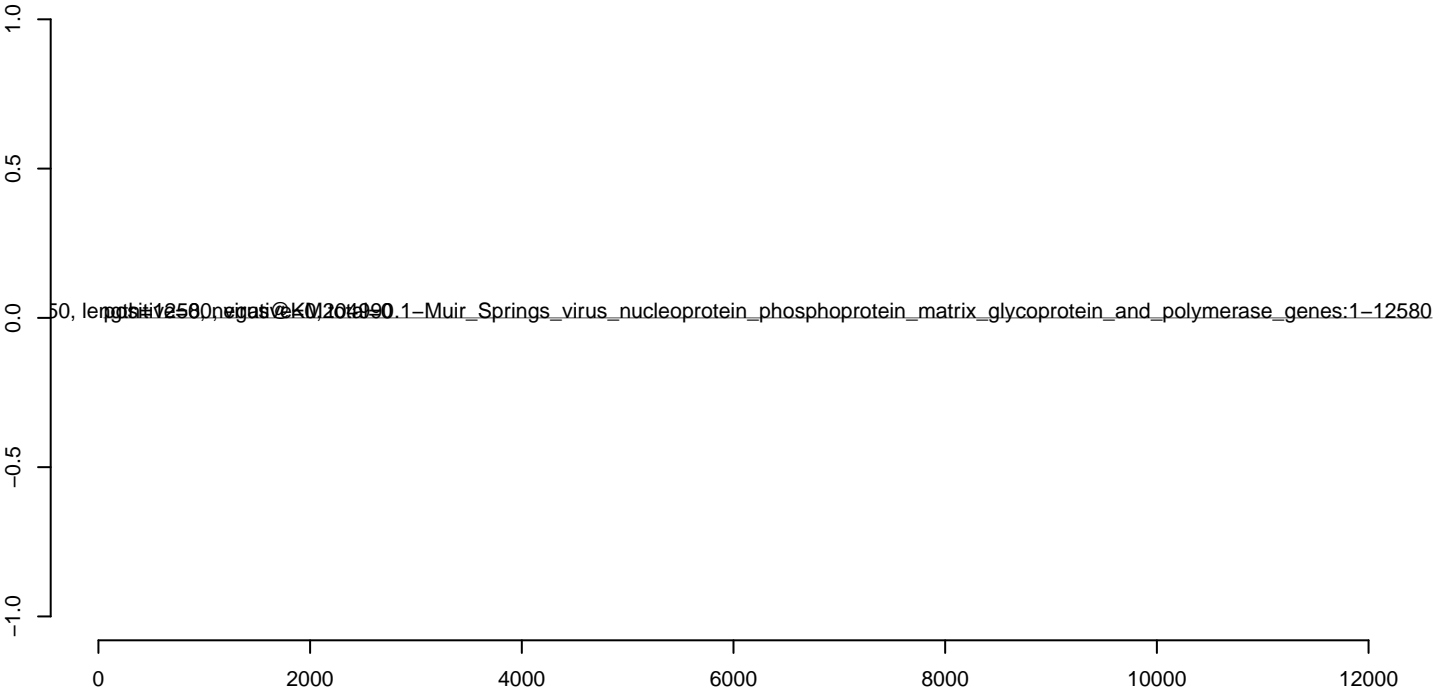
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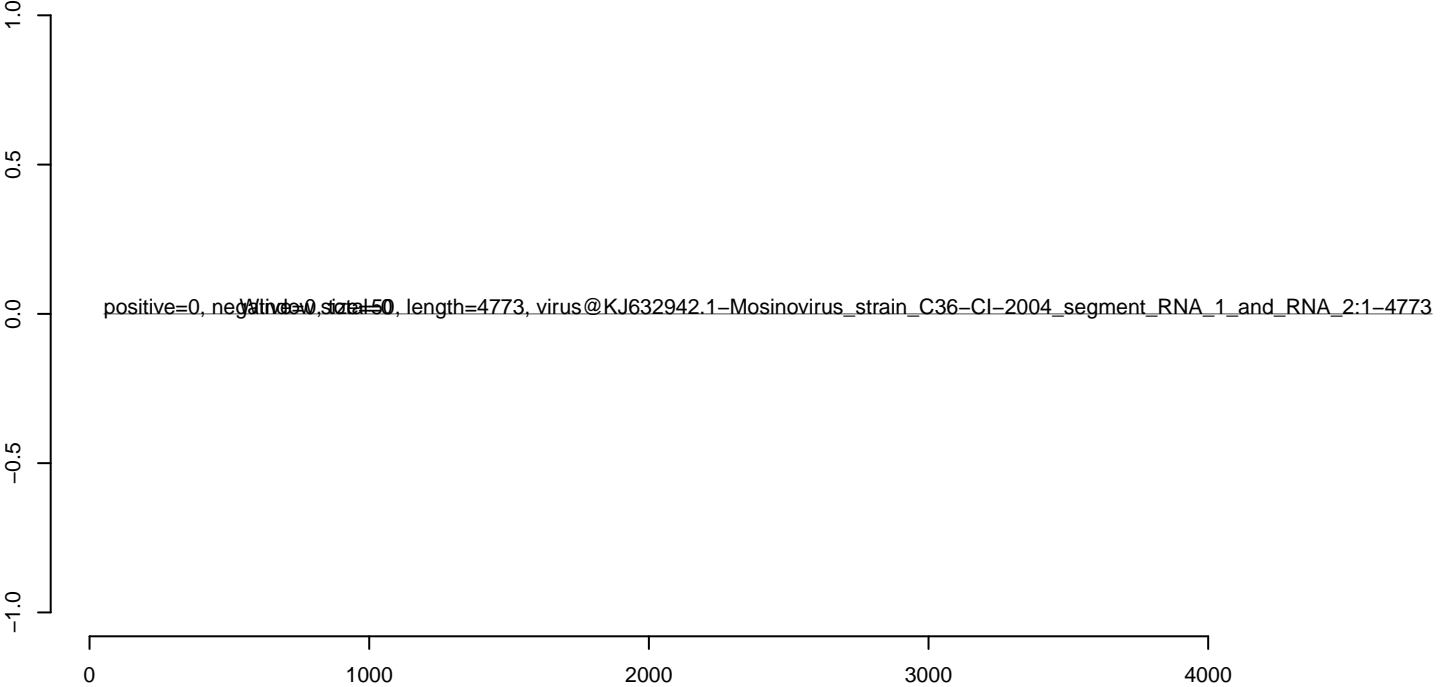
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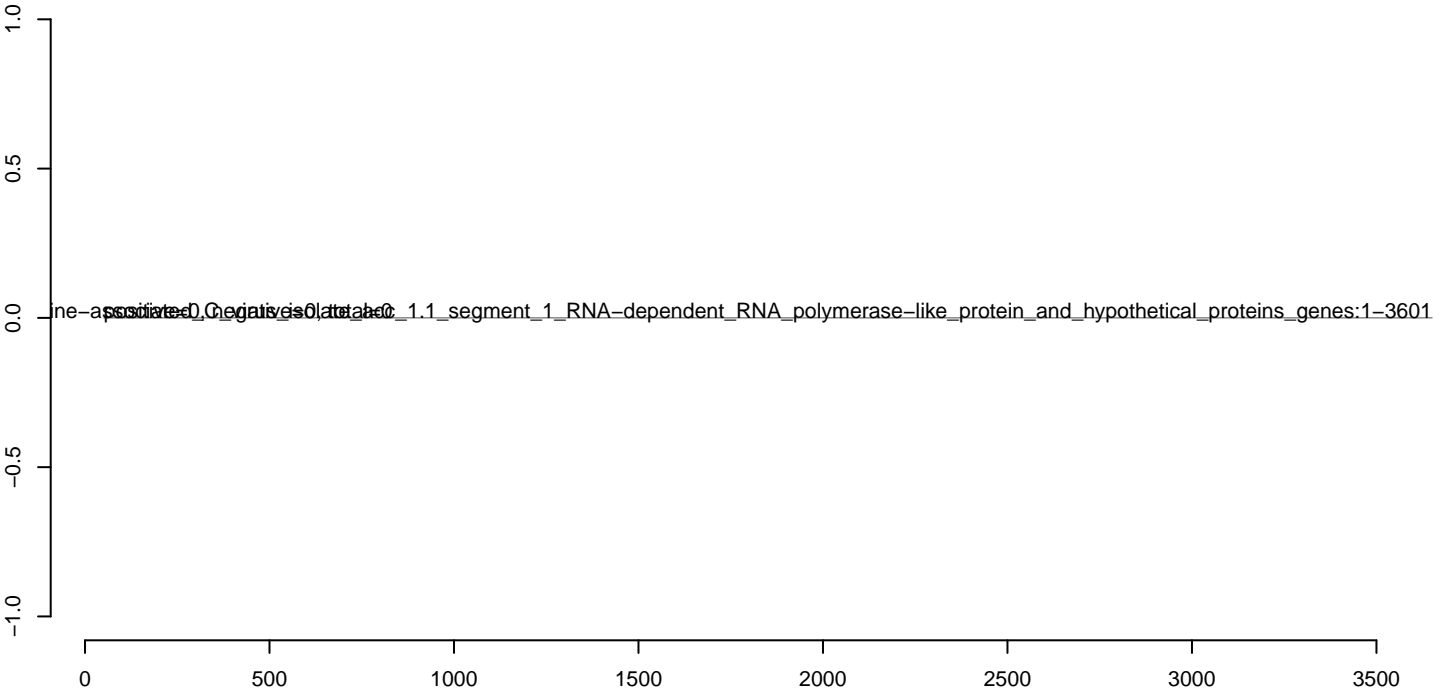
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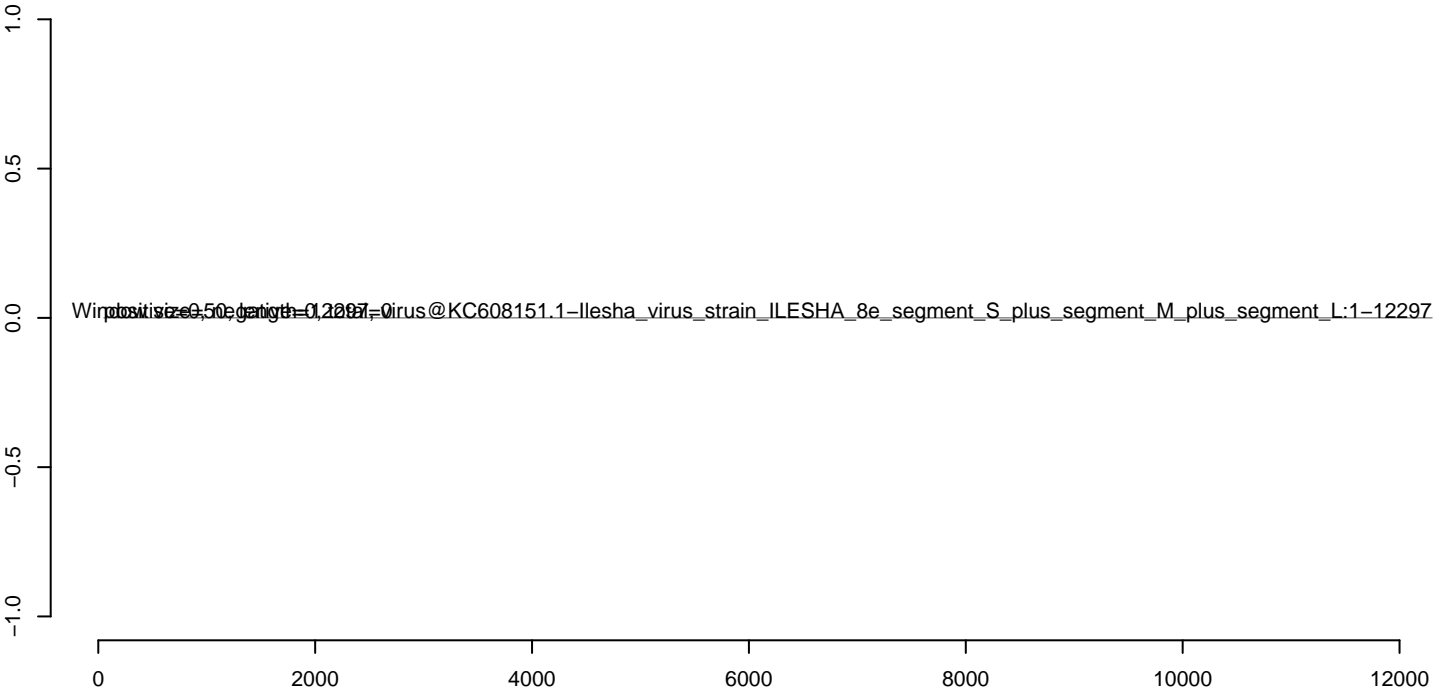
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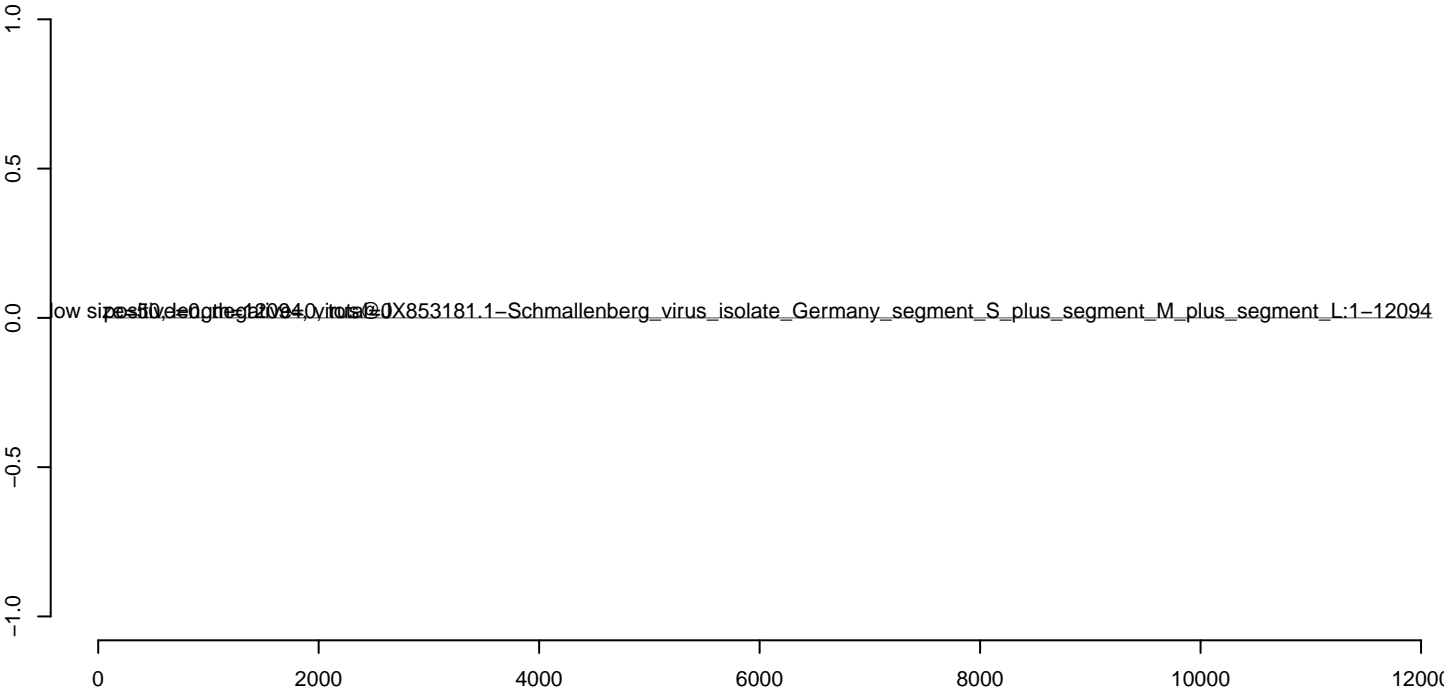
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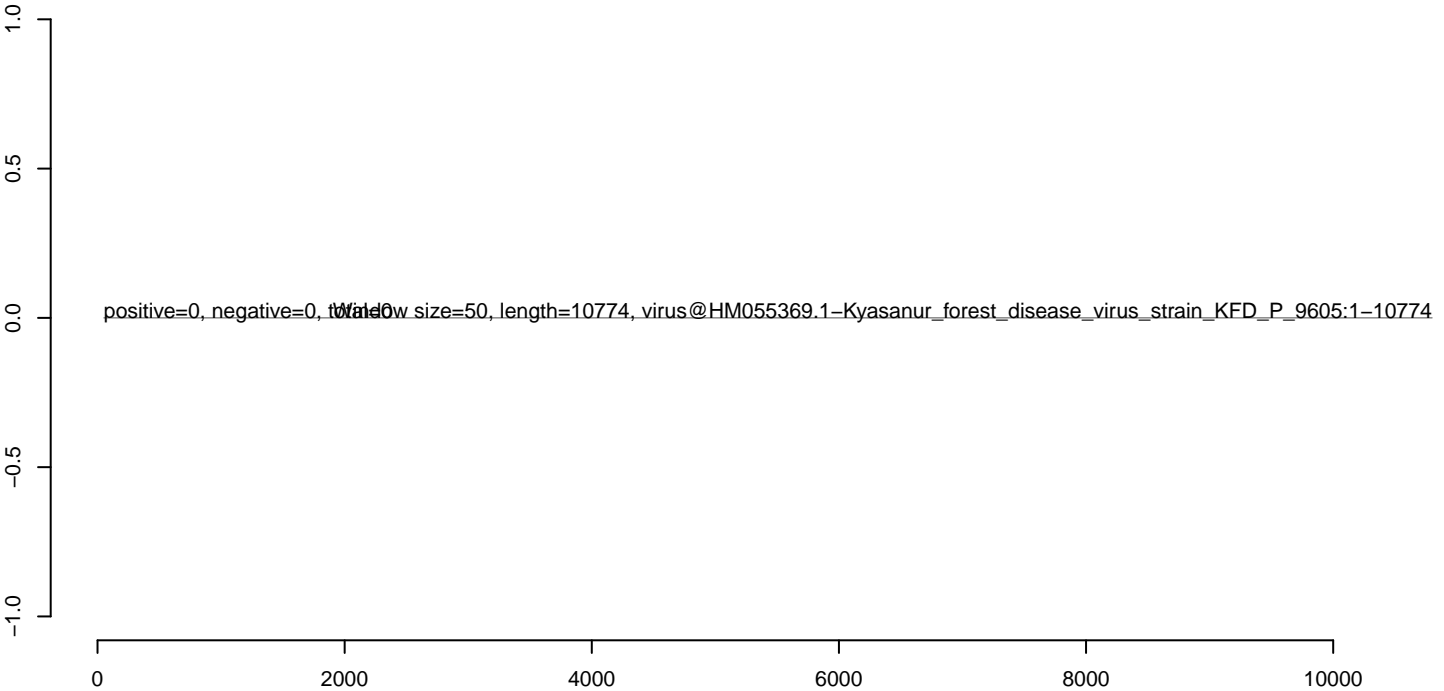
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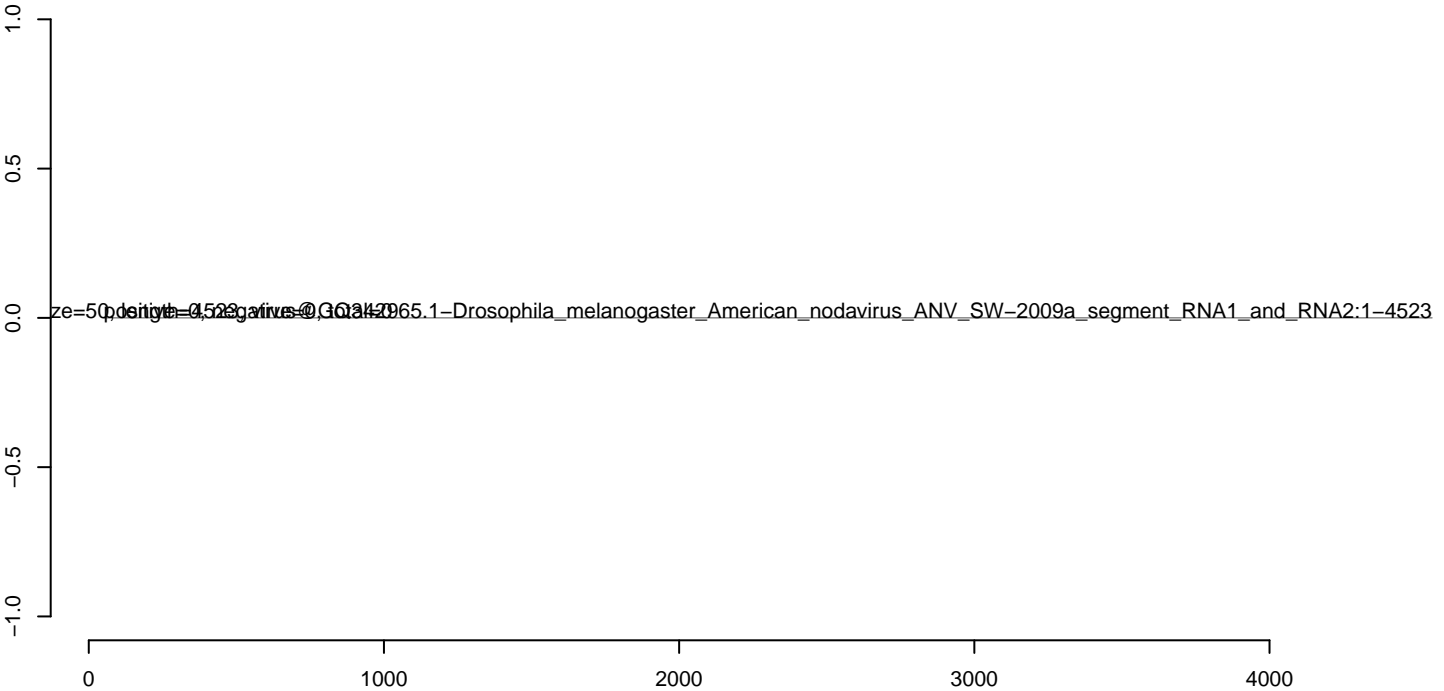
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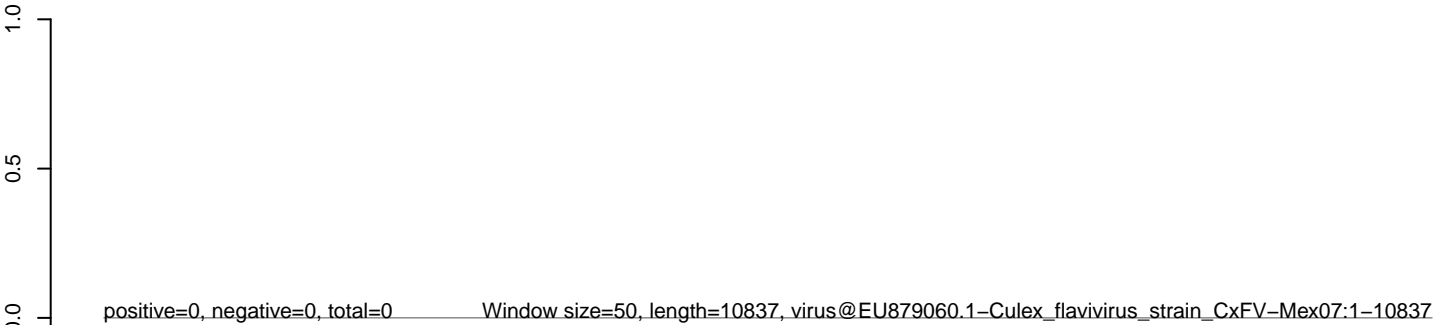
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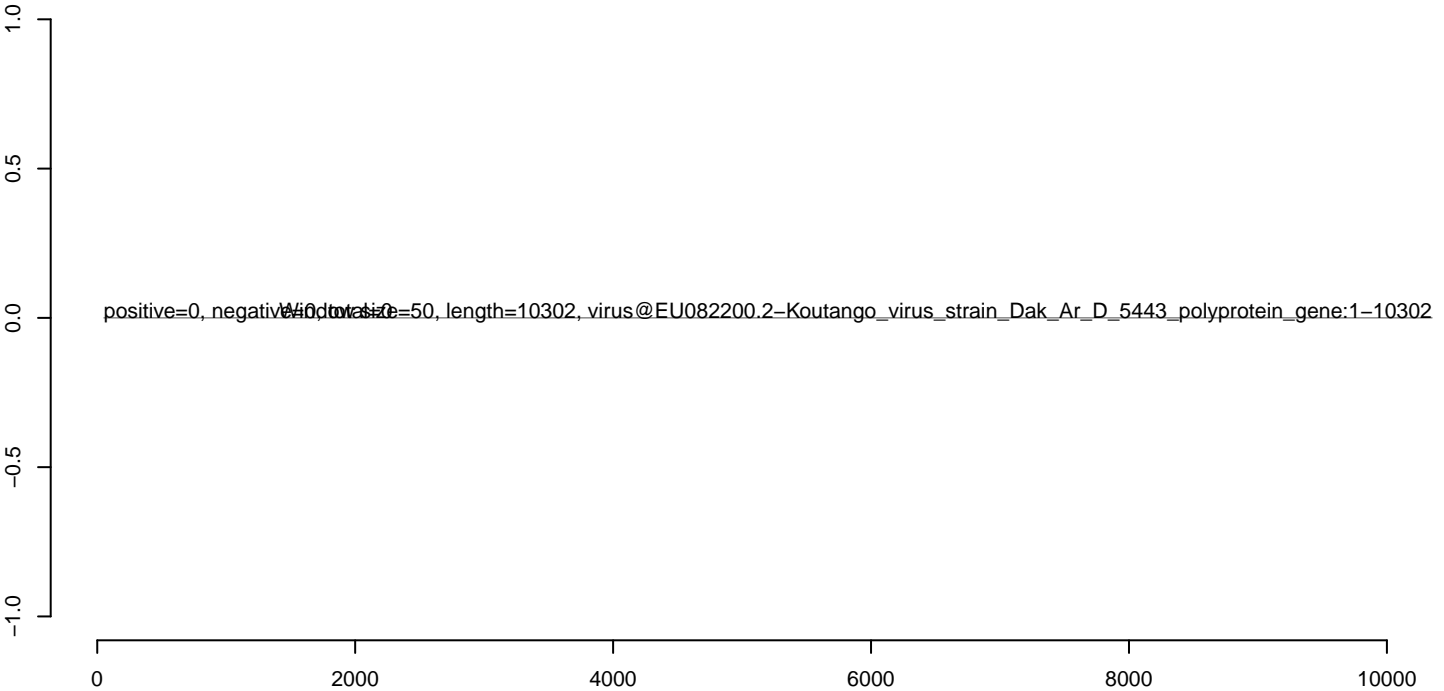
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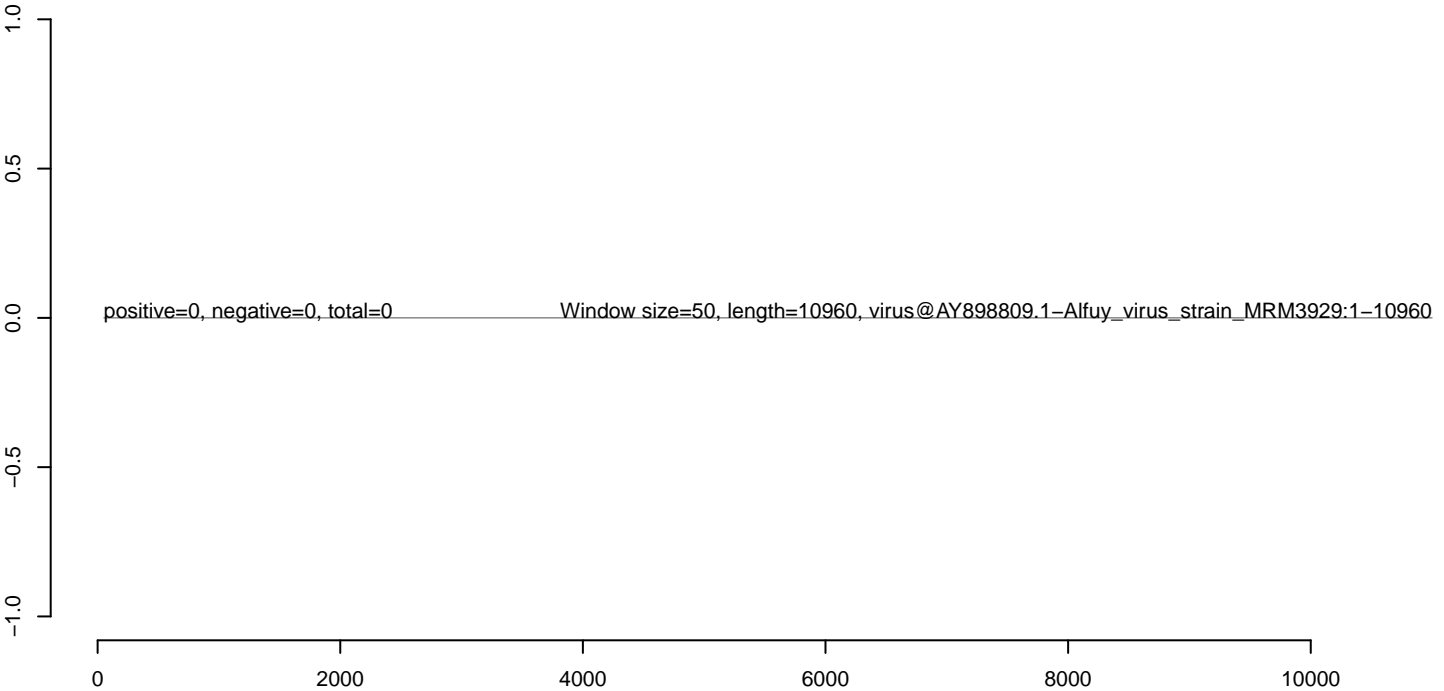
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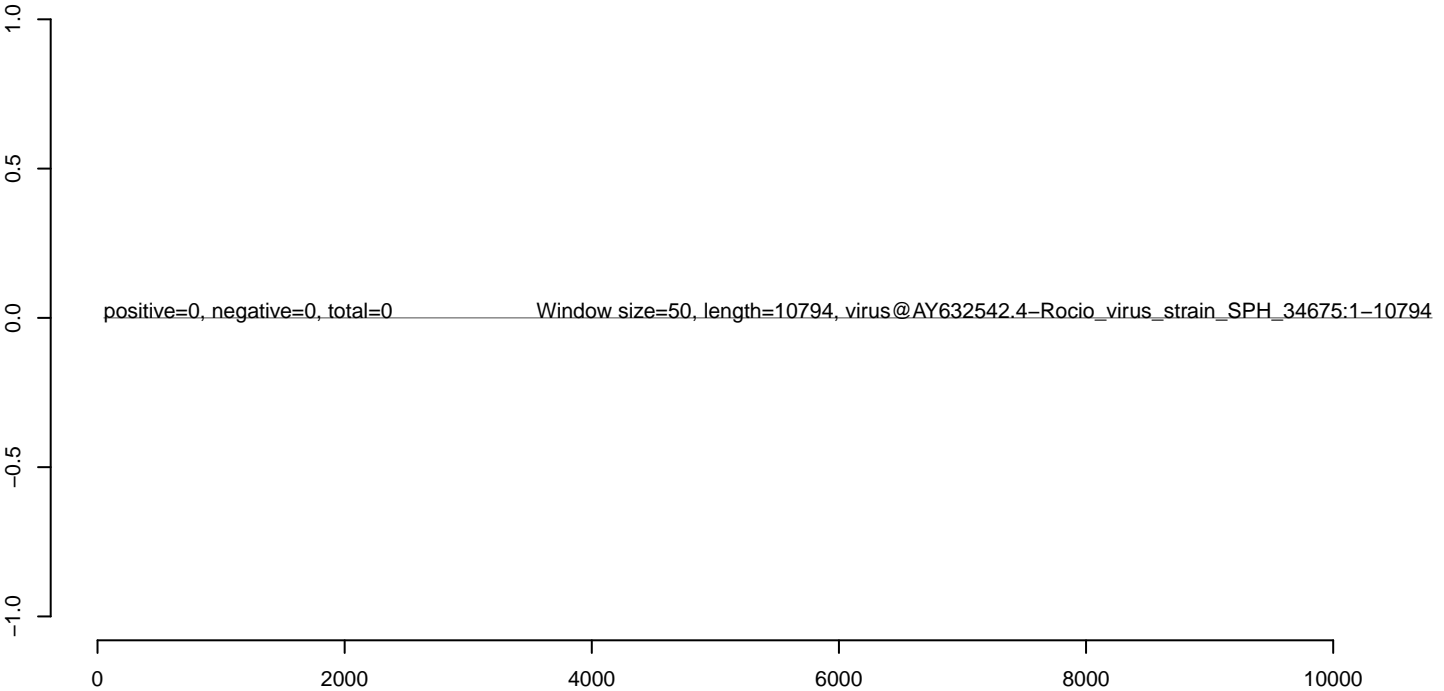
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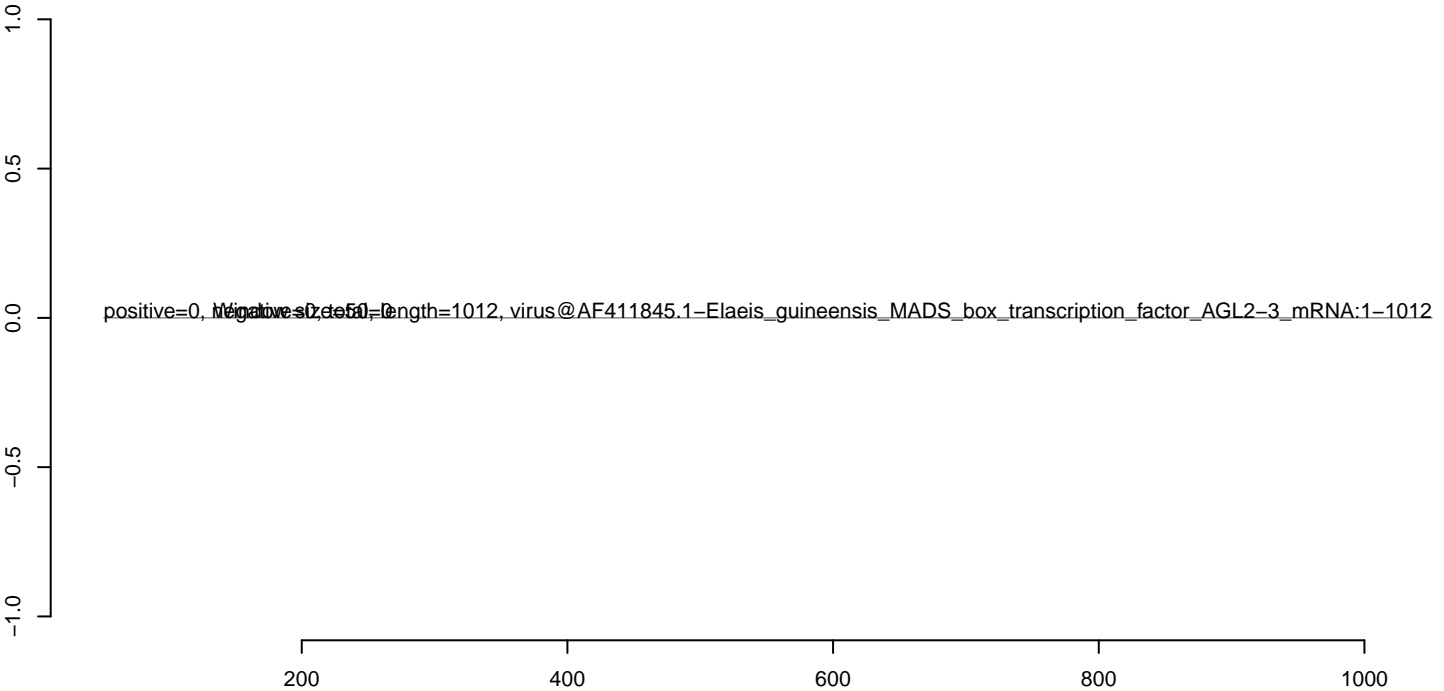
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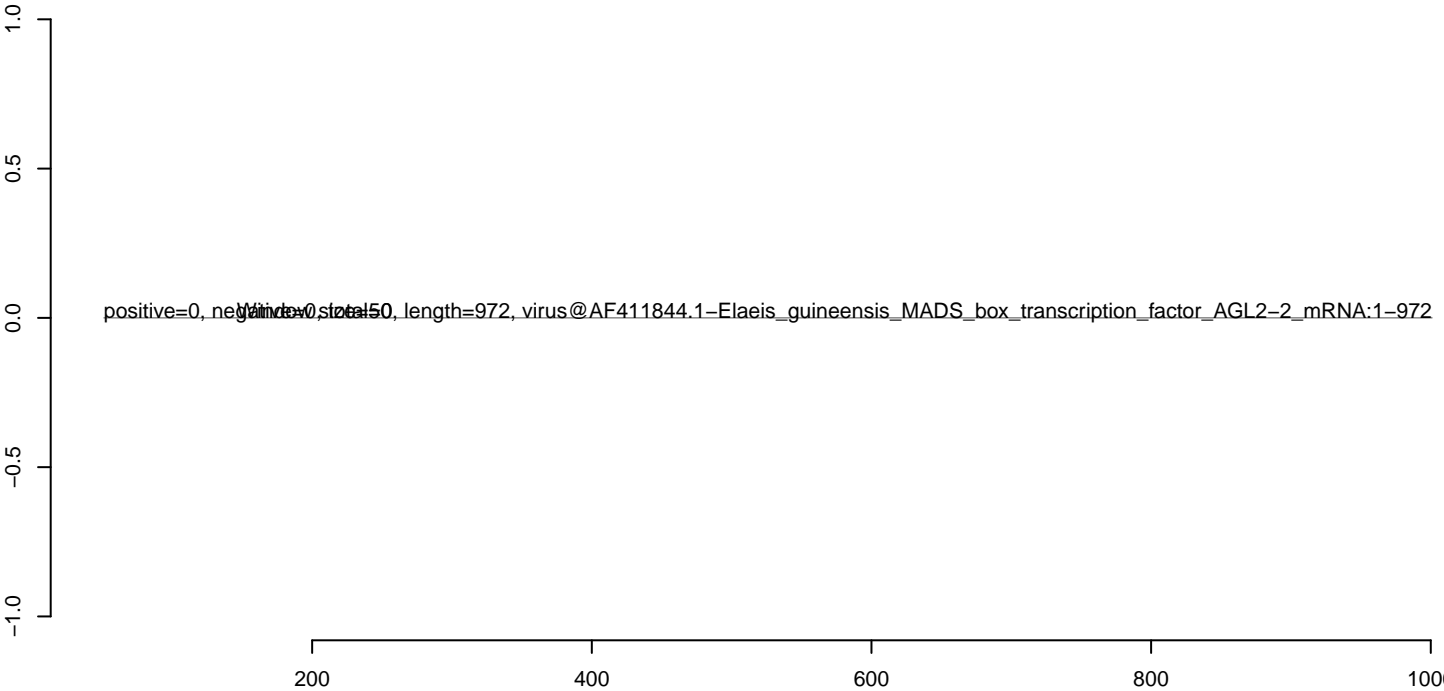
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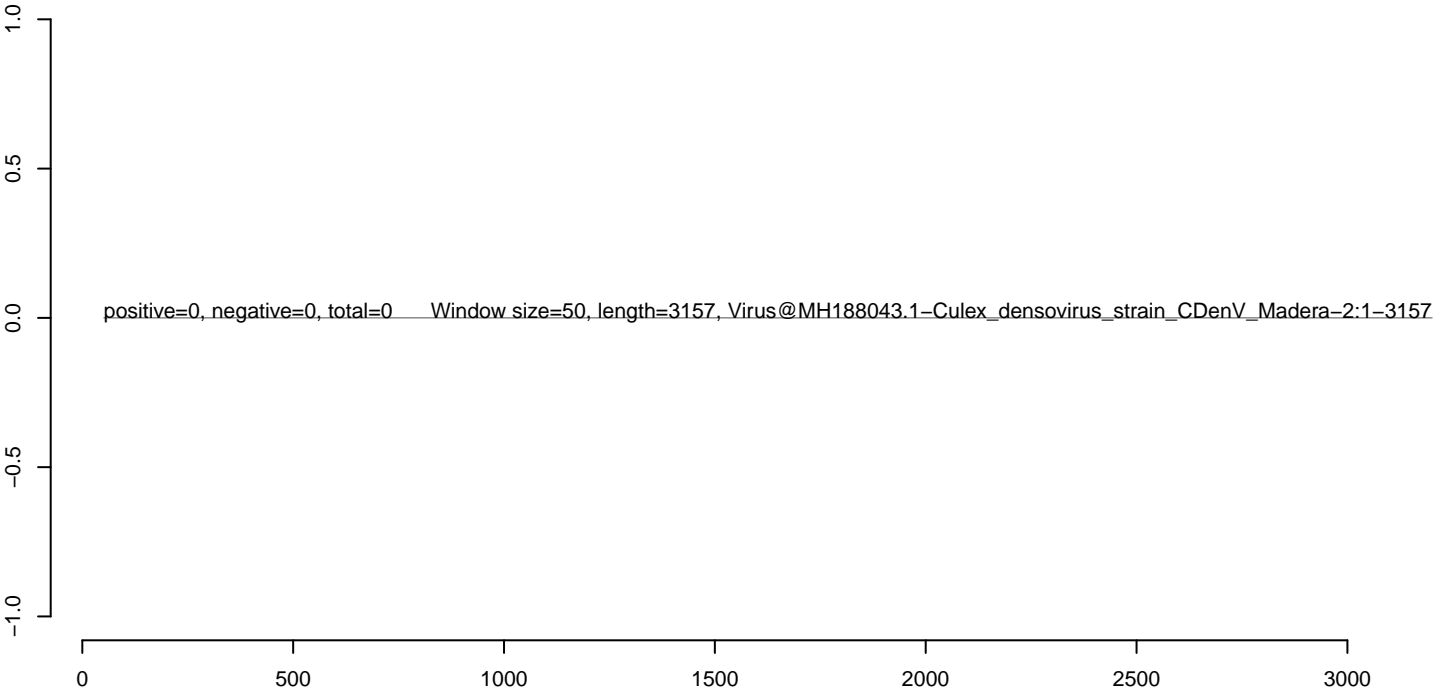
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