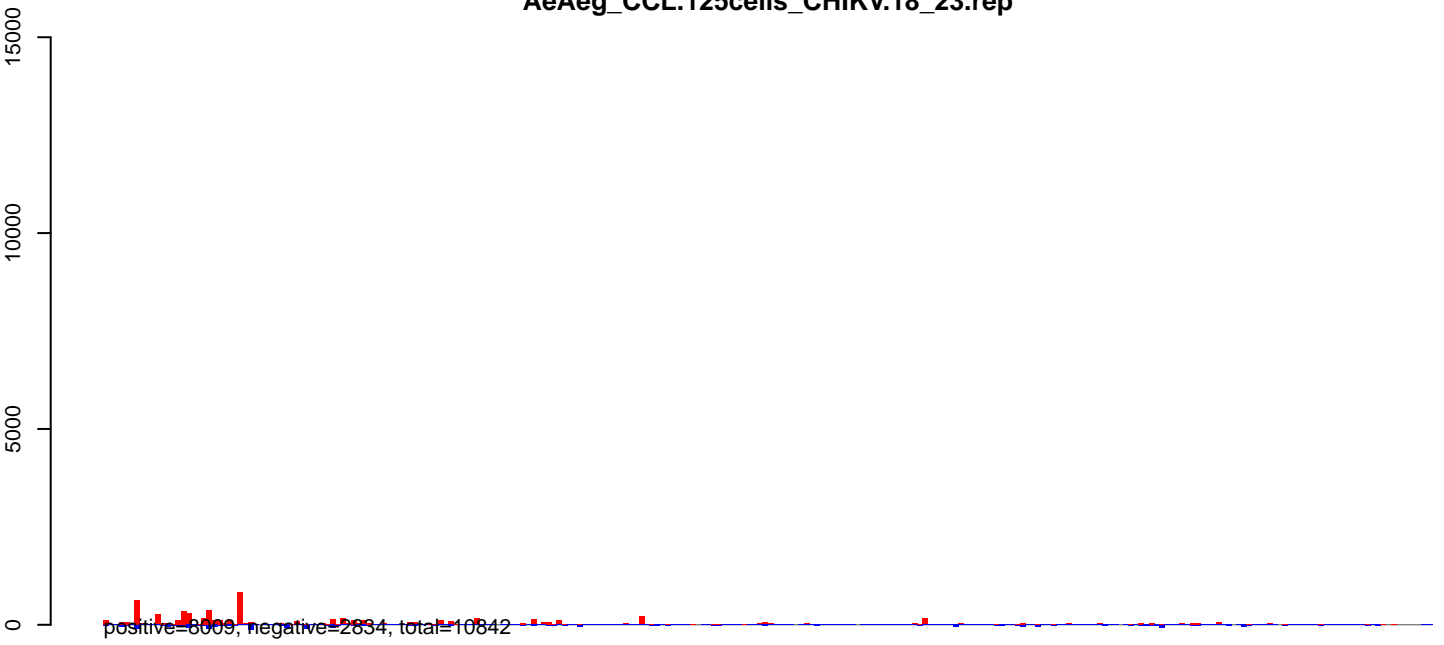
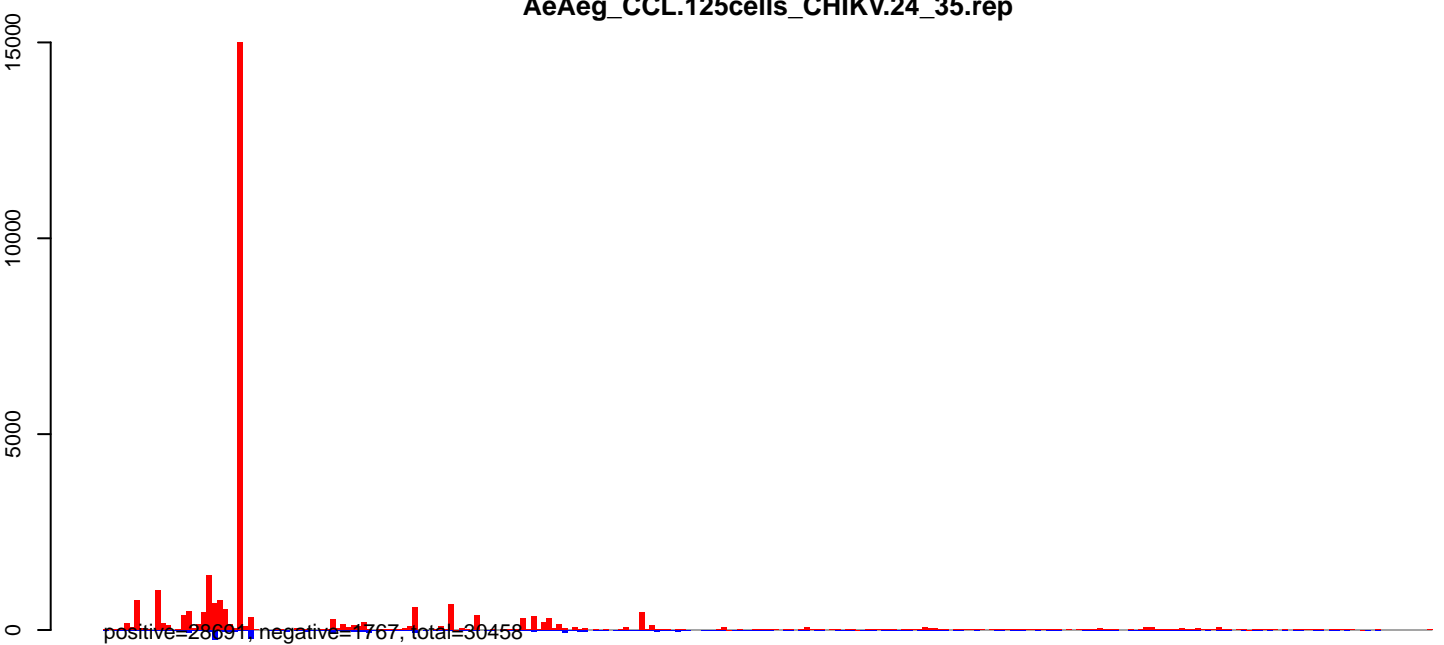


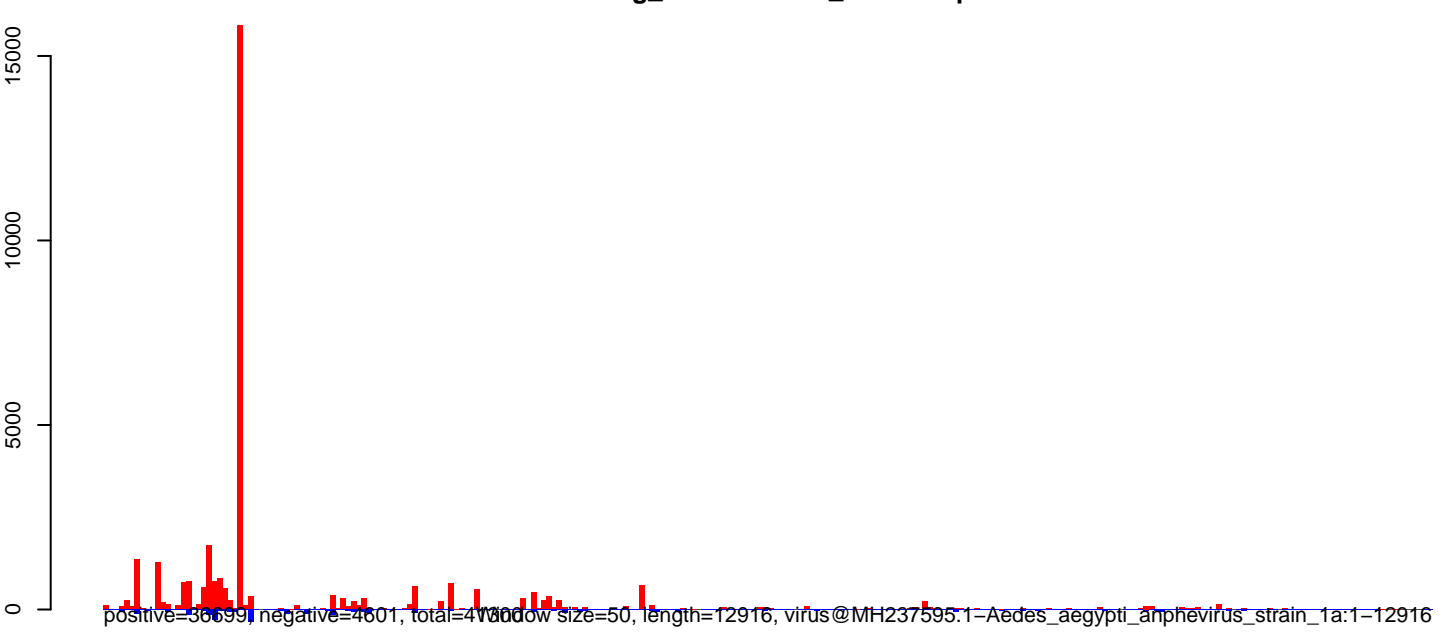
AeAeg_CCL.125cells_CHIKV.18_23.rep



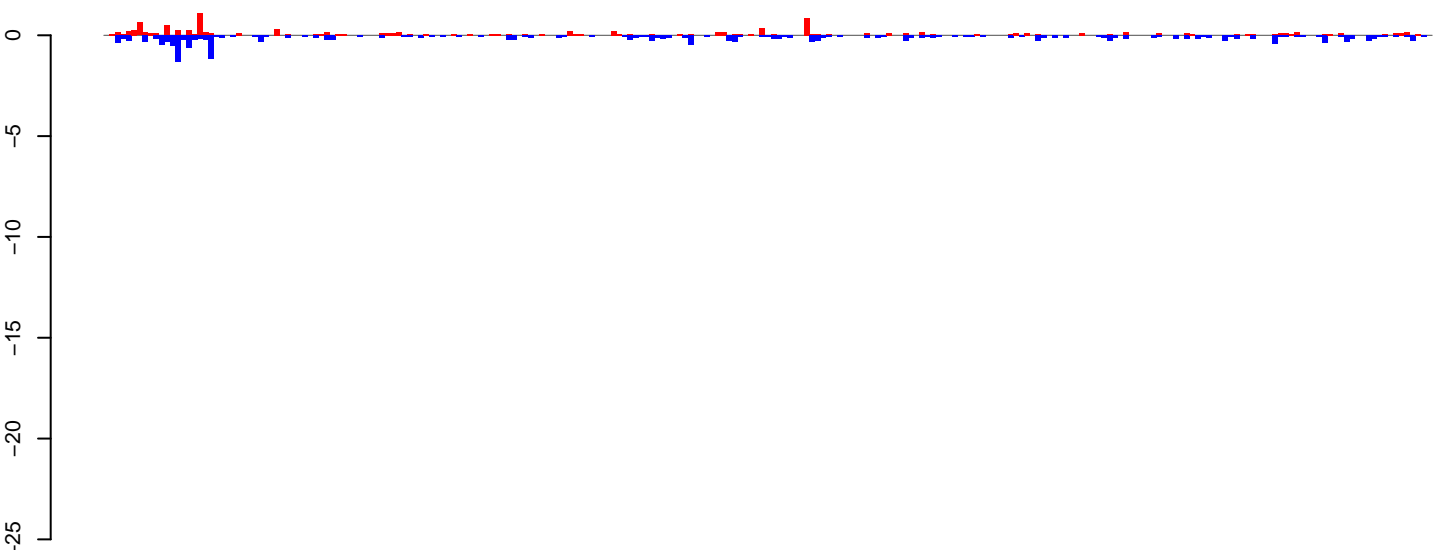
AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep

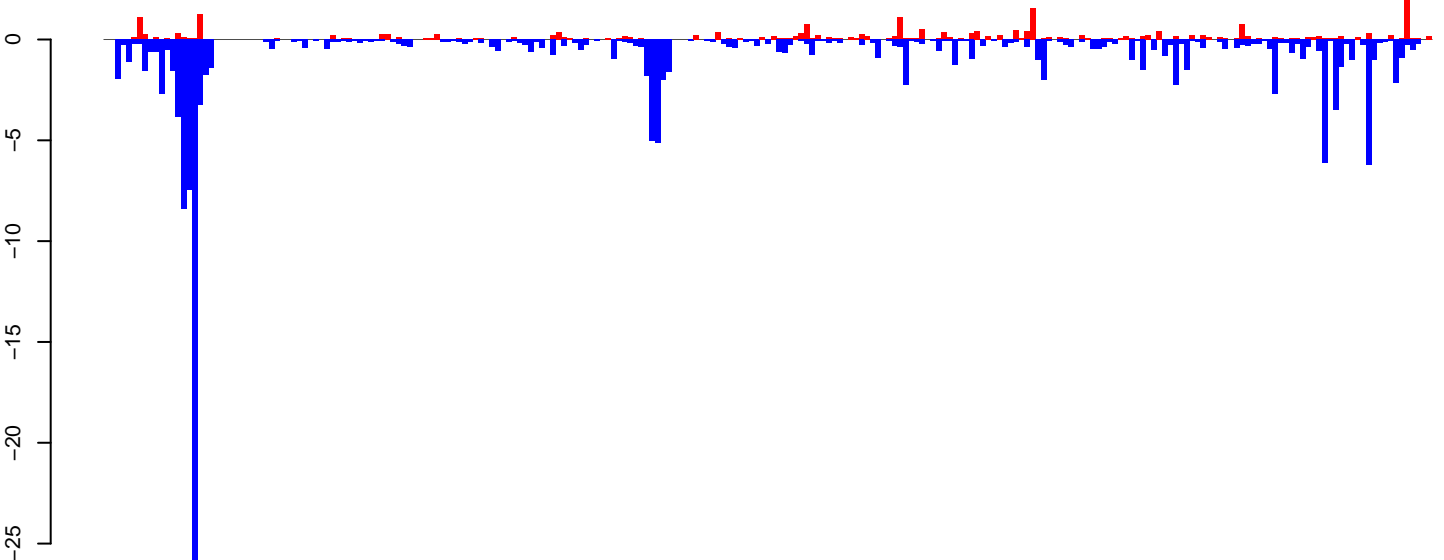


AeAeg_CCL.125cells_CHIKV.18_23.rep



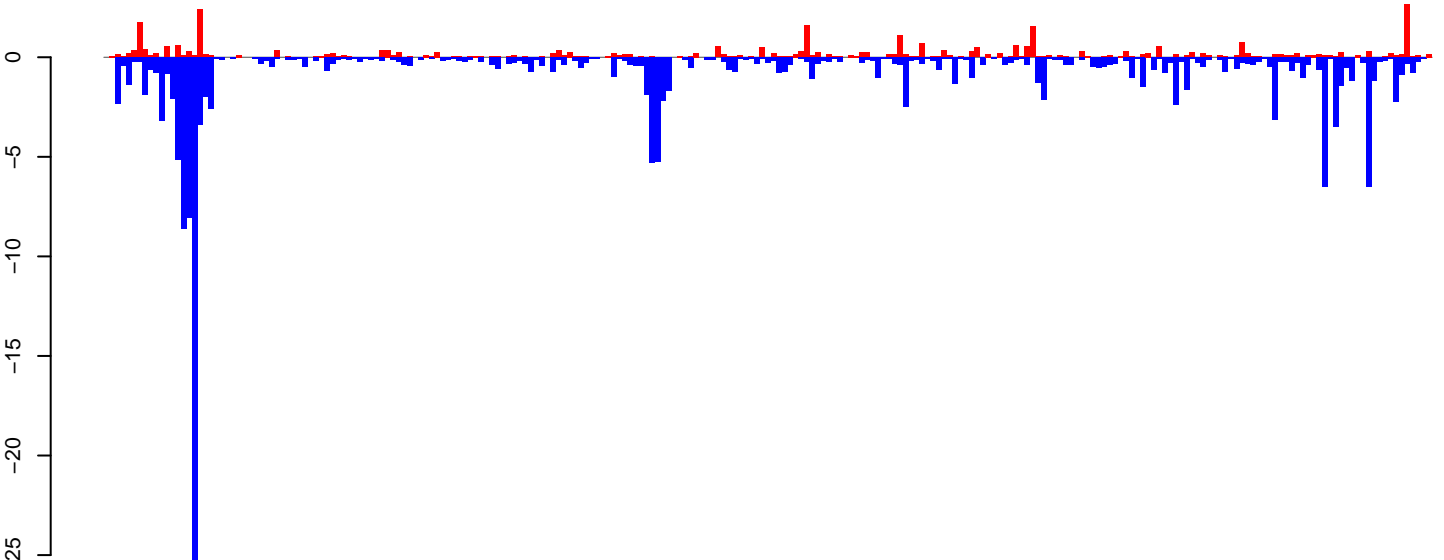
positive=12, negative=20, total=31

AeAeg_CCL.125cells_CHIKV.24_35.rep

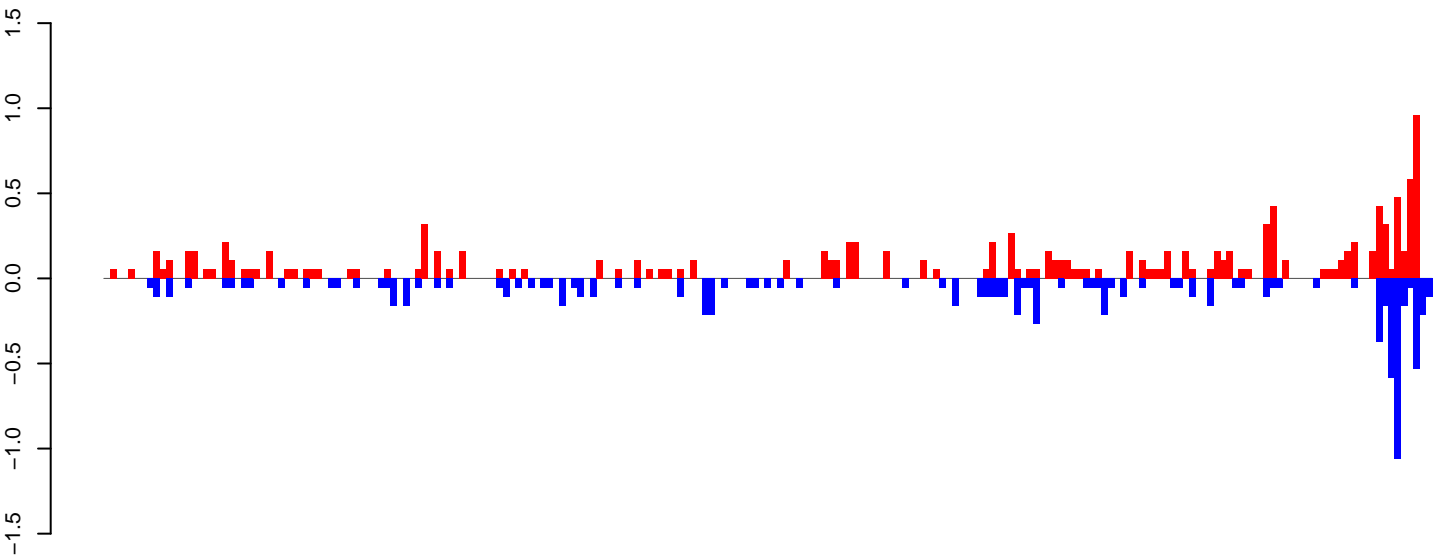


positive=24, negative=152, total=177

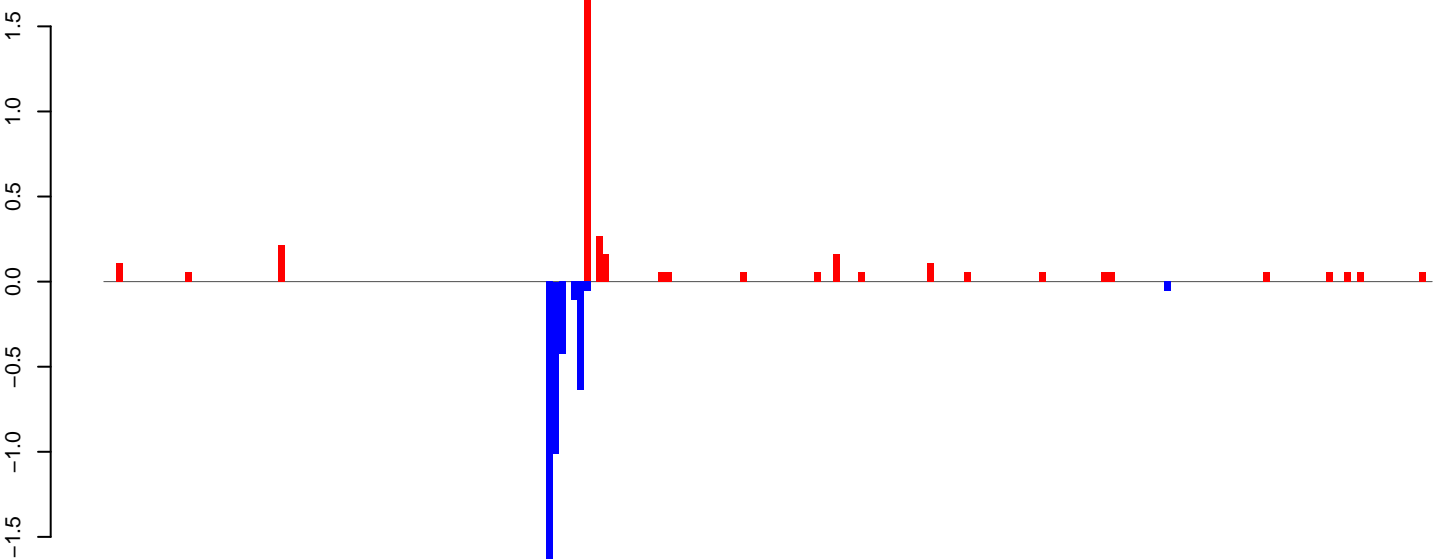
AeAeg_CCL.125cells_CHIKV.rep



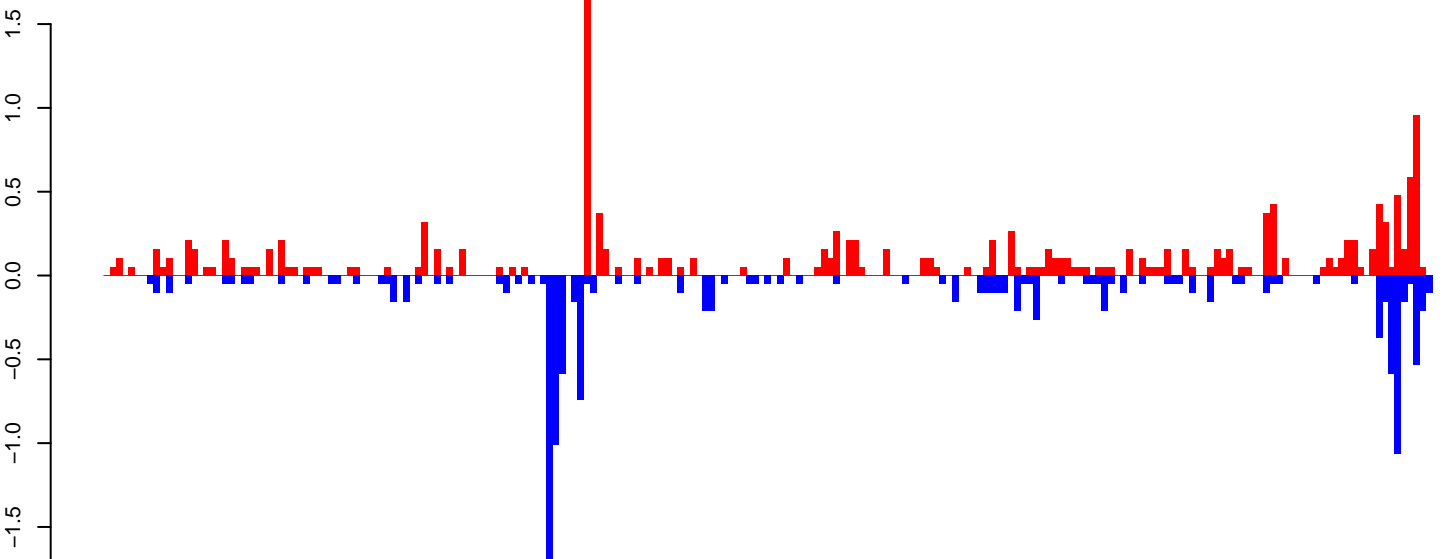
AeAeg_CCL.125cells_CHIKV.18_23.rep



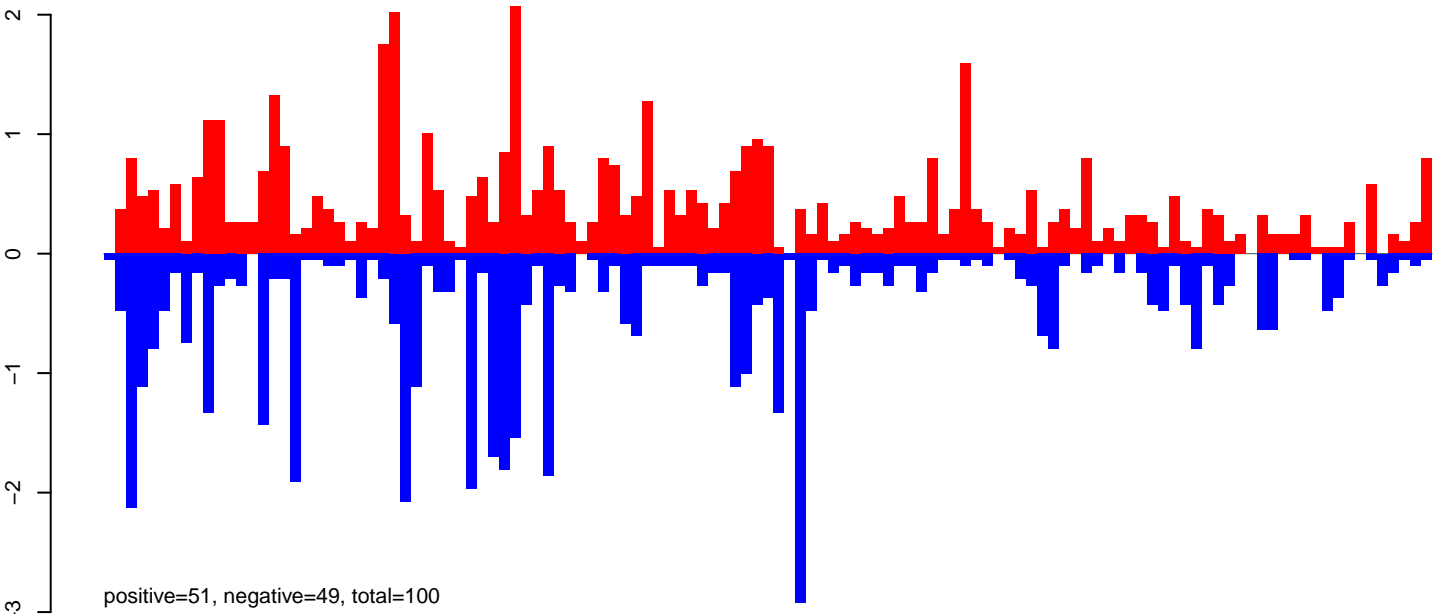
AeAeg_CCL.125cells_CHIKV.24_35.rep



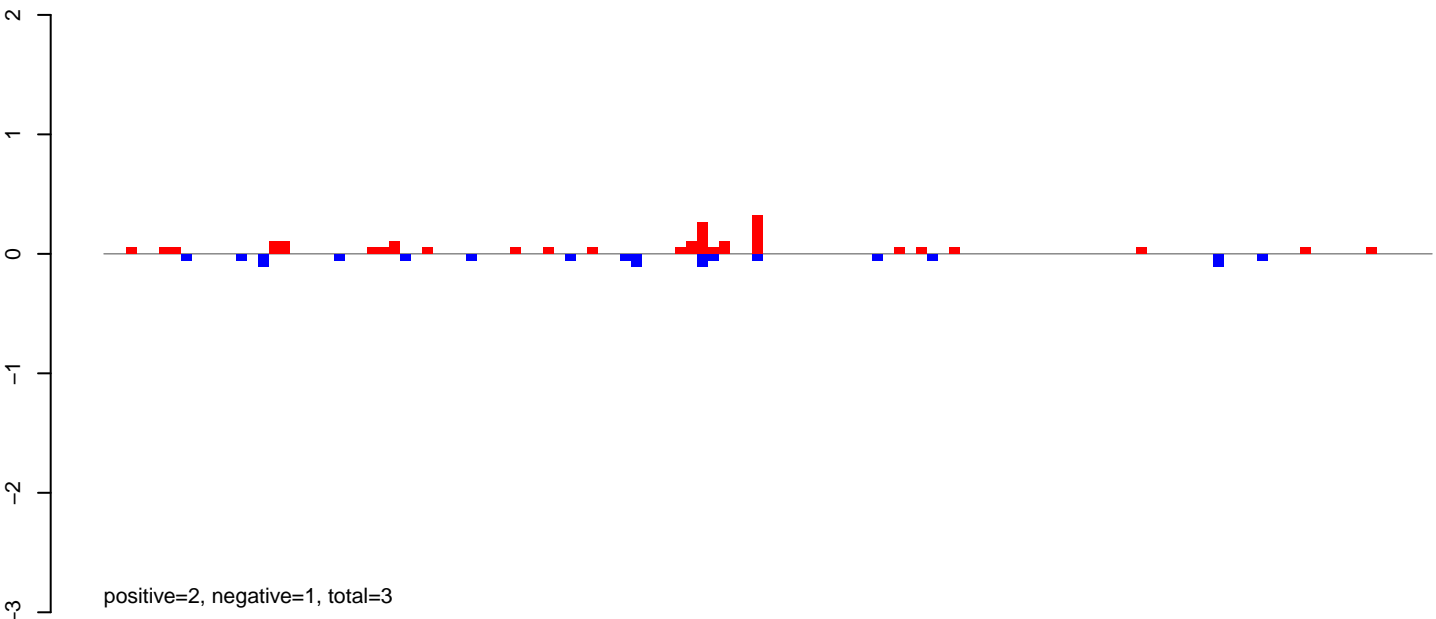
AeAeg_CCL.125cells_CHIKV.rep



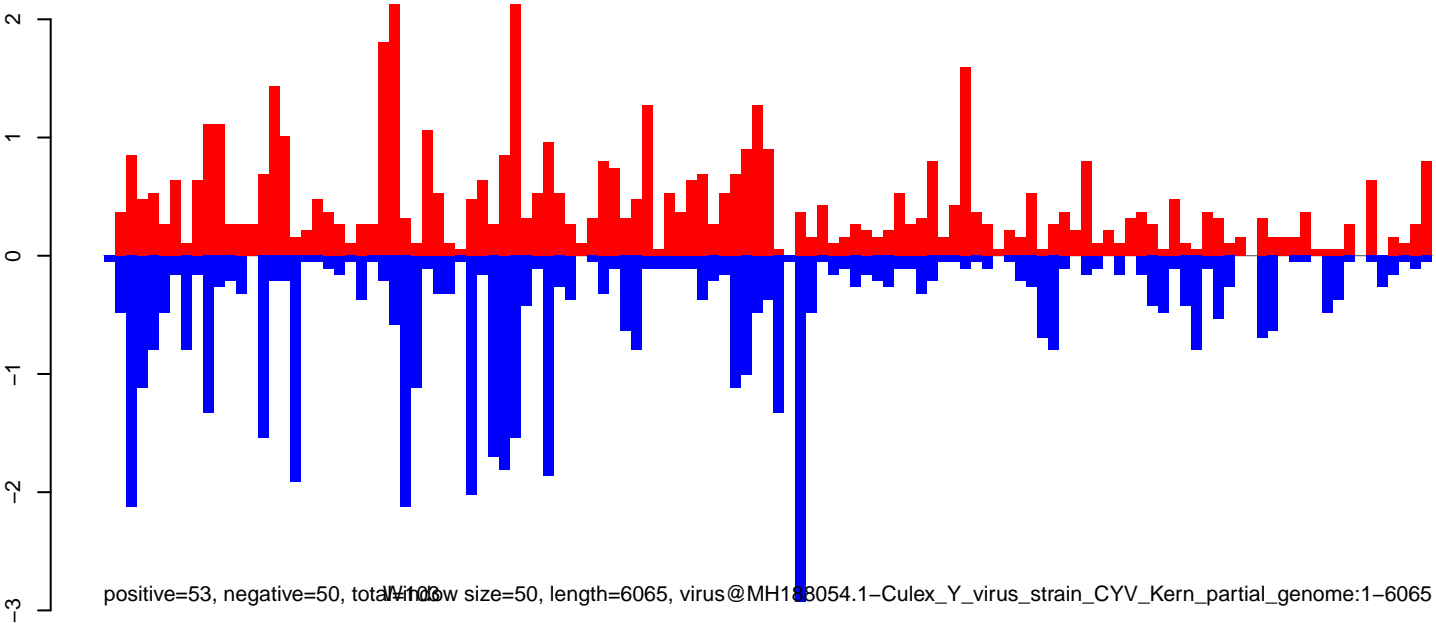
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

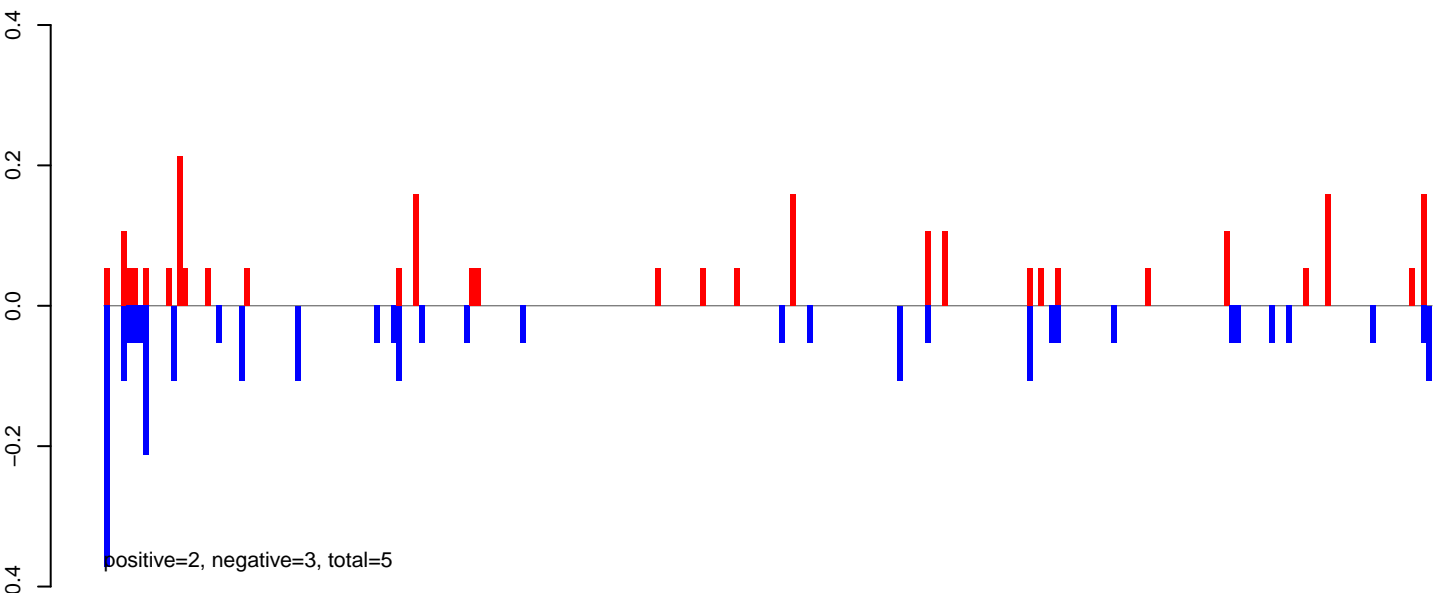


AeAeg_CCL.125cells_CHIKV.rep

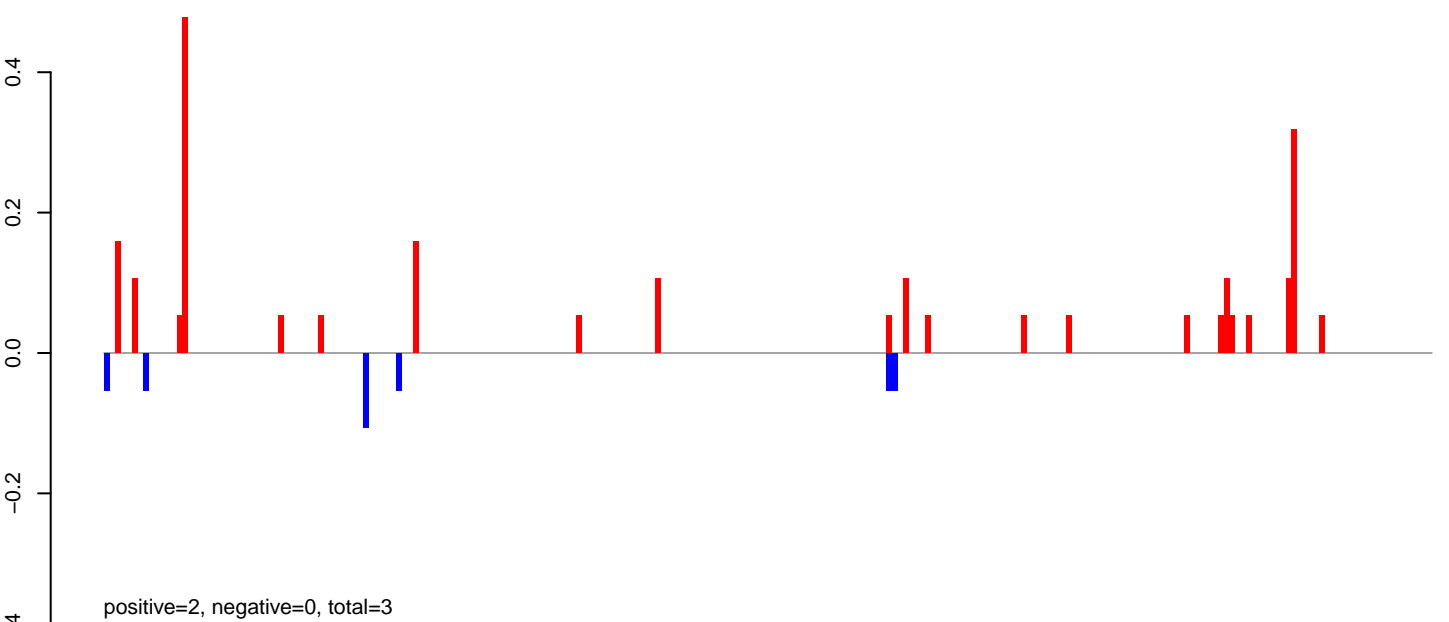


Window size=50, length=6065, virus@MH18054.1-Culex_Y_virus_strain_CYV_Kern_partial_genome:1-6065

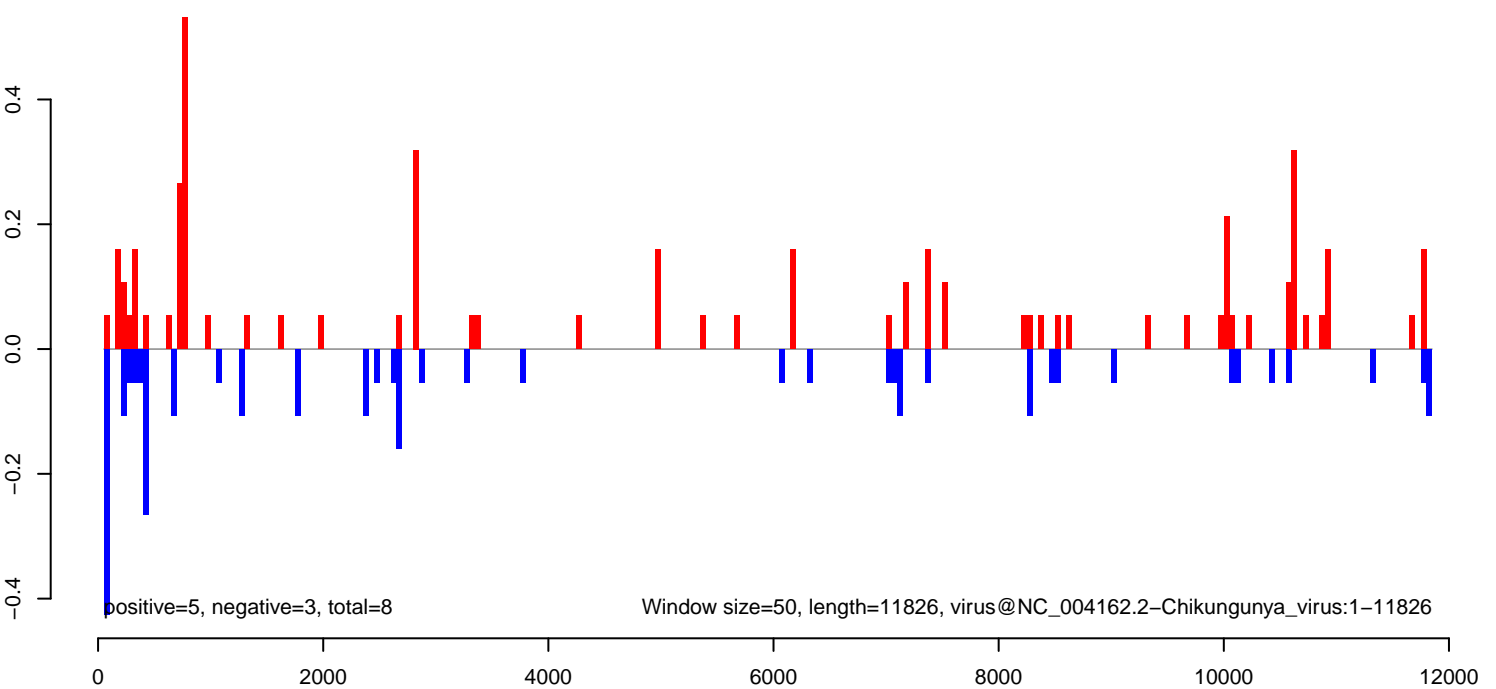
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep



AeAeg_CCL.125cells_CHIKV.18_23.rep



positive=4, negative=43, total=47

AeAeg_CCL.125cells_CHIKV.24_35.rep



positive=6, negative=341, total=346

AeAeg_CCL.125cells_CHIKV.rep

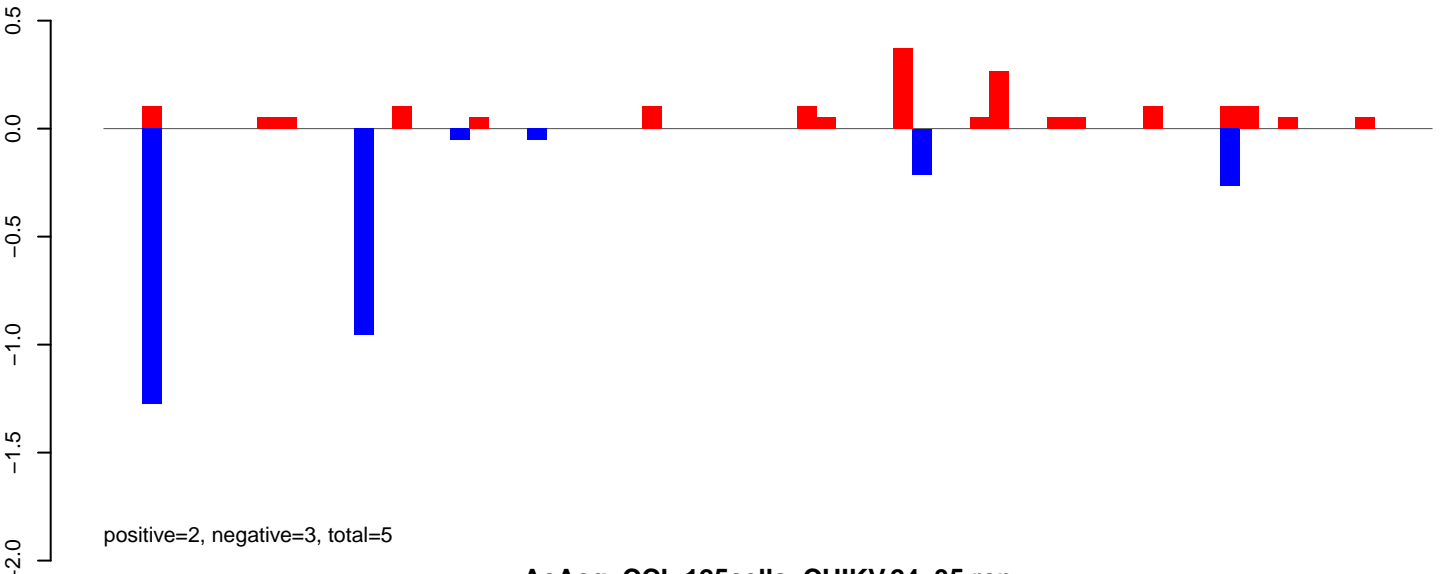


positive=10, negative=384, total=394

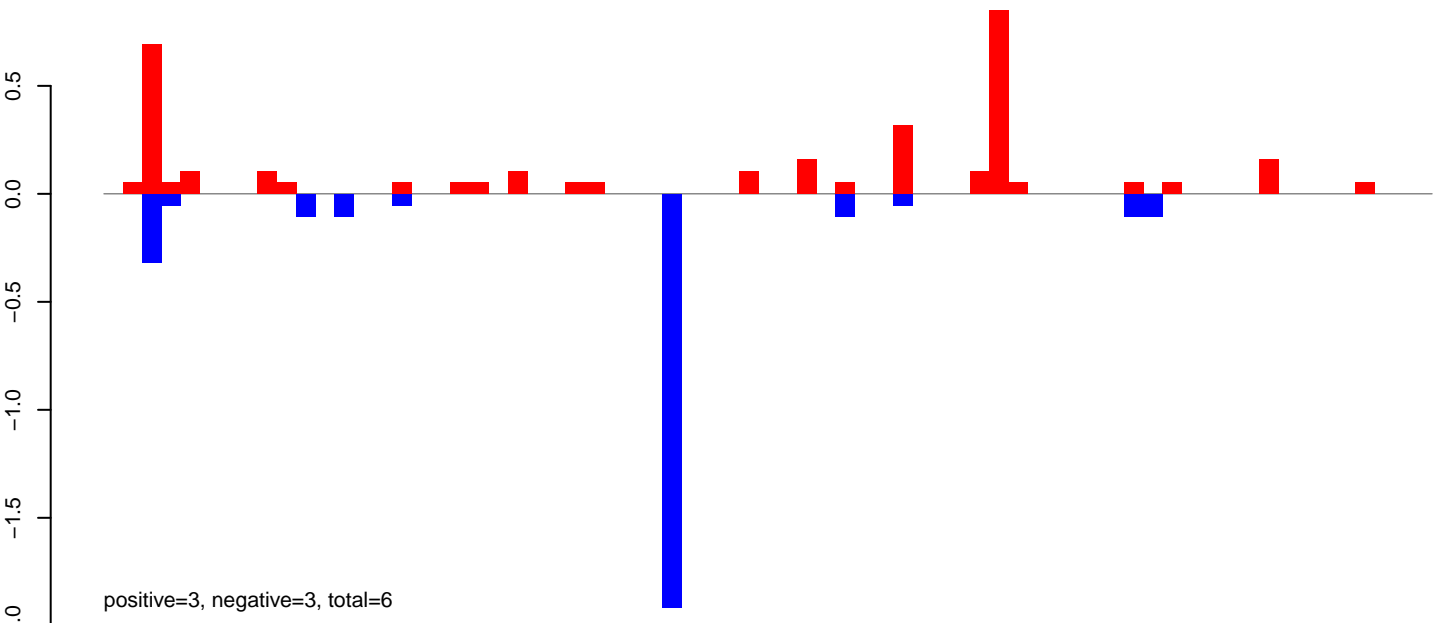
Window size=50, length=2965, virus@AY347953.2-Aedes_aegypti_genomic_sequence:1-2965

0 500 1000 1500 2000 2500 3000

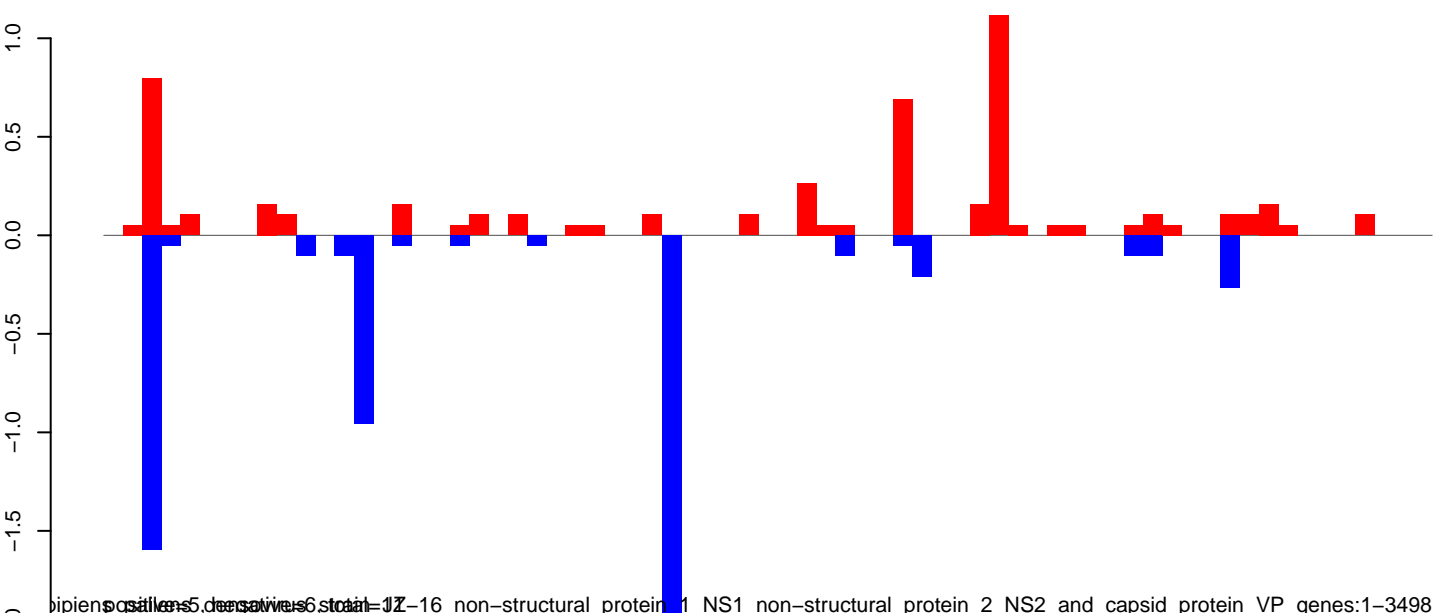
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

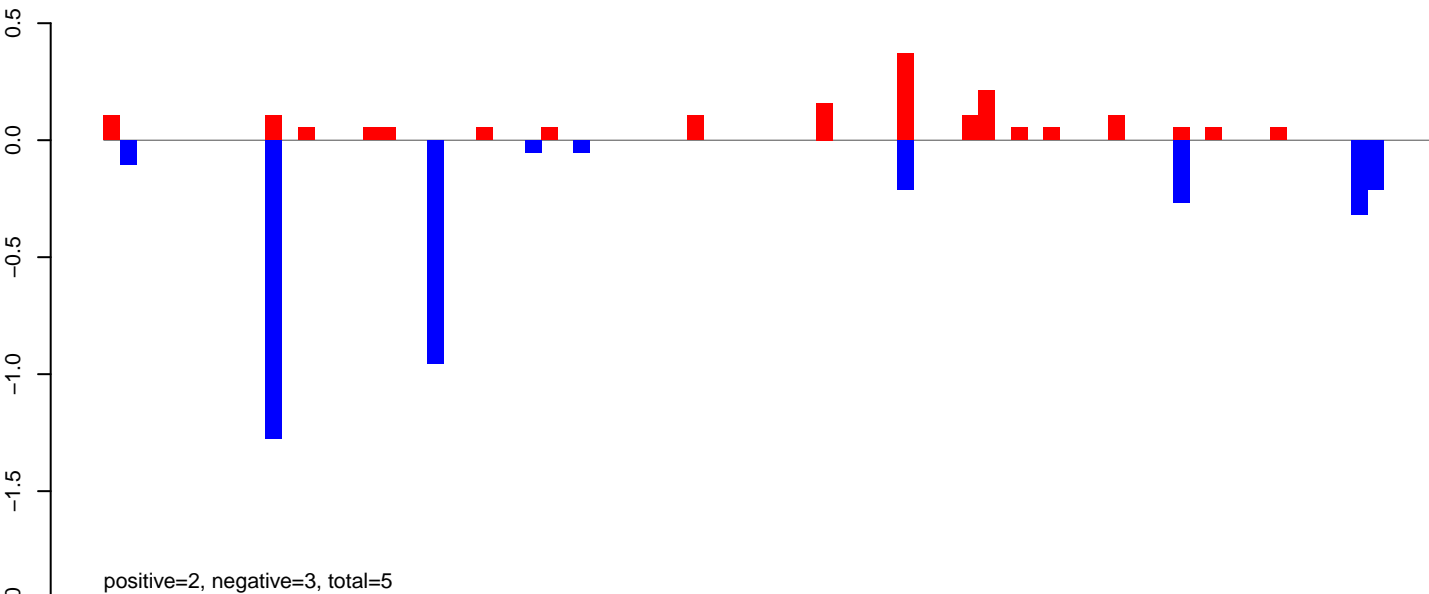


AeAeg_CCL.125cells_CHIKV.rep

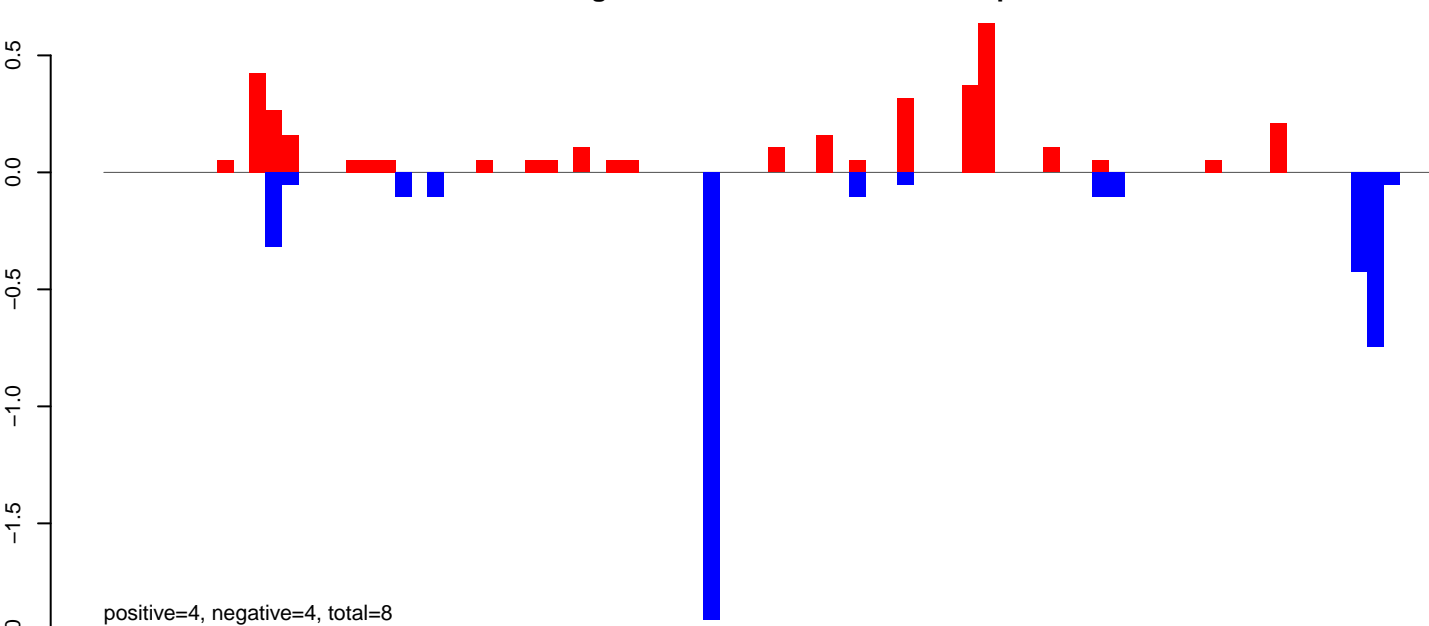


ipienposilens5, negatoves6, total=11-16_non-structural_protein_1_NS1_non-structural_protein_2_NS2_and_capsid_protein_VP_genes:1-3498

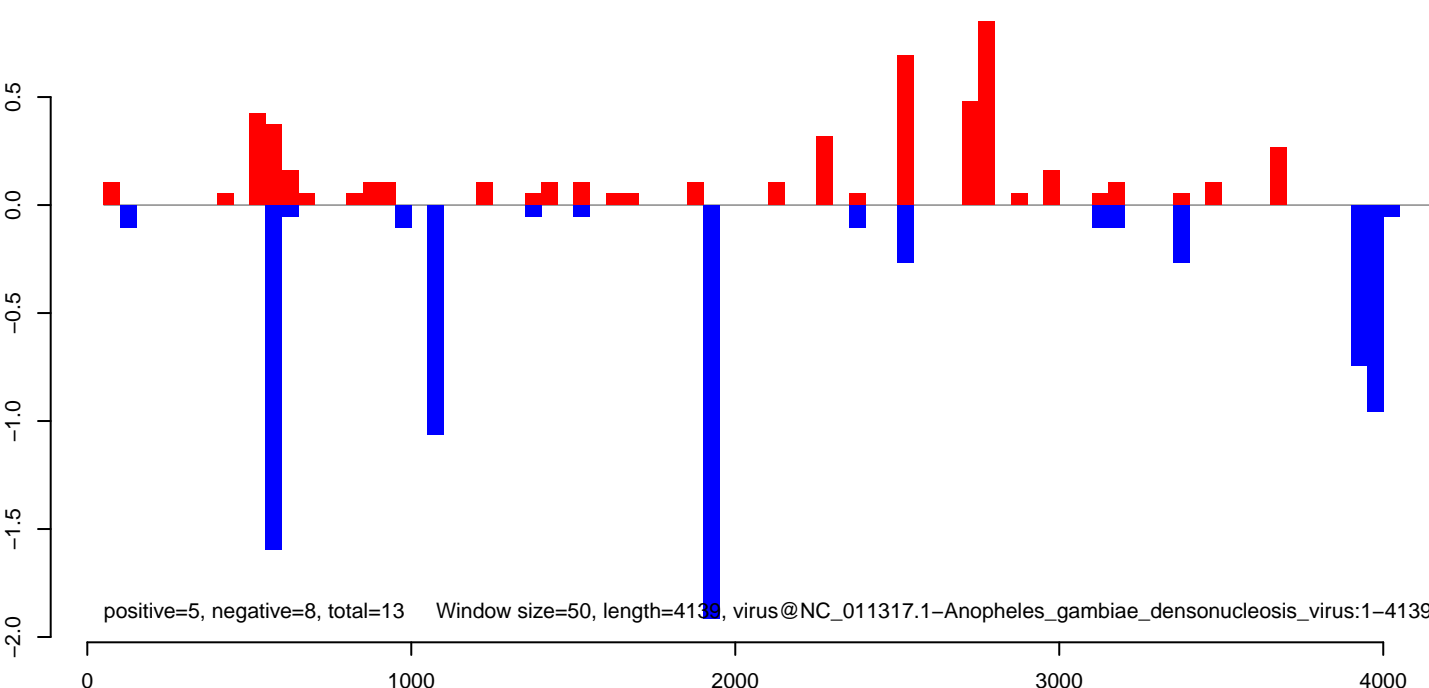
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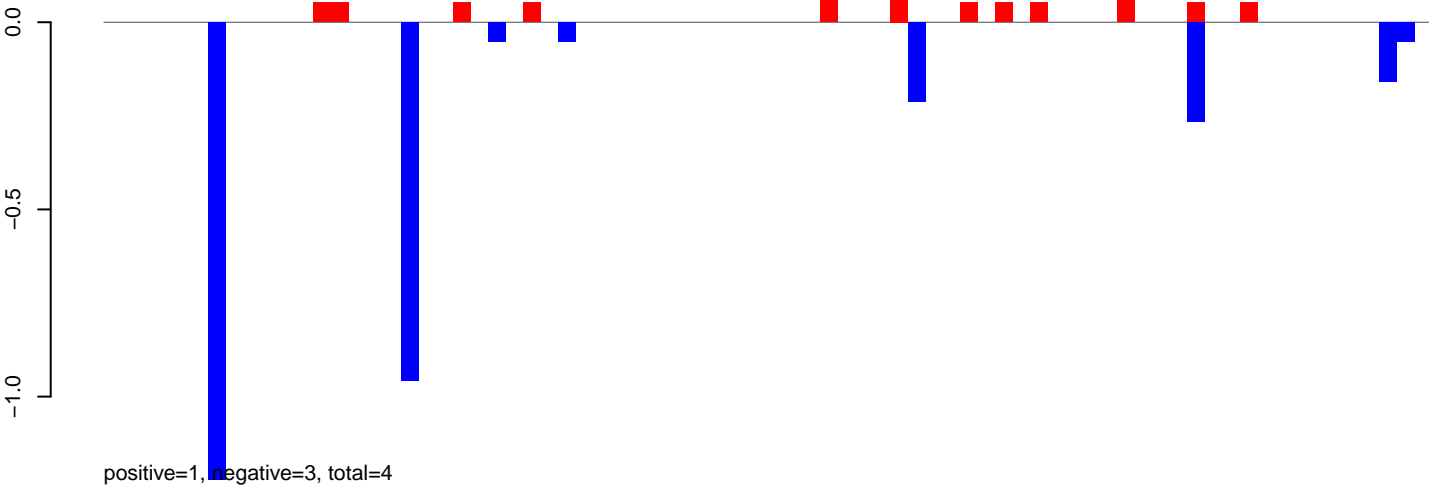
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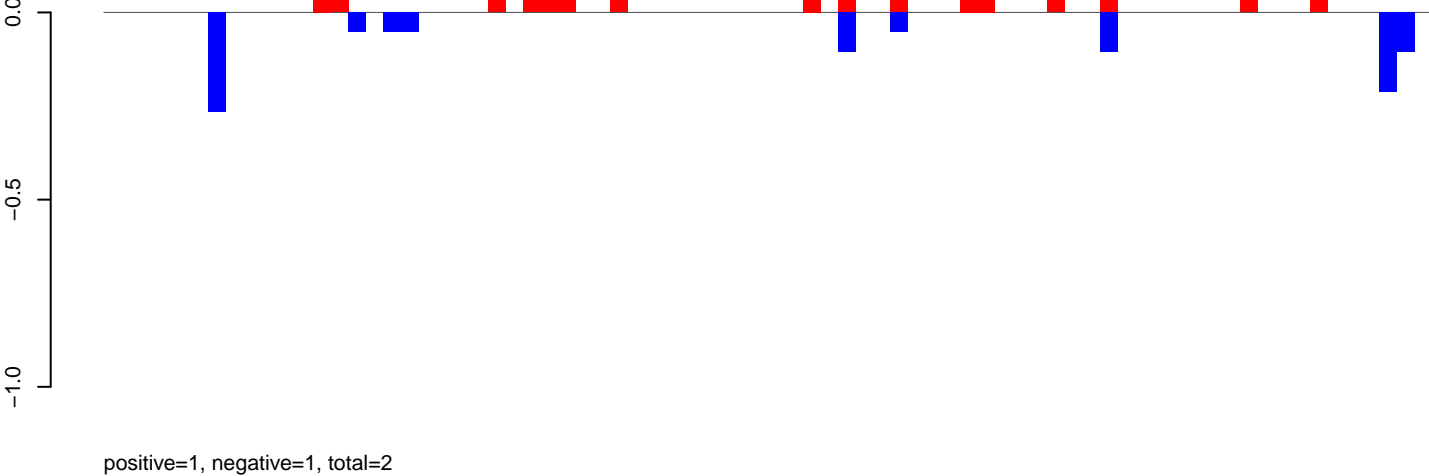
AeAeg_CCL.125cells_CHIKV.rep



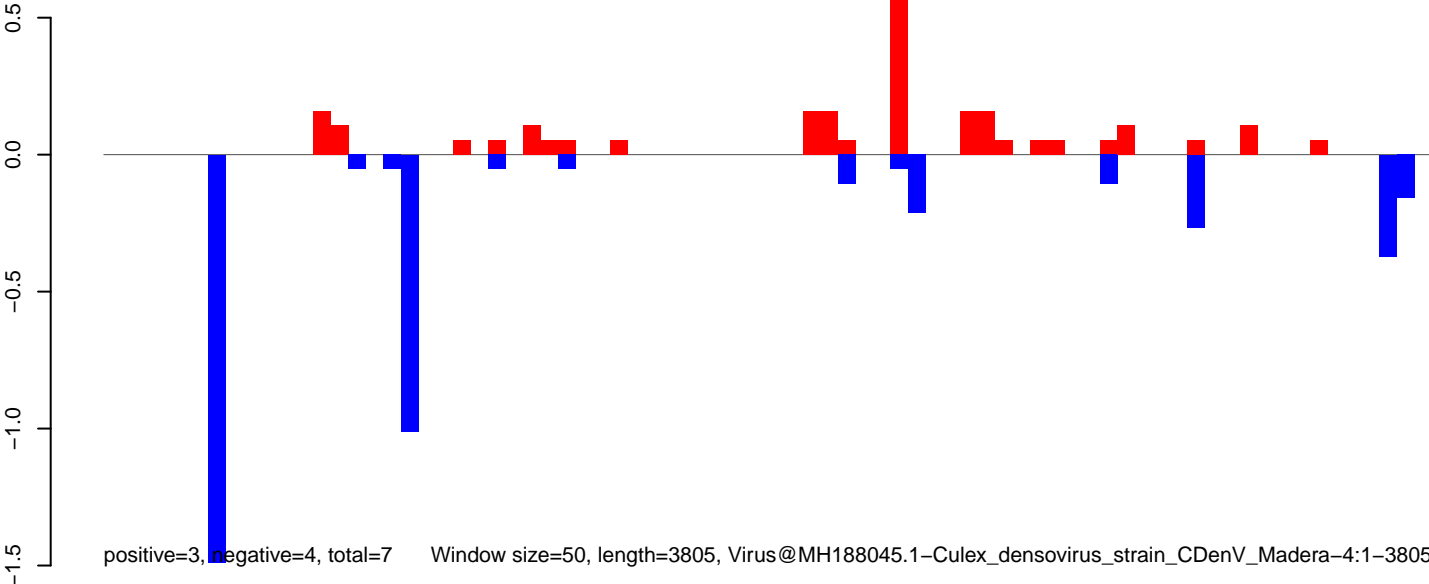
AeAeg_CCL.125cells_CHIKV.18_23.rep



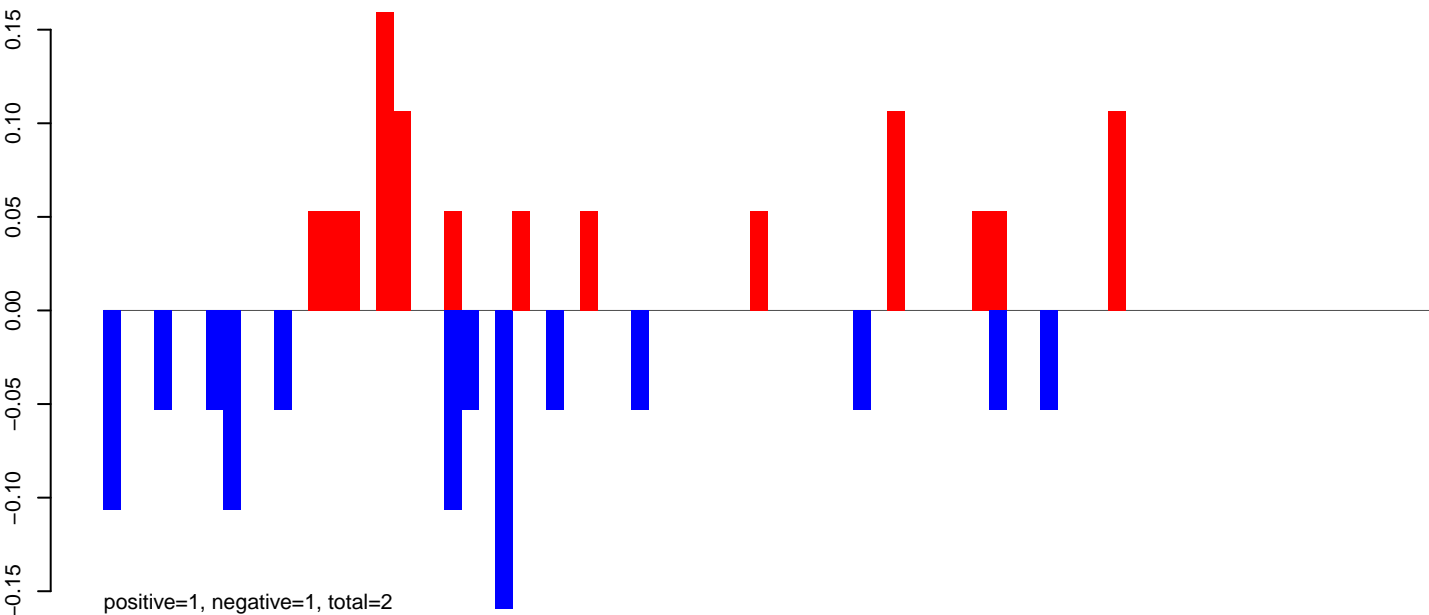
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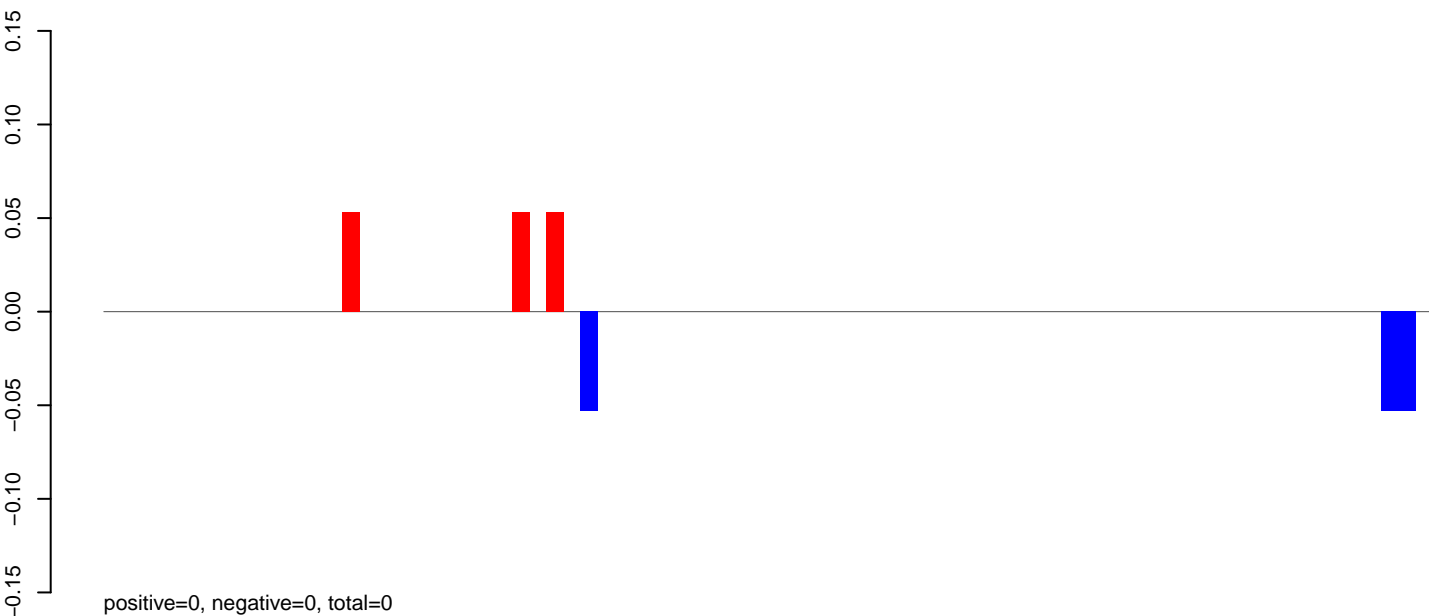
AeAeg_CCL.125cells_CHIKV.rep



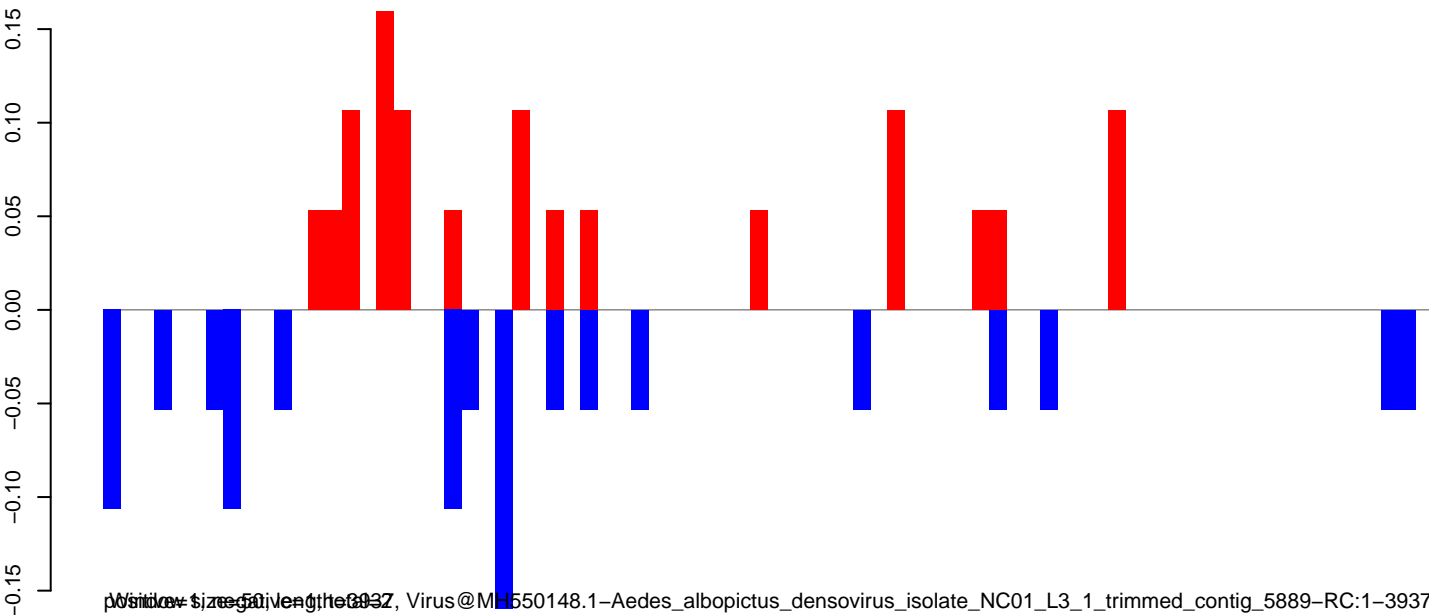
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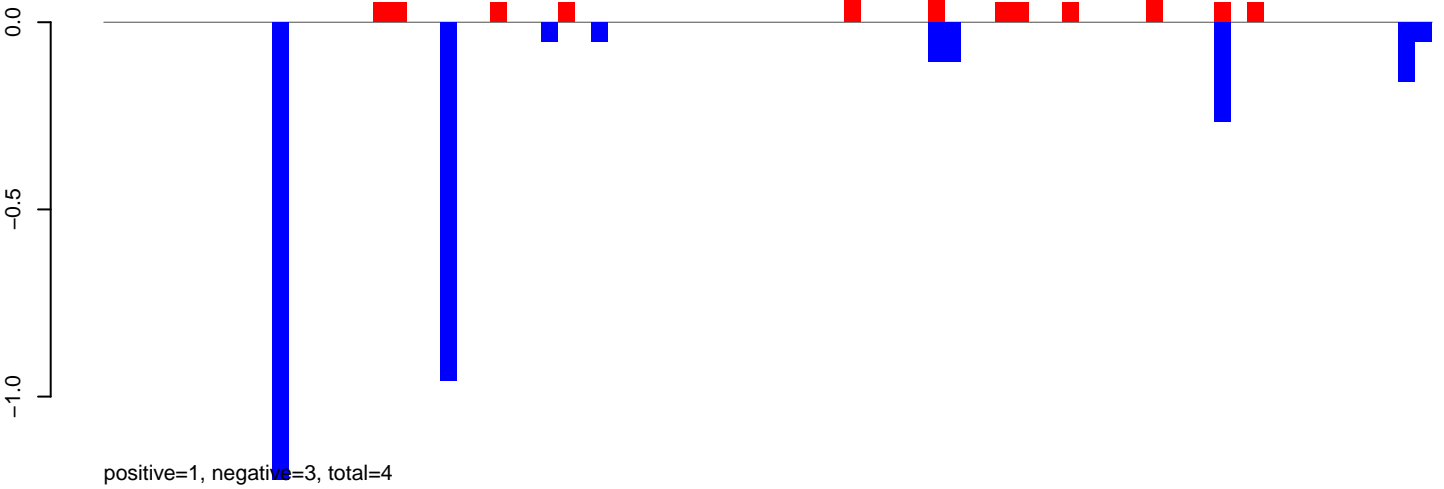
AeAeg_CCL.125cells_CHIKV.24_35.rep



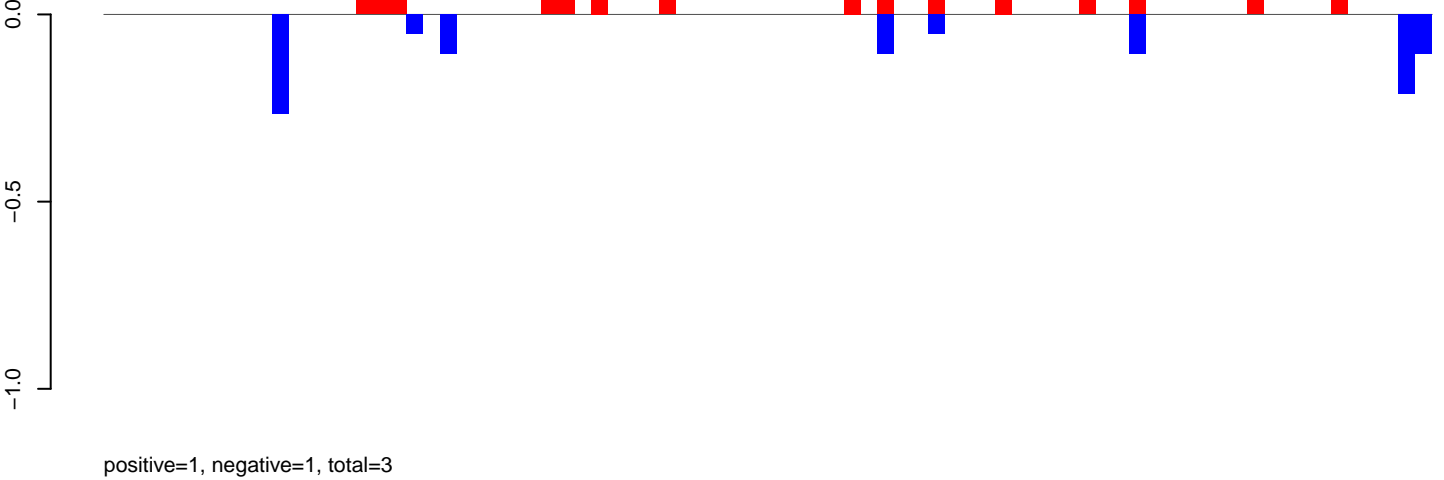
AeAeg_CCL.125cells_CHIKV.rep



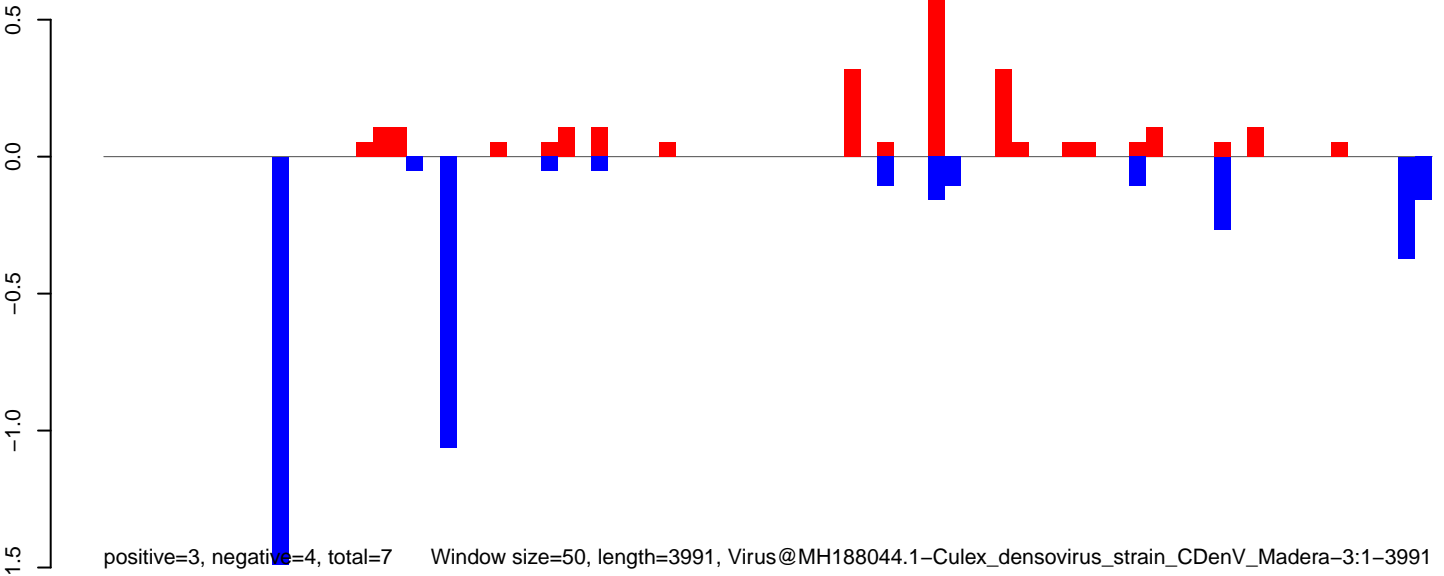
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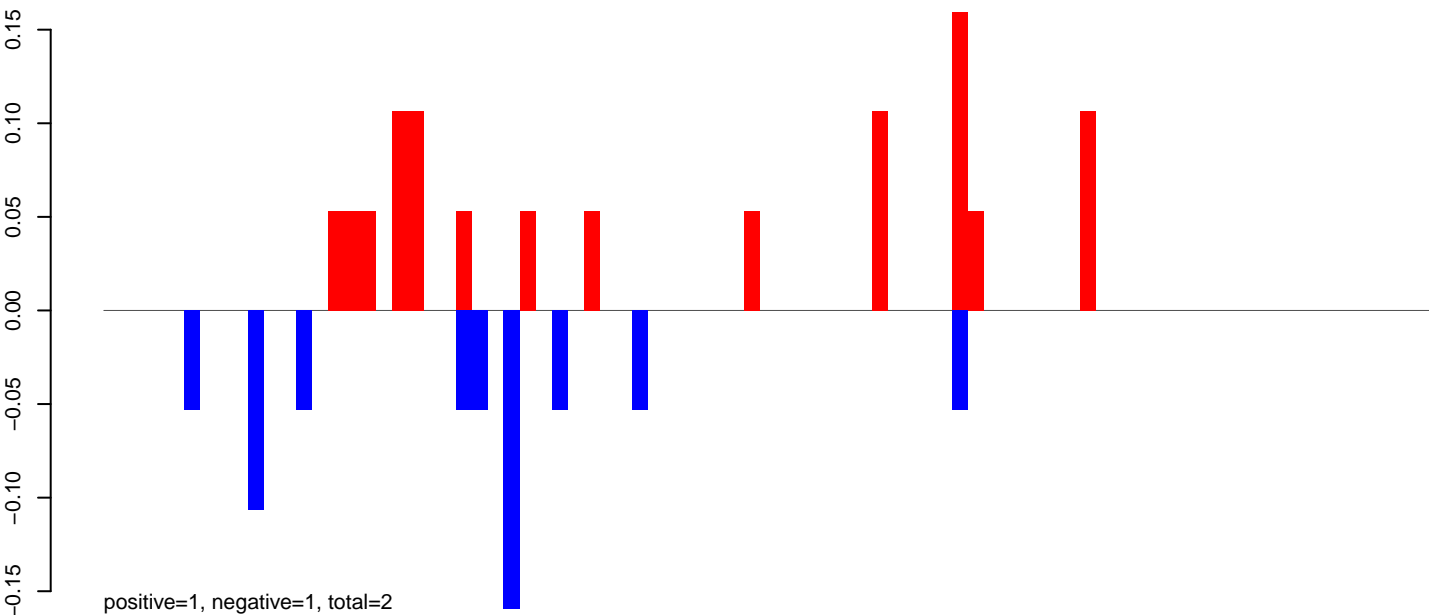
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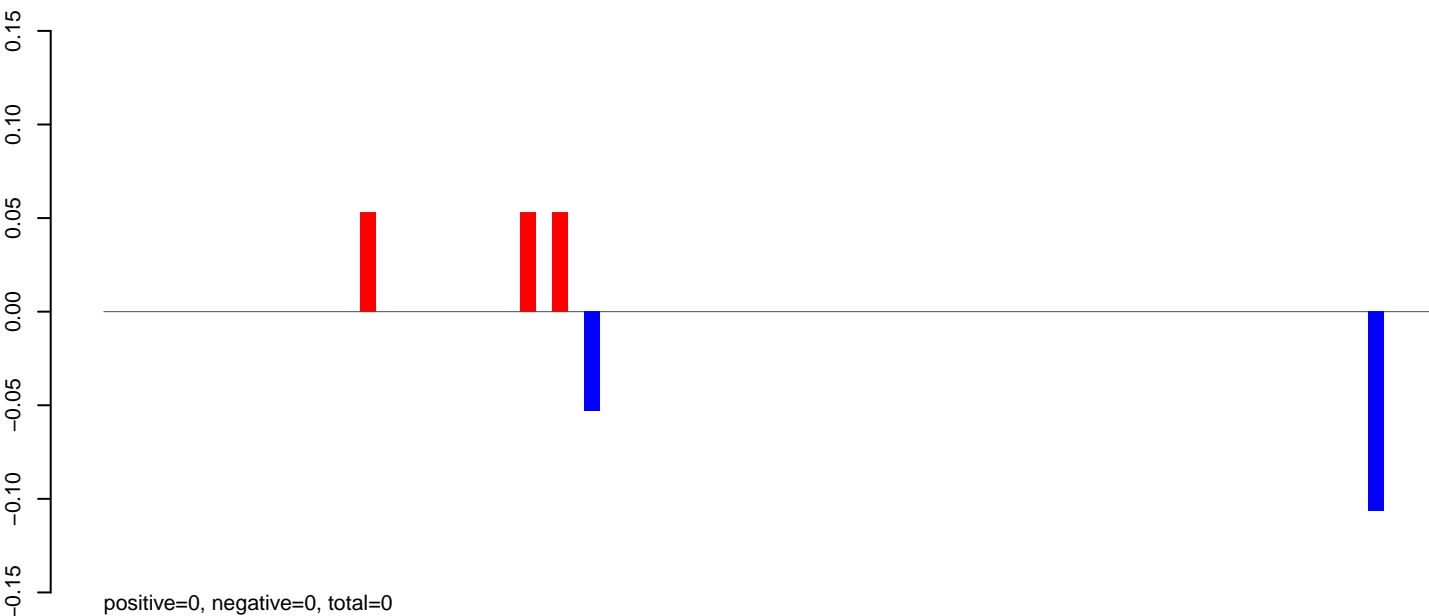
AeAeg_CCL.125cells_CHIKV.rep



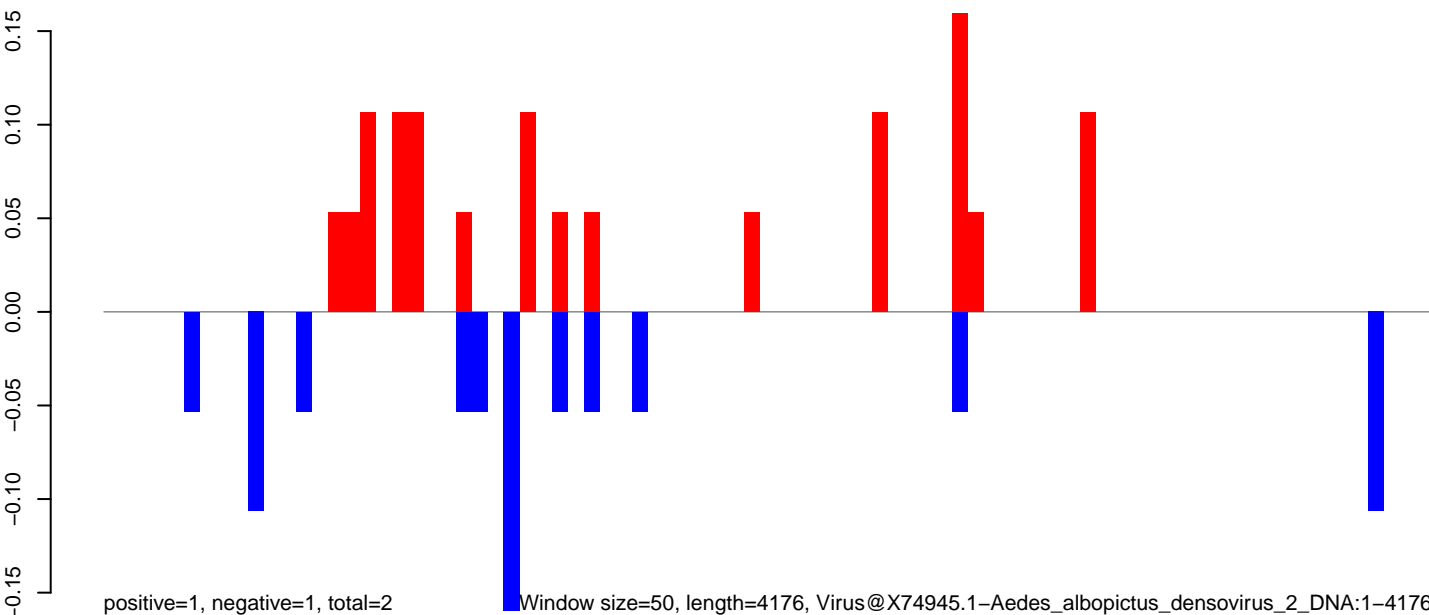
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

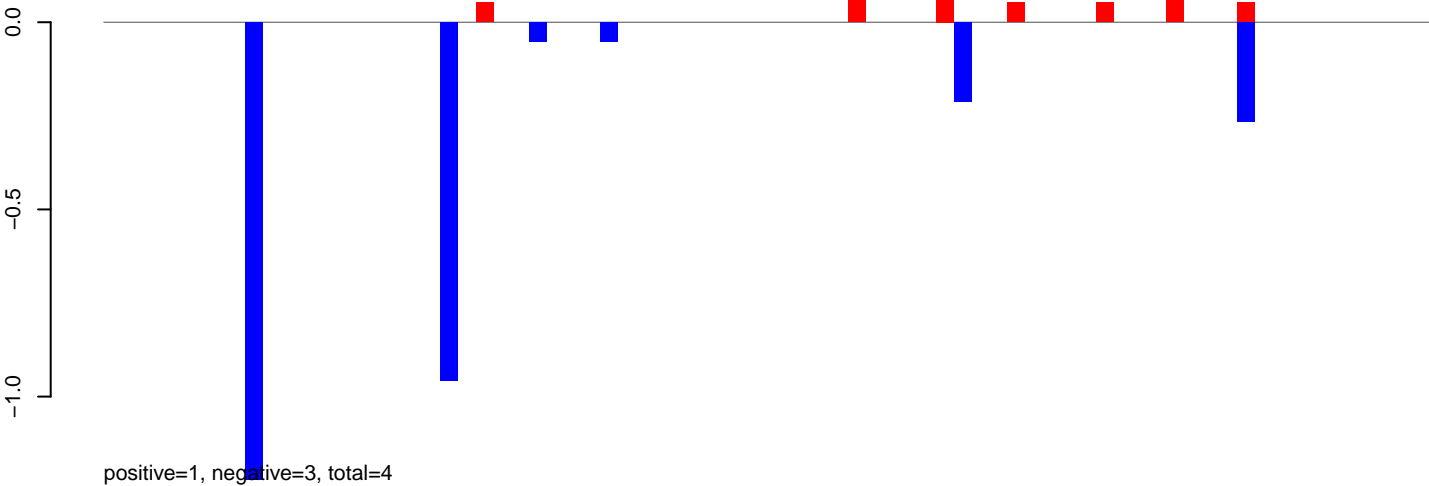


AeAeg_CCL.125cells_CHIKV.rep

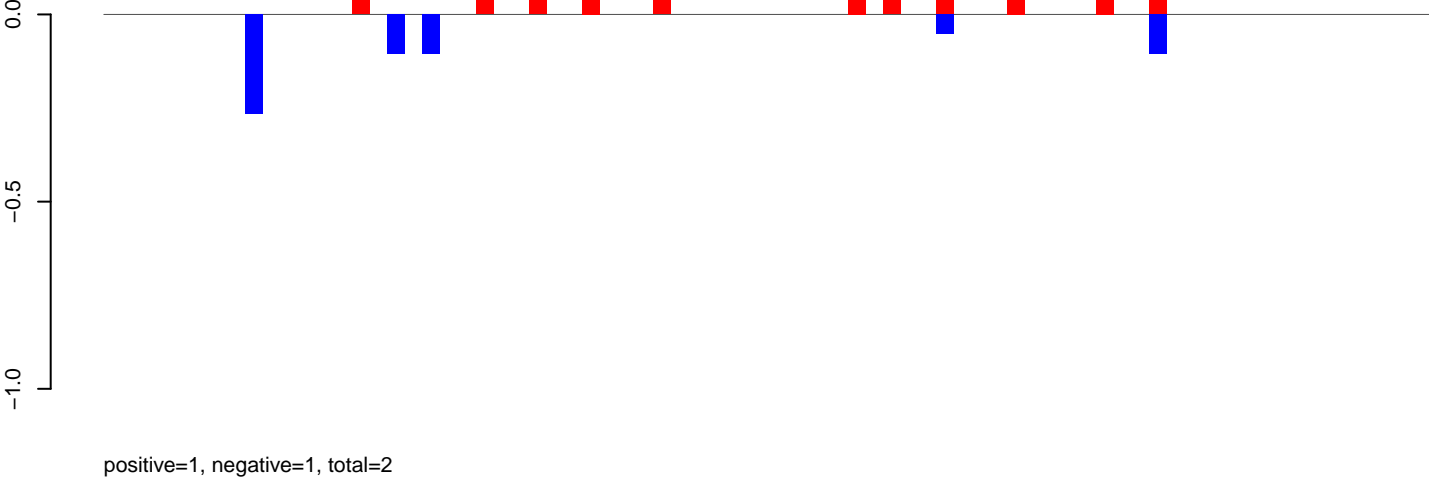


Window size=50, length=4176, Virus@X74945.1-Aedes_albopictus_densovirus_2_DNA:1-4176

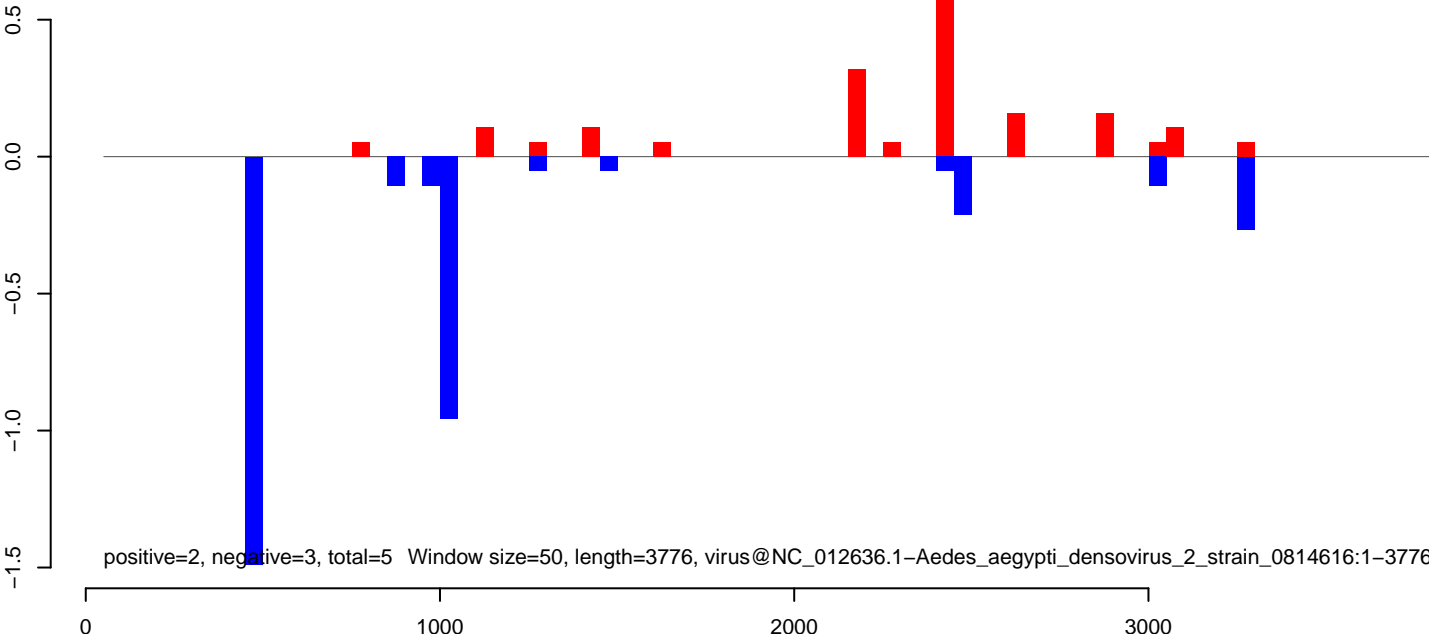
AeAeg_CCL.125cells_CHIKV.18_23.rep



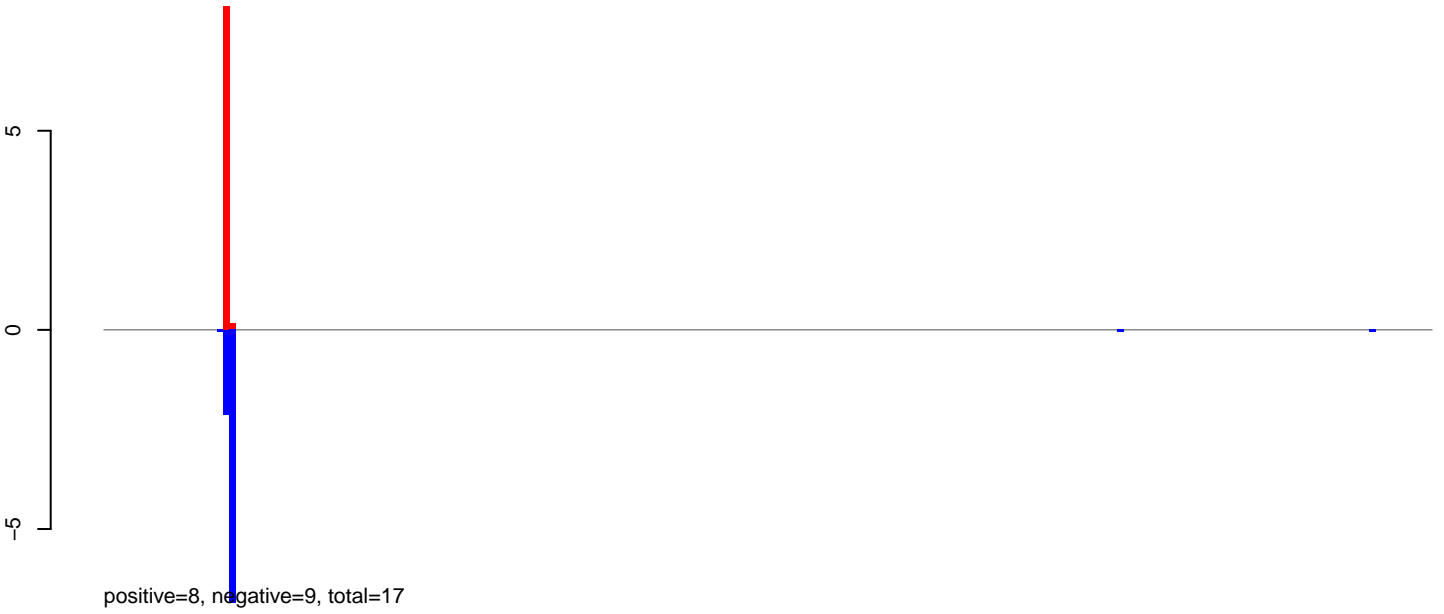
AeAeg_CCL.125cells_CHIKV.24_35.rep



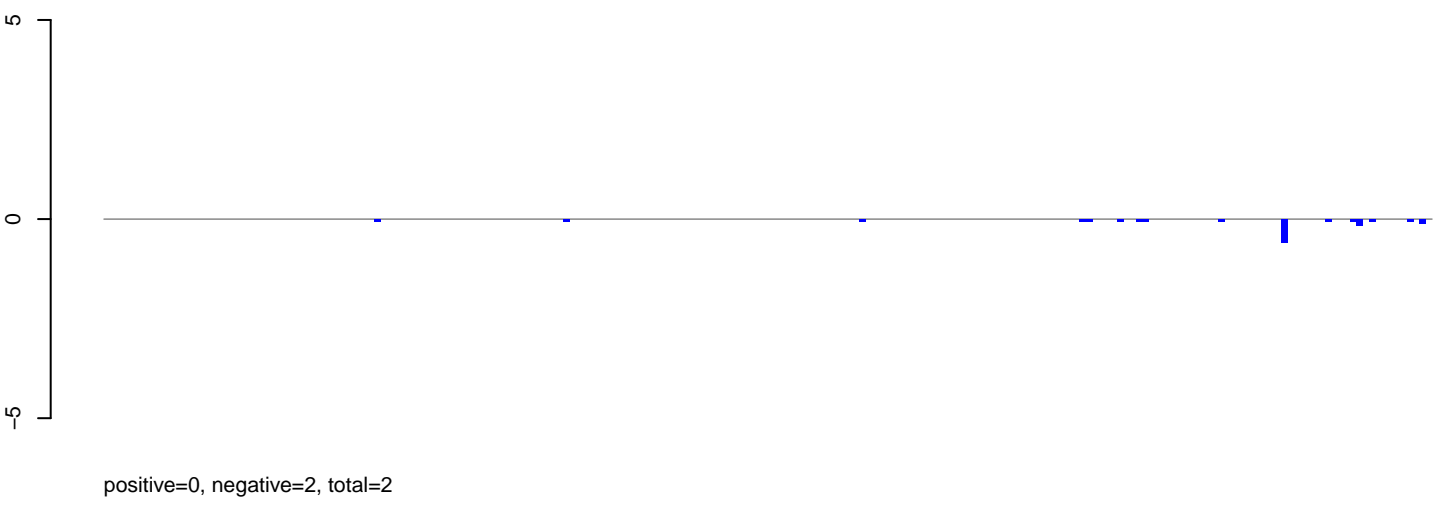
AeAeg_CCL.125cells_CHIKV.rep



AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



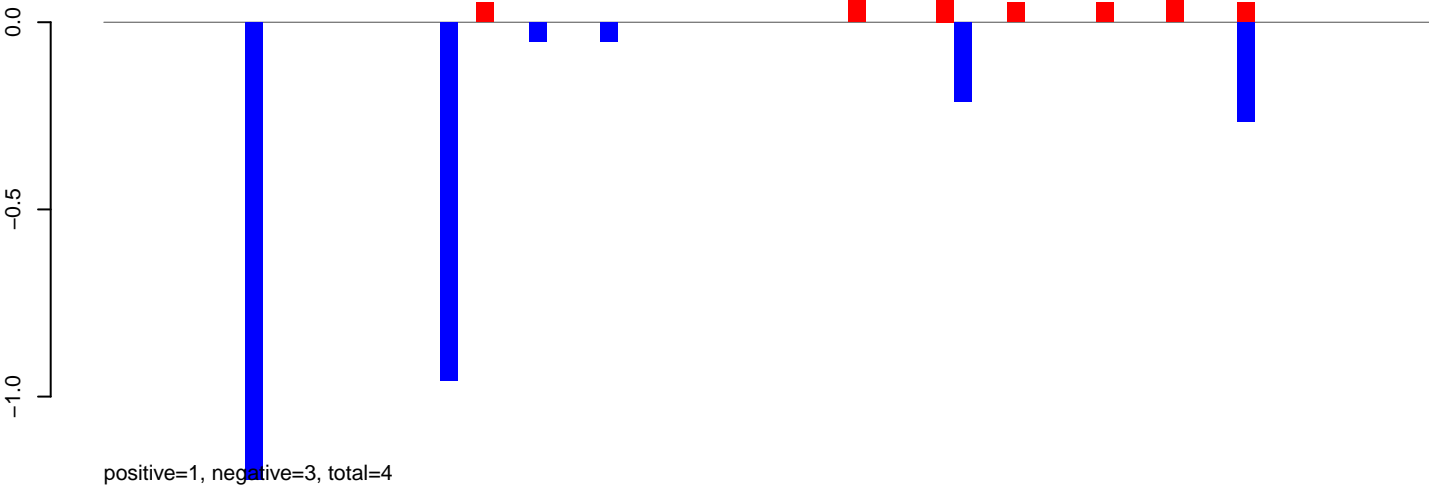
AeAeg_CCL.125cells_CHIKV.rep



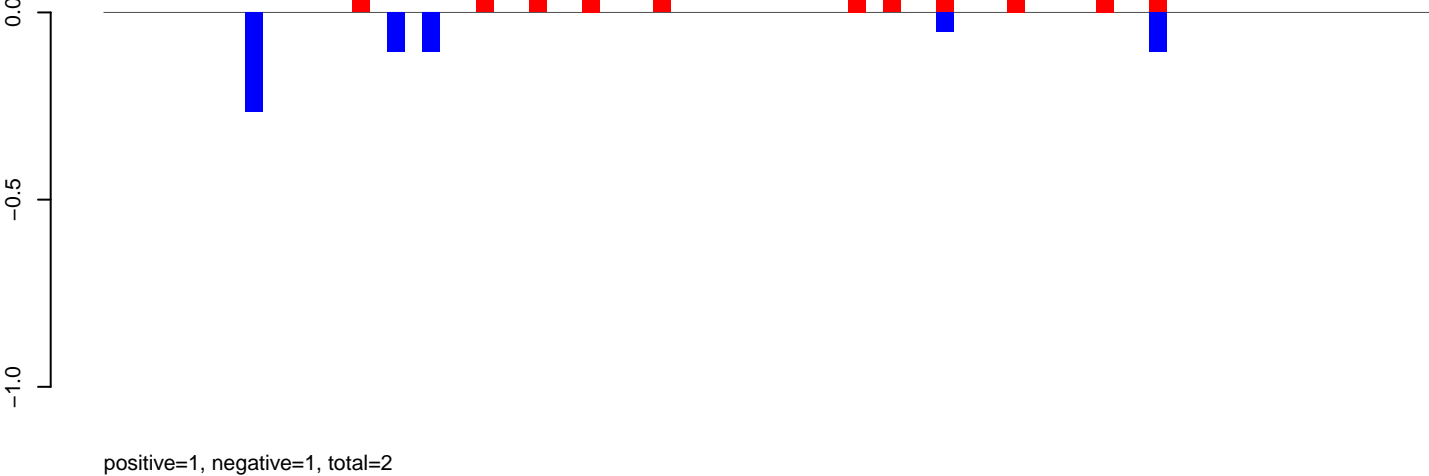
AF416352-8 deduced from AF416352-9 using agent_virus_polyprotein-like_protein_gene_and_agCP8252-like_protein_gene_partial_cds:1-10587

0 2000 4000 6000 8000 10000

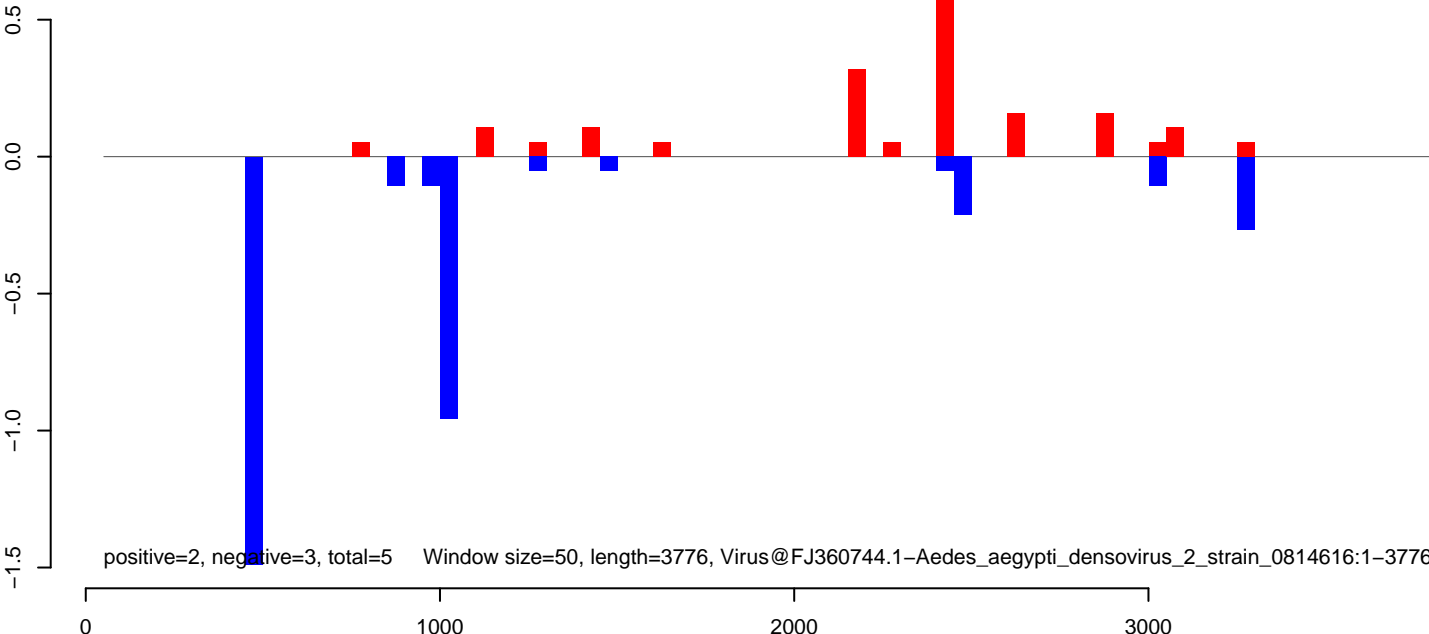
AeAeg_CCL.125cells_CHIKV.18_23.rep



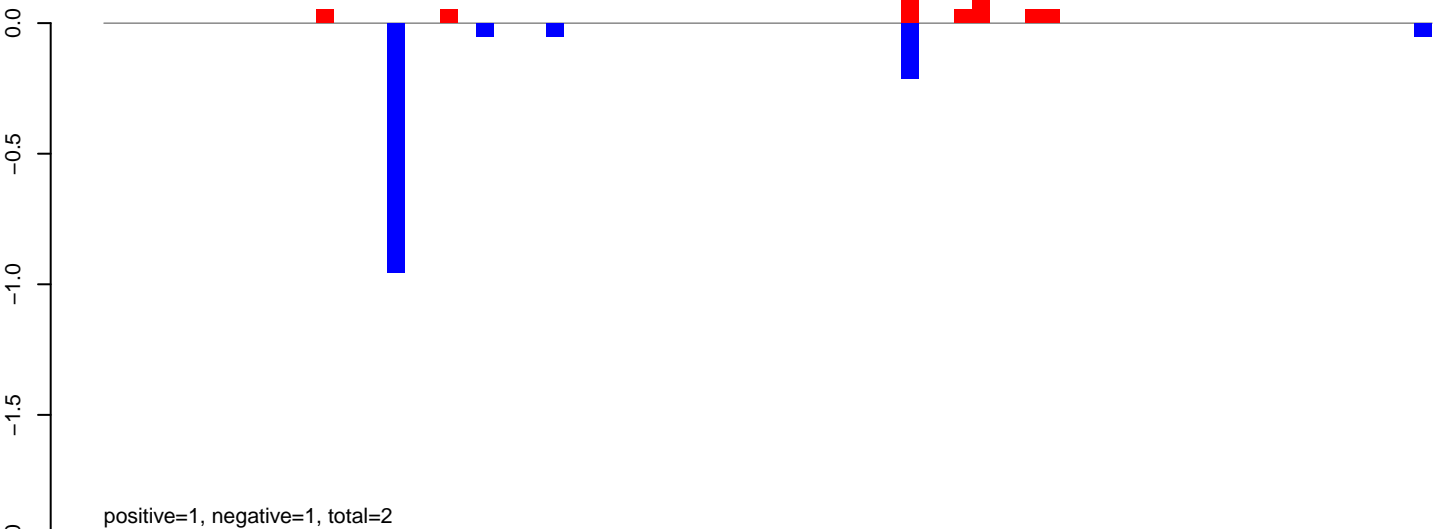
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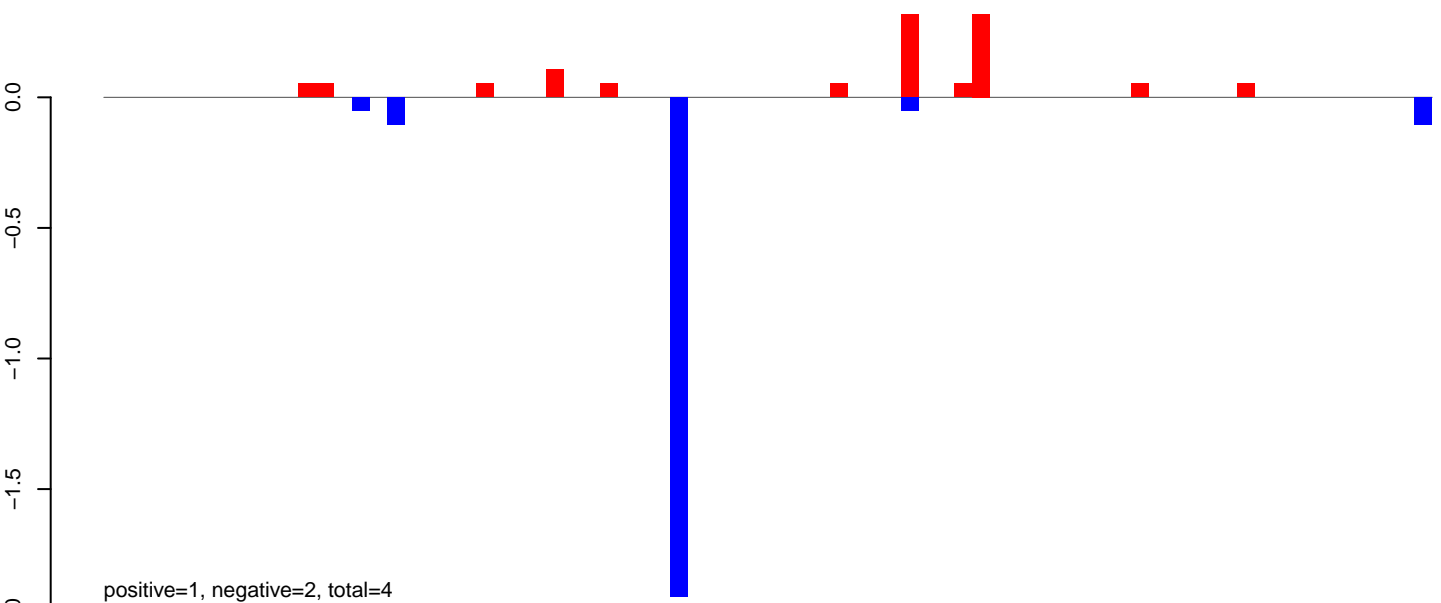
AeAeg_CCL.125cells_CHIKV.rep



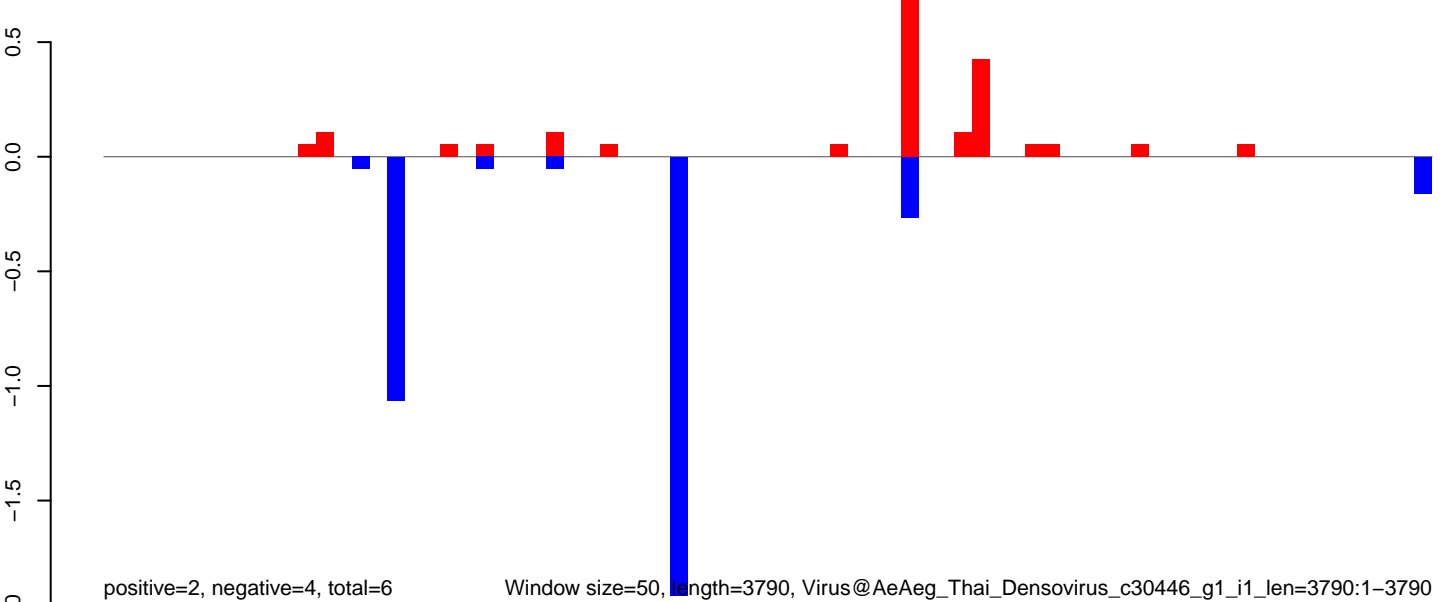
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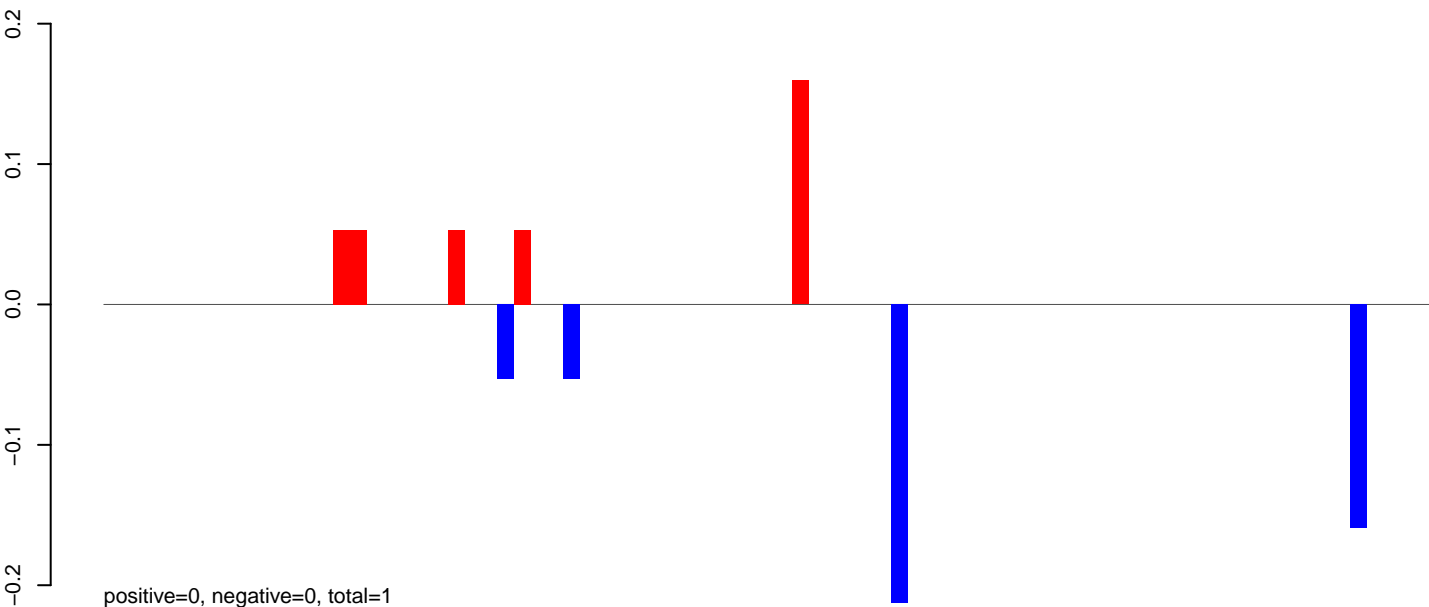
AeAeg_CCL.125cells_CHIKV.24_35.rep



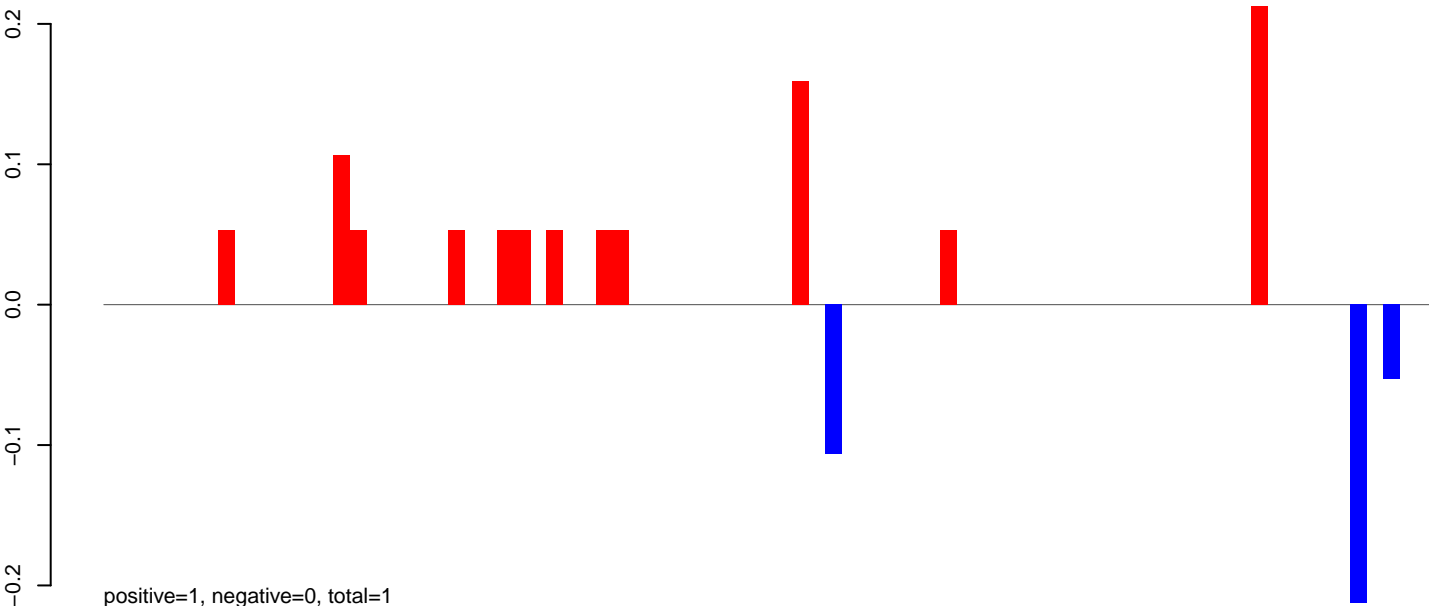
AeAeg_CCL.125cells_CHIKV.rep



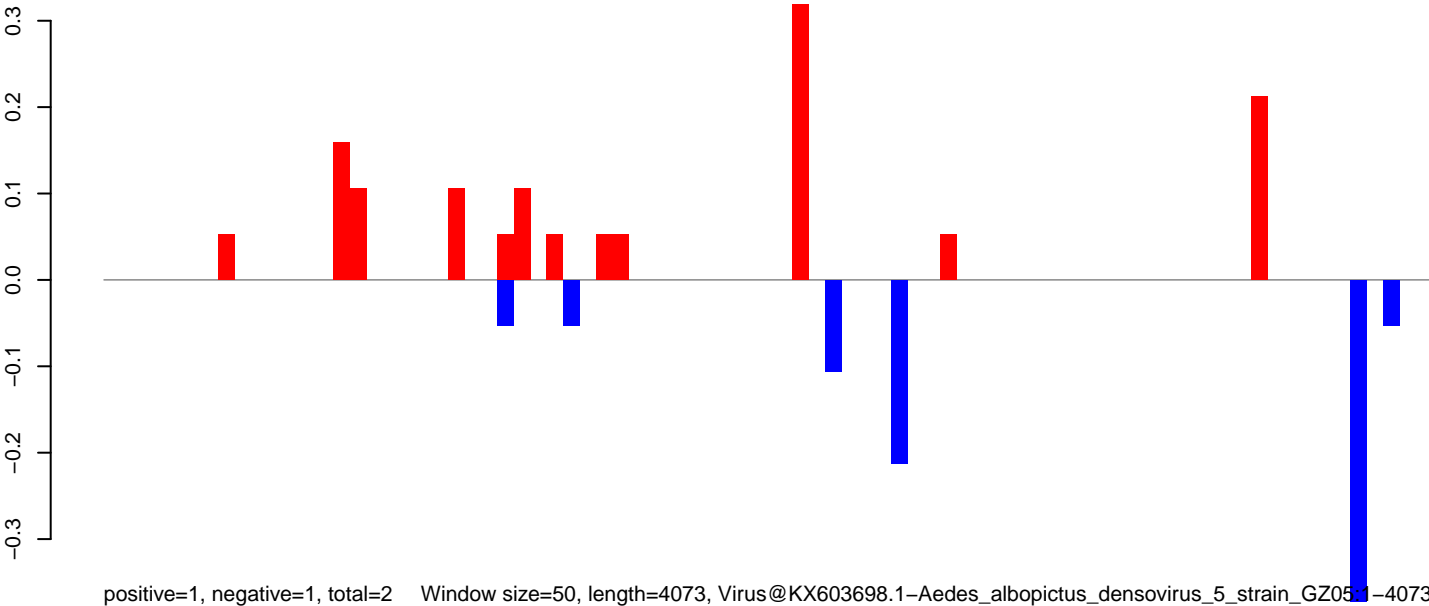
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

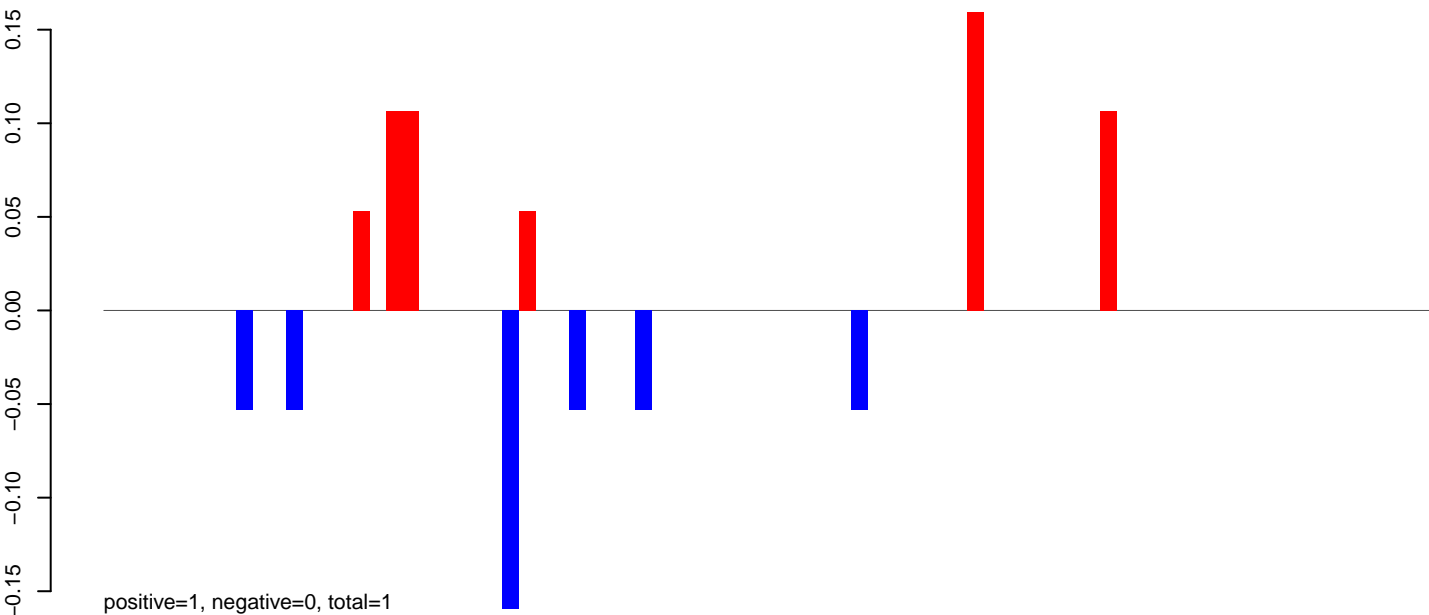


AeAeg_CCL.125cells_CHIKV.rep

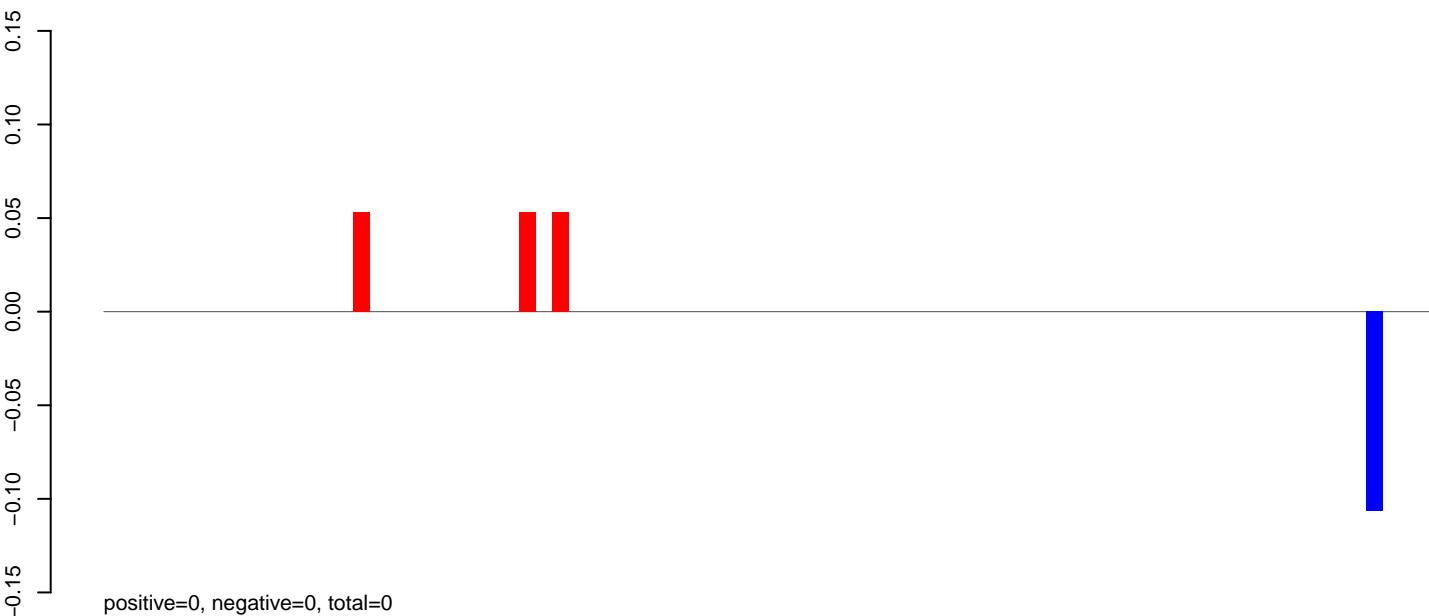


Window size=50, length=4073, Virus@KX603698.1-Aedes_albopictus_densovirus_5_strain_GZ051-4073

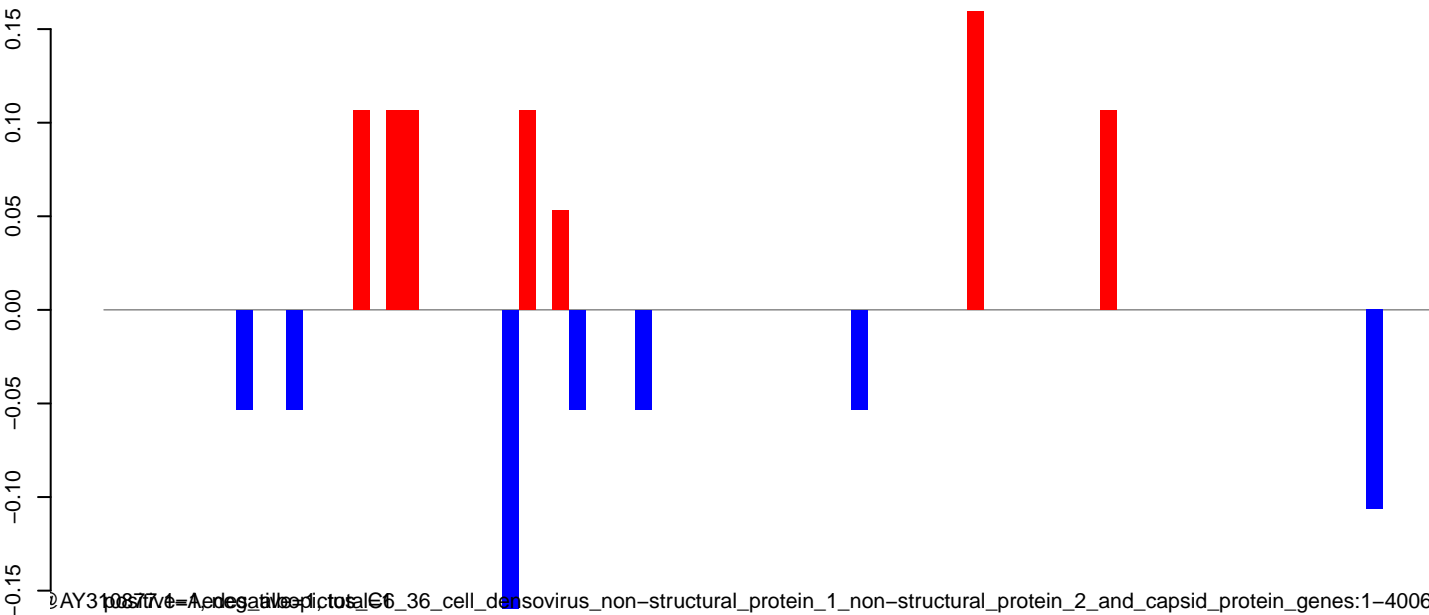
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

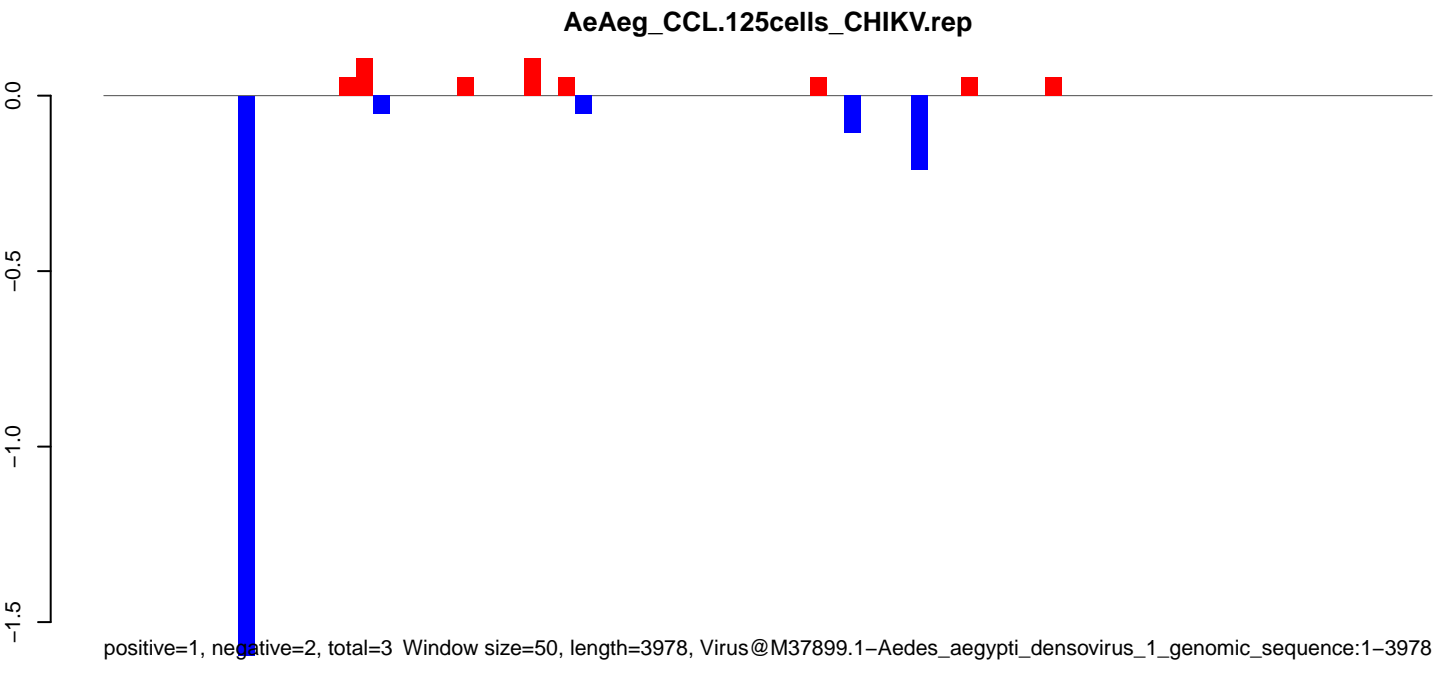
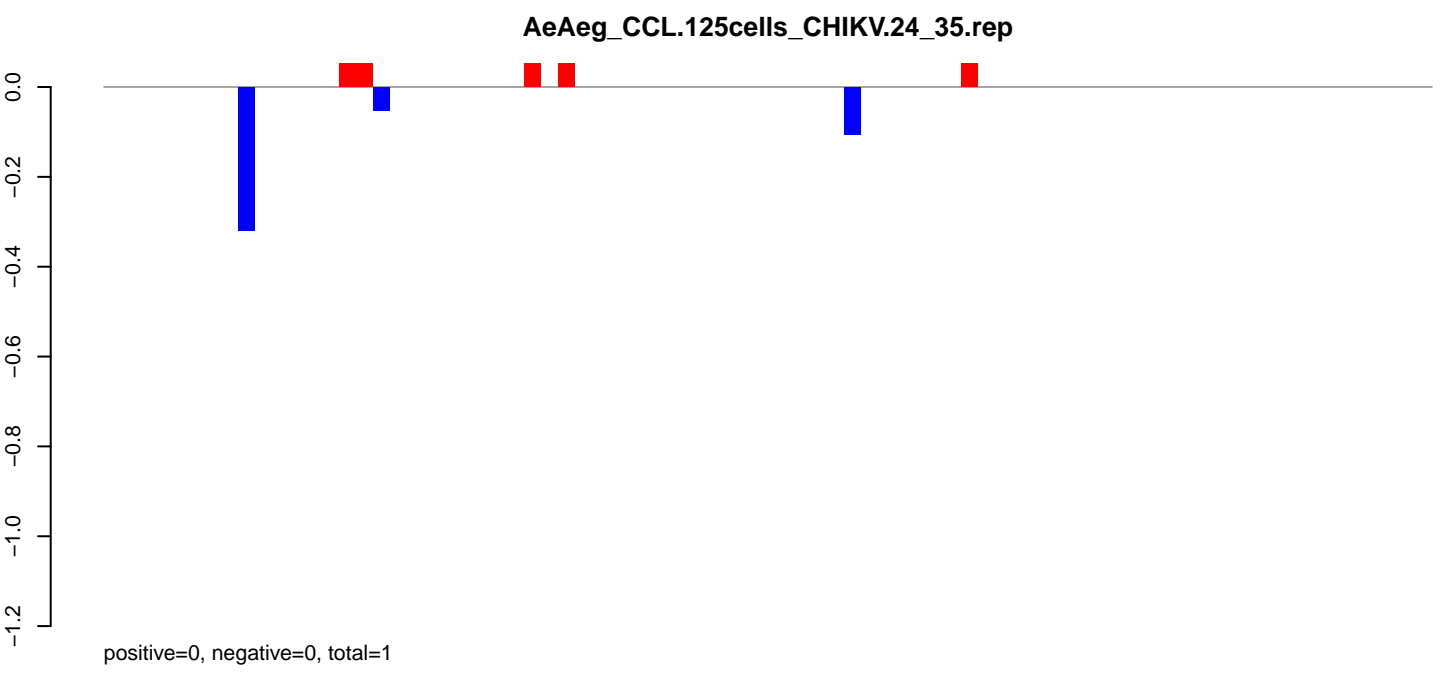
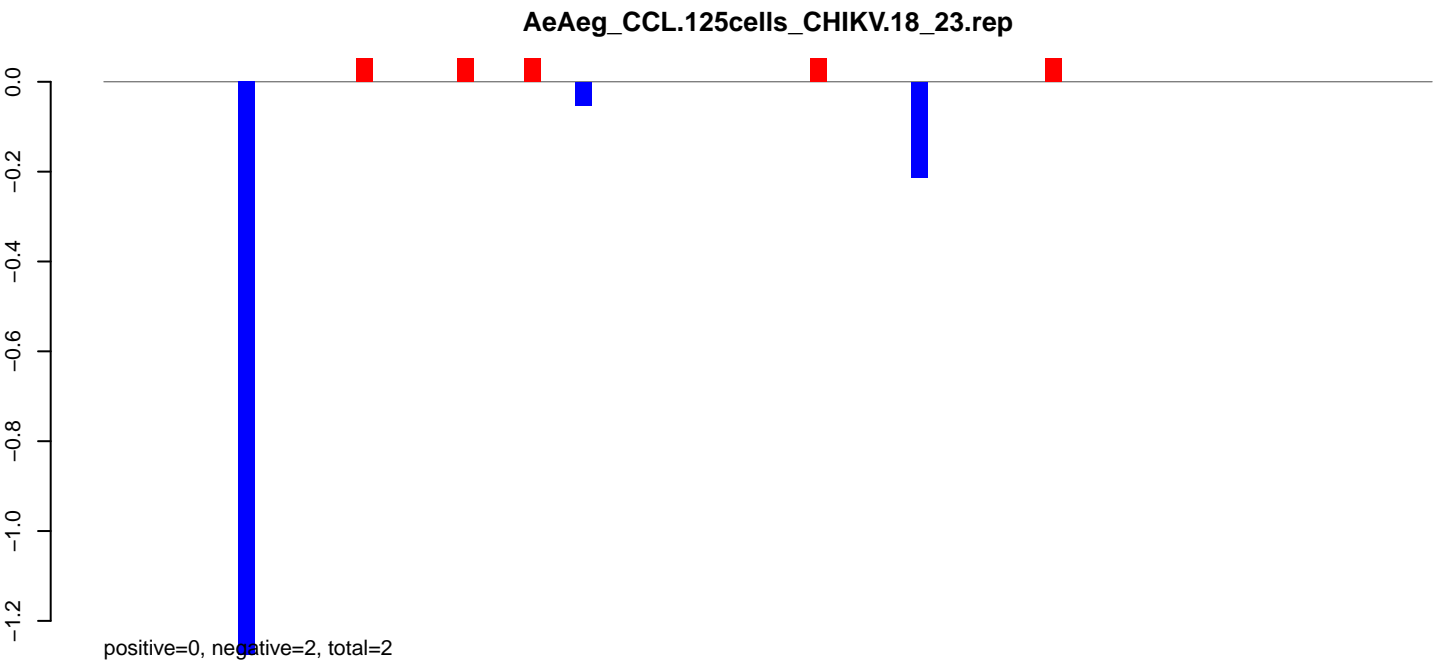


AeAeg_CCL.125cells_CHIKV.rep

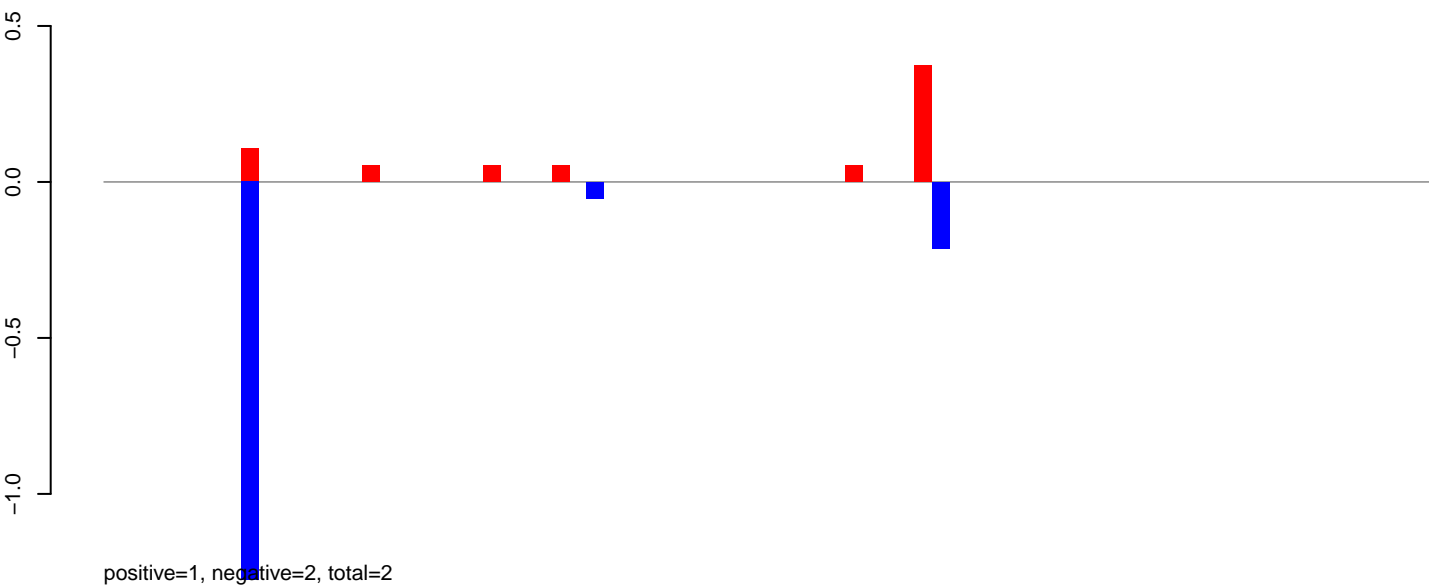


AY310874-Aedes albopictus C6_36_cell_densovirus_non-structural_protein_1_non-structural_protein_2_and_capsid_protein_genes:1-4006

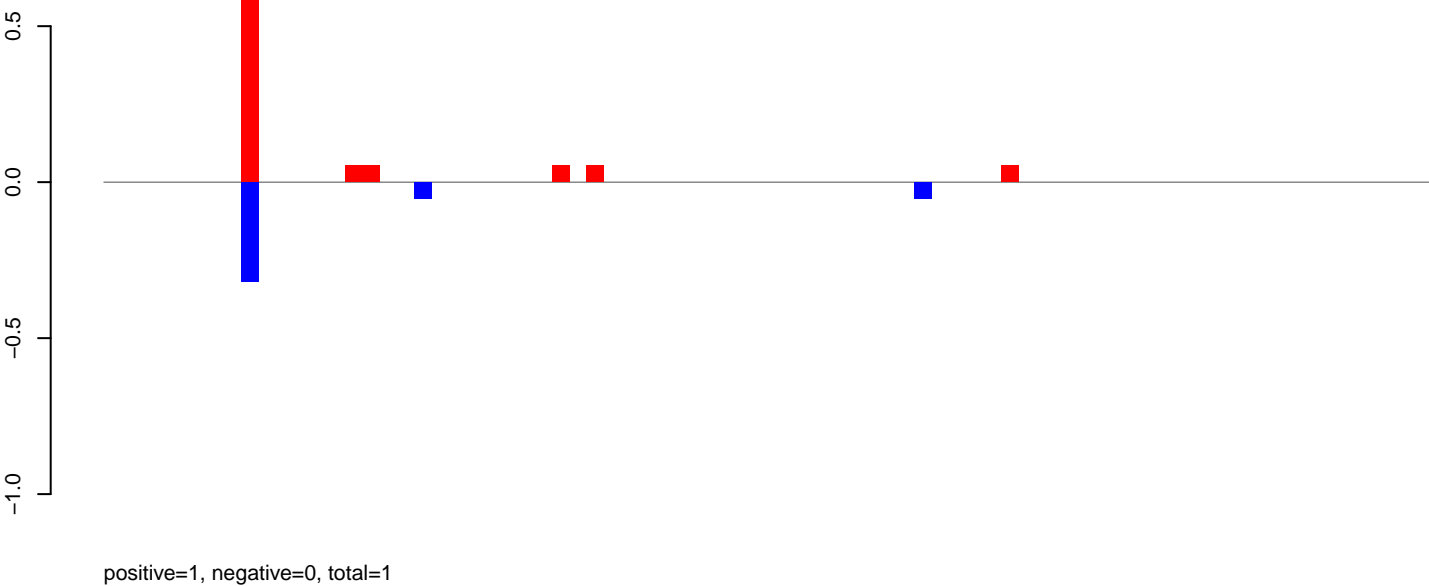
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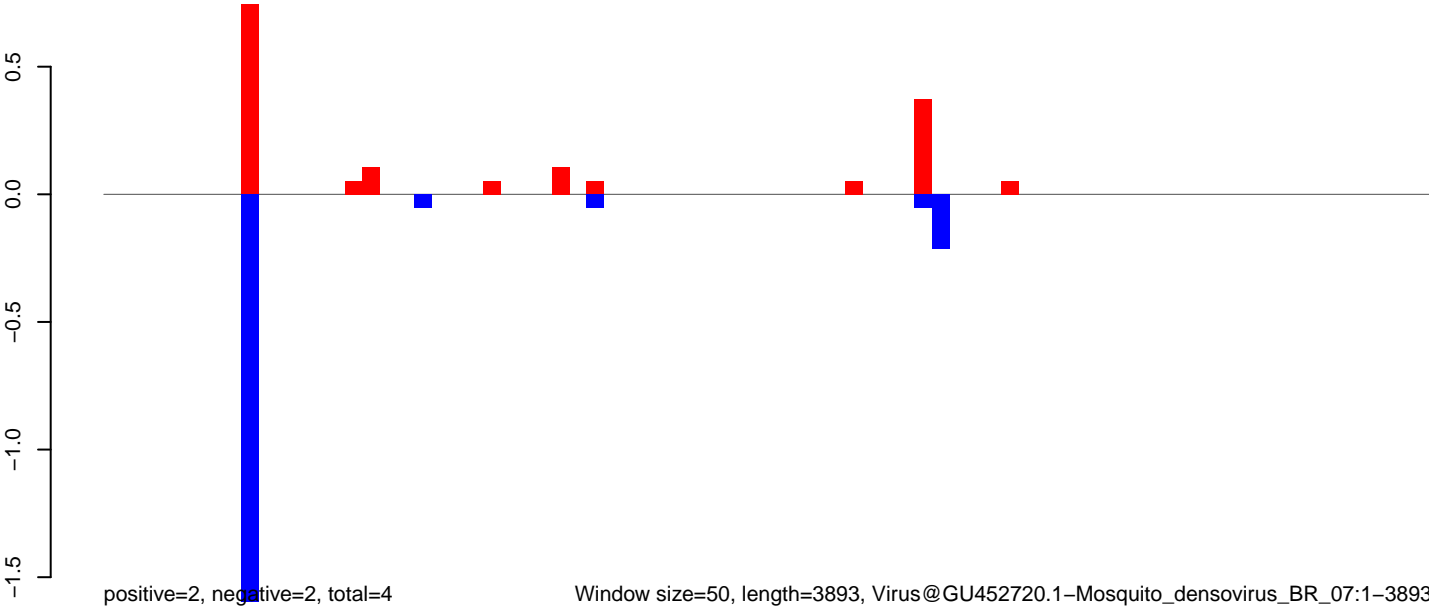
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



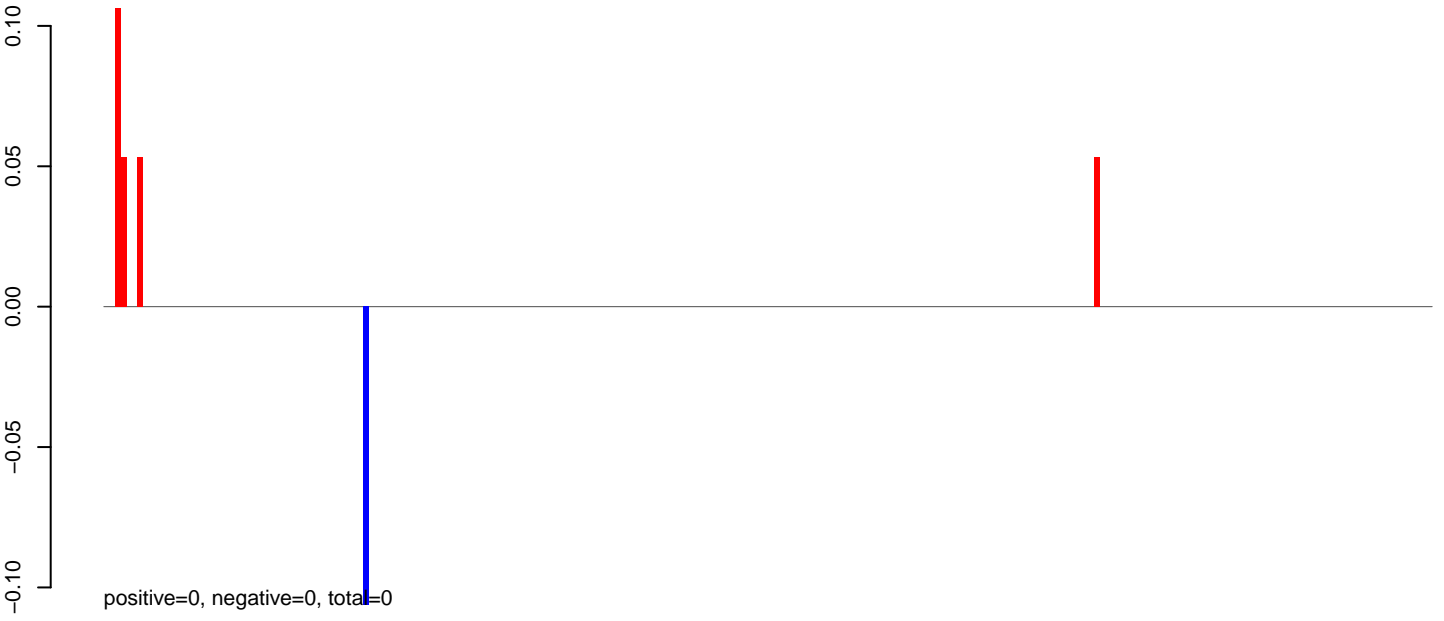
AeAeg_CCL.125cells_CHIKV.rep



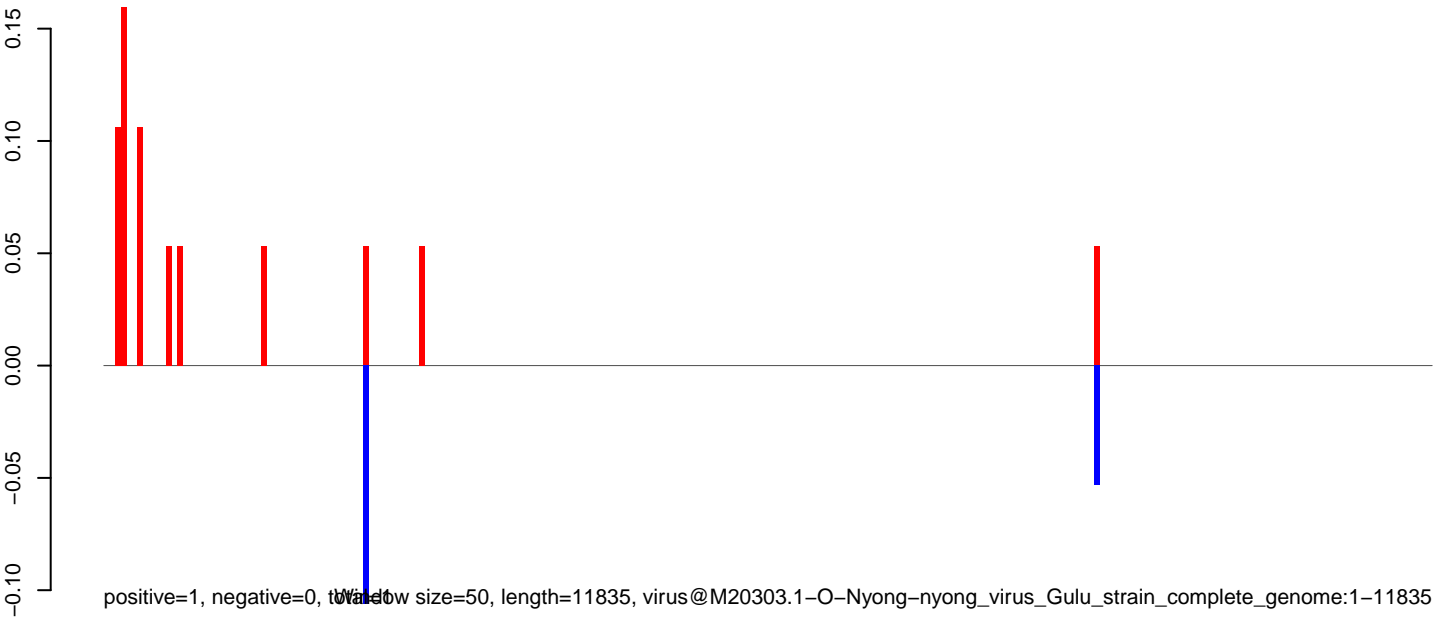
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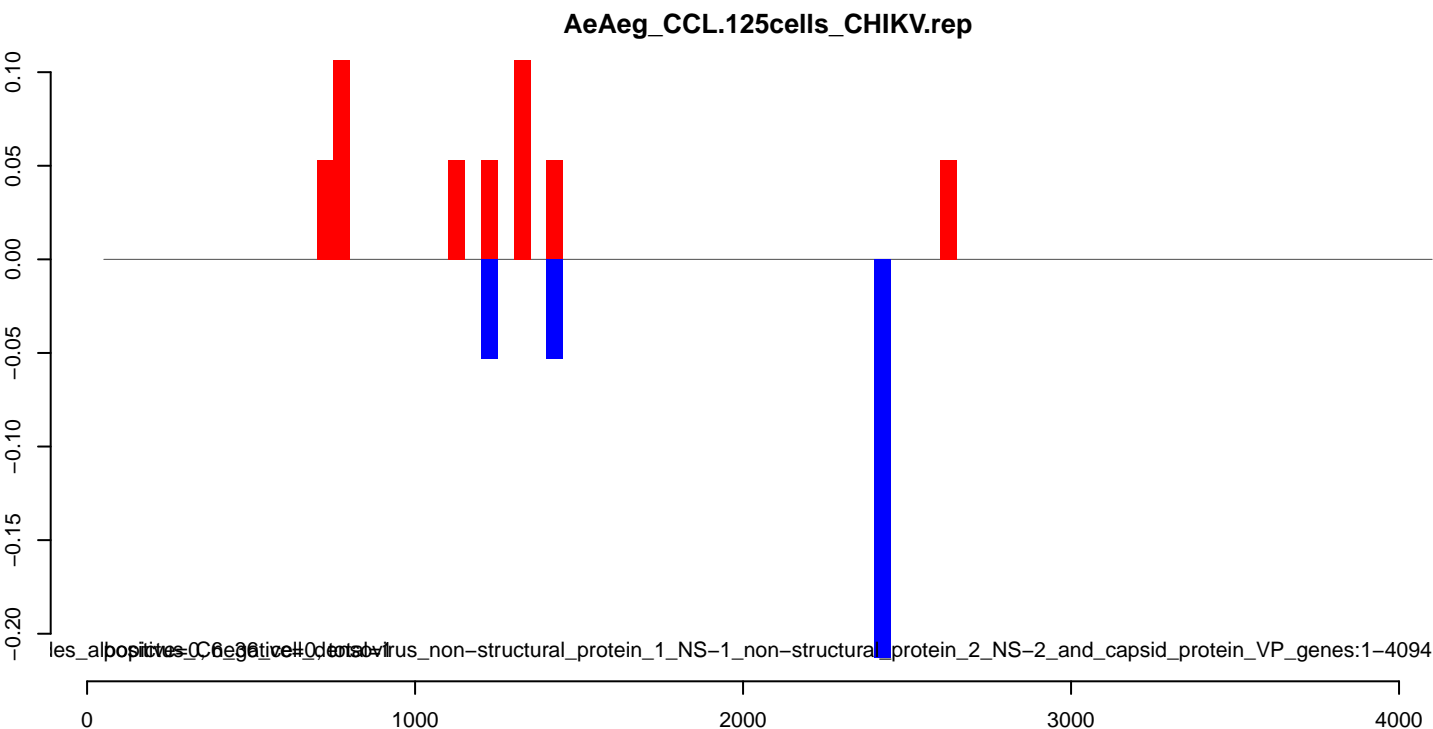
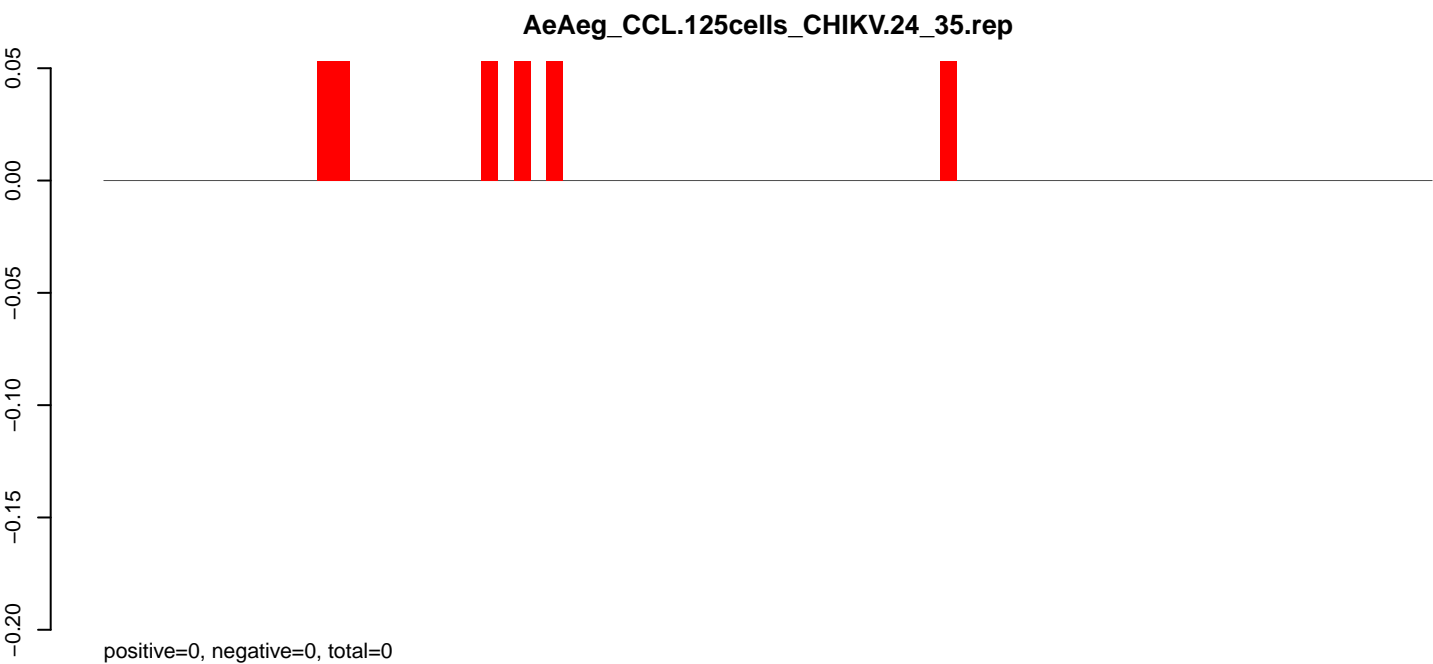
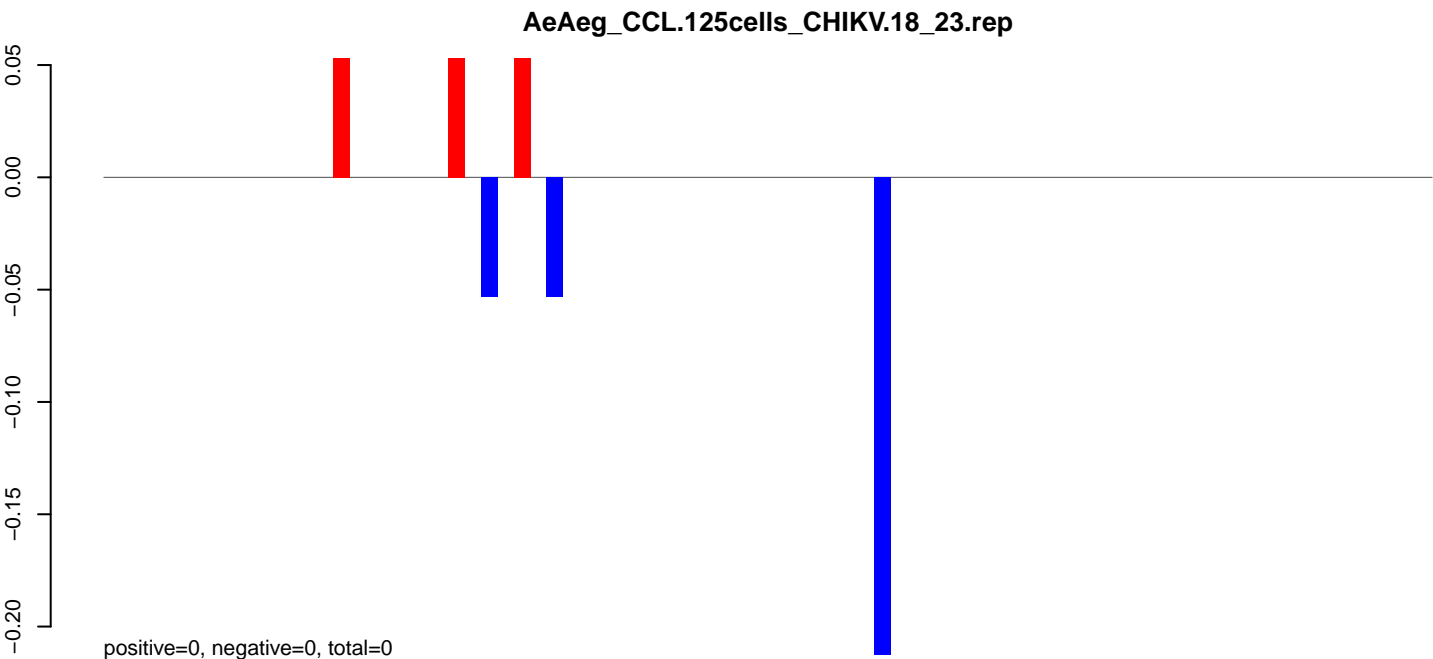


AeAeg_CCL.125cells_CHIKV.24_35.rep

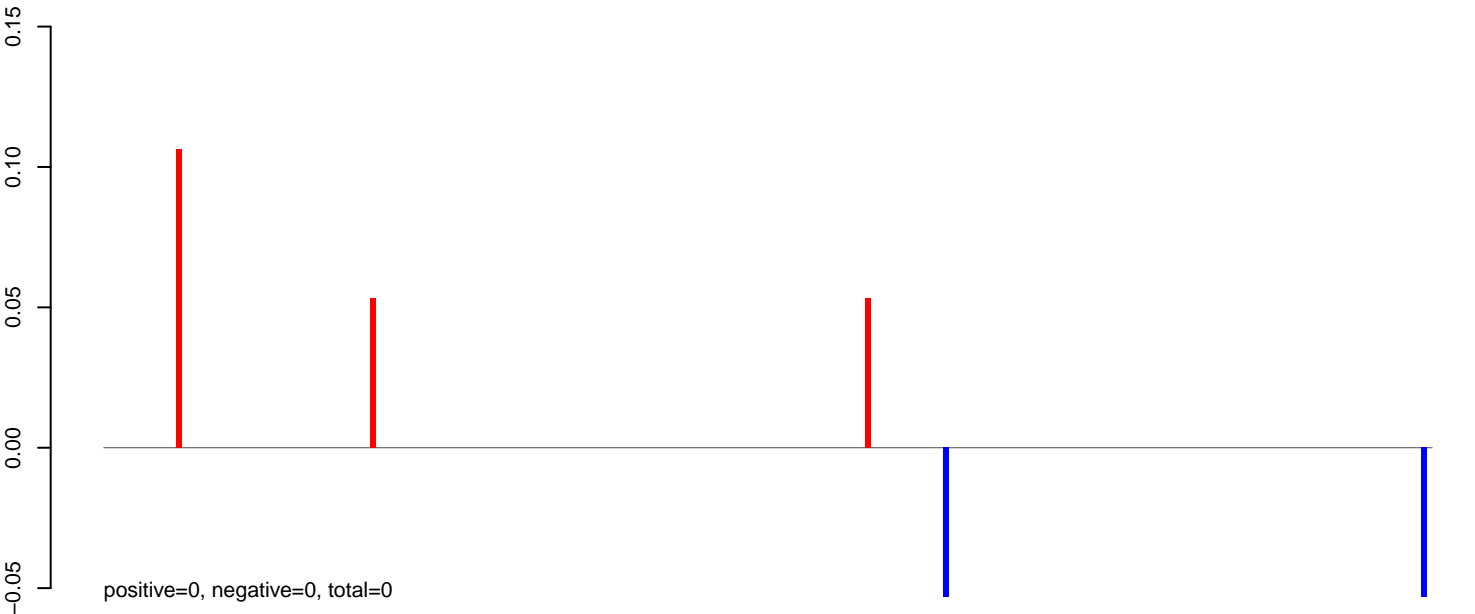


AeAeg_CCL.125cells_CHIKV.rep





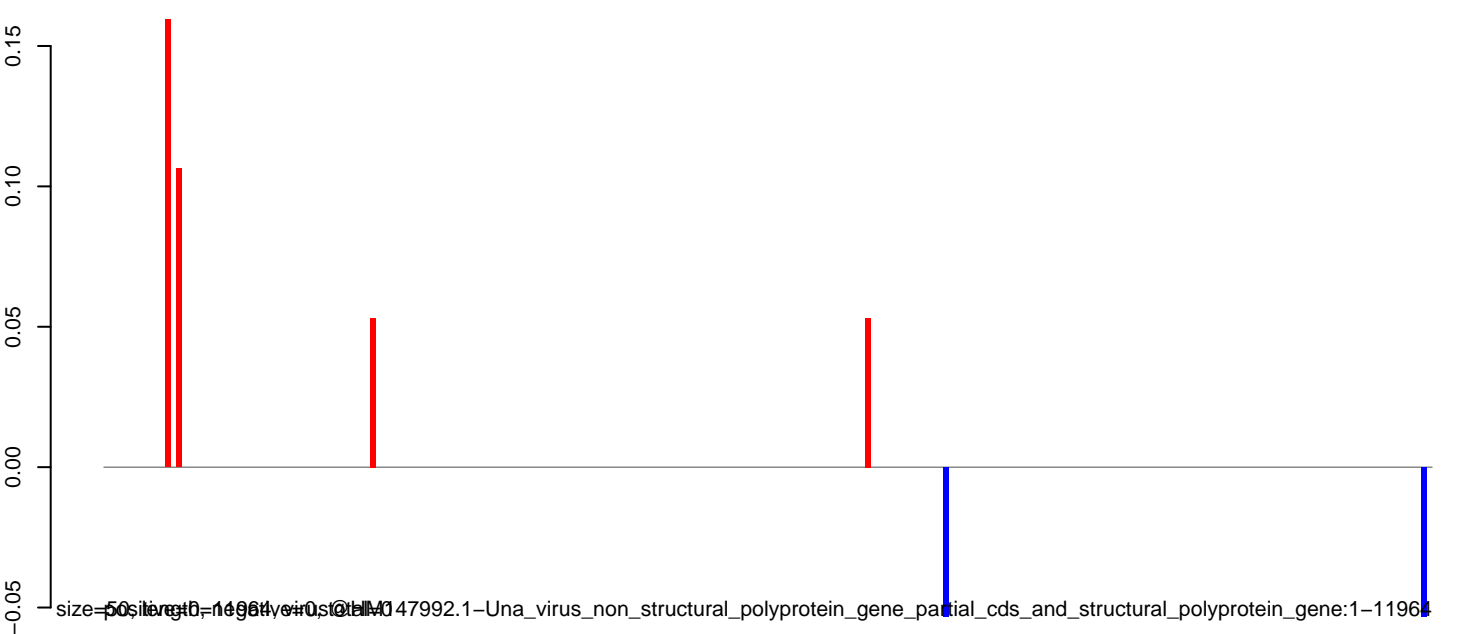
AeAeg_CCL.125cells_CHIKV.18_23.rep



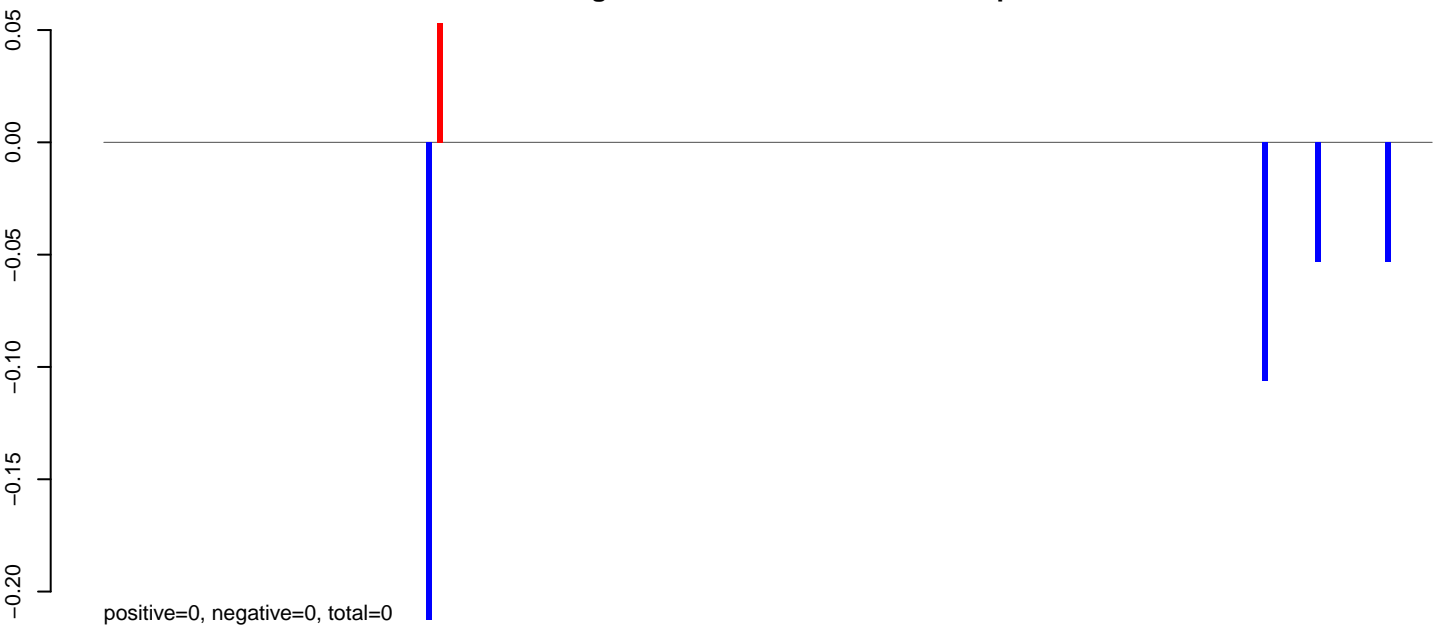
AeAeg_CCL.125cells_CHIKV.24_35.rep



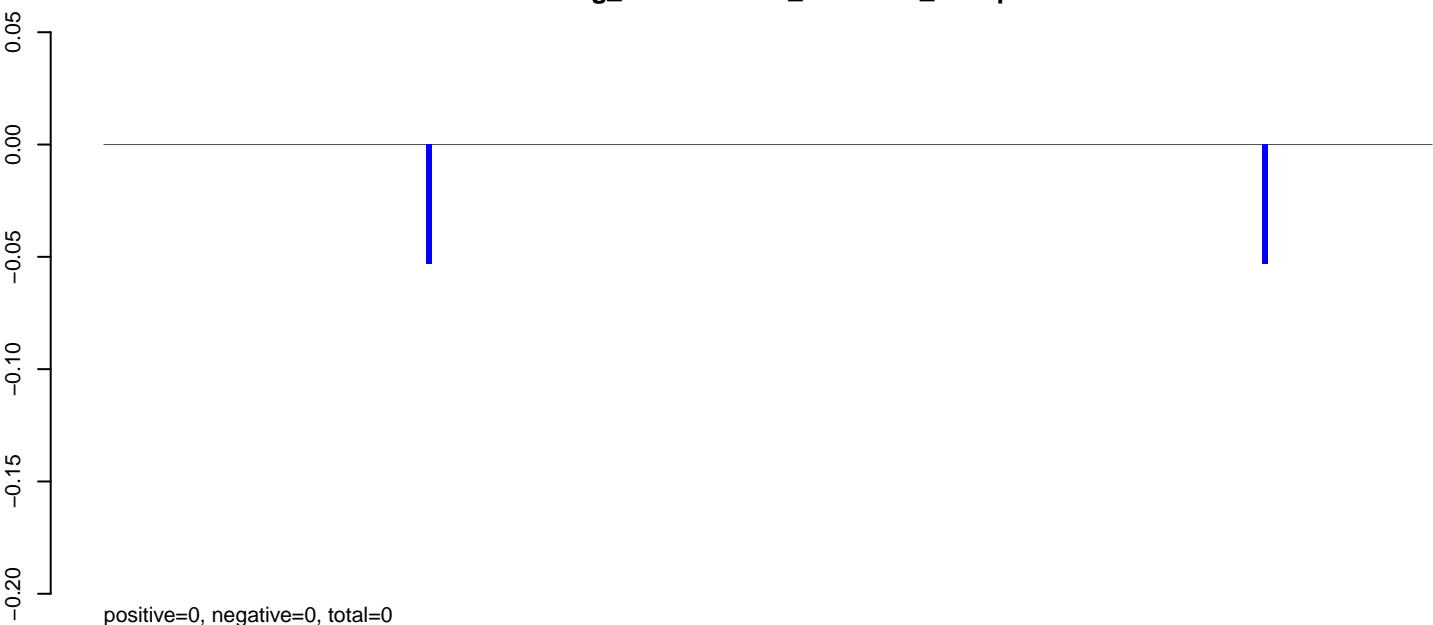
AeAeg_CCL.125cells_CHIKV.rep



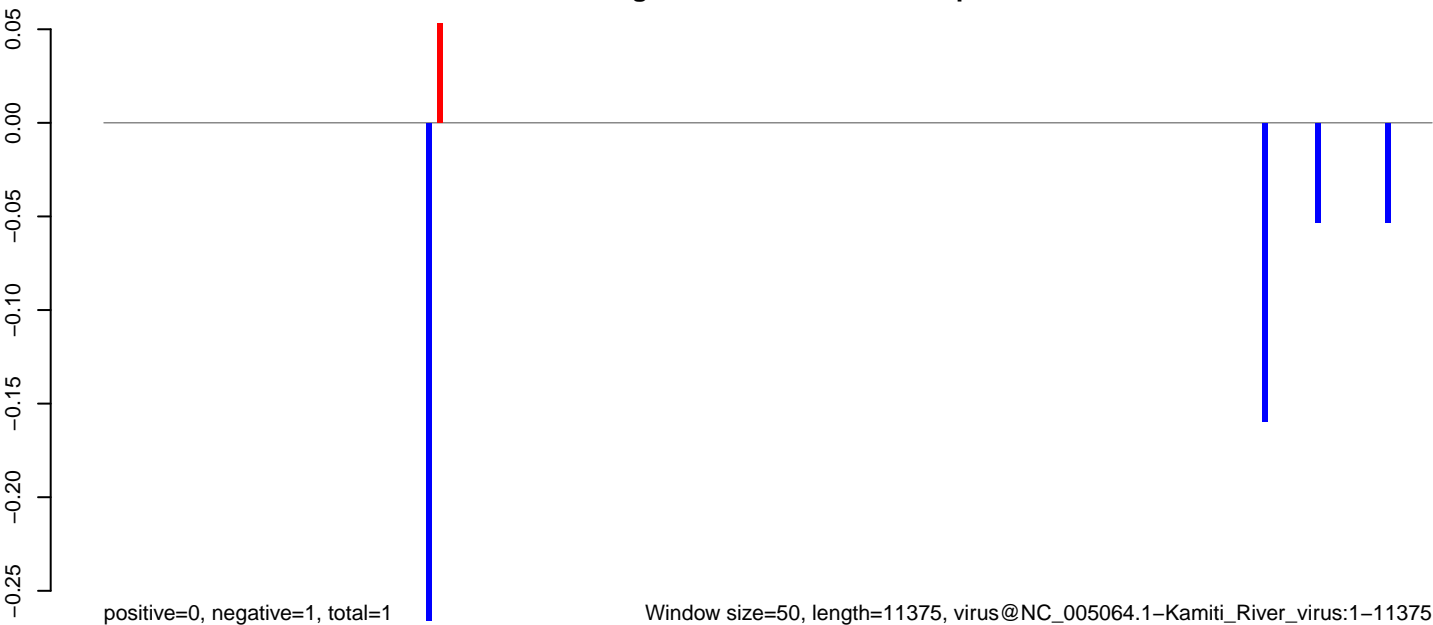
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep



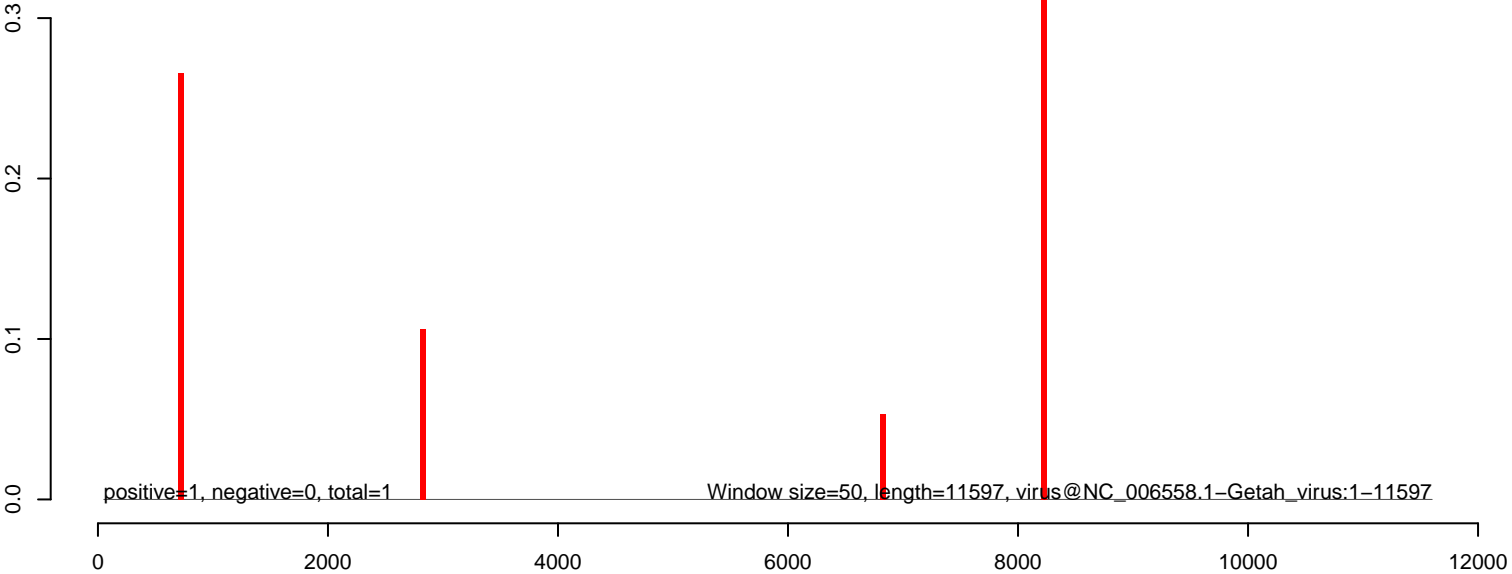
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AeAeg_CCL.125cells_CHIKV.24_35.rep



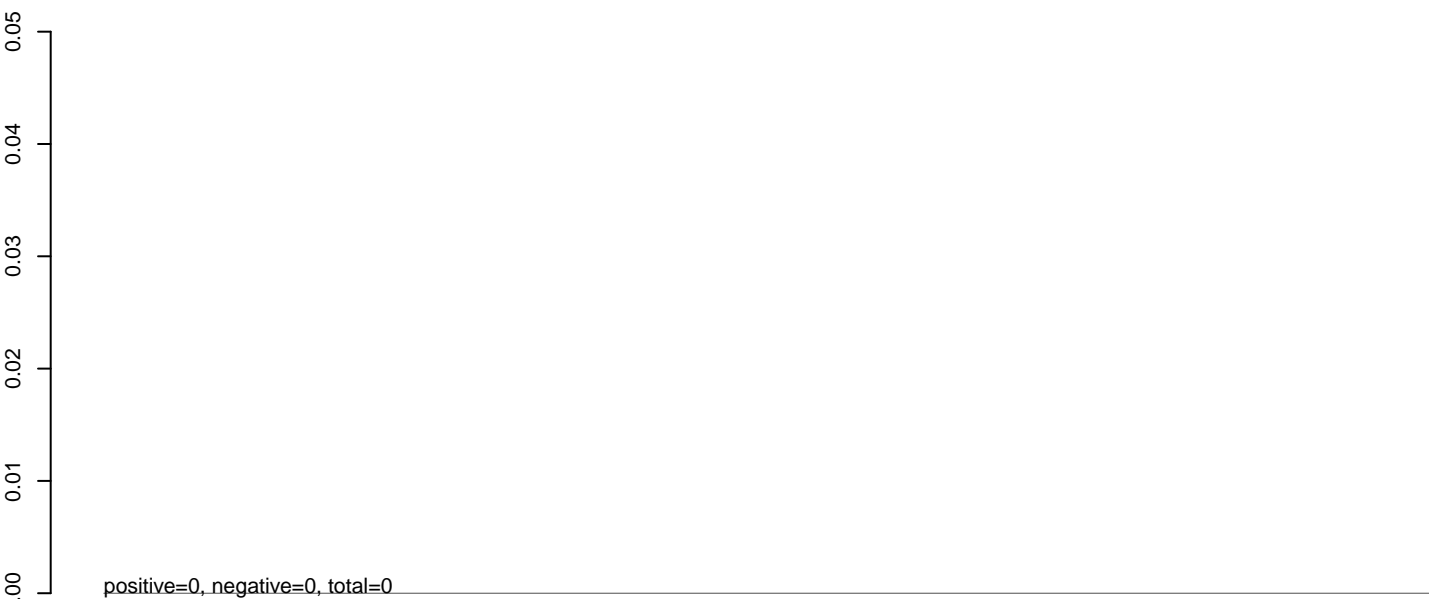
AeAeg_CCL.125cells_CHIKV.rep



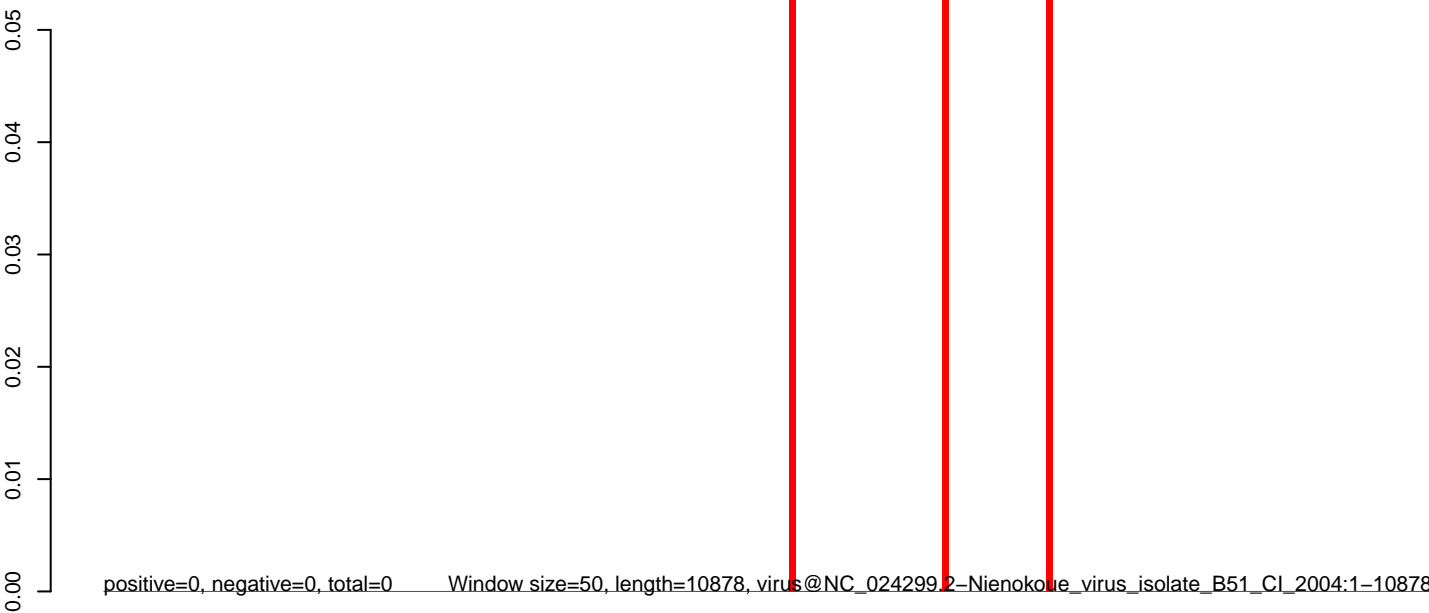
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

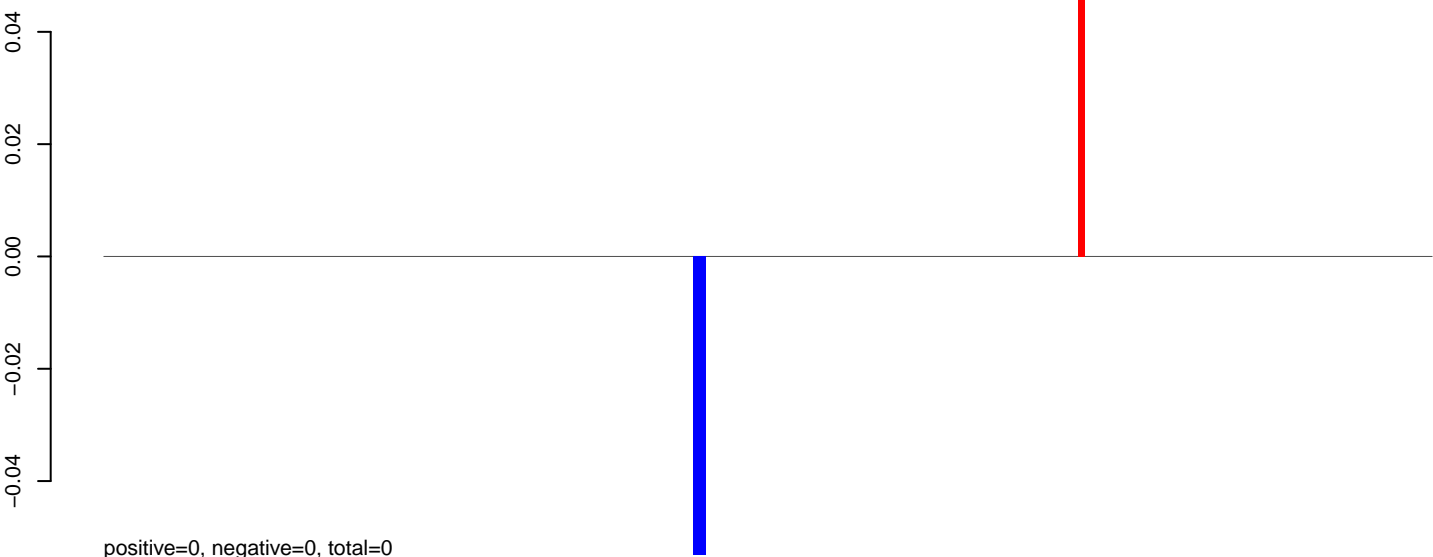


AeAeg_CCL.125cells_CHIKV.rep

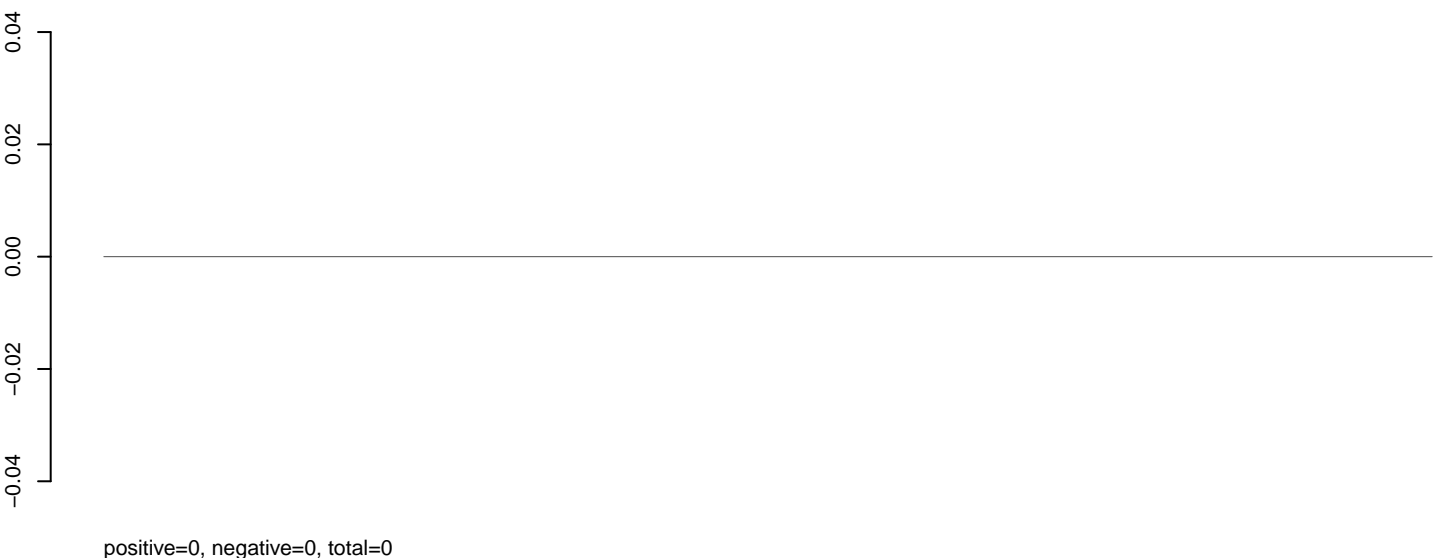


0 2000 4000 6000 8000 10000

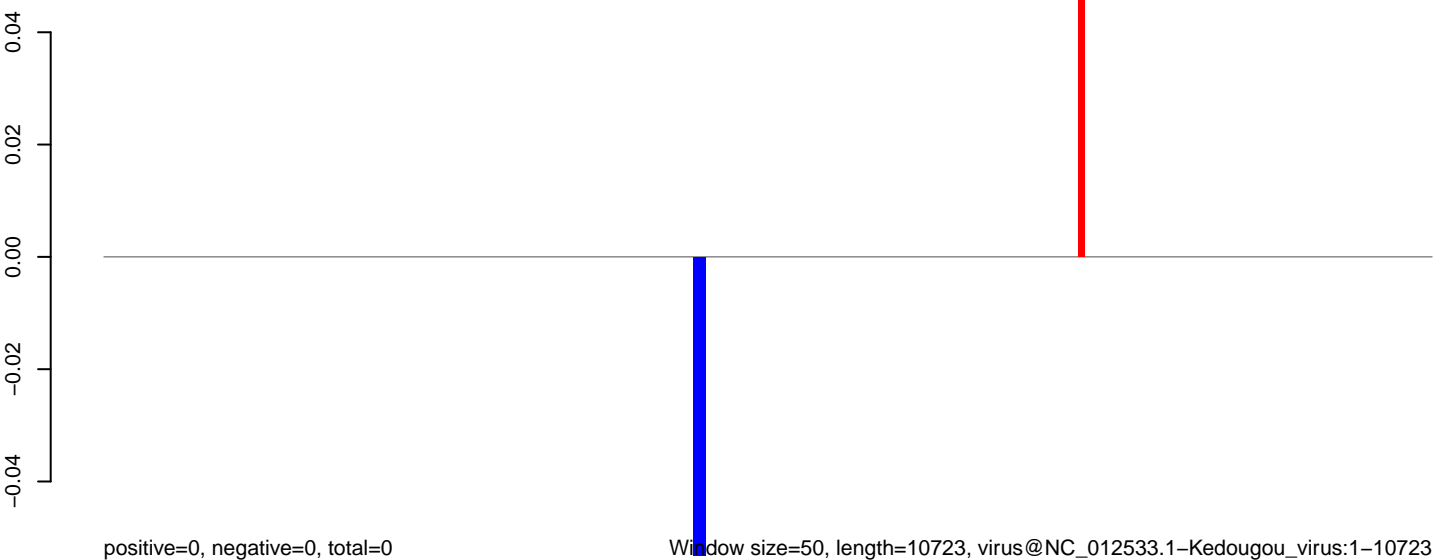
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep



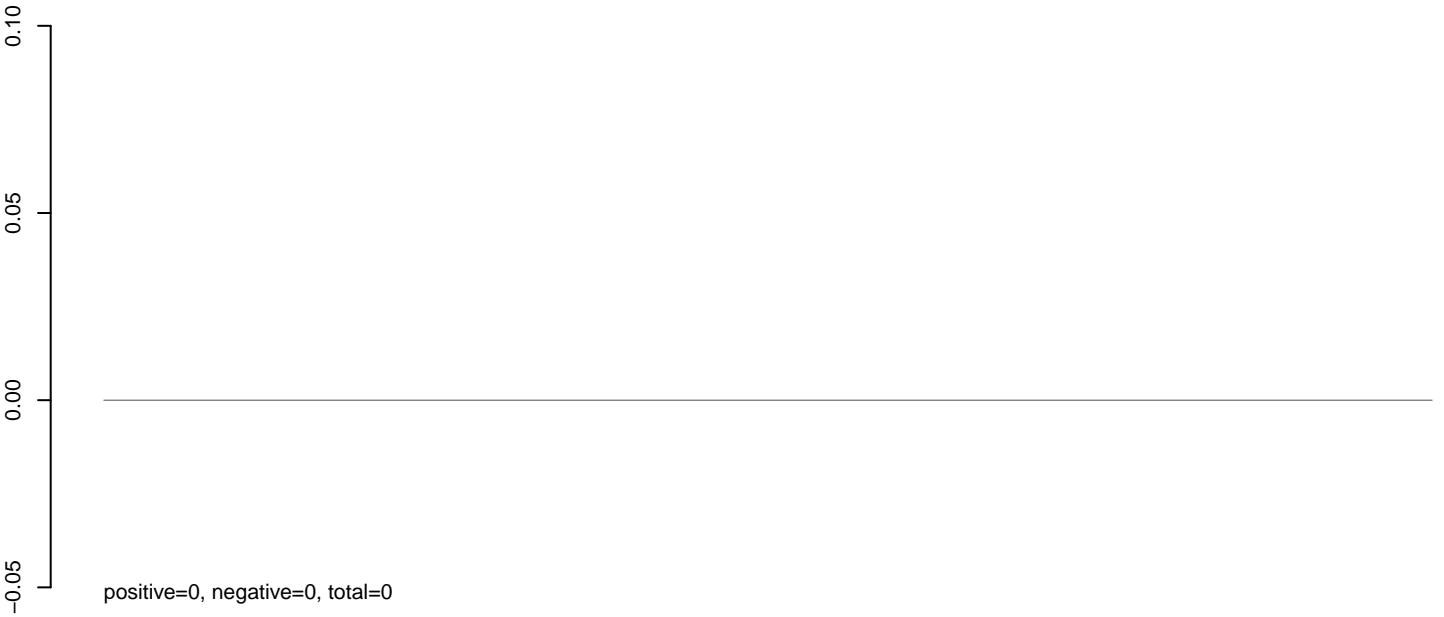
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0 2000 4000 6000 8000 10000

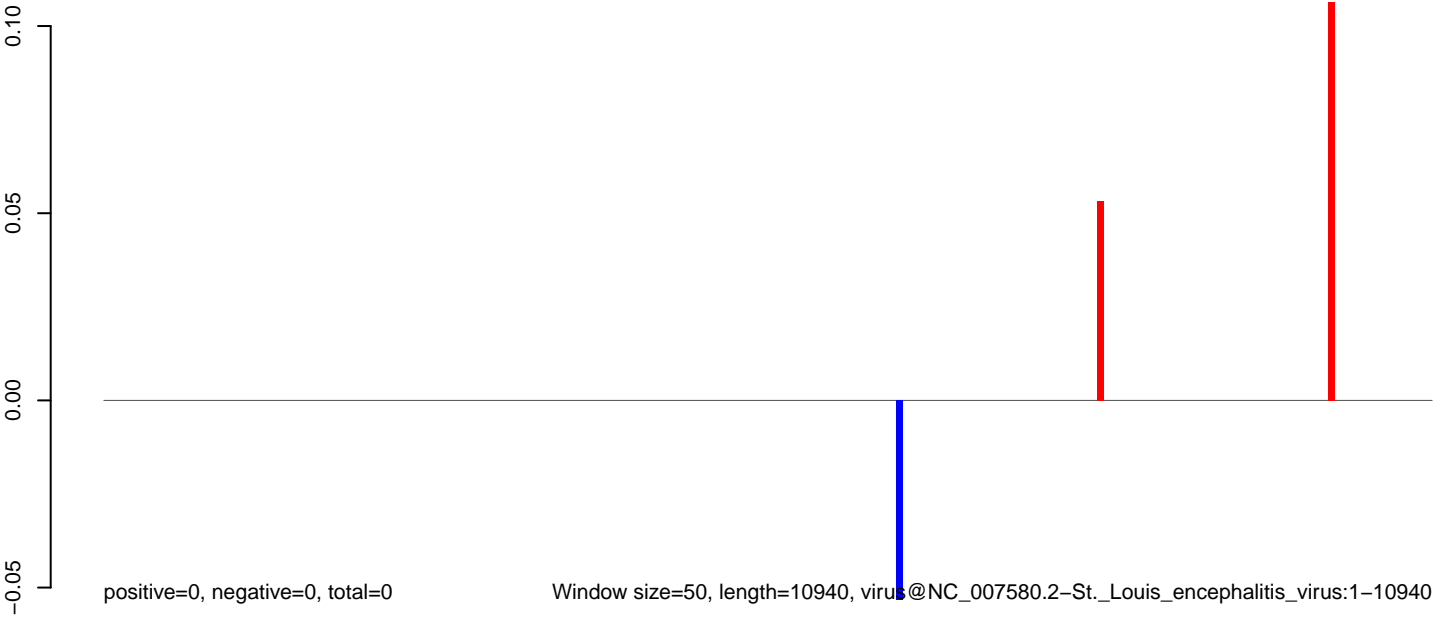
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep



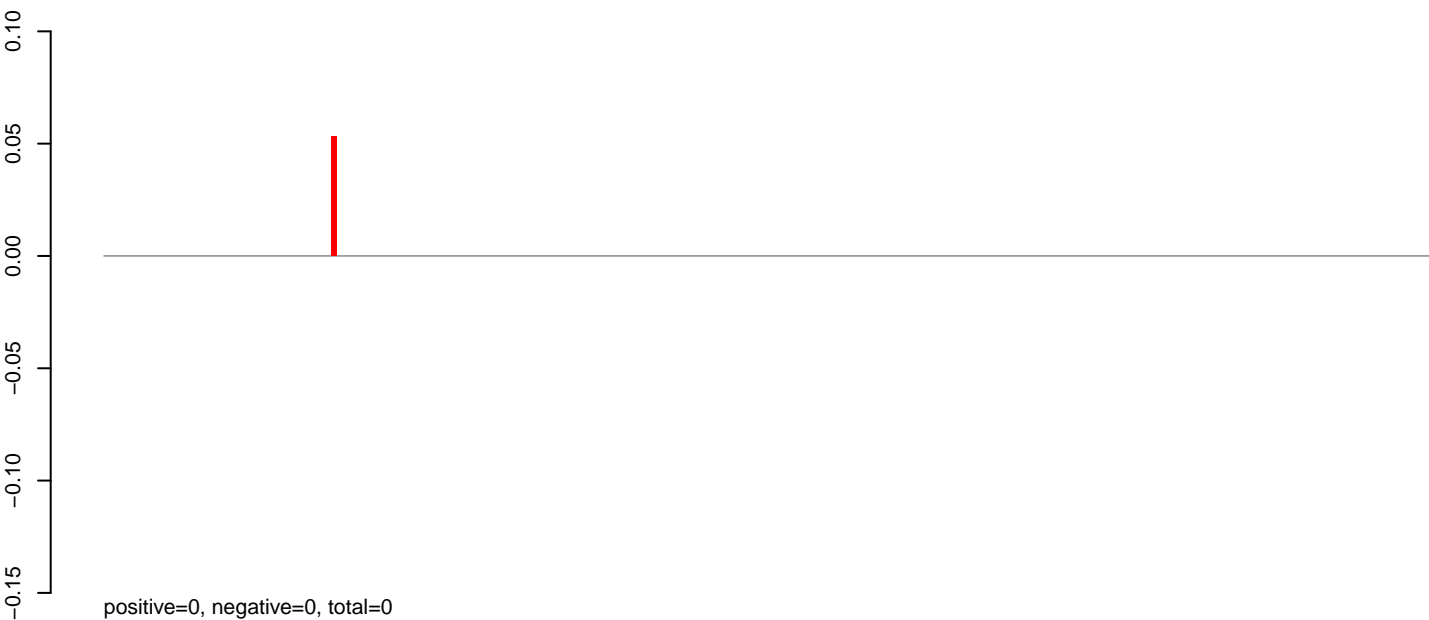
Window size=50, length=10940, virus@NC_007580.2-St._Louis_encephalitis_virus:1-10940

0 2000 4000 6000 8000 10000

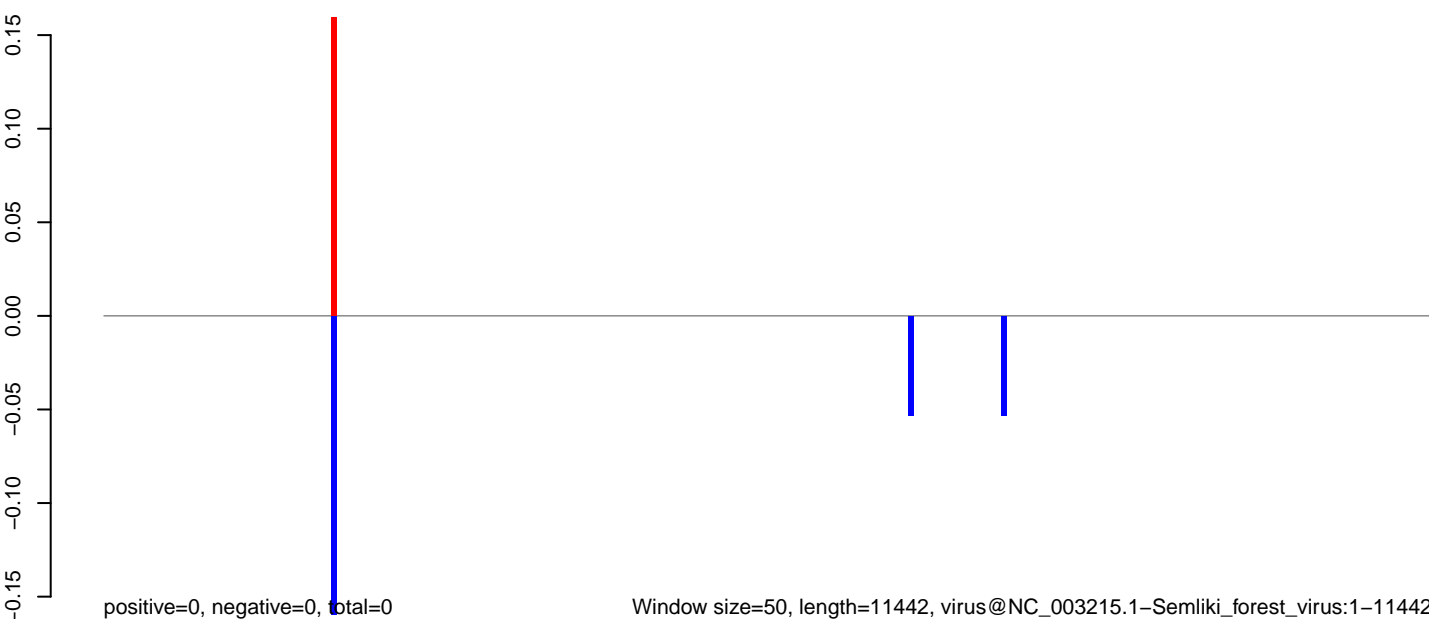
AeAeg_CCL.125cells_CHIKV.18_23.rep



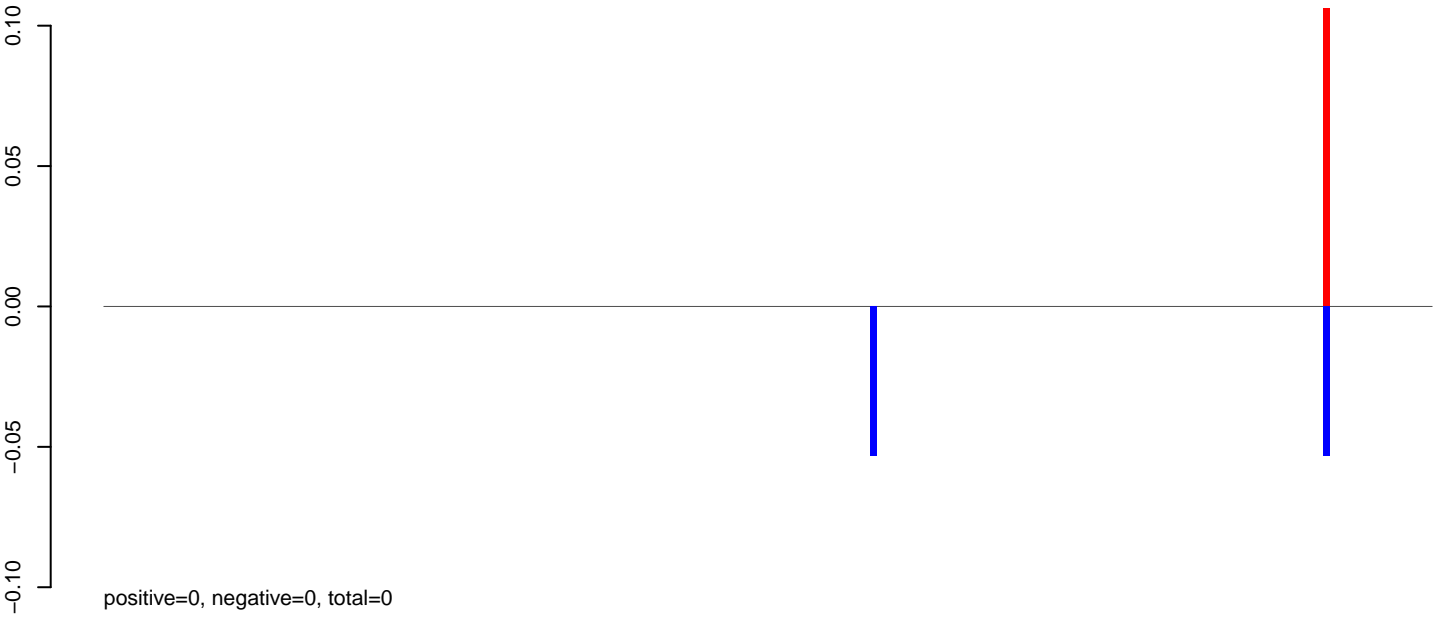
AeAeg_CCL.125cells_CHIKV.24_35.rep



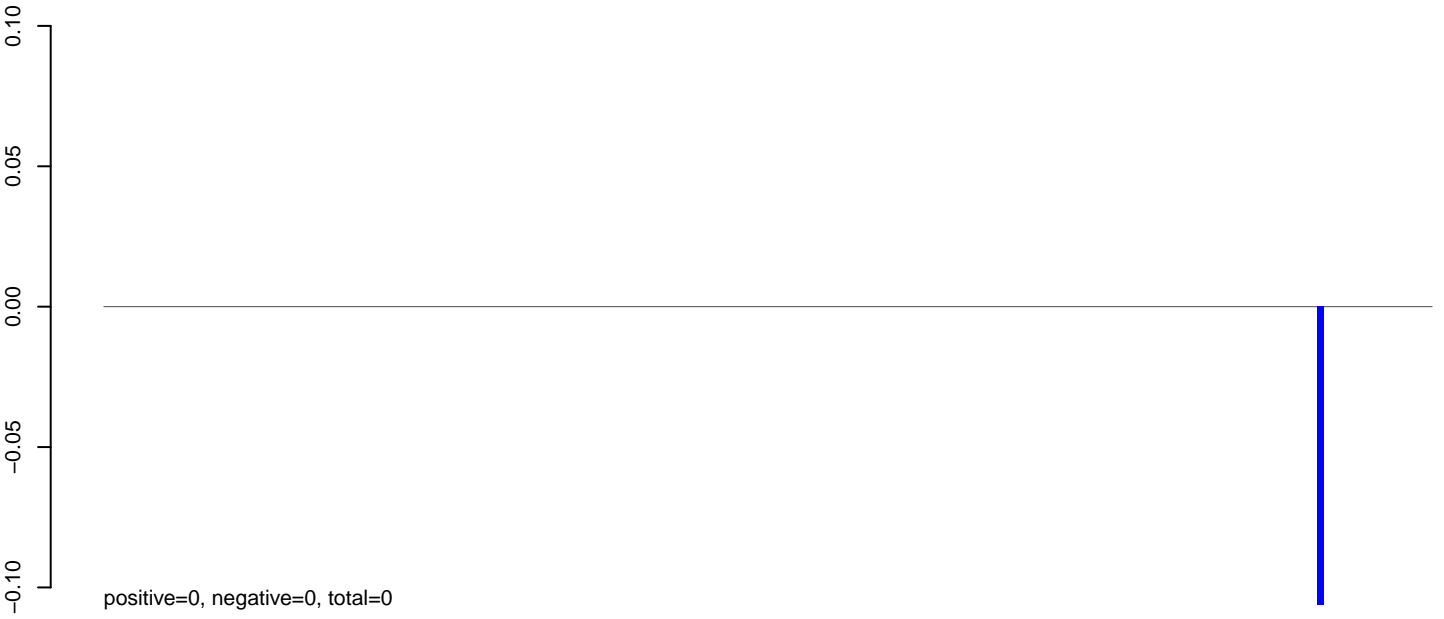
AeAeg_CCL.125cells_CHIKV.rep



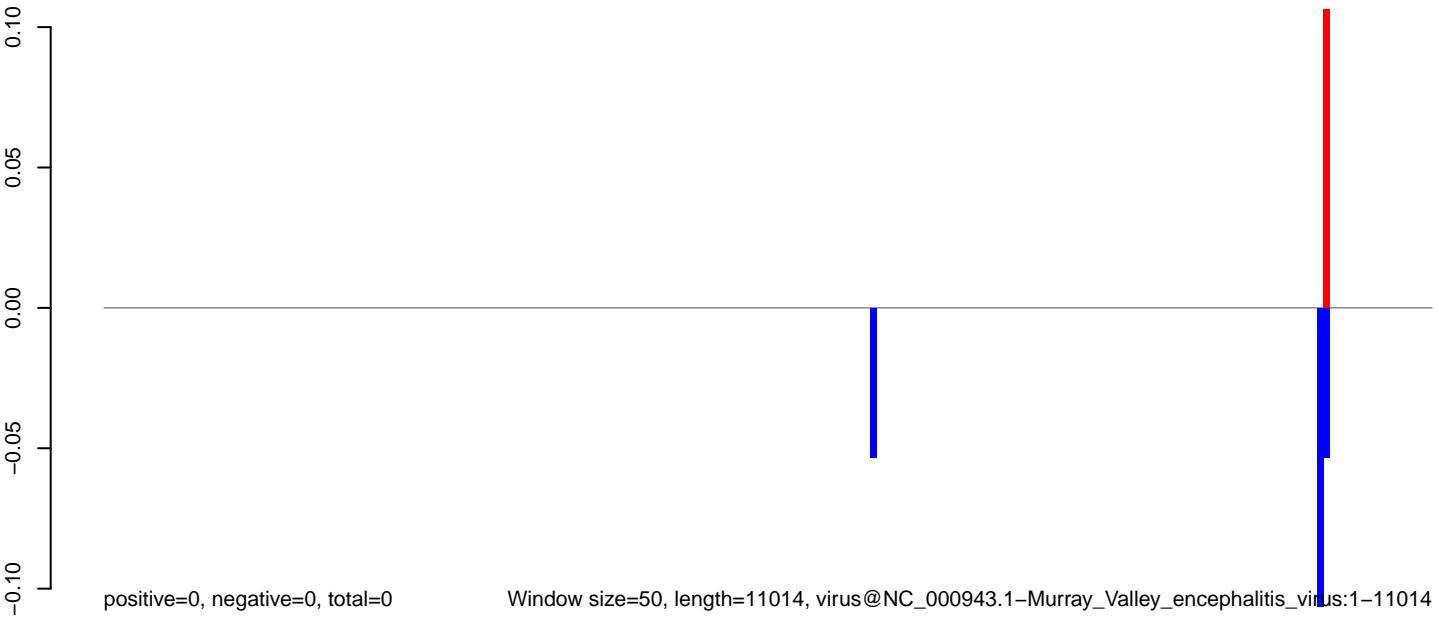
AeAeg_CCL.125cells_CHIKV.18_23.rep



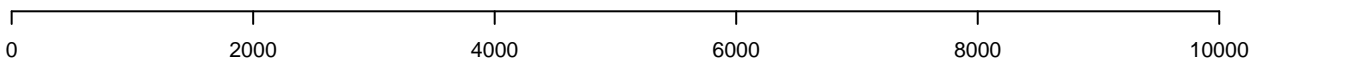
AeAeg_CCL.125cells_CHIKV.24_35.rep



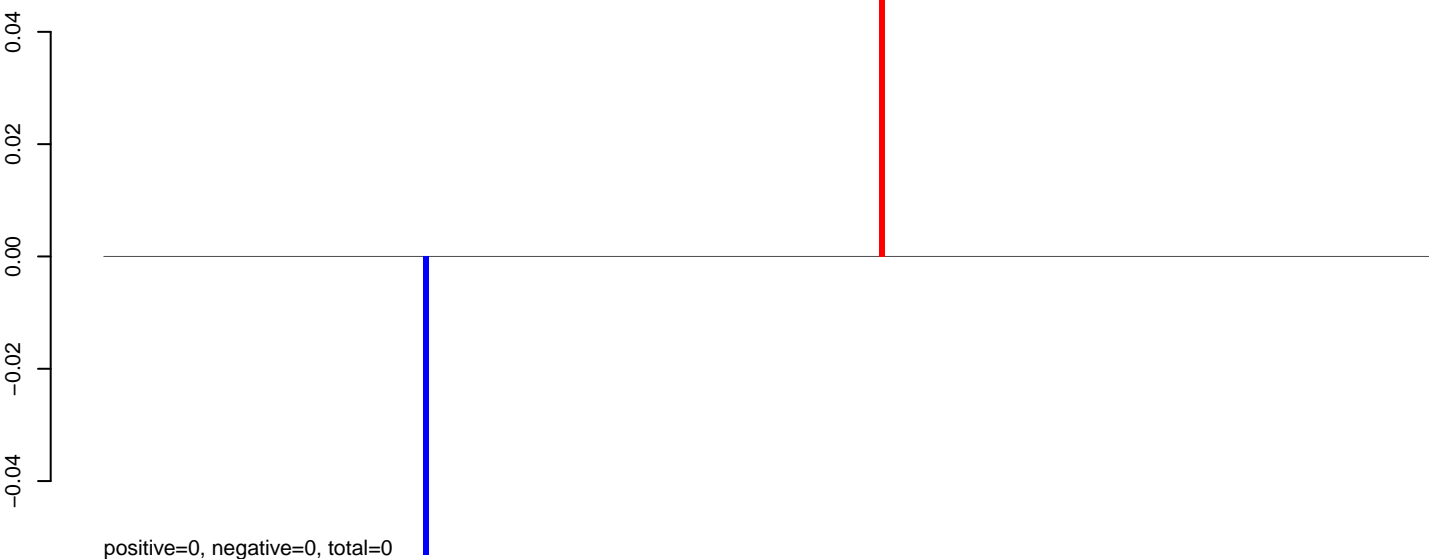
AeAeg_CCL.125cells_CHIKV.rep



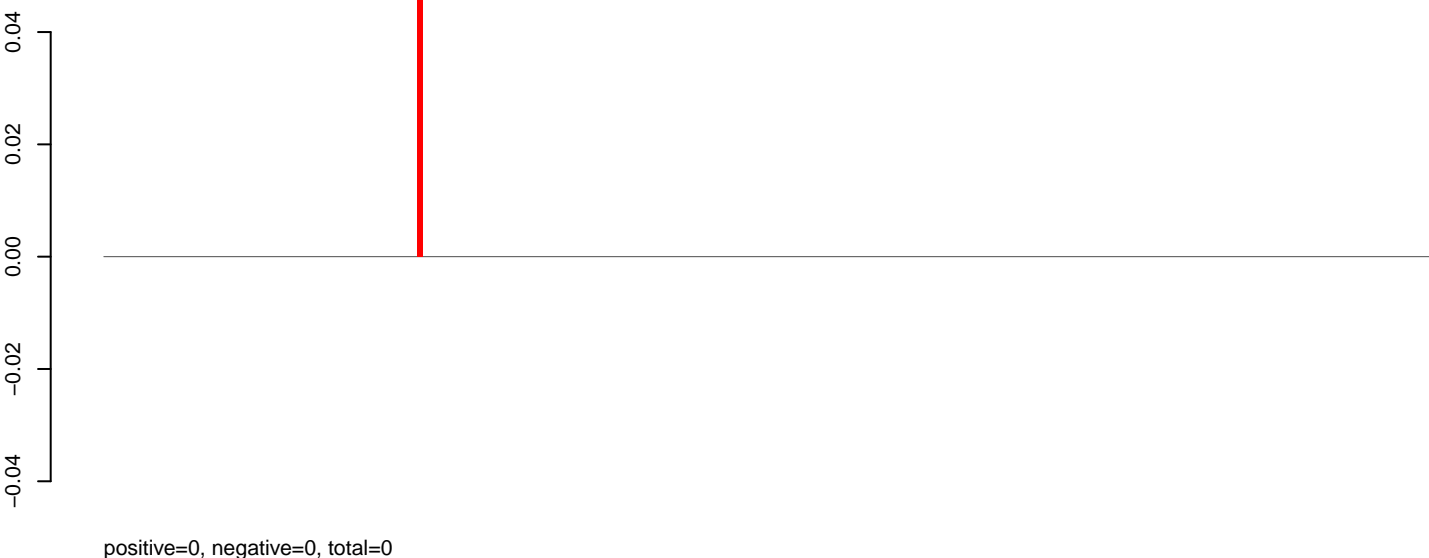
Window size=50, length=11014, virus@NC_000943.1-Murray_Valley_encephalitis_virus:1-11014



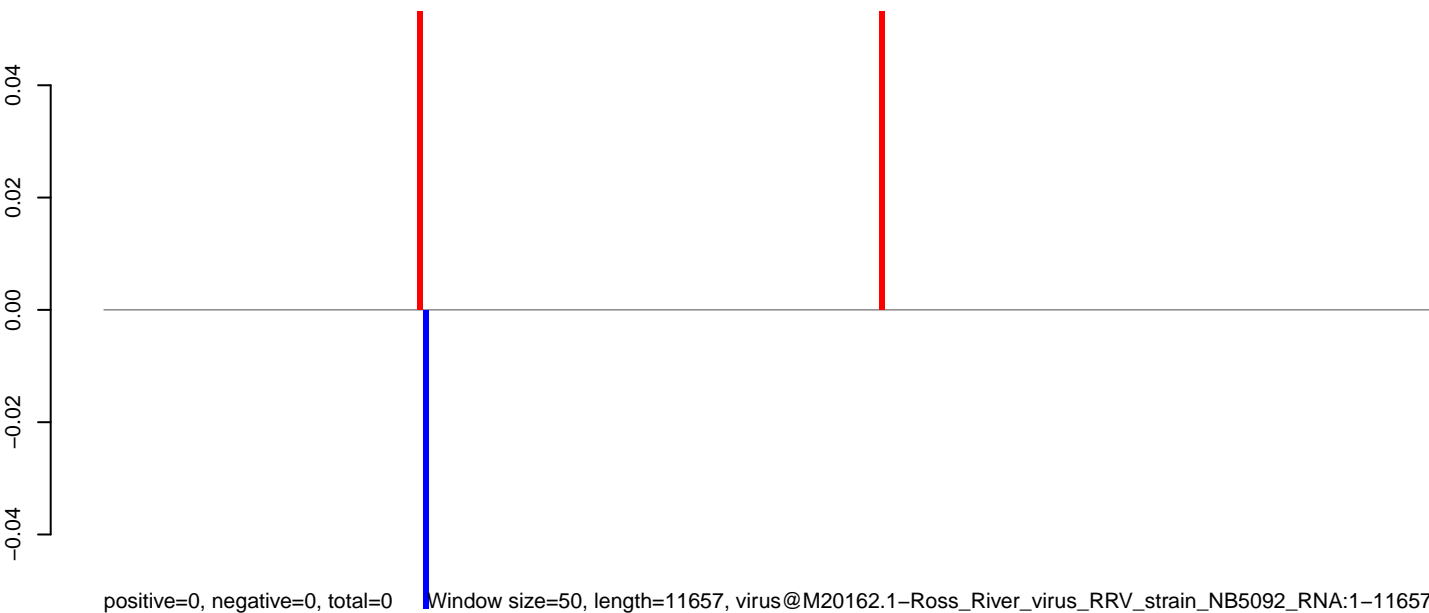
AeAeg_CCL.125cells_CHIKV.18_23.rep



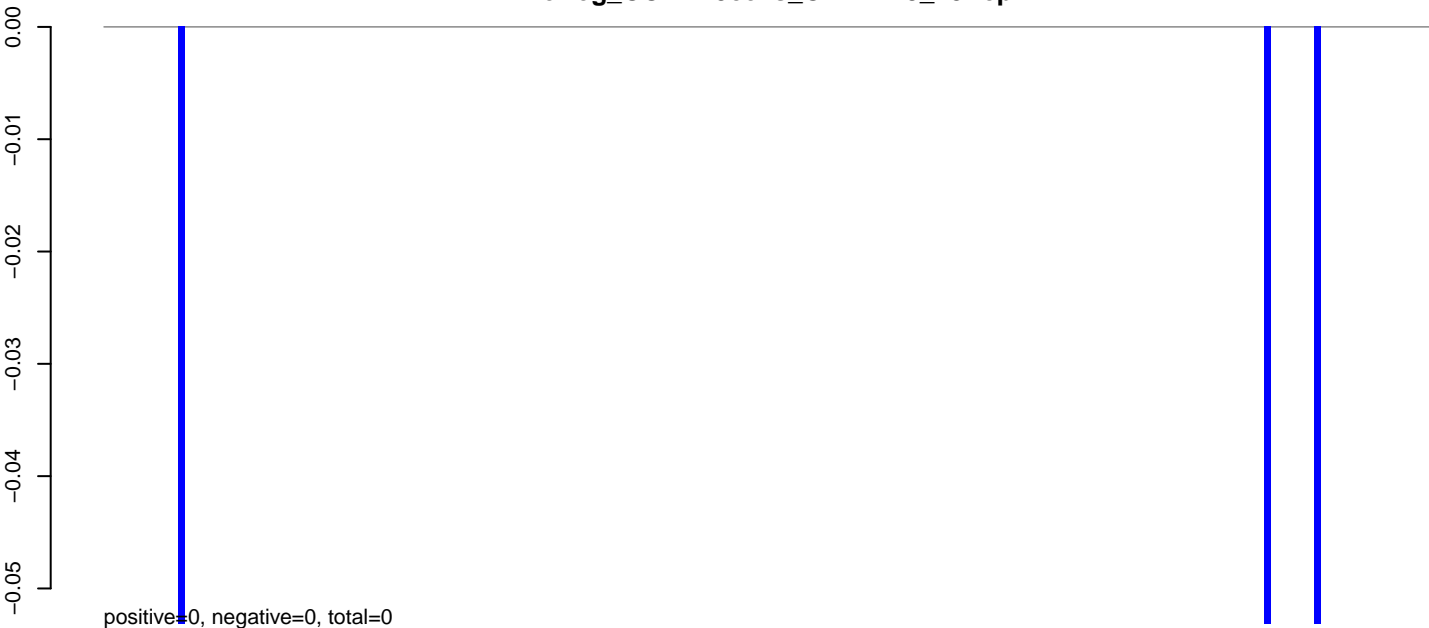
AeAeg_CCL.125cells_CHIKV.24_35.rep



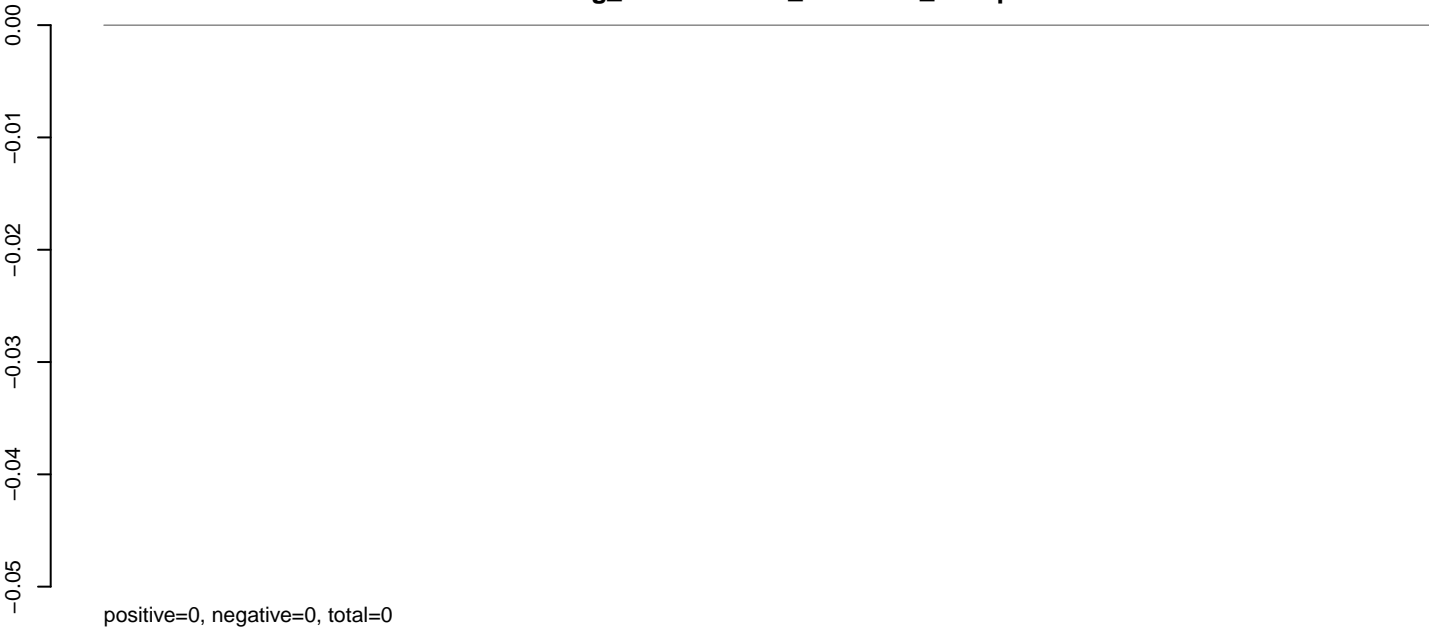
AeAeg_CCL.125cells_CHIKV.rep



AeAeg_CCL.125cells_CHIKV.18_23.rep



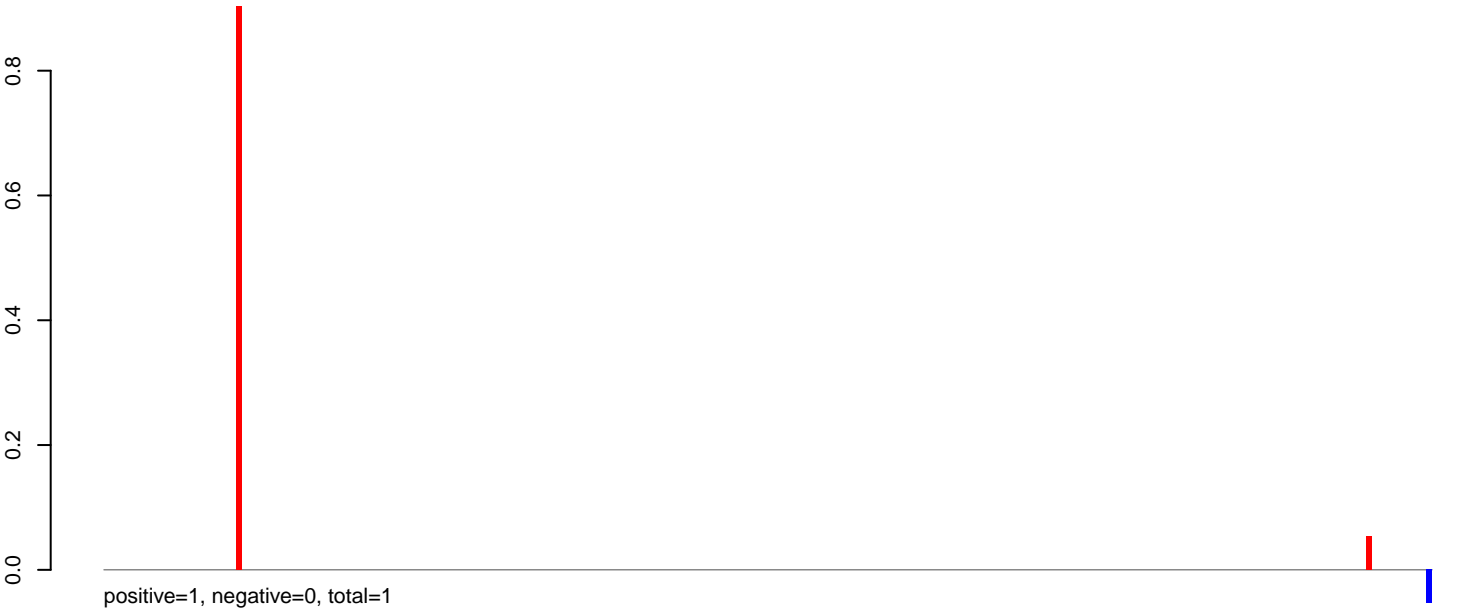
AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep



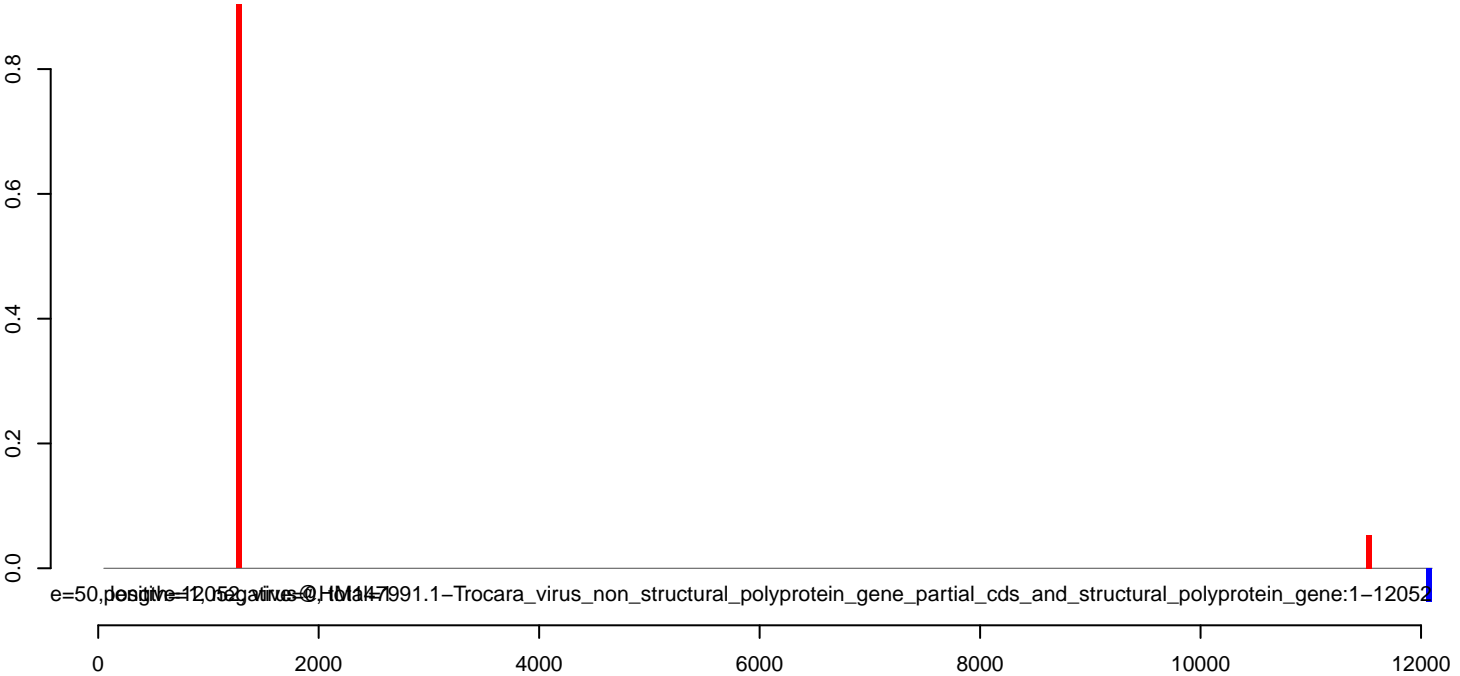
AeAeg_CCL.125cells_CHIKV.18_23.rep



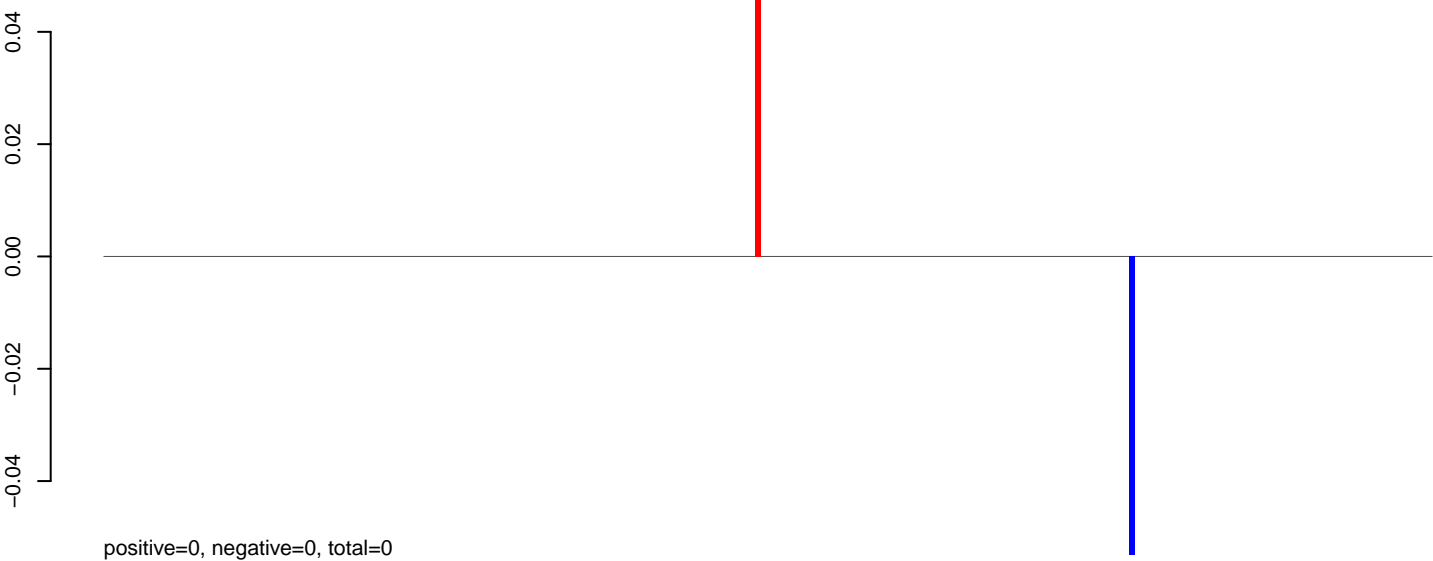
AeAeg_CCL.125cells_CHIKV.24_35.rep



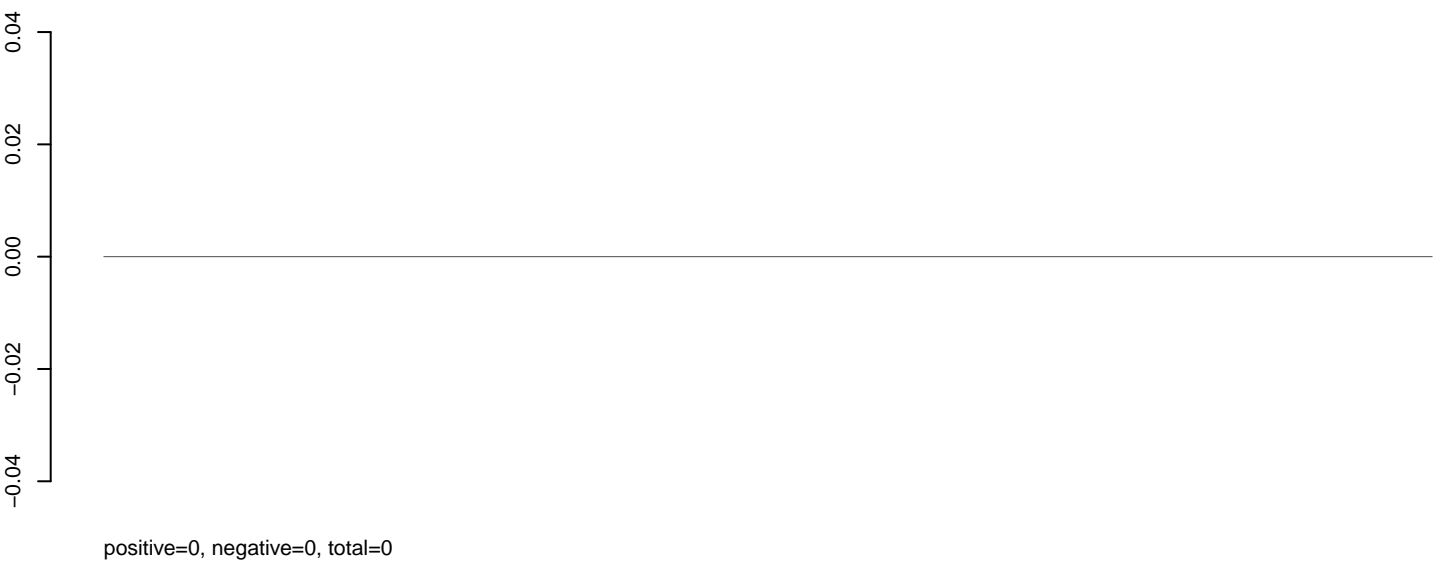
AeAeg_CCL.125cells_CHIKV.rep



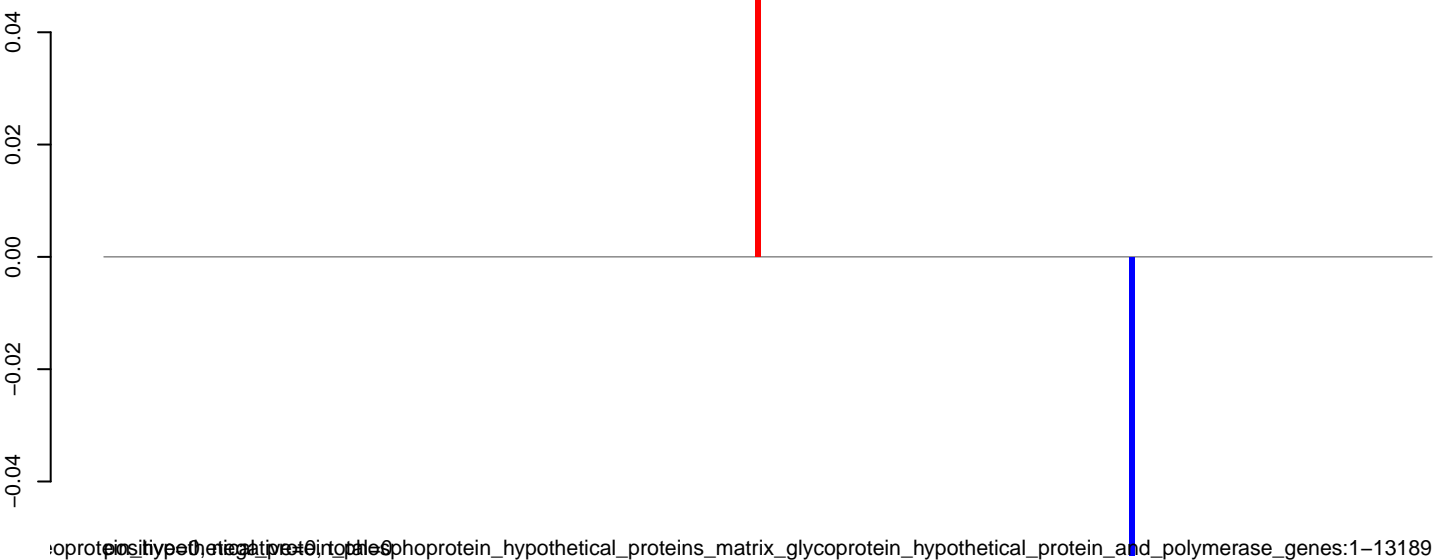
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



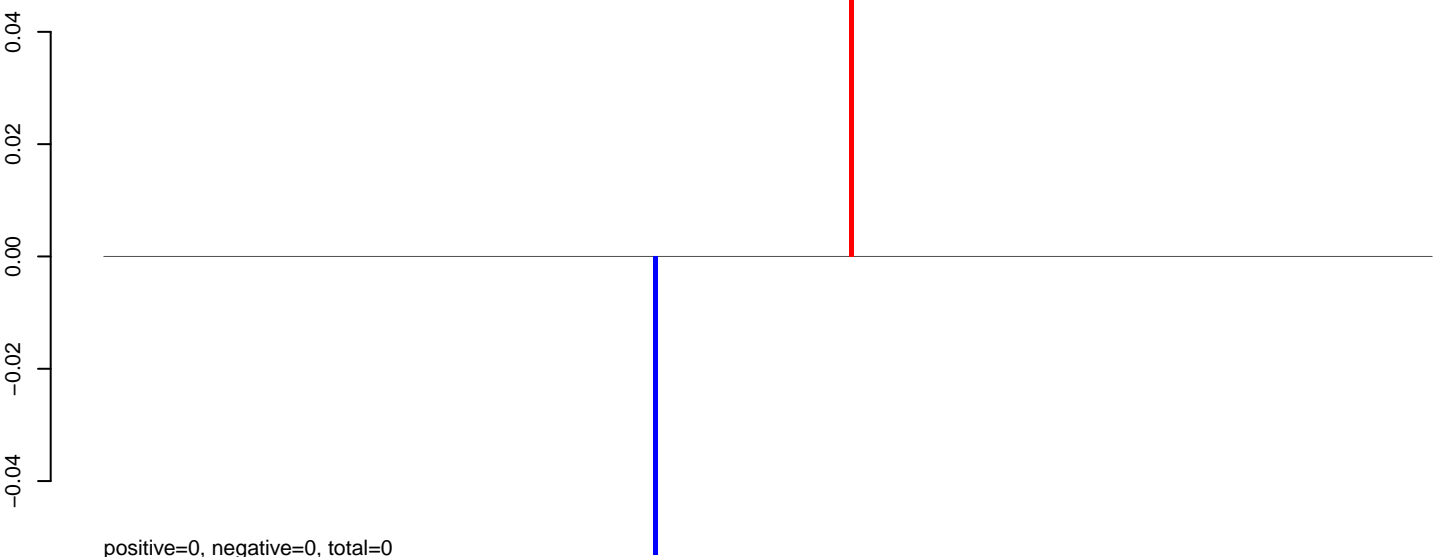
AeAeg_CCL.125cells_CHIKV.rep



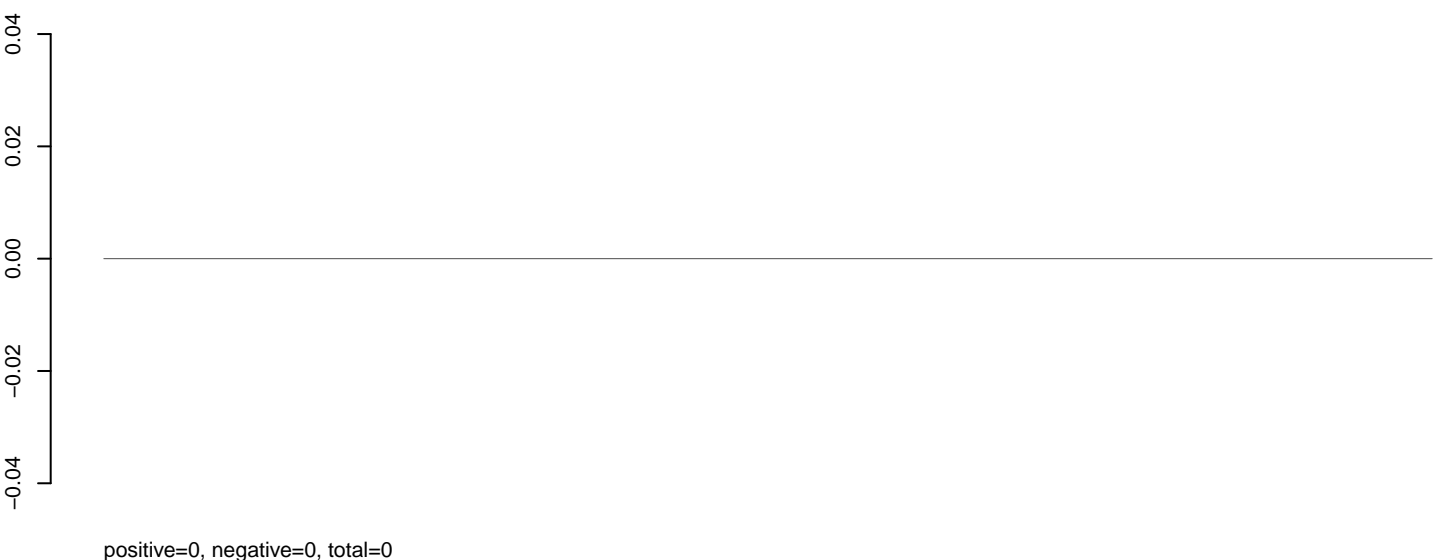
protein_hypothetical_protein_13189

0 2000 4000 6000 8000 10000 12000

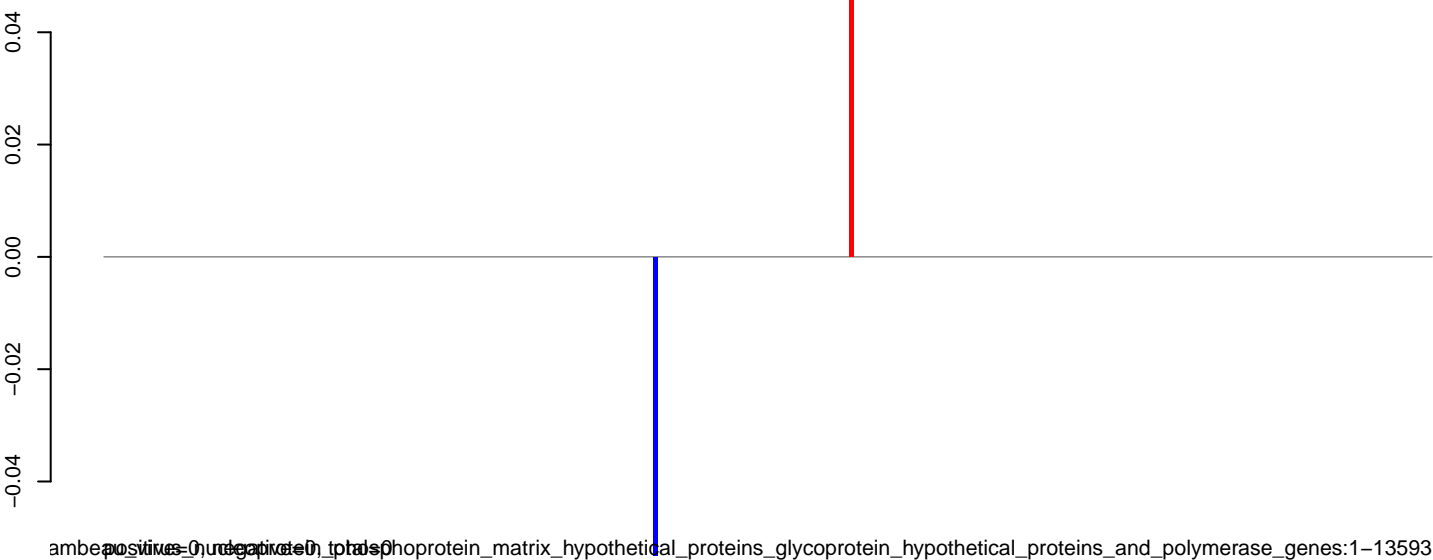
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep



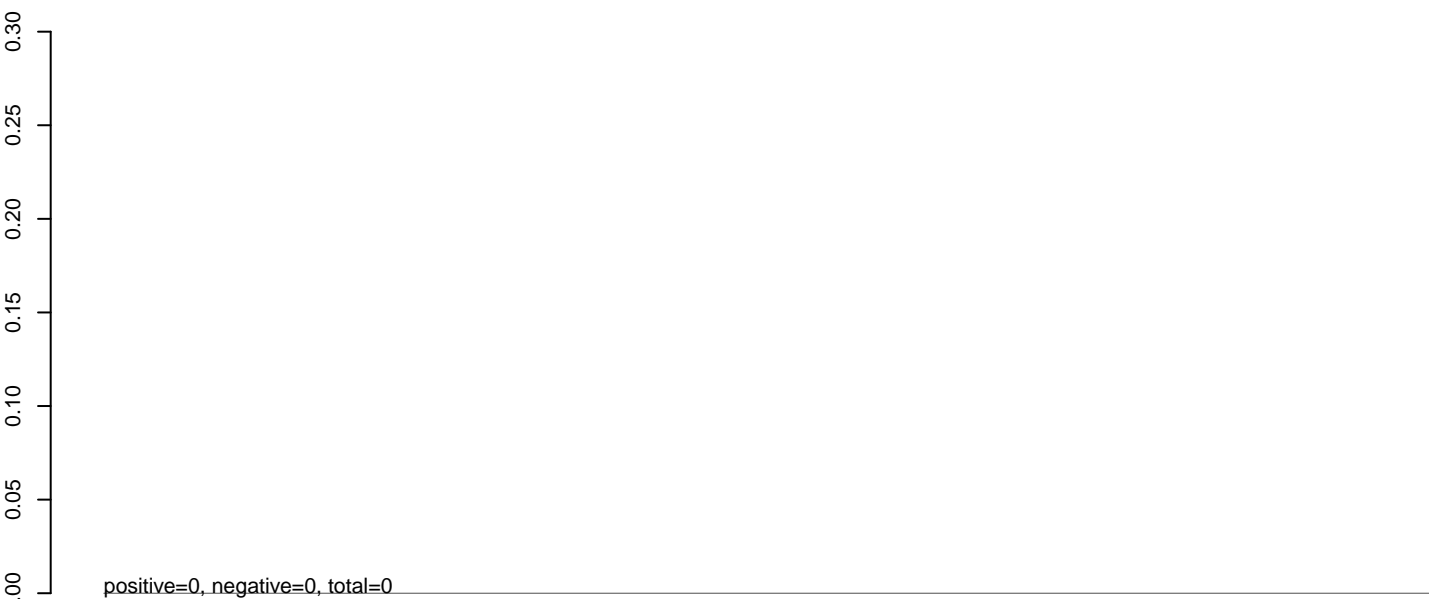
ambee... protein_matrix_hypothetical_proteins_glycoprotein_hypothetical_proteins_and_polymerase_genes:1-13593

0 2000 4000 6000 8000 10000 12000 14000

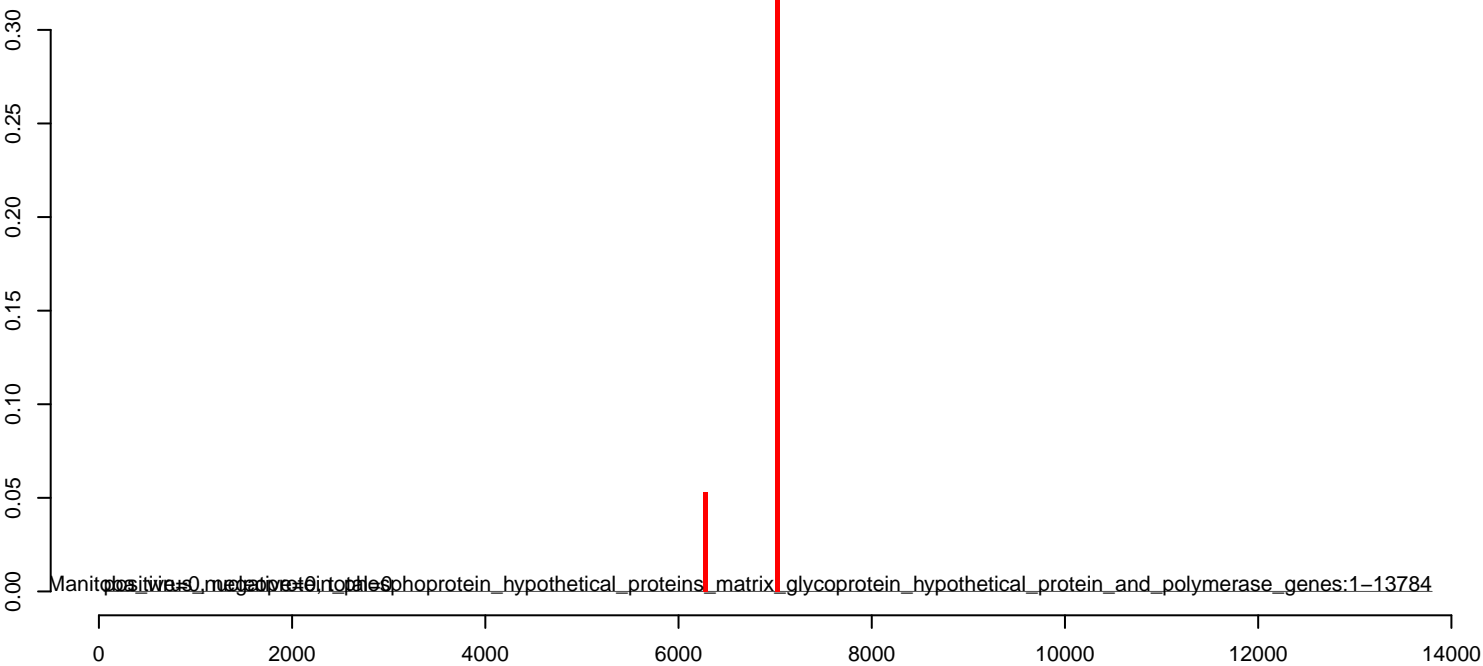
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



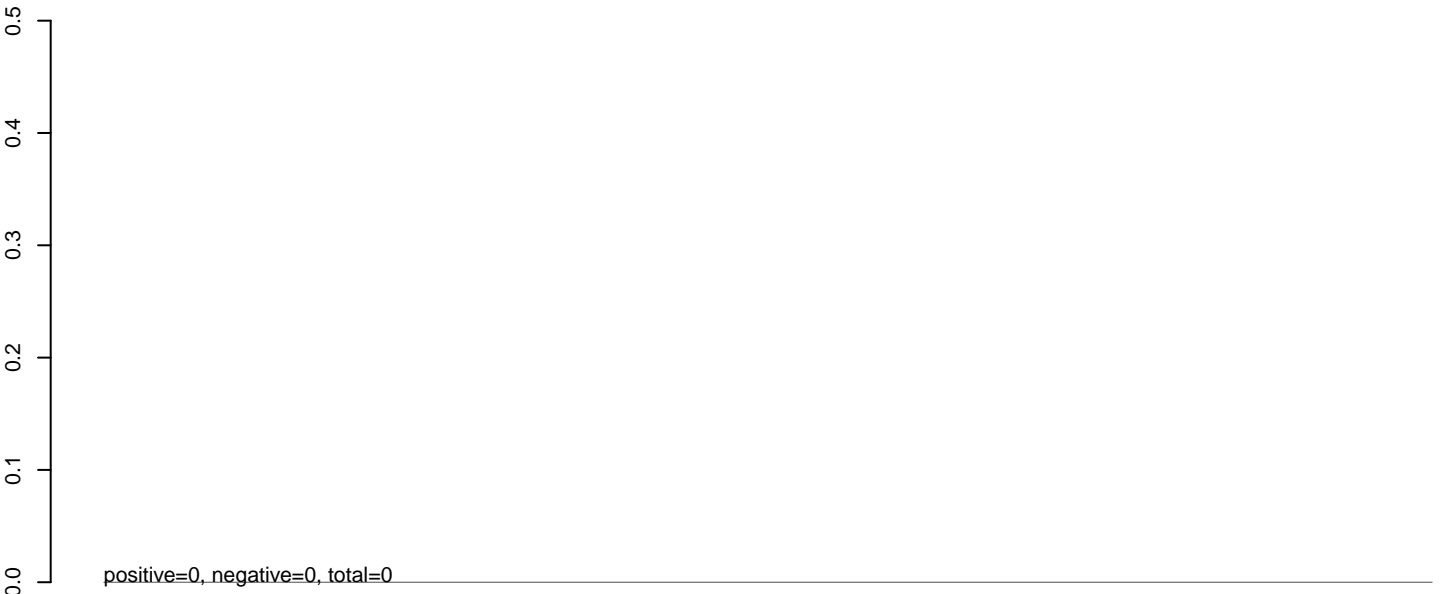
AeAeg_CCL.125cells_CHIKV.rep



AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep



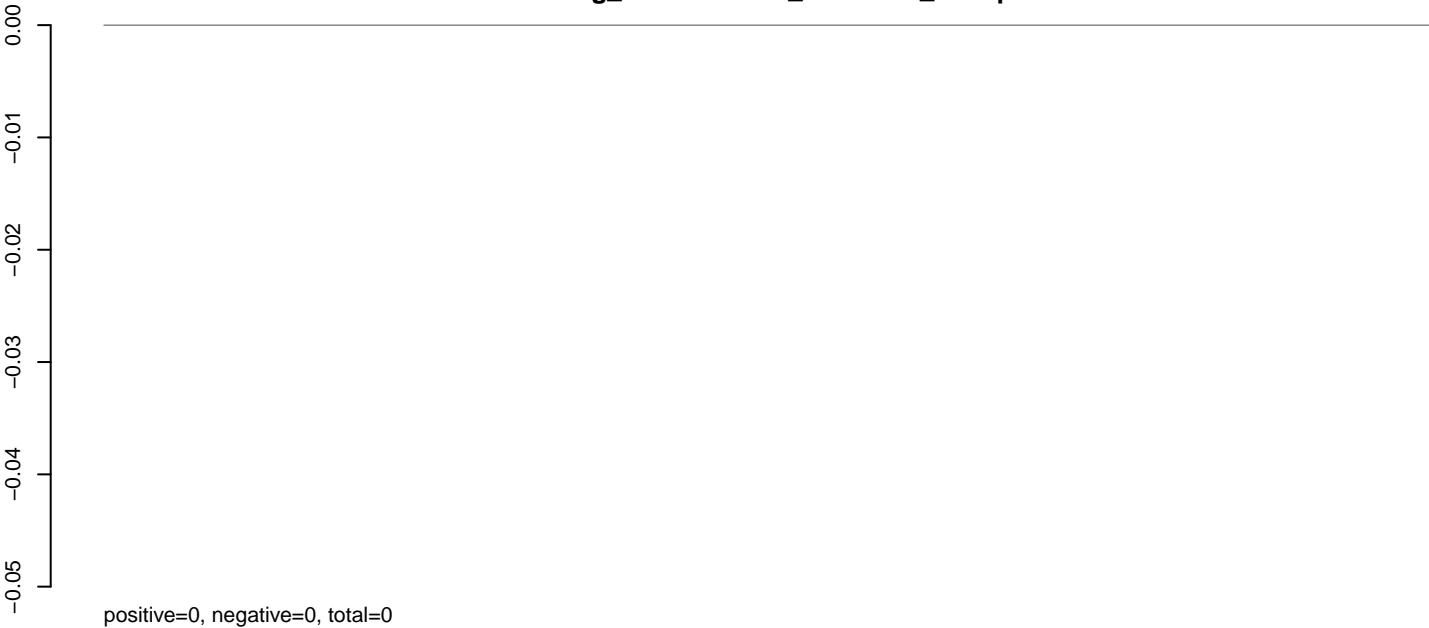
0 2000 4000 6000 8000 10000 12000

art_positives1_negaives0_totales phosphoprotein_hypothetical_proteins_matrix_glycoprotein_hypothetical_protein_and_polymerase_genes:1-13104

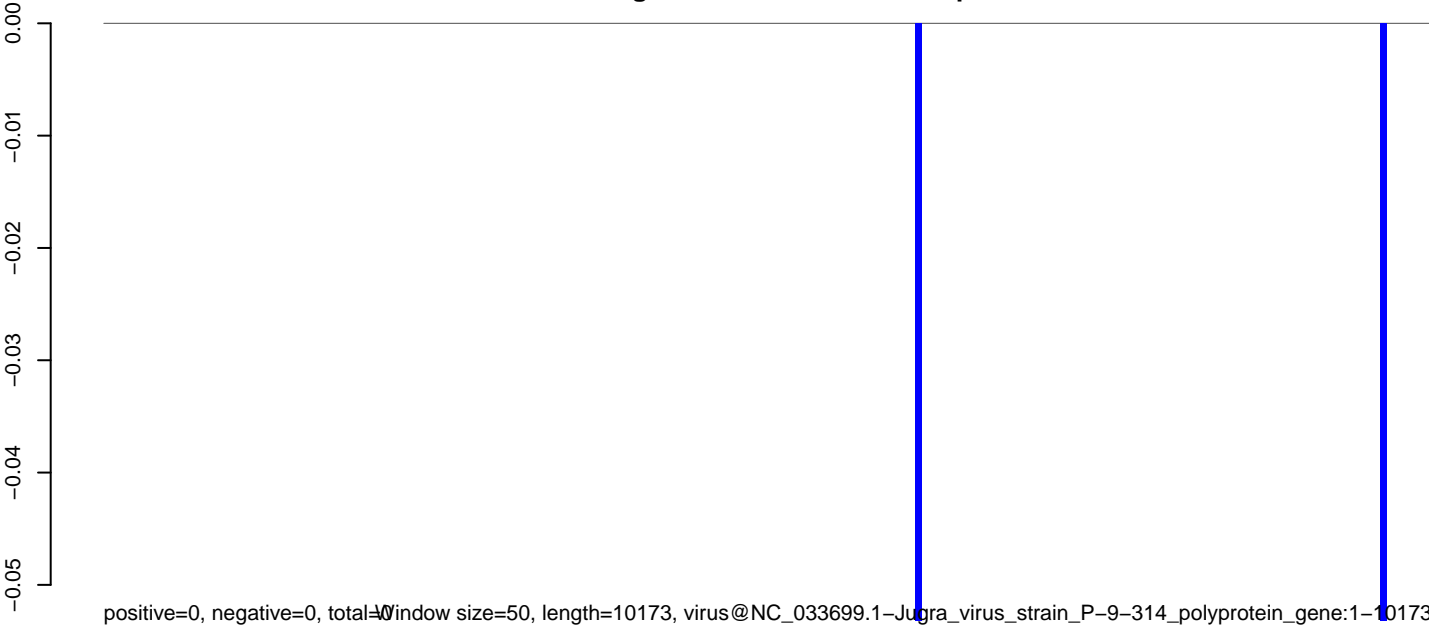
AeAeg_CCL.125cells_CHIKV.18_23.rep



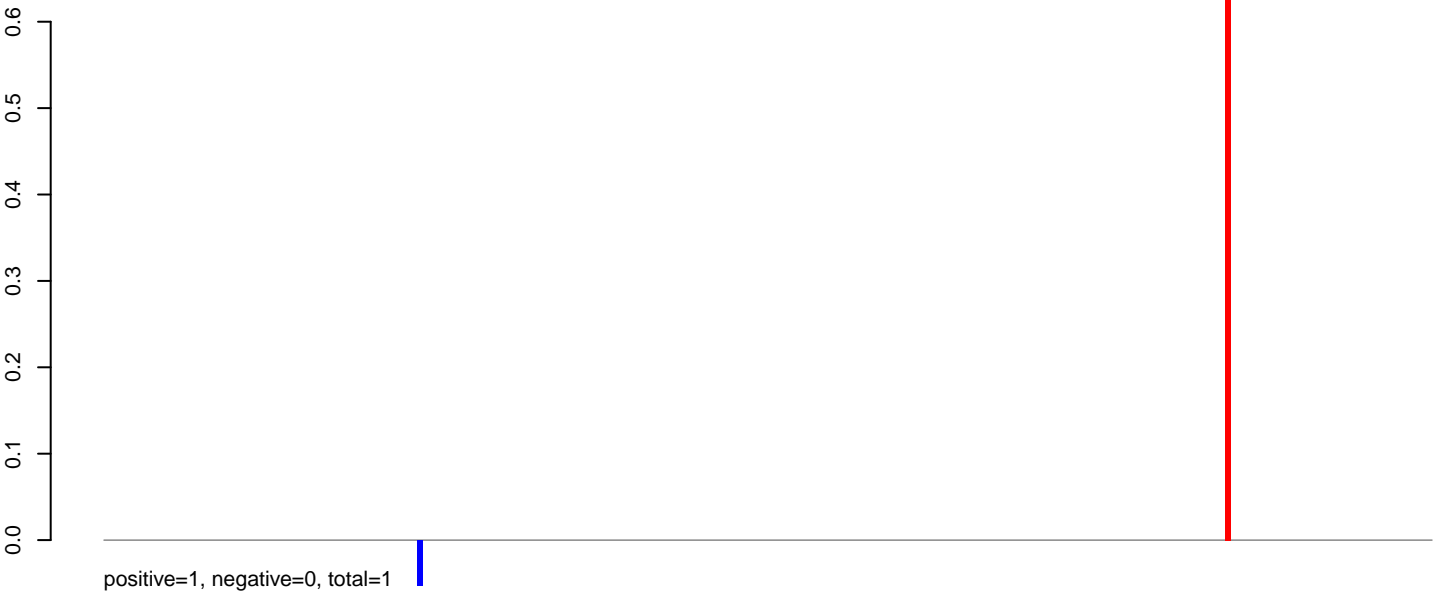
AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep



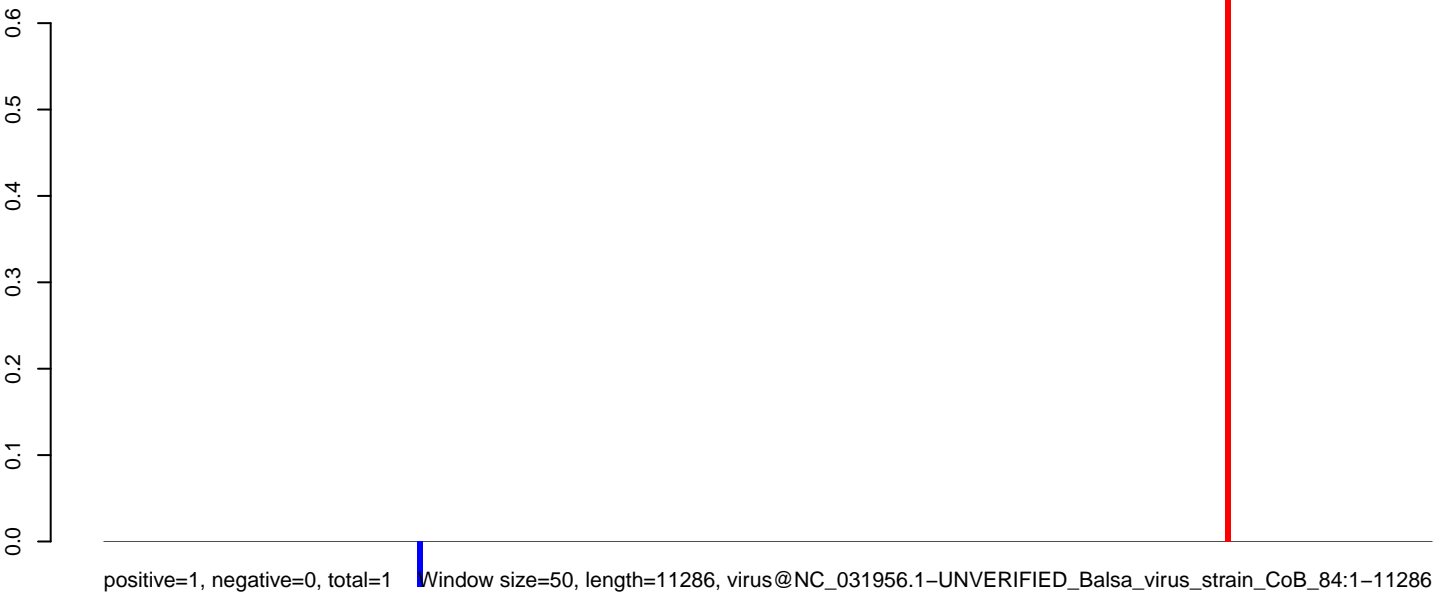
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



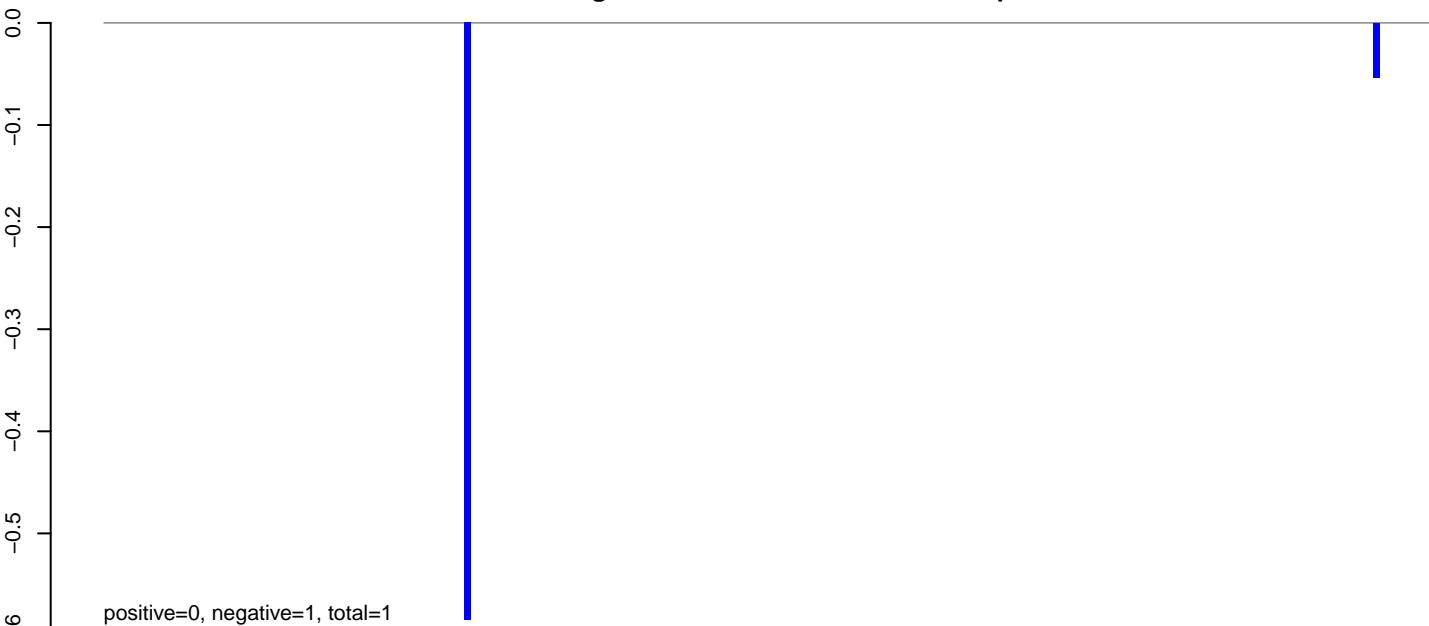
AeAeg_CCL.125cells_CHIKV.rep



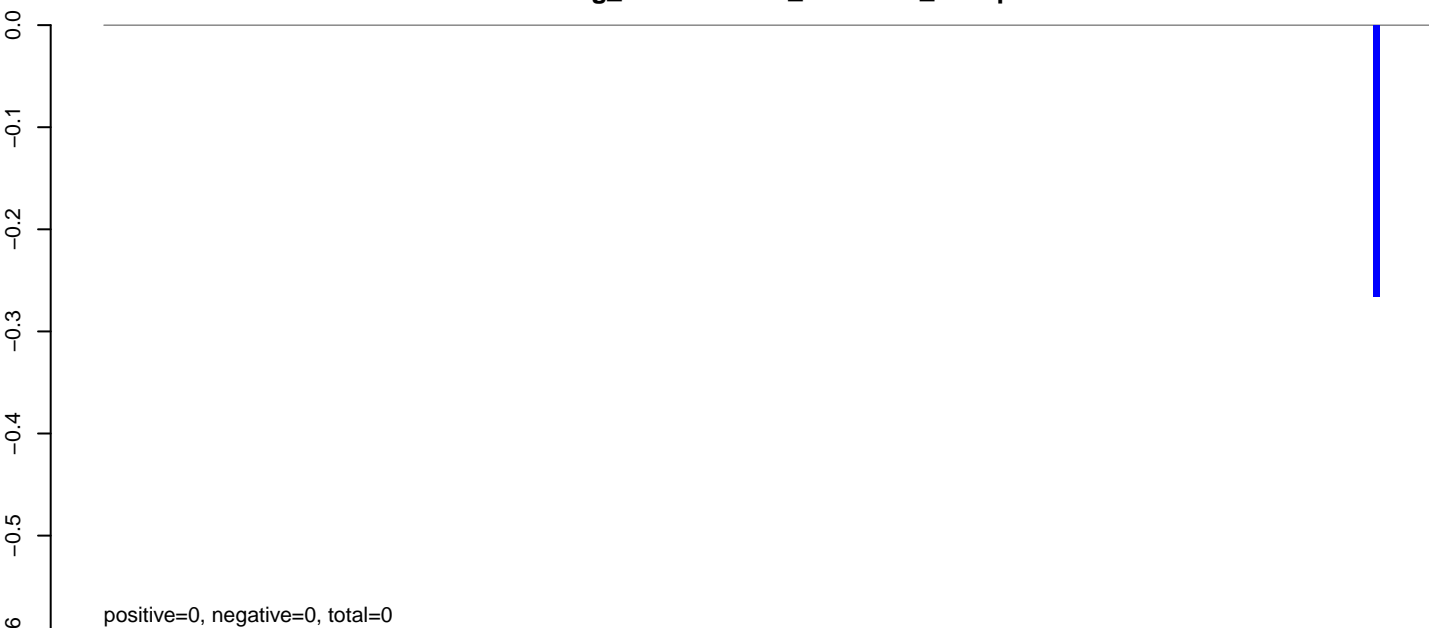
Window size=50, length=11286, virus@NC_031956.1-UNVERIFIED_Balsa_virus_strain_CoB_84:1-11286

0 2000 4000 6000 8000 10000

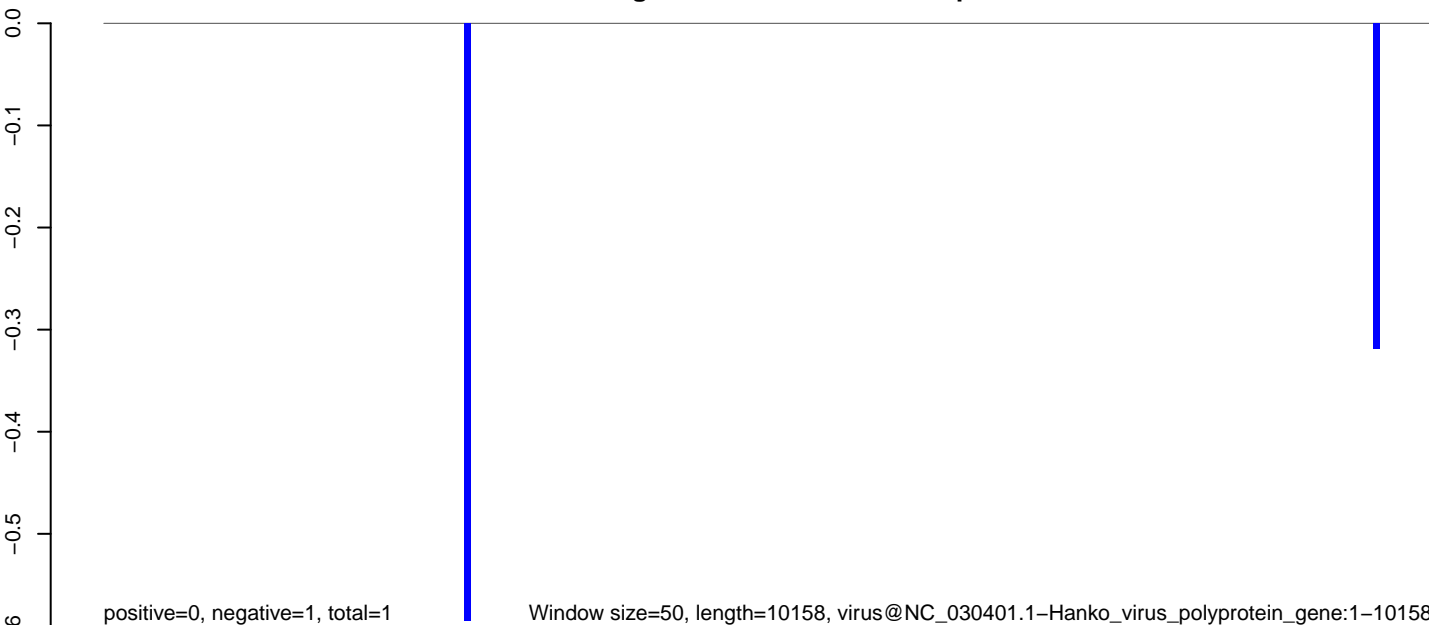
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



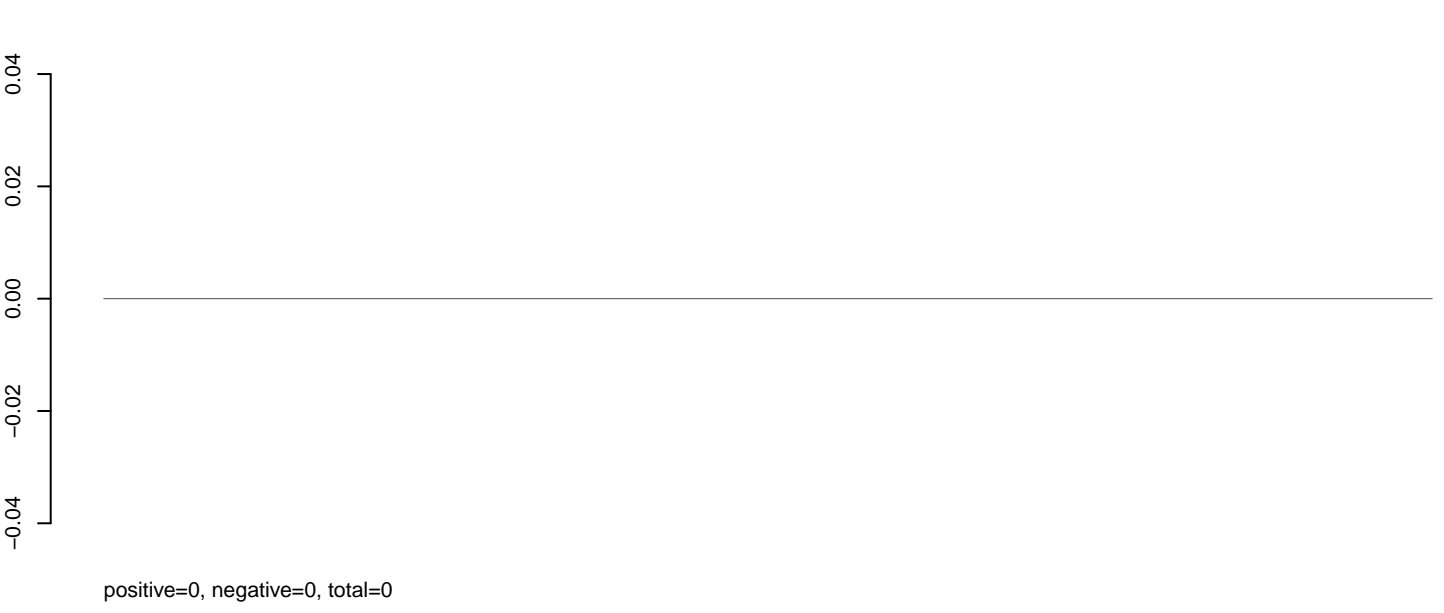
AeAeg_CCL.125cells_CHIKV.rep



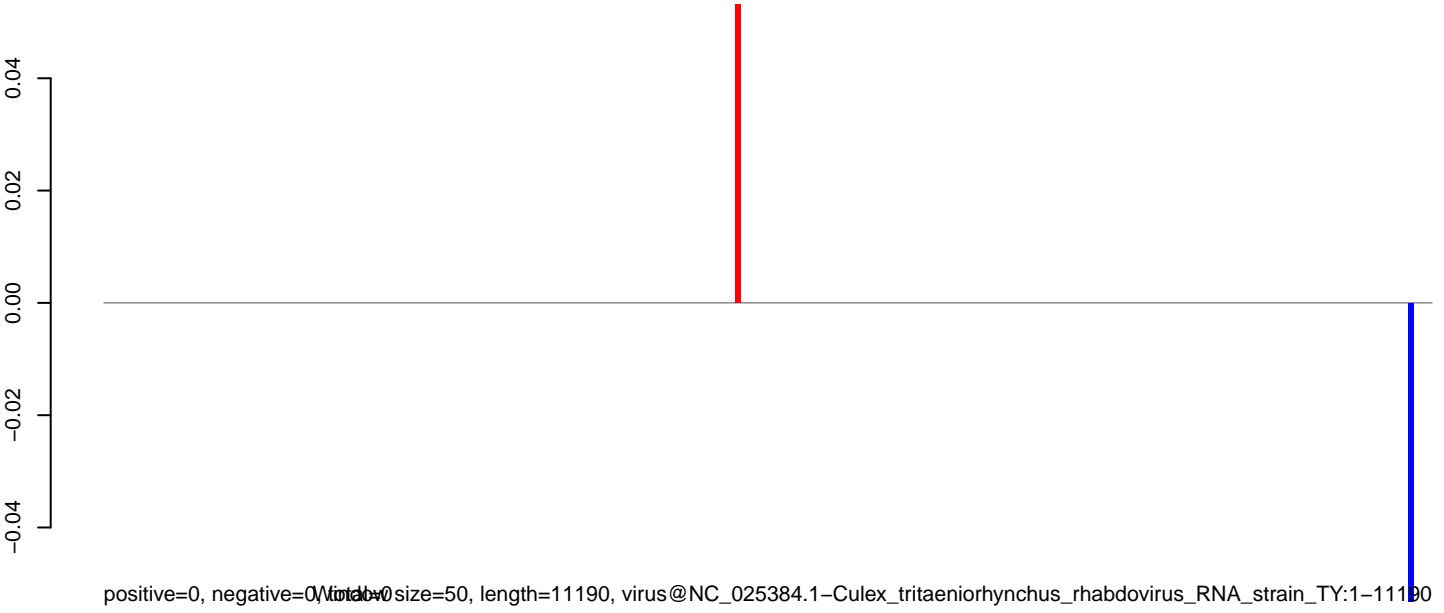
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep



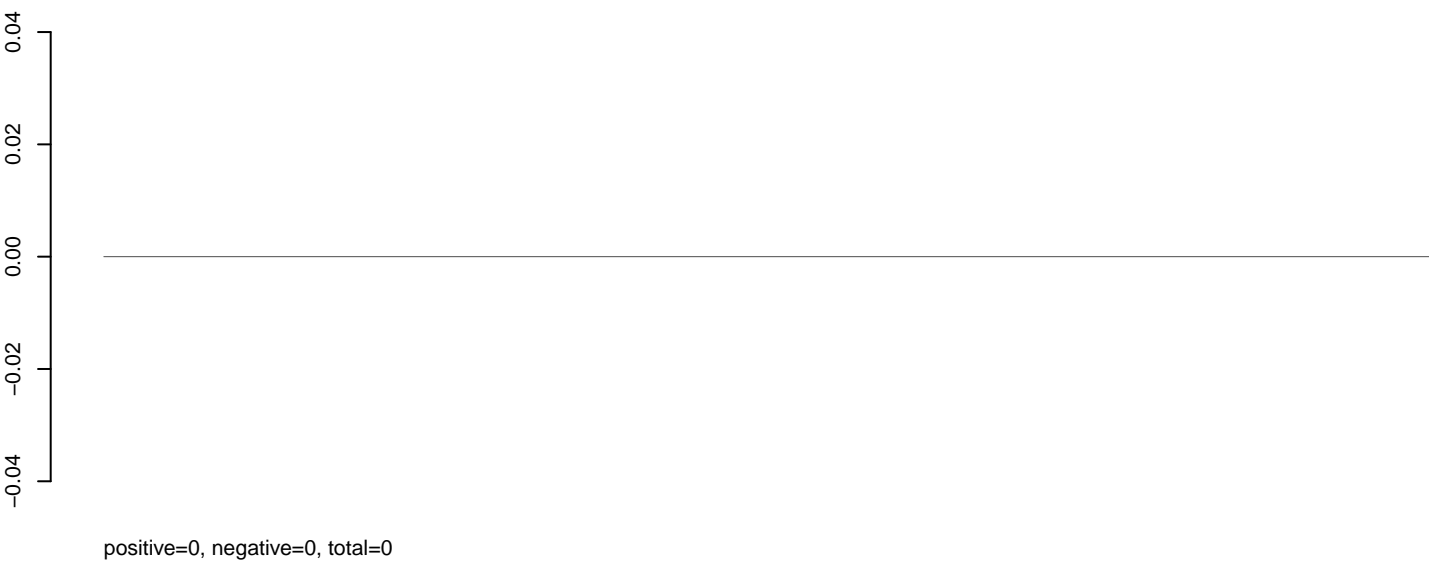
width=50, length=11190, virus@NC_025384.1-Culex tritaeniorhynchus rhabdovirus RNA strain TY:1-11190

0 2000 4000 6000 8000 10000

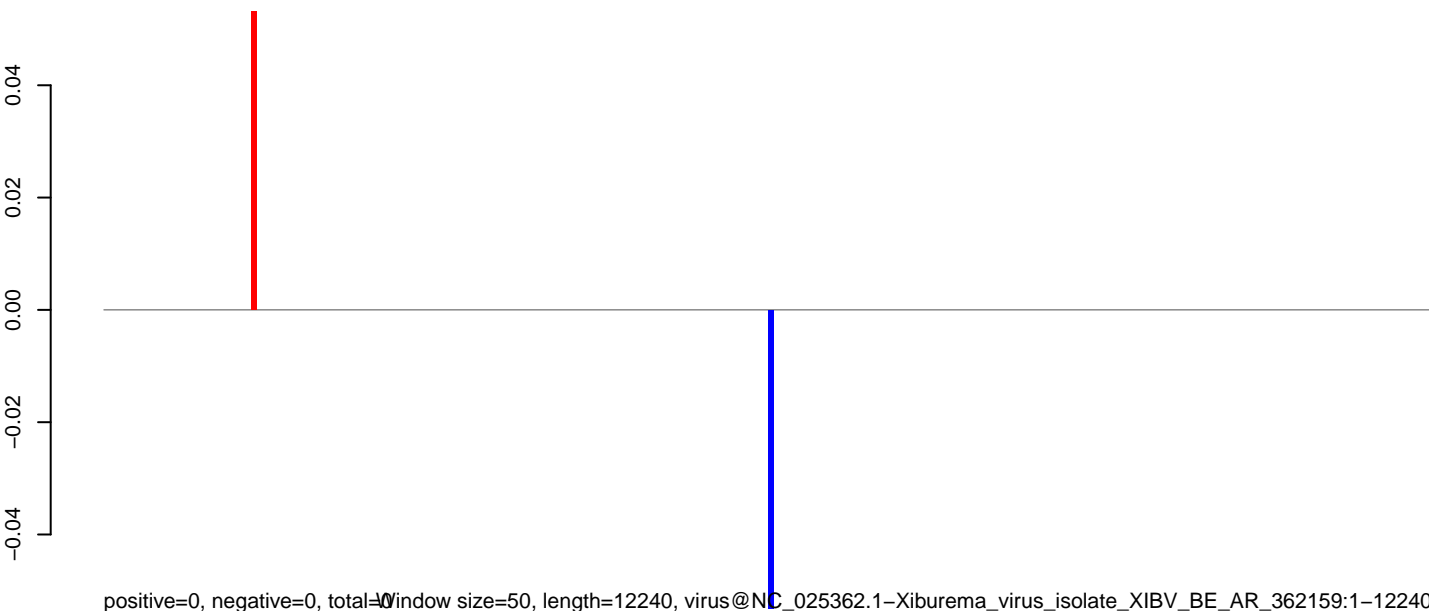
AeAeg_CCL.125cells_CHIKV.18_23.rep



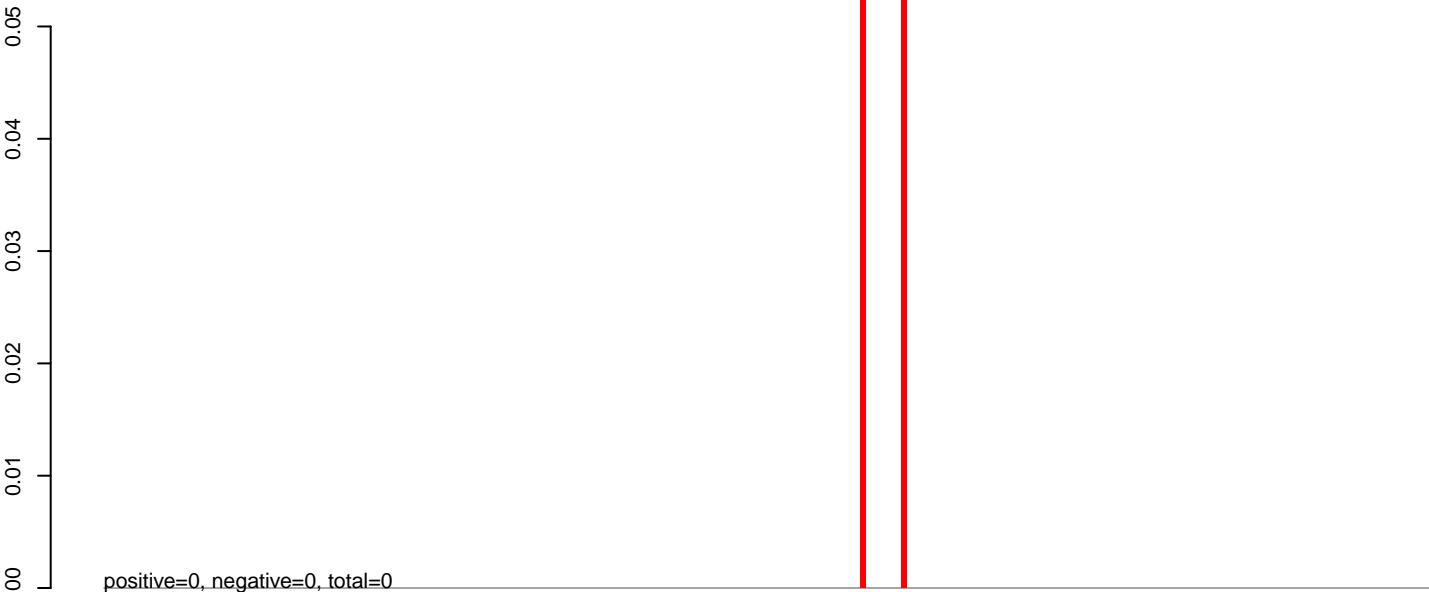
AeAeg_CCL.125cells_CHIKV.24_35.rep



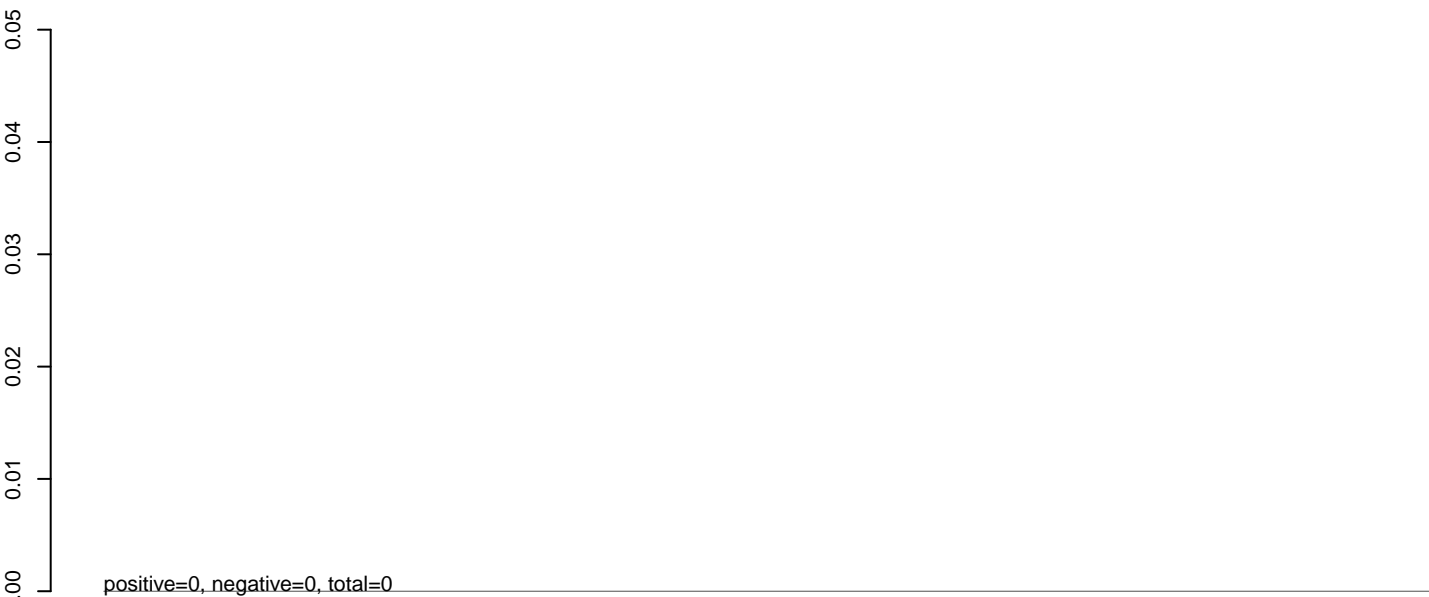
AeAeg_CCL.125cells_CHIKV.rep



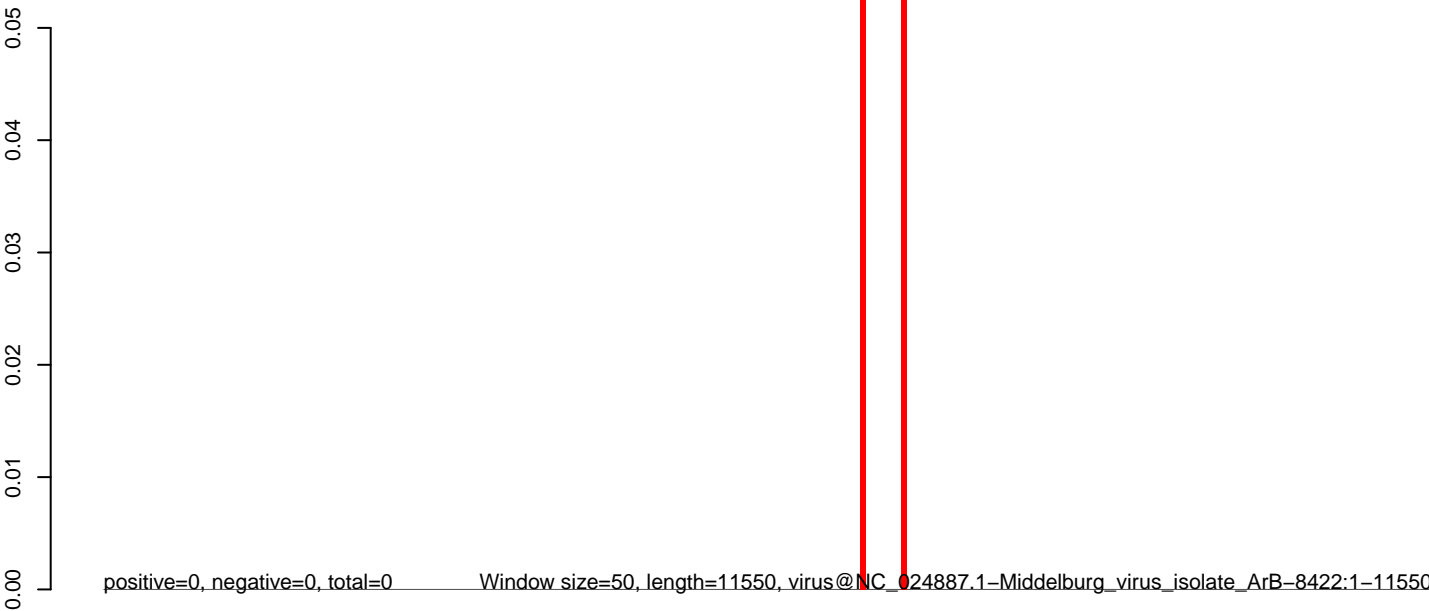
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



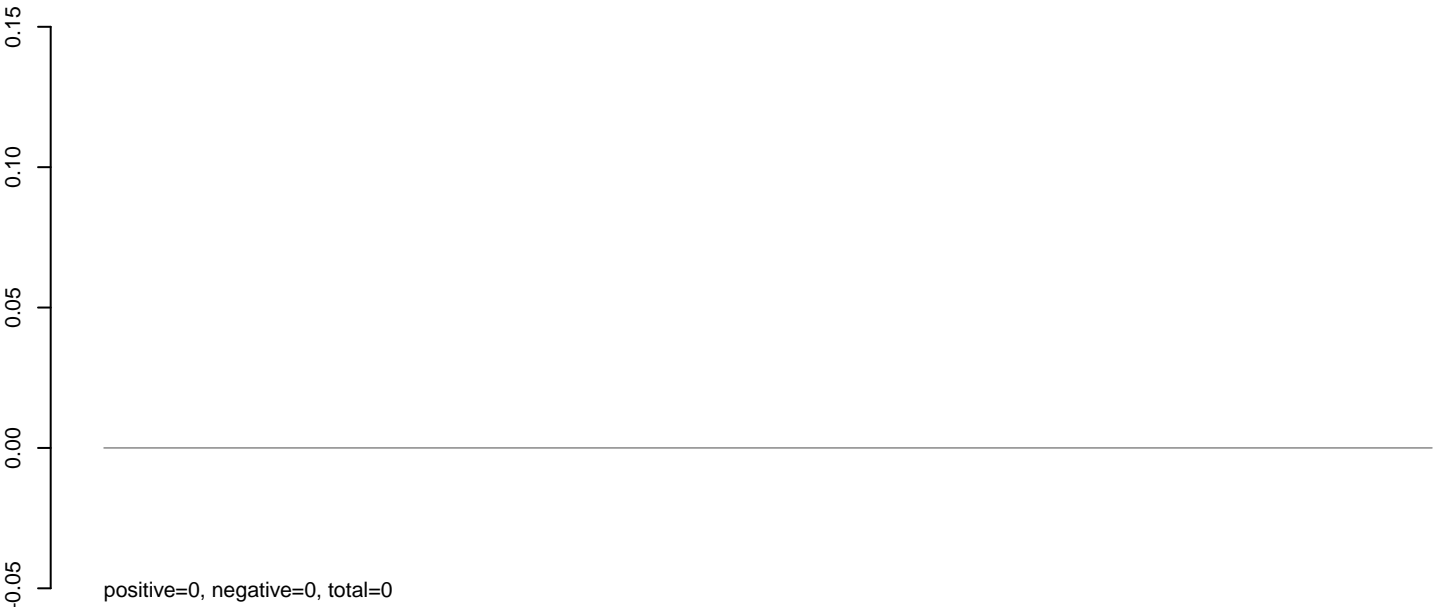
AeAeg_CCL.125cells_CHIKV.rep



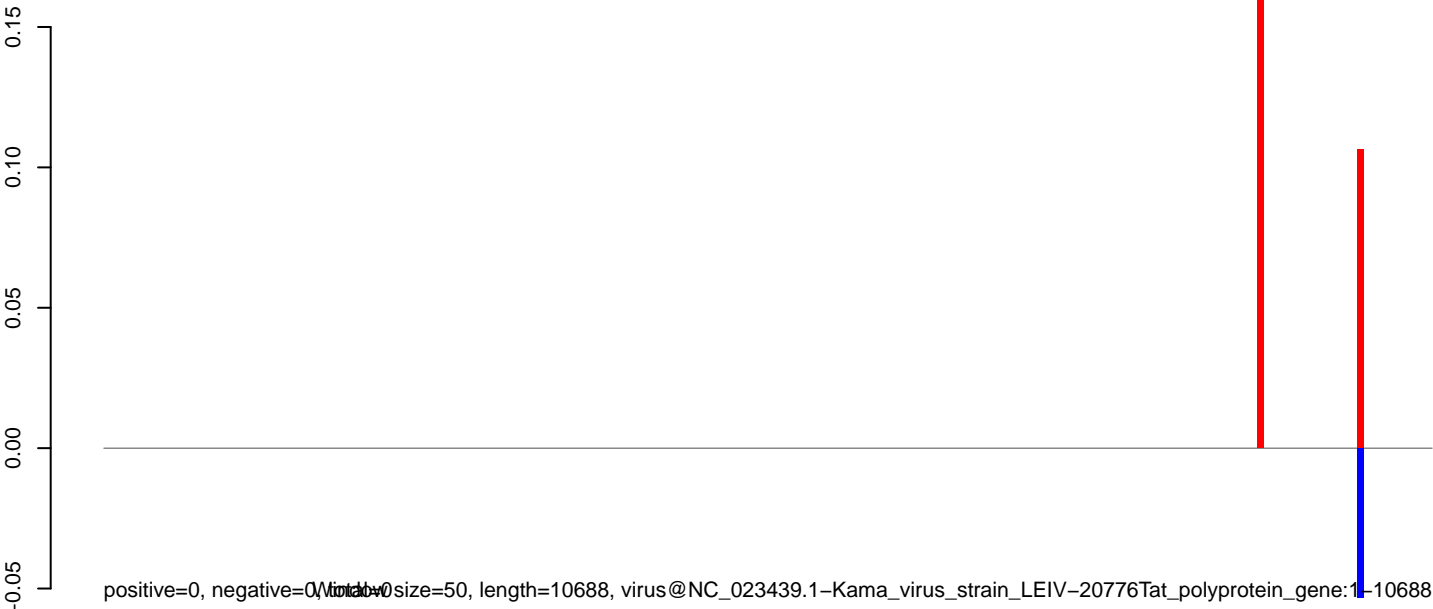
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

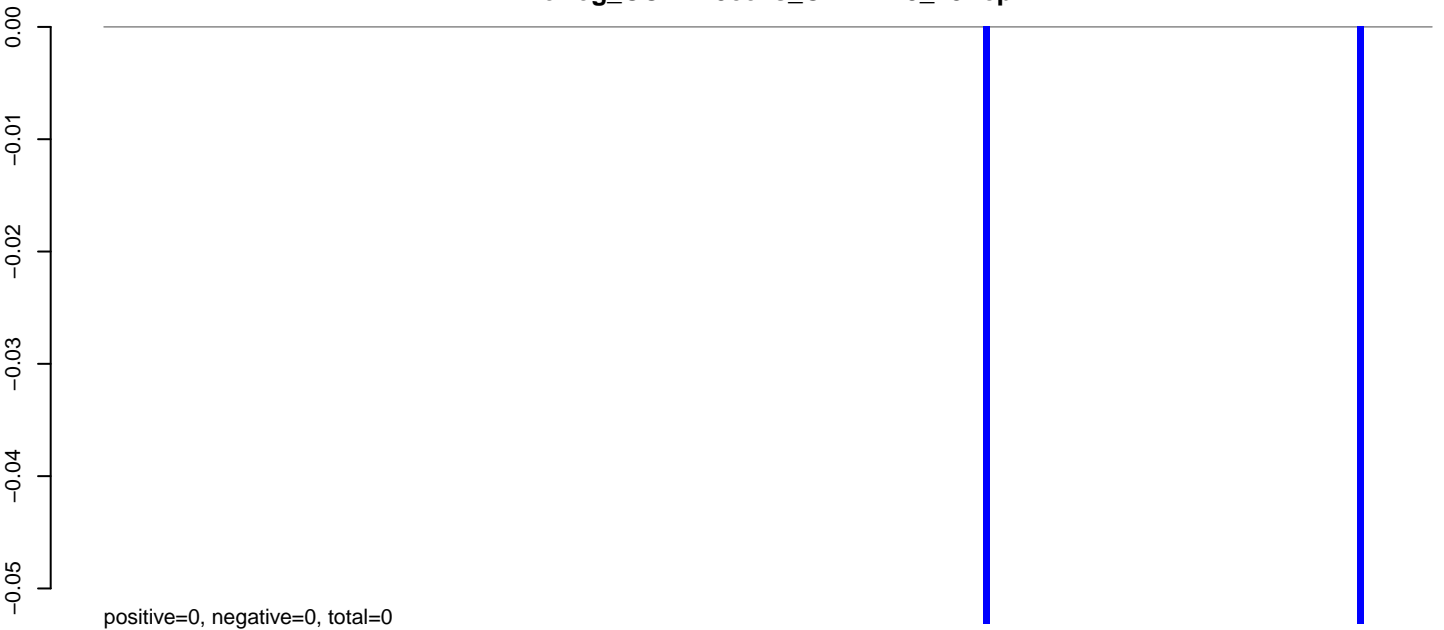


AeAeg_CCL.125cells_CHIKV.rep

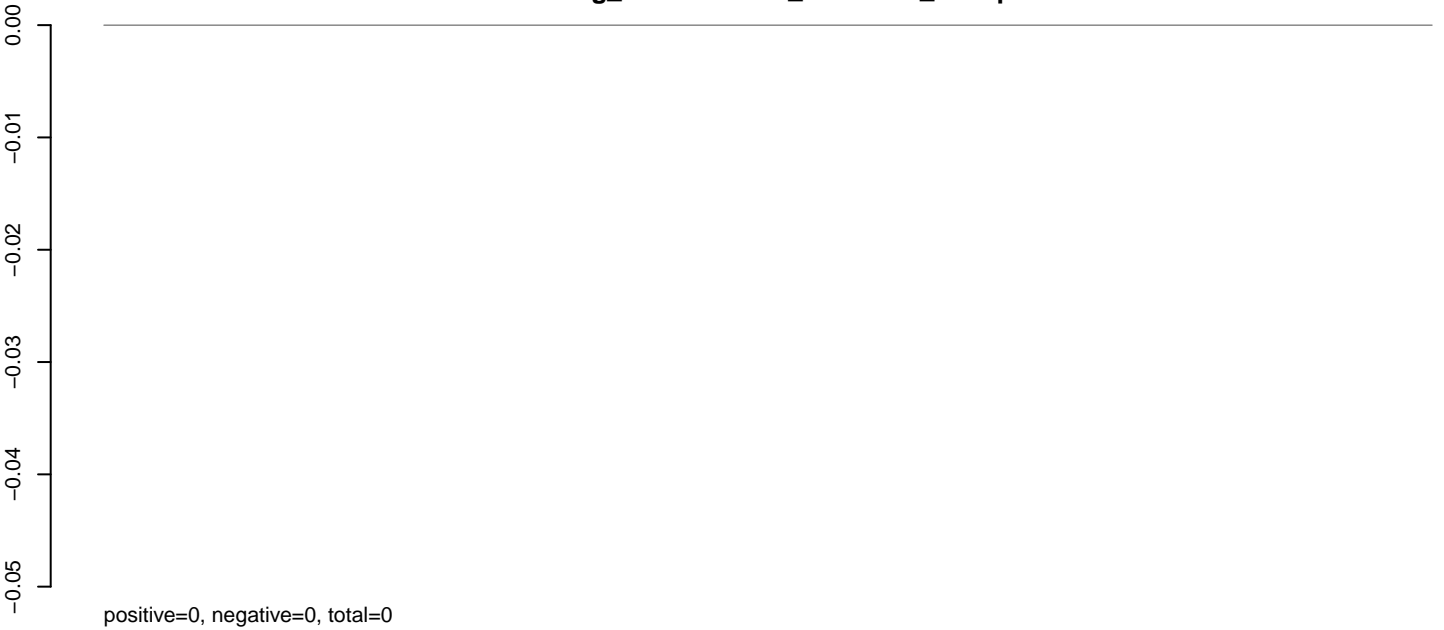


0 2000 4000 6000 8000 10000
positive=0, negative=0, total=0, window size=50, length=10688, virus@NC_023439.1-Kama_virus_strain_LEIV-20776Tat_polyprotein_gene:1-10688

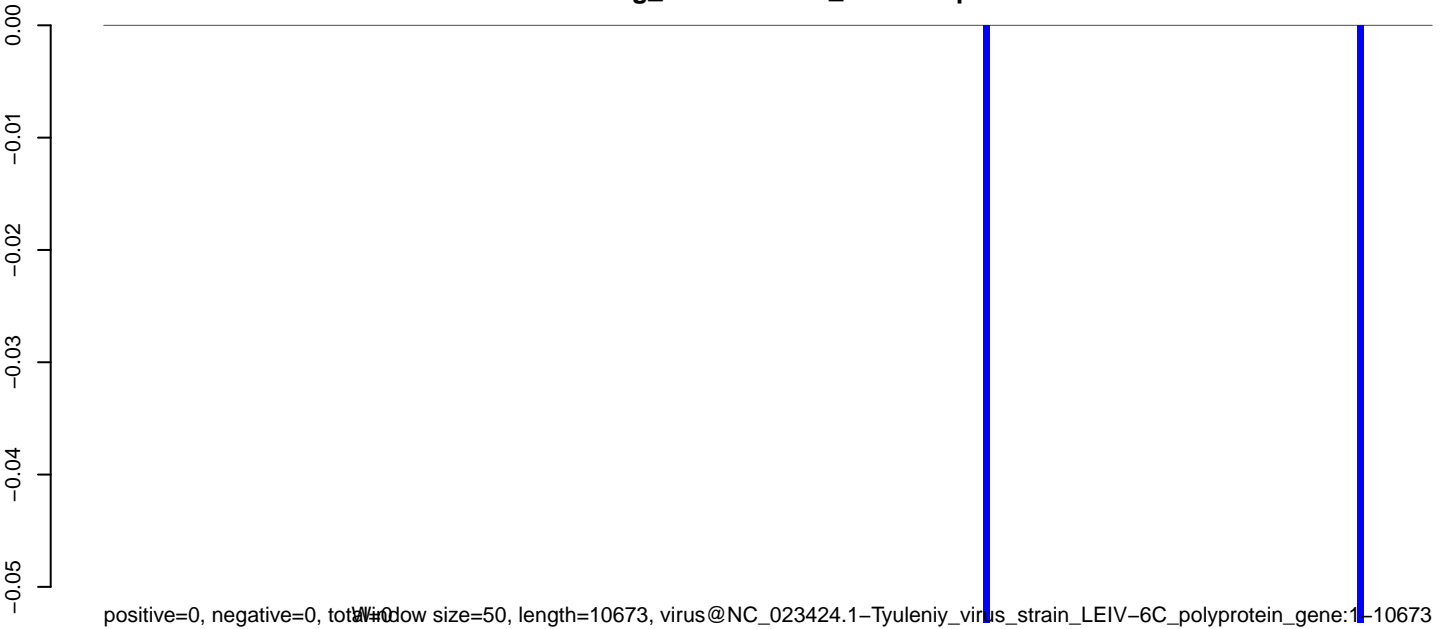
AeAeg_CCL.125cells_CHIKV.18_23.rep



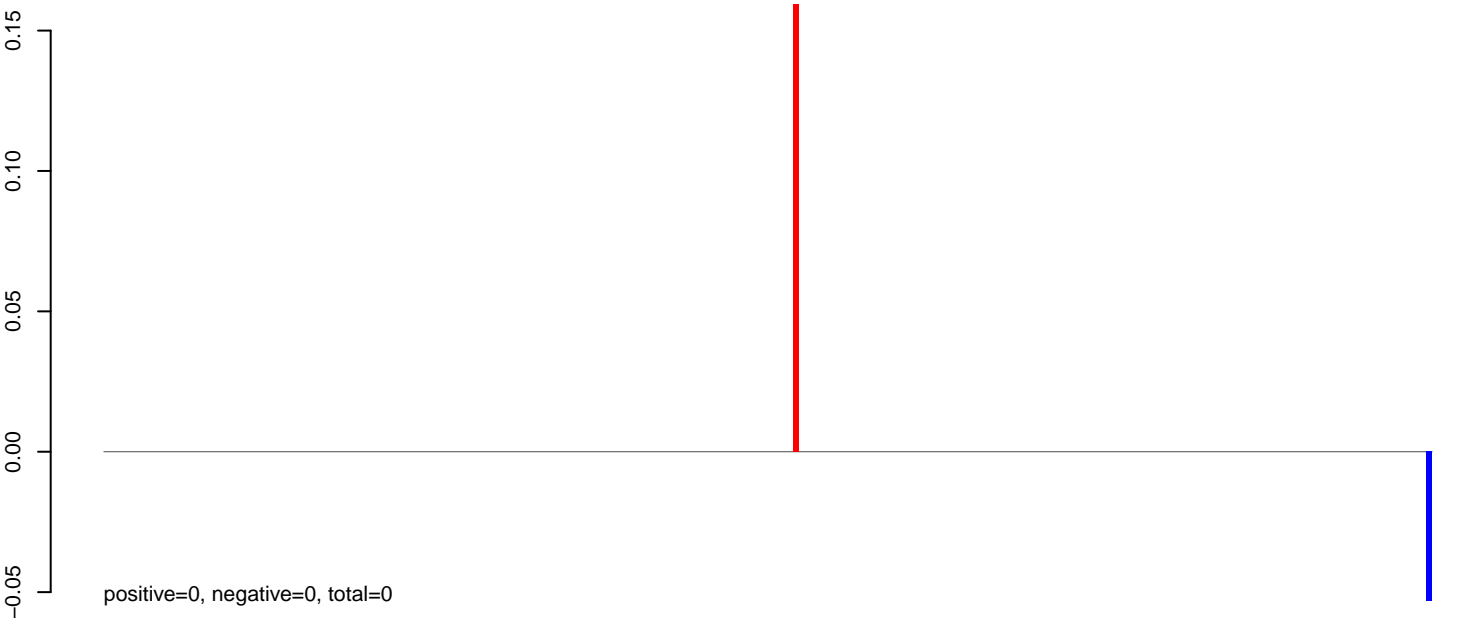
AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep



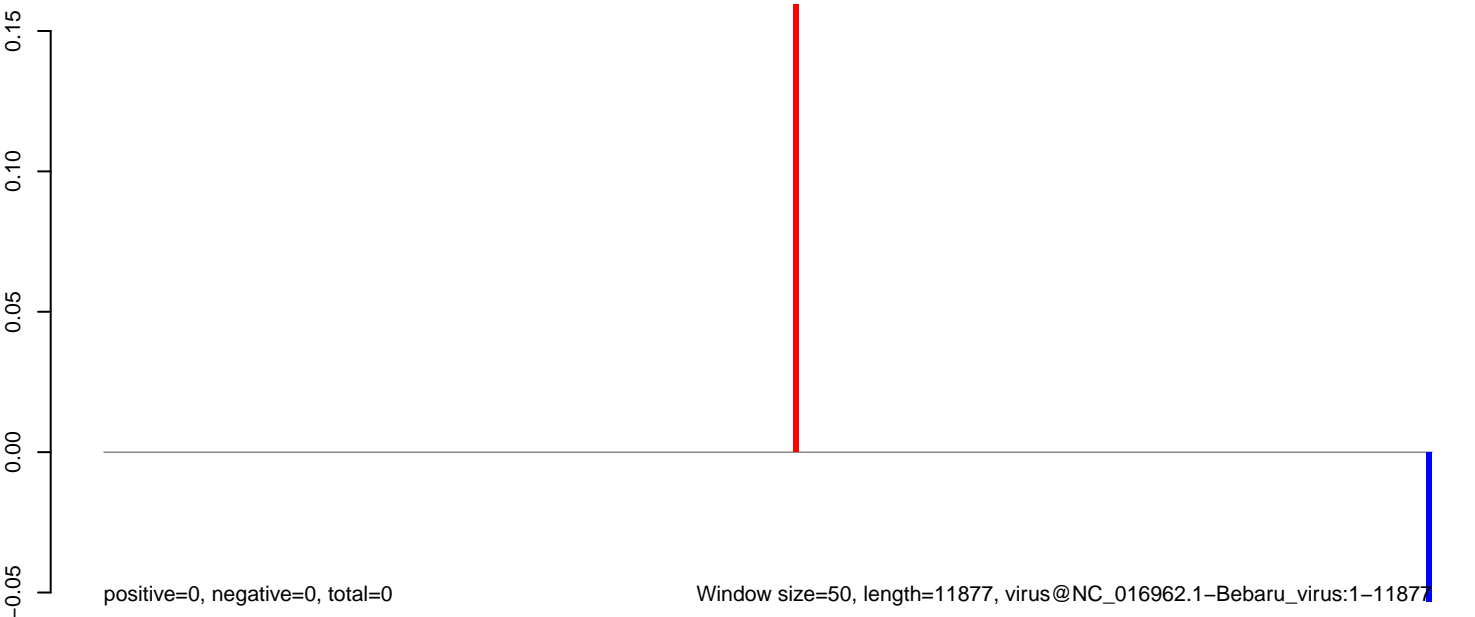
AeAeg_CCL.125cells_CHIKV.18_23.rep



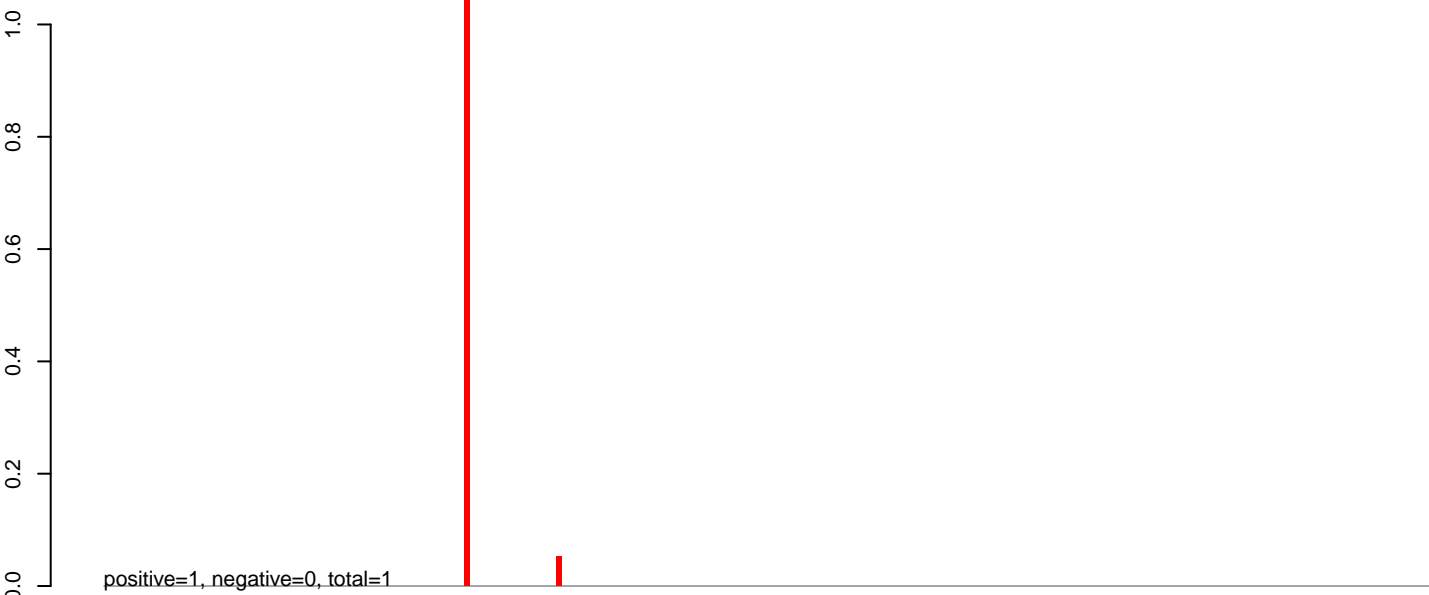
AeAeg_CCL.125cells_CHIKV.24_35.rep



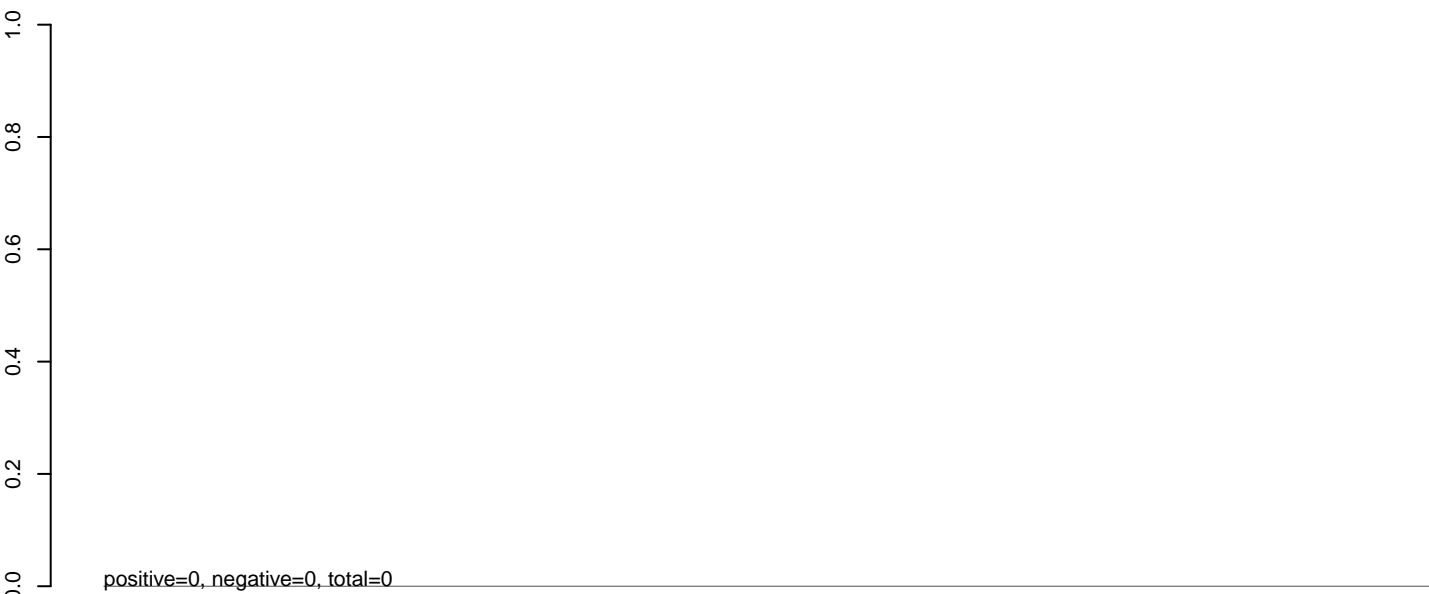
AeAeg_CCL.125cells_CHIKV.rep



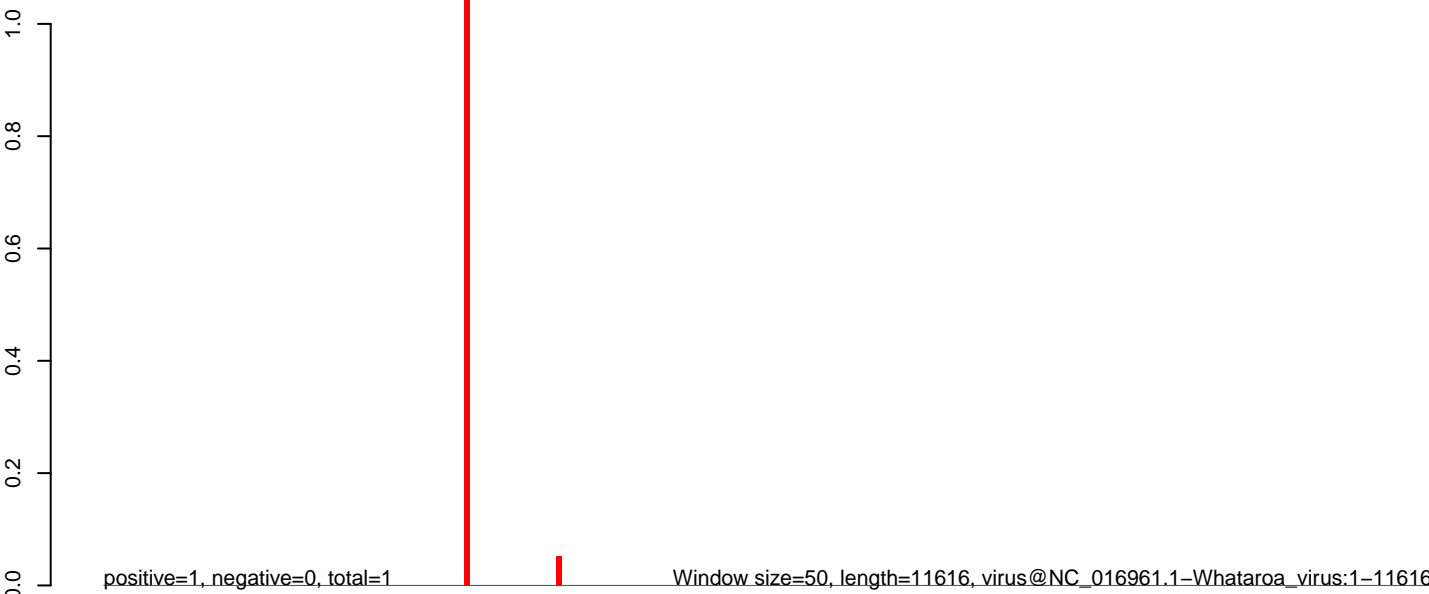
AeAeg_CCL.125cells_CHIKV.18_23.rep



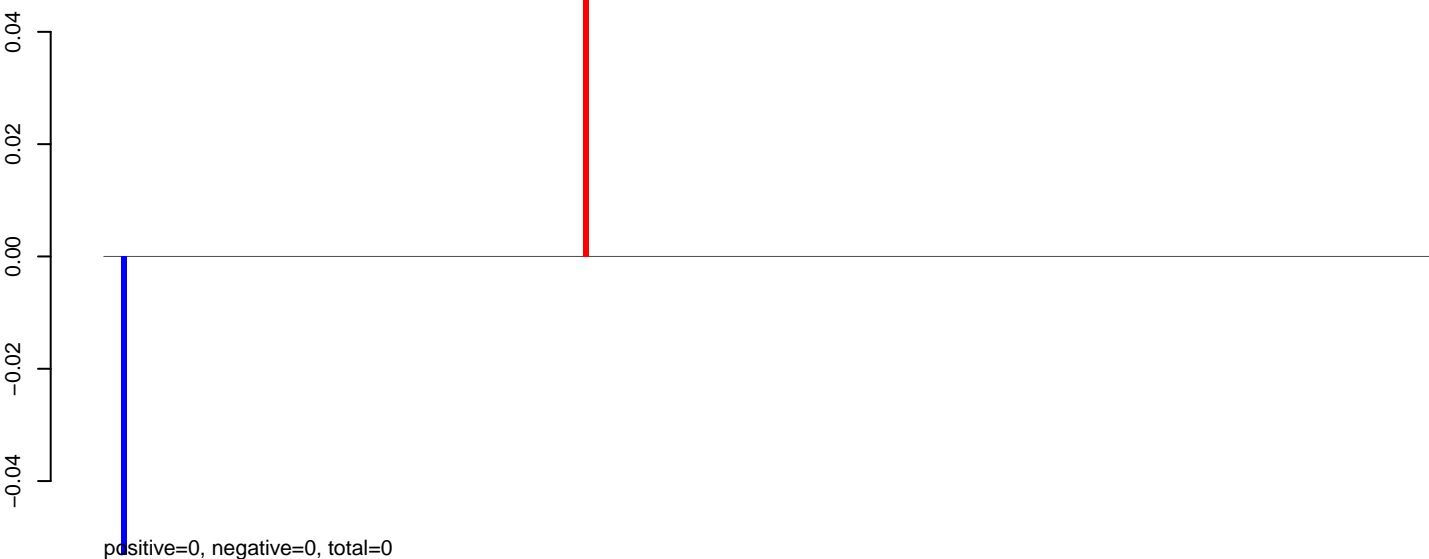
AeAeg_CCL.125cells_CHIKV.24_35.rep



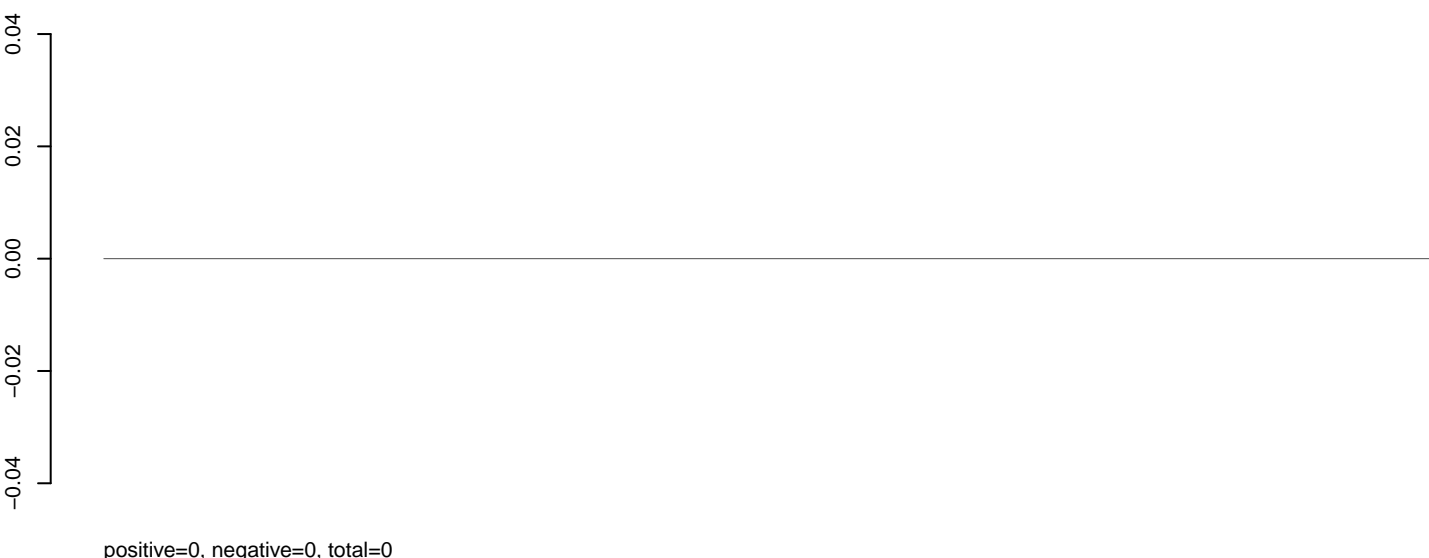
AeAeg_CCL.125cells_CHIKV.rep



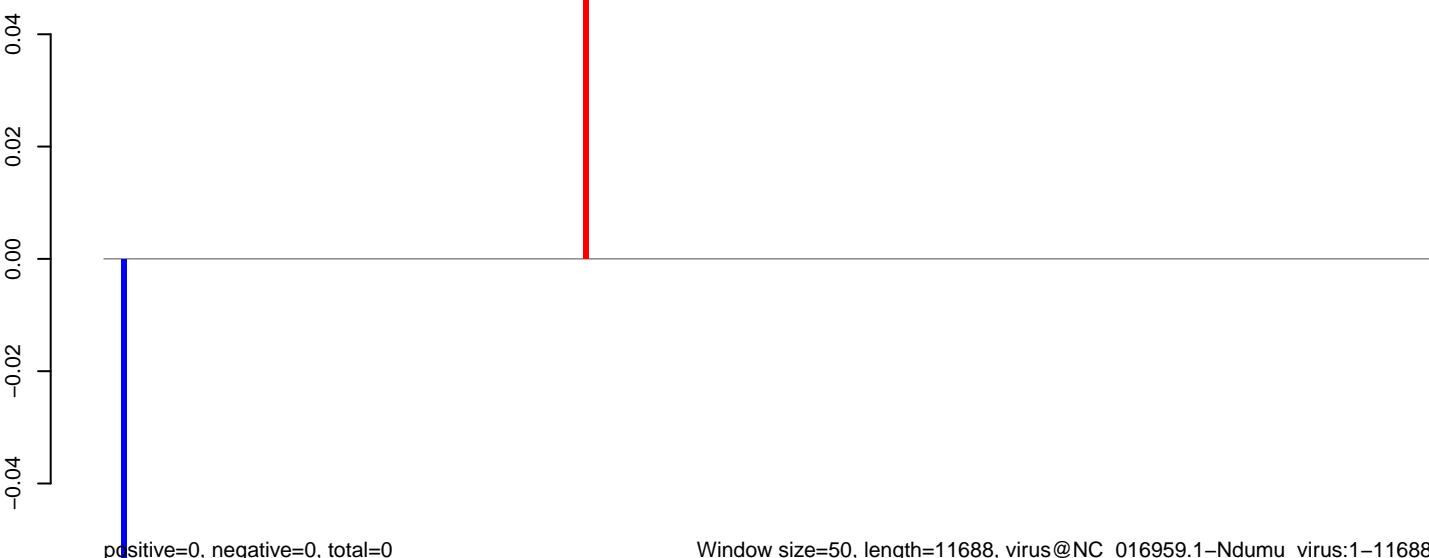
AeAeg_CCL.125cells_CHIKV.18_23.rep



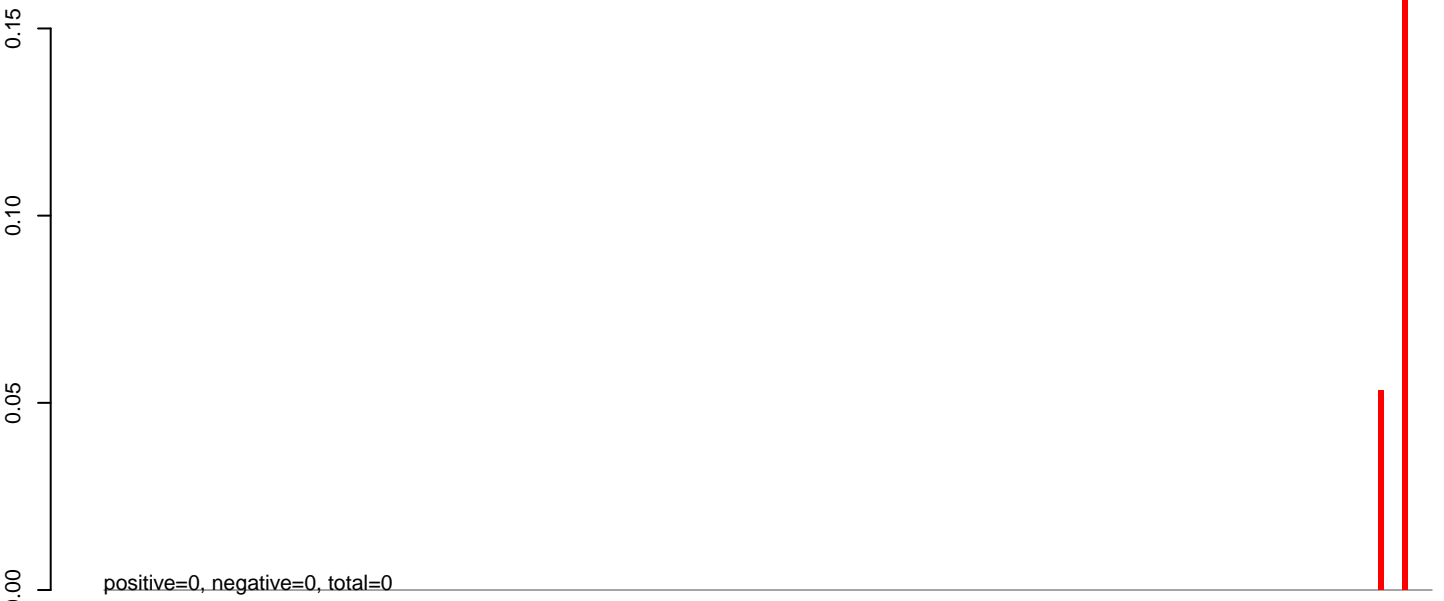
AeAeg_CCL.125cells_CHIKV.24_35.rep



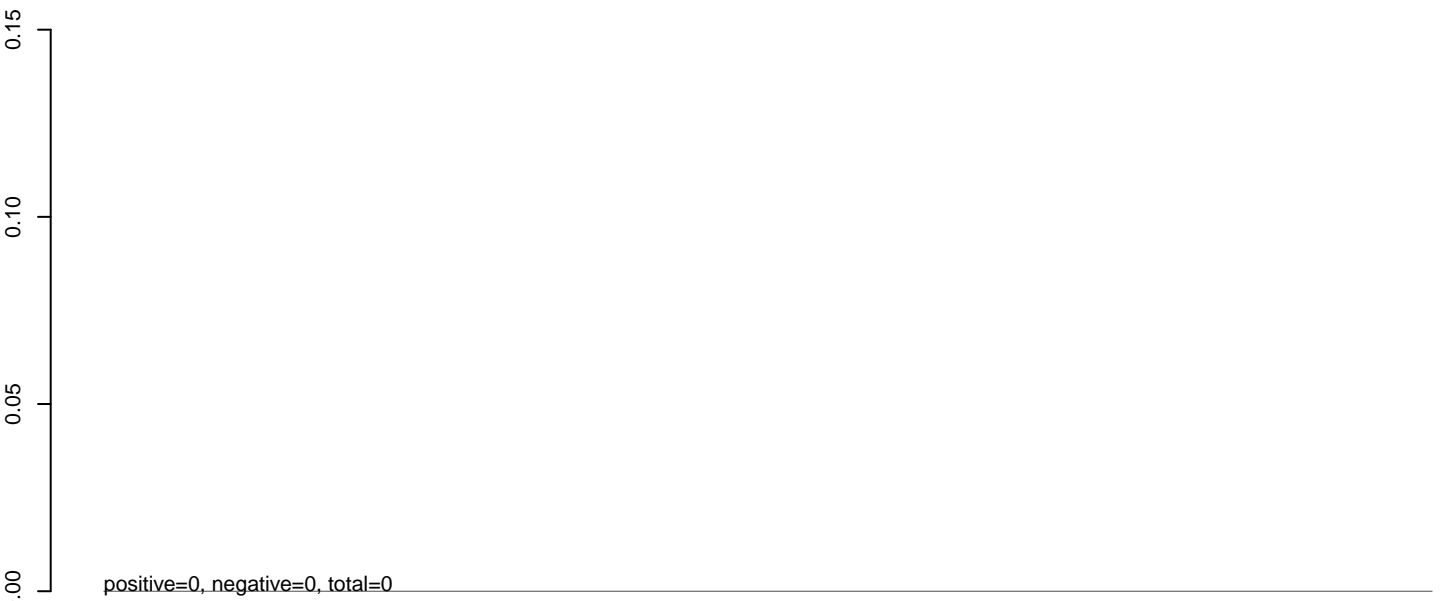
AeAeg_CCL.125cells_CHIKV.rep



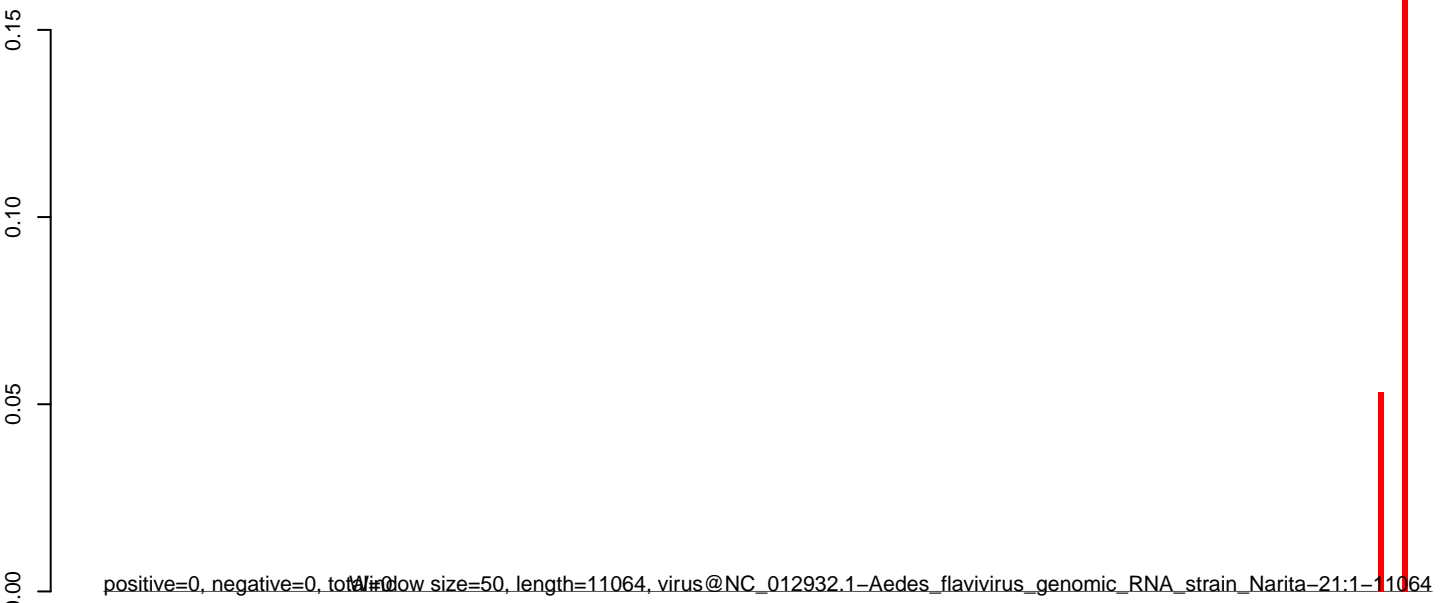
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

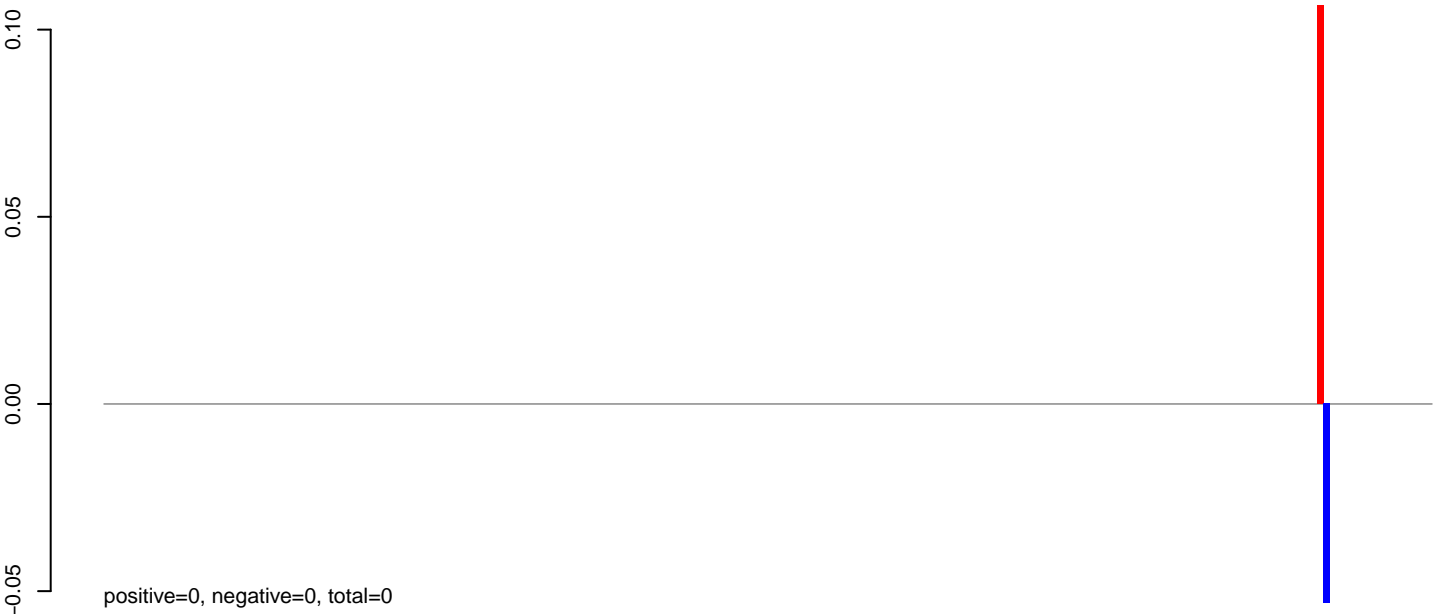


AeAeg_CCL.125cells_CHIKV.rep

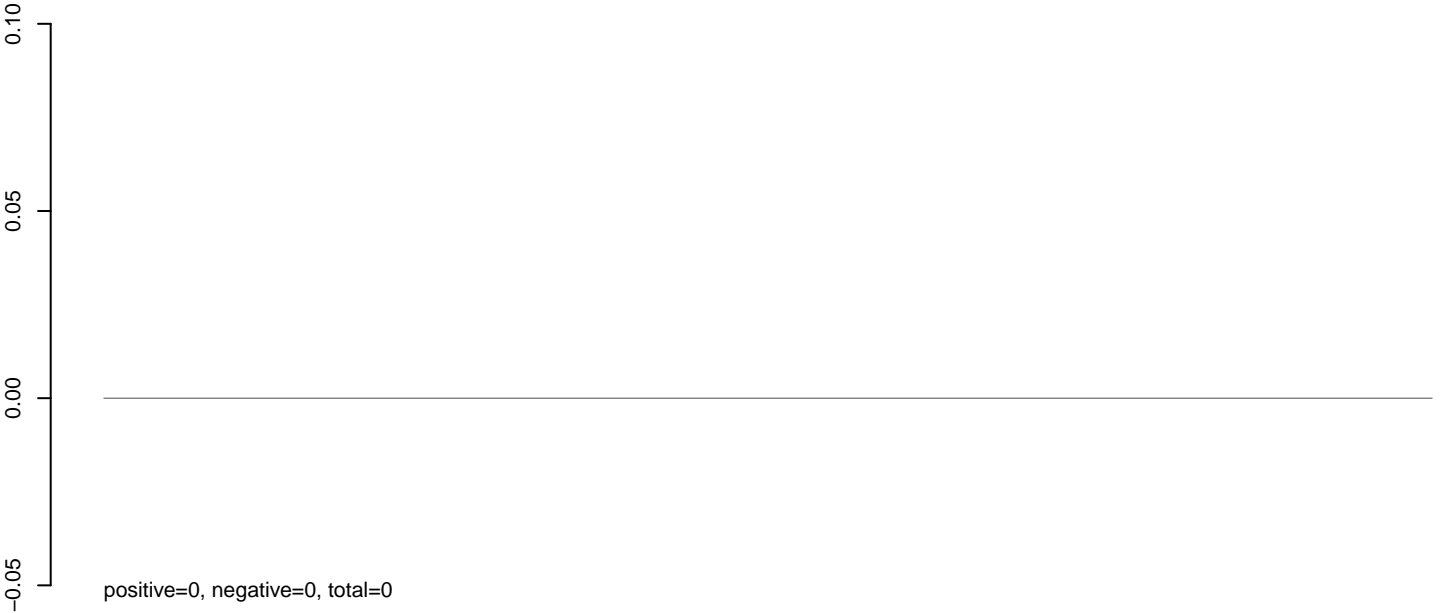


low size=50, length=11064, virus@NC_012932.1-Aedes_flavivirus_genomic_RNA_strain_Narita-21:1-11064

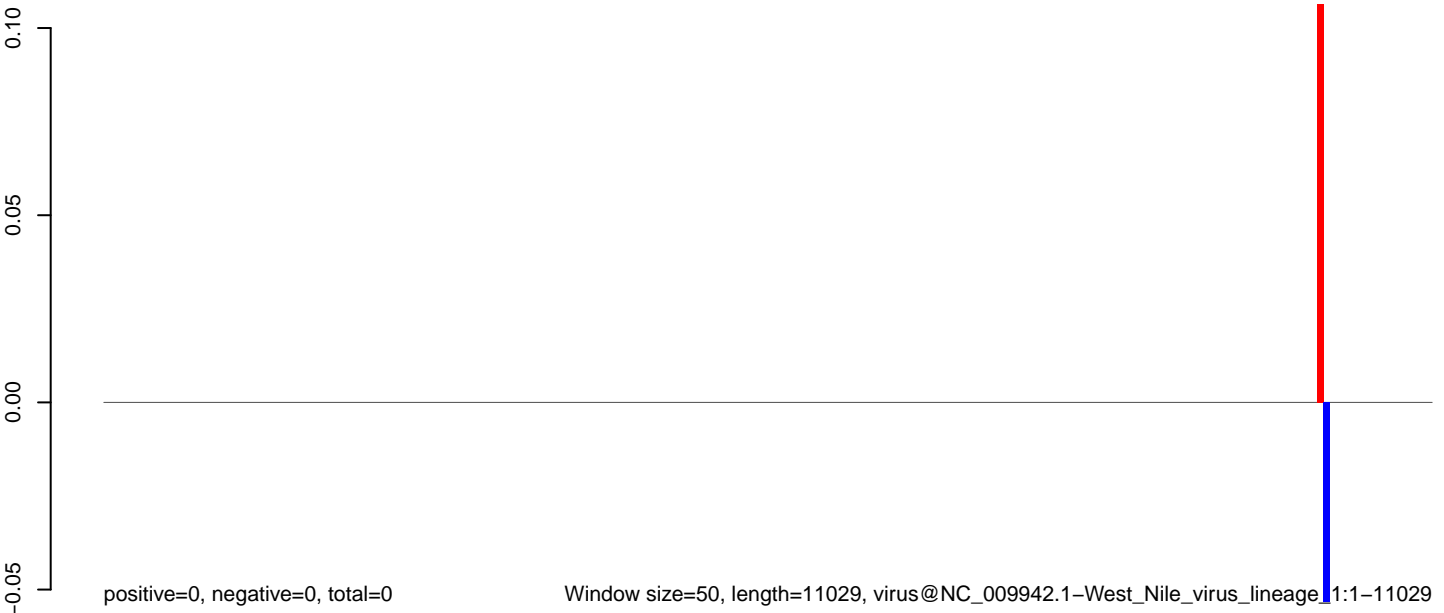
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

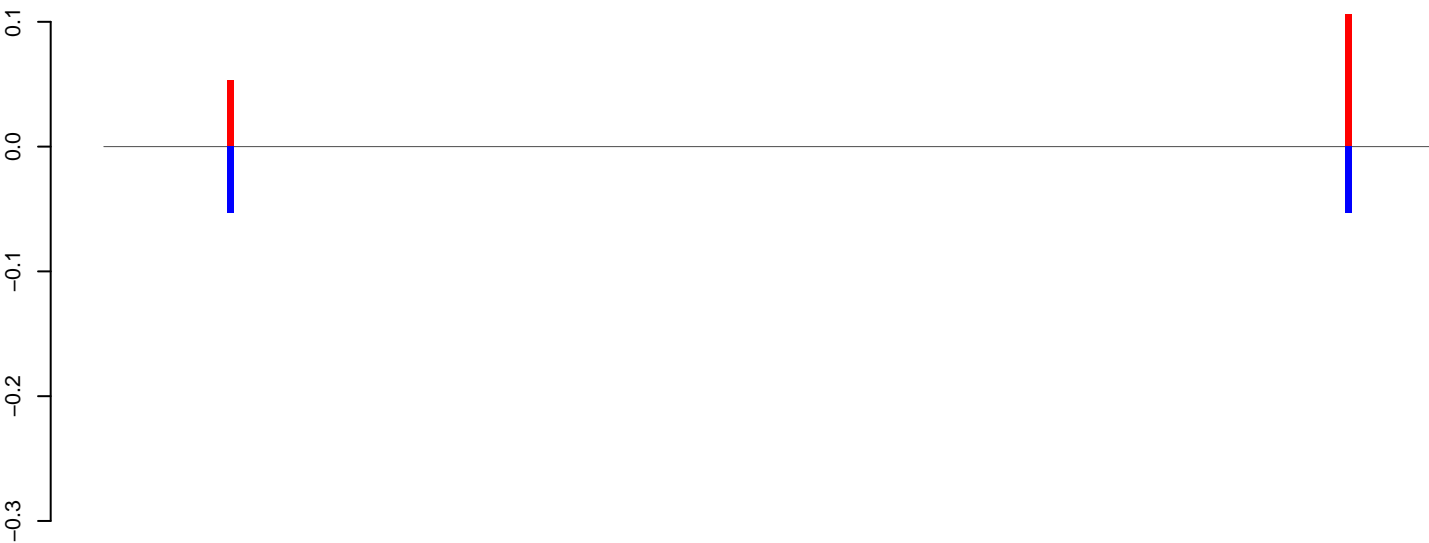


AeAeg_CCL.125cells_CHIKV.rep



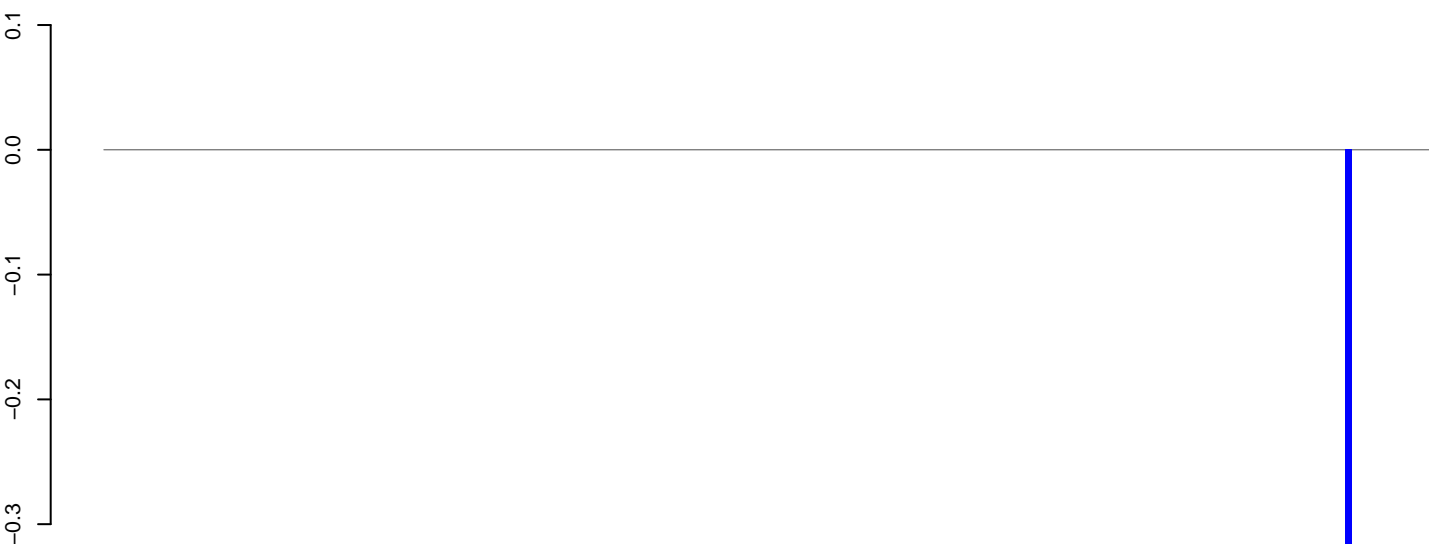
0 2000 4000 6000 8000 10000

AeAeg_CCL.125cells_CHIKV.18_23.rep



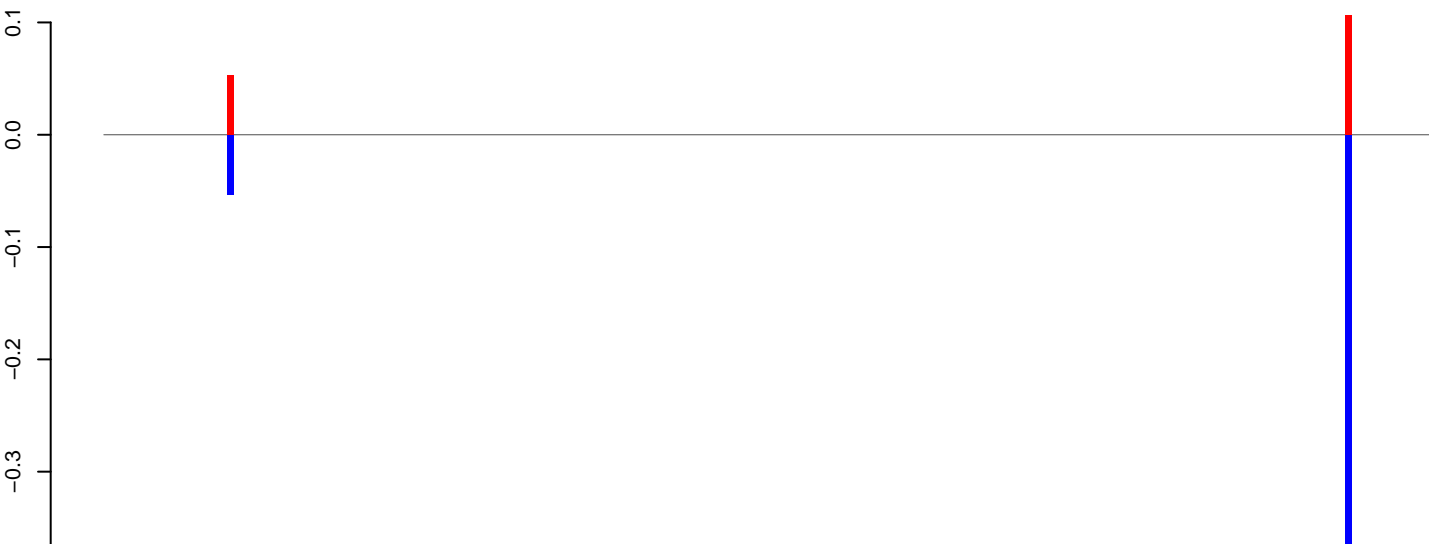
positive=0, negative=0, total=0

AeAeg_CCL.125cells_CHIKV.24_35.rep



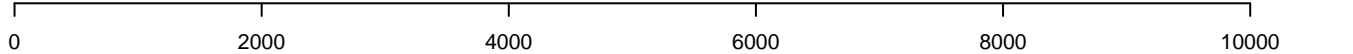
positive=0, negative=0, total=0

AeAeg_CCL.125cells_CHIKV.rep

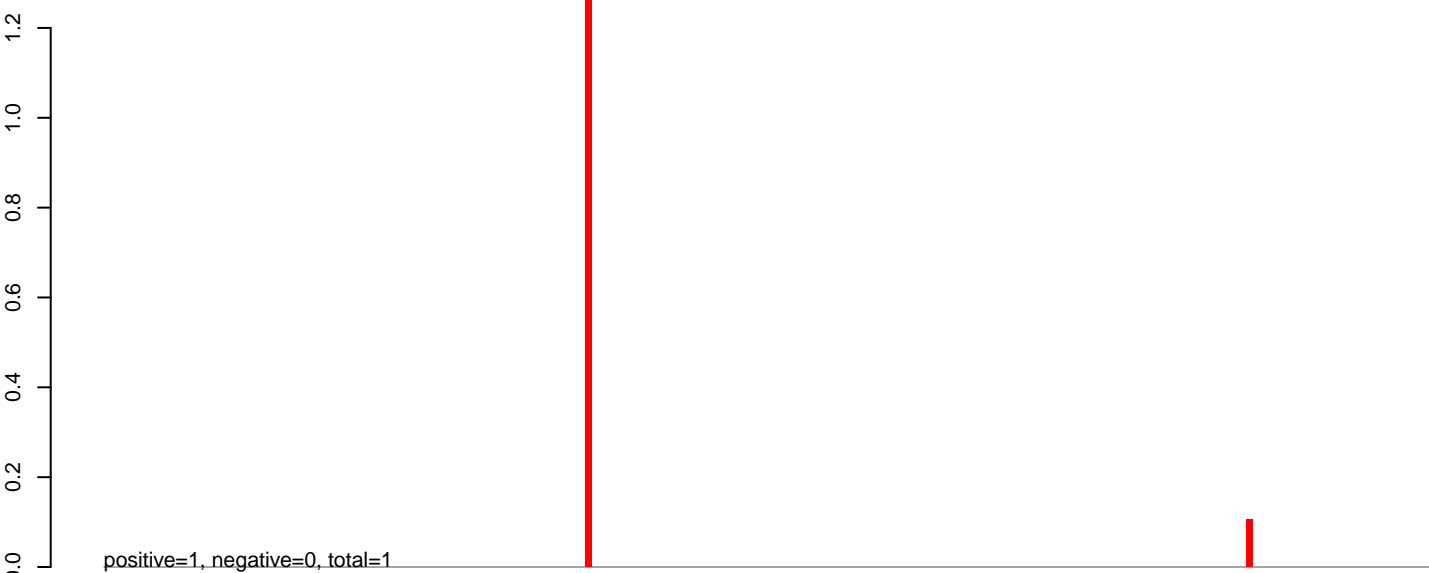


positive=0, negative=0, total=1

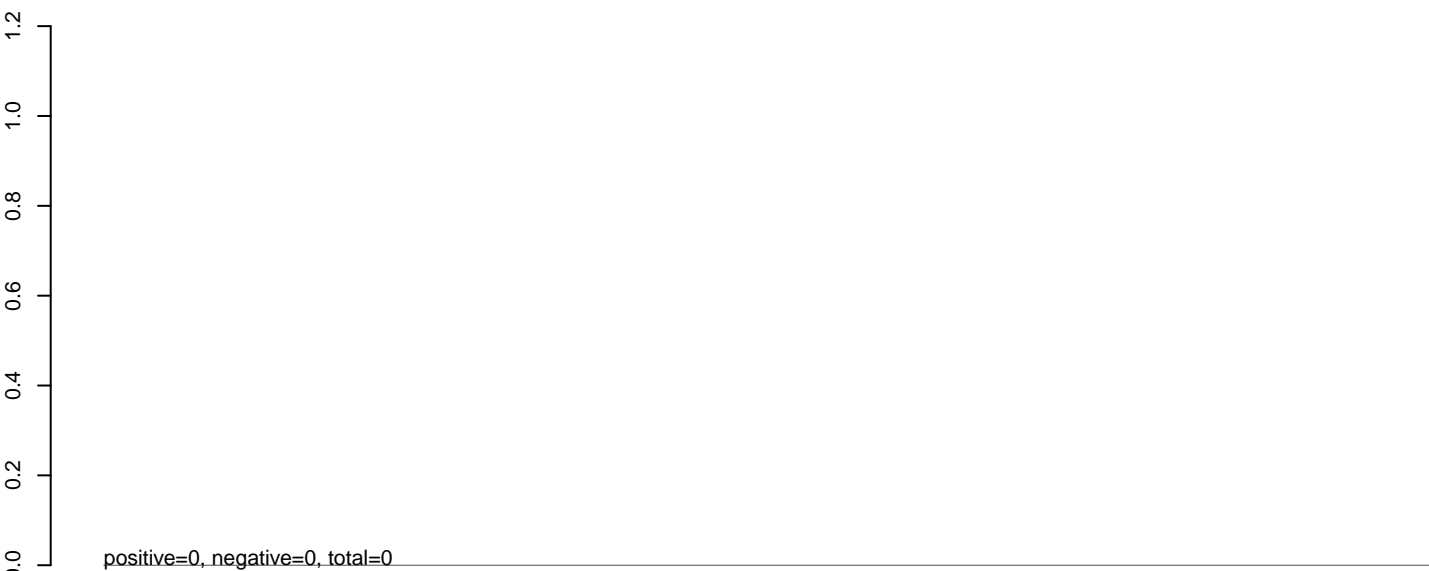
Window size=50, length=10755, virus@NC_009028.2-Ilheus_virus1-10755



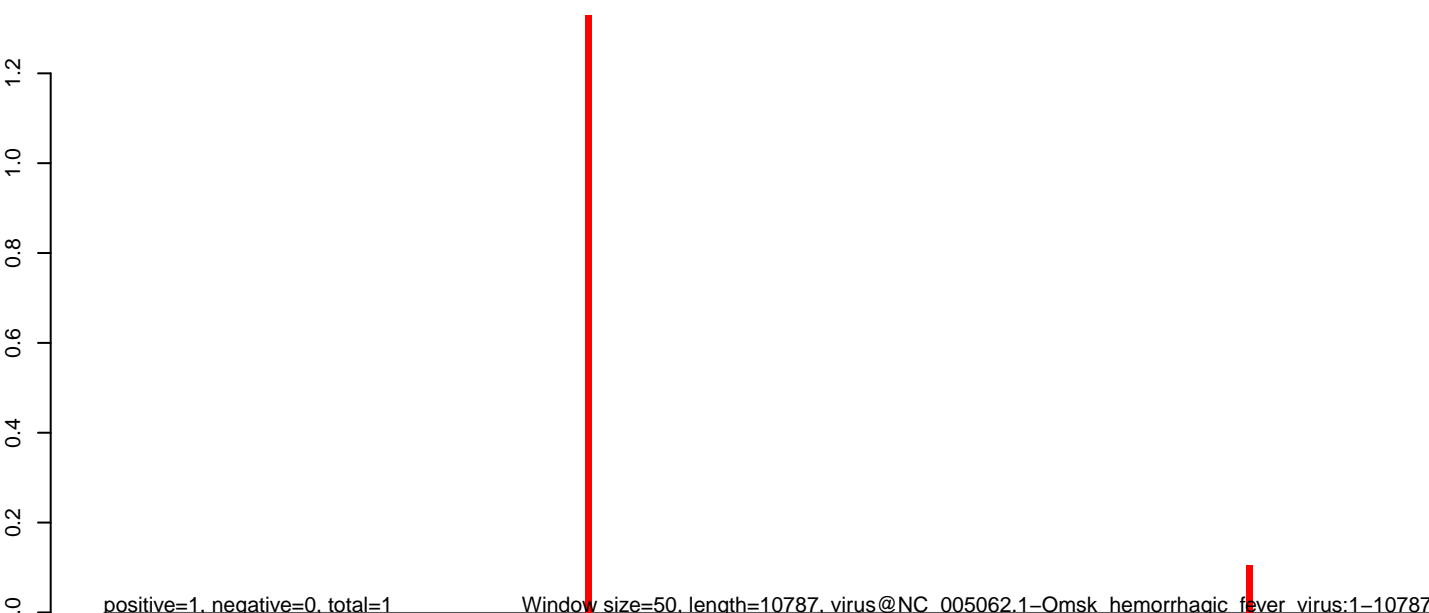
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

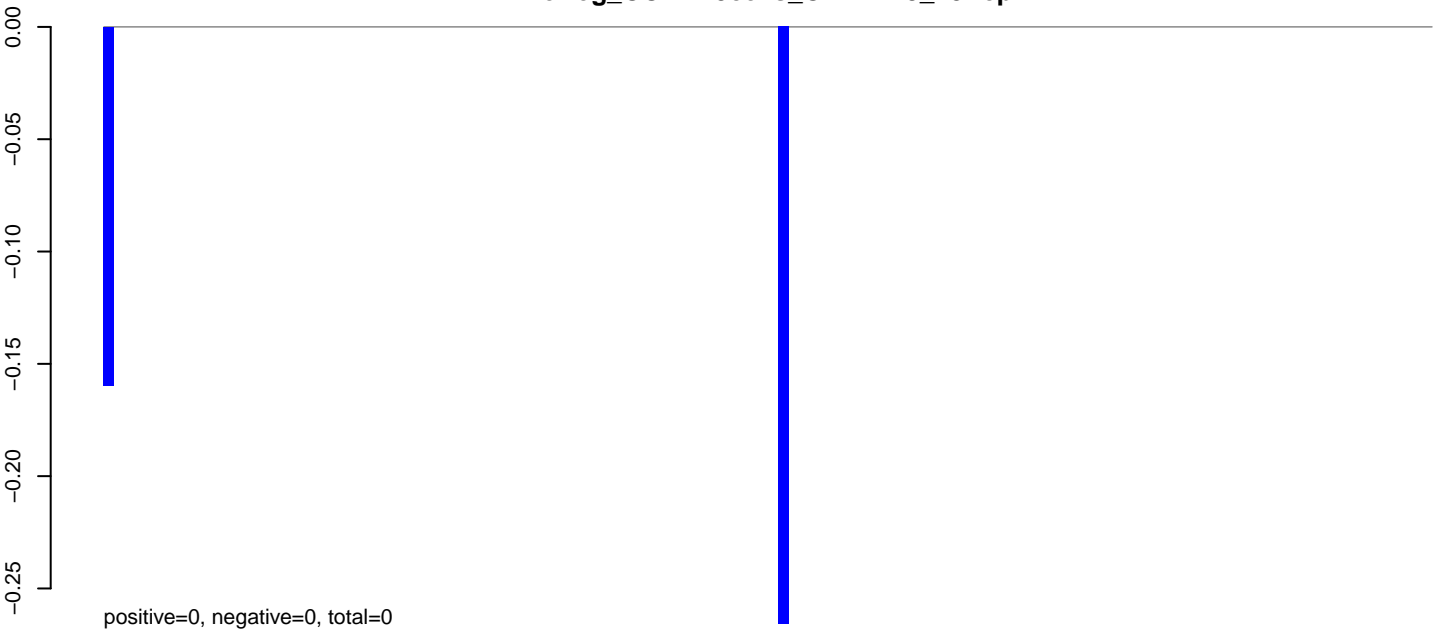


AeAeg_CCL.125cells_CHIKV.rep

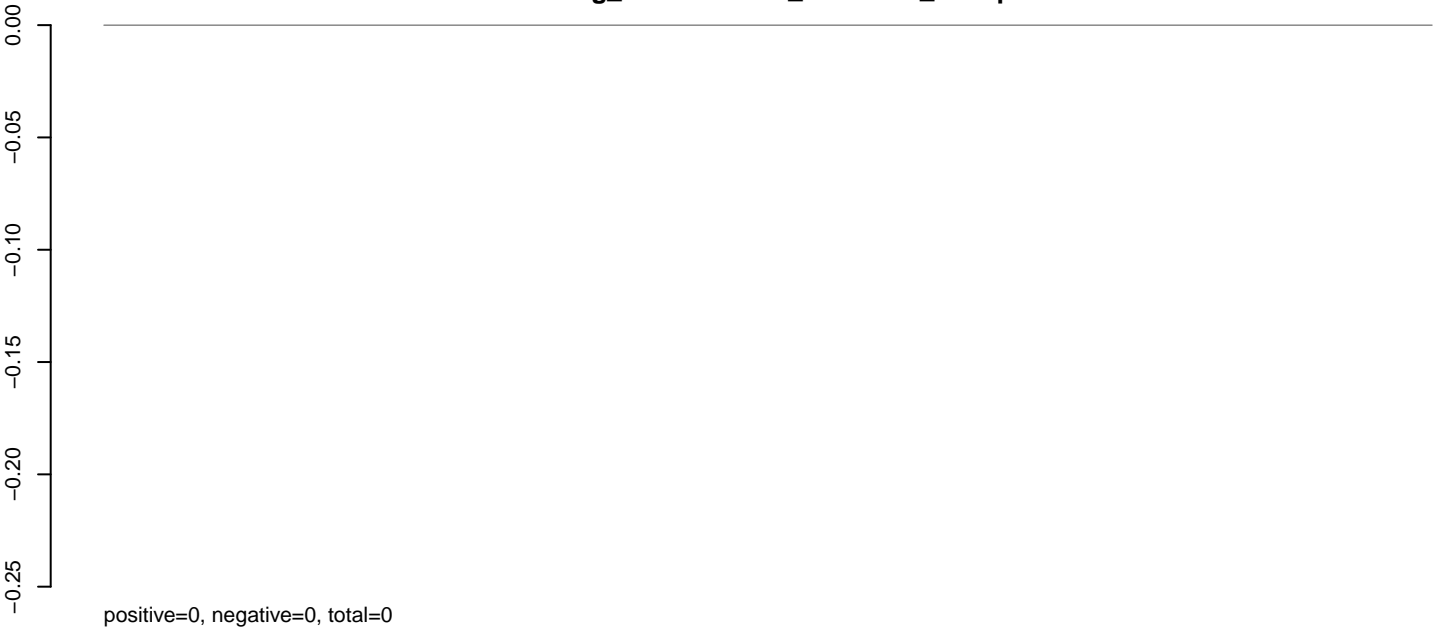


0 2000 4000 6000 8000 10000

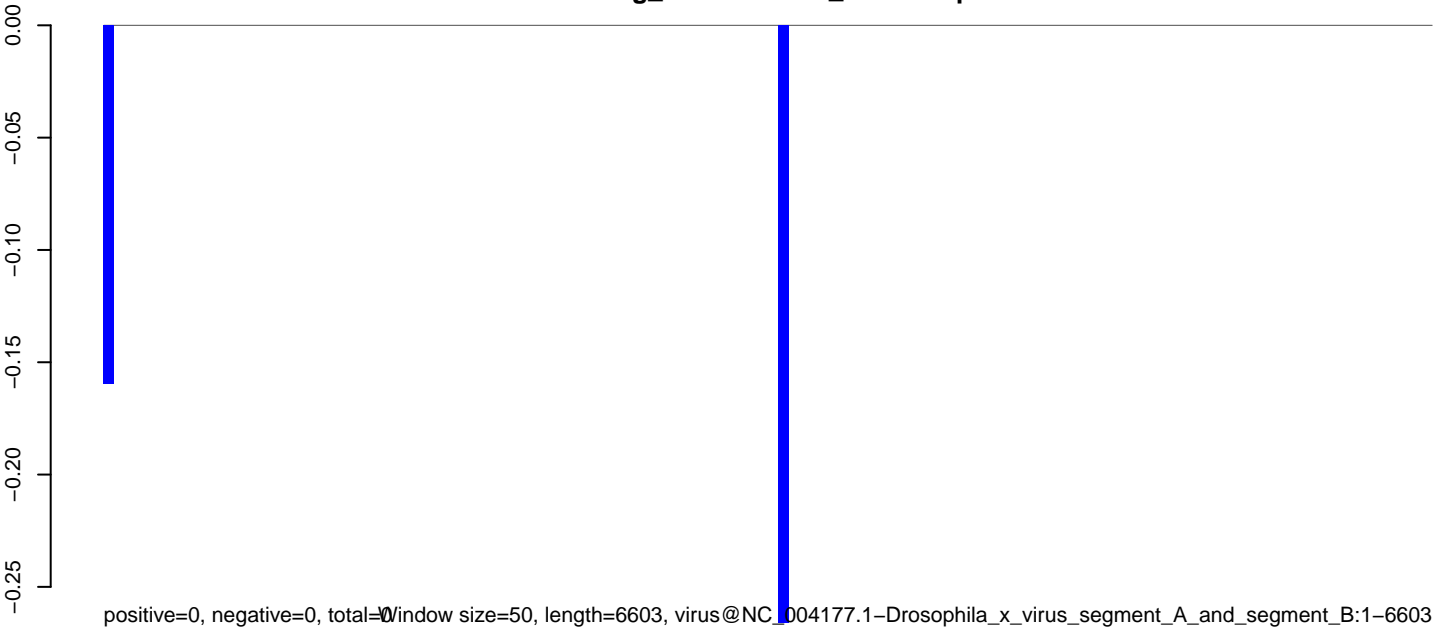
AeAeg_CCL.125cells_CHIKV.18_23.rep



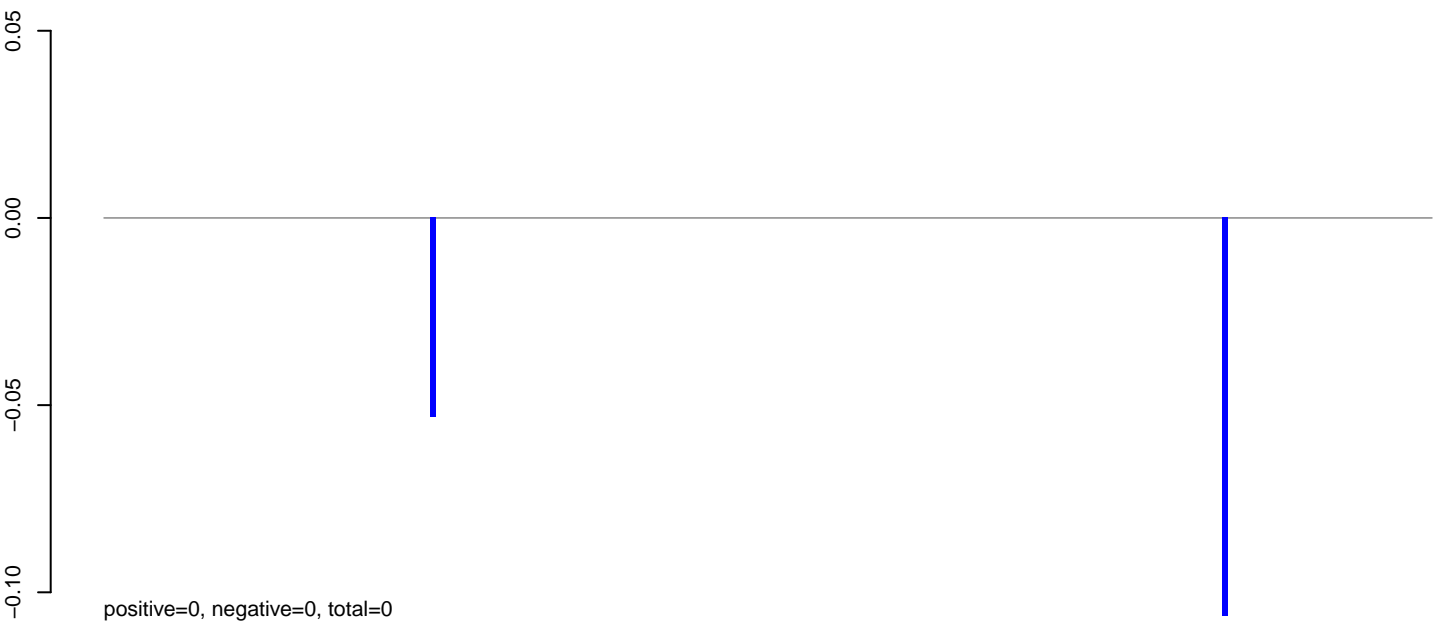
AeAeg_CCL.125cells_CHIKV.24_35.rep



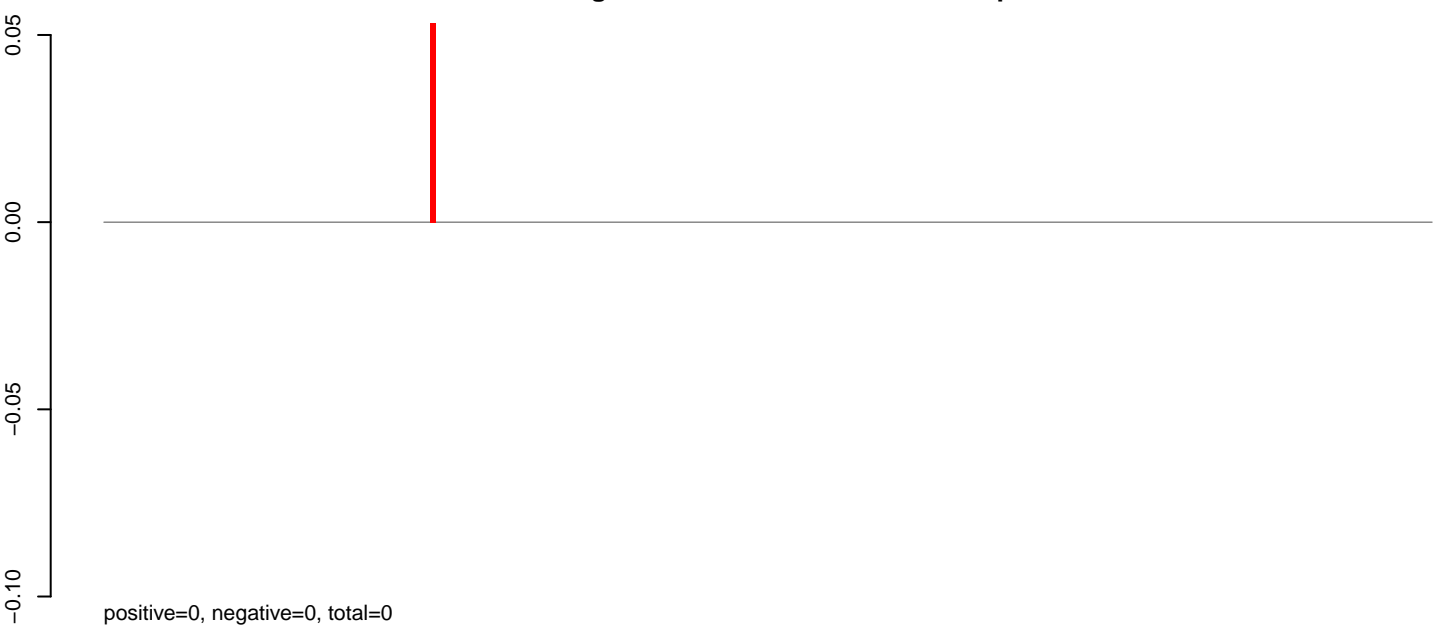
AeAeg_CCL.125cells_CHIKV.rep



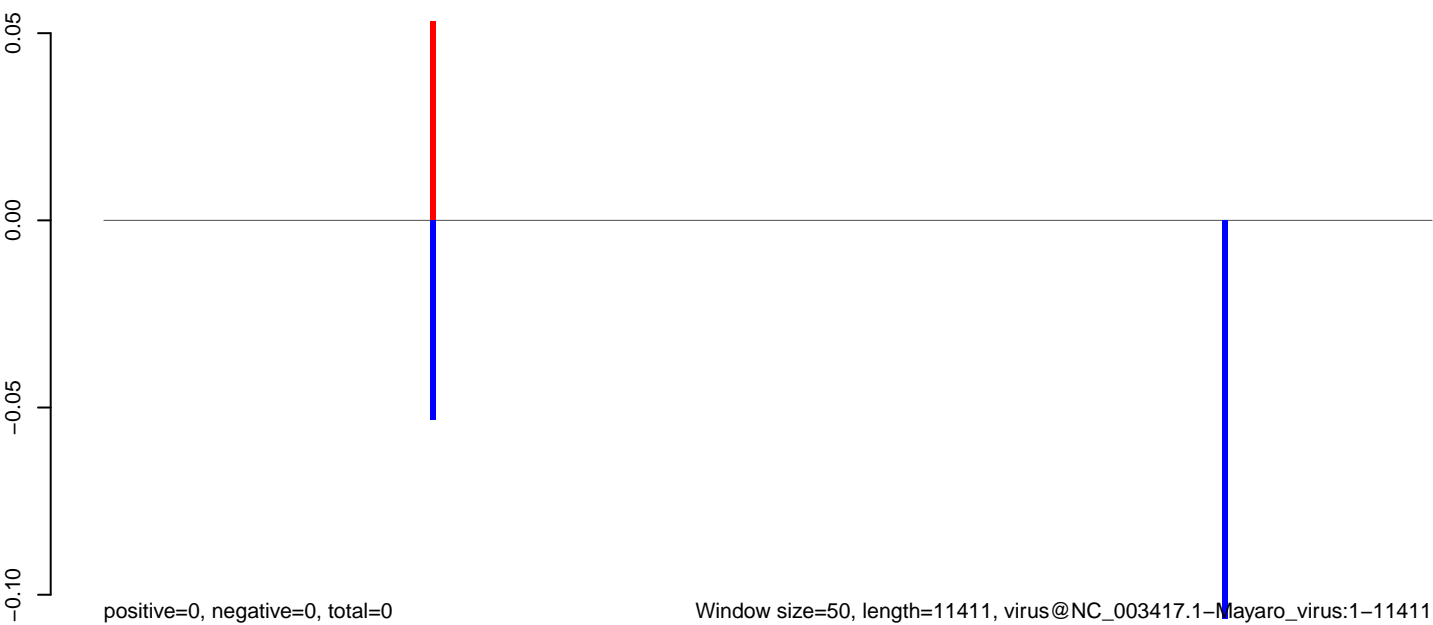
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

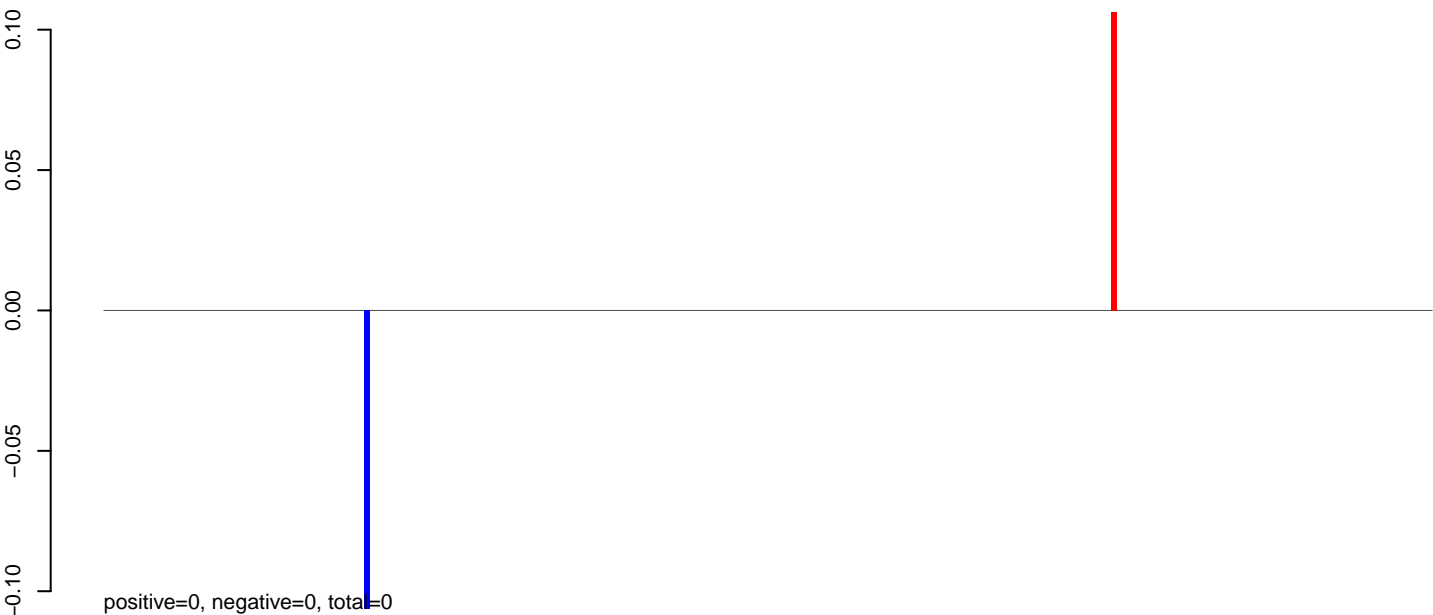


AeAeg_CCL.125cells_CHIKV.rep

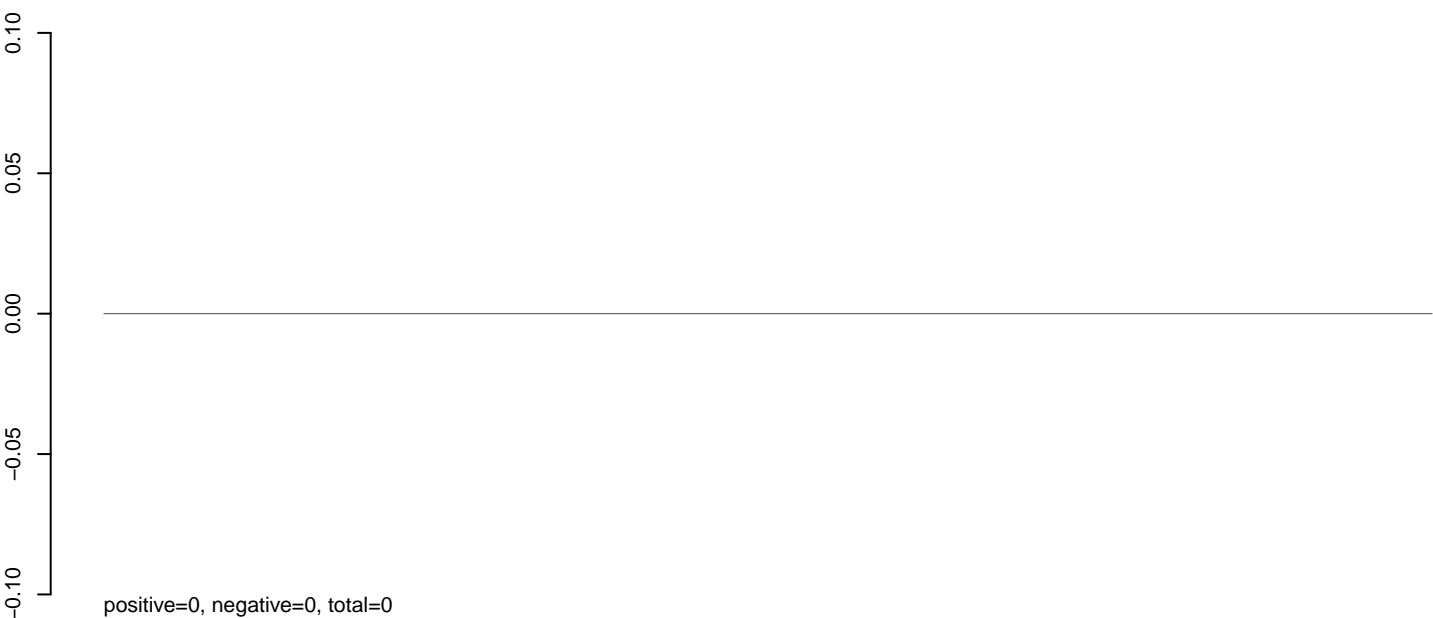


0 2000 4000 6000 8000 10000

AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

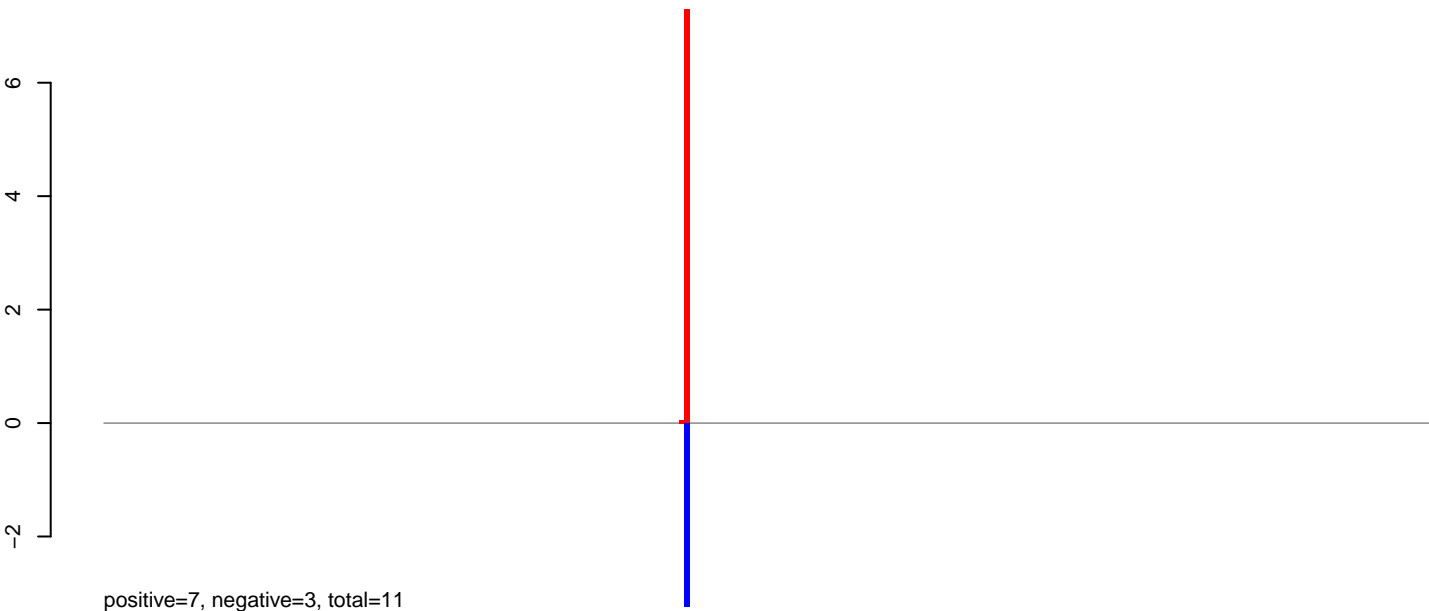


AeAeg_CCL.125cells_CHIKV.rep

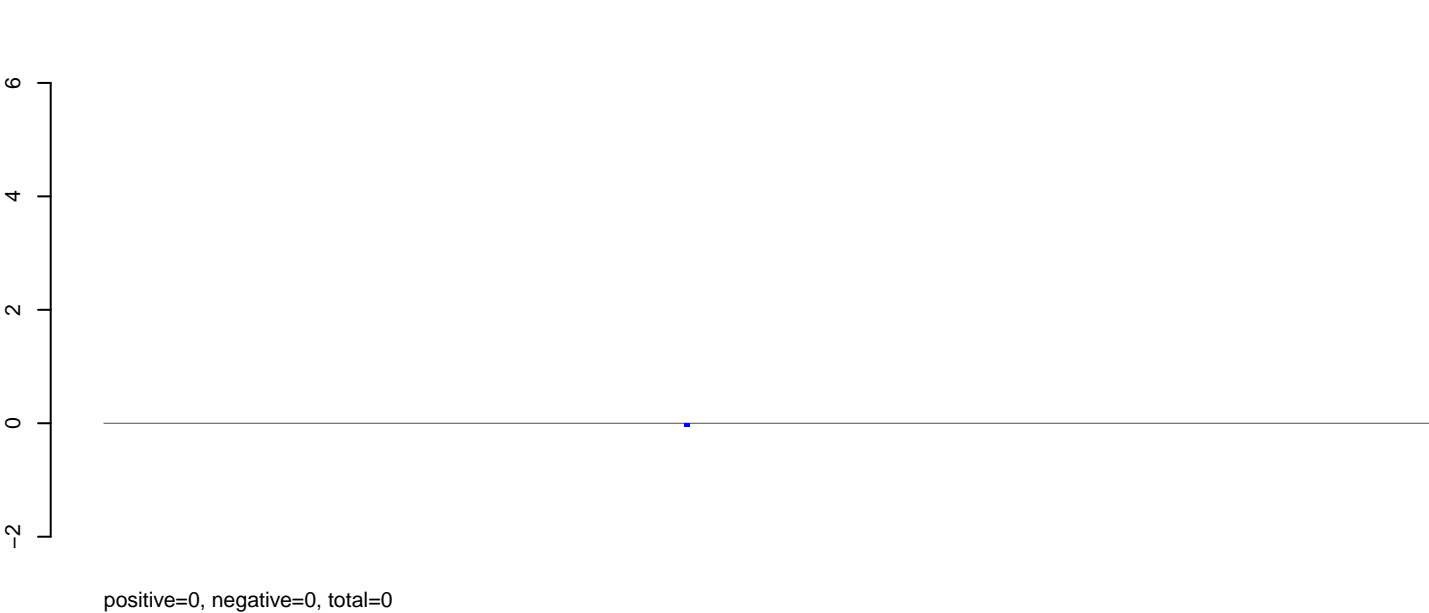


0 2000 4000 6000 8000 10000 12000
positive=0, negative=0, total=0, width=50, length=13297, virus@MF176298.1-Culex_mononega-like_virus_2_strain_mos191gb23464:1-13297

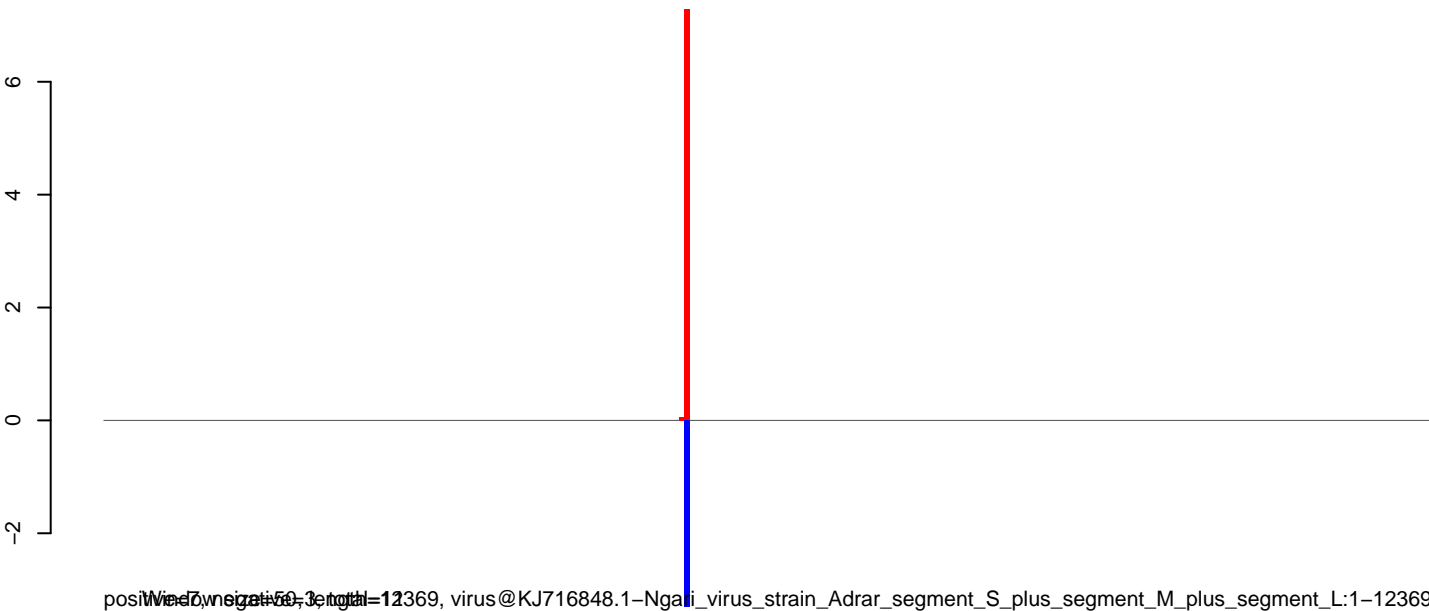
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



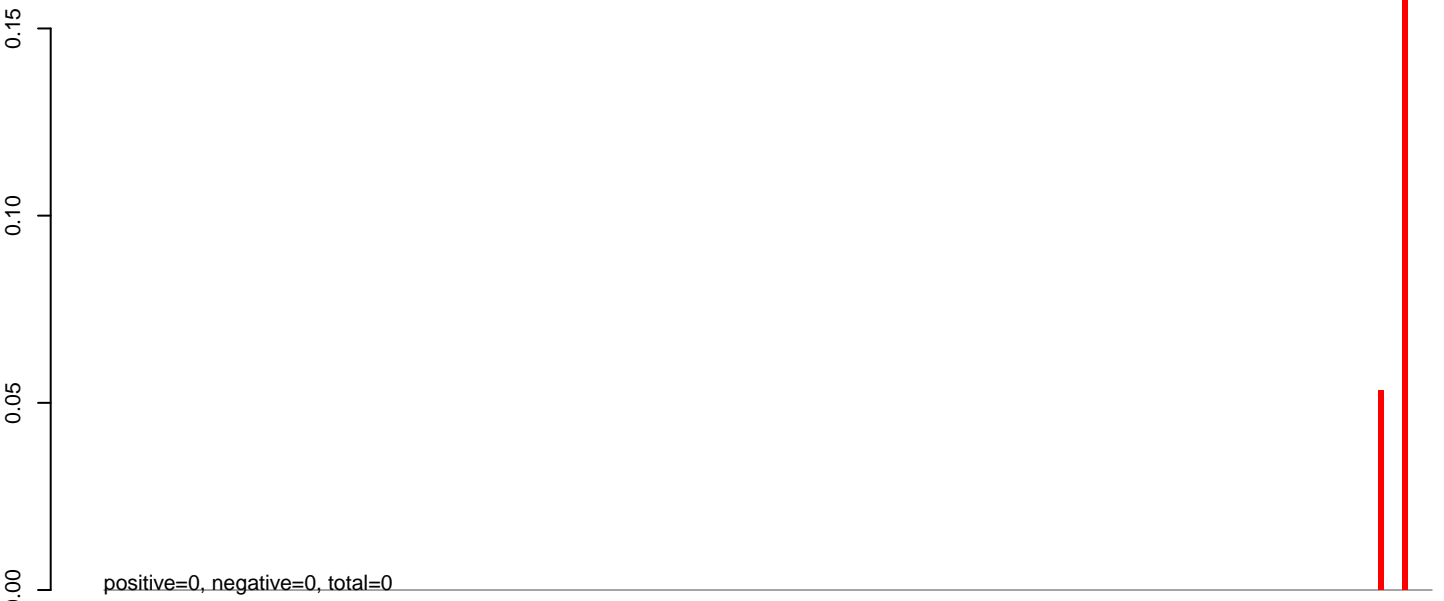
AeAeg_CCL.125cells_CHIKV.rep



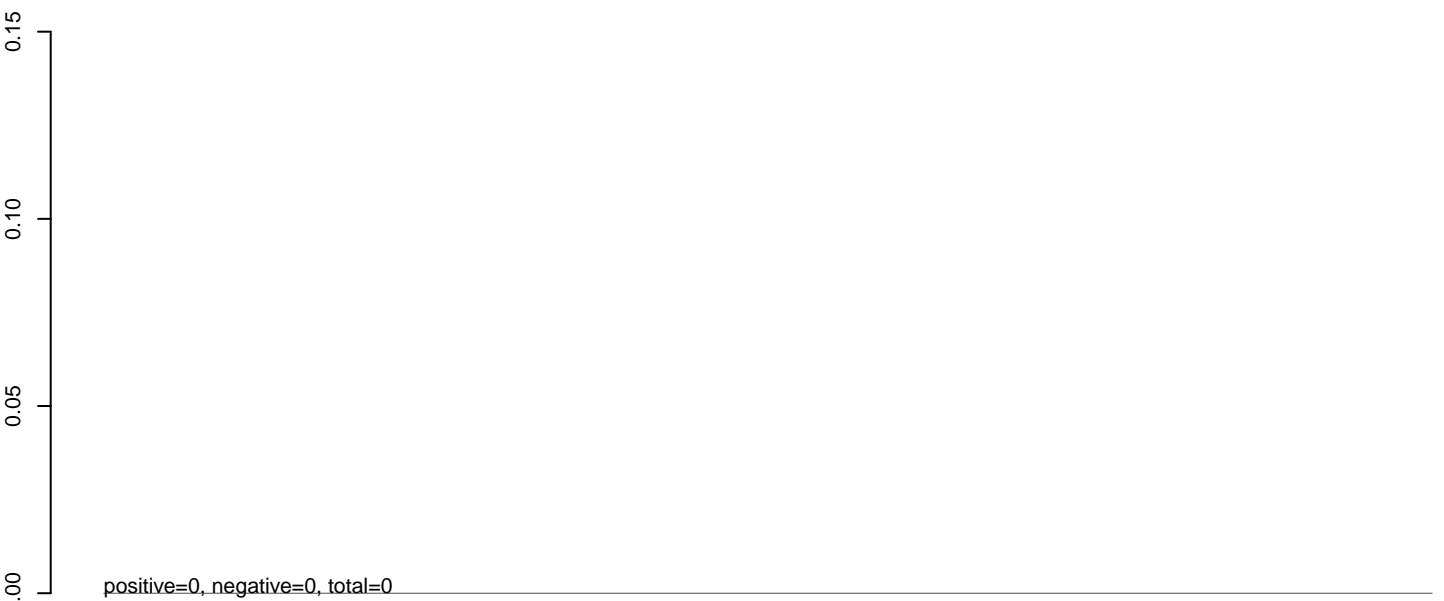
12369, virus@KJ716848.1-Ngali_virus_strain_Adrar_segment_S_plus_segment_M_plus_segment_L:1-12369

0 2000 4000 6000 8000 10000 12000

AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

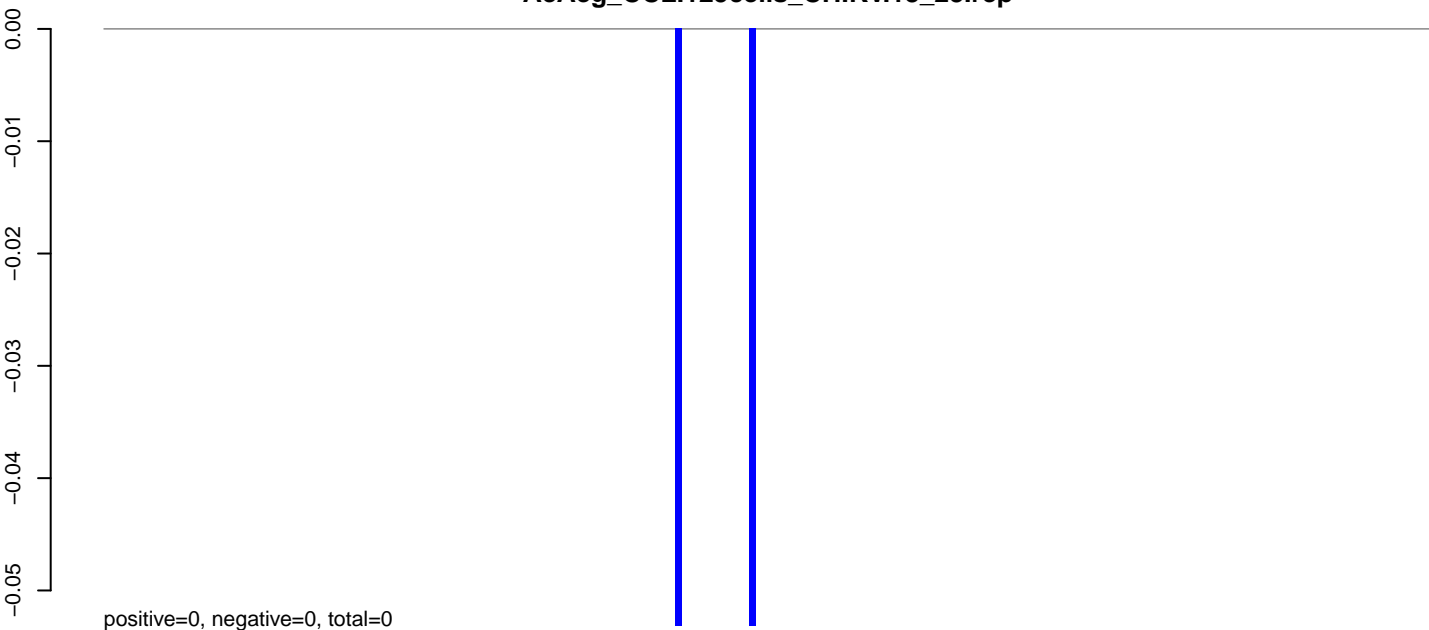


AeAeg_CCL.125cells_CHIKV.rep

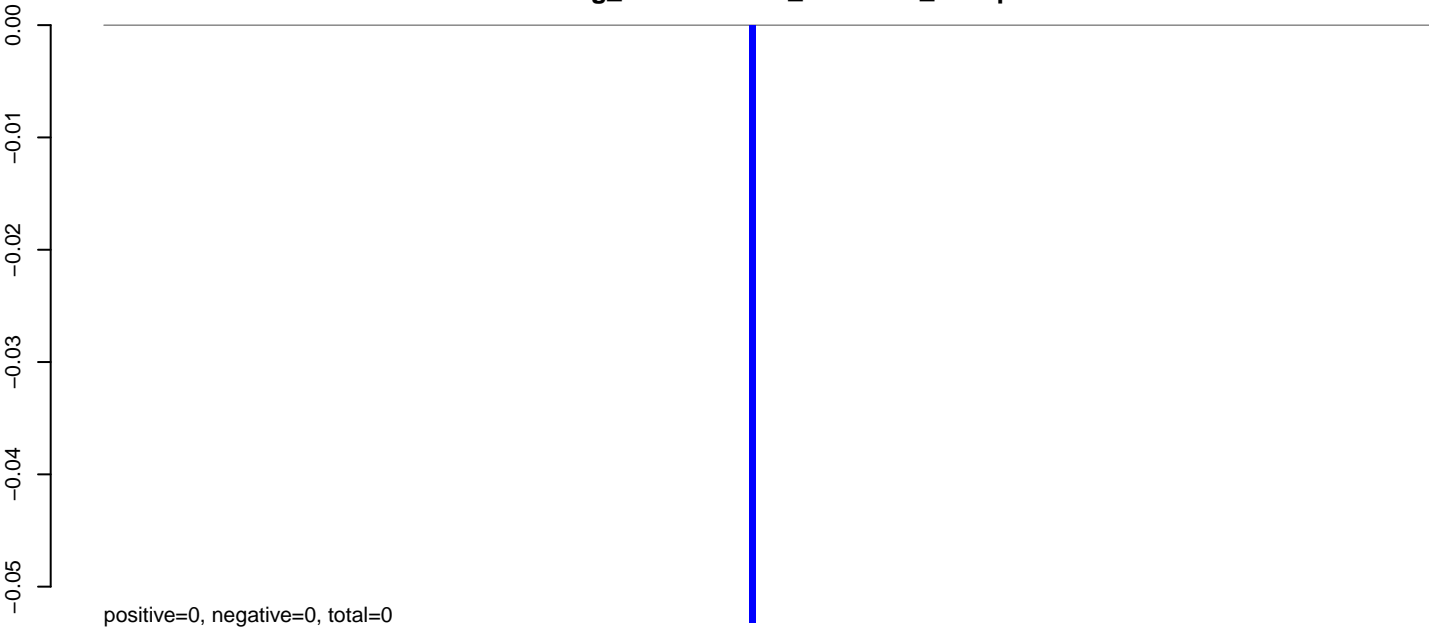


Window size=50, length=11079, virus@KC181923.1-Aedes flavivirus_strain_AEFV-SPFLD-MO-2011-MP6:1-11079

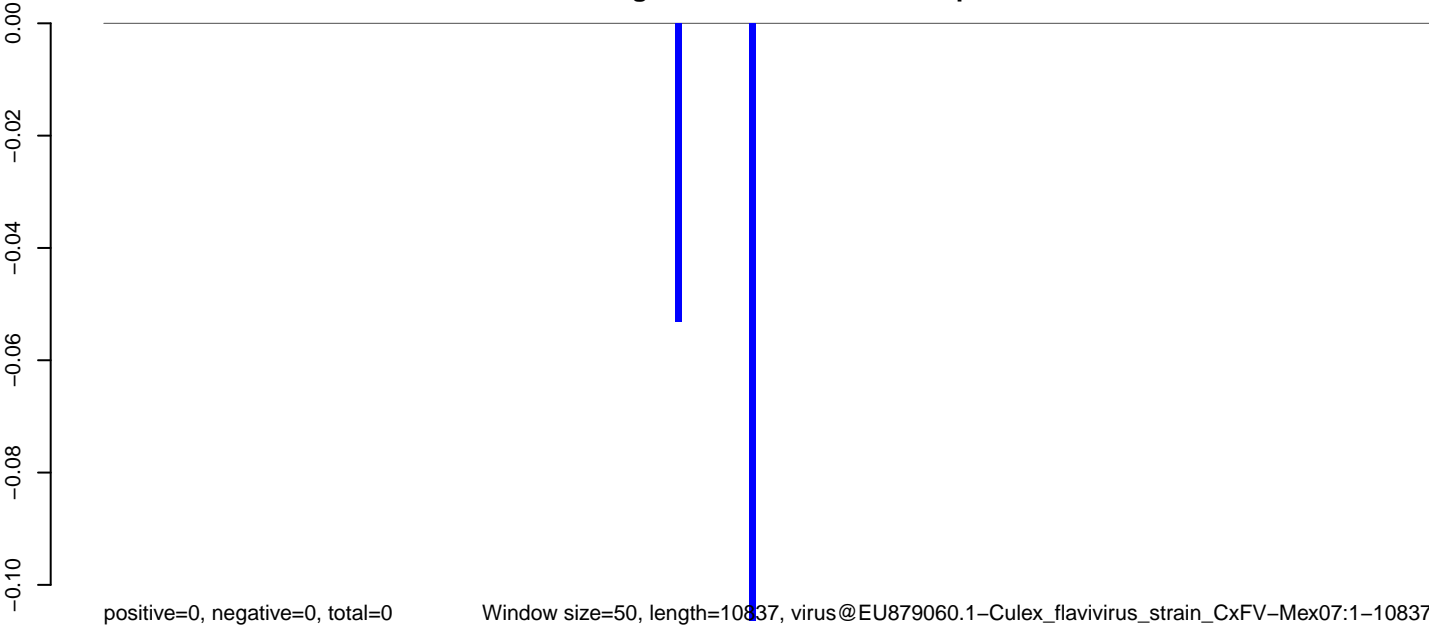
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

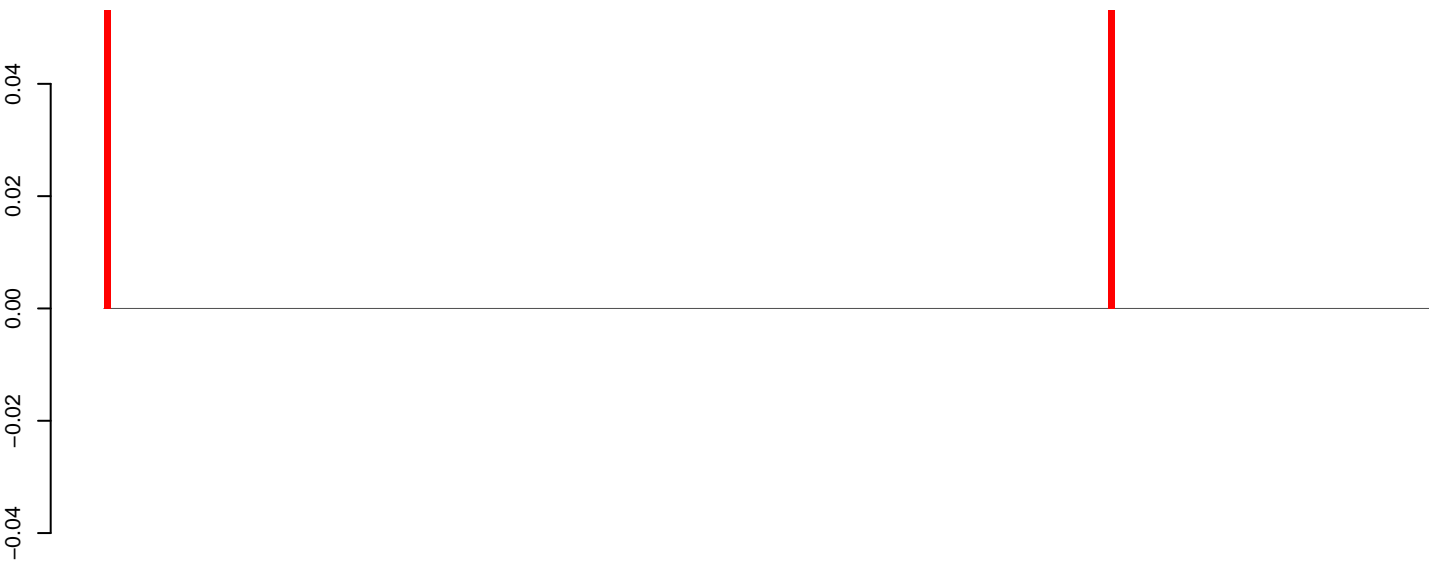


AeAeg_CCL.125cells_CHIKV.rep



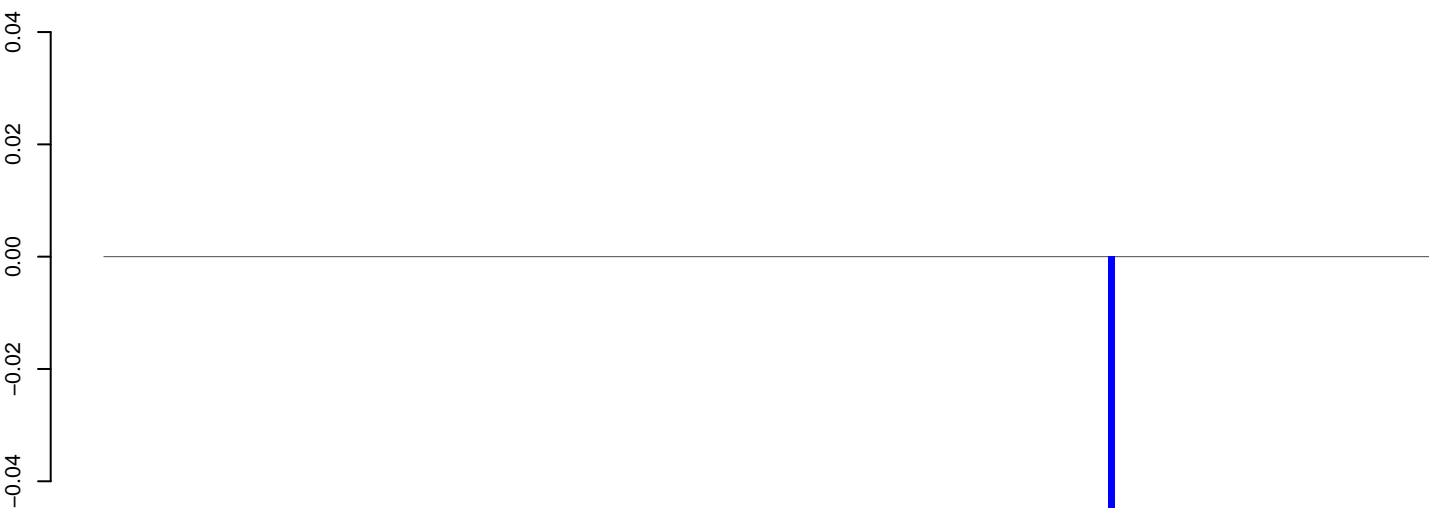
0 2000 4000 6000 8000 10000

AeAeg_CCL.125cells_CHIKV.18_23.rep



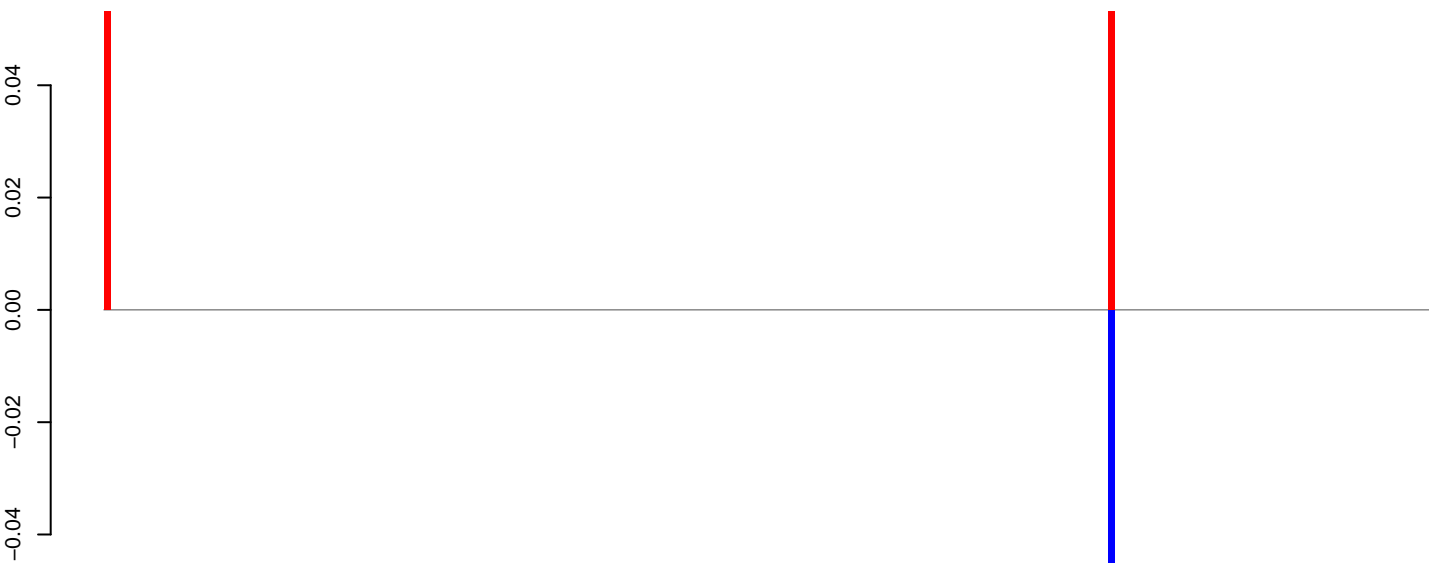
positive=0, negative=0, total=0

AeAeg_CCL.125cells_CHIKV.24_35.rep



positive=0, negative=0, total=0

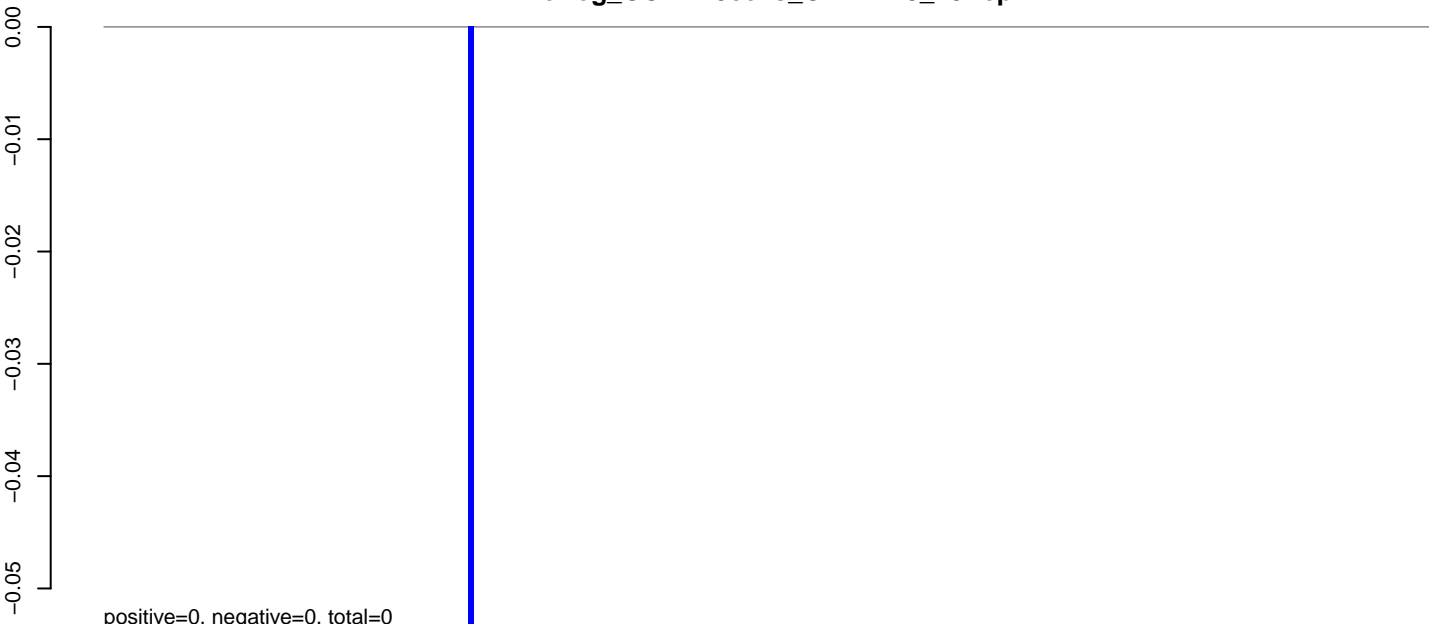
AeAeg_CCL.125cells_CHIKV.rep



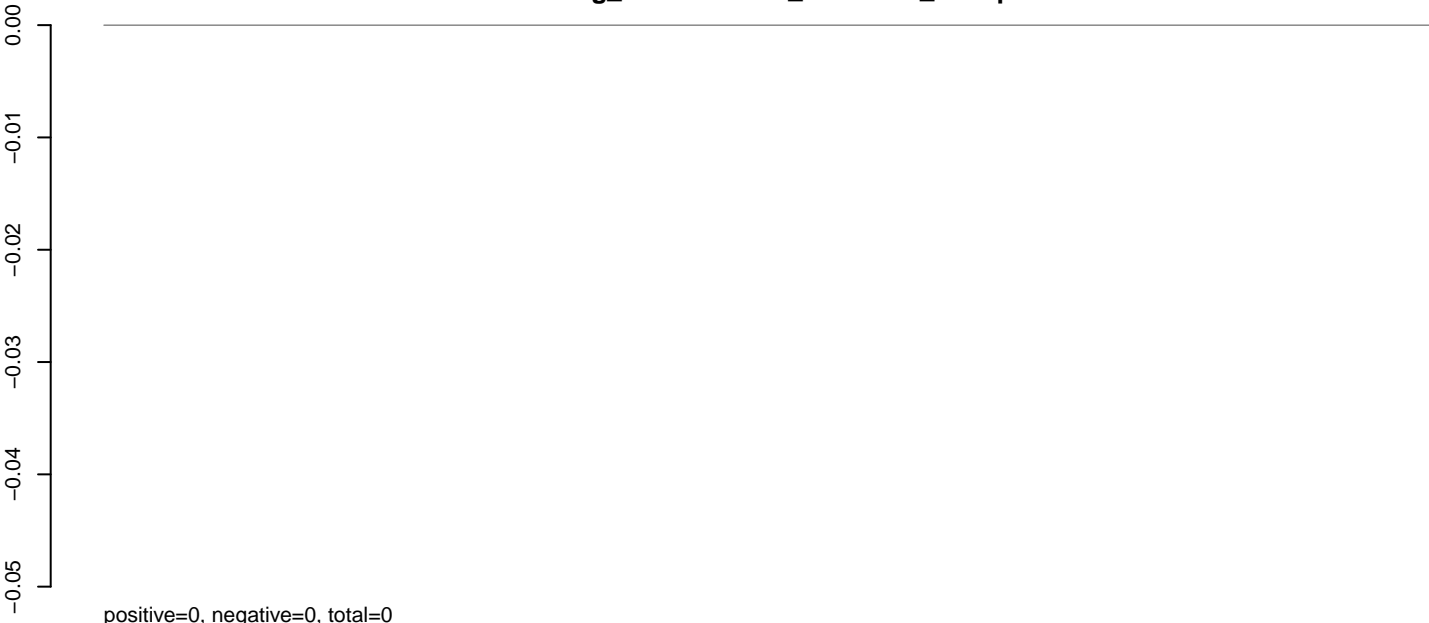
positive=0, negative=0, total=0, window size=50, length=10254, virus@DQ235149.1-Royal_Farm_virus_from_Afghanistan_polyprotein_gene:1-10254

0 2000 4000 6000 8000 10000

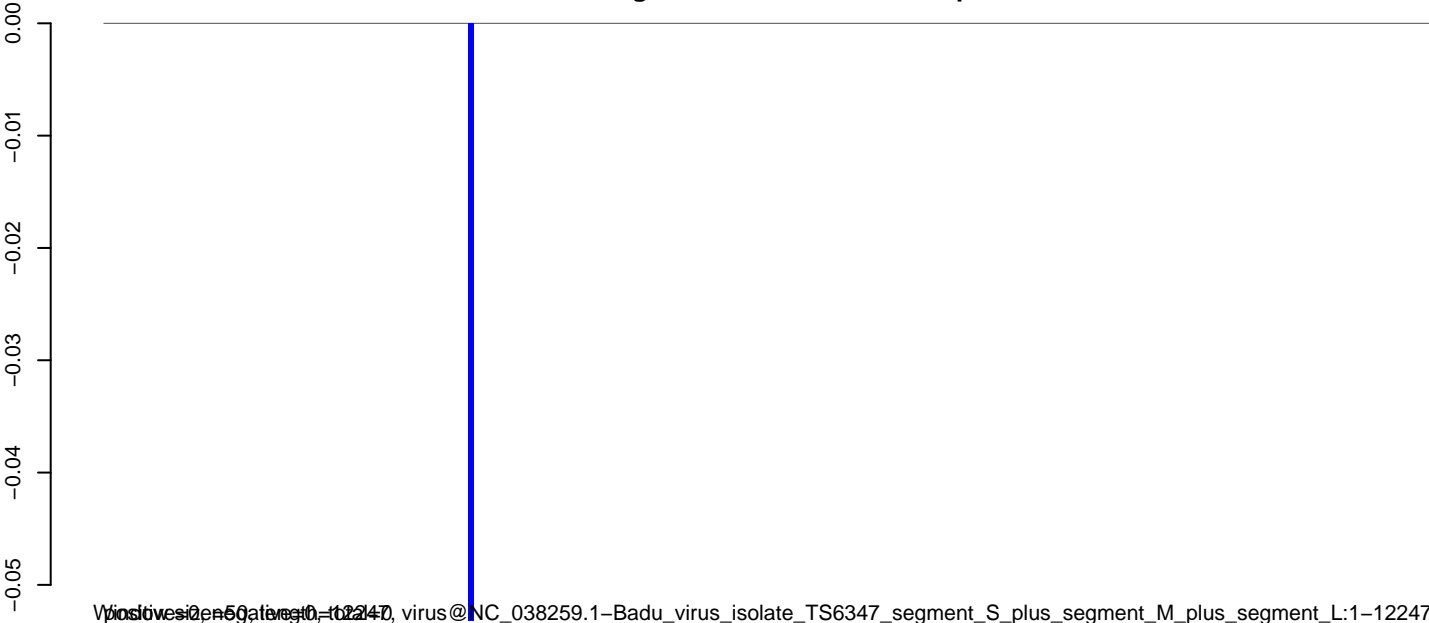
AeAeg_CCL.125cells_CHIKV.18_23.rep



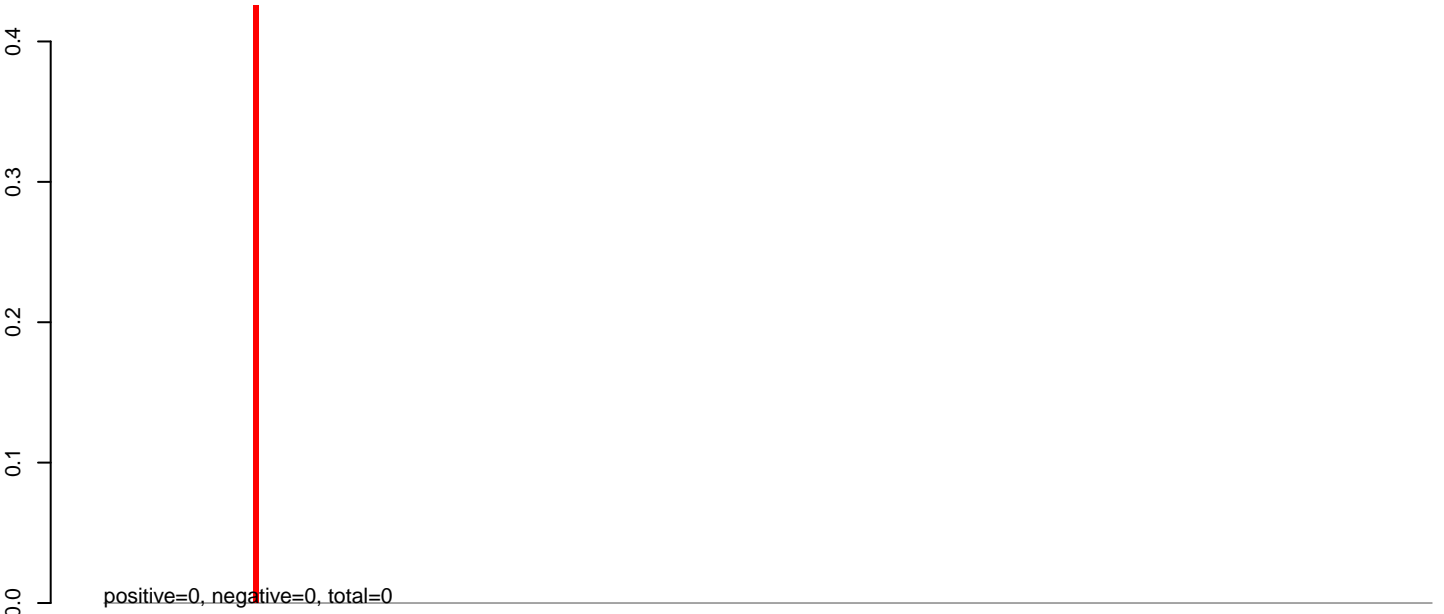
AeAeg_CCL.125cells_CHIKV.24_35.rep



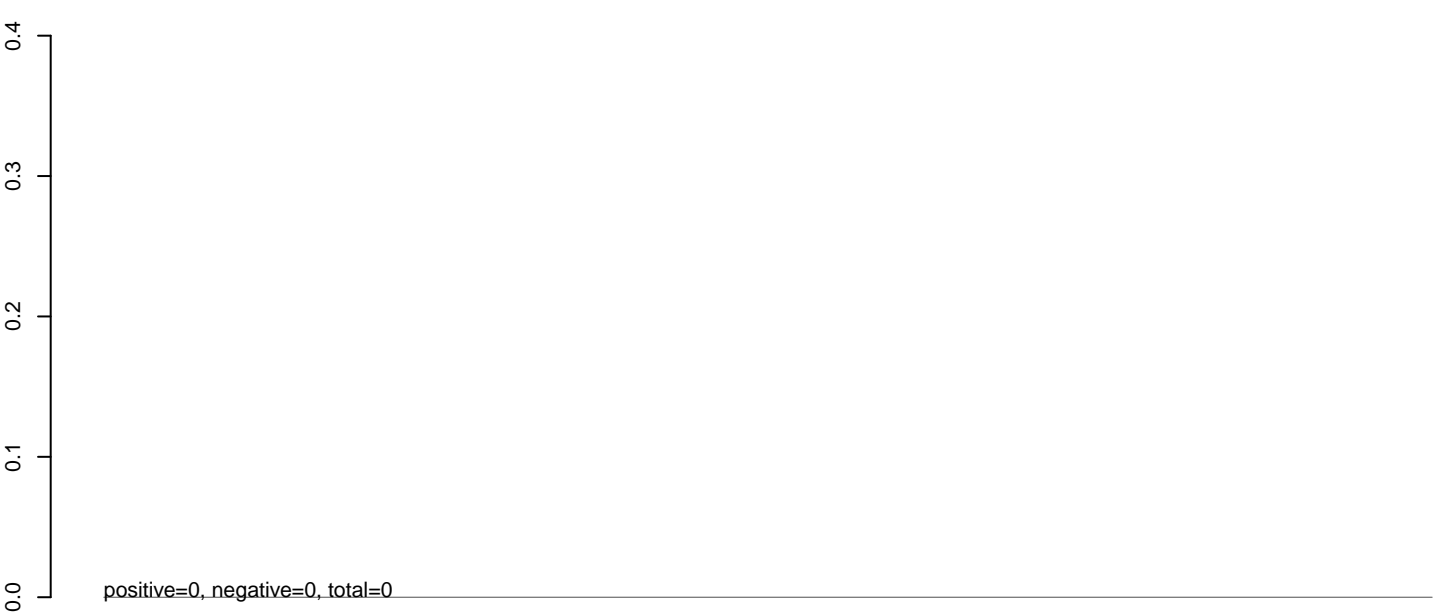
AeAeg_CCL.125cells_CHIKV.rep



AeAeg_CCL.125cells_CHIKV.18_23.rep



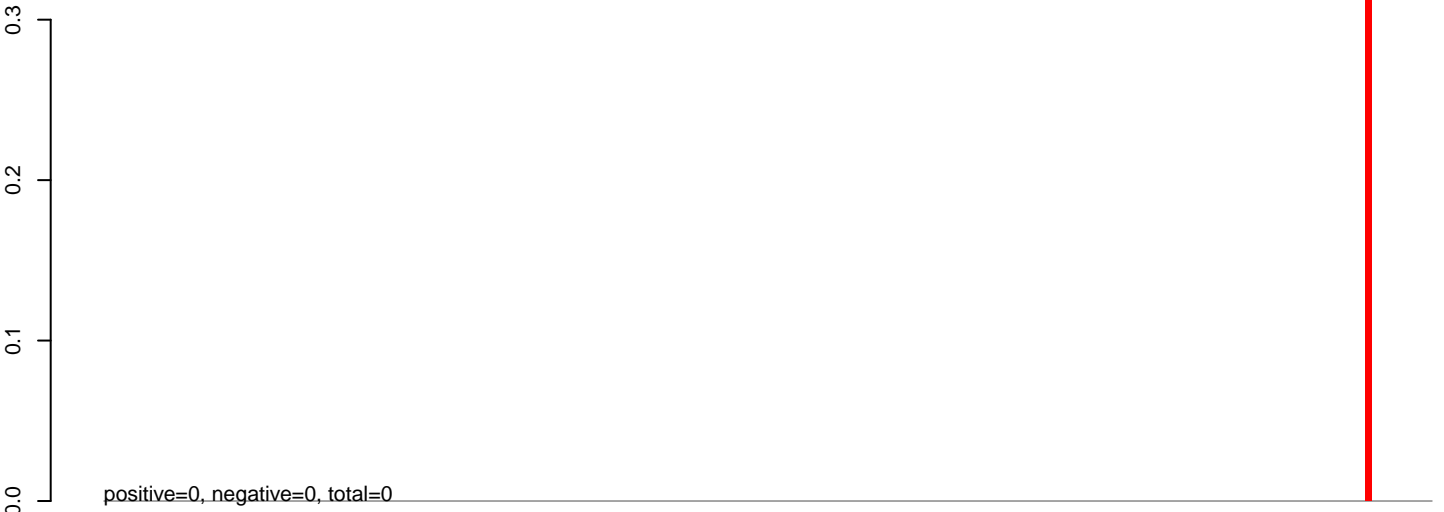
AeAeg_CCL.125cells_CHIKV.24_35.rep



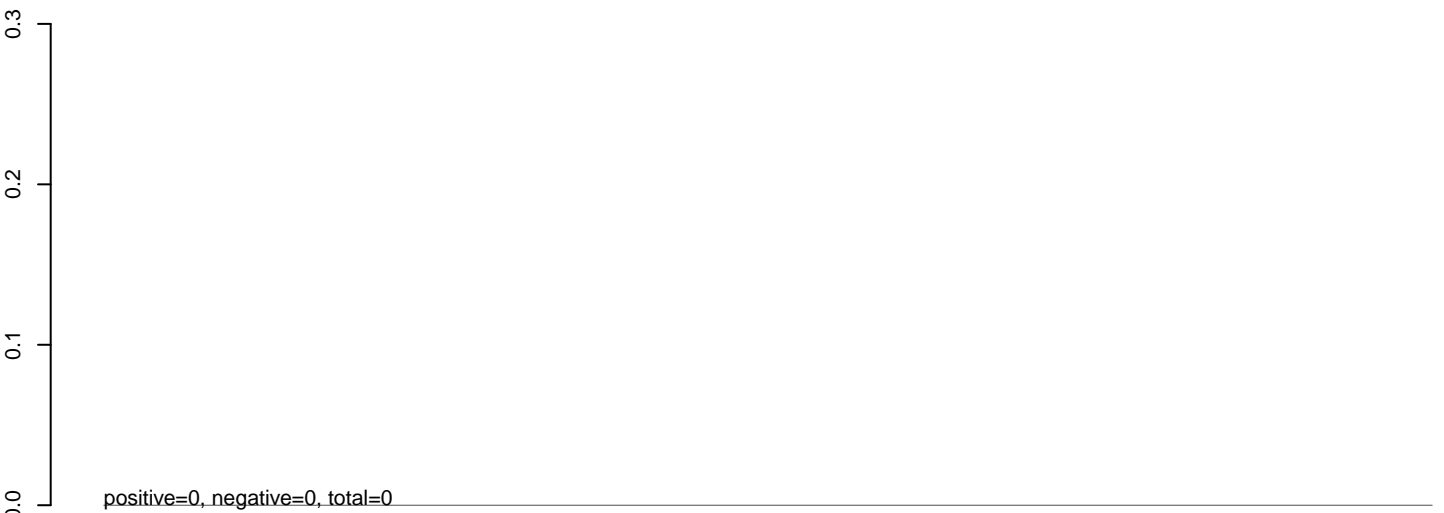
AeAeg_CCL.125cells_CHIKV.rep



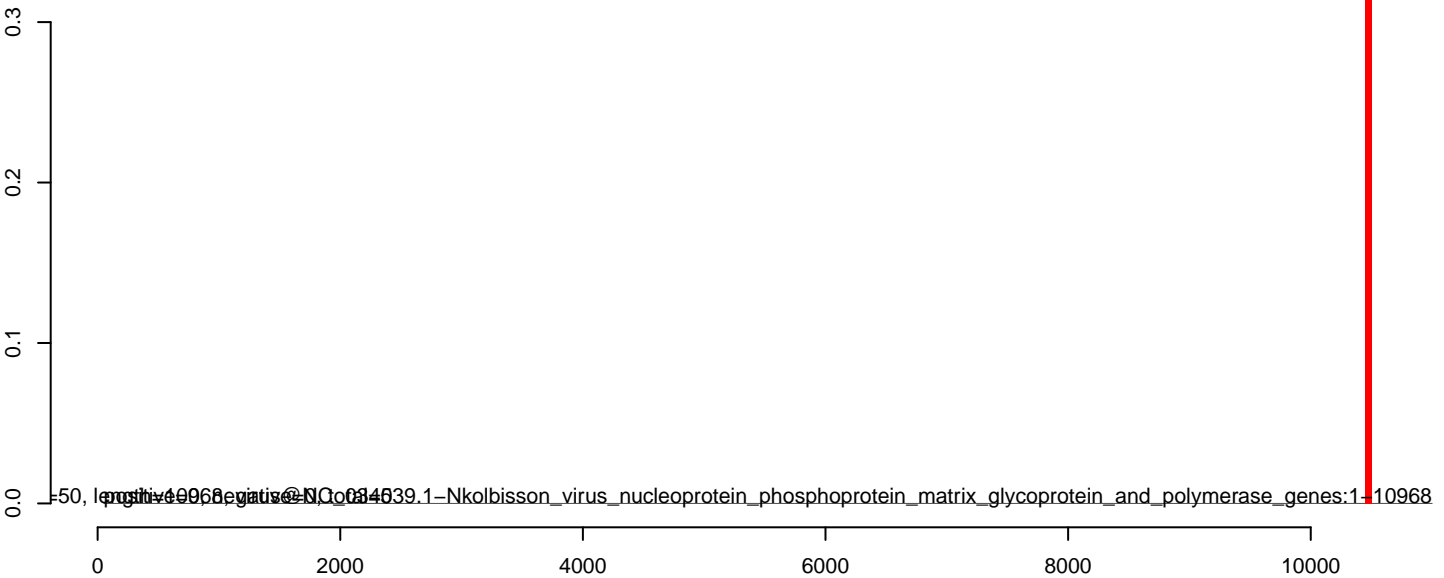
AeAeg_CCL.125cells_CHIKV.18_23.rep



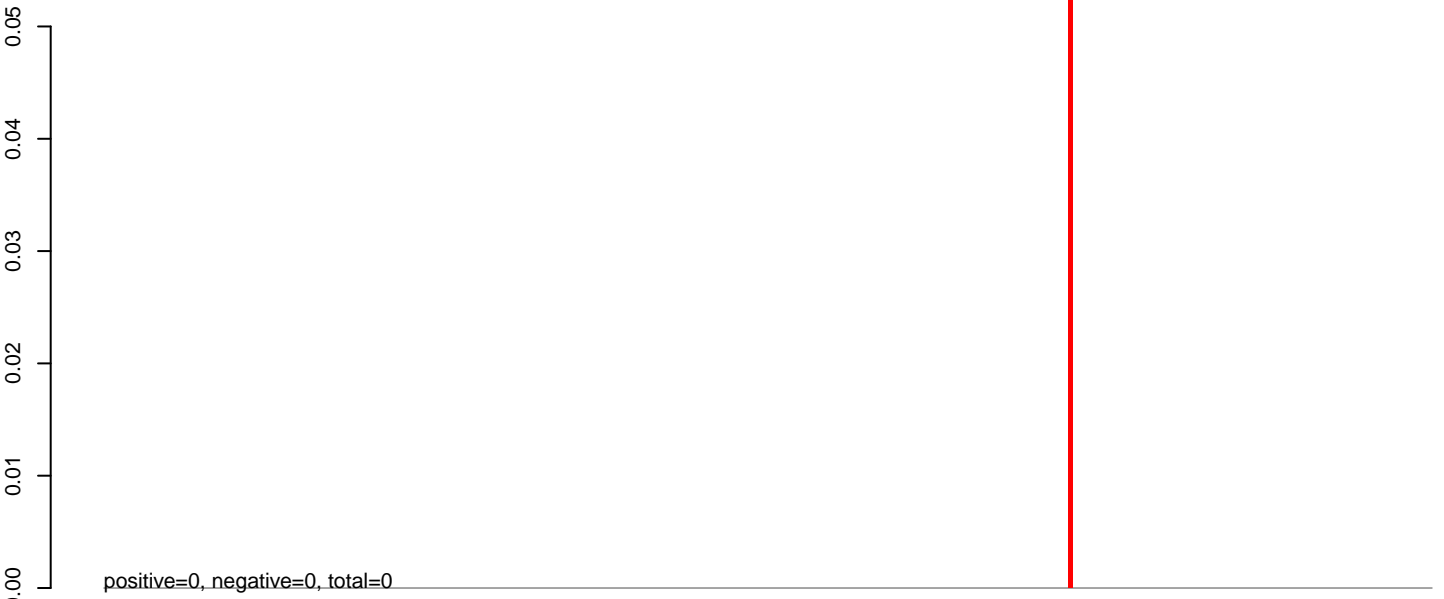
AeAeg_CCL.125cells_CHIKV.24_35.rep



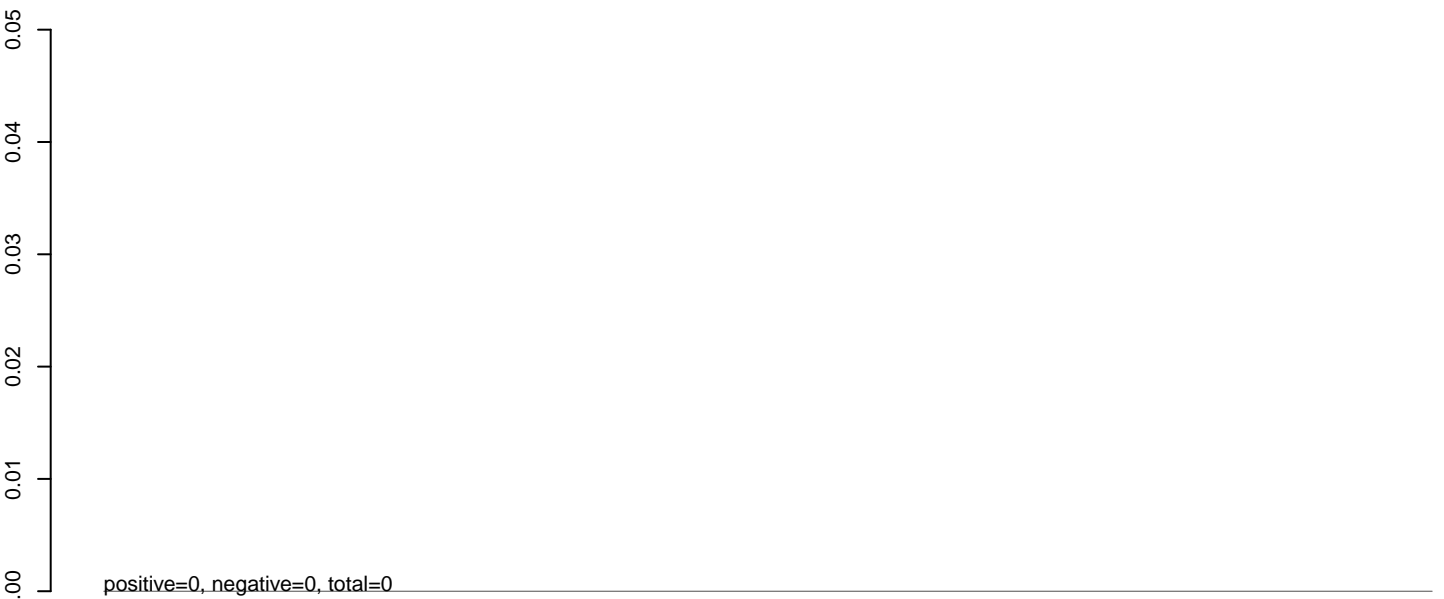
AeAeg_CCL.125cells_CHIKV.rep



AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



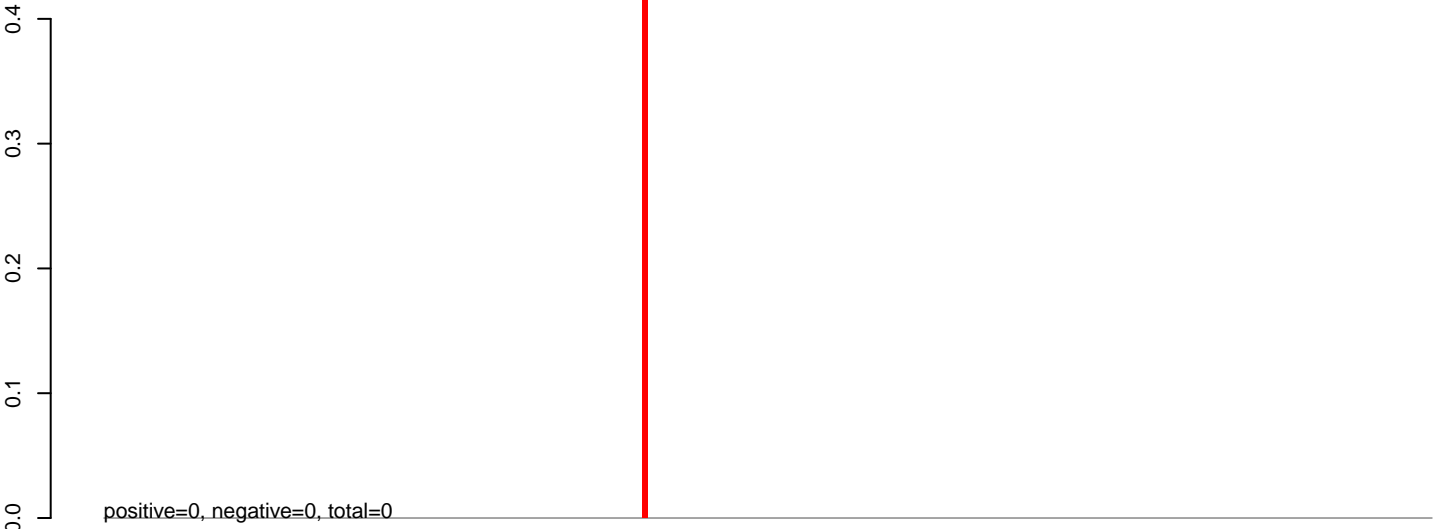
AeAeg_CCL.125cells_CHIKV.rep



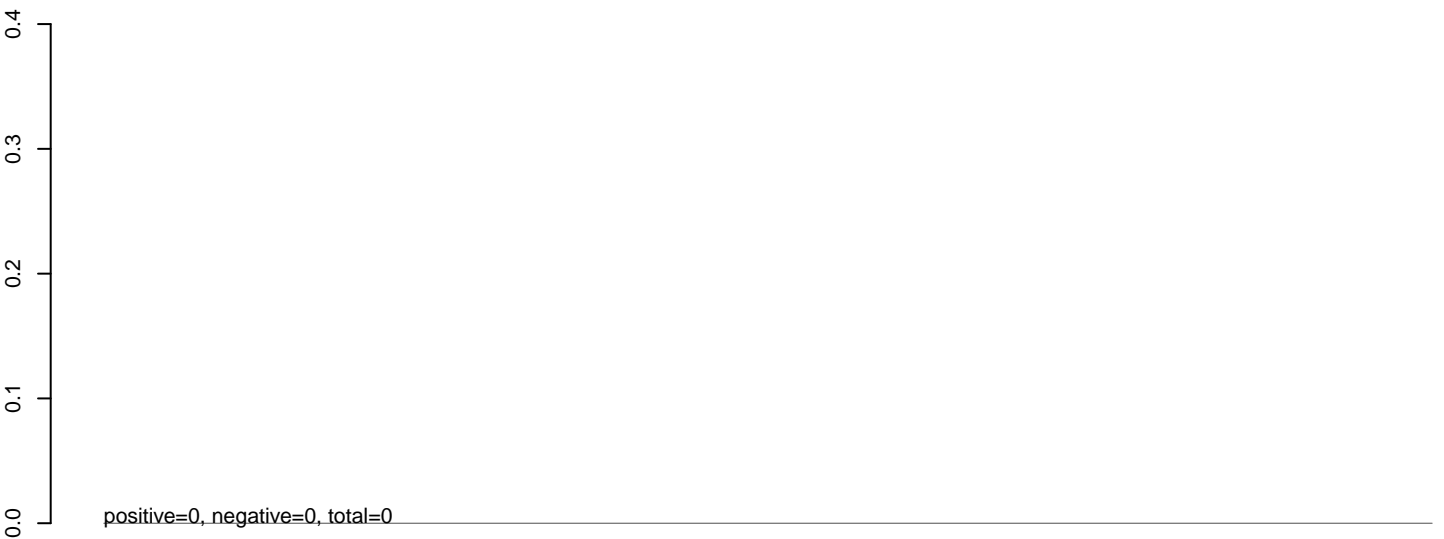
protein_coding_hypothetical_proteins_matrix_hypothetical_proteins_glycoprotein_hypothetical_proteins_and_polymerase_genes:1-15721

0 5000 10000 15000

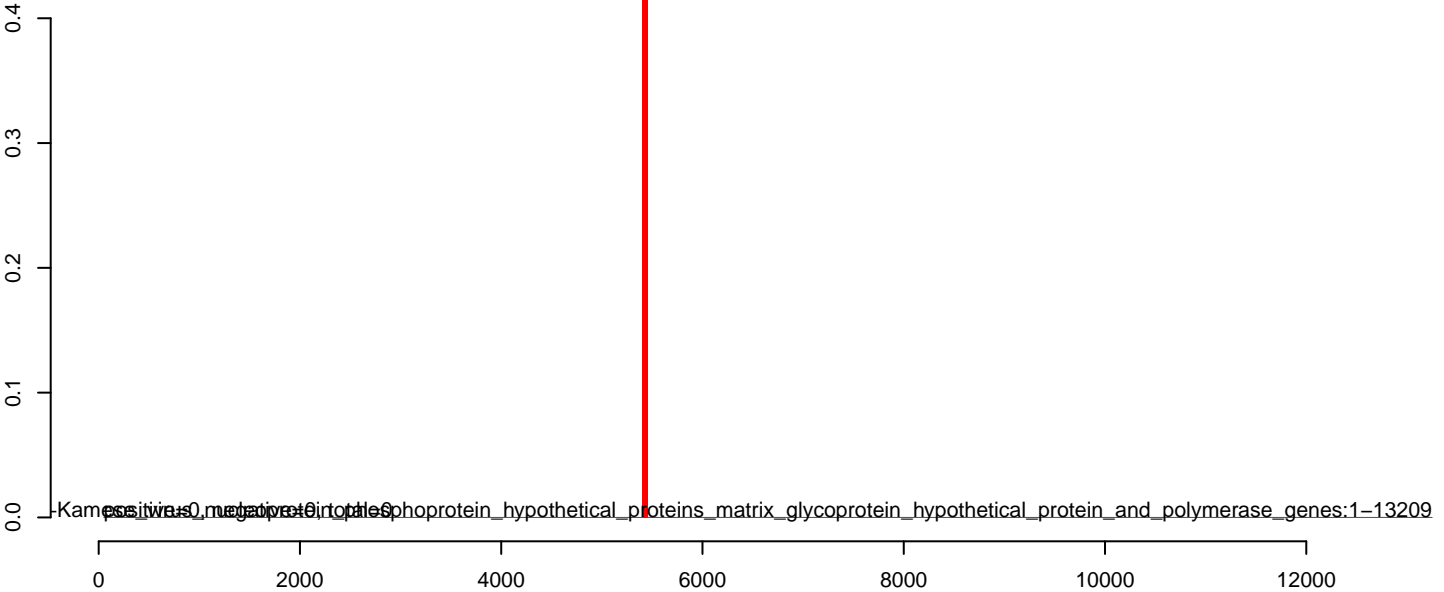
AeAeg_CCL.125cells_CHIKV.18_23.rep



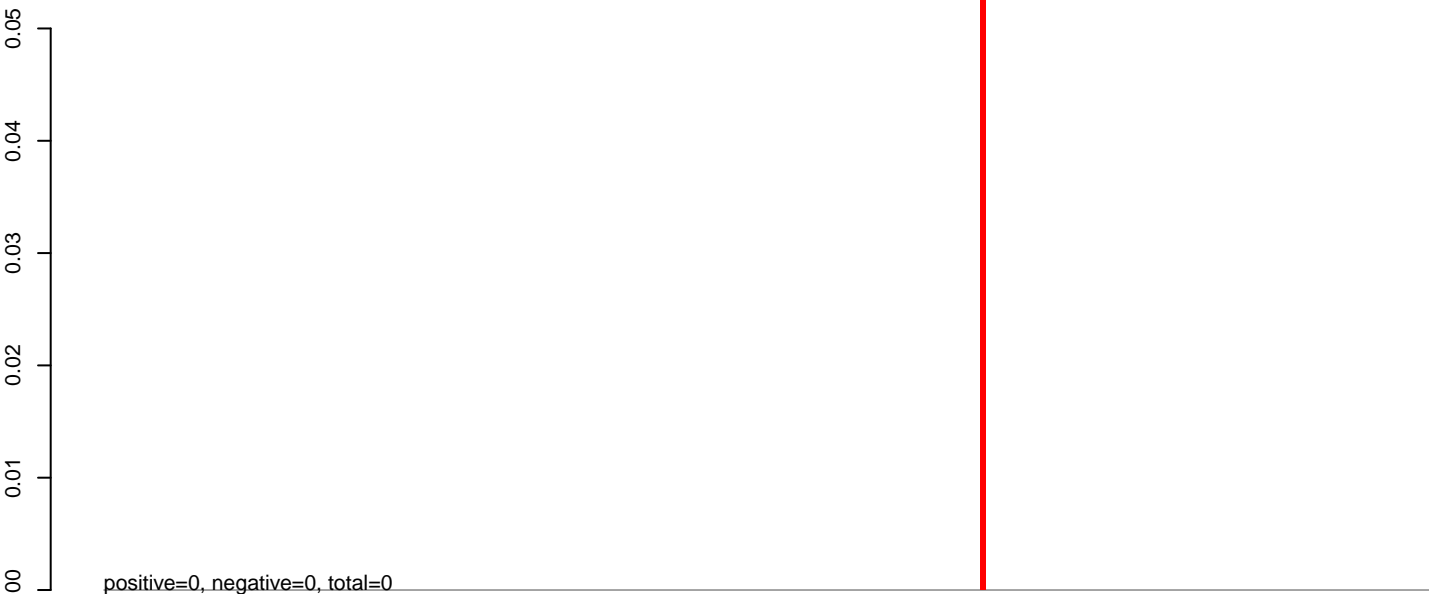
AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep



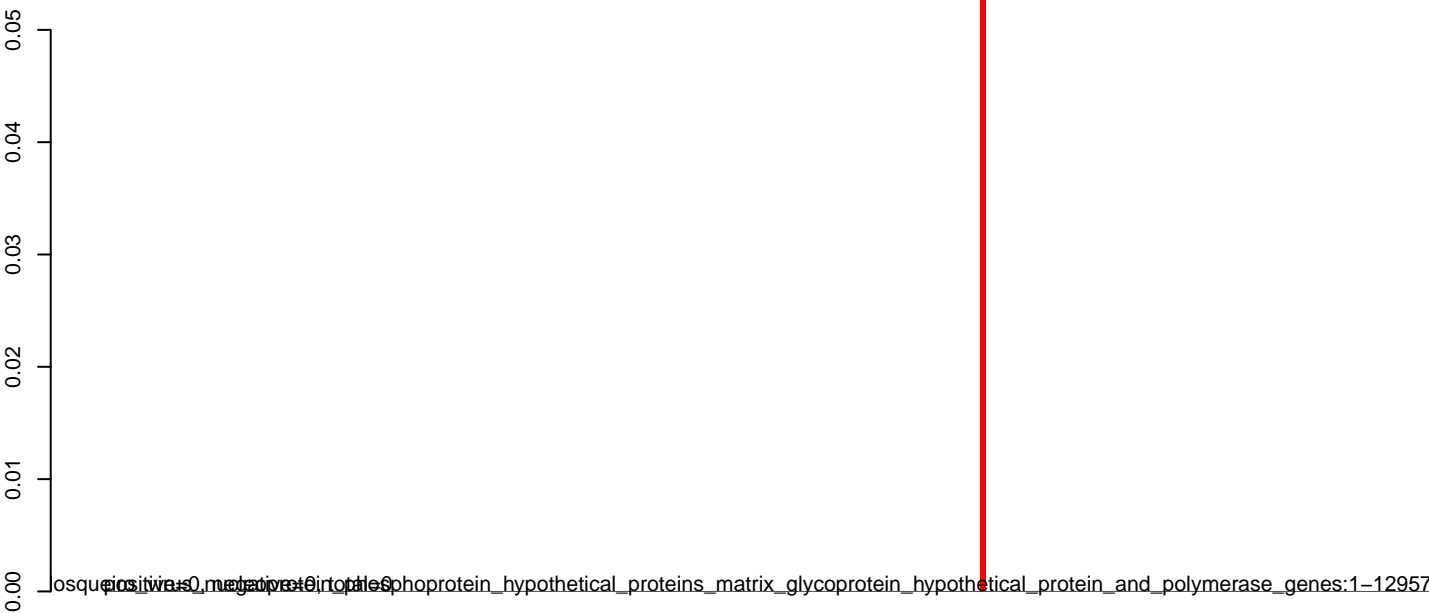
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

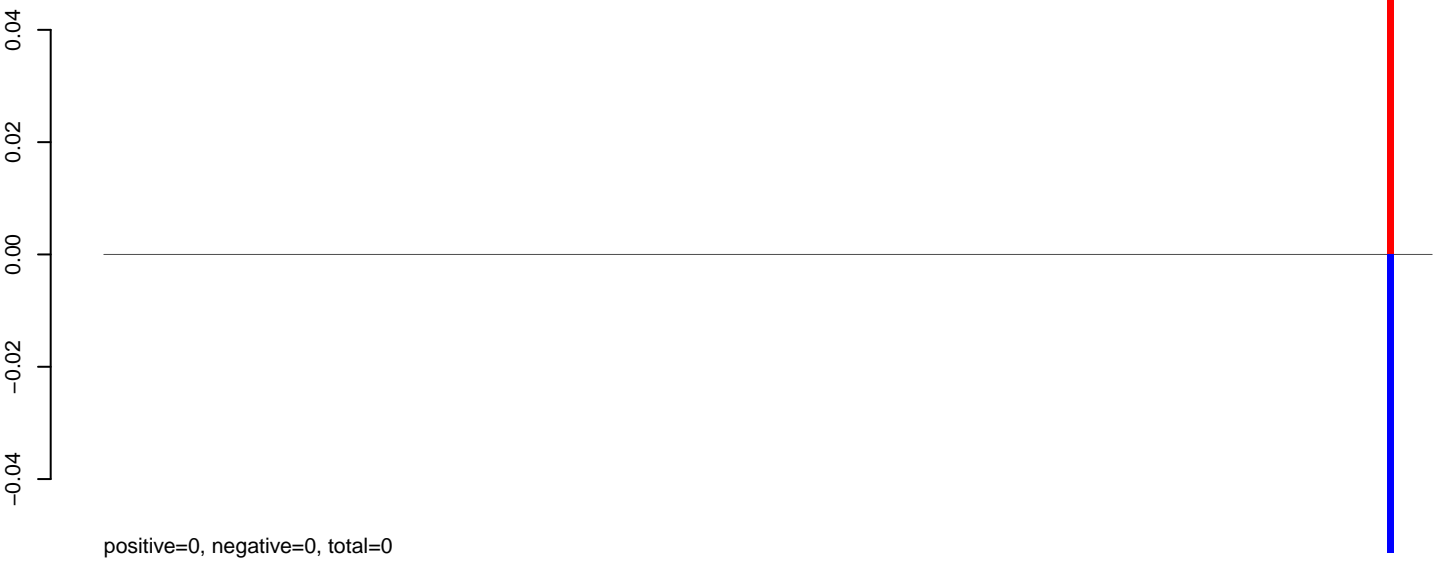


AeAeg_CCL.125cells_CHIKV.rep

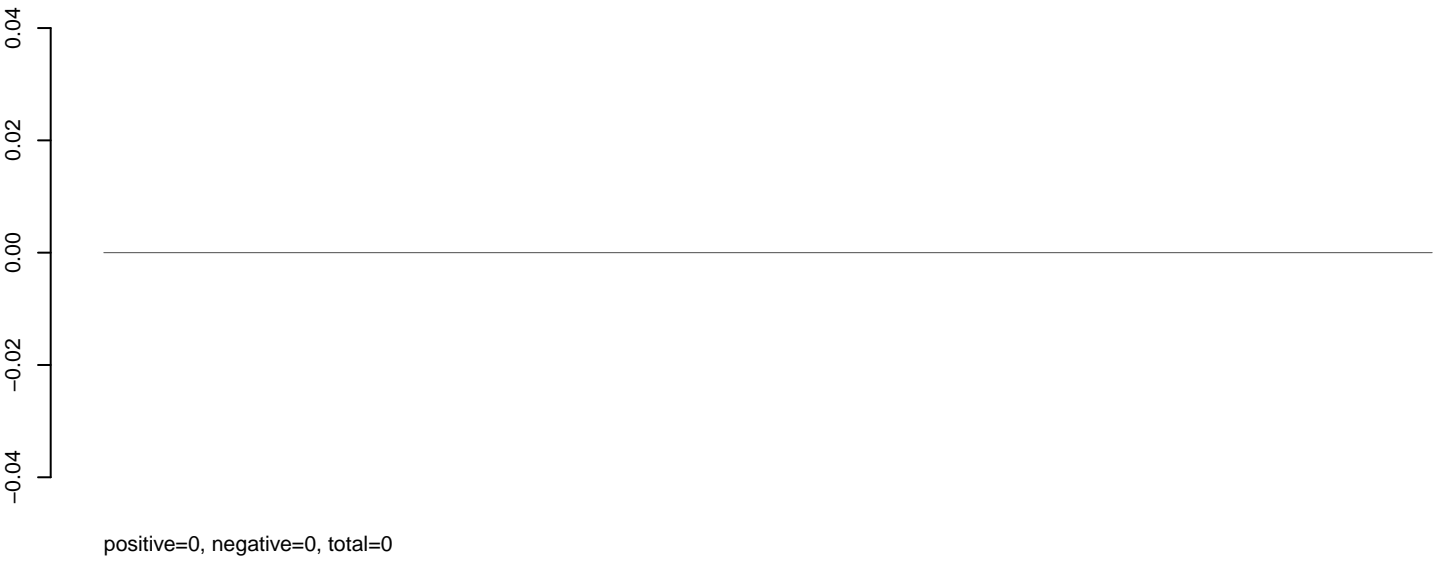


osquepinosin_megastrolinone_phosphoprotein_hypothetical_proteins_matrix_glycoprotein_hypothetical_protein_and_polymerase_genes:1-12957

AeAeg_CCL.125cells_CHIKV.18_23.rep



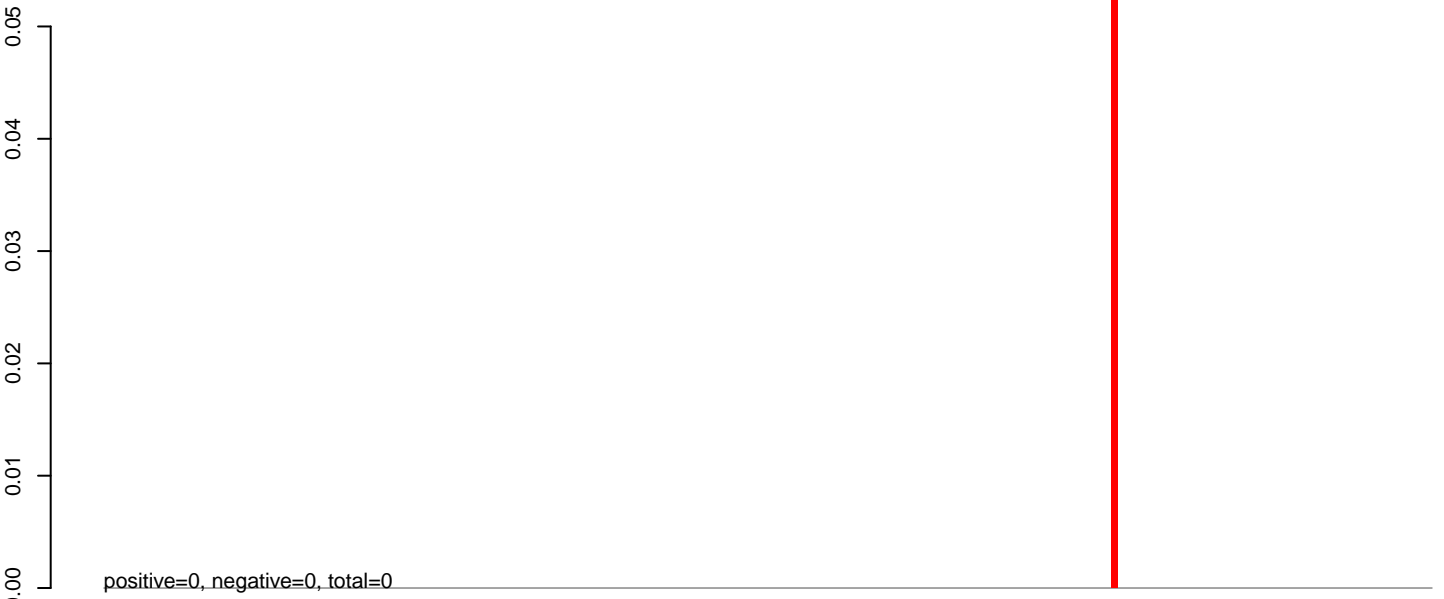
AeAeg_CCL.125cells_CHIKV.24_35.rep



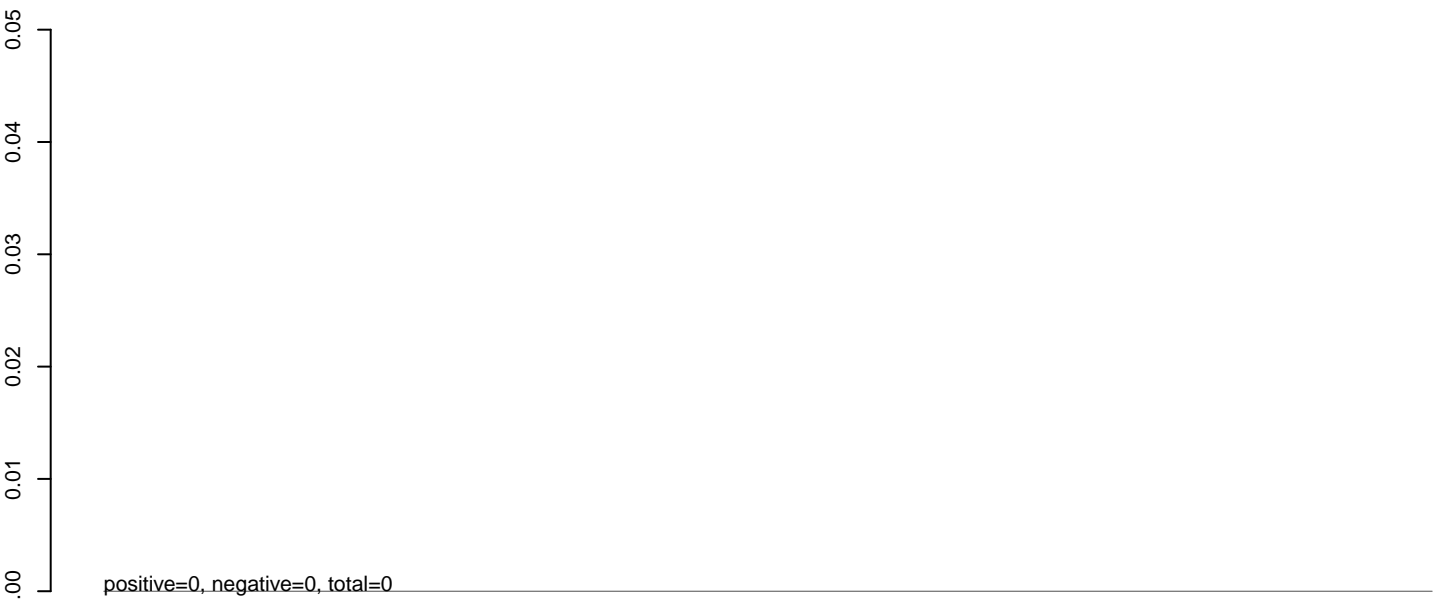
AeAeg_CCL.125cells_CHIKV.rep



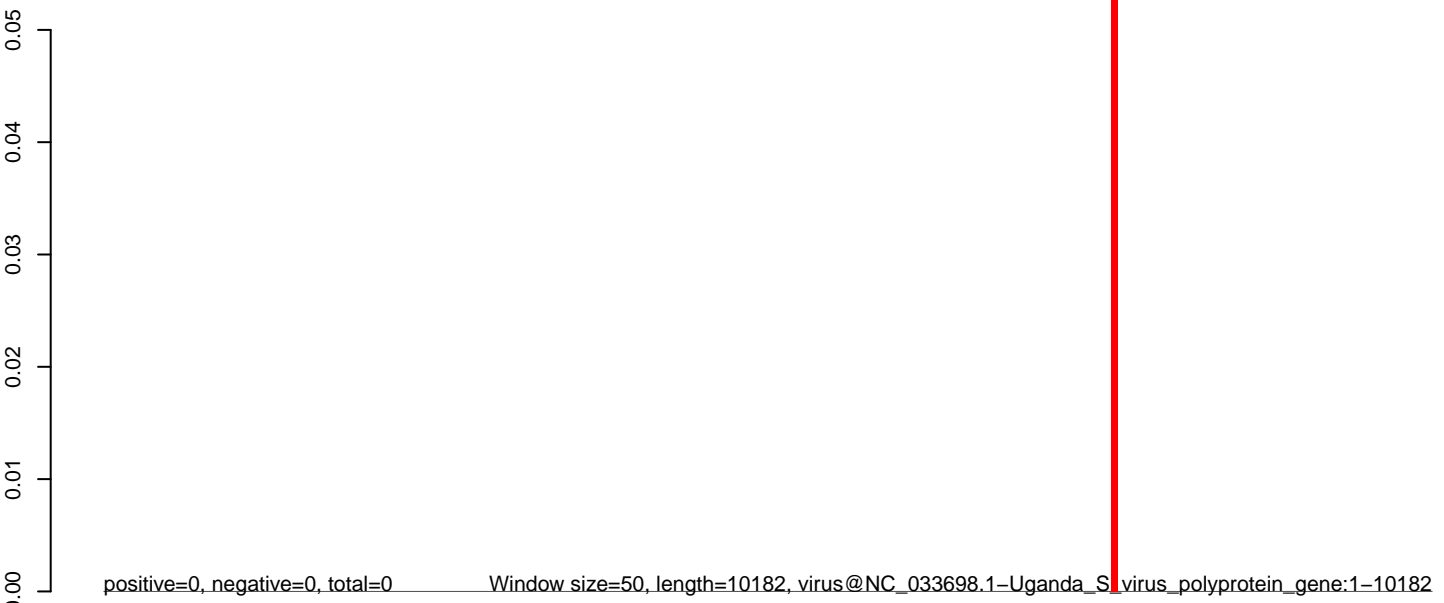
AeAeg_CCL.125cells_CHIKV.18_23.rep



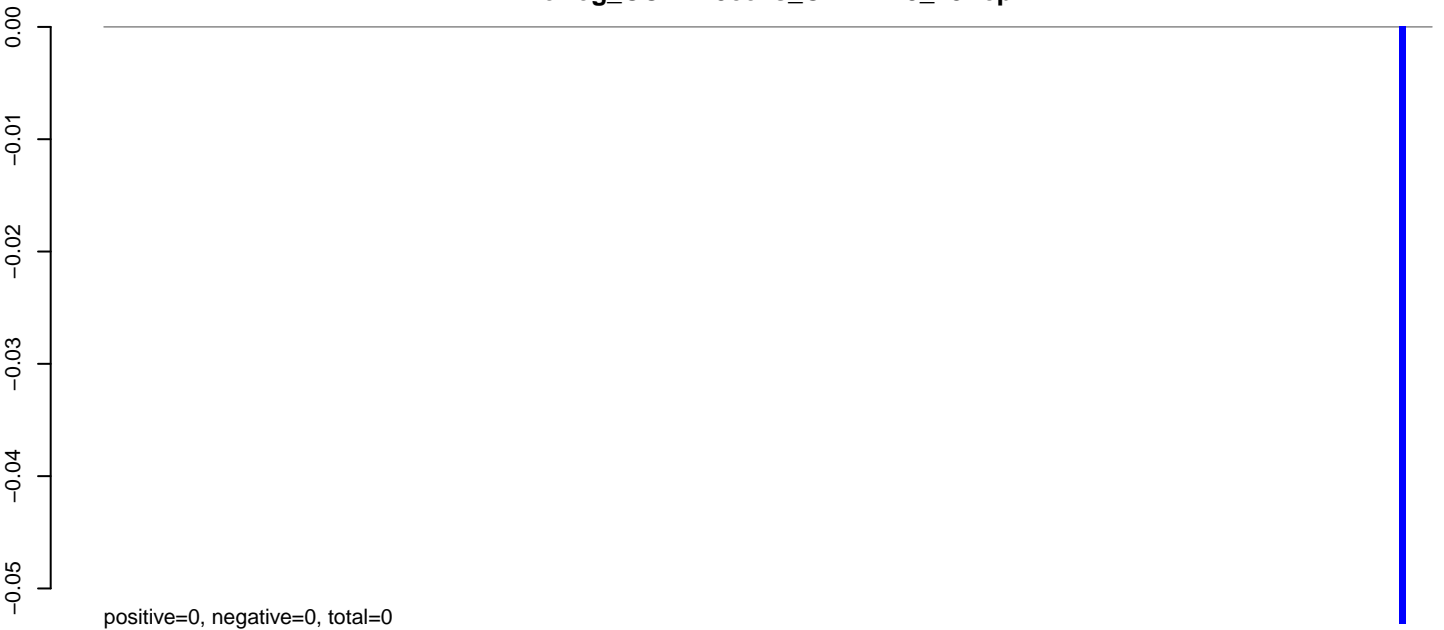
AeAeg_CCL.125cells_CHIKV.24_35.rep



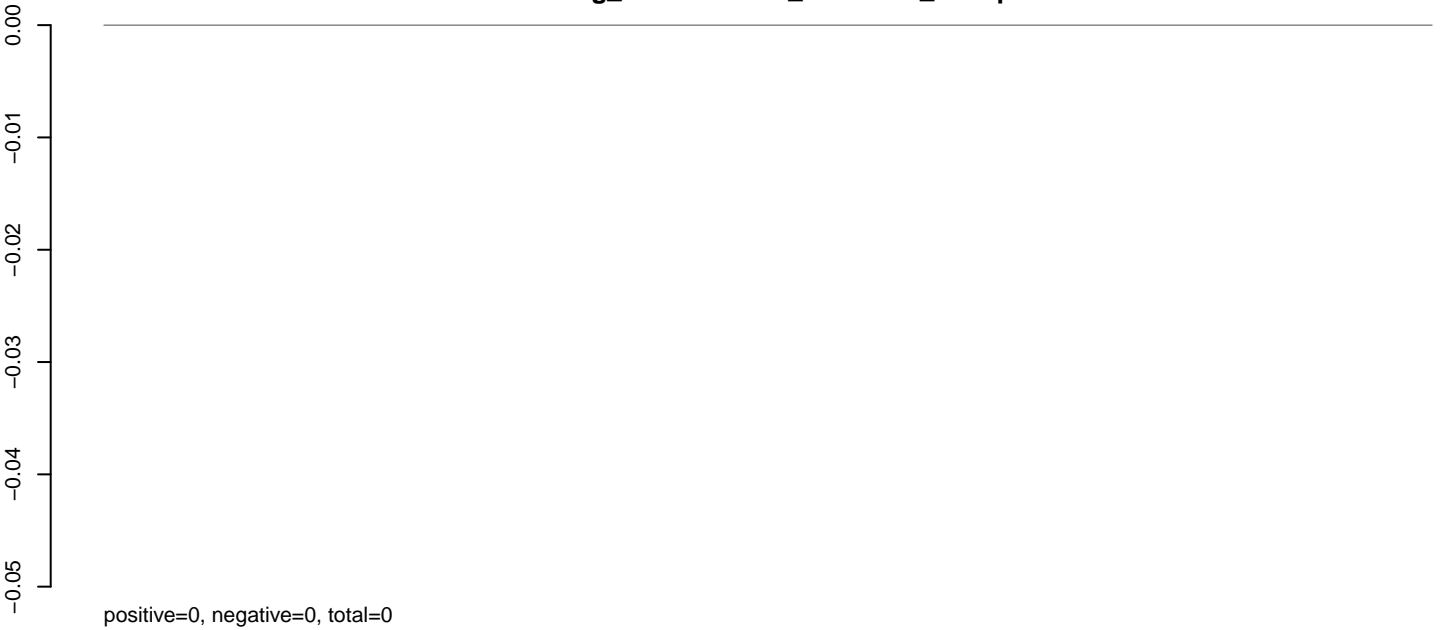
AeAeg_CCL.125cells_CHIKV.rep



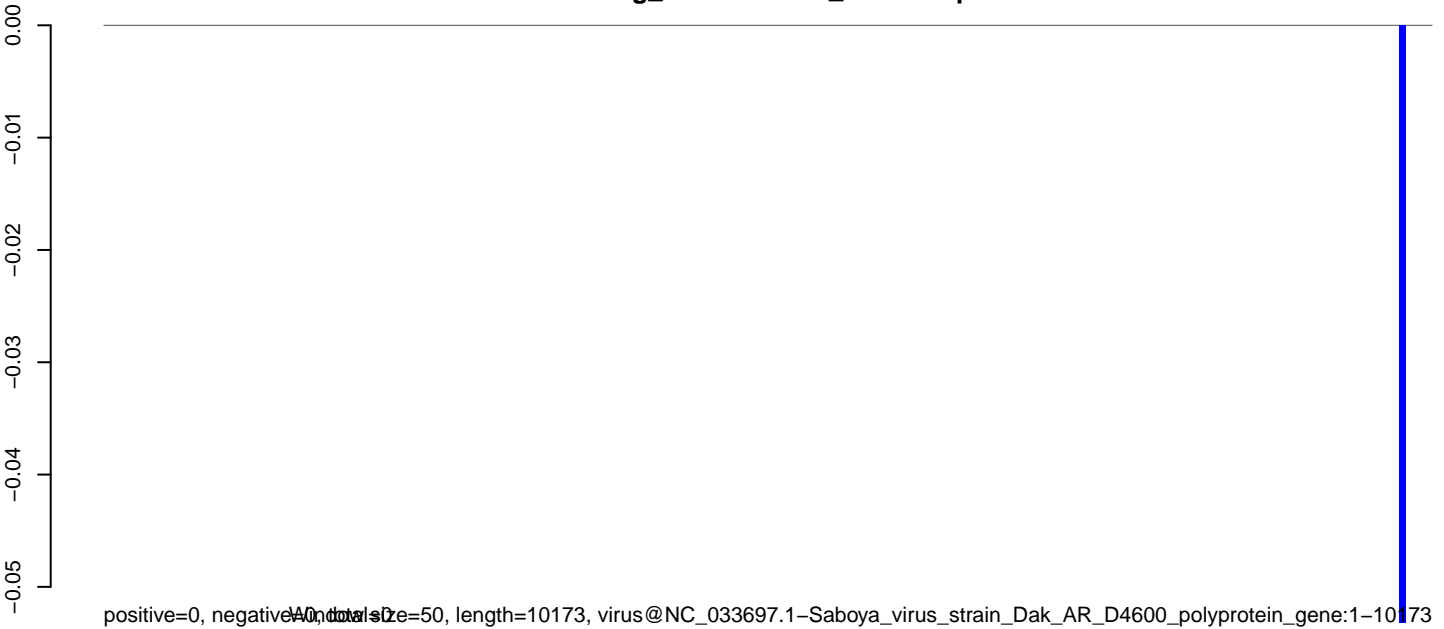
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



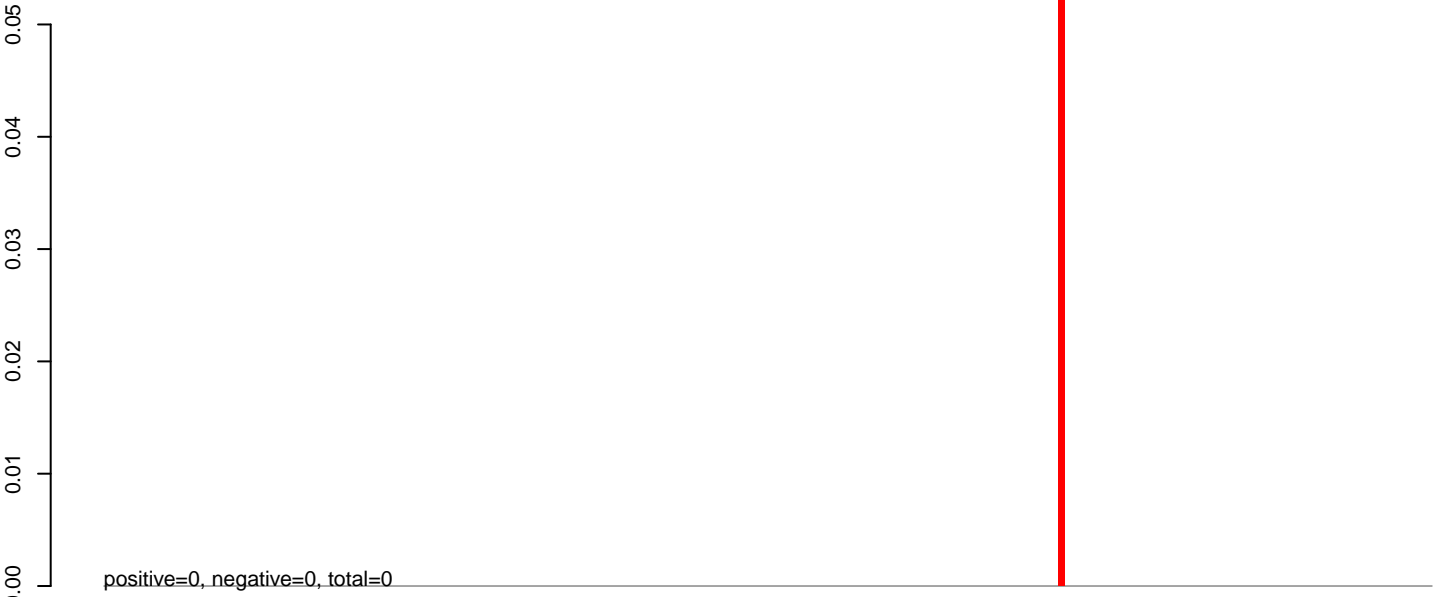
AeAeg_CCL.125cells_CHIKV.rep



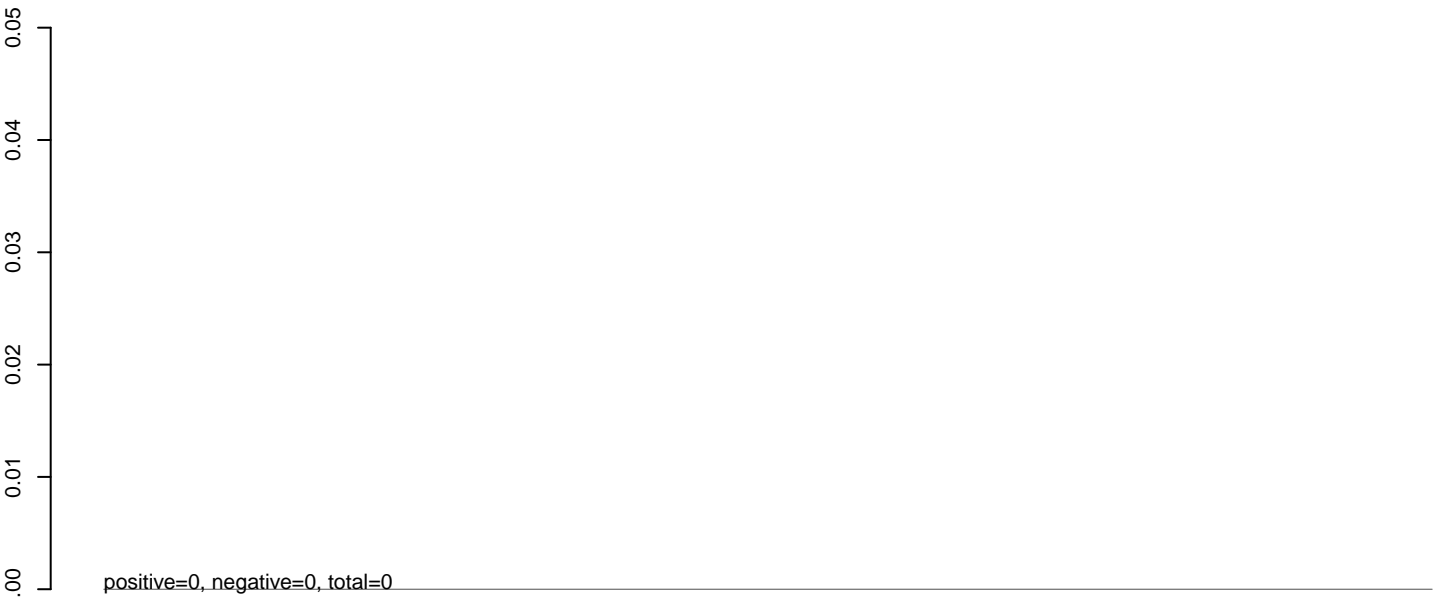
positive=0, negative=0, total=0, length=10173, virus@NC_033697.1-Saboya_virus_strain_Dak_AR_D4600_polyprotein_gene:1-10173

0 2000 4000 6000 8000 10000

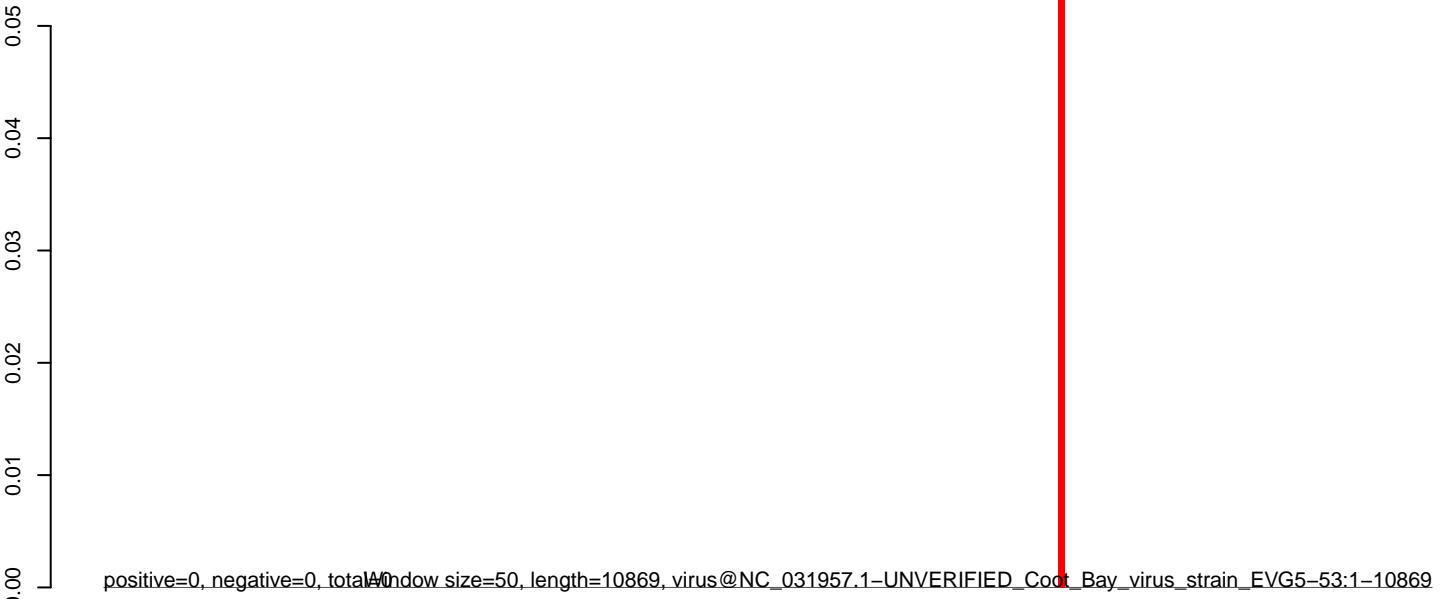
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep

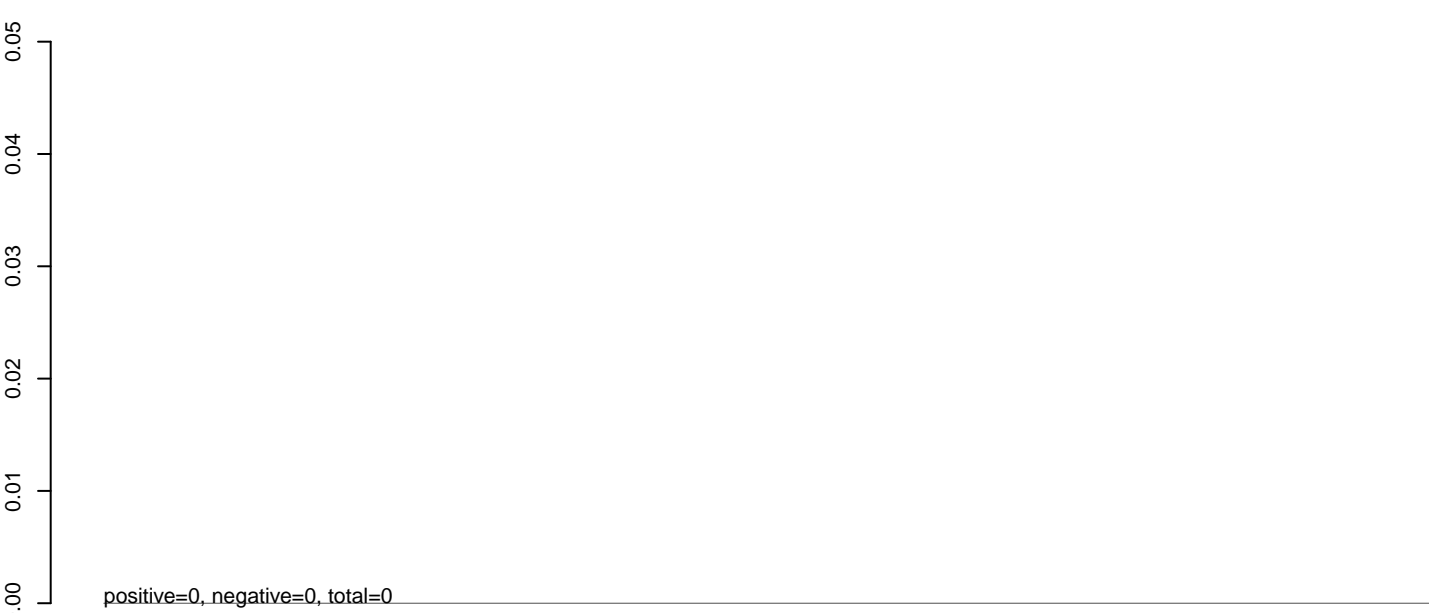


Window size=50, length=10869, virus@NC_031957.1-UNVERIFIED_Coot_Bay_virus_strain_EVG5-53:1-10869

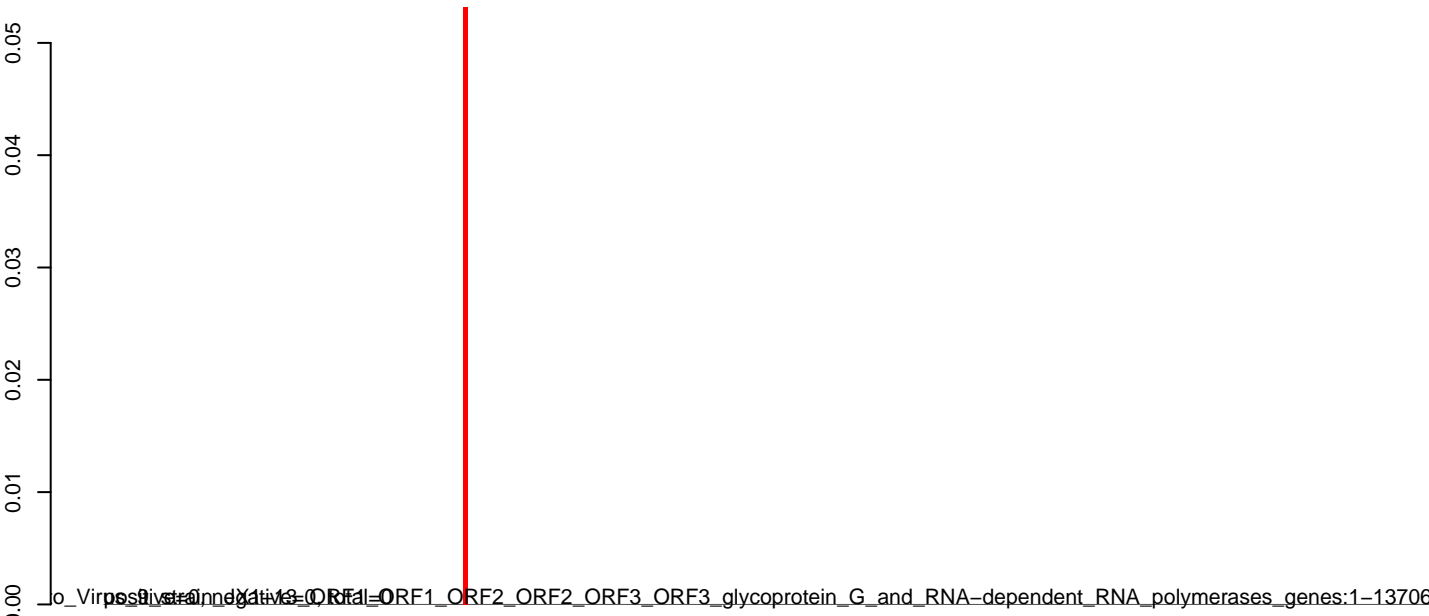
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



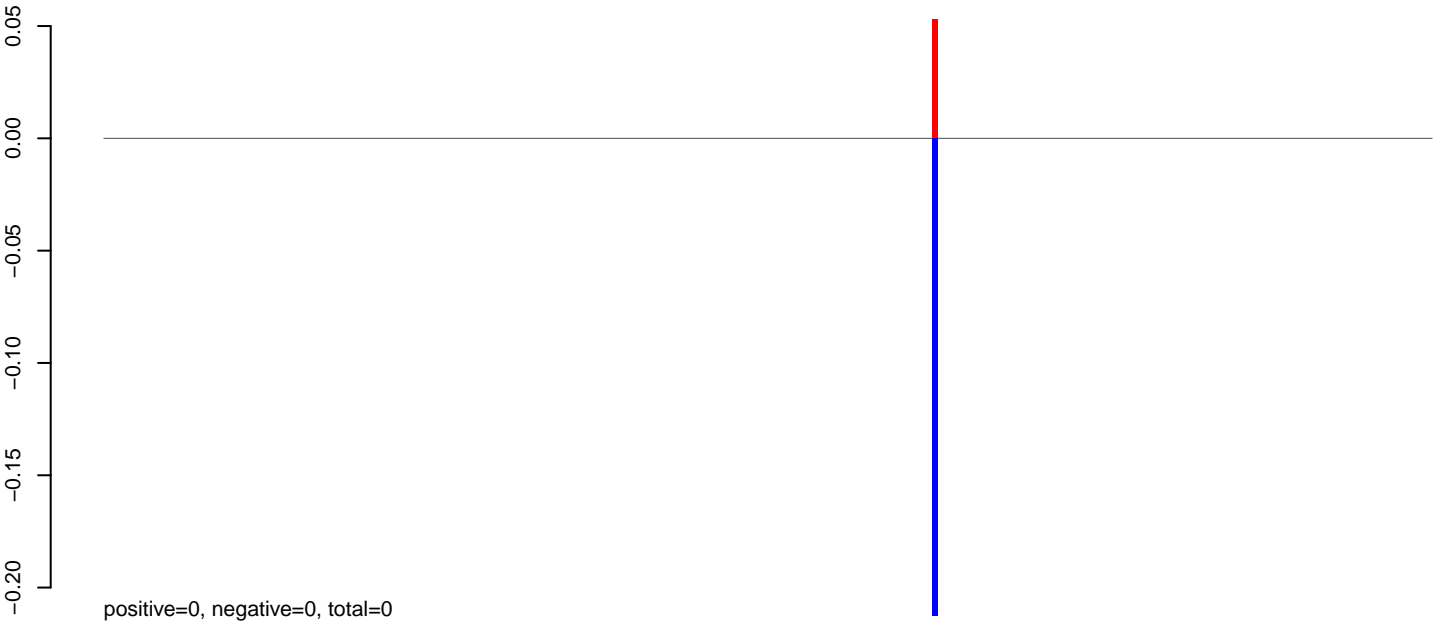
AeAeg_CCL.125cells_CHIKV.rep



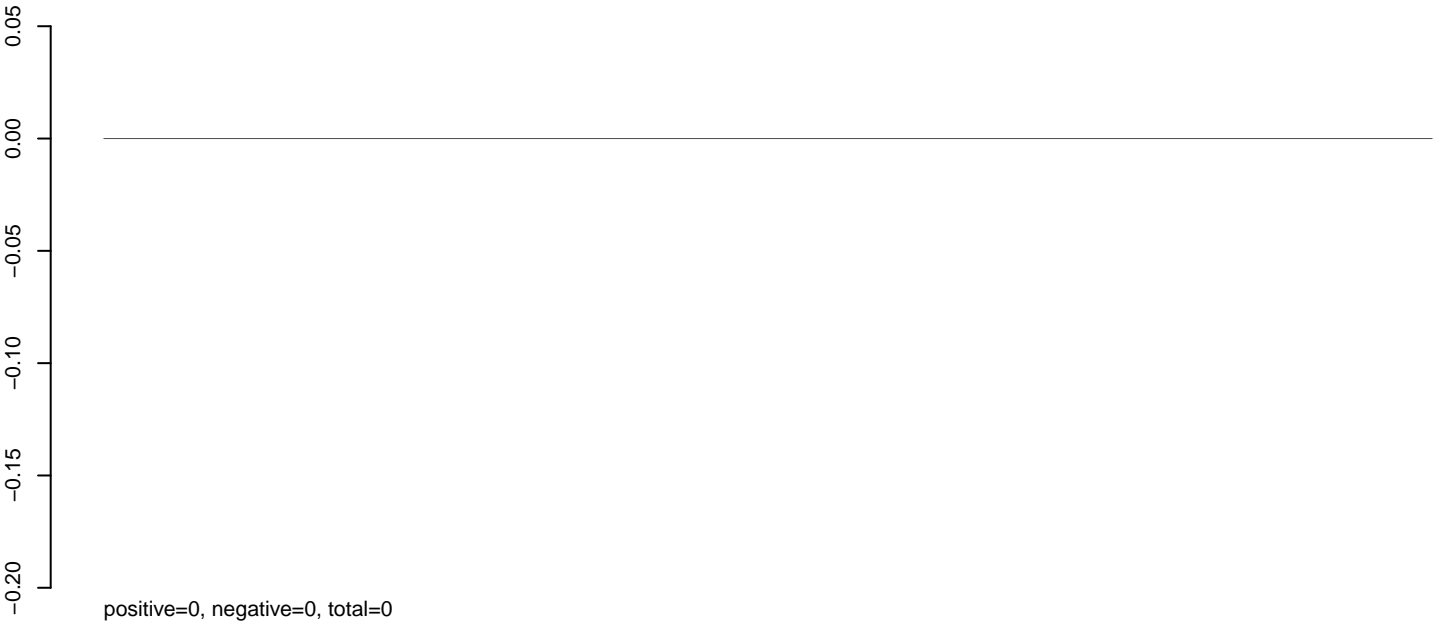
o_Virus_innogy113_ORF1_ORF1_ORF2_ORF2_ORF3_ORF3_glycoprotein_G_and_RNA-dependent_RNA_polymerases_genes:1-13706

0 2000 4000 6000 8000 10000 12000 14000

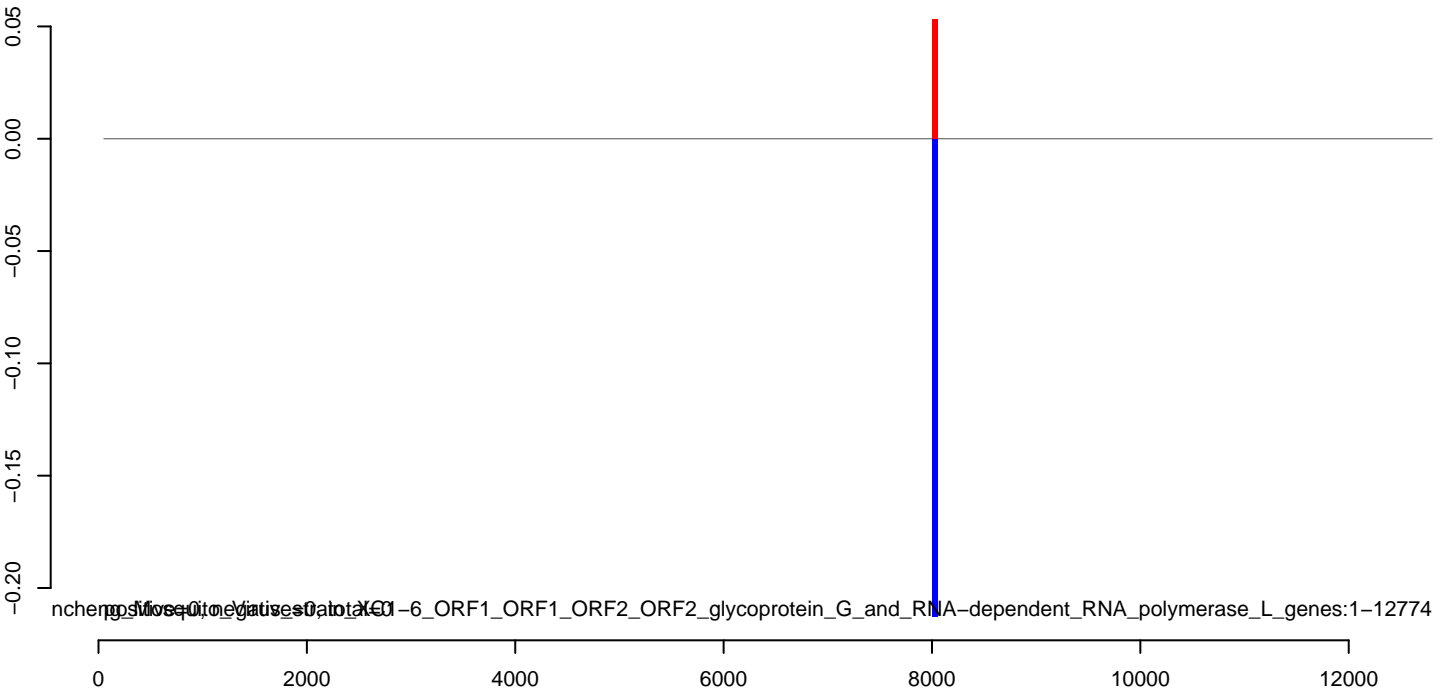
AeAeg_CCL.125cells_CHIKV.18_23.rep



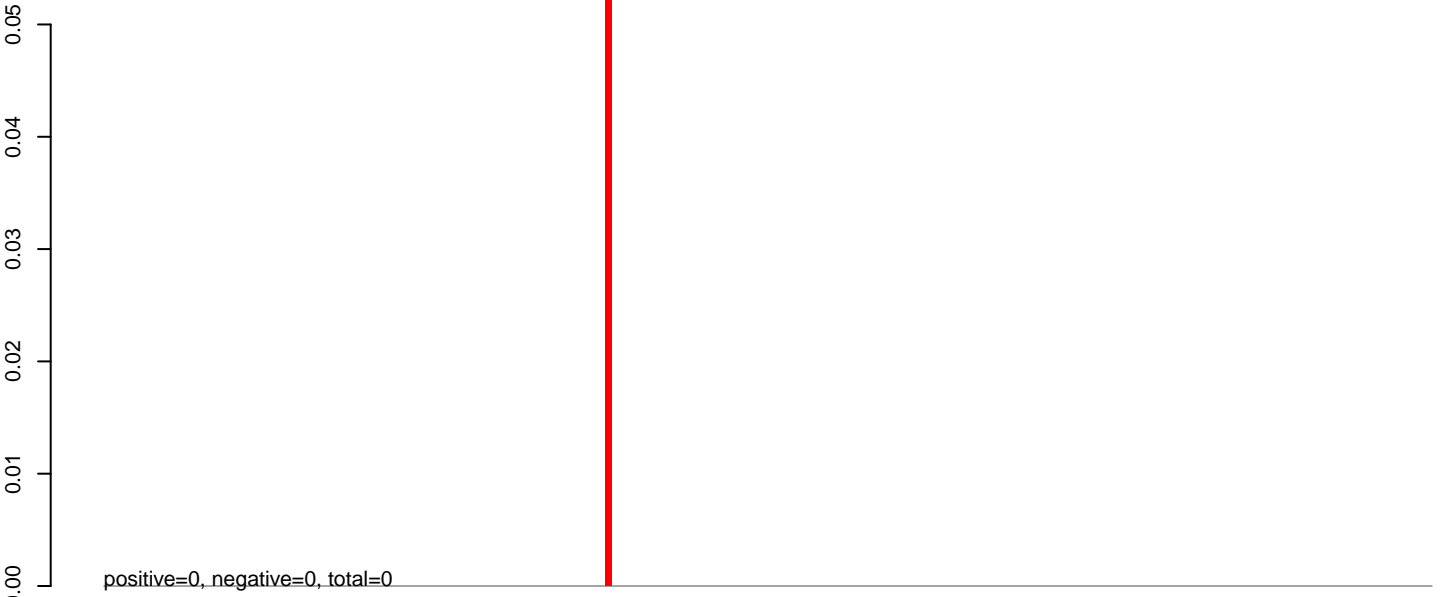
AeAeg_CCL.125cells_CHIKV.24_35.rep



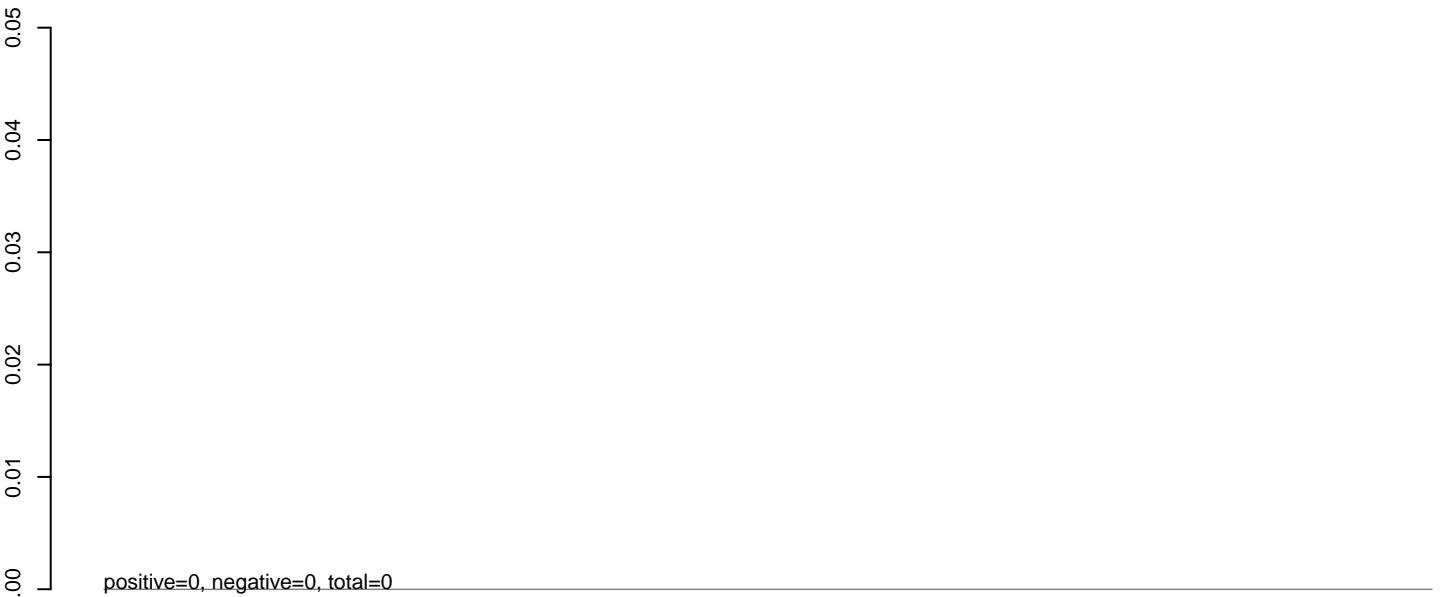
AeAeg_CCL.125cells_CHIKV.rep



AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



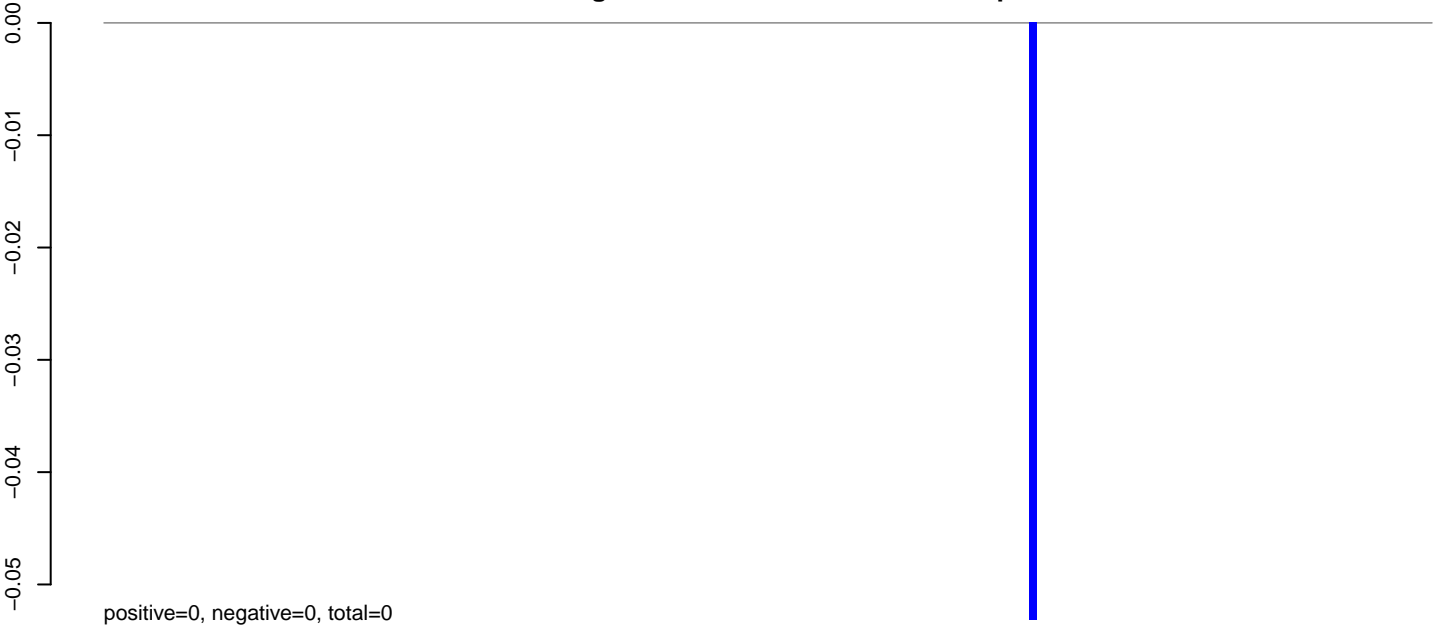
AeAeg_CCL.125cells_CHIKV.rep



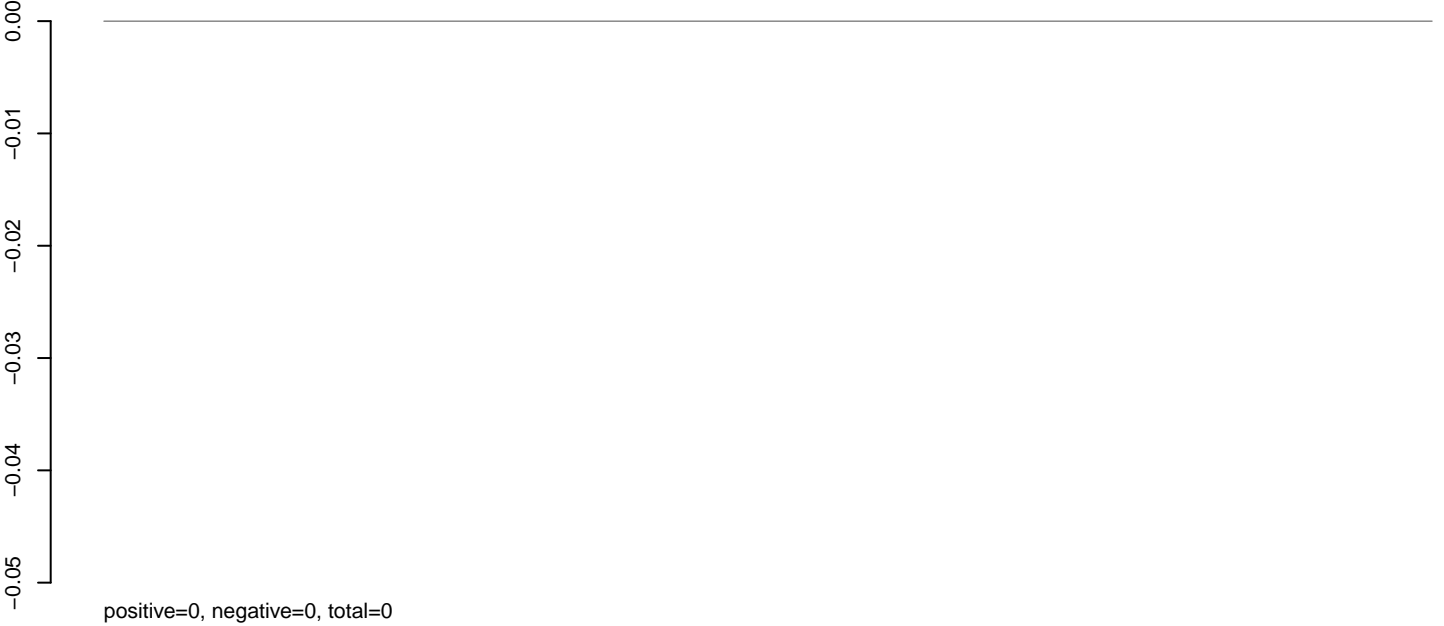
Window size=50, length=10206, virus@NC_030289.1-Edge_Hill_virus_strain_YMP_48_polyprotein_gene:1-10206

0 2000 4000 6000 8000 10000

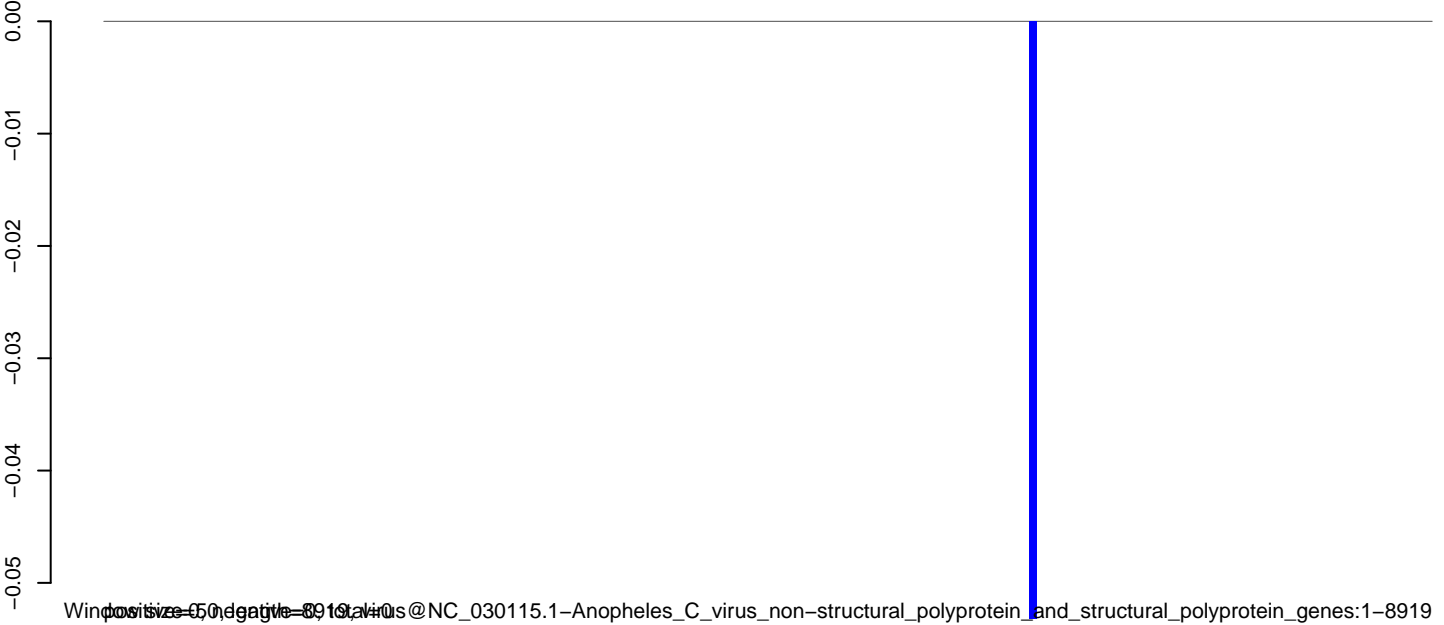
AeAeg_CCL.125cells_CHIKV.18_23.rep



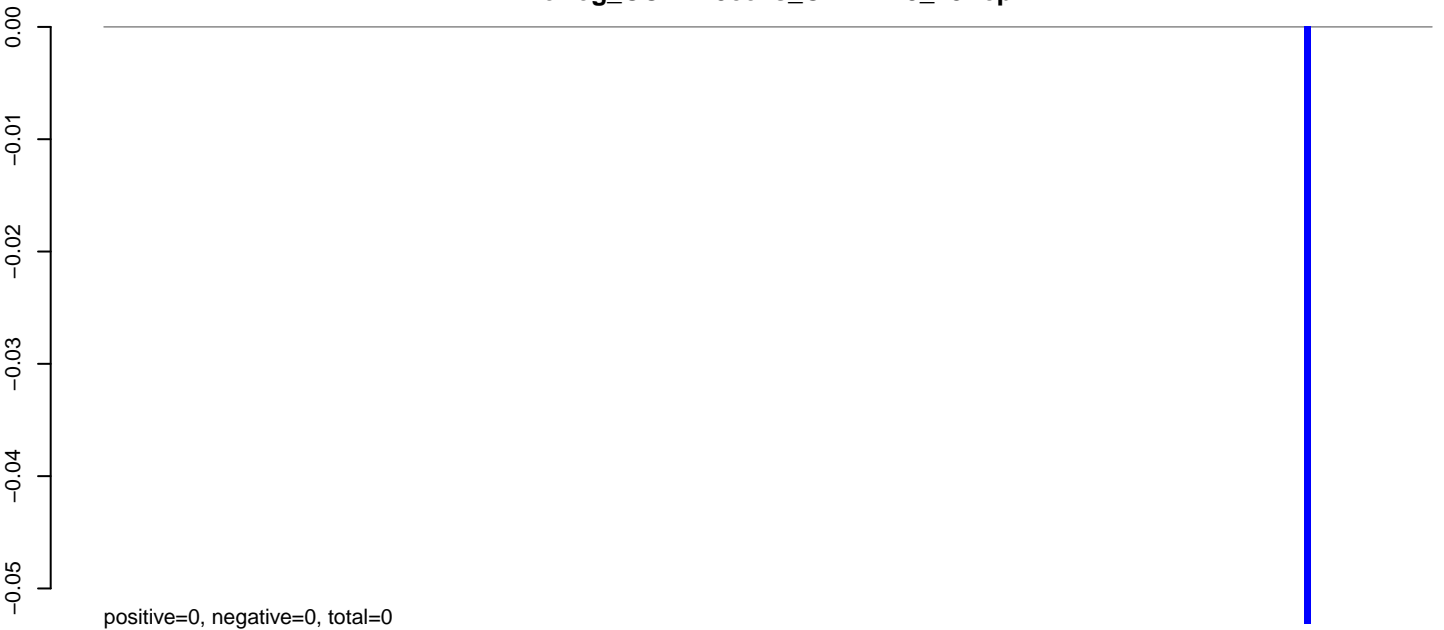
AeAeg_CCL.125cells_CHIKV.24_35.rep



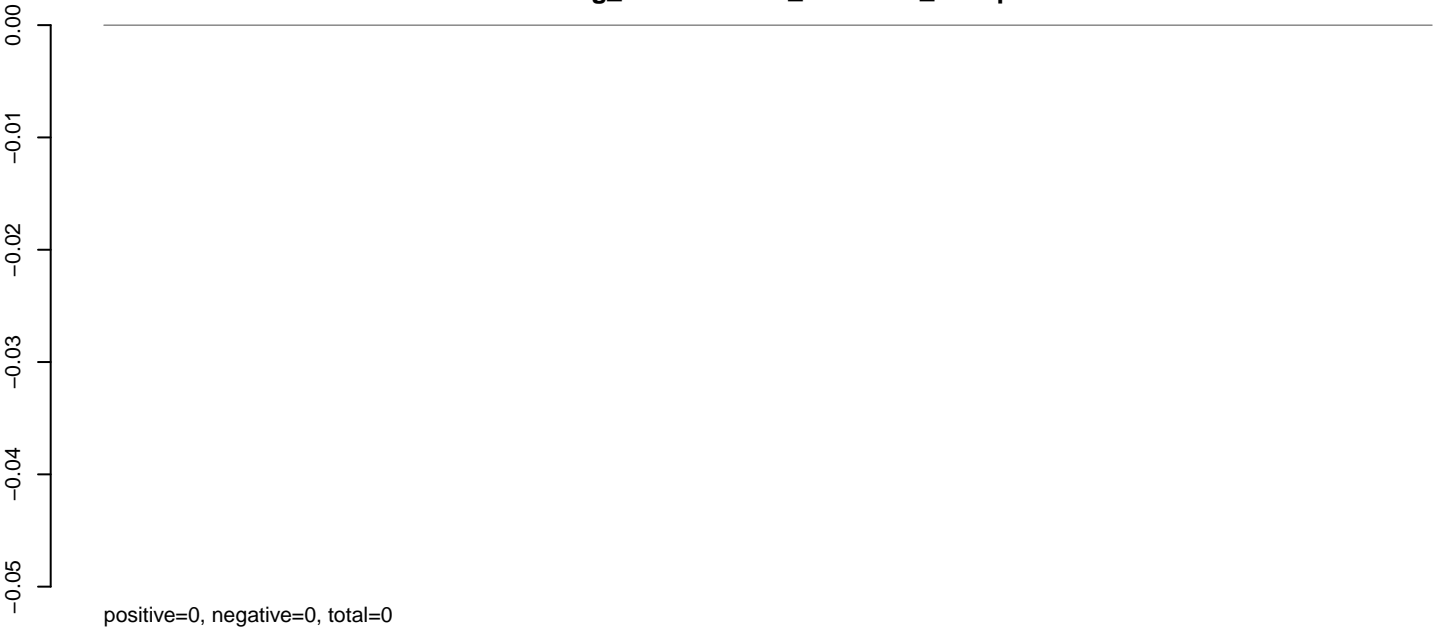
AeAeg_CCL.125cells_CHIKV.rep



AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

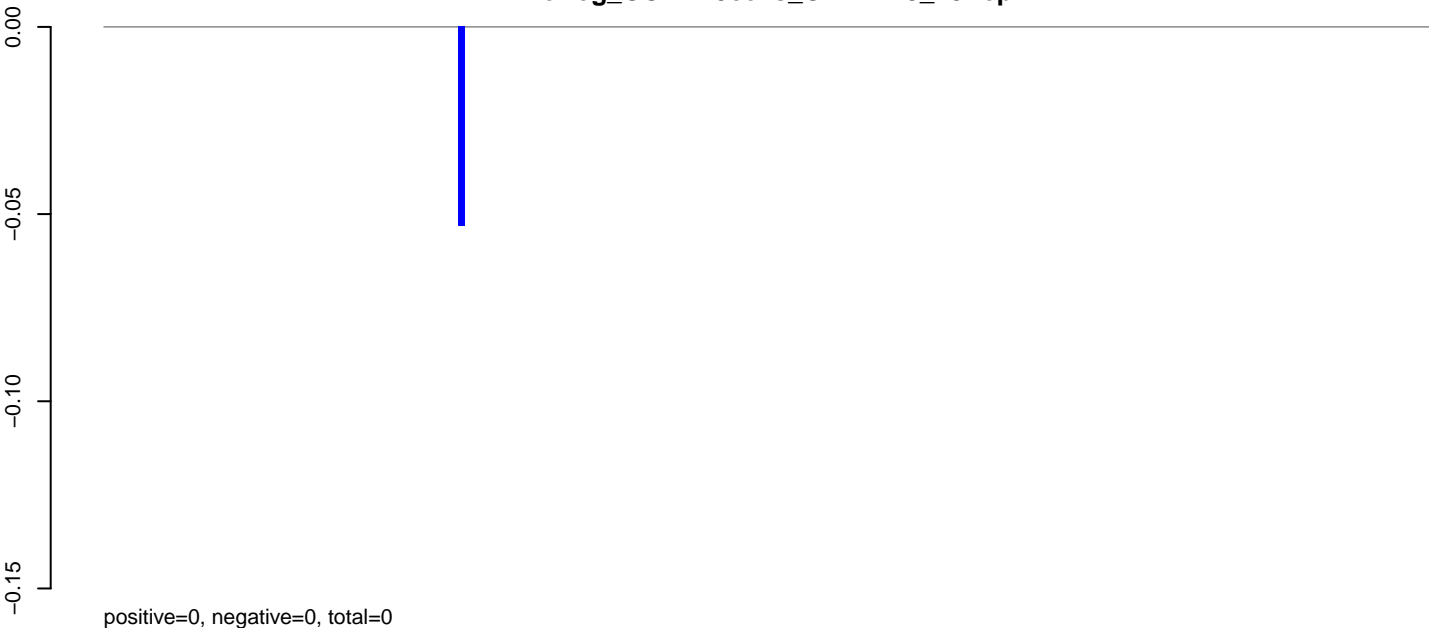


AeAeg_CCL.125cells_CHIKV.rep

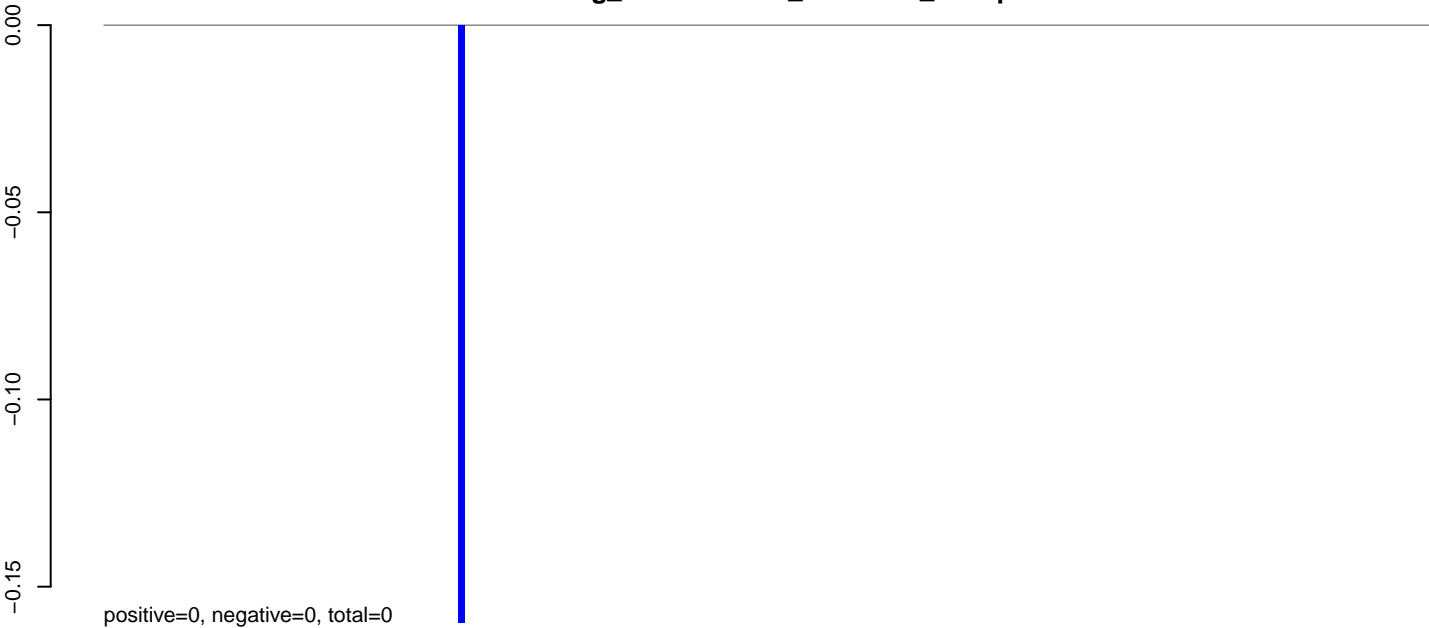


0 2000 4000 6000 8000 10000

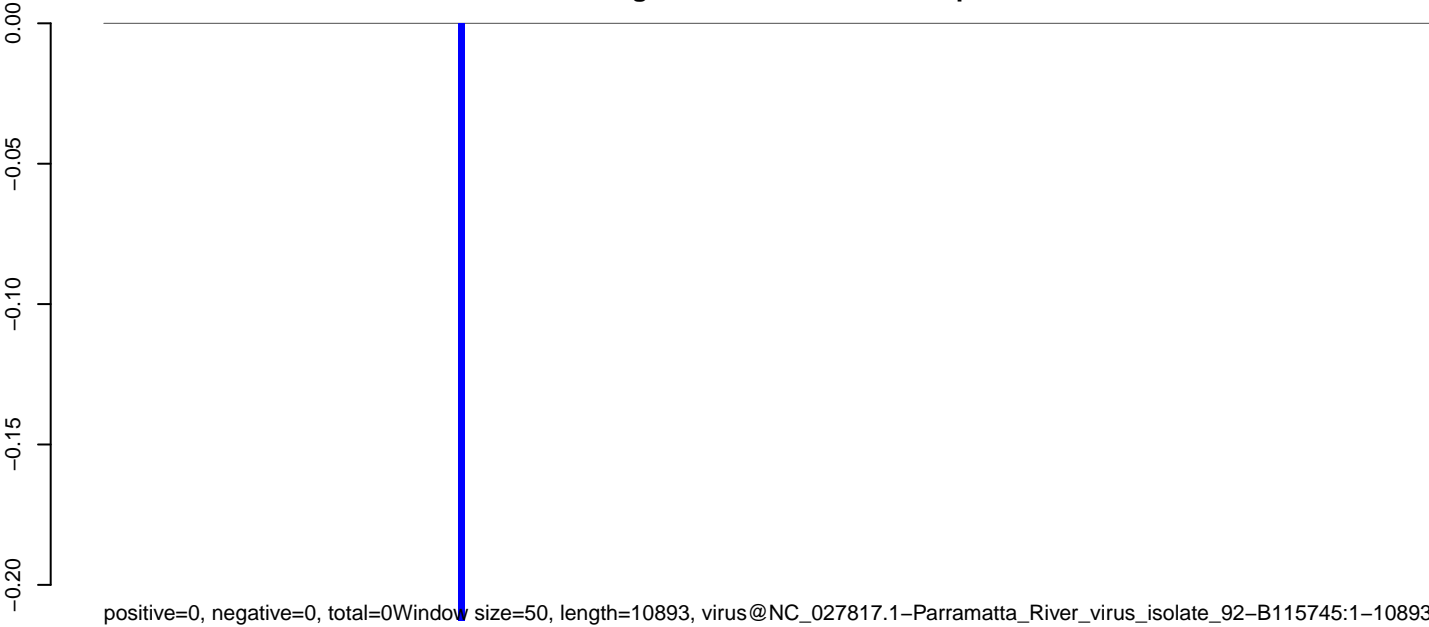
AeAeg_CCL.125cells_CHIKV.18_23.rep



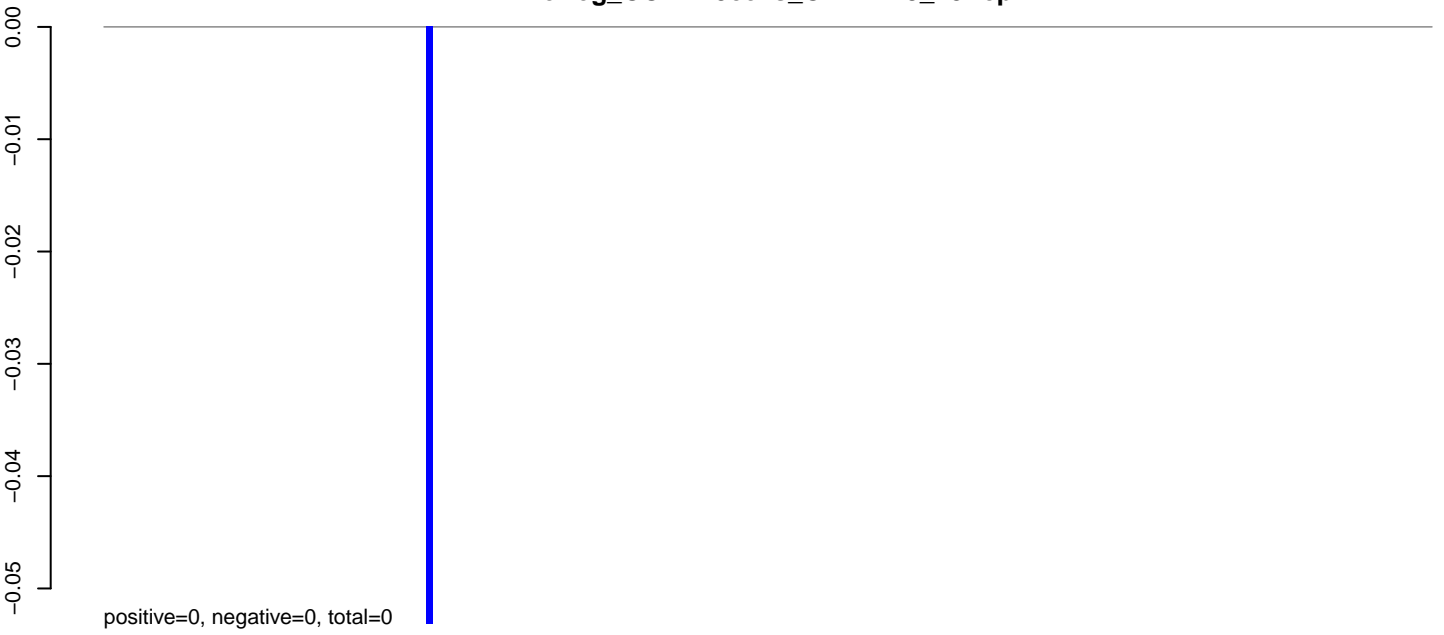
AeAeg_CCL.125cells_CHIKV.24_35.rep



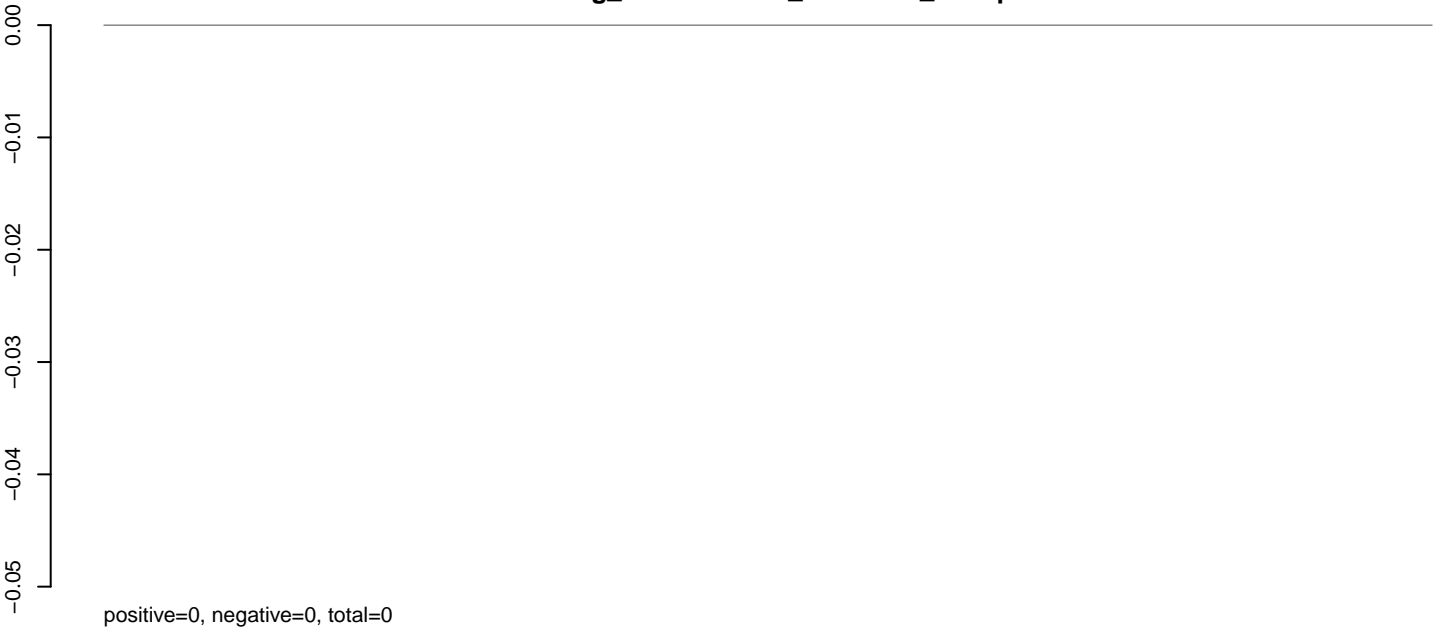
AeAeg_CCL.125cells_CHIKV.rep



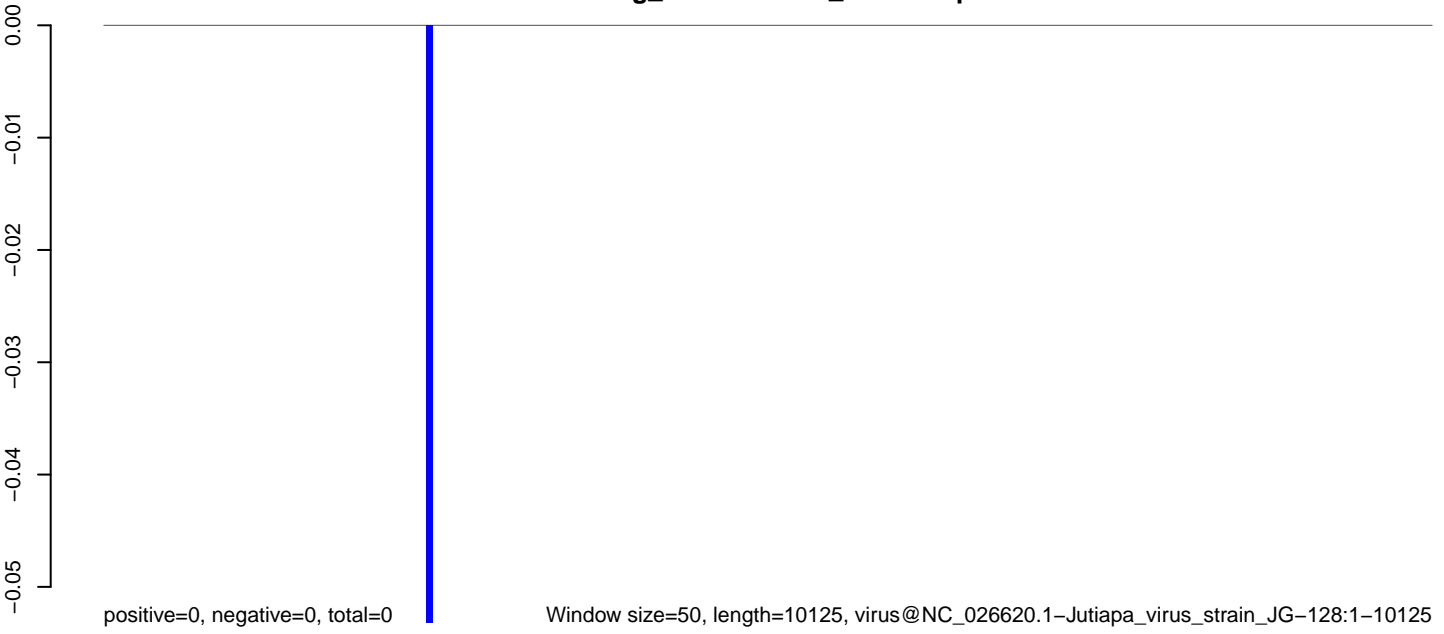
AeAeg_CCL.125cells_CHIKV.18_23.rep



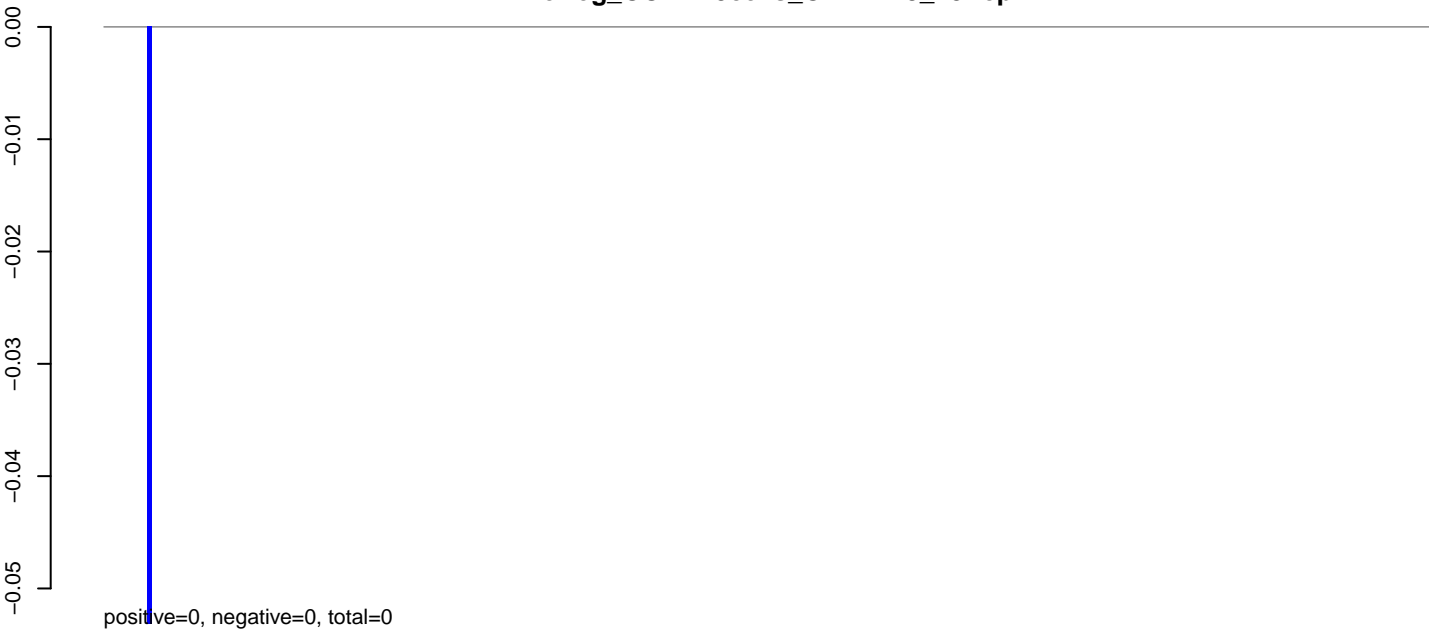
AeAeg_CCL.125cells_CHIKV.24_35.rep



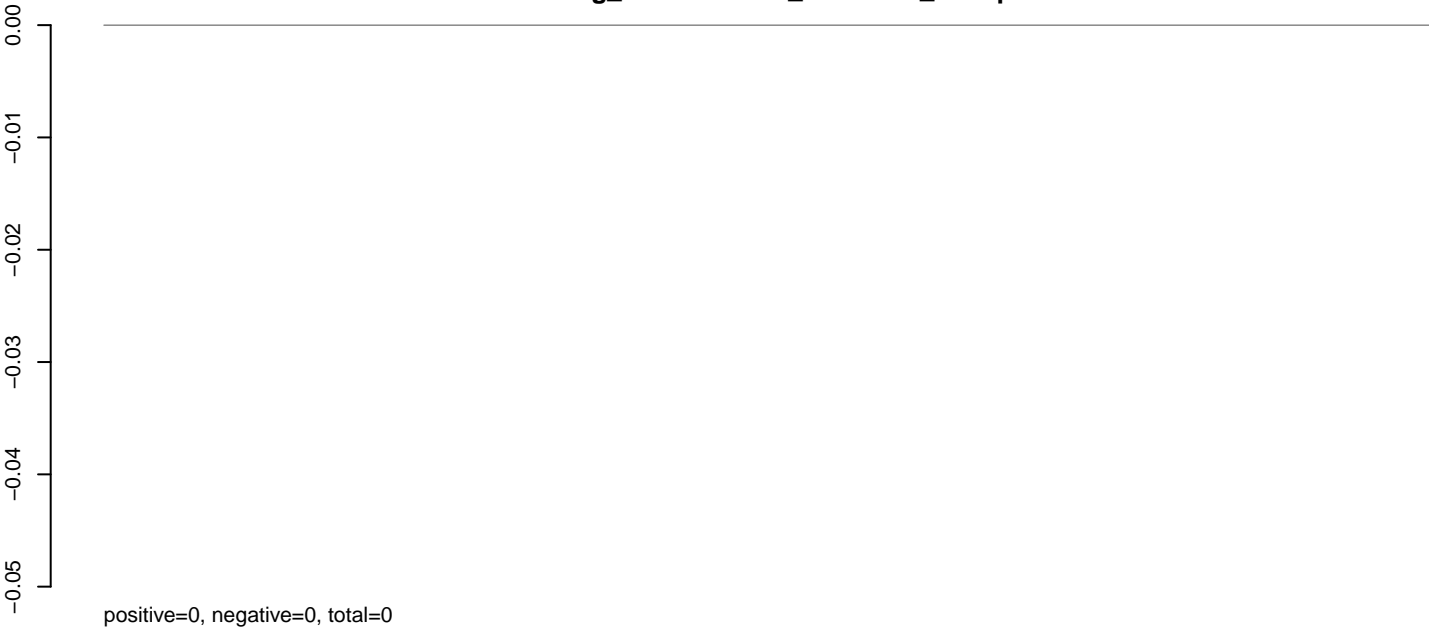
AeAeg_CCL.125cells_CHIKV.rep



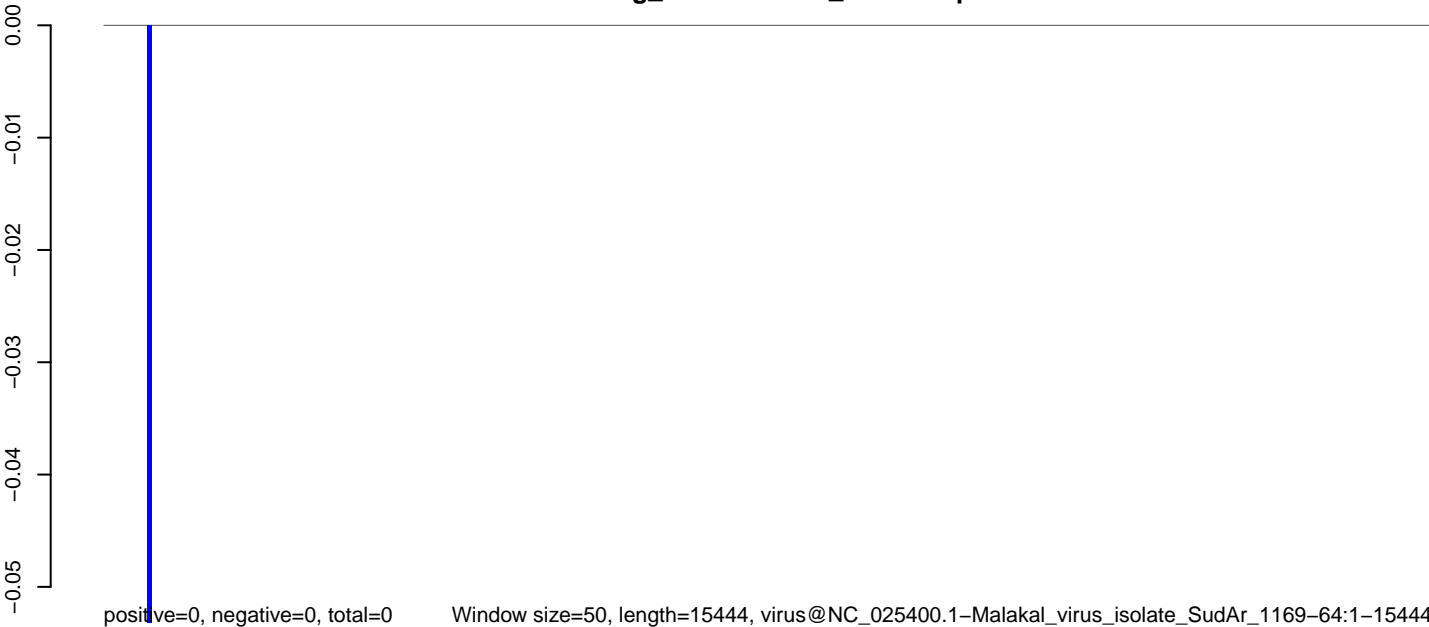
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

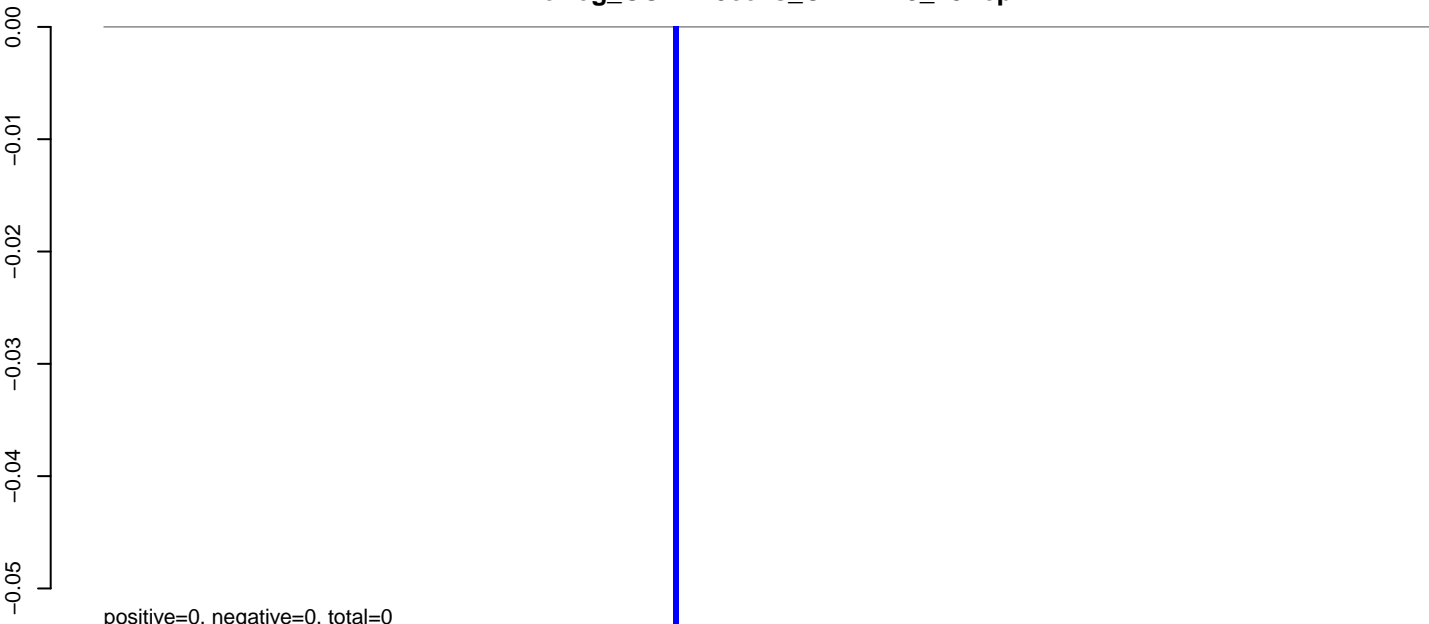


AeAeg_CCL.125cells_CHIKV.rep

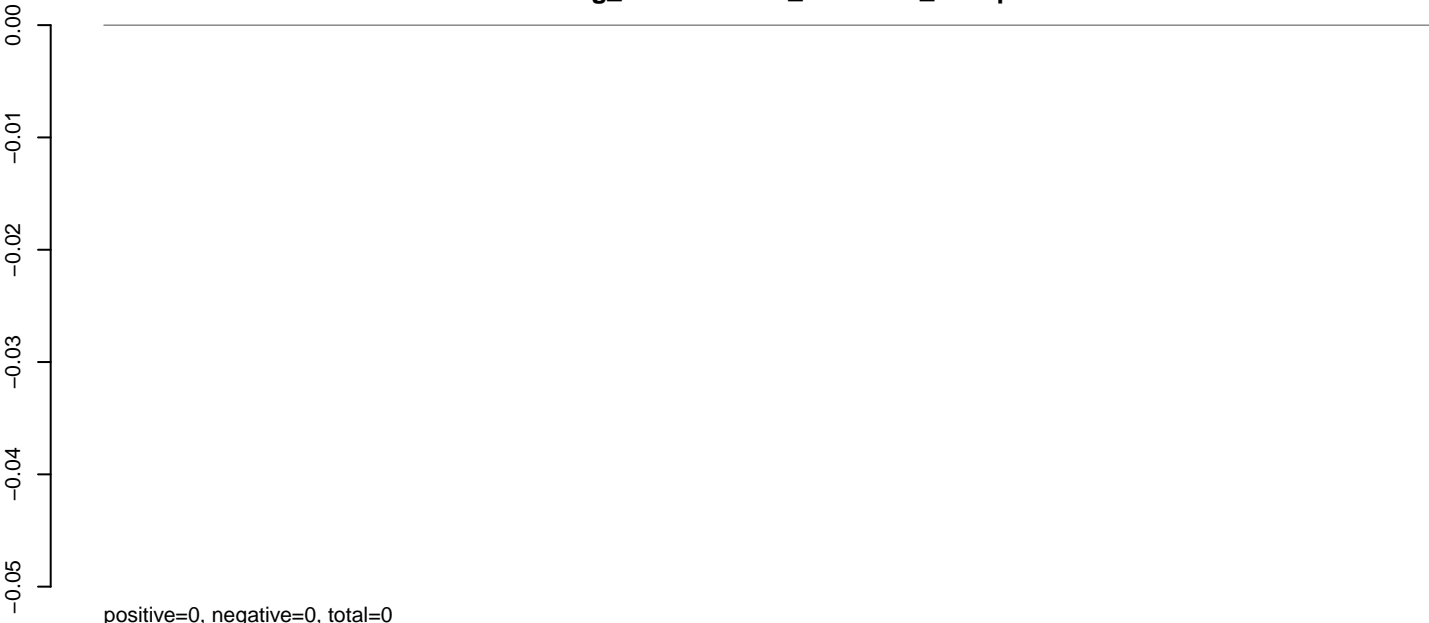


Window size=50, length=15444, virus@NC_025400.1-Malaka1_virus_isolate_SudAr_1169-64:1-15444

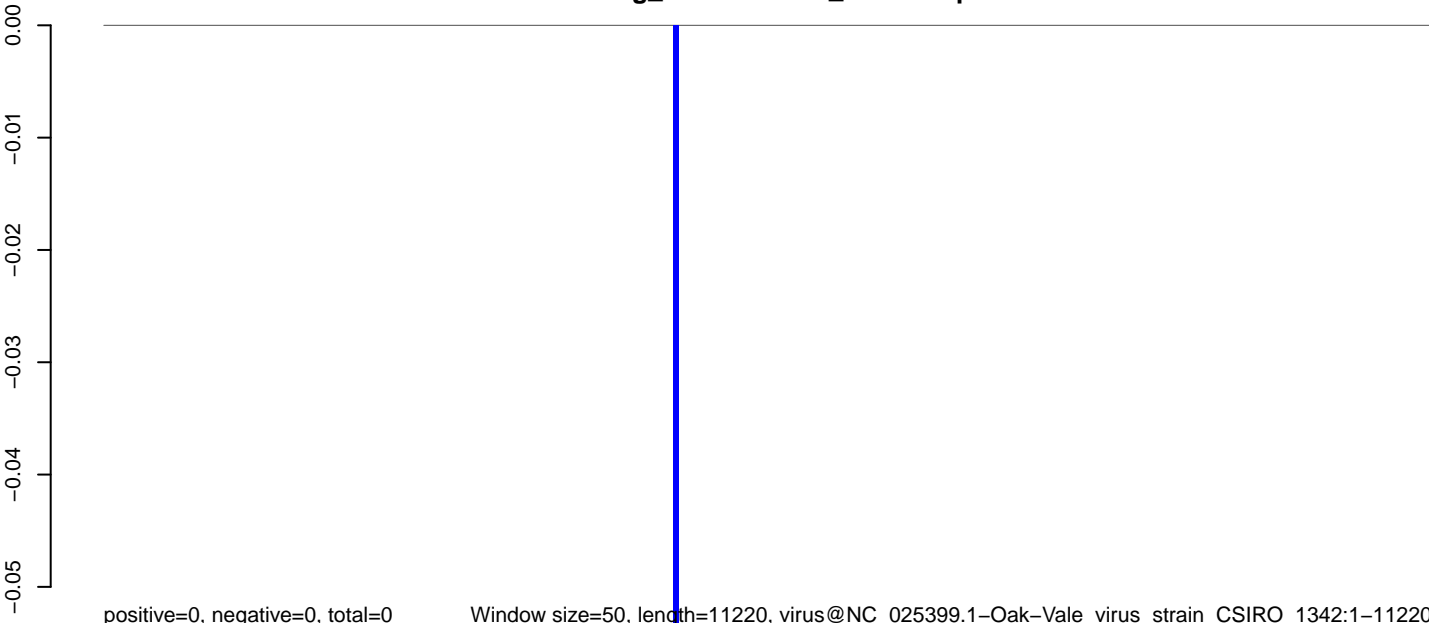
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

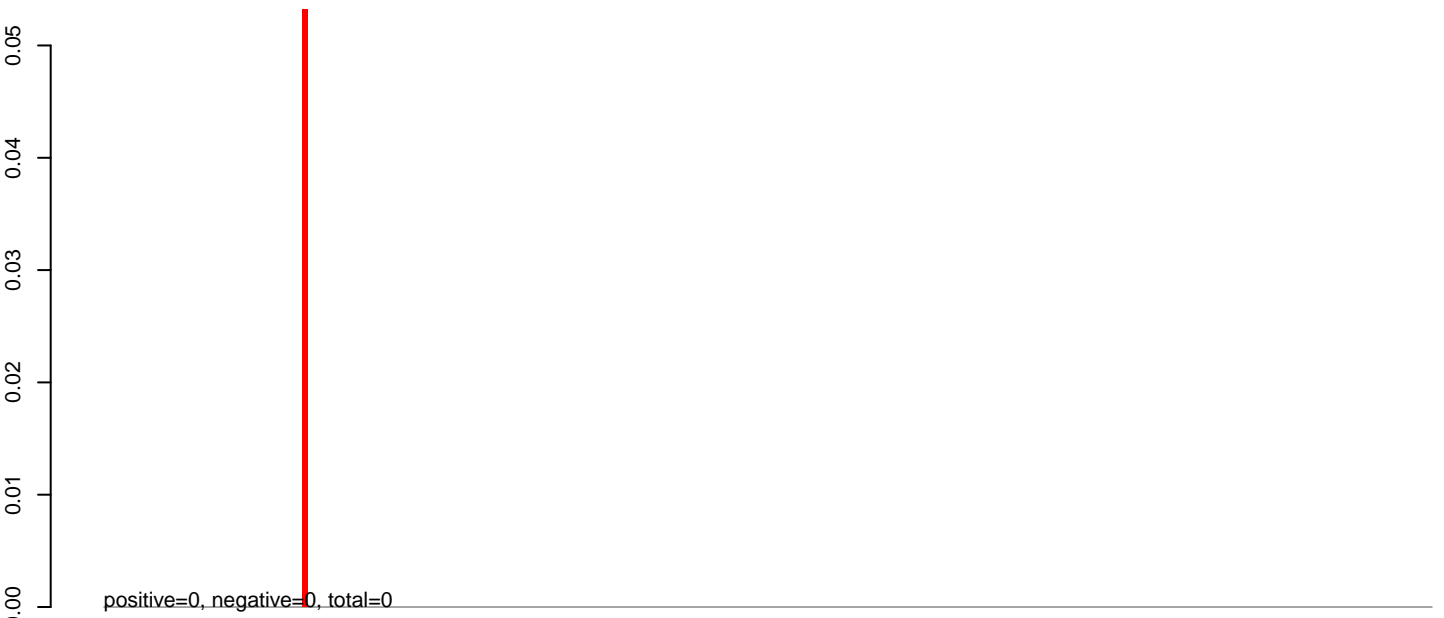


AeAeg_CCL.125cells_CHIKV.rep

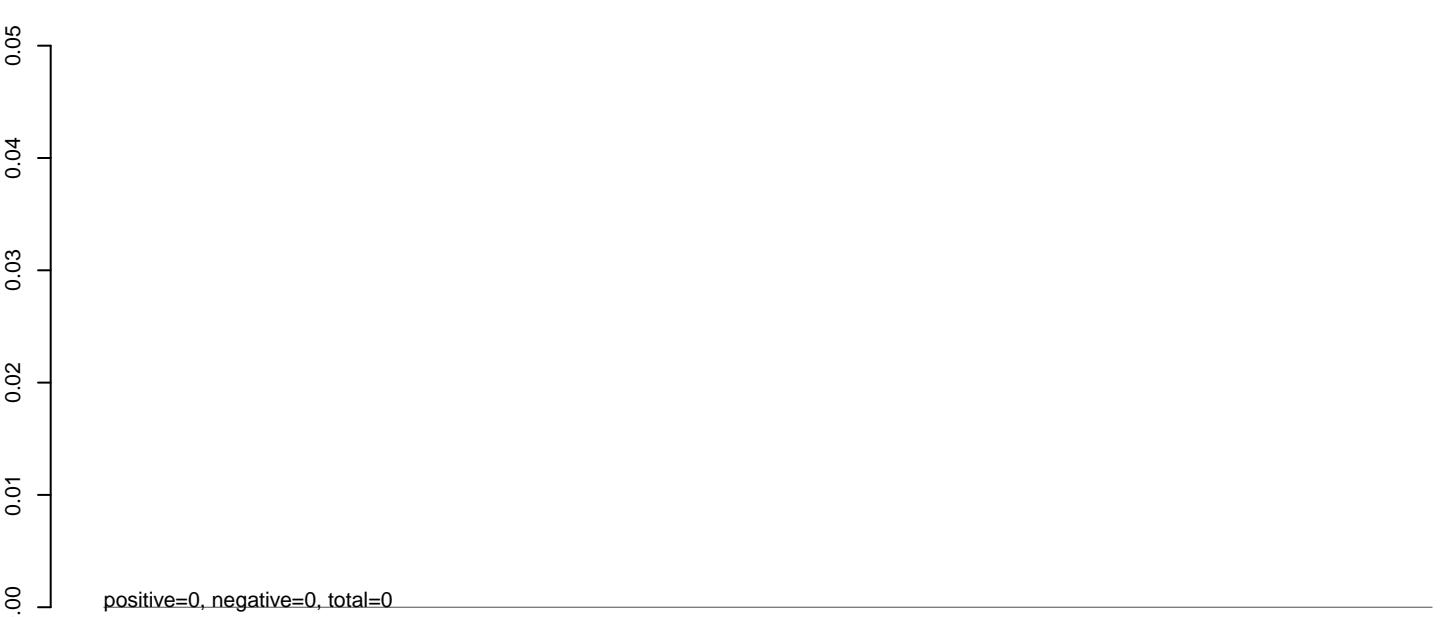


0 2000 4000 6000 8000 10000

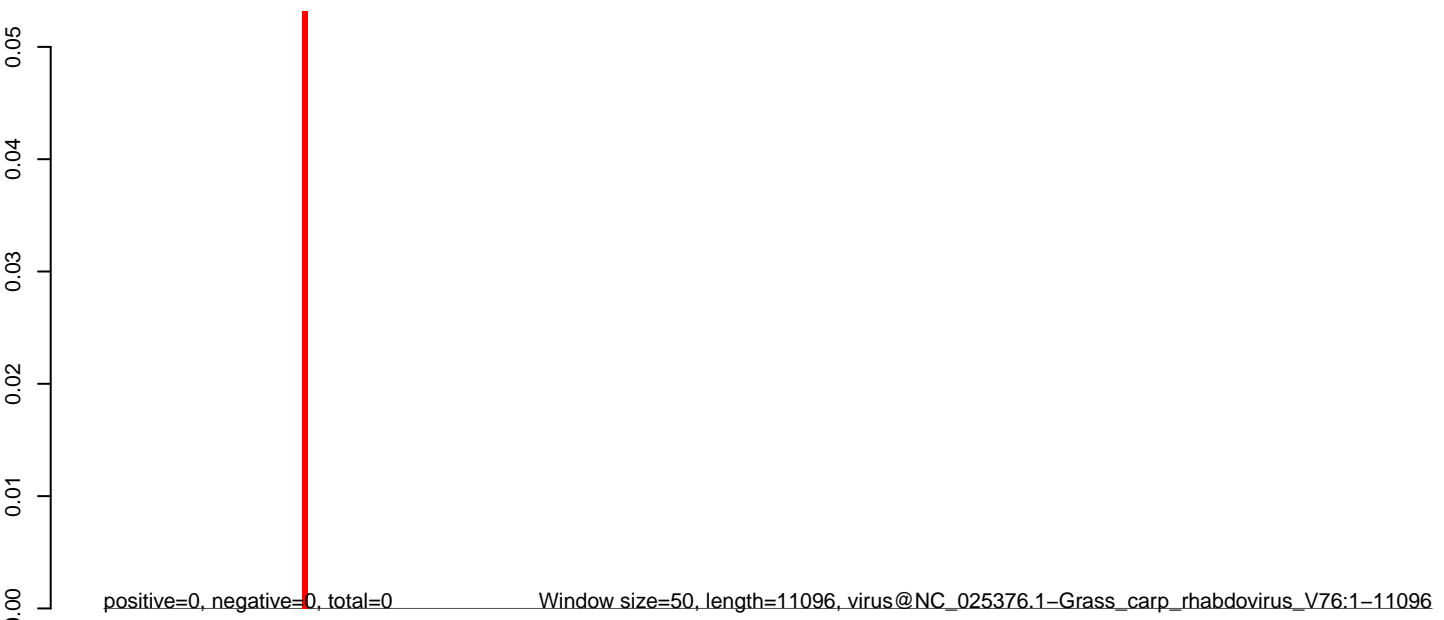
AeAeg_CCL.125cells_CHIKV.18_23.rep



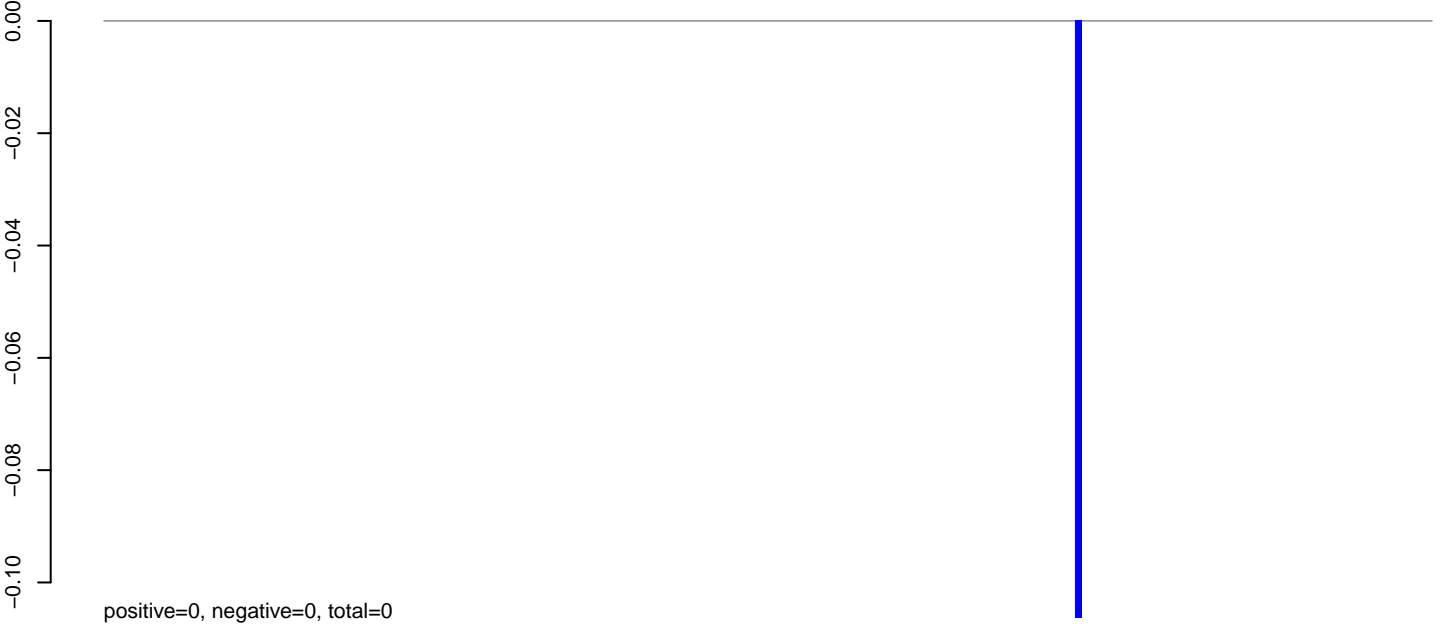
AeAeg_CCL.125cells_CHIKV.24_35.rep



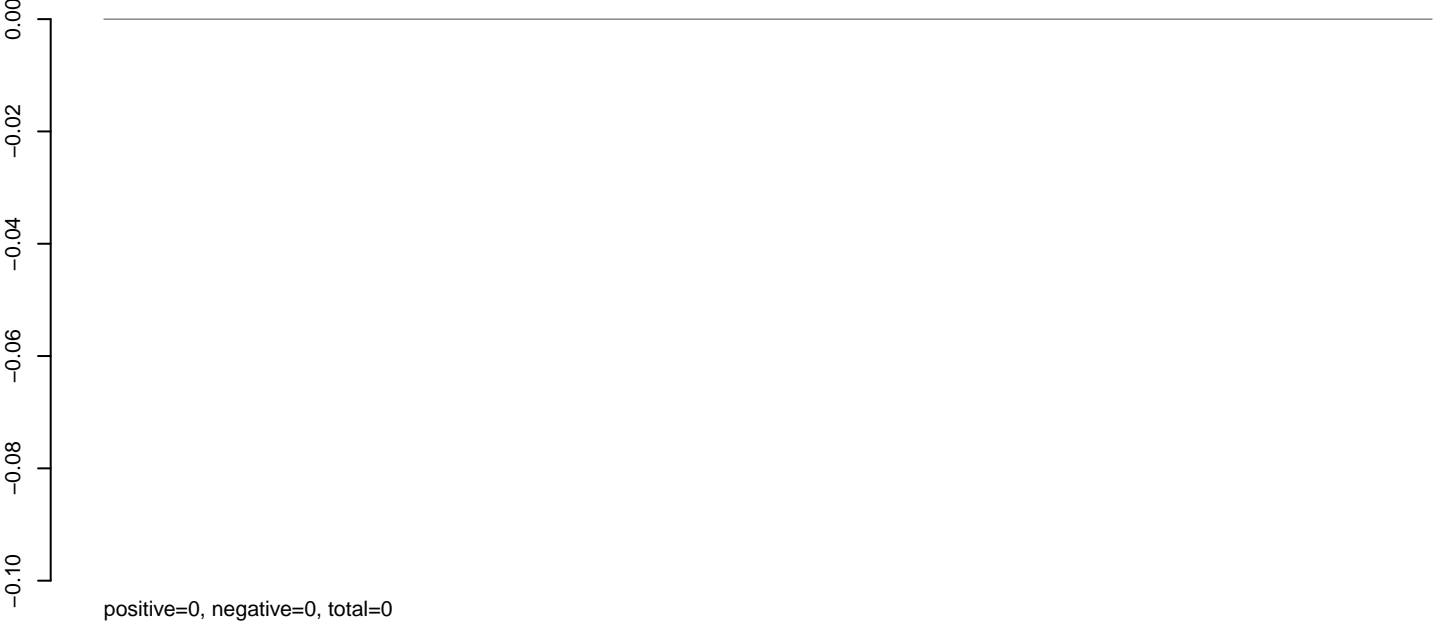
AeAeg_CCL.125cells_CHIKV.rep



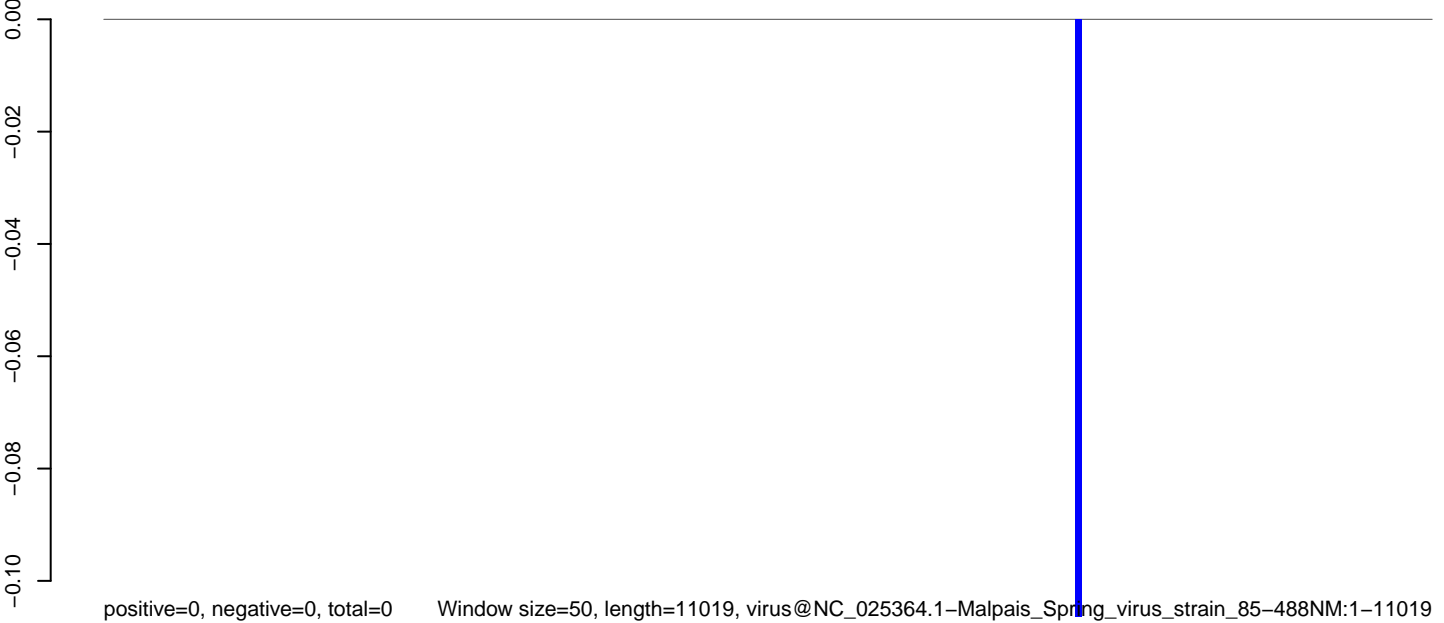
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

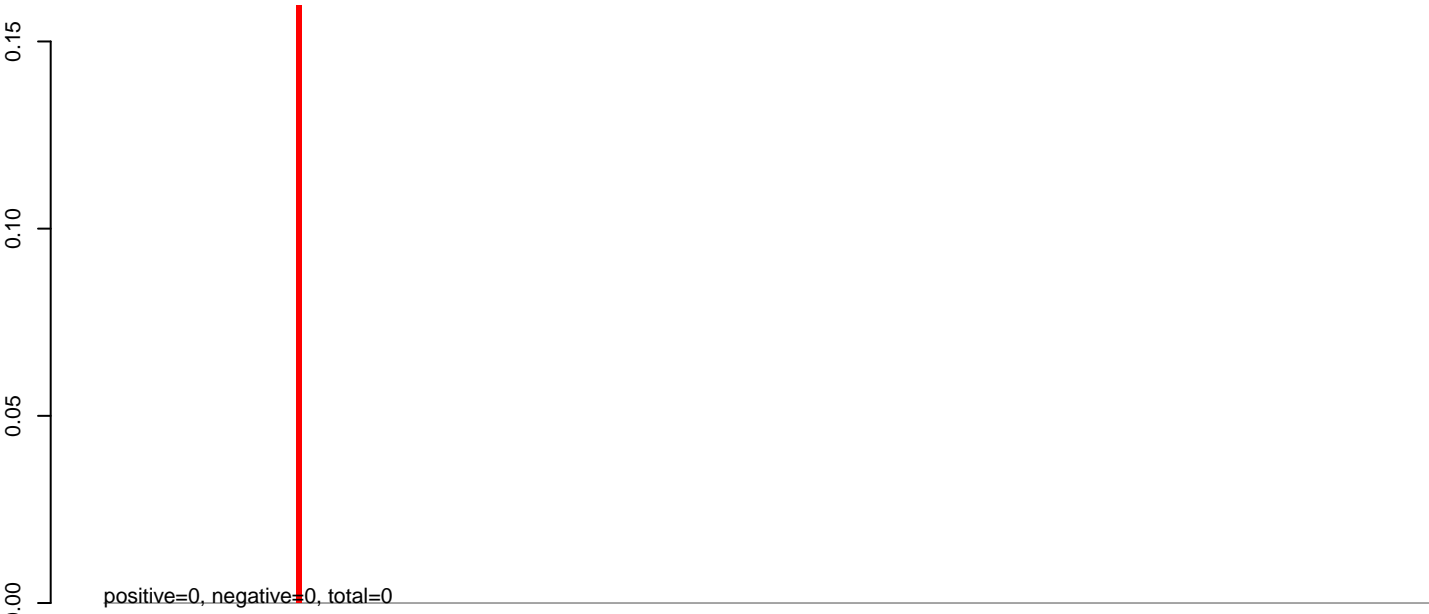


AeAeg_CCL.125cells_CHIKV.rep

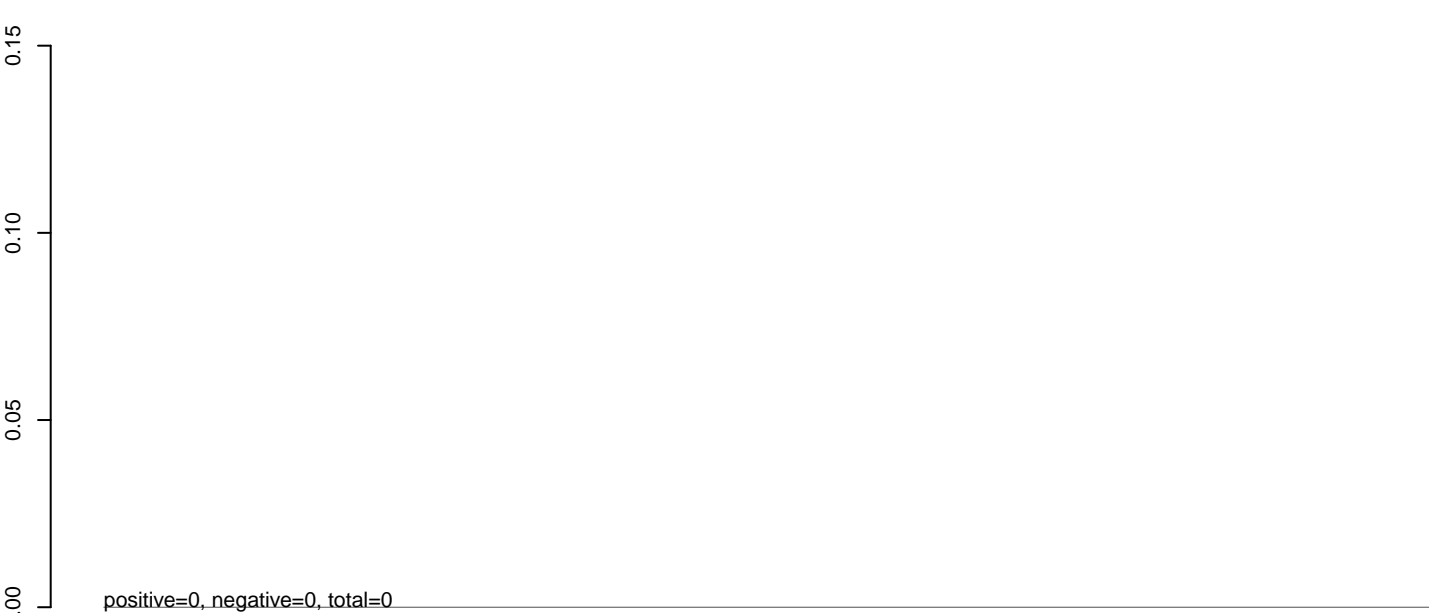


0 2000 4000 6000 8000 10000

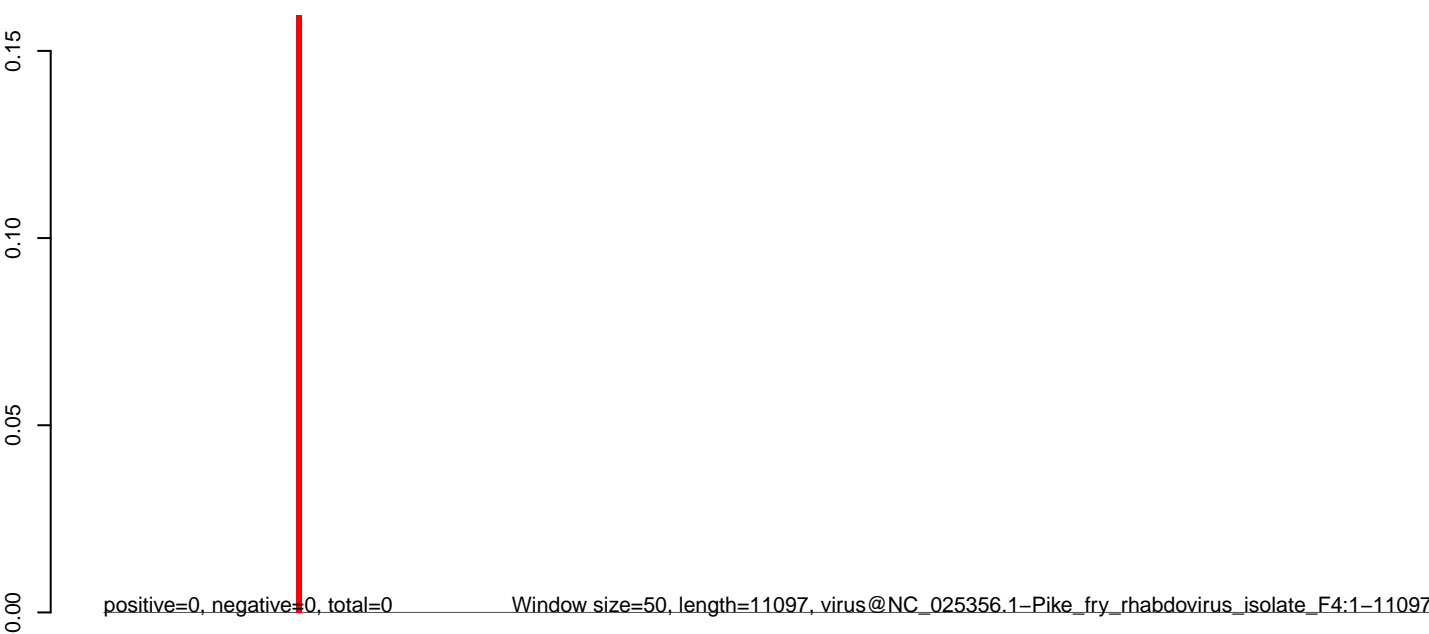
AeAeg_CCL.125cells_CHIKV.18_23.rep



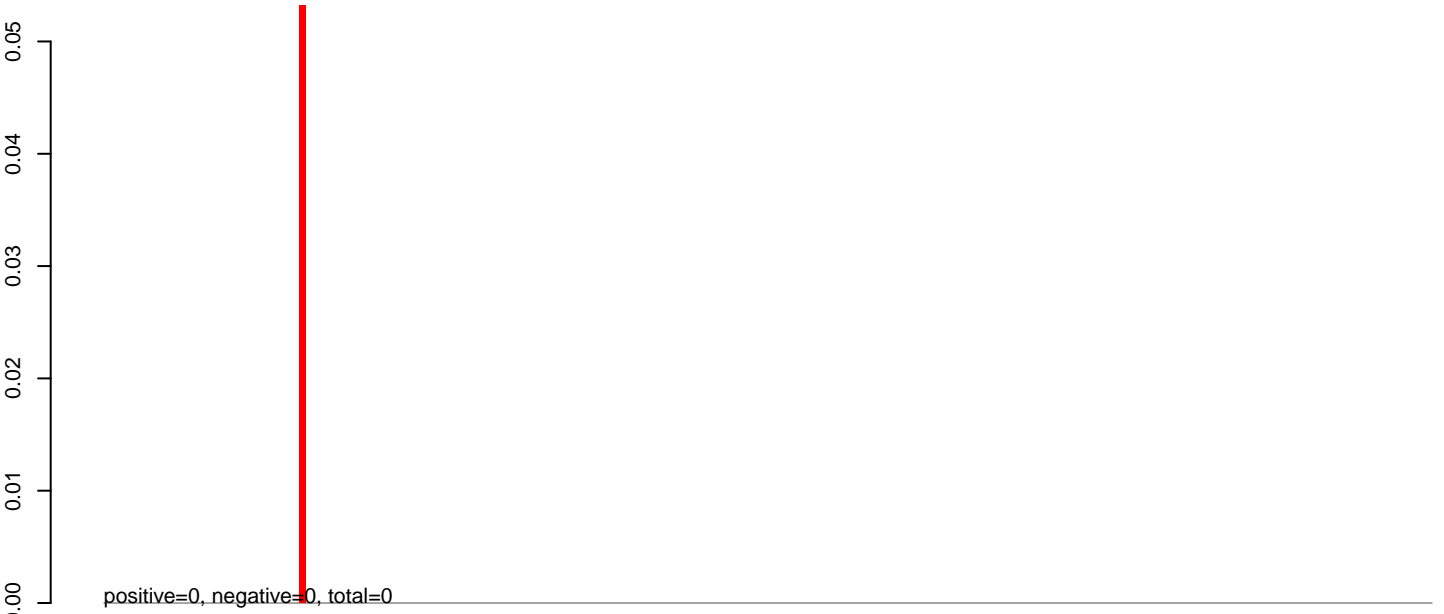
AeAeg_CCL.125cells_CHIKV.24_35.rep



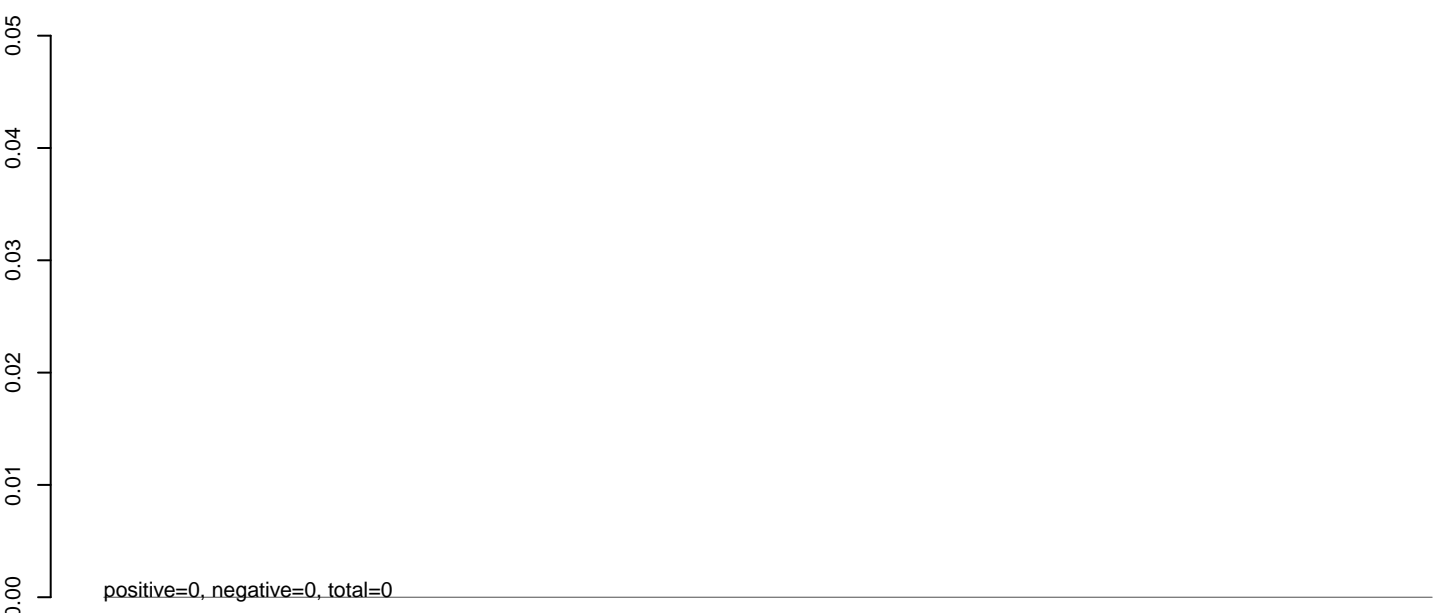
AeAeg_CCL.125cells_CHIKV.rep



AeAeg_CCL.125cells_CHIKV.18_23.rep



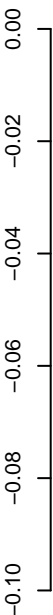
AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep

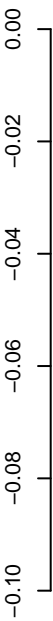


AeAeg_CCL.125cells_CHIKV.18_23.rep



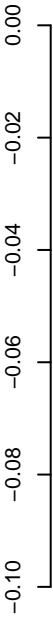
positive=0, negative=0, total=0

AeAeg_CCL.125cells_CHIKV.24_35.rep



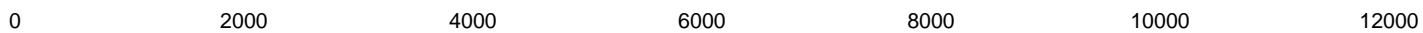
positive=0, negative=0, total=0

AeAeg_CCL.125cells_CHIKV.rep

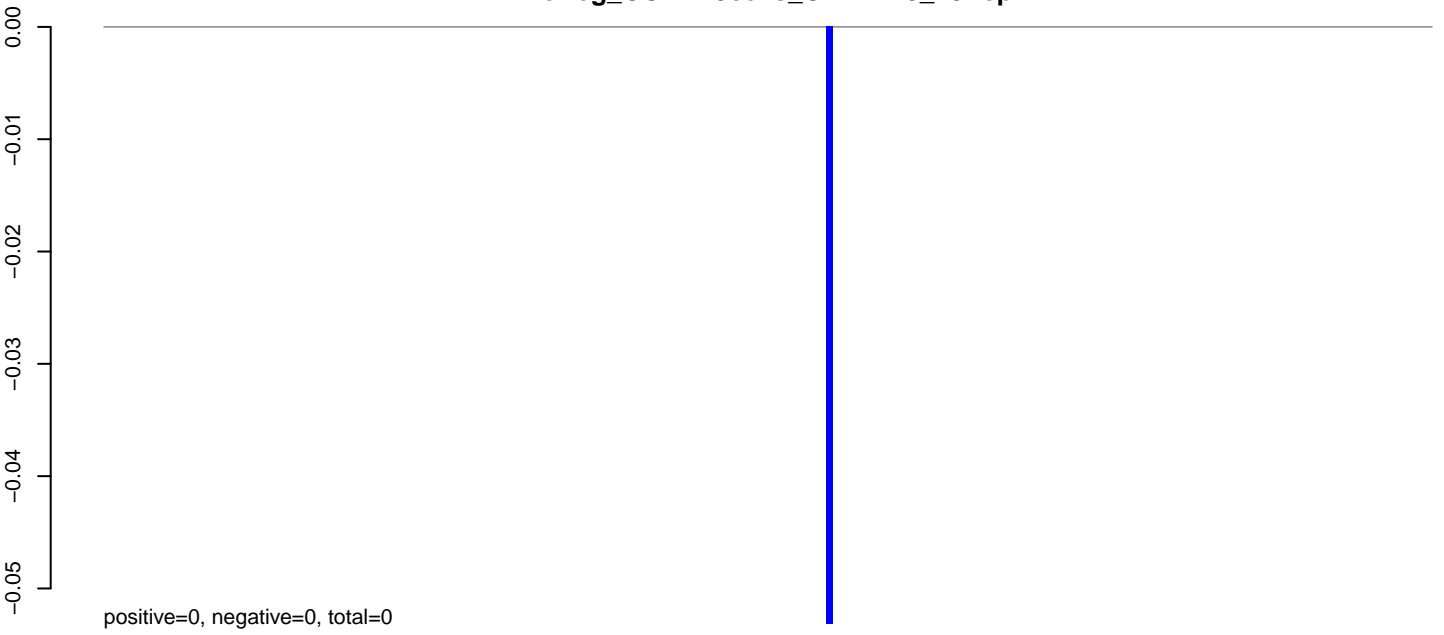


positive=0, negative=0, total=0

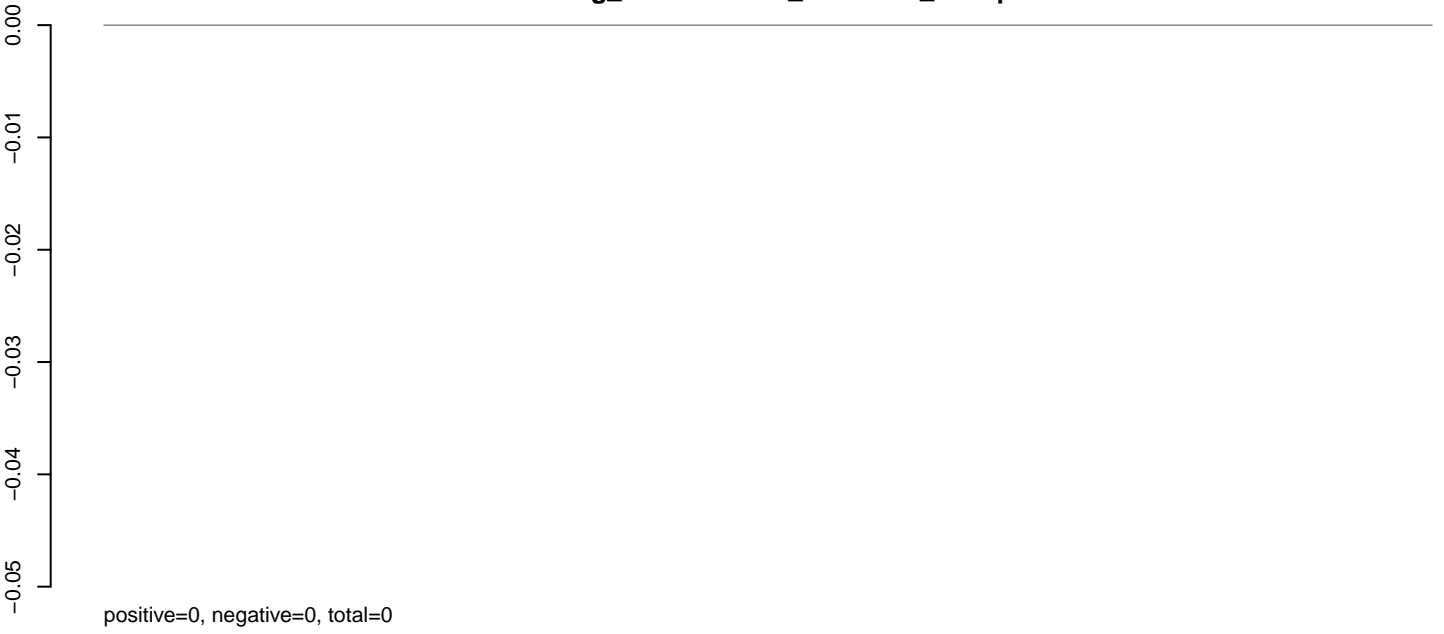
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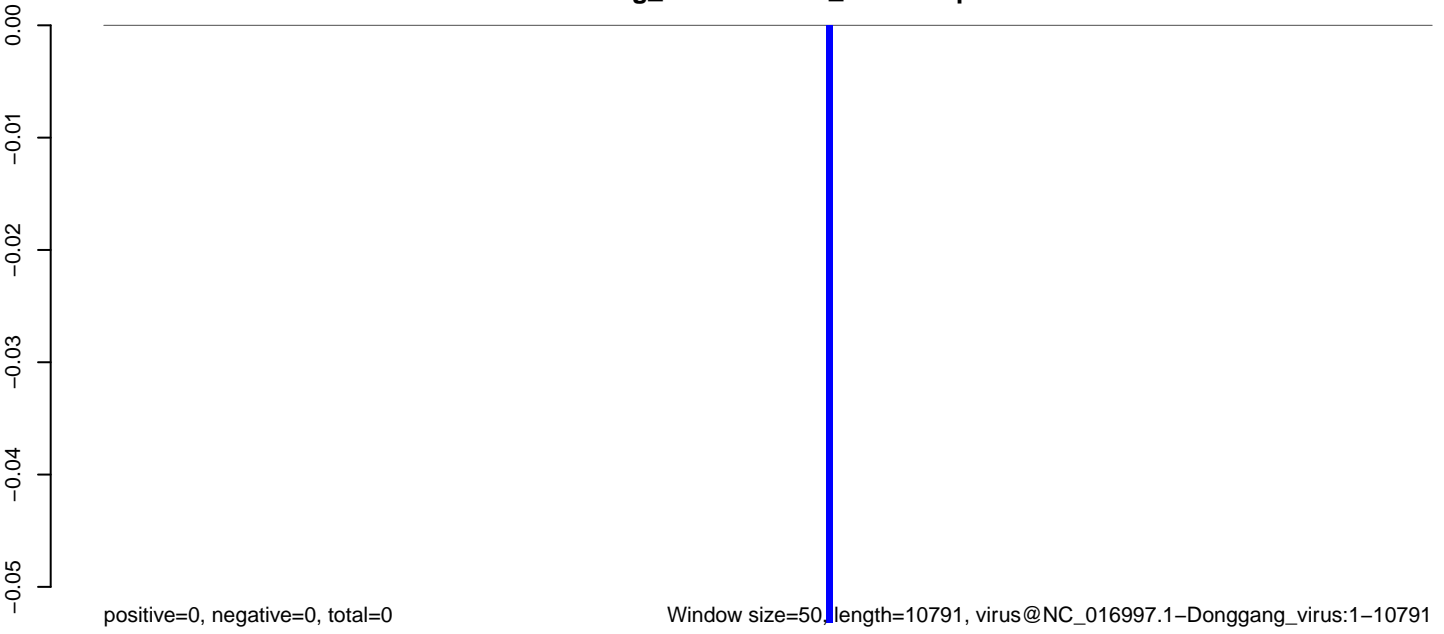
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

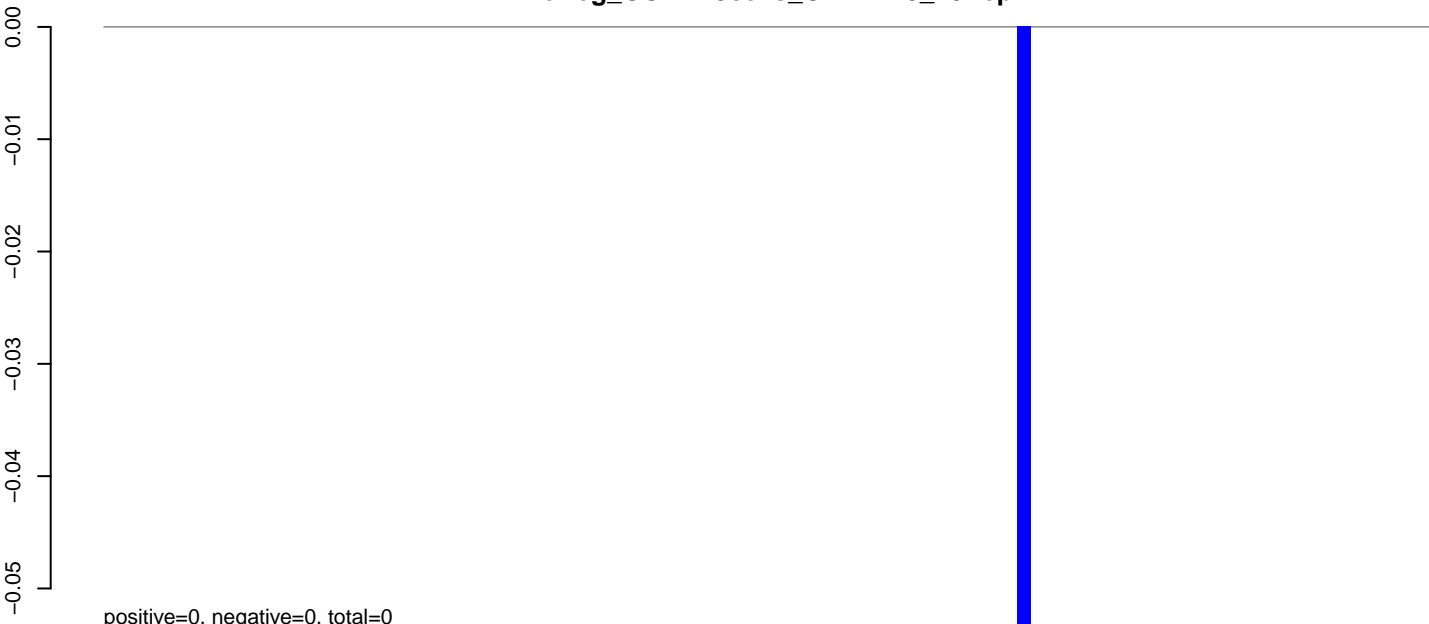


AeAeg_CCL.125cells_CHIKV.rep

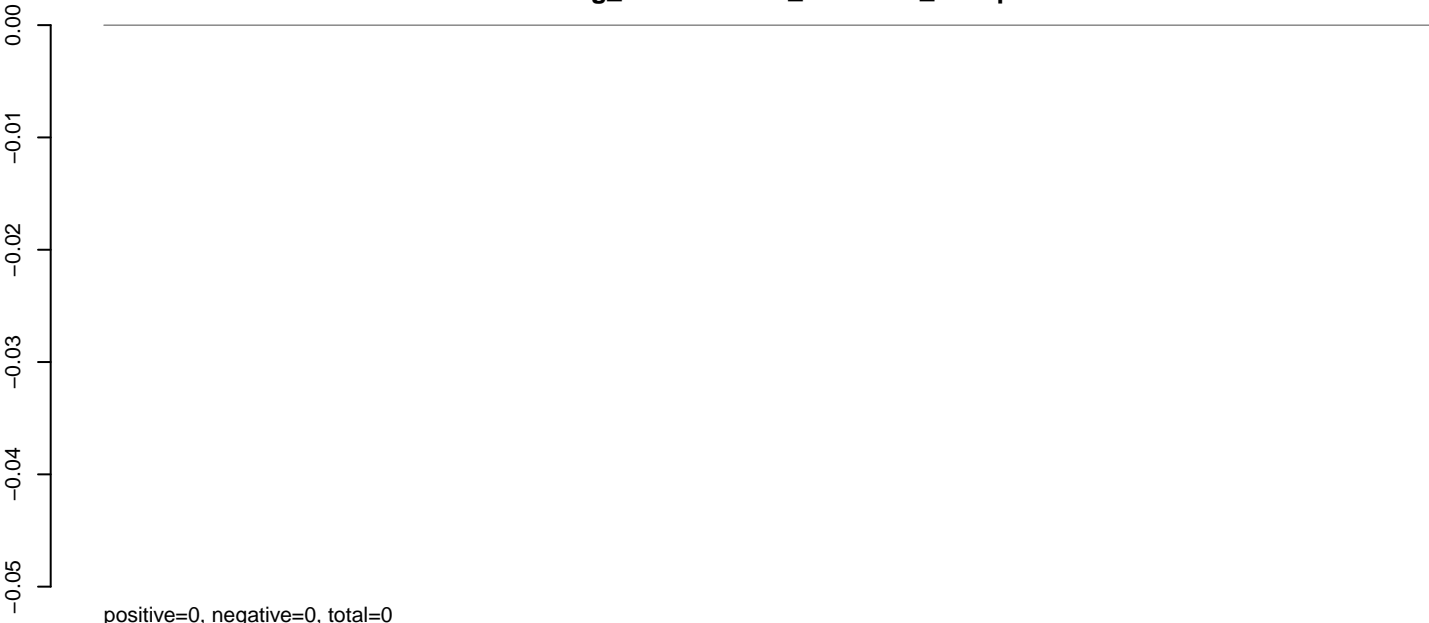


0 2000 4000 6000 8000 10000

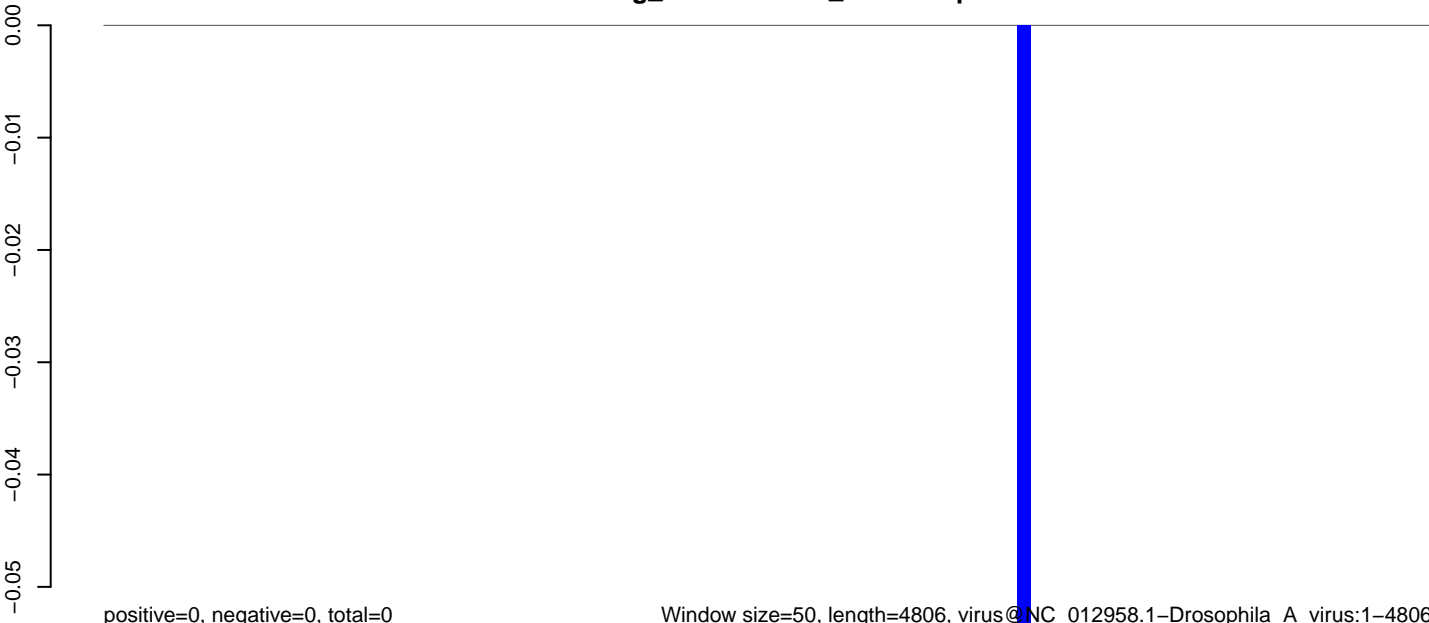
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

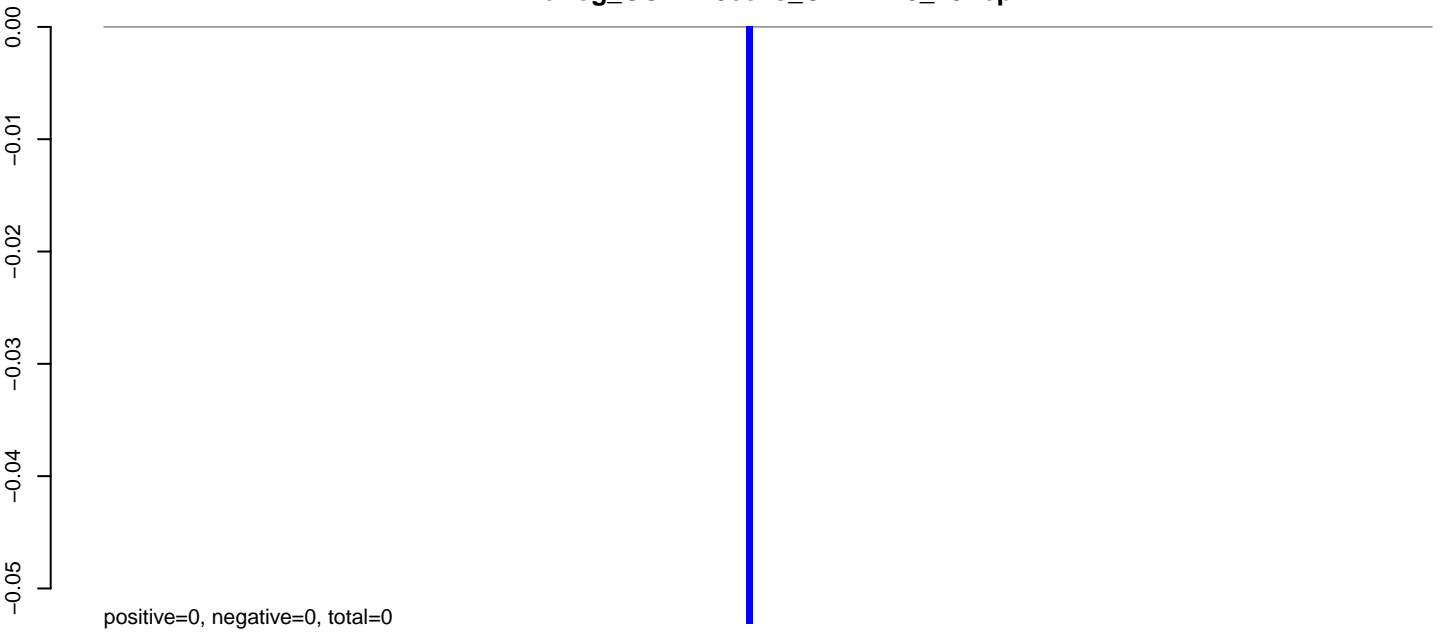


AeAeg_CCL.125cells_CHIKV.rep

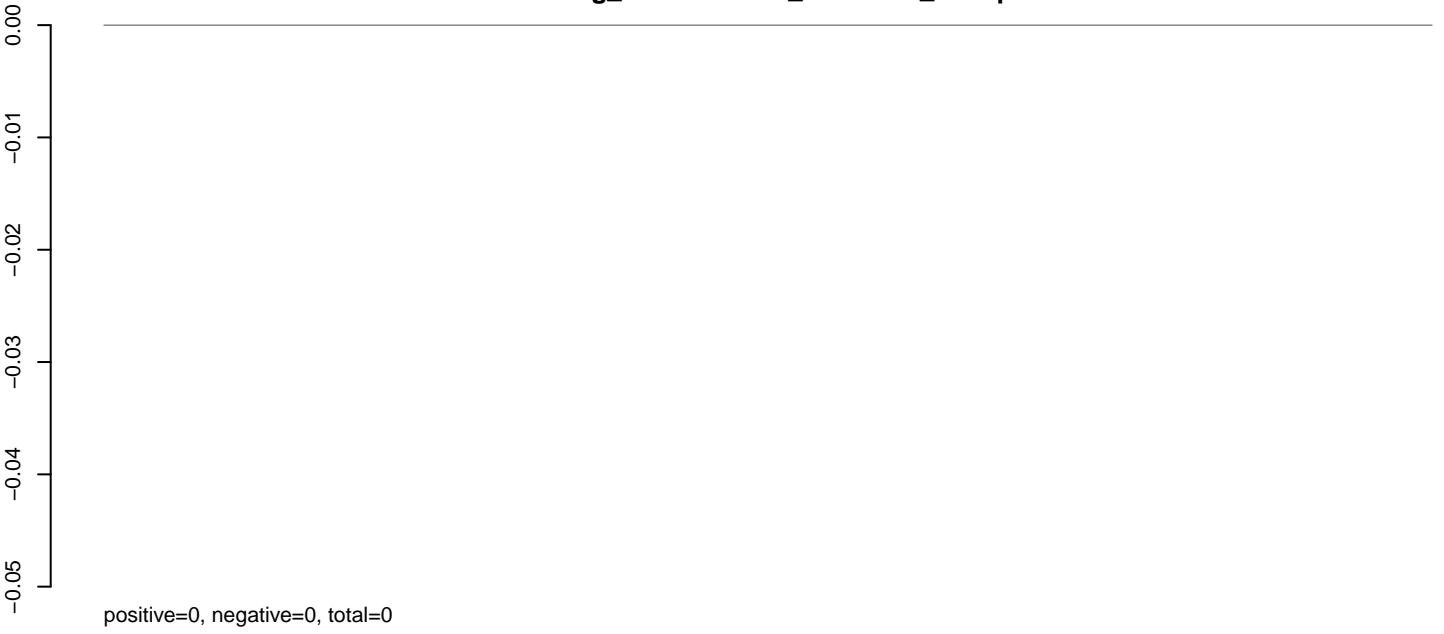


0 1000 2000 3000 4000 5000

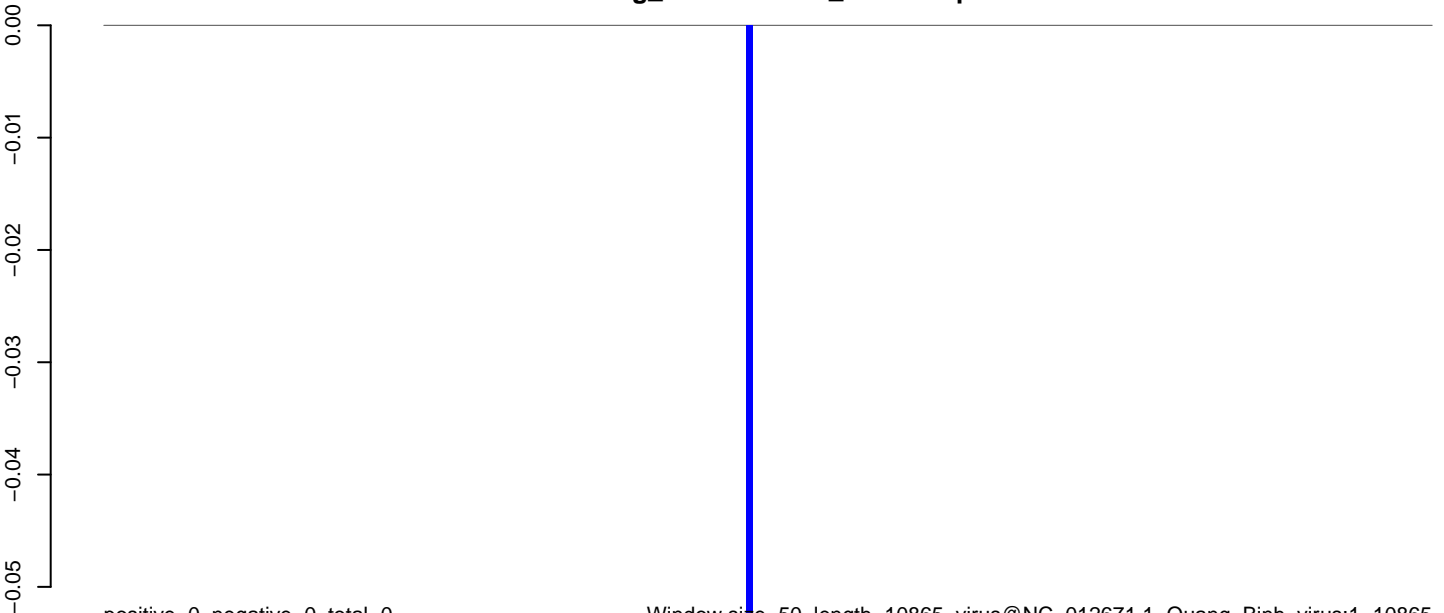
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

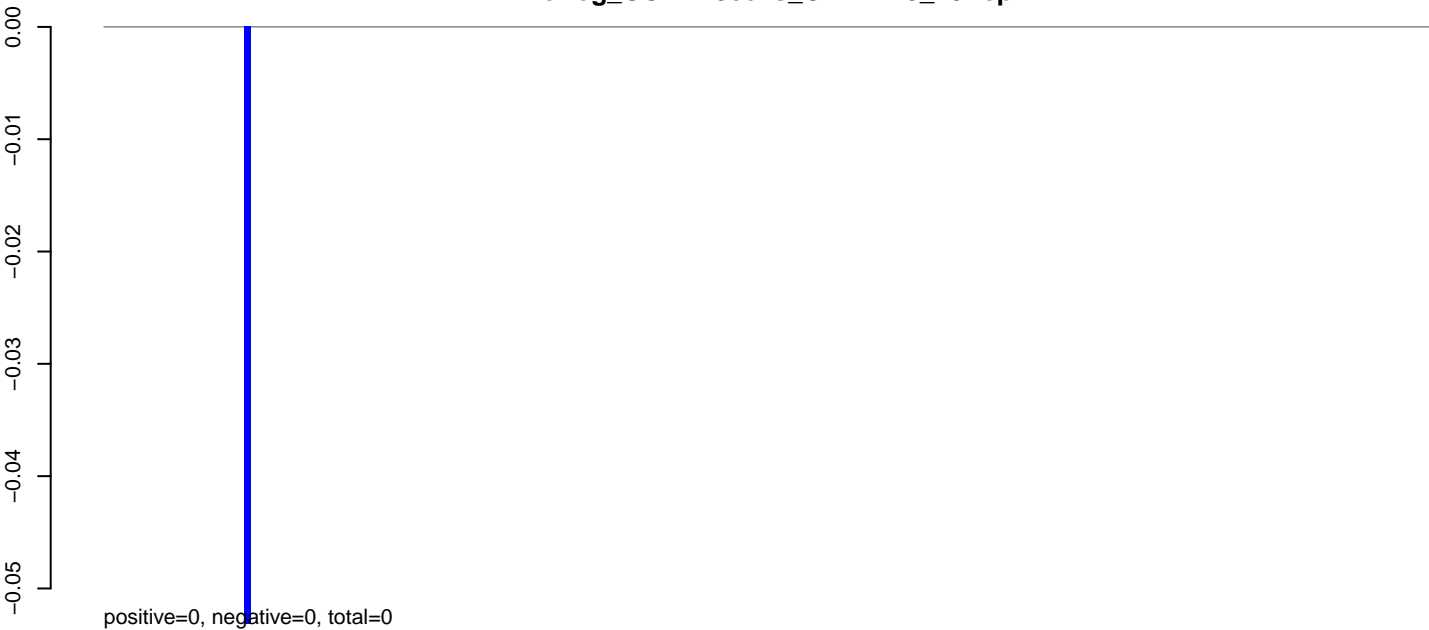


AeAeg_CCL.125cells_CHIKV.rep

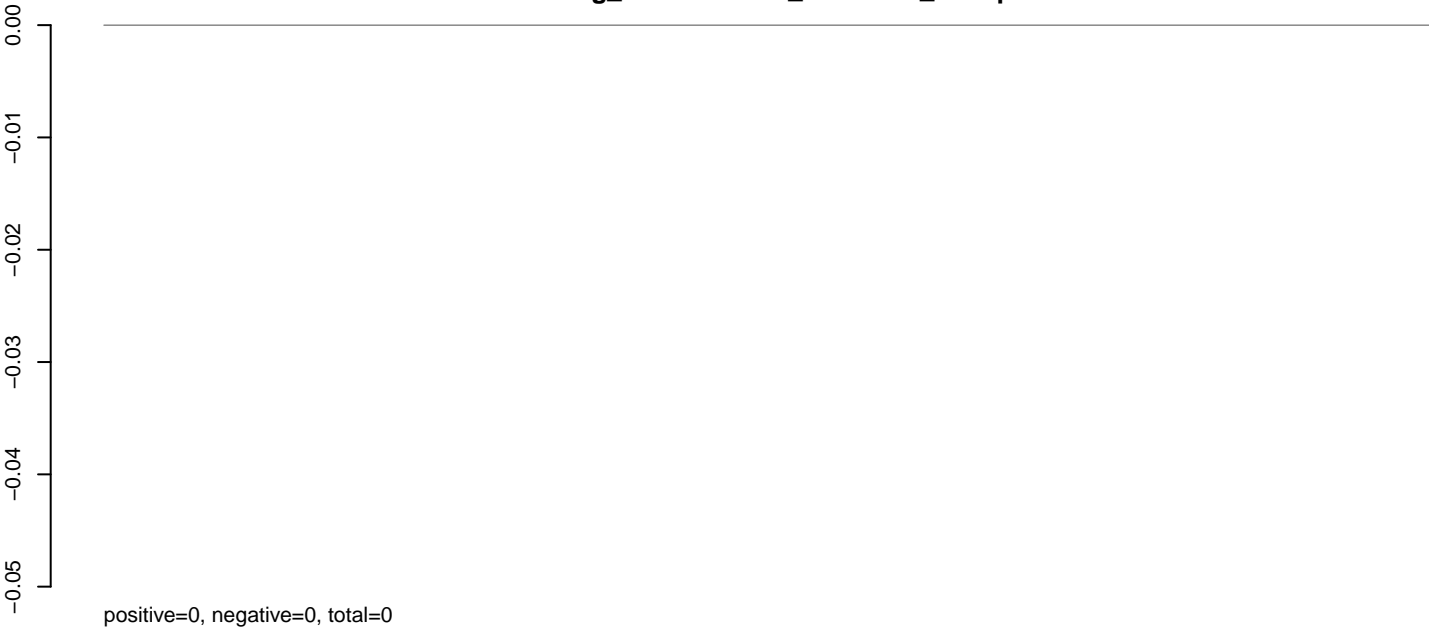


0 2000 4000 6000 8000 10000

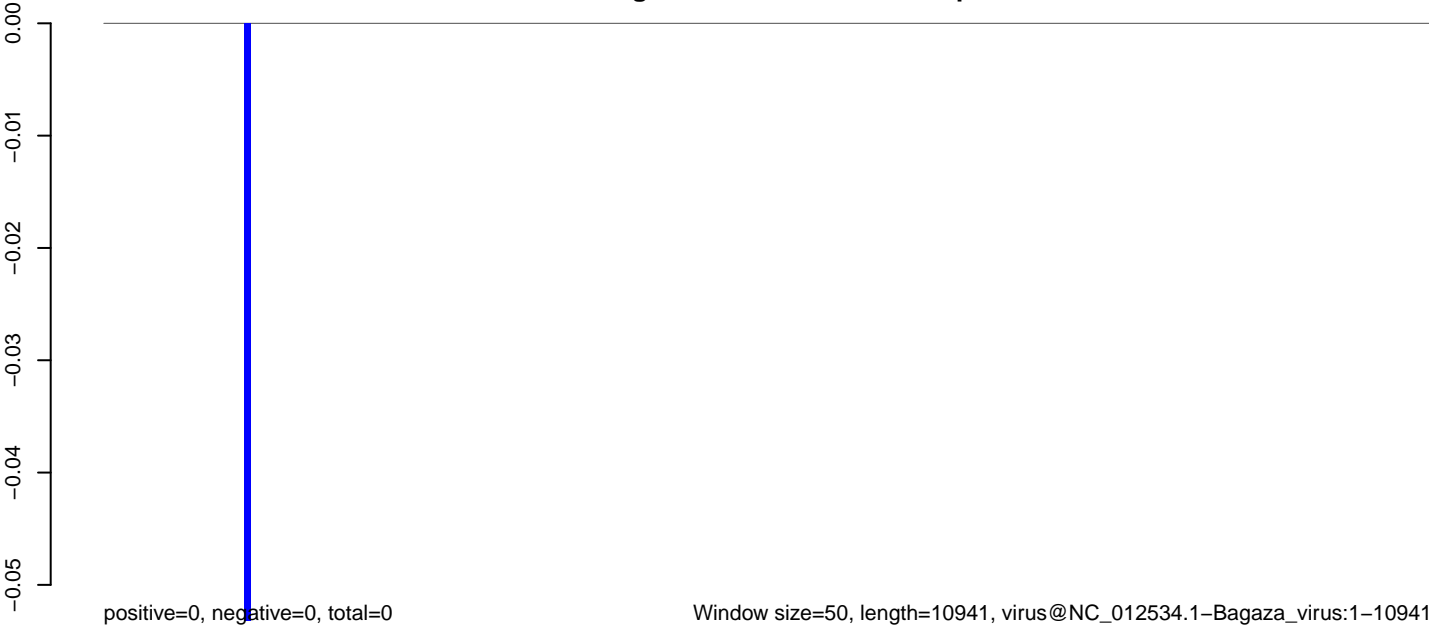
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

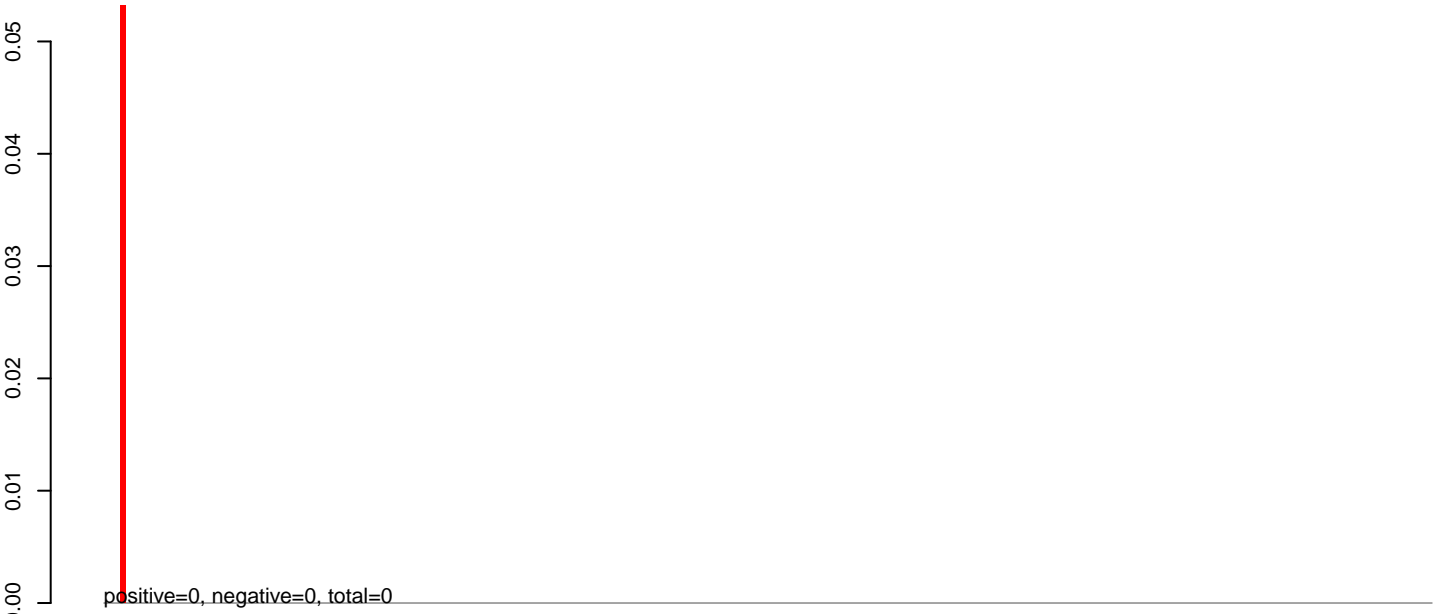


AeAeg_CCL.125cells_CHIKV.rep

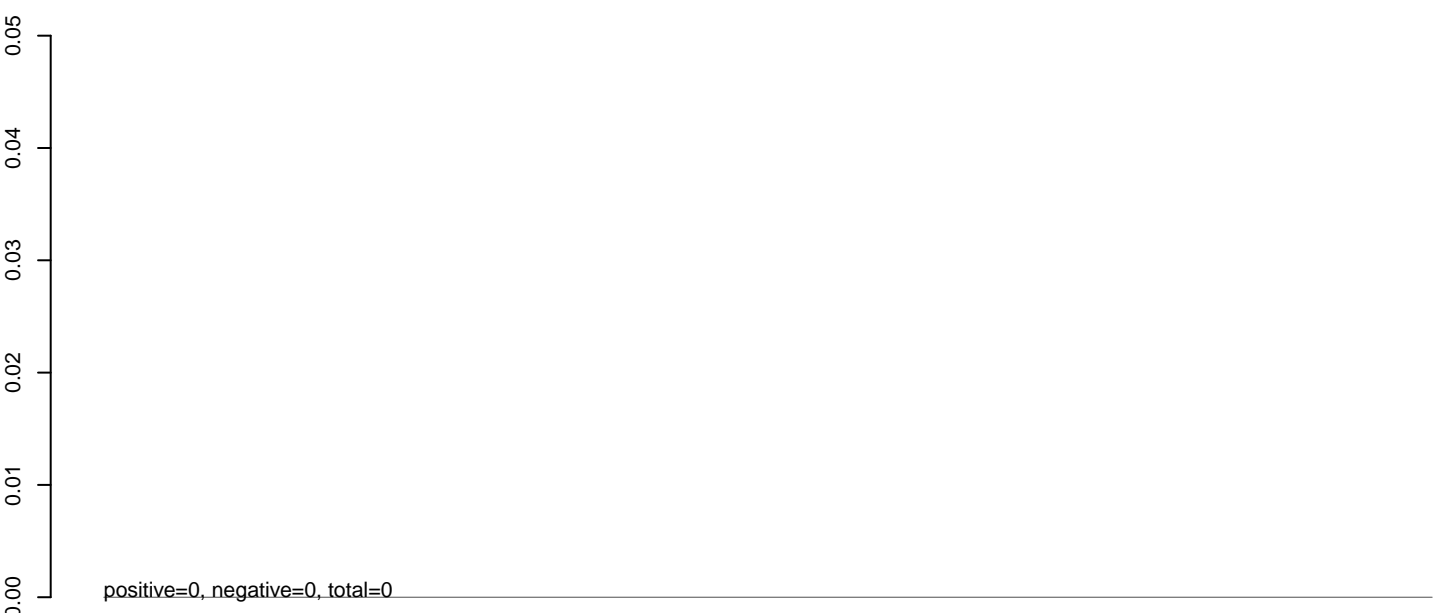


0 2000 4000 6000 8000 10000

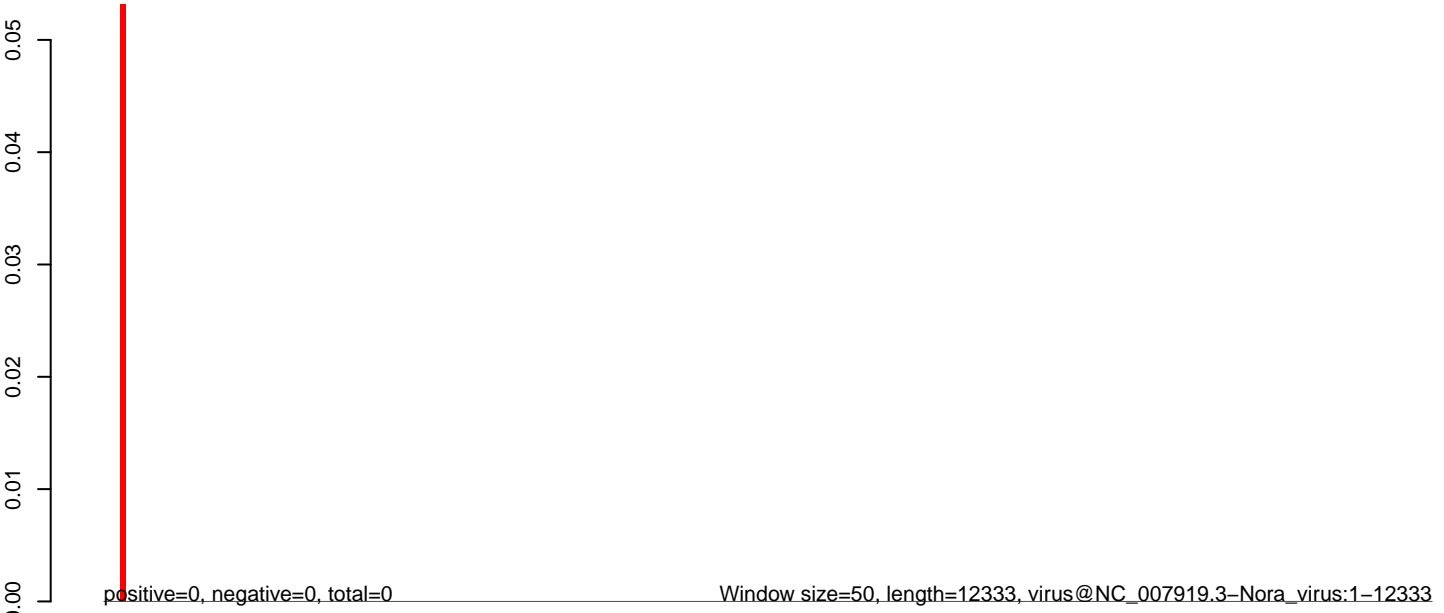
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

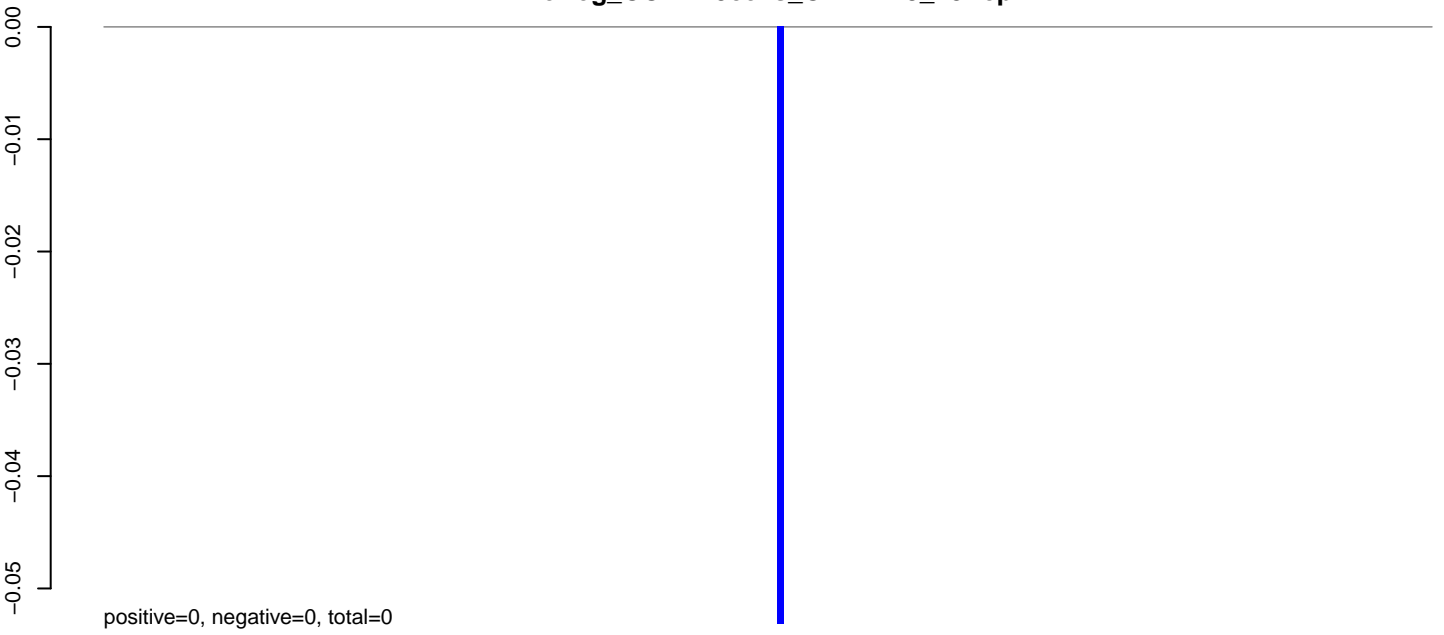


AeAeg_CCL.125cells_CHIKV.rep

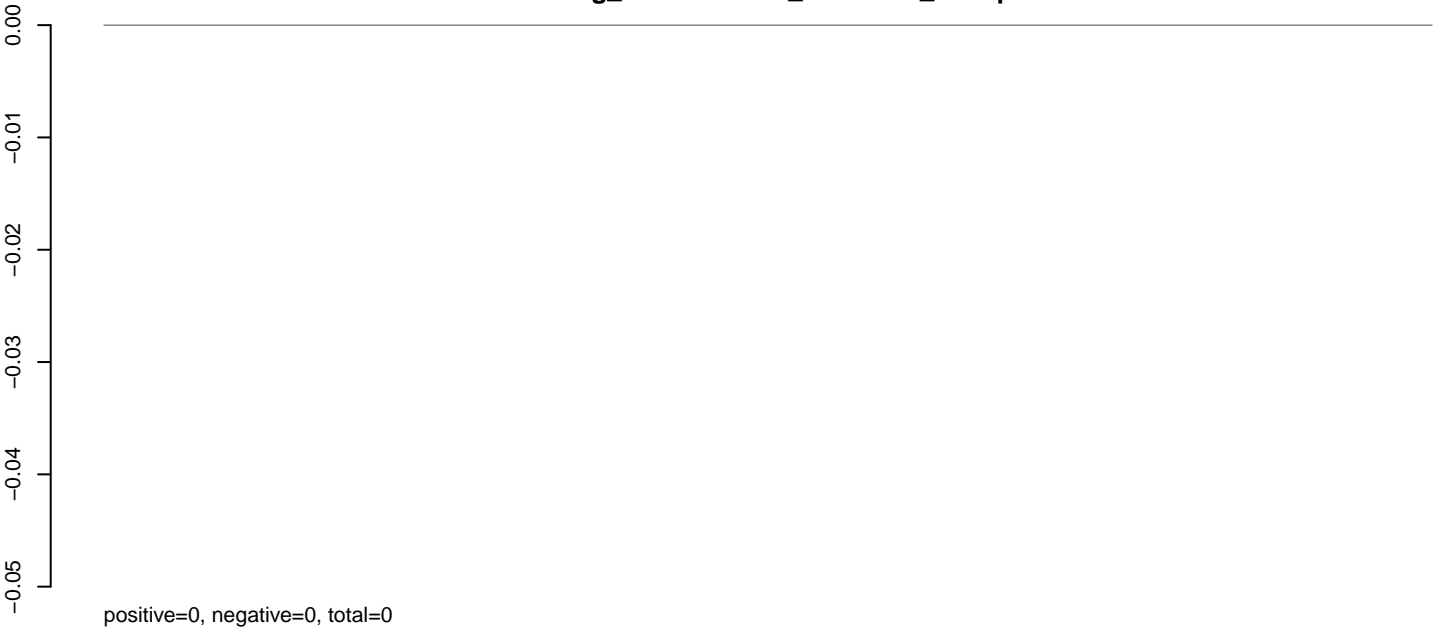


0 2000 4000 6000 8000 10000 12000

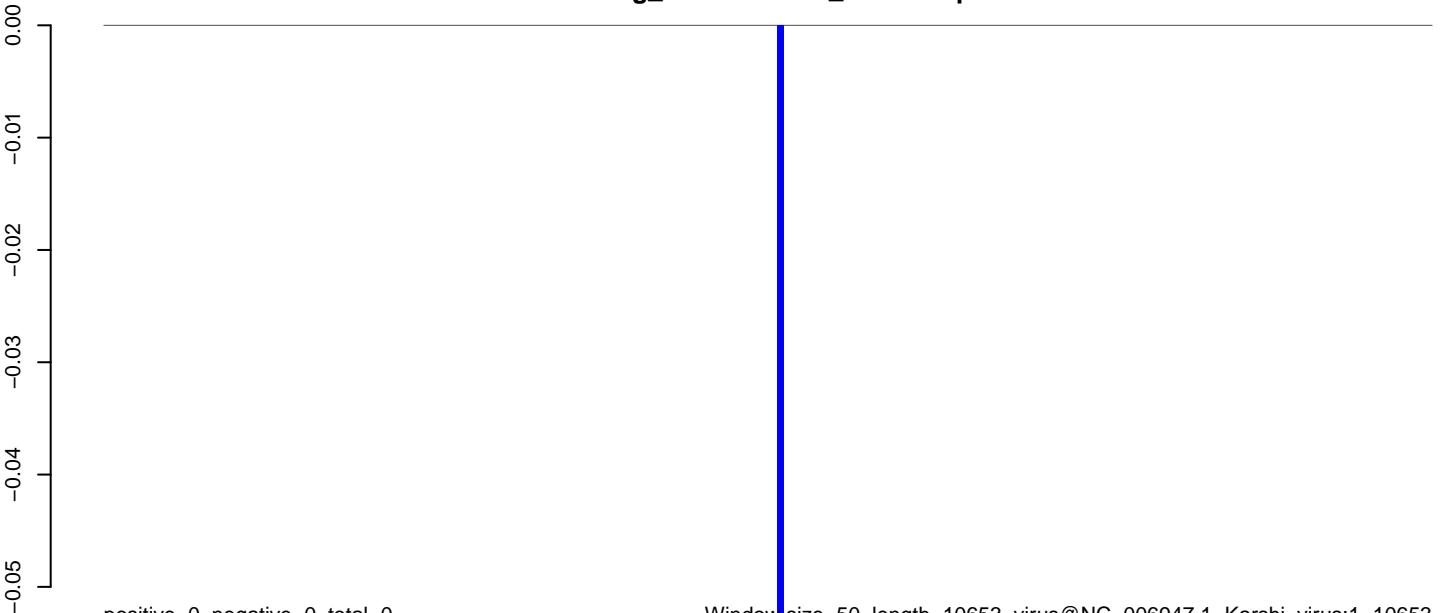
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

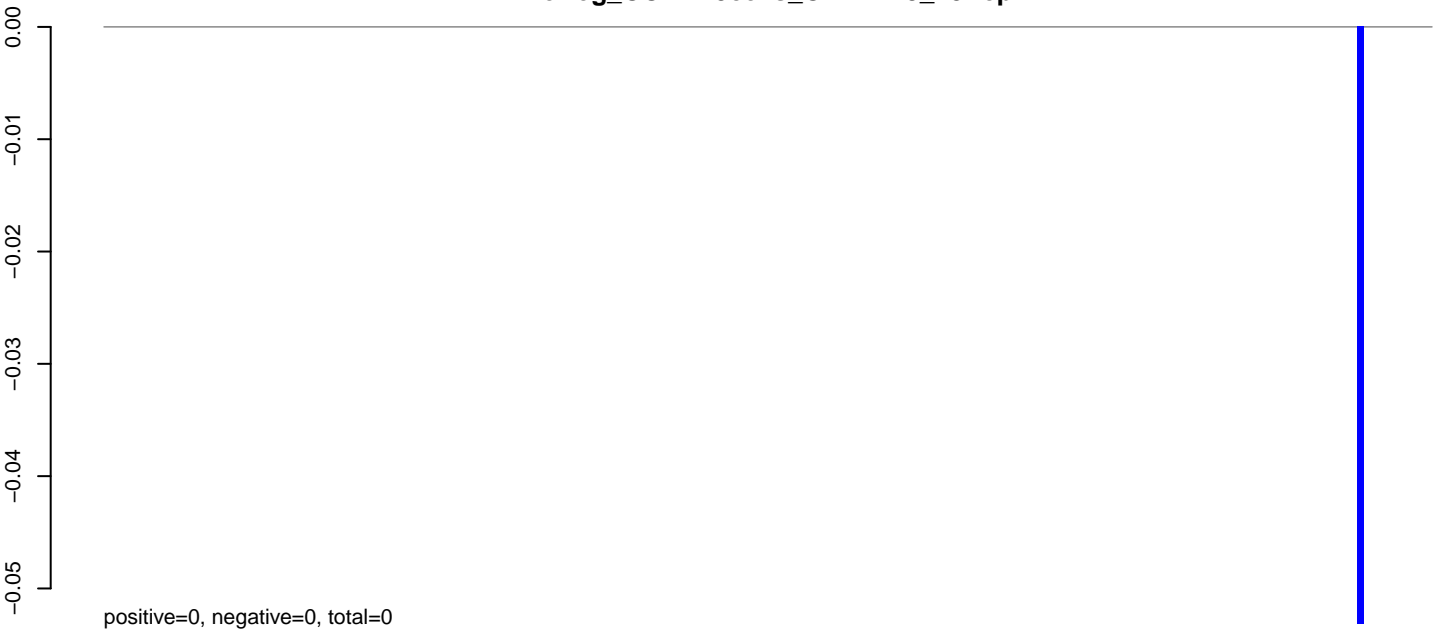


AeAeg_CCL.125cells_CHIKV.rep

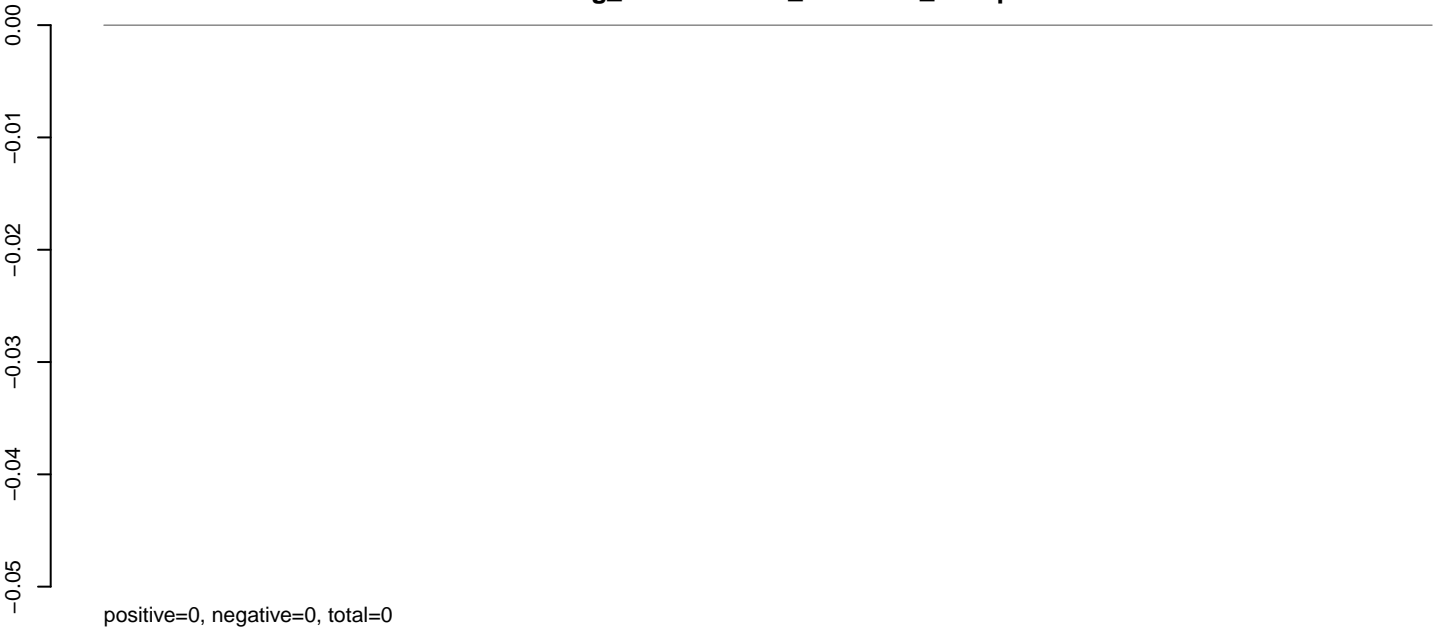


0 2000 4000 6000 8000 10000

AeAeg_CCL.125cells_CHIKV.18_23.rep



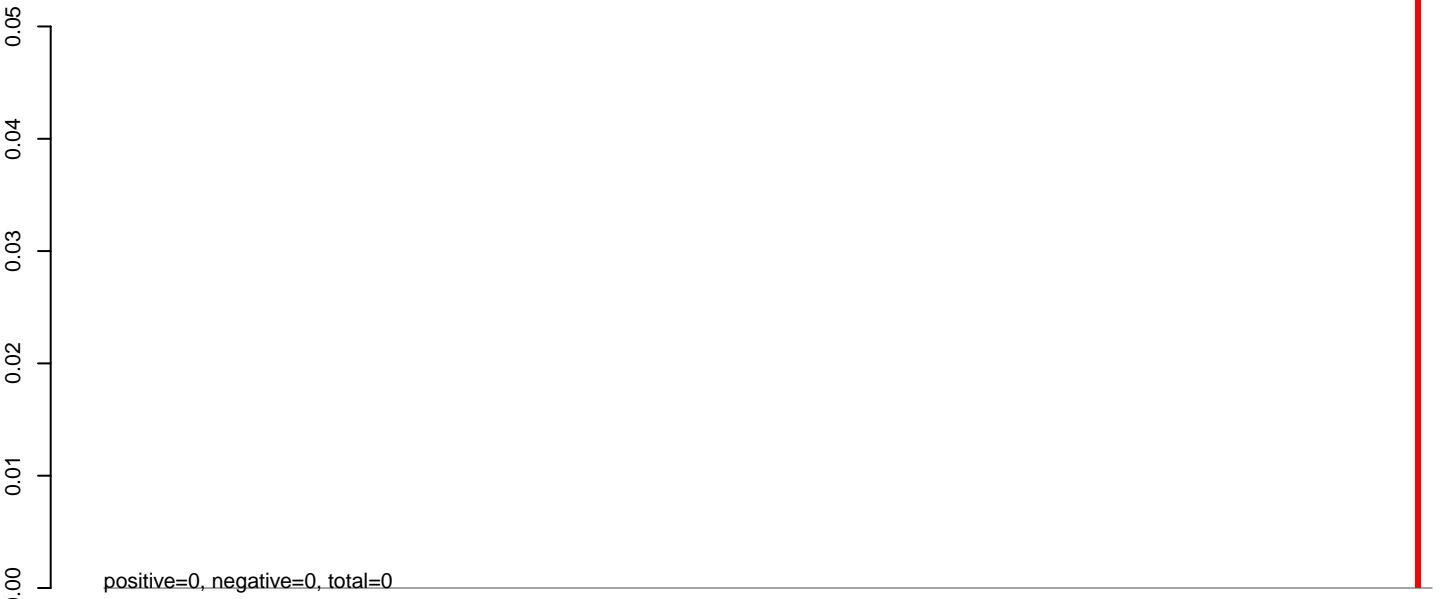
AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep

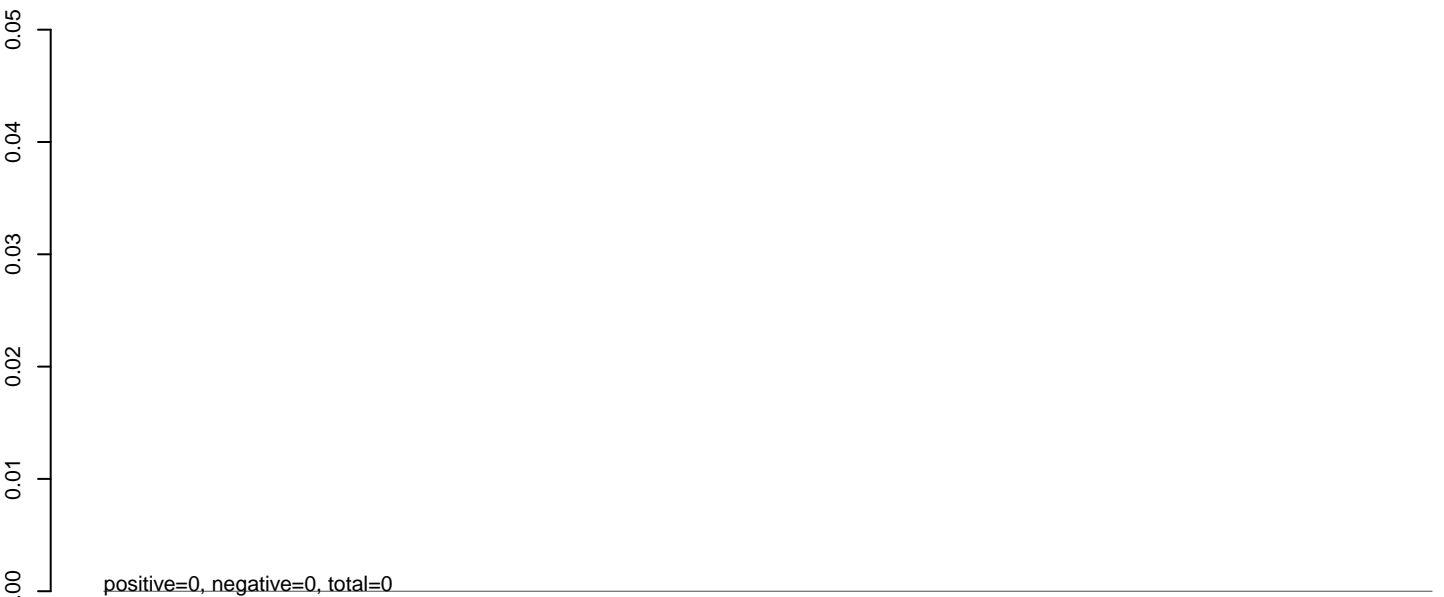


AeAeg_CCL.125cells_CHIKV.18_23.rep



positive=0, negative=0, total=0

AeAeg_CCL.125cells_CHIKV.24_35.rep



positive=0, negative=0, total=0

AeAeg_CCL.125cells_CHIKV.rep

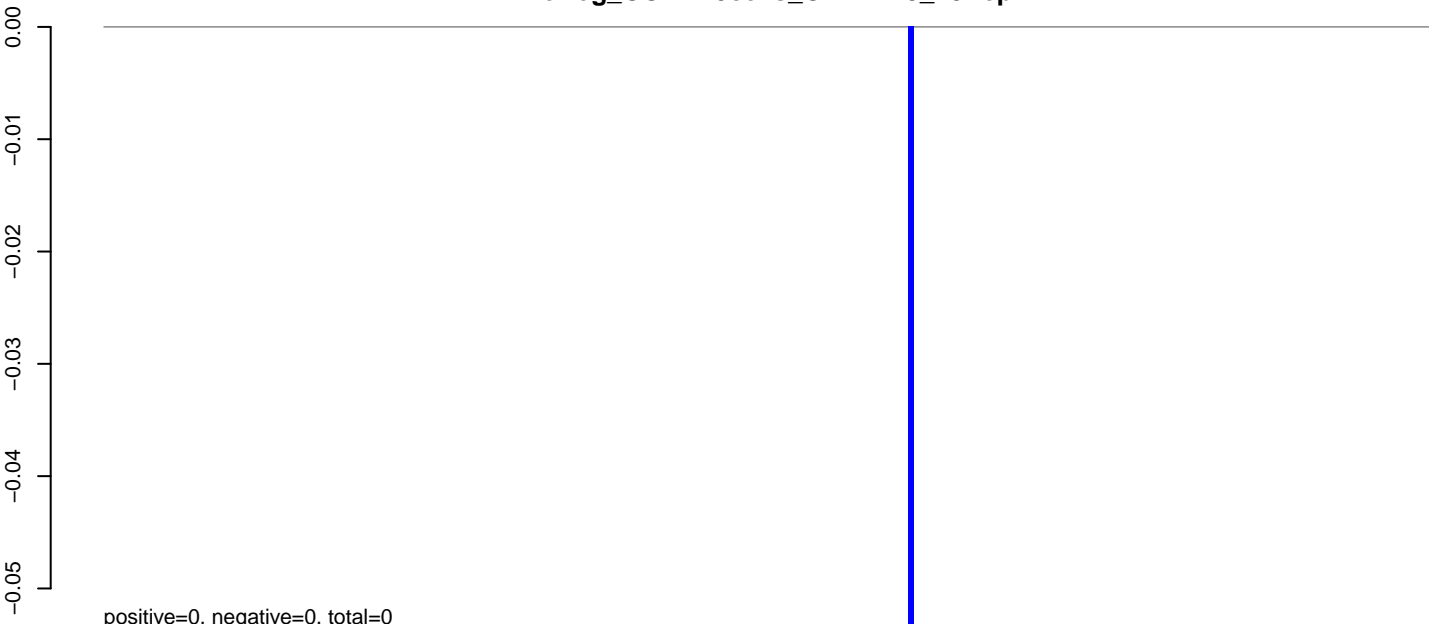


positive=0, negative=0, total=0

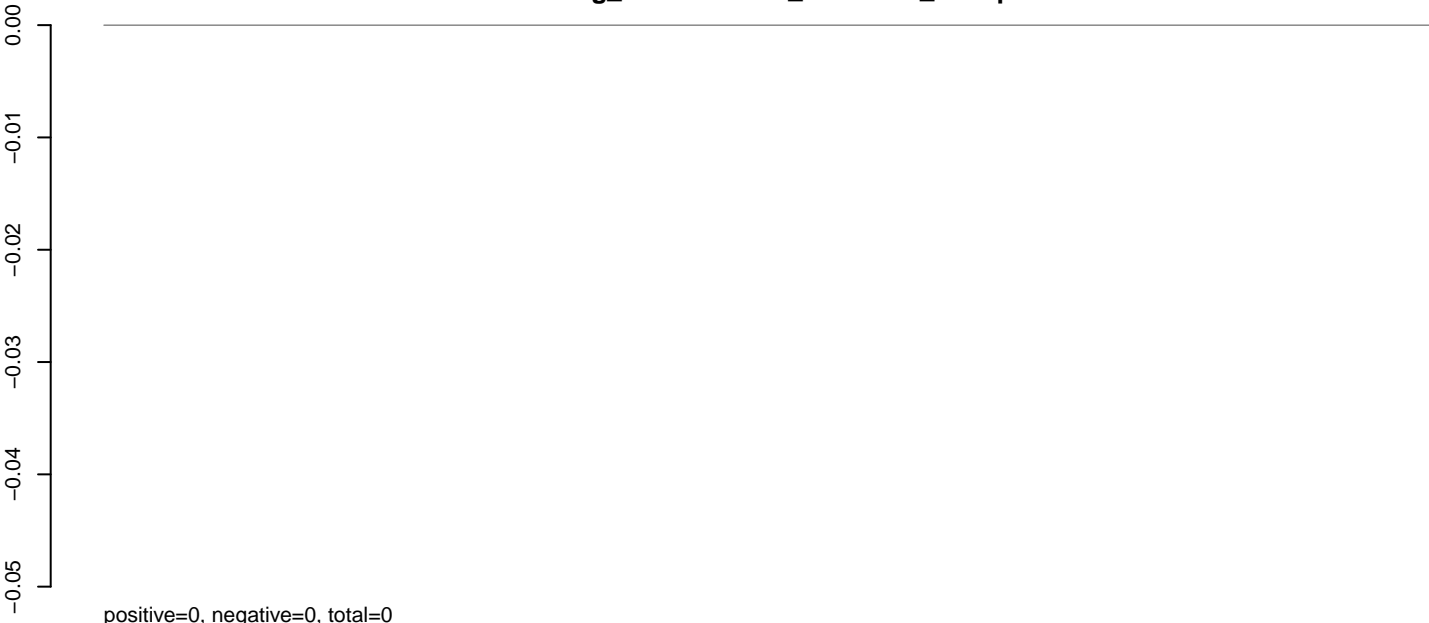
Window size=50, length=11824, virus@NC_003900.1-Aura_virus:1-11824

0 2000 4000 6000 8000 10000 12000

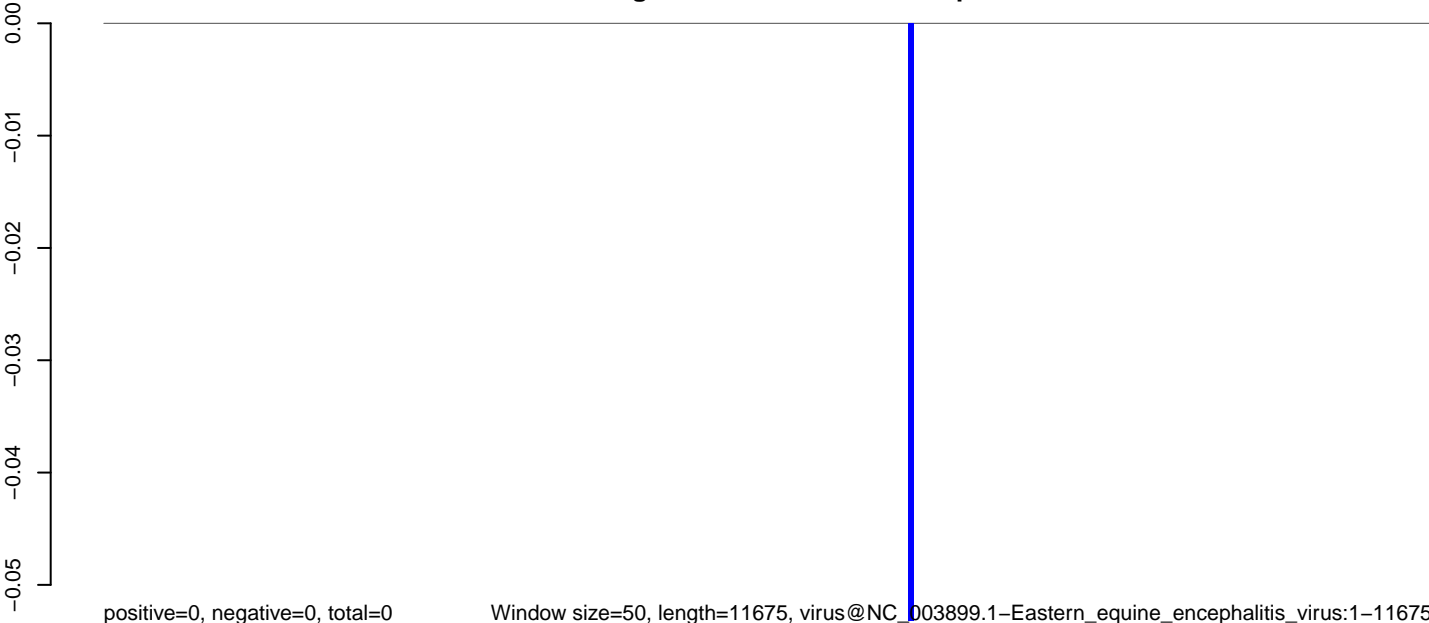
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



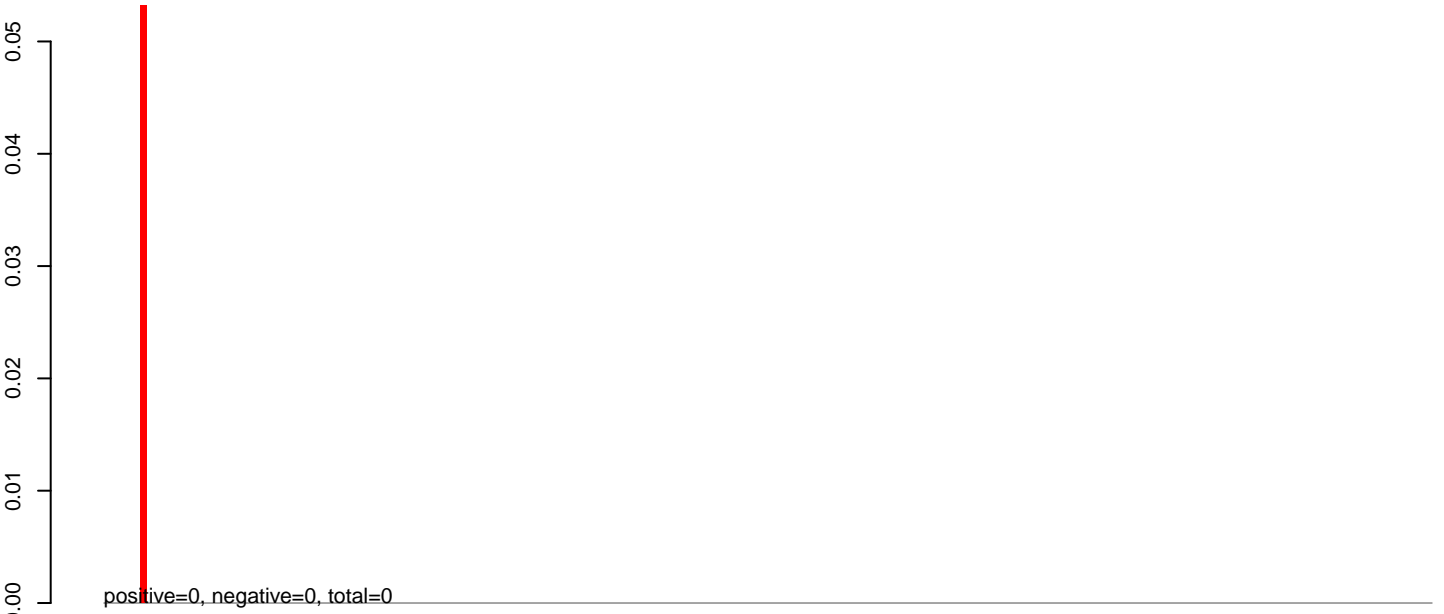
AeAeg_CCL.125cells_CHIKV.rep



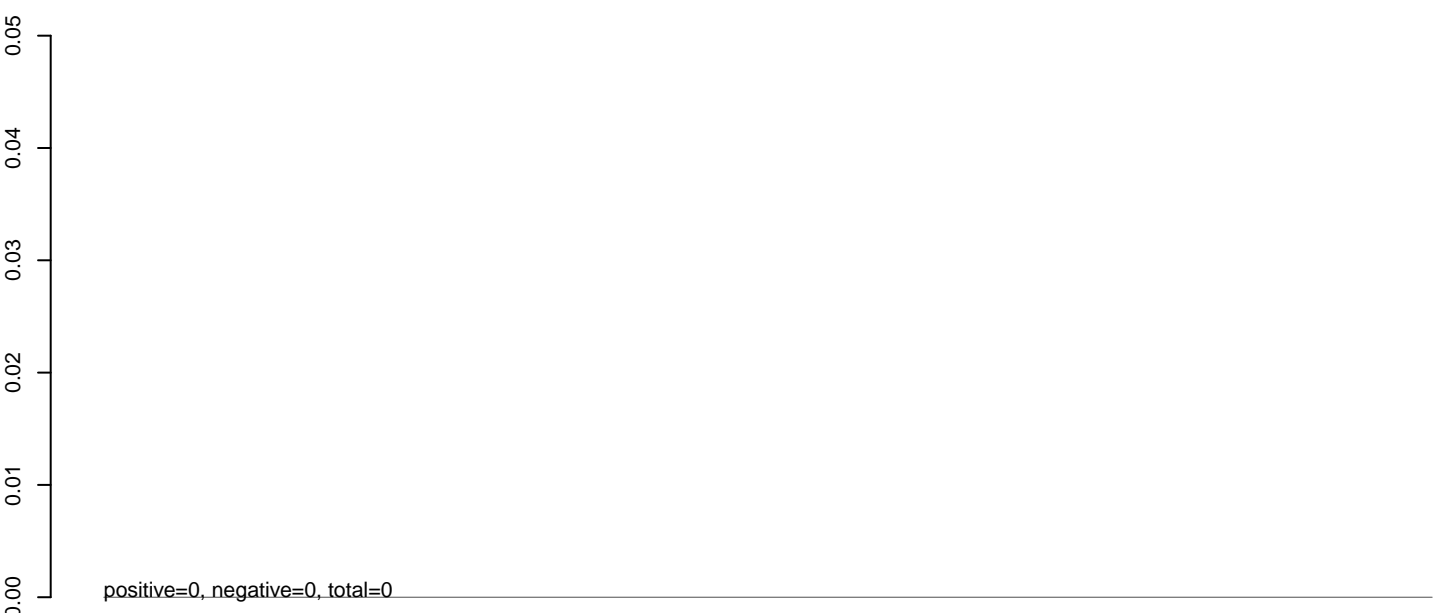
Window size=50, length=11675, virus@NC_003899.1-Eastern_equine_encephalitis_virus:1-11675

0 2000 4000 6000 8000 10000 12000

AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



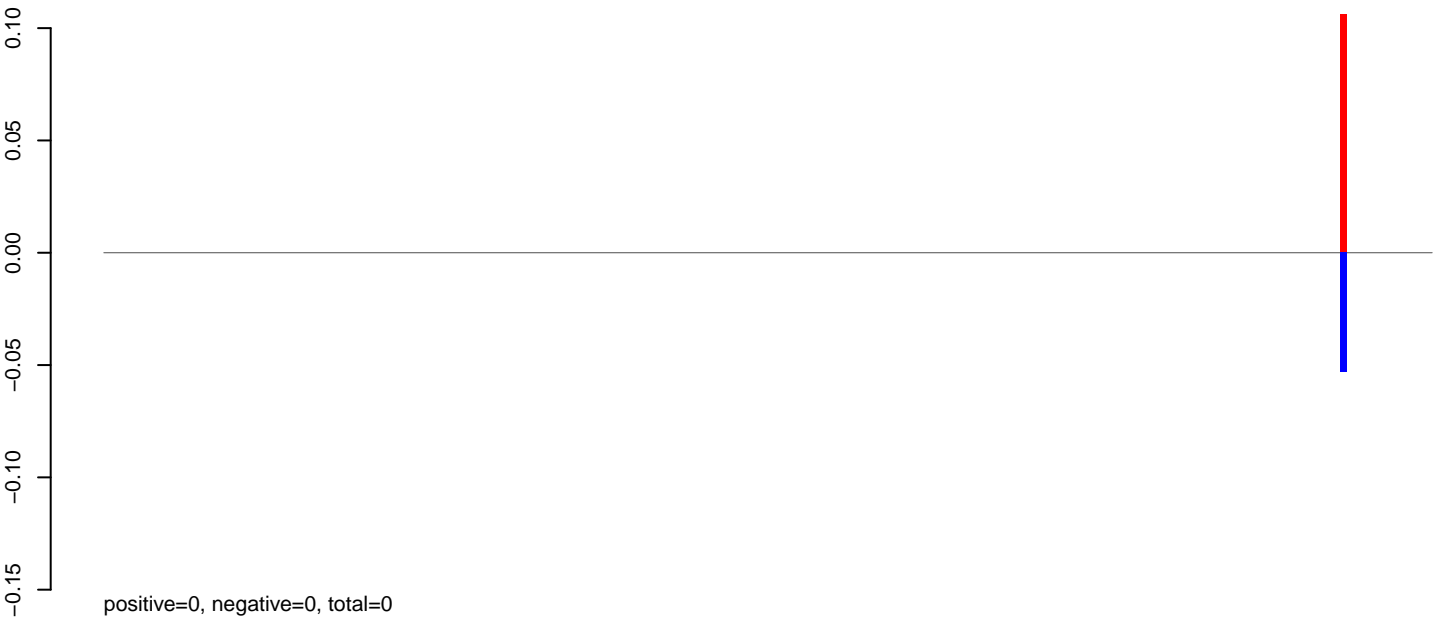
AeAeg_CCL.125cells_CHIKV.rep



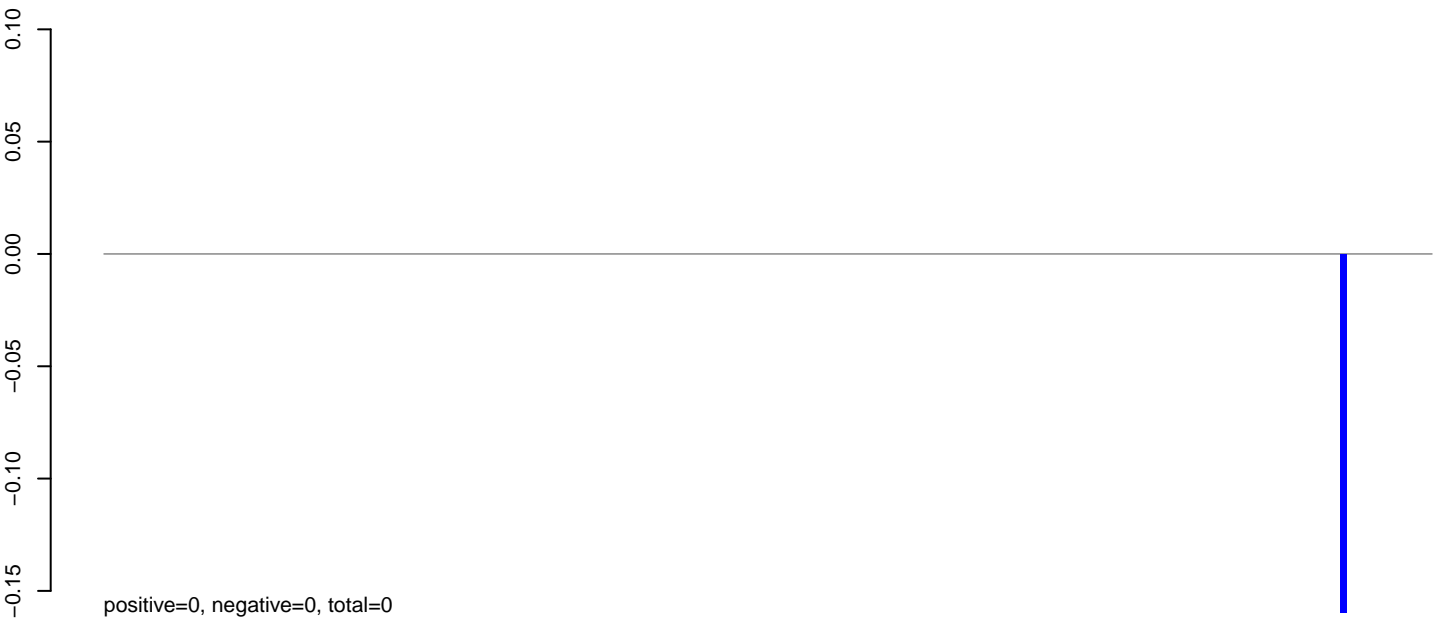
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0 2000 4000 6000 8000 10000

AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



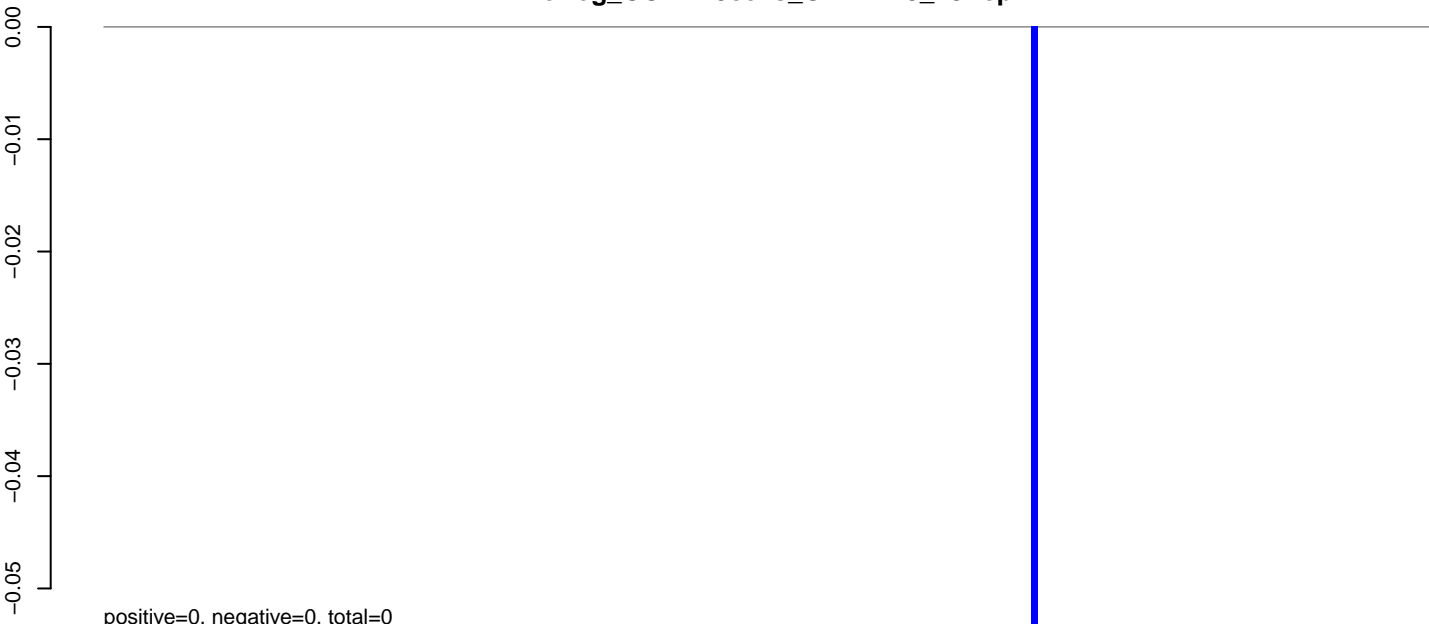
AeAeg_CCL.125cells_CHIKV.rep



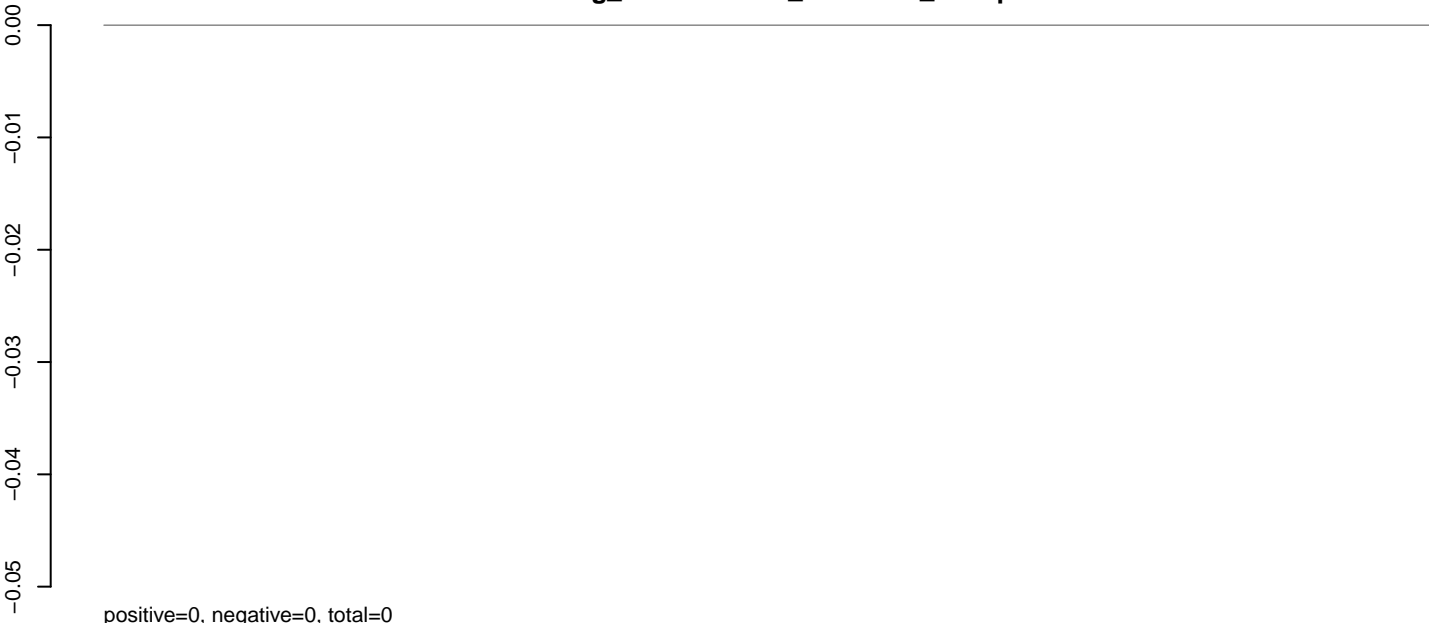
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0 2000 4000 6000 8000 10000

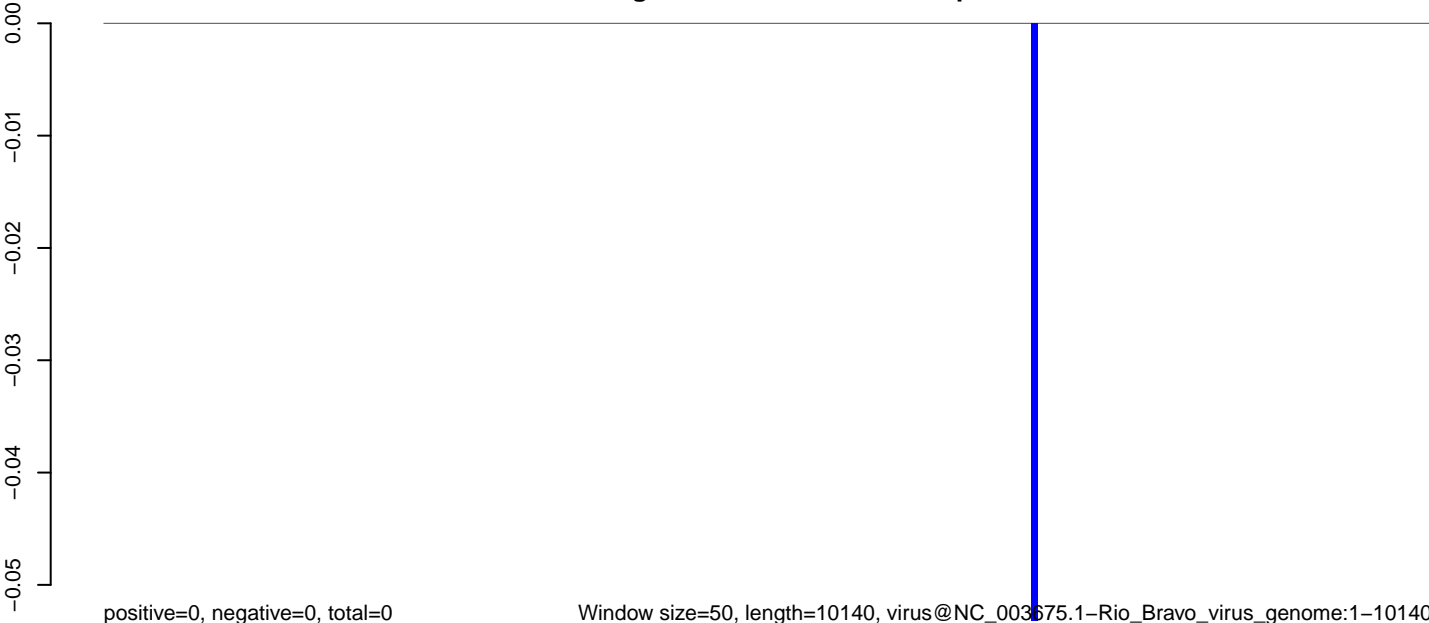
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



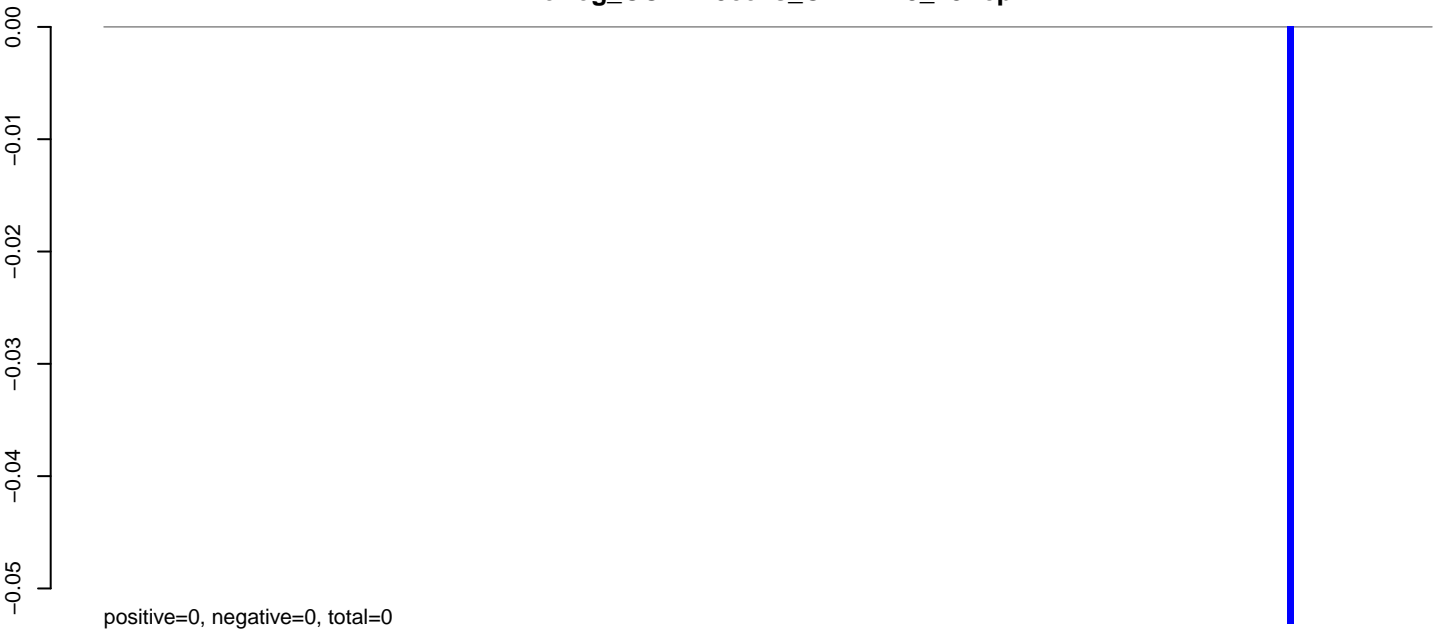
AeAeg_CCL.125cells_CHIKV.rep



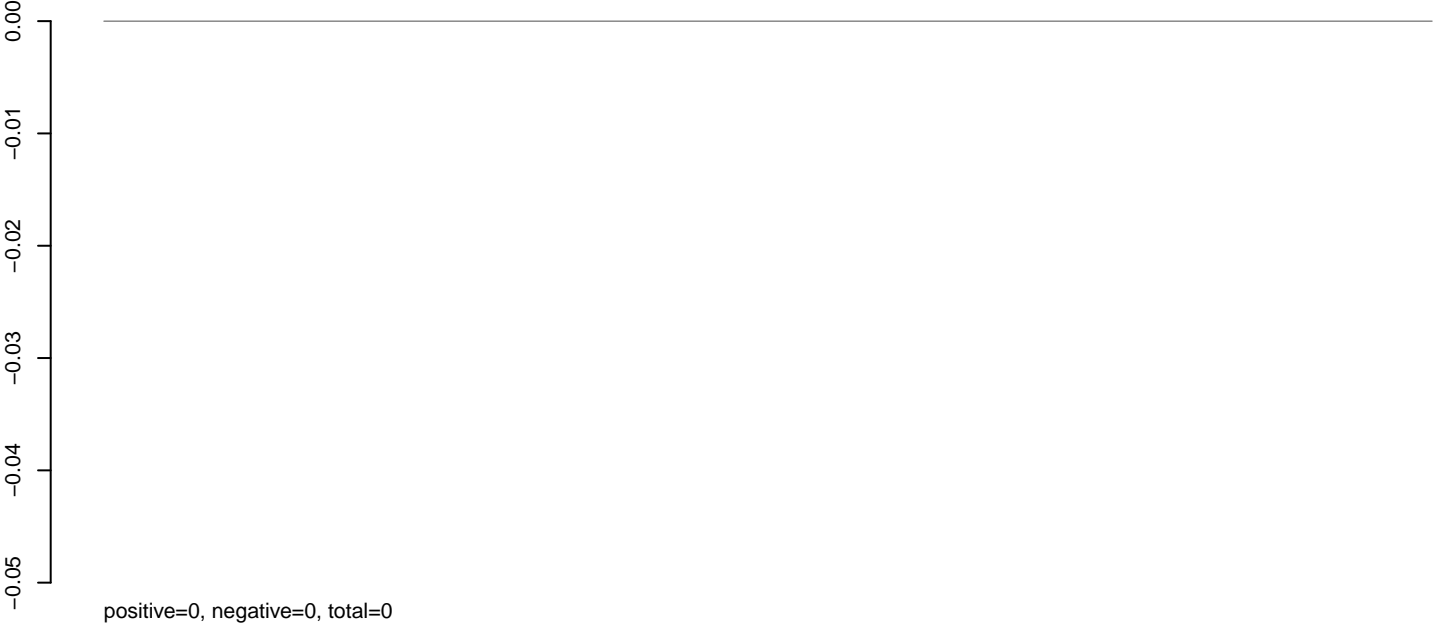
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0 2000 4000 6000 8000 10000

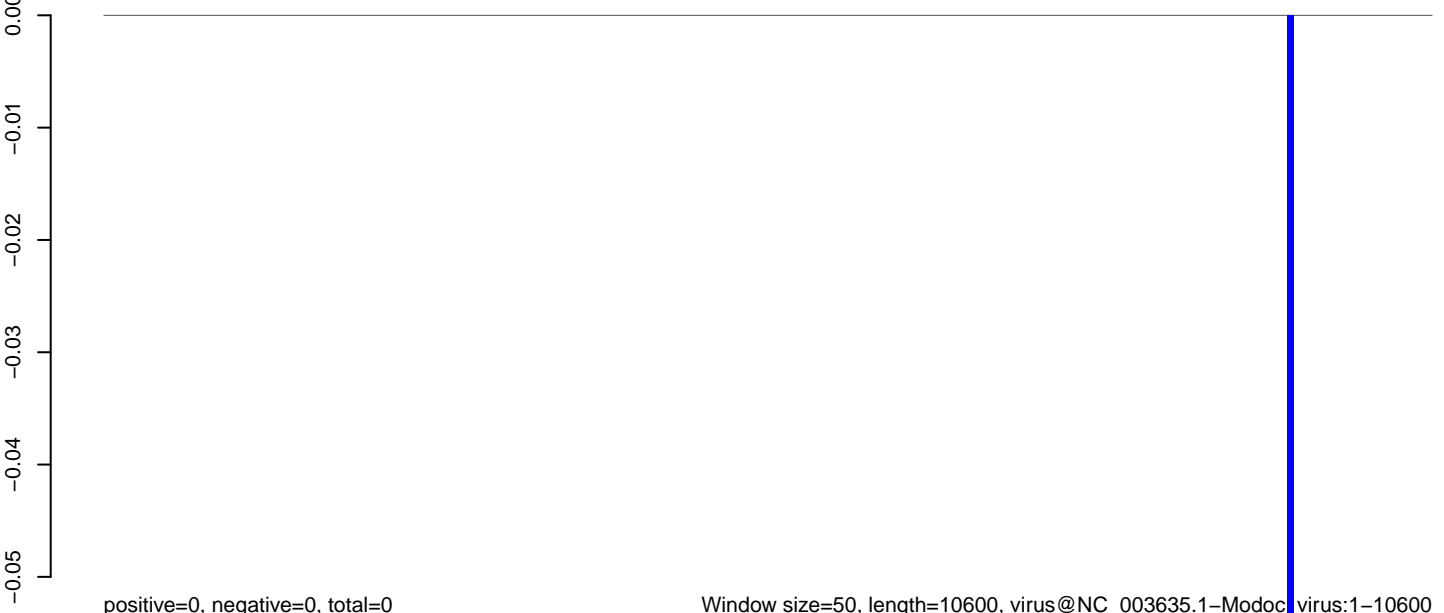
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

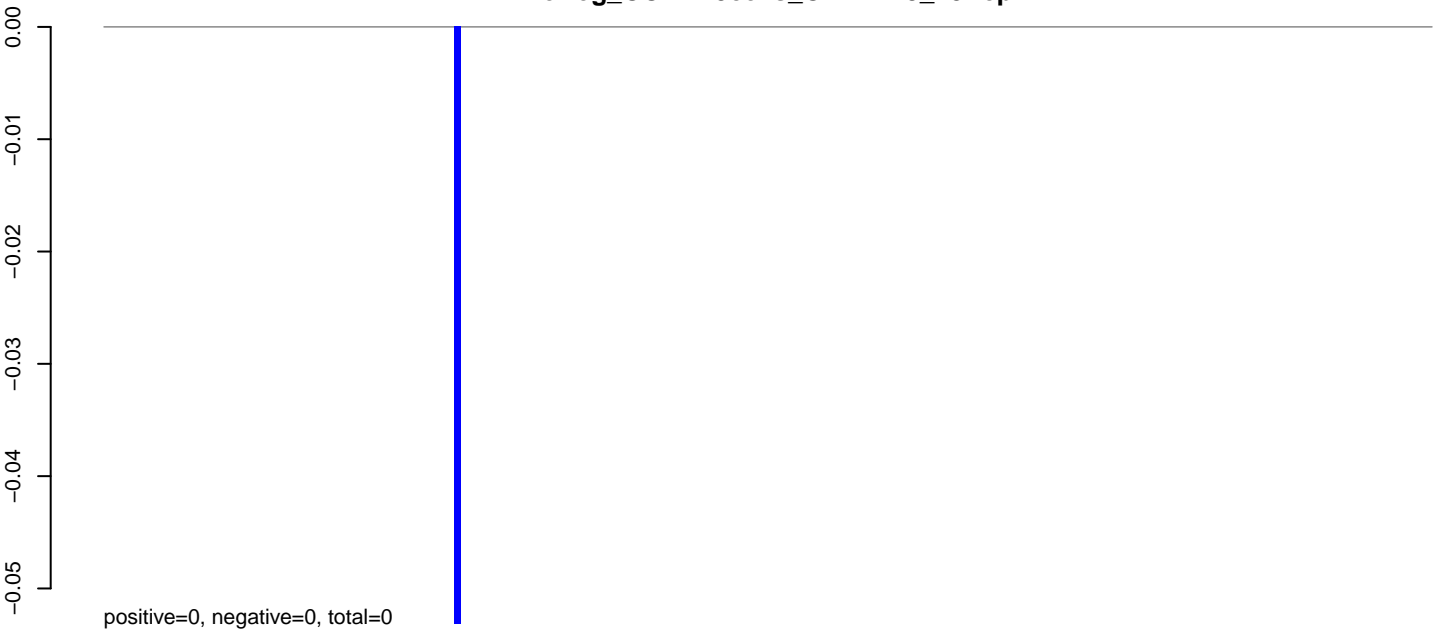


AeAeg_CCL.125cells_CHIKV.rep

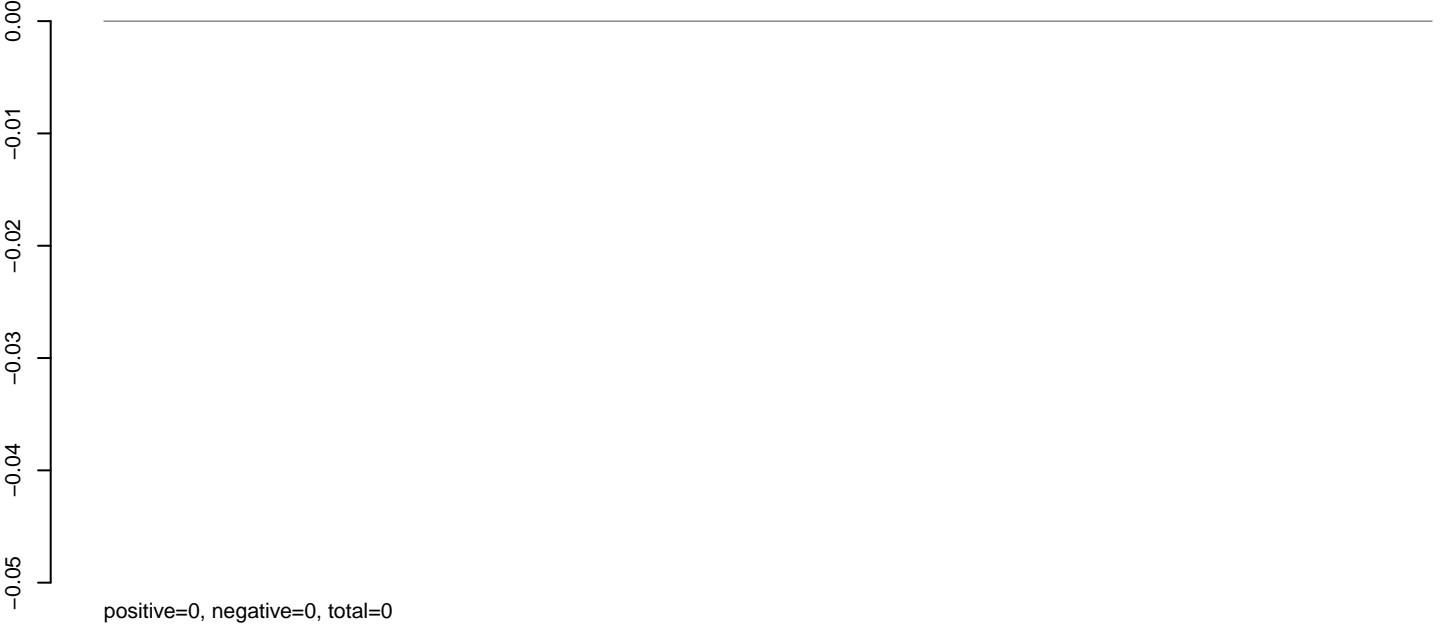


0 2000 4000 6000 8000 10000

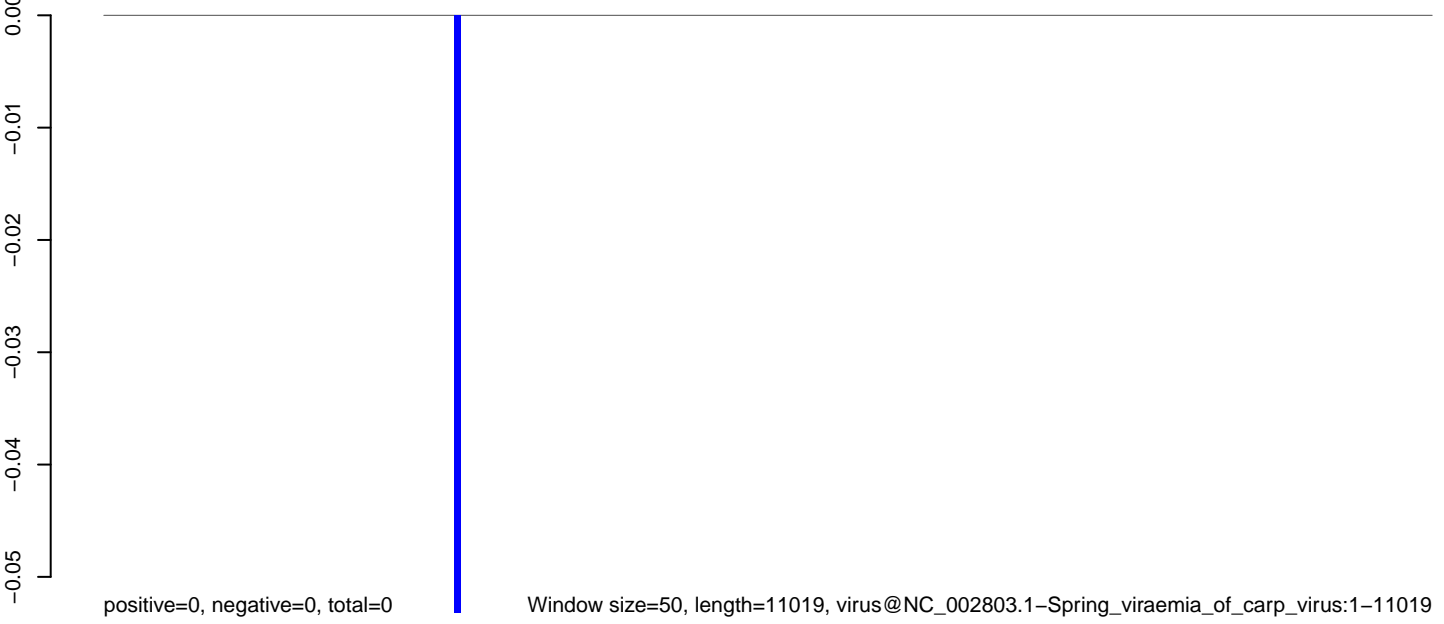
AeAeg_CCL.125cells_CHIKV.18_23.rep



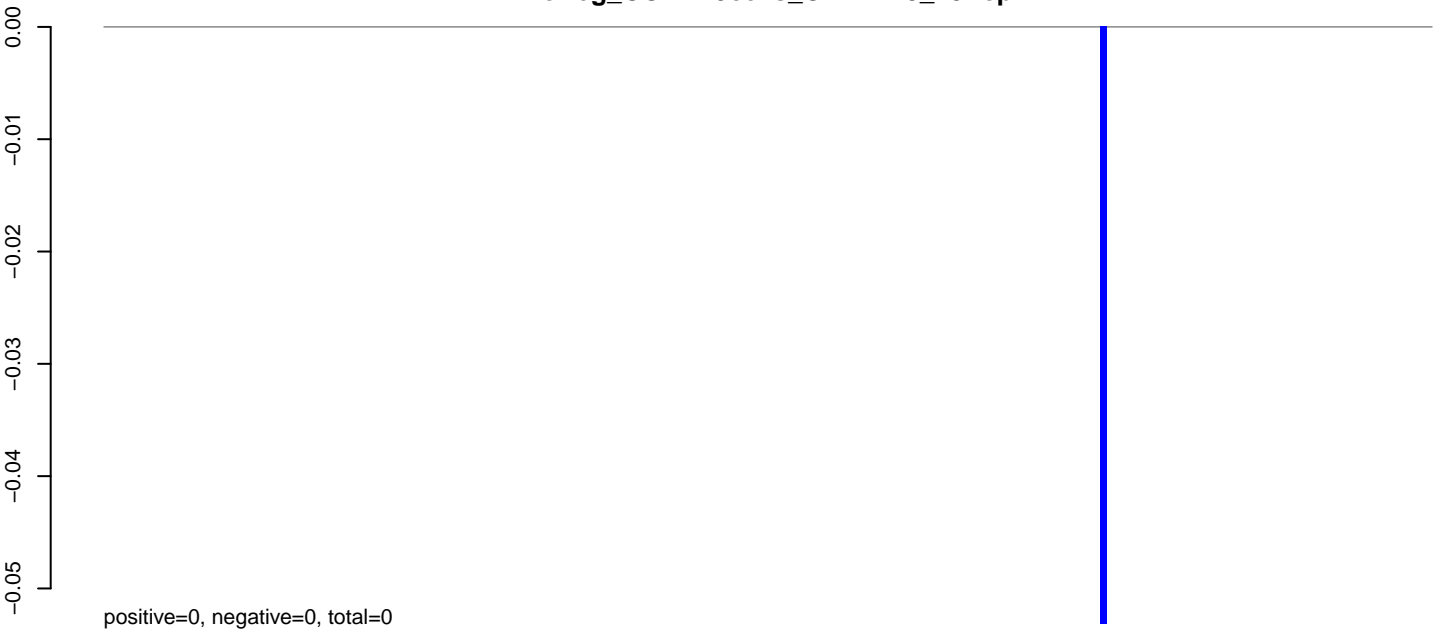
AeAeg_CCL.125cells_CHIKV.24_35.rep



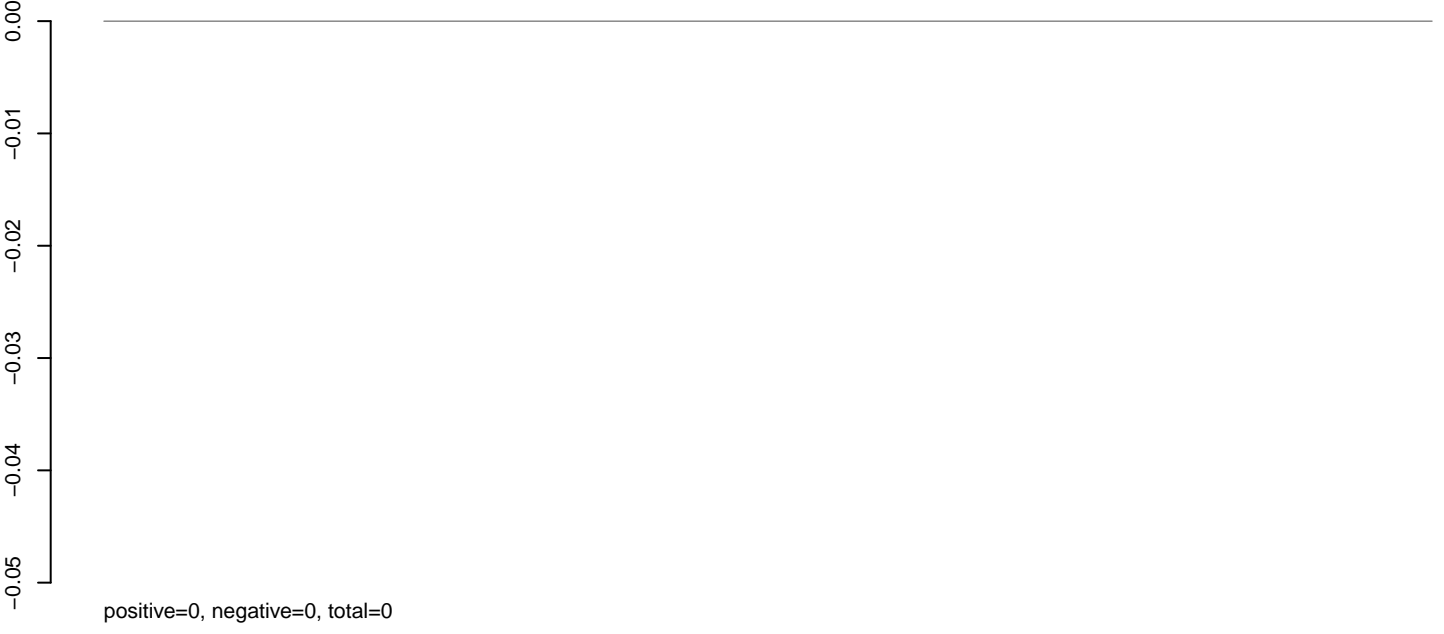
AeAeg_CCL.125cells_CHIKV.rep



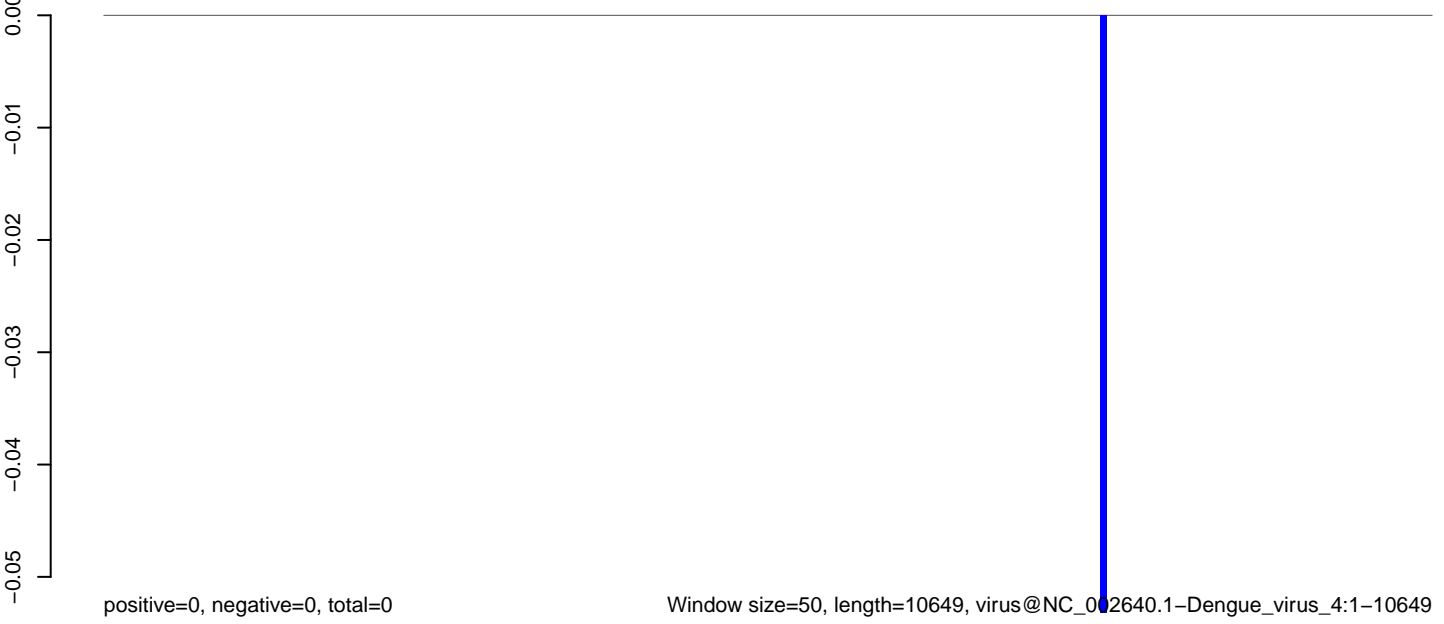
AeAeg_CCL.125cells_CHIKV.18_23.rep



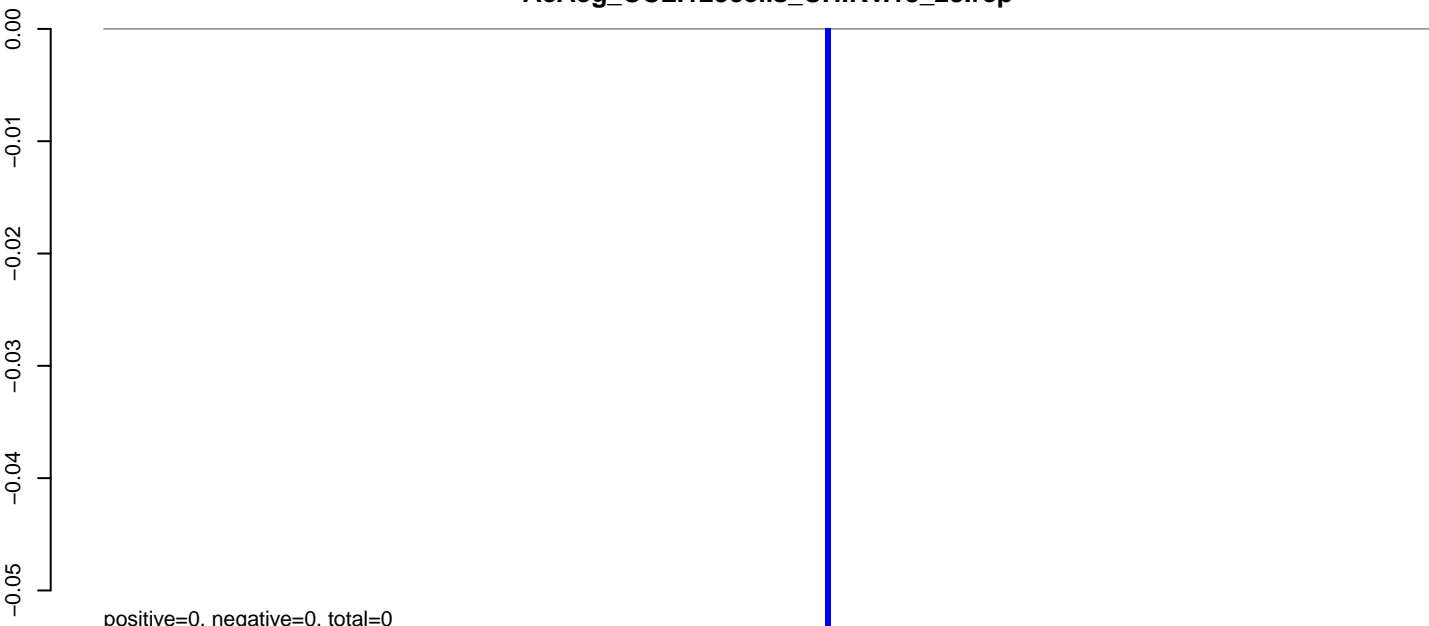
AeAeg_CCL.125cells_CHIKV.24_35.rep



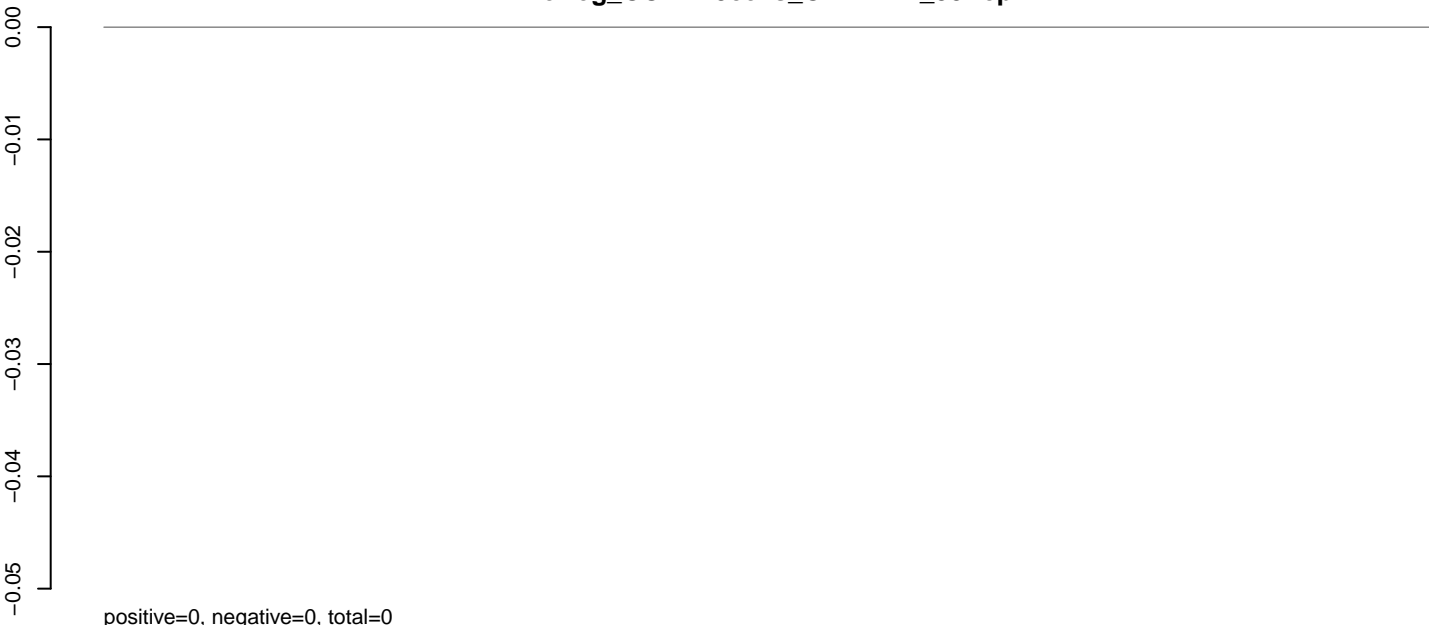
AeAeg_CCL.125cells_CHIKV.rep



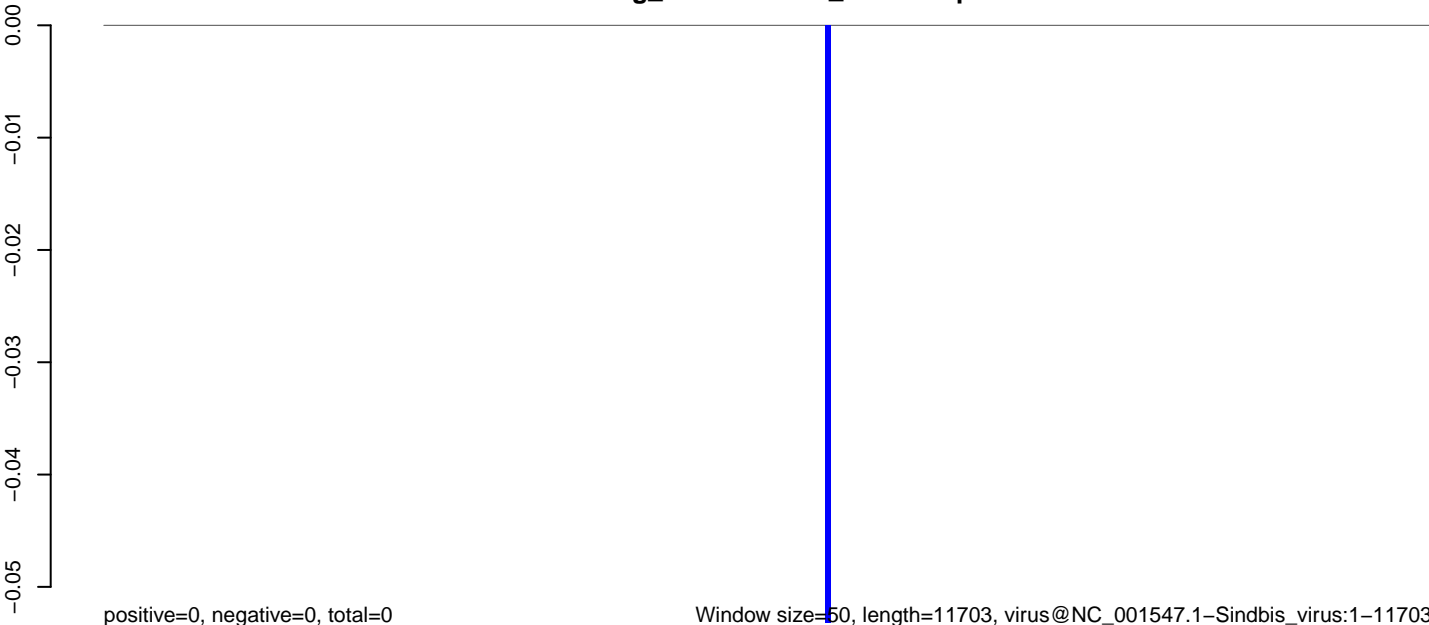
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

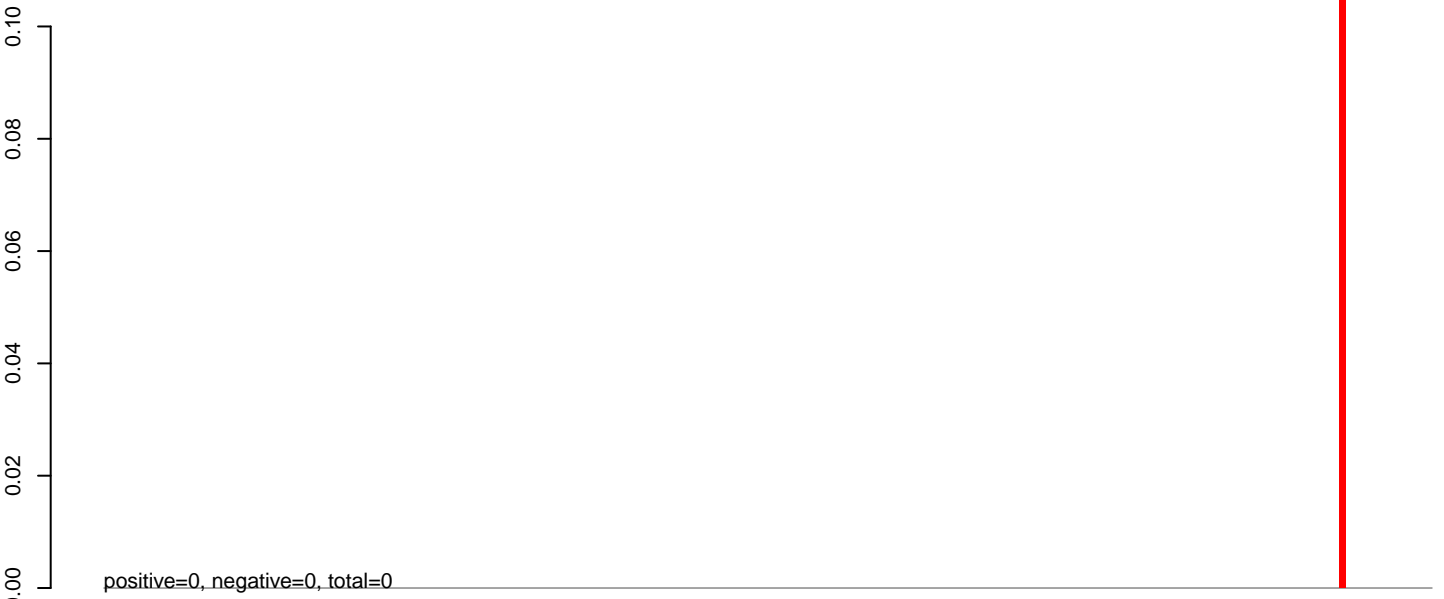


AeAeg_CCL.125cells_CHIKV.rep

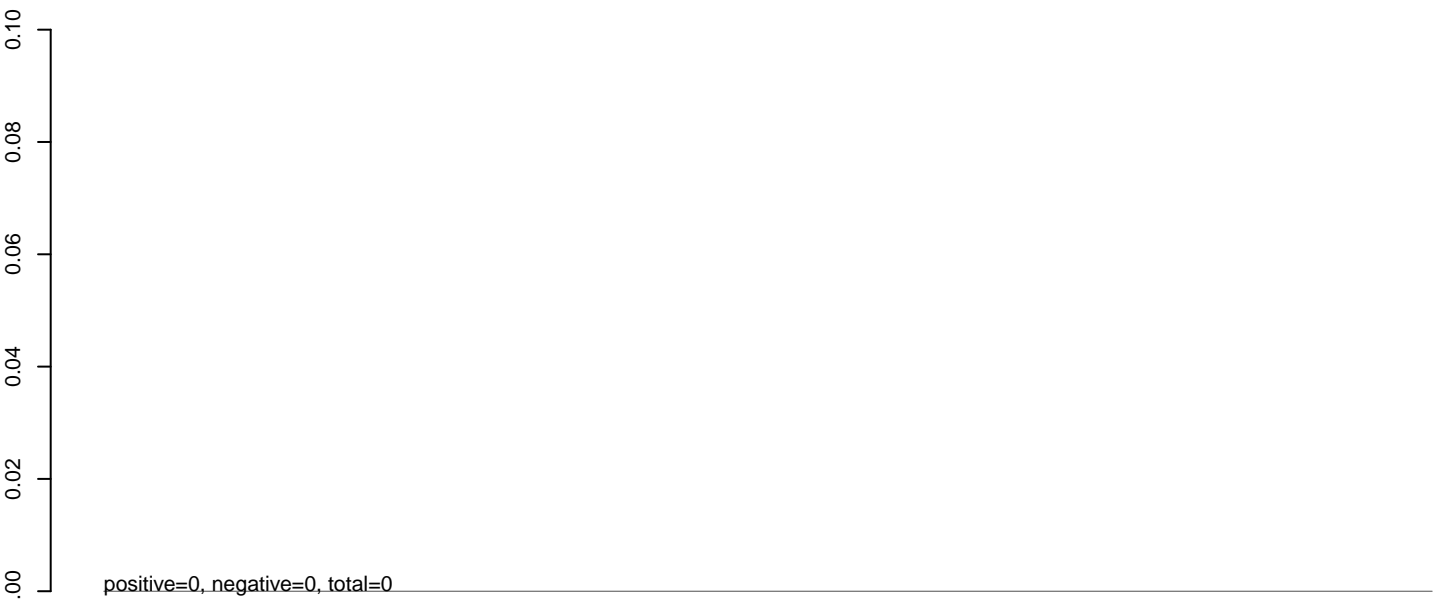


0 2000 4000 6000 8000 10000 12000

AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



AeAeg_CCL.125cells_CHIKV.rep

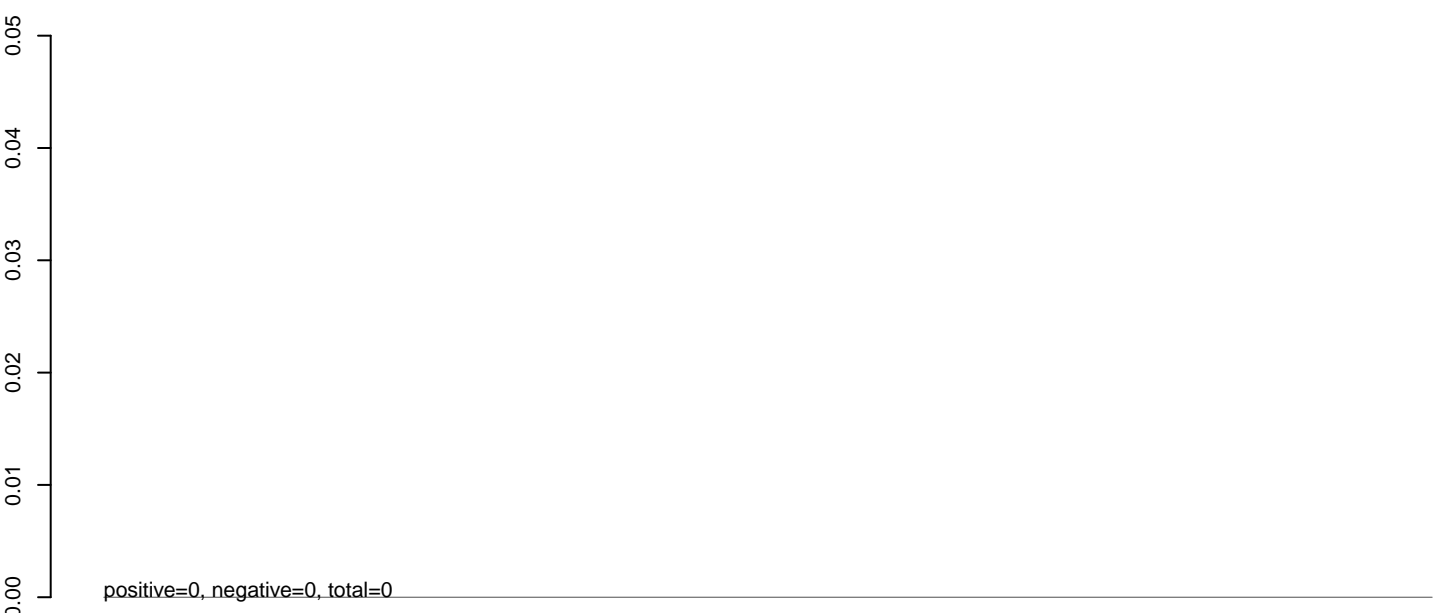


0 2000 4000 6000 8000 10000

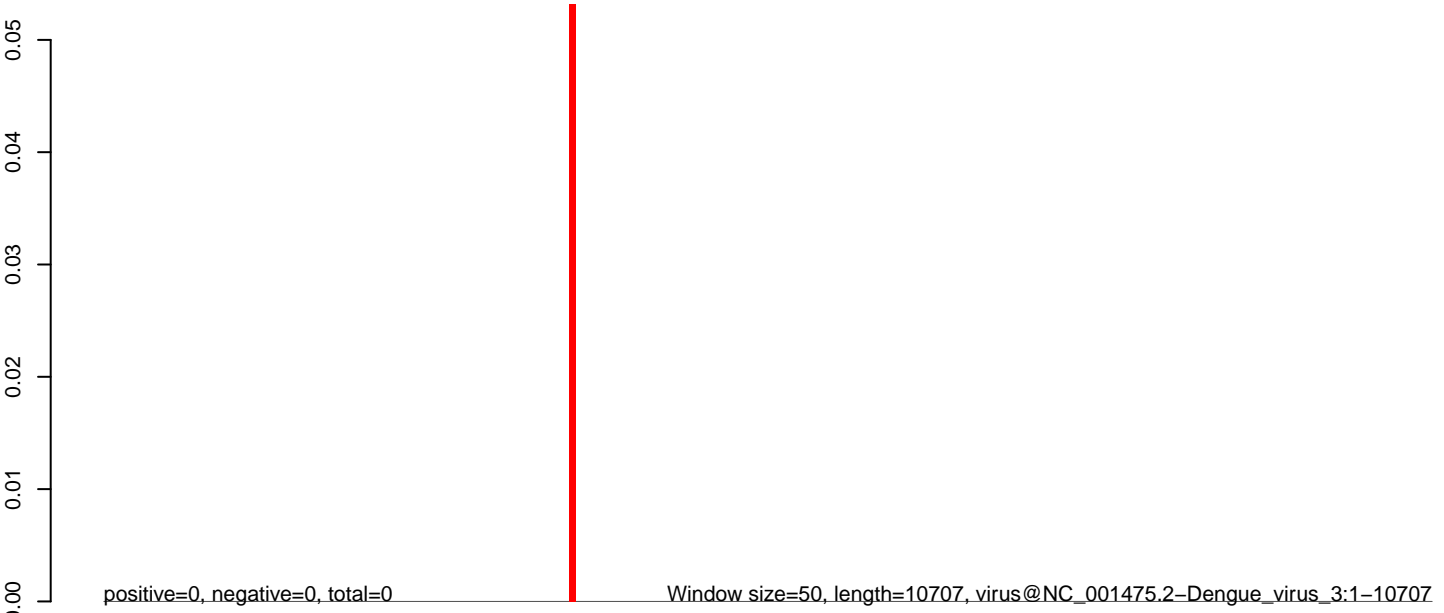
AeAeg_CCL.125cells_CHIKV.18_23.rep



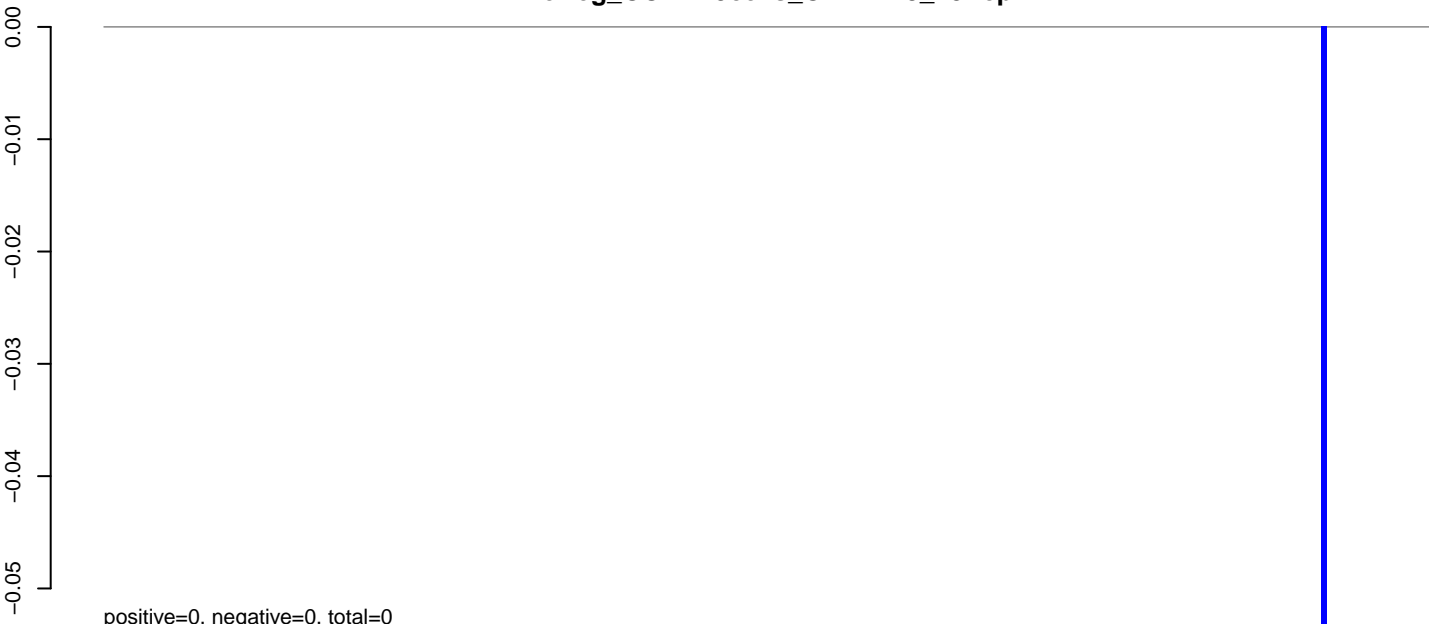
AeAeg_CCL.125cells_CHIKV.24_35.rep



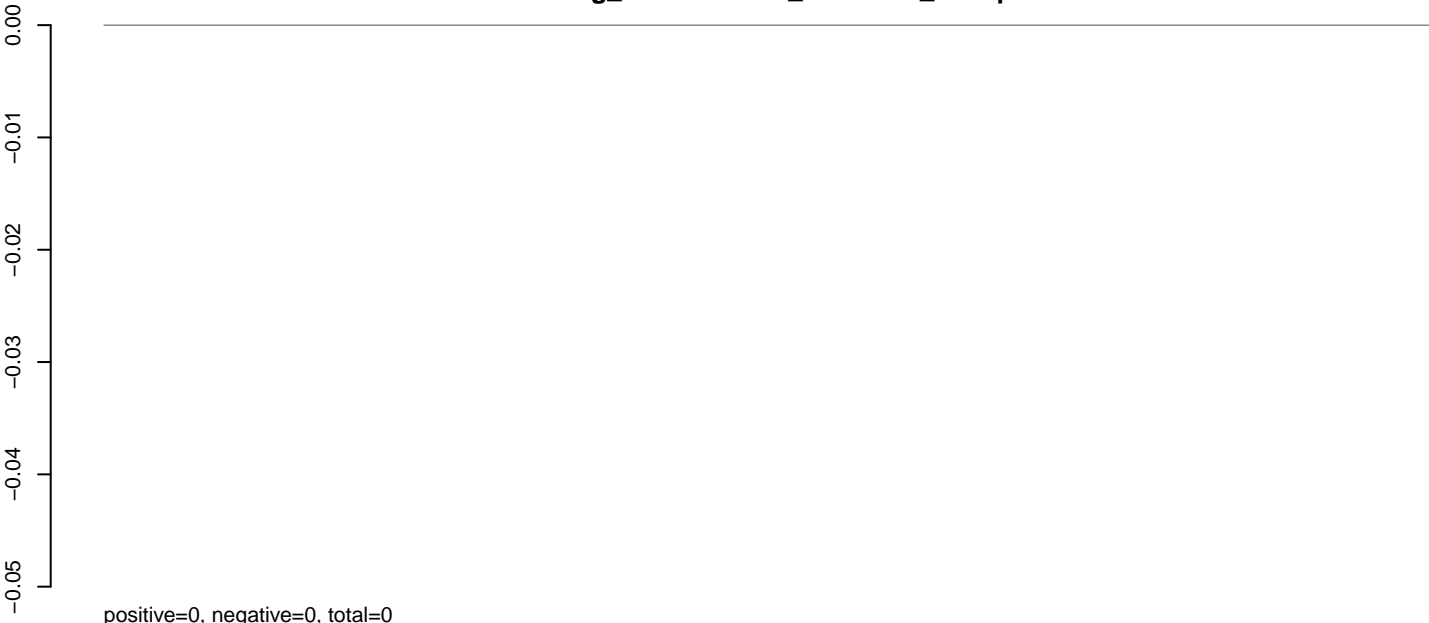
AeAeg_CCL.125cells_CHIKV.rep



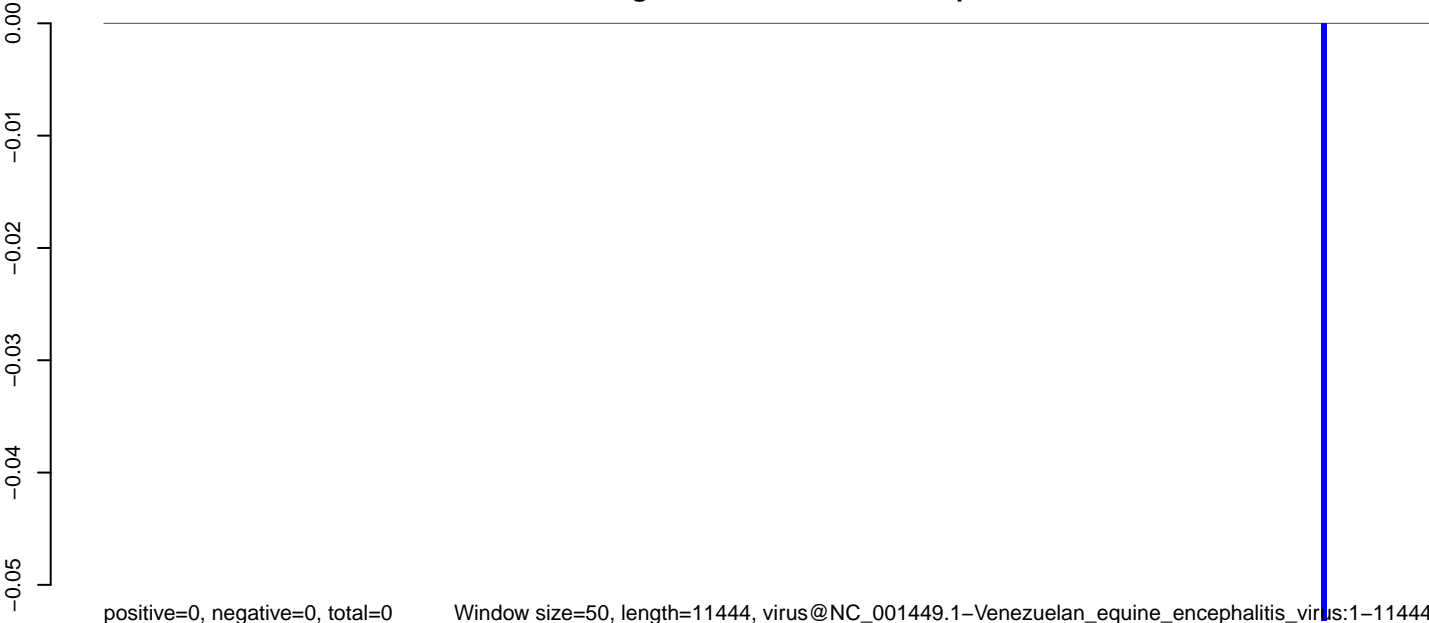
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



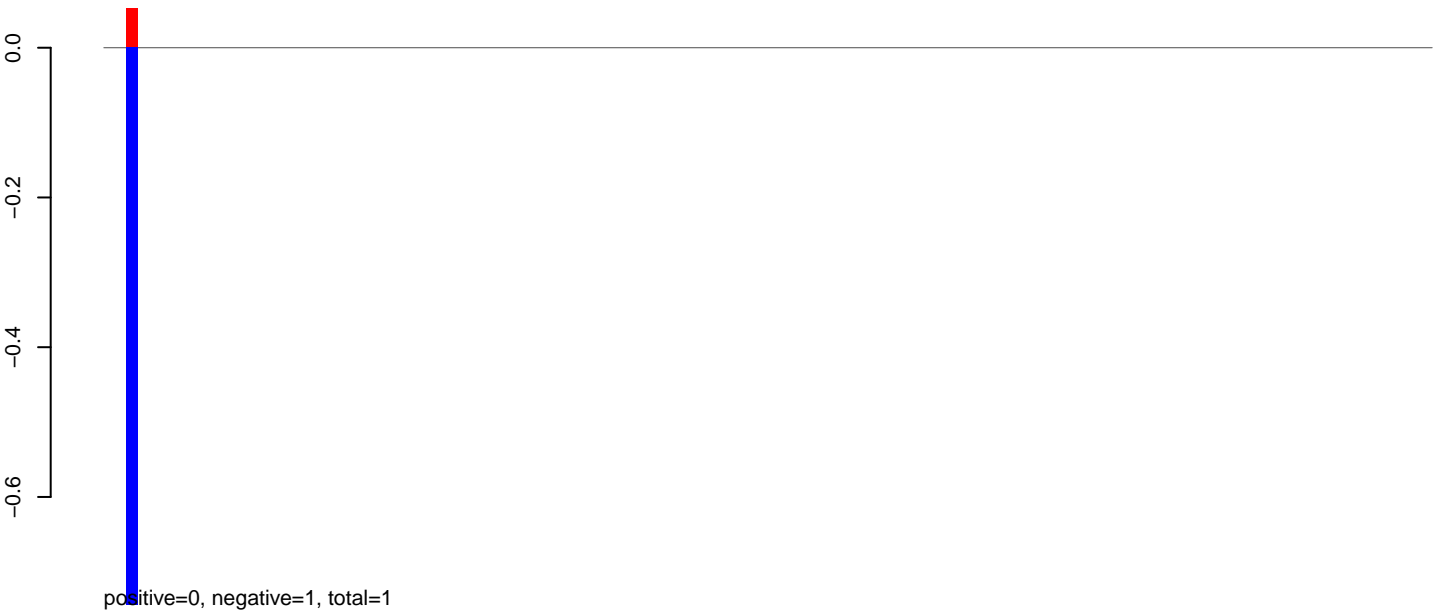
AeAeg_CCL.125cells_CHIKV.rep



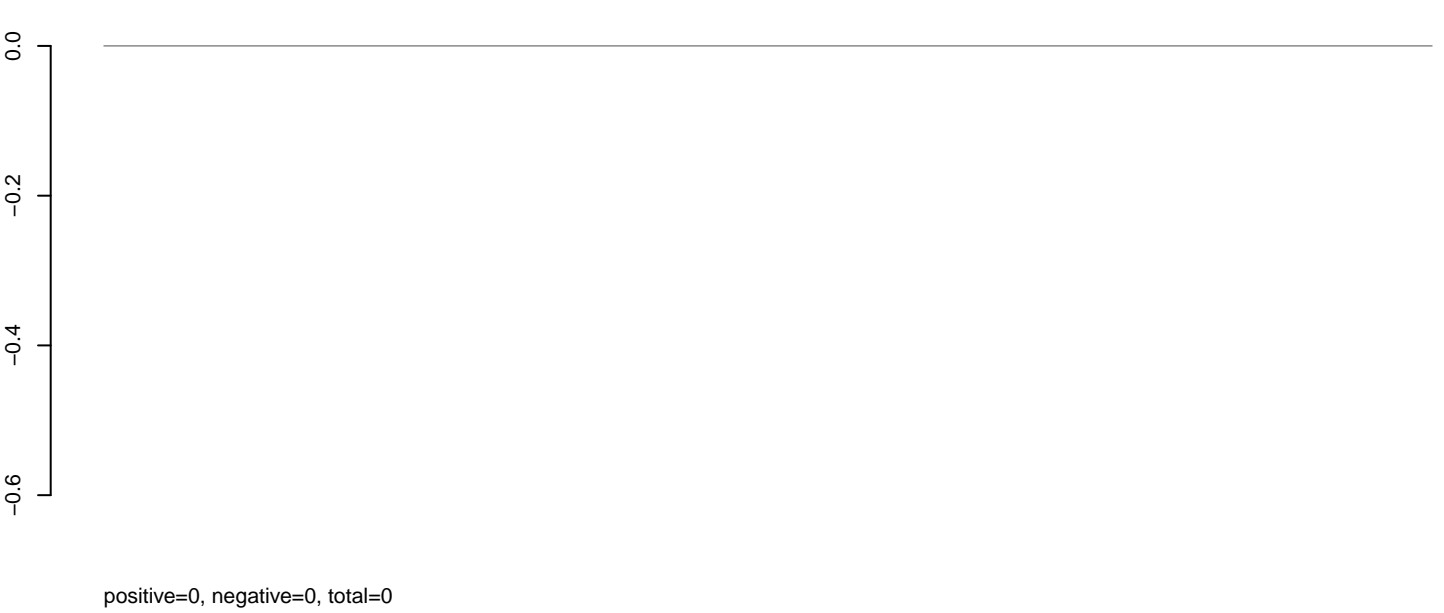
Window size=50, length=11444, virus@NC_001449.1-Venezuelan_equine_encephalitis_virus:1-11444

0 2000 4000 6000 8000 10000

AeAeg_CCL.125cells_CHIKV.18_23.rep



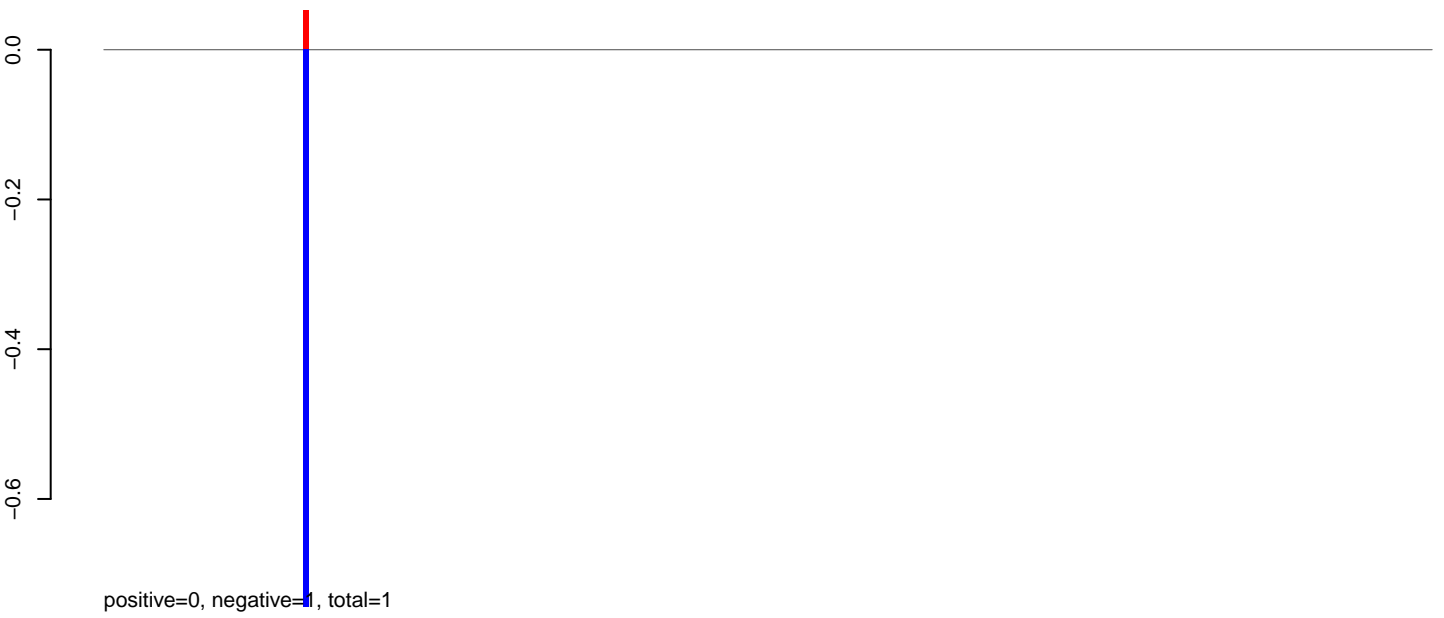
AeAeg_CCL.125cells_CHIKV.24_35.rep



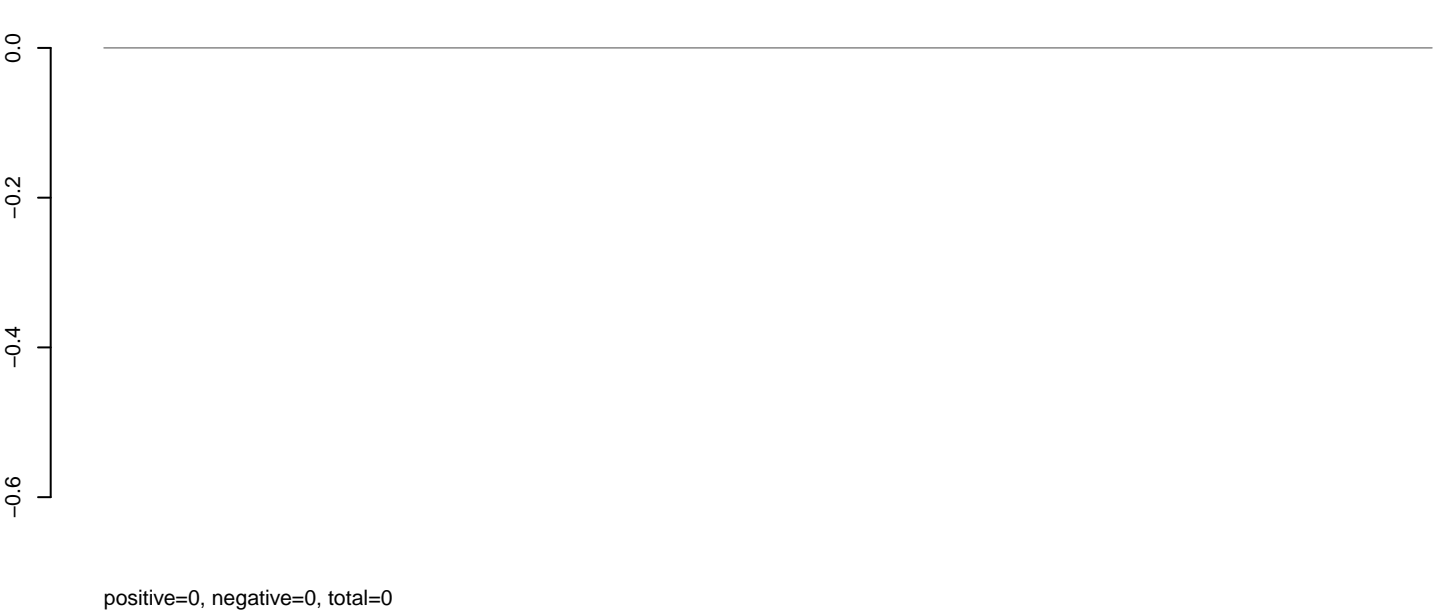
AeAeg_CCL.125cells_CHIKV.rep



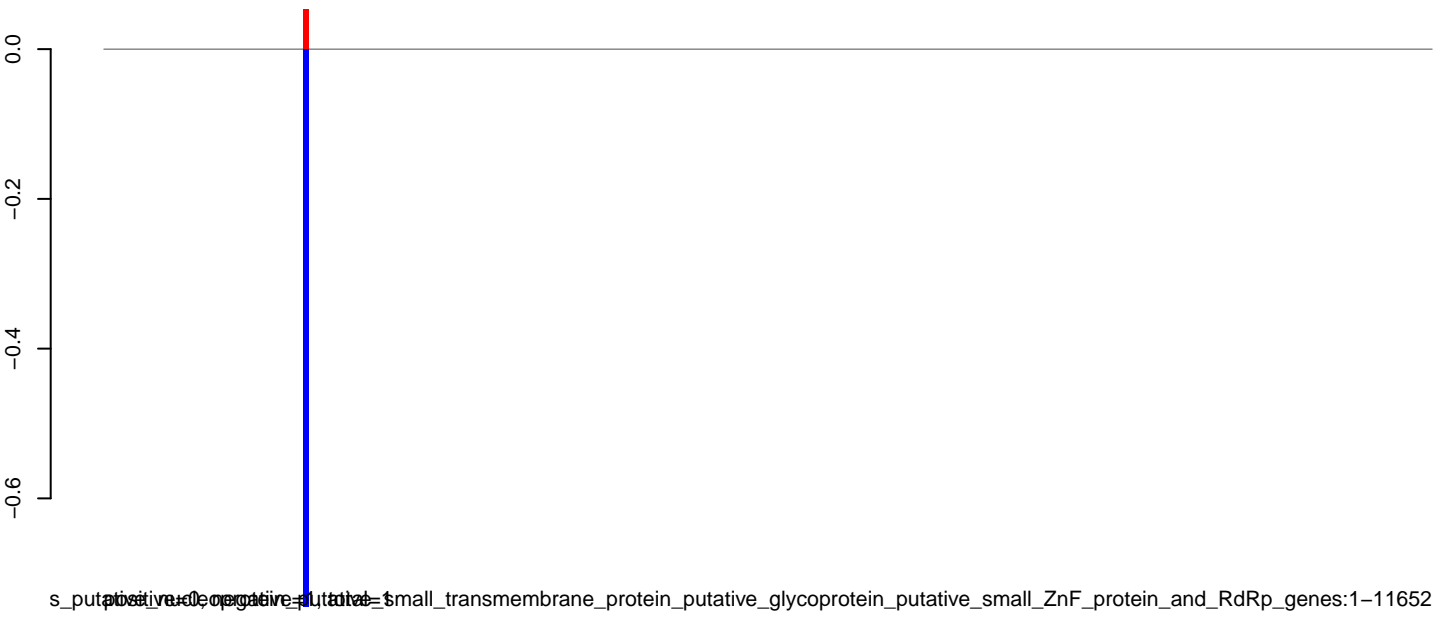
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

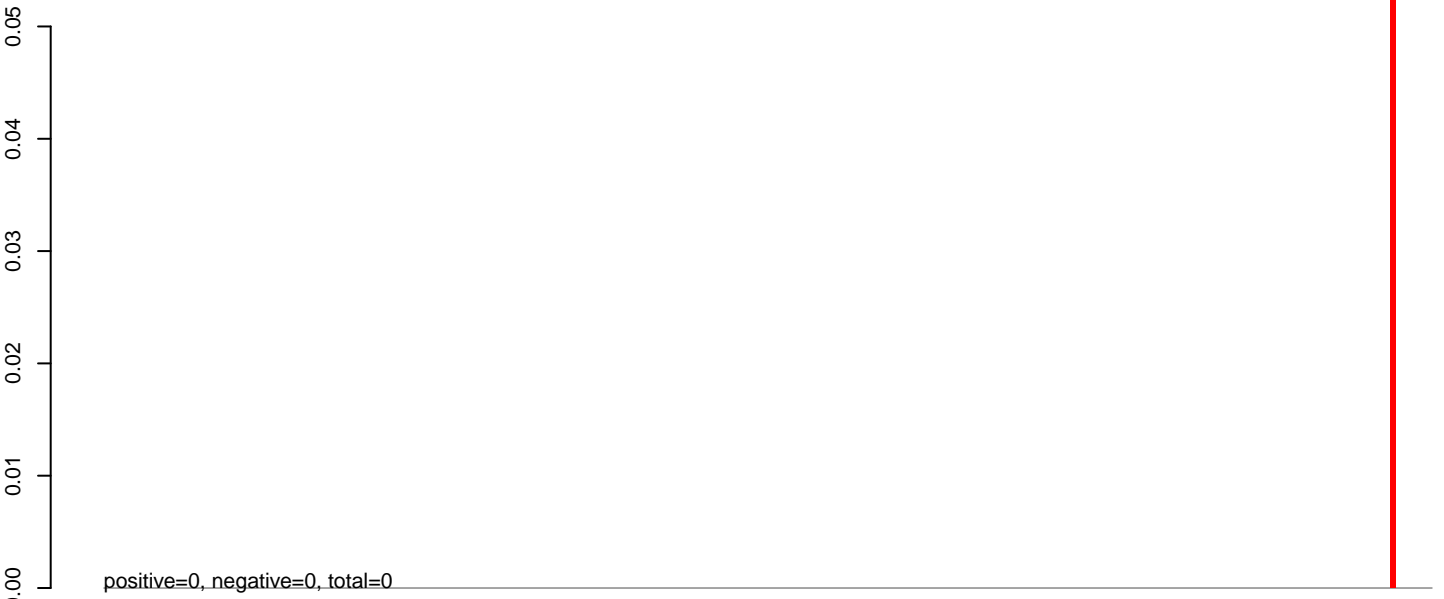


AeAeg_CCL.125cells_CHIKV.rep

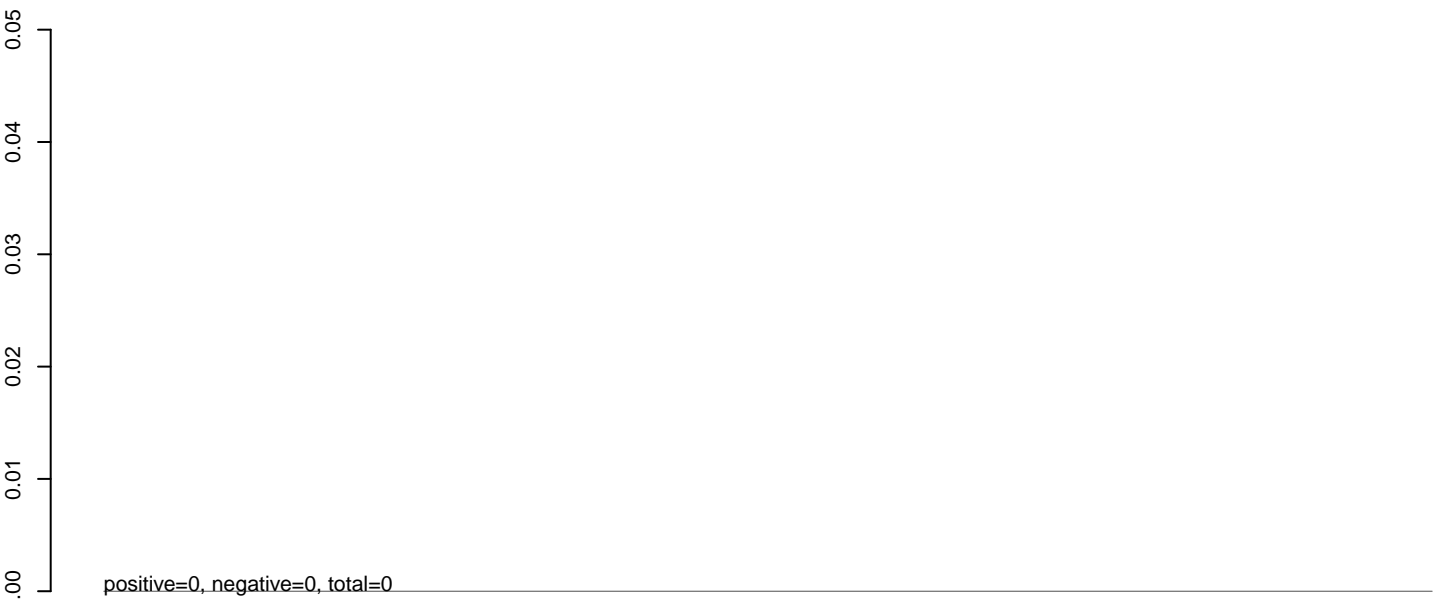


0 2000 4000 6000 8000 10000 12000

AeAeg_CCL.125cells_CHIKV.18_23.rep



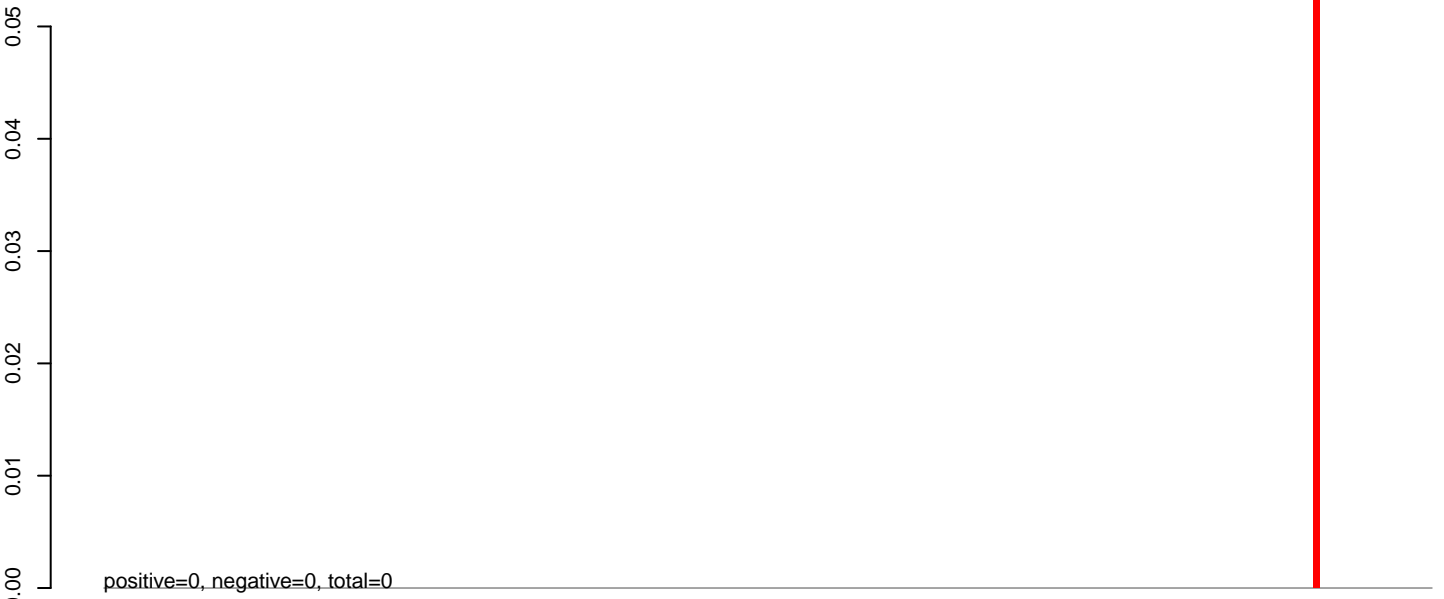
AeAeg_CCL.125cells_CHIKV.24_35.rep



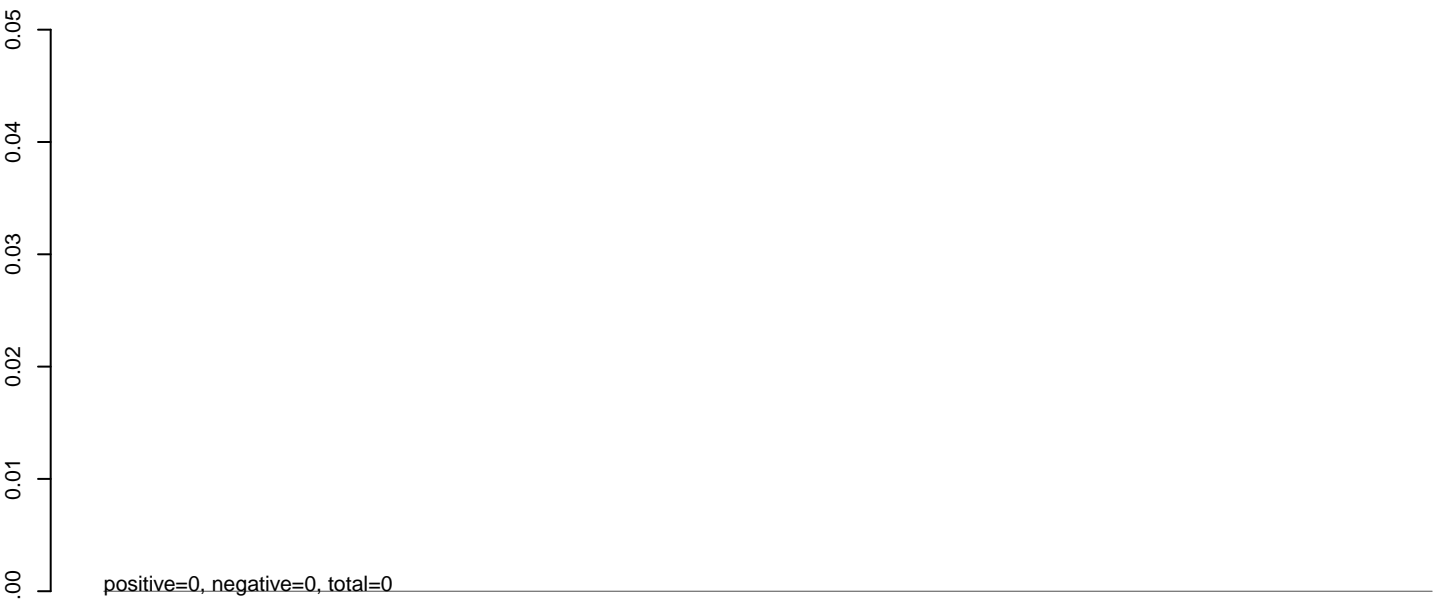
AeAeg_CCL.125cells_CHIKV.rep



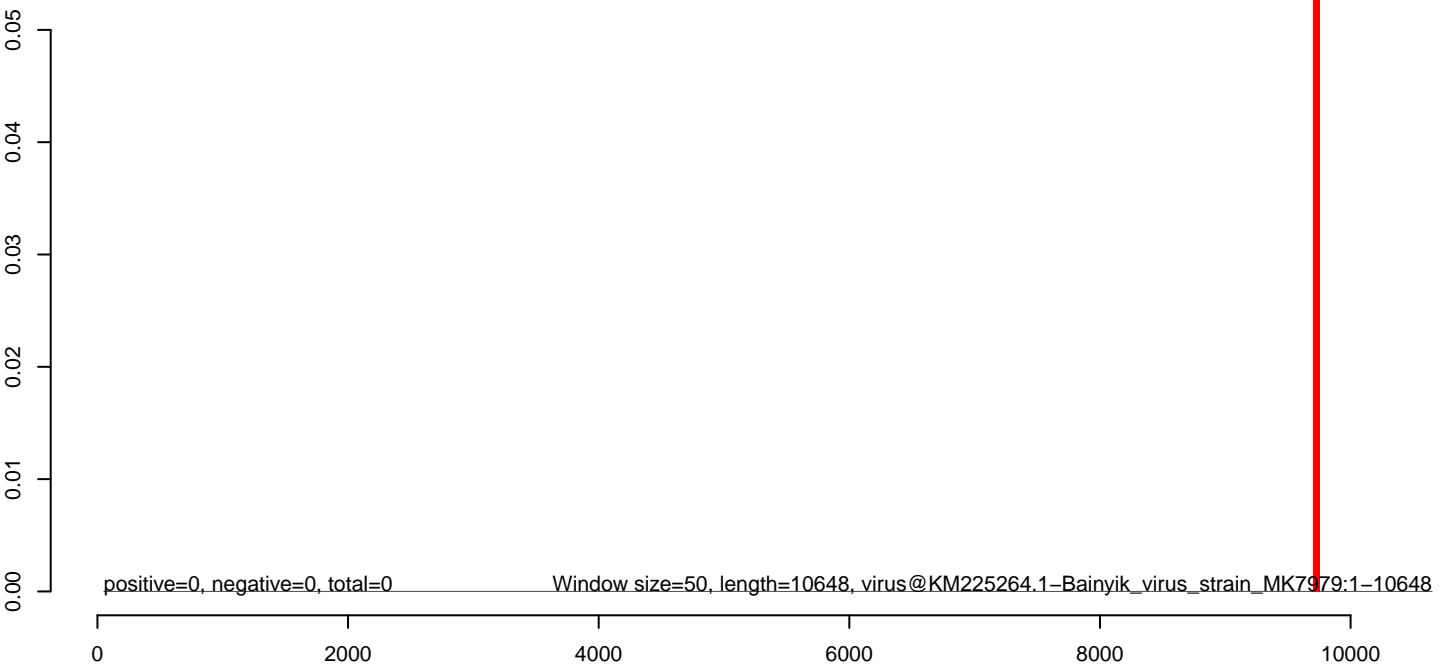
AeAeg_CCL.125cells_CHIKV.18_23.rep



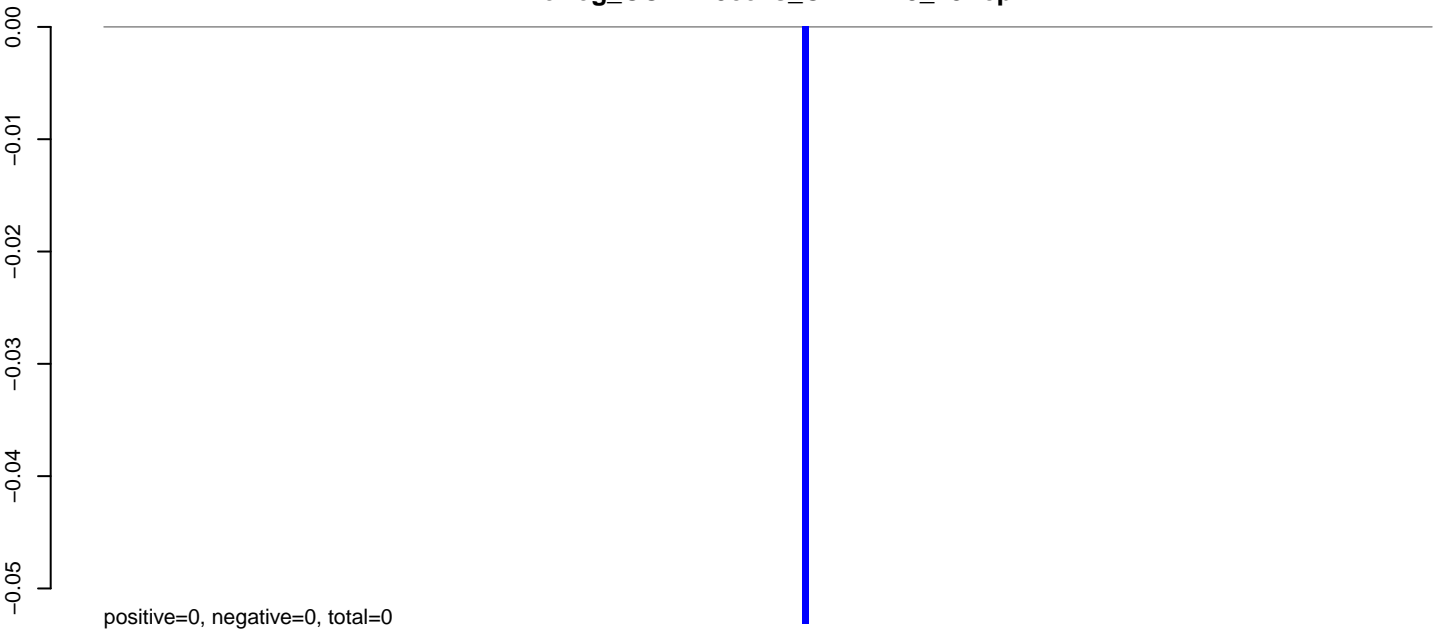
AeAeg_CCL.125cells_CHIKV.24_35.rep



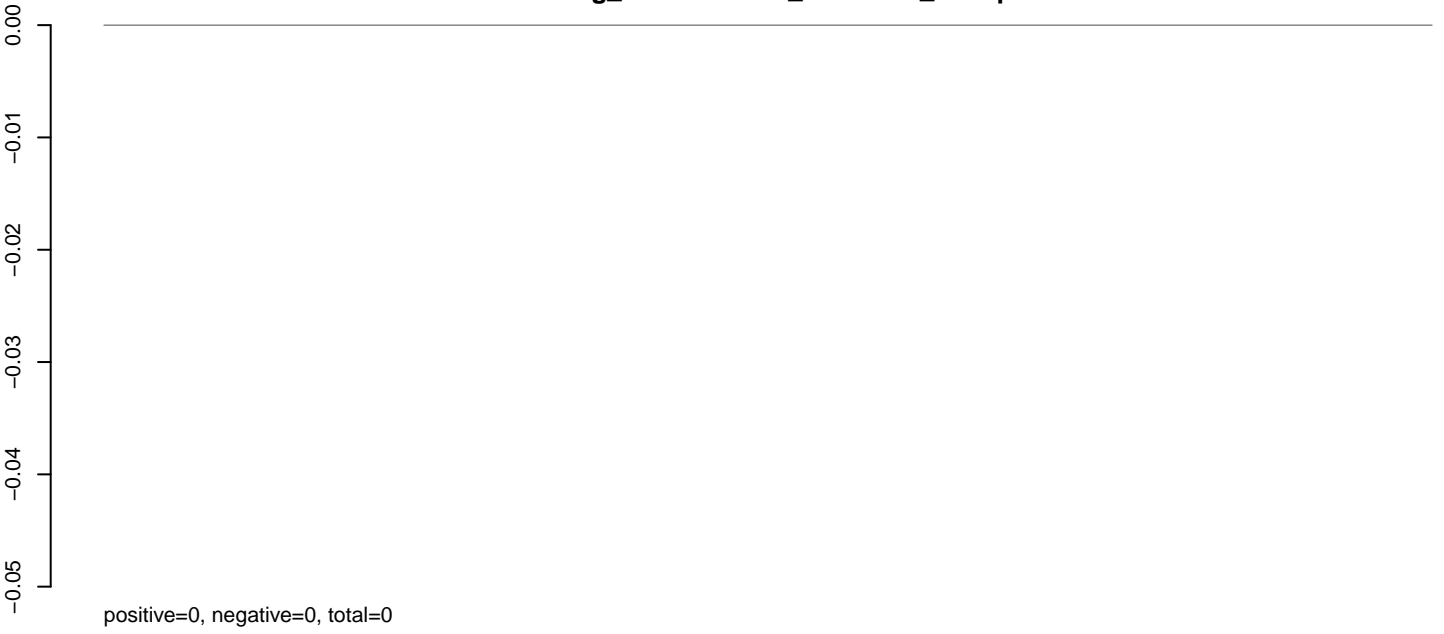
AeAeg_CCL.125cells_CHIKV.rep



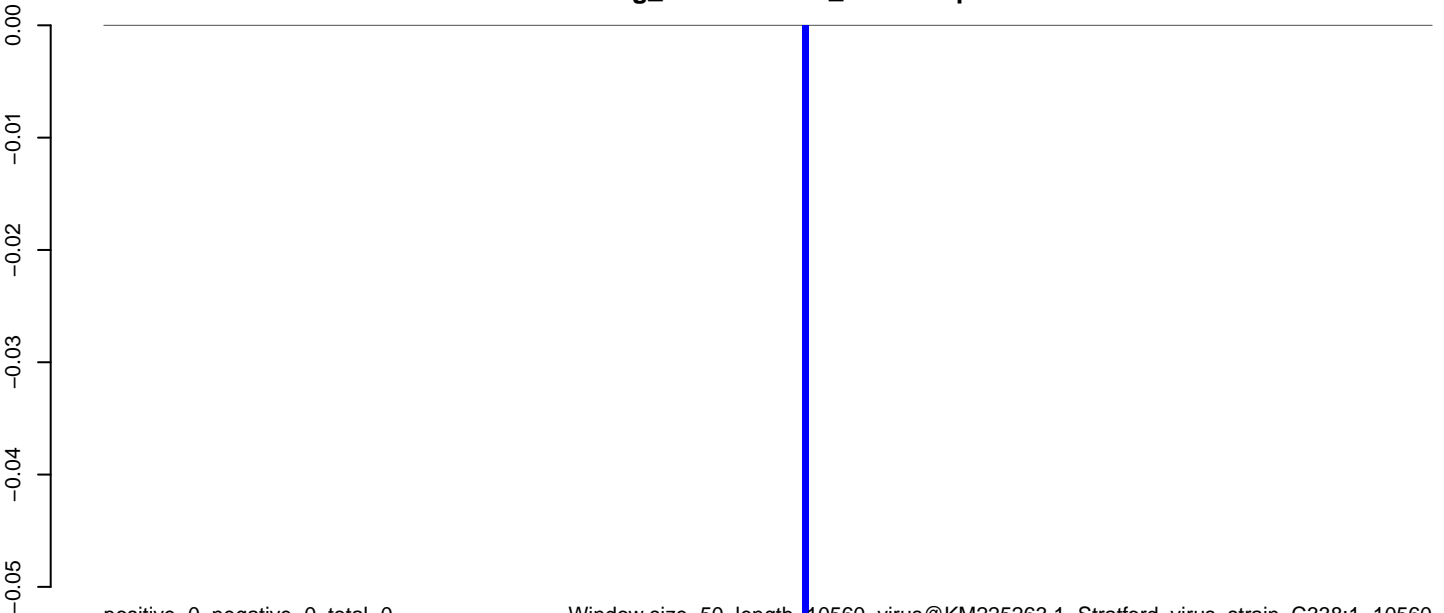
AeAeg_CCL.125cells_CHIKV.18_23.rep



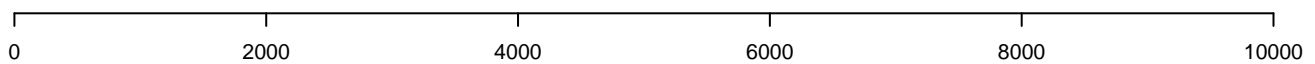
AeAeg_CCL.125cells_CHIKV.24_35.rep



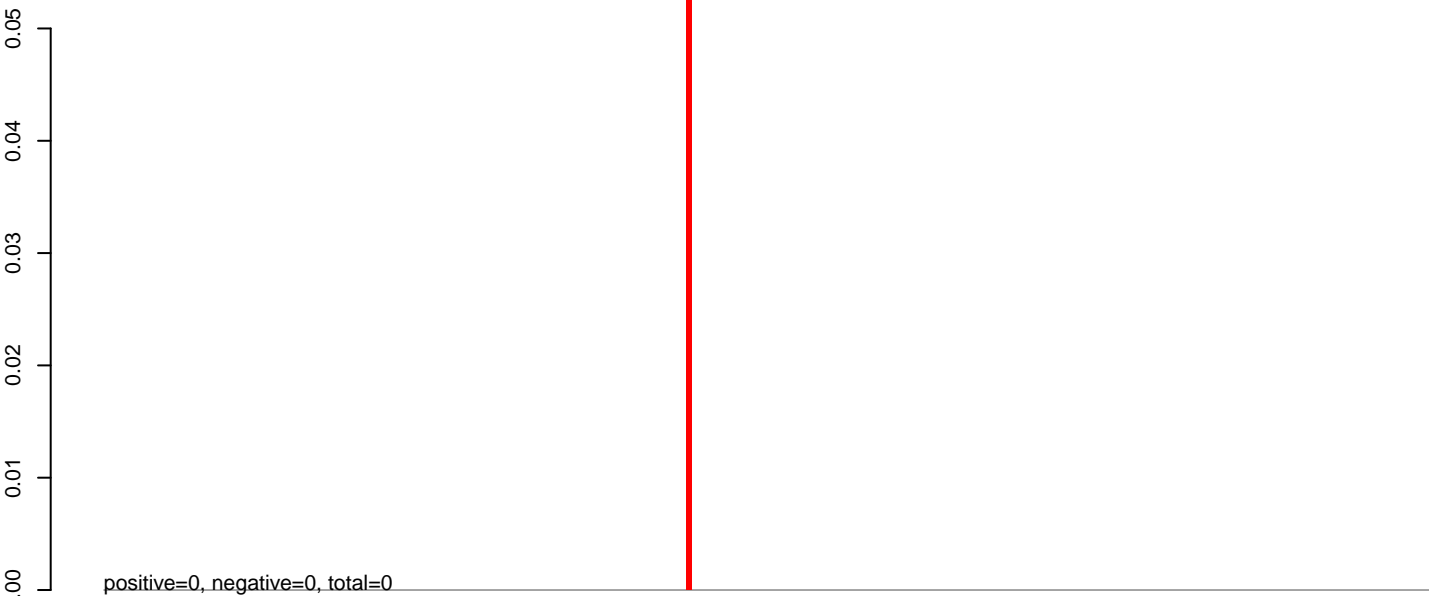
AeAeg_CCL.125cells_CHIKV.rep



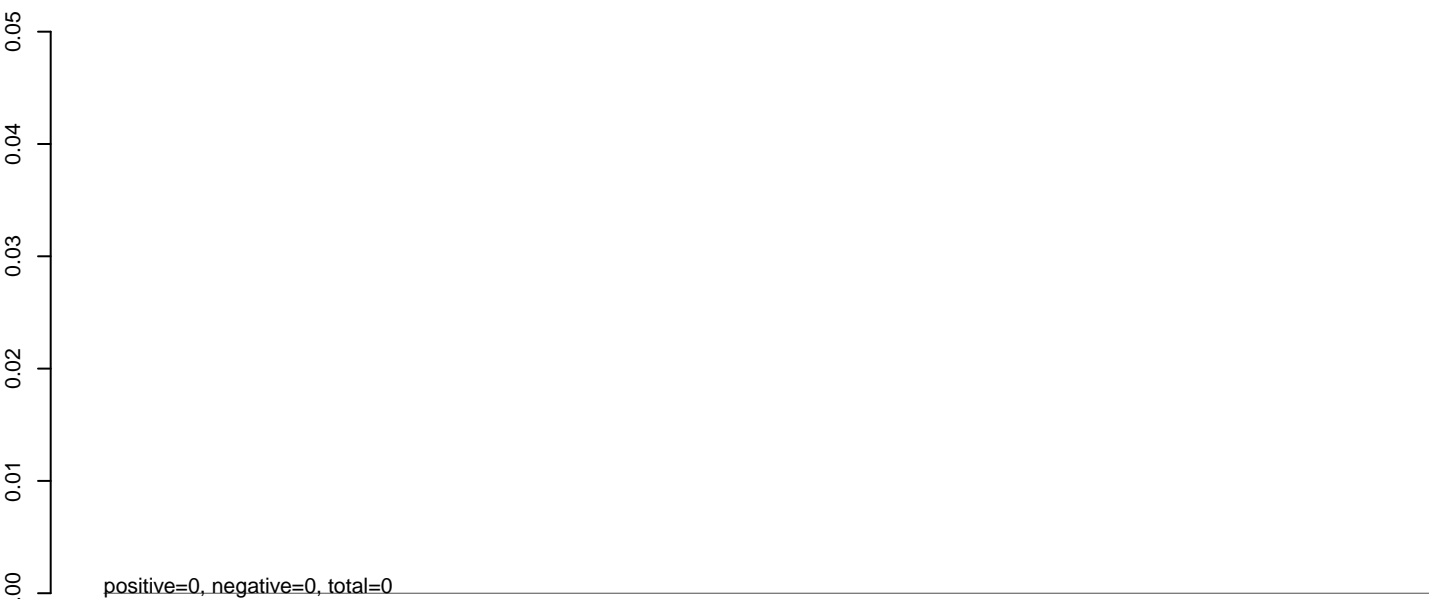
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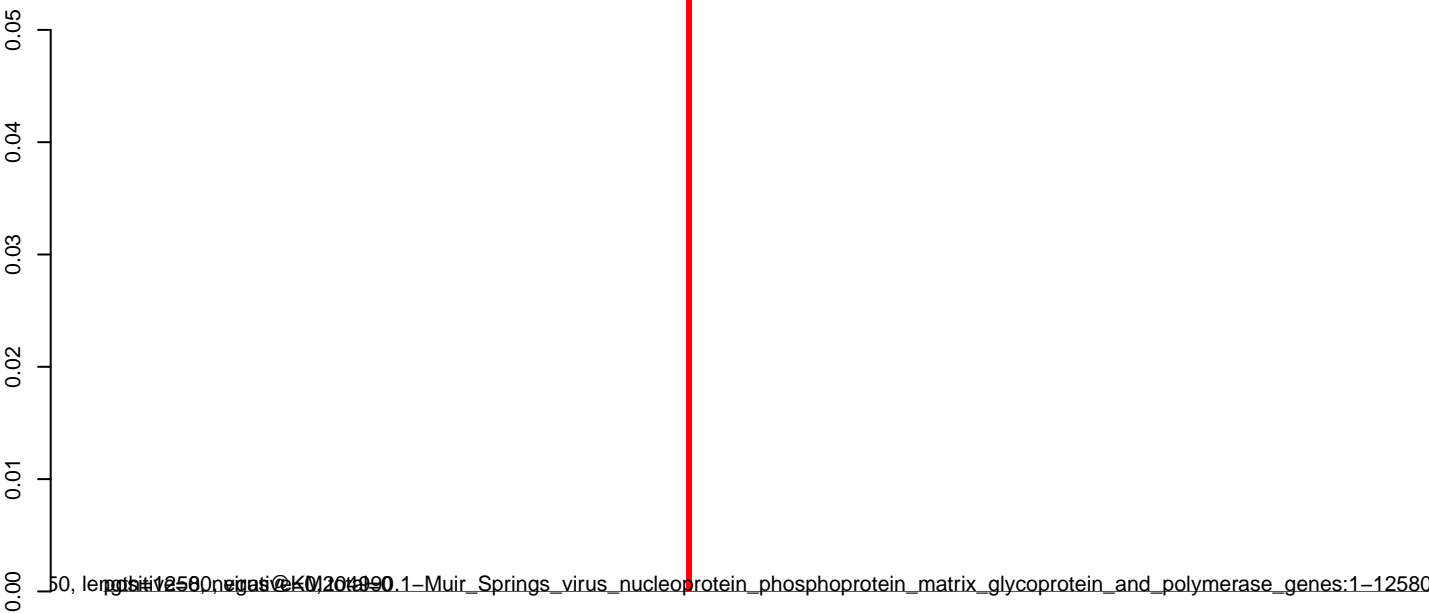
AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep



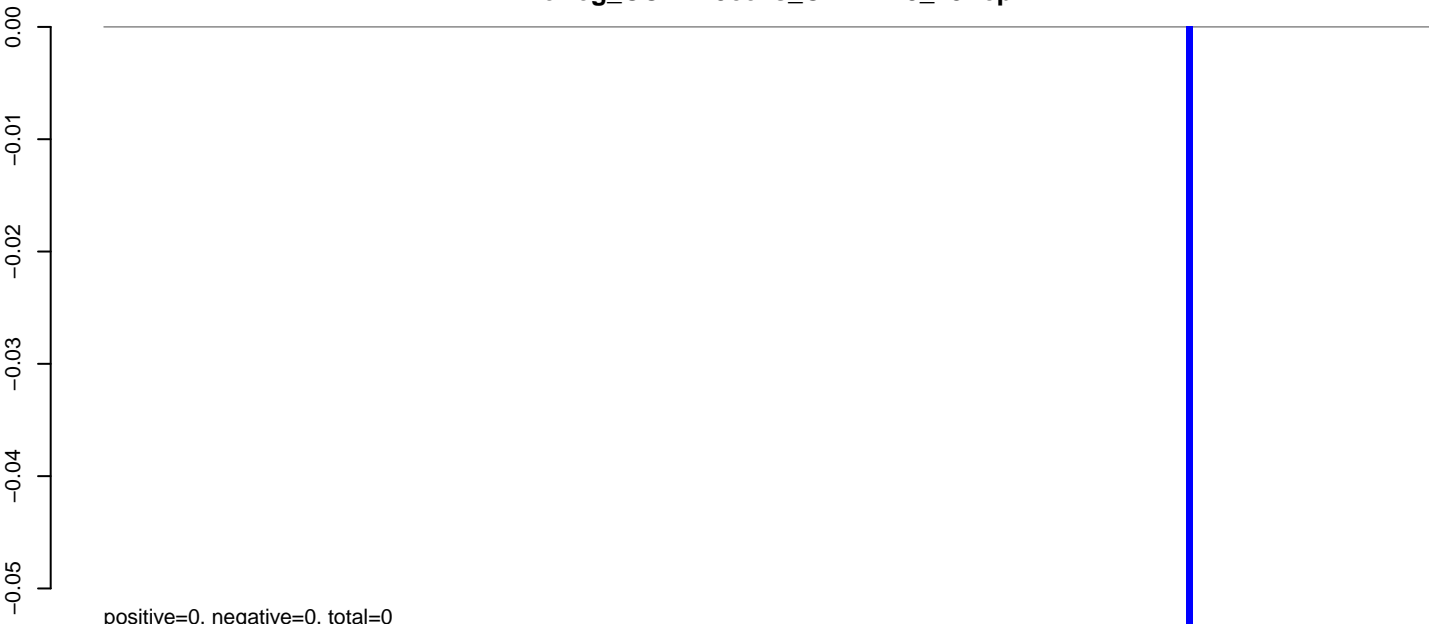
AeAeg_CCL.125cells_CHIKV.rep



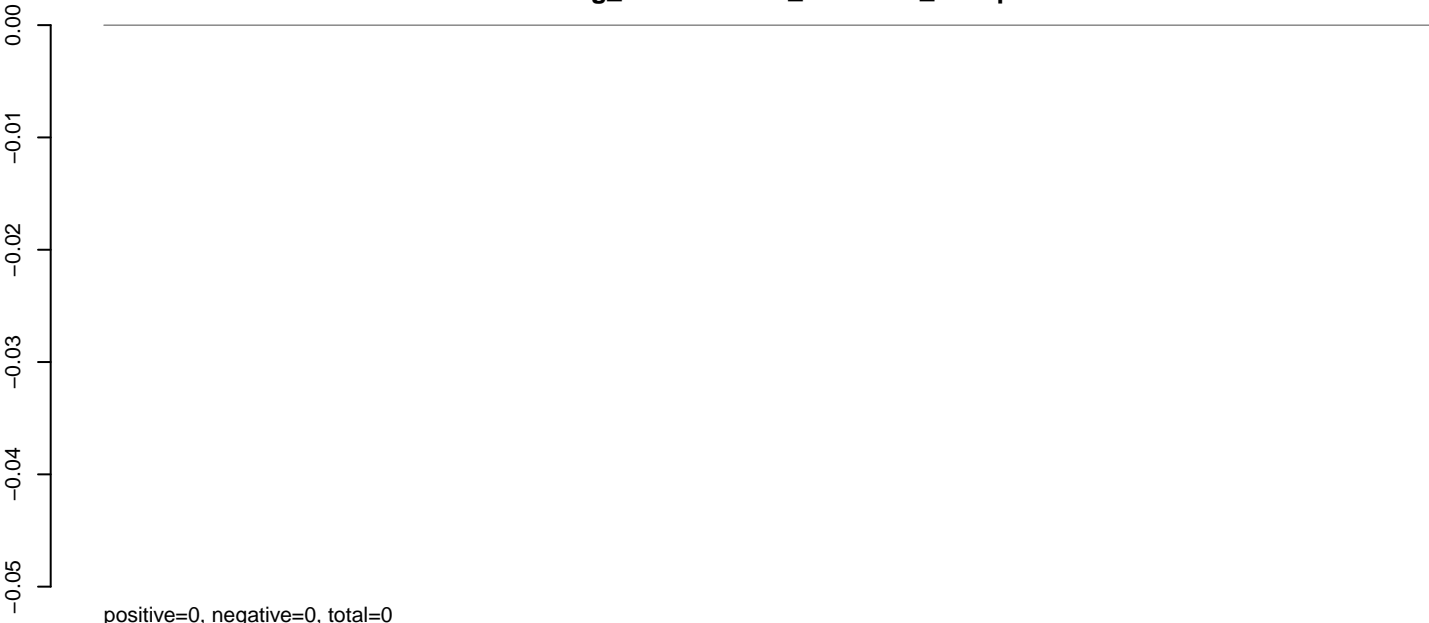
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0 2000 4000 6000 8000 10000 12000

AeAeg_CCL.125cells_CHIKV.18_23.rep



AeAeg_CCL.125cells_CHIKV.24_35.rep

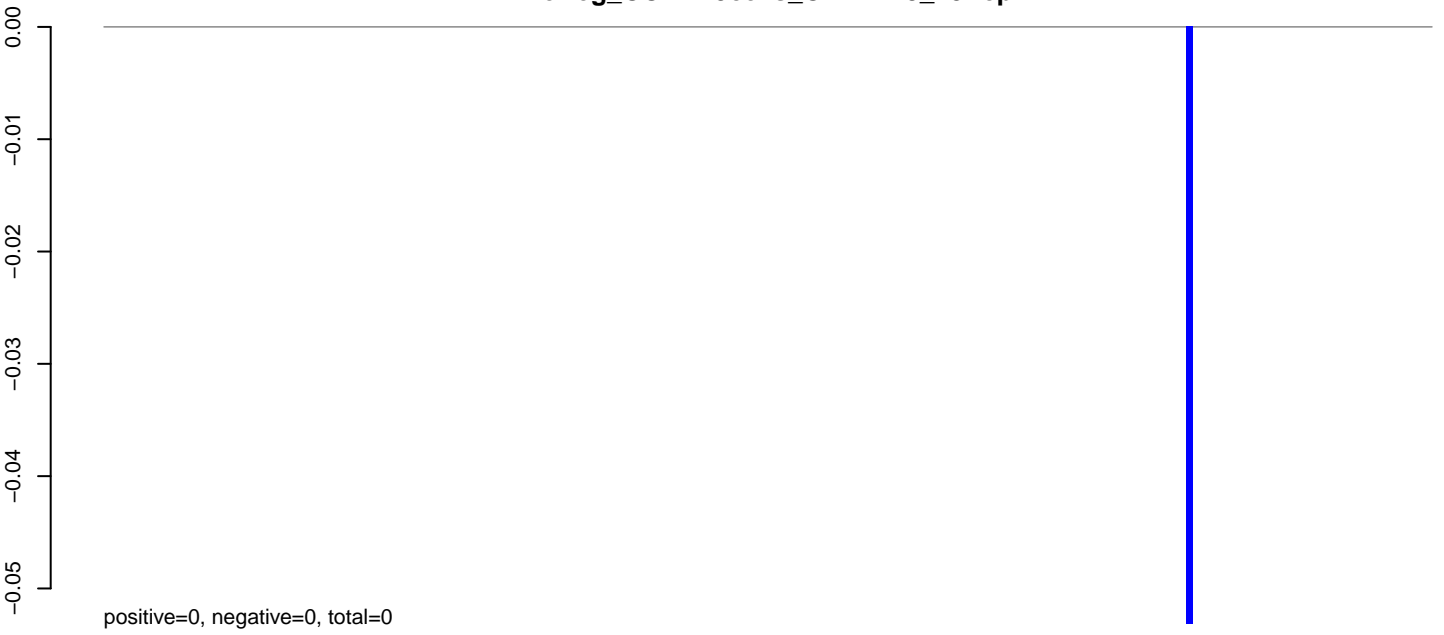


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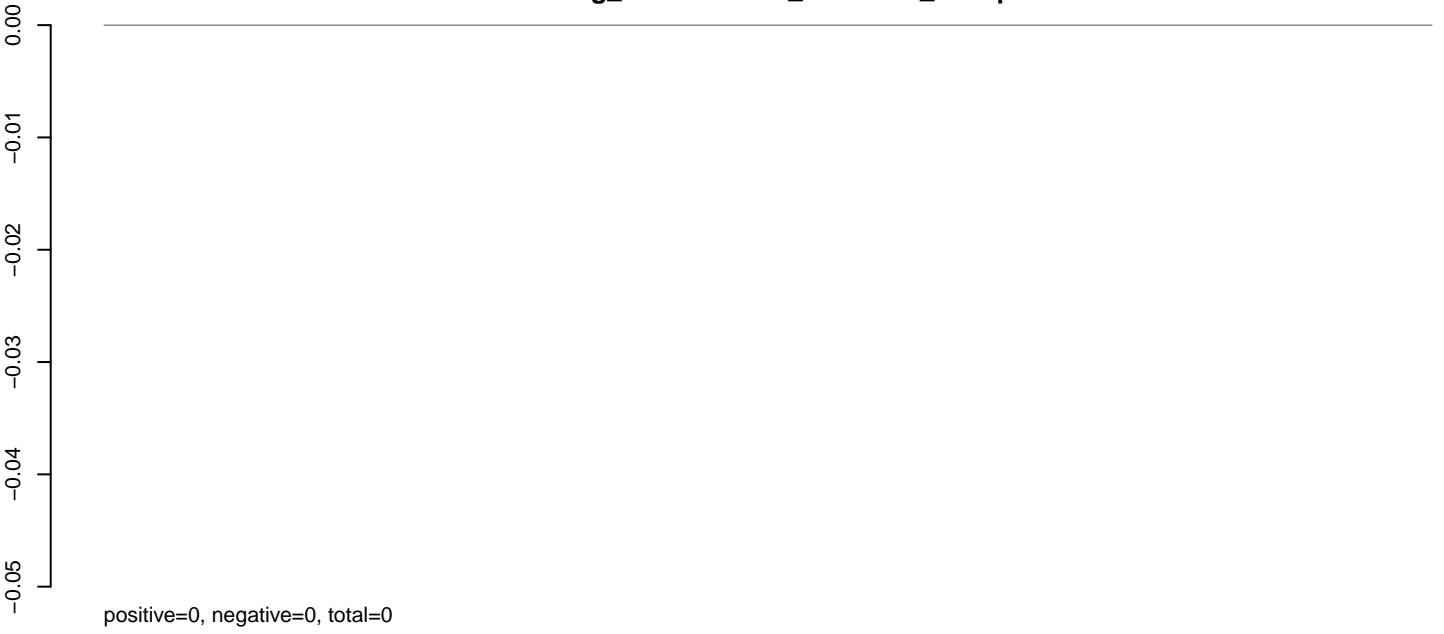


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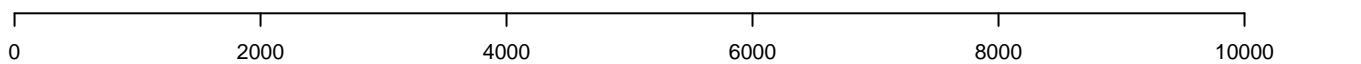
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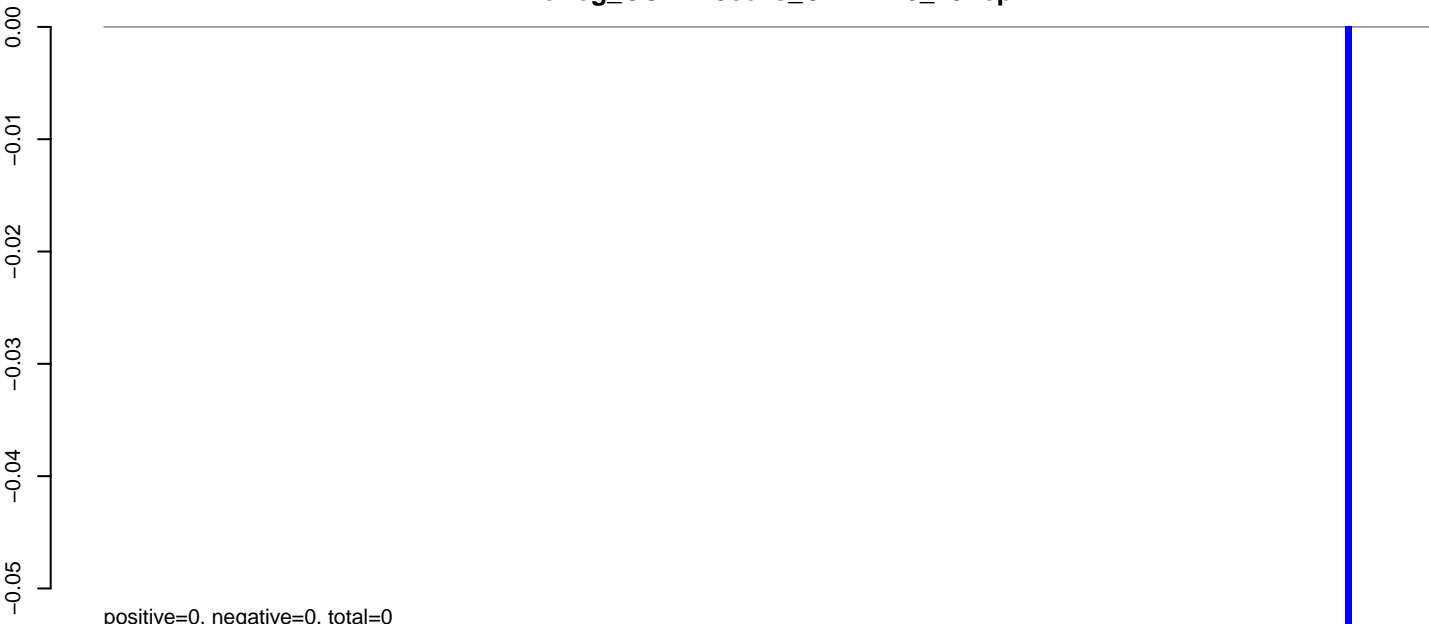
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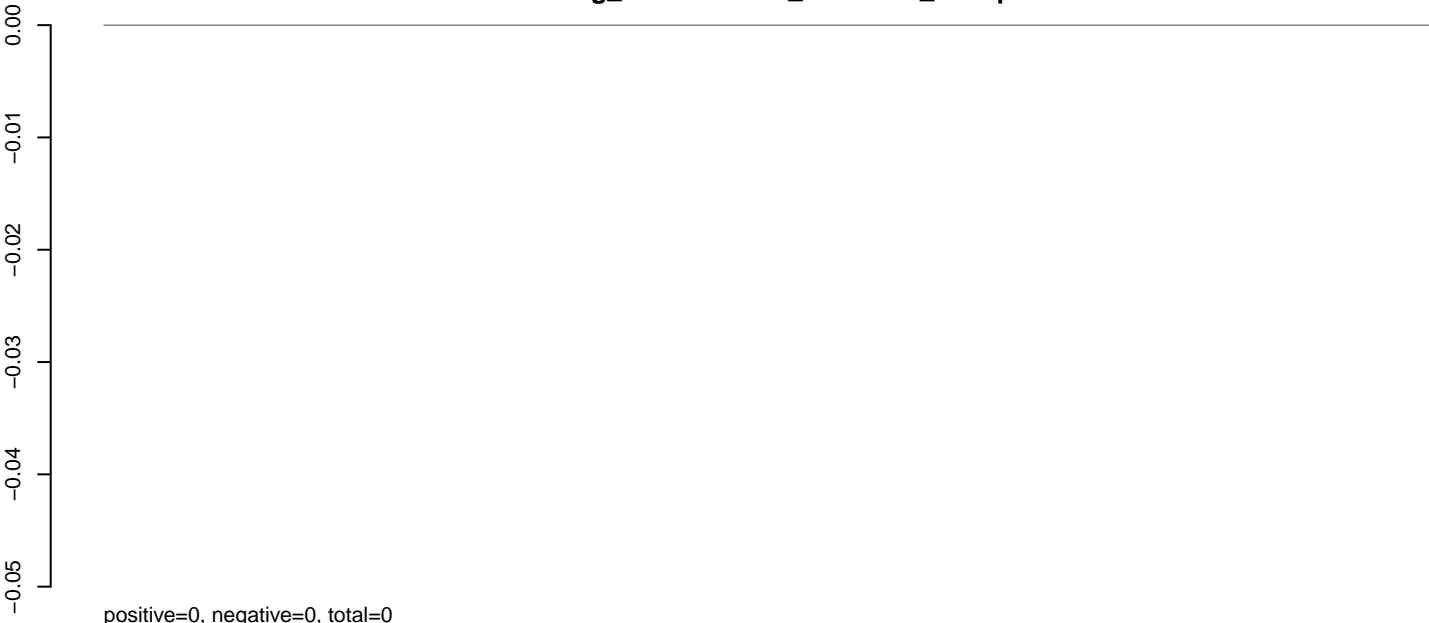
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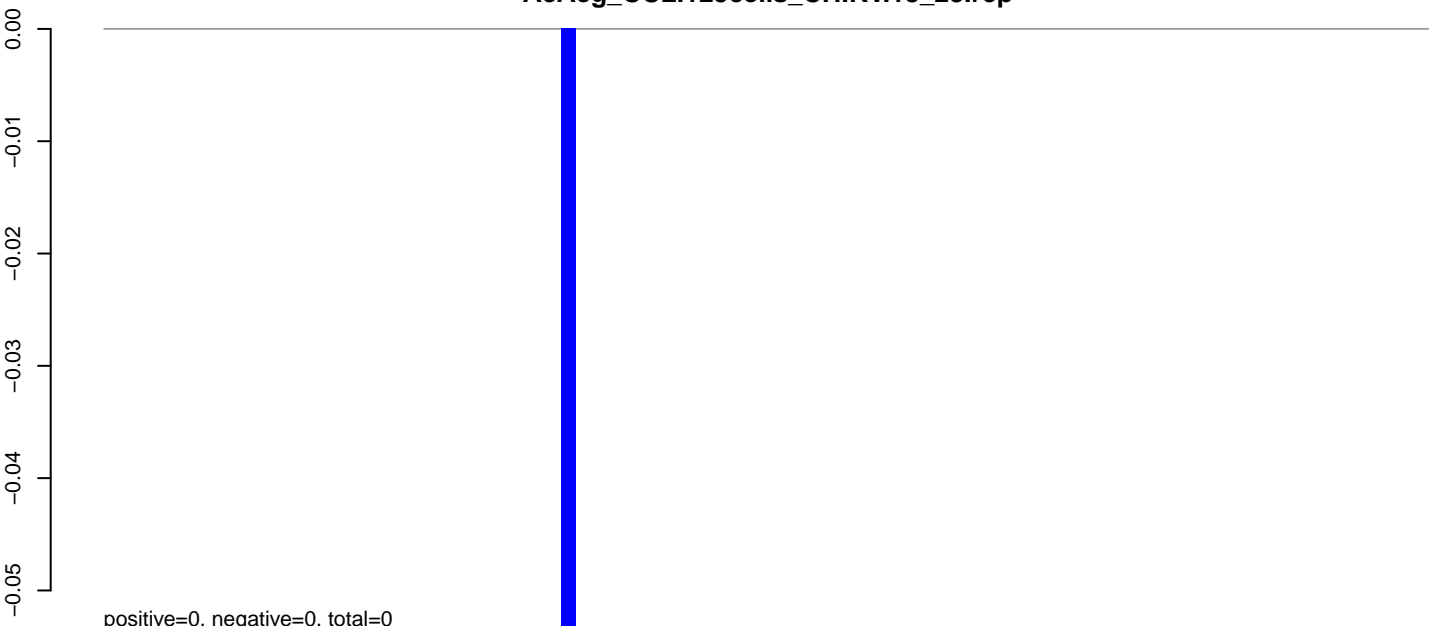
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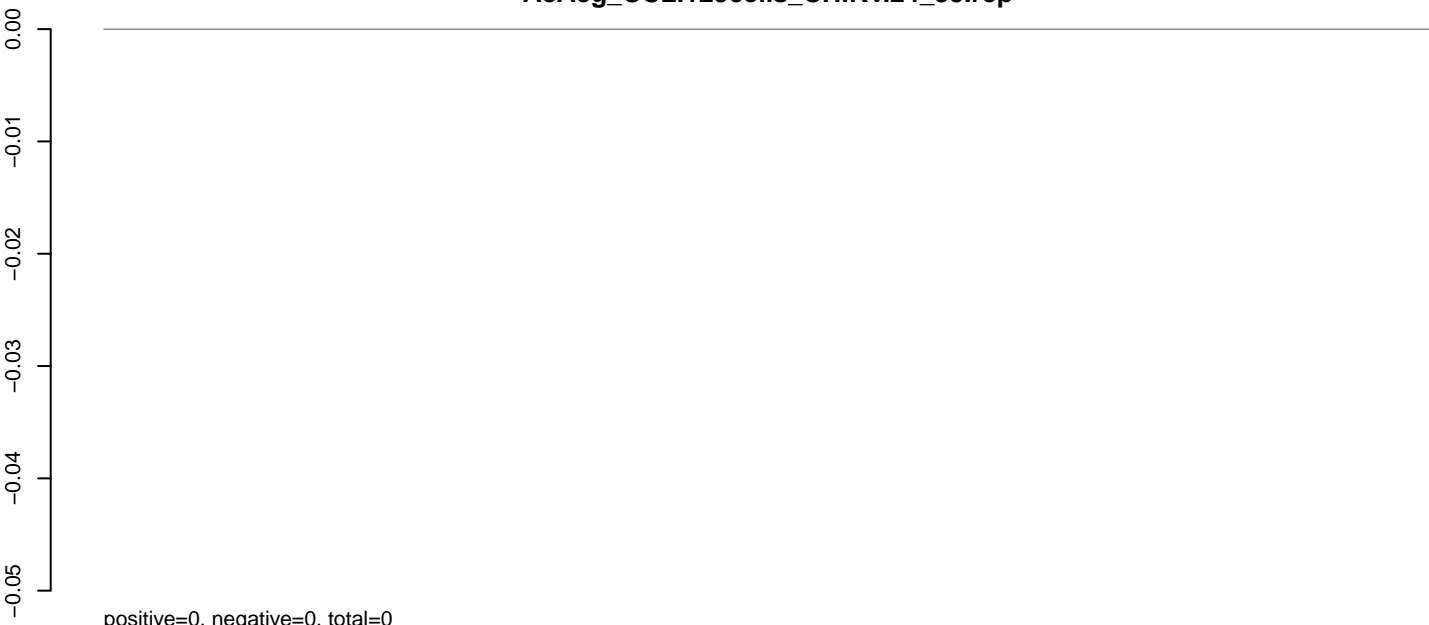
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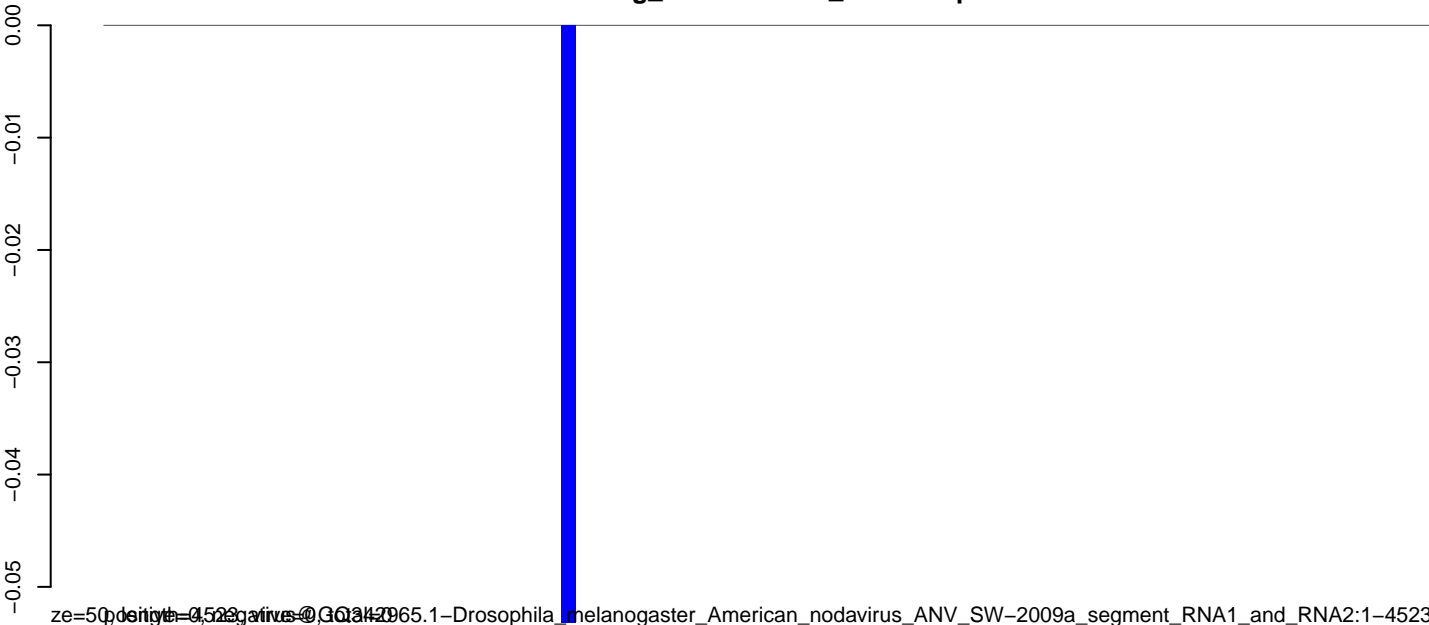
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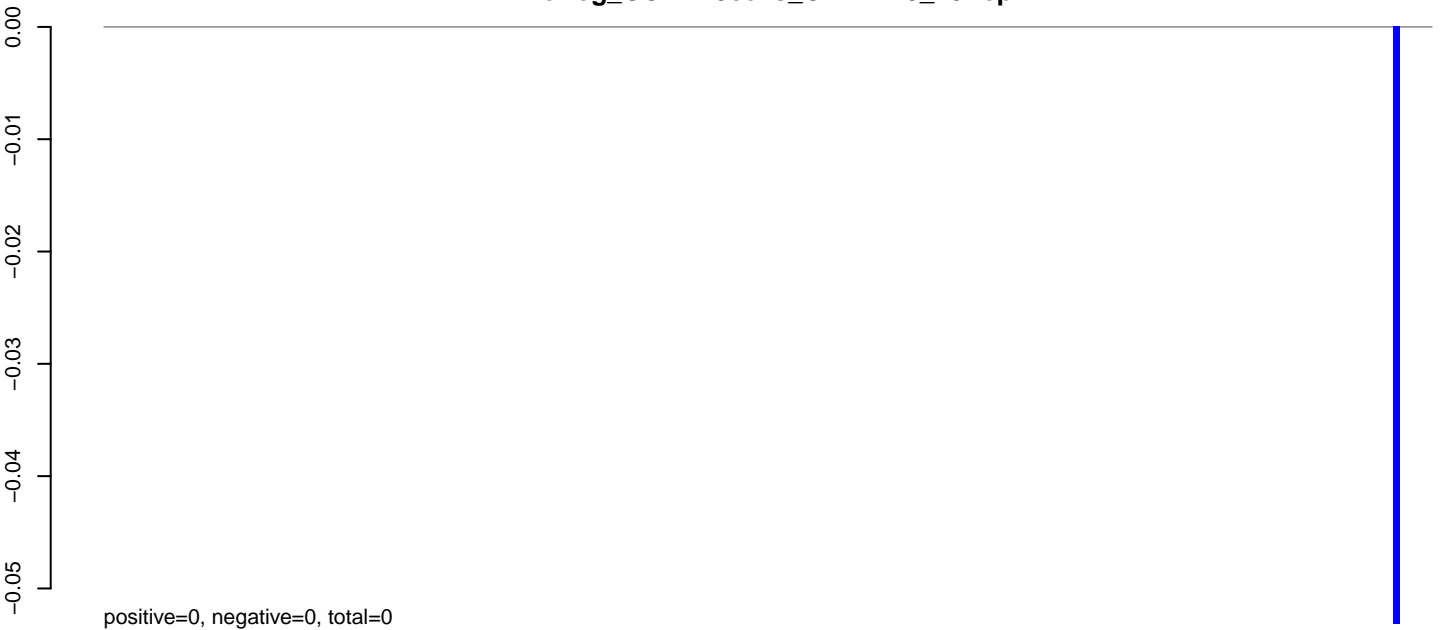
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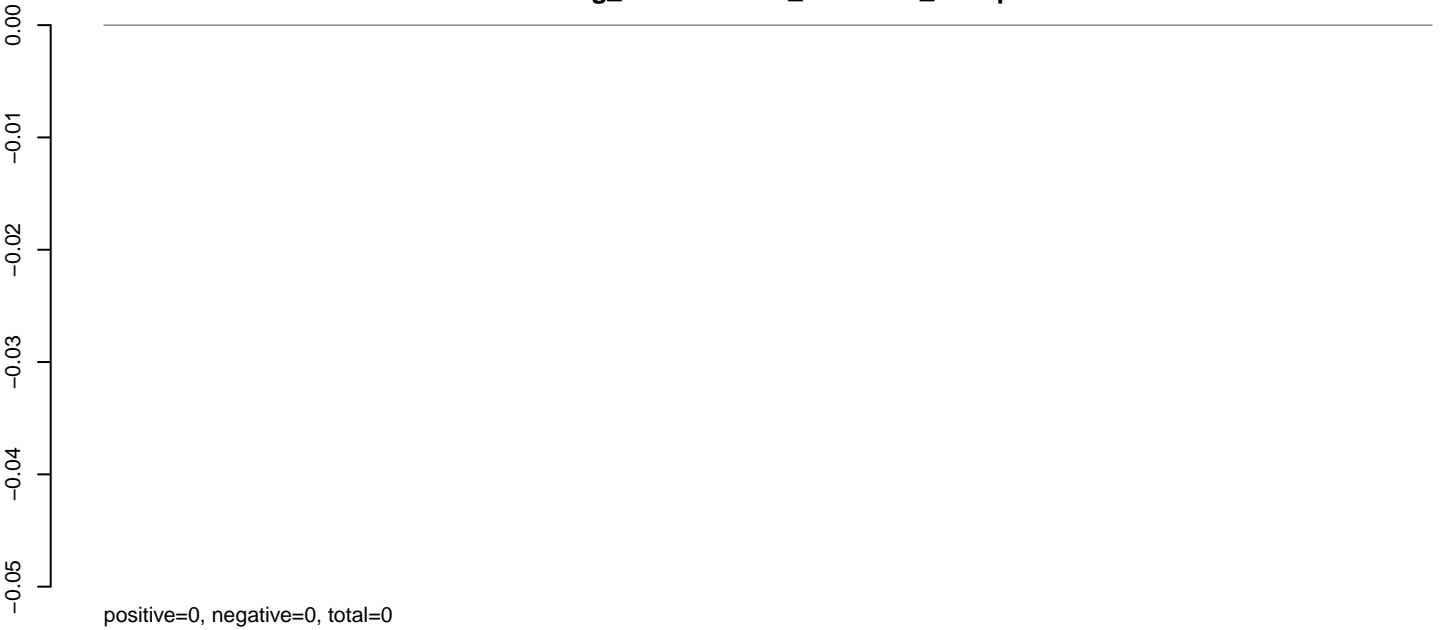
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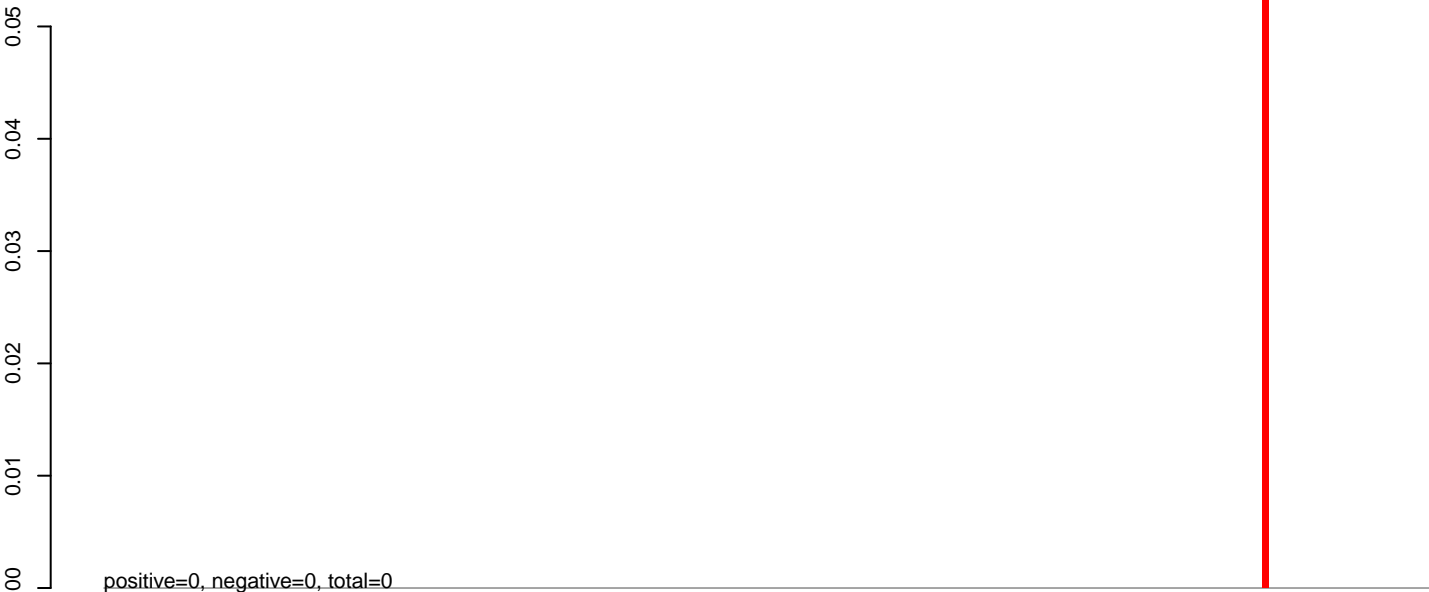
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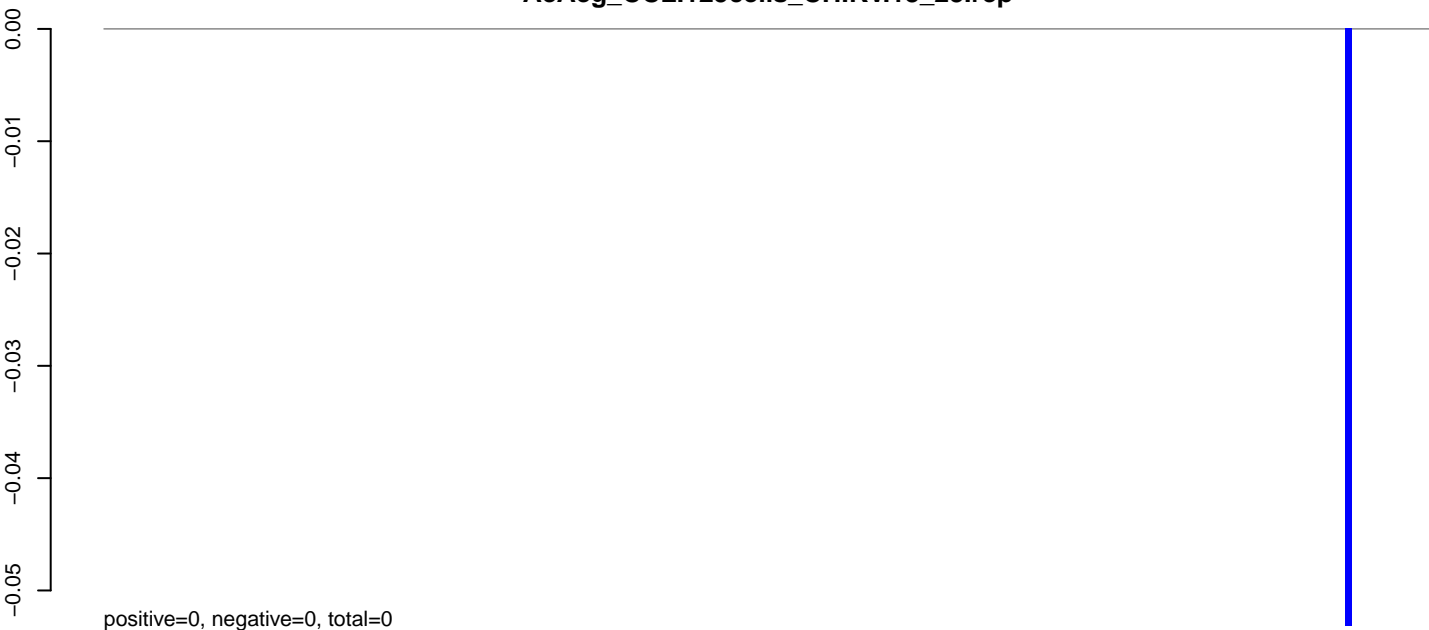


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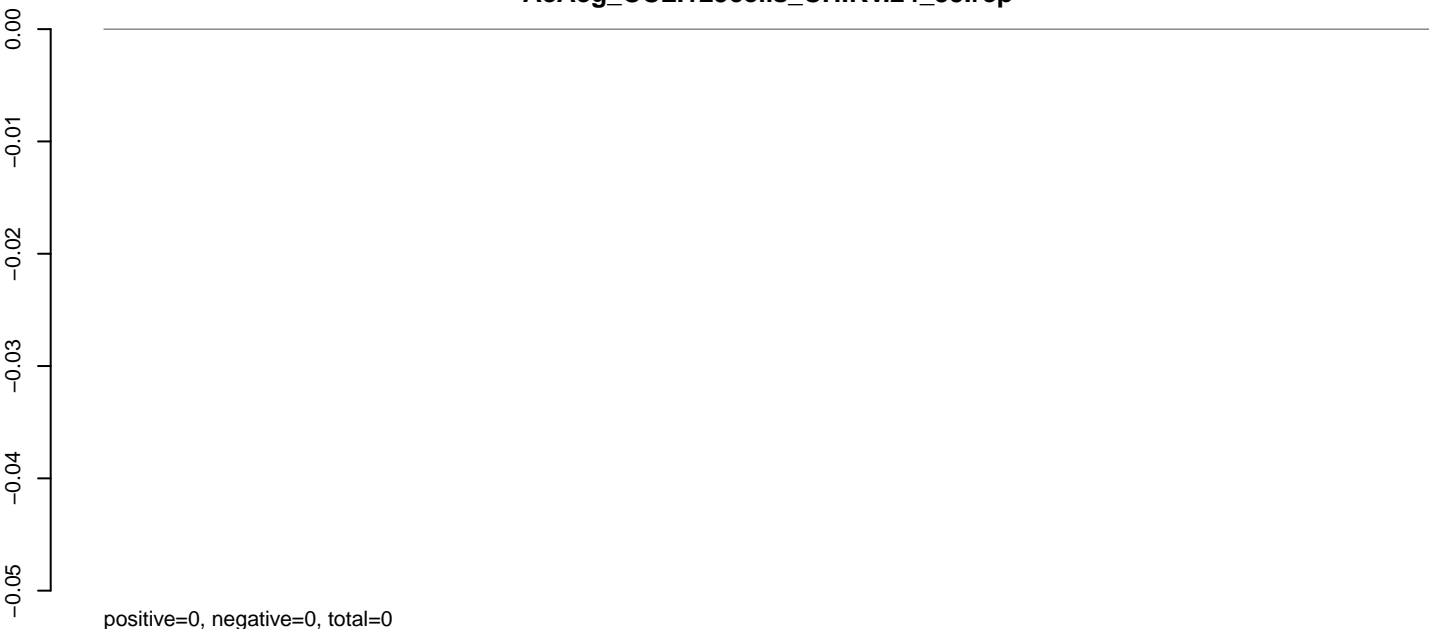


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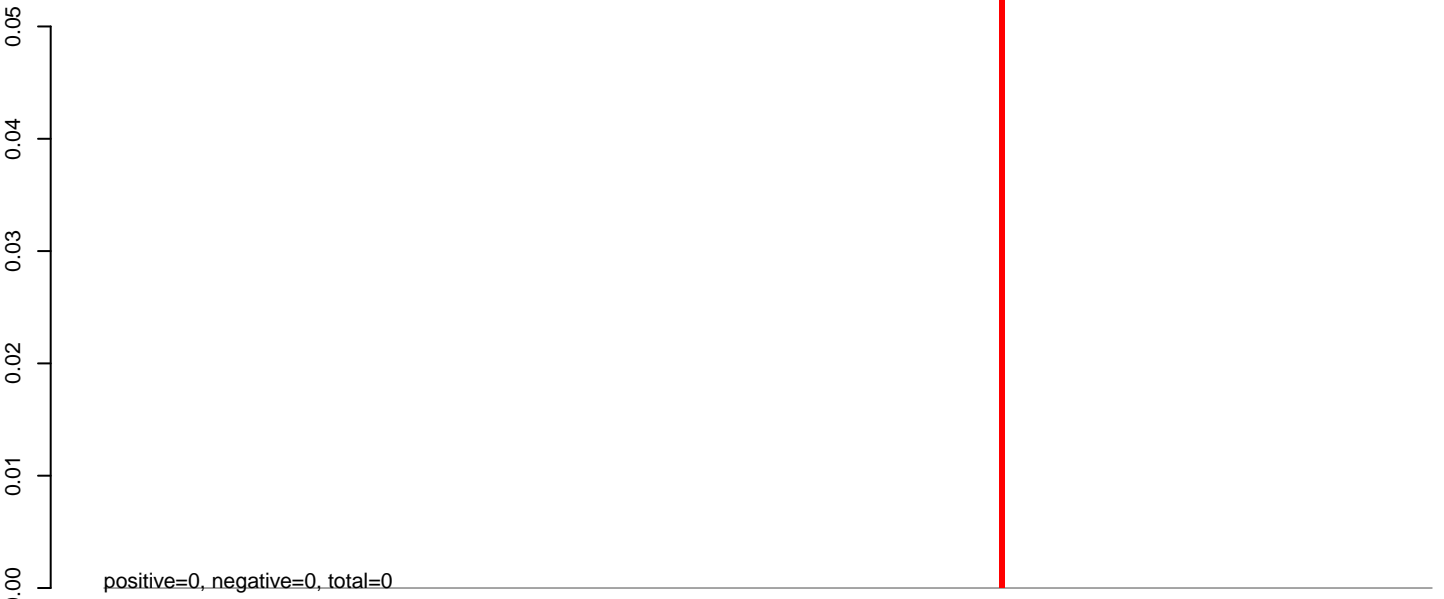
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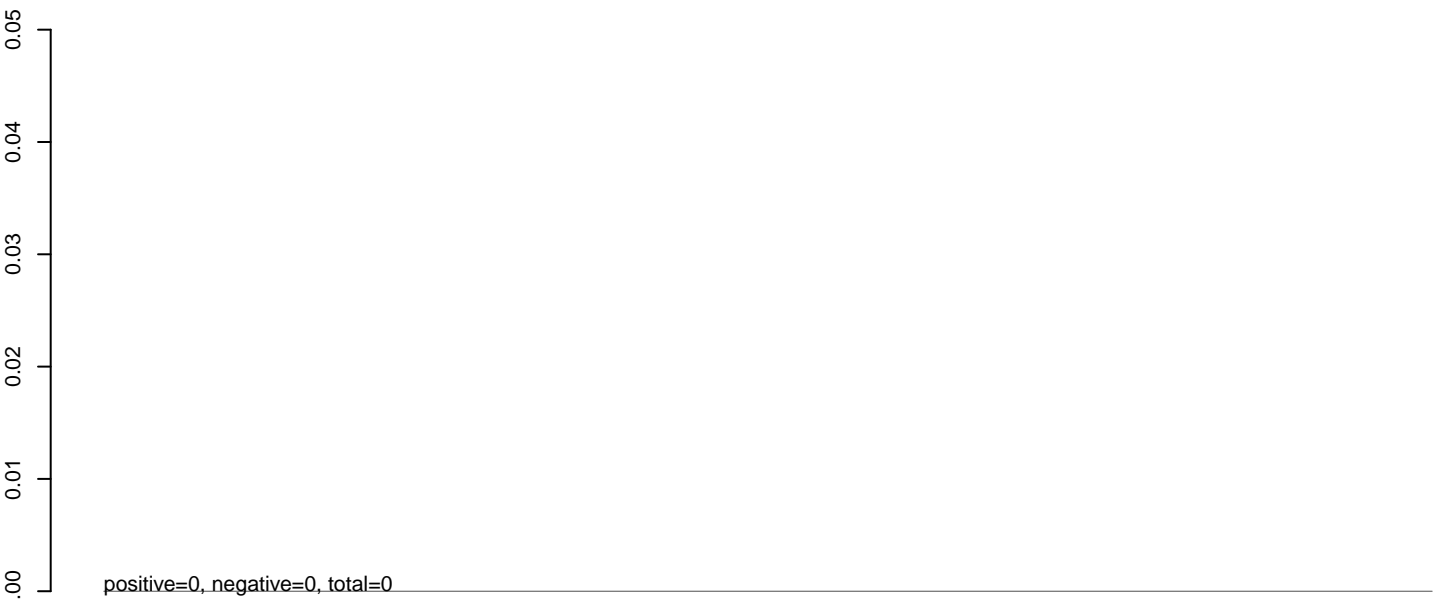
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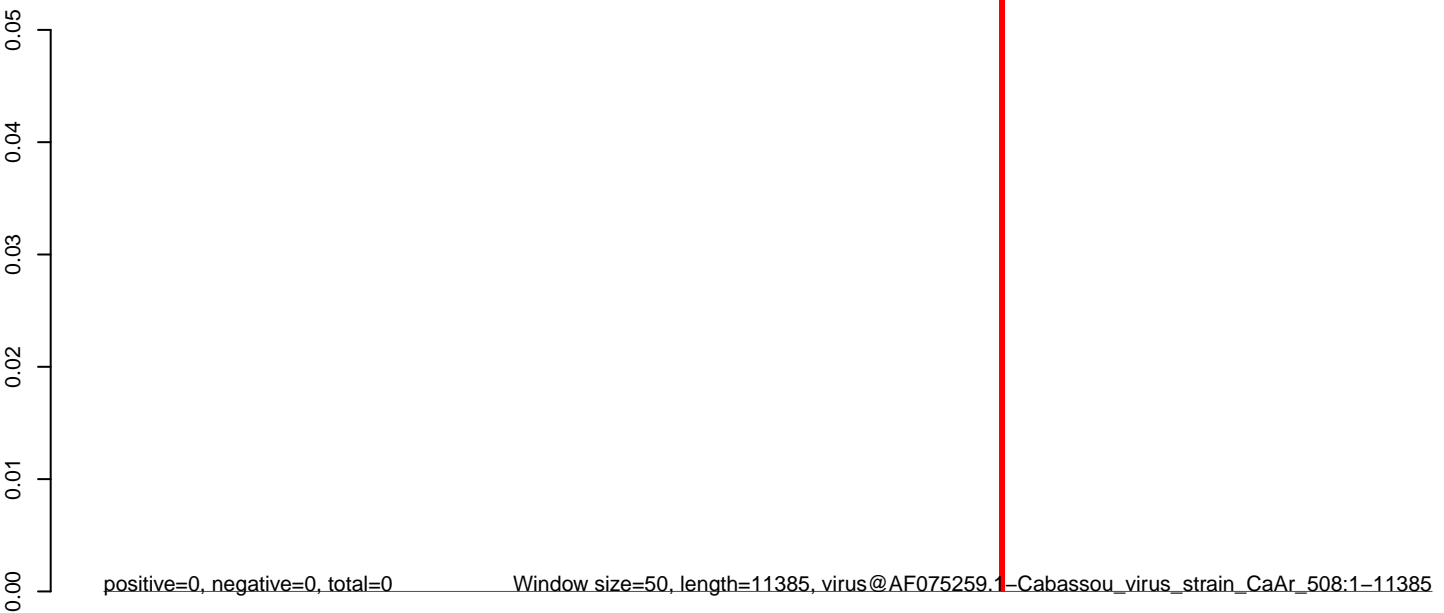
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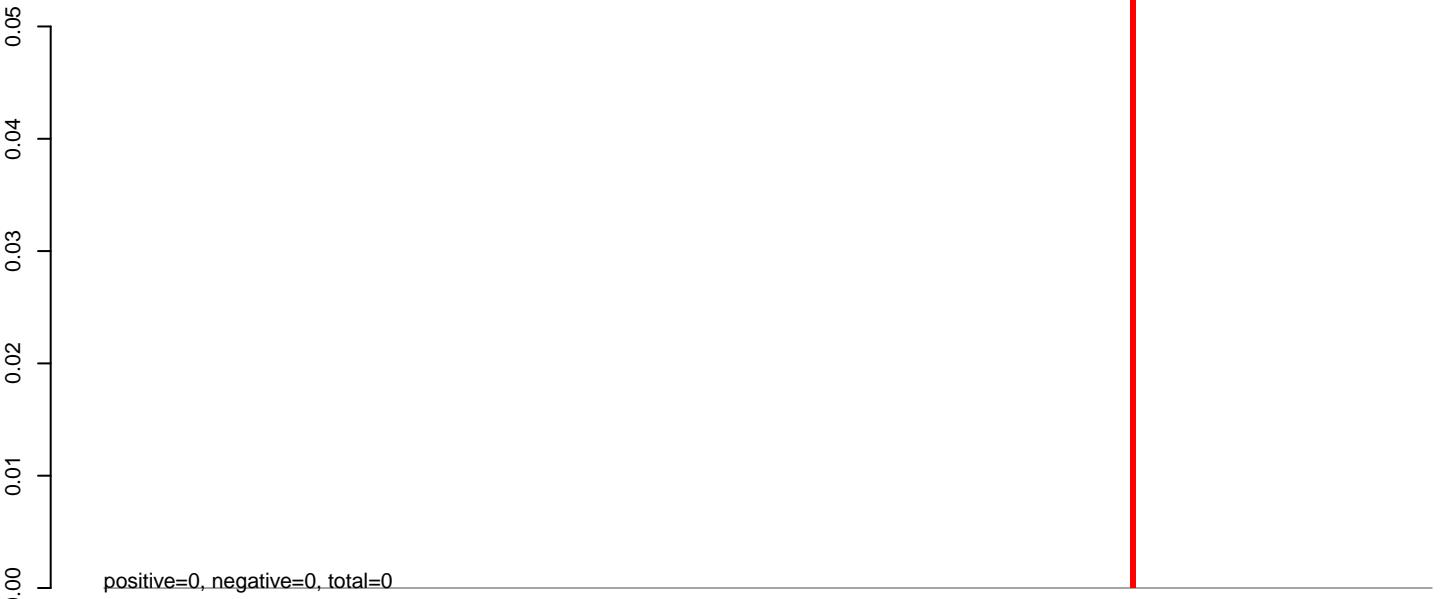


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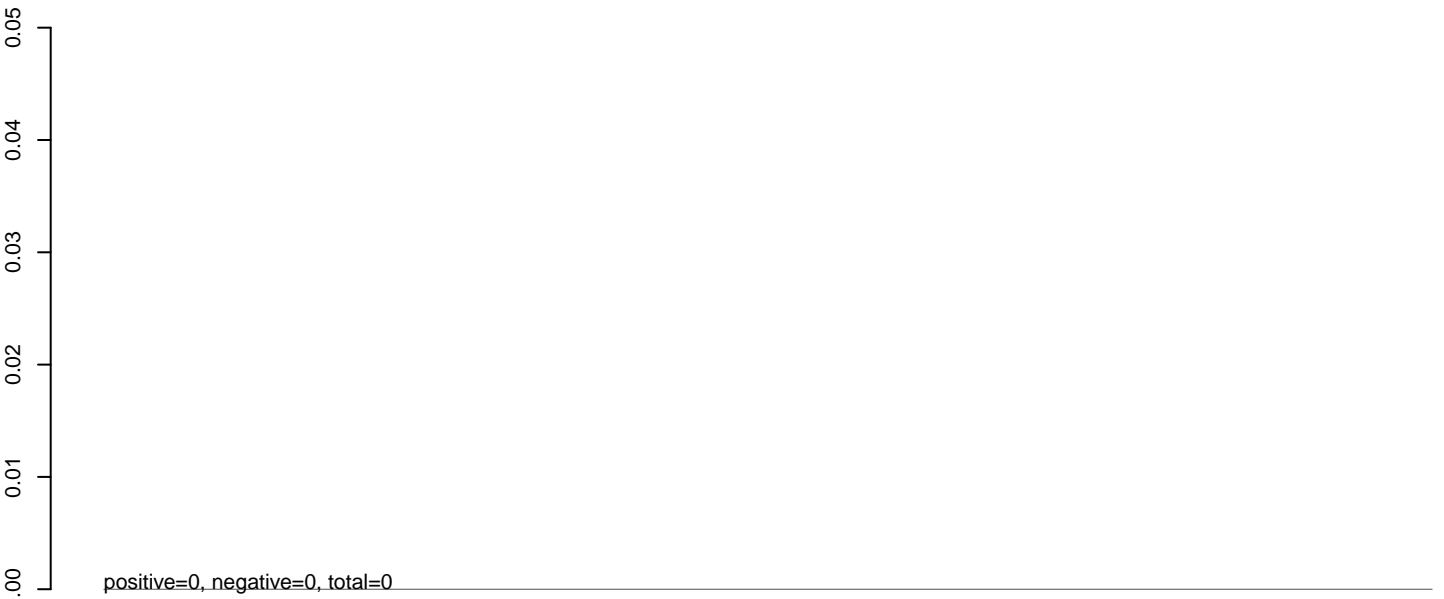


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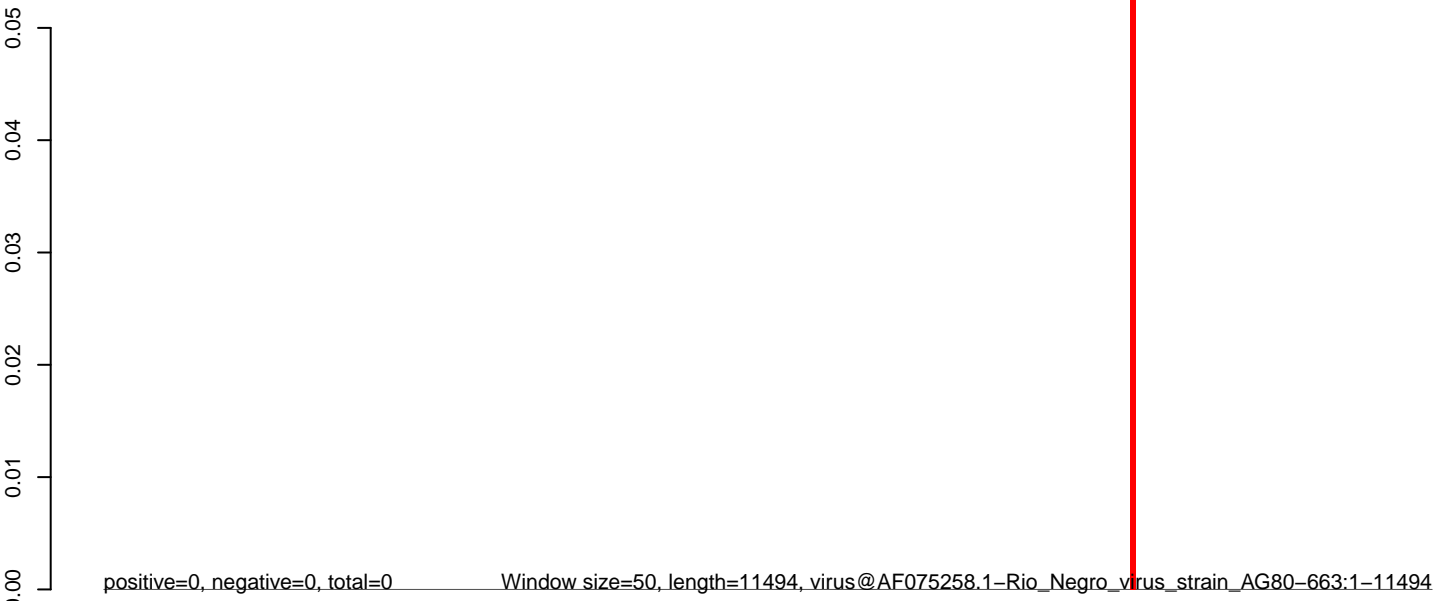
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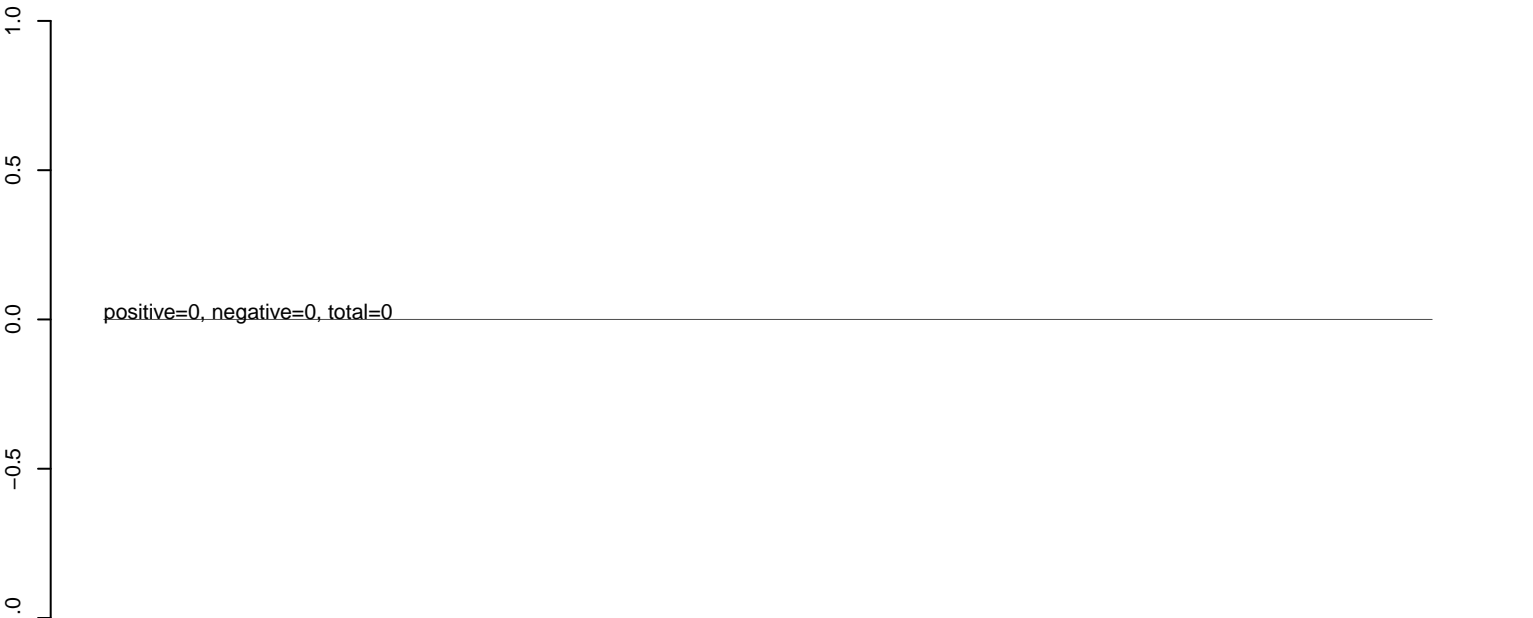
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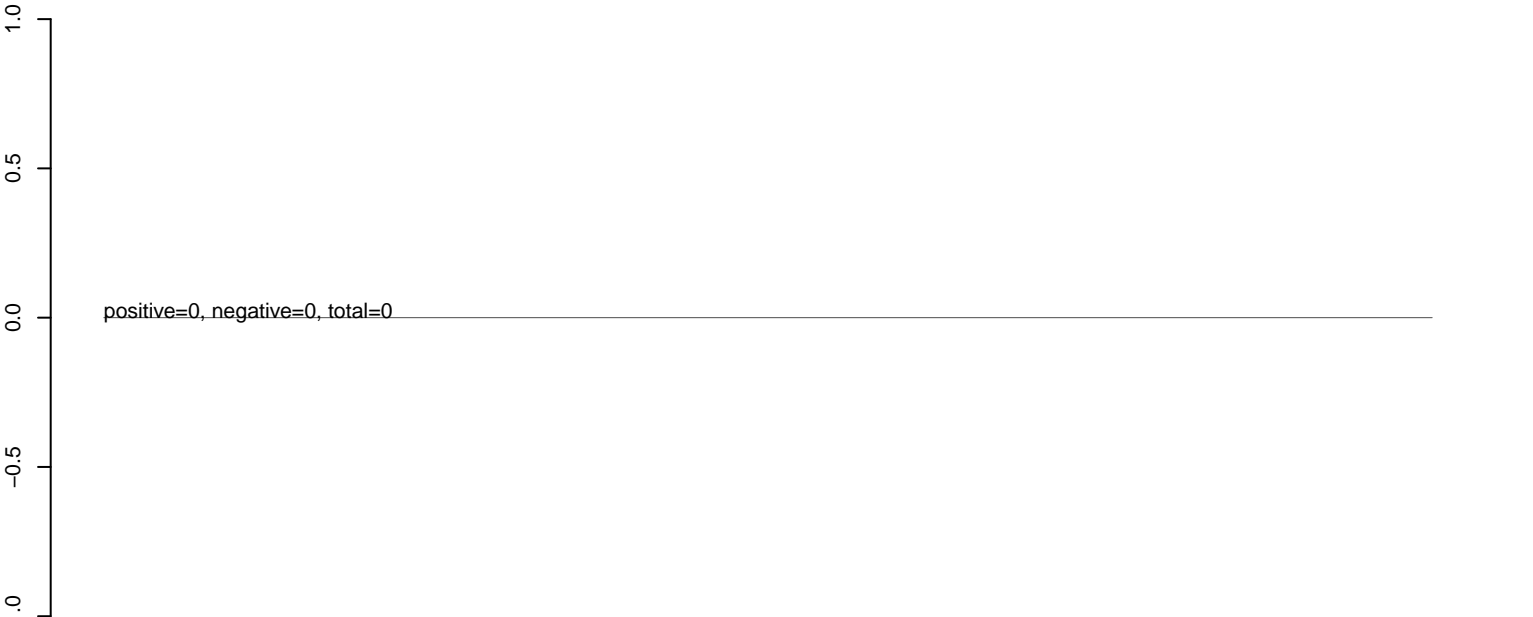
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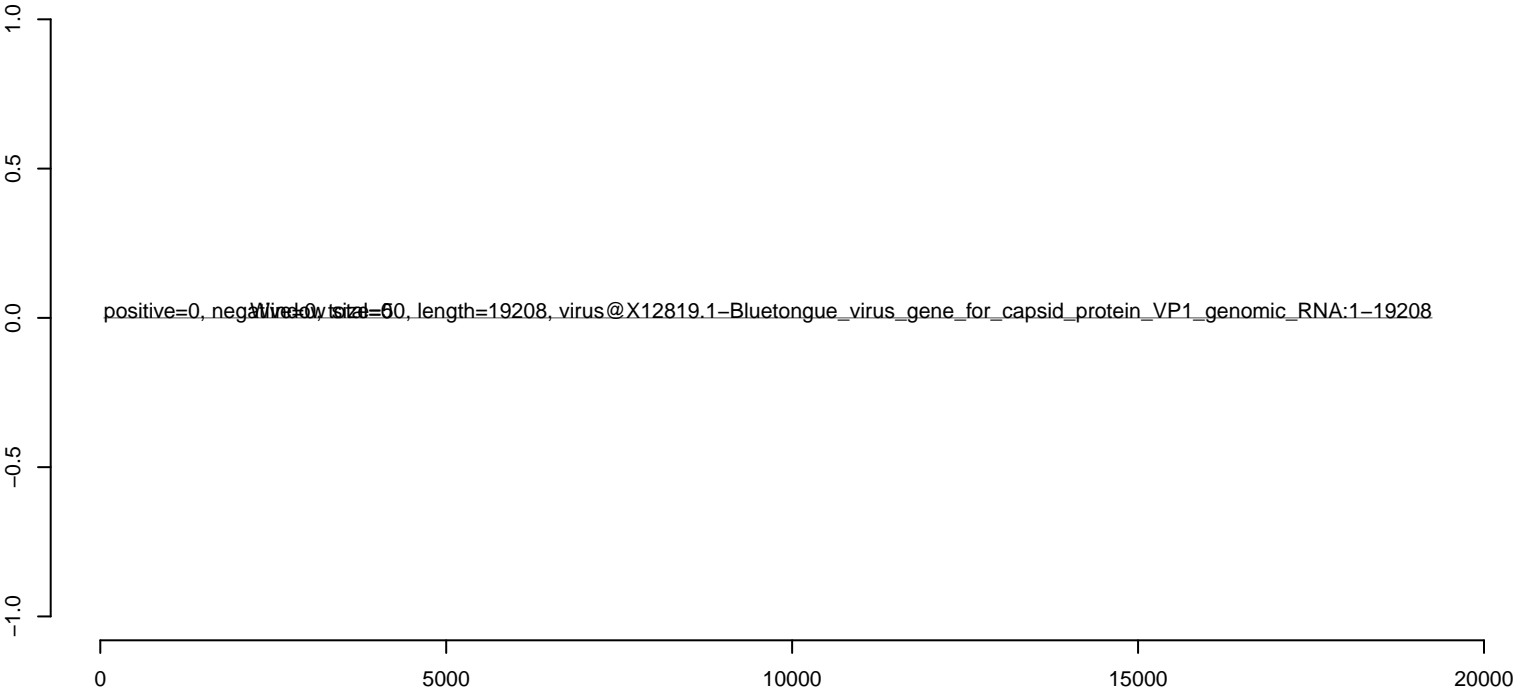
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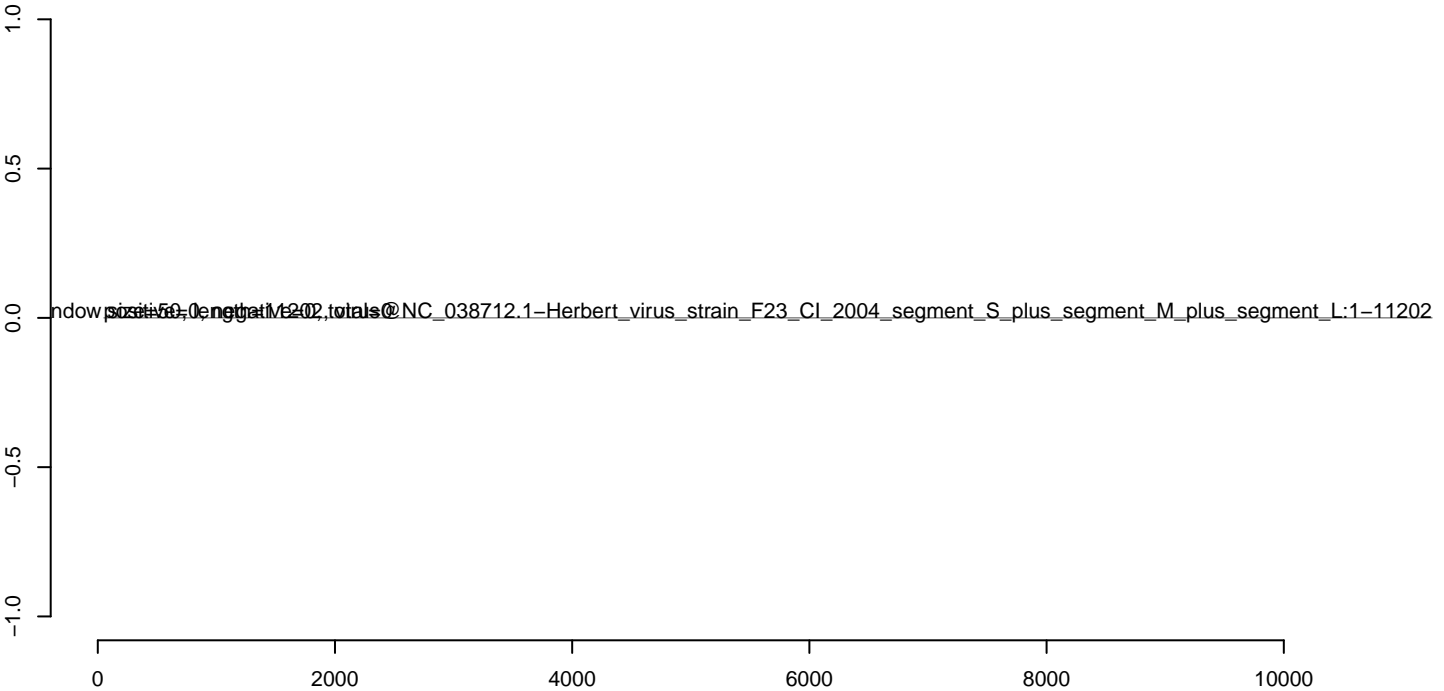
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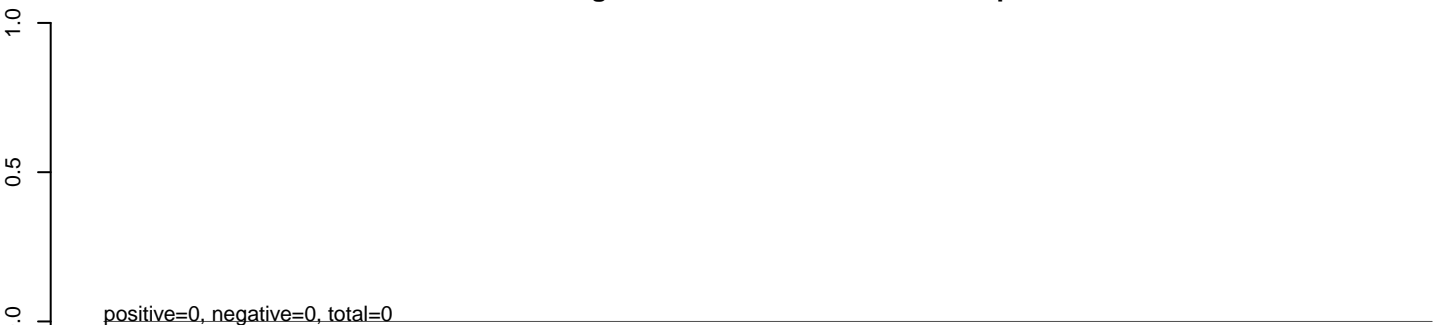
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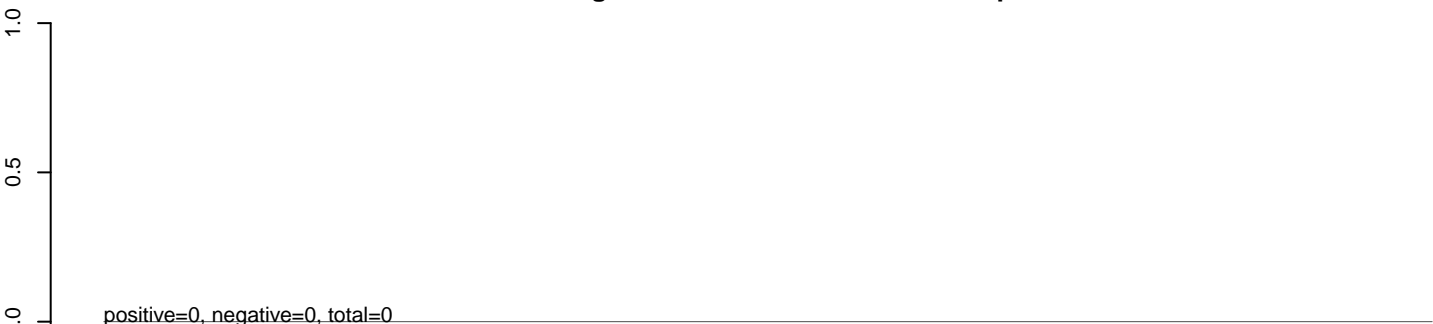
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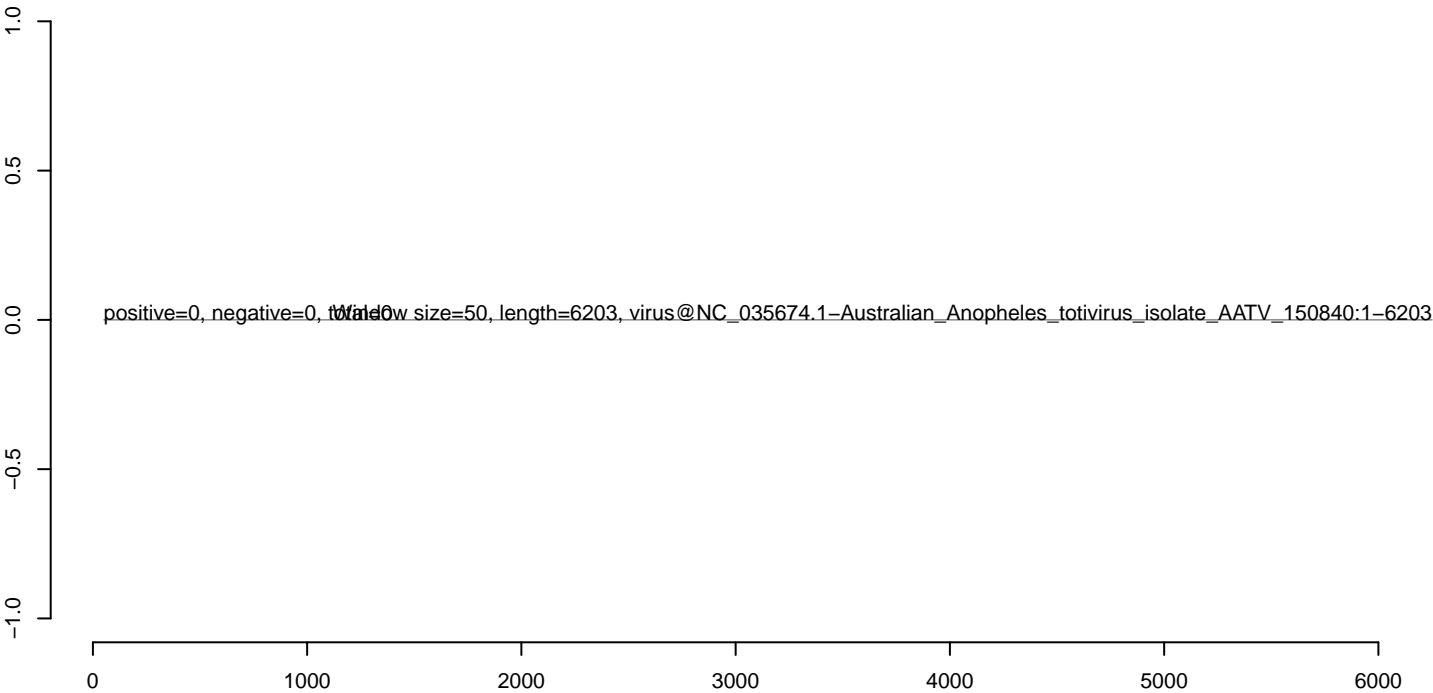
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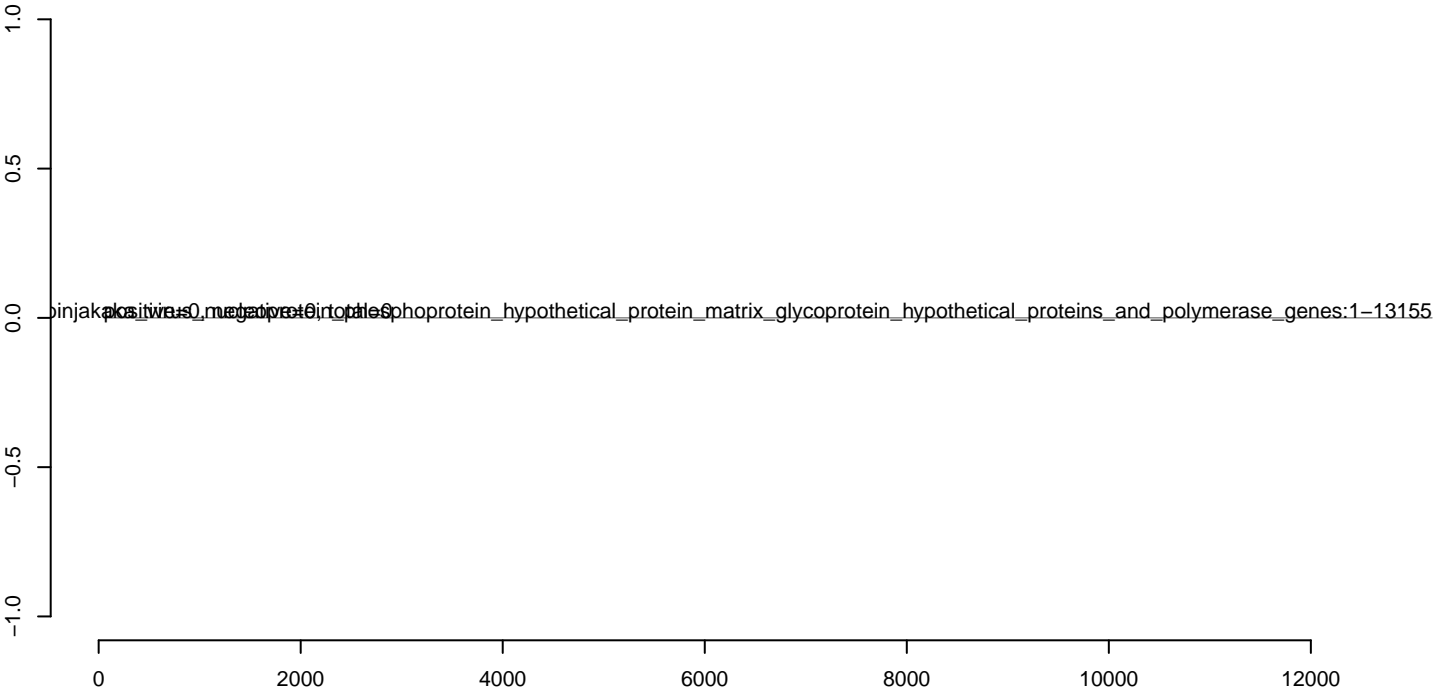
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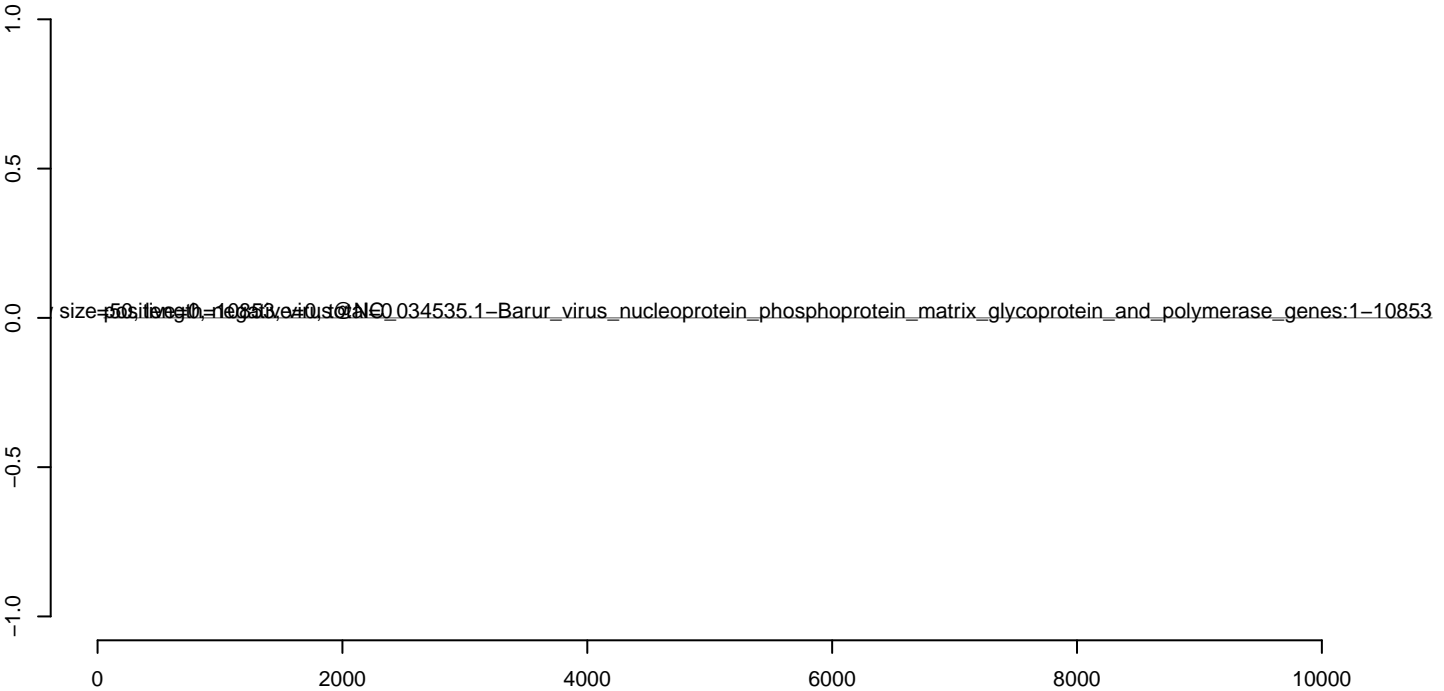
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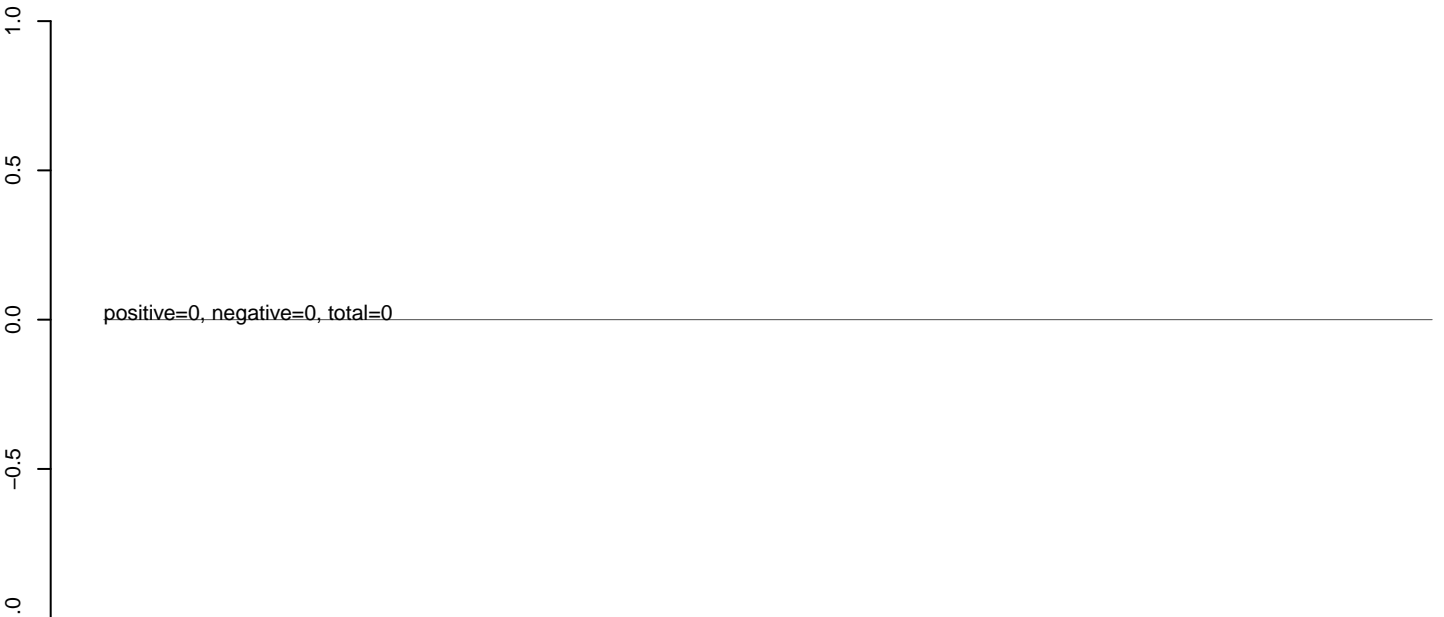
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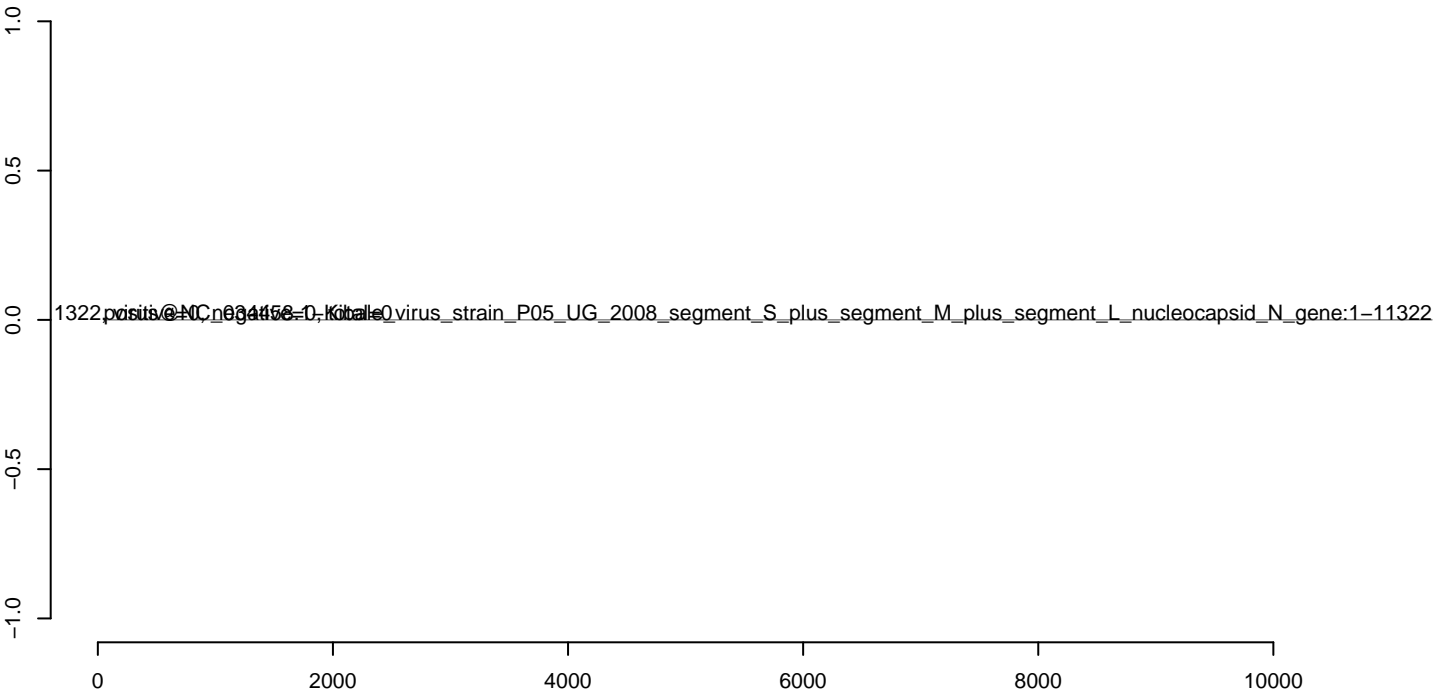
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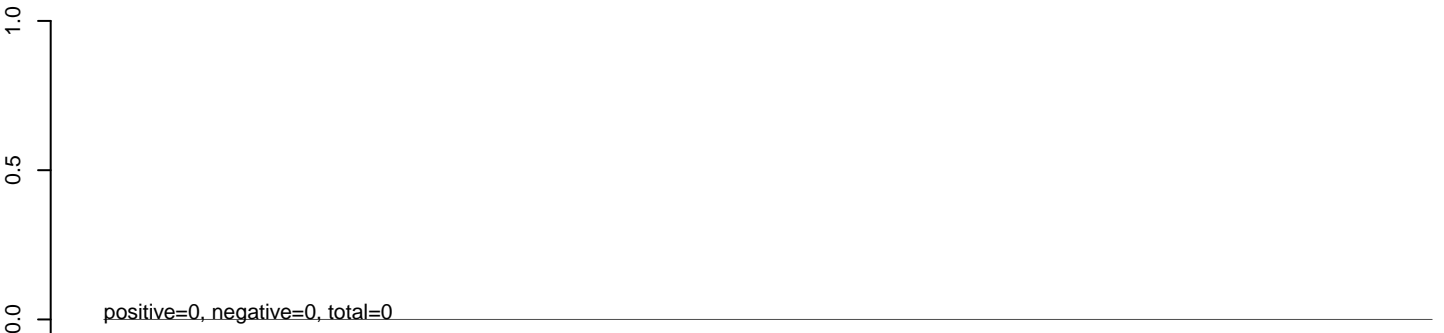
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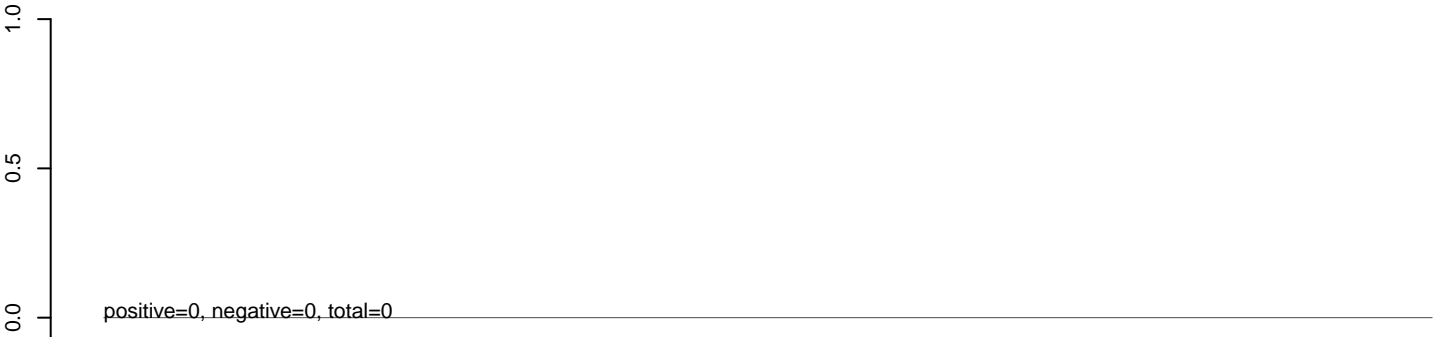
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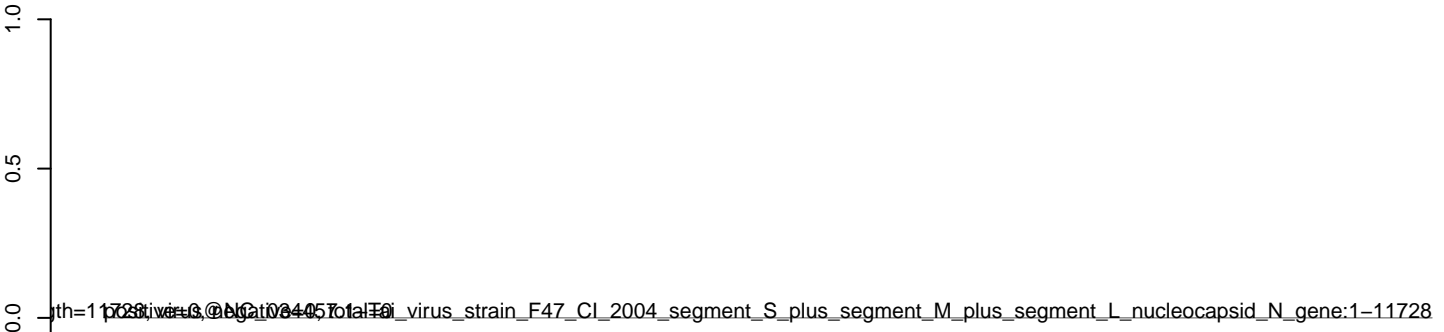
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AeAeg_CCL.125cells_CHIKV.rep



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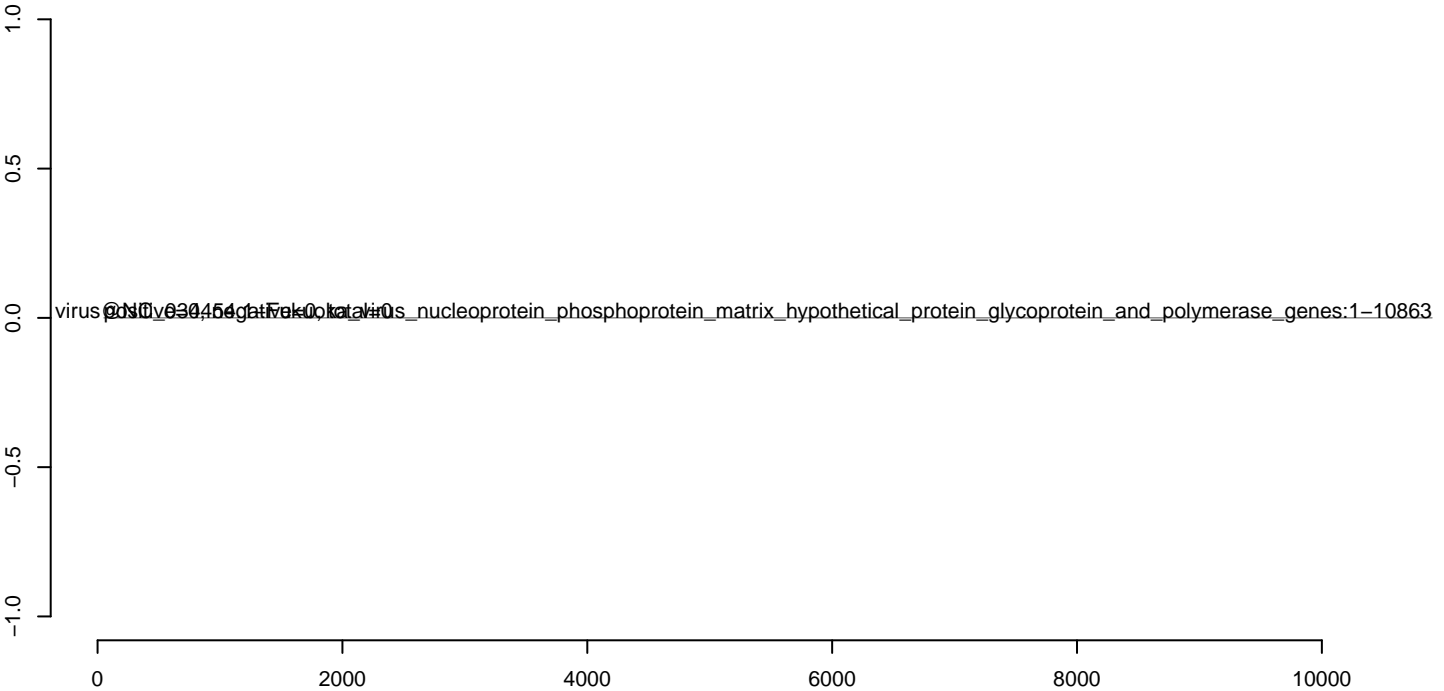
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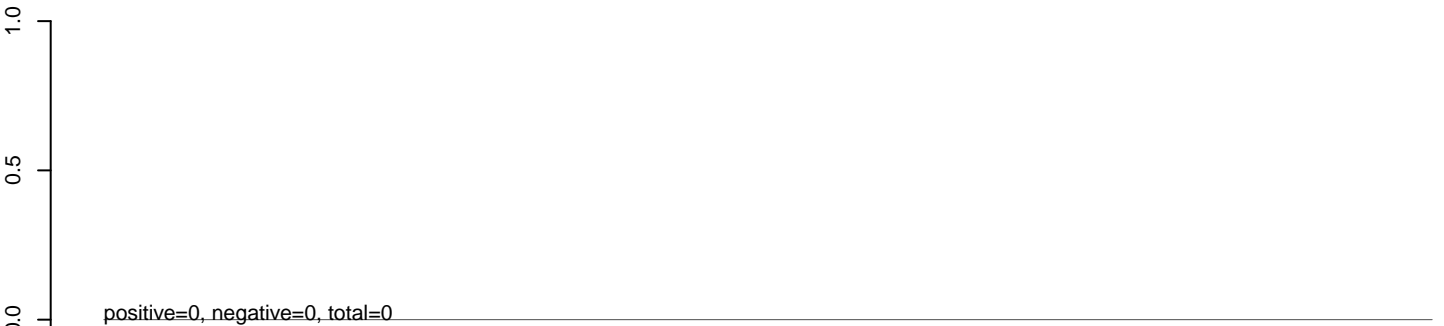
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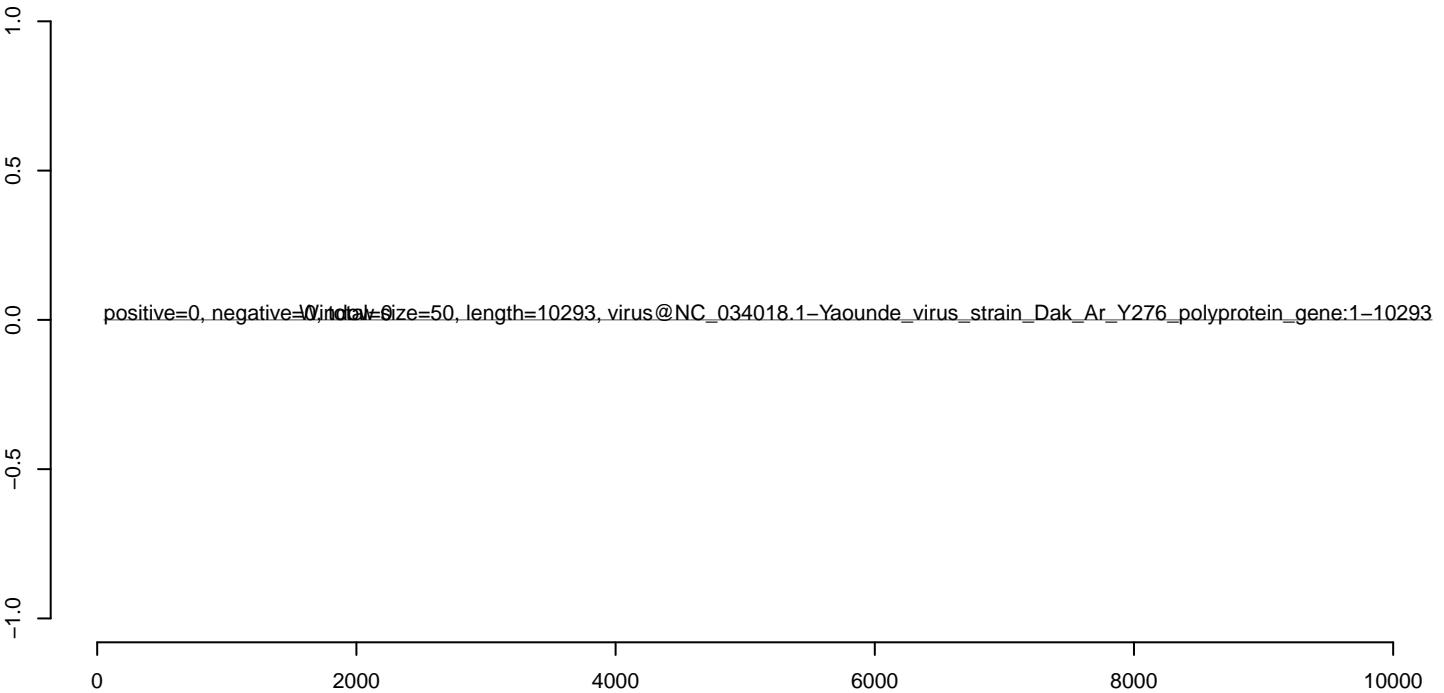
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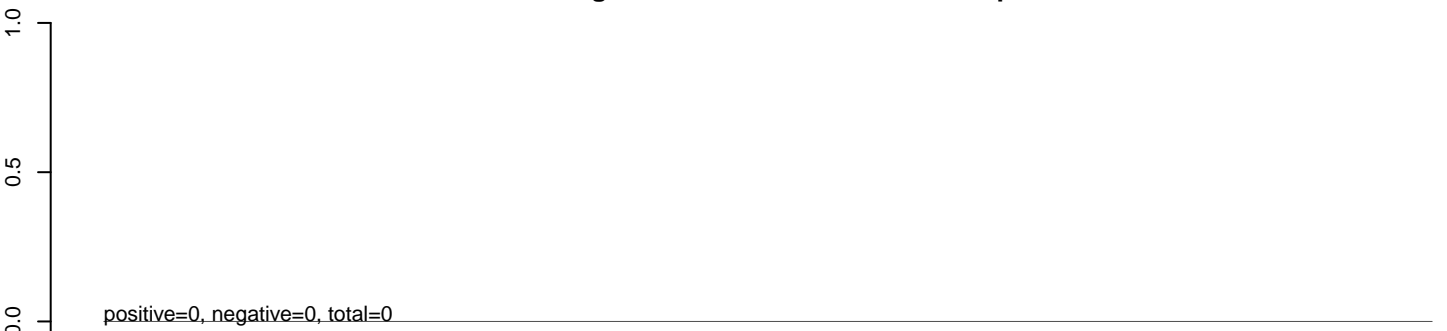
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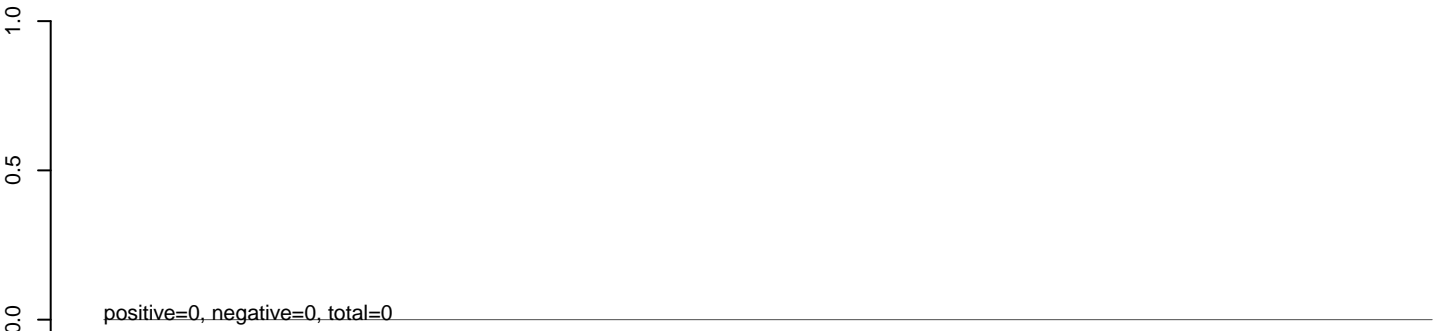
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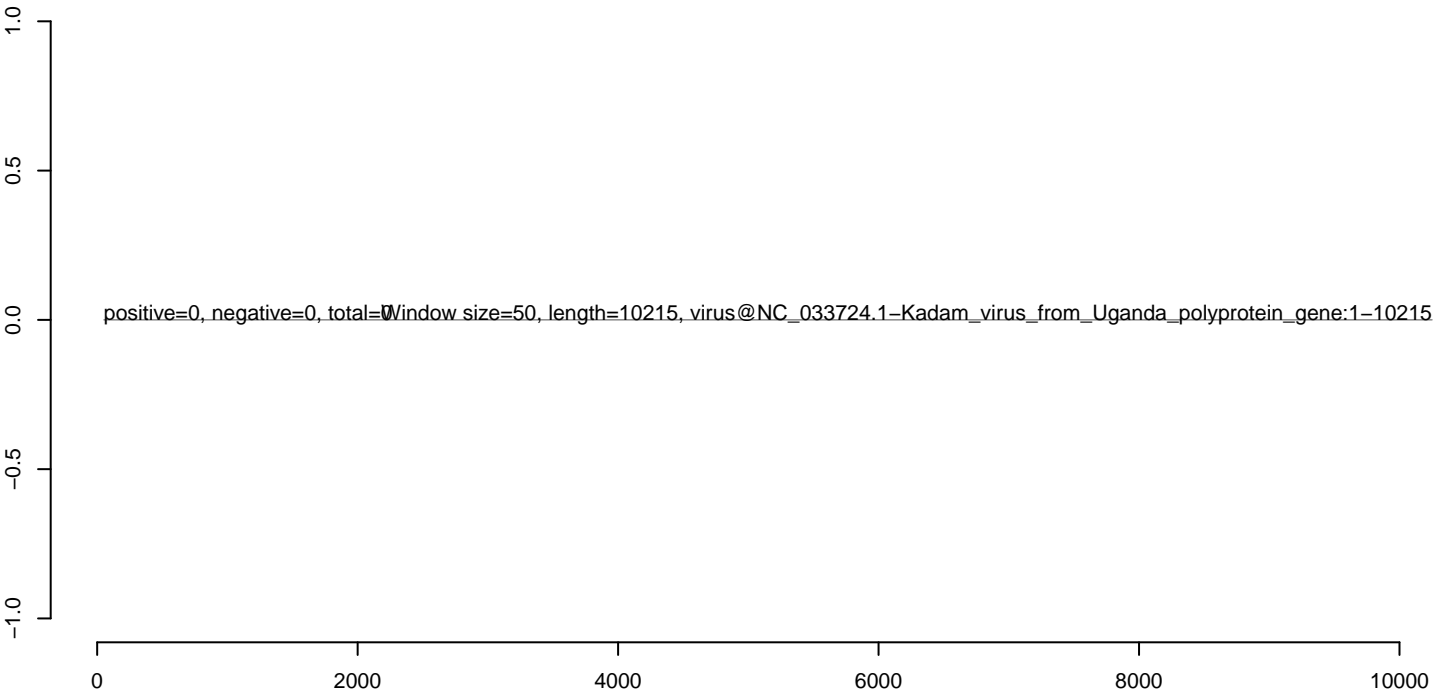
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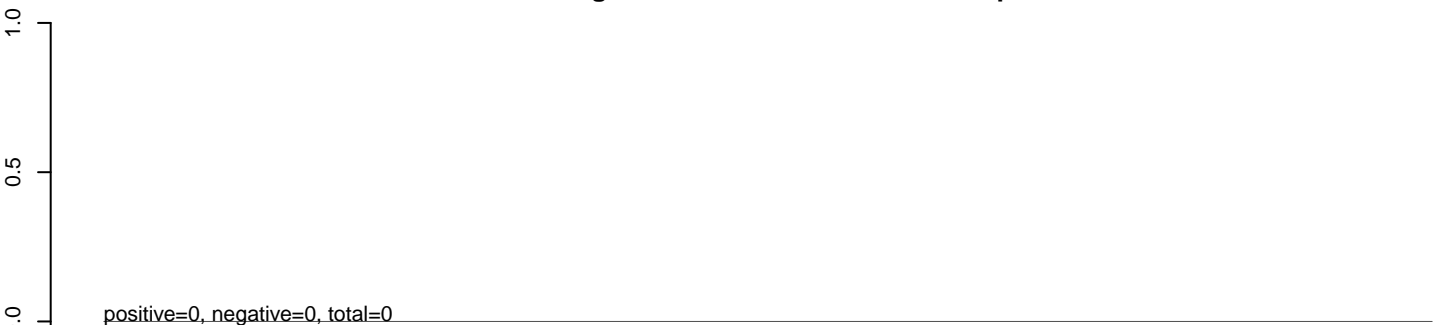
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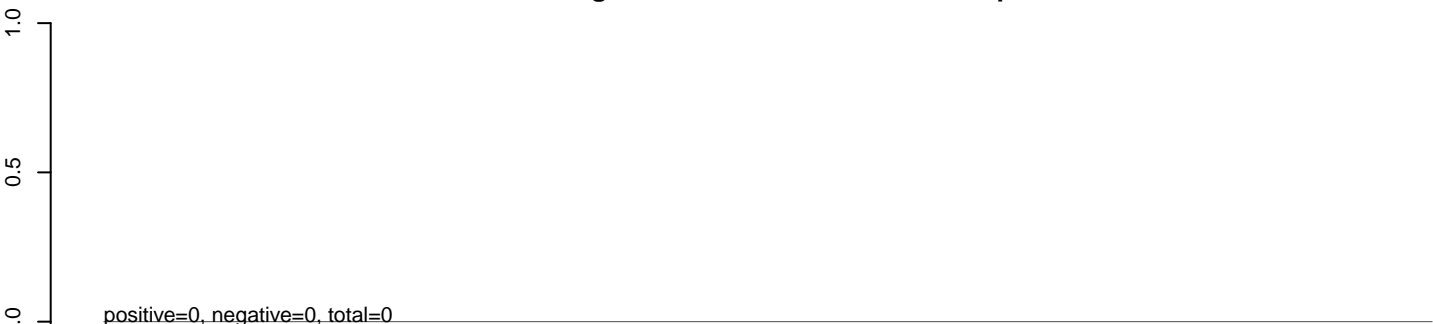
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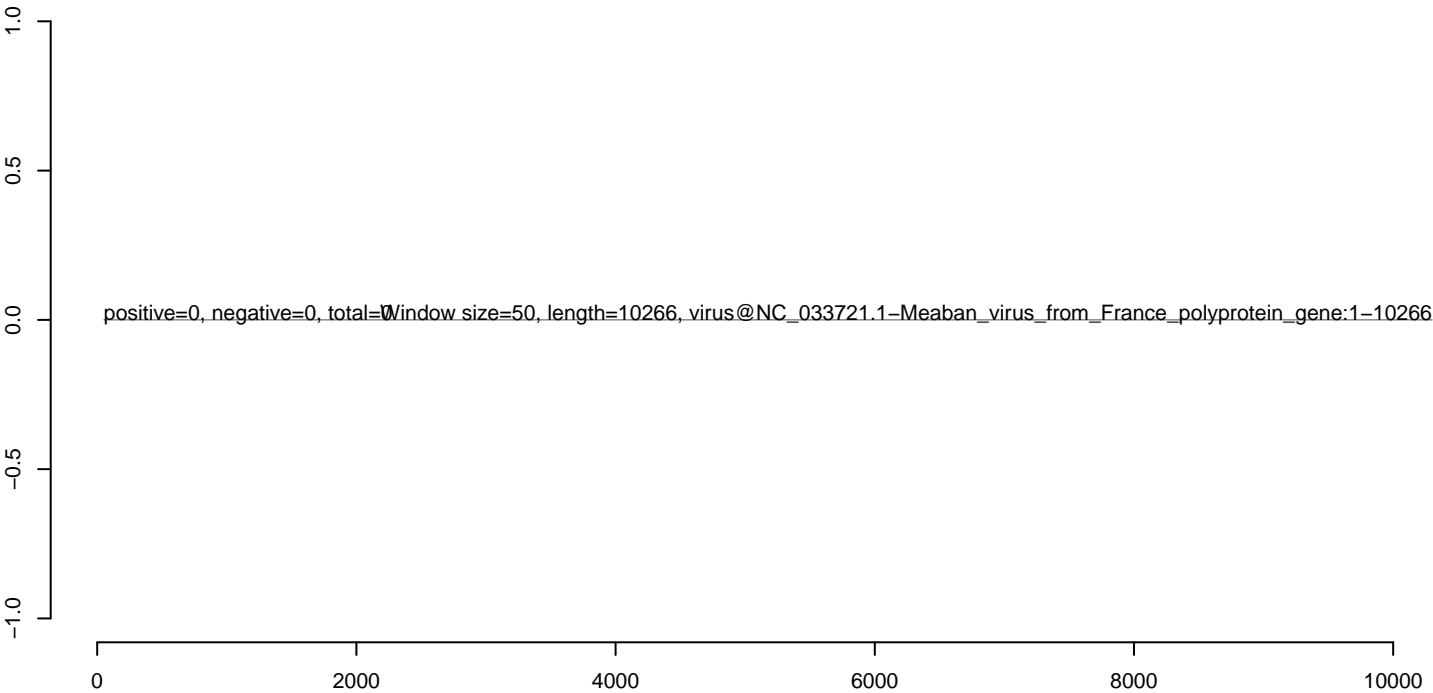
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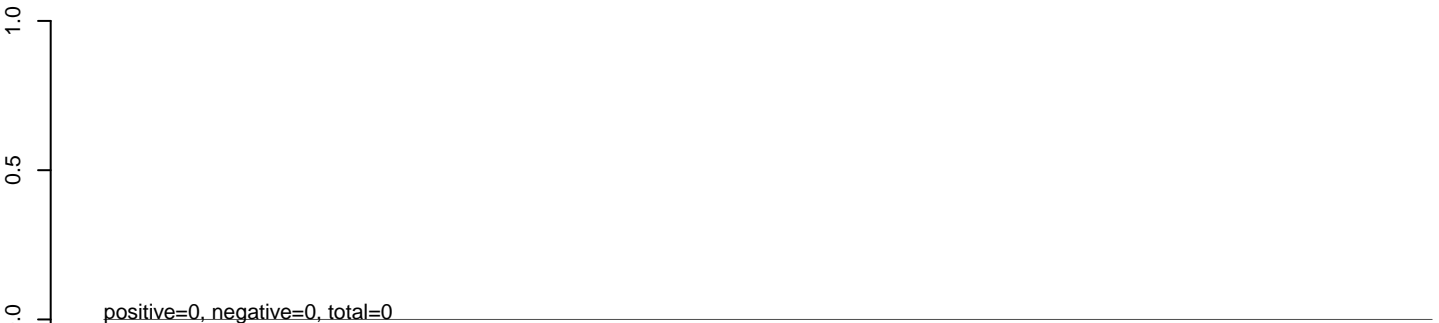
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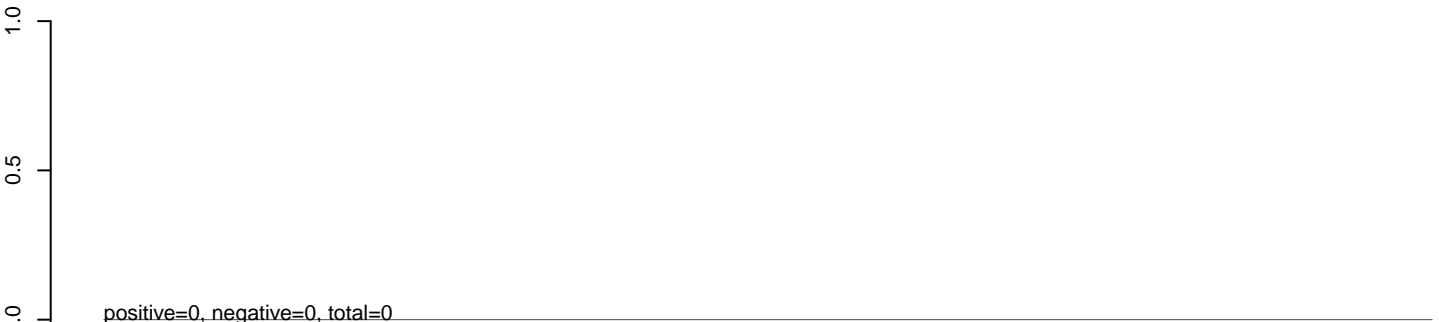
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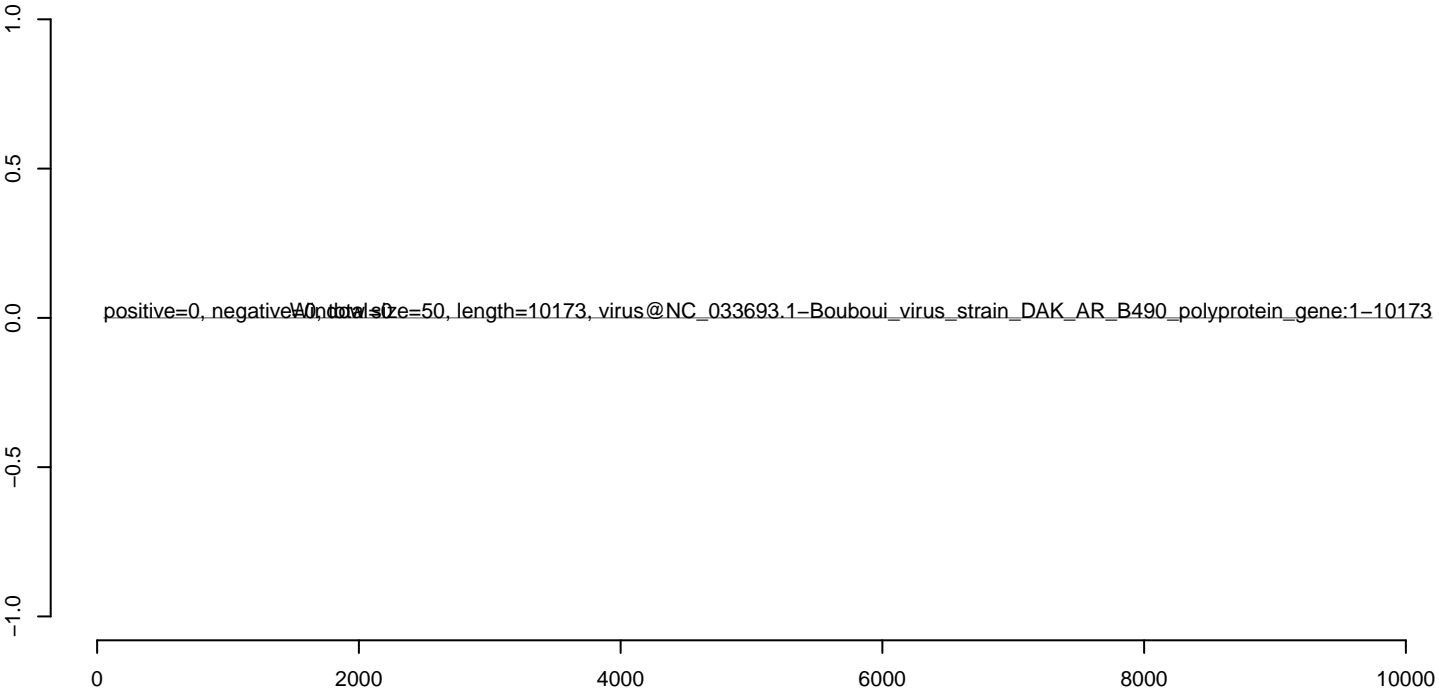
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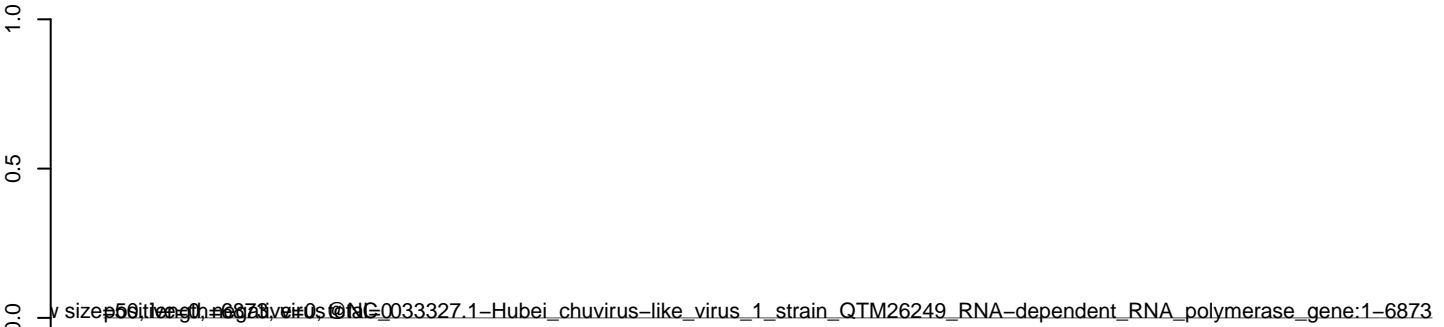
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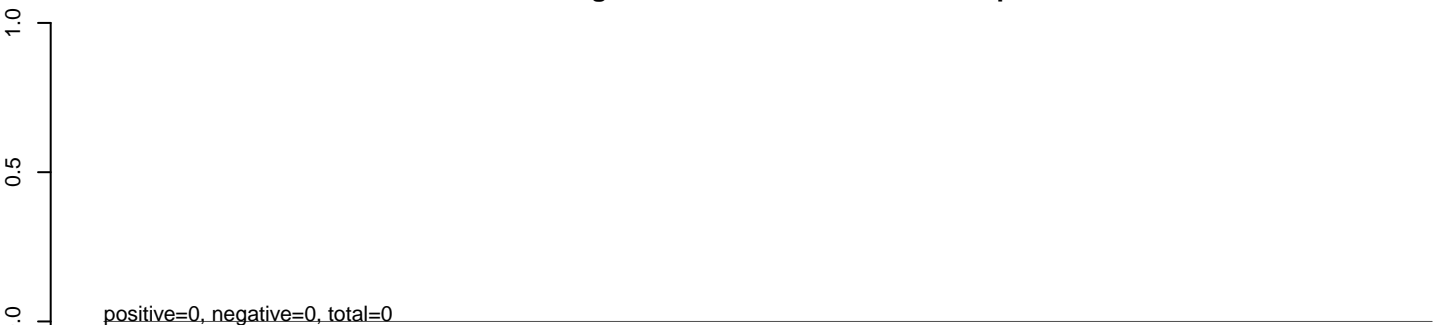


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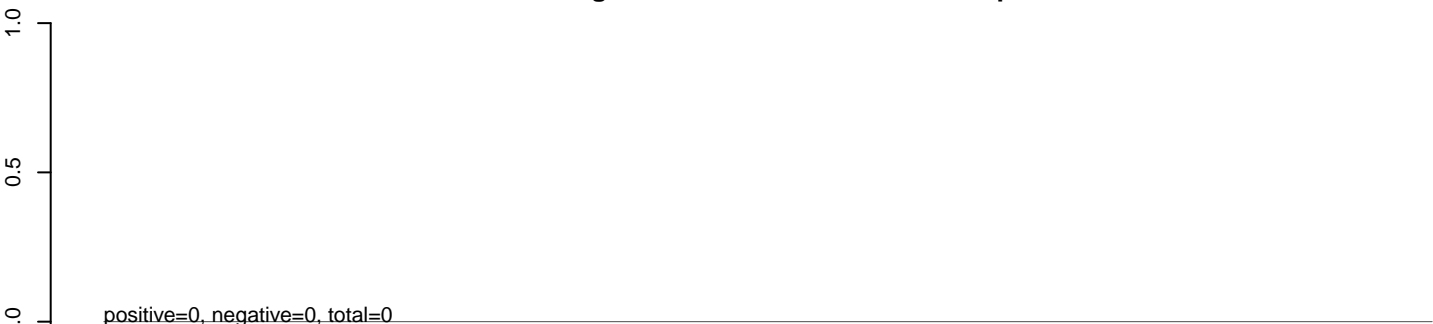


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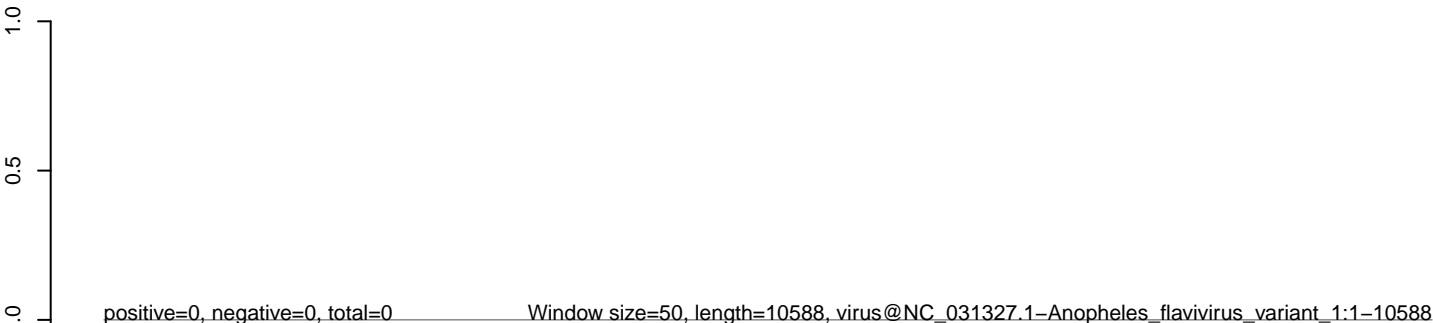
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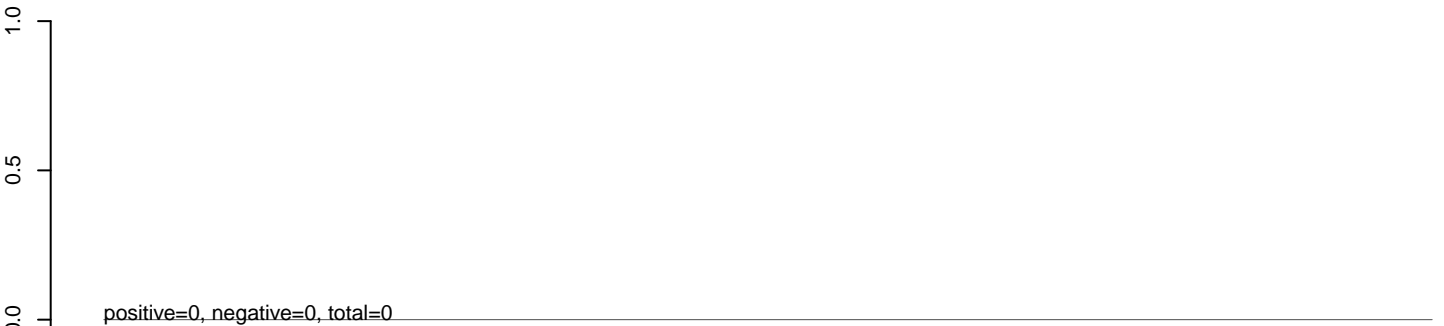


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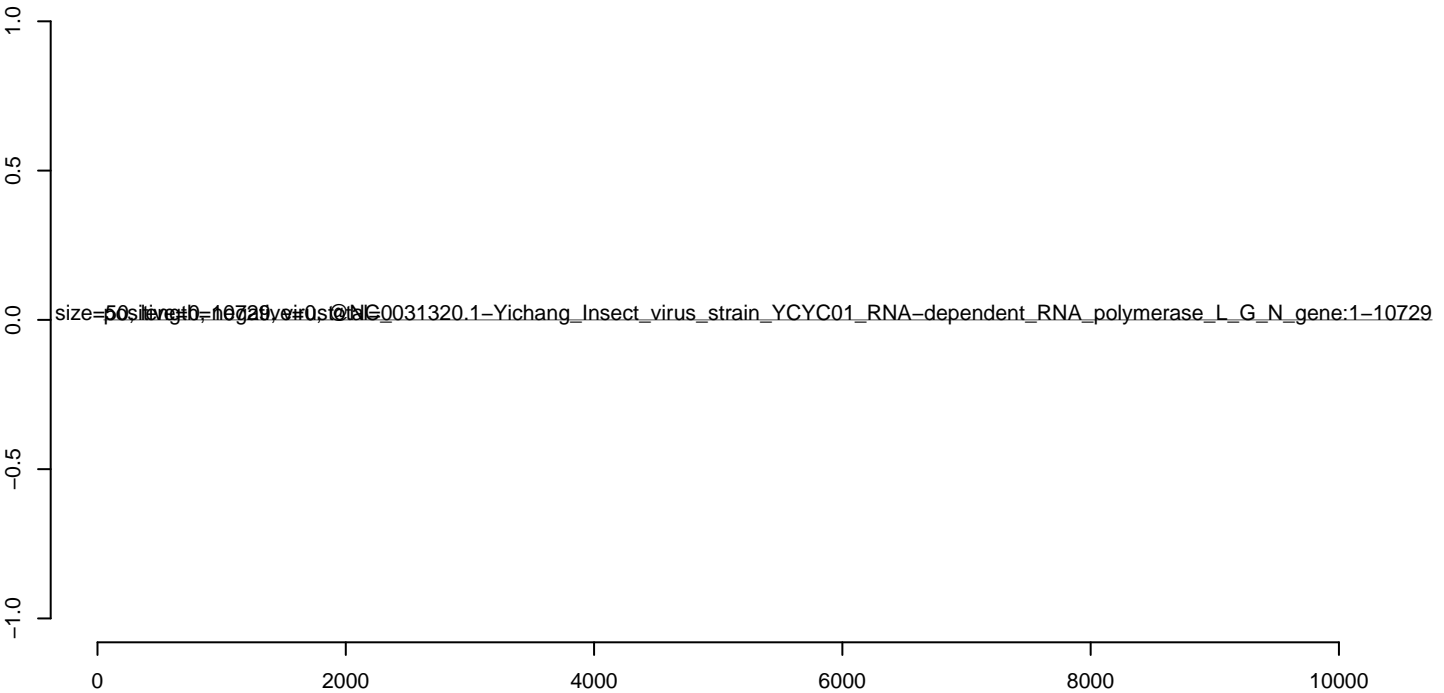
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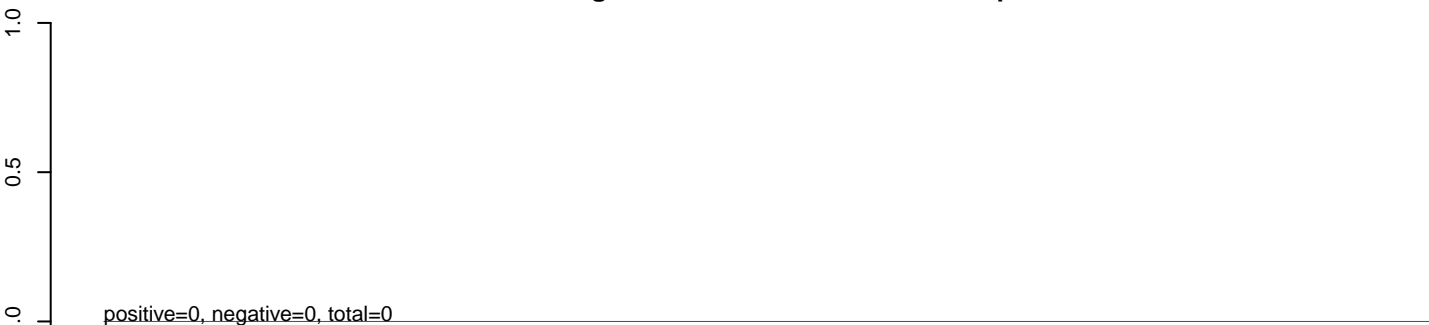
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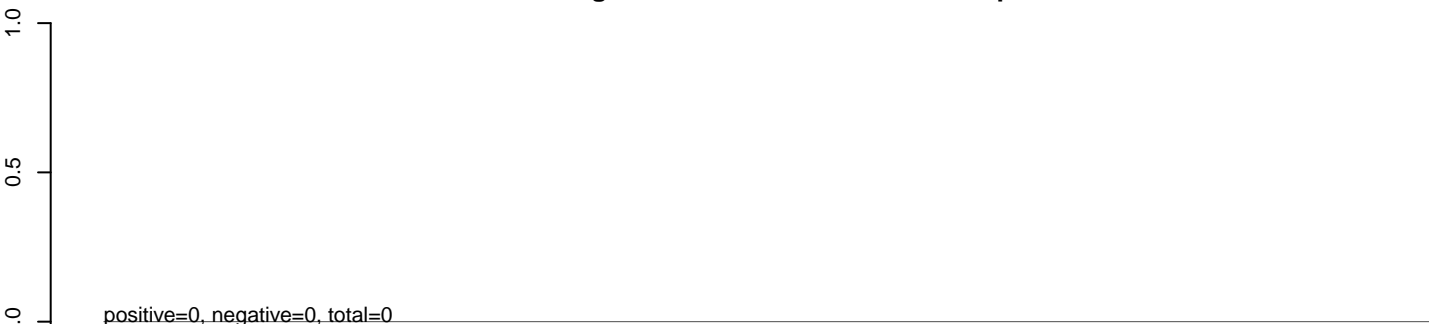
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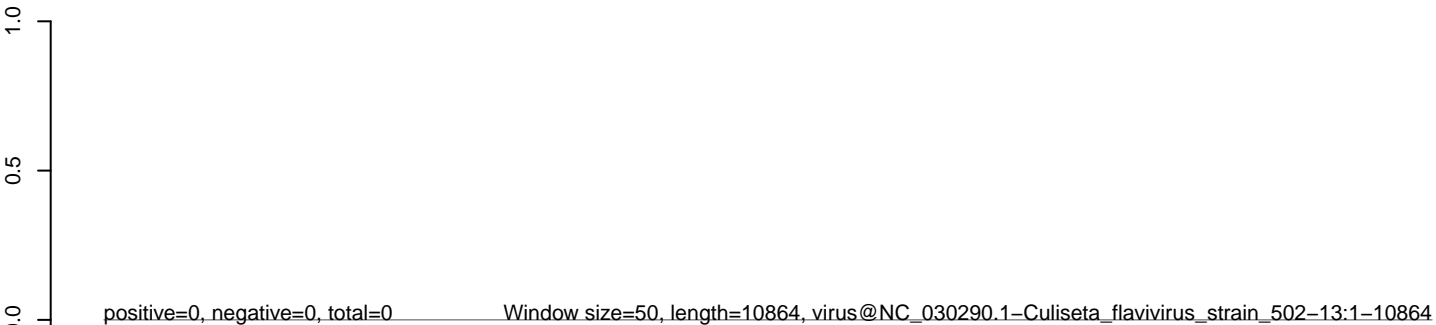
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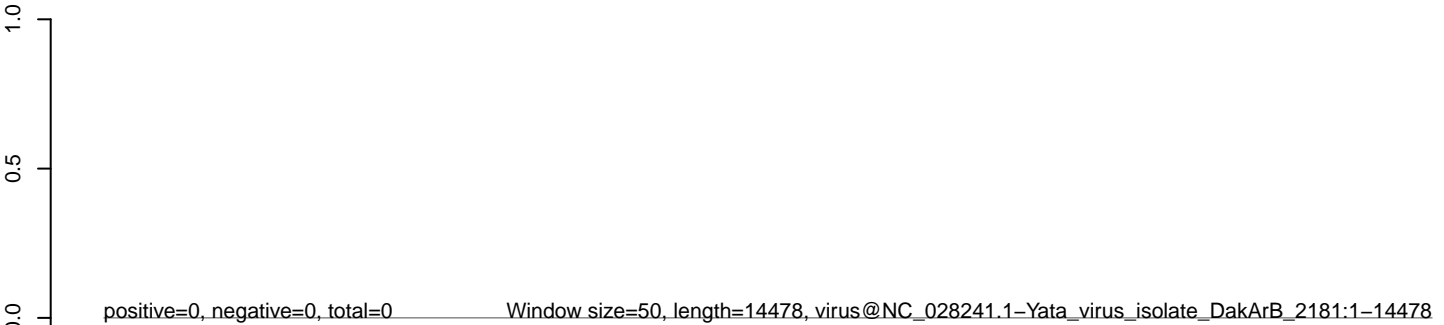
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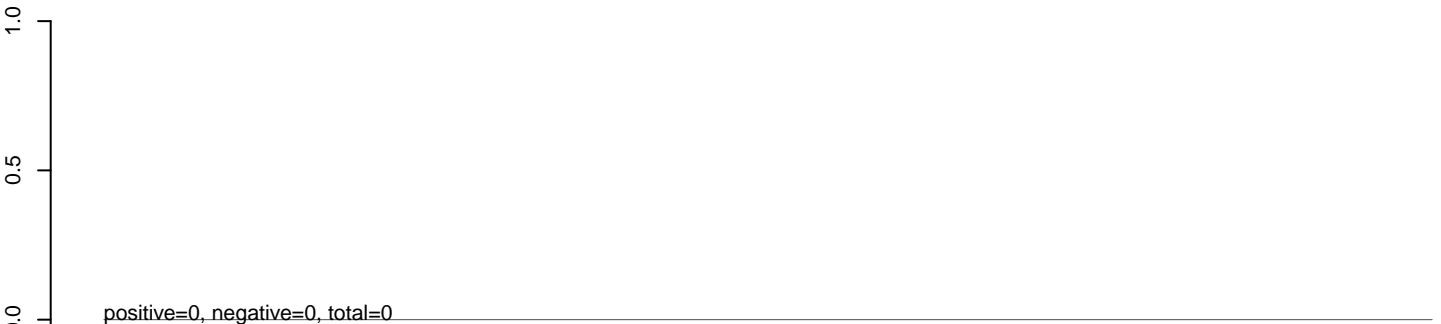
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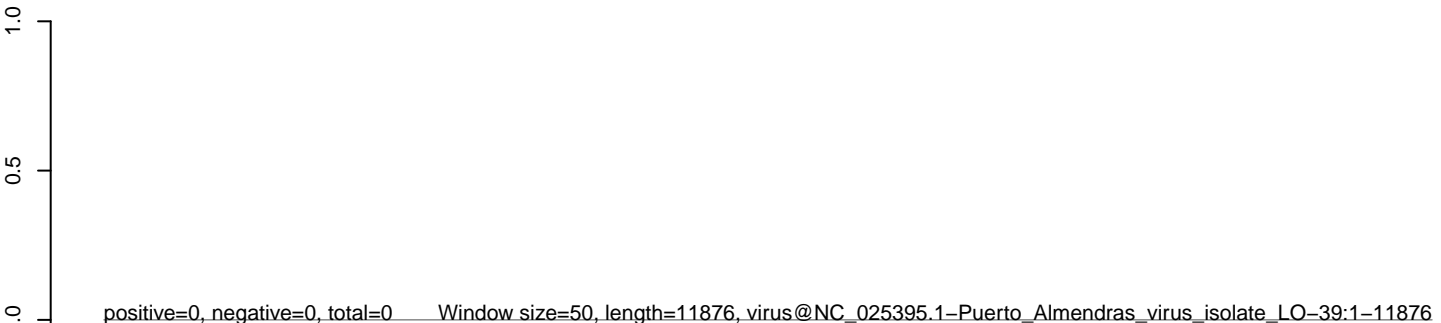
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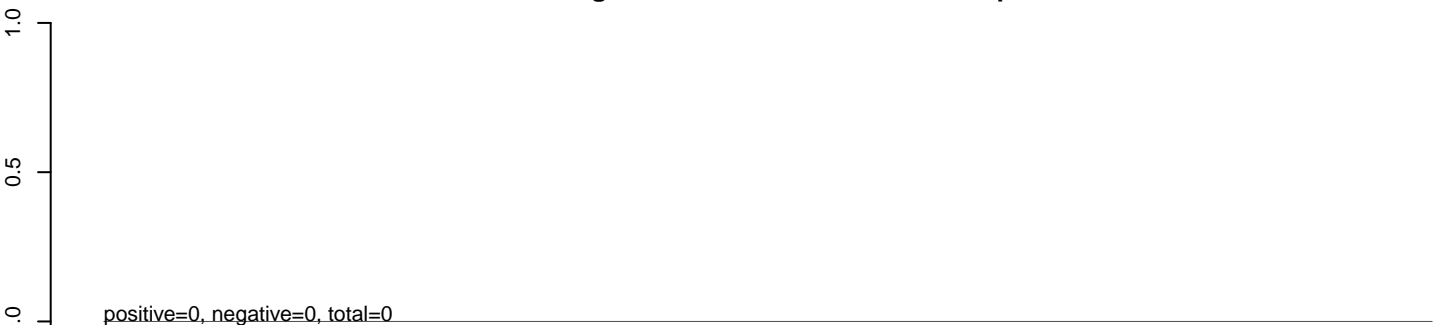
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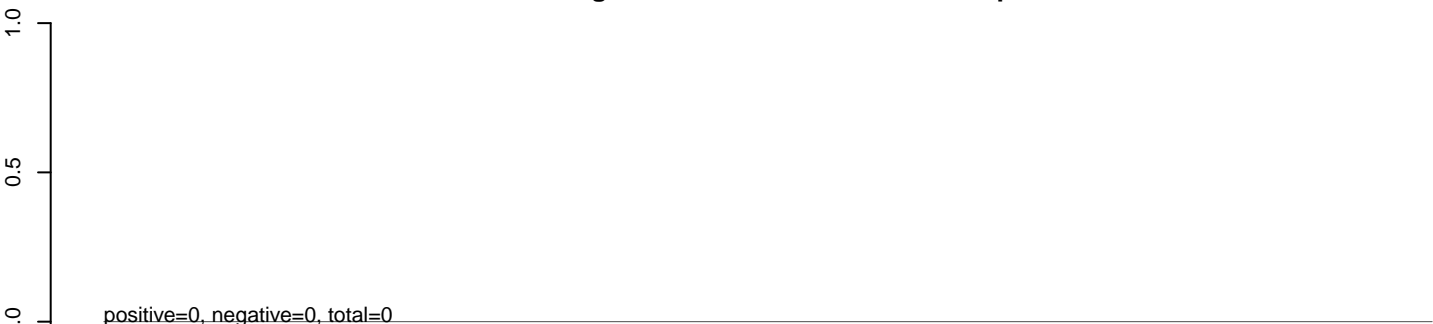
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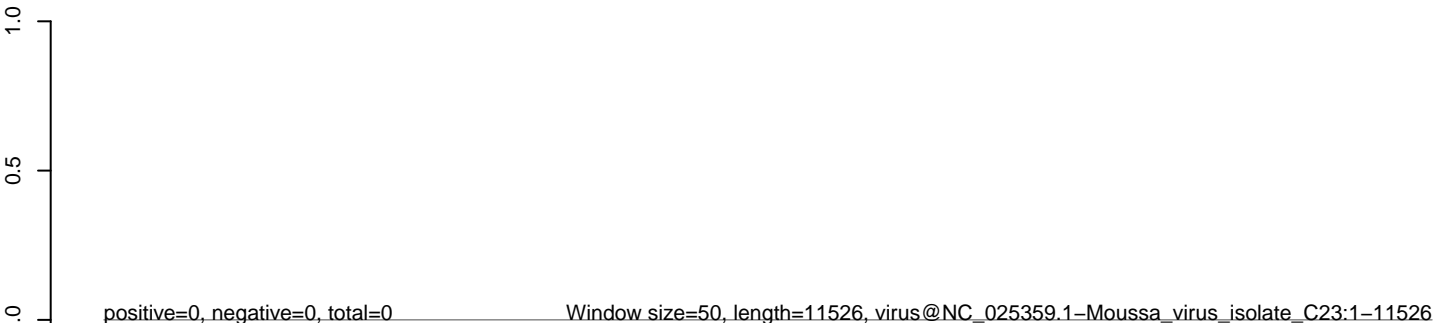
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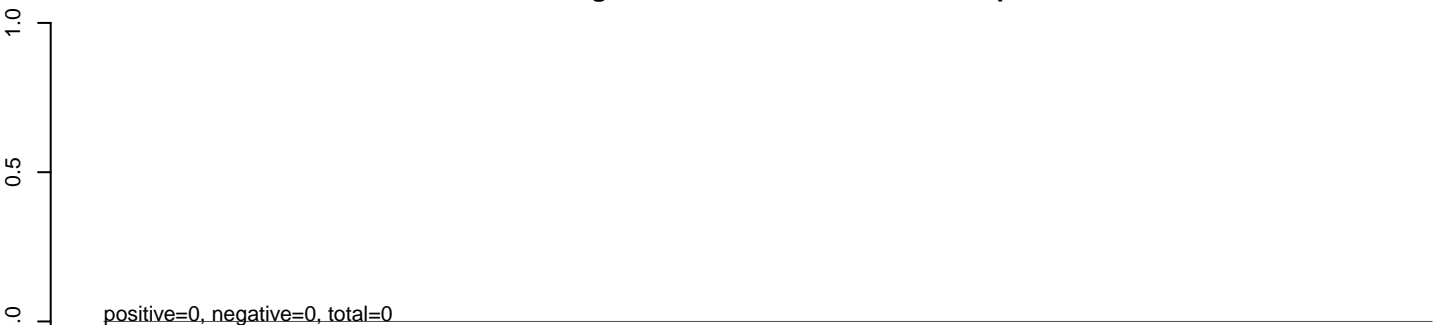


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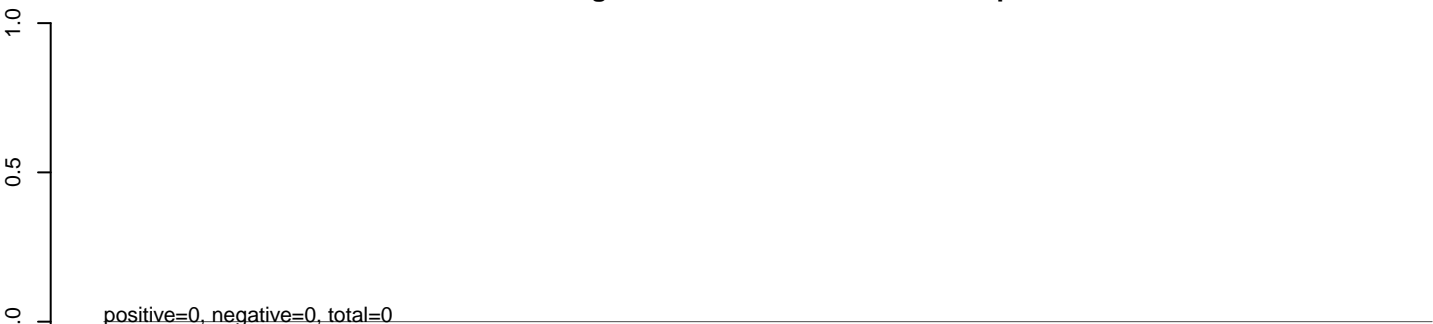


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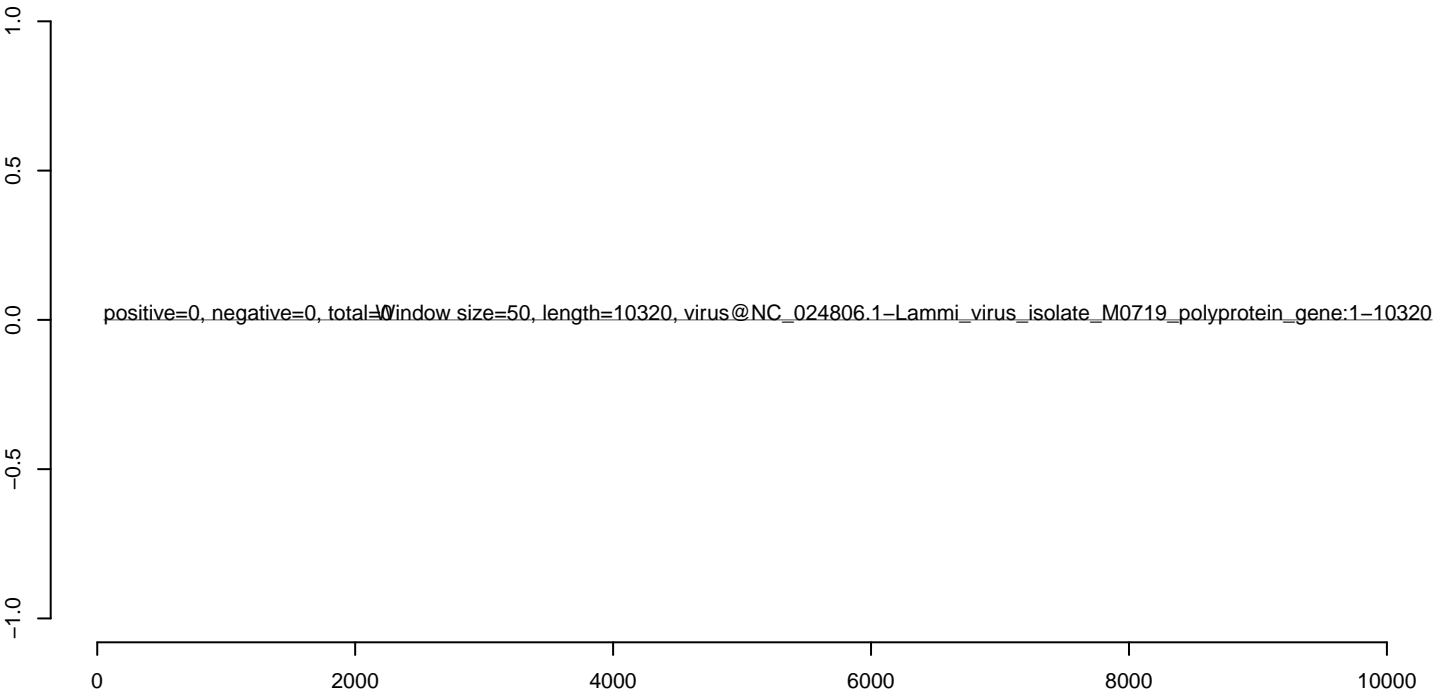
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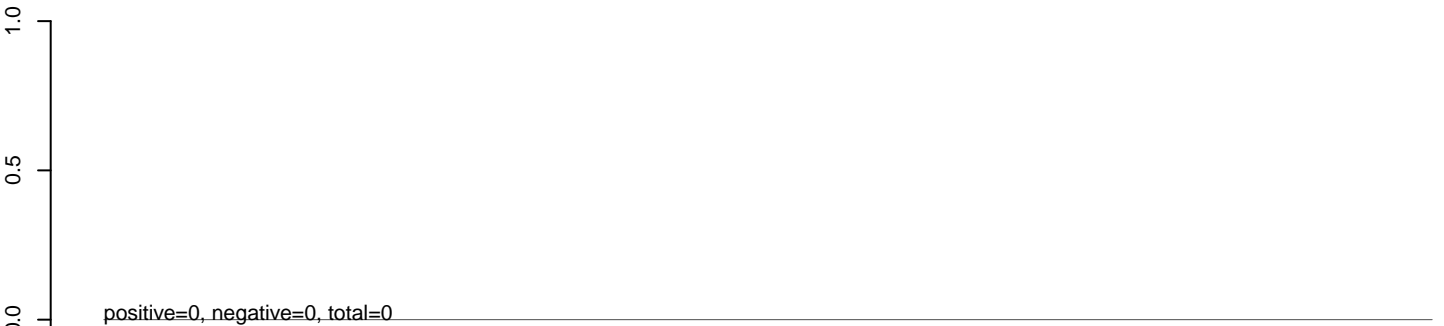
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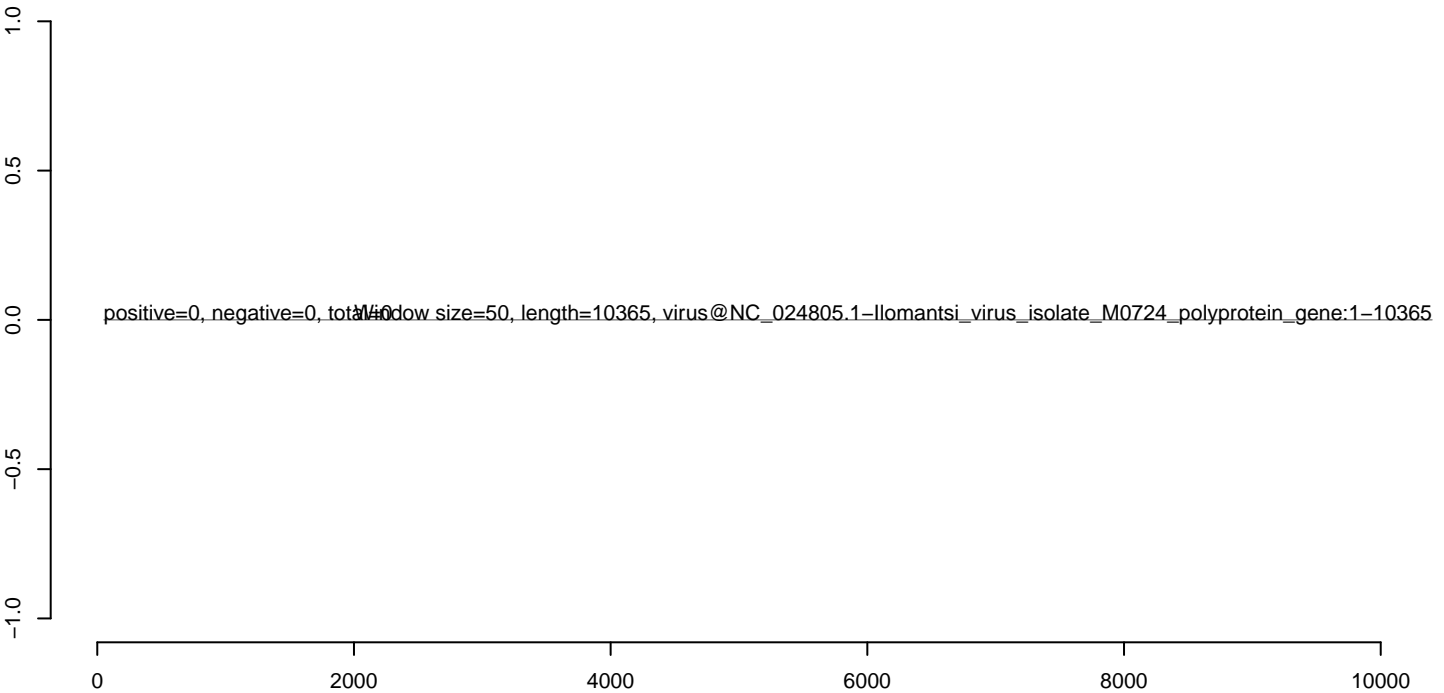
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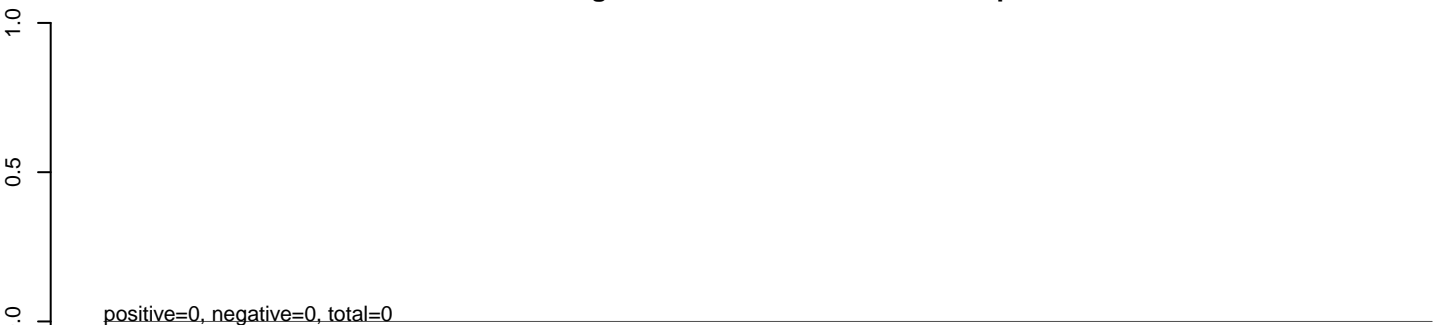
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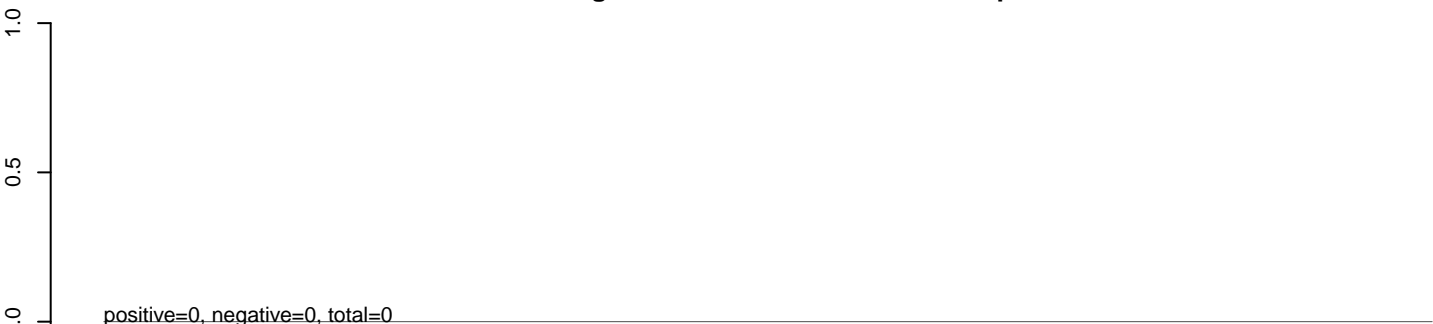
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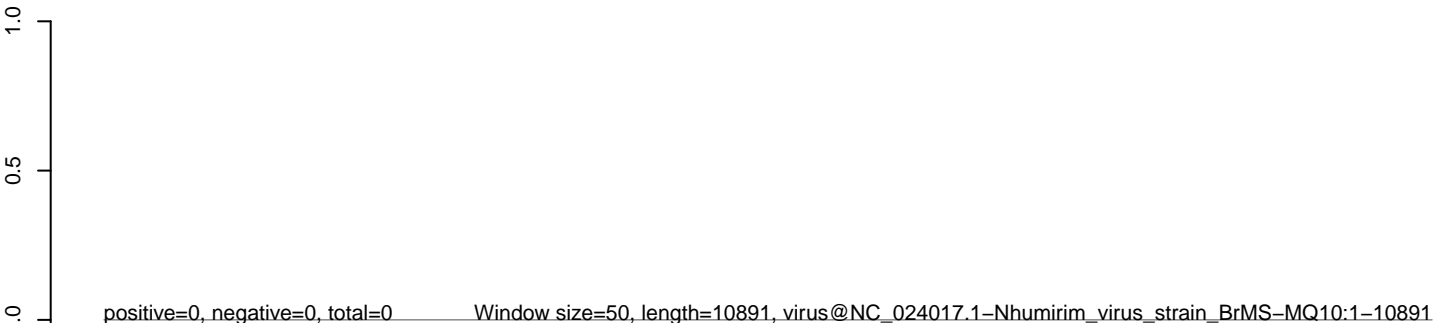
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AeAeg_CCL.125cells_CHIKV.rep



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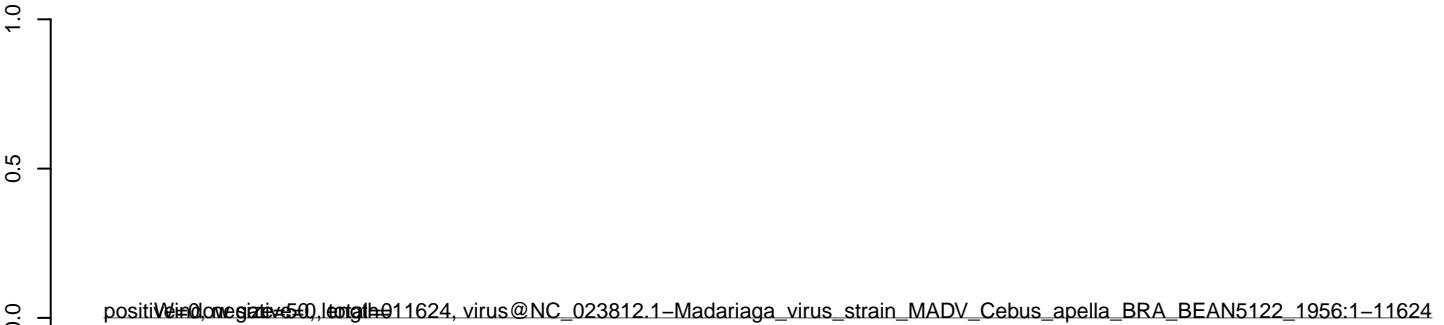
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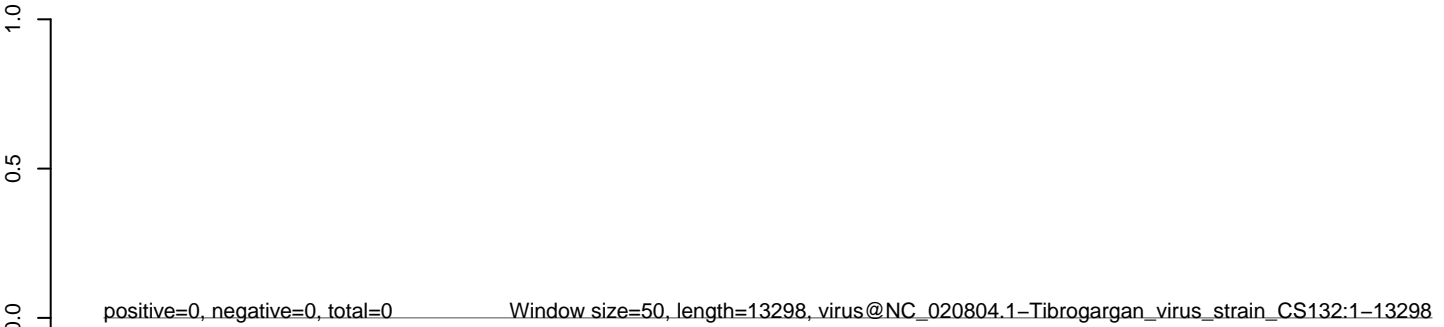
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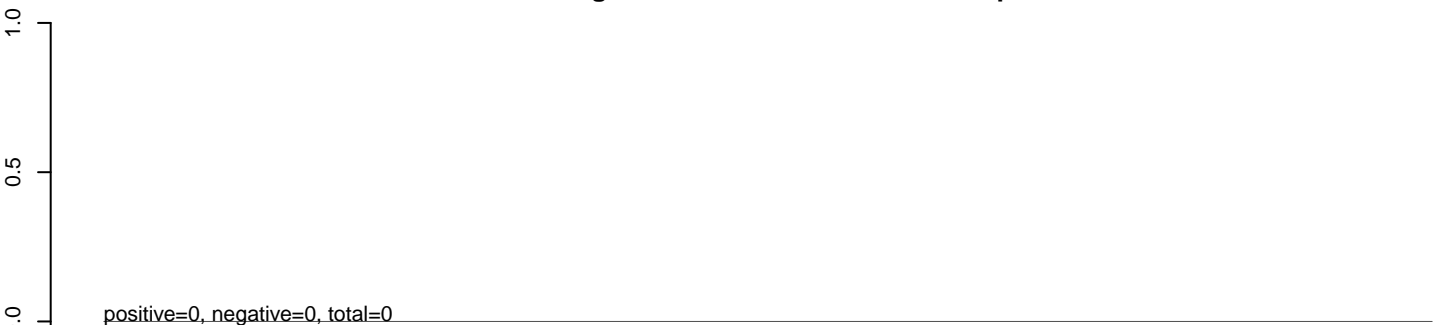


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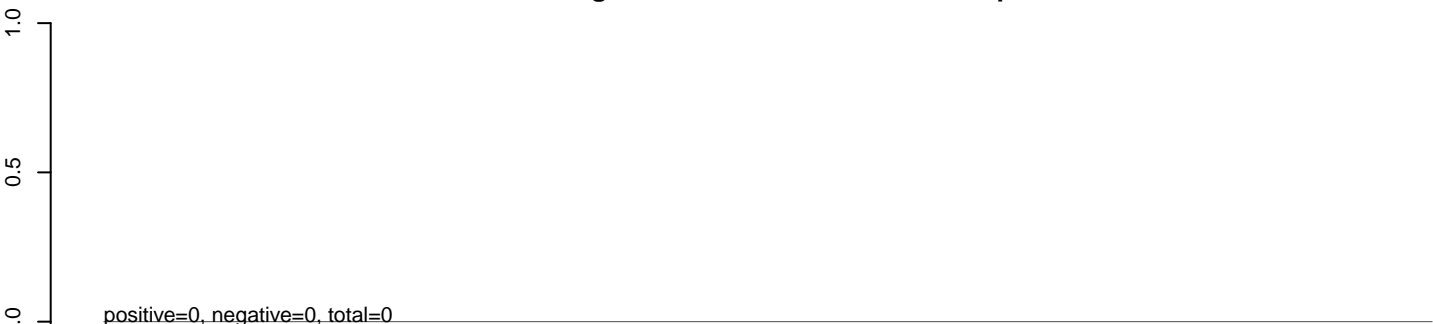


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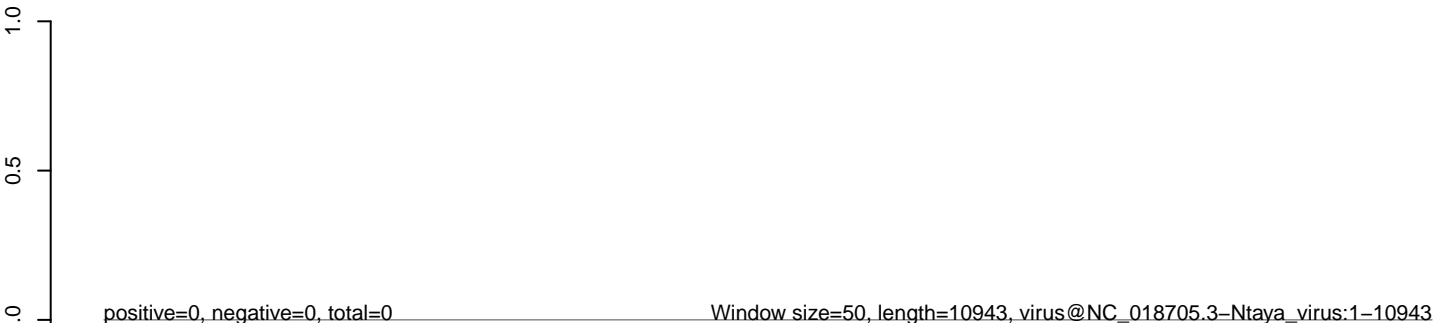
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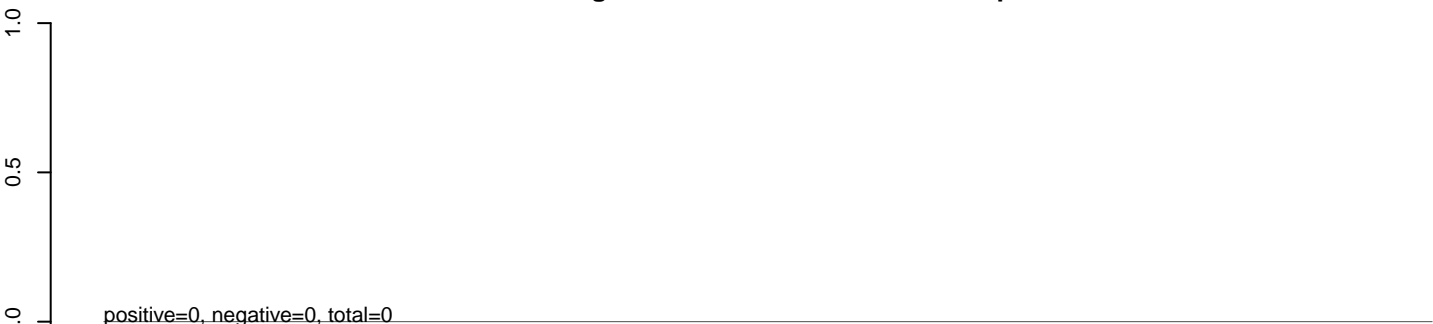


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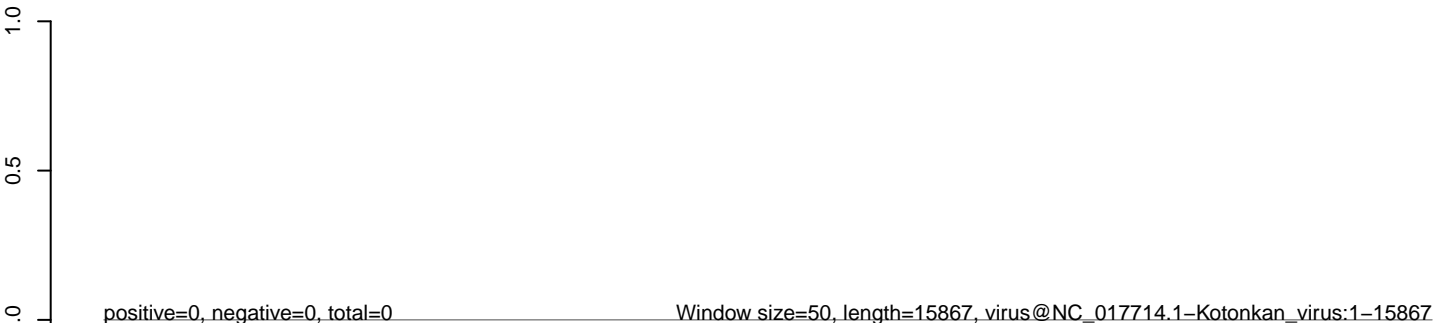
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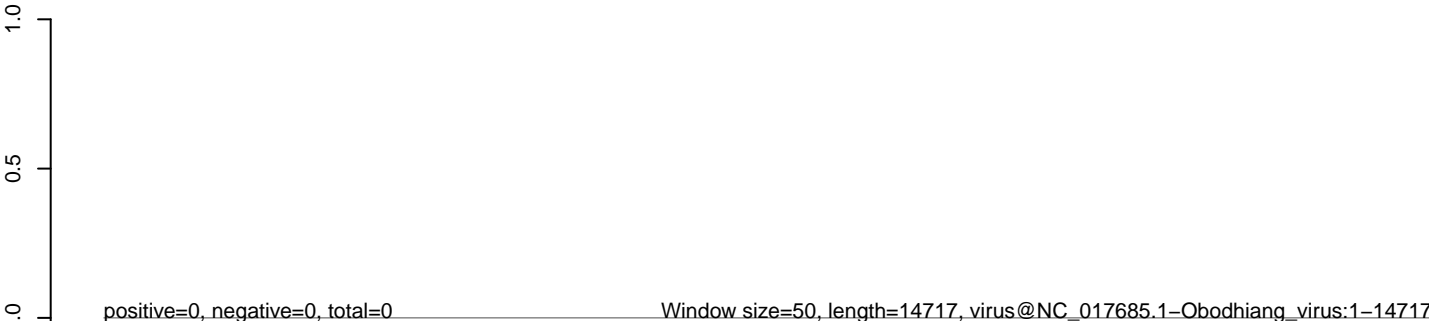
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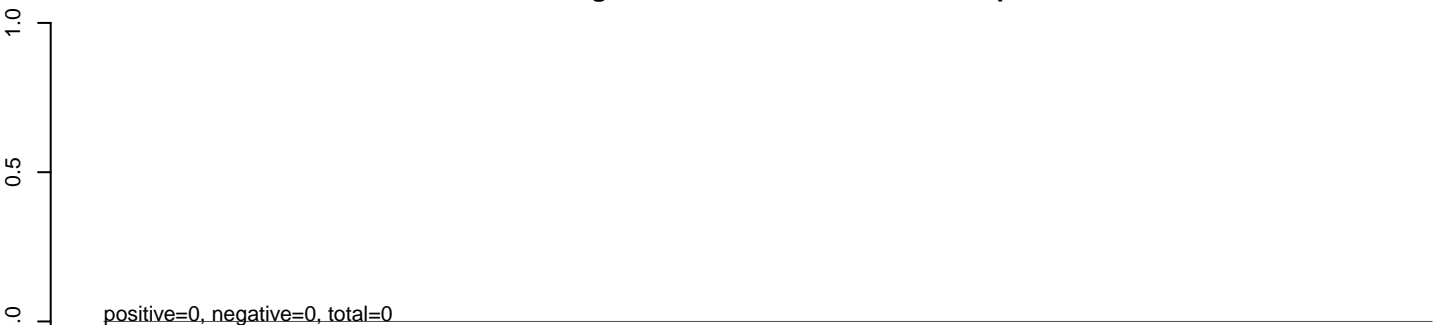
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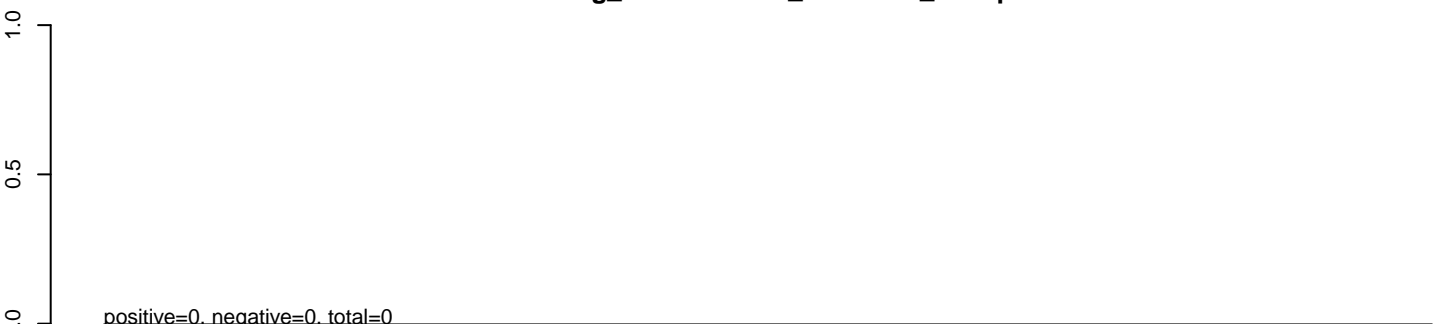
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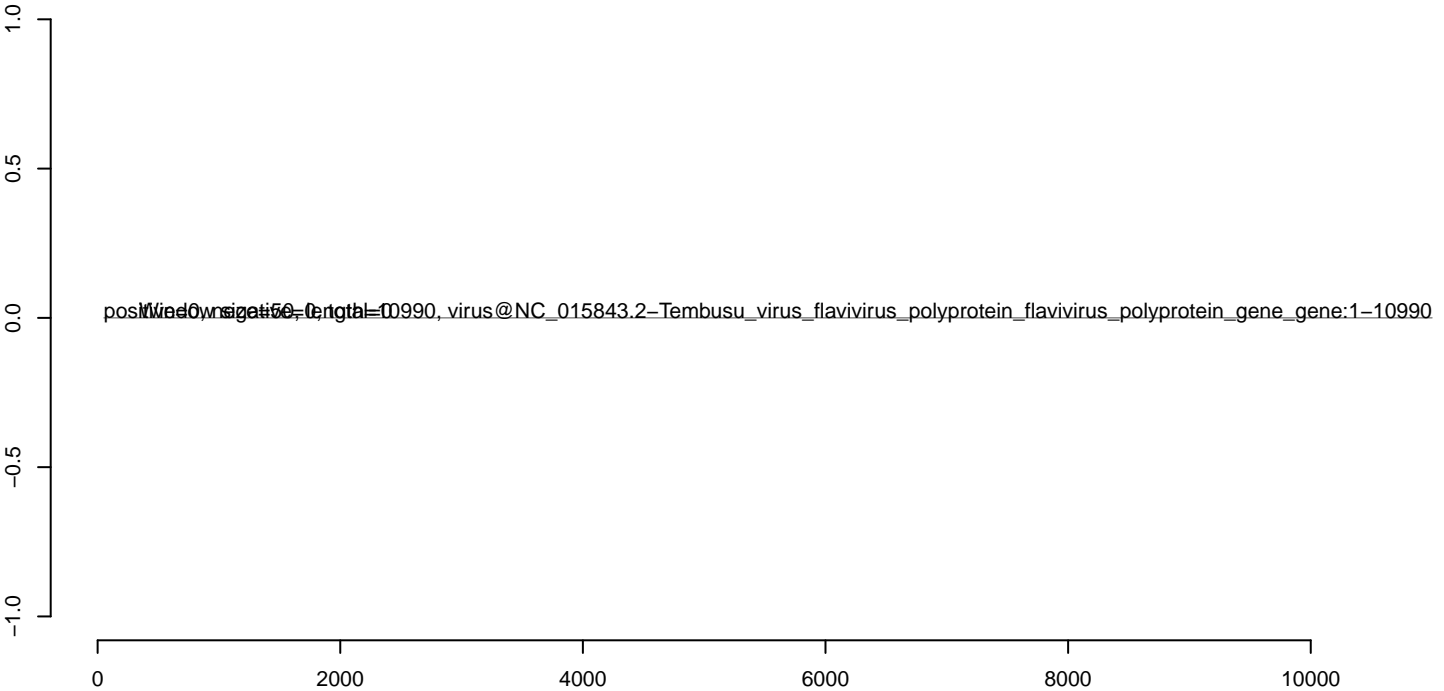
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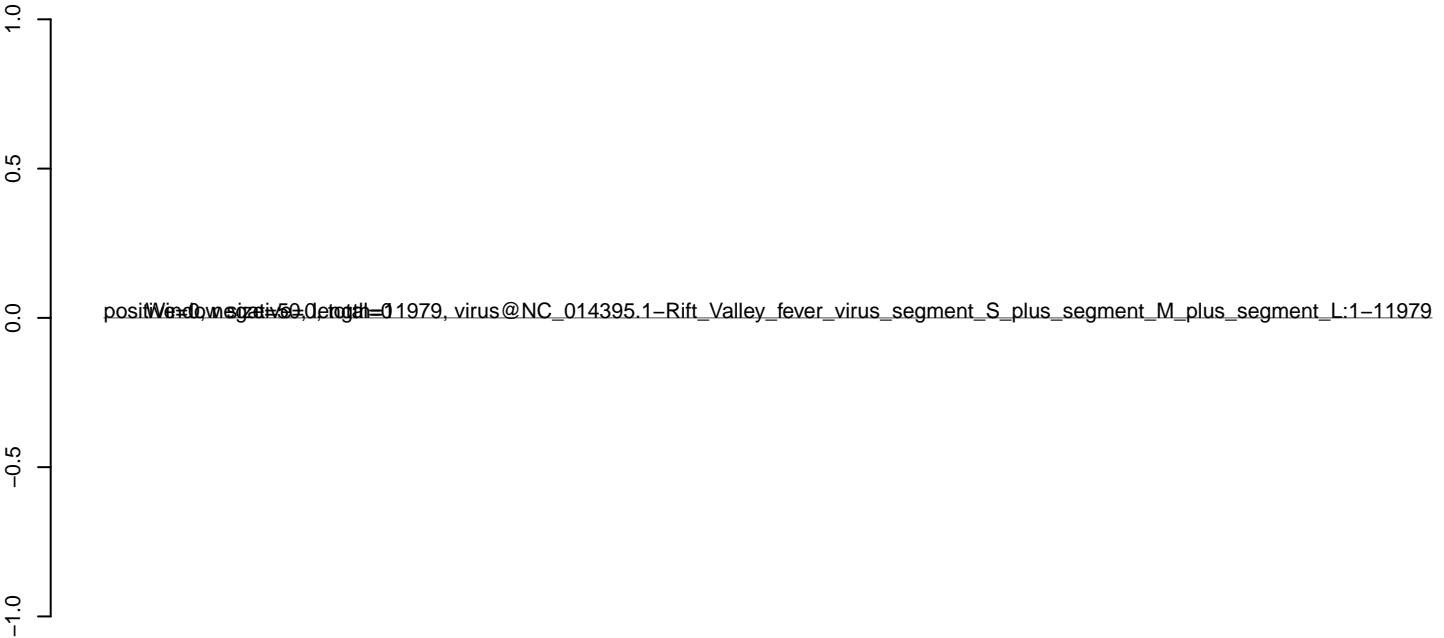
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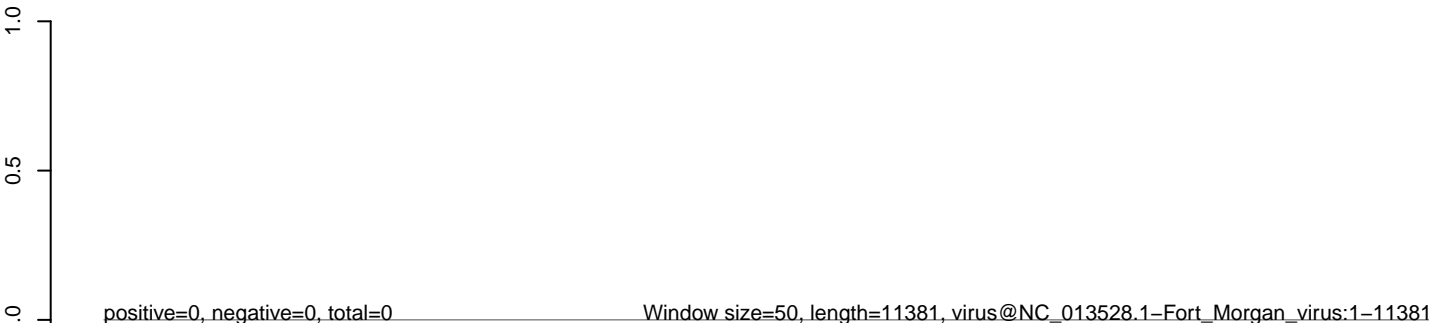
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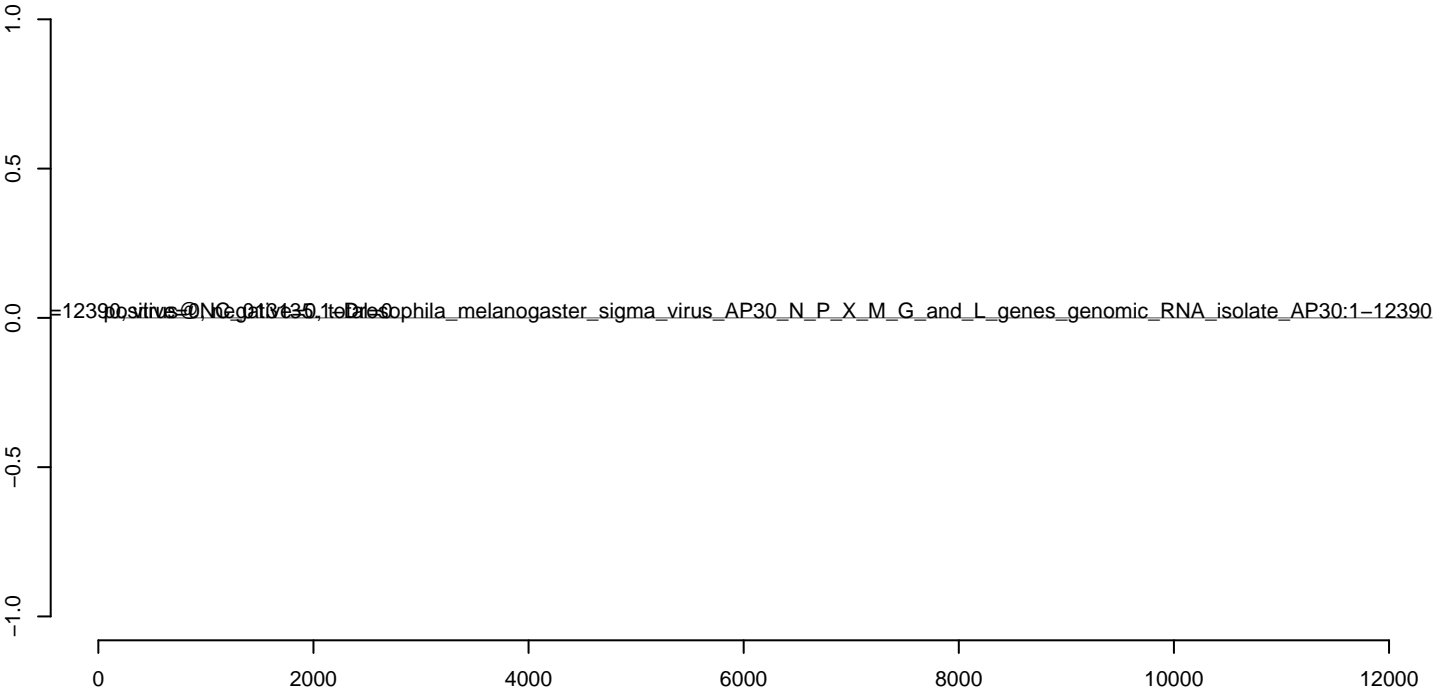
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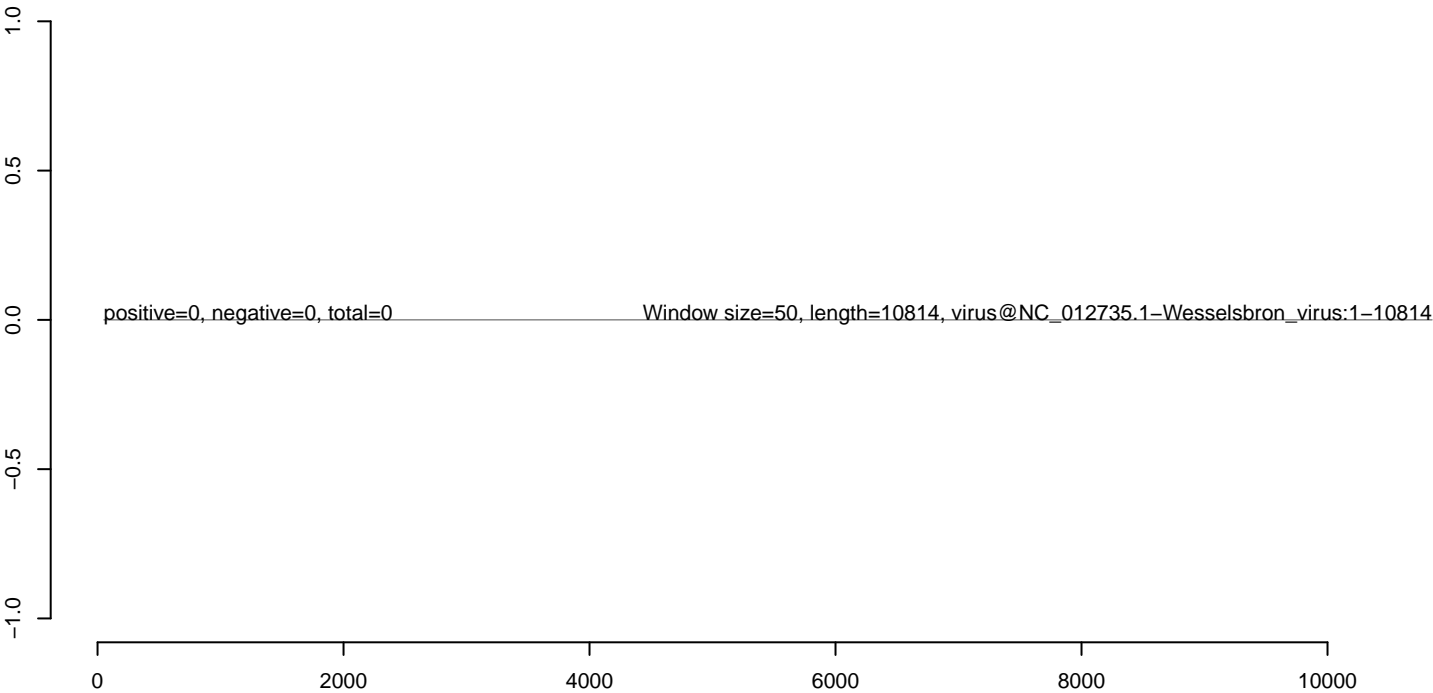
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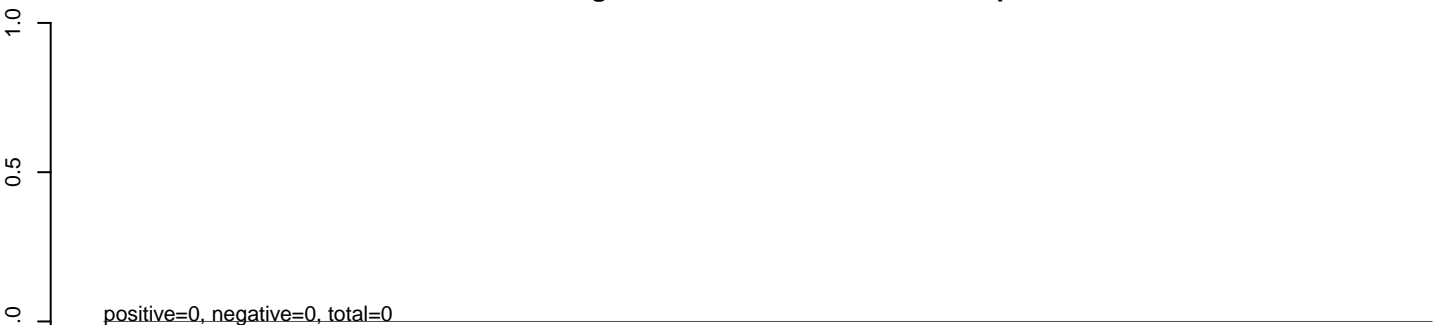
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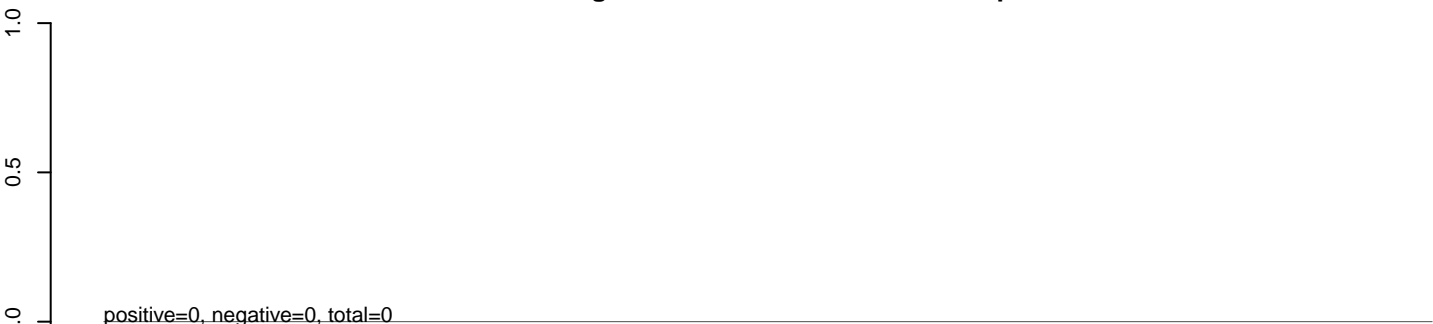
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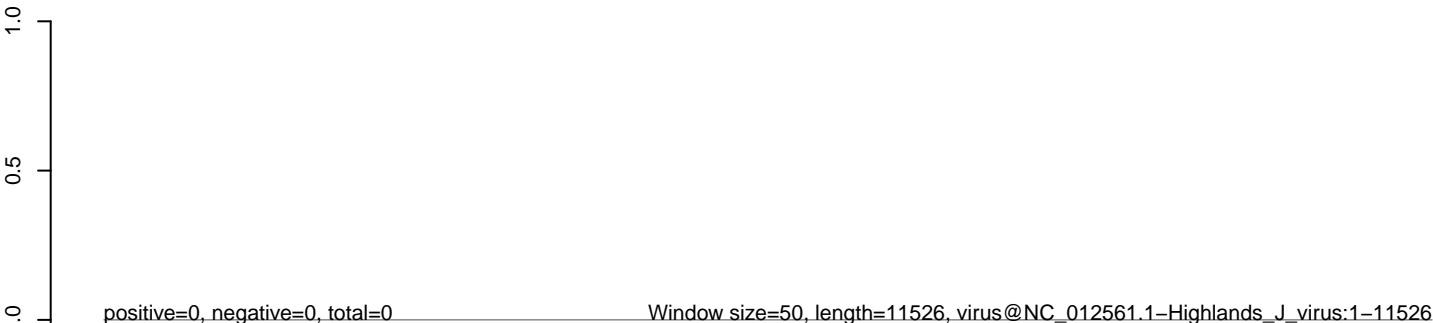
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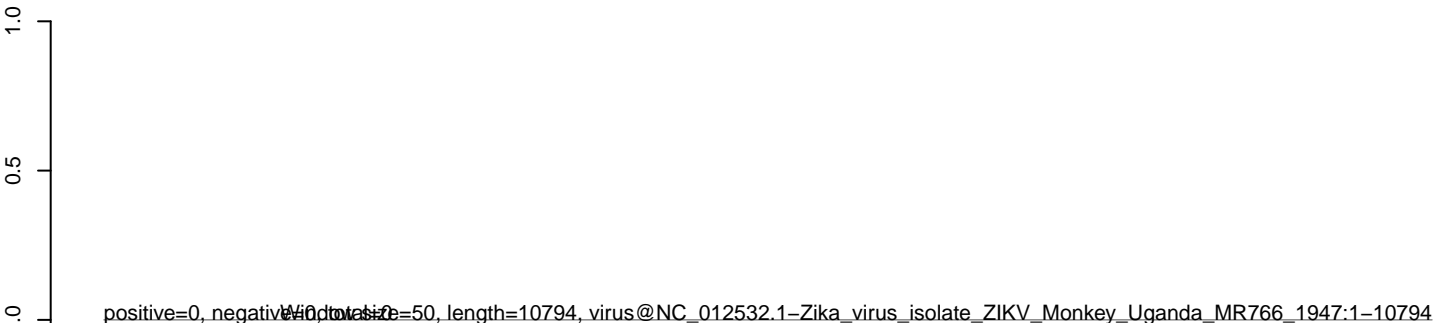
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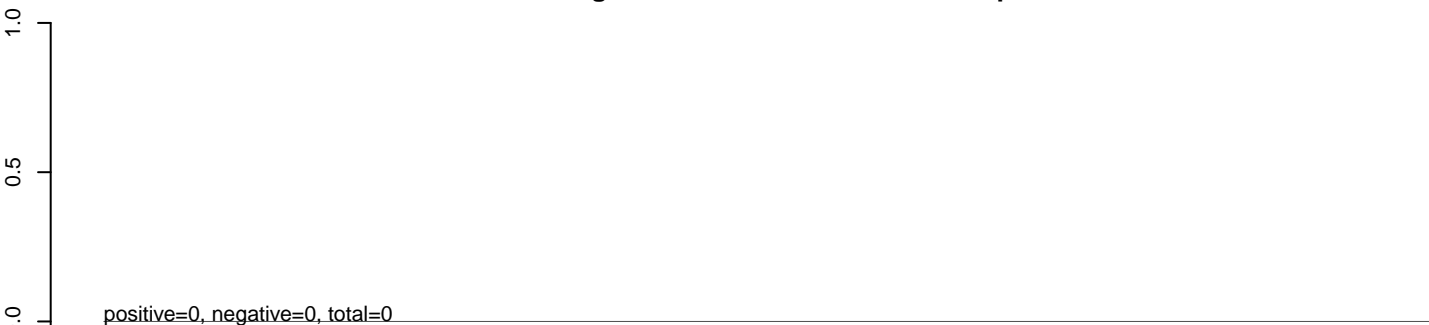


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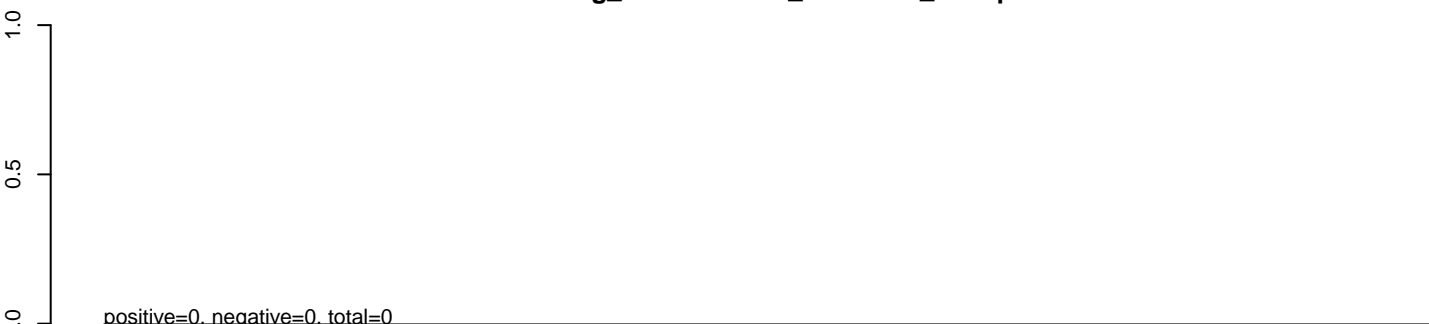


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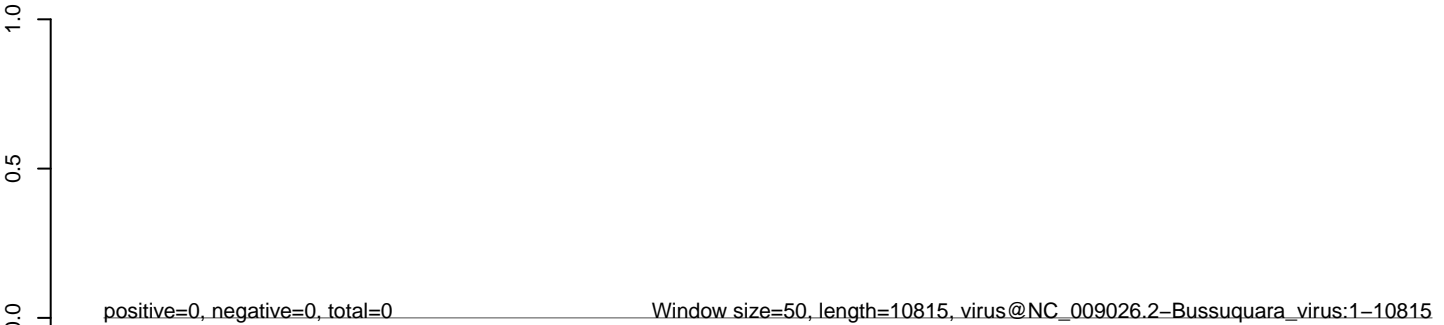
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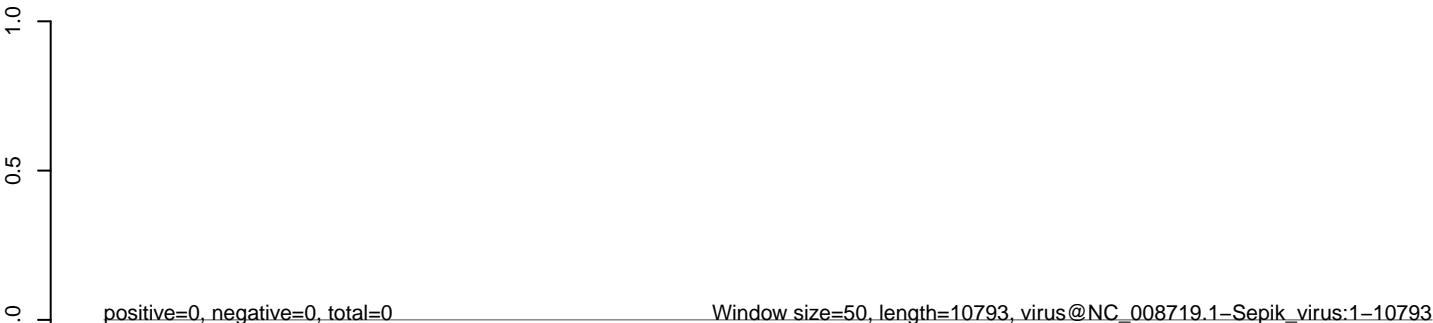
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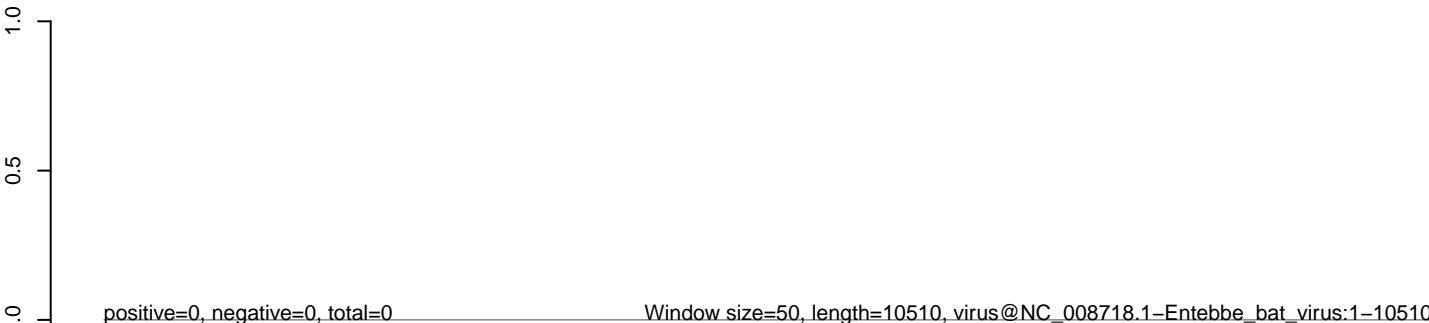
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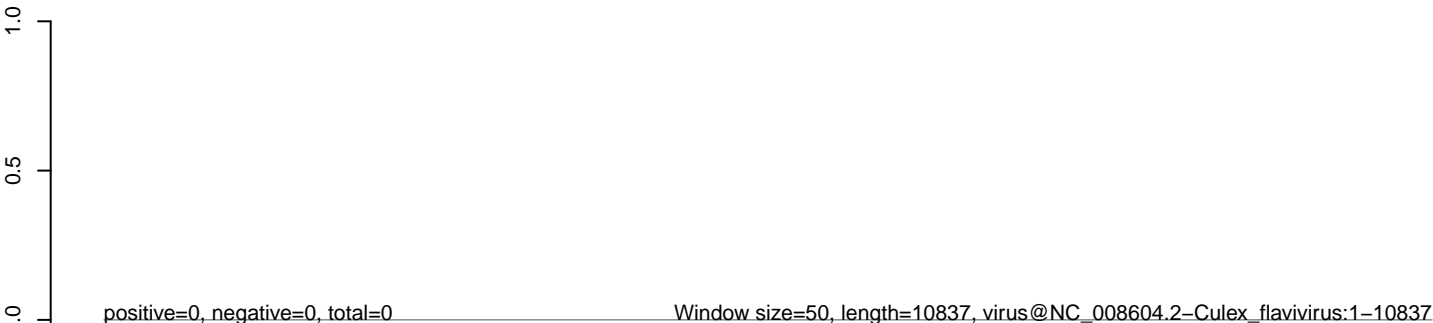
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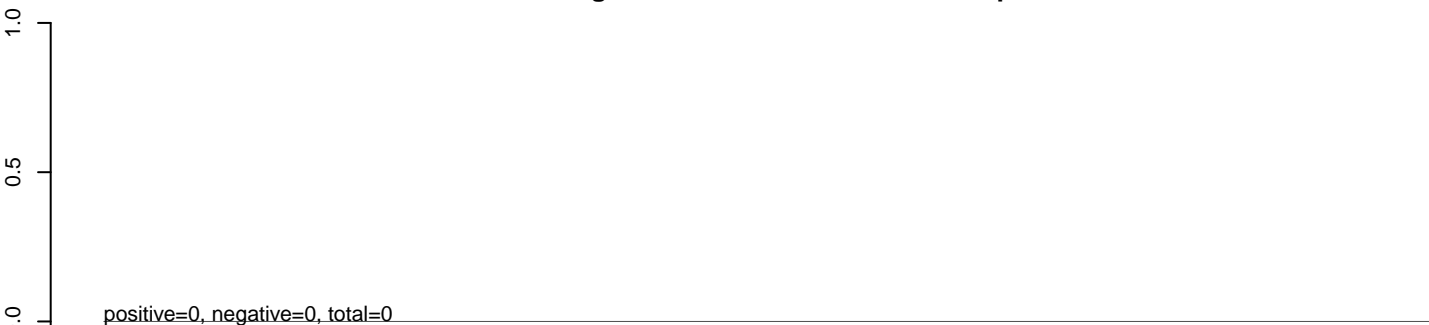


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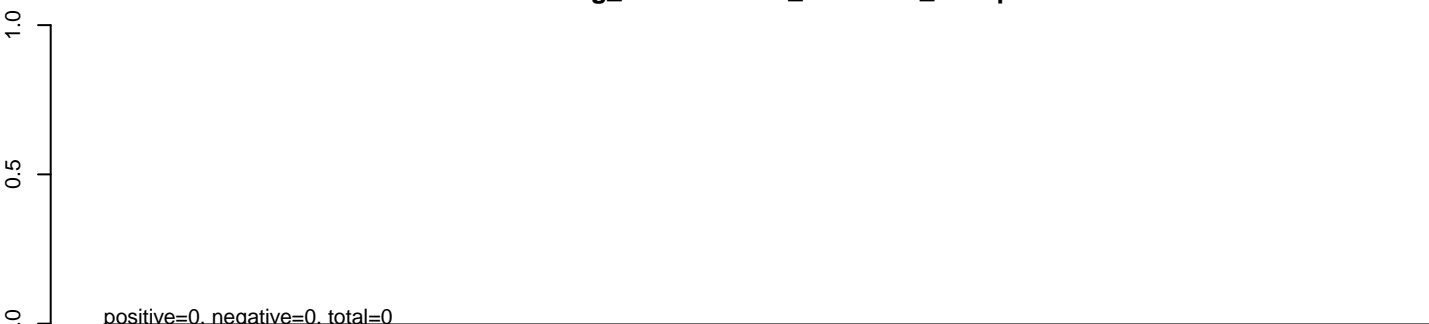


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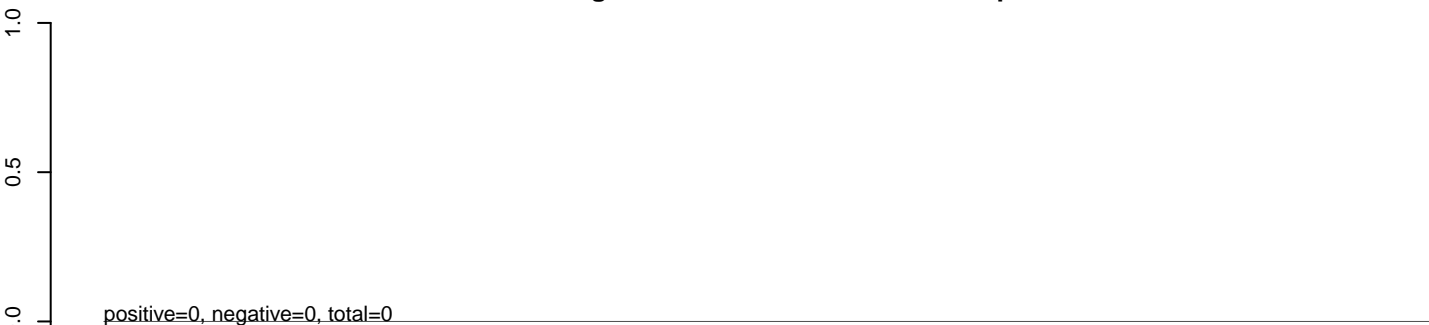


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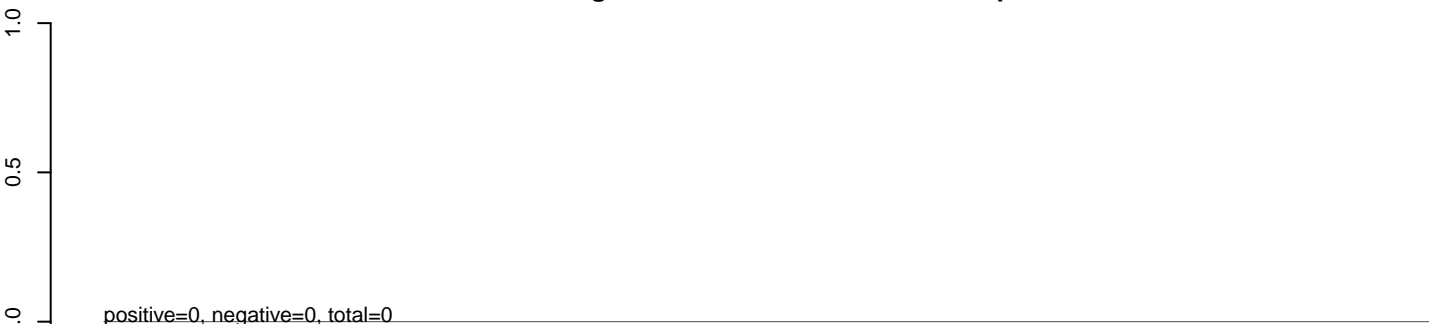


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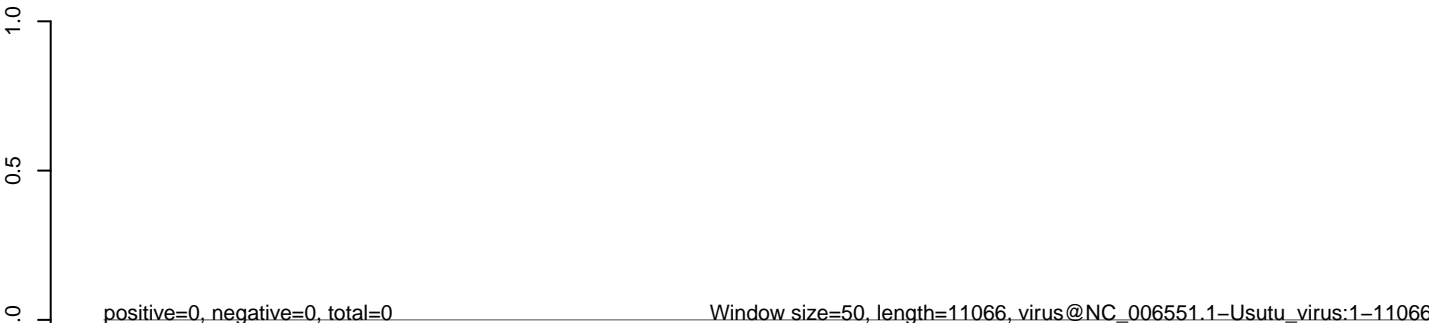
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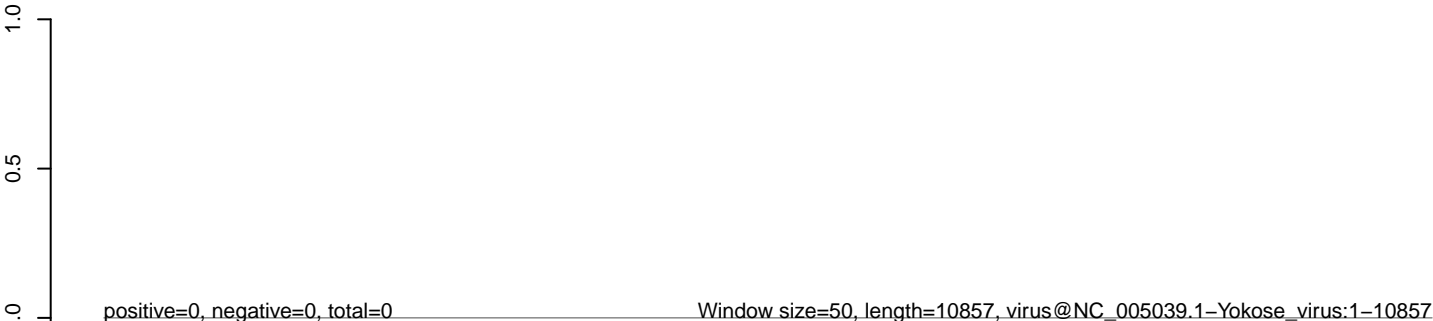
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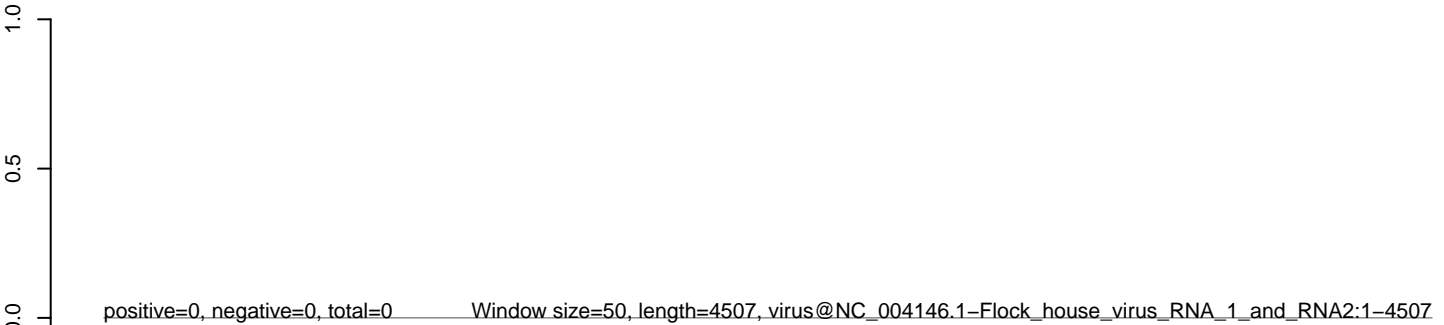
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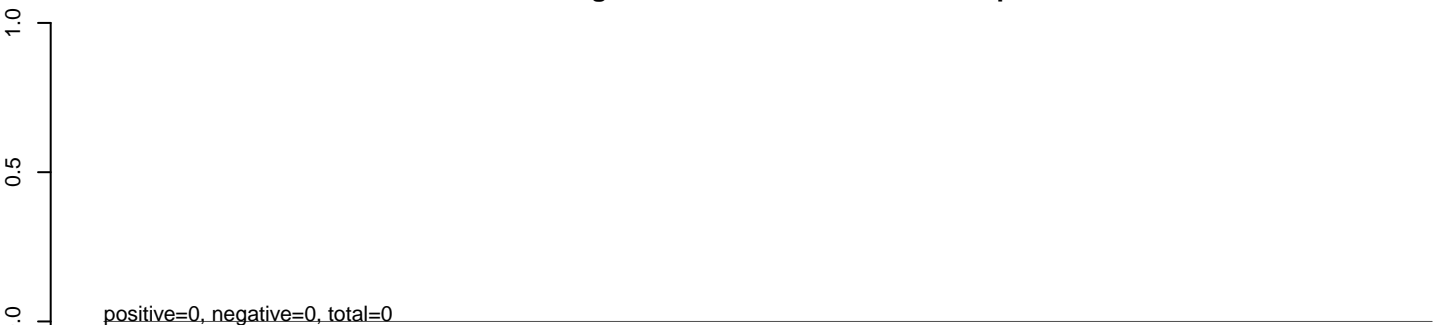


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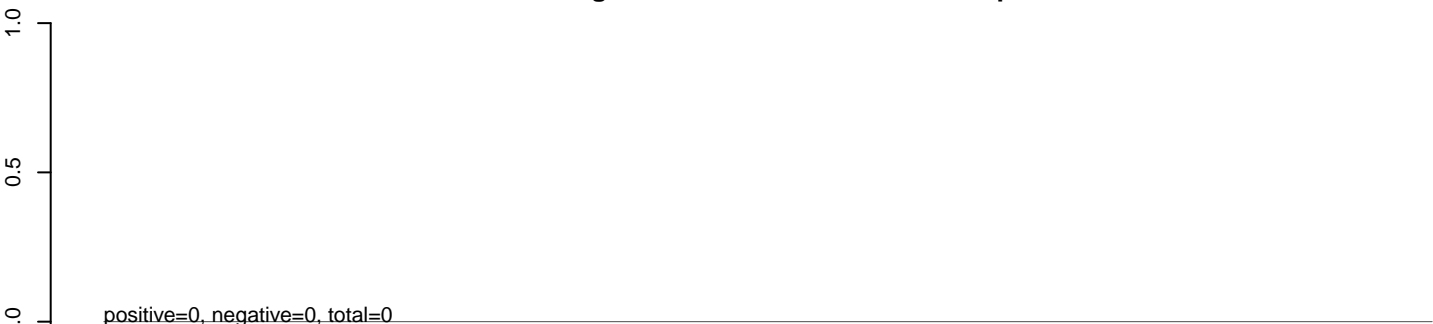


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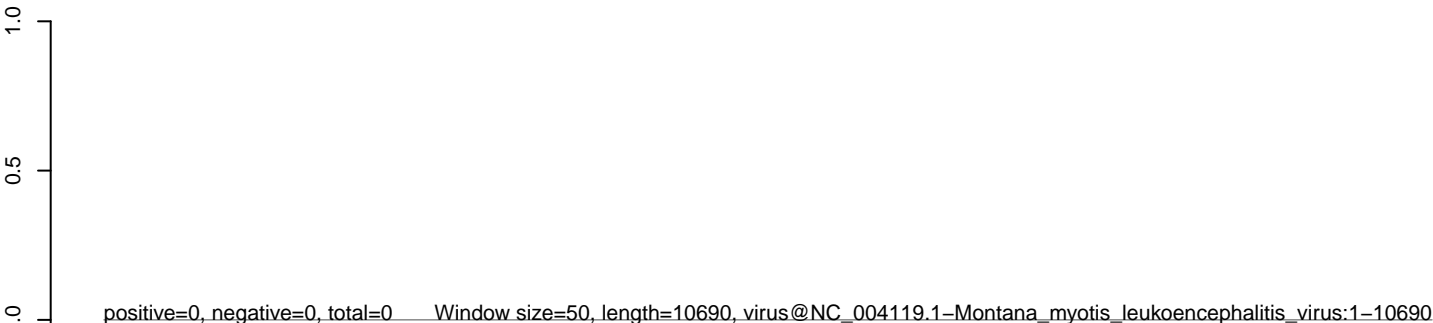
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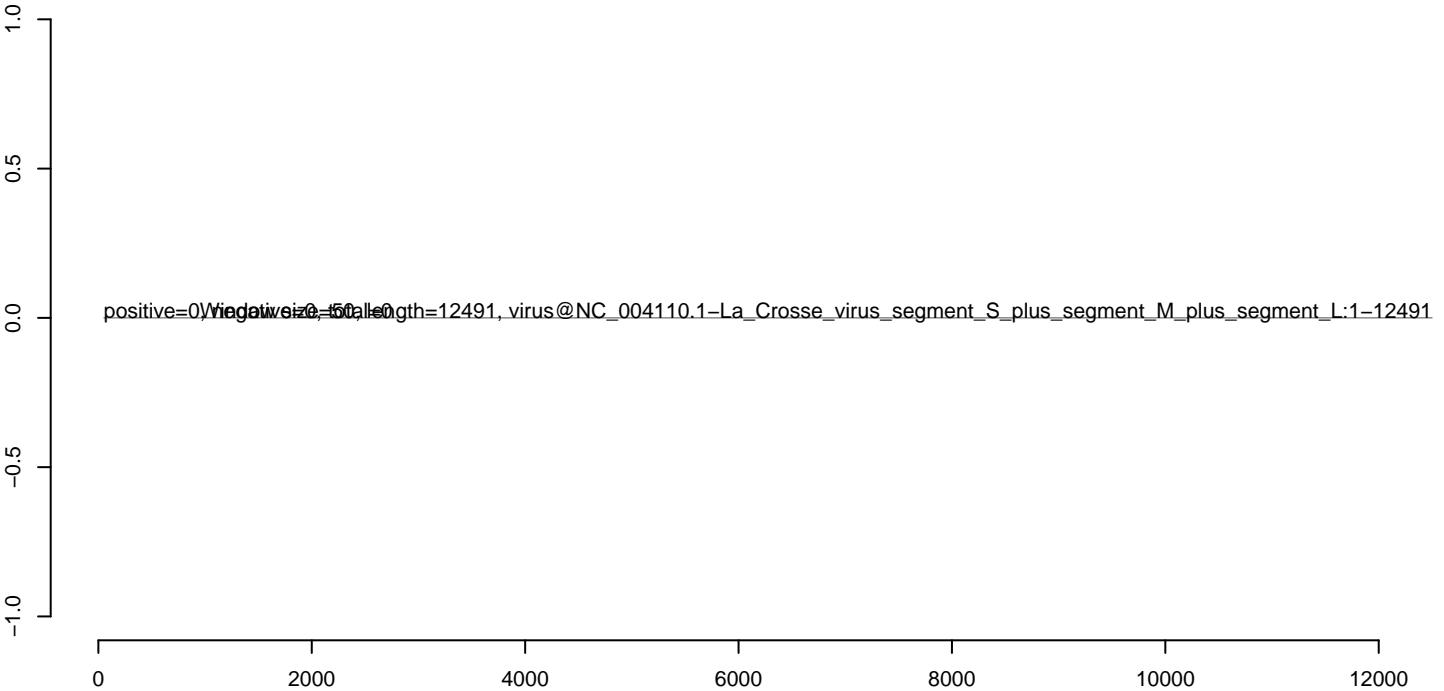
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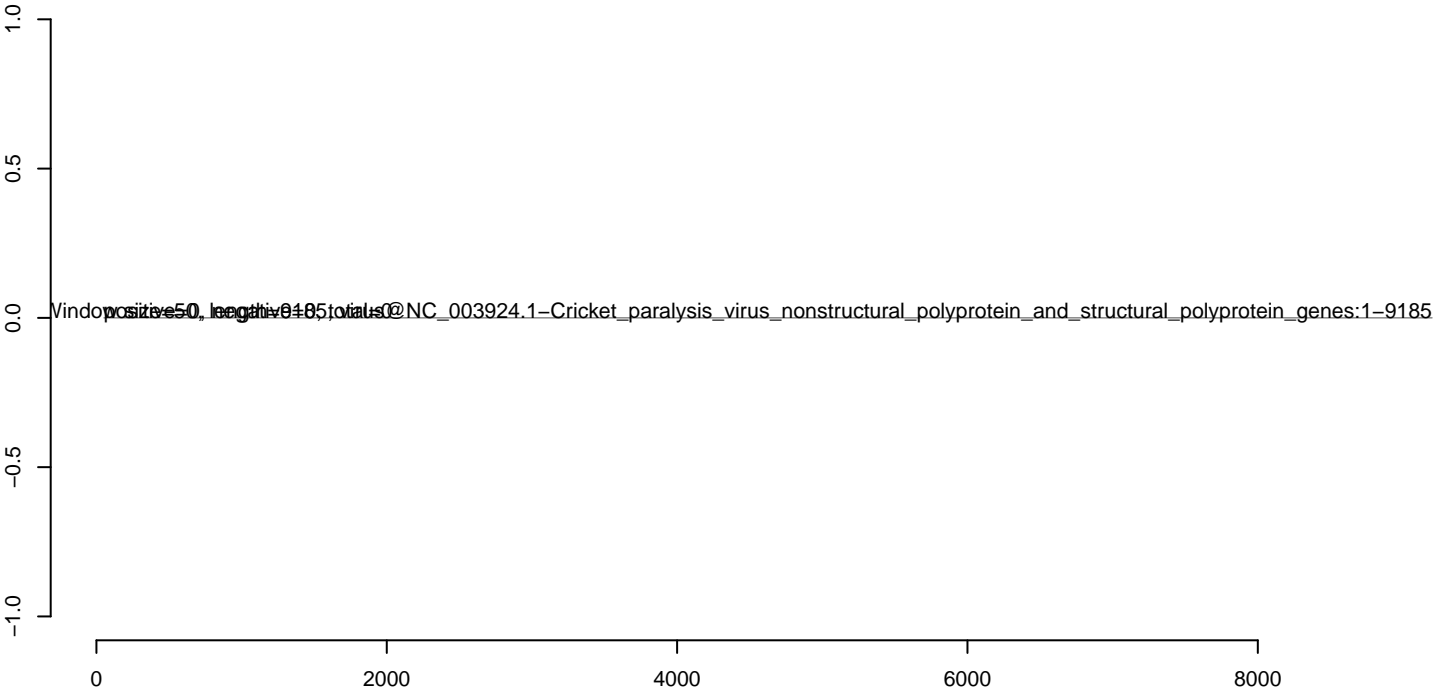
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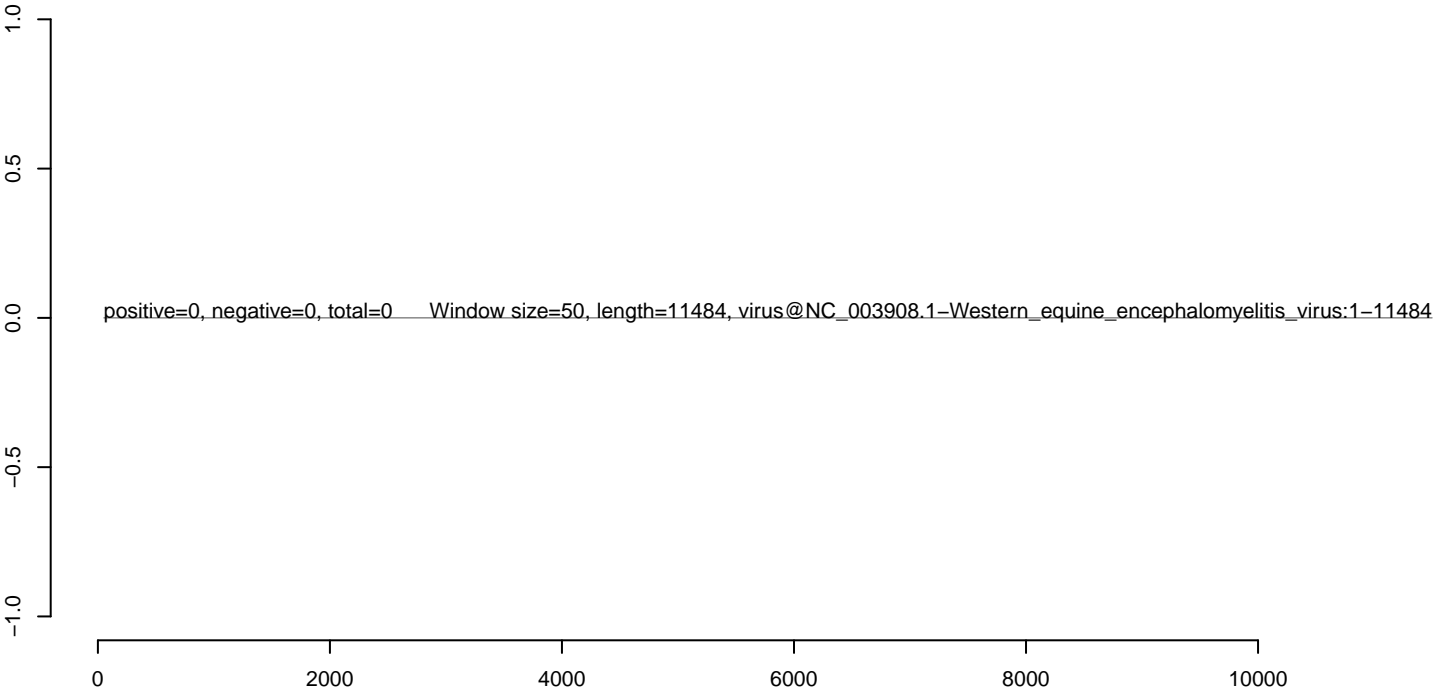
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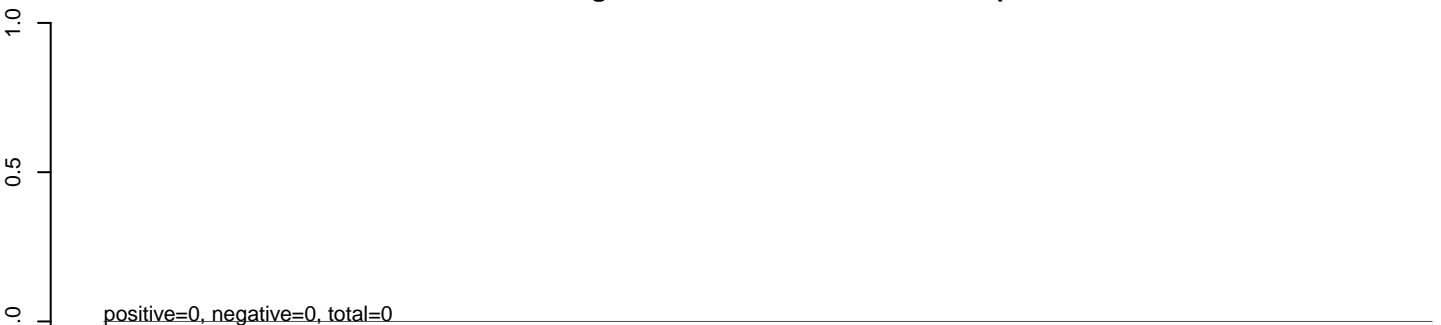
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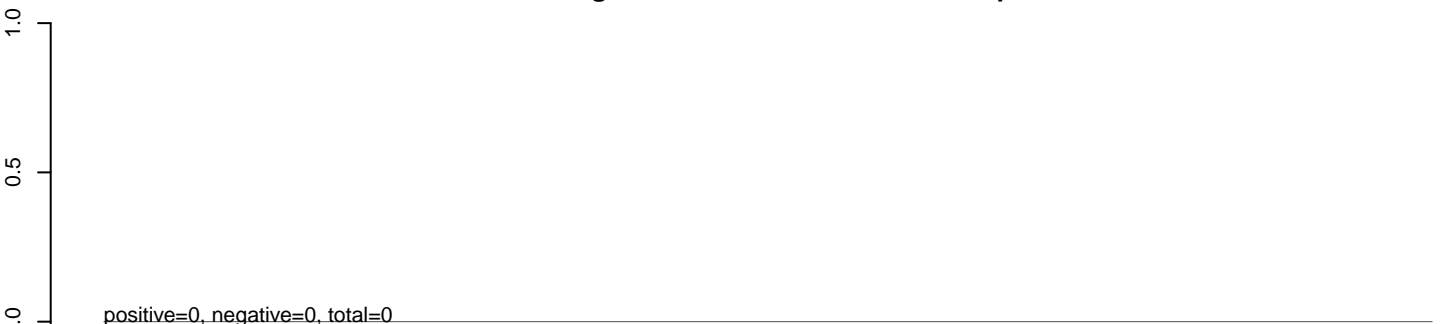
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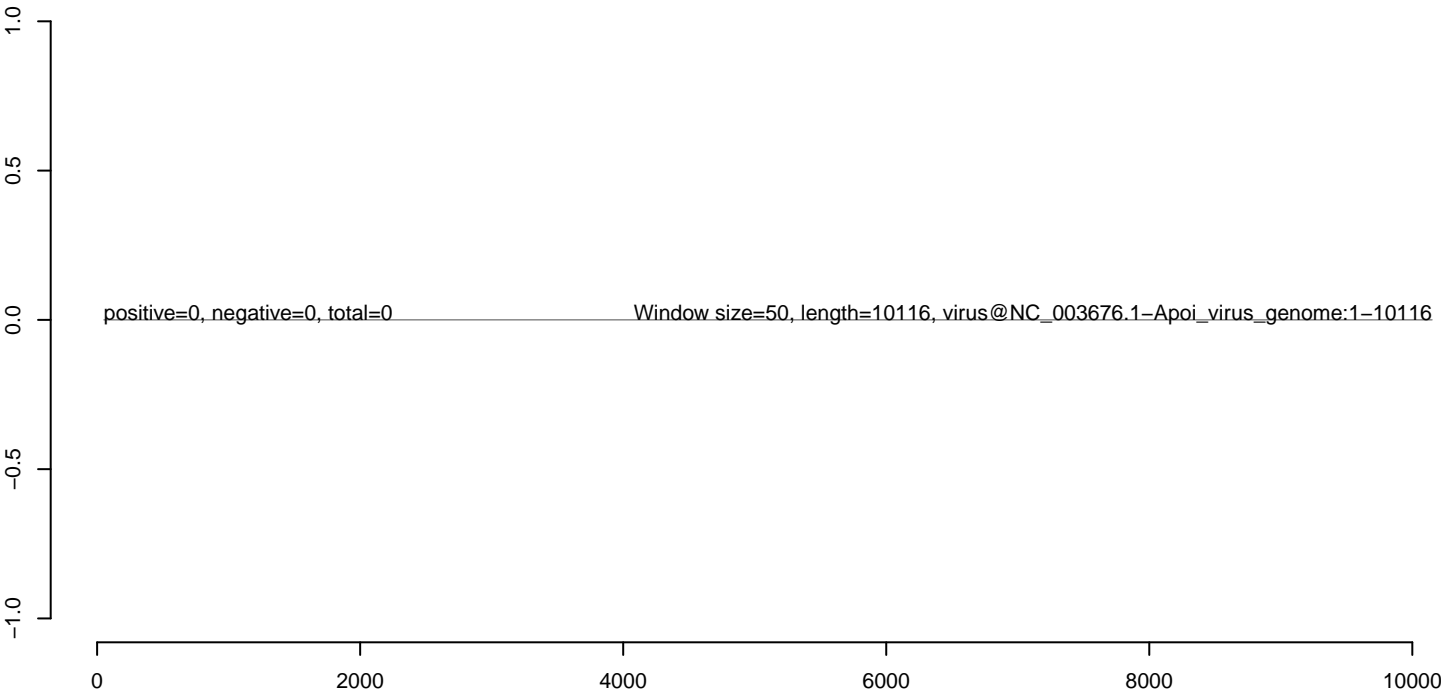
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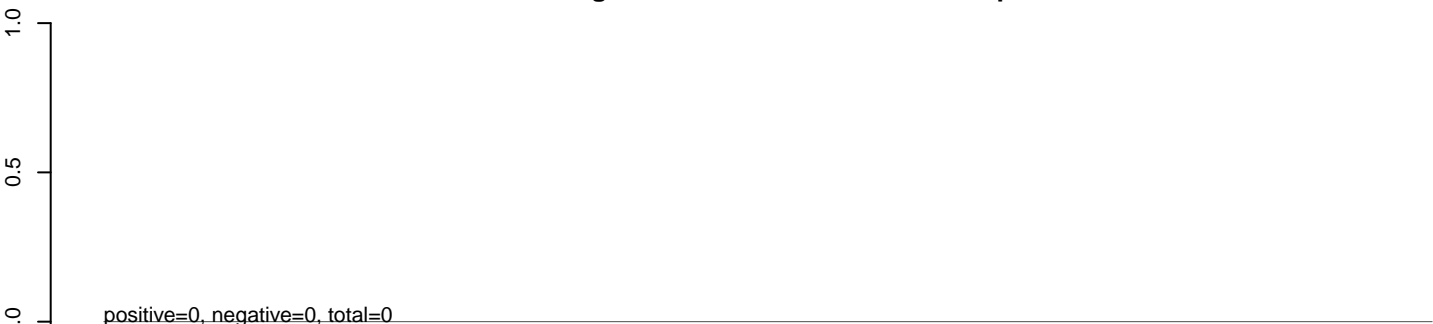
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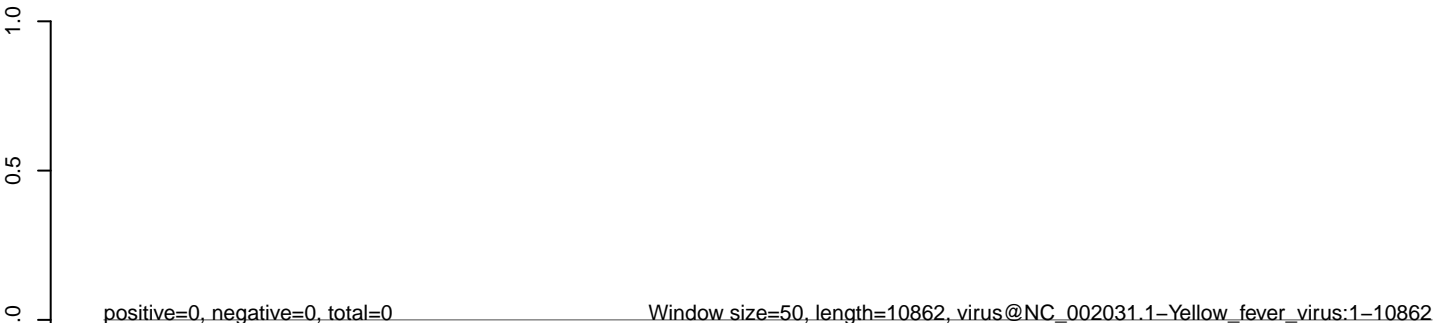
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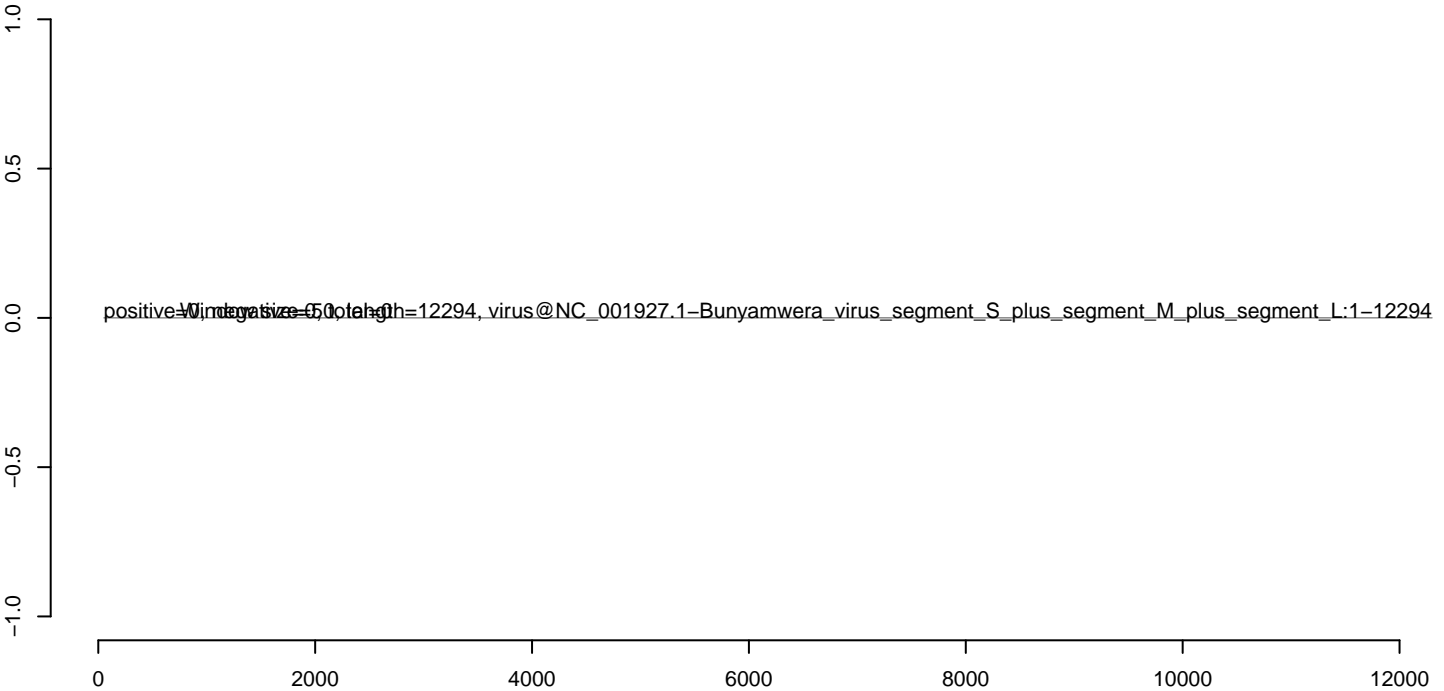
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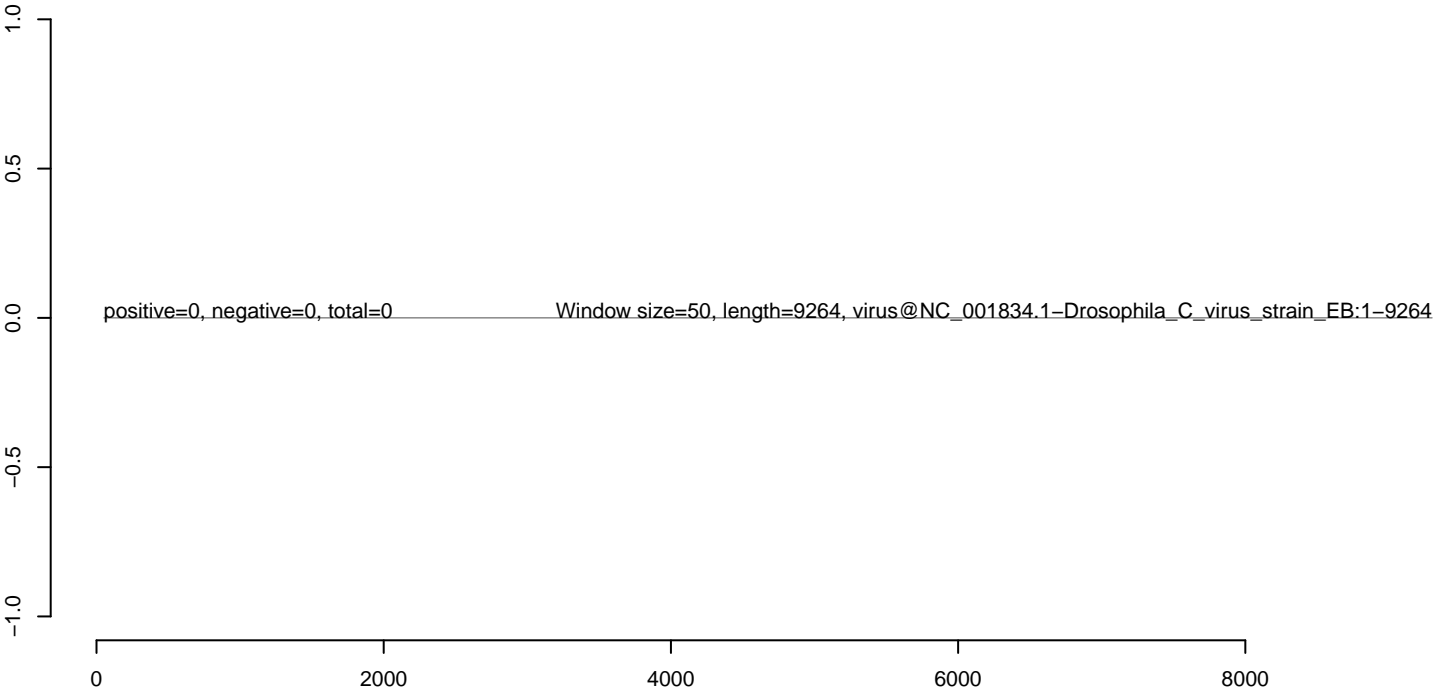
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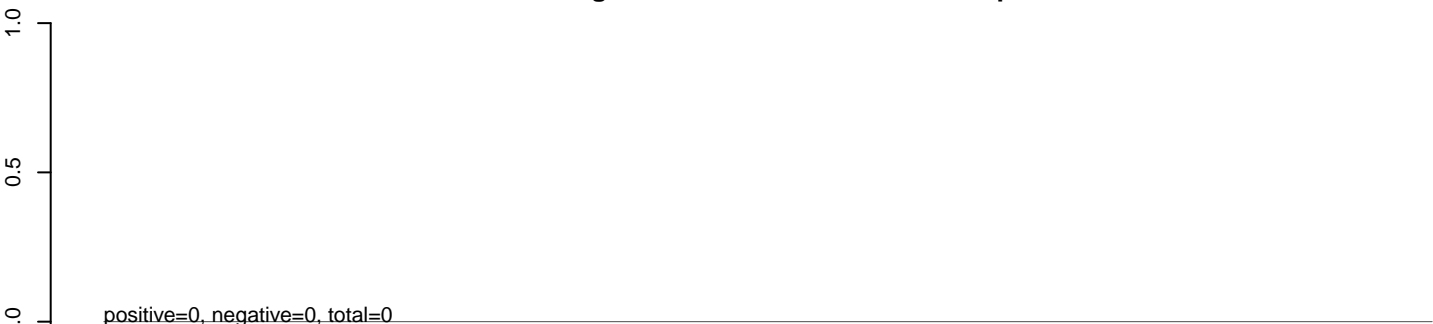
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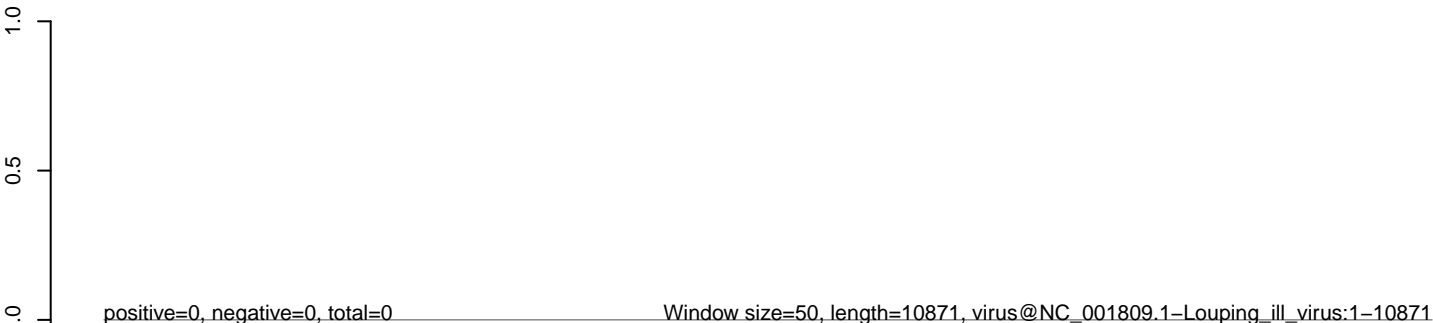
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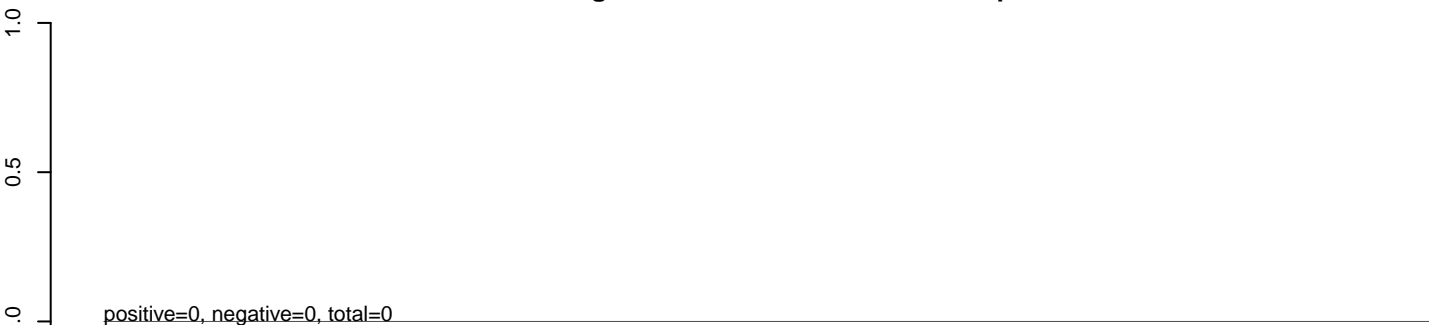


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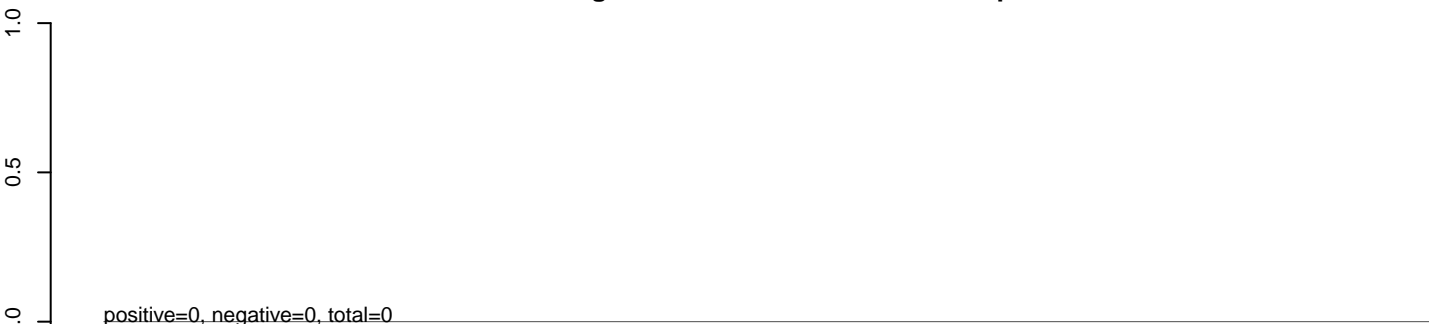


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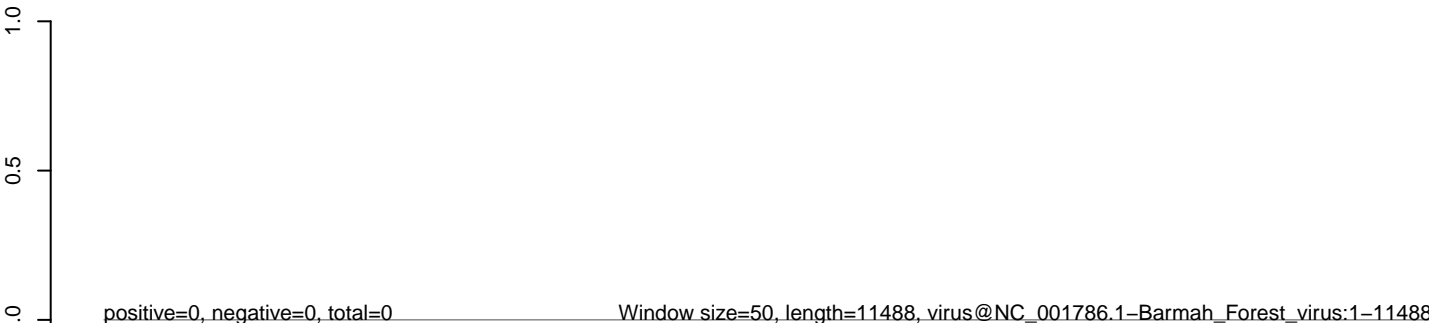
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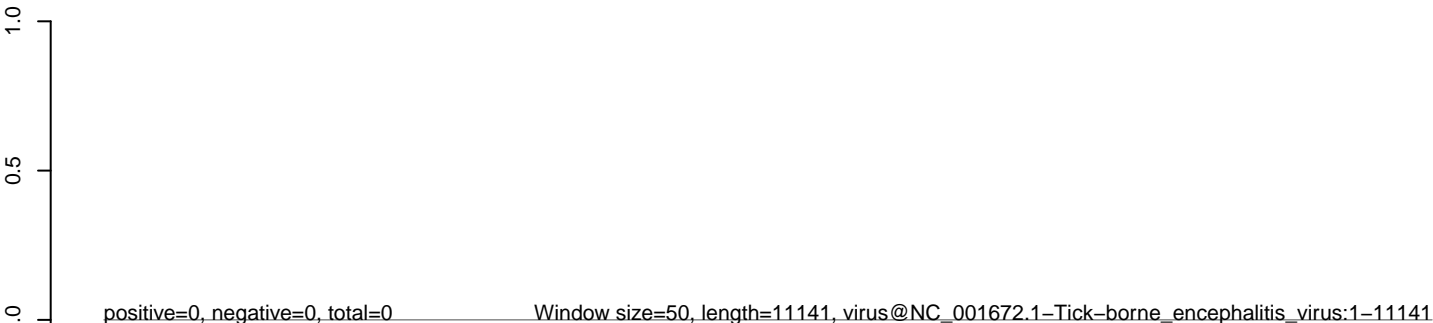
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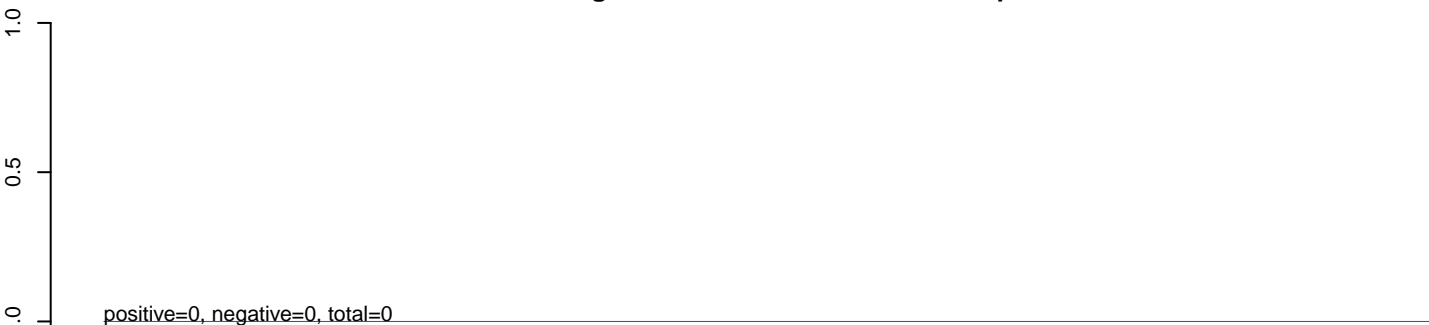


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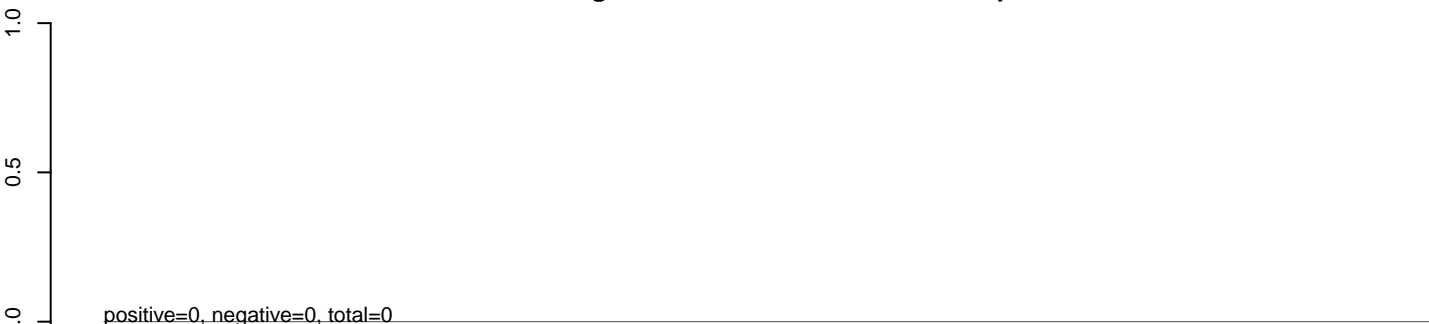


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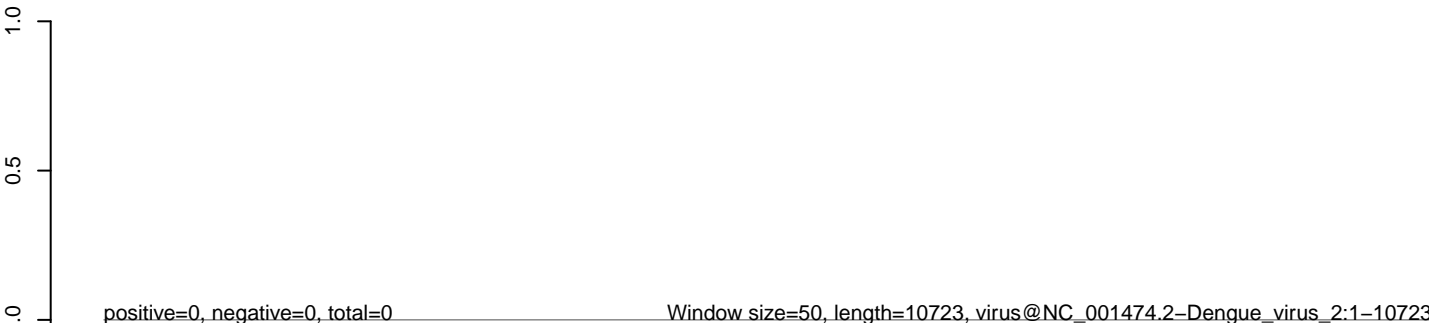
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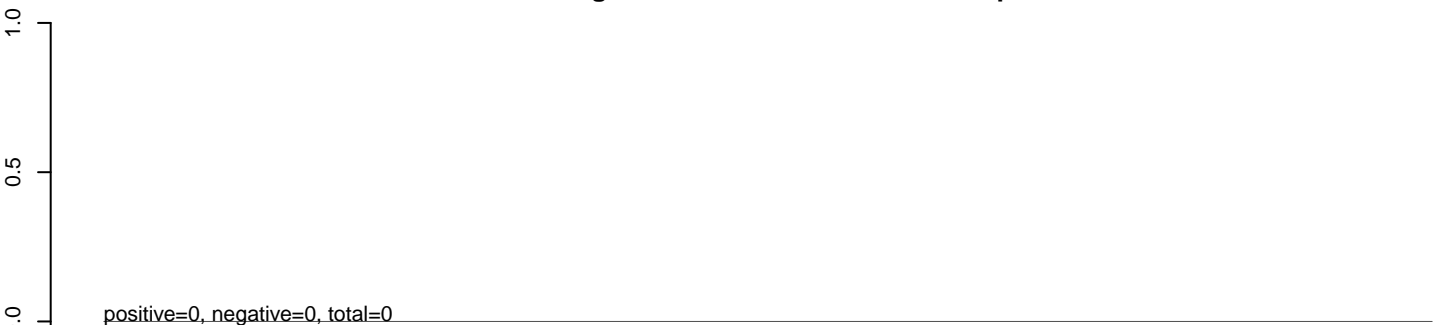


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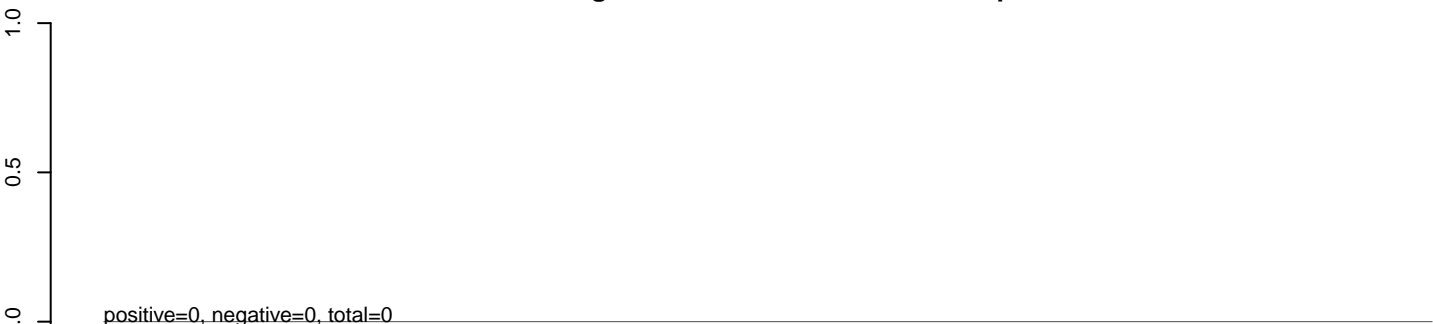


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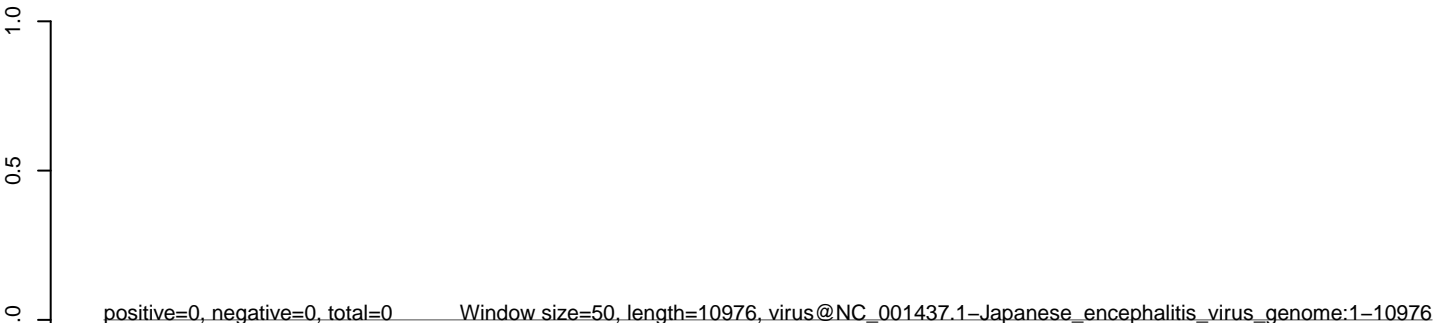
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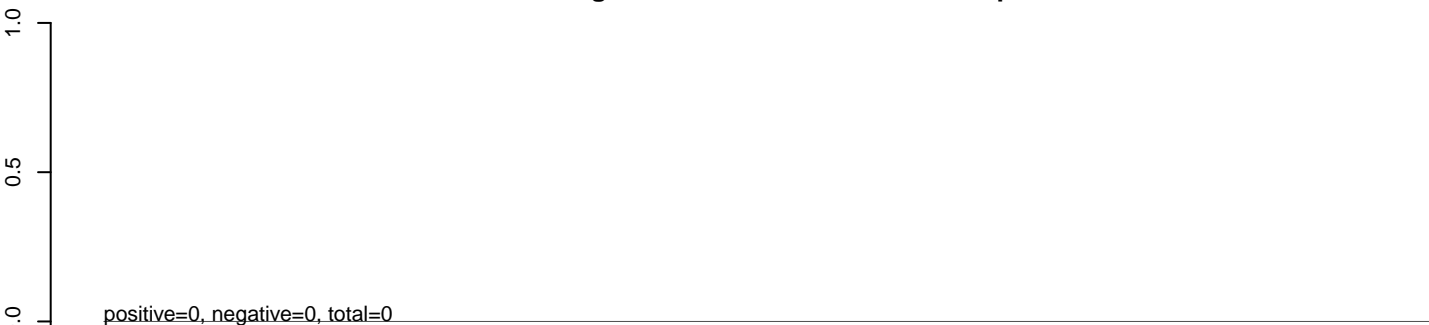


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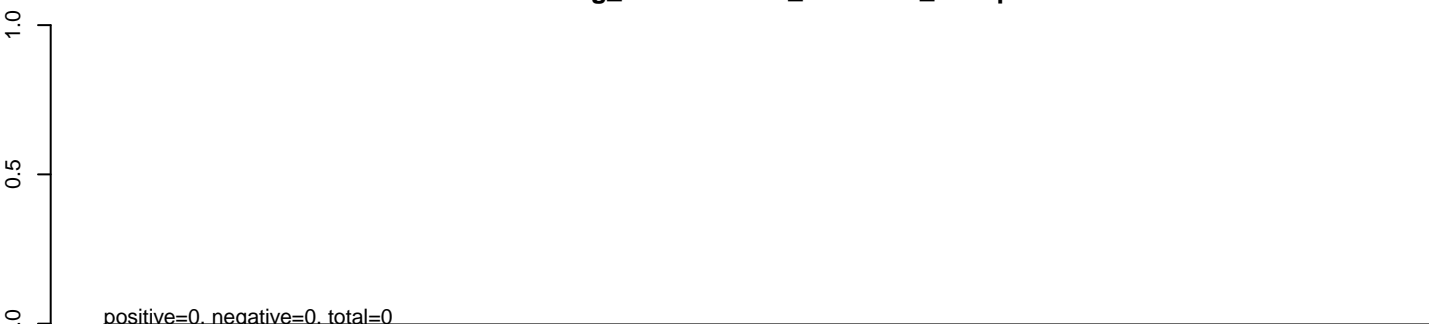


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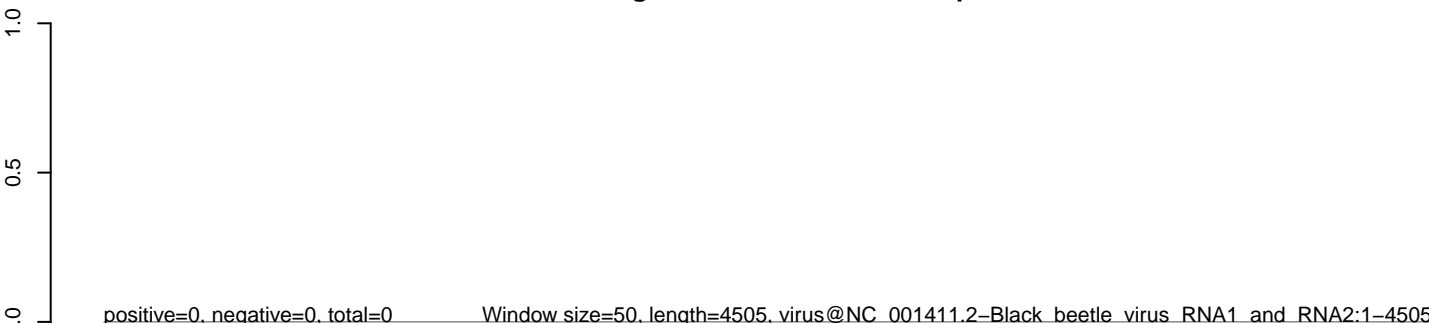
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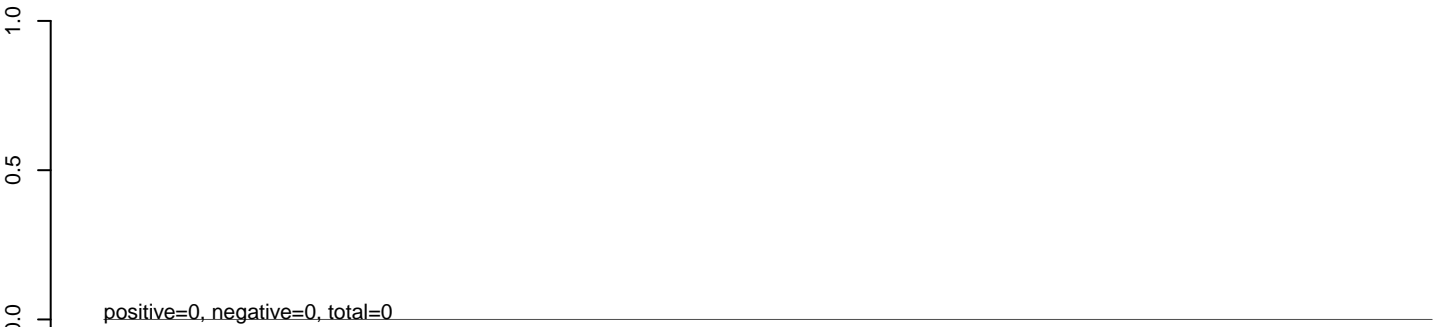


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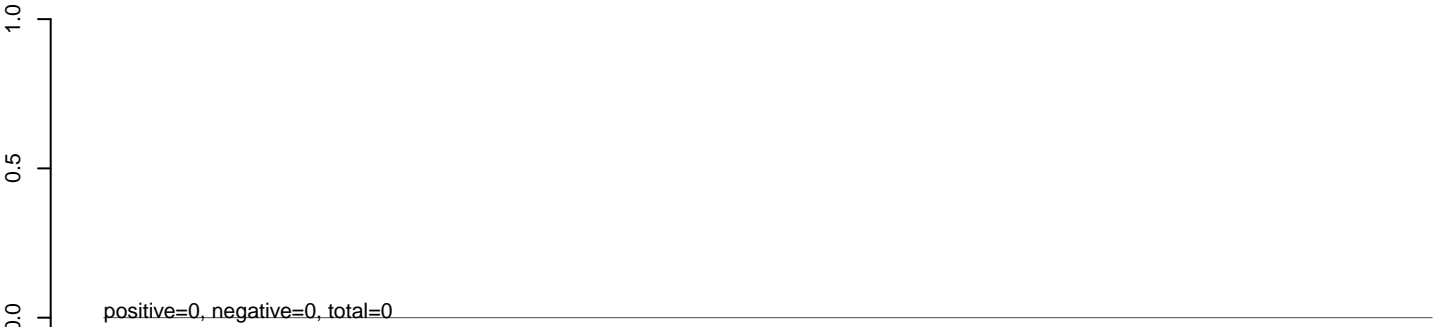


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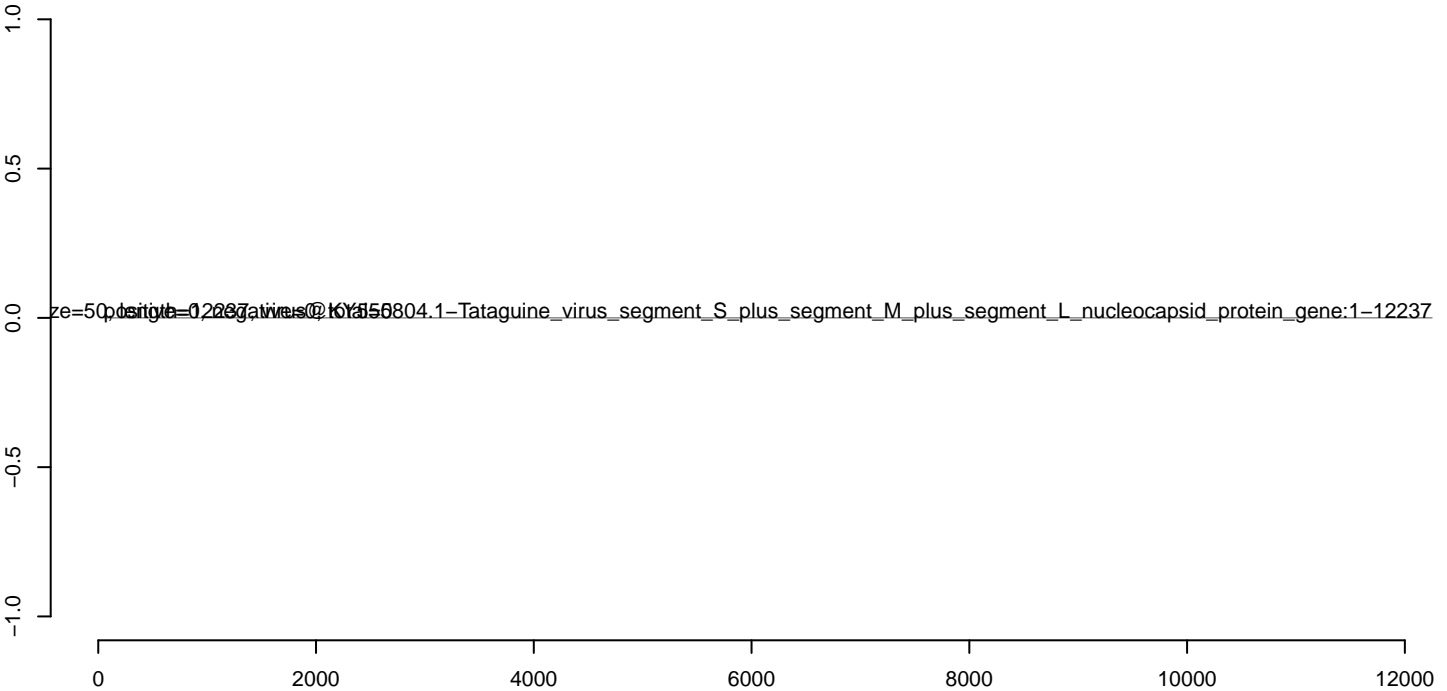
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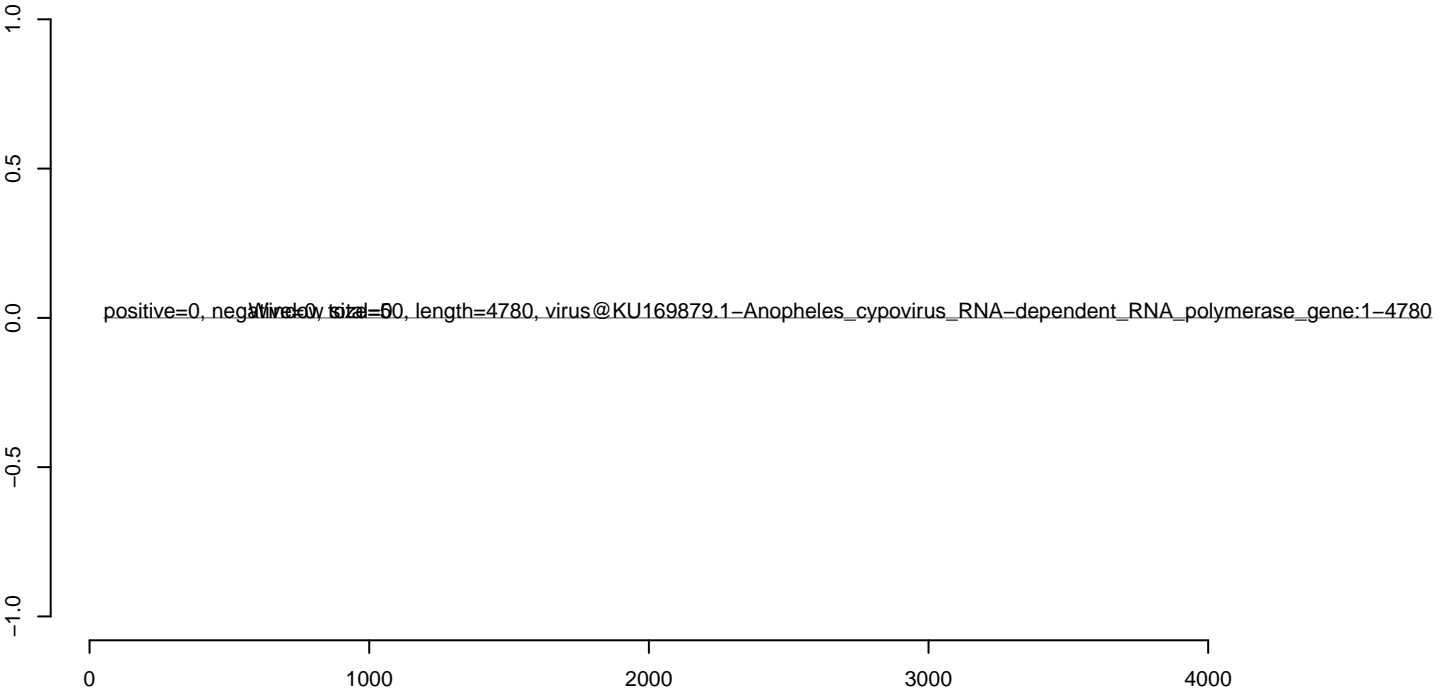
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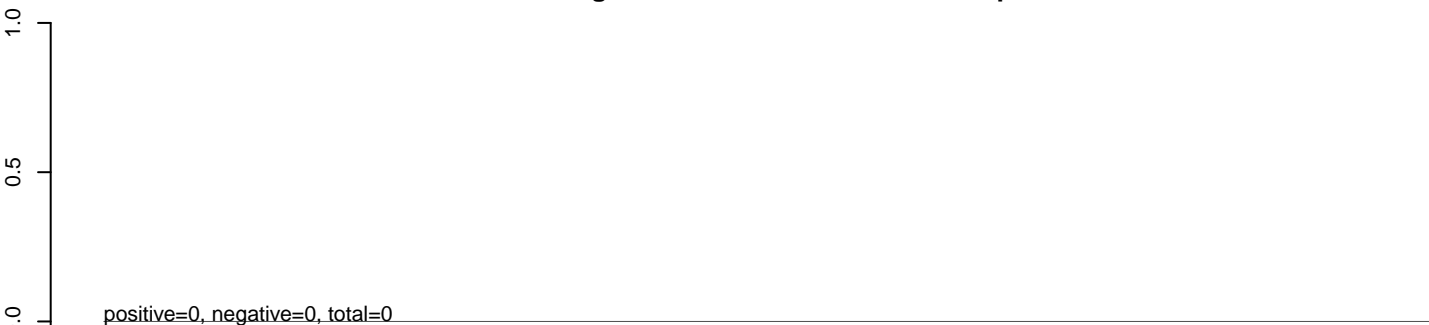
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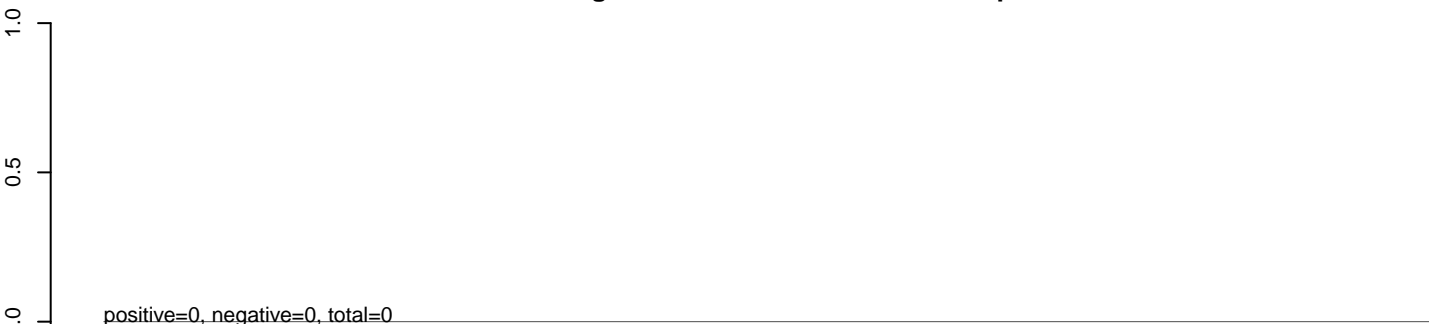
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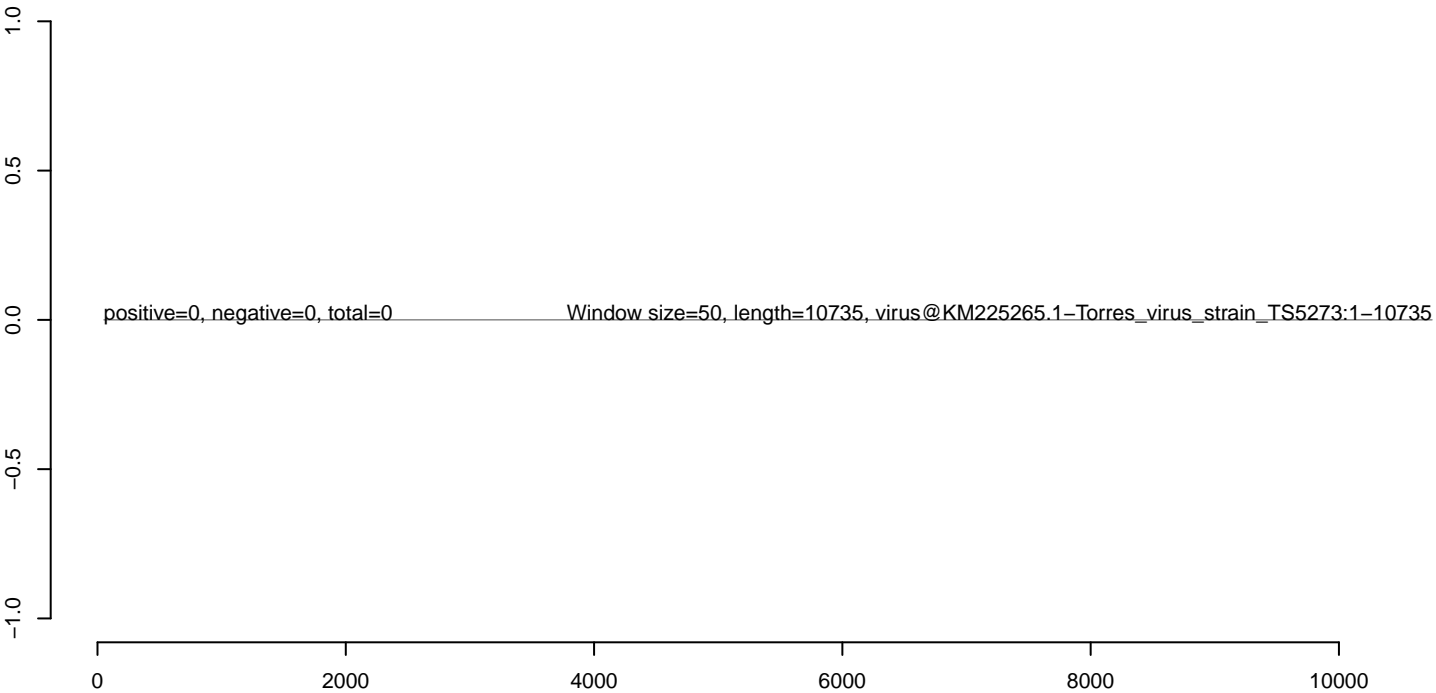
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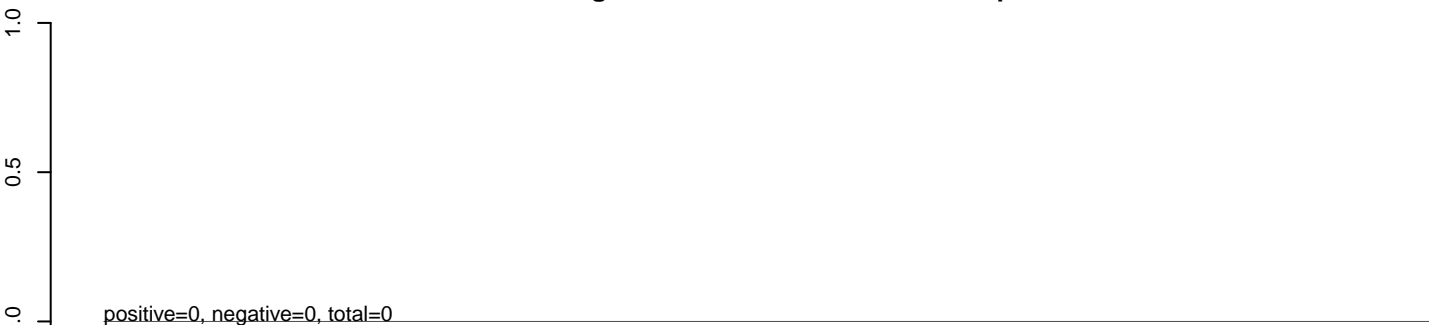
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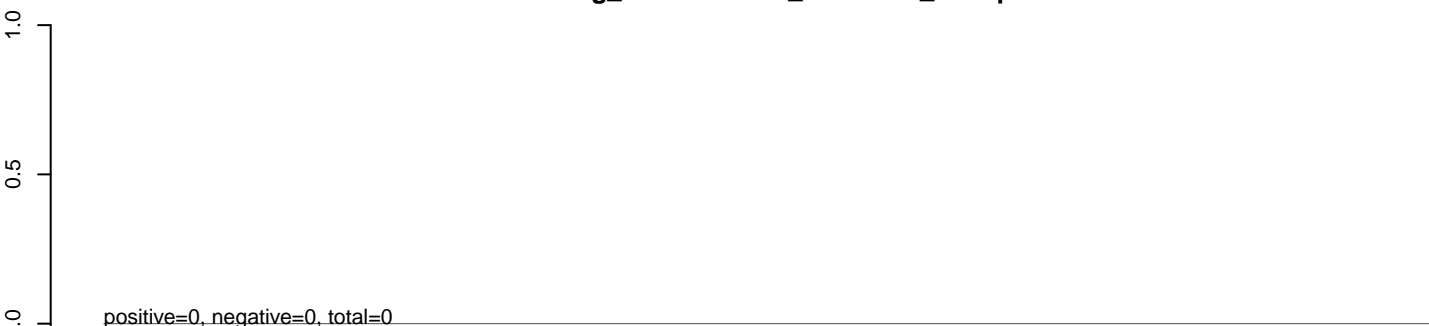
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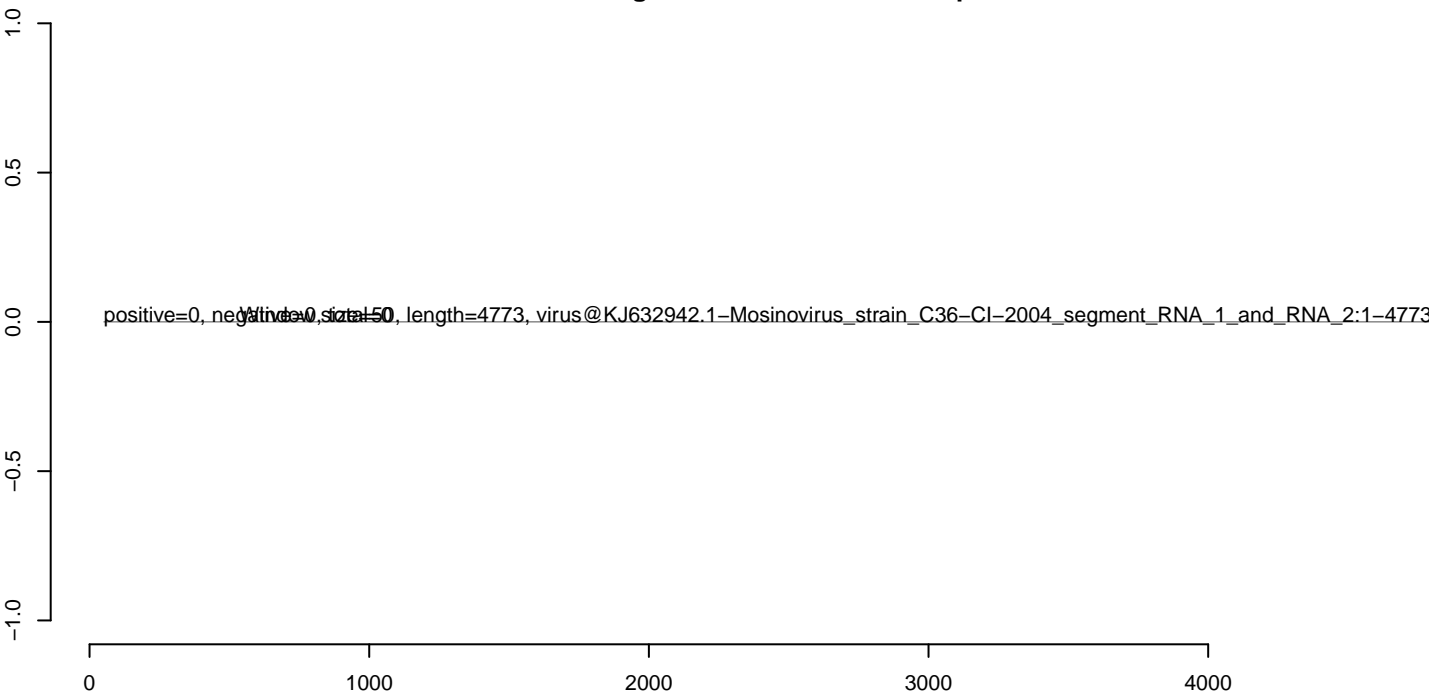
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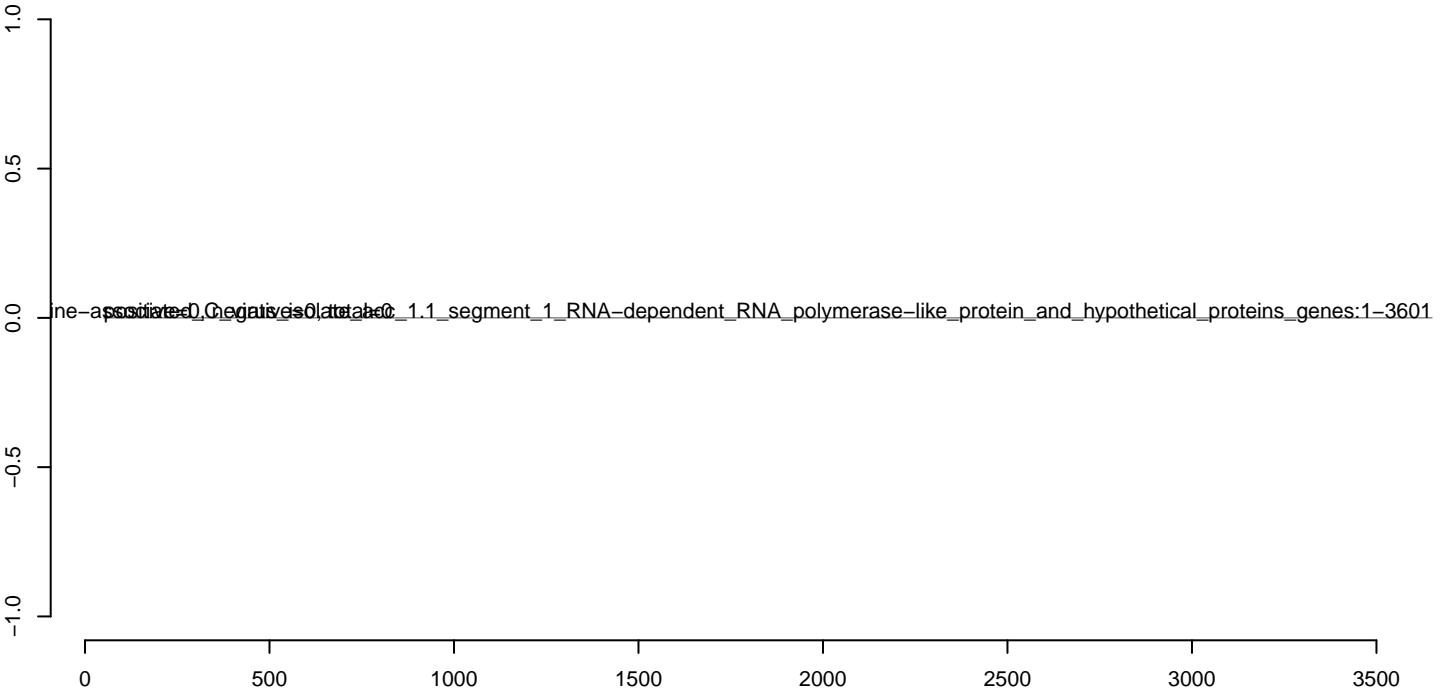
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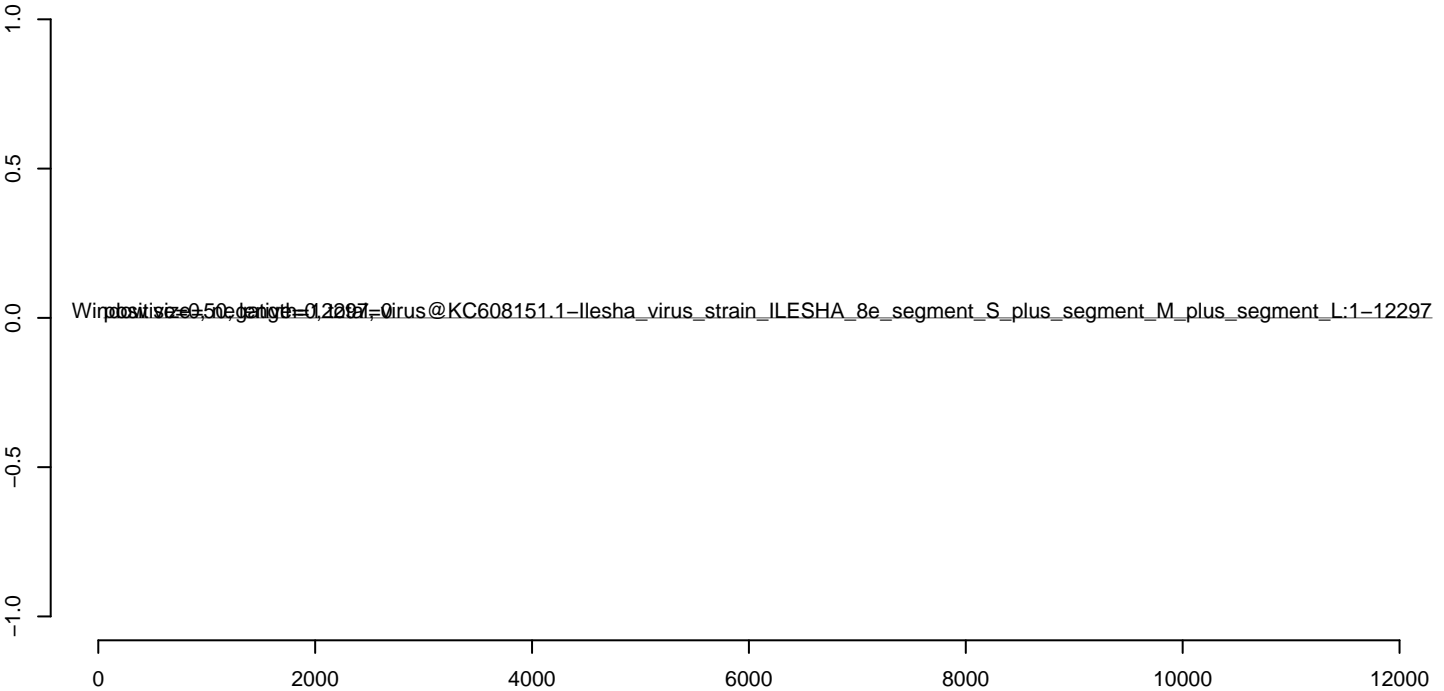
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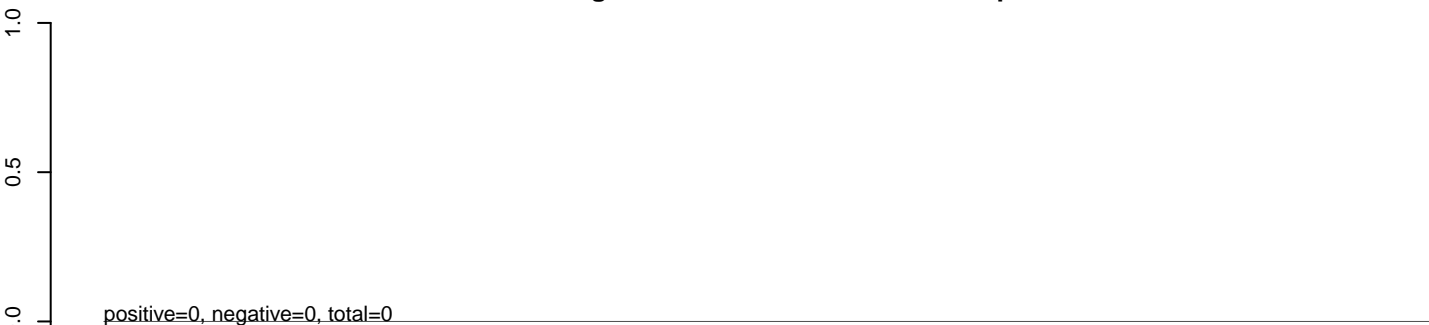
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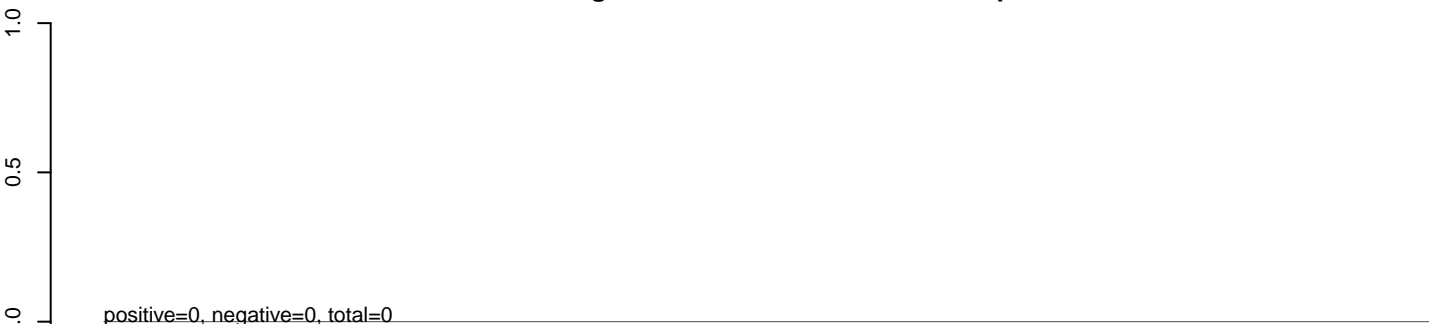
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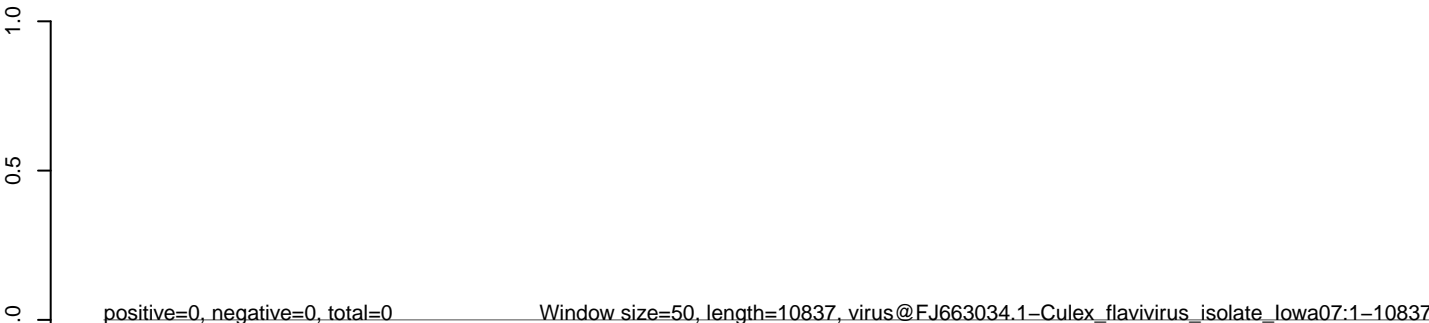
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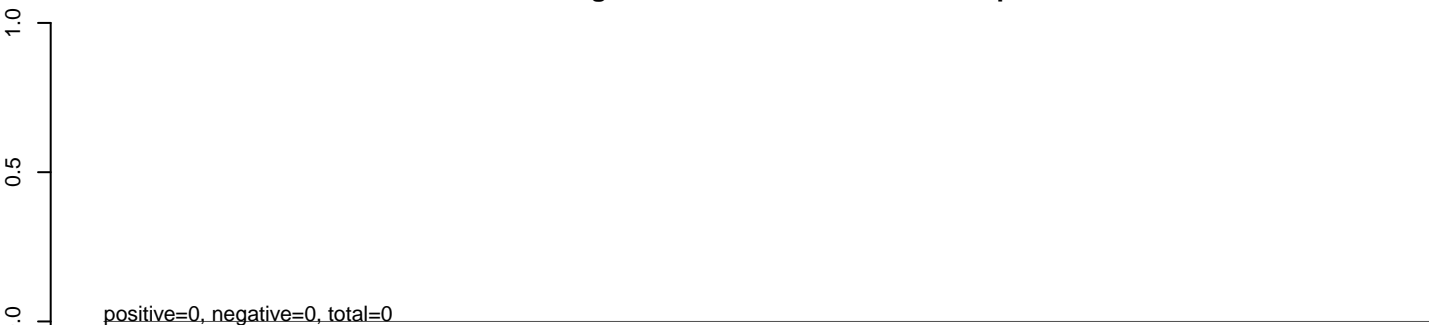


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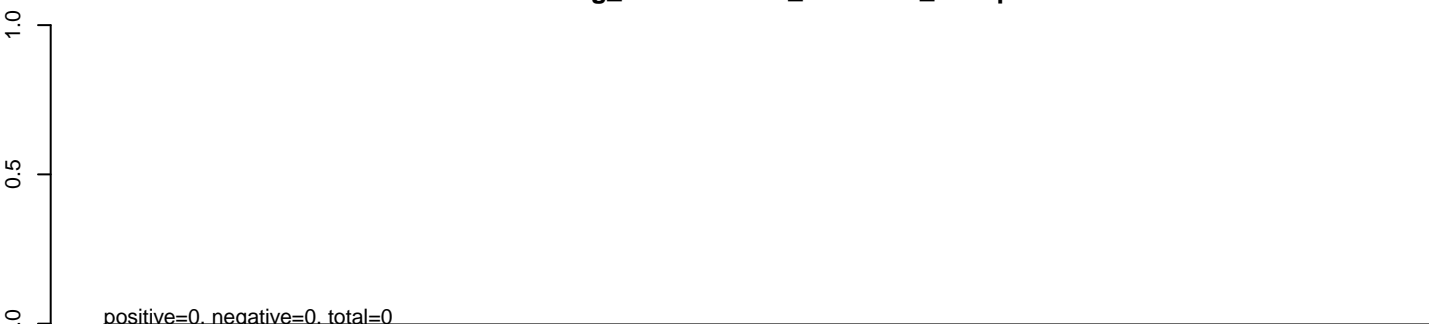


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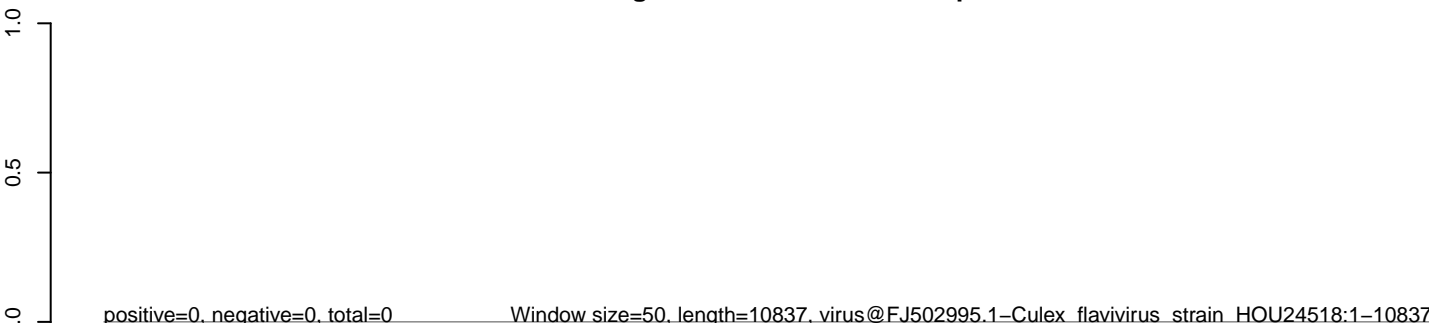
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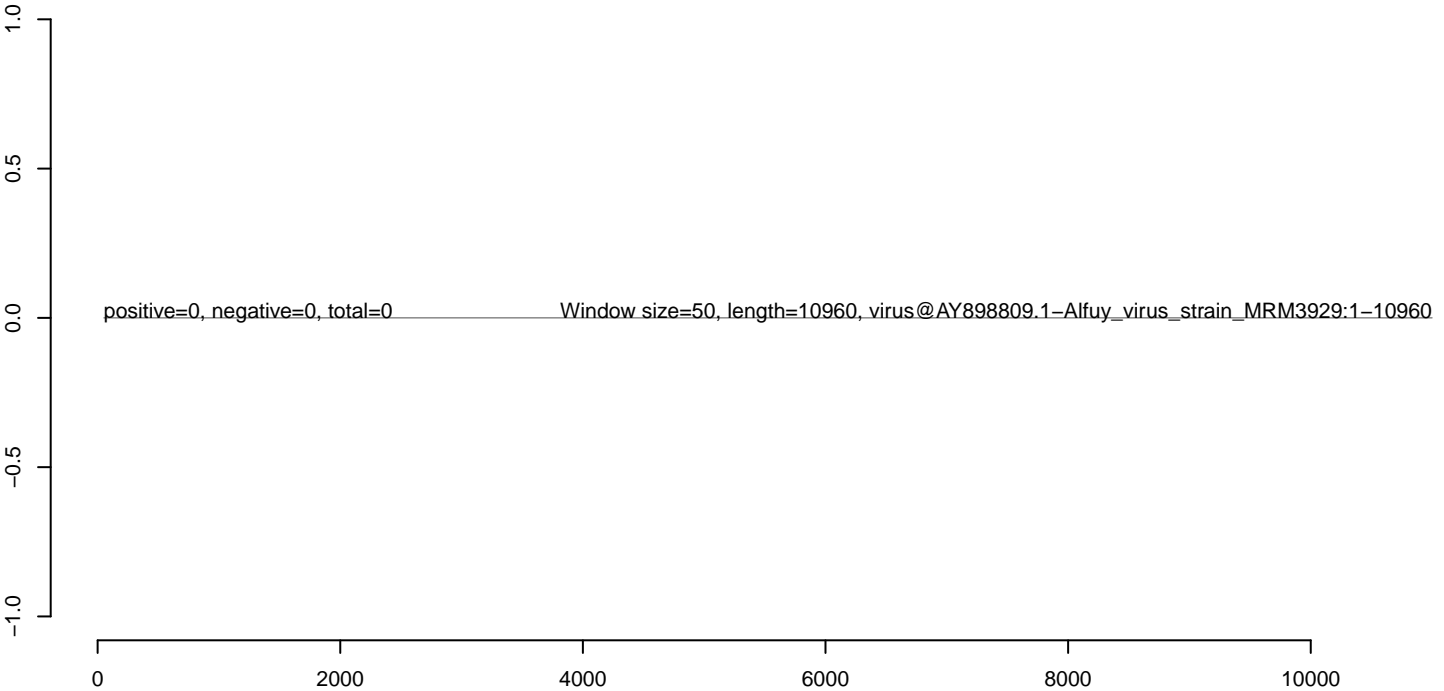
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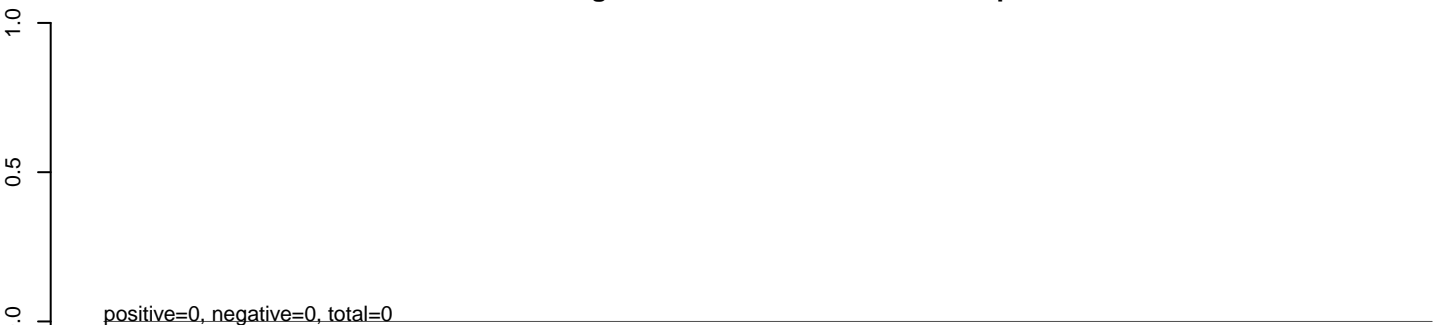
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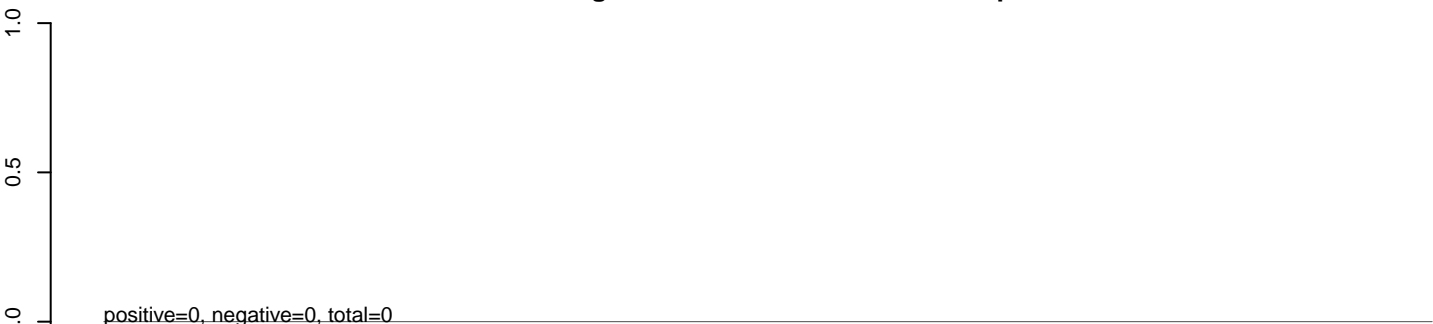
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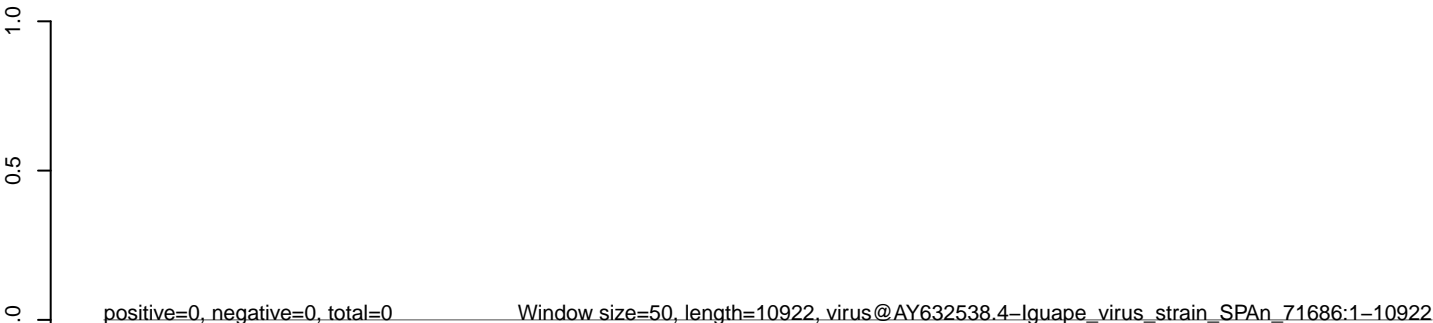
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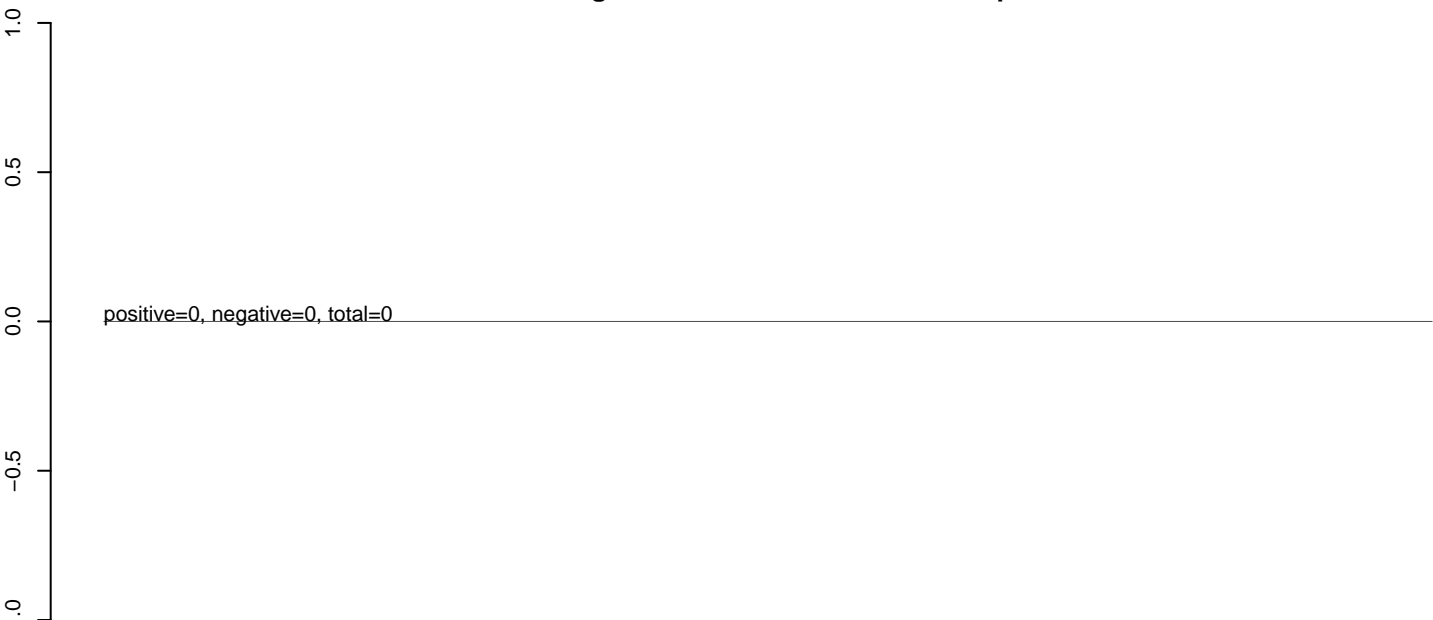
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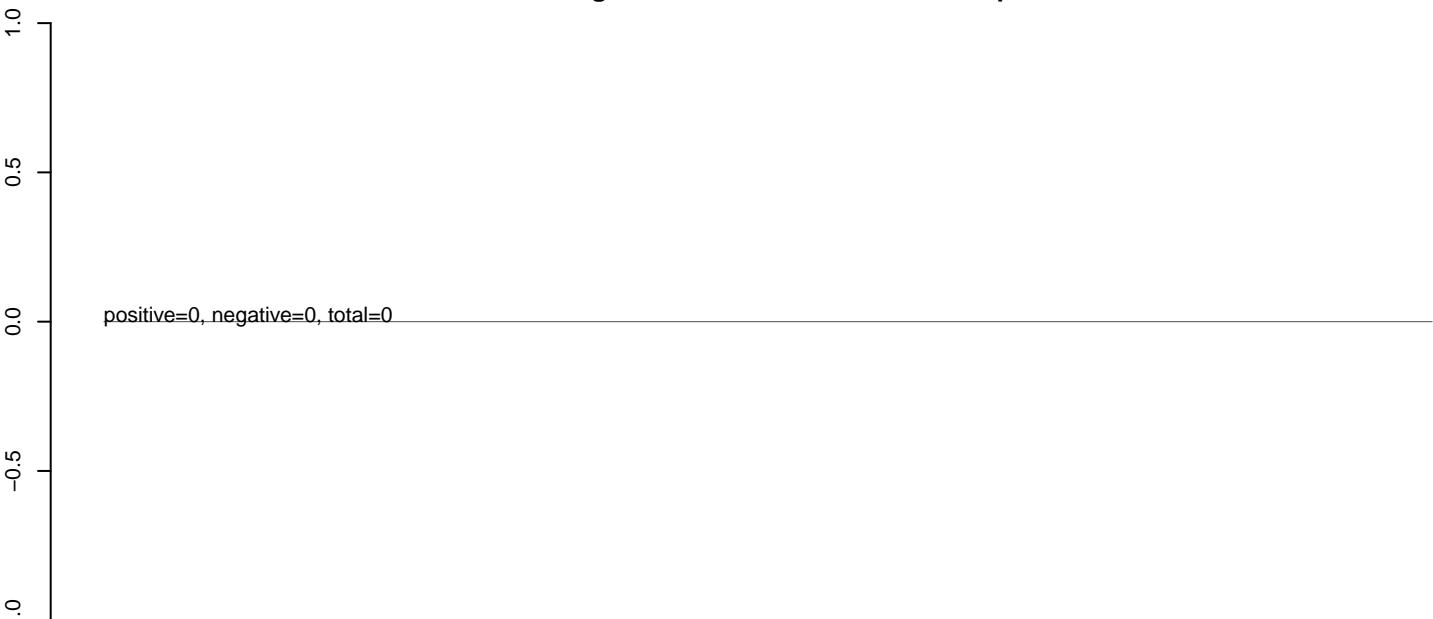
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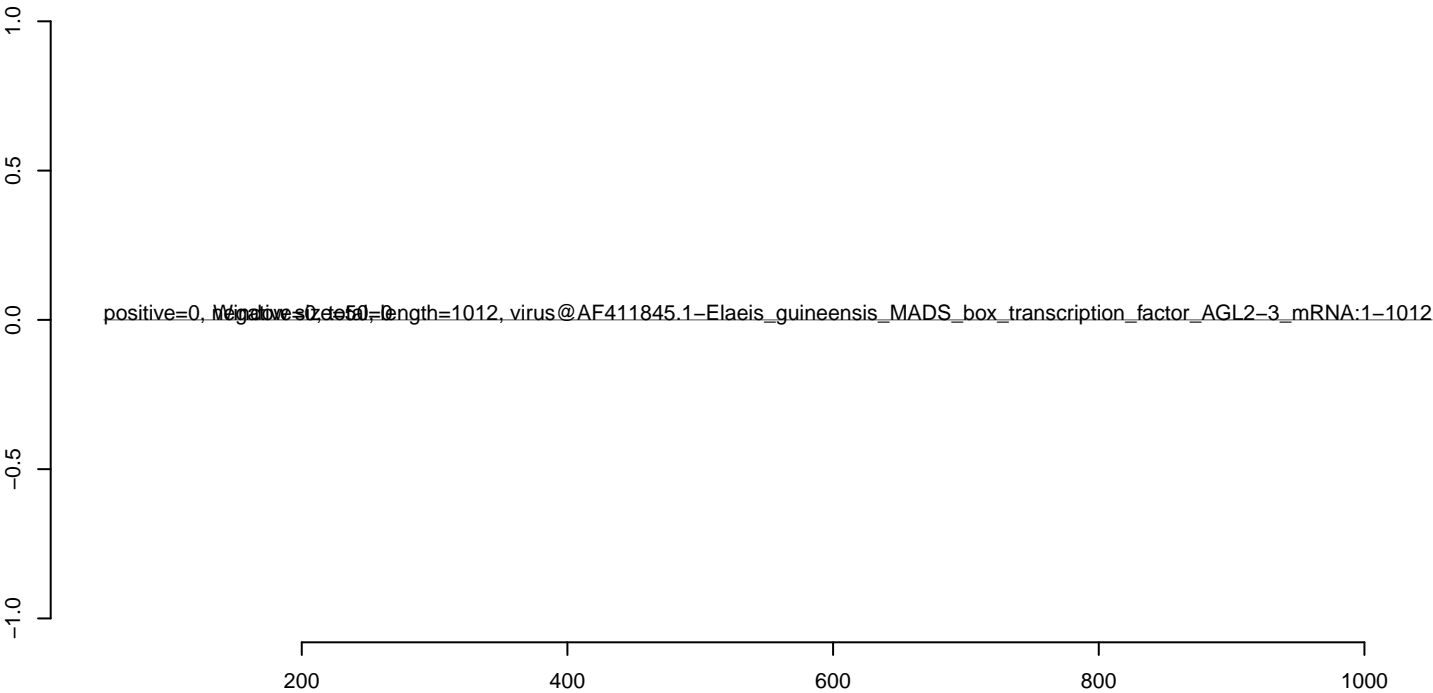
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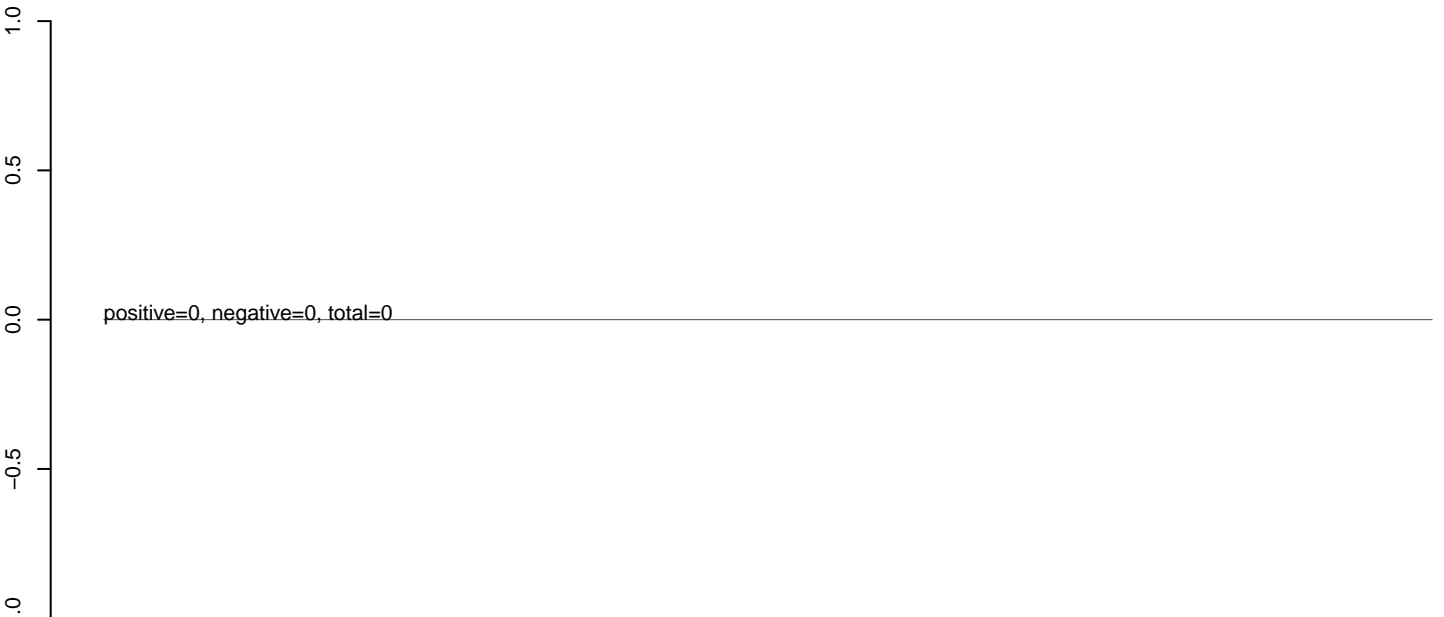
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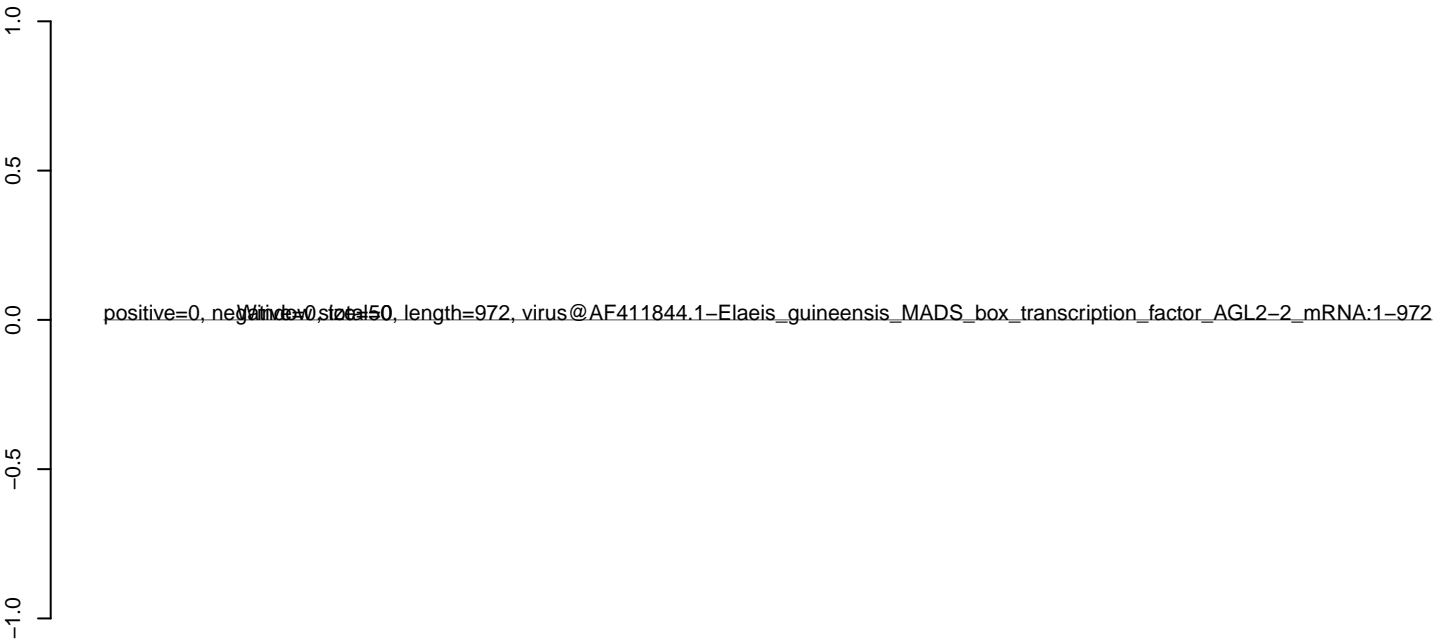
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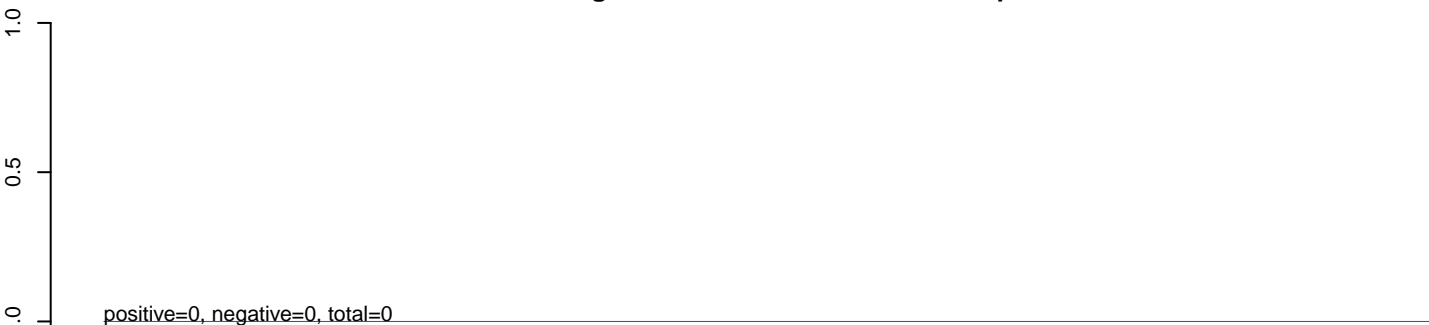
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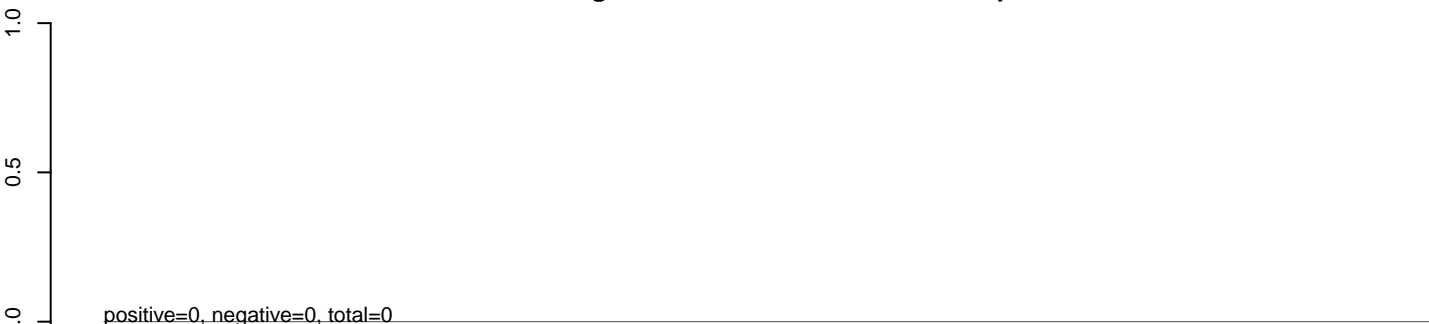
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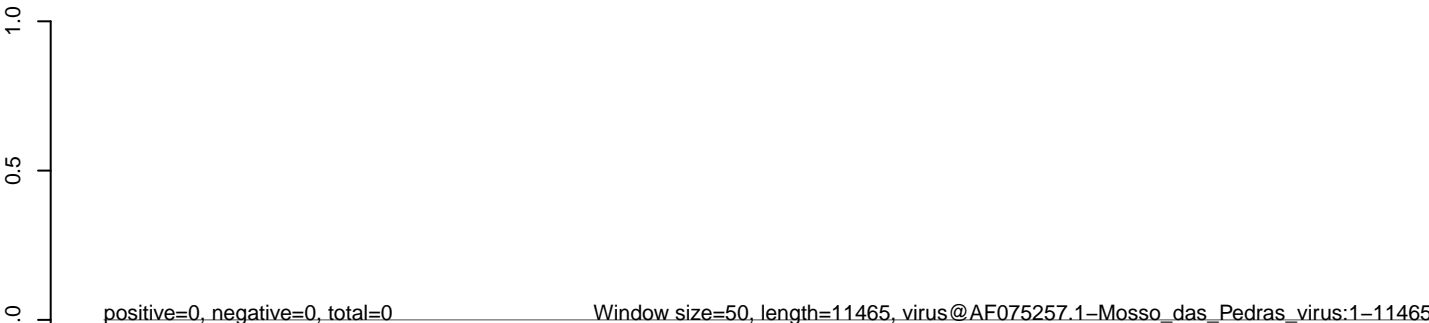
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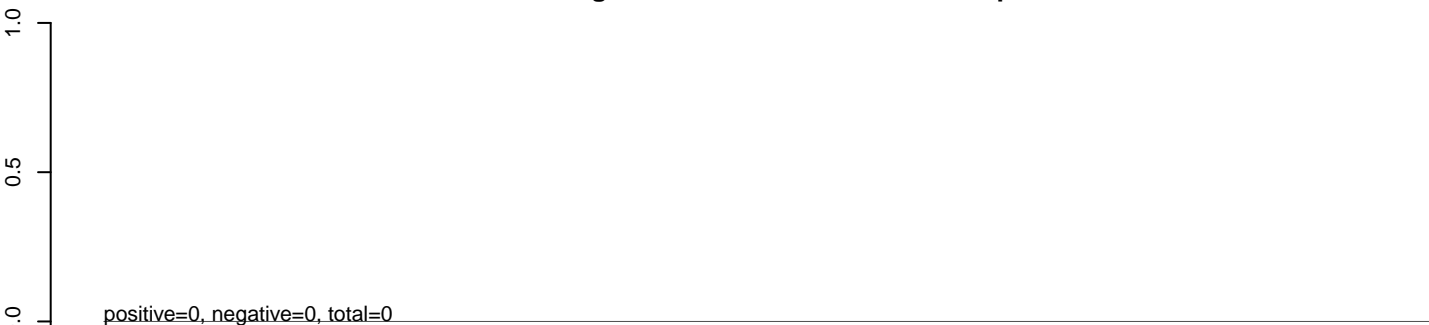


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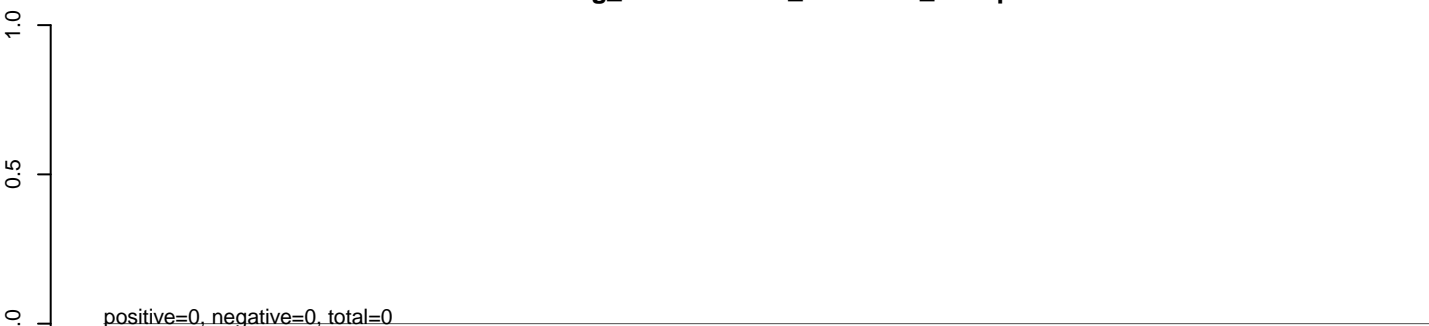


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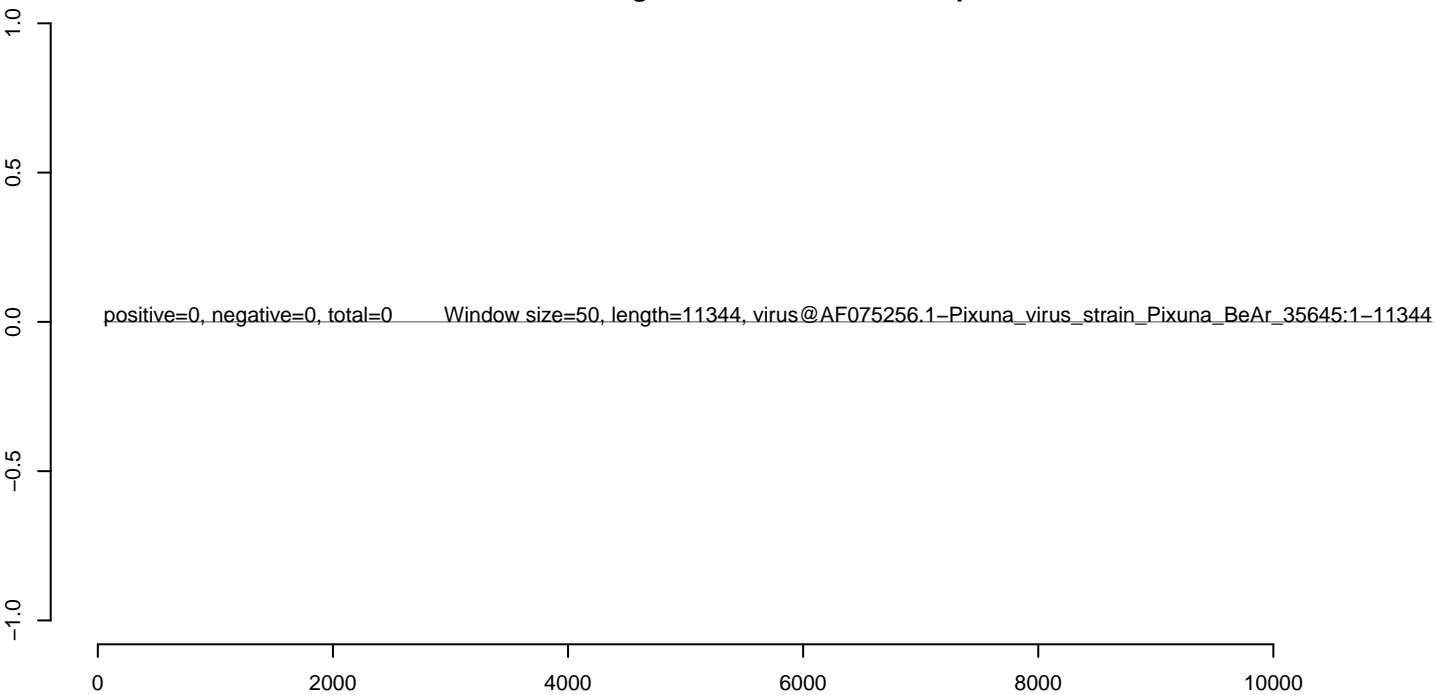
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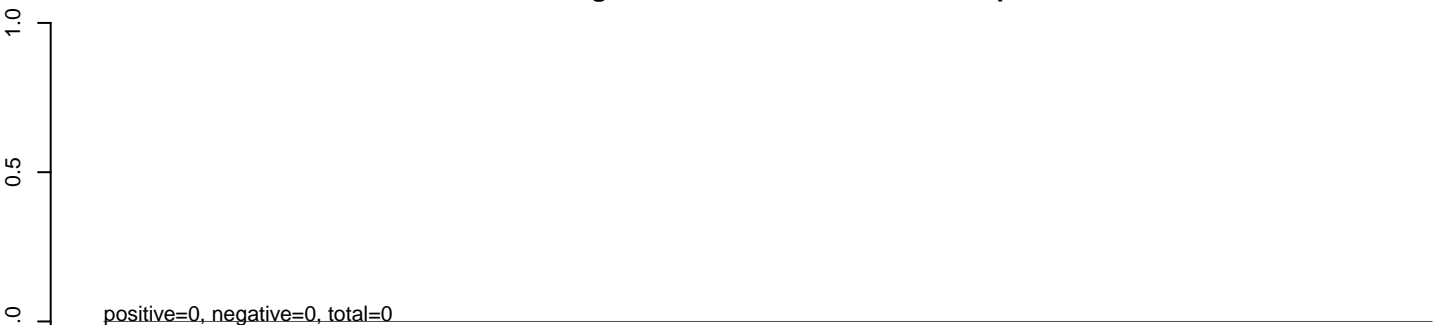
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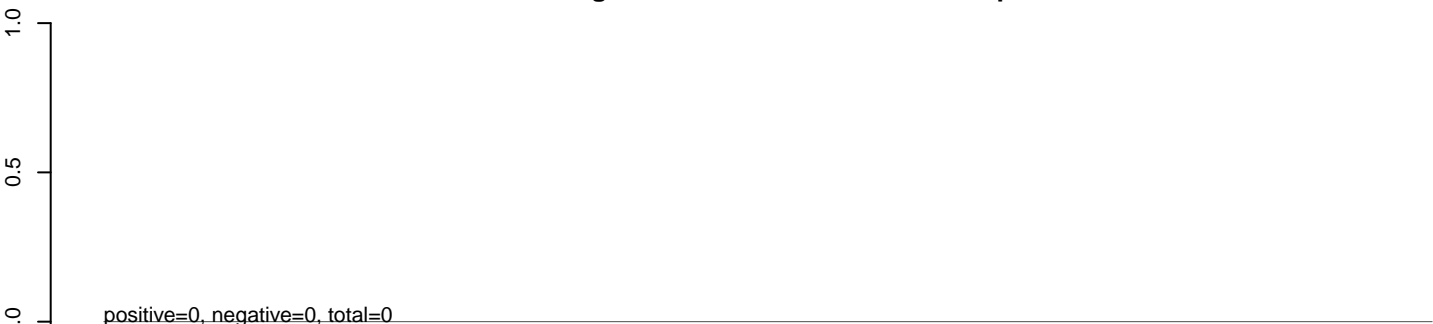
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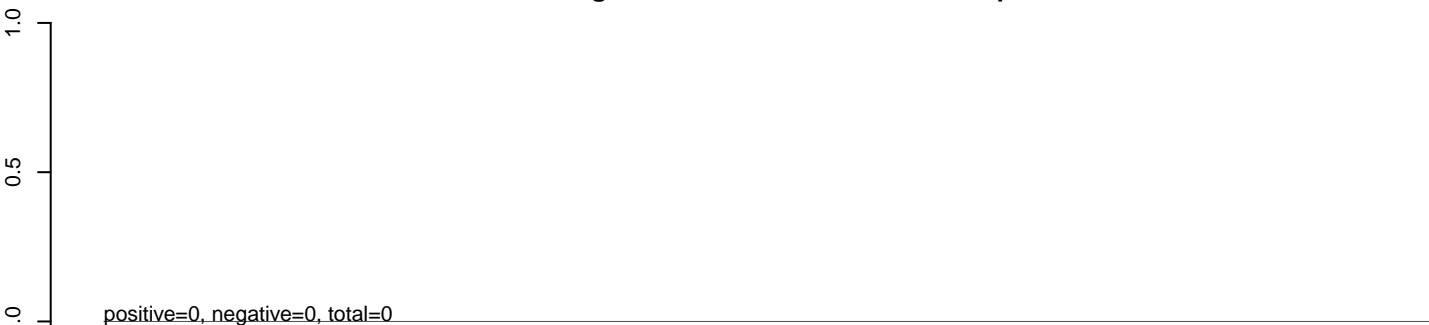


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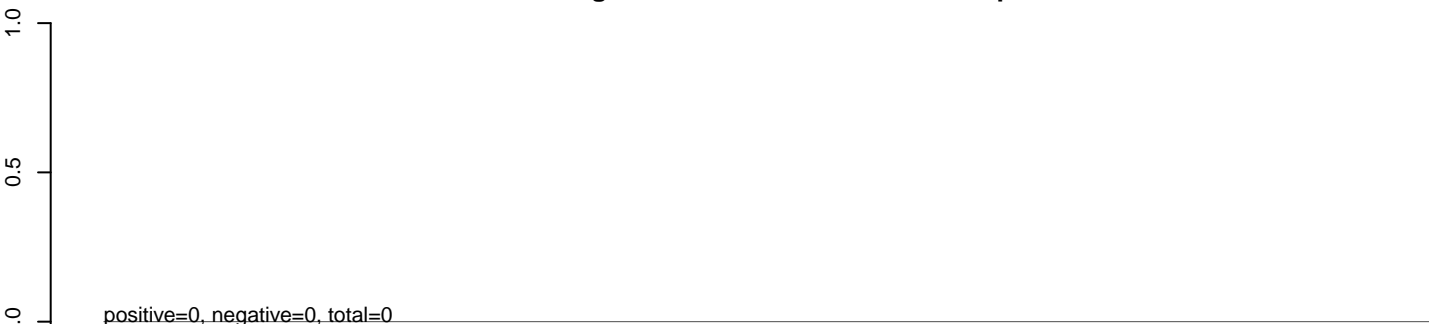


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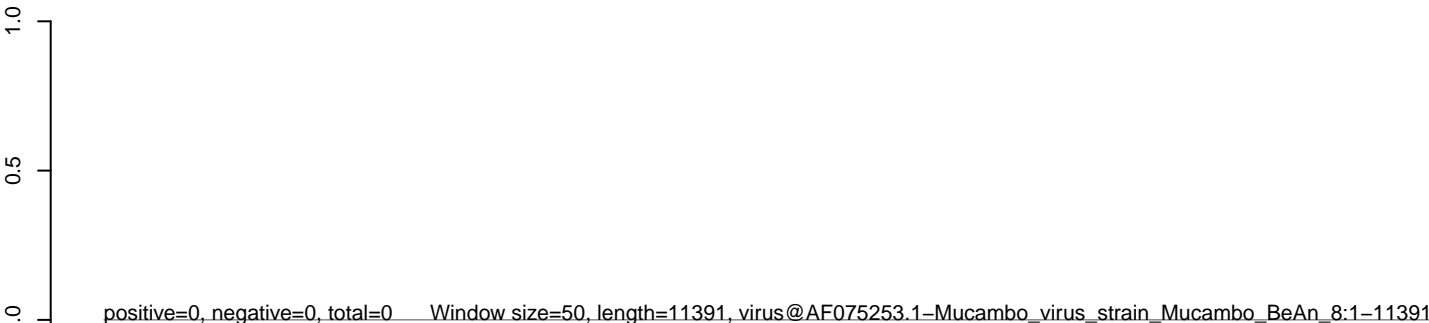
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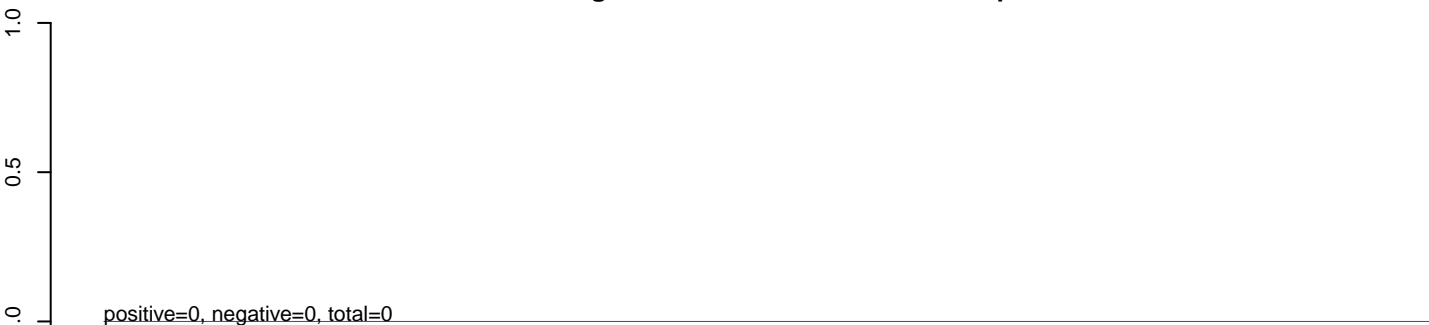


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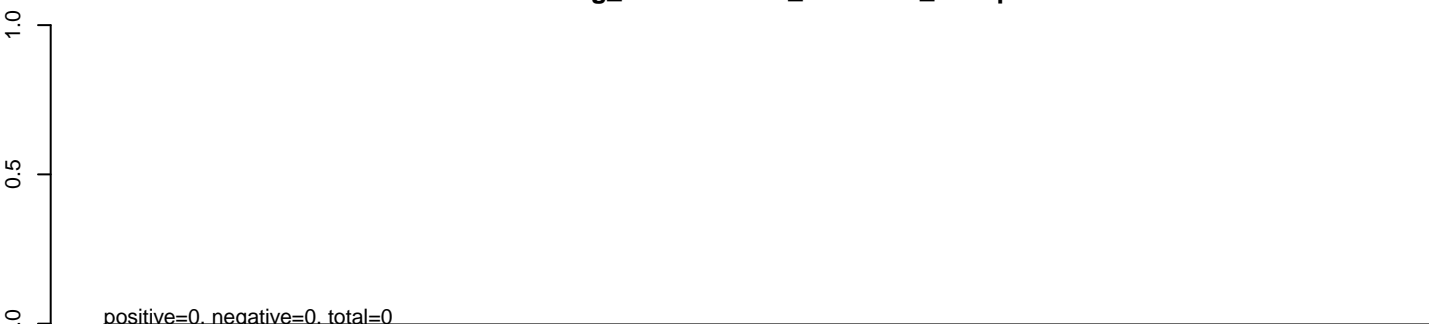


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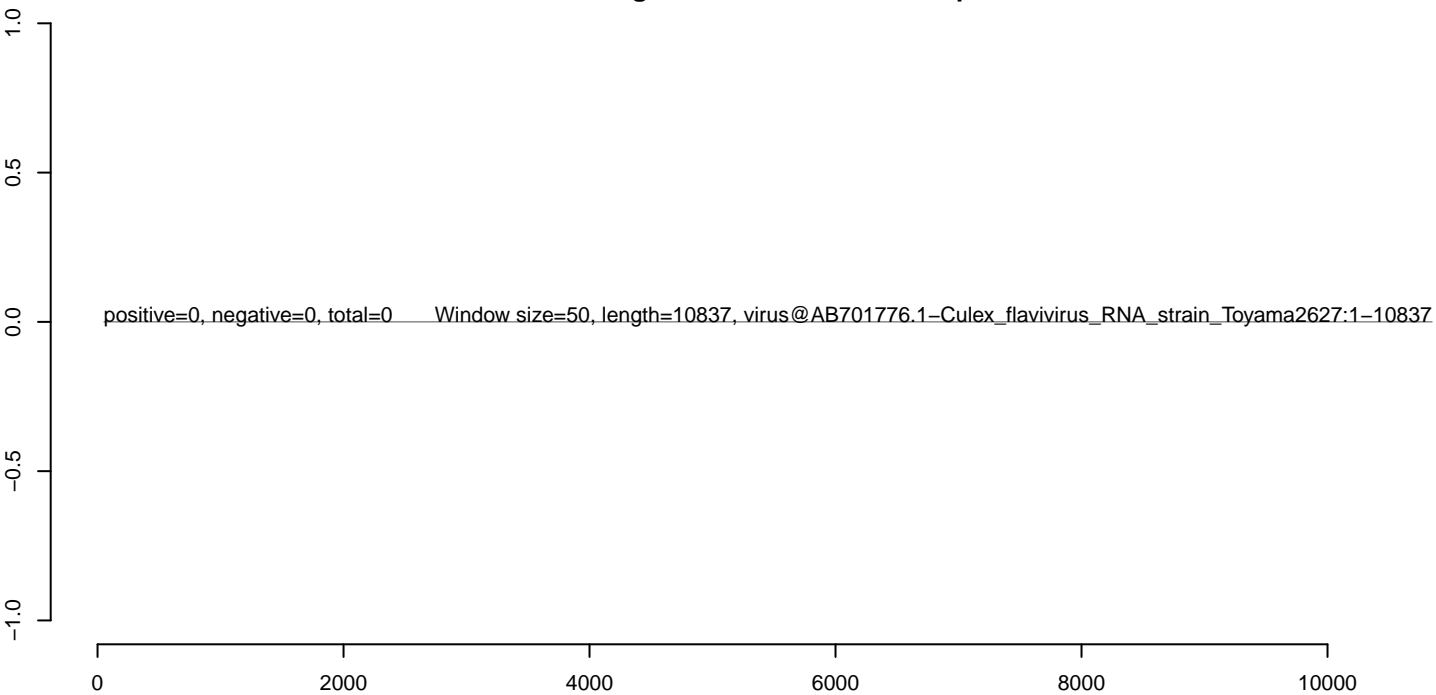
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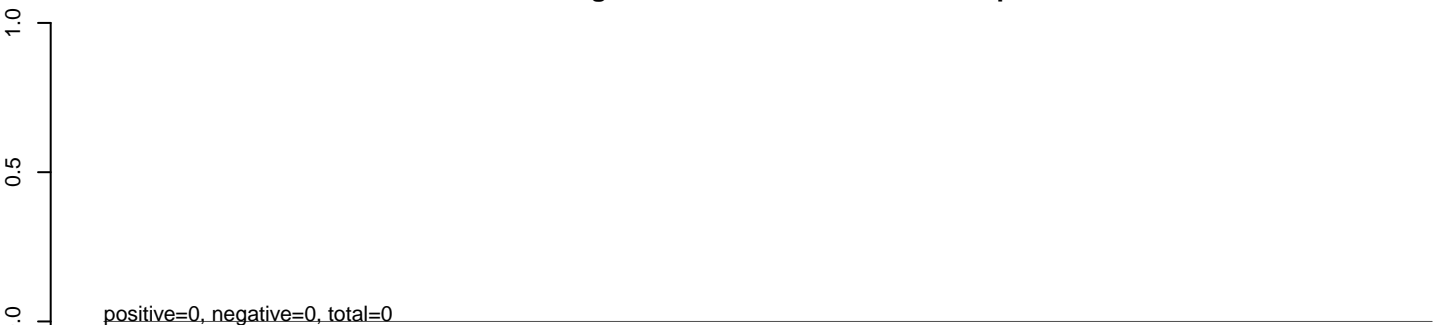
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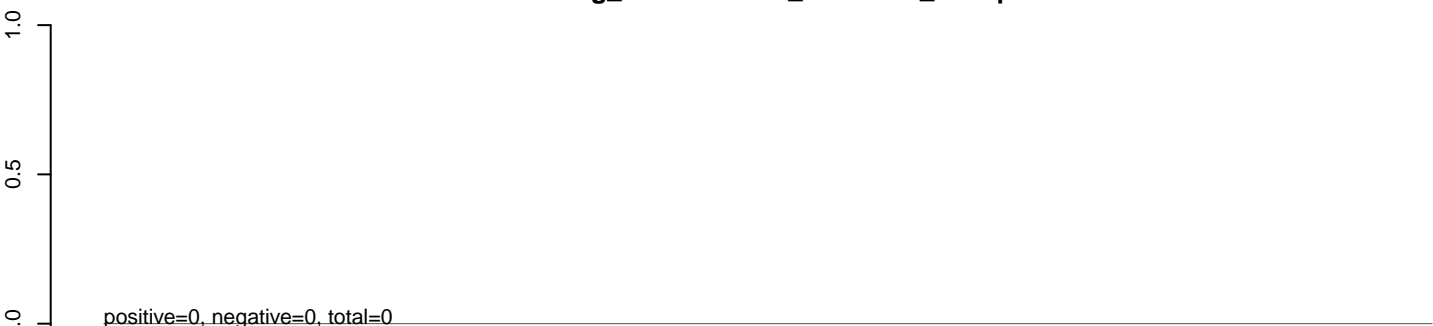
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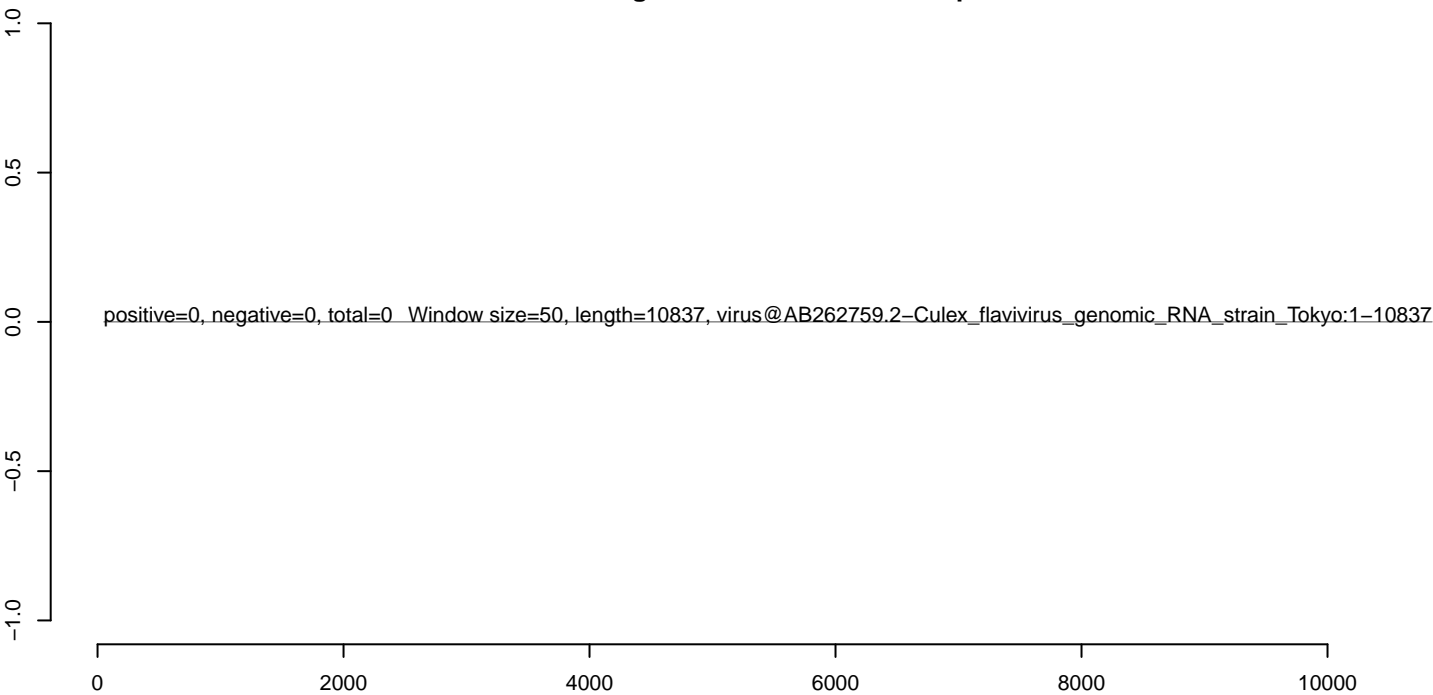
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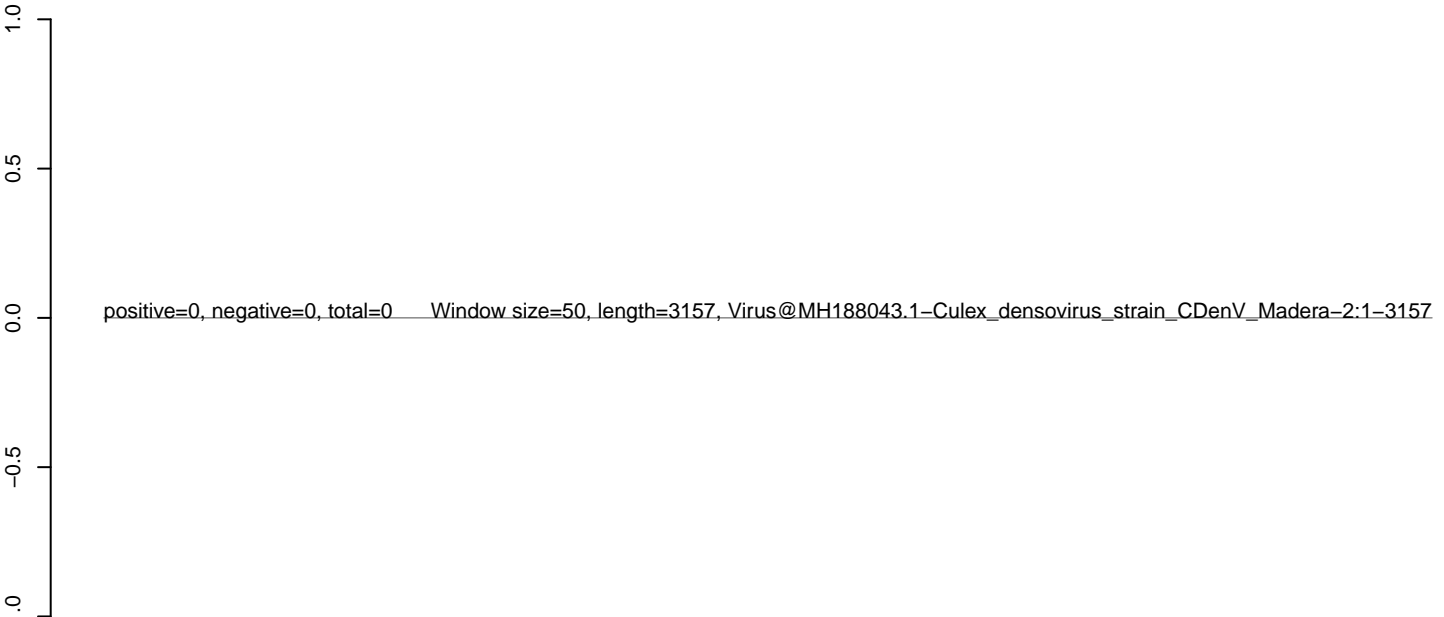
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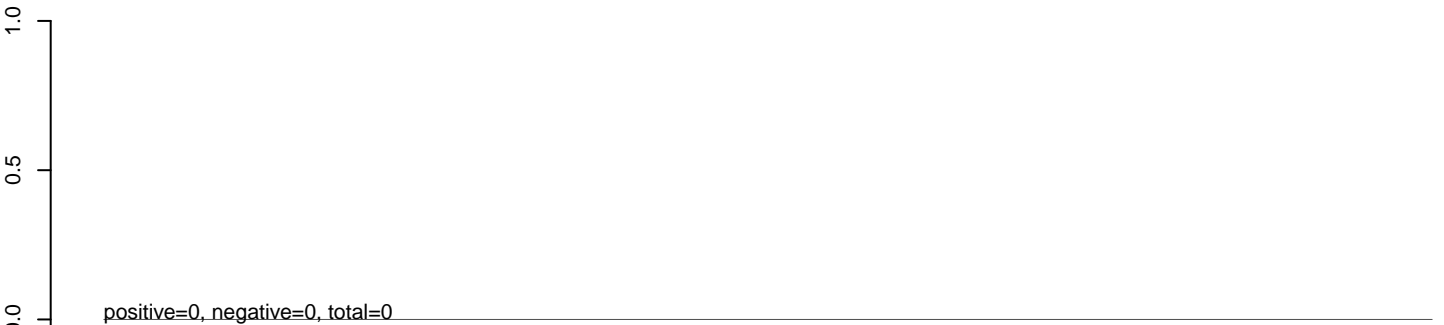
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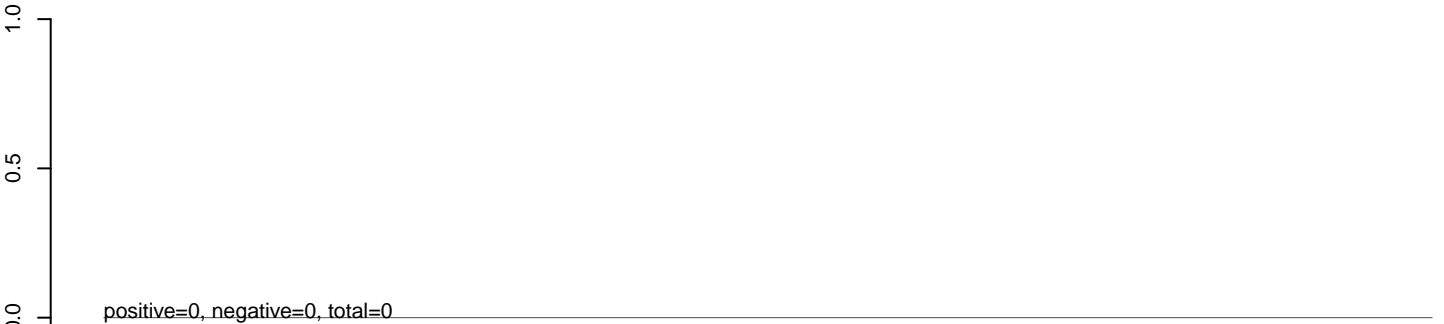
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