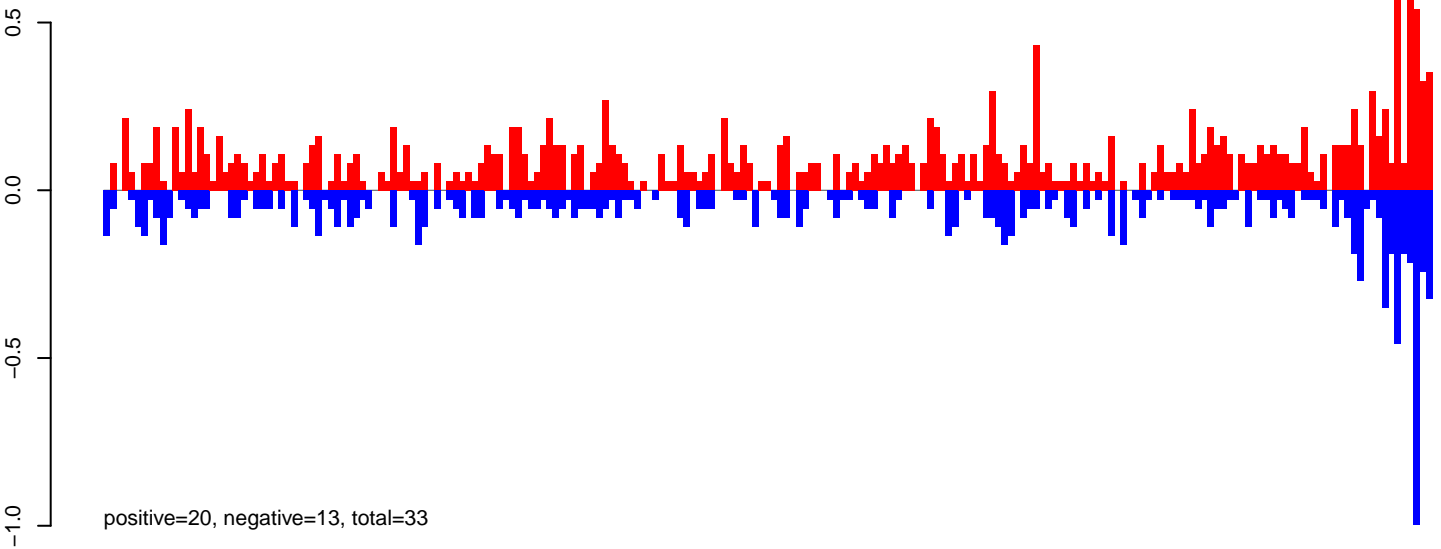
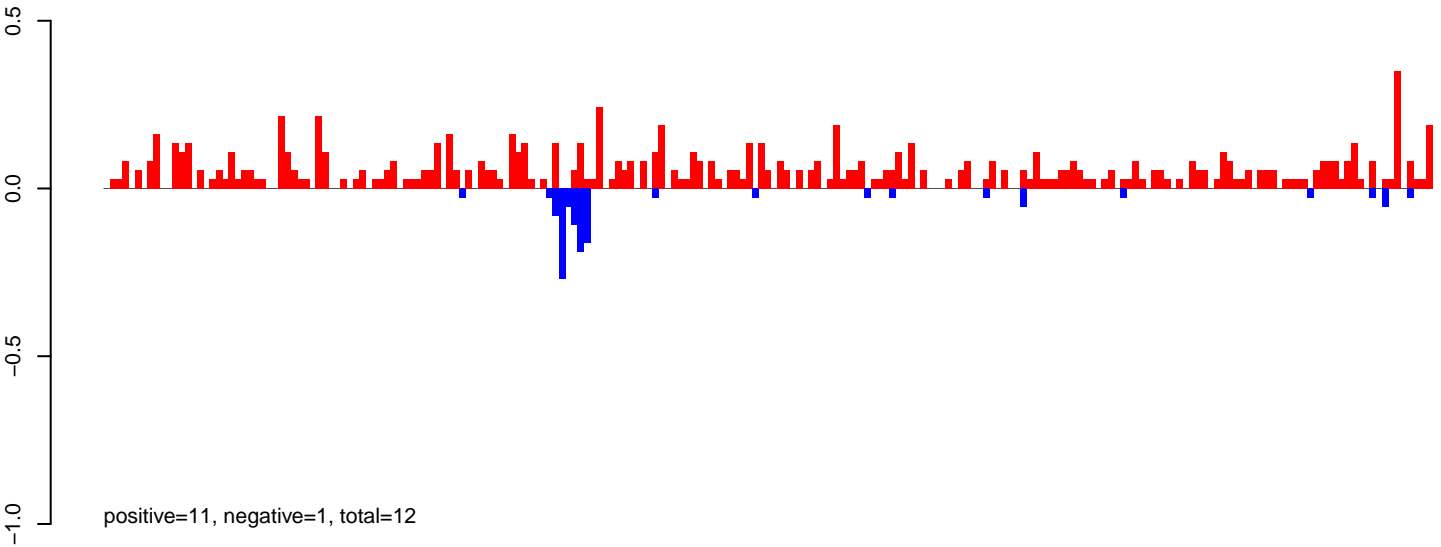


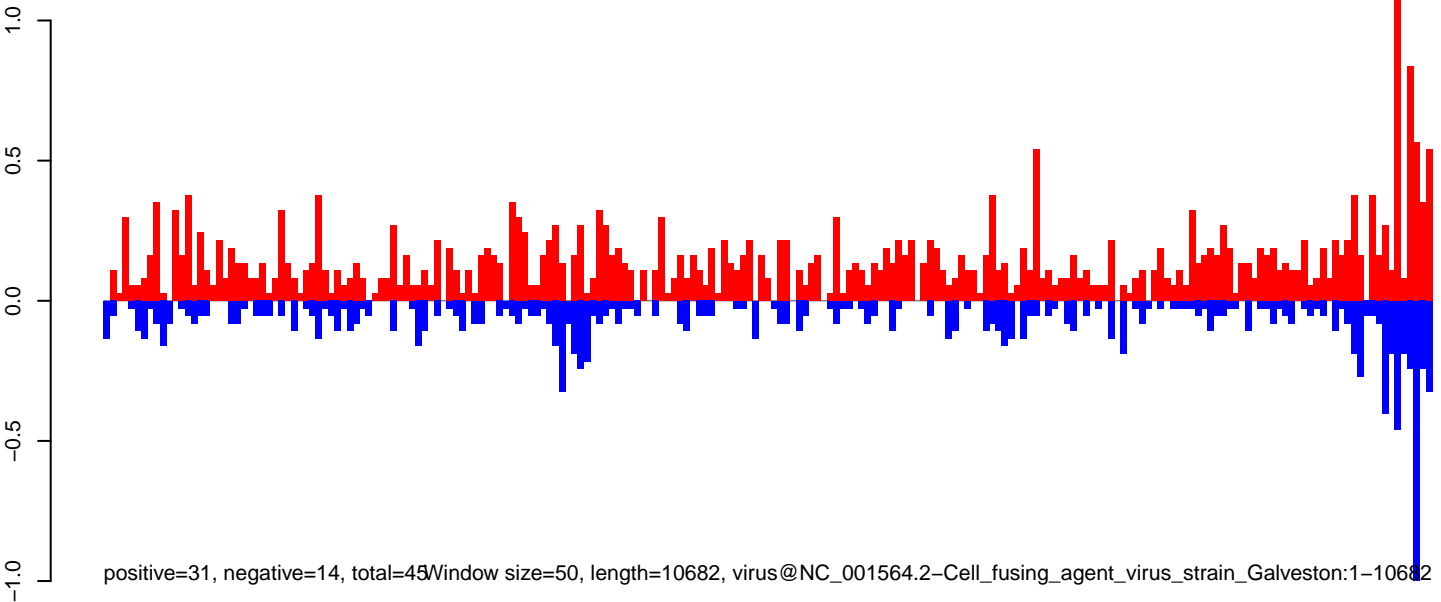
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



Window size=50, length=10682, virus@NC_001564.2-Cell_fusing_agent_virus_strain_Galveston:1-10682

0 2000 4000 6000 8000 10000

AeAeg_Ovary_GH.18_23.rep

0
-5
-10
-15

positive=0, negative=29, total=29

AeAeg_Ovary_GH.24_35.rep

0
-5
-10
-15

positive=1, negative=170, total=171

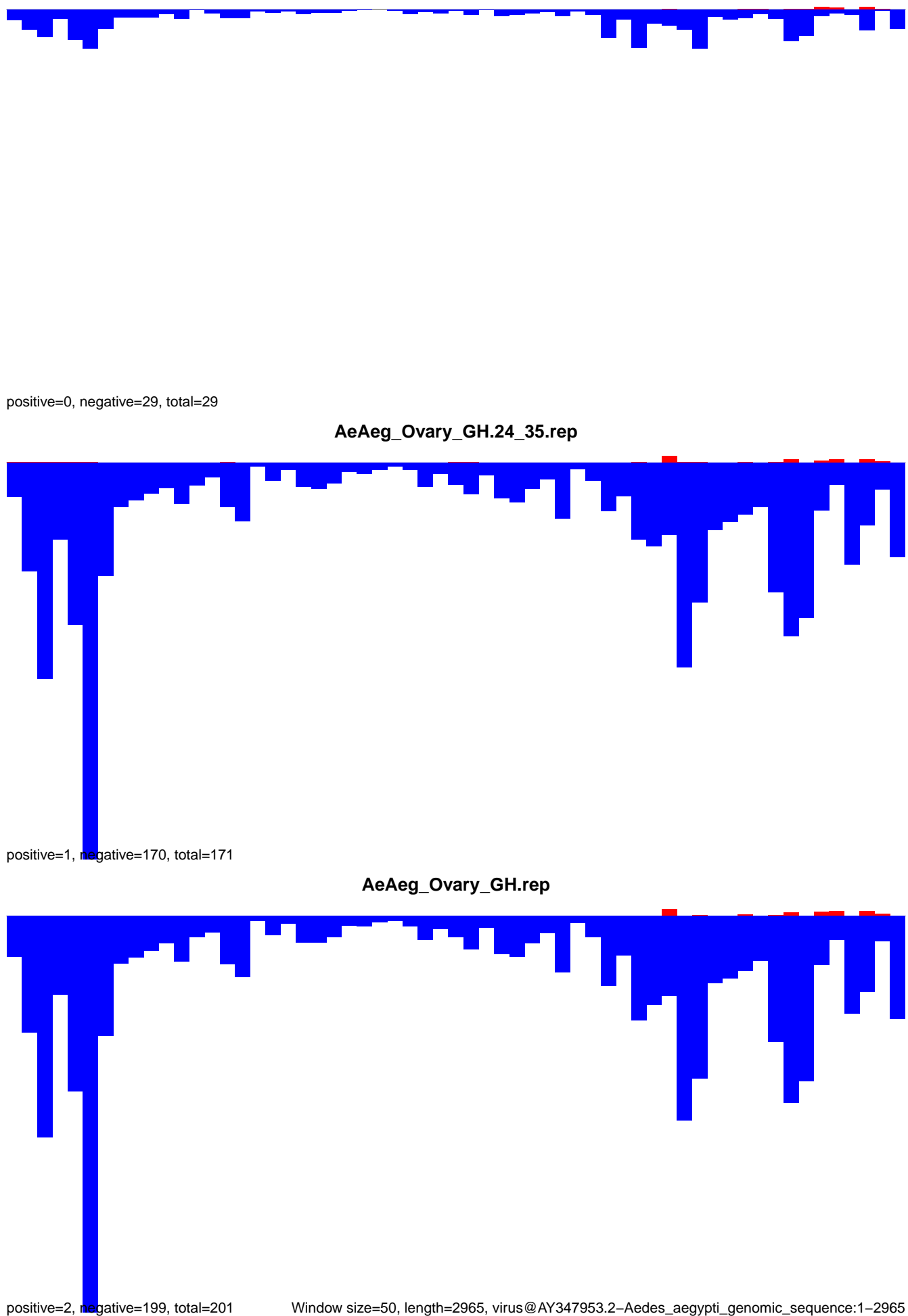
AeAeg_Ovary_GH.rep

0
-5
-10
-15
-20

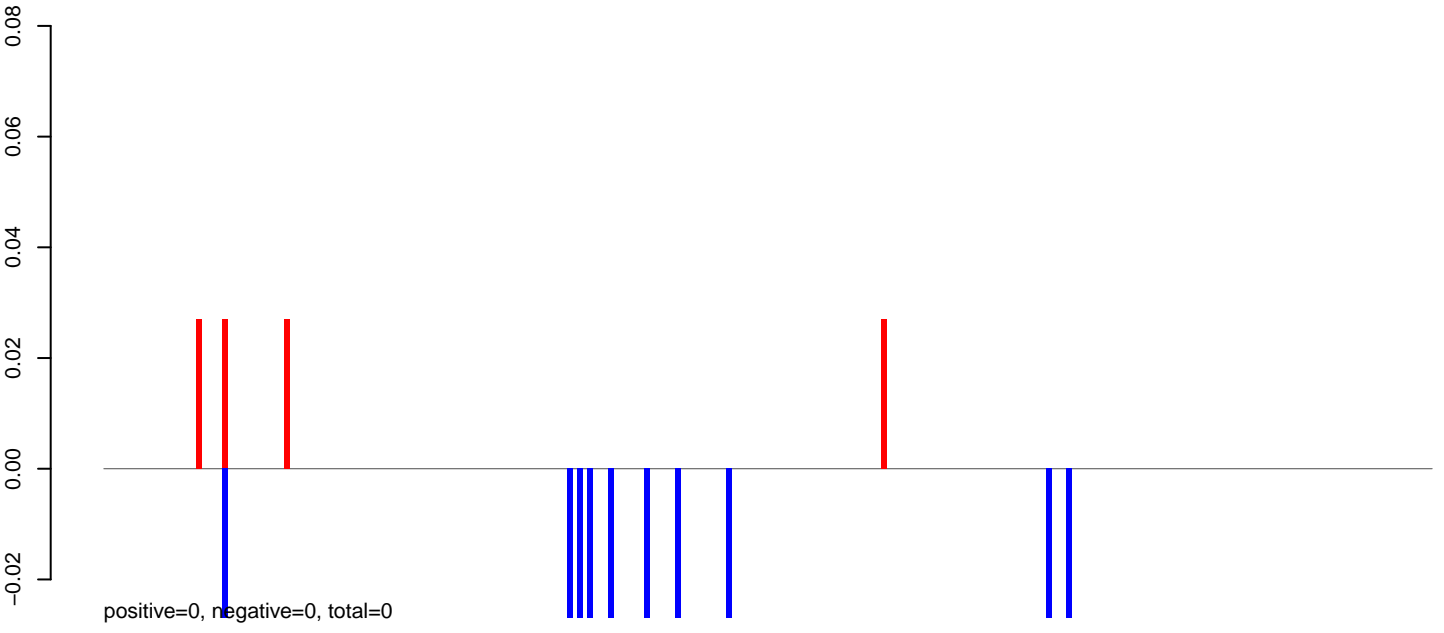
positive=2, negative=199, total=201

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

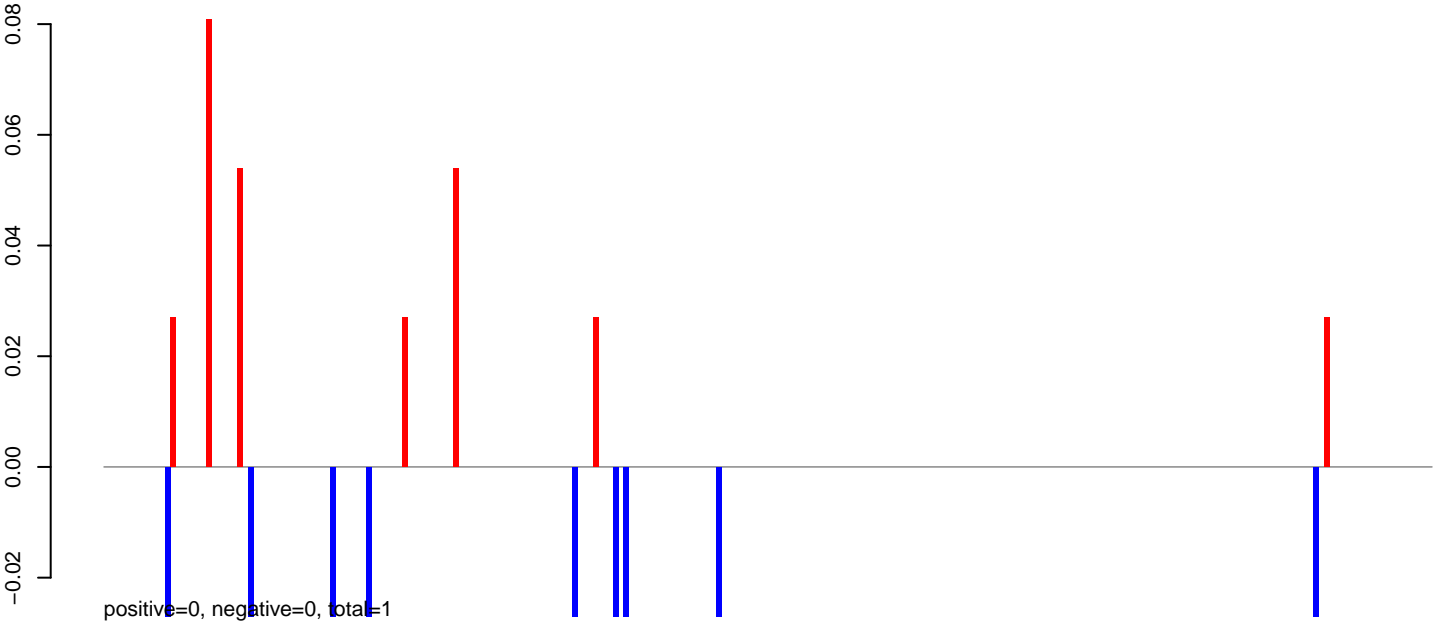
0 500 1000 1500 2000 2500 3000



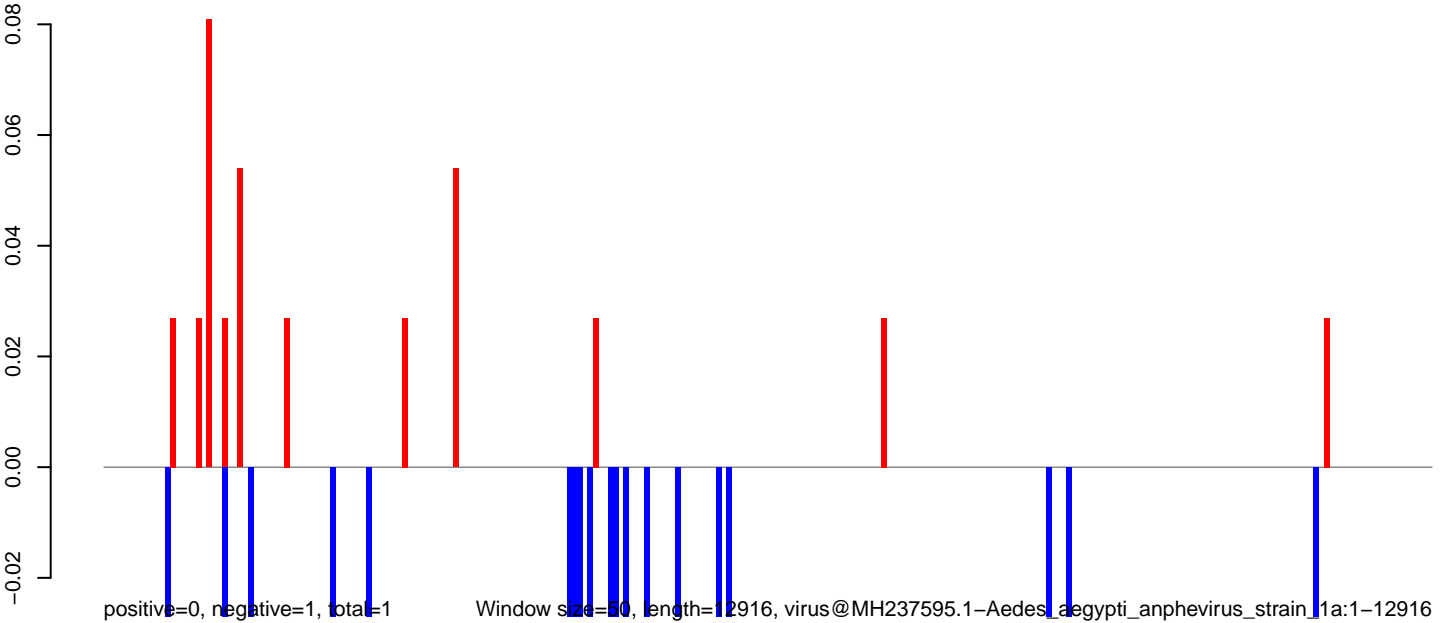
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

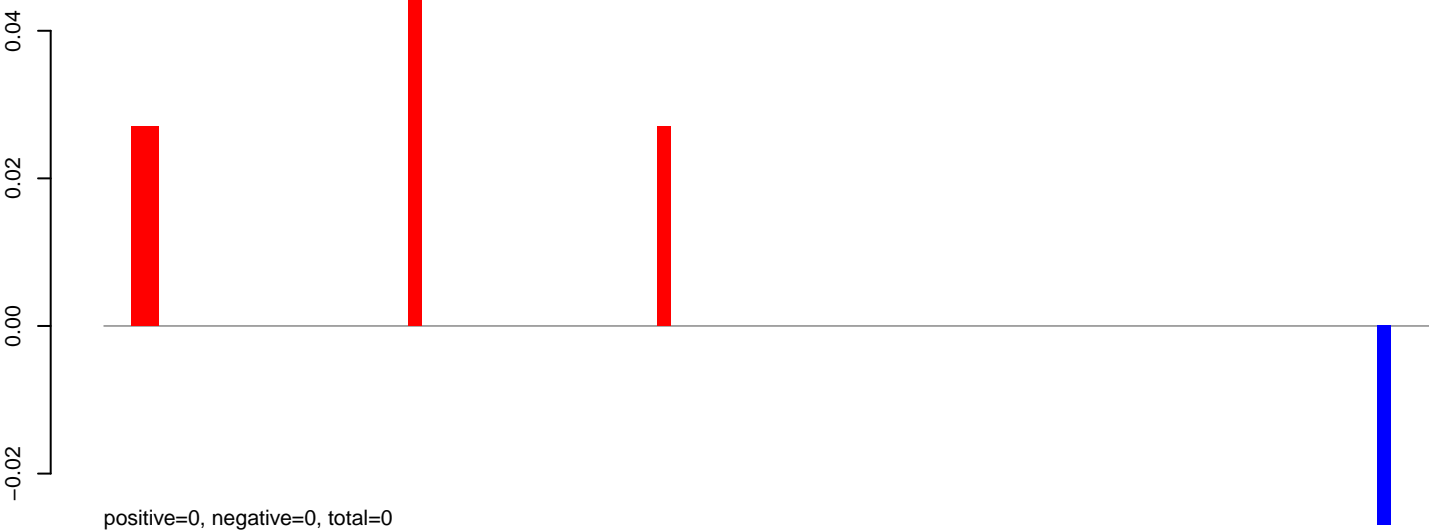


AeAeg_Ovary_GH.rep

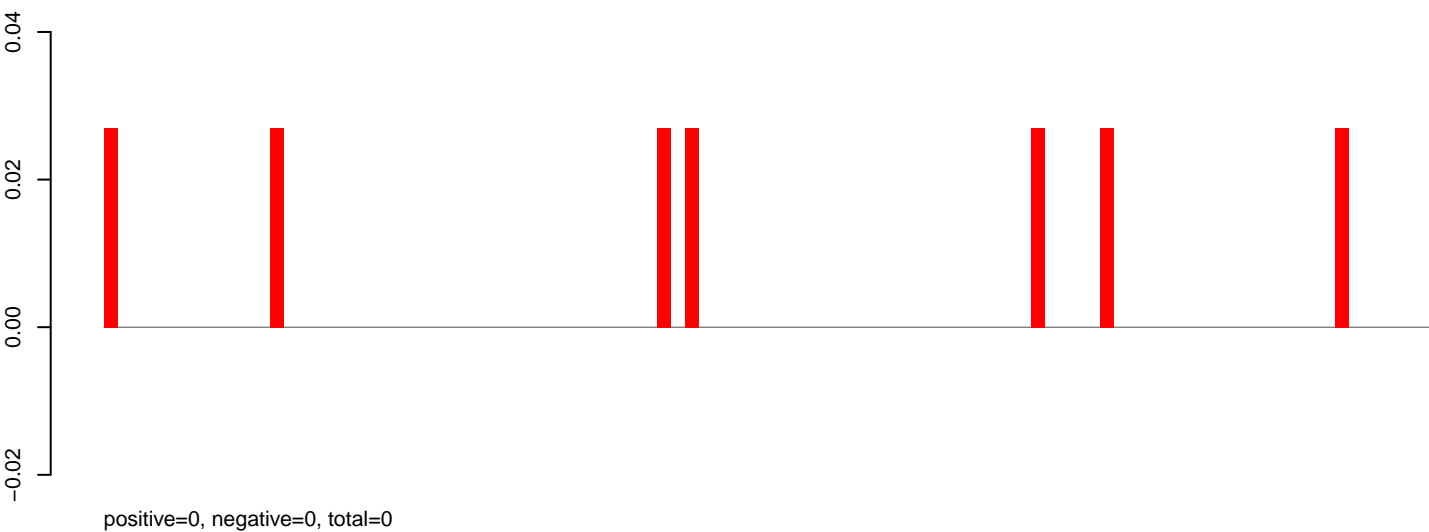


0 2000 4000 6000 8000 10000 12000

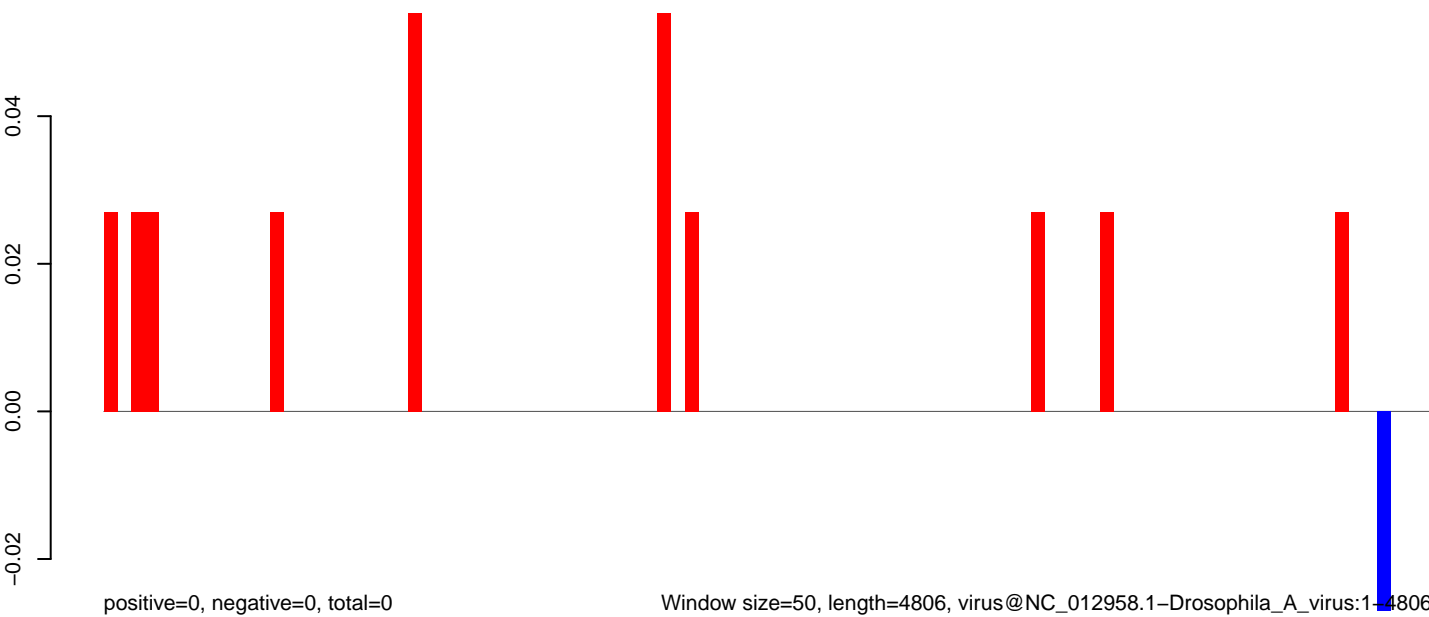
AeAeg_Ovary_GH.18_23.rep



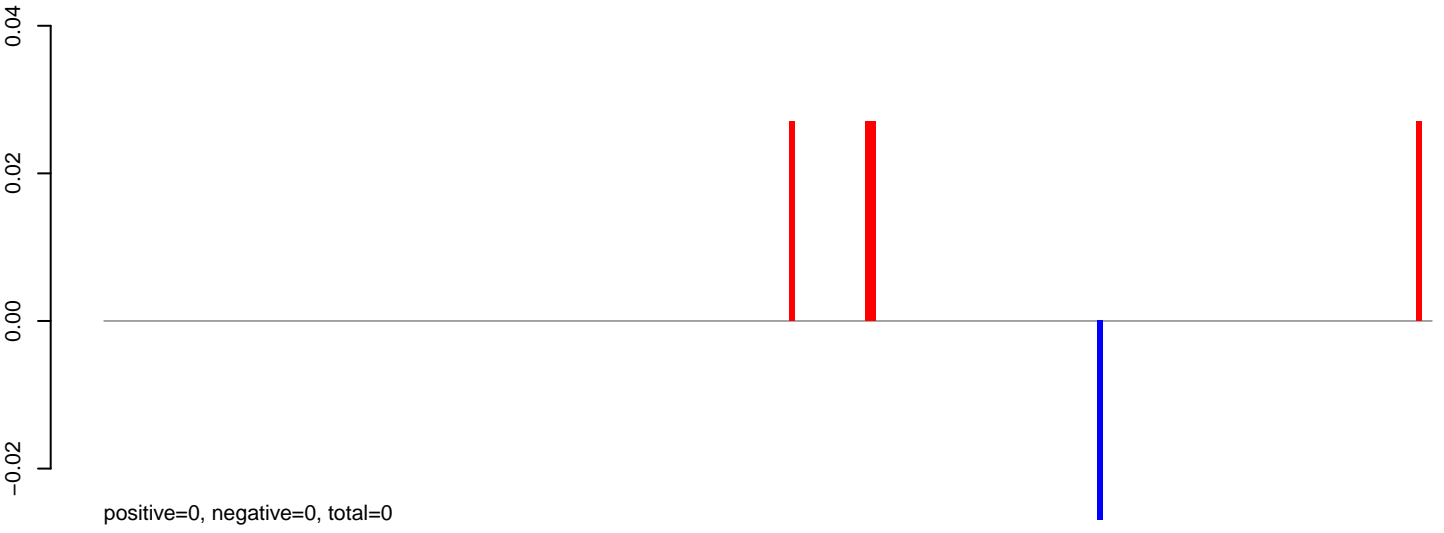
AeAeg_Ovary_GH.24_35.rep



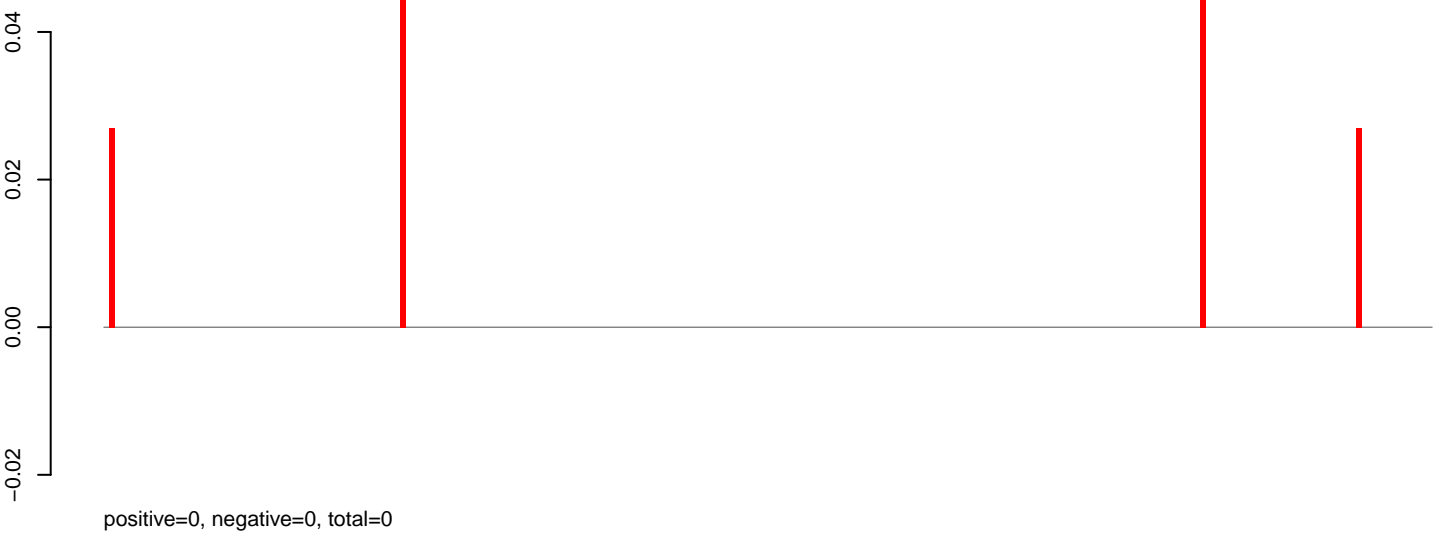
AeAeg_Ovary_GH.rep



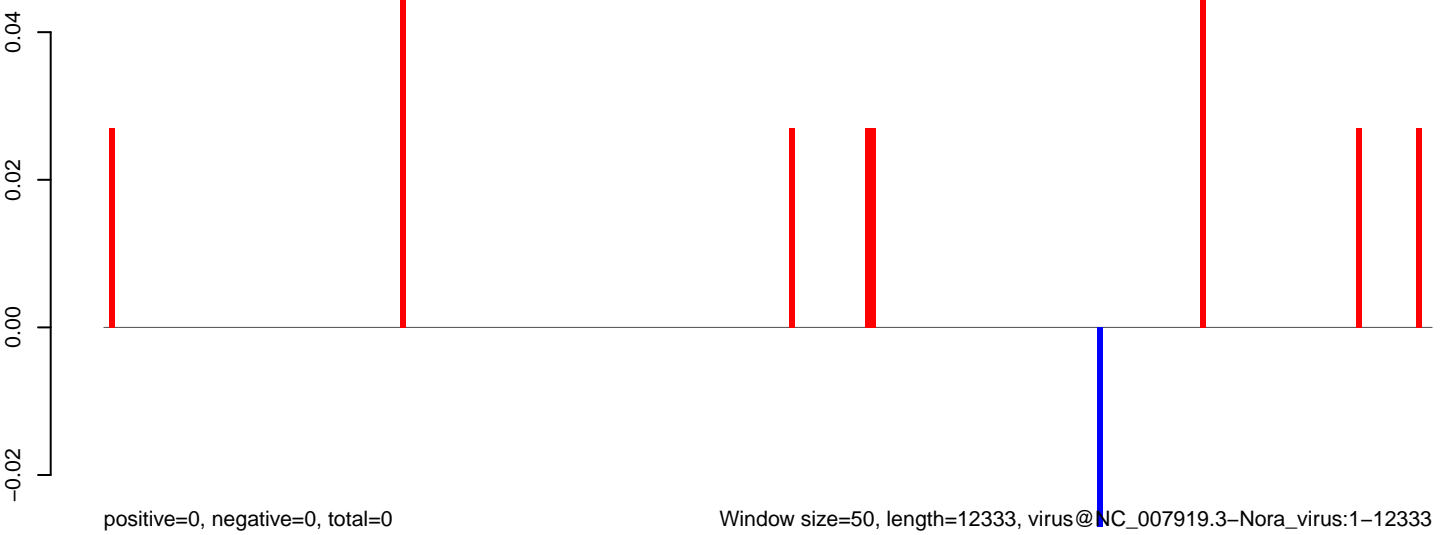
AeAeg_Ovary_GH.18_23.rep



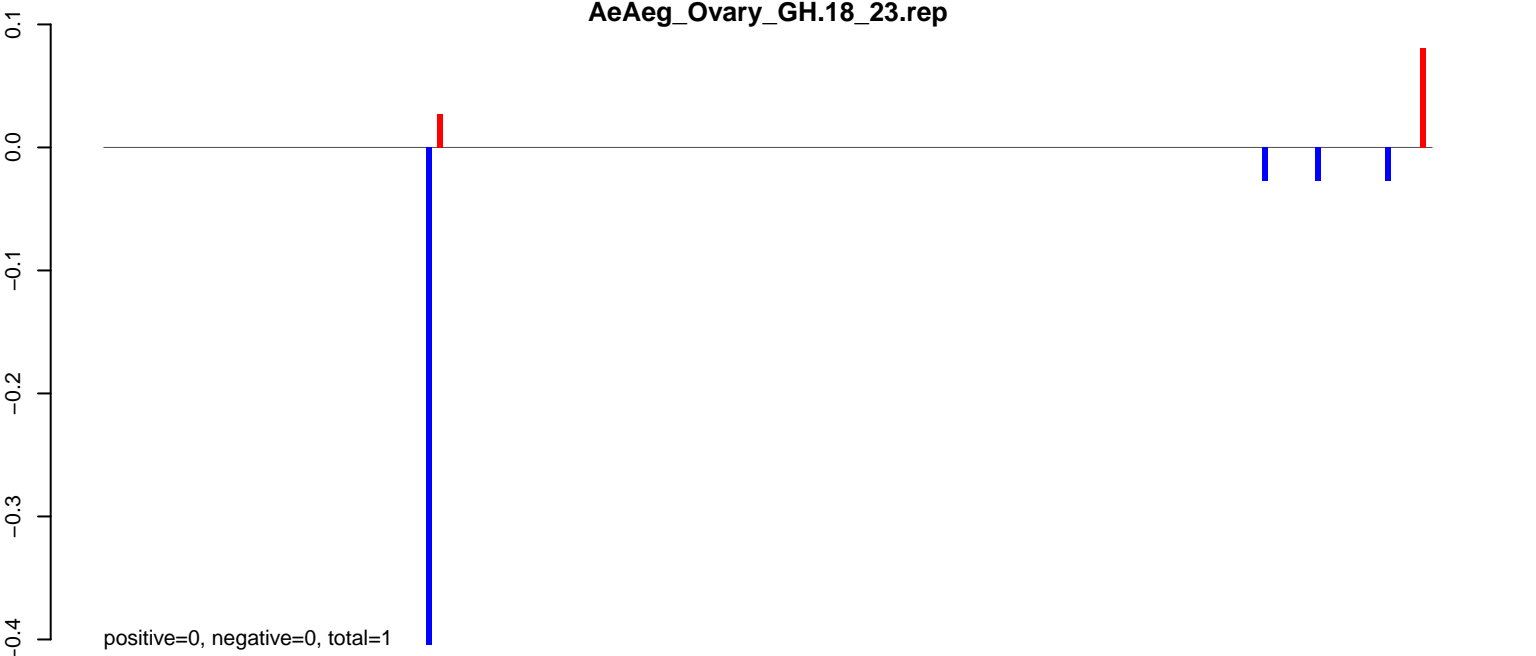
AeAeg_Ovary_GH.24_35.rep



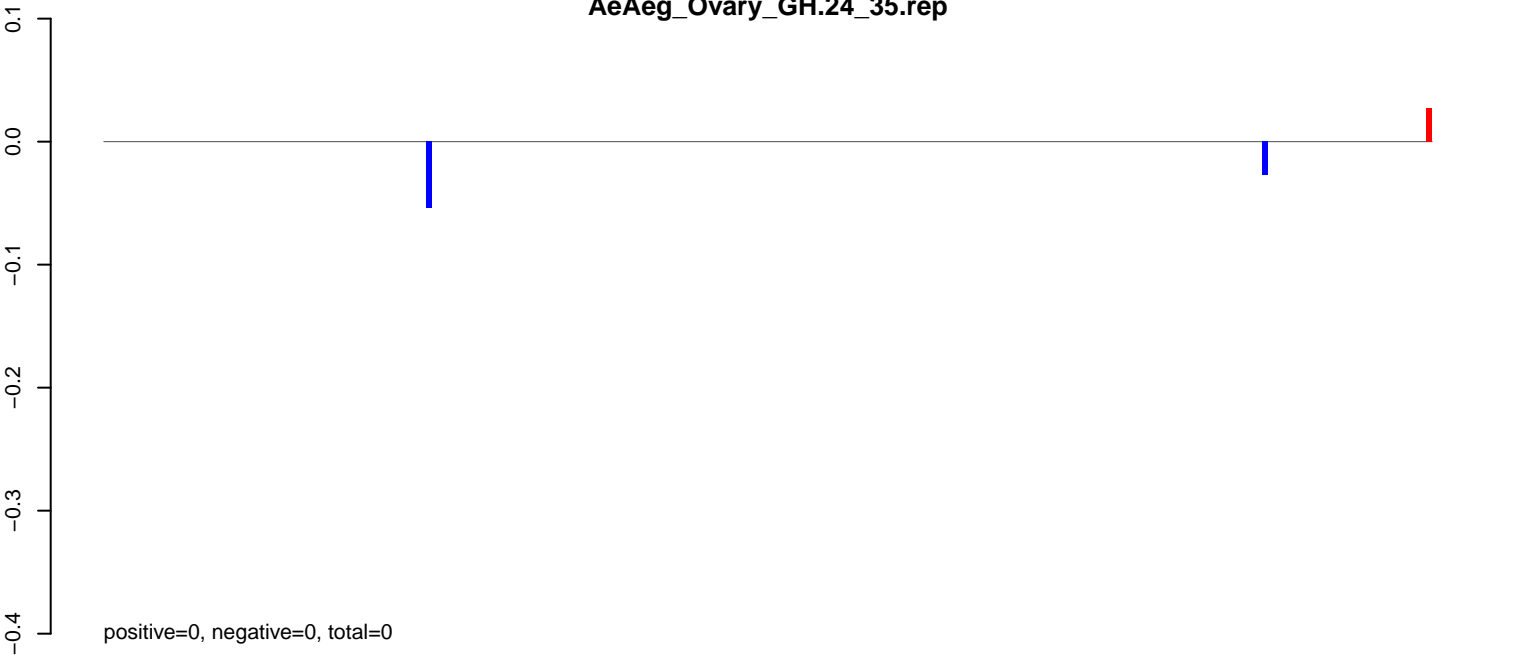
AeAeg_Ovary_GH.rep



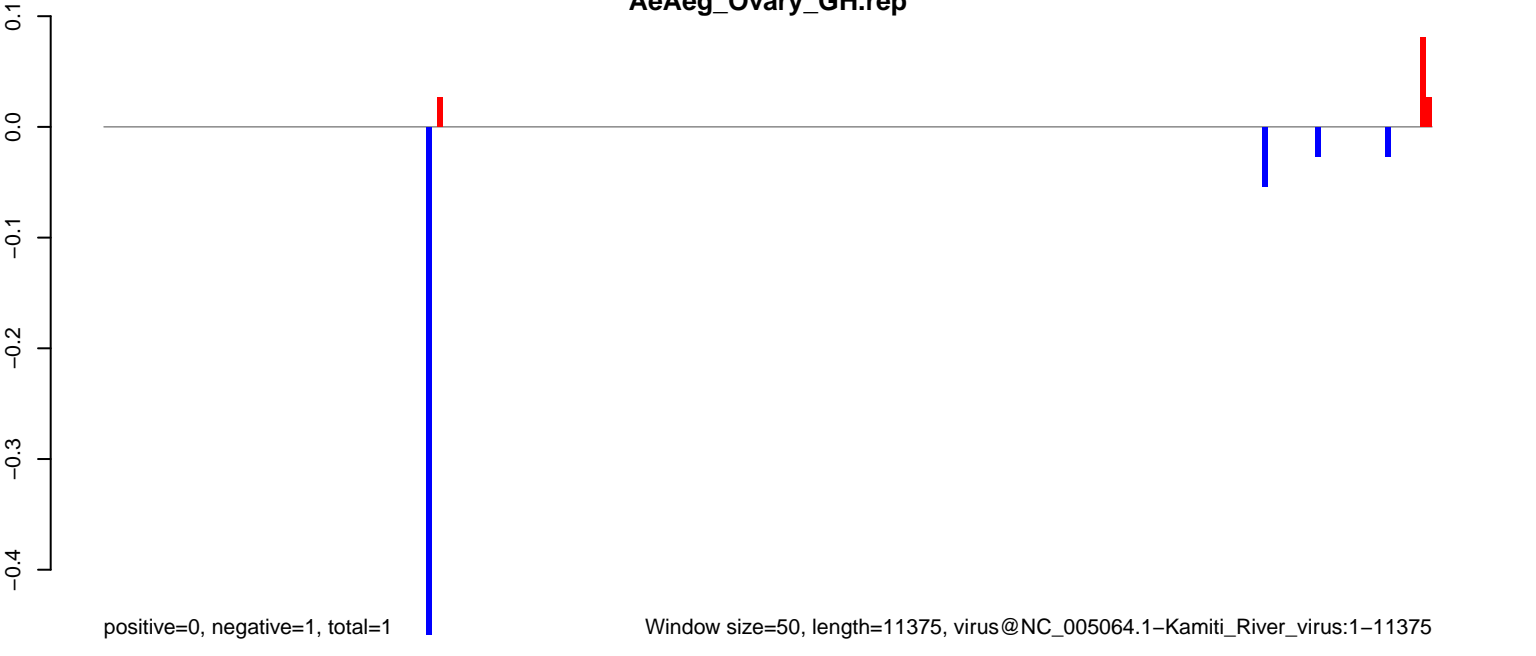
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep

0.3
0.2
0.1
0.0
-0.1

positive=1, negative=0, total=1

AeAeg_Ovary_GH.24_35.rep

0.3
0.2
0.1
0.0
-0.1

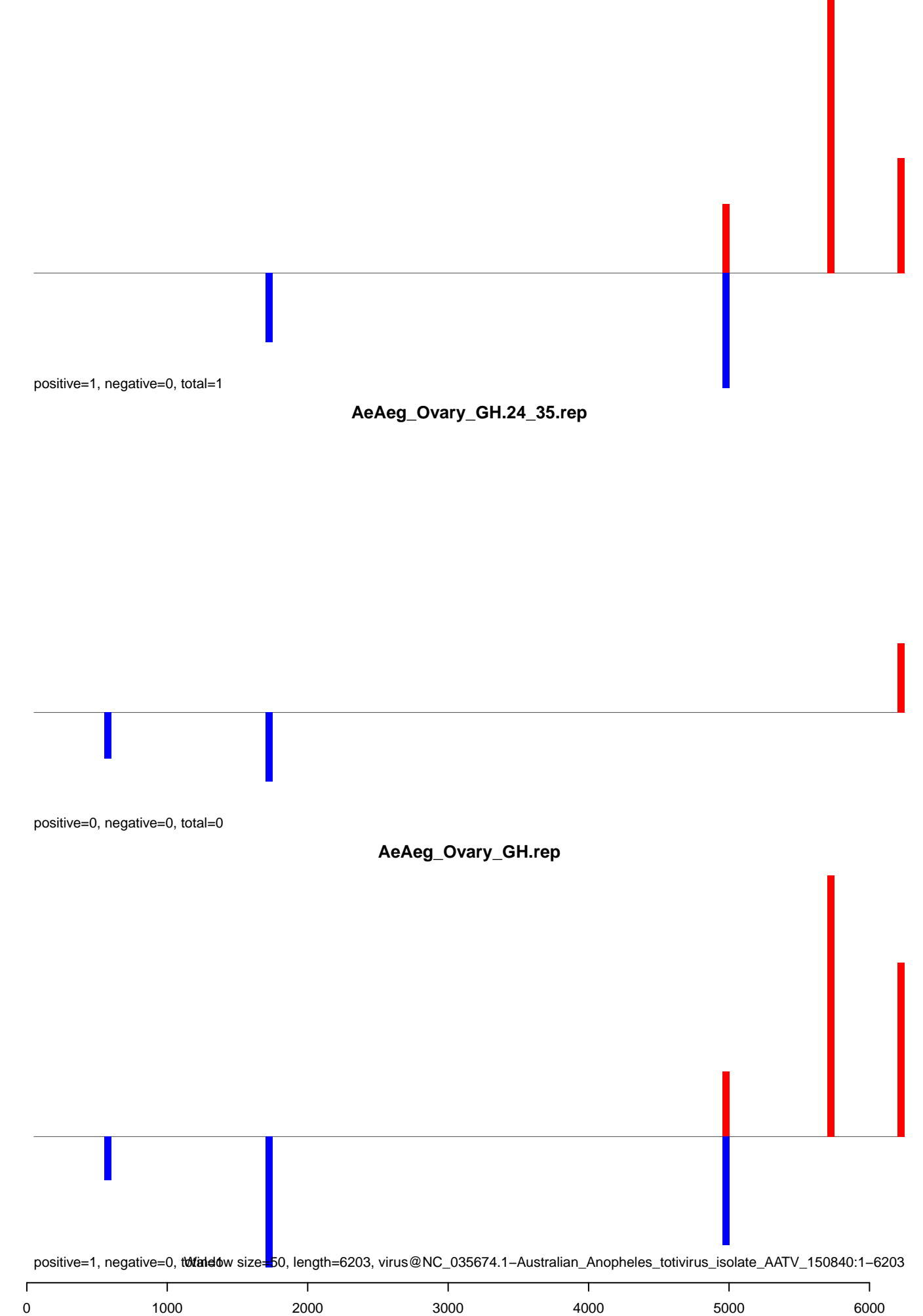
positive=0, negative=0, total=0

AeAeg_Ovary_GH.rep

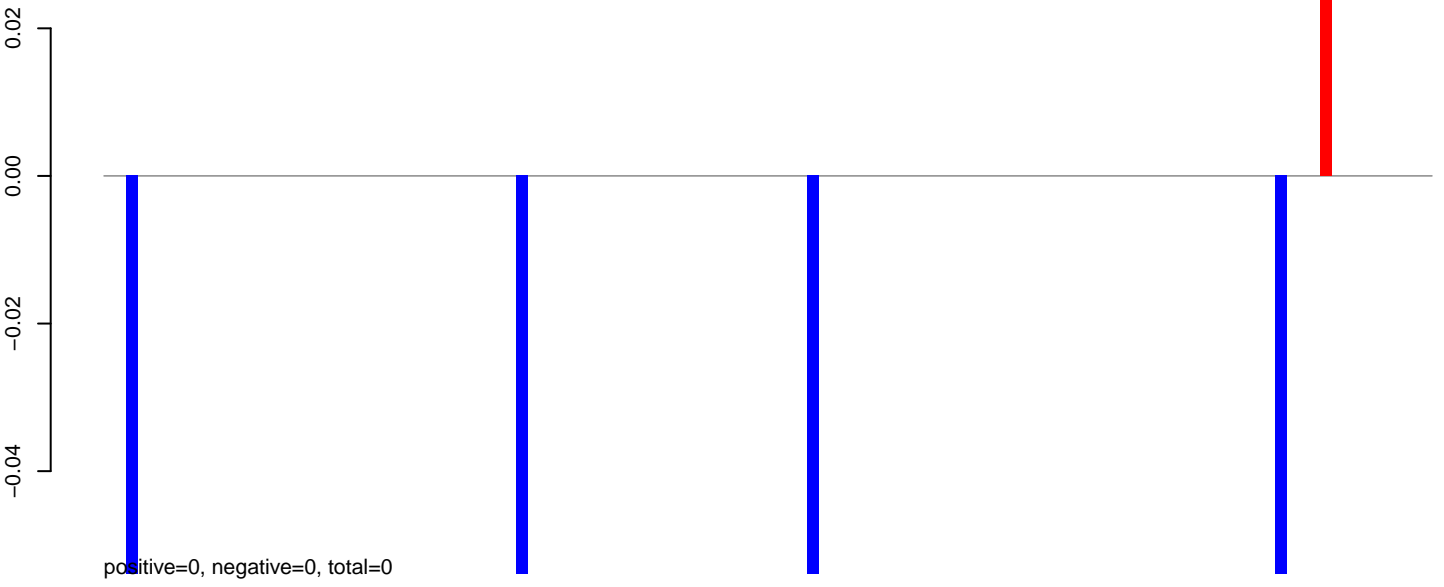
0.3
0.2
0.1
0.0
-0.1

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

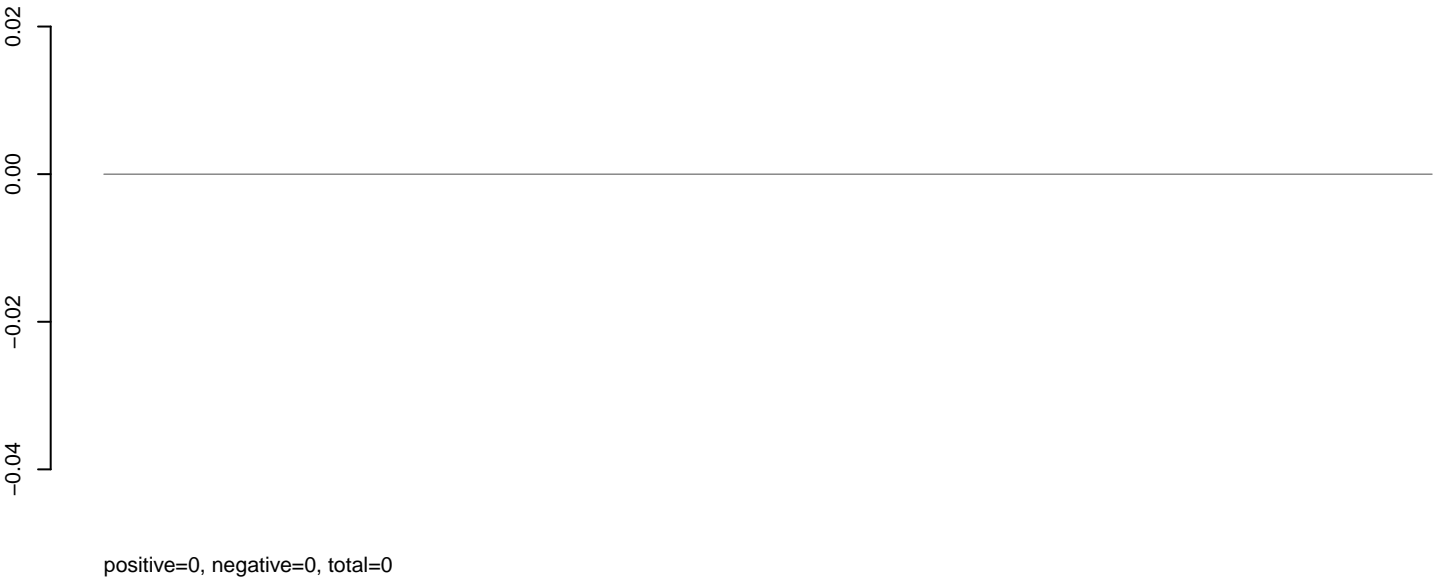
0 1000 2000 3000 4000 5000 6000



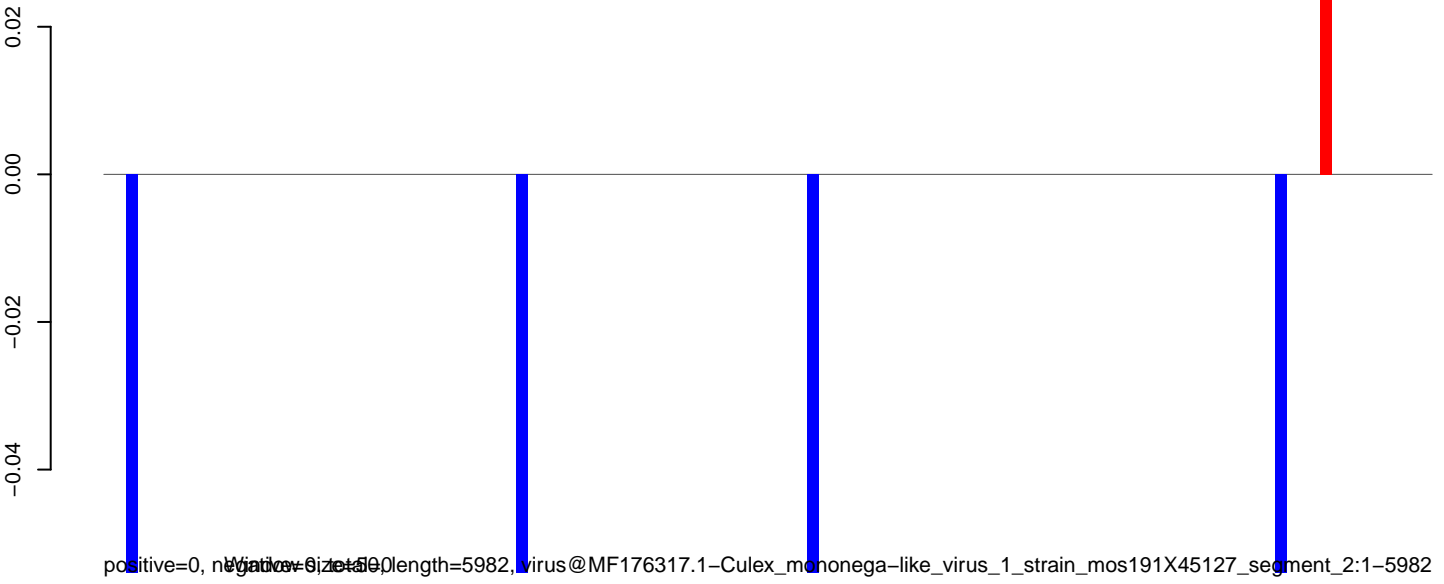
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



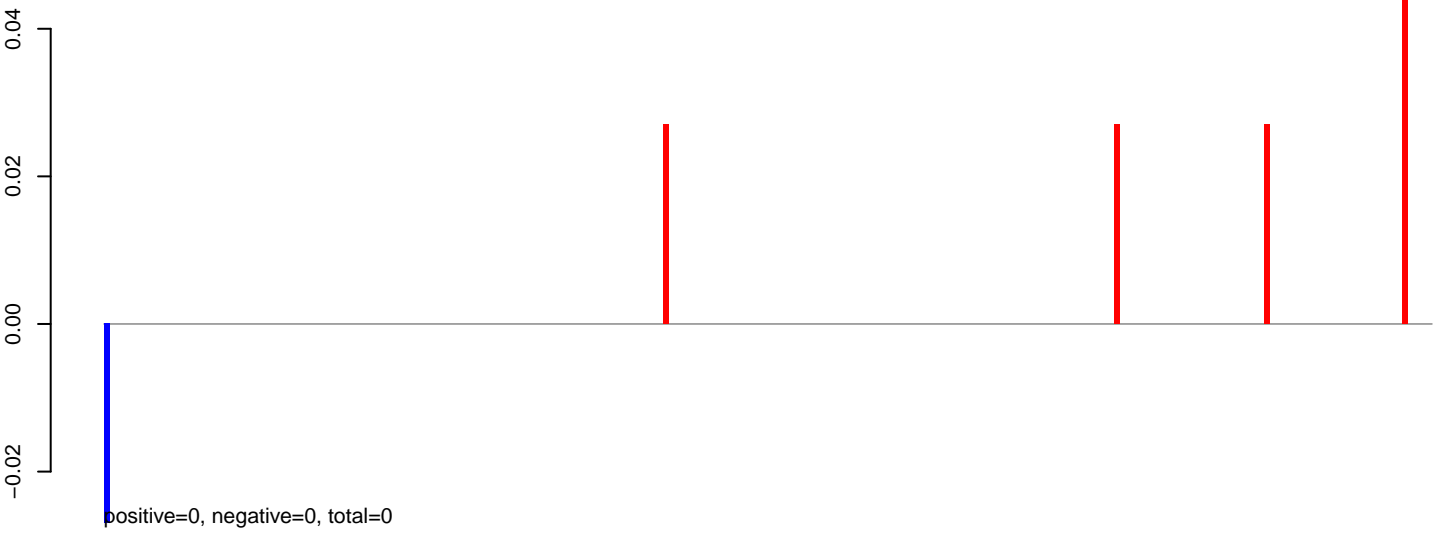
AeAeg_Ovary_GH.rep



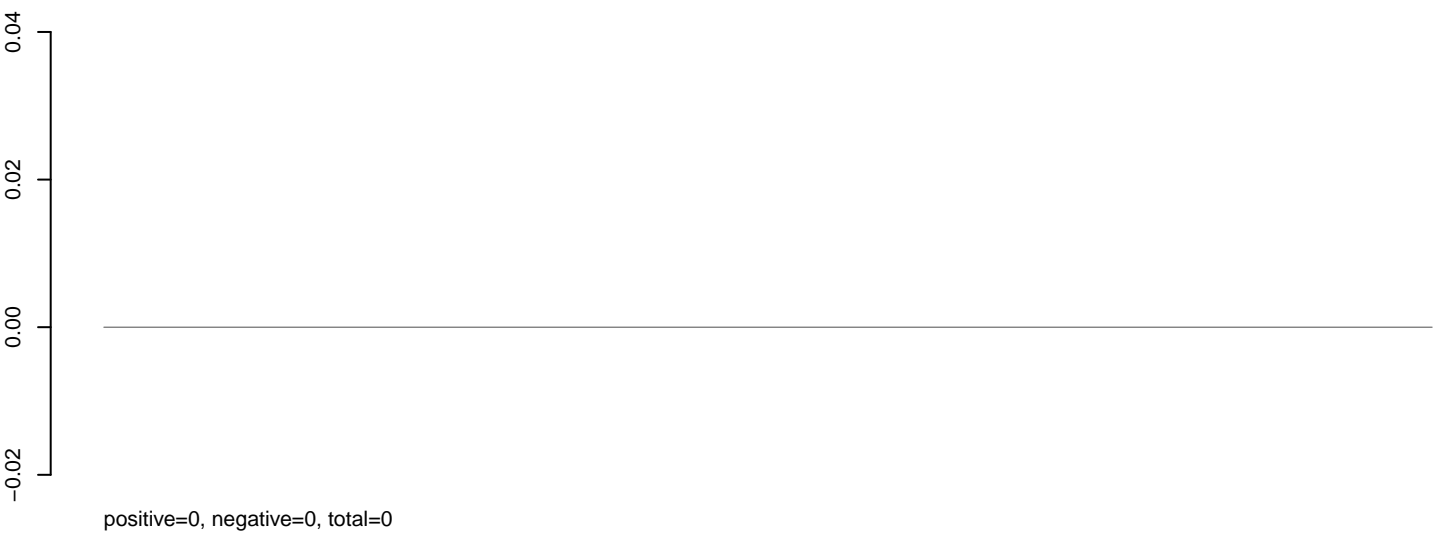
positive=0, negative=0, total=0
negative=0, positive=0, total=0
length=5982, virus@MF176317.1-Culex_mohonega-like_virus_1_strain_mos191X45127_segment_2:1-5982

0 1000 2000 3000 4000 5000 6000

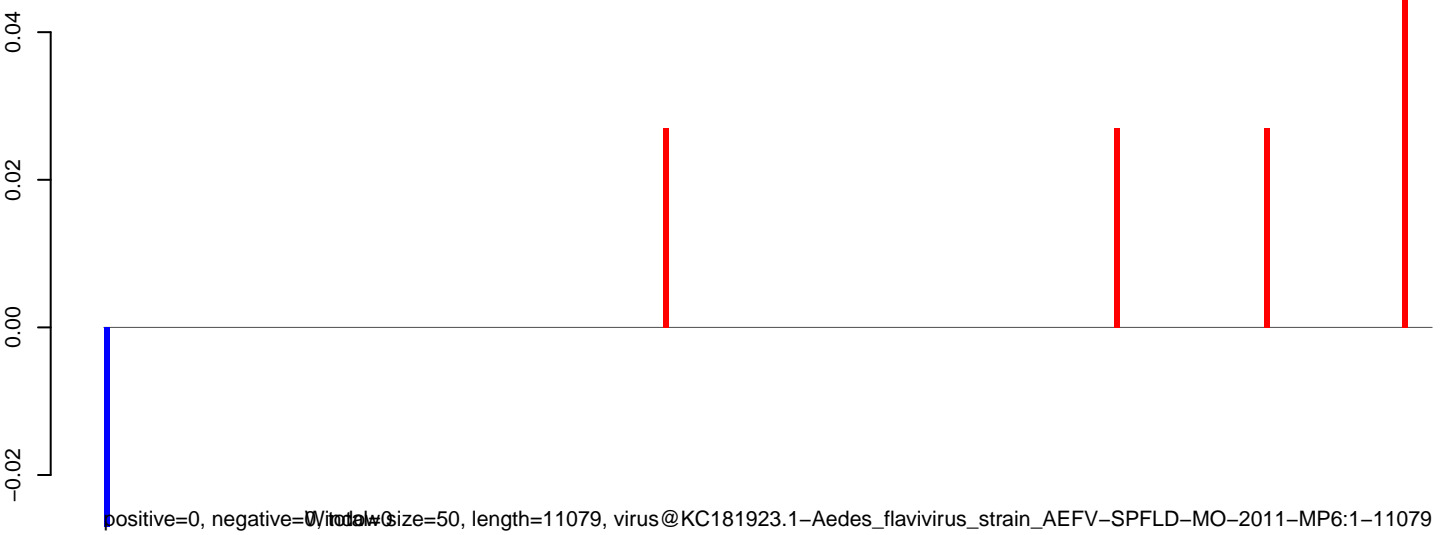
AeAeg_Ovary_GH.18_23.rep



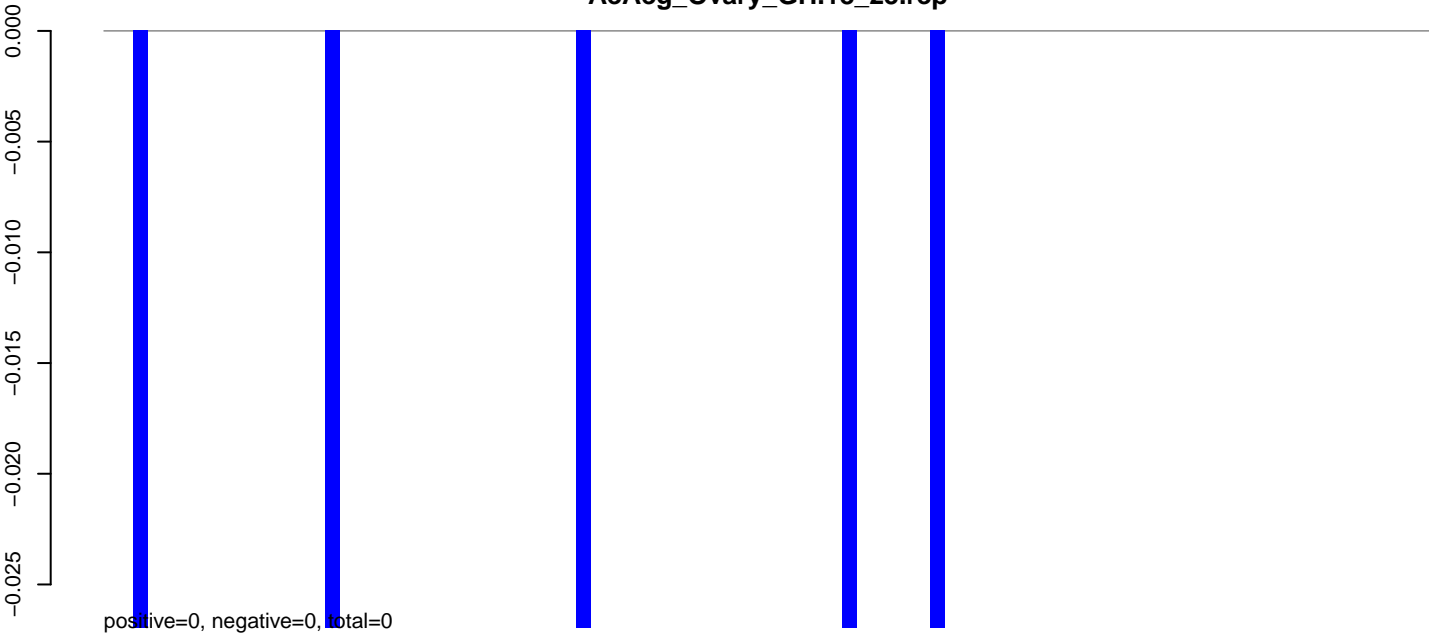
AeAeg_Ovary_GH.24_35.rep



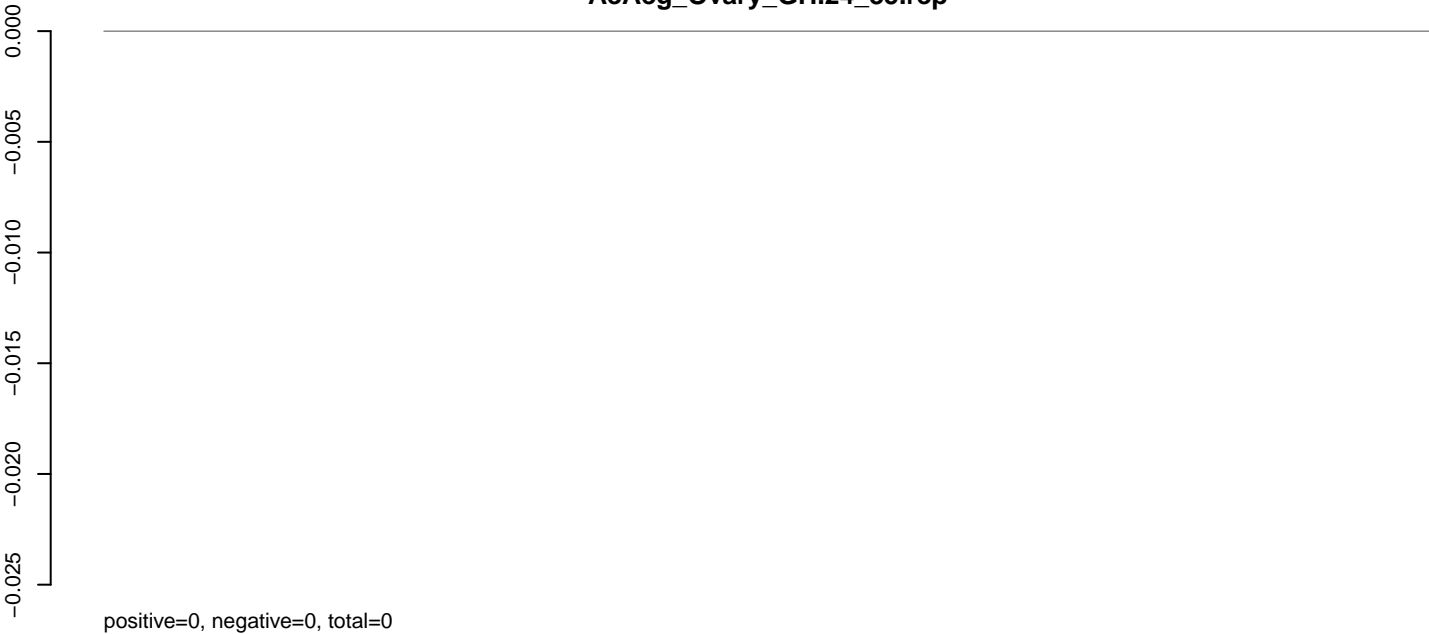
AeAeg_Ovary_GH.rep



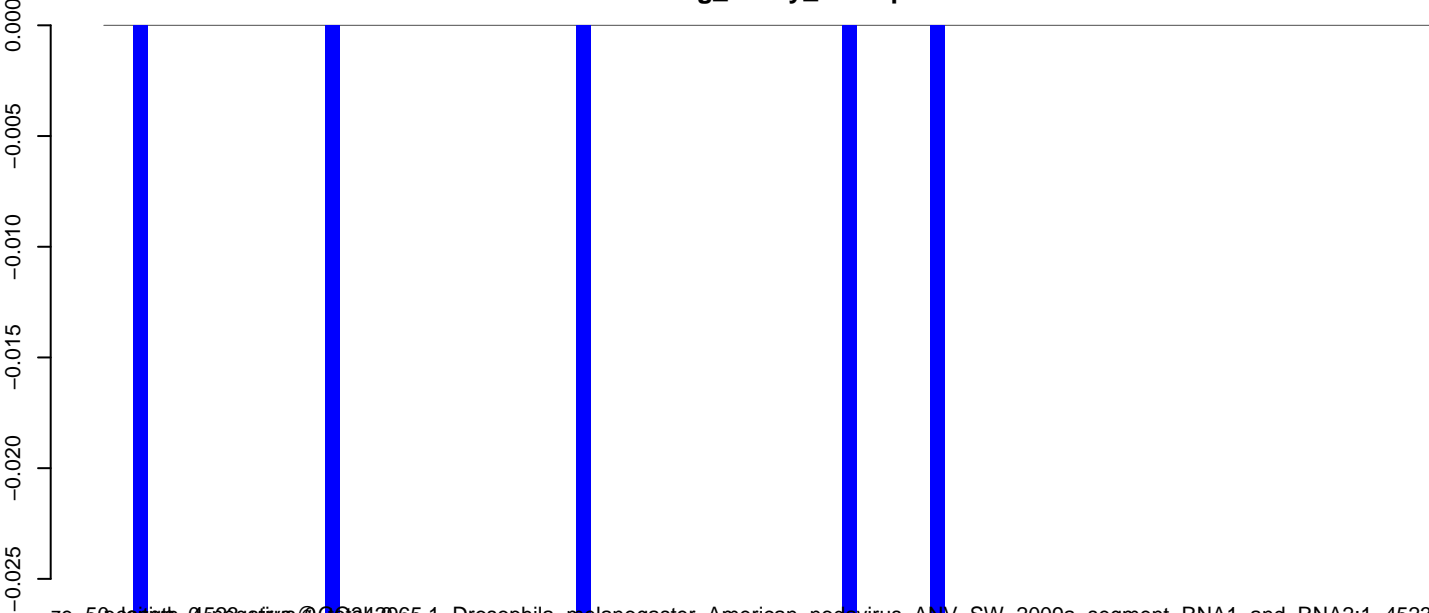
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



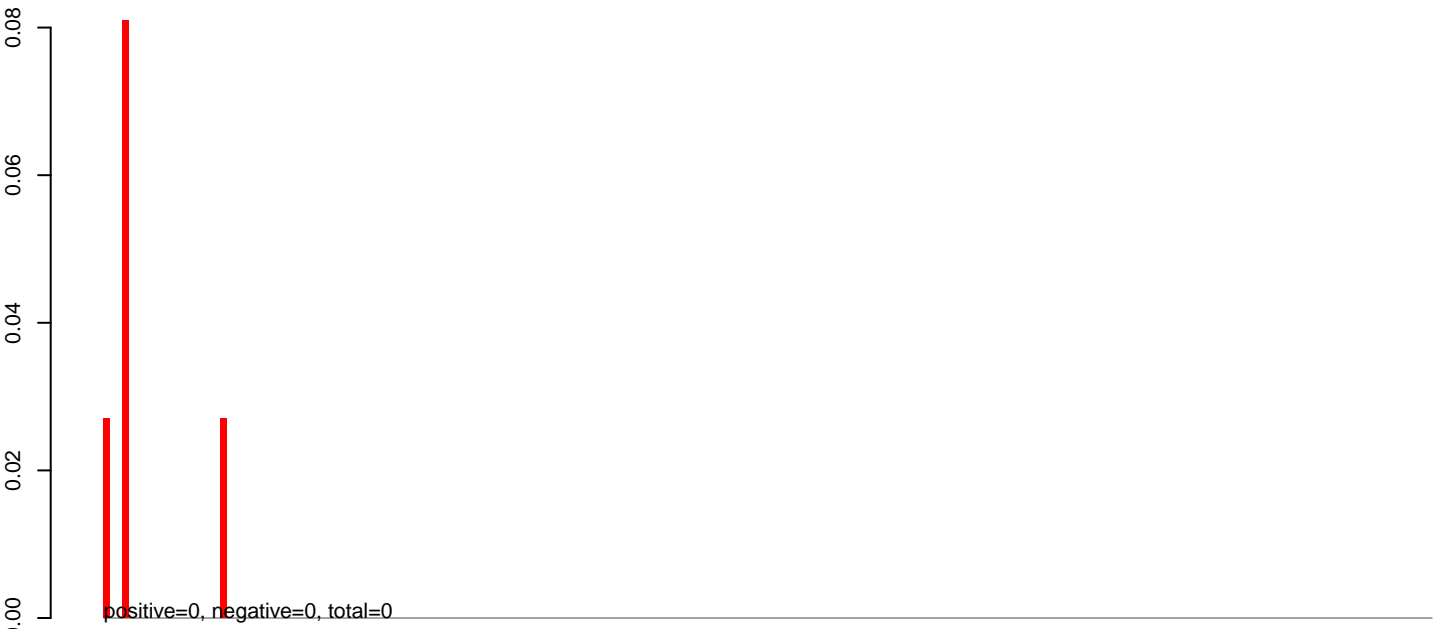
AeAeg_Ovary_GH.rep



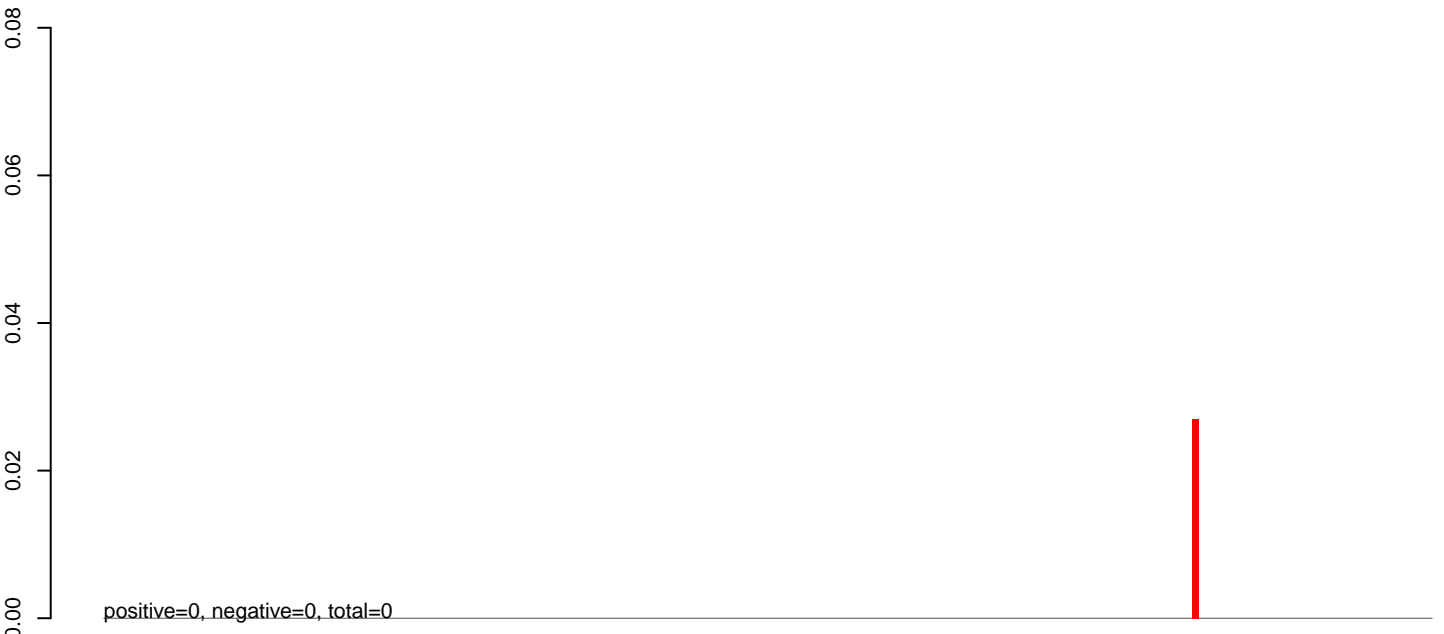
ze=50, depth=4,523, givis=0,03442, 65.1-Drosophila_melanogaster_American_nodavirus_ANV_SW-2009a_segment_RNA1_and_RNA2:1-4523

0 1000 2000 3000 4000

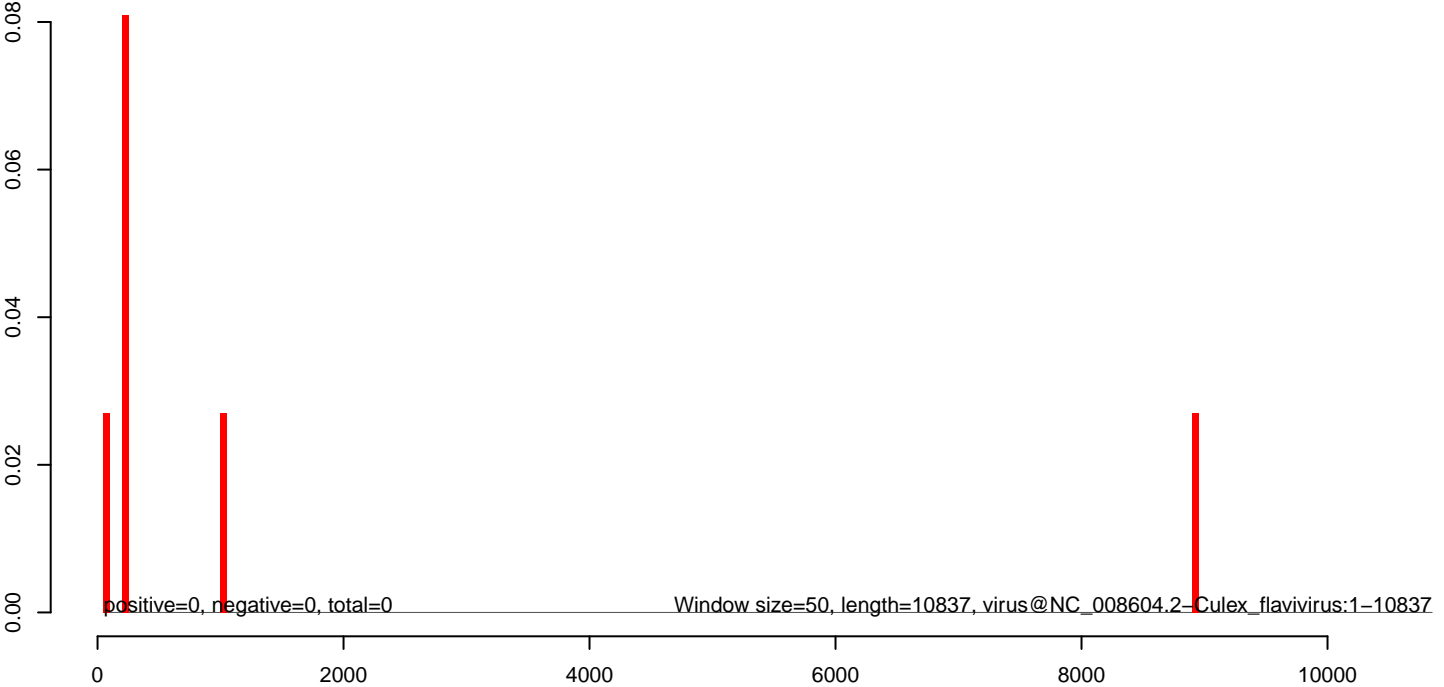
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



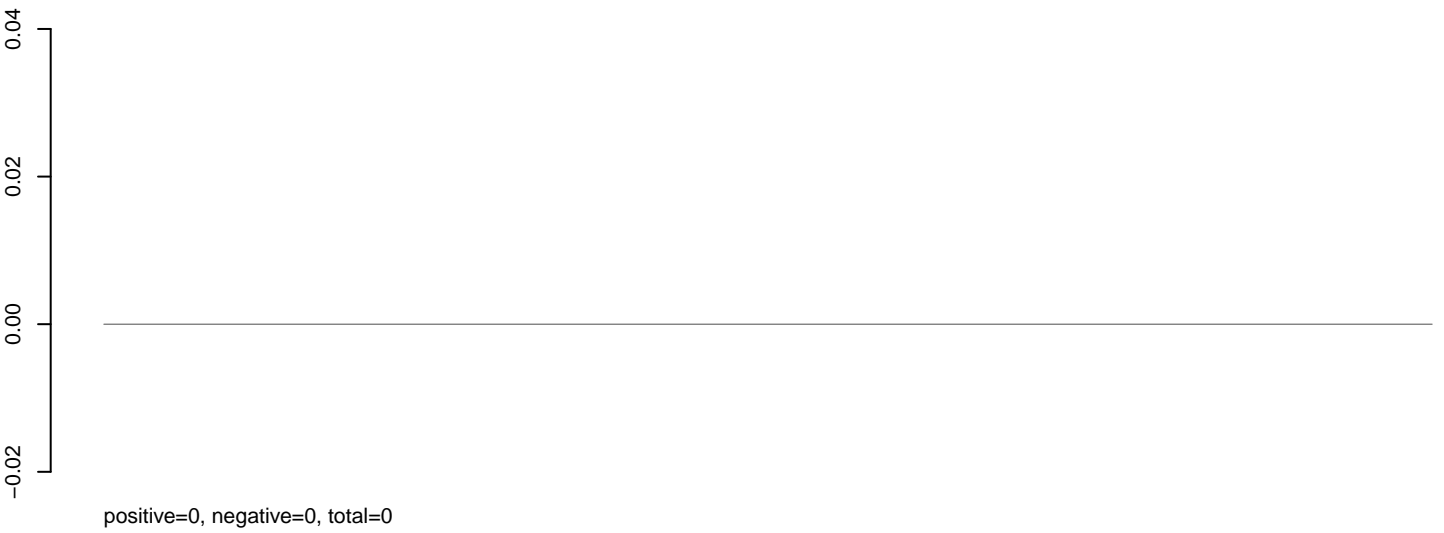
AeAeg_Ovary_GH.rep



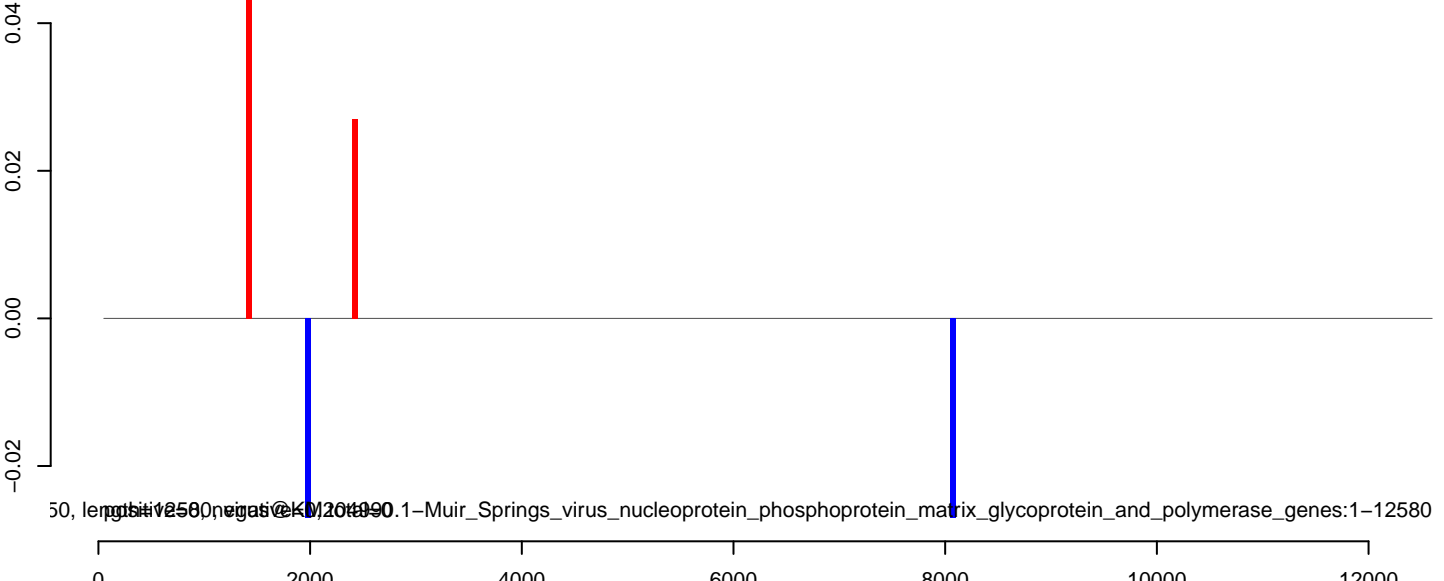
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

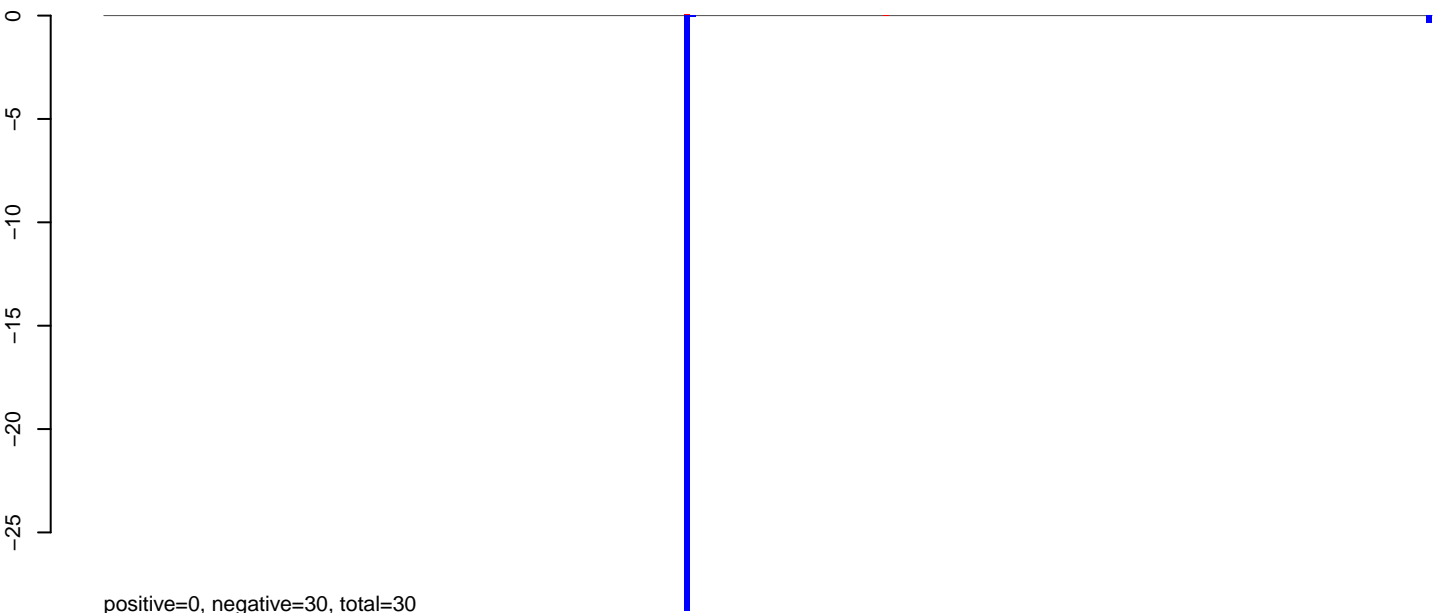


AeAeg_Ovary_GH.rep

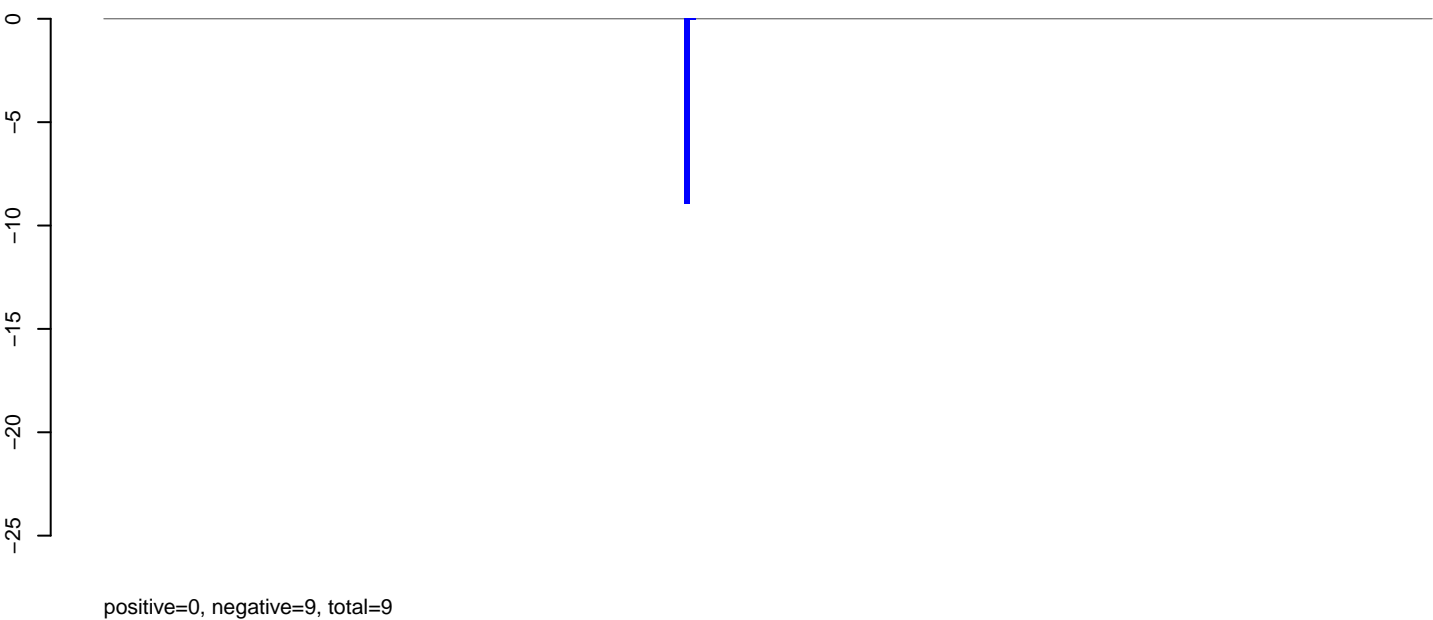


50, length=12580, negative=4, total=90.1-Muir_Springs_virus_nucleoprotein_phosphoprotein_matrix_glycoprotein_and_polymerase_genes:1-12580

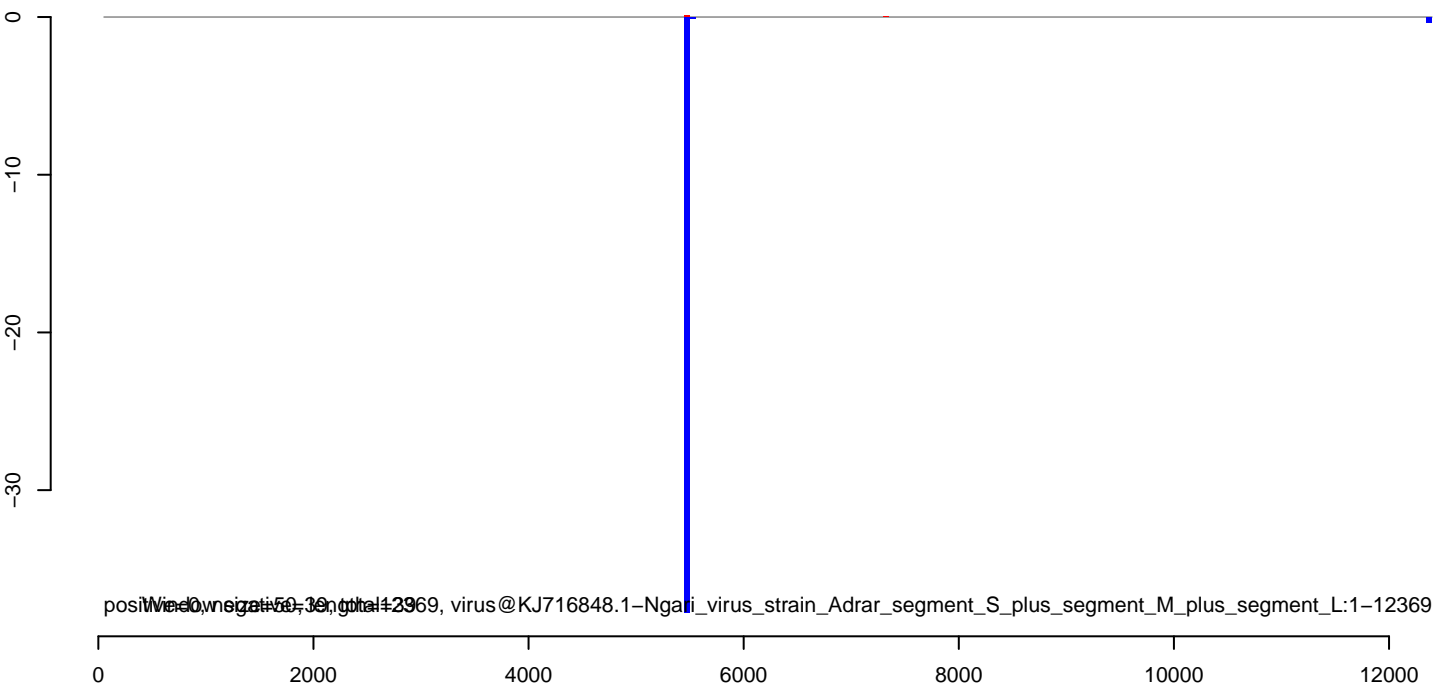
AeAeg_Ovary_GH.18_23.rep



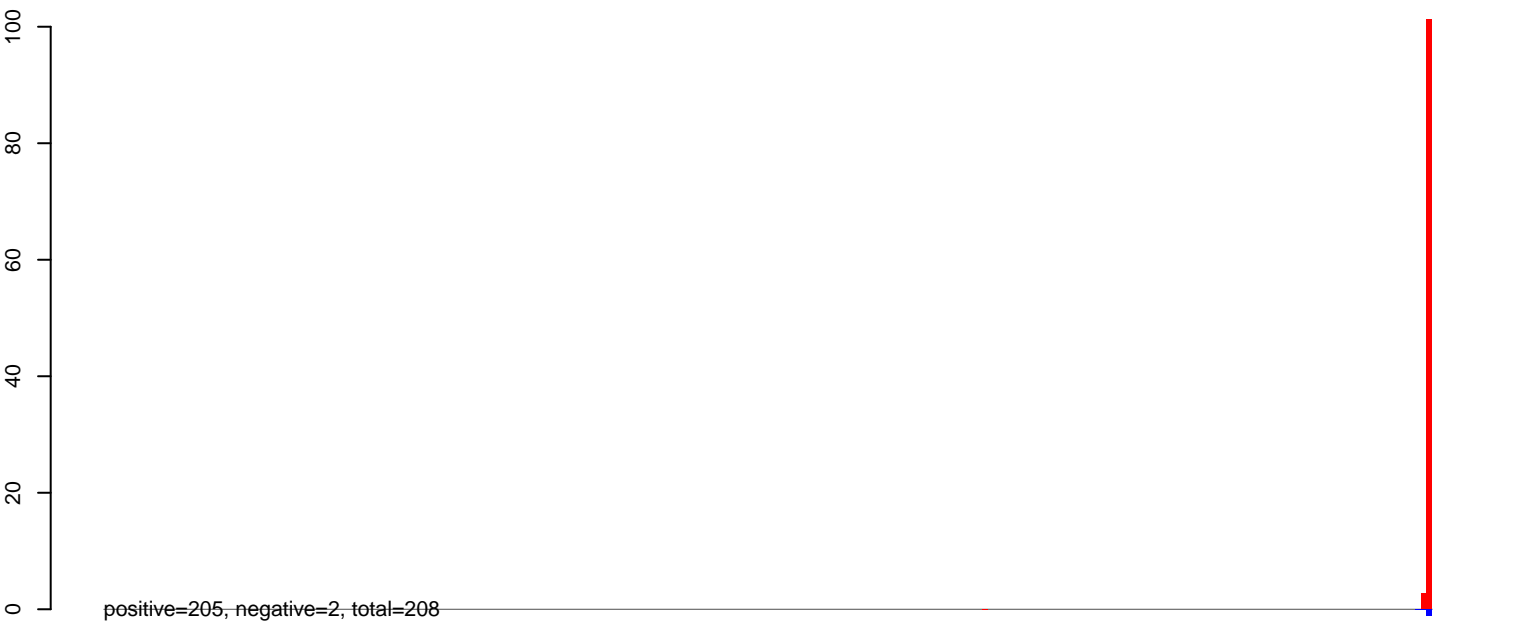
AeAeg_Ovary_GH.24_35.rep



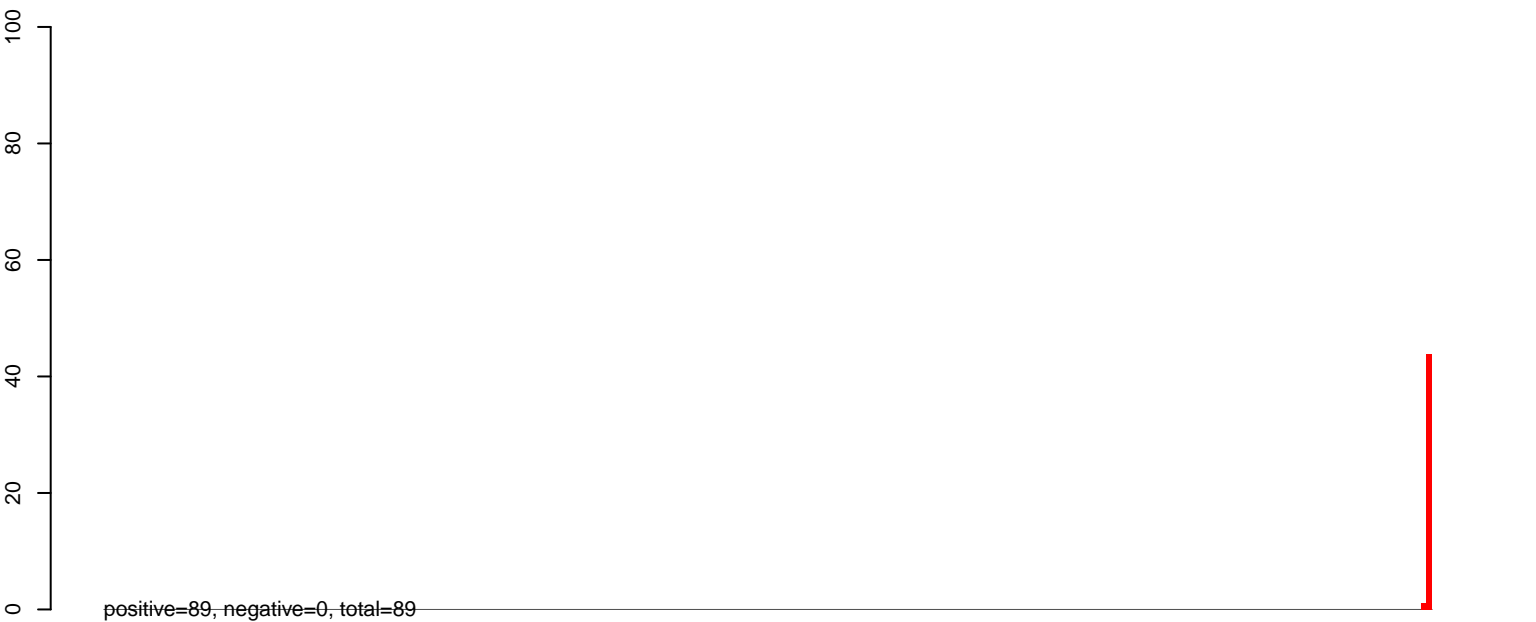
AeAeg_Ovary_GH.rep



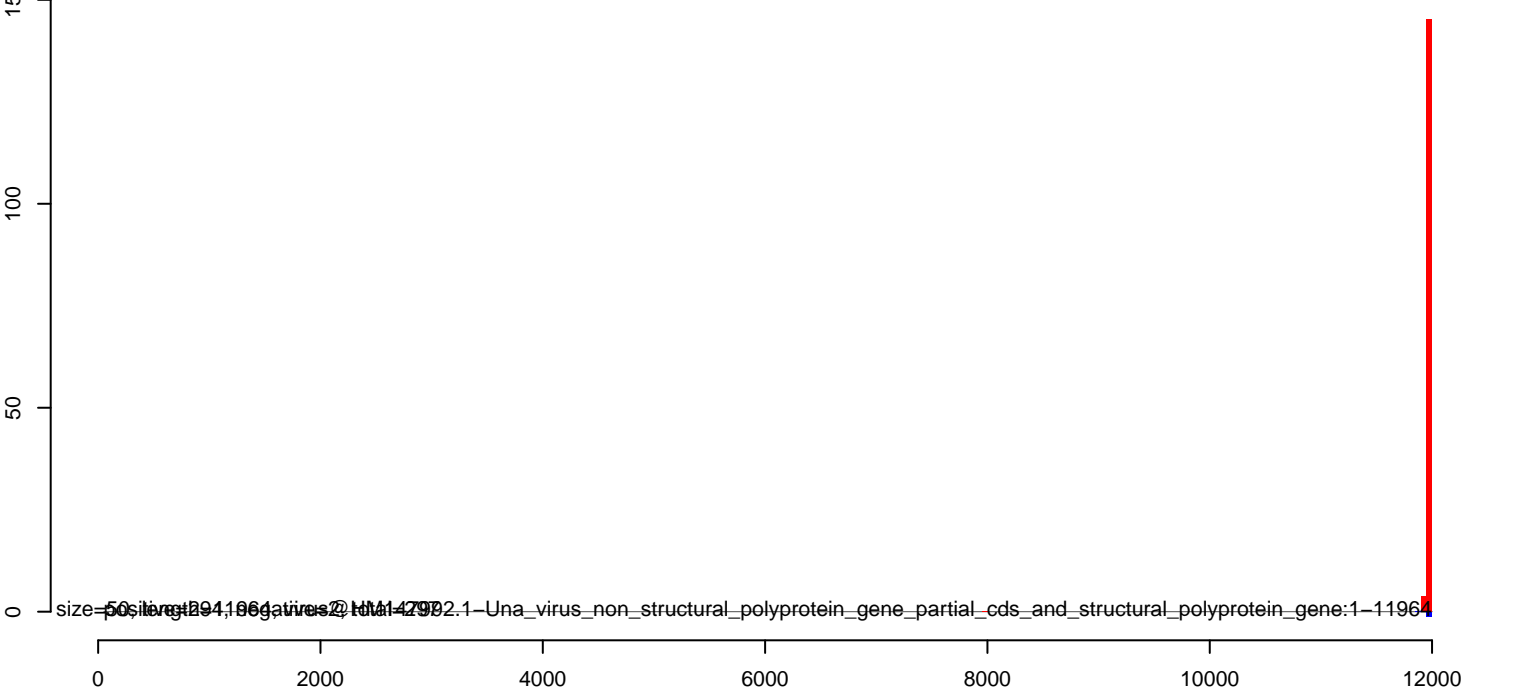
AeAeg_Ovary_GH.18_23.rep



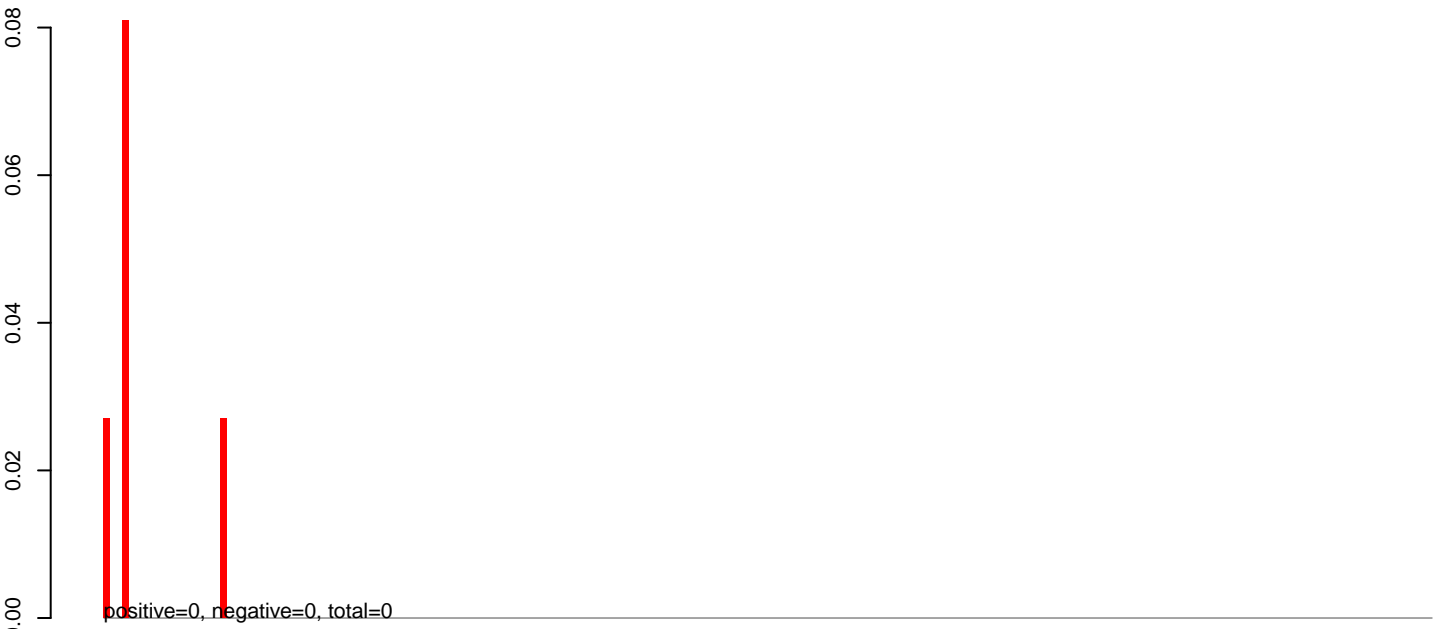
AeAeg_Ovary_GH.24_35.rep



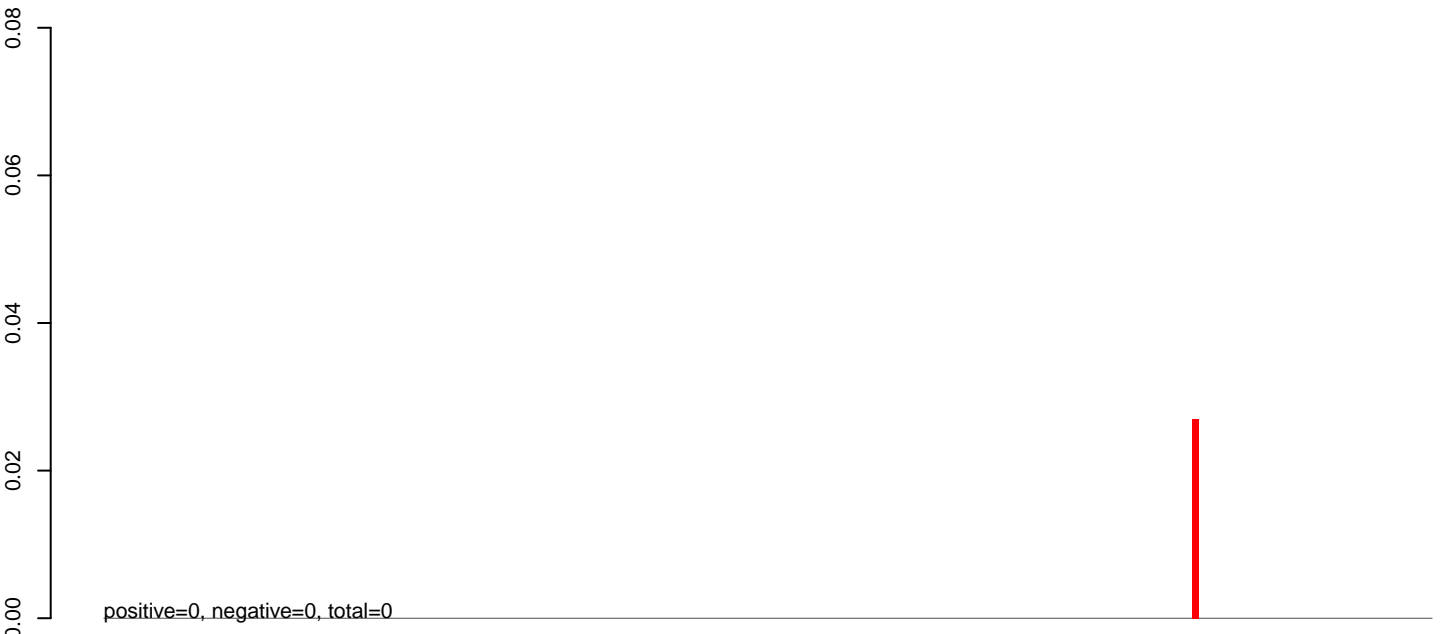
AeAeg_Ovary_GH.rep



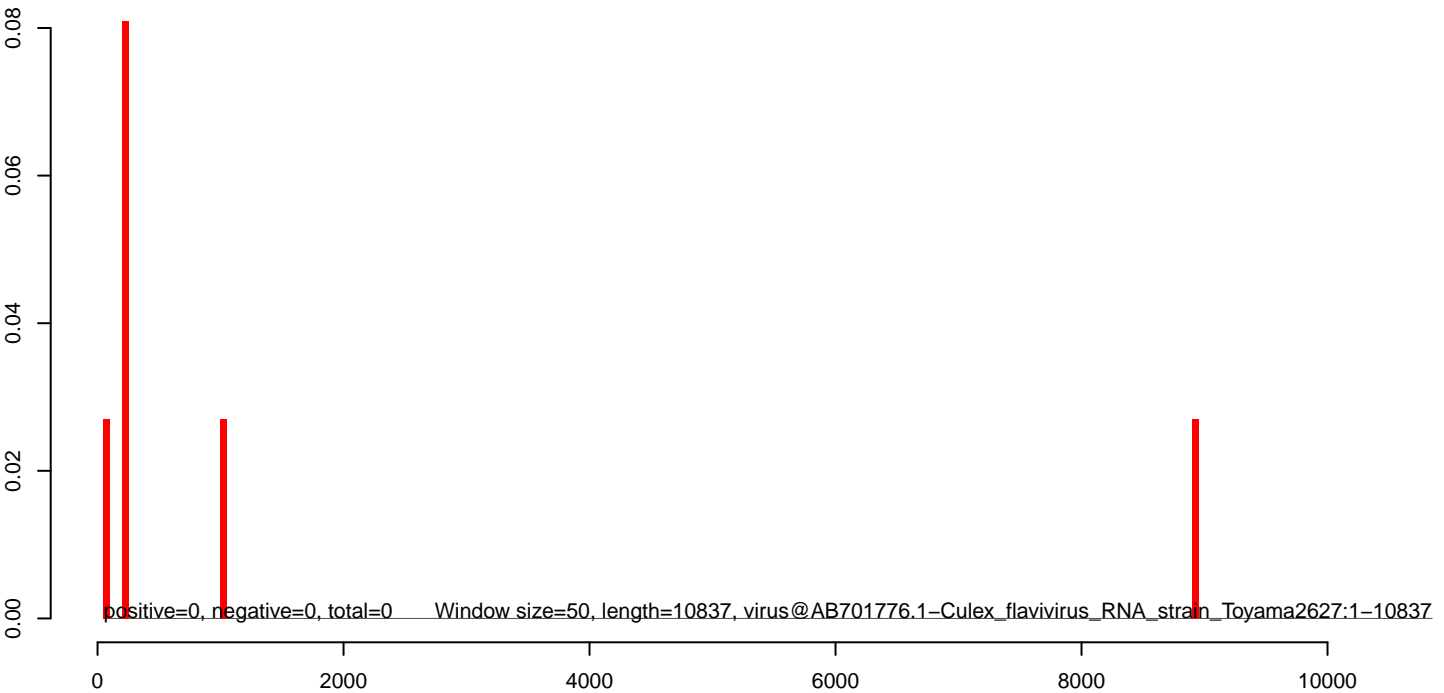
AeAeg_Ovary_GH.18_23.rep



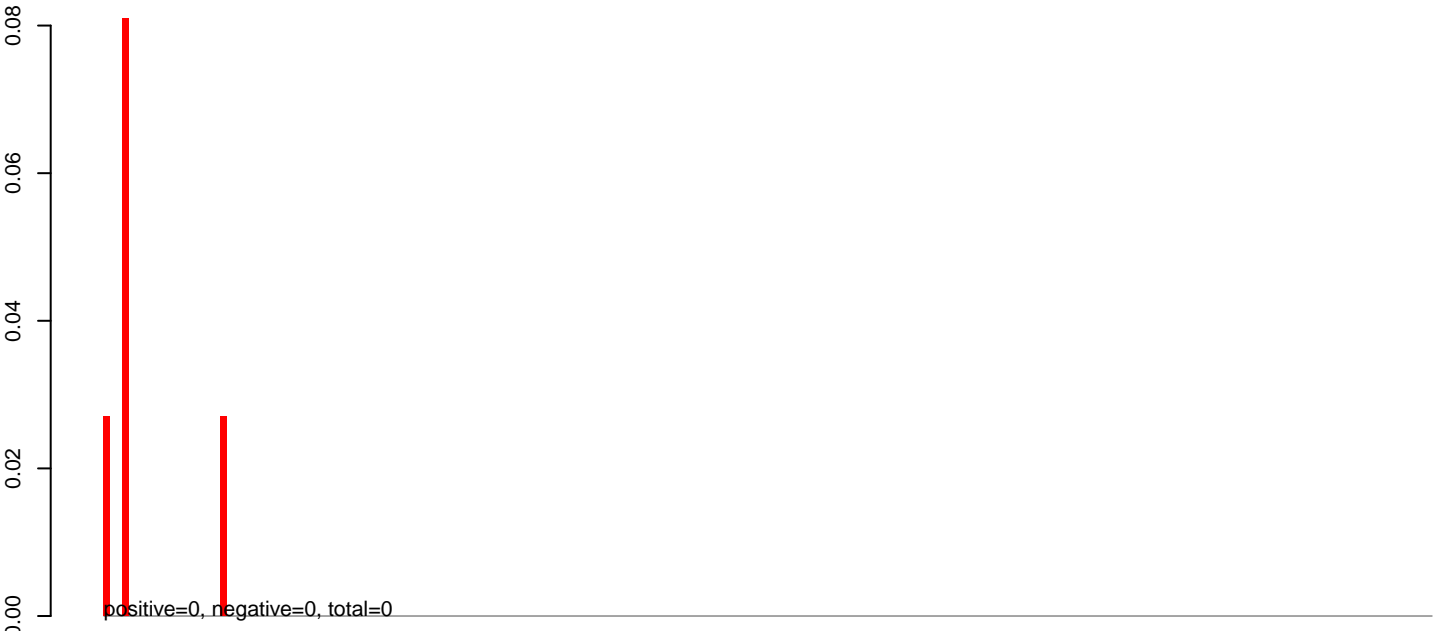
AeAeg_Ovary_GH.24_35.rep



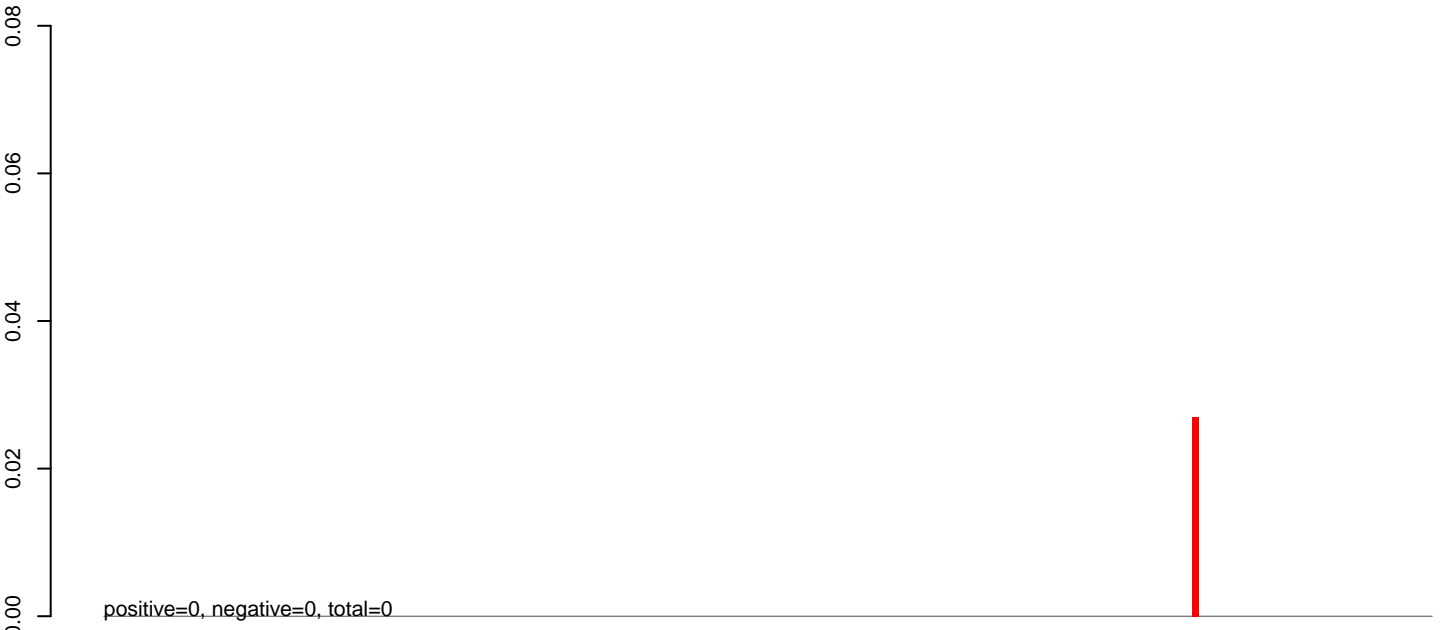
AeAeg_Ovary_GH.rep



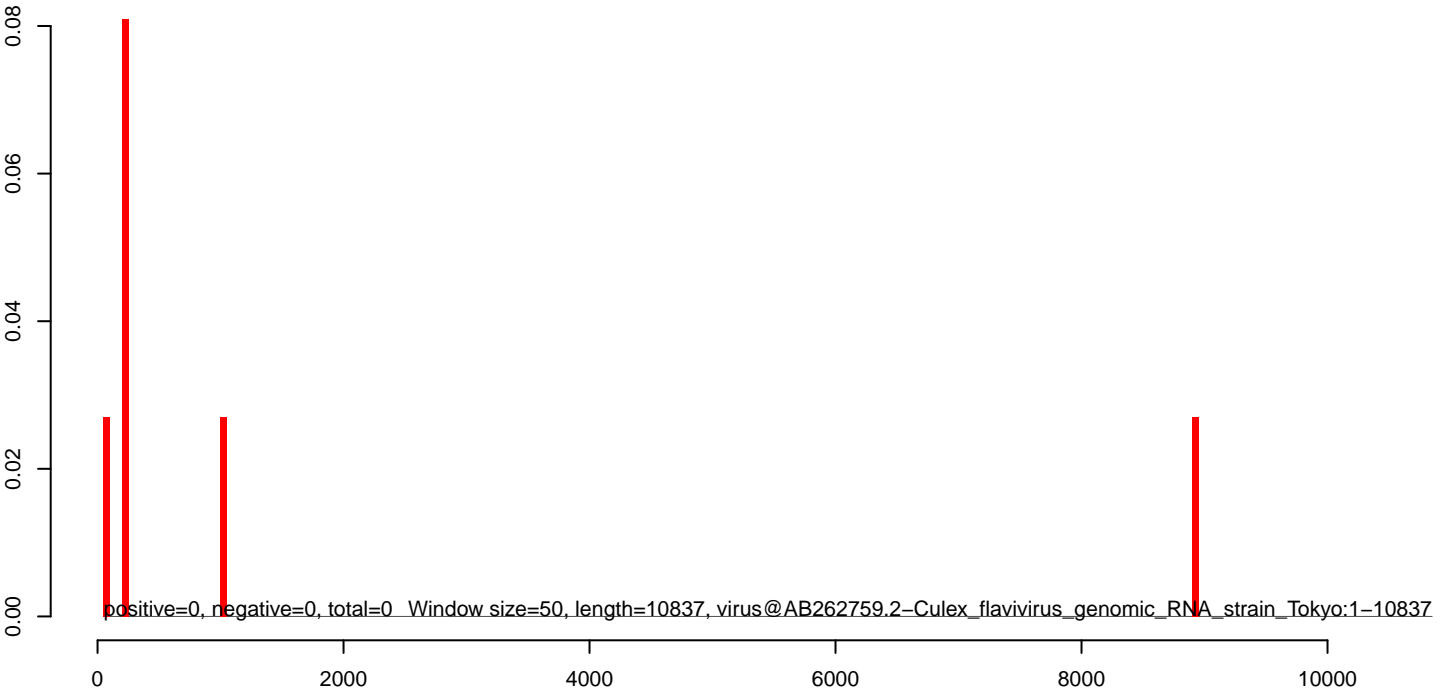
AeAeg_Ovary_GH.18_23.rep



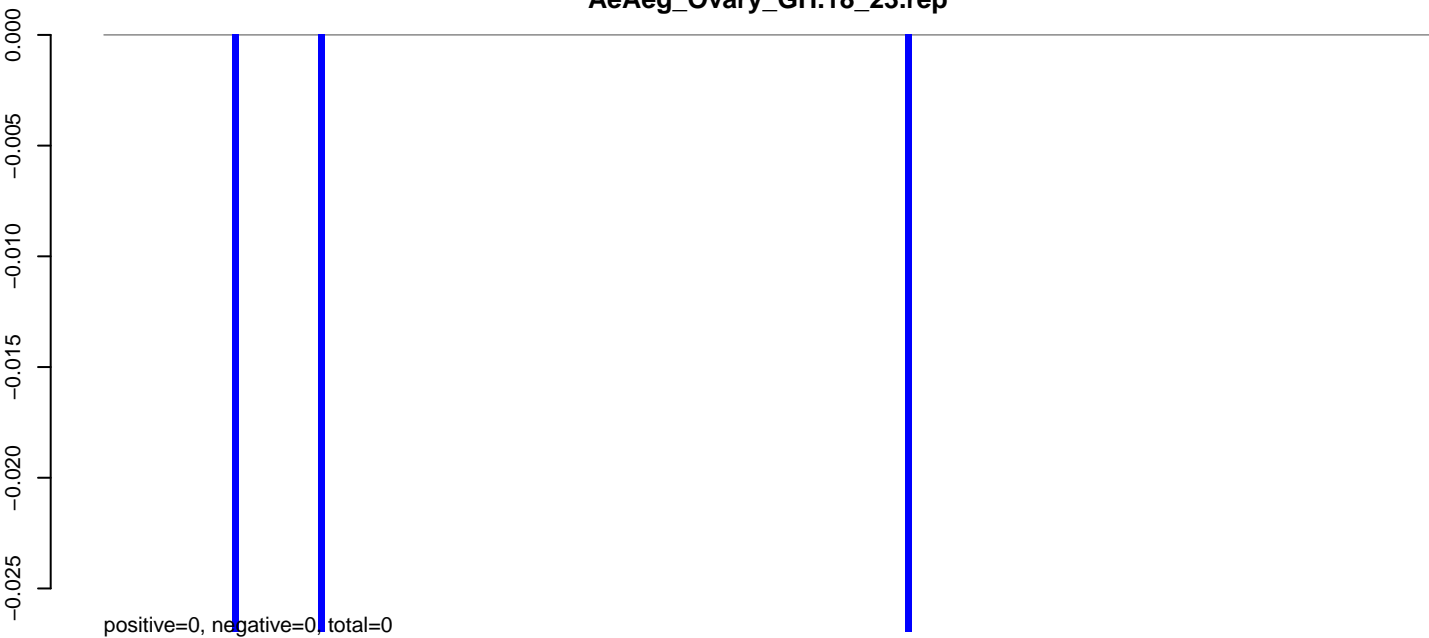
AeAeg_Ovary_GH.24_35.rep



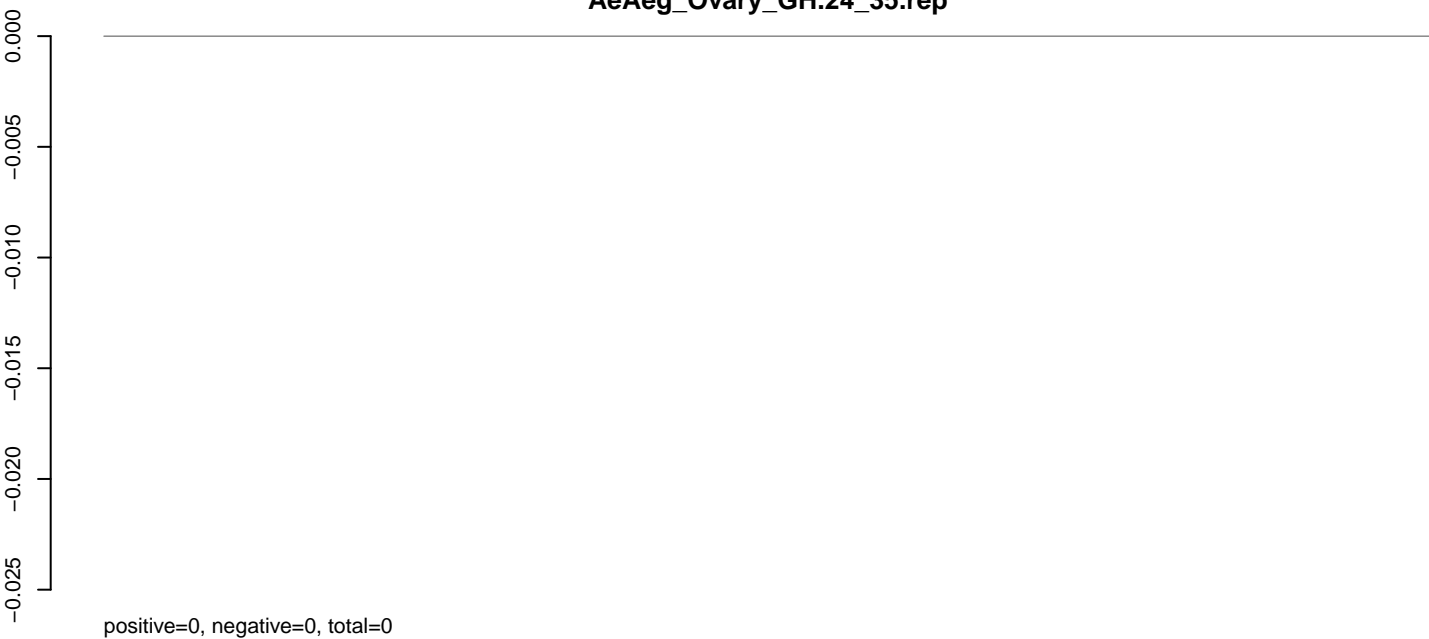
AeAeg_Ovary_GH.rep



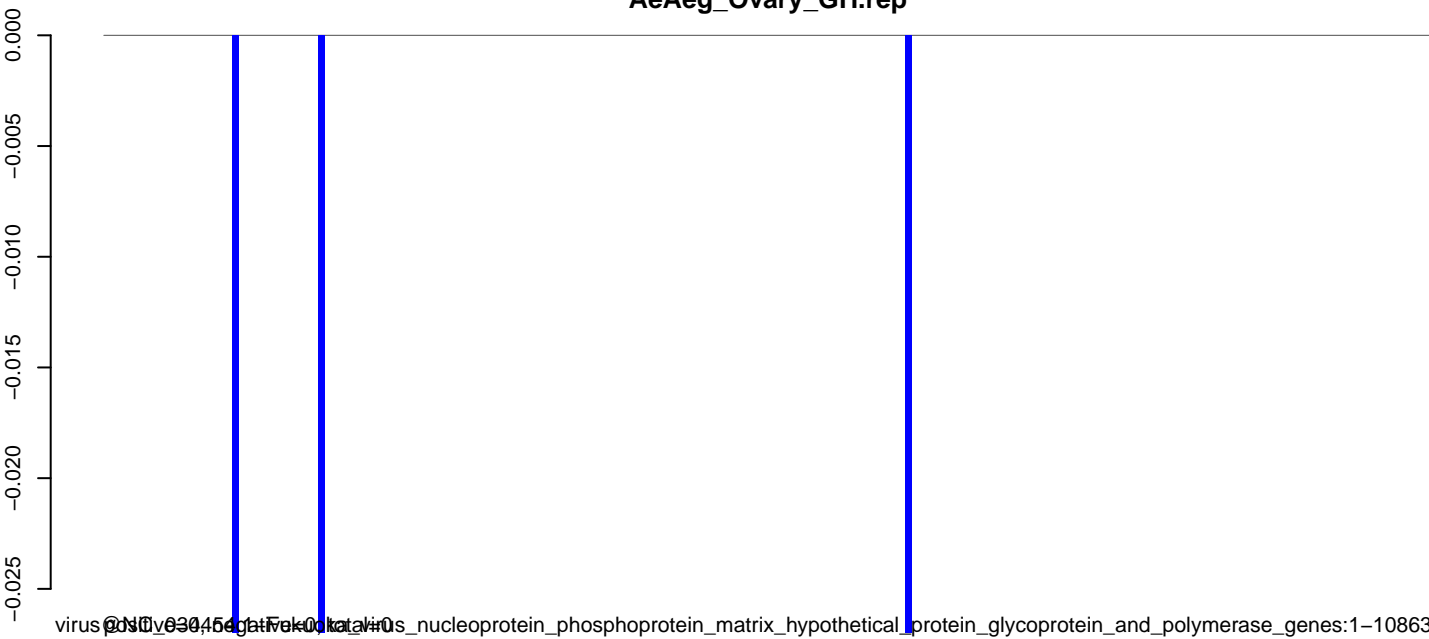
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

AeAeg_Ovary_GH.18_23.rep

positive=1, negative=0, total=1

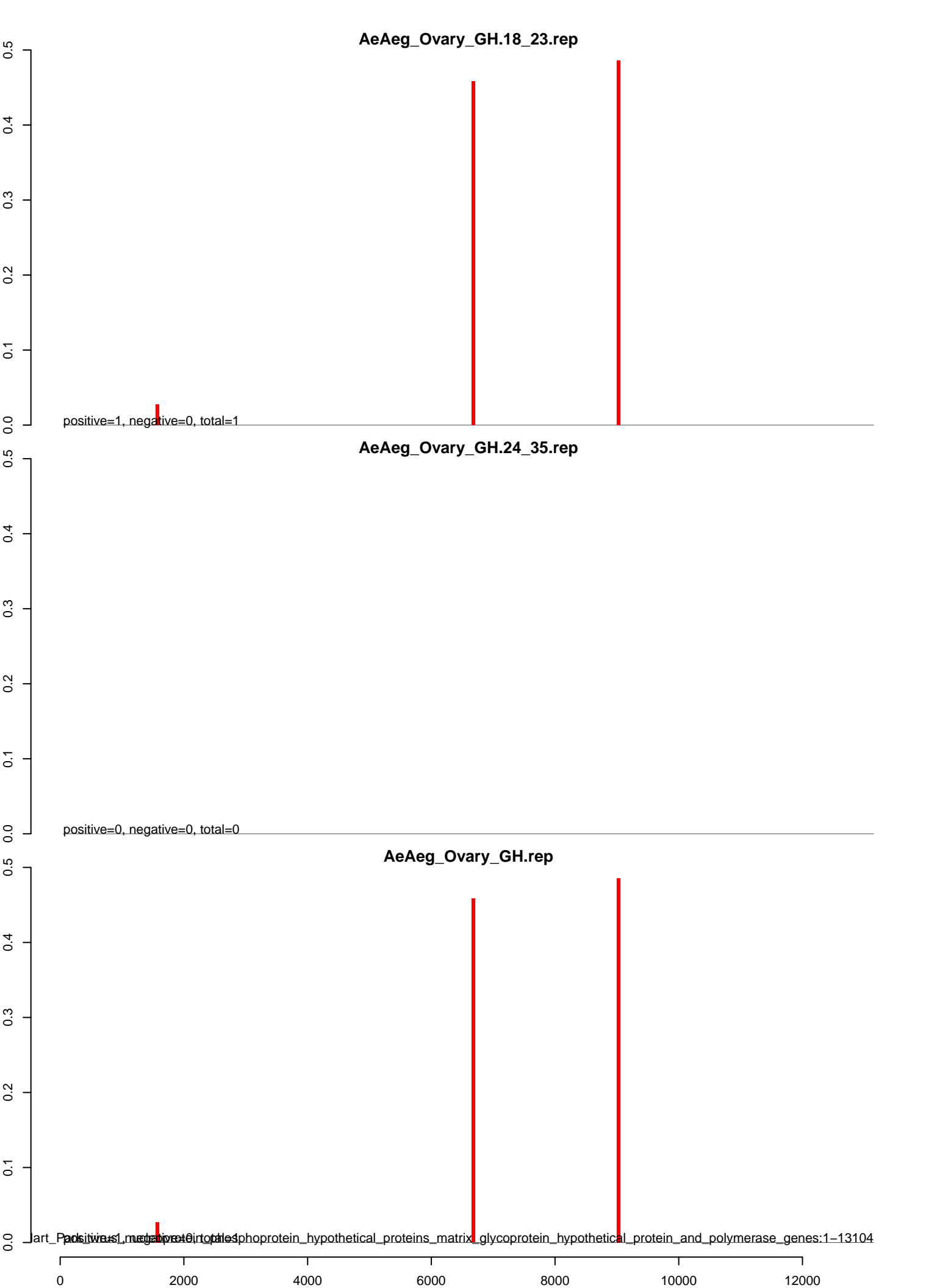
AeAeg_Ovary_GH.24_35.rep

positive=0, negative=0, total=0

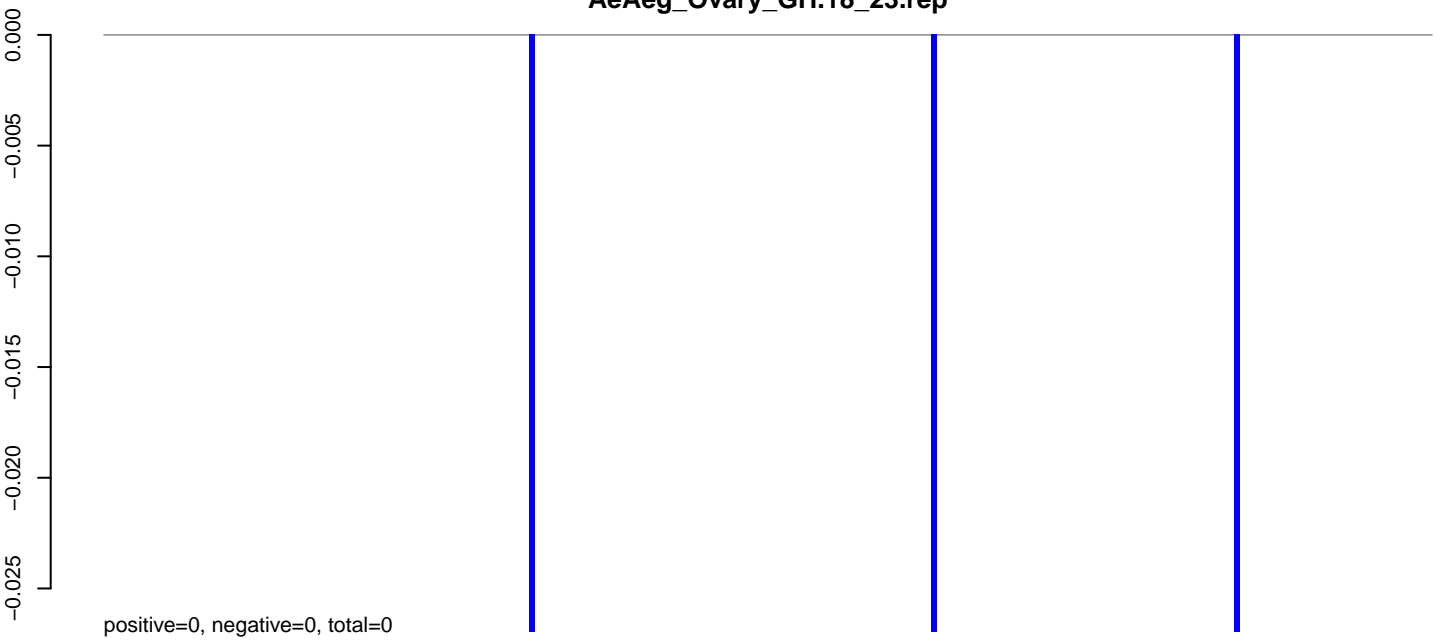
AeAeg_Ovary_GH.rep

part_positive=1, negative=0, total=1
phosphoprotein_hypothetical_proteins_matrix_glycoprotein_hypothetical_protein_and_polymerase_genes:1-13104

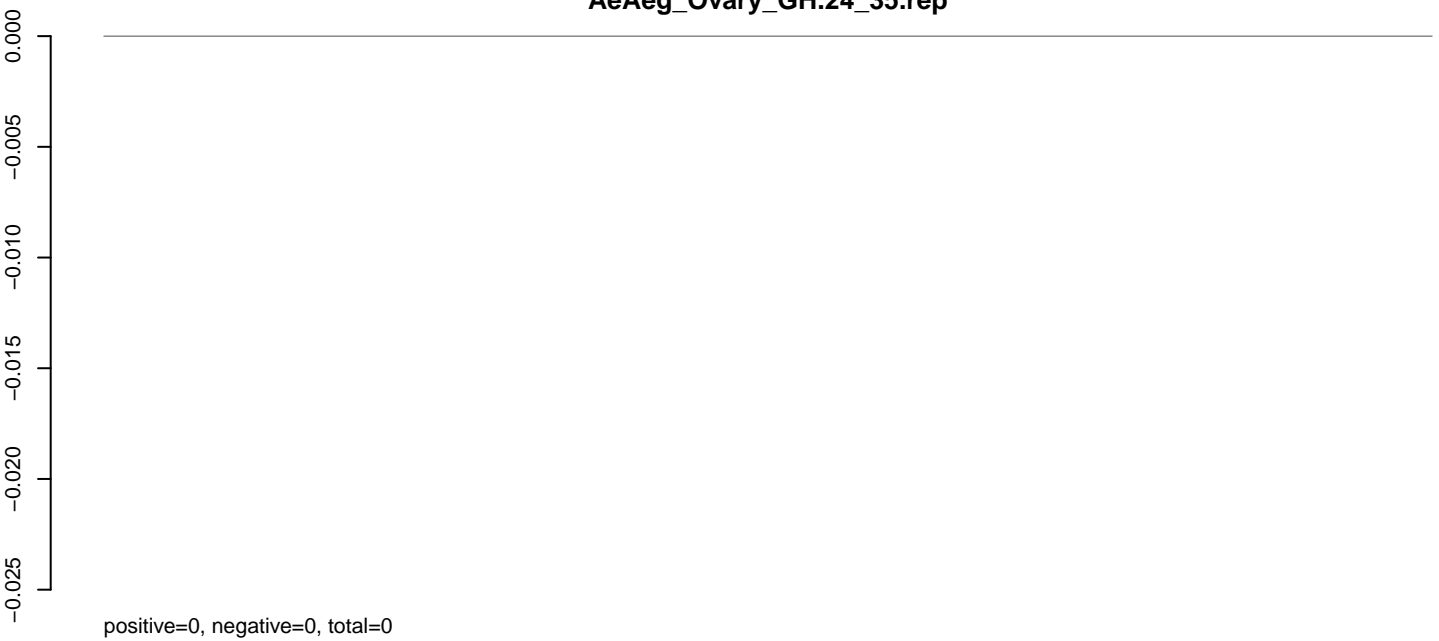
0 2000 4000 6000 8000 10000 12000



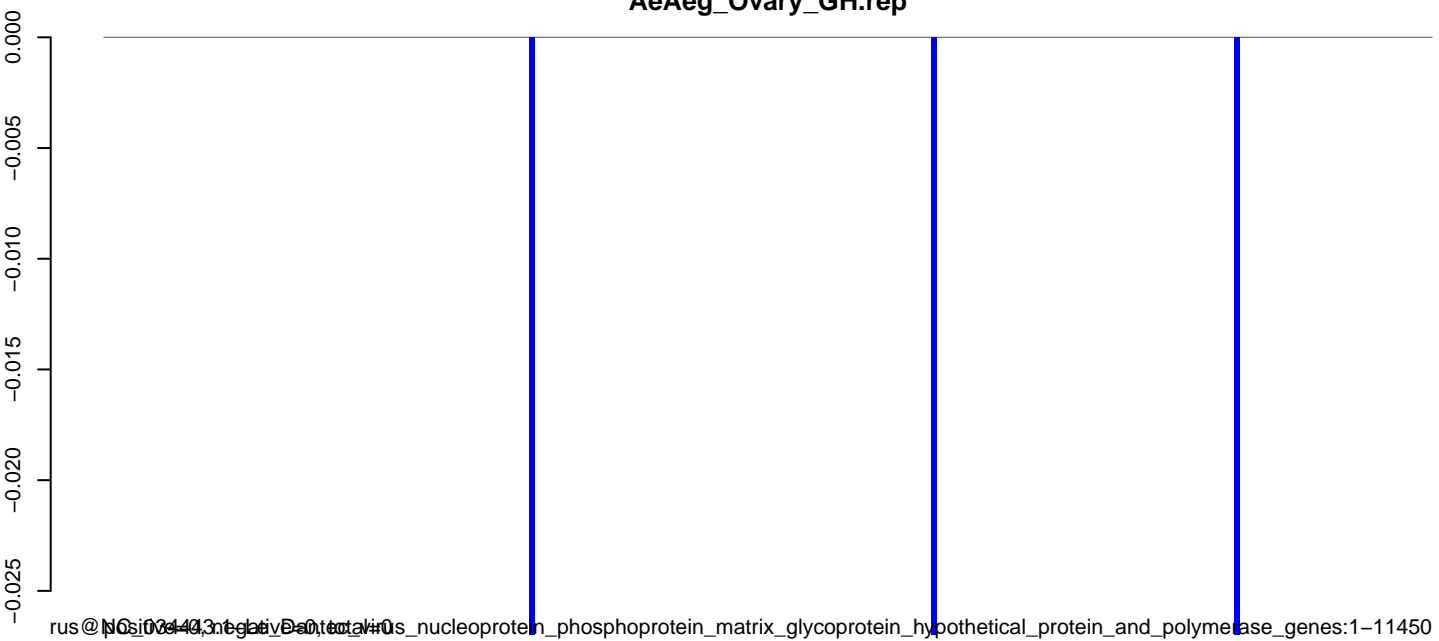
AeAeg_Ovary_GH.18_23.rep



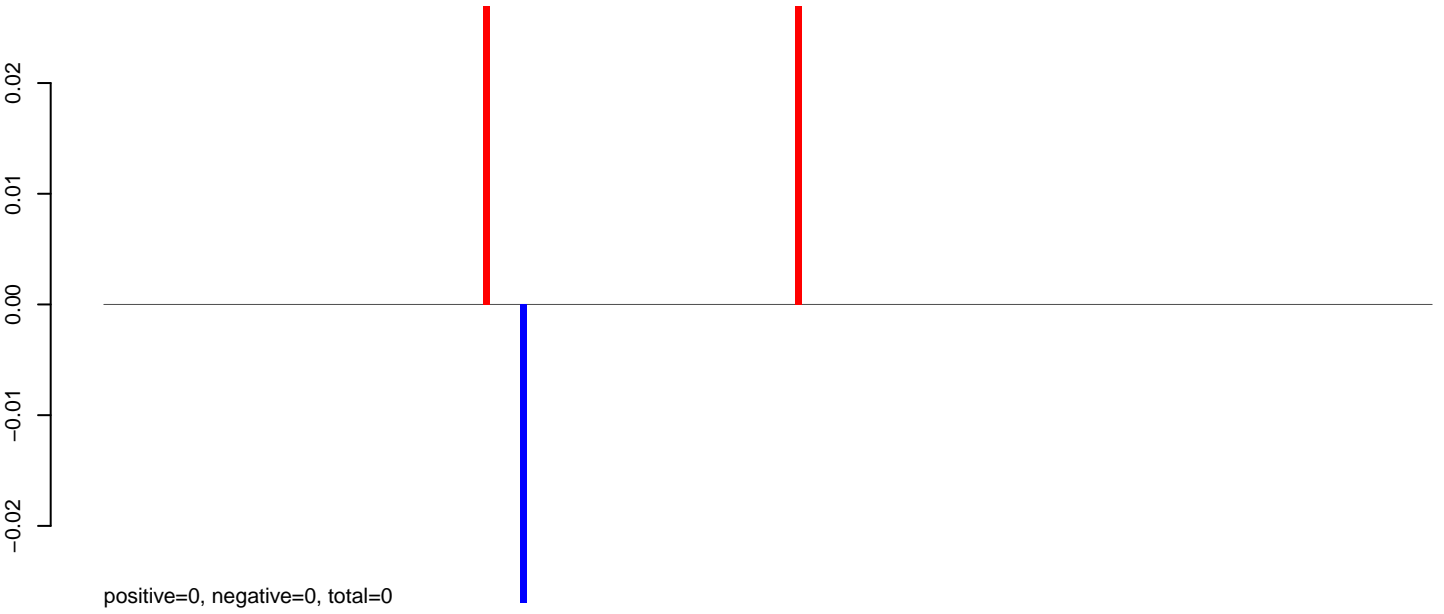
AeAeg_Ovary_GH.24_35.rep



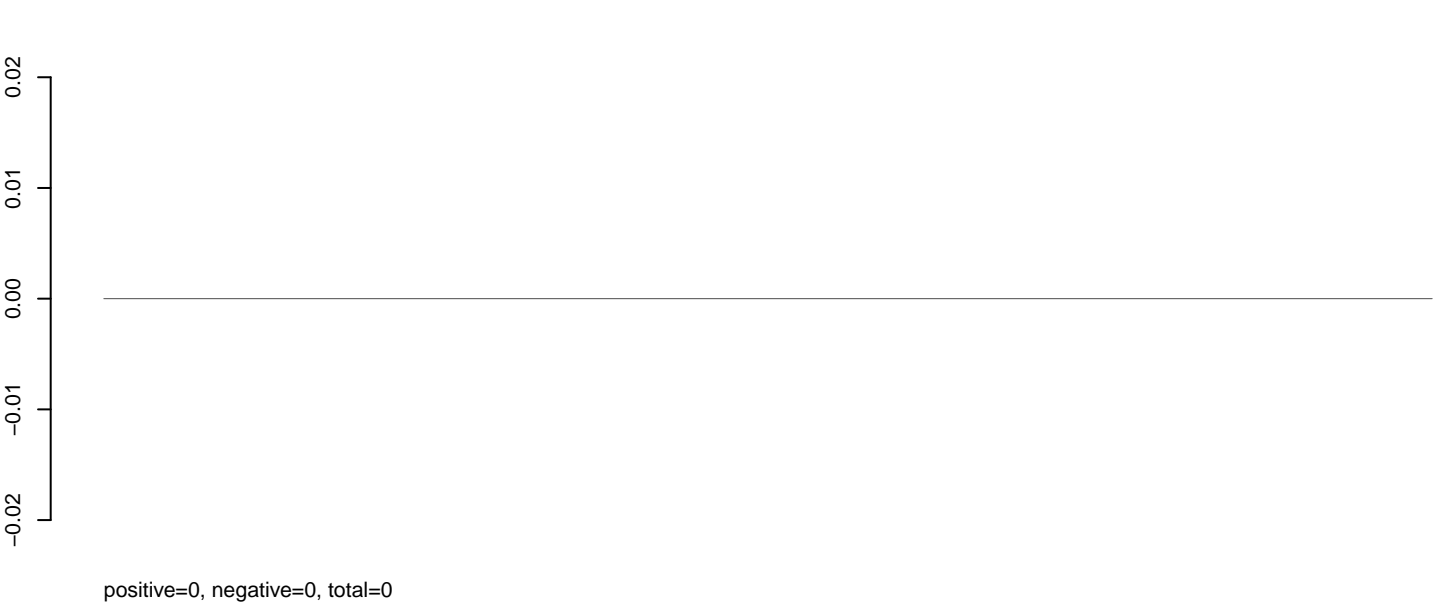
AeAeg_Ovary_GH.rep



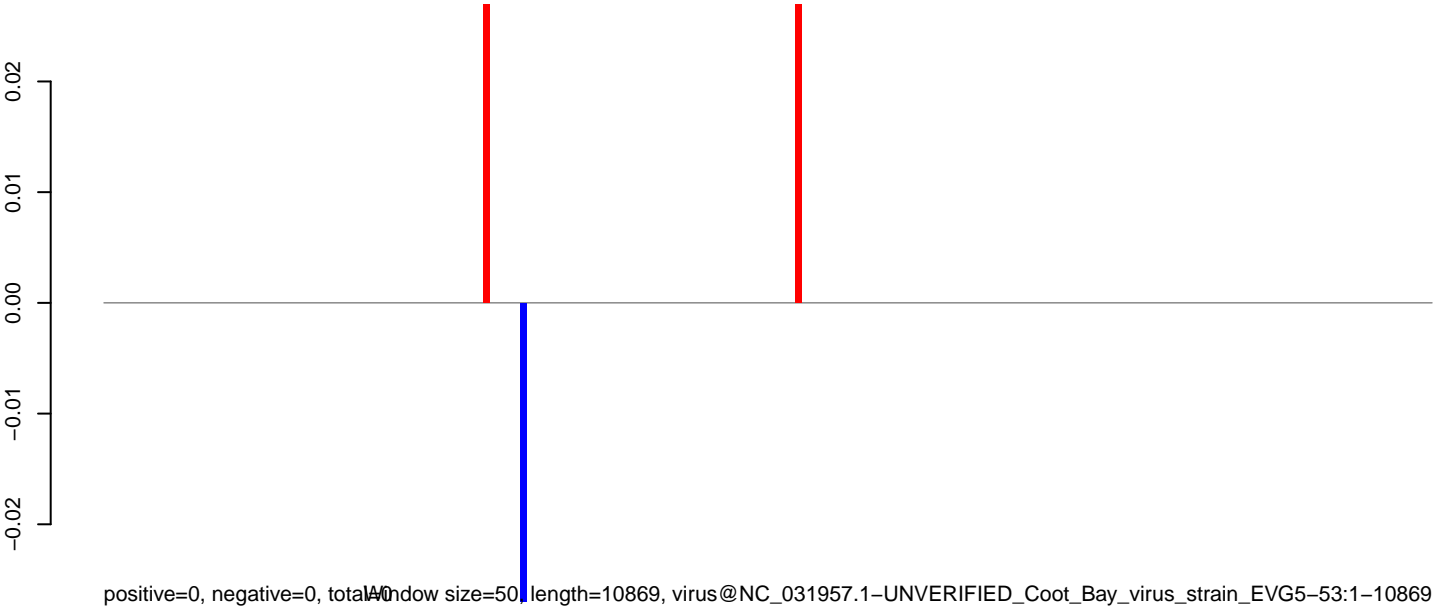
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



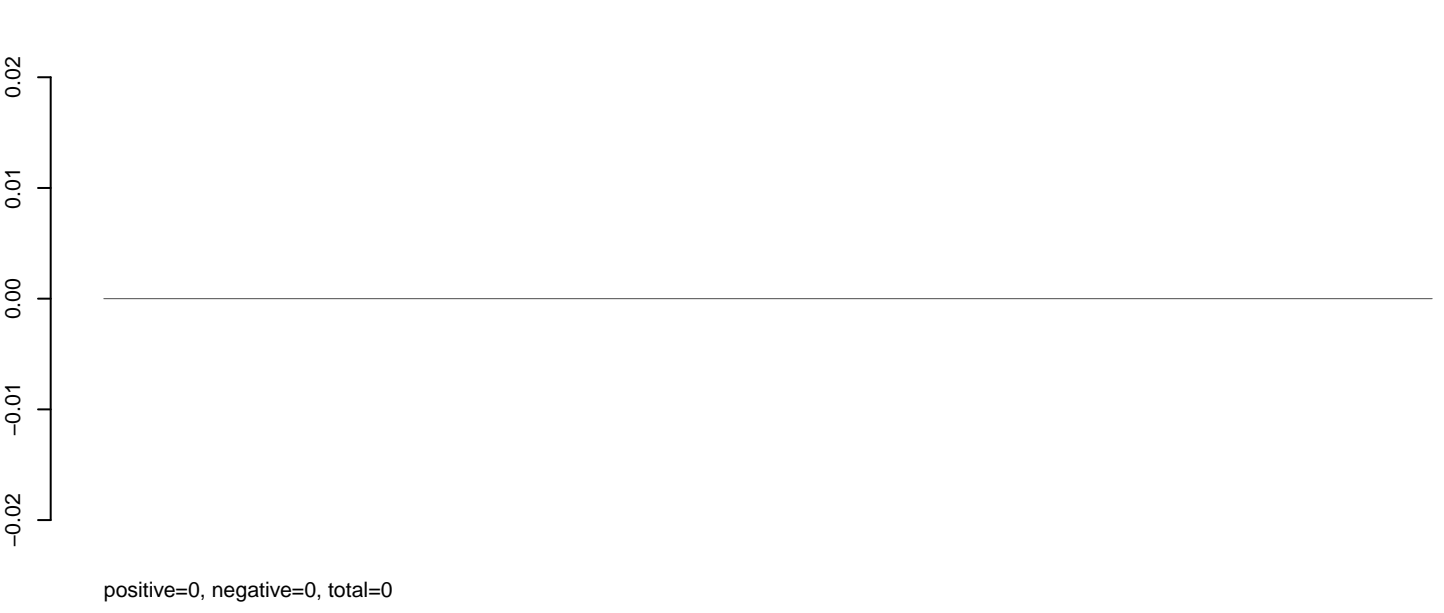
Window size=50, length=10869, virus@NC_031957.1-UNVERIFIED_Coot_Bay_virus_strain_EVG5-53:1-10869

0 2000 4000 6000 8000 10000

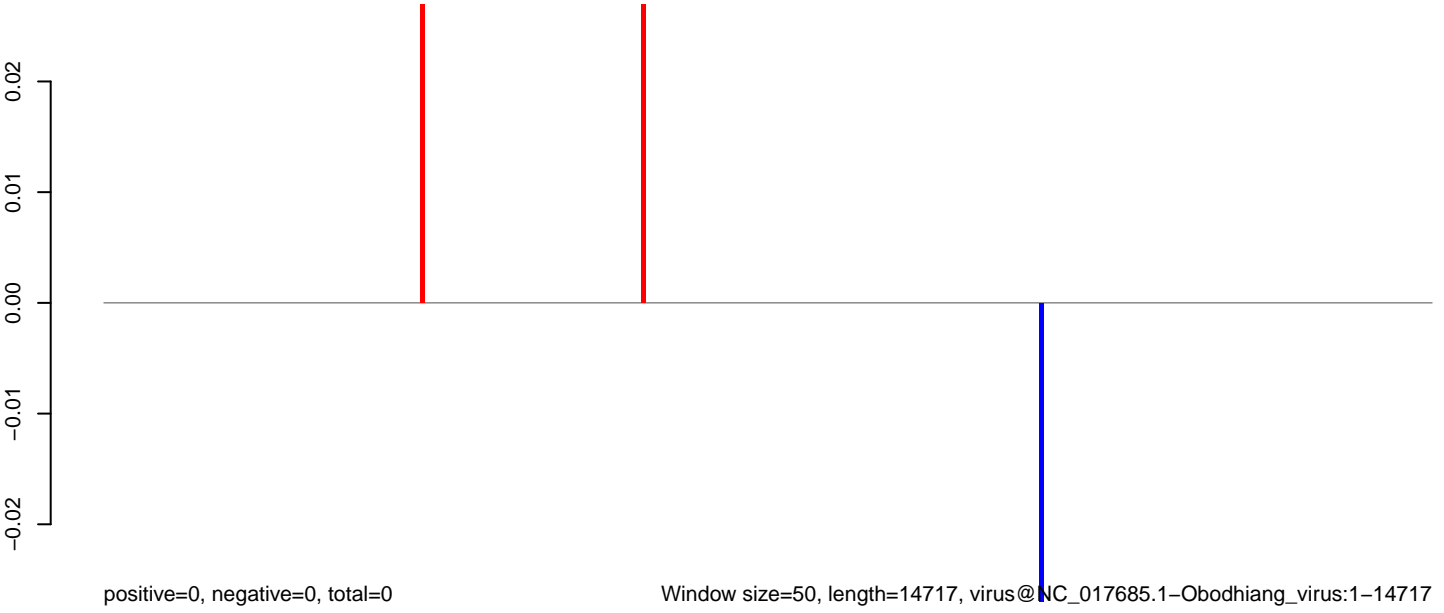
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



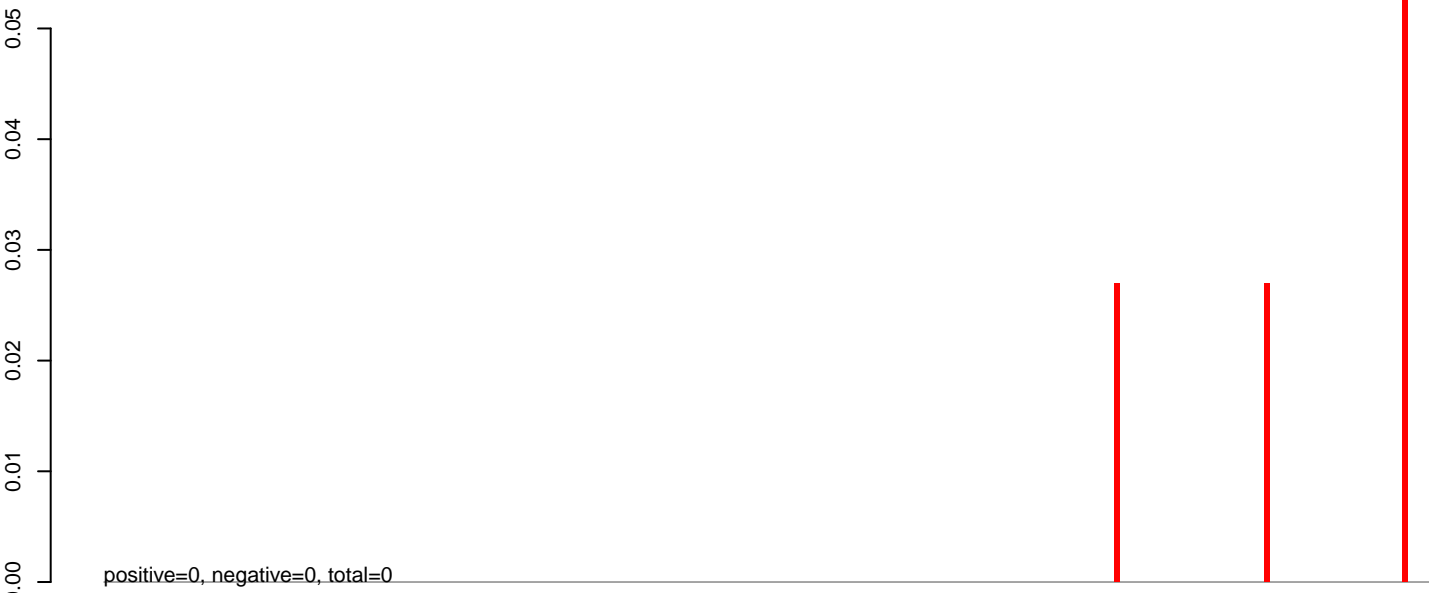
AeAeg_Ovary_GH.rep



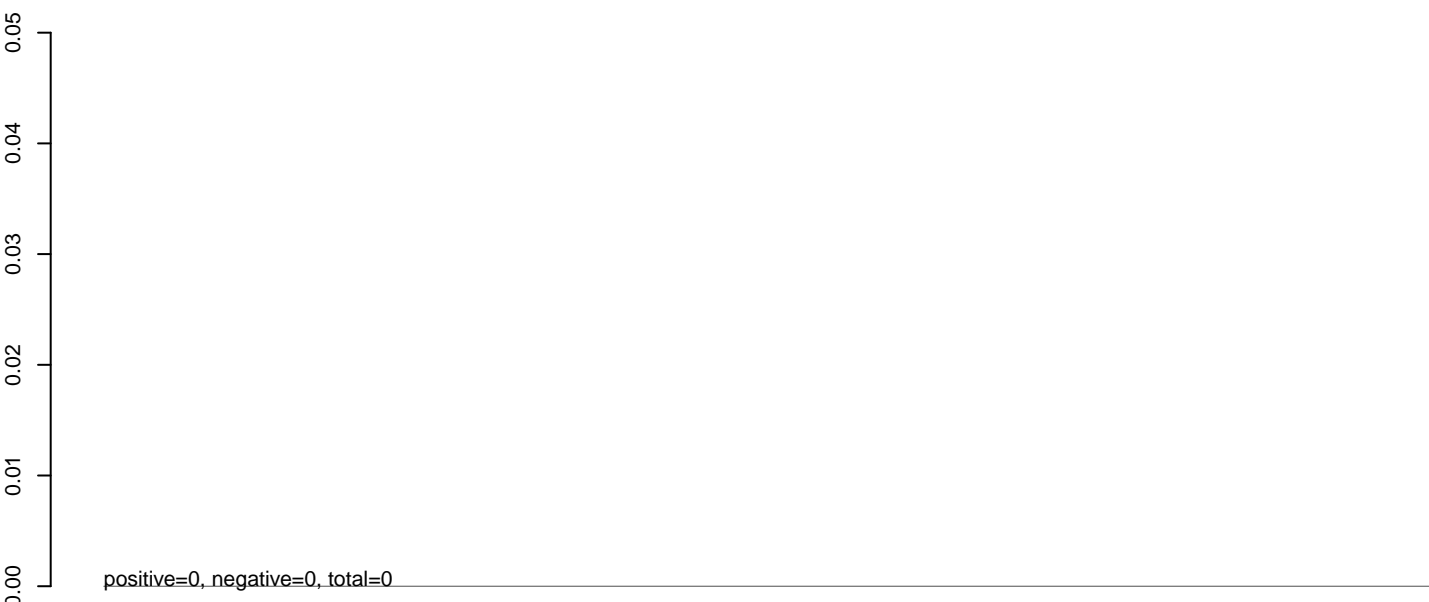
Window size=50, length=14717, virus@NC_017685.1-Obodhiang_virus:1-14717

0 5000 10000 15000

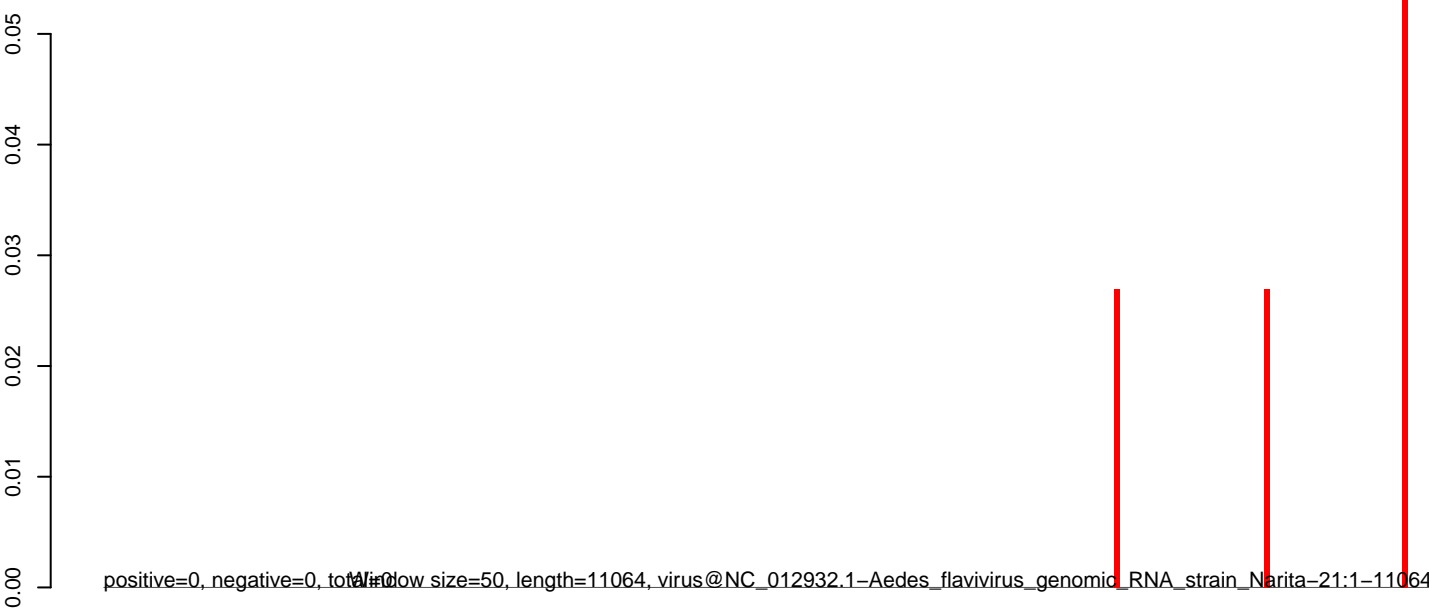
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



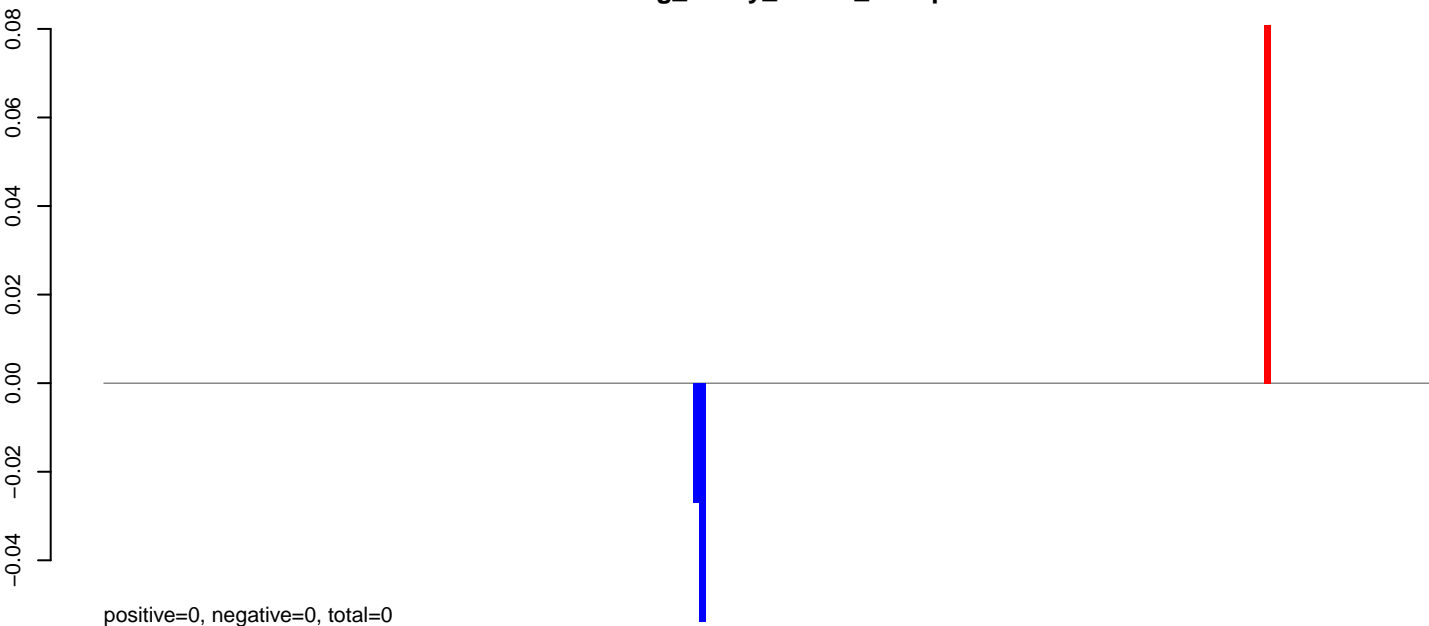
AeAeg_Ovary_GH.rep



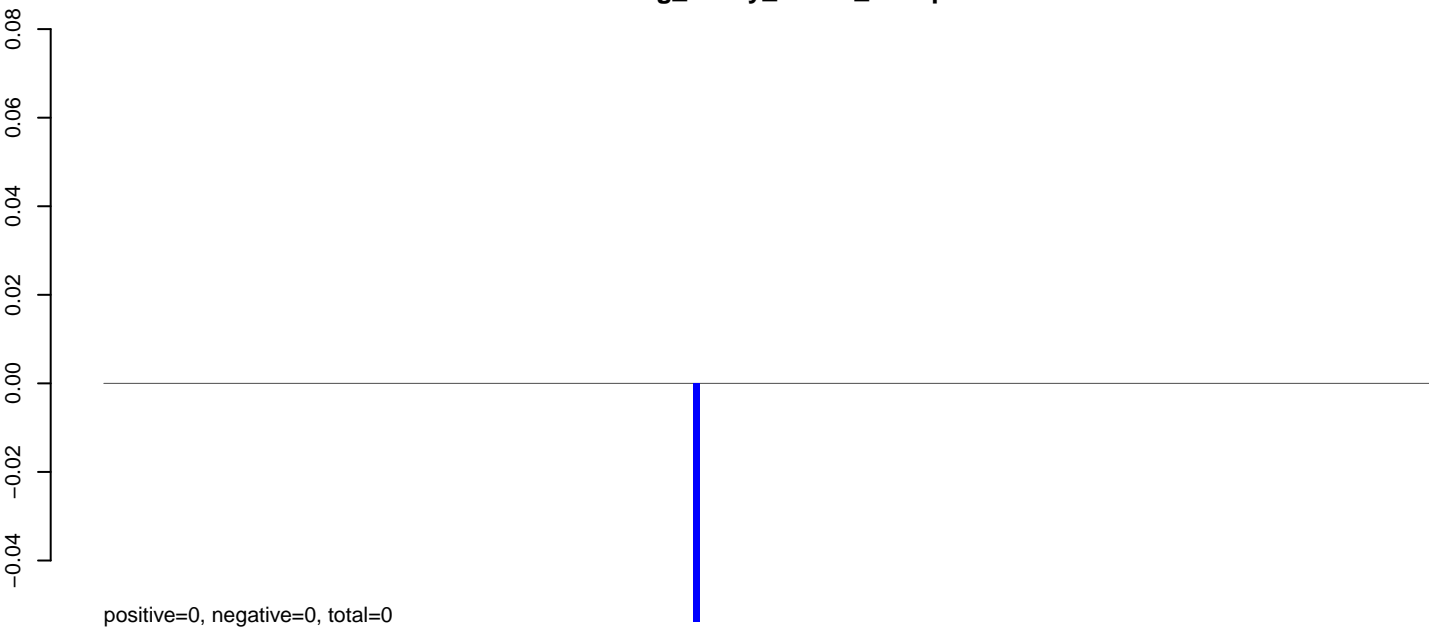
low size=50, length=11064, virus@NC_012932.1-Aedes_flavivirus_genomic_RNA_strain_Narita-21:1-11064

0 2000 4000 6000 8000 10000

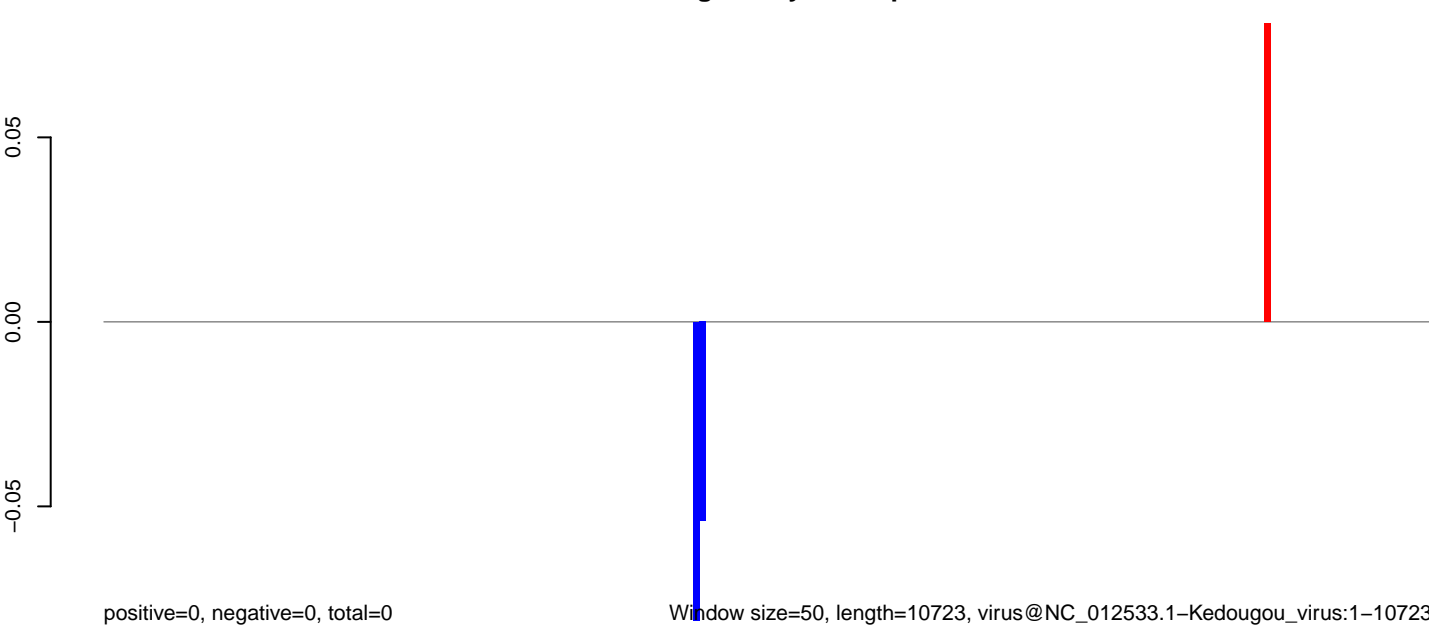
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



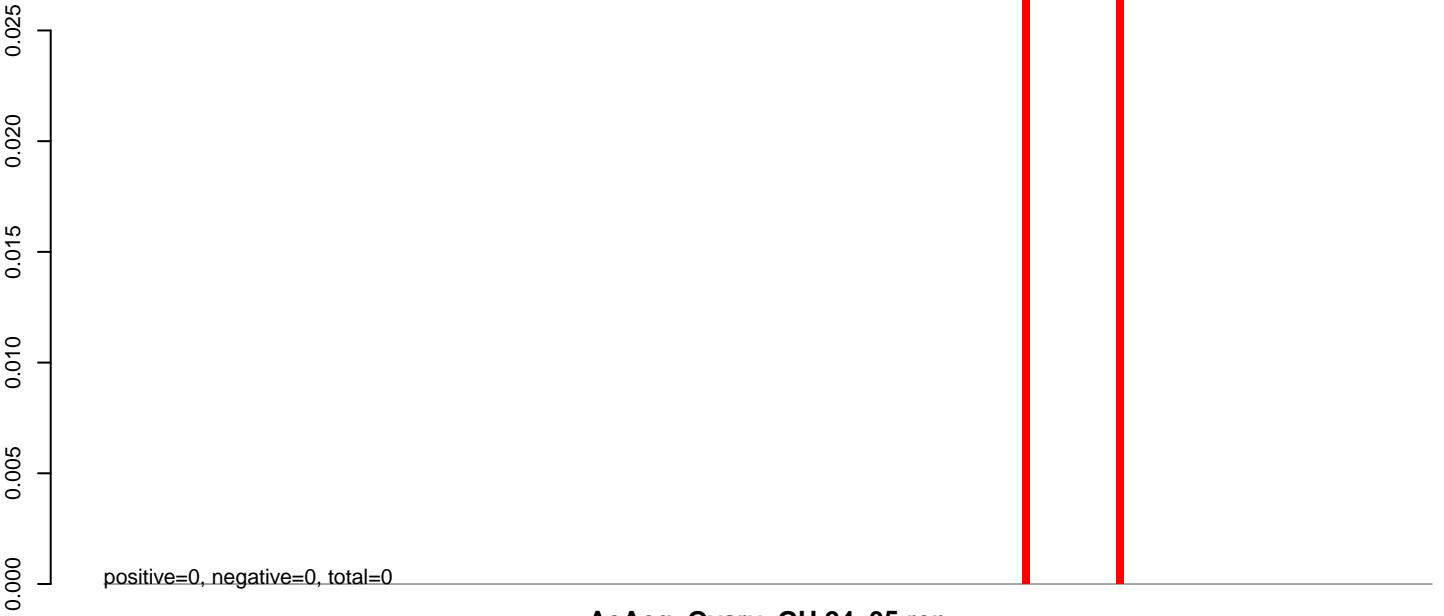
AeAeg_Ovary_GH.rep



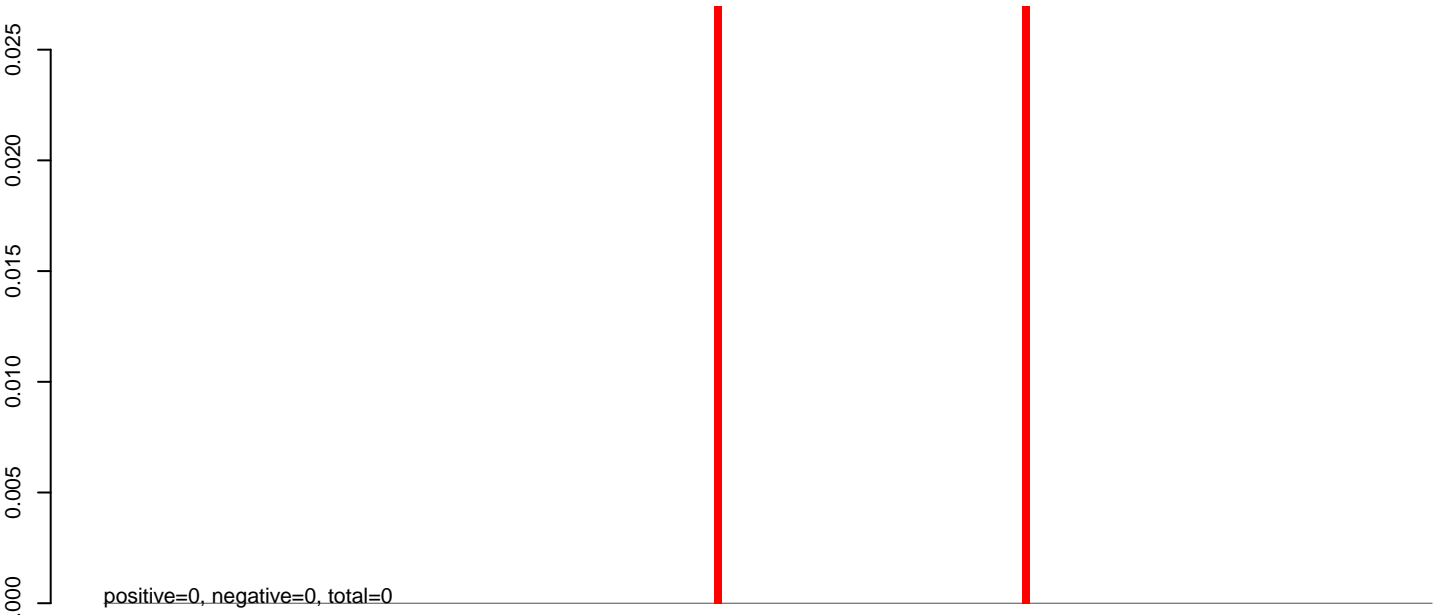
Window size=50, length=10723, virus@NC_012533.1-Kedougou_virus:1-10723

0 2000 4000 6000 8000 10000

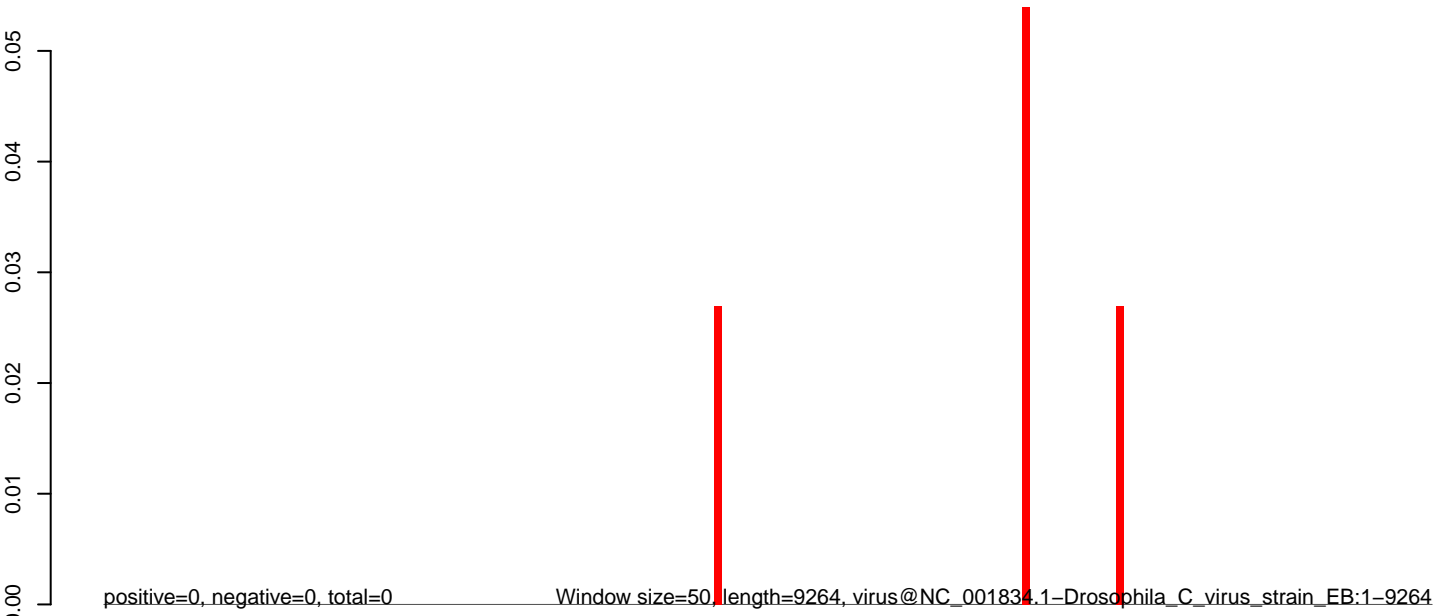
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

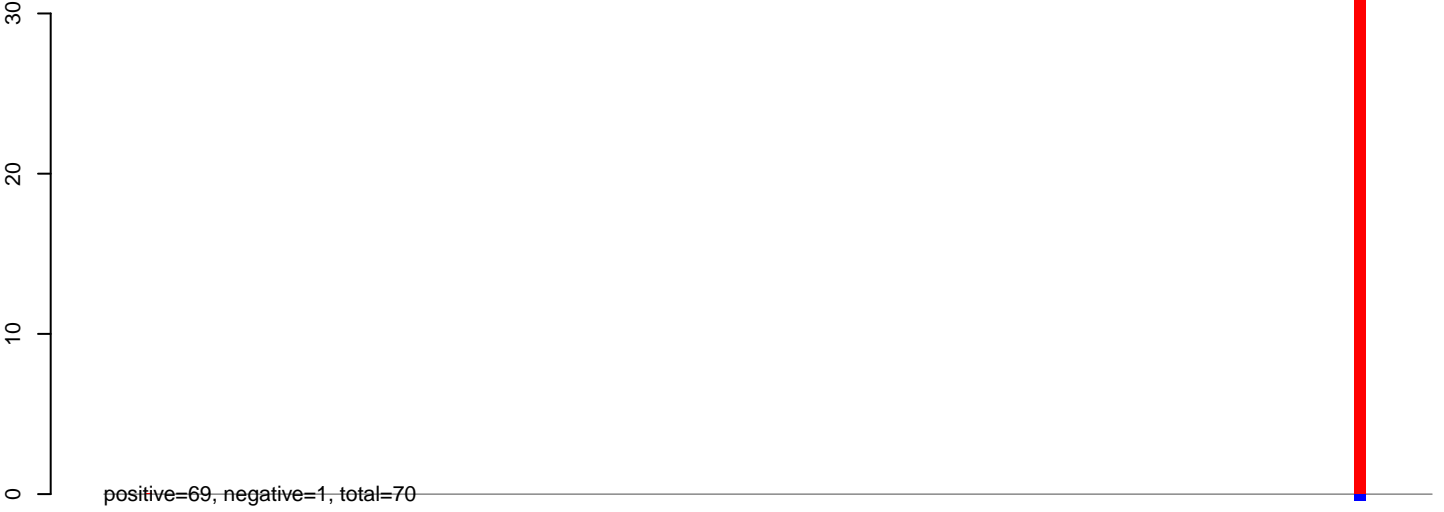


AeAeg_Ovary_GH.rep

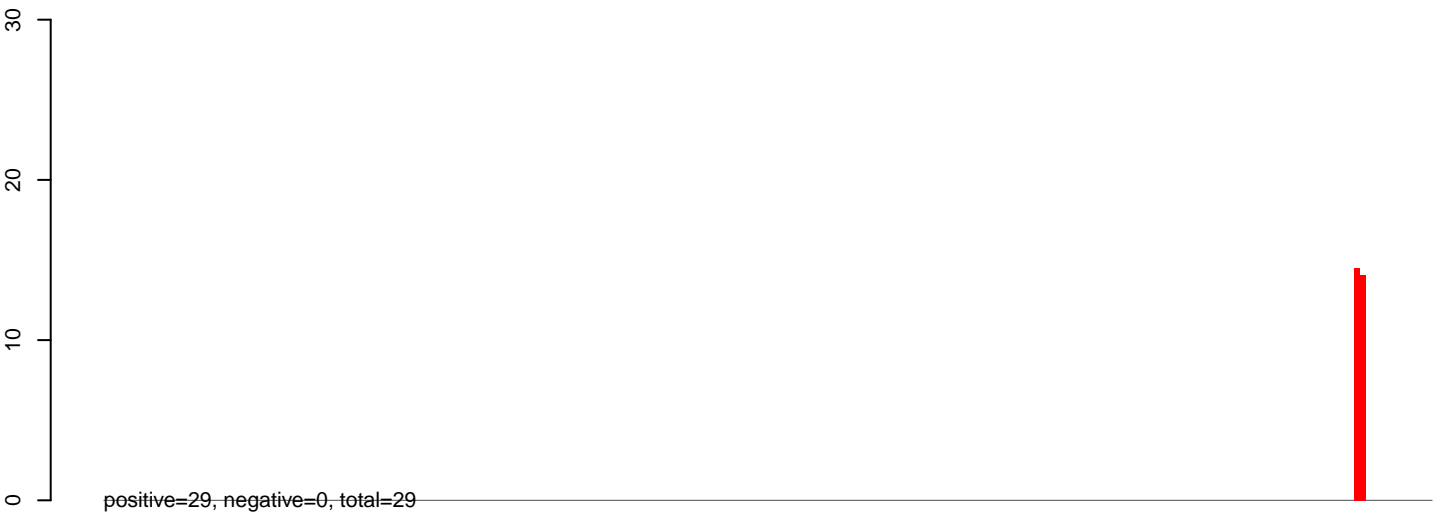


0 2000 4000 6000 8000

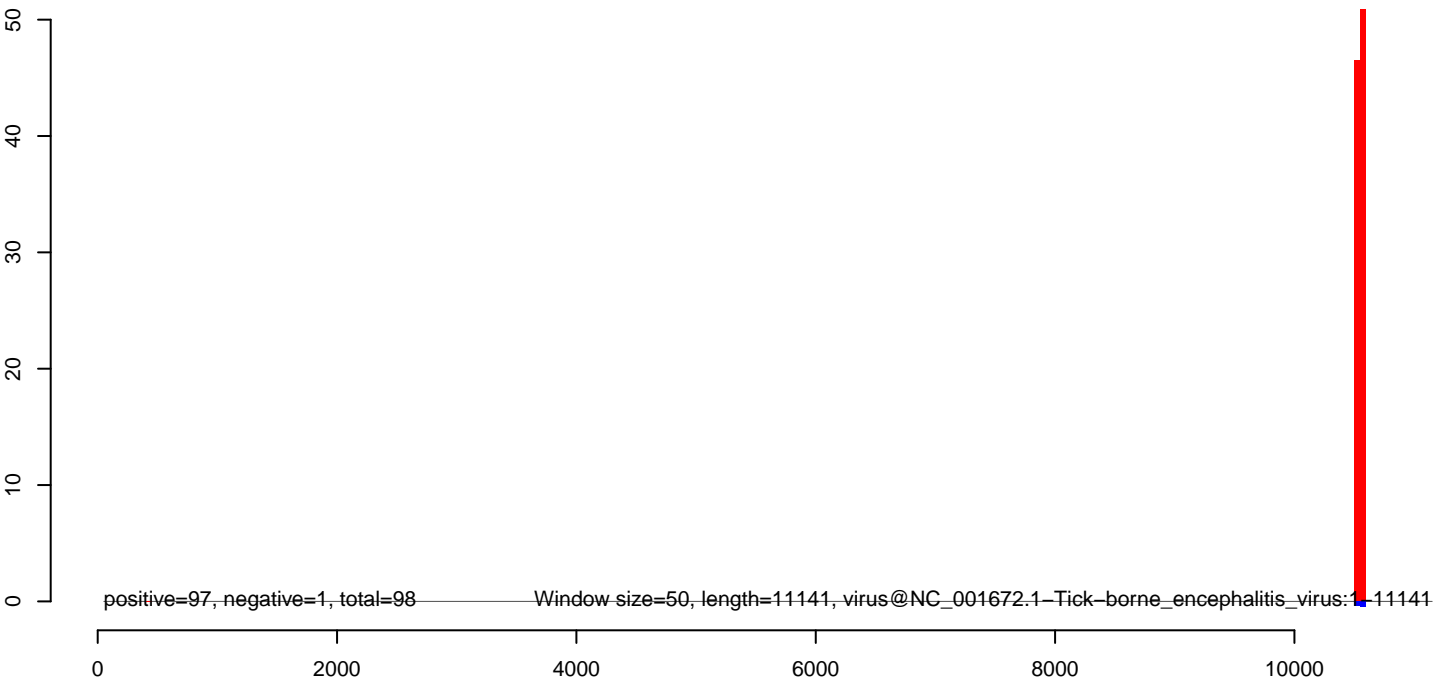
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



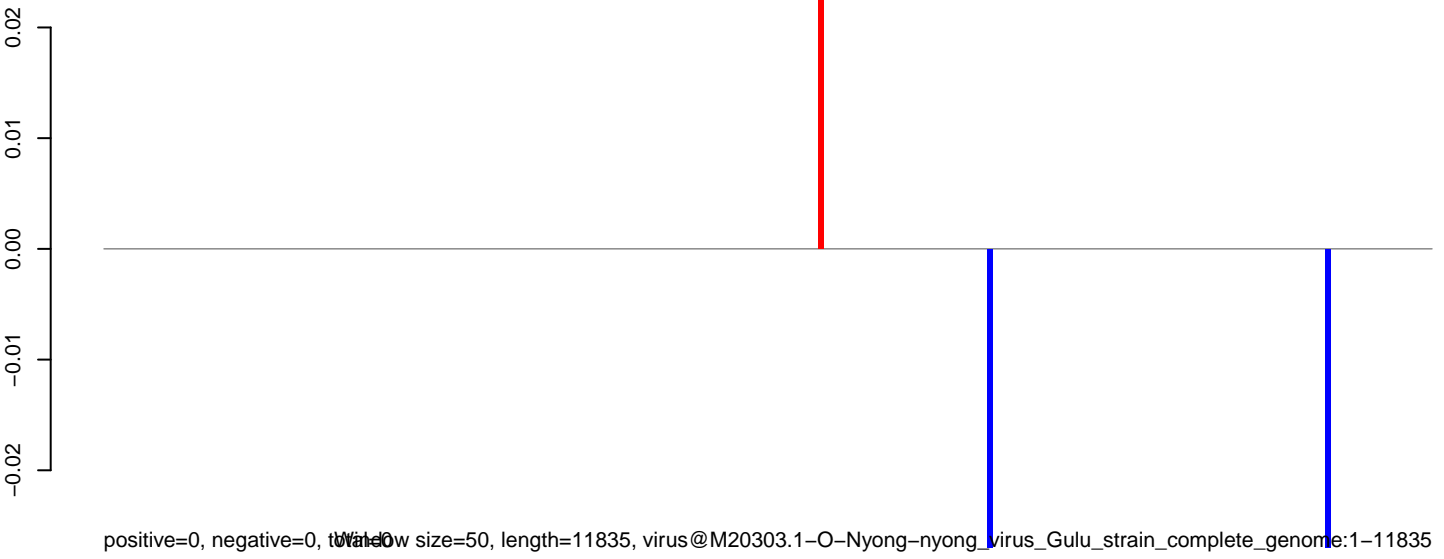
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



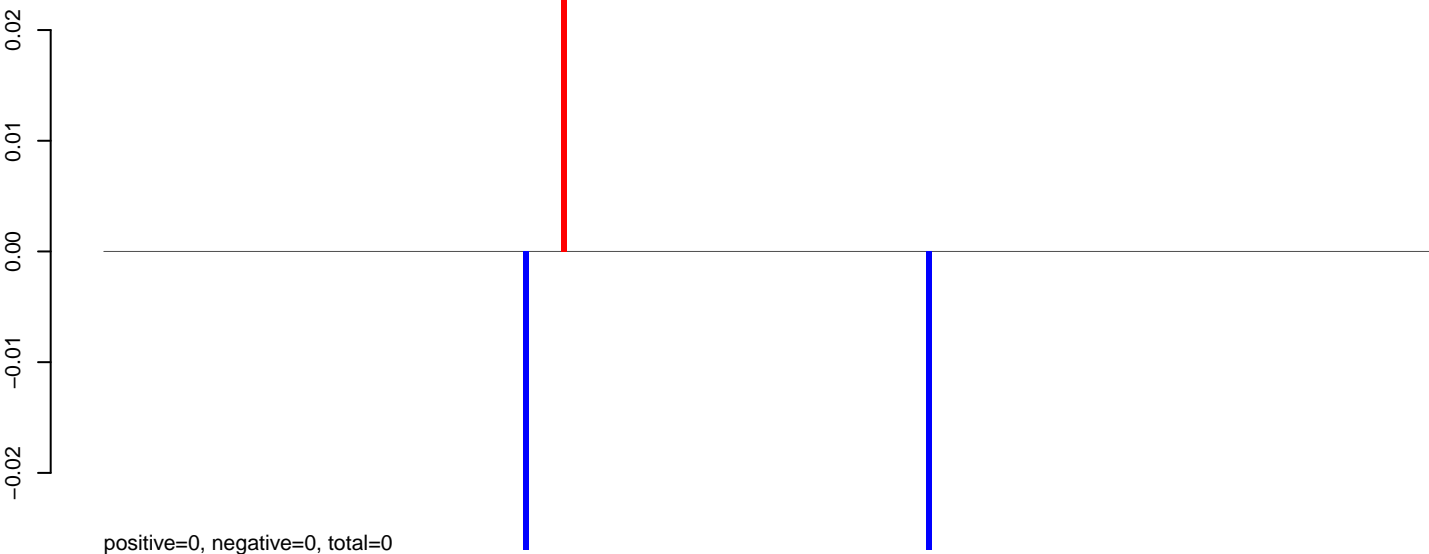
AeAeg_Ovary_GH.rep



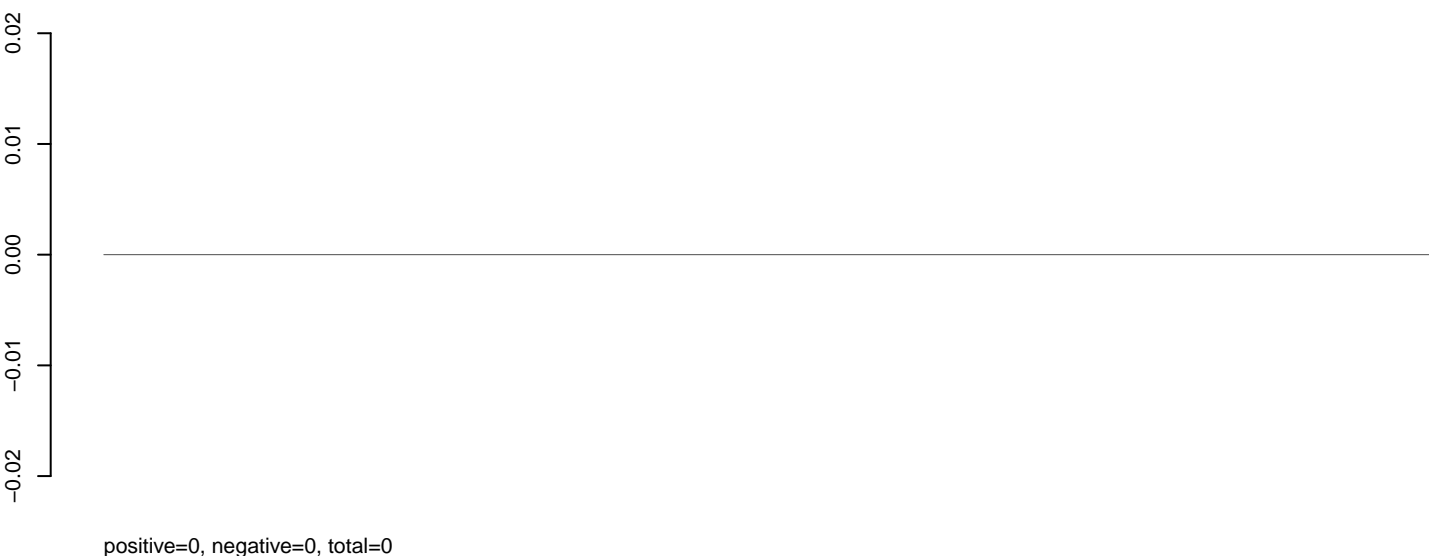
Window size=50, length=11835, virus@M20303.1-O-Nyong-nyong_virus_Gulu_strain_complete_genome:1-11835

0 2000 4000 6000 8000 10000 12000

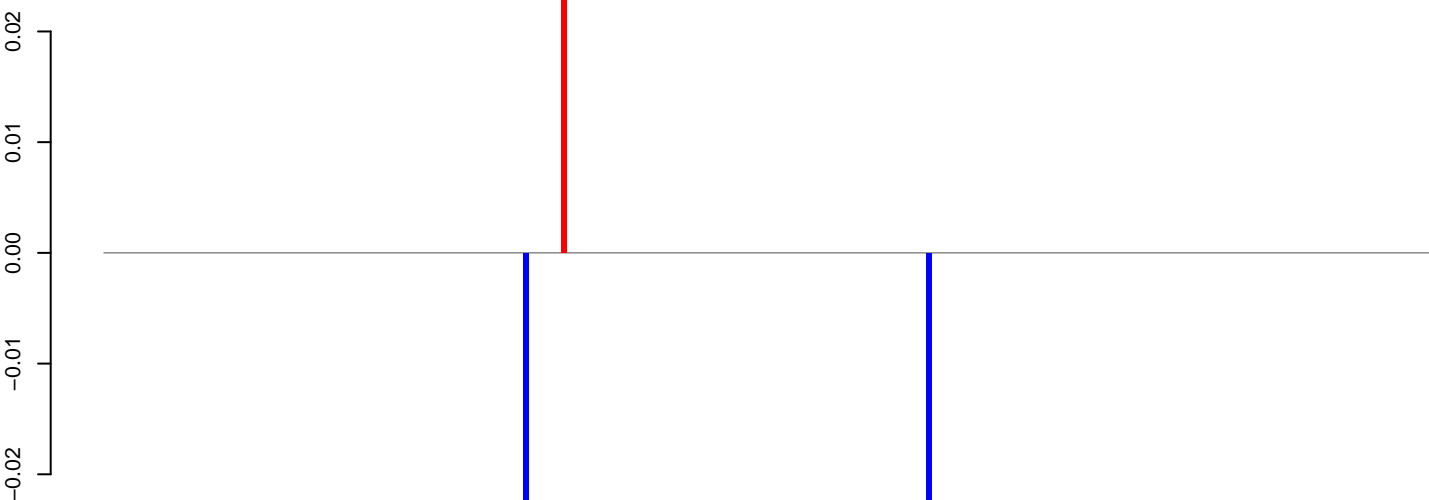
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



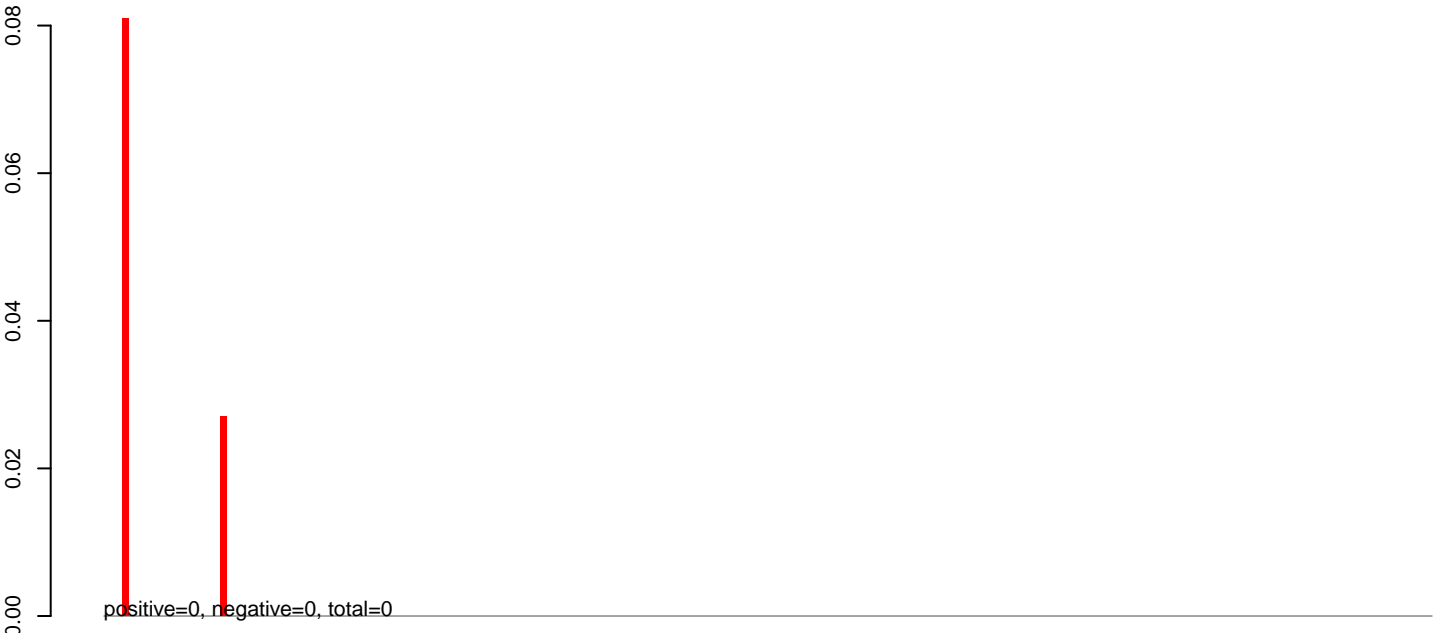
AeAeg_Ovary_GH.rep



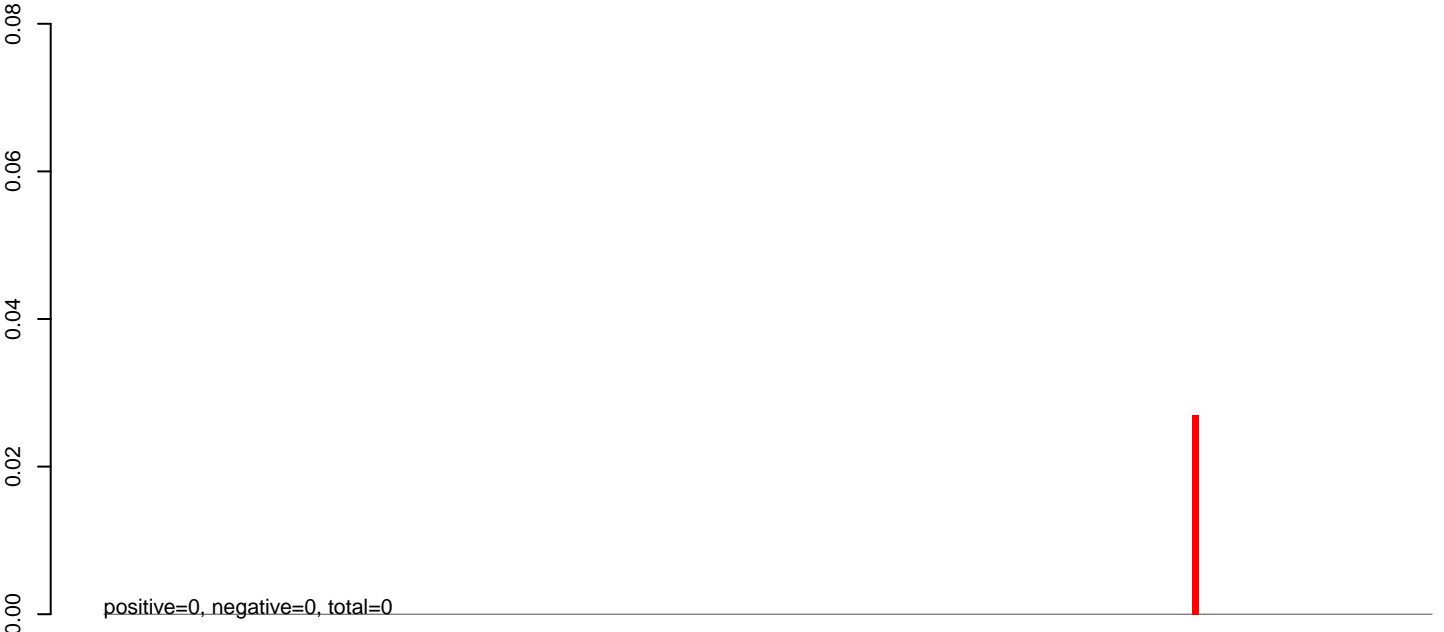
ze=50, size=12237, virus@16156804.1-Tatagline_virus_segment_S_plus_segment_M_plus_segment_L_nucleocapsid_protein_gene:1-12237

0 2000 4000 6000 8000 10000 12000

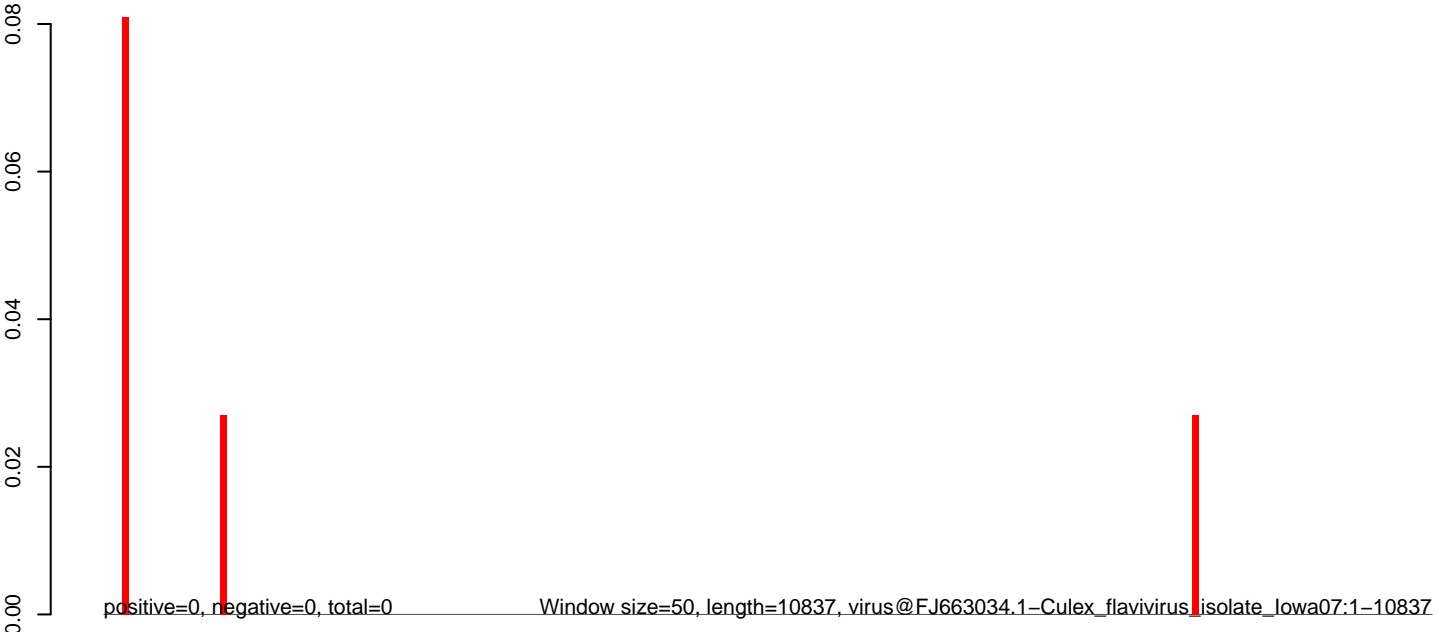
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



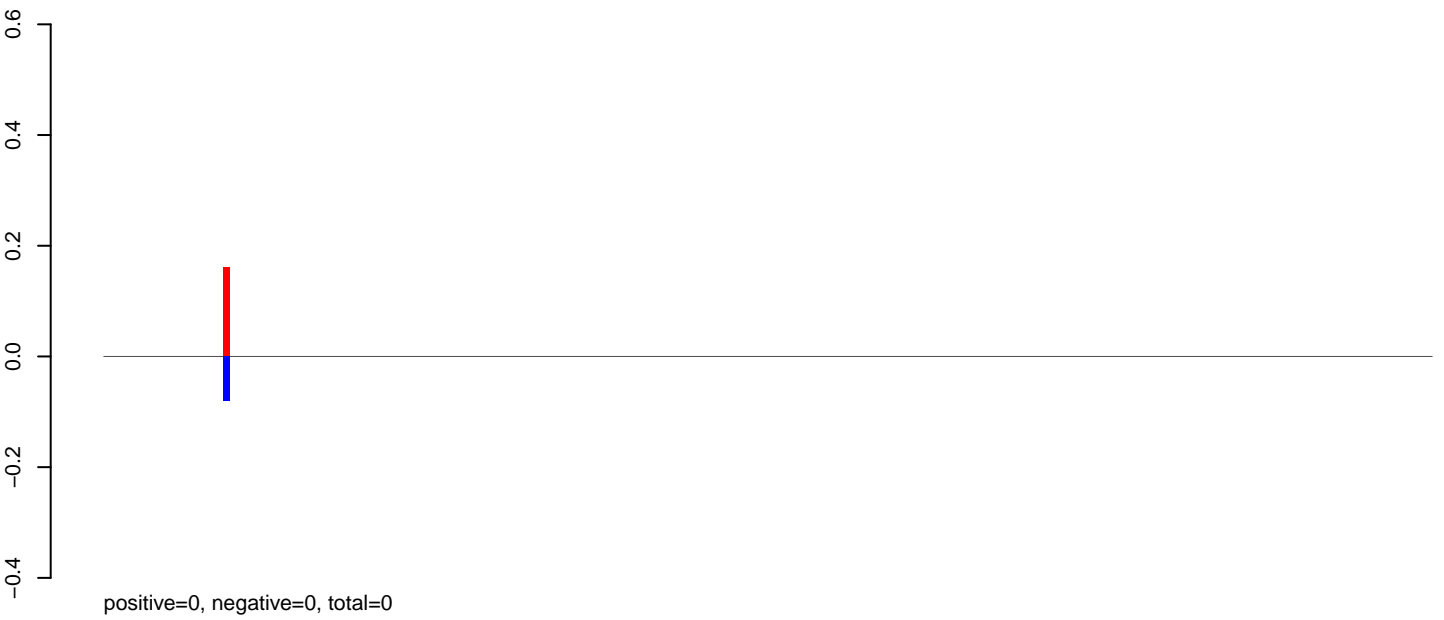
AeAeg_Ovary_GH.rep



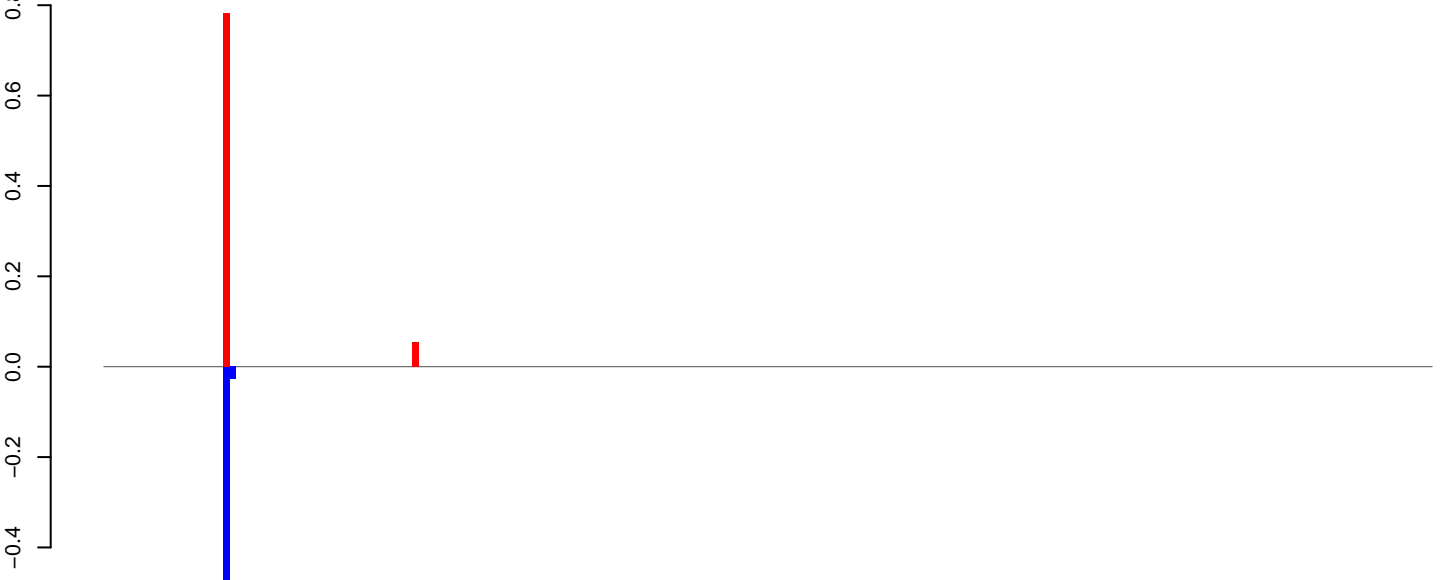
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



AF416352=Aedes albopictus cell_fusing_agent_virus_polyprotein-like_protein_gene_and_agCP8252-like_protein_gene_partial_cds:1-10587

0 2000 4000 6000 8000 10000

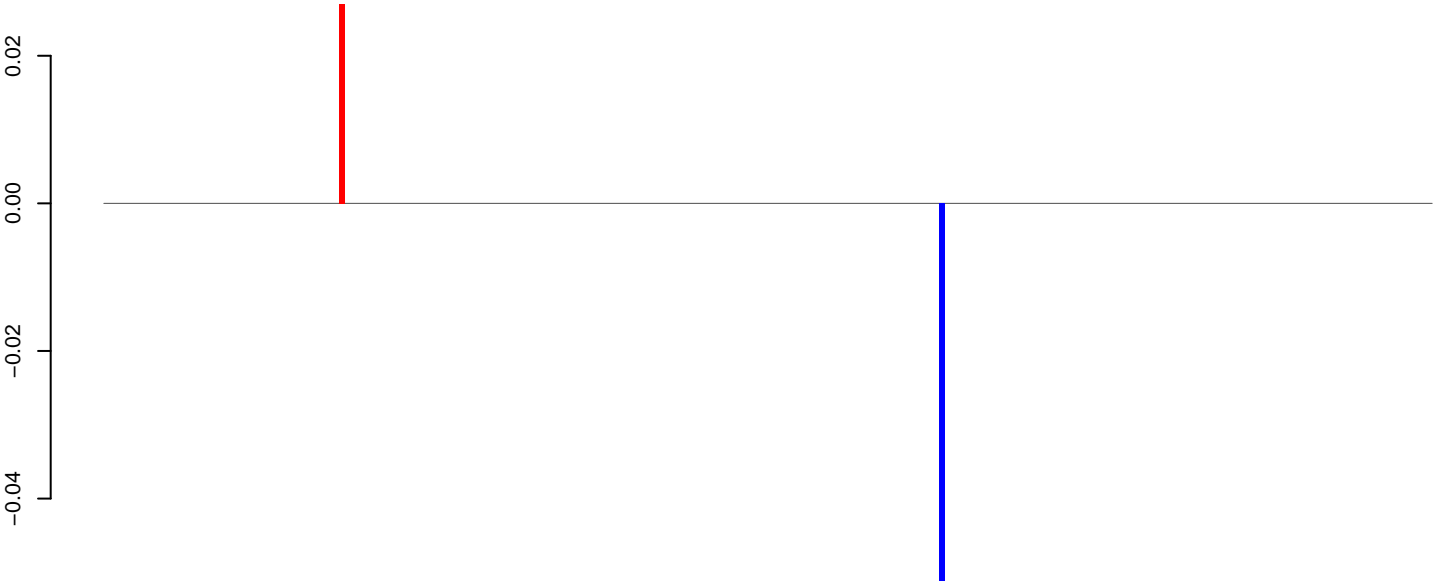
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



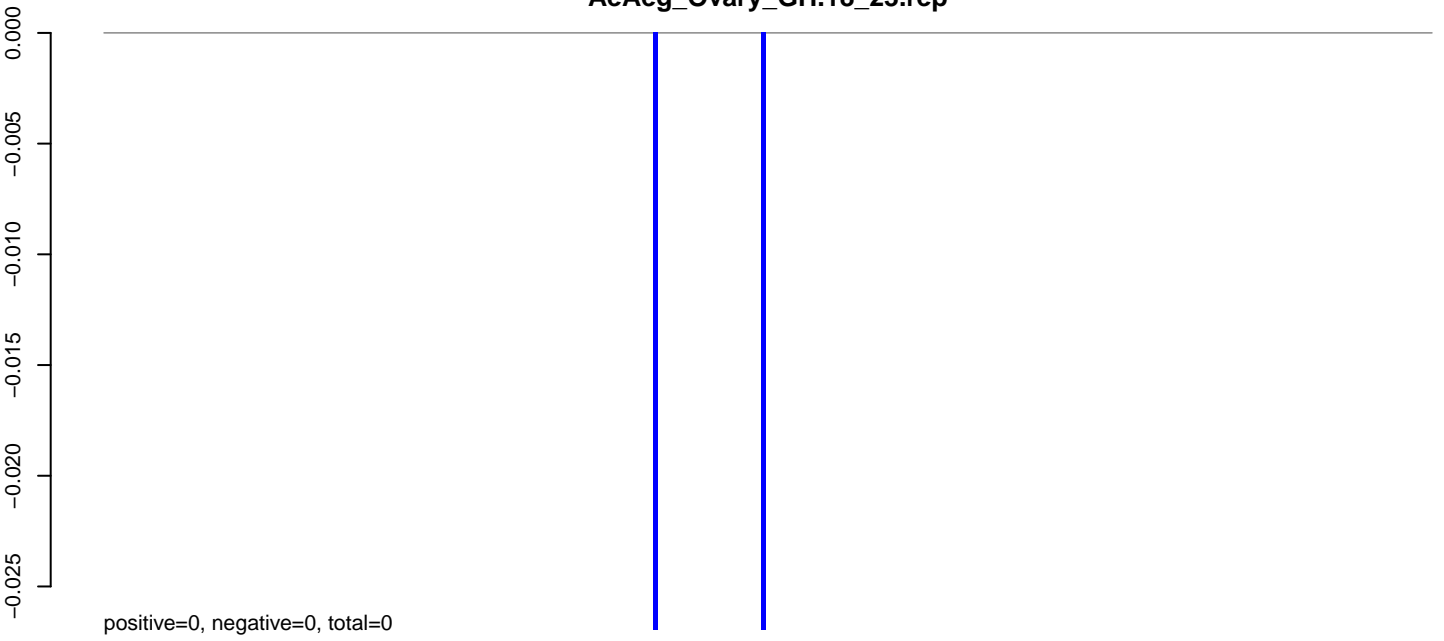
AeAeg_Ovary_GH.rep



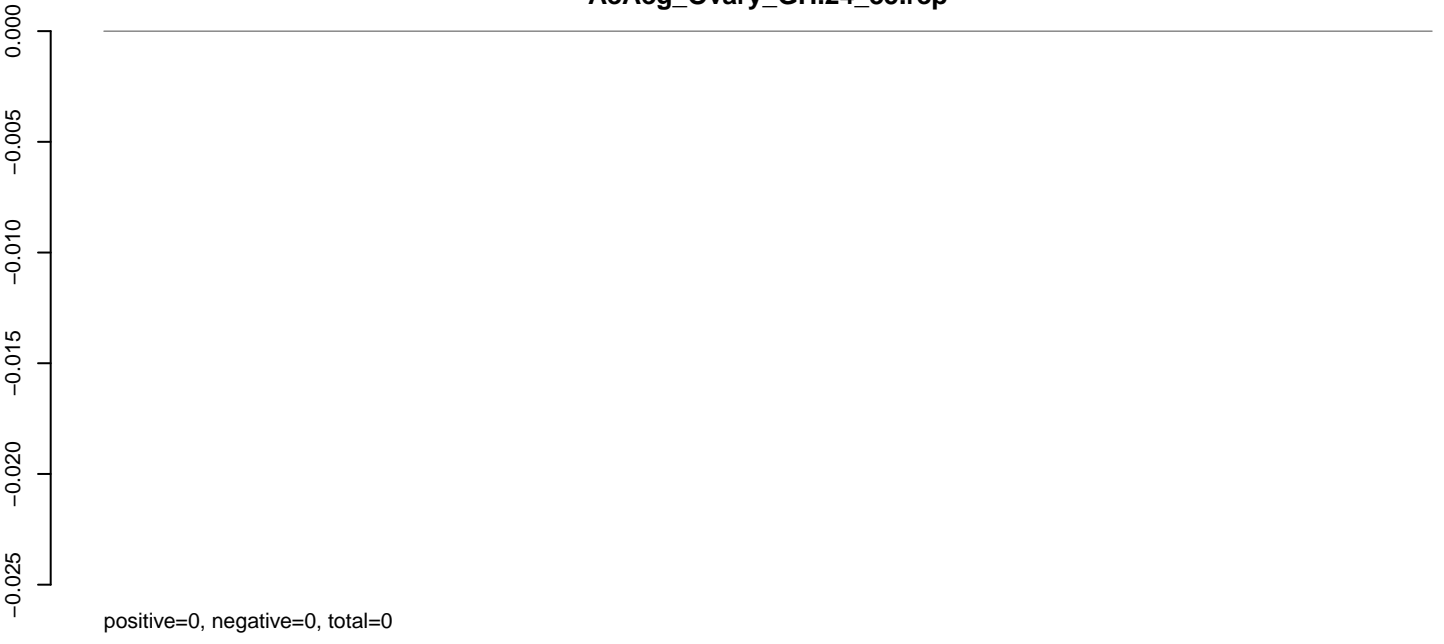
phosphoprotein_hypothetical_proteins_matrix_glycoprotein_hypothetical_protein_and_polymerase_genes:1-12403

0 2000 4000 6000 8000 10000 12000

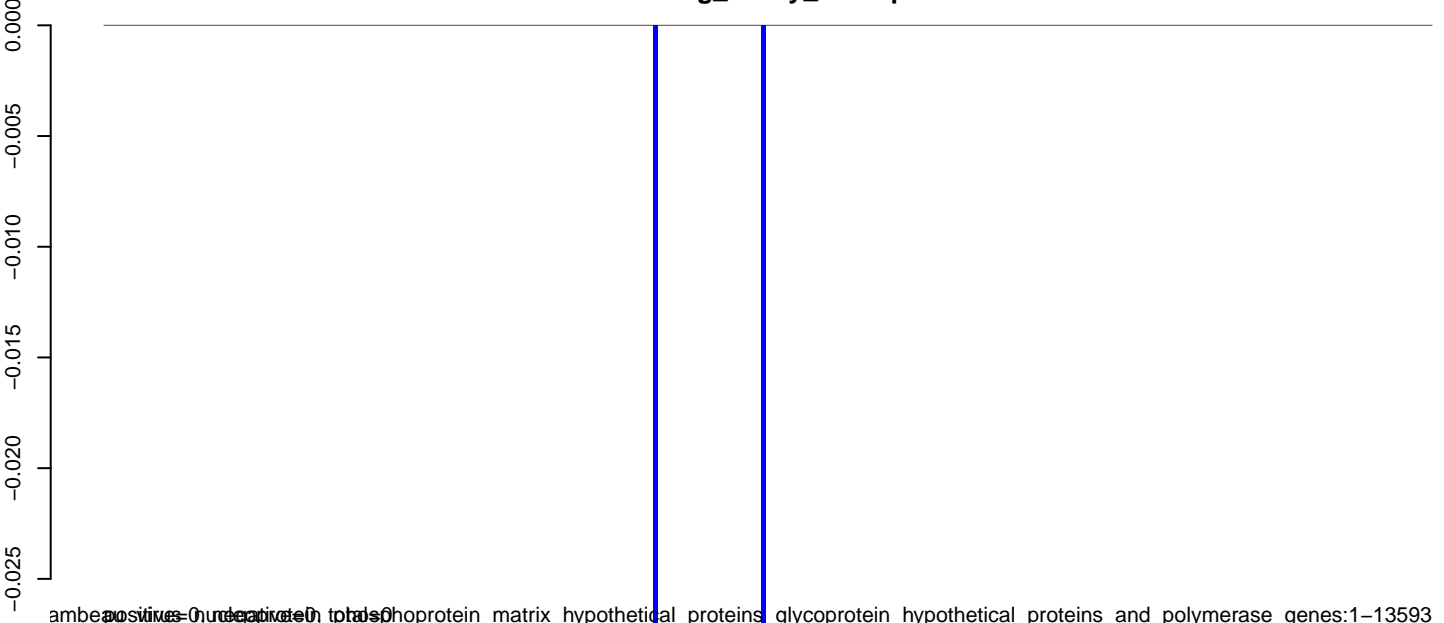
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



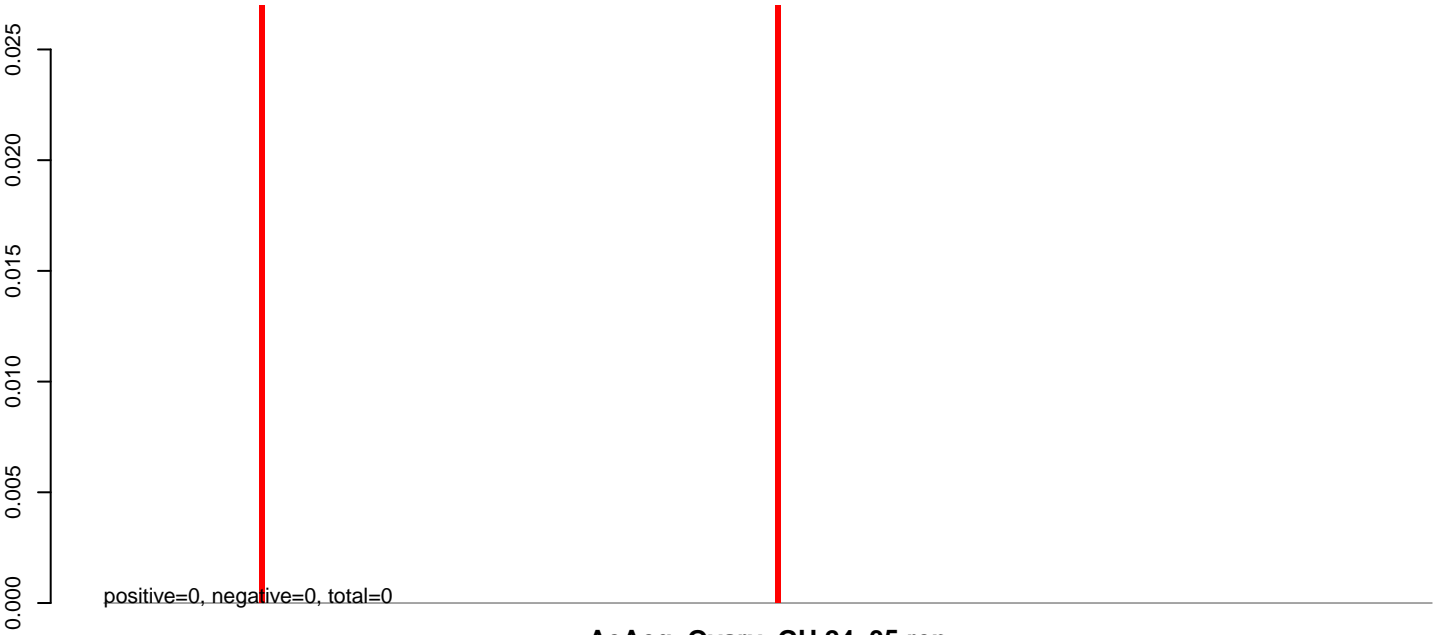
AeAeg_Ovary_GH.rep



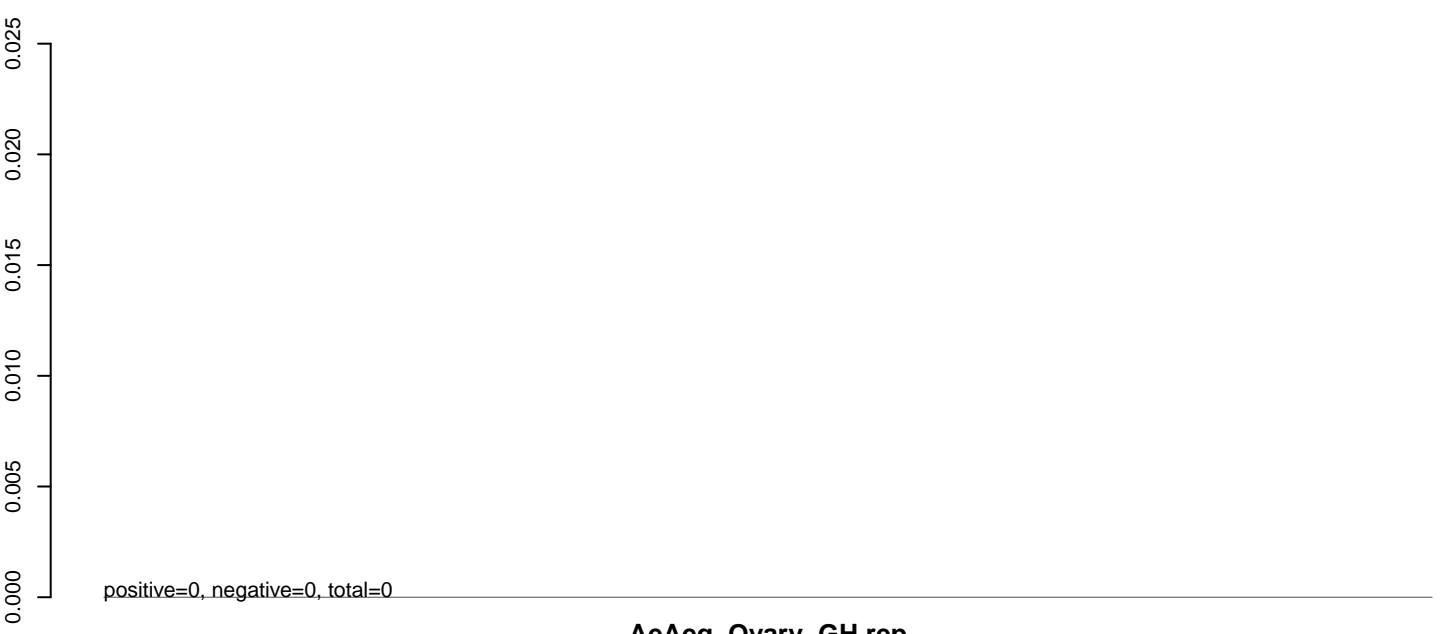
ambee protein_matrix_hypothetical_proteins_glycoprotein_hypothetical_proteins_and_polymerase_genes:1-13593

0 2000 4000 6000 8000 10000 12000 14000

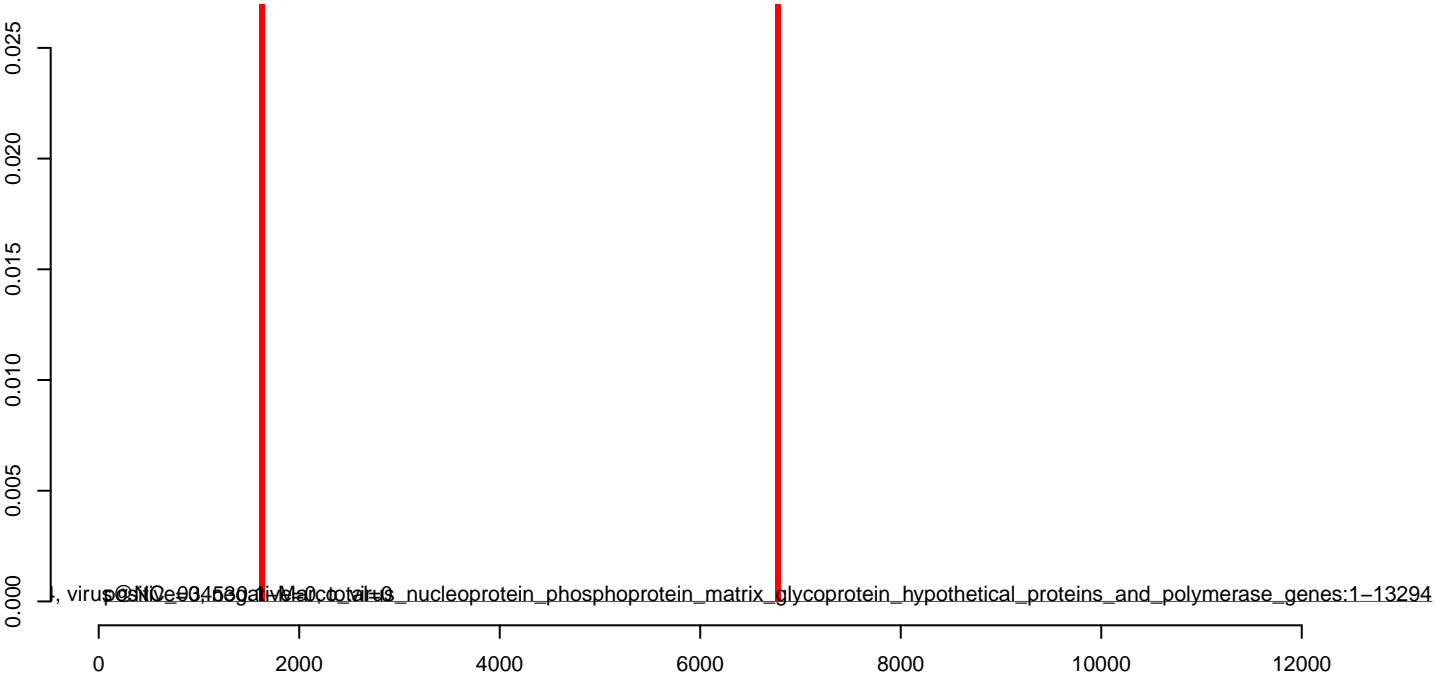
AeAeg_Ovary_GH.18_23.rep



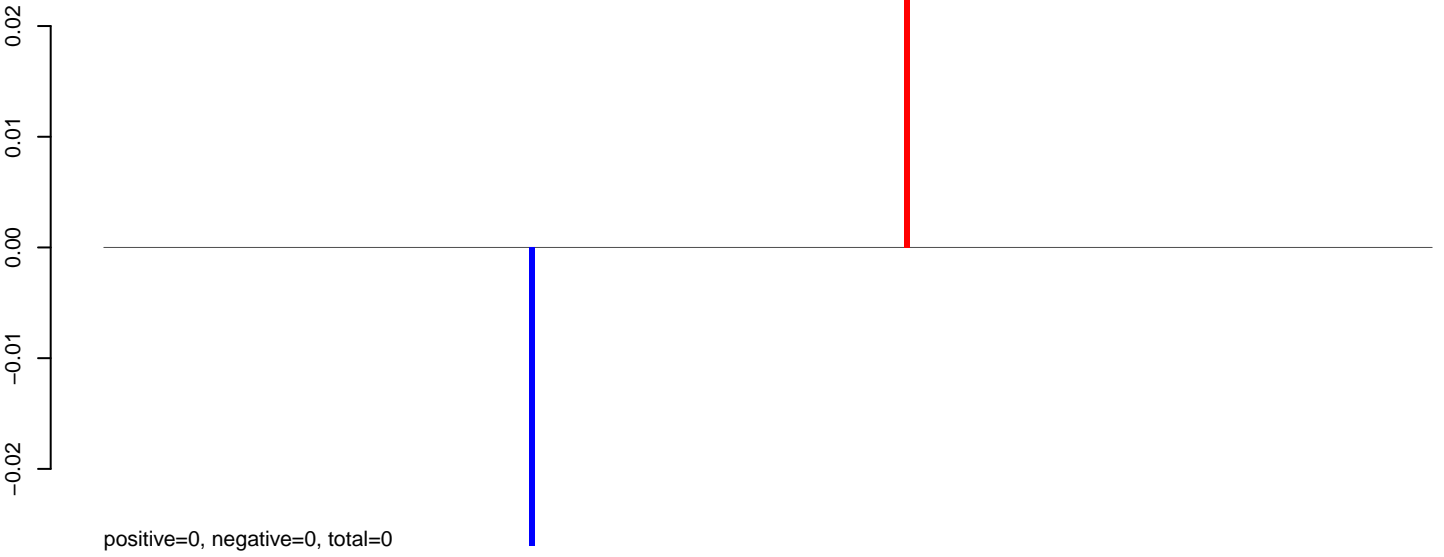
AeAeg_Ovary_GH.24_35.rep



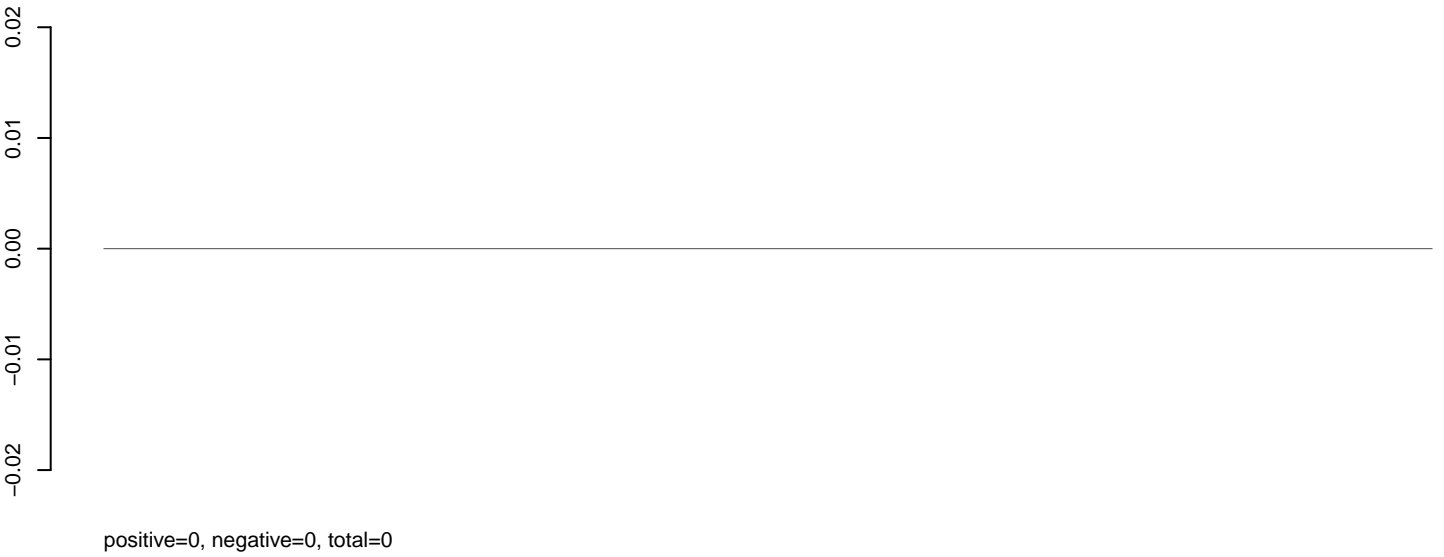
AeAeg_Ovary_GH.rep



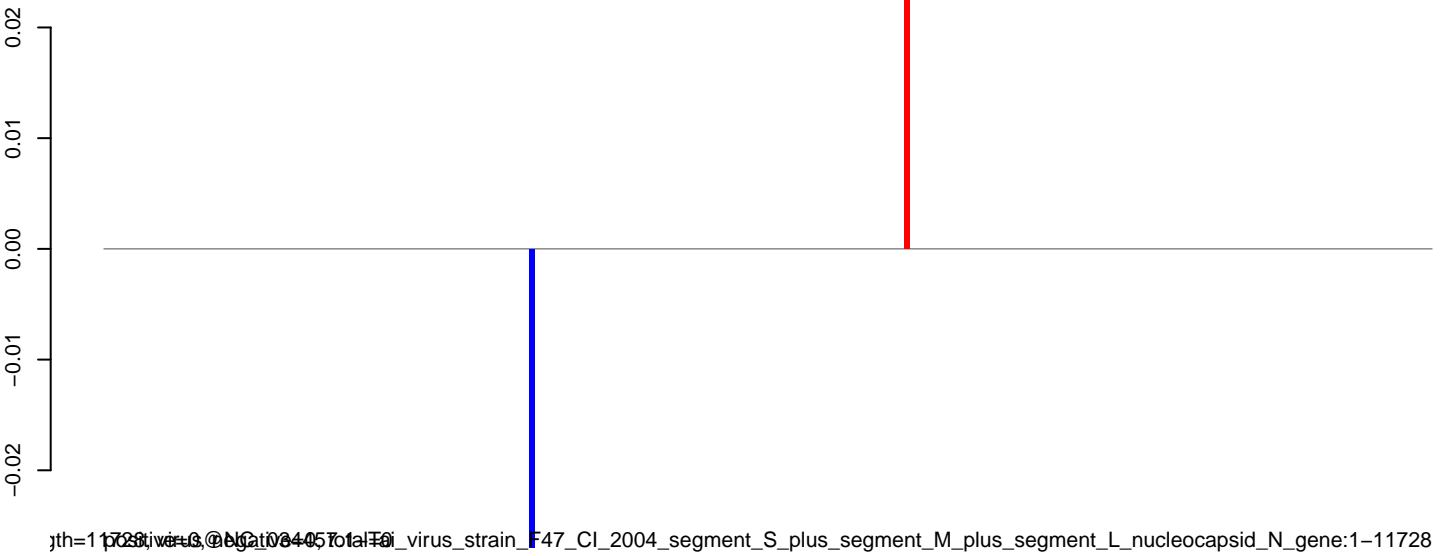
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



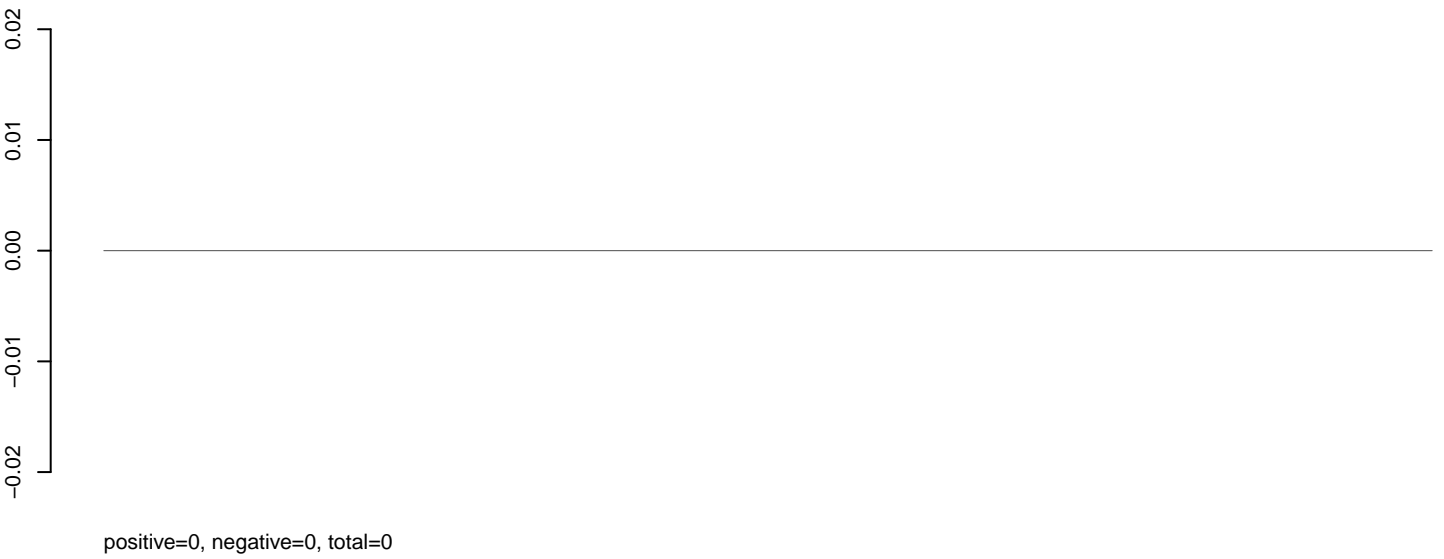
AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



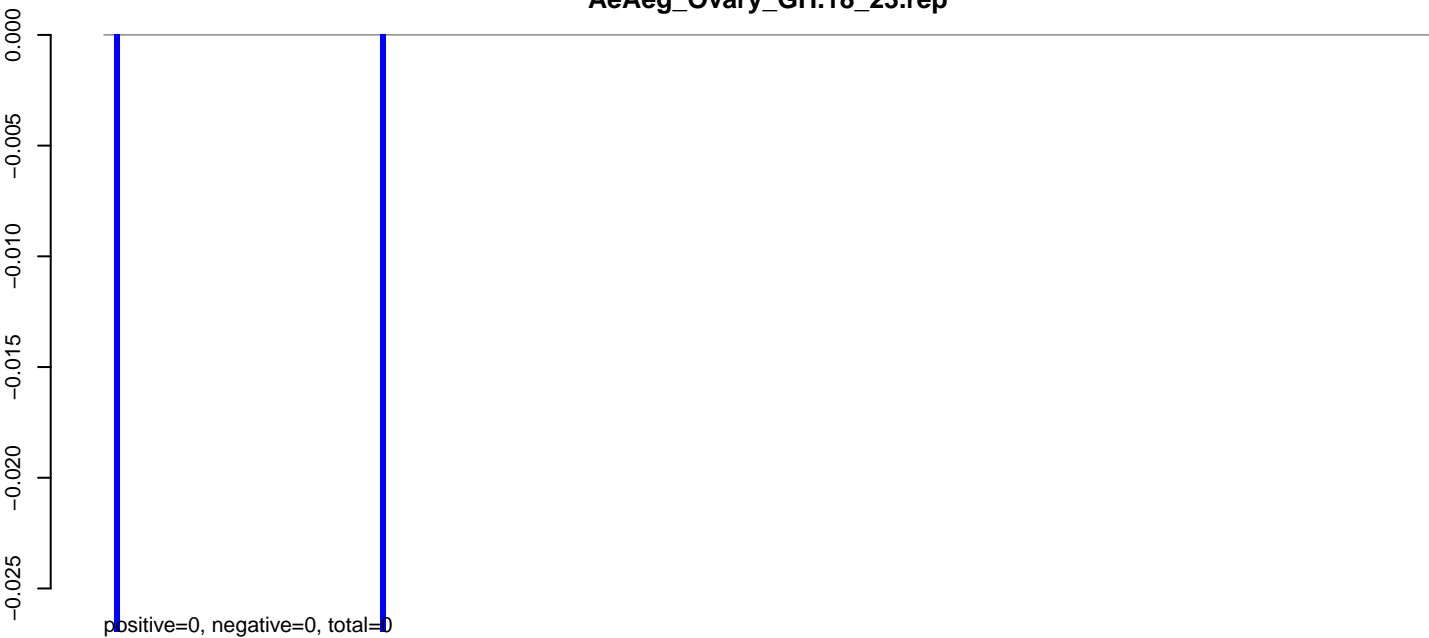
AeAeg_Ovary_GH.rep



-Kame... phosphoprotein_hypothetical_proteins_matrix_glycoprotein_hypothetical_protein_and_polymerase_genes:1-13209

0 2000 4000 6000 8000 10000 12000

AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



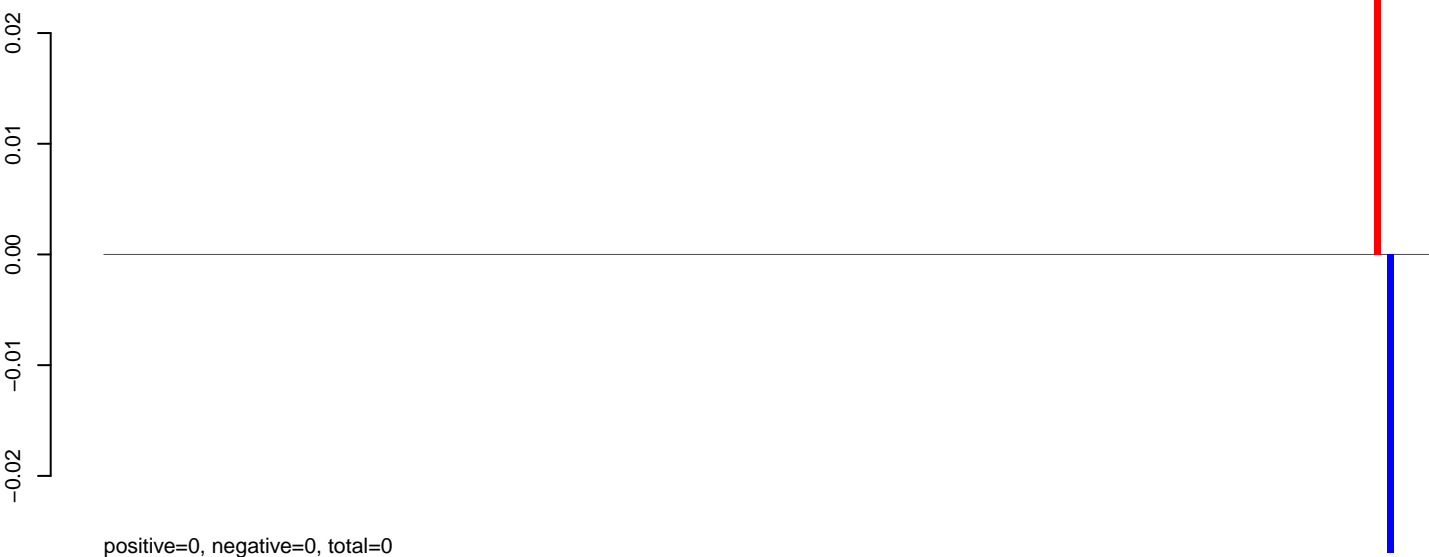
AeAeg_Ovary_GH.rep



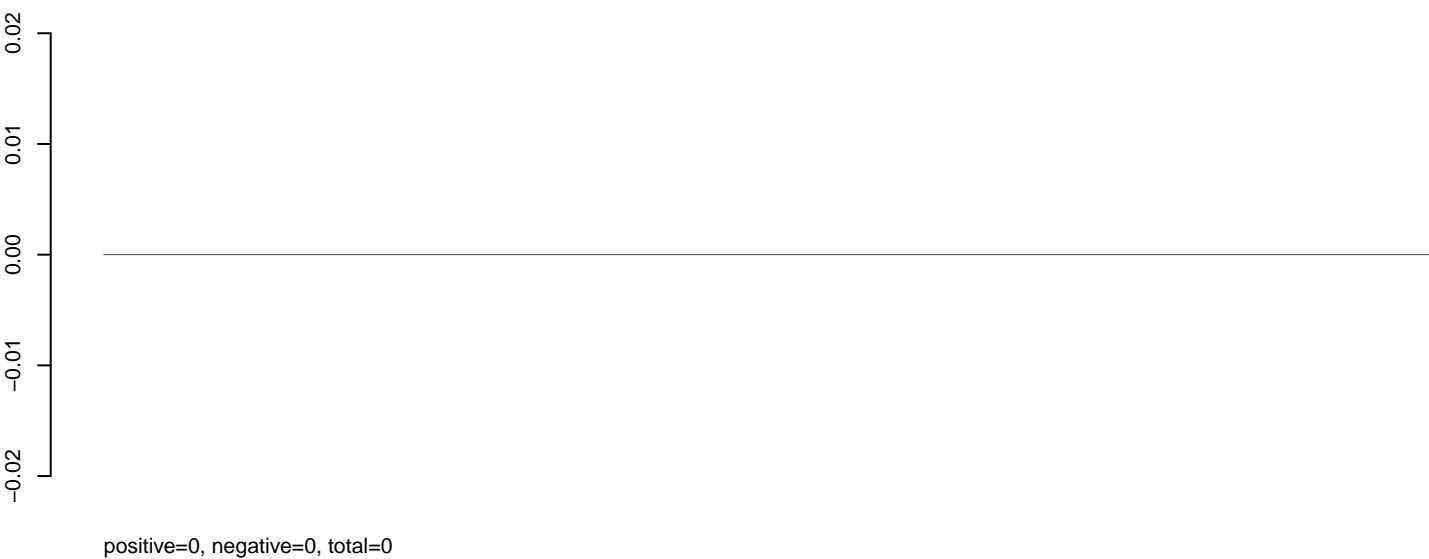
losque positive=0, negative=0, total=0 phosphoprotein_hypothetical_proteins_matrix_glycoprotein_hypothetical_protein_and_polymerase_genes:1-12957

0 2000 4000 6000 8000 10000 12000

AeAeg_Ovary_GH.18_23.rep



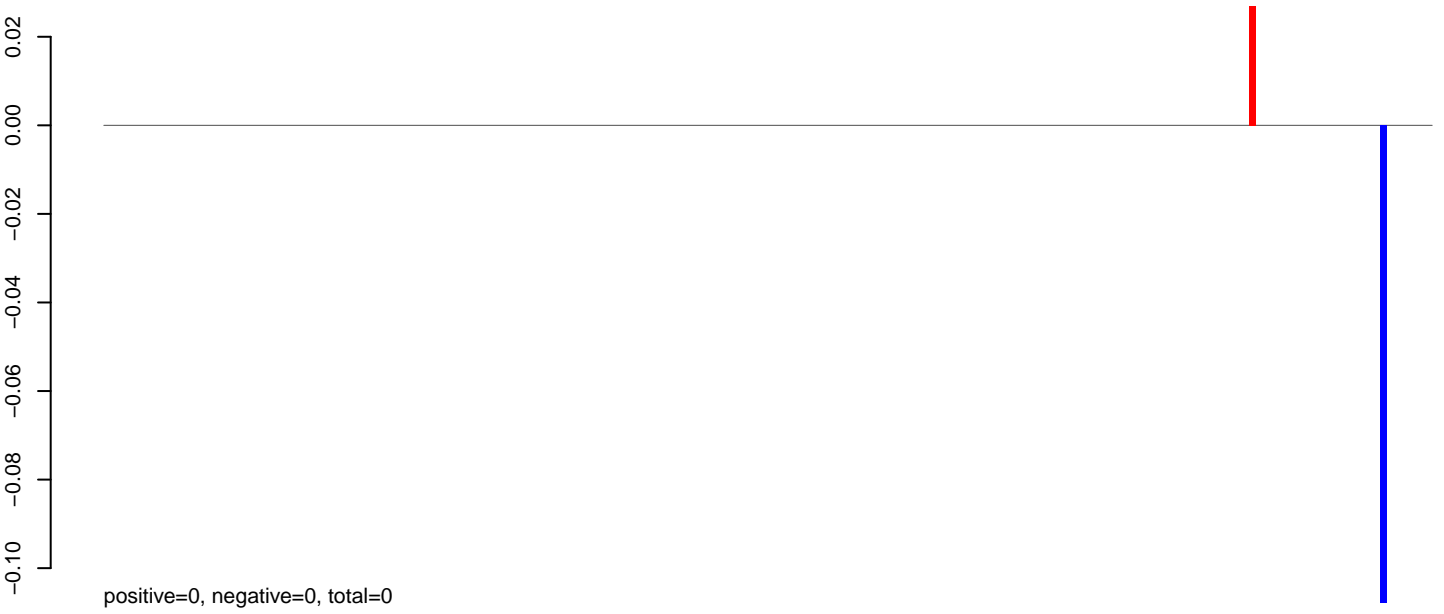
AeAeg_Ovary_GH.24_35.rep



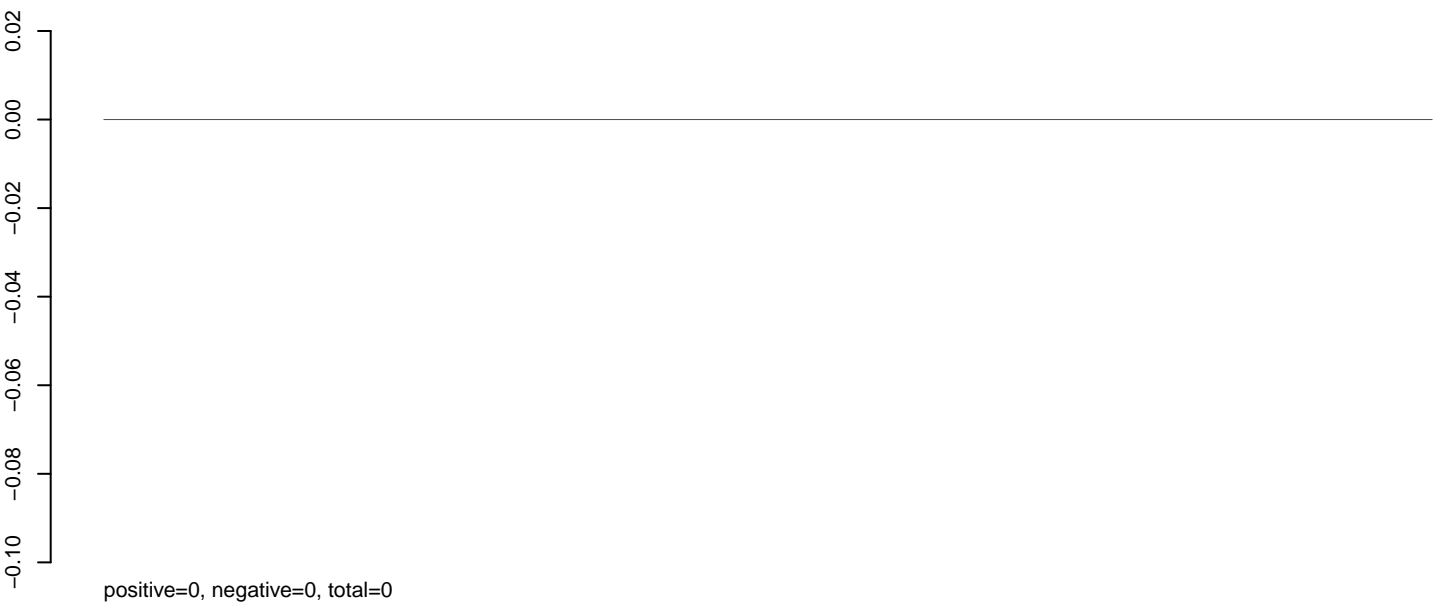
AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



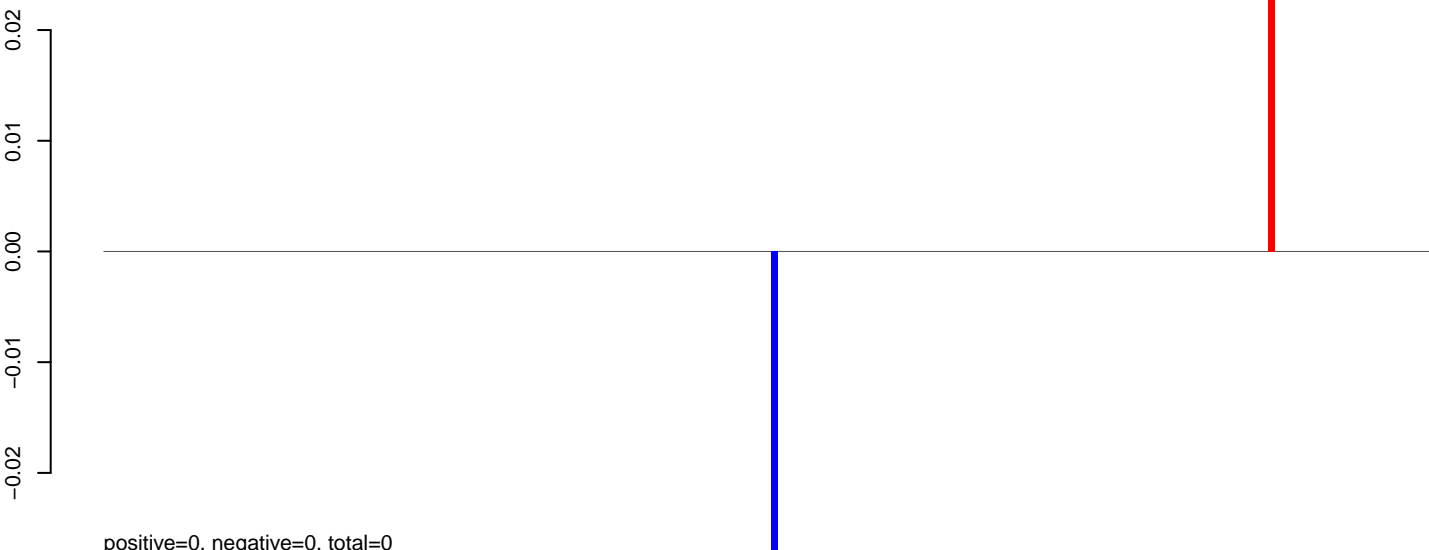
AeAeg_Ovary_GH.rep



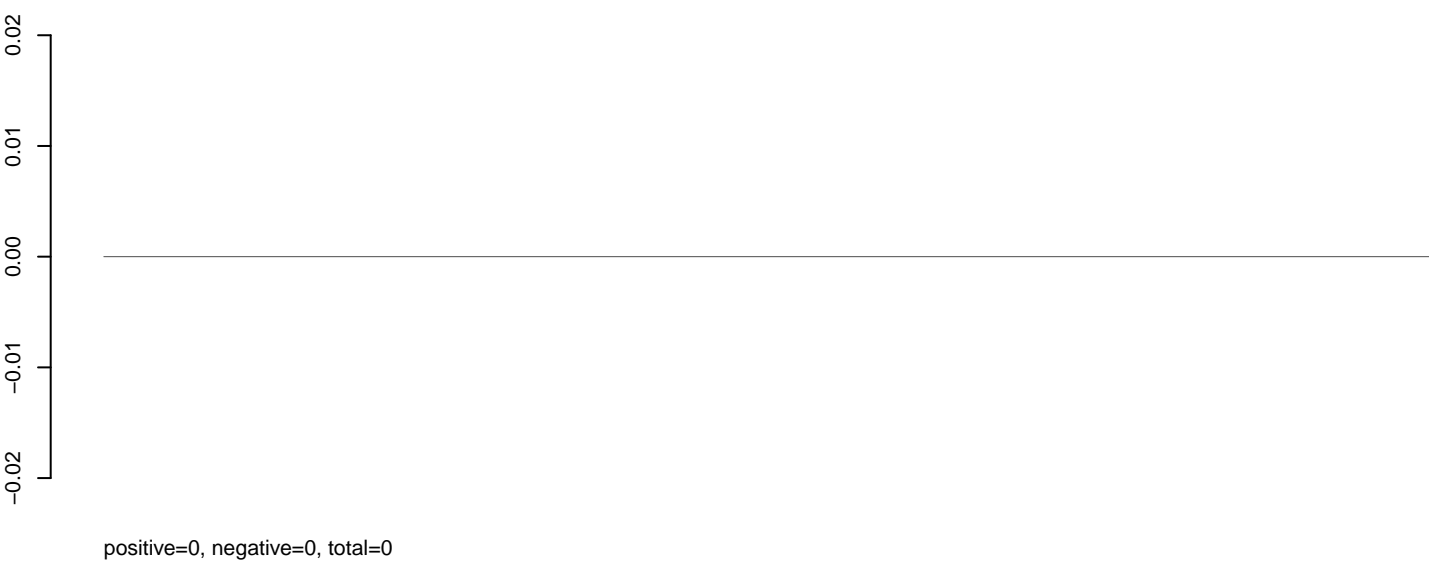
Window size=50, length=10173, virus@NC_033699.1-Jugra_virus_strain_P-9-314_polyprotein_gene:1-10173

0 2000 4000 6000 8000 10000

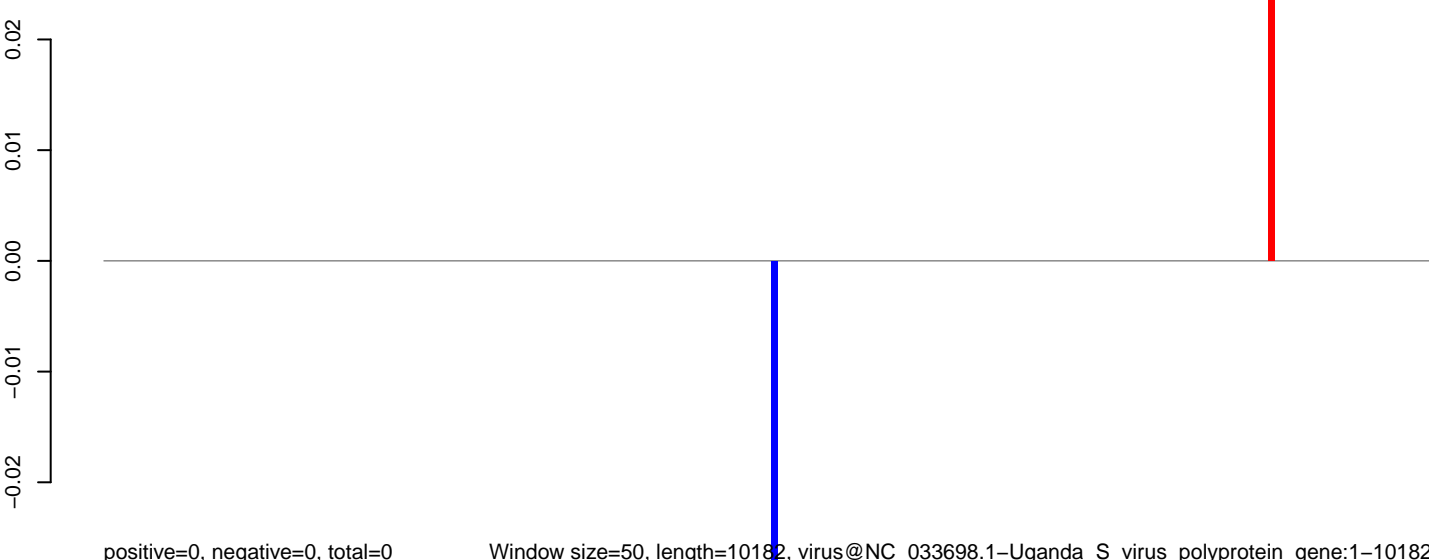
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



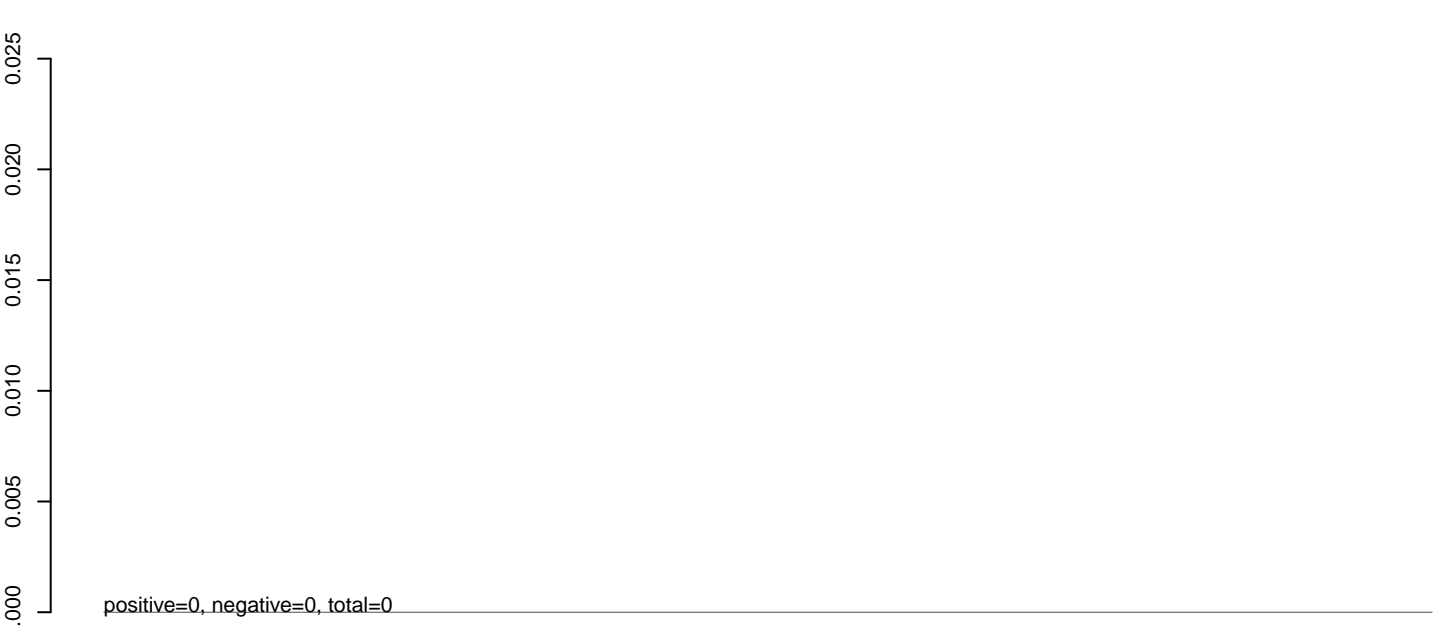
Window size=50, length=10182, virus@NC_033698.1-Uganda_S_virus_polyprotein_gene:1-10182

0 2000 4000 6000 8000 10000

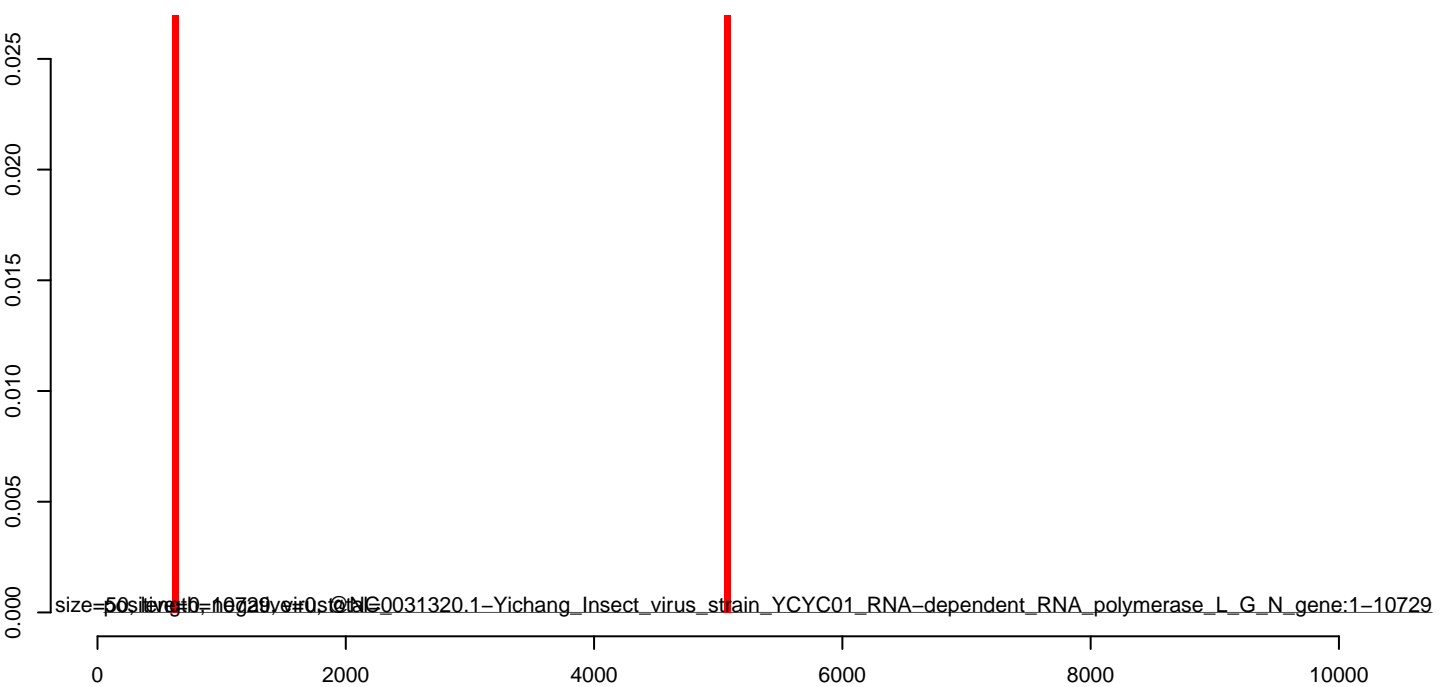
AeAeg_Ovary_GH.18_23.rep



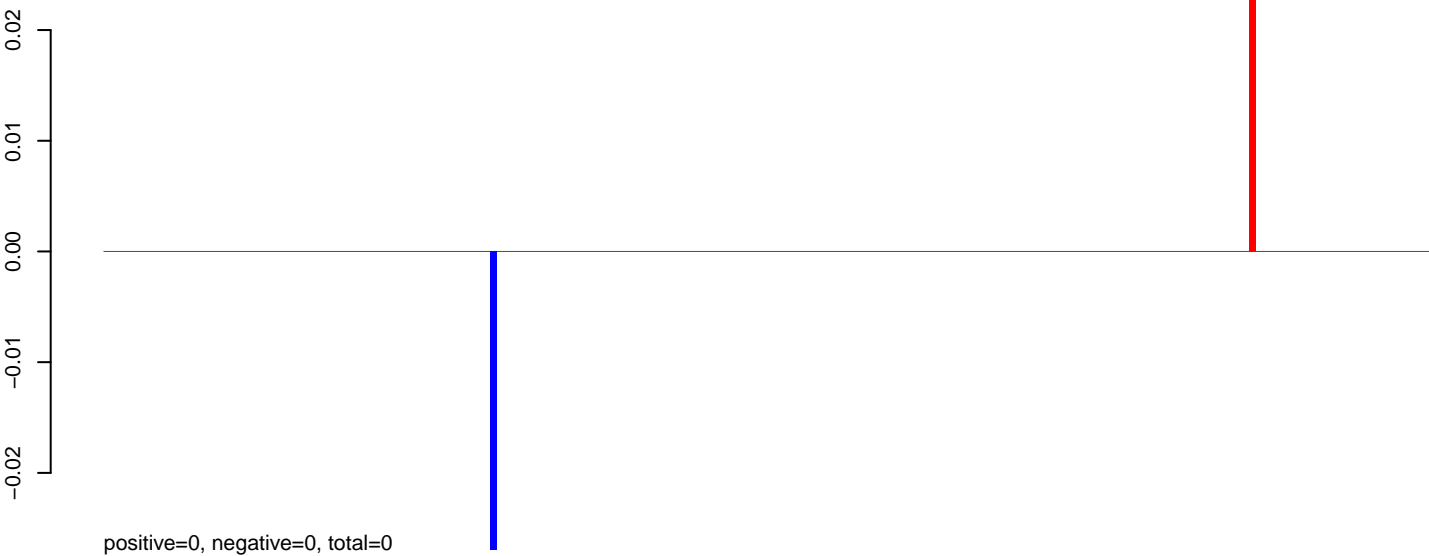
AeAeg_Ovary_GH.24_35.rep



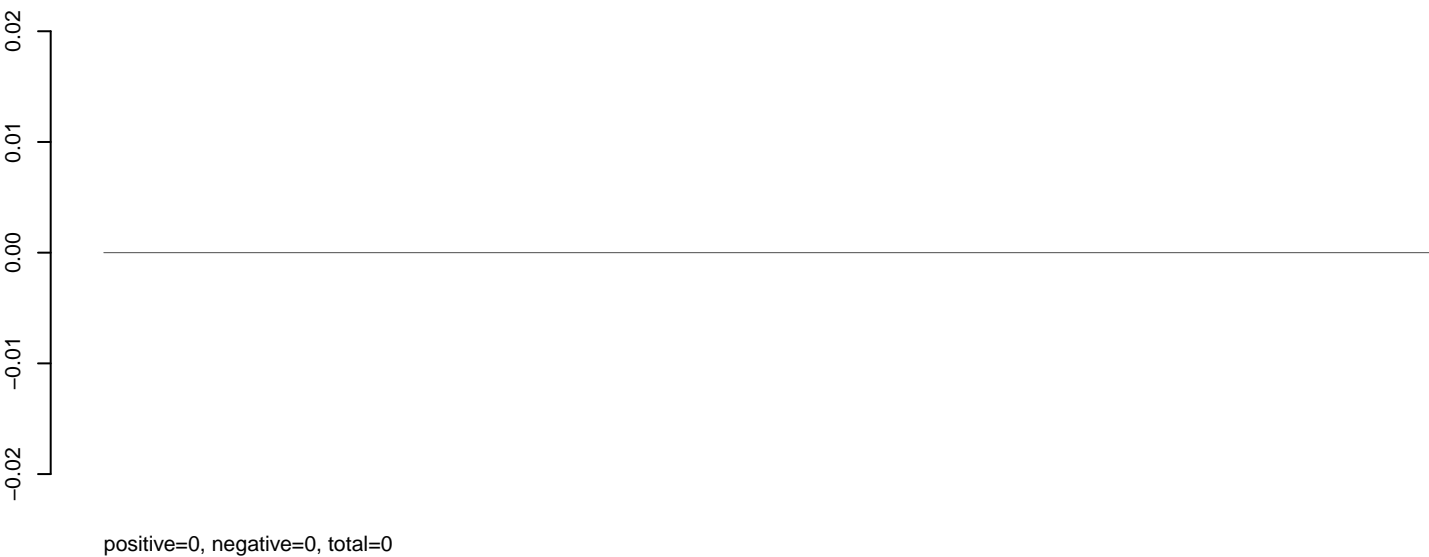
AeAeg_Ovary_GH.rep



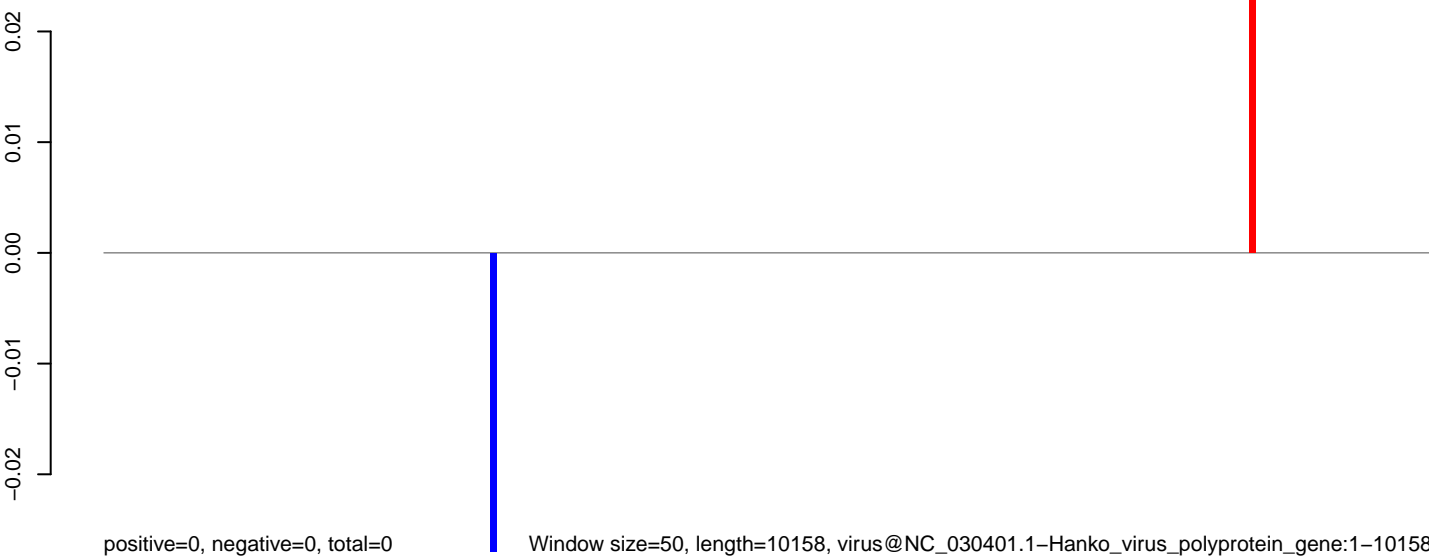
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

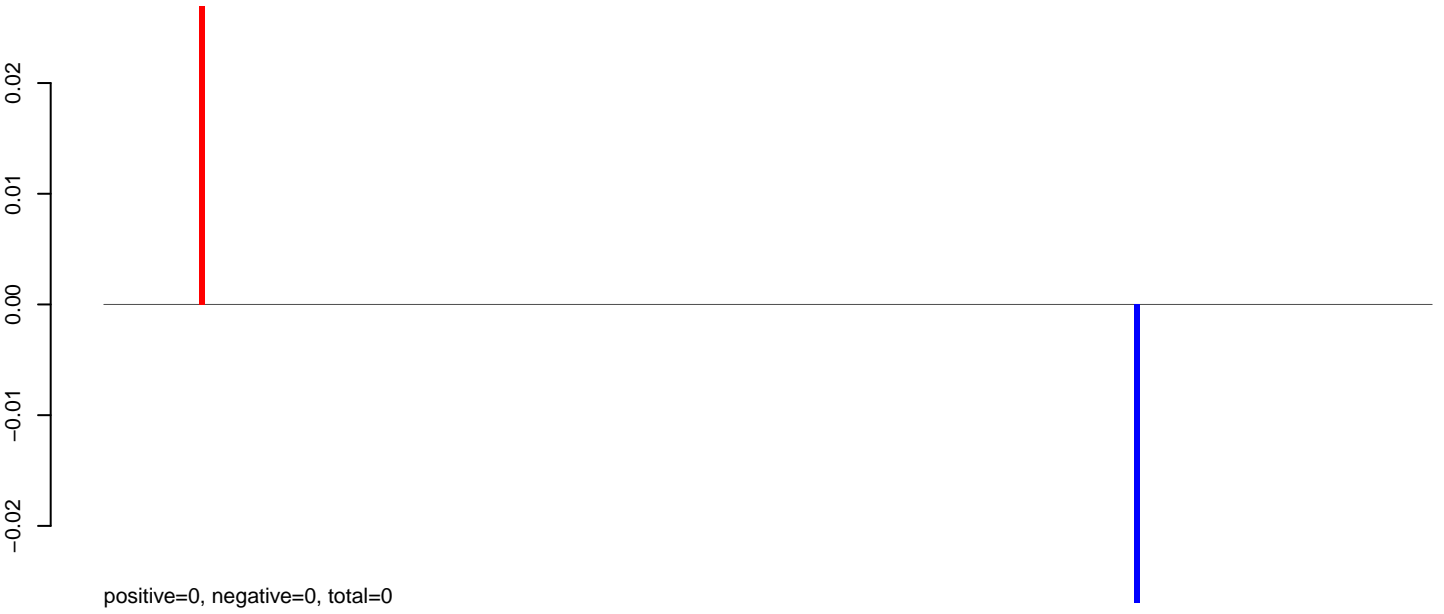


AeAeg_Ovary_GH.rep

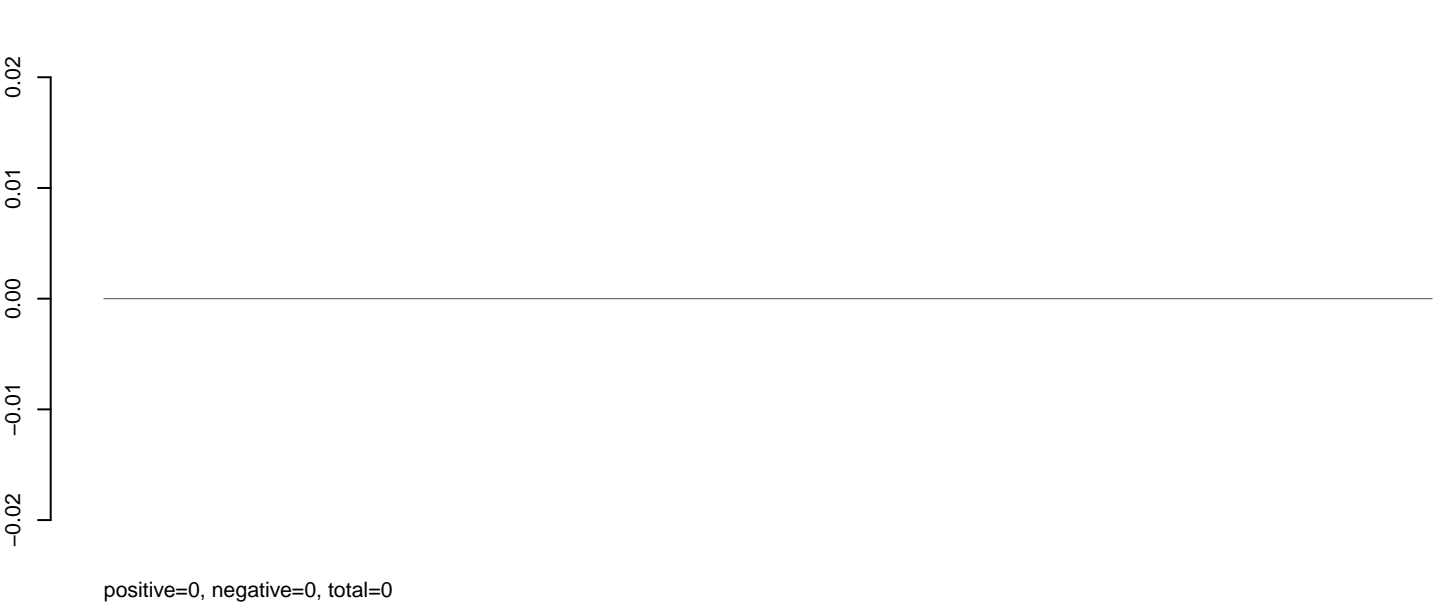


0 2000 4000 6000 8000 10000

AeAeg_Ovary_GH.18_23.rep



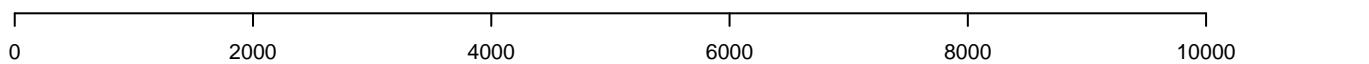
AeAeg_Ovary_GH.24_35.rep



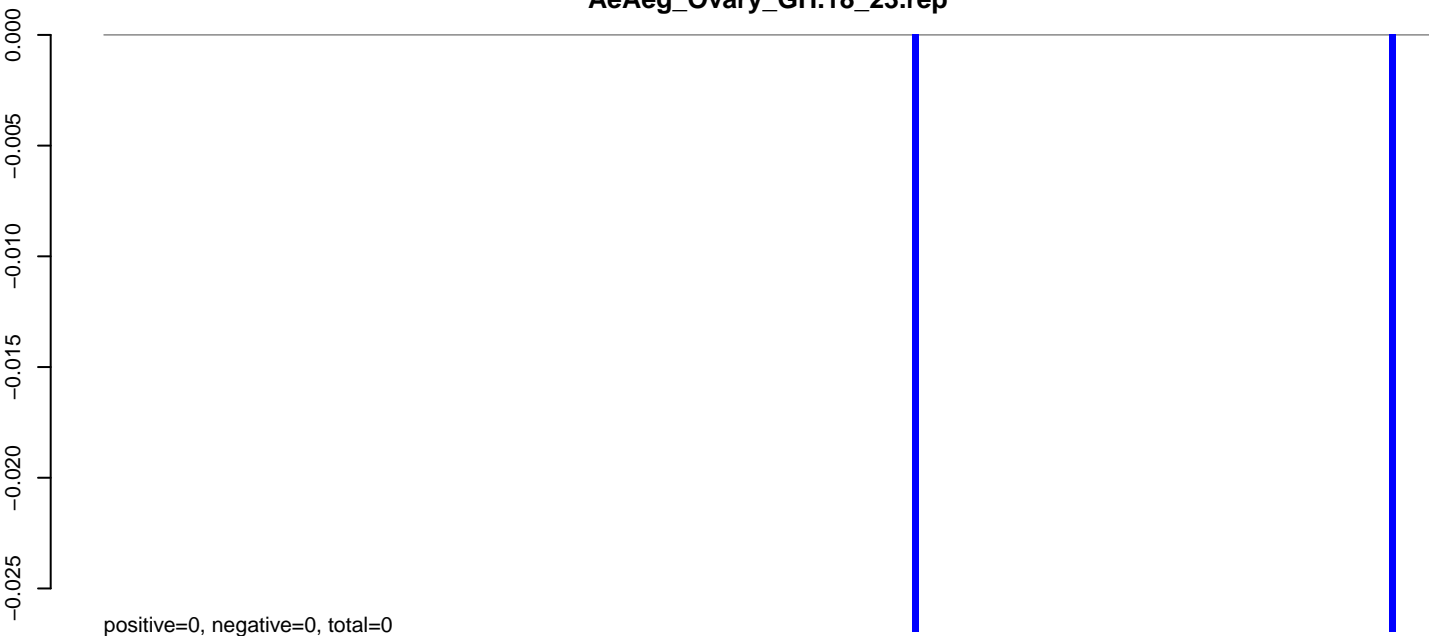
AeAeg_Ovary_GH.rep



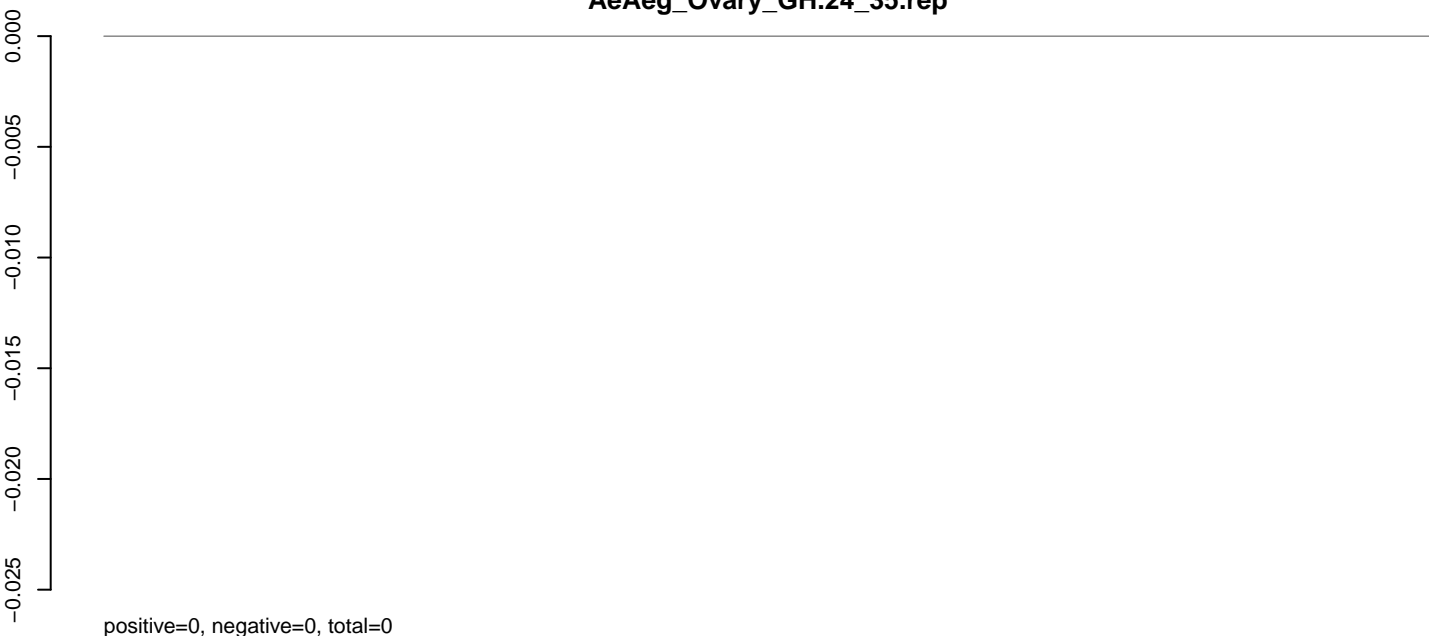
Window size=50, length=11190, virus@NC_025384.1-Culex tritaeniorhynchus rhabdovirus RNA strain TY:1-11190



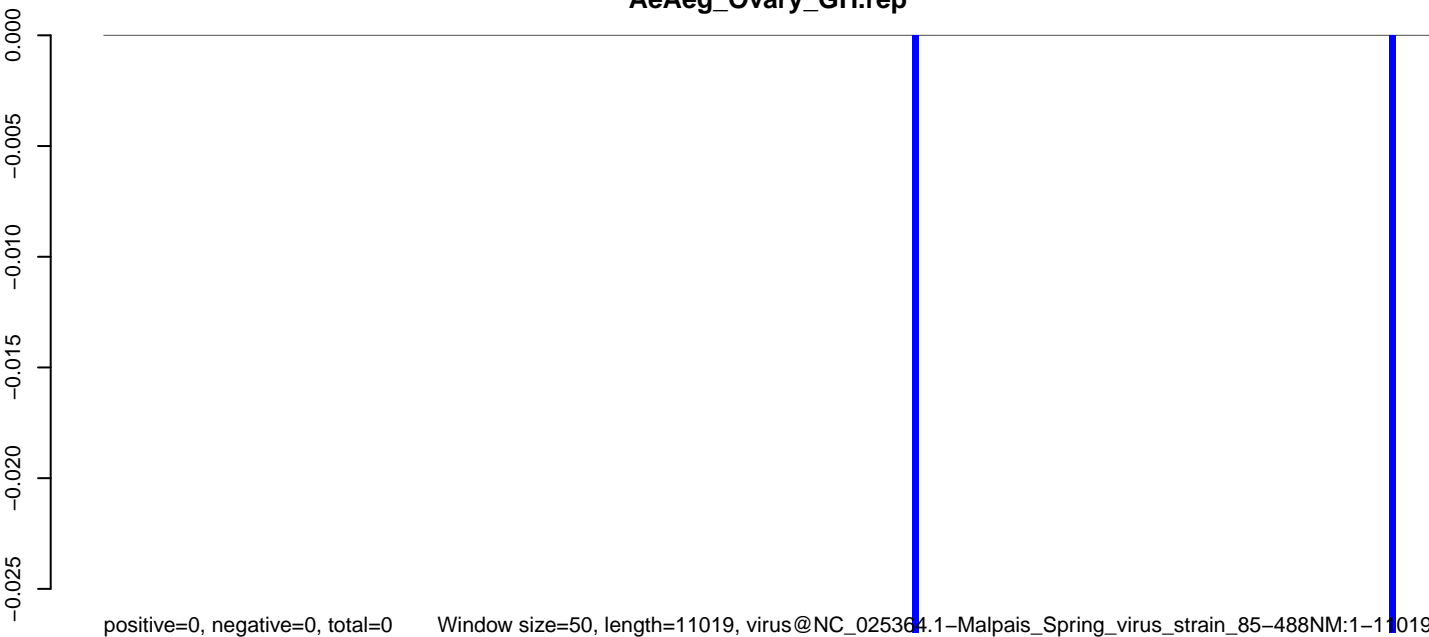
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



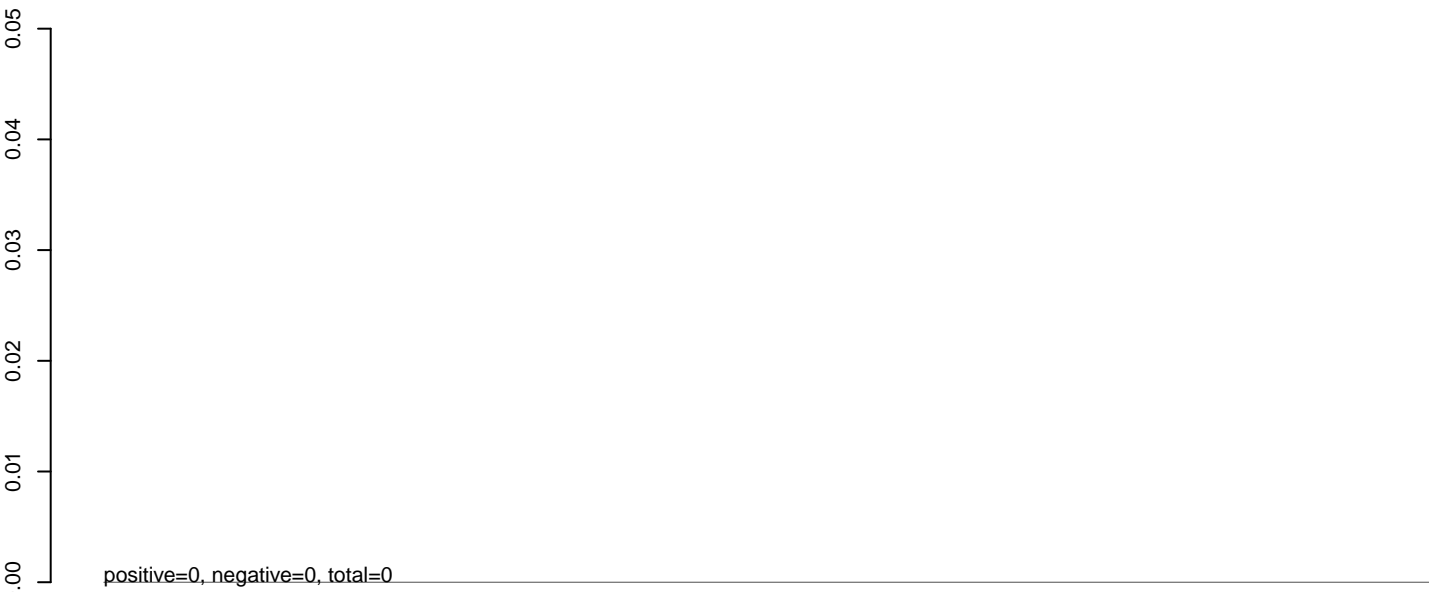
Window size=50, length=11019, virus@NC_025364.1-Malpais_Spring_virus_strain_85-488NM:1-11019

0 2000 4000 6000 8000 10000

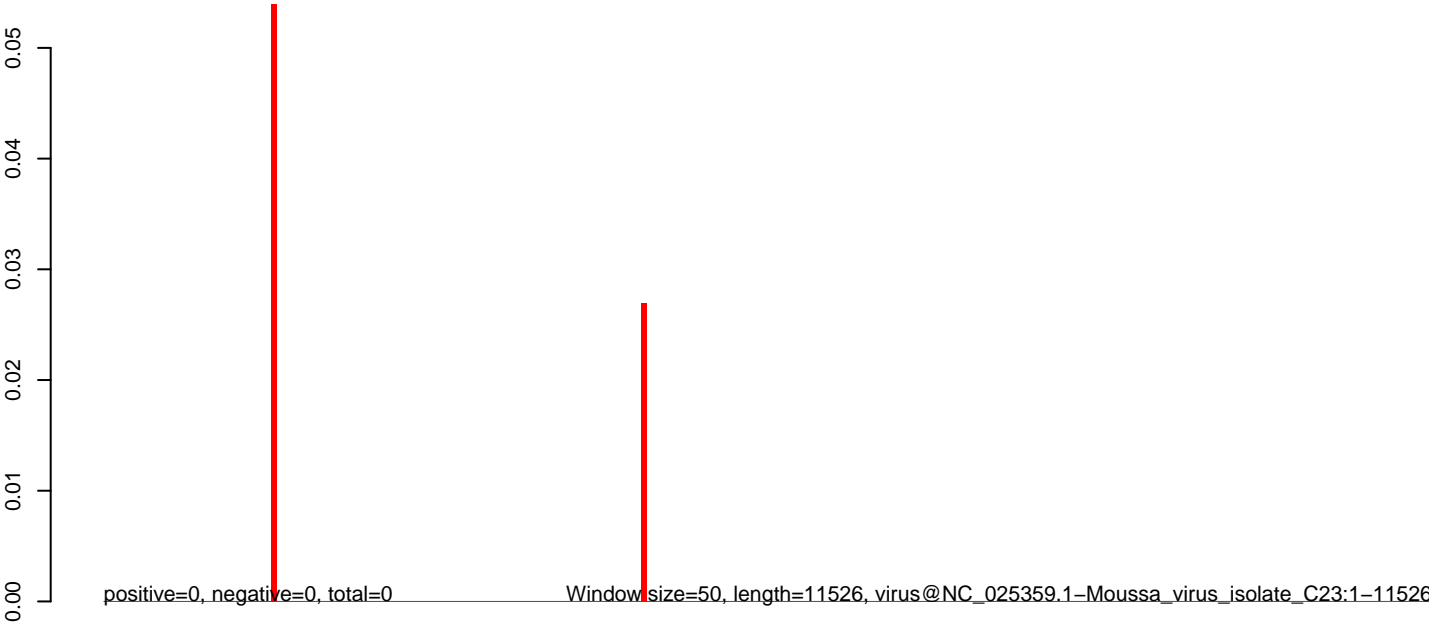
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

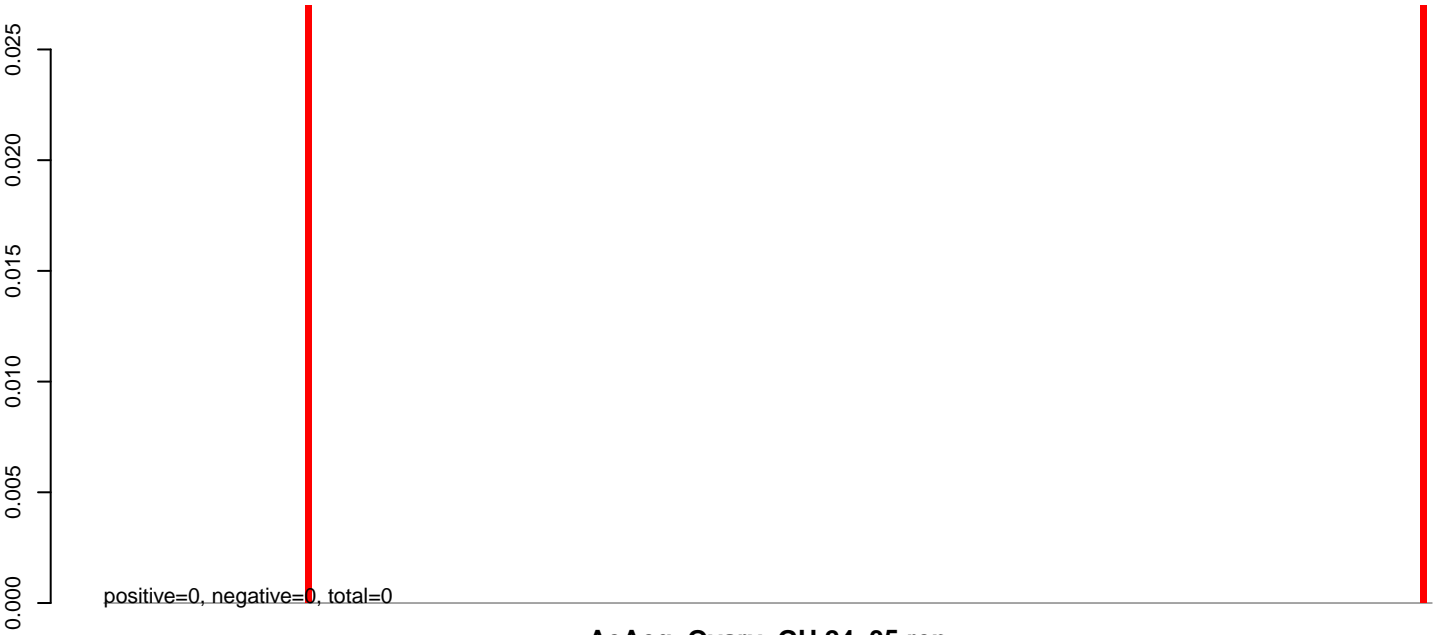


AeAeg_Ovary_GH.rep

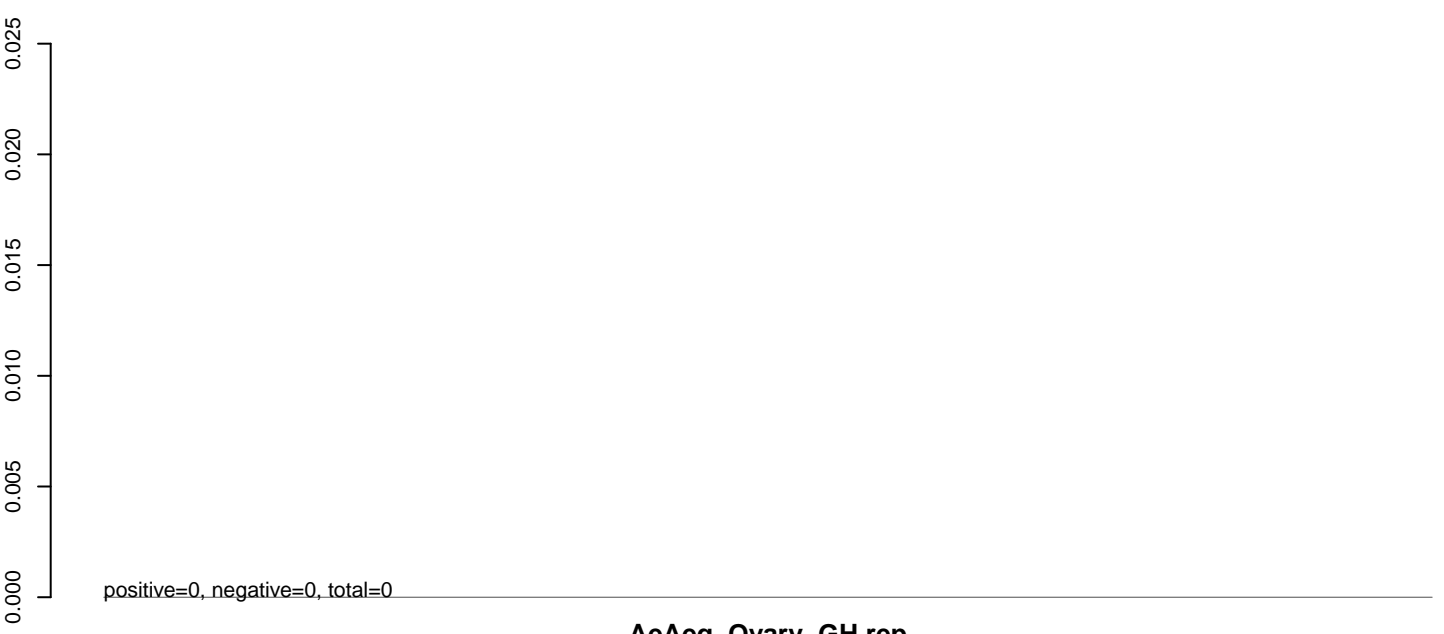


0 2000 4000 6000 8000 10000 12000

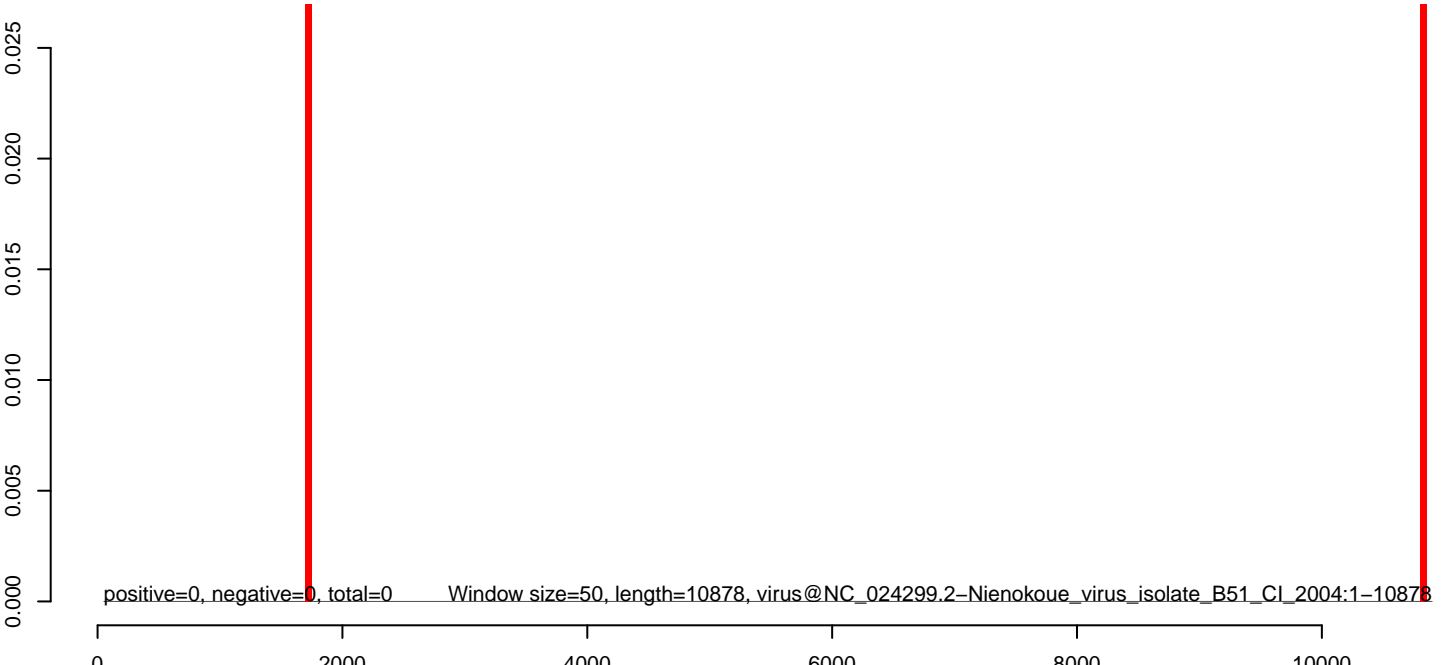
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



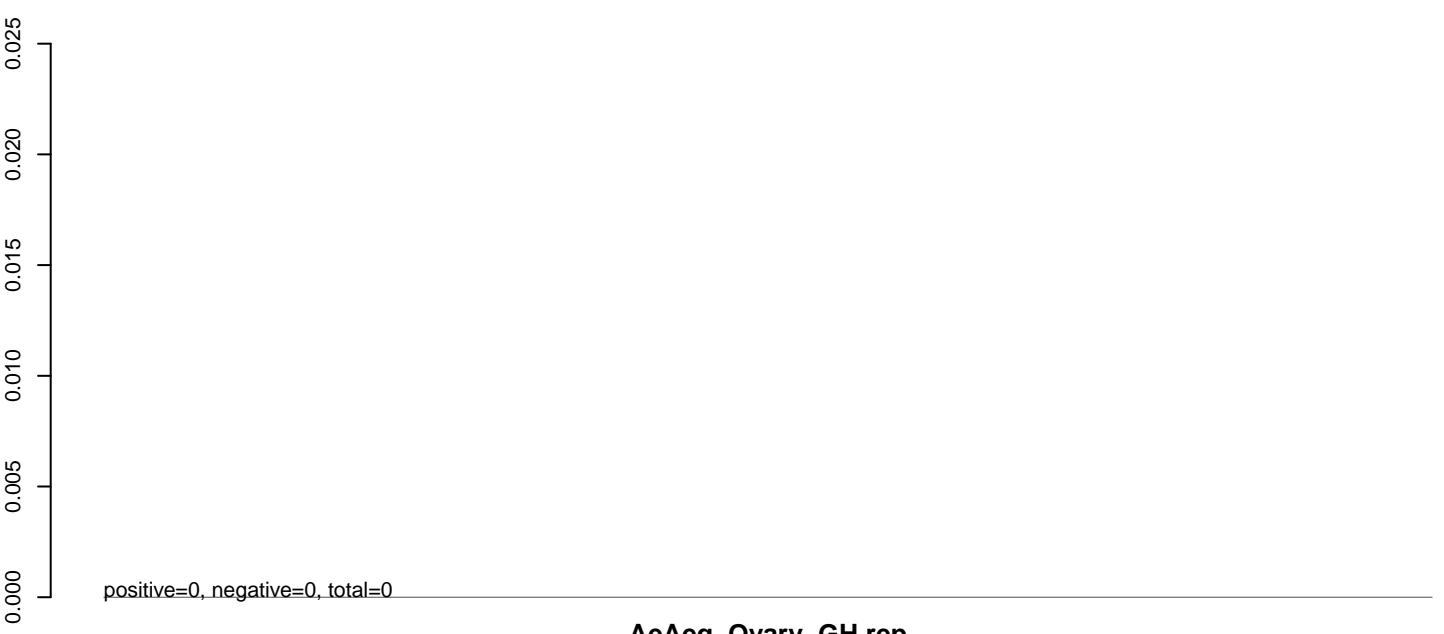
AeAeg_Ovary_GH.rep



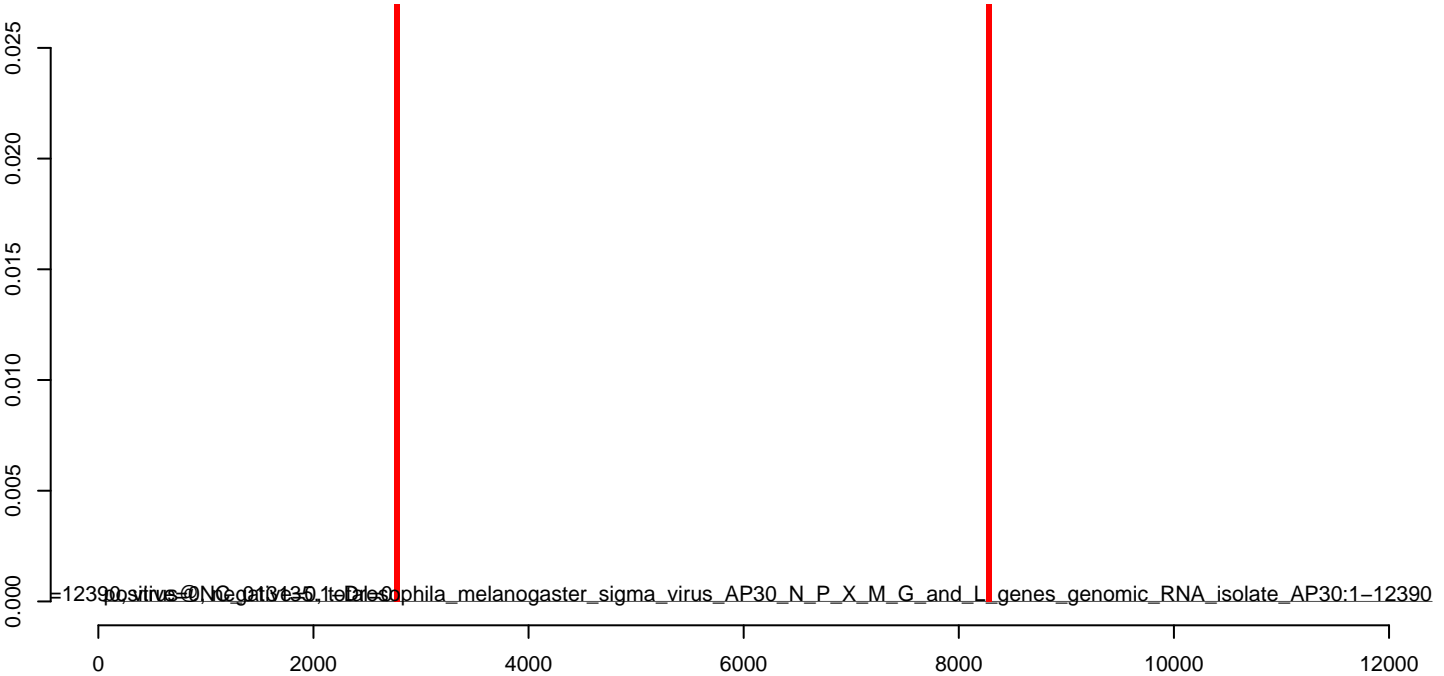
AeAeg_Ovary_GH.18_23.rep



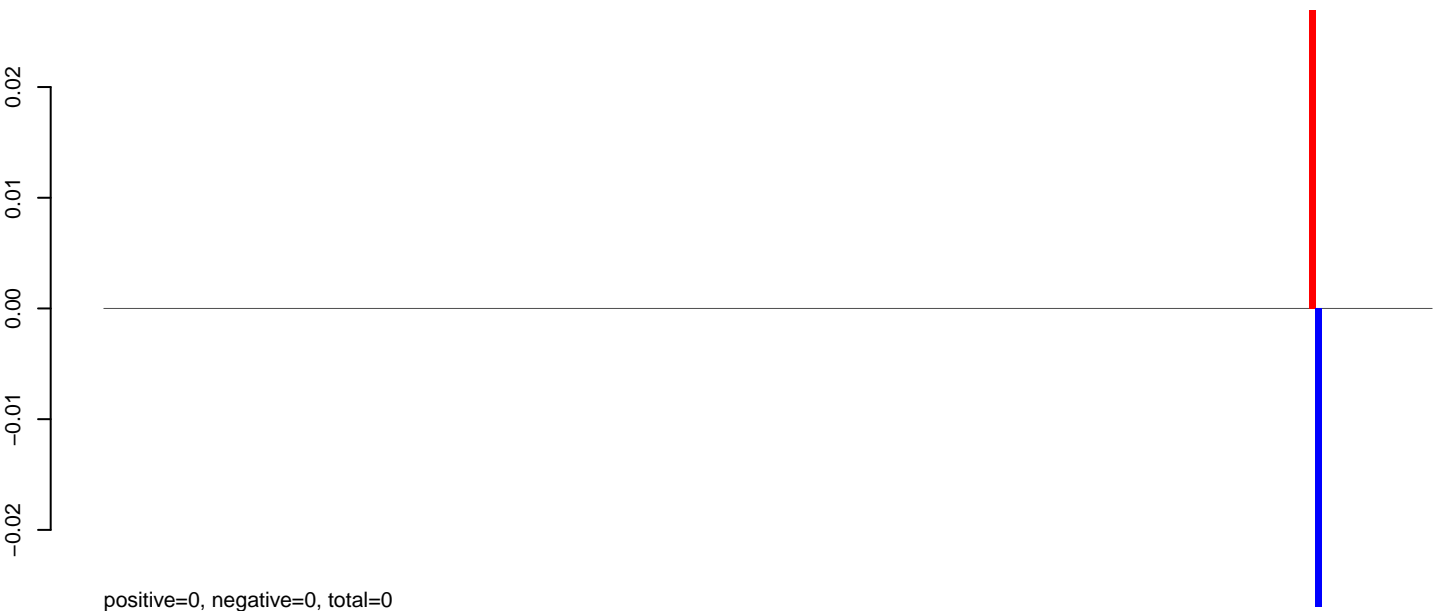
AeAeg_Ovary_GH.24_35.rep



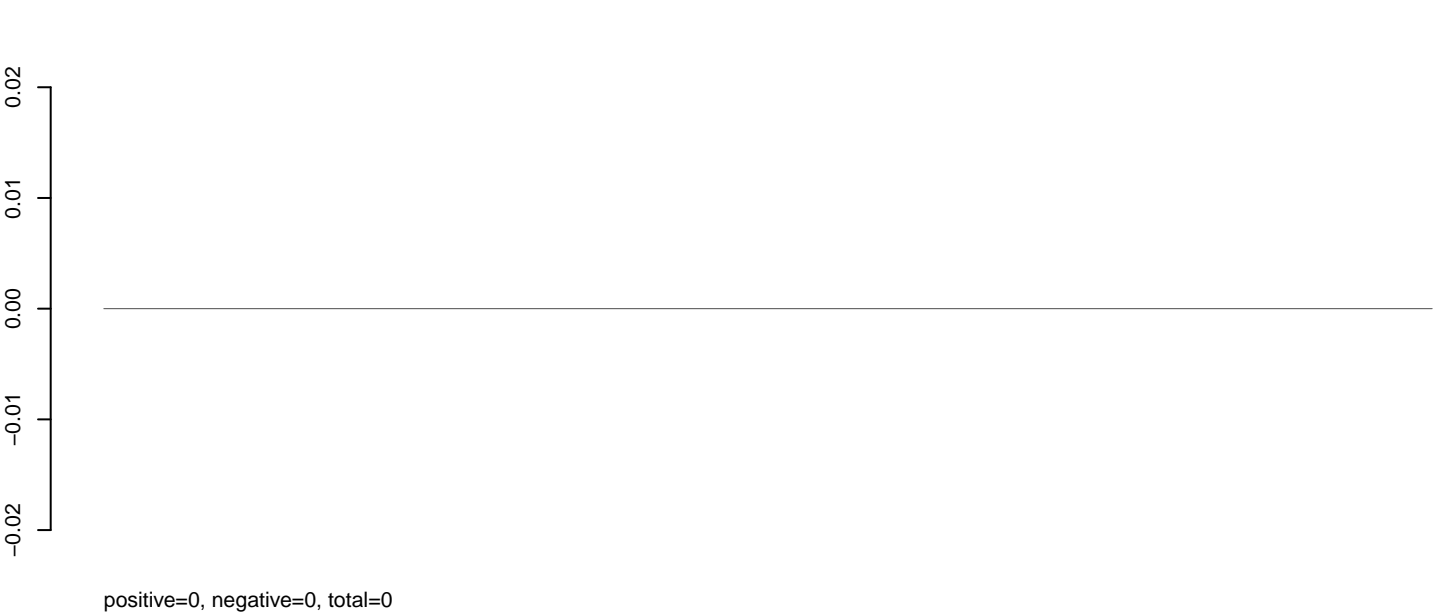
AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

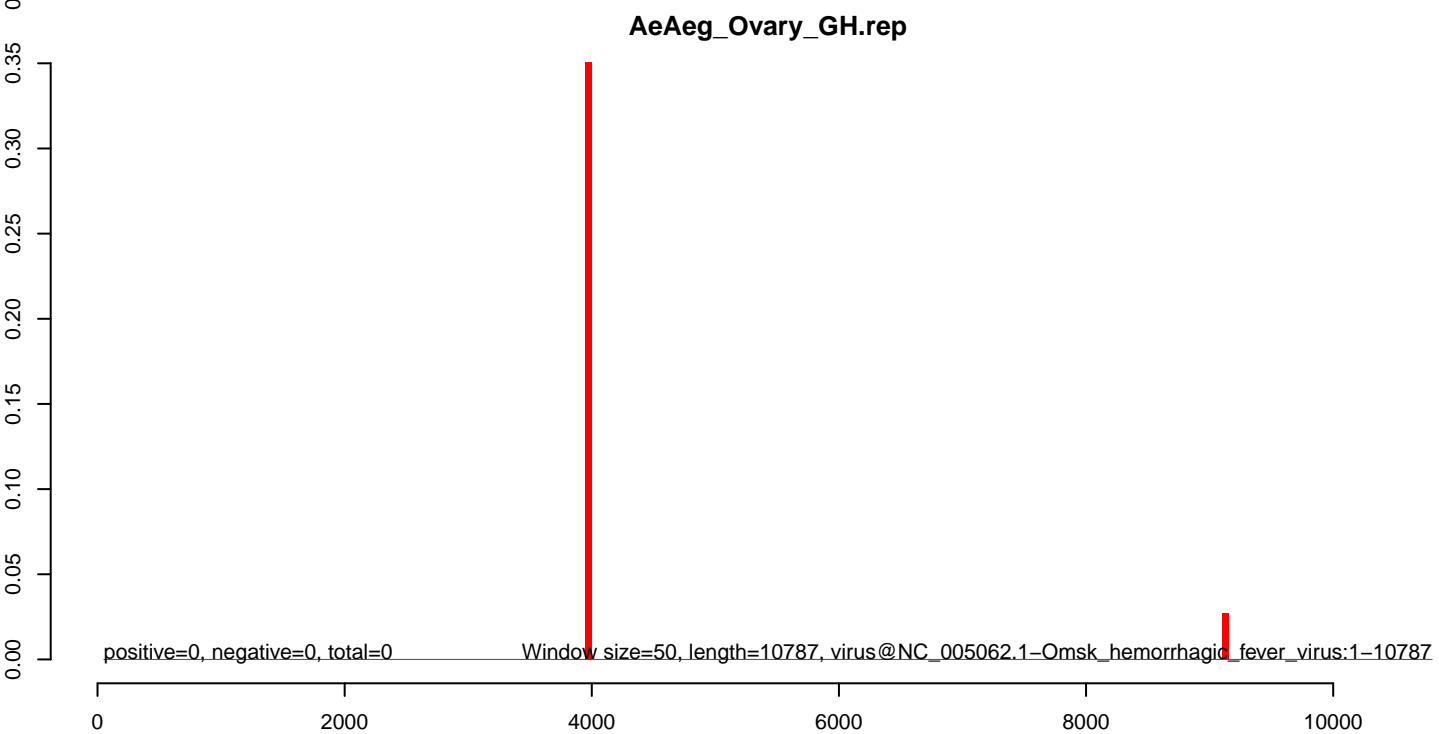
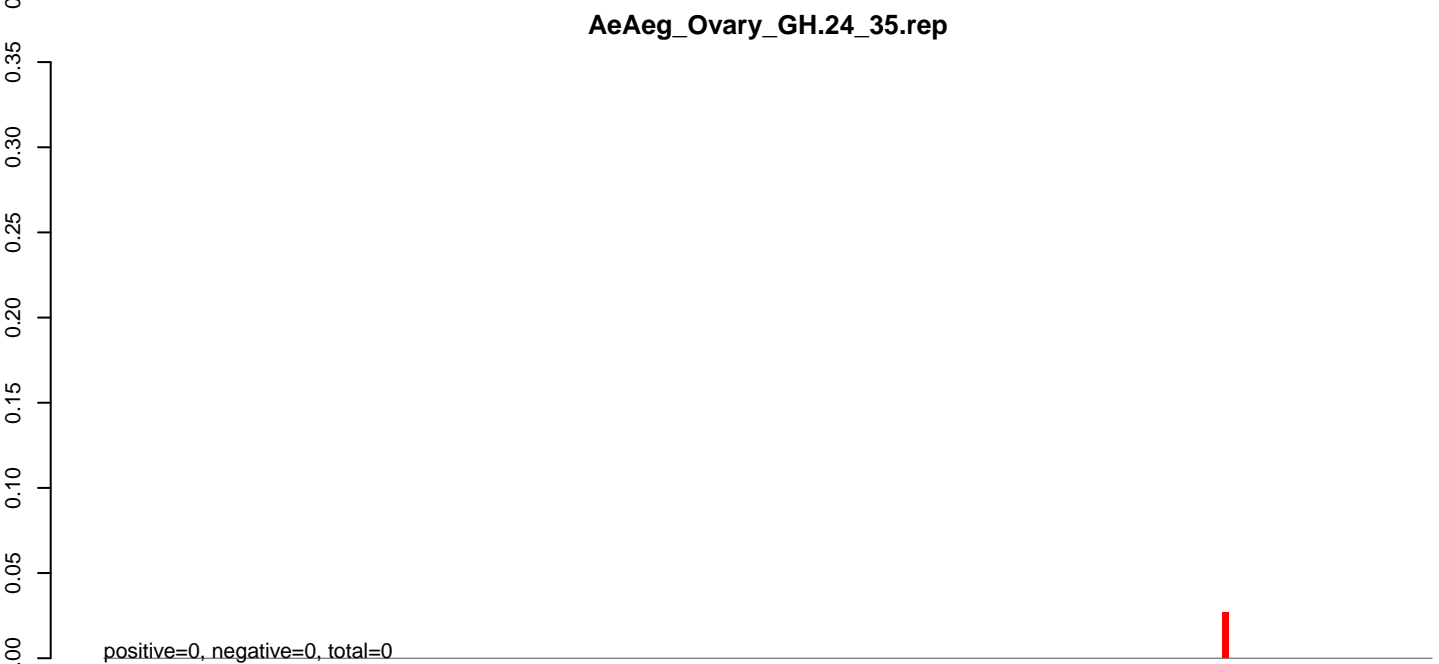
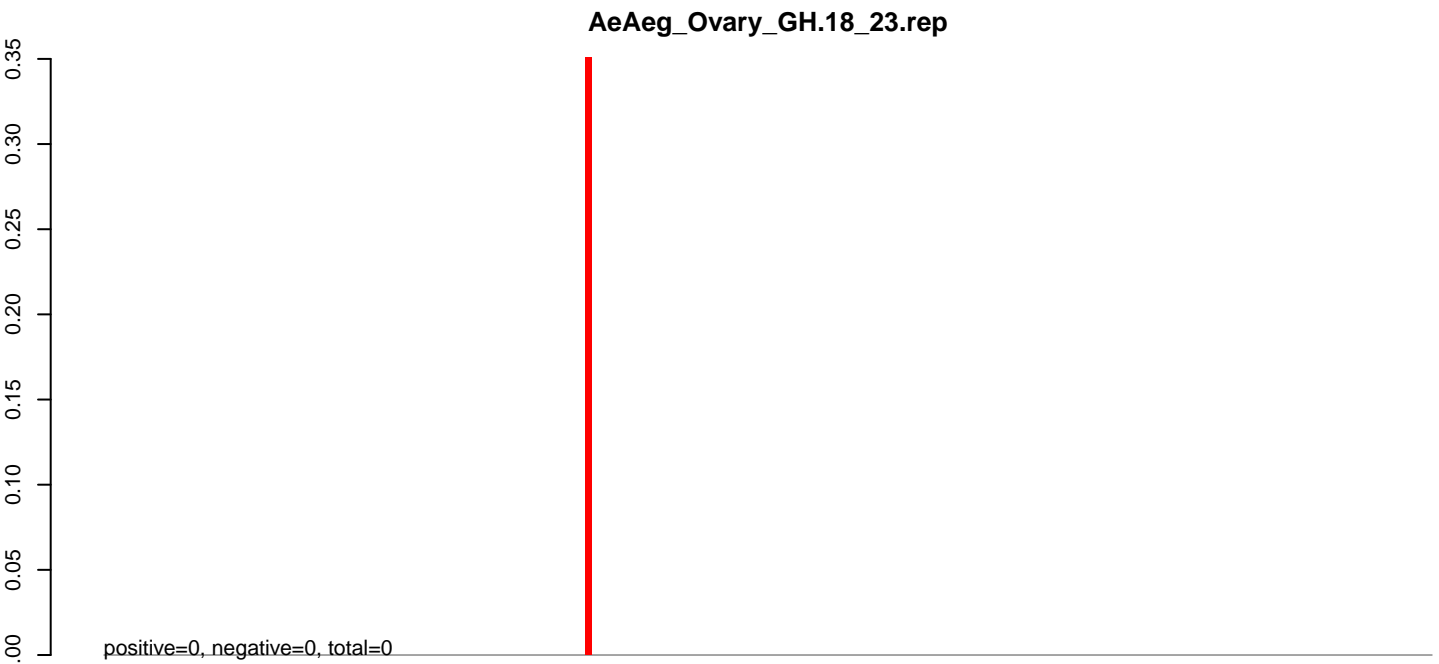


AeAeg_Ovary_GH.rep

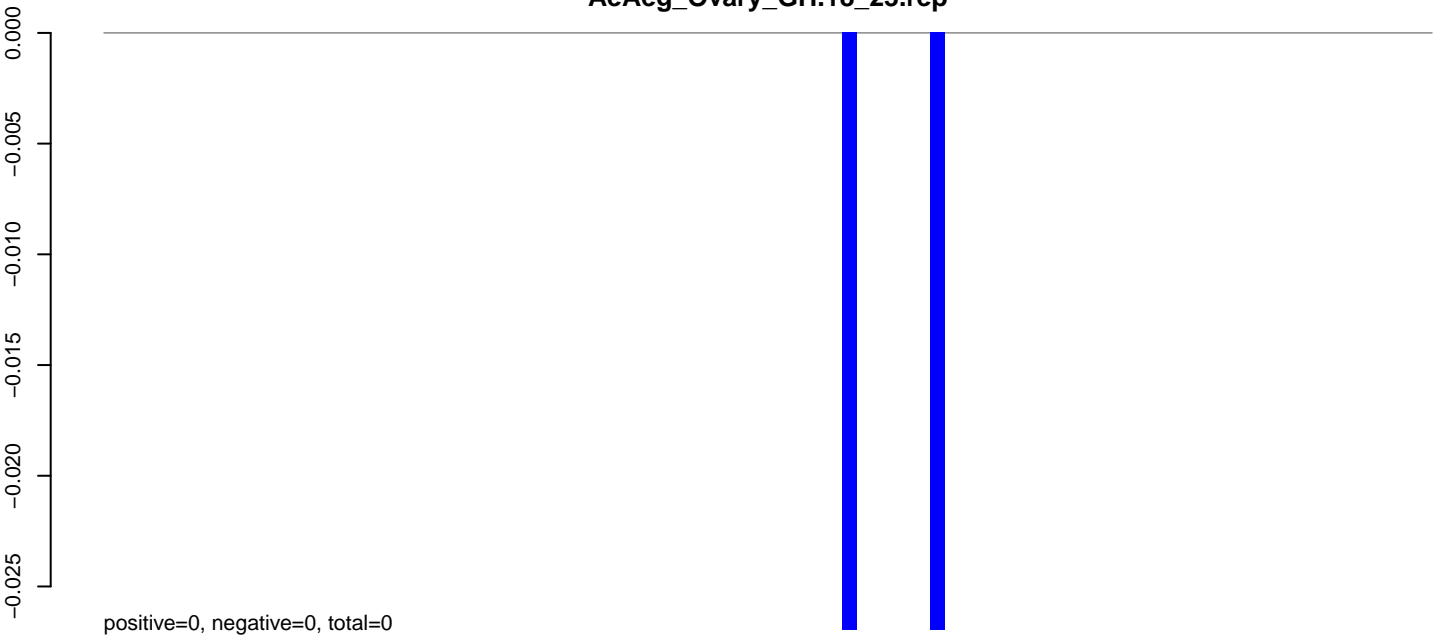


Window size=50, length=10874, virus@NC_009029.2-Kokobera_virus:1-10874

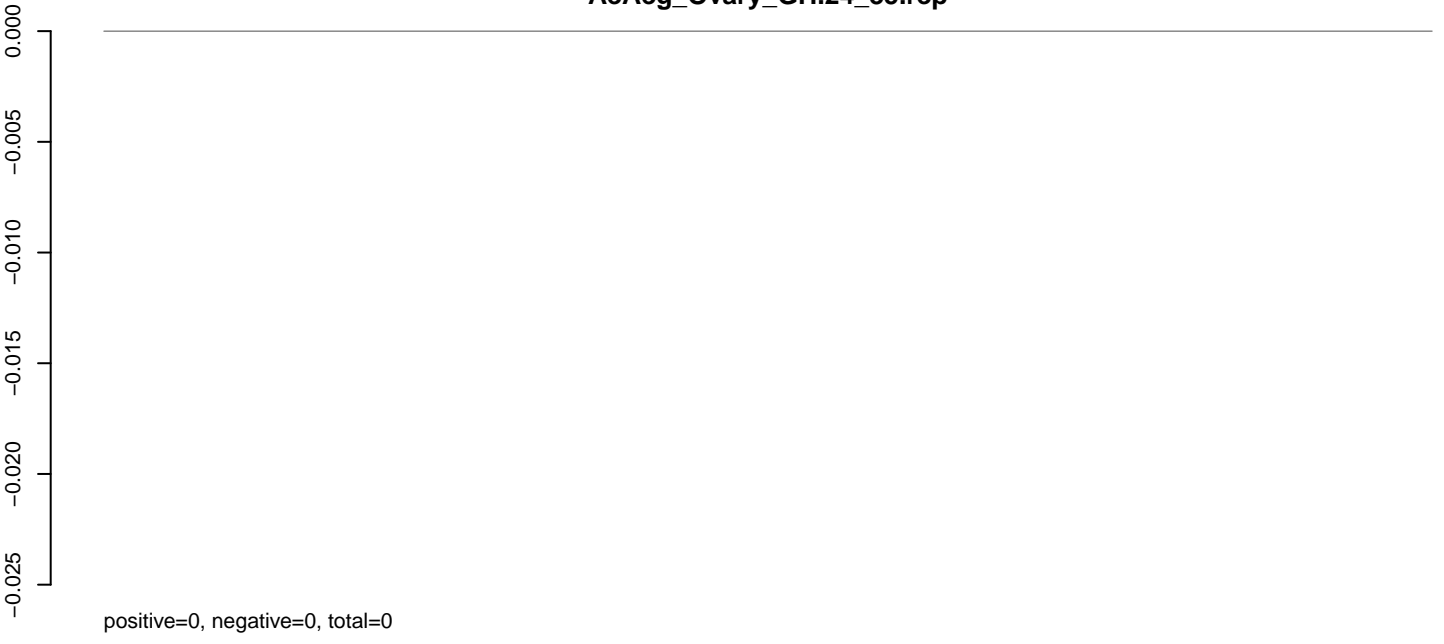
0 2000 4000 6000 8000 10000



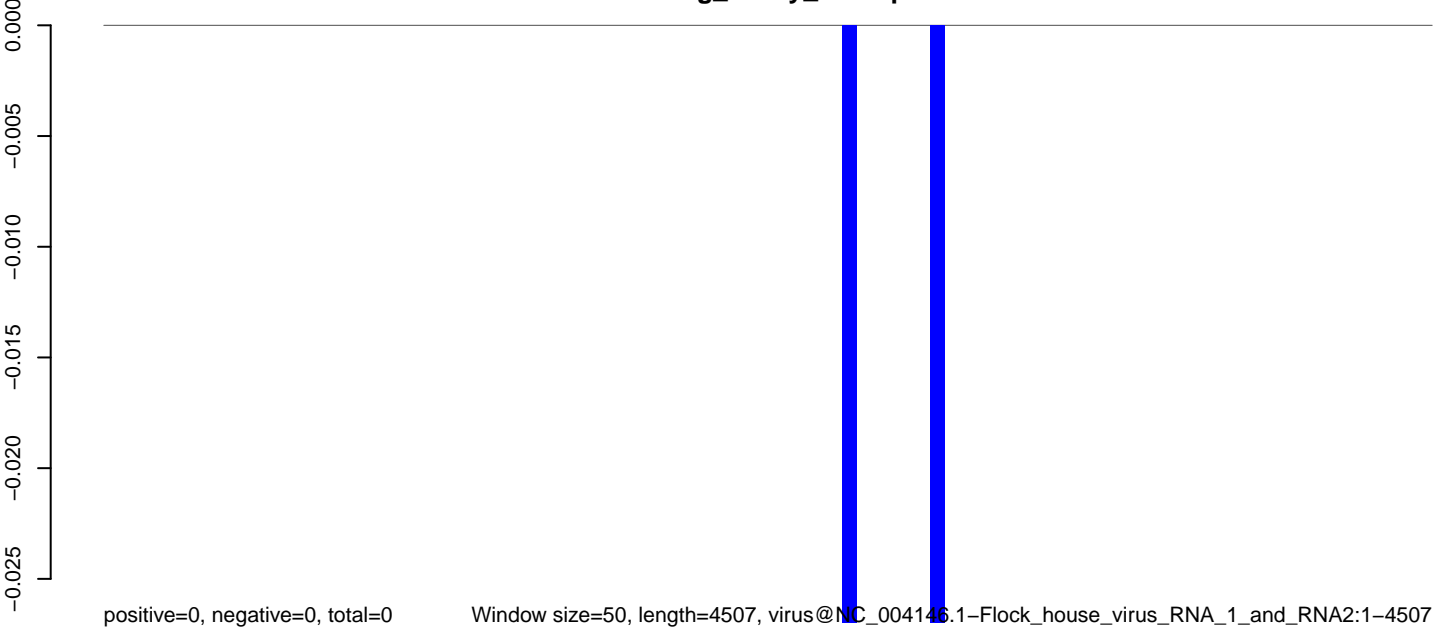
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

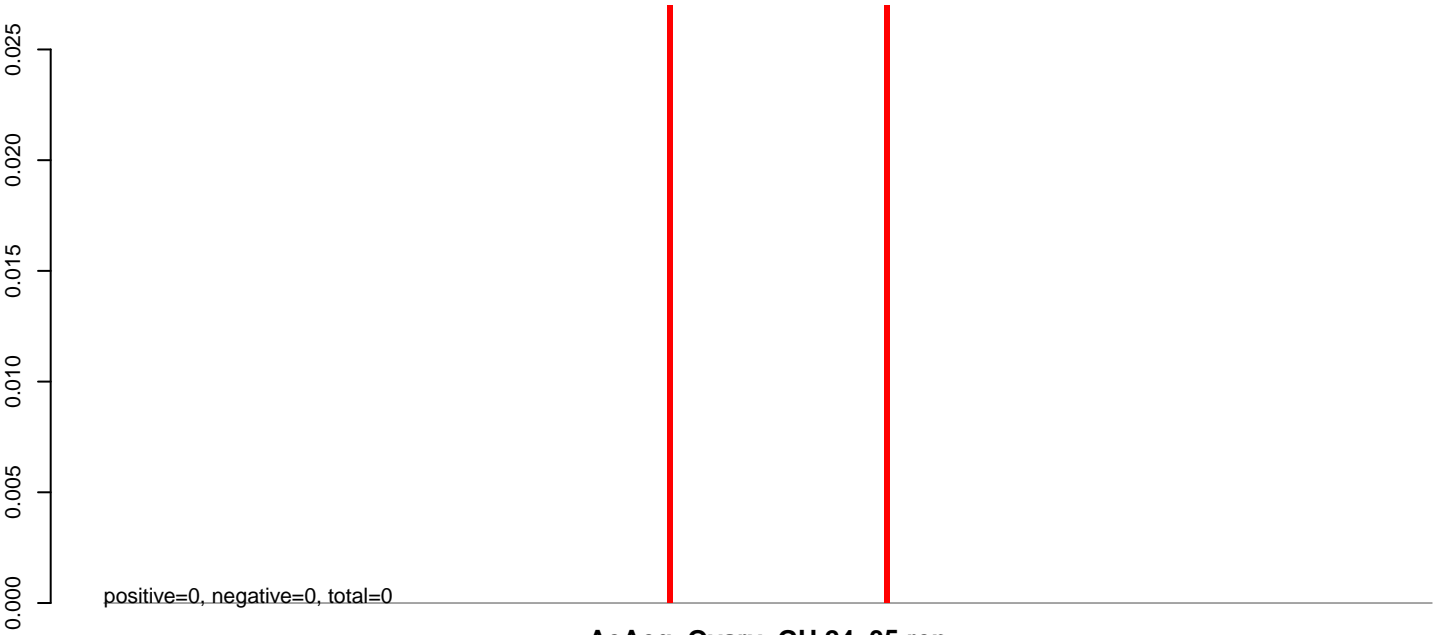


AeAeg_Ovary_GH.rep

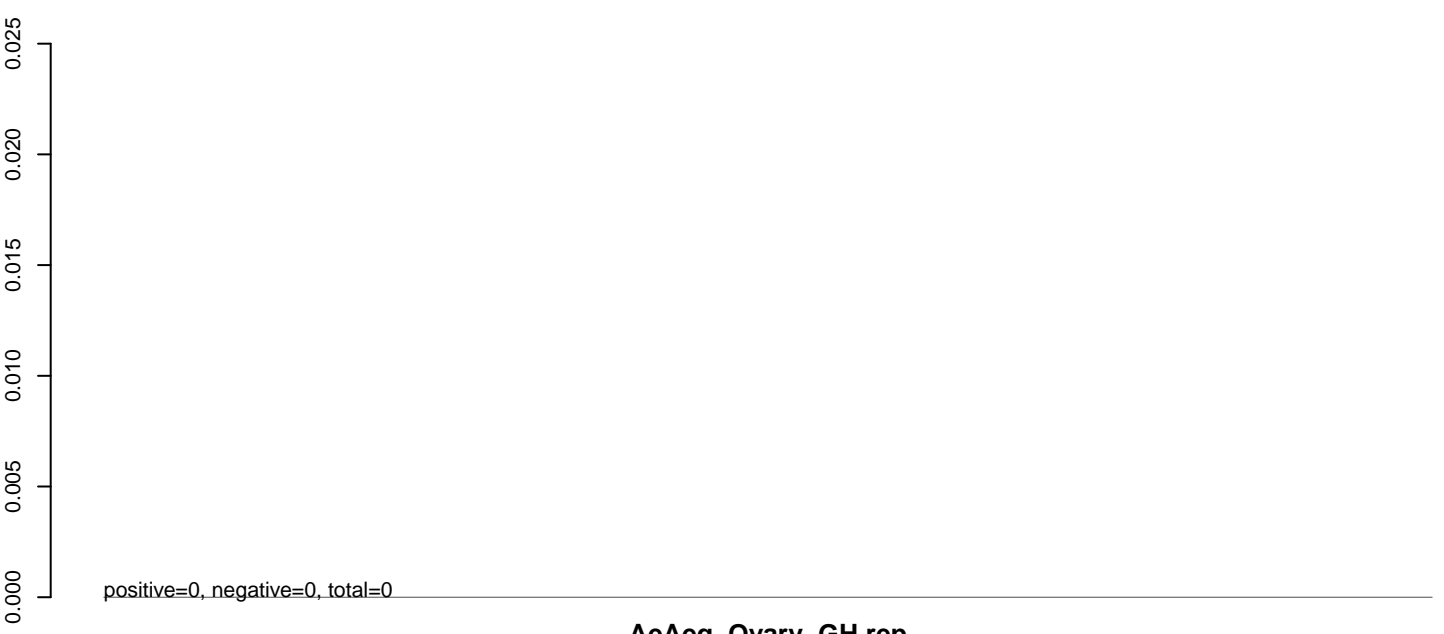


0 1000 2000 3000 4000

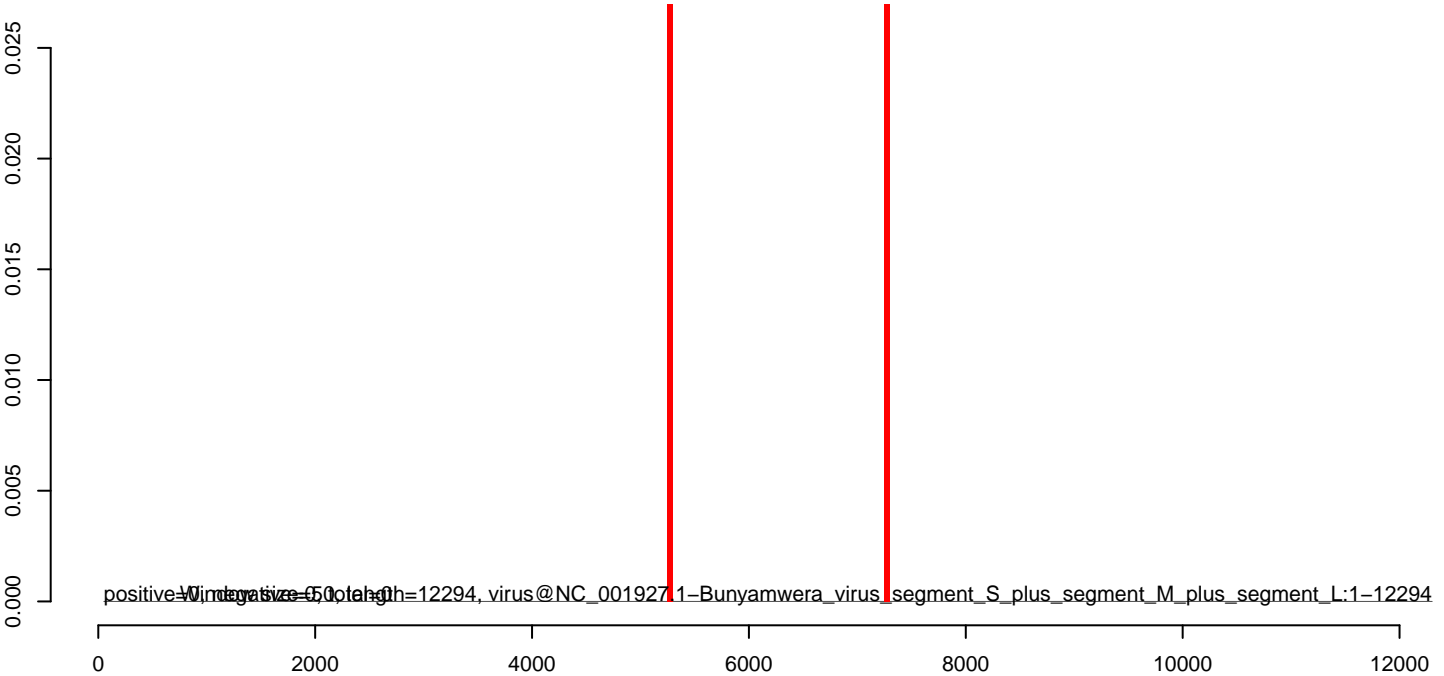
AeAeg_Ovary_GH.18_23.rep



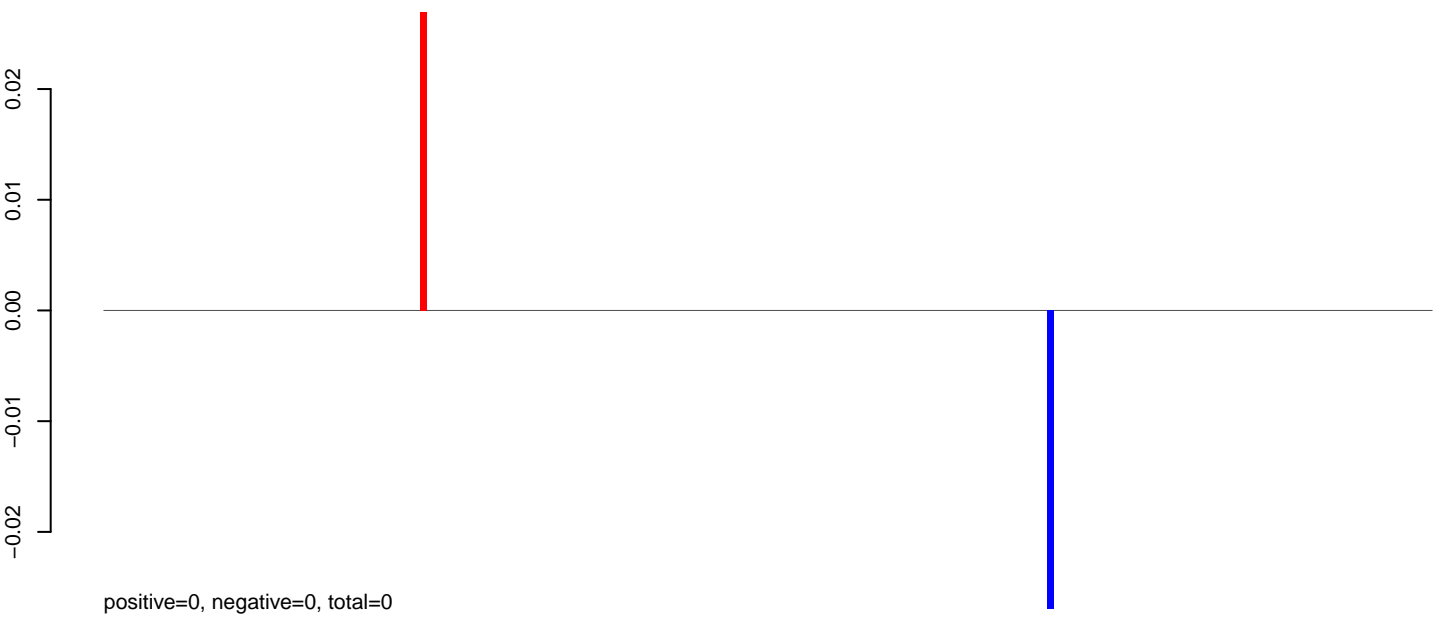
AeAeg_Ovary_GH.24_35.rep



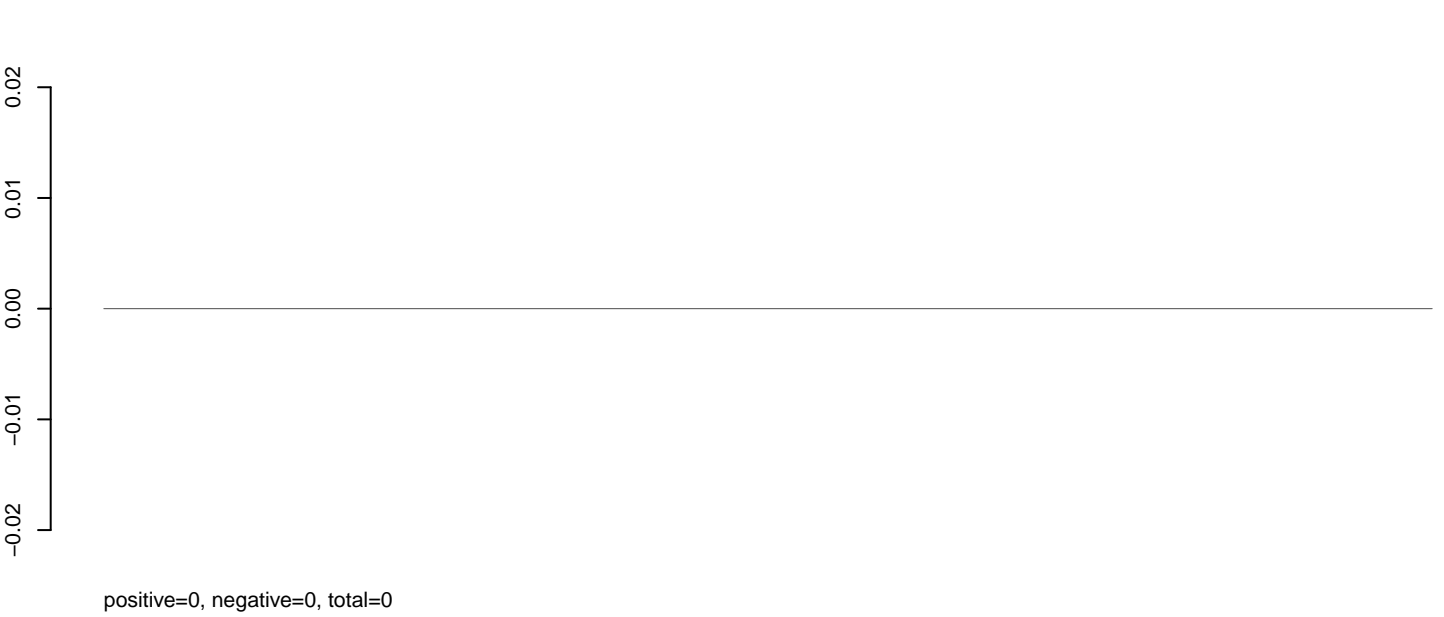
AeAeg_Ovary_GH.rep



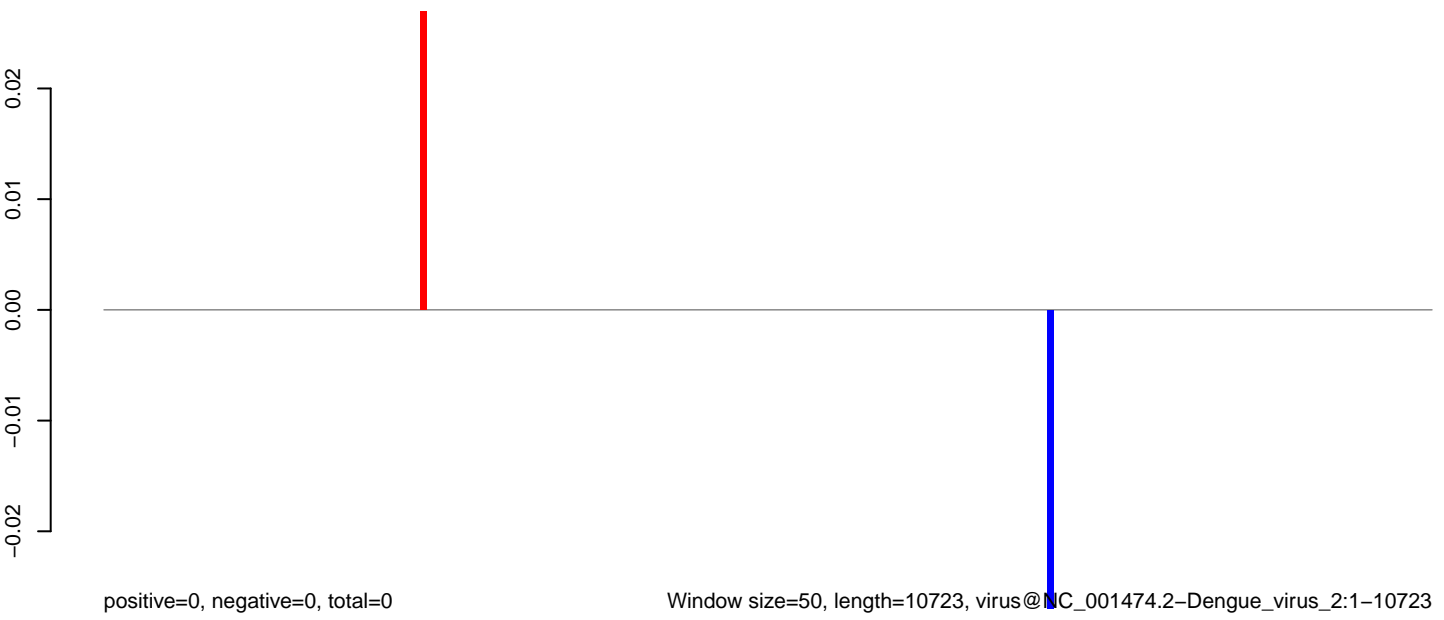
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

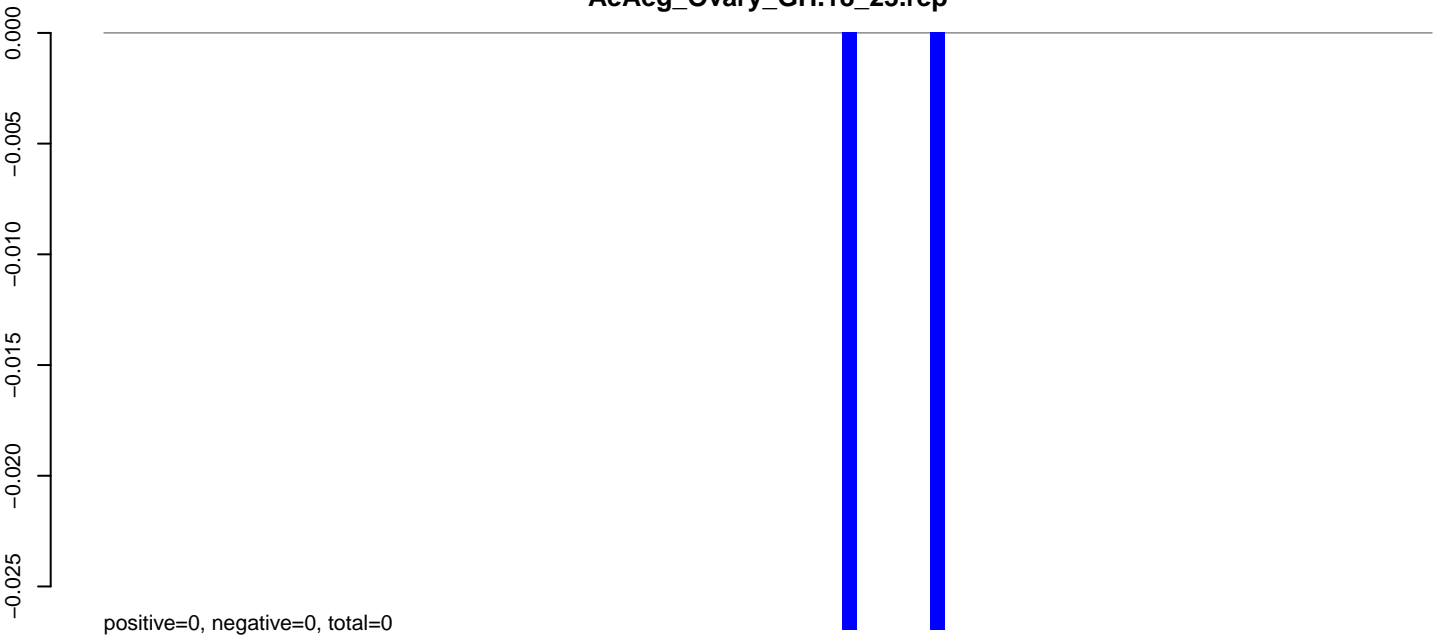


AeAeg_Ovary_GH.rep

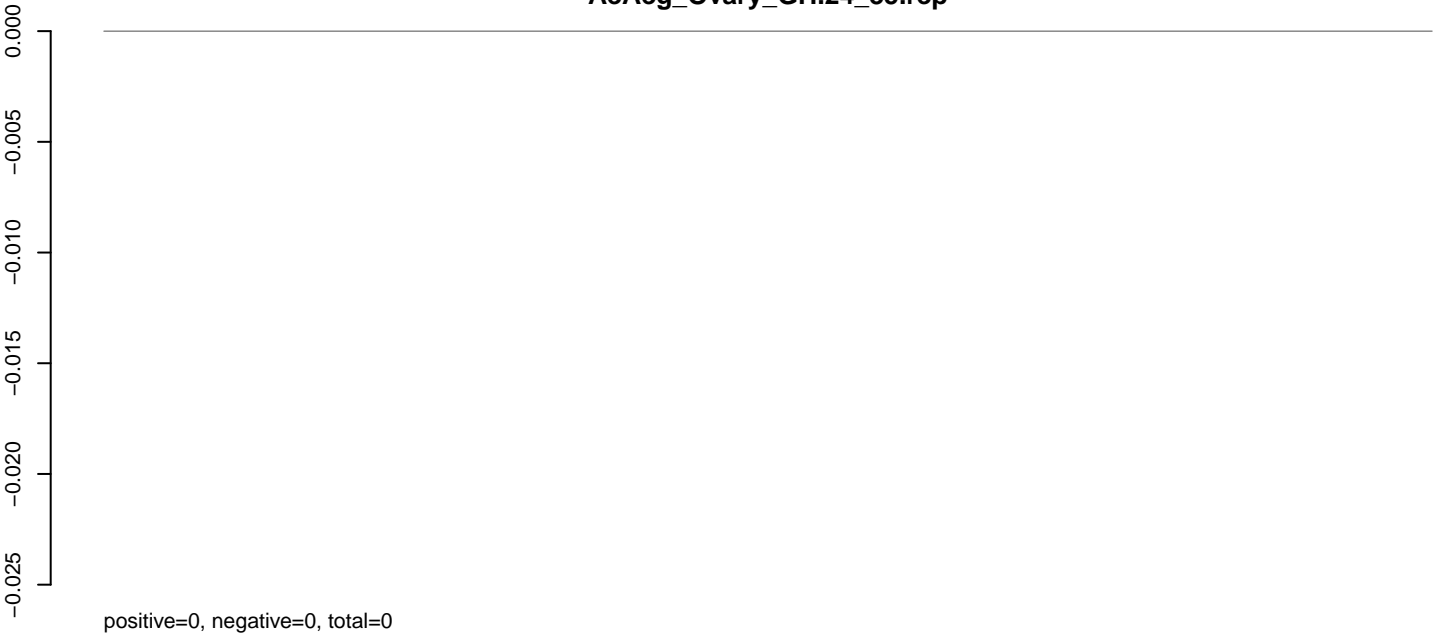


0 2000 4000 6000 8000 10000

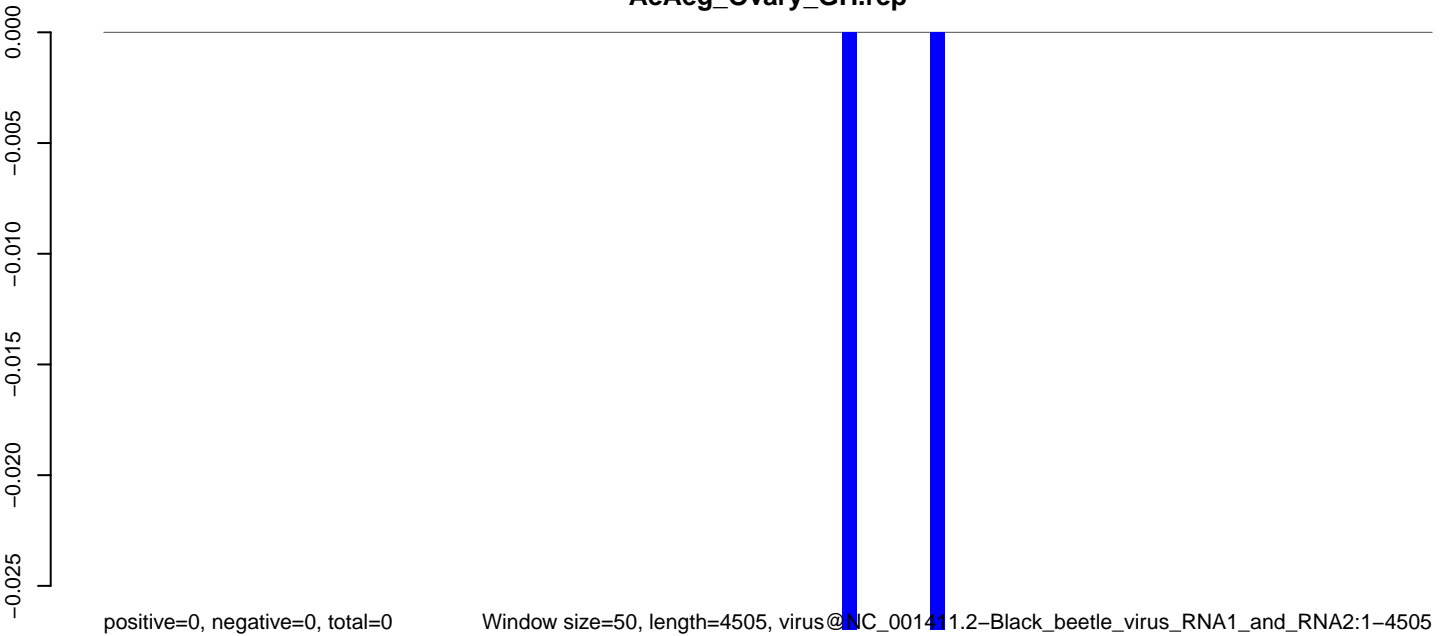
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

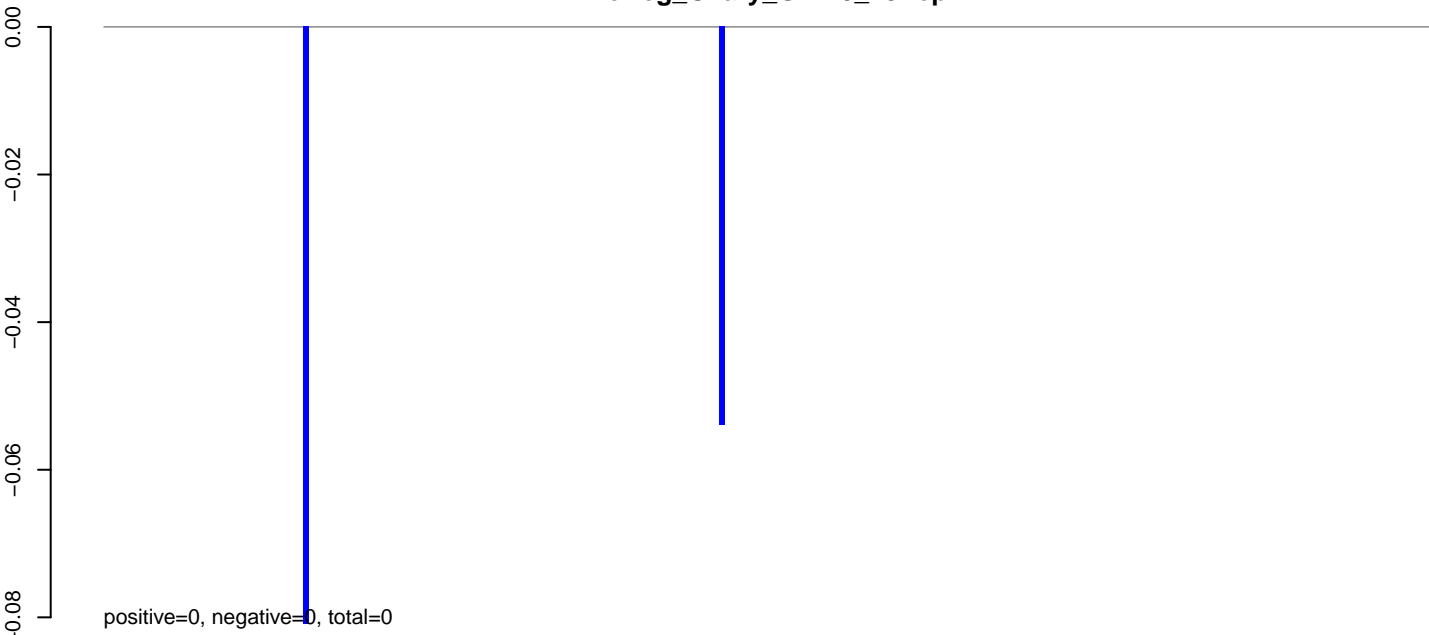


AeAeg_Ovary_GH.rep

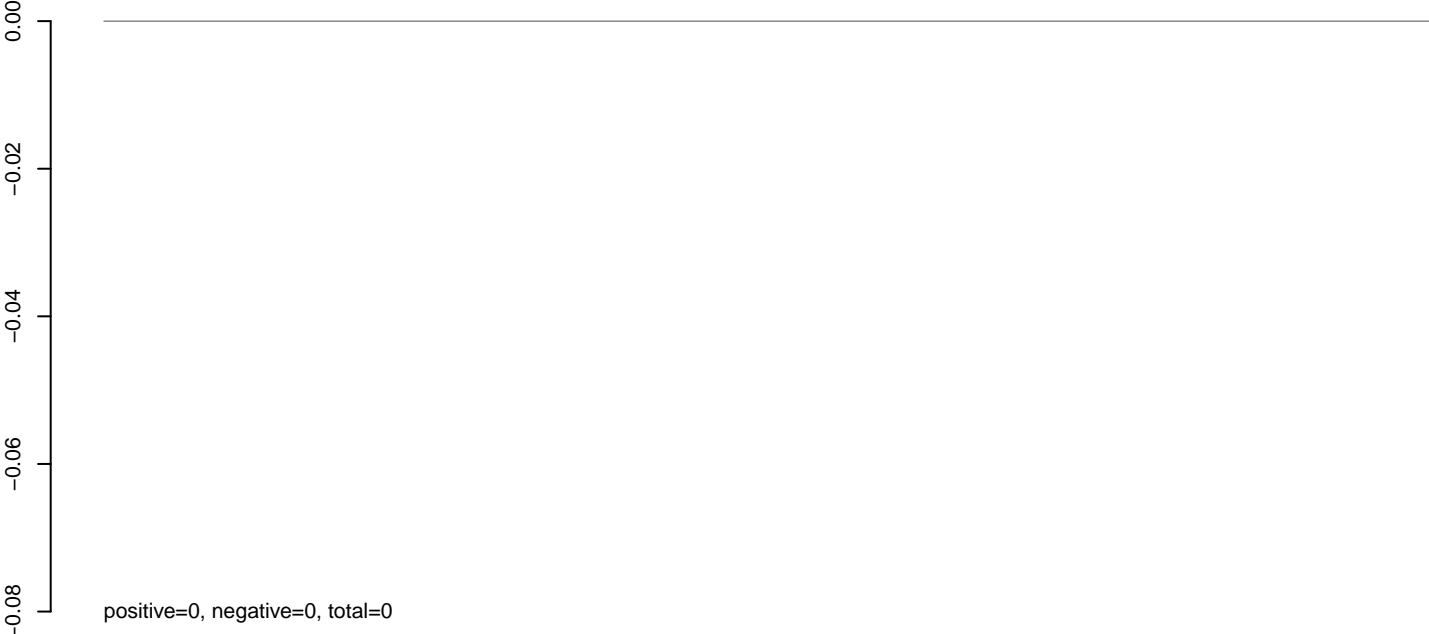


0 1000 2000 3000 4000

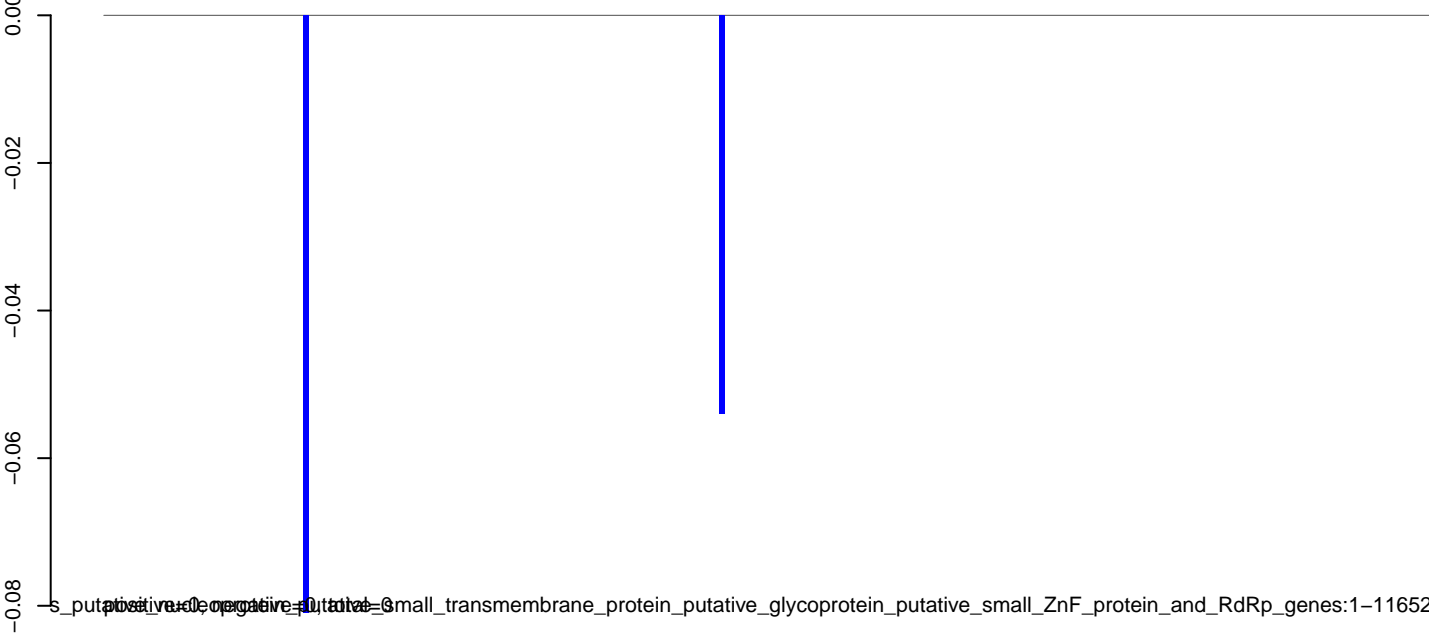
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



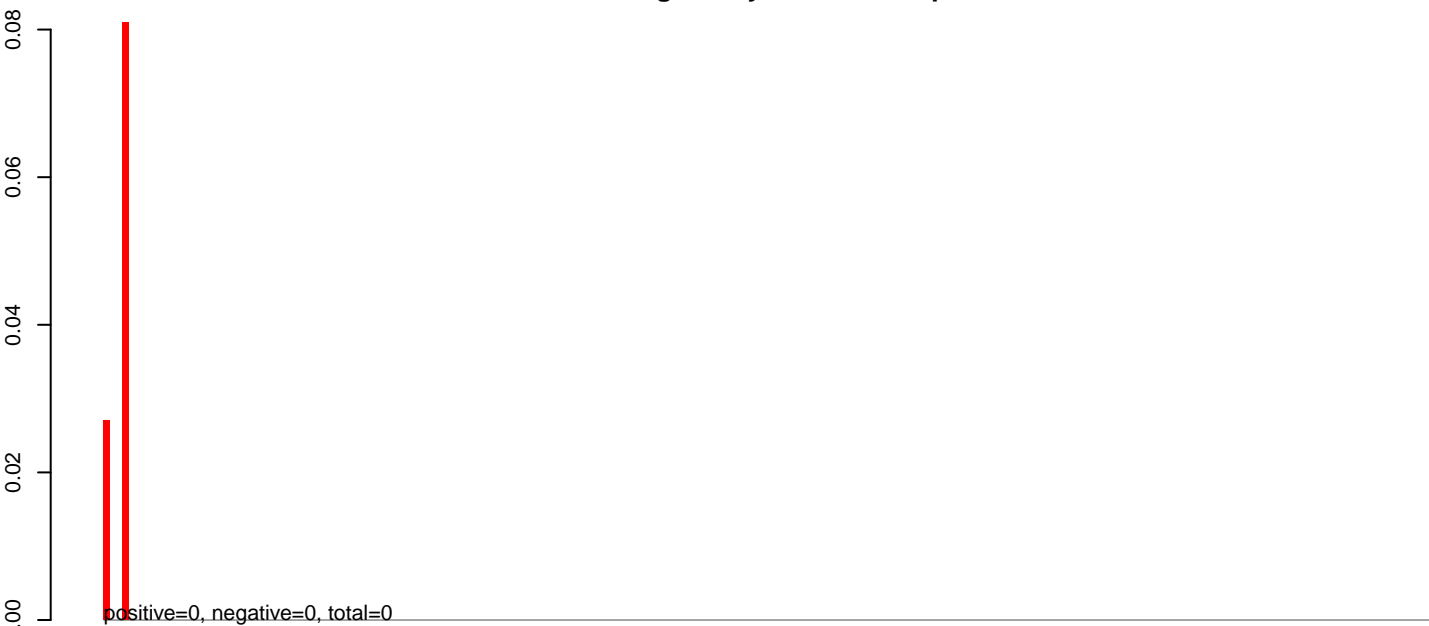
AeAeg_Ovary_GH.rep



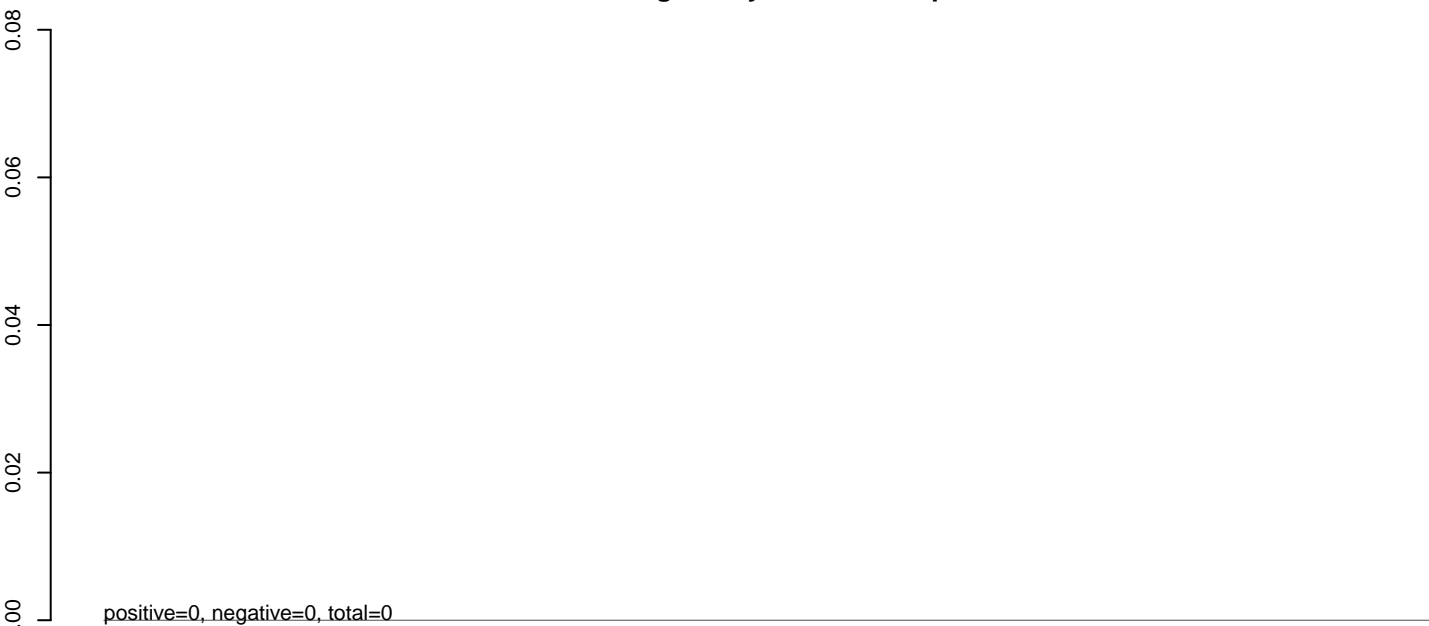
s_putative_small_transmembrane_protein_putative_glycoprotein_putative_small_ZnF_protein_and_RdRp_genes:1-11652

0 2000 4000 6000 8000 10000 12000

AeAeg_Ovary_GH.18_23.rep



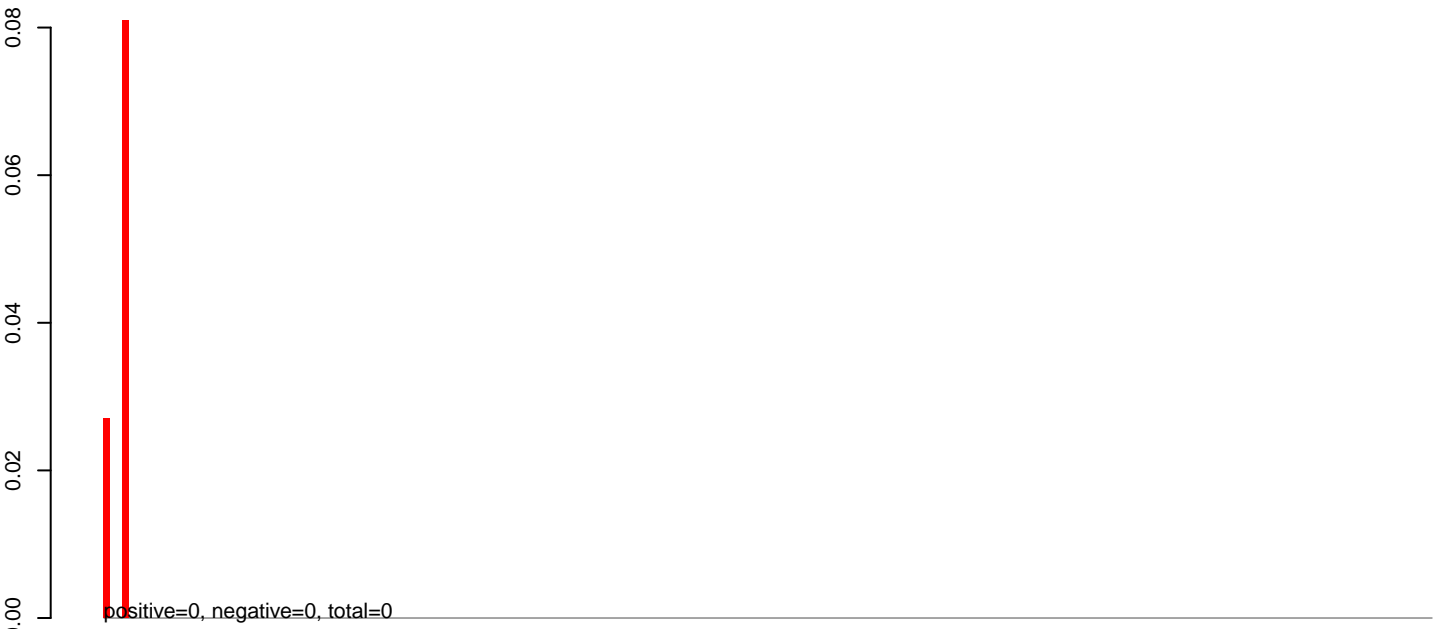
AeAeg_Ovary_GH.24_35.rep



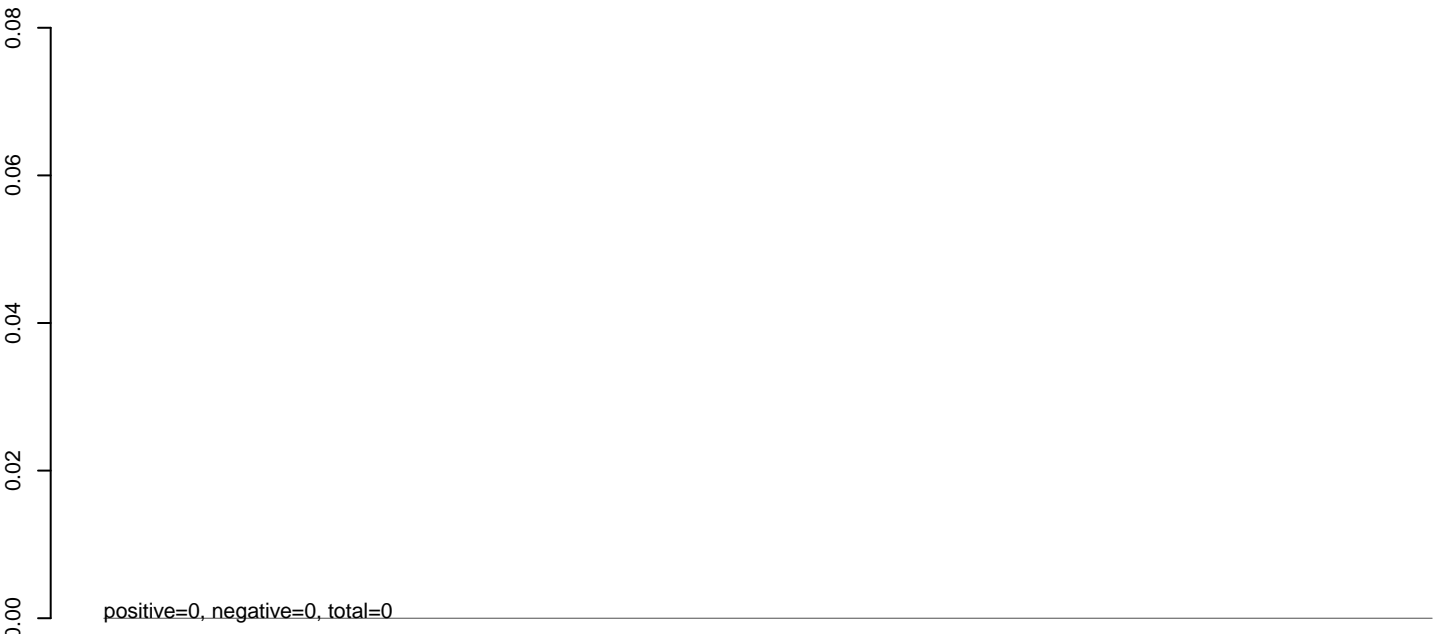
AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

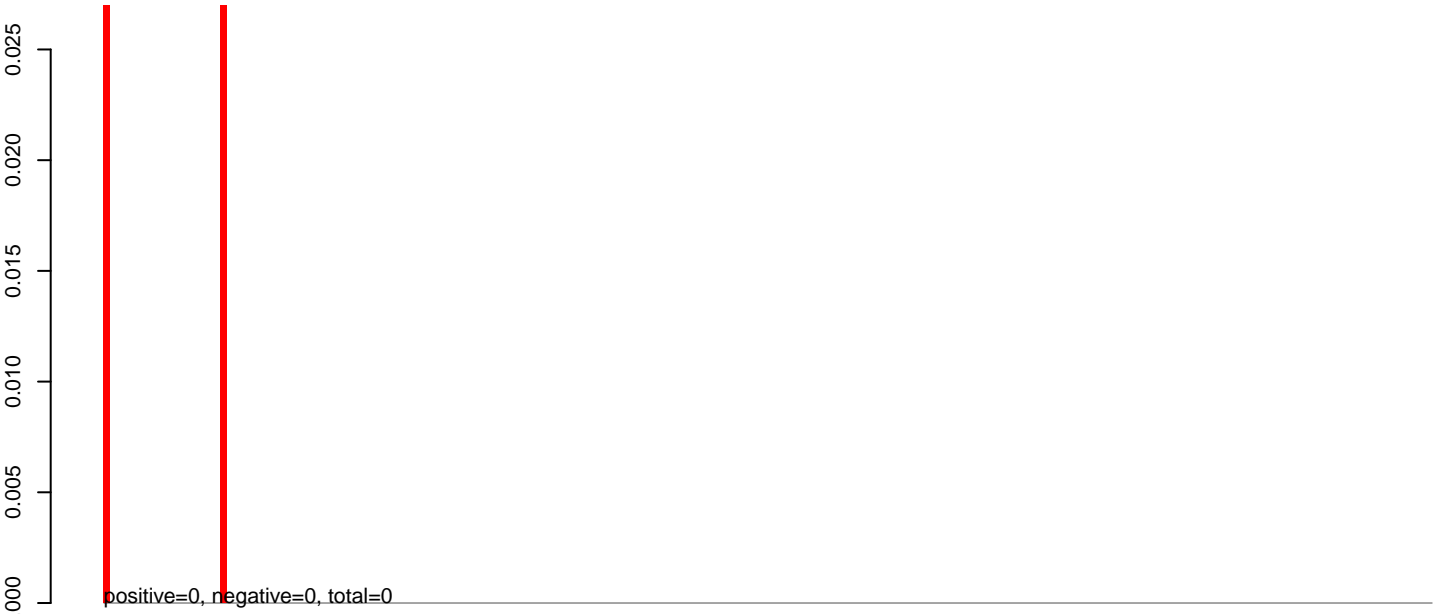


AeAeg_Ovary_GH.rep

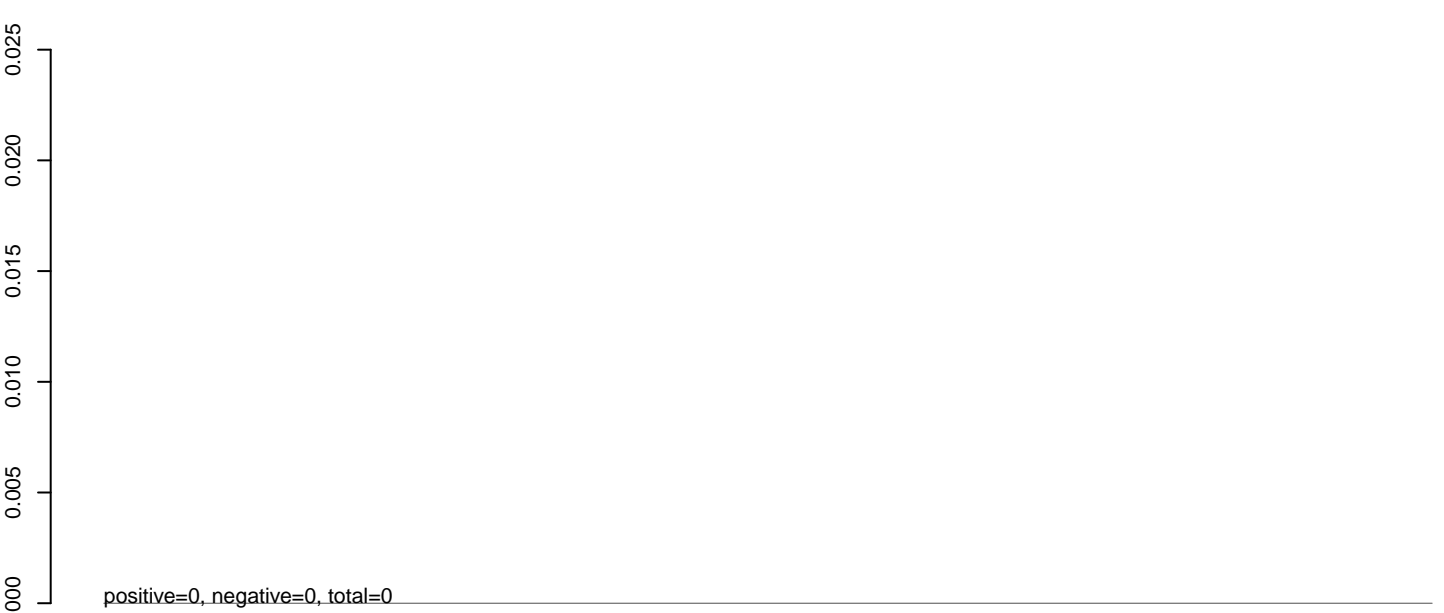


0 2000 4000 6000 8000 10000

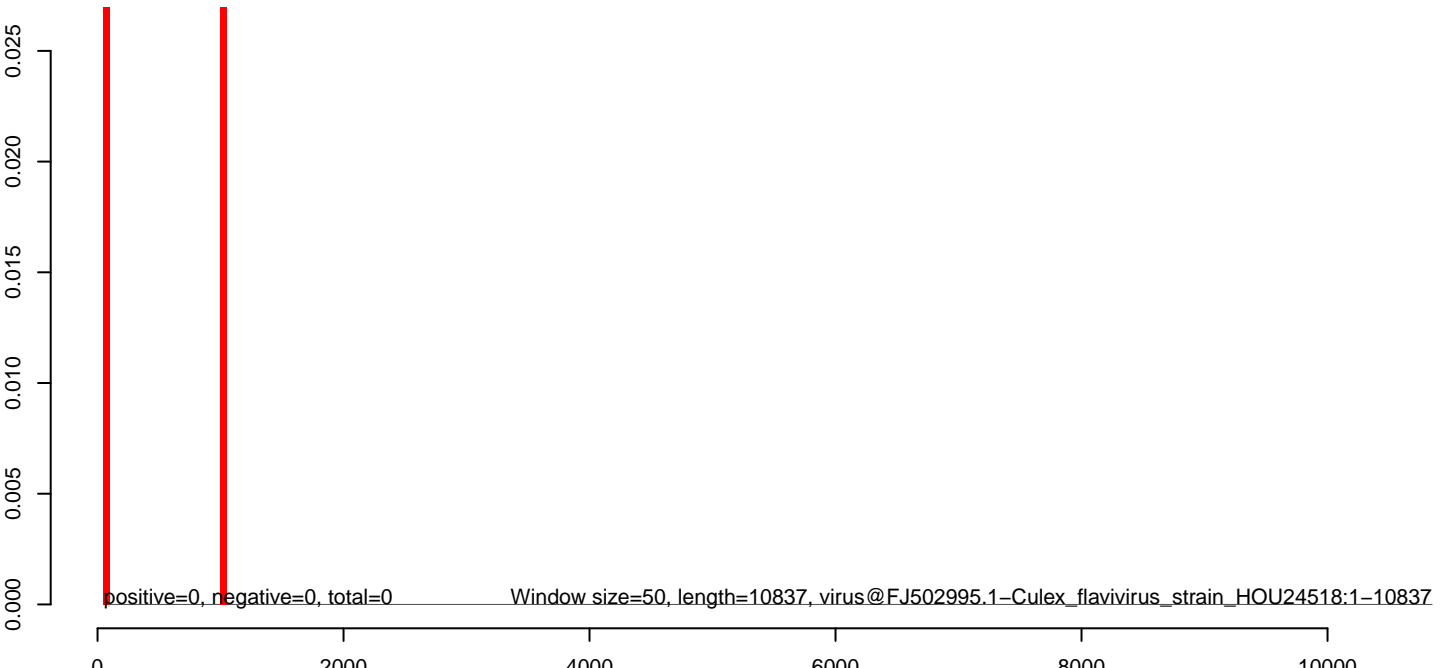
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



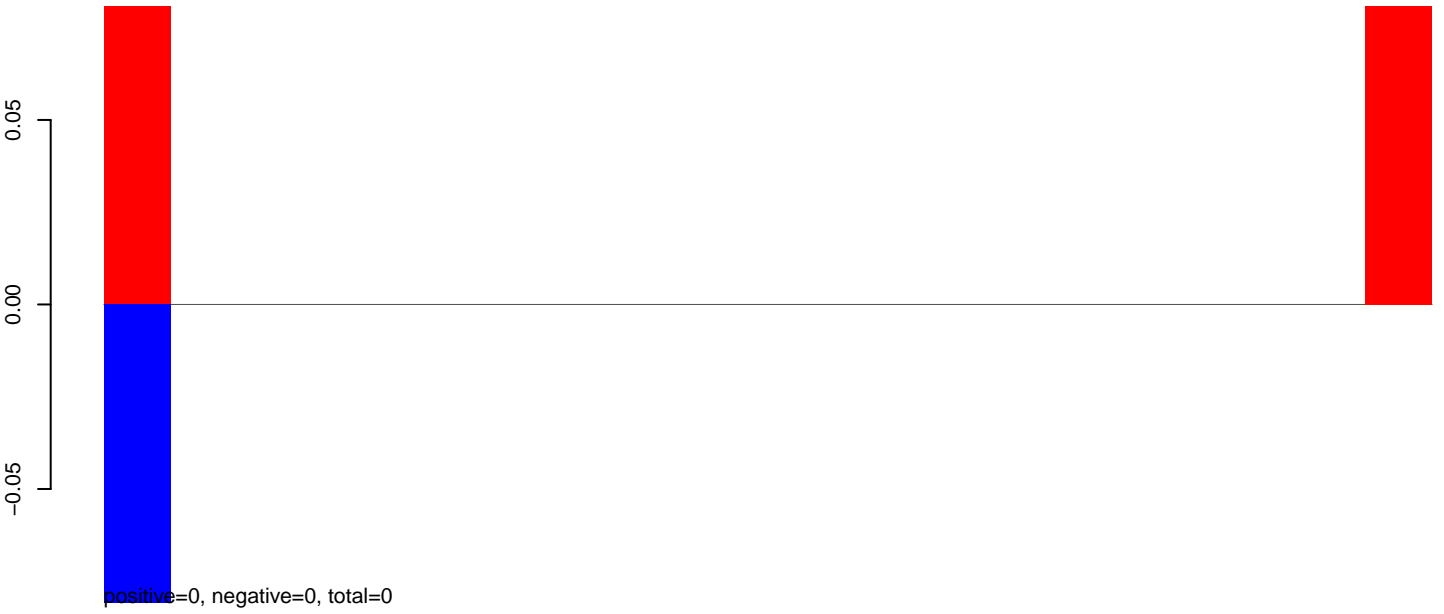
AeAeg_Ovary_GH.24_35.rep



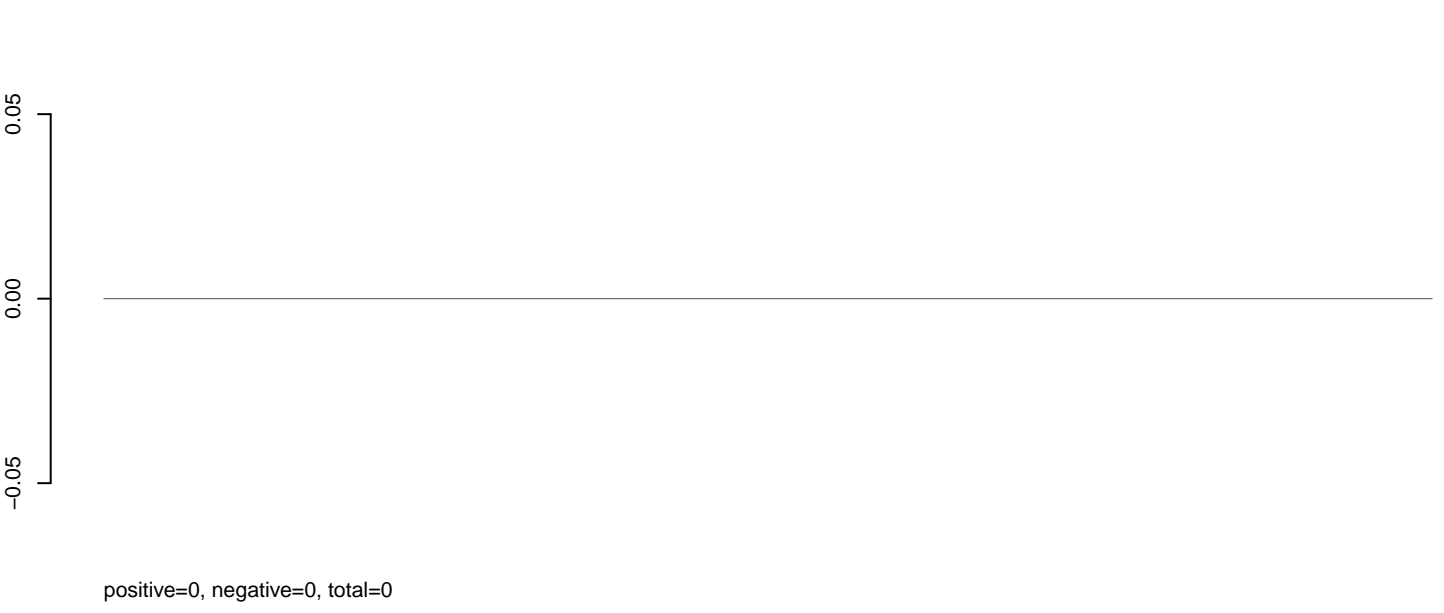
AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



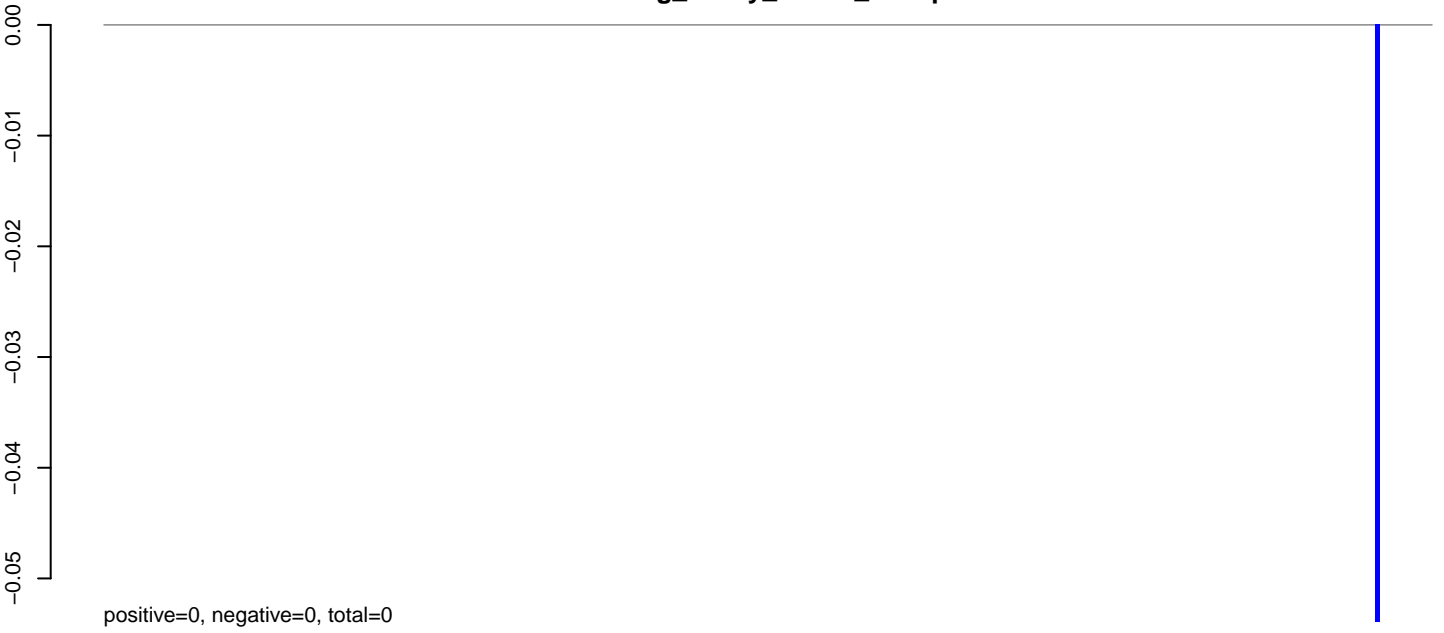
AeAeg_Ovary_GH.rep



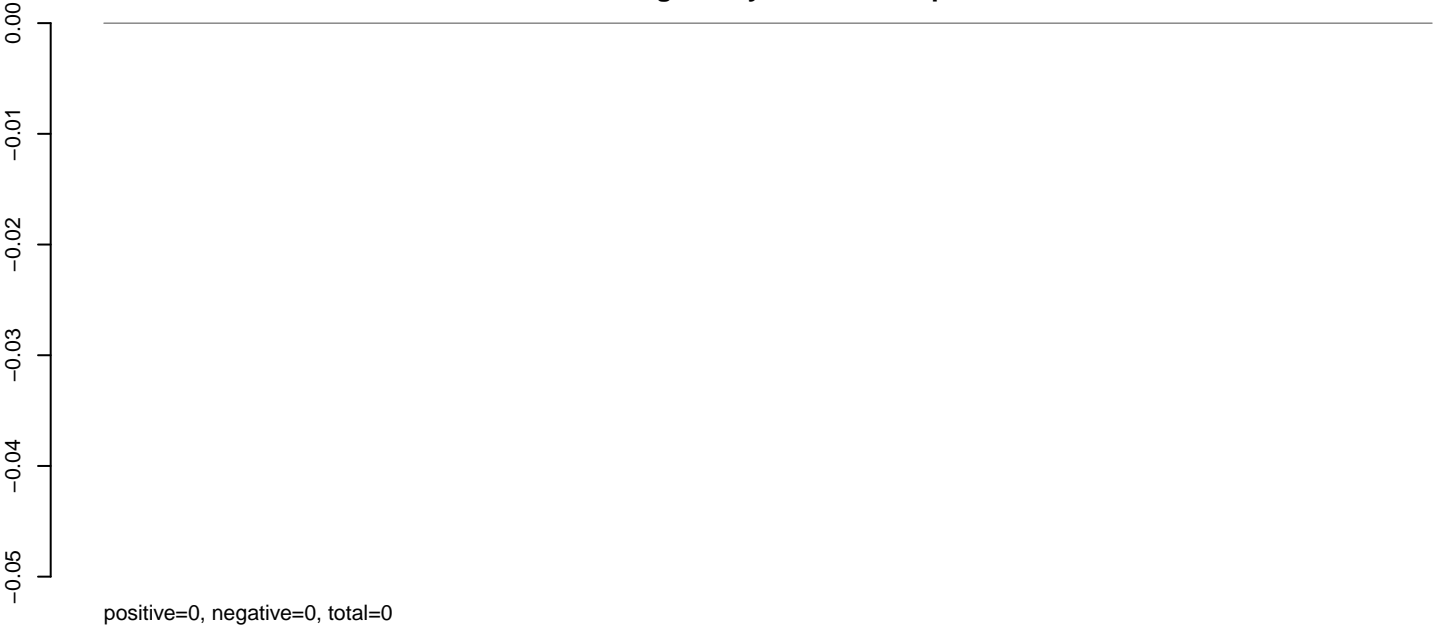
200 400 600 800 1000

positive=0, negative=0, total=0, length=1012, virus@AF411845.1-Elaeis_guineensis_MADS_box_transcription_factor_AGL2-3_mRNA:1-1012

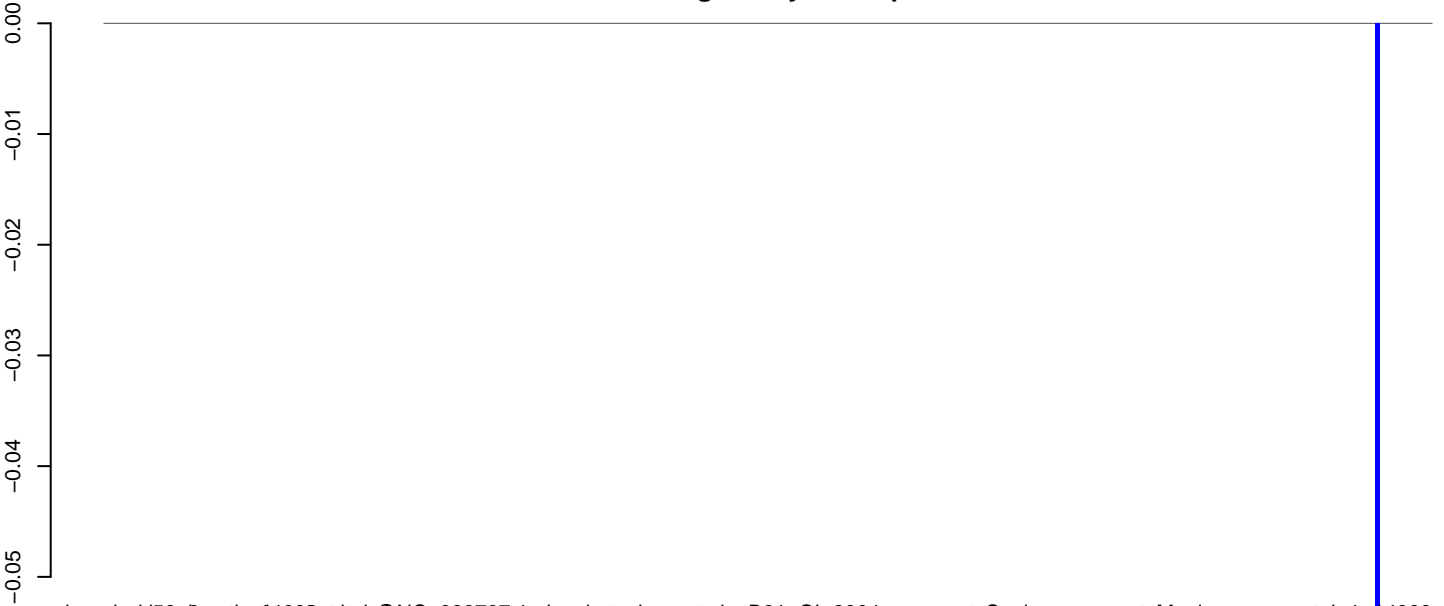
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

positive=0, negative=0, total=0

AeAeg_Ovary_GH.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

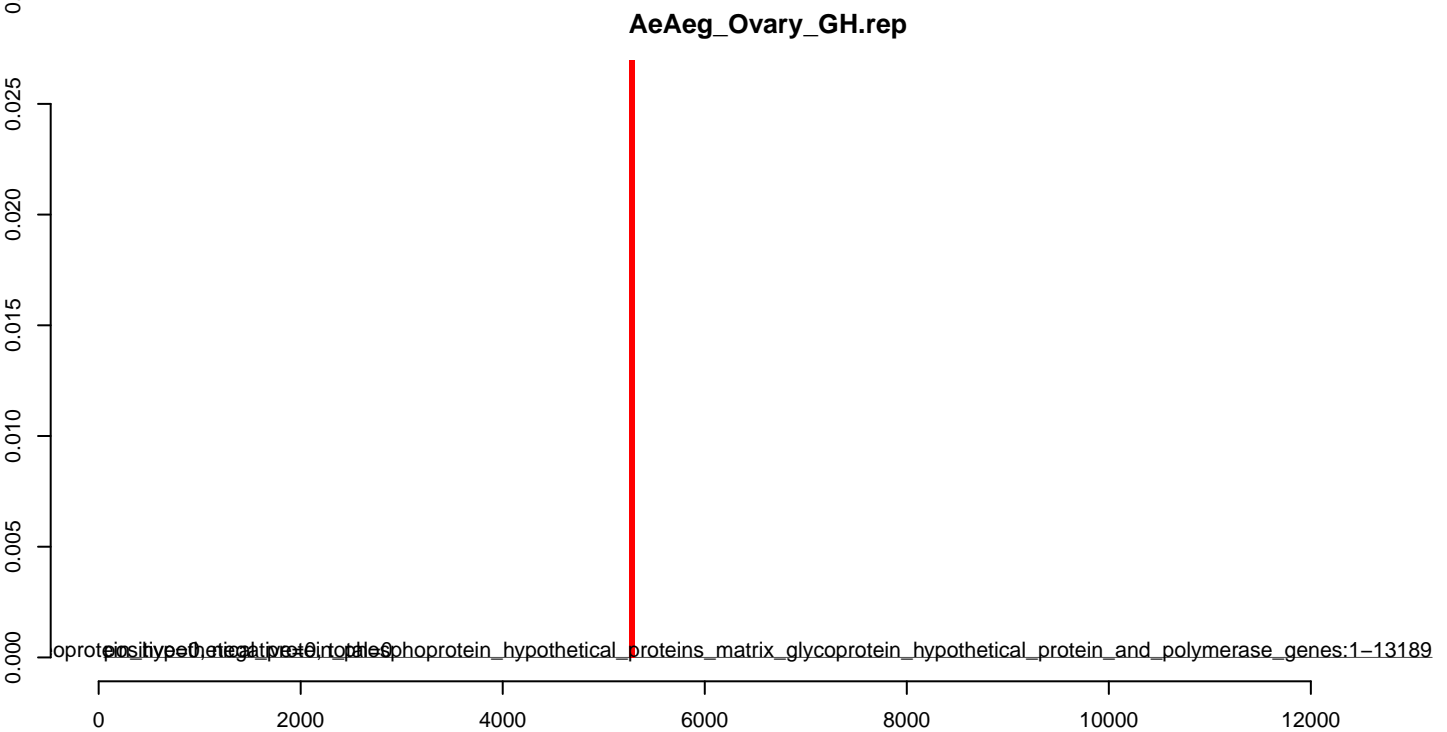
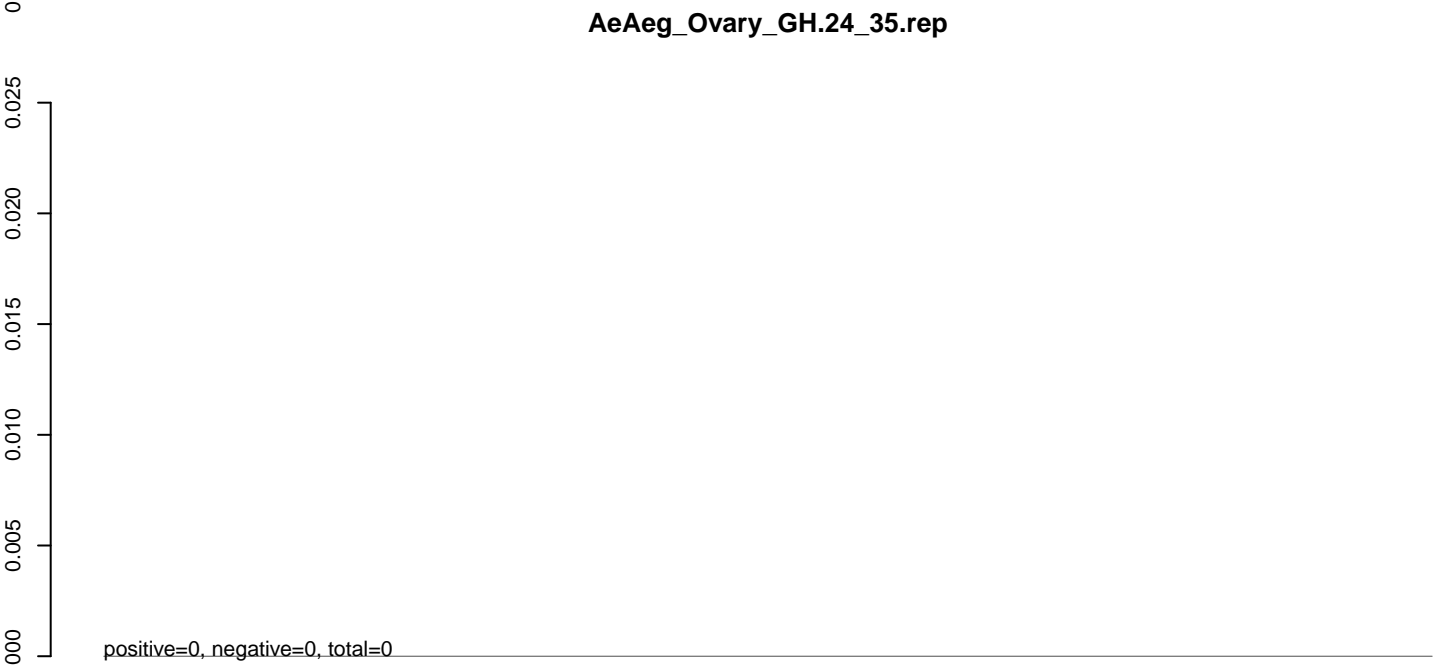
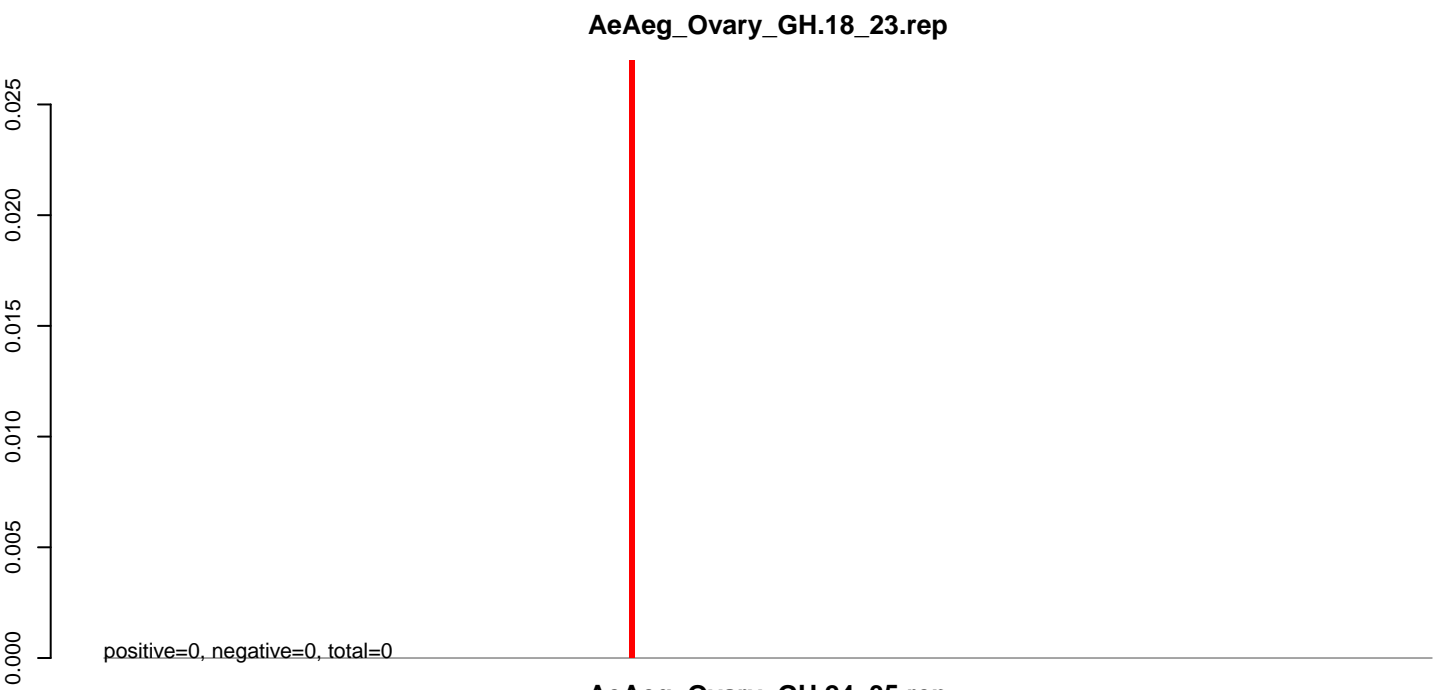
positive=0, negative=0, total=0

AeAeg_Ovary_GH.rep

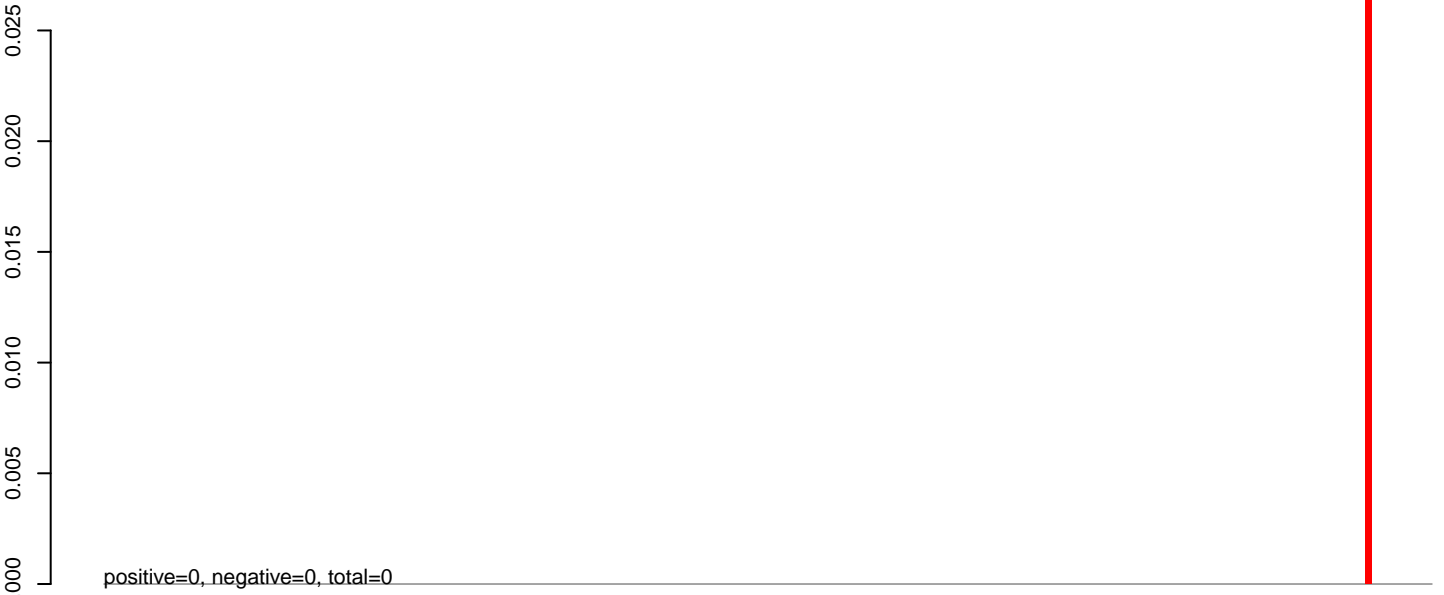
0.000
-0.005
-0.010
-0.015
-0.020
-0.025

2099, positive=0, negative=0, total=0 Charoen-like_virus_isolate_Rio_segment_S_plus_segment_M_plus_segment_L_Nucleocapsid_gene:1-12099

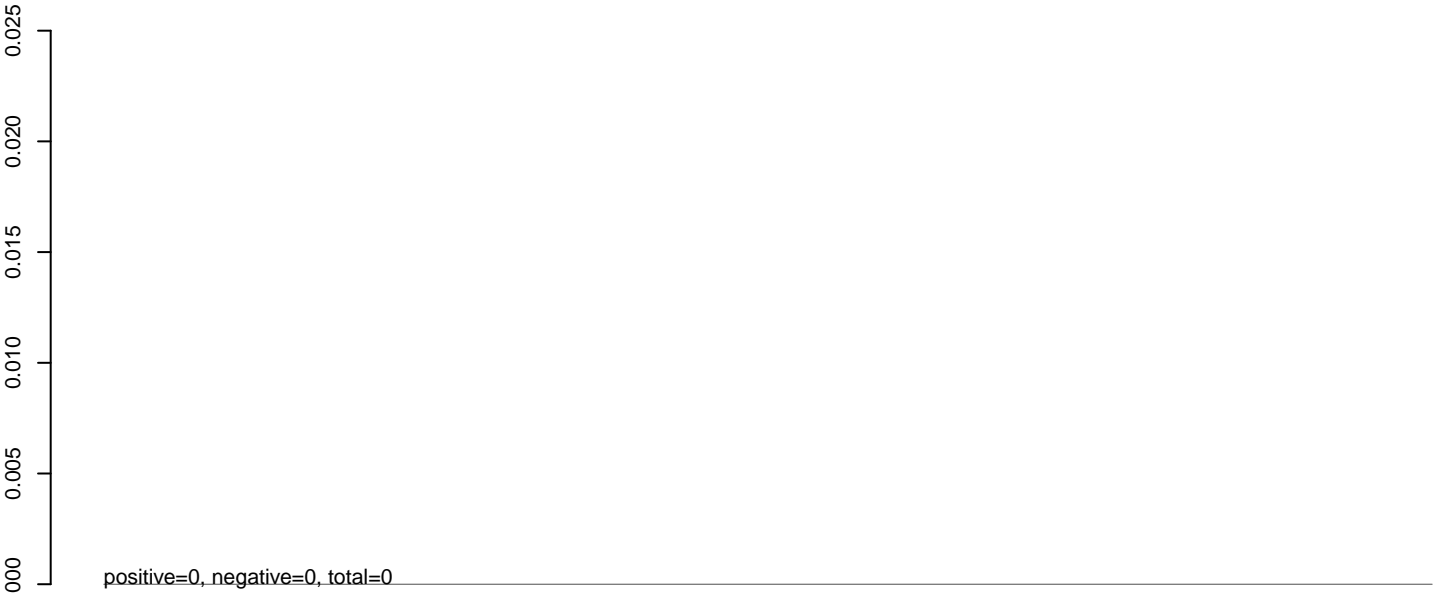
0 2000 4000 6000 8000 10000 12000



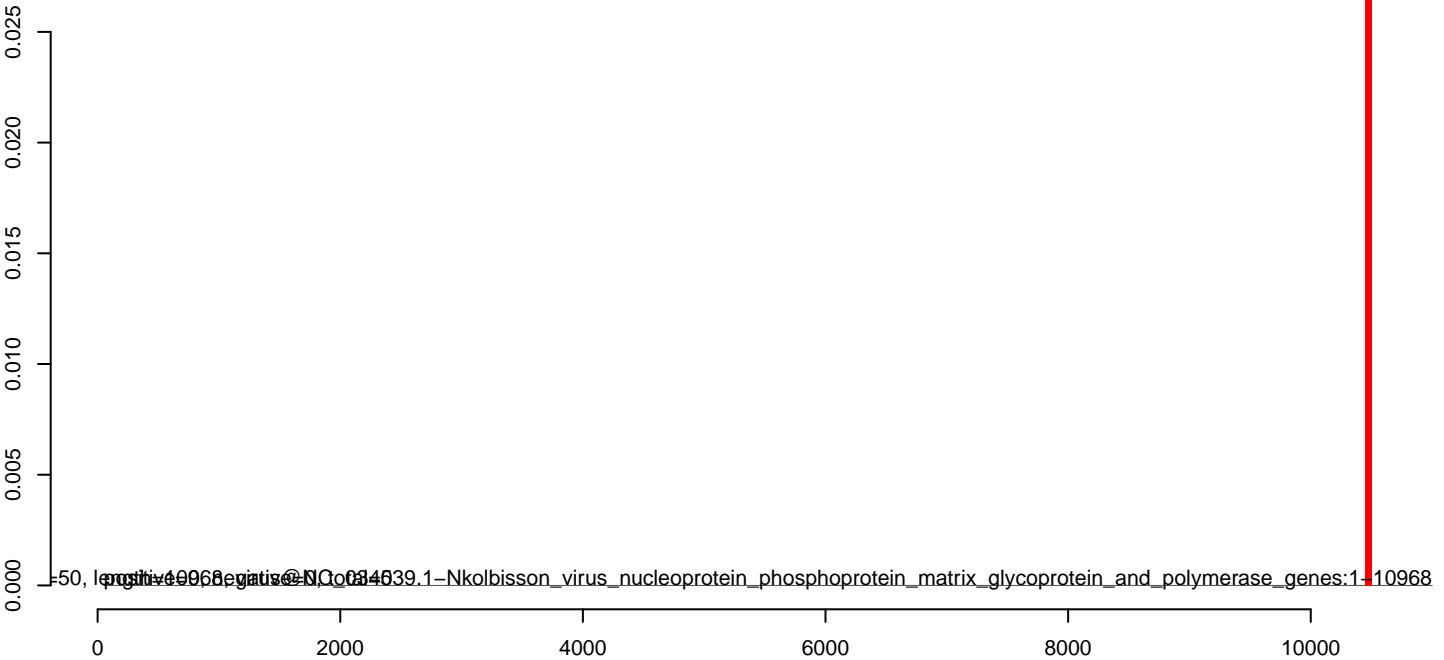
AeAeg_Ovary_GH.18_23.rep



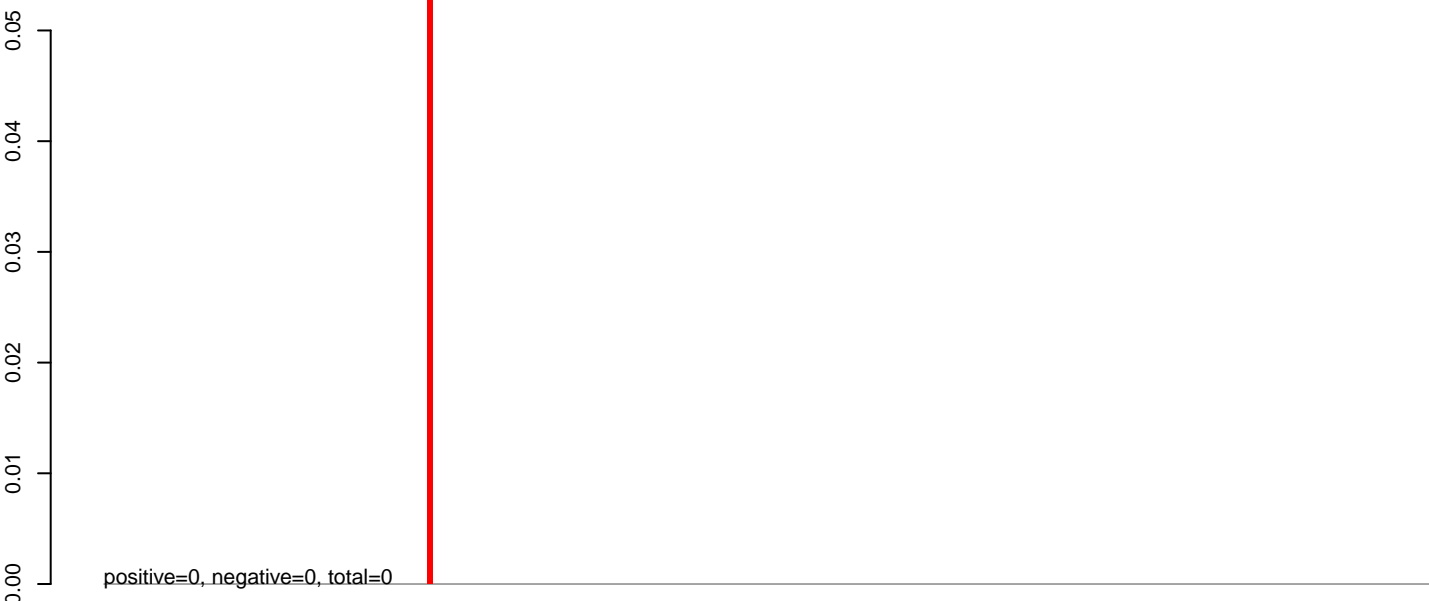
AeAeg_Ovary_GH.24_35.rep



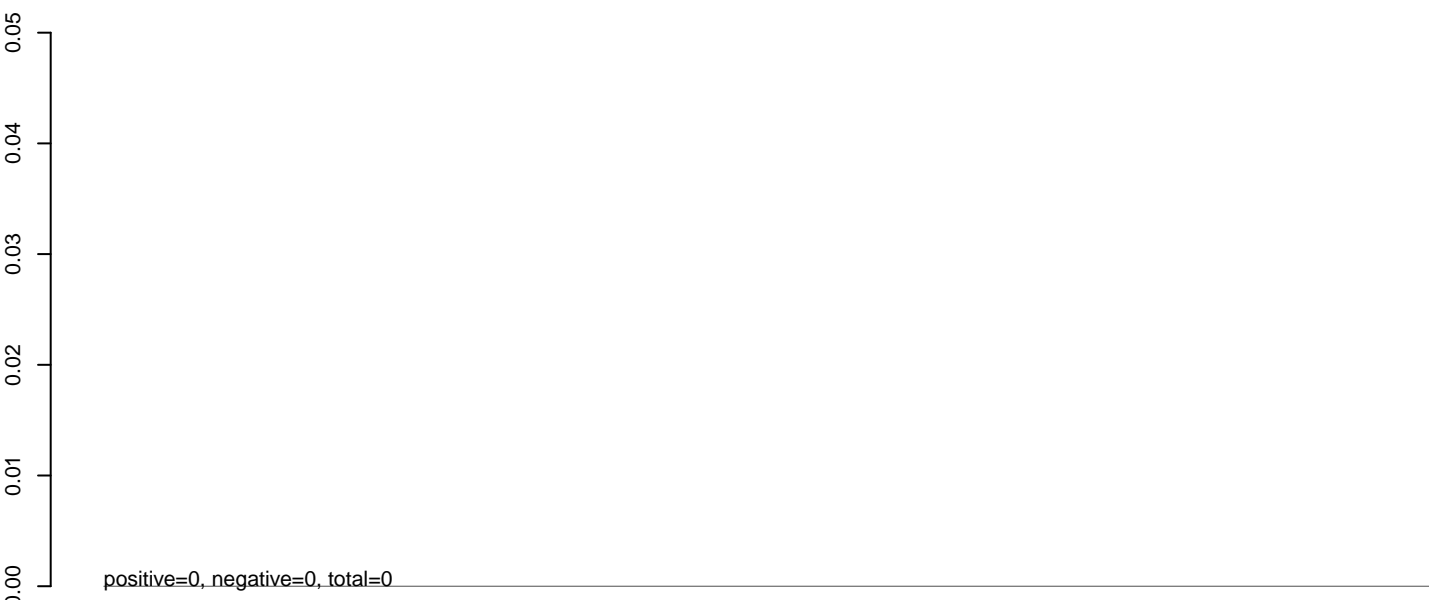
AeAeg_Ovary_GH.rep



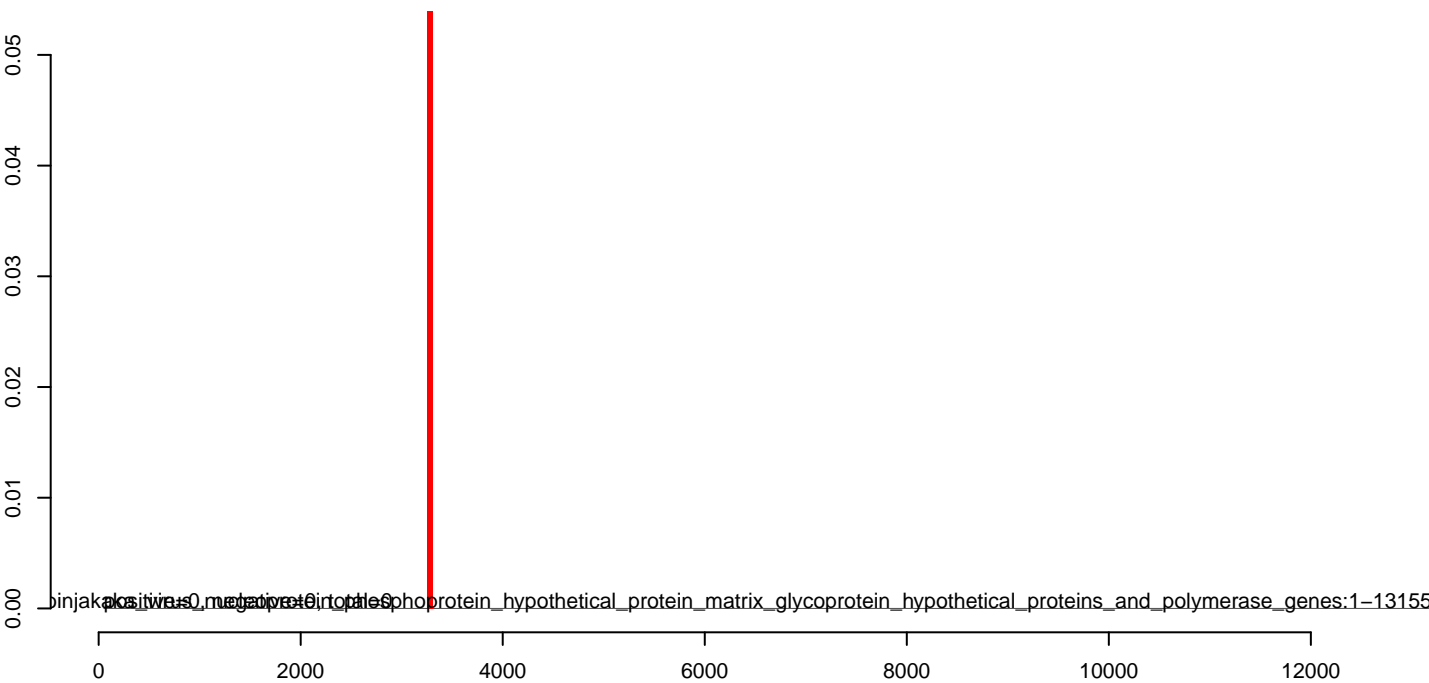
AeAeg_Ovary_GH.18_23.rep



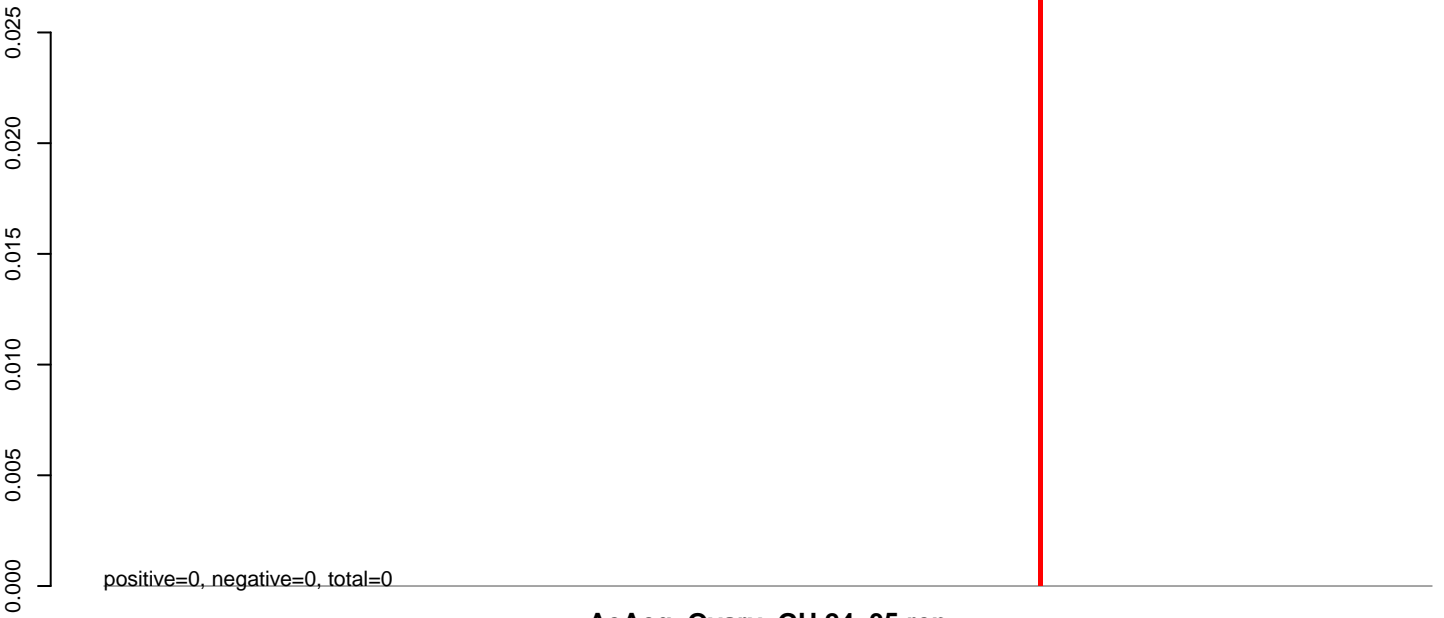
AeAeg_Ovary_GH.24_35.rep



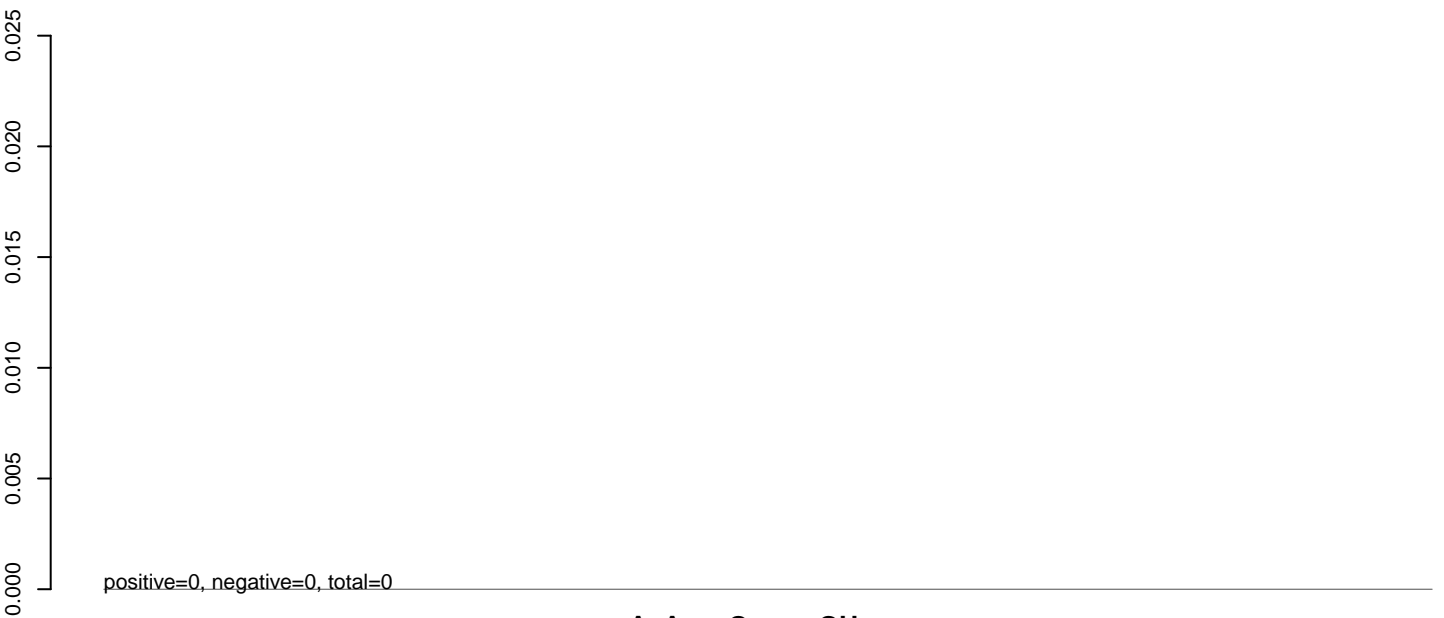
AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



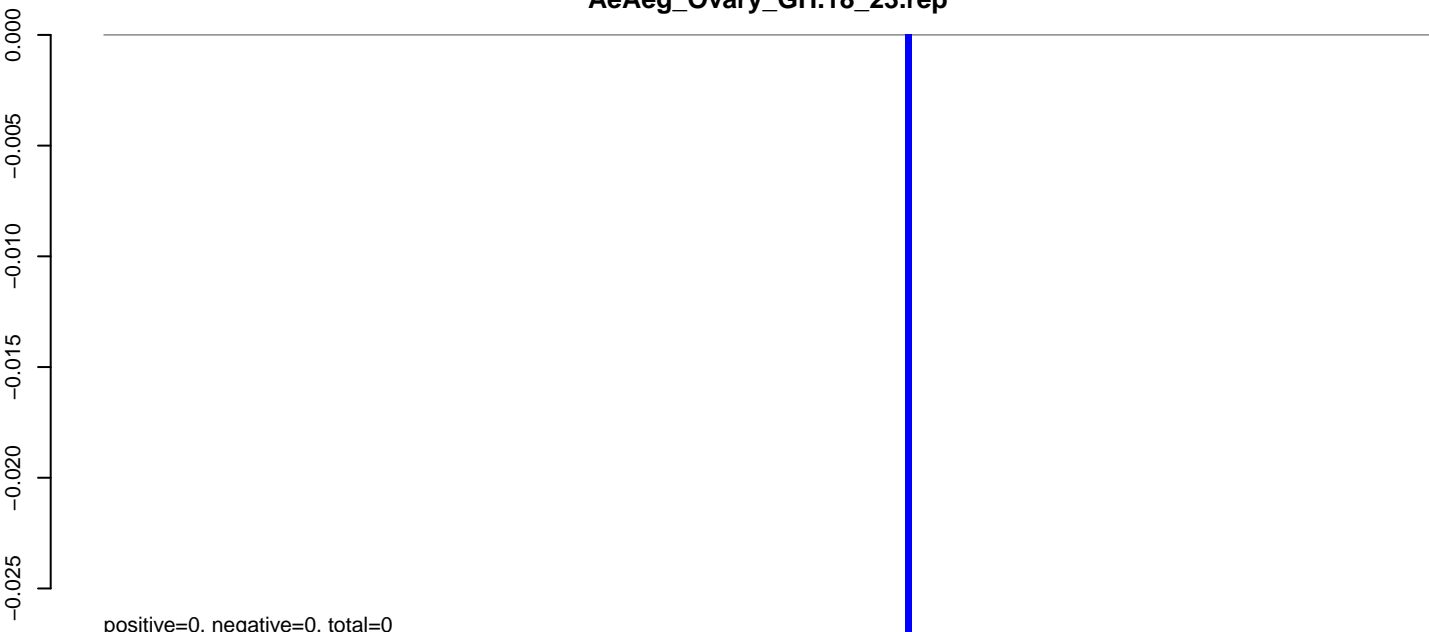
AeAeg_Ovary_GH.24_35.rep



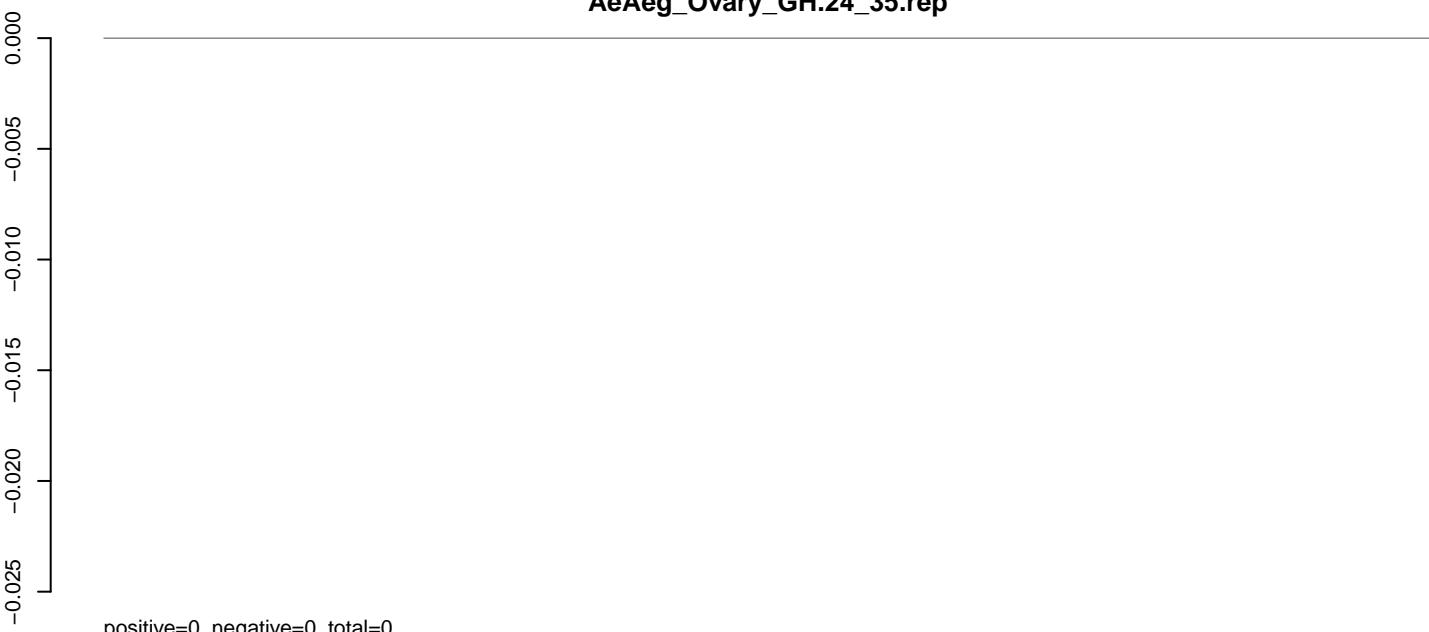
AeAeg_Ovary_GH.rep



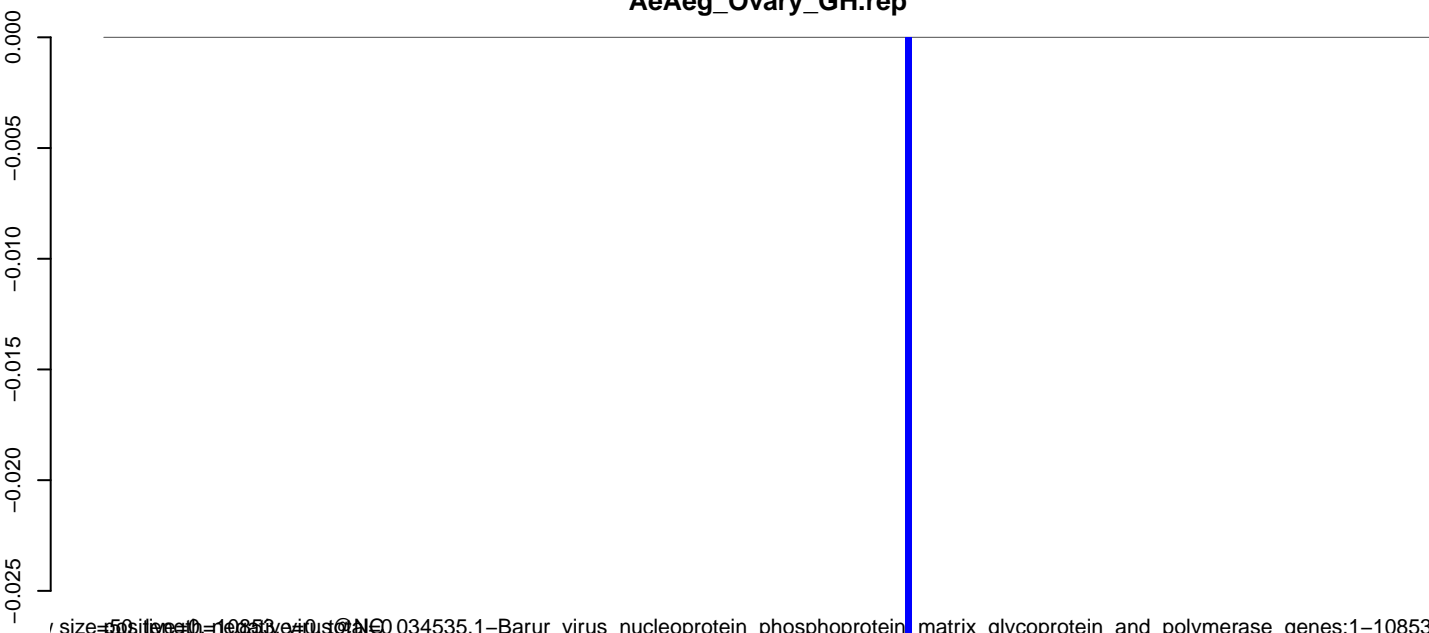
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



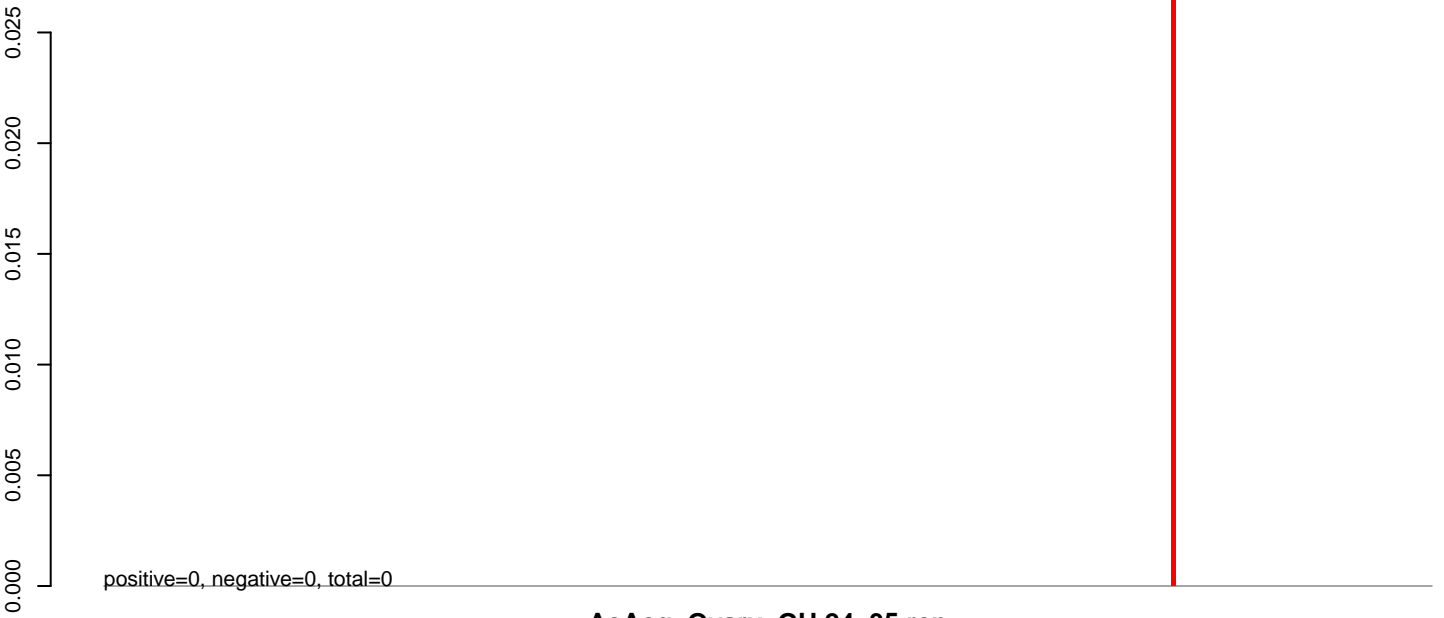
AeAeg_Ovary_GH.rep



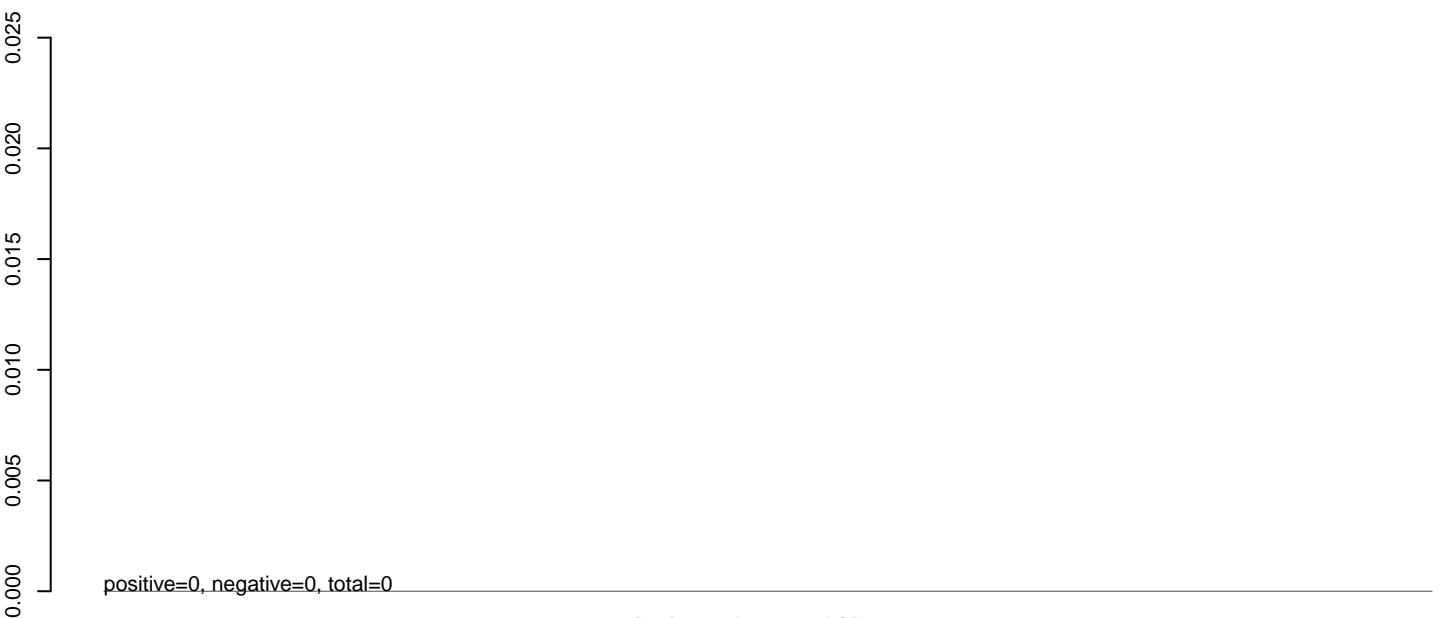
size=50, line=0, fill=0, stroke=0, @NE_034535.1-Barur_virus_nucleoprotein_phosphoprotein_matrix_glycoprotein_and_polymerase_genes:1-10853

0 2000 4000 6000 8000 10000

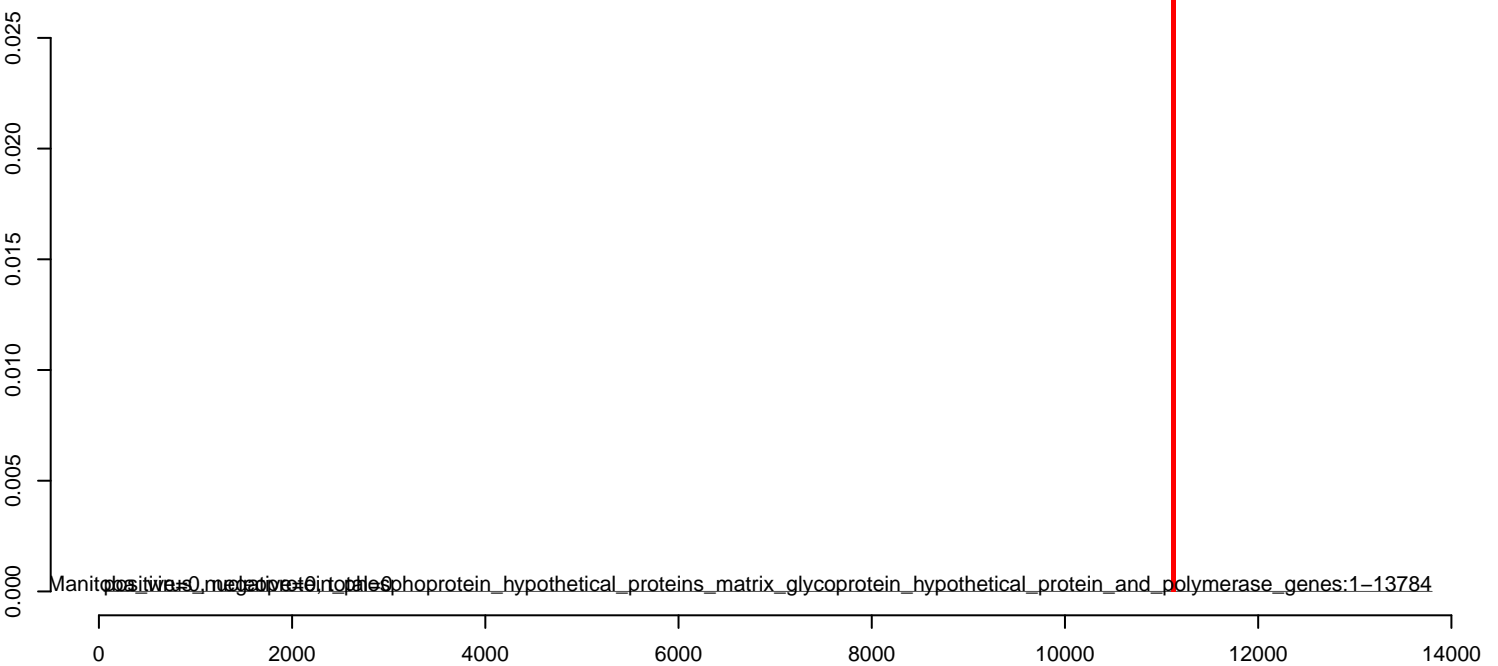
AeAeg_Ovary_GH.18_23.rep



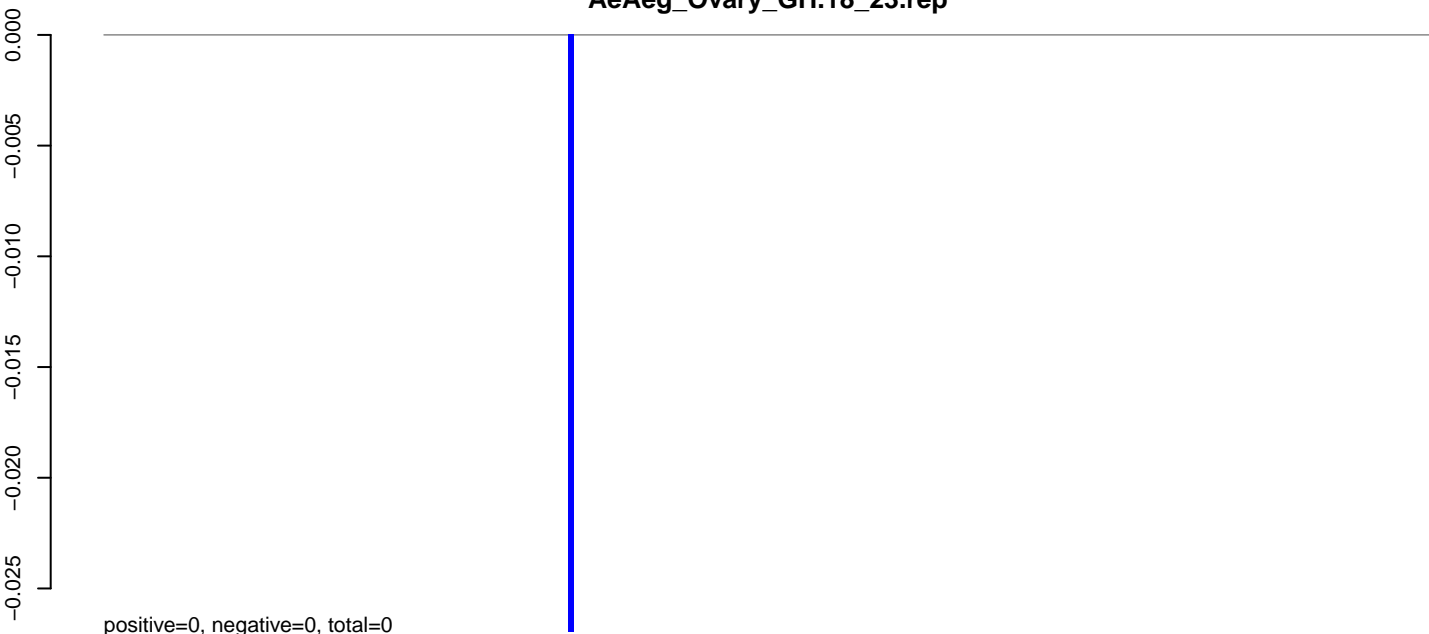
AeAeg_Ovary_GH.24_35.rep



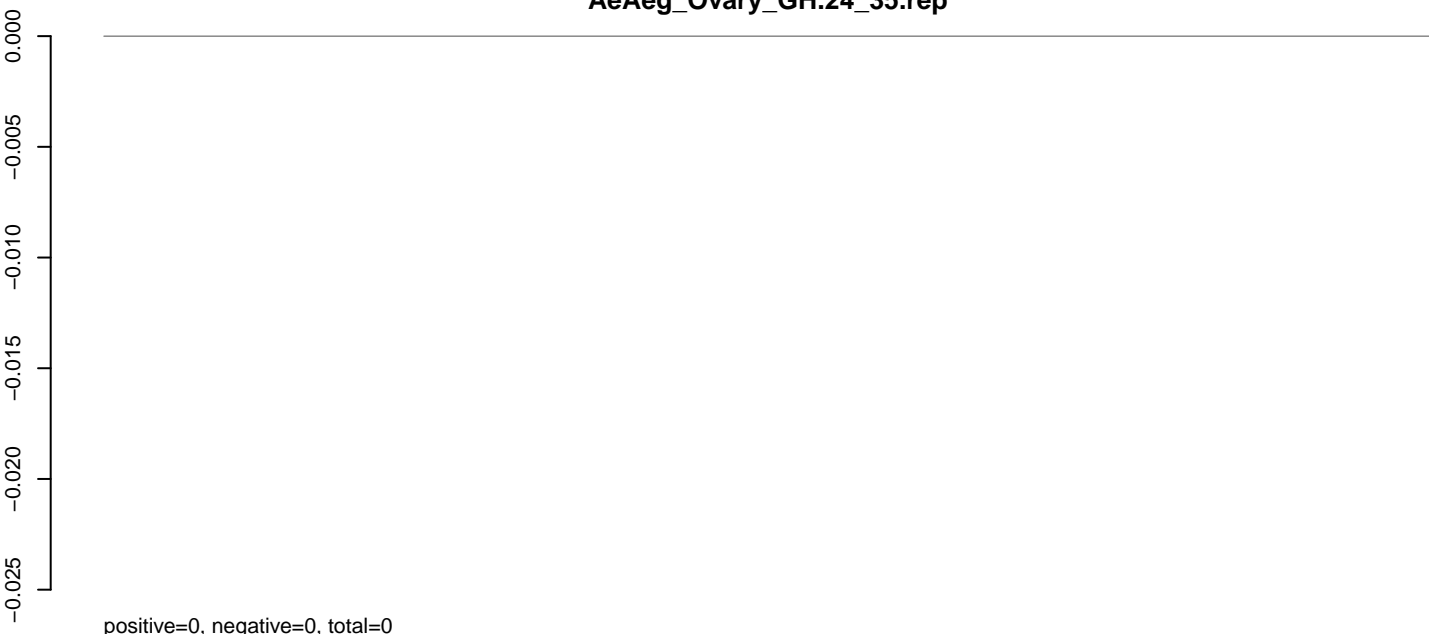
AeAeg_Ovary_GH.rep



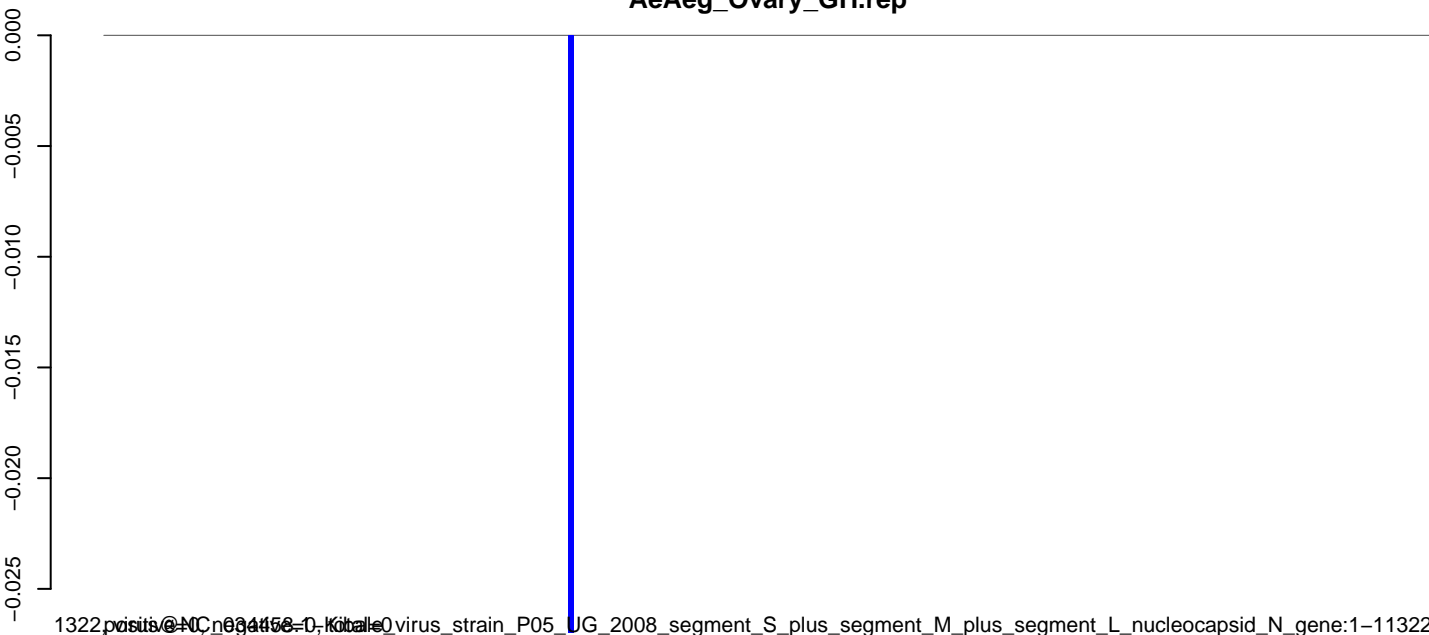
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



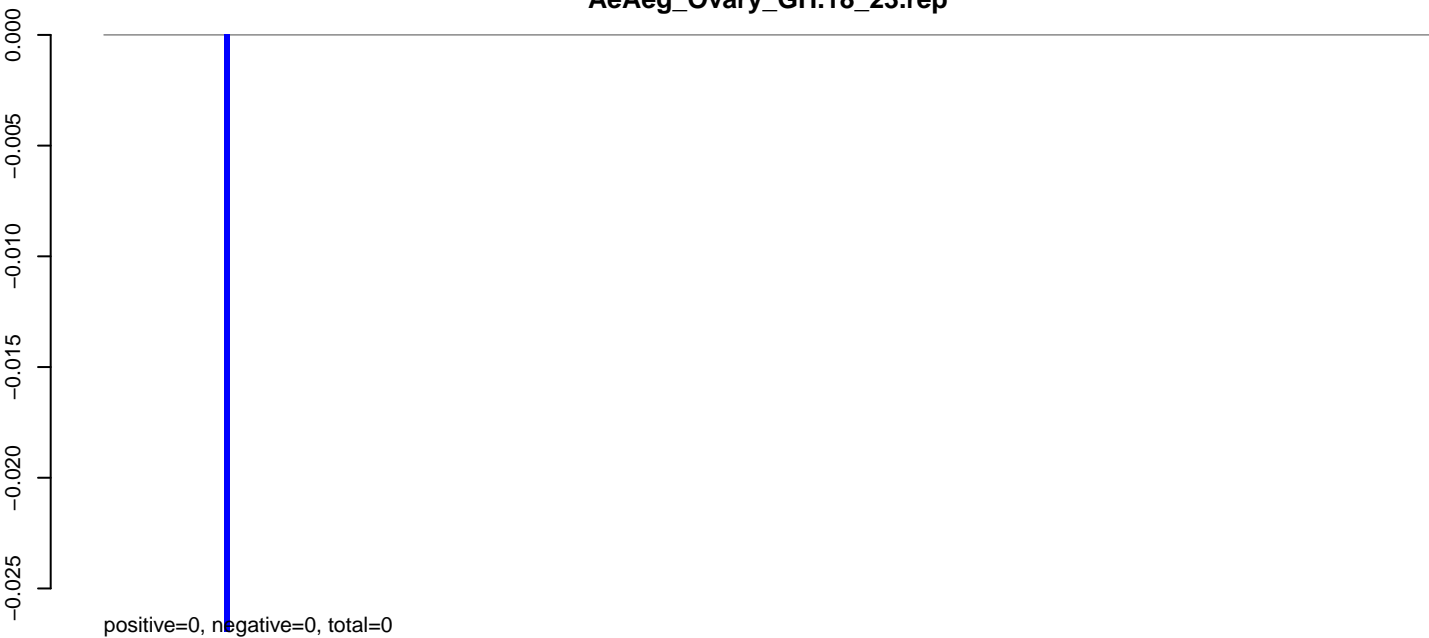
AeAeg_Ovary_GH.rep



1322_positiv@NC_034458-1_KOala0virus_strain_P05_UG_2008_segment_S_plus_segment_M_plus_segment_L_nucleocapsid_N_gene:1-11322

0 2000 4000 6000 8000 10000

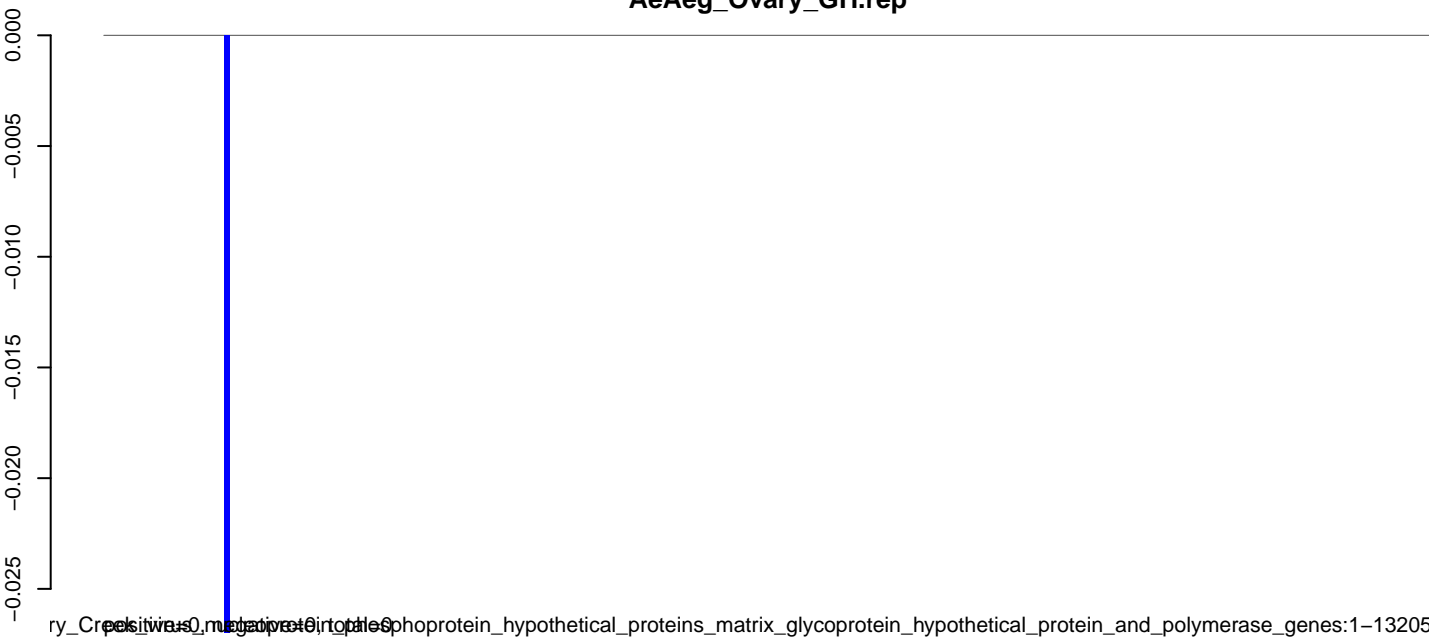
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



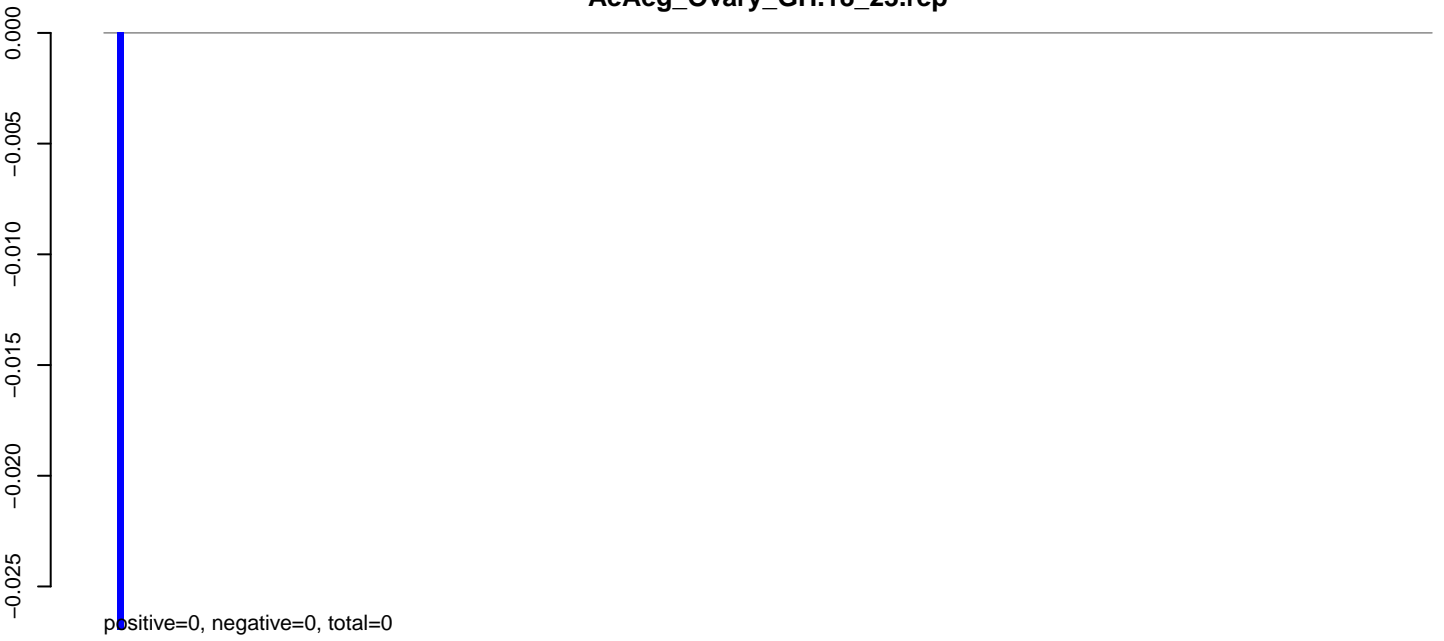
AeAeg_Ovary_GH.rep



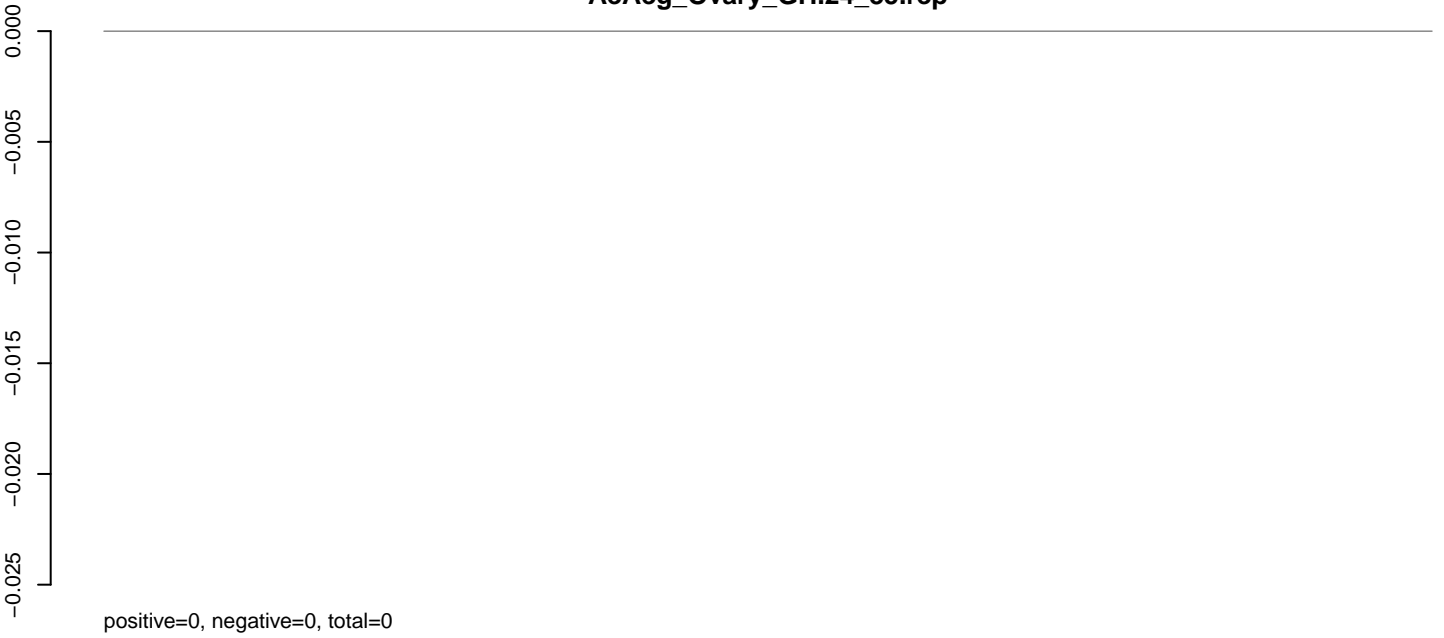
ry_Creptinus_megaprotin_phosphoprotein_hypothetical_proteins_matrix_glycoprotein_hypothetical_protein_and_polymerase_genes:1-13205

0 2000 4000 6000 8000 10000 12000

AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



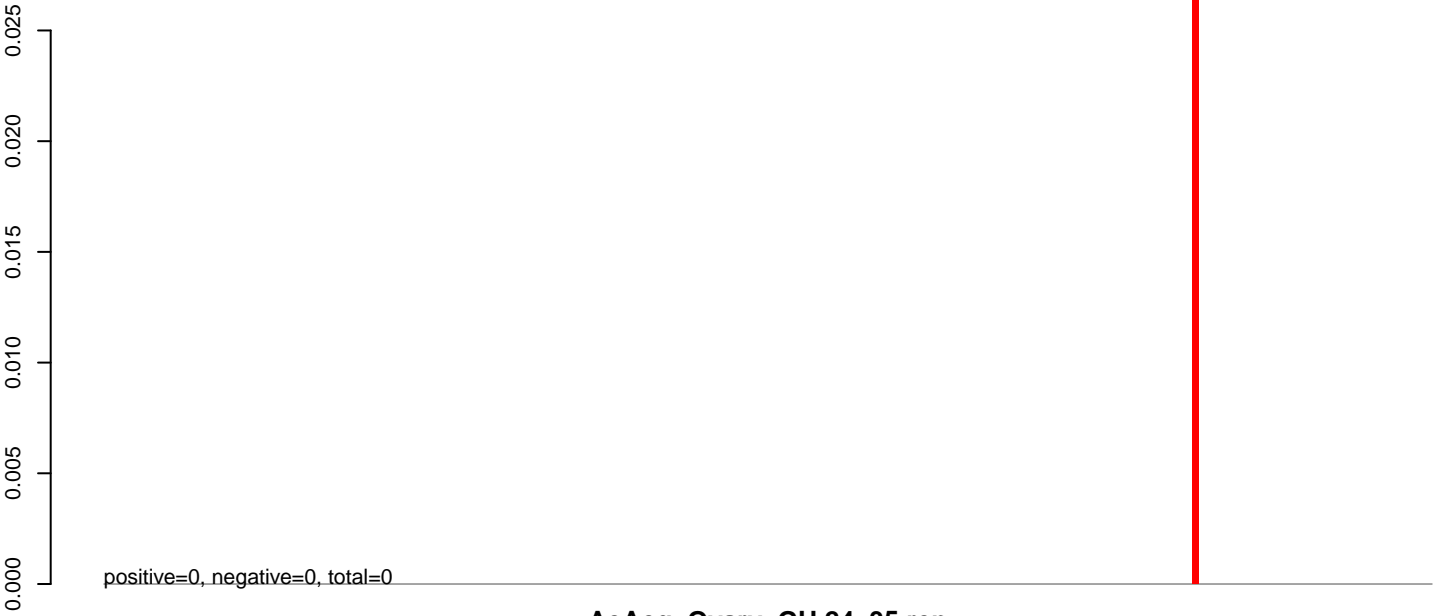
AeAeg_Ovary_GH.rep



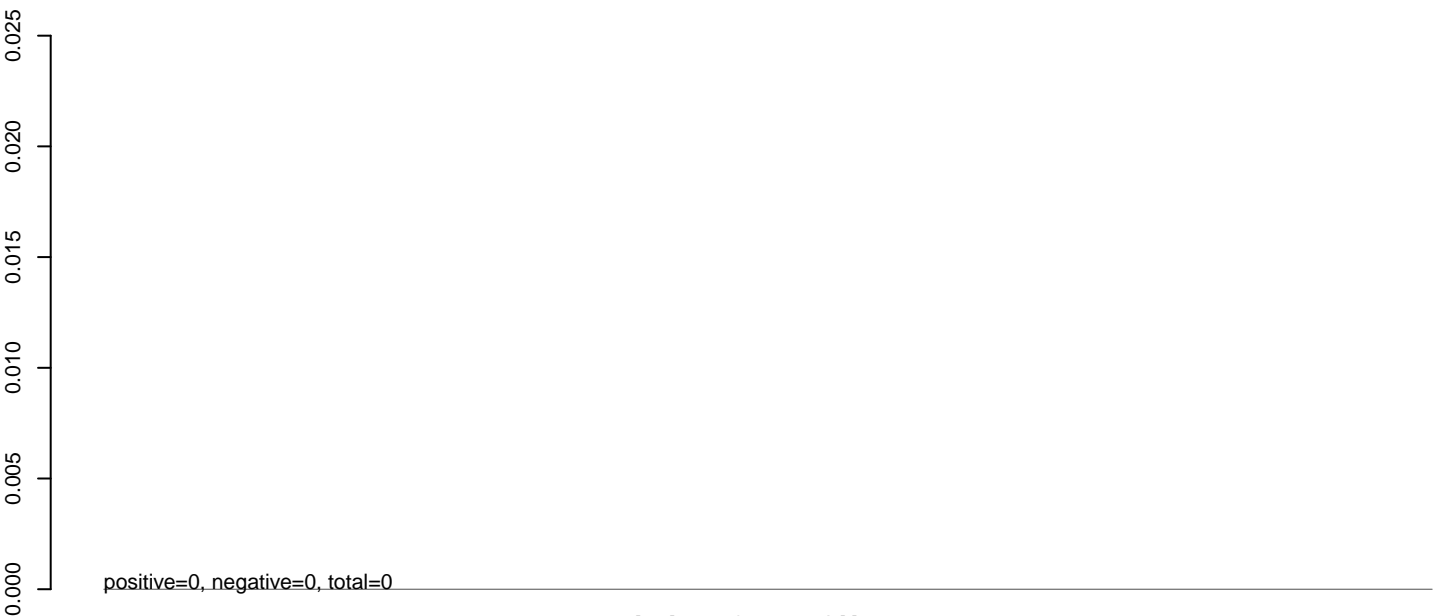
Window size=50, length=10293, virus@NC_034018.1-Yaounde_virus_strain_Dak_Ar_Y276_polyprotein_gene:1-10293

0 2000 4000 6000 8000 10000

AeAeg_Ovary_GH.18_23.rep



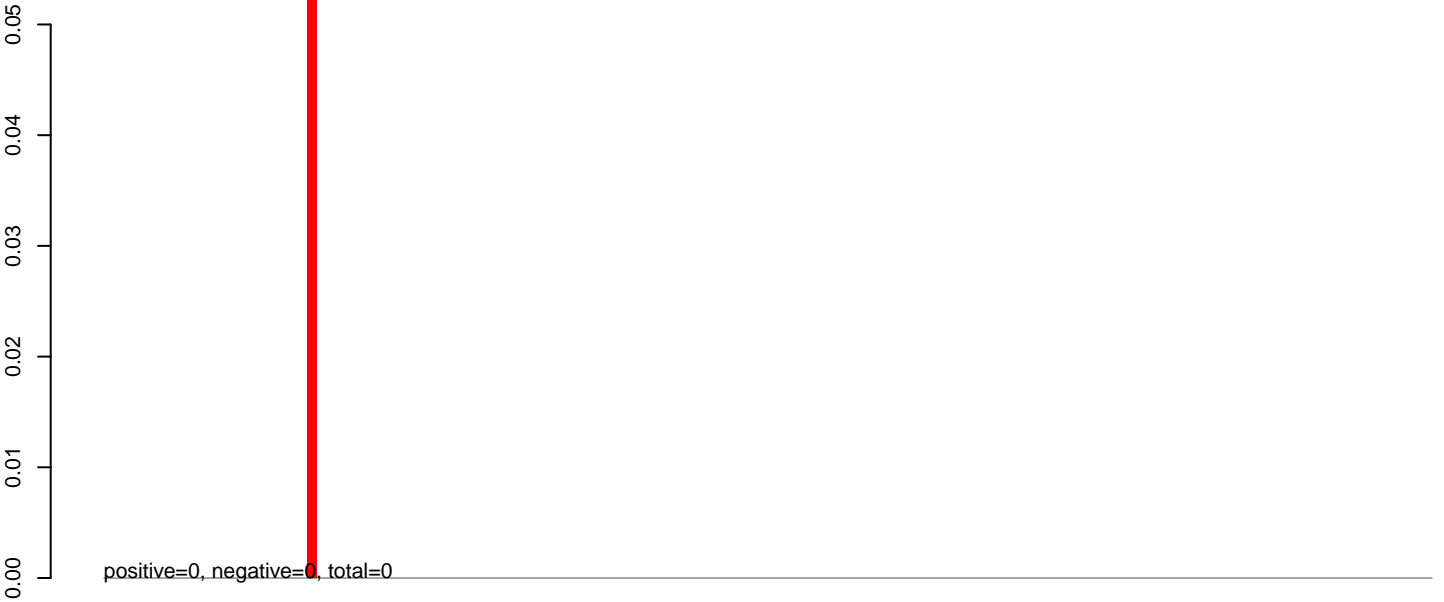
AeAeg_Ovary_GH.24_35.rep



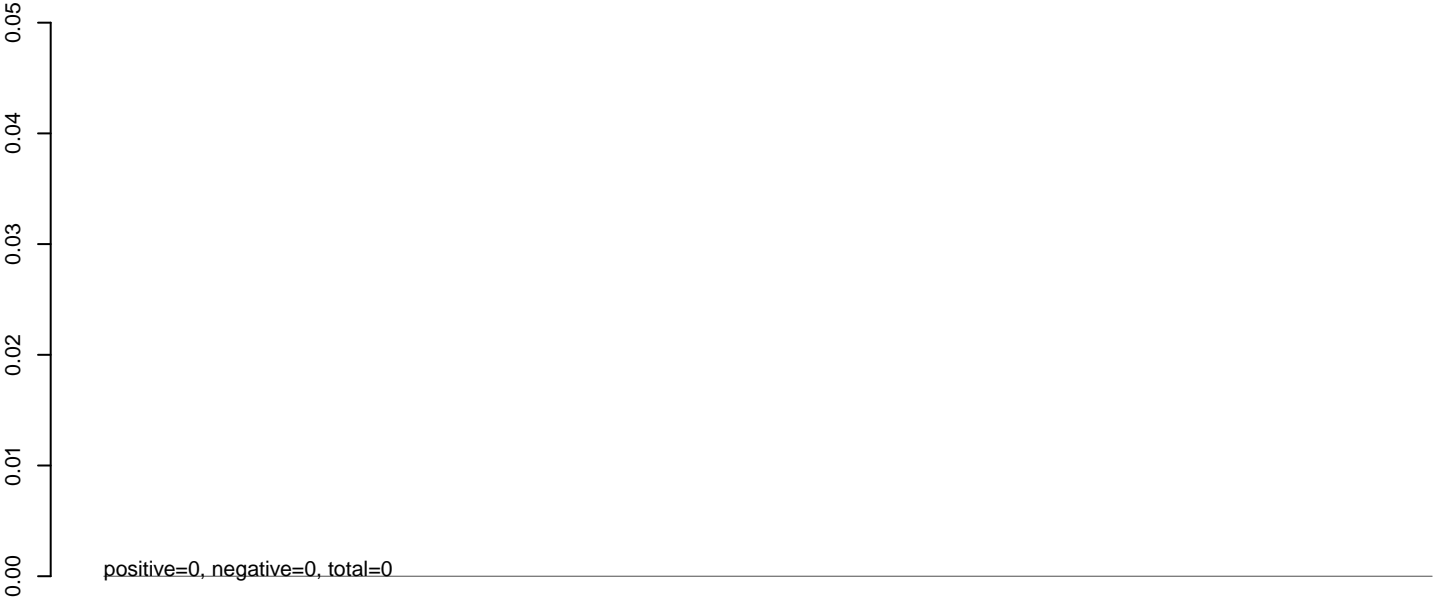
AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



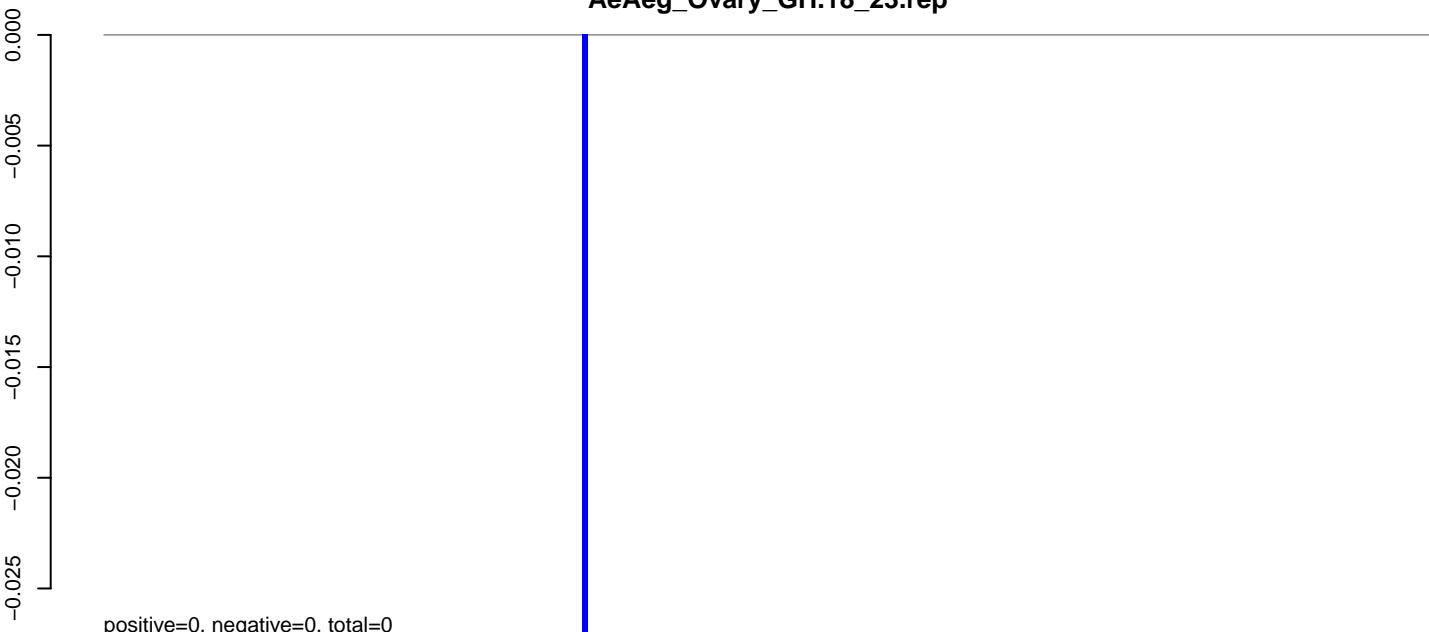
AeAeg_Ovary_GH.24_35.rep



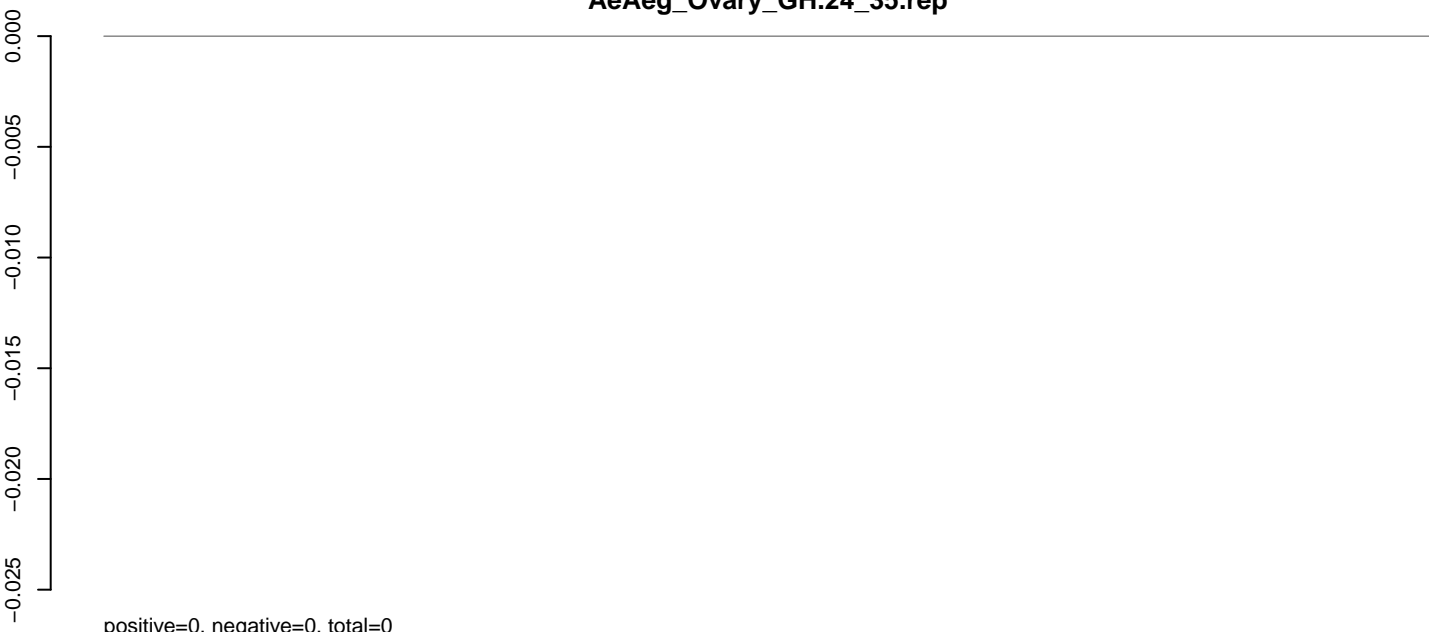
AeAeg_Ovary_GH.rep



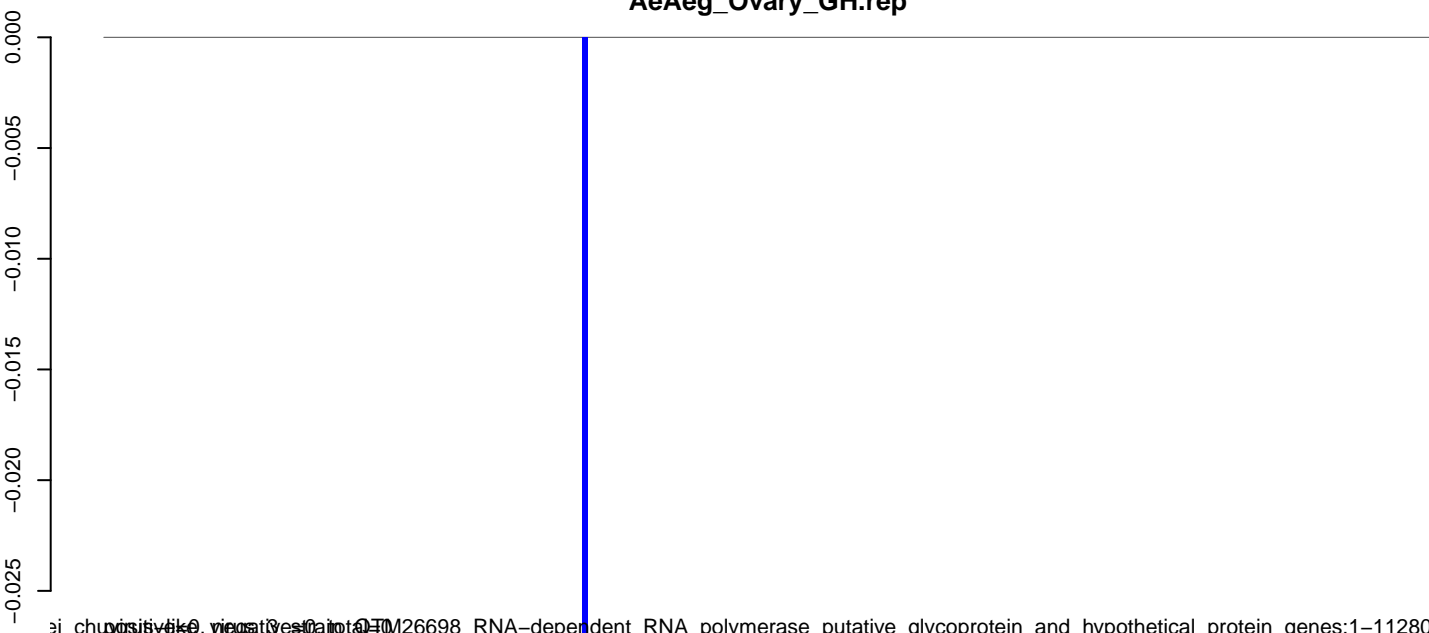
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



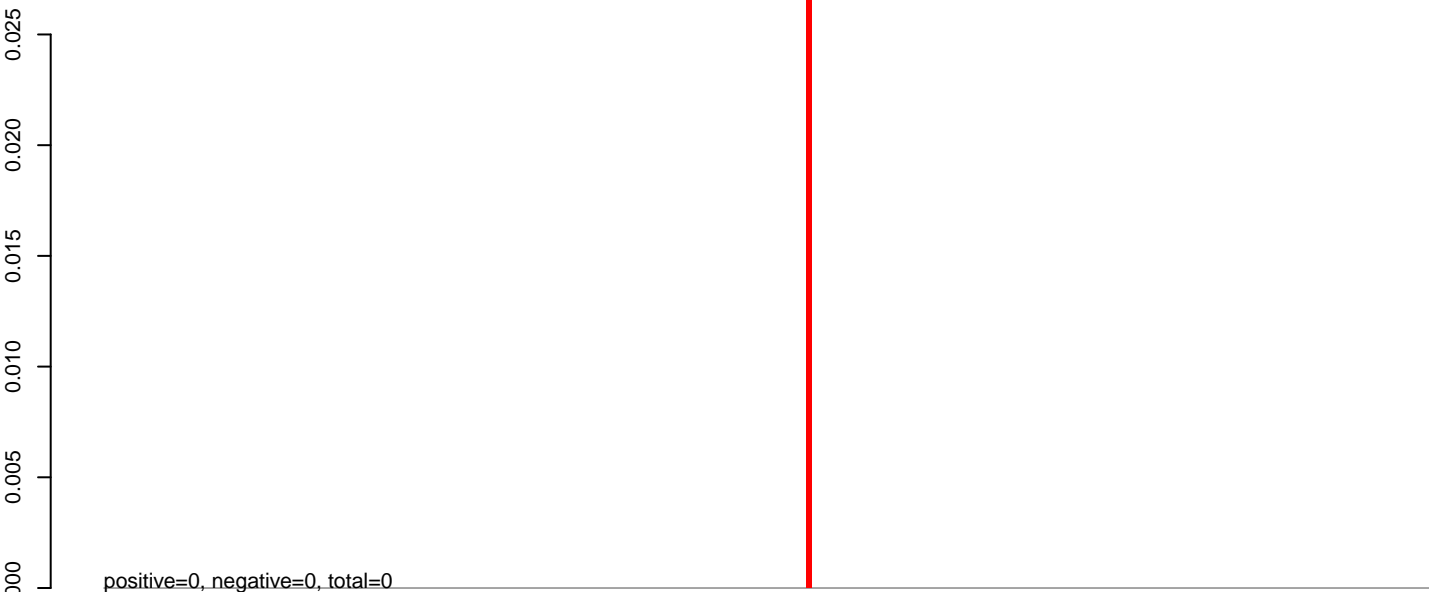
AeAeg_Ovary_GH.rep



si_chupusiko_viegat050102411M26698_RNA-dependent_RNA_polymerase_putative_glycoprotein_and_hypothetical_protein_genes:1-11280

0 2000 4000 6000 8000 10000

AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

positive=0, negative=0, total=0

AeAeg_Ovary_GH.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

positive=0, negative=0, total=0

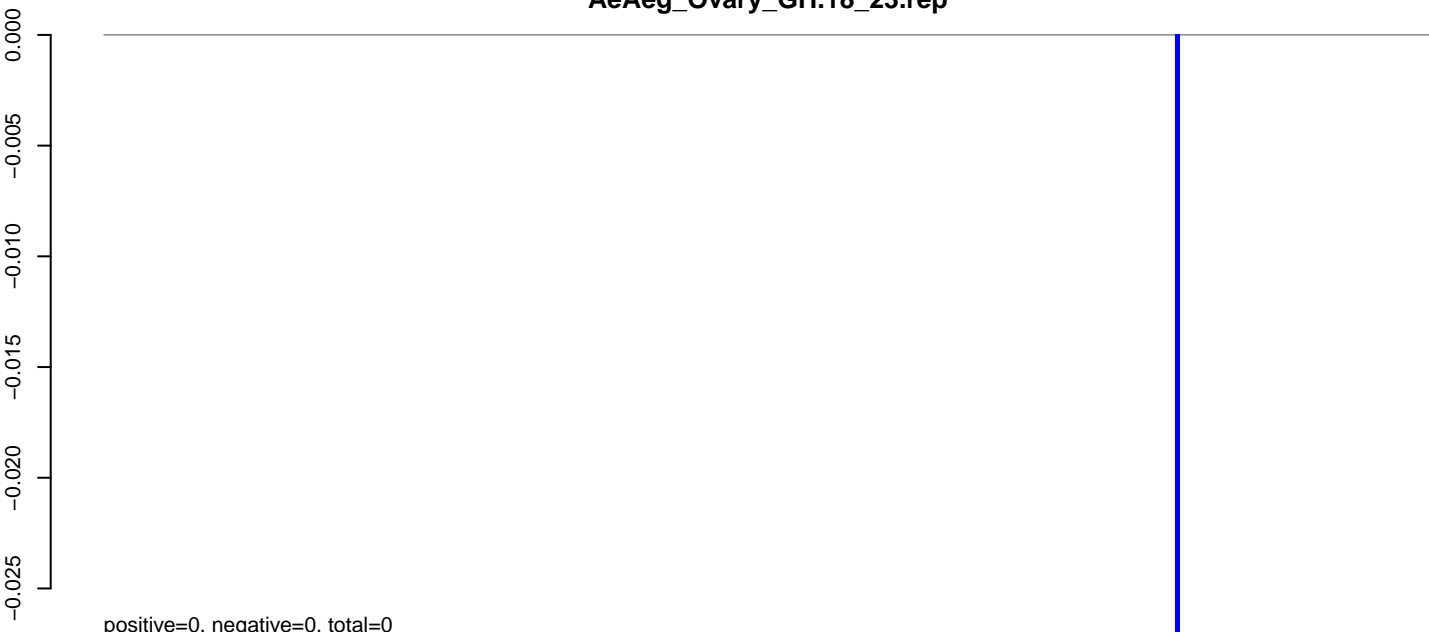
AeAeg_Ovary_GH.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

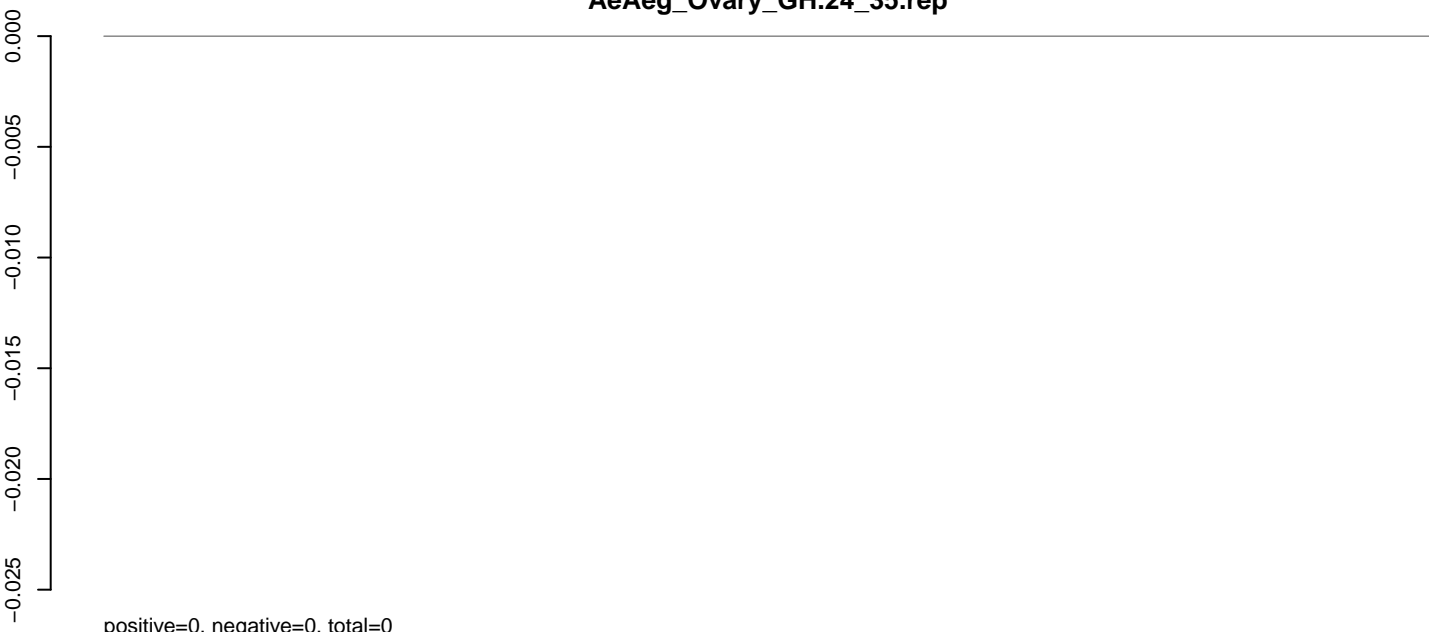
io_Viruses_innby413-ORF1-ORF1_ORF2_ORF2_ORF3_ORF3_glycoprotein_G_and_RNA-dependent_RNA_polymerases_genes:1-13706

0 2000 4000 6000 8000 10000 12000 14000

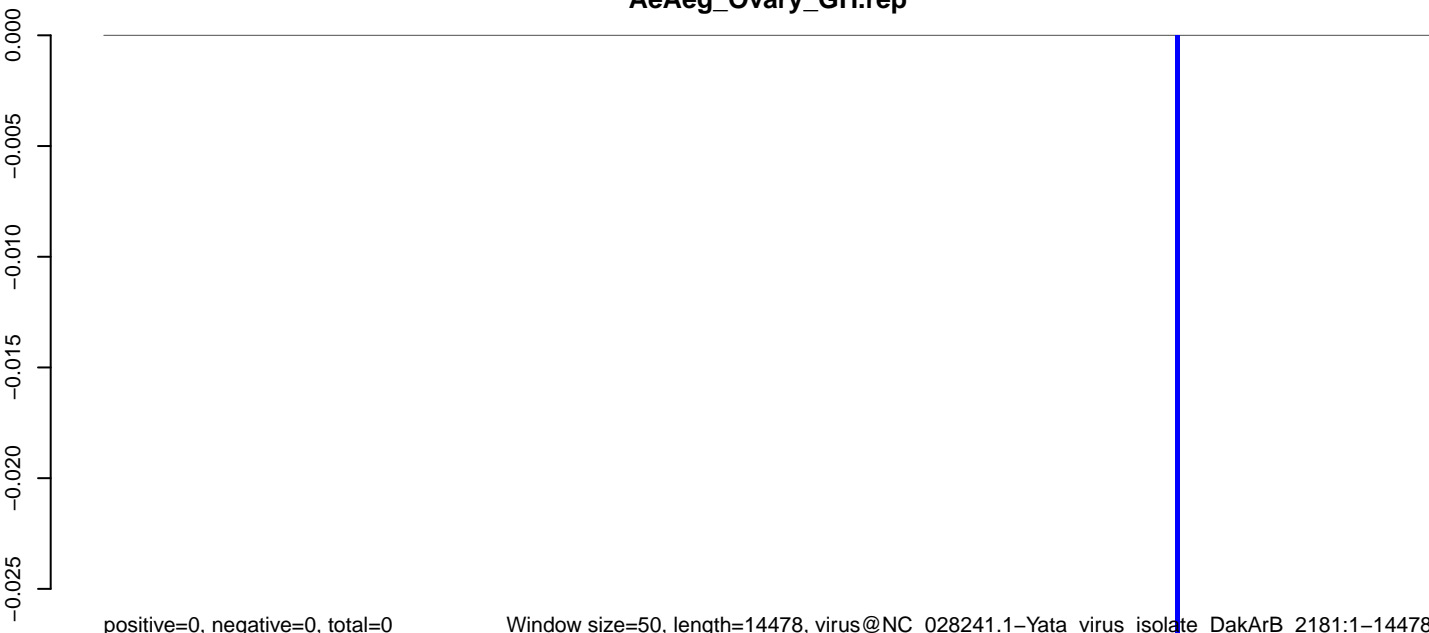
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



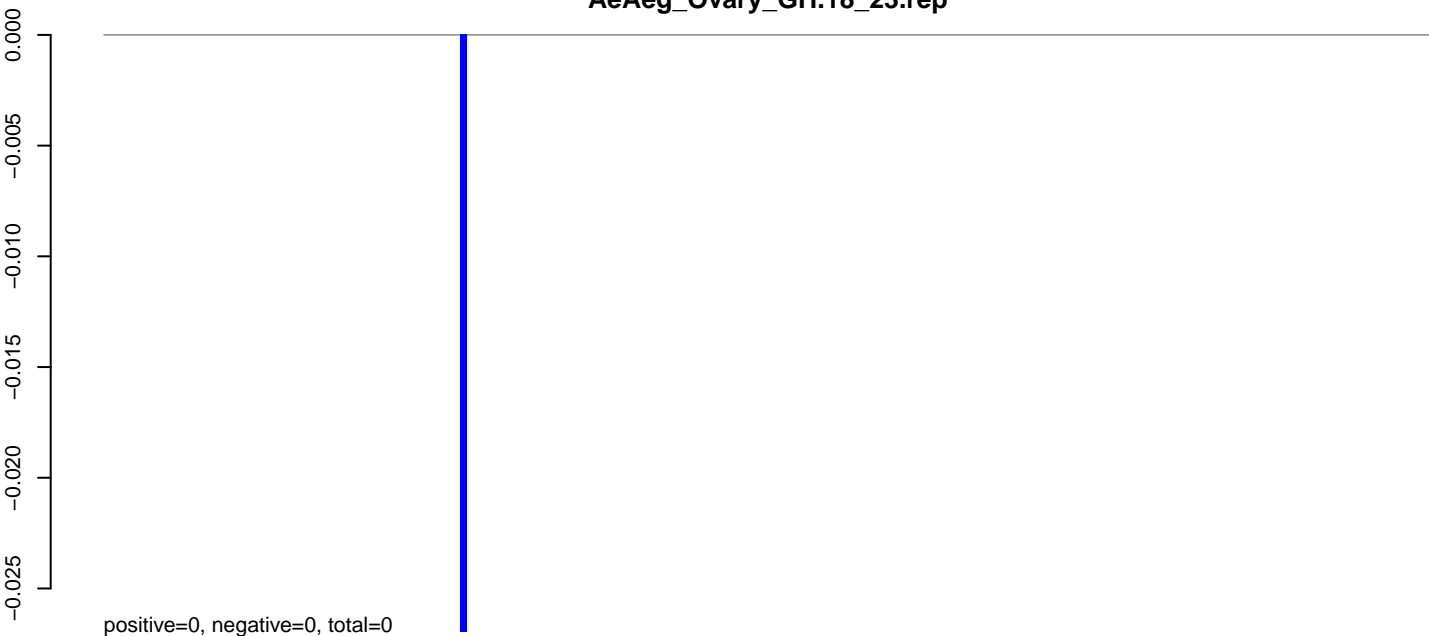
AeAeg_Ovary_GH.rep



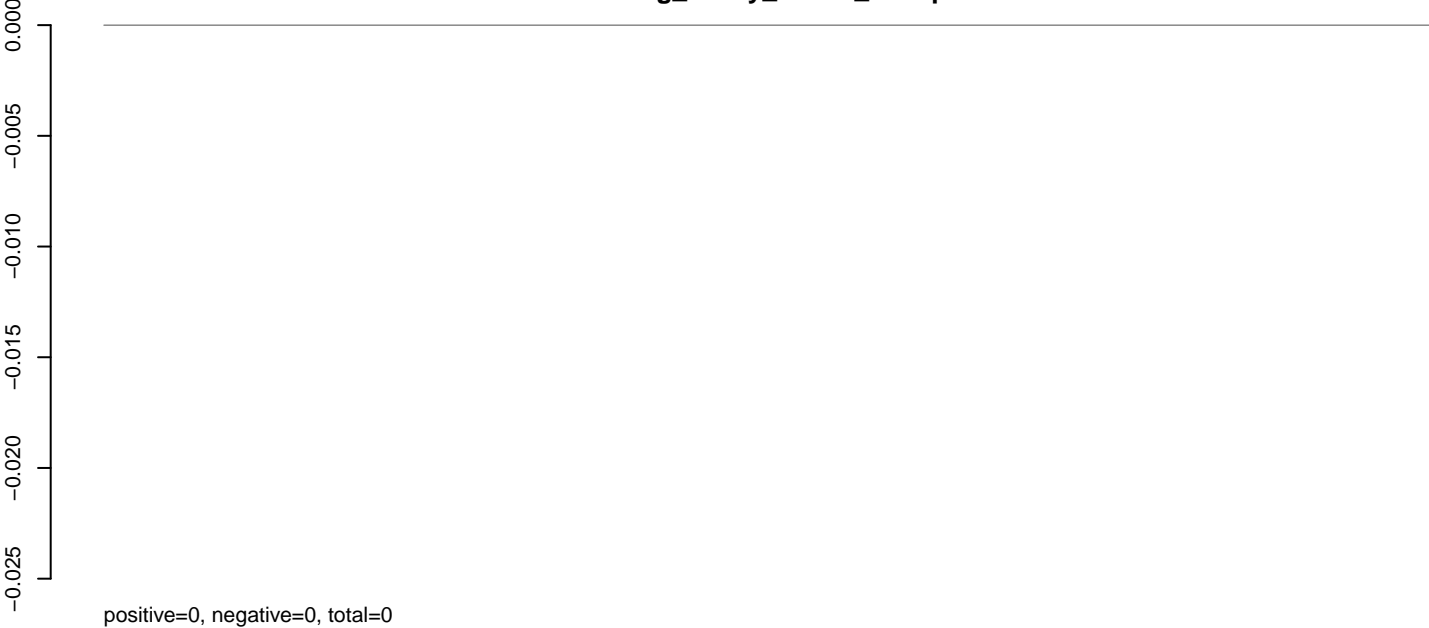
Window size=50, length=14478, virus@NC_028241.1-Yata_virus_isolate_DakArB_2181:1-14478

0 5000 10000 15000

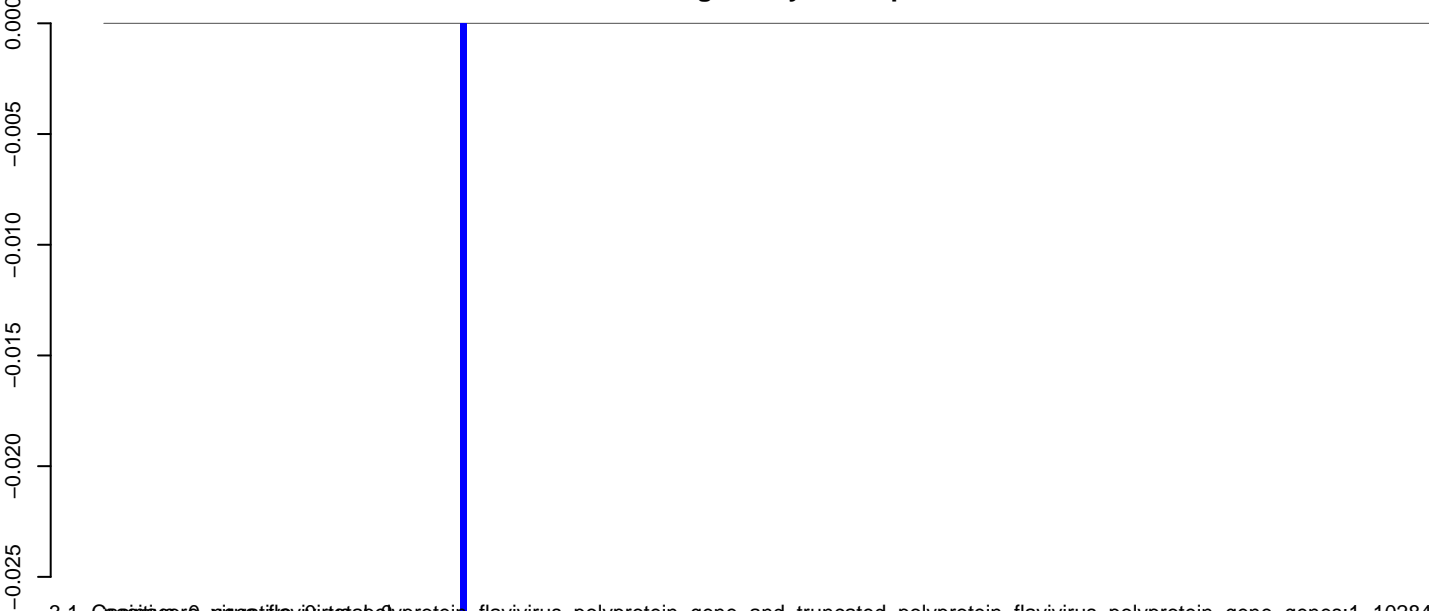
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



3.1-Opisao=0, negativ=0, total=0 protein_flavivirus_polyprotein_gene_and_truncated_polyprotein_flavivirus_polyprotein_gene_genes:1-10284

0 2000 4000 6000 8000 10000

AeAeg_Ovary_GH.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

positive=0, negative=0, total=0



AeAeg_Ovary_GH.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

positive=0, negative=0, total=0

AeAeg_Ovary_GH.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

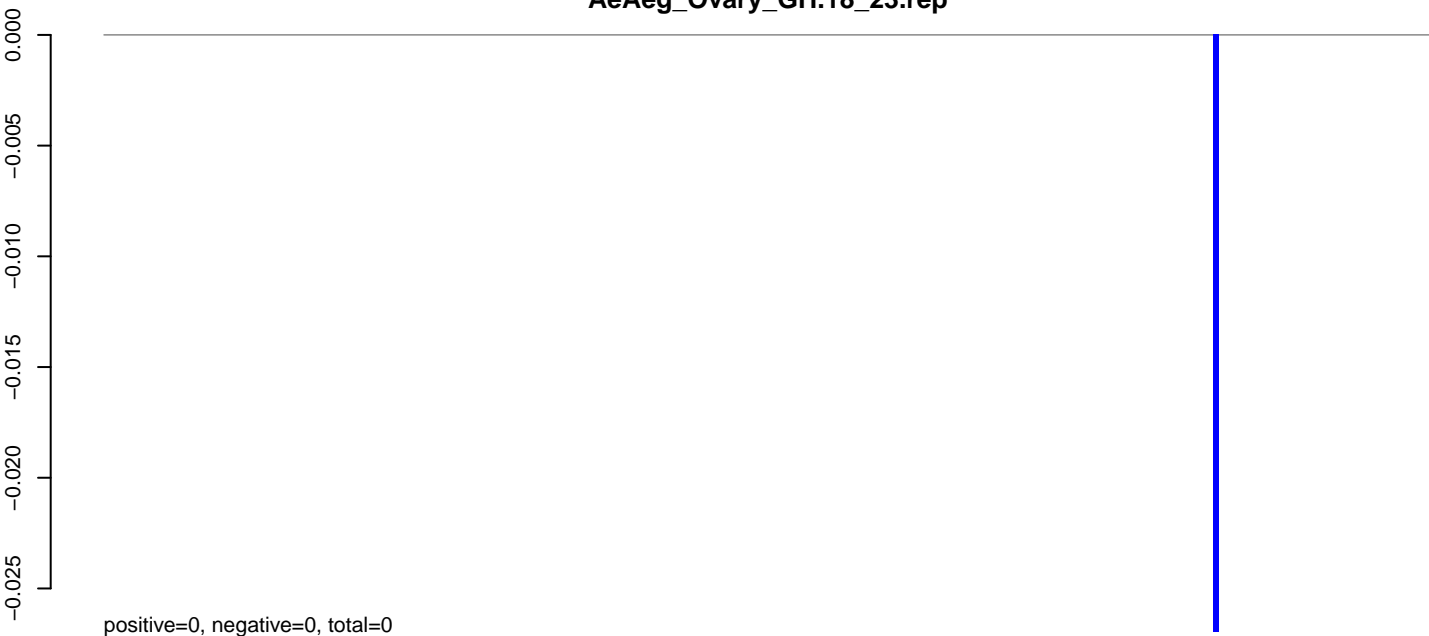
positive=0, negative=0, total=0

Window size=50, length=10125 virus@NC_026620.1-Jutiapa_virus_strain_JG-128:1-10125

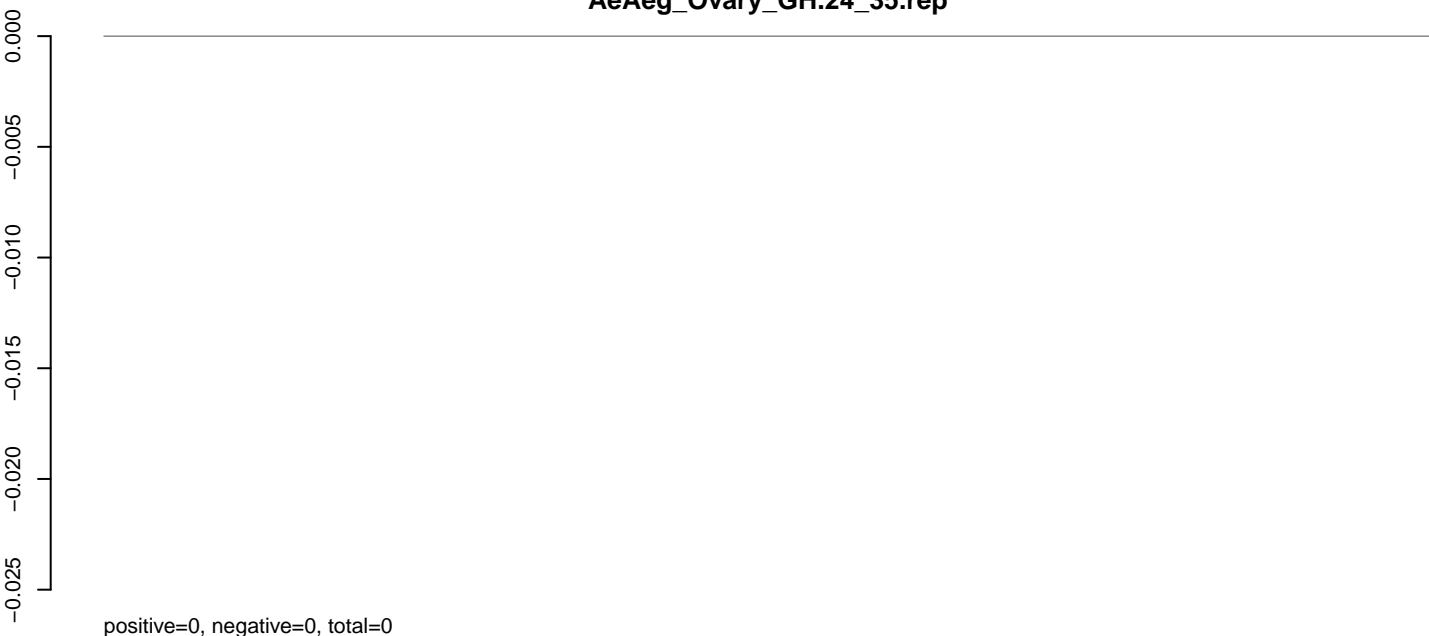


0 2000 4000 6000 8000 10000

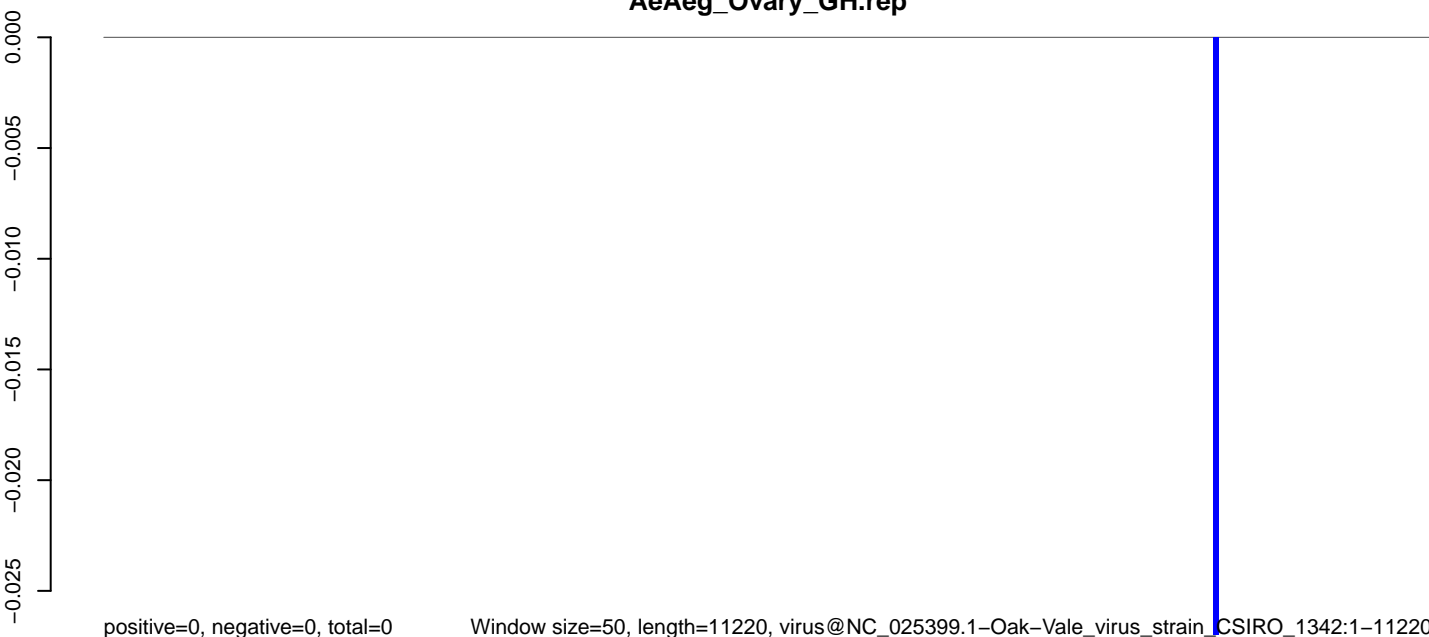
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

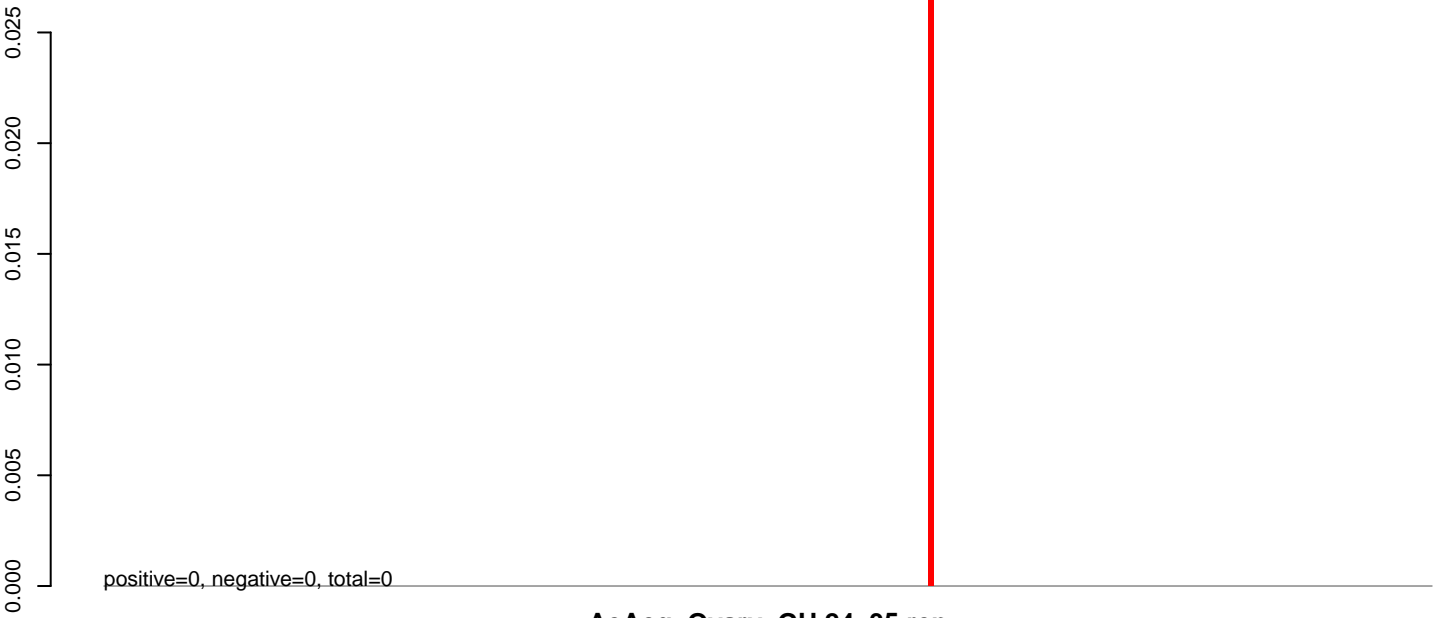


AeAeg_Ovary_GH.rep

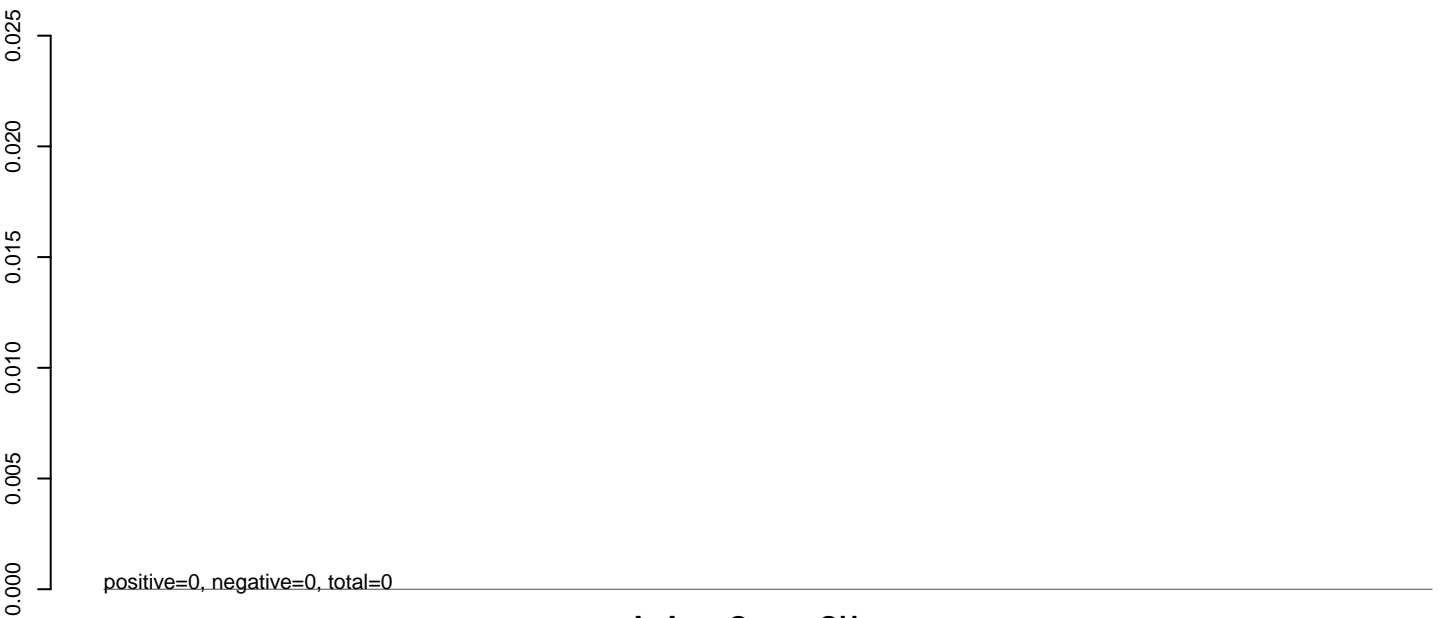


0 2000 4000 6000 8000 10000

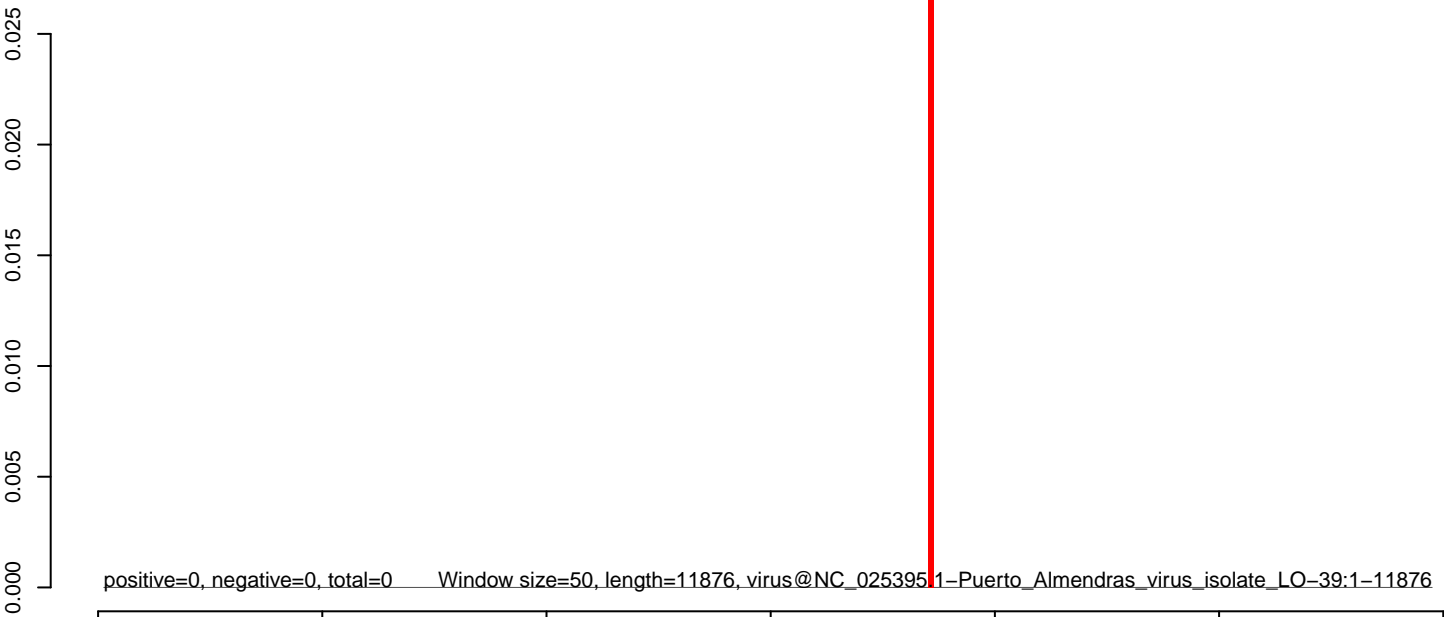
AeAeg_Ovary_GH.18_23.rep



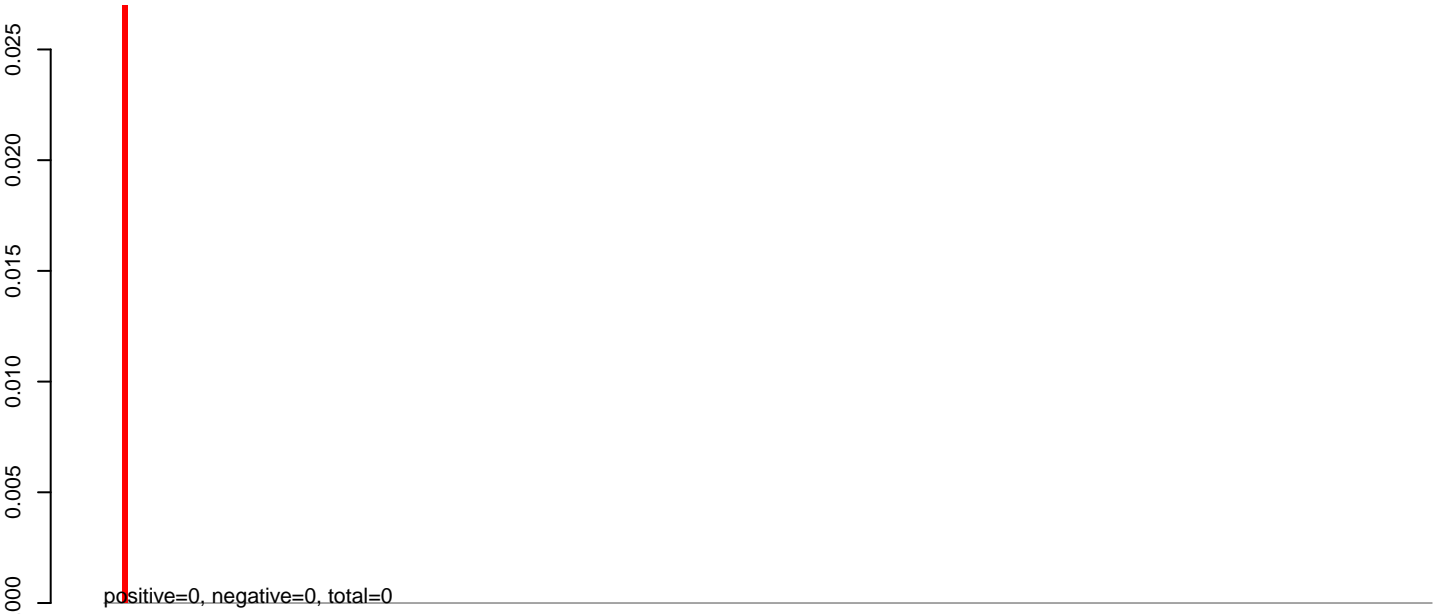
AeAeg_Ovary_GH.24_35.rep



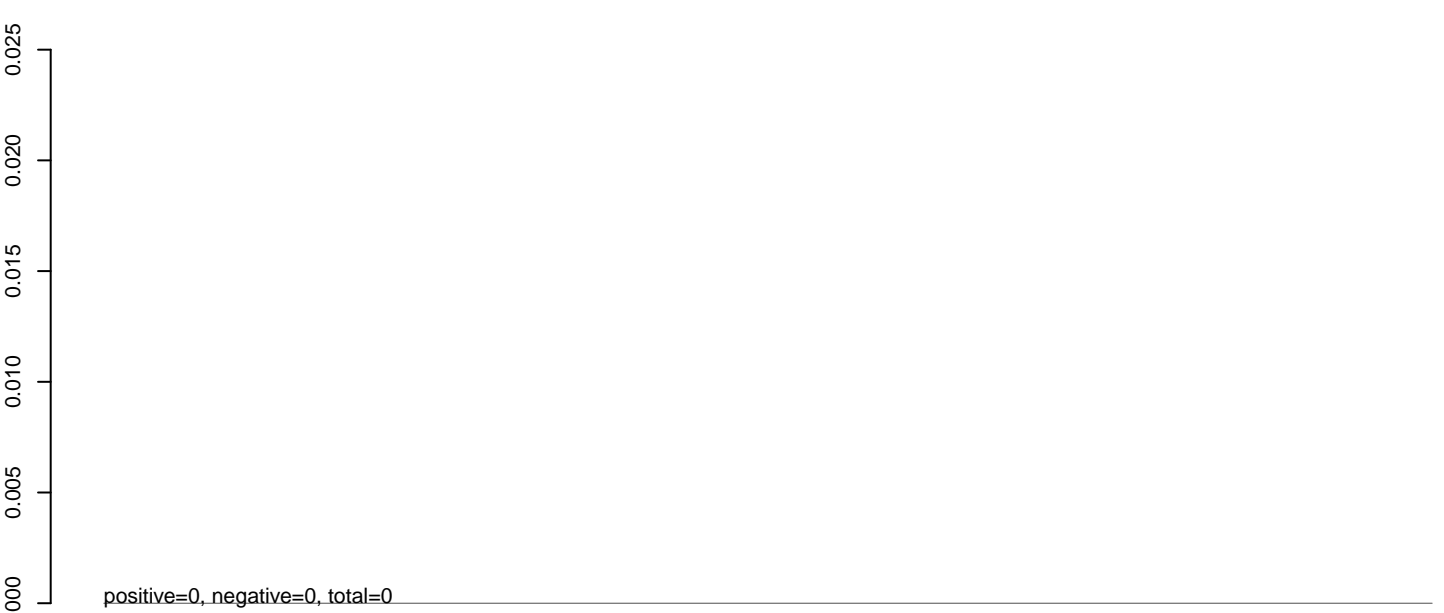
AeAeg_Ovary_GH.rep



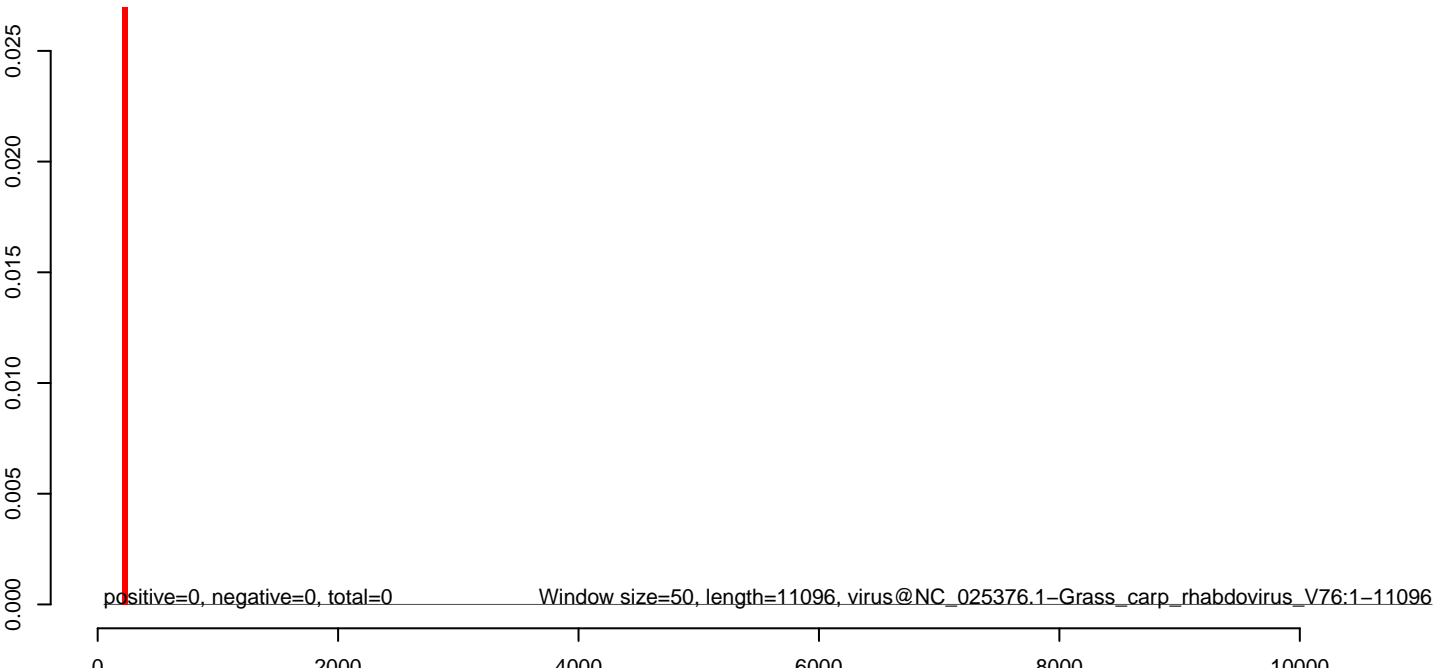
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



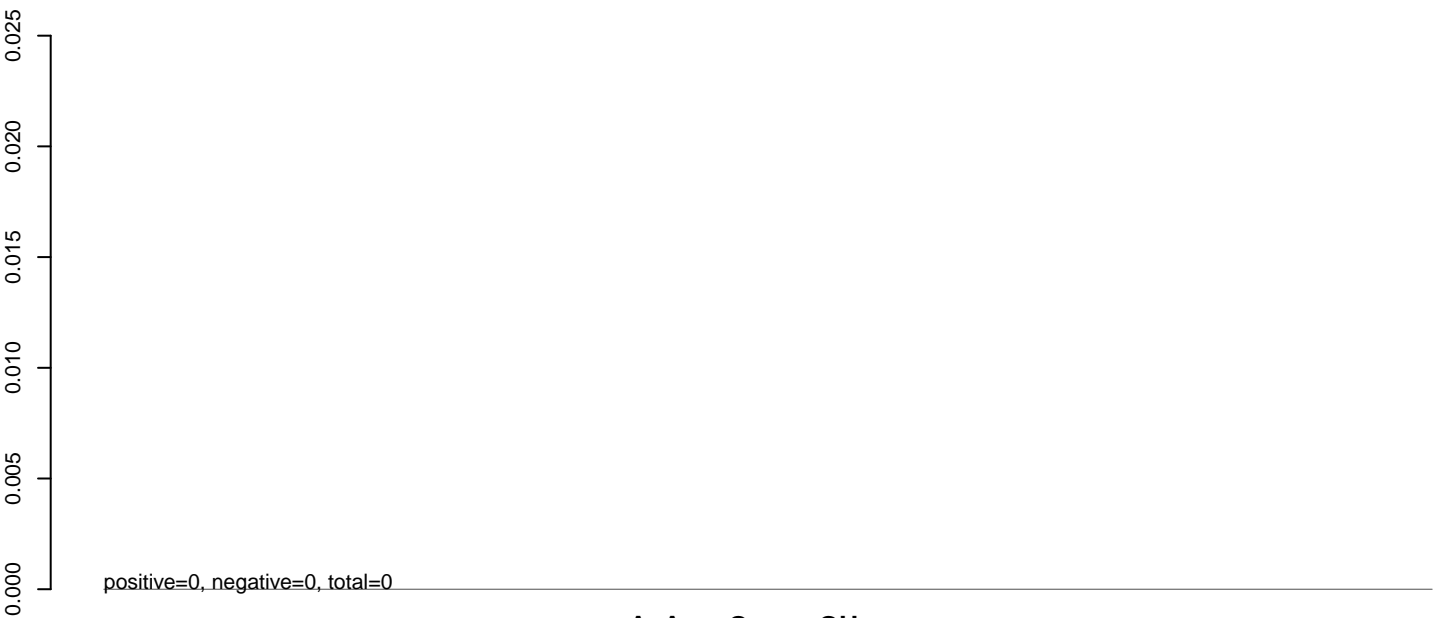
AeAeg_Ovary_GH.rep



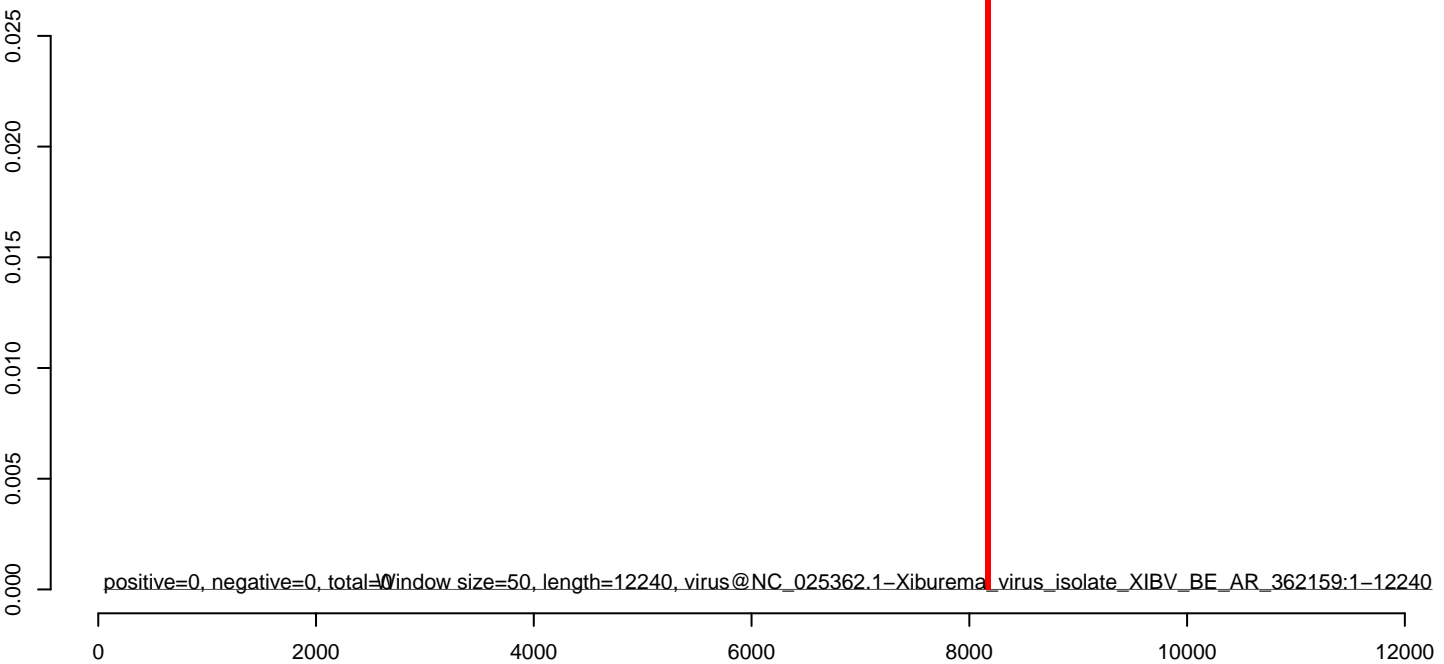
AeAeg_Ovary_GH.18_23.rep



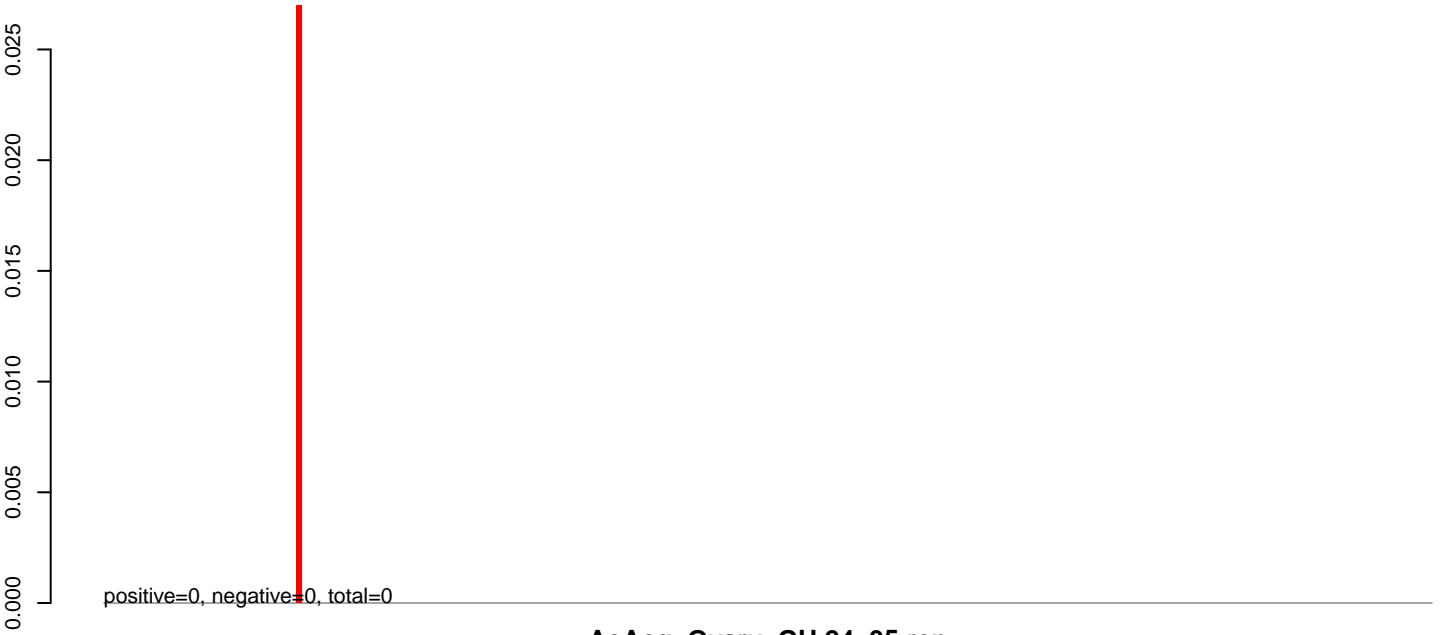
AeAeg_Ovary_GH.24_35.rep



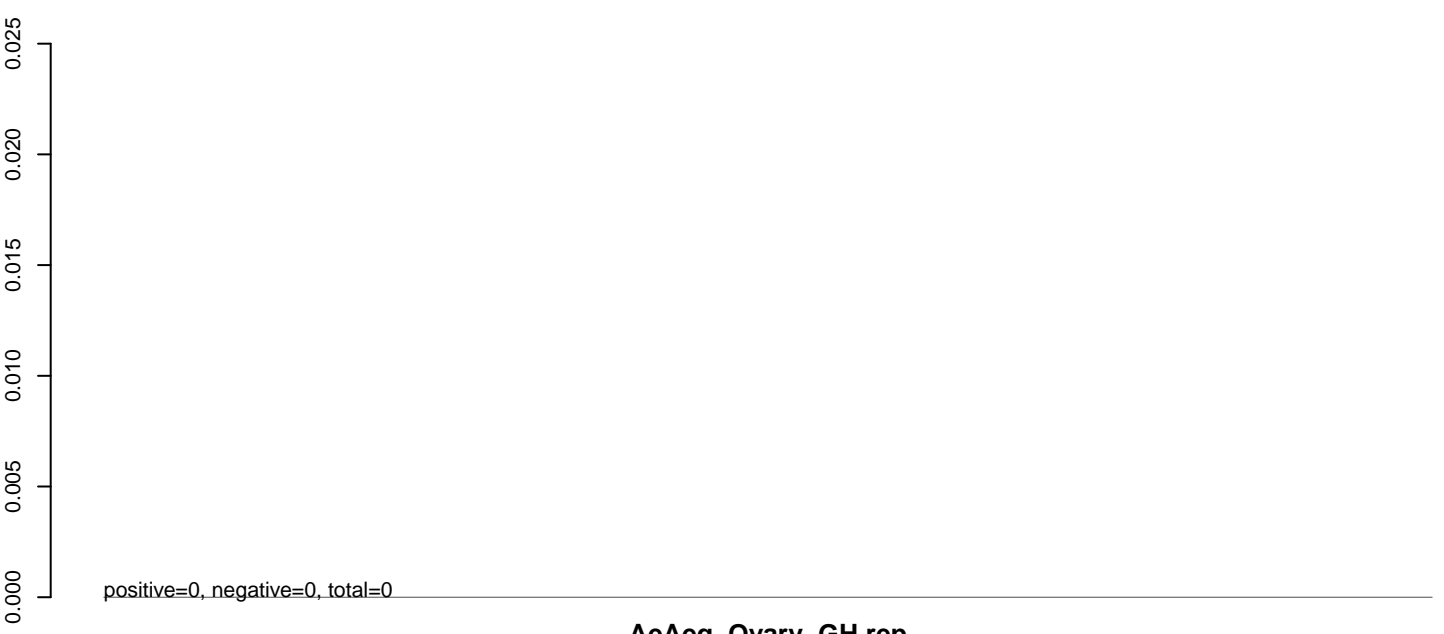
AeAeg_Ovary_GH.rep



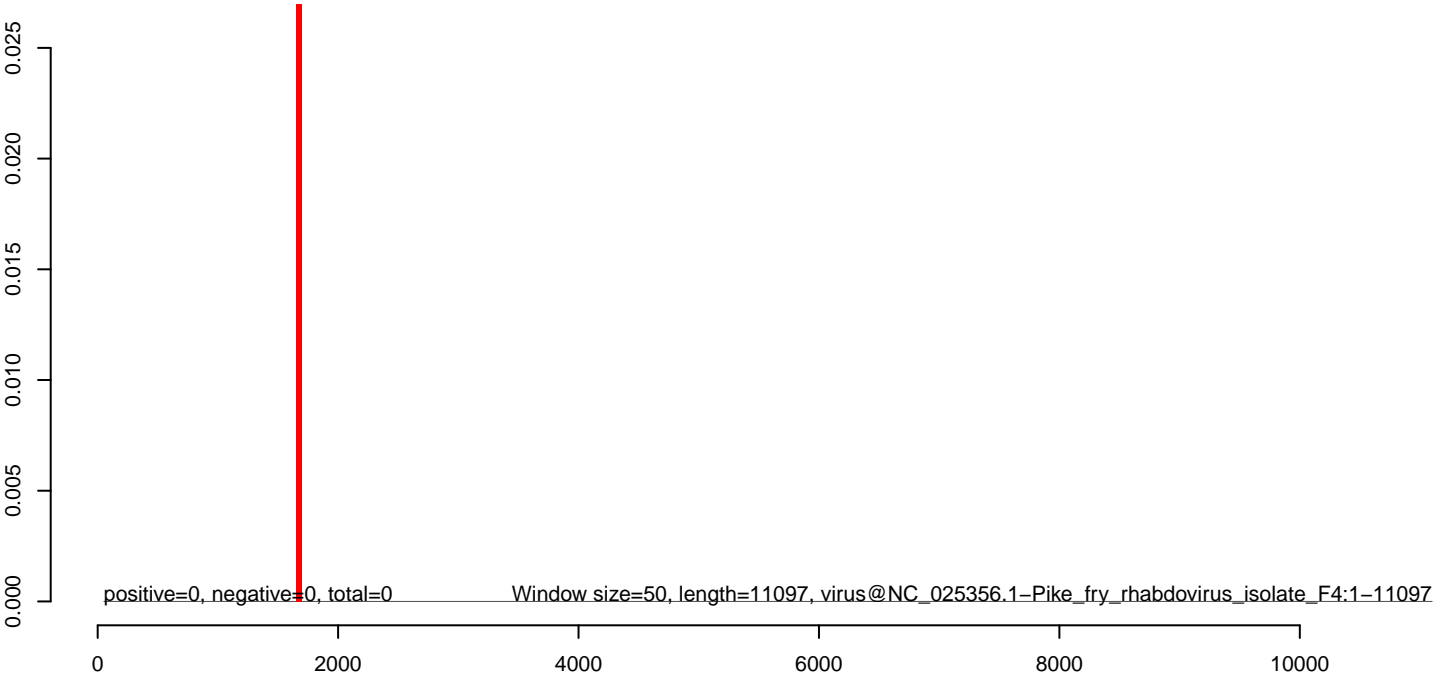
AeAeg_Ovary_GH.18_23.rep



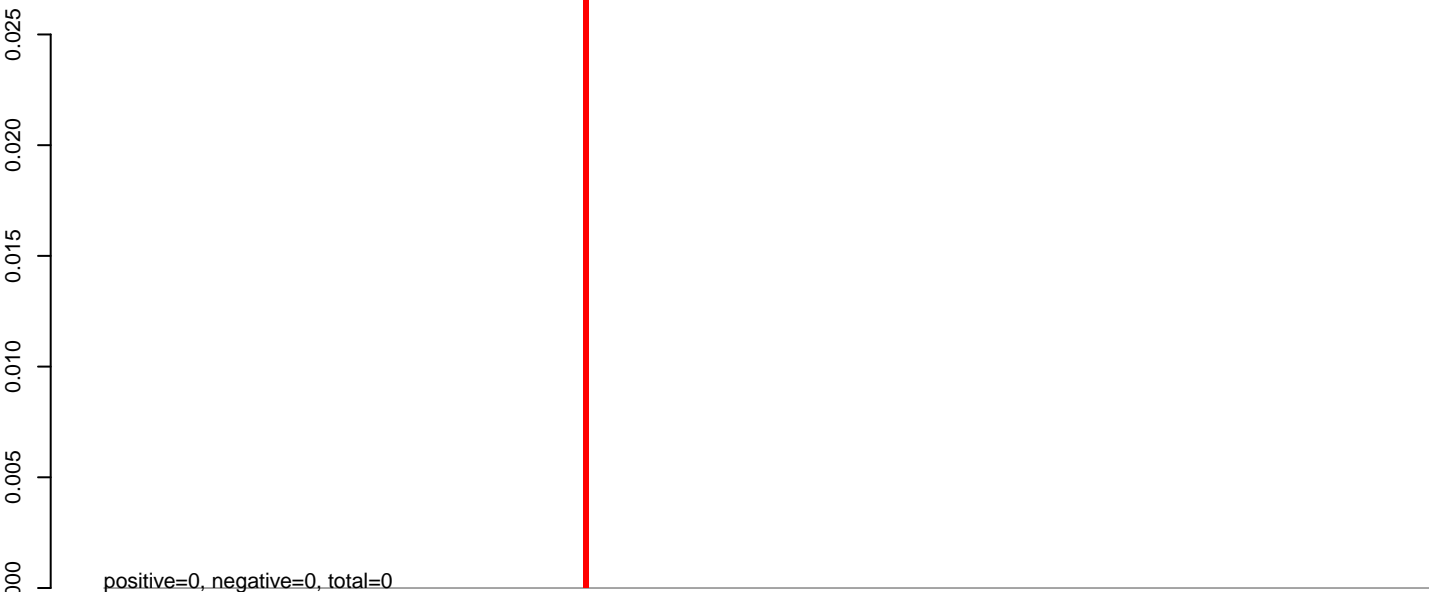
AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



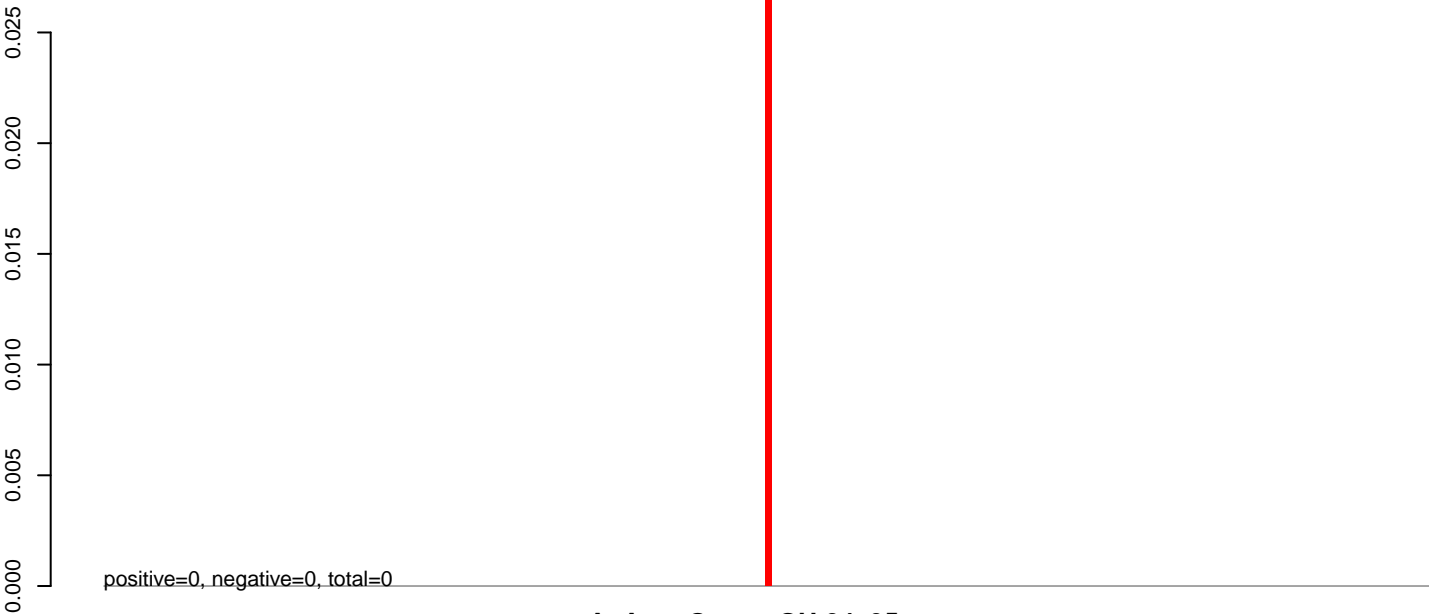
AeAeg_Ovary_GH.24_35.rep



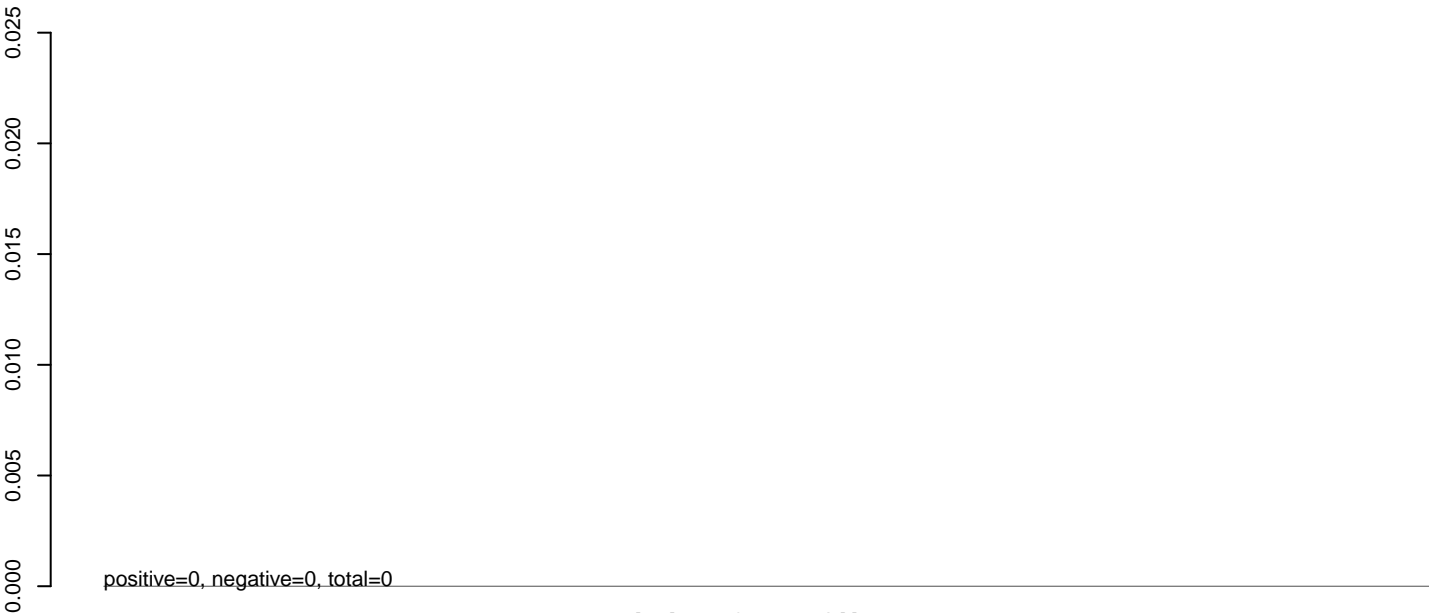
AeAeg_Ovary_GH.rep



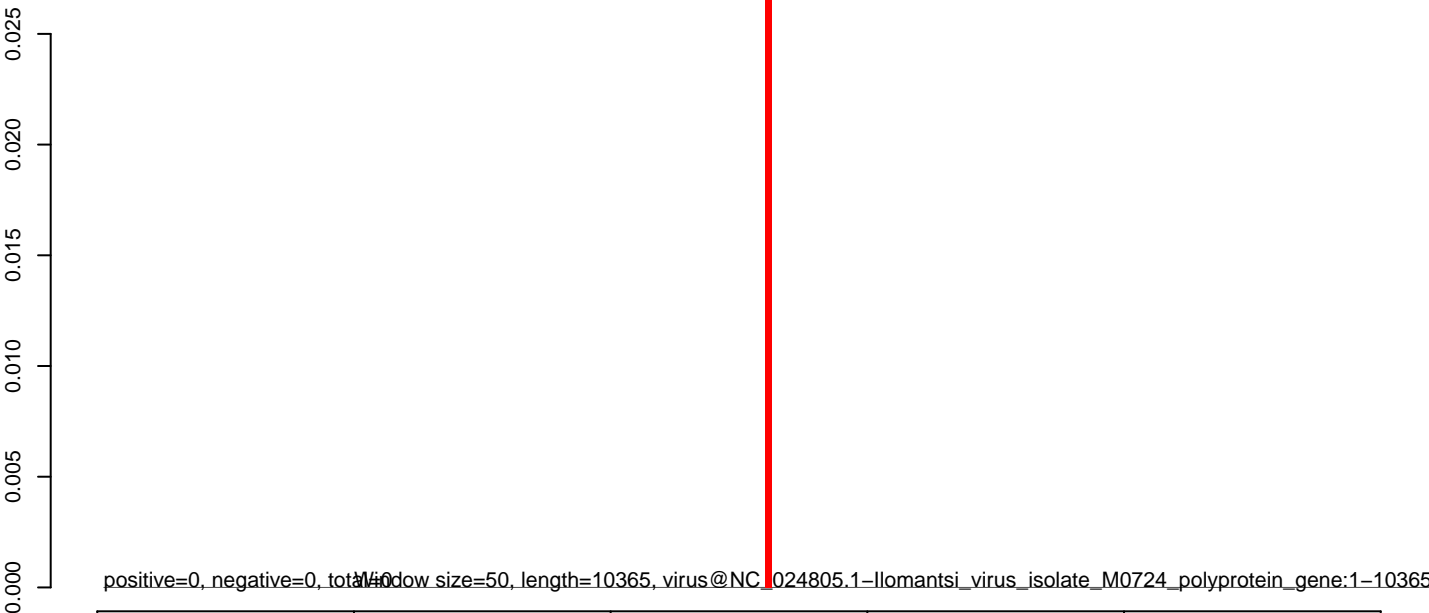
AeAeg_Ovary_GH.18_23.rep



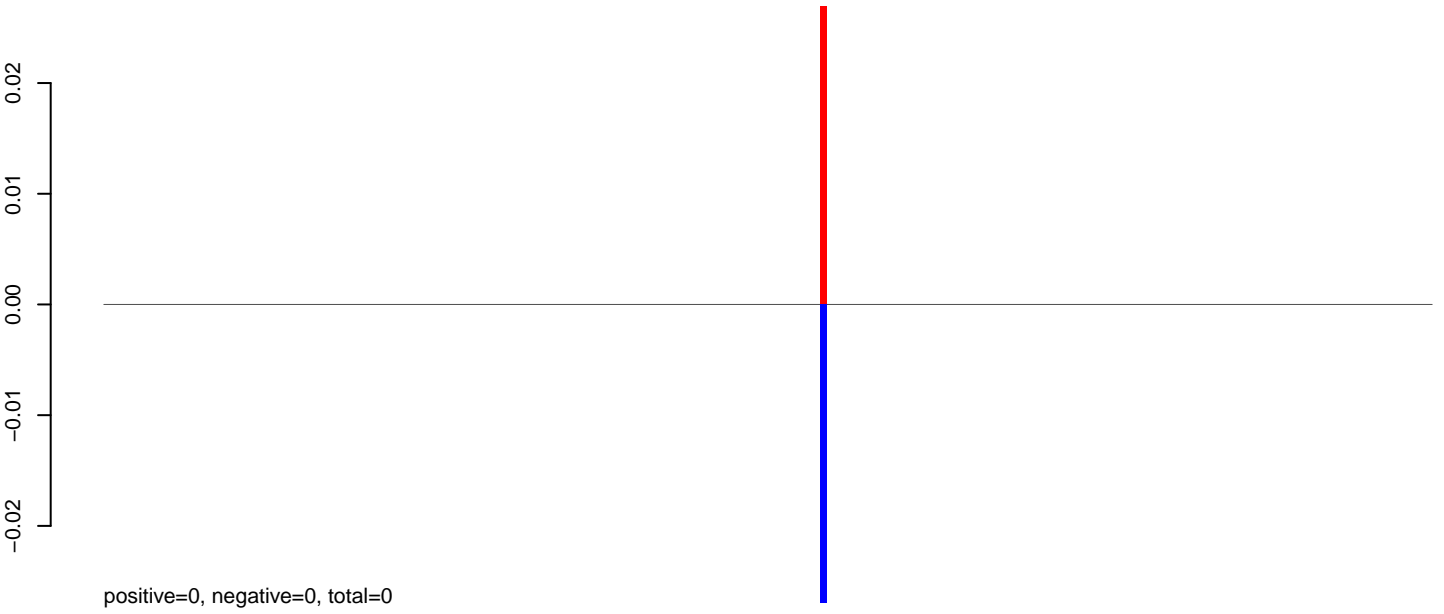
AeAeg_Ovary_GH.24_35.rep



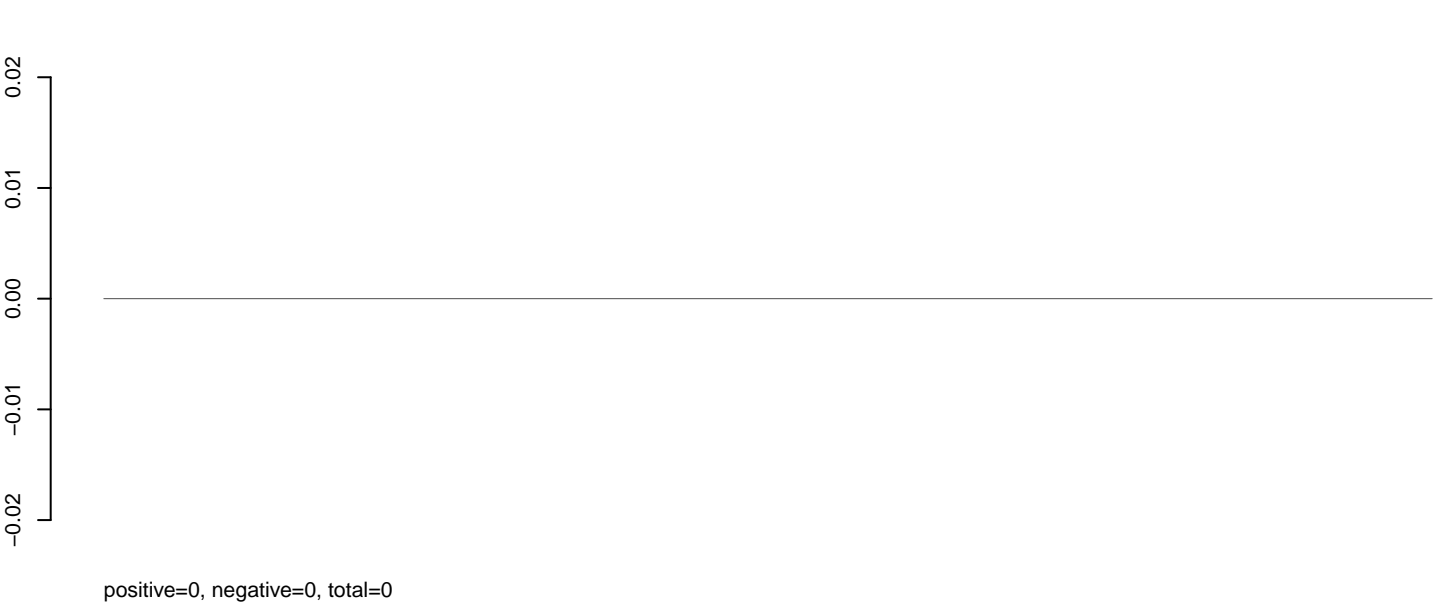
AeAeg_Ovary_GH.rep



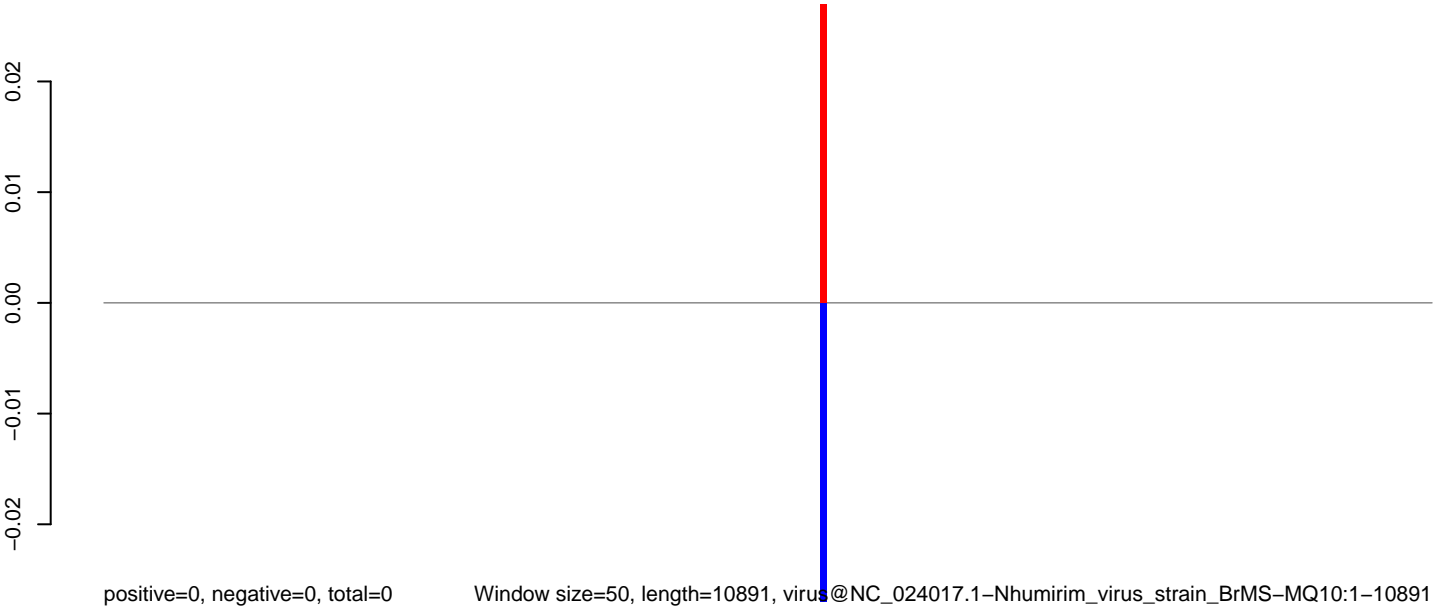
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



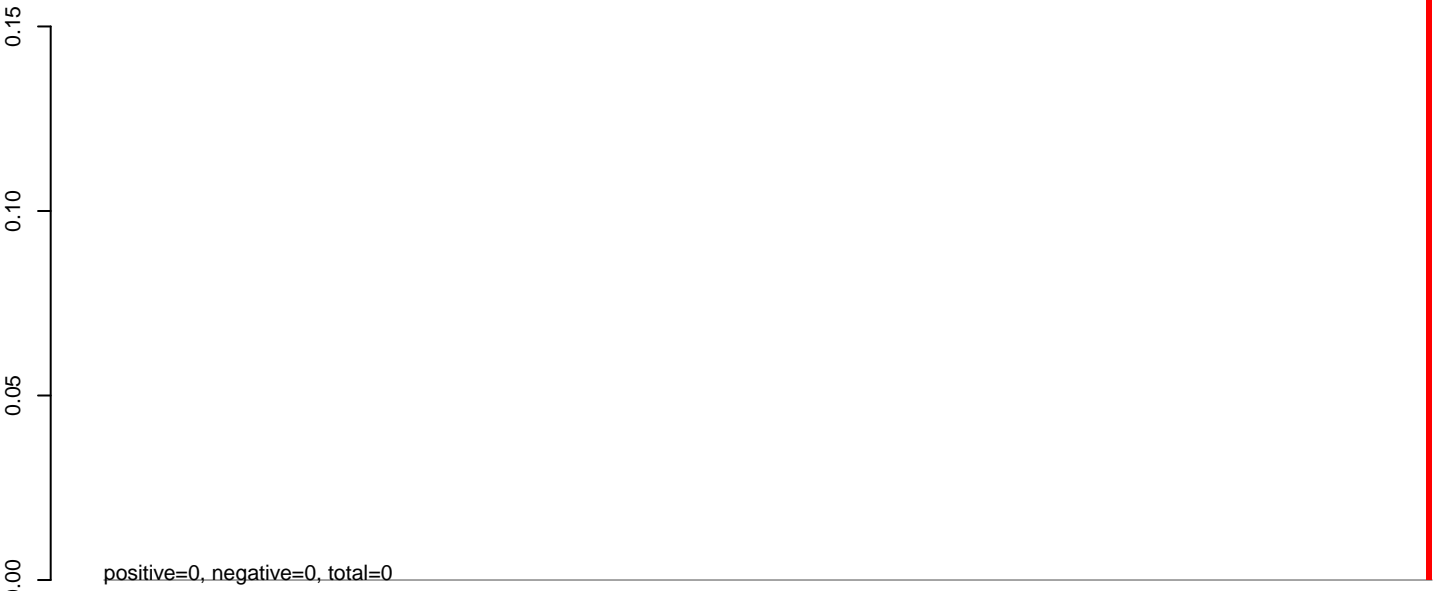
AeAeg_Ovary_GH.rep



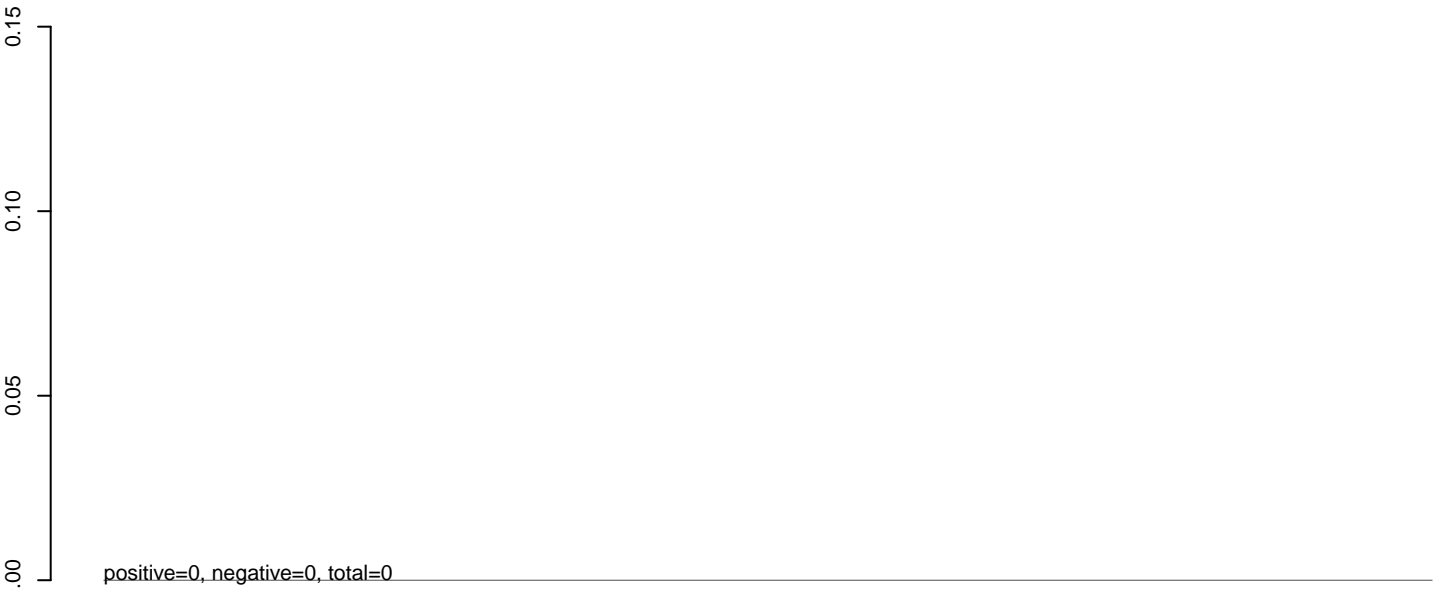
Window size=50, length=10891, virus@NC_024017.1-Nhumirim_virus_strain_BrMS-MQ10:1-10891

0 2000 4000 6000 8000 10000

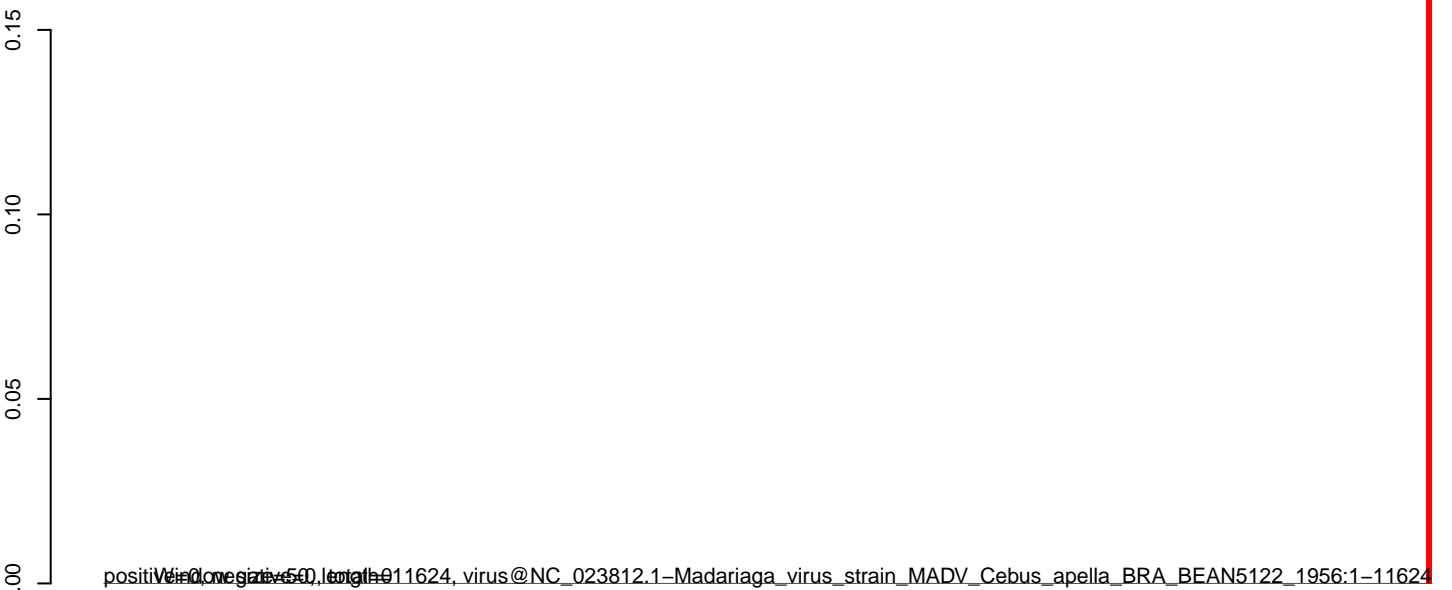
AeAeg_Ovary_GH.18_23.rep



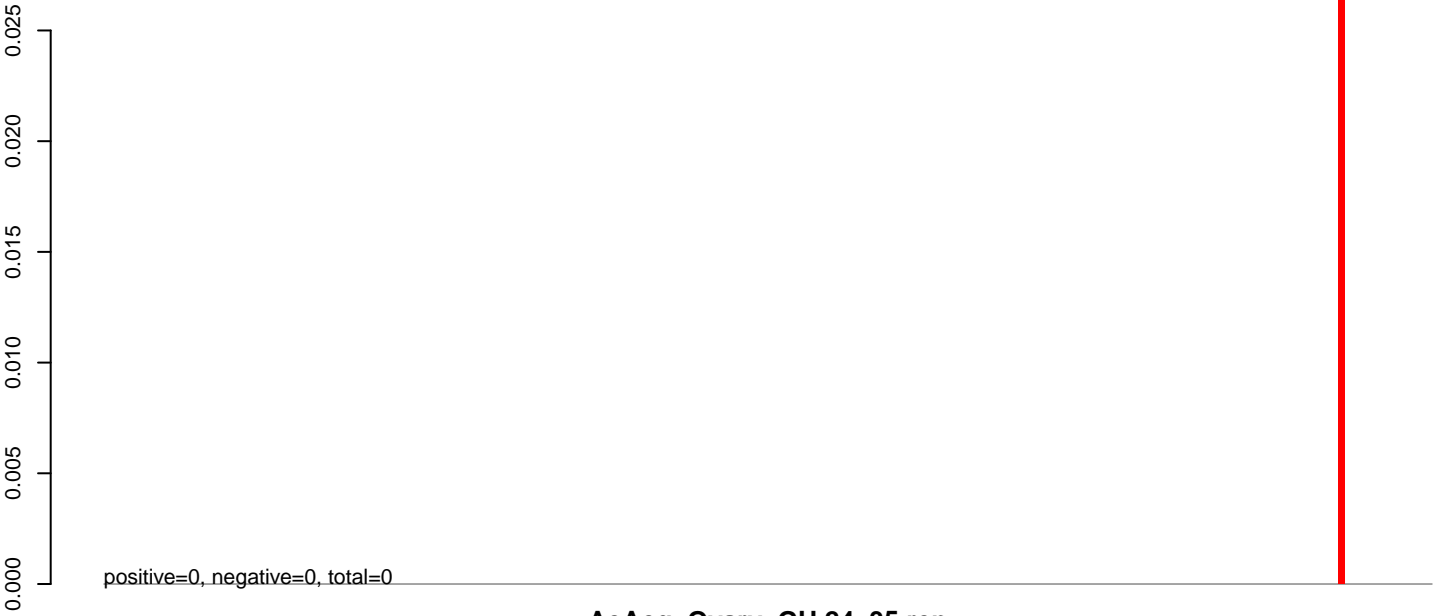
AeAeg_Ovary_GH.24_35.rep



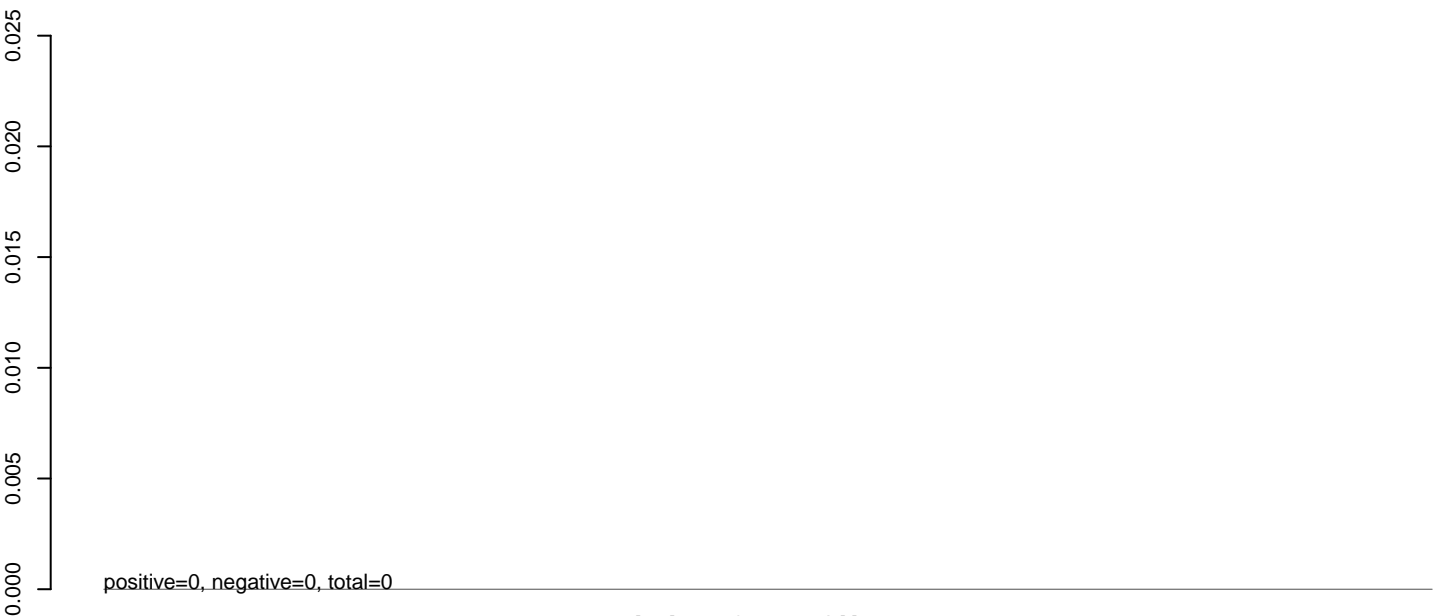
AeAeg_Ovary_GH.rep



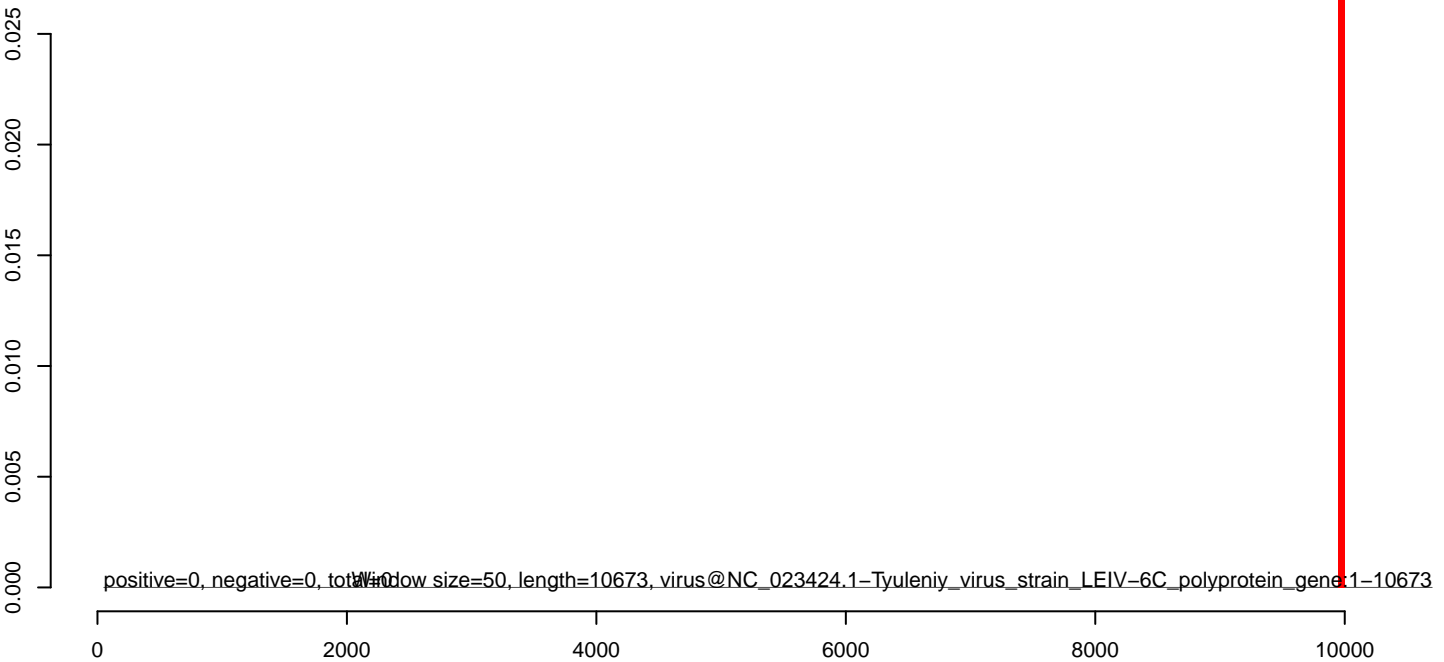
AeAeg_Ovary_GH.18_23.rep



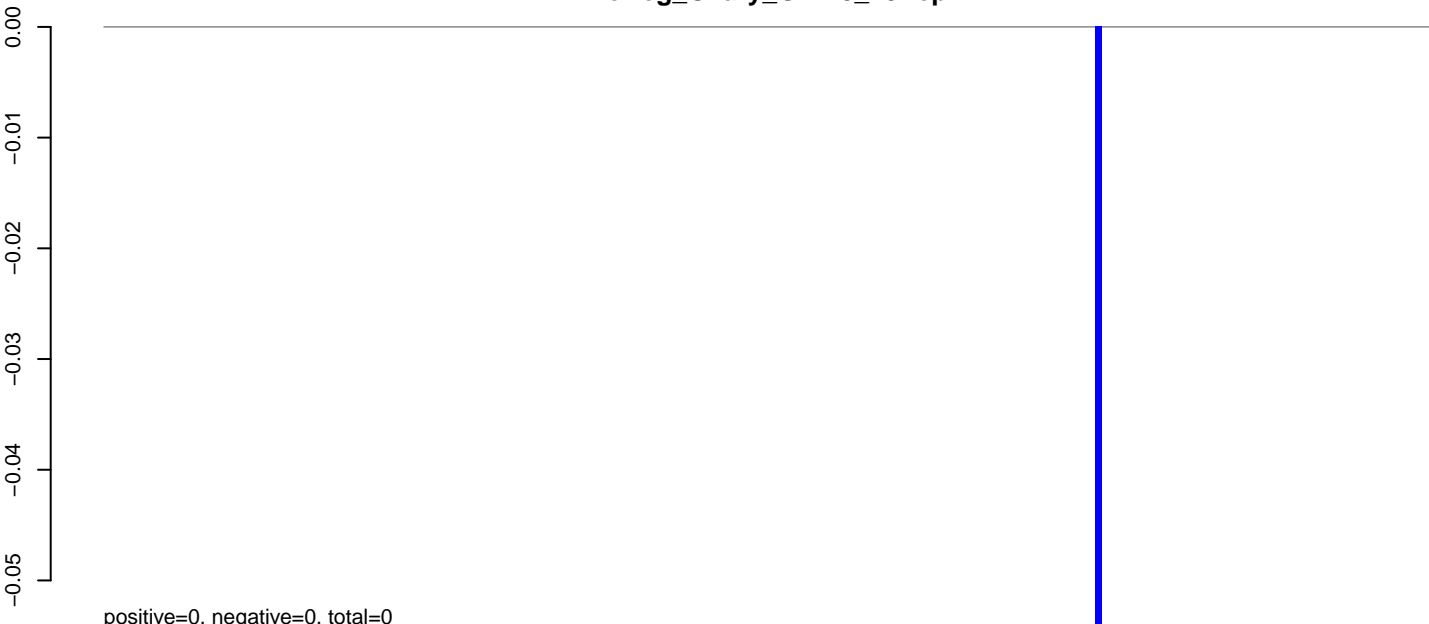
AeAeg_Ovary_GH.24_35.rep



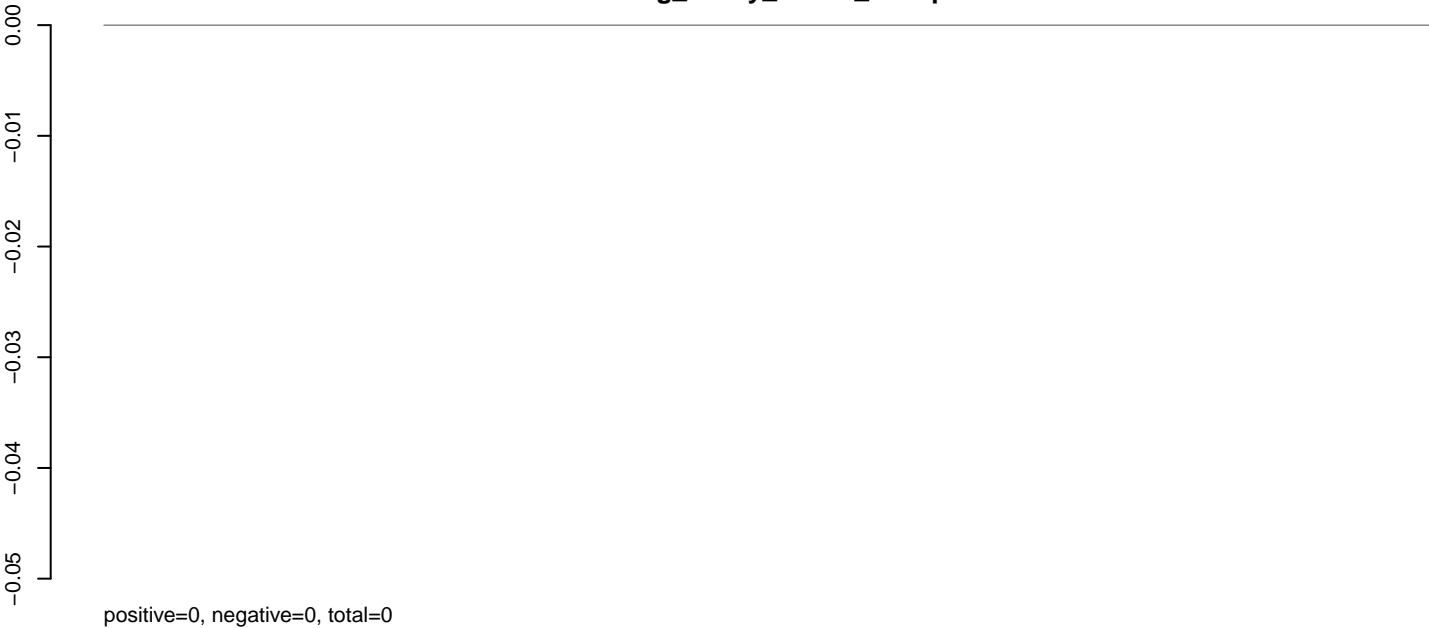
AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



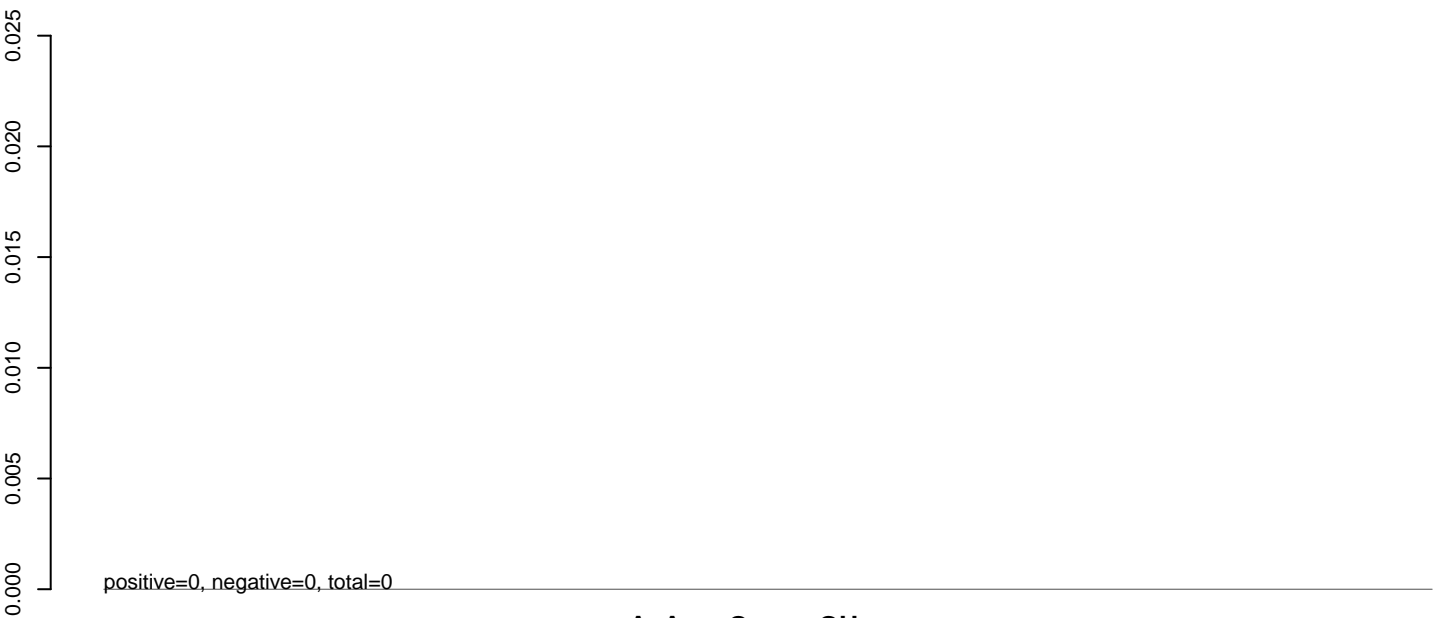
AeAeg_Ovary_GH.rep



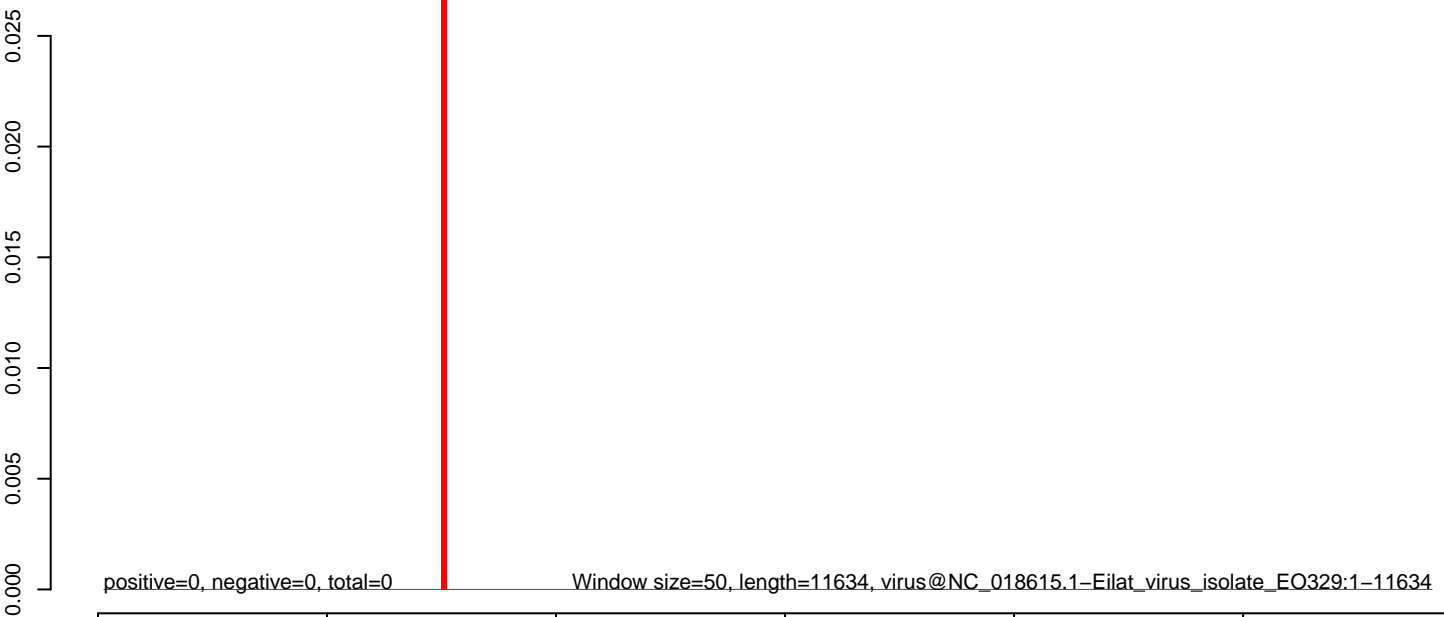
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

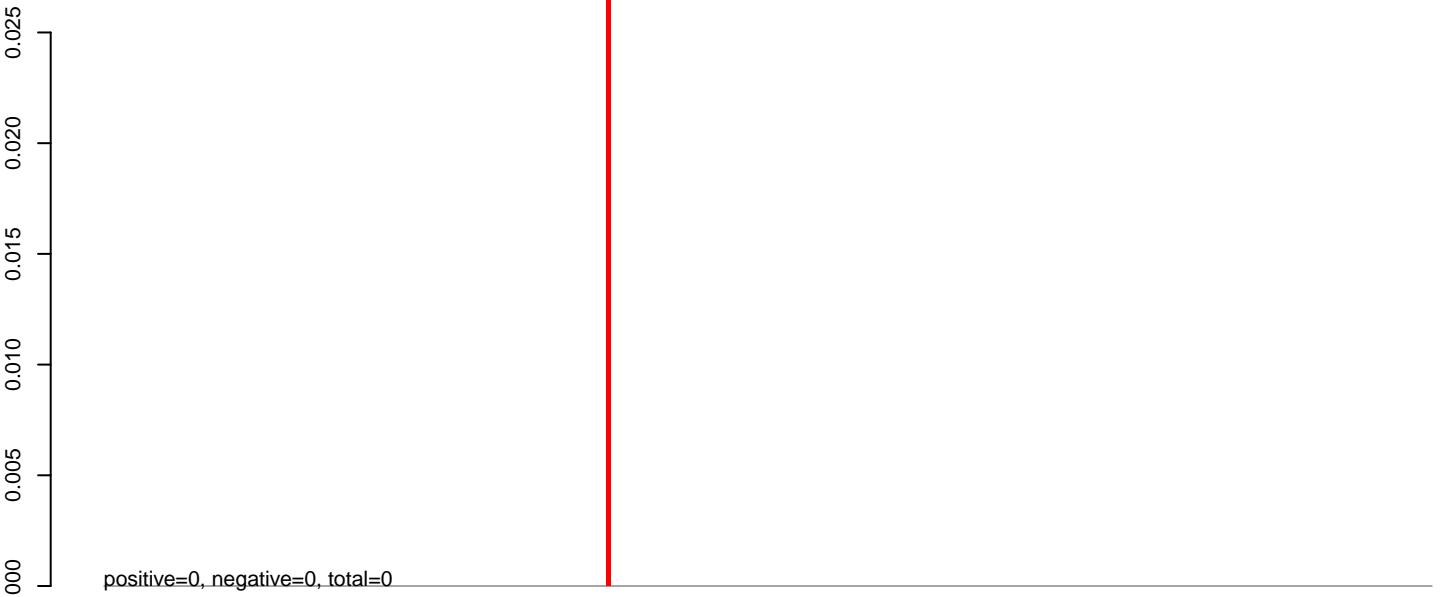


AeAeg_Ovary_GH.rep

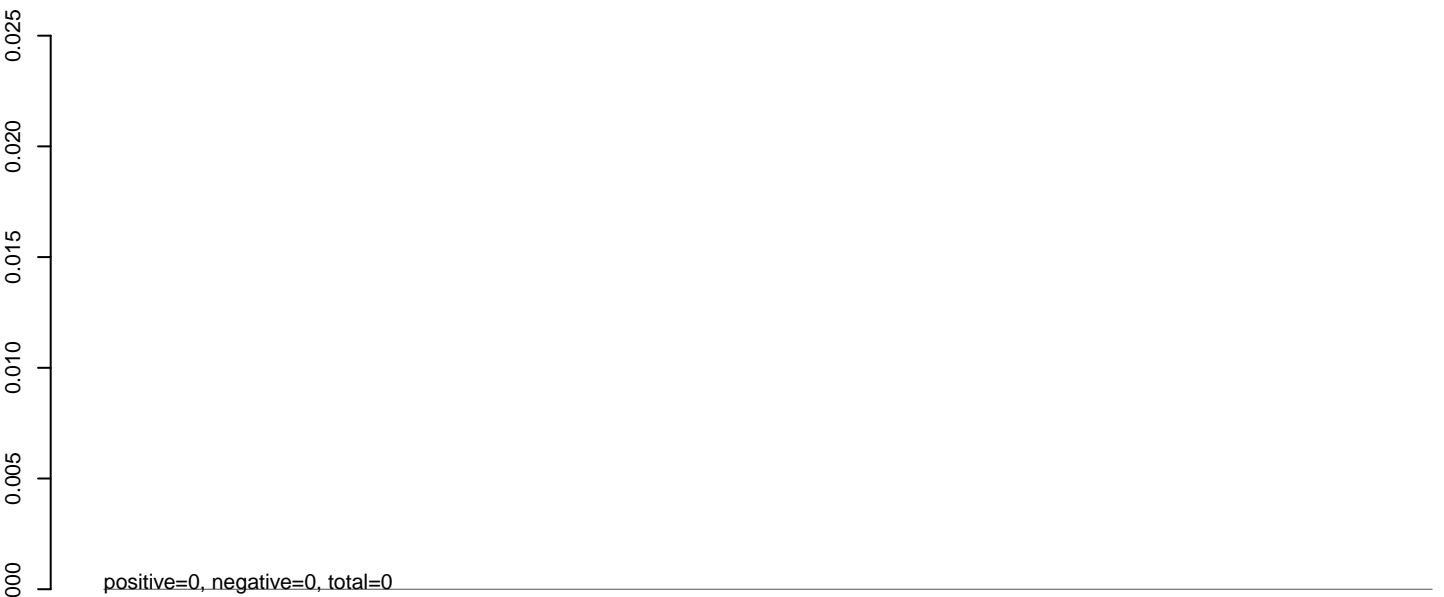


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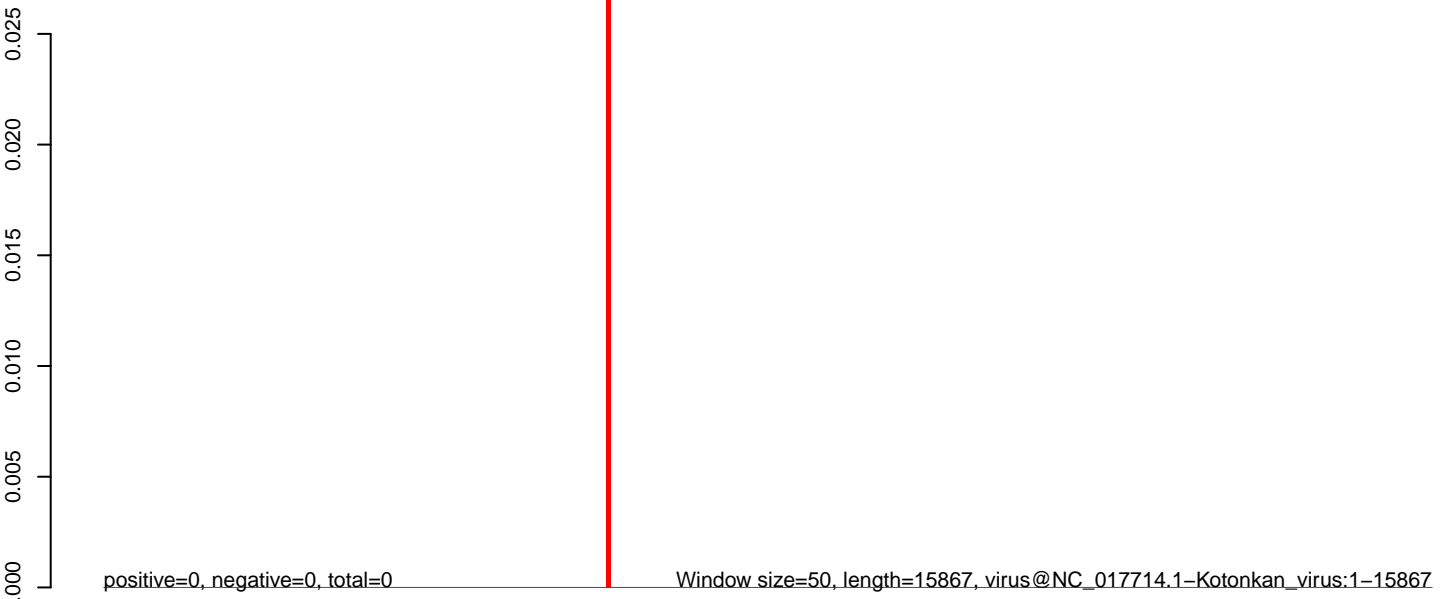
AeAeg_Ovary_GH.18_23.rep



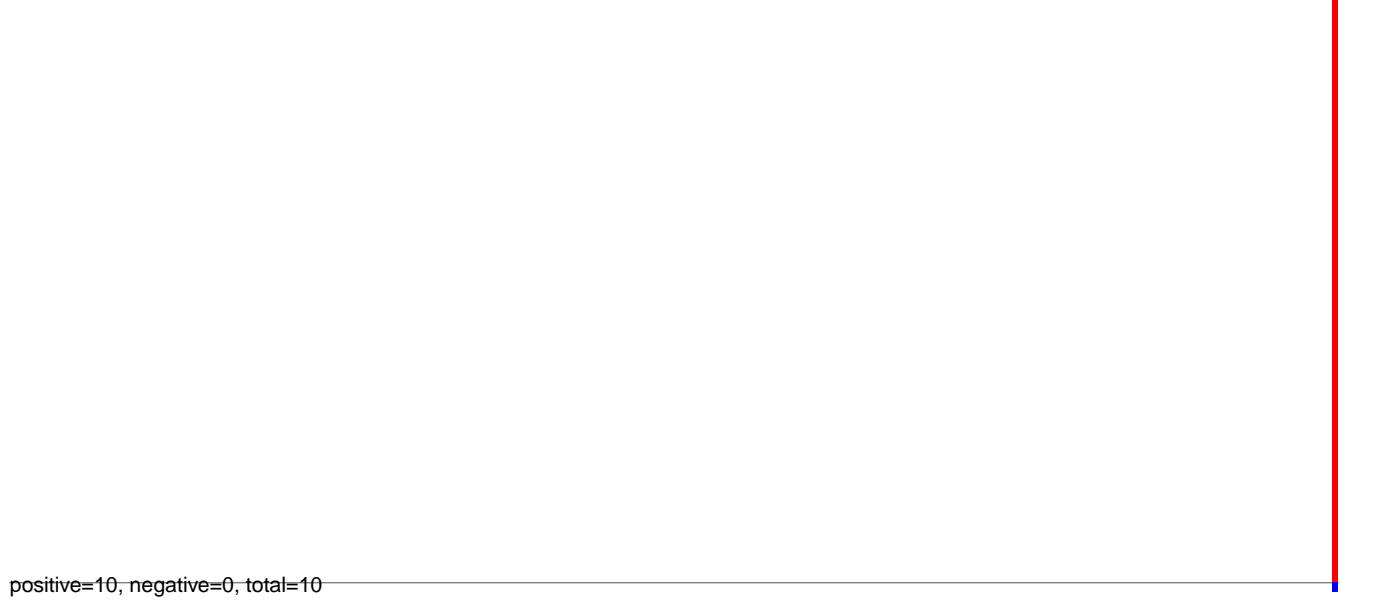
AeAeg_Ovary_GH.24_35.rep



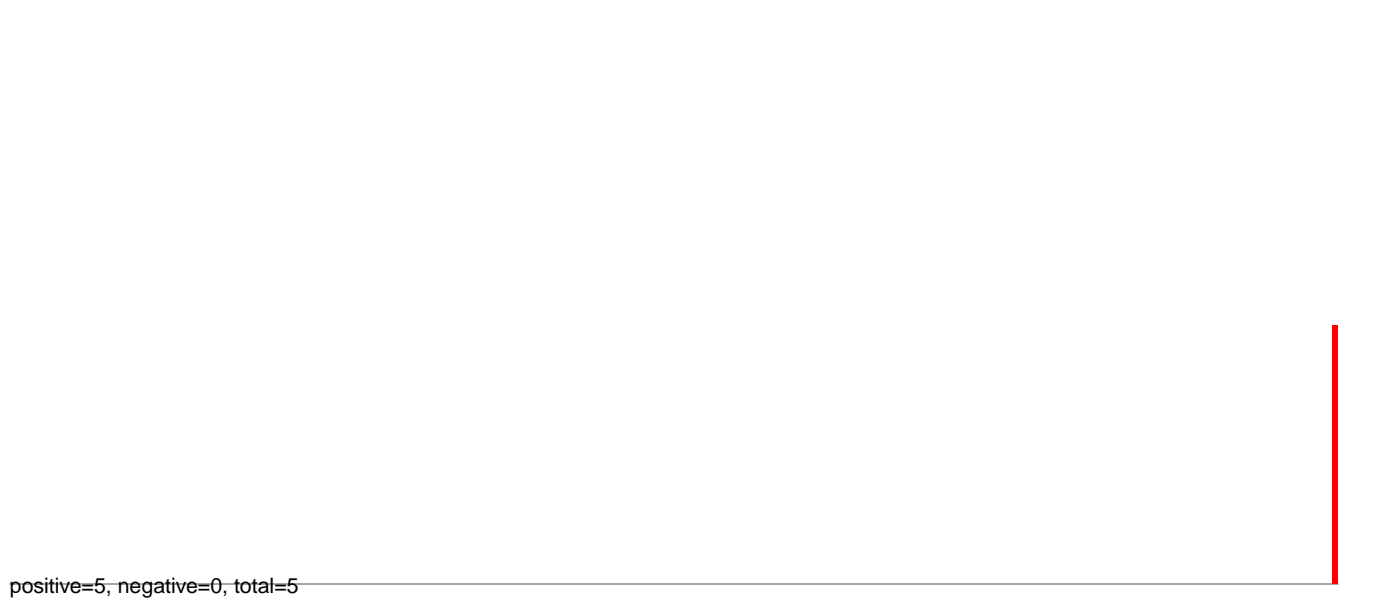
AeAeg_Ovary_GH.rep



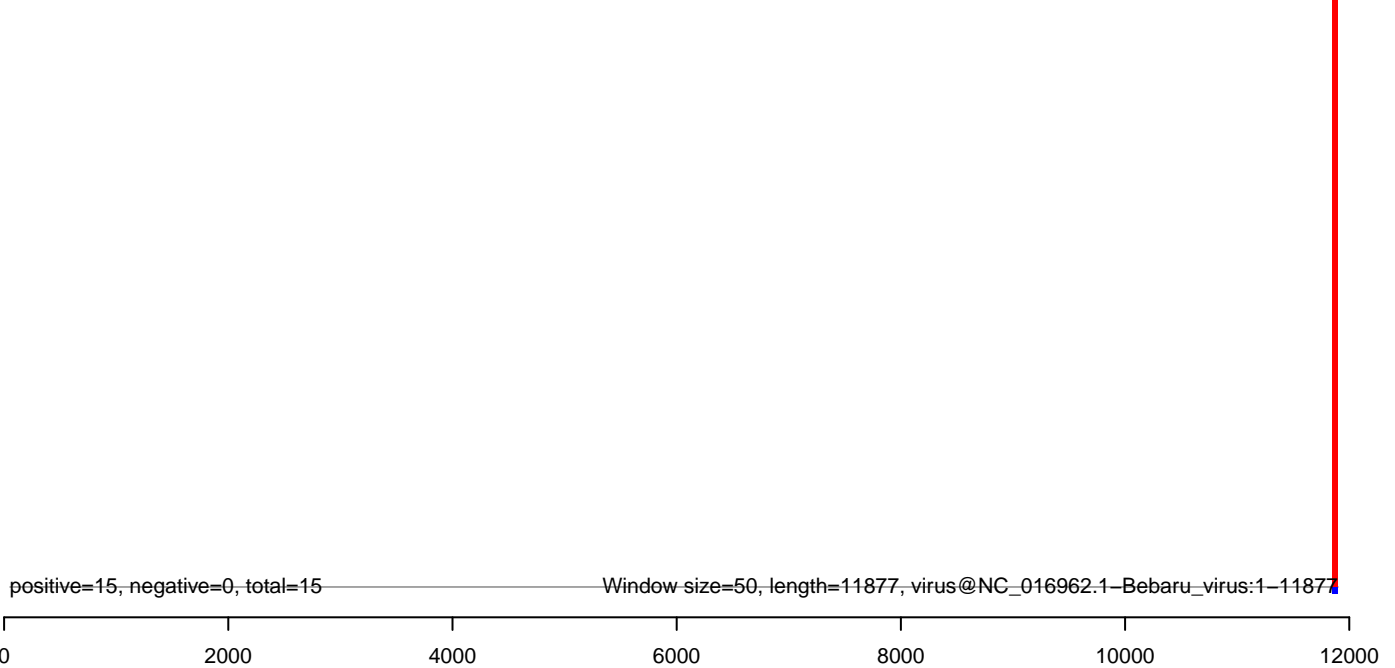
AeAeg_Ovary_GH.18_23.rep



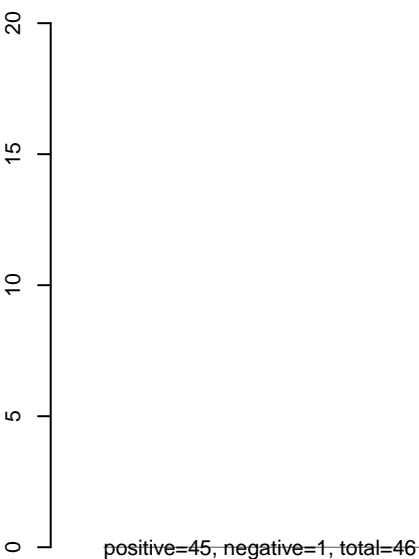
AeAeg_Ovary_GH.24_35.rep



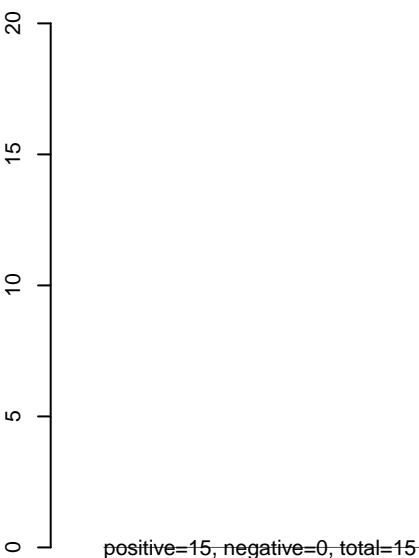
AeAeg_Ovary_GH.rep



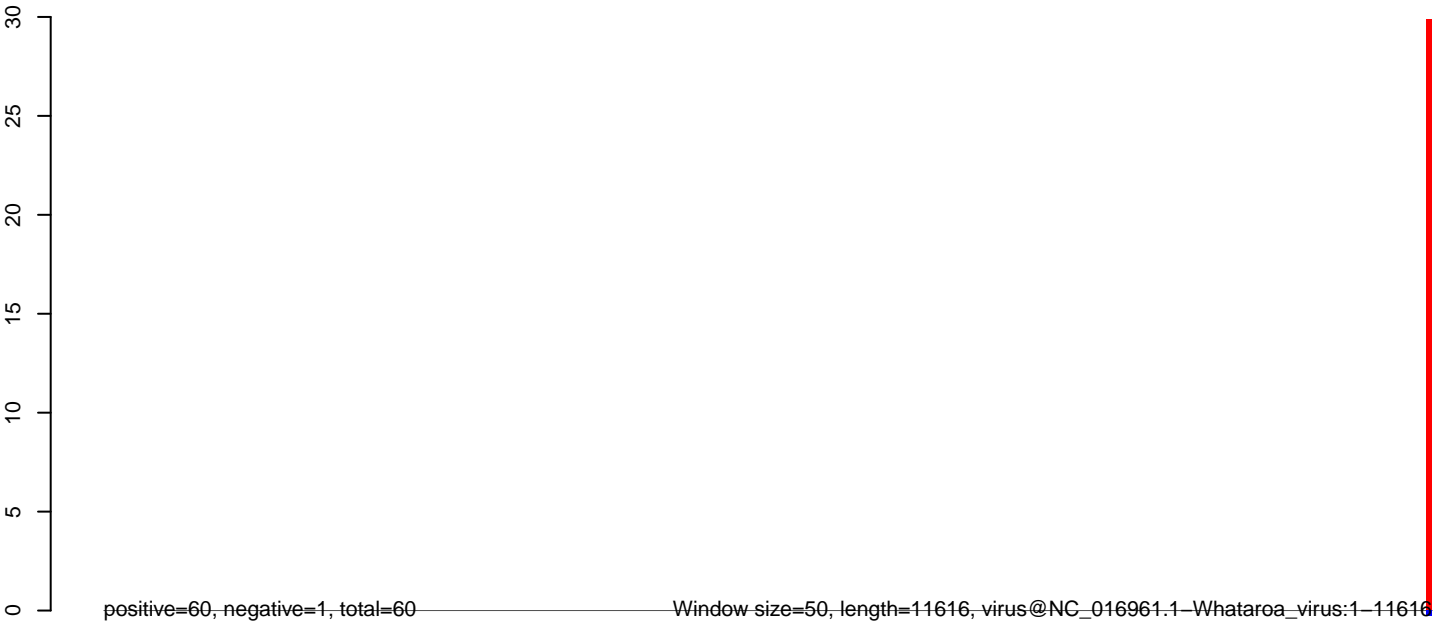
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

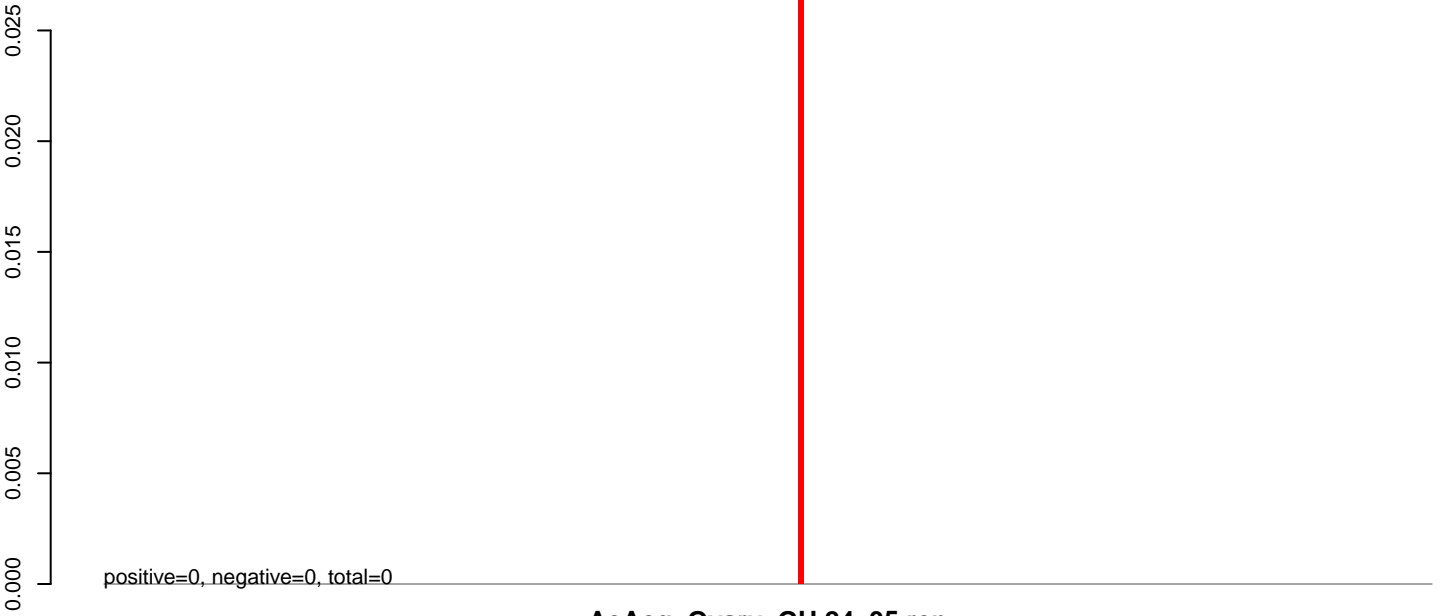


AeAeg_Ovary_GH.rep

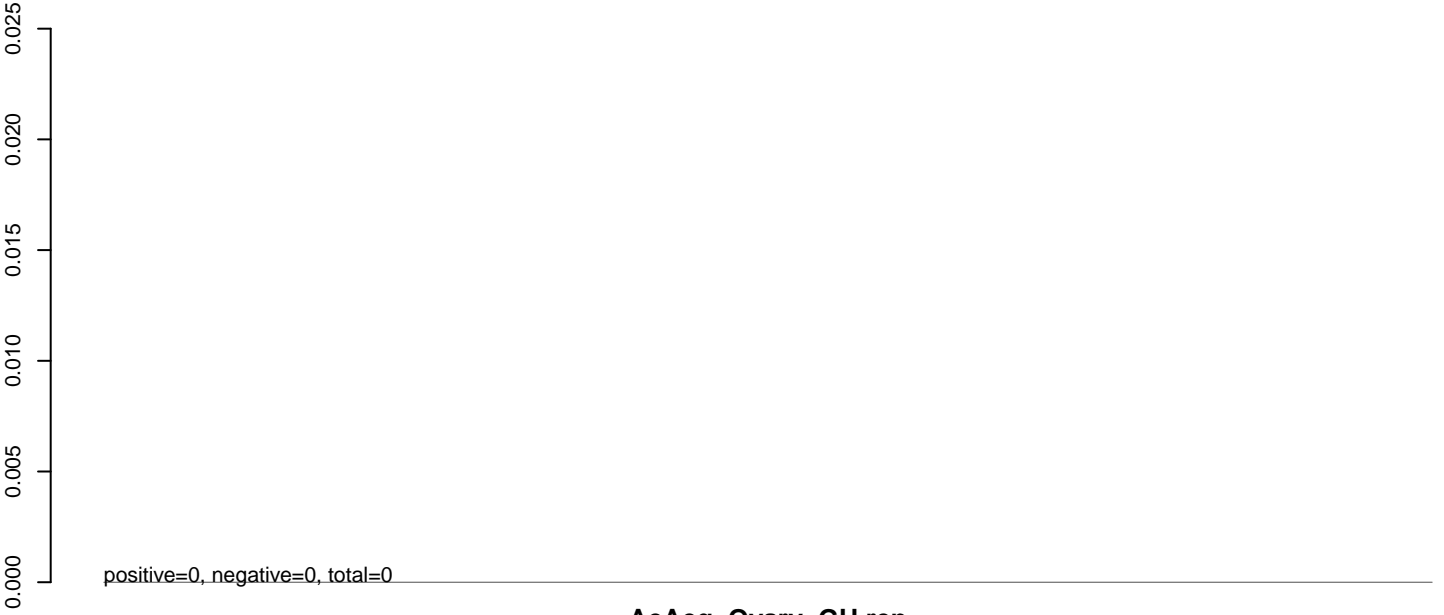


0 2000 4000 6000 8000 10000 12000

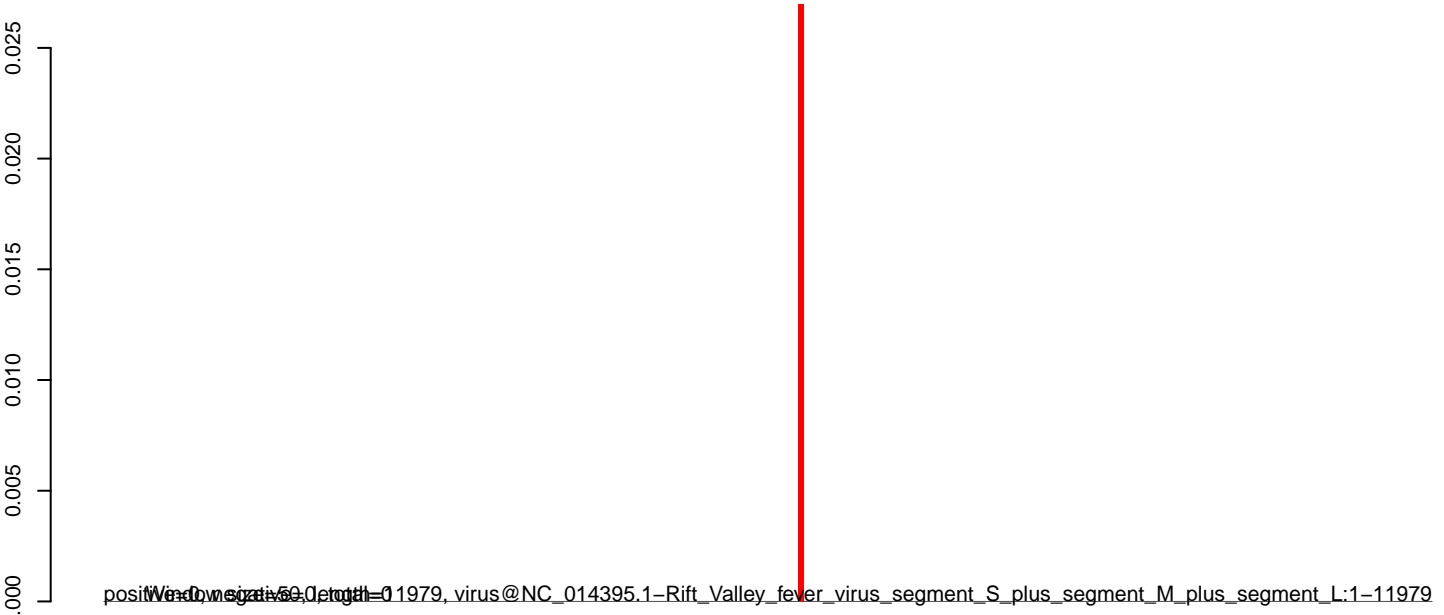
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

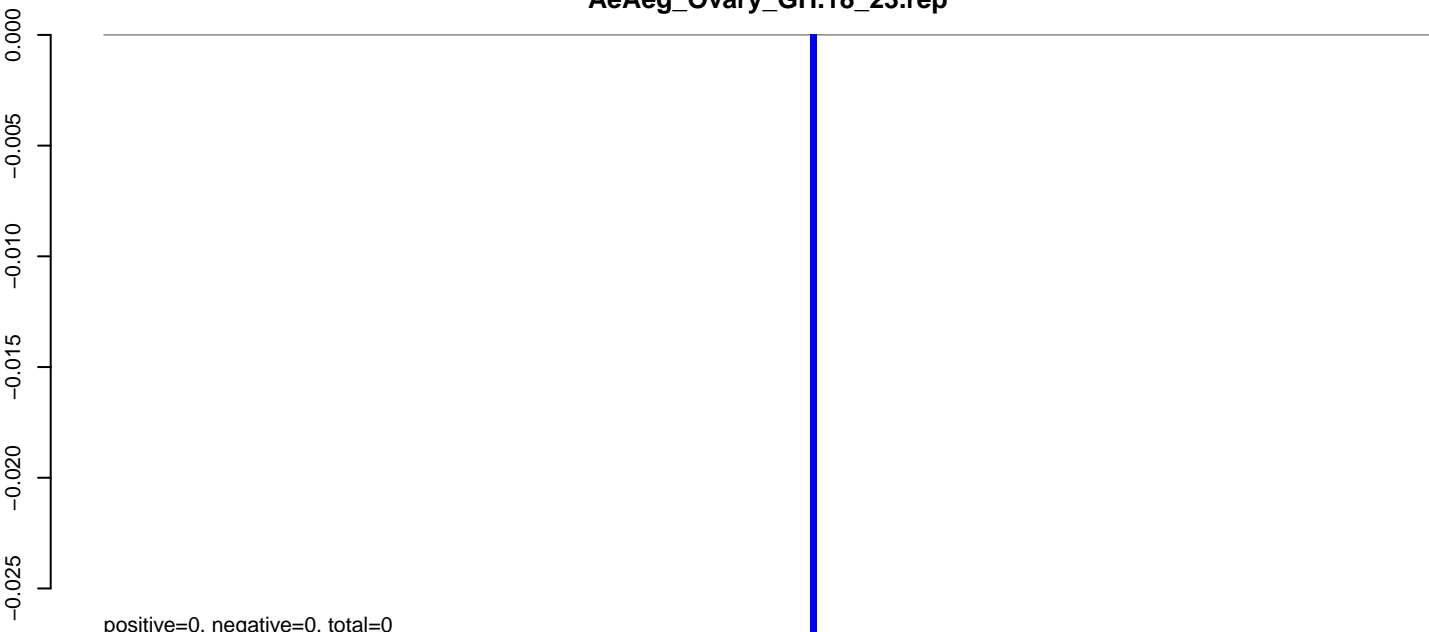


AeAeg_Ovary_GH.rep

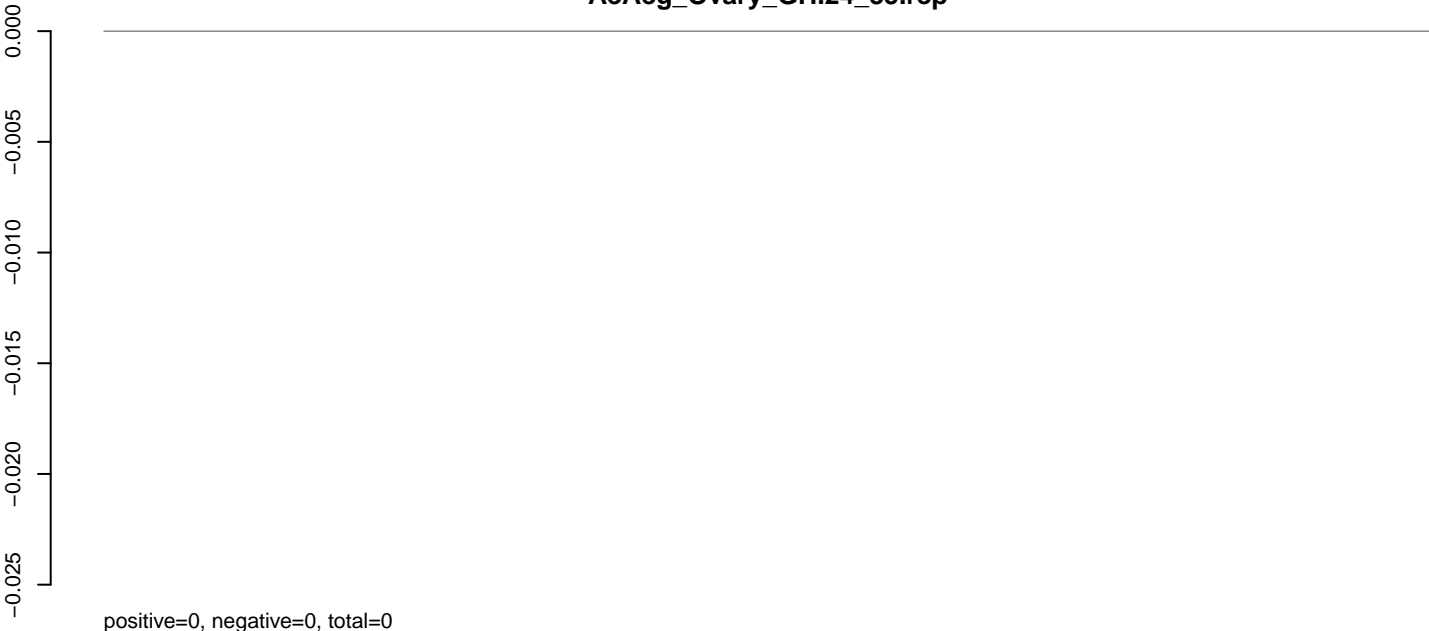


Wendong Zhang, et al. 2019, virus@NC_014395.1-Rift_Valley_fever_virus_segment_S_plus_segment_M_plus_segment_L:1-11979

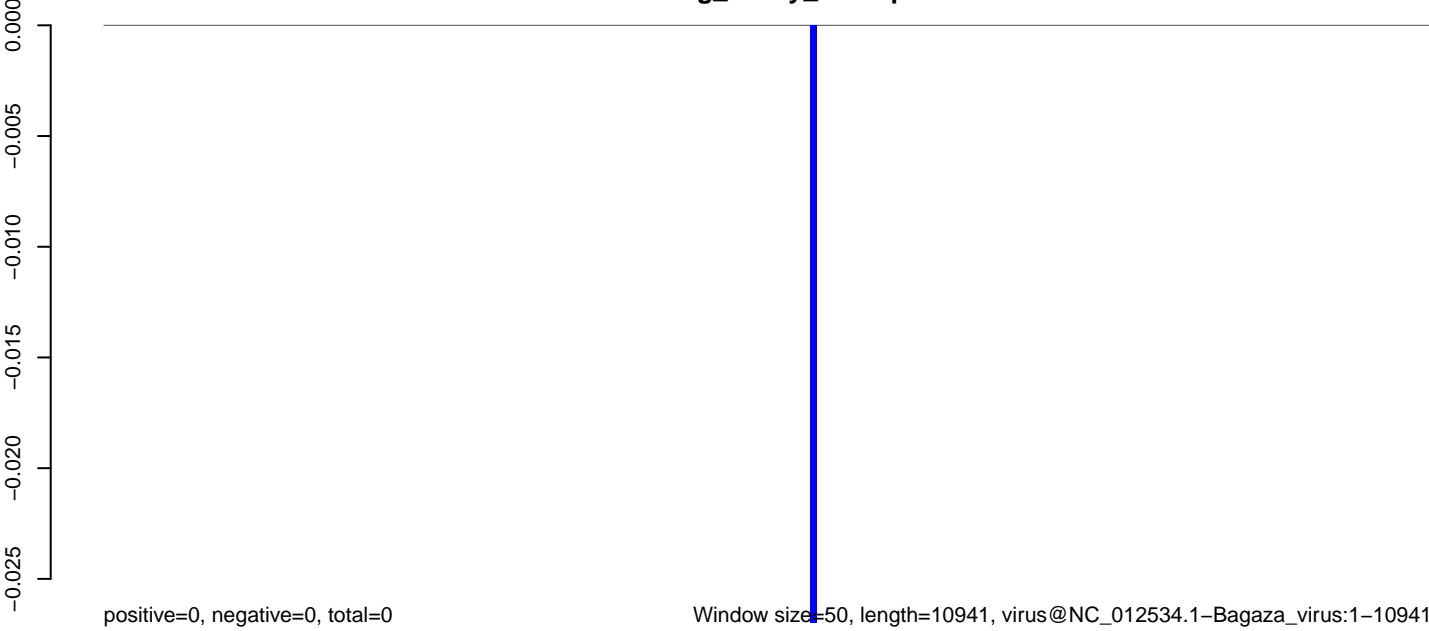
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

AeAeg_Ovary_GH.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AeAeg_Ovary_GH.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

AeAeg_Ovary_GH.rep

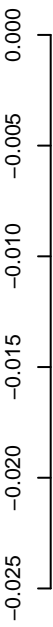
0.00
-0.02
-0.04
-0.06
-0.08
-0.10

positive=0, negative=0, total=0

Window size=50, length=10755, virus@NC_009028.2-Ilheus_virus:1-10755

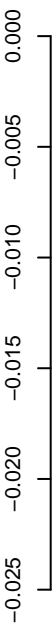
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AeAeg_Ovary_GH.18_23.rep



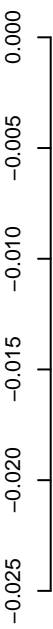
positive=0, negative=0, total=0

AeAeg_Ovary_GH.24_35.rep



positive=0, negative=0, total=0

AeAeg_Ovary_GH.rep

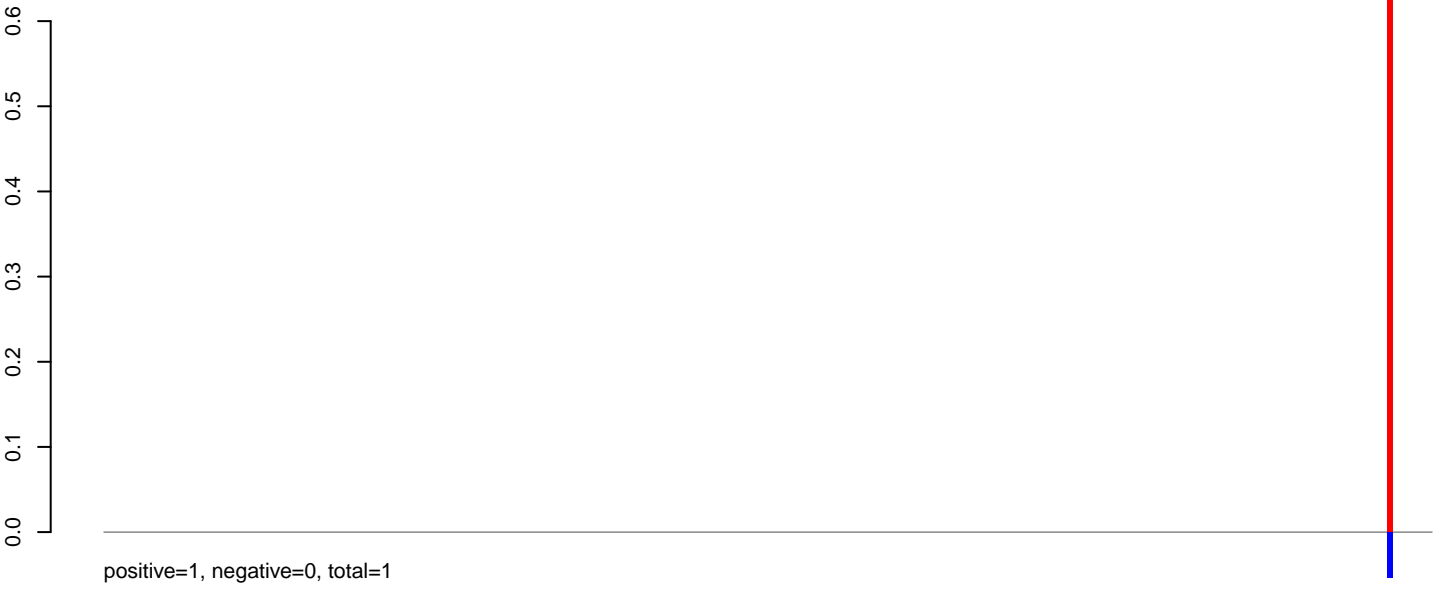


positive=0, negative=0, total=0

Window size=50, length=11066, virus@NC_006551.1-Uсутu_virus:1-11066



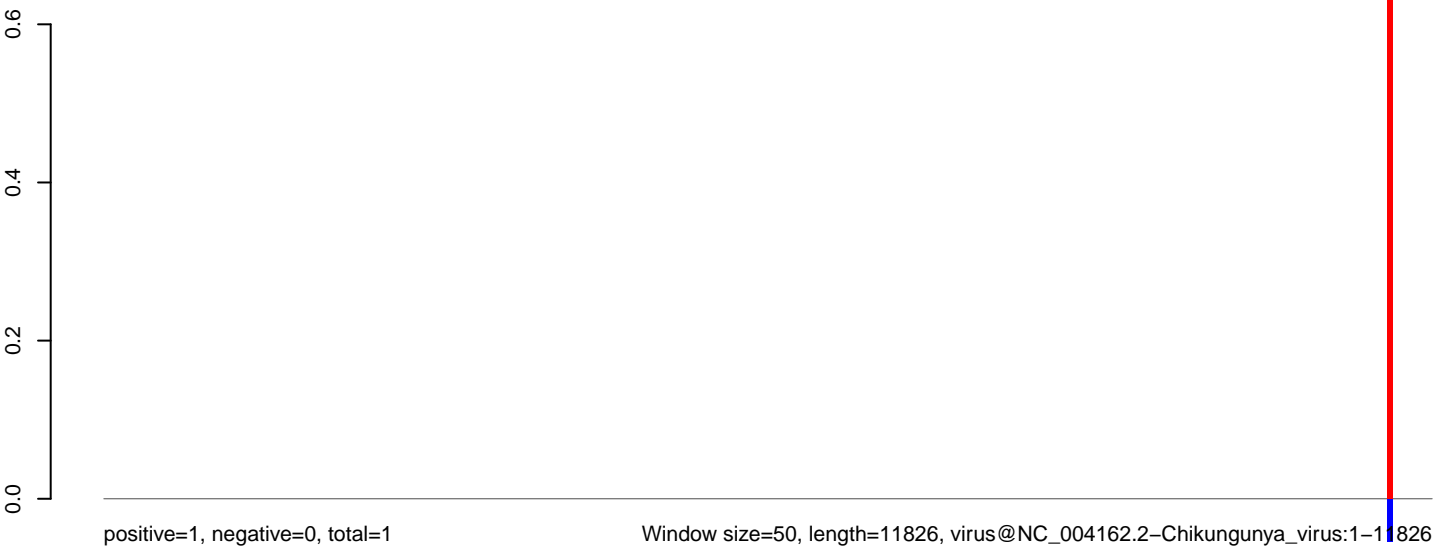
AeAeg_Ovary_GH.18_23.rep



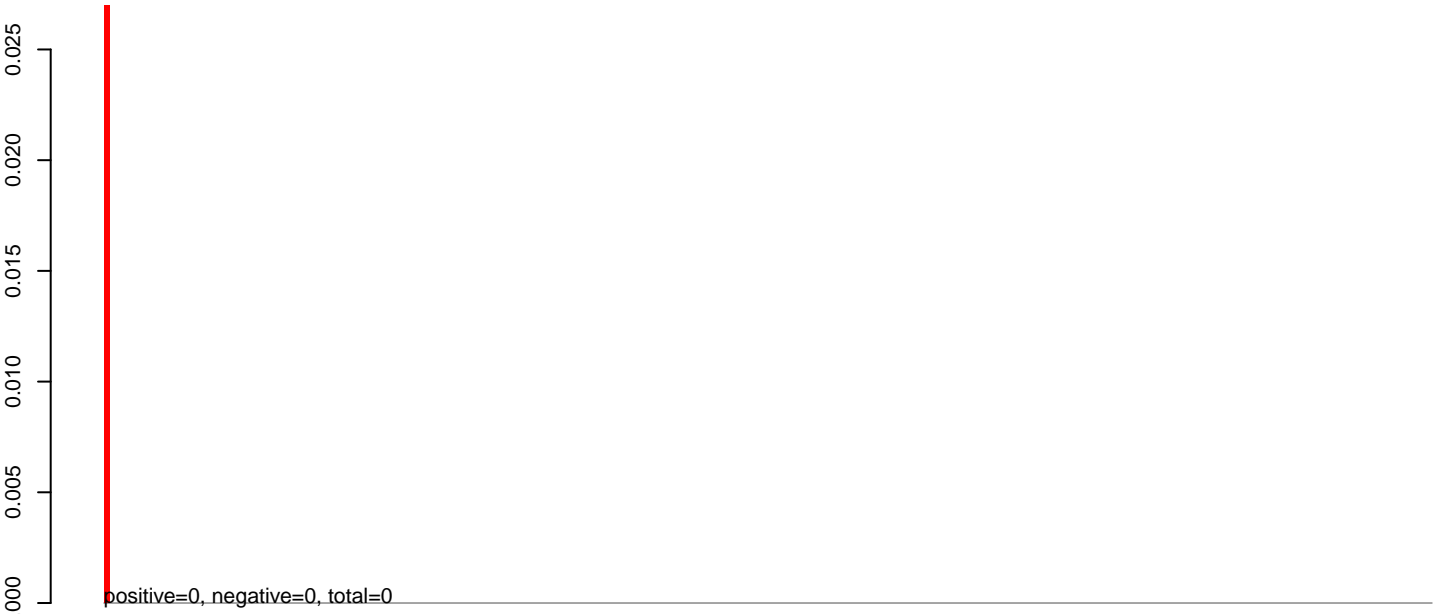
AeAeg_Ovary_GH.24_35.rep



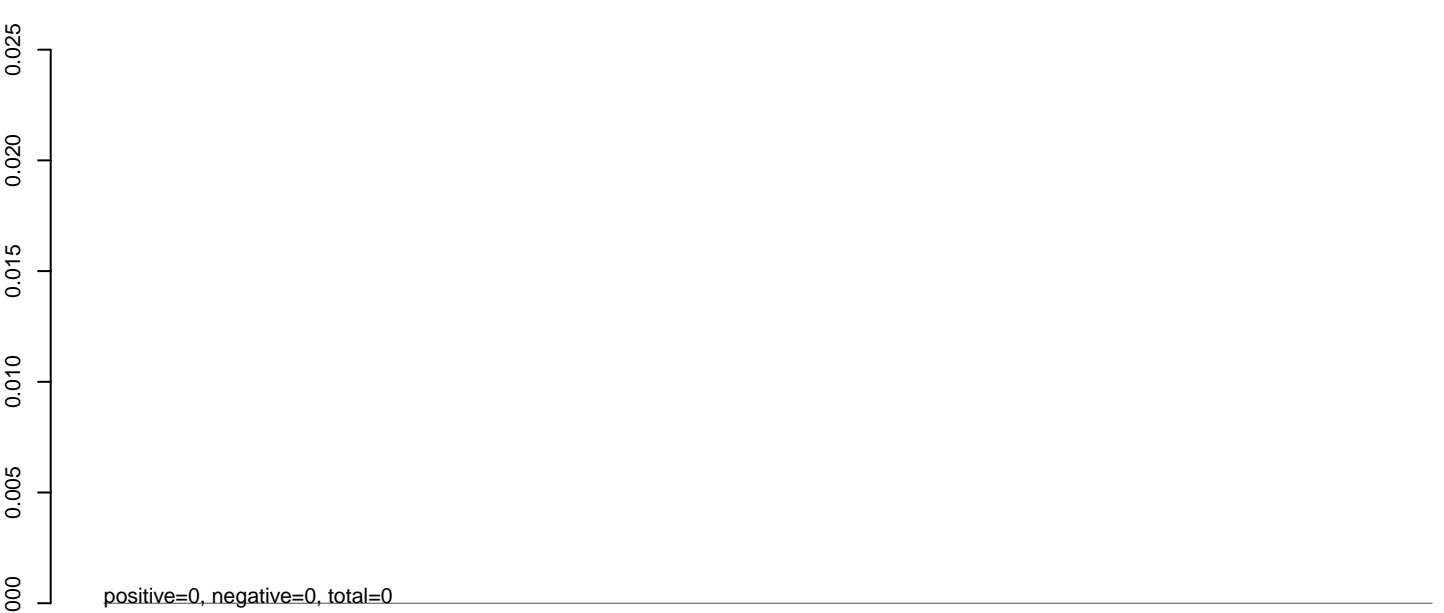
AeAeg_Ovary_GH.rep



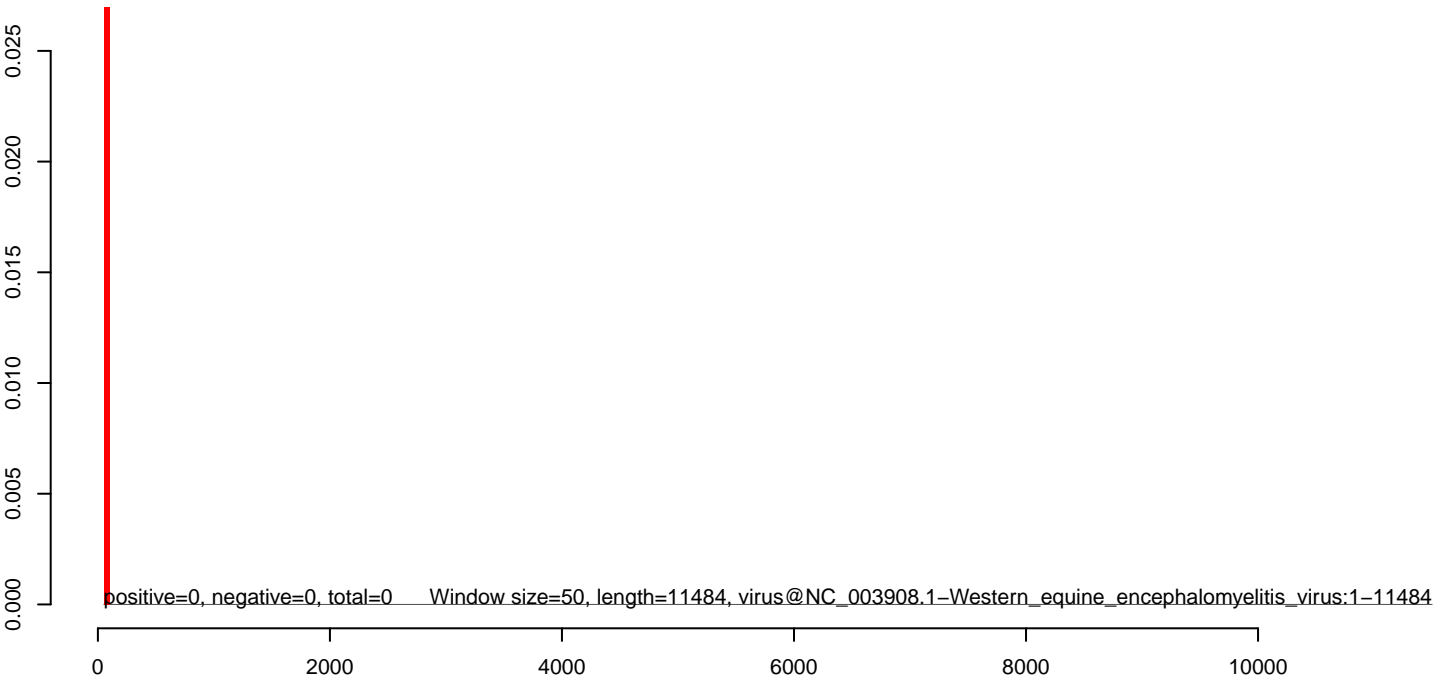
AeAeg_Ovary_GH.18_23.rep



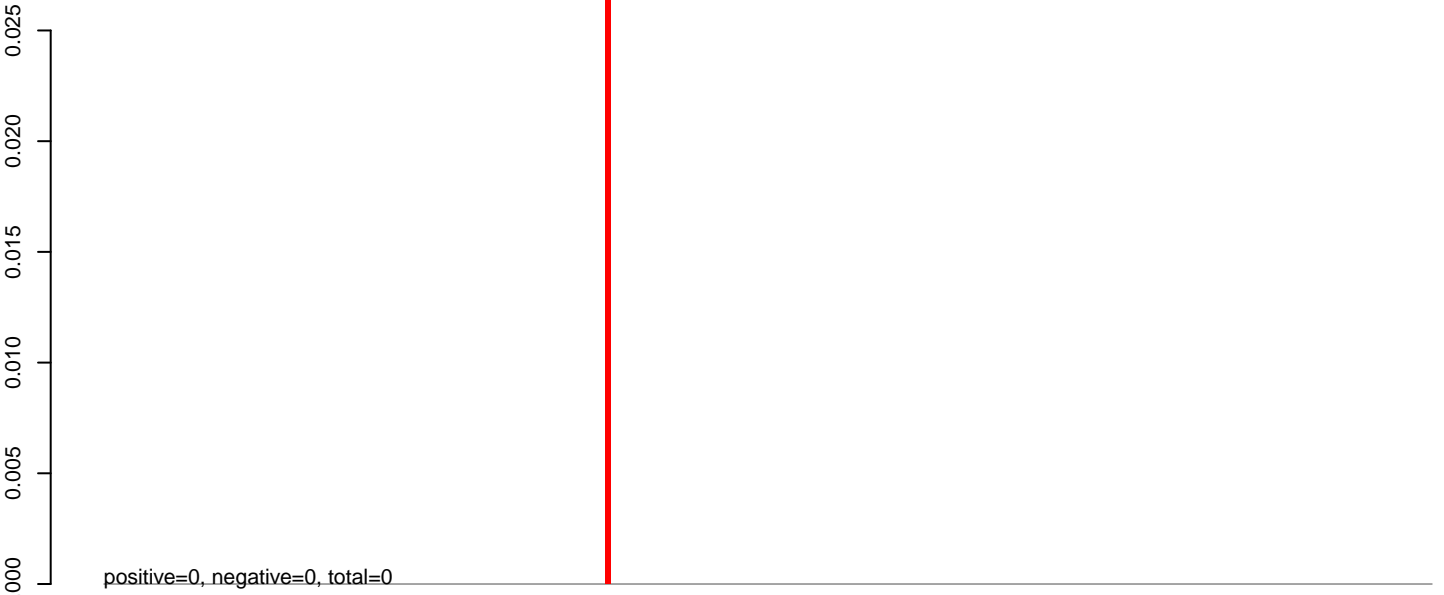
AeAeg_Ovary_GH.24_35.rep



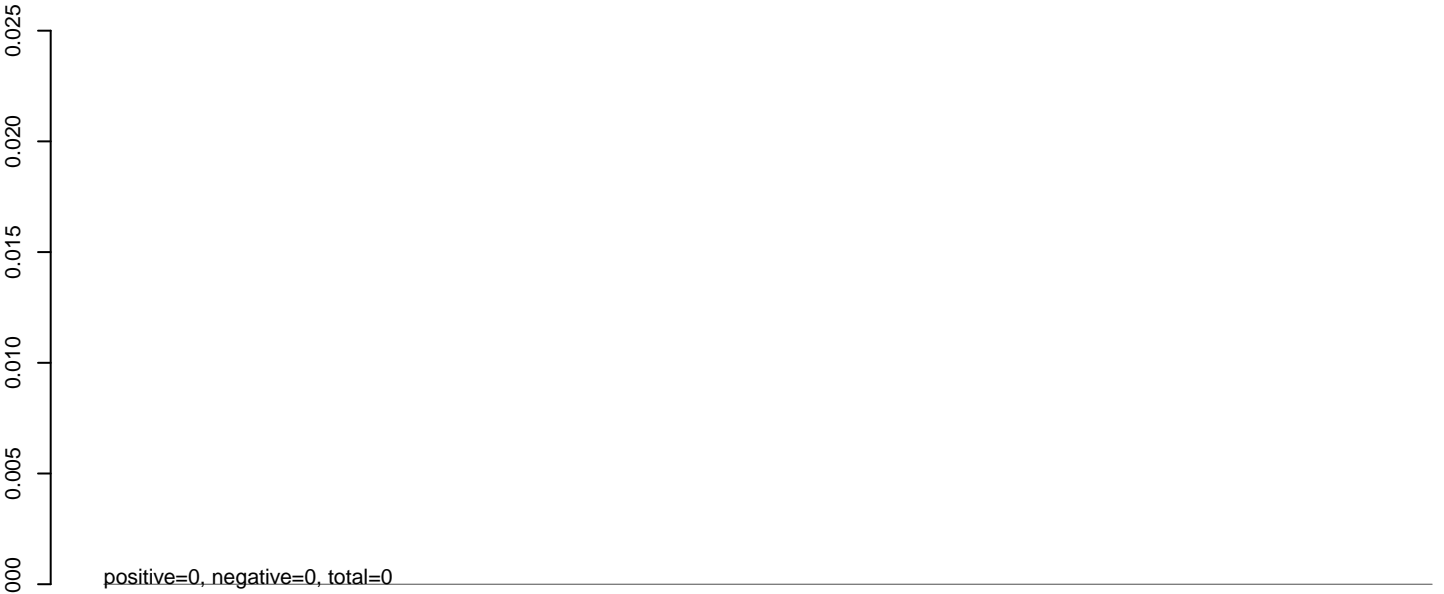
AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



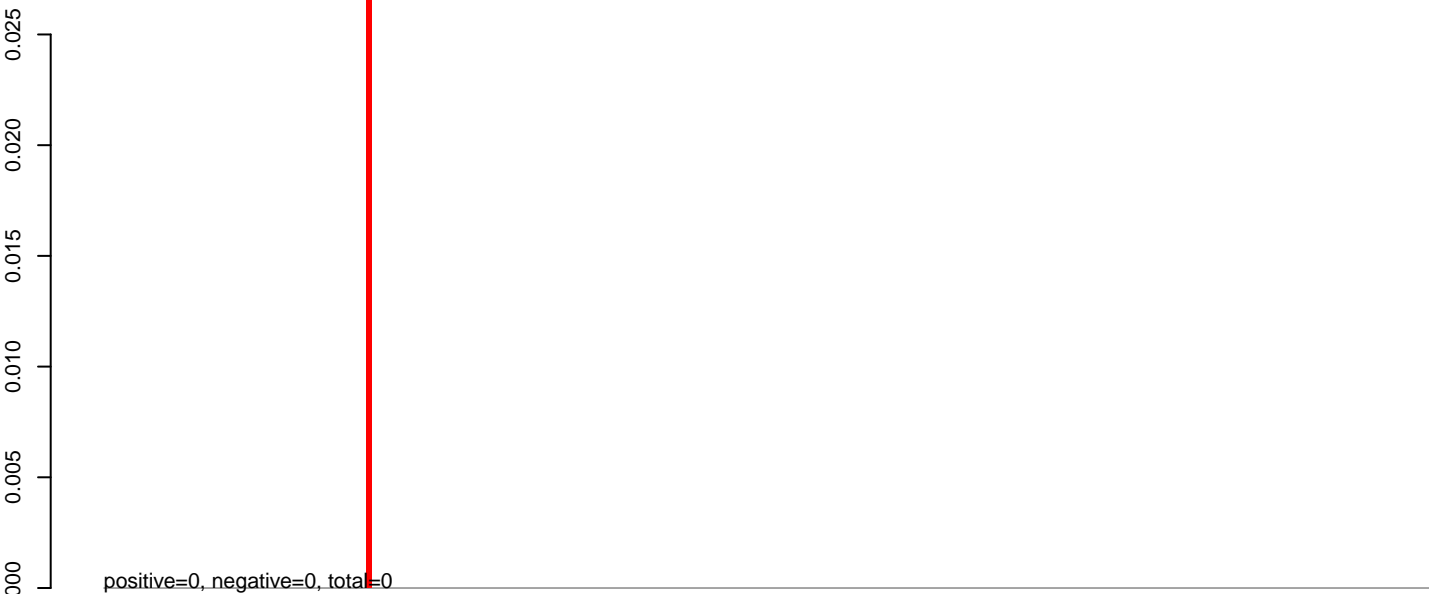
AeAeg_Ovary_GH.24_35.rep



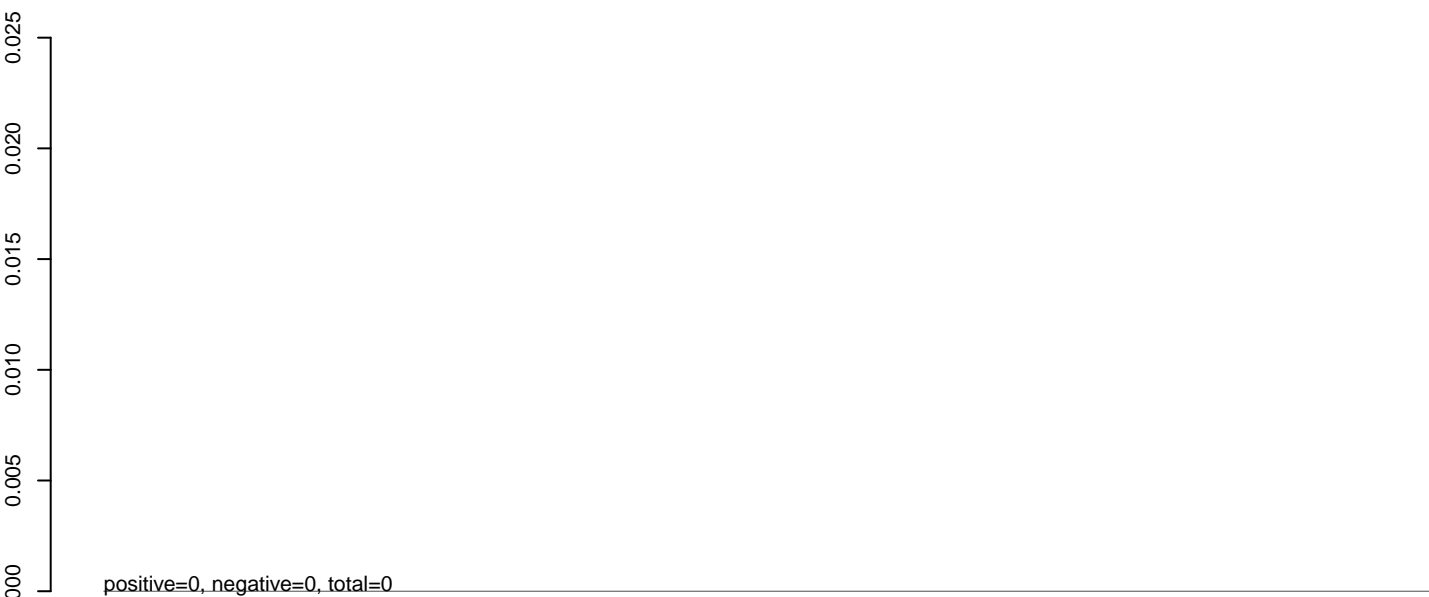
AeAeg_Ovary_GH.rep



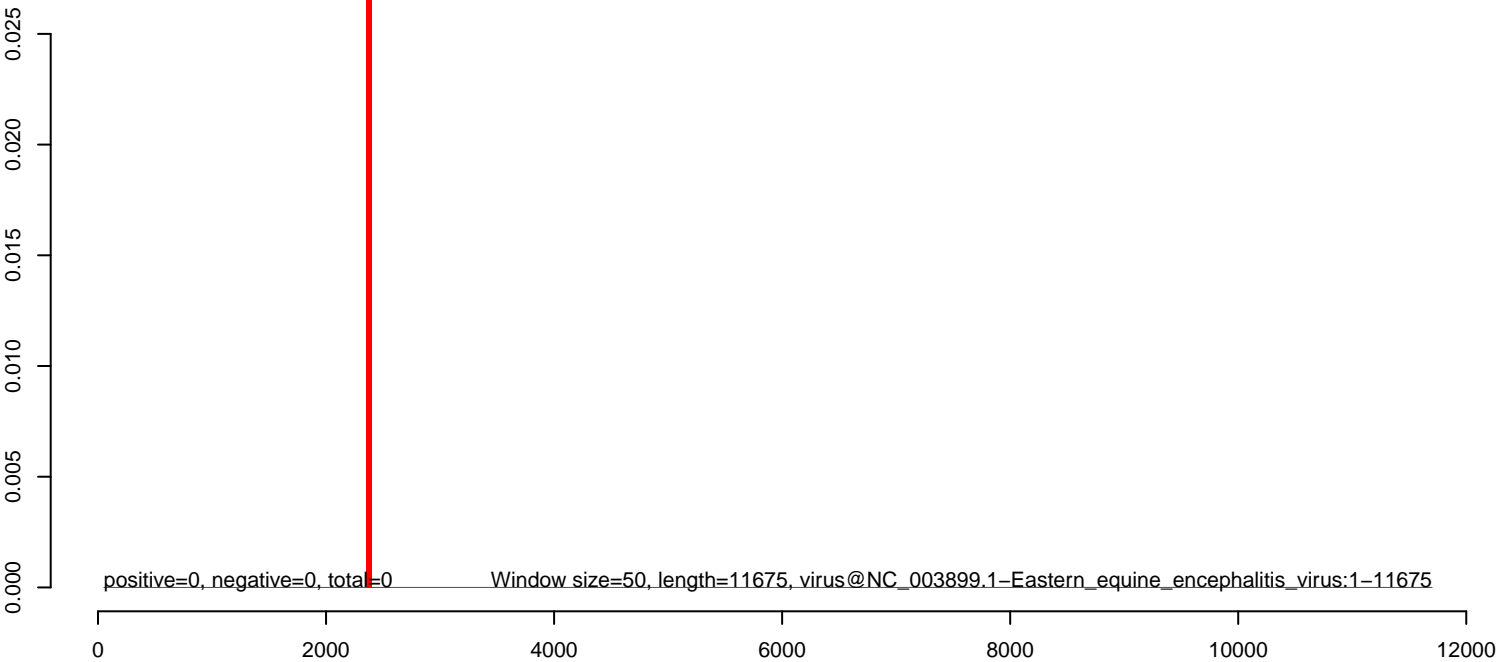
AeAeg_Ovary_GH.18_23.rep



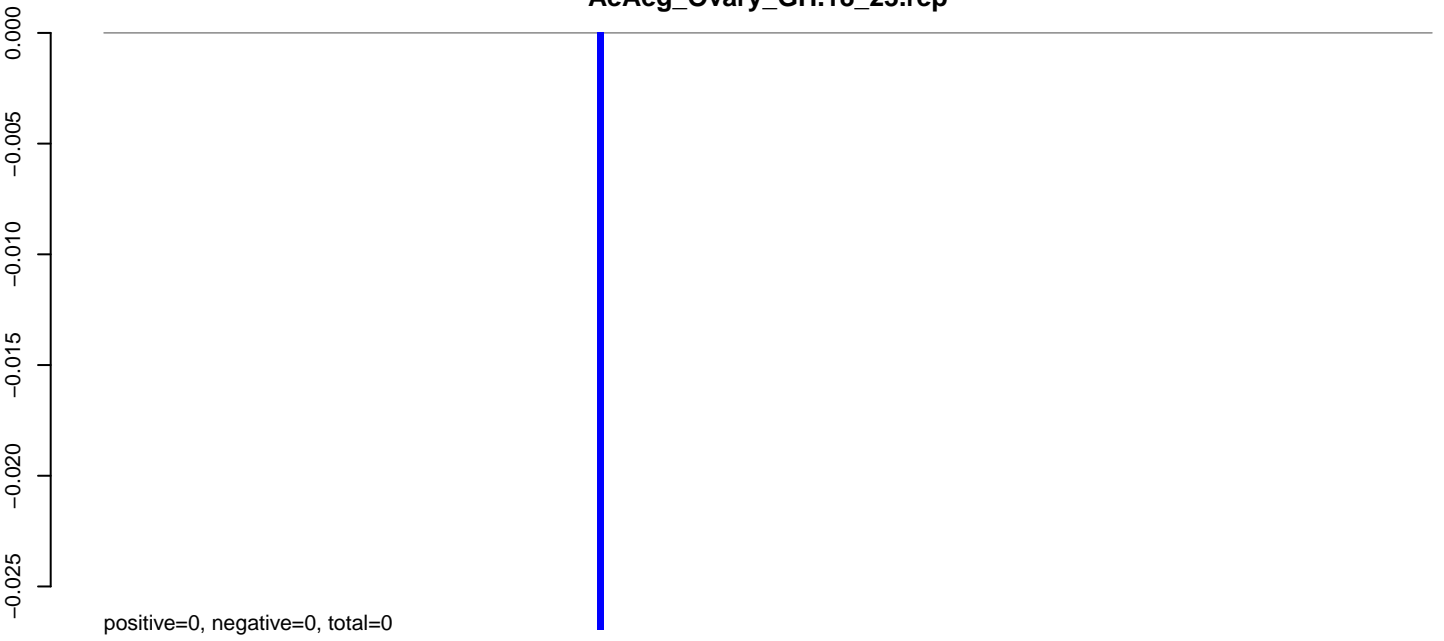
AeAeg_Ovary_GH.24_35.rep



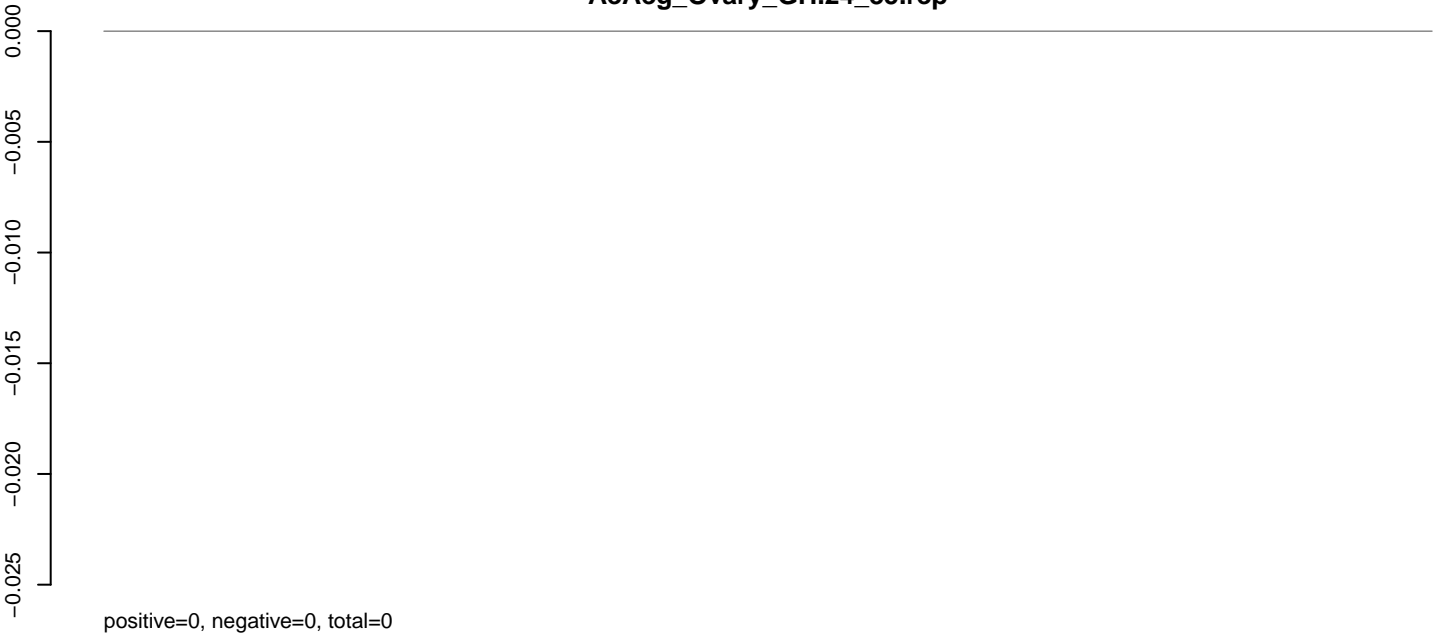
AeAeg_Ovary_GH.rep



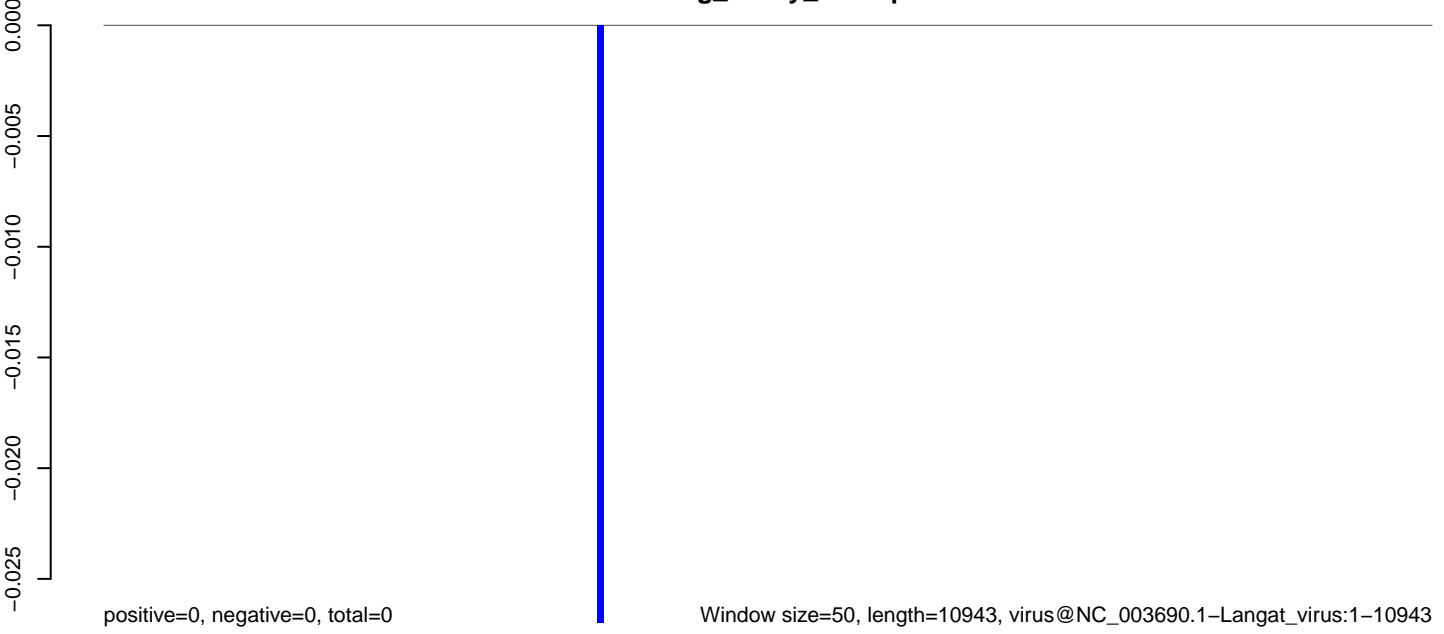
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

positive=0, negative=0, total=0

AeAeg_Ovary_GH.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

positive=0, negative=0, total=0

AeAeg_Ovary_GH.rep

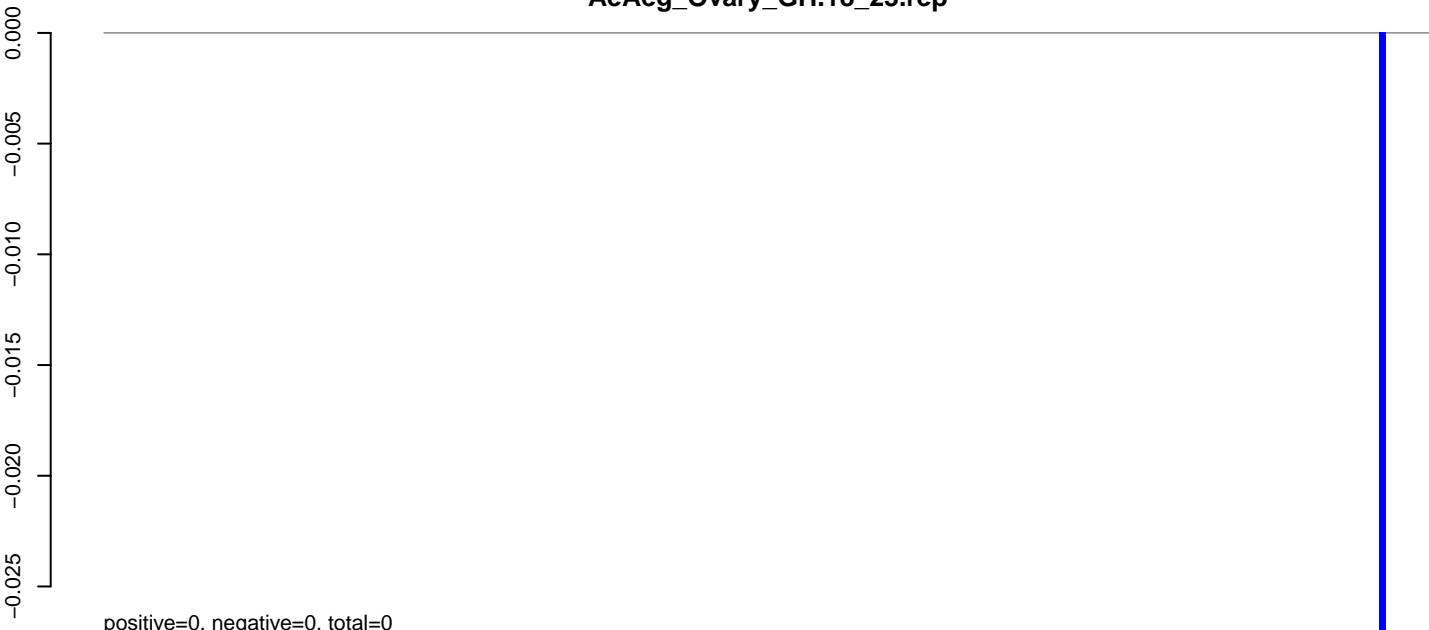
0.000
-0.005
-0.010
-0.015
-0.020
-0.025

positive=0, negative=0, total=0

Window size=50, length=10839, virus@NC_003687.1-Powassan_virus_1-10839

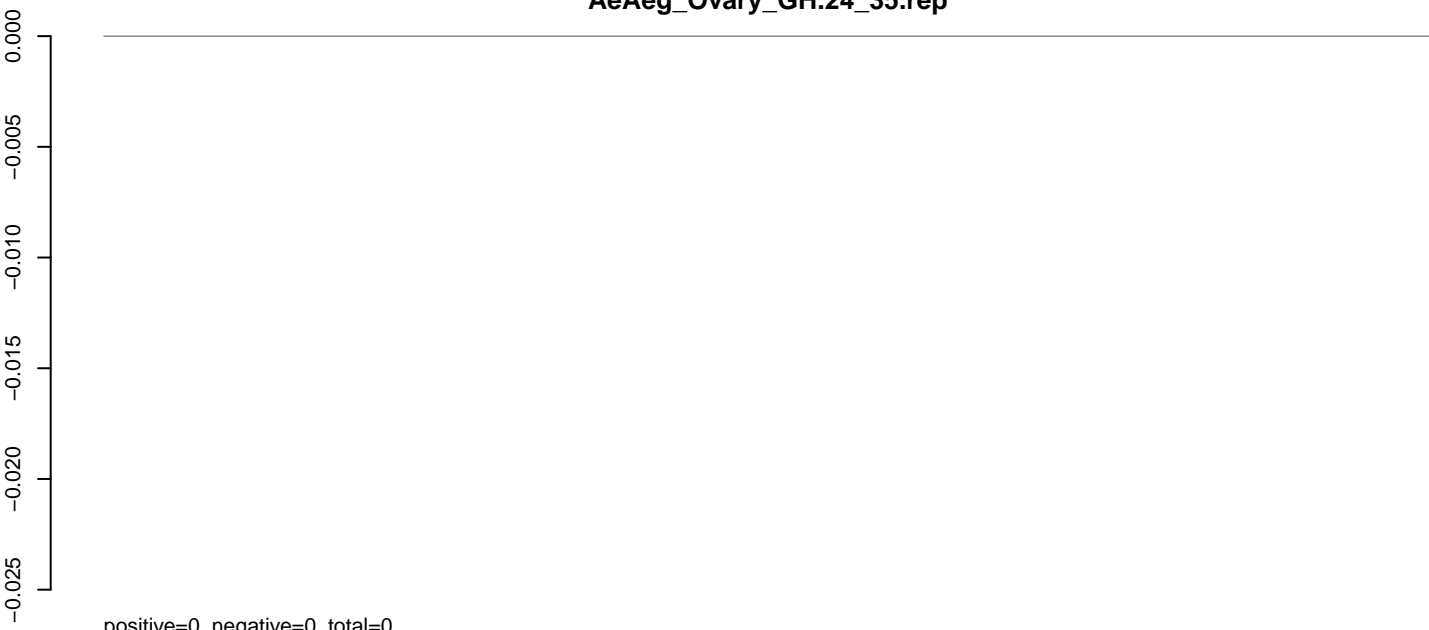
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AeAeg_Ovary_GH.18_23.rep



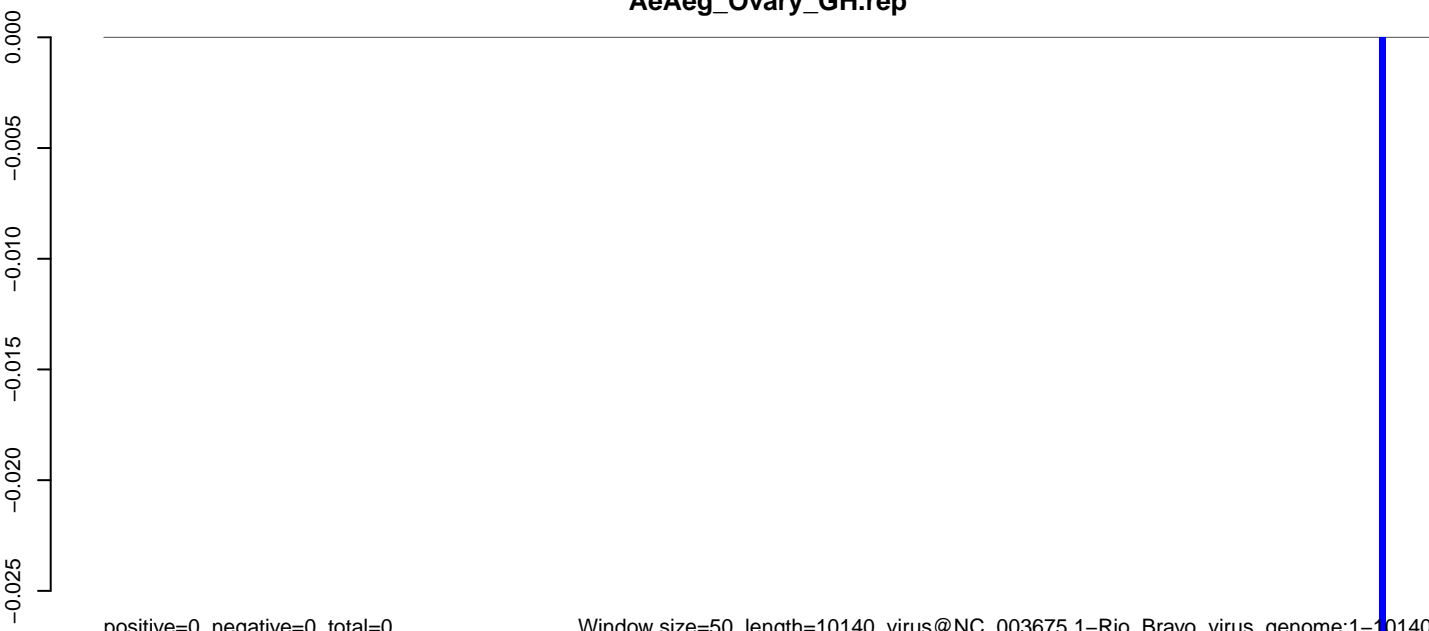
positive=0, negative=0, total=0

AeAeg_Ovary_GH.24_35.rep



positive=0, negative=0, total=0

AeAeg_Ovary_GH.rep

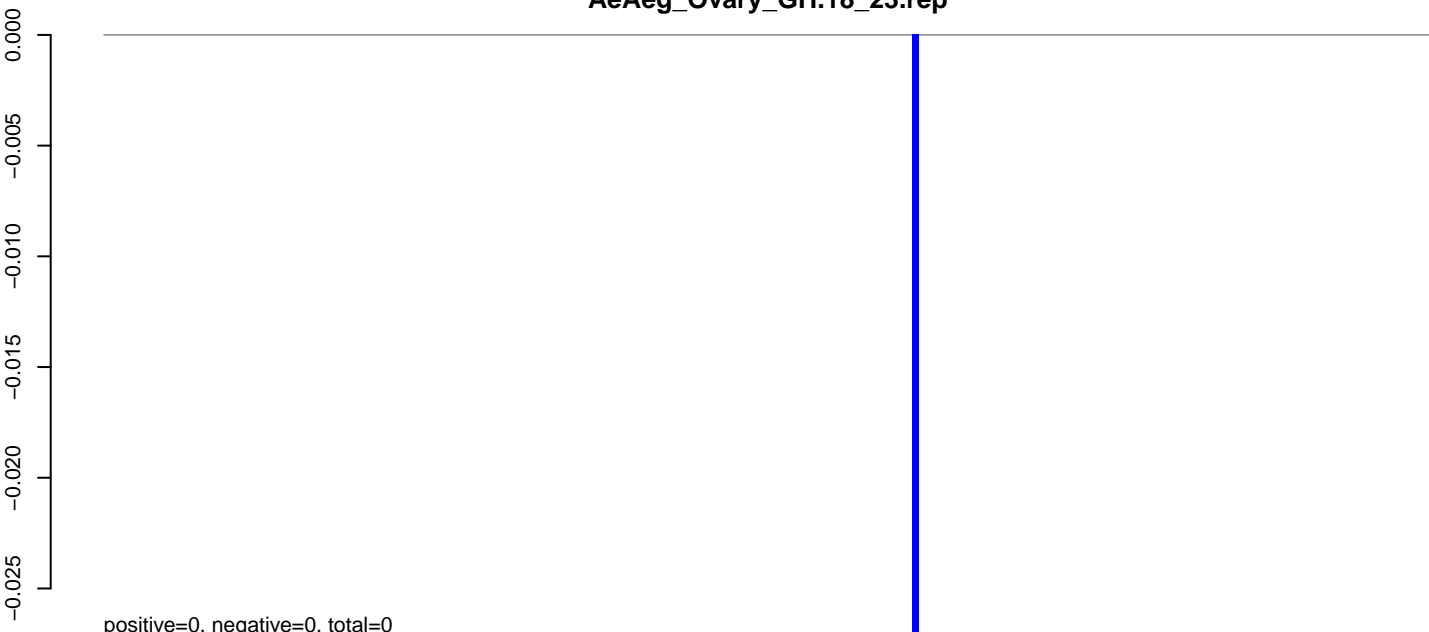


positive=0, negative=0, total=0

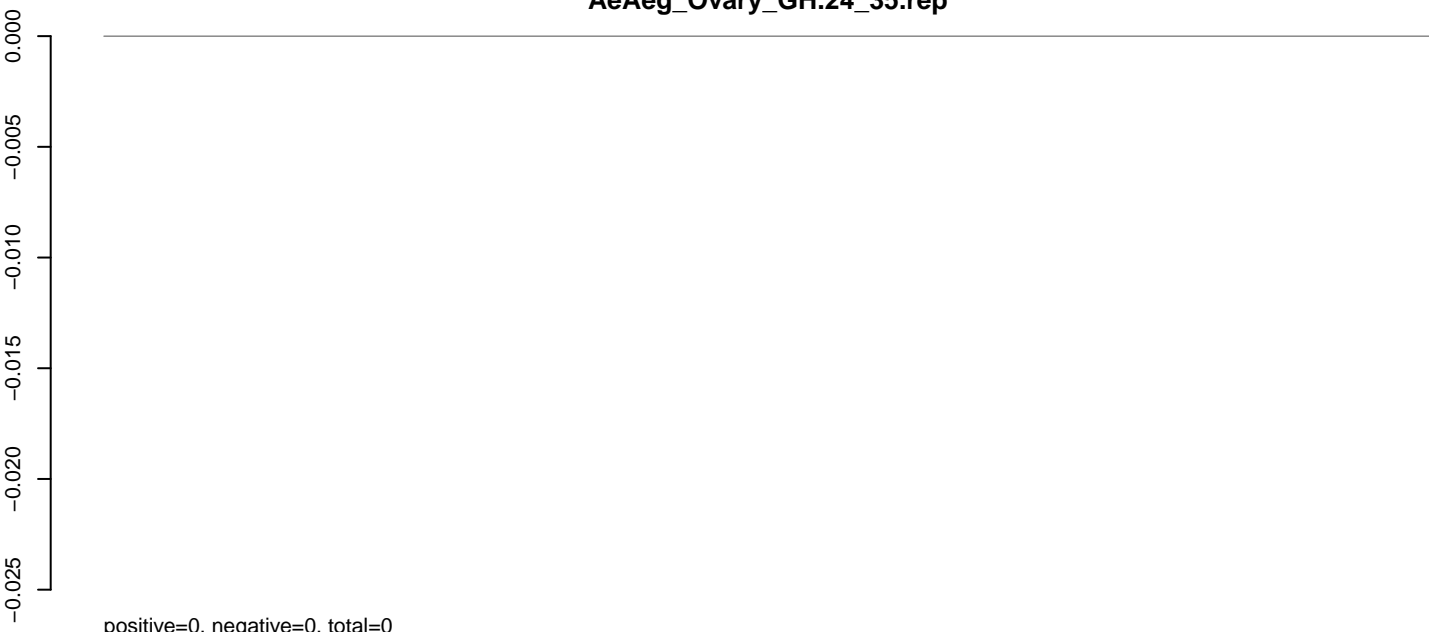
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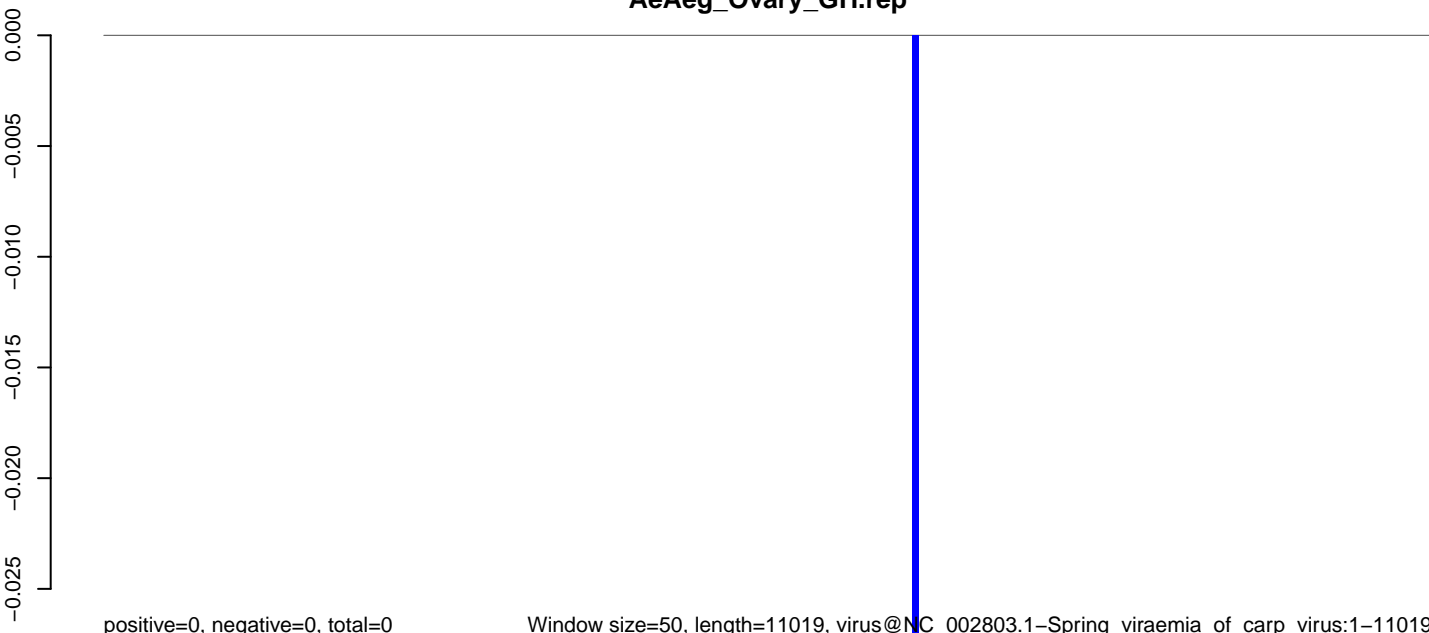
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



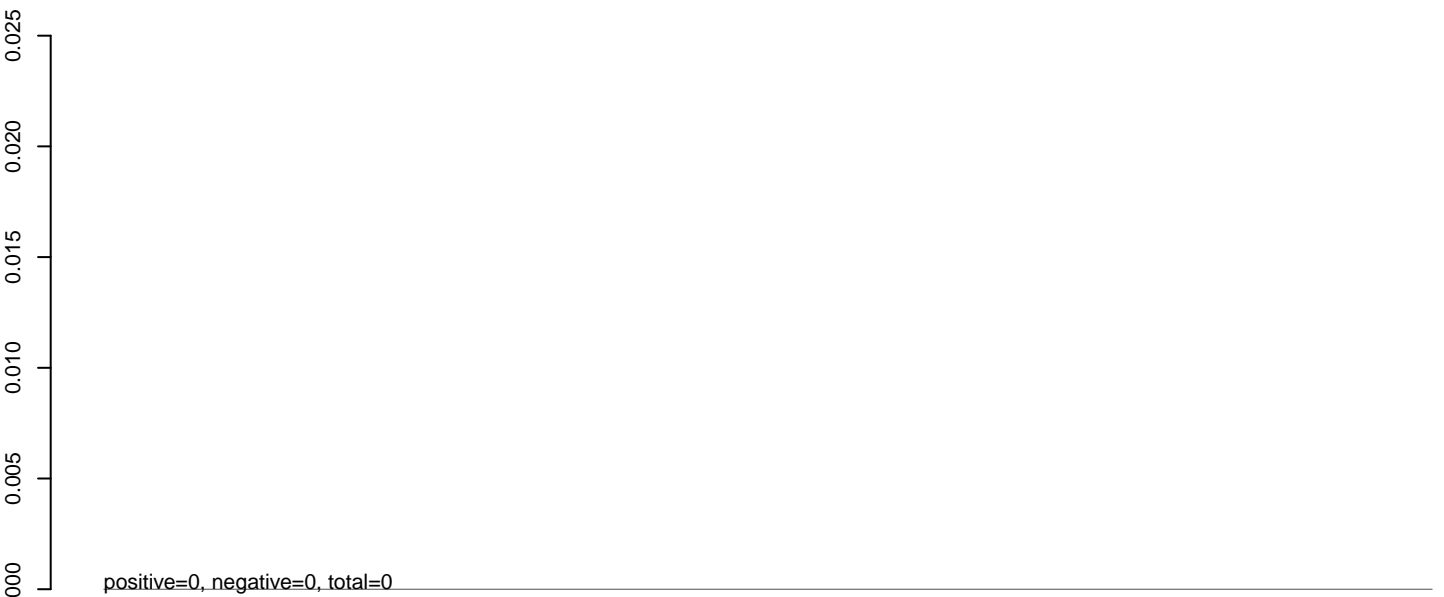
Window size=50, length=11019, virus@NC_002803.1-Spring_viraemia_of_carp_virus:1-11019

0 2000 4000 6000 8000 10000

AeAeg_Ovary_GH.18_23.rep



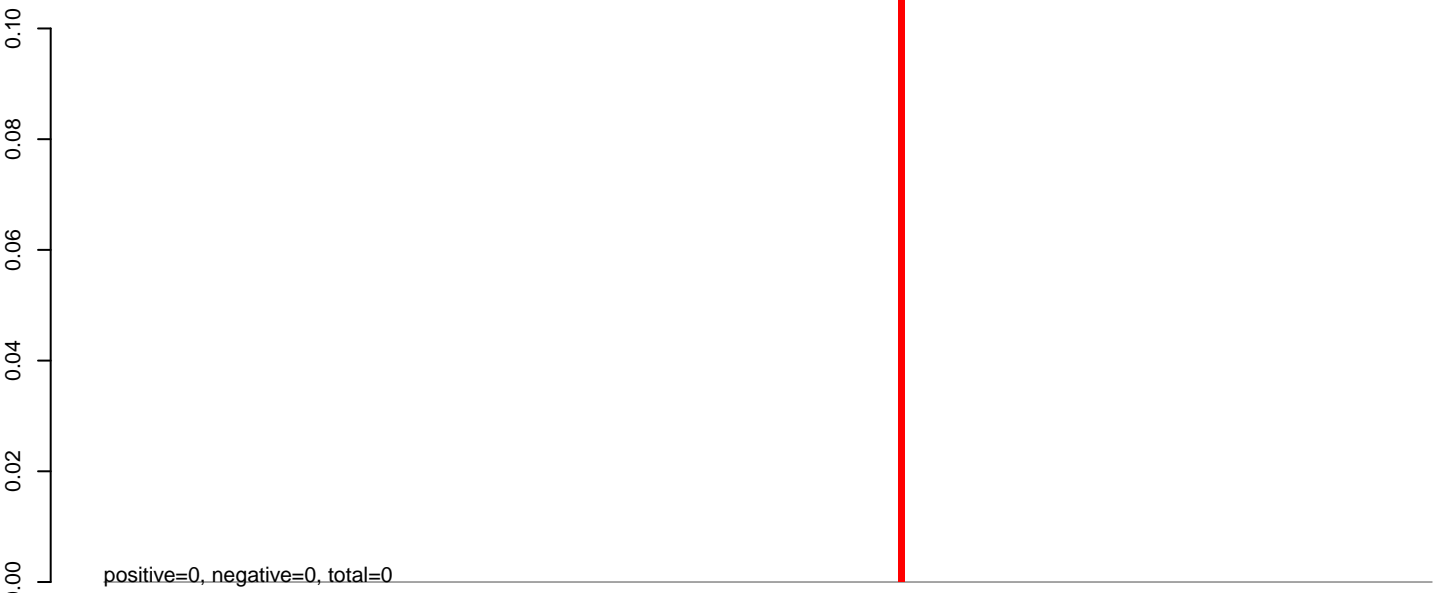
AeAeg_Ovary_GH.24_35.rep



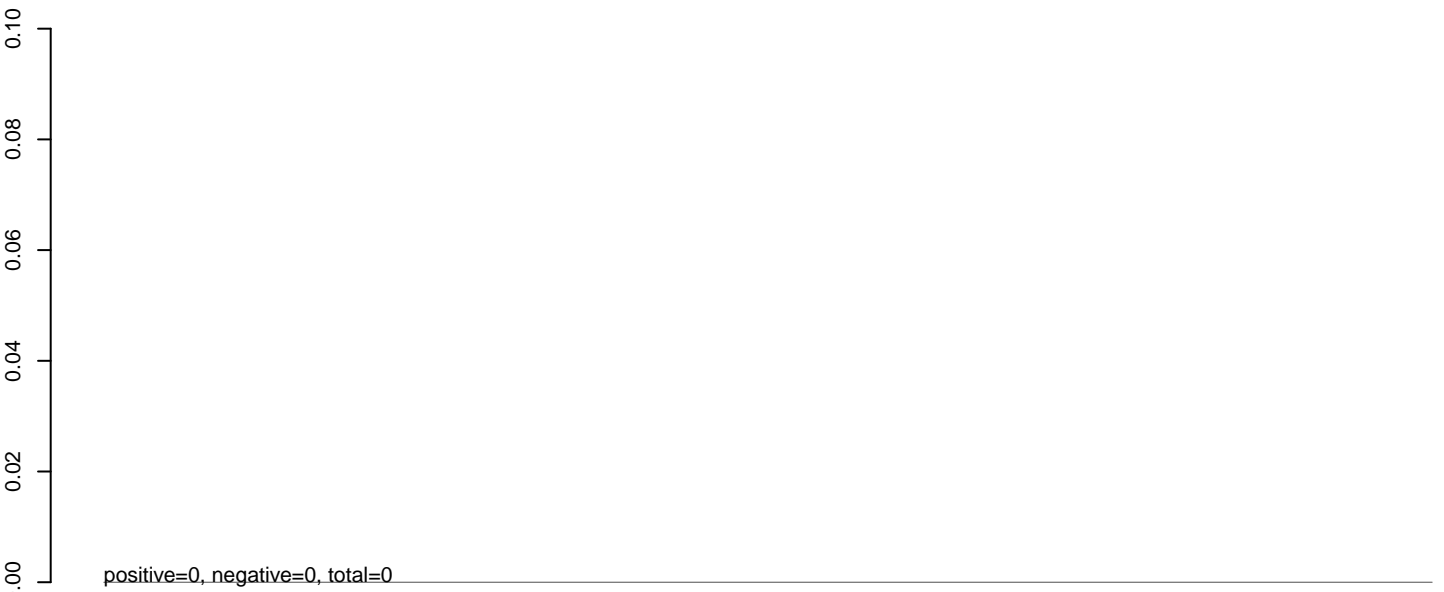
AeAeg_Ovary_GH.rep



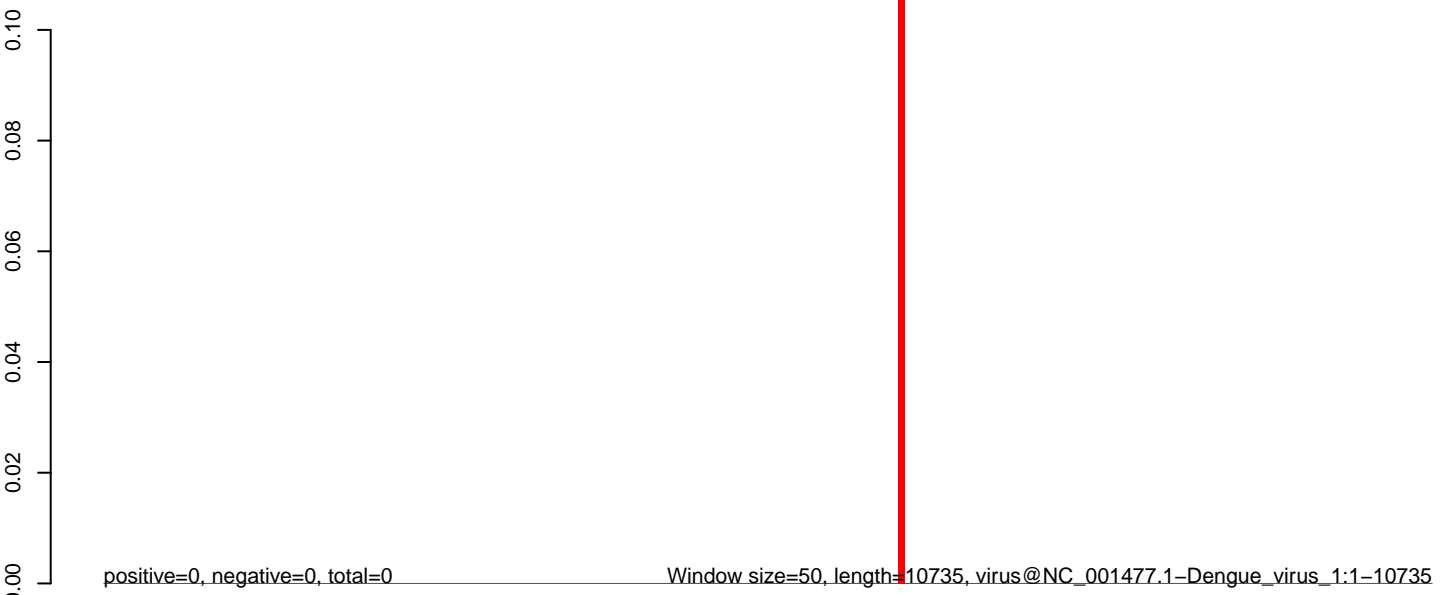
AeAeg_Ovary_GH.18_23.rep



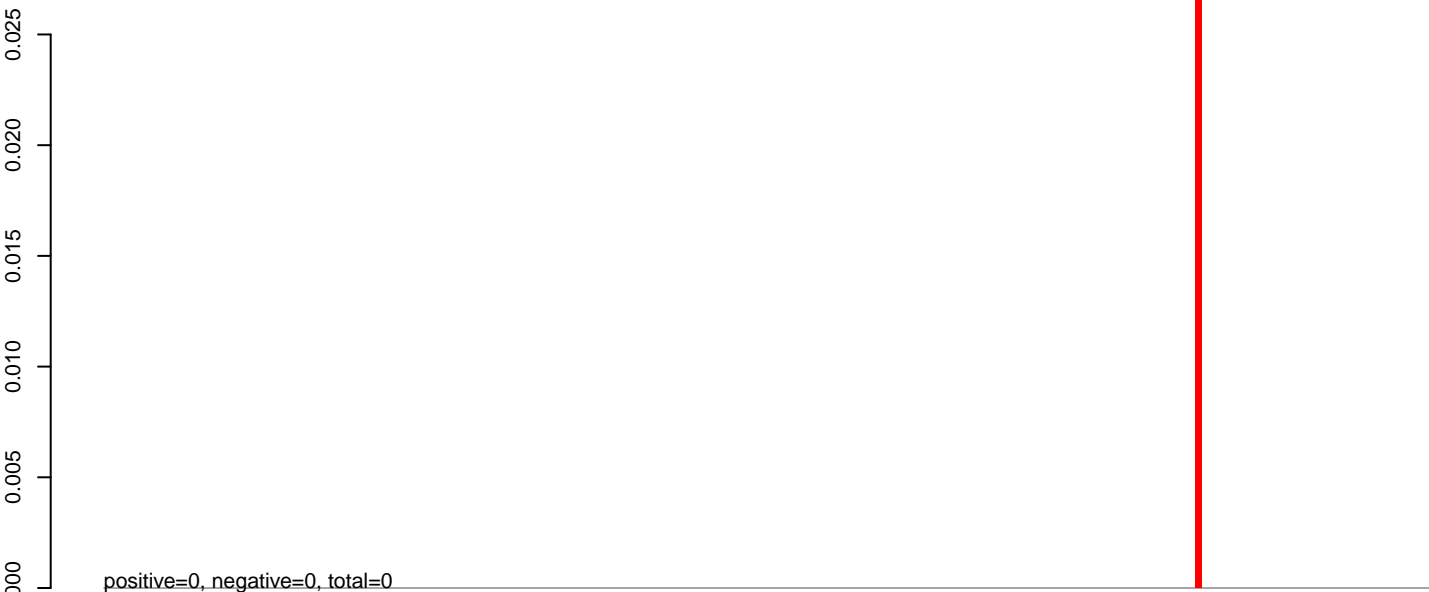
AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



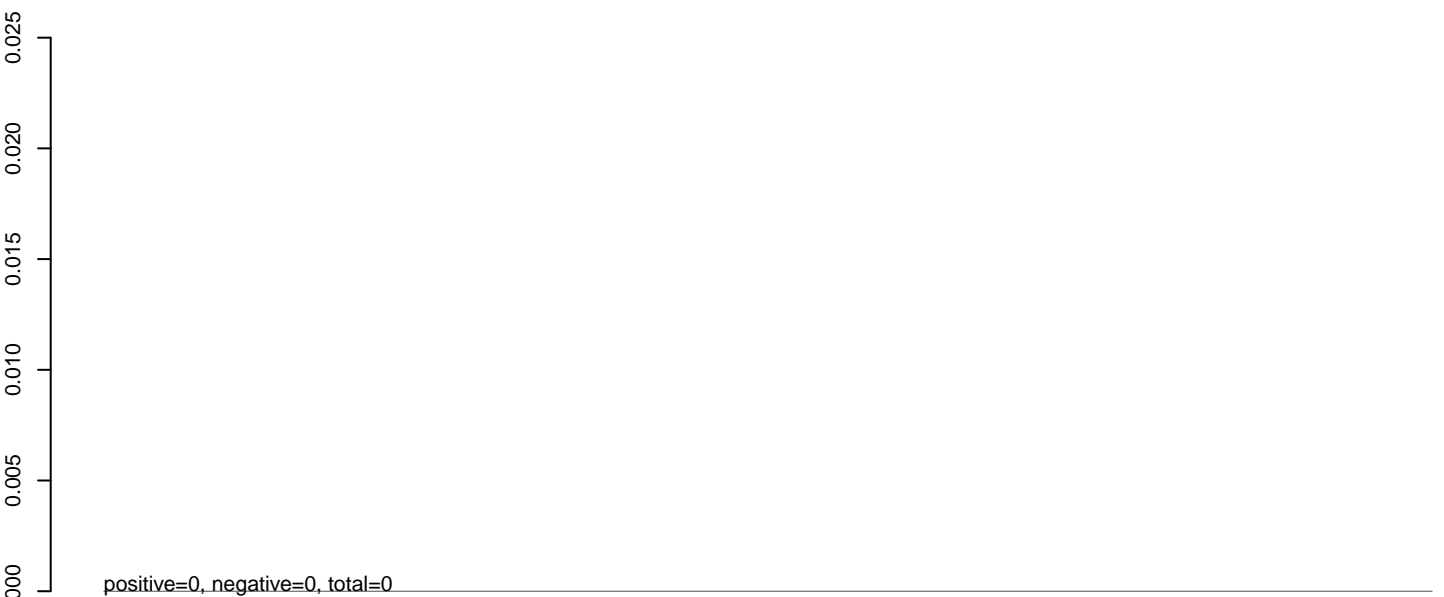
AeAeg_Ovary_GH.rep



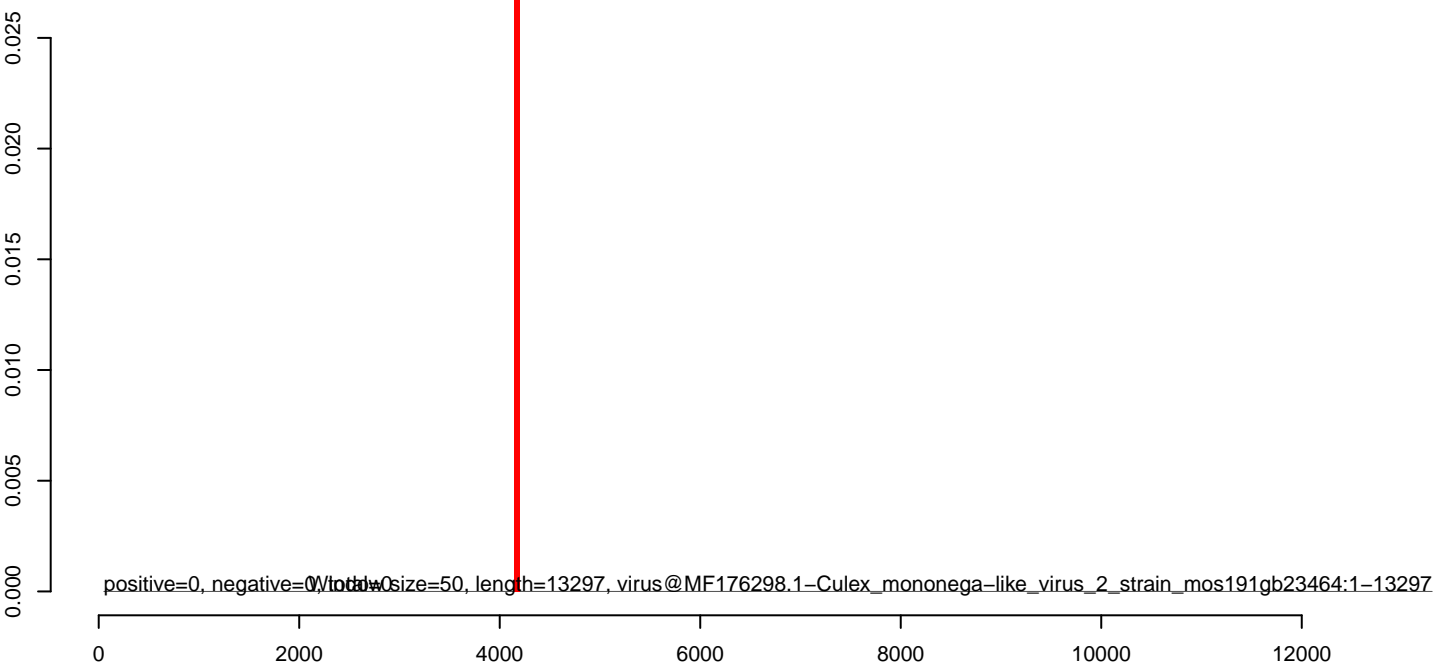
AeAeg_Ovary_GH.18_23.rep



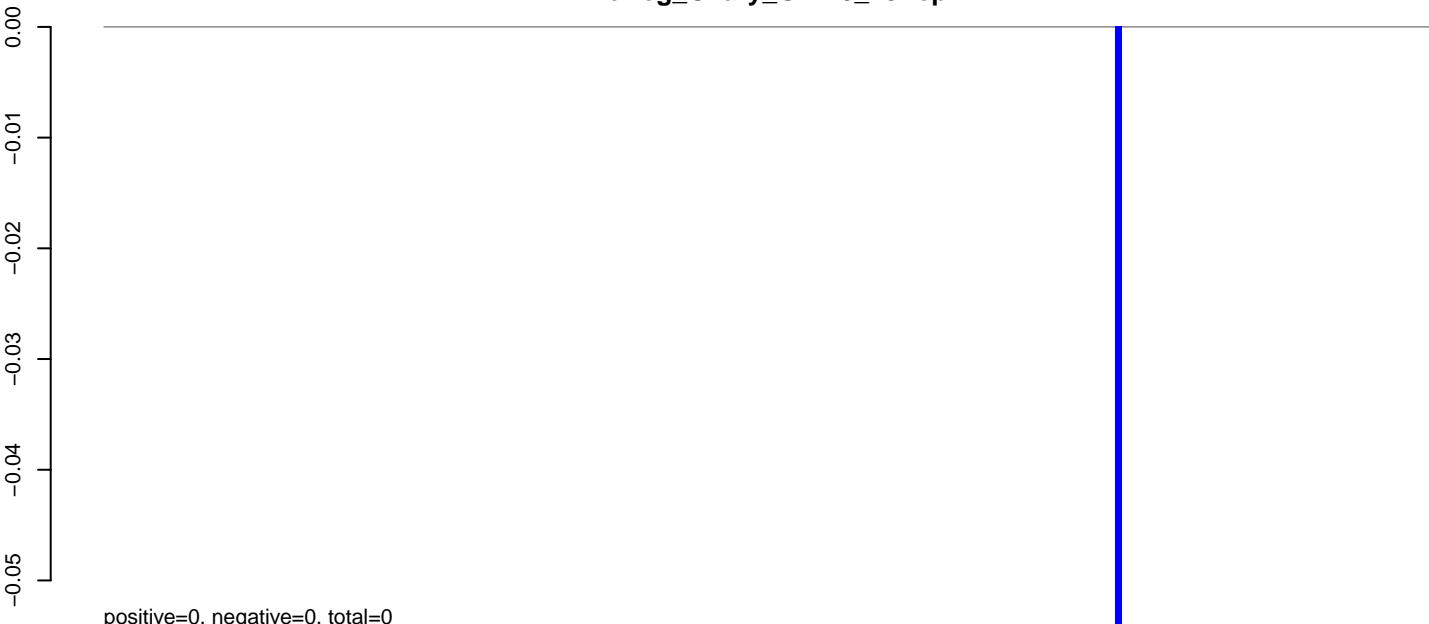
AeAeg_Ovary_GH.24_35.rep



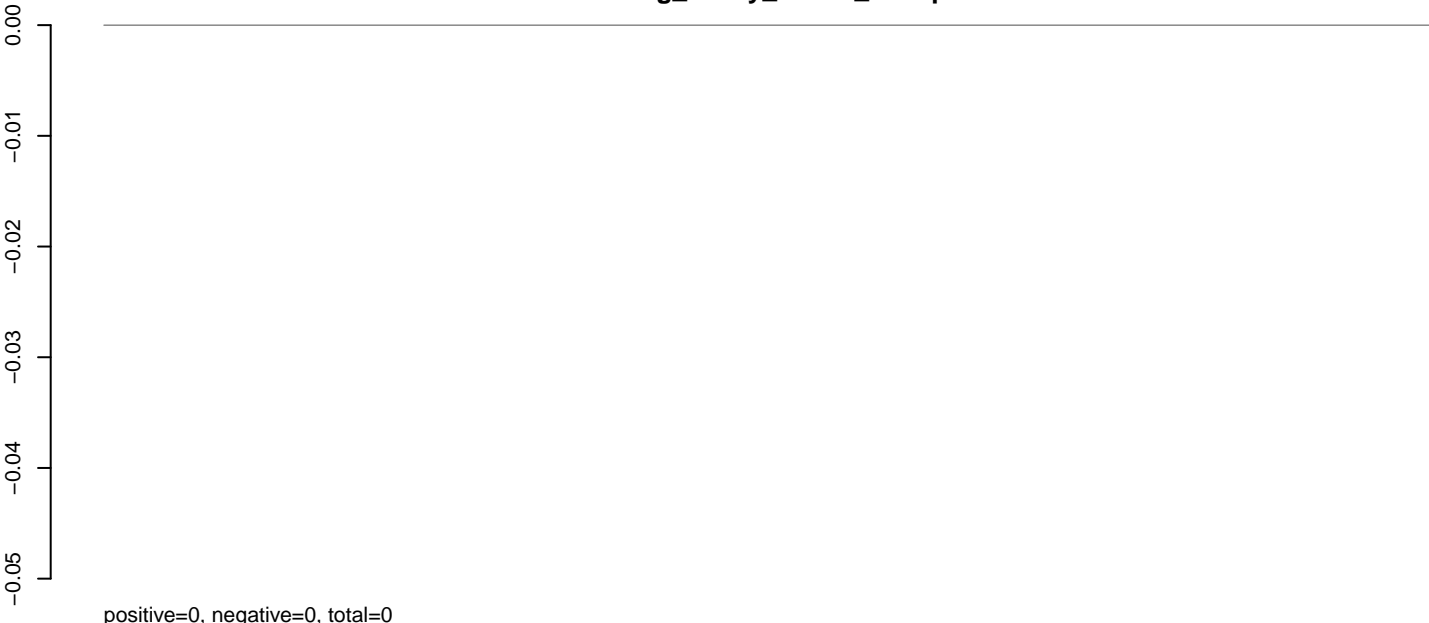
AeAeg_Ovary_GH.rep



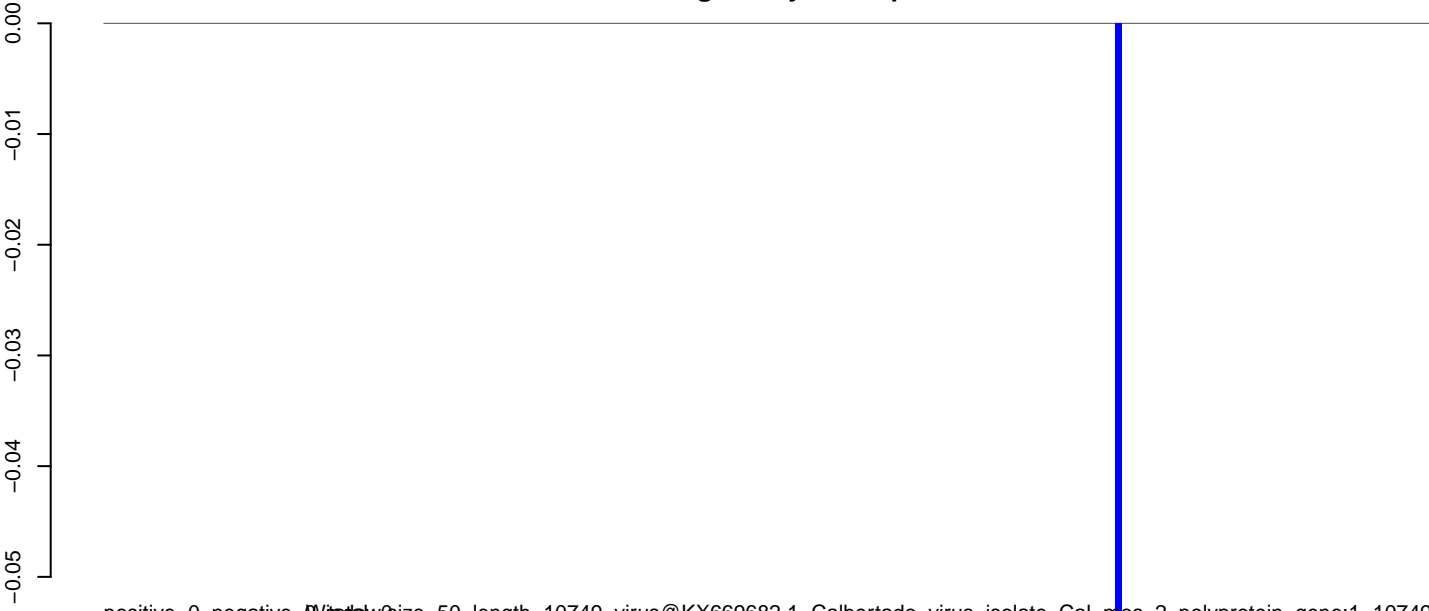
AeAeg_Ovary_GH.18_23.rep



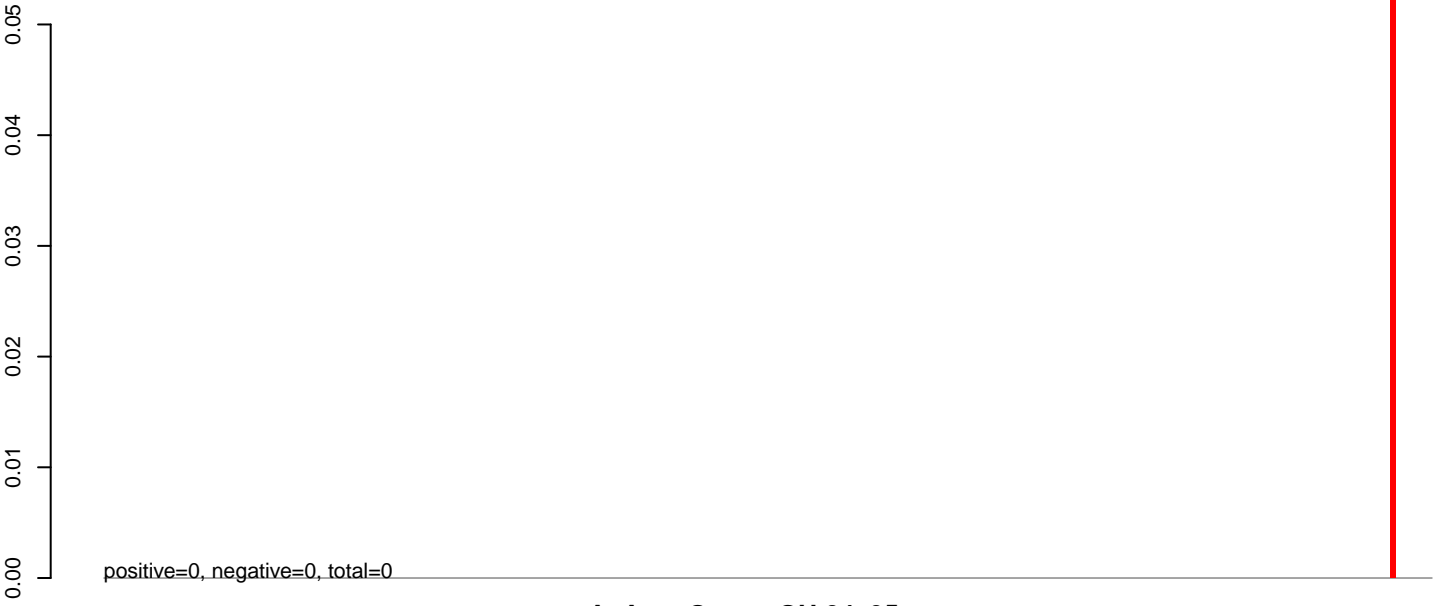
AeAeg_Ovary_GH.24_35.rep



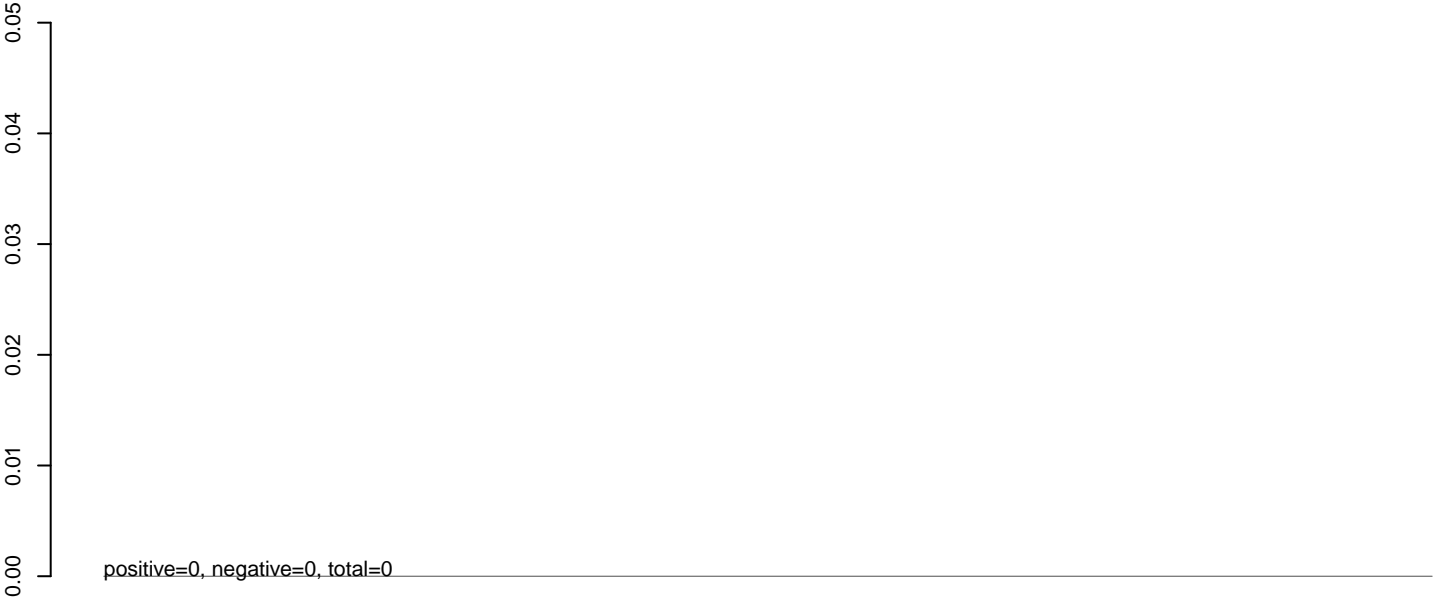
AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



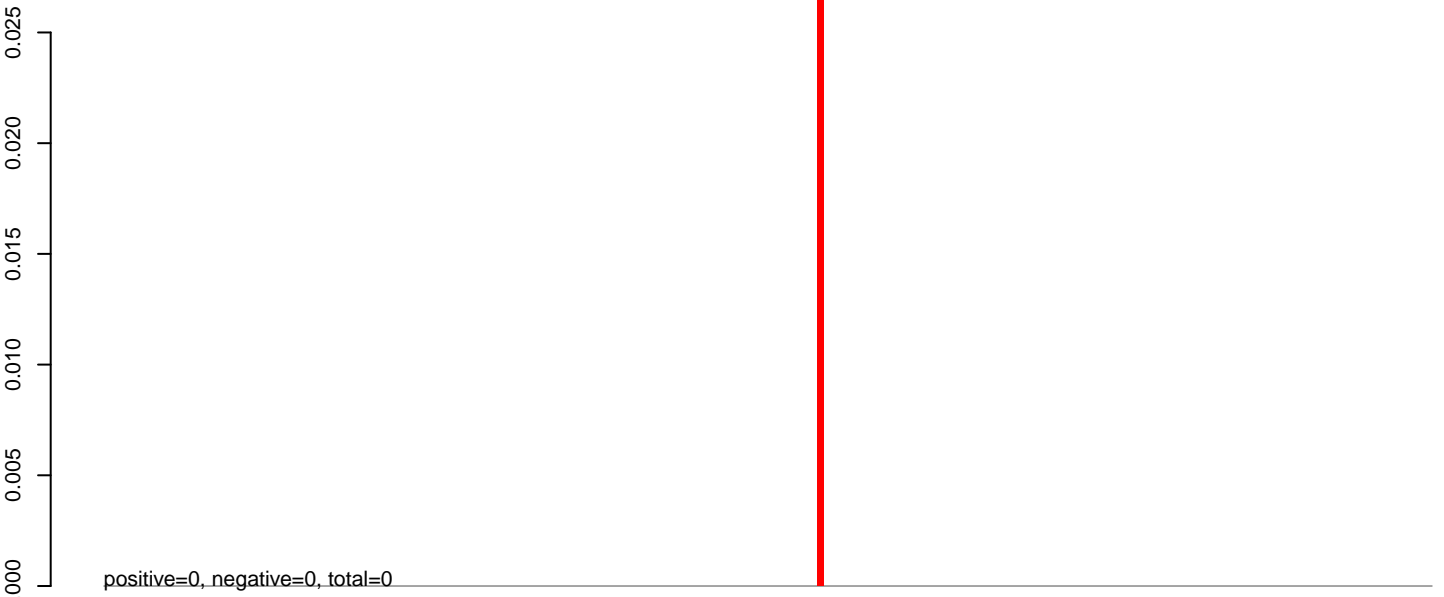
AeAeg_Ovary_GH.24_35.rep



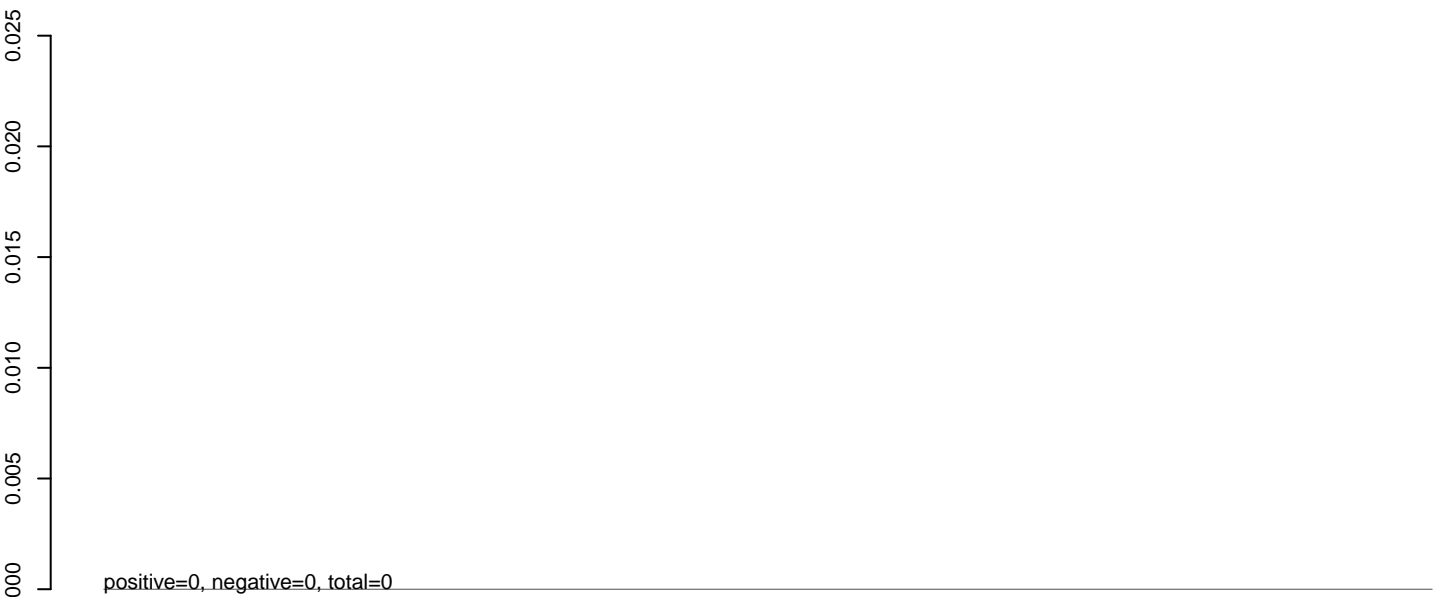
AeAeg_Ovary_GH.rep



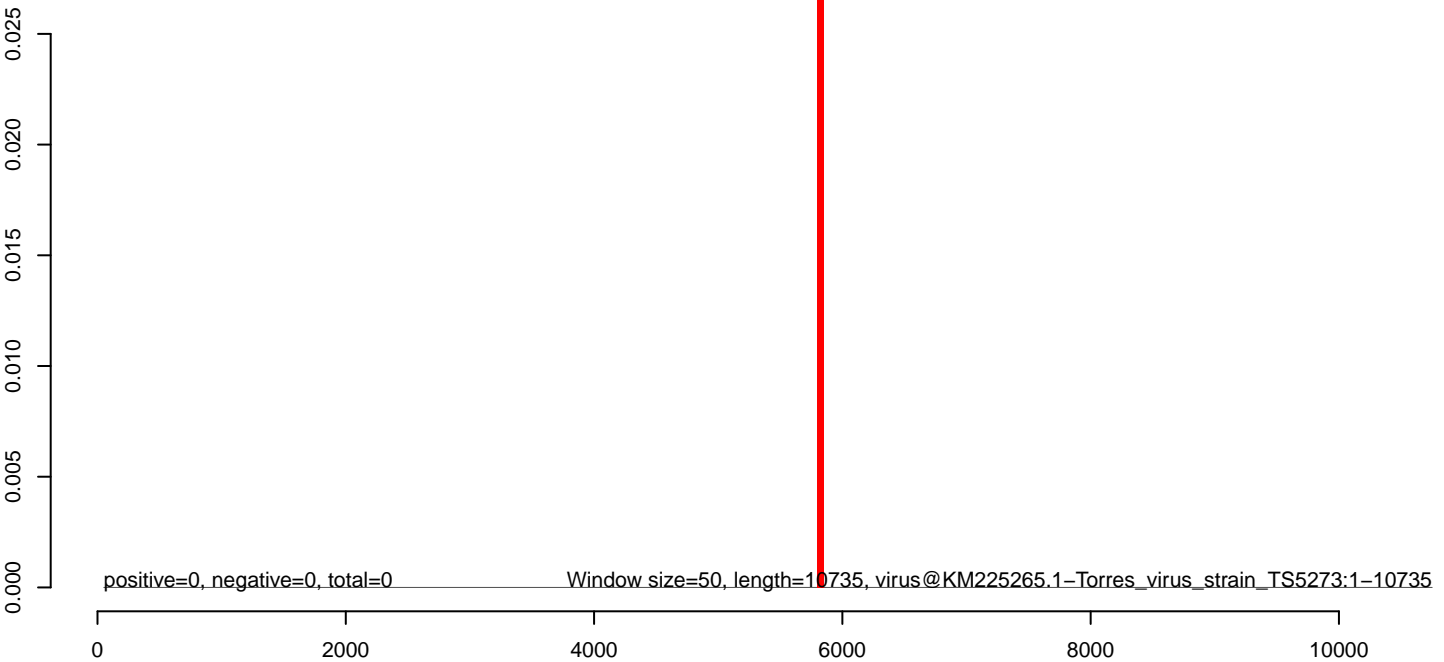
AeAeg_Ovary_GH.18_23.rep



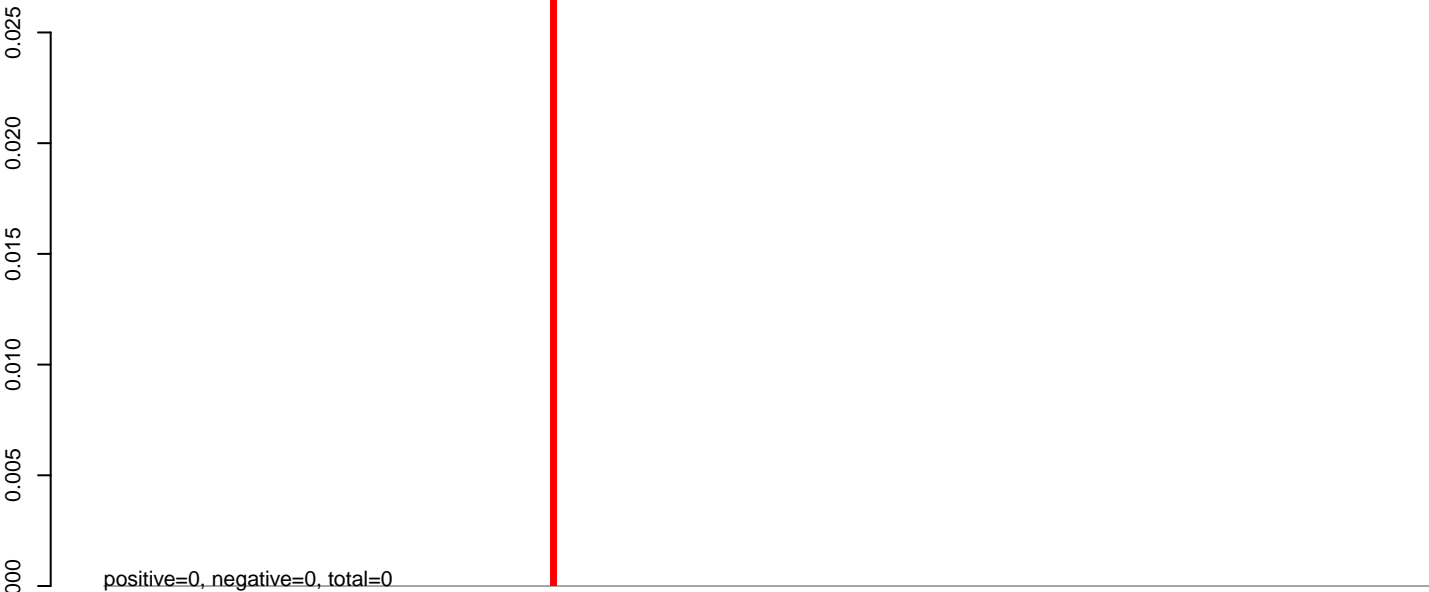
AeAeg_Ovary_GH.24_35.rep



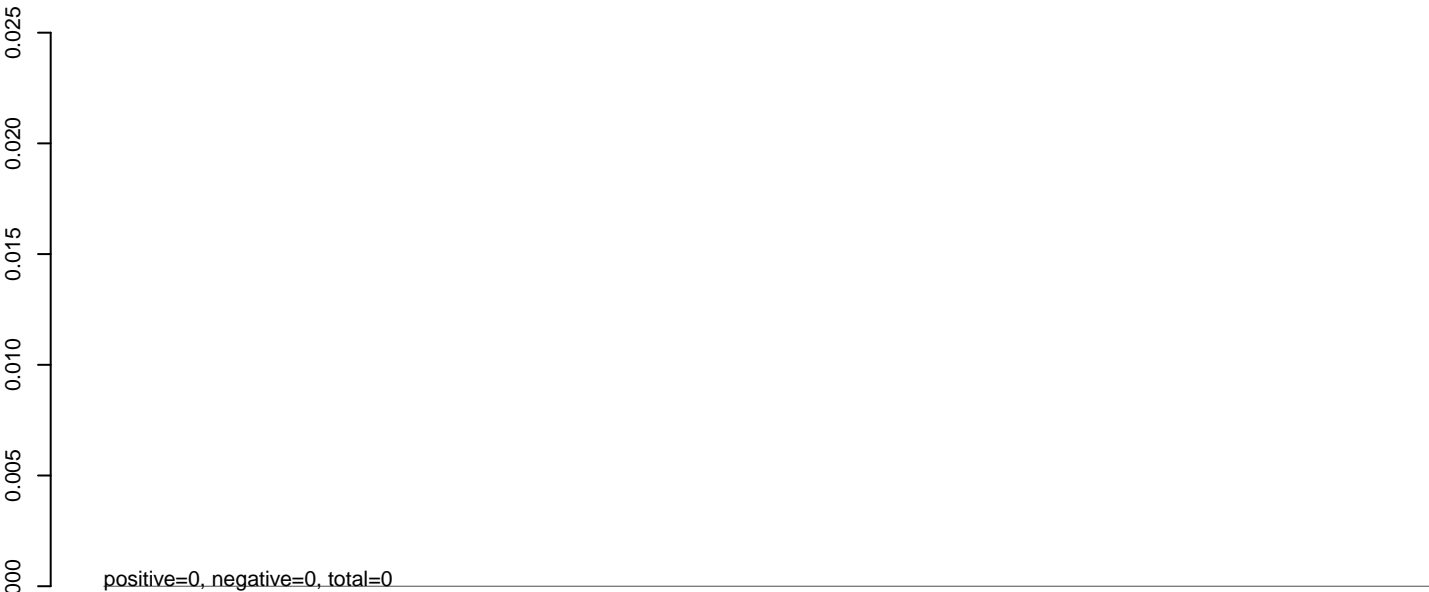
AeAeg_Ovary_GH.rep



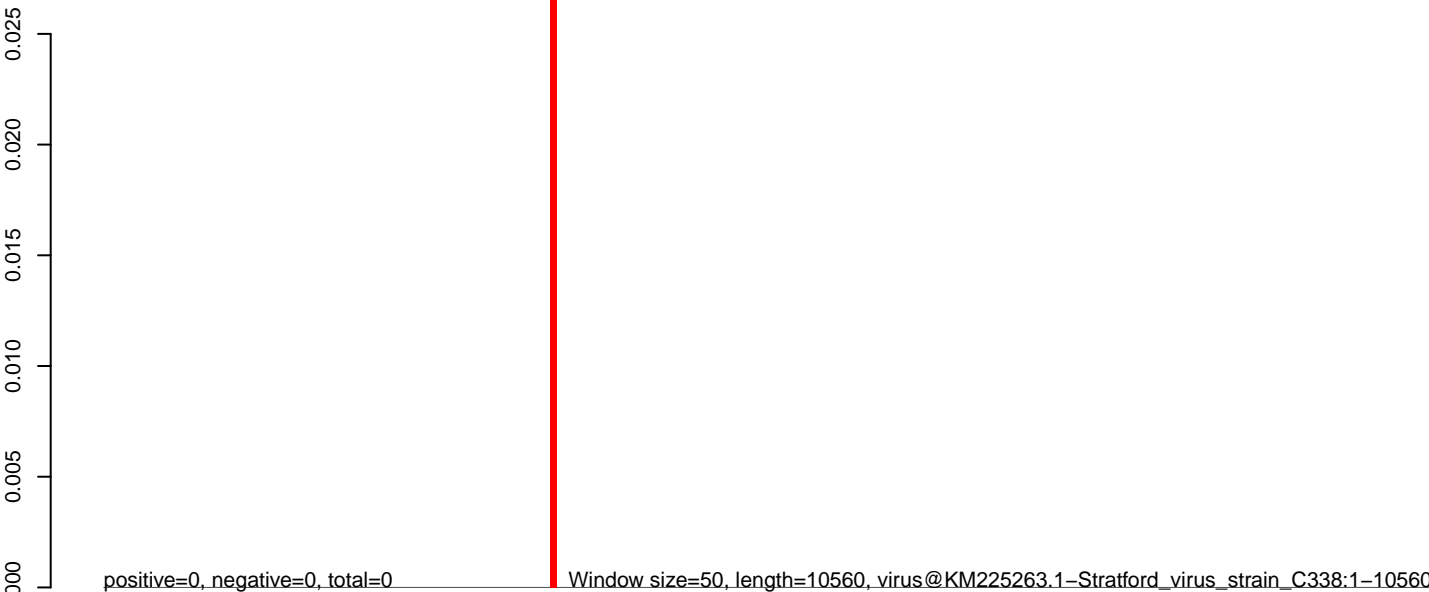
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

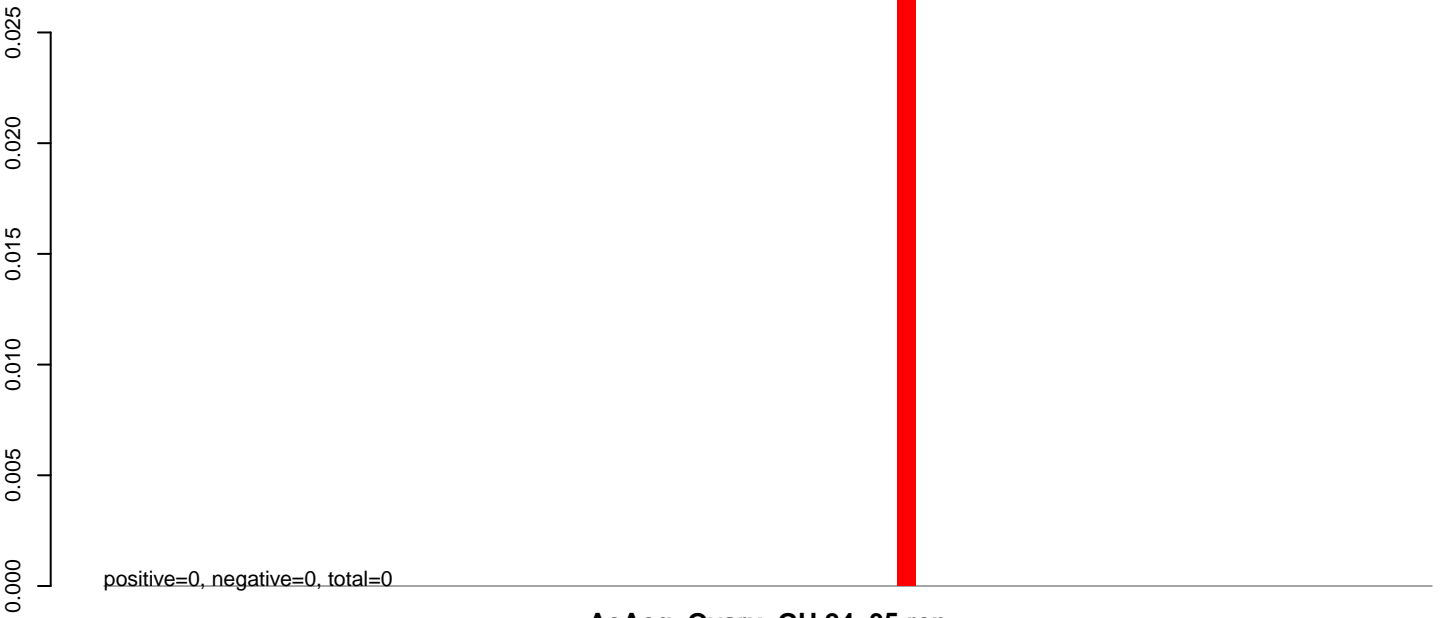


AeAeg_Ovary_GH.rep

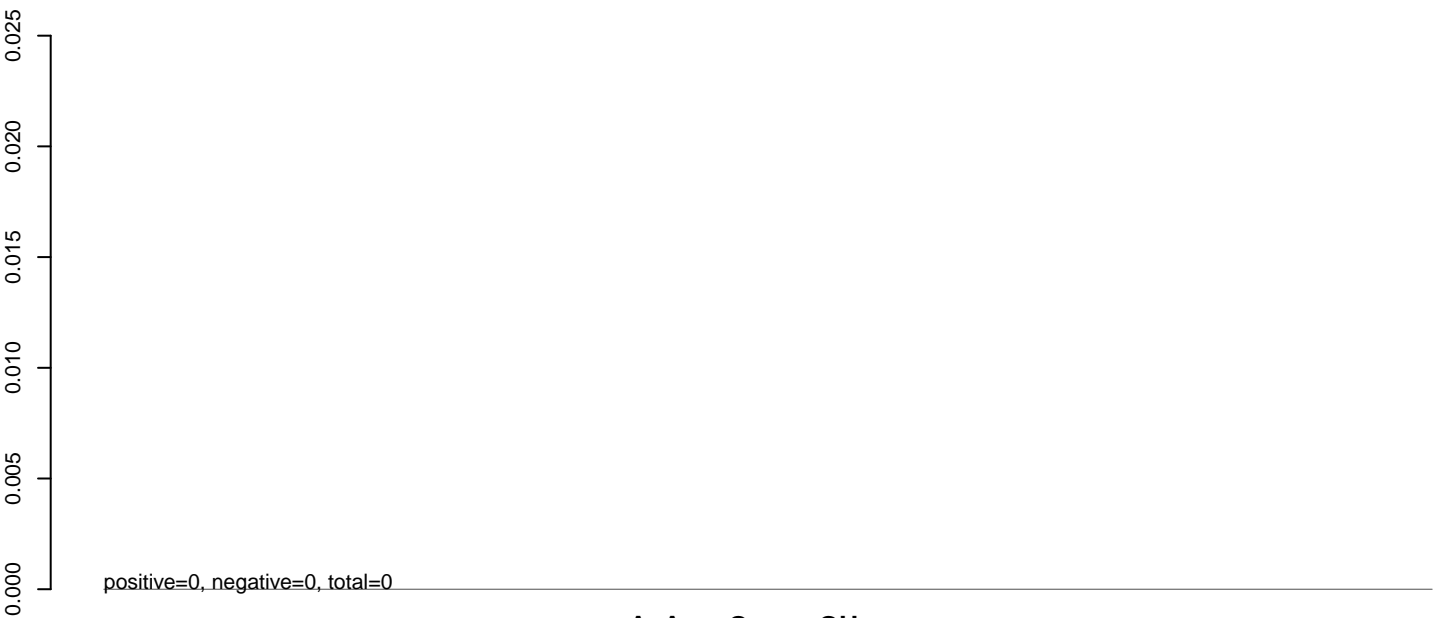


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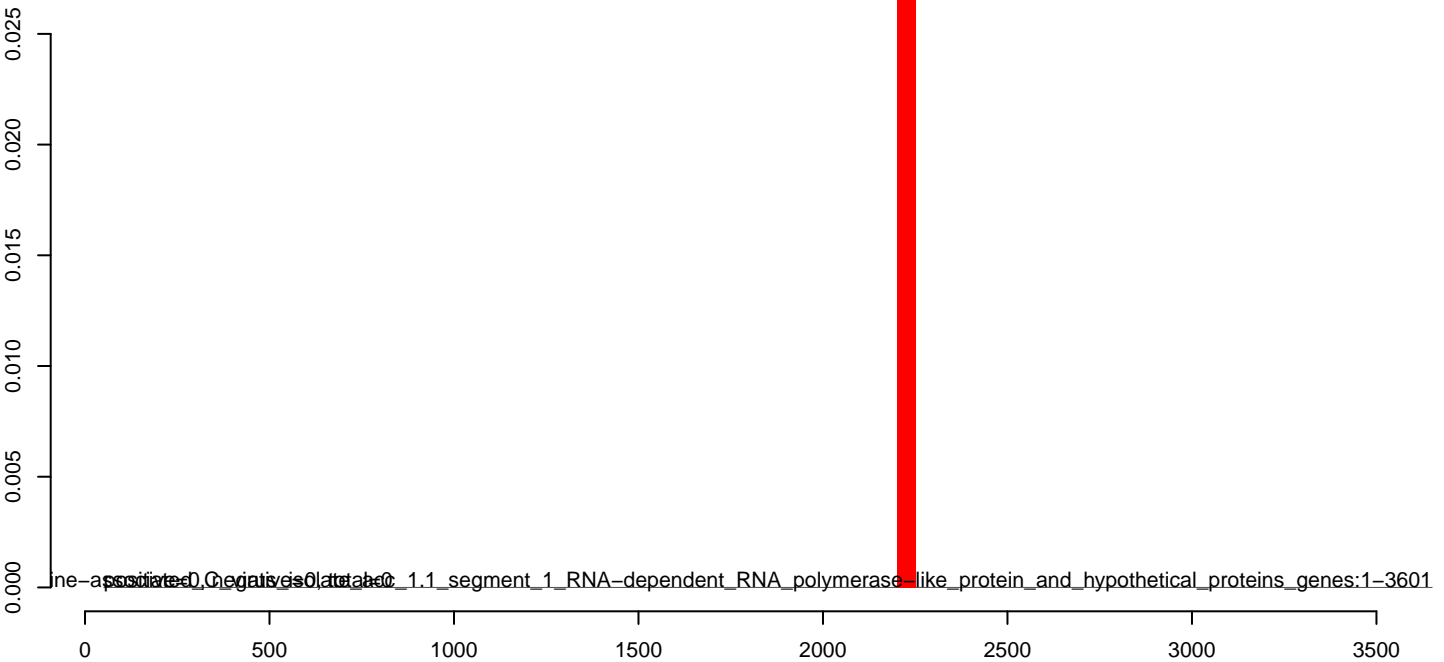
AeAeg_Ovary_GH.18_23.rep



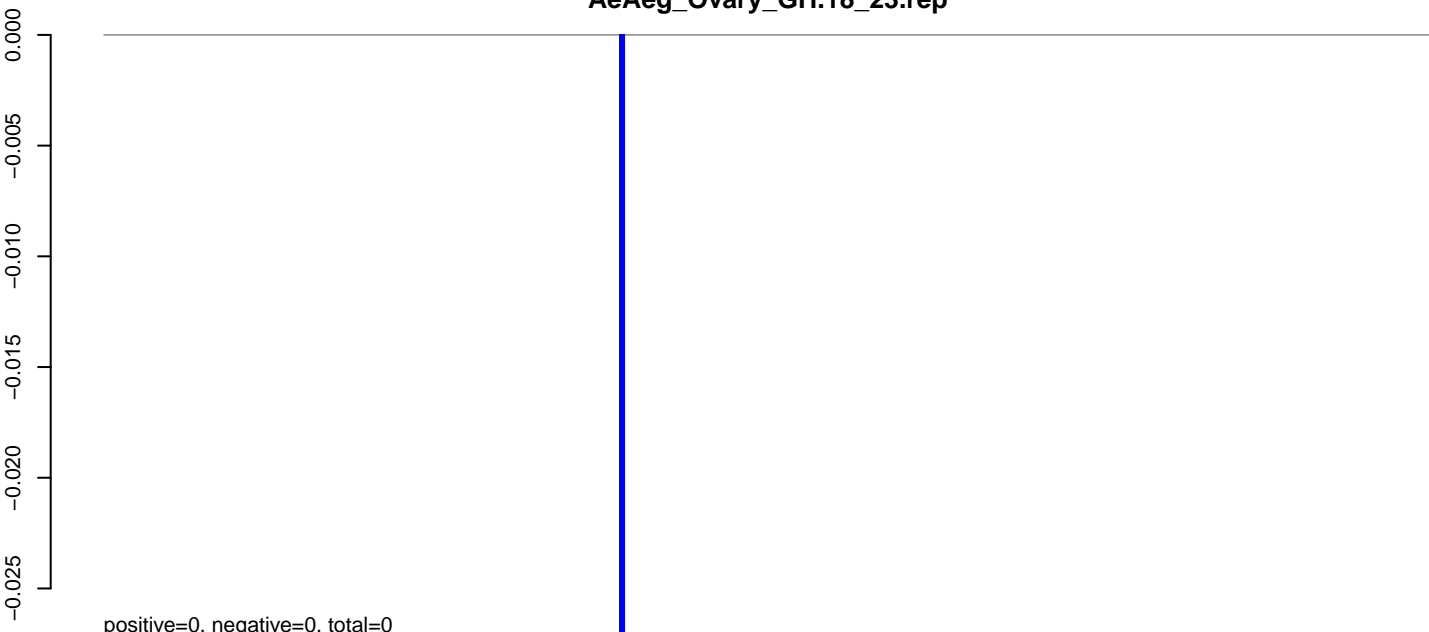
AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



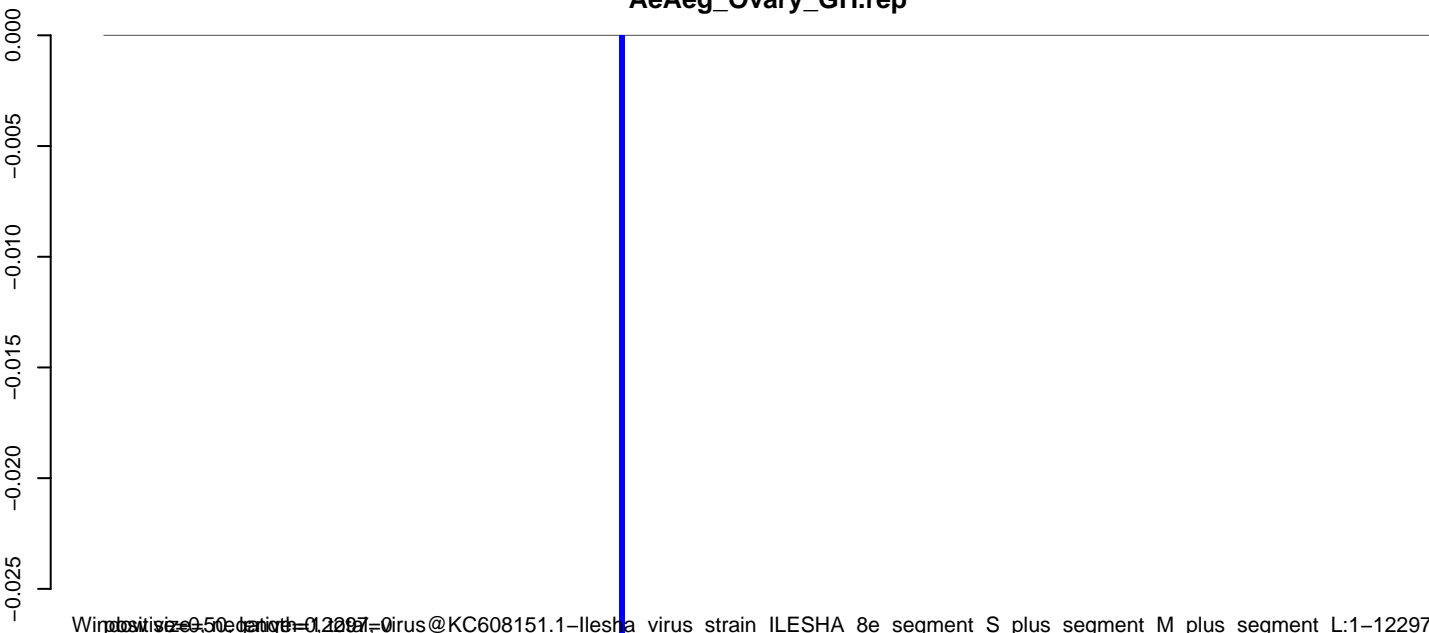
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



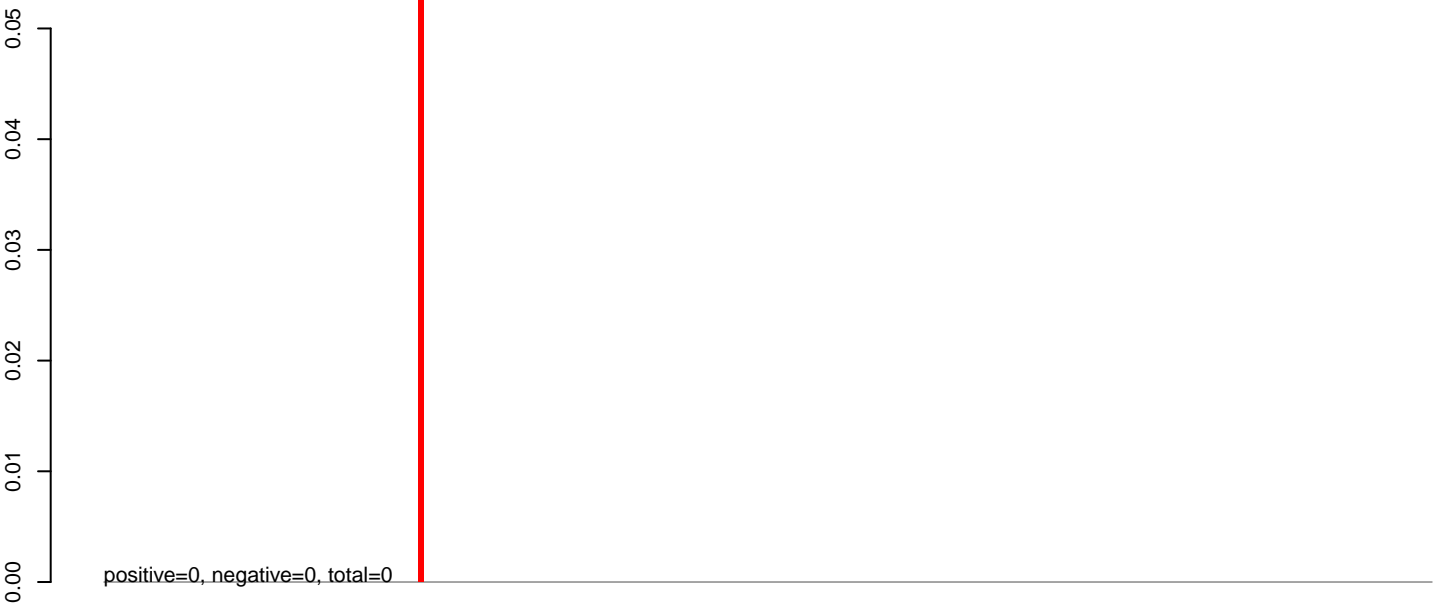
AeAeg_Ovary_GH.rep



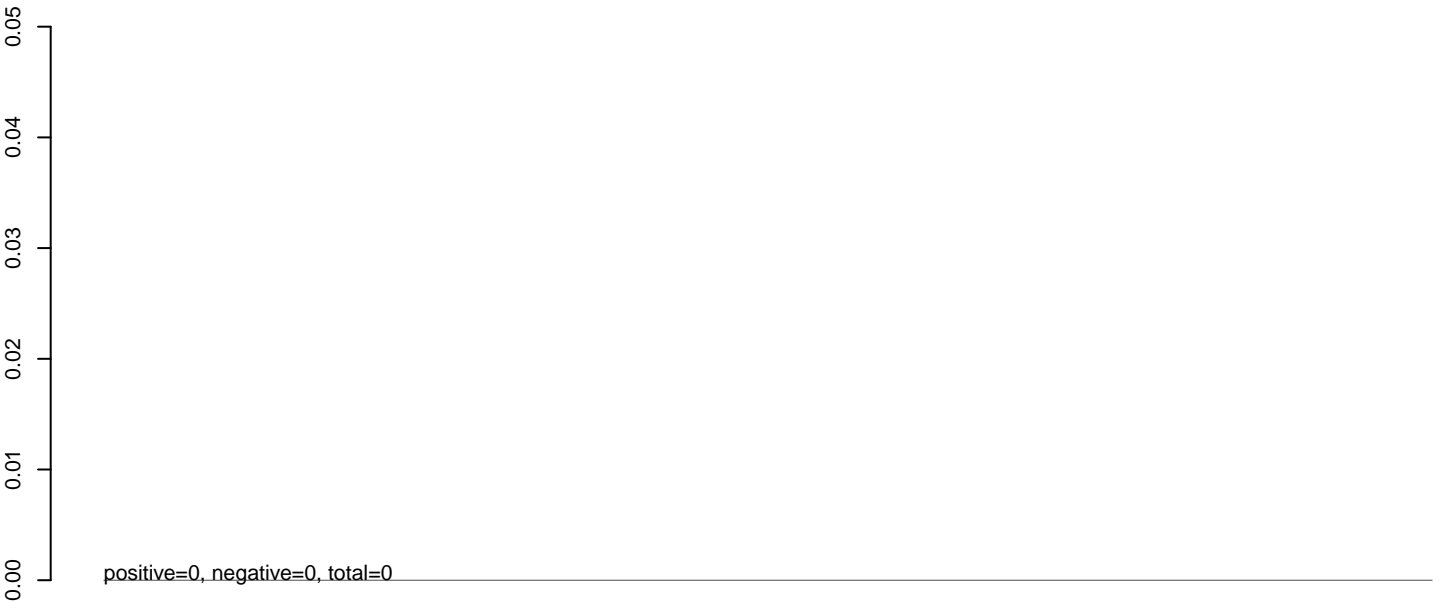
Winnipeg, 50, length=0, 2097, Virus@KC608151.1-Ilesha_virus_strain_ILESHA_8e_segment_S_plus_segment_M_plus_segment_L:1-12297

0 2000 4000 6000 8000 10000 12000

AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

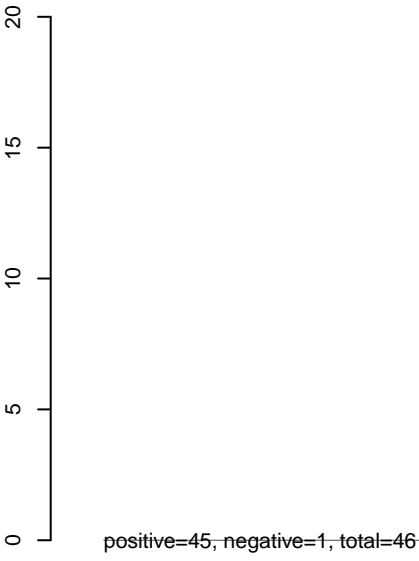


AeAeg_Ovary_GH.rep

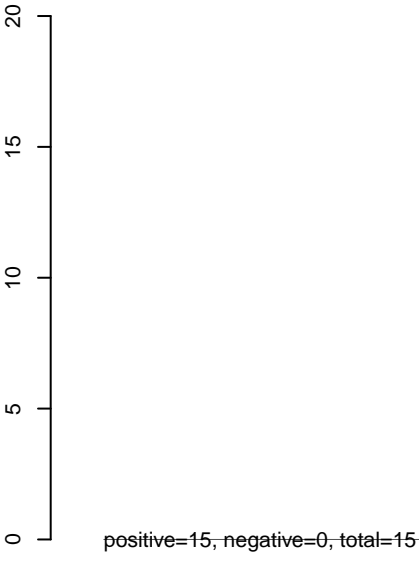


0 2000 4000 6000 8000 10000 12000

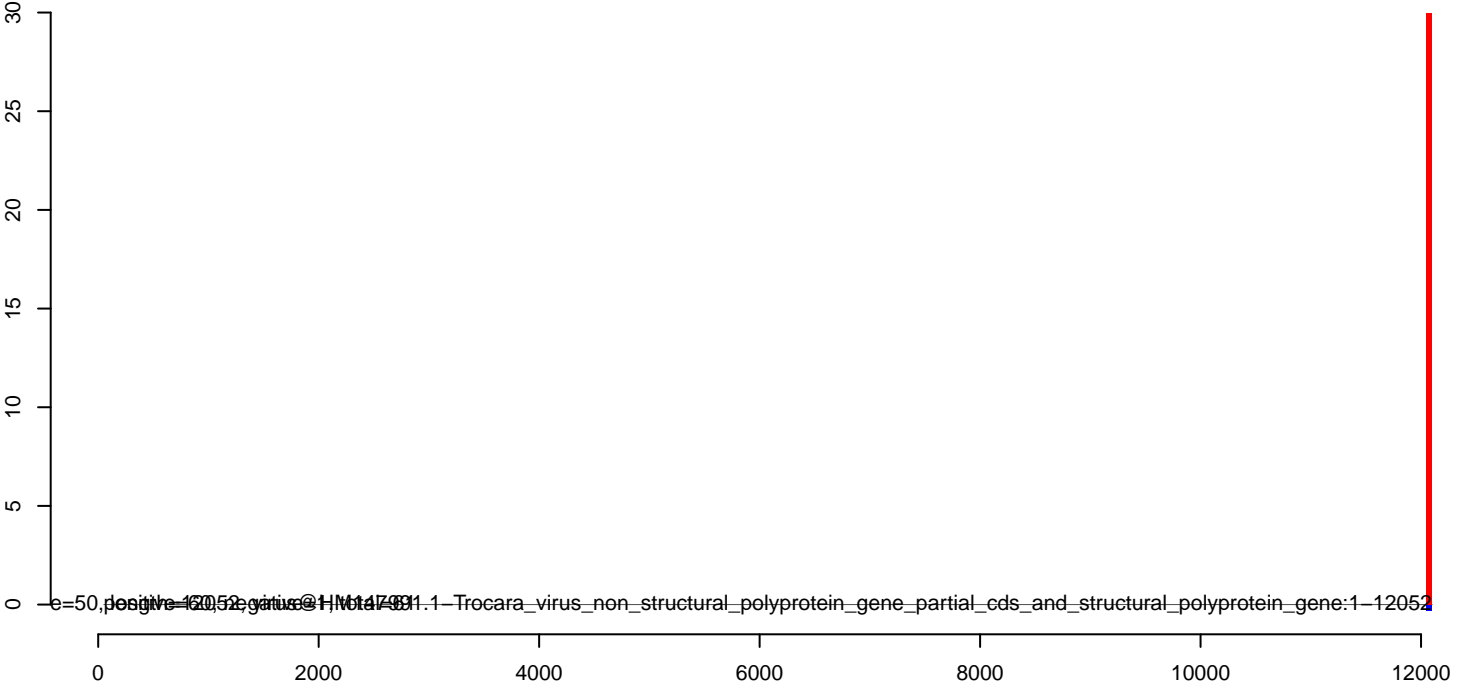
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



12052-HM14791.1-Trocaravirus non-structural polyprotein gene partial cds and structural polyprotein gene:1-12052

AeAeg_Ovary_GH.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

positive=0, negative=0, total=0

AeAeg_Ovary_GH.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

positive=0, negative=0, total=0

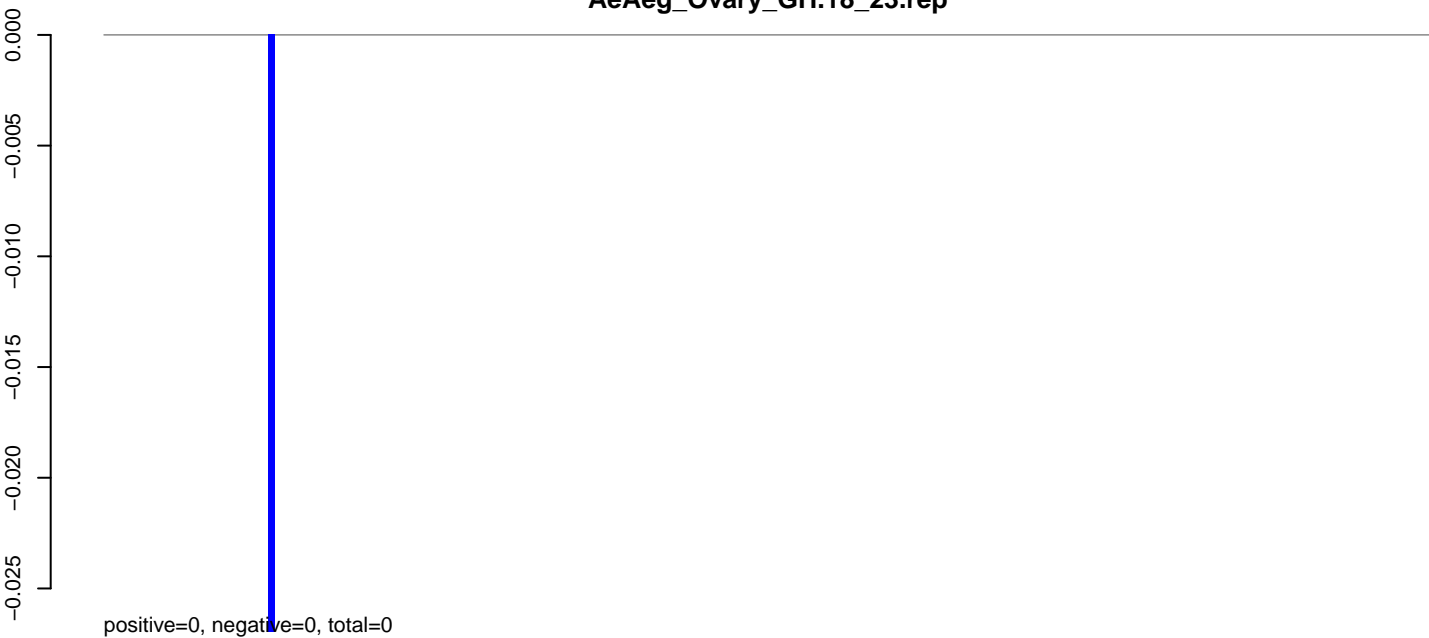
AeAeg_Ovary_GH.rep

0.000
-0.005
-0.010
-0.015
-0.020
-0.025

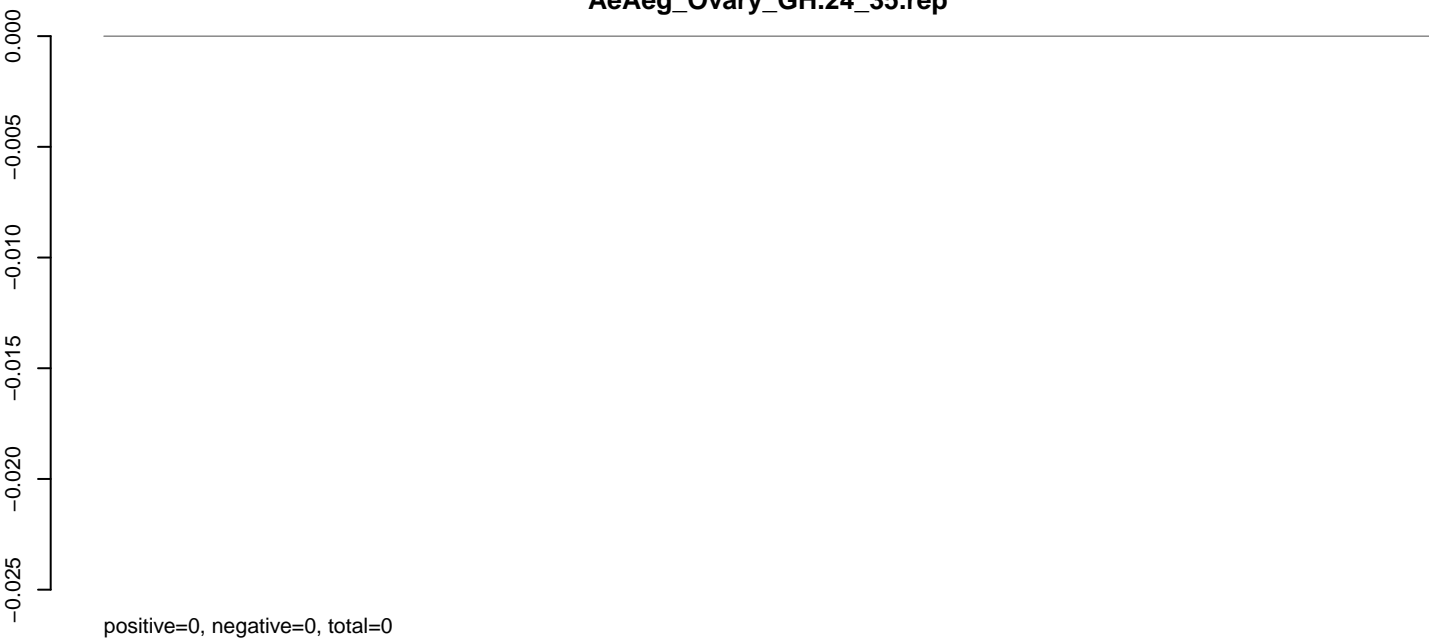
positive=0, negative=0, total=0, window size=50, length=10254, virus@DQ235149.1-Royal_Farm_virus_from_Afghanistan_polyprotein_gene:1-10254

0 2000 4000 6000 8000 10000

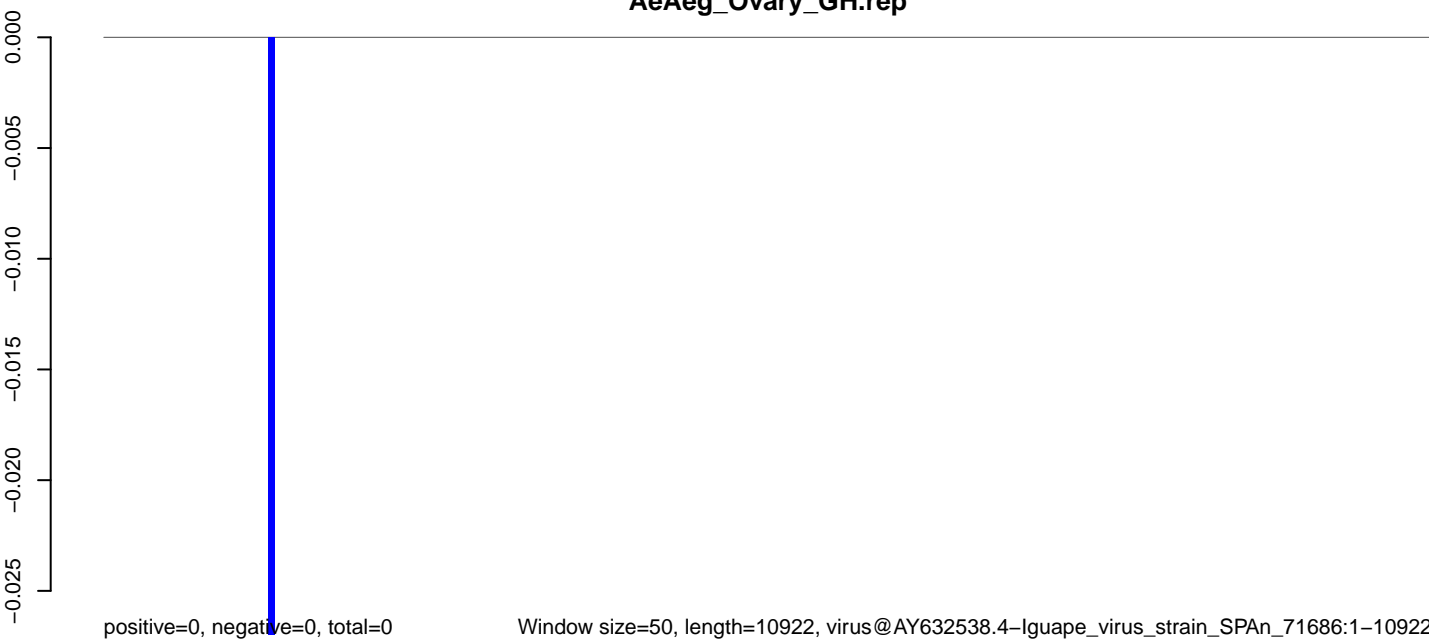
AeAeg_Ovary_GH.18_23.rep



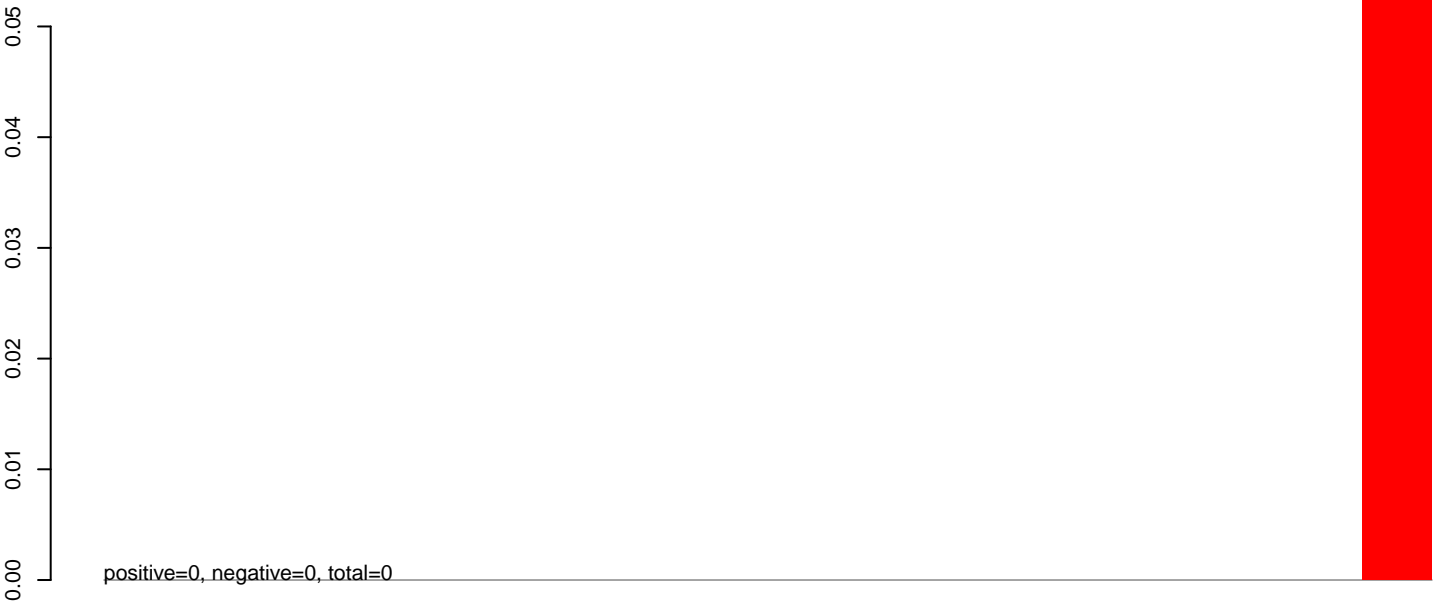
AeAeg_Ovary_GH.24_35.rep



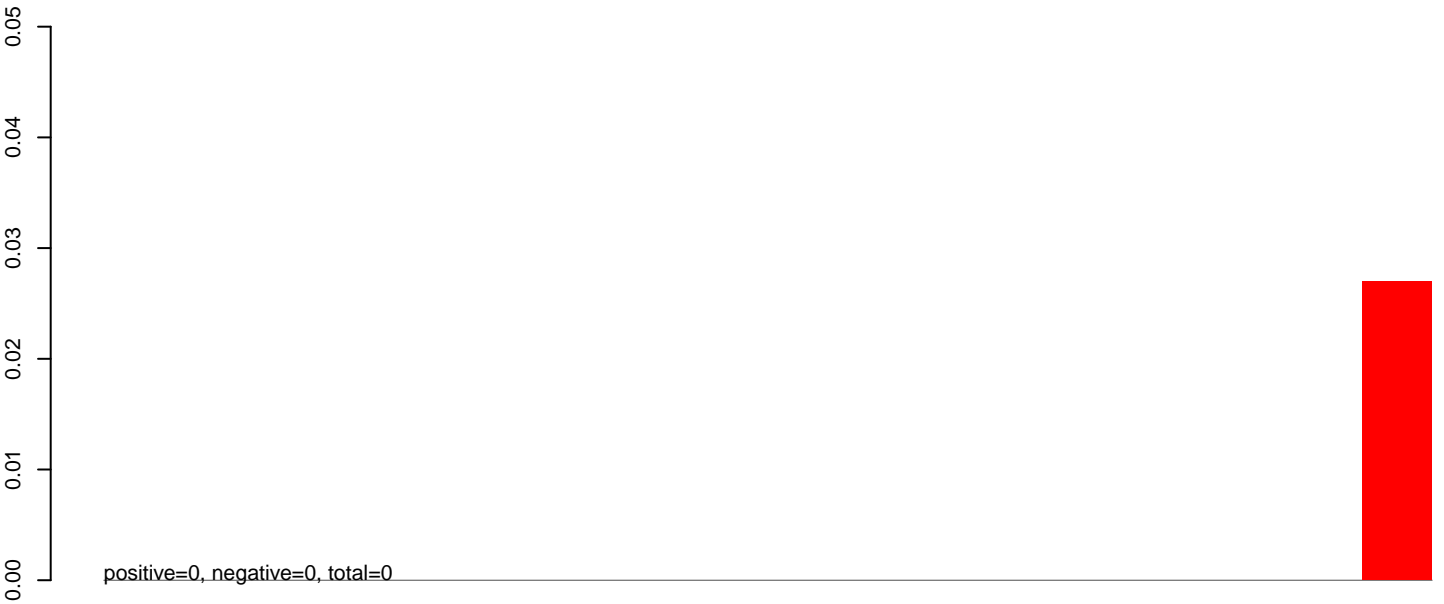
AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



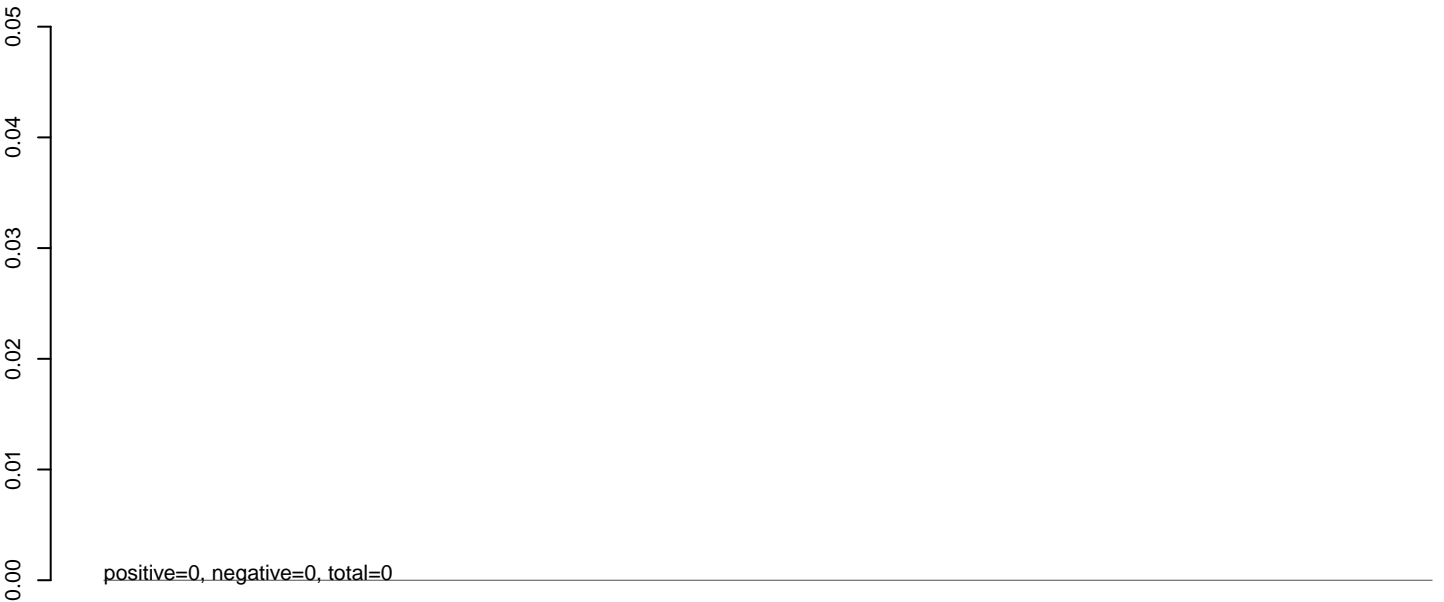
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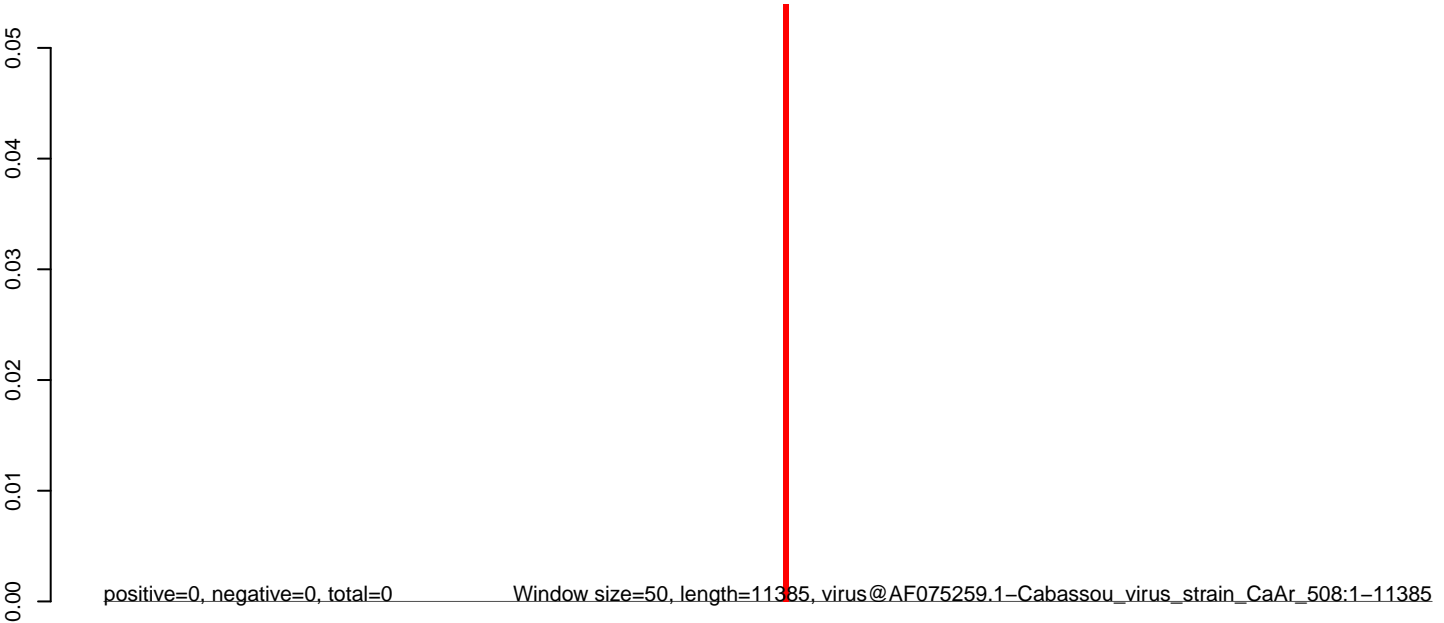
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

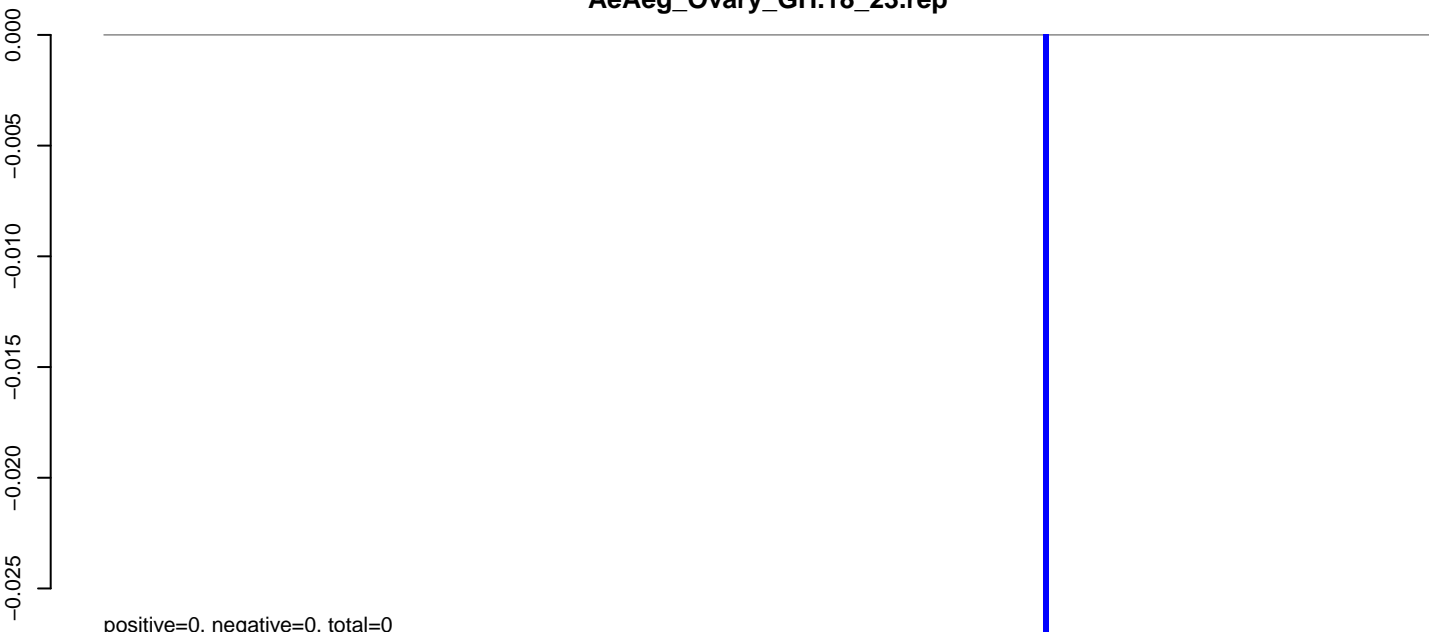


AeAeg_Ovary_GH.rep

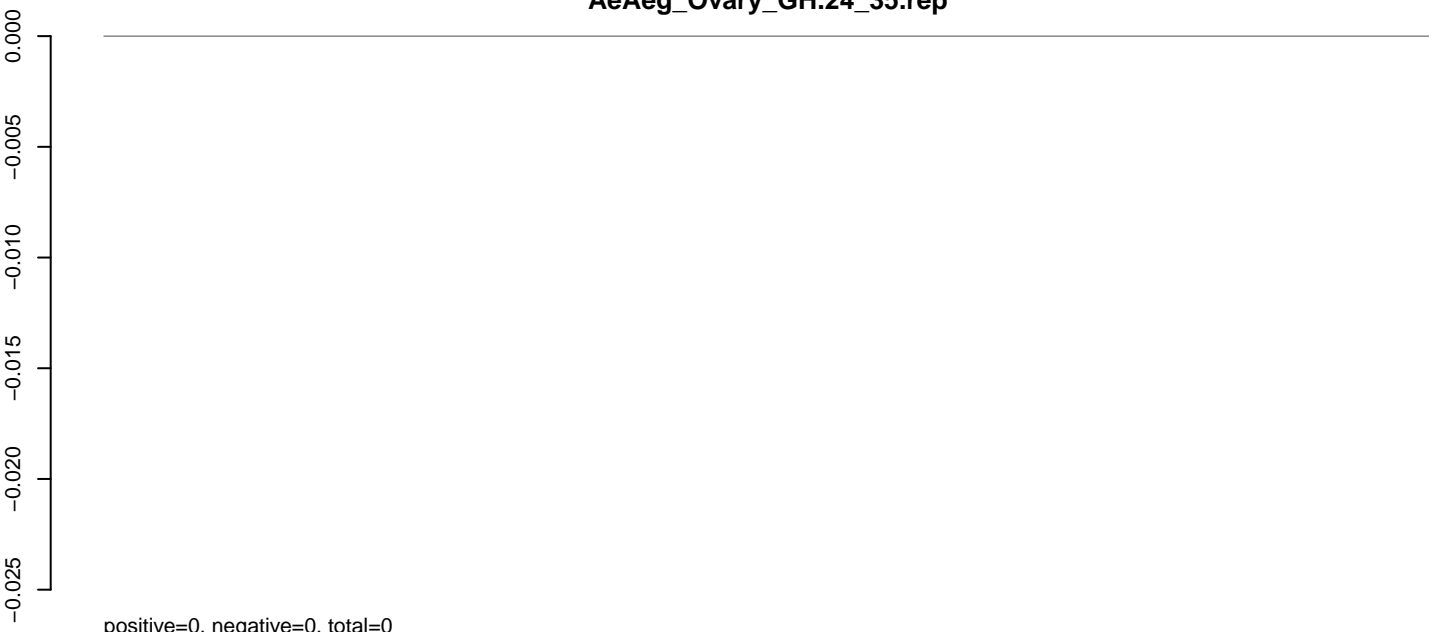


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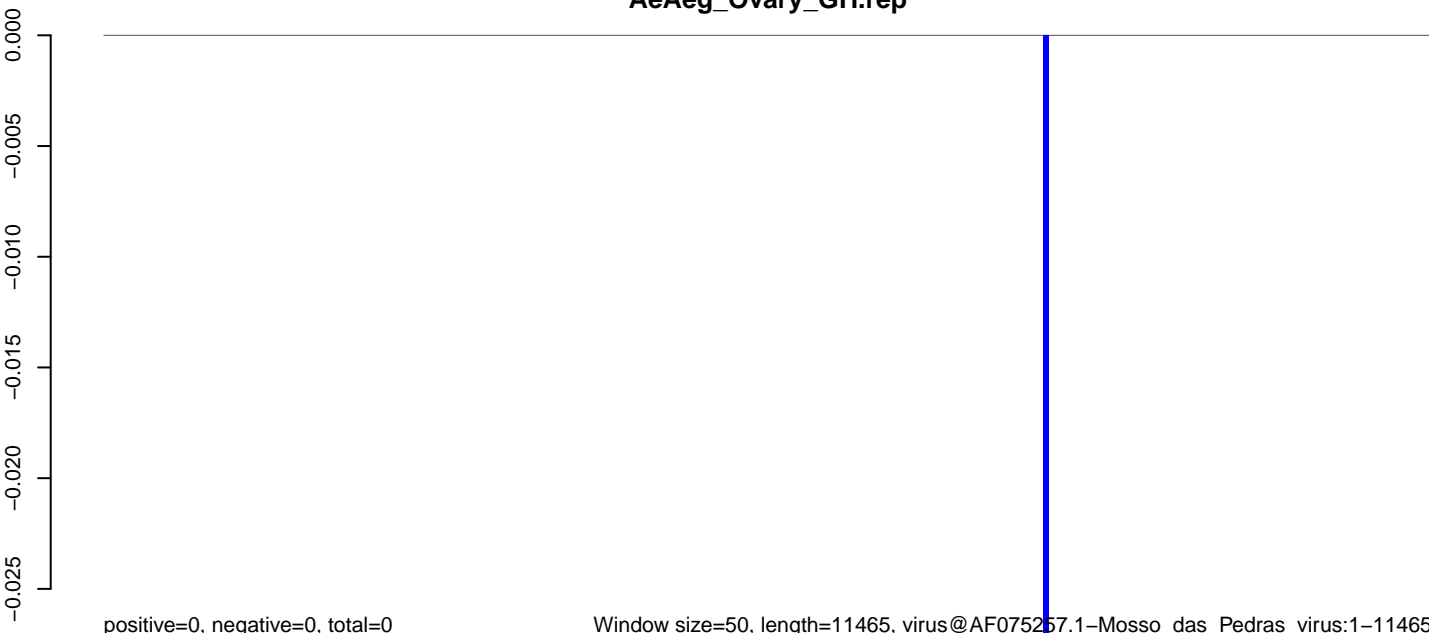
AeAeg_Ovary_GH.18_23.rep



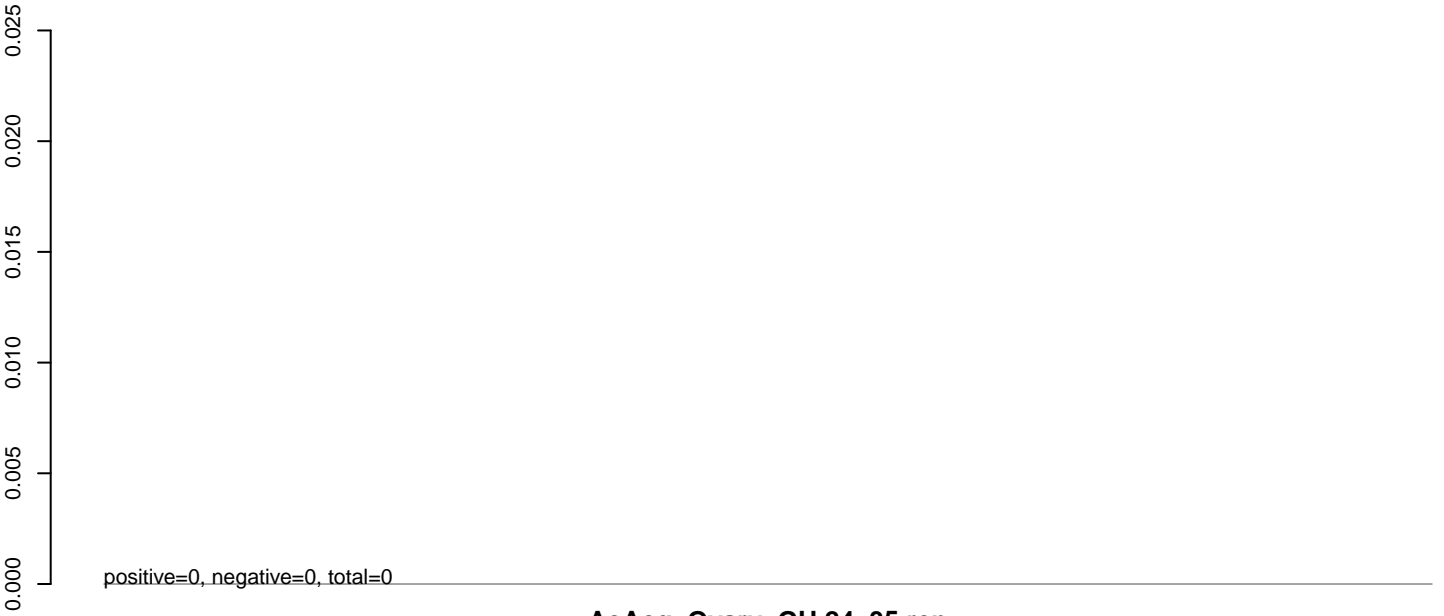
AeAeg_Ovary_GH.24_35.rep



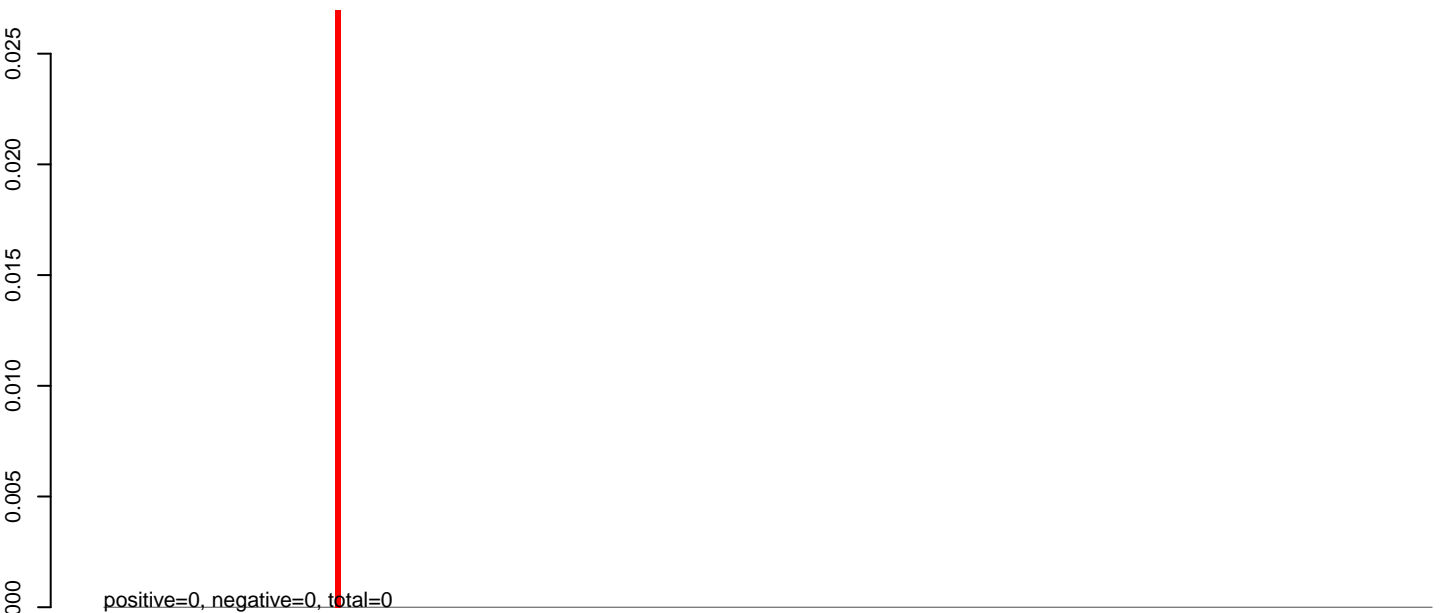
AeAeg_Ovary_GH.rep



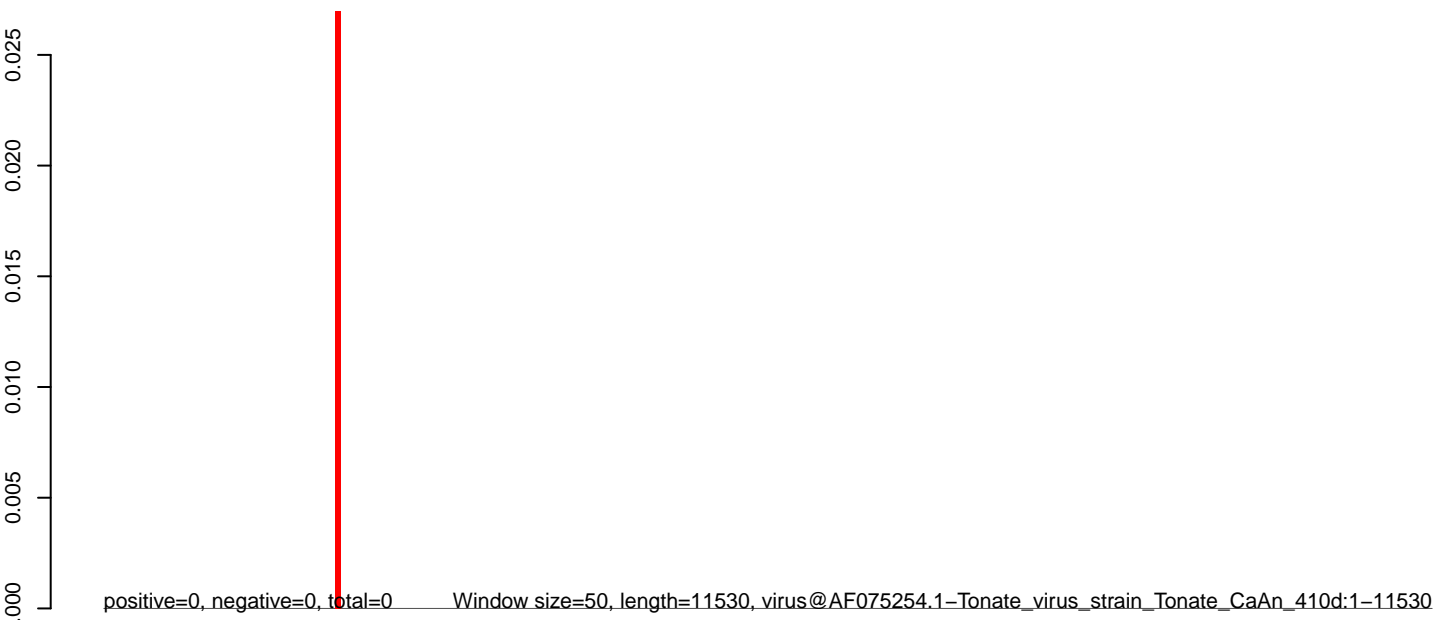
AeAeg_Ovary_GH.18_23.rep



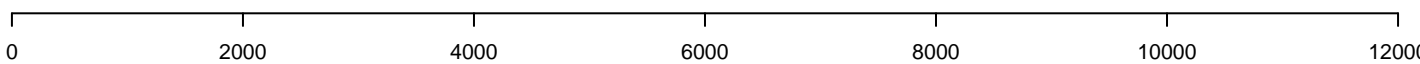
AeAeg_Ovary_GH.24_35.rep



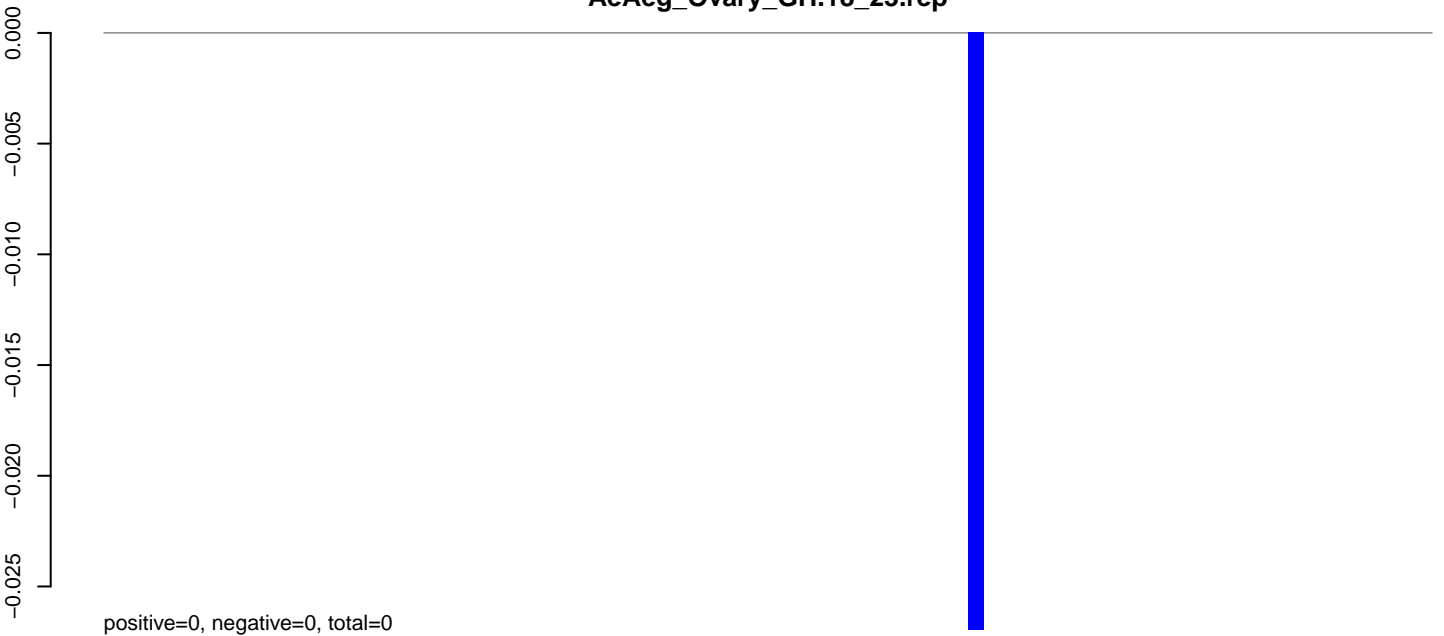
AeAeg_Ovary_GH.rep



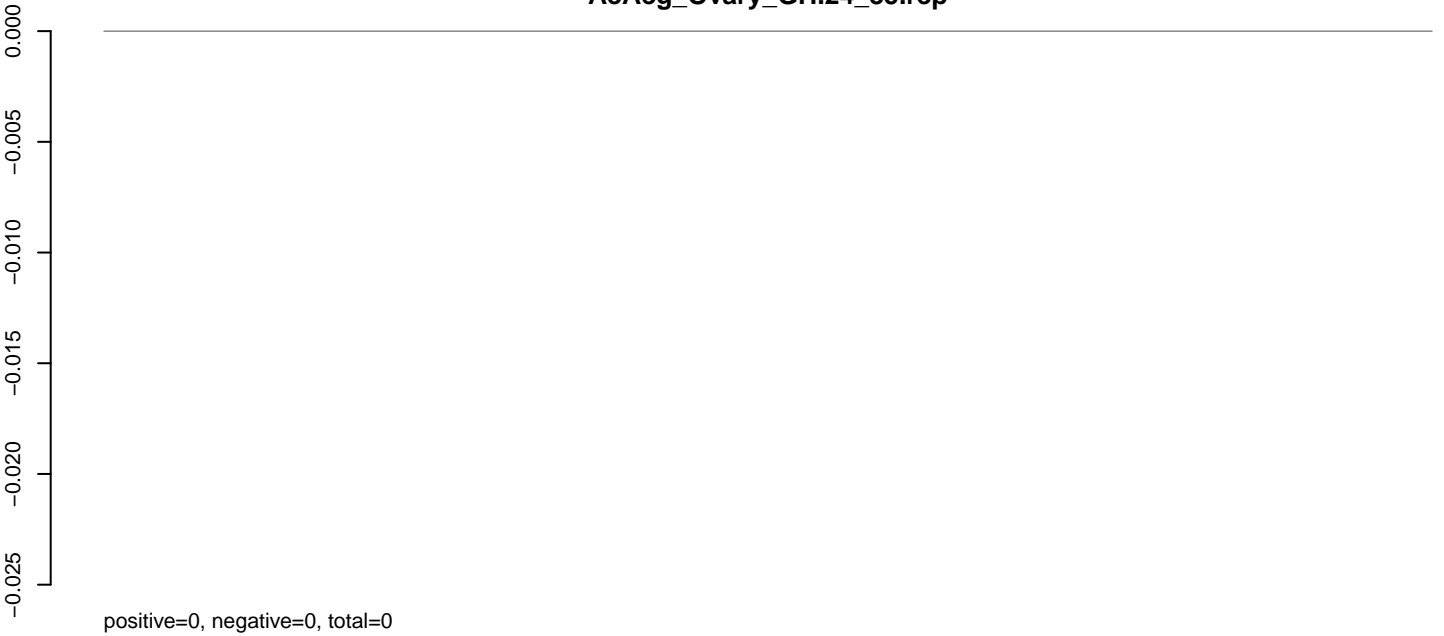
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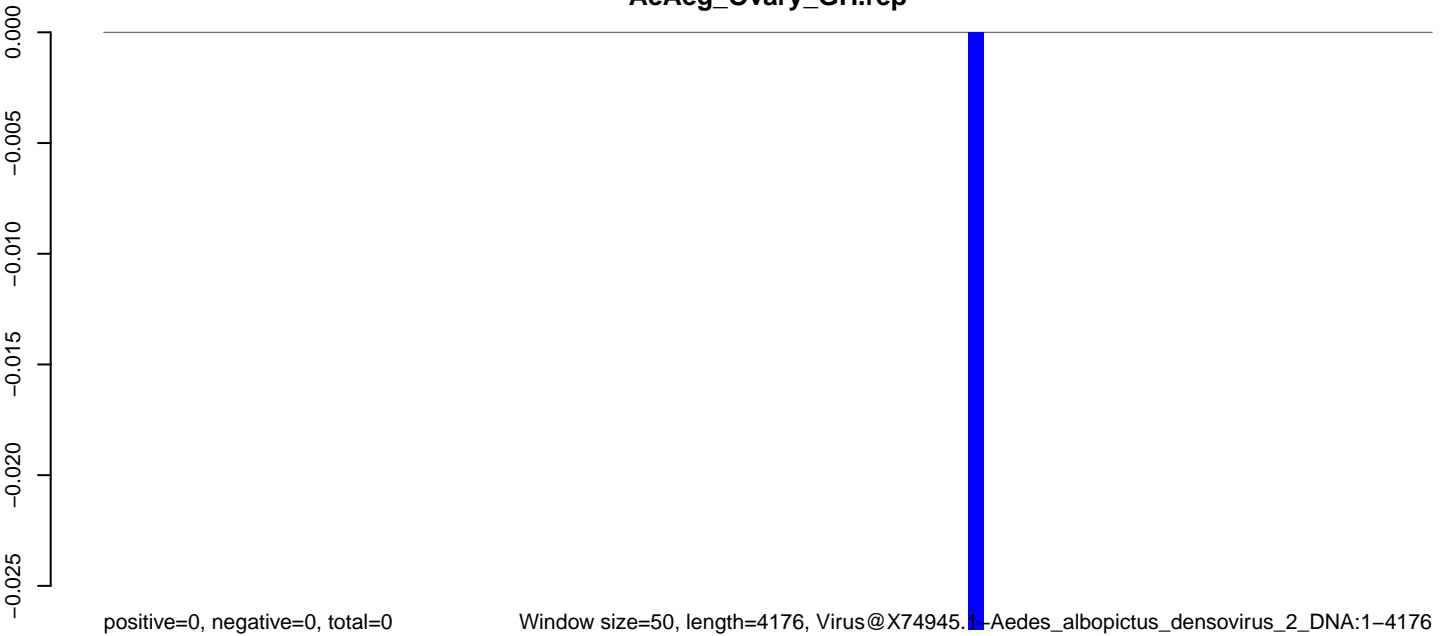
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



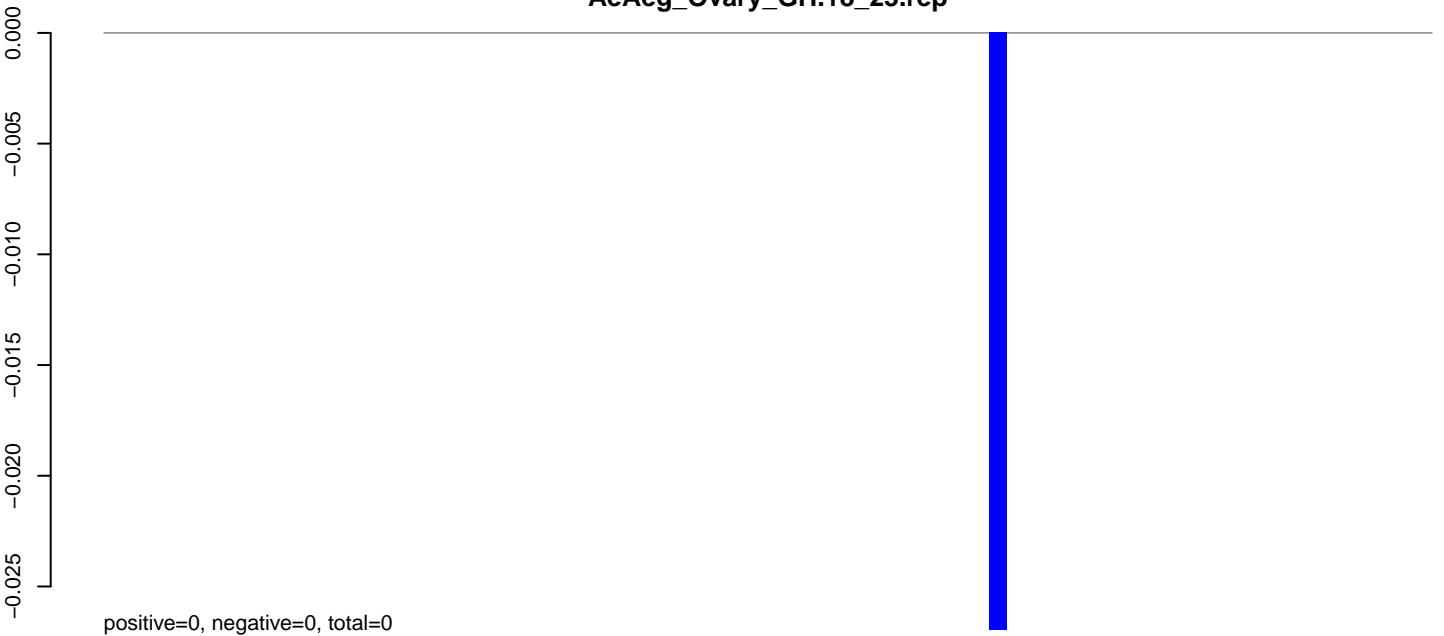
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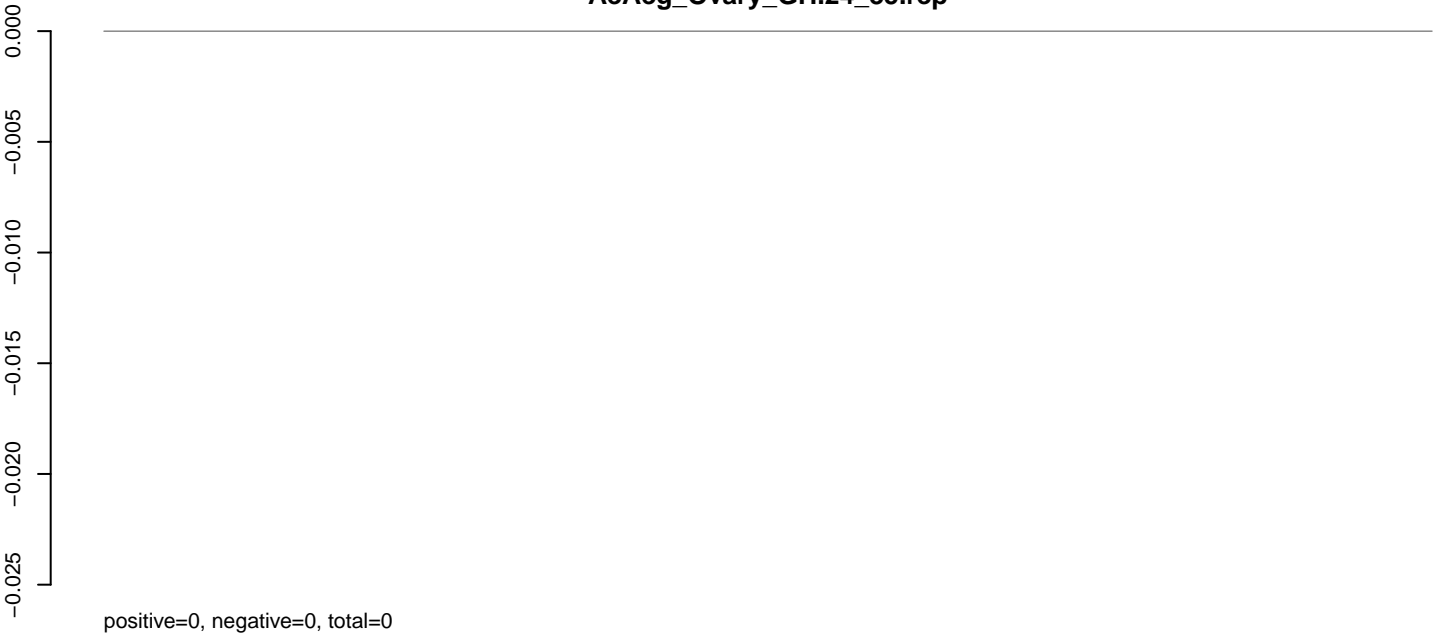
Window size=50, length=4176, Virus@X74945.1-Aedes_albopictus_densovirus_2_DNA:1-4176

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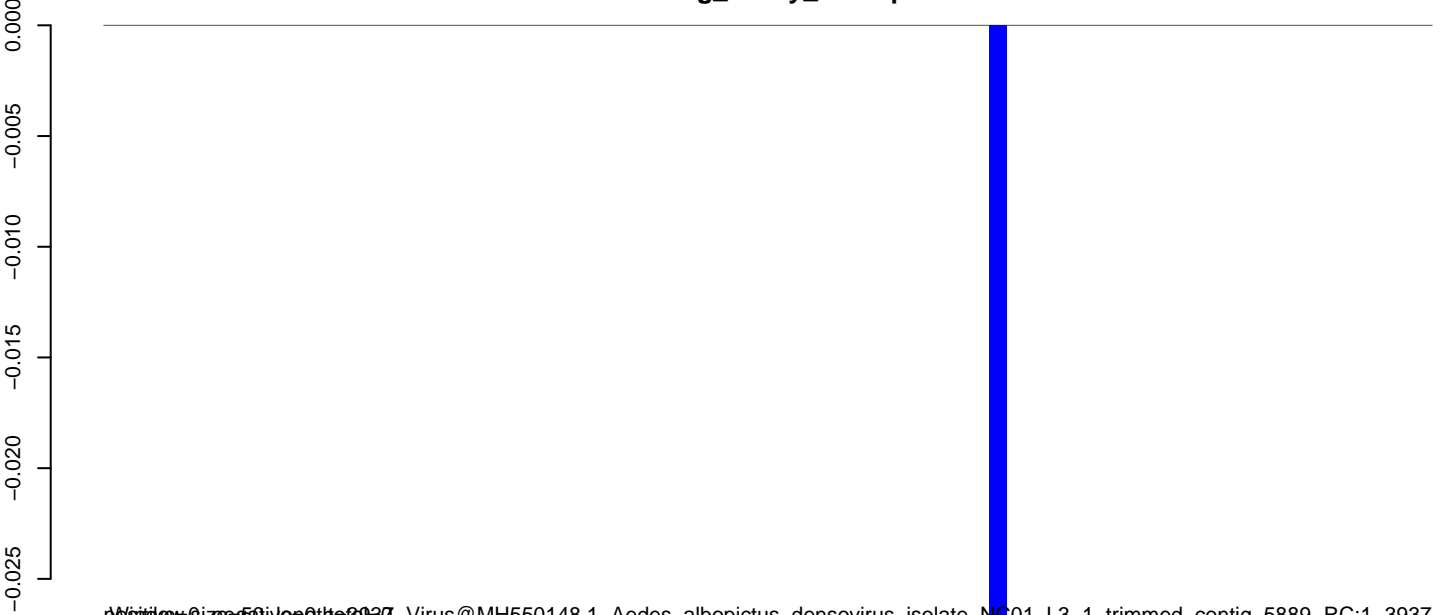
AeAeg_Ovary_GH.18_23.rep



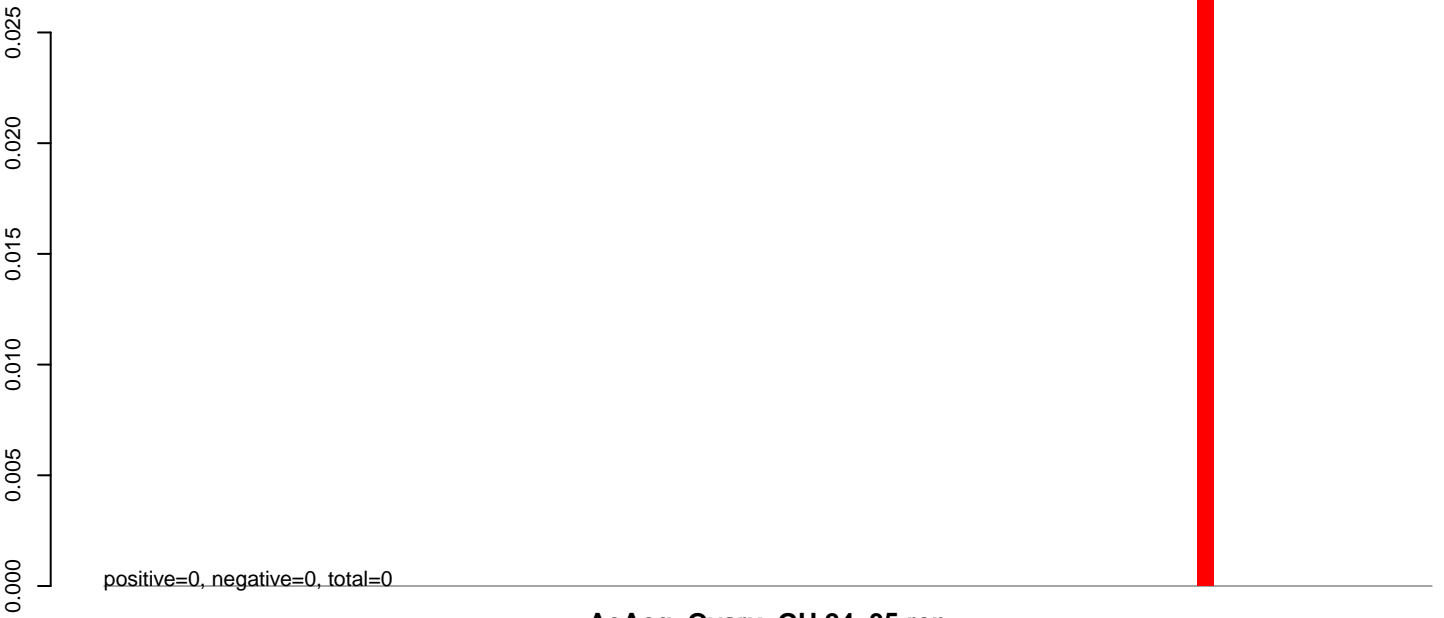
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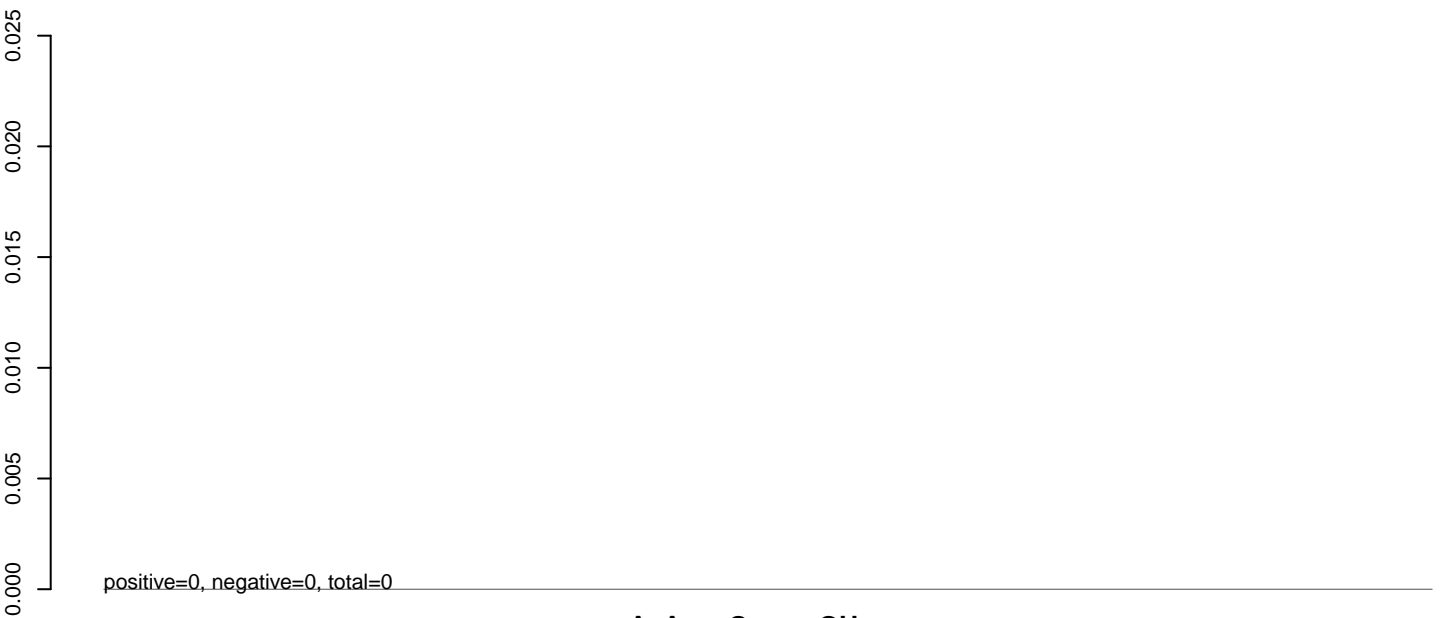
AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



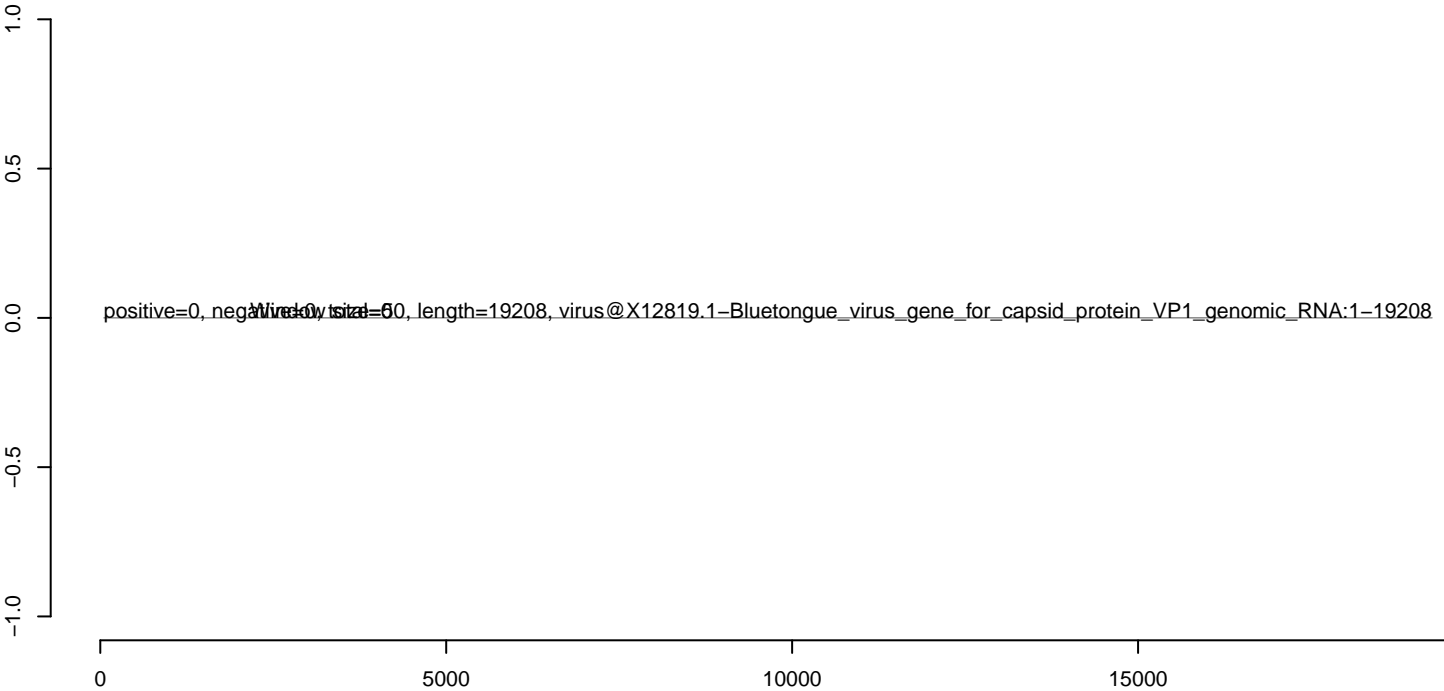
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



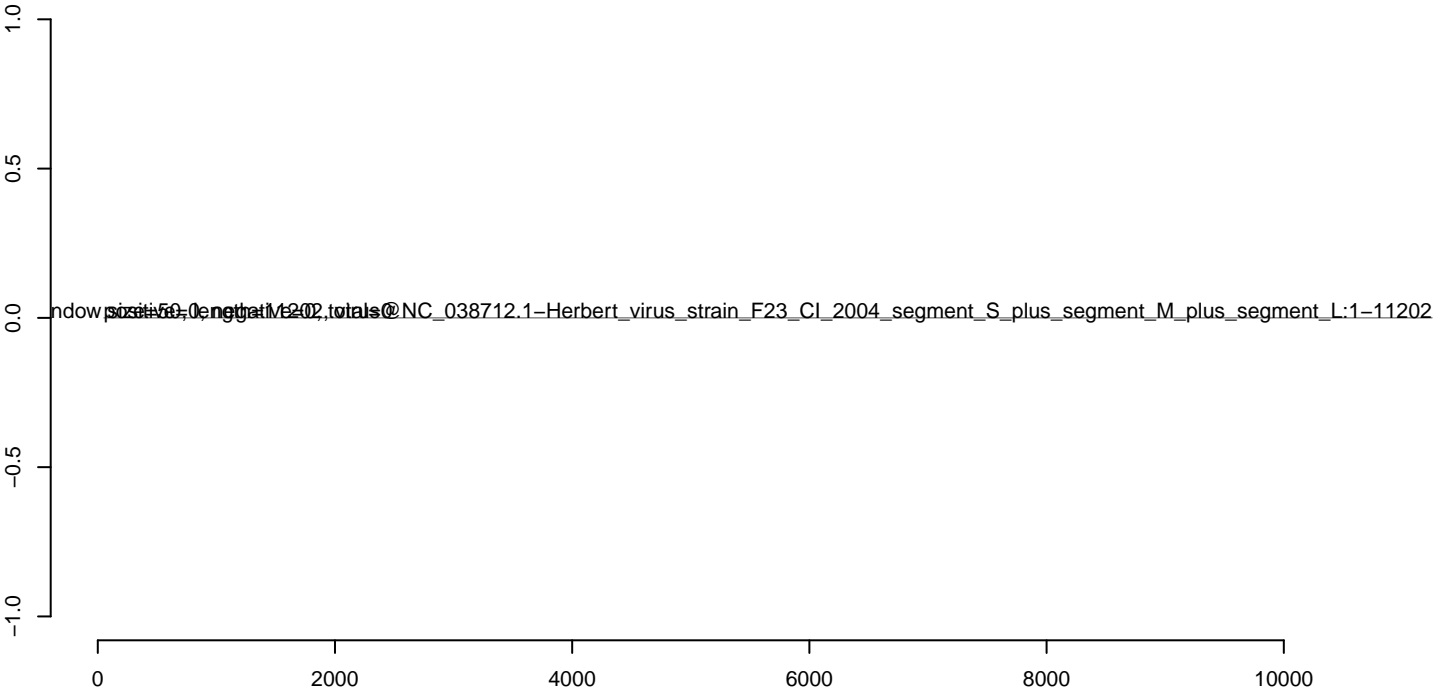
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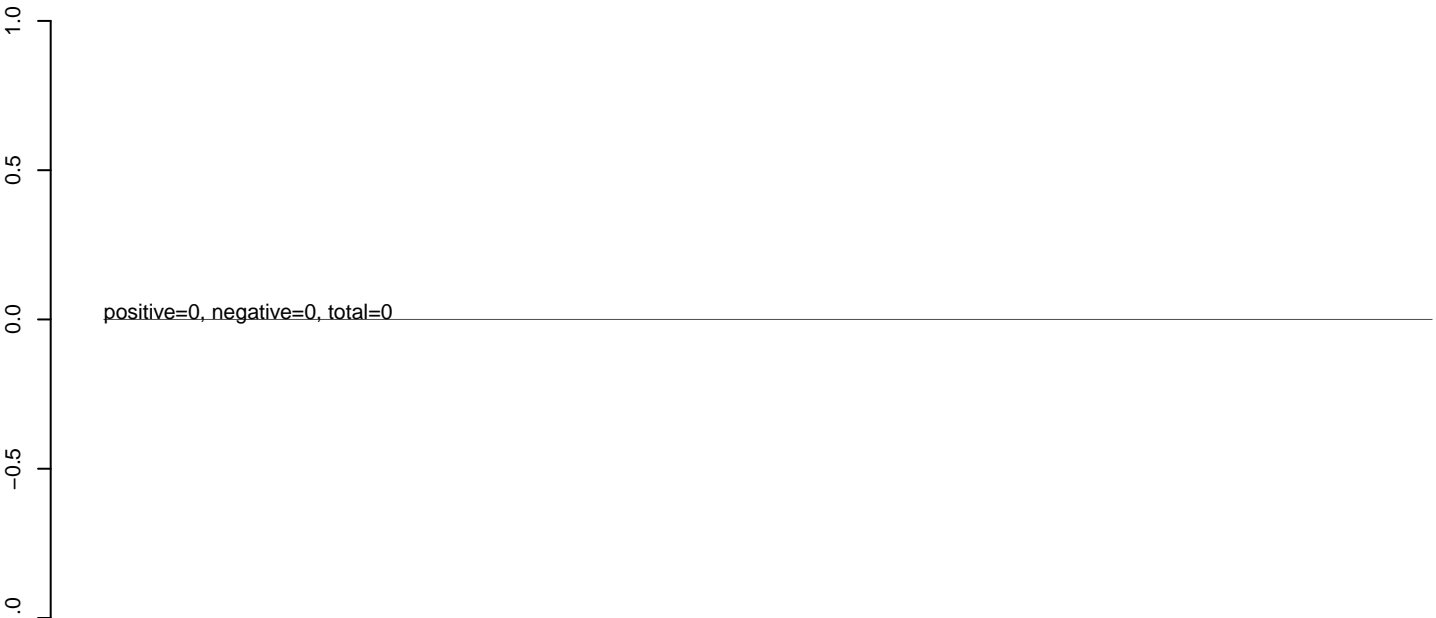
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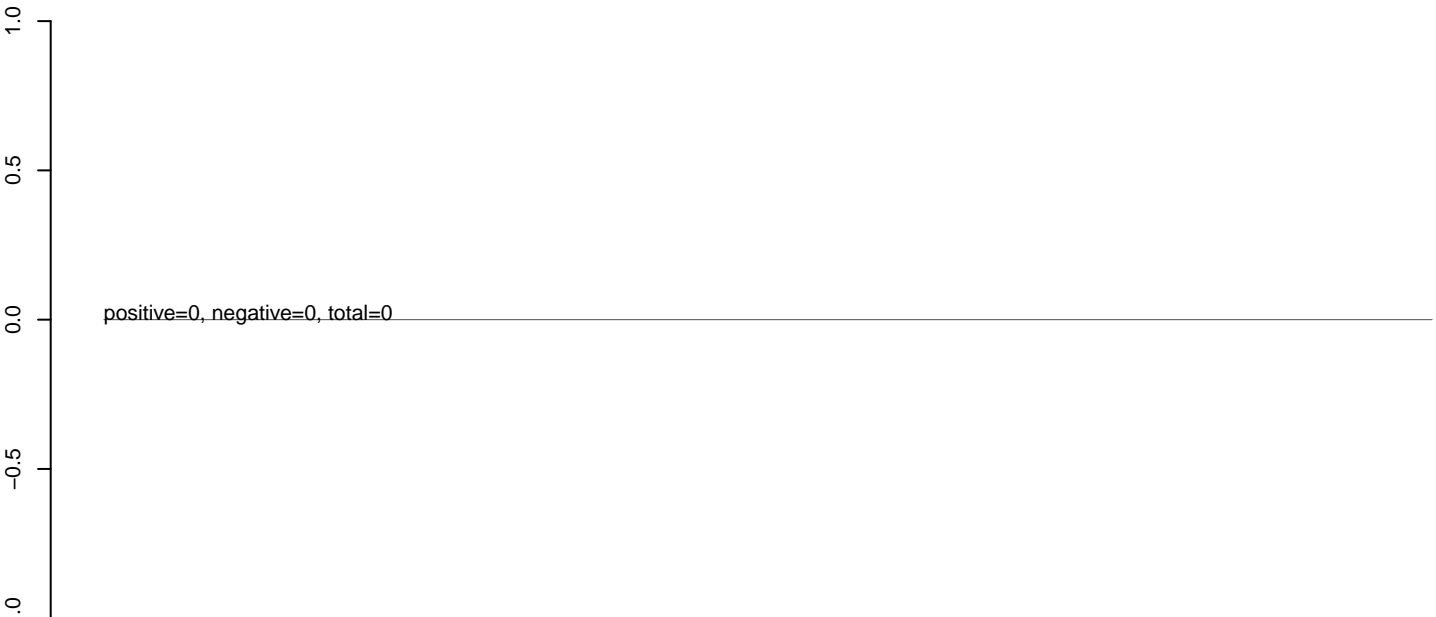
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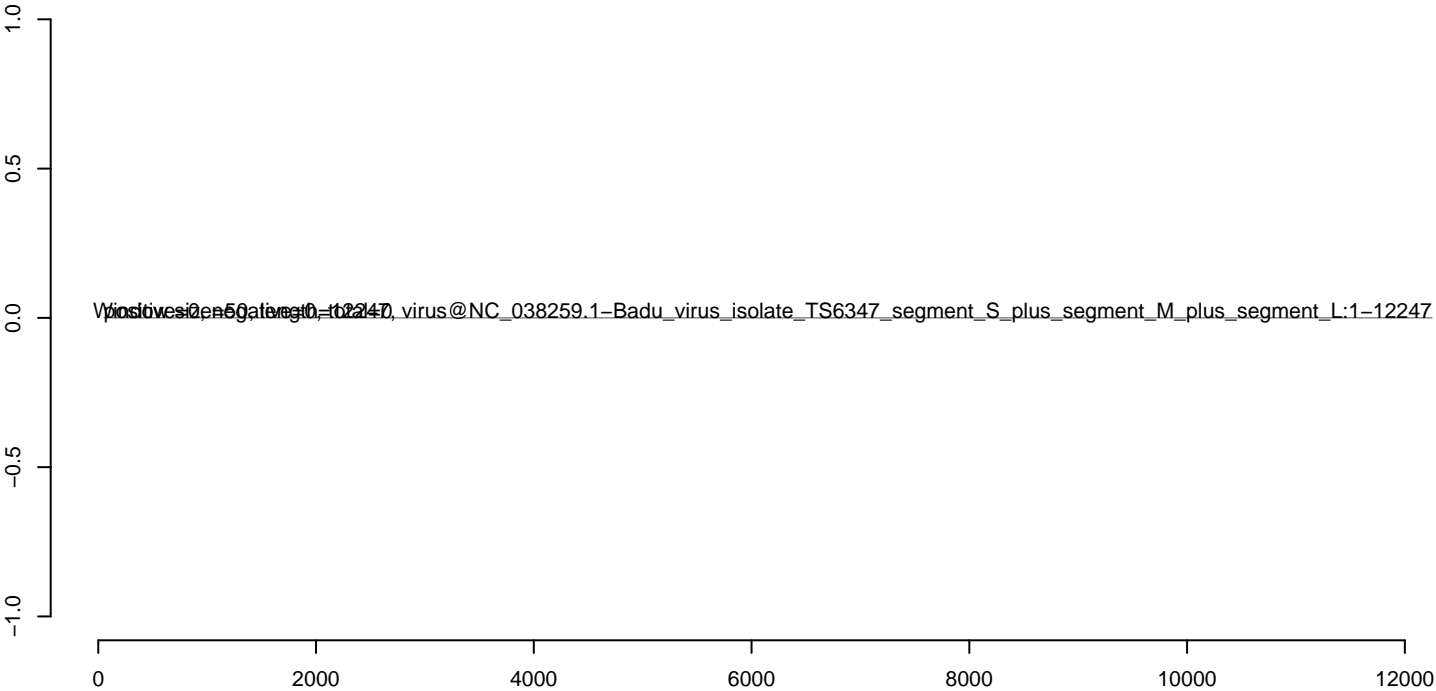
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



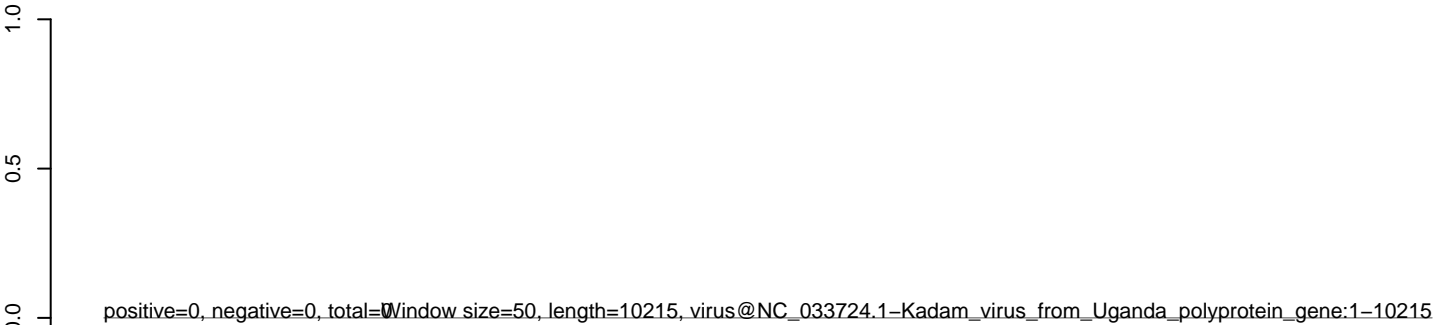
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



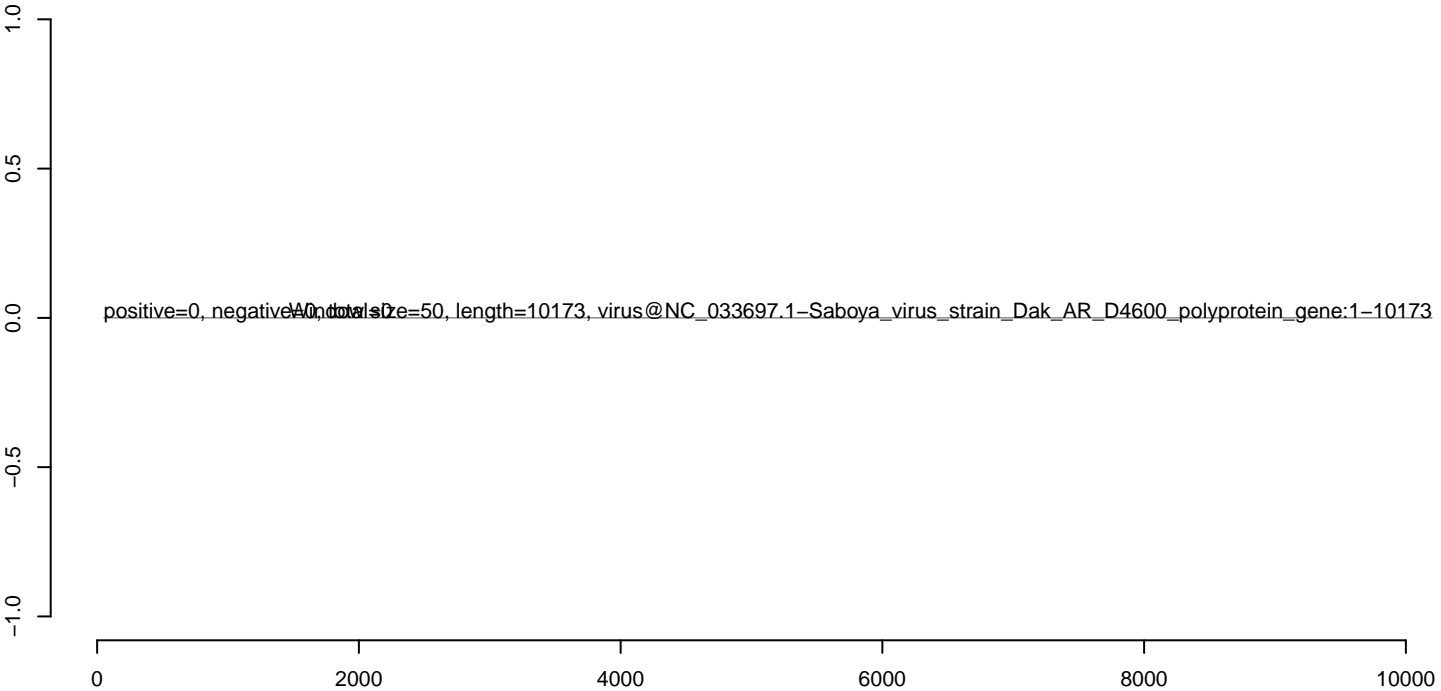
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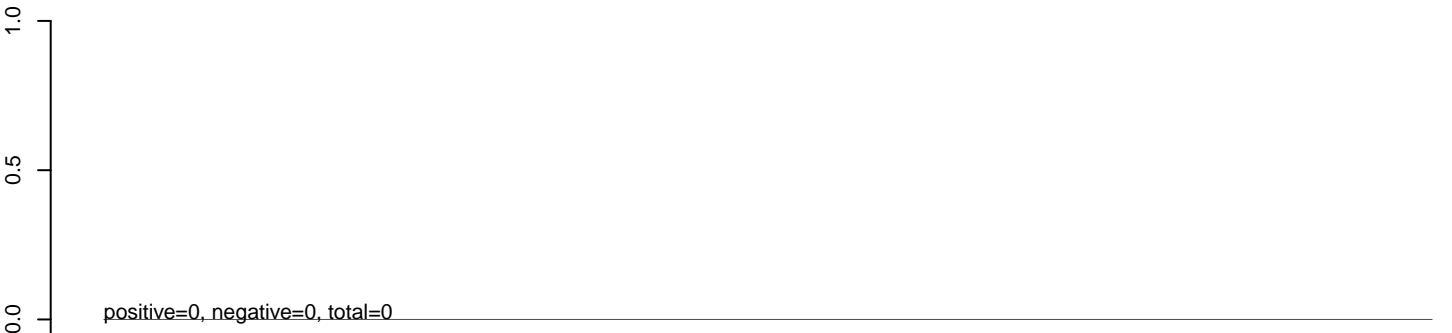
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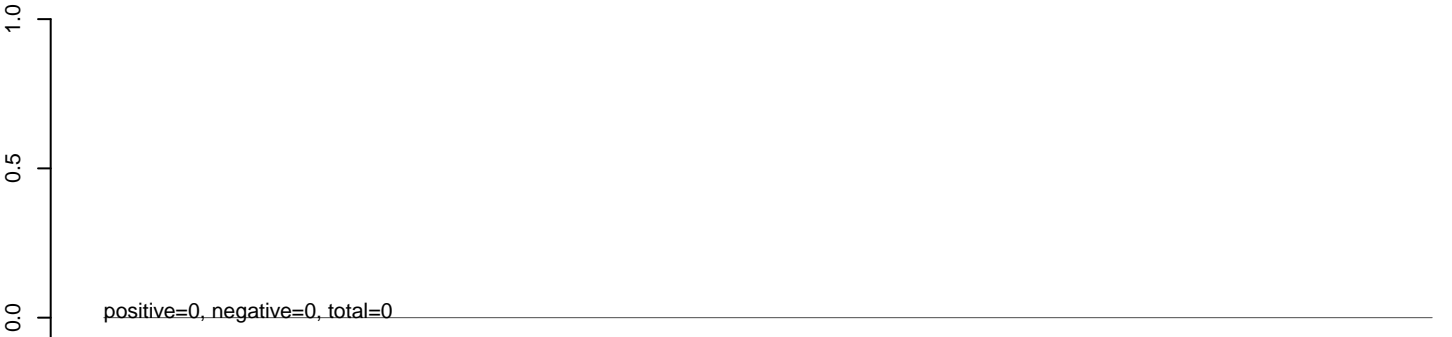
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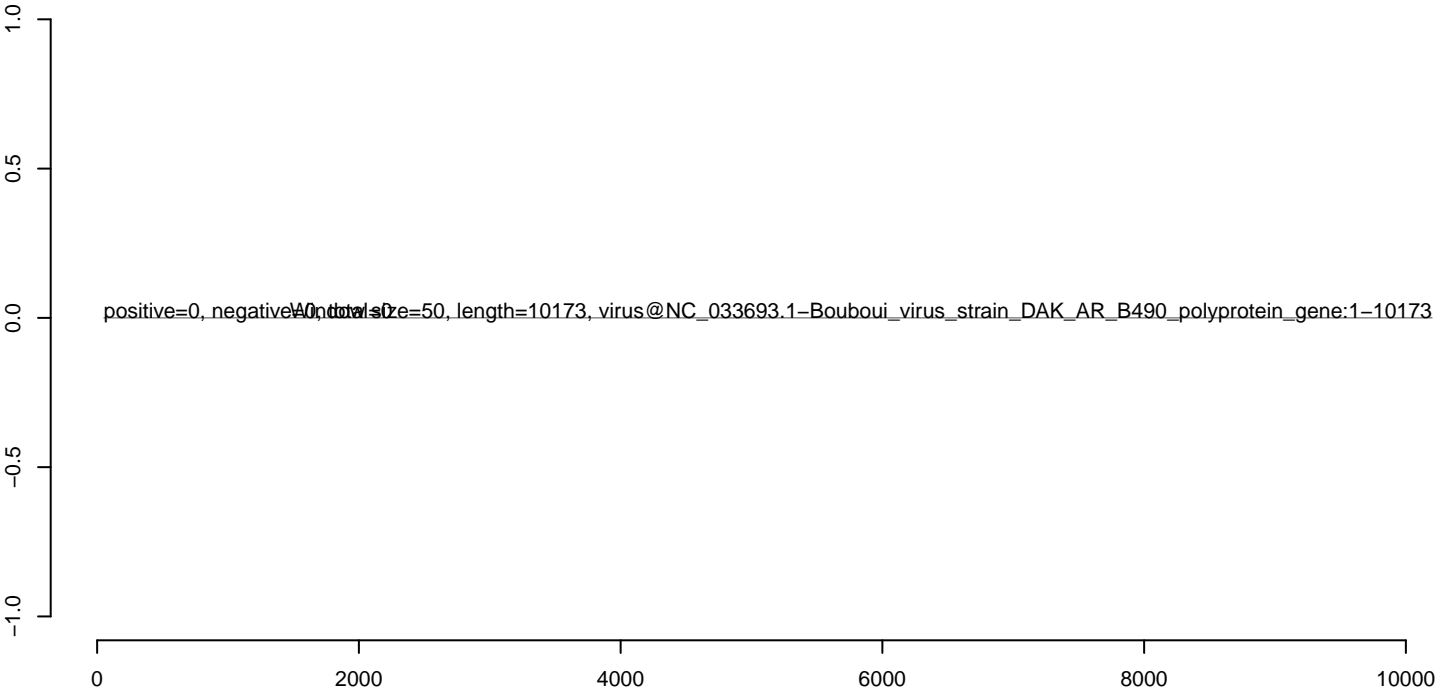
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



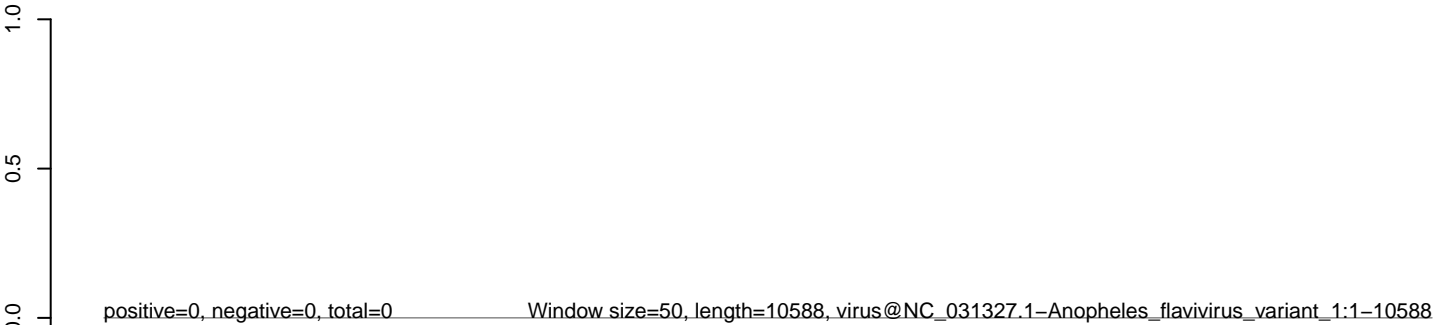
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AeAeg_Ovary_GH.24_35.rep

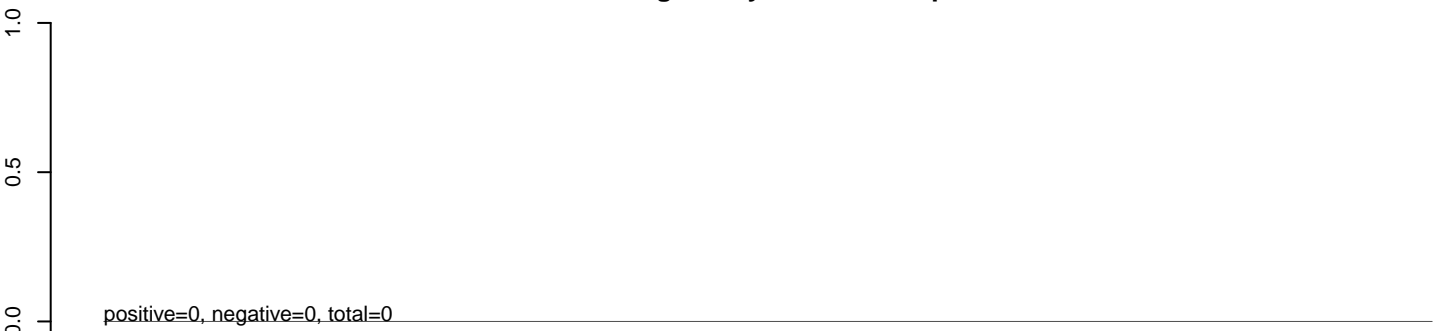


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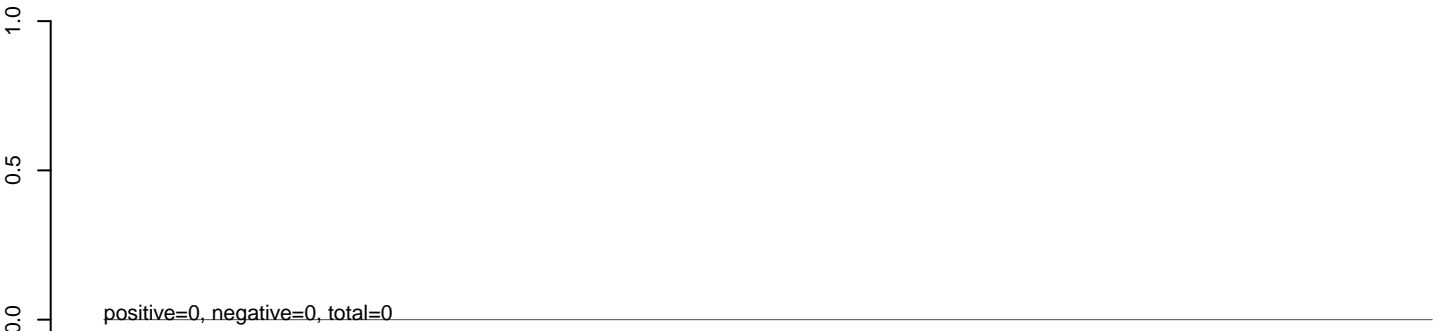


0 2000 4000 6000 8000 10000

AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep

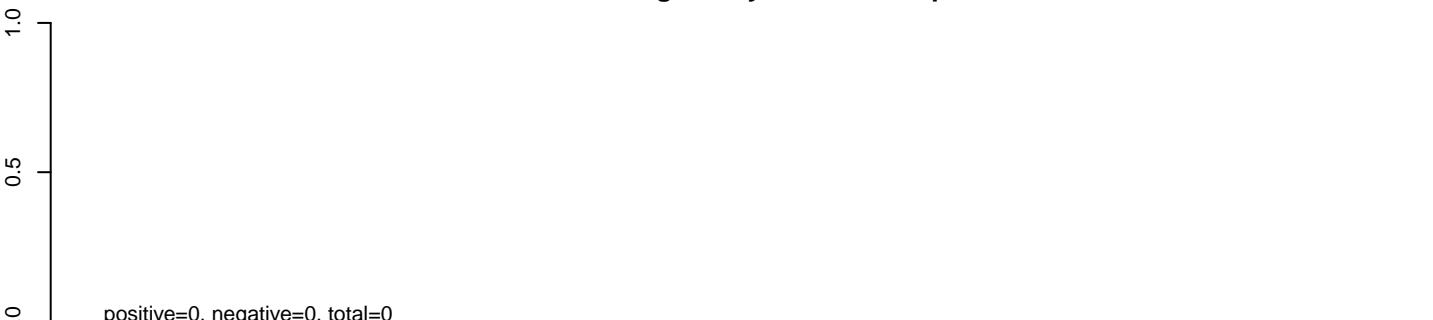


AeAeg_Ovary_GH.rep

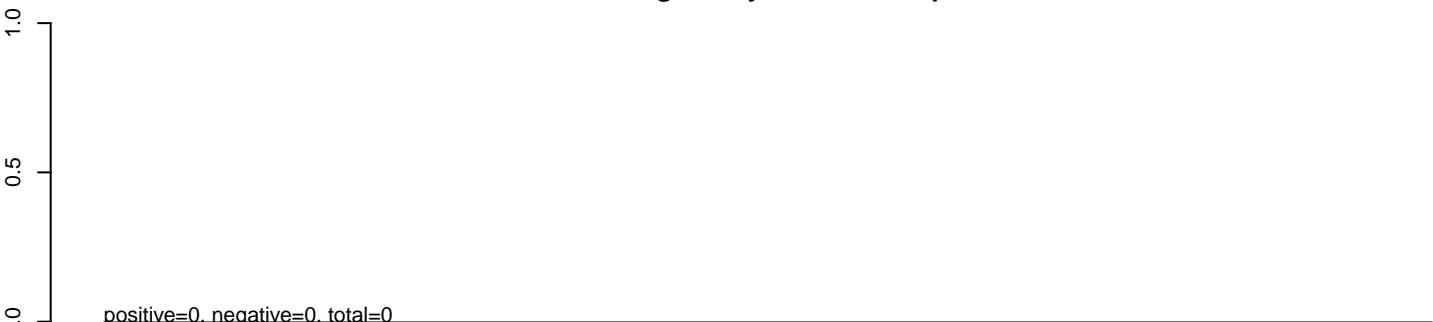


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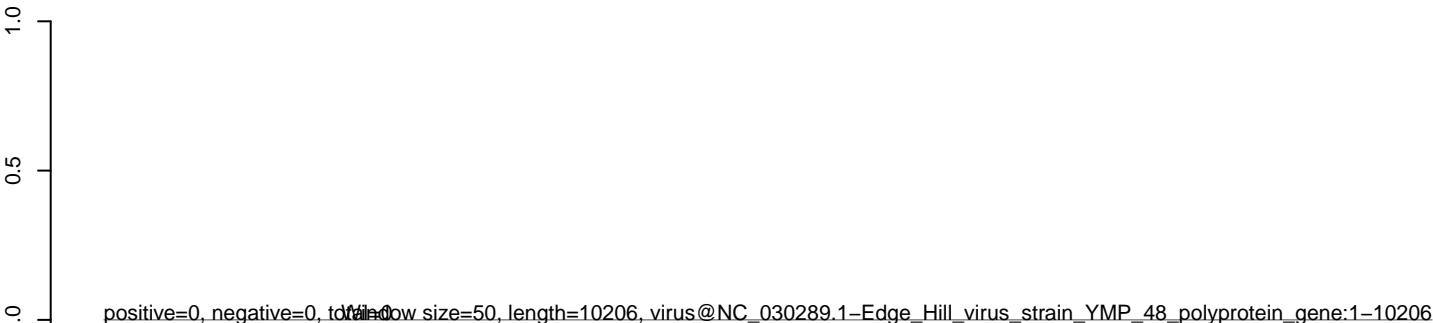
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

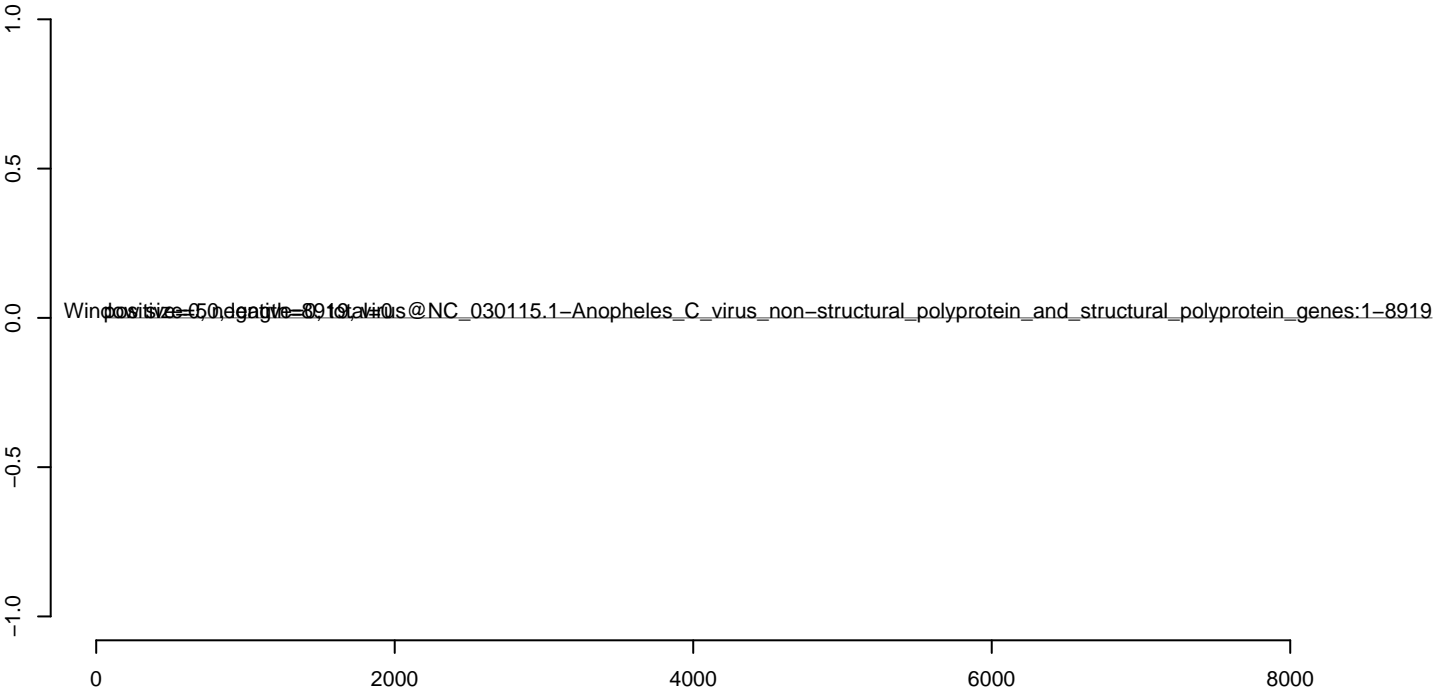
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



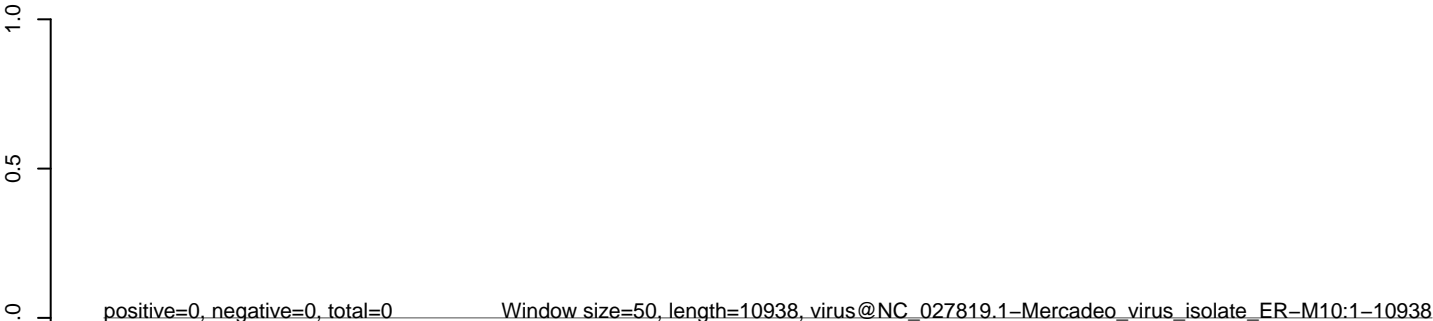
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

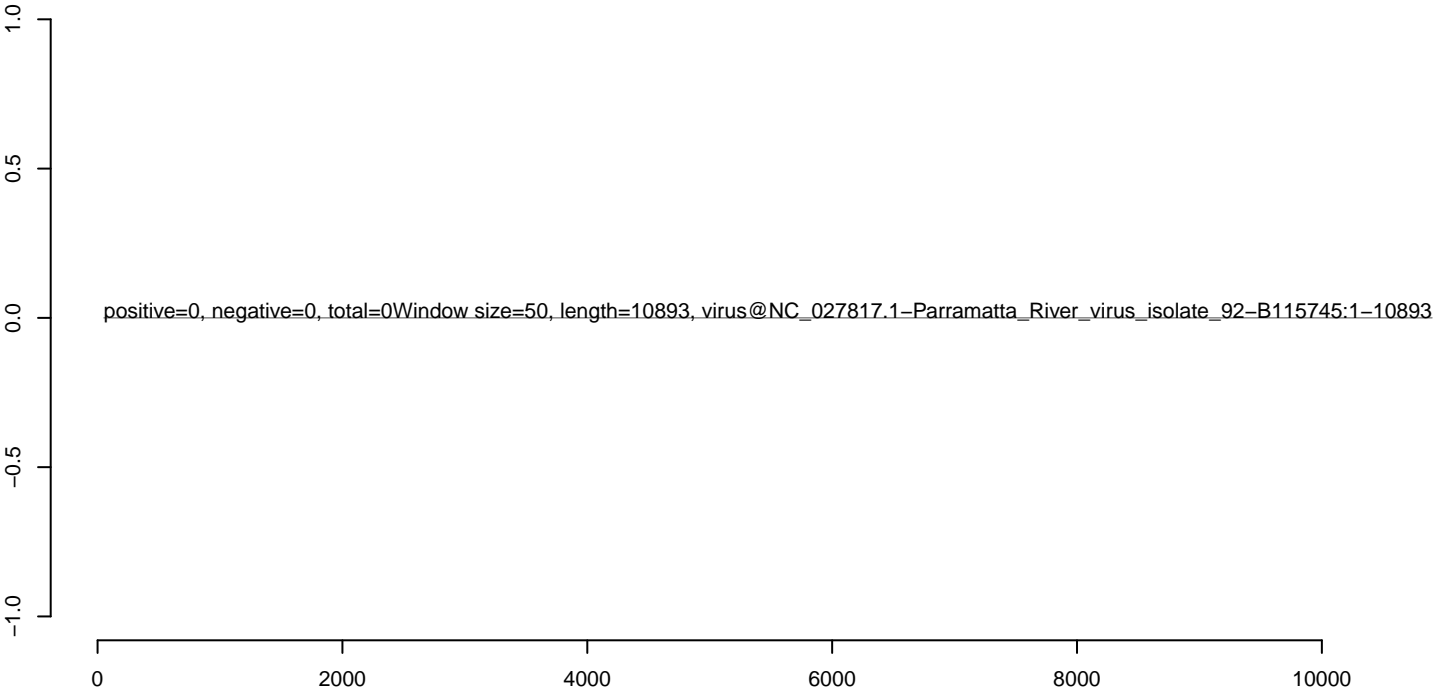
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



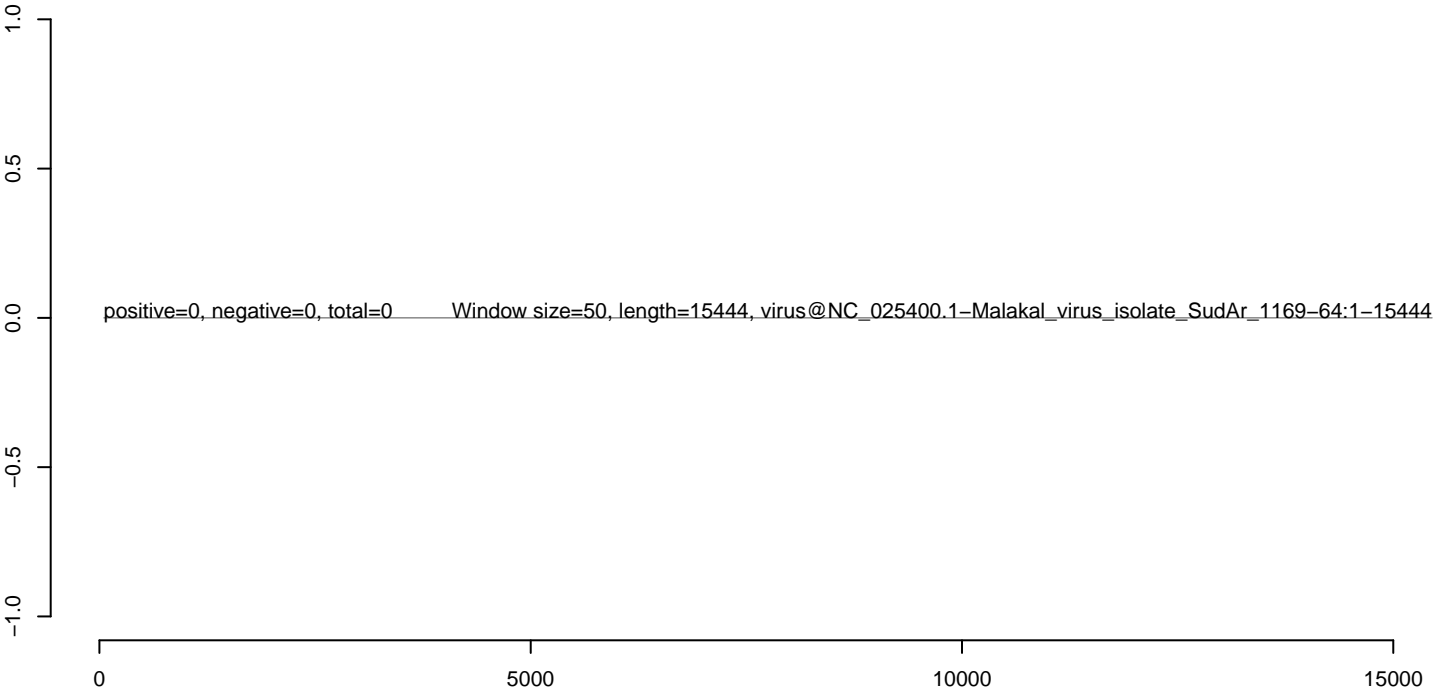
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



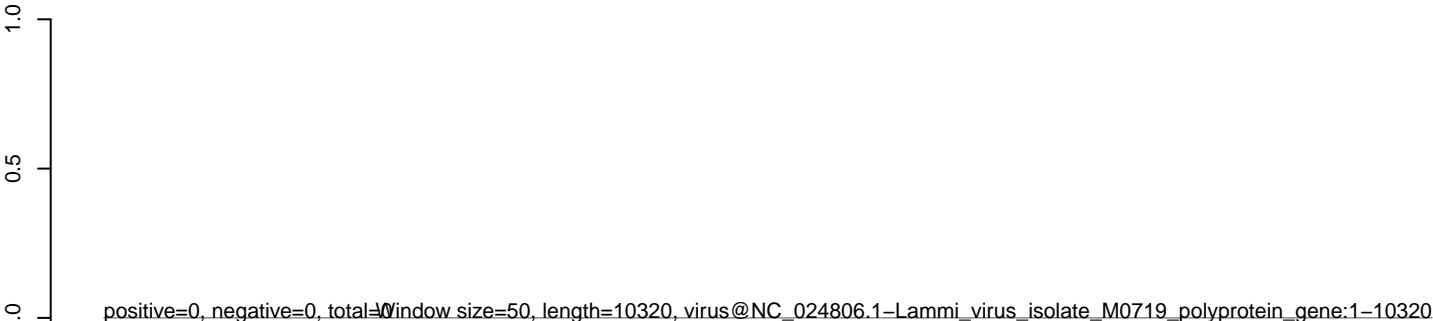
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

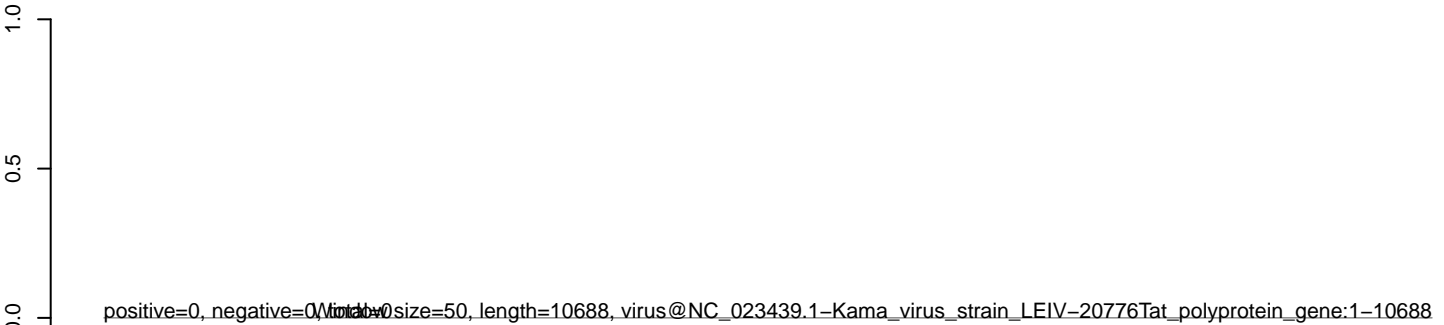
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

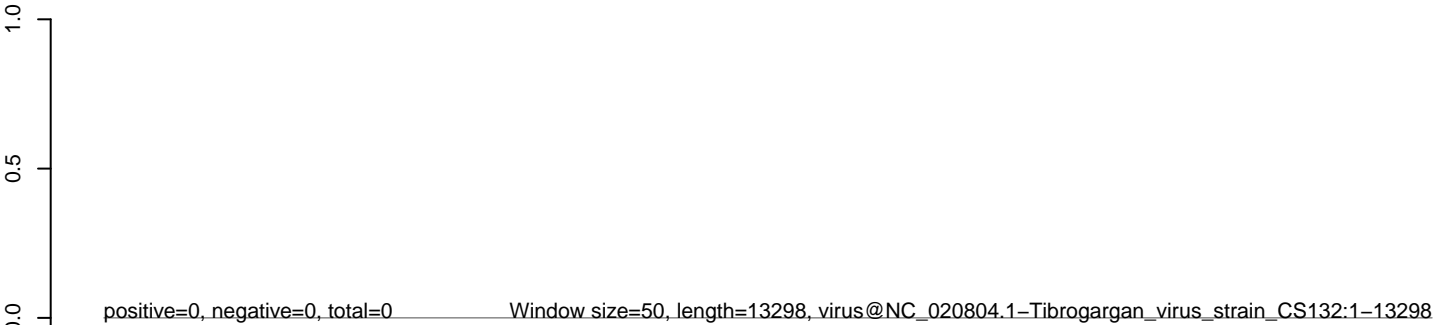
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000 12000

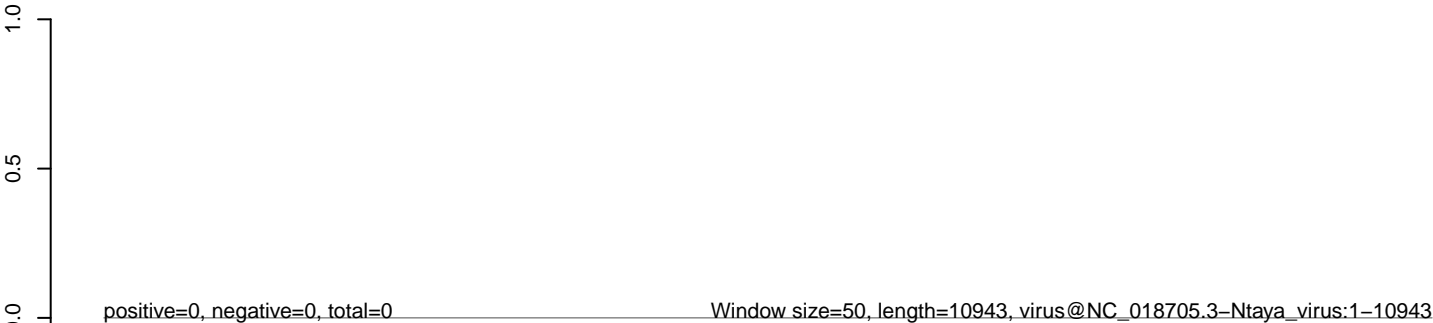
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

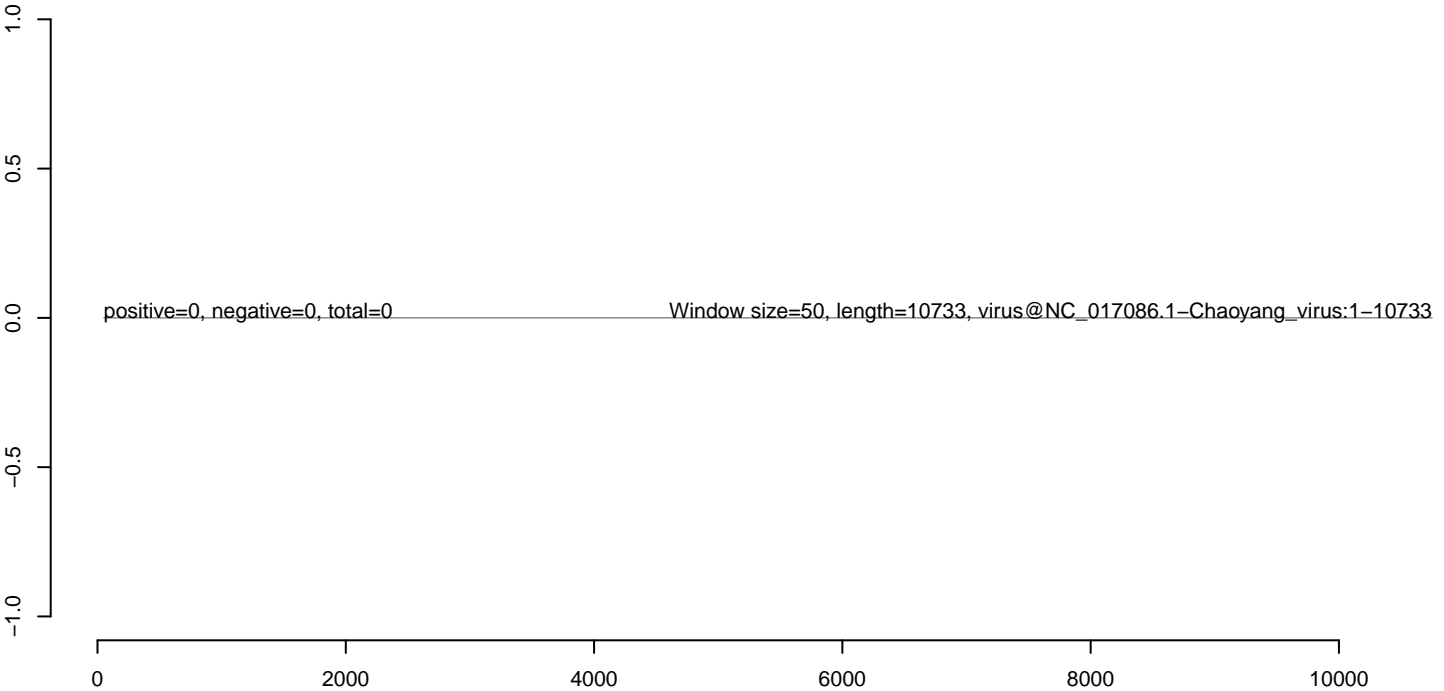
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



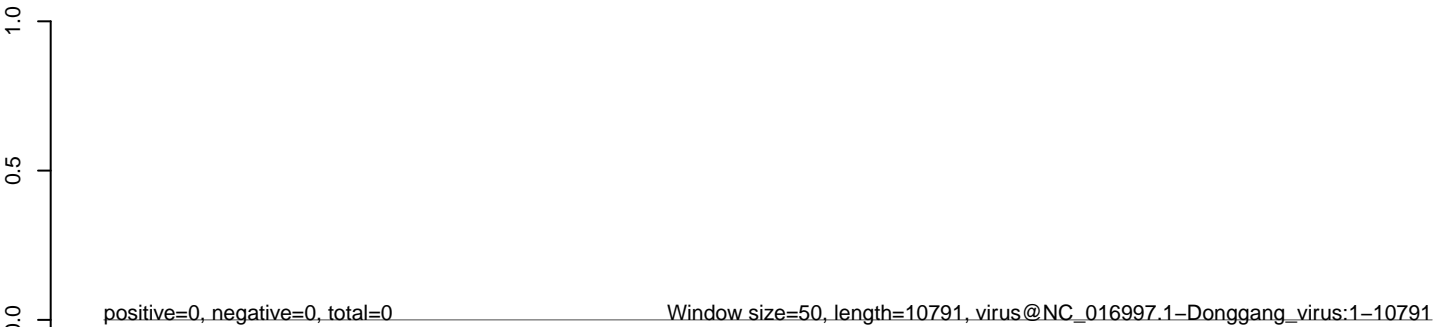
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

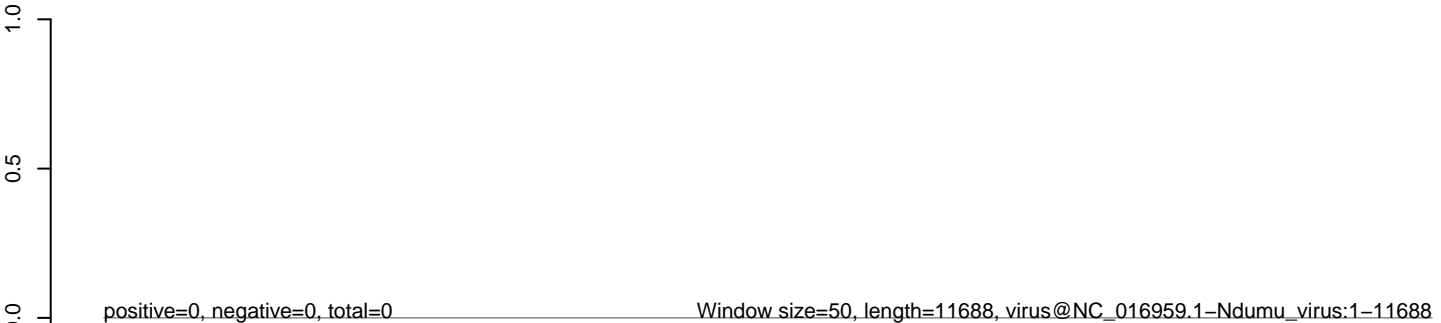
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000 12000

AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

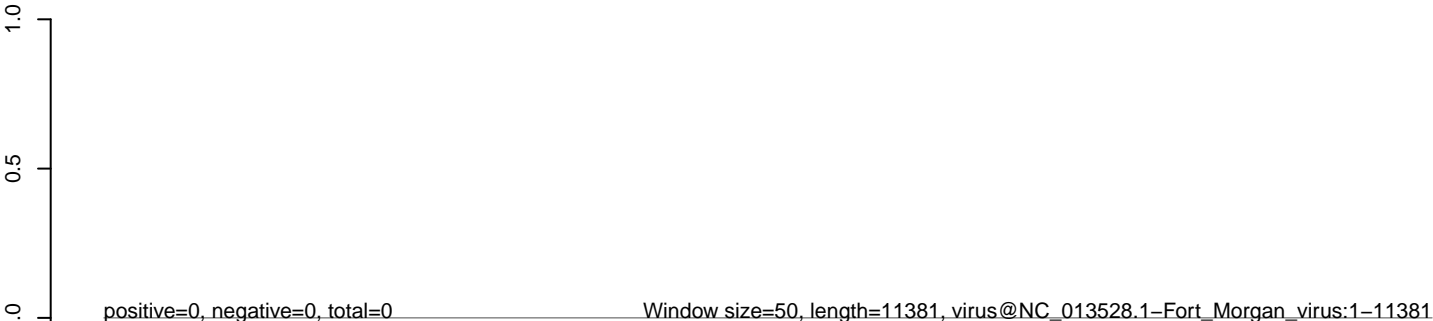
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

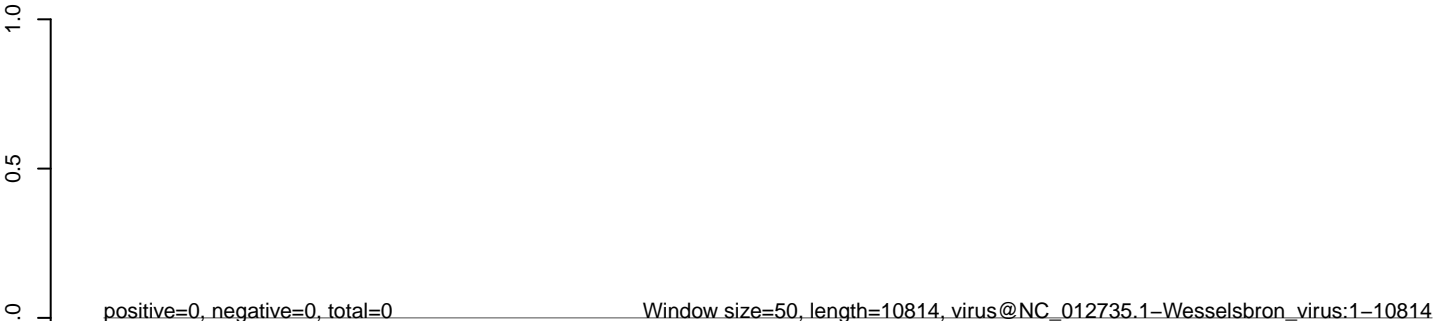
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

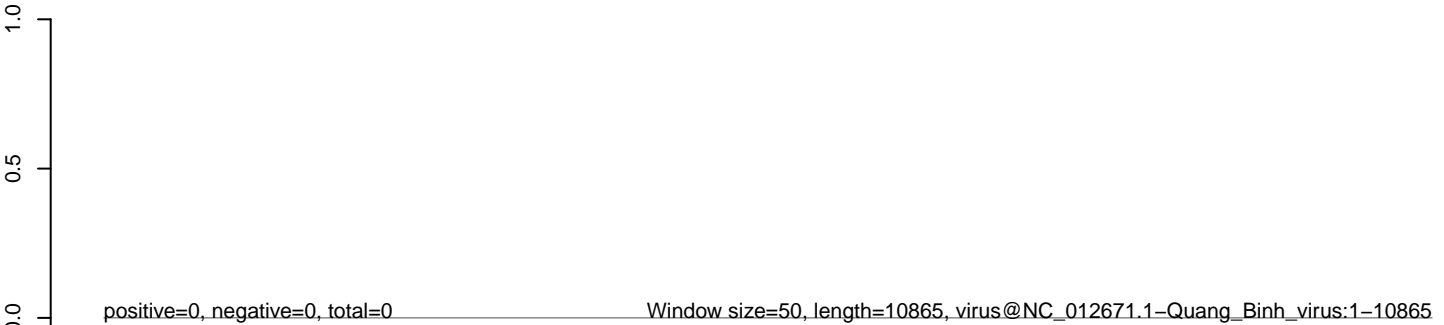
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

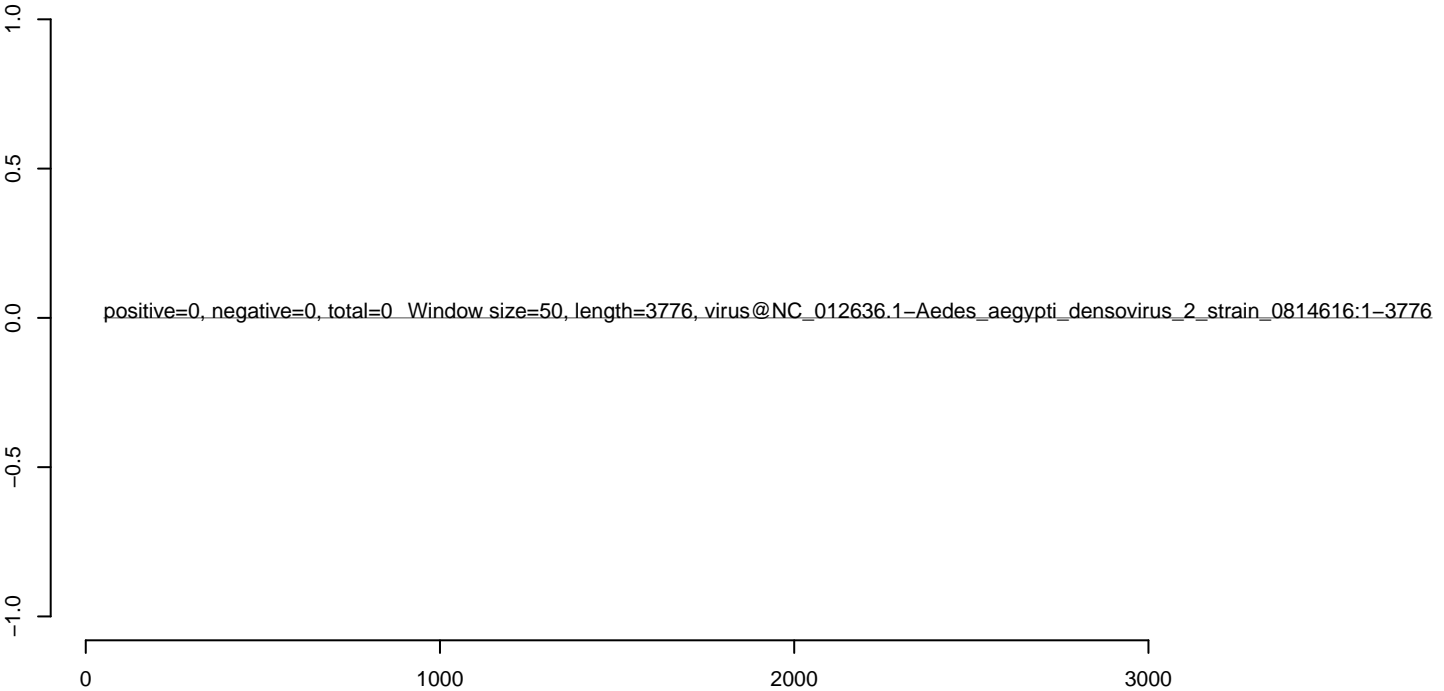
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



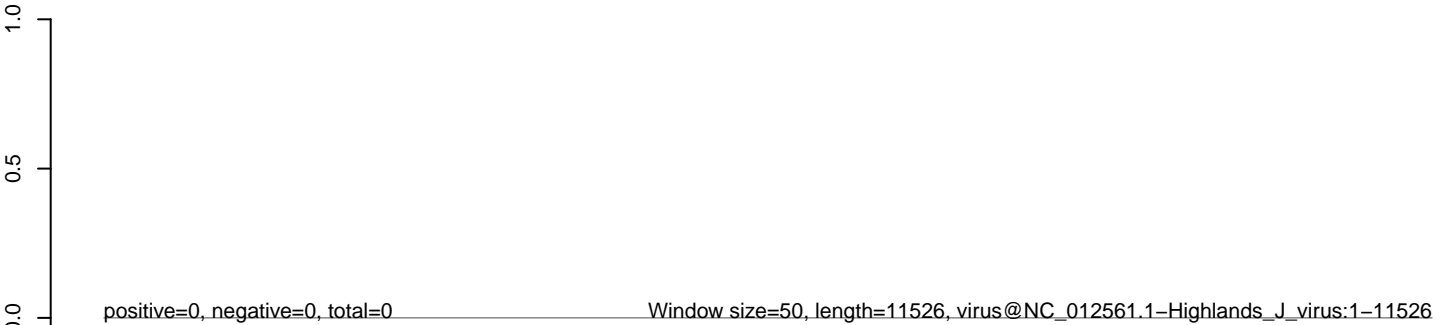
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000 12000

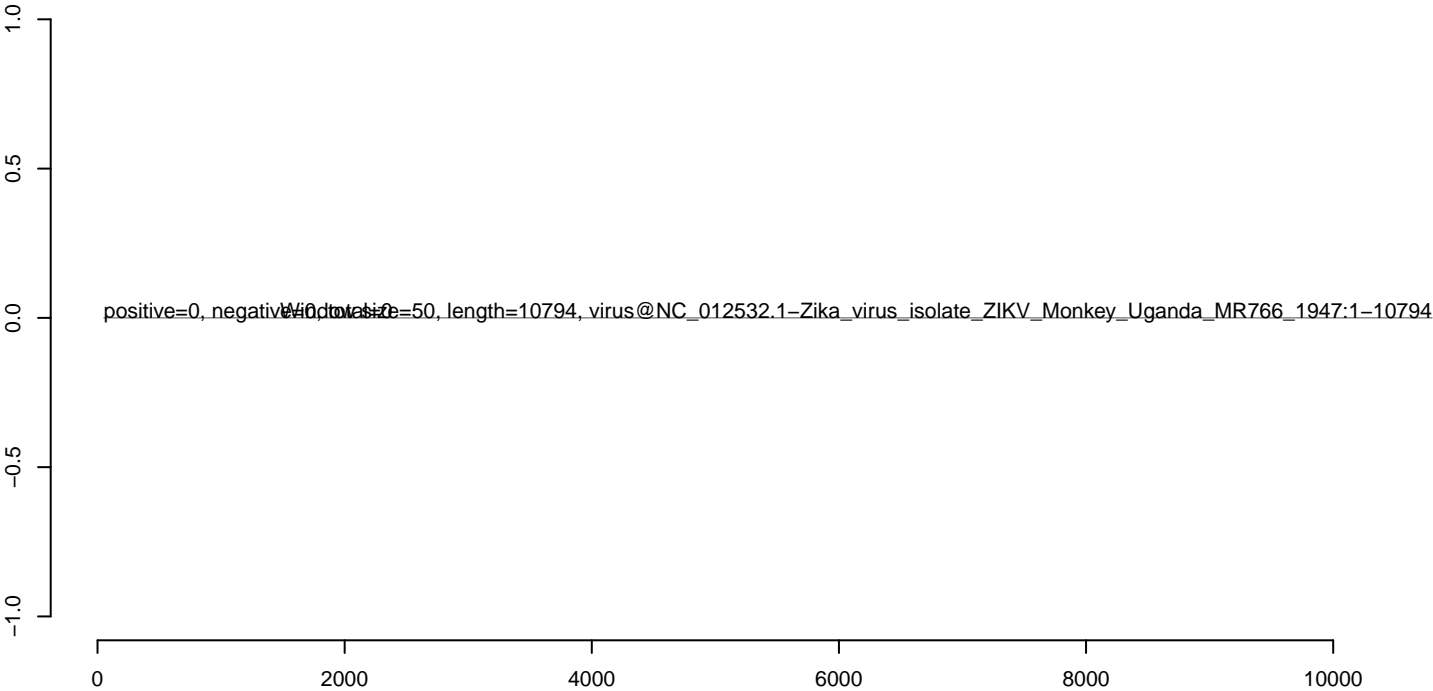
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



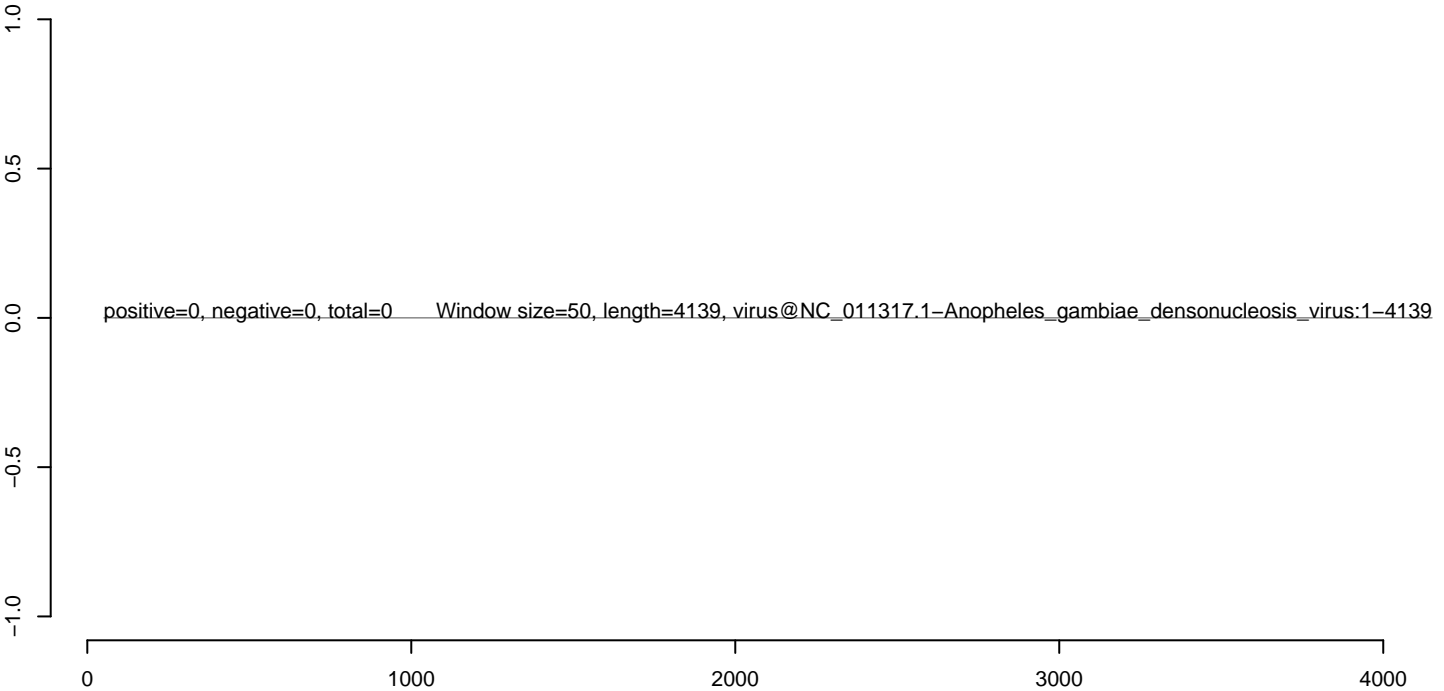
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



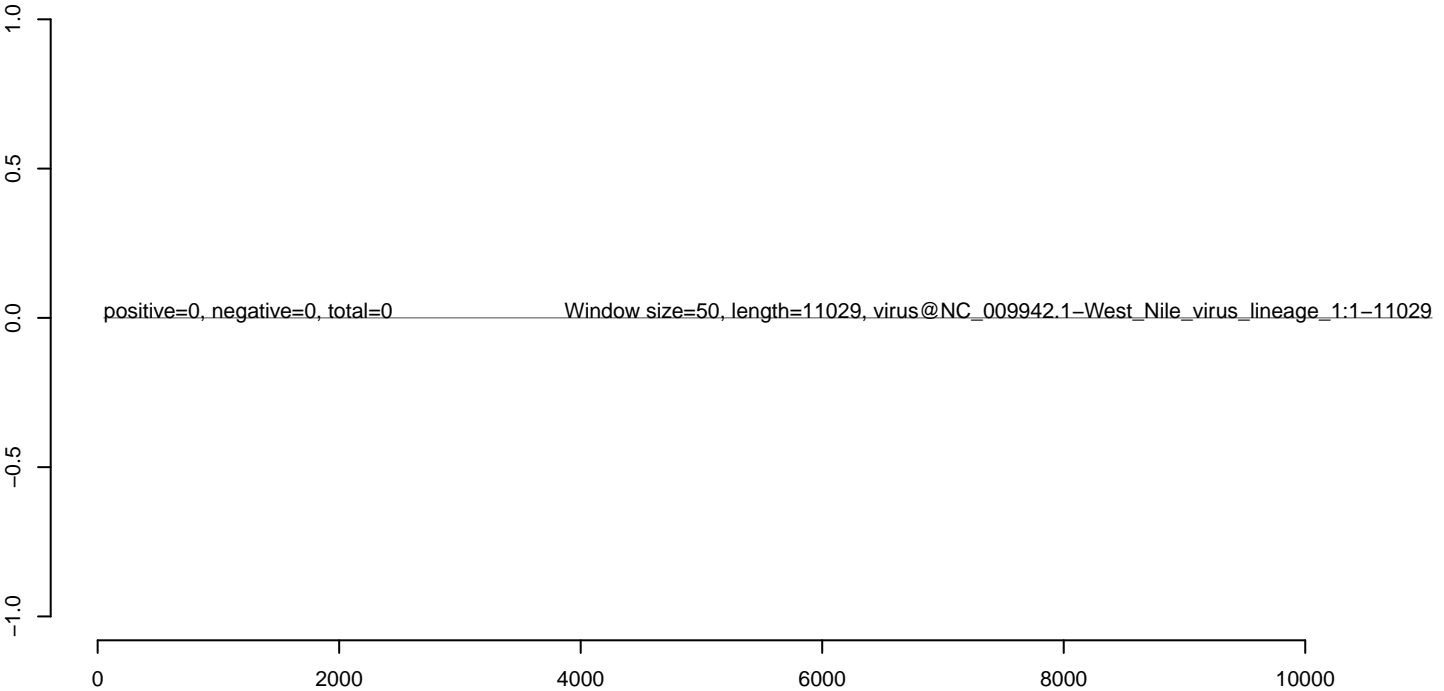
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



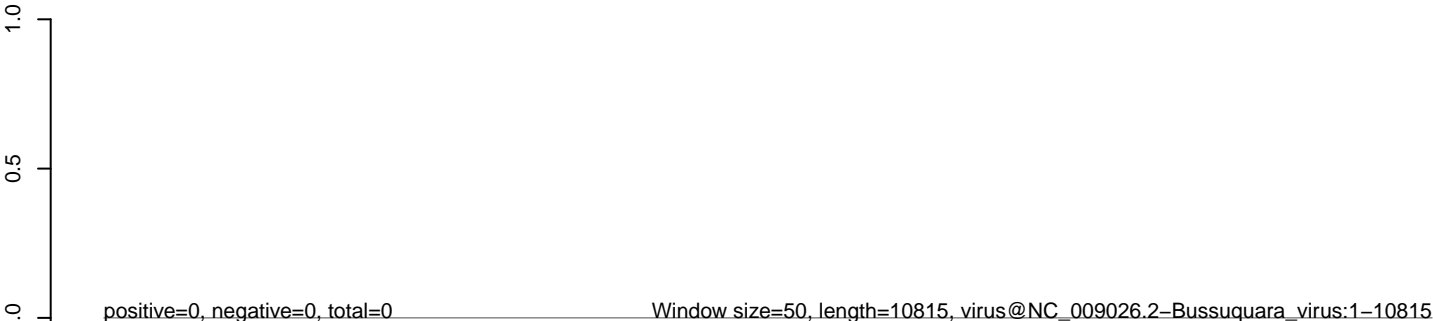
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

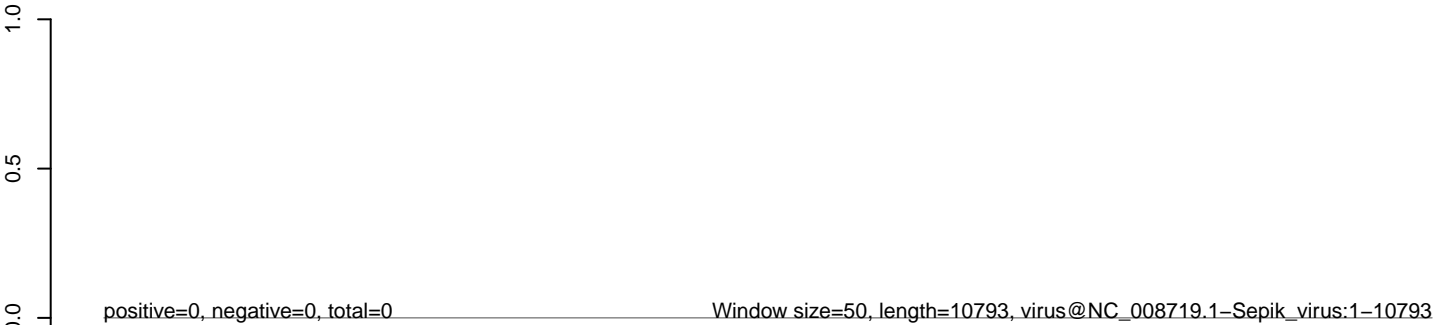
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

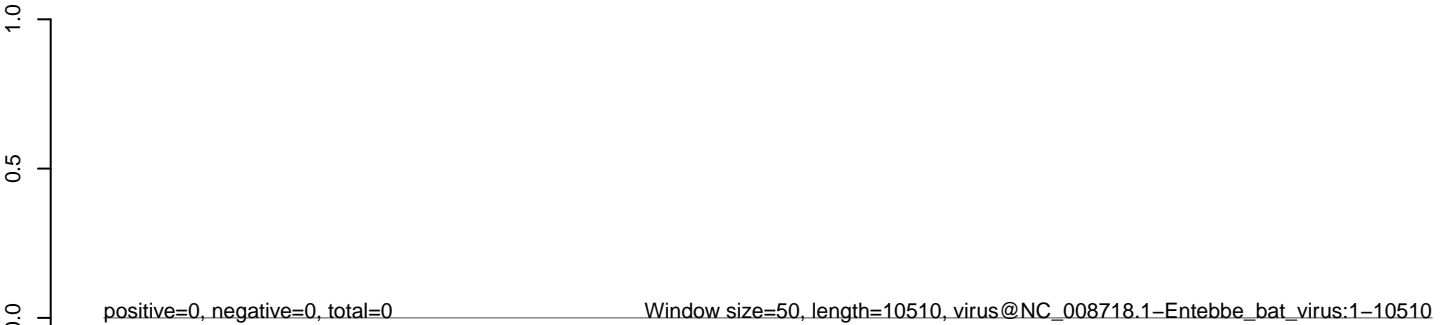
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

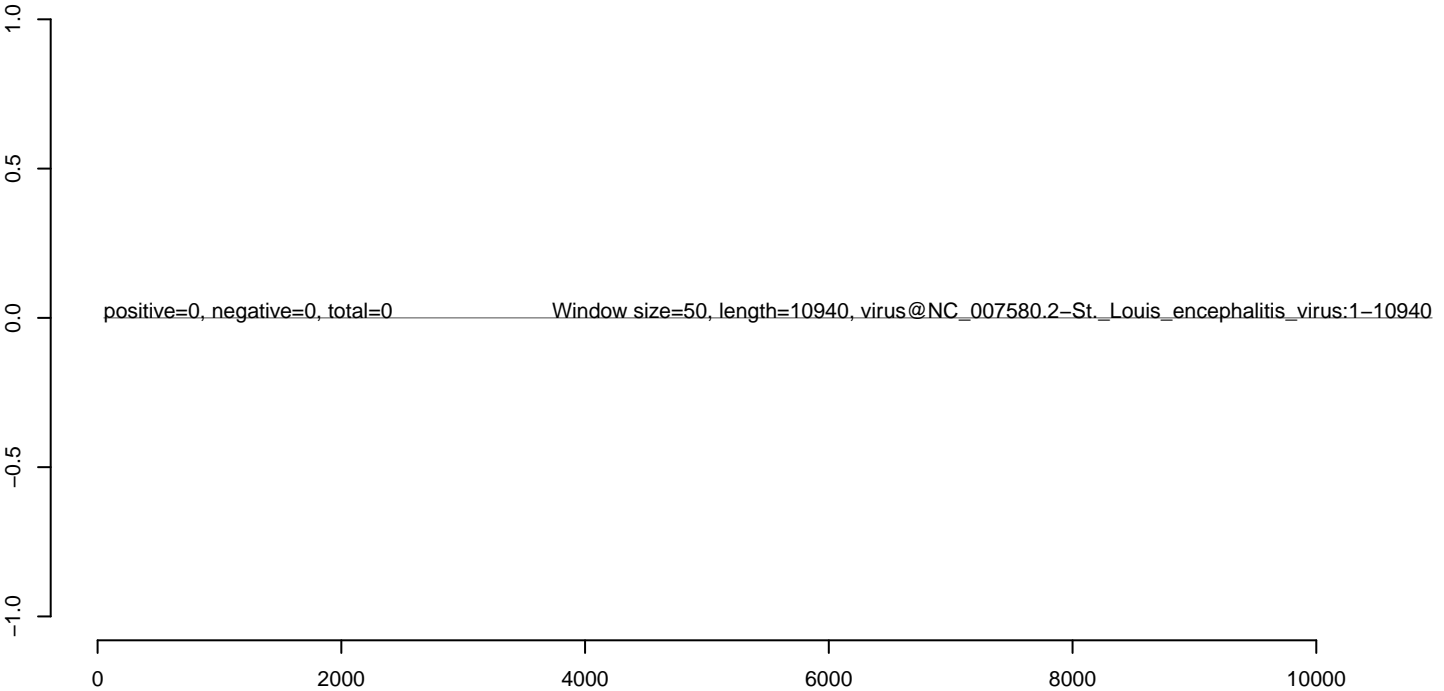
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AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



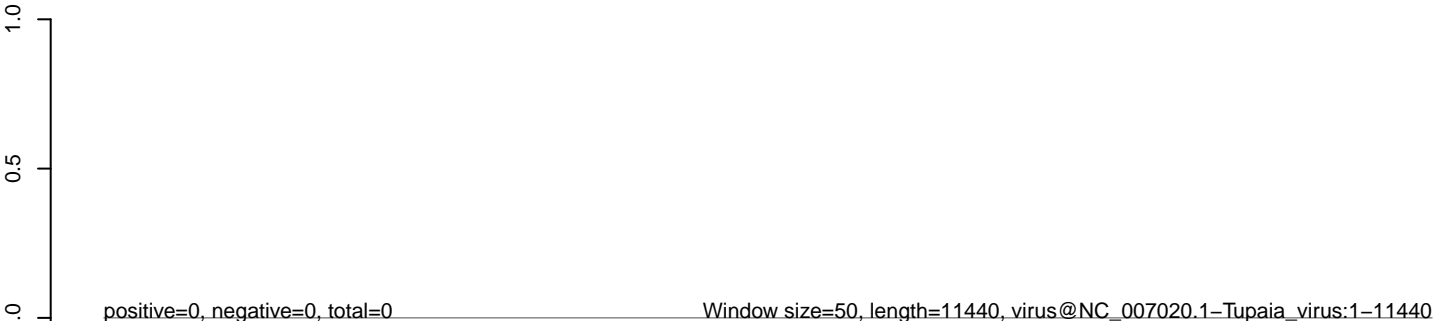
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

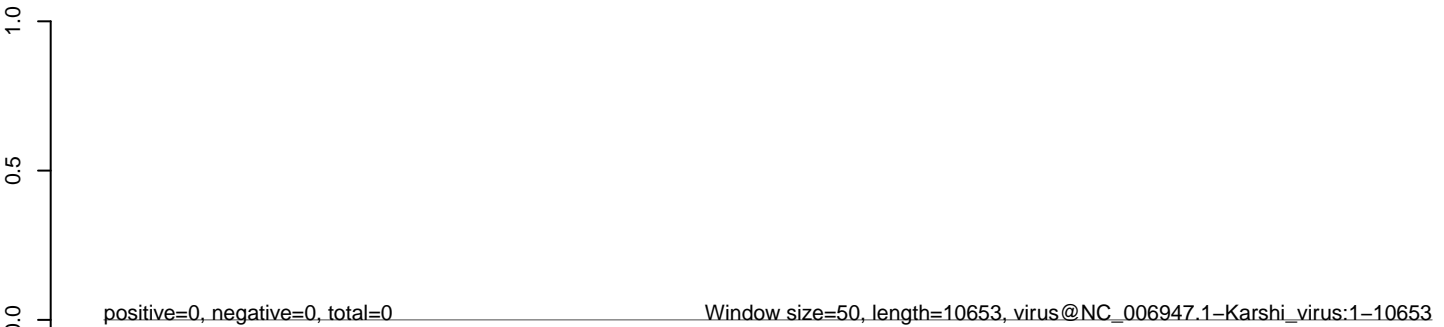
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000

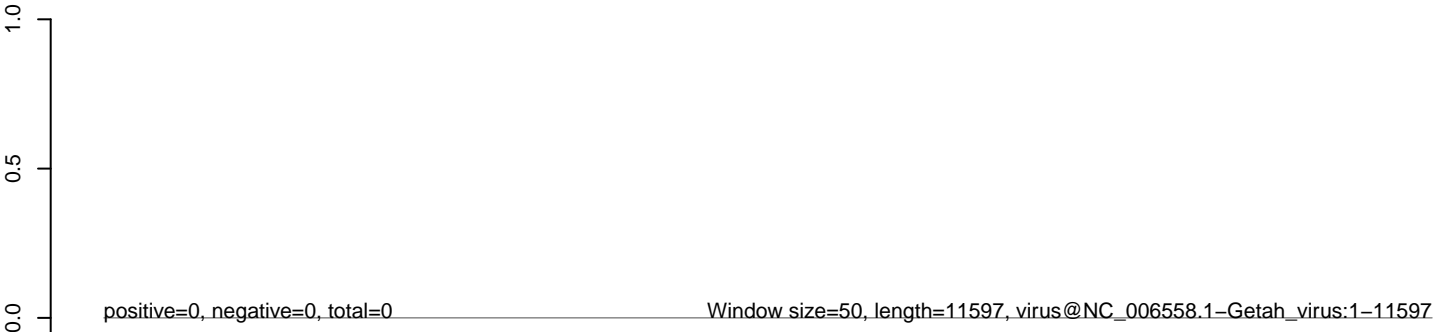
AeAeg_Ovary_GH.18_23.rep



AeAeg_Ovary_GH.24_35.rep



AeAeg_Ovary_GH.rep



0 2000 4000 6000 8000 10000 12000

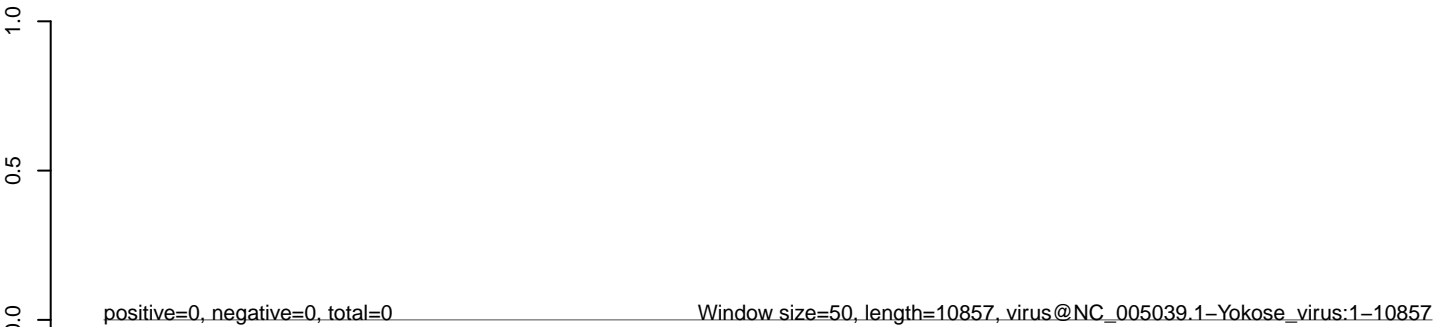
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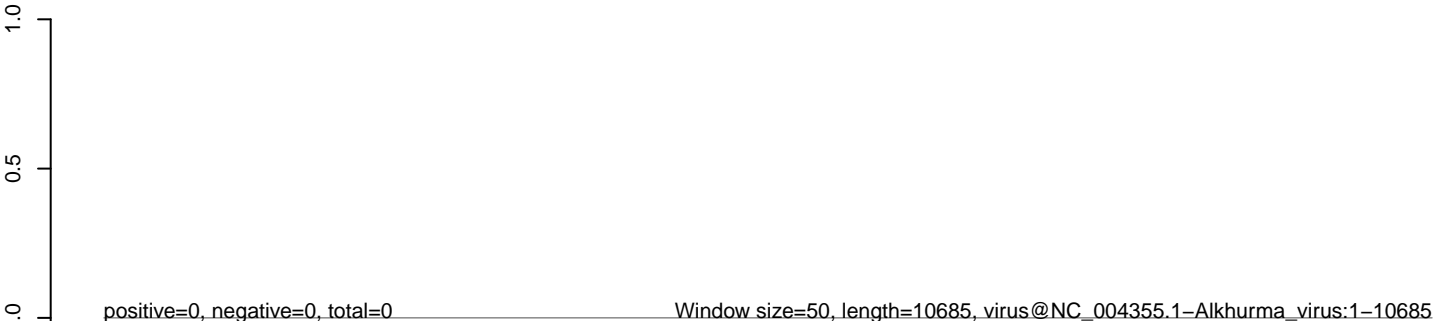
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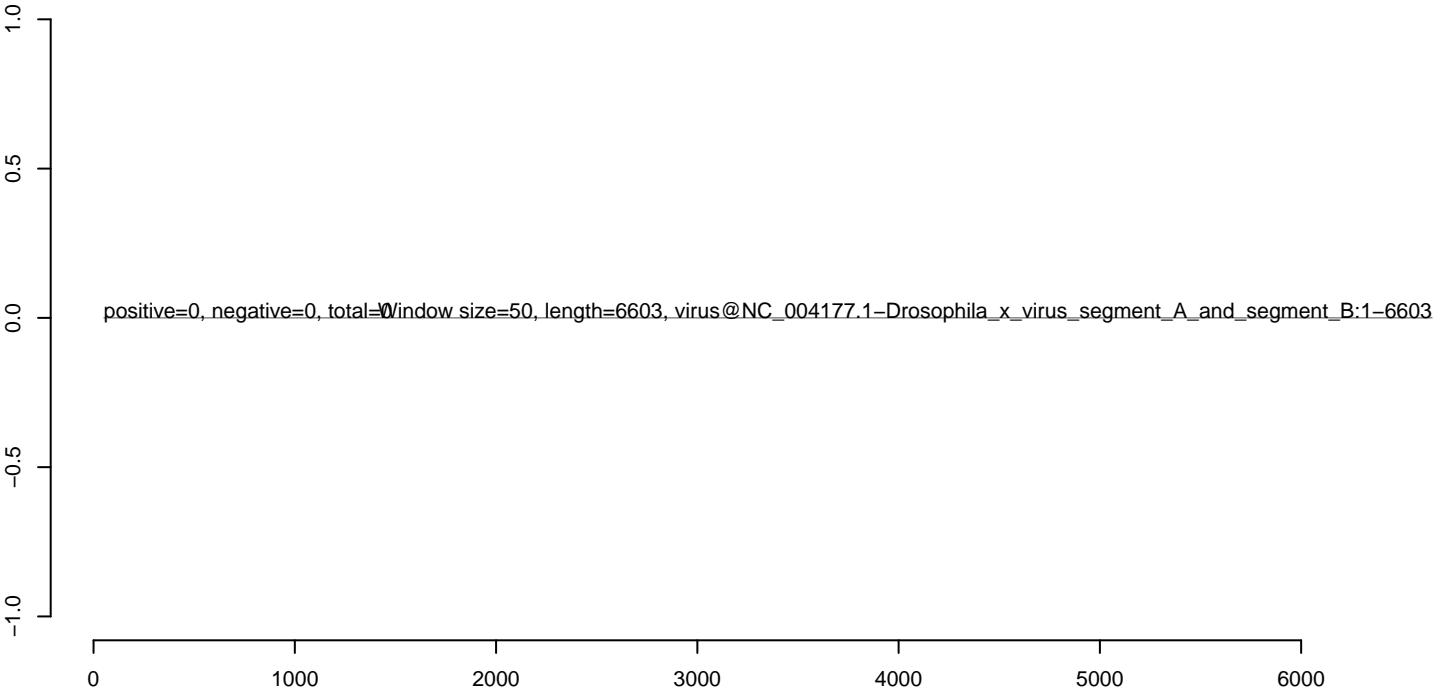
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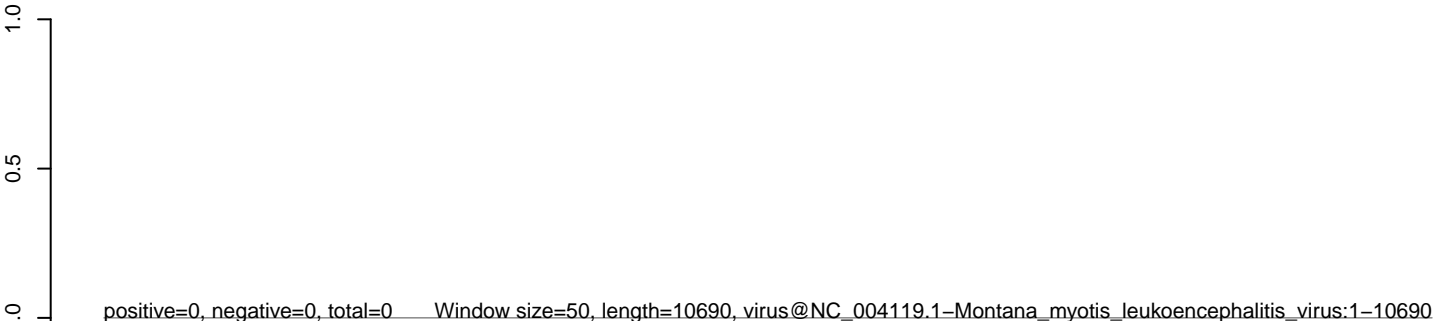
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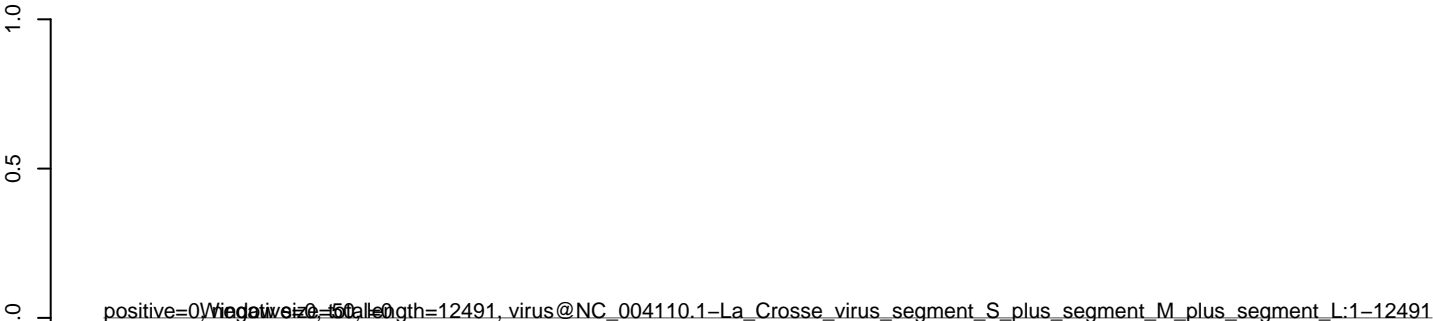
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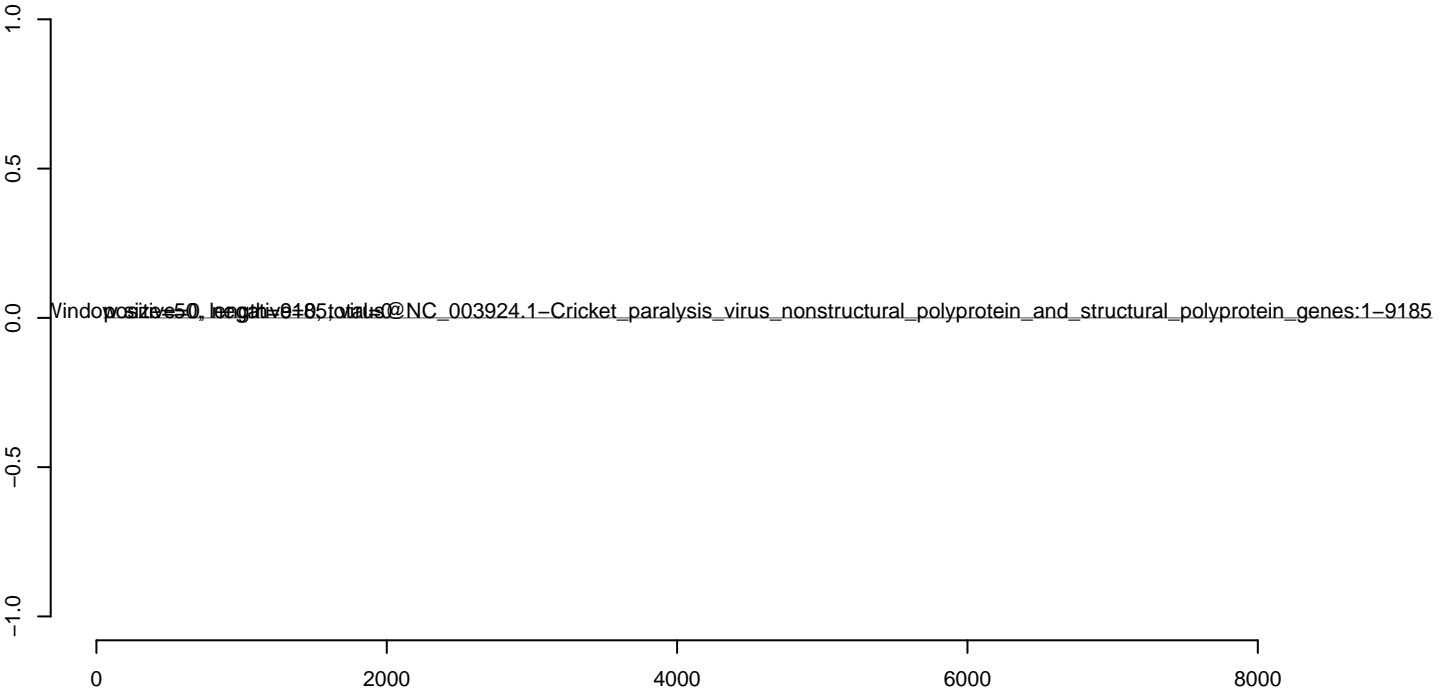
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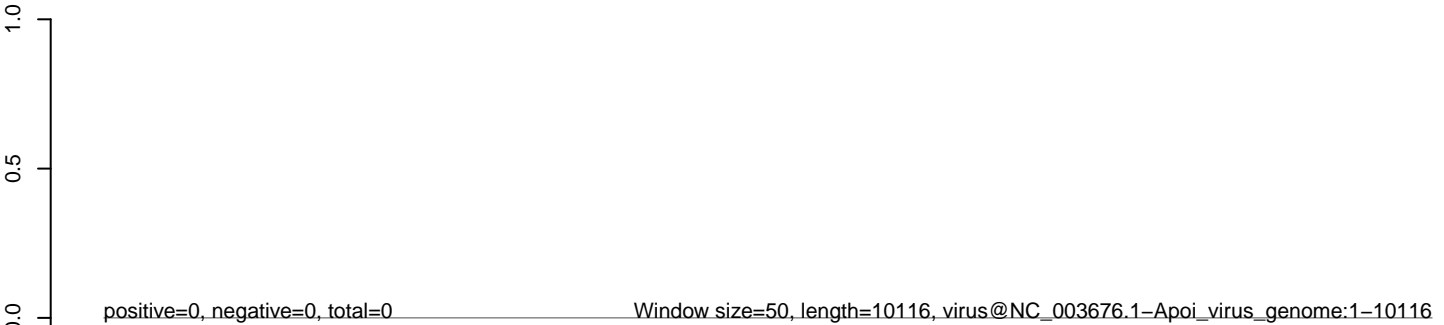
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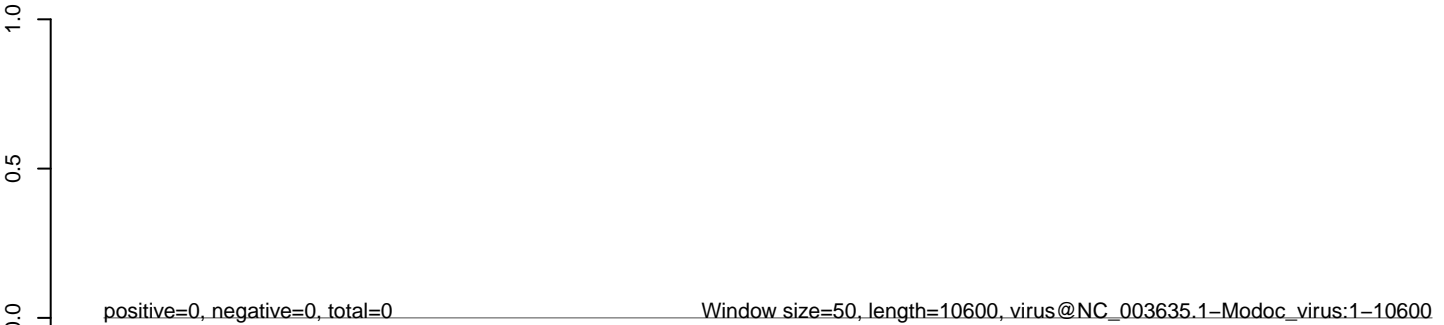
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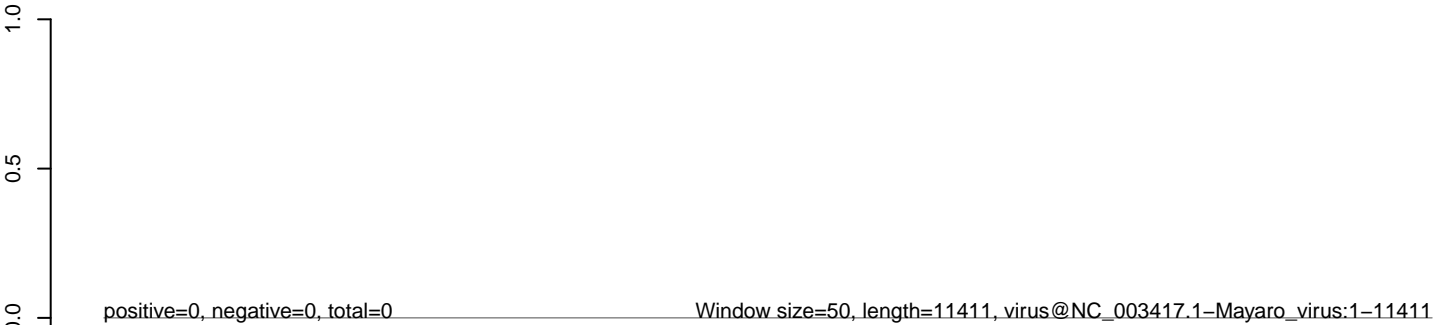
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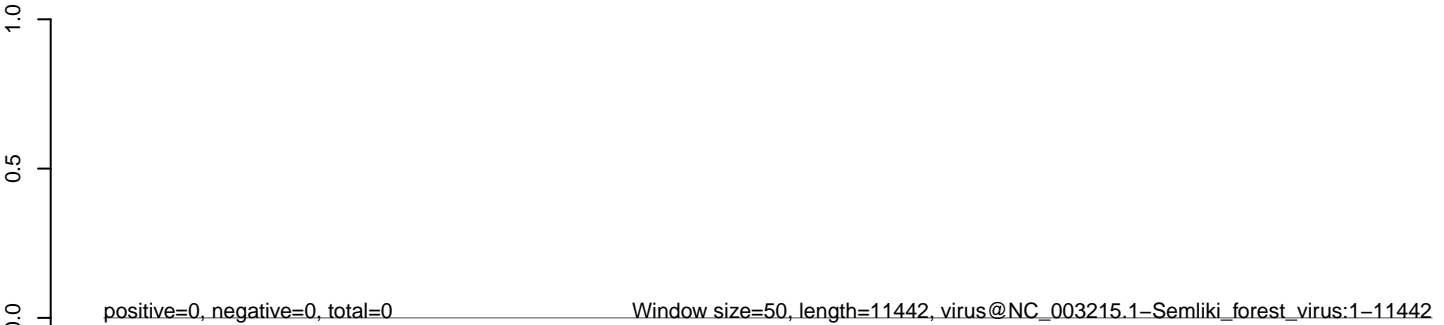
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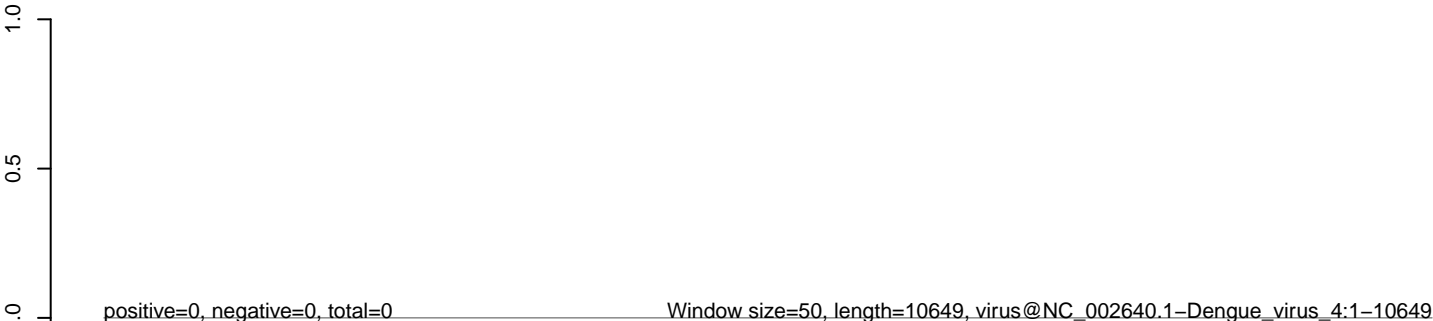
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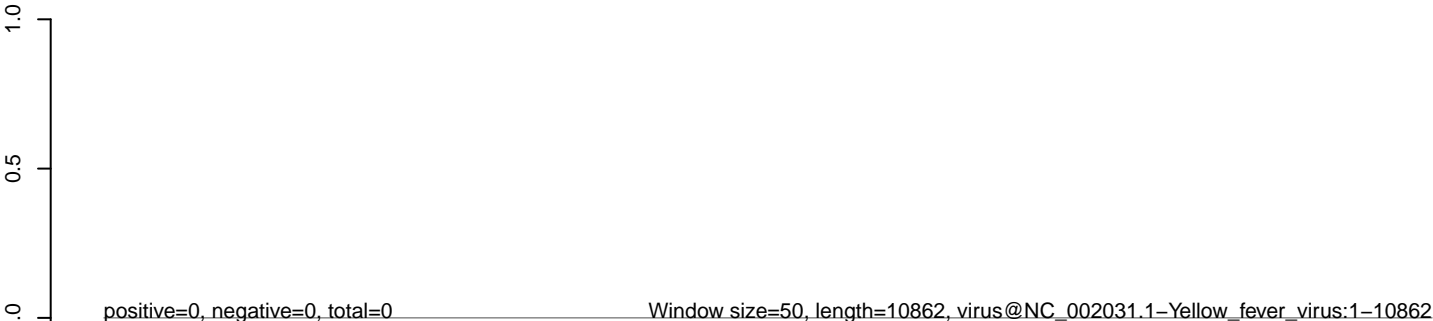
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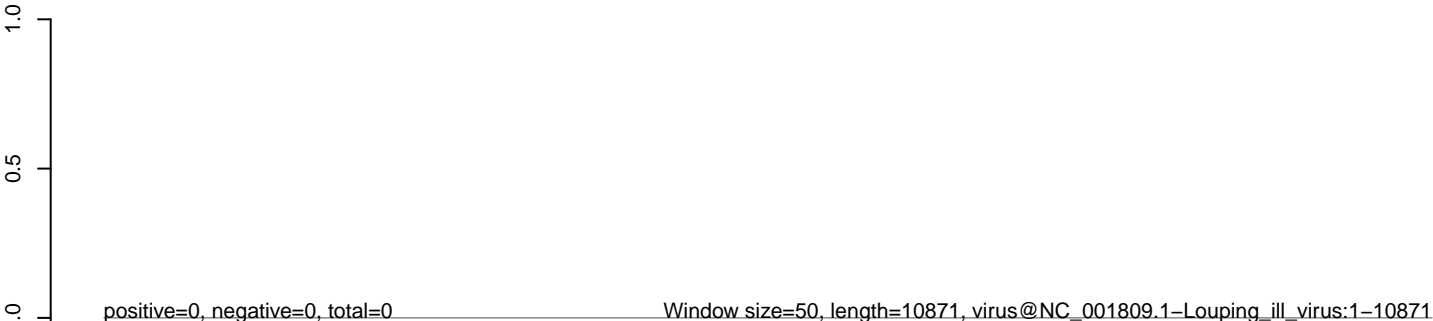
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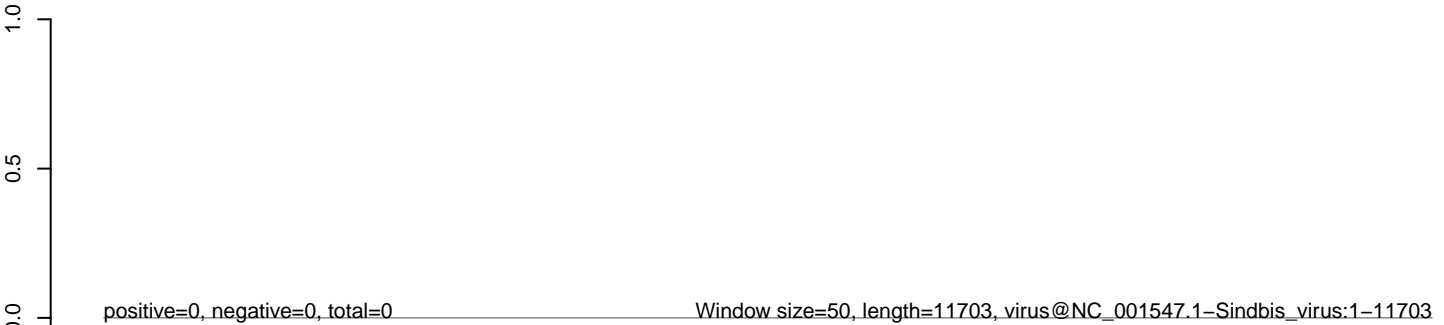
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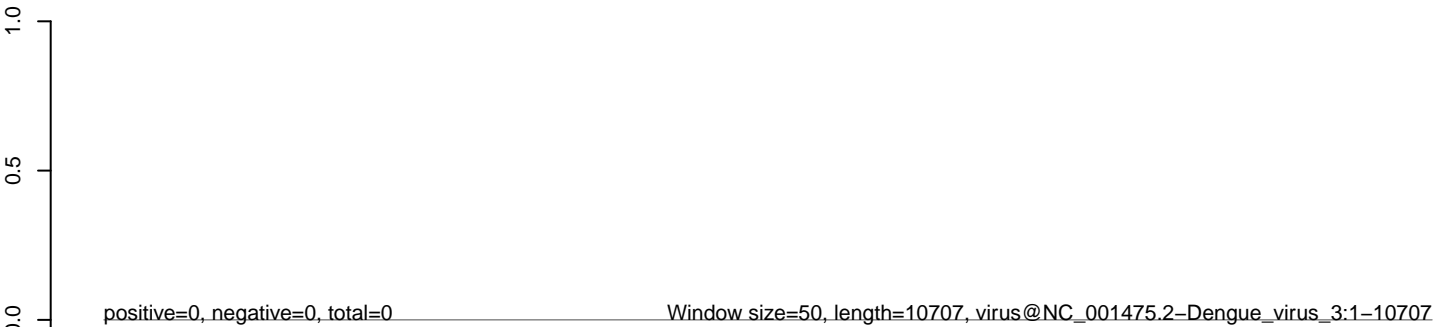
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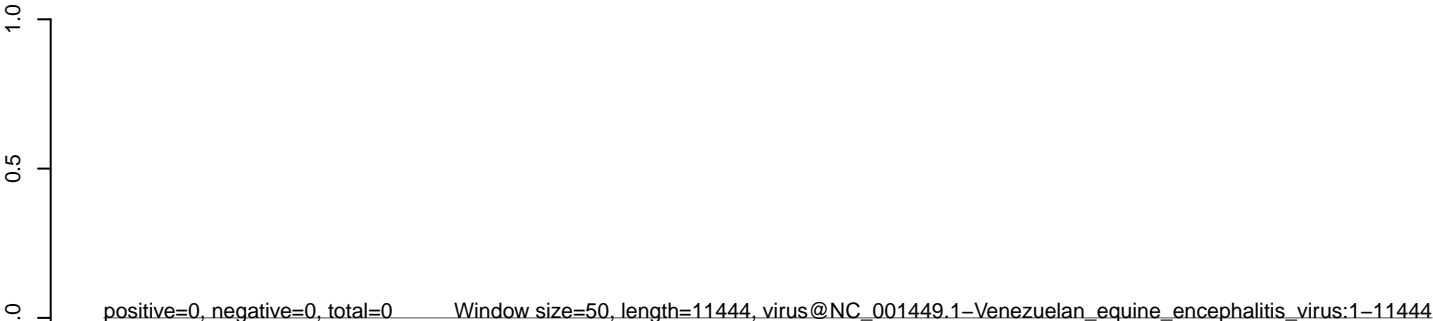
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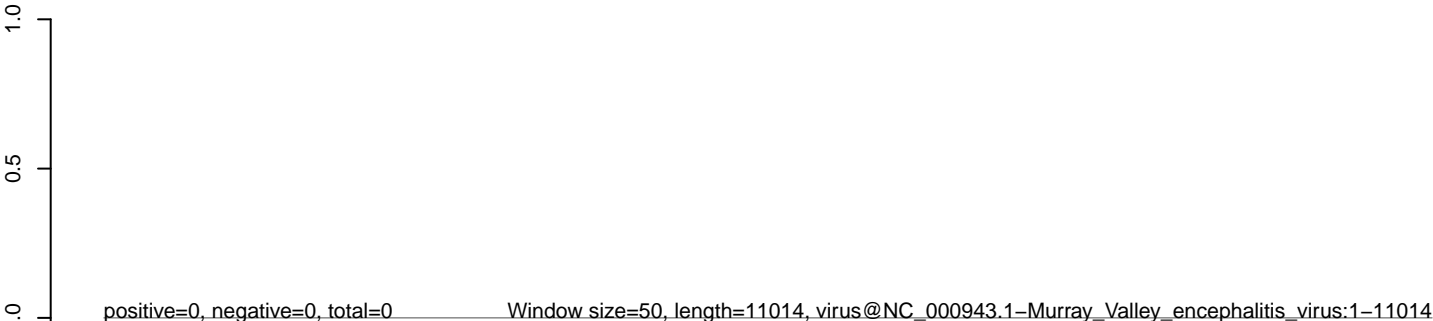
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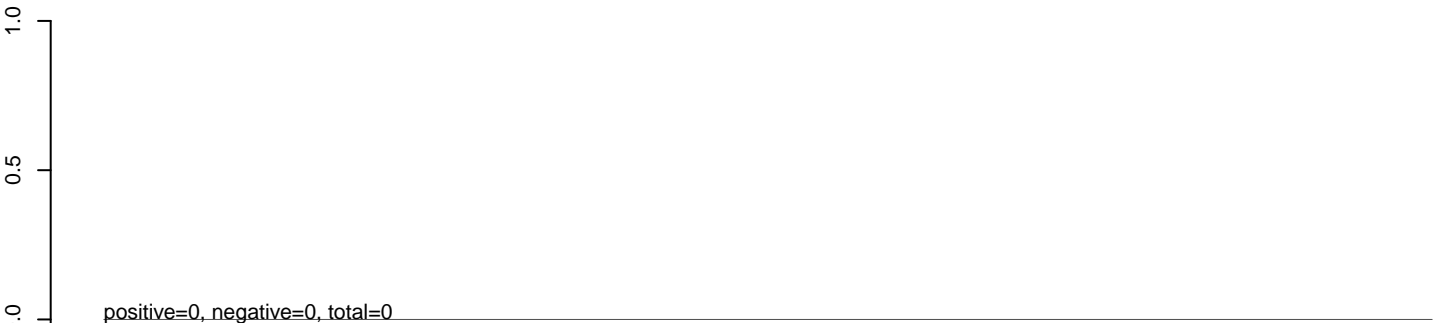


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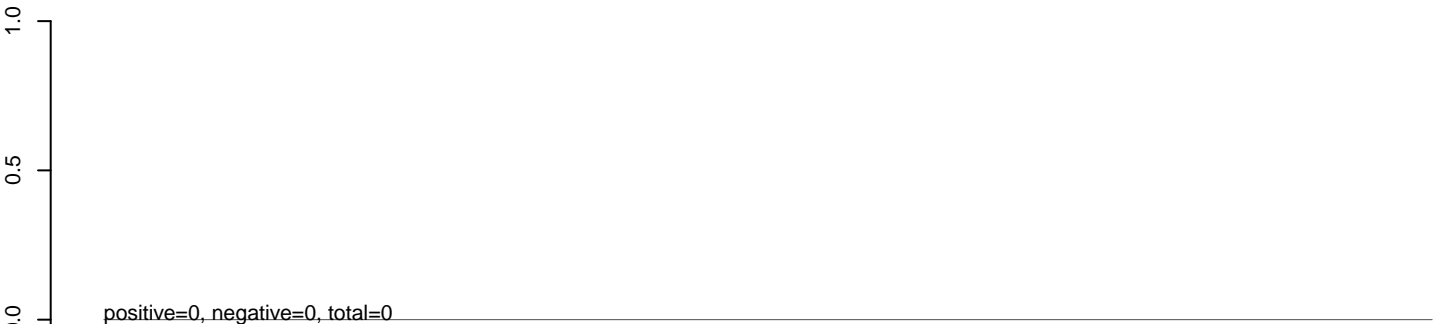


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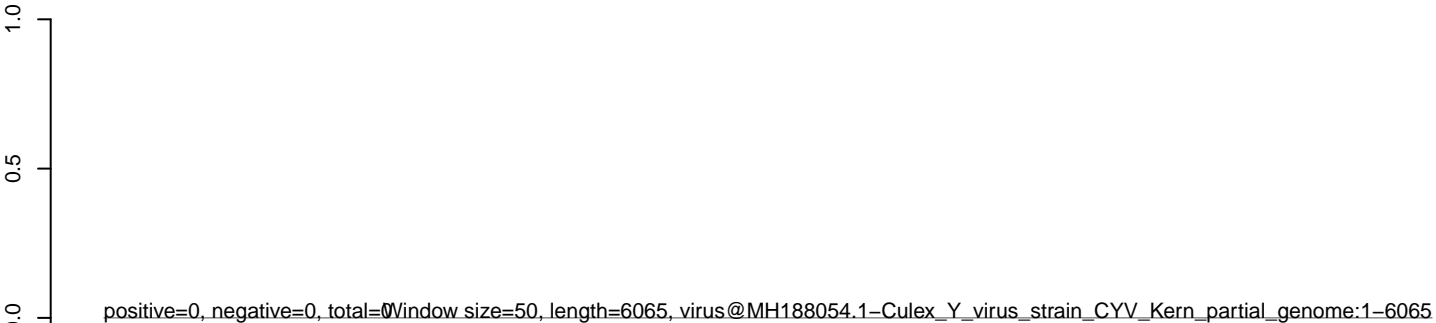
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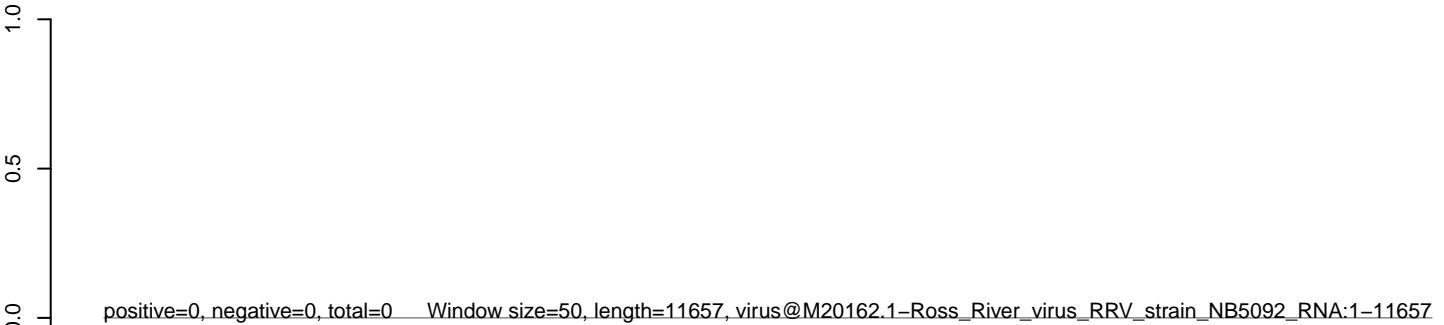
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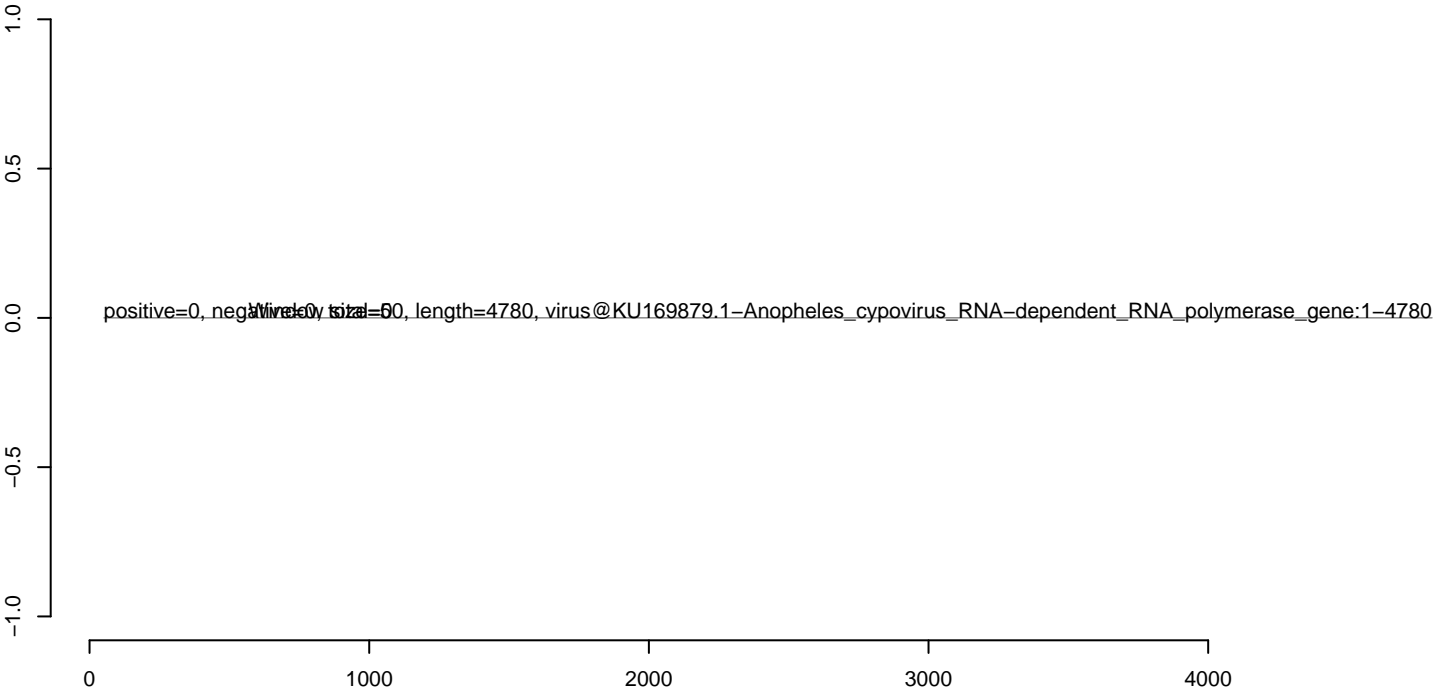
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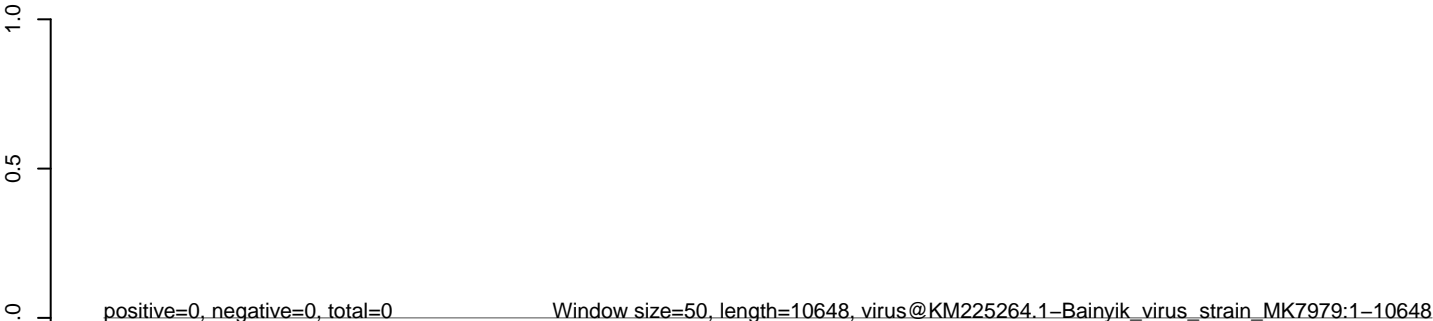
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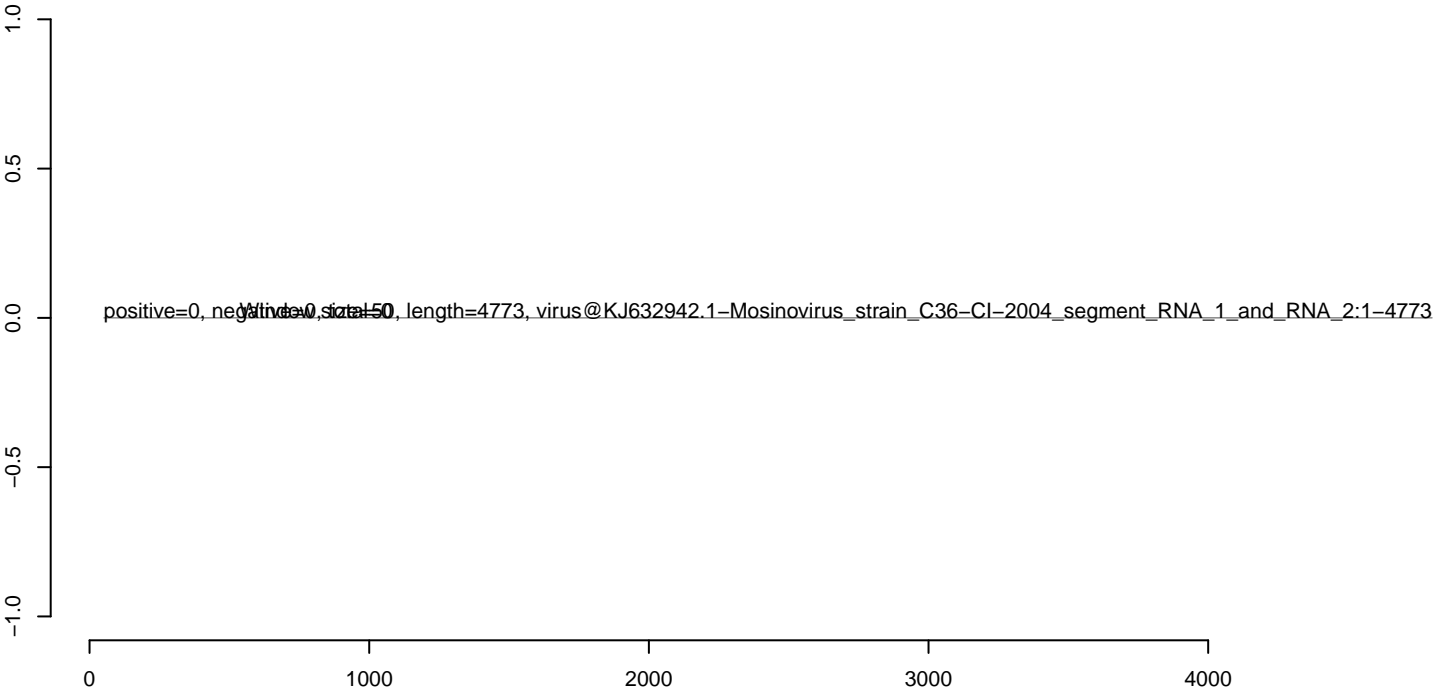
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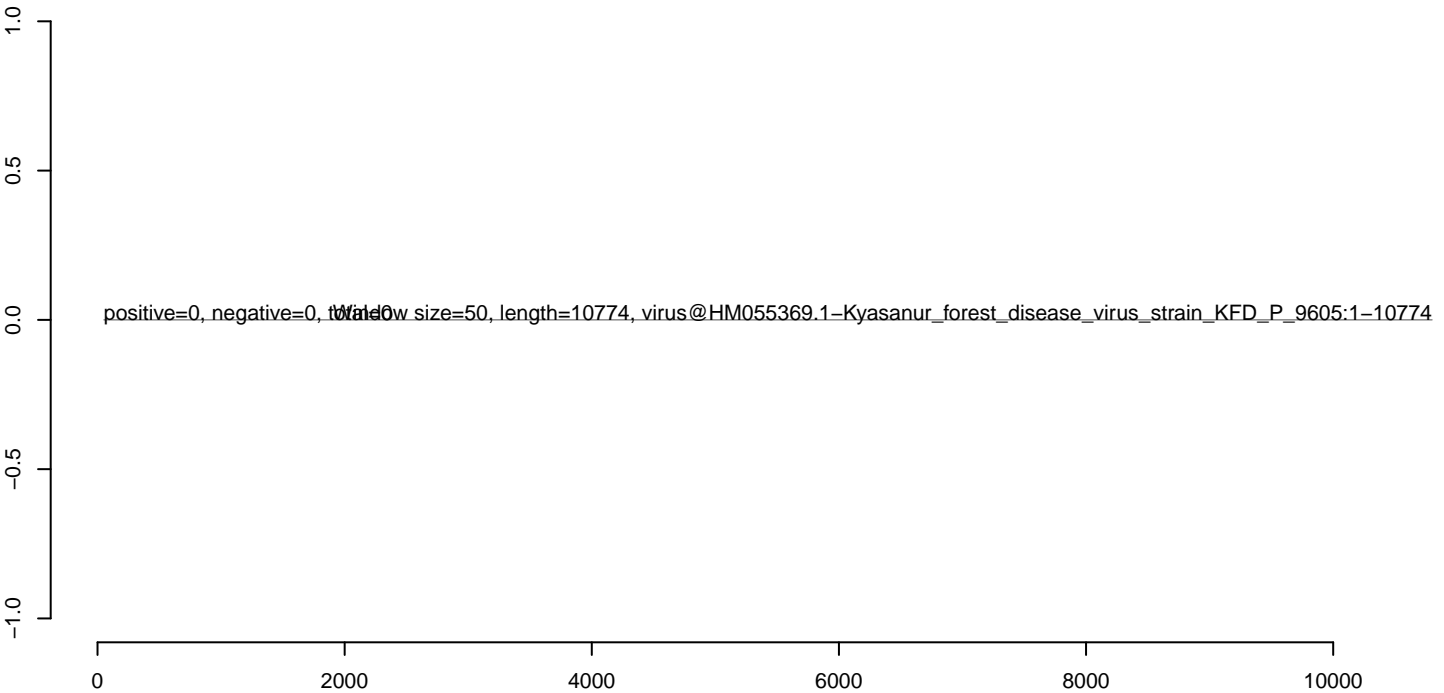
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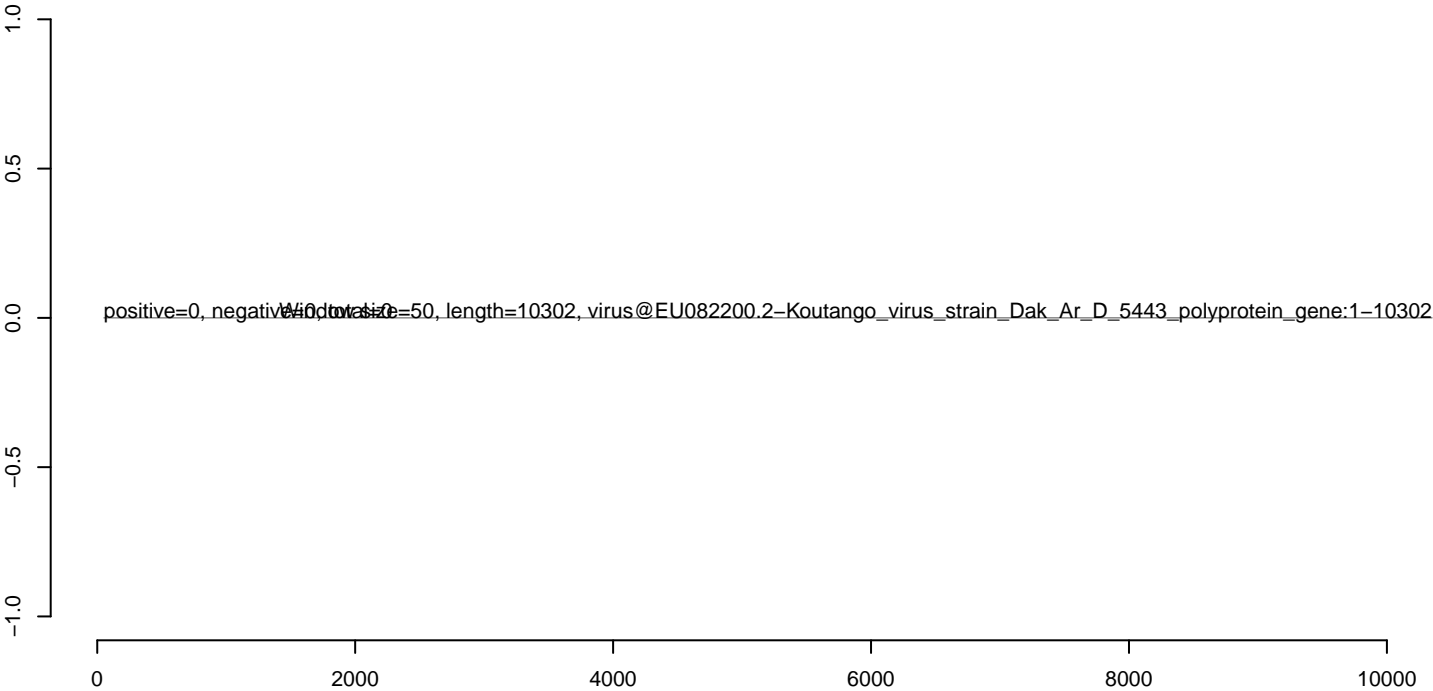
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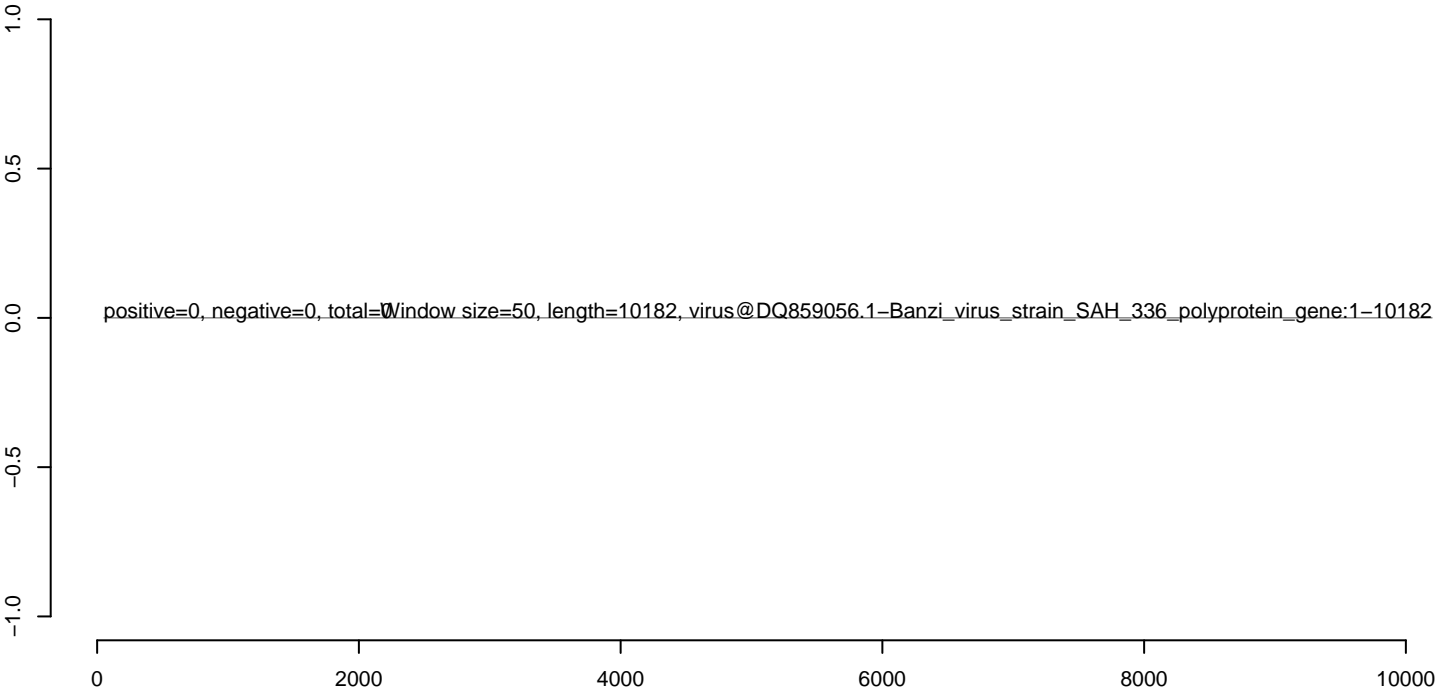
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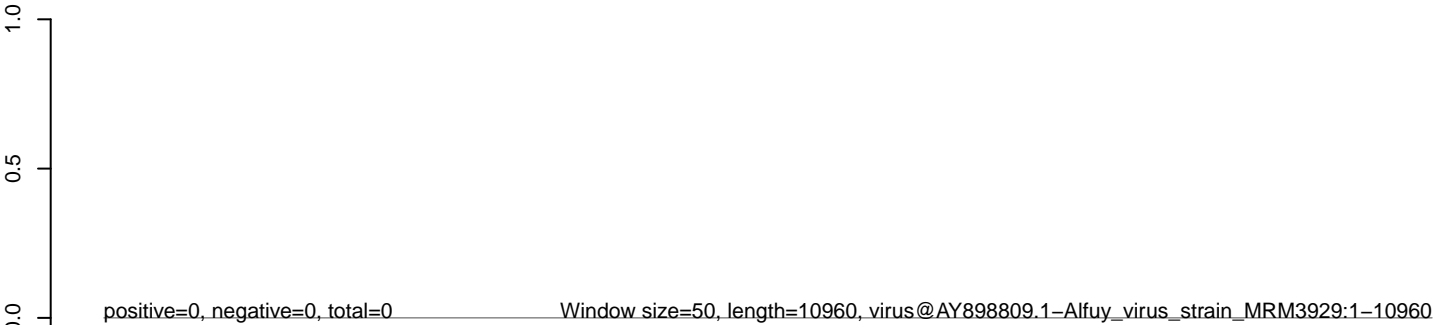
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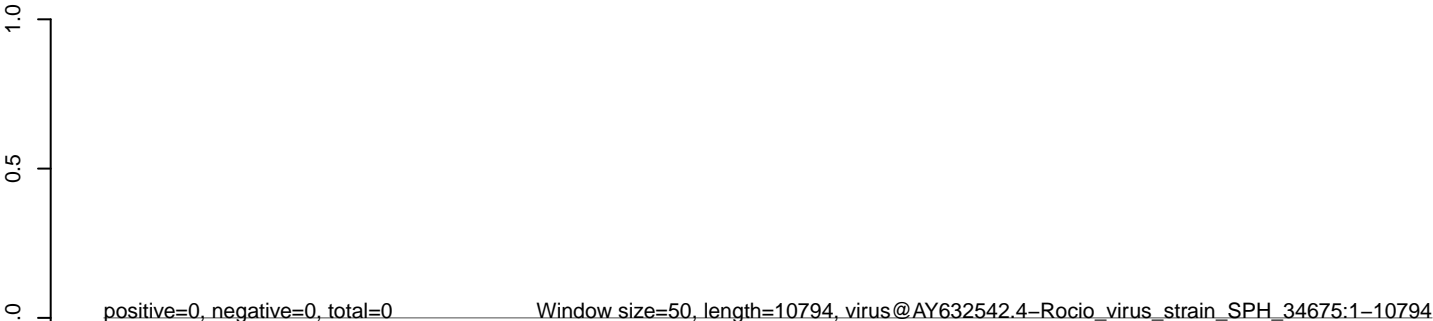
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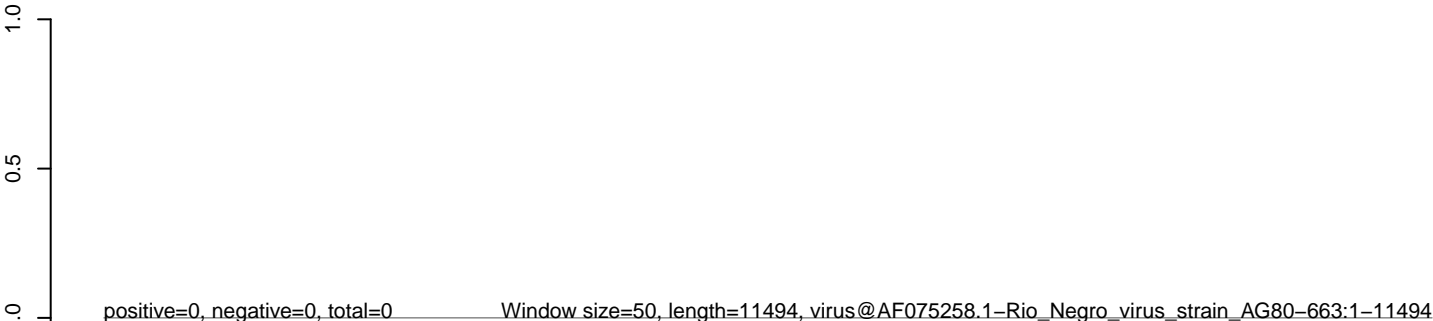
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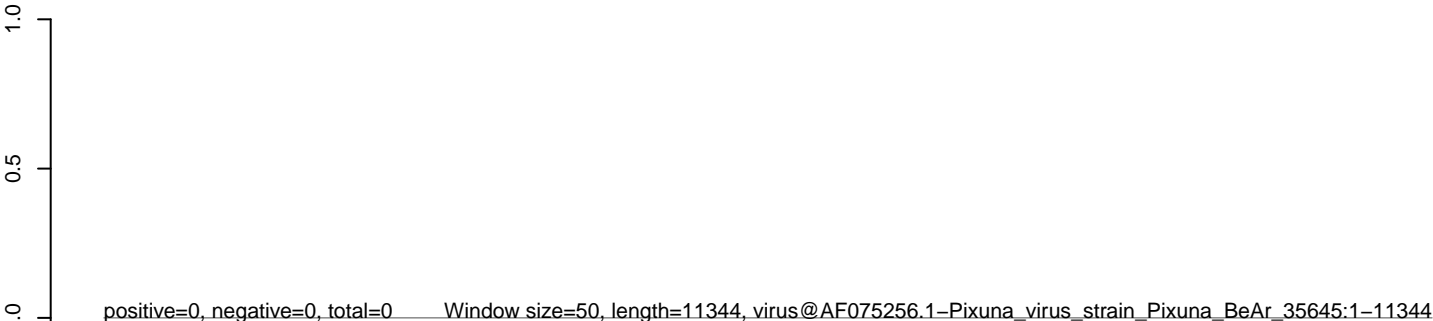
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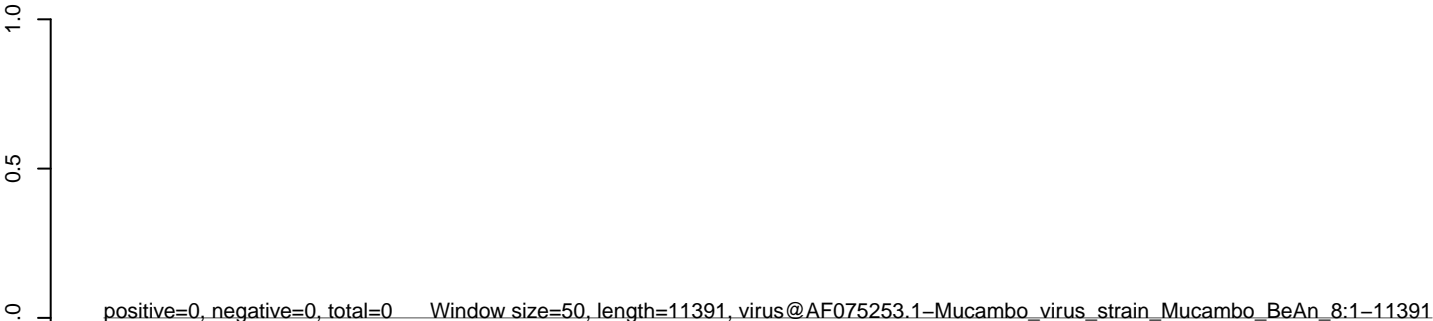
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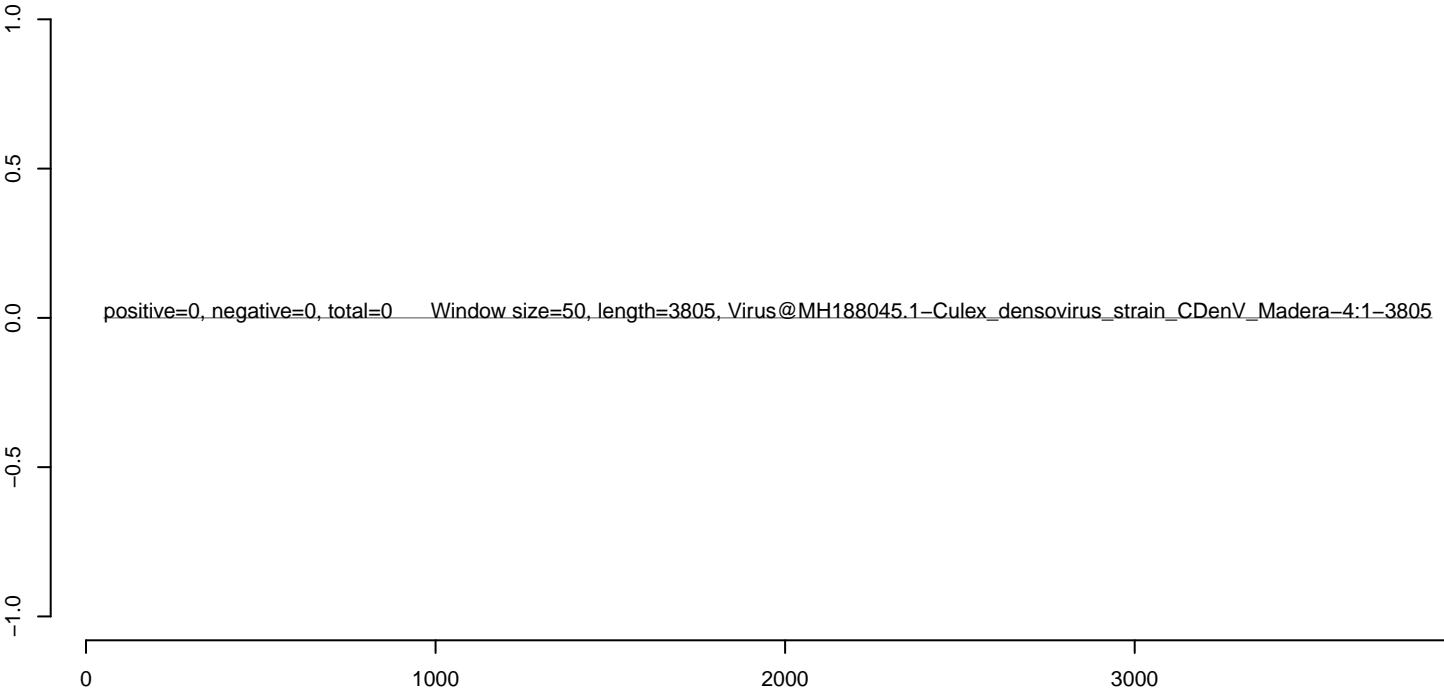
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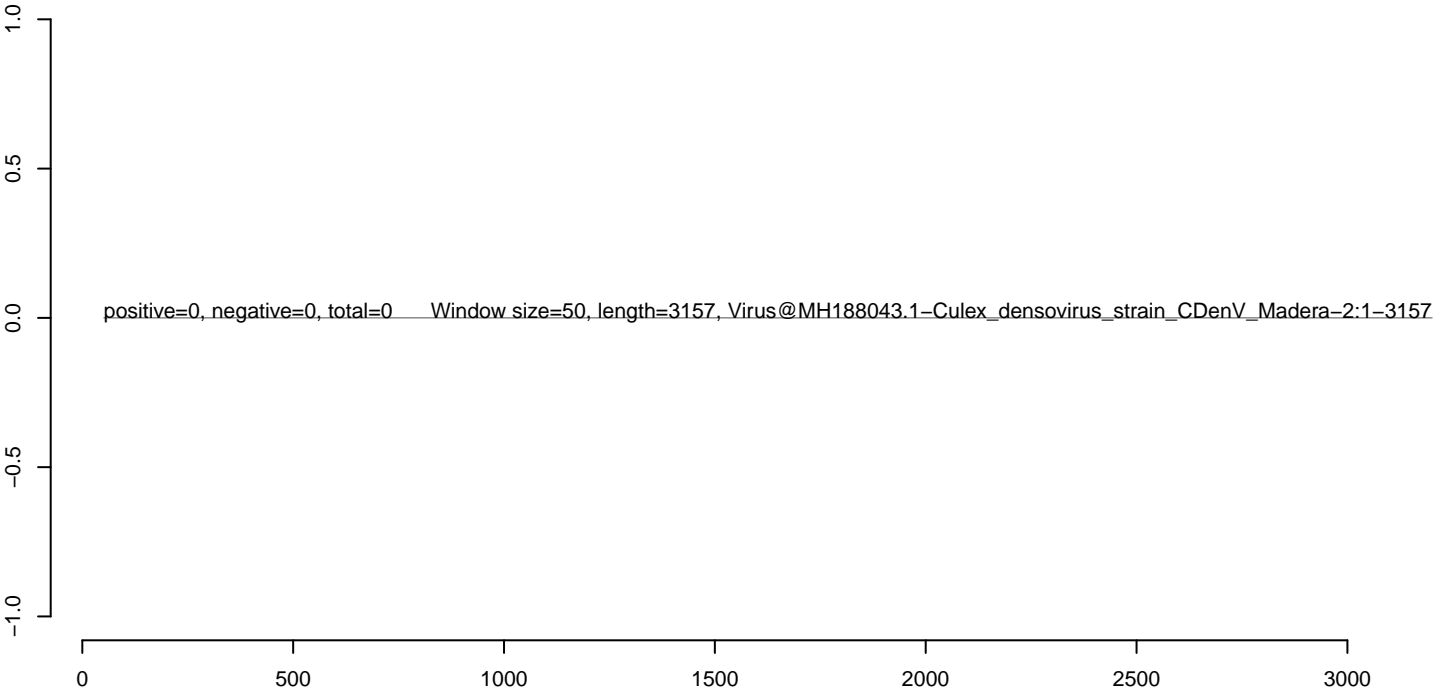
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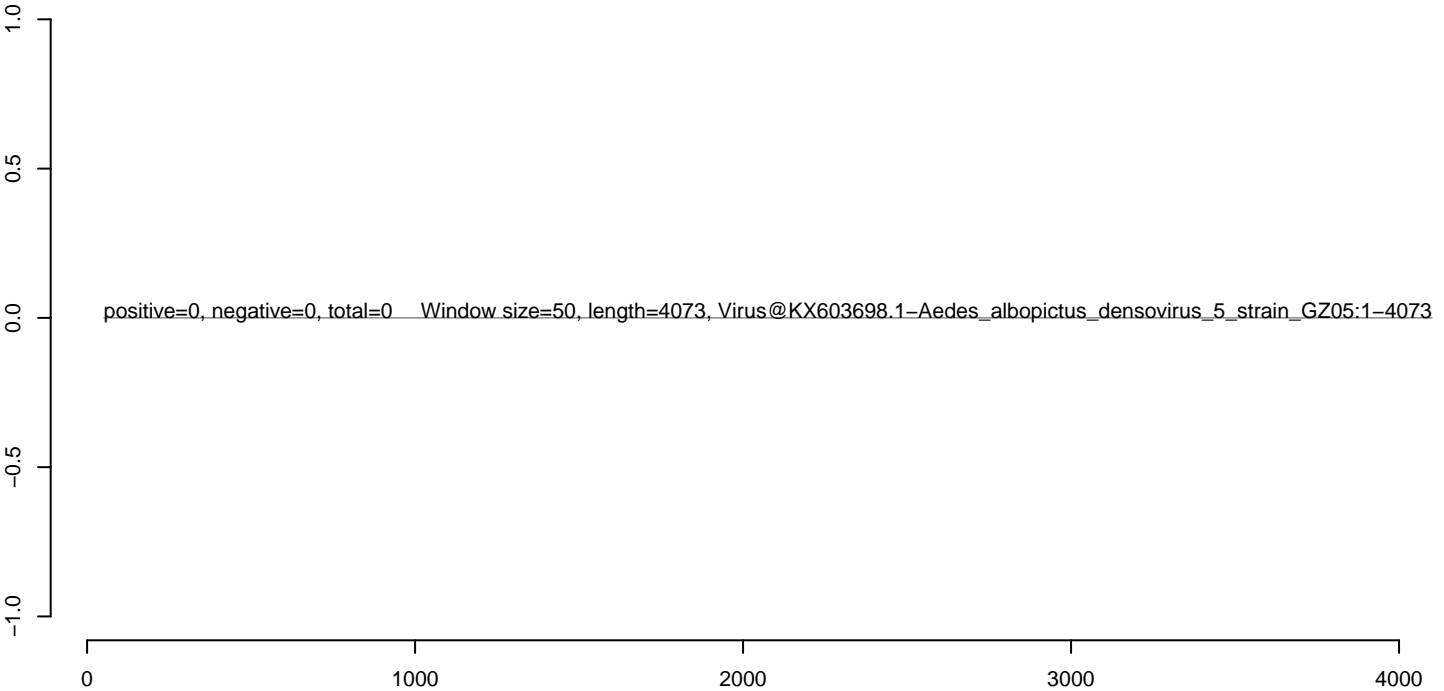
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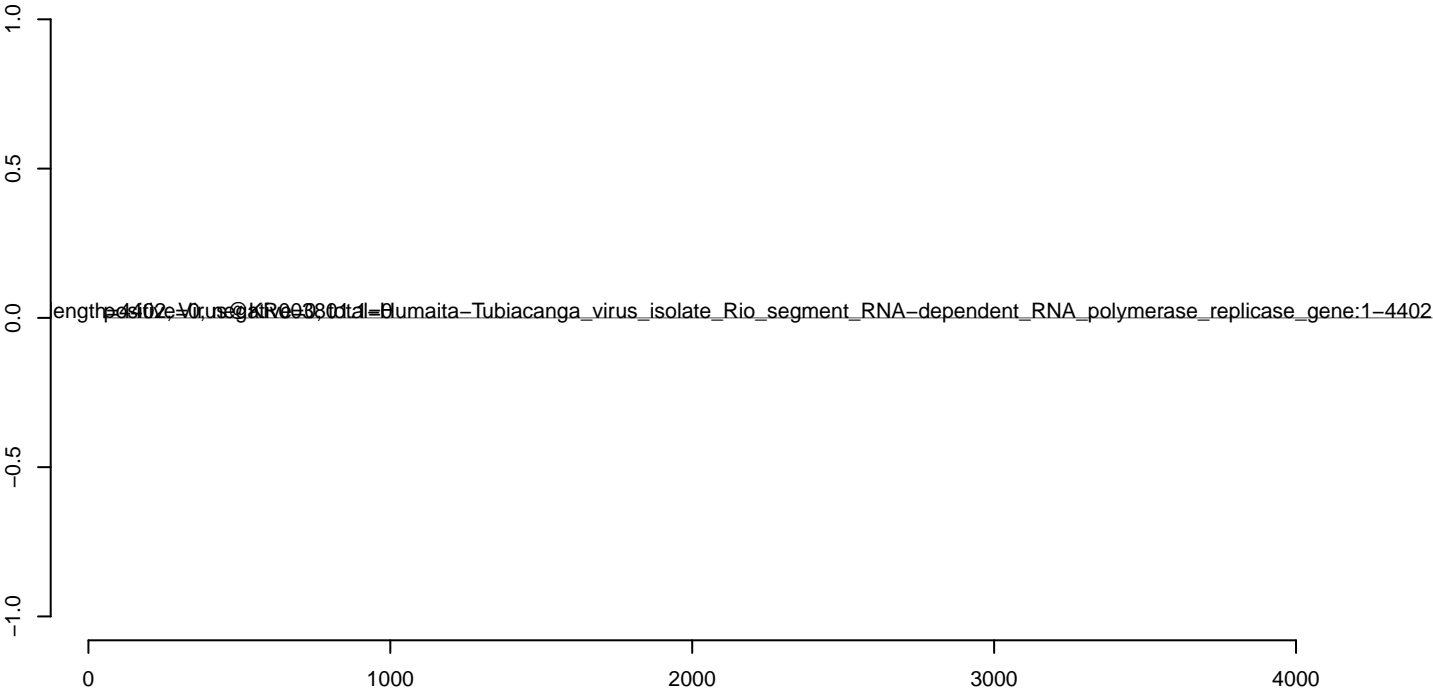
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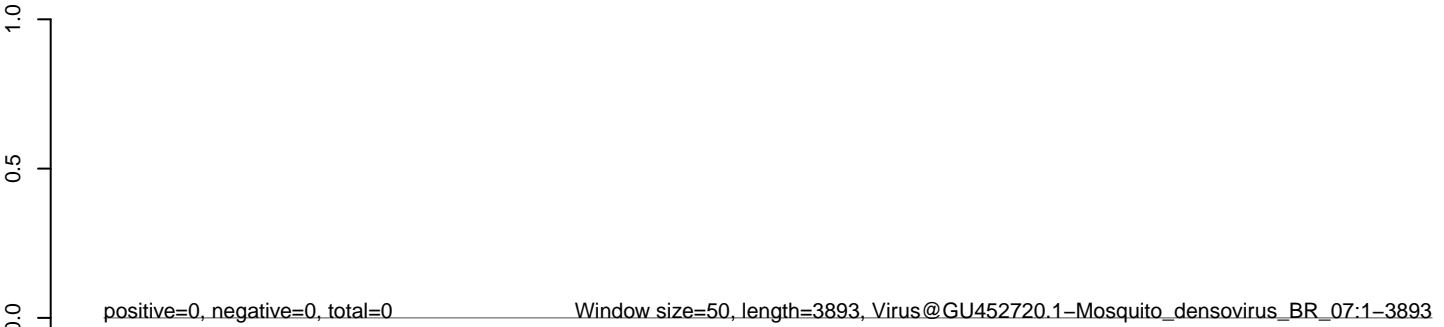
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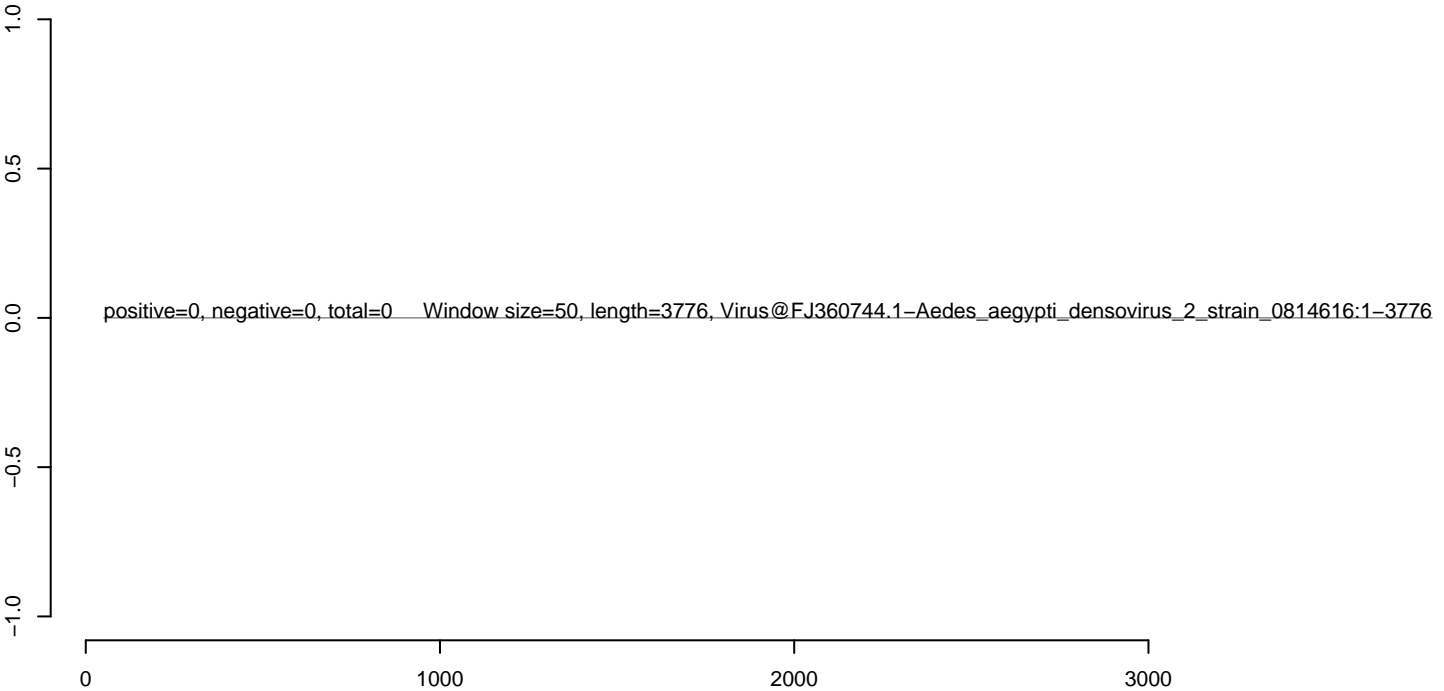
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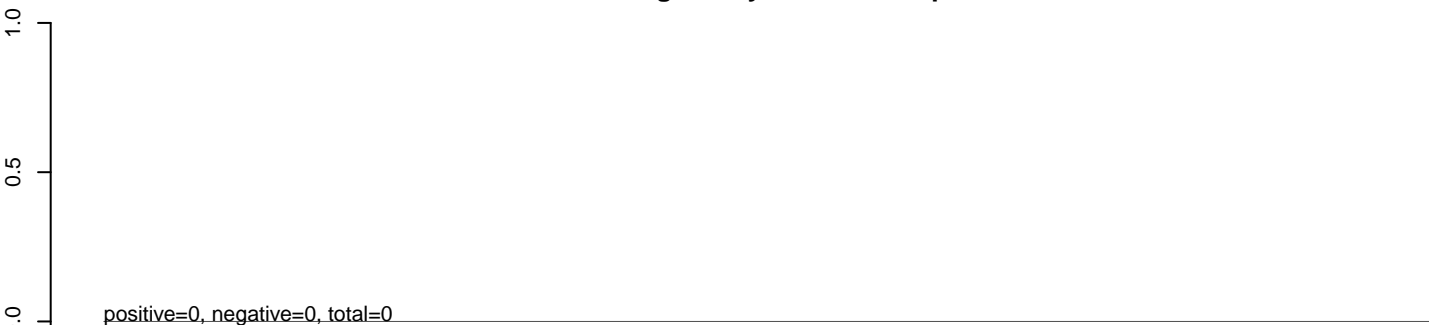
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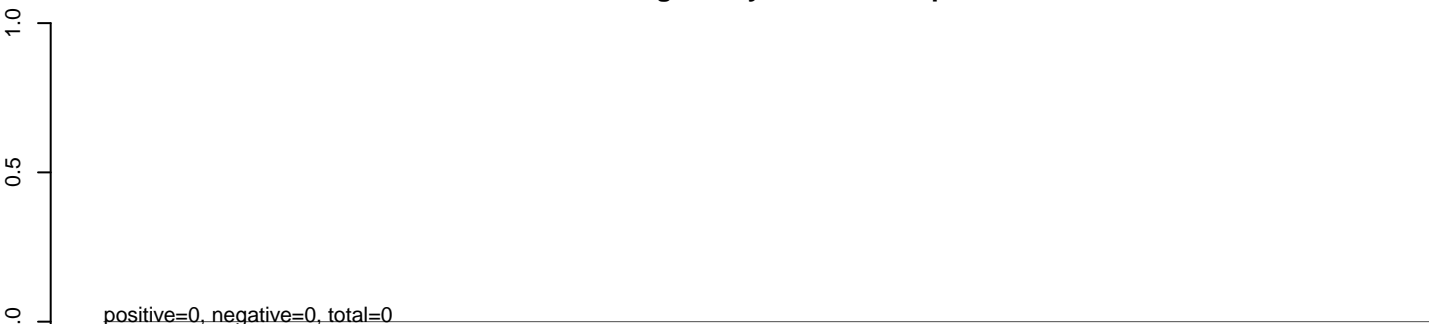
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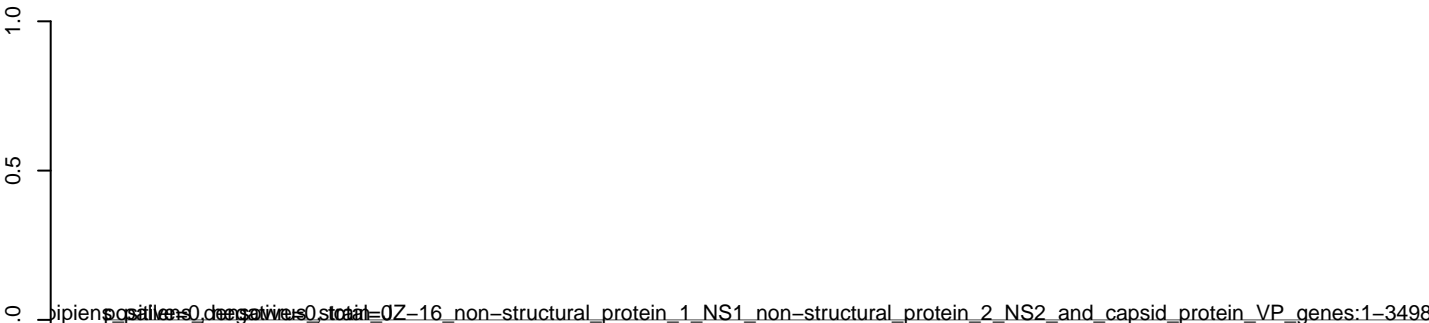
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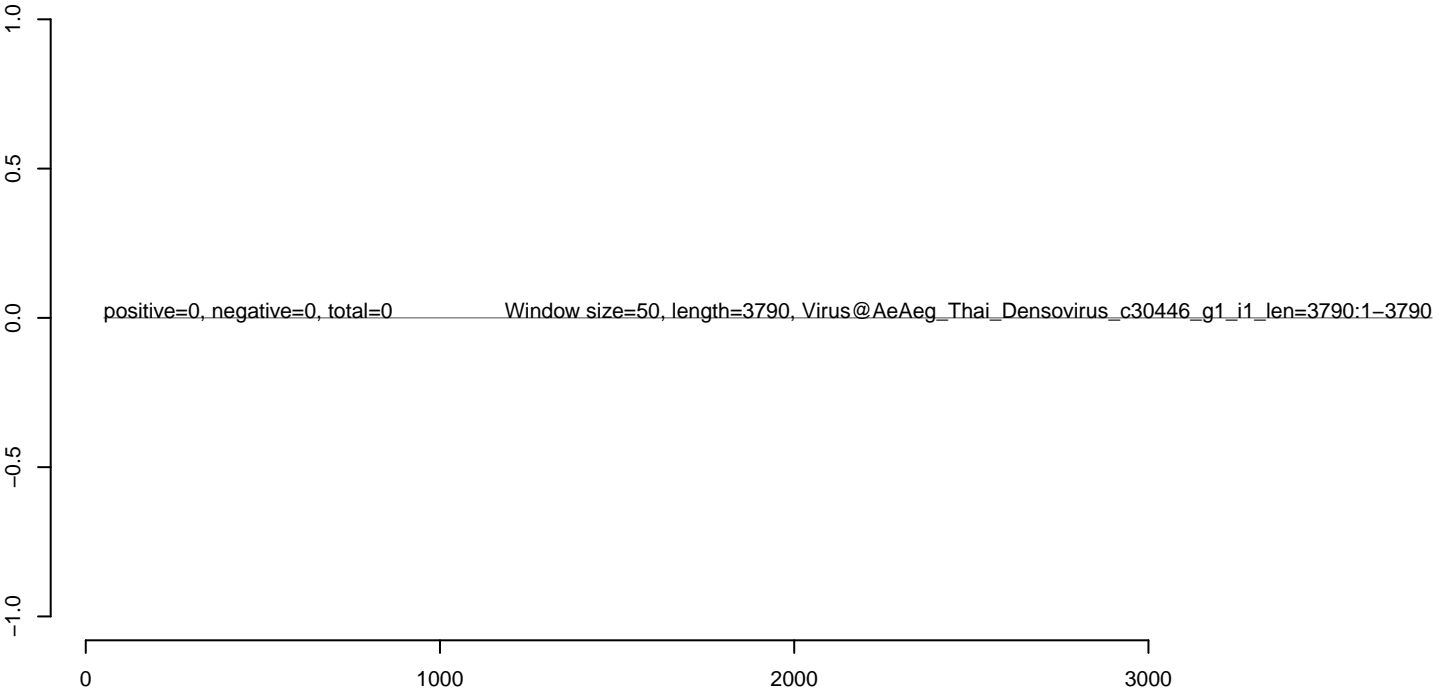
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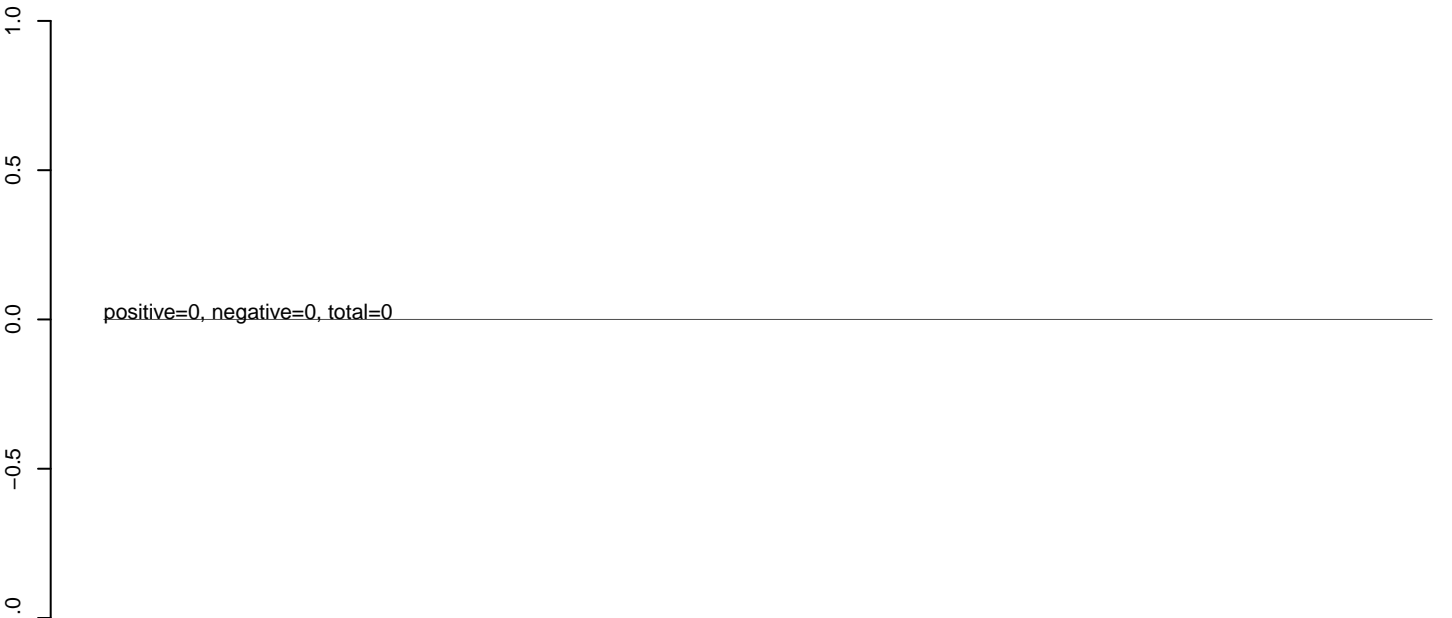
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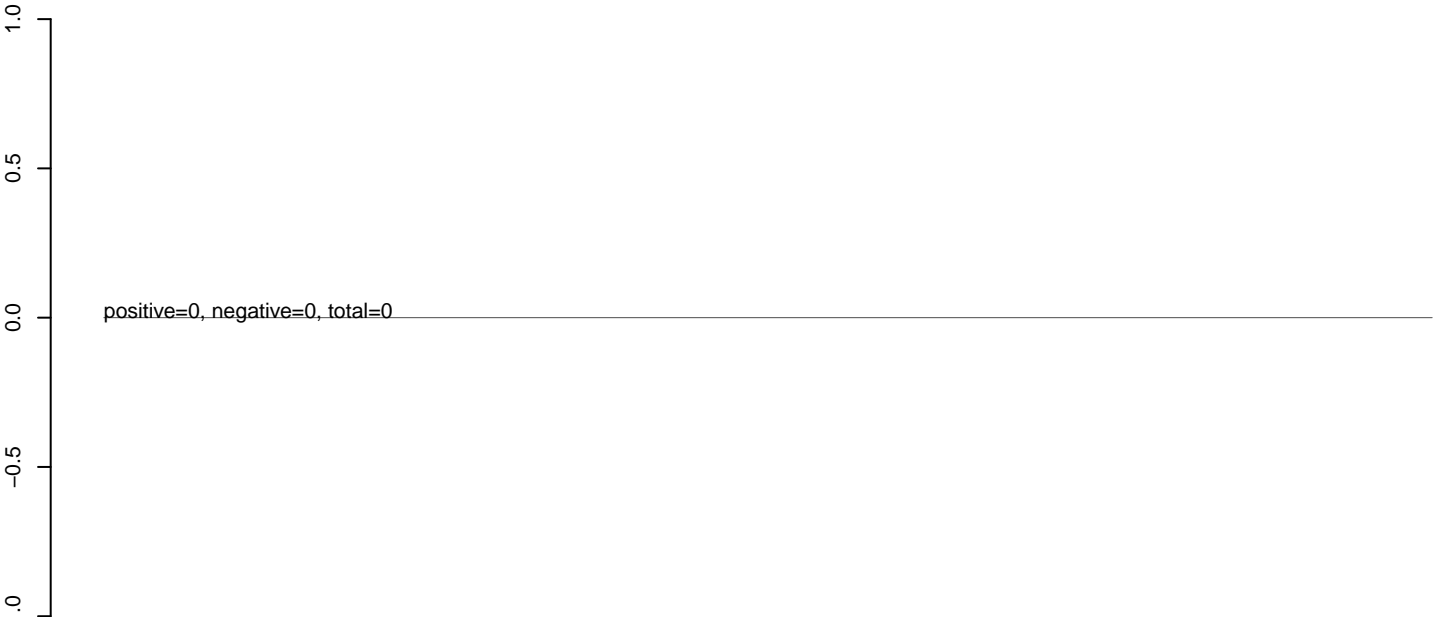
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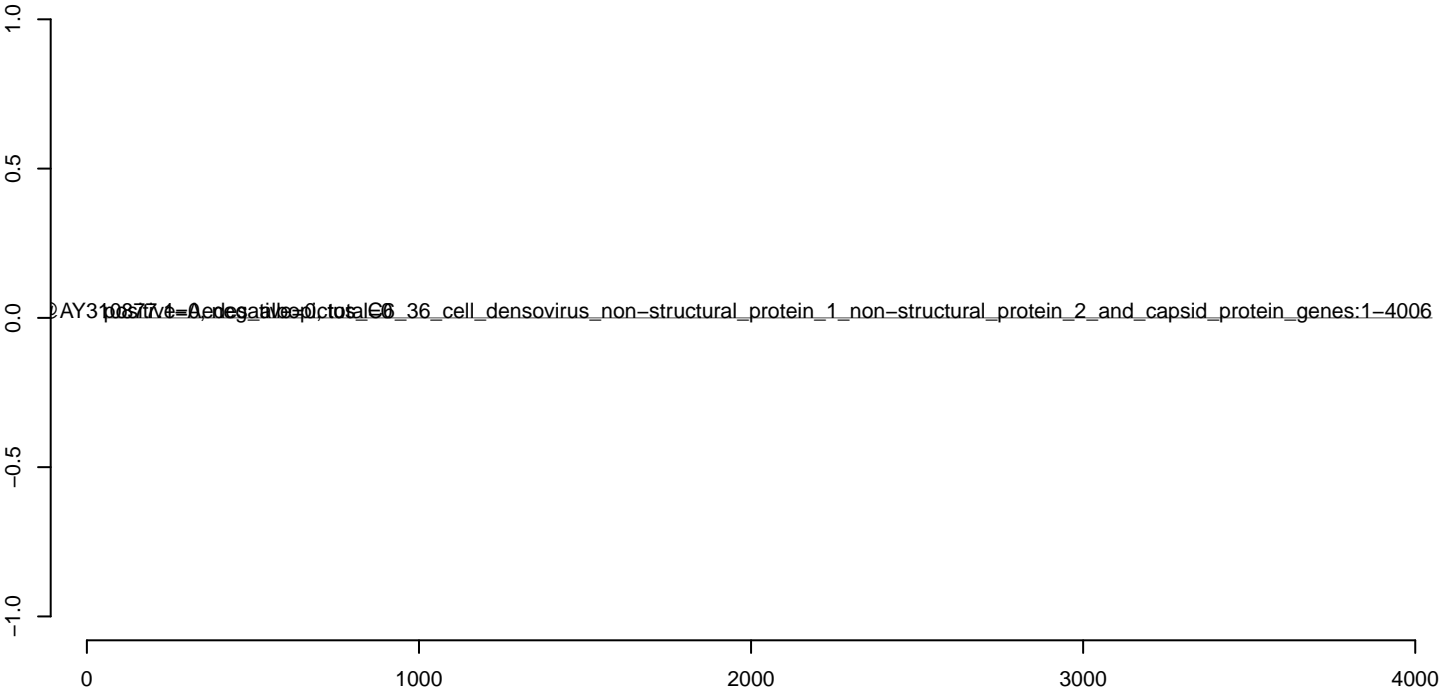
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AeAeg_Ovary_GH.rep



AeAeg_Ovary_GH.18_23.rep



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AeAeg_Ovary_GH.rep

