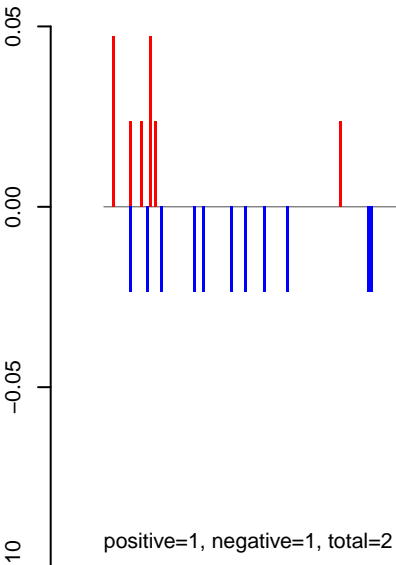
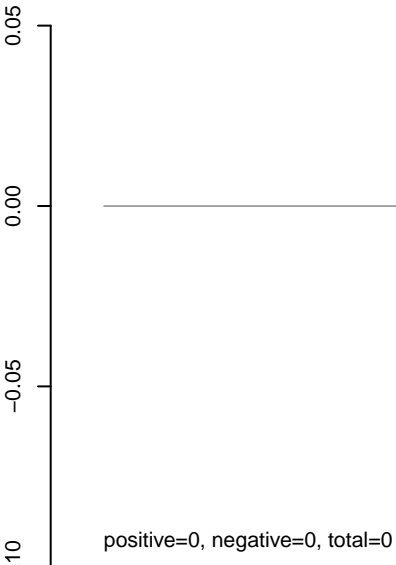


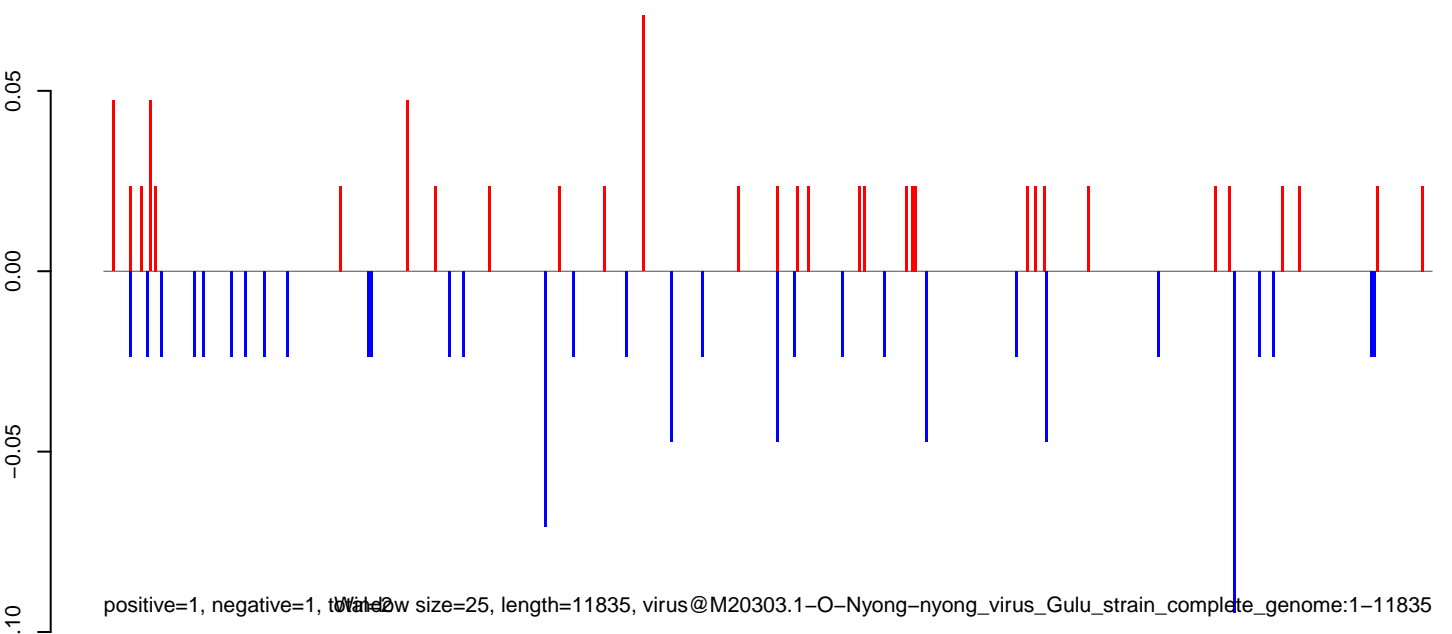
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

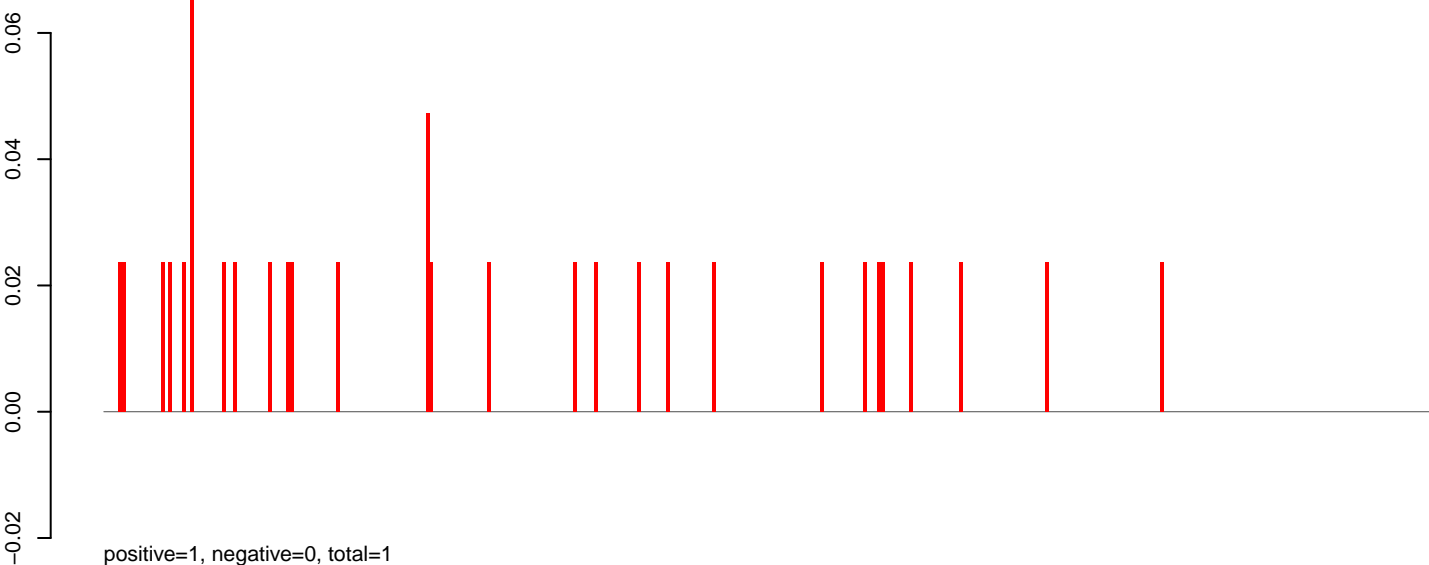


AnGam_Ovary_BetaE.rep

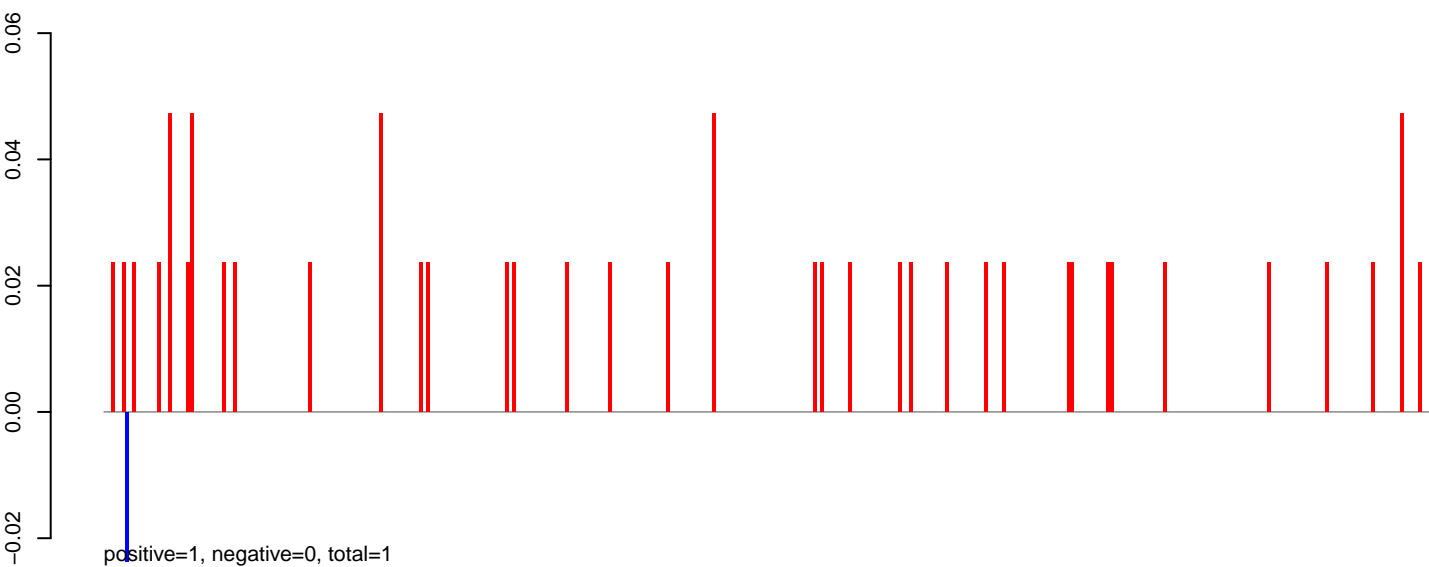


Window size=25, length=11835, virus@M20303.1-O-Nyong-nyong_virus_Gulu_strain_complete_genome:1-11835

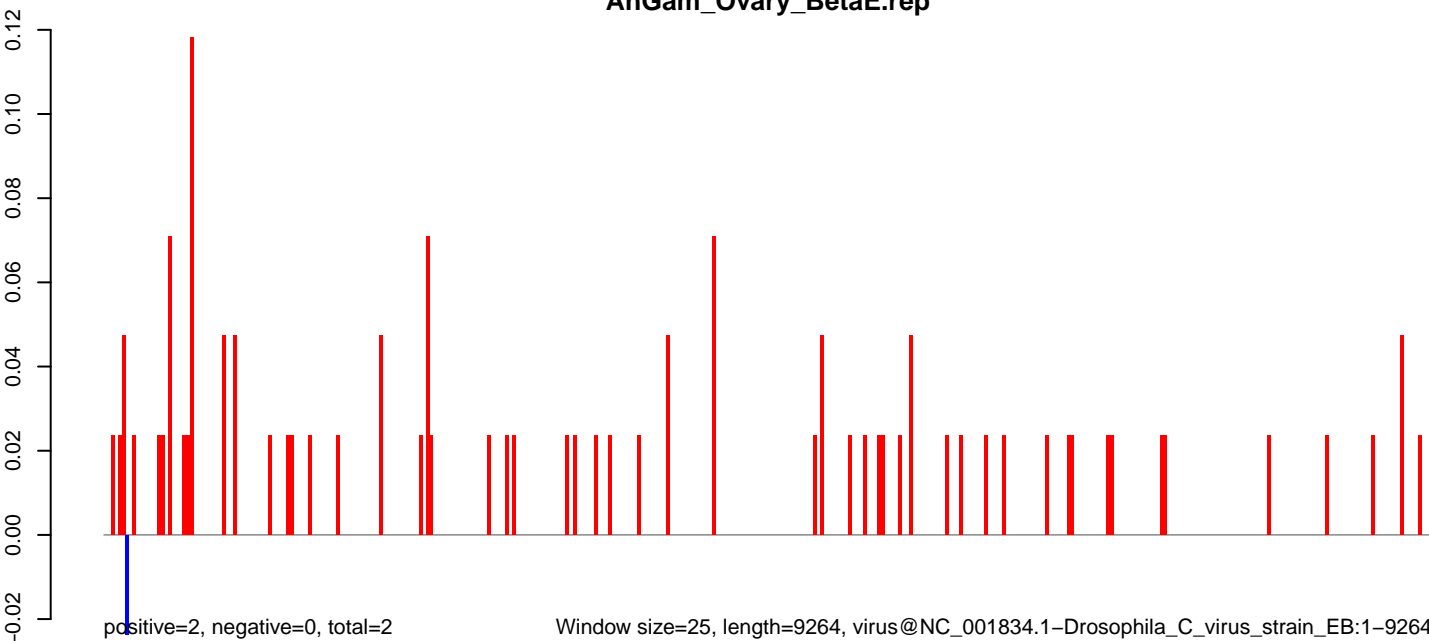
AnGam_Ovary_BetaE.18_23.rep



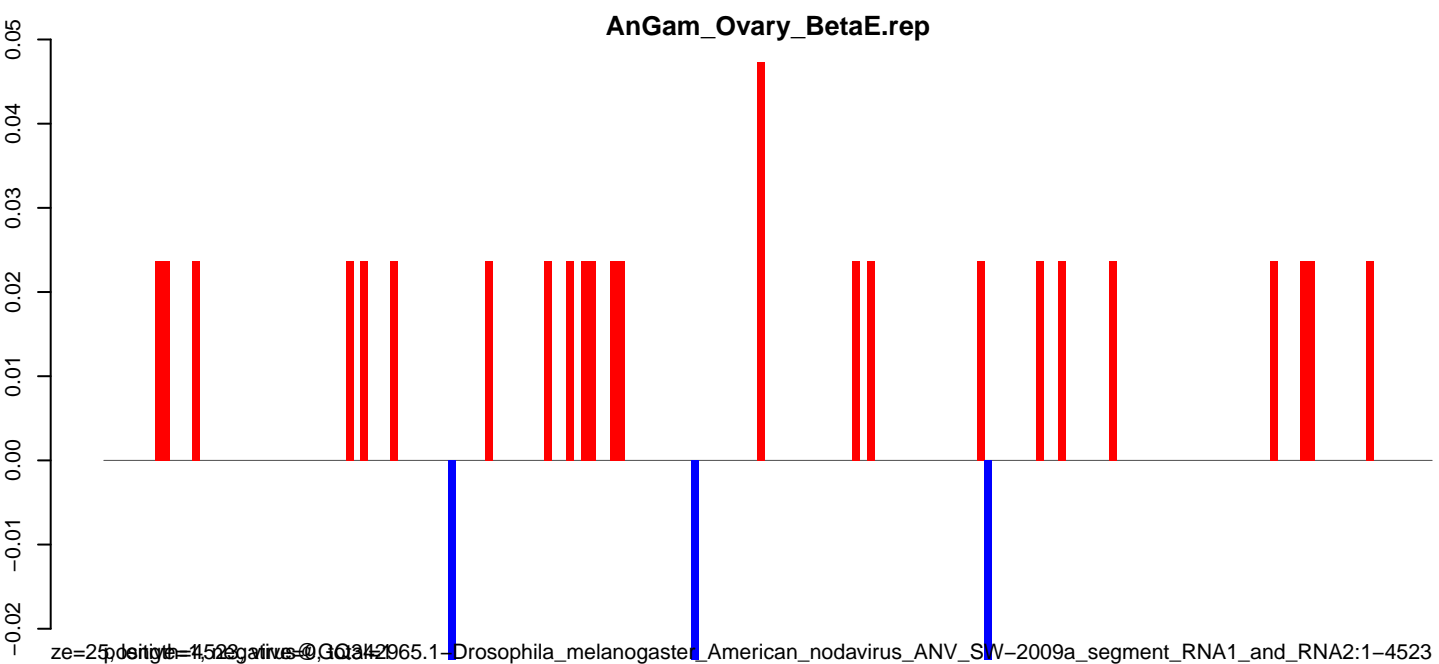
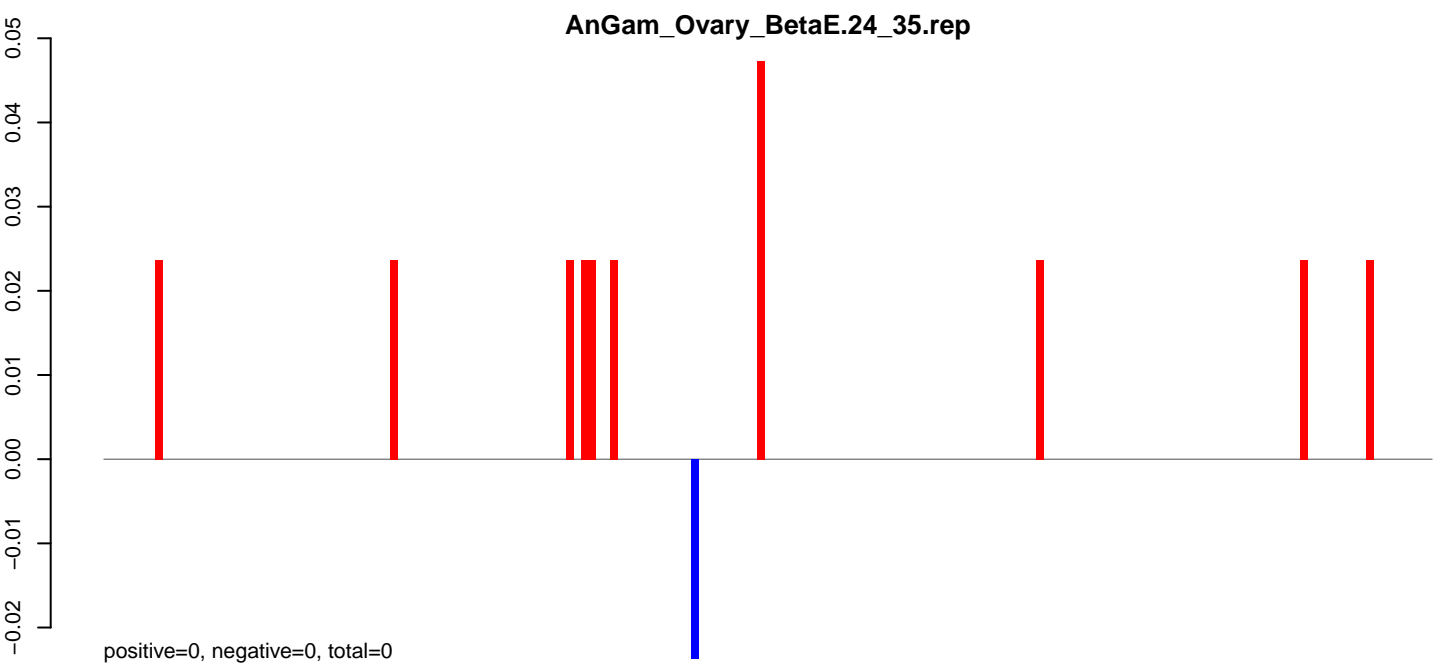
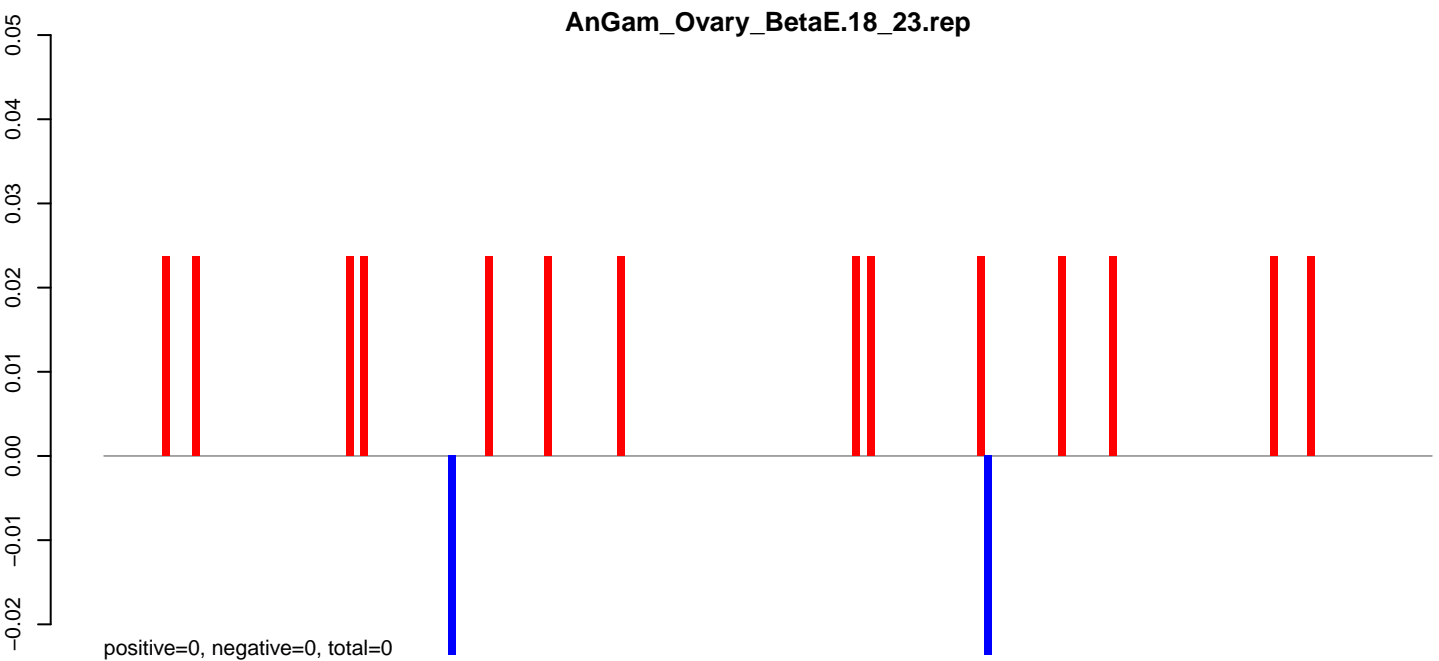
AnGam_Ovary_BetaE.24_35.rep



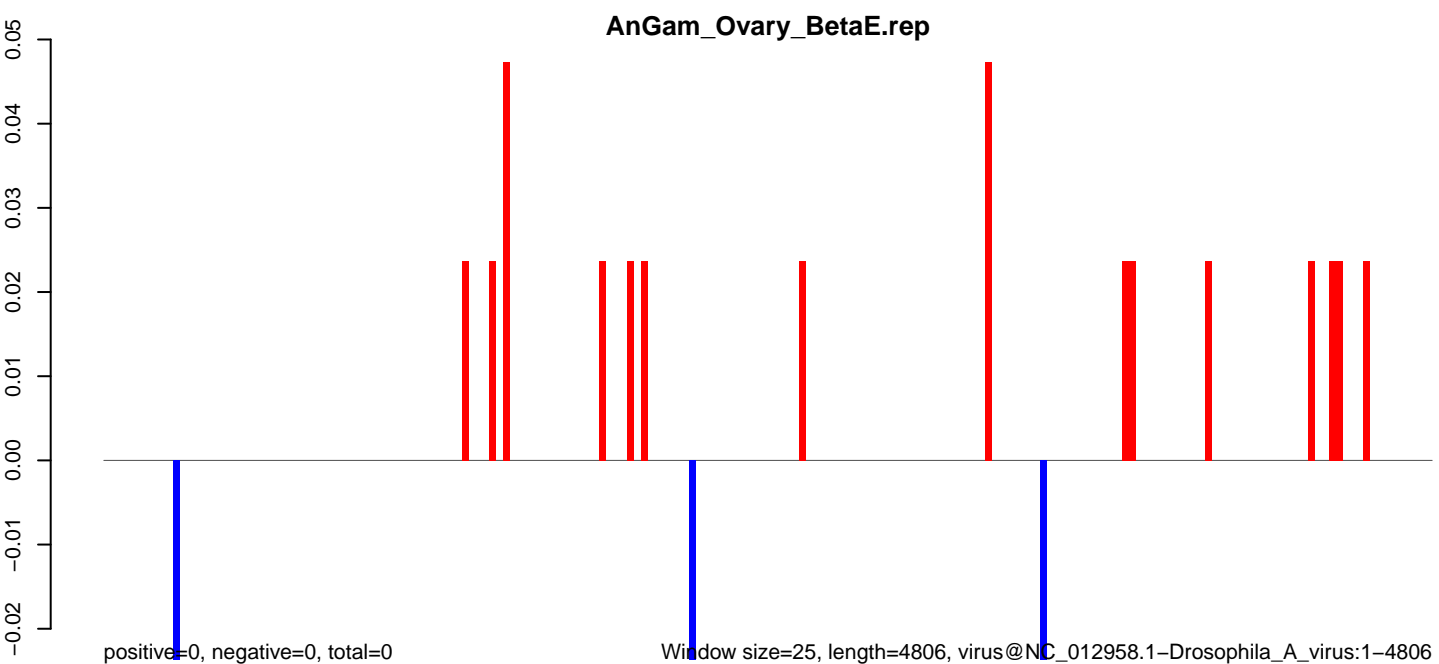
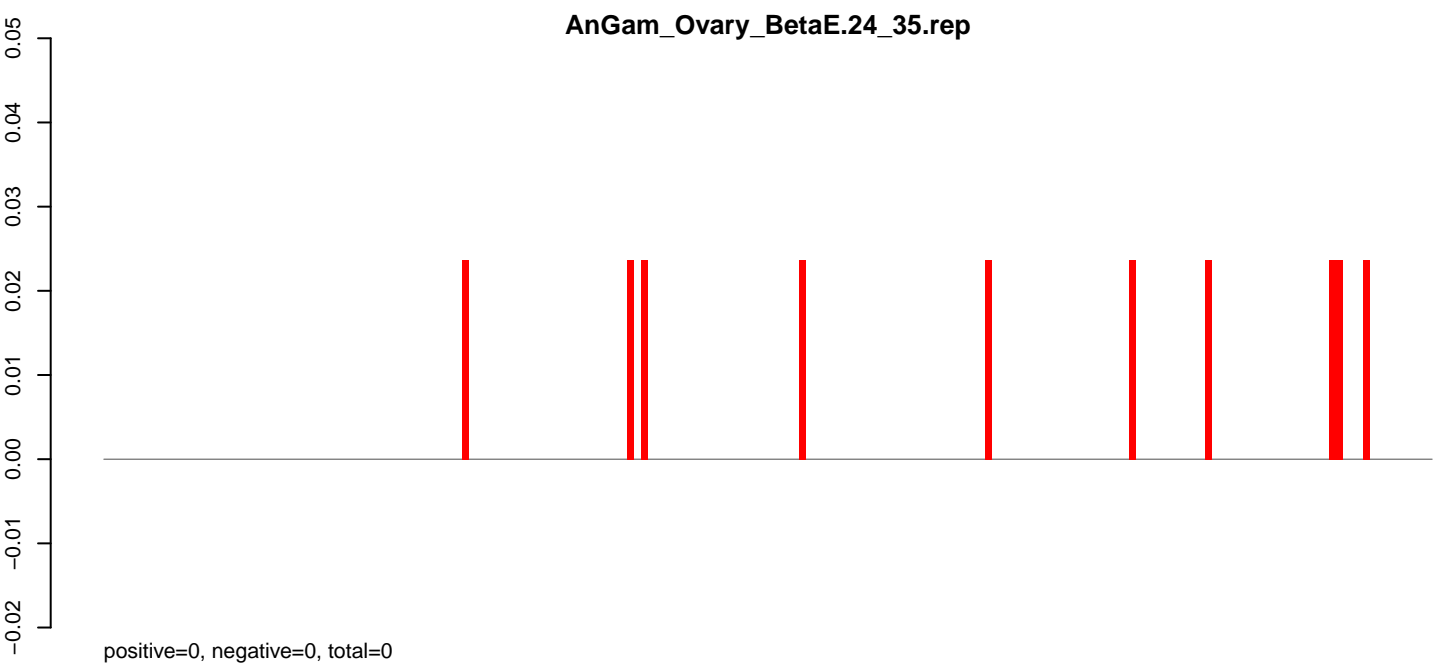
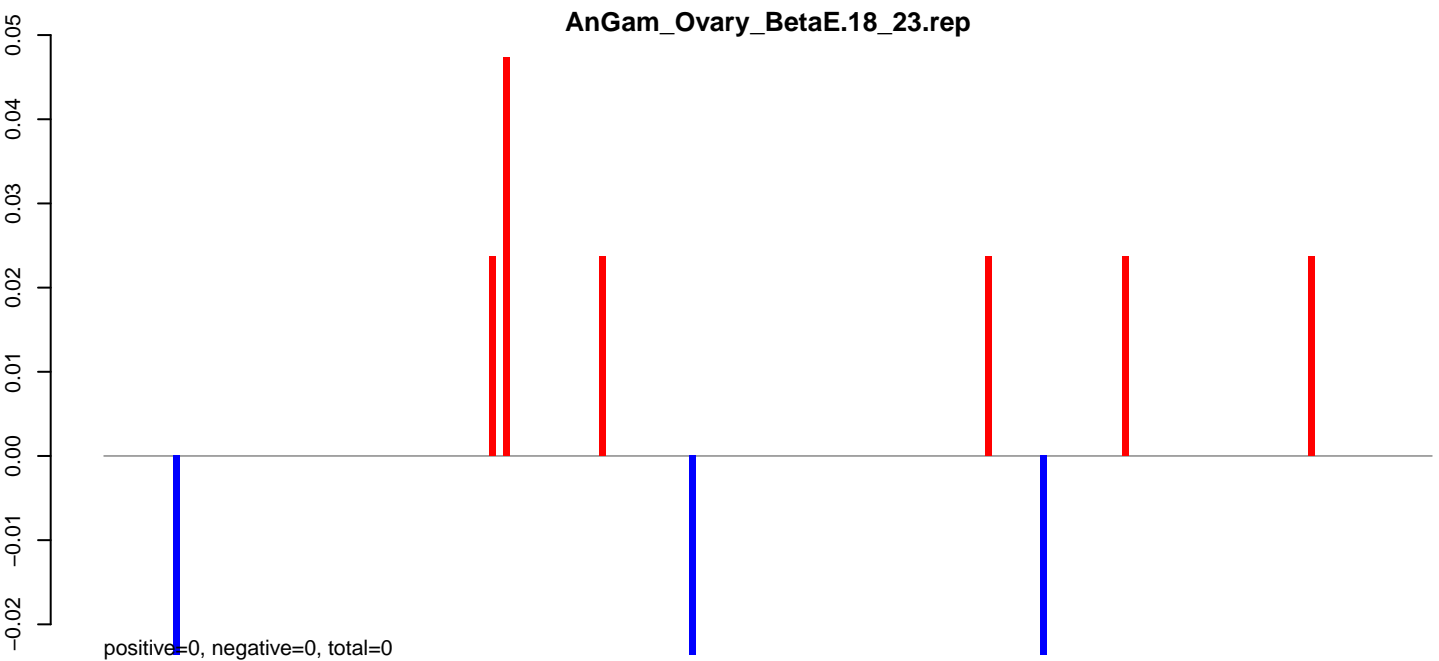
AnGam_Ovary_BetaE.rep



0 2000 4000 6000 8000

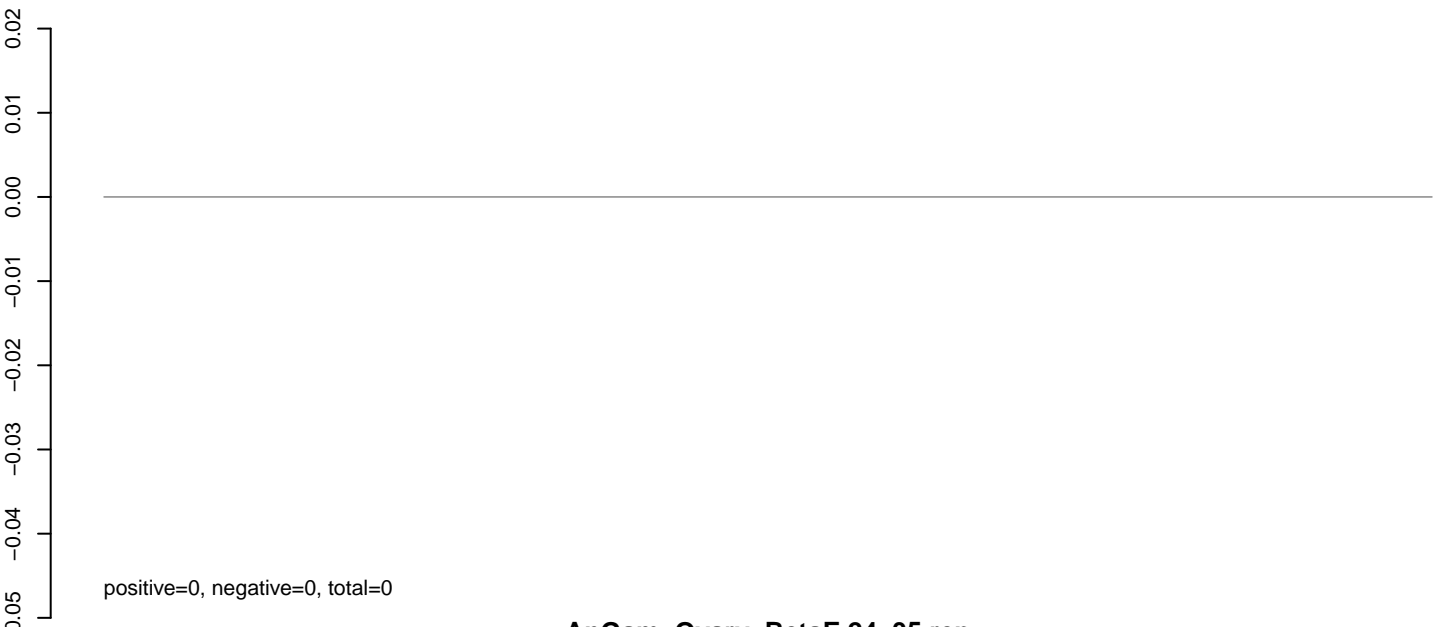


ze=25 positive=4, negative=3, total=7
 0 1000 2000 3000 4000
 Drosophila_melanogaster_American_nodavirus_ANV_SW-2009a_segment_RNA1_and_RNA2:1-4523

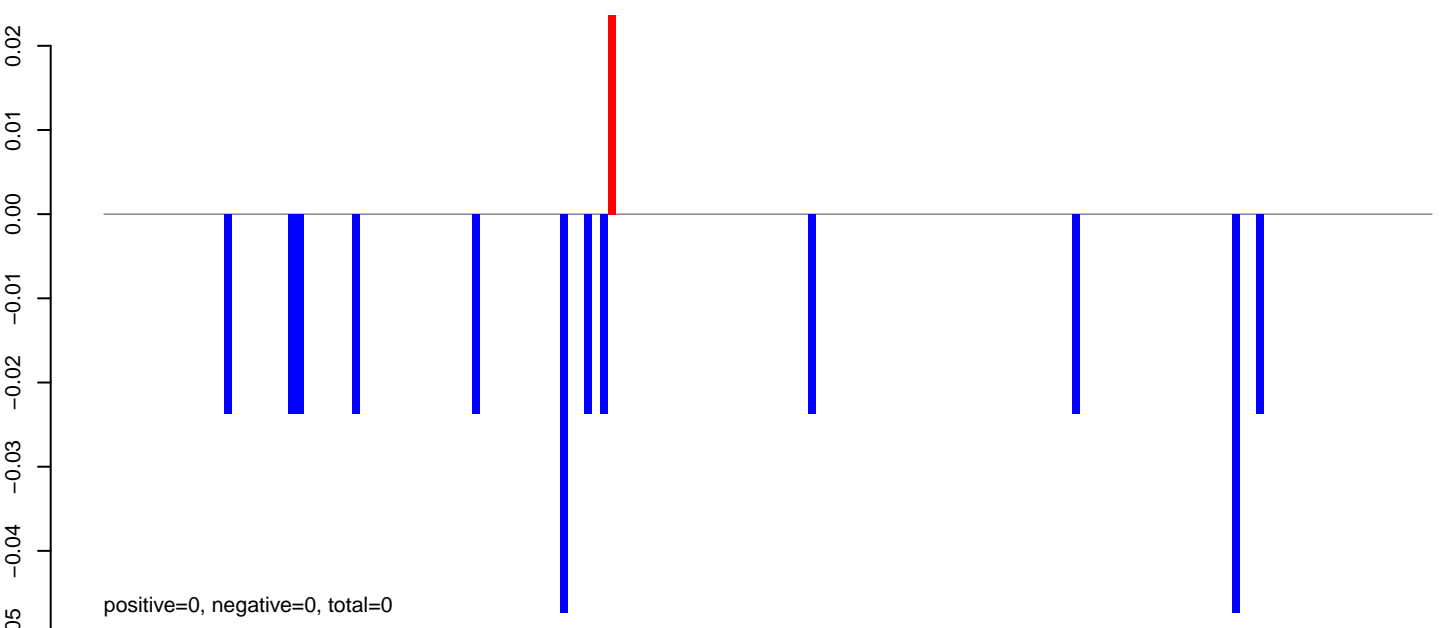


0 1000 2000 3000 4000 5000

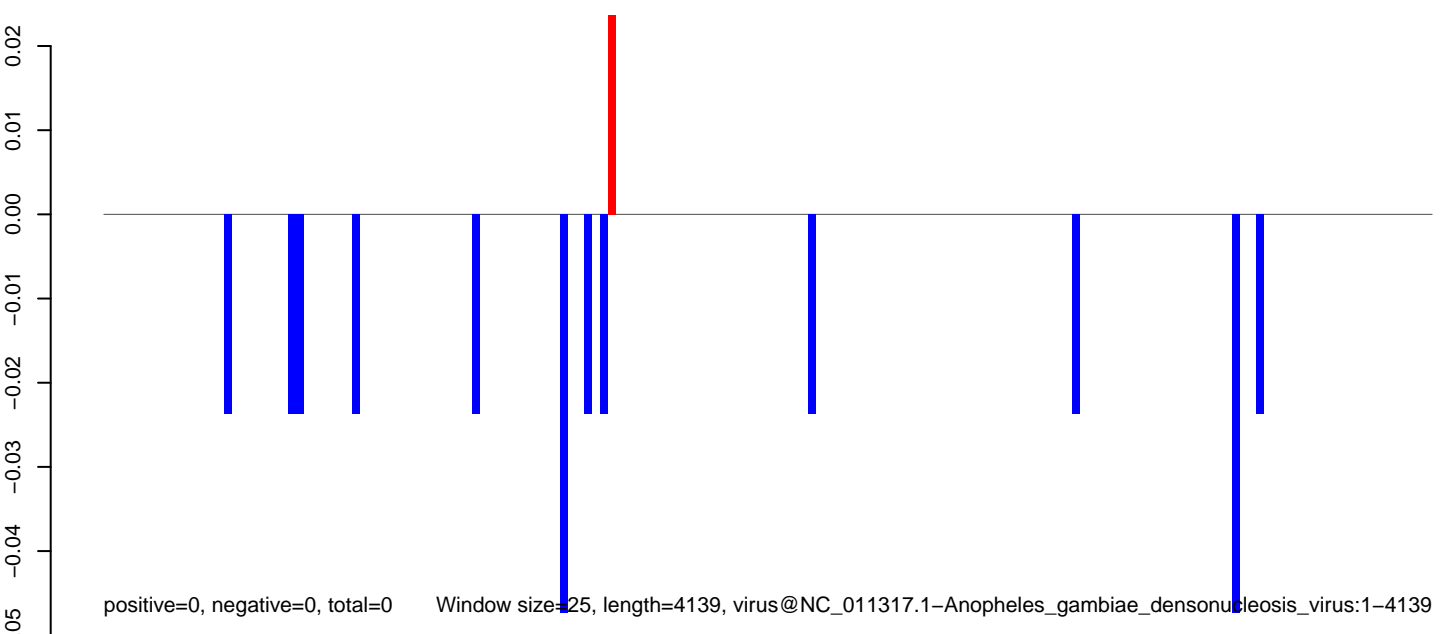
AnGam_Ovary_BetaE.18_23.rep



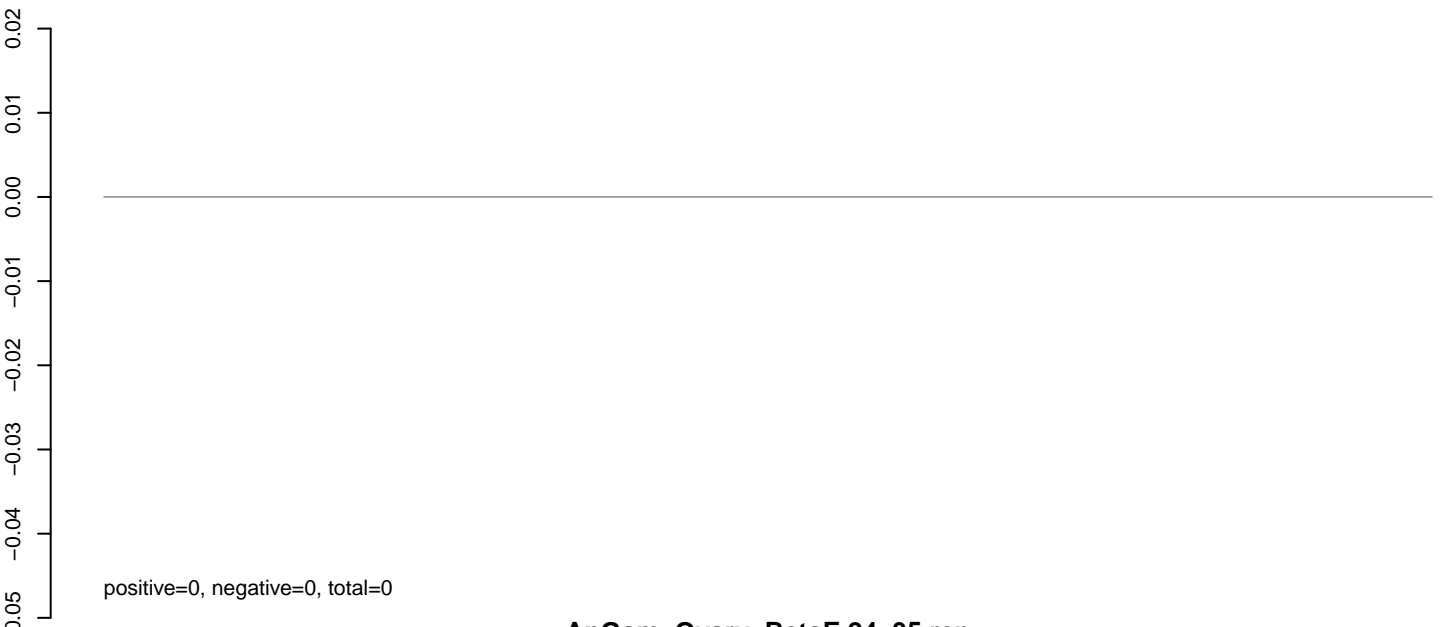
AnGam_Ovary_BetaE.24_35.rep



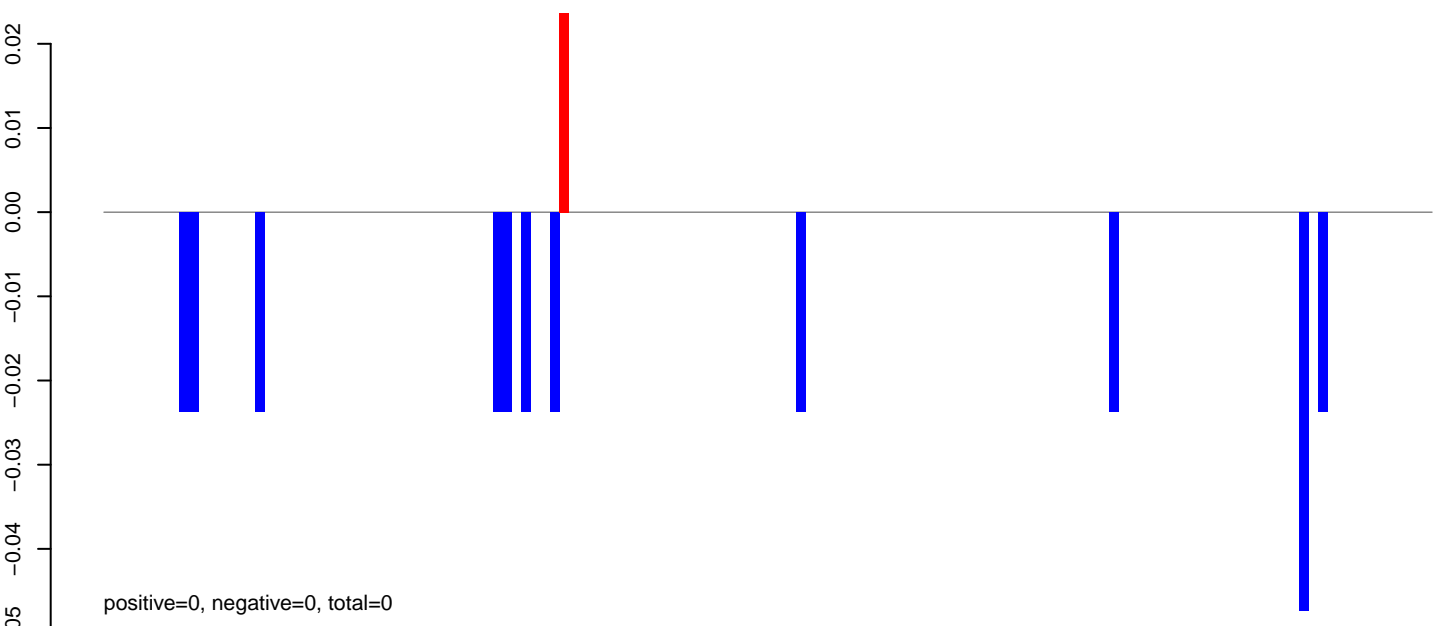
AnGam_Ovary_BetaE.rep



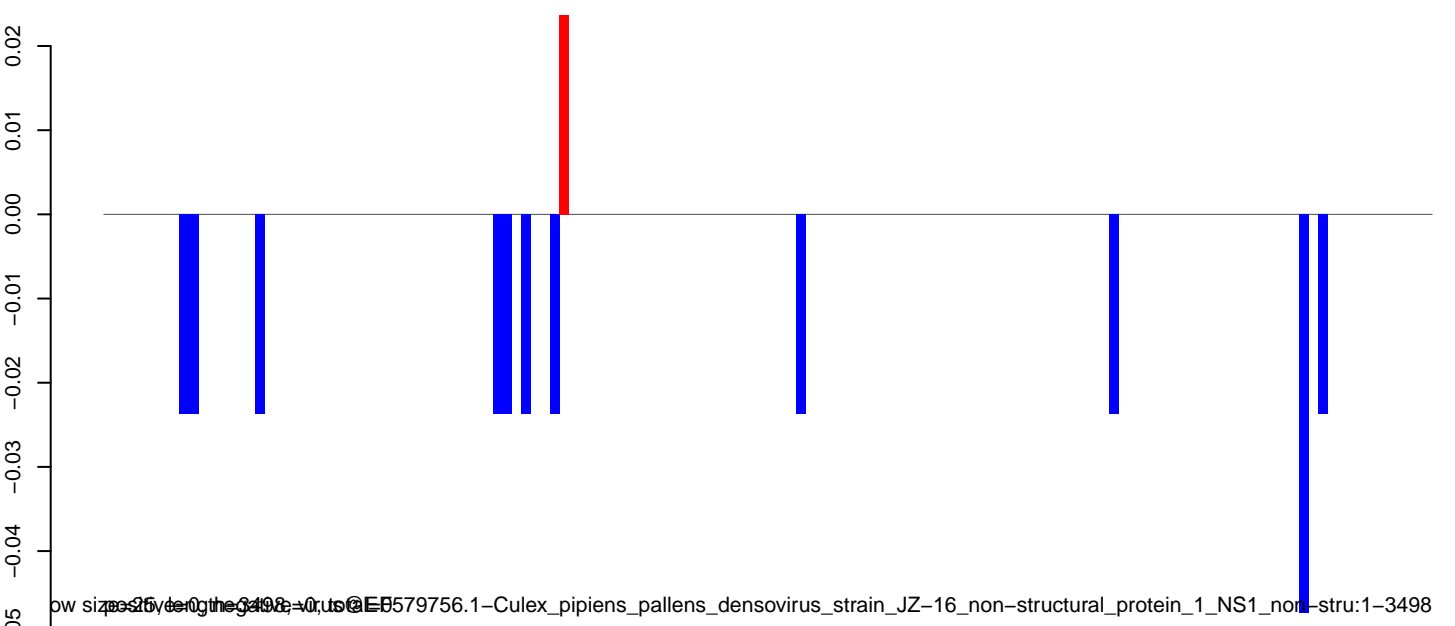
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

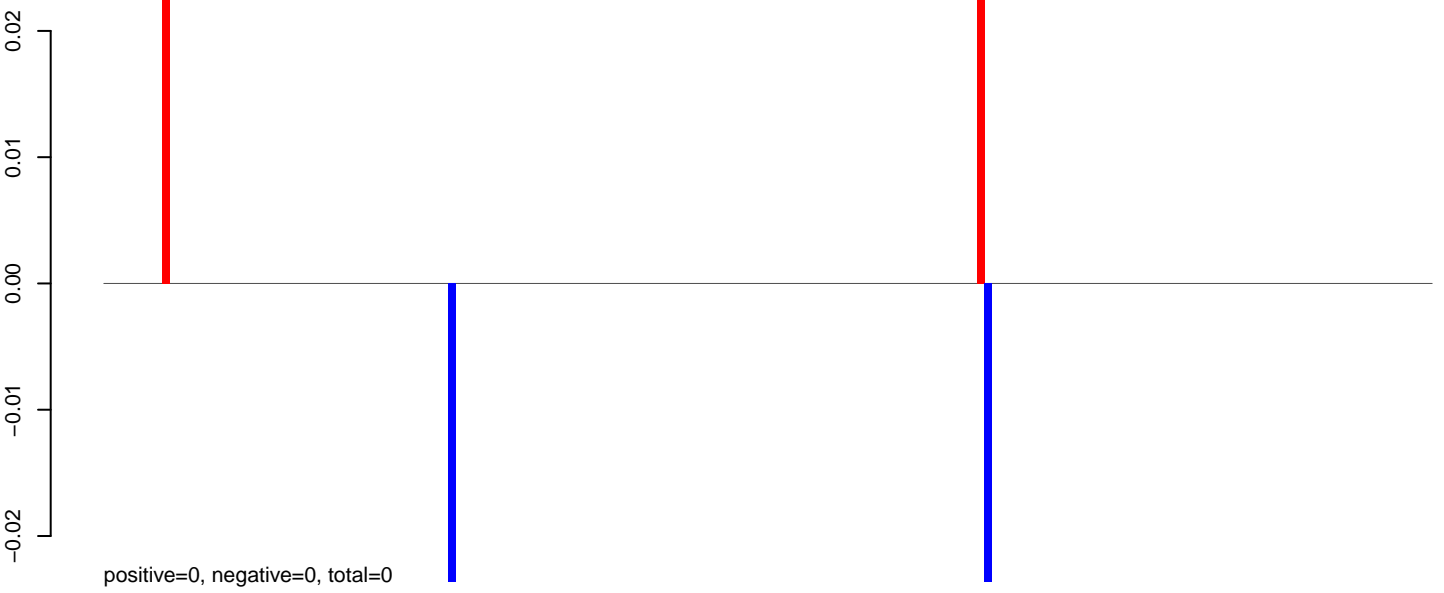


AnGam_Ovary_BetaE.rep

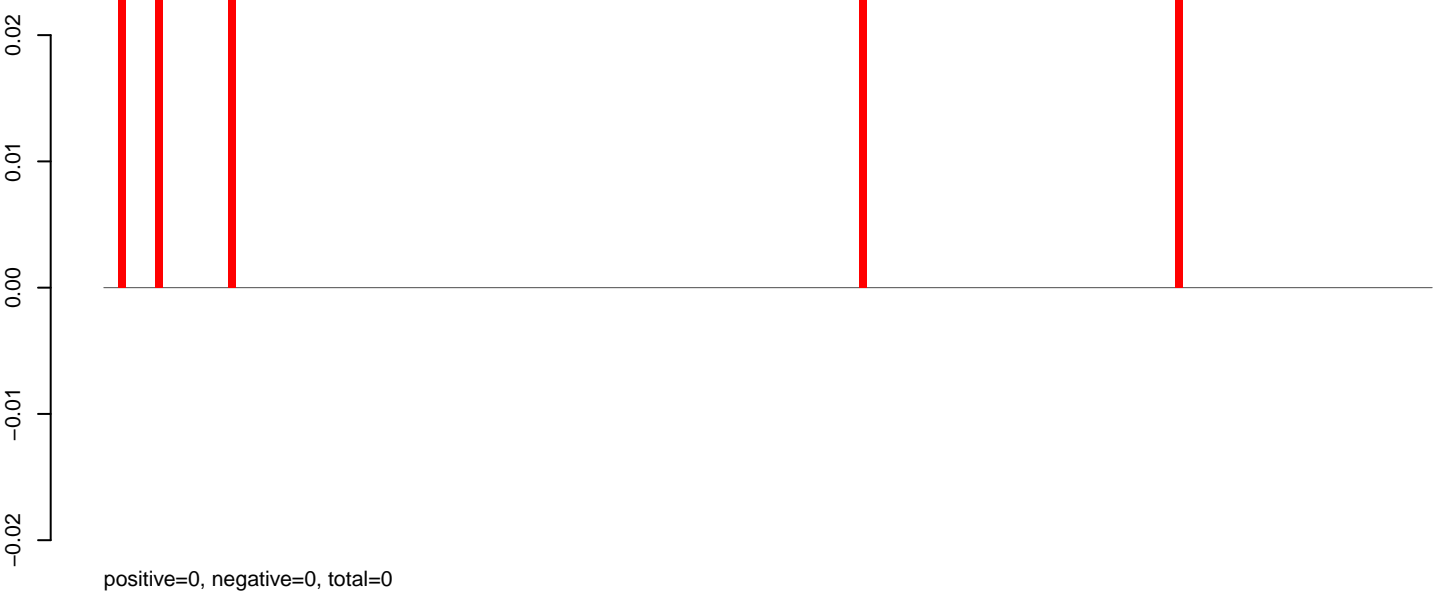


bow size=25, the 3498, use @EB579756.1-Culex_pipiens_pallens_densovirus_strain_JZ-16_non-structural_protein_1_NS1_non-stru:1-3498

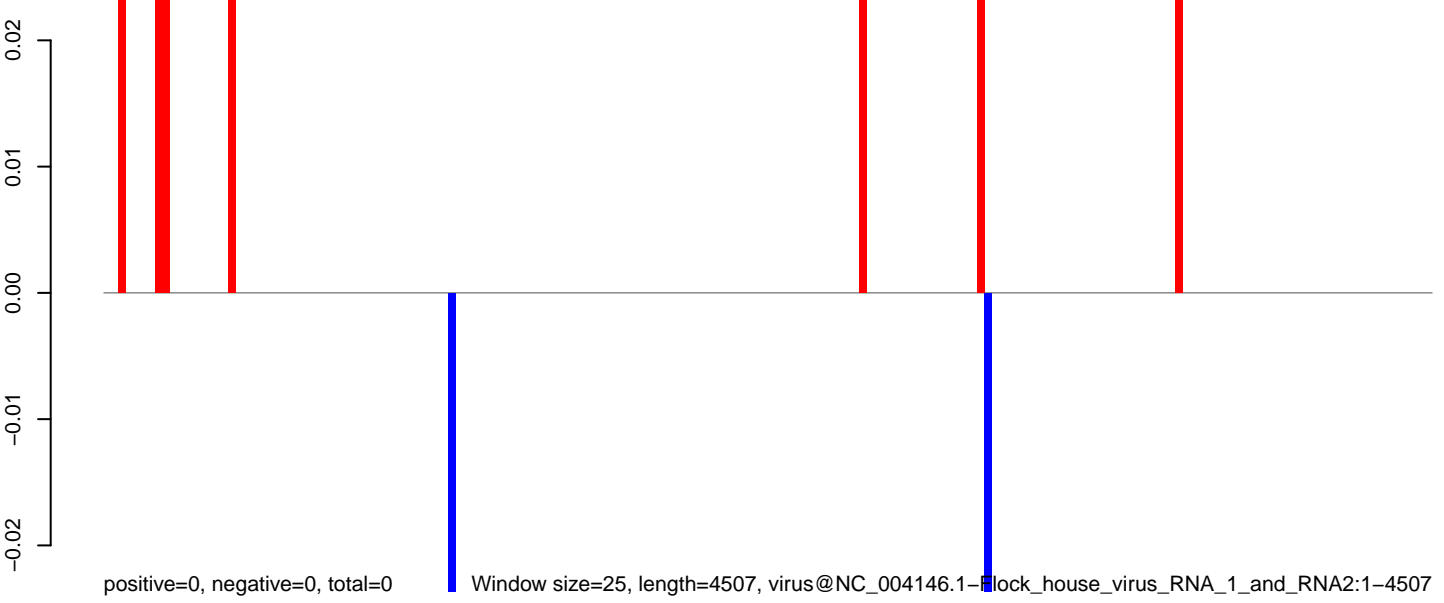
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

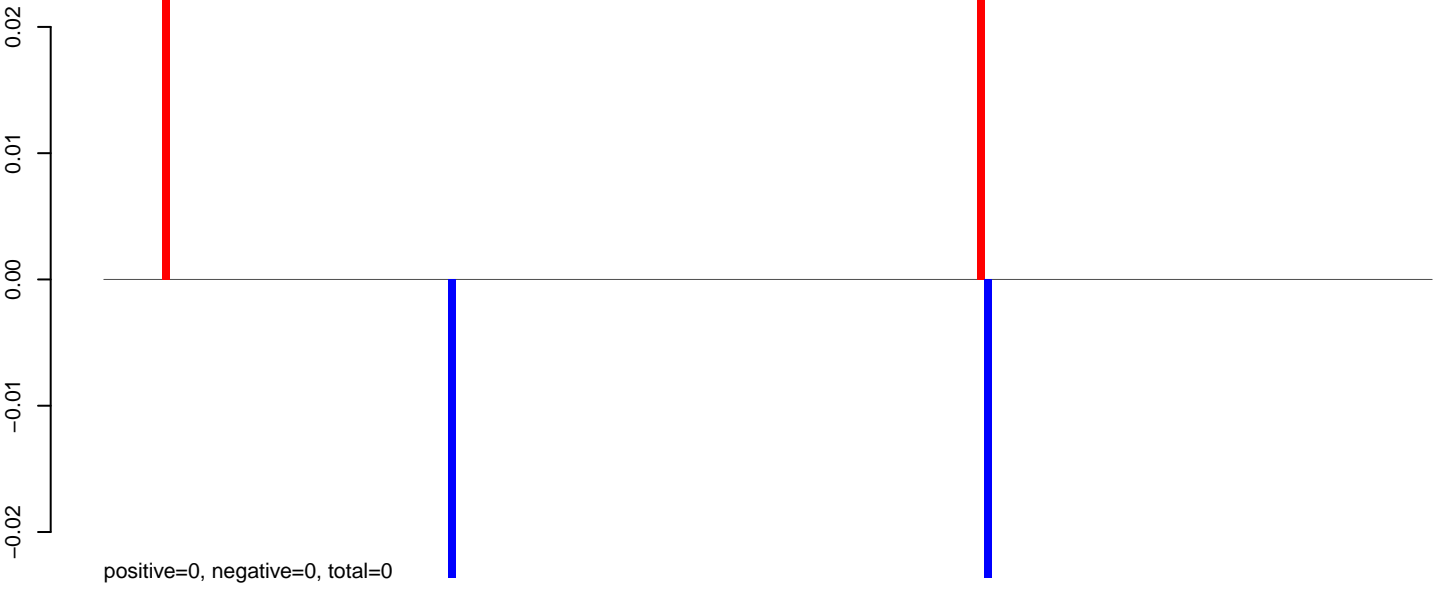


AnGam_Ovary_BetaE.rep



0 1000 2000 3000 4000

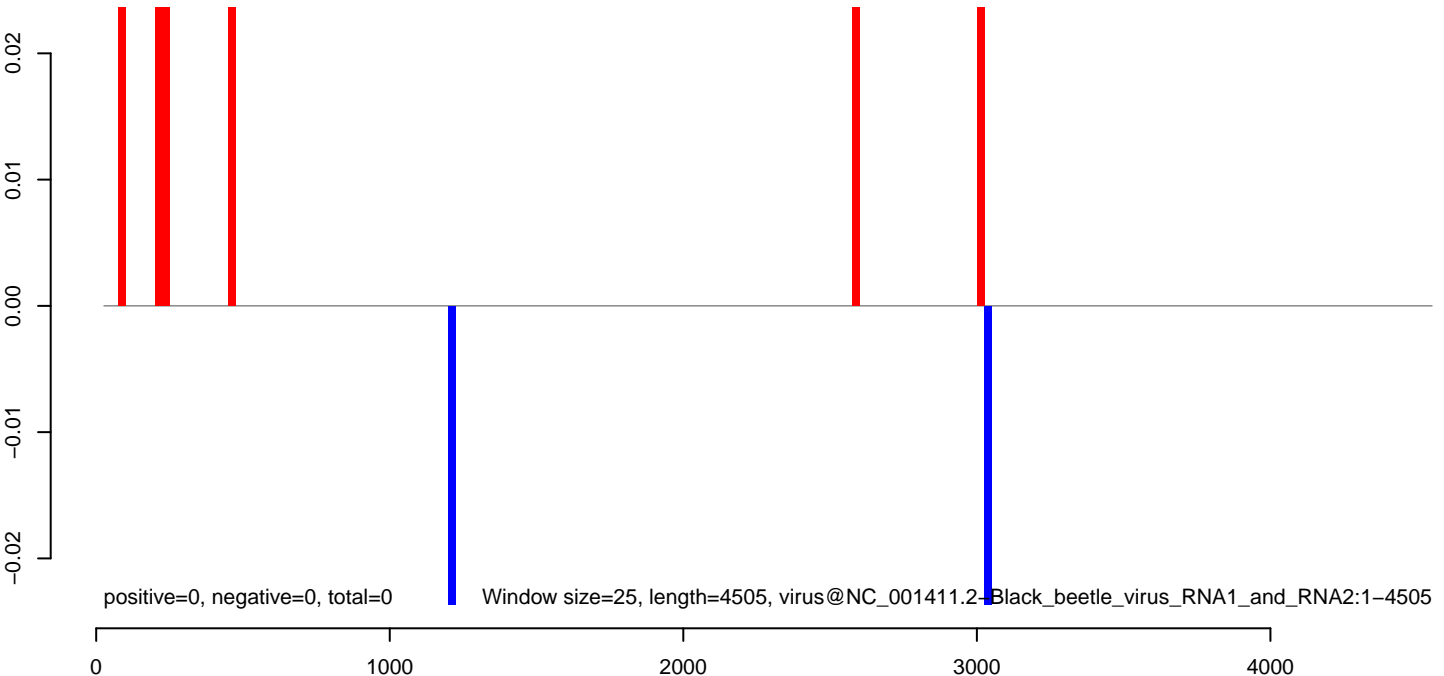
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



AnGam_Ovary_BetaE.18_23.rep

0.00
-0.01
-0.02
-0.03
-0.04

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep

0.00
-0.01
-0.02
-0.03
-0.04

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep

0.00
-0.01
-0.02
-0.03
-0.04

positive=0, negative=0, total=0

Window size=25, length=3805, virus@MH188045.1-Culex_densovirus_strain_CDENV_Madera-4:1-3805

0 1000 2000 3000 4000

AnGam_Ovary_BetaE.18_23.rep

0.00
-0.01
-0.02
-0.03
-0.04

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep

0.00
-0.01
-0.02
-0.03
-0.04

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep

0.00
-0.01
-0.02
-0.03
-0.04

positive=0, negative=0, total=0

Window size=25, length=3991, virus@MH188044.1-Culex_densovirus_strain_CDenV_Madera-3:1-3991

0 1000 2000 3000 4000

AnGam_Ovary_BetaE.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

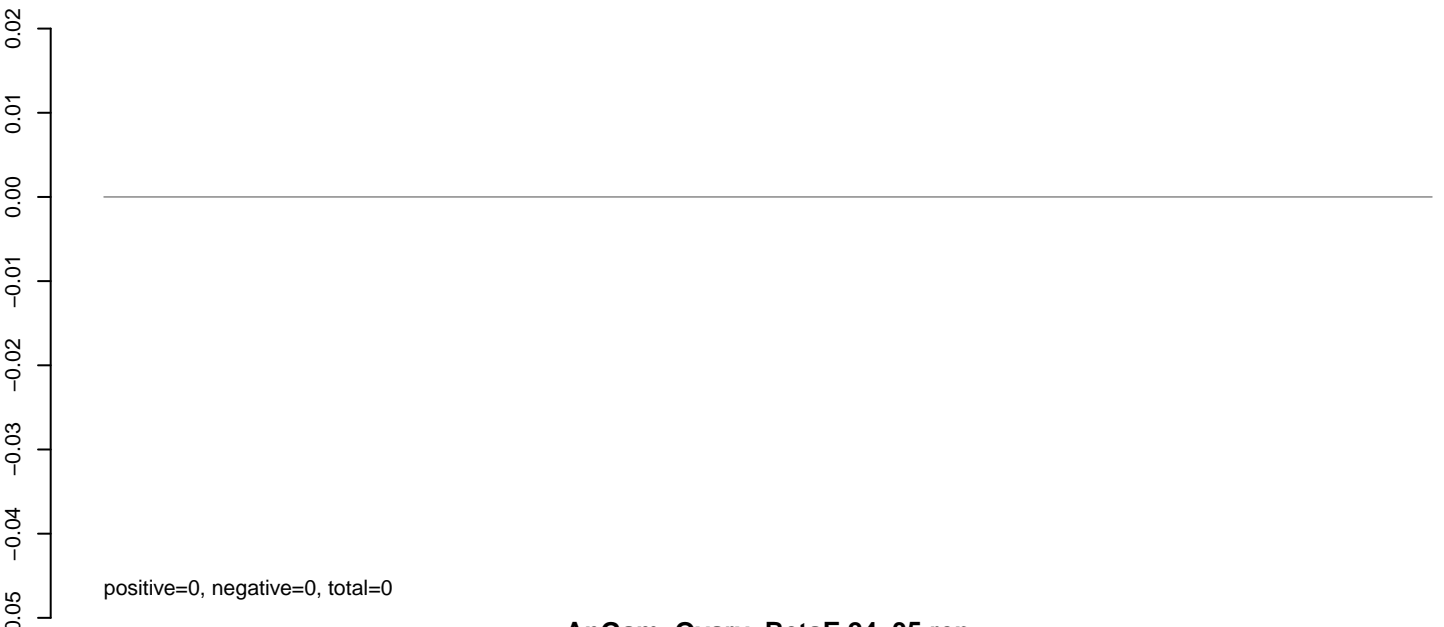
AnGam_Ovary_BetaE.rep

0.000
-0.005
-0.010
-0.015
-0.020

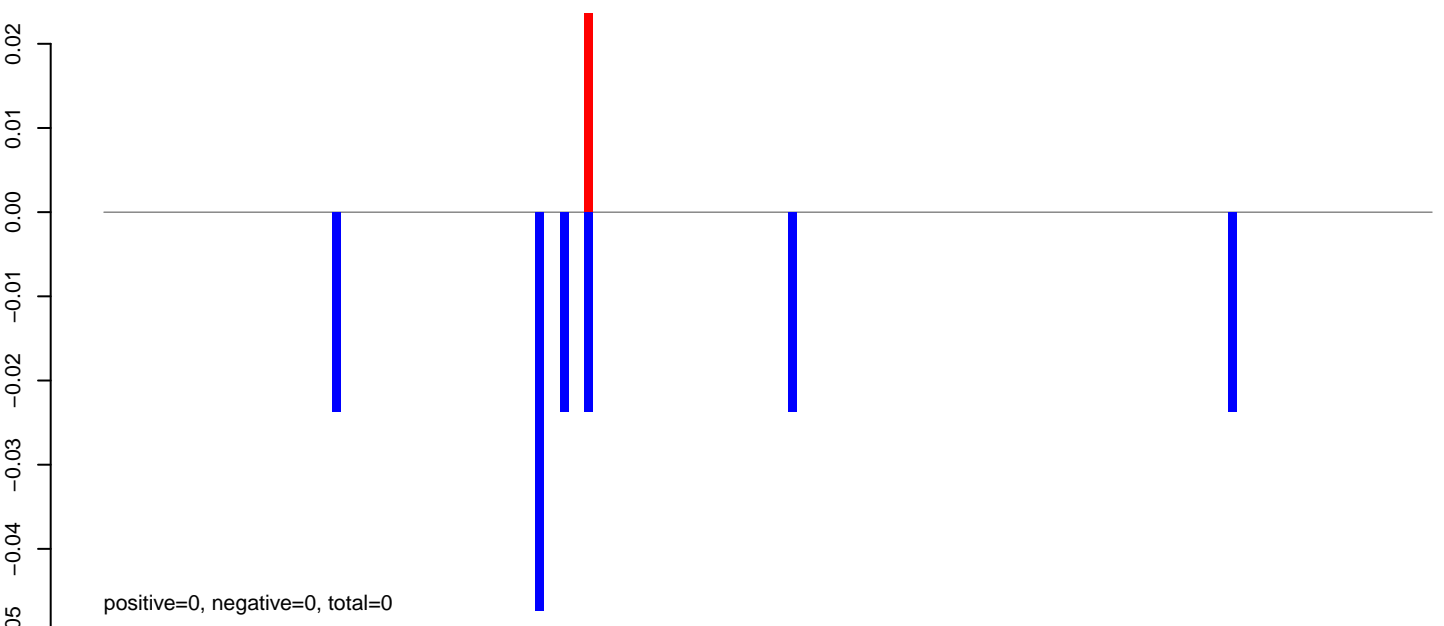
positive=0, negative=0, total=0 Window size=25, length=3776, virus@NC_012636.1-Aedes_aegypti_densovirus_2_strain_0814616:1-3776

0 1000 2000 3000

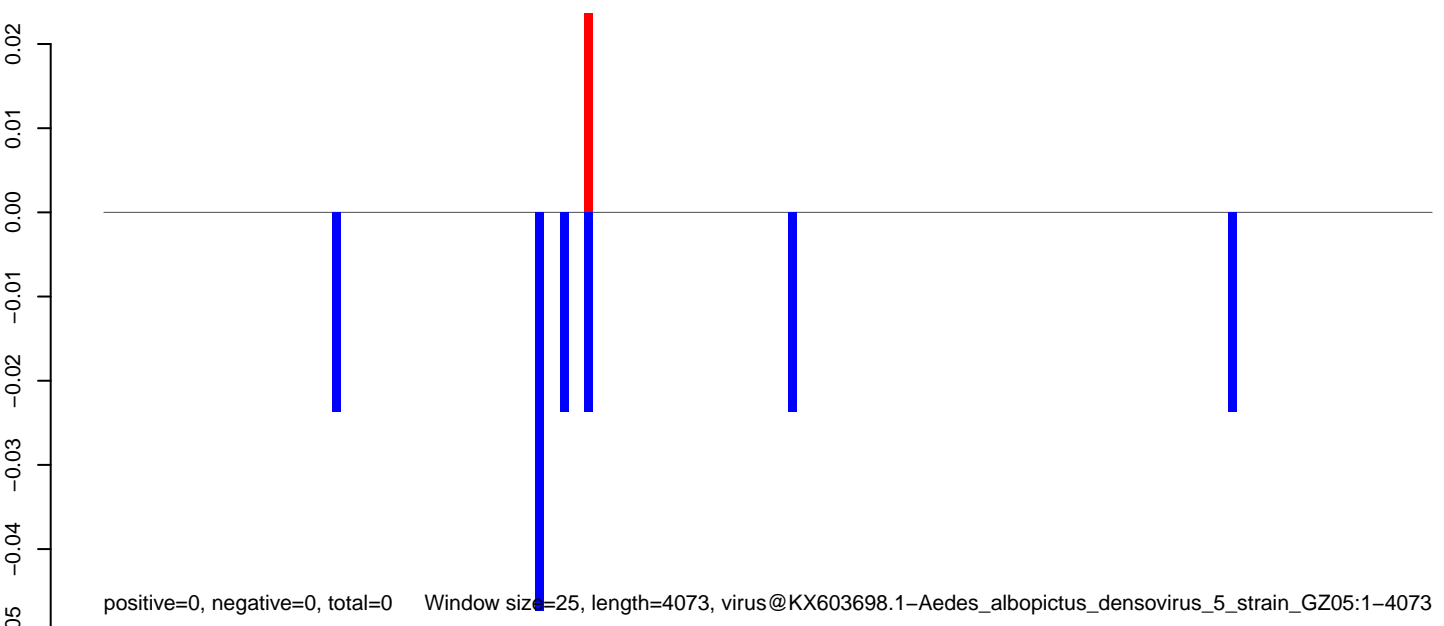
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

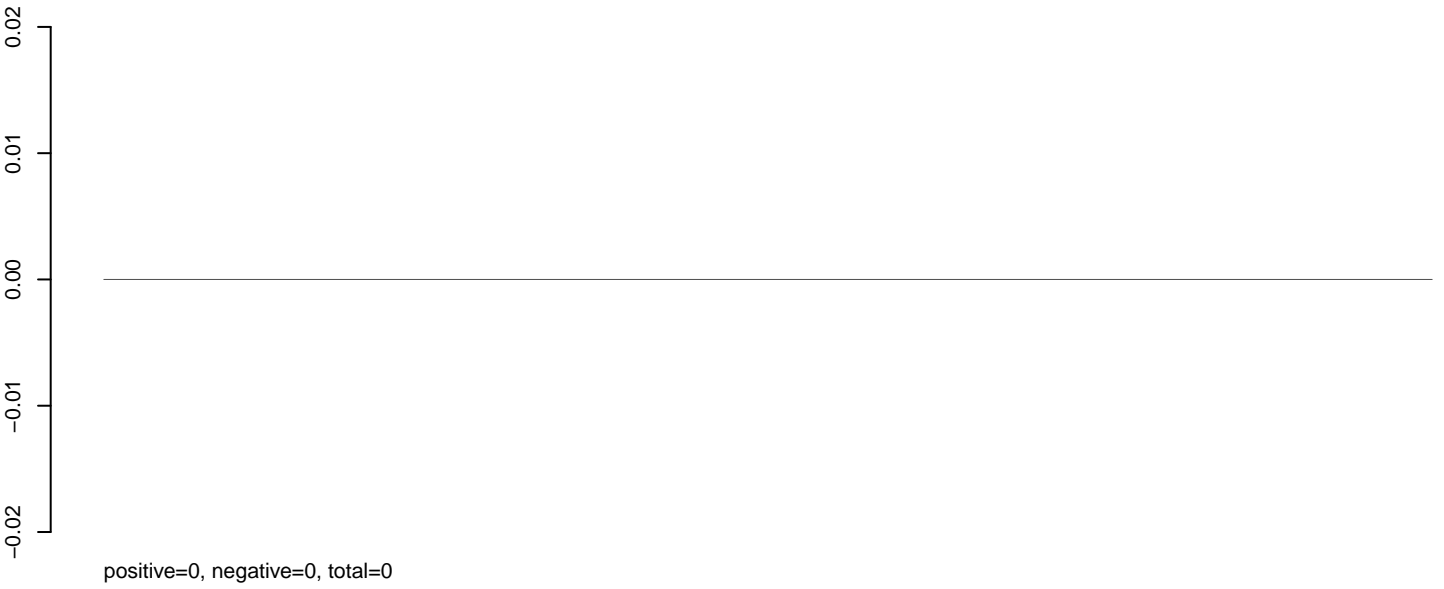


AnGam_Ovary_BetaE.rep

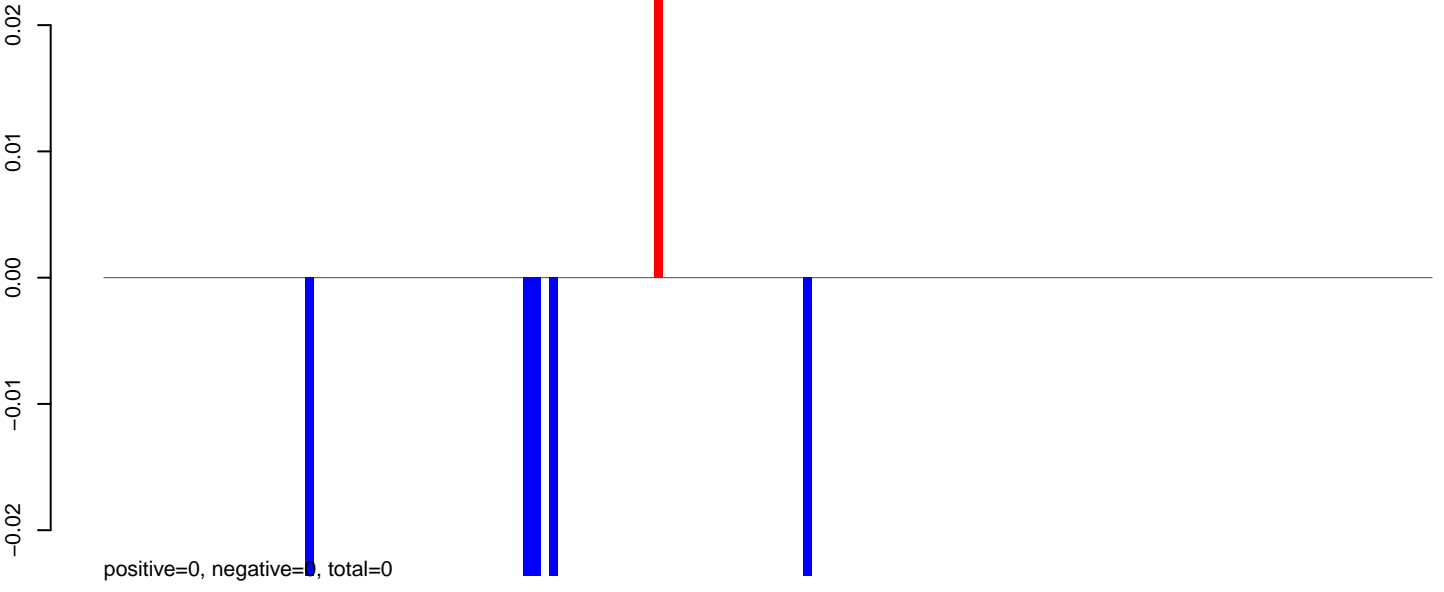


0 1000 2000 3000 4000

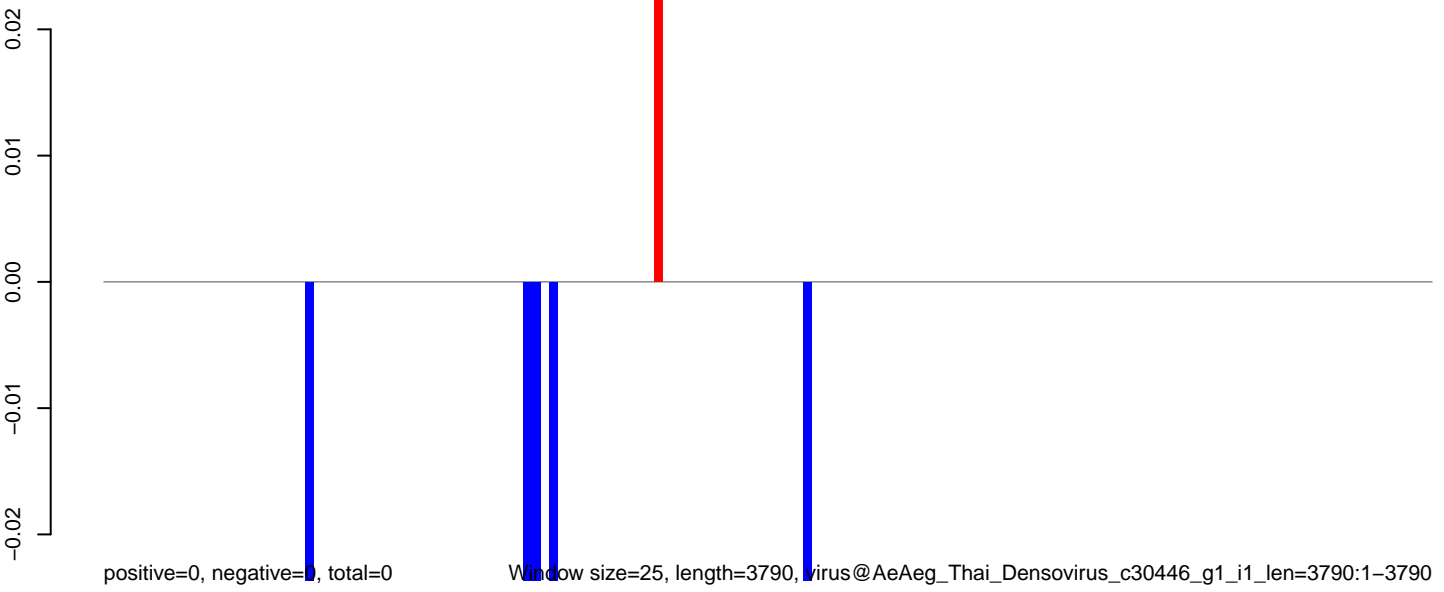
AnGam_Ovary_BetaE.18_23.rep



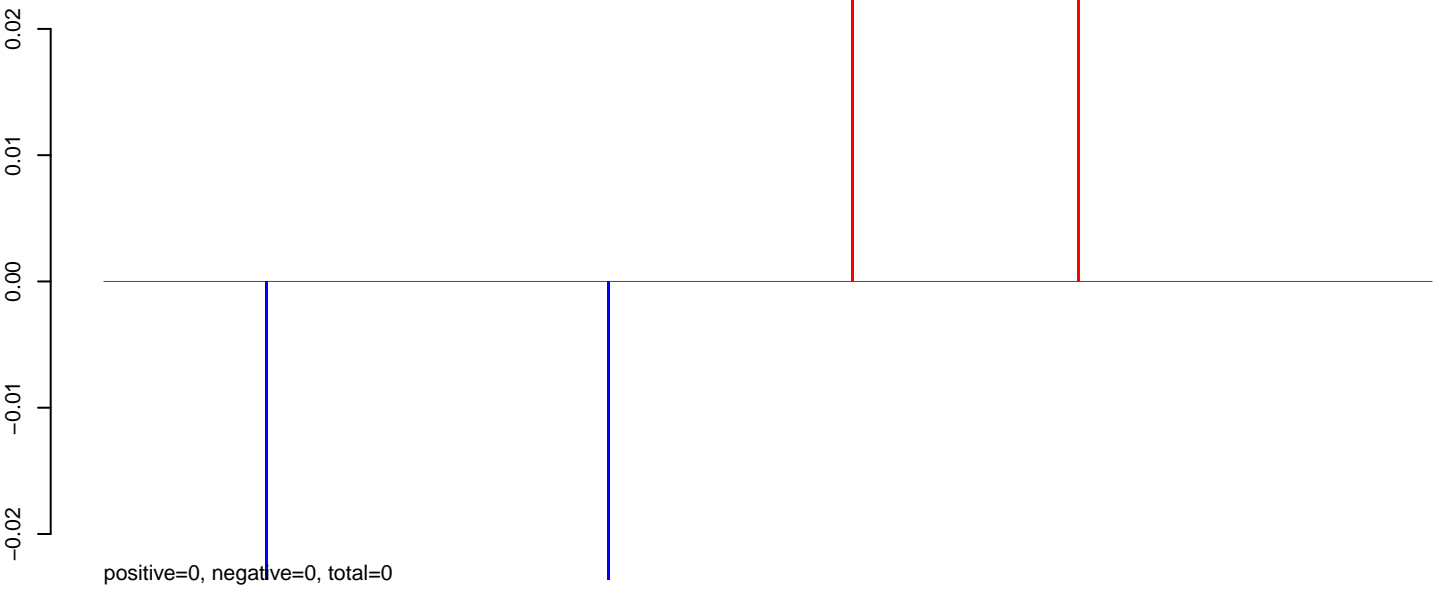
AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



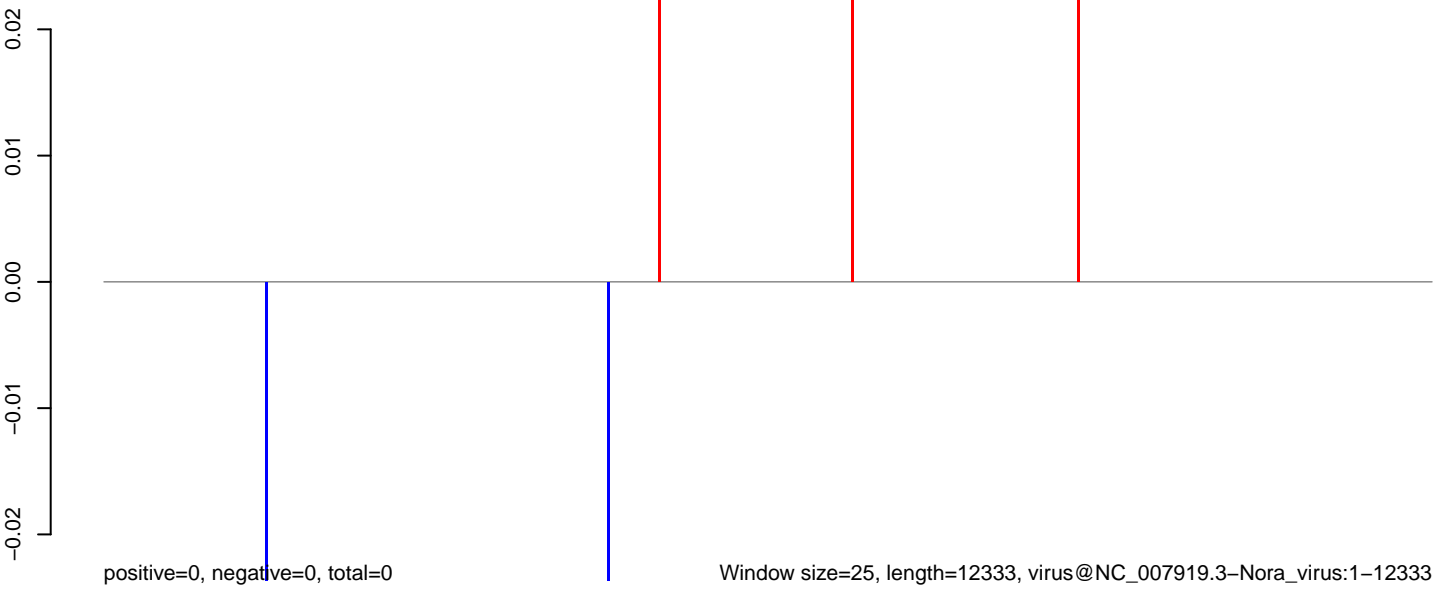
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

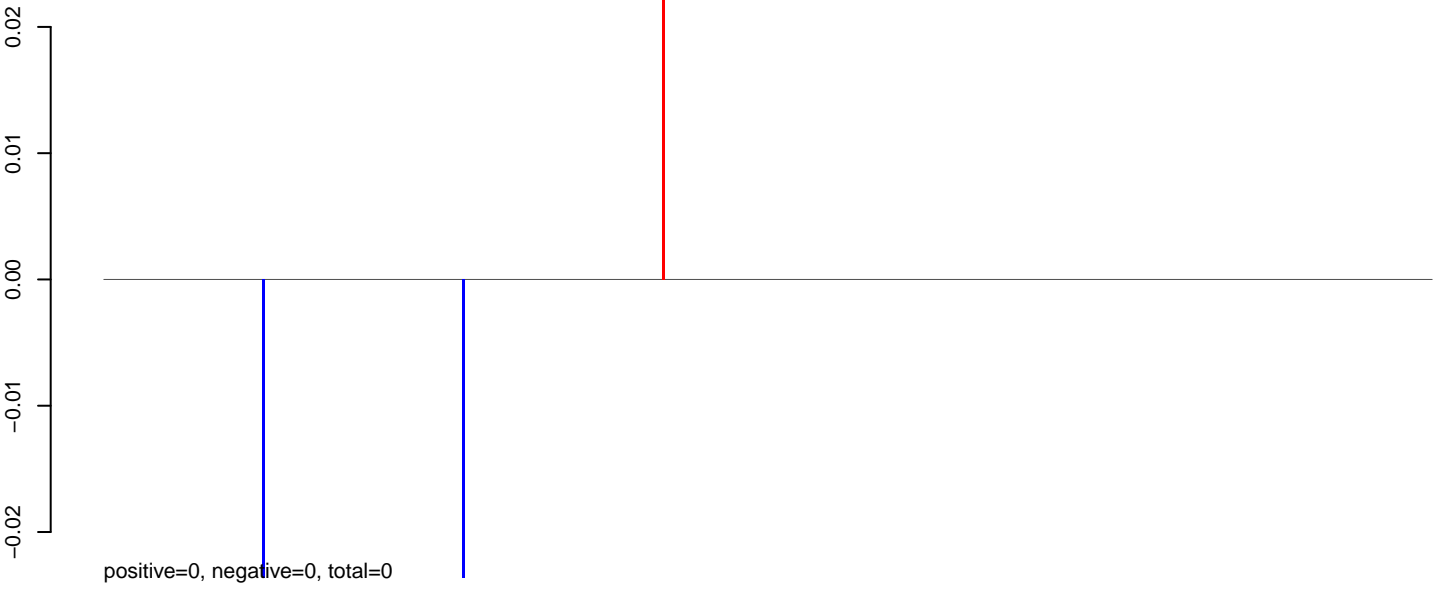


AnGam_Ovary_BetaE.rep

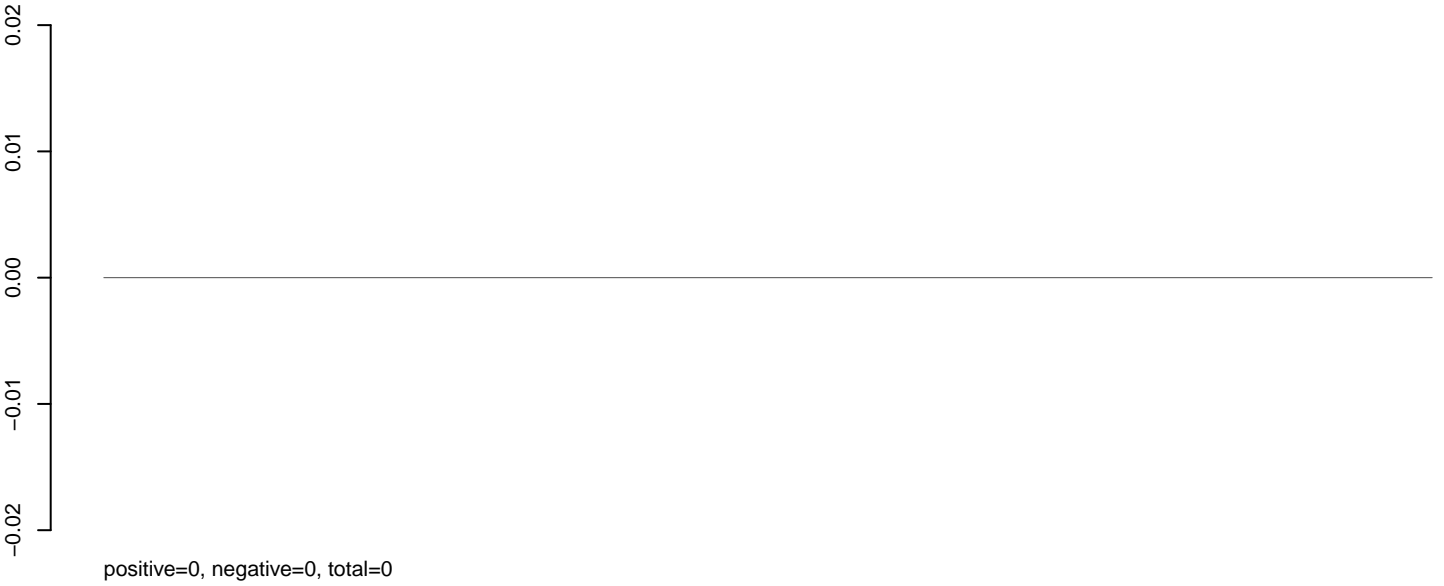


0 2000 4000 6000 8000 10000 12000

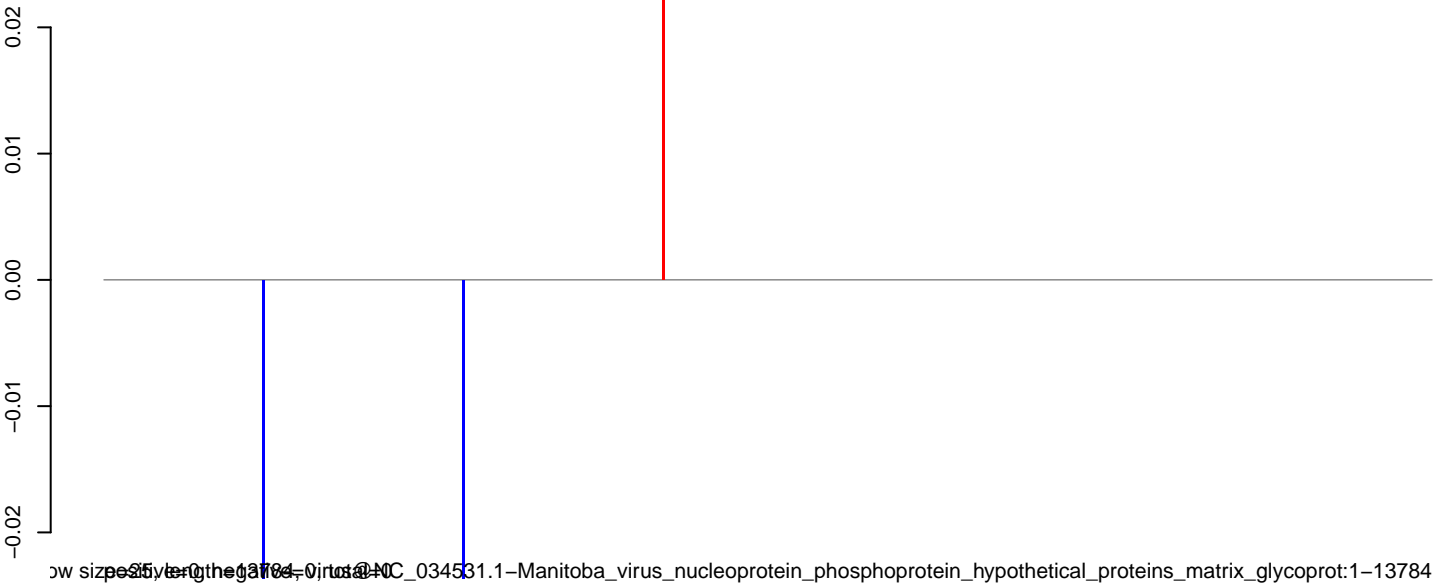
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



AnGam_Ovary_BetaE.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

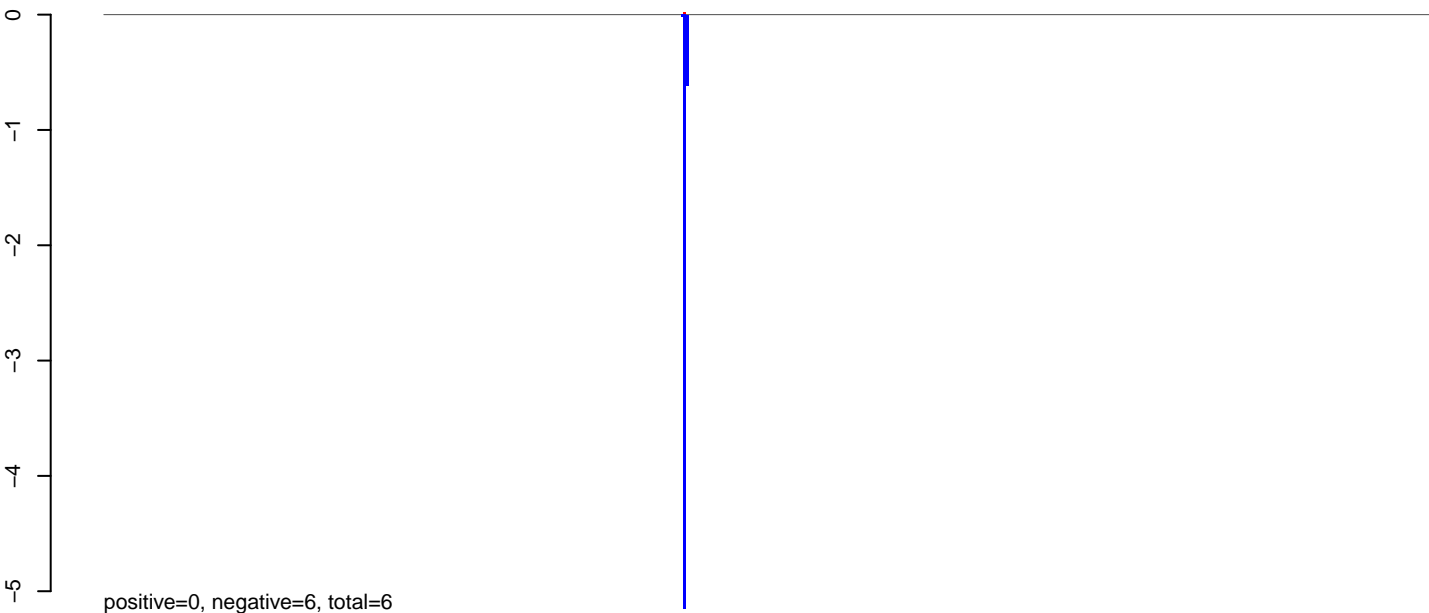
AnGam_Ovary_BetaE.rep

0.000
-0.005
-0.010
-0.015
-0.020

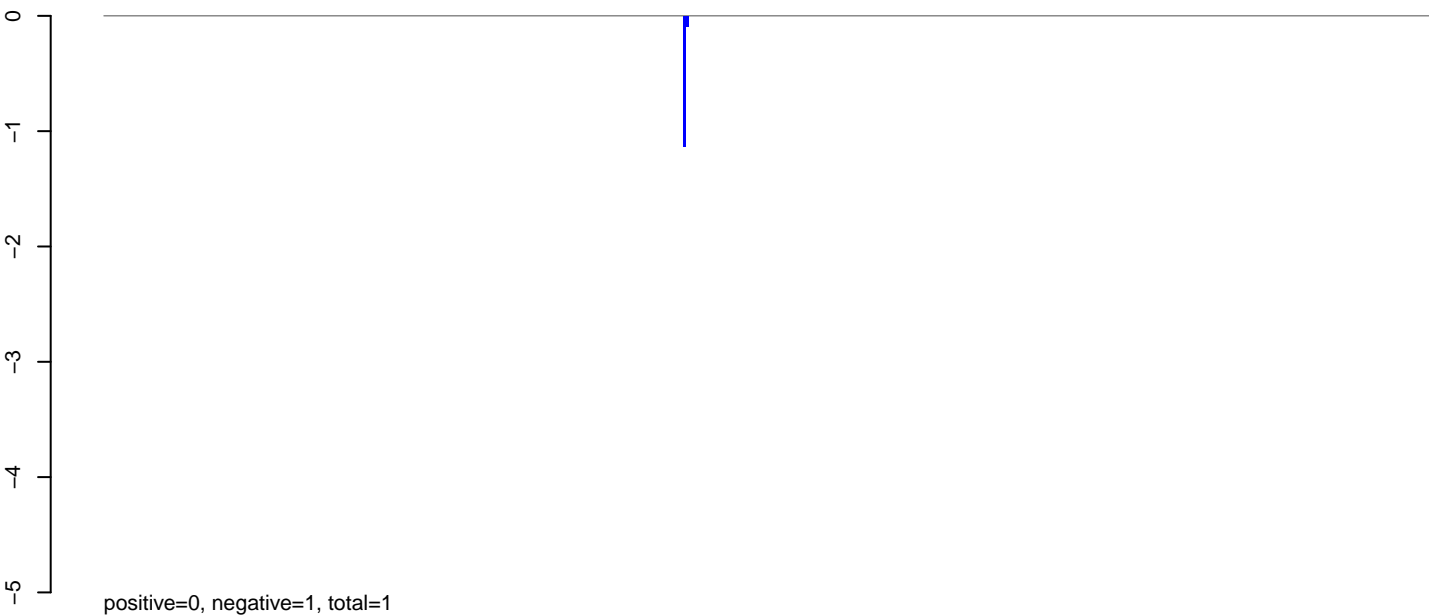
positive=0, negative=0, total=0 Window size=25, length=3978, virus@M37899.1-Aedes_aegypti_densovirus_1_genomic_sequence:1-3978

0 1000 2000 3000 4000

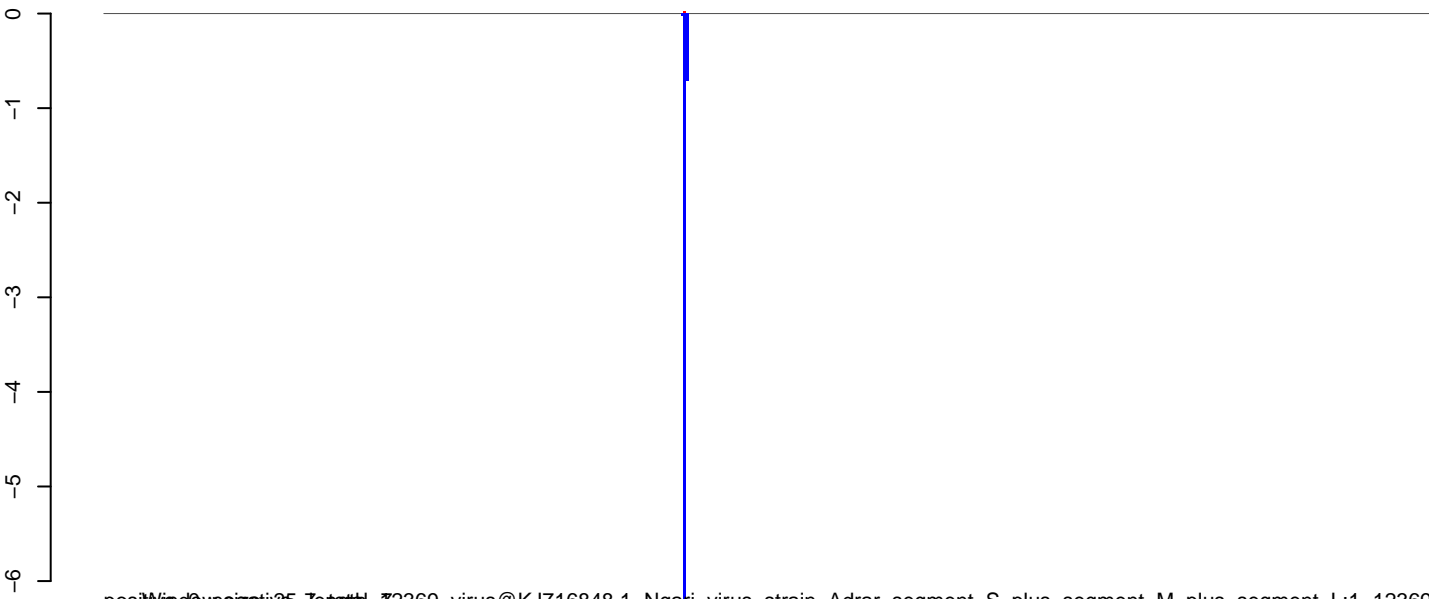
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

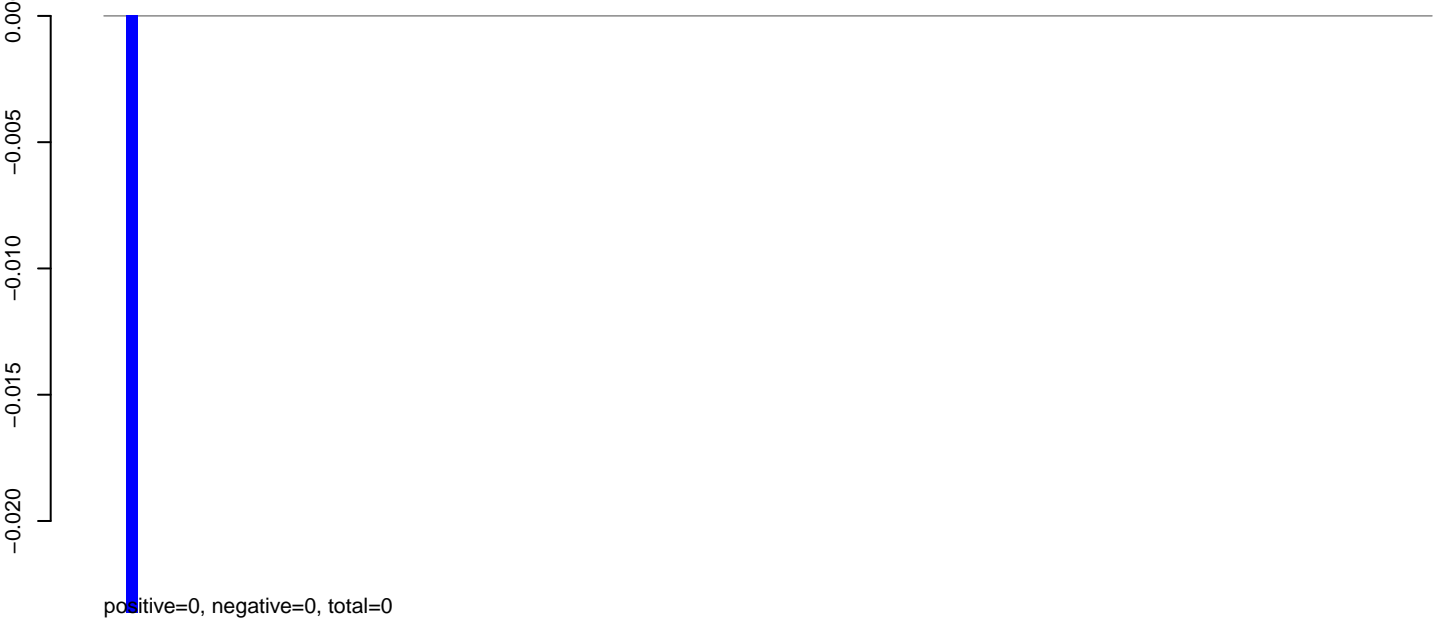


AnGam_Ovary_BetaE.rep

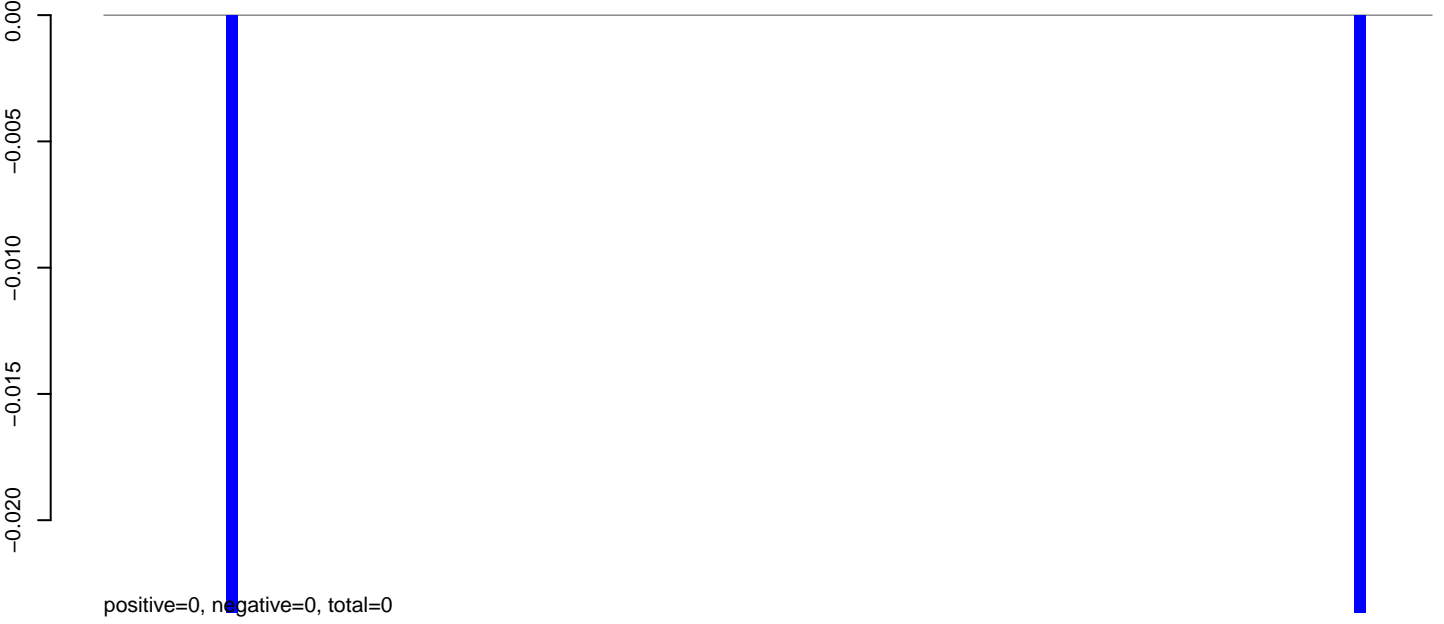


0 2000 4000 6000 8000 10000 12000

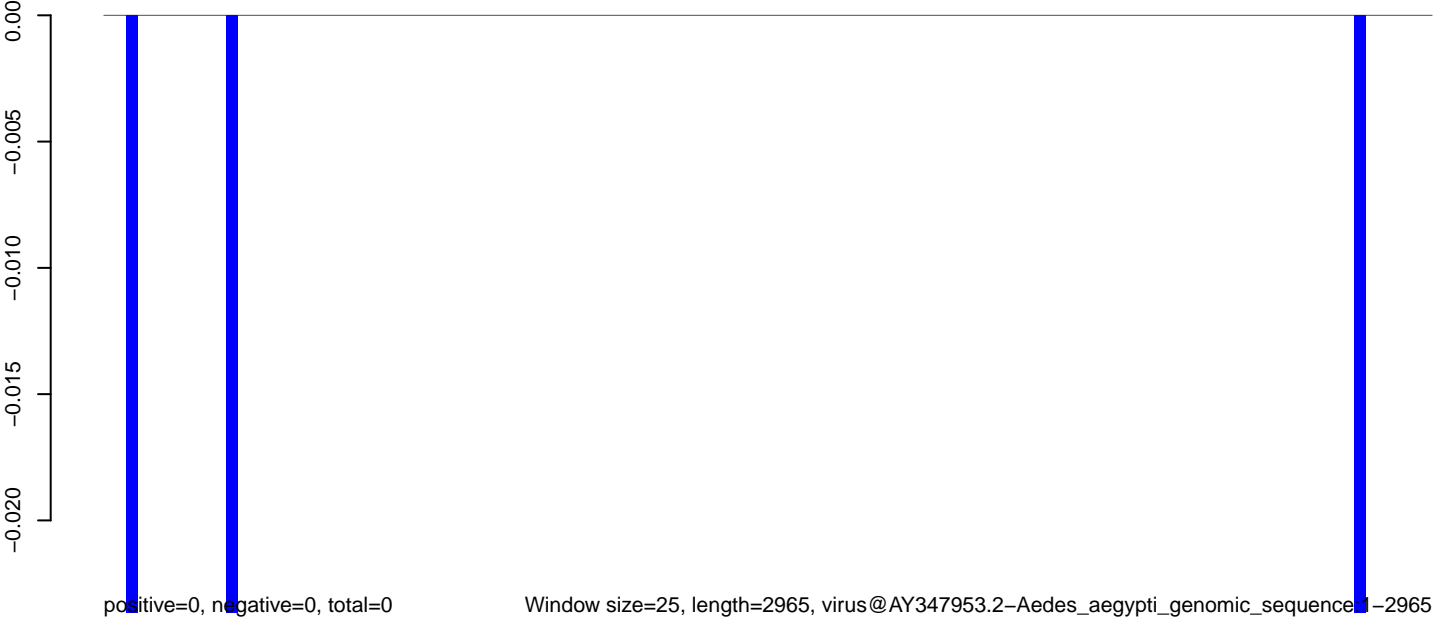
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



AnGam_Ovary_BetaE.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep

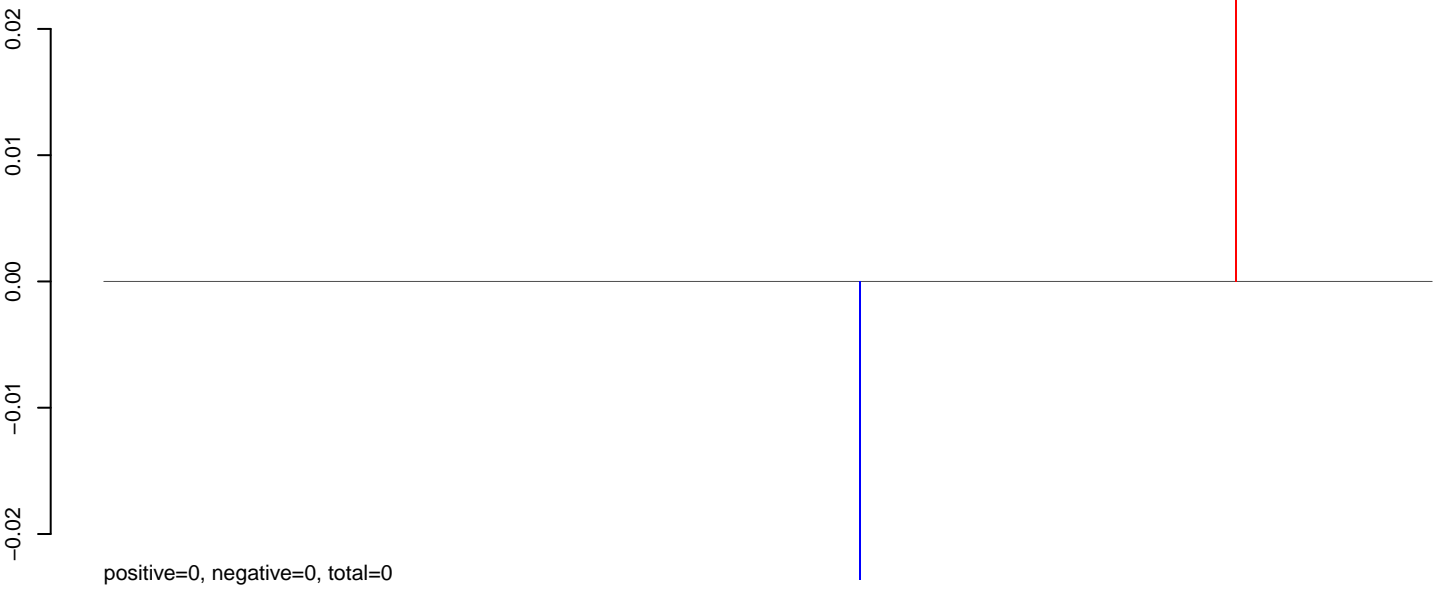
0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

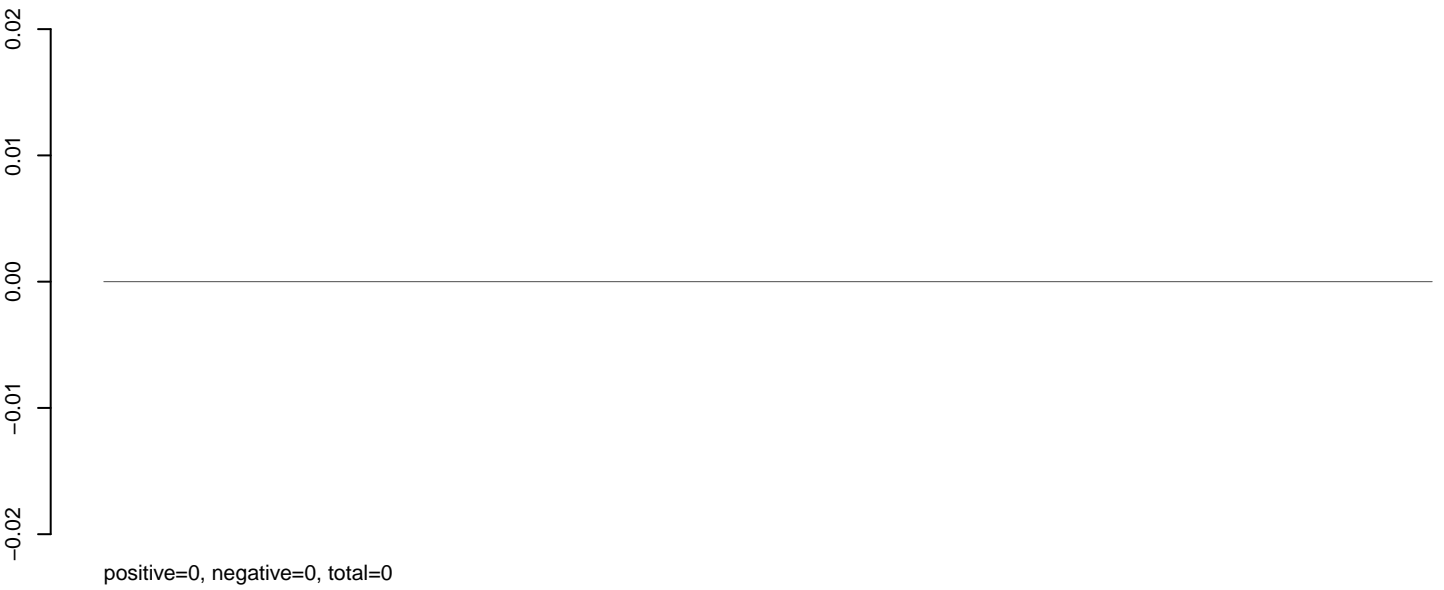
Window size=25, length=4176, virus@X74945.1-Aedes_albopictus_densovirus_2_DNA:1-4176

0 1000 2000 3000 4000

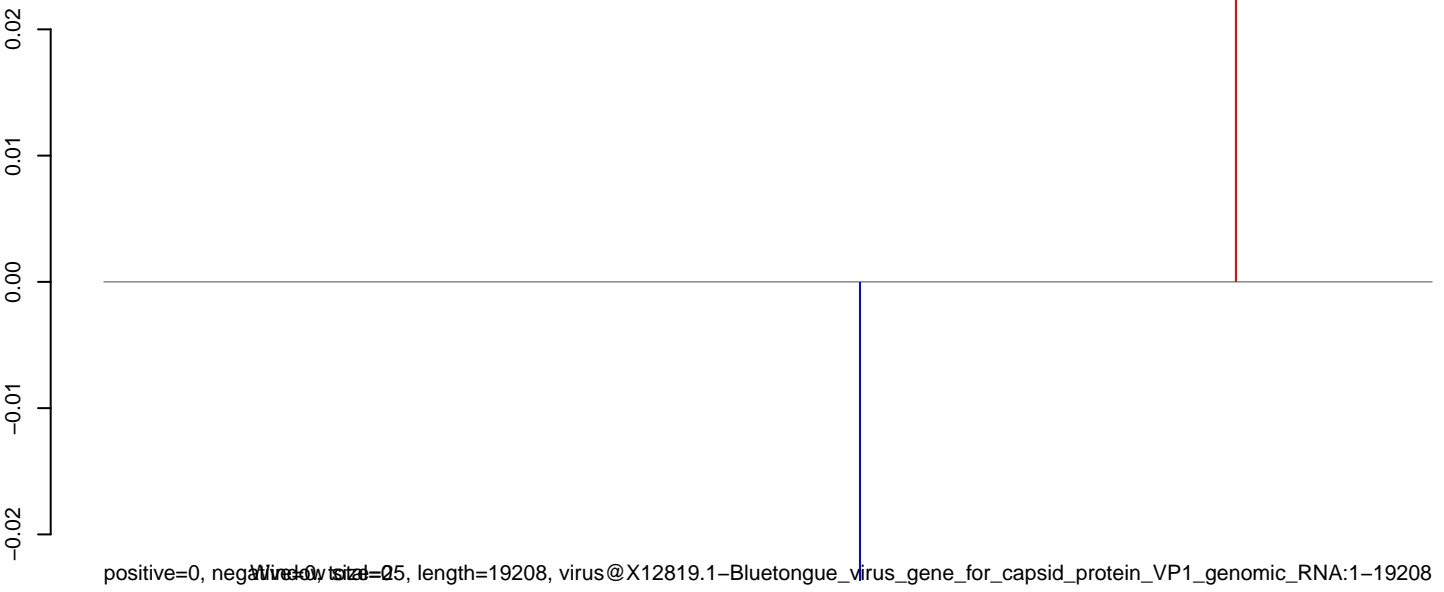
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

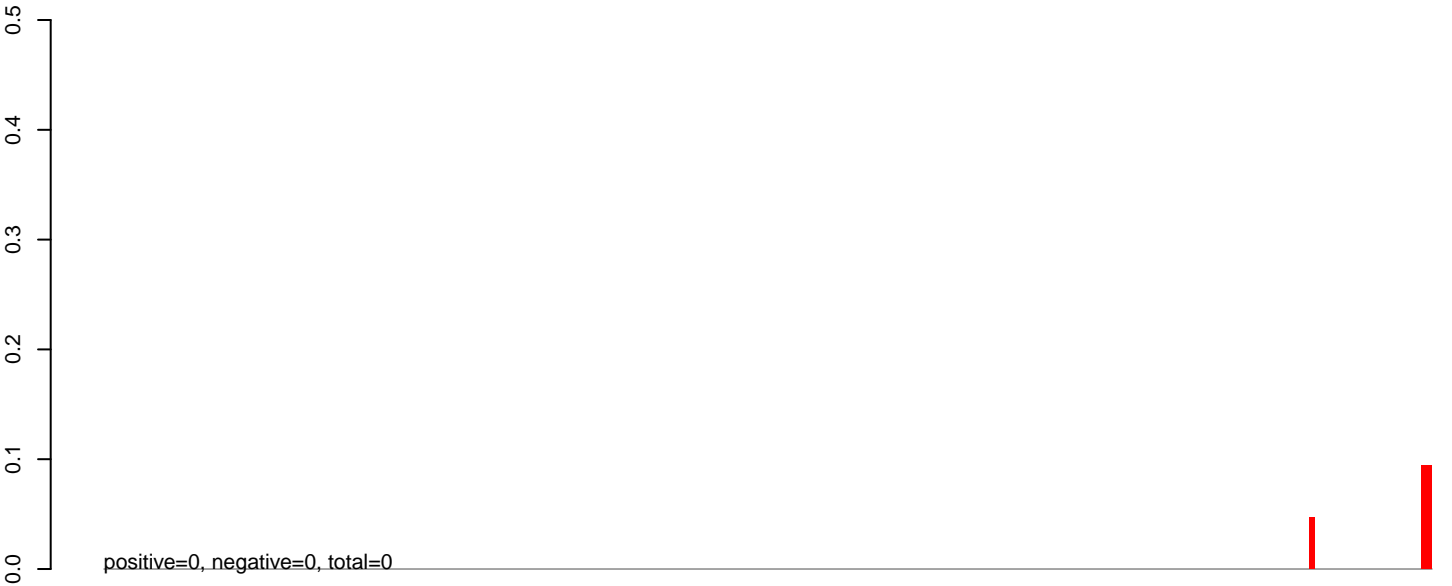


AnGam_Ovary_BetaE.rep

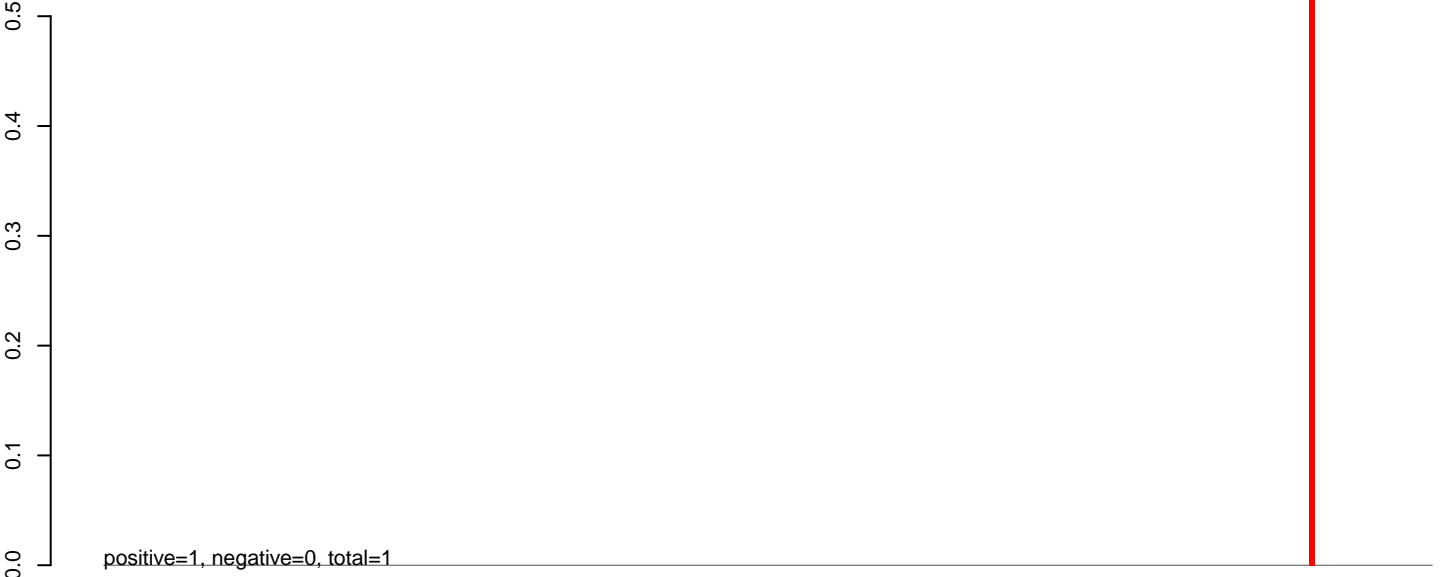


0 5000 10000 15000 20000

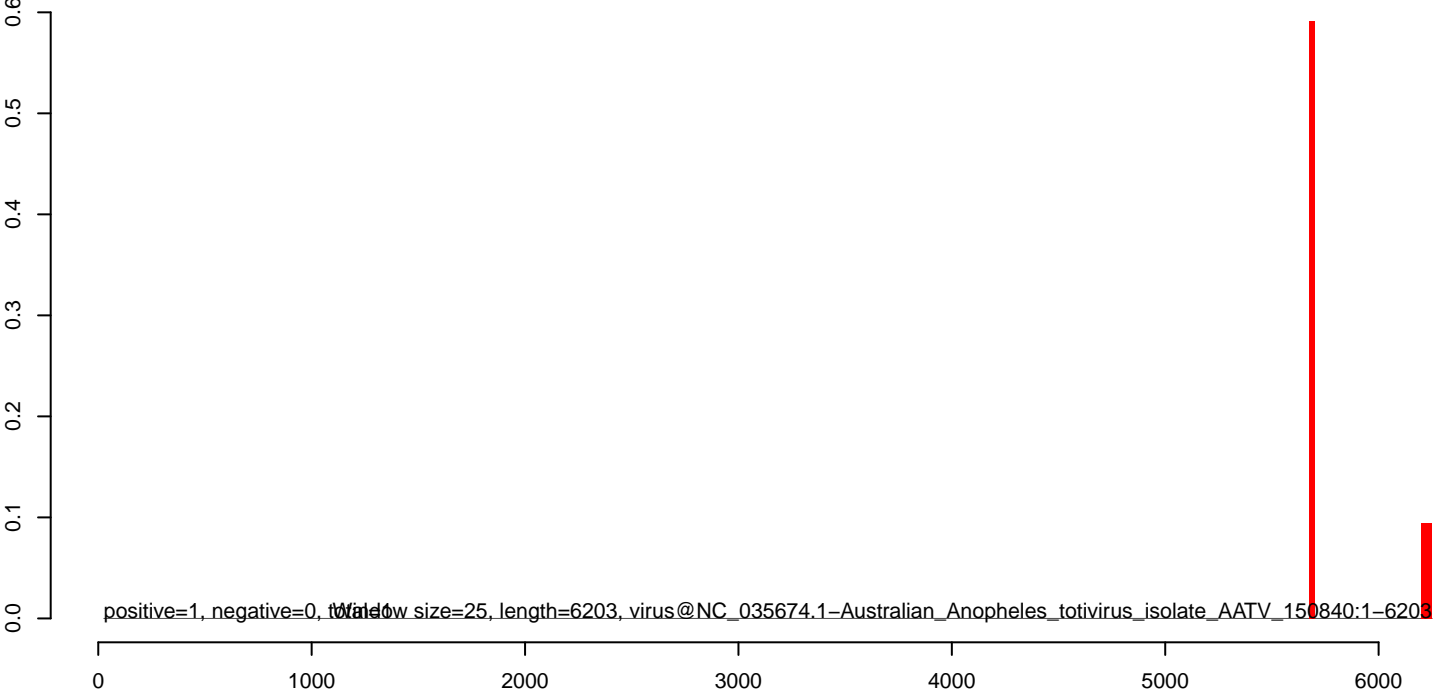
AnGam_Ovary_BetaE.18_23.rep



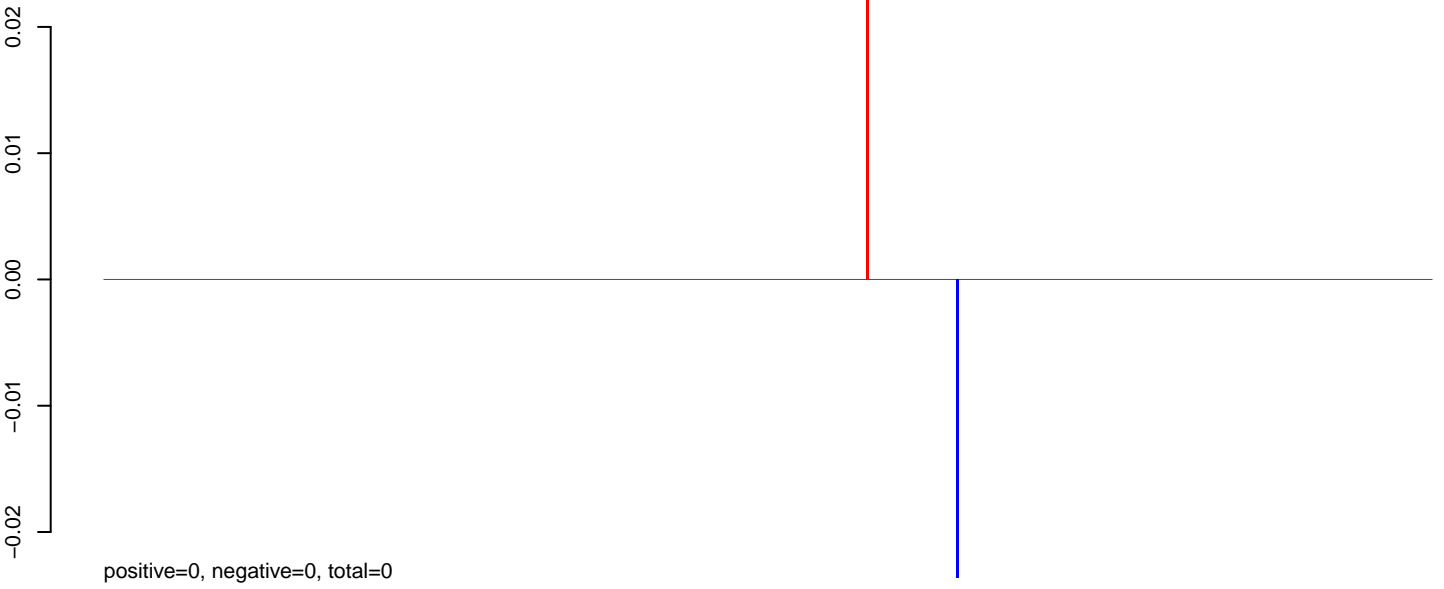
AnGam_Ovary_BetaE.24_35.rep



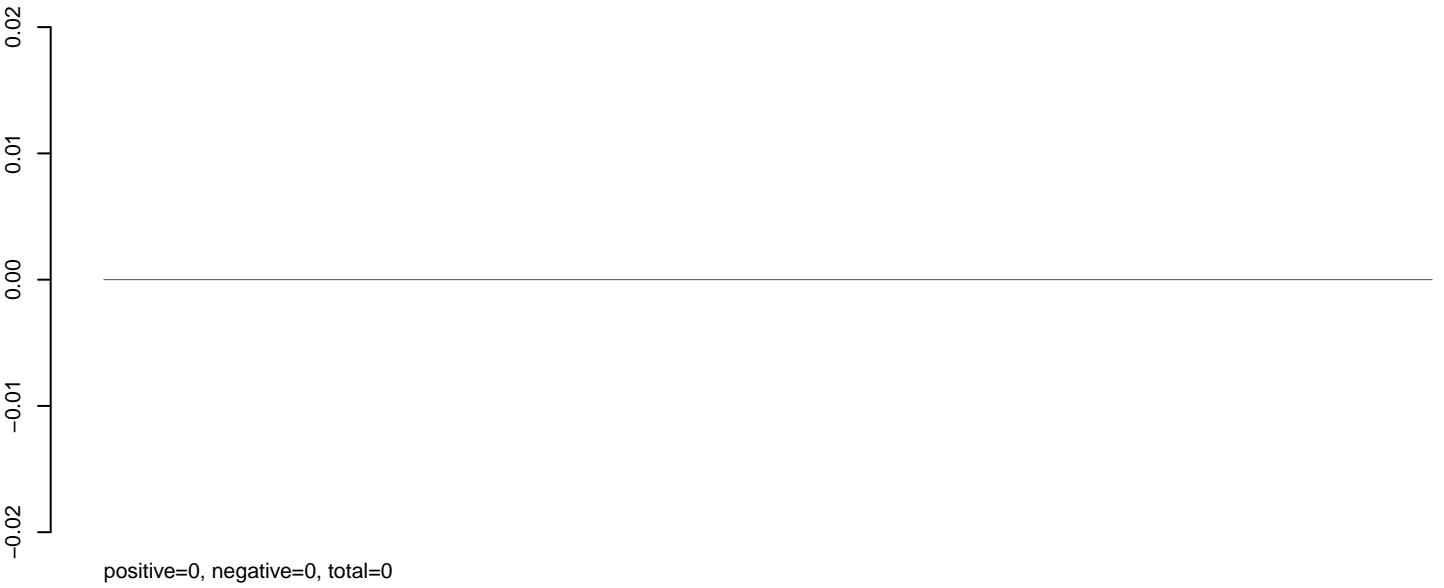
AnGam_Ovary_BetaE.rep



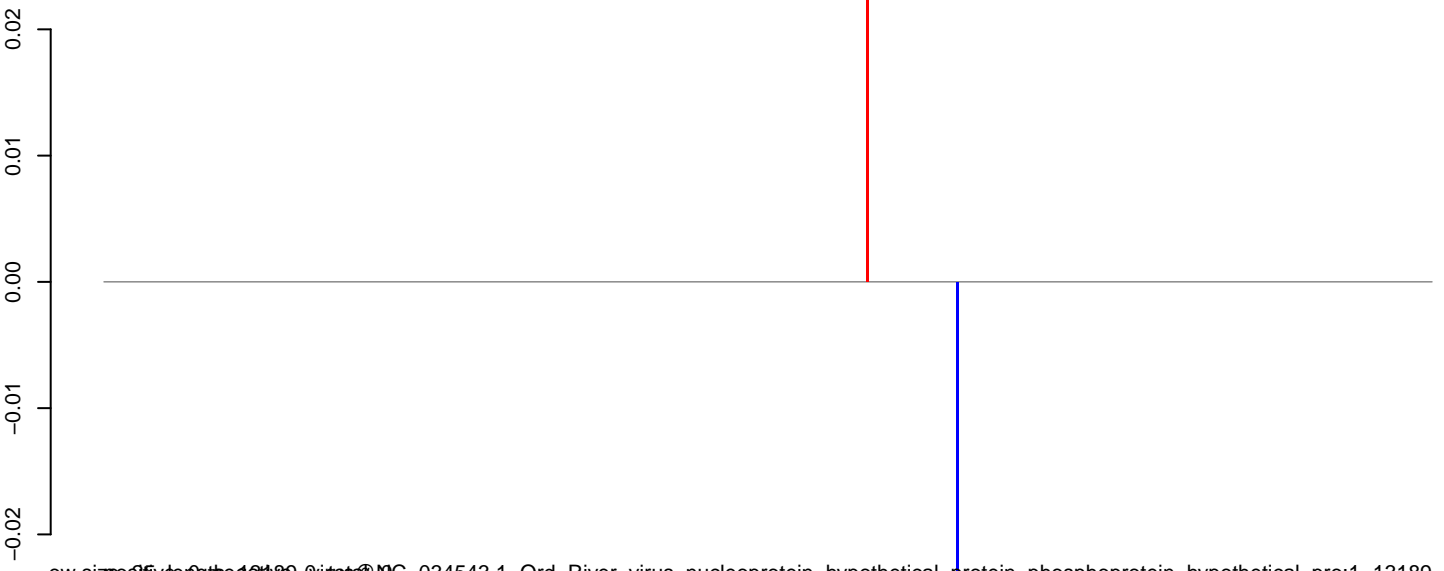
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



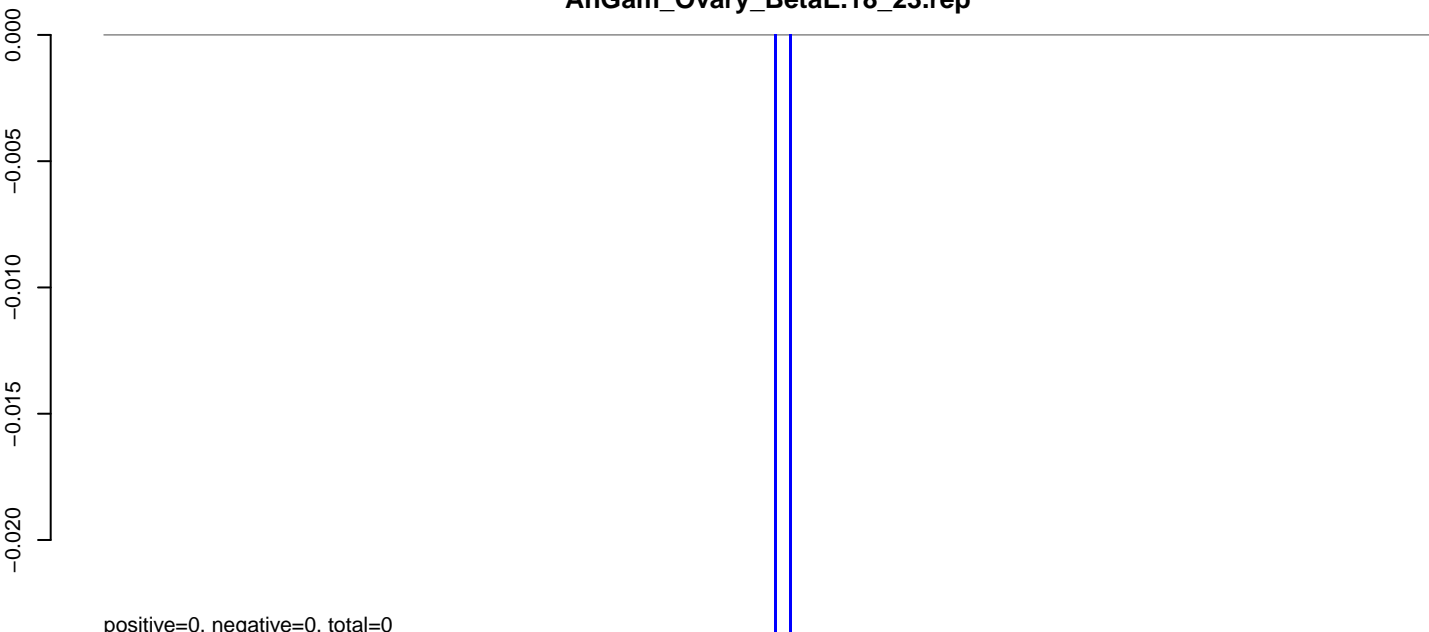
AnGam_Ovary_BetaE.rep



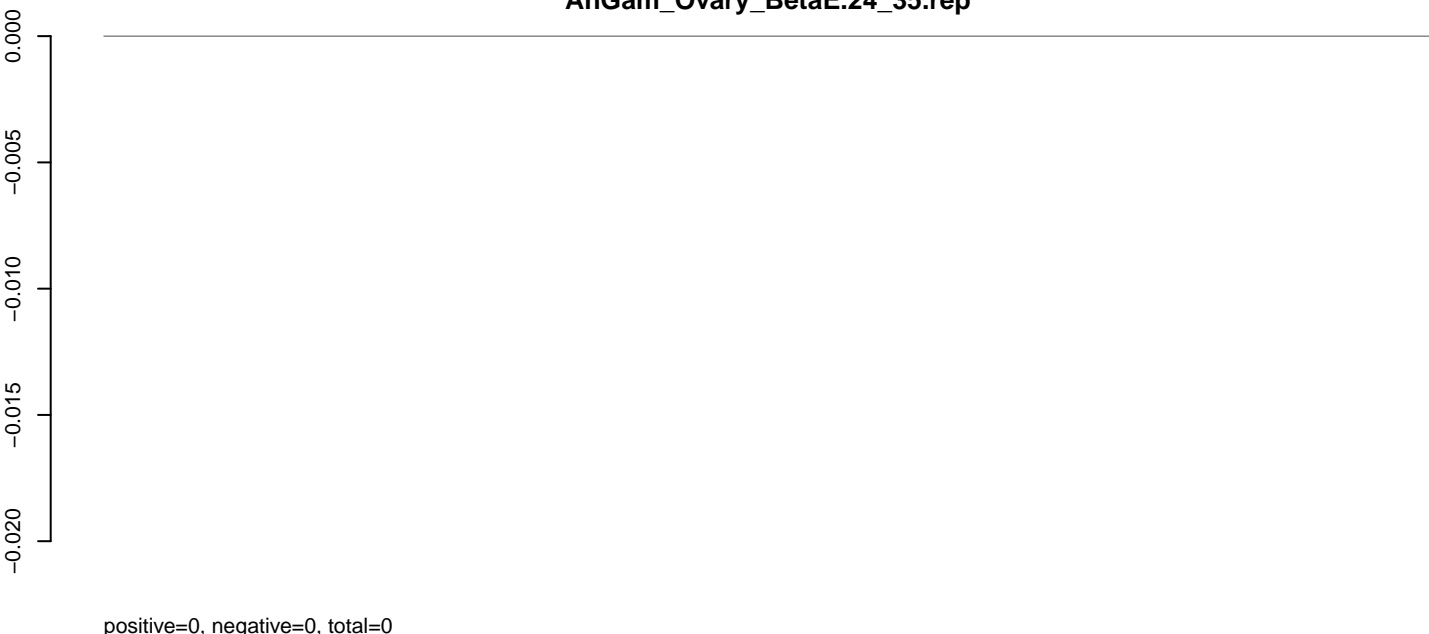
row sizes: 25, 1189, 1189, 0, total=2400 NC_034543.1-Ord_River_virus_nucleoprotein_hypothetical_protein_phosphoprotein_hypothetical_pro:1-13189

0 2000 4000 6000 8000 10000 12000

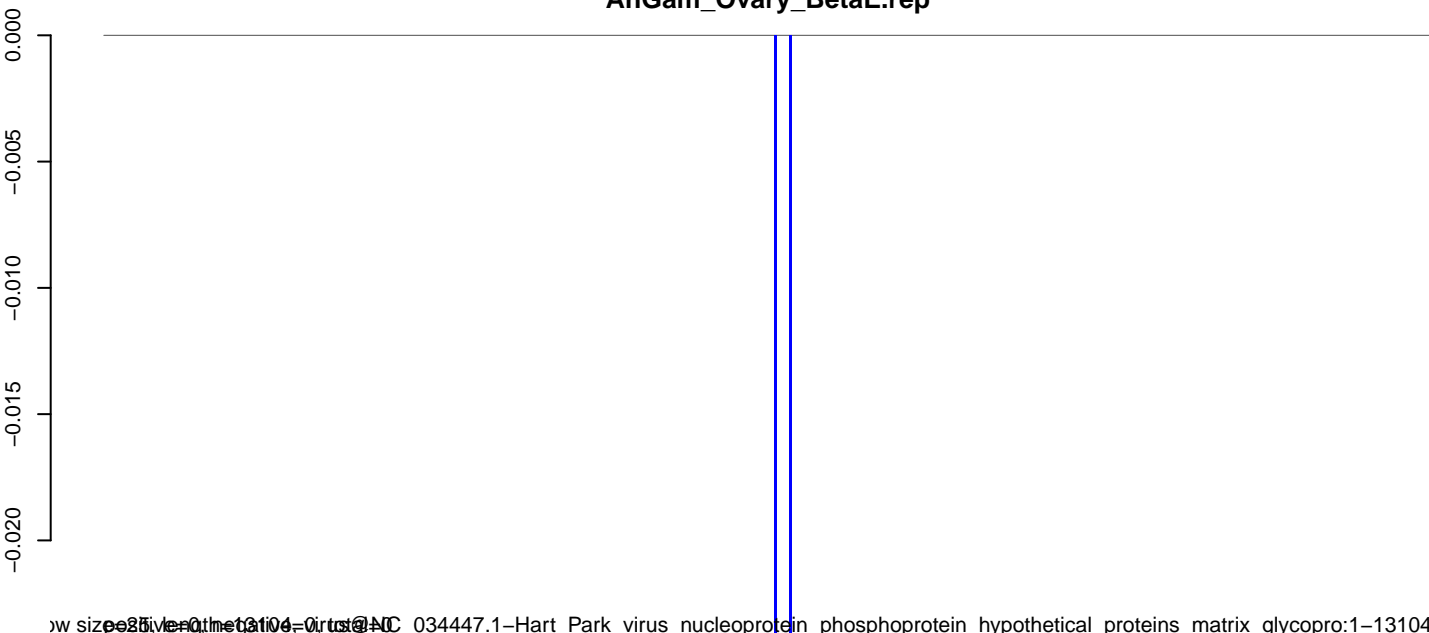
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



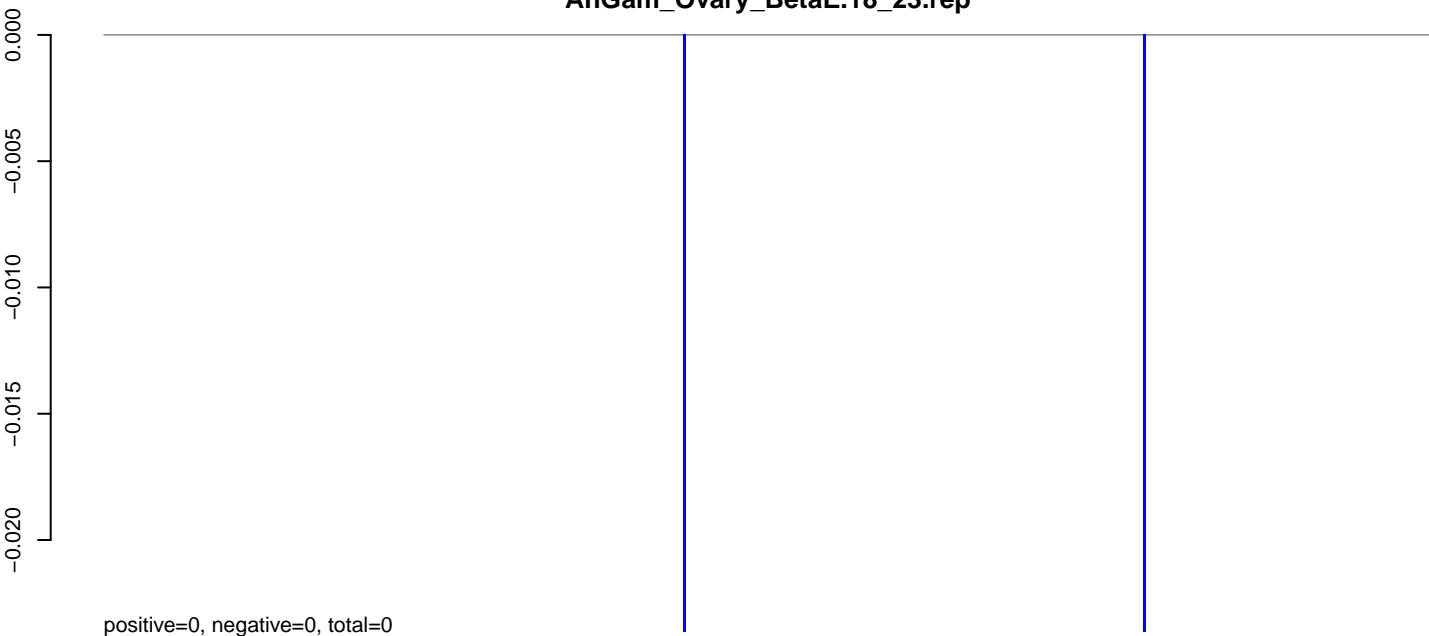
AnGam_Ovary_BetaE.rep



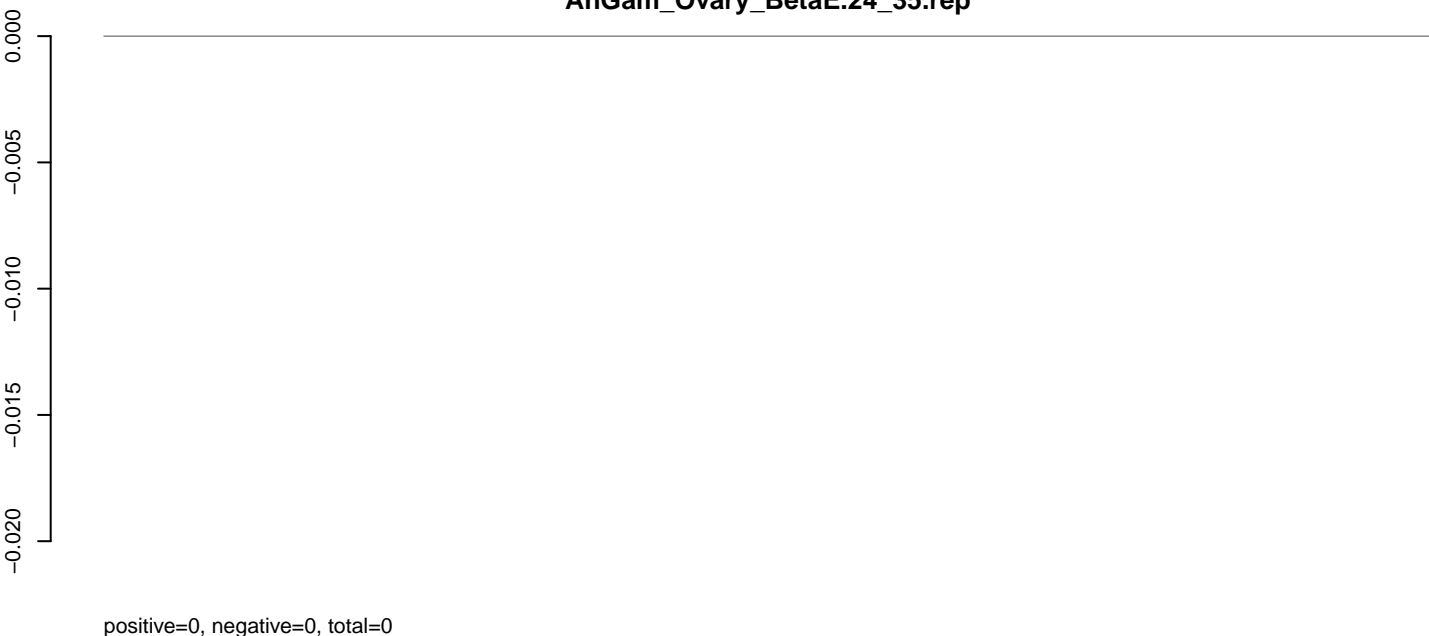
www.ebi.ac.uk/ena/browser/view/PRINCE_034447.1-Hart_Park_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopro:1-13104

0 2000 4000 6000 8000 10000 12000

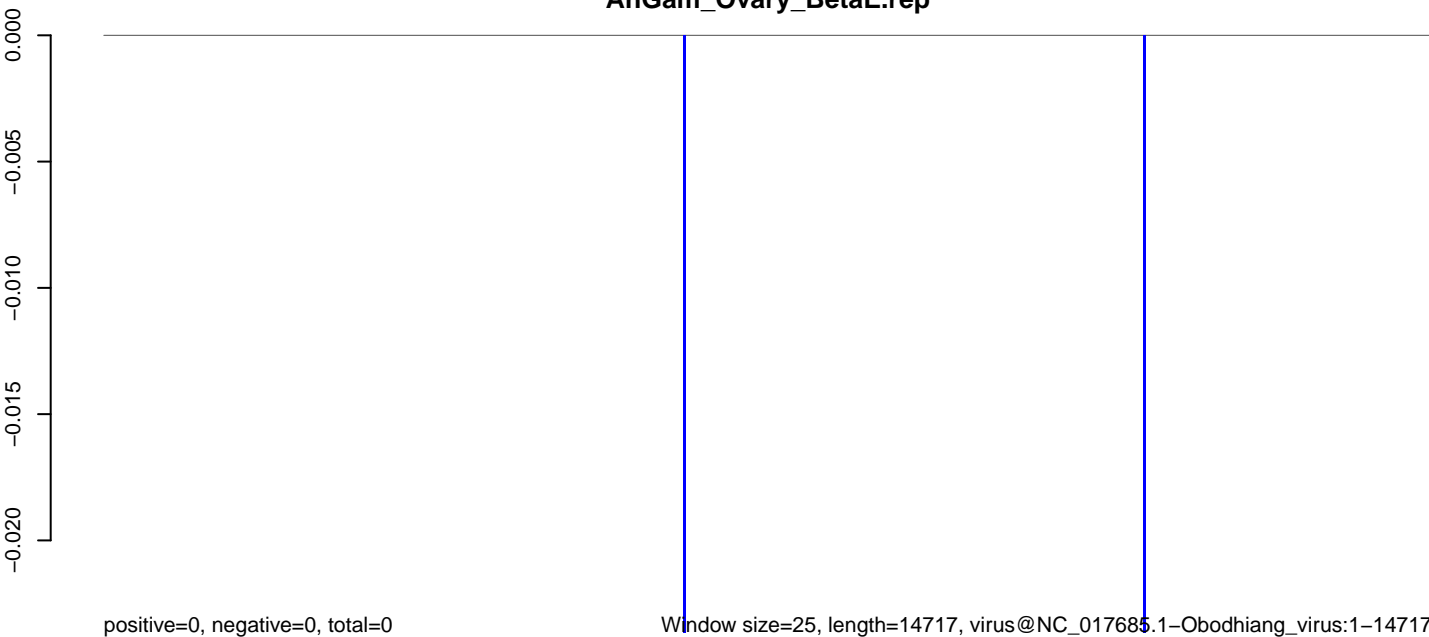
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



Window size=25, length=14717, virus@NC_017685.1-Obodhiang_virus:1-14717

AnGam_Ovary_BetaE.18_23.rep

0.2
0.1
0.0
-0.1

positive=1, negative=0, total=1

AnGam_Ovary_BetaE.24_35.rep

0.2
0.1
0.0
-0.1

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep

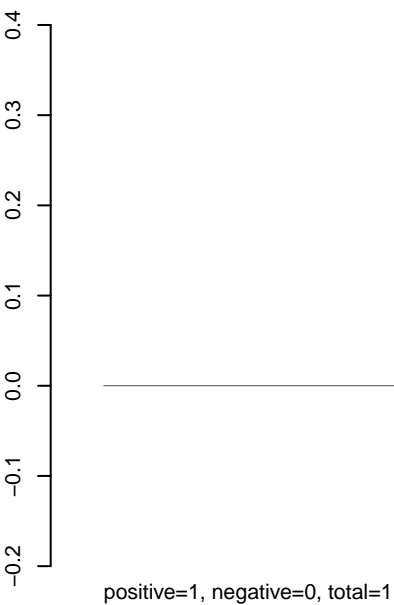
0.3
0.2
0.1
0.0
-0.1

positive=1, negative=0, total=1

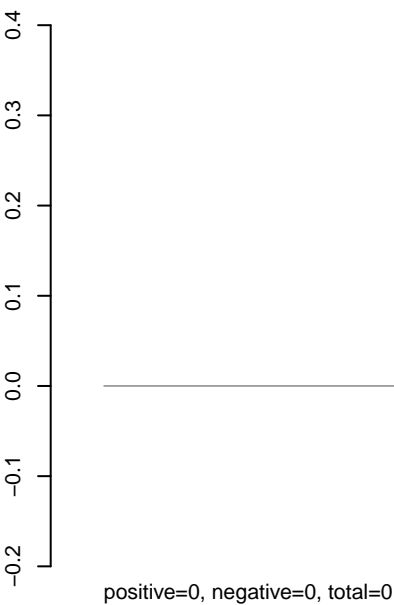
Window size=25, length=11877, virus@NC_016962.1-Bebaru_virus:1-11877

0 2000 4000 6000 8000 10000 12000

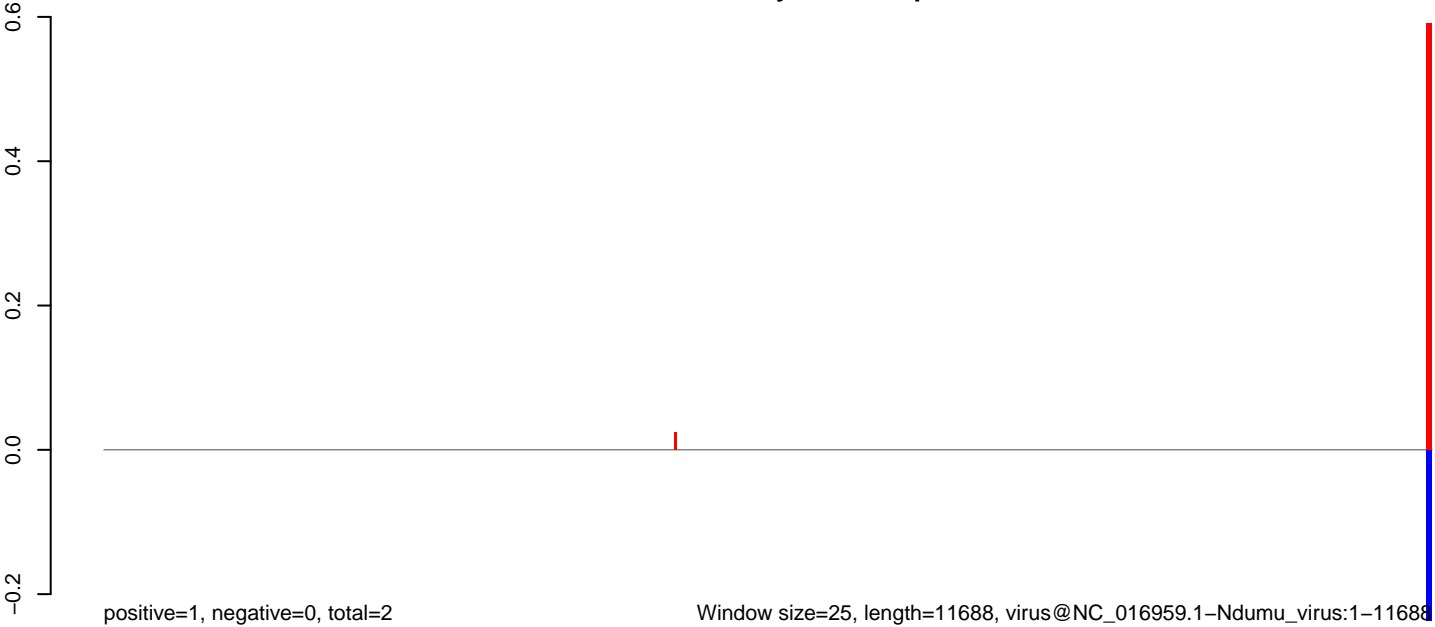
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

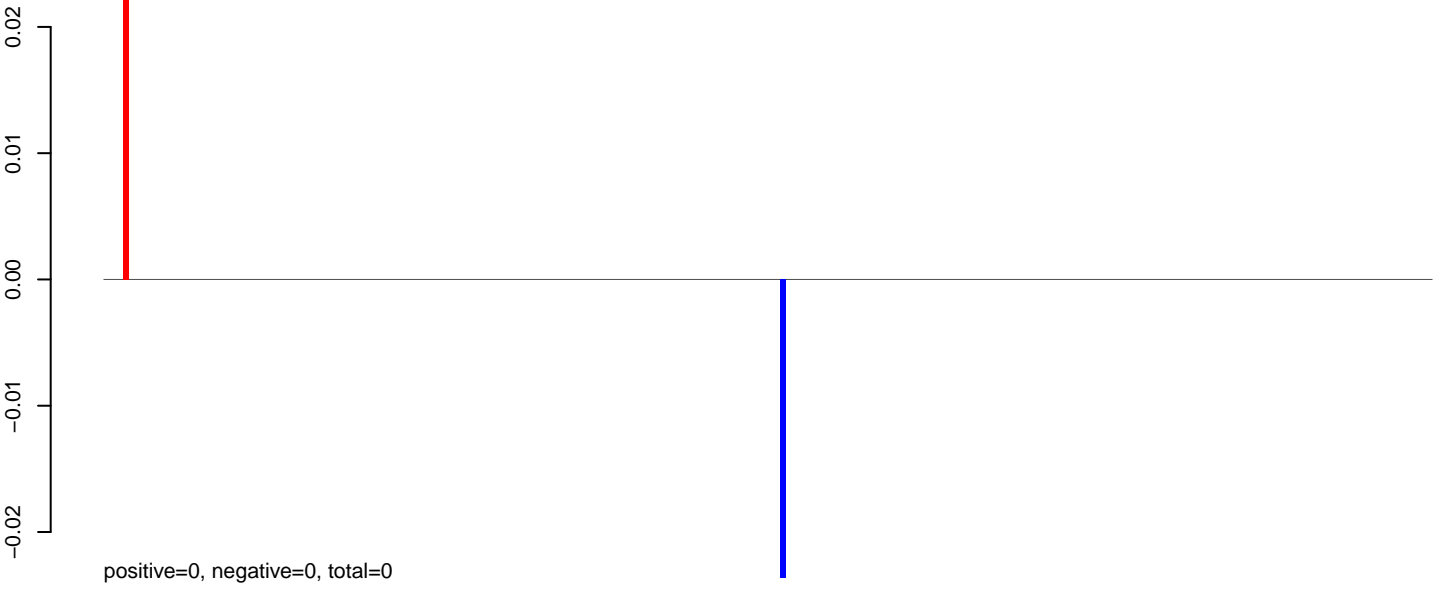


AnGam_Ovary_BetaE.rep

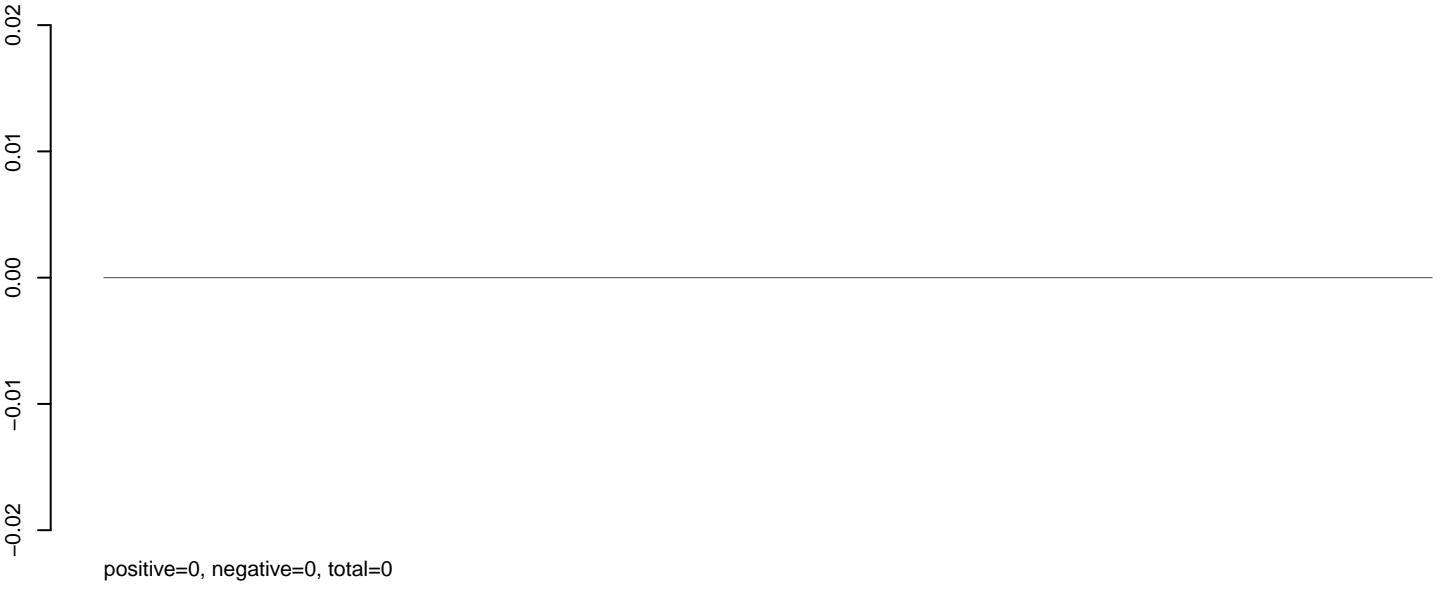


0 2000 4000 6000 8000 10000 12000

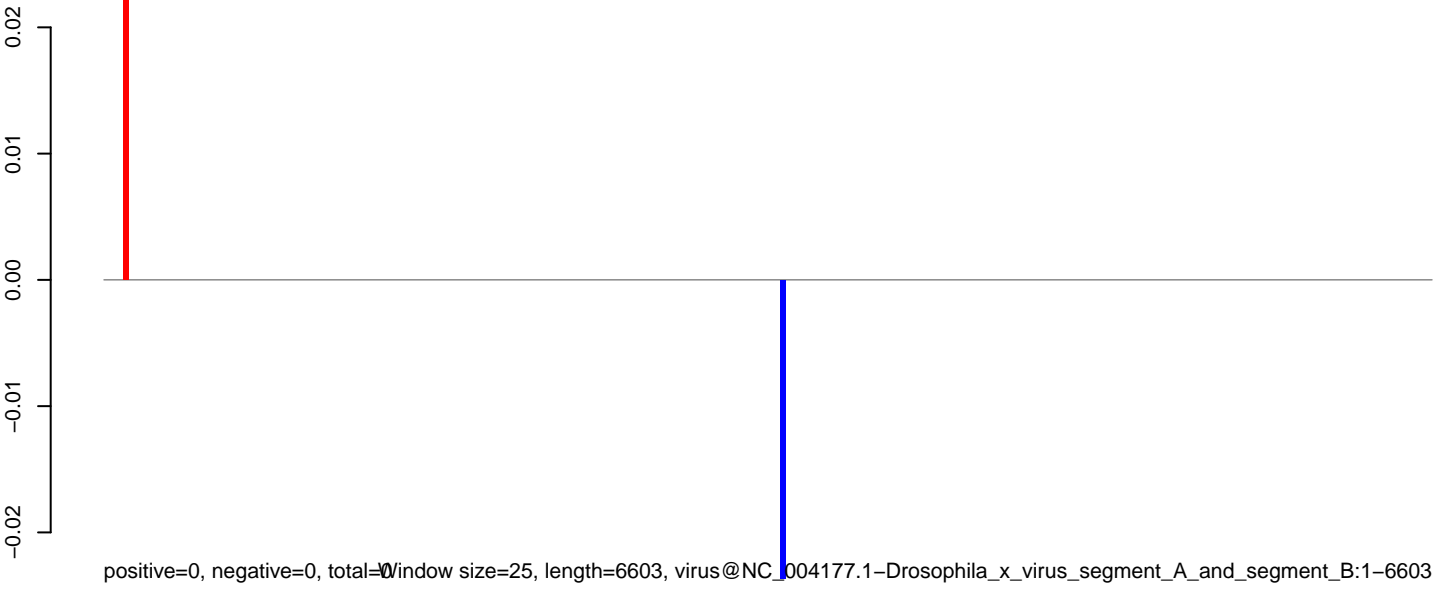
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



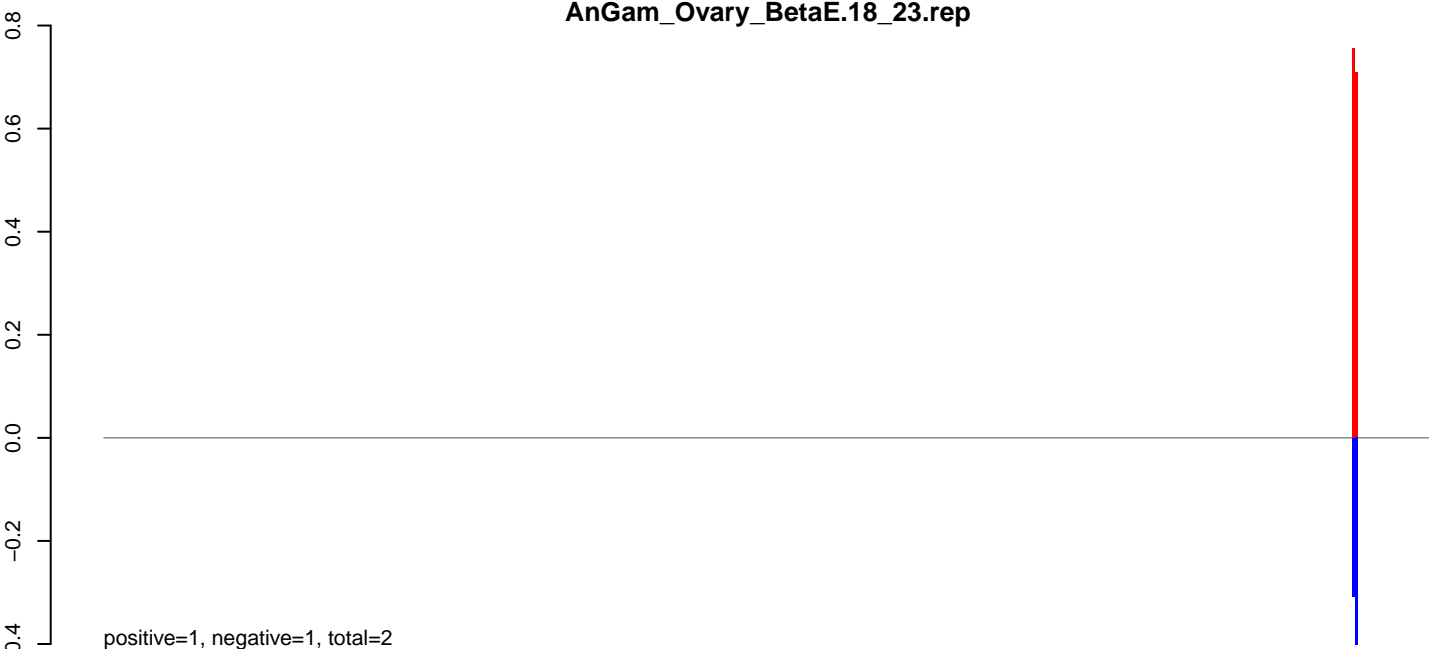
AnGam_Ovary_BetaE.rep



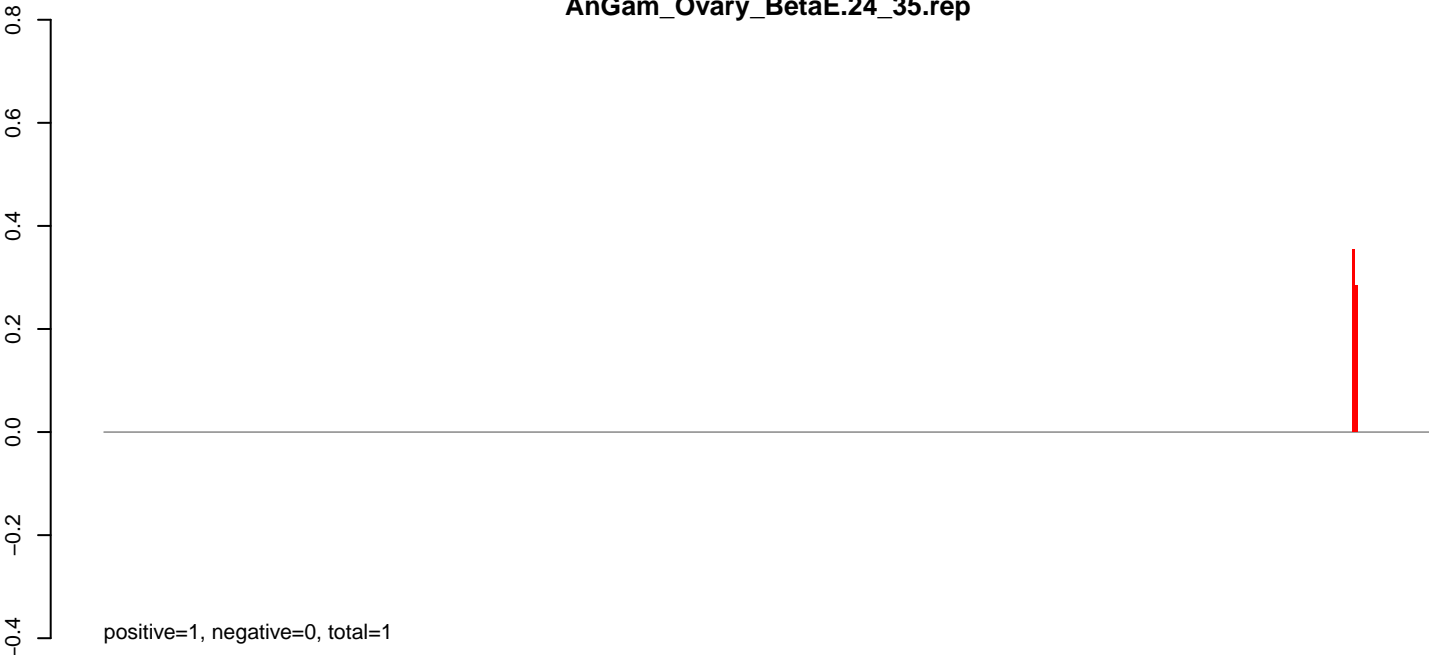
Window size=25, length=6603, virus@NC_004177.1-Drosophila_x_virus_segment_A_and_segment_B:1-6603

0 1000 2000 3000 4000 5000 6000

AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



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

0 2000 4000 6000 8000 10000

AnGam_Ovary_BetaE.18_23.rep

0.00
-0.01
-0.02
-0.03
-0.04

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep

0.00
-0.01
-0.02
-0.03
-0.04

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep

0.00
-0.01
-0.02
-0.03
-0.04

Window size=25, n=0, total=3937, virus@MH550148.1-Aedes_albopictus_densovirus_isolate_NC01_L3_1_trimmed_contig_5889-RC:1-3937

0 1000 2000 3000 4000

AnGam_Ovary_BetaE.18_23.rep

0.2
0.1
0.0
-0.1

positive=1, negative=0, total=1

AnGam_Ovary_BetaE.24_35.rep

0.2
0.1
0.0
-0.1

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep

0.3
0.2
0.1
0.0
-0.1

downloaded from SRA on 2025-10-10 12:05:20, via @HM147991.1-Trocaravirus_non_structural_polyprotein_gene_partial_cds_and_structural_polyprotei:1-12052

0 2000 4000 6000 8000 10000 12000

AnGam_Ovary_BetaE.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

Window size=25, length=3893, virus@GU452720.1-Mosquito_densovirus_BR_07:1-3893

0 1000 2000 3000 4000

AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep

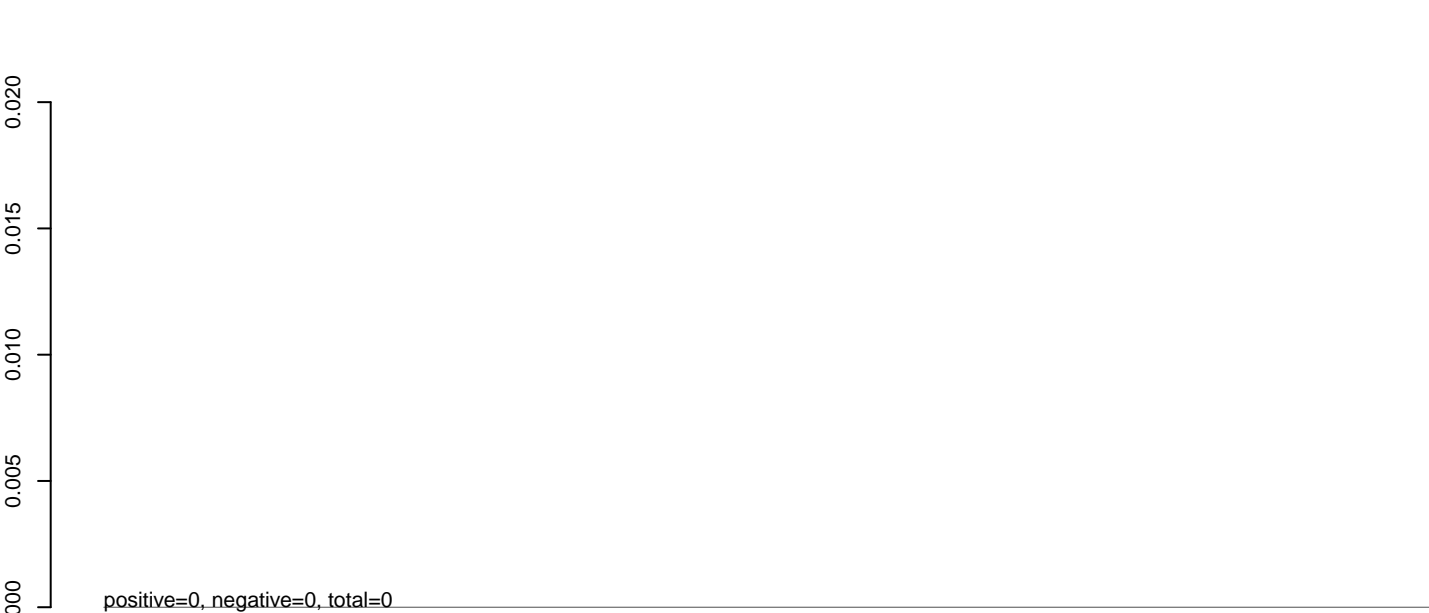


0 200 400 600 800 1000

AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



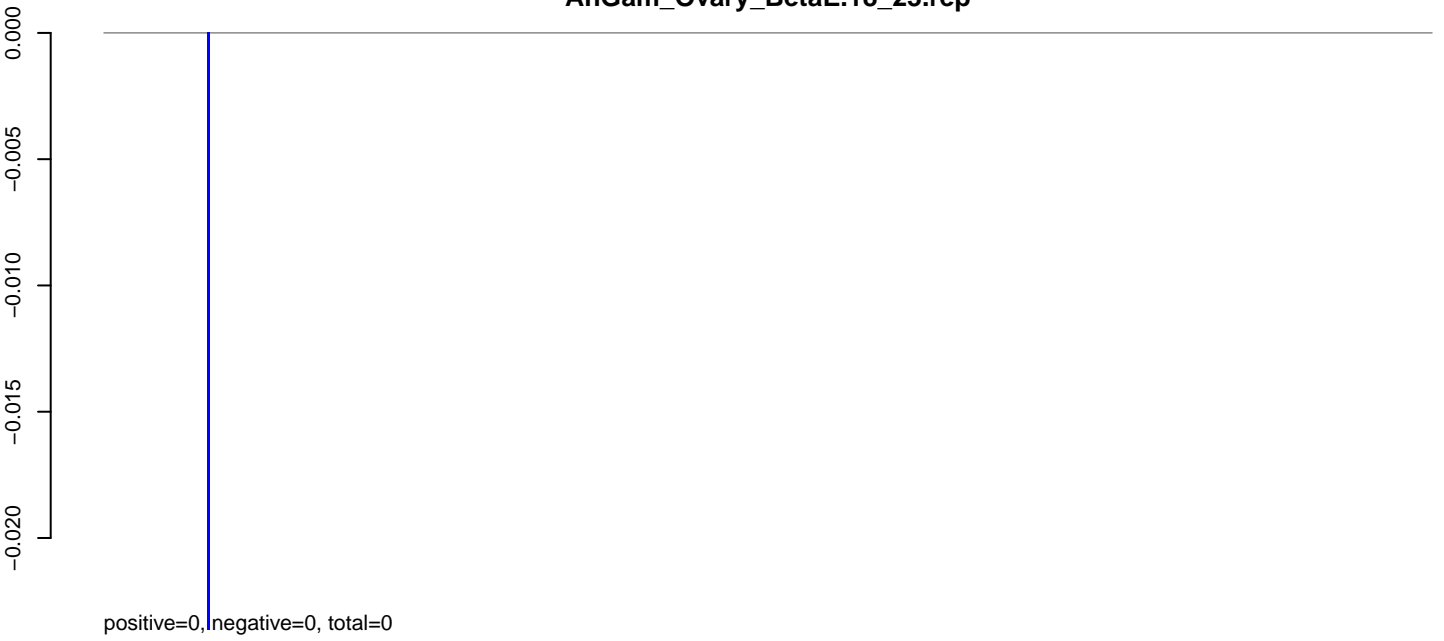
AnGam_Ovary_BetaE.rep



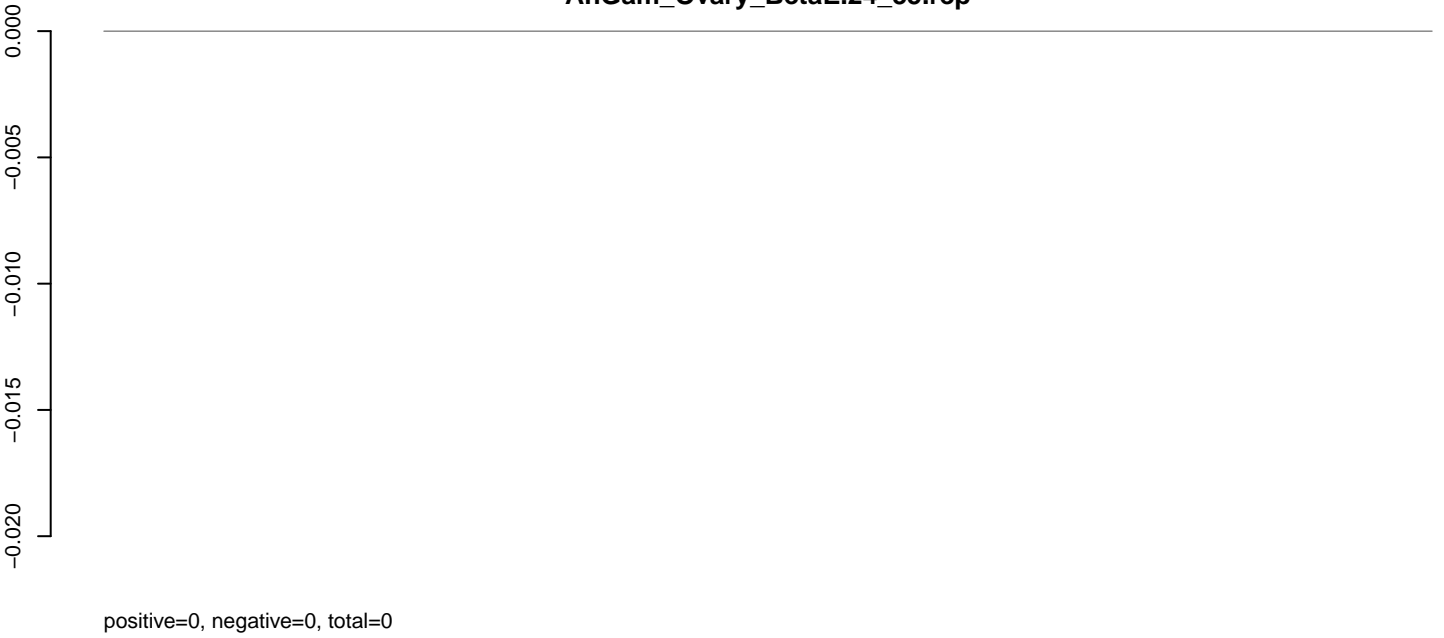
0 2000 4000 6000 8000 10000

11835.2-Aedes_albopictus_cell_fusing_agent_virus_polyprotein-like_protein_gene_and_agCP8252:1-10587

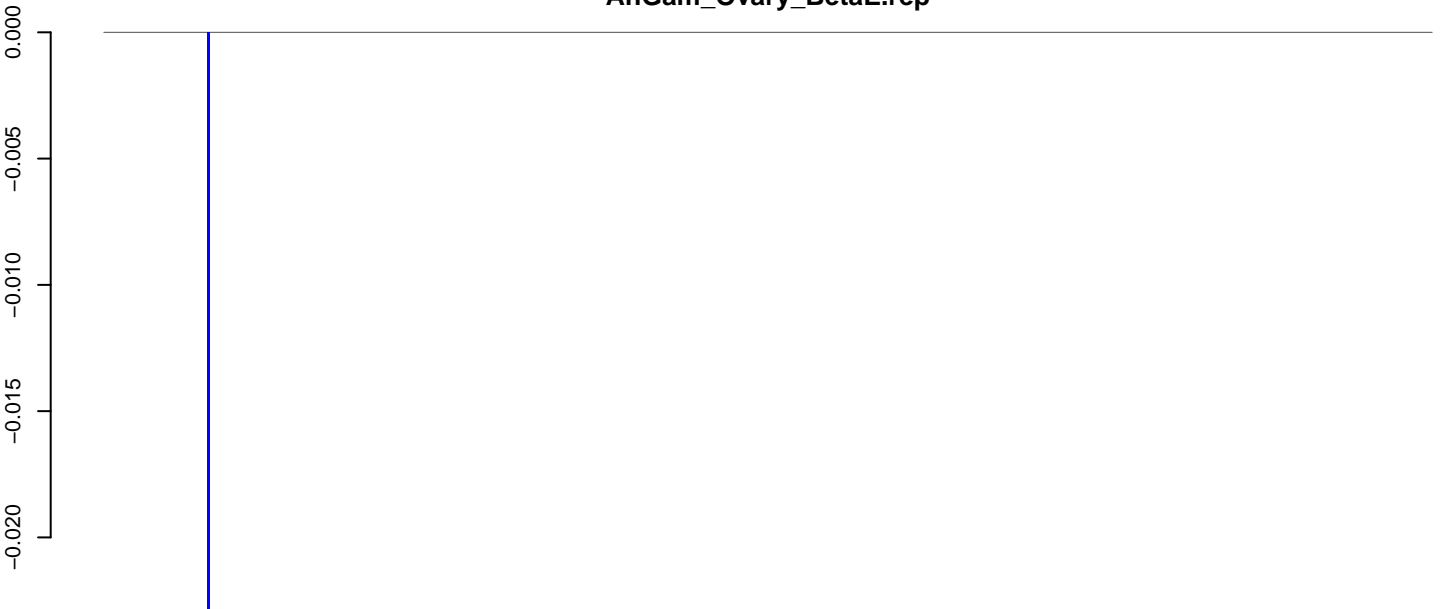
AnGam_Ovary_BetaE.18_23.rep



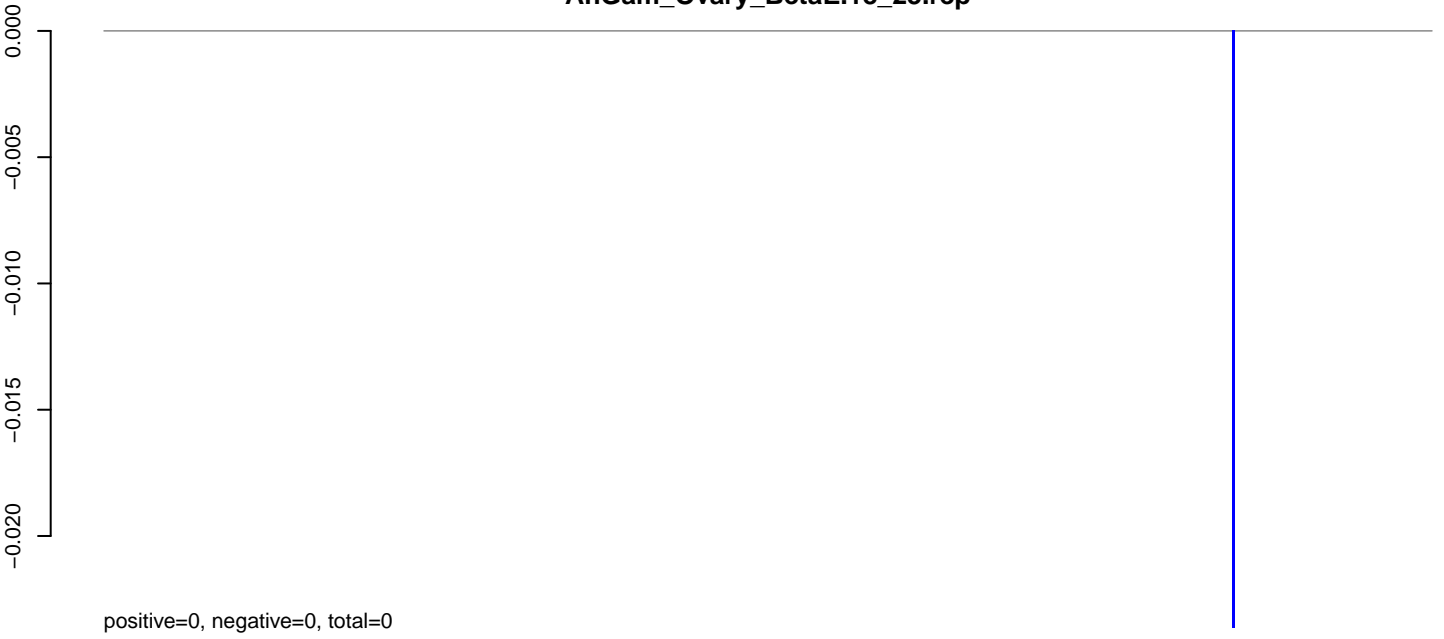
AnGam_Ovary_BetaE.24_35.rep



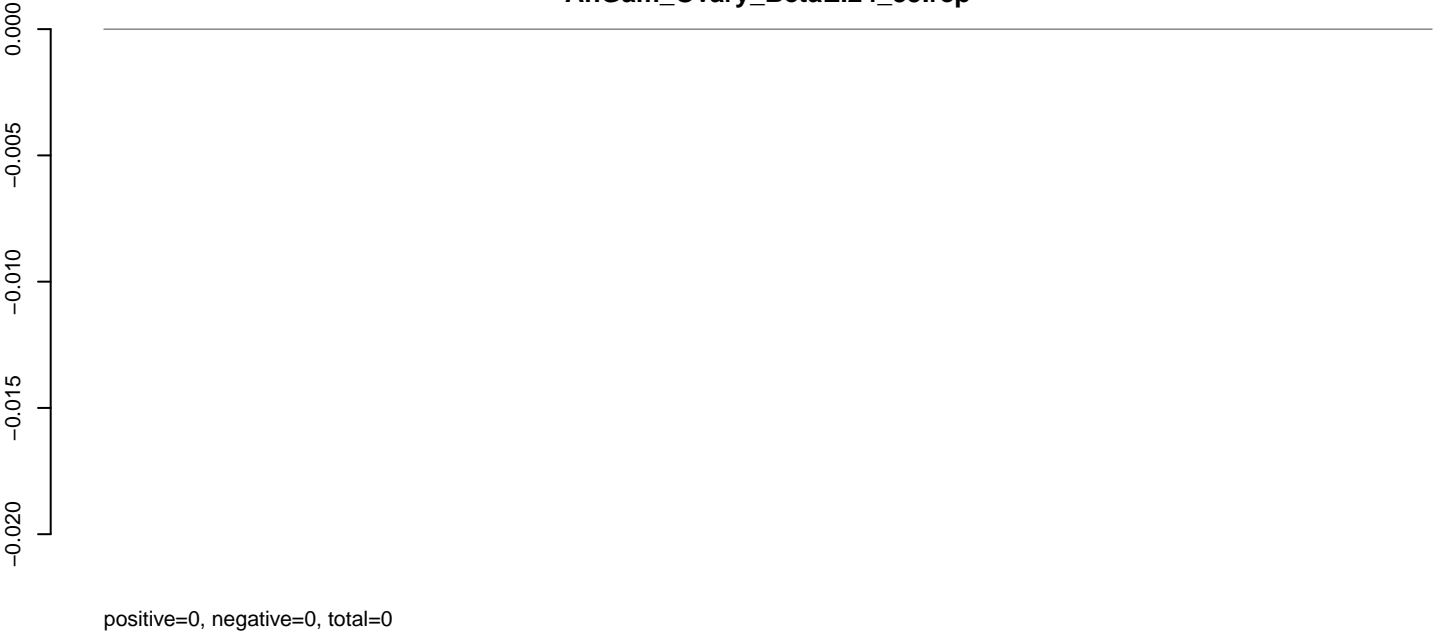
AnGam_Ovary_BetaE.rep



AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

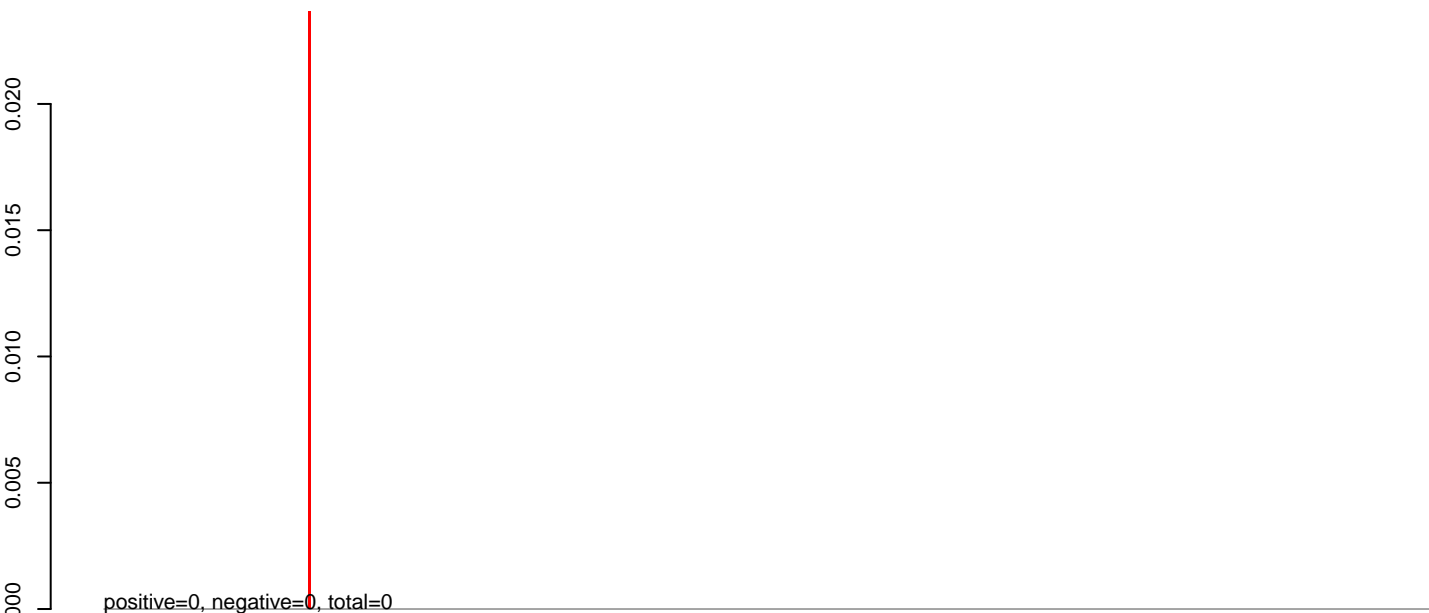


AnGam_Ovary_BetaE.rep



0 2000 4000 6000 8000 10000 12000 14000

AnGam_Ovary_BetaE.18_23.rep



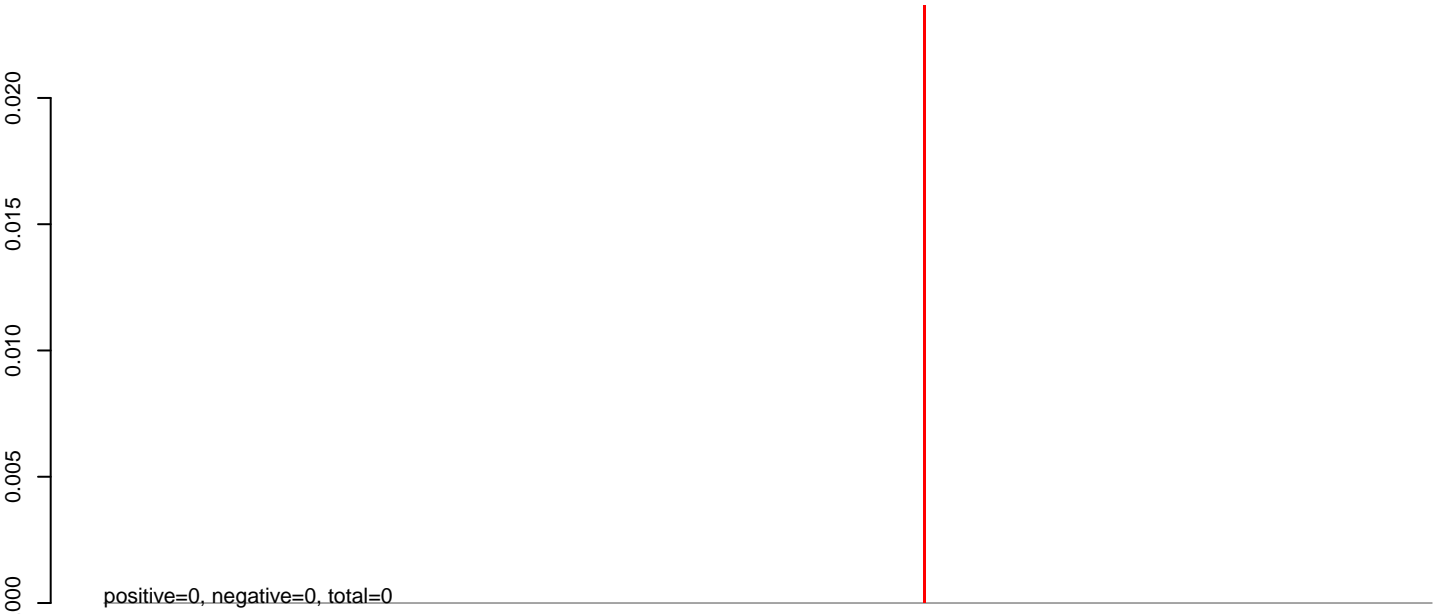
AnGam_Ovary_BetaE.24_35.rep



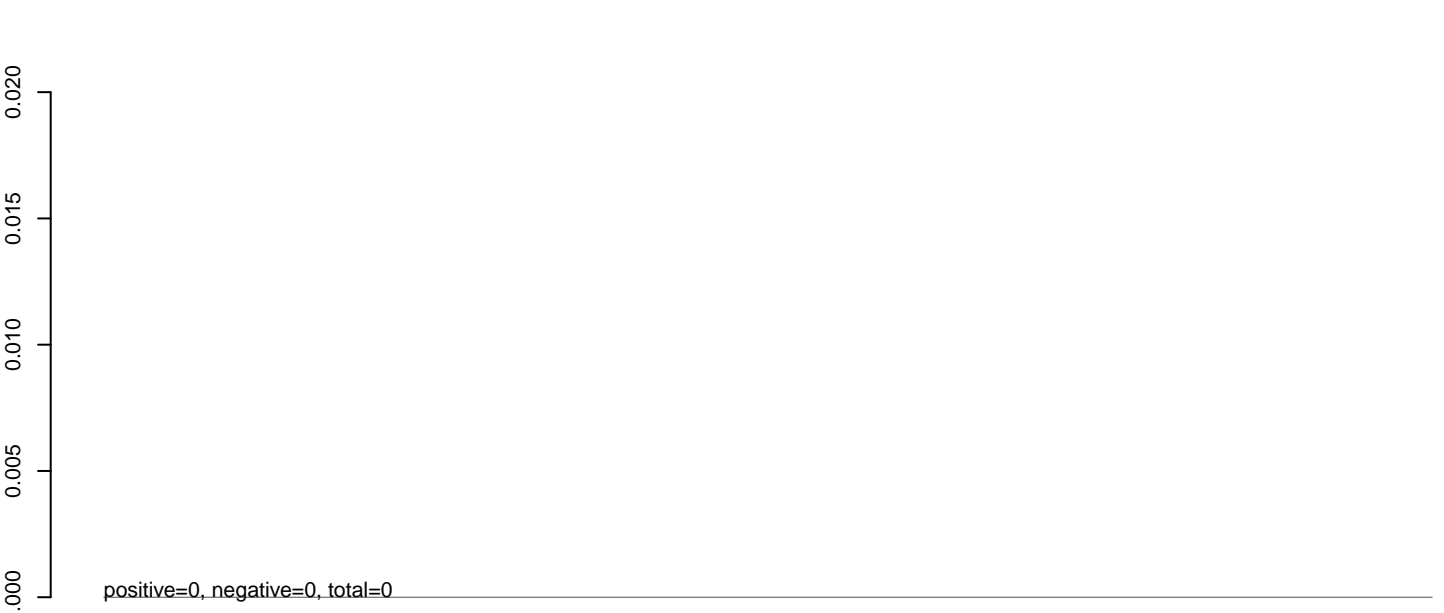
AnGam_Ovary_BetaE.rep



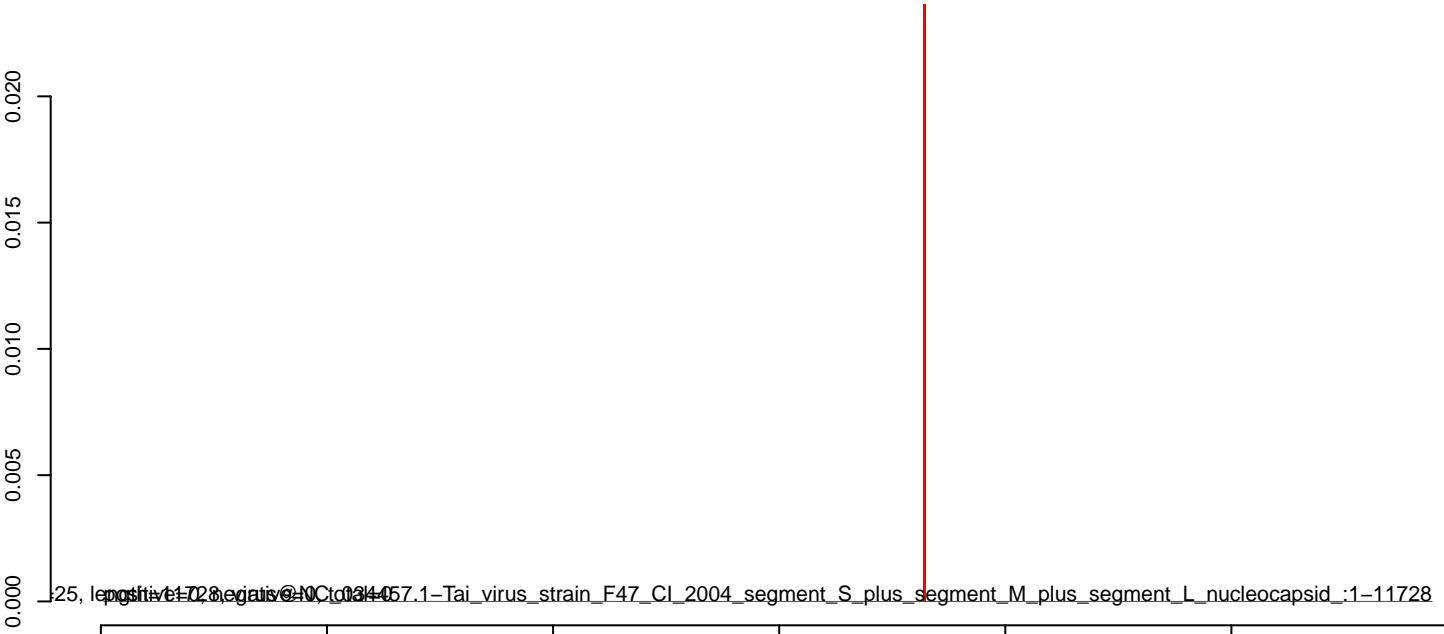
AnGam_Ovary_BetaE.18_23.rep



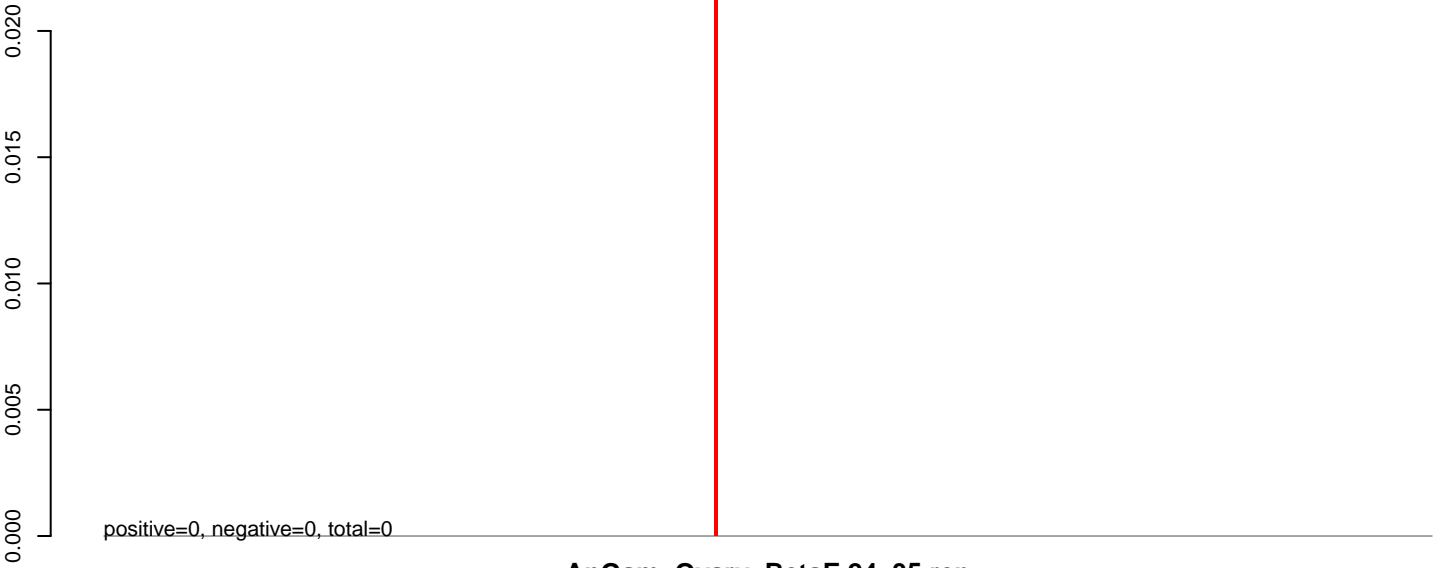
AnGam_Ovary_BetaE.24_35.rep



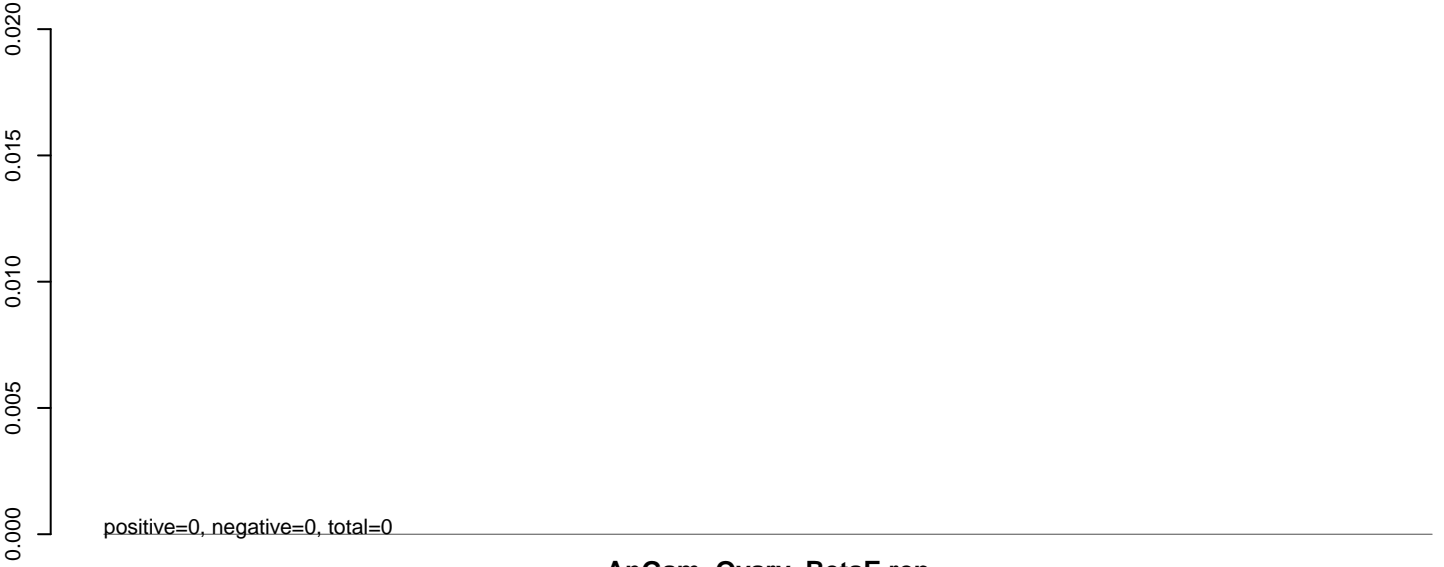
AnGam_Ovary_BetaE.rep



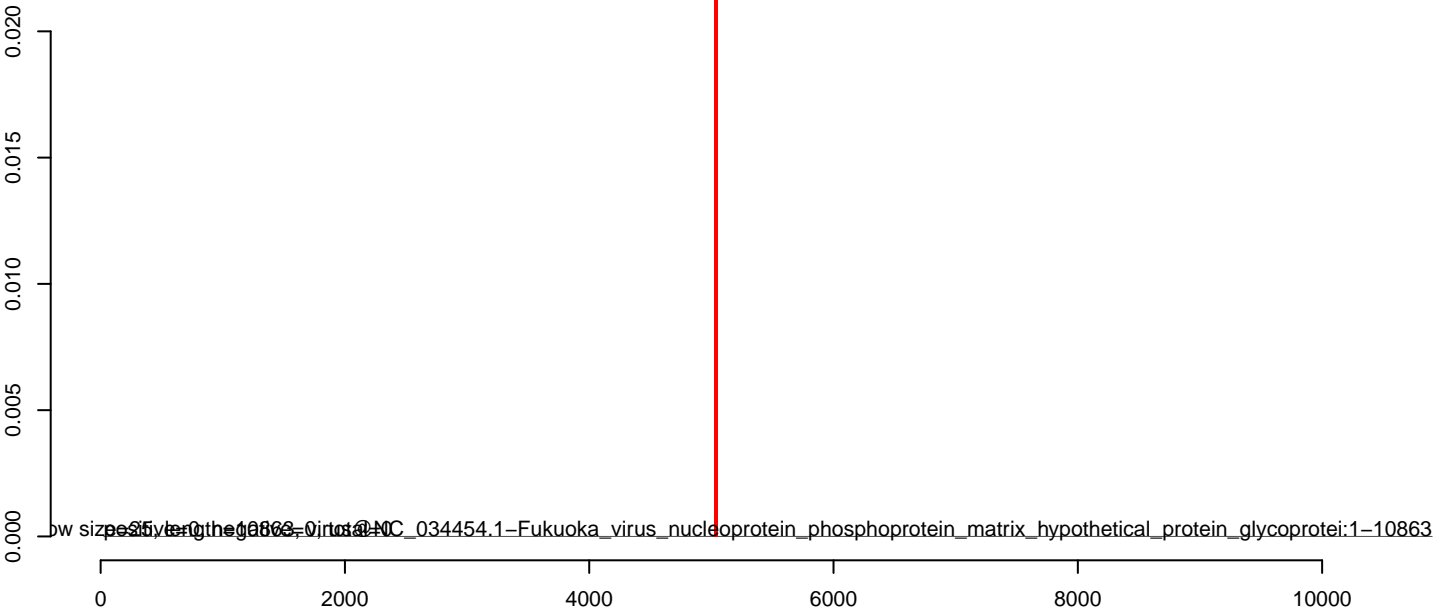
AnGam_Ovary_BetaE.18_23.rep



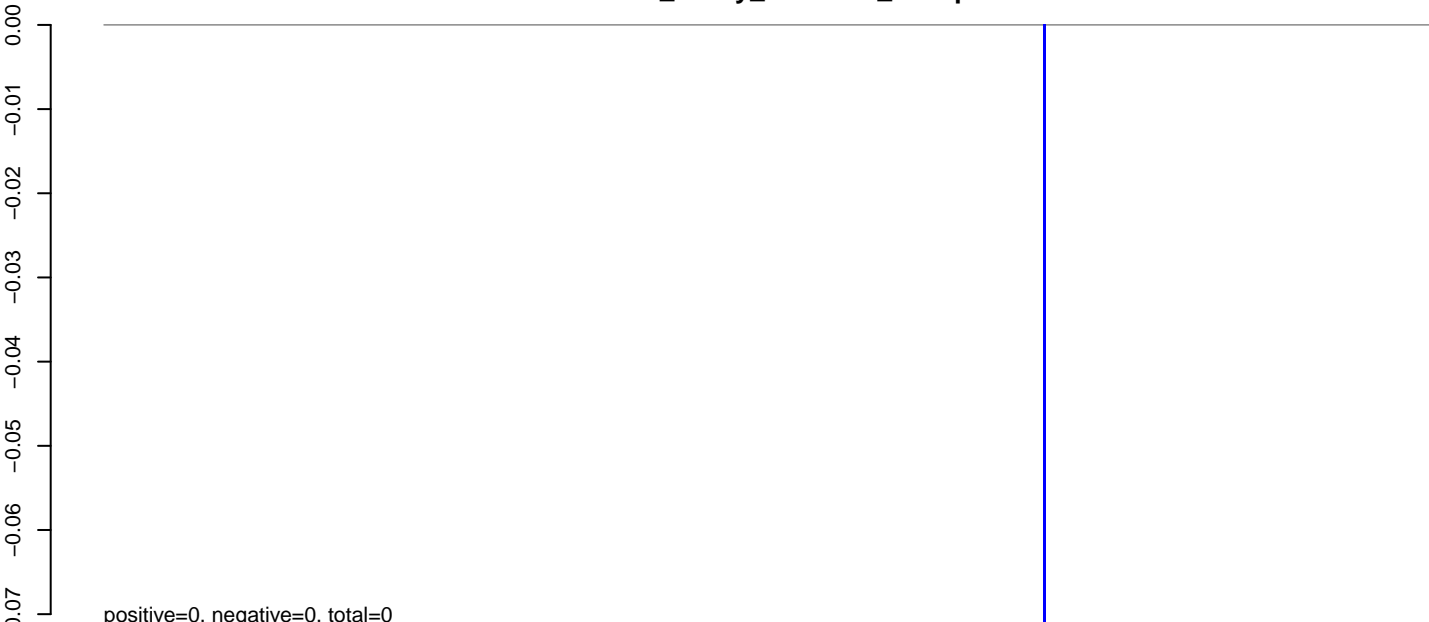
AnGam_Ovary_BetaE.24_35.rep



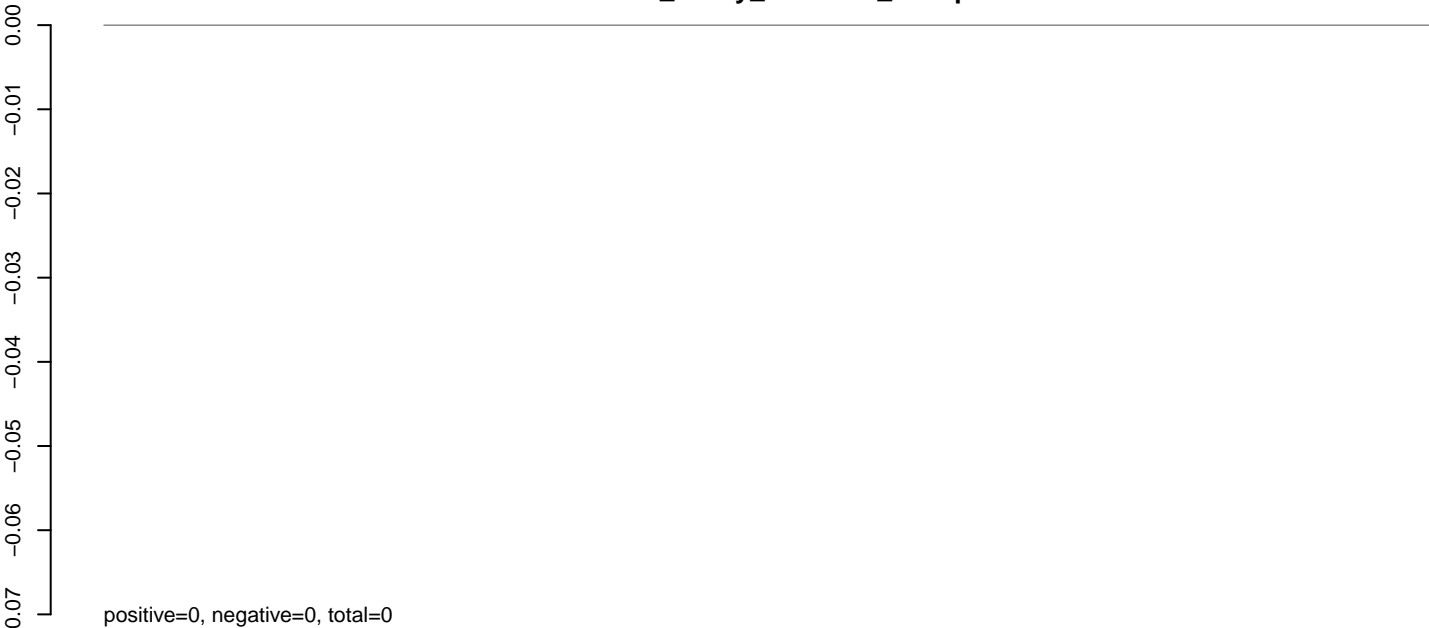
AnGam_Ovary_BetaE.rep



AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

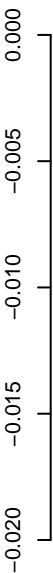


AnGam_Ovary_BetaE.rep



0 2000 4000 6000 8000 10000

AnGam_Ovary_BetaE.18_23.rep



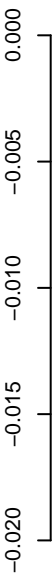
positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep

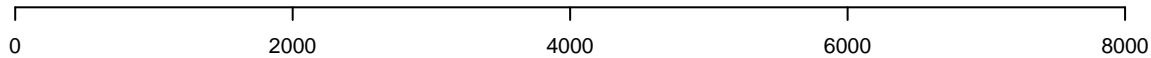


positive=0, negative=0, total=0

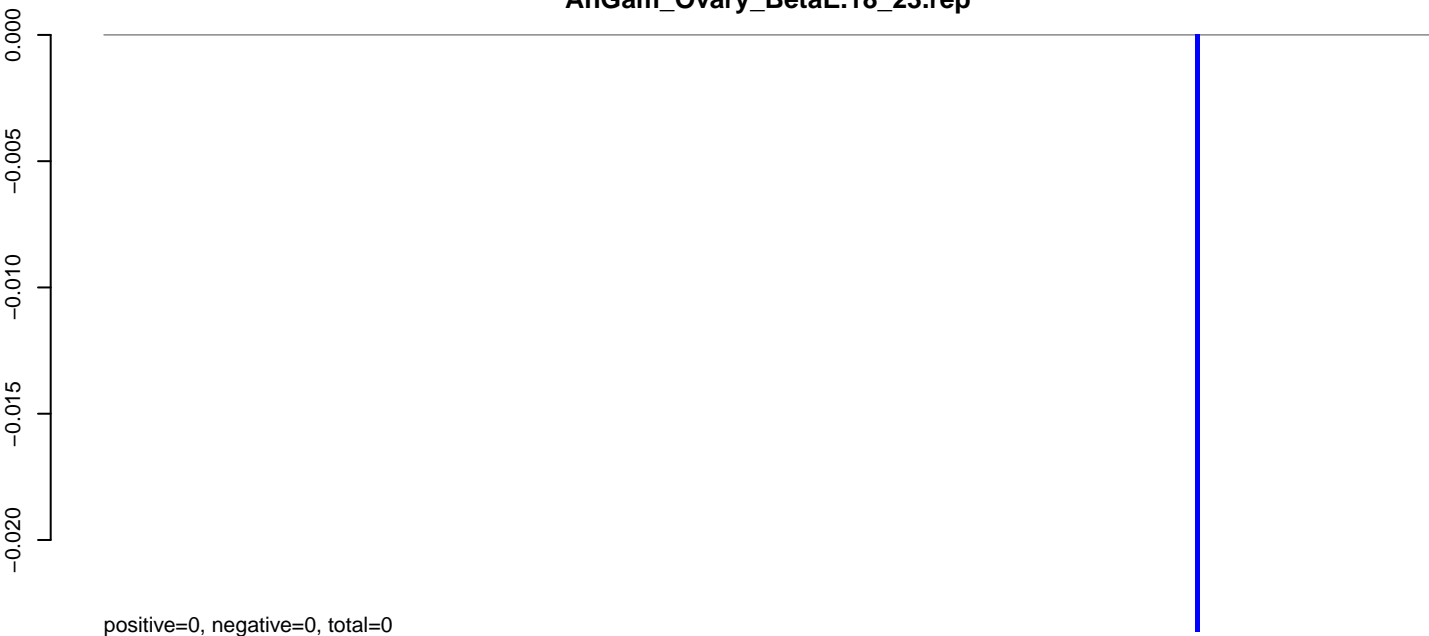
AnGam_Ovary_BetaE.rep



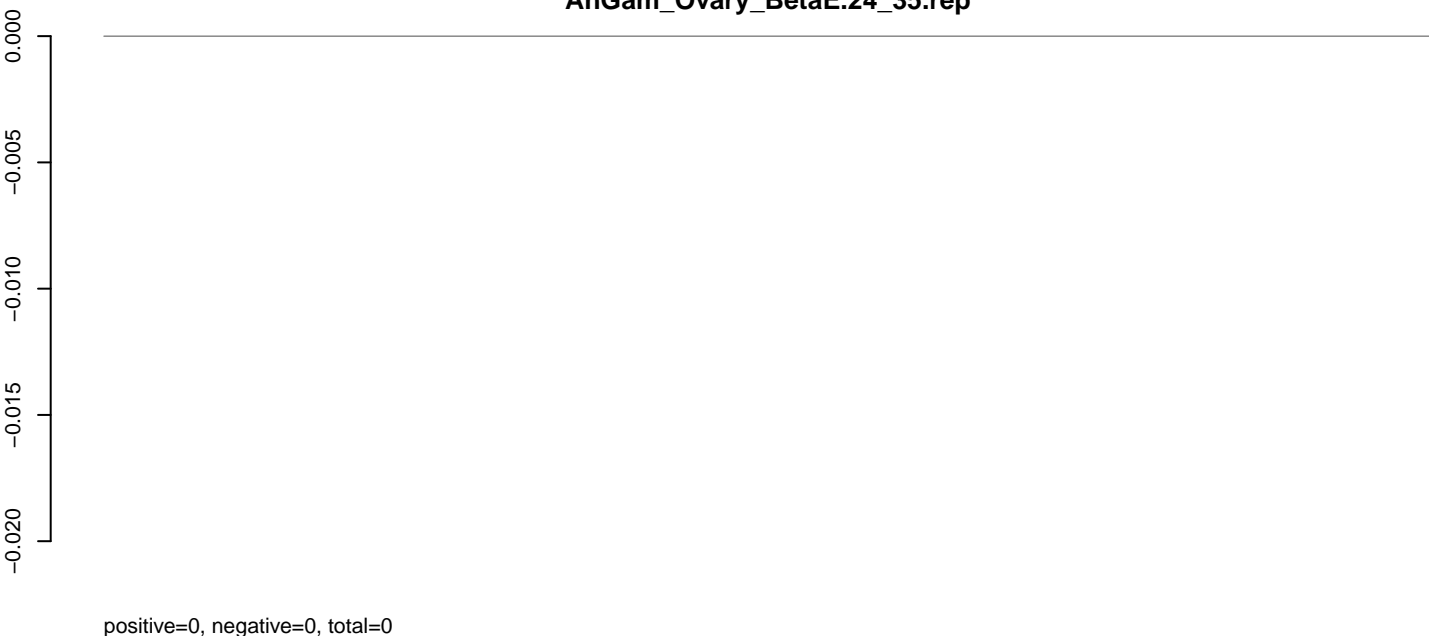
positive=0, negative=0, total=0, virus@NC_033716.1_Zhejiang_mosquito_virus_1_strain_mosZJ35497_hypothetical_protein_gene:1-9558



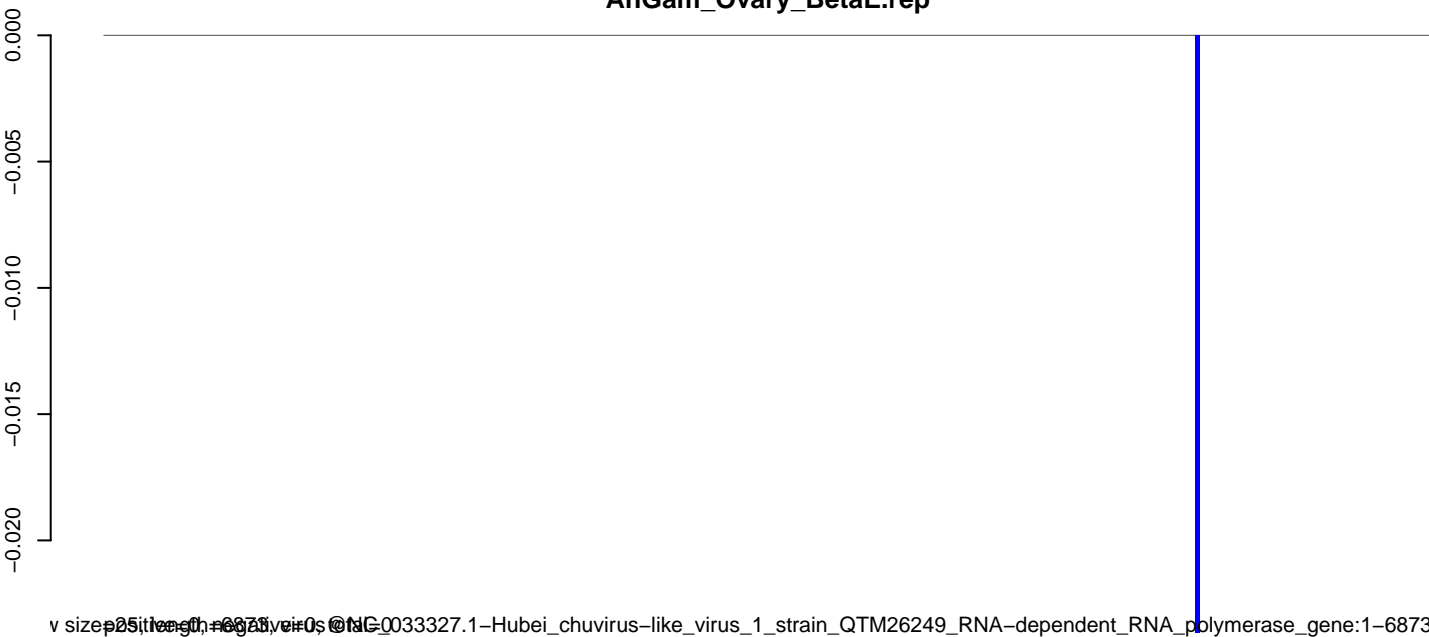
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



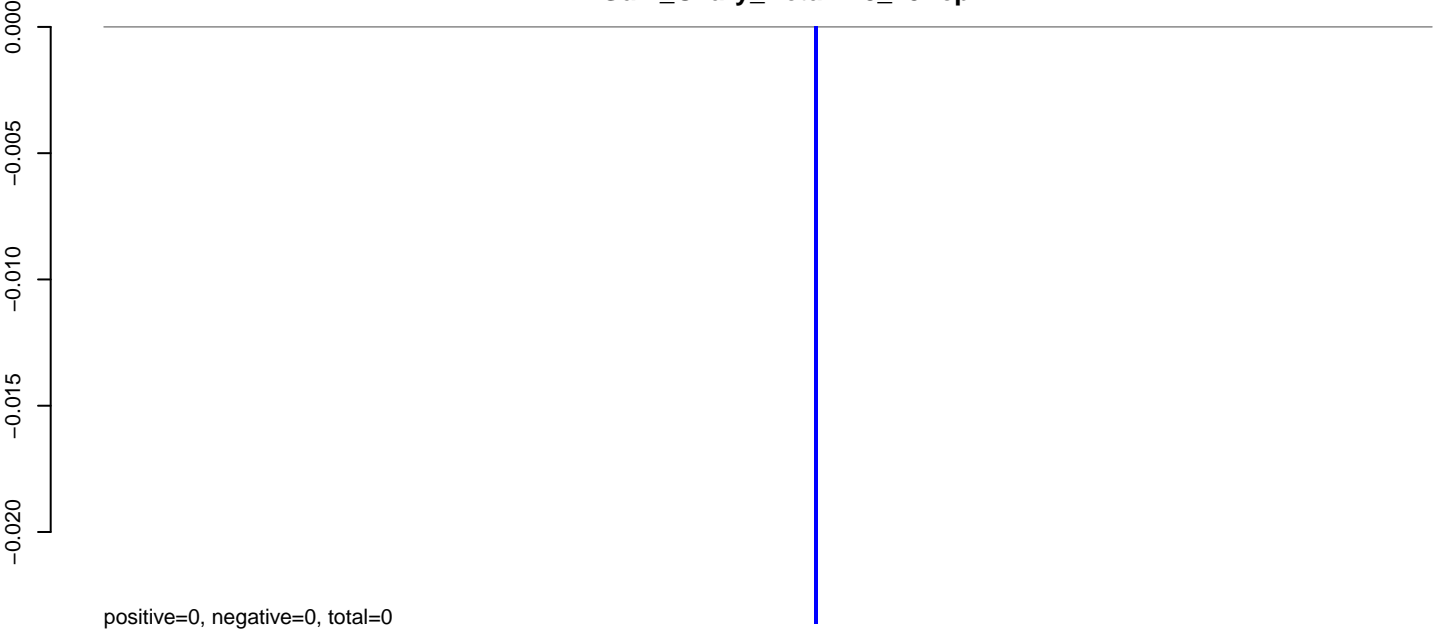
AnGam_Ovary_BetaE.rep



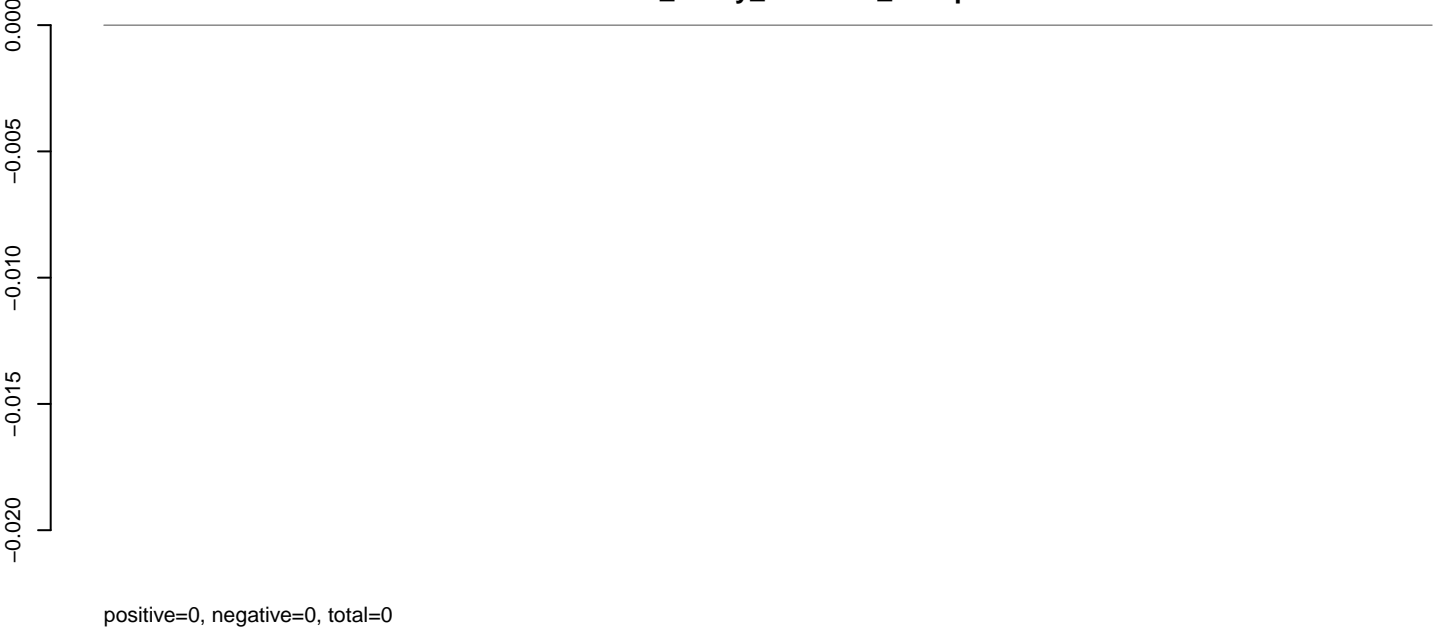
NC_033327.1-Hubei_chuvirus-like_virus_1_strain_QTM26249_RNA-dependent_RNA_polymerase_gene:1-6873

0 1000 2000 3000 4000 5000 6000 7000

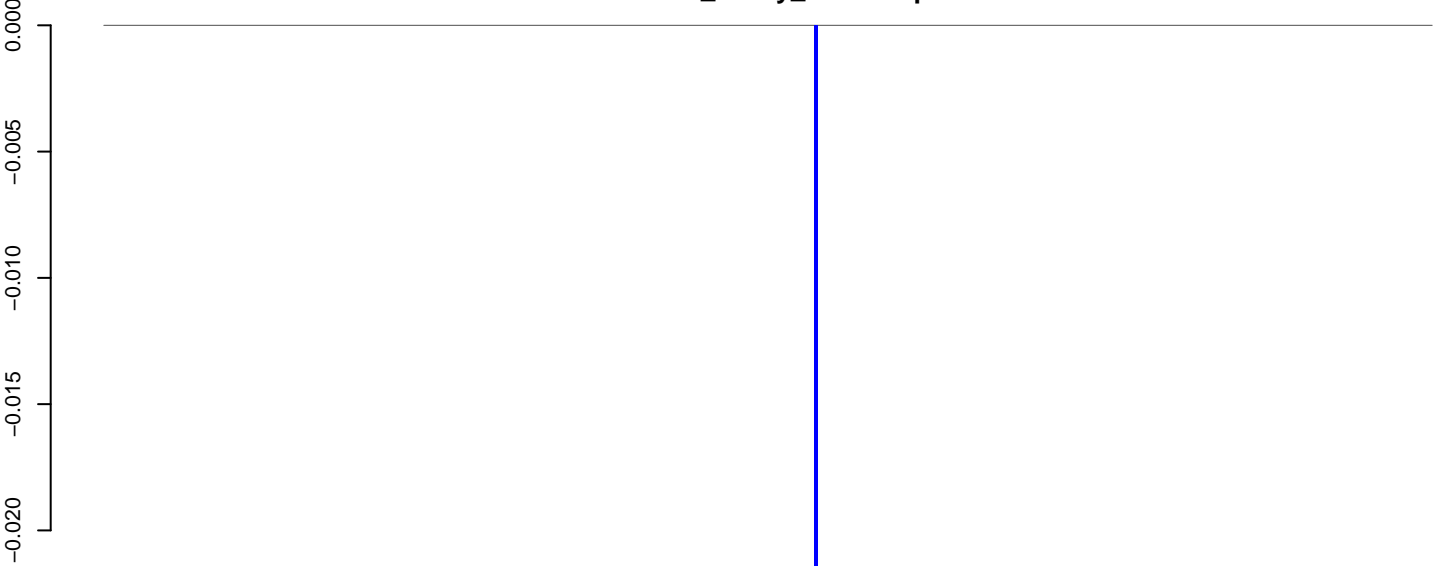
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



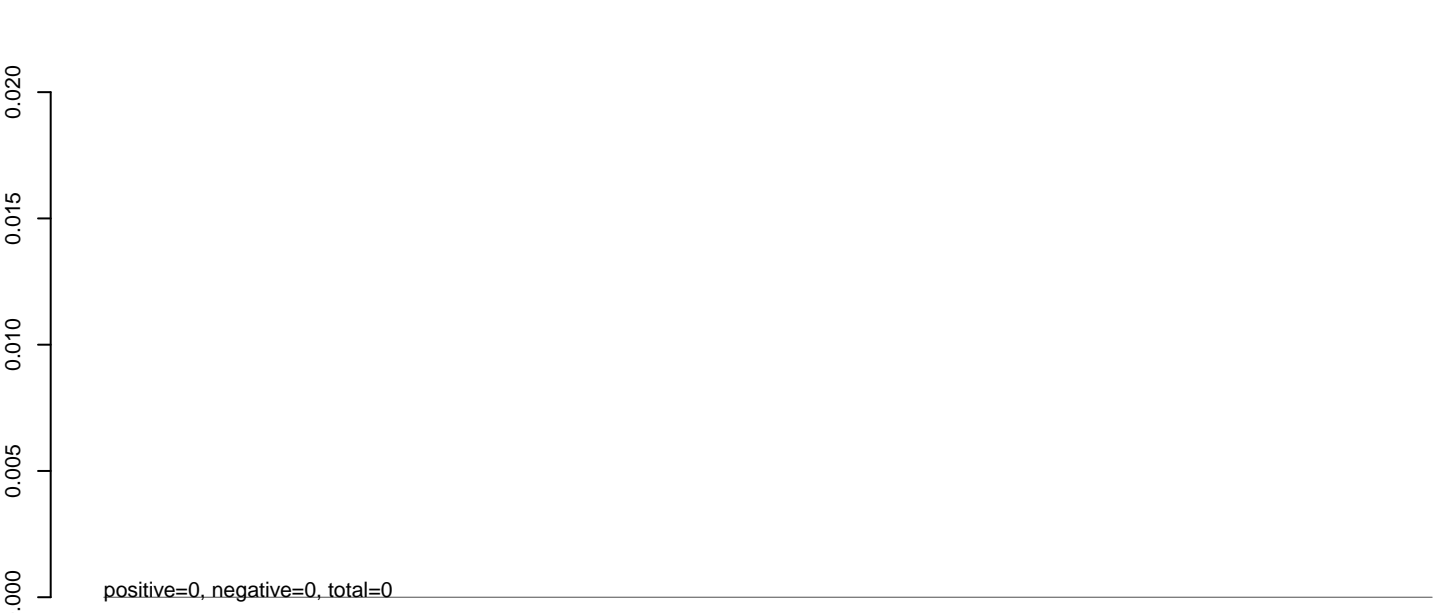
AnGam_Ovary_BetaE.rep



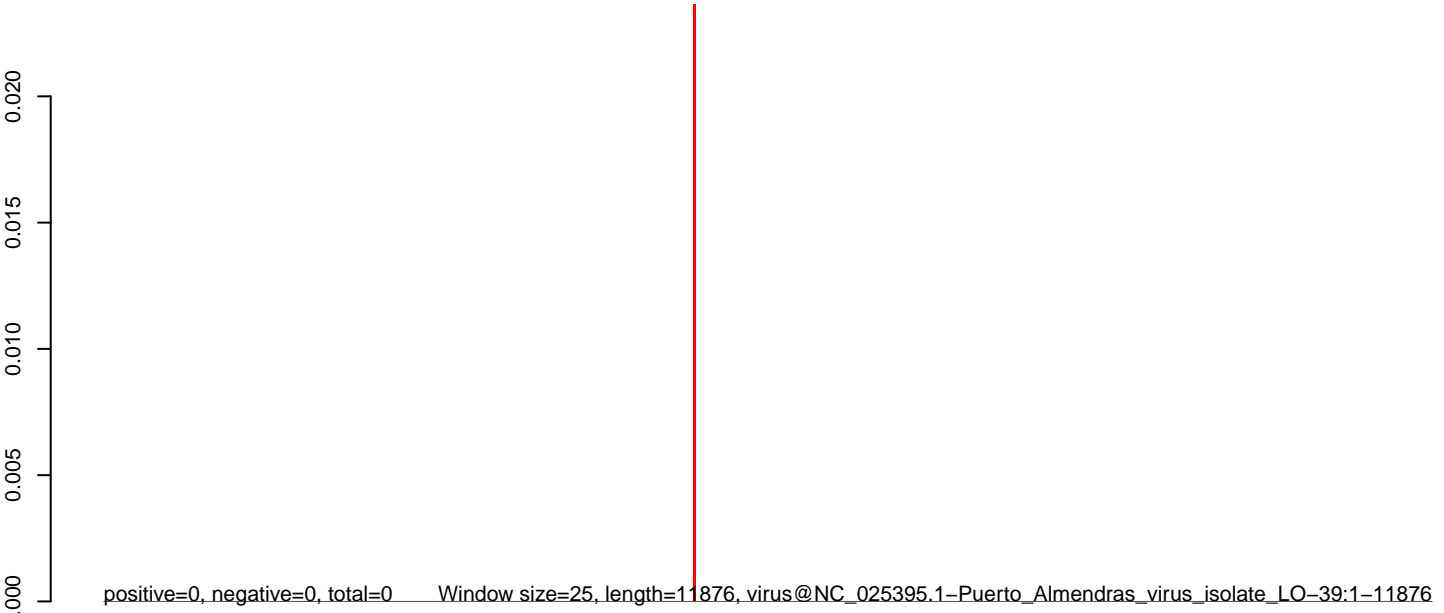
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

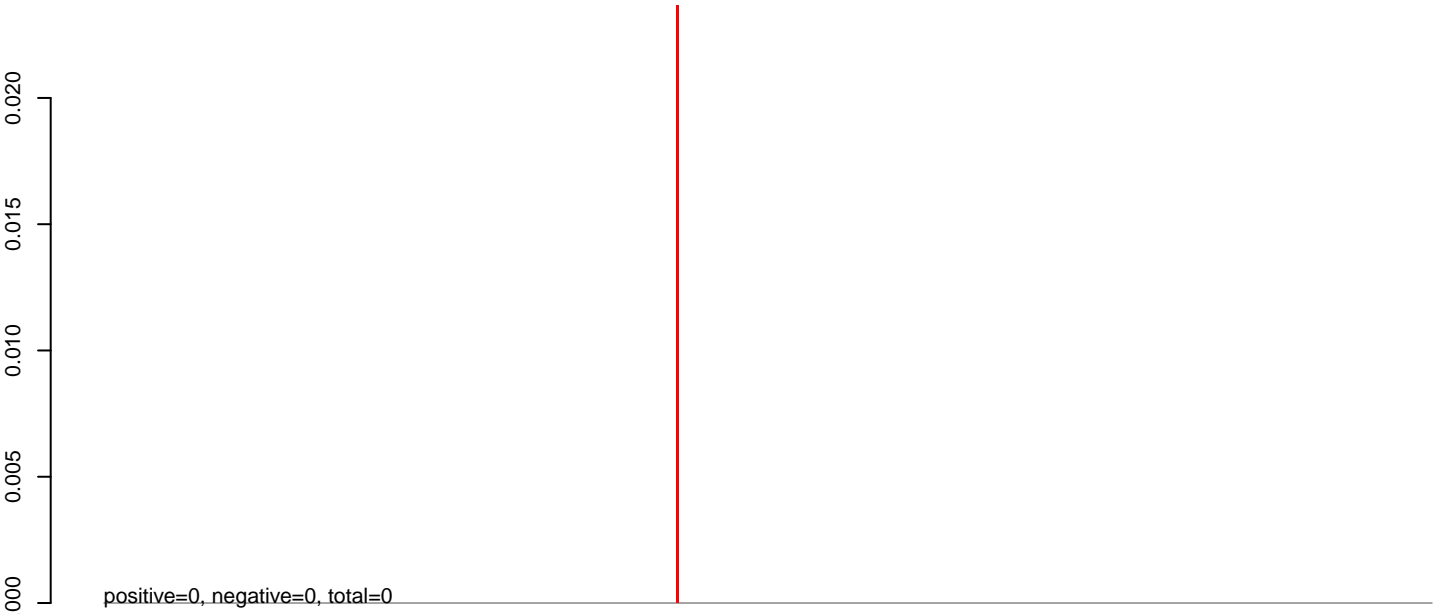


AnGam_Ovary_BetaE.rep

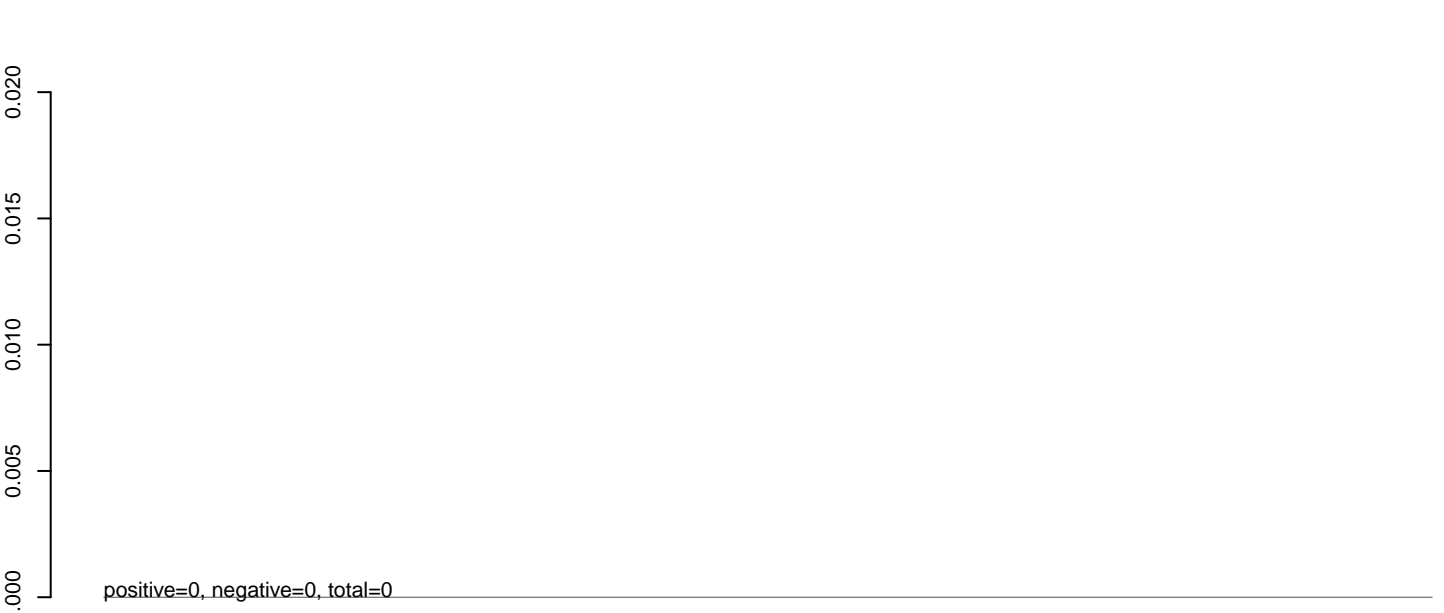


0 2000 4000 6000 8000 10000 12000

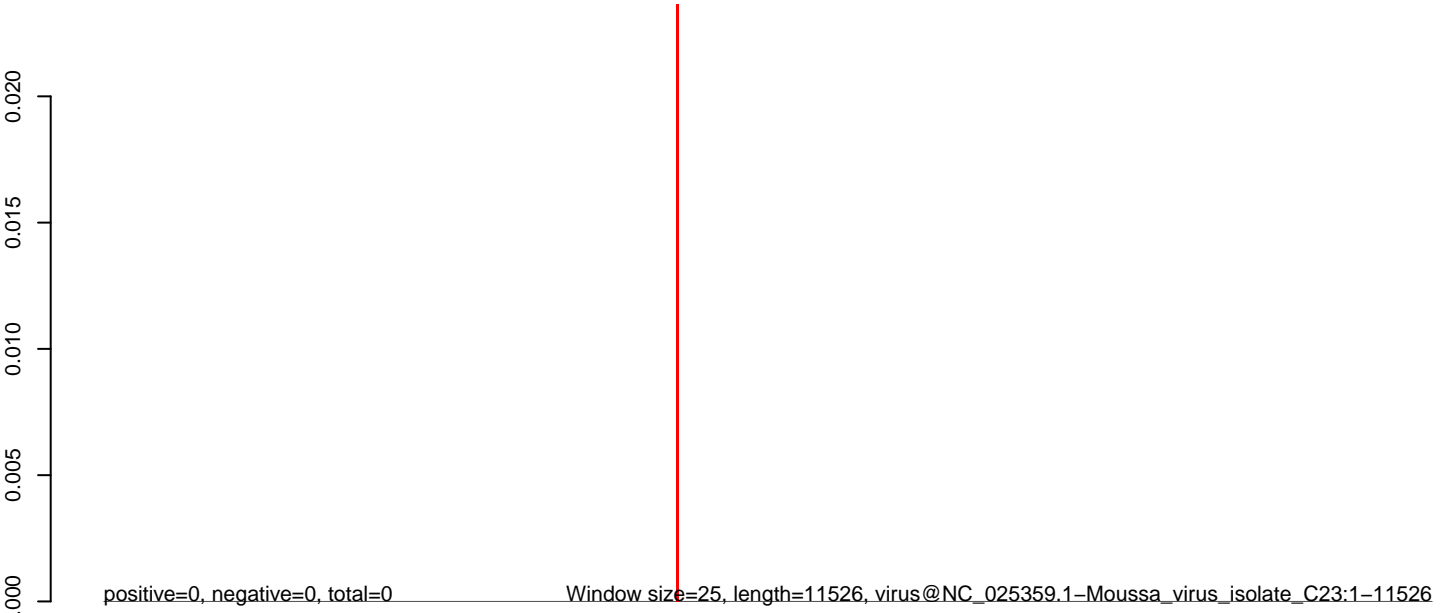
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

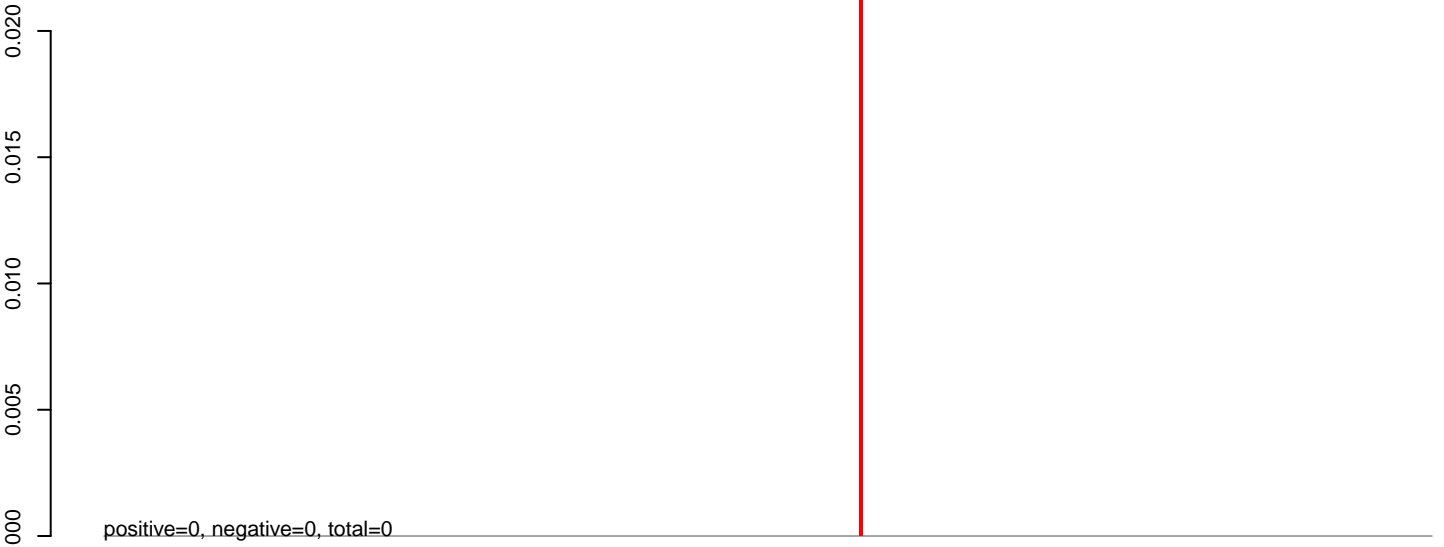


AnGam_Ovary_BetaE.rep

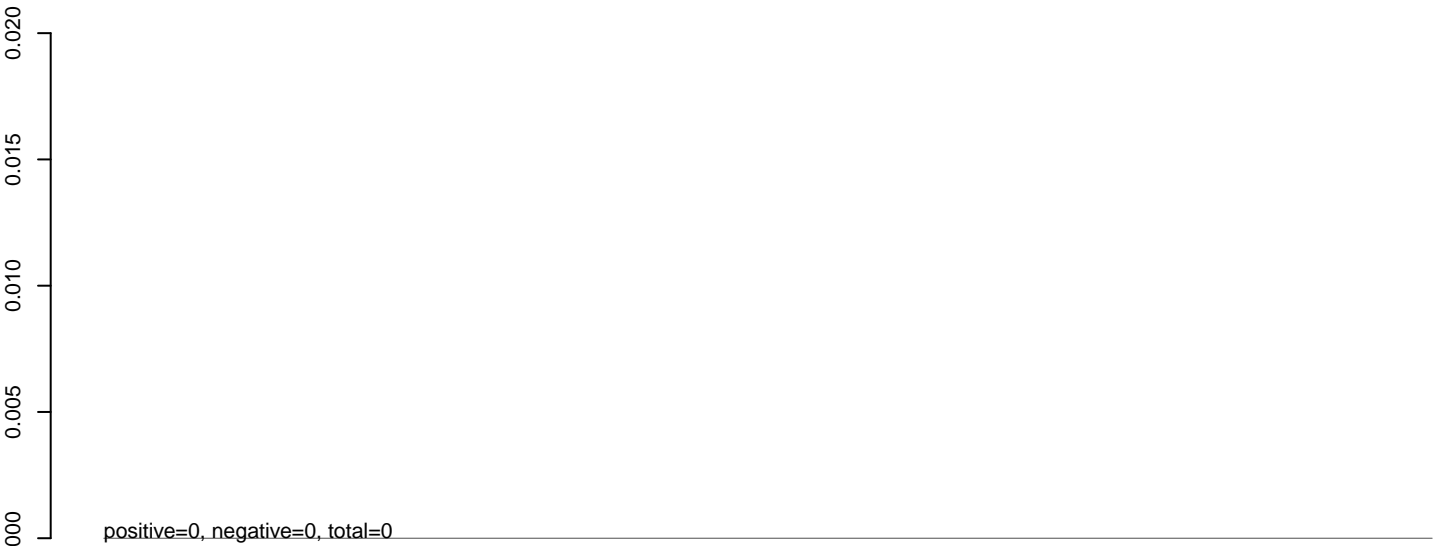


0 2000 4000 6000 8000 10000 12000

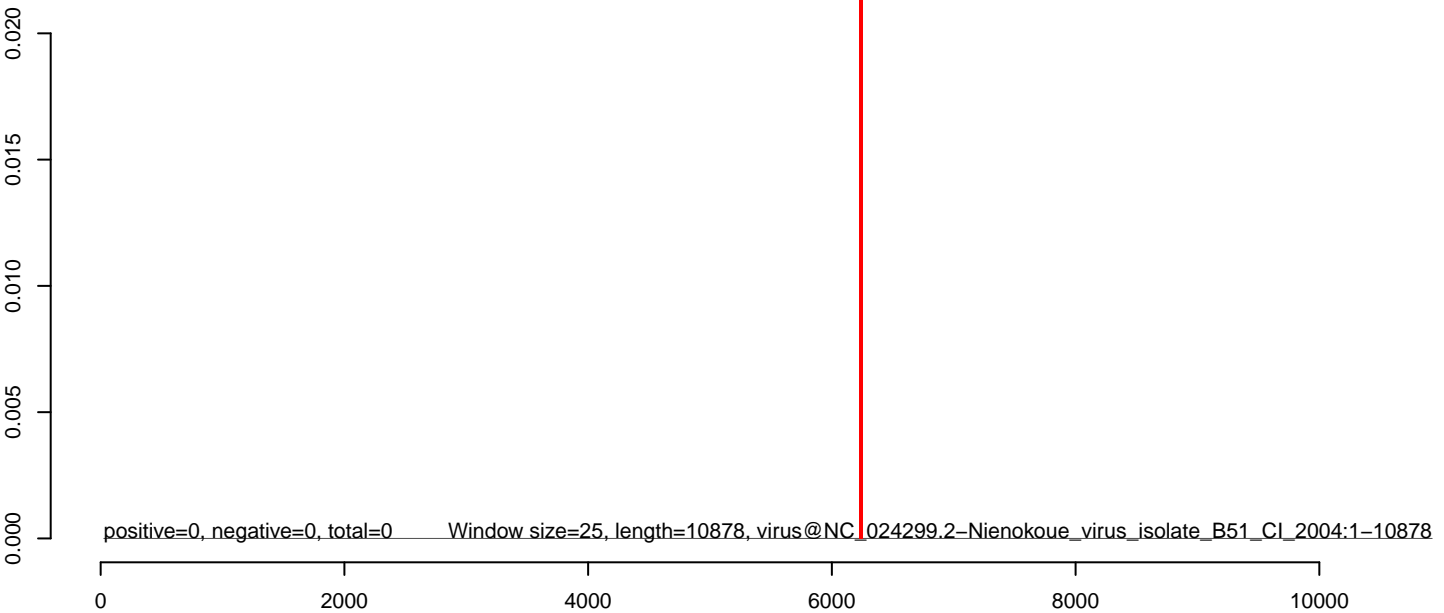
AnGam_Ovary_BetaE.18_23.rep



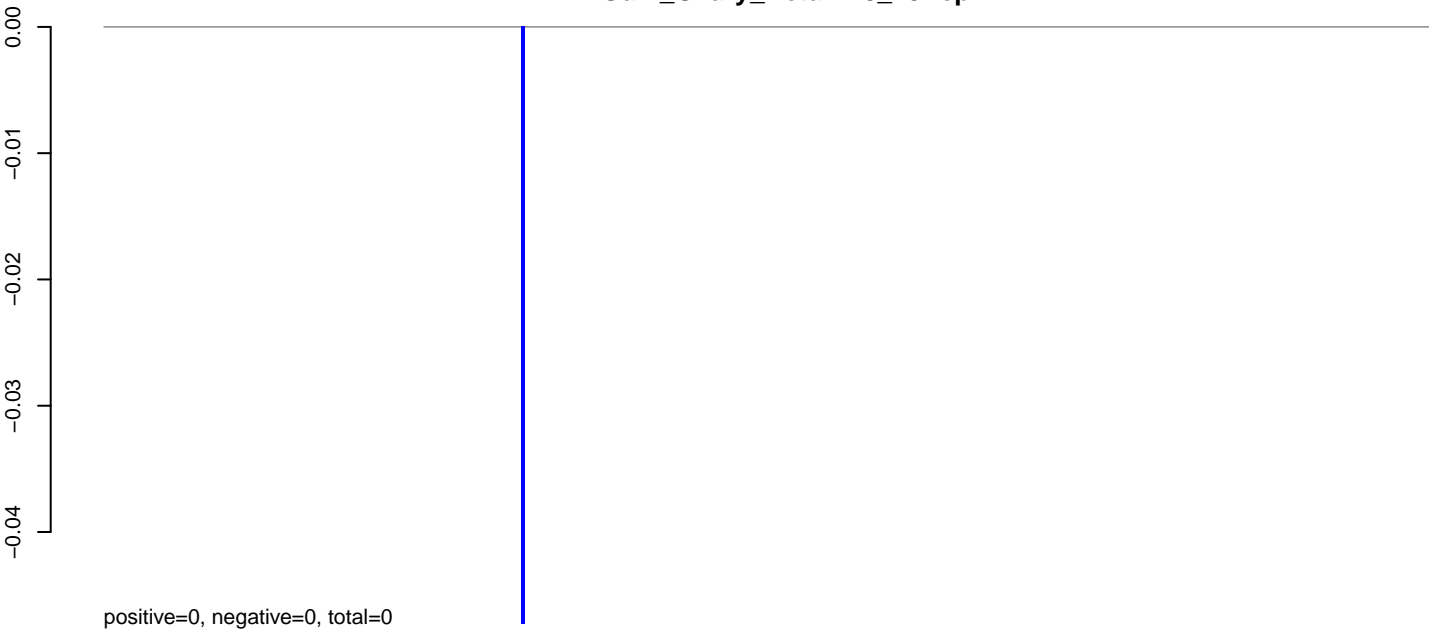
AnGam_Ovary_BetaE.24_35.rep



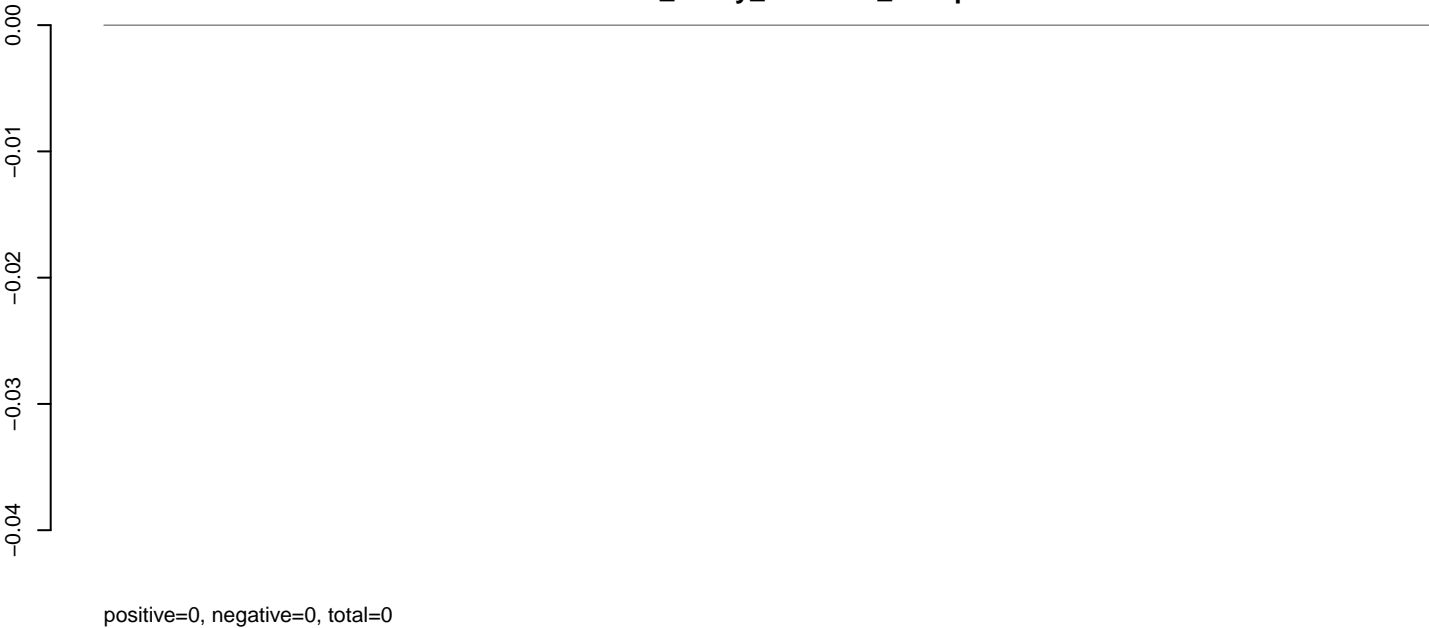
AnGam_Ovary_BetaE.rep



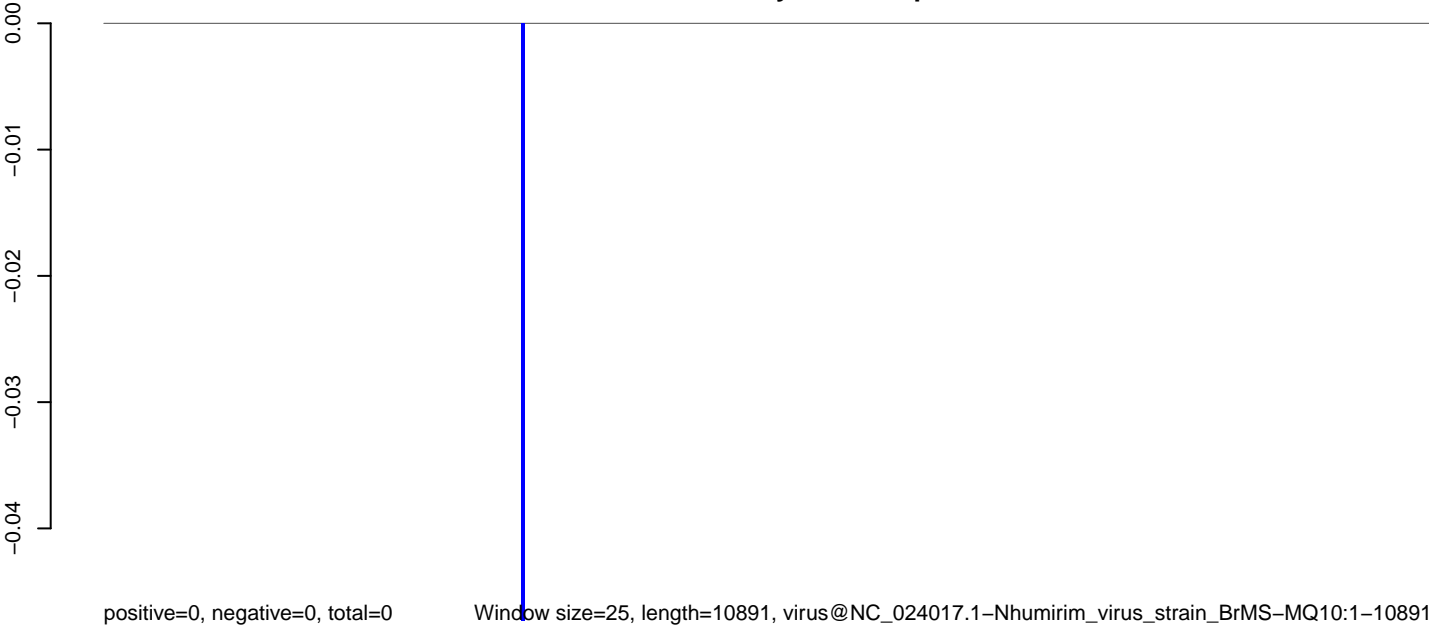
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

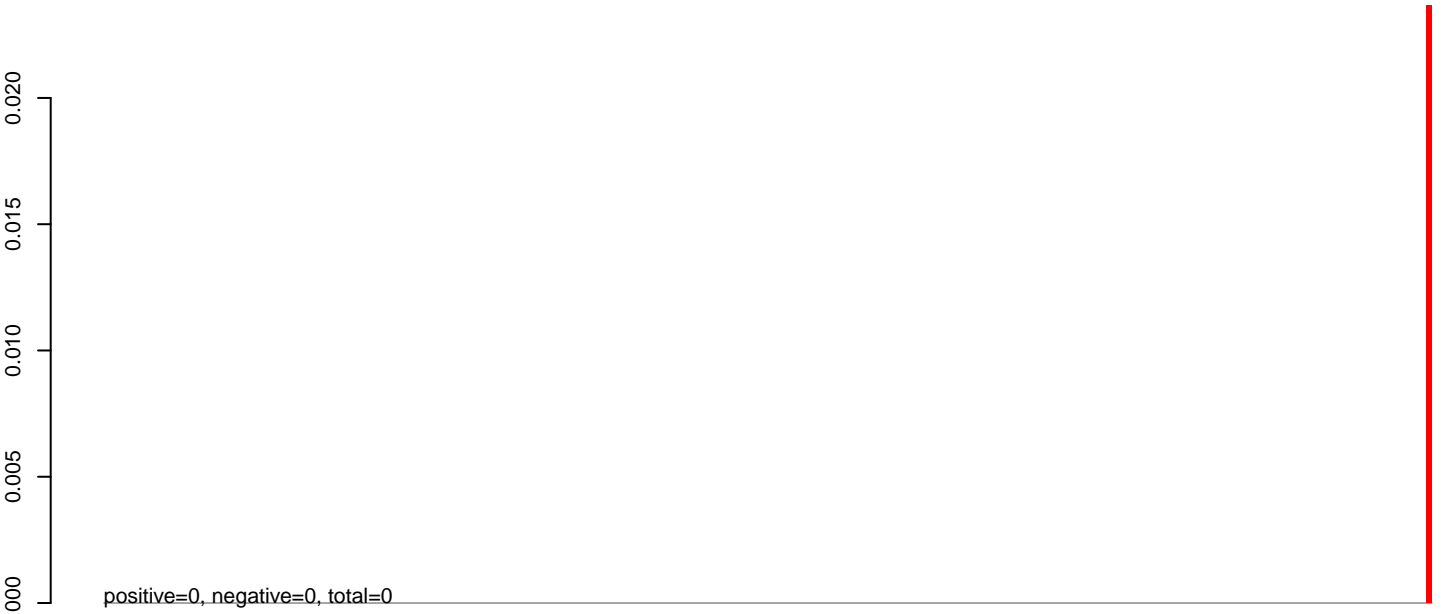


AnGam_Ovary_BetaE.rep

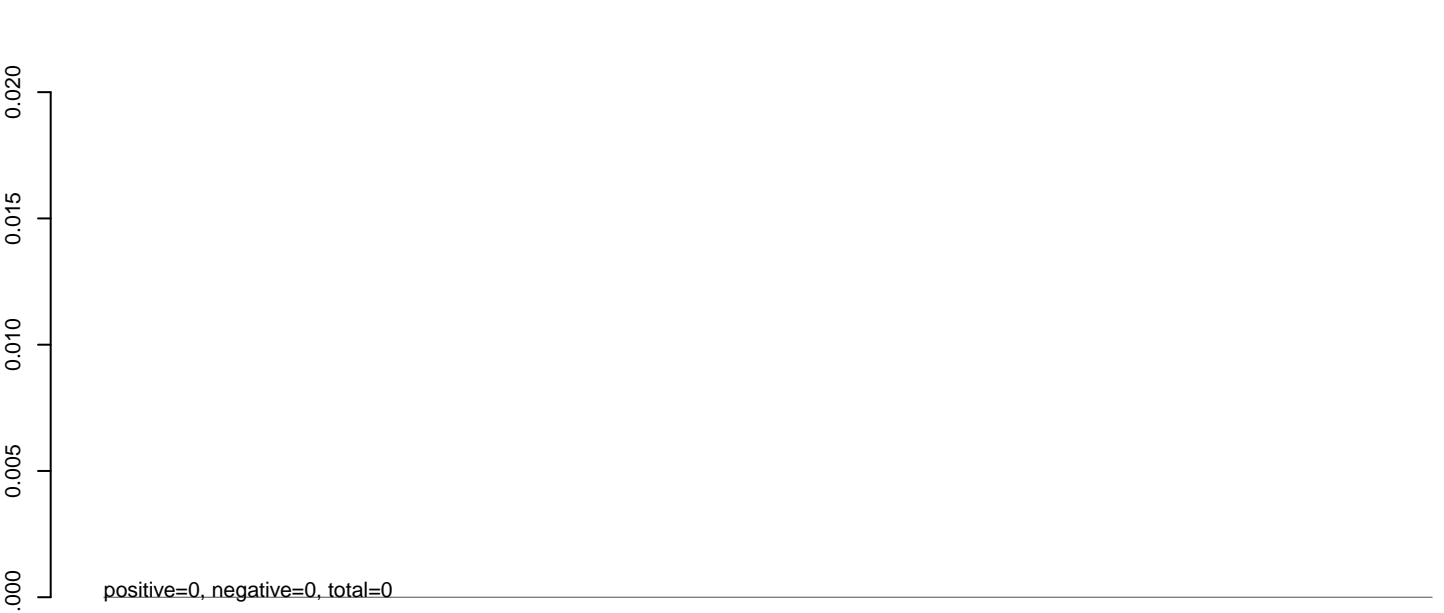


0 2000 4000 6000 8000 10000

AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



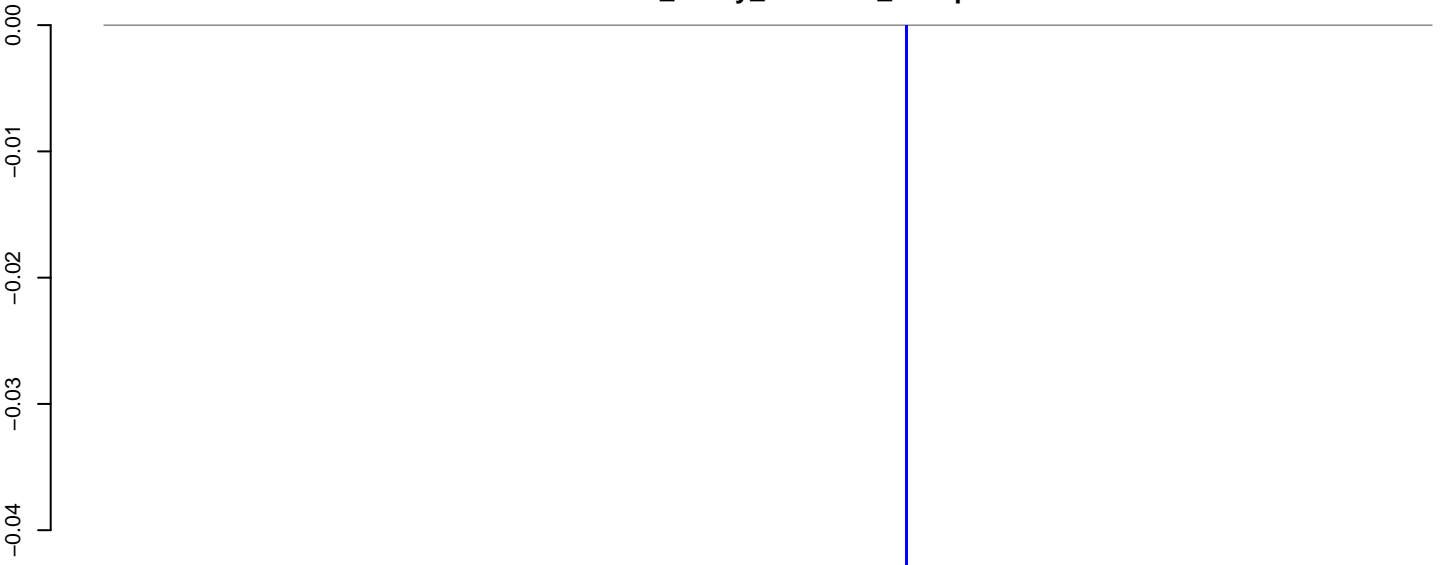
0 2000 4000 6000 8000 10000 12000
Virus: 11624, length=911624, virus@NC_023812.1-Madariaga_virus_strain_MADV_Cebus_apella_BRA_BEAN5122_1956:1-11624

AnGam_Ovary_BetaE.18_23.rep



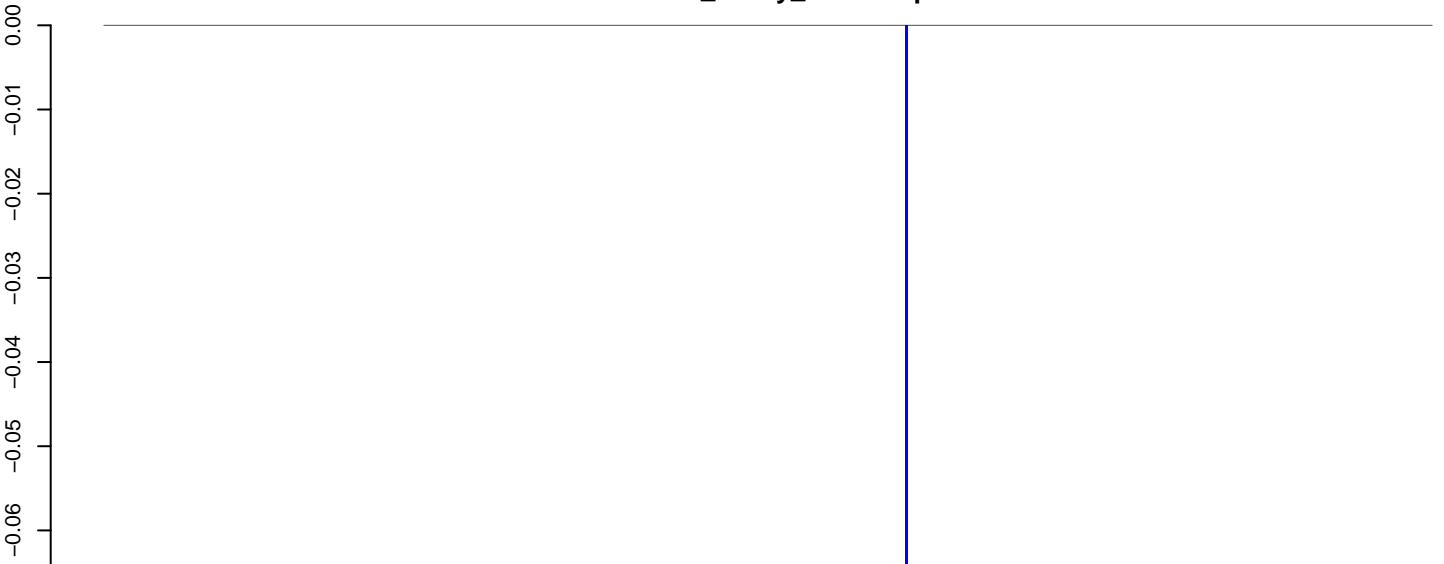
positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep



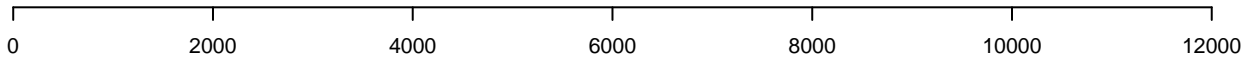
positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep



positive=0, negative=0, total=0

Window size=25, length=13298, virus@NC_020804.1-Tibrogargan_virus_strain_CS132:1-13298



AnGam_Ovary_BetaE.18_23.rep

0.3
0.2
0.1
0.0
-0.1

positive=1, negative=0, total=1



AnGam_Ovary_BetaE.24_35.rep

0.3
0.2
0.1
0.0
-0.1

positive=0, negative=0, total=0



AnGam_Ovary_BetaE.rep

0.3
0.2
0.1
0.0
-0.1

positive=1, negative=0, total=1

Window size=25, length=11634, virus@NC_018615.1-Eilat_virus_isolate_EO329:1-11634



0 2000 4000 6000 8000 10000 12000

AnGam_Ovary_BetaE.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep

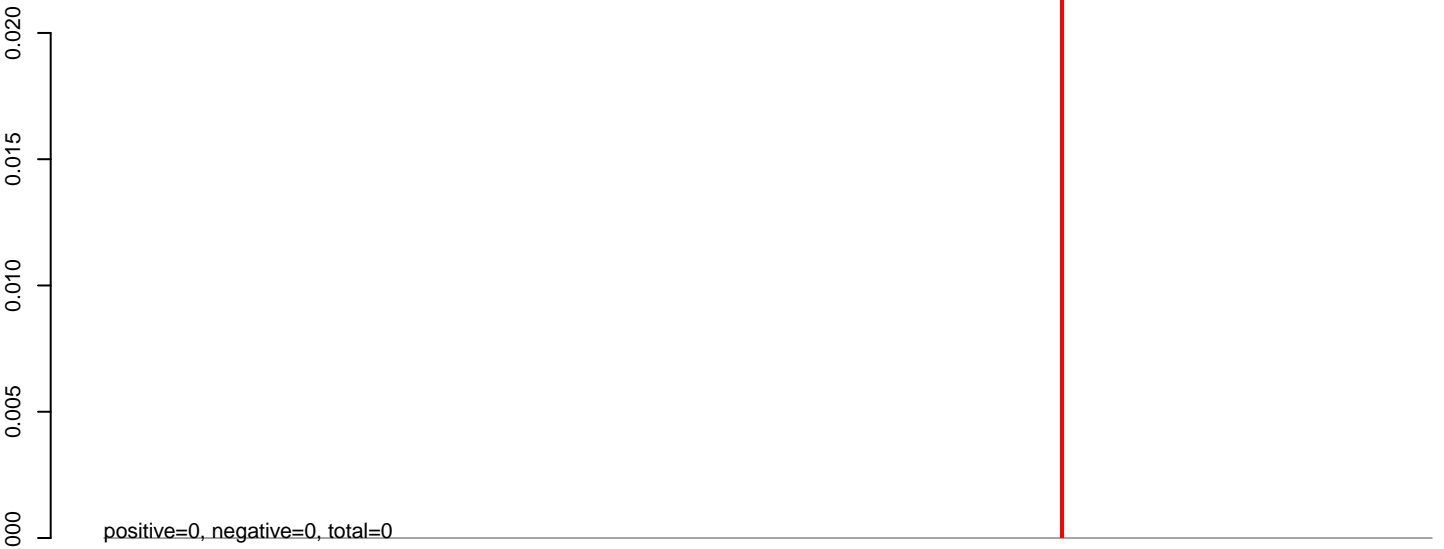
0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

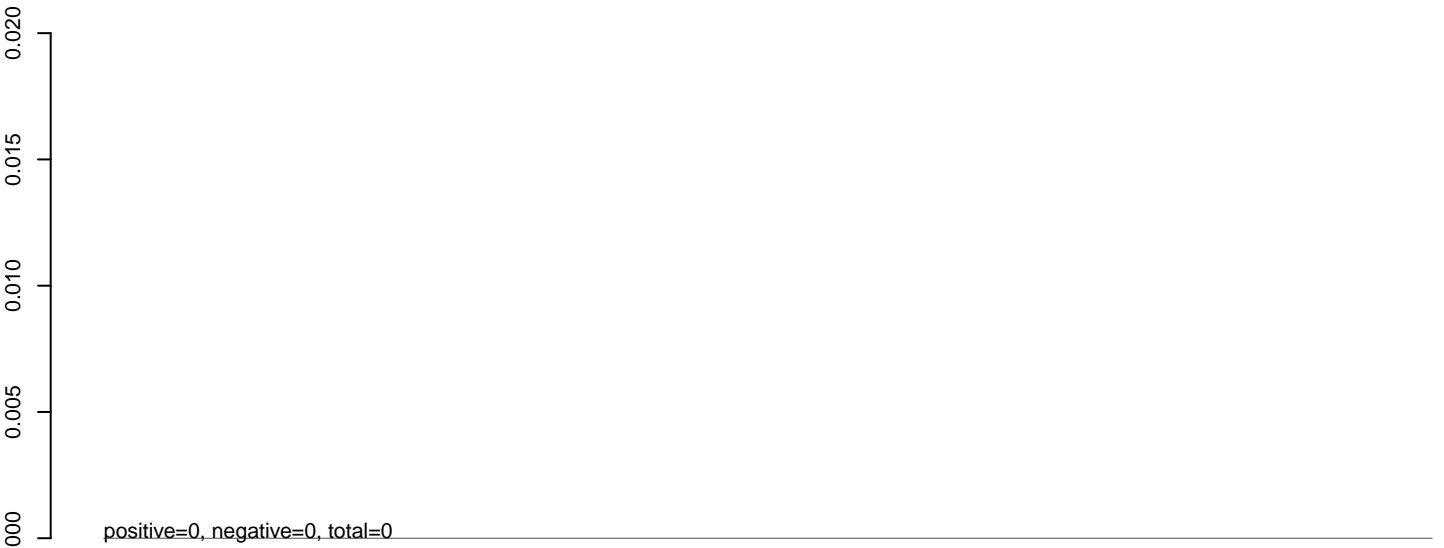
Window size=25, length=15867, virus@NC_017714.1-Kotonkan_virus:1-15867

0 5000 10000 15000

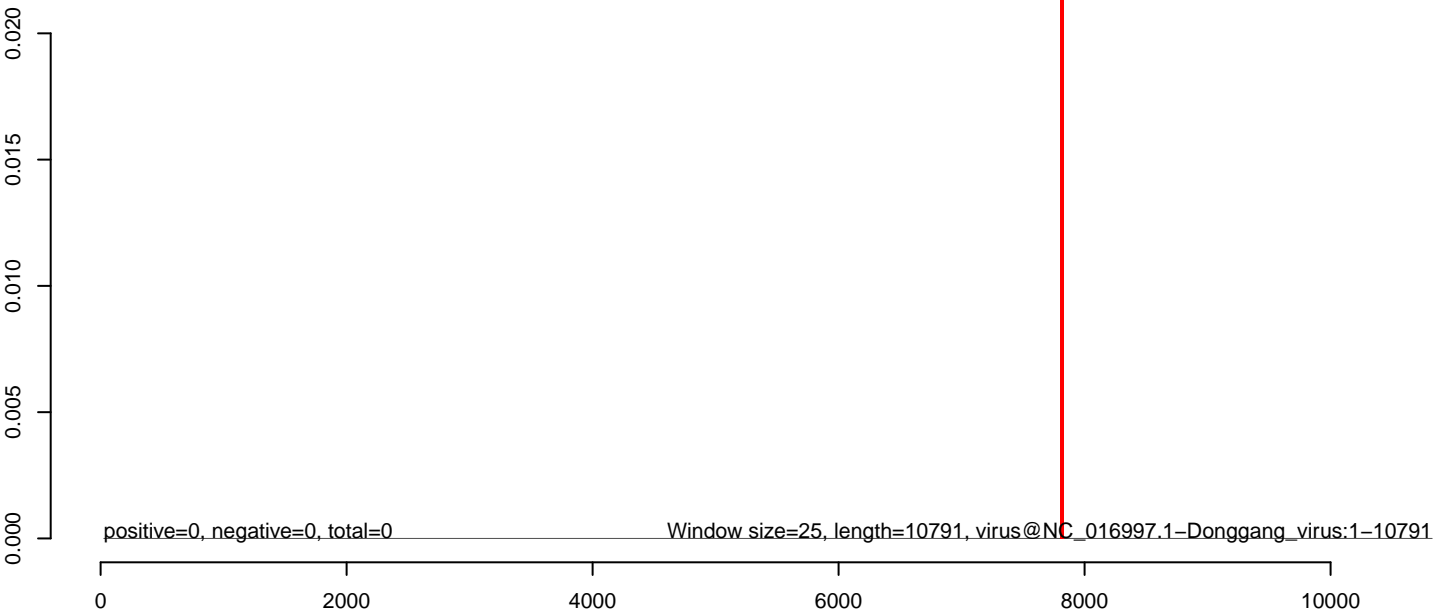
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



AnGam_Ovary_BetaE.18_23.rep

0.4
0.2
0.0
-0.2

positive=1, negative=0, total=2

AnGam_Ovary_BetaE.24_35.rep

0.4
0.2
0.0
-0.2

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep

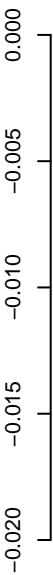
0.6
0.4
0.2
0.0
-0.2

positive=1, negative=0, total=2

Window size=25, length=11616, virus@NC_016961.1-Whataroa_virus:1-11616

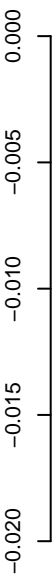
0 2000 4000 6000 8000 10000 12000

AnGam_Ovary_BetaE.18_23.rep



positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep



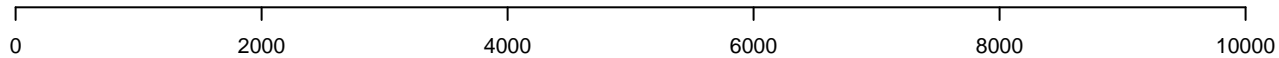
positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep

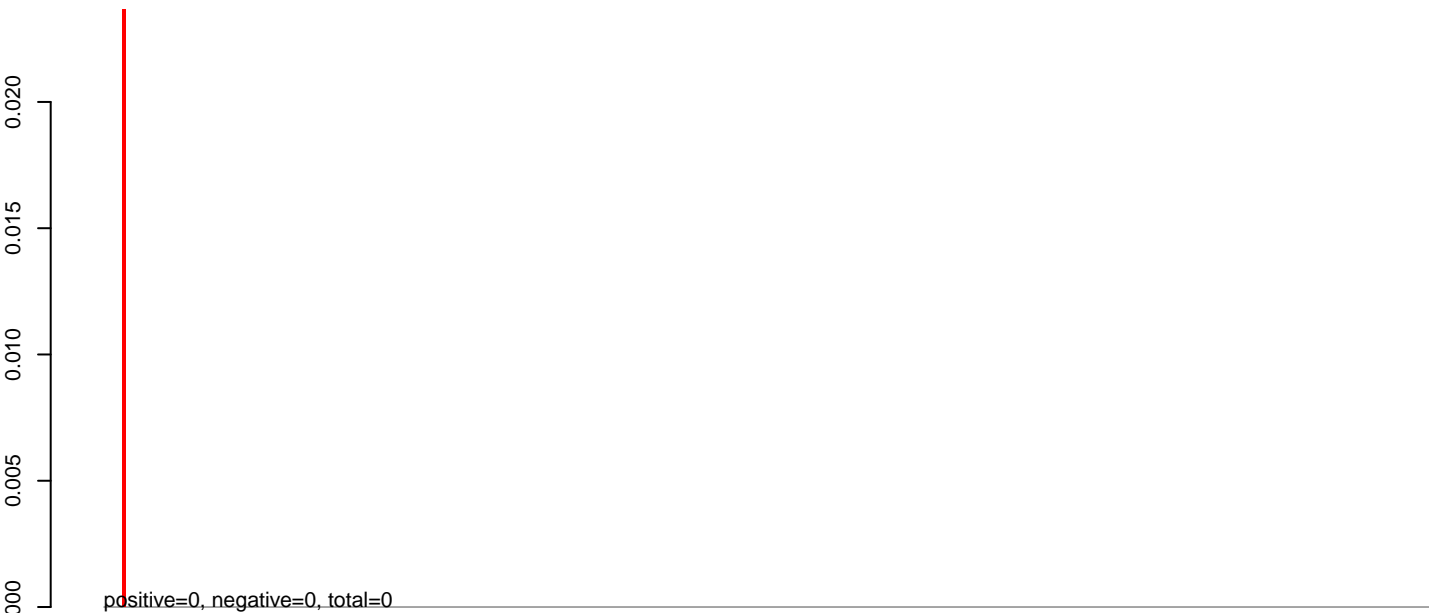


positive=0, negative=0, total=0

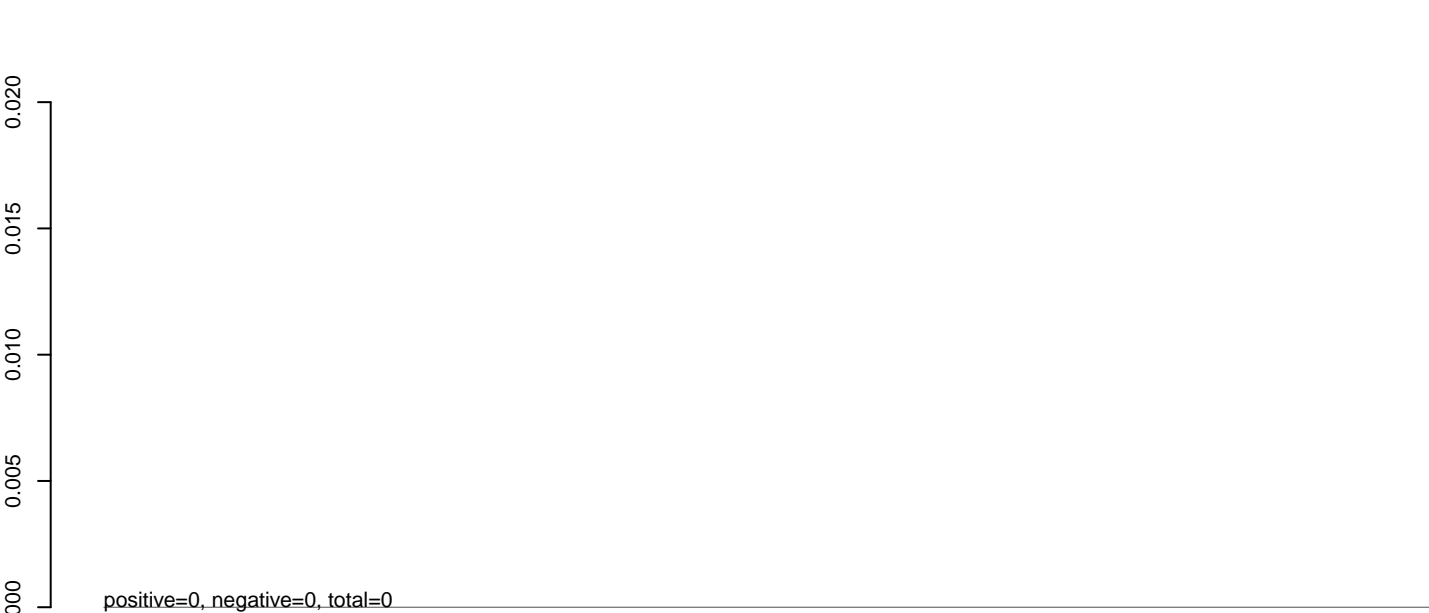
Window size=25, length=10793, virus@NC_008719.1-Sepik_virus:1-10793



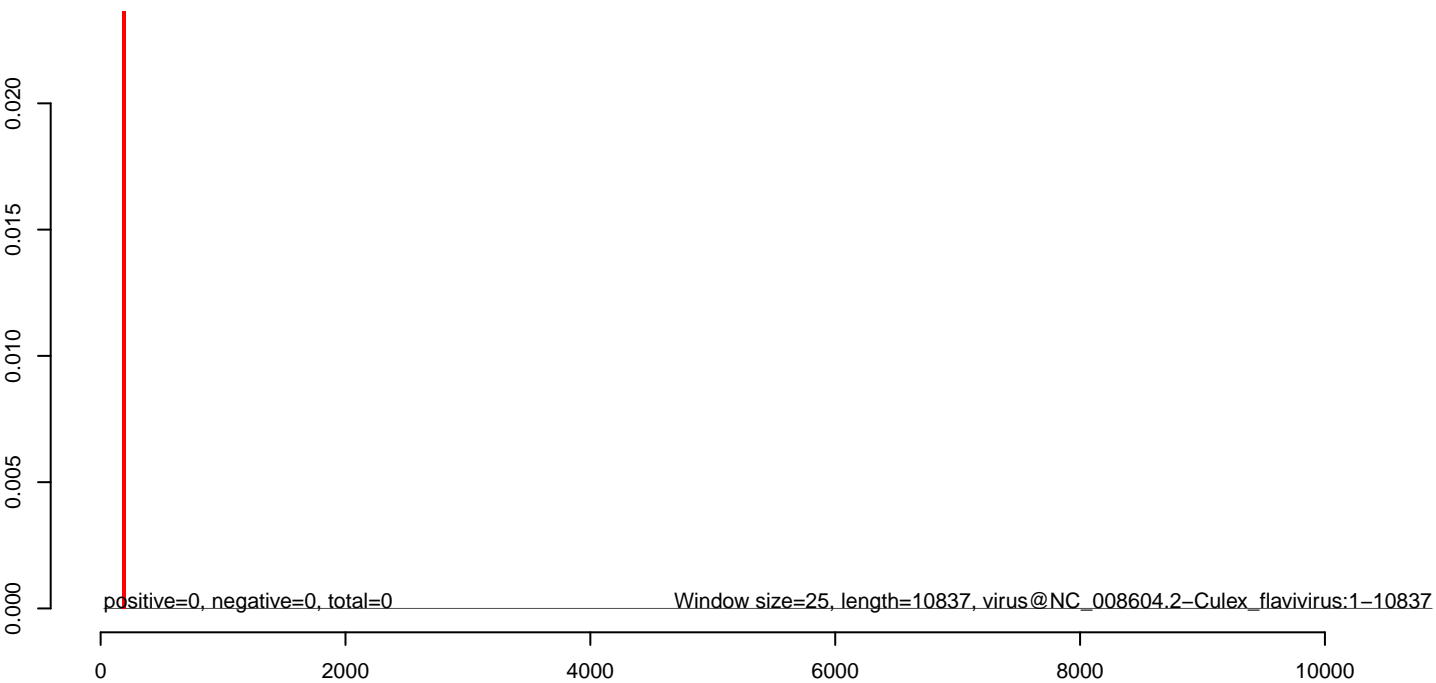
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



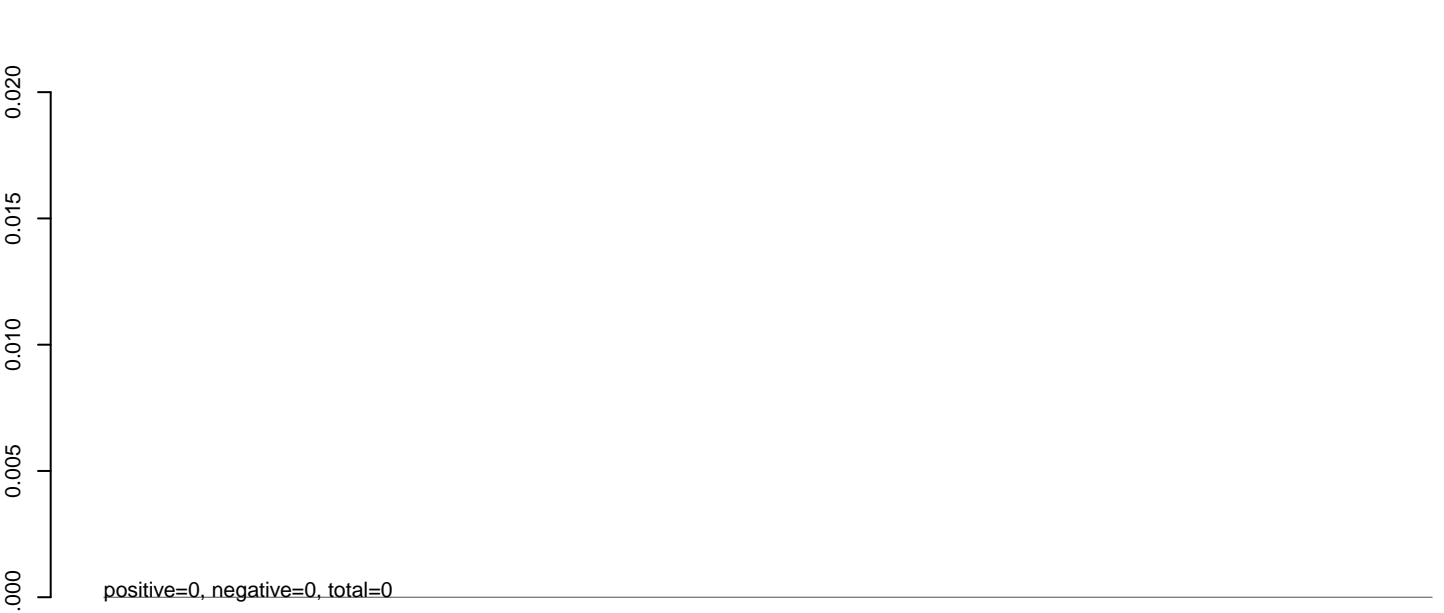
AnGam_Ovary_BetaE.rep



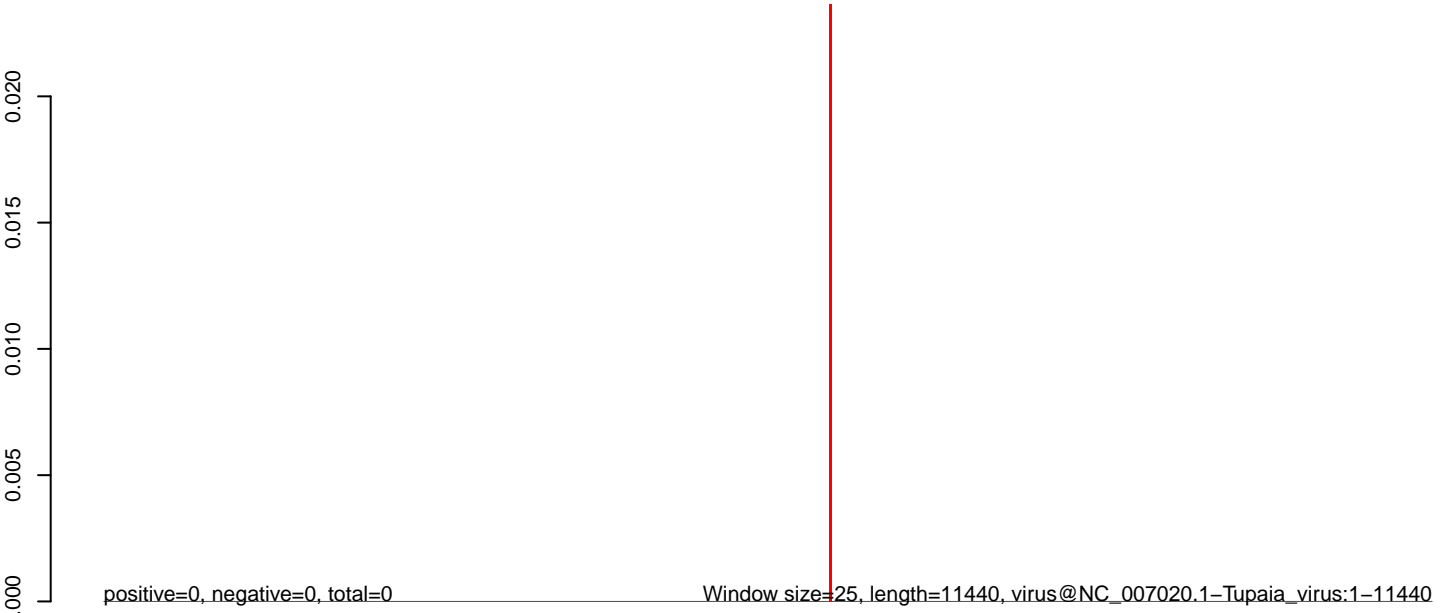
AnGam_Ovary_BetaE.18_23.rep



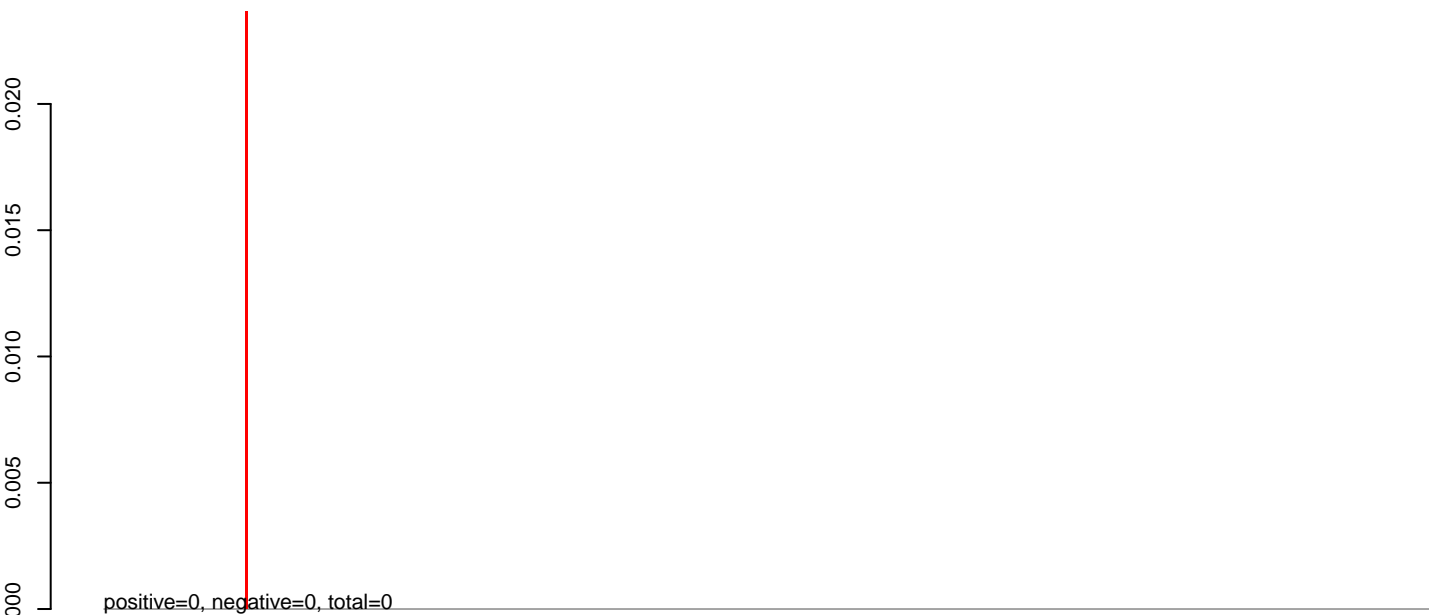
AnGam_Ovary_BetaE.24_35.rep



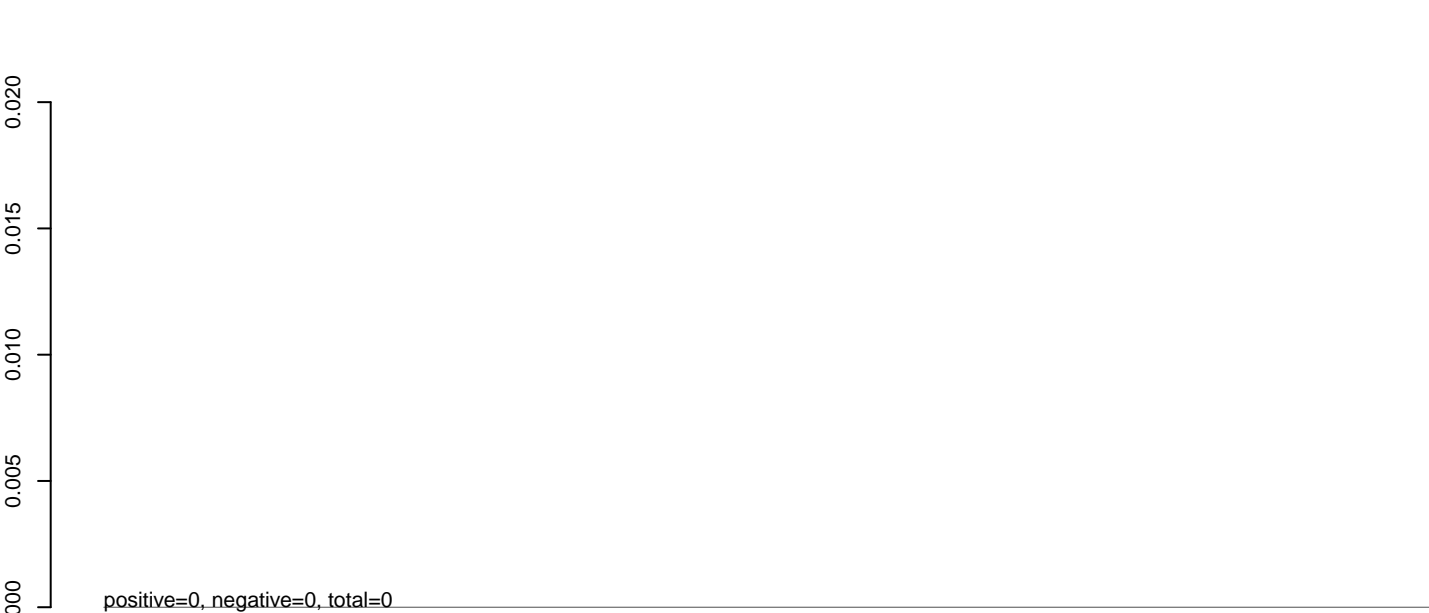
AnGam_Ovary_BetaE.rep



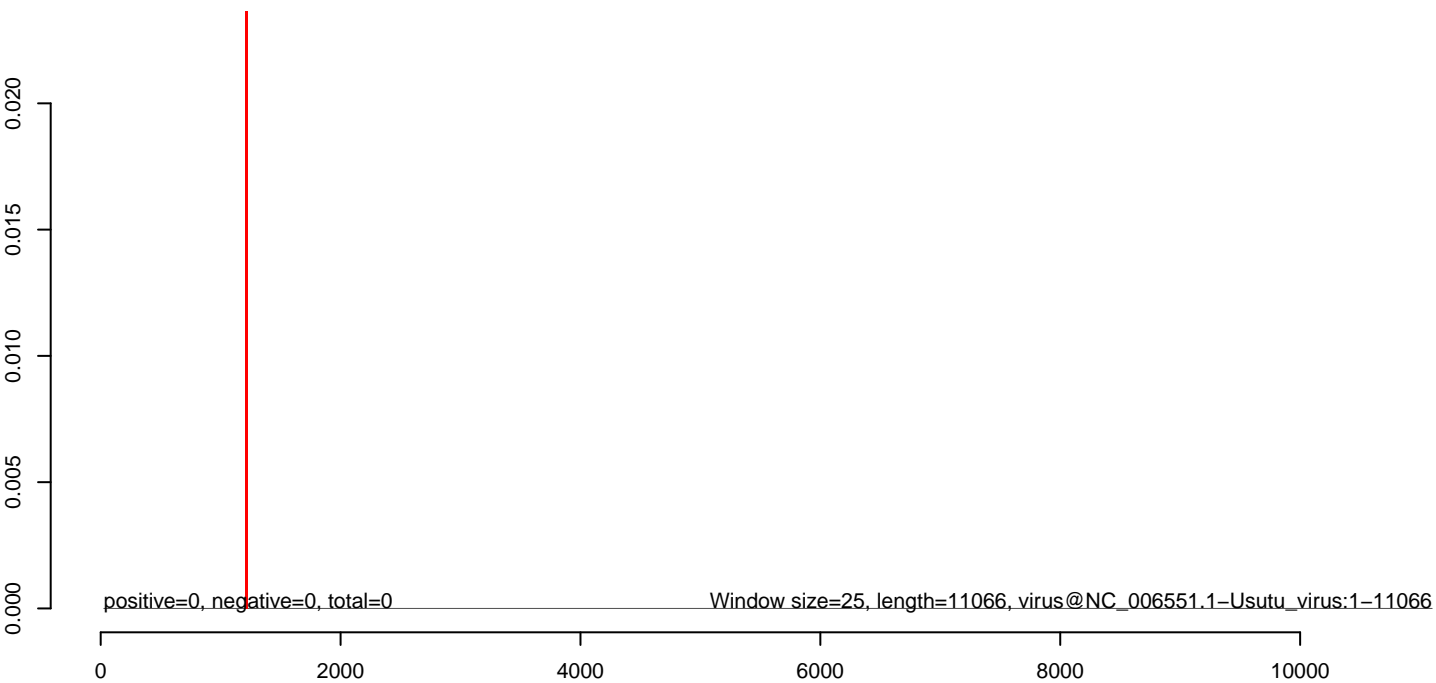
AnGam_Ovary_BetaE.18_23.rep



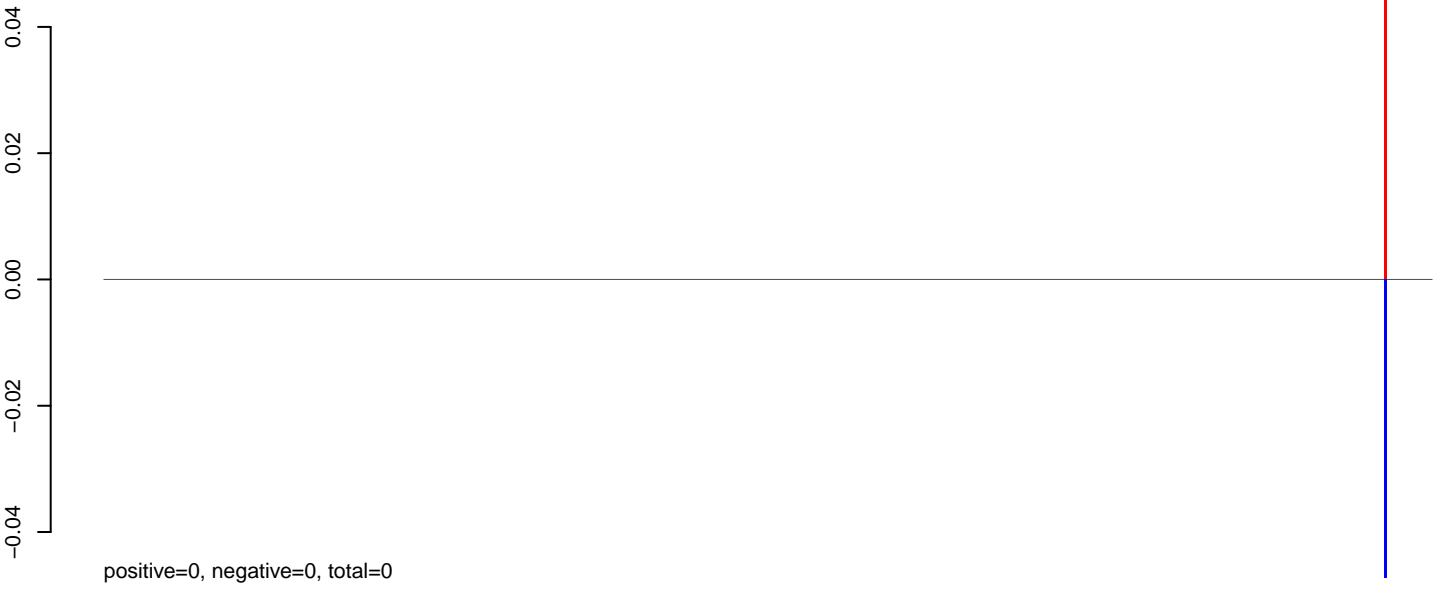
AnGam_Ovary_BetaE.24_35.rep



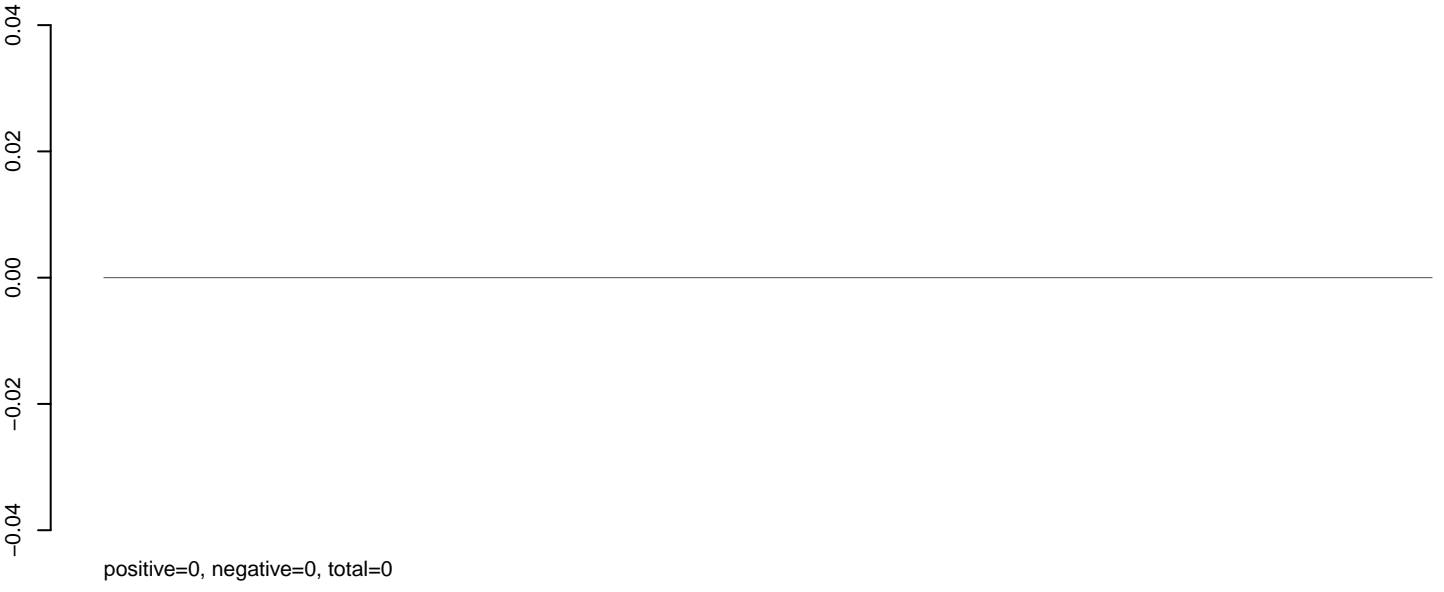
AnGam_Ovary_BetaE.rep



AnGam_Ovary_BetaE.18_23.rep



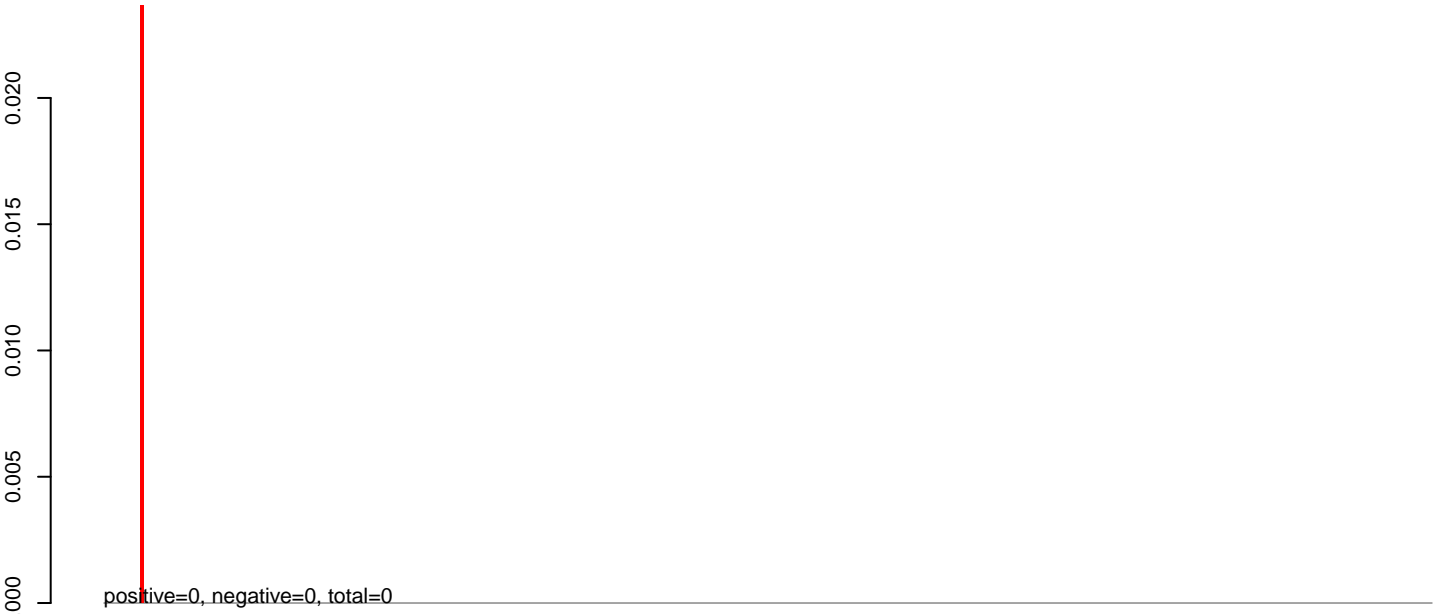
AnGam_Ovary_BetaE.24_35.rep



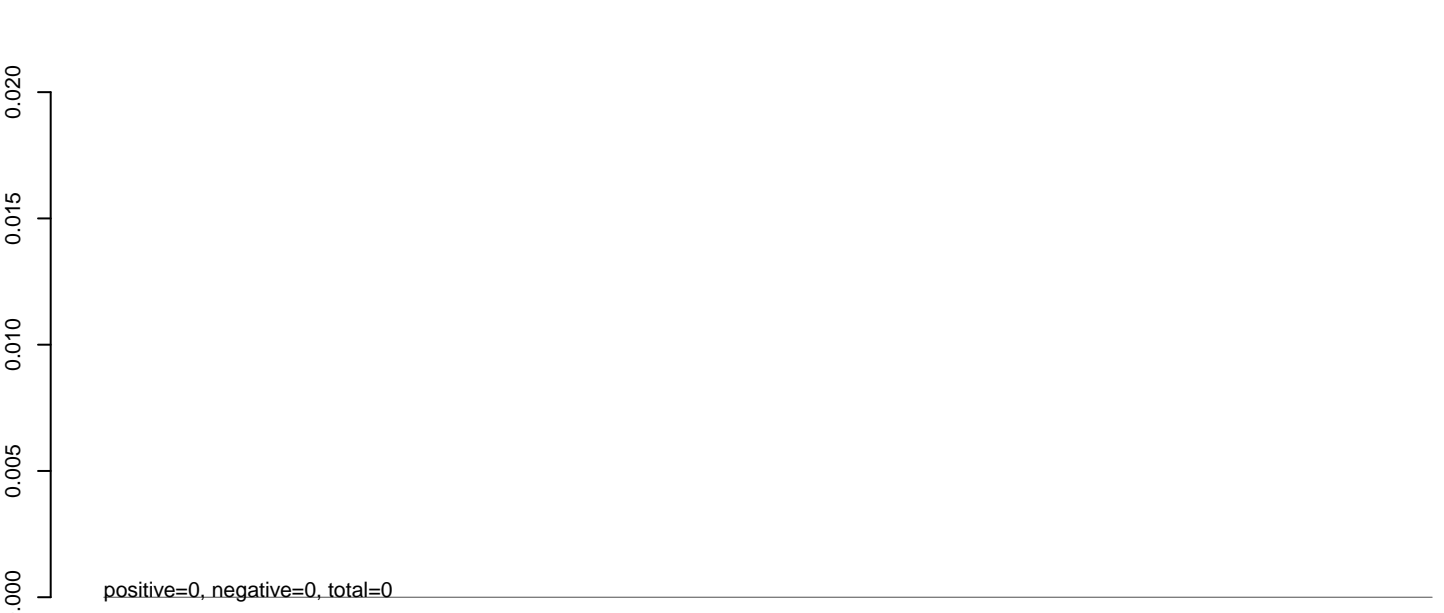
AnGam_Ovary_BetaE.rep



AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

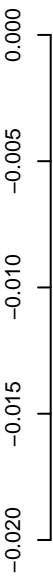


AnGam_Ovary_BetaE.rep



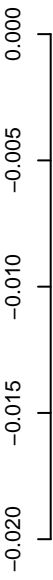
WINDOMAS, R. (2015) NC_003924.1-Cricket paralysis virus nonstructural polyprotein and structural polyprotein genes:1-9185

AnGam_Ovary_BetaE.18_23.rep



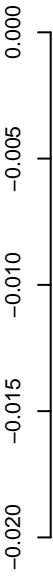
positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep



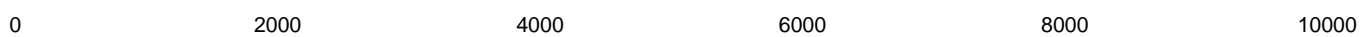
positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep

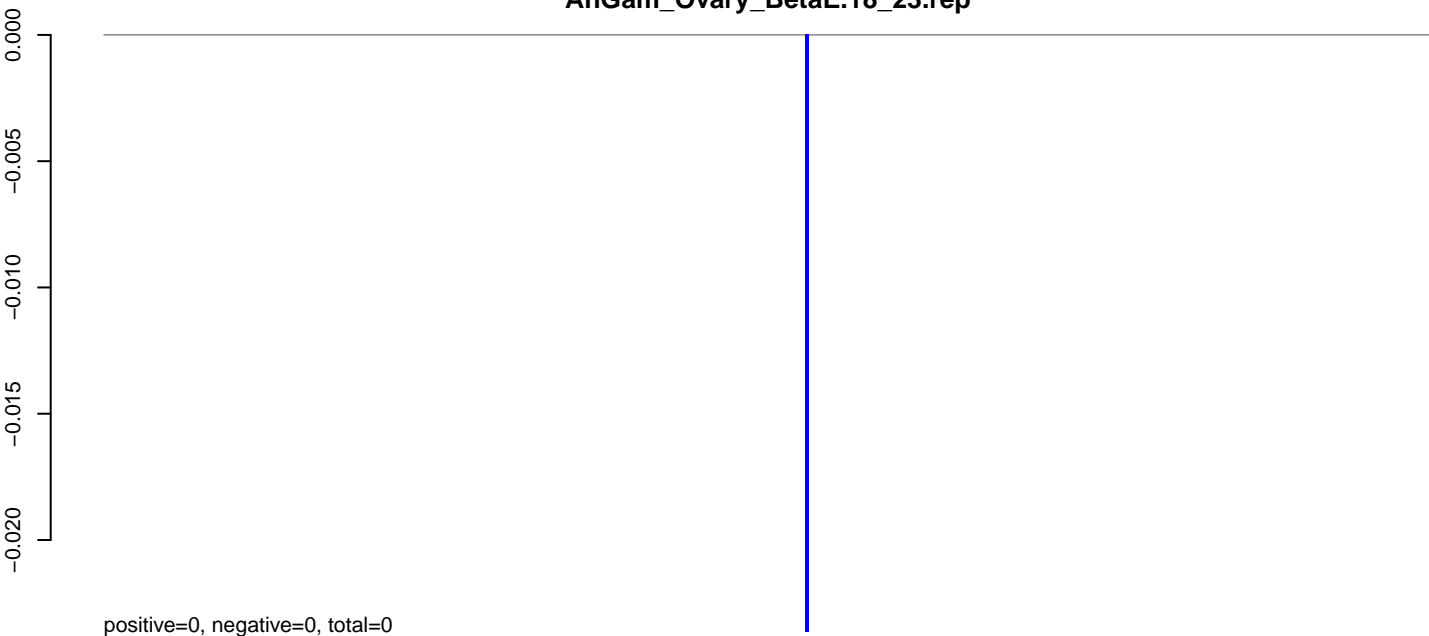


positive=0, negative=0, total=0

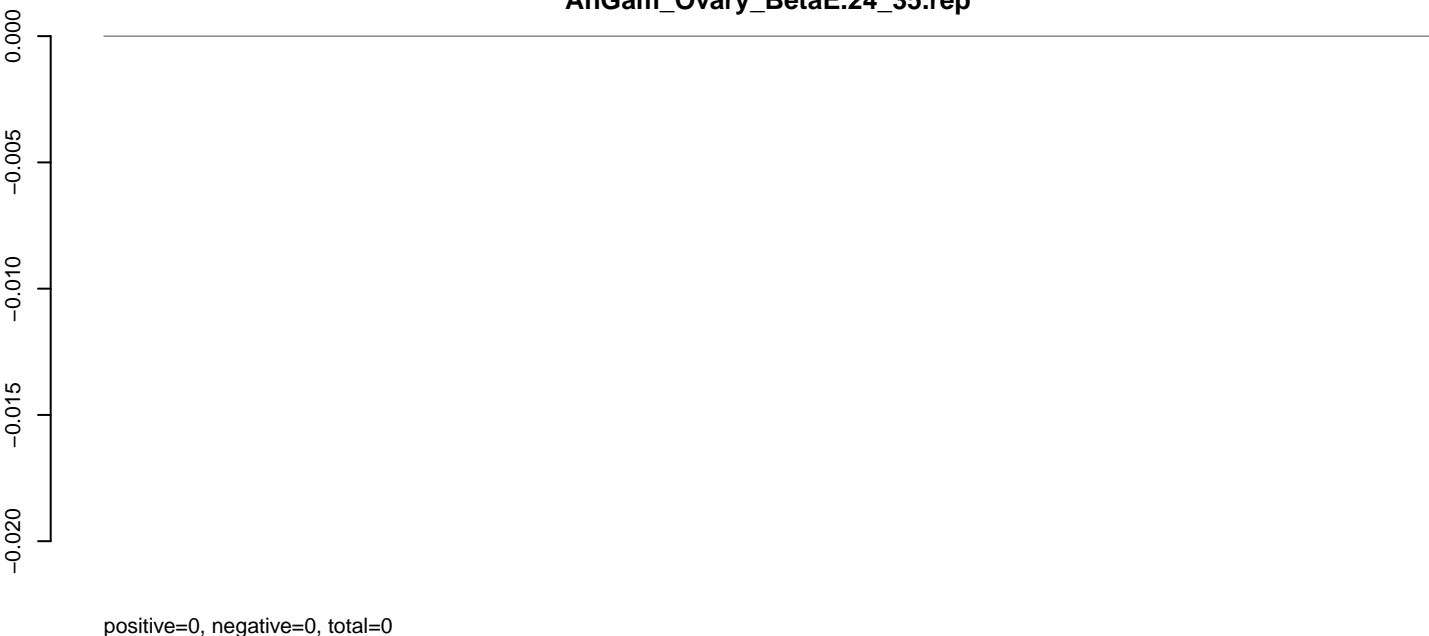
Window size=25, length=10116, virus@NC_003676.1-Apoi_virus_genome:1-10116



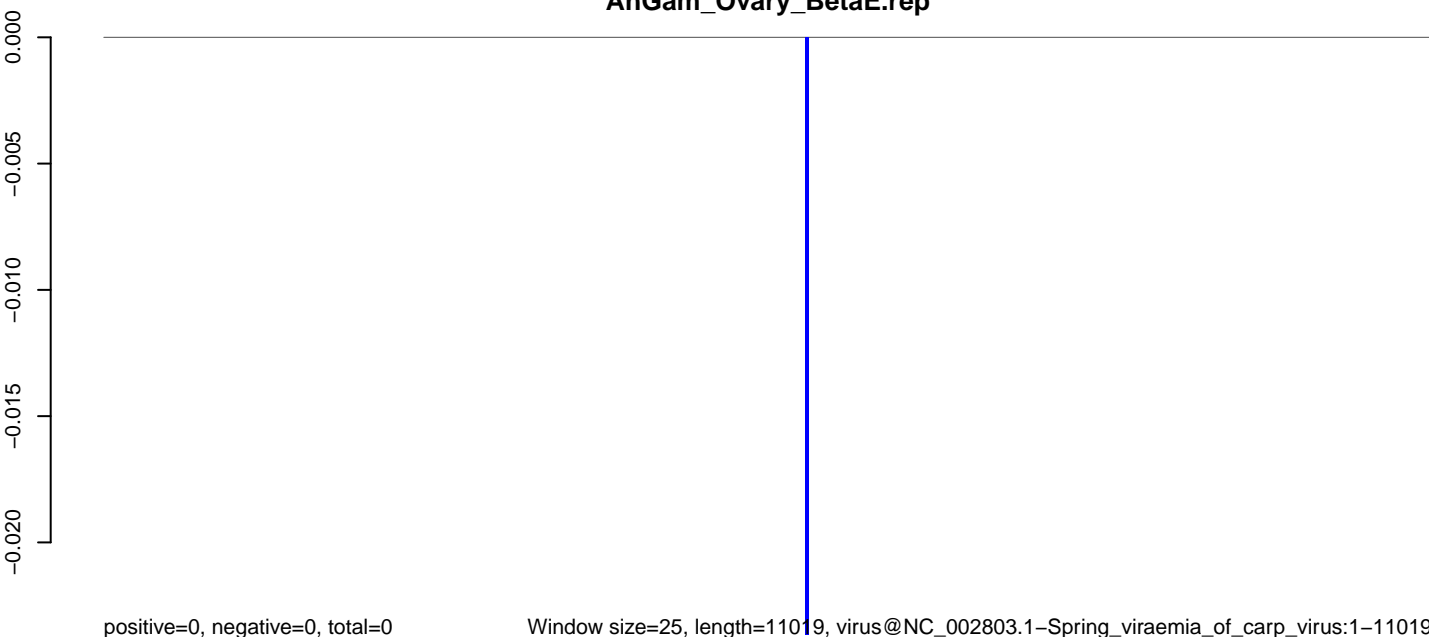
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



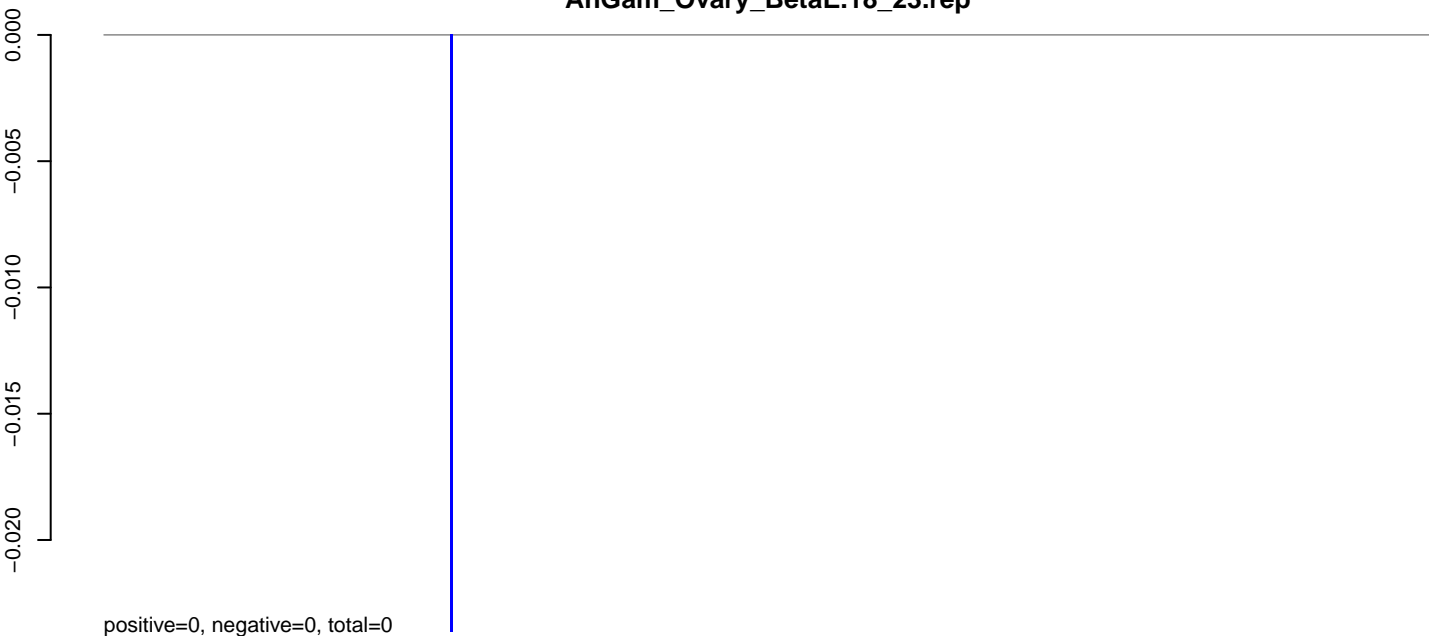
AnGam_Ovary_BetaE.rep



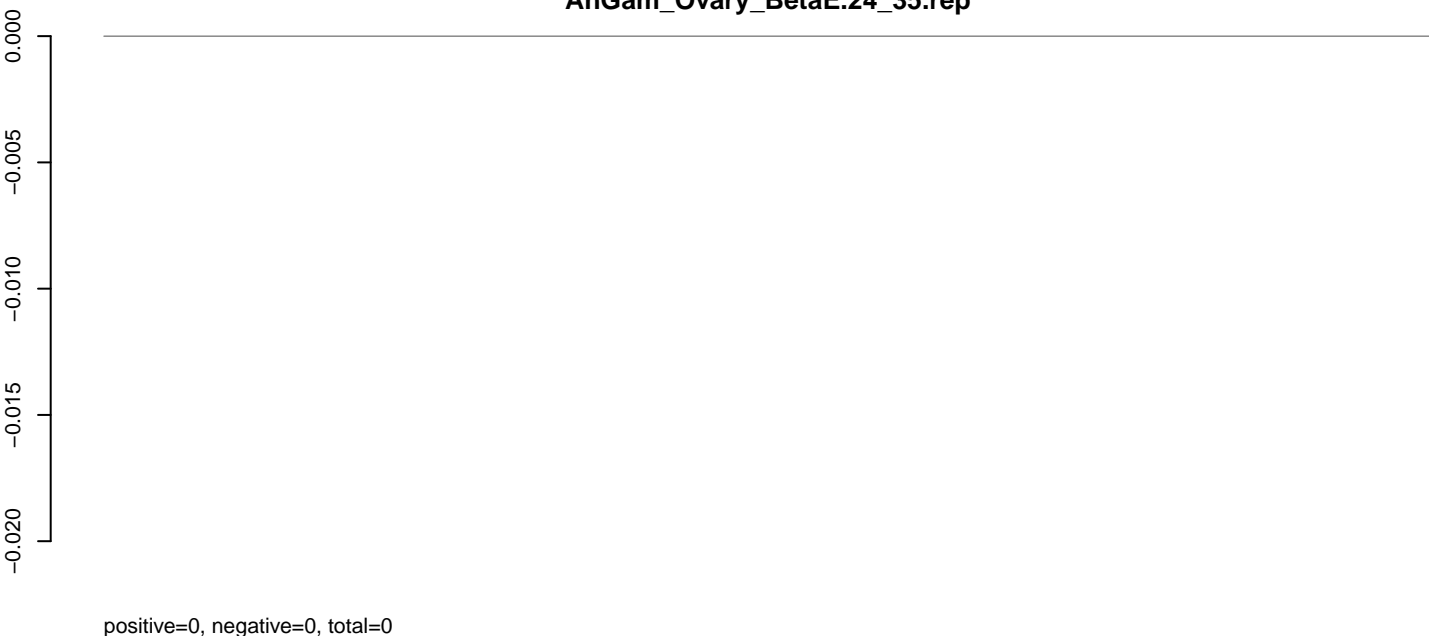
Window size=25, length=11019, virus@NC_002803.1-Spring_viraemia_of_carp_virus:1-11019

0 2000 4000 6000 8000 10000

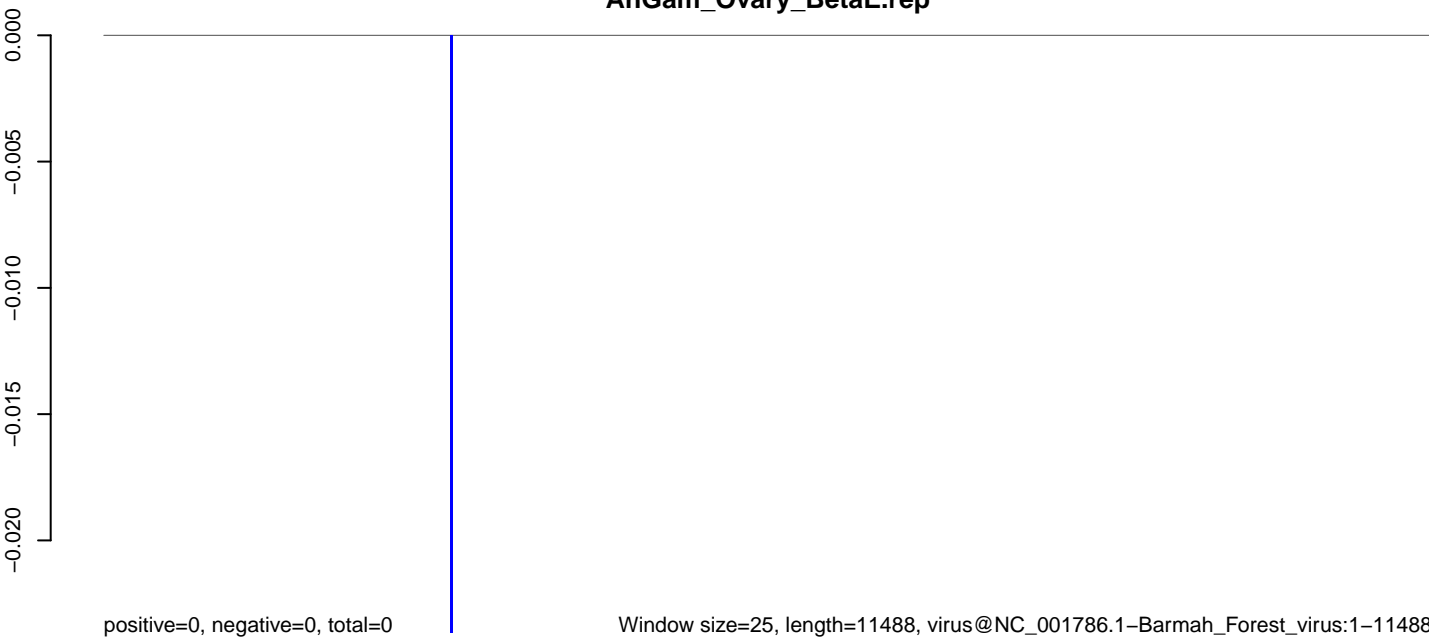
AnGam_Ovary_BetaE.18_23.rep



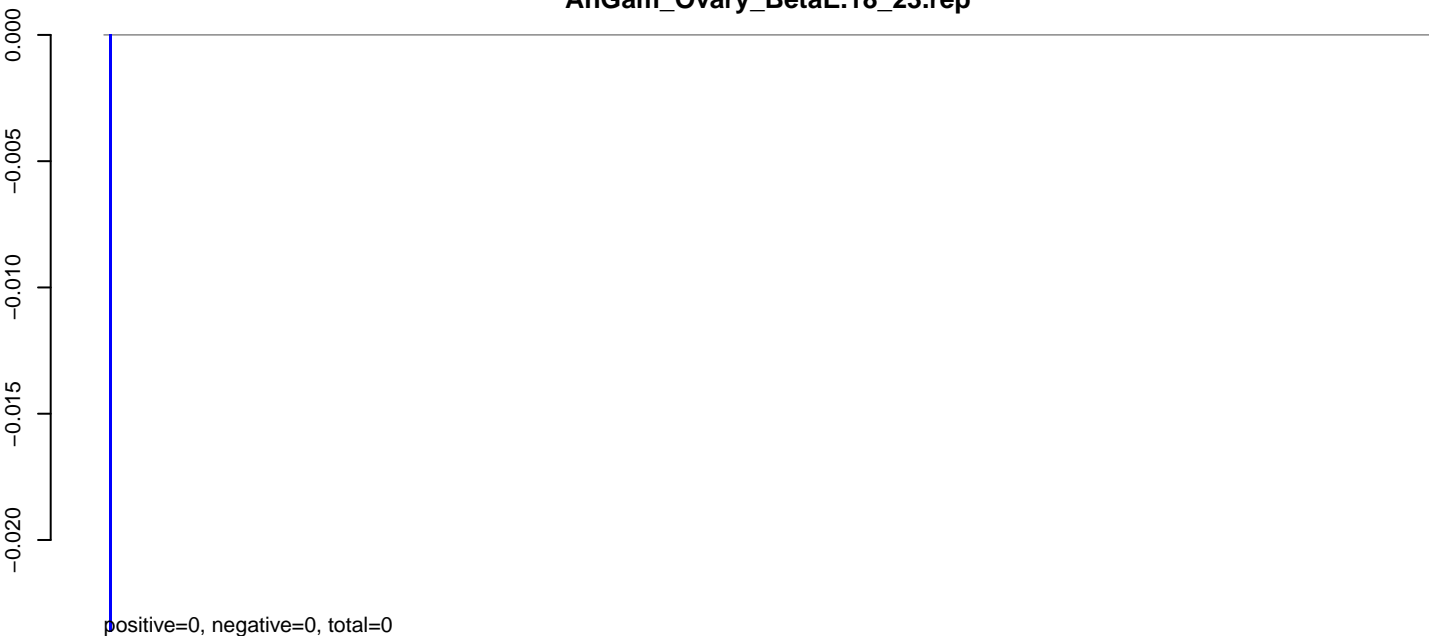
AnGam_Ovary_BetaE.24_35.rep



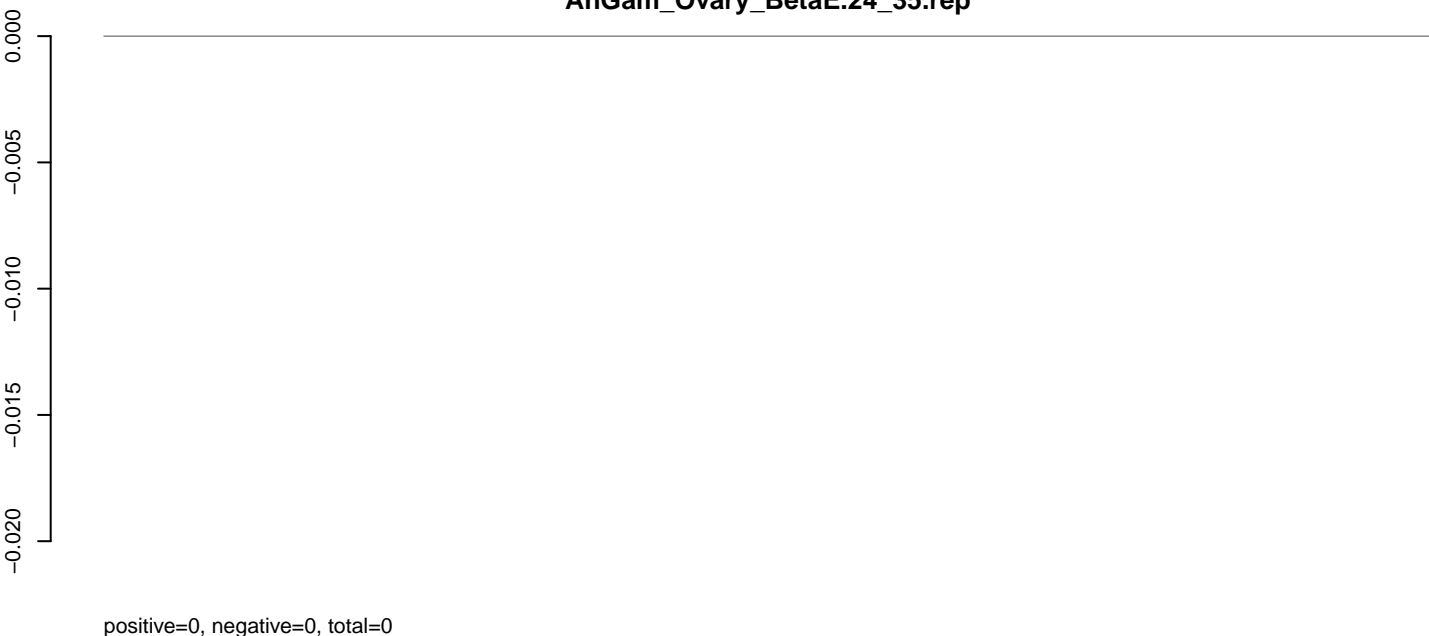
AnGam_Ovary_BetaE.rep



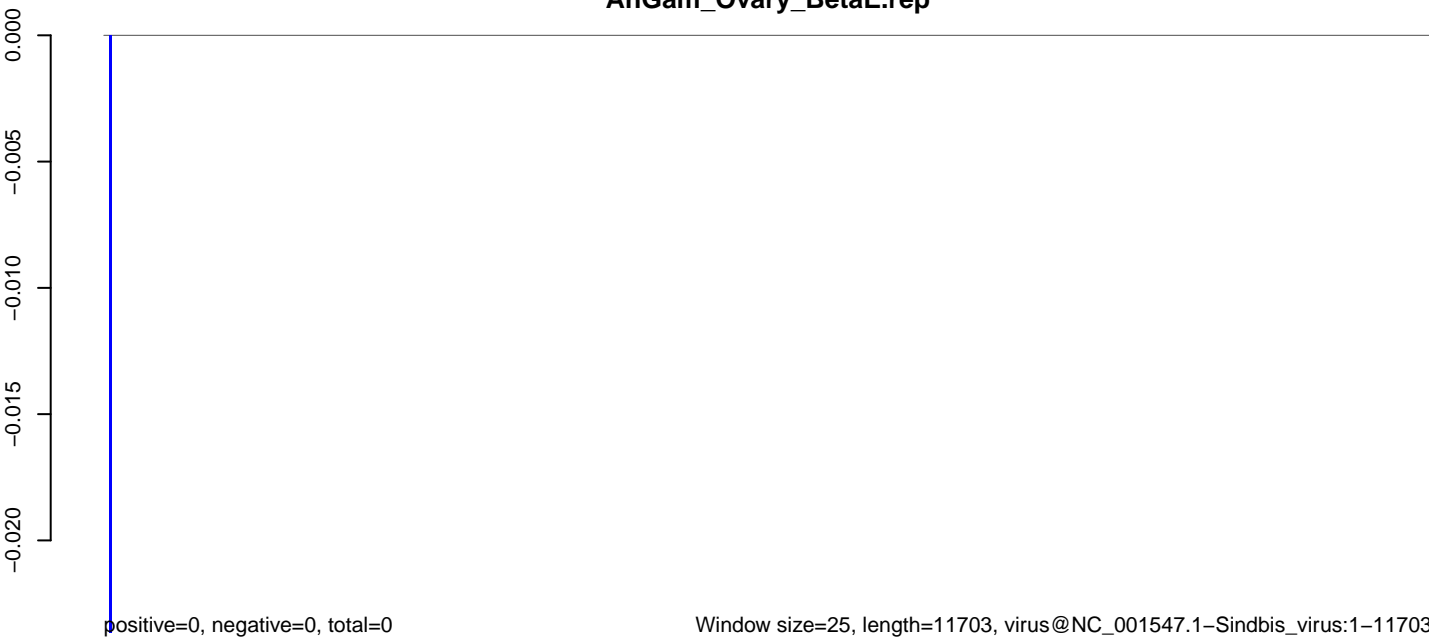
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



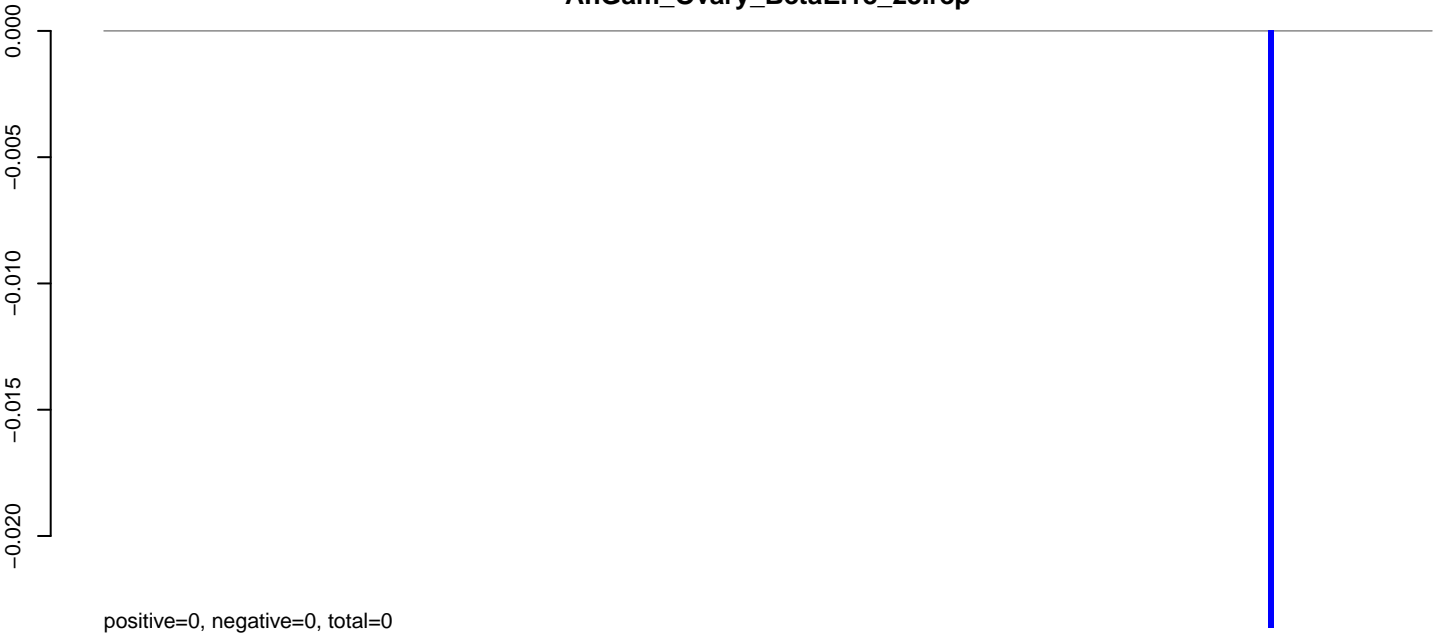
AnGam_Ovary_BetaE.rep



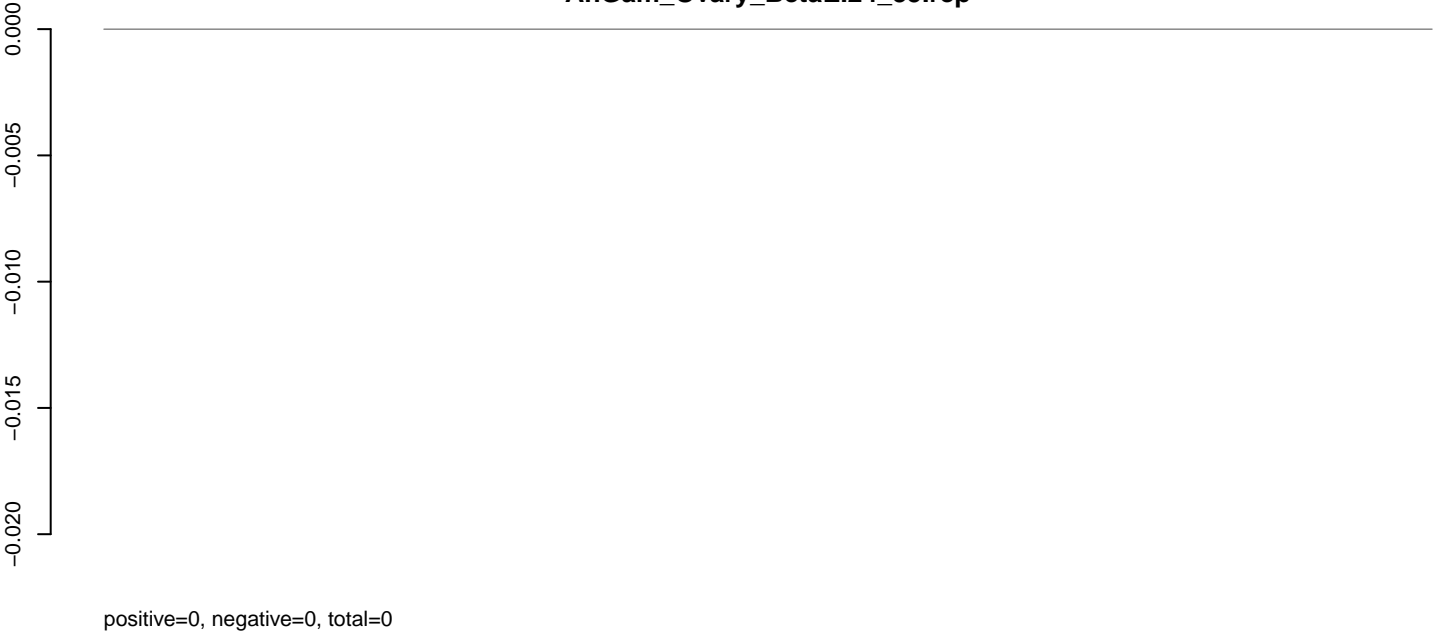
Window size=25, length=11703, virus@NC_001547.1-Sindbis_virus:1-11703

0 2000 4000 6000 8000 10000 12000

AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



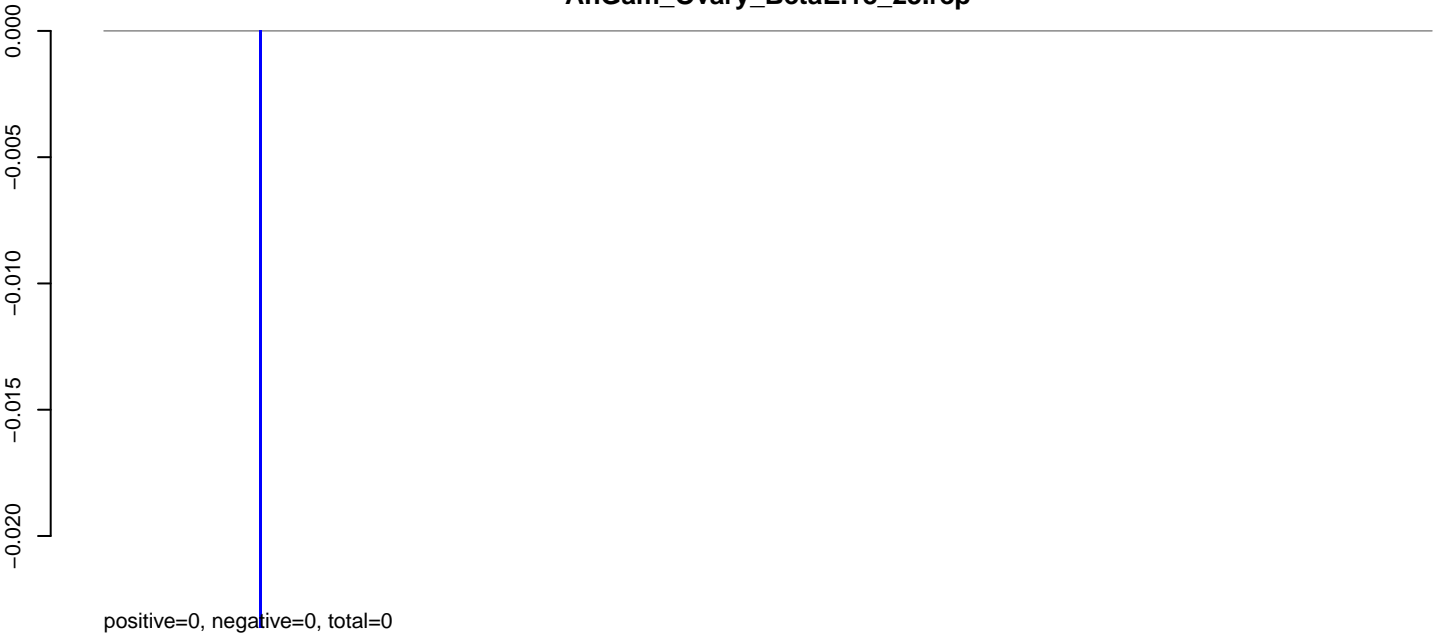
AnGam_Ovary_BetaE.rep



Window size=25, length=6065, virus@MH188054.1-Culex_Y_virus_strain_CYV_Kern_partial.genome:1-6065

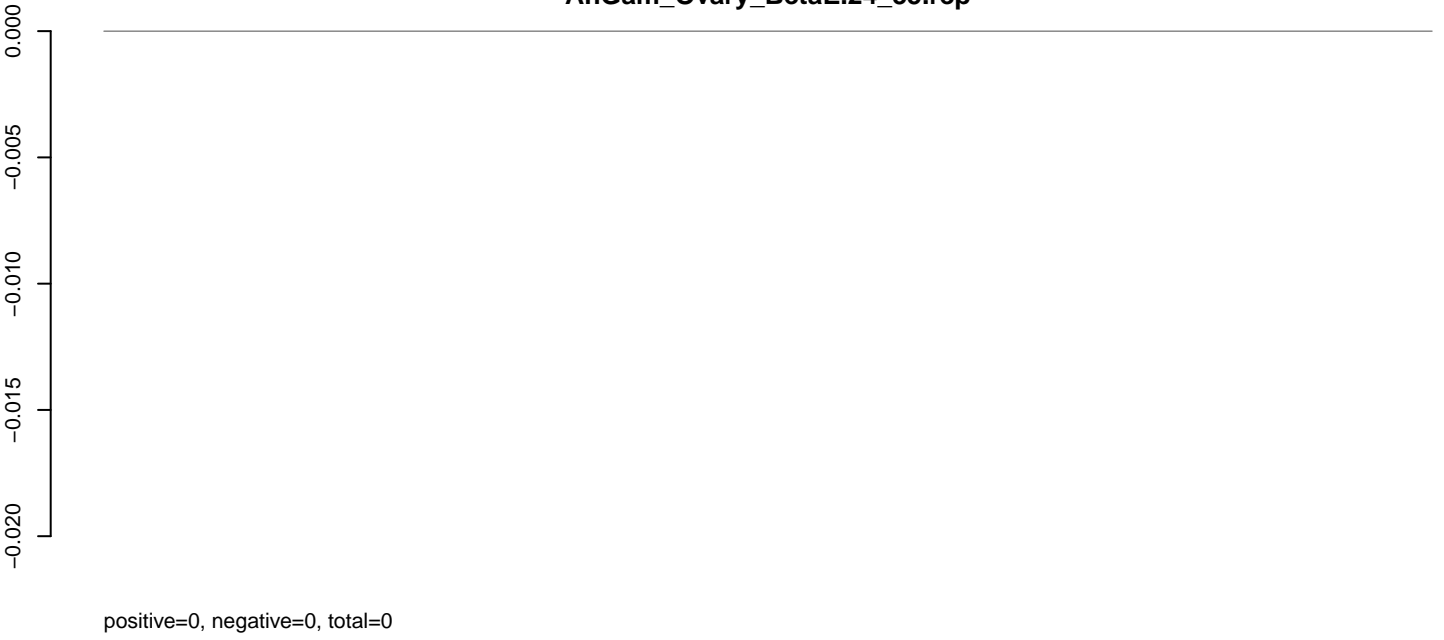
0 1000 2000 3000 4000 5000 6000

AnGam_Ovary_BetaE.18_23.rep



positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep



positive=0, negative=0, total=0

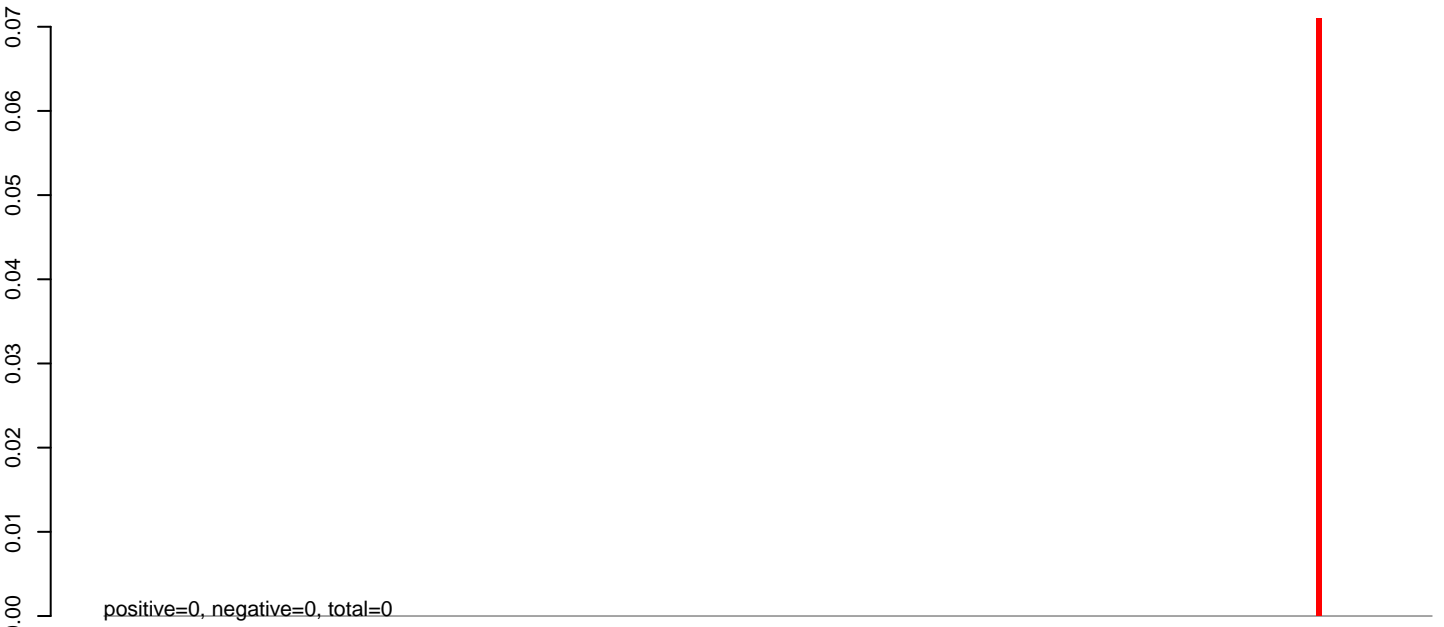
AnGam_Ovary_BetaE.rep



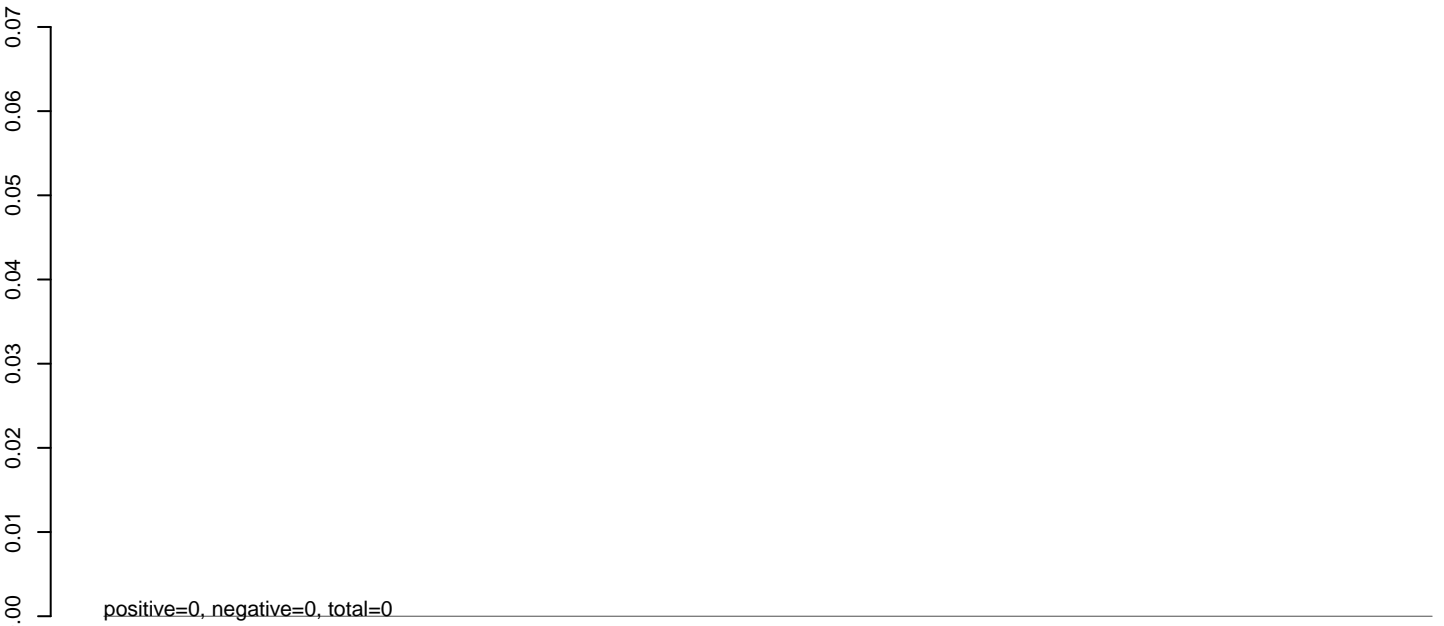
positive=0, negative=0, total=0, window size=25, length=12195, virus@MG747536.1_Oropouche_virus_isolate_BeAr_136921_segment_S_M_L_:1-12195

0 2000 4000 6000 8000 10000 12000

AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



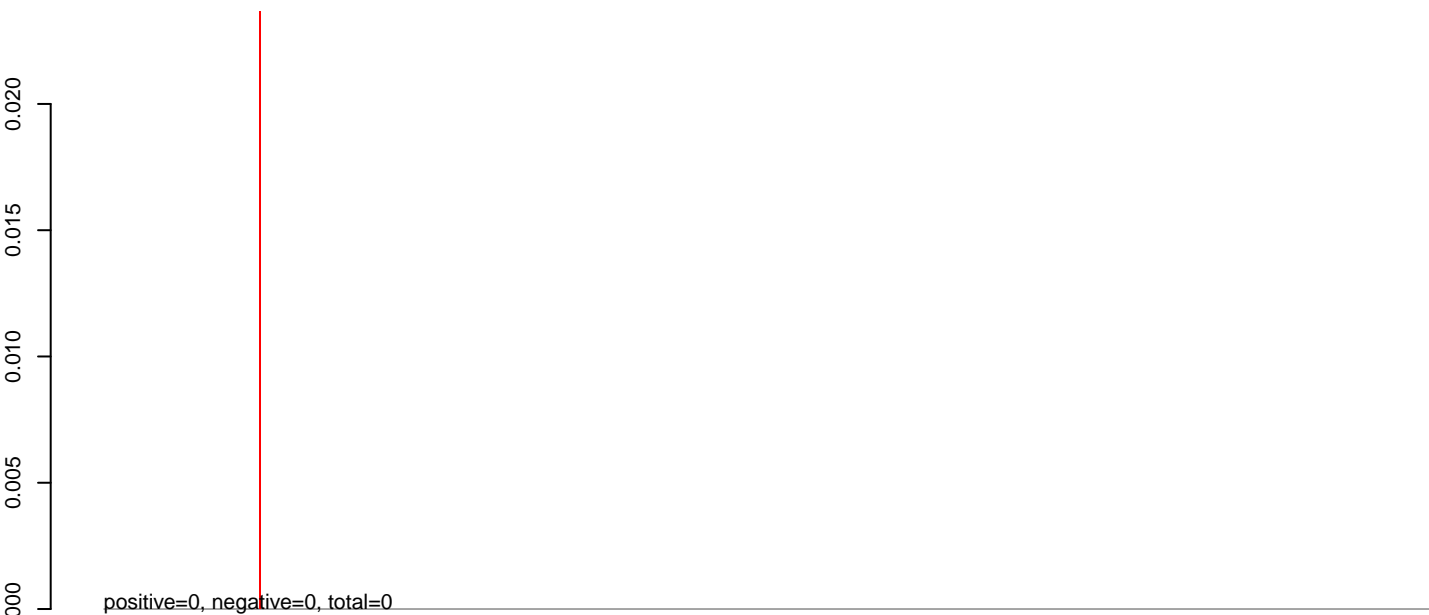
AnGam_Ovary_BetaE.rep



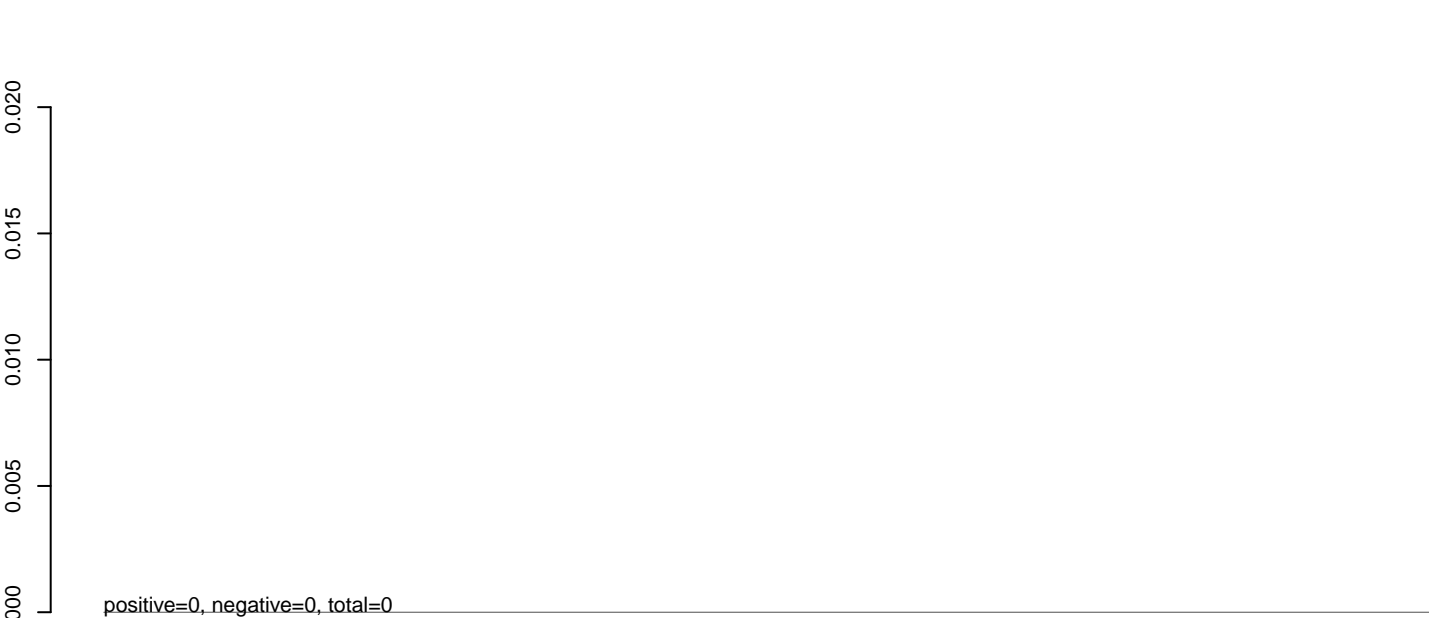
negative size=25, length=5982, virus@MF176317.1-Culex_mononega-like_virus_1_strain_mos191X45127_segment_2:1-5982

0 1000 2000 3000 4000 5000 6000

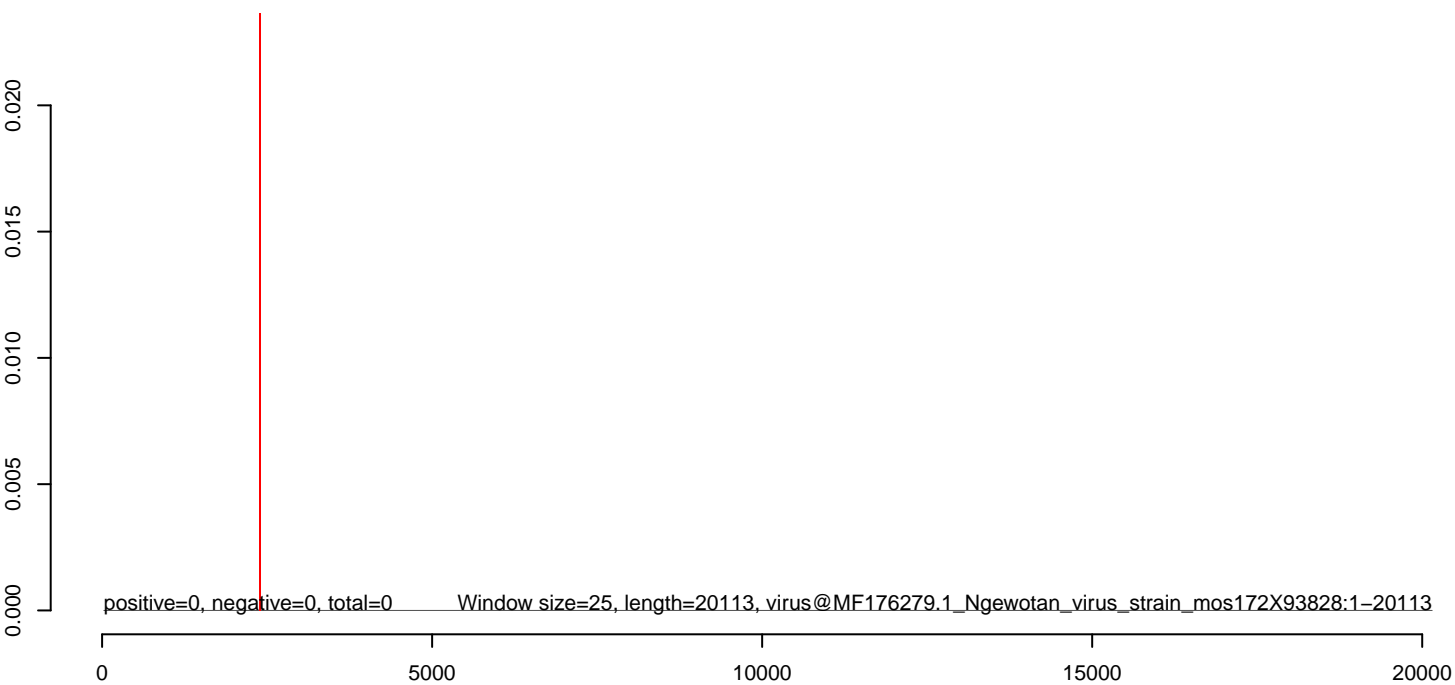
AnGam_Ovary_BetaE.18_23.rep



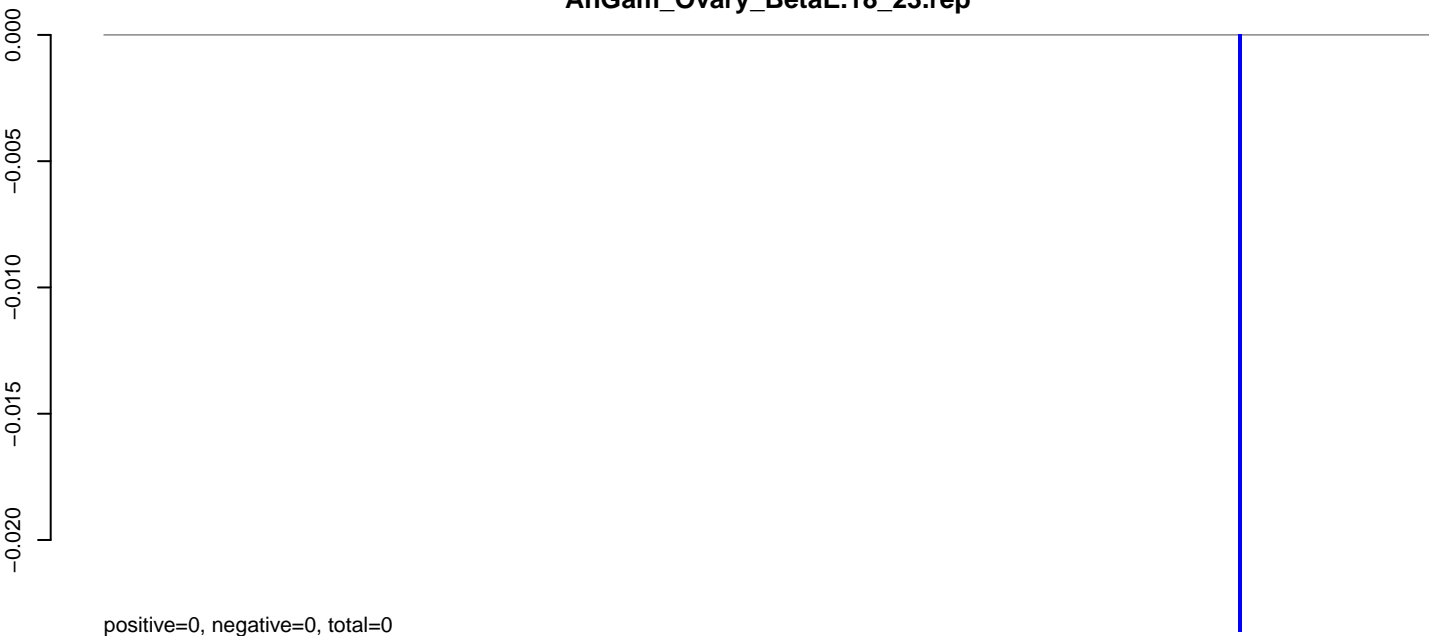
AnGam_Ovary_BetaE.24_35.rep



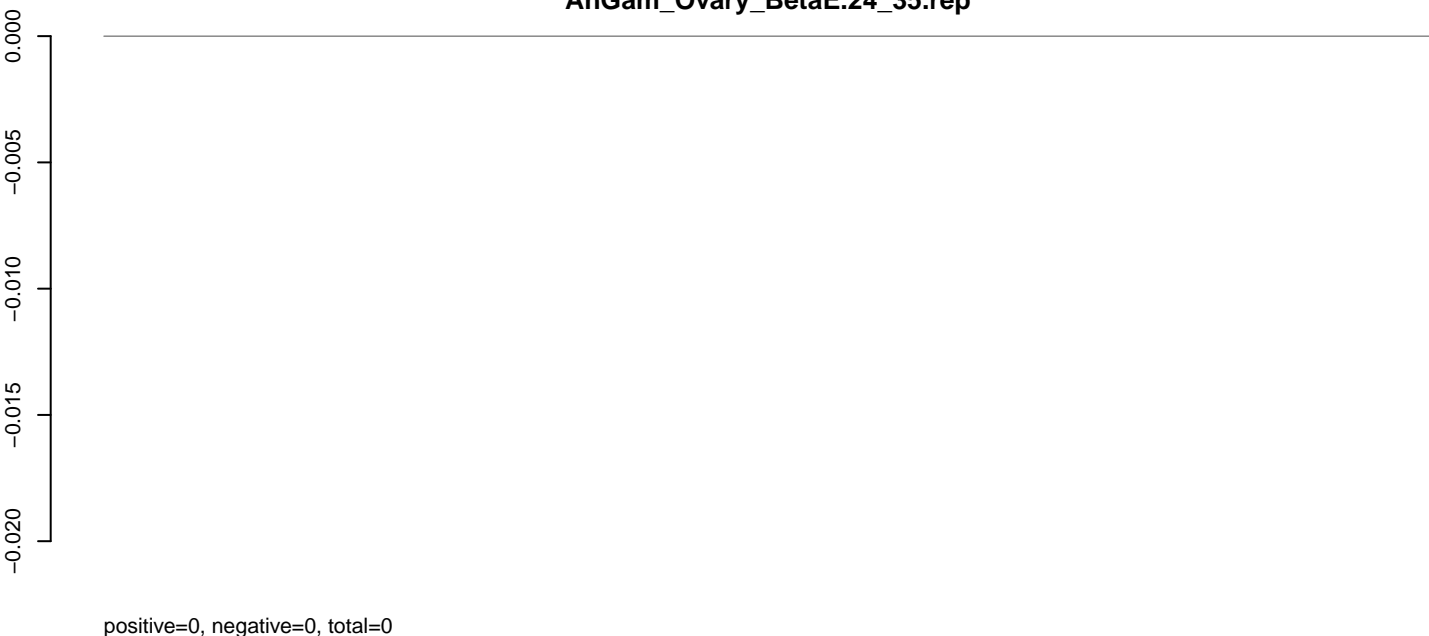
AnGam_Ovary_BetaE.rep



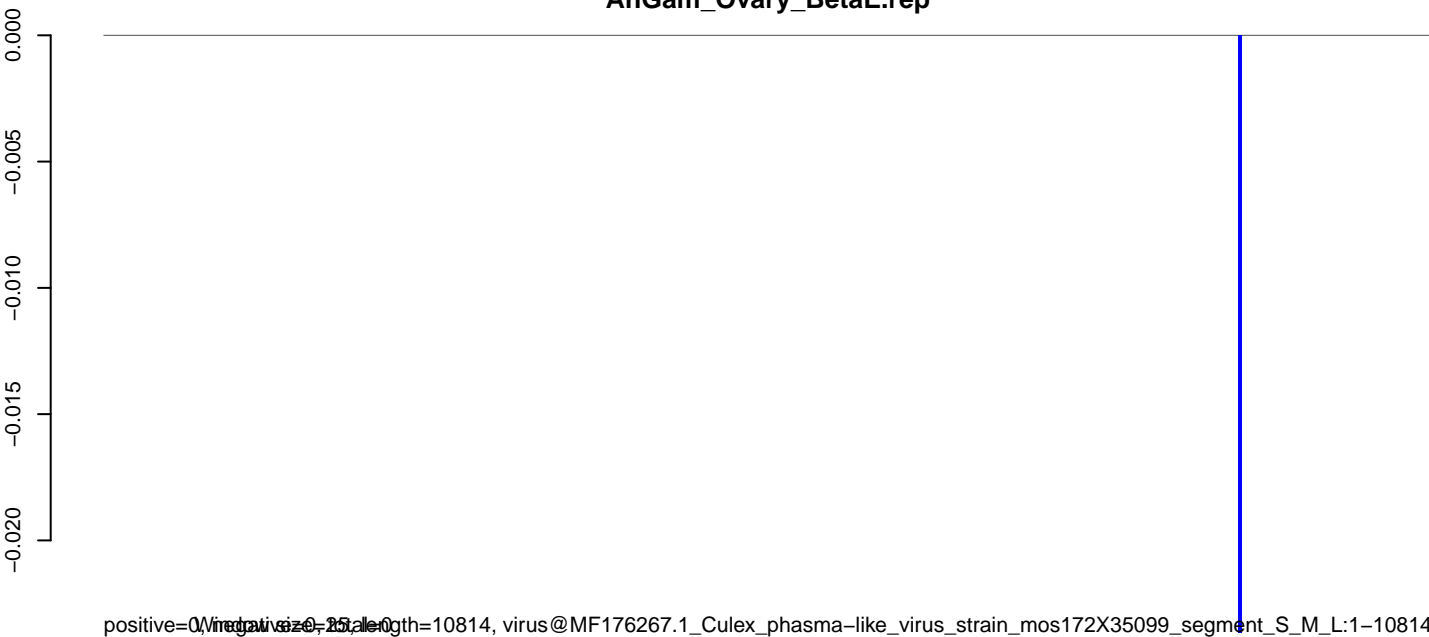
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



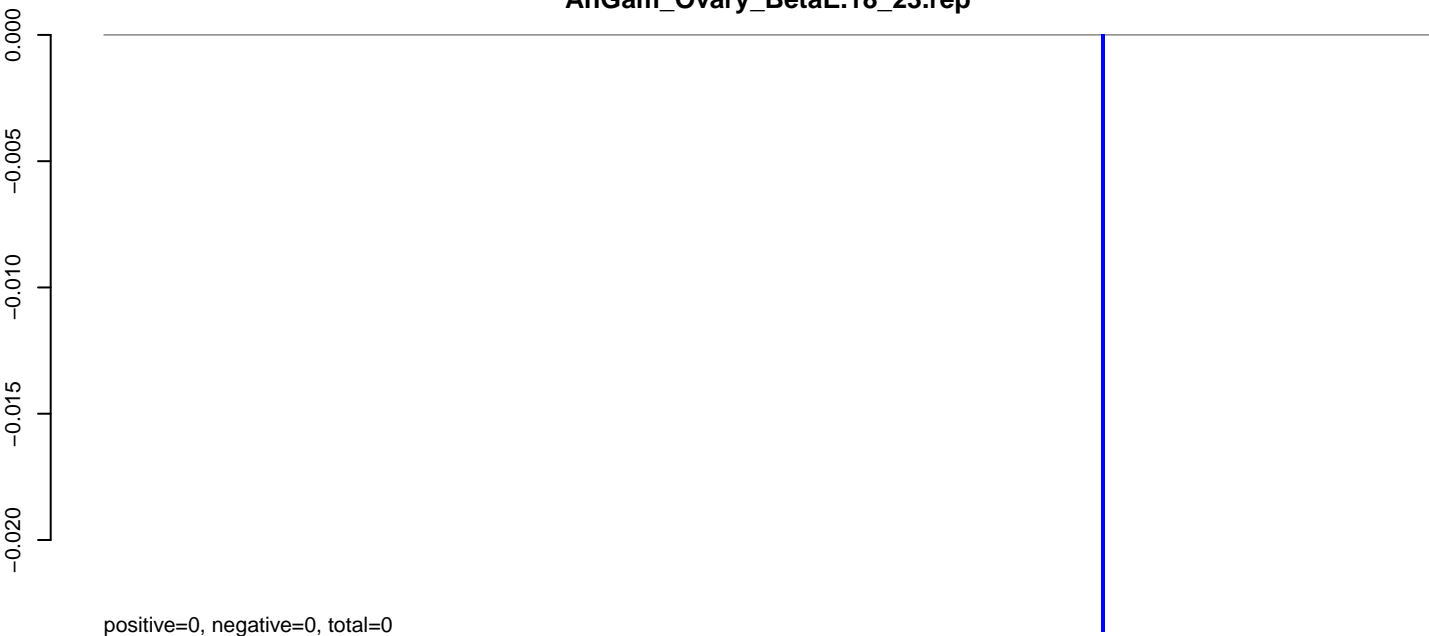
AnGam_Ovary_BetaE.rep



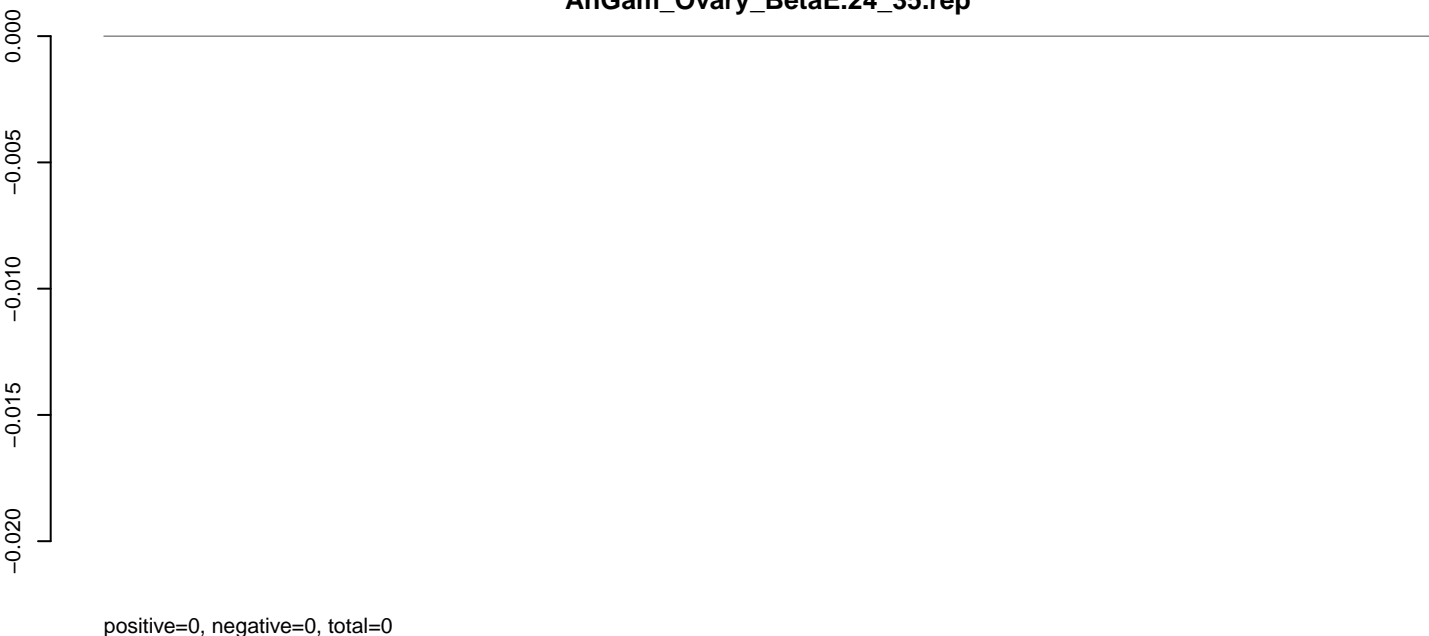
0 2000 4000 6000 8000 10000

positive=0, negative=0, total=0, virus@MF176267.1_Culex phasma-like virus_strain_mos172X35099_segment_S_M_L:1-10814

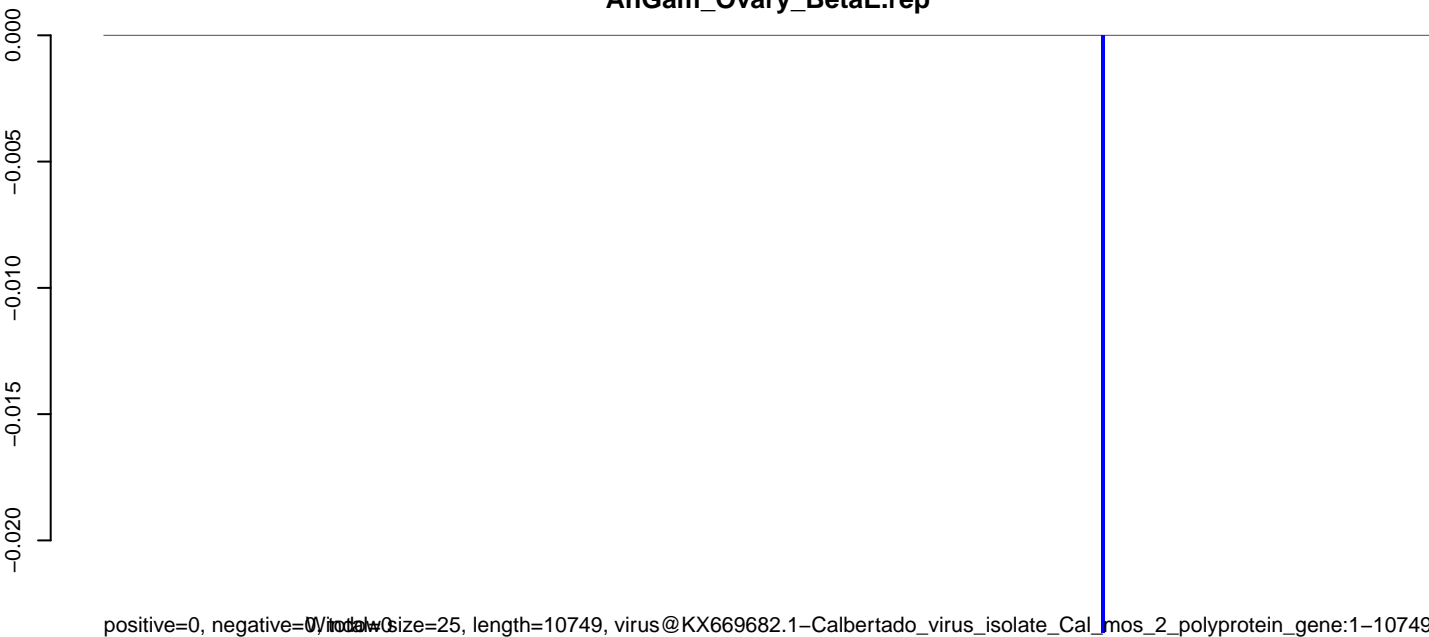
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



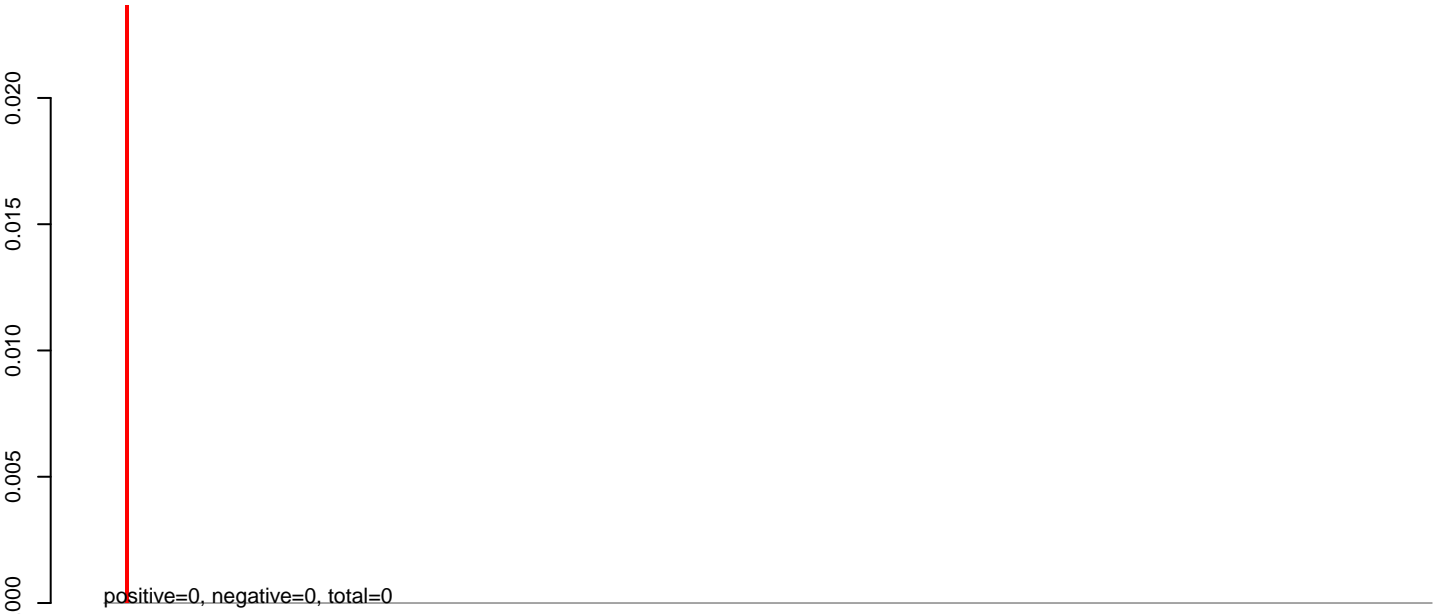
AnGam_Ovary_BetaE.rep



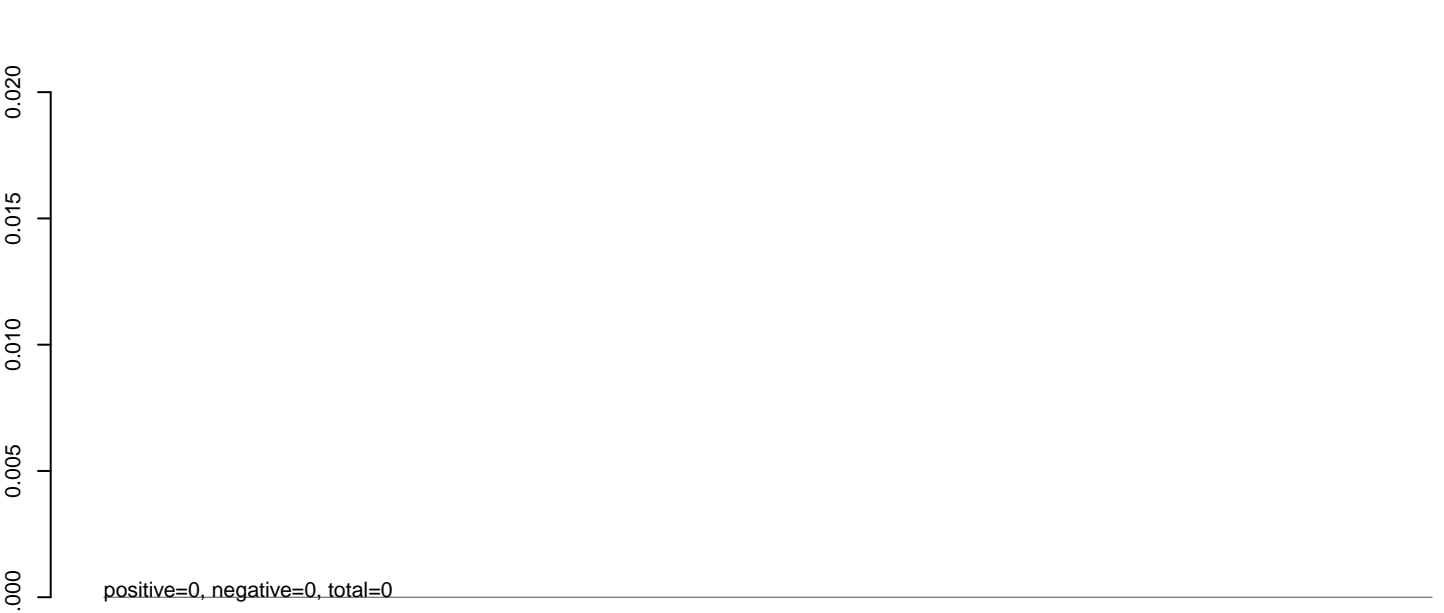
Window size=25, length=10749, virus@KX669682.1-Calbertado_virus_isolate_Cal_mos_2_polyprotein_gene:1-10749

0 2000 4000 6000 8000 10000

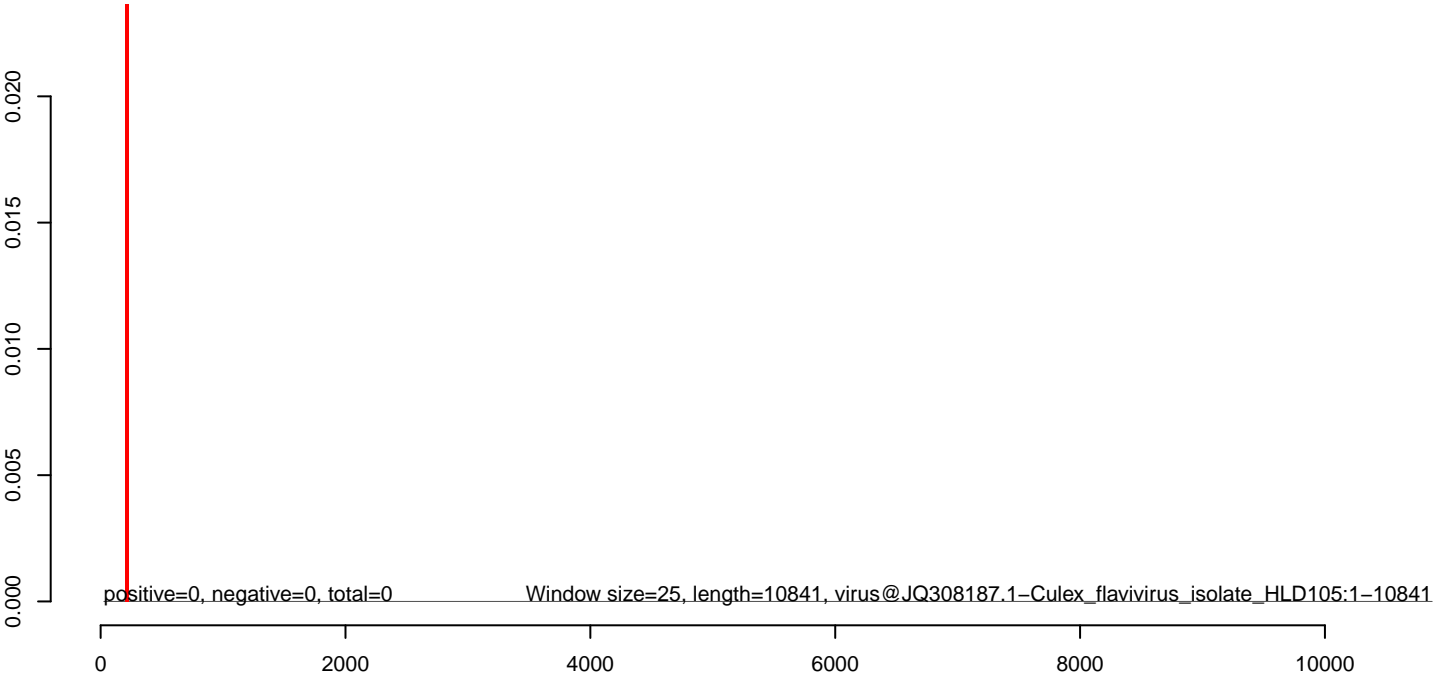
AnGam_Ovary_BetaE.18_23.rep



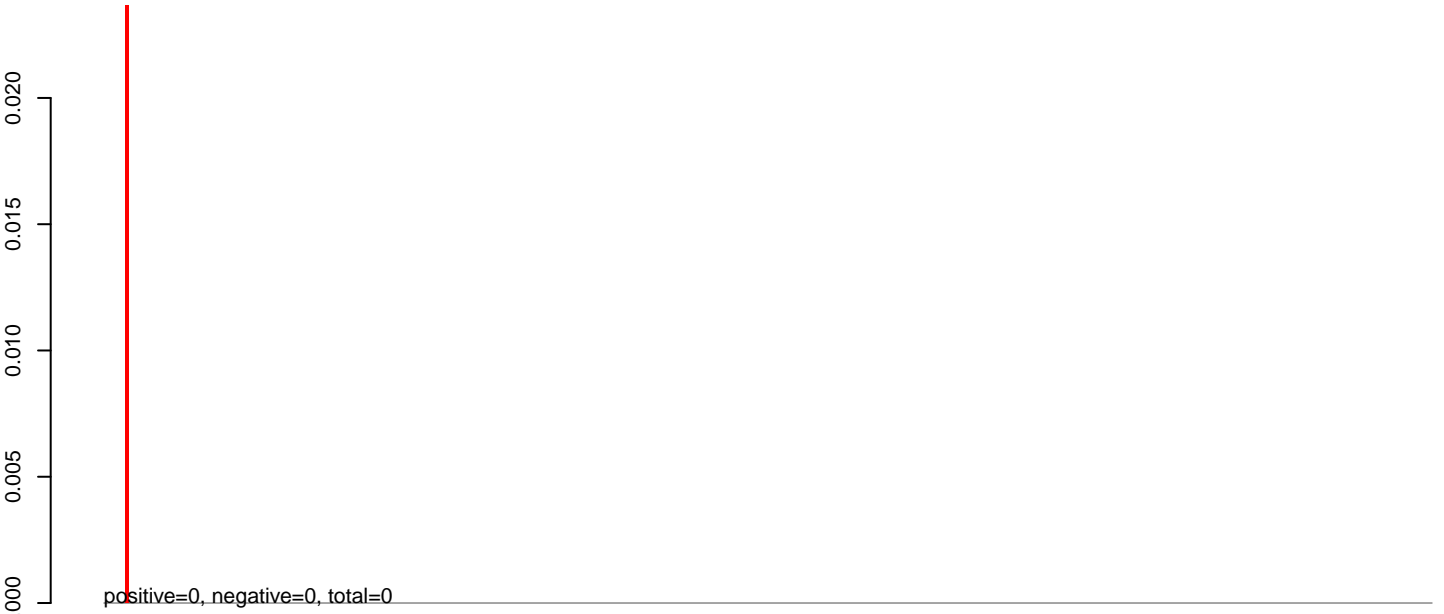
AnGam_Ovary_BetaE.24_35.rep



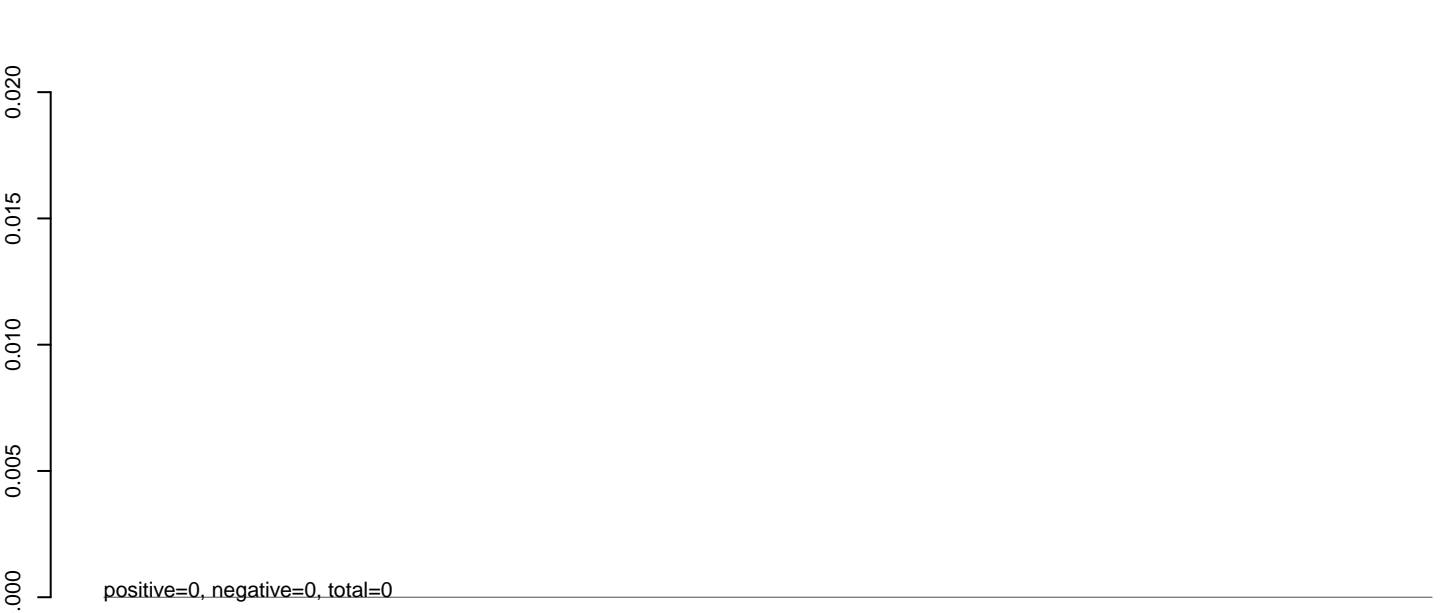
AnGam_Ovary_BetaE.rep



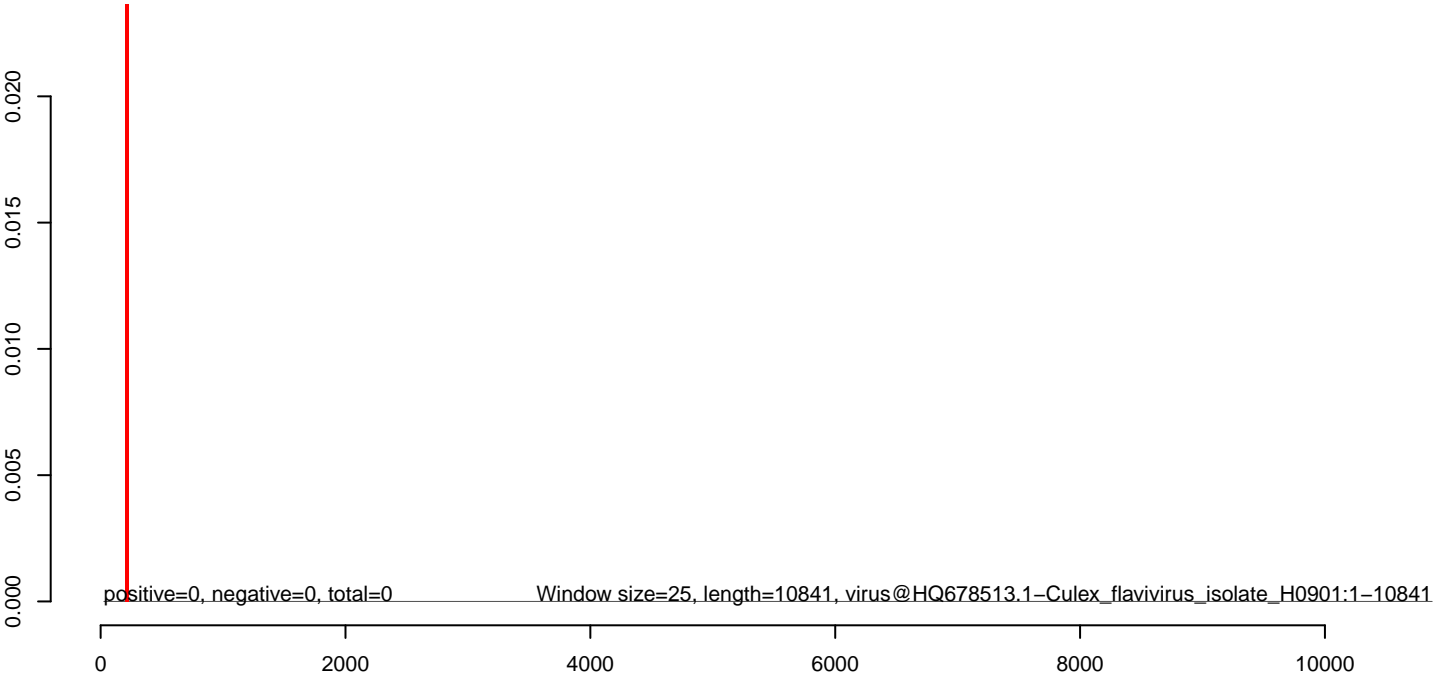
AnGam_Ovary_BetaE.18_23.rep



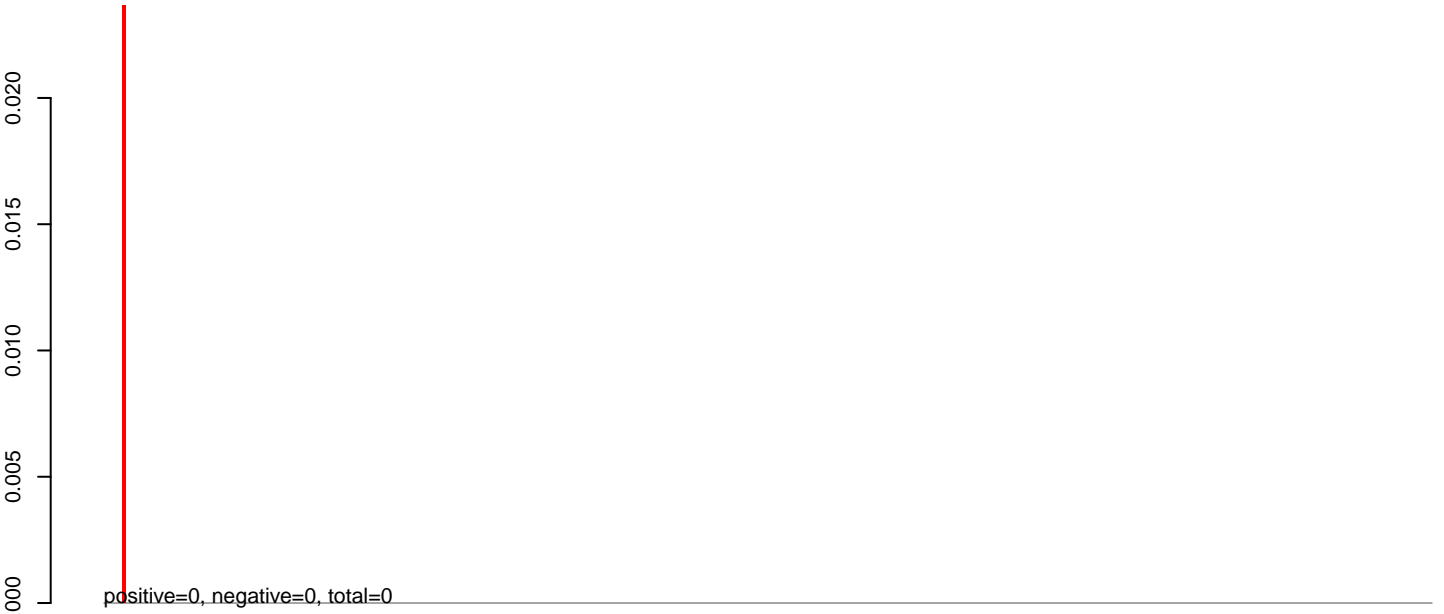
AnGam_Ovary_BetaE.24_35.rep



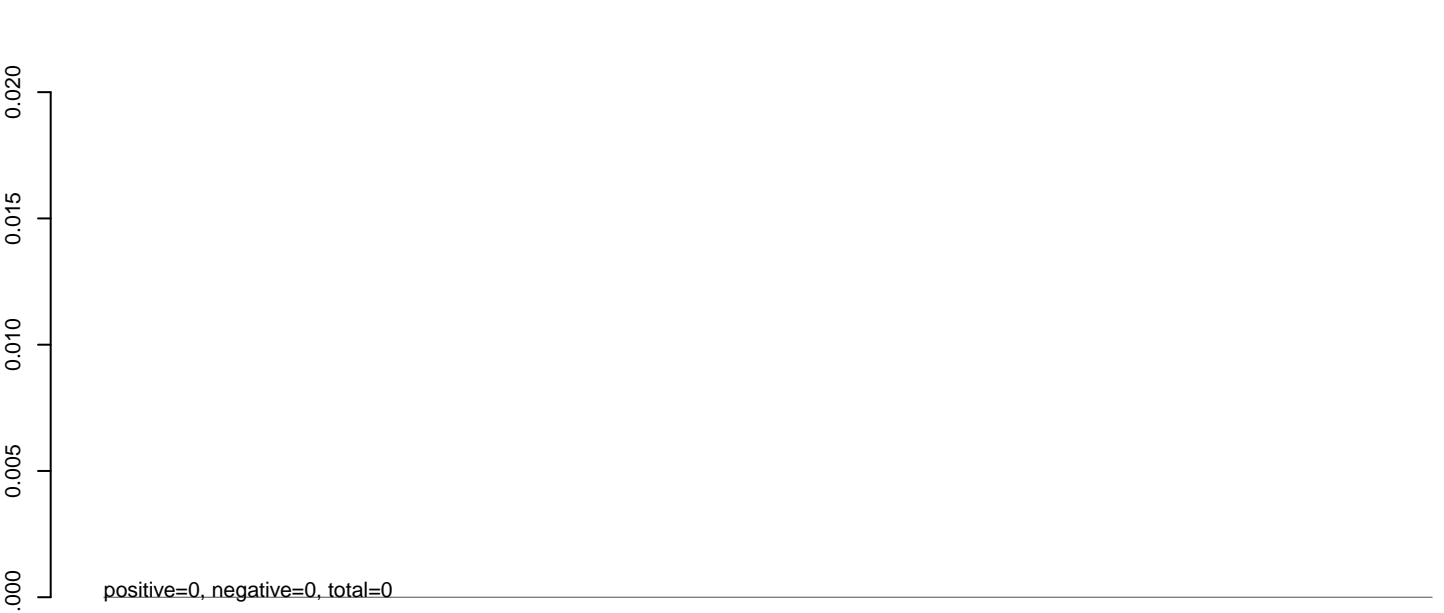
AnGam_Ovary_BetaE.rep



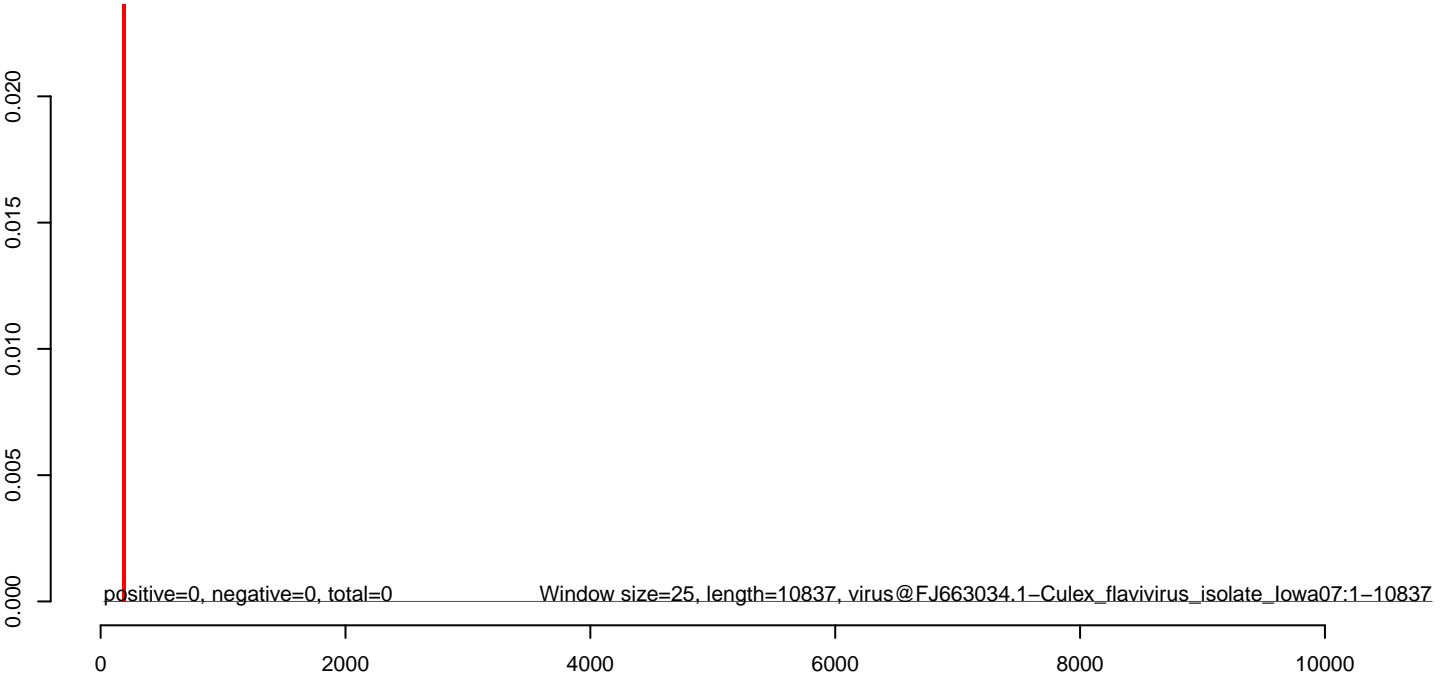
AnGam_Ovary_BetaE.18_23.rep



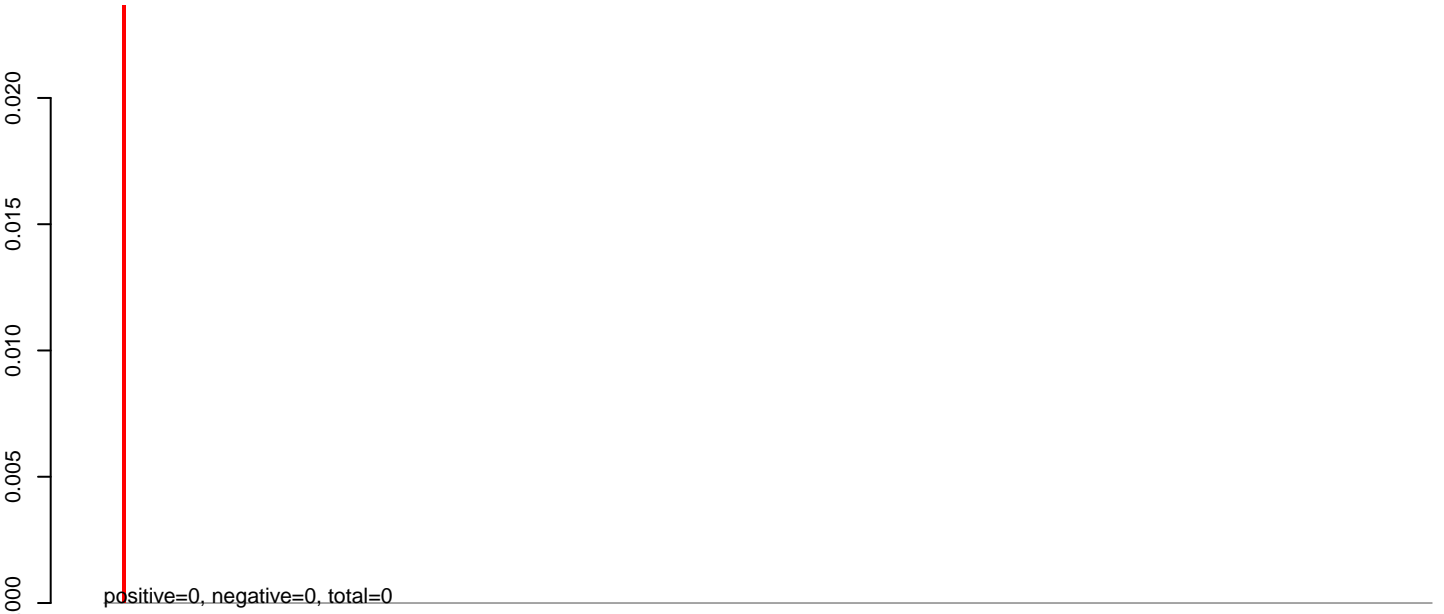
AnGam_Ovary_BetaE.24_35.rep



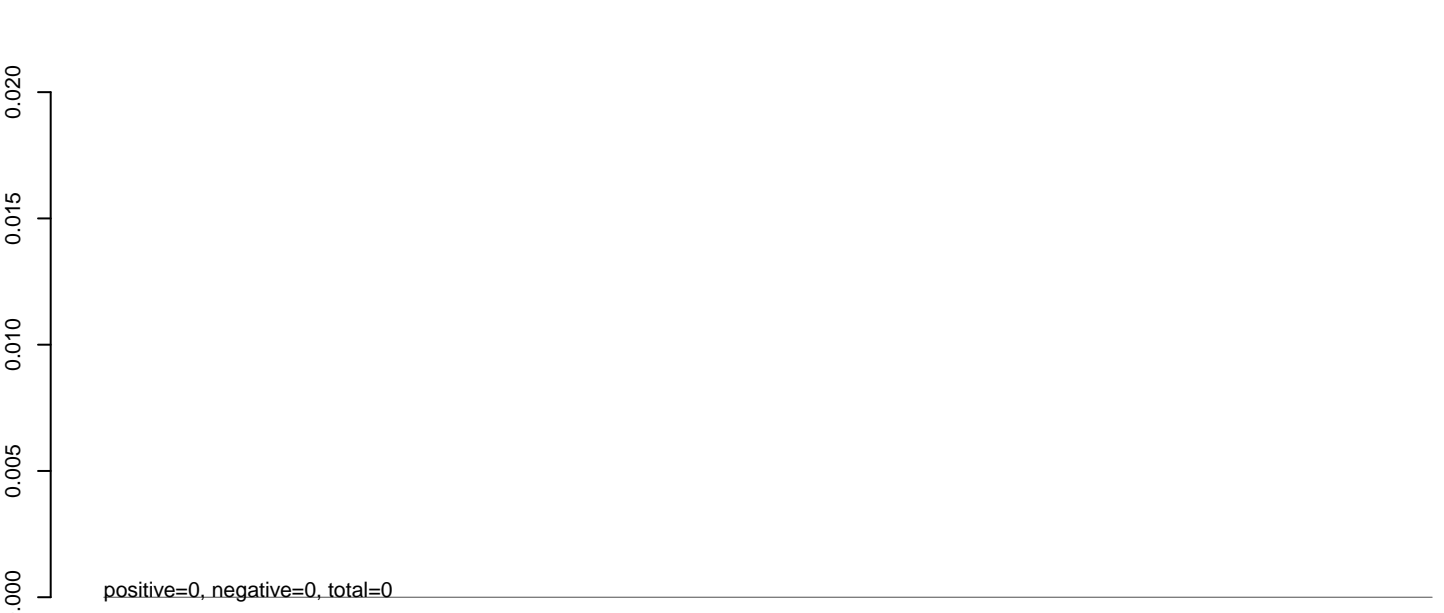
AnGam_Ovary_BetaE.rep



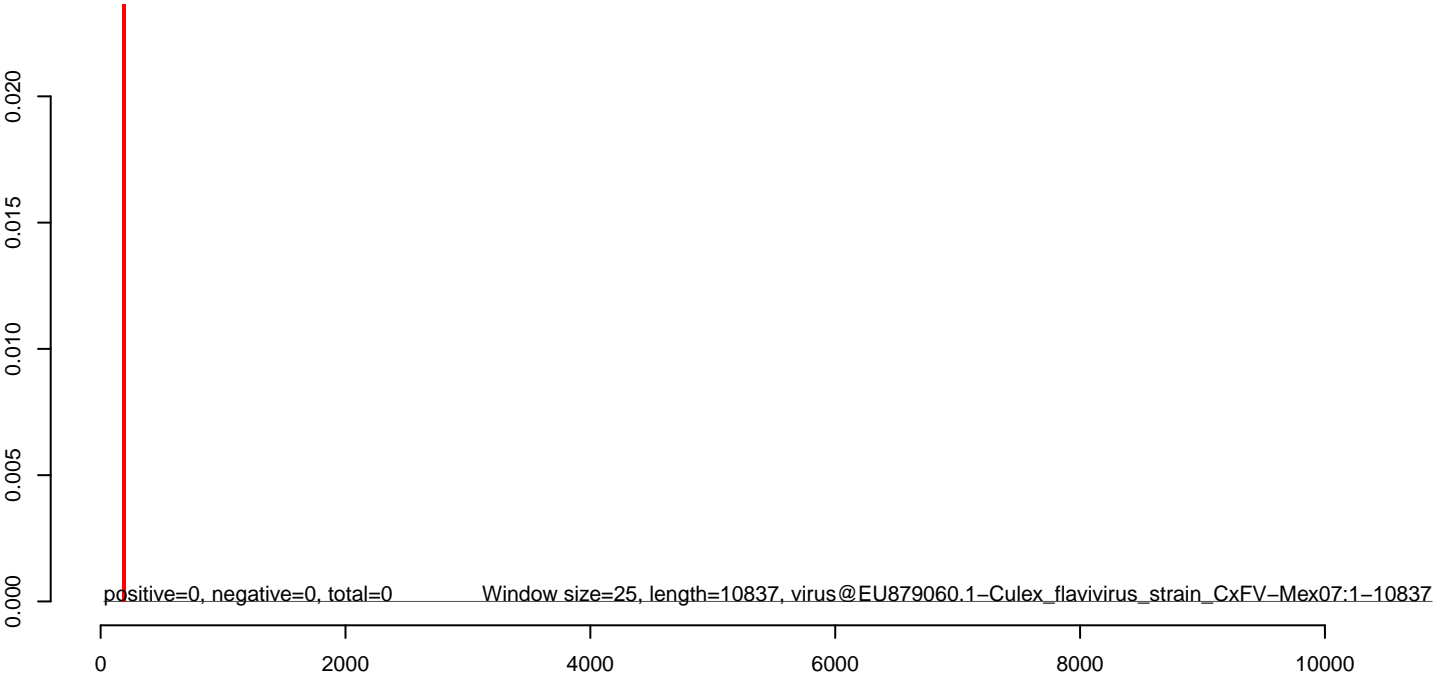
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



AnGam_Ovary_BetaE.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep

0.000
-0.005
-0.010
-0.015
-0.020

with size 251, length 3986, cv=0, score=0.005055.1-Haemagogus_equinus_densovirus_nonstructural_protein_1_nonstructural_protein_2_and_c:1-3986

0 1000 2000 3000 4000

AnGam_Ovary_BetaE.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

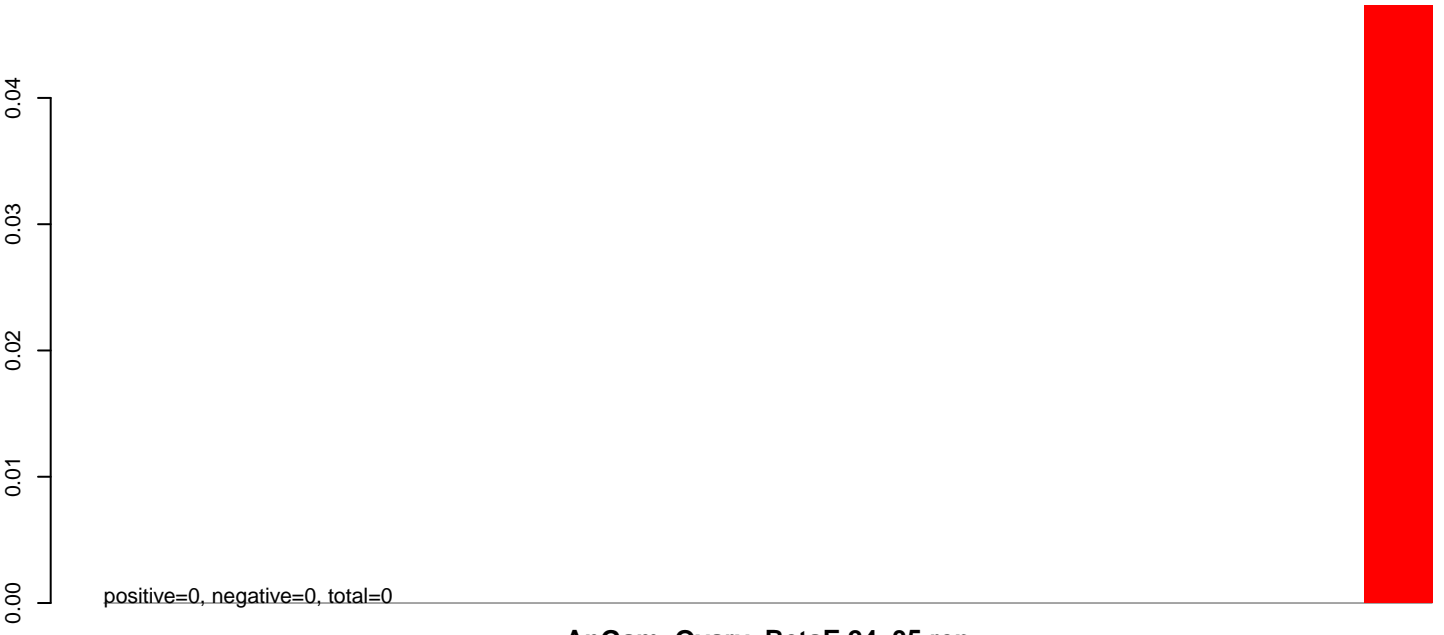
AnGam_Ovary_BetaE.rep

0.000
-0.005
-0.010
-0.015
-0.020

downloaded from https://www.ncbi.nlm.nih.gov/trace/view_alignment.html?accession=AY310877.1 - Aedes albopictus_C6_36_cell_densovirus_non-structural_protein_1_non-structural_prot:1-4006

0 1000 2000 3000 4000

AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



AnGam_Ovary_BetaE.18_23.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.24_35.rep

0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

AnGam_Ovary_BetaE.rep

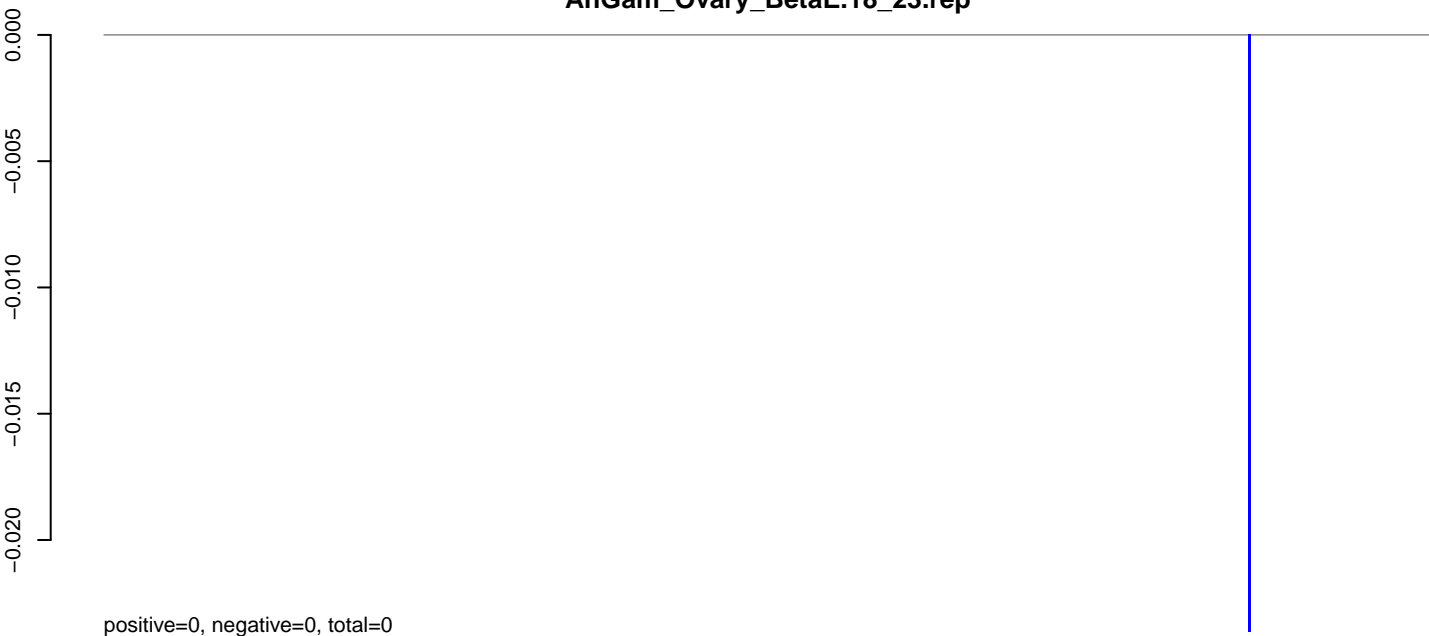
0.000
-0.005
-0.010
-0.015
-0.020

positive=0, negative=0, total=0

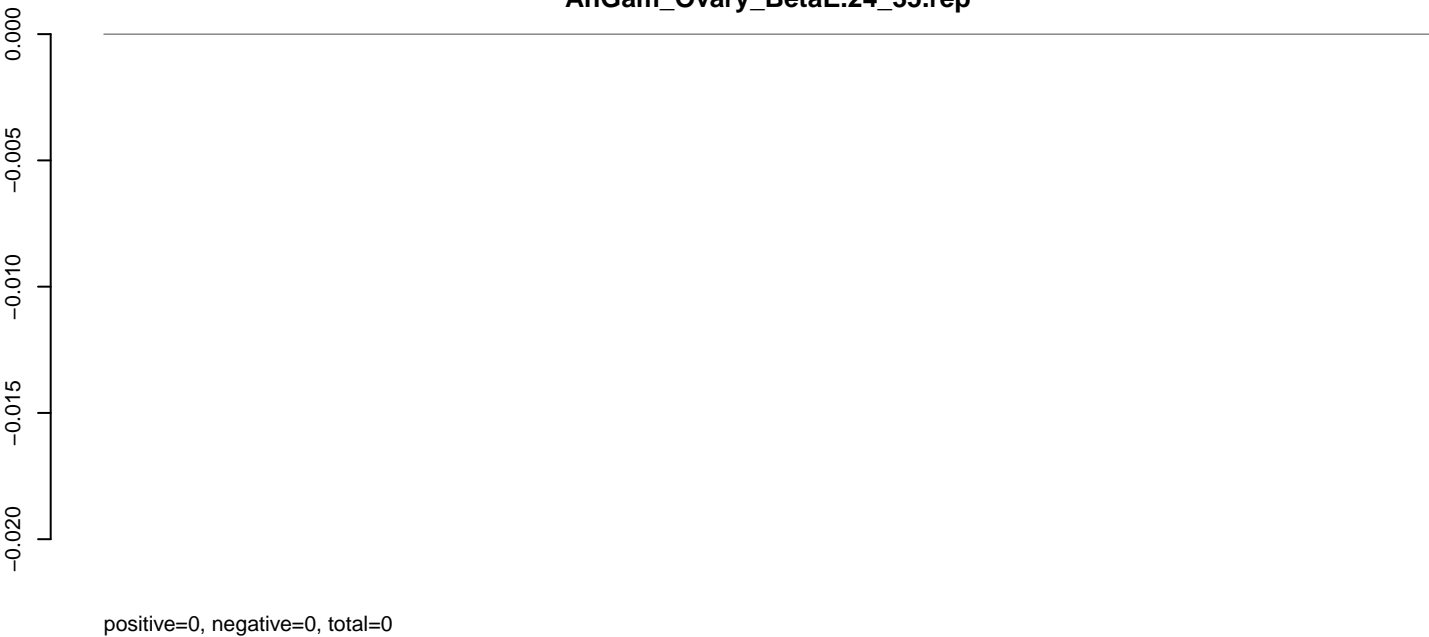
Window size=25, length=11344, virus@AF075256.1-Pixuna_virus_strain_Pixuna_BeAr_35645:1-11344

0 2000 4000 6000 8000 10000

AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



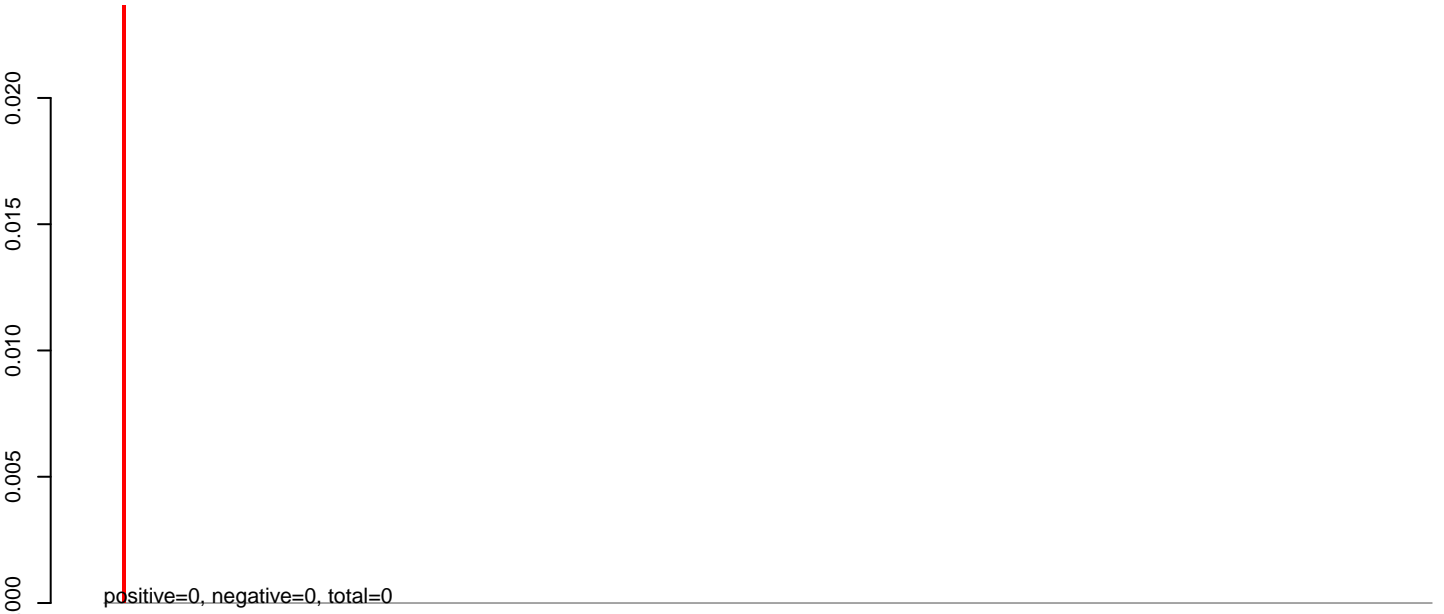
AnGam_Ovary_BetaE.rep



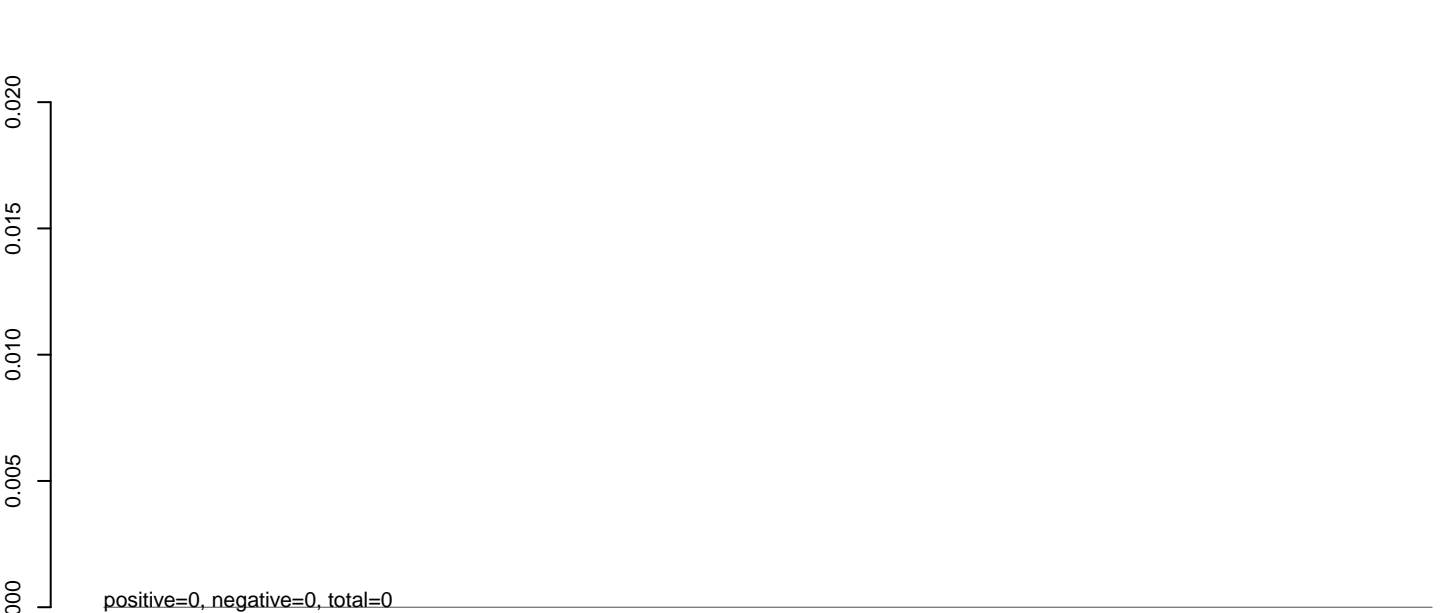
Window size=25, length=11530, virus@AF075254.1-Tonate_virus_strain_Tonate_CaAn_410d:1-11530

0 2000 4000 6000 8000 10000 12000

AnGam_Ovary_BetaE.18_23.rep



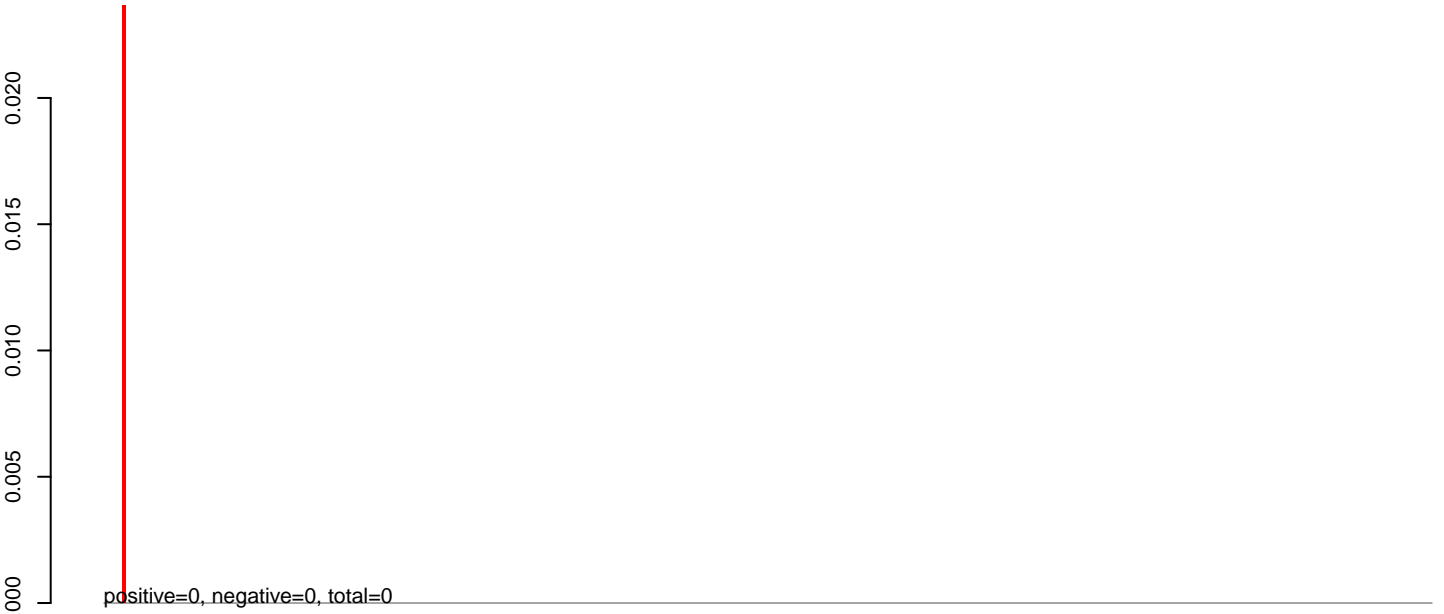
AnGam_Ovary_BetaE.24_35.rep



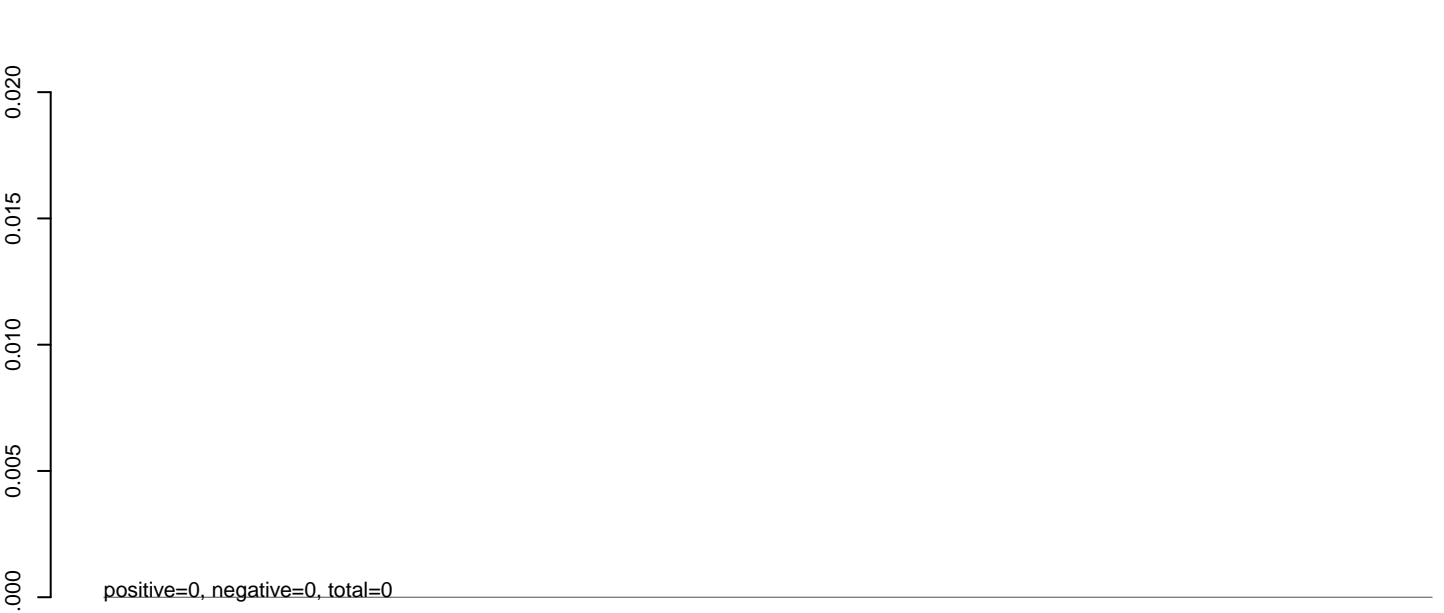
AnGam_Ovary_BetaE.rep



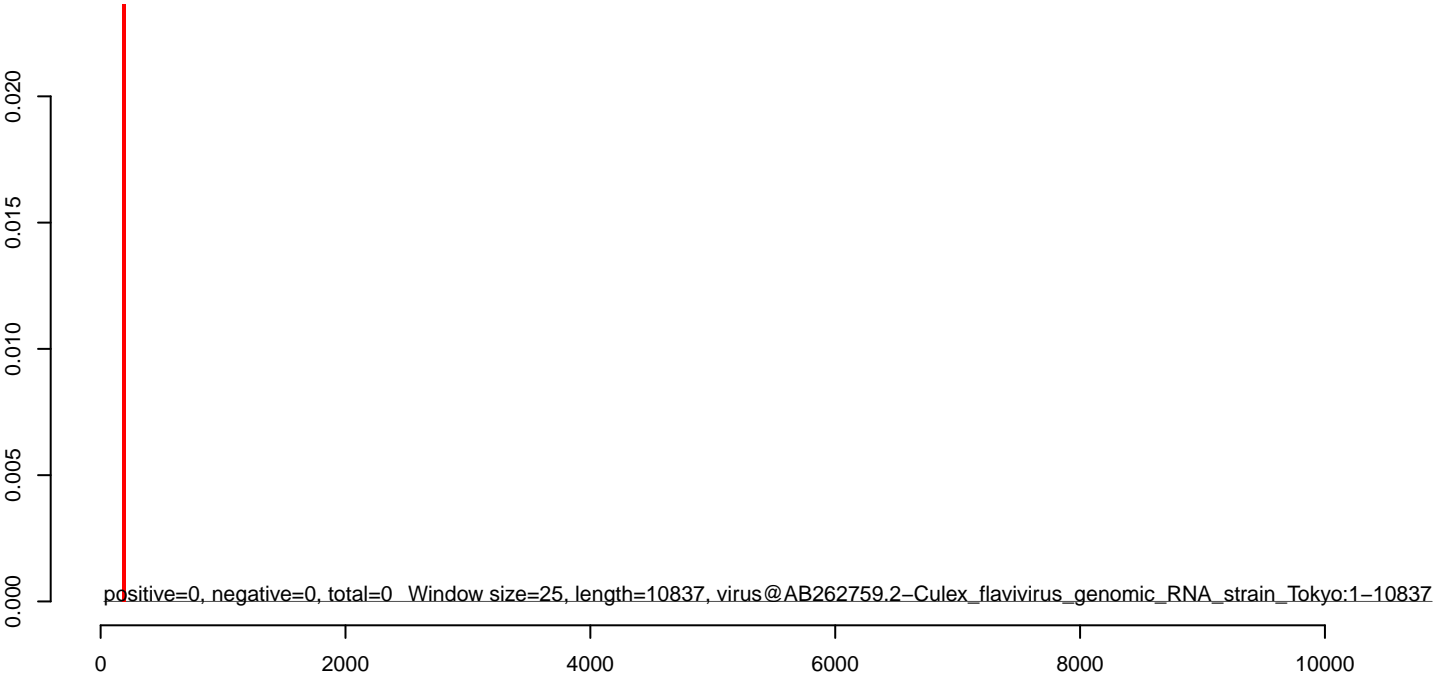
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



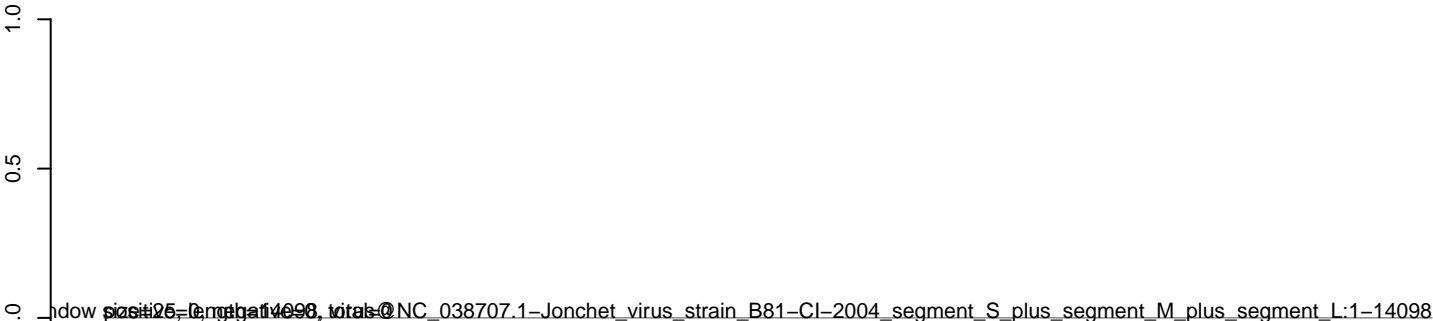
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



0 2000 4000 6000 8000 10000 12000 14000

AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

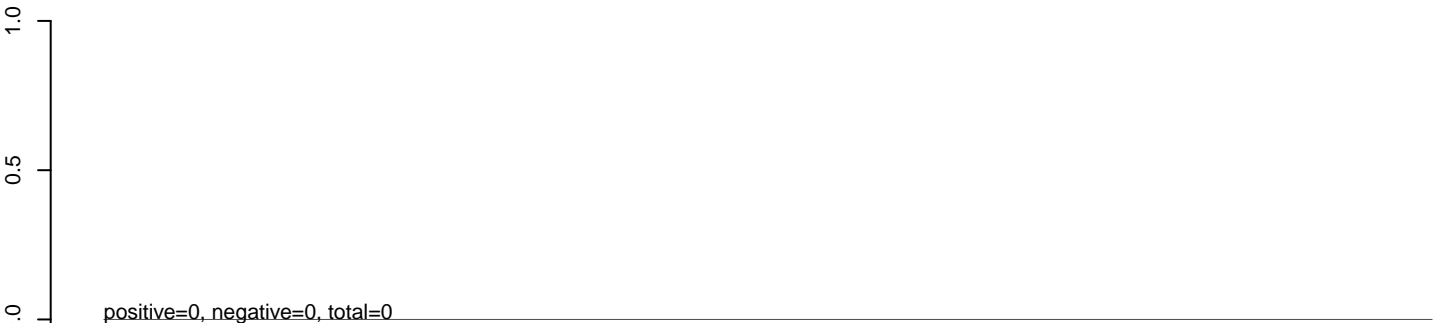


AnGam_Ovary_BetaE.rep

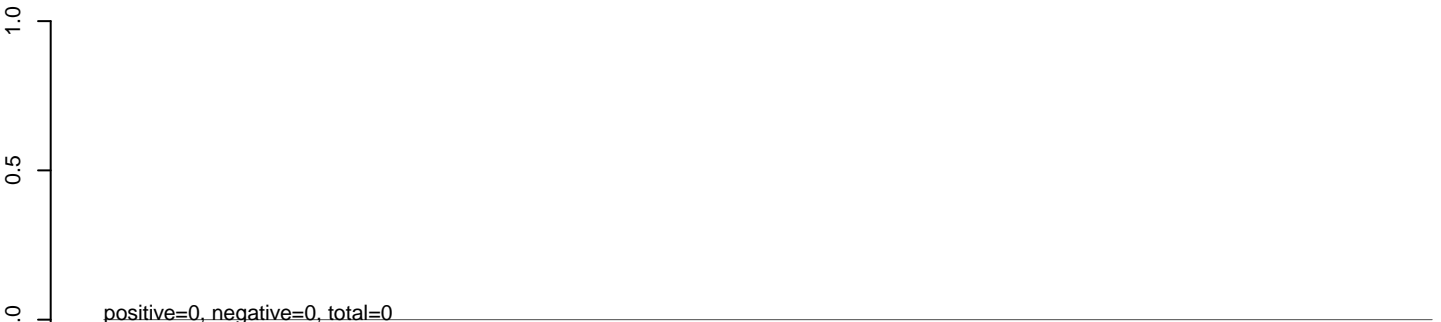


0 2000 4000 6000 8000 10000 12000

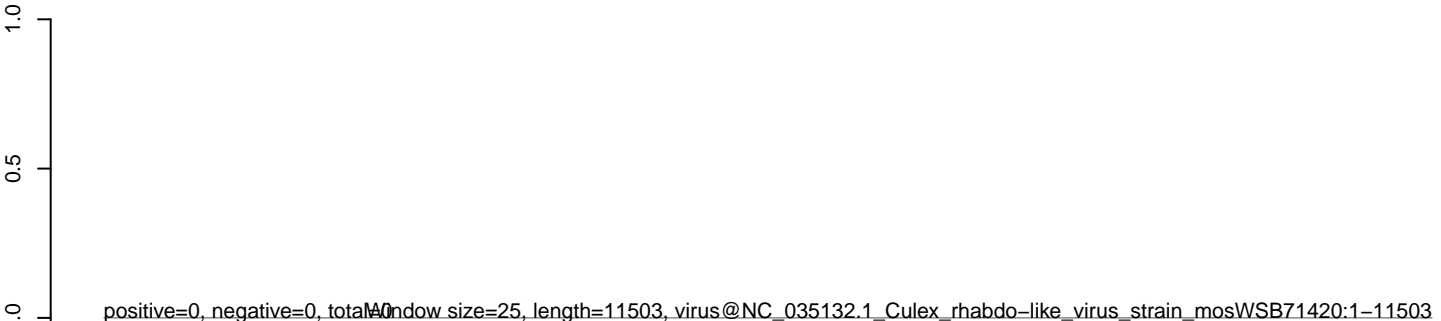
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



0 2000 4000 6000 8000 10000 12000

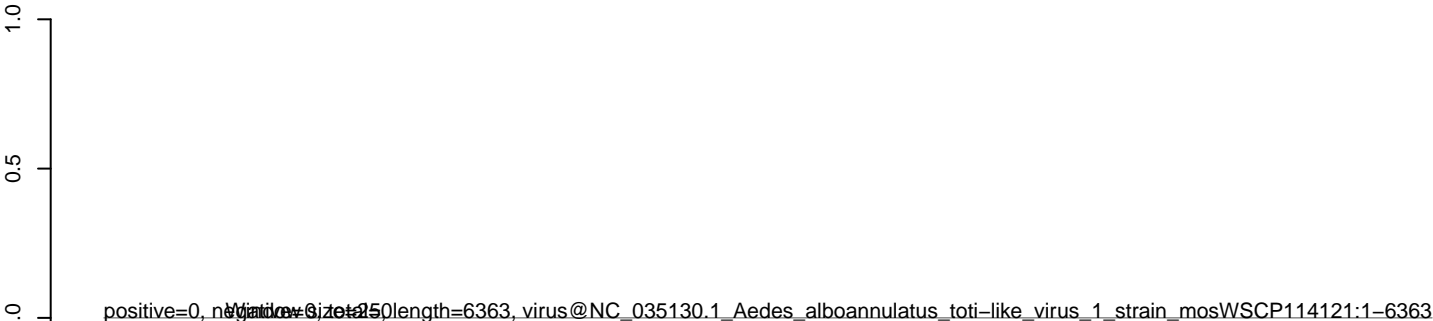
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

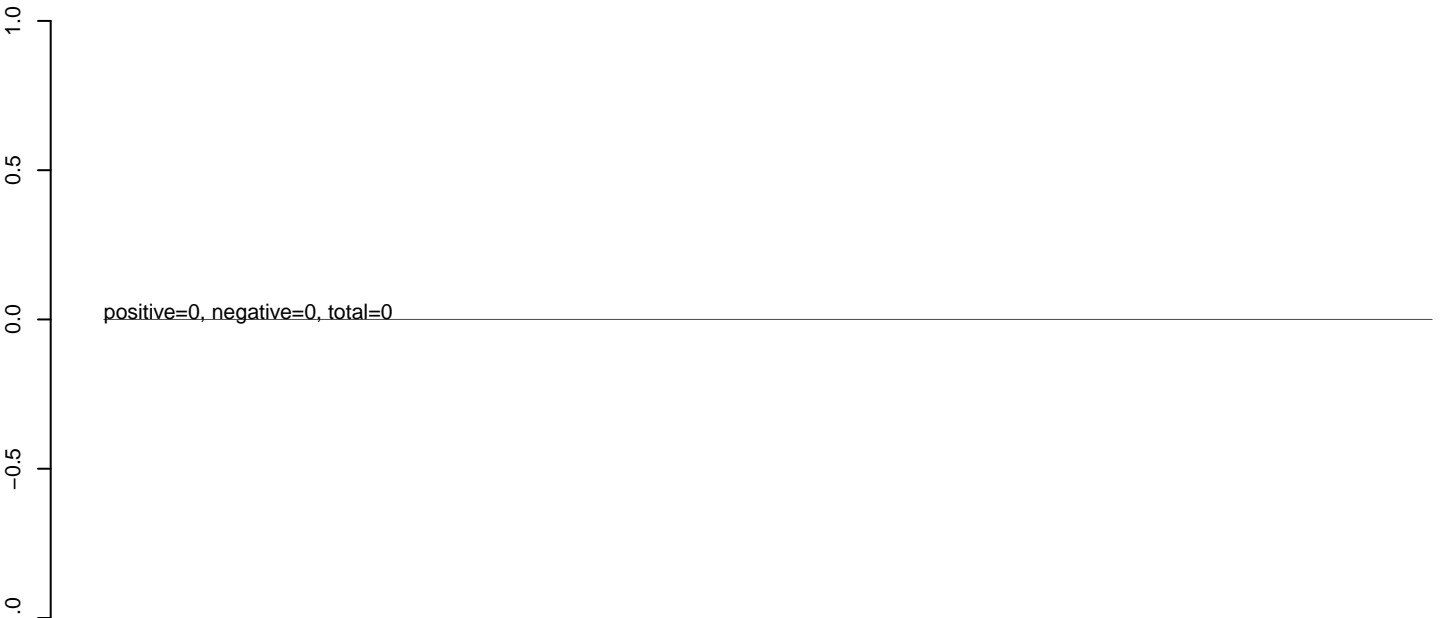


AnGam_Ovary_BetaE.rep

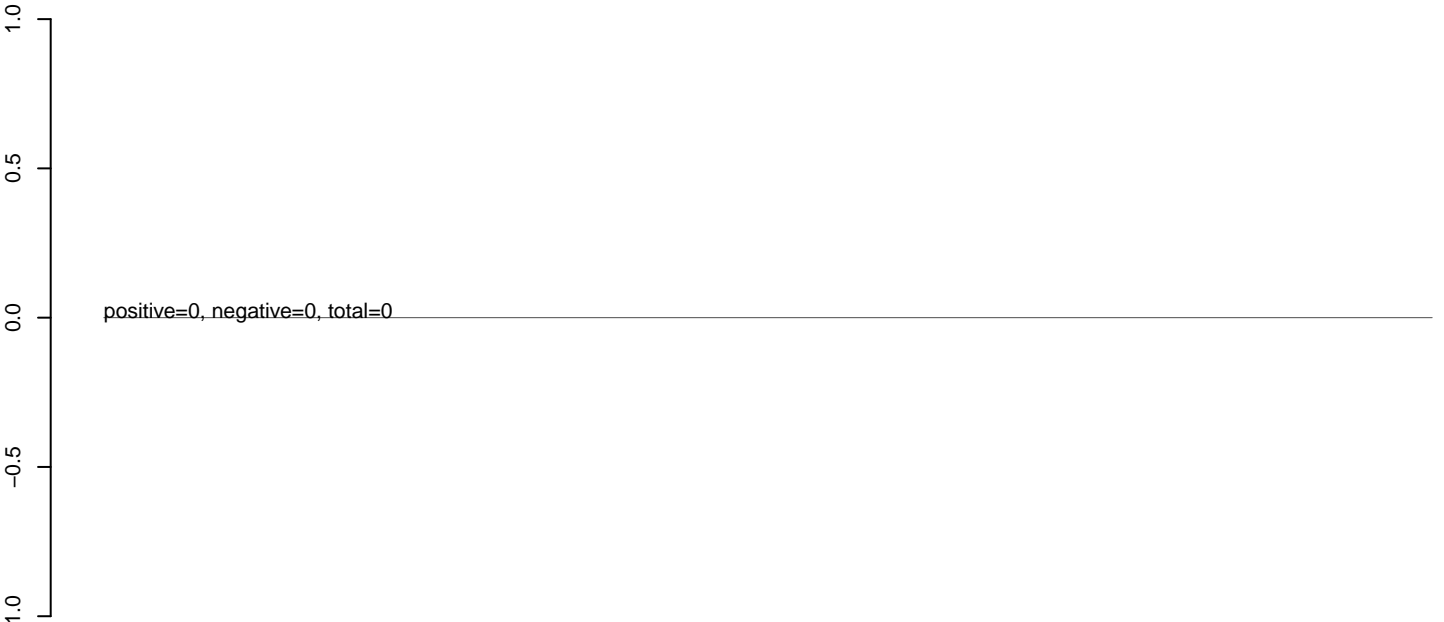


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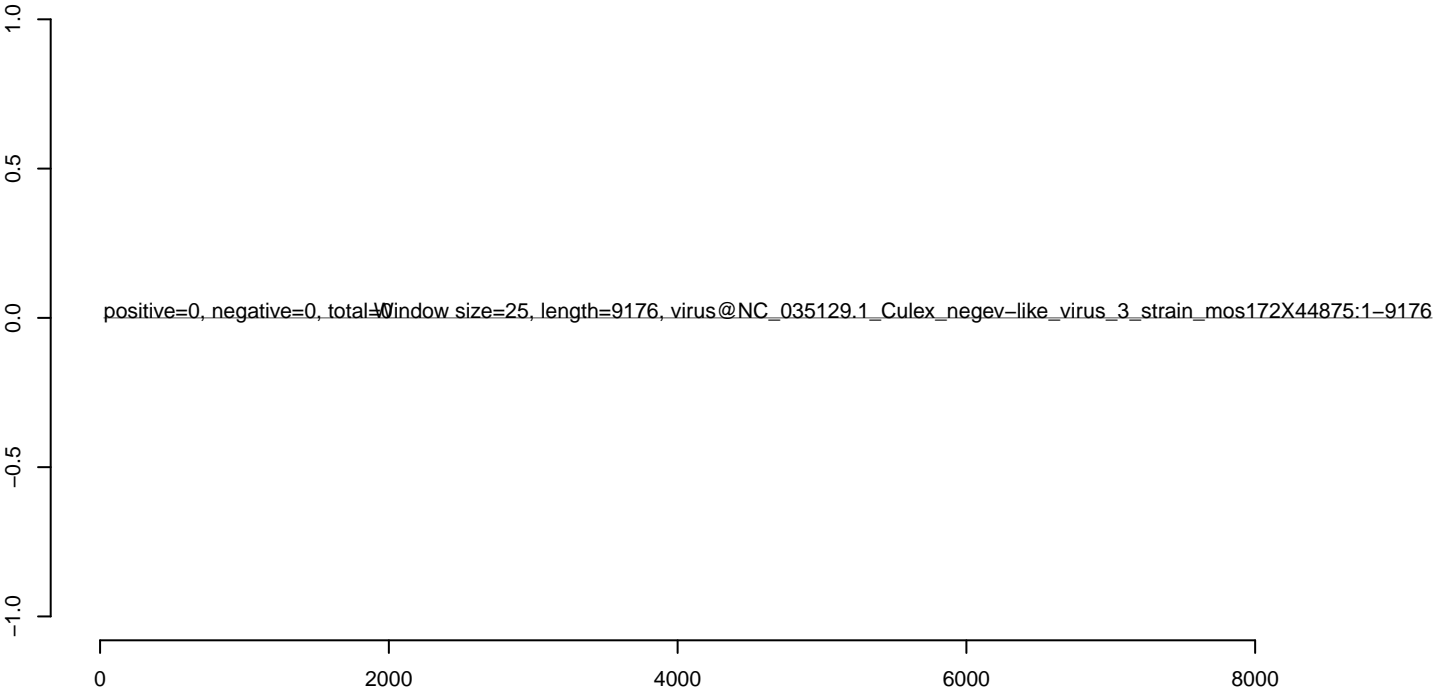
AnGam_Ovary_BetaE.18_23.rep



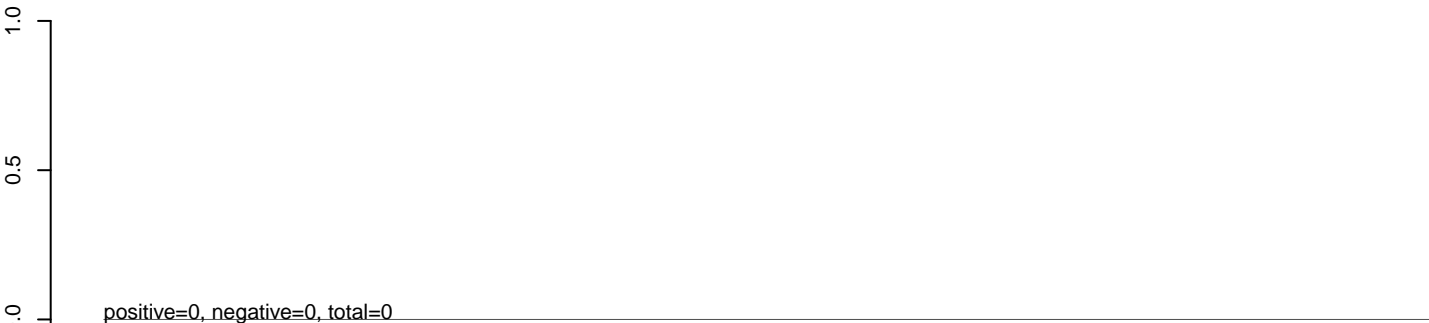
AnGam_Ovary_BetaE.24_35.rep



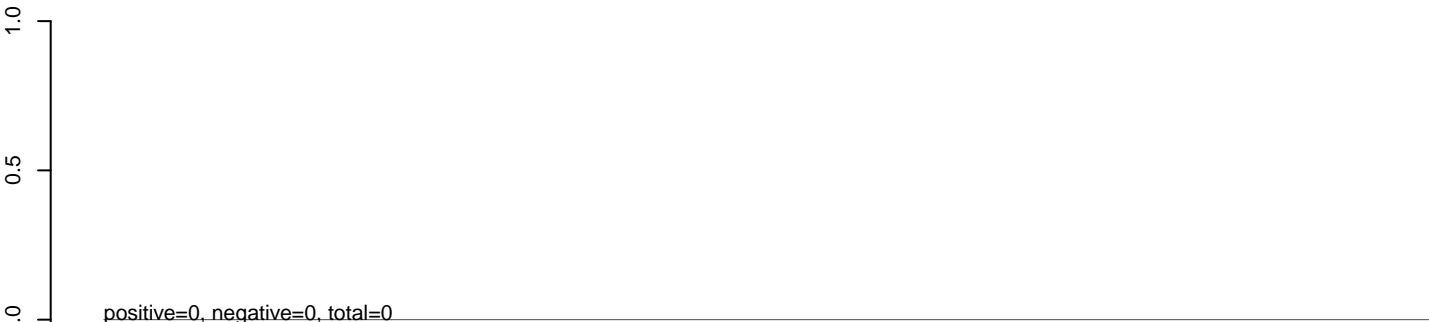
AnGam_Ovary_BetaE.rep



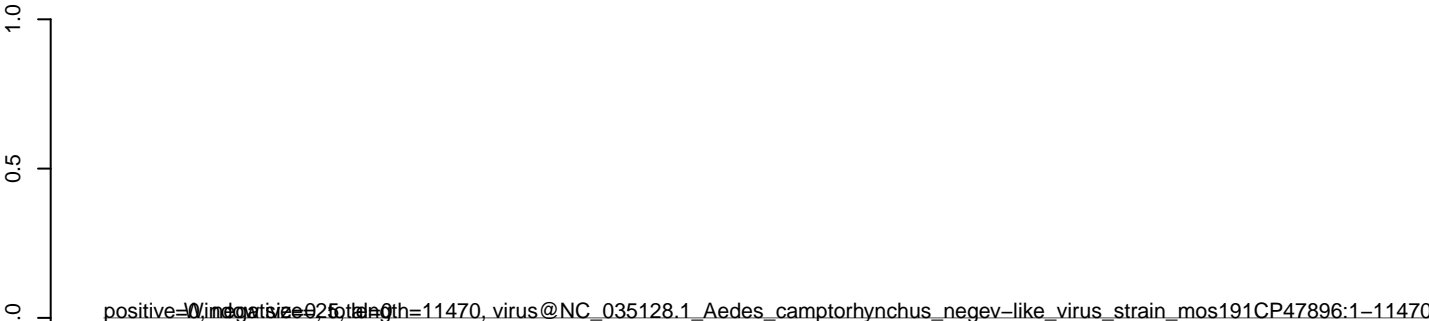
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

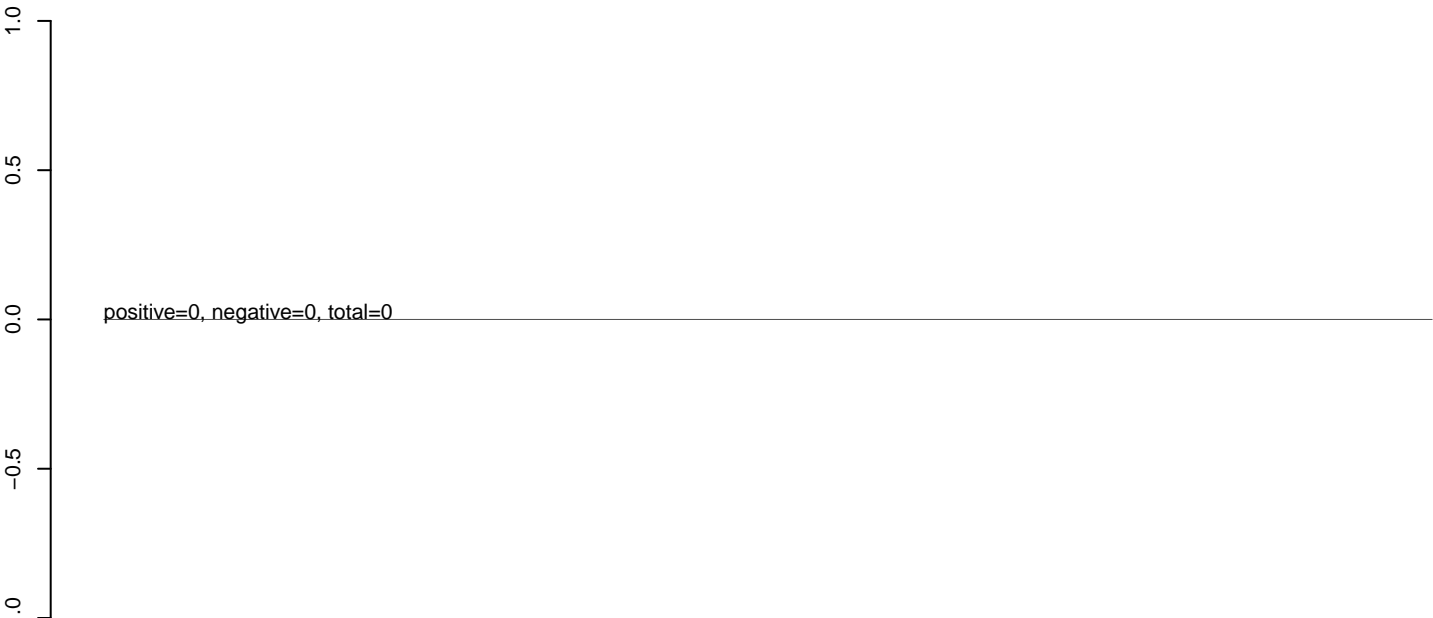


AnGam_Ovary_BetaE.rep

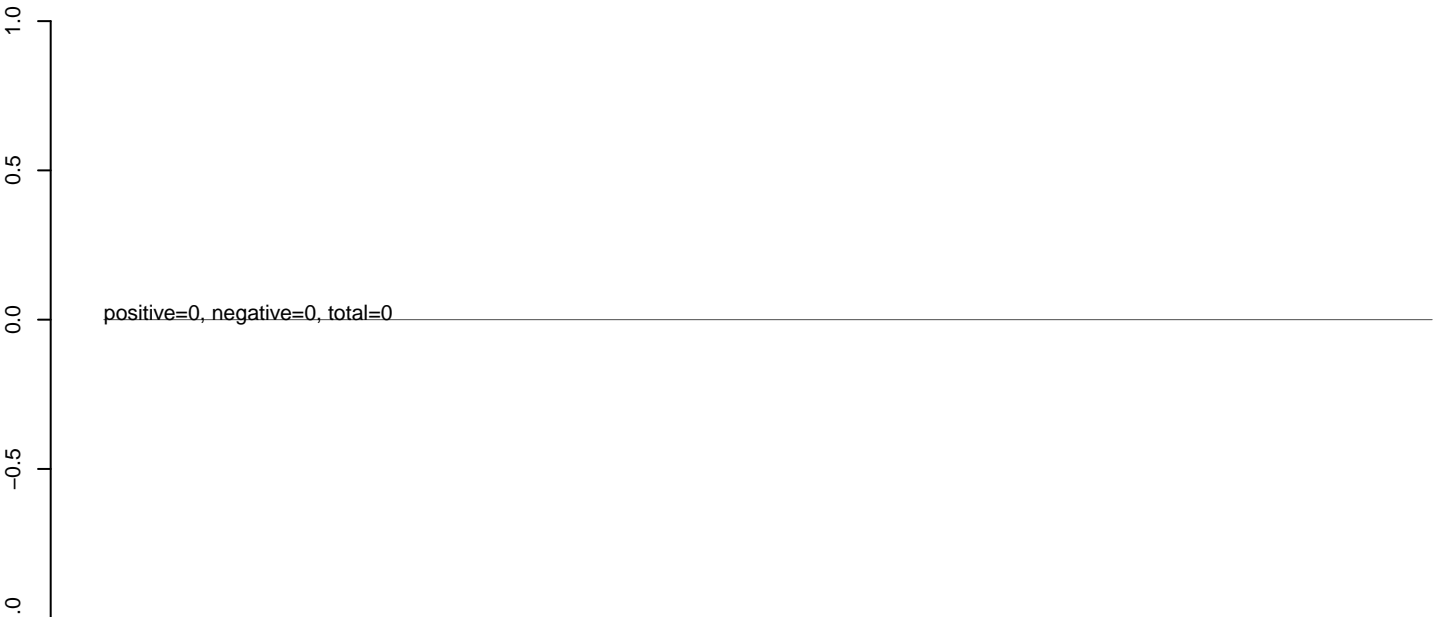


0 2000 4000 6000 8000 10000

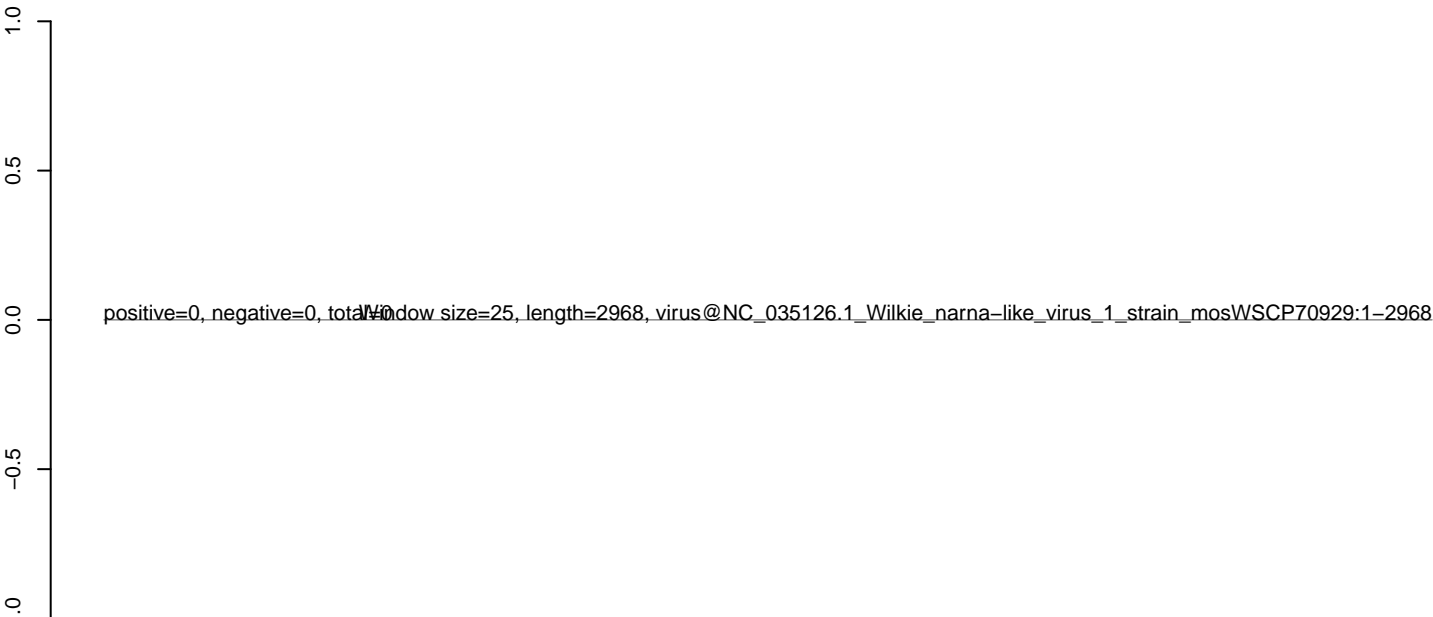
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

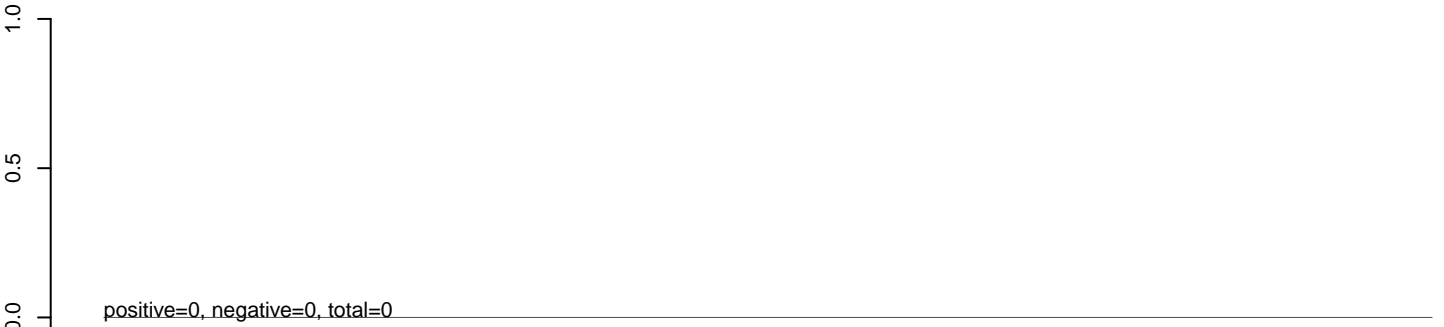


AnGam_Ovary_BetaE.rep

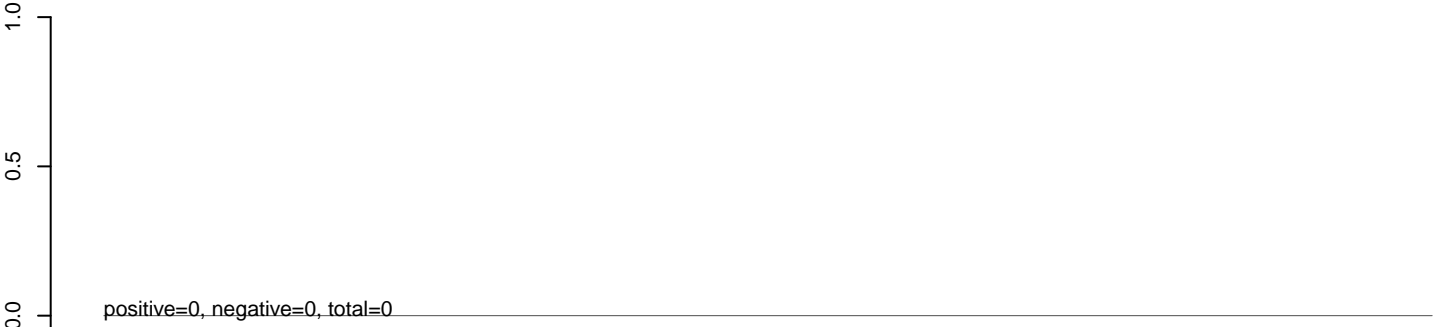


0 500 1000 1500 2000 2500 3000

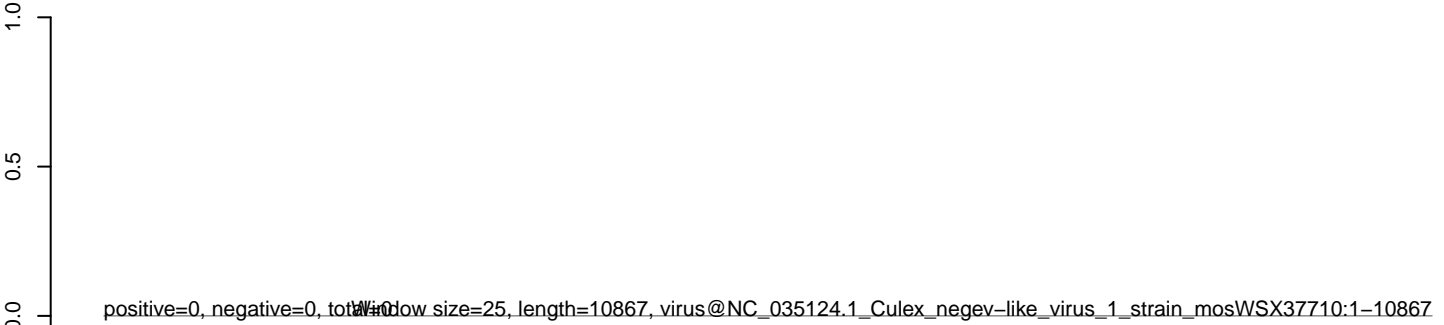
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

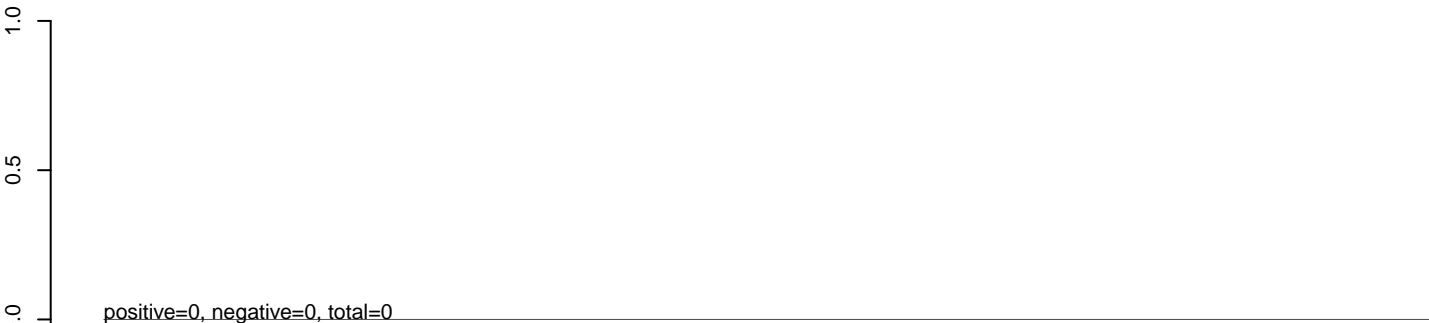


AnGam_Ovary_BetaE.rep

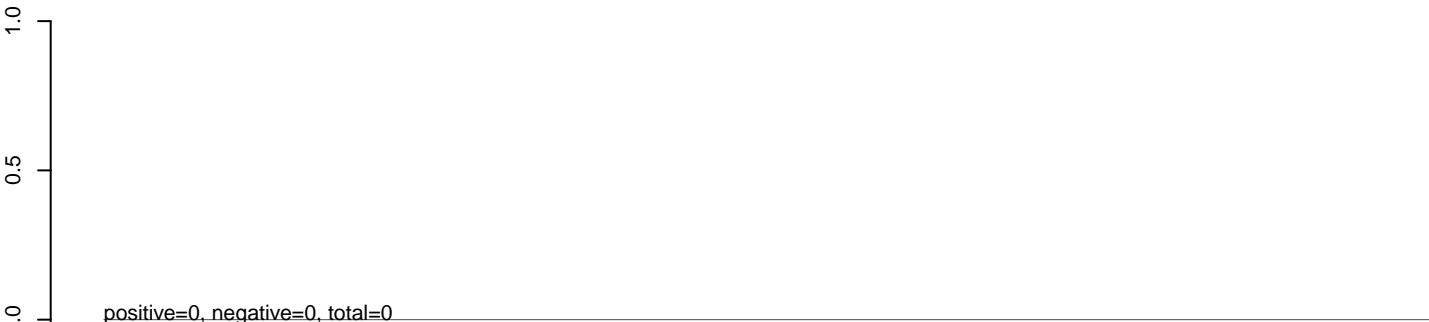


0 2000 4000 6000 8000 10000

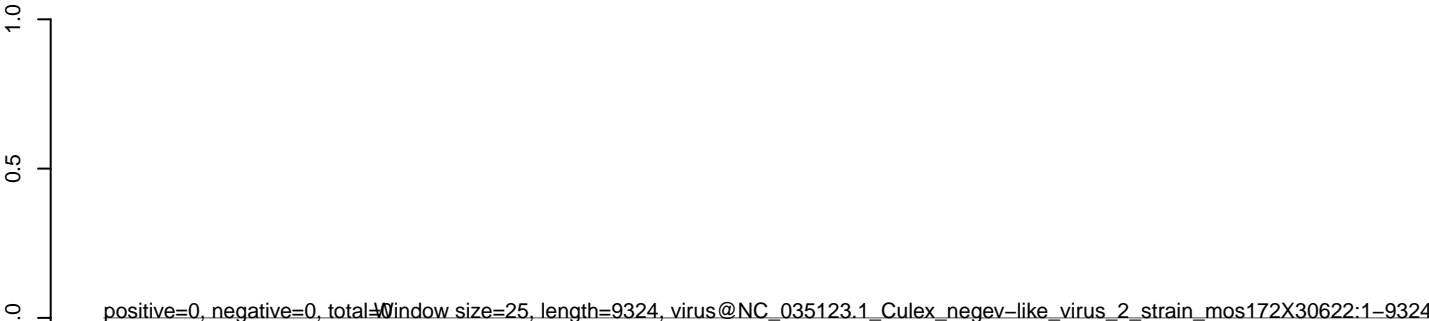
AnGam_Ovary_BetaE.18_23.rep



AnGam_Ovary_BetaE.24_35.rep

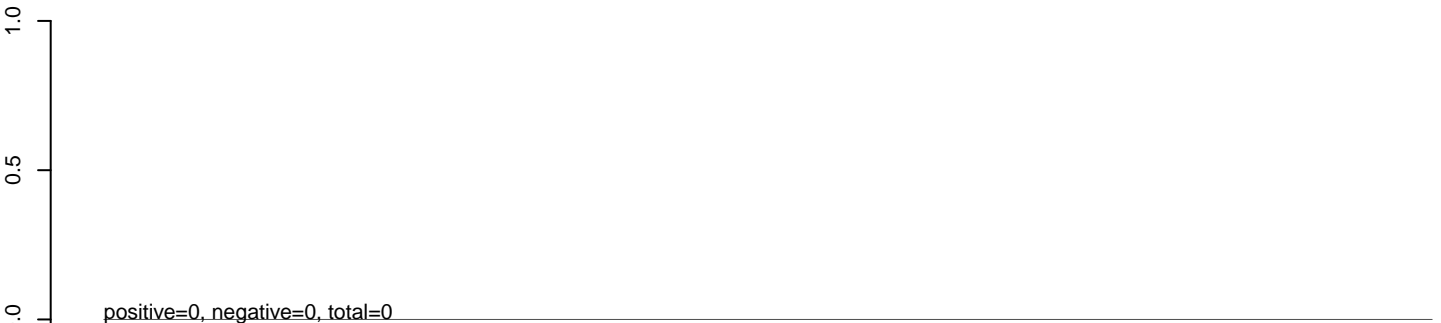


AnGam_Ovary_BetaE.rep

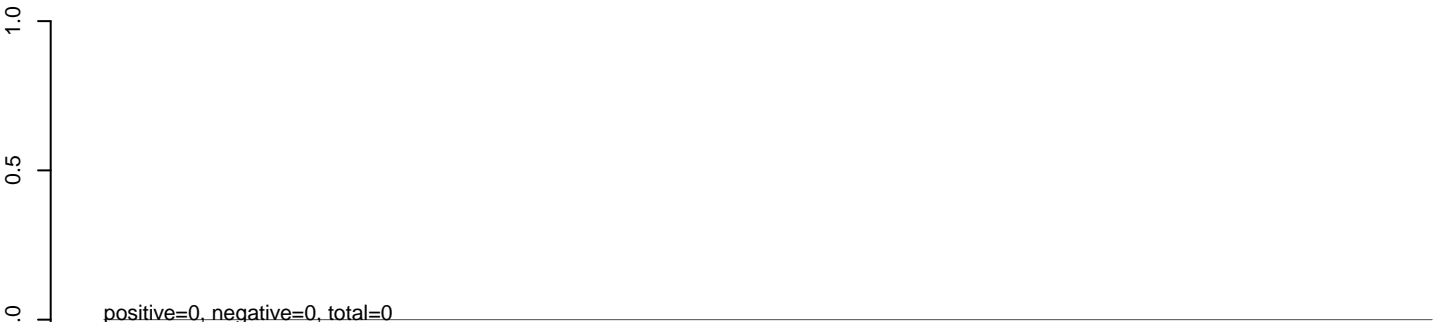


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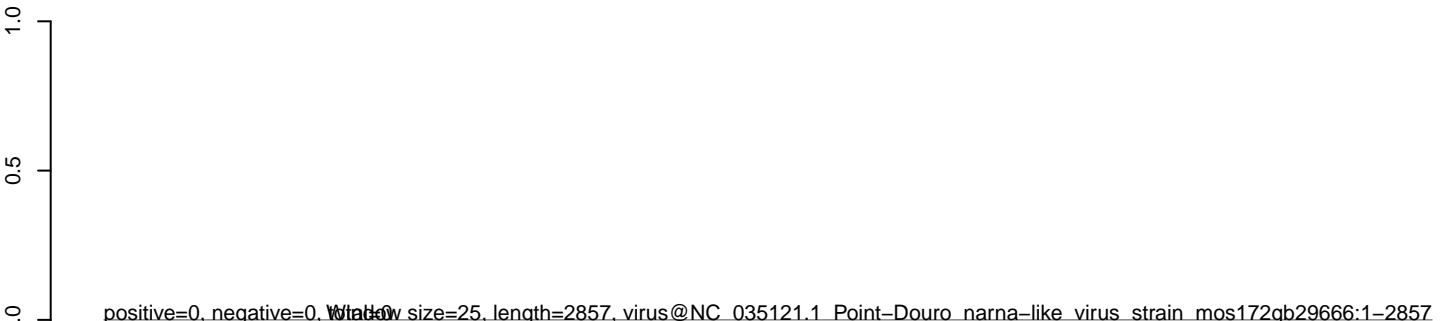
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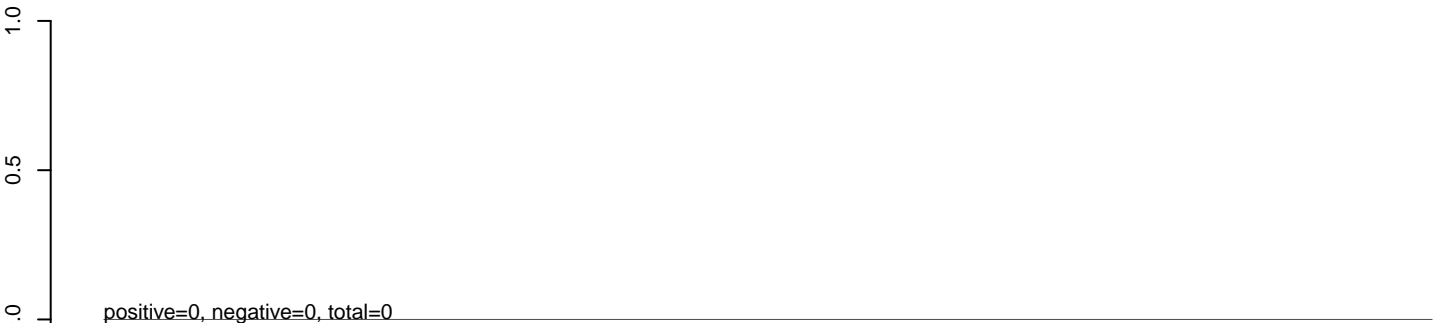


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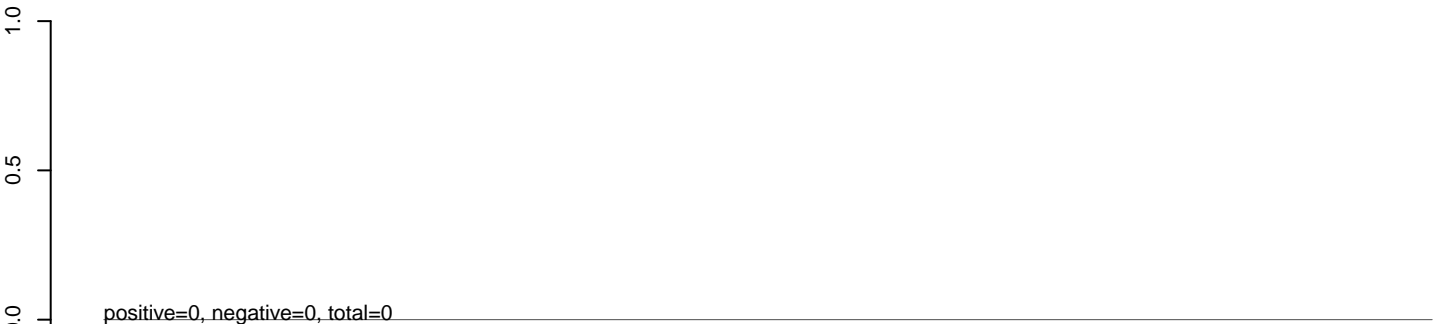


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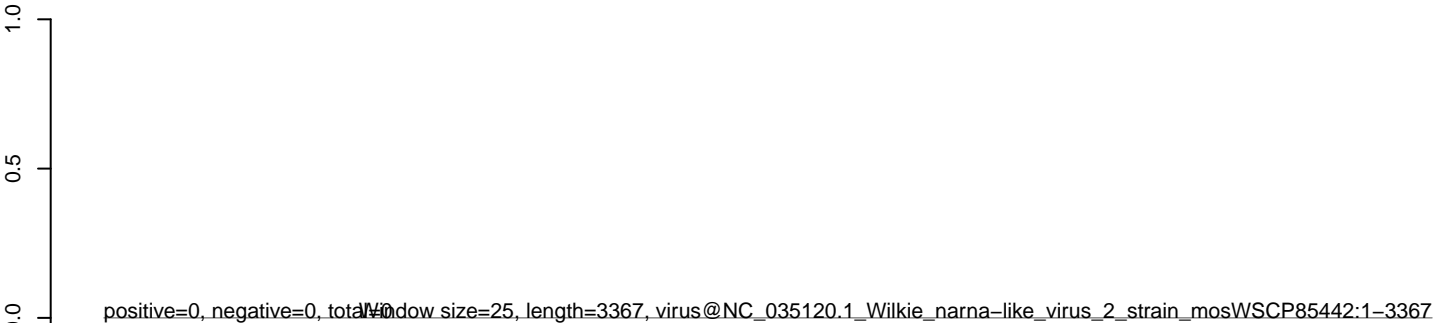
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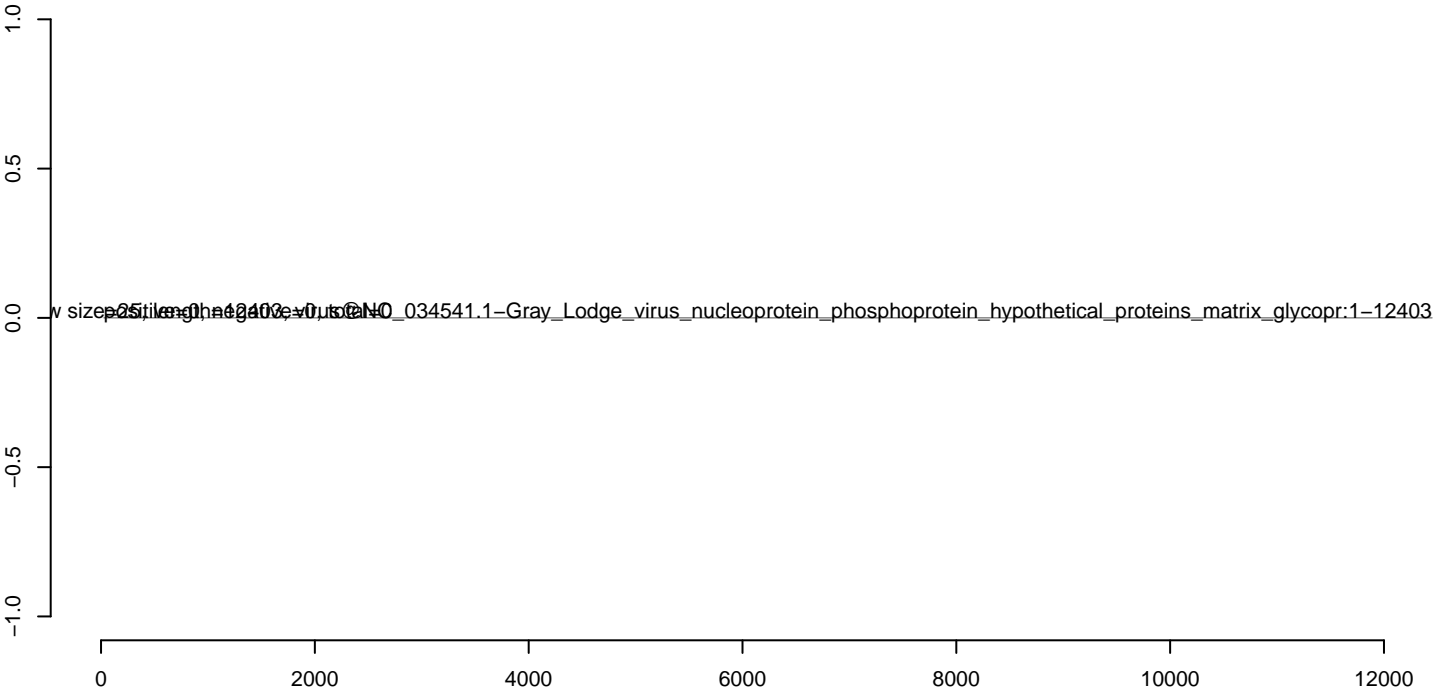
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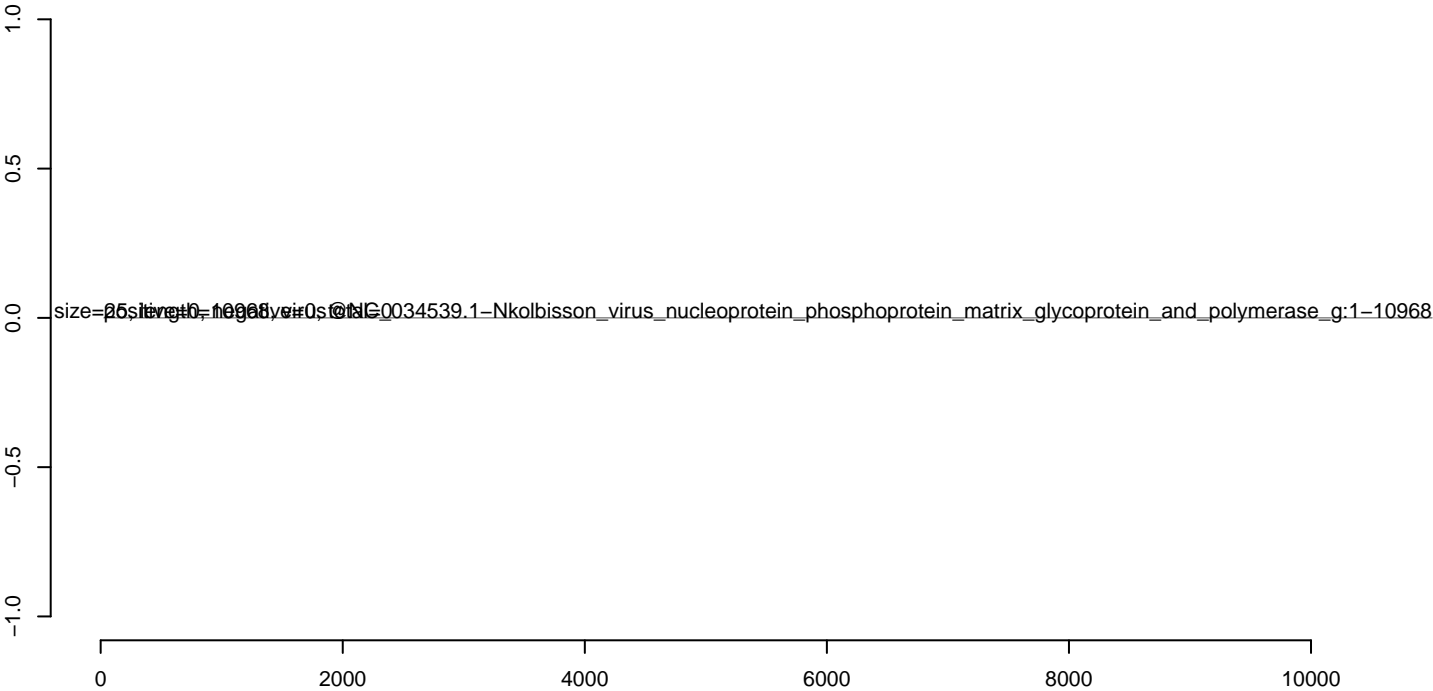
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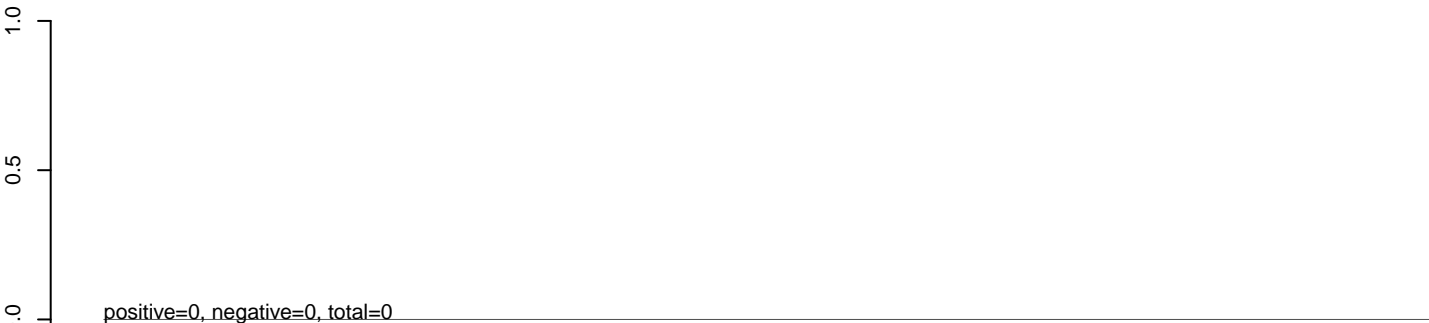
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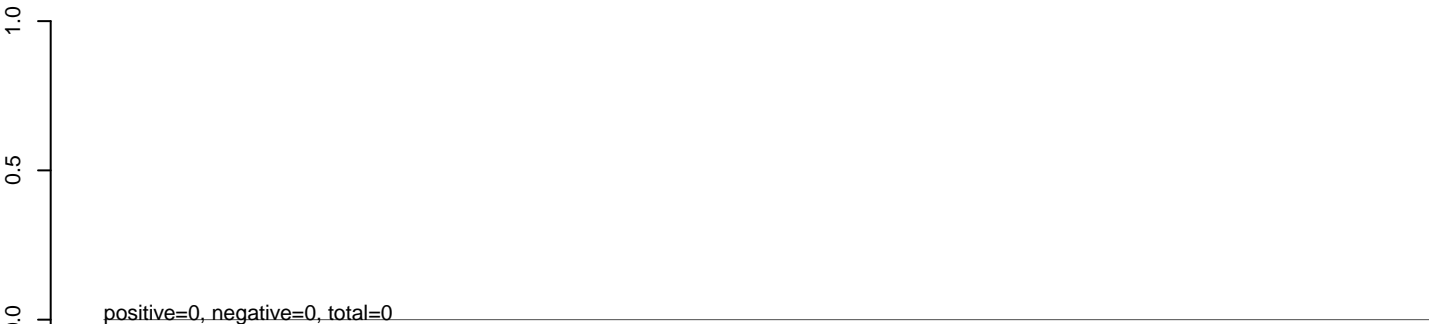
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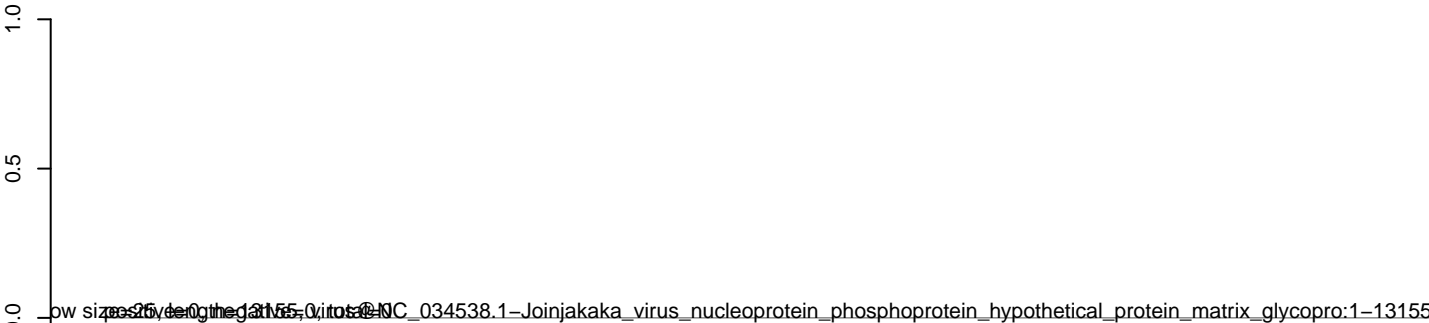
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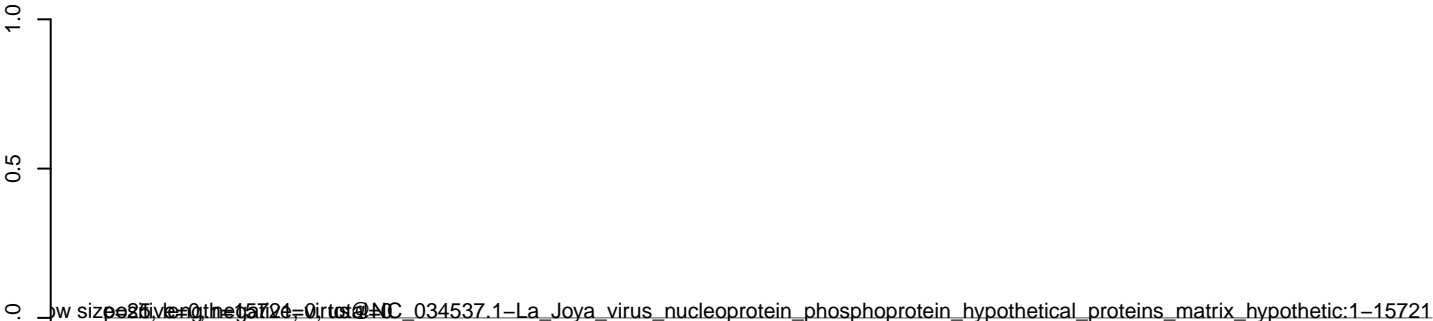
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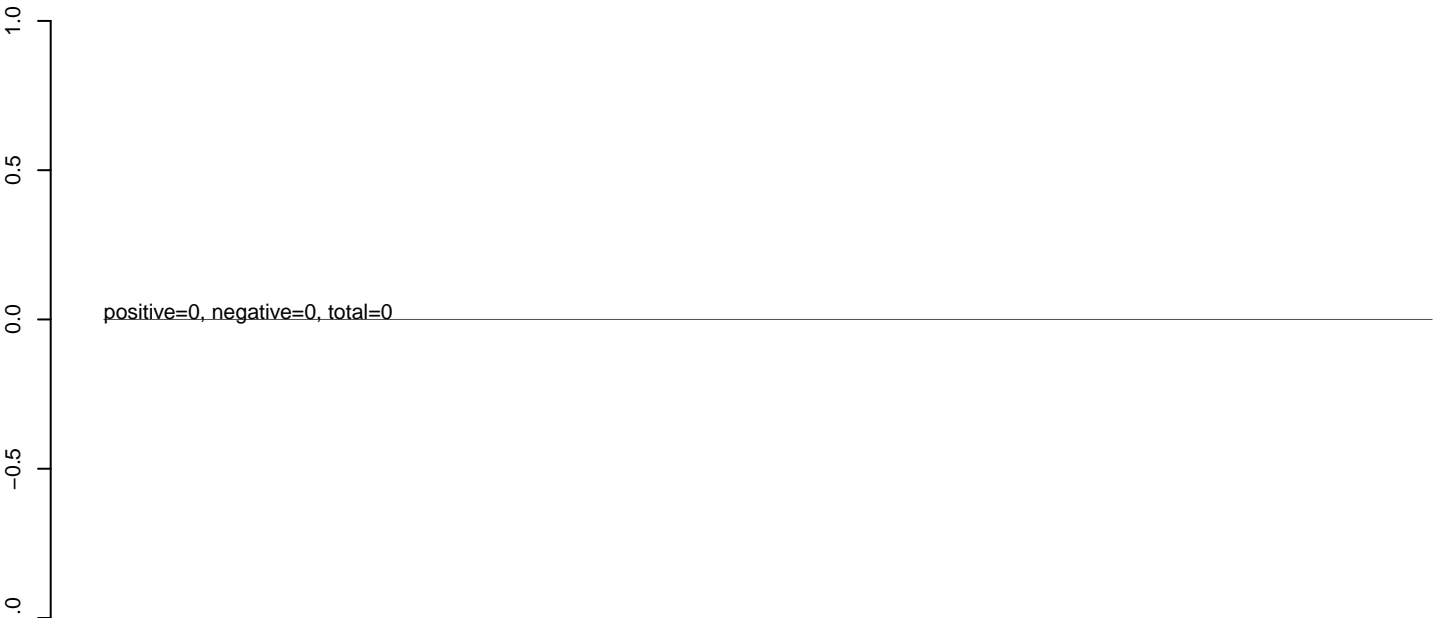
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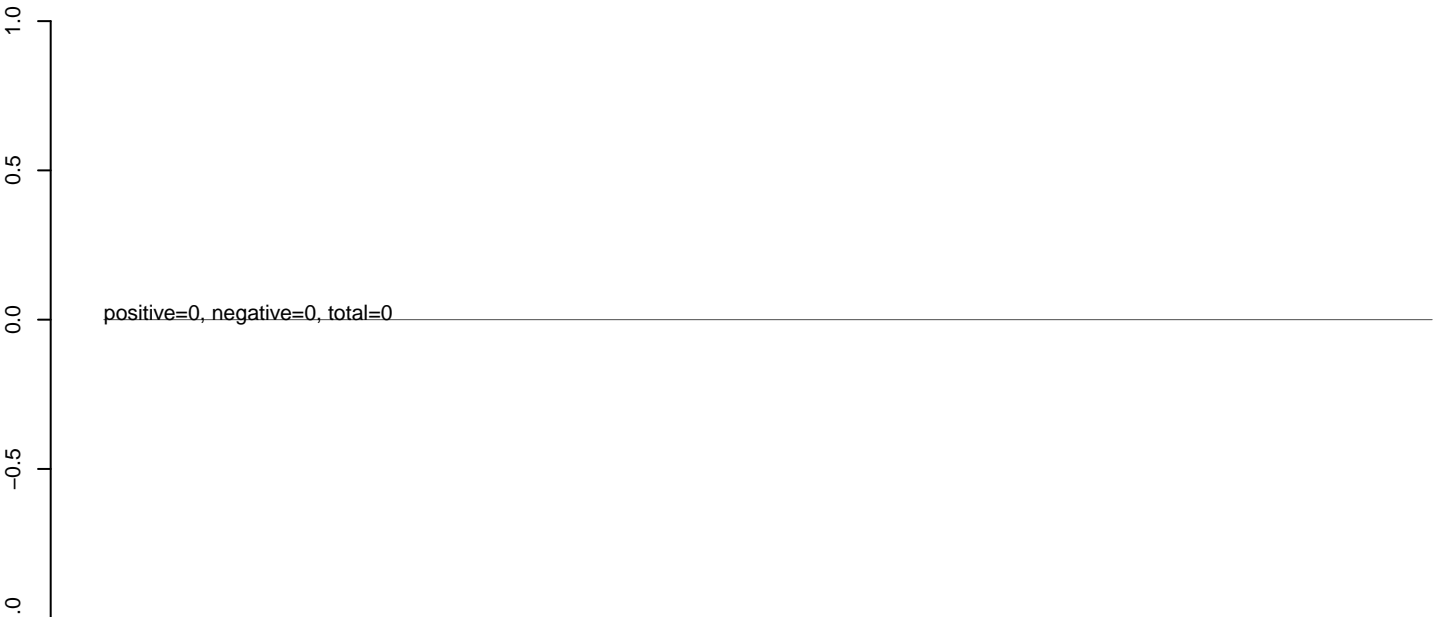
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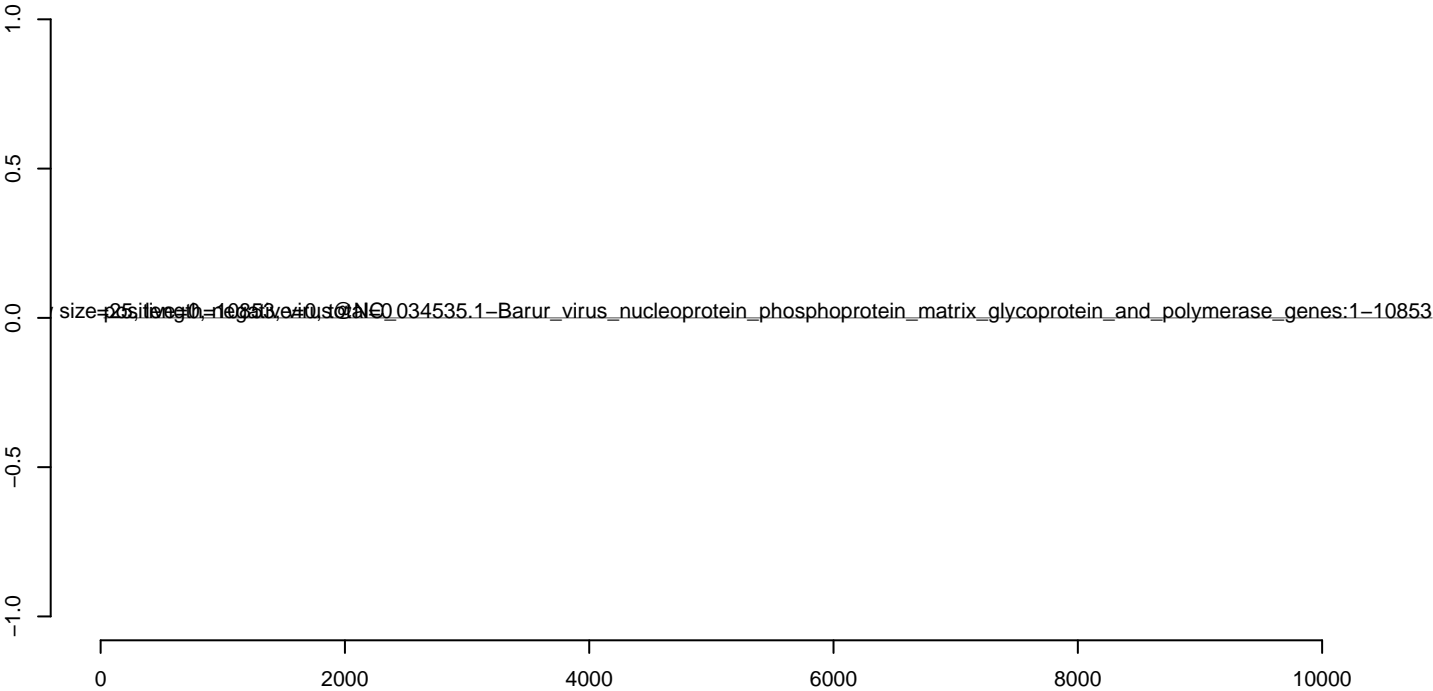
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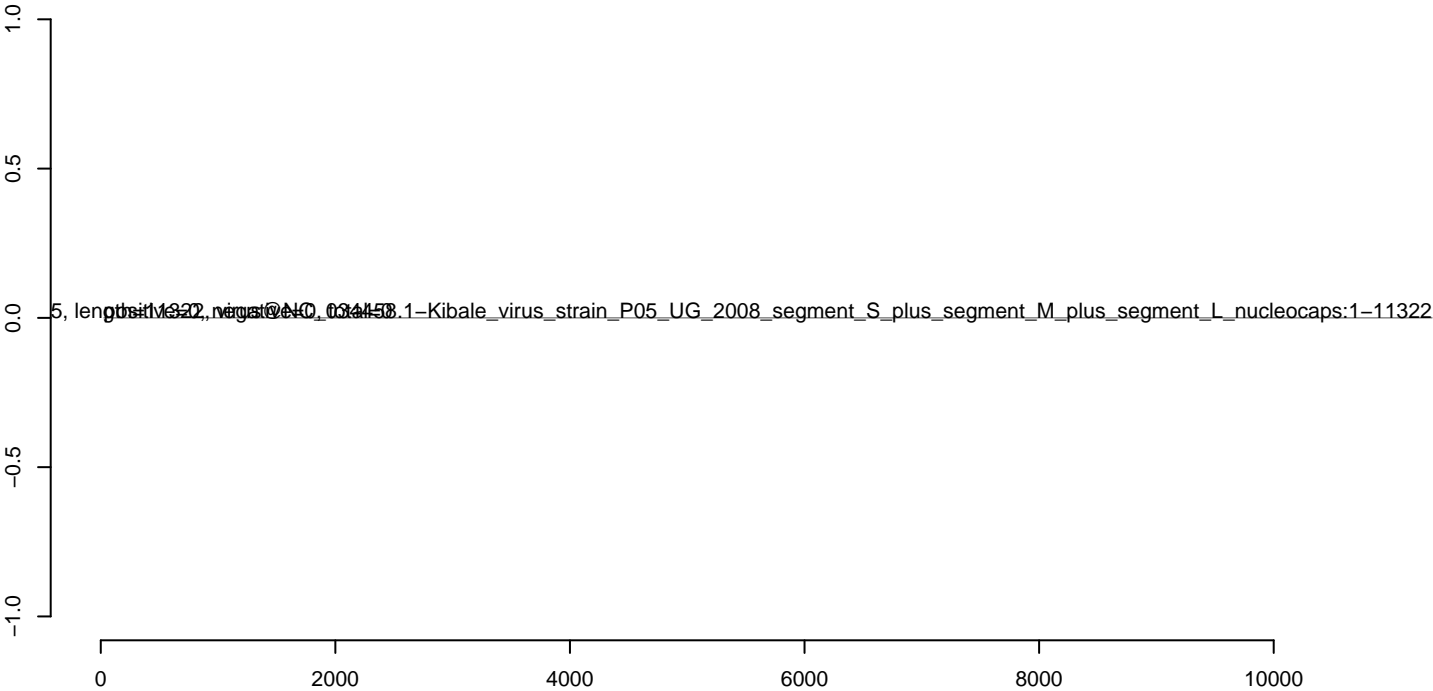
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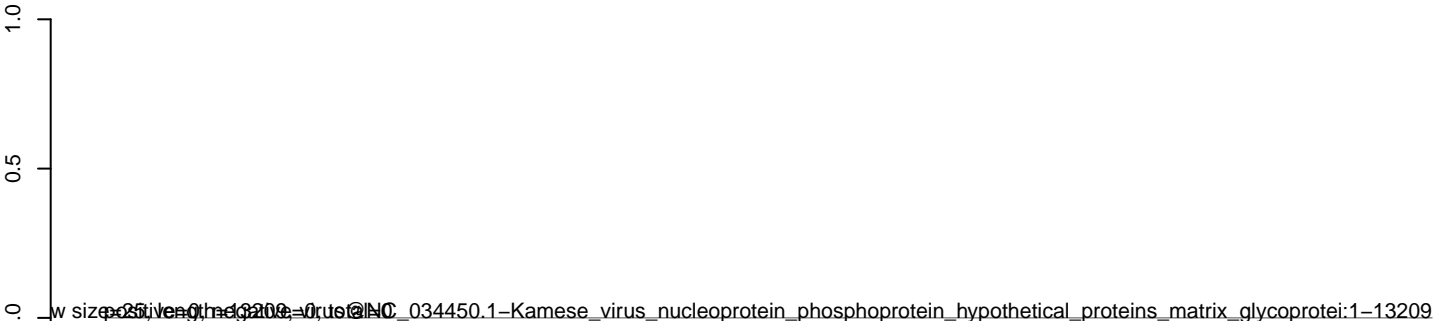
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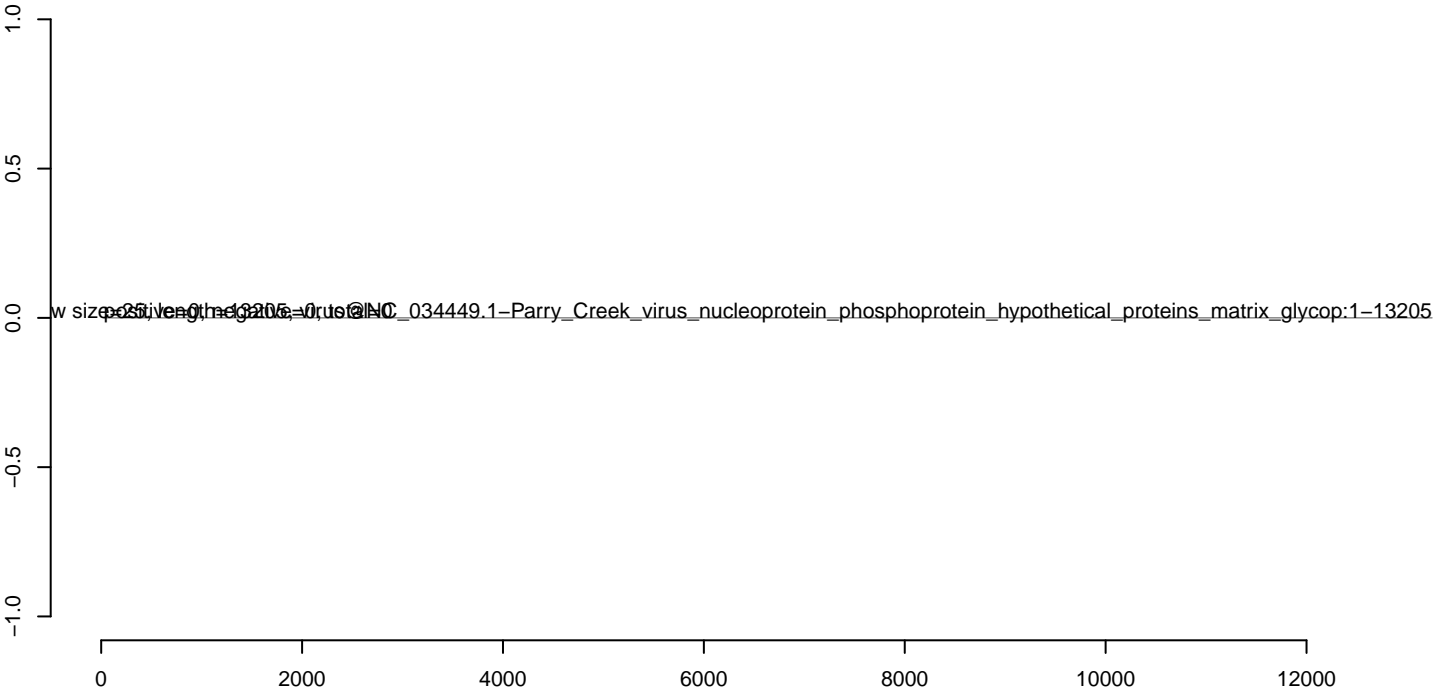
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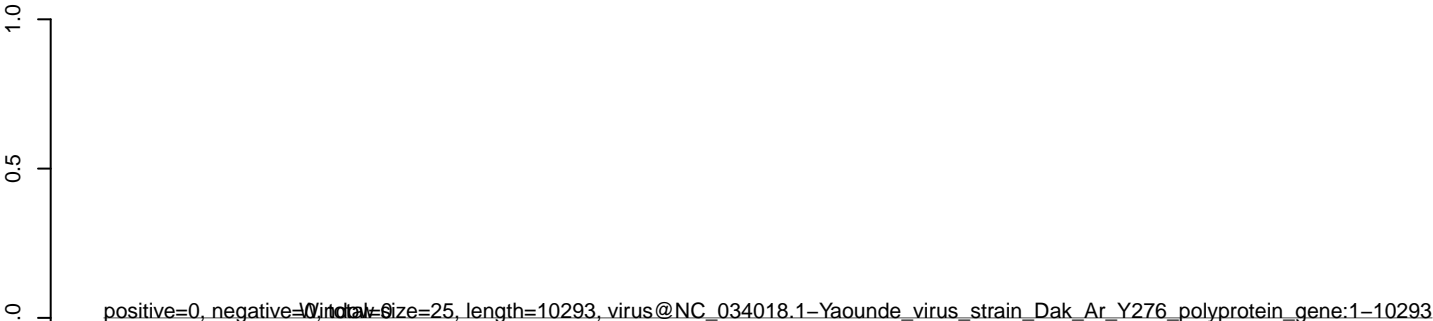
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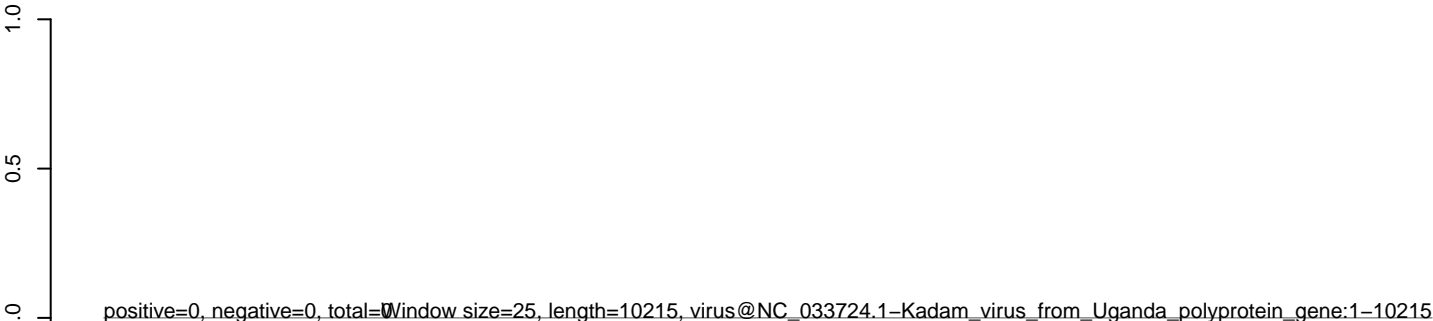
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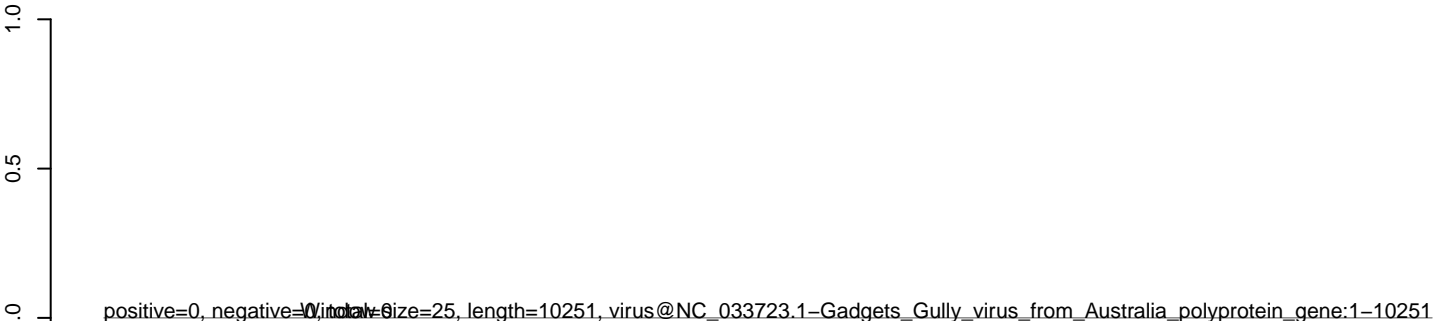
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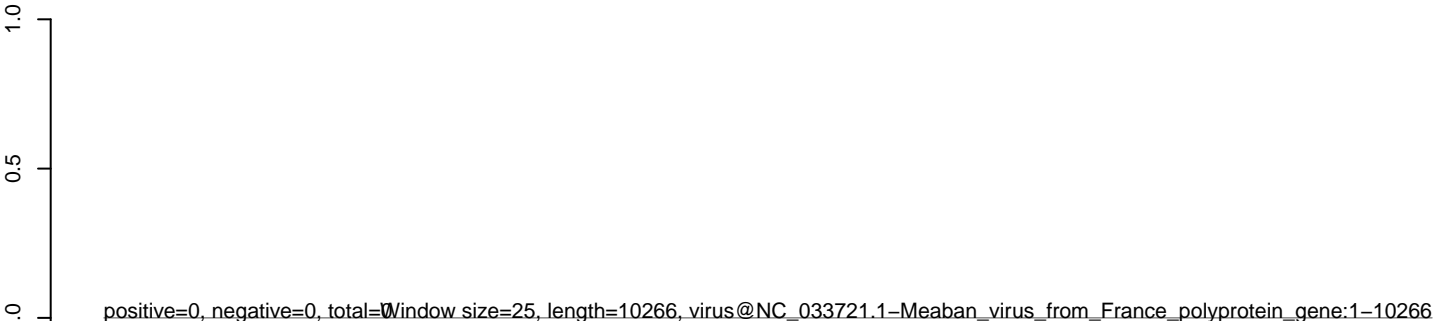
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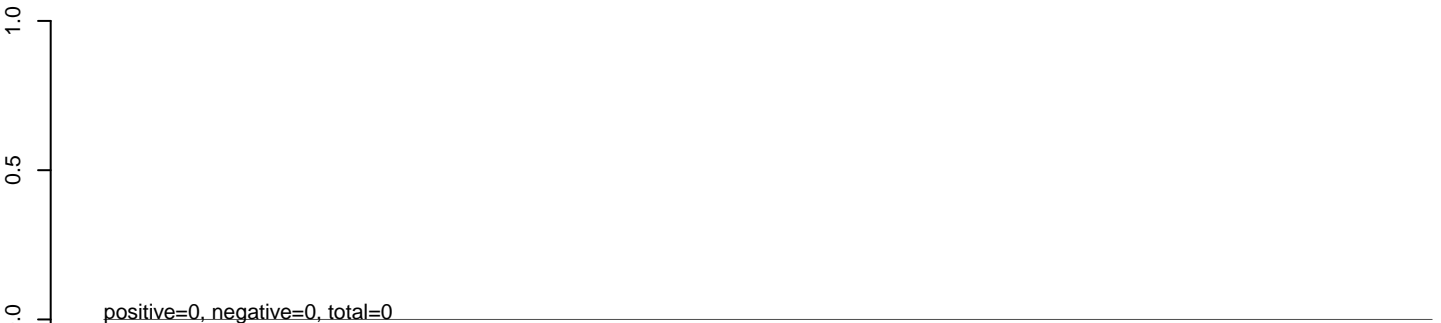
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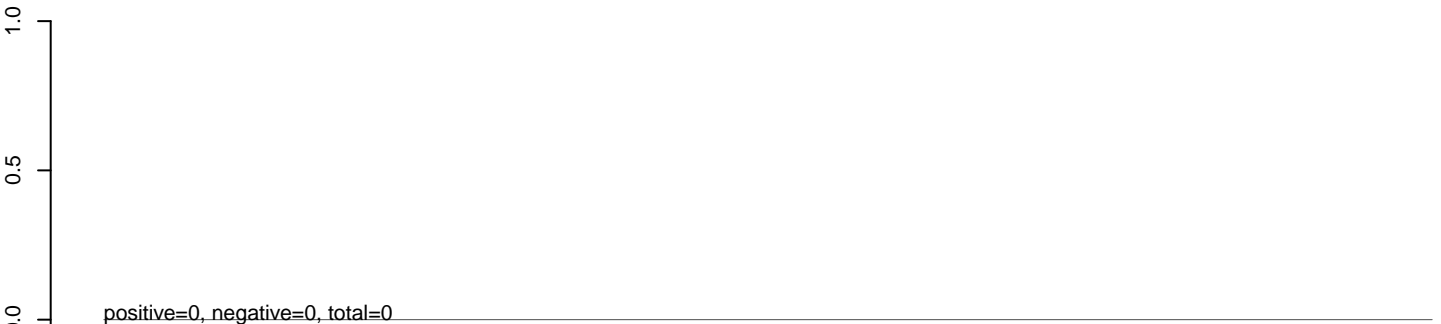
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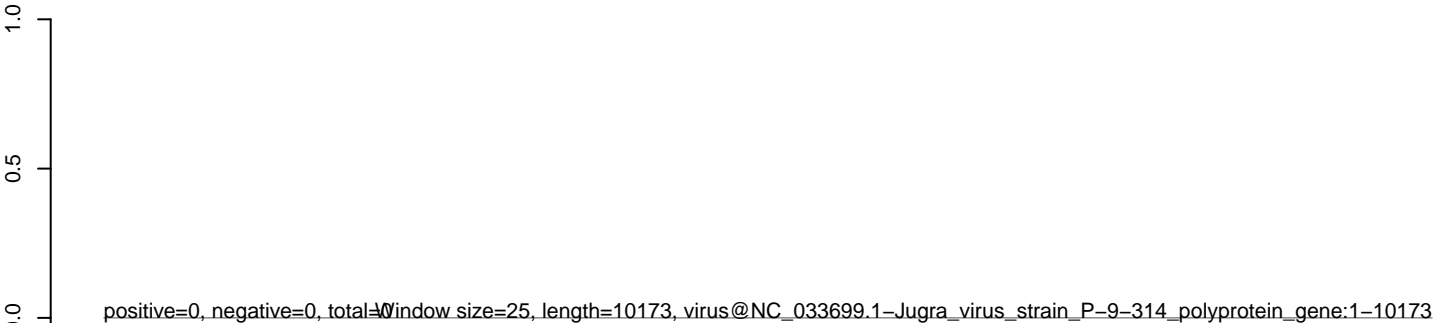
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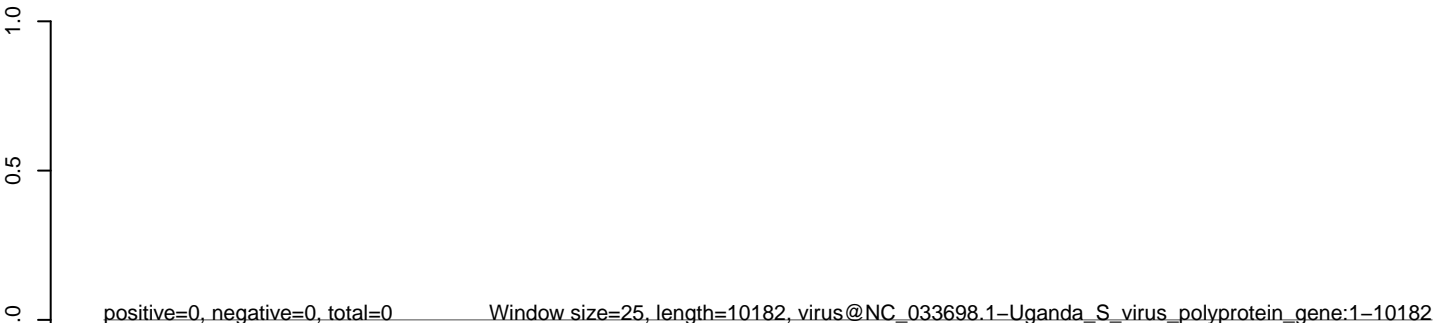
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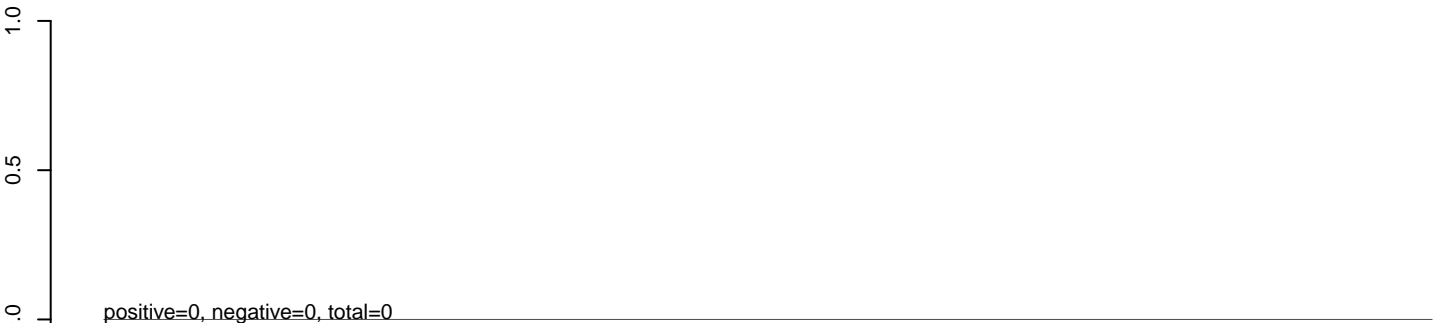


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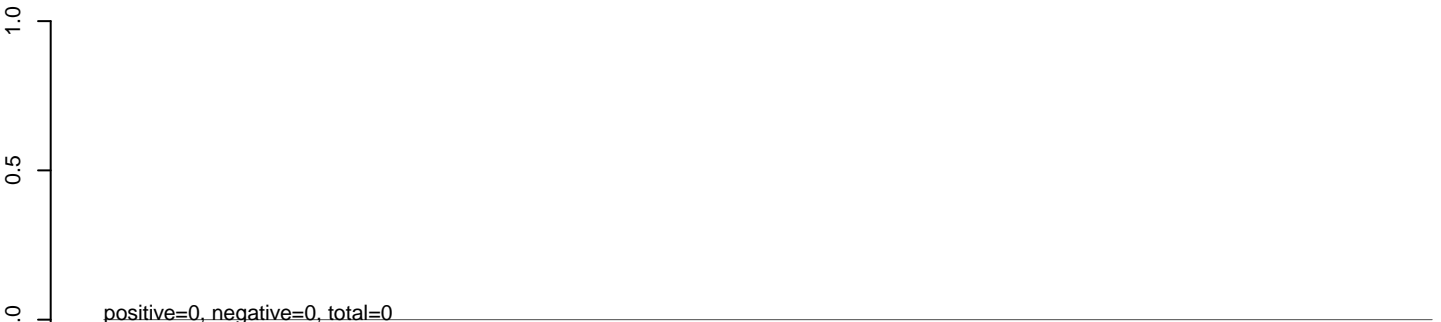


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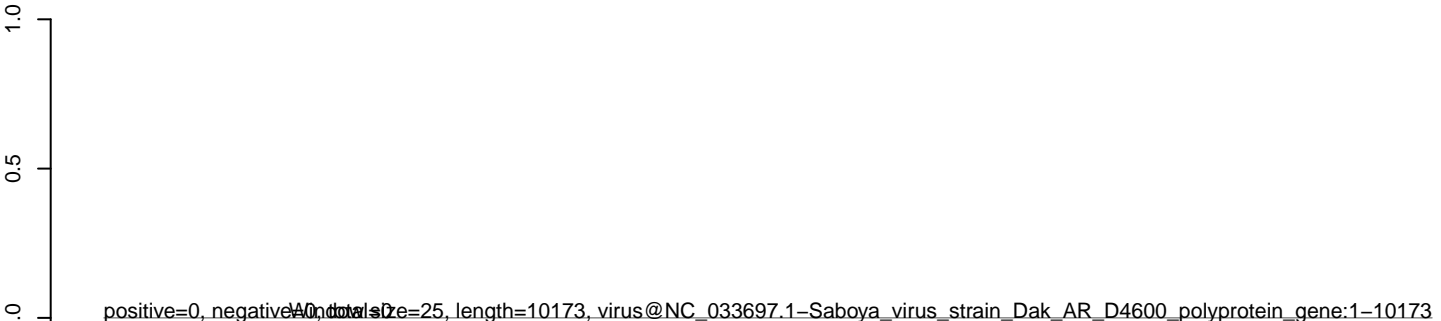
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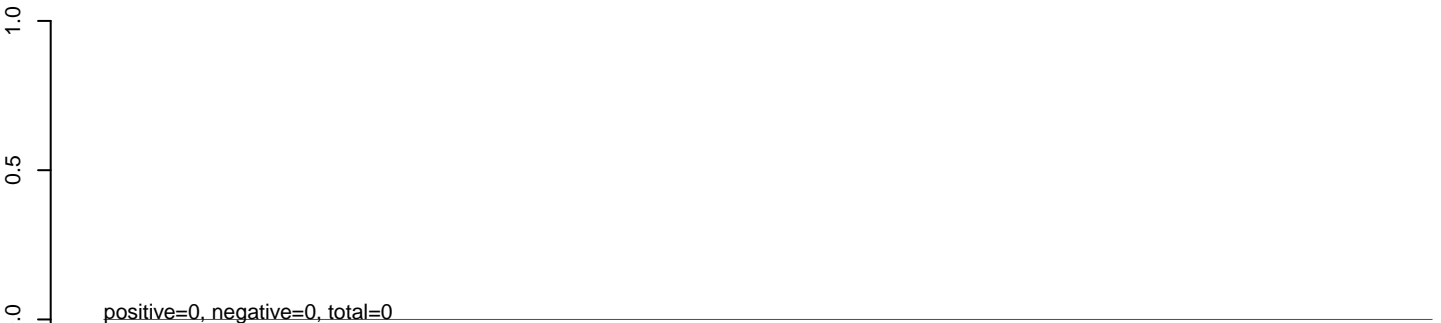
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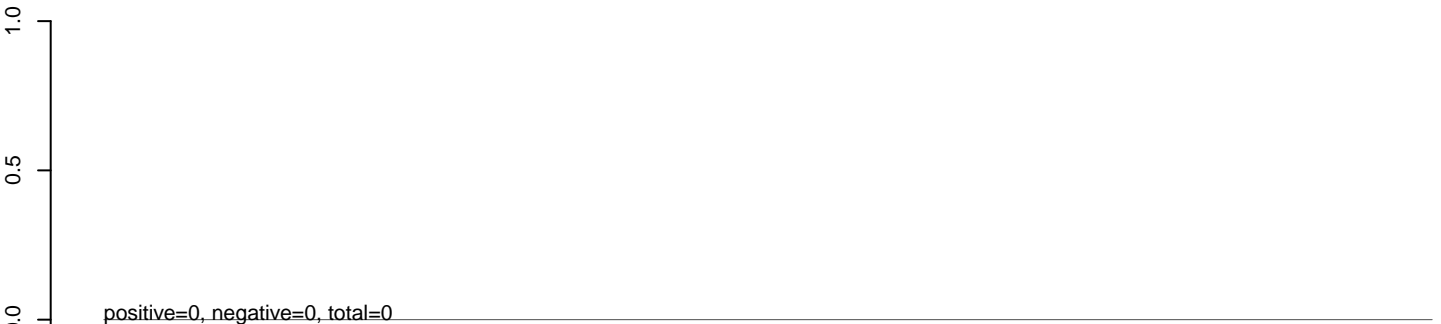
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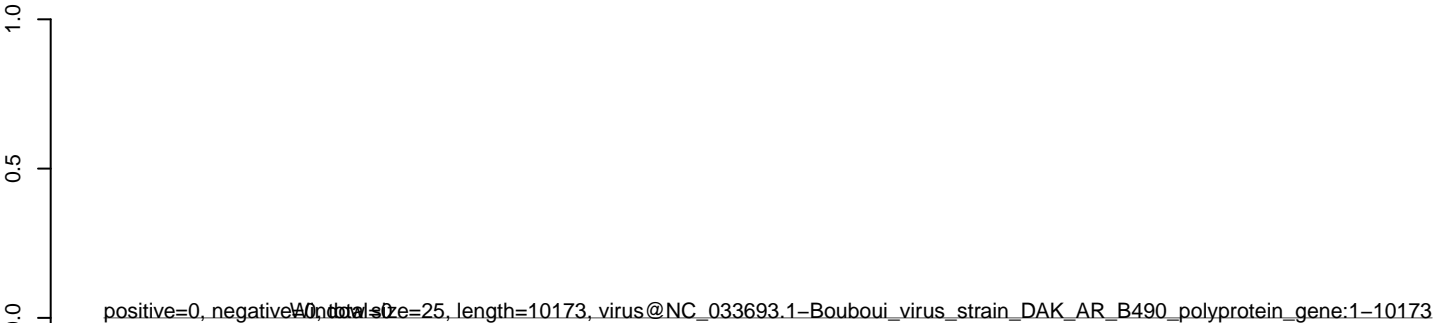
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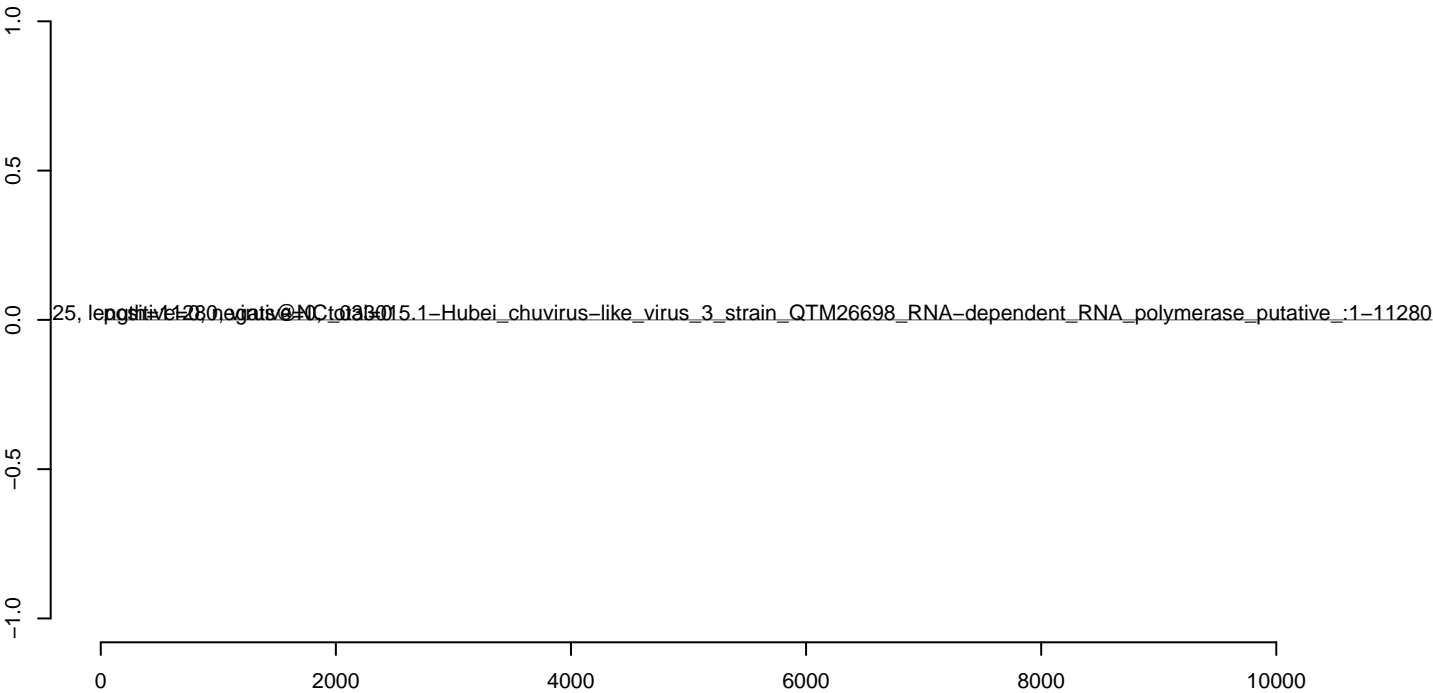
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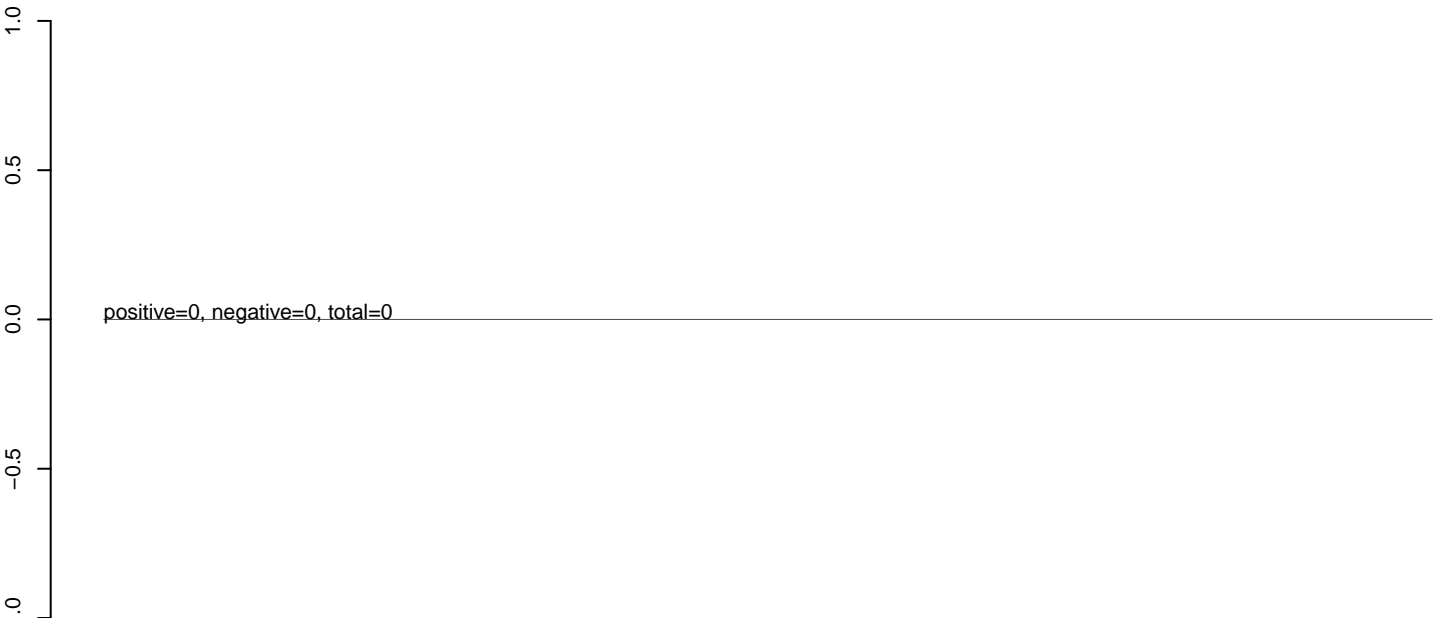
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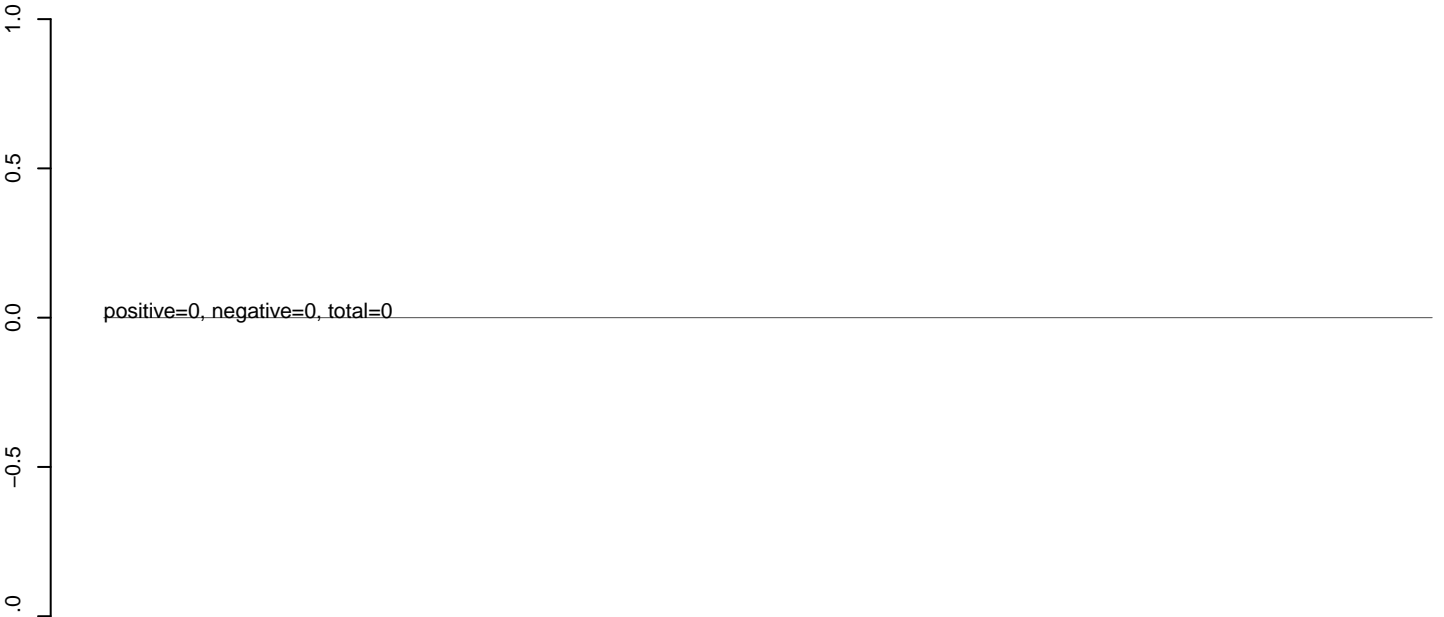
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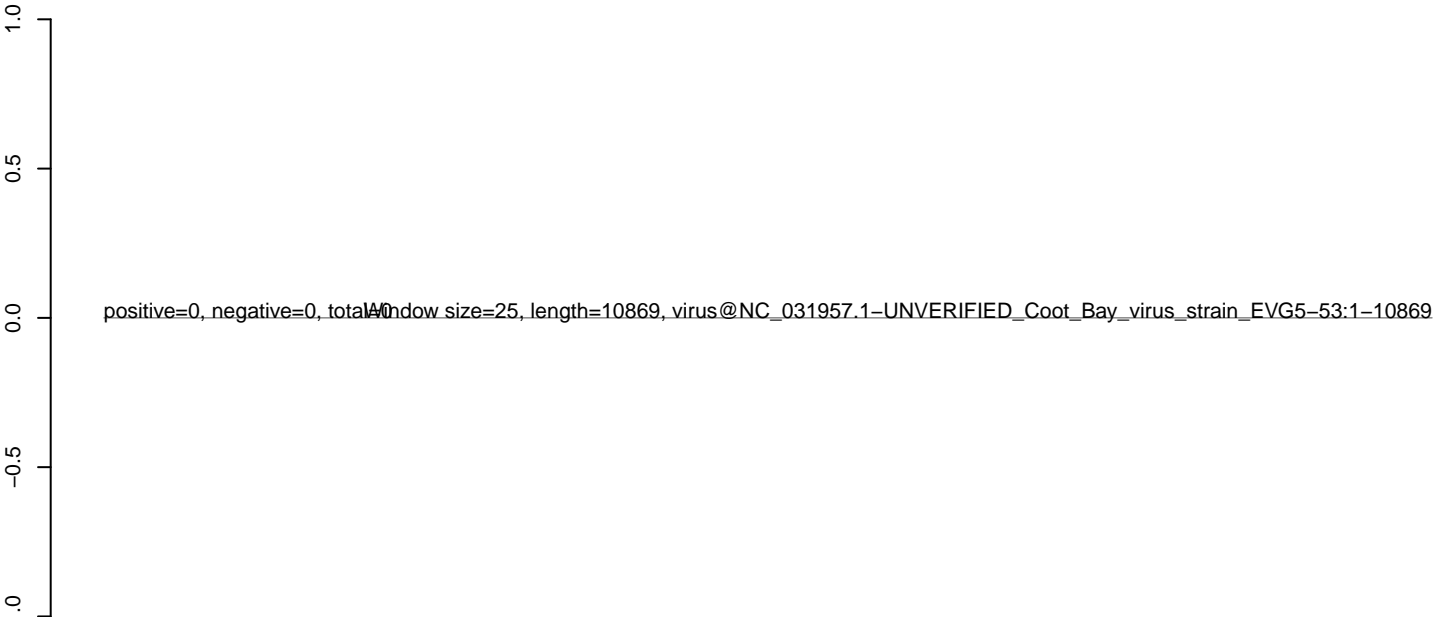
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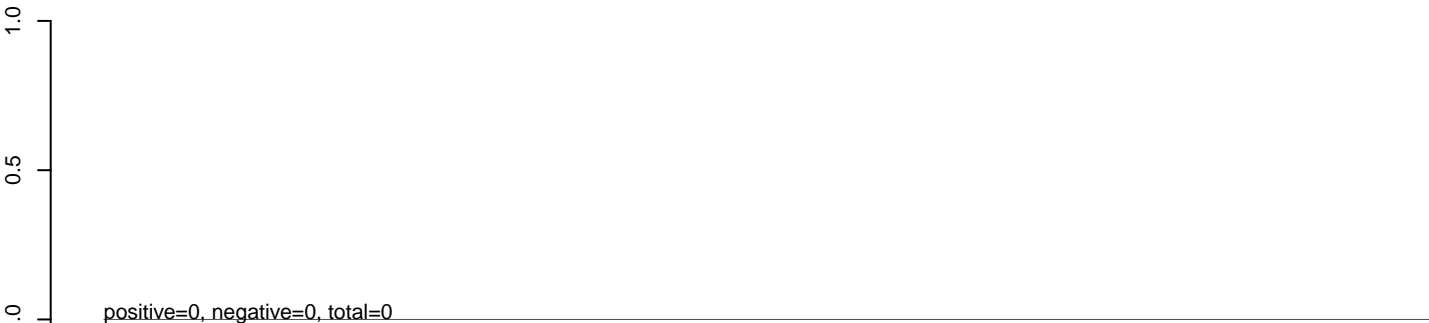


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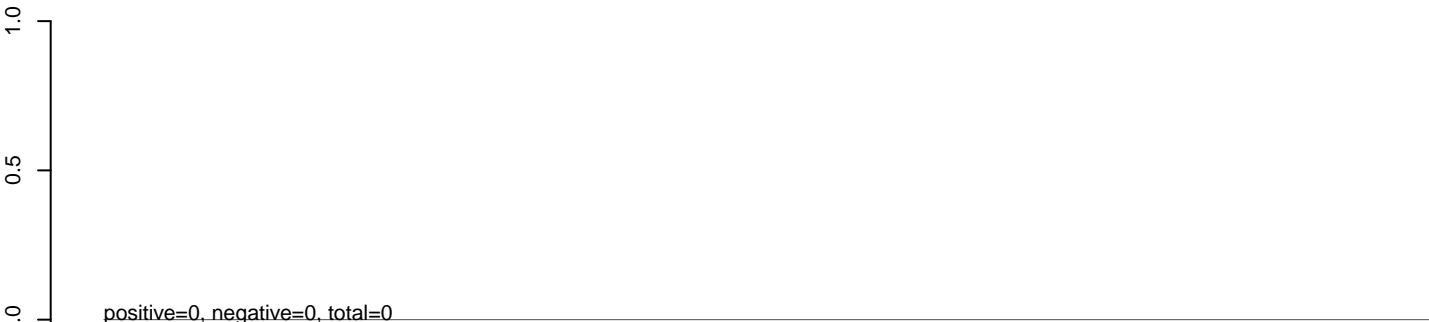


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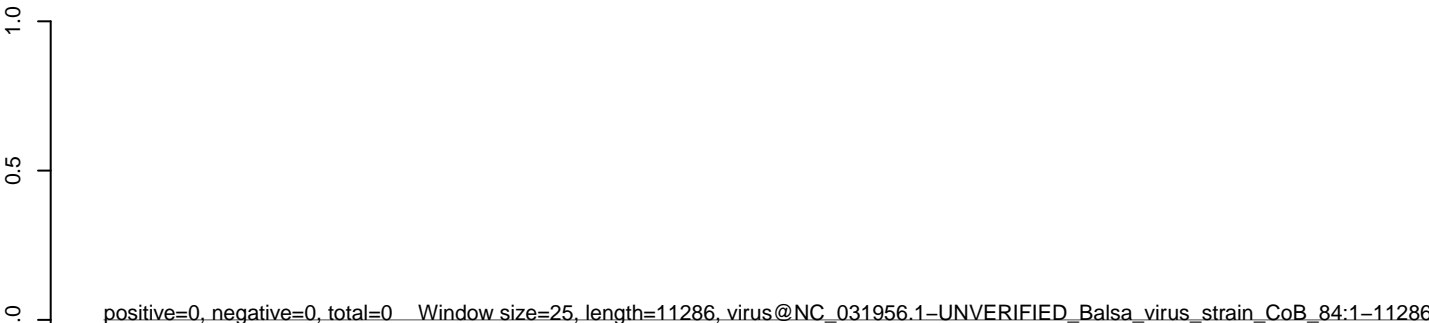
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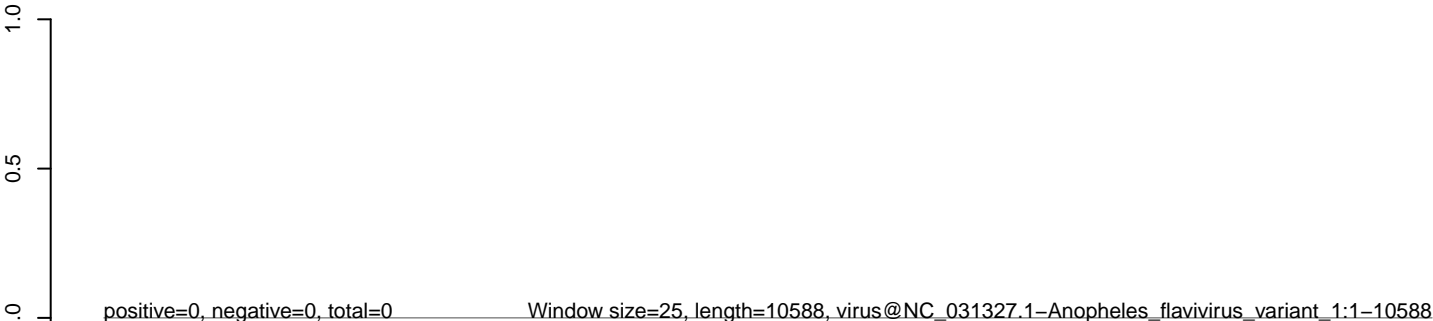
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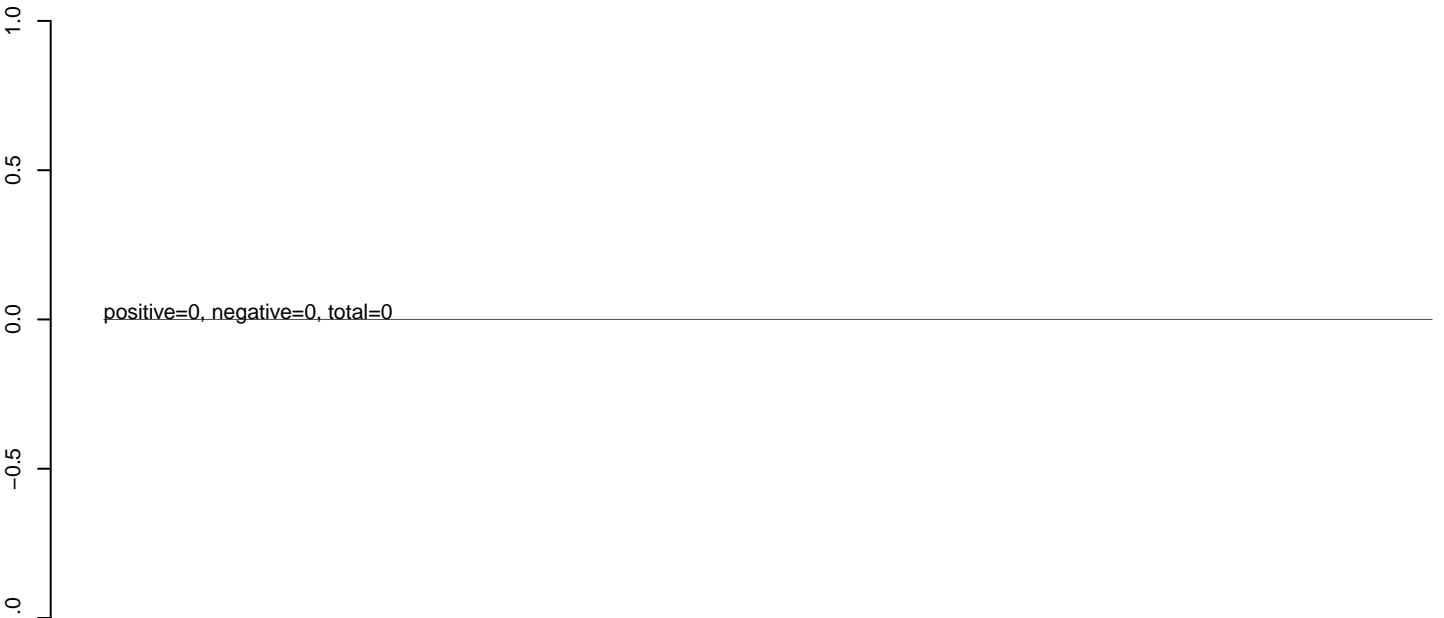


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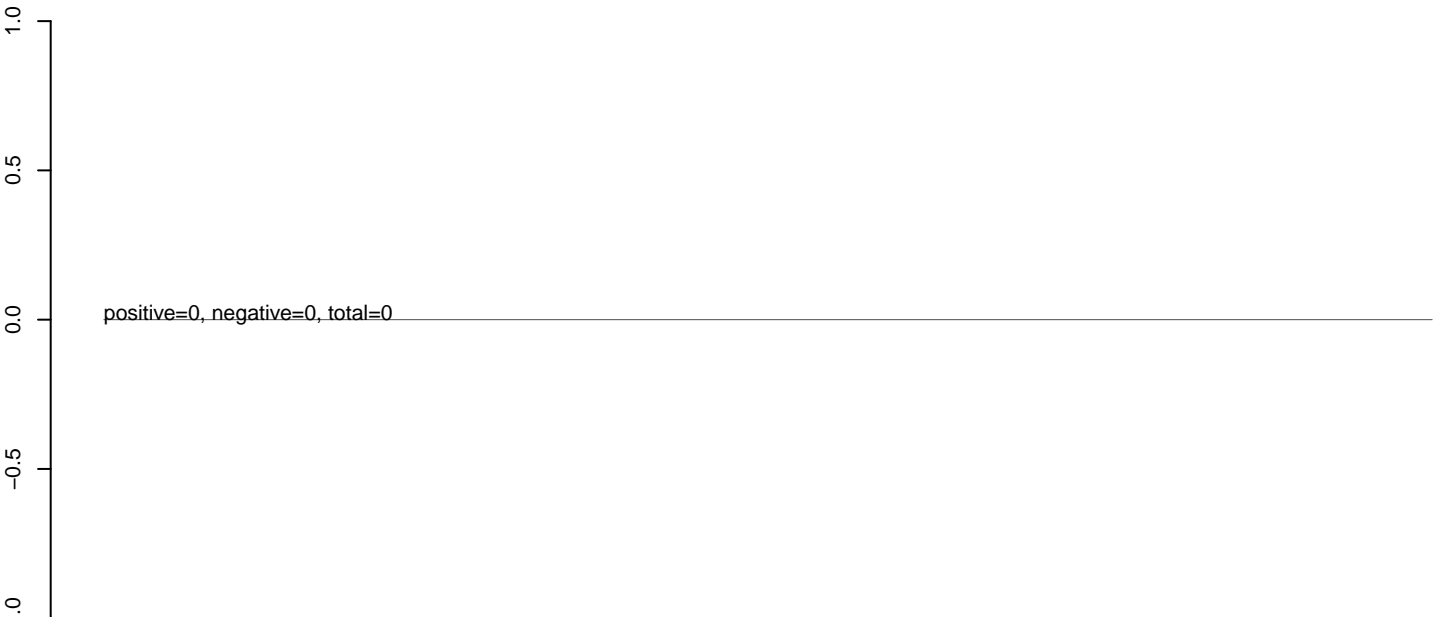


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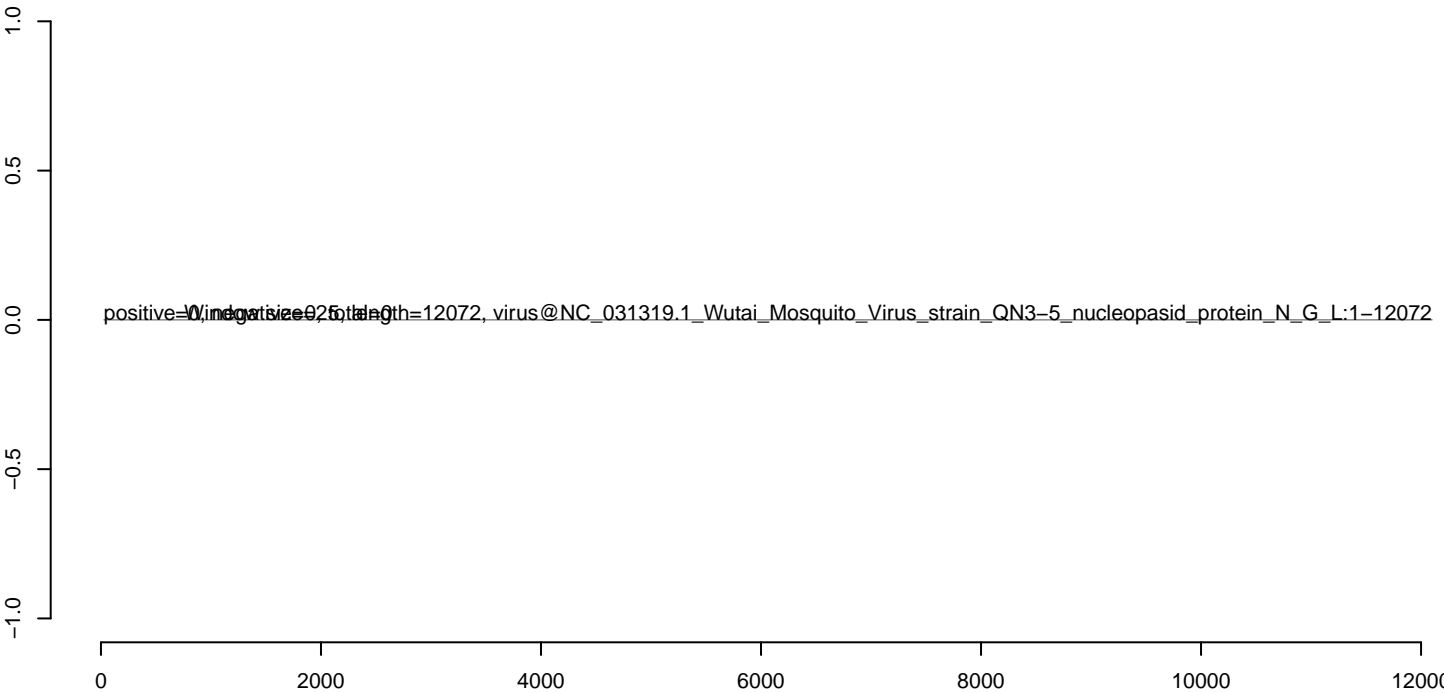
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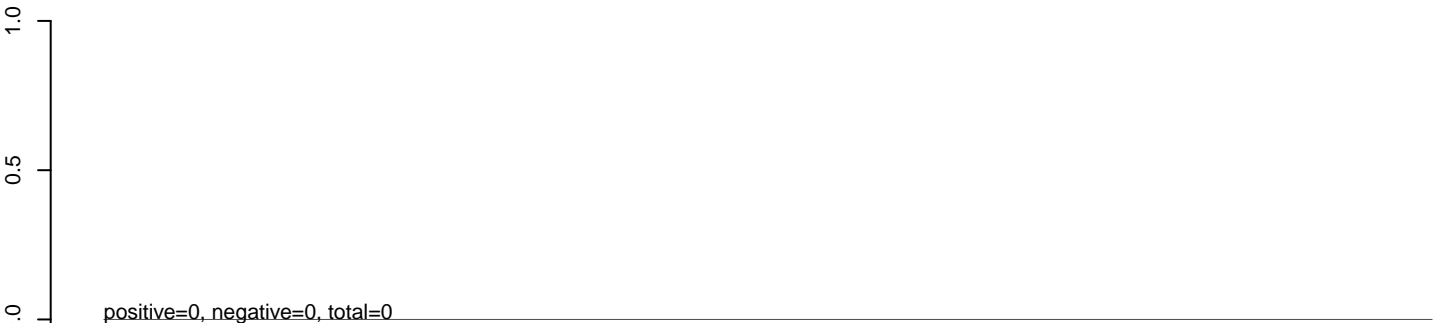
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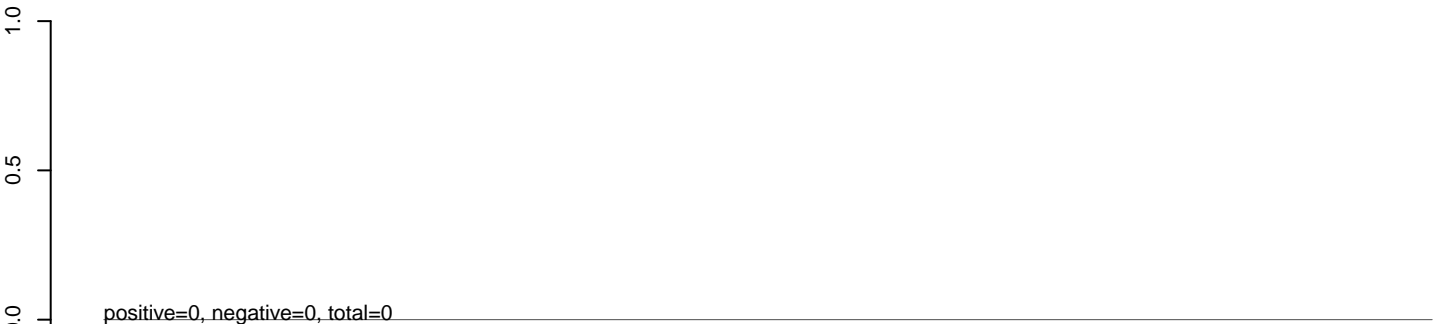
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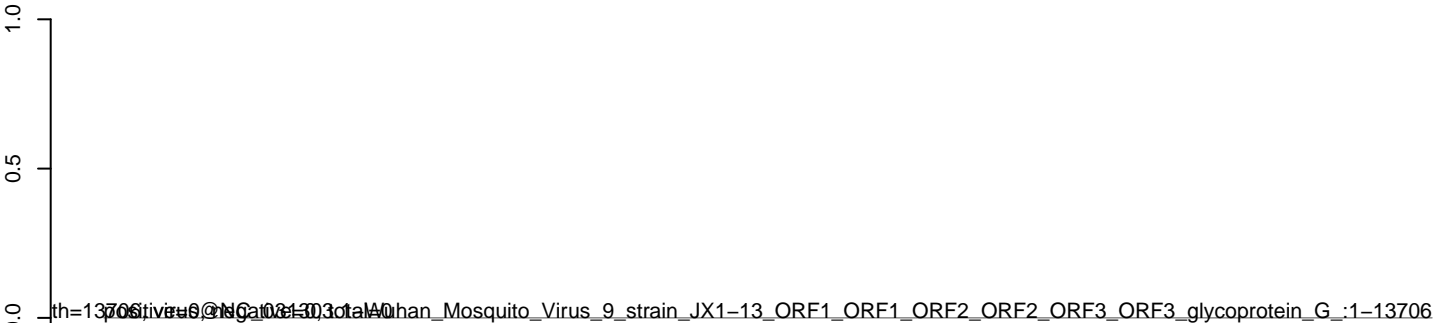
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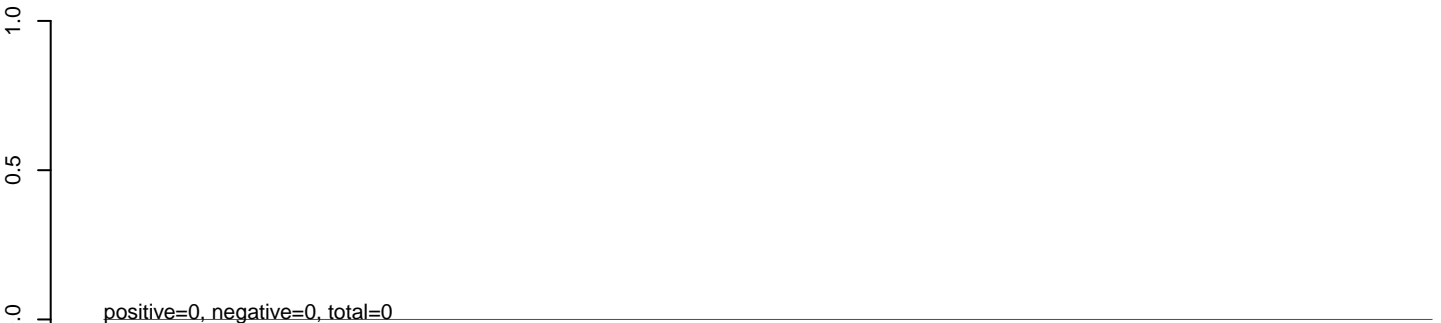


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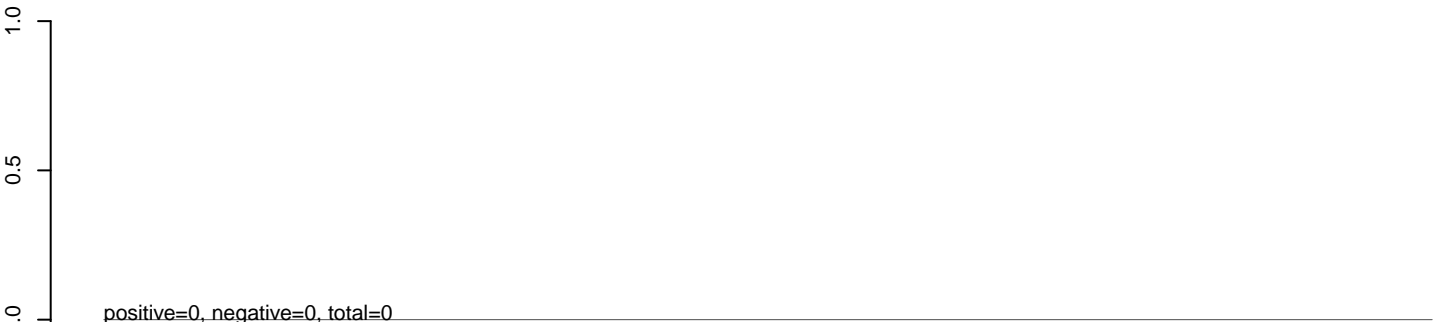


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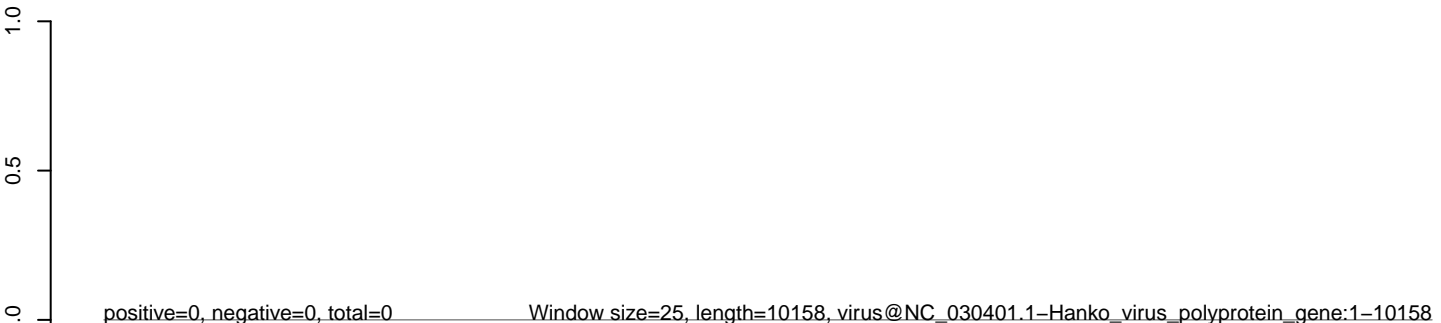
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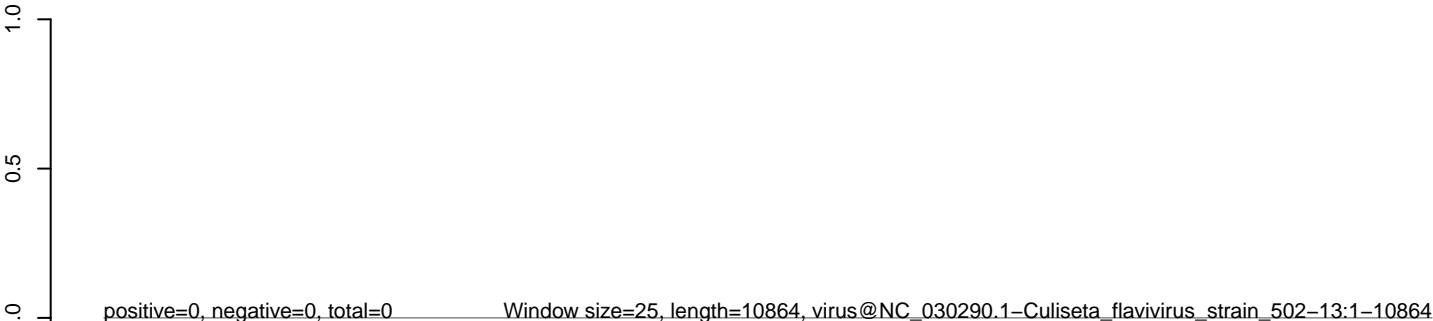
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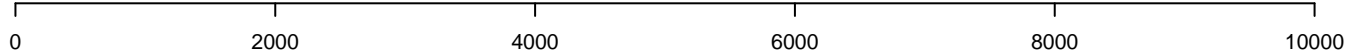
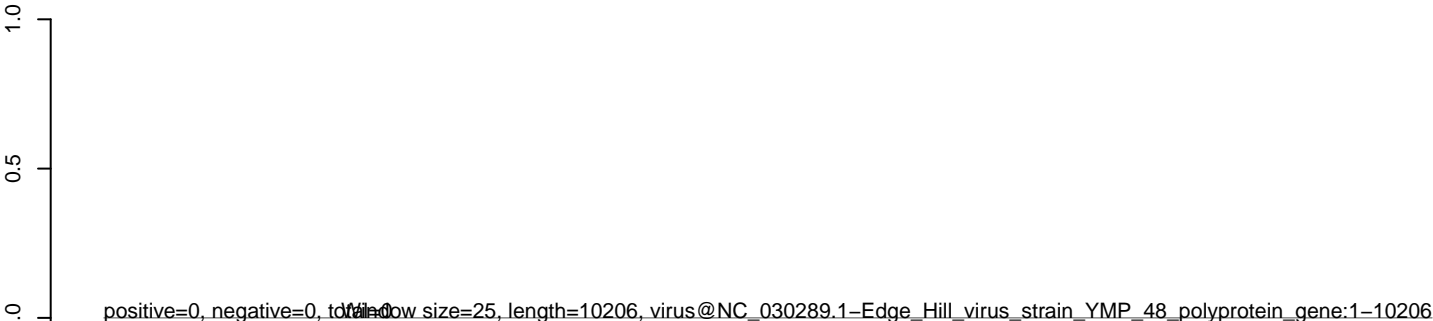
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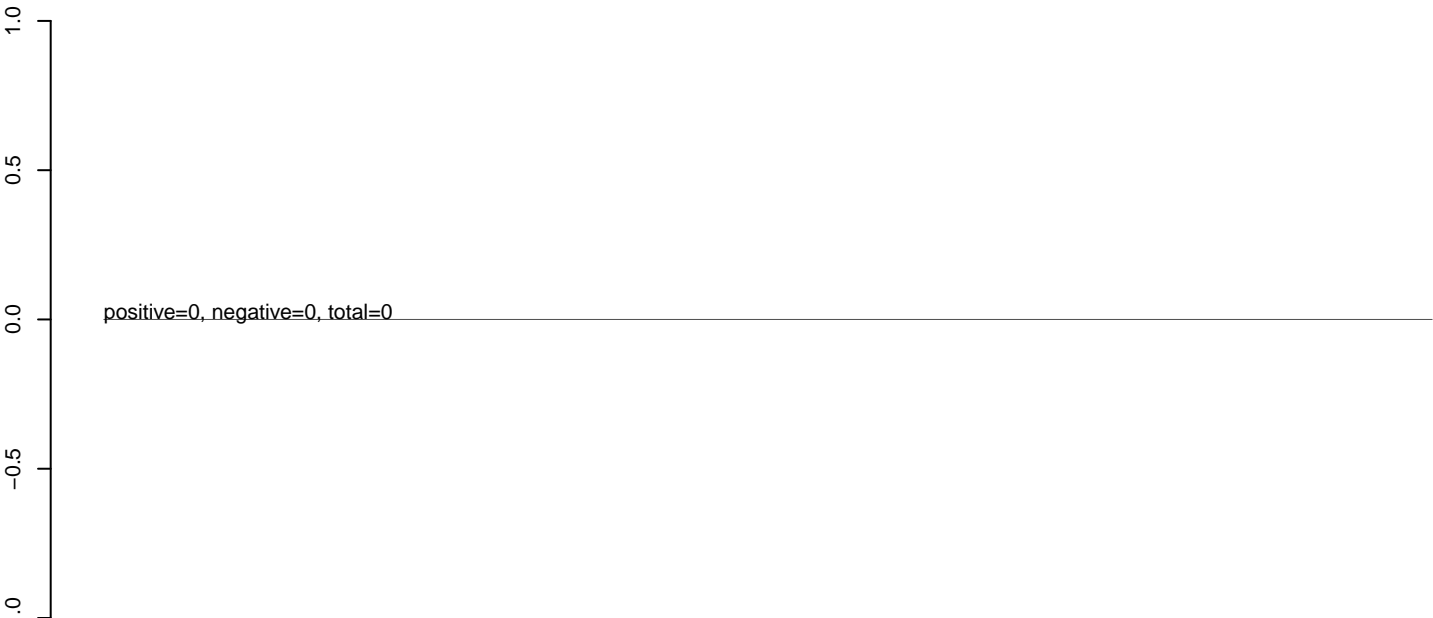
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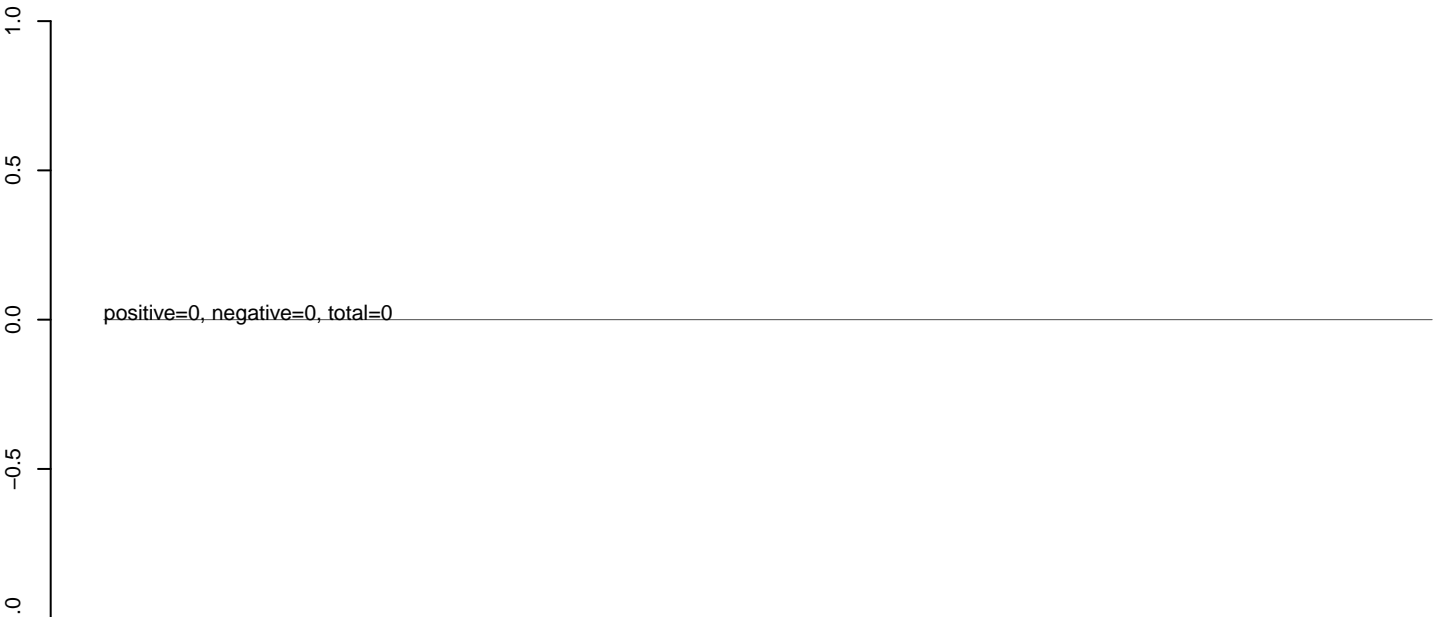
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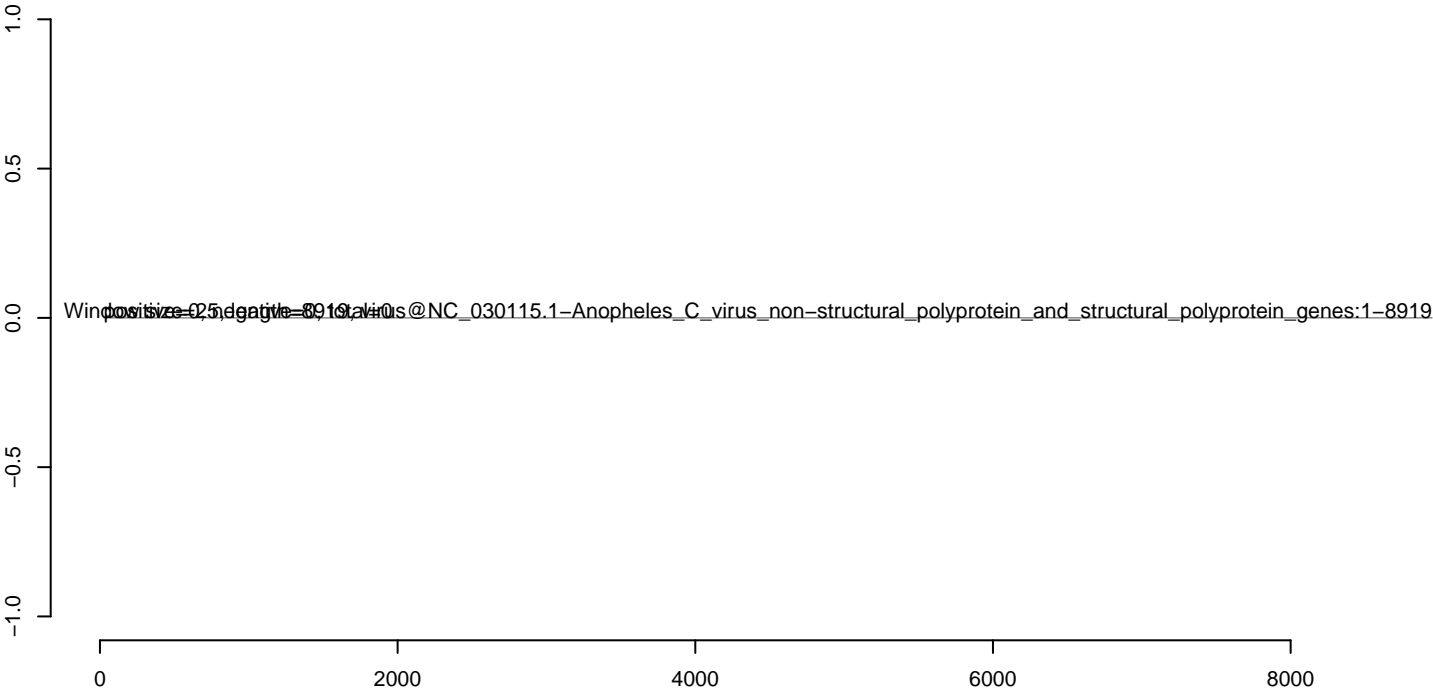
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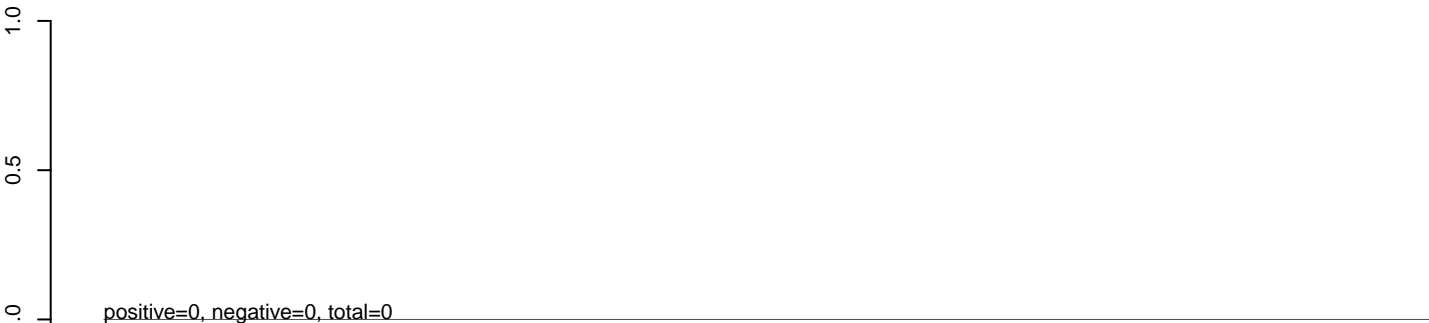
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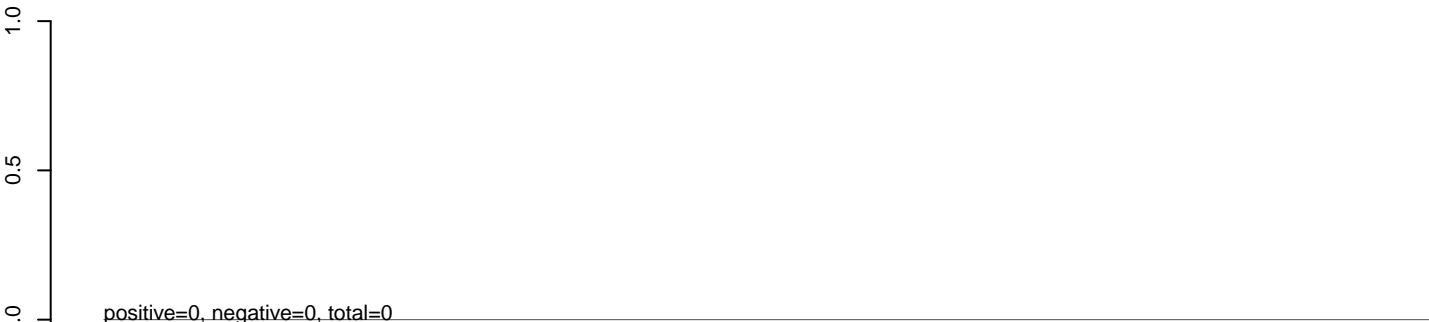
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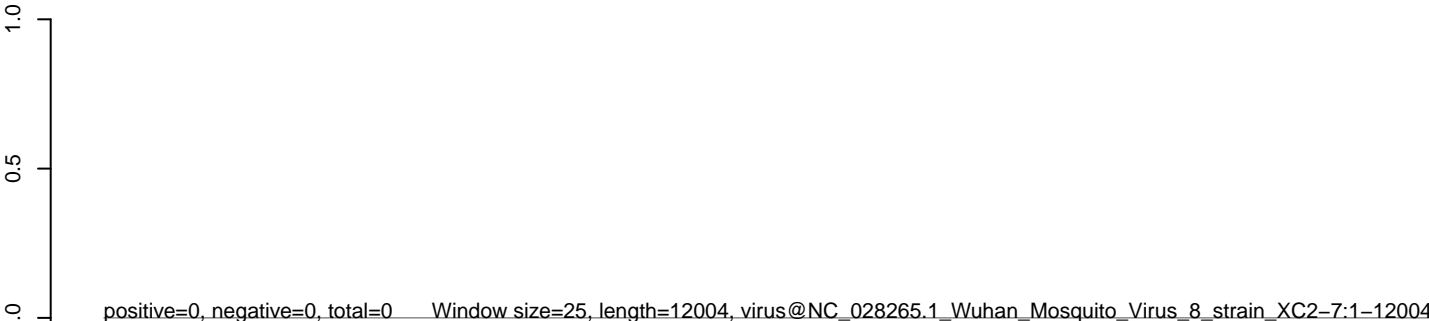
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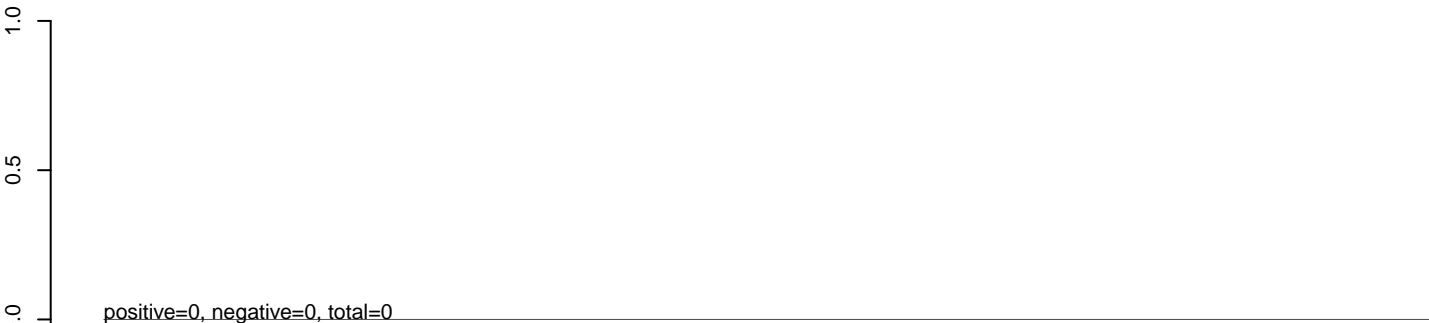
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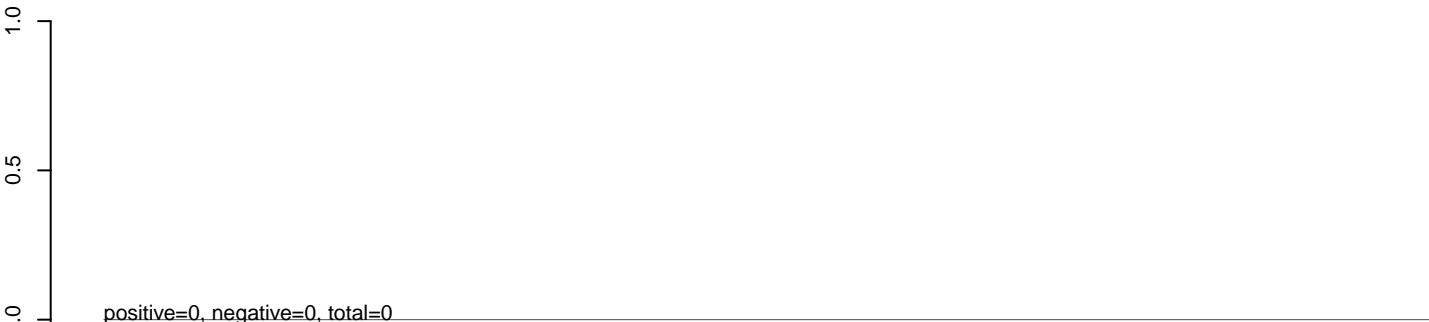
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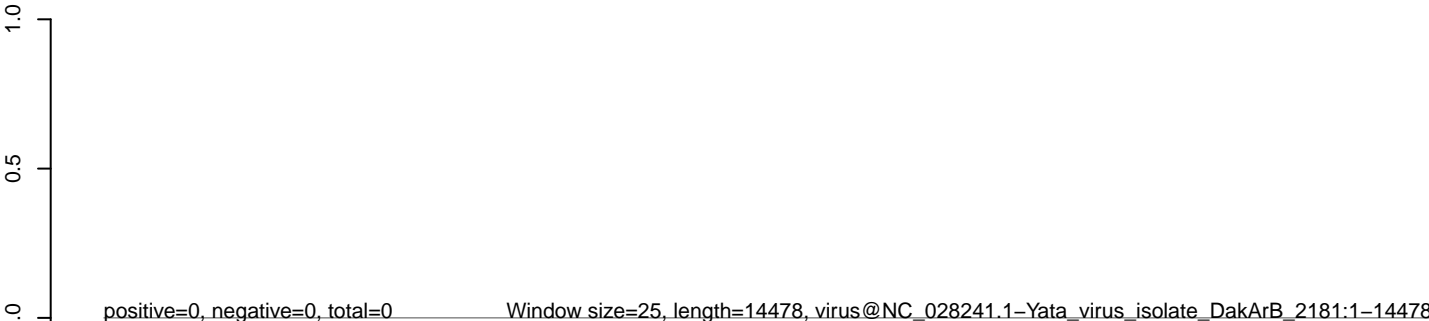
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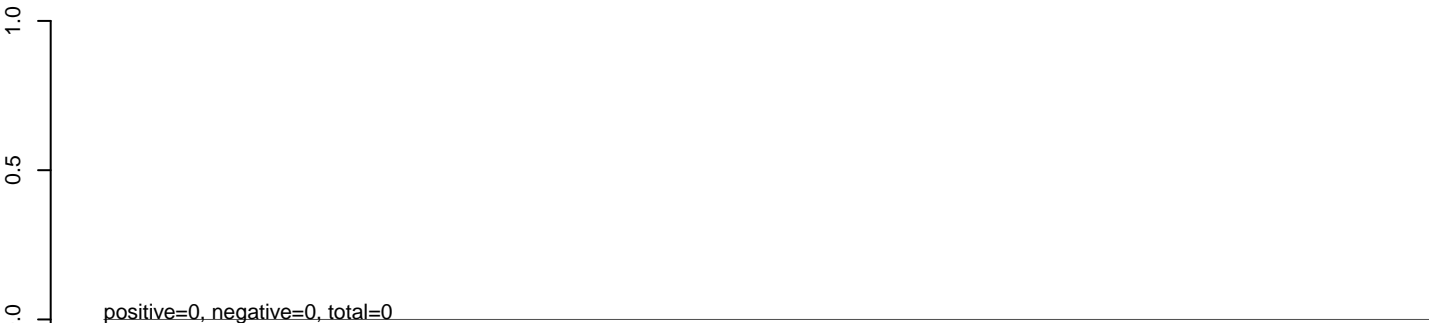
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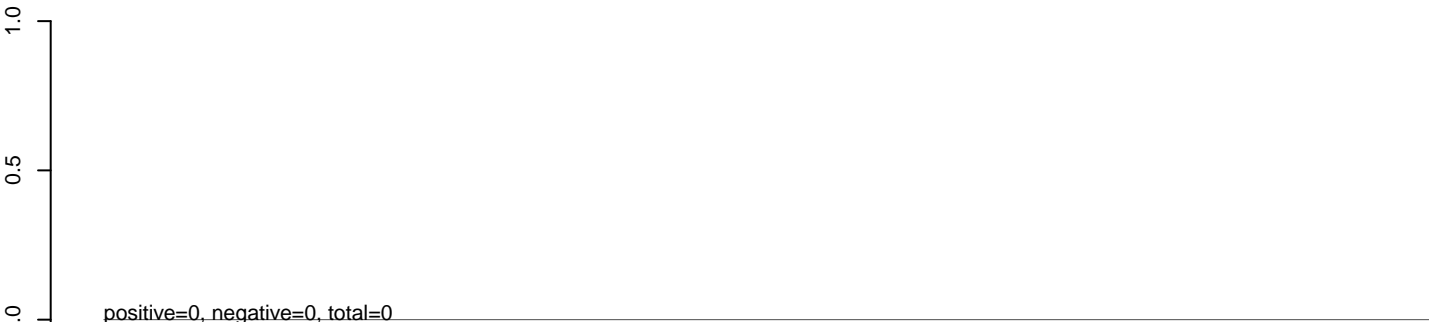
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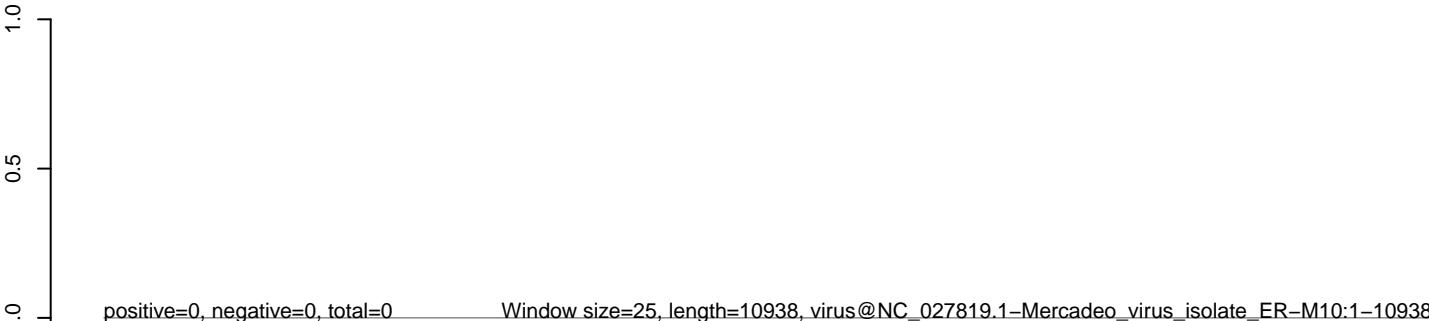
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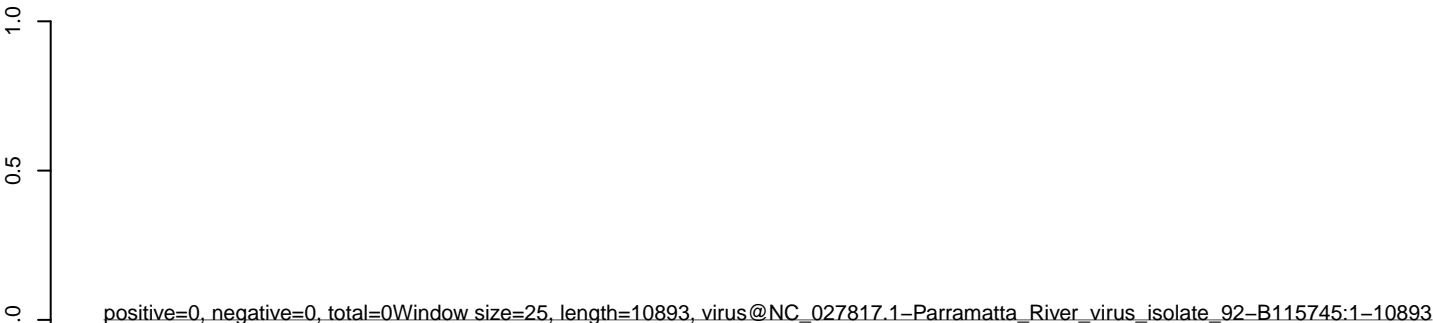
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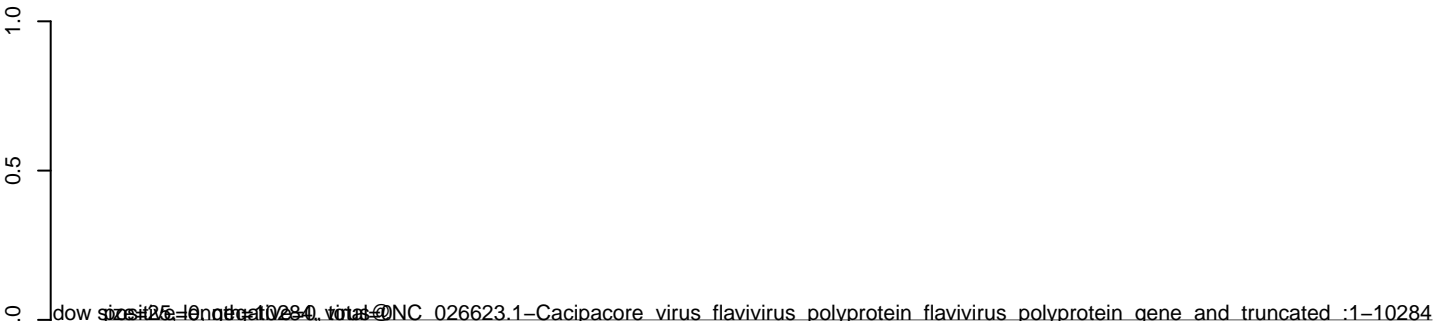
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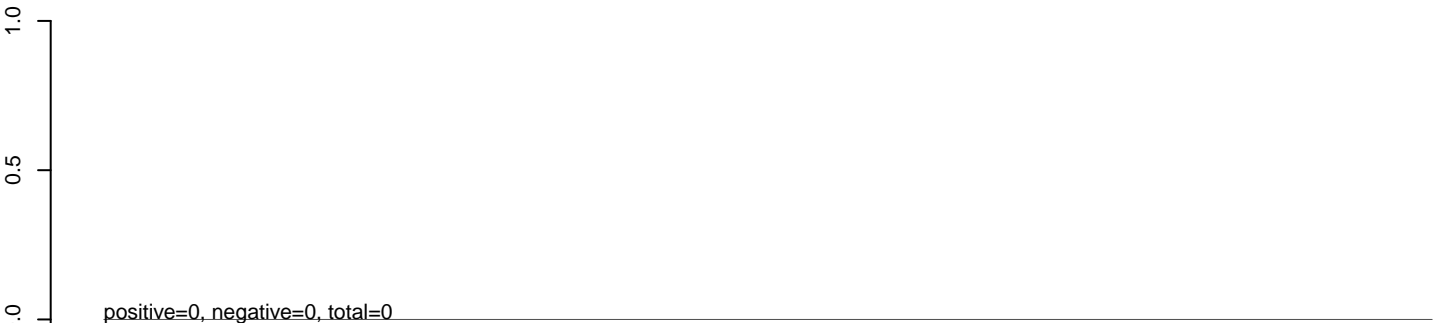
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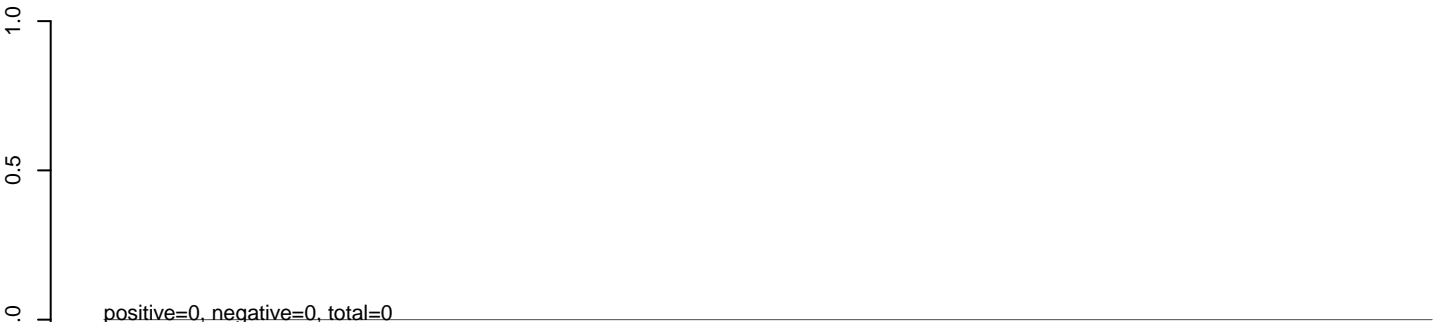
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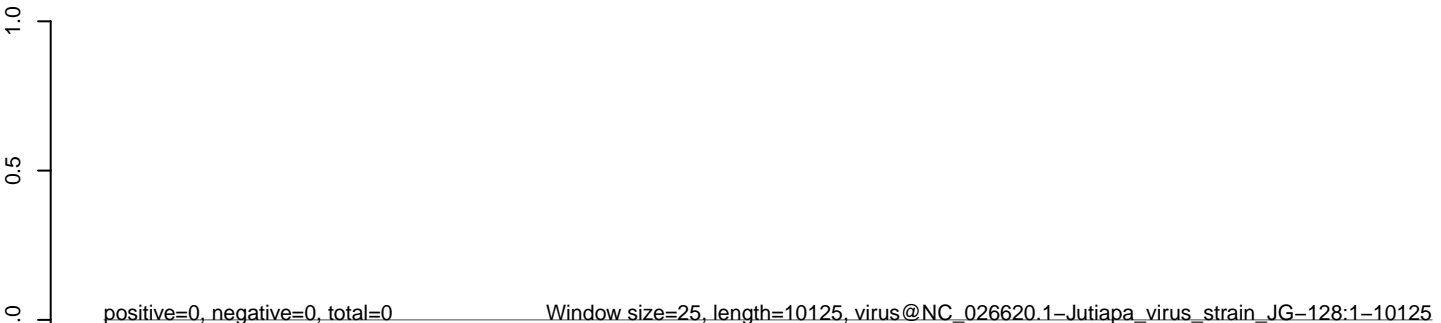
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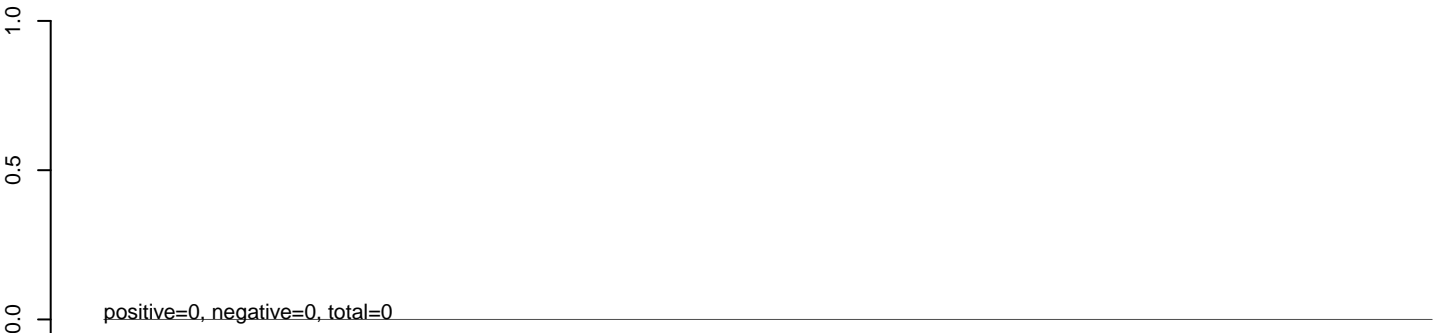


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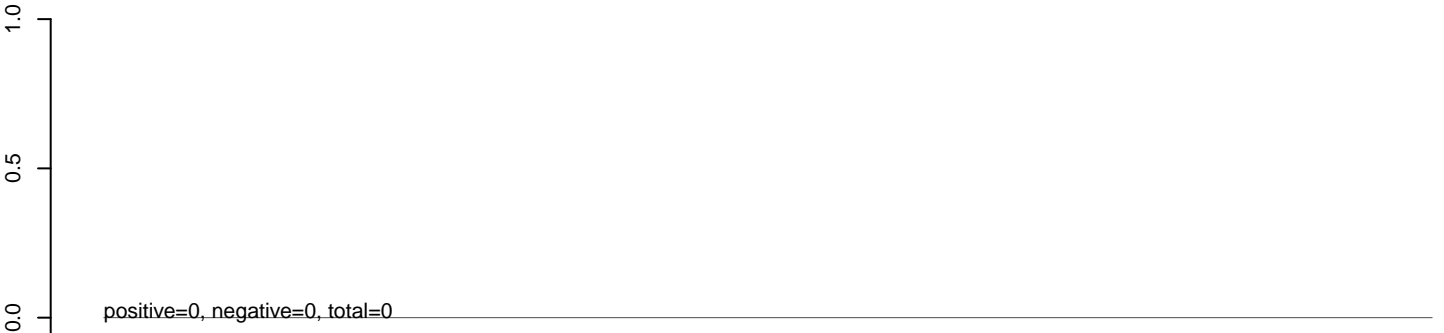


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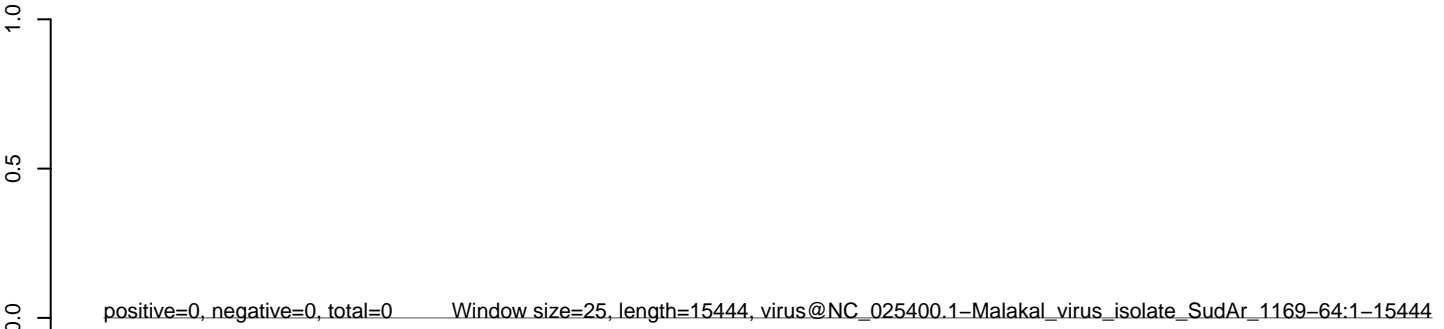
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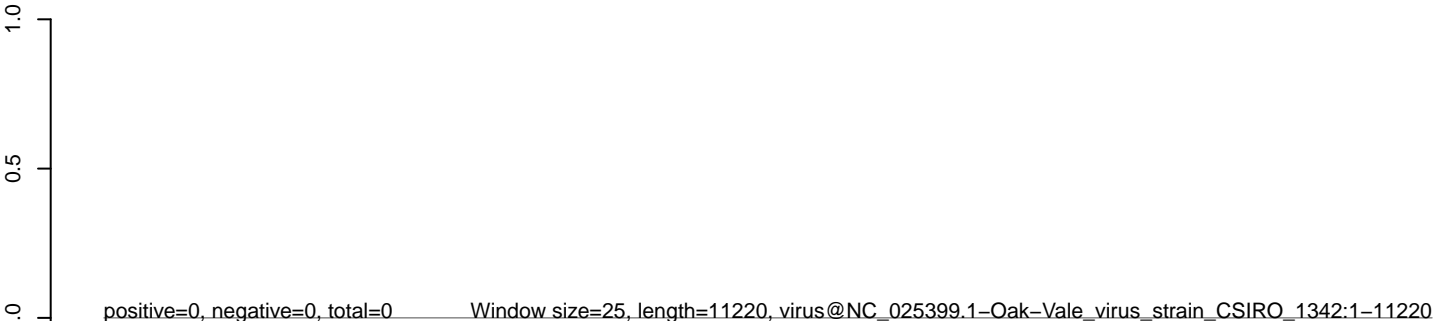
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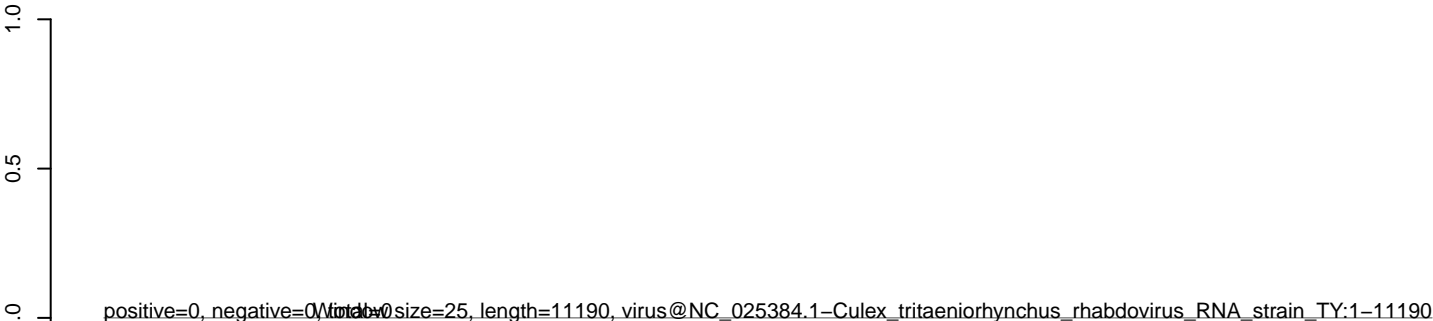
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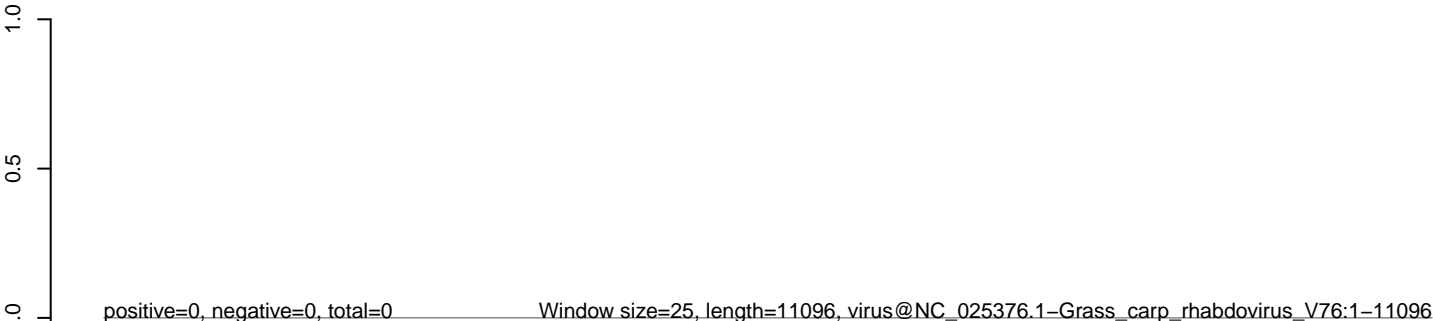
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AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



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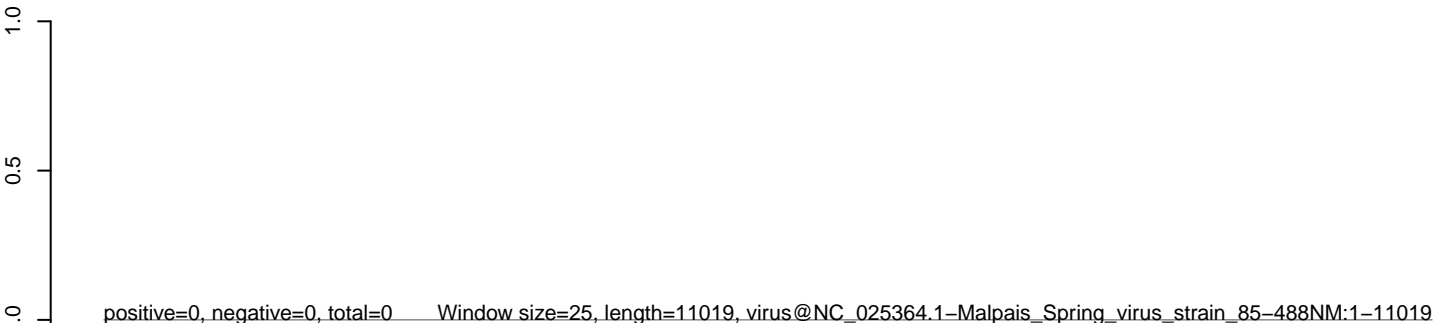
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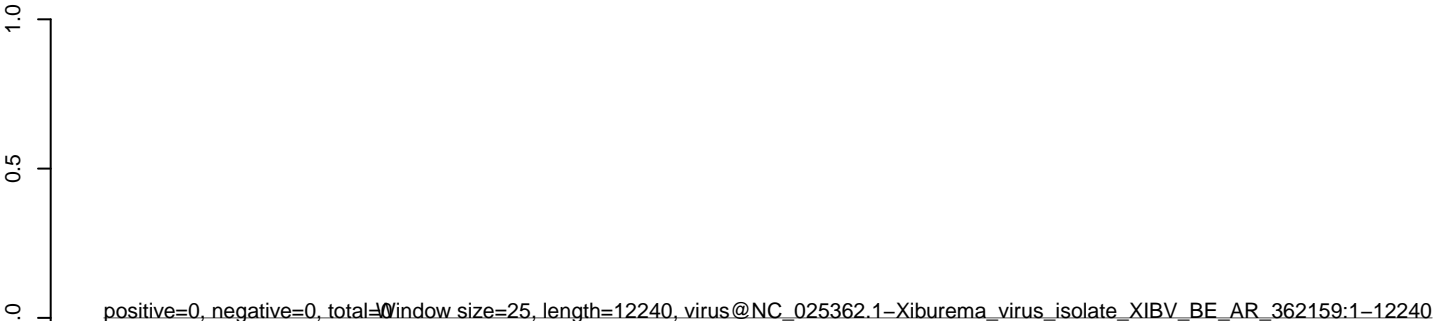
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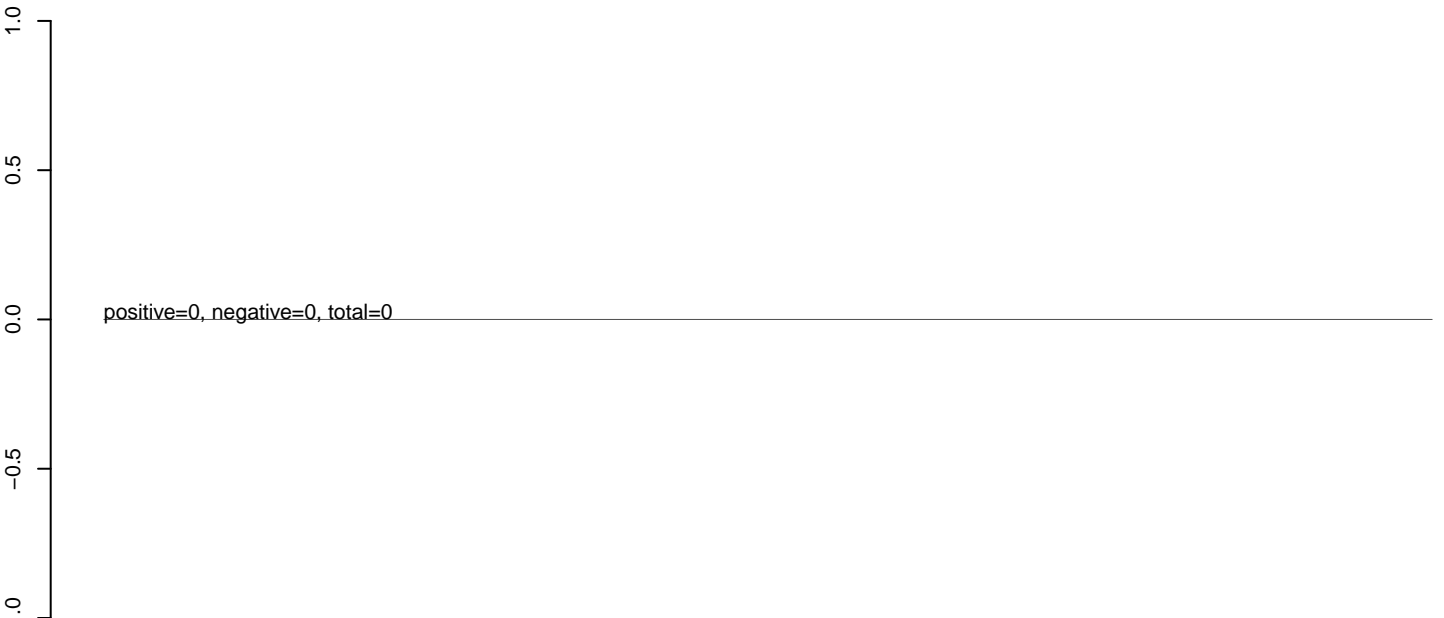
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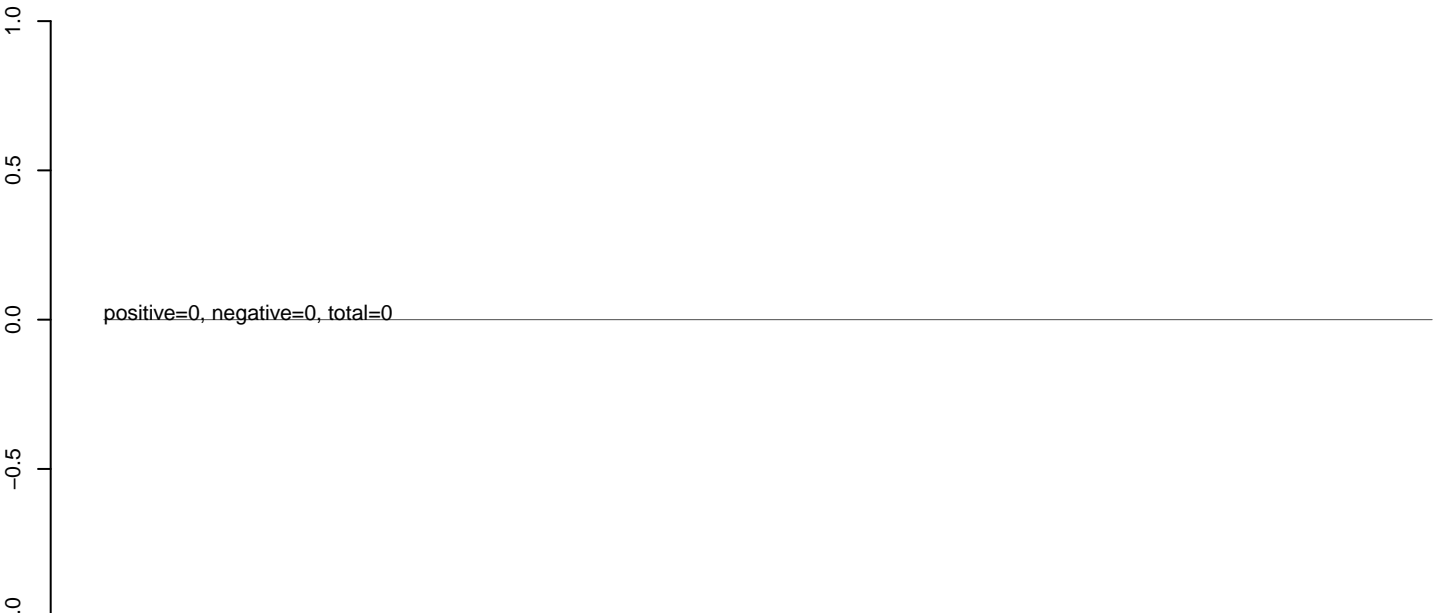
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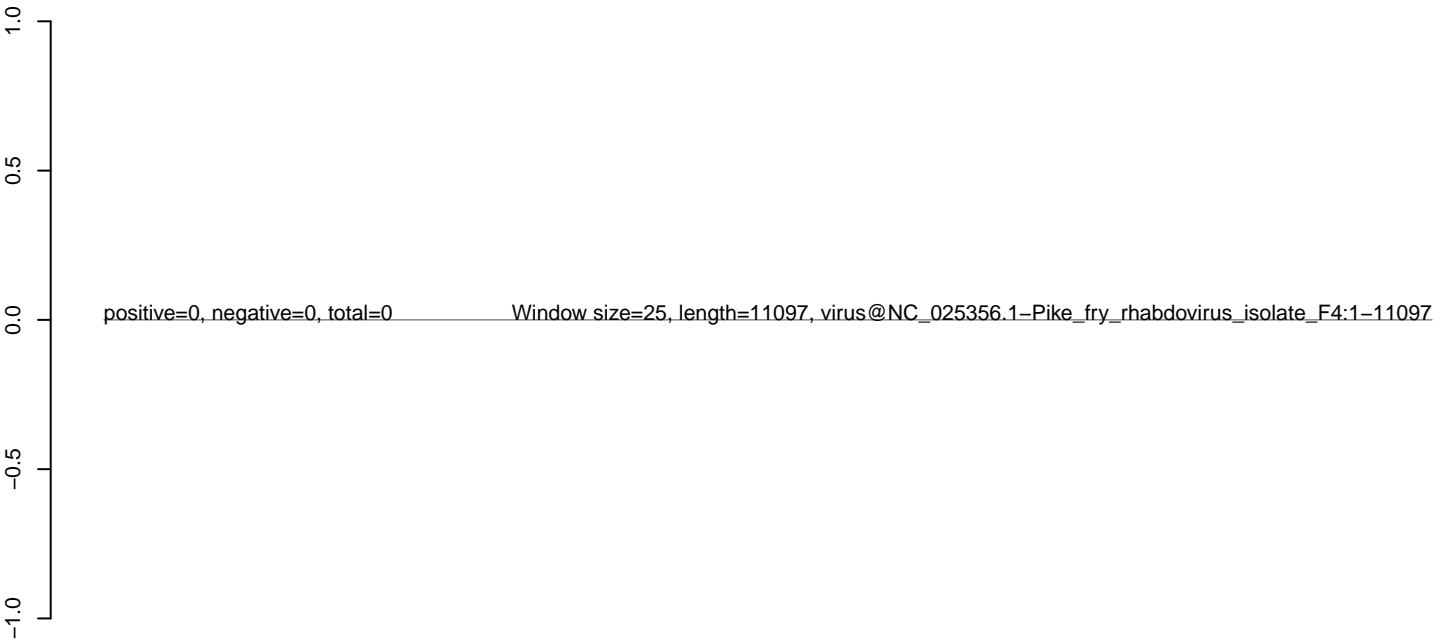
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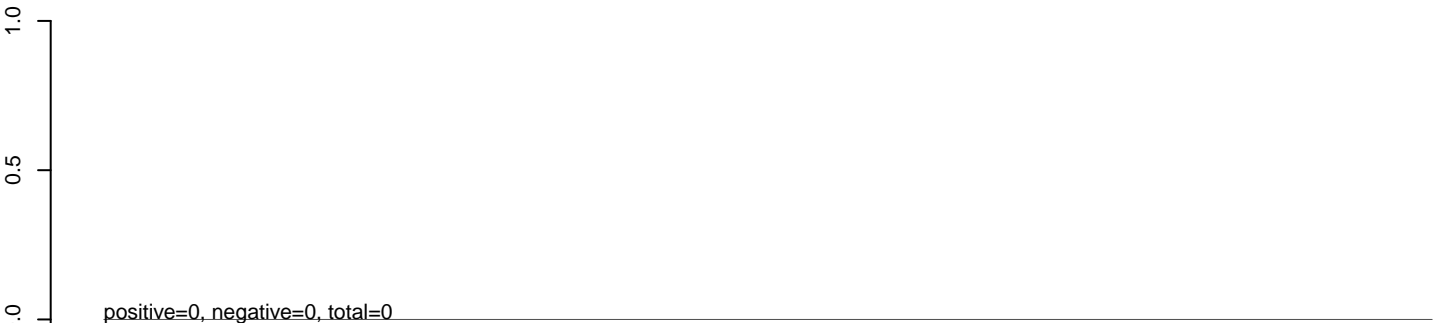
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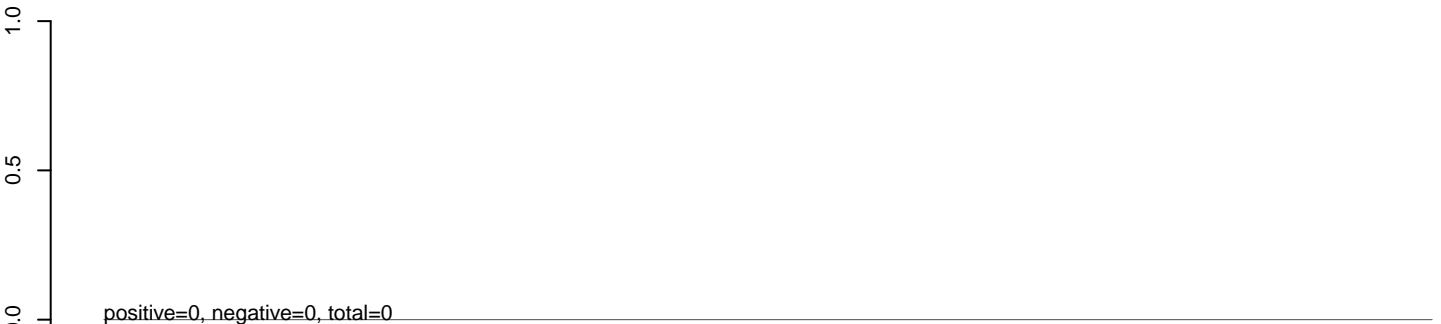
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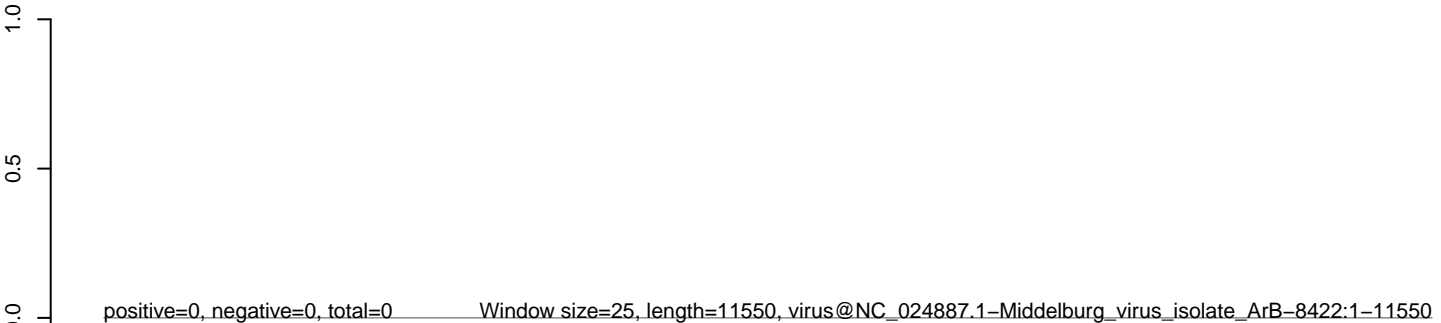
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AnGam_Ovary_BetaE.rep



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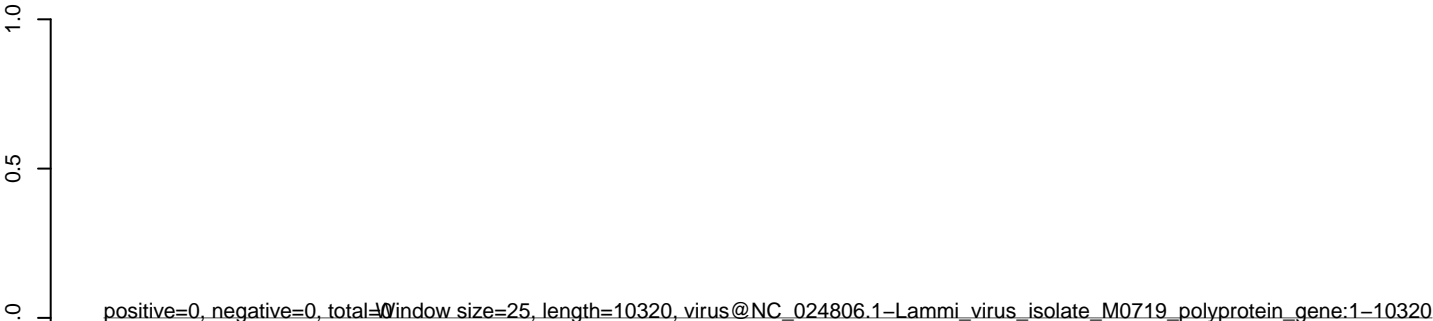
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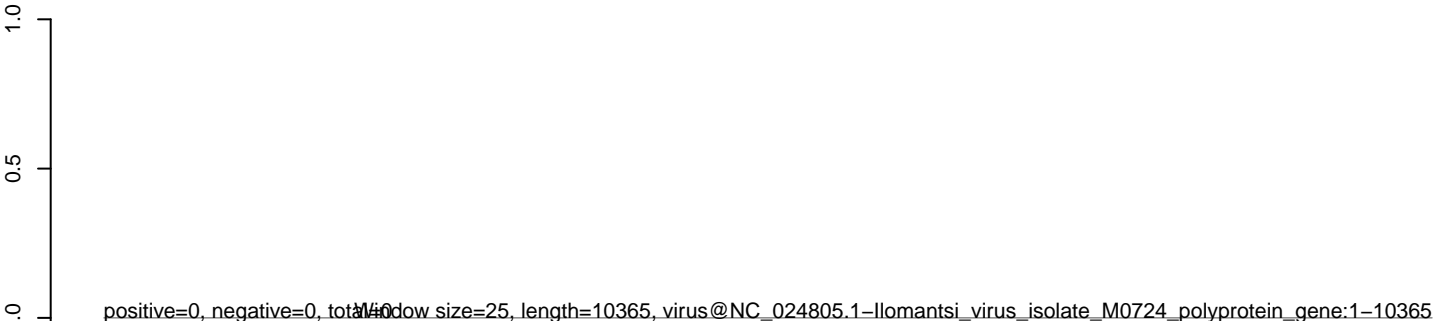
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AnGam_Ovary_BetaE.rep



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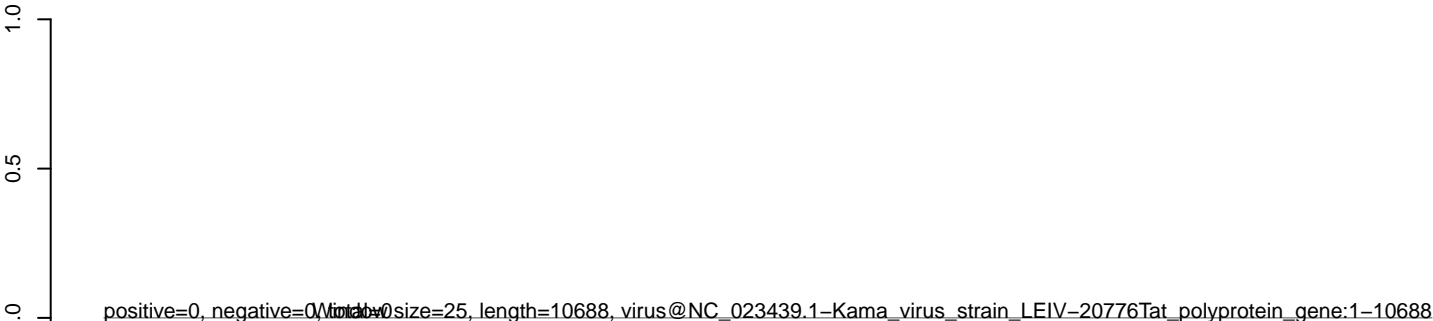
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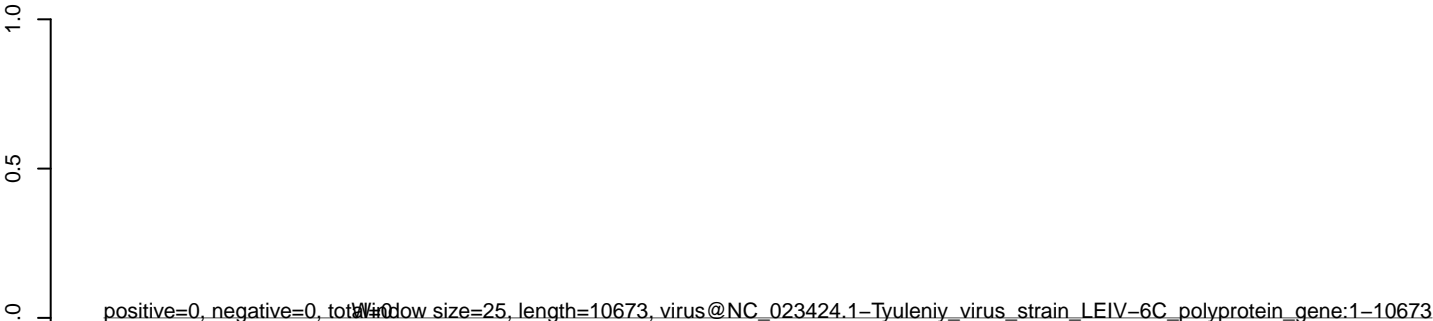
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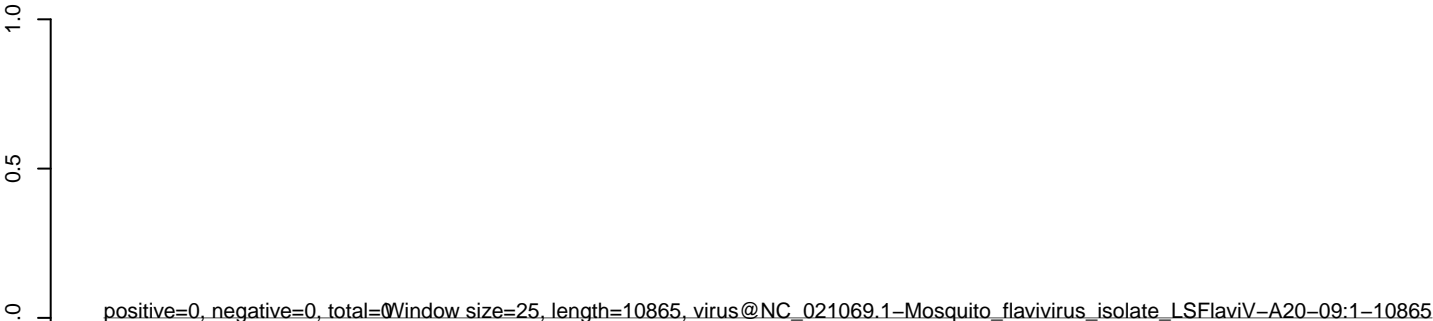
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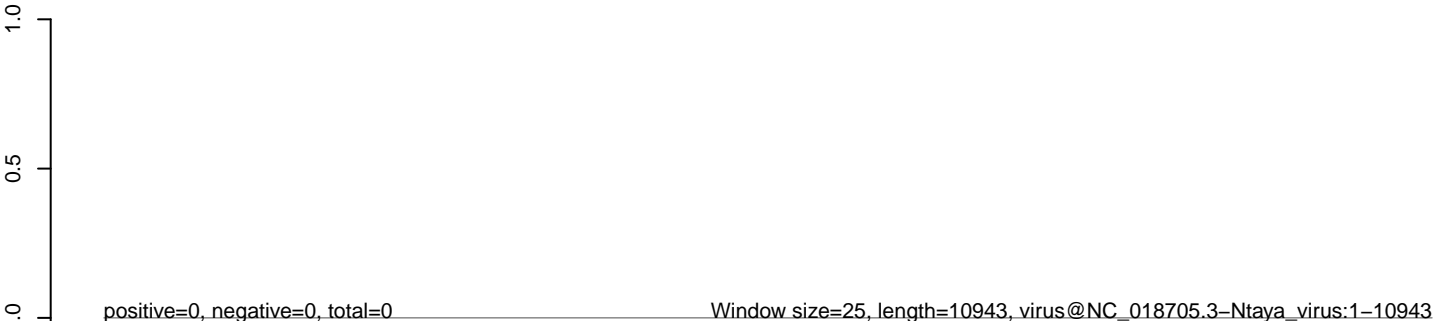
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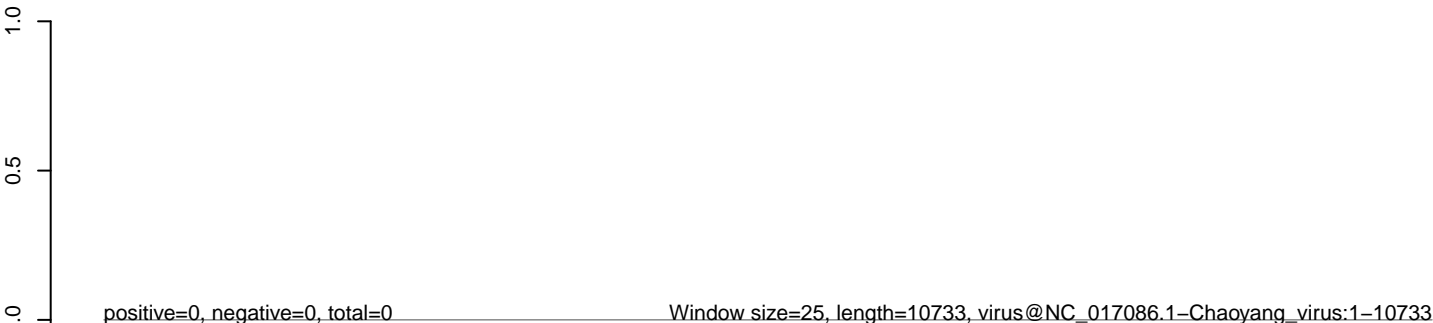
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AnGam_Ovary_BetaE.rep



0 2000 4000 6000 8000 10000

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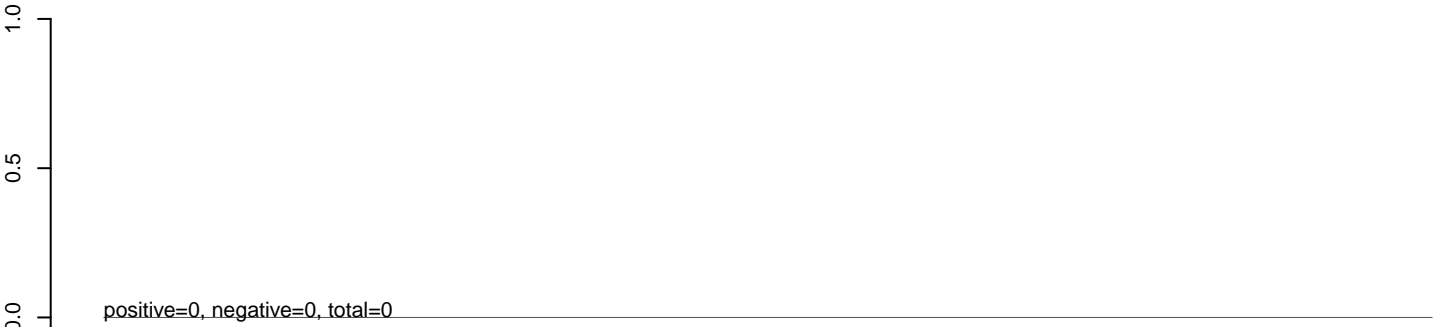


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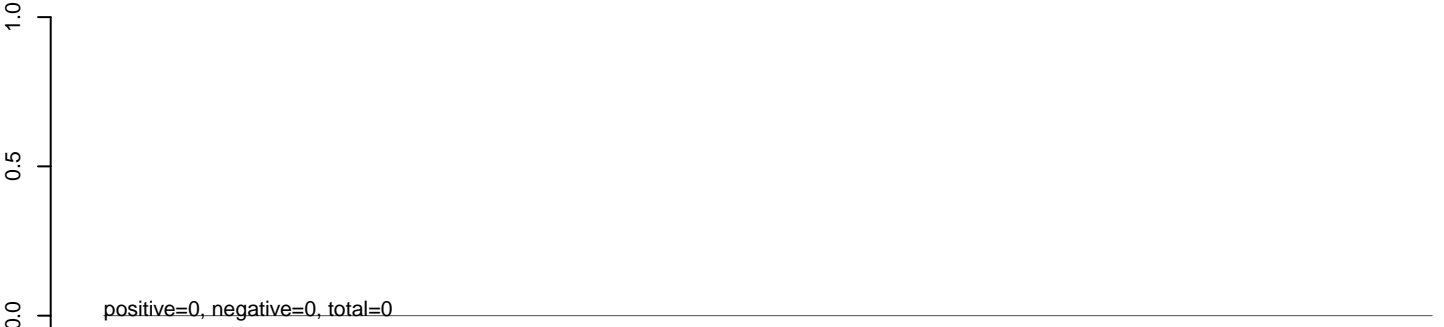


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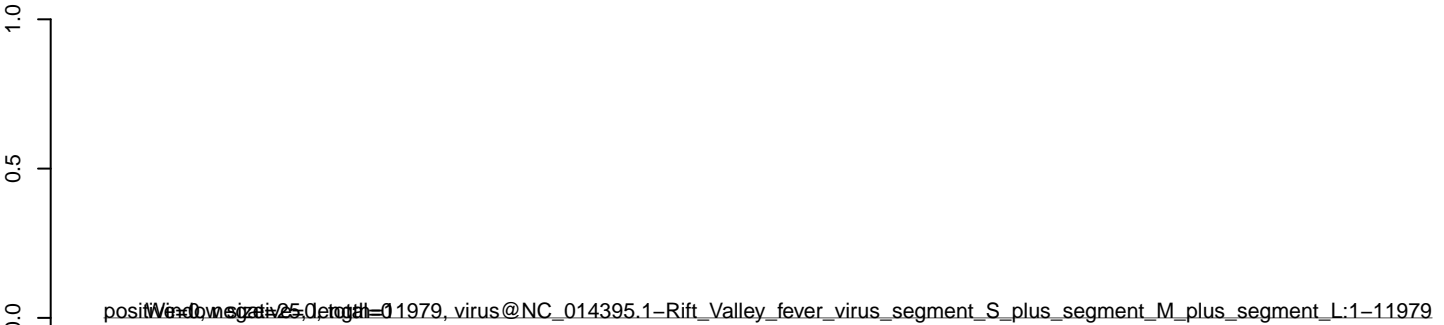
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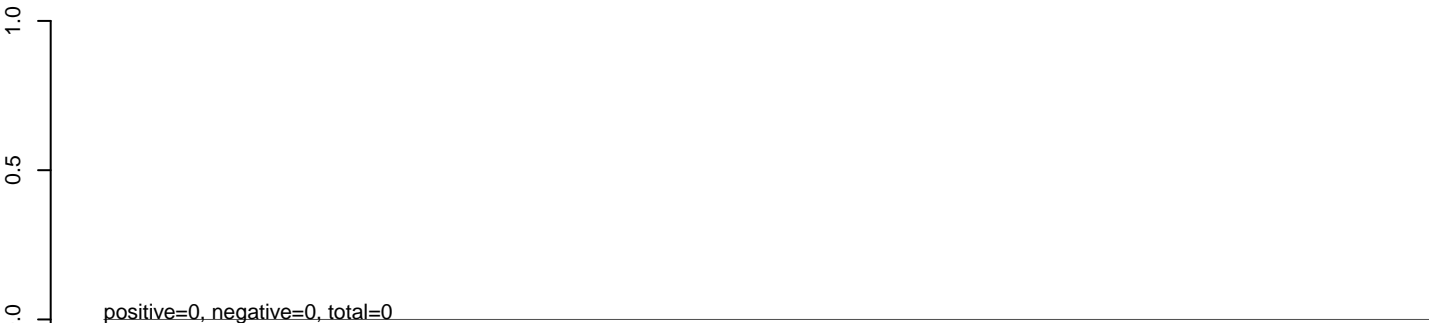
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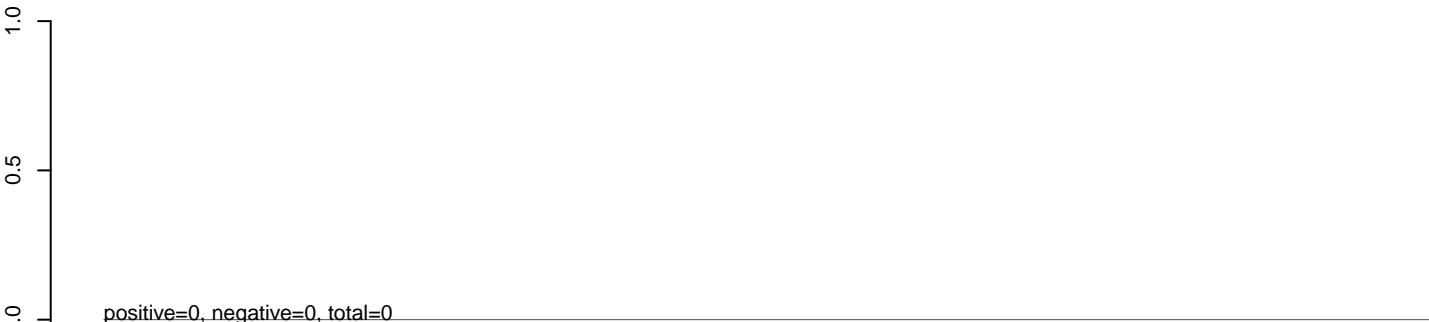
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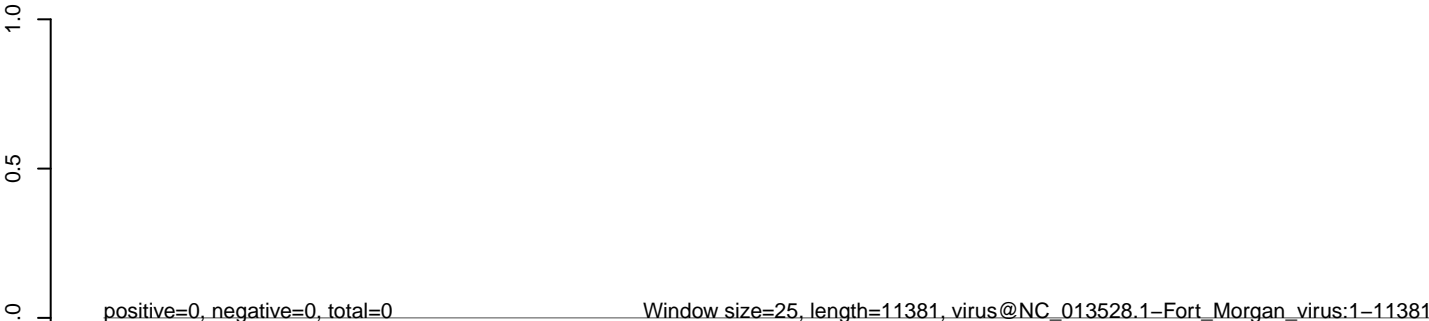
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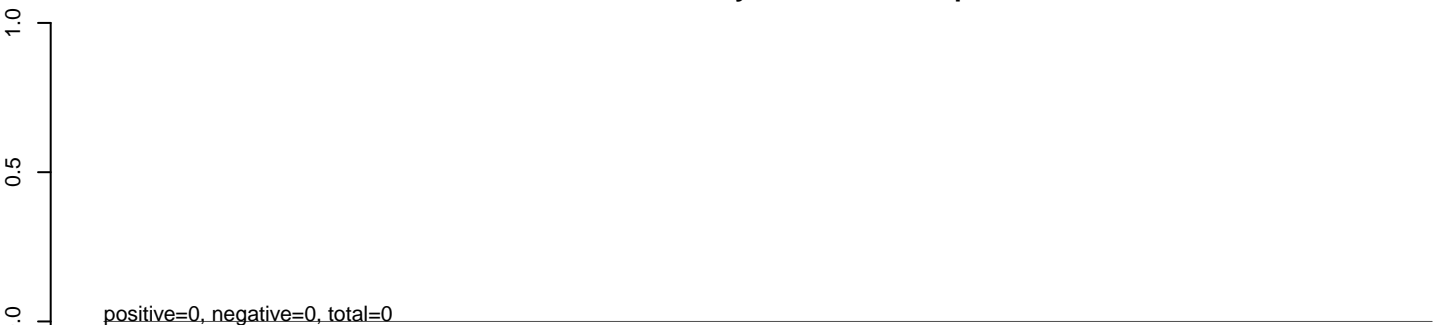


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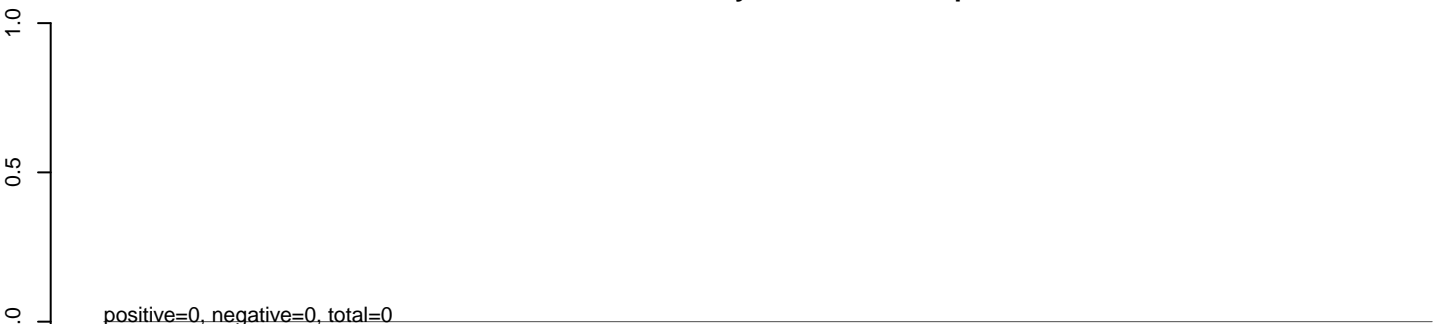


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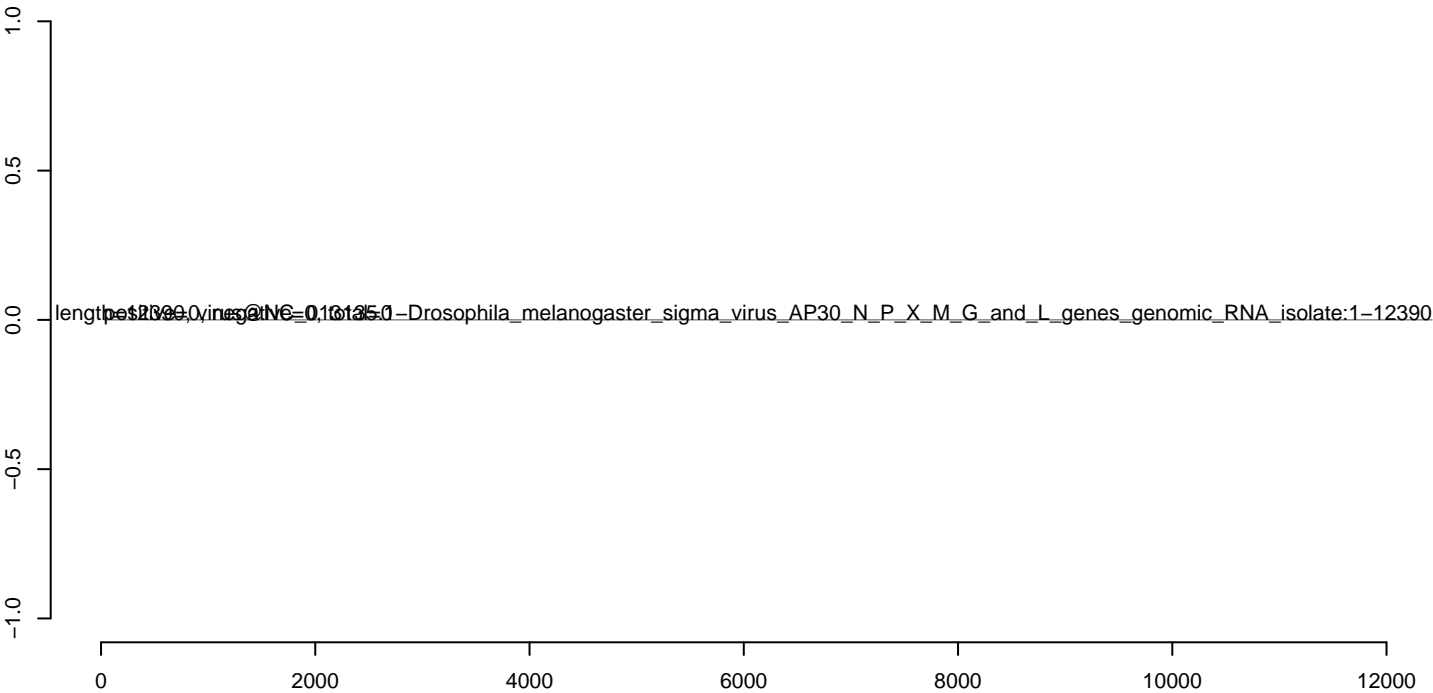
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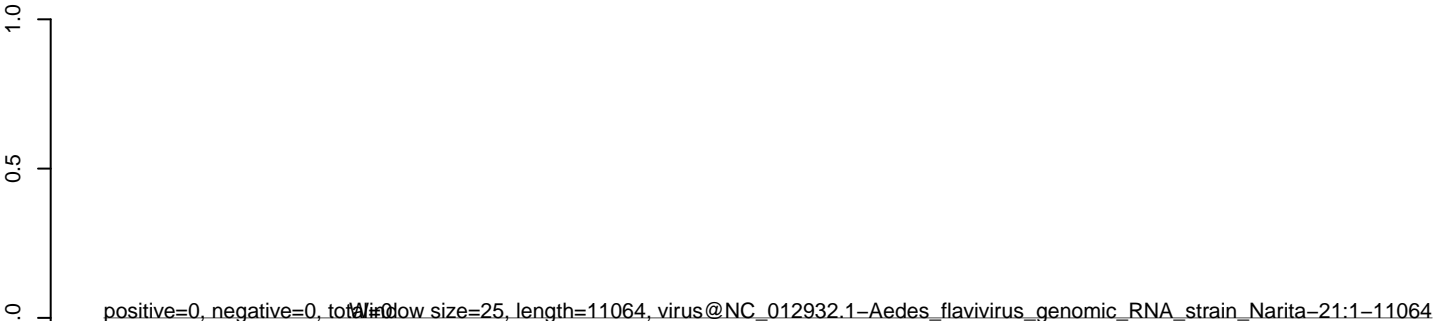
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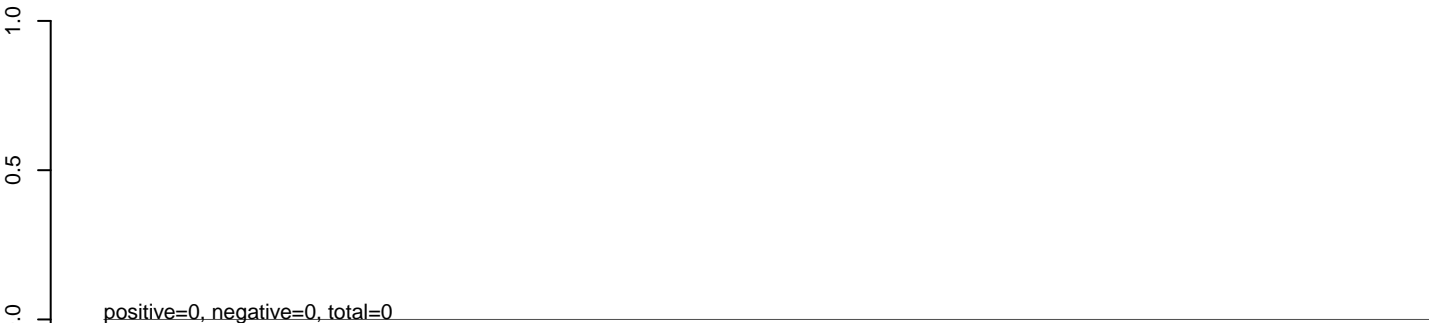


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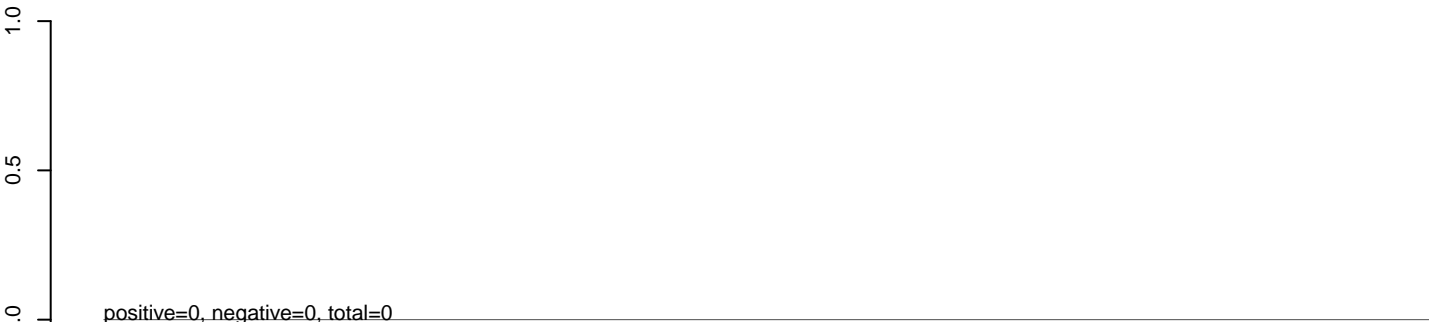


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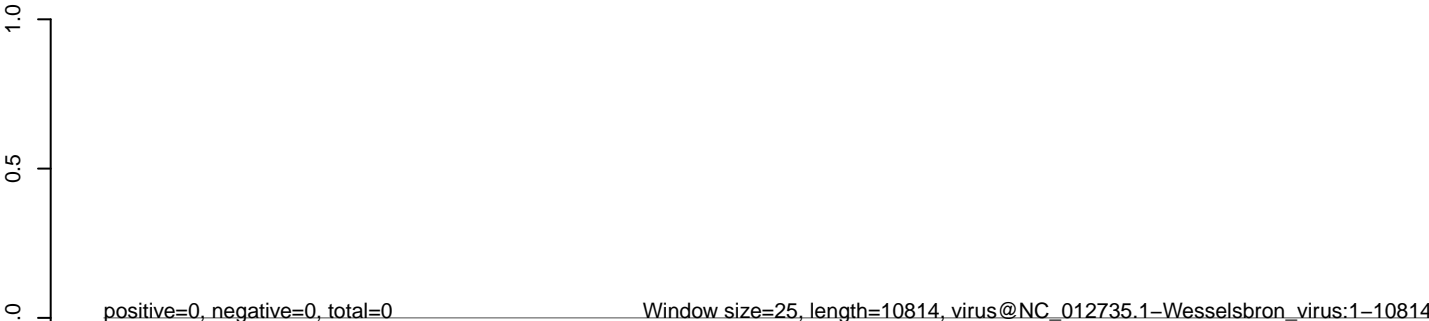
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AnGam_Ovary_BetaE.rep



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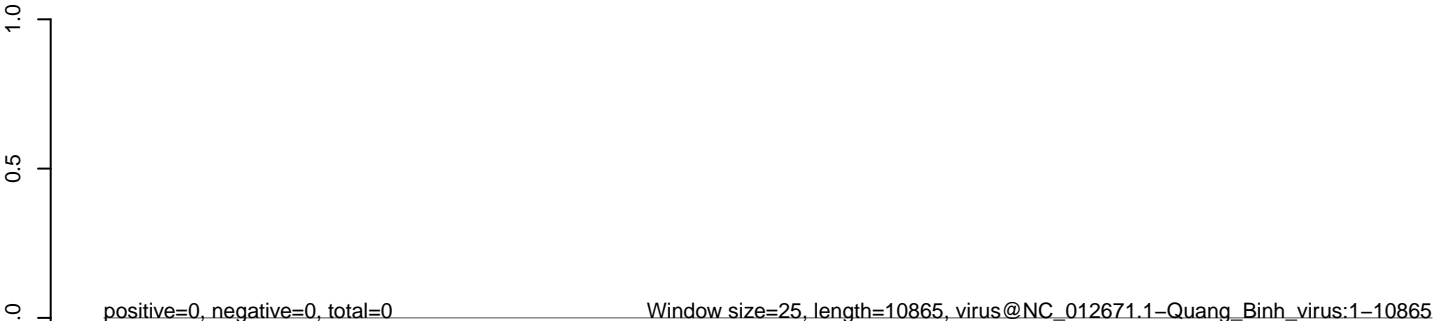
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AnGam_Ovary_BetaE.rep



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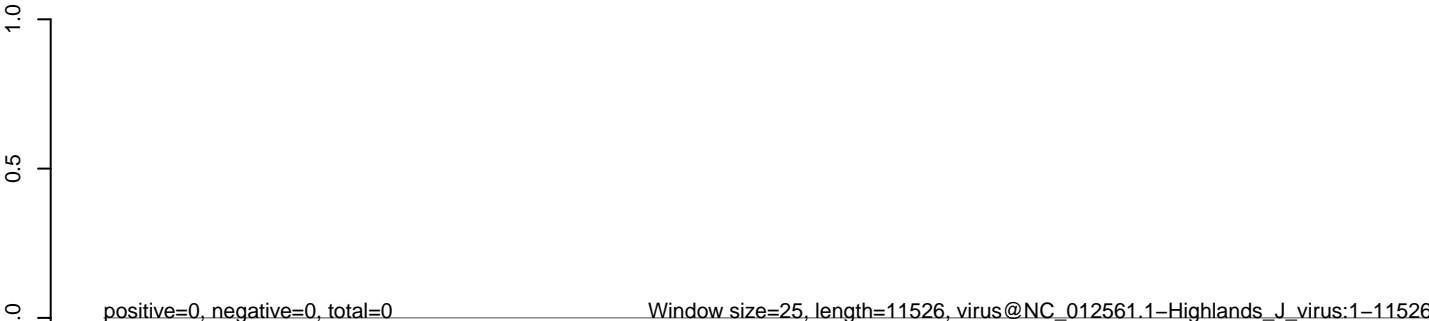
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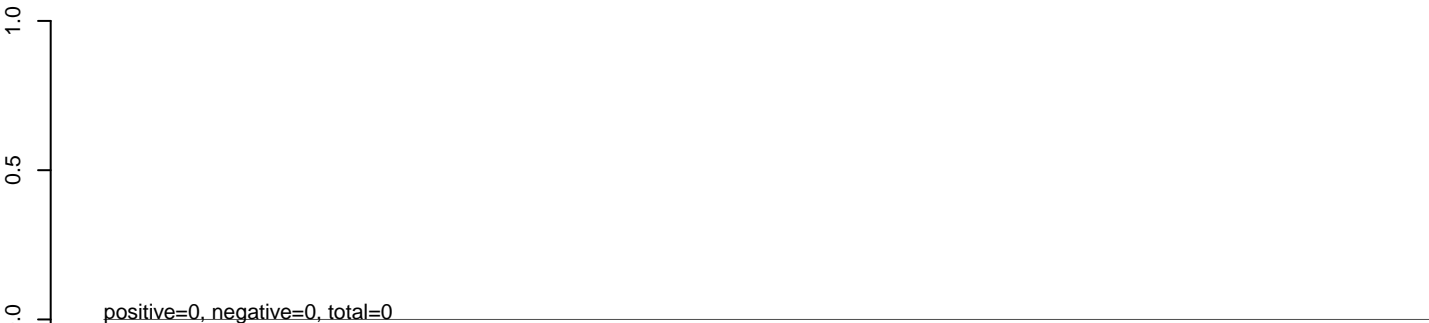


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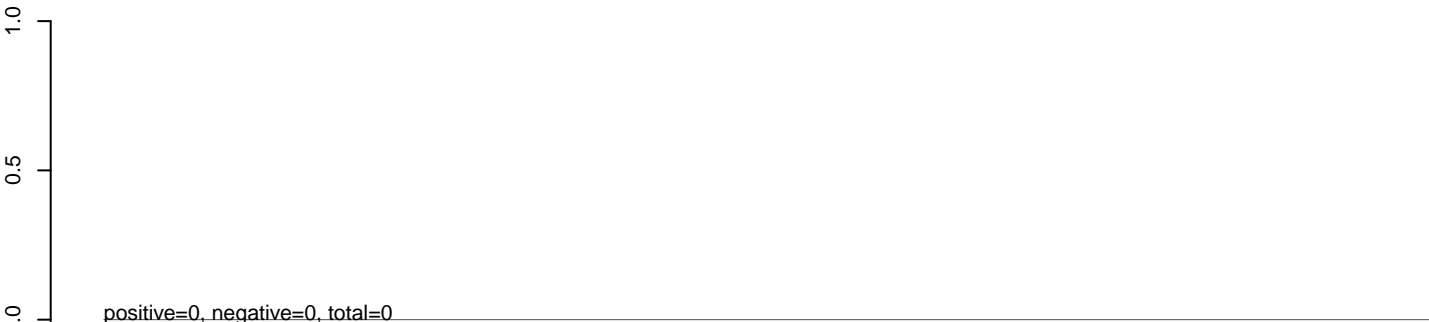


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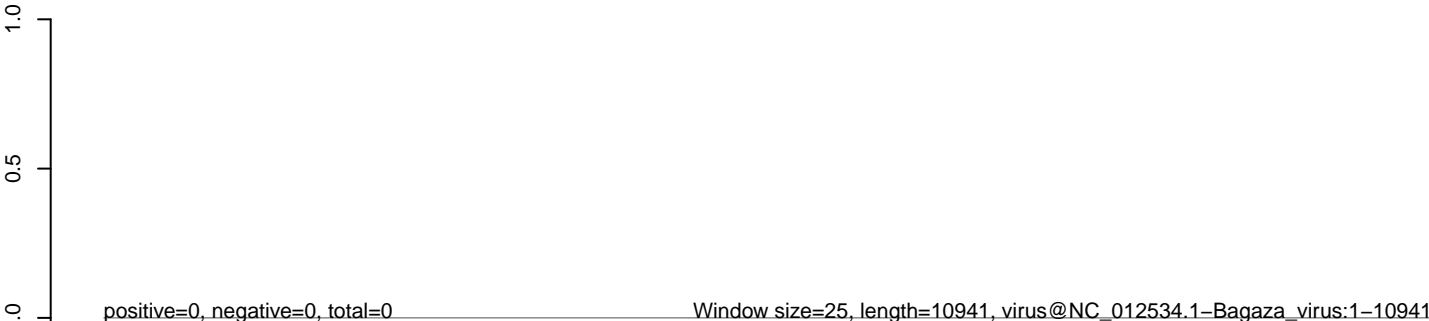
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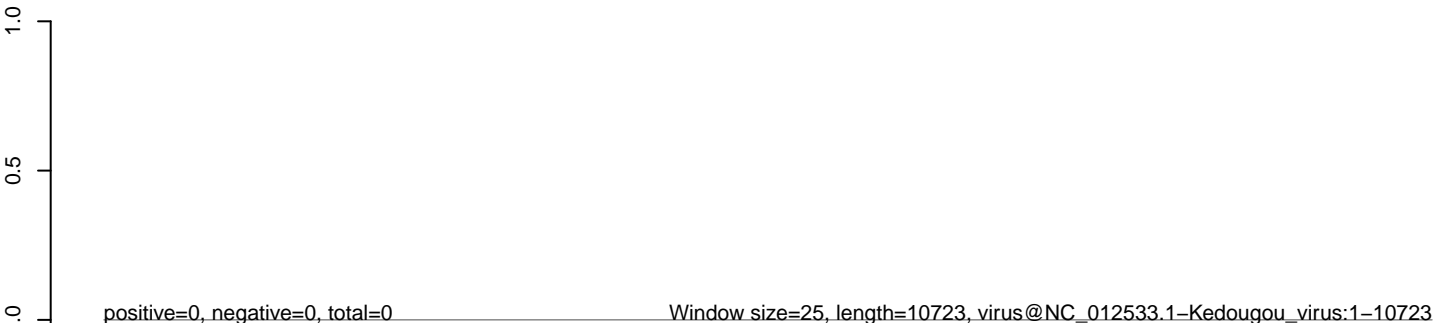
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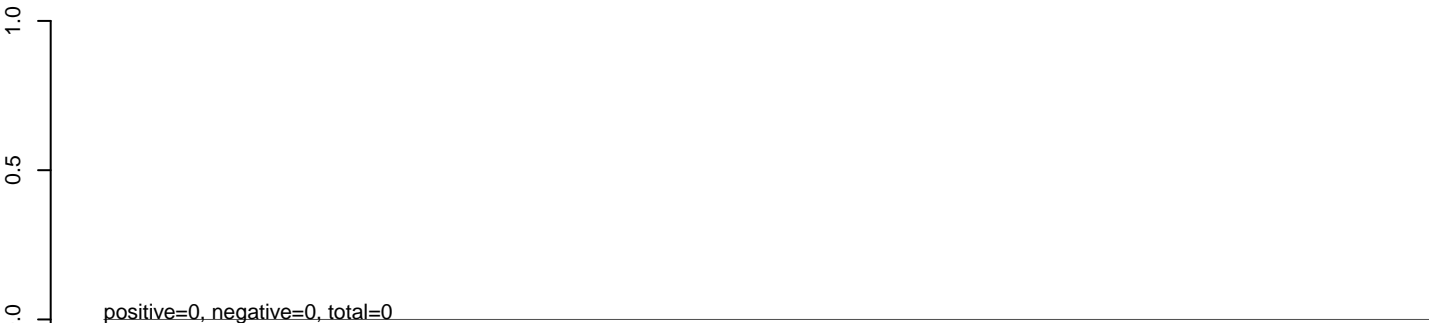


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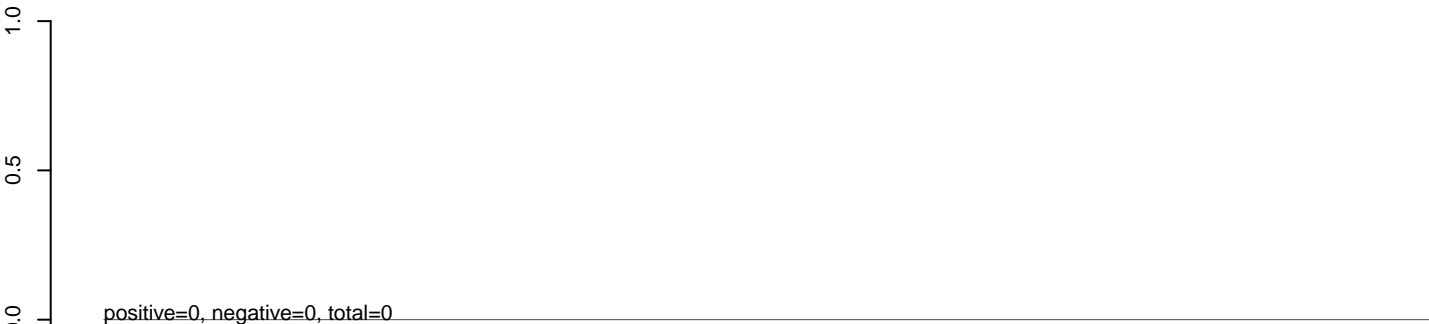


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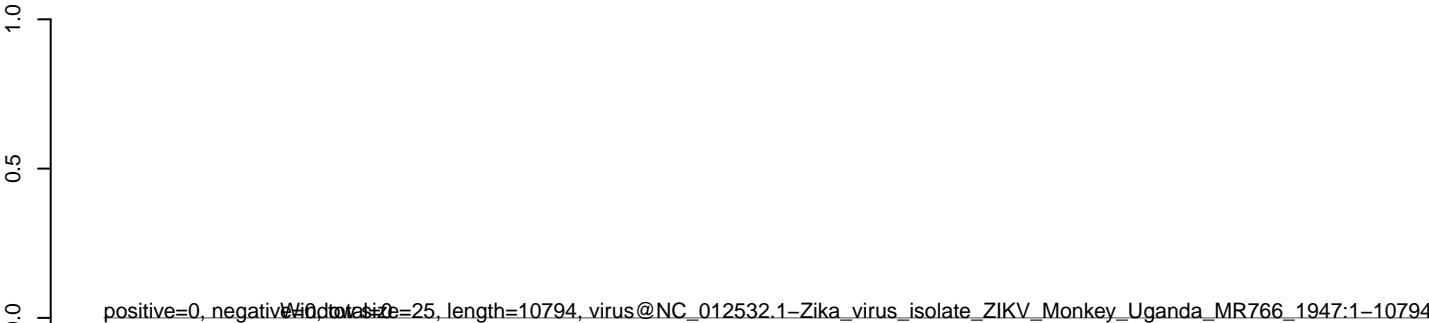
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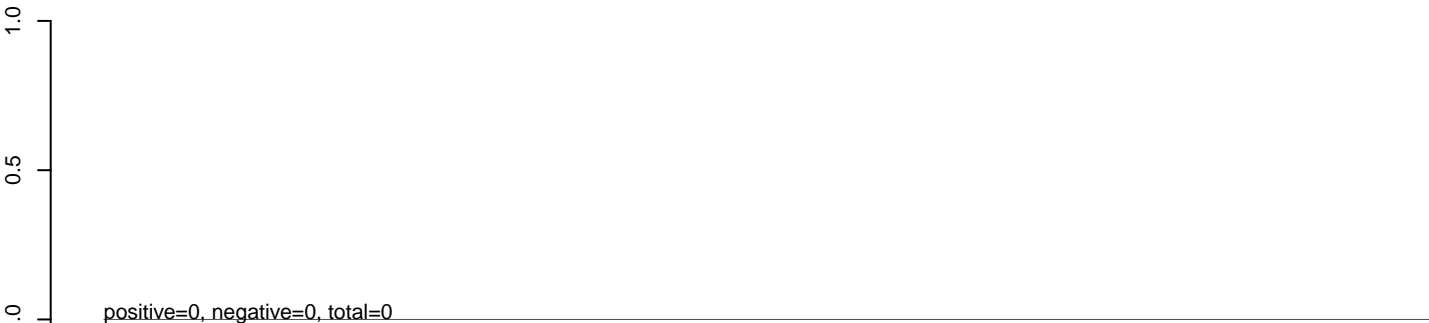


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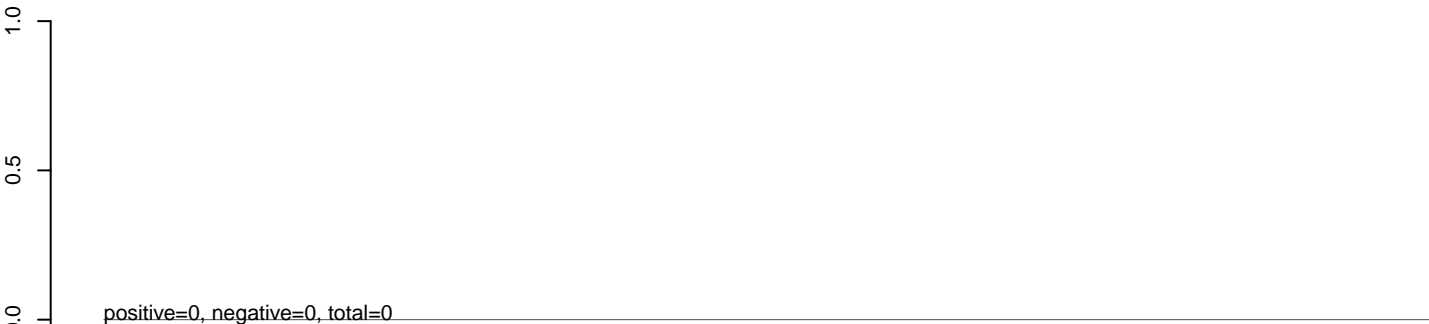


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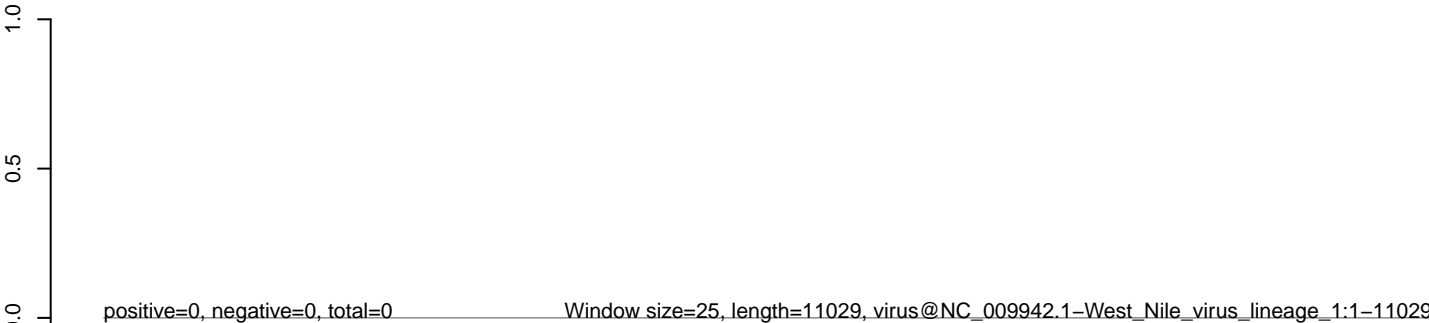
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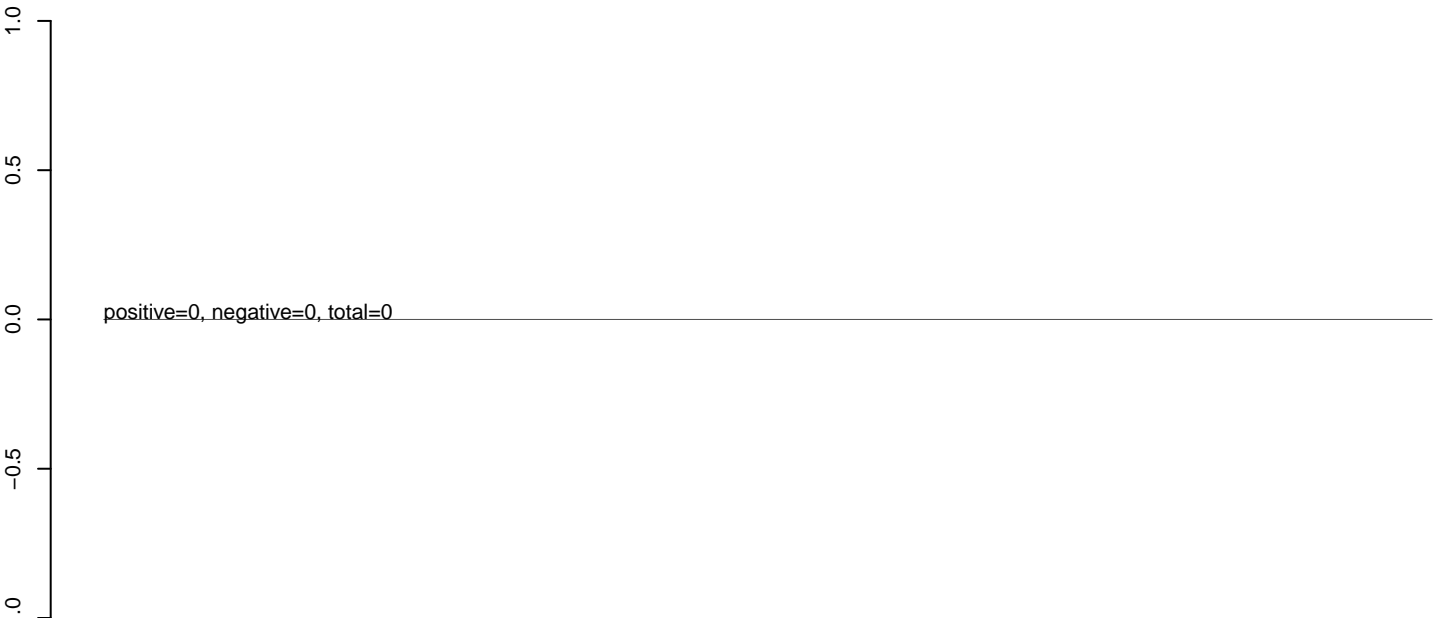


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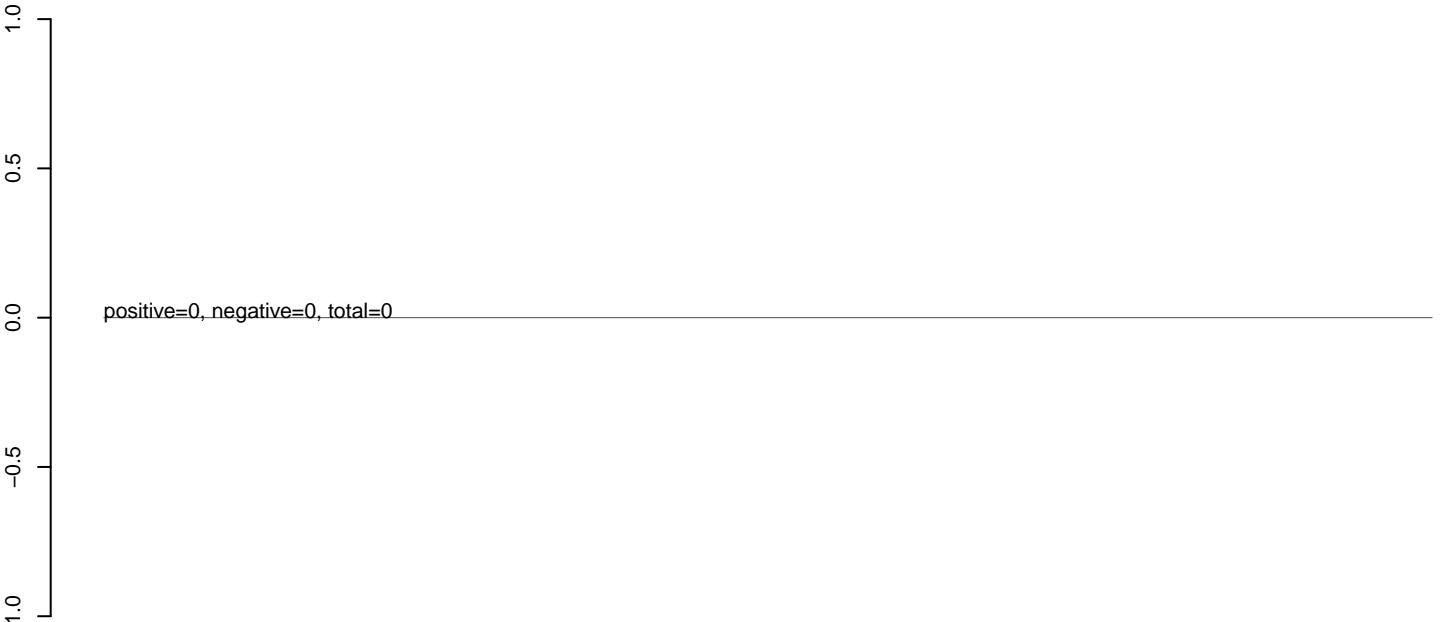


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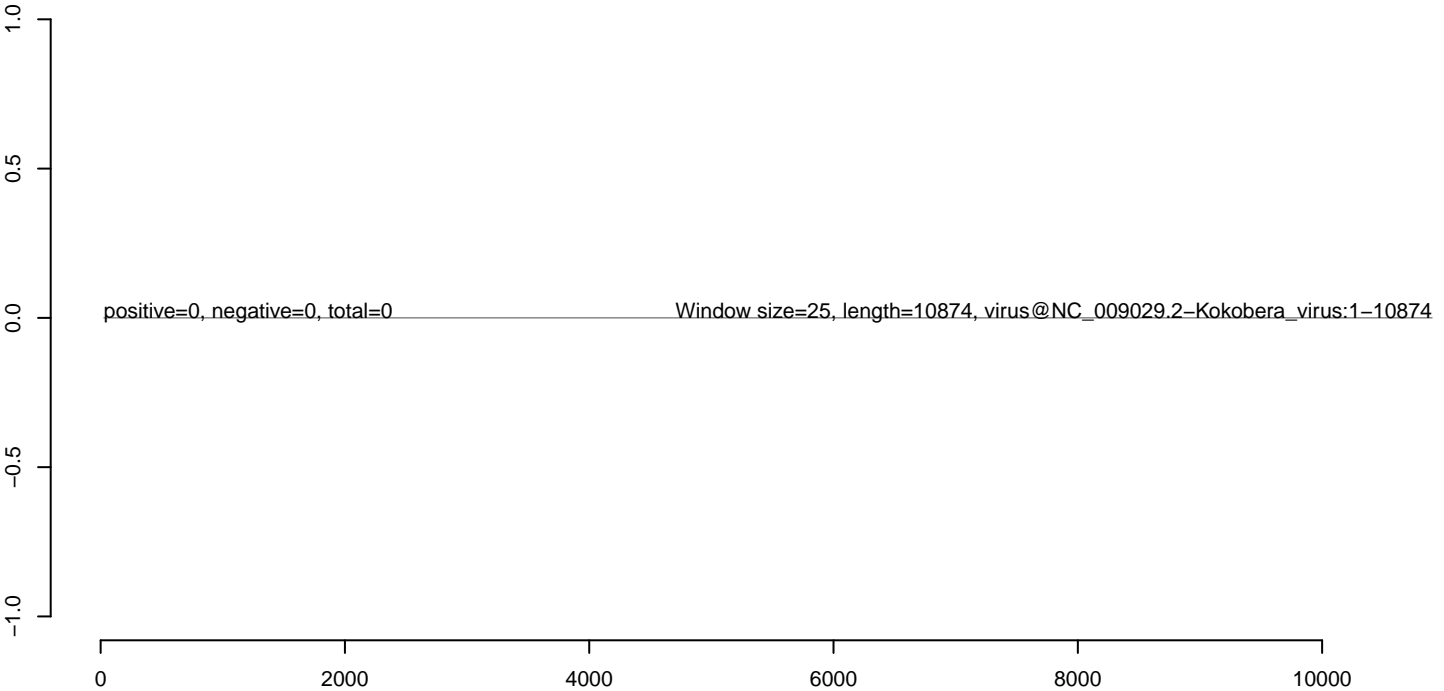
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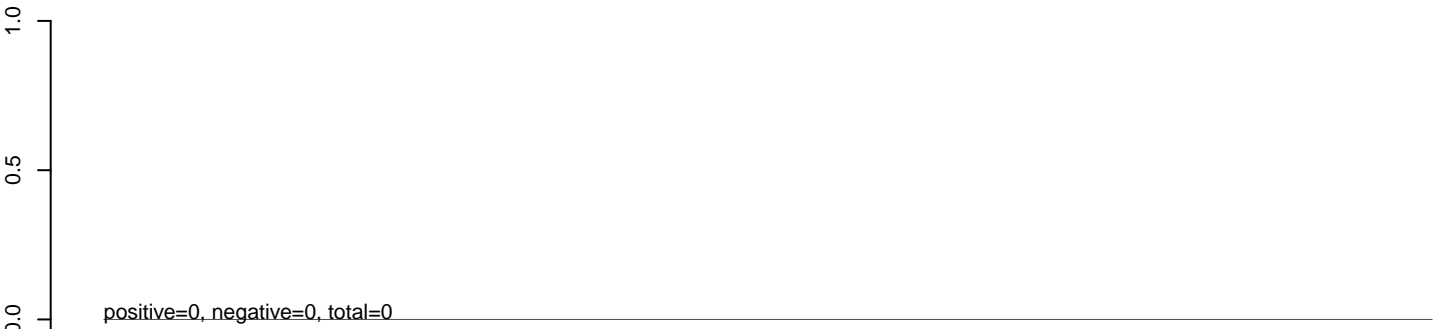
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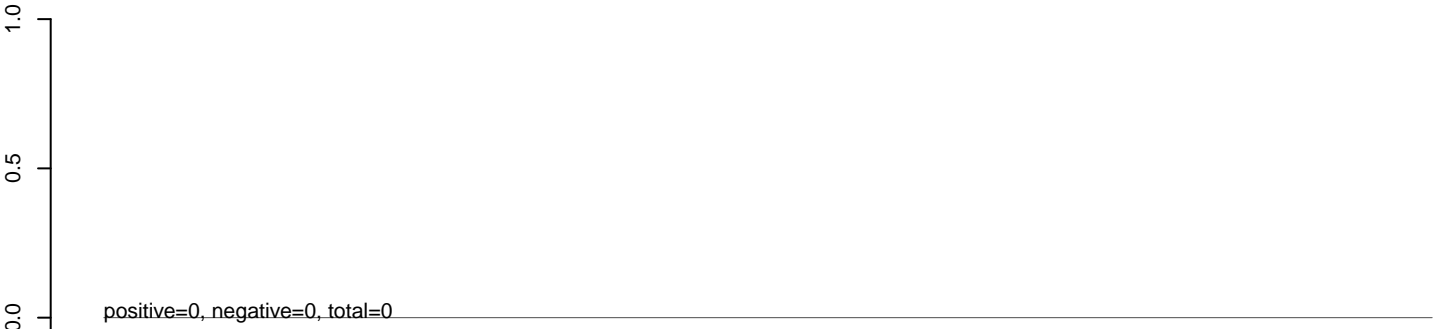
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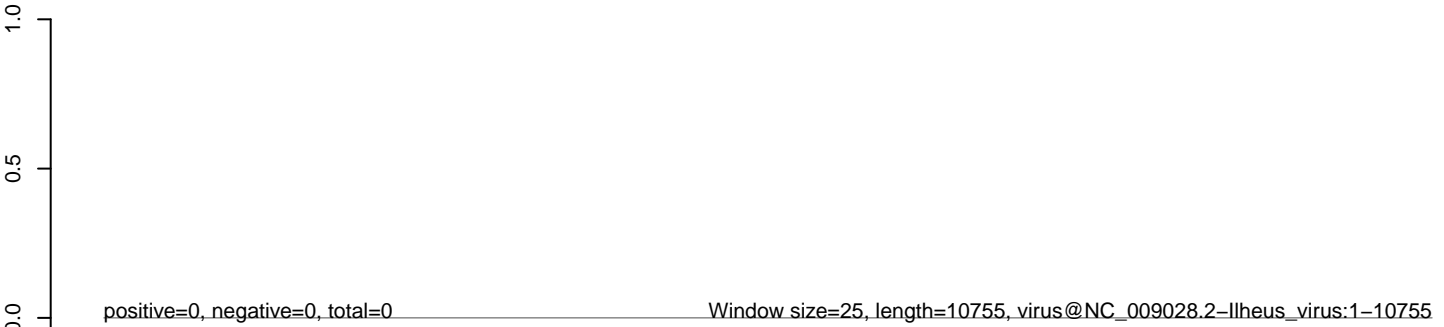
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AnGam_Ovary_BetaE.rep



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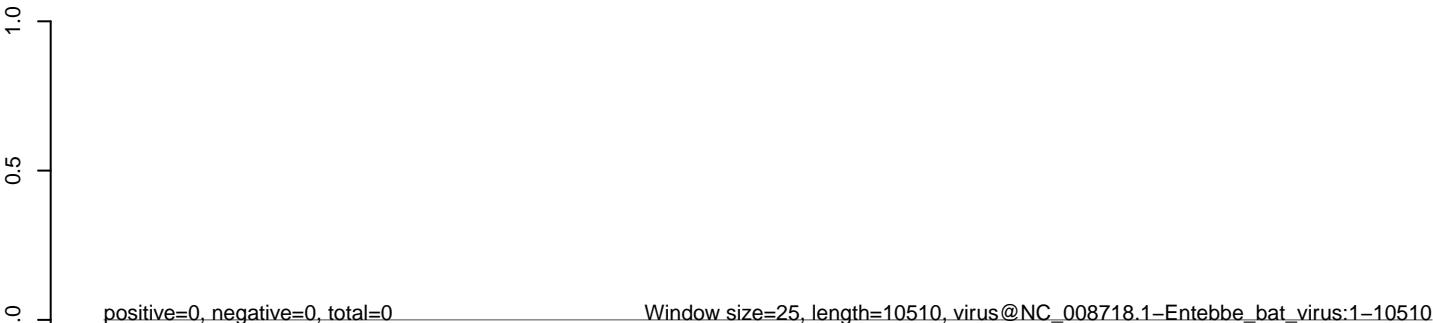
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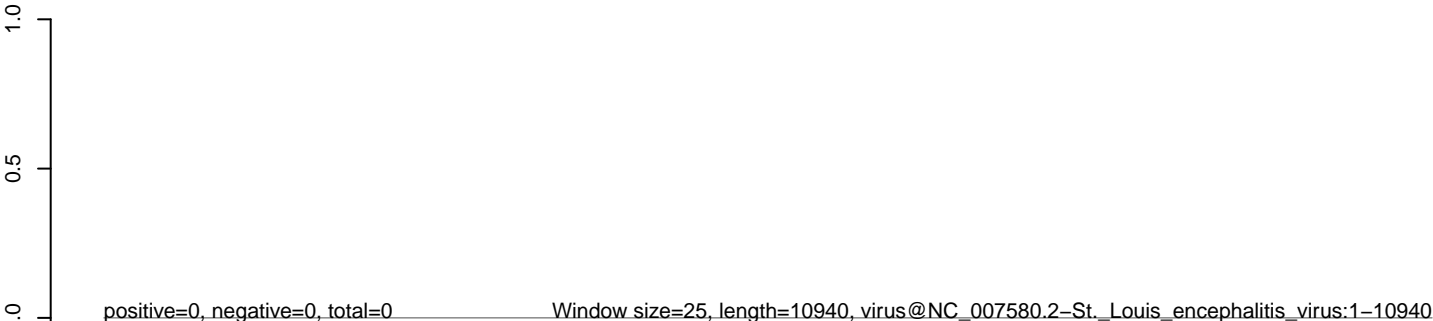
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AnGam_Ovary_BetaE.rep



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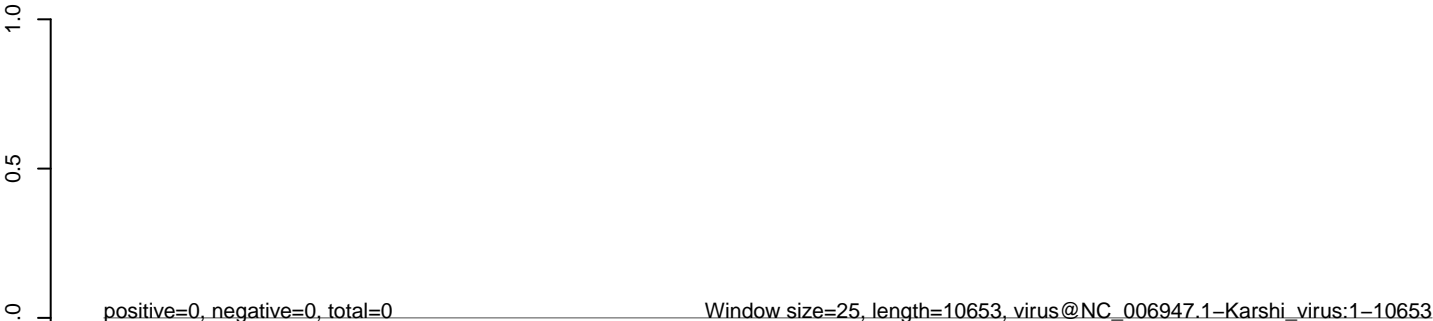
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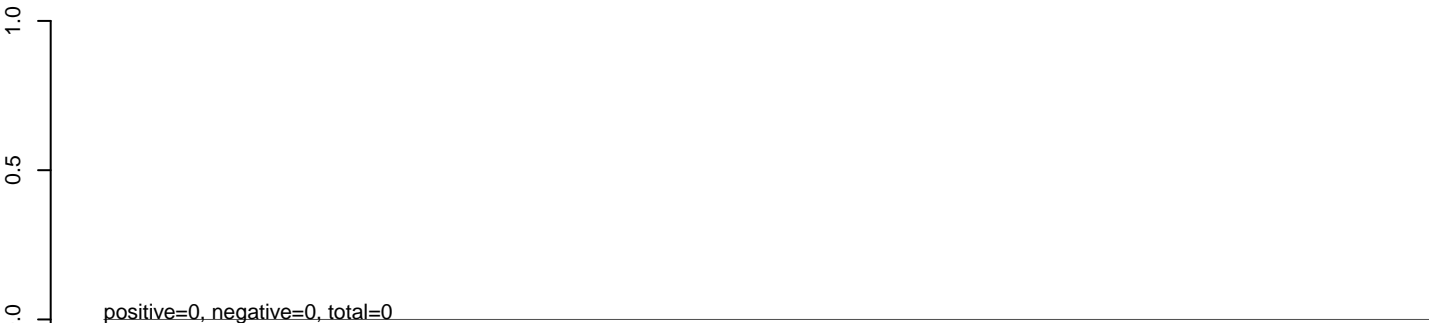
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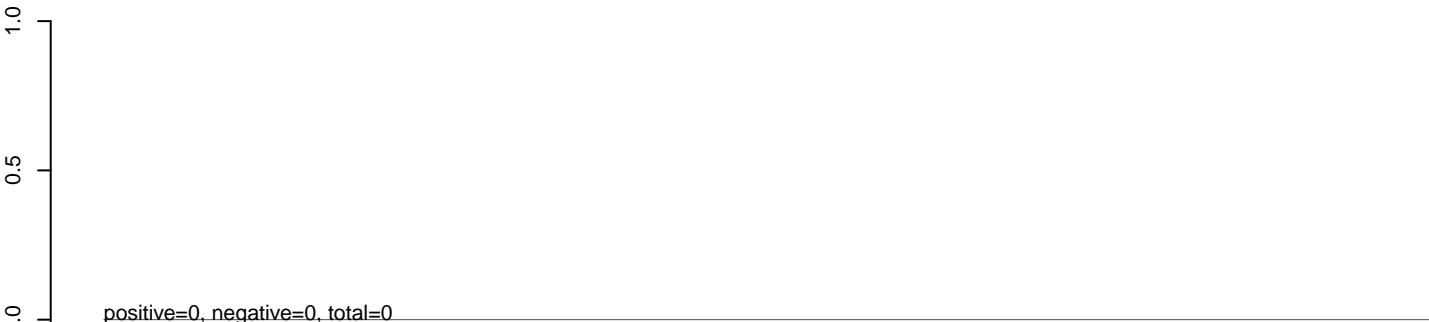
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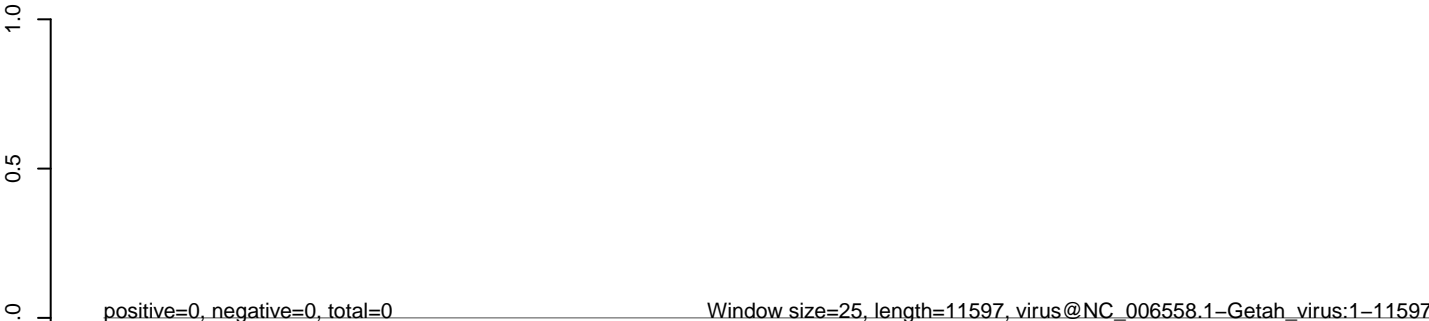
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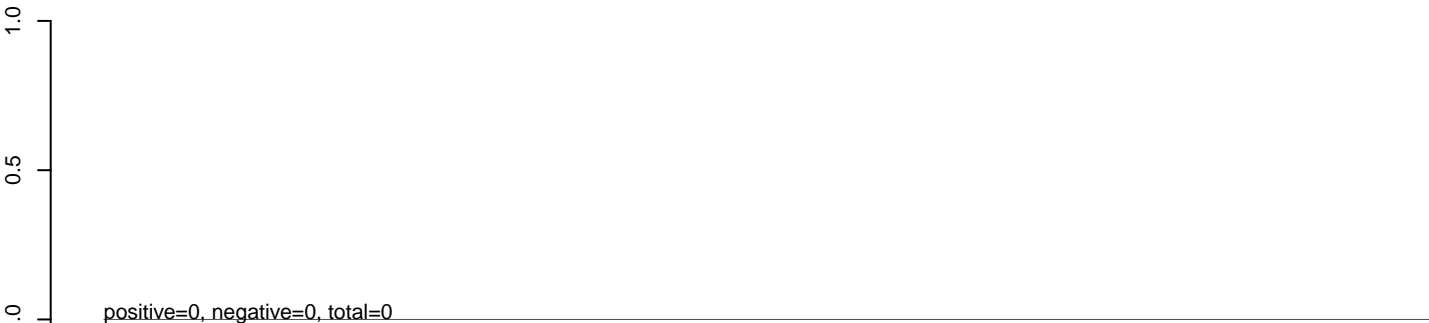
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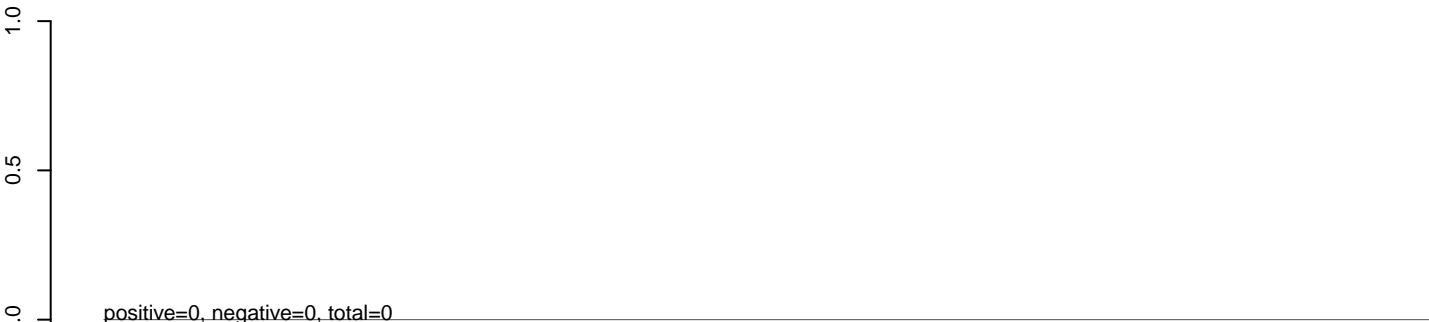
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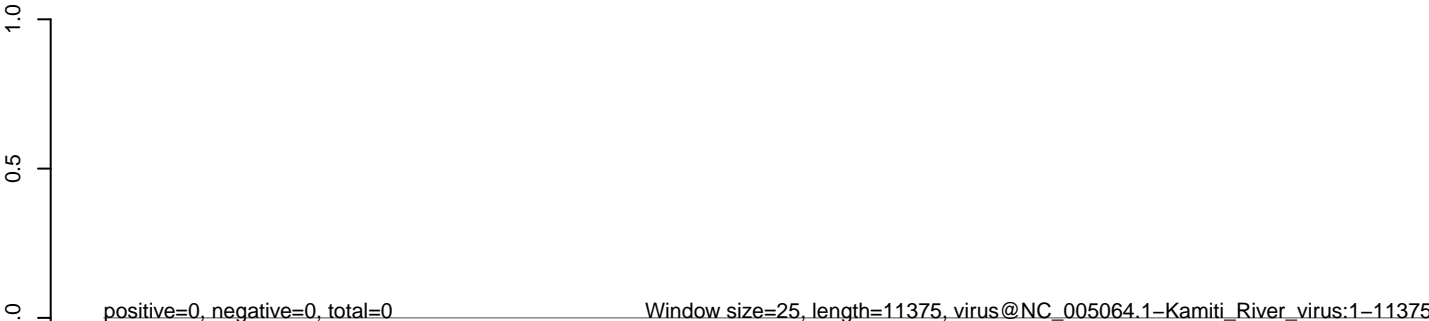
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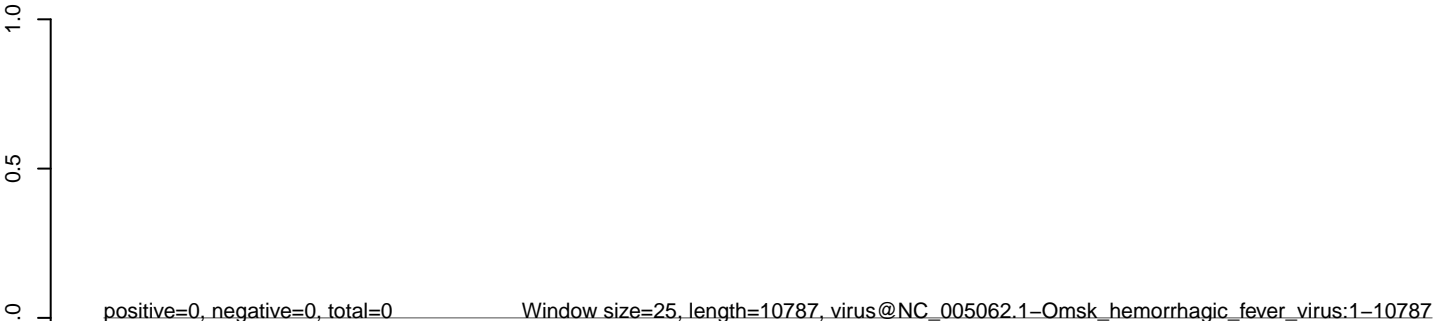
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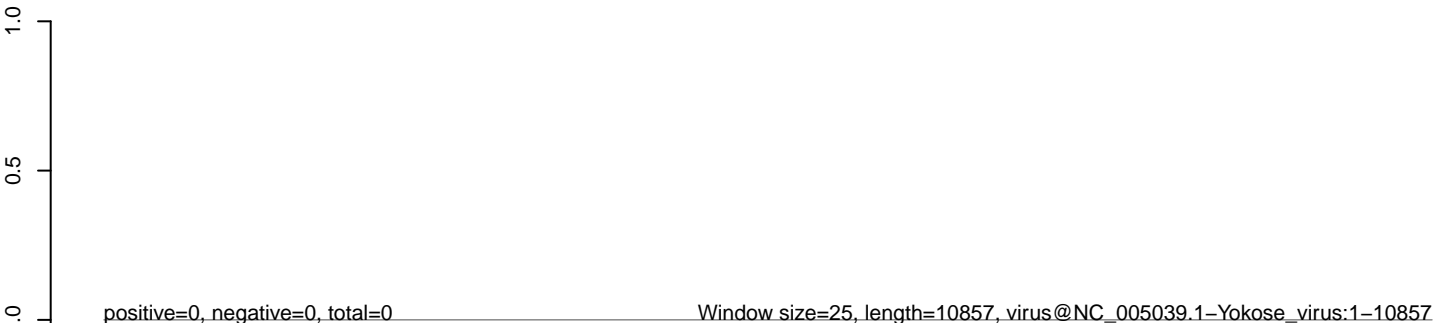
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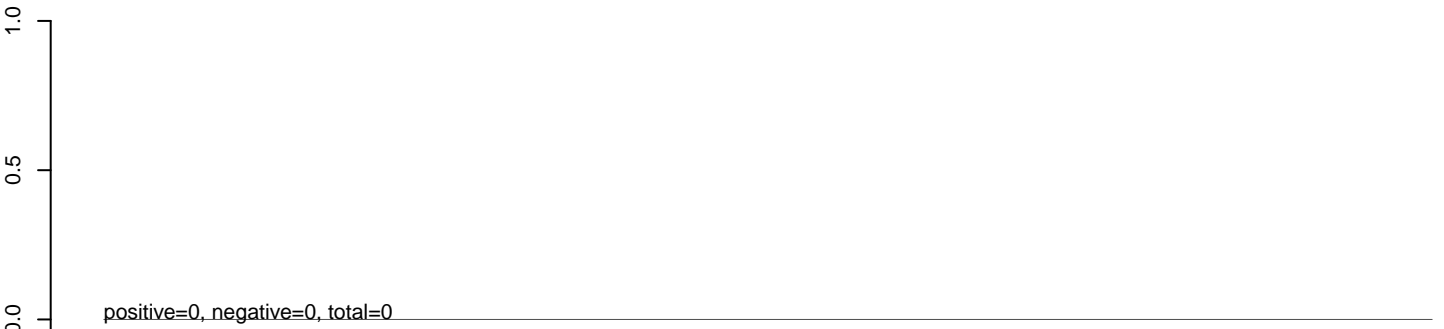


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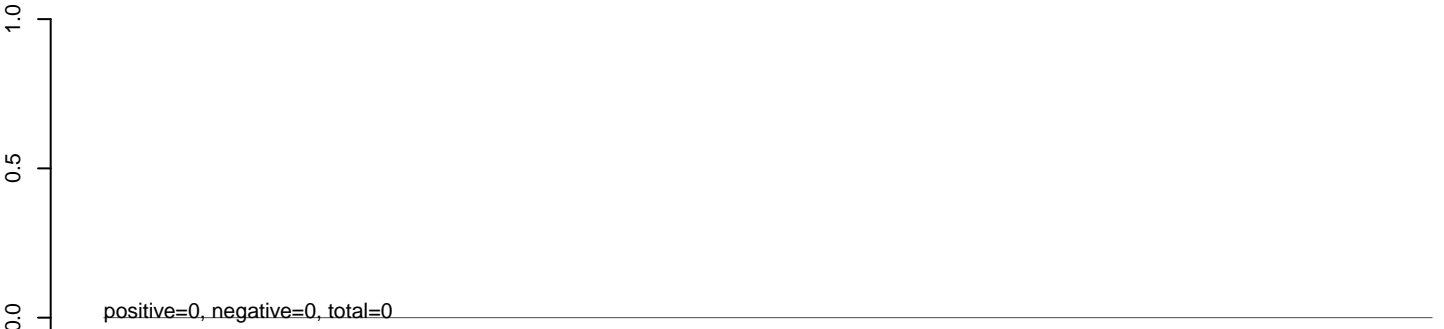


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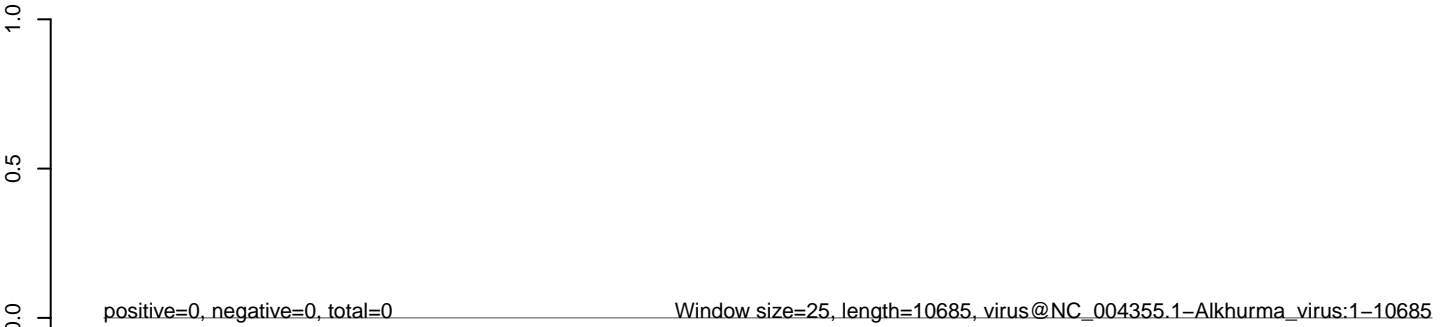
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AnGam_Ovary_BetaE.rep



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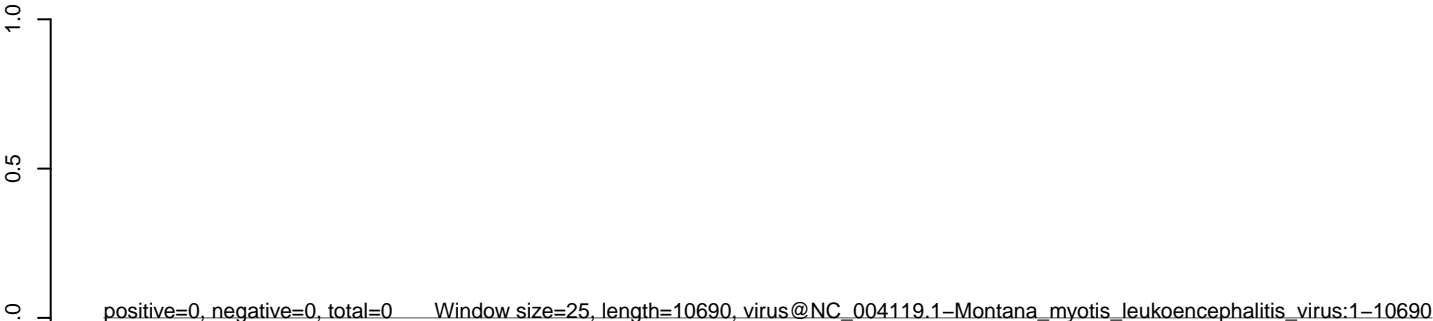
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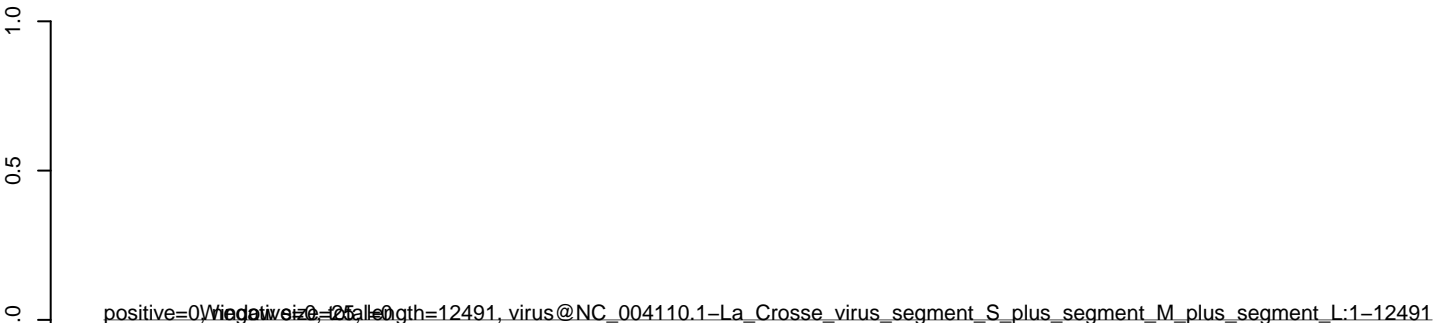
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AnGam_Ovary_BetaE.rep



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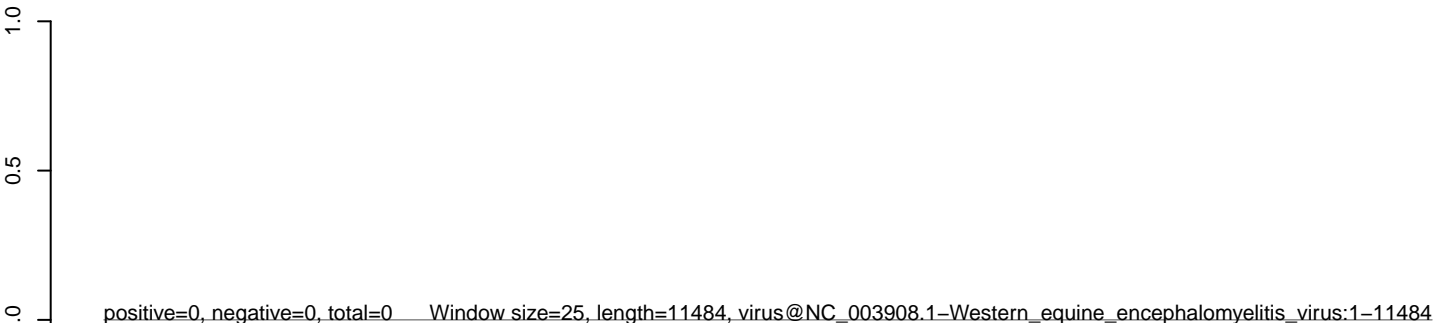
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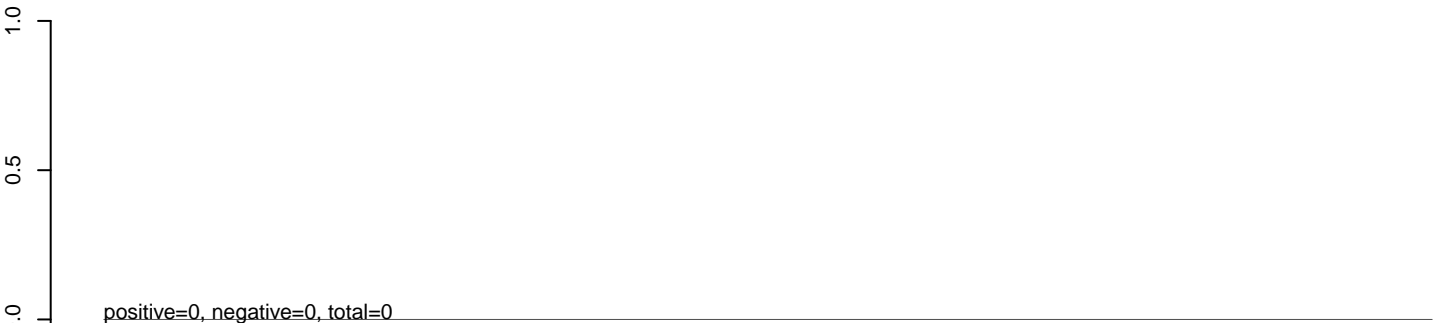


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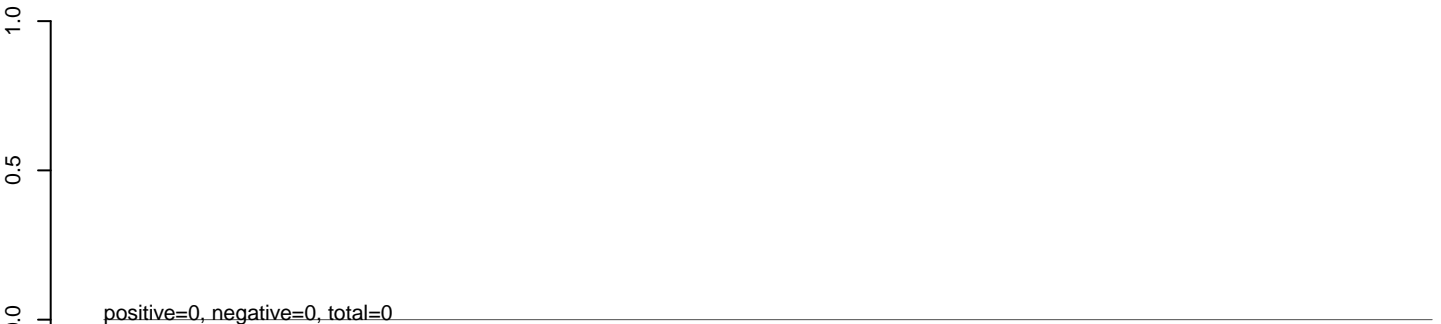


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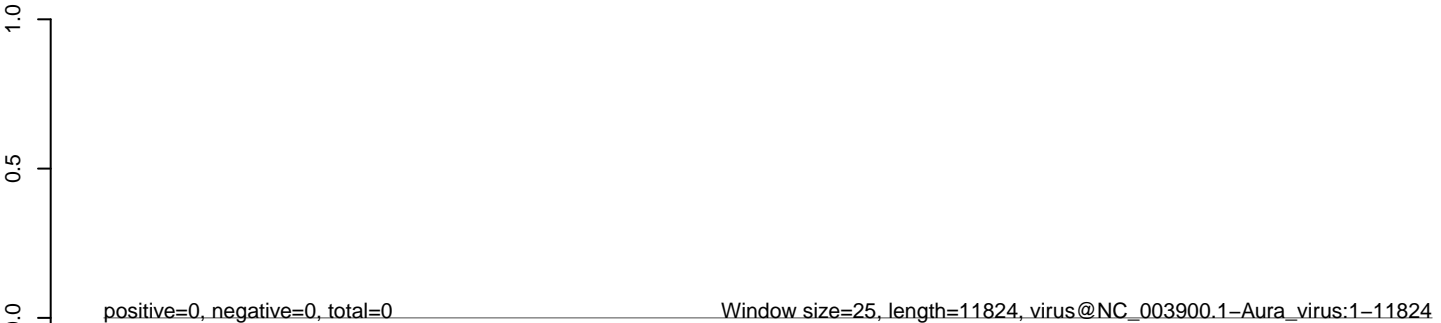
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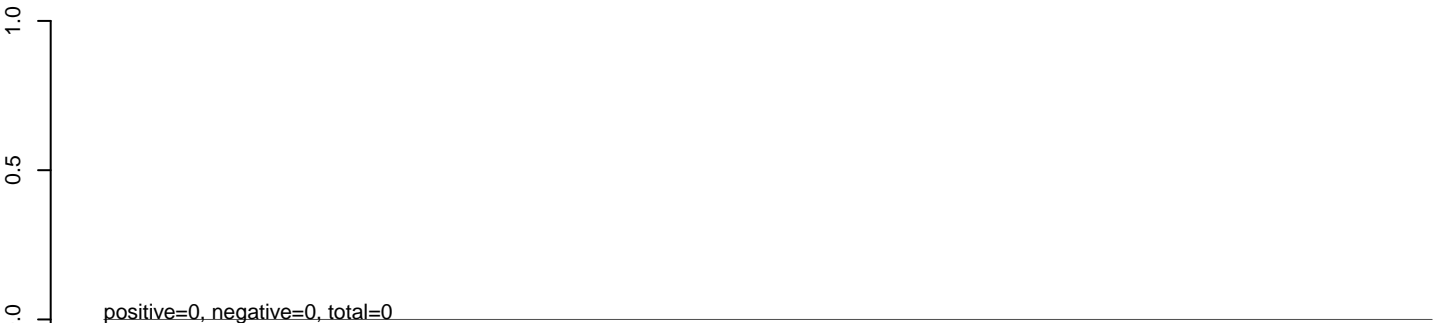


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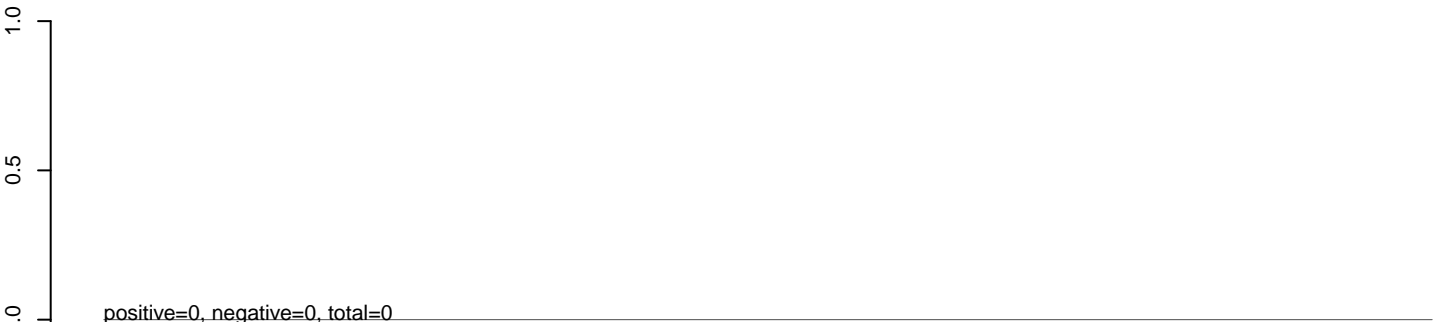


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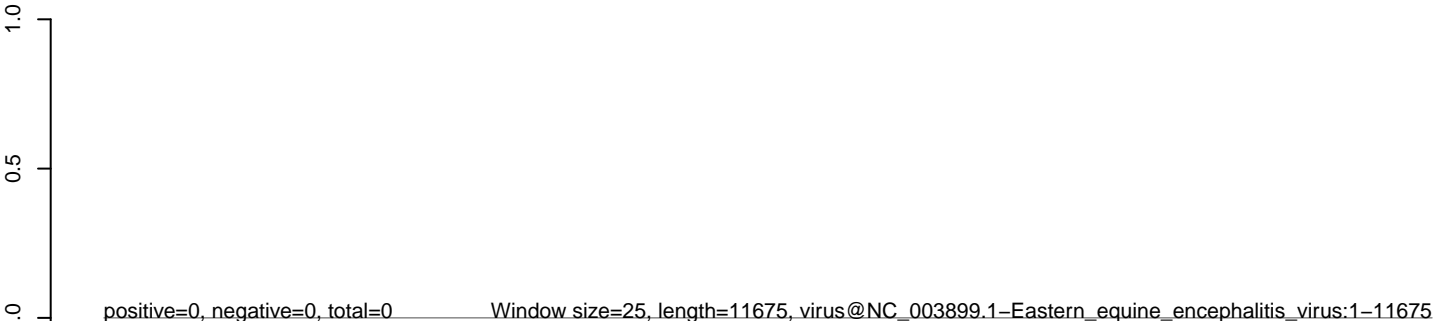
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AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



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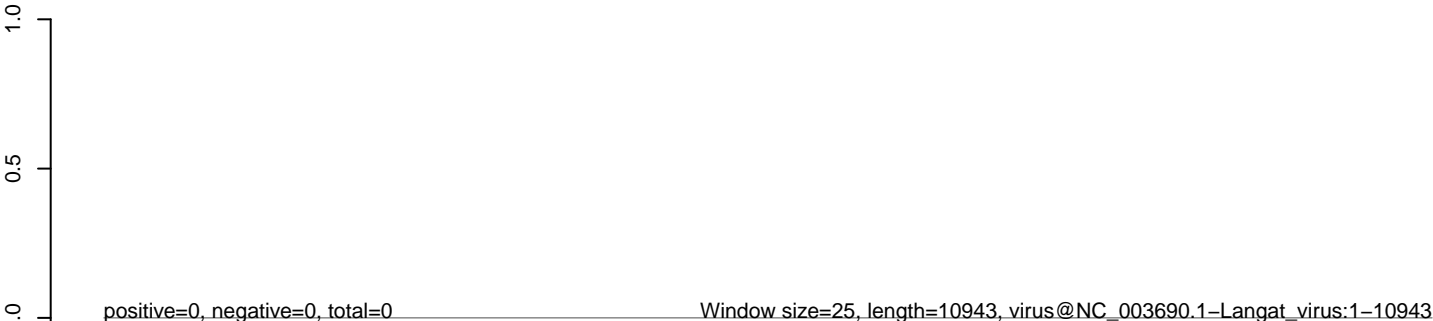
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AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



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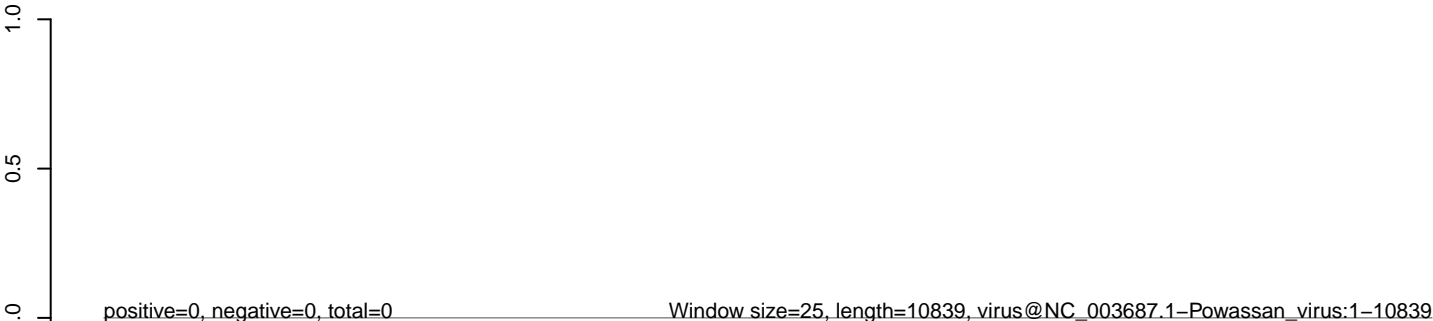
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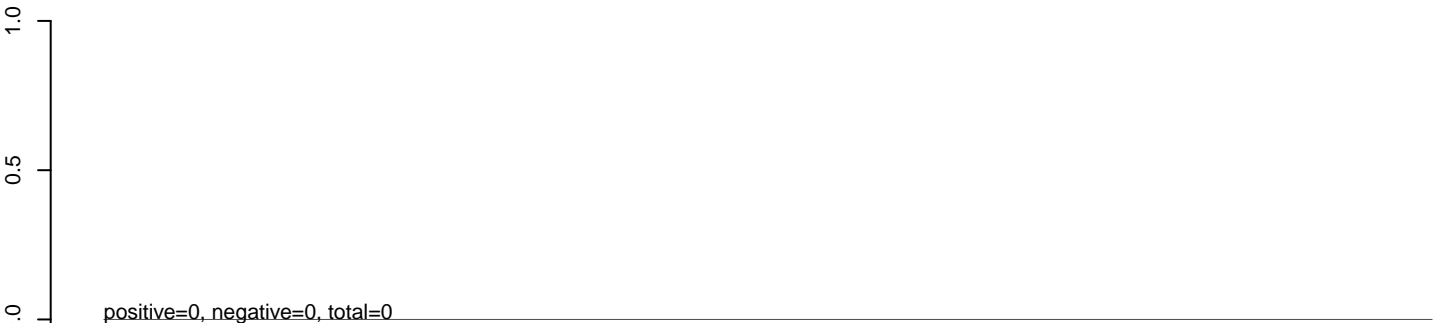


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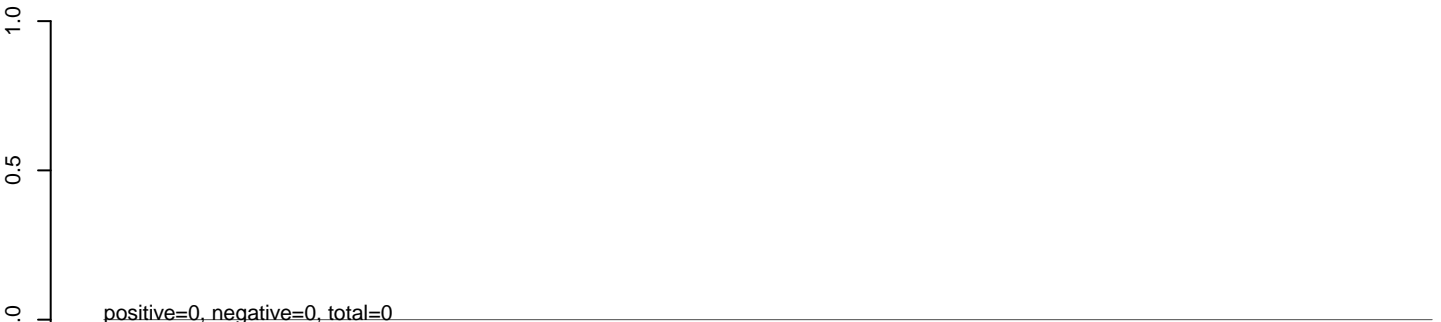


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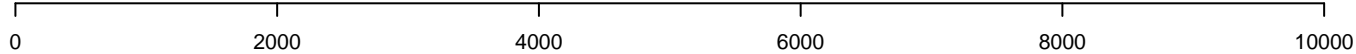
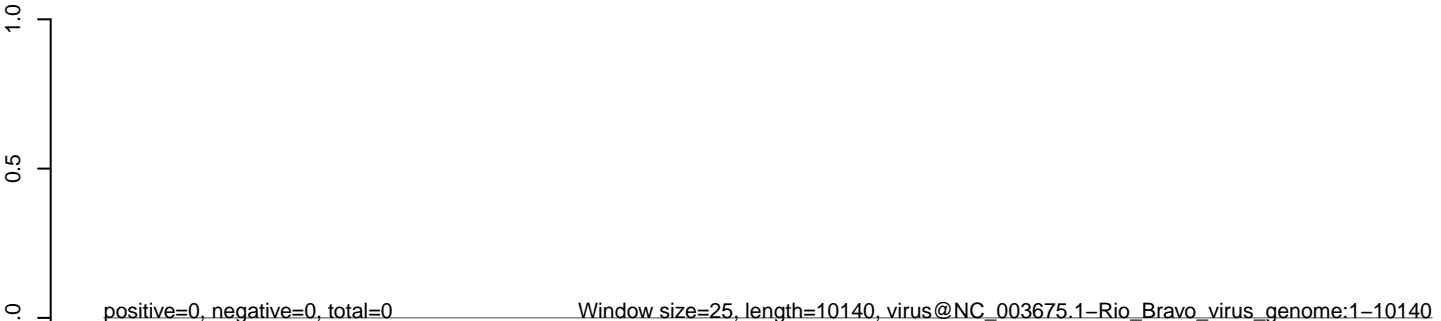
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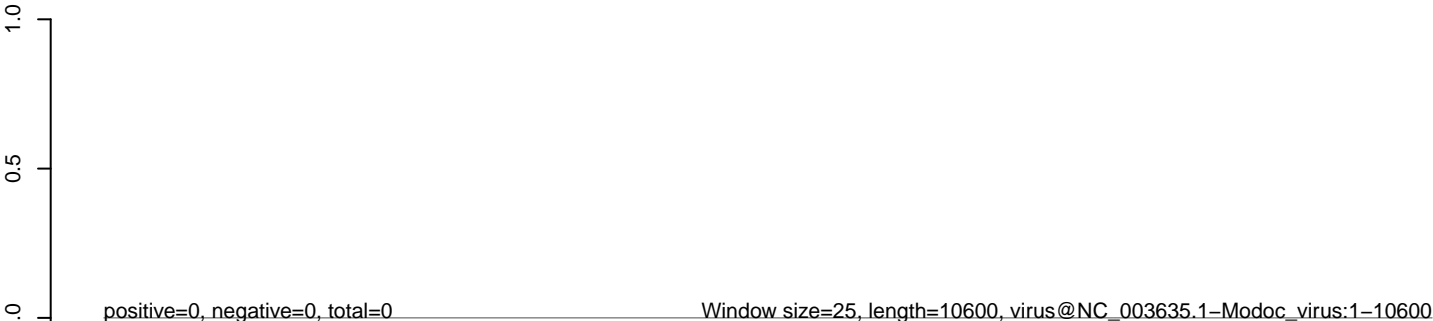
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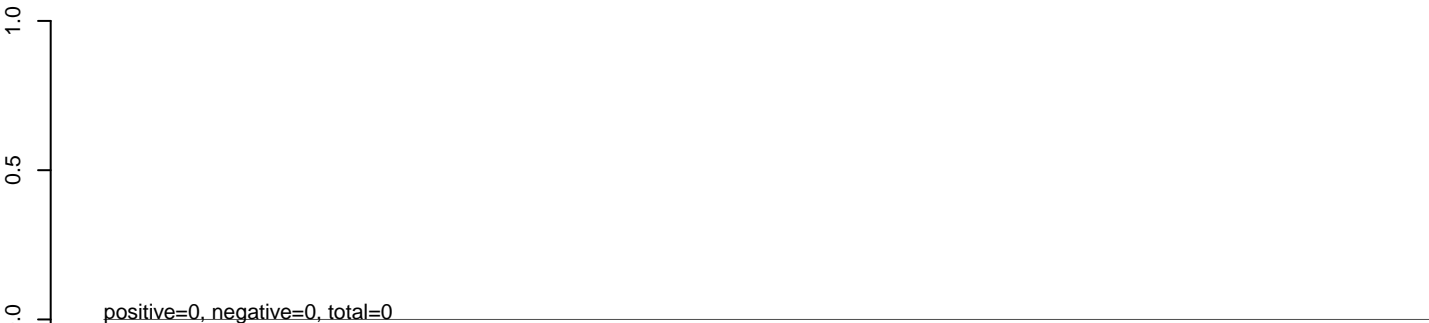


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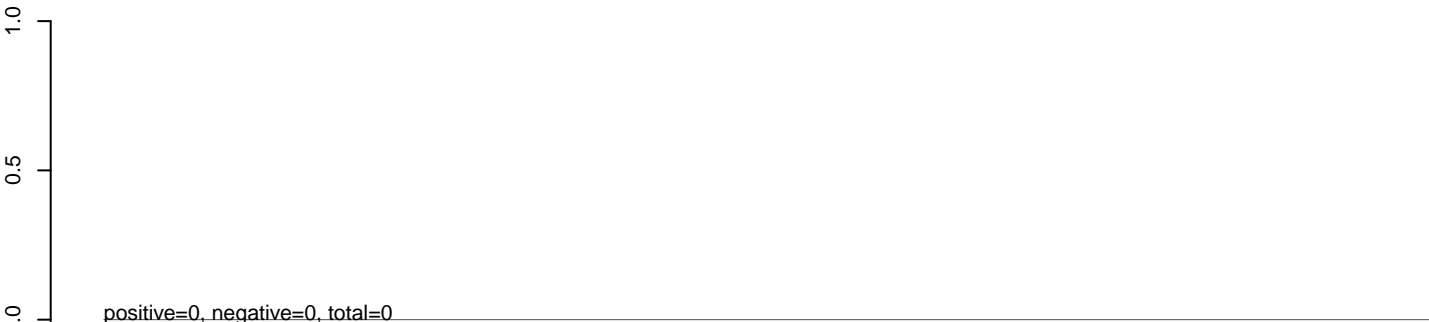


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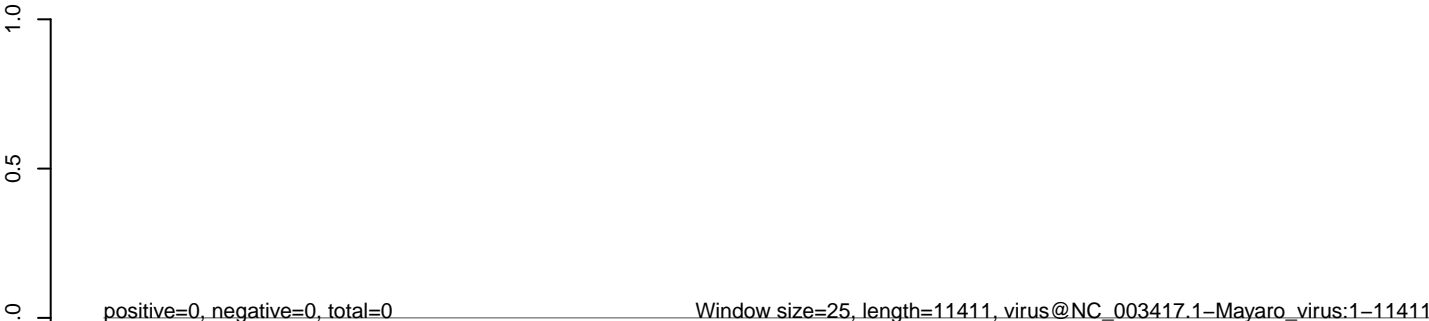
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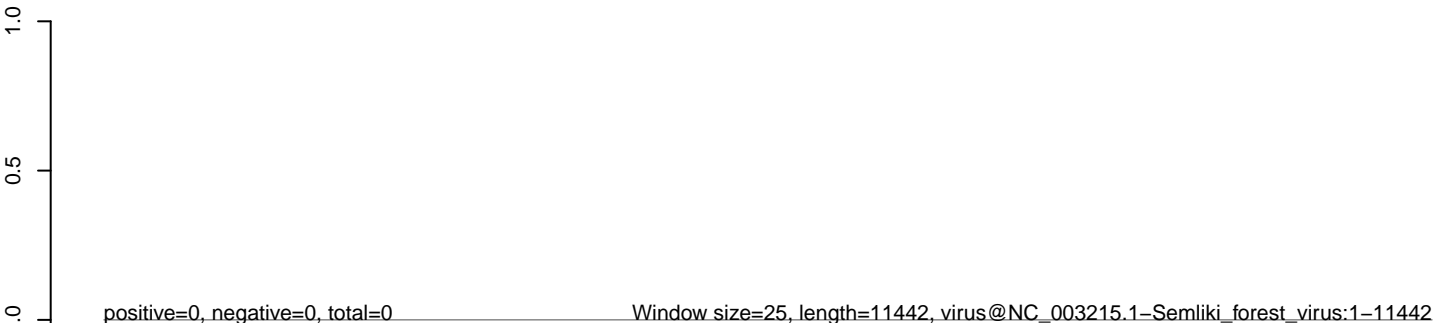
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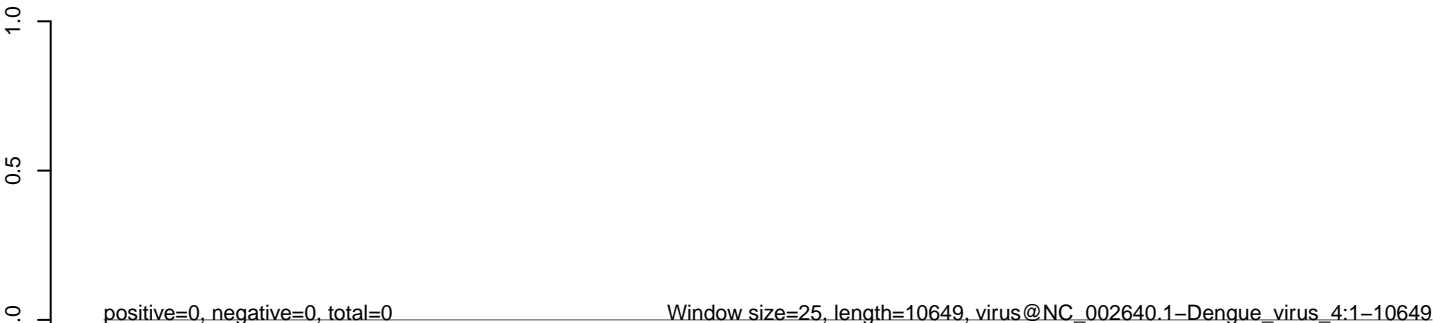
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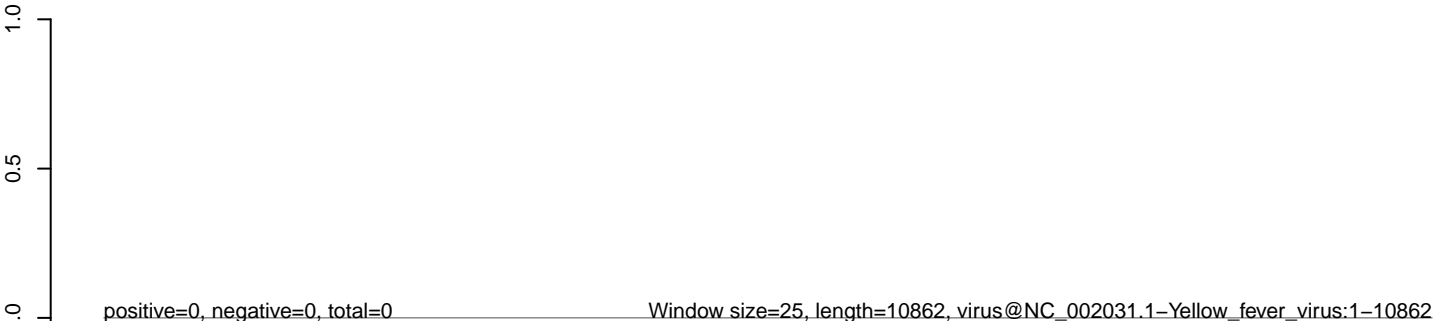
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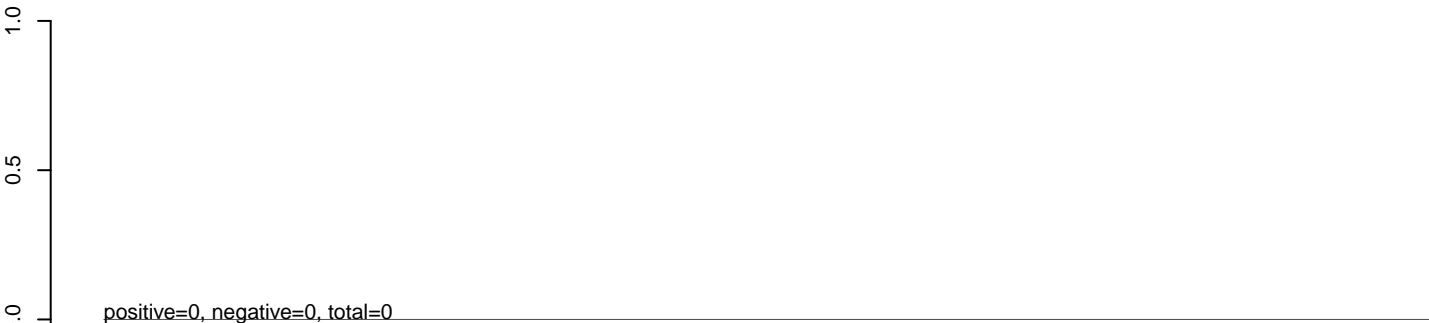


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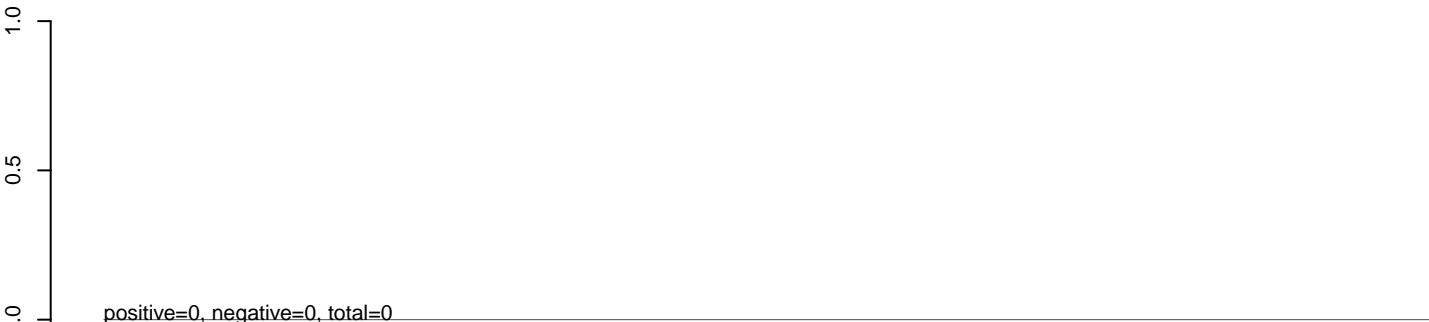


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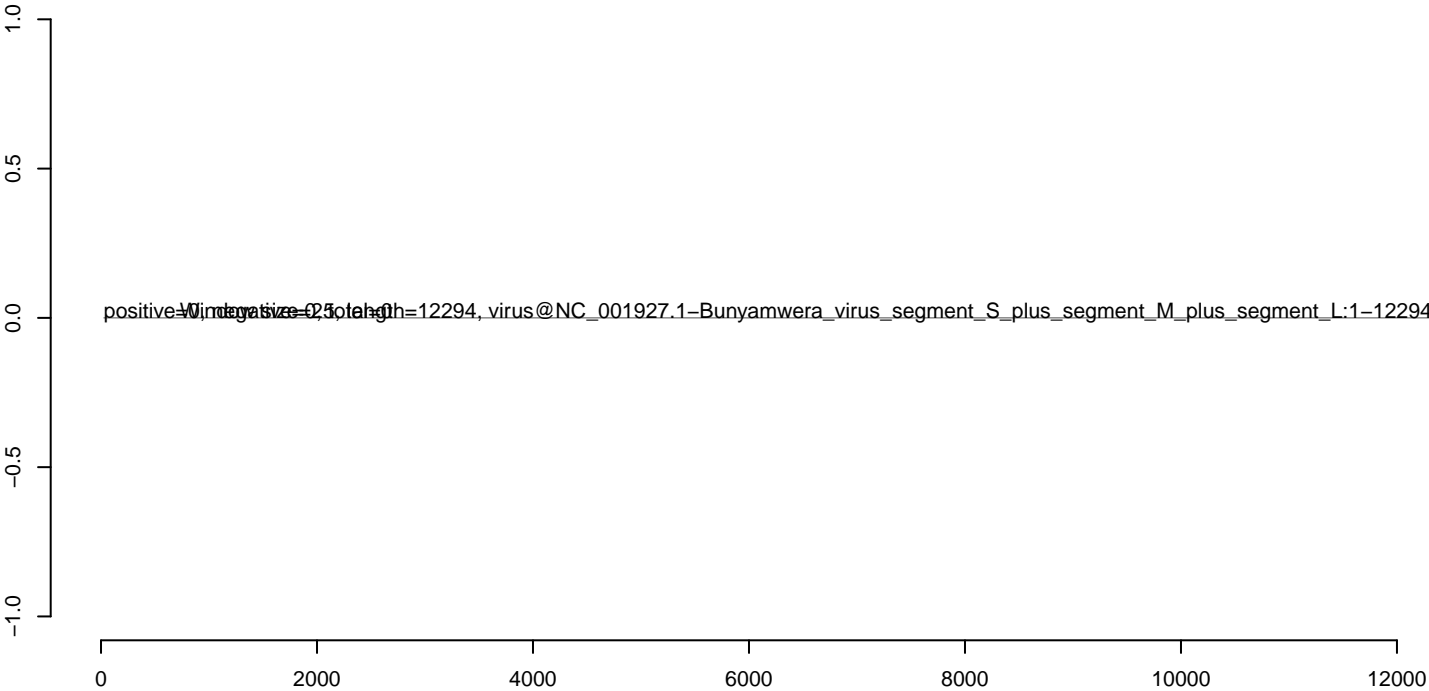
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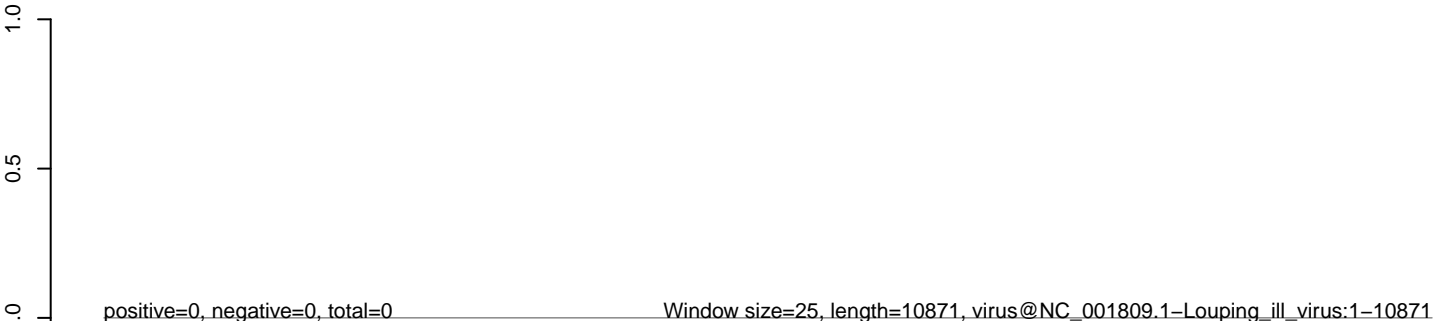
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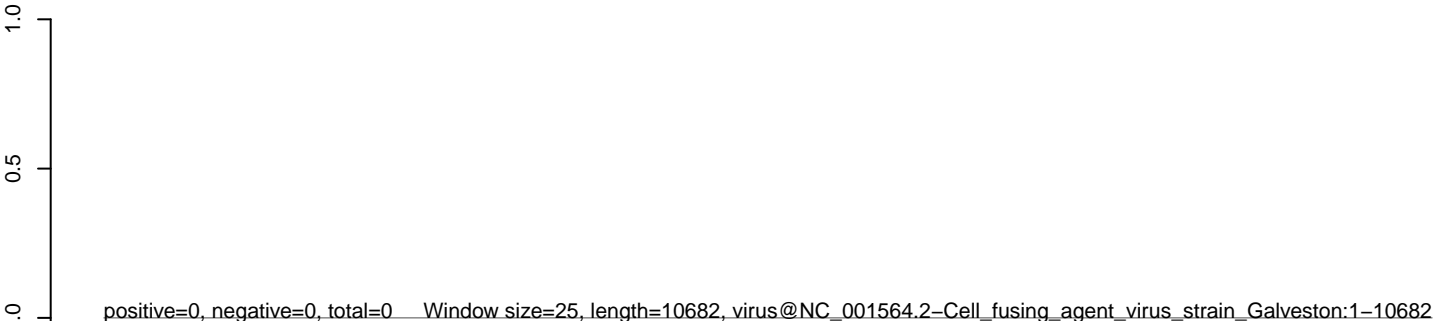
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AnGam_Ovary_BetaE.rep



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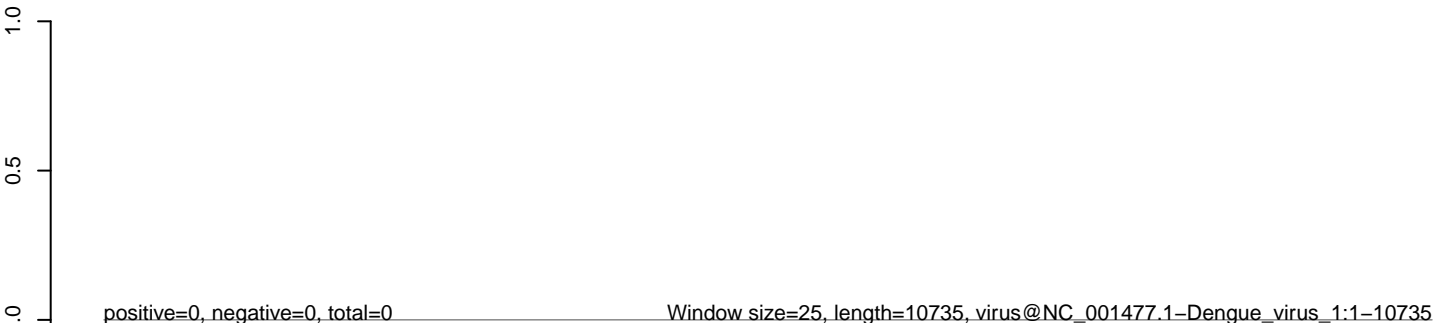
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AnGam_Ovary_BetaE.rep



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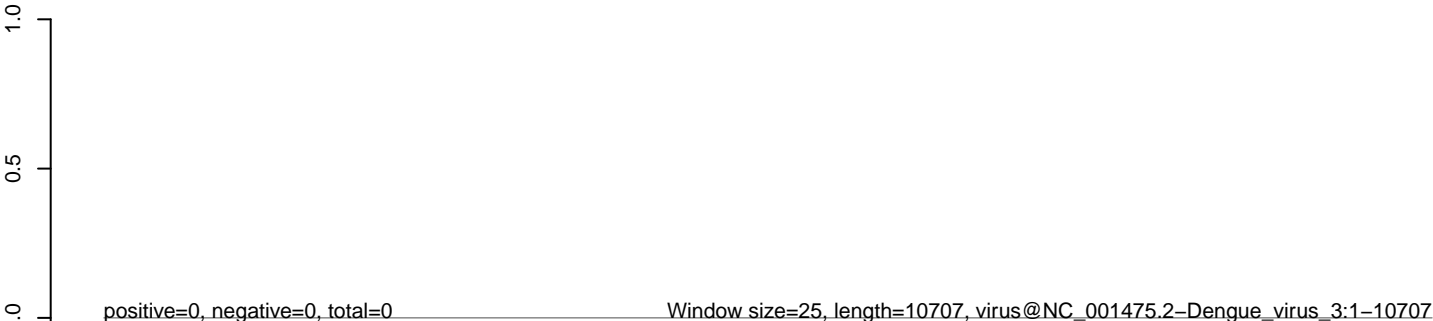
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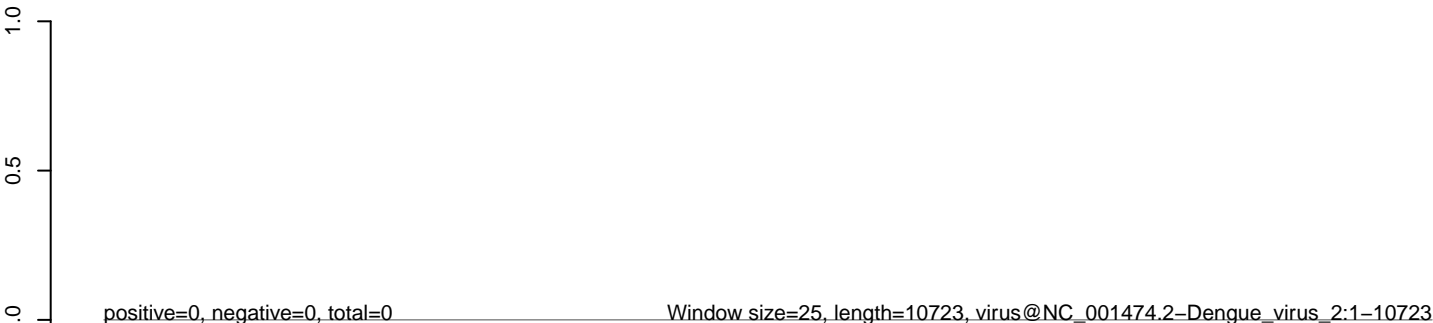
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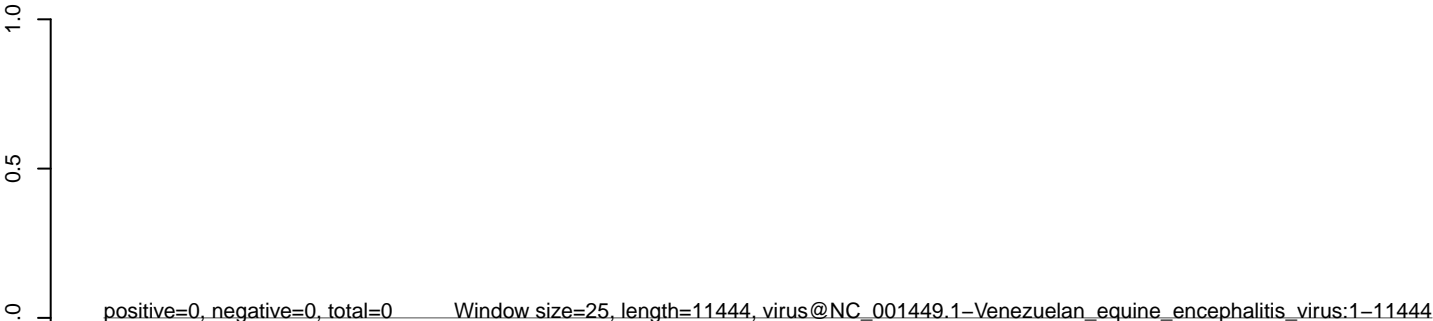
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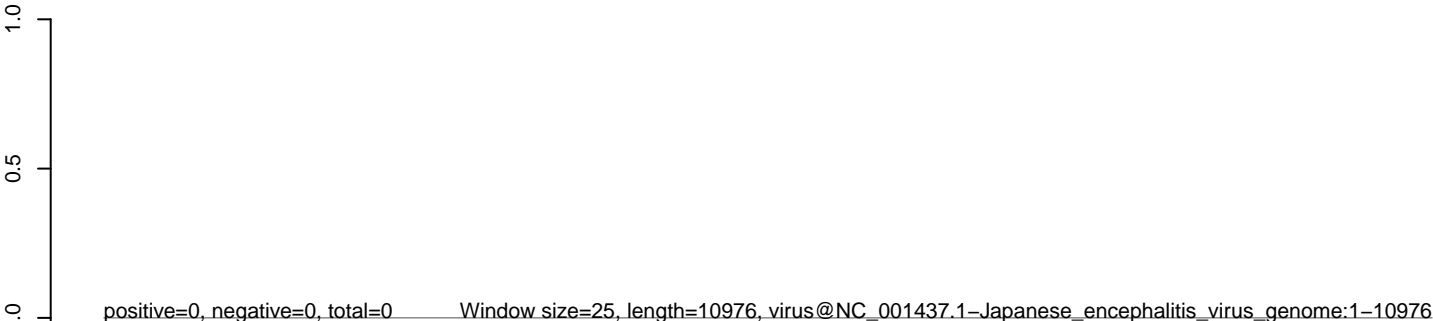
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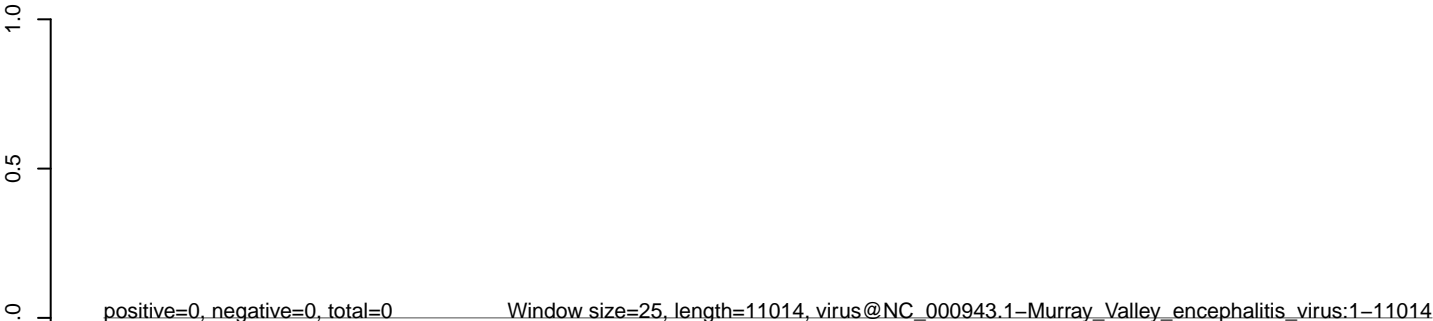
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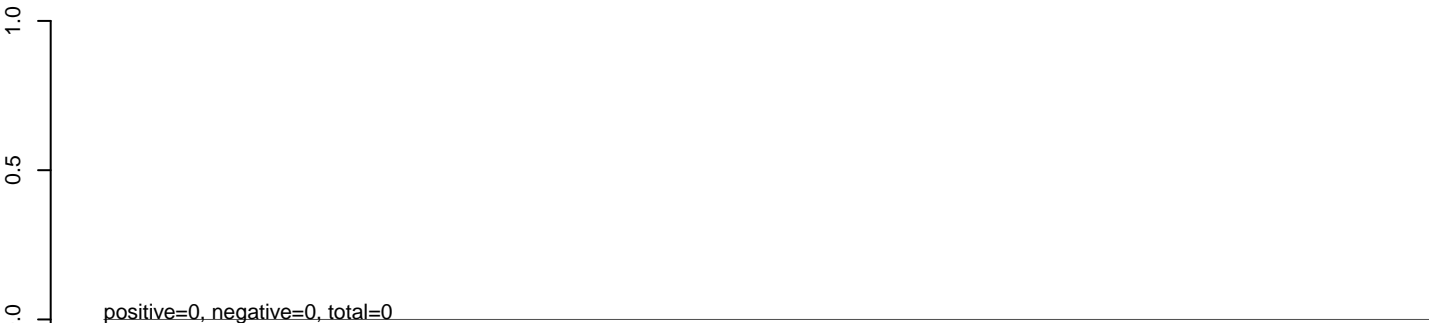


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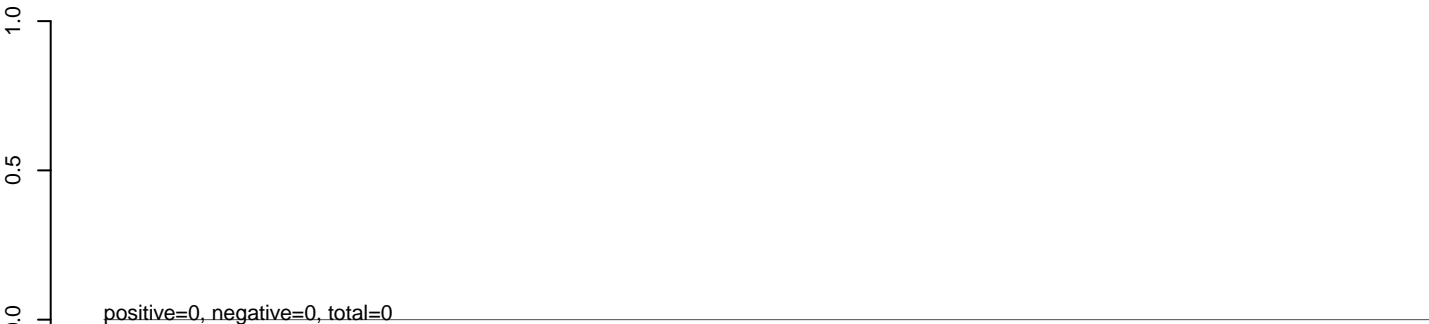


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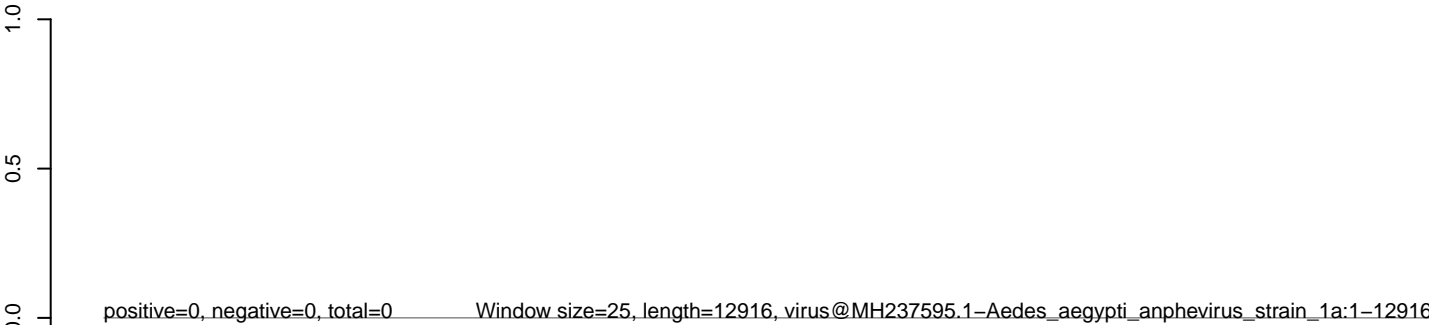
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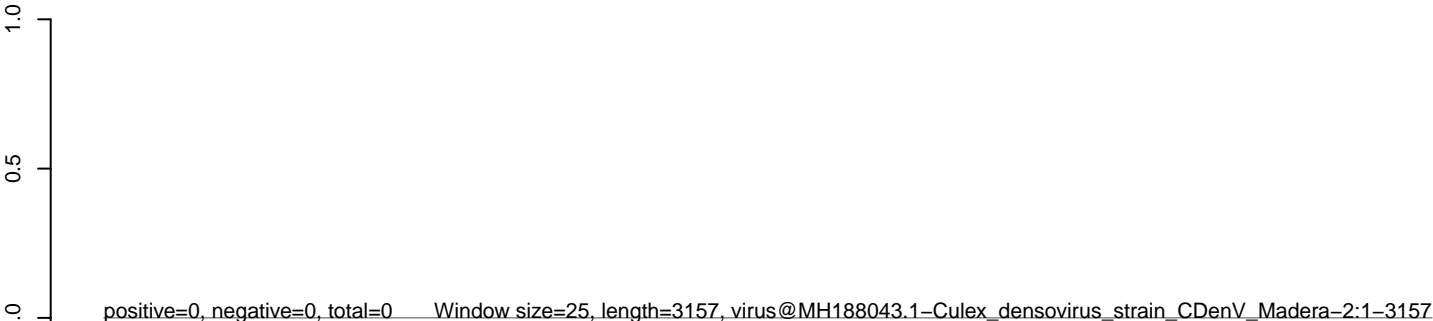
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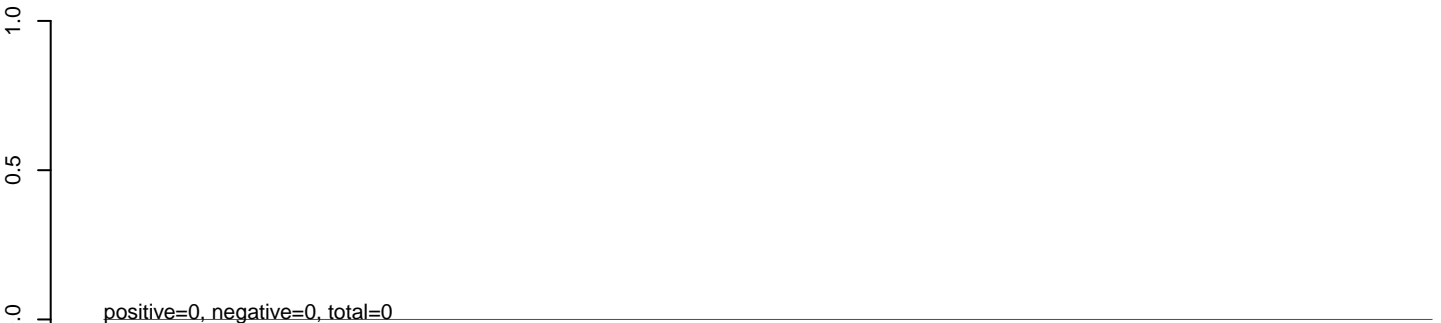


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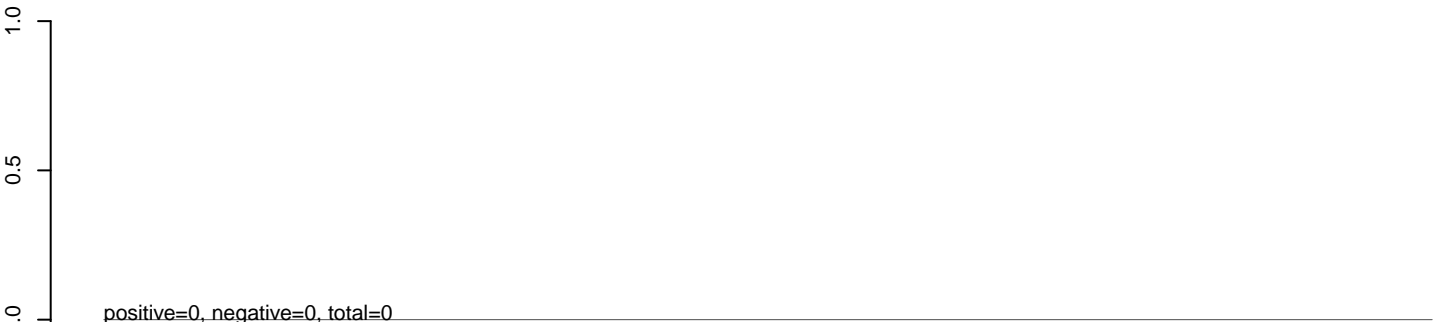


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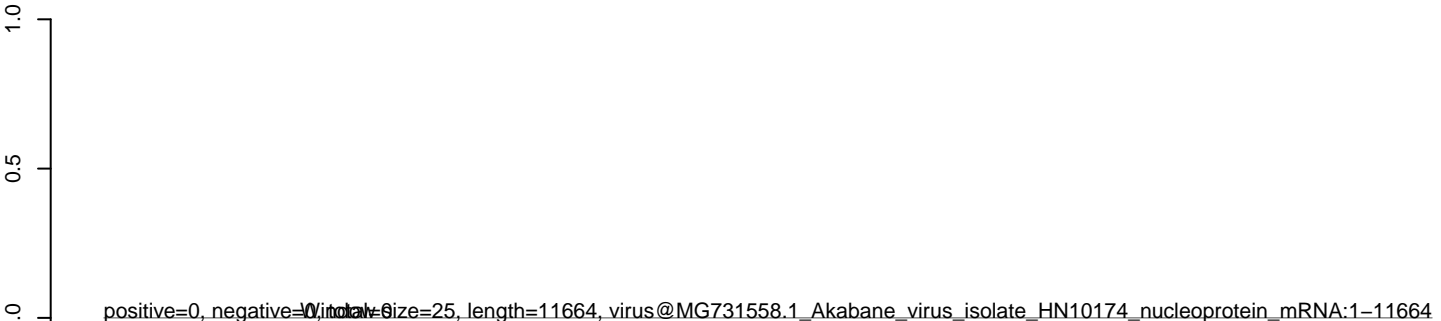
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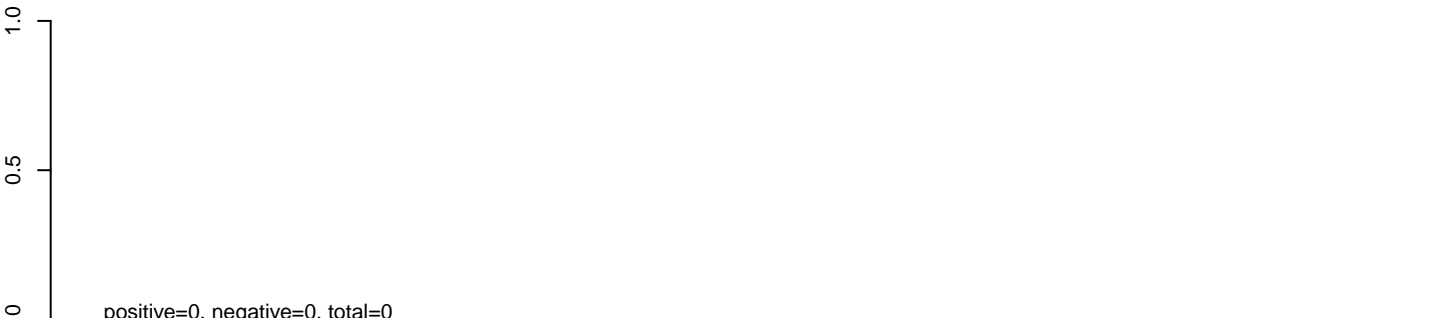


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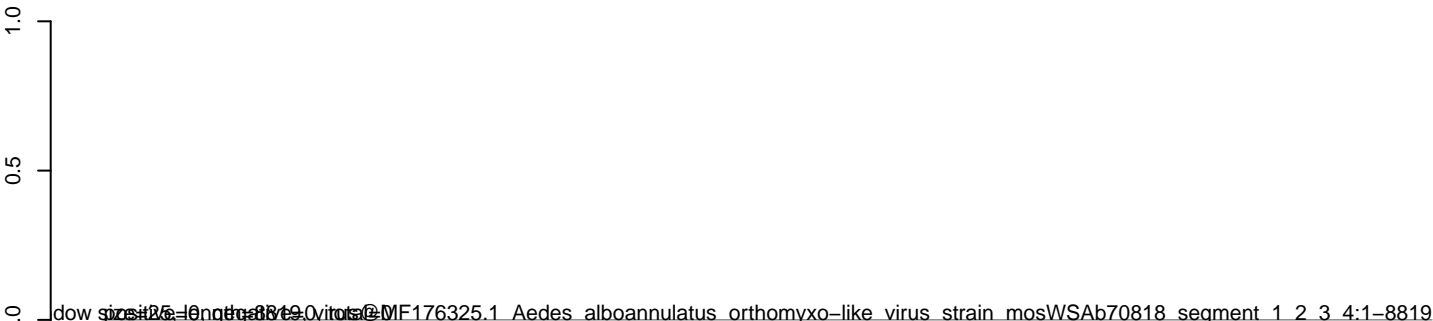
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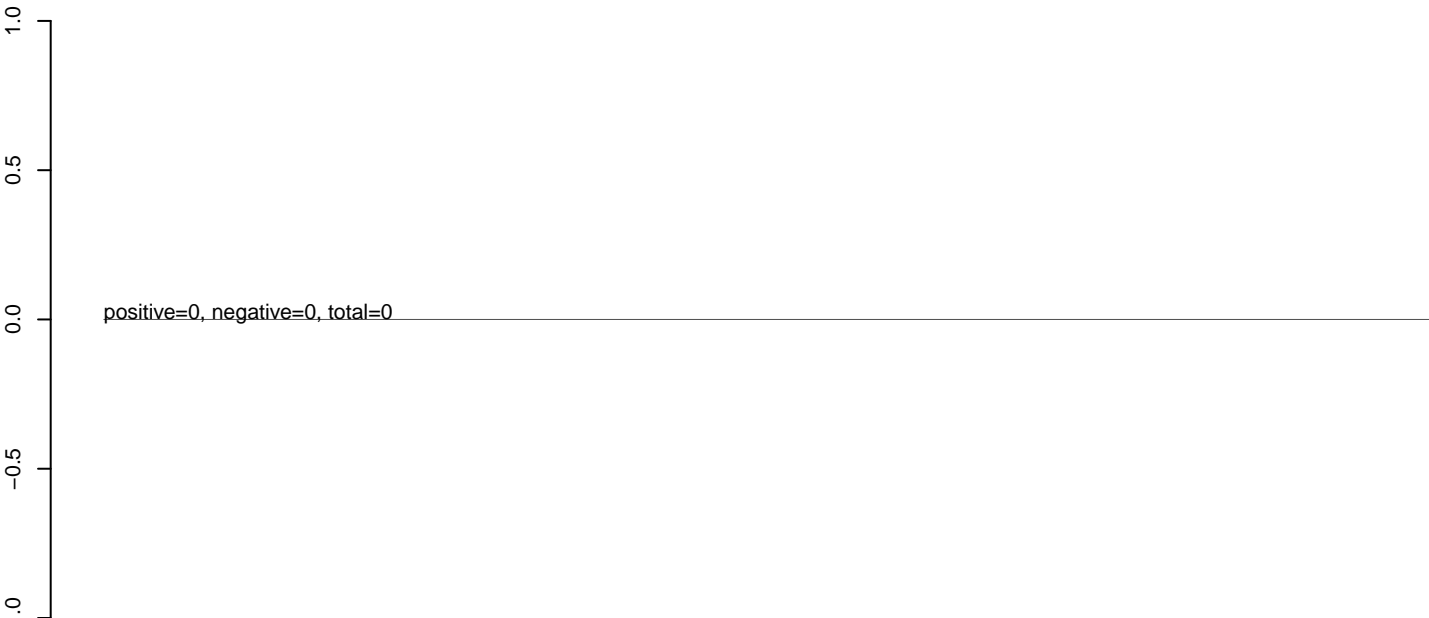


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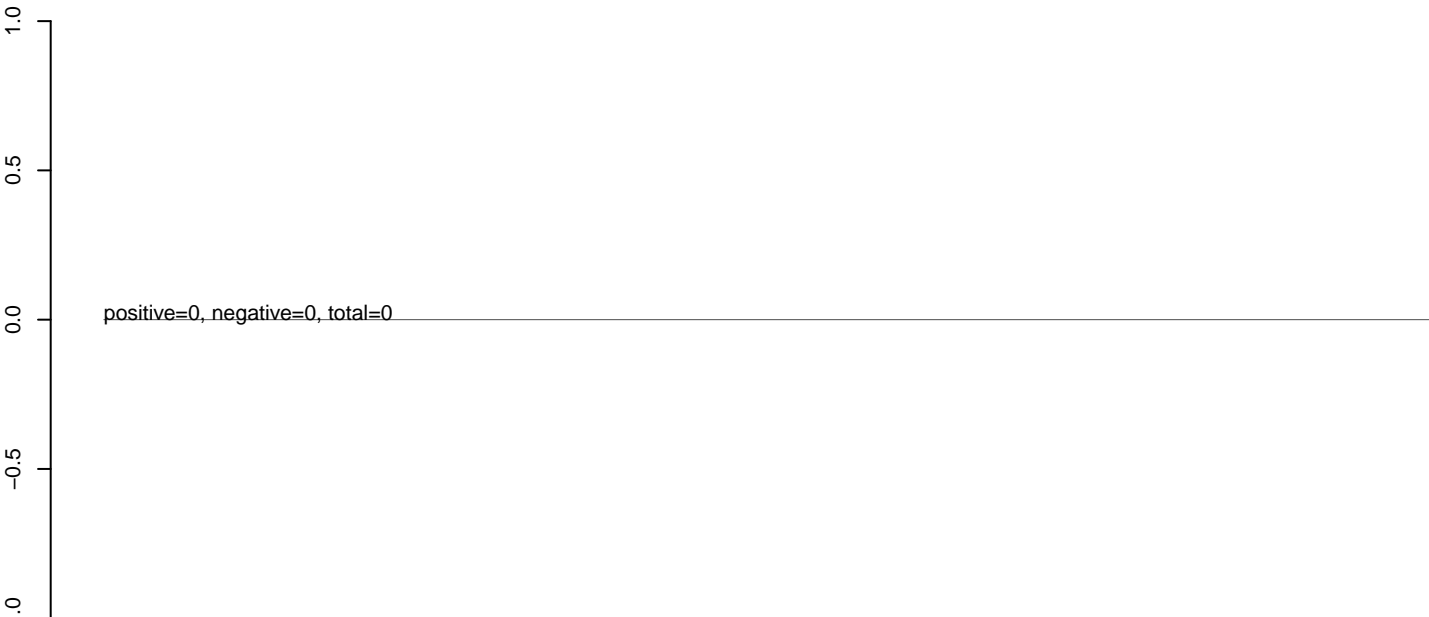


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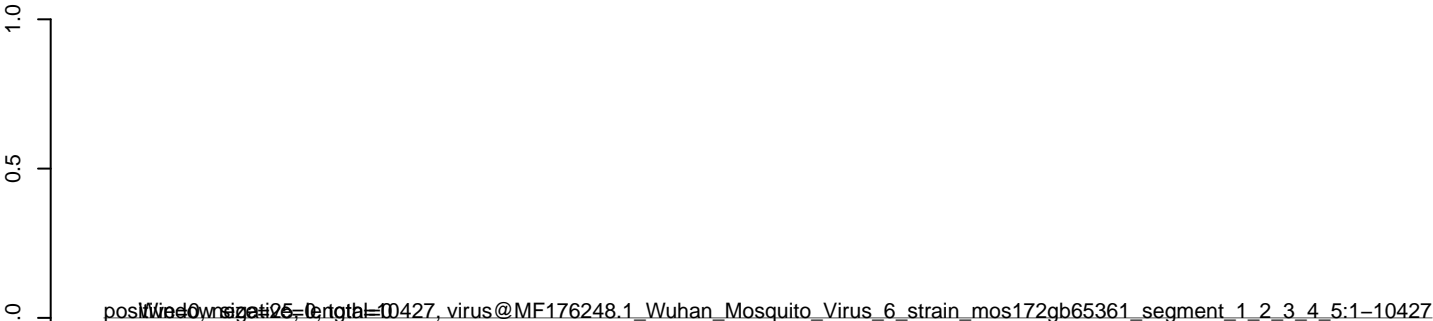
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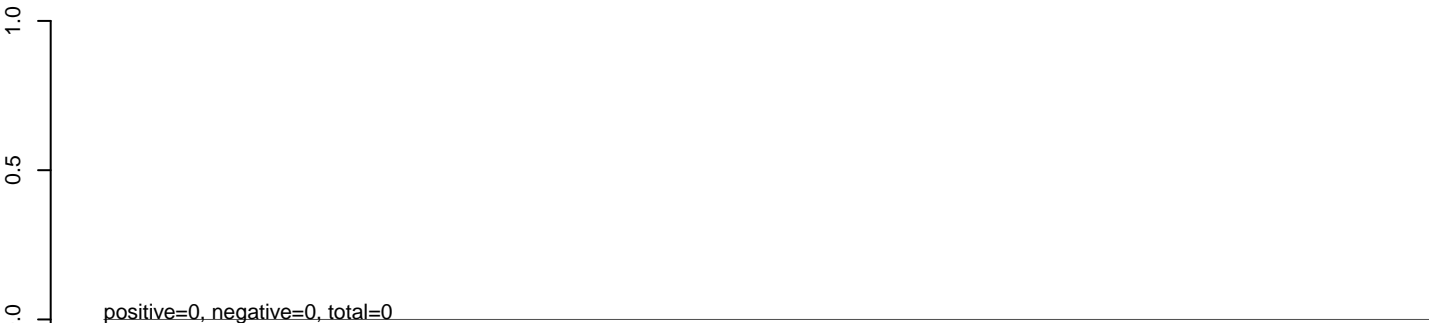


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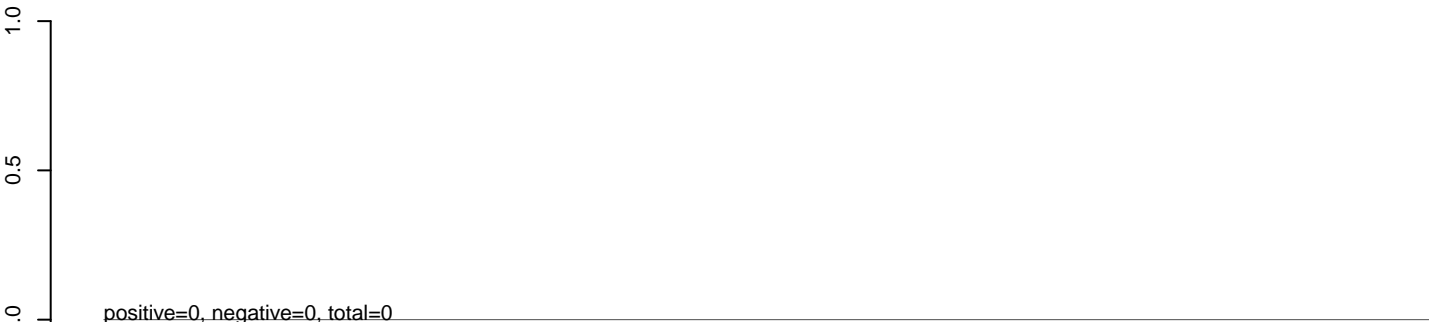


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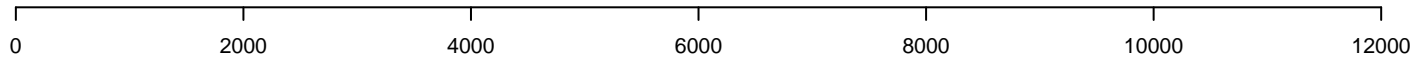
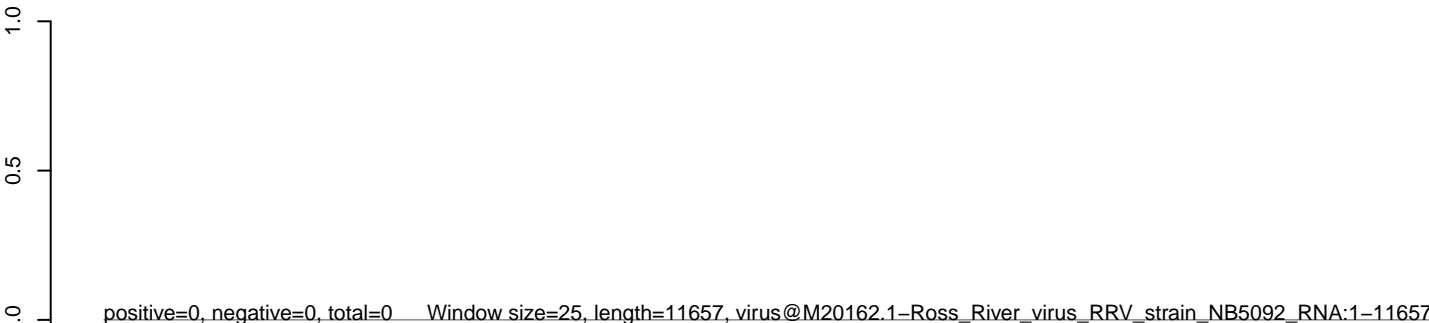
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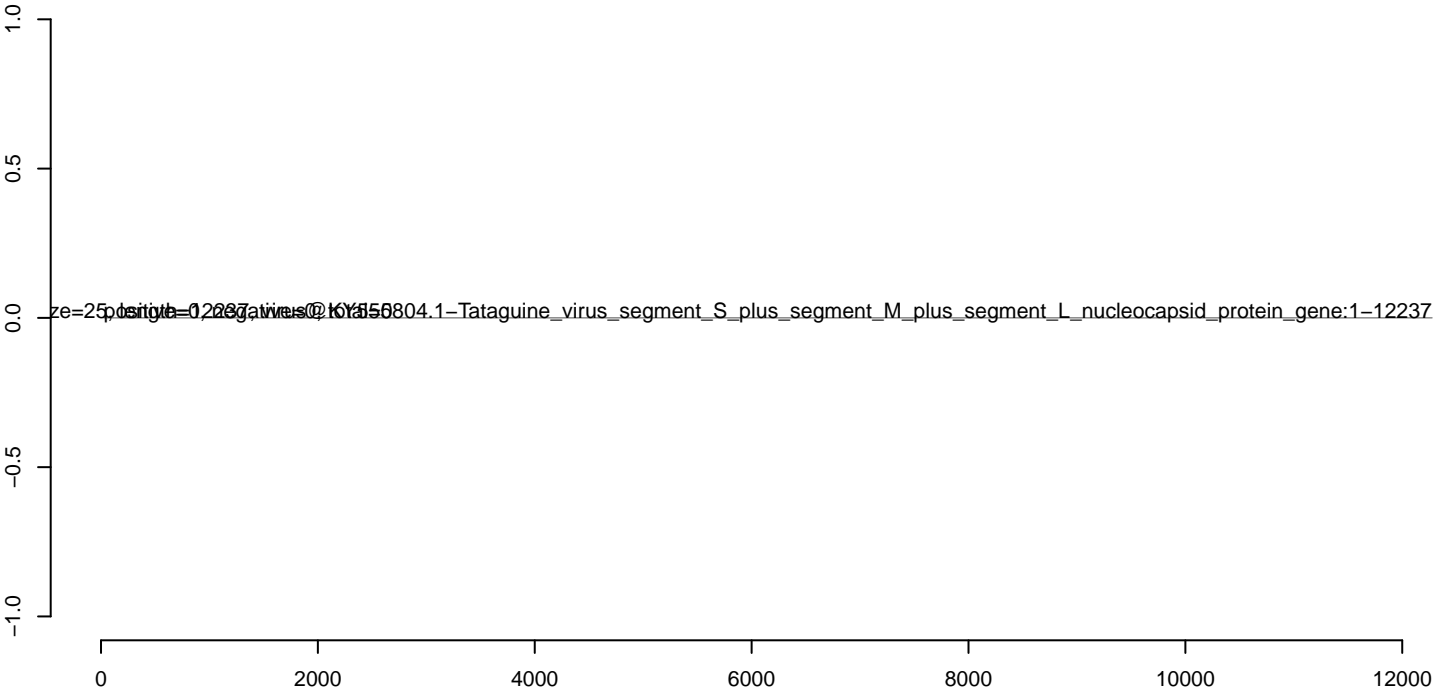
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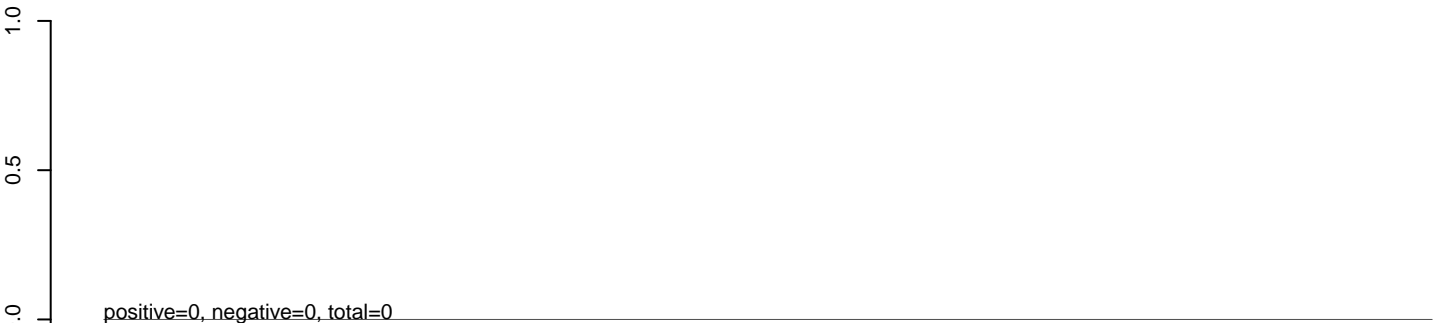
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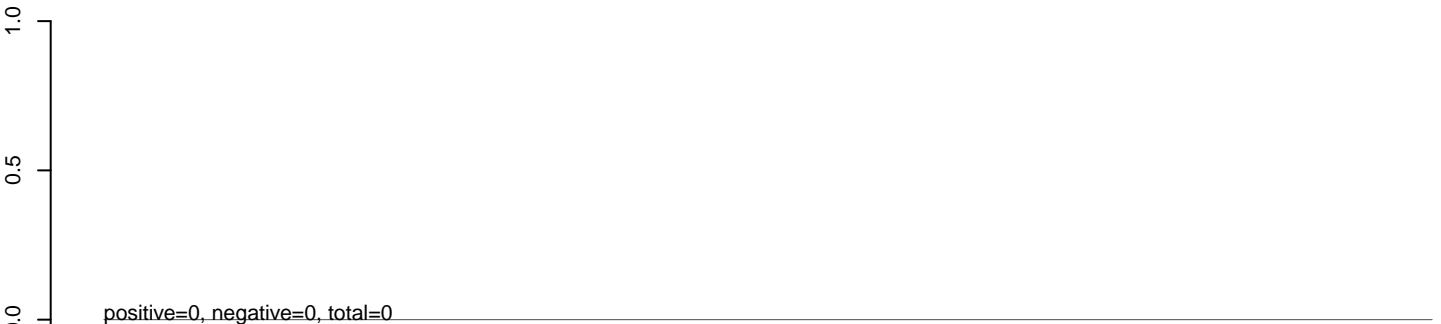
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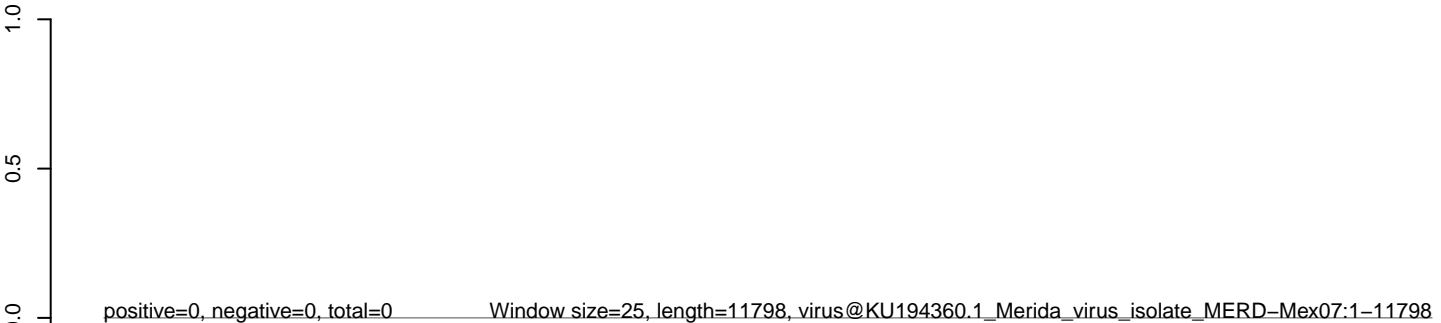
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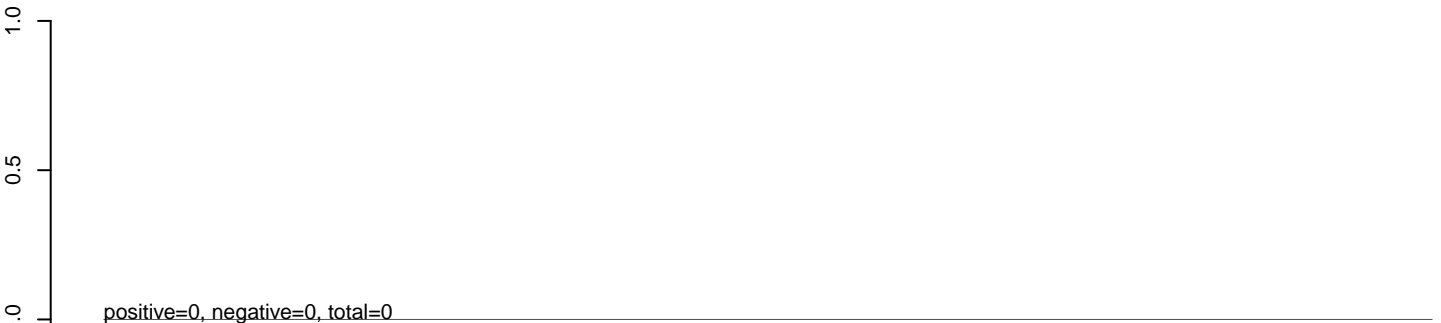


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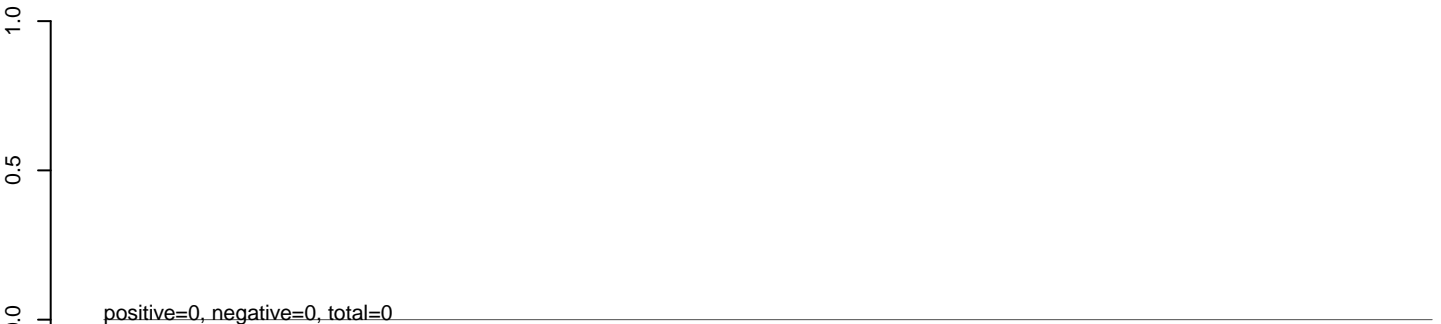


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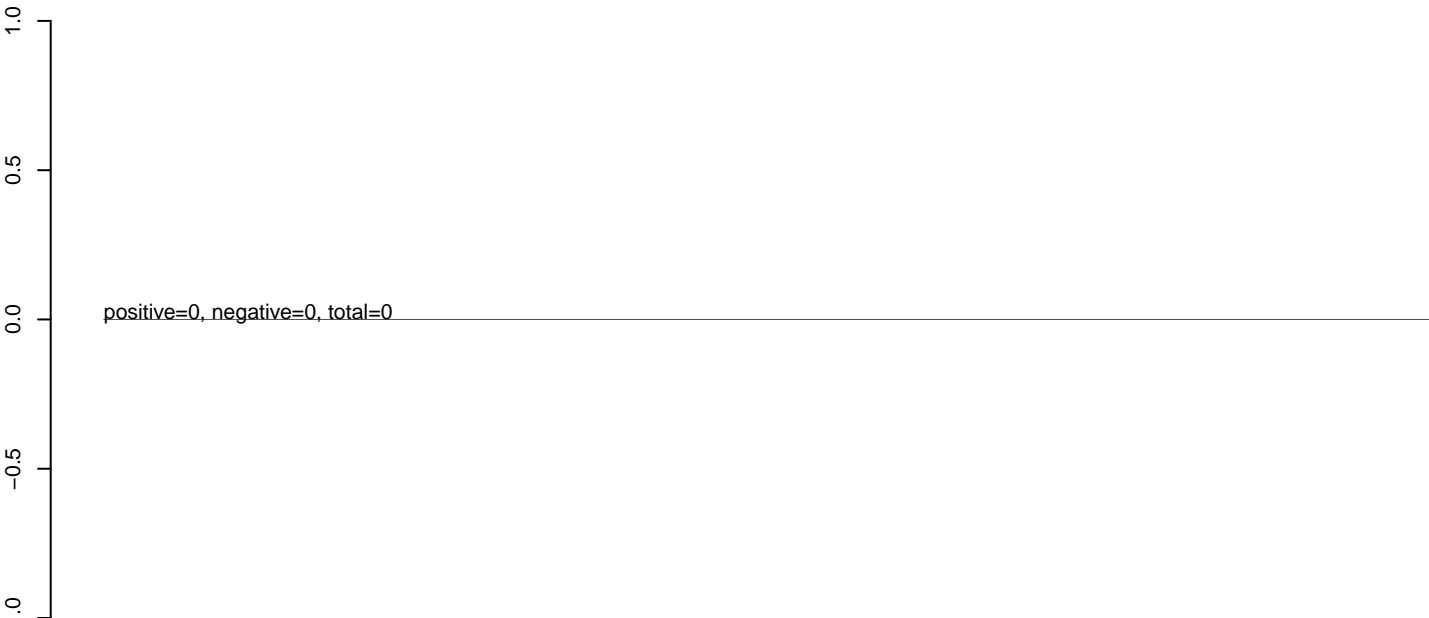
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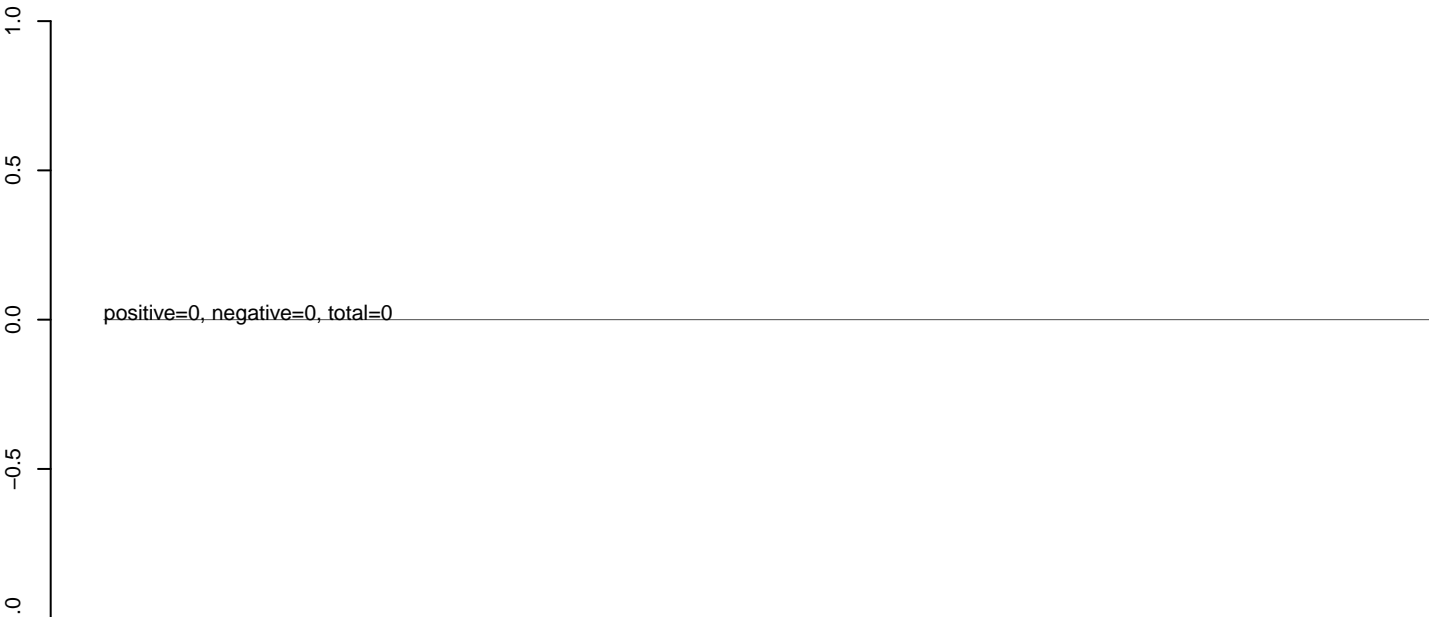
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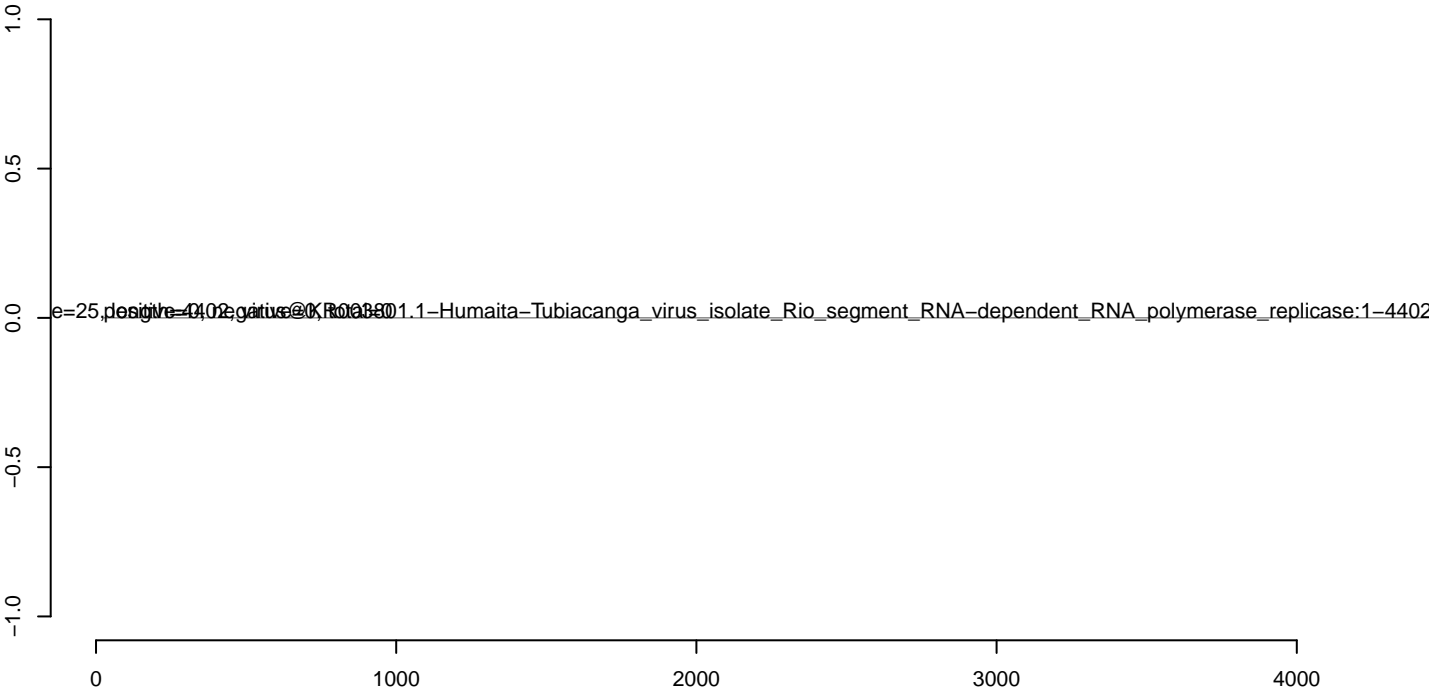
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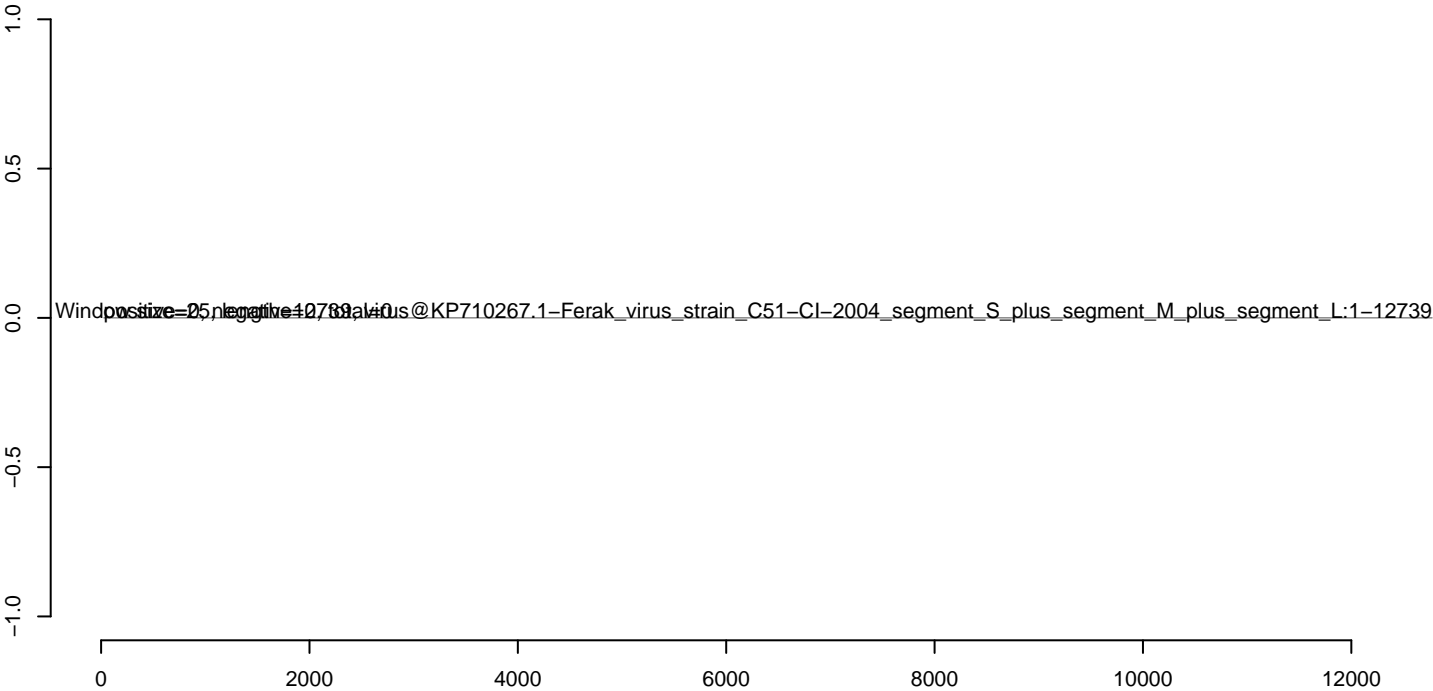
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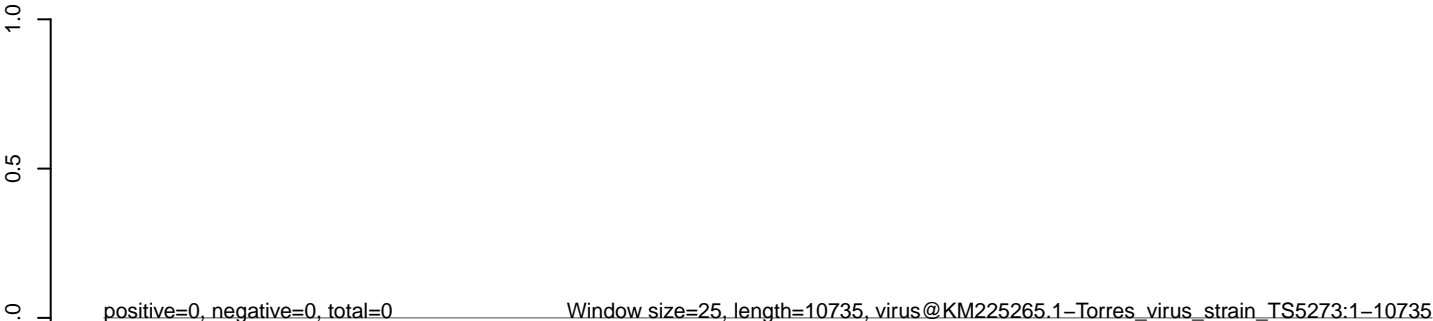
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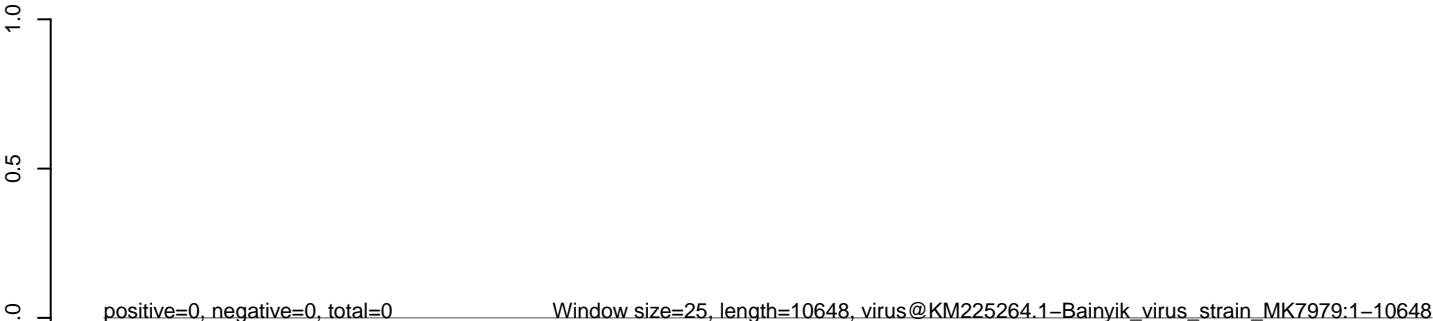
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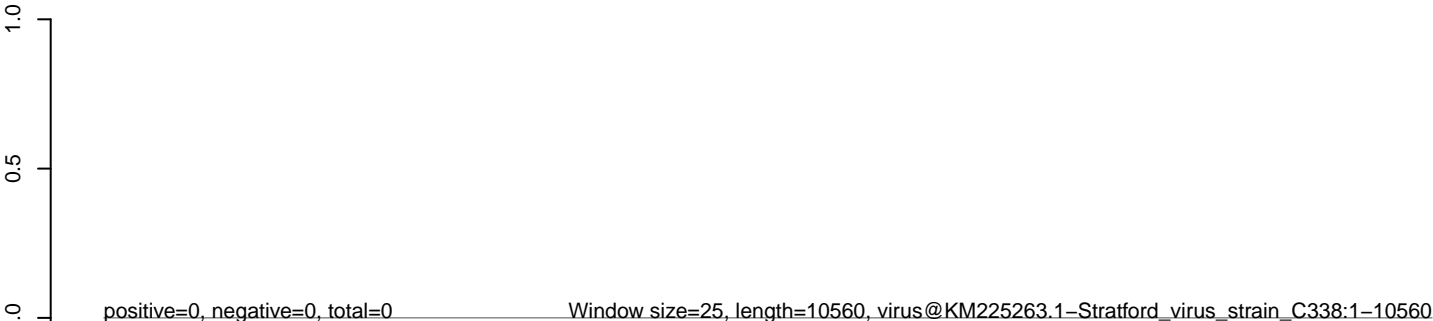
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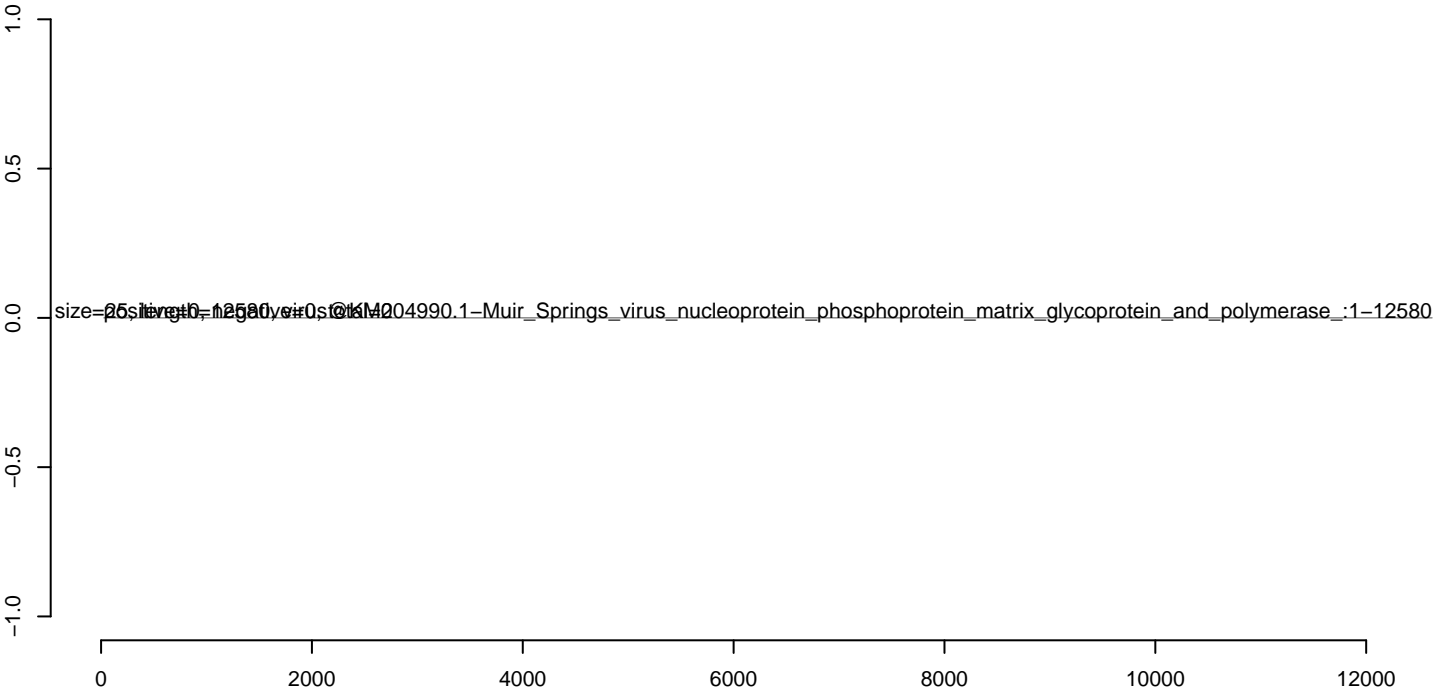
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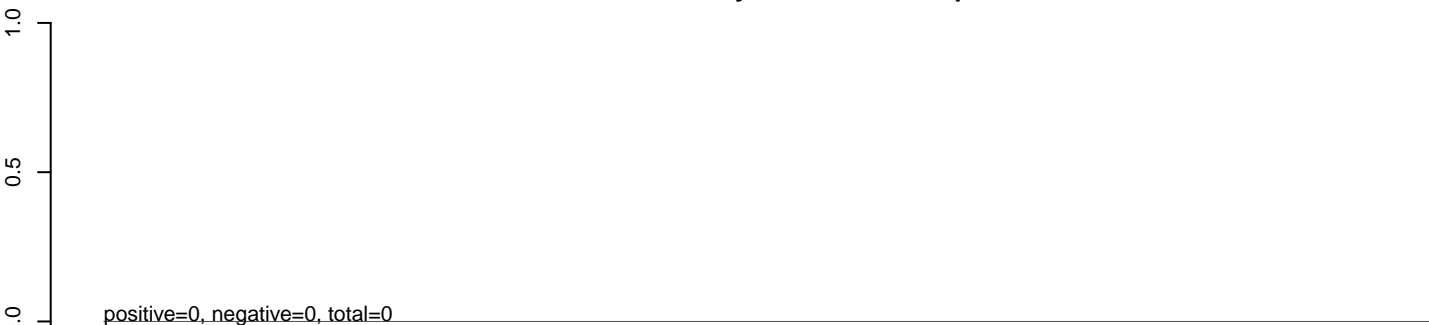
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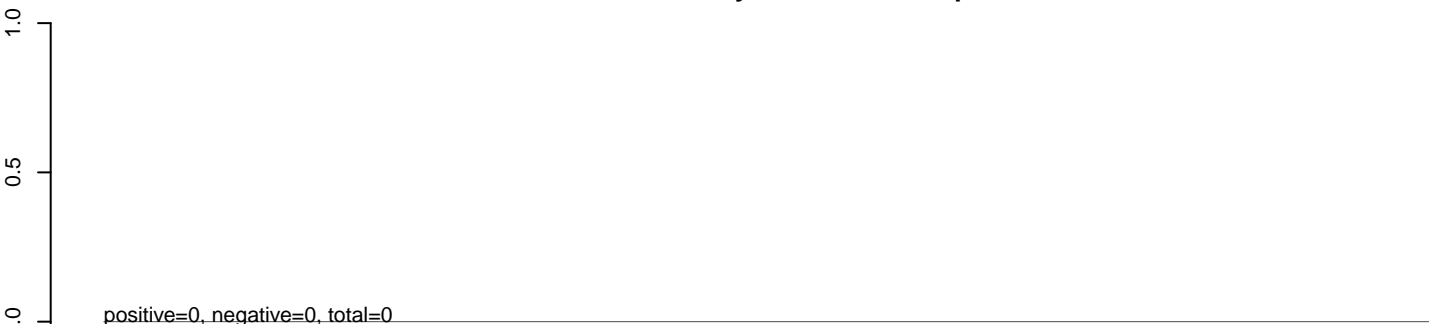
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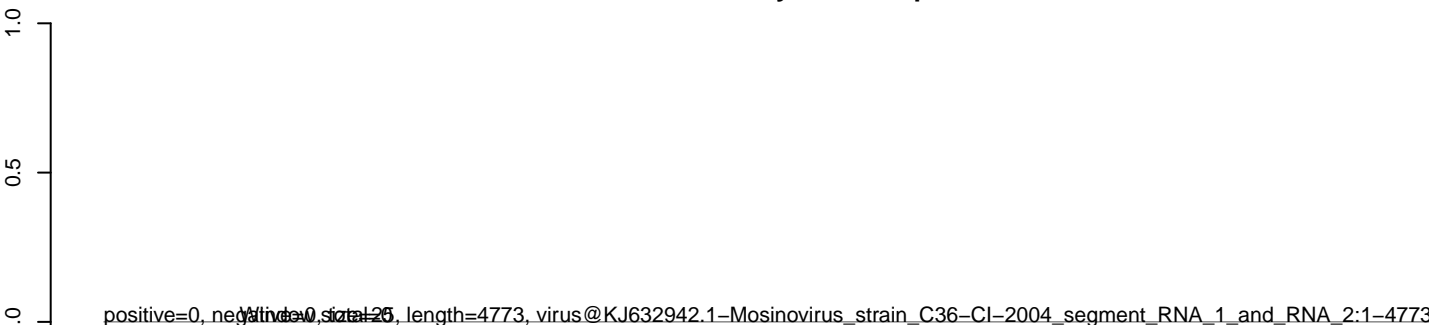
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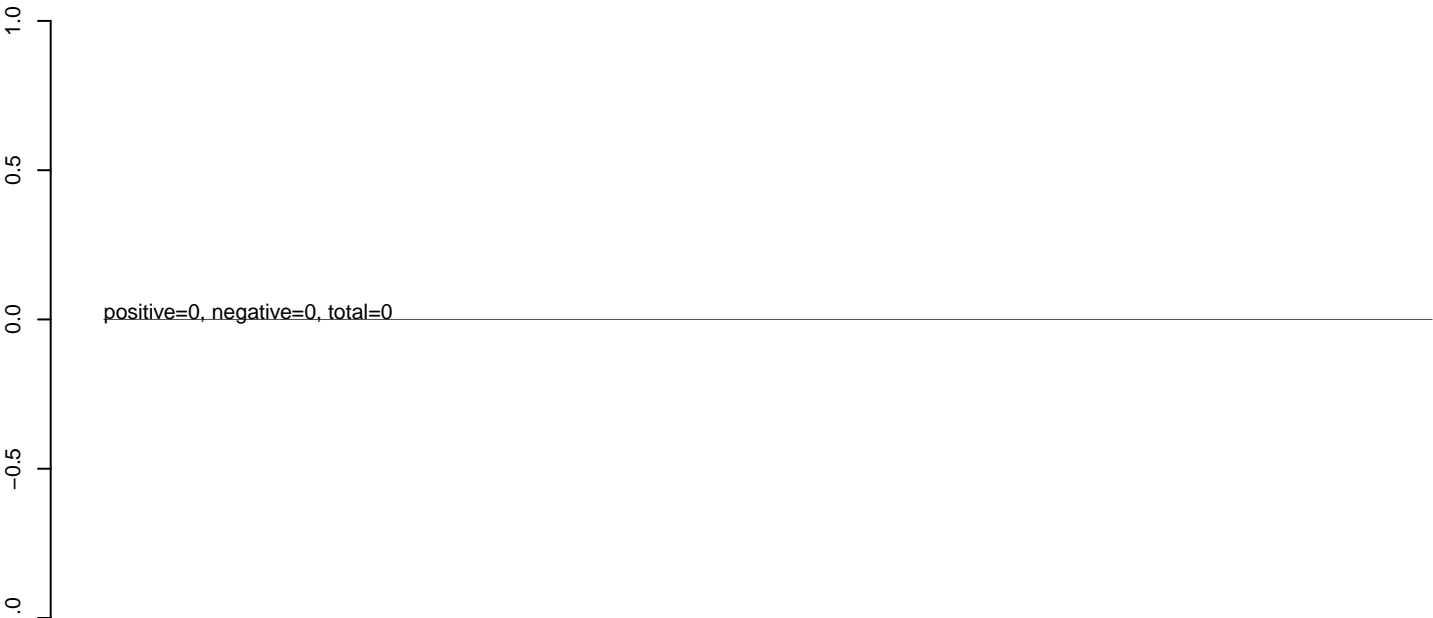


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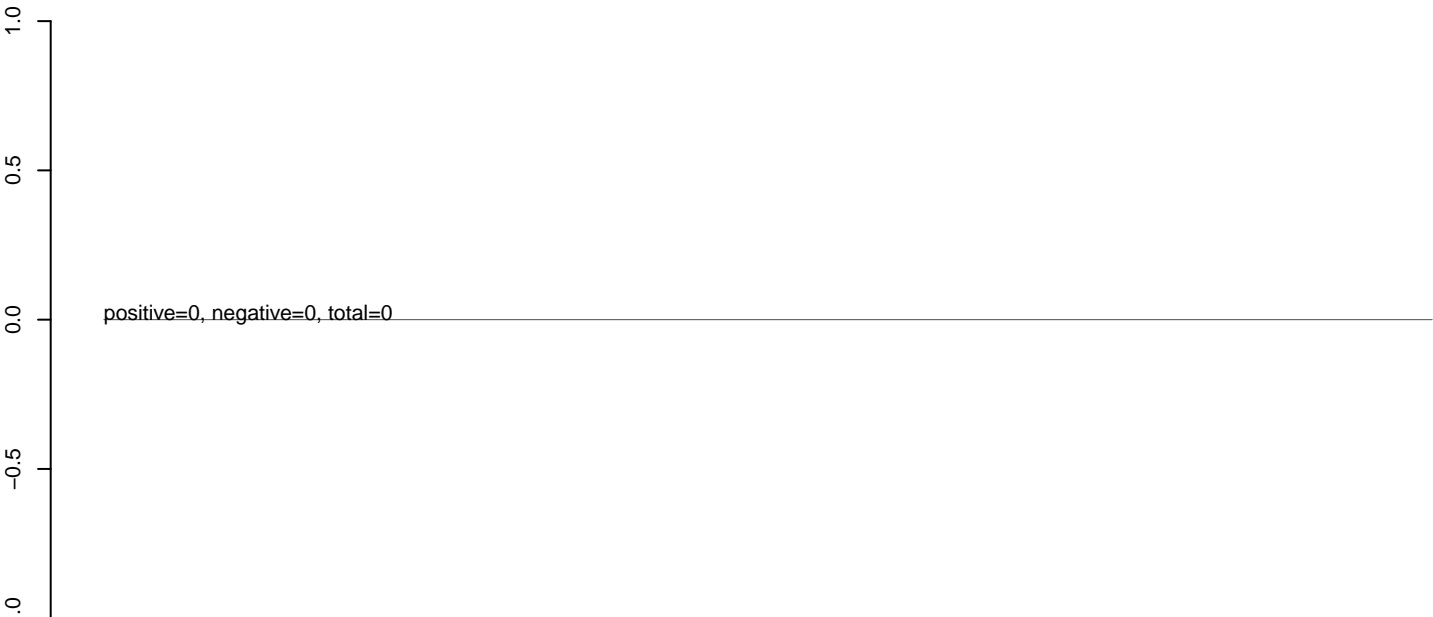


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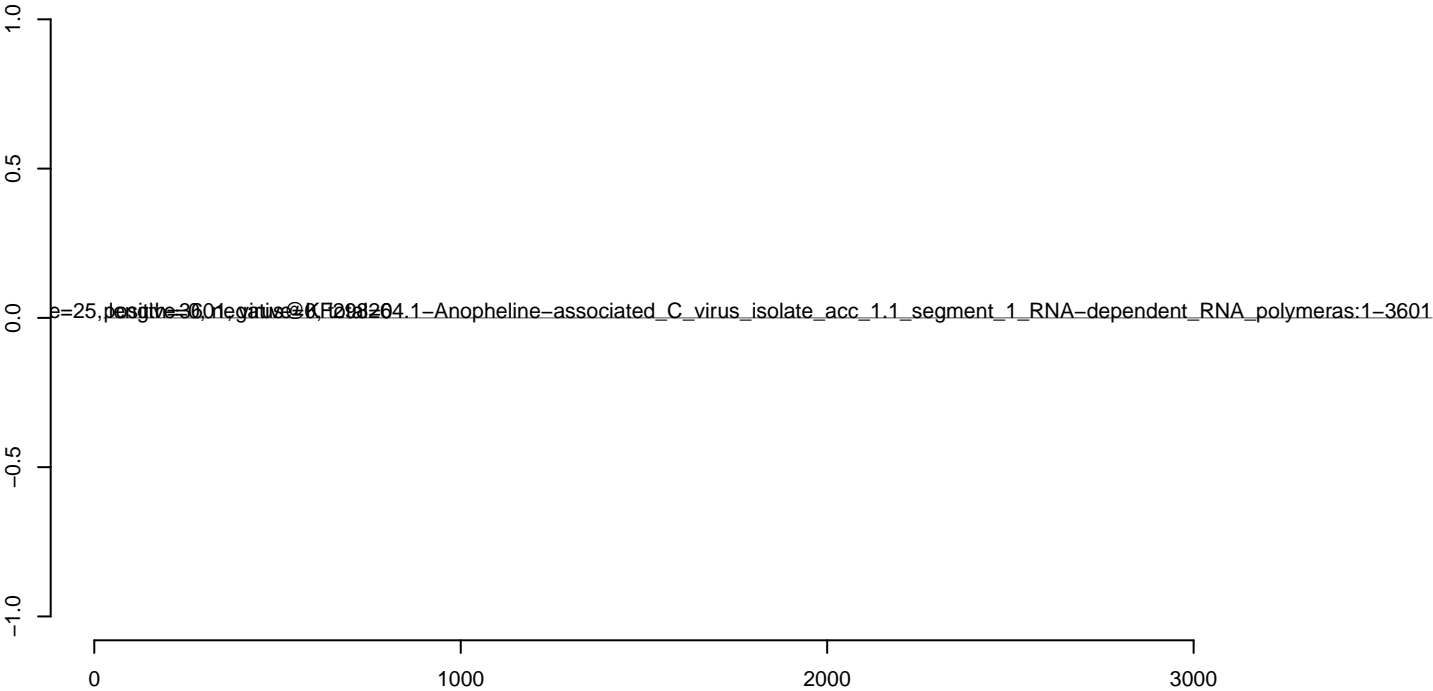
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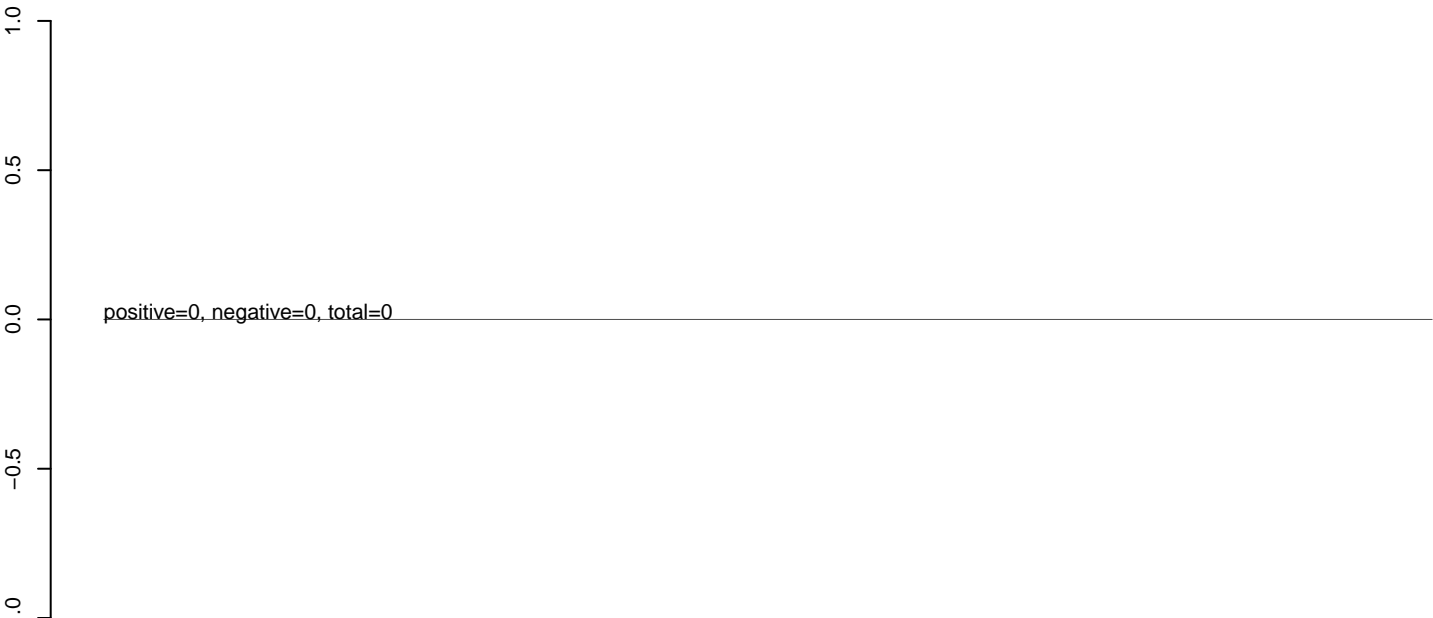
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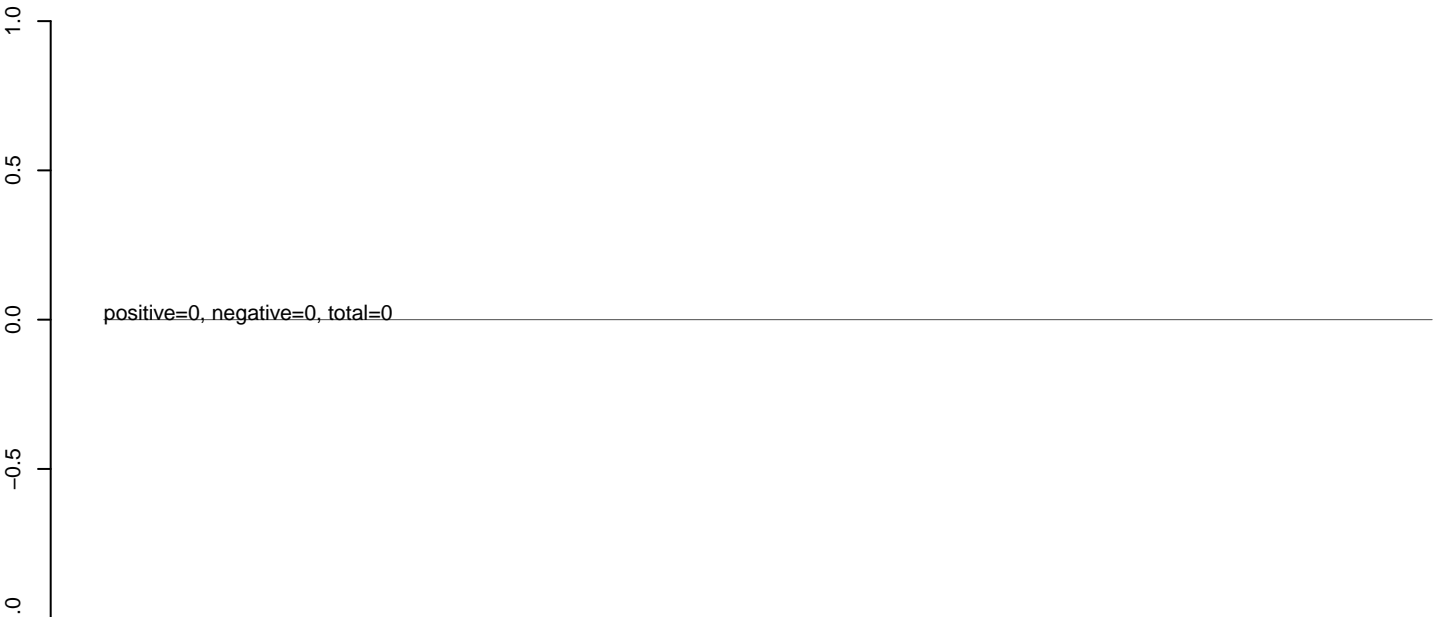
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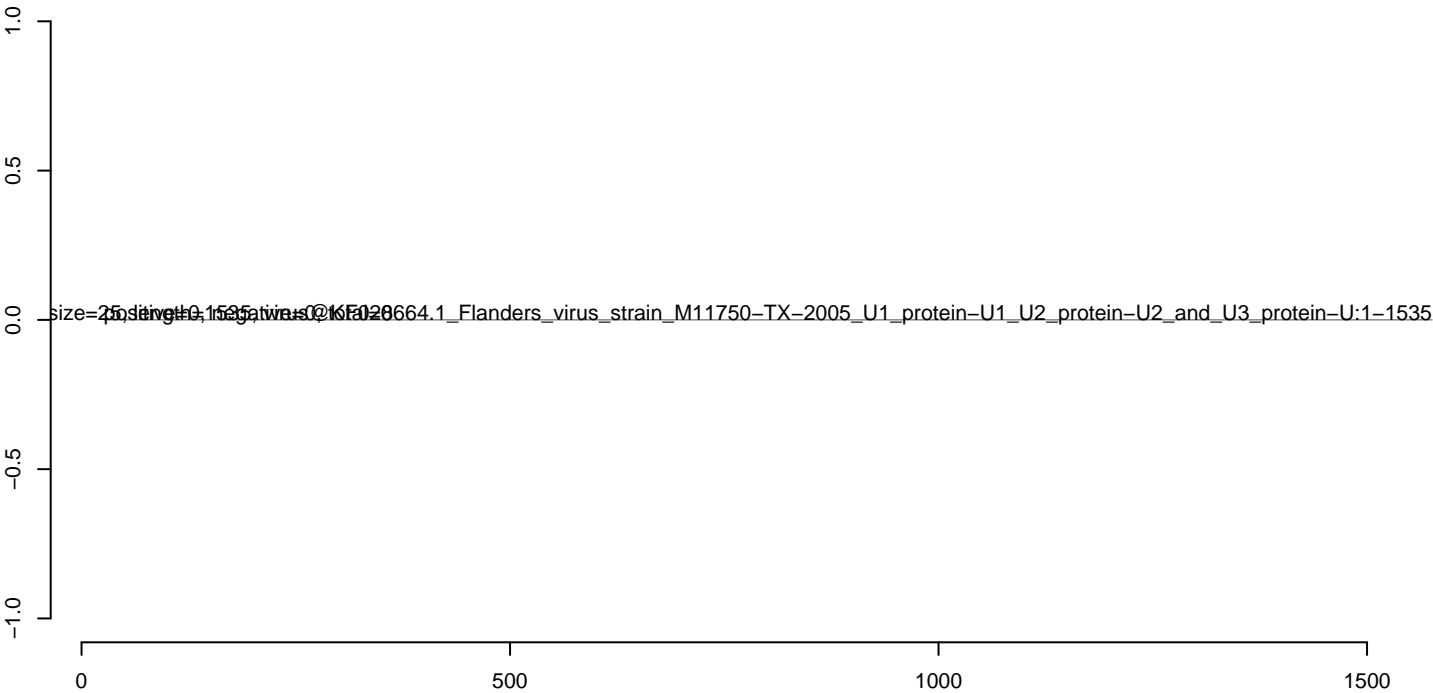
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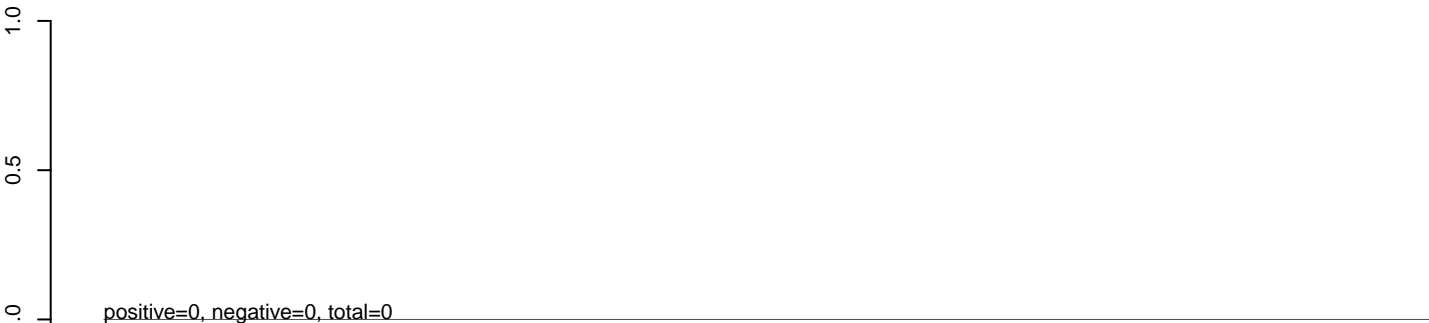
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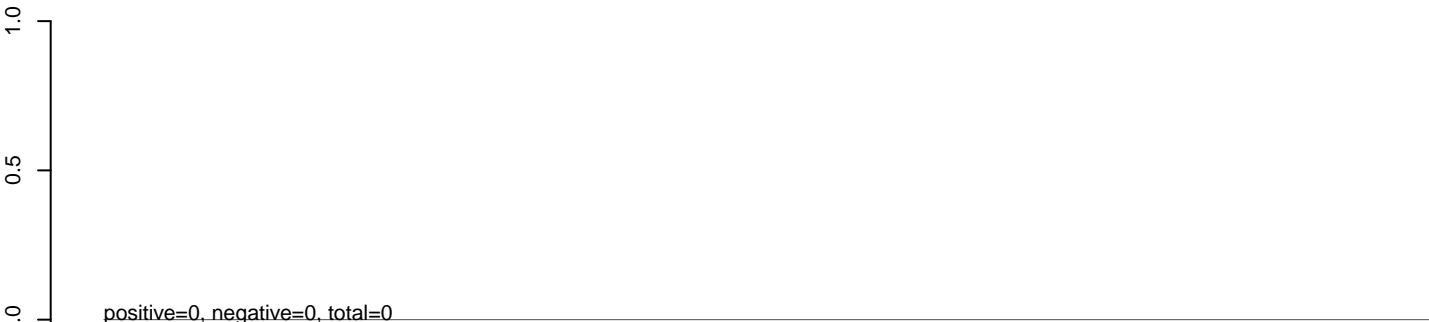
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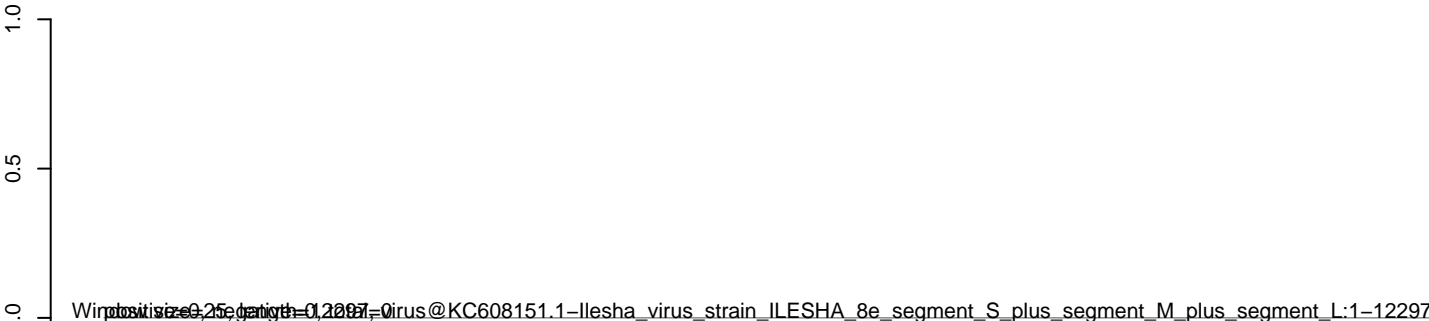
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AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



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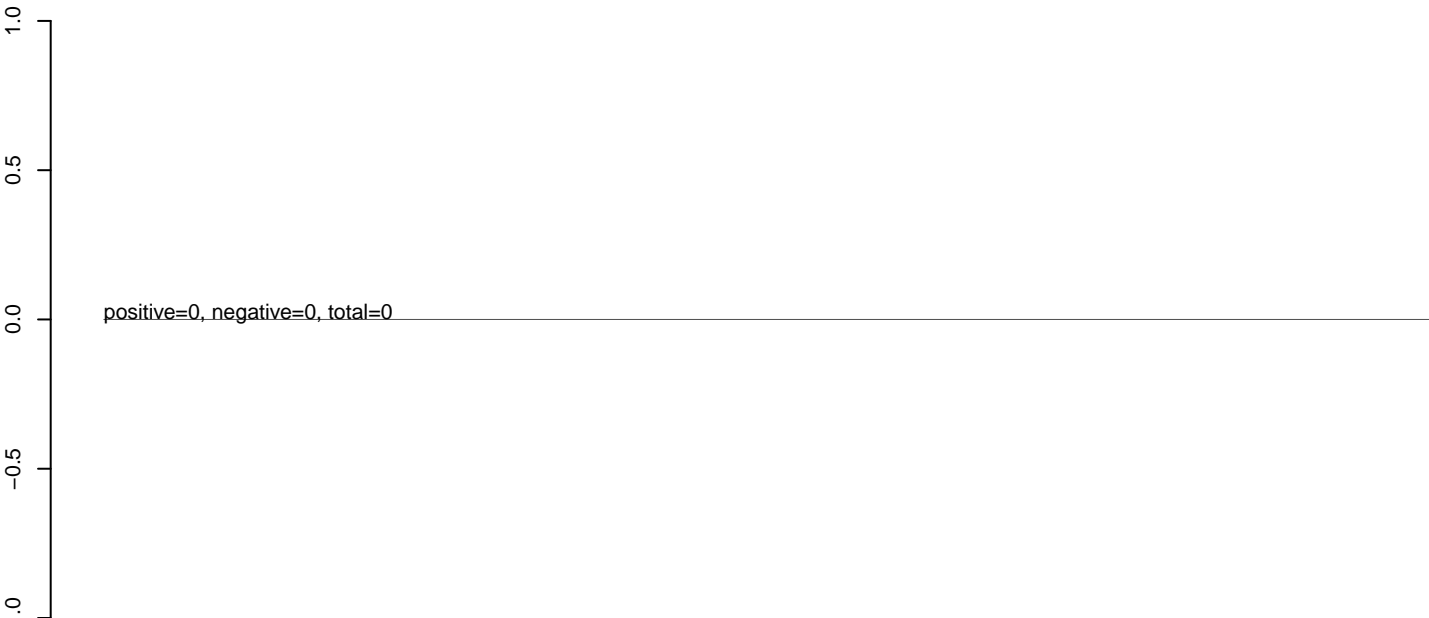


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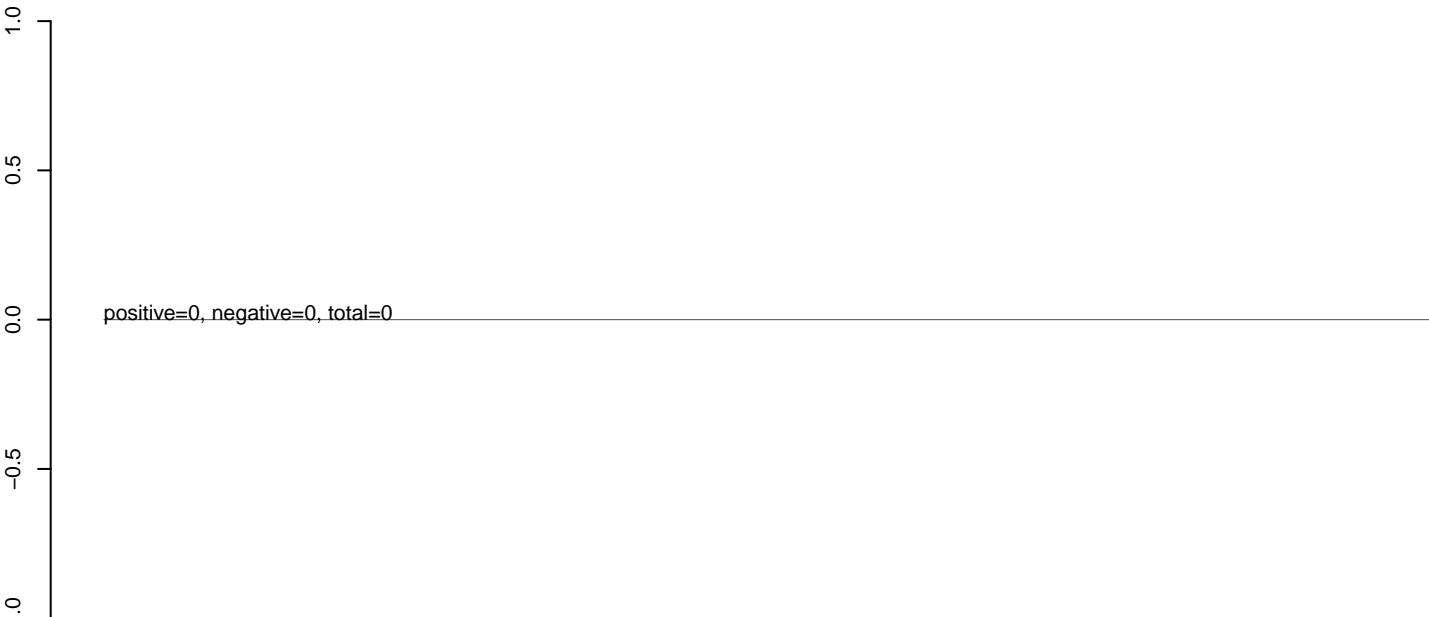


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AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



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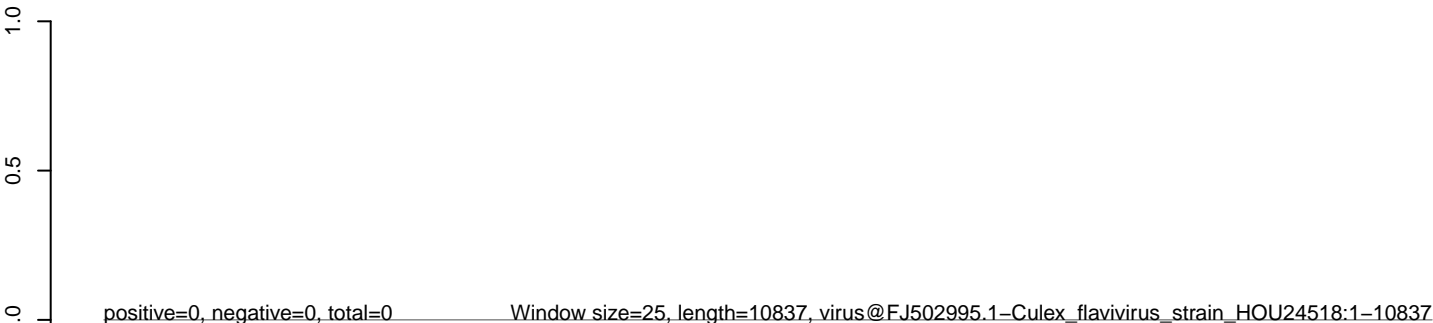
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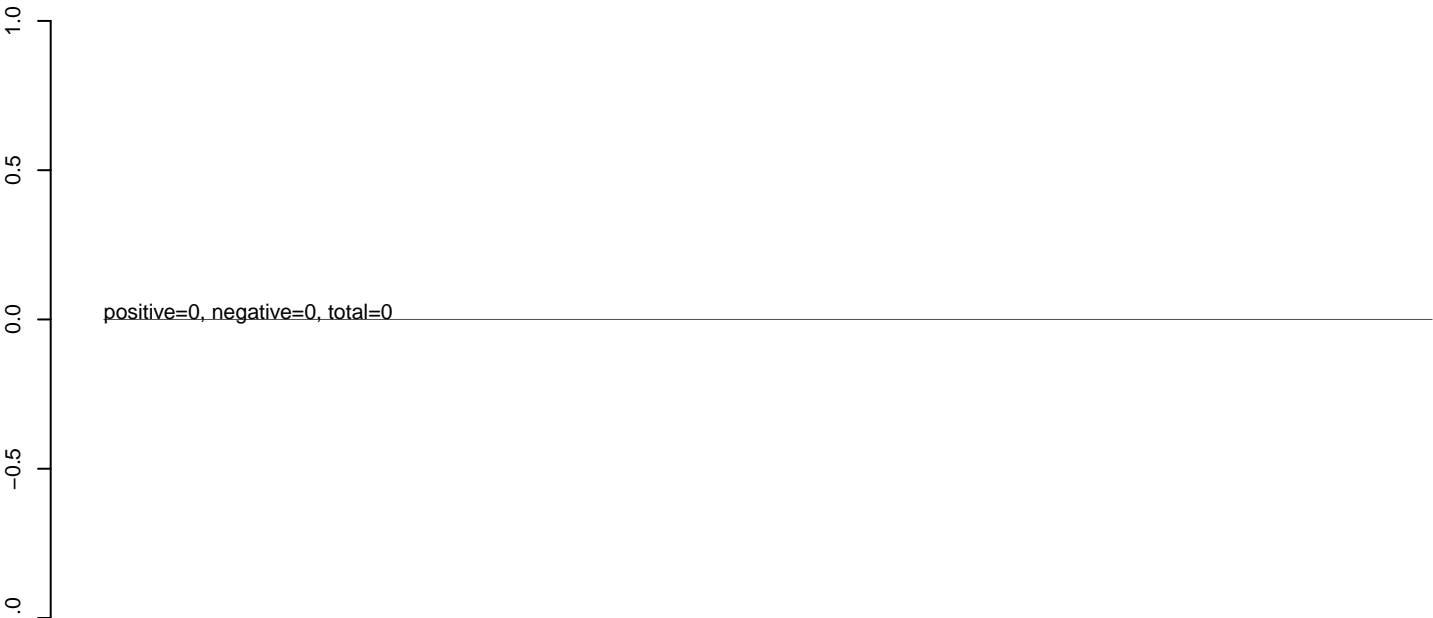


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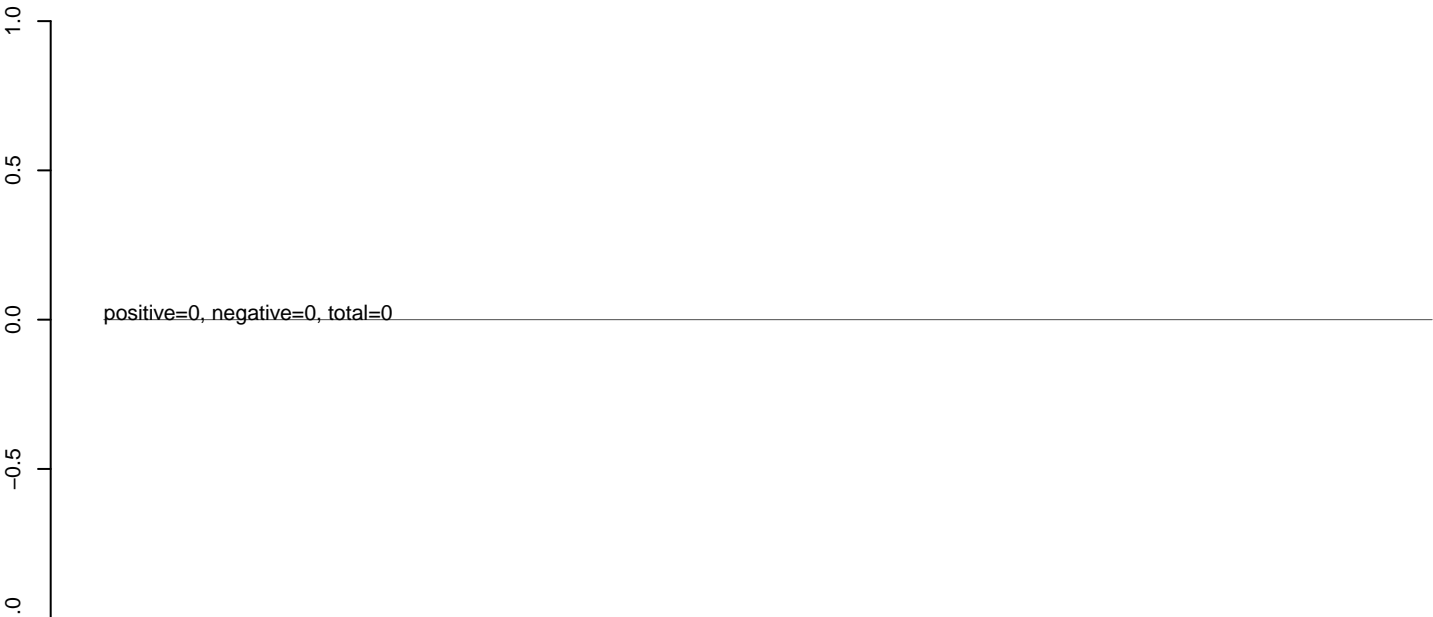


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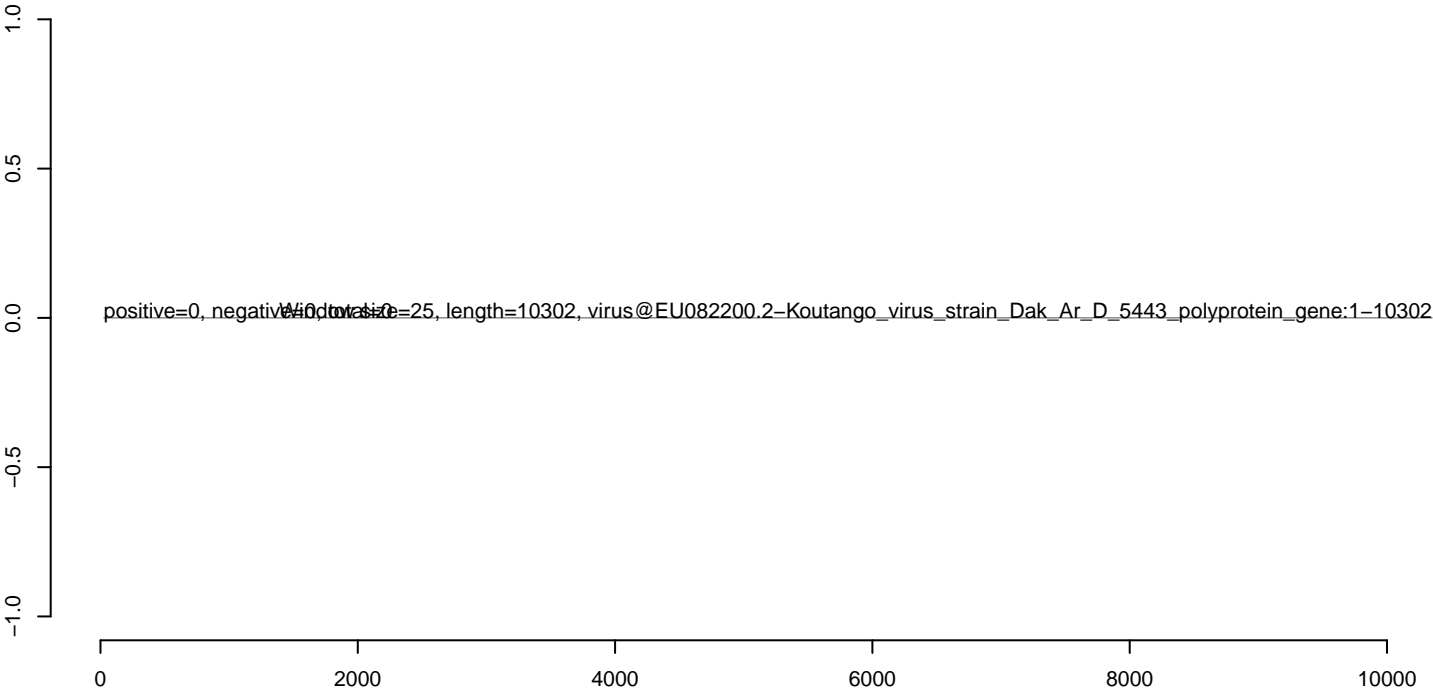
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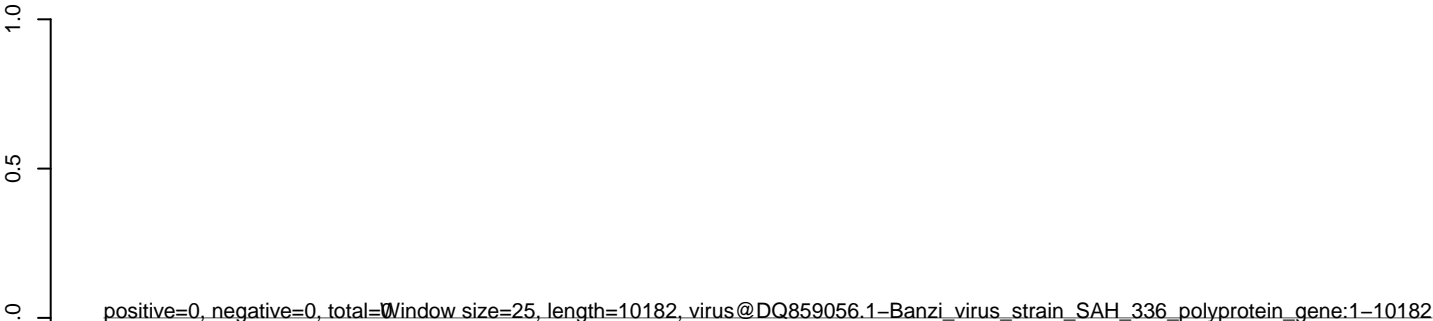
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AnGam_Ovary_BetaE.rep



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AnGam_Ovary_BetaE.24_35.rep



AnGam_Ovary_BetaE.rep



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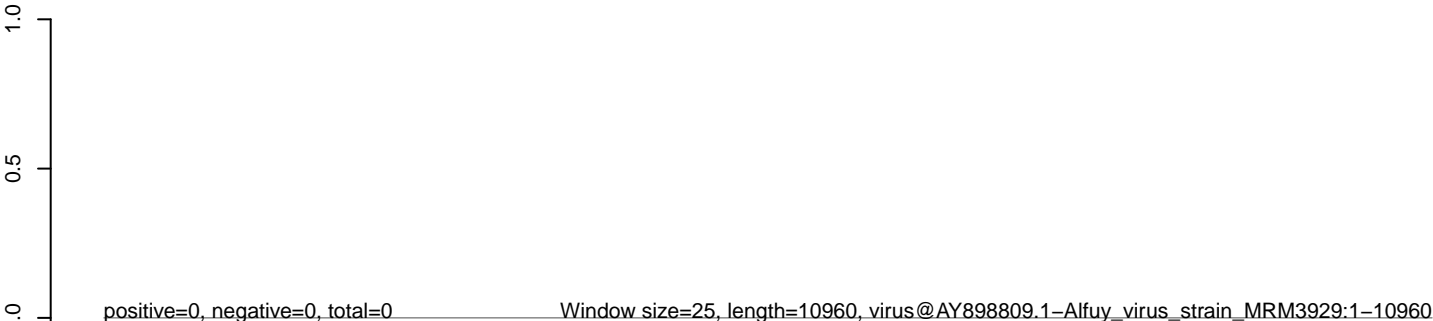
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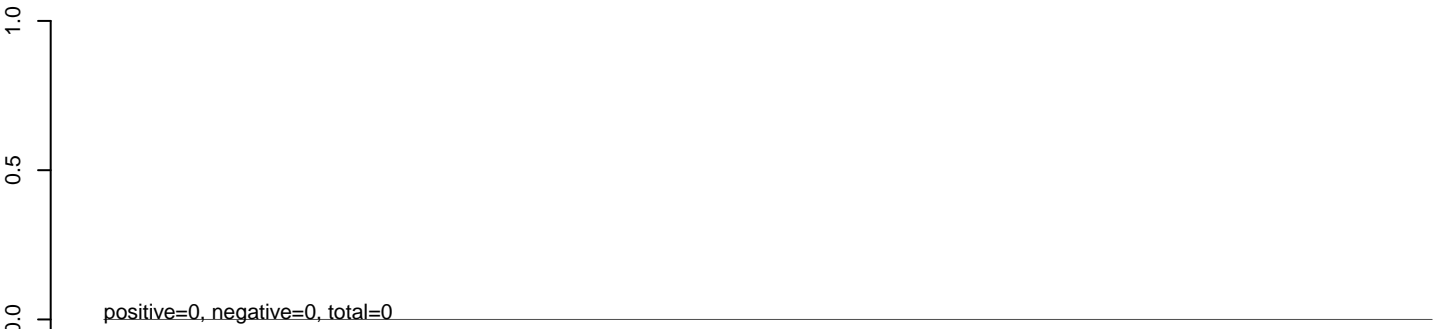


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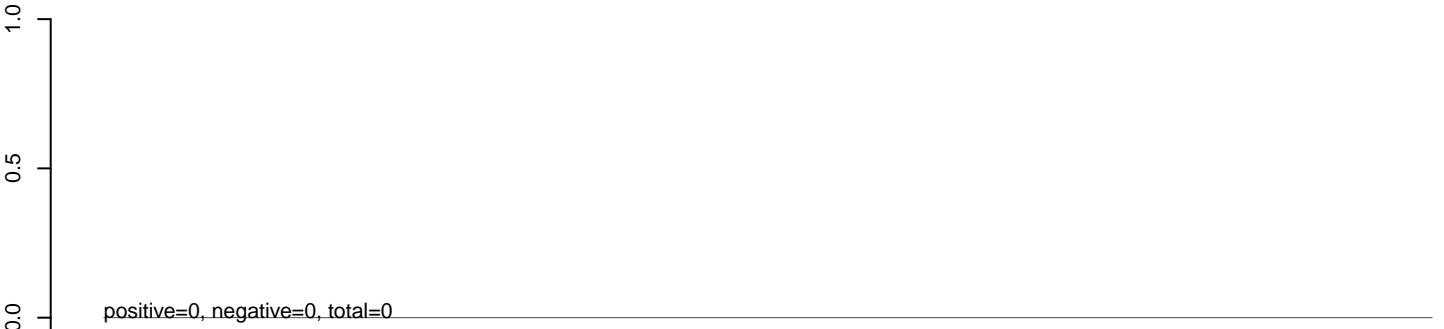


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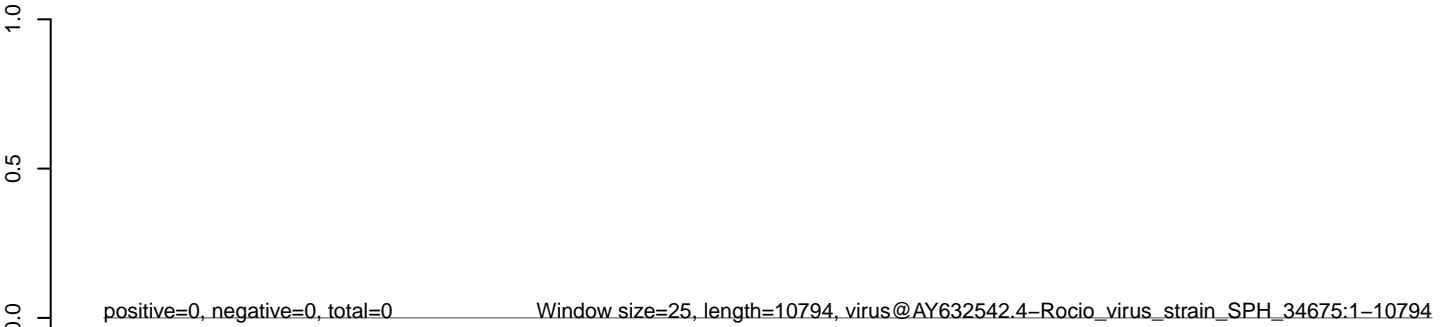
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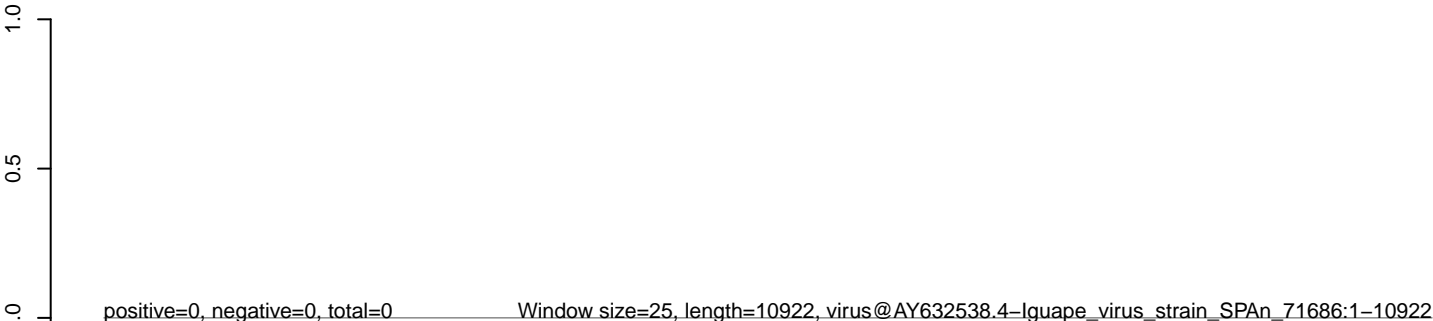
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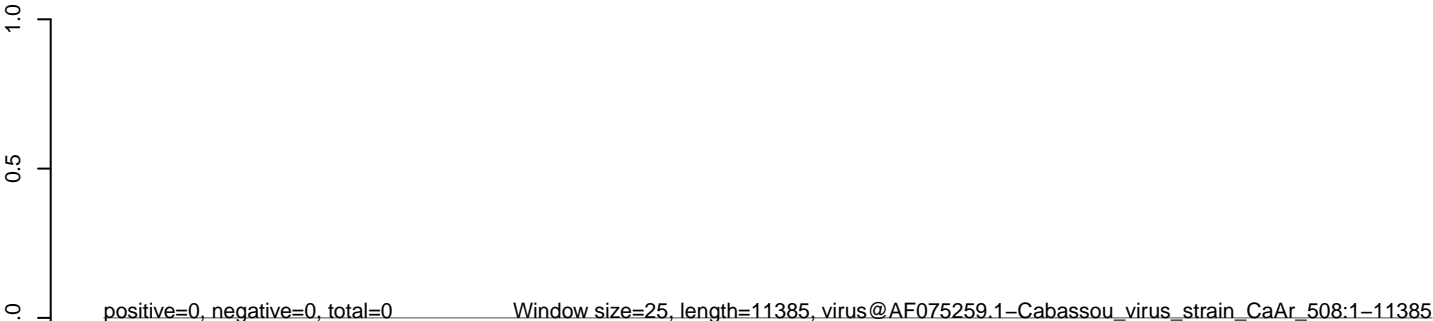
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AnGam_Ovary_BetaE.rep



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AnGam_Ovary_BetaE.rep



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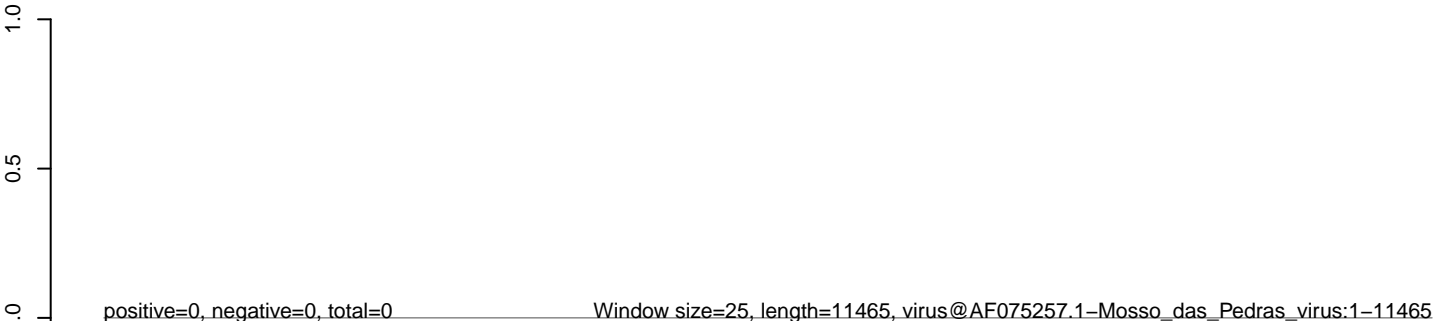
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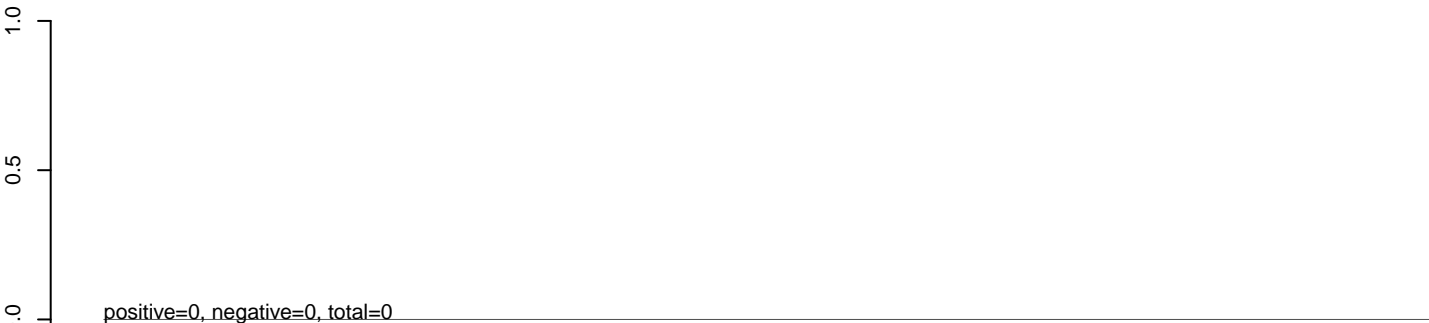


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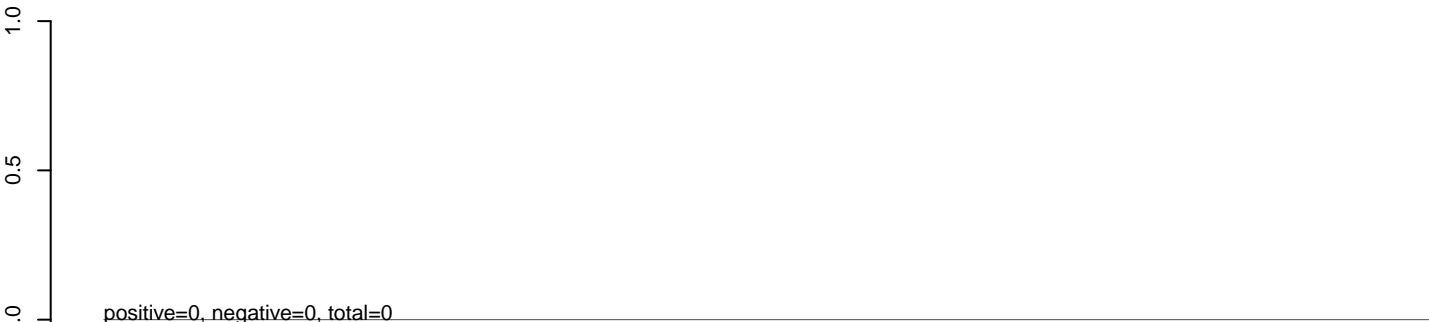


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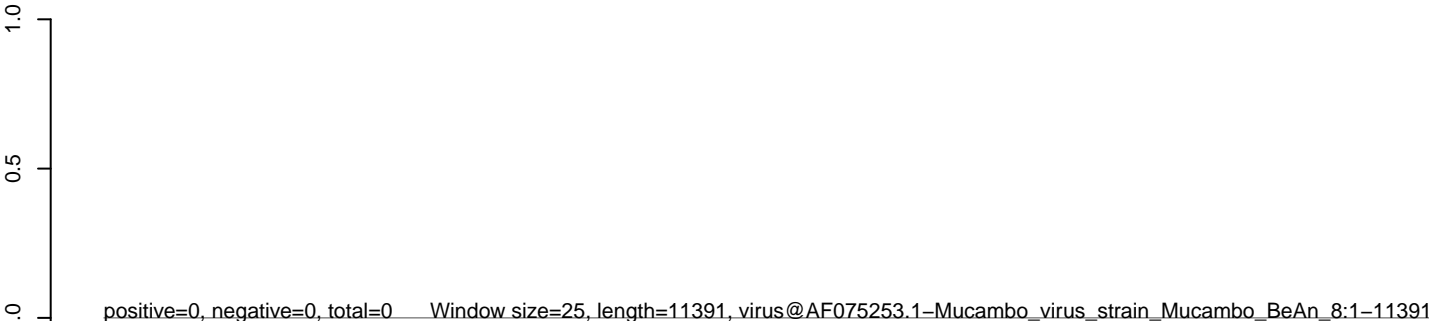
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