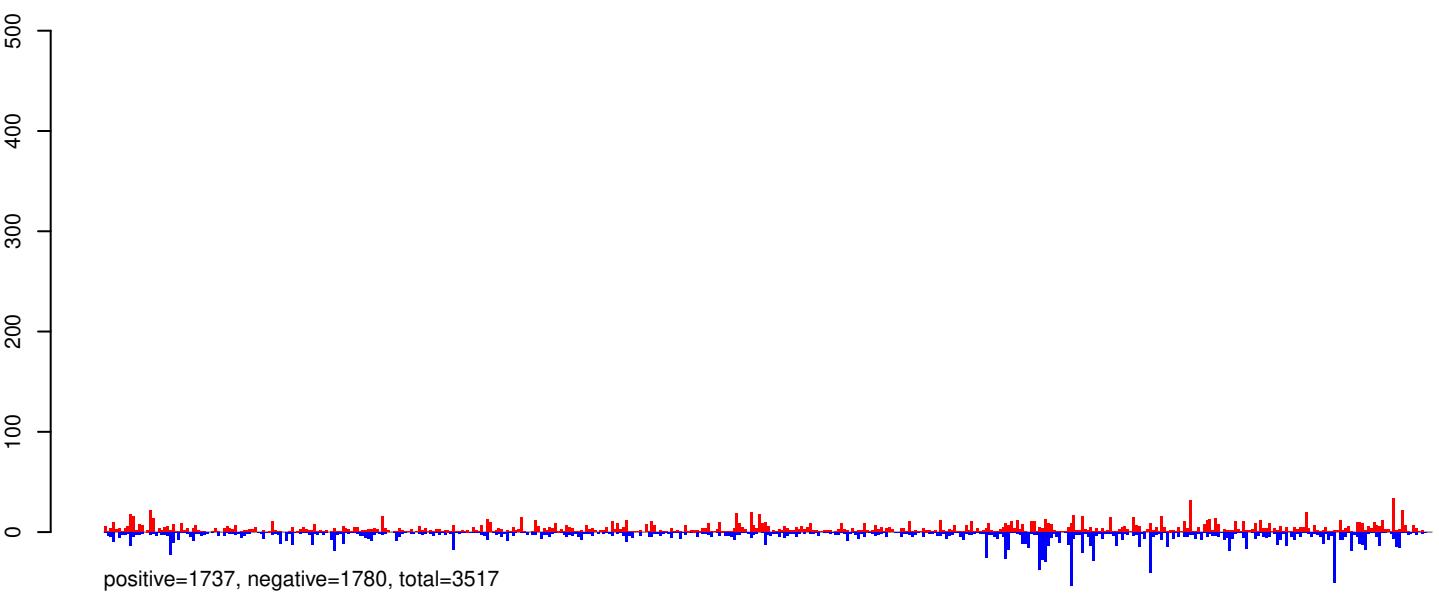
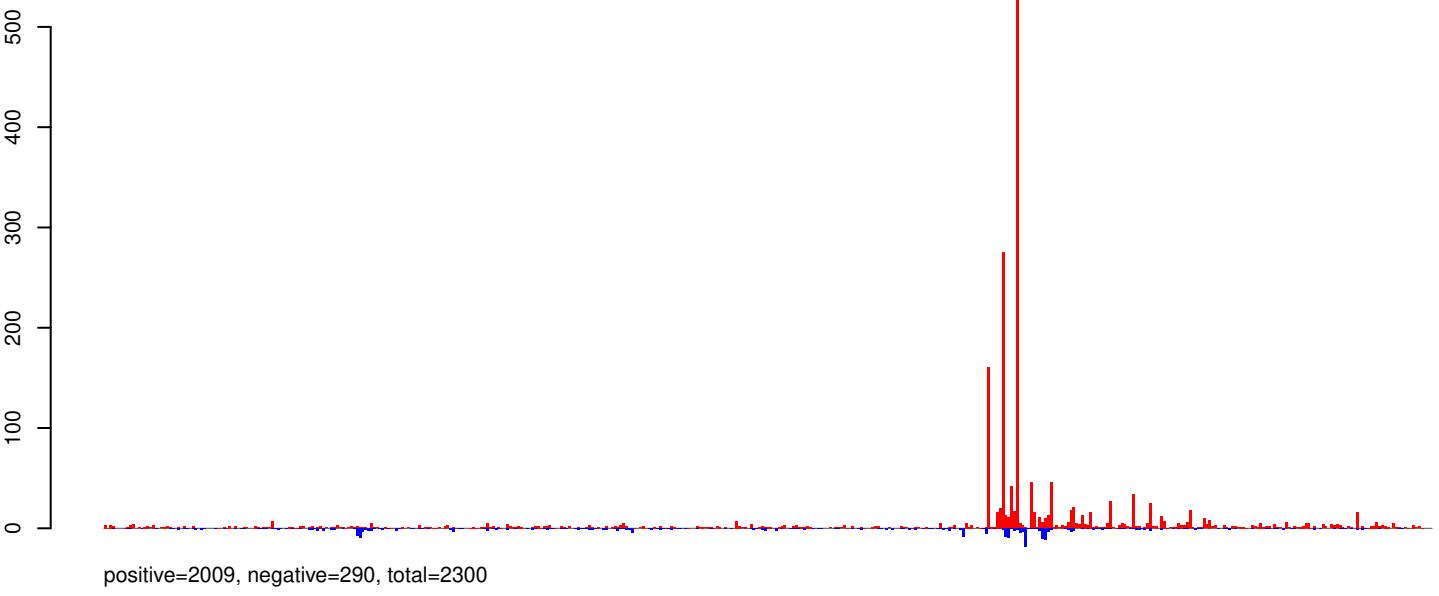


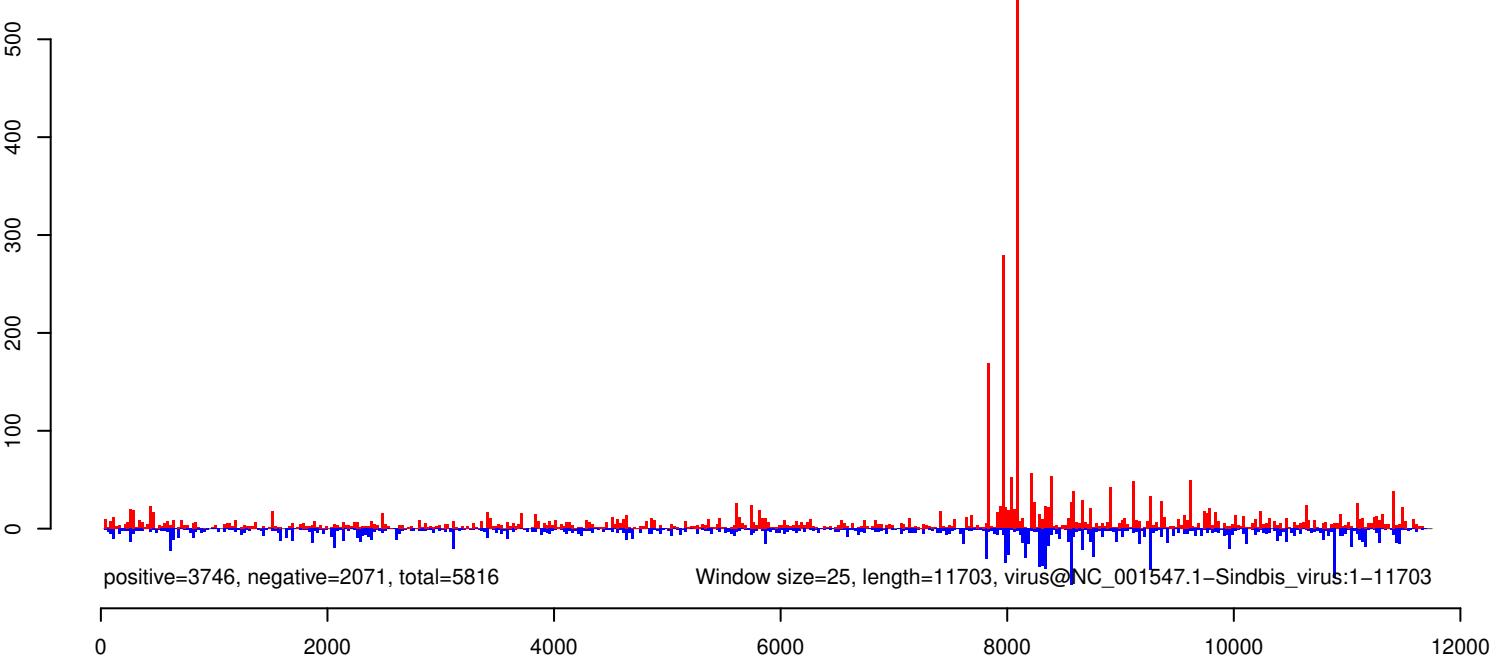
AeAeg_Aag2SINV_GFP.IP.18_23.rep



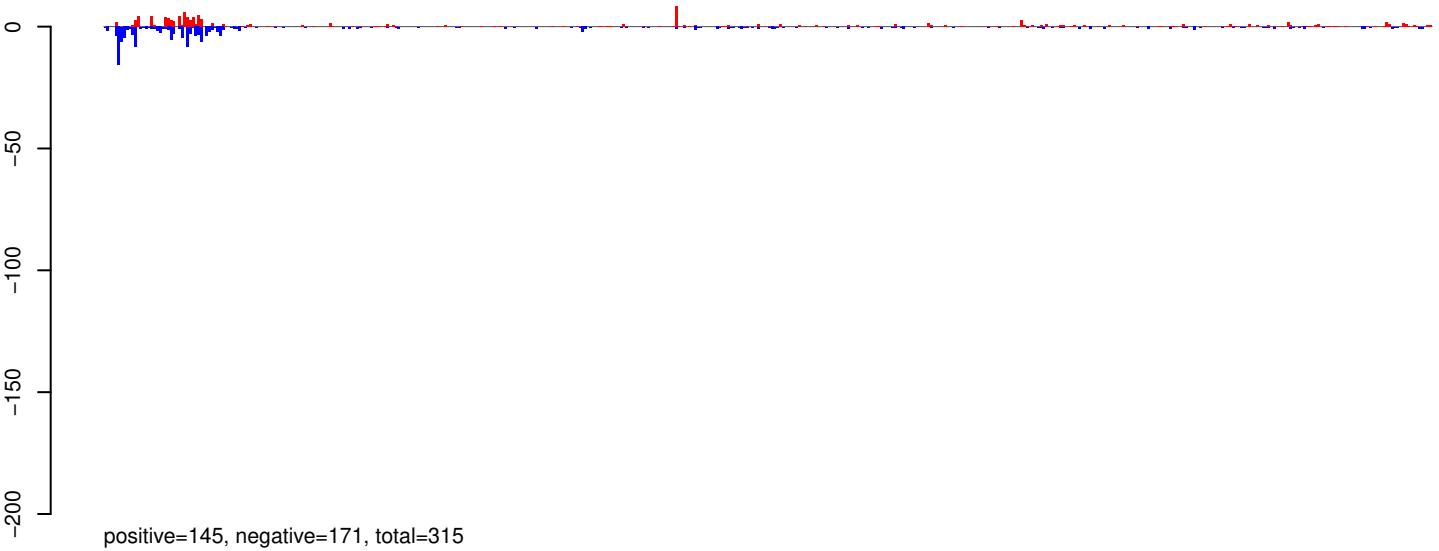
AeAeg_Aag2SINV_GFP.IP.24_35.rep



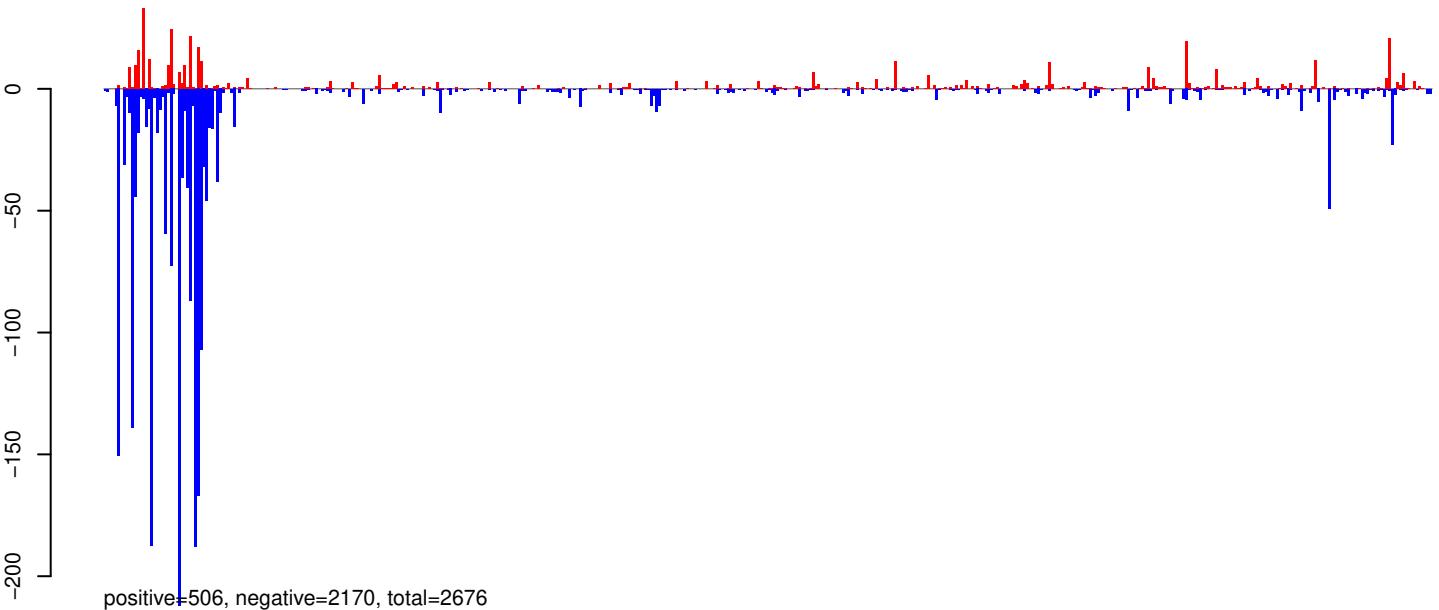
AeAeg_Aag2SINV_GFP.IP.rep



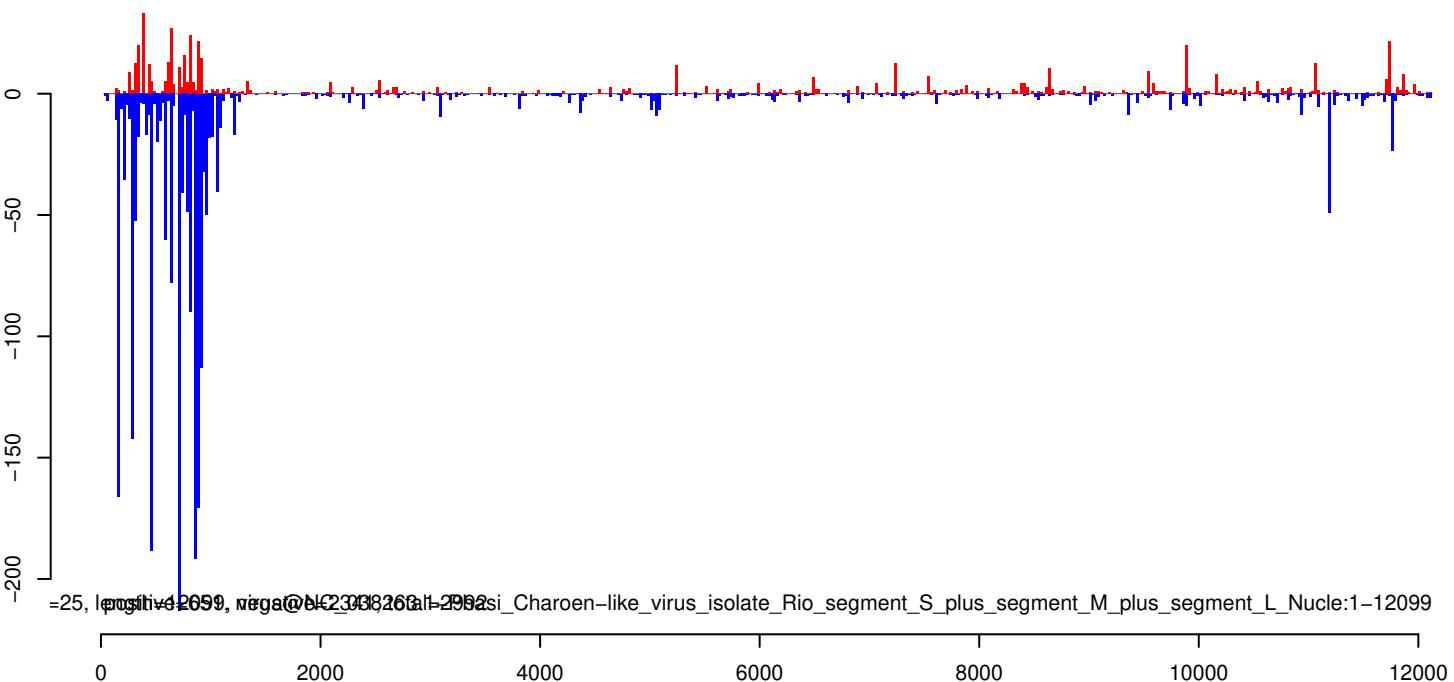
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=224, negative=260, total=484

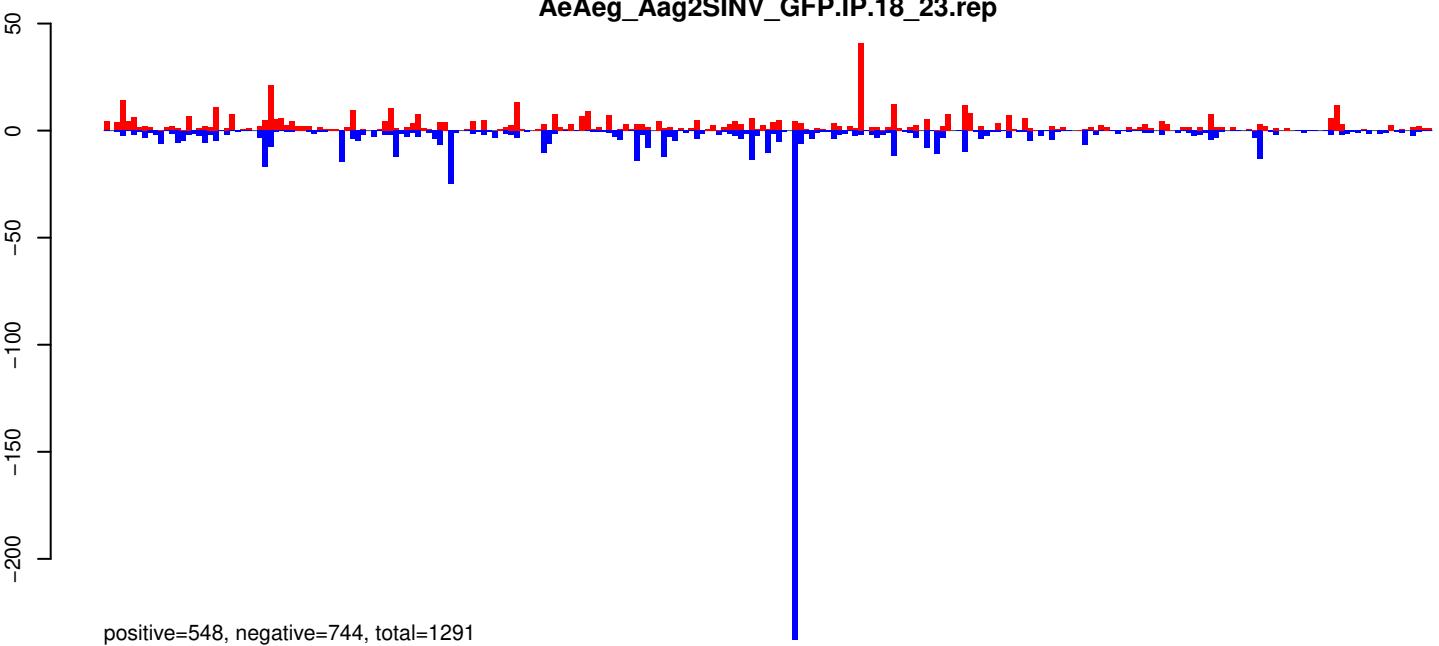
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=44, negative=113, total=157

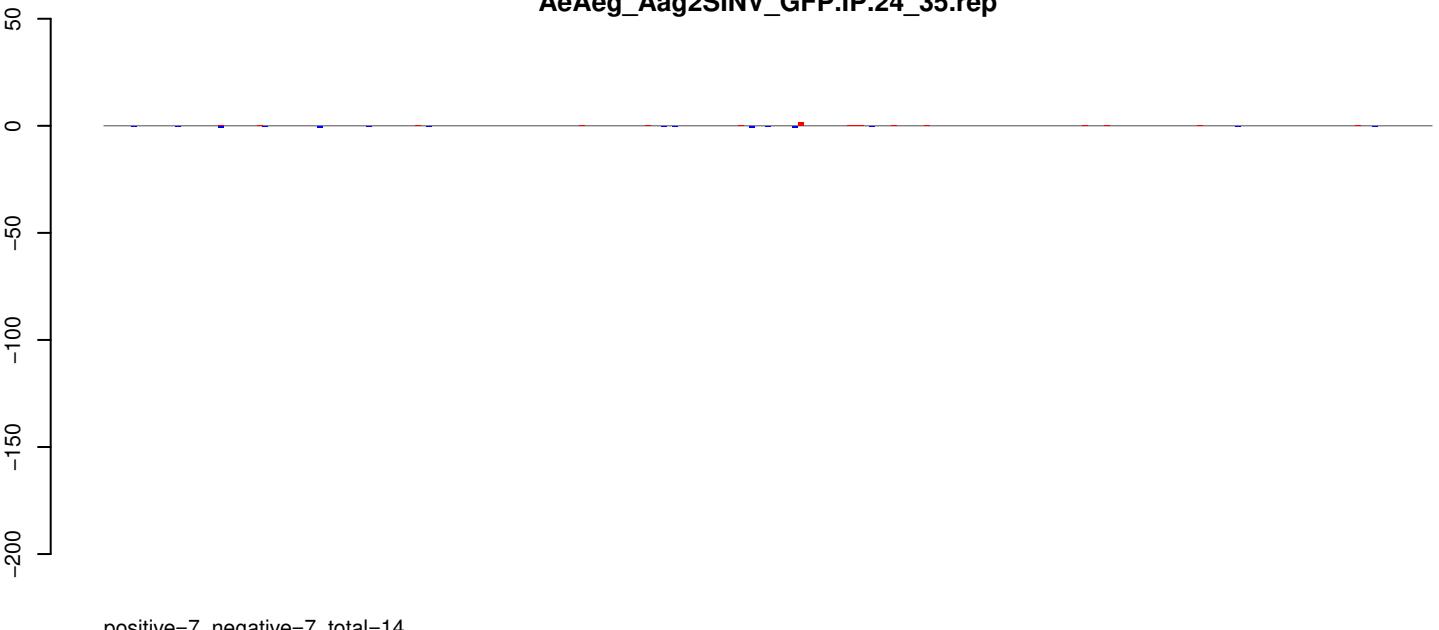
AeAeg_Aag2SINV_GFP.IP.rep

positive=268, negative=374, total=642, window size=25, length=10682, virus@NC_001564.2-Cell_fusing_agent_virus_strain_Galveston:1-10682

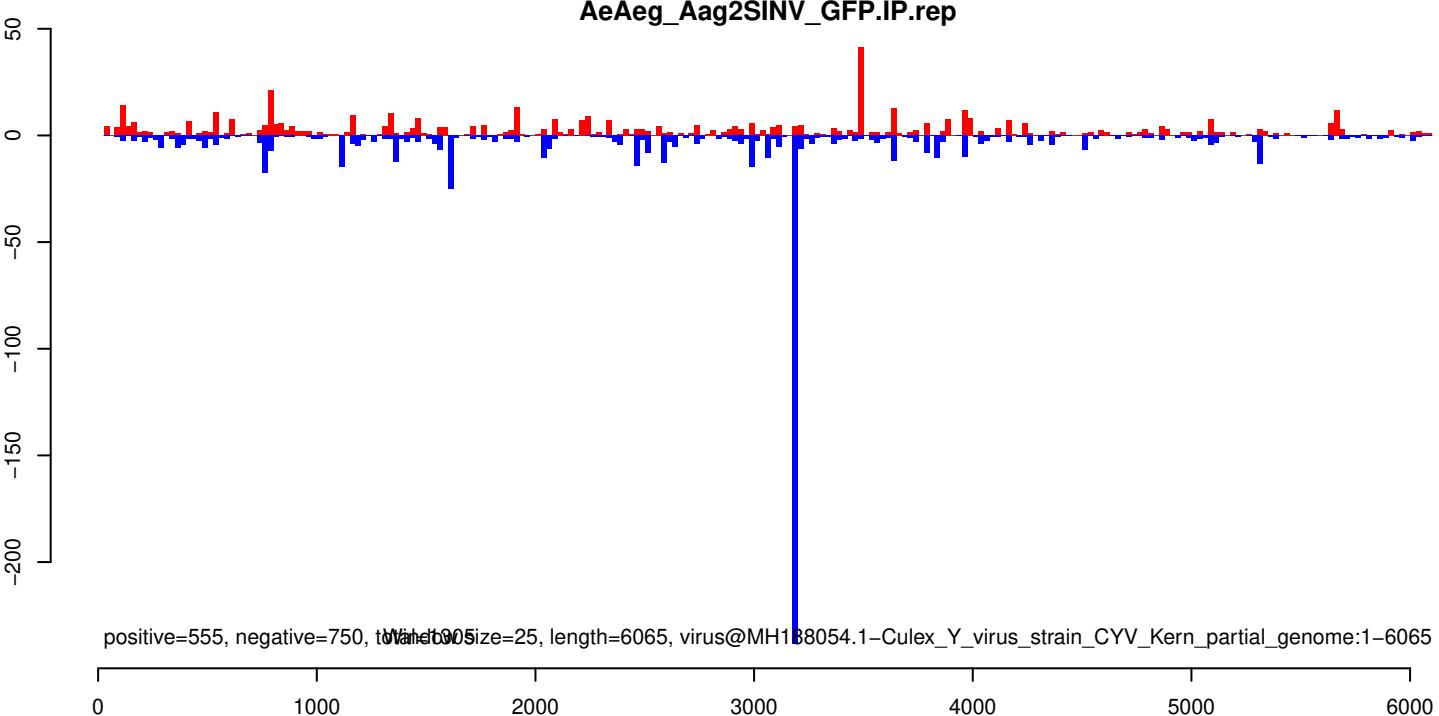
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



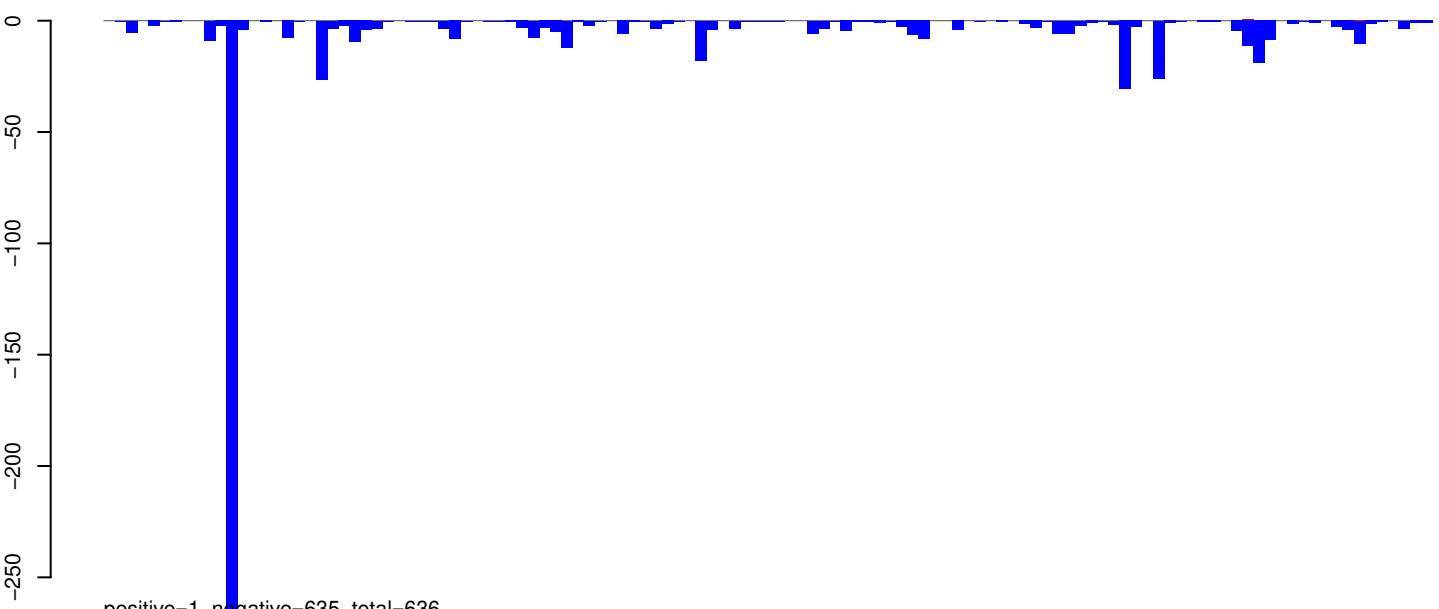
AeAeg_Aag2SINV_GFP.IP.rep



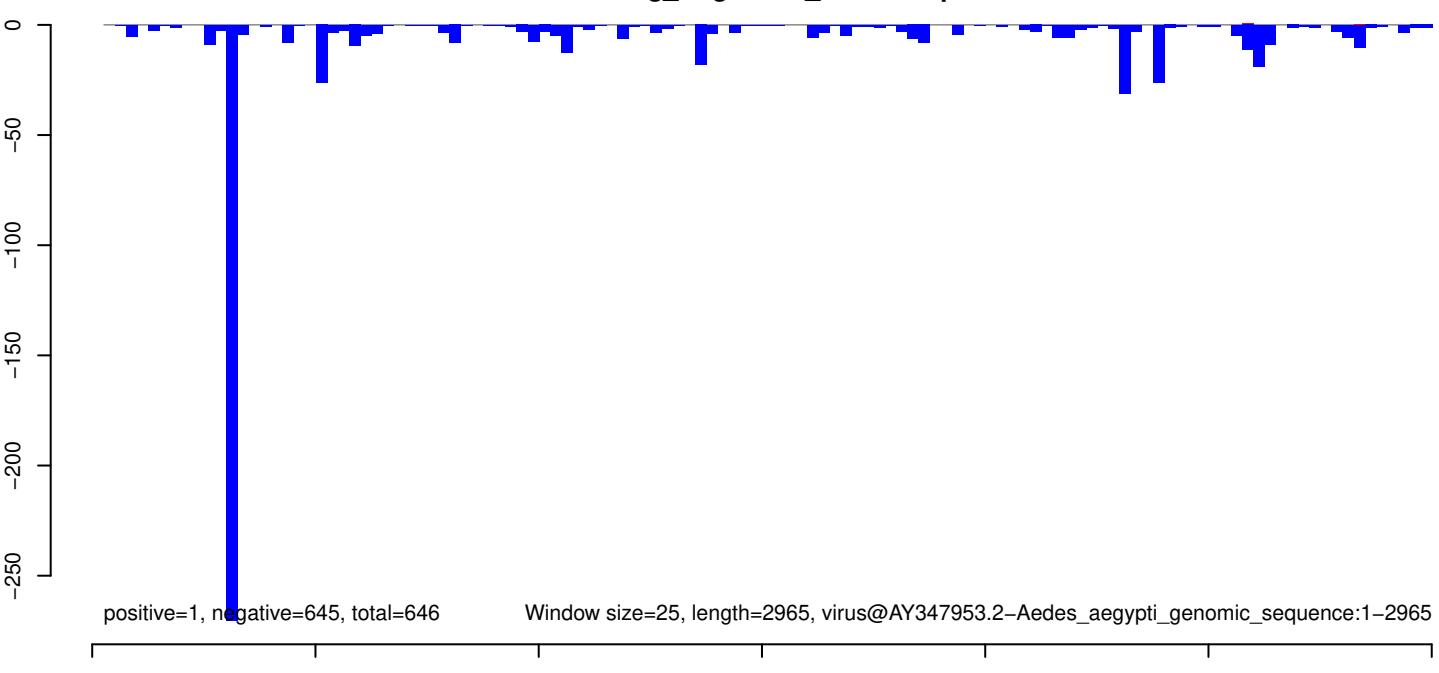
AeAeg_Aag2SINV_GFP.IP.18_23.rep



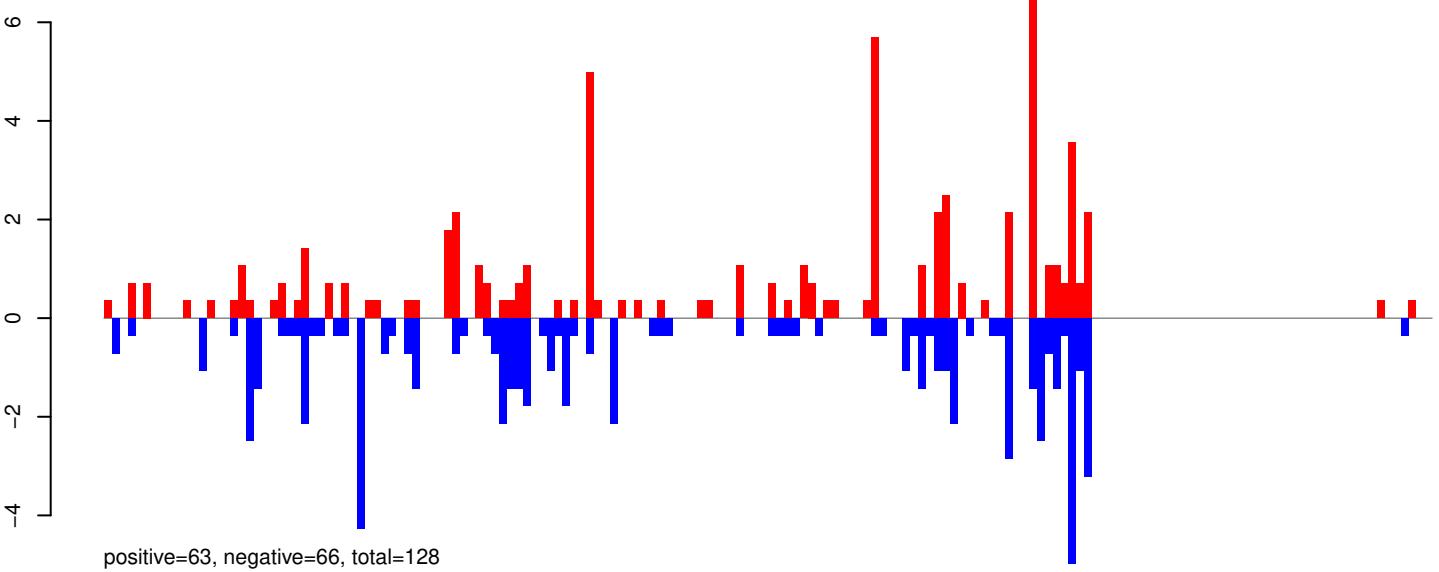
AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



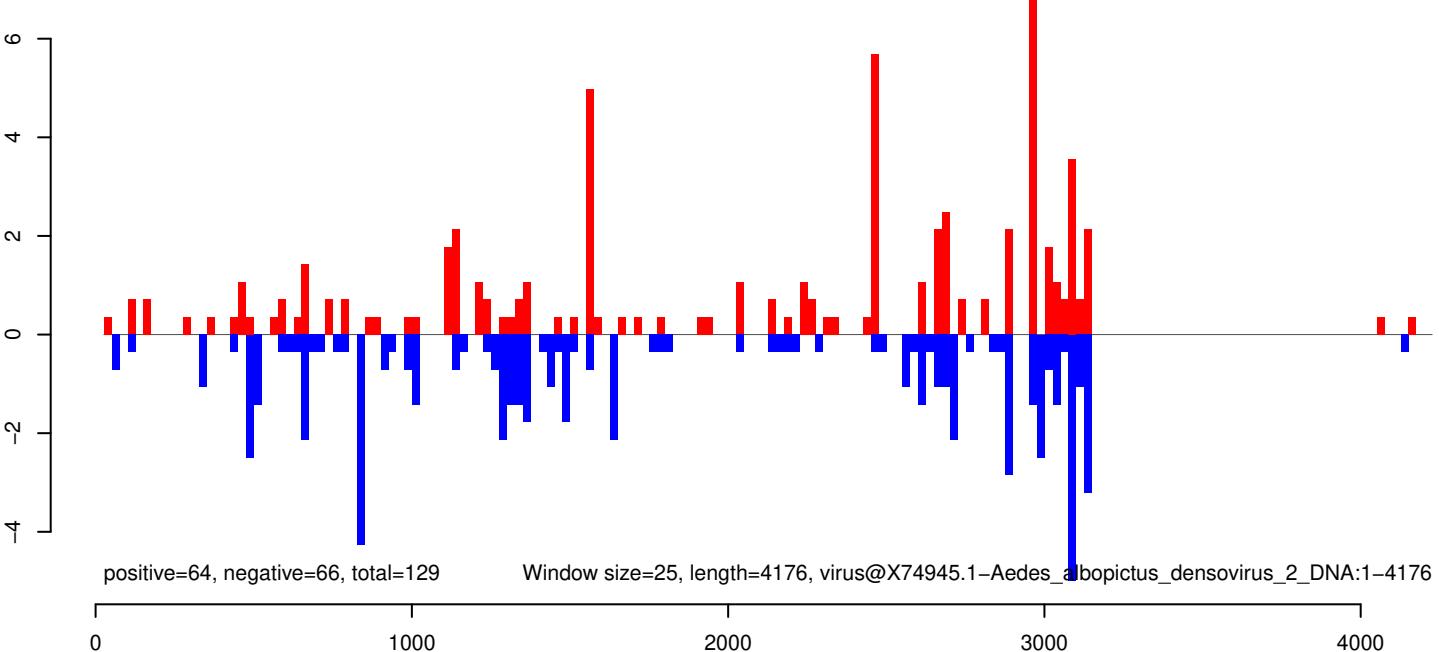
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep

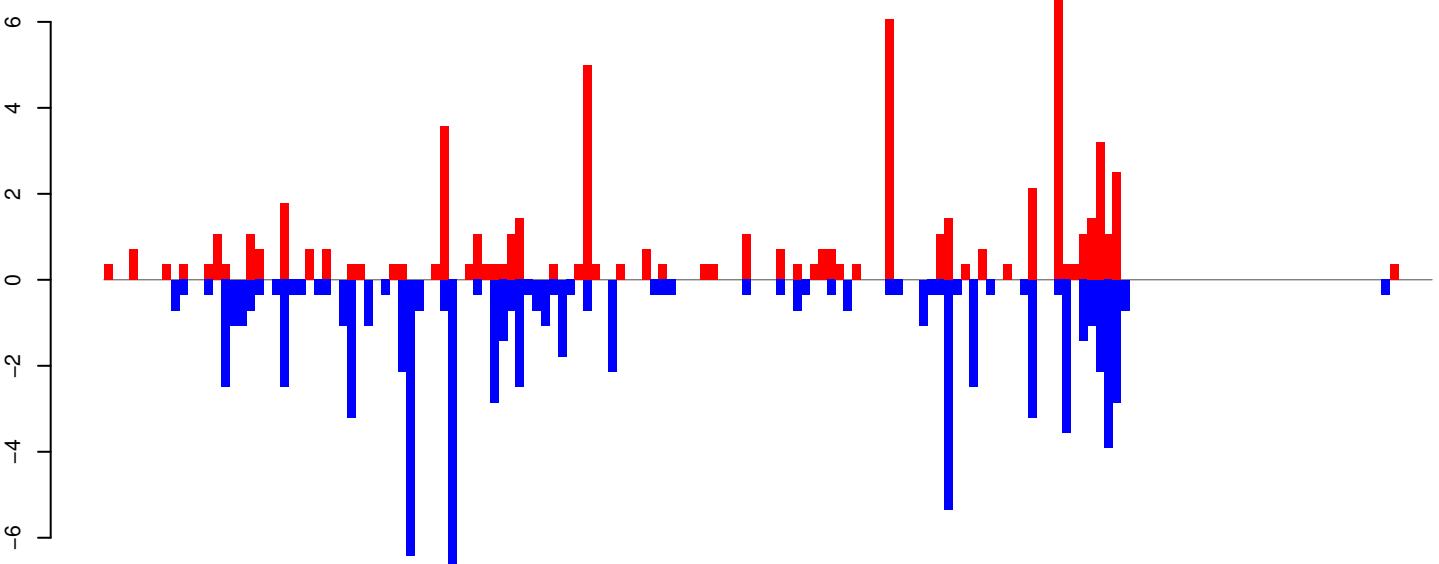


AeAeg_Aag2SINV_GFP.IP.rep



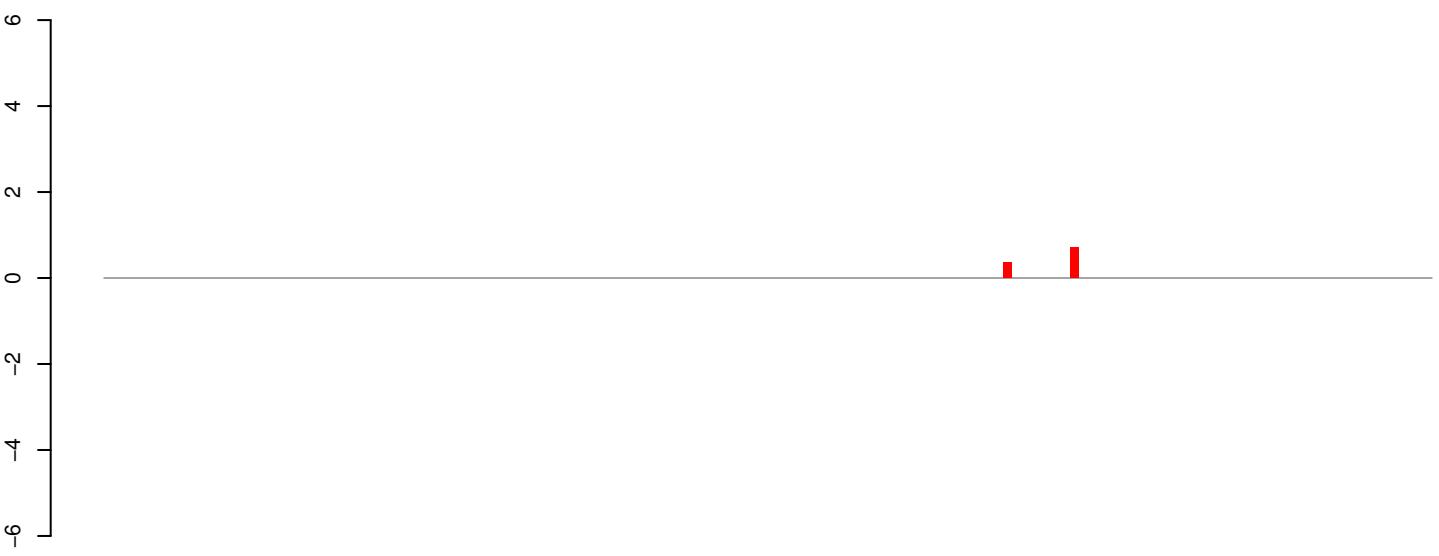
AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=61, negative=84, total=145



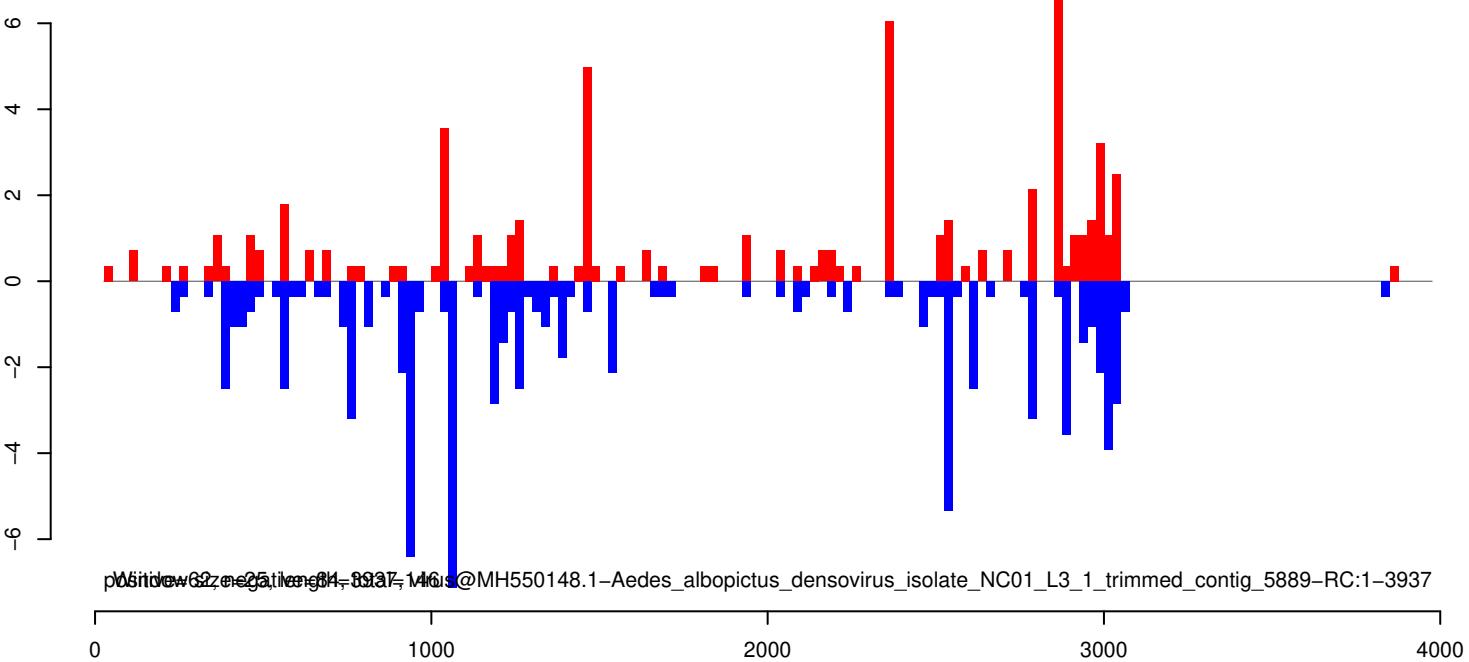
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=1, negative=0, total=1

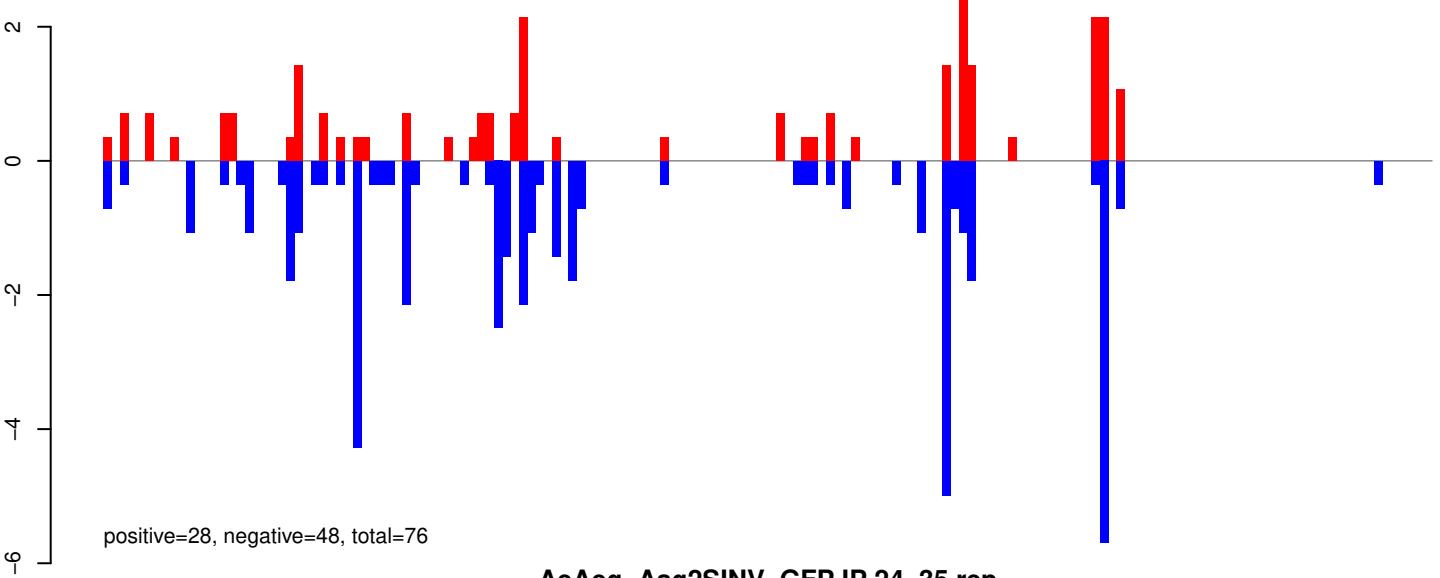


AeAeg_Aag2SINV_GFP.IP.rep

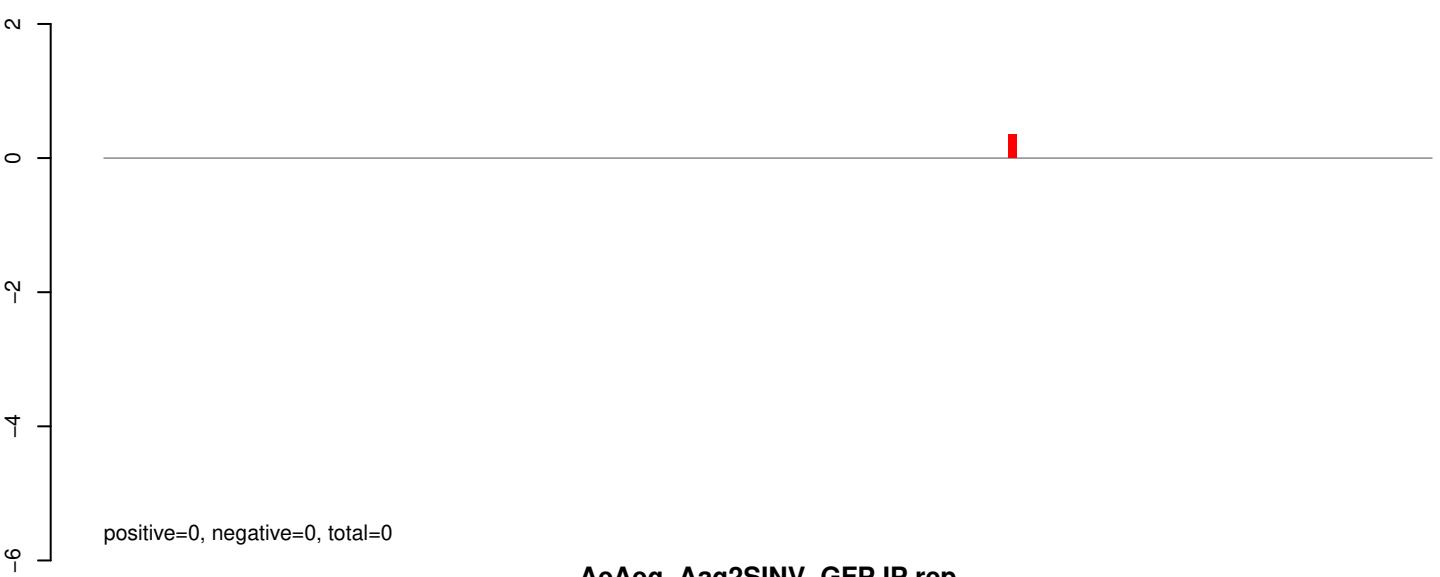
positive=62, negative=84, total=146 @MH550148.1-Aedes_albopictus_densovirus_isolate_NC01_L3_1_trimmed_contig_5889-RC:1-3937



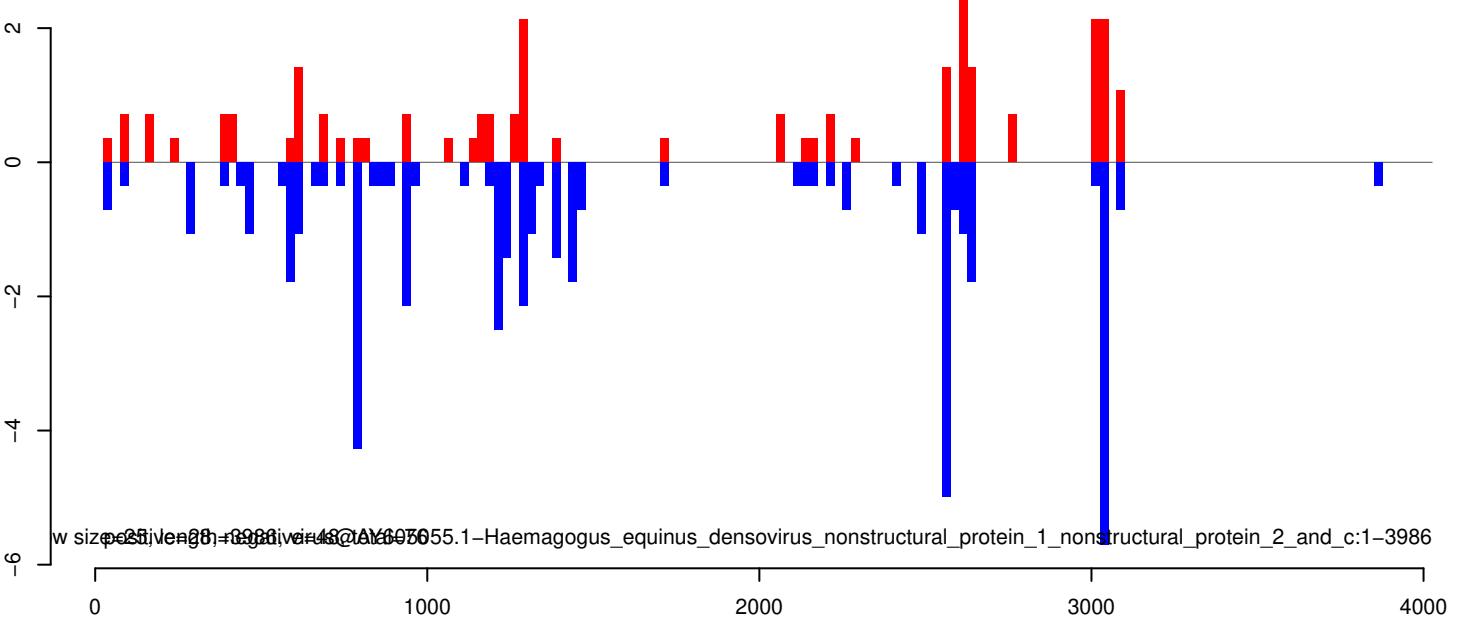
AeAeg_Aag2SINV_GFP.IP.18_23.rep



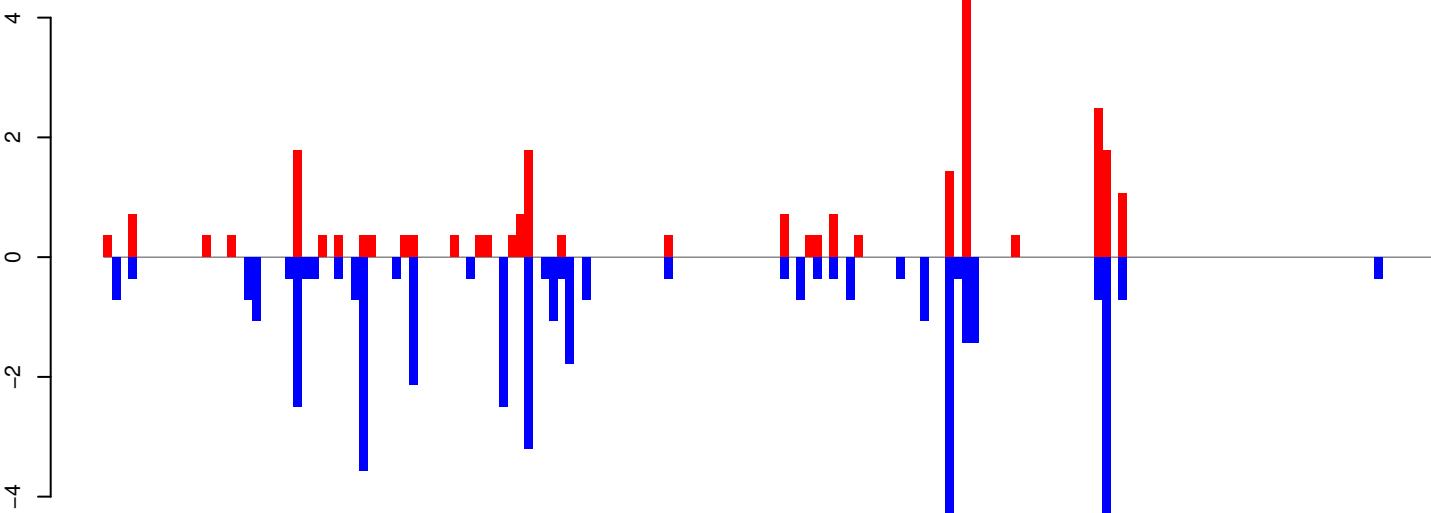
AeAeg_Aag2SINV_GFP.IP.24_35.rep



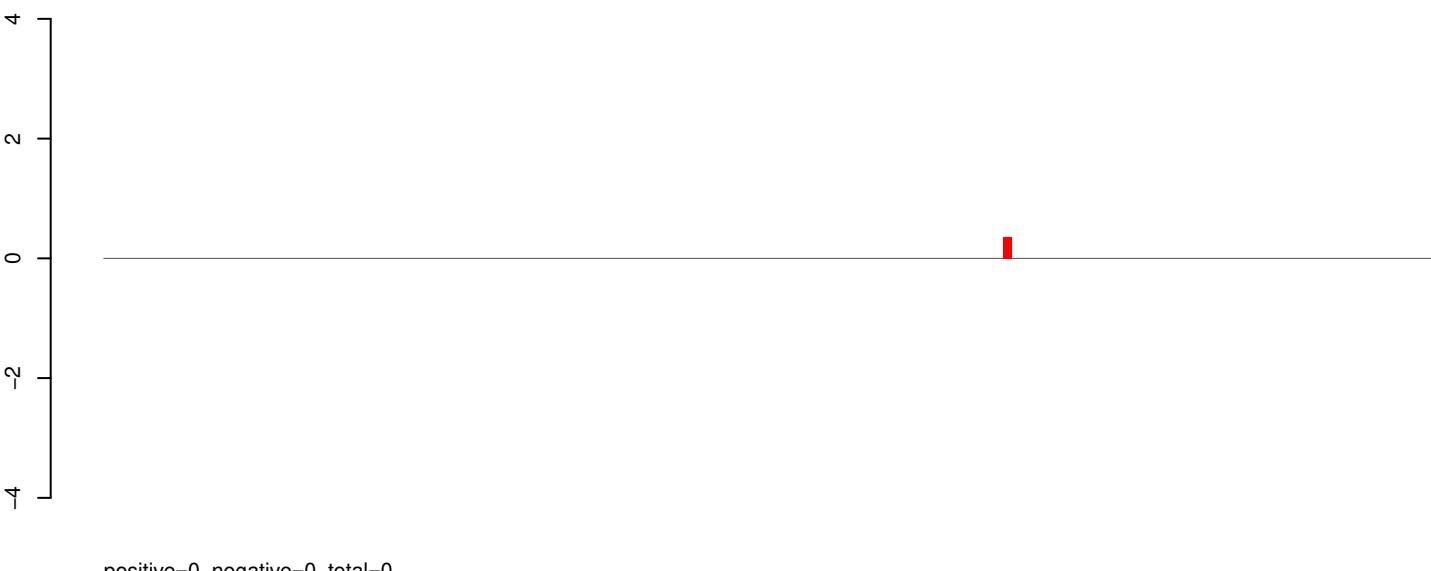
AeAeg_Aag2SINV_GFP.IP.rep



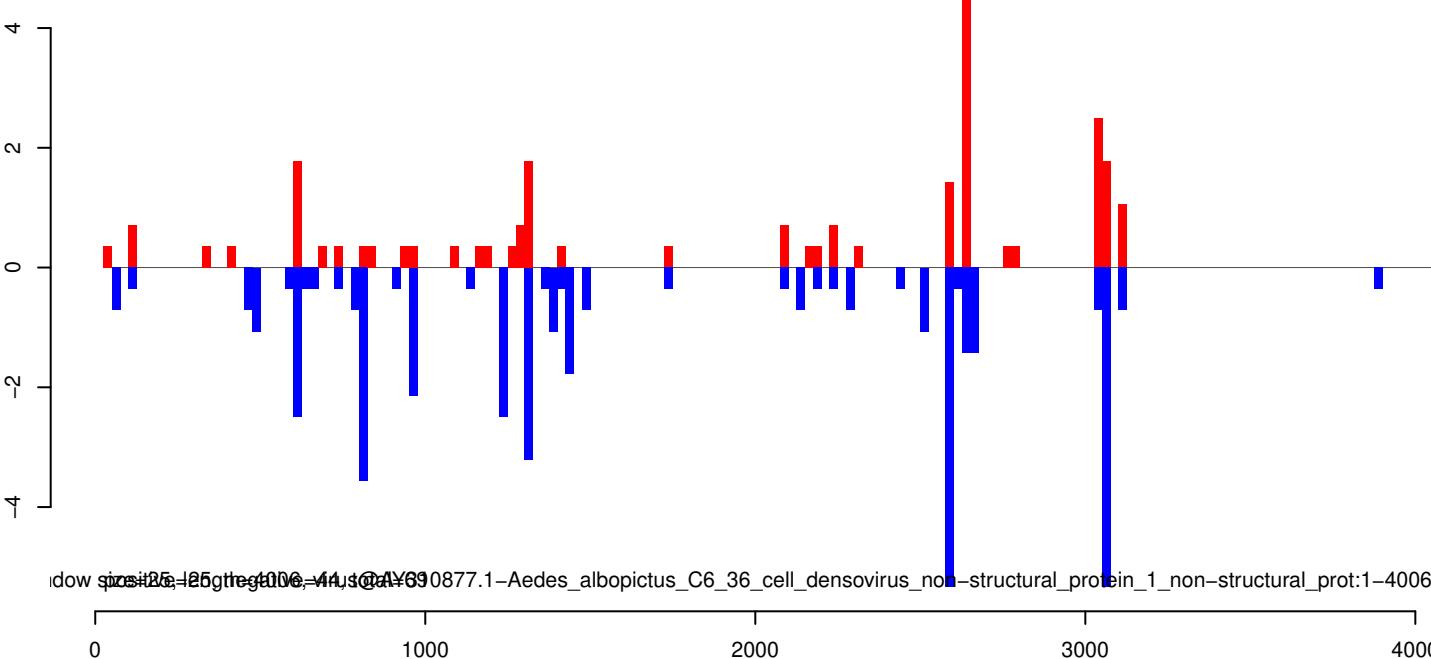
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=4, negative=14, total=18

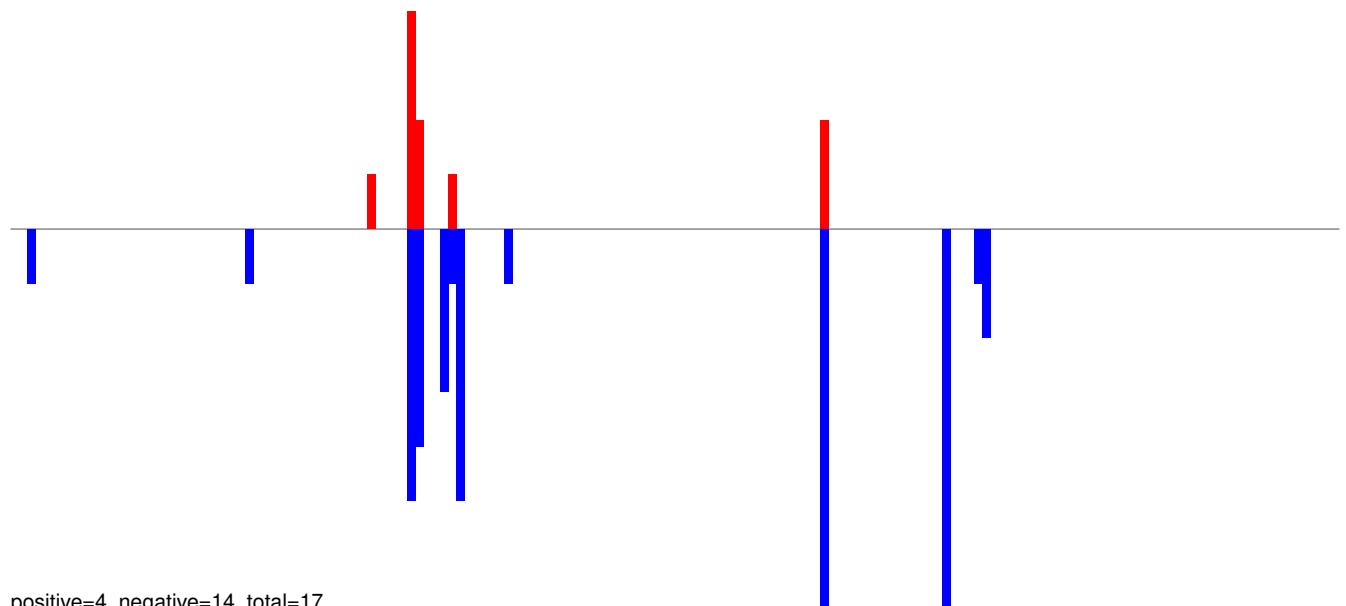
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

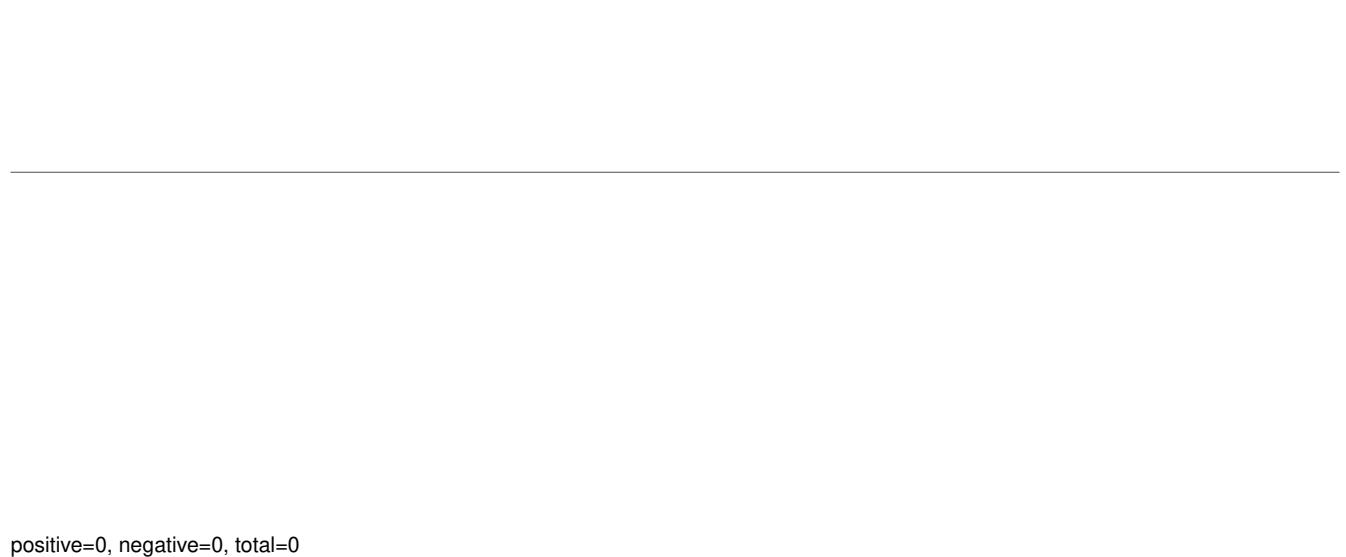
AeAeg_Aag2SINV_GFP.IP.rep

positive=4, negative=14, total=19 Window size=25, length=4073, virus@KX603698.1–Aedes_albopictus_densovirus_5_strain_GZ05:1–4073

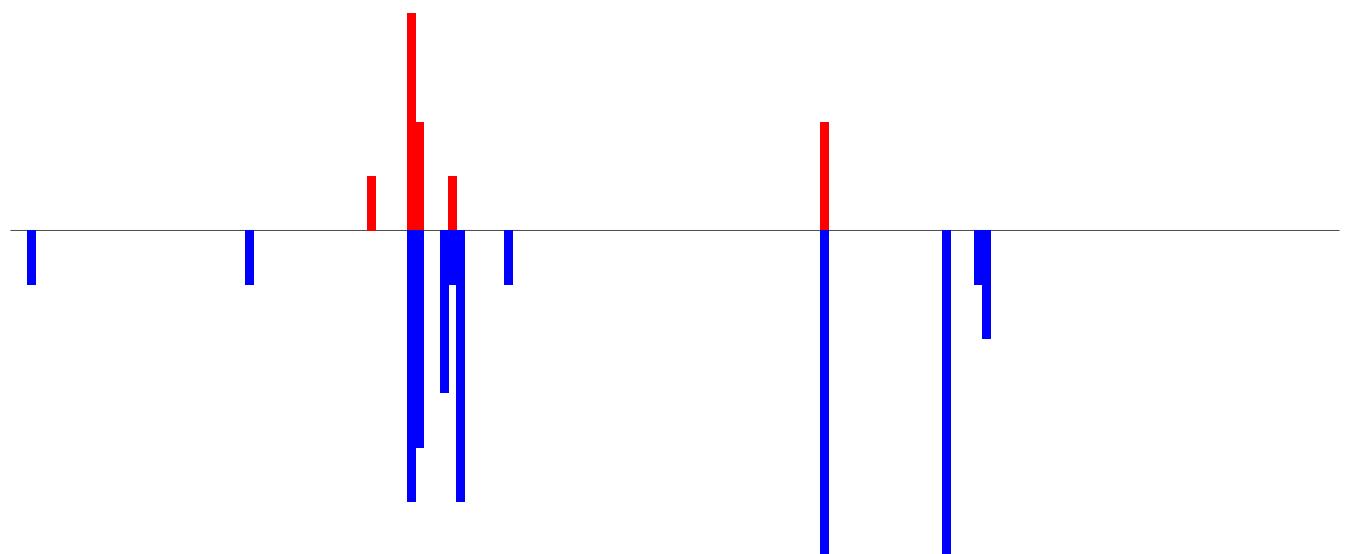
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



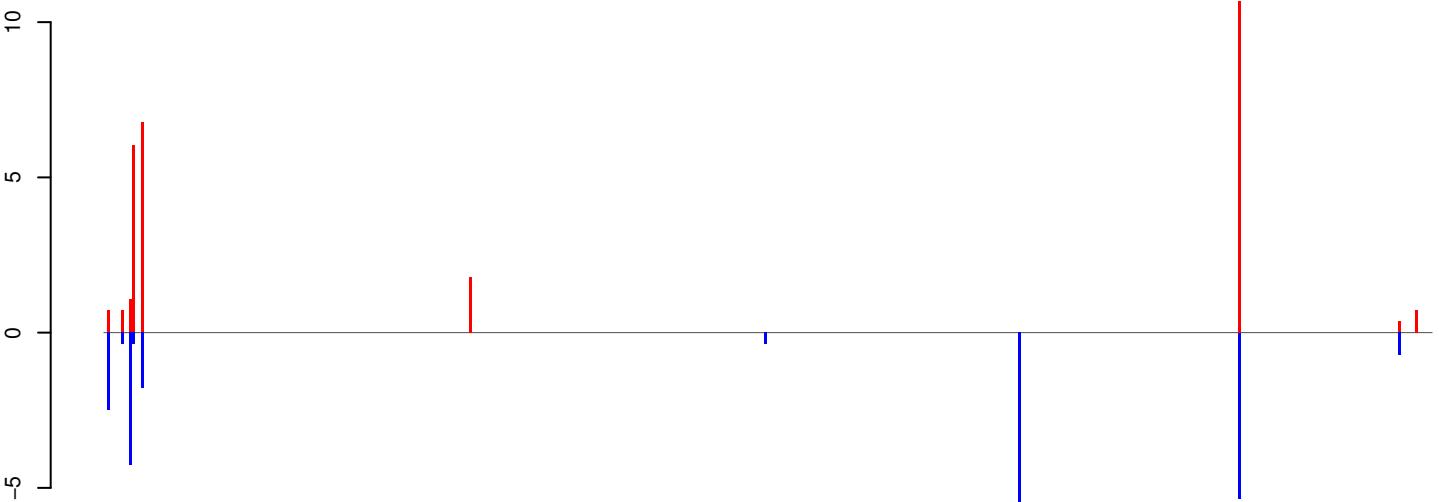
AeAeg_Aag2SINV_GFP.IP.rep



cow size=25 by length=1000 width=141@AY095351.2-Aedes_albopictus_C6_36_cell_densovirus_non-structural_protein_1_NS-1_non-structural:1-4094

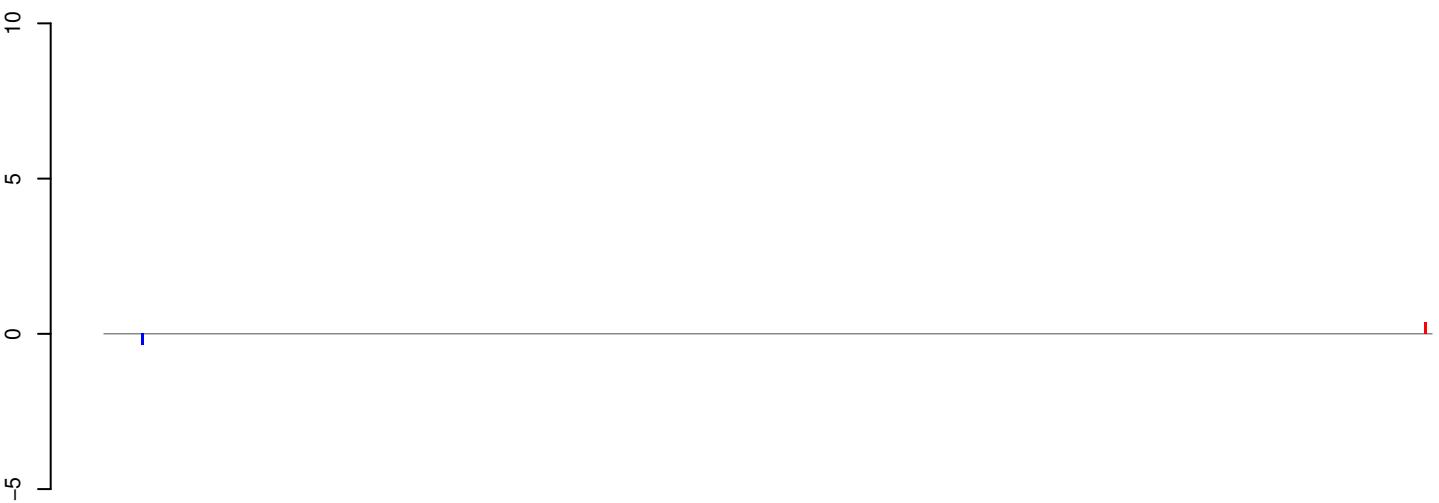
0 1000 2000 3000 4000

AeAeg_Aag2SINV_GFP.IP.18_23.rep



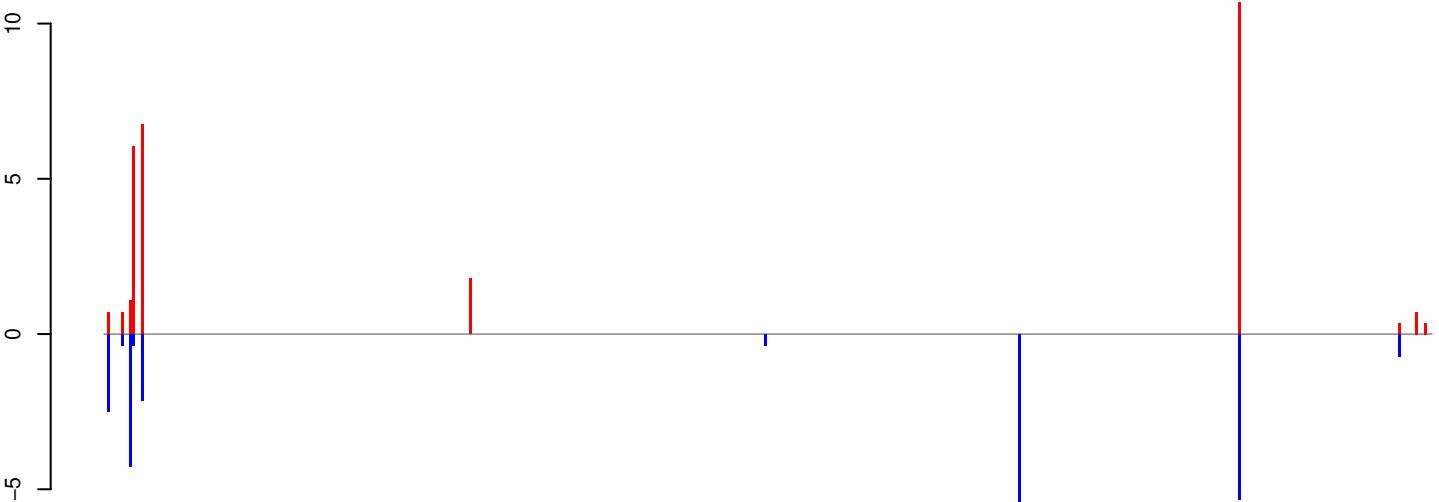
positive=29, negative=24, total=53

AeAeg_Aag2SINV_GFP.IP.24_35.rep



positive=0, negative=0, total=1

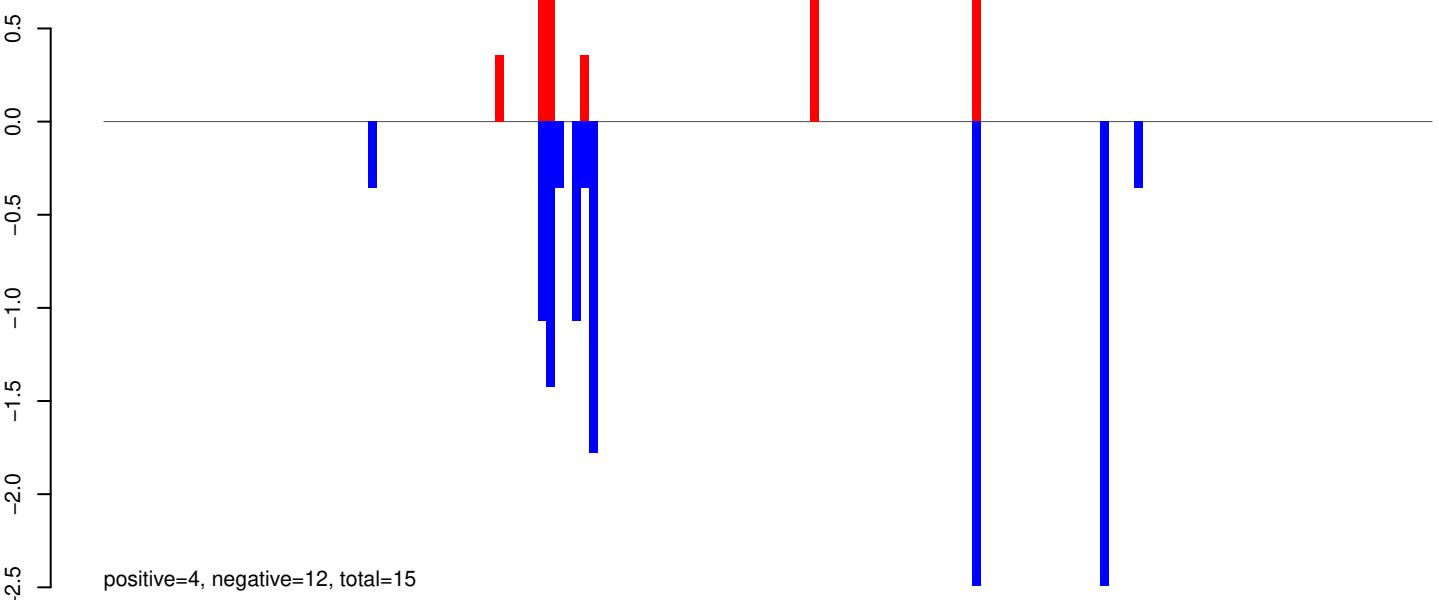
AeAeg_Aag2SINV_GFP.IP.rep



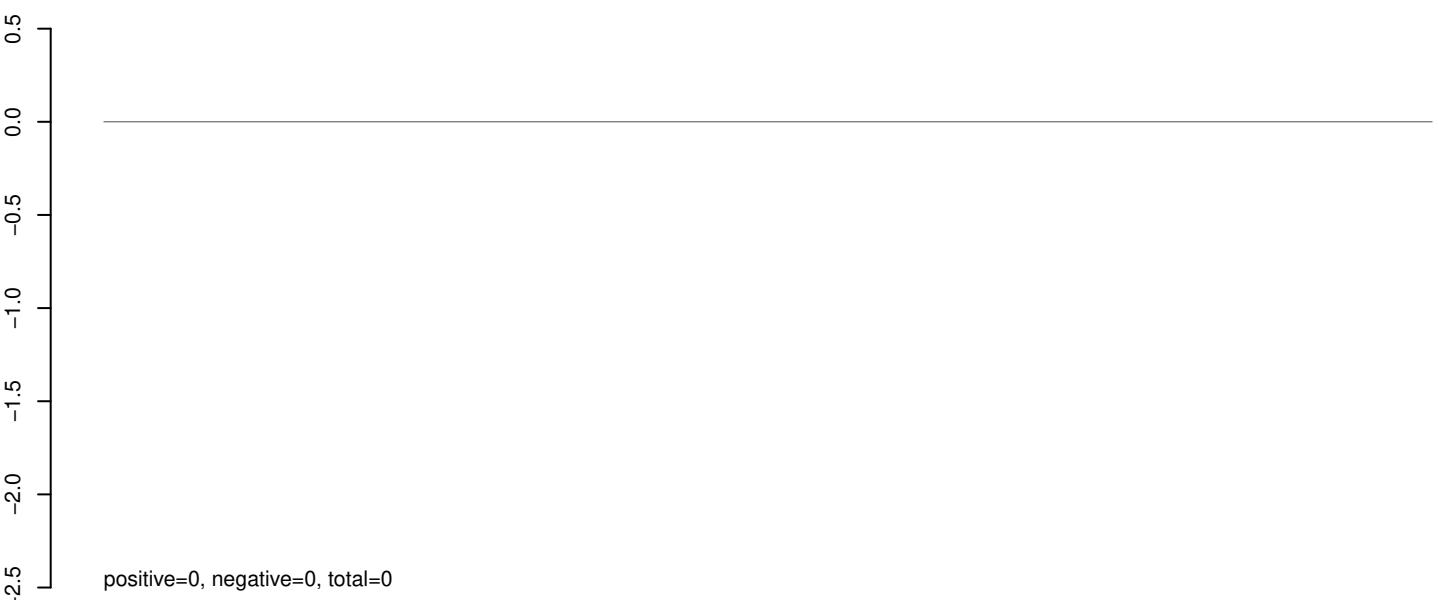
positive=29, negative=25, total=54

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

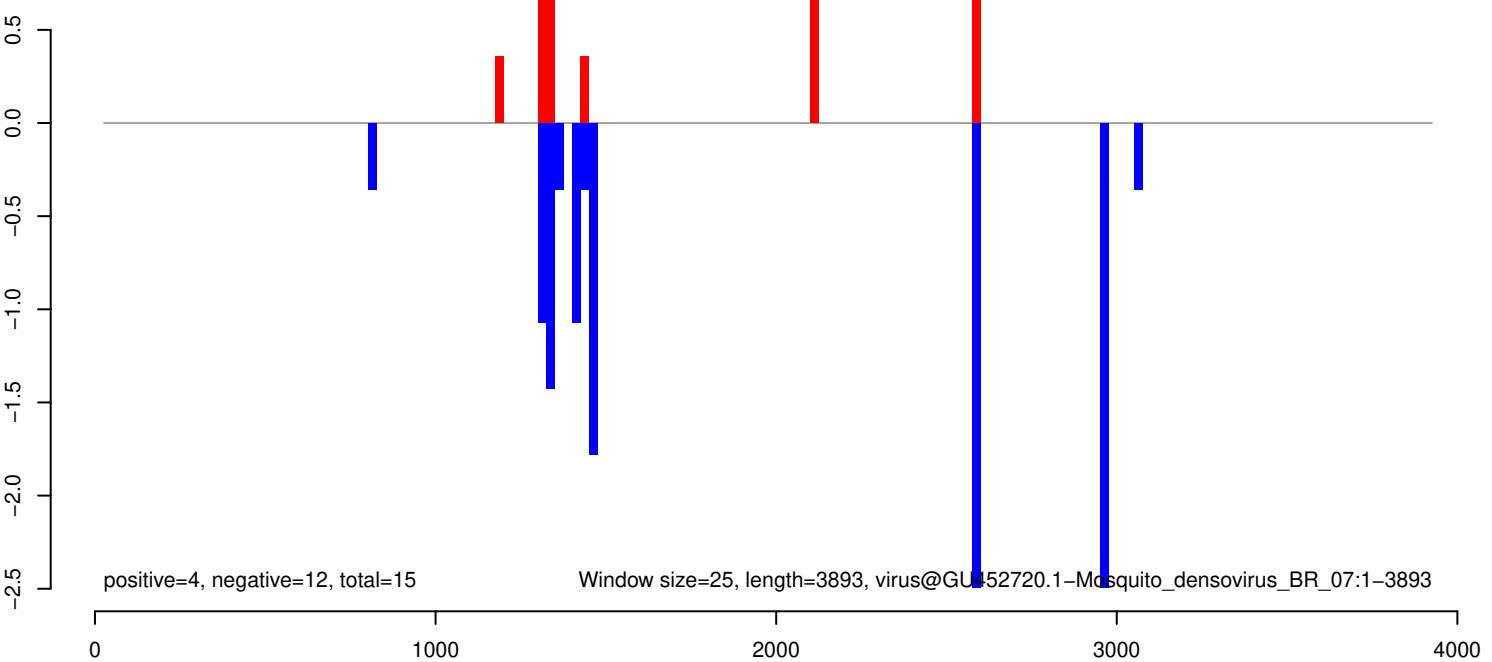
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=6, negative=12, total=18

AeAeg_Aag2SINV_GFP.IP.24_35.rep

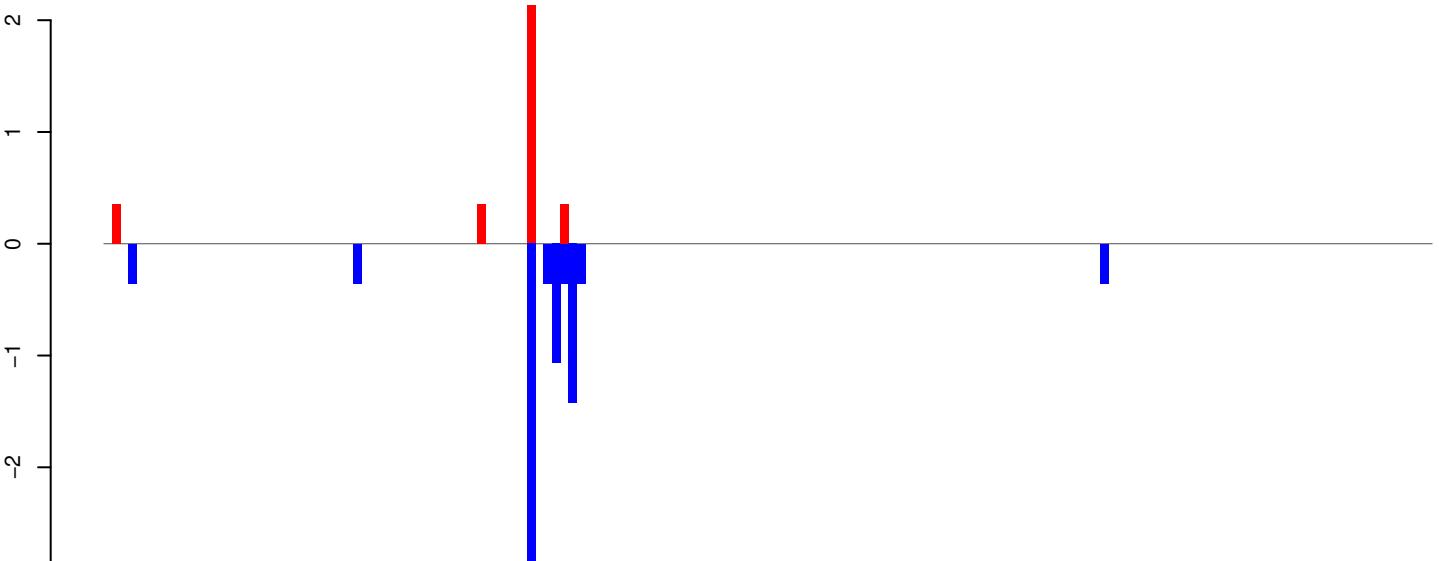
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=6, negative=12, total=18 Window size=25, length=4139, virus@NC_011317.1-Anopheles_gambiae_densonucleosis_virus:1-4139

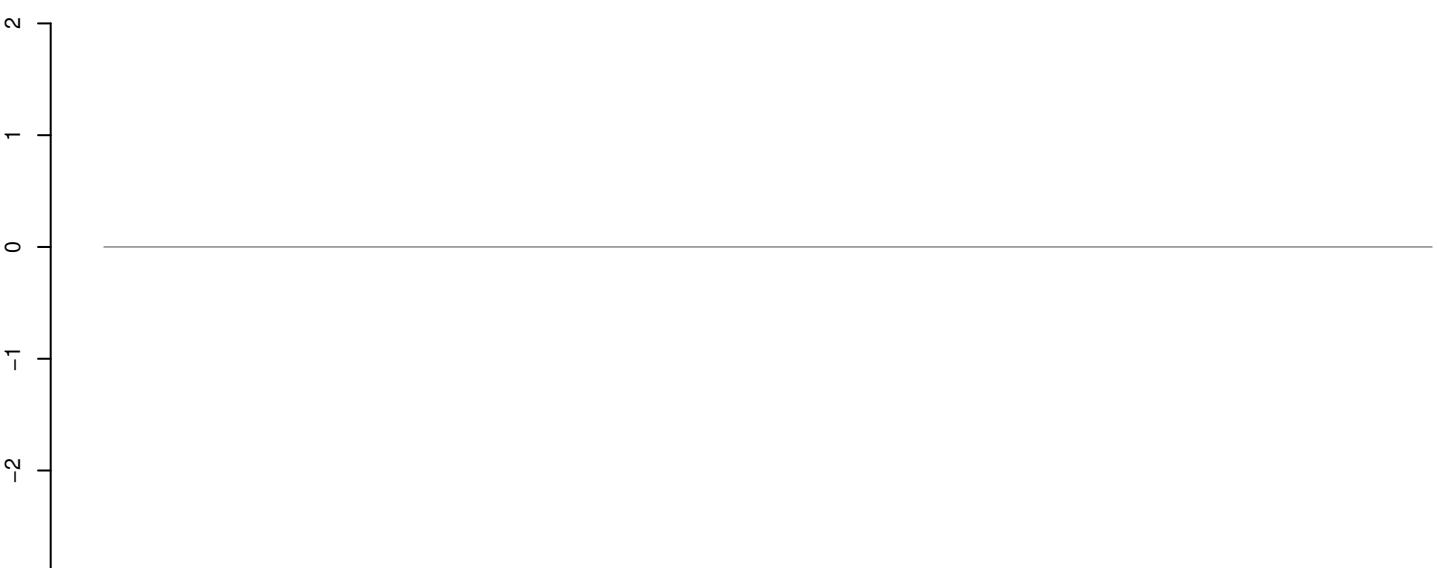
AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=3, negative=8, total=11



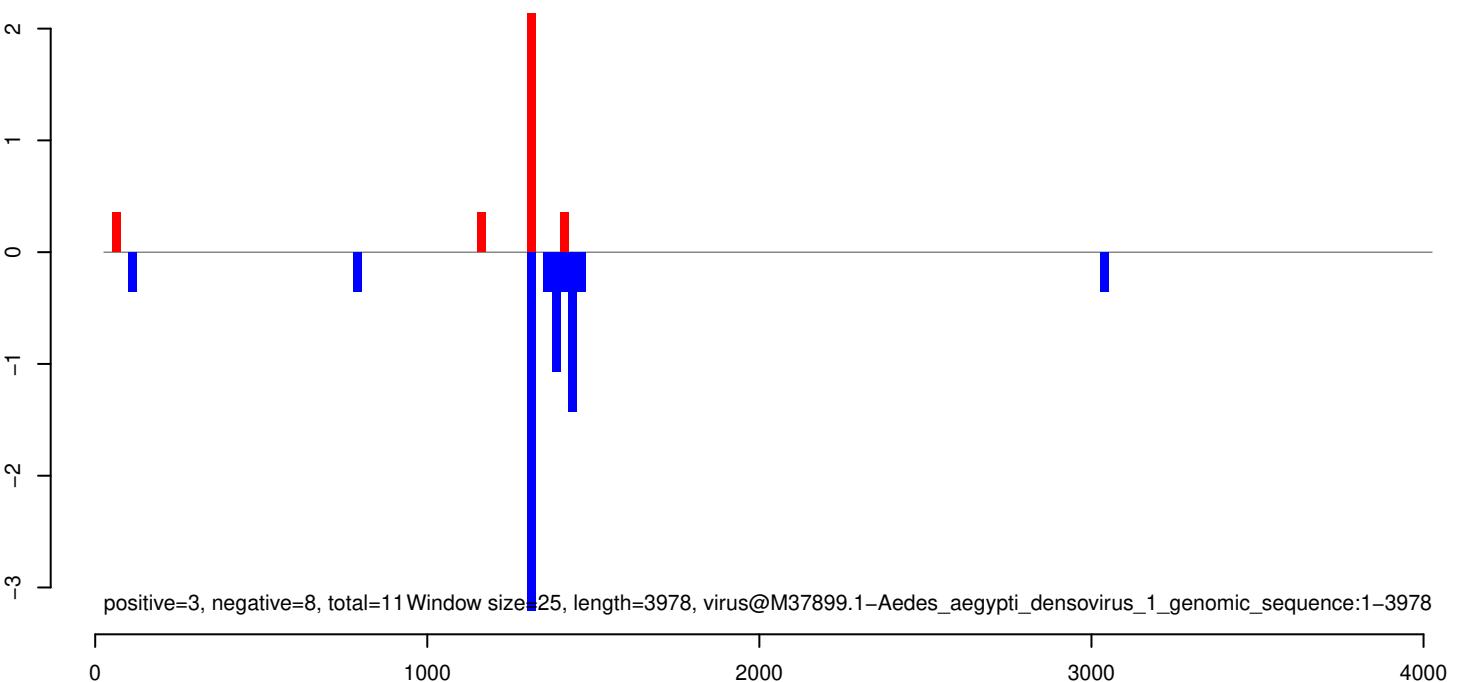
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

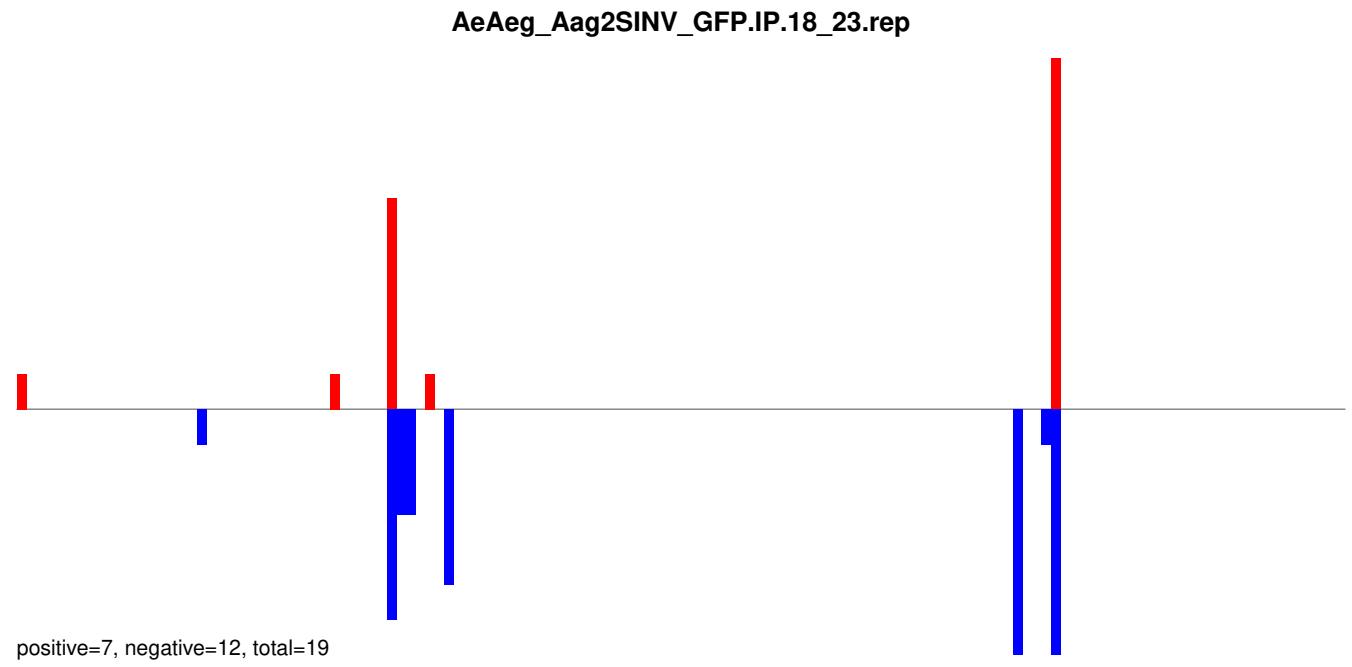


AeAeg_Aag2SINV_GFP.IP.rep

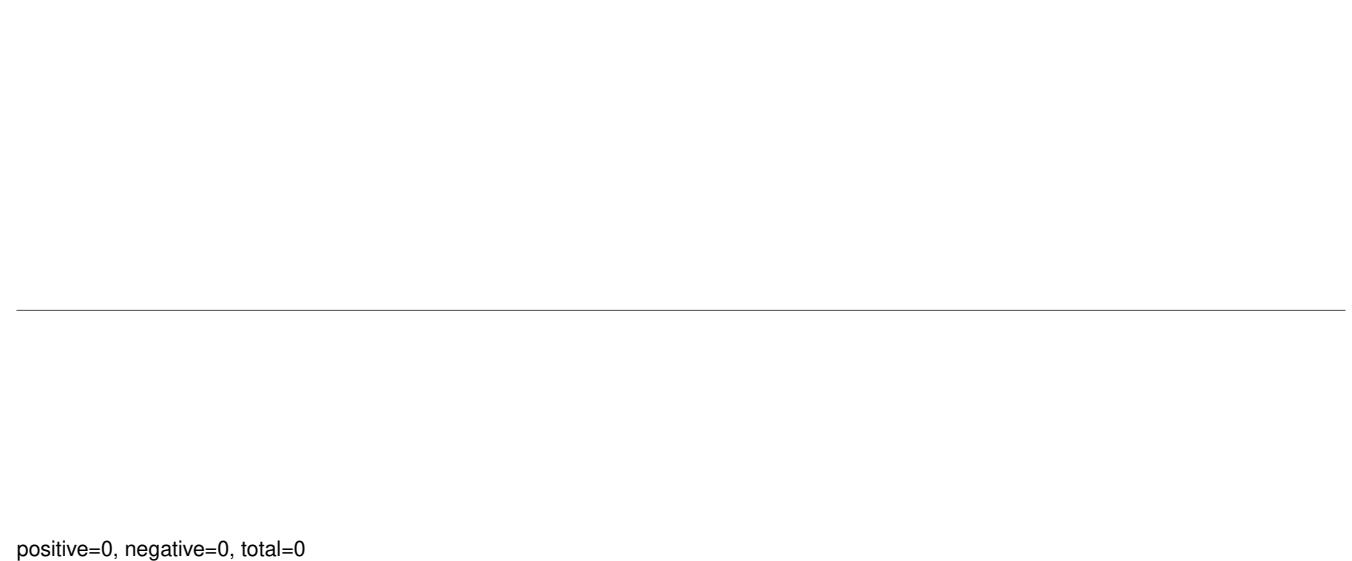
positive=3, negative=8, total=11 Window size=25, length=3978, virus@M37899.1-Aedes_aegypti_densovirus_1_genomic_sequence:1-3978



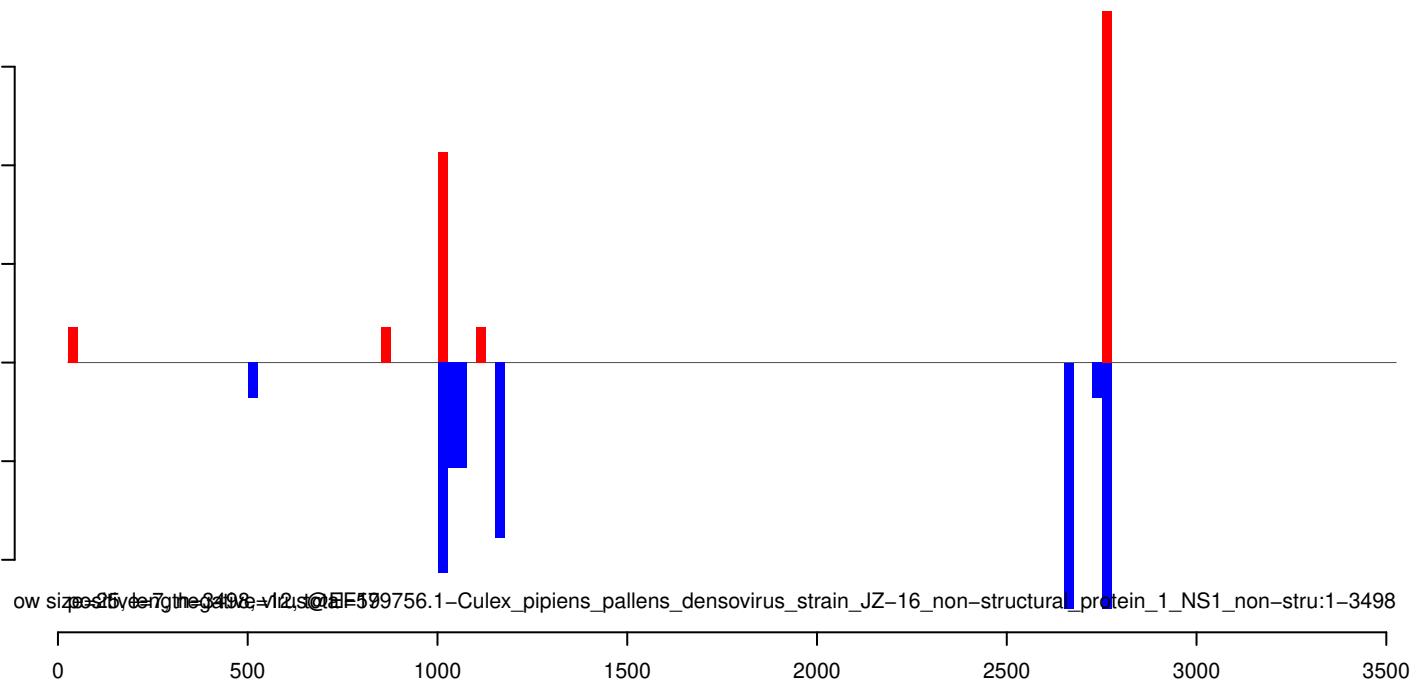
AeAeg_Aag2SINV_GFP.IP.18_23.rep



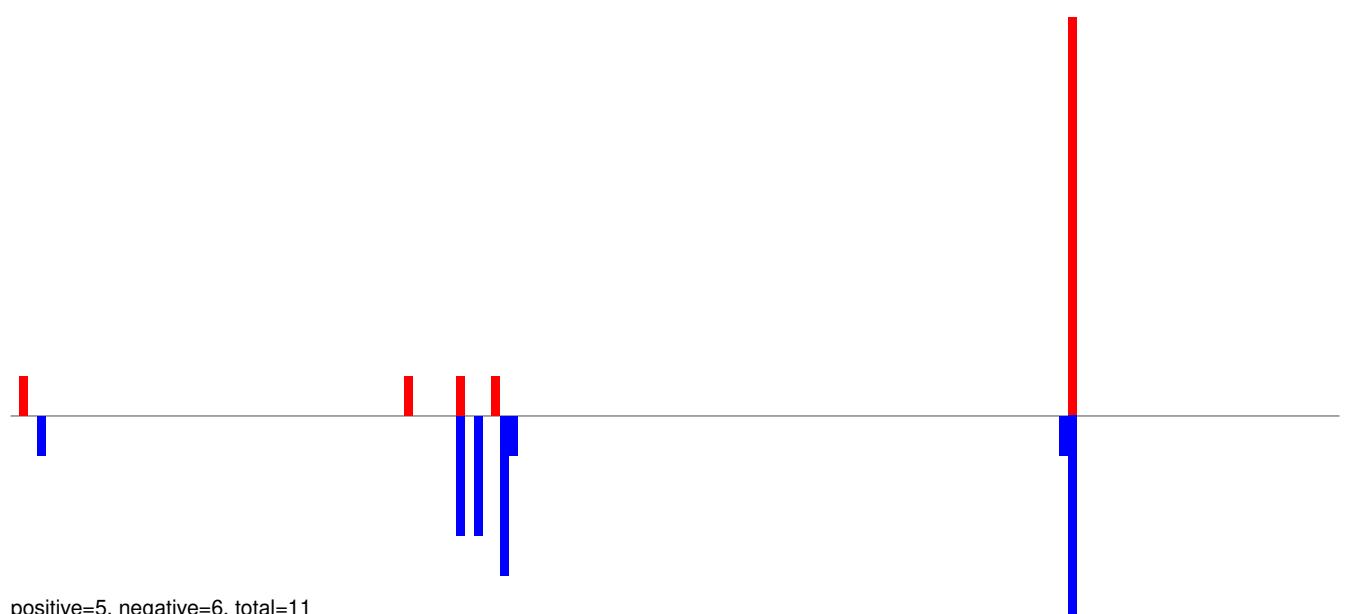
AeAeg_Aag2SINV_GFP.IP.24_35.rep



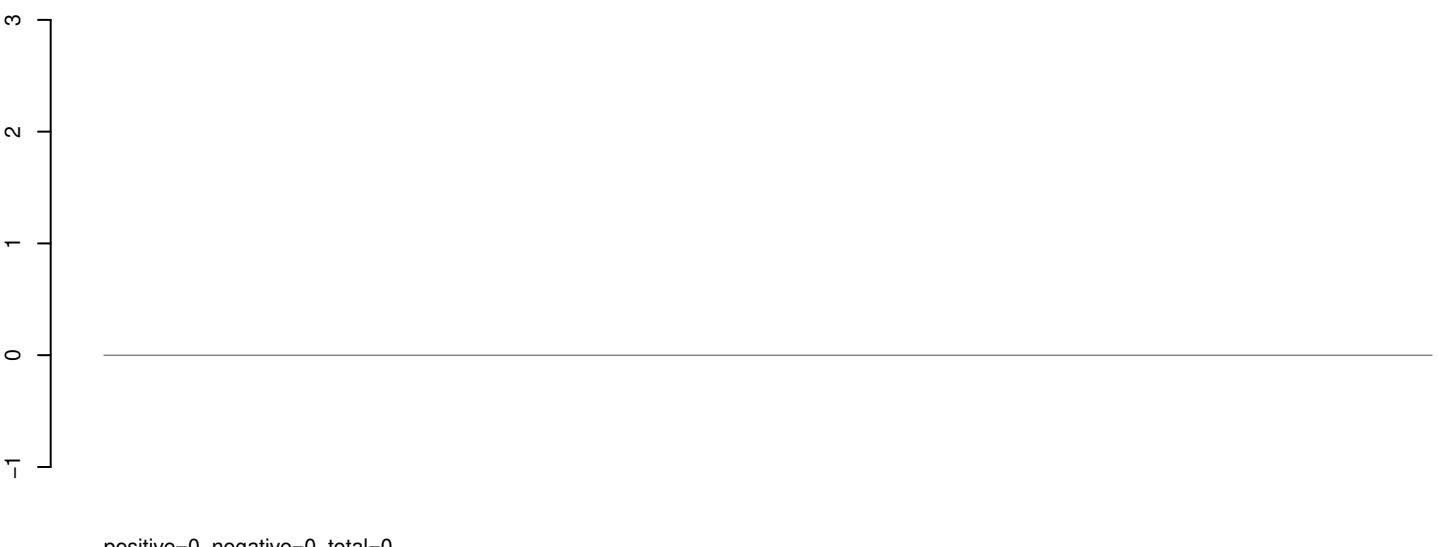
AeAeg_Aag2SINV_GFP.IP.rep



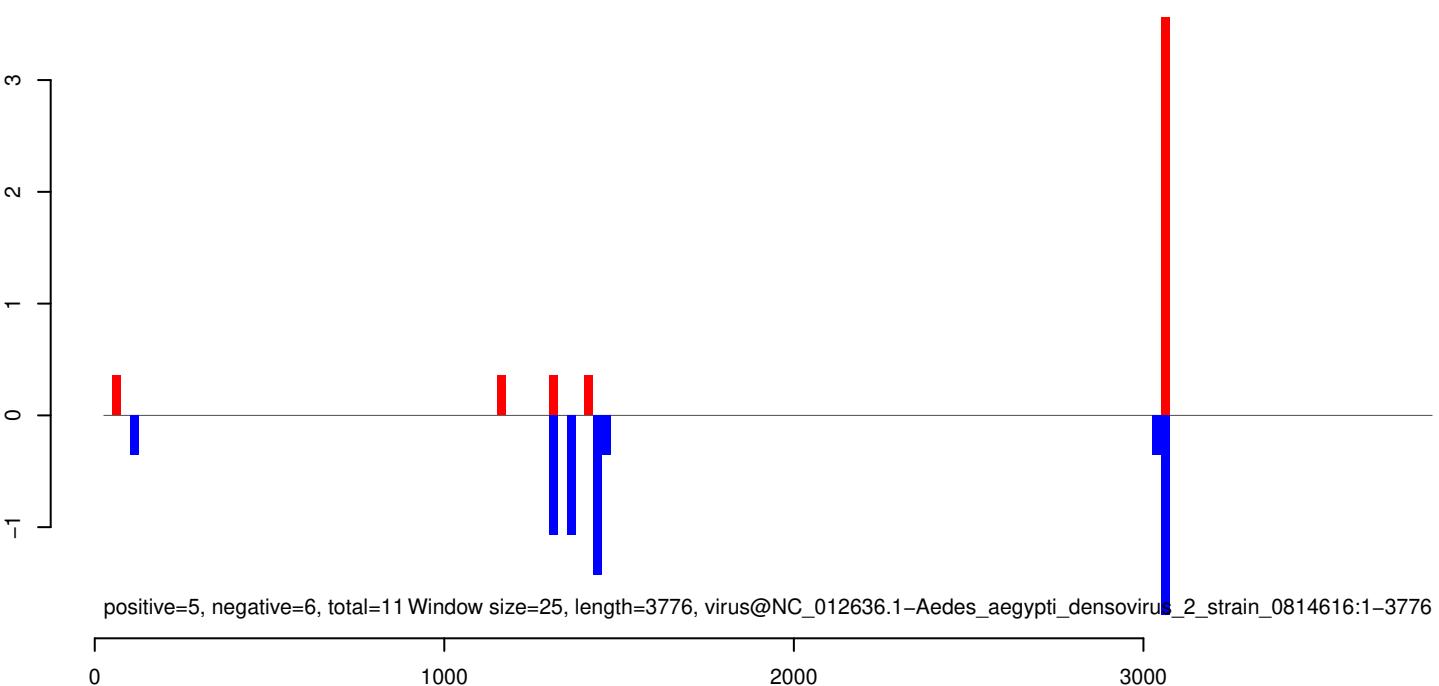
AeAeg_Aag2SINV_GFP.IP.18_23.rep



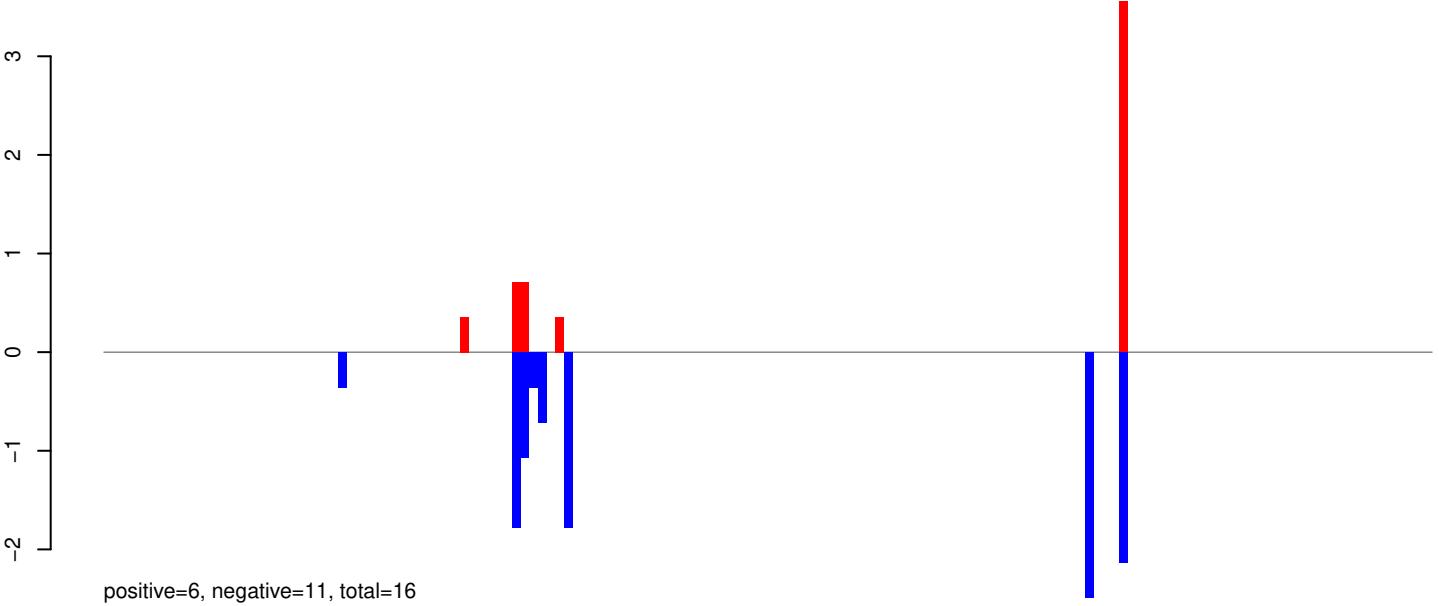
AeAeg_Aag2SINV_GFP.IP.24_35.rep



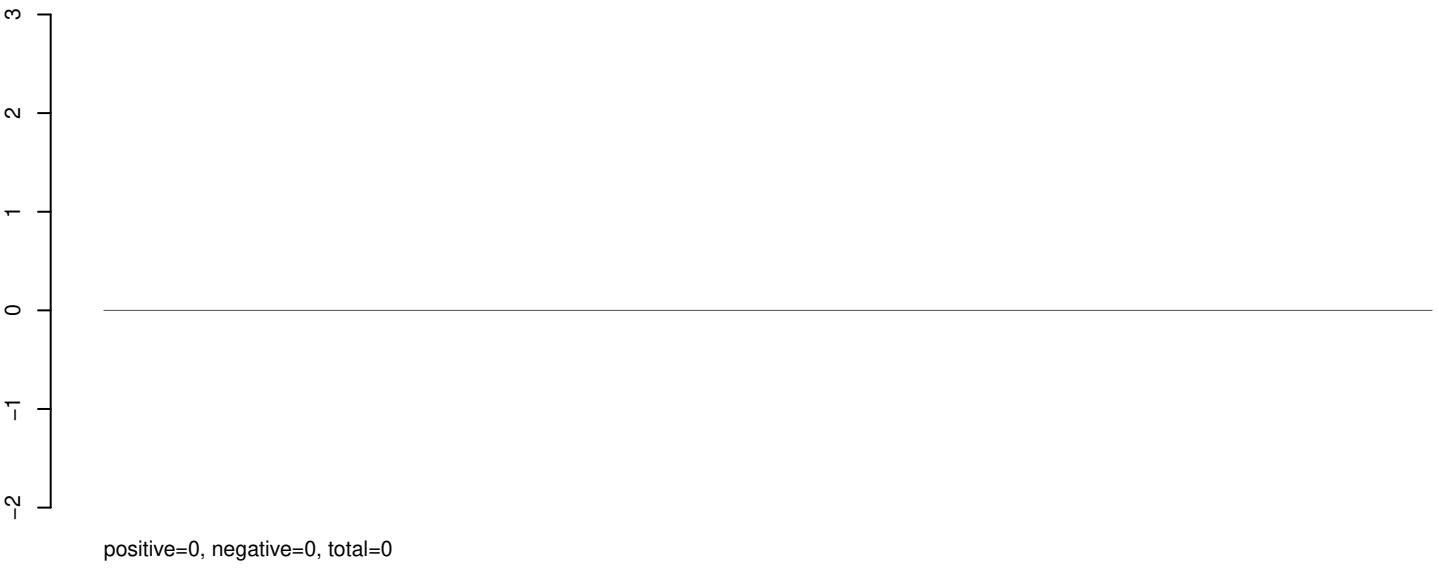
AeAeg_Aag2SINV_GFP.IP.rep



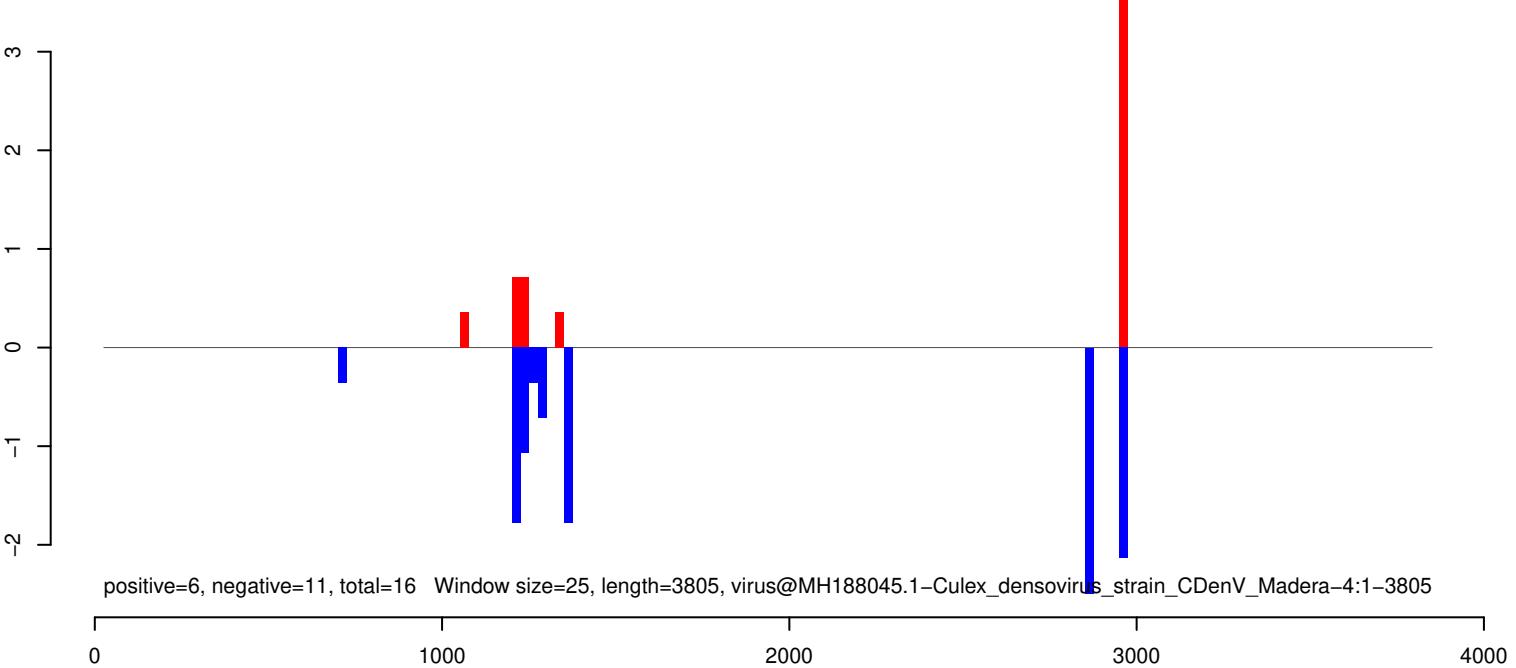
AeAeg_Aag2SINV_GFP.IP.18_23.rep



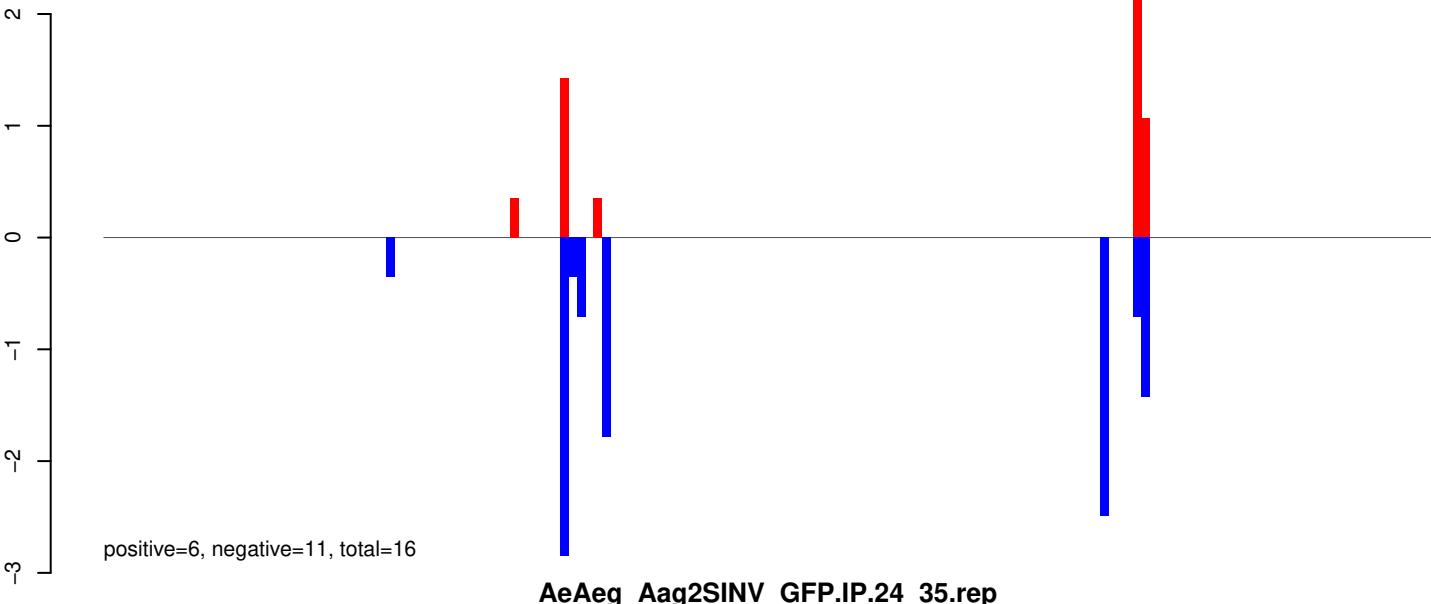
AeAeg_Aag2SINV_GFP.IP.24_35.rep



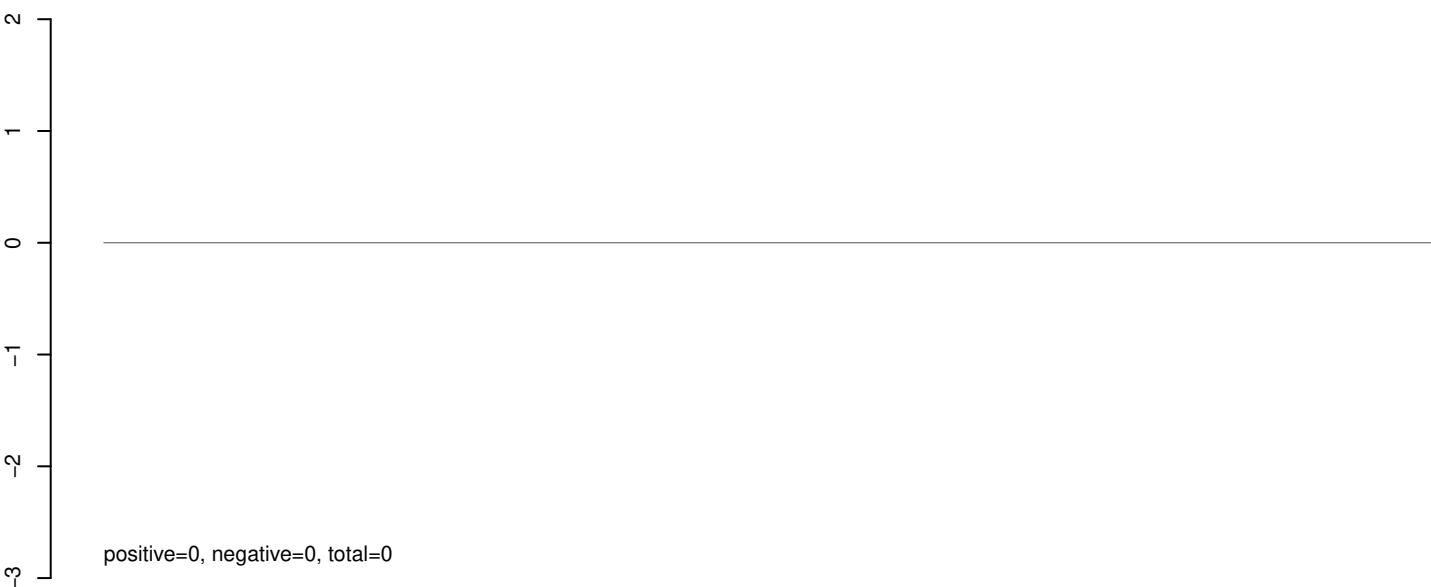
AeAeg_Aag2SINV_GFP.IP.rep



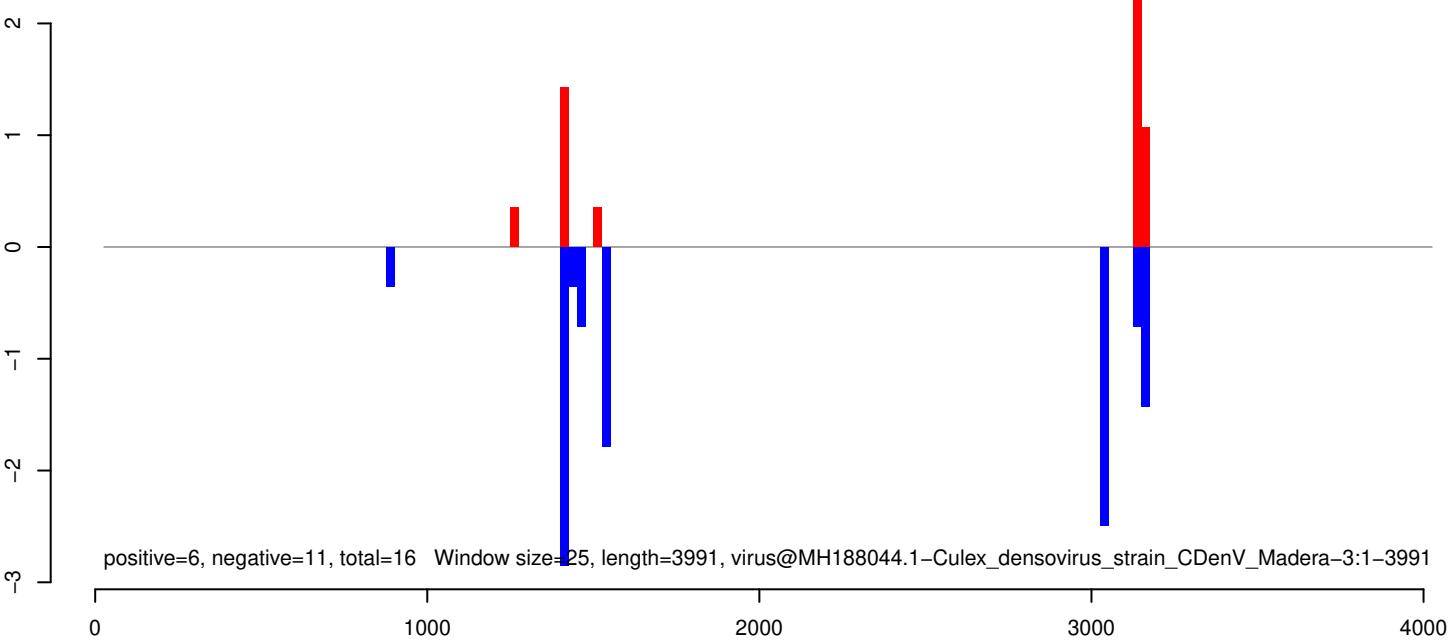
AeAeg_Aag2SINV_GFP.IP.18_23.rep



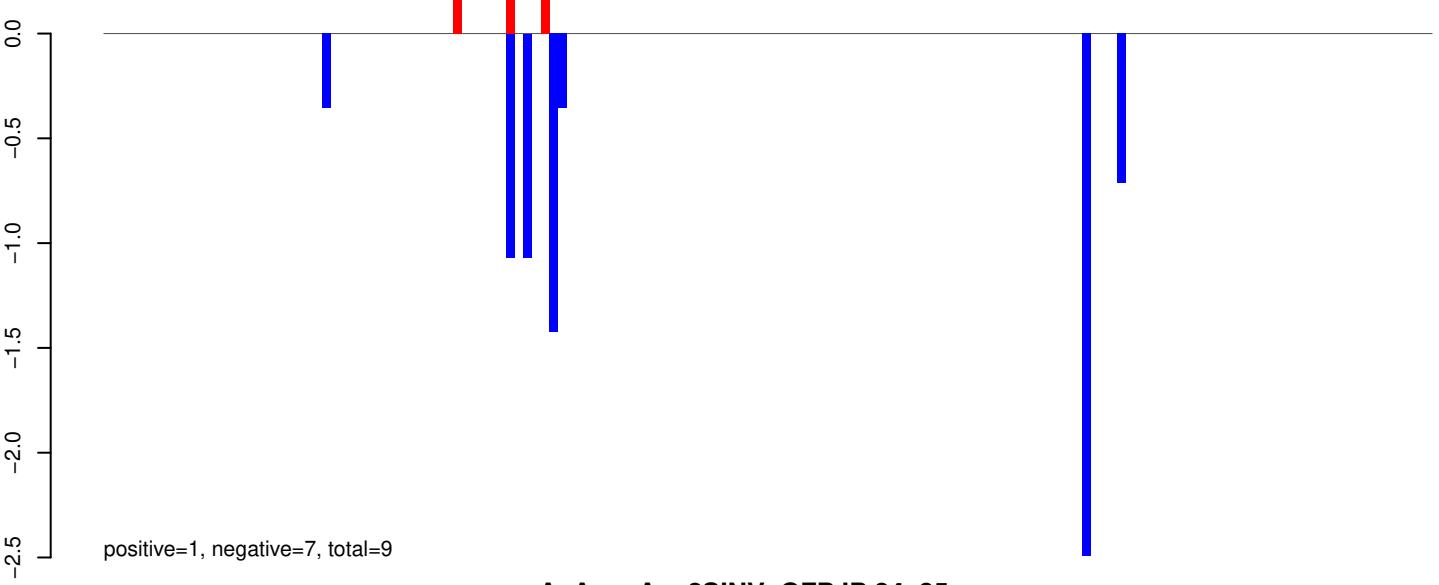
AeAeg_Aag2SINV_GFP.IP.24_35.rep



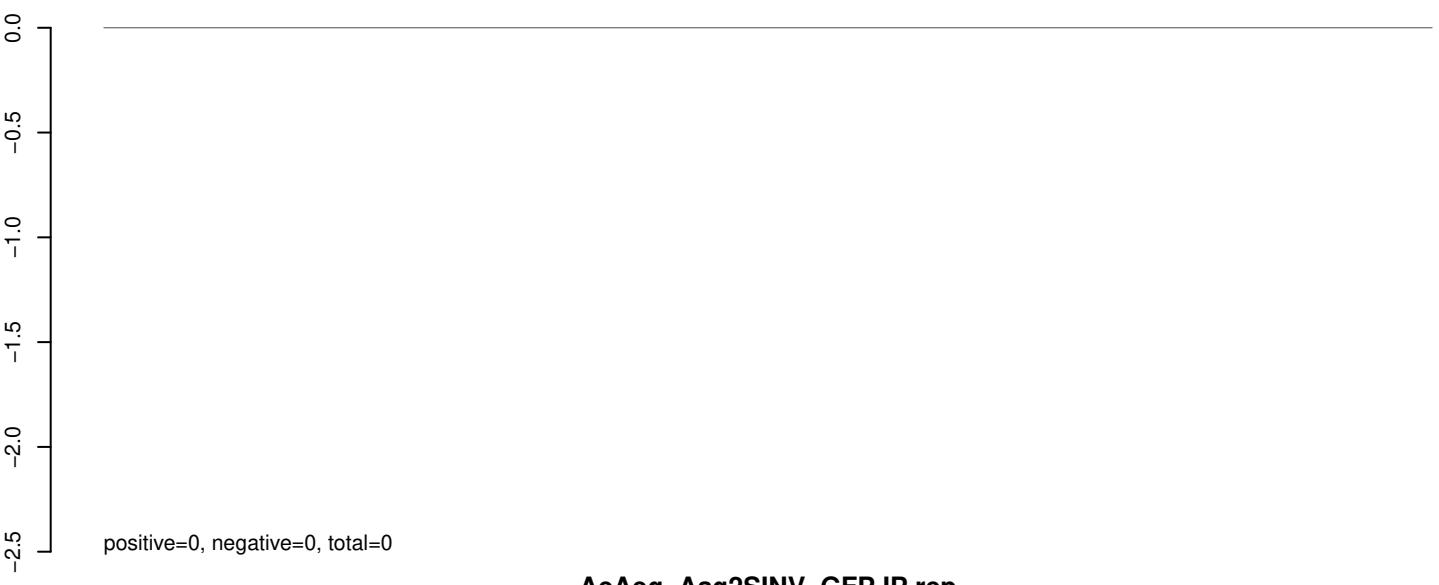
AeAeg_Aag2SINV_GFP.IP.rep



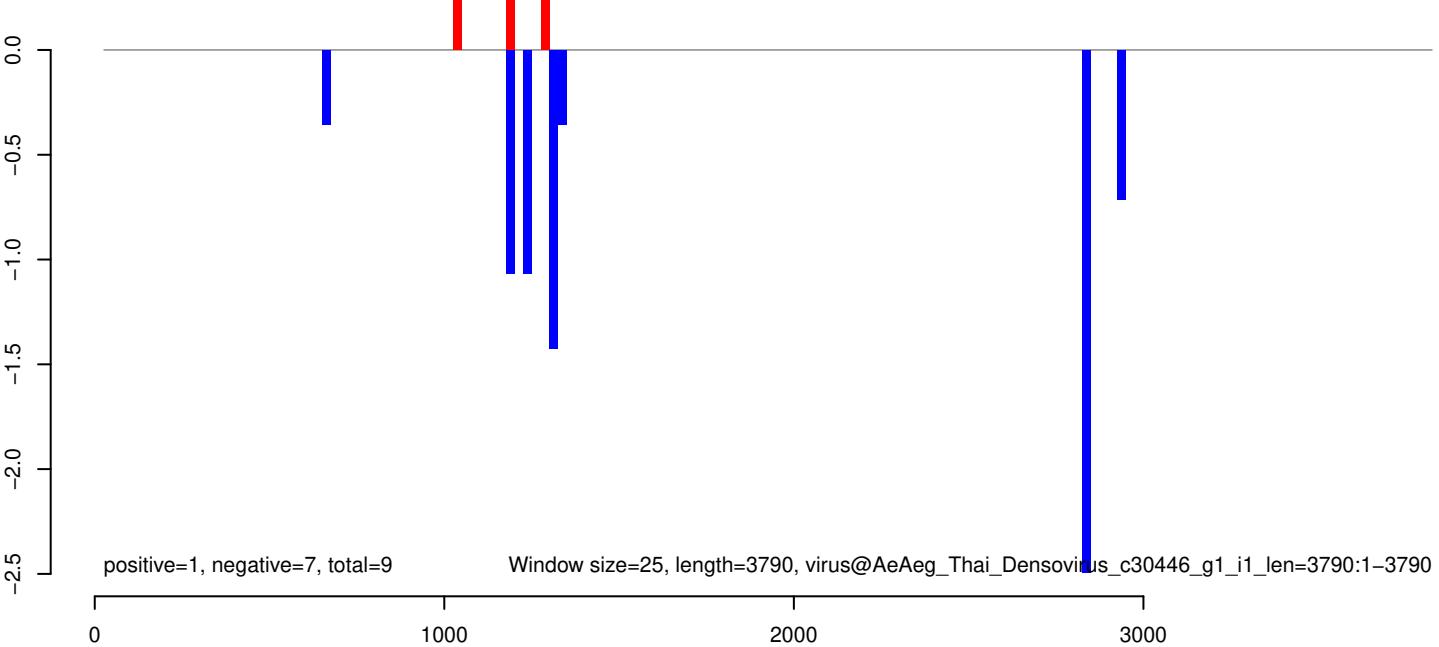
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=3, negative=6, total=9

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=1, negative=0, total=1

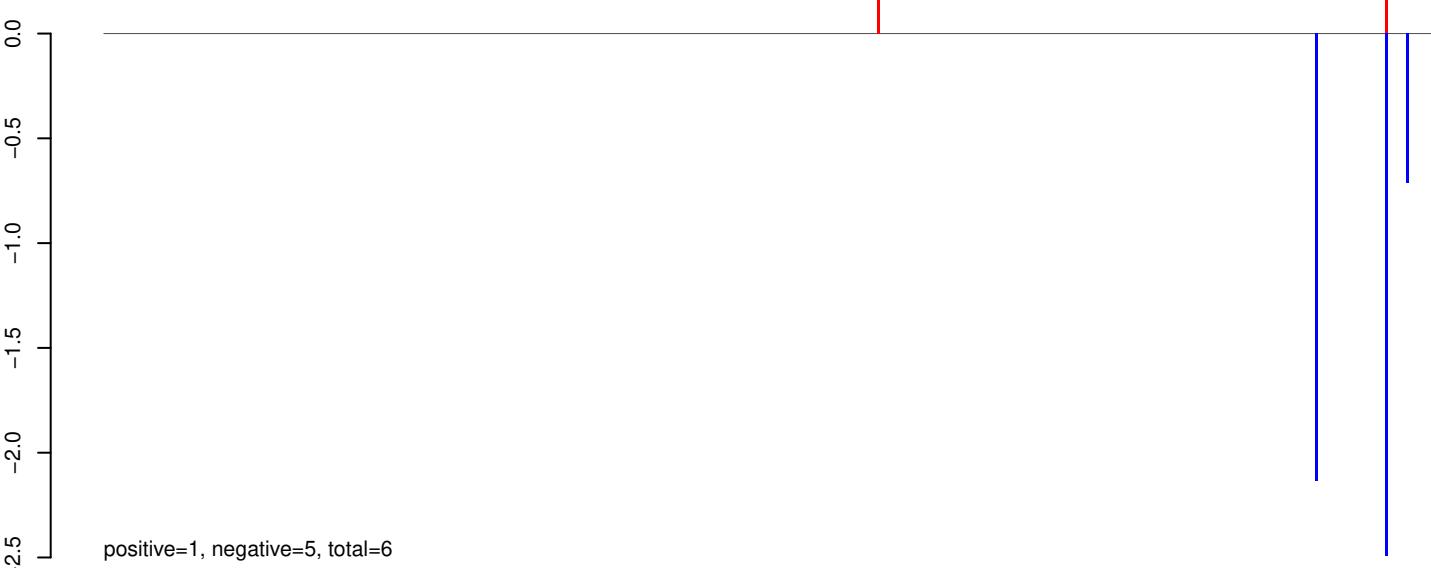
AeAeg_Aag2SINV_GFP.IP.rep

positive=4, negative=6, total=10

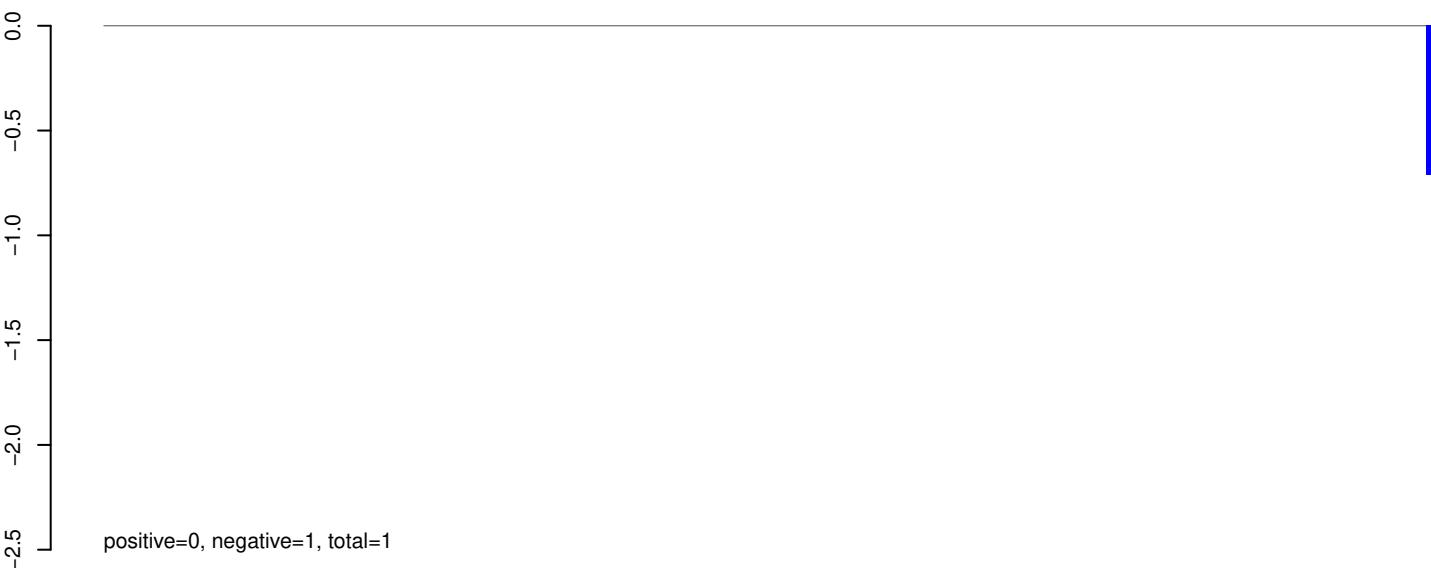
Window size=25, length=11381, virus@NC_013528.1–Fort_Morgan_virus:1–11381



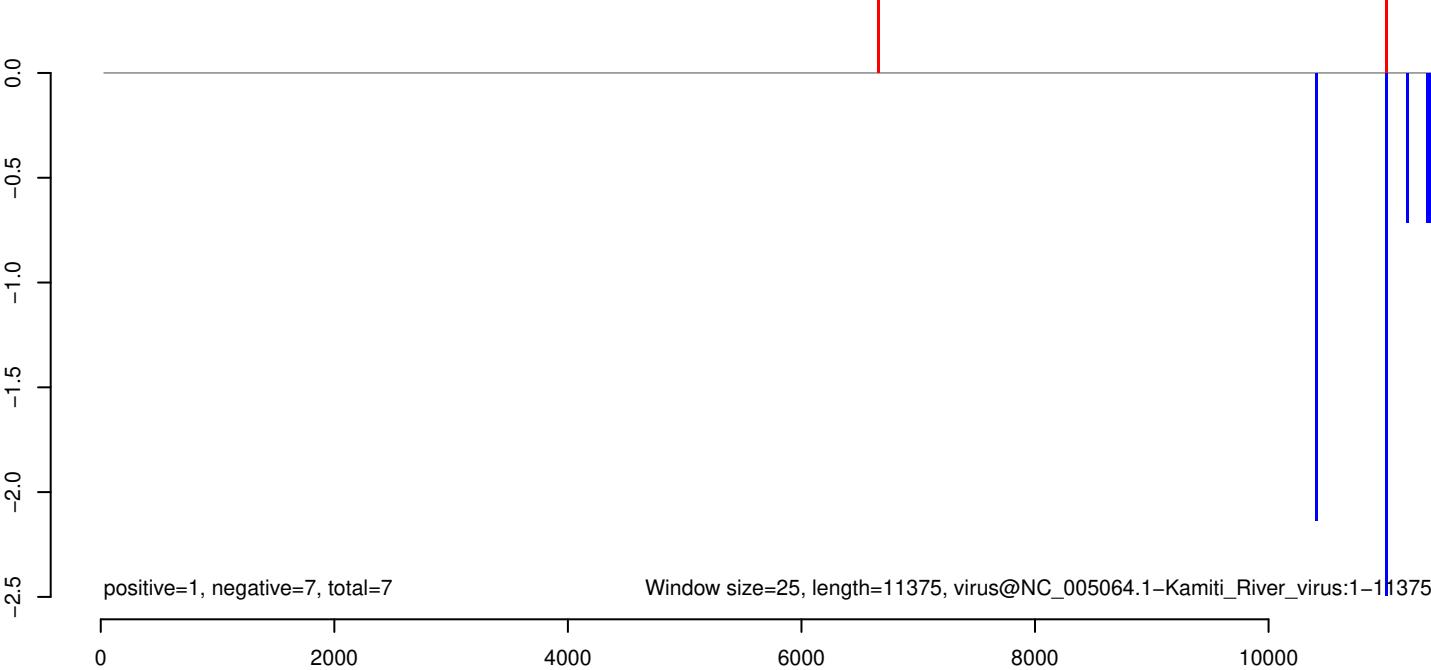
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep

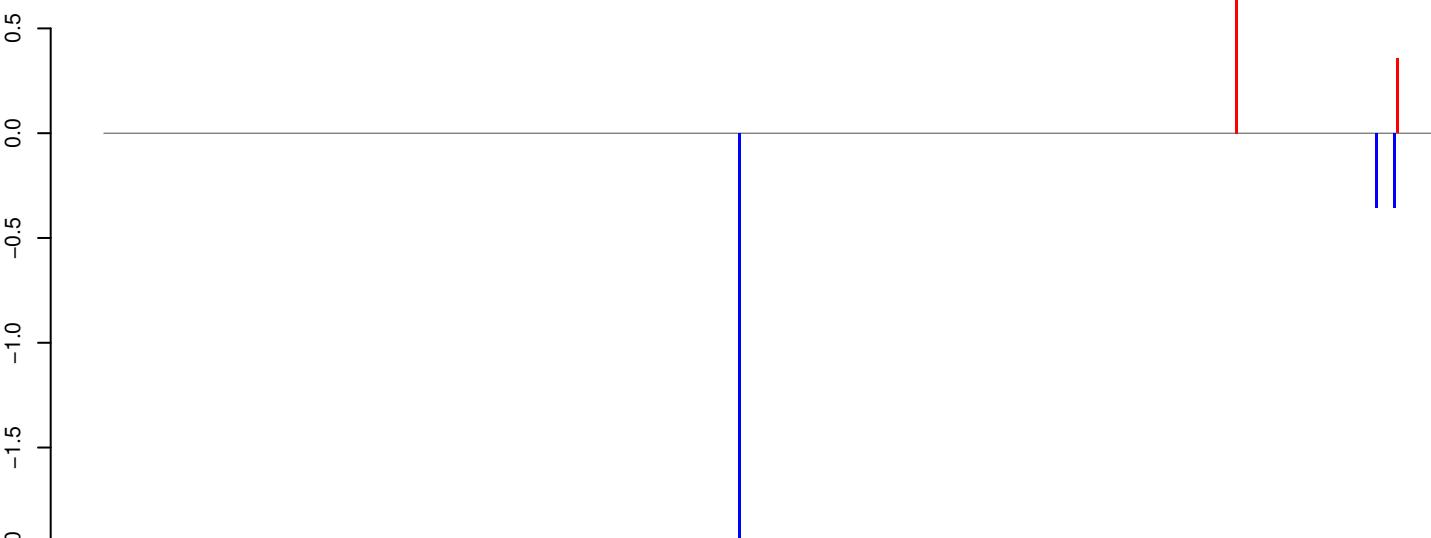


AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=1, negative=3, total=4

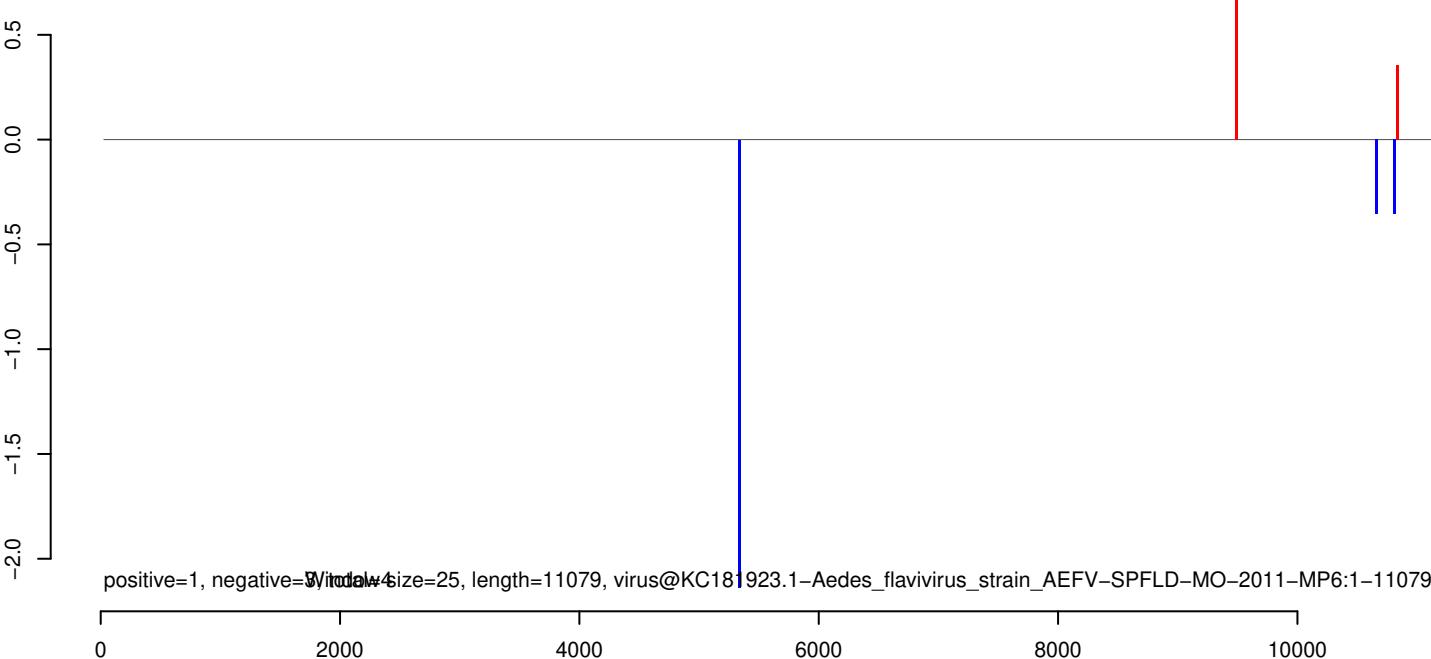


AeAeg_Aag2SINV_GFP.IP.24_35.rep

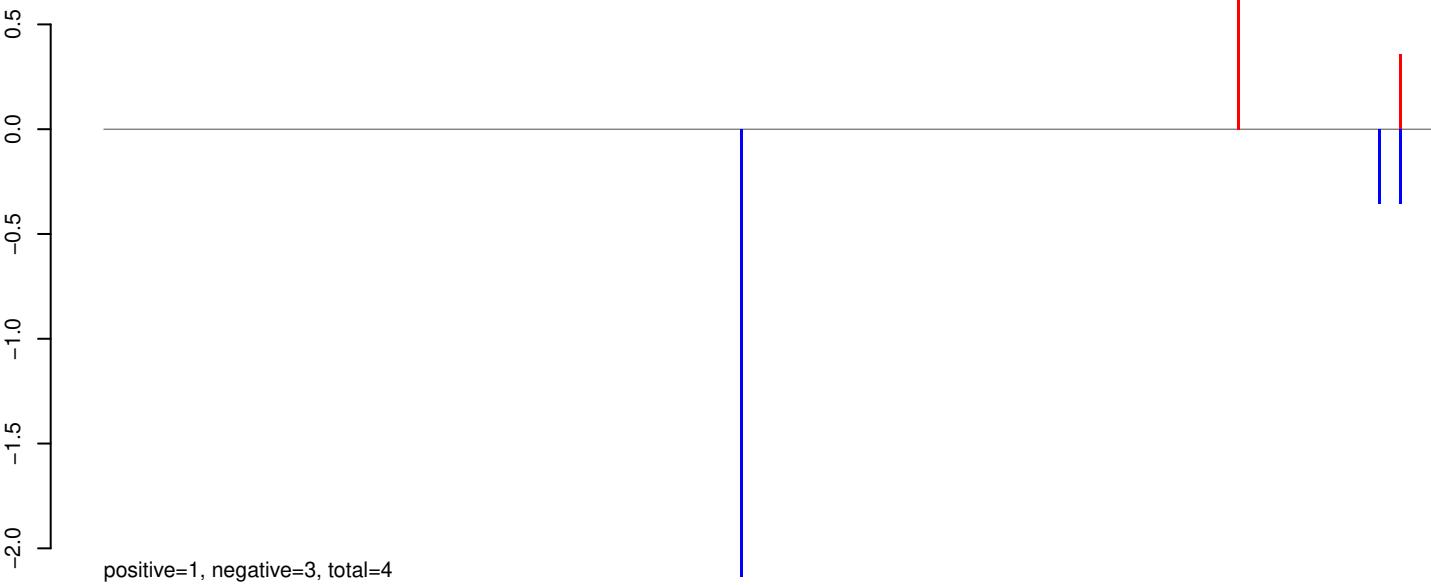
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

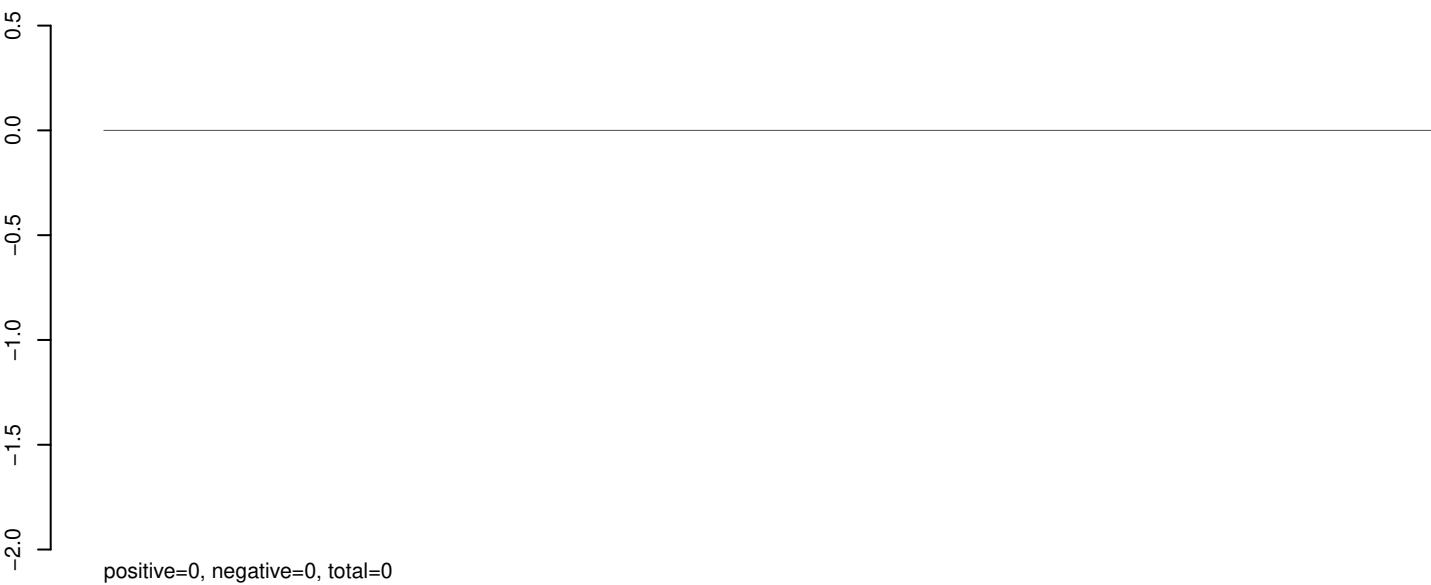
positive=1, negative=5, window=4, size=25, length=11079, virus@KC181923.1-Aedes_flavivirus_strain_AEFV-SPFLD-MO-2011-MP6:1-11079



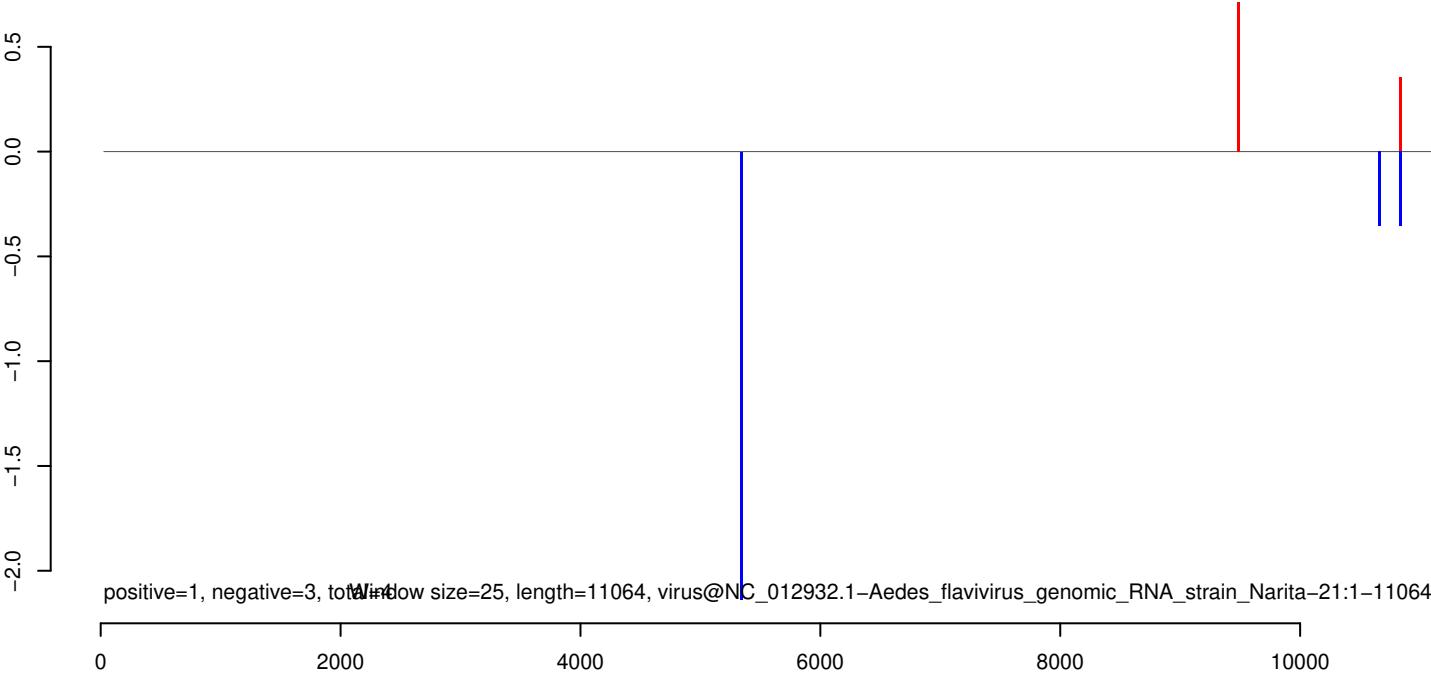
AeAeg_Aag2SINV_GFP.IP.18_23.rep



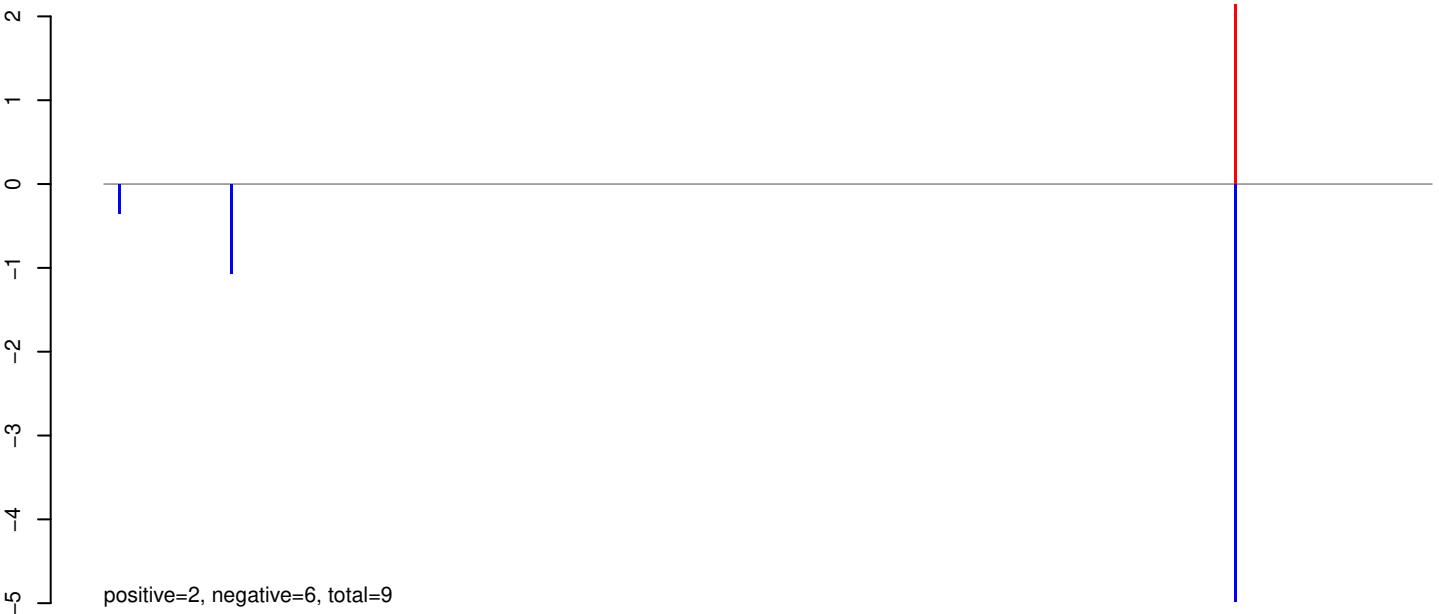
AeAeg_Aag2SINV_GFP.IP.24_35.rep



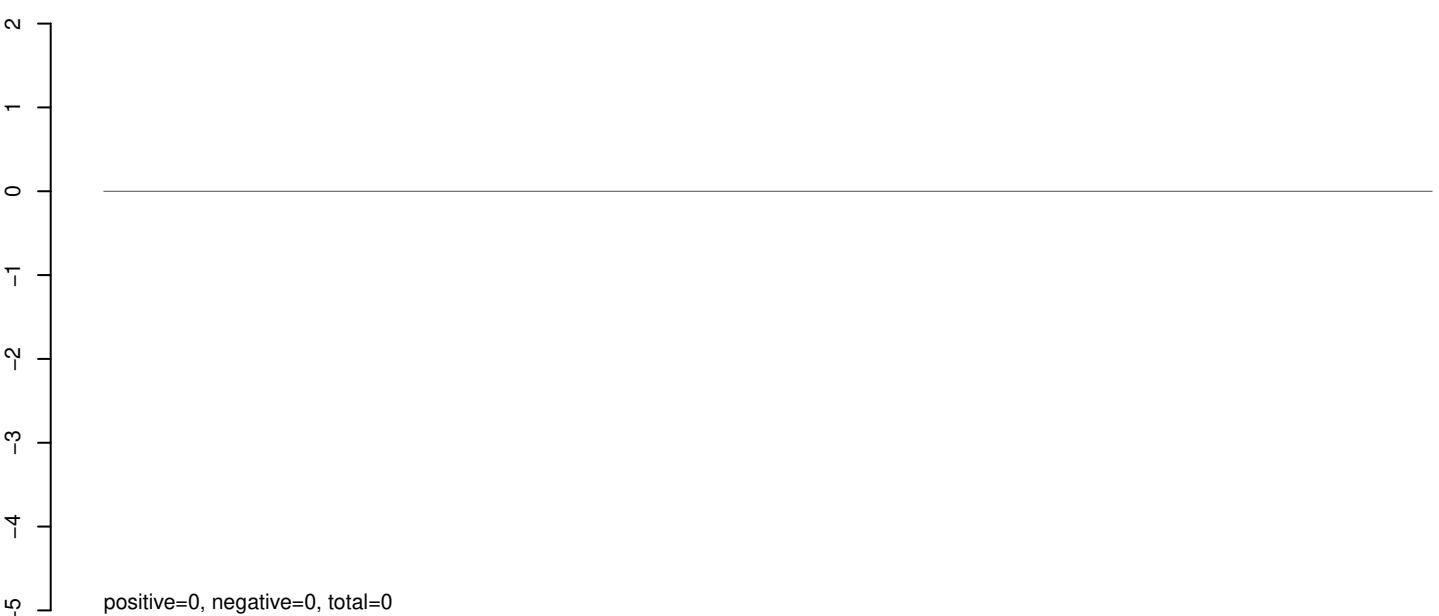
AeAeg_Aag2SINV_GFP.IP.rep



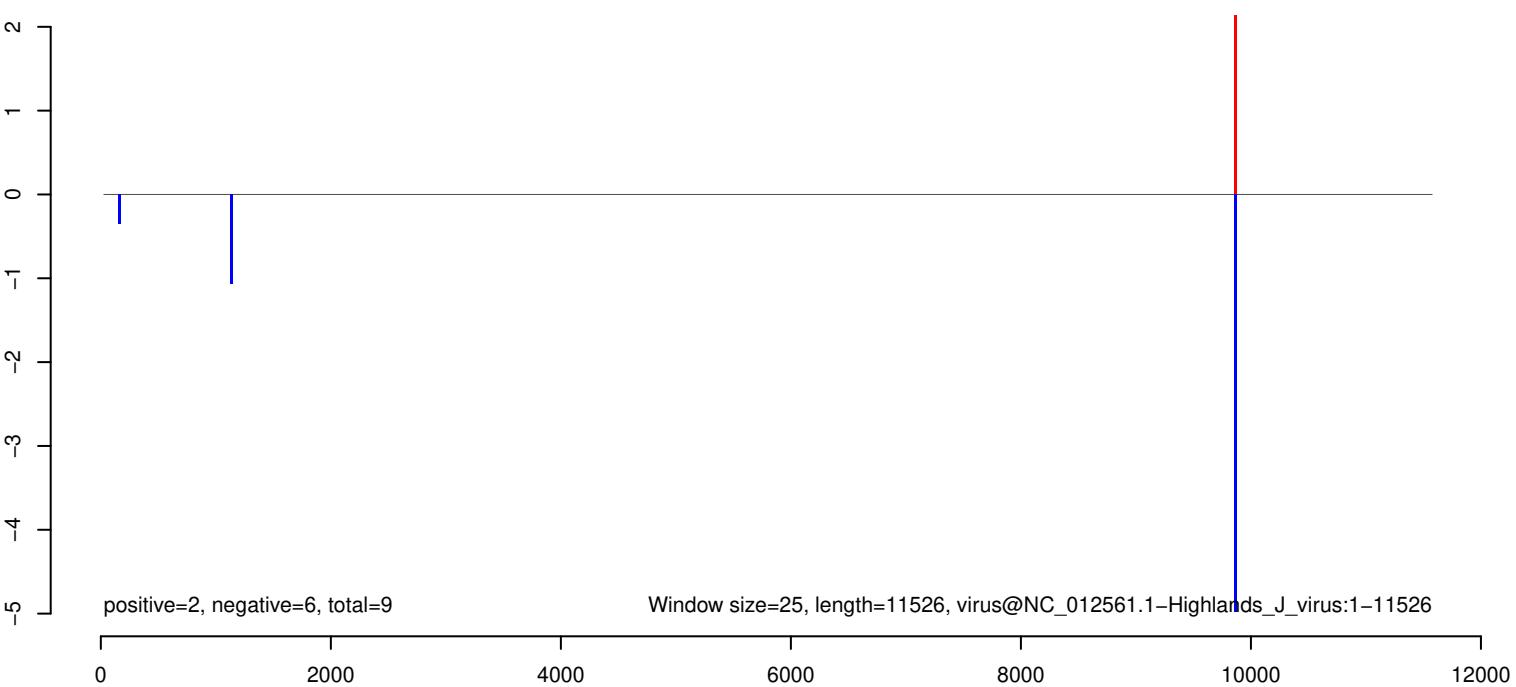
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep

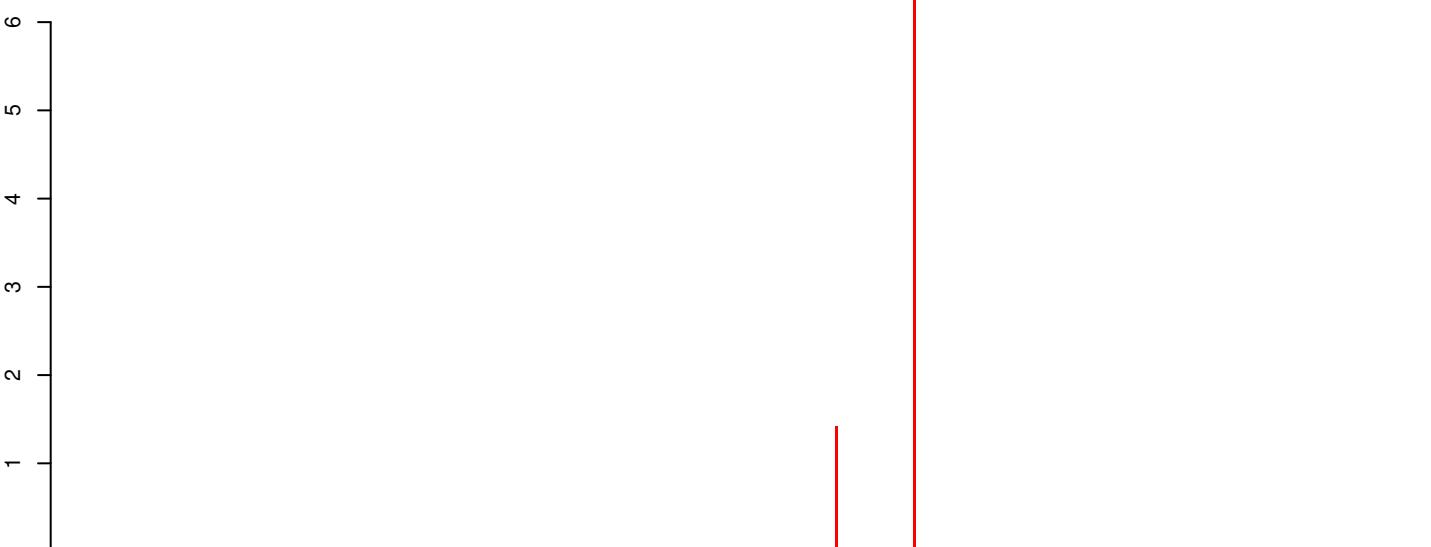


AeAeg_Aag2SINV_GFP.IP.rep



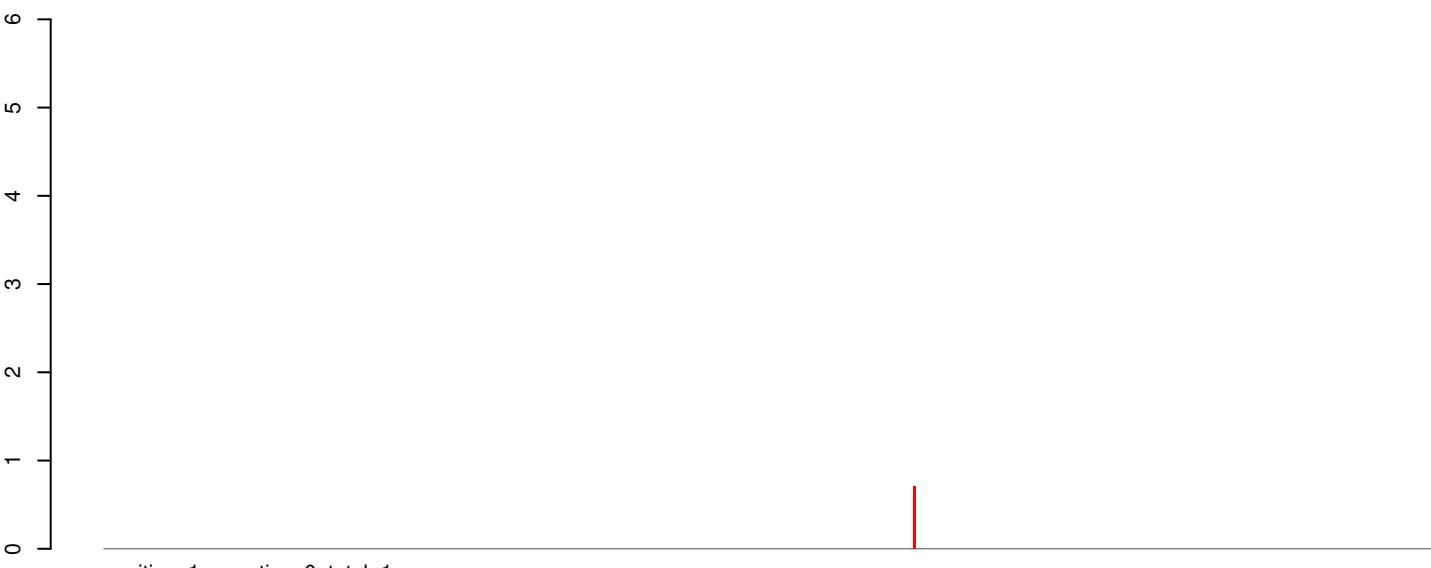
AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=8, negative=0, total=8



AeAeg_Aag2SINV_GFP.IP.24_35.rep

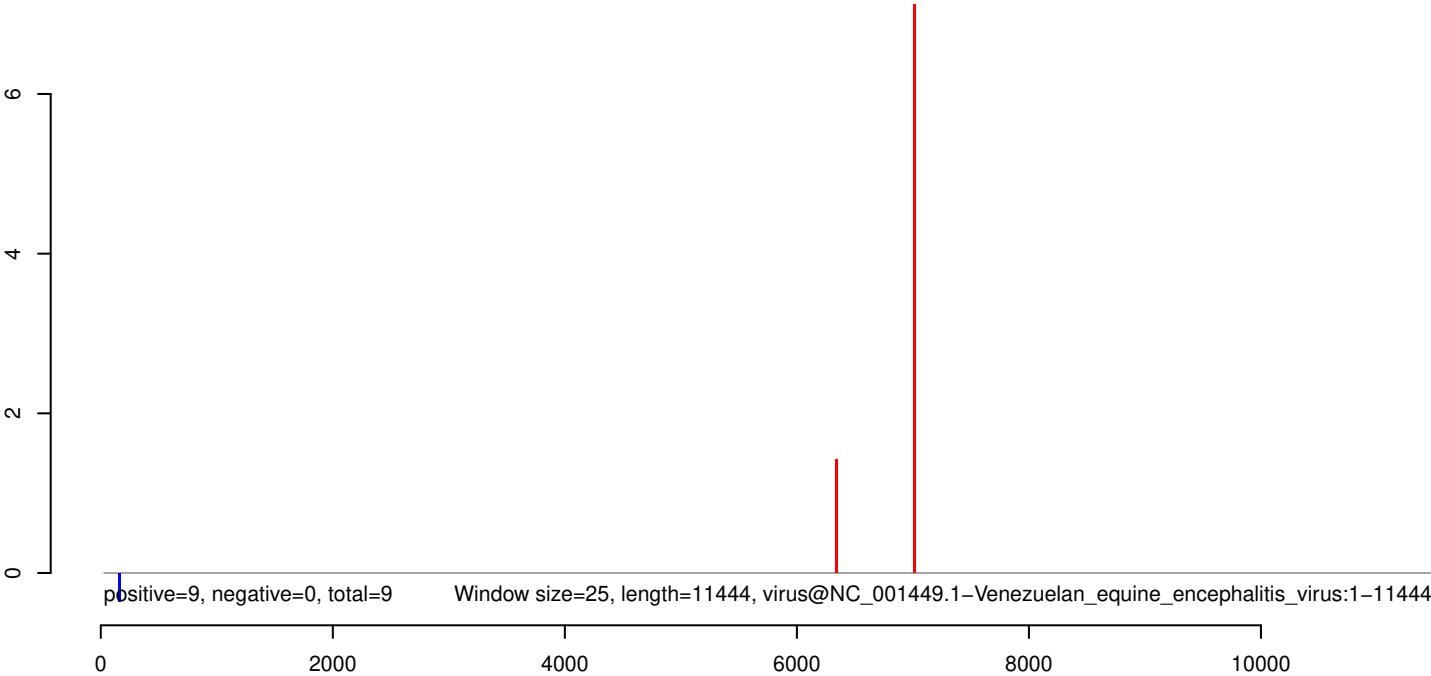
positive=1, negative=0, total=1



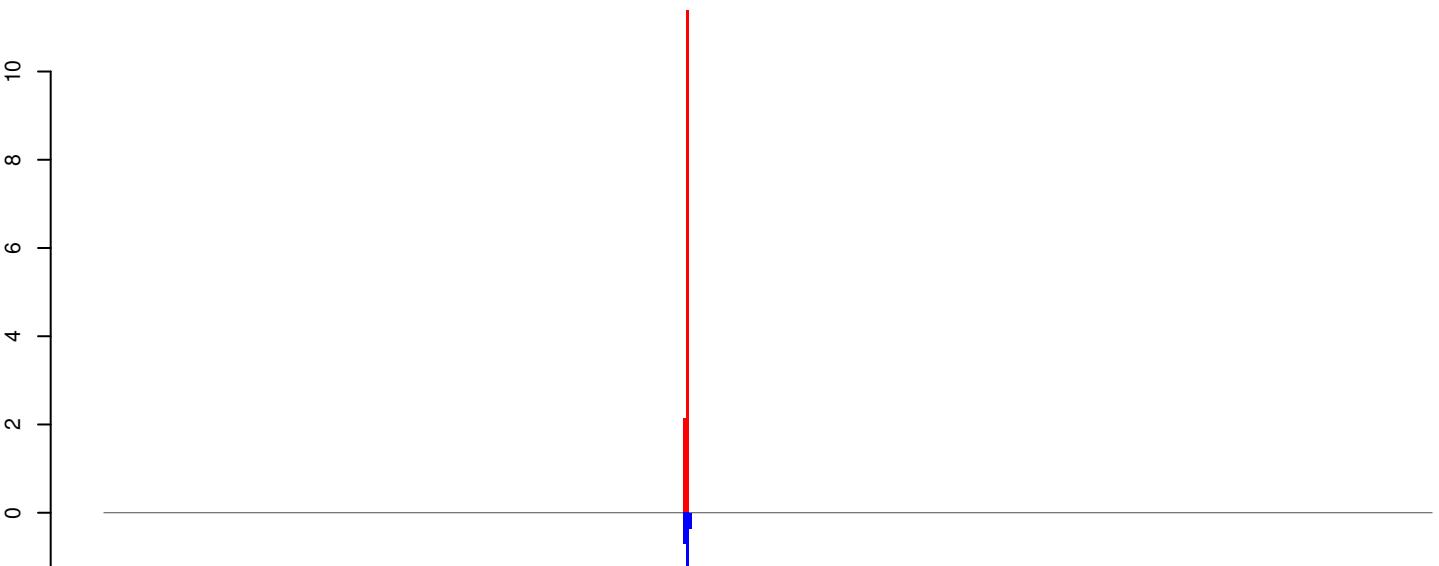
AeAeg_Aag2SINV_GFP.IP.rep

positive=9, negative=0, total=9

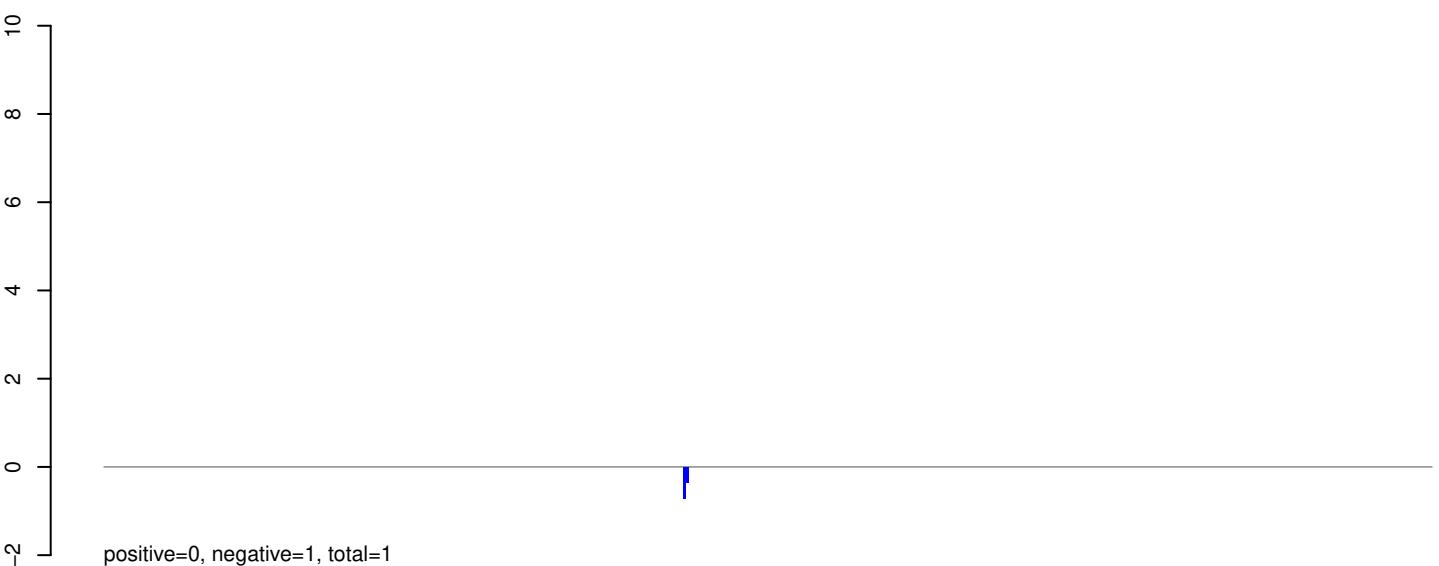
Window size=25, length=11444, virus@NC_001449.1–Venezuelan equine encephalitis virus:1–11444



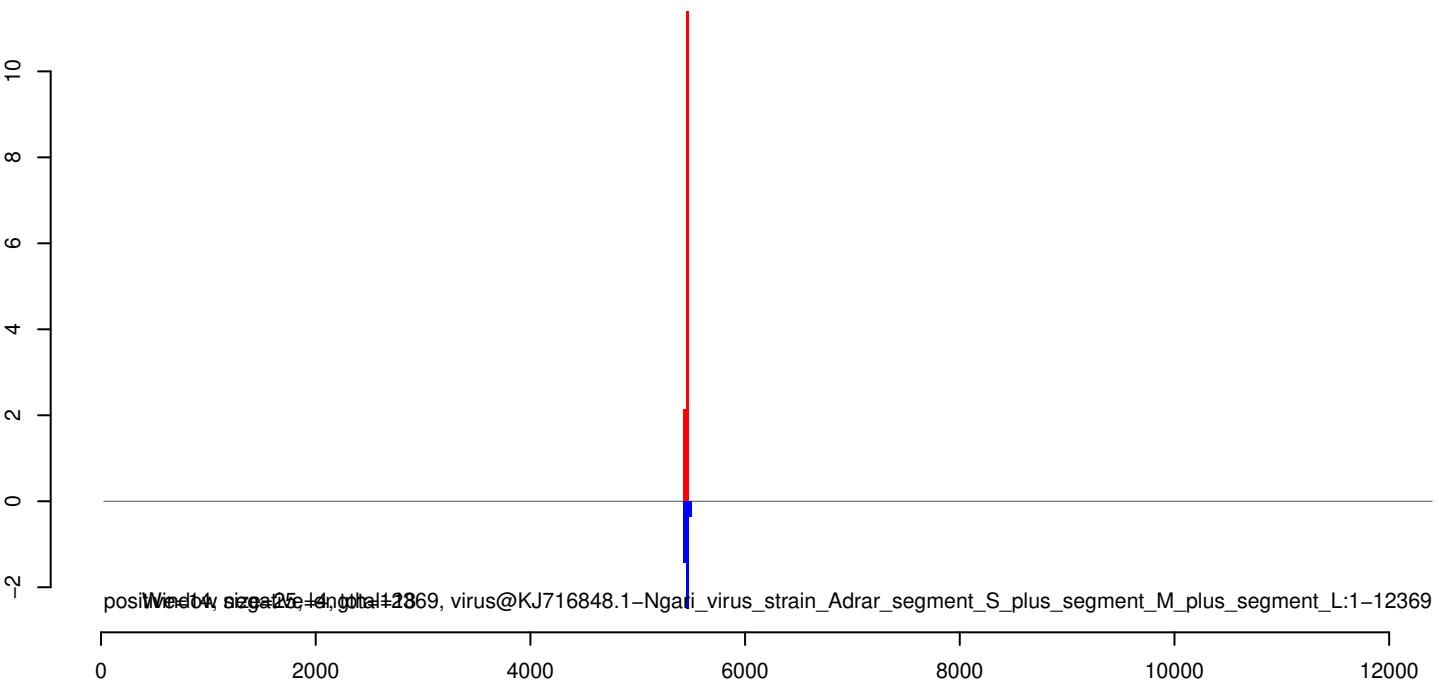
AeAeg_Aag2SINV_GFP.IP.18_23.rep



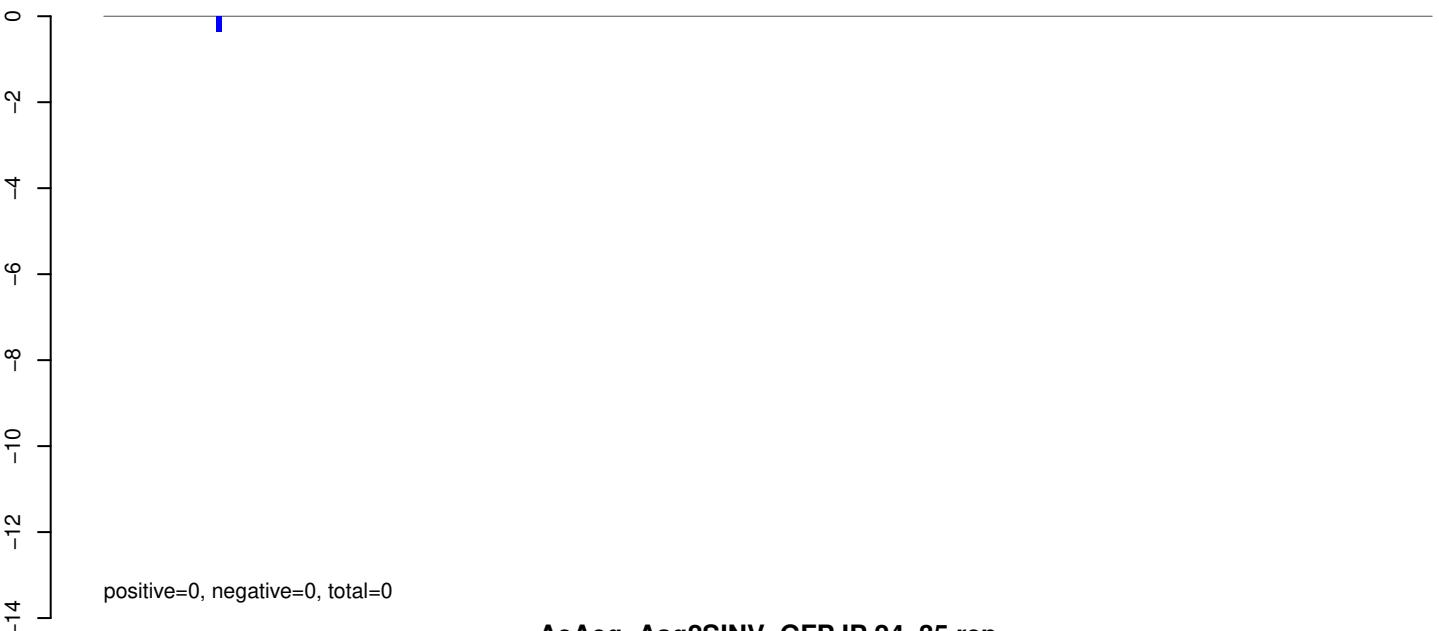
AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



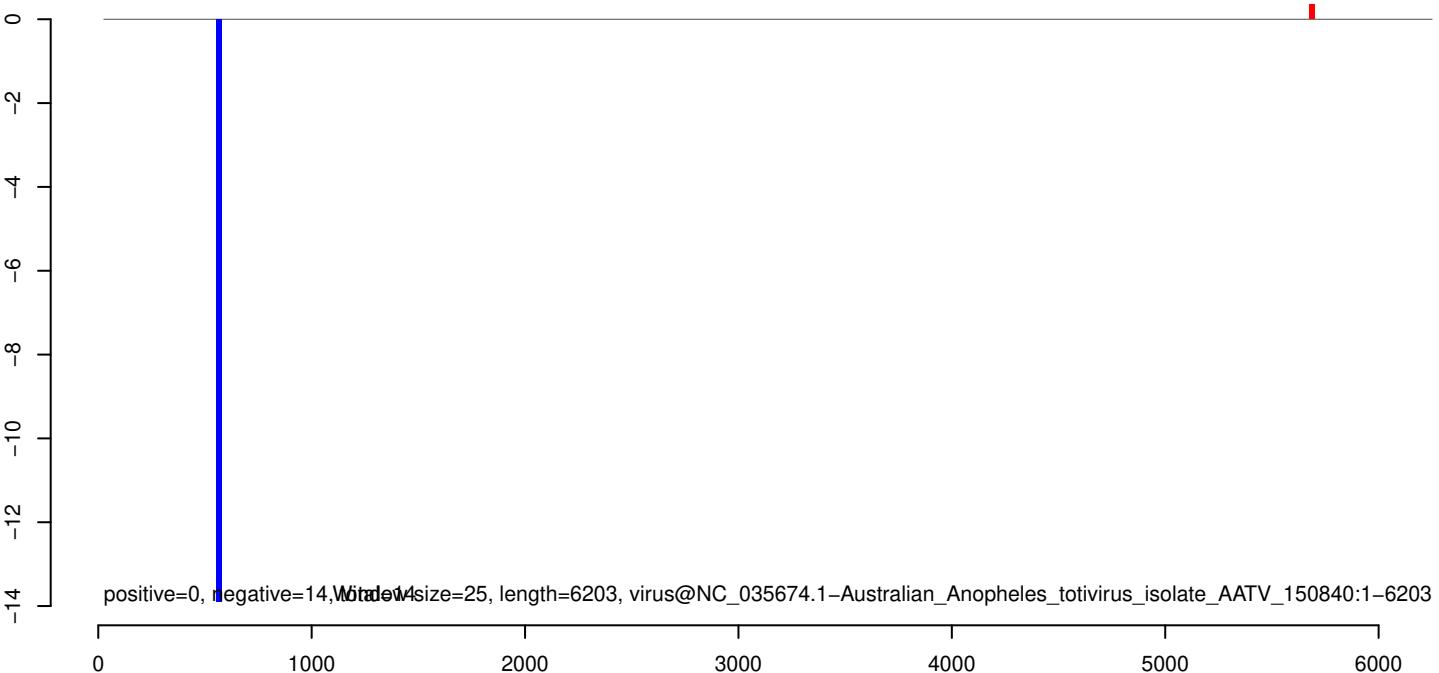
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=2, negative=0, total=2

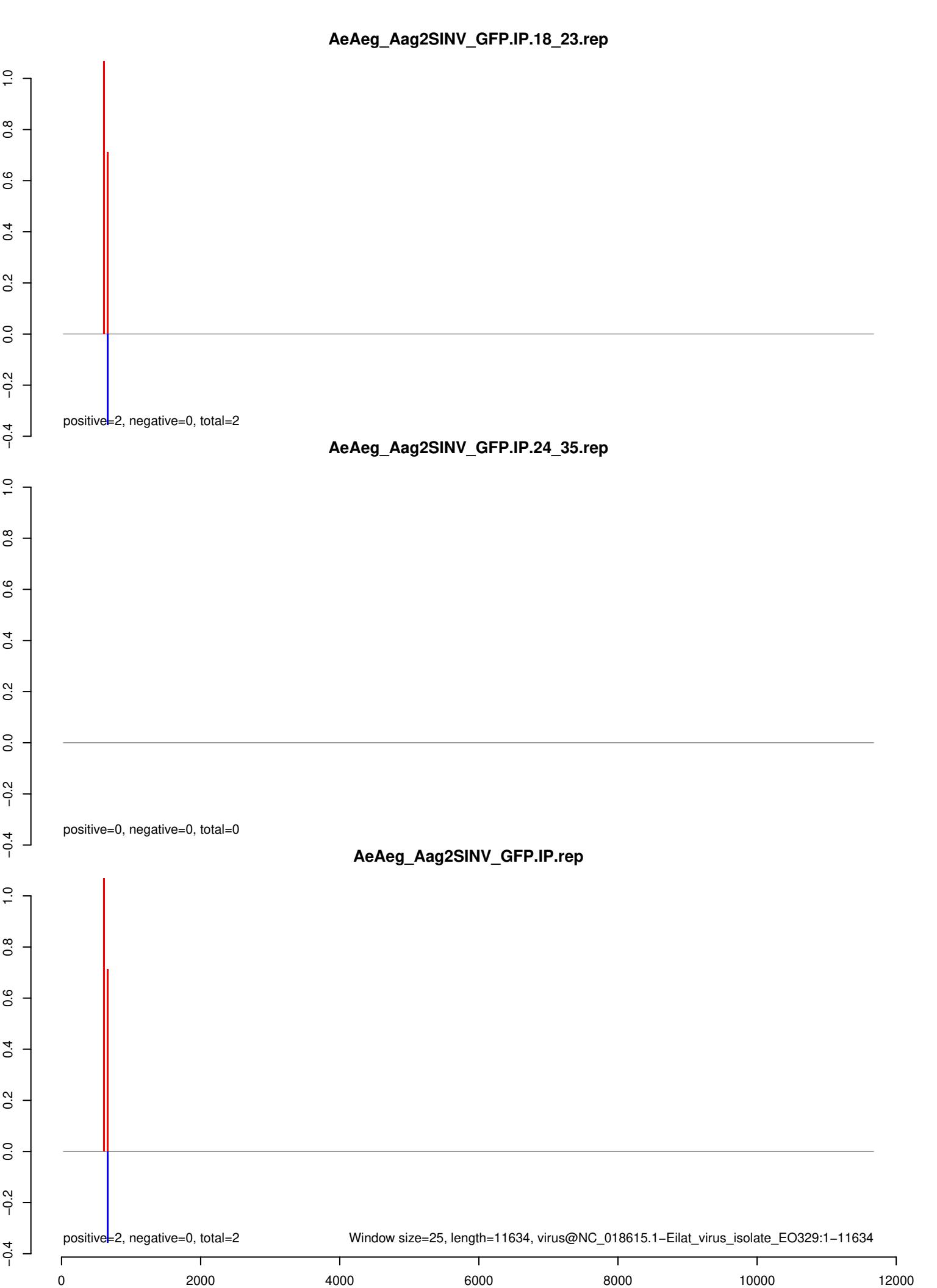
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=2, negative=0, total=2

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



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=2, negative=0, total=2



AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0



AeAeg_Aag2SINV_GFP.IP.rep

positive=2, negative=0, total=2

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

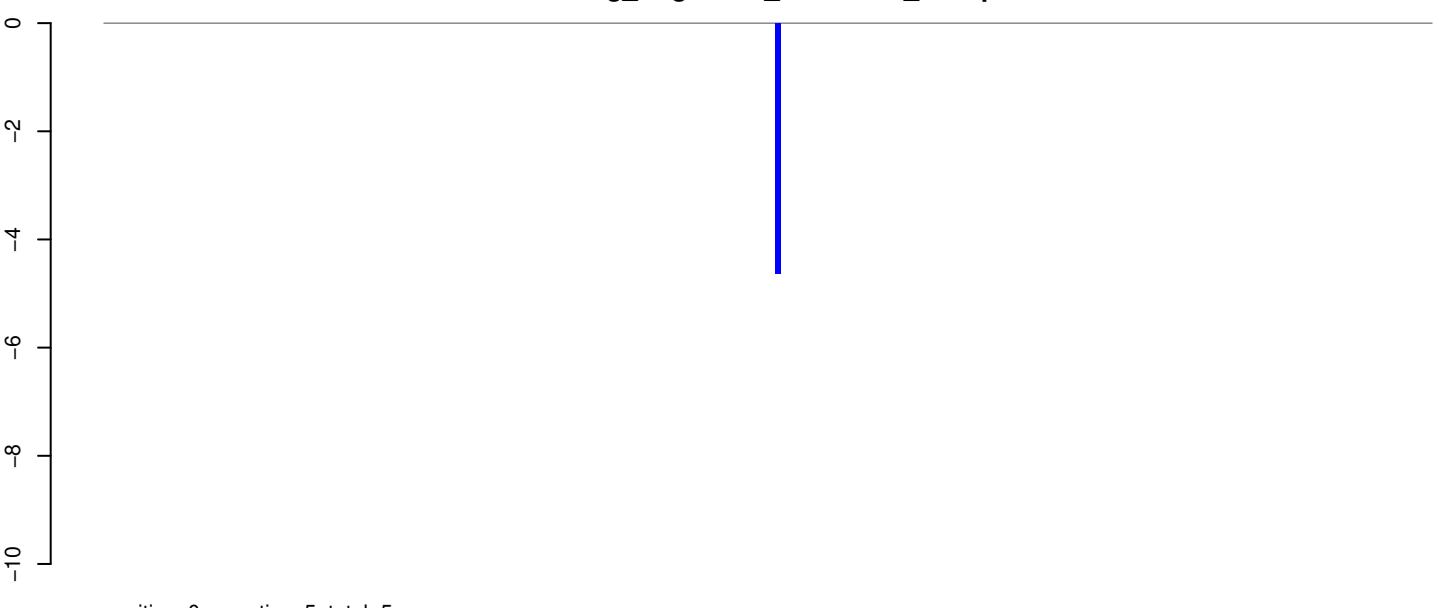


0 2000 4000 6000 8000 10000 12000

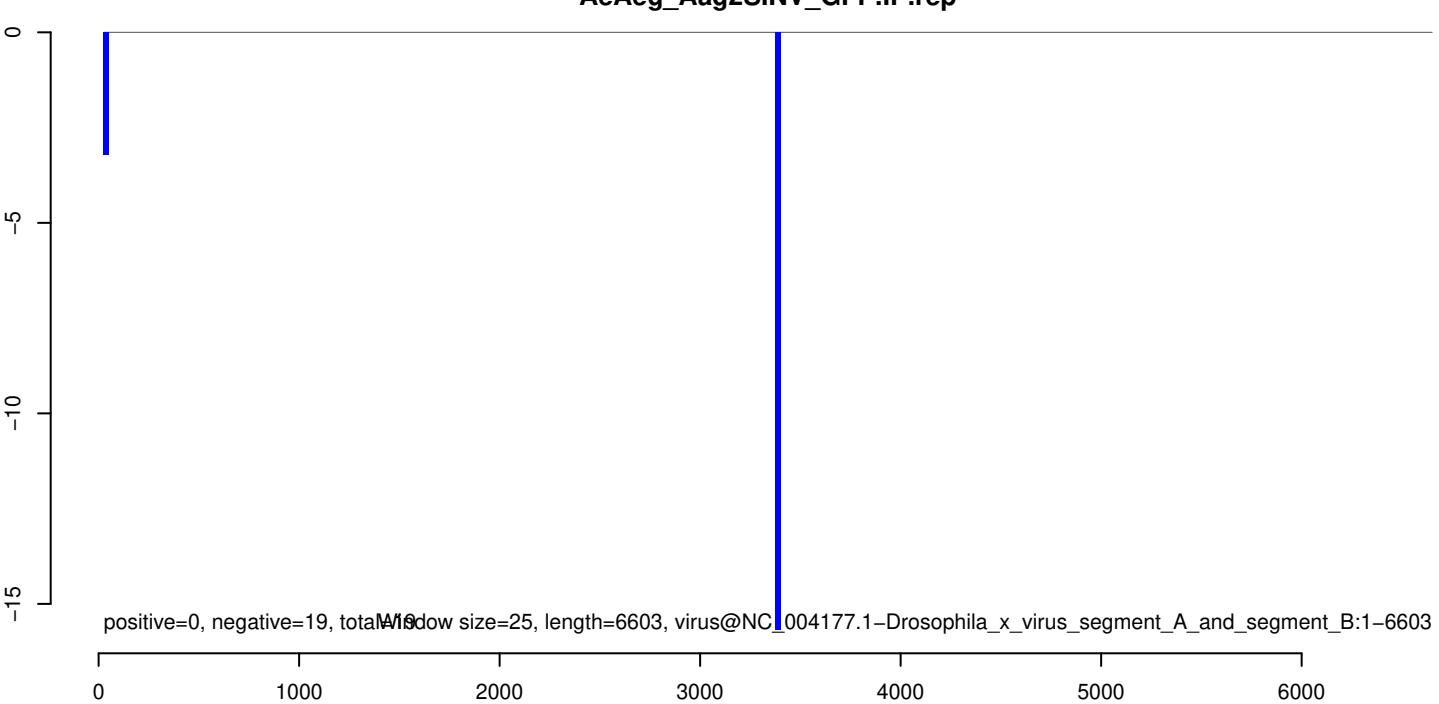
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=1, total=1

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=1, total=1

Window size=25, length=11488, virus@NC_001786.1-Barmah_Forest_virus:1-11488

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=1



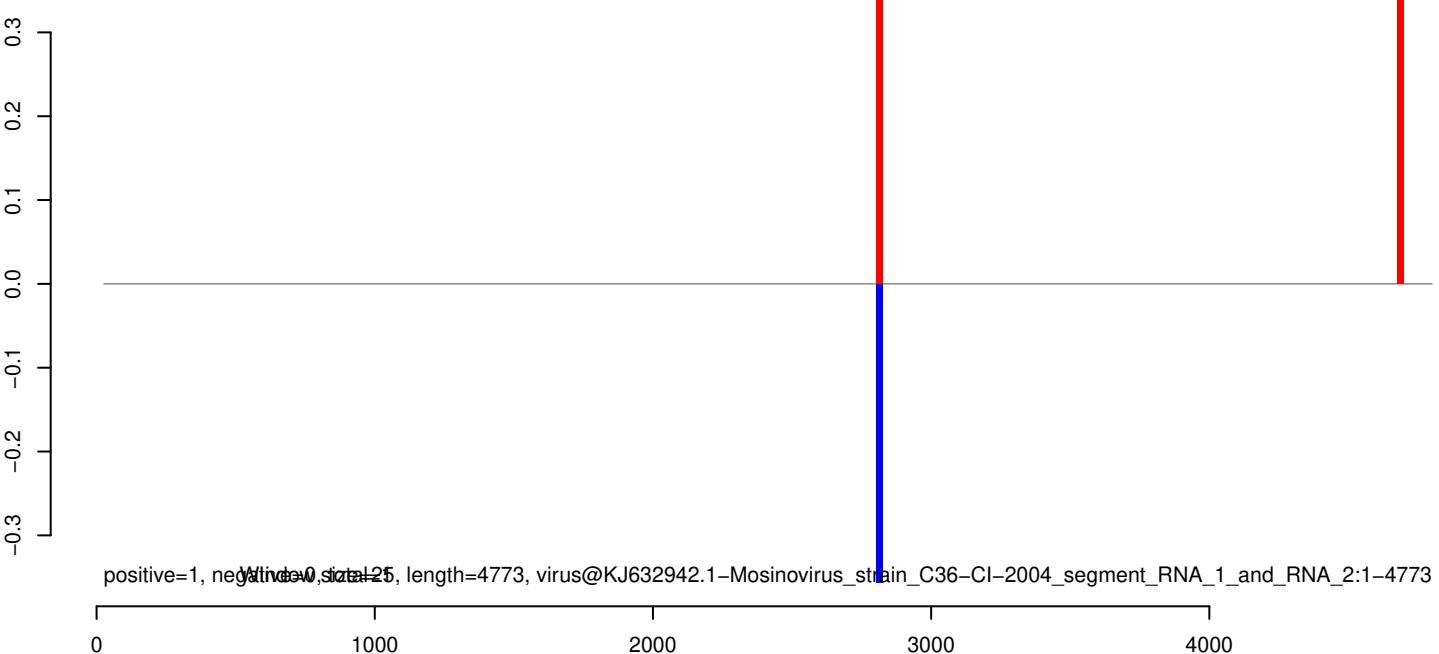
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0



AeAeg_Aag2SINV_GFP.IP.rep

positive=1, negative=0, size=25, length=4773, virus@KJ632942.1-Mosinovirus_stain_C36-Cl-2004_segment_RNA_1_and_RNA_2:1-4773



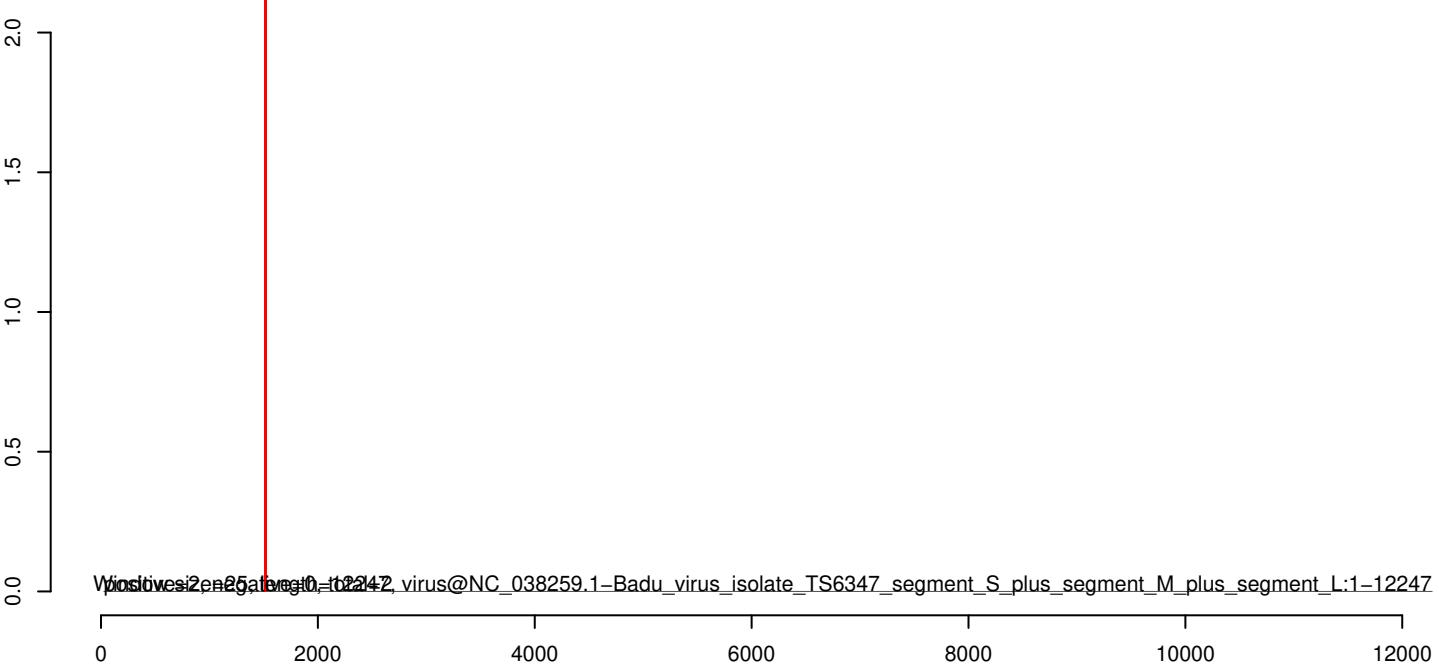
AeAeg_Aag2SINV_GFP.IP.18_23.rep



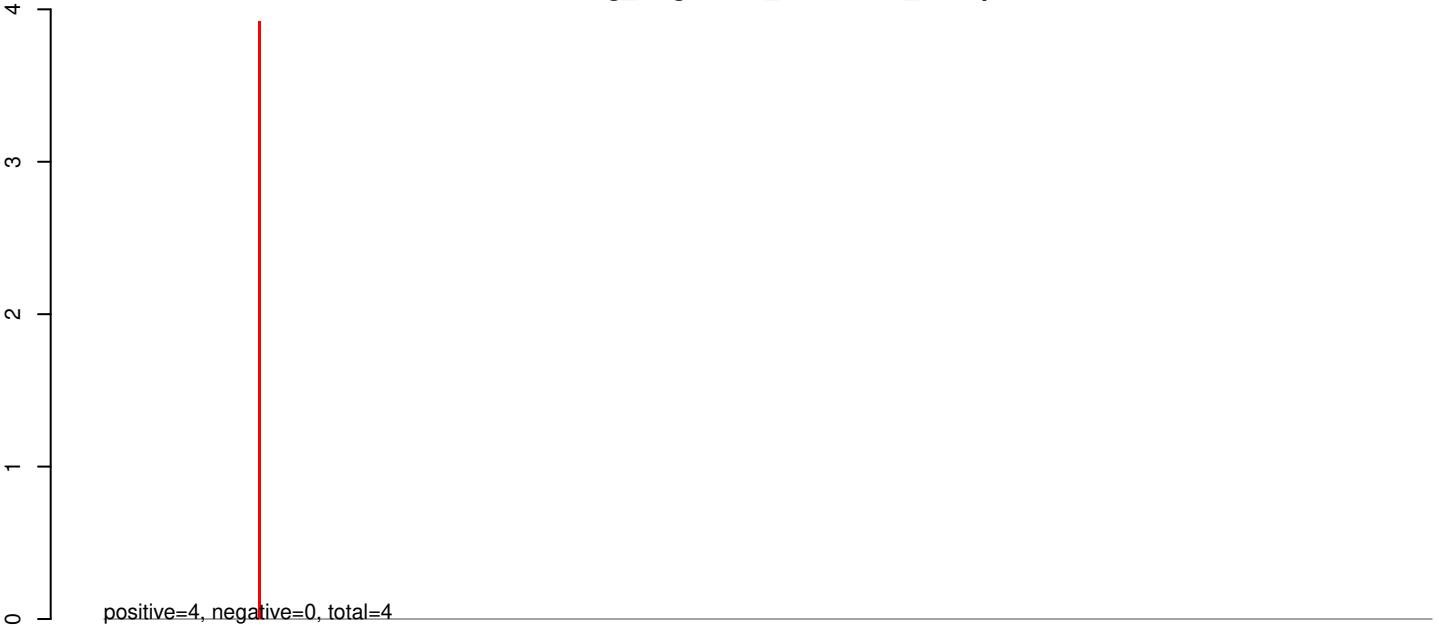
AeAeg_Aag2SINV_GFP.IP.24_35.rep



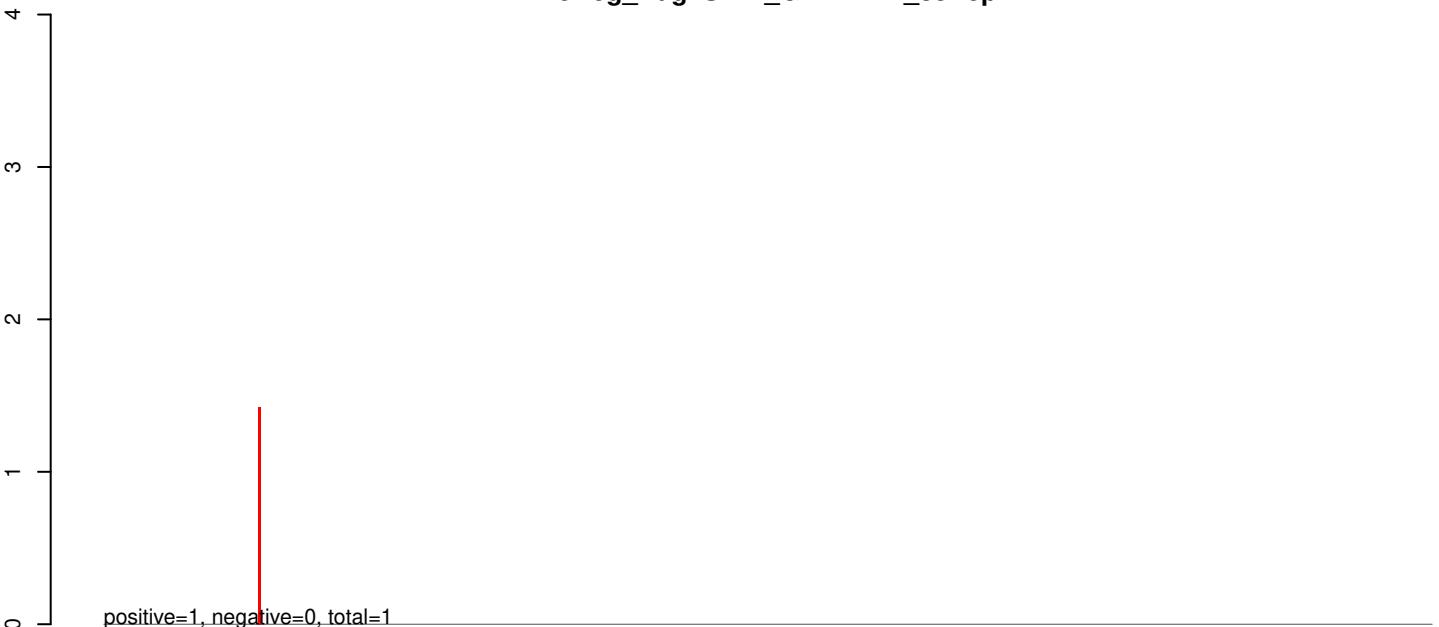
AeAeg_Aag2SINV_GFP.IP.rep



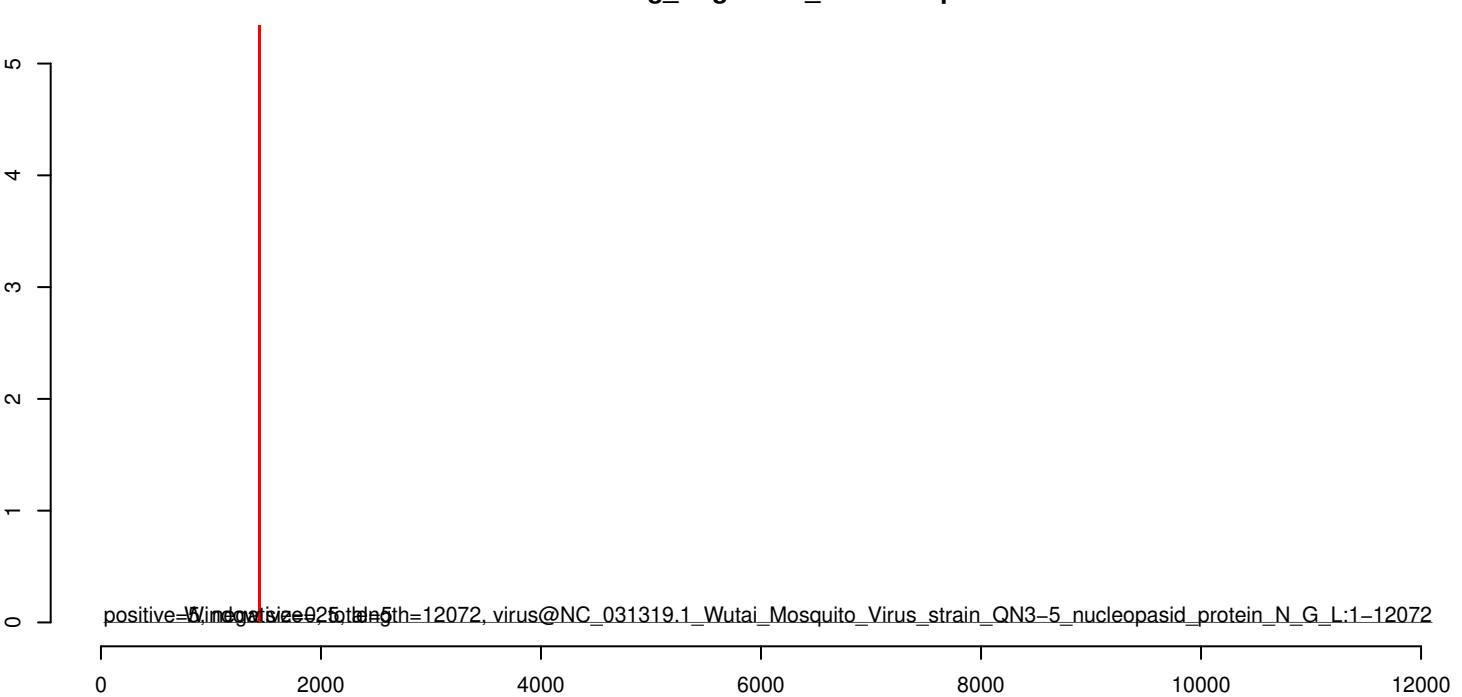
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=1, total=1

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=1, total=1

Window size=25, length=10158, virus@NC_030401.1-Hanko_virus_polyprotein_gene:1-10158

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

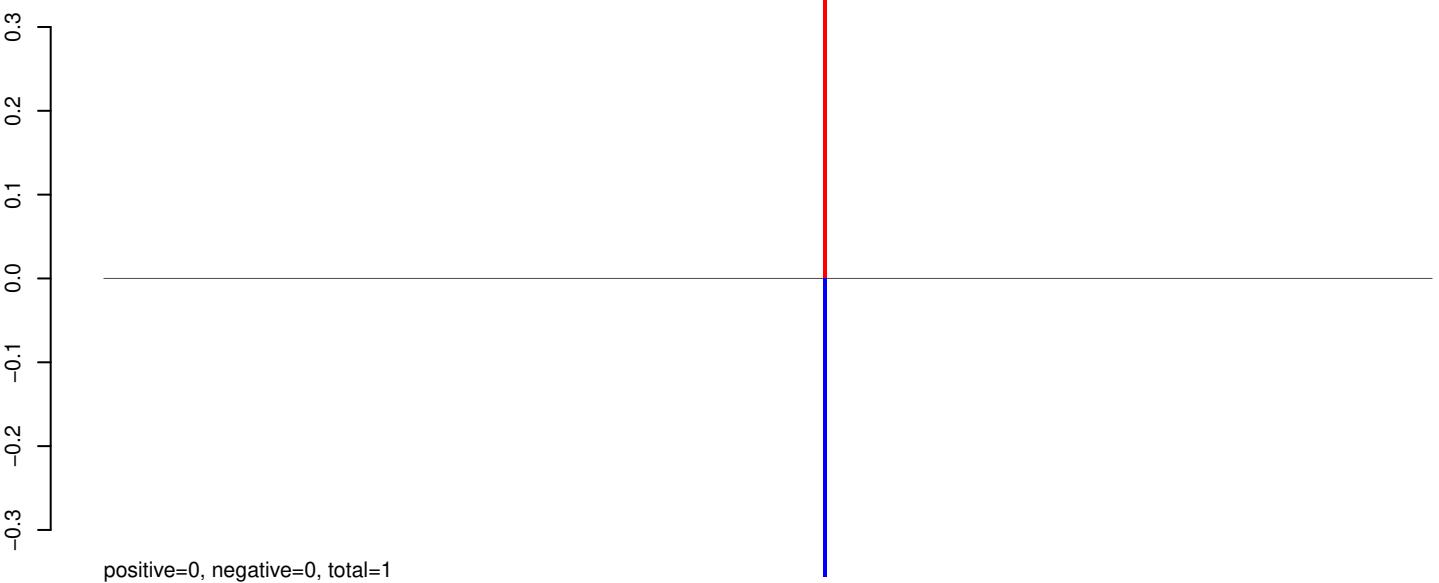
AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

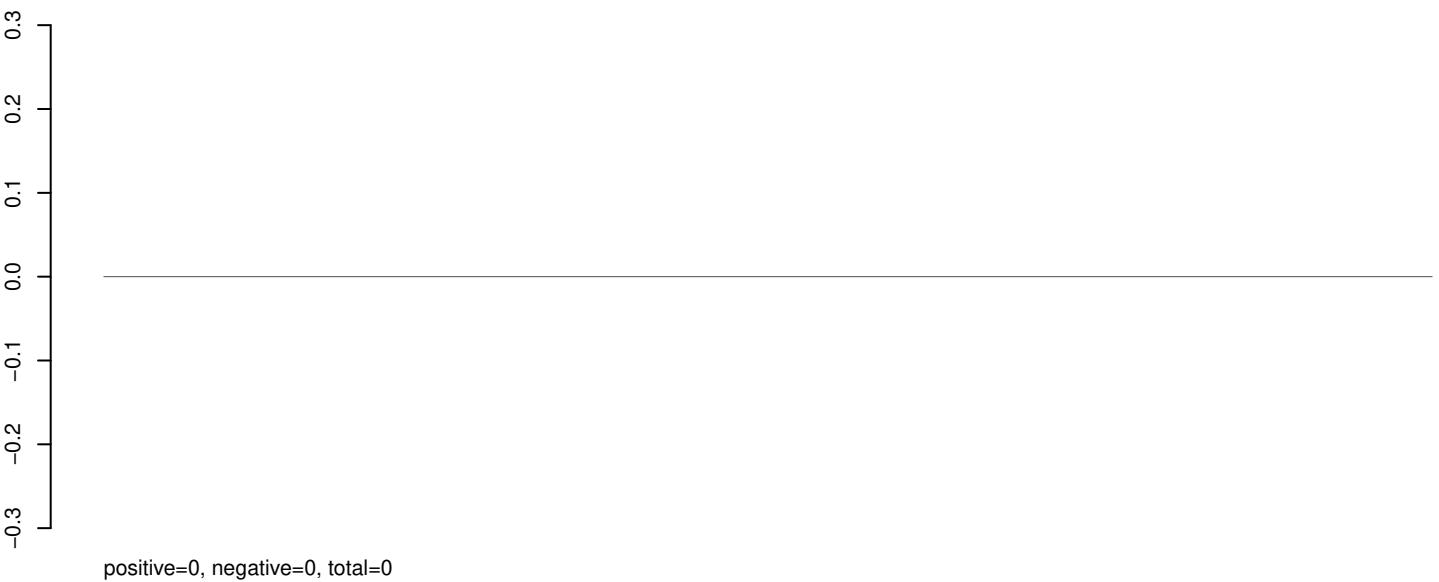
Window size=25, length=11550, virus@NC_024887.1-Middleburg_virus_isolate_ArB-8422:1-11550

0 2000 4000 6000 8000 10000 12000

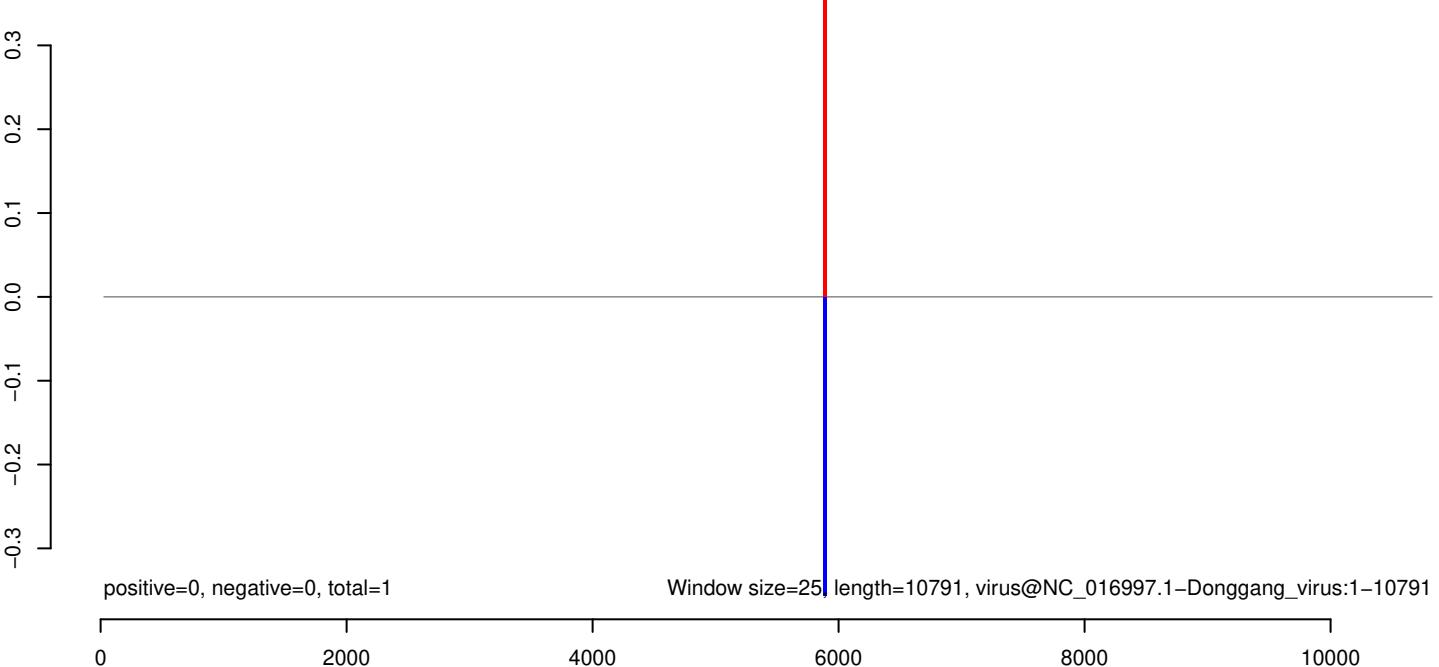
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10874, virus@NC_009029.2-Kokobera_virus:1-10874

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0



AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0



AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=11597, virus@NC_006558.1-Getah_virus:1-11597



0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=11826, virus@NC_004162.2-Chikungunya_virus:1-11826

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

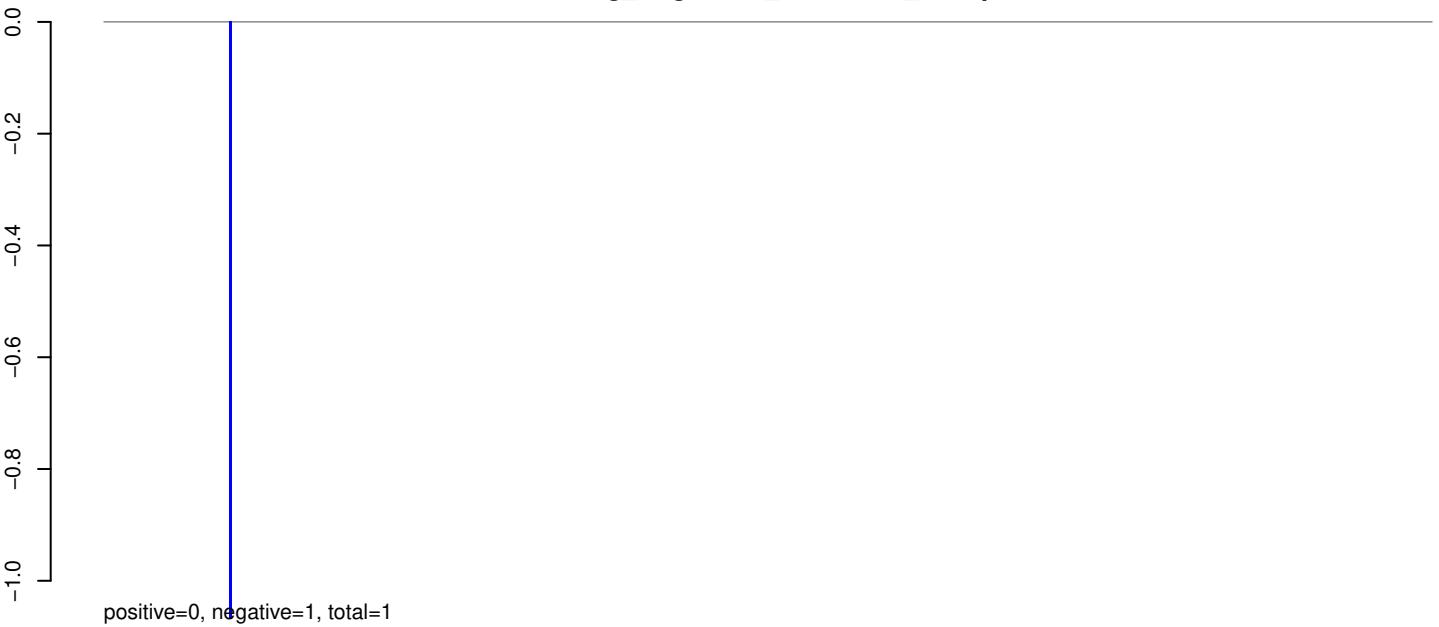
AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

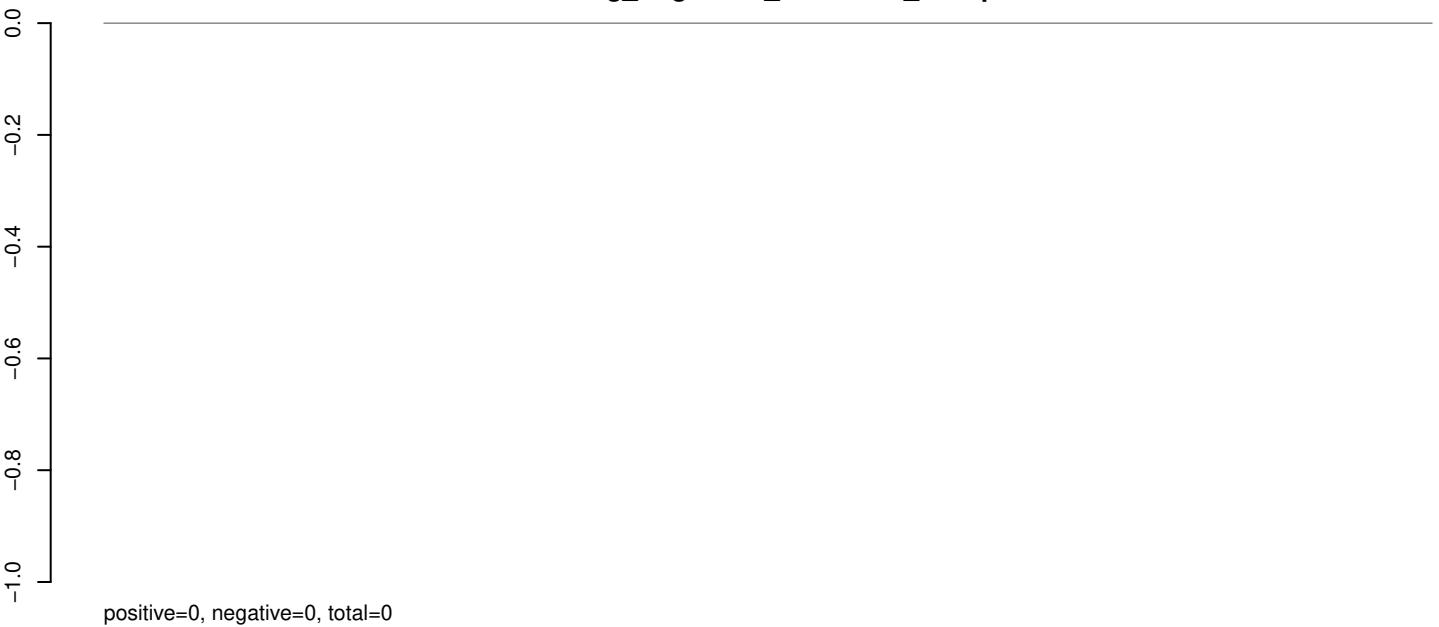
Window size=25, length=11824, virus@NC_003900.1-Aura_virus:1-11824

0 2000 4000 6000 8000 10000 12000

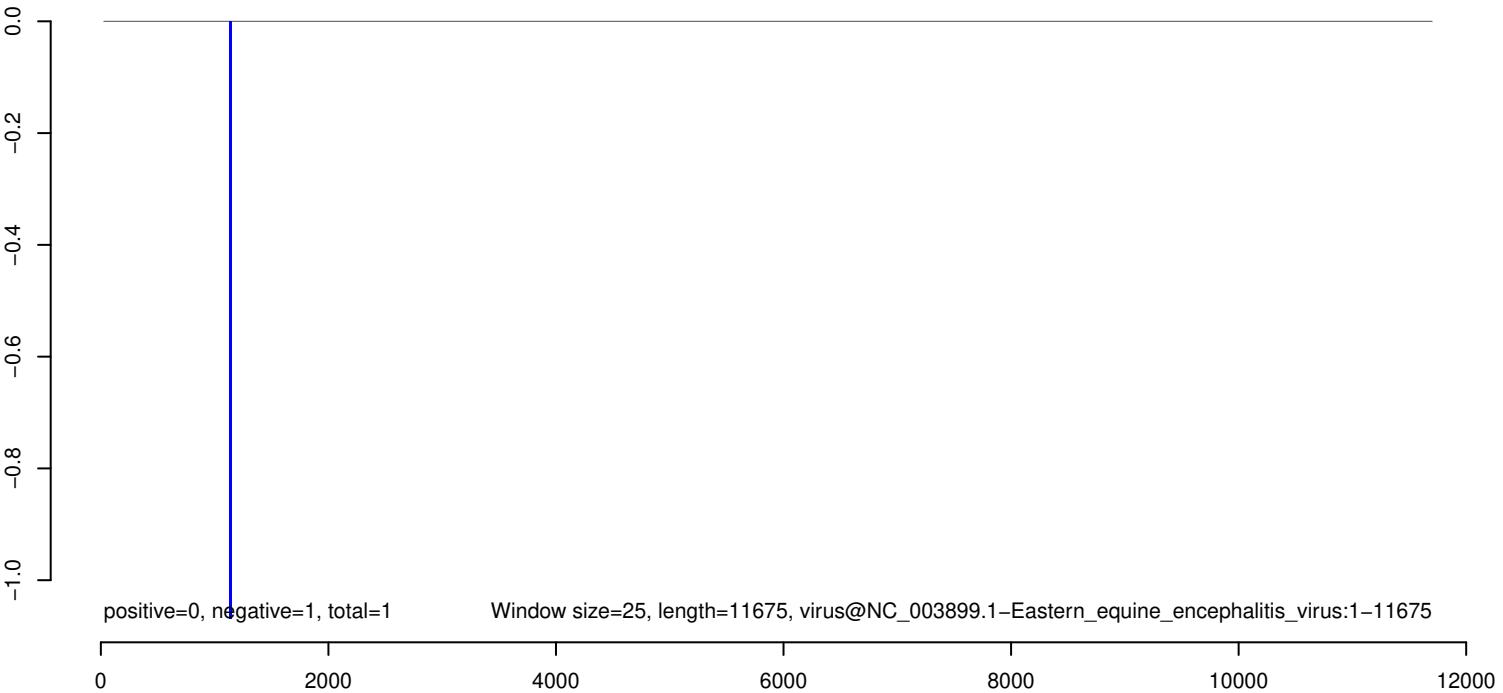
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

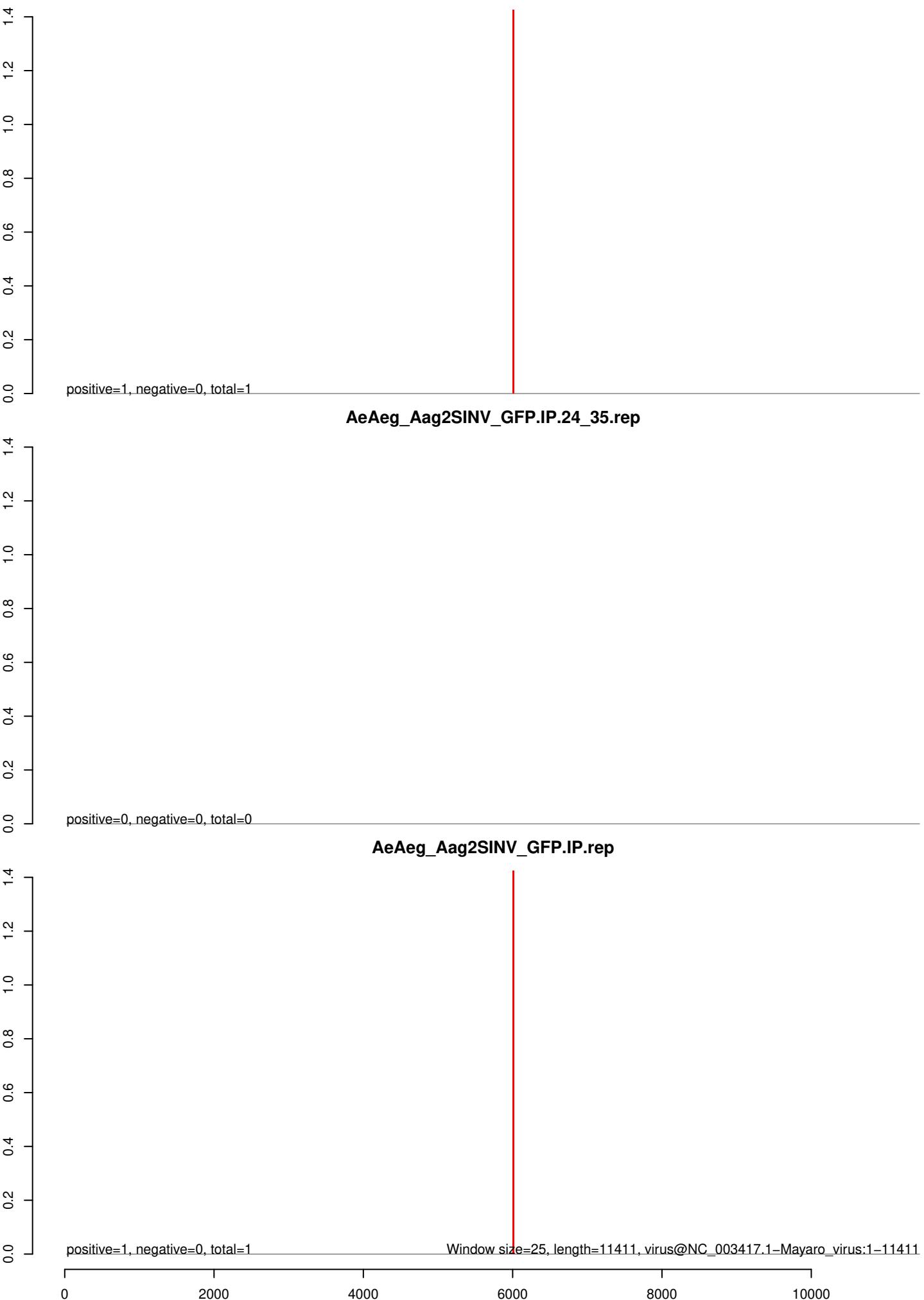
positive=1, negative=0, total=1

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=1, negative=0, total=1

Window size=25, length=11411, virus@NC_003417.1-Mayaro_virus:1-11411



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=1, negative=0, total=1



AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0



AeAeg_Aag2SINV_GFP.IP.rep

positive=1, negative=0, total=1, window size=25, length=11835, virus@M20303.1-O-Nyong-nyong_virus_Gulu_strain_complete_genome:1-11835



0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0 Window size=25, length=11657, virus@M20162.1–Ross_River_virus_RRV_strain_NB5092_RNA:1–11657

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

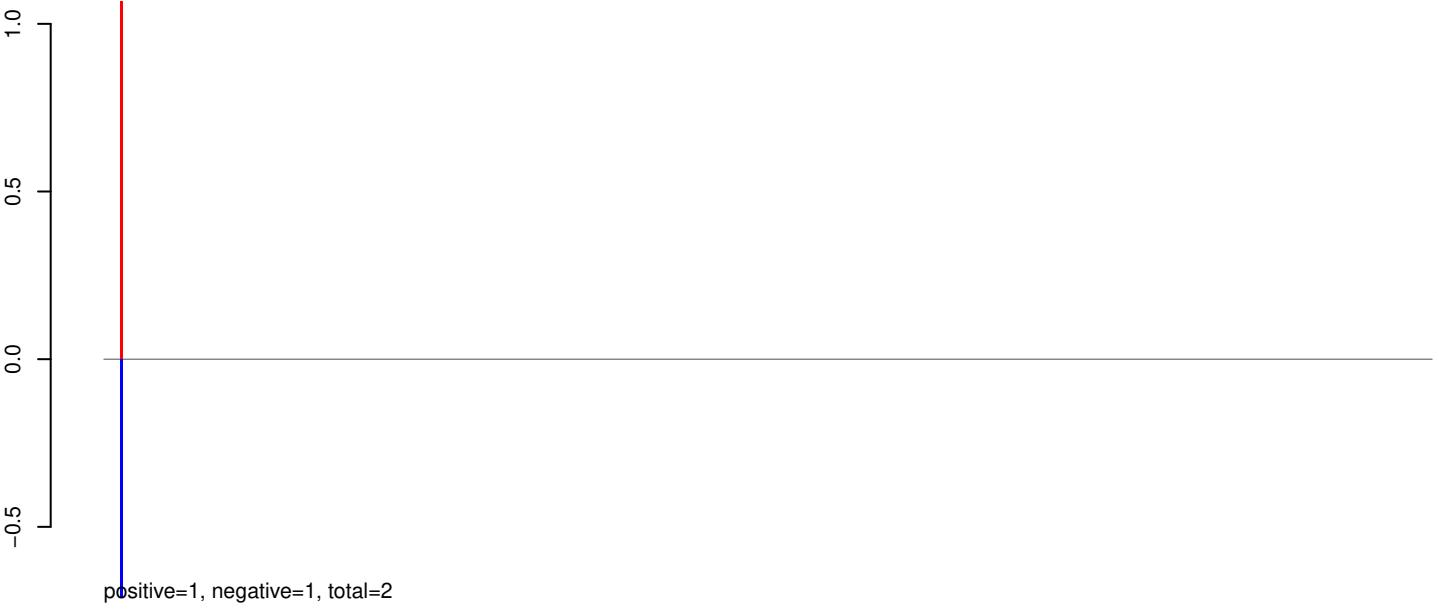
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

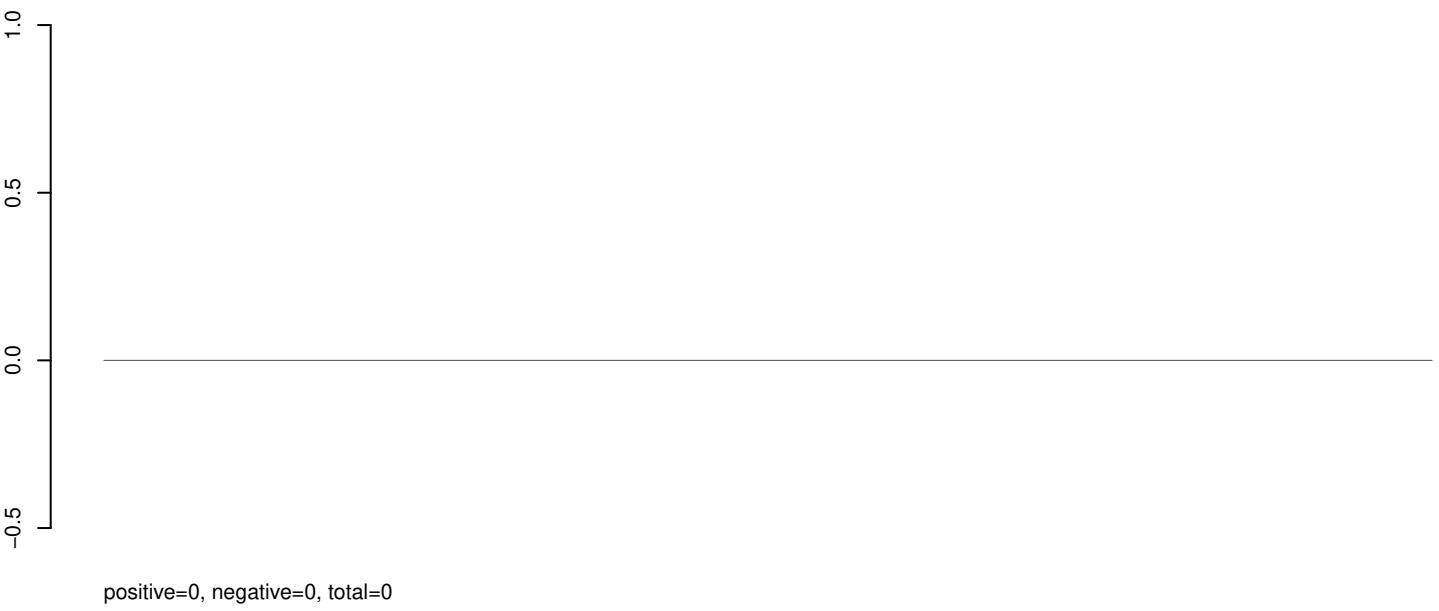
Wirus positive size 0.25 fold length = 0.12027 Virus@KC608151.1-llesha_virus_strain_ILESCHA_8e_segment_S_plus_segment_M_plus_segment_L:1-12297

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

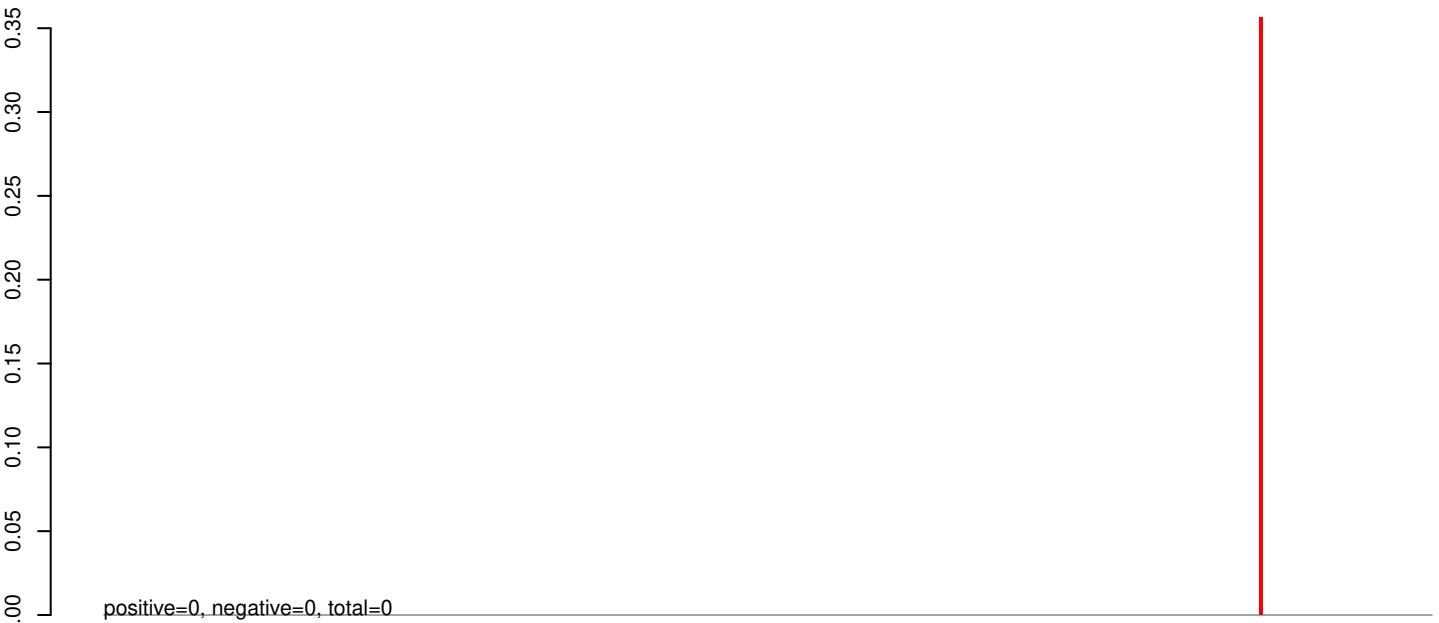
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

dow_spe125e_10_gnega12052, total@HM147991.1-Trocara_virus_non_structural_polyprotein_gene_partial_cds_and_structural_polyprotein:1-12052

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=1, negative=0, total=1

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

size=10587 length=10587 version=0 user=AEG111835.2-Aedes_albopictus_cell_fusing_agent_virus_polyprotein-like_protein_gene_and_agCP8252:1-10587

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=11385, virus@AF075259.1-Cabassou_virus_strain_CaAr_508:1-11385

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=11465, virus@AF075257.1-Mosso_das_Pedras_virus:1-11465

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

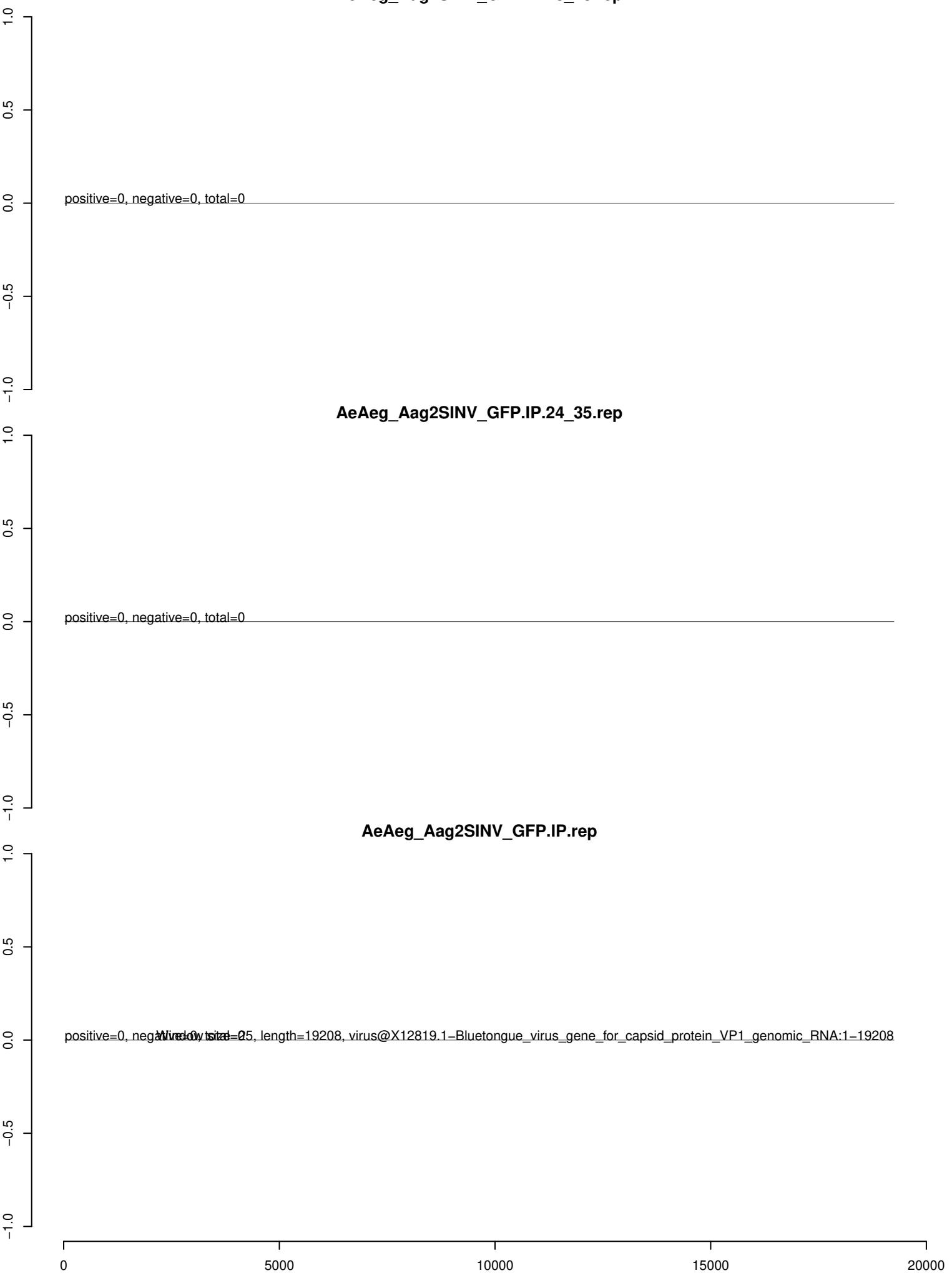
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, virus@X12819.1-Bluetongue_virus_gene_for_capsid_protein_VP1_genomic_RNA:1-19208



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

window positive 25,0 negative 1202, total 1227 @ NC_038712.1-Herbert_virus_strain_F23_CI_2004_segment_S_plus_segment_M_plus_segment_L:1-11202

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

window positive=0, negative=14098, total=0 NC_038707.1-Jonchet_virus_strain_B81-CI-2004_segment_S_plus_segment_M_plus_segment_L:1-14098

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, window_size=25, length=11503, virus@NC_035132.1_Culex_rhabdo-like_virus_strain_mosWSB71420:1-11503

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=250 length=6363, virus@NC_035130.1_Aedes_alboannulatus_toti-like_virus_1_strain_mosWSCP114121:1-6363

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, Window size=25, length=9176, virus@NC_035129.1_Culex_negev-like_virus_3_strain_mos172X44875:1-9176

0 2000 4000 6000 8000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=925, total=11470, virus@NC_035128.1_Aedes_camptorhynchus_negev-like_virus_strain_mos191CP47896:1–11470

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

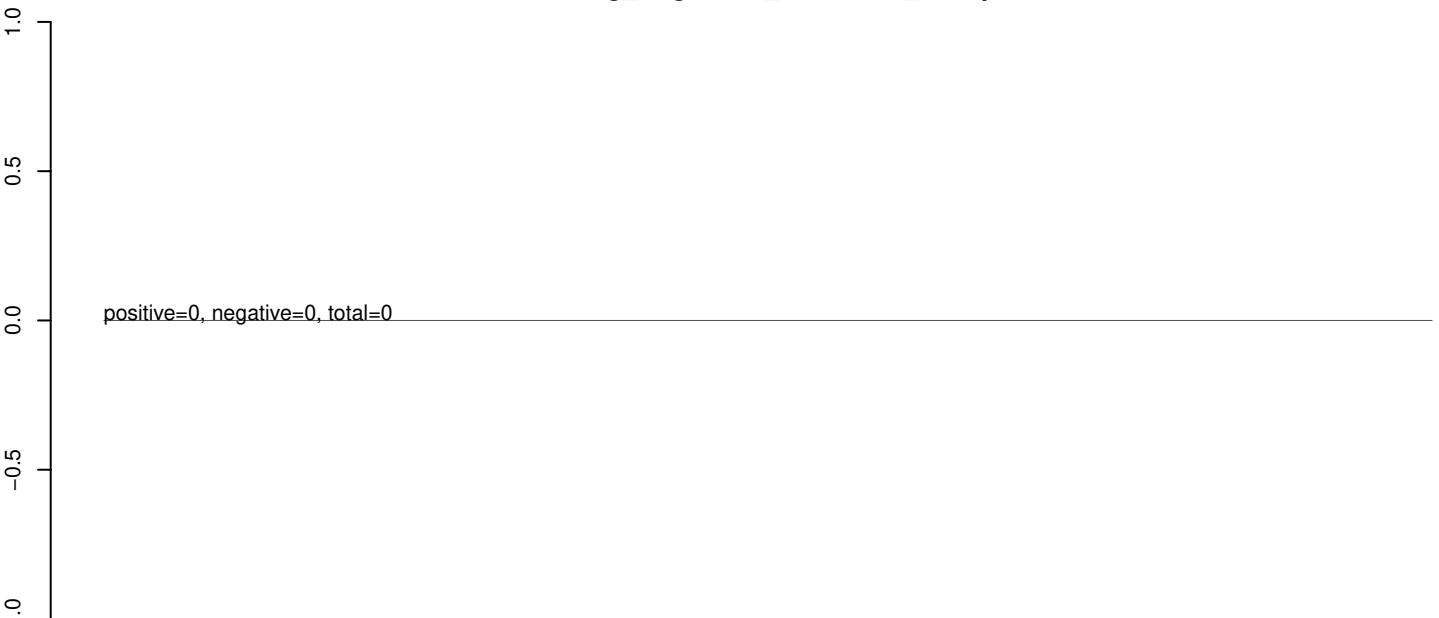
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

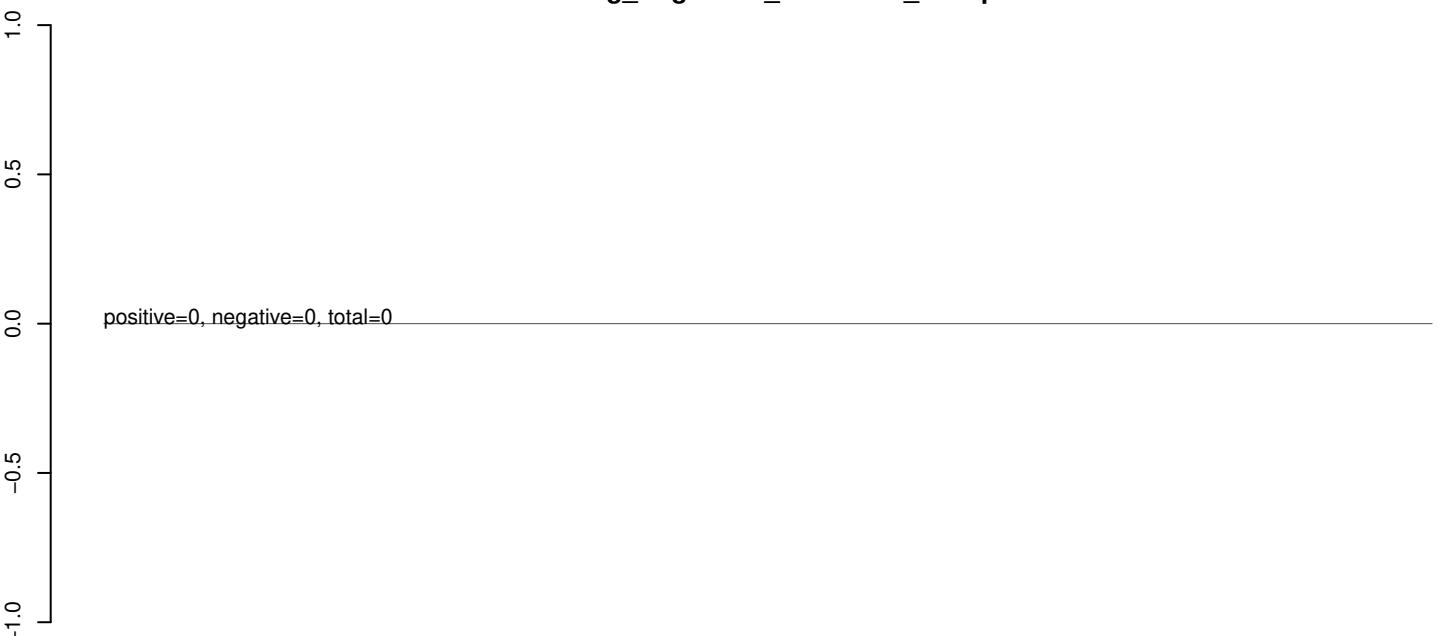
AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, window.size=25, length=2968, virus@NC_035126.1_Wilkie_narna-like_virus_1_strain_mosWSCP70929:1-2968

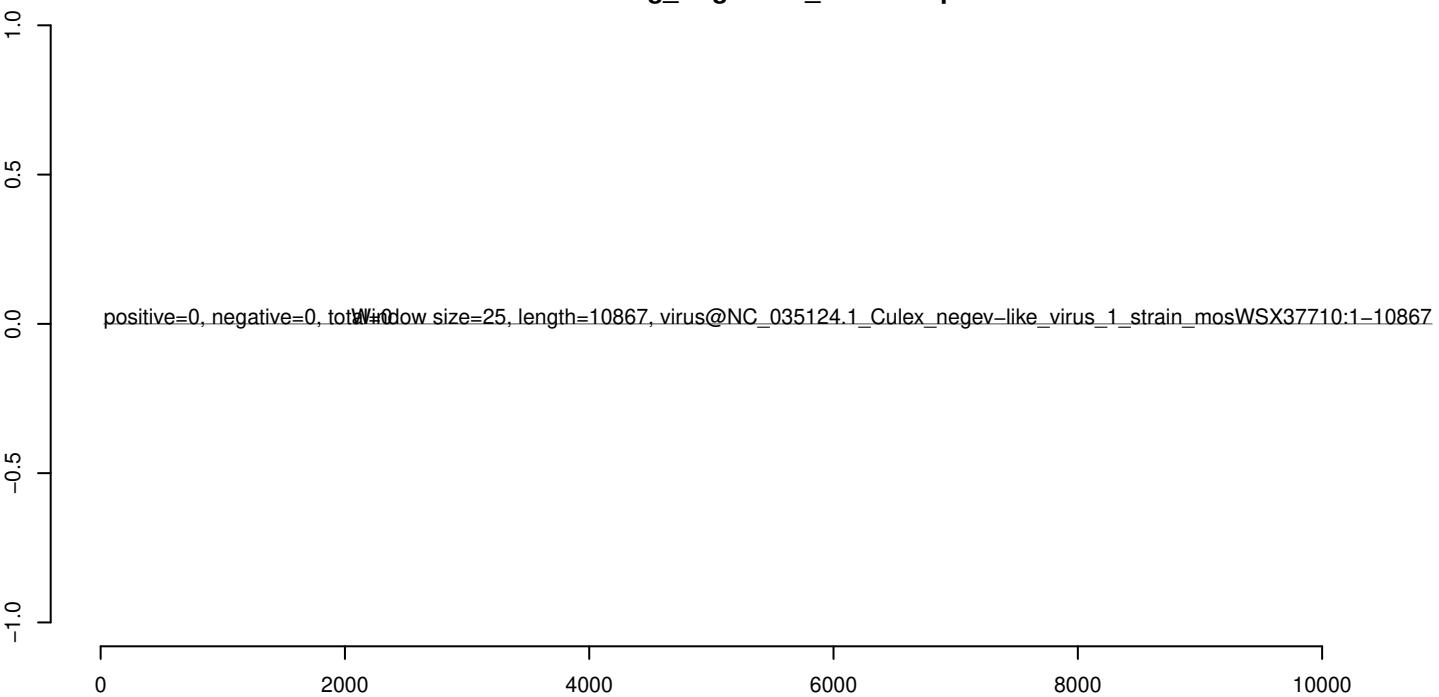
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, Window size=25, length=9324, virus@NC_035123.1_Culex_negev-like_virus_2_strain_mos172X30622:1-9324

0 2000 4000 6000 8000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, Window size=25, length=2857, virus@NC_035121.1_Point-Douro_narna-like_virus_strain_mos172gb29666:1-2857

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, window.size=25, length=3367, virus@NC_035120.1_Wilkie_narna-like_virus_2_strain_mosWSCP85442:1-3367

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

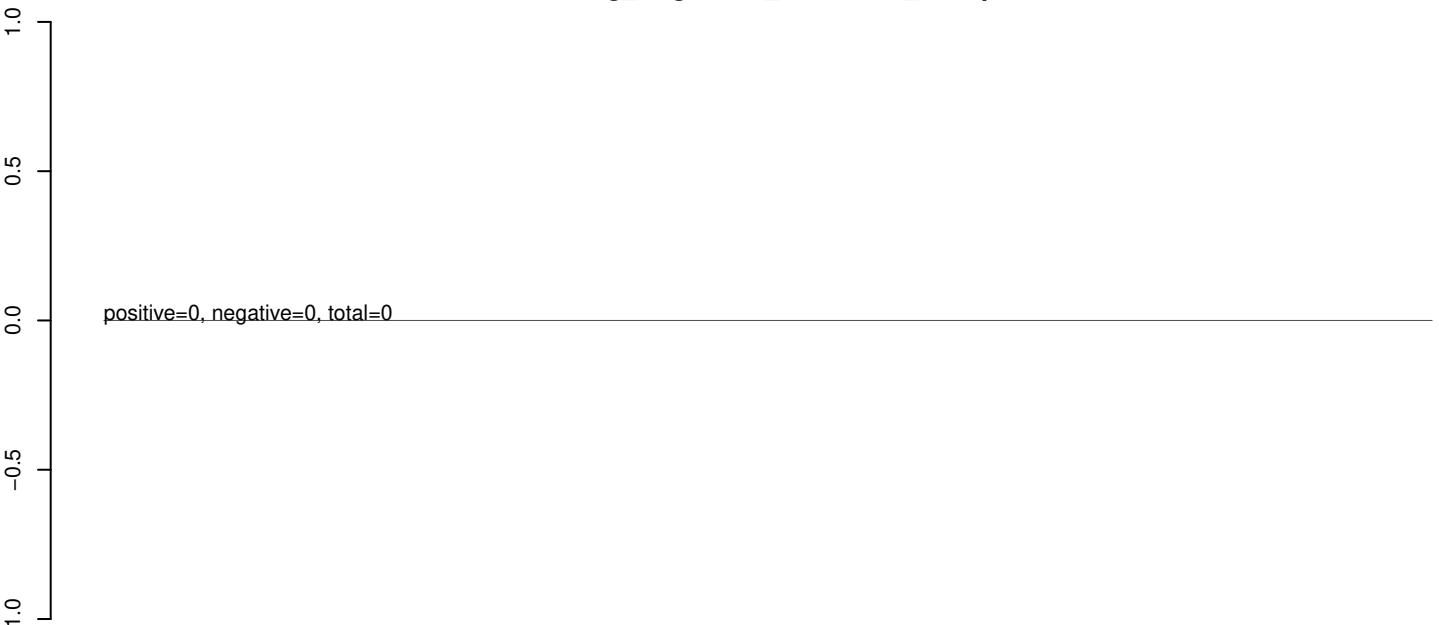
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

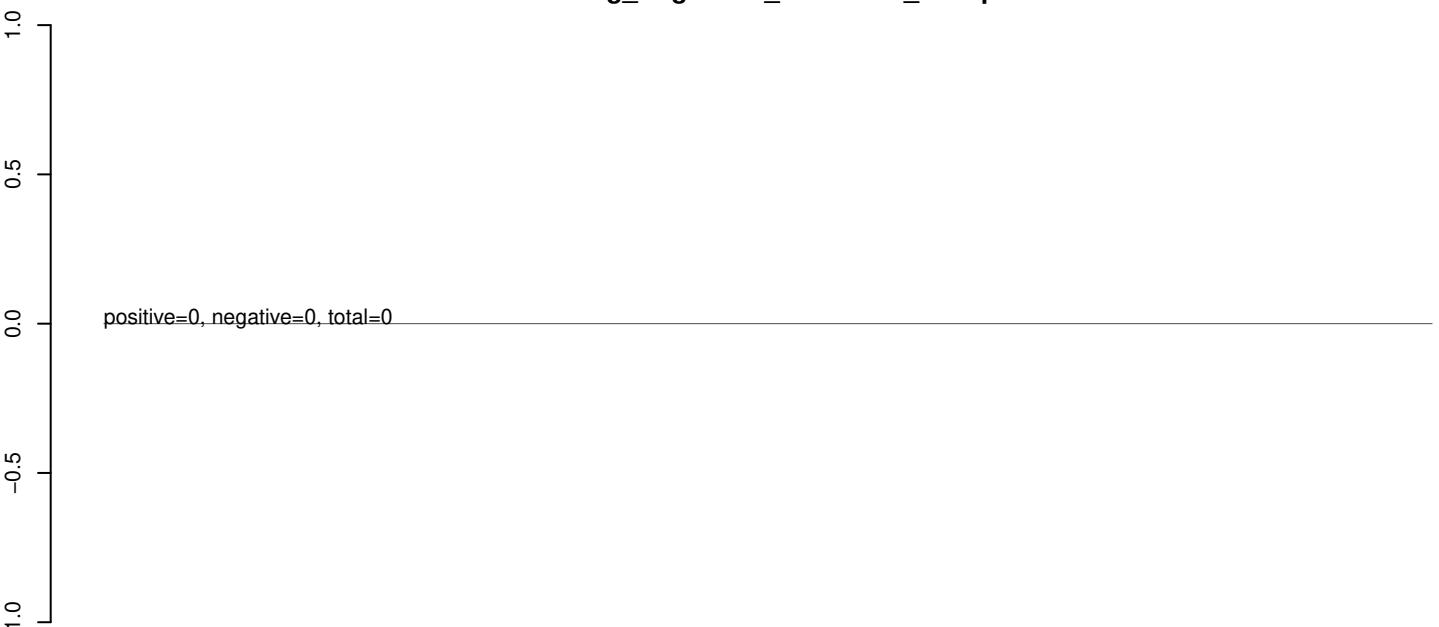
low size:25 type:0 theta:189 virus:NC_034543.1-Ord_River_virus_nucleoprotein_hypothetical_protein_phosphoprotein_hypothetical_pro:1-13189

0 2000 4000 6000 8000 10000 12000

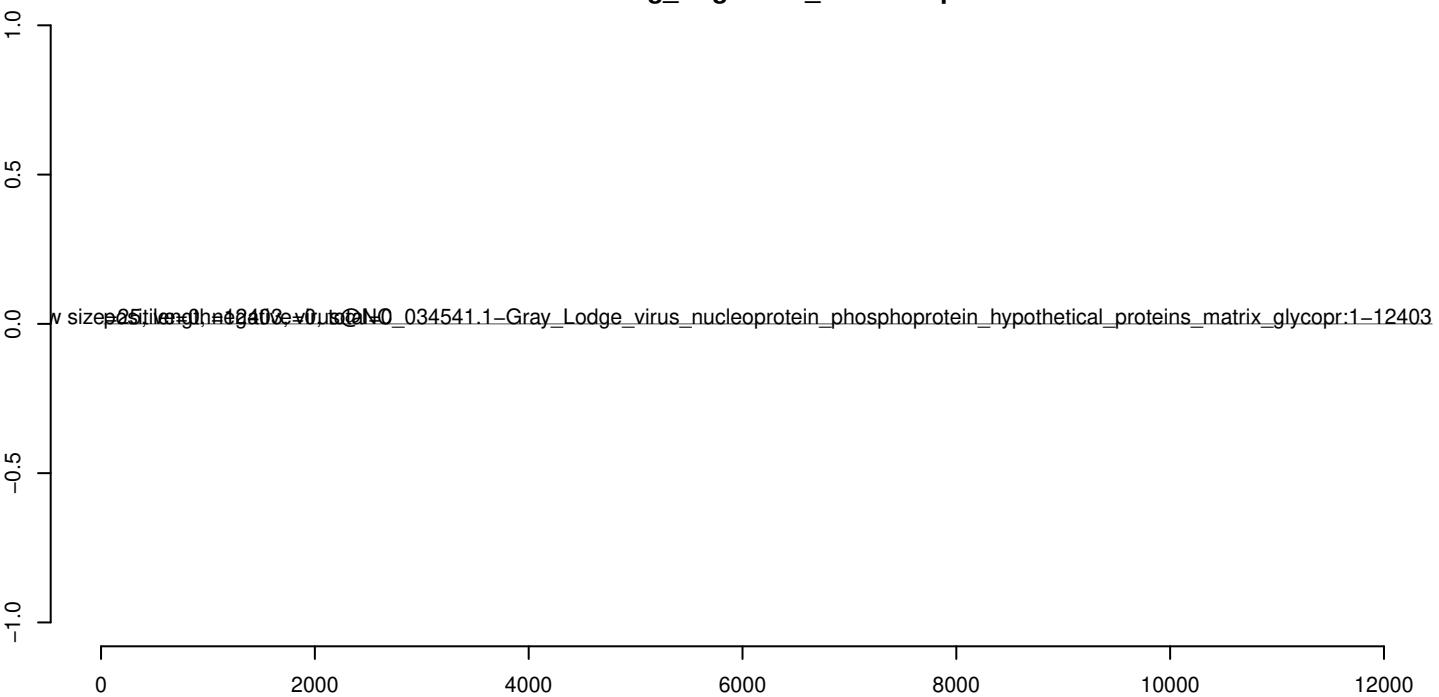
AeAeg_Aag2SINV_GFP.IP.18_23.rep



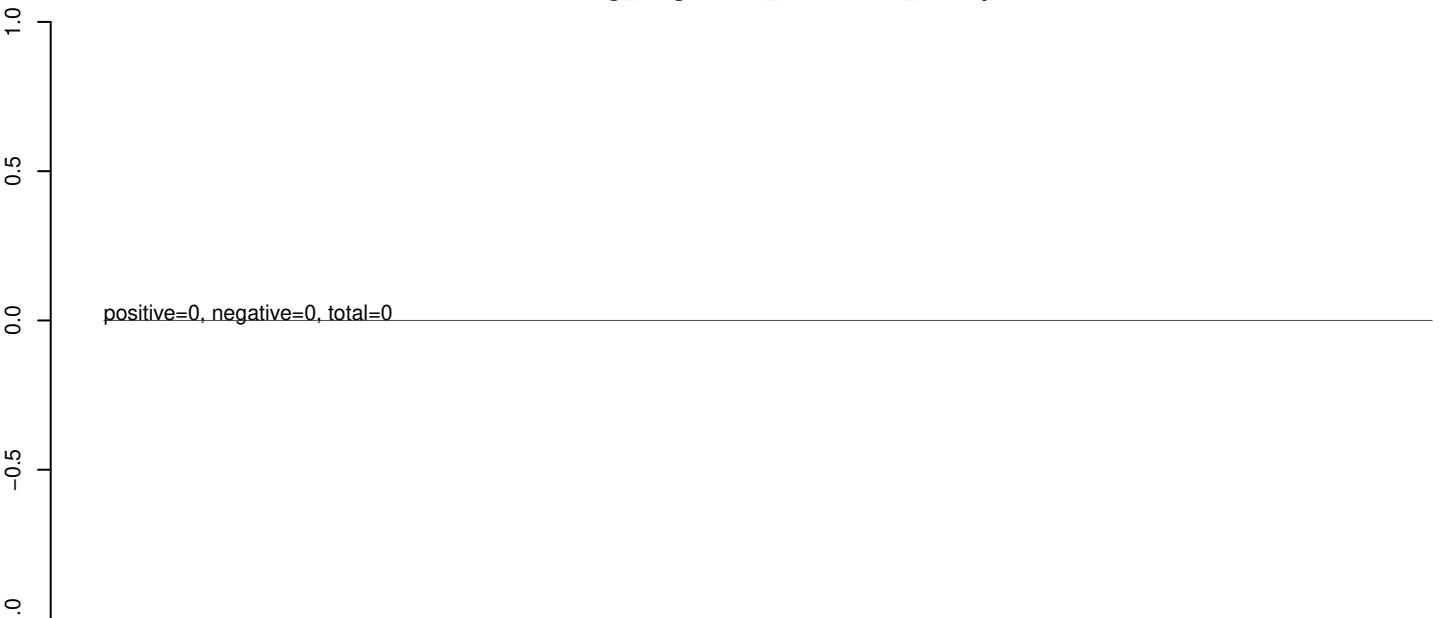
AeAeg_Aag2SINV_GFP.IP.24_35.rep



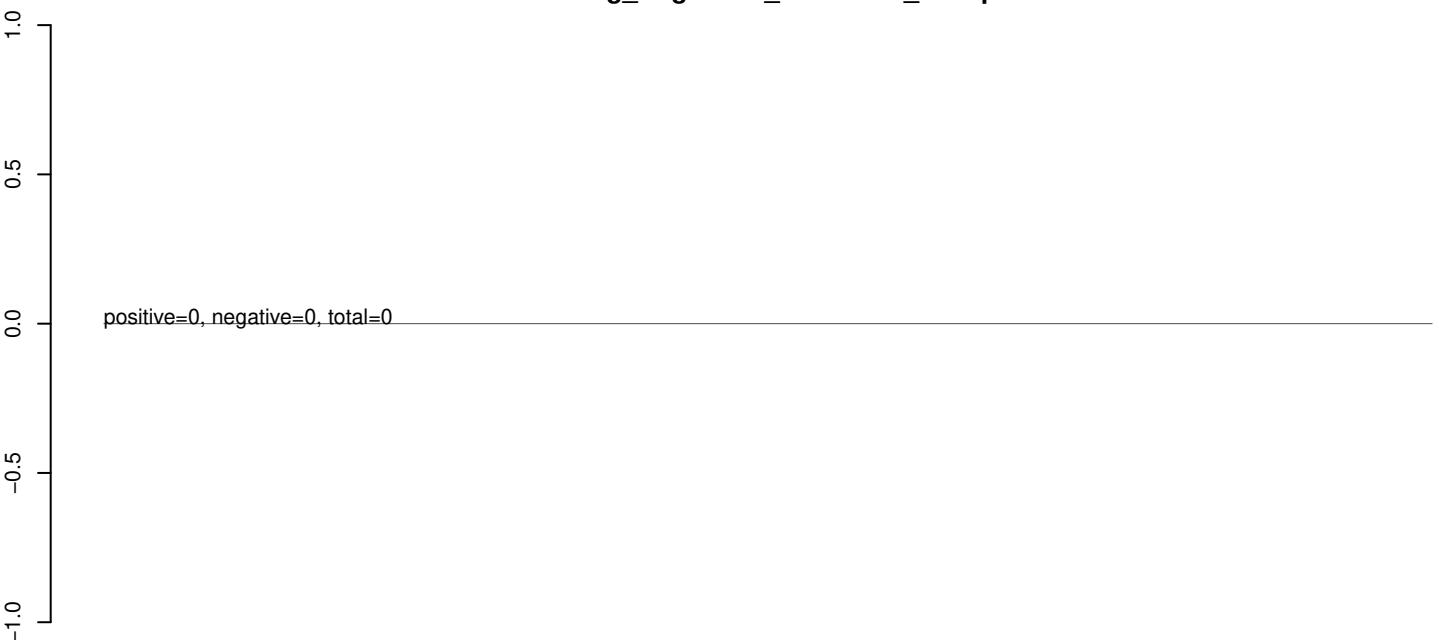
AeAeg_Aag2SINV_GFP.IP.rep



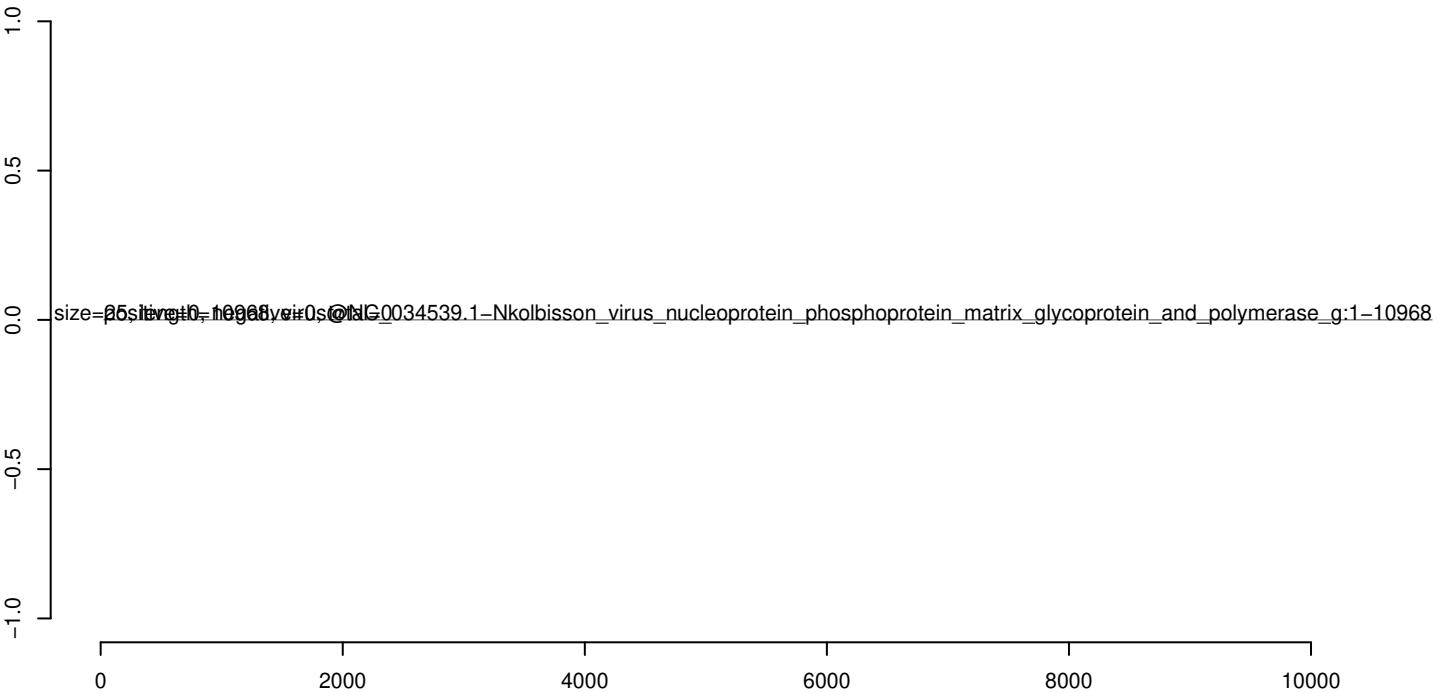
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

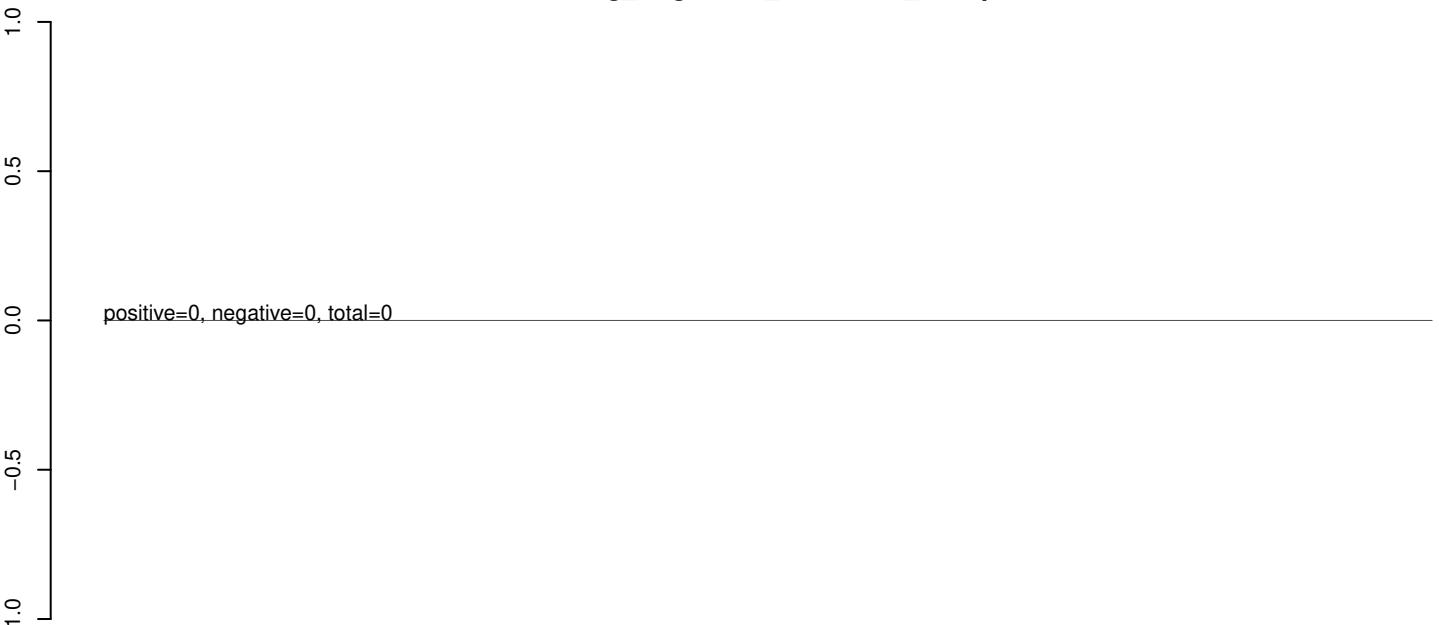
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

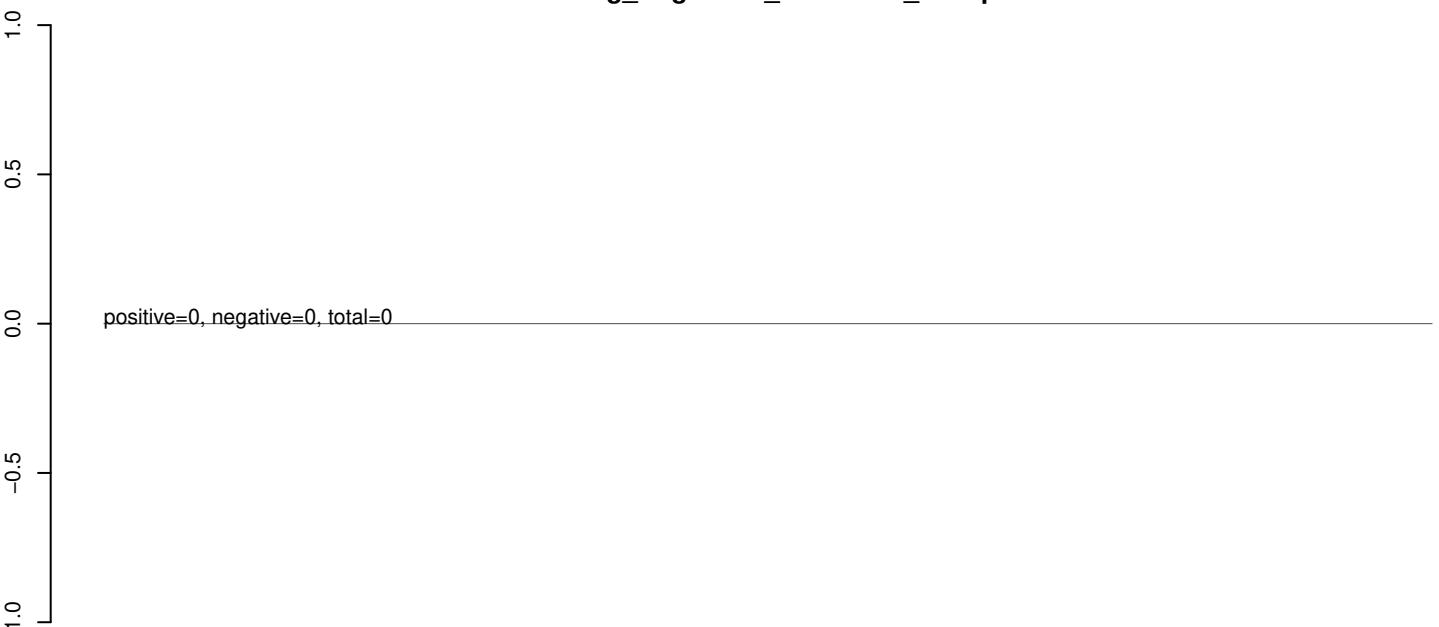
AeAeg_Aag2SINV_GFP.IP.rep

low size:25 deg:0 theta:2155 0 total@NC_034538.1-Joinjakaka_virus_nucleoprotein_phosphoprotein_hypothetical_protein_matrix_glycopro:1-13155

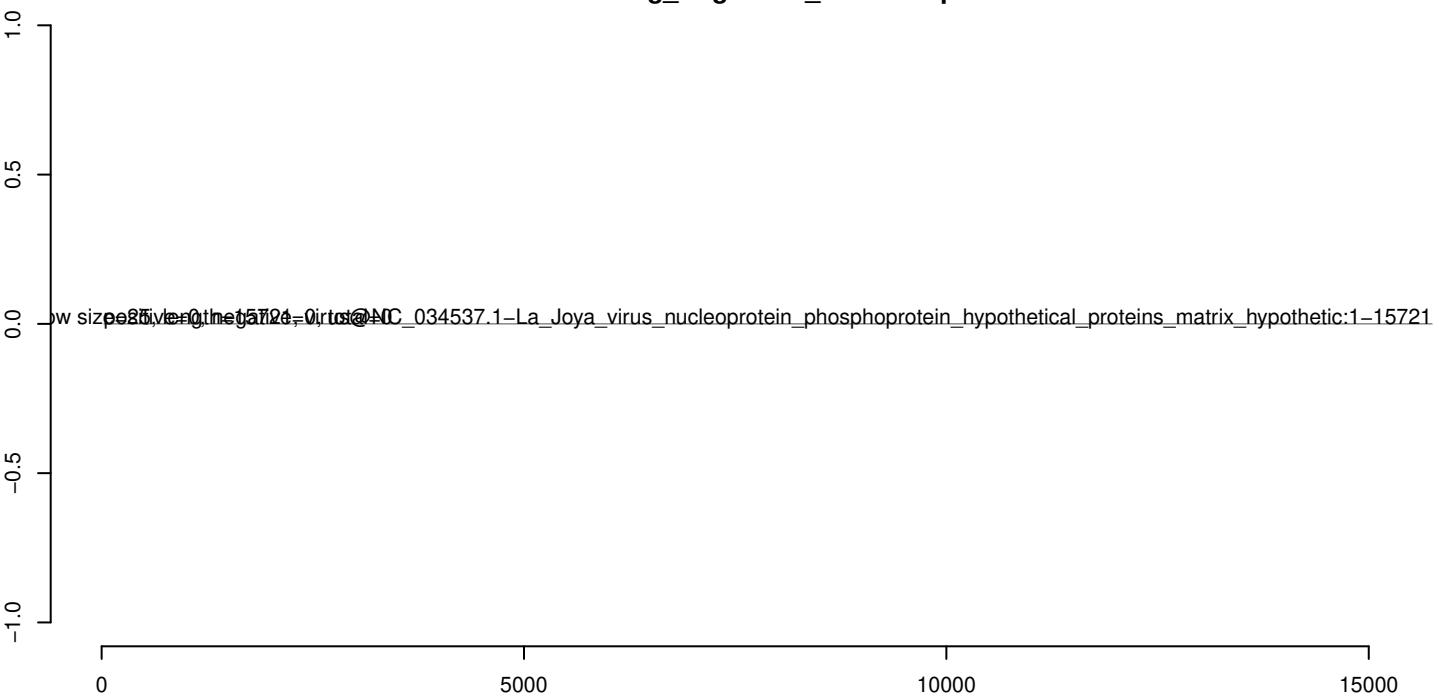
AeAeg_Aag2SINV_GFP.IP.18_23.rep



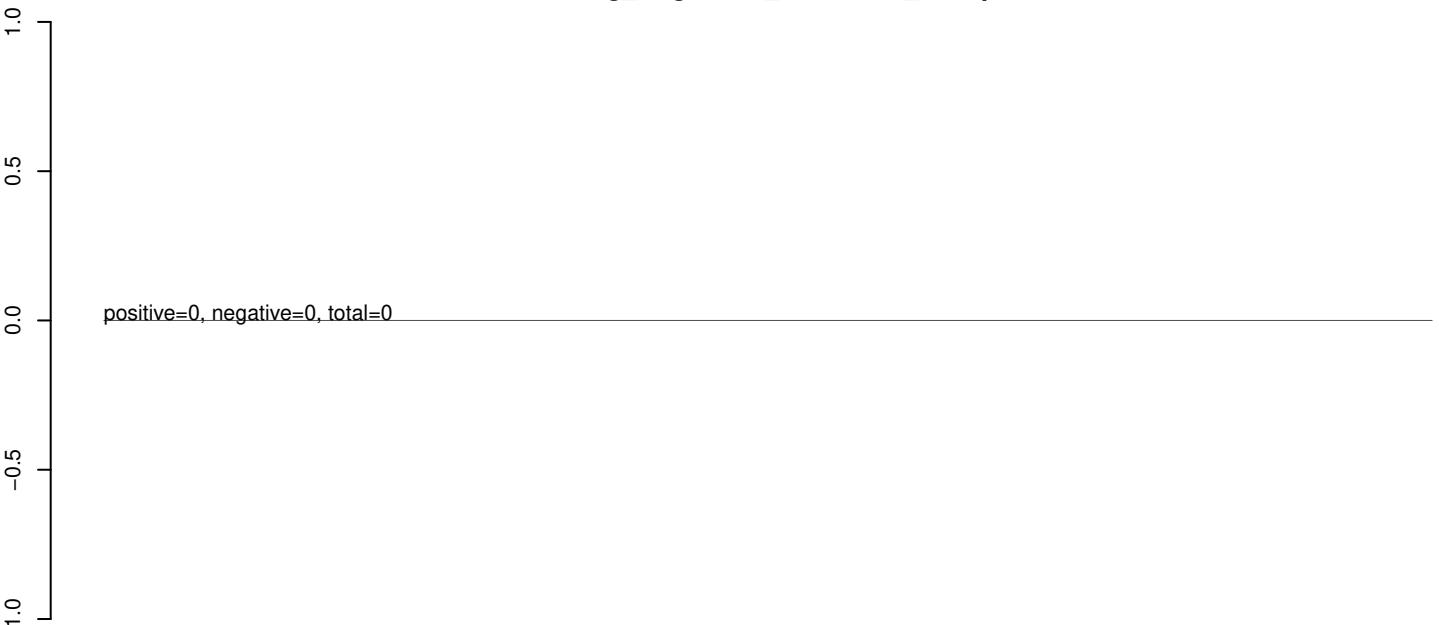
AeAeg_Aag2SINV_GFP.IP.24_35.rep



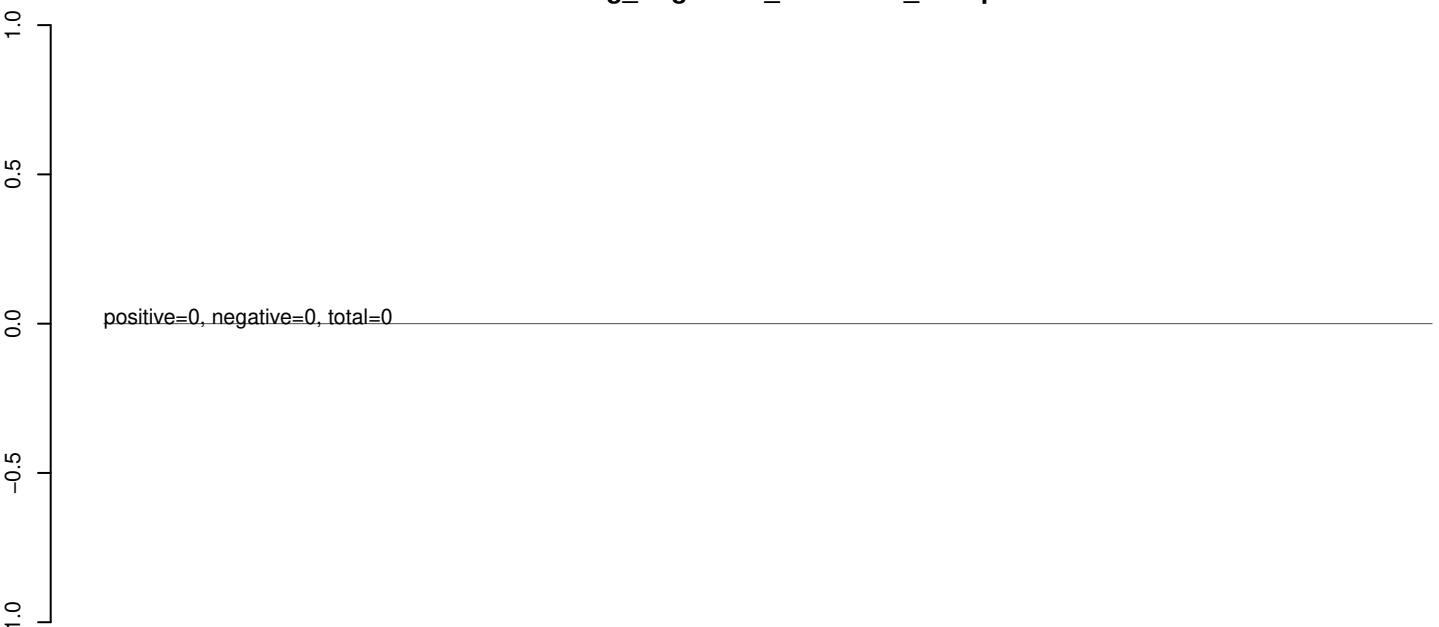
AeAeg_Aag2SINV_GFP.IP.rep



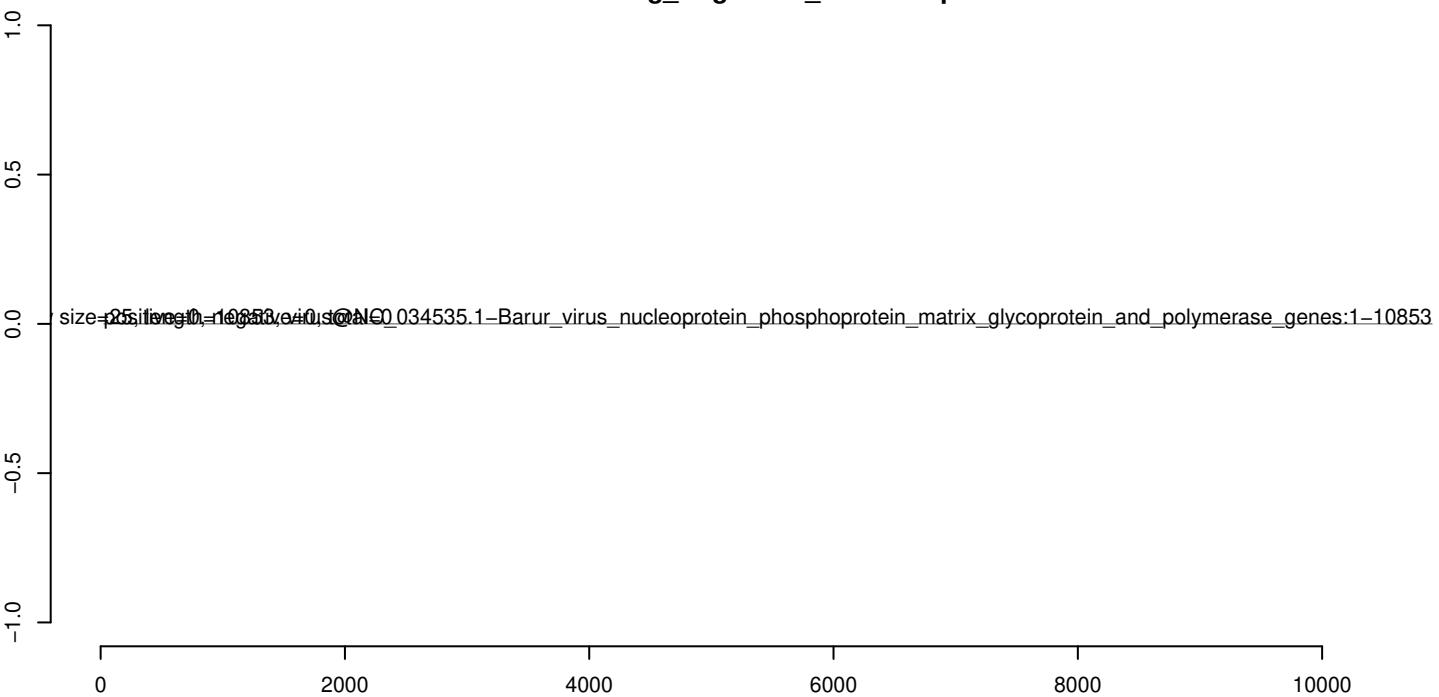
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

size=25 items=0, 16580 views=0, total=0 034534.1_Rochambeau_virus_nucleoprotein_phosphoprotein_matrix_hypothetical_proteins_glycopr:1-13593

0 2000 4000 6000 8000 10000 12000 14000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

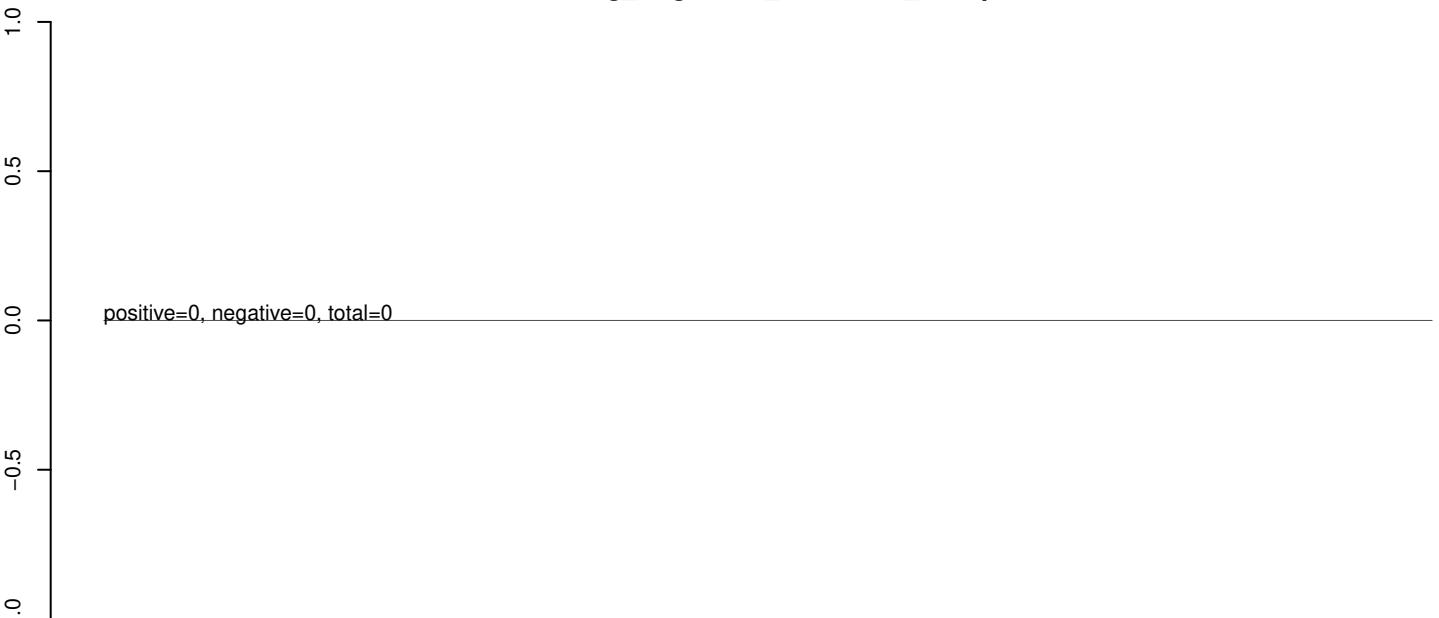
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

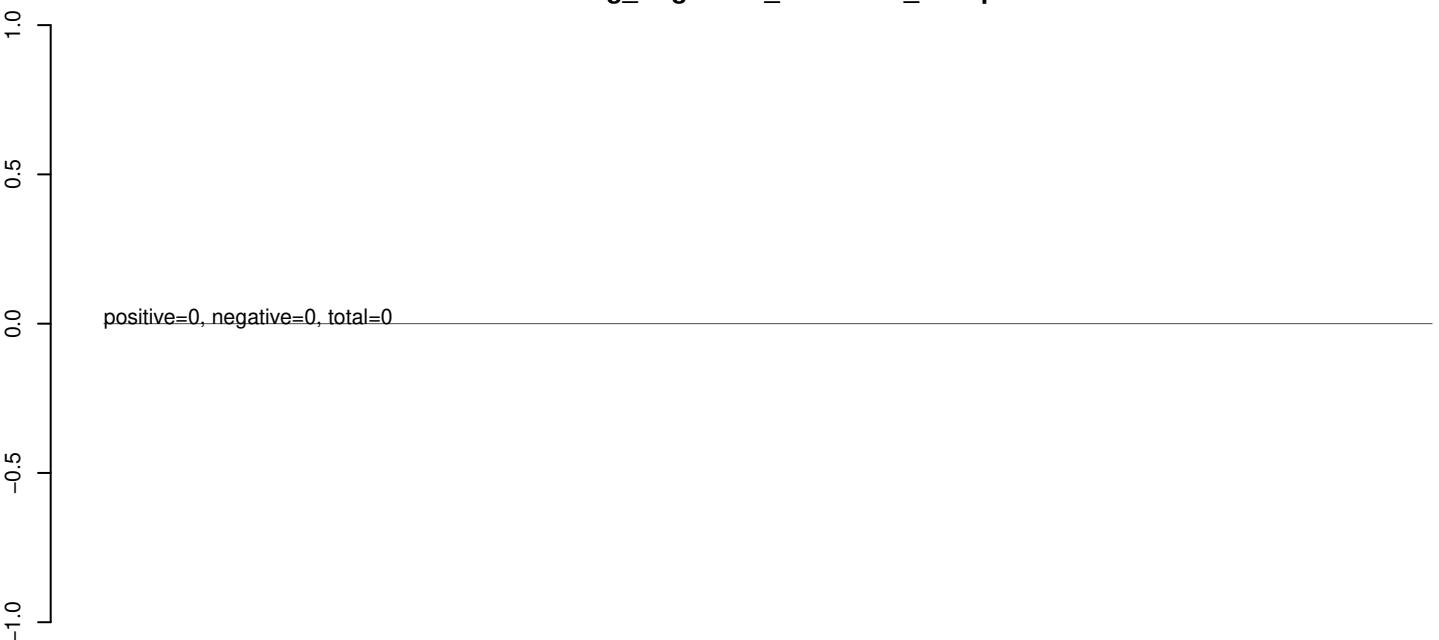
low size peptide 25 kDa or less 1704 0 viral NC_034531.1-Manitoba_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycoprot:1-13784

0 2000 4000 6000 8000 10000 12000 14000

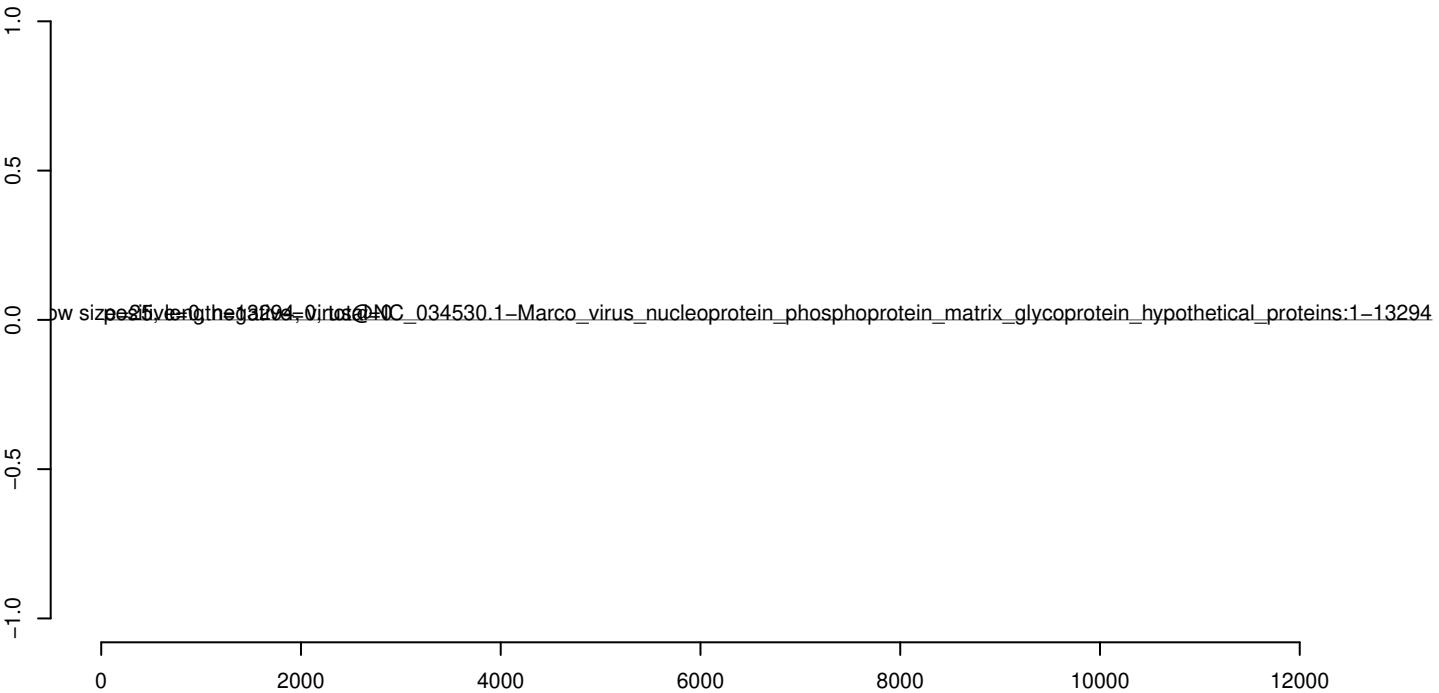
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

length=11322, negative=0, positive=0, total=0
5, length=11322, negative=0, positive=0, total=0
Kibale_virus_strain_P05_UG_2008_segment_S_plus_segment_M_plus_segment_L_nucleocaps:1-11322

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

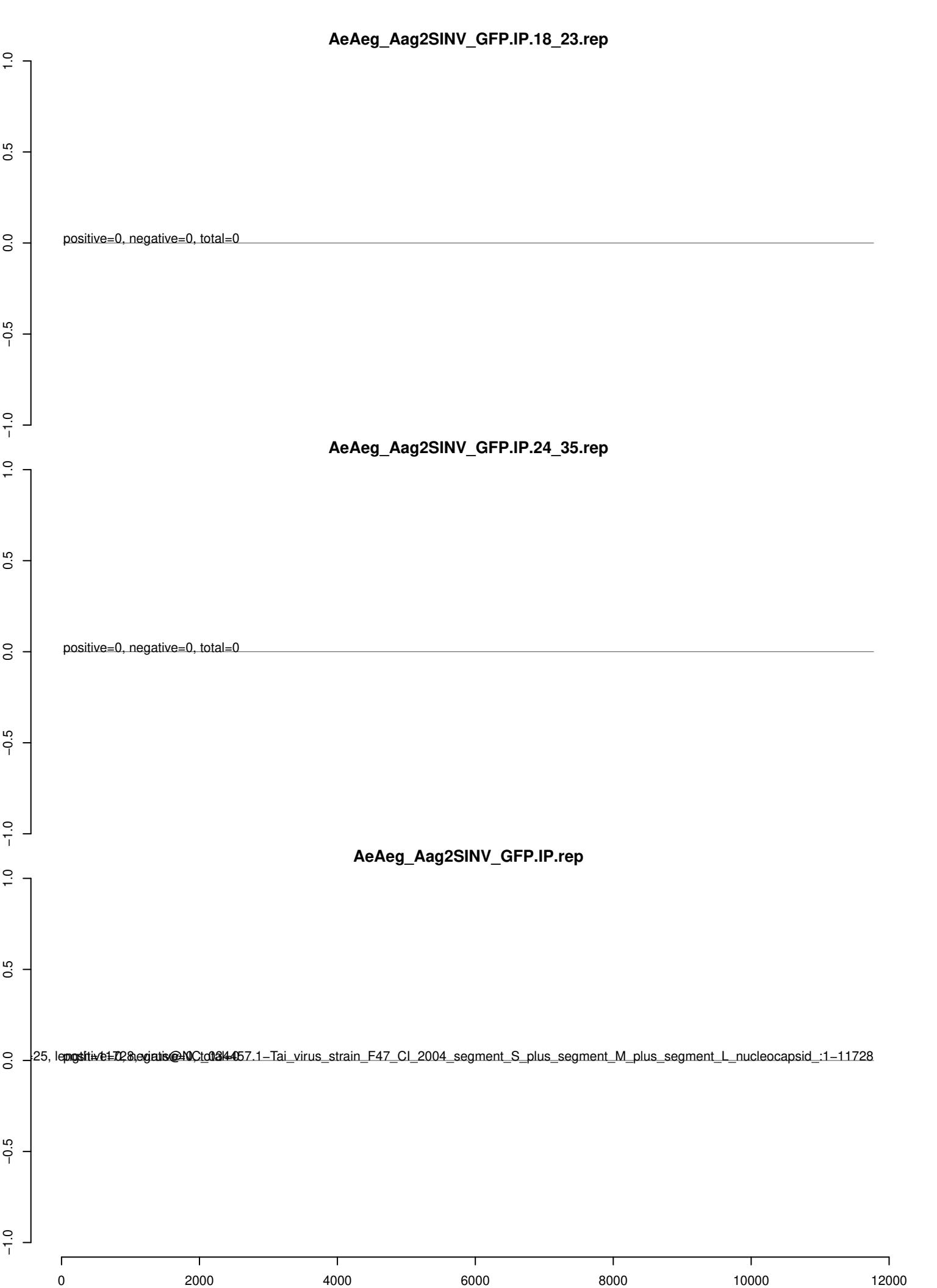
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

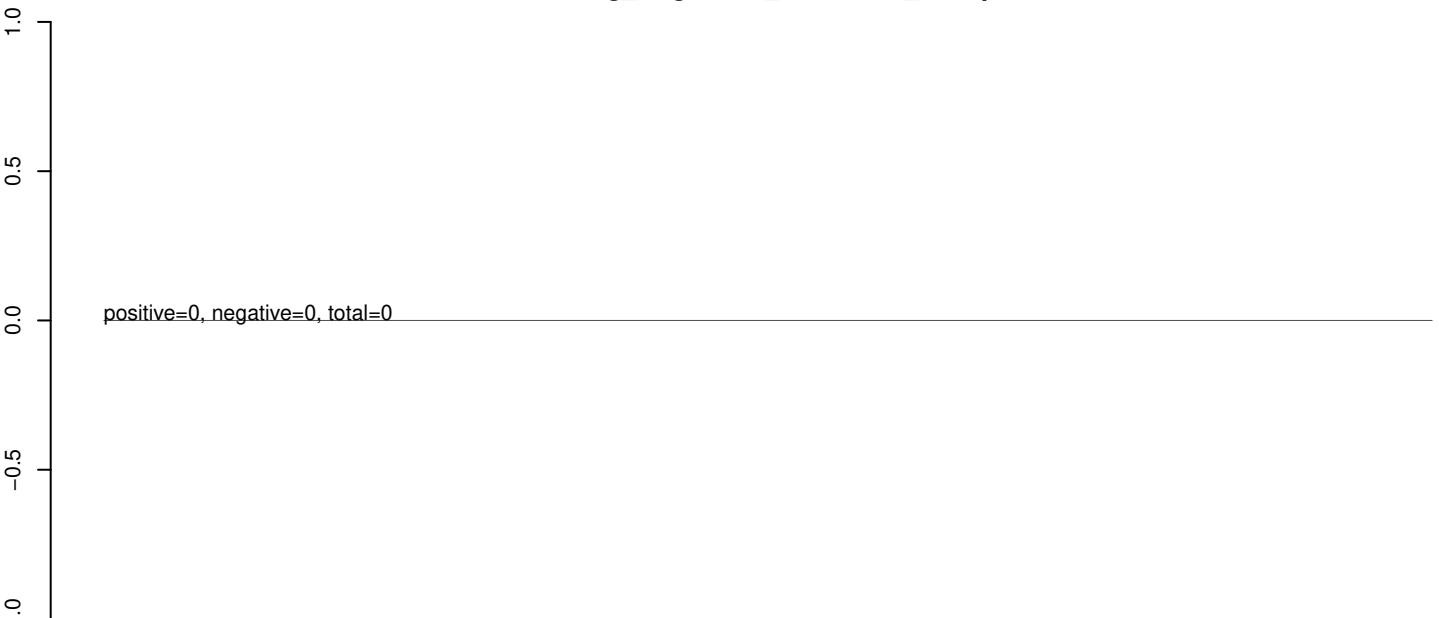
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

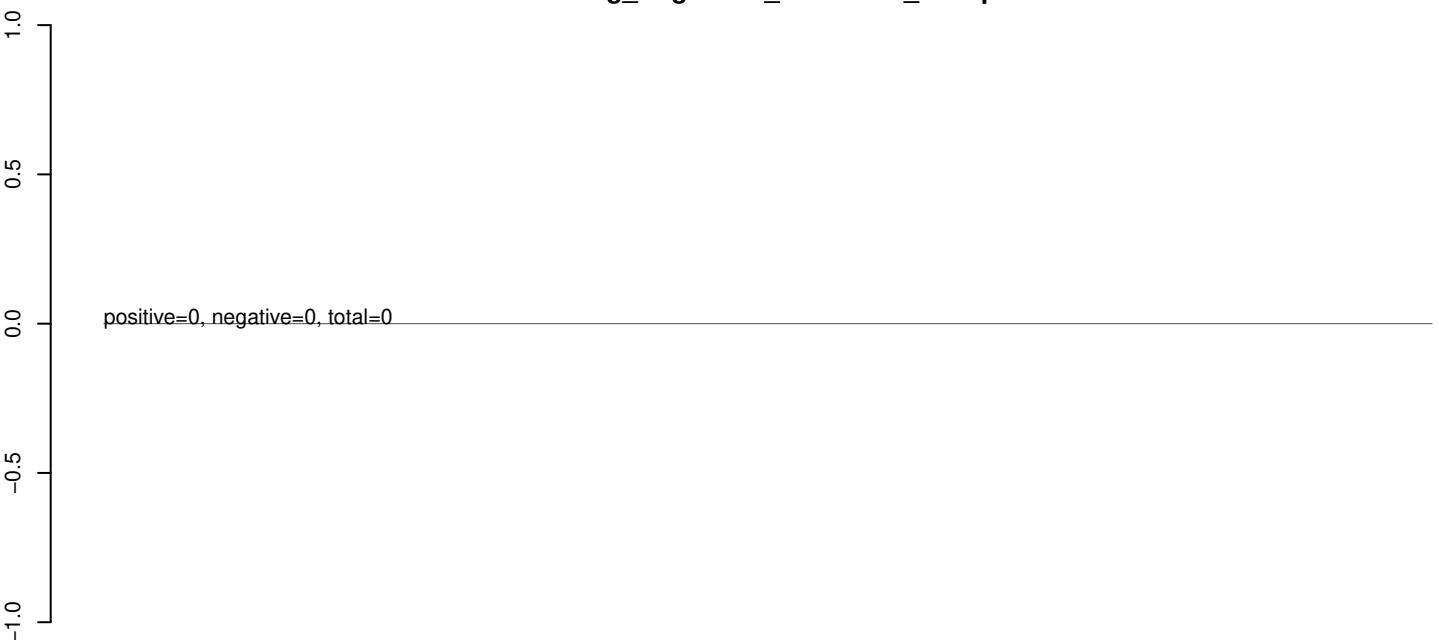
-25, length: 11728, negative@NC_01024457.1-Tai_virus_strain_F47_Cl_2004_segment_S_plus_segment_M_plus_segment_L_nucleocapsid_1-11728



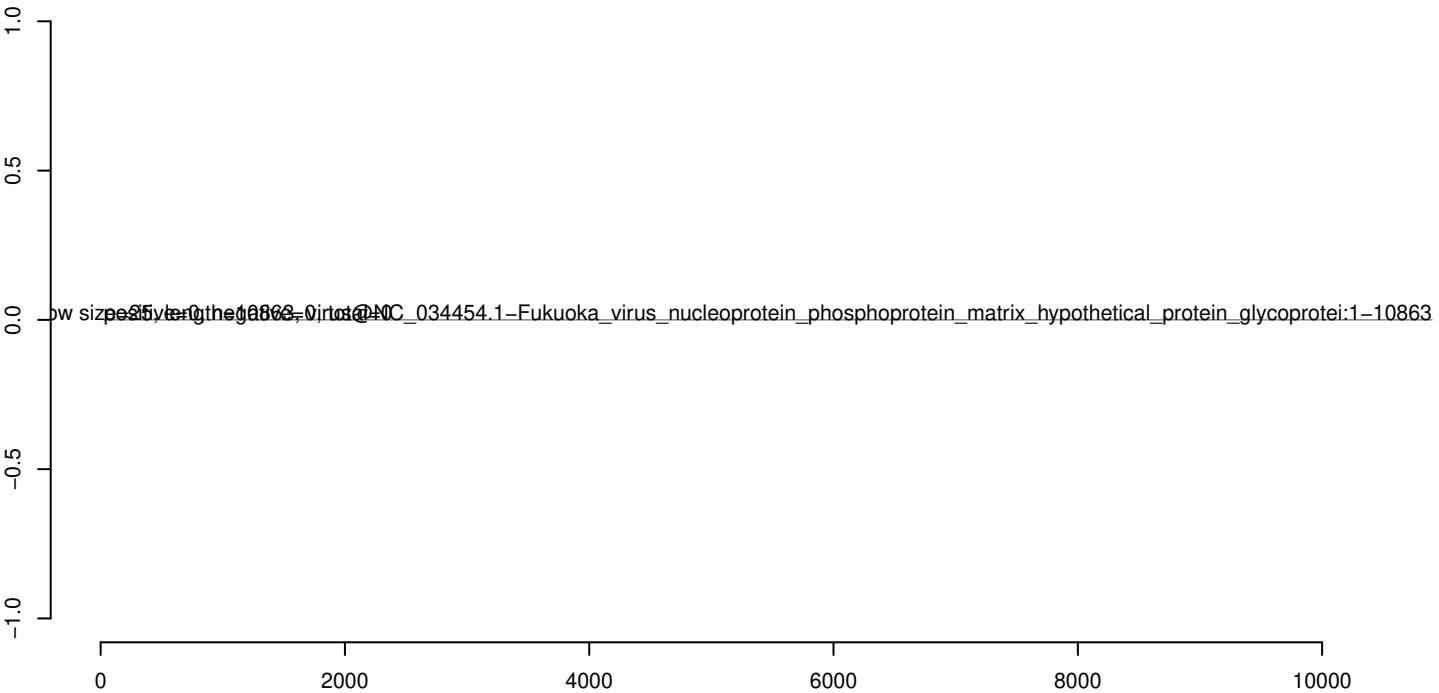
AeAeg_Aag2SINV_GFP.IP.18_23.rep



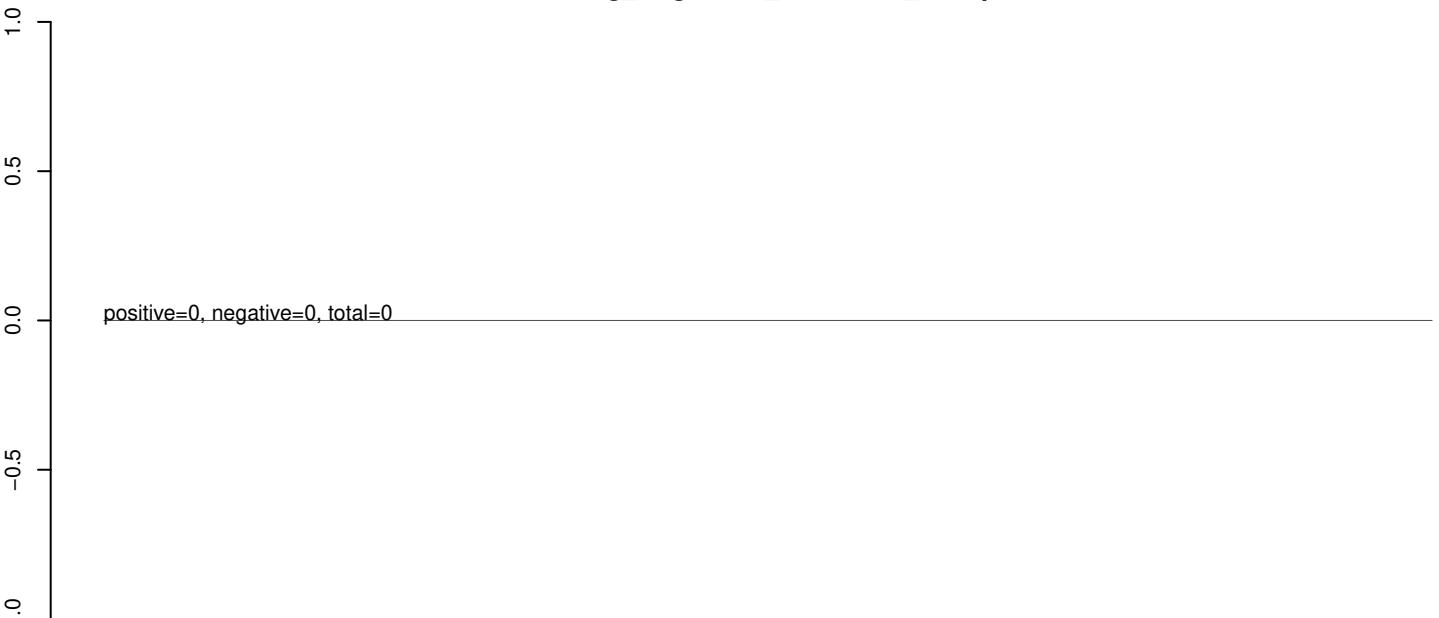
AeAeg_Aag2SINV_GFP.IP.24_35.rep



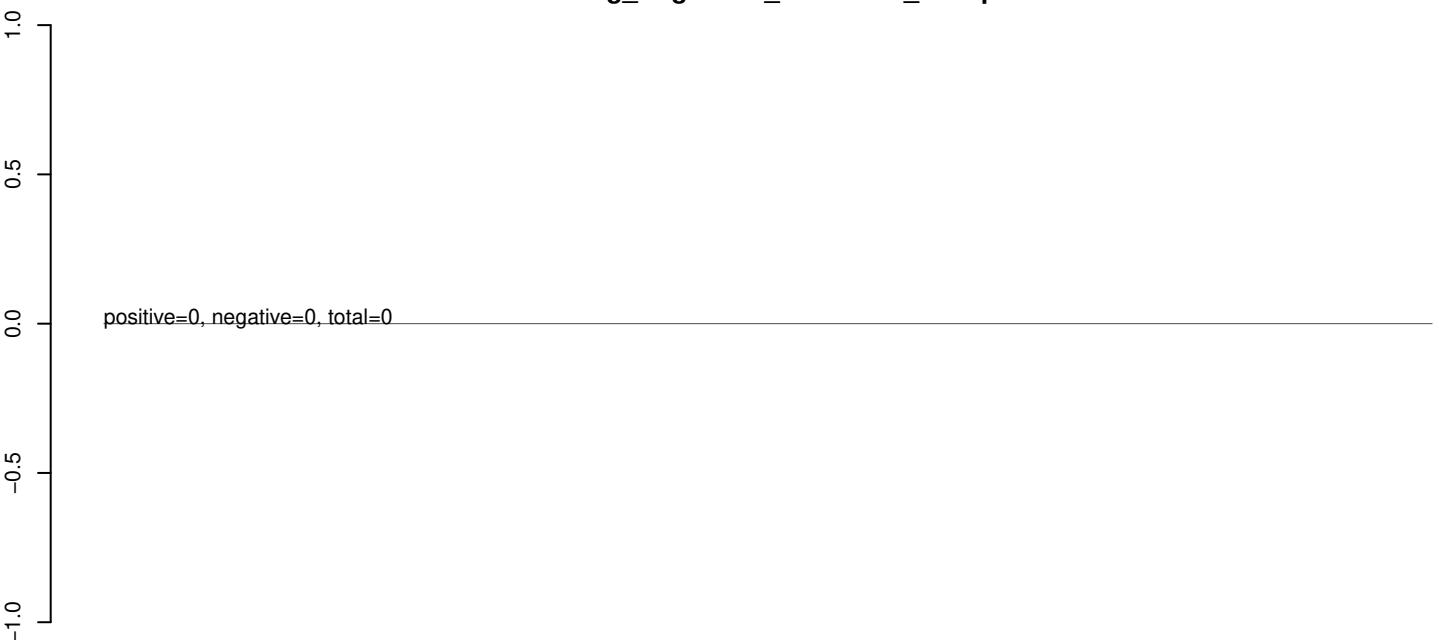
AeAeg_Aag2SINV_GFP.IP.rep



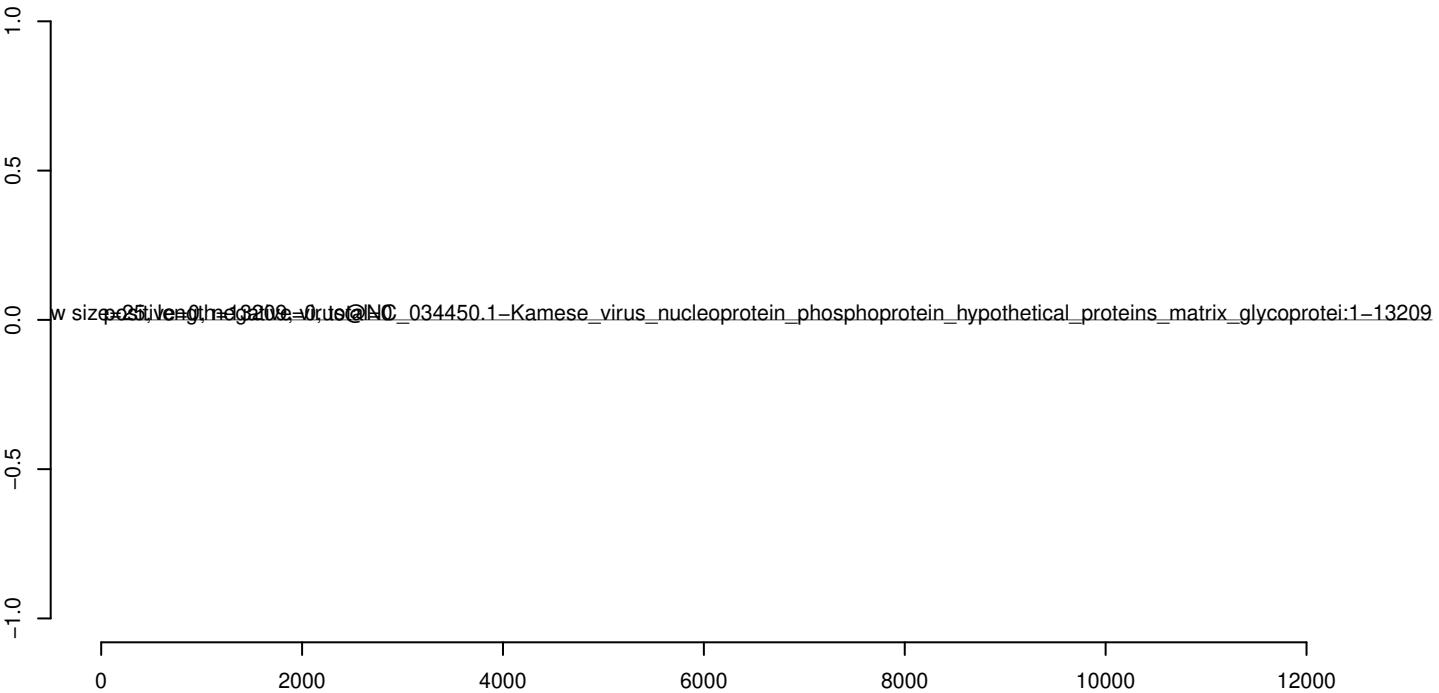
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

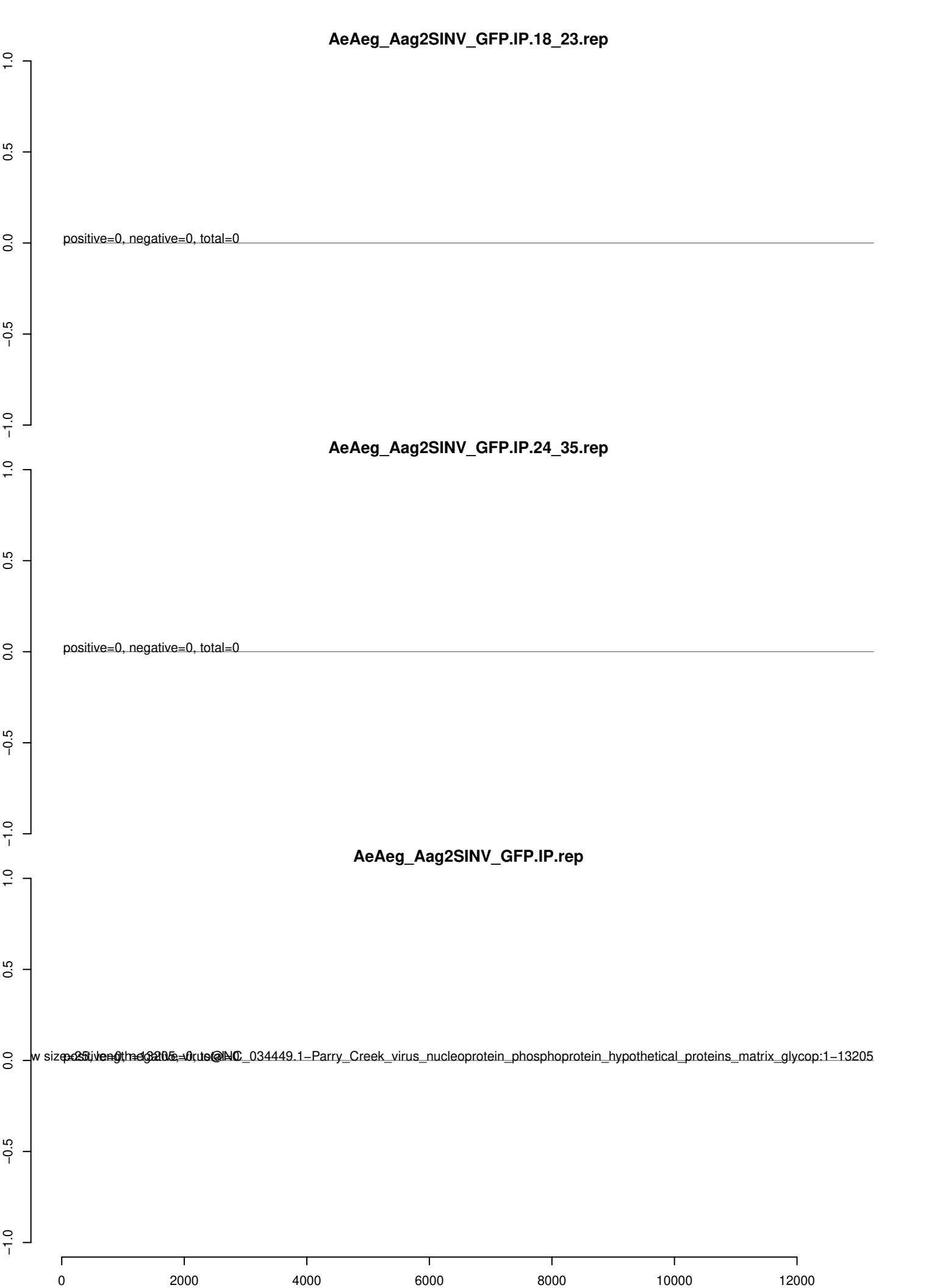
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

w size:25 length:13205 positive=0 negative=0 total@NC_034449.1_Parry_Creek_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycop:1-13205



AeAeg_Aag2SINV_GFP.IP.18_23.rep

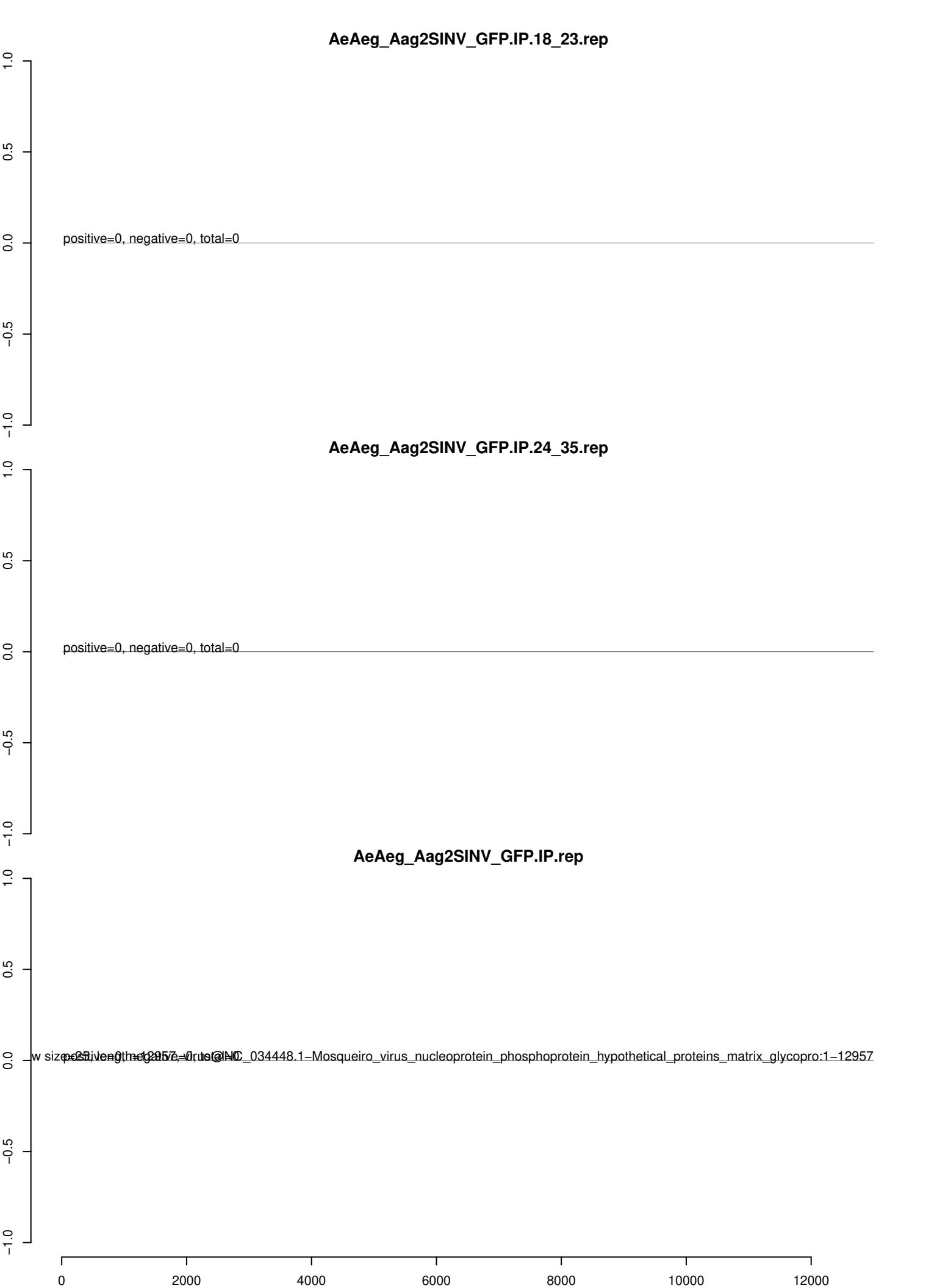
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

w size=25, length=12957, 0 hits@INC_034448.1-Mosqueiro_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopro:1-12957



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0
size=251, mean=0.01, median=0.01, min=-0.01, max=0.01
orthologs@NC_034447.1-Hart_Park_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopro:1-13104



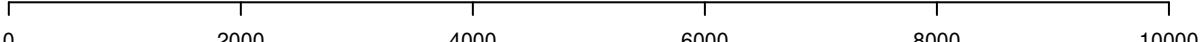
positive=0, negative=0, total=0



positive=0, negative=0, total=0



W size=251 seq=0 meta=1450 -Drust@NC_034443.1-Le_Dantec_virus_nucleoprotein_phosphoprotein_matrix_glycoprotein_hypothetical_prot:1-11450



AeAeg_Aag2SINV_GFP.IP.18_23.rep

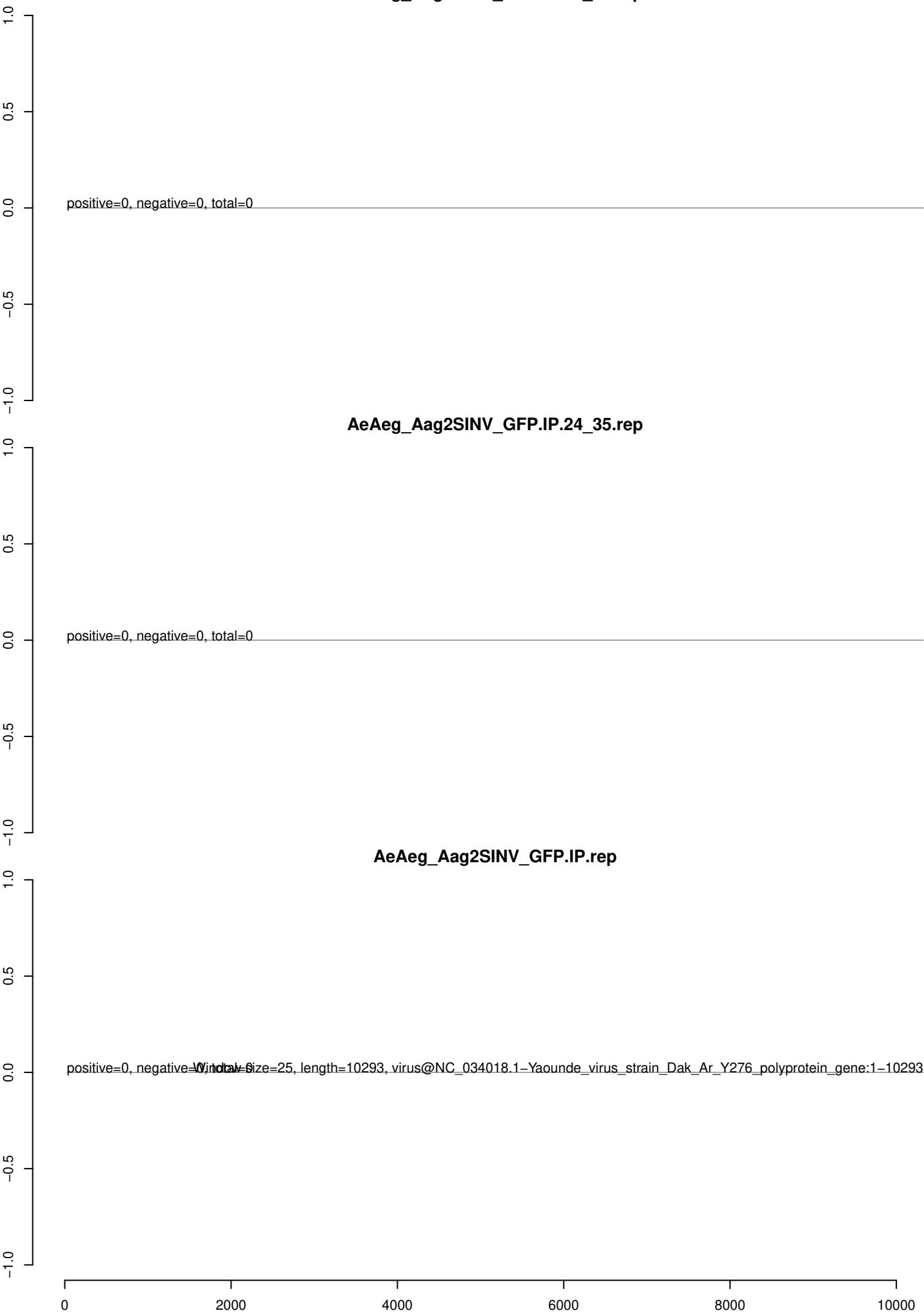
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=~~Window~~, size=25, length=10293, virus@NC_034018.1_Yaounde_virus_strain_Dak_Ar_Y276_polyprotein_gene:1-10293



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

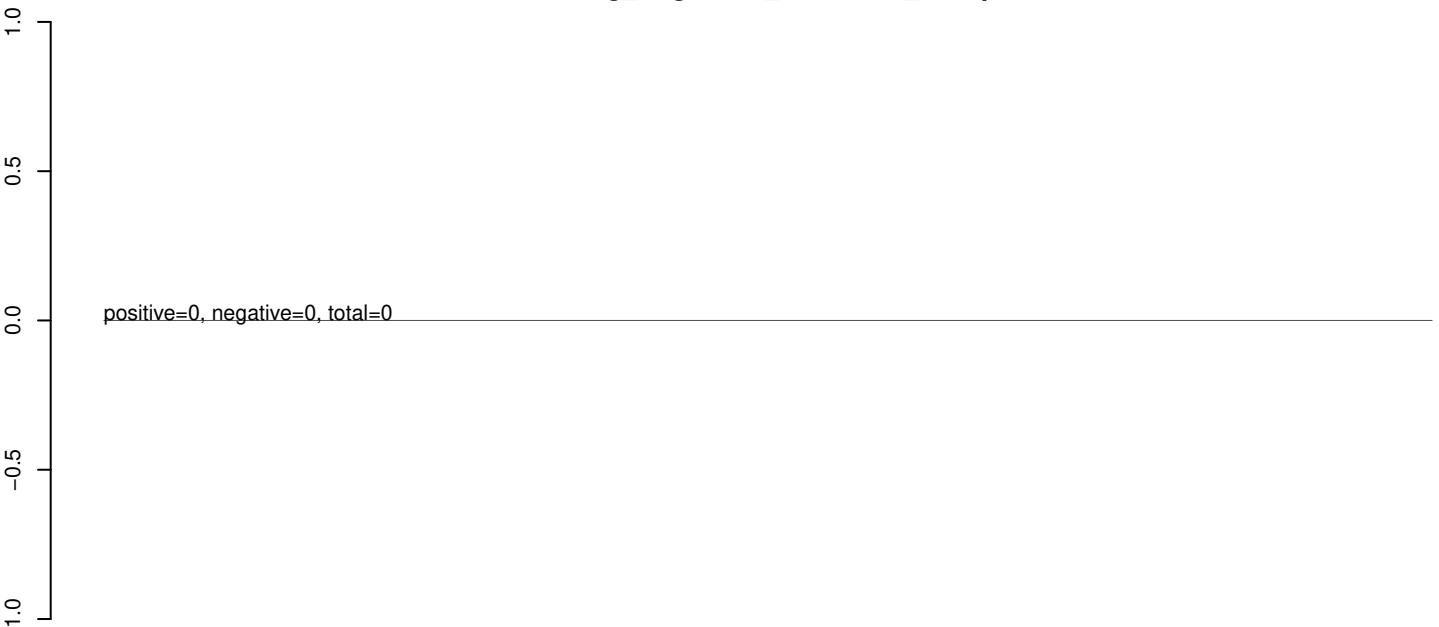
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

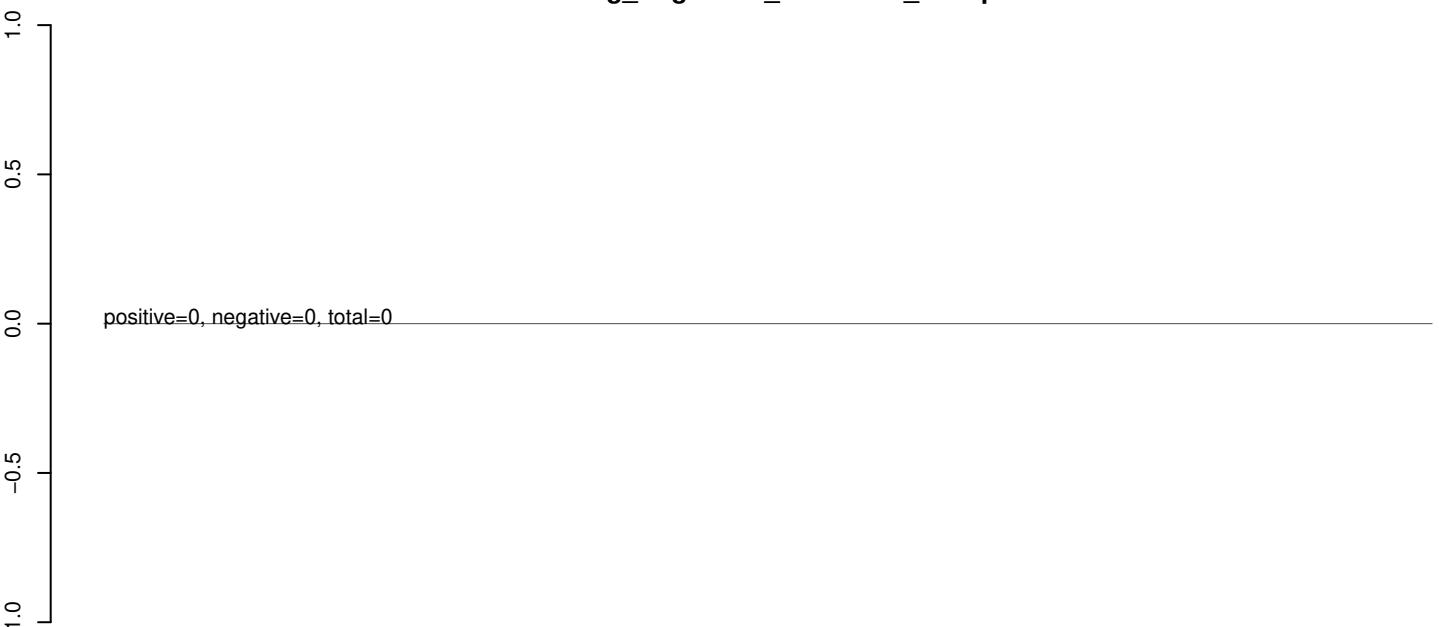
AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, Window size=25, length=10215, virus@NC_033724.1-Kadam_virus_from_Uganda_polyprotein_gene:1-10215

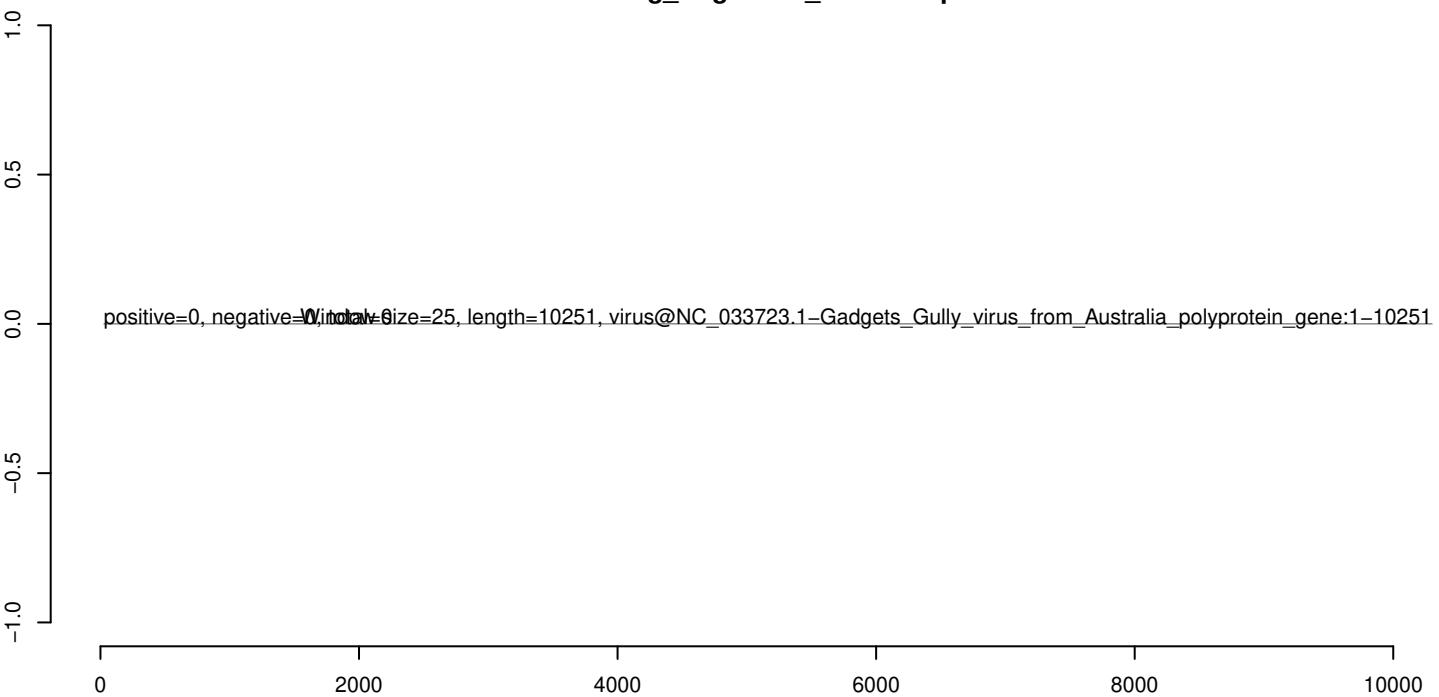
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=Window size=25, length=10266, virus@NC_033721.1-Meaban_virus_from_France_polyprotein_gene:1-10266

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

Window size:256, Overlap:0, virus@NC_033716.1_Zhejiang_mosquito_virus_1_strain_mosZJ35497_hypothetical_protein_gene:1-9558

0 2000 4000 6000 8000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=Window size=25, length=10173, virus@NC_033699.1-Jugra_virus_strain_P-9-314_polyprotein_gene:1-10173

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

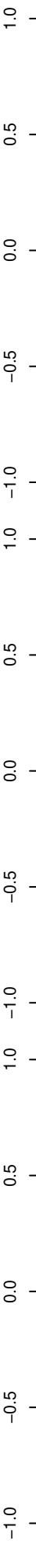
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10182, virus@NC_033698.1-Uganda_S_virus_polyprotein_gene:1-10182



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, size=25, length=10173, virus@NC_033697.1-Saboya_virus_strain_Dak_AR_D4600_polyprotein_gene:1-10173

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

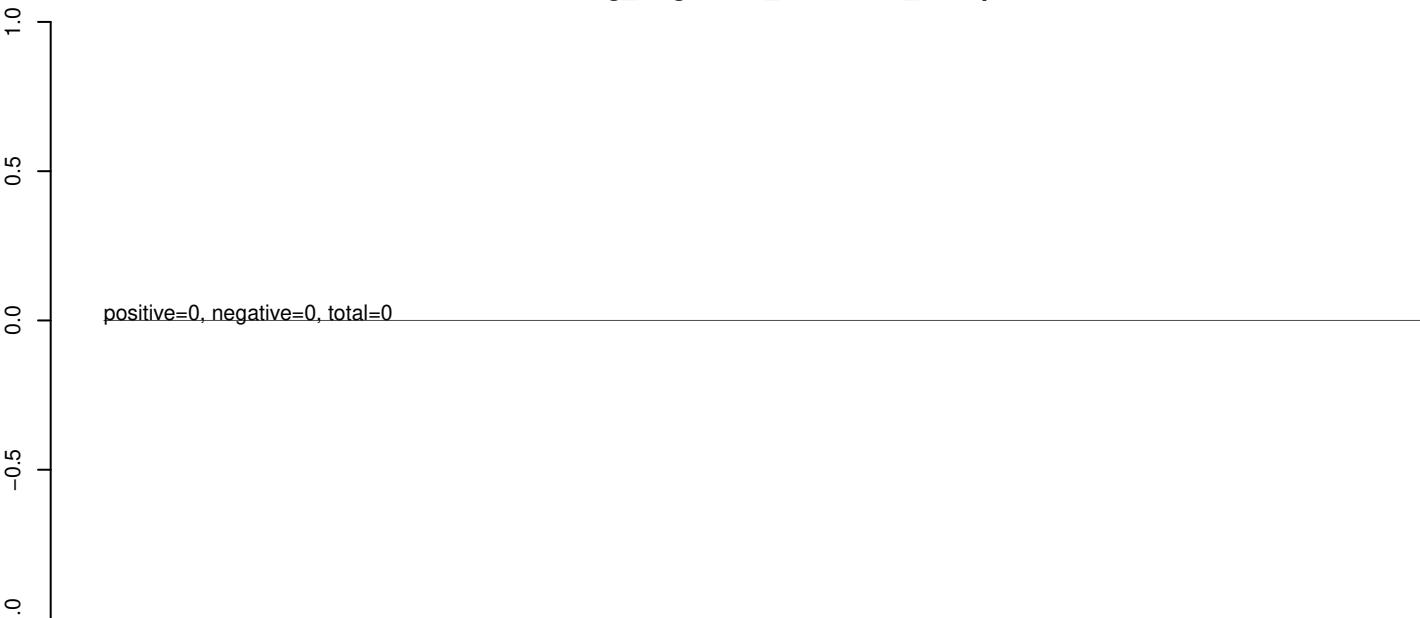
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

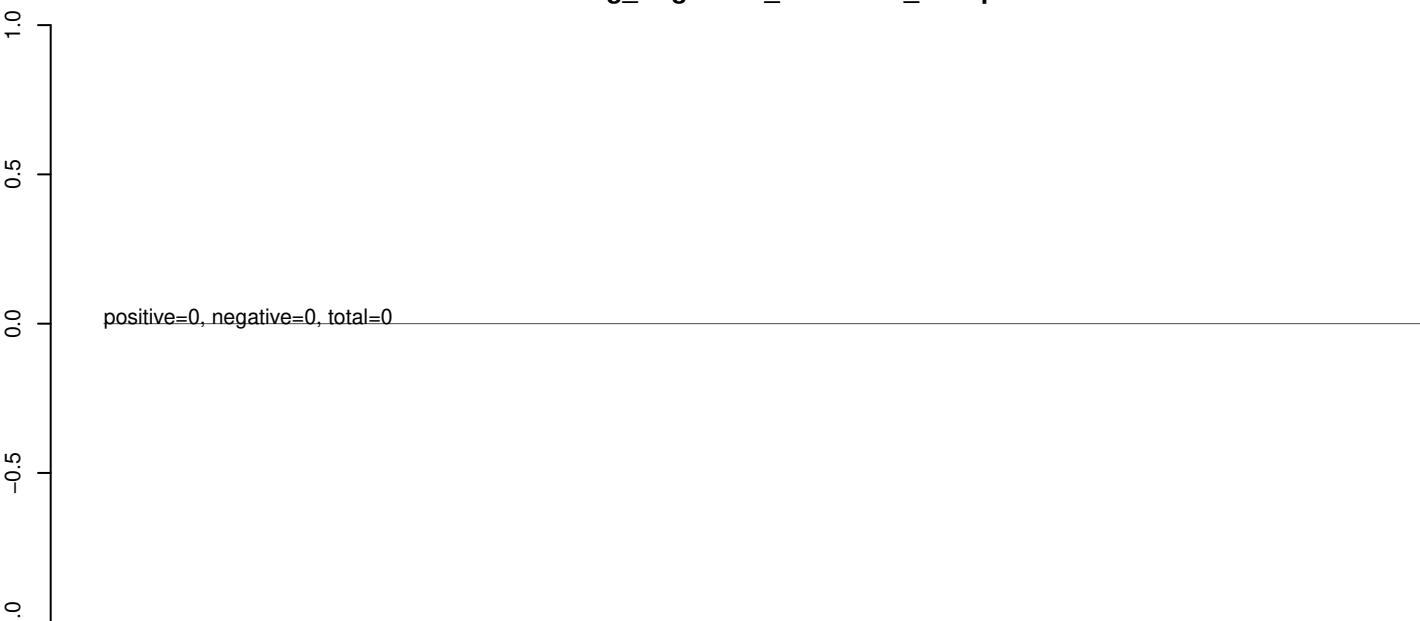
AeAeg_Aag2SINV_GFP.IP.rep

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

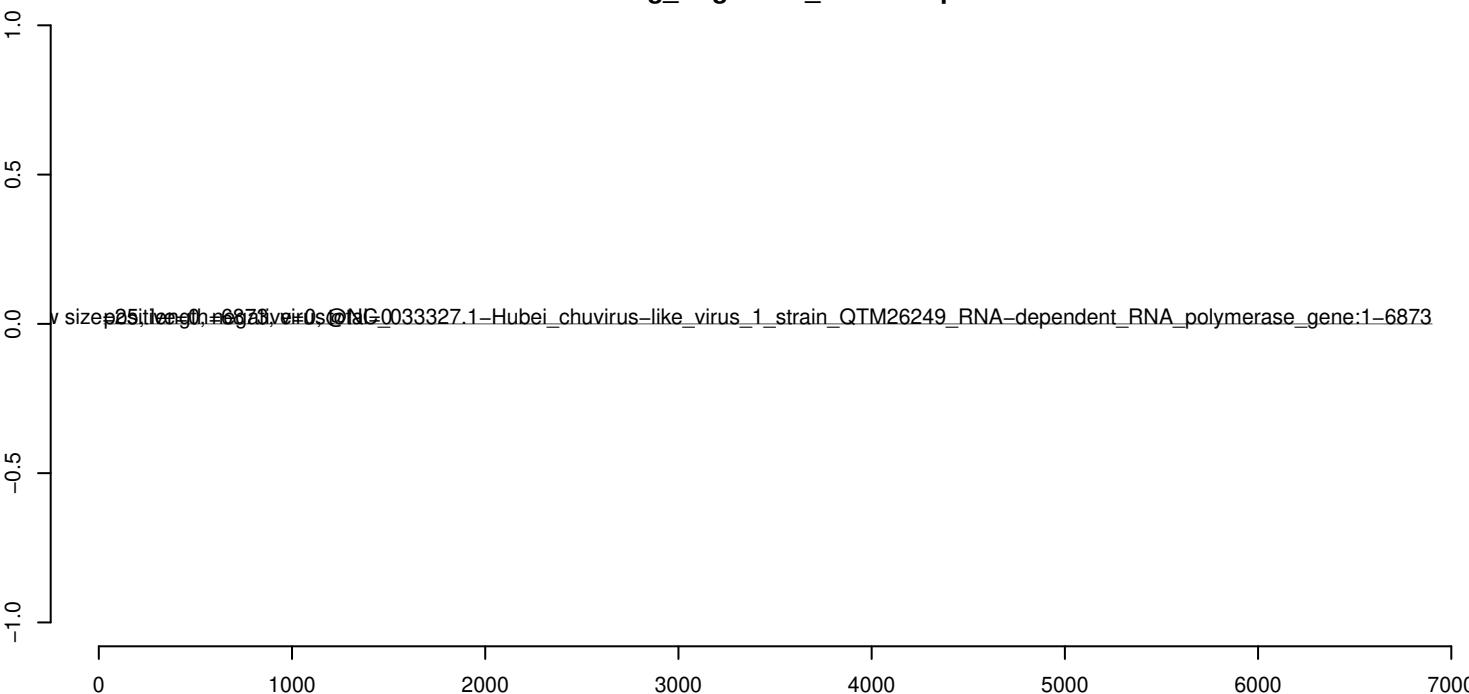
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

25, length=11280, positive=0, negative=0, total=0
NC_0123015.1-Hubei_chuvirus-like_virus_3_strain_QTM26698_RNA-dependent_RNA_polymerase_putative_1-11280

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

size=25 length=3226 positive=0 negative=0
Q1011032491.1_Zhejiang_mosquito_virus_3_strain_mosZJ35354_hypothetical_protein_and_RNA-dependent:1-3226

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, window size=25, length=10869, virus@NC_031957.1-UNVERIFIED_Coot_Bay_virus_strain_EVG5-53:1-10869

AeAeg_Aag2SINV_GFP.IP.18_23.rep

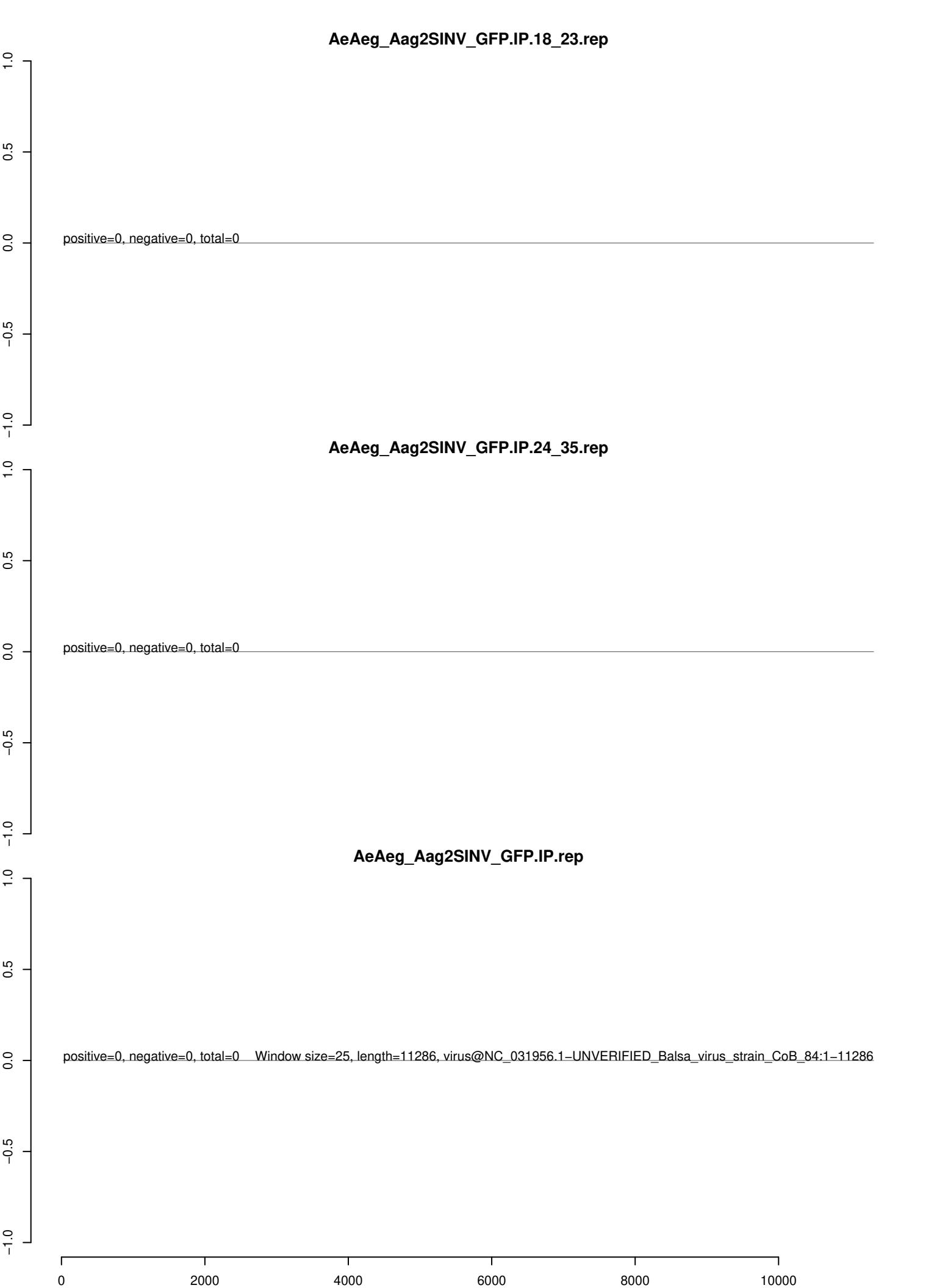
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0 Window.size=25, length=11286, virus@NC_031956.1-UNVERIFIED_Balsa_virus_strain_CoB_84:1-11286



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

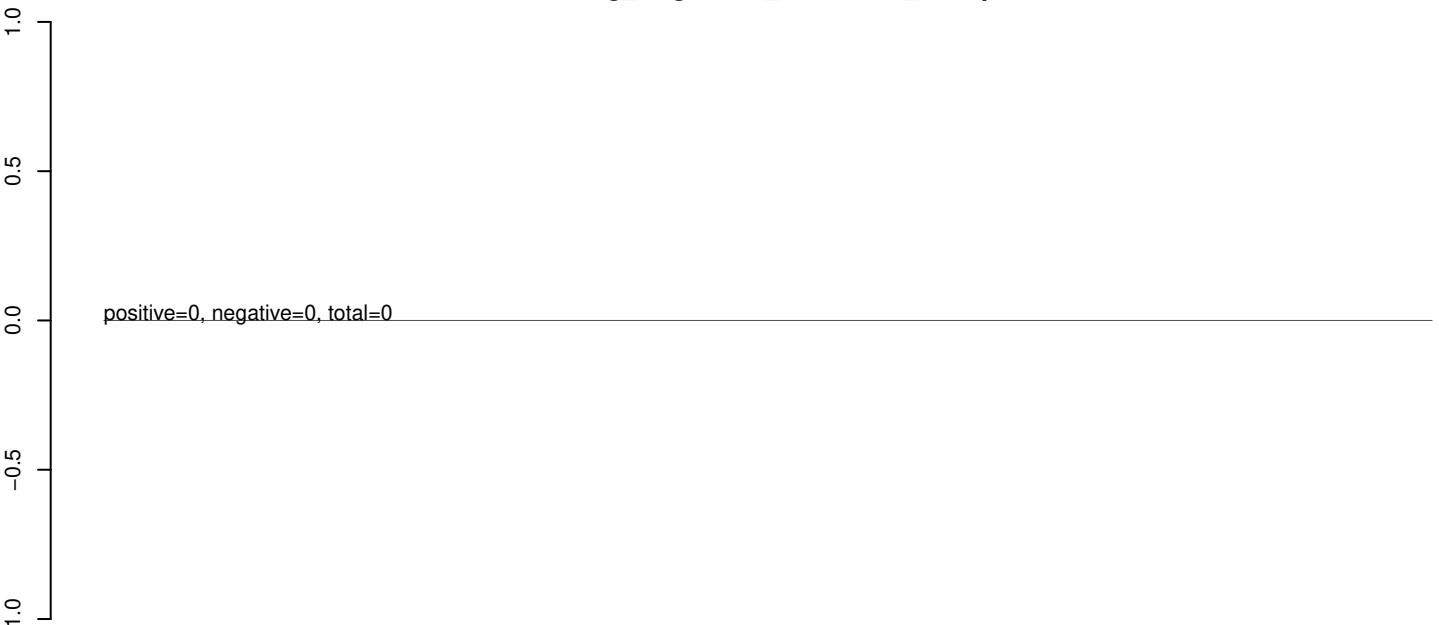
AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

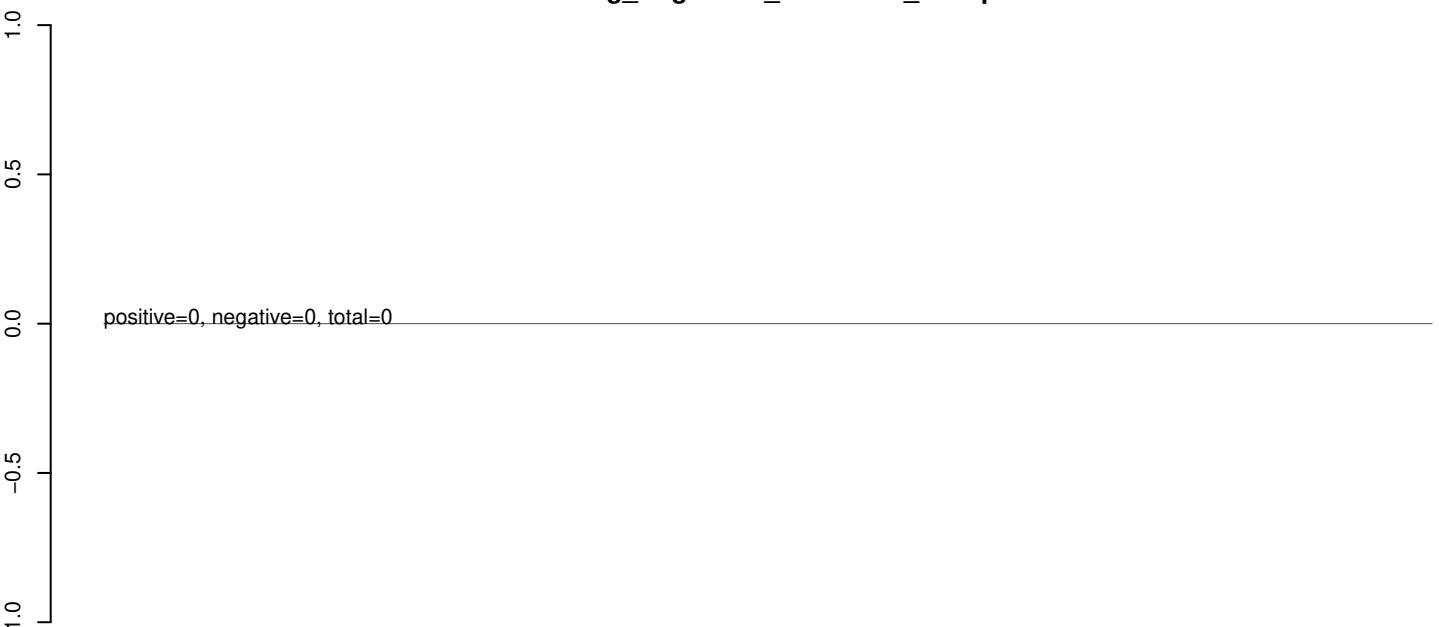
Window size=25, length=10588, virus@NC_031327.1-Anopheles_flavivirus_variant_1:1-10588

0 2000 4000 6000 8000 10000

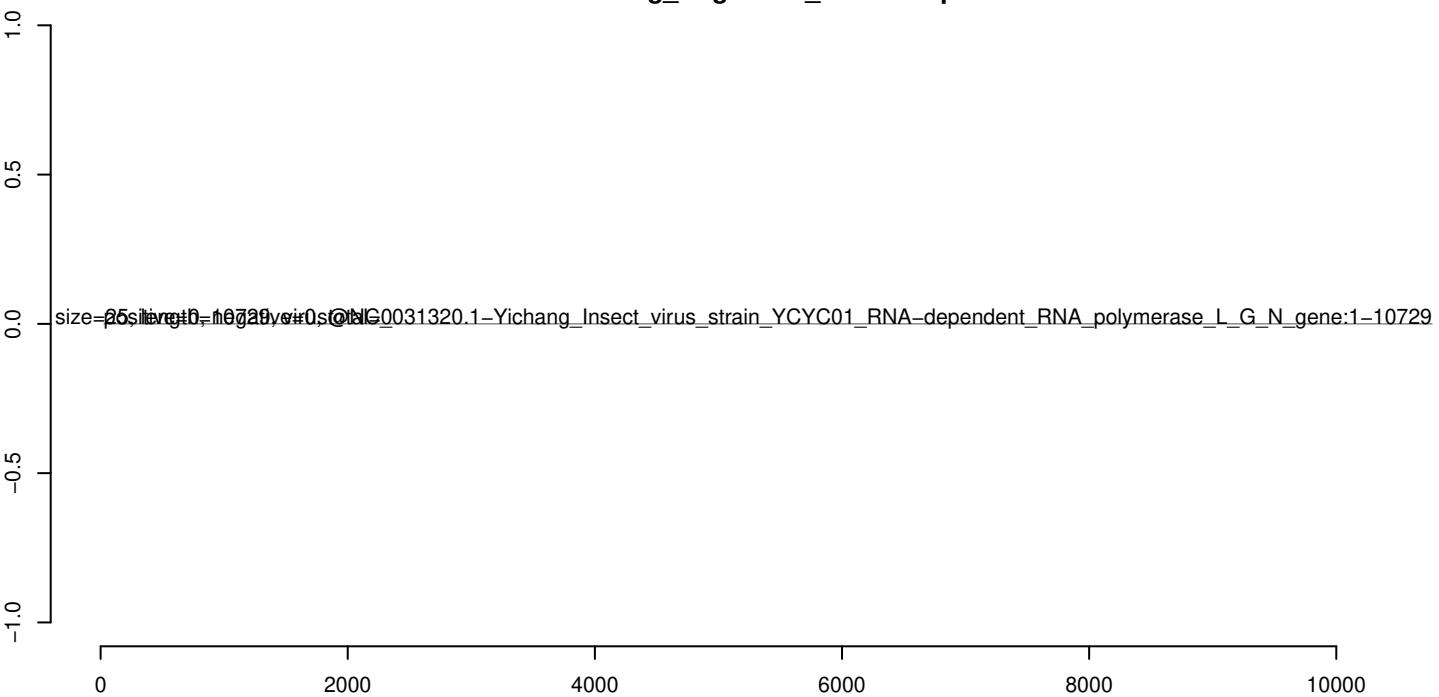
AeAeg_Aag2SINV_GFP.IP.18_23.rep



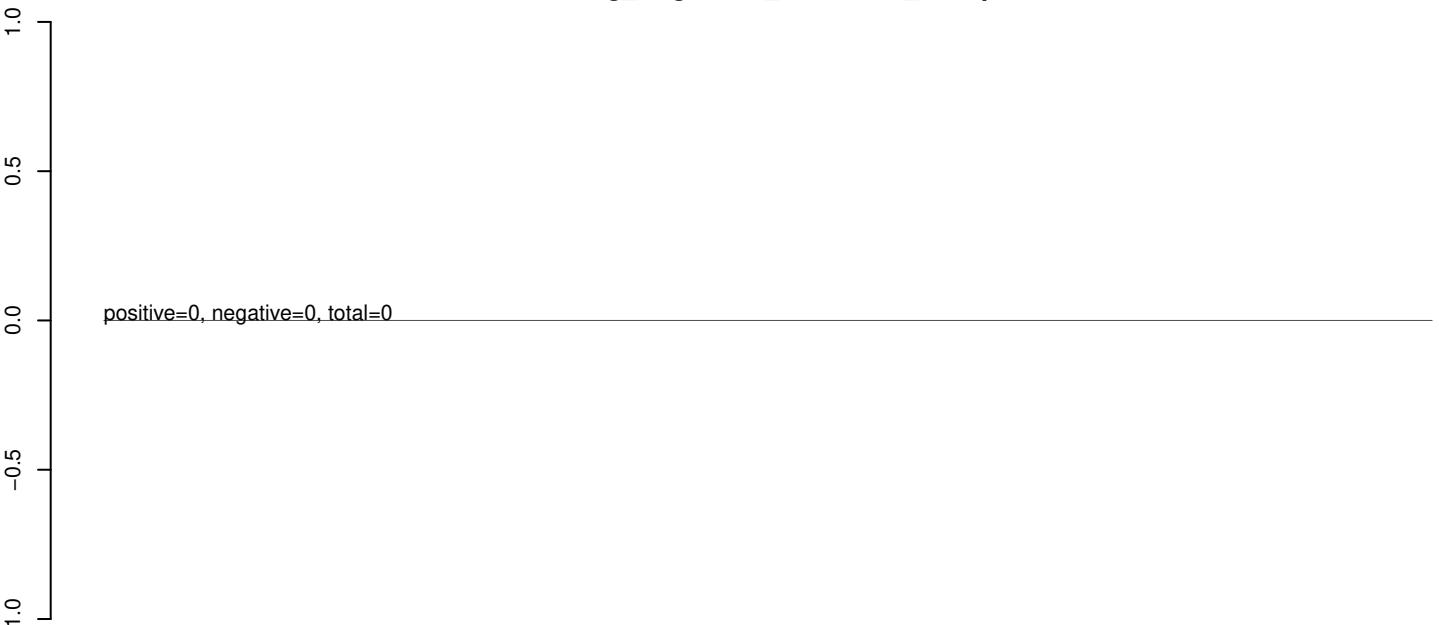
AeAeg_Aag2SINV_GFP.IP.24_35.rep



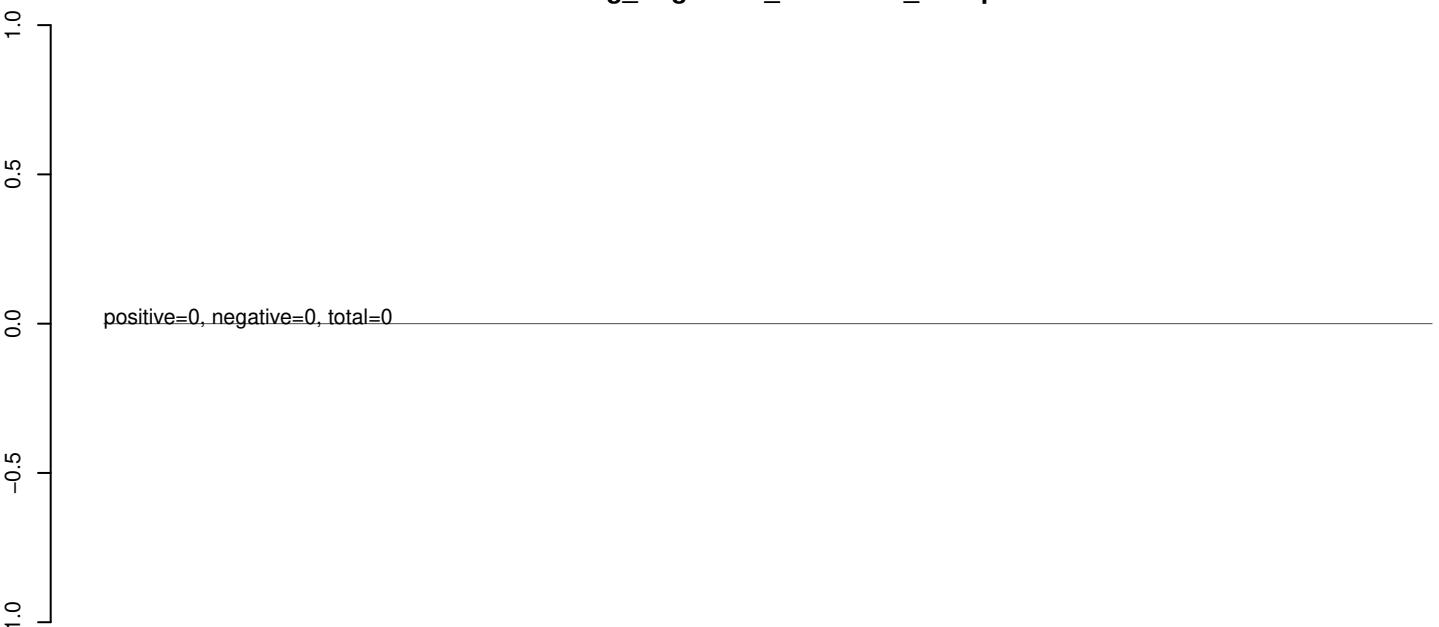
AeAeg_Aag2SINV_GFP.IP.rep



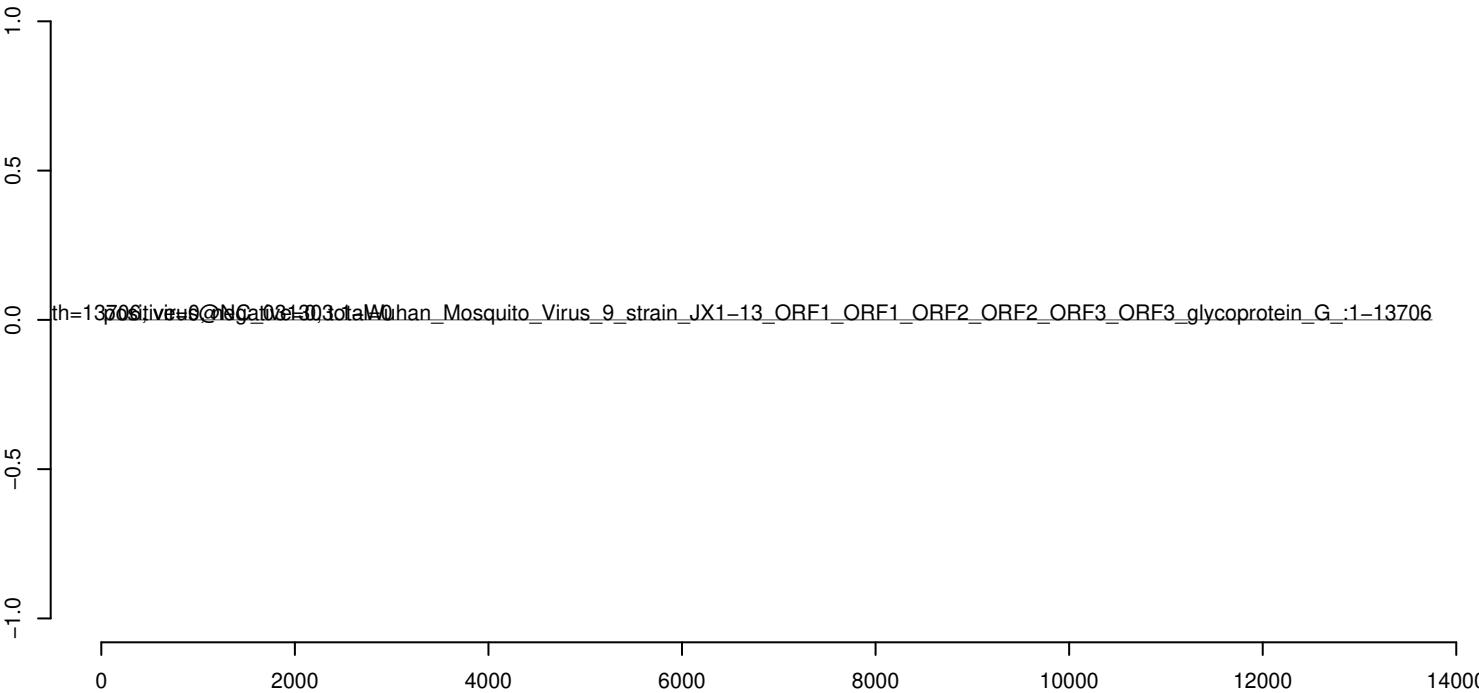
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

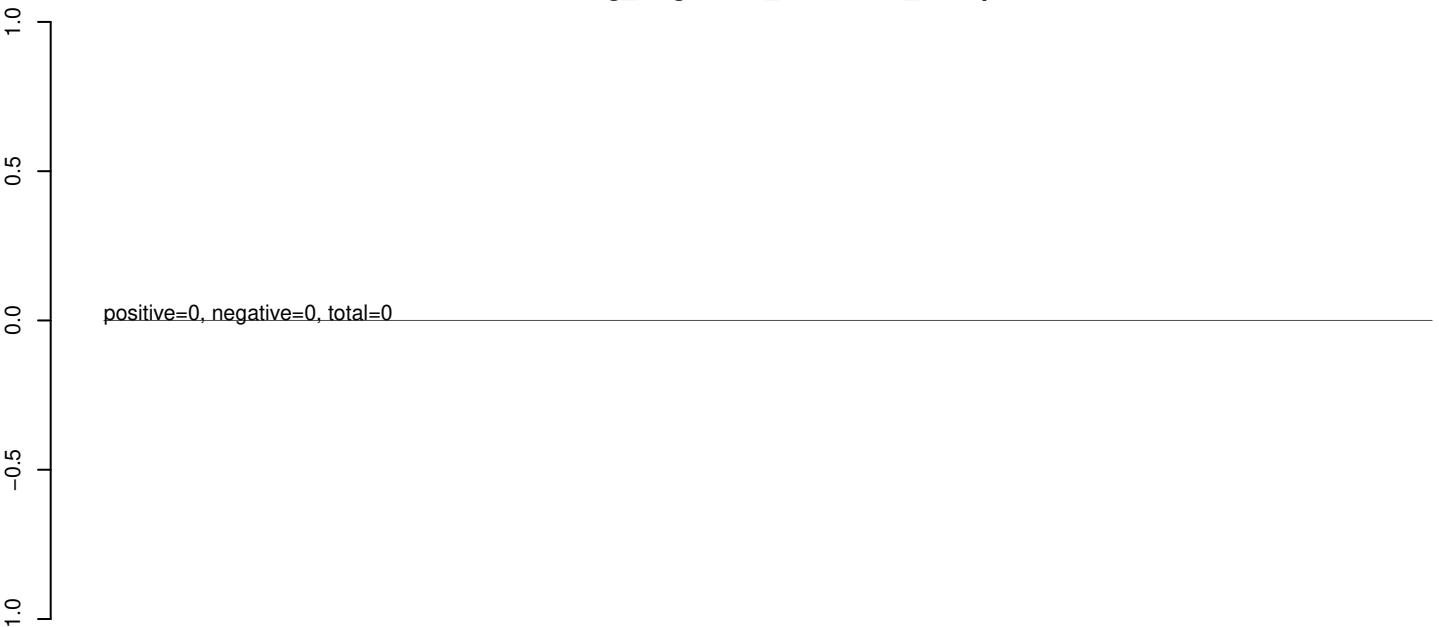
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

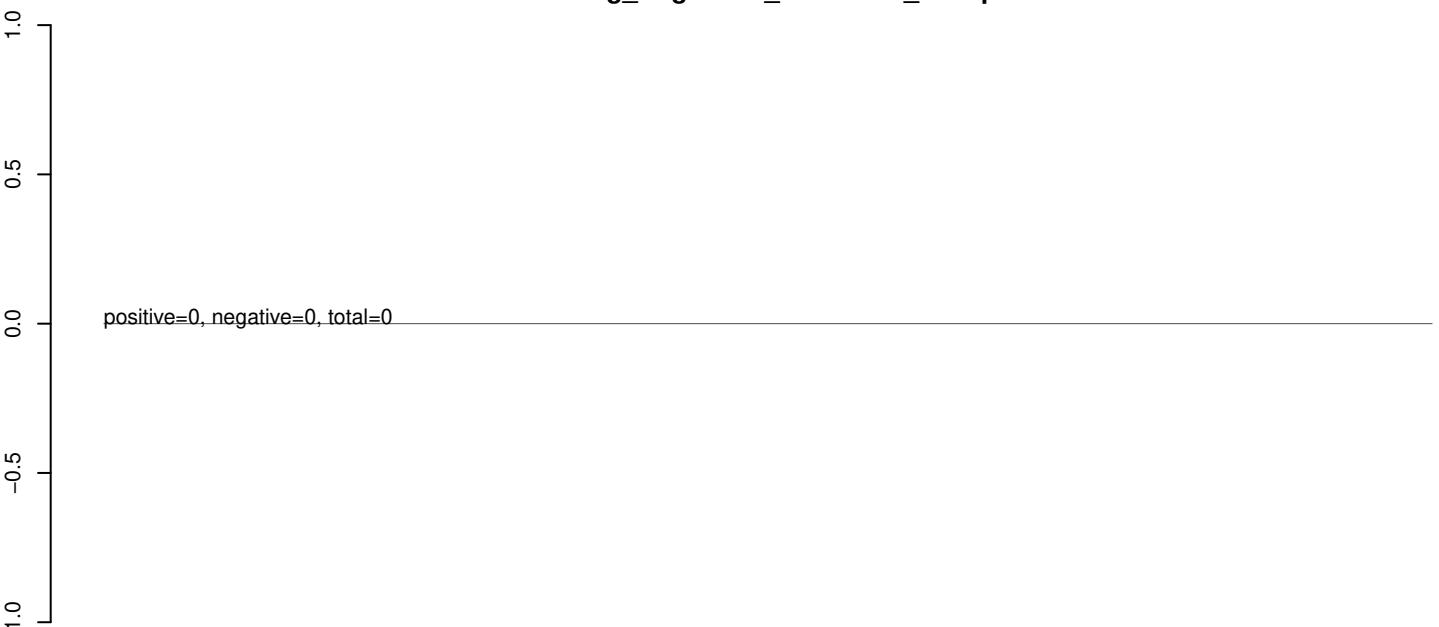
length=12774, e-value=0.0, p-value=0.0302641, Xincheng_Mosquito_Virus_strain_XC1-6_ORF1_ORF1_ORF2_ORF2_glycoprotein_G_and_RNA-de:1-12774

0 2000 4000 6000 8000 10000 12000

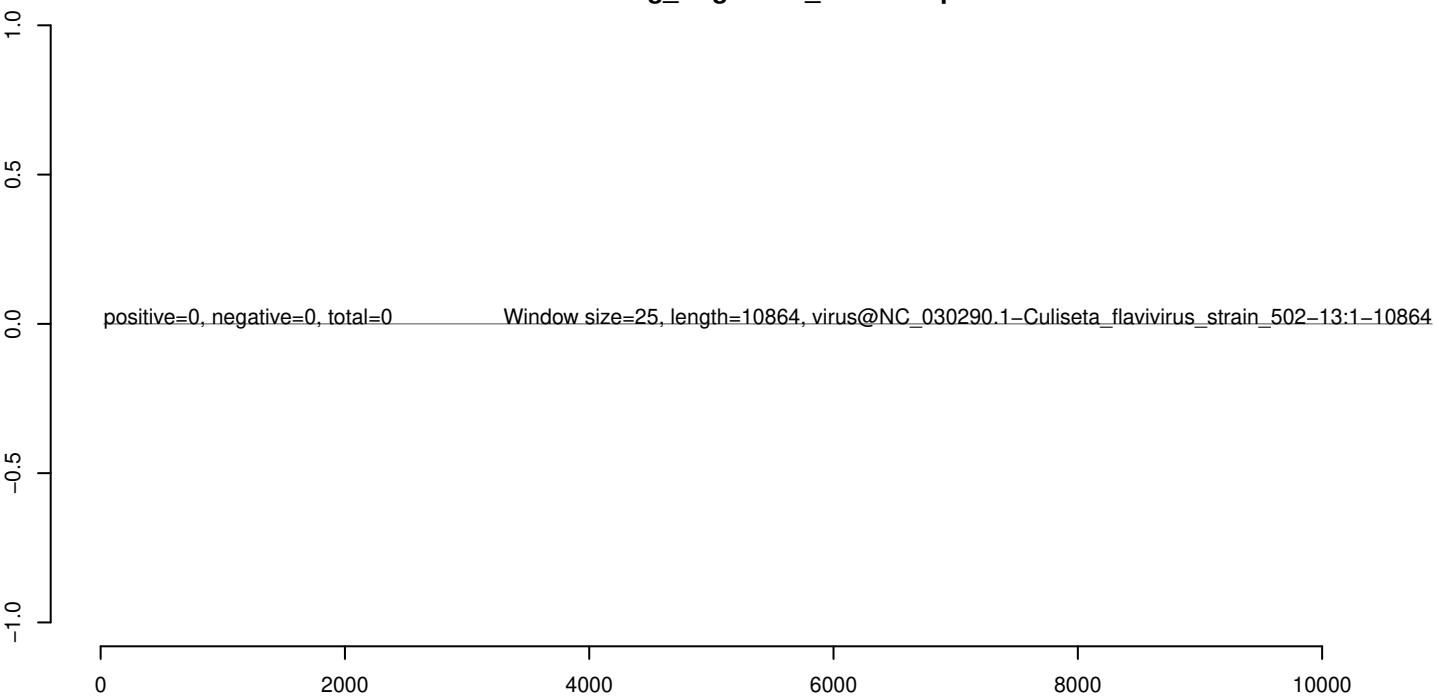
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, window size=25, length=10206, virus@NC_030289.1-Edge_Hill_virus_strain_YMP_48_polyprotein_gene:1-10206

0 2000 4000 6000 8000 10000



positive=0, negative=0, total=0



positive=0, negative=0, total=0



Windows-1252|negative_8919|virus@NC_030115.1-Anopheles_C_virus_non-structural_polyprotein_and_structural_polyprotein_genes:1-8919

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0 Window.size=25, length=12004, virus@NC_028265.1_Wuhan_Mosquito_Virus_8_strain_XC2-7:1-12004

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

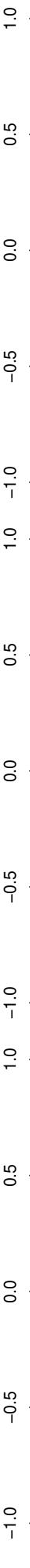
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=14478, virus@NC_028241.1-Yata_virus_isolate_DakArB_2181:1-14478



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window.size=25, length=10938, virus@NC_027819.1-Mercadeo_virus_isolate_ER-M10:1-10938

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

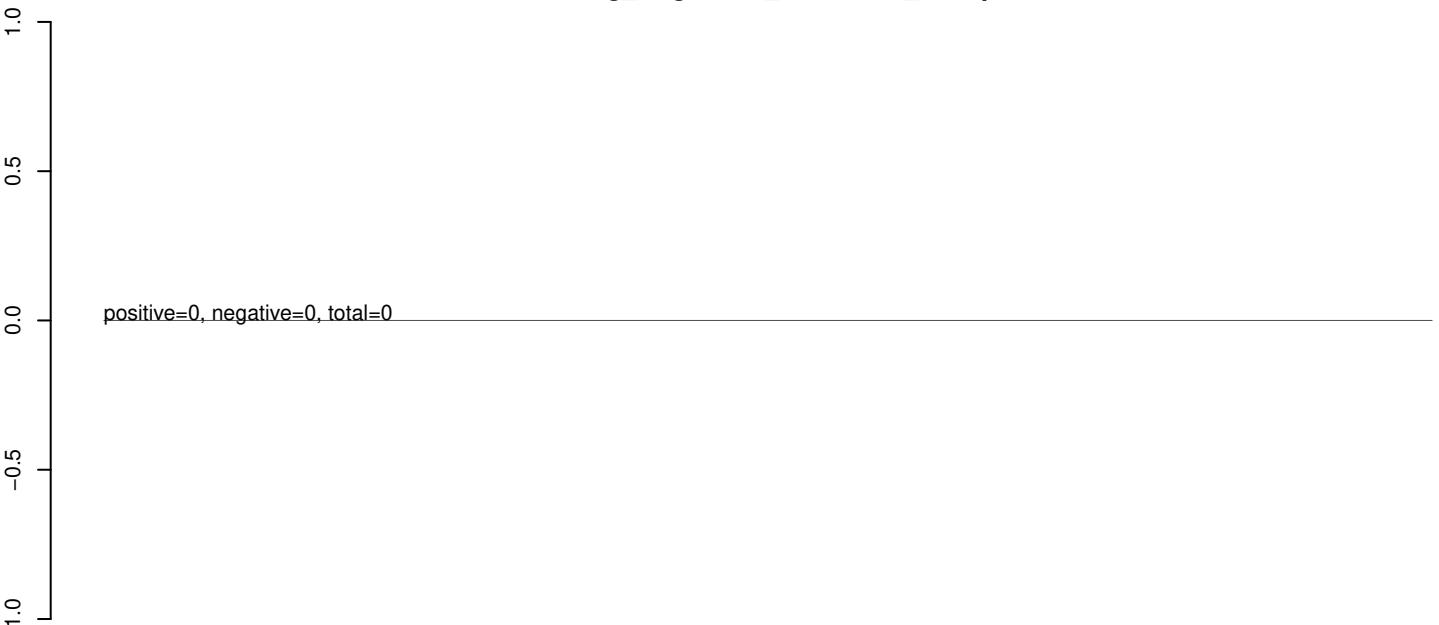
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

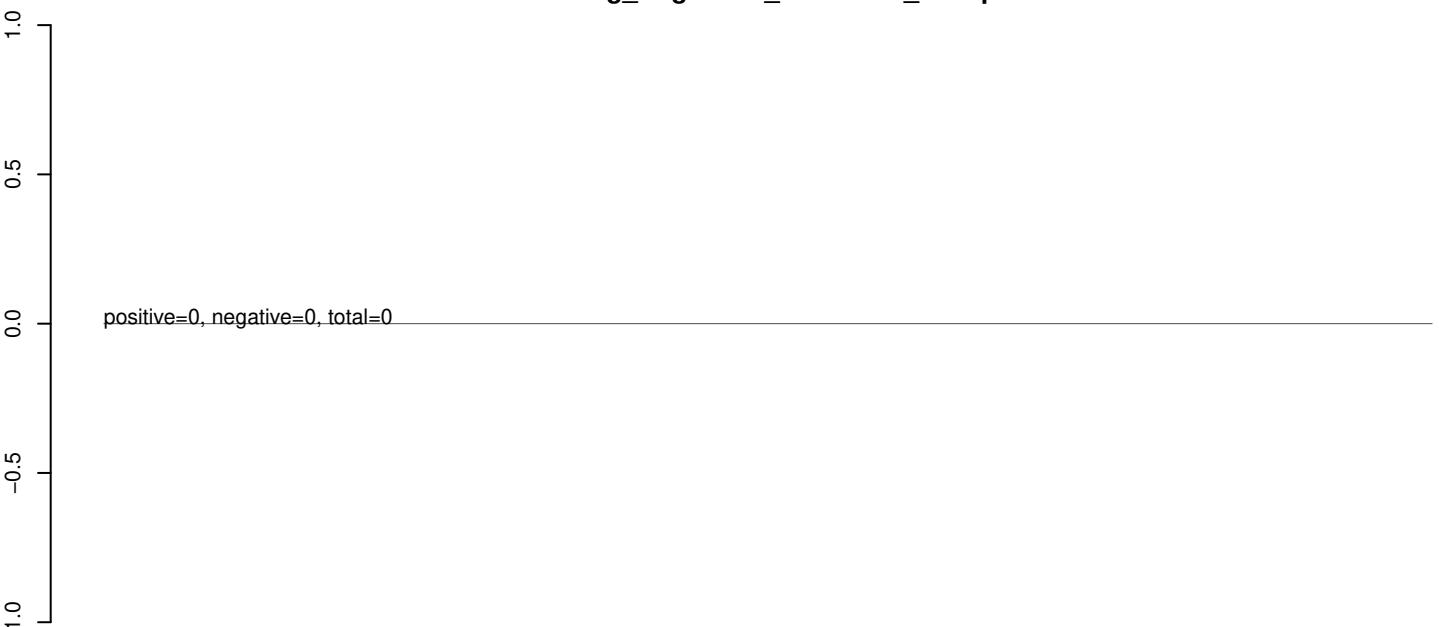
positive=0, negative=0, total=0 Window size=25, length=10893, virus@NC_027817.1_Parramatta_River_virus_isolate_92_B115745:1-10893

0 2000 4000 6000 8000 10000

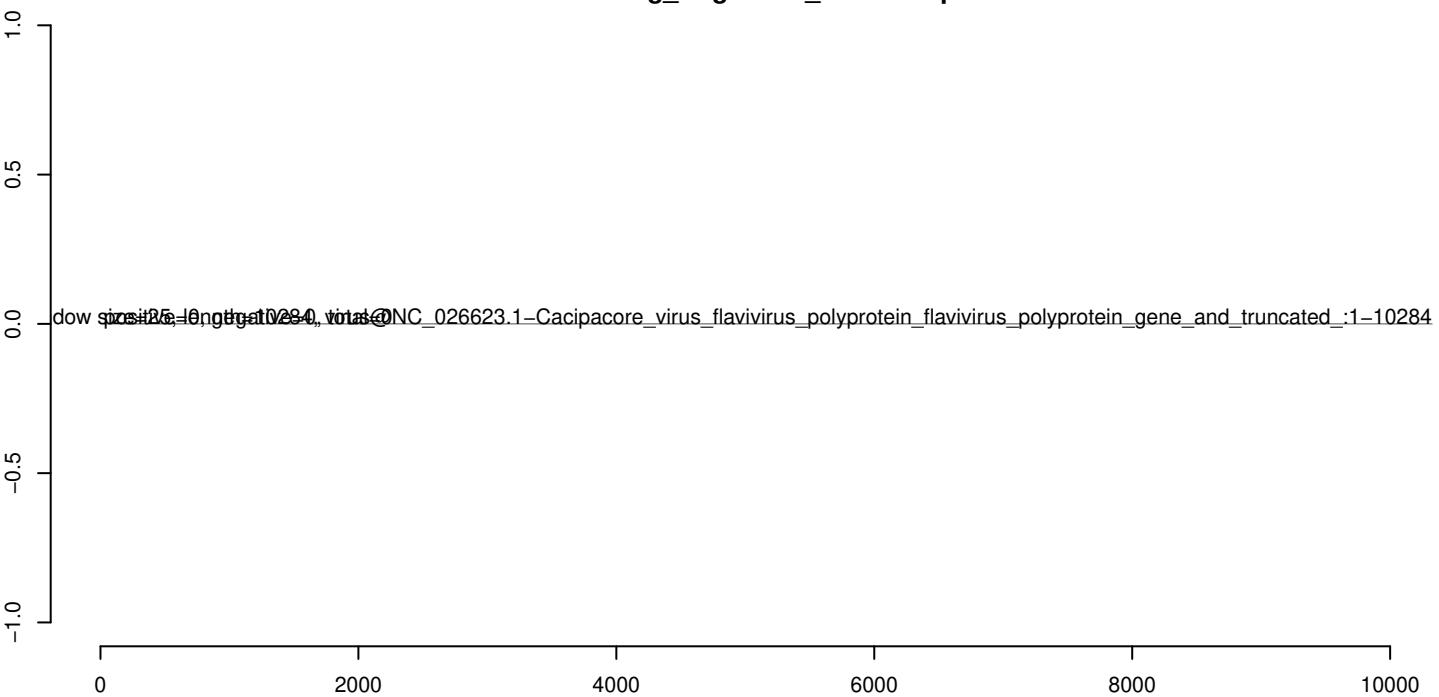
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10125, virus@NC_026620.1-Jutiapa_virus_strain_JG-128:1-10125

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

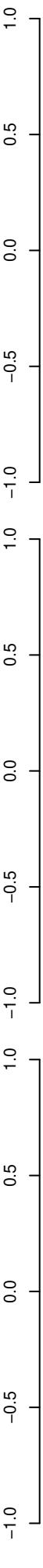
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=15444, virus@NC_025400.1-Malakal_virus_isolate_SudAr_1169-64:1-15444



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=11220, virus@NC_025399.1-Oak-Vale_virus_strain_CSIRO_1342:1-11220

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

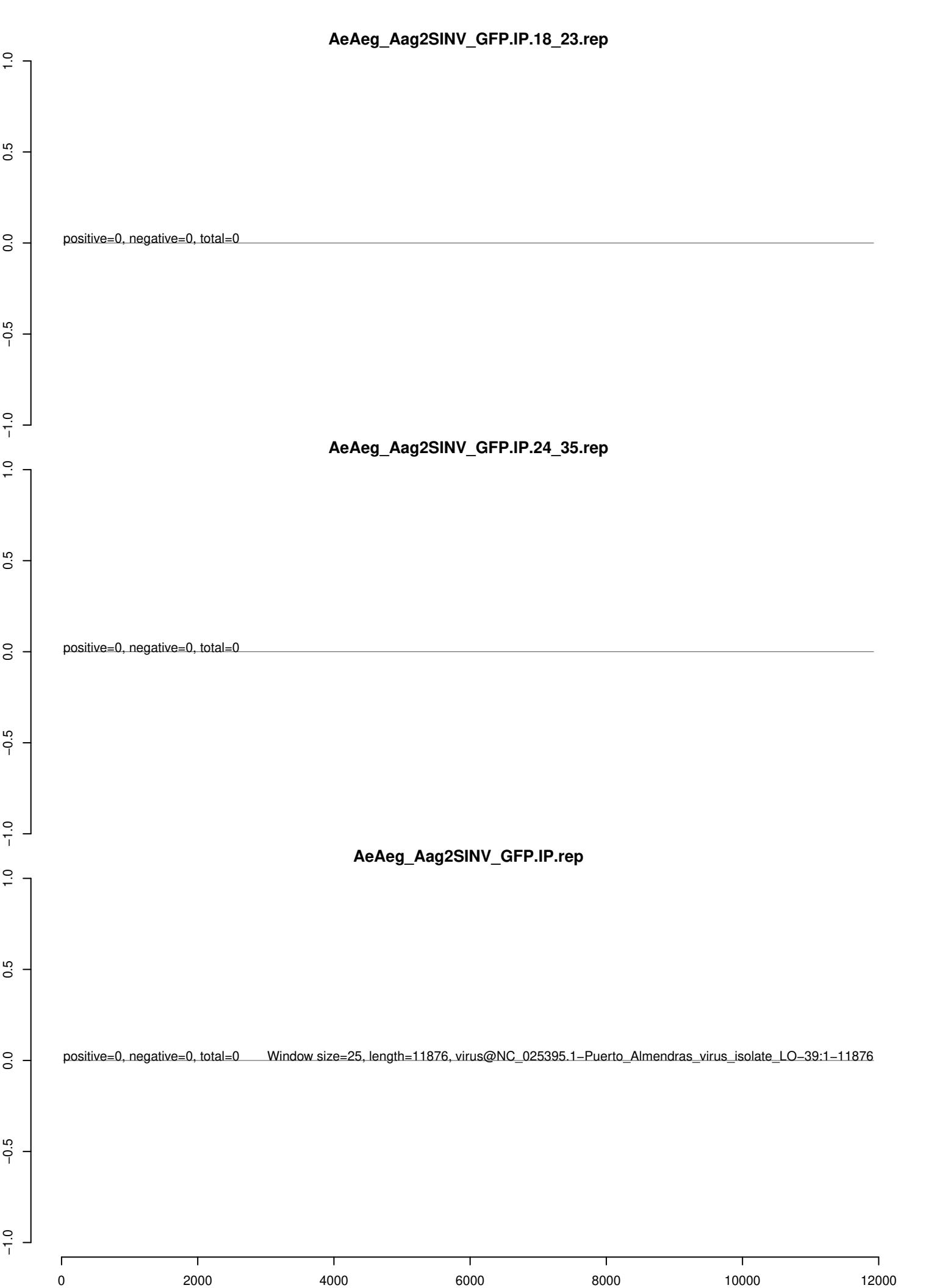
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0 Window size=25, length=11876, virus@NC_025395.1–Puerto_Almendras_virus_isolate_LO-39:1–11876



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0 size=25, length=11190, virus@NC_025384.1_Culex_tritaeniorhynchus_rhabdovirus_RNA_strain_TY:1-11190

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

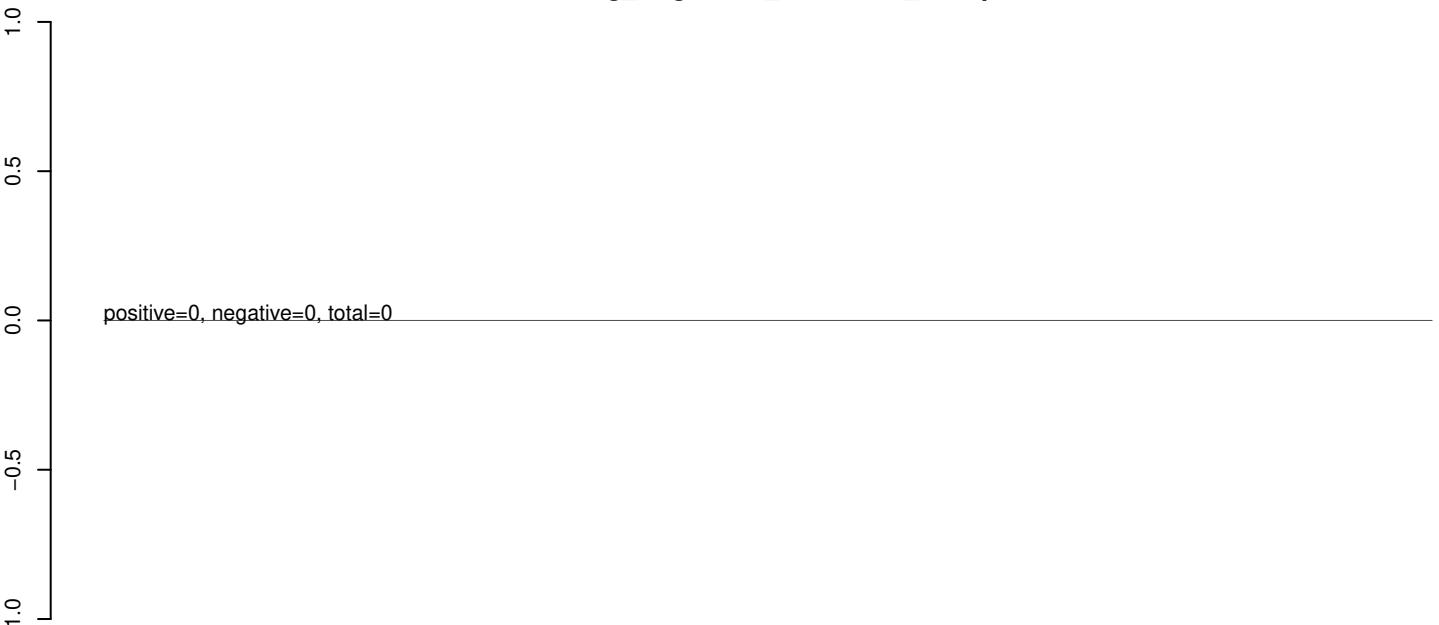
AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

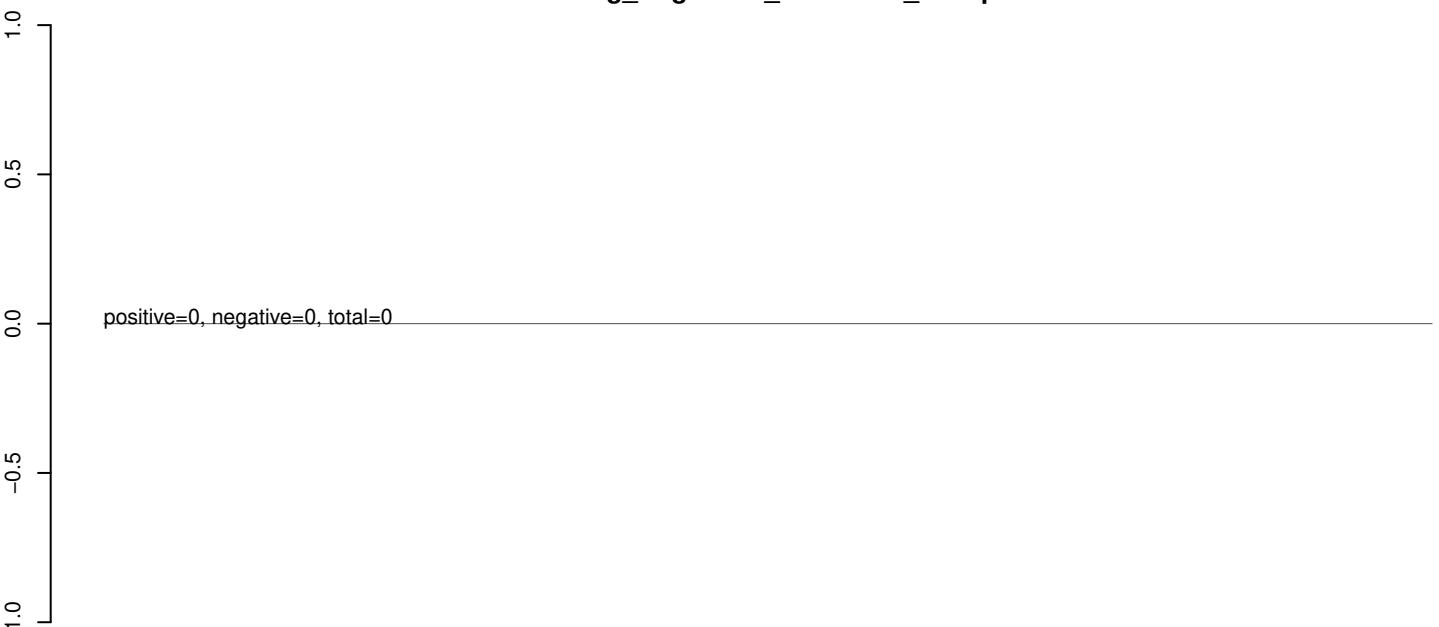
Window size=25, length=11096, virus@NC_025376.1–Grass_carp_rhabdovirus_V76:1–11096

0 2000 4000 6000 8000 10000

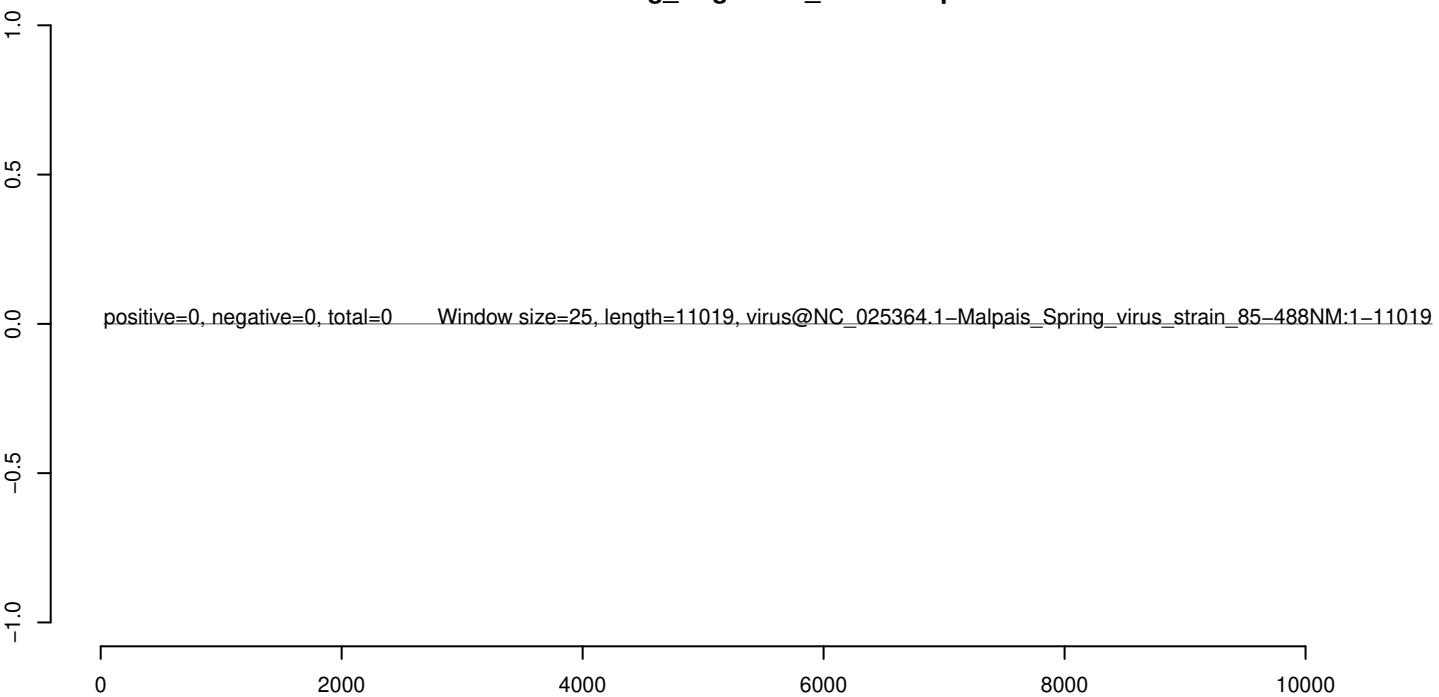
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=Window size=25, length=12240, virus@NC_025362.1-Xiburema_virus_isolate_XIBV_BE_AR_362159:1-12240

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

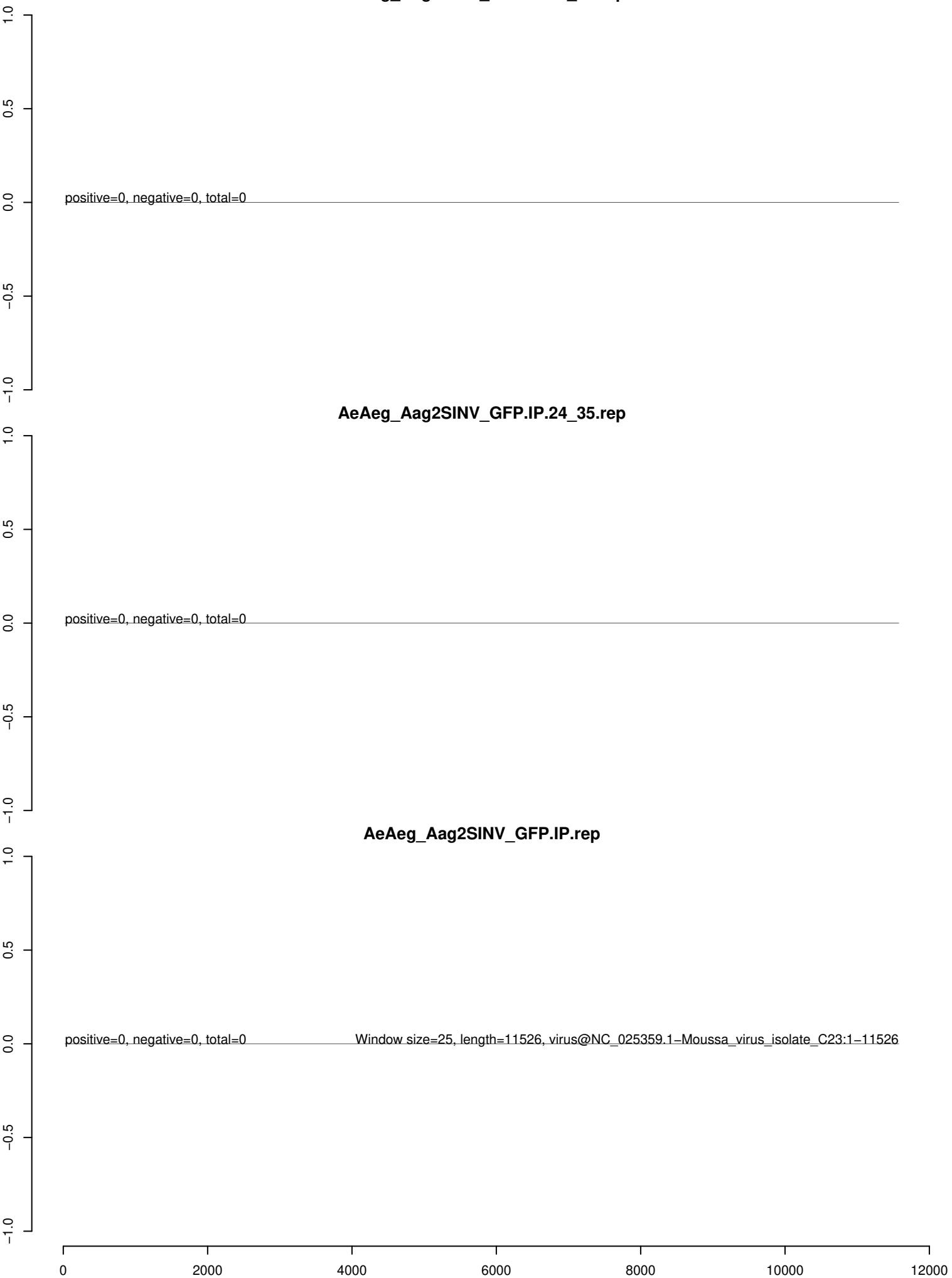
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=11526, virus@NC_025359.1-Moussa_virus_isolate_C23:1-11526



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=11097, virus@NC_025356.1–Pike_fry_rhabdovirus_isolate_F4:1–11097

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

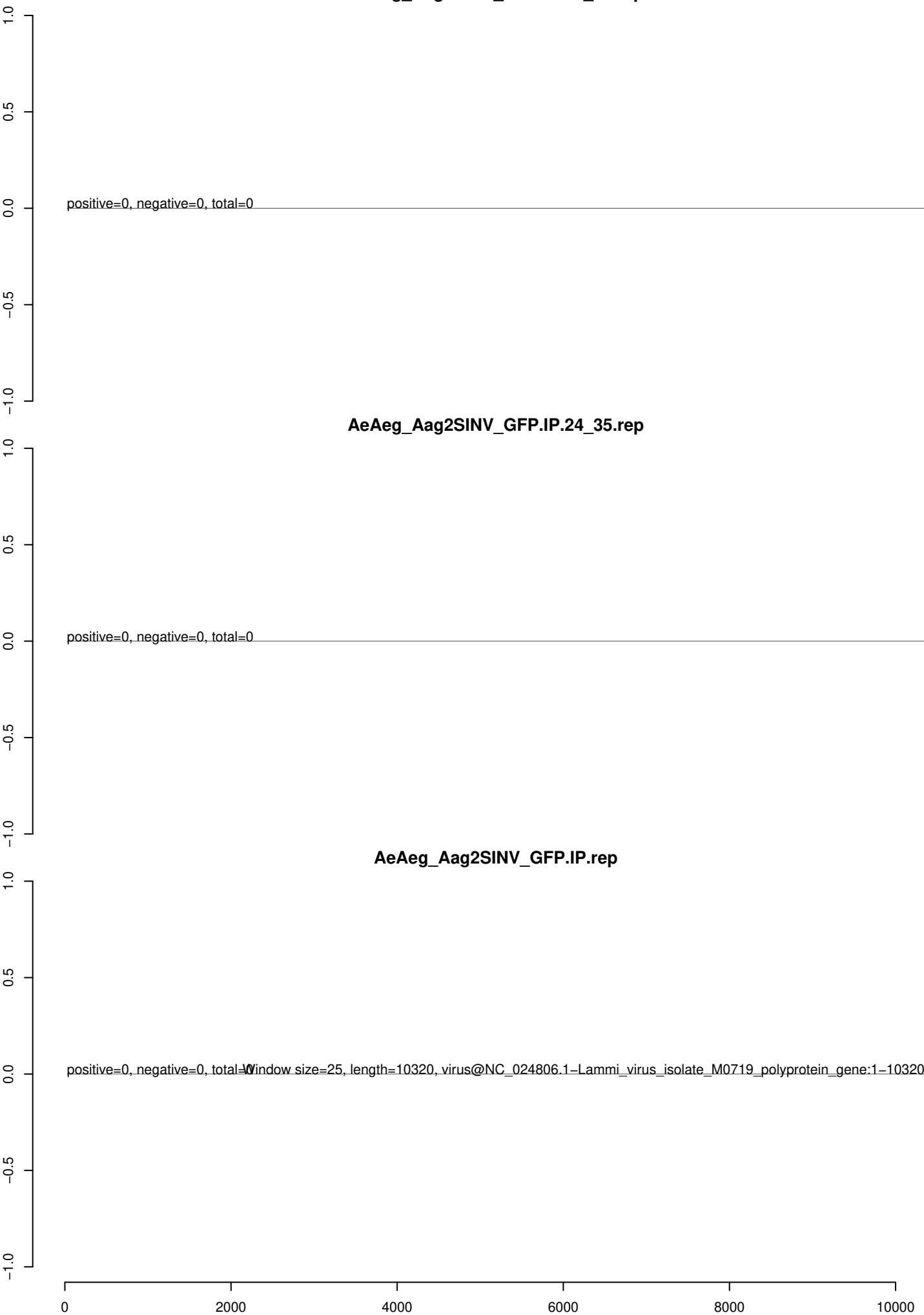
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=Window size=25, length=10320, virus@NC_024806.1-Lammi_virus_isolate_M0719_polyprotein_gene:1-10320



AeAeg_Aag2SINV_GFP.IP.18_23.rep

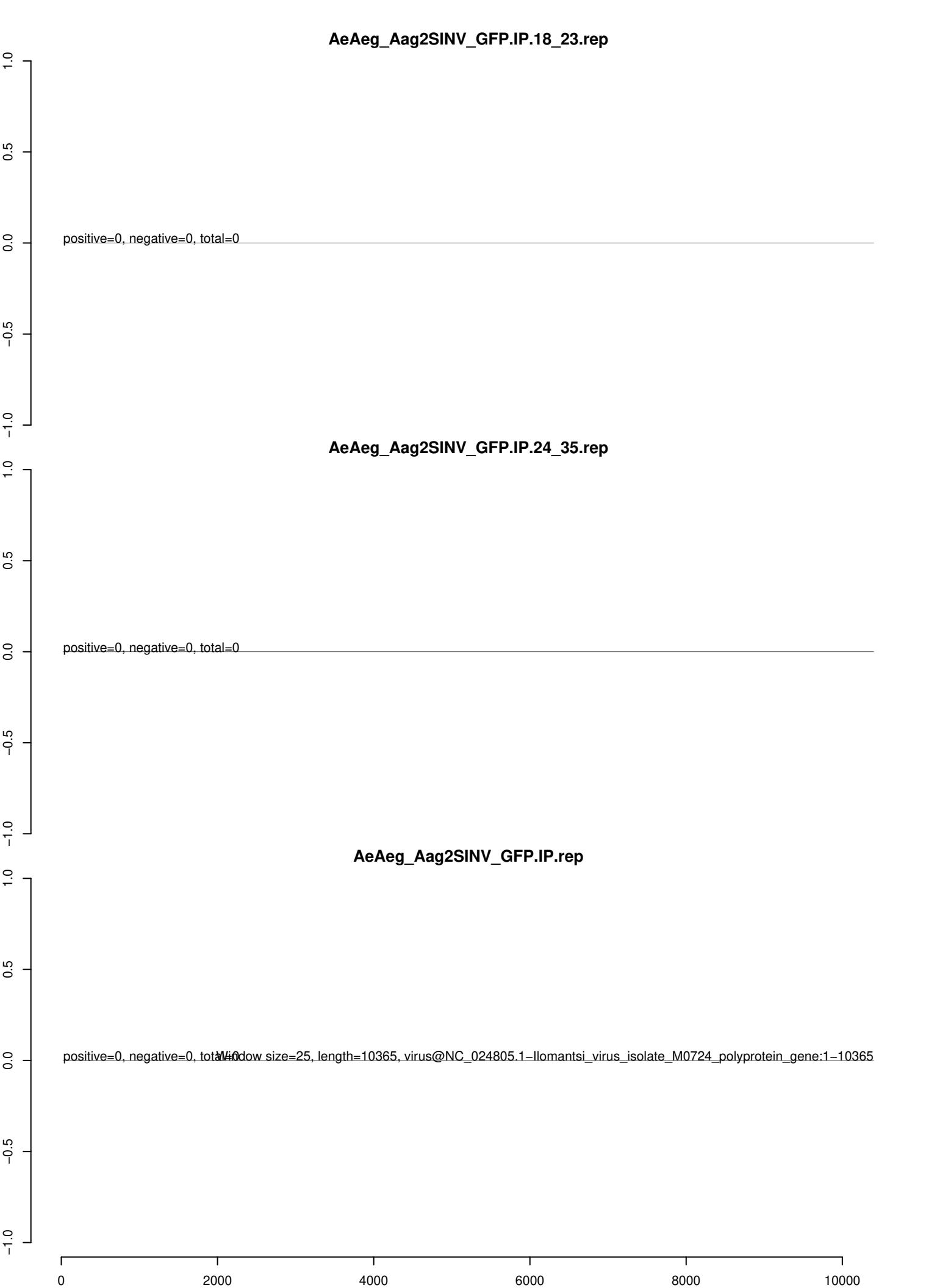
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

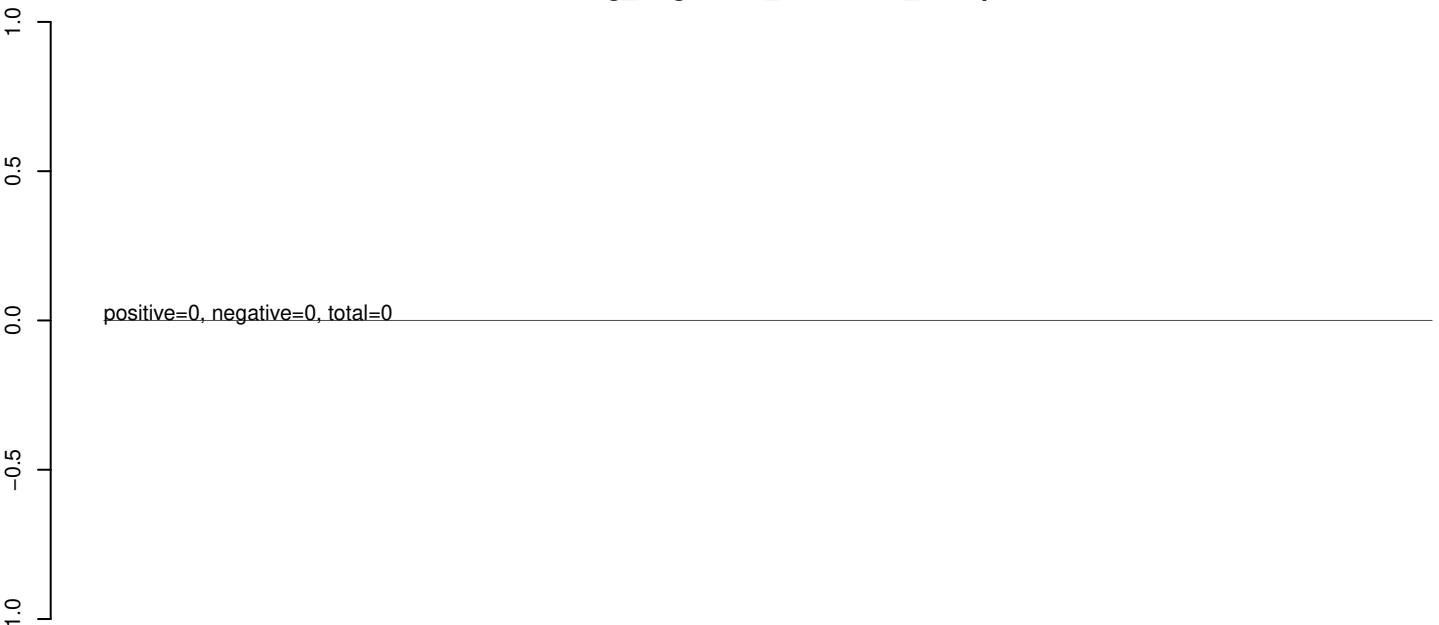
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

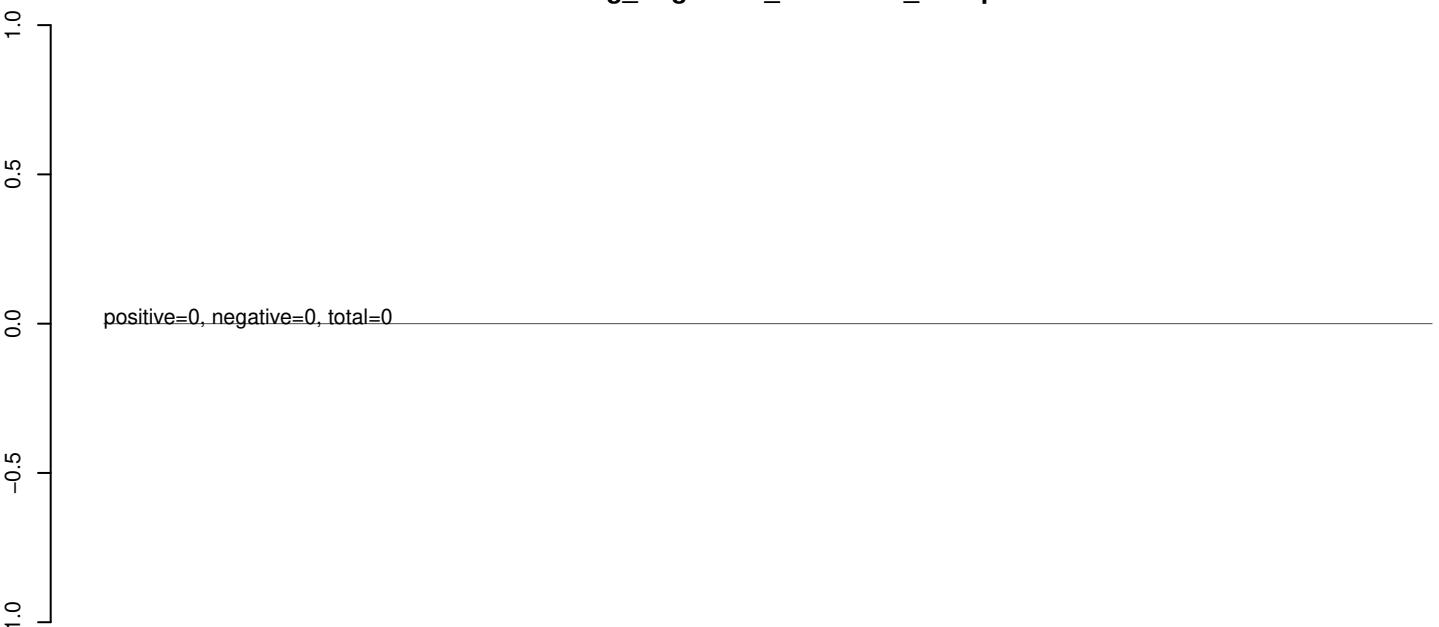
positive=0, negative=0, total=0, window size=25, length=10365, virus@NC_024805.1_llomantsi_virus_isolate_M0724_polyprotein_gene:1-10365



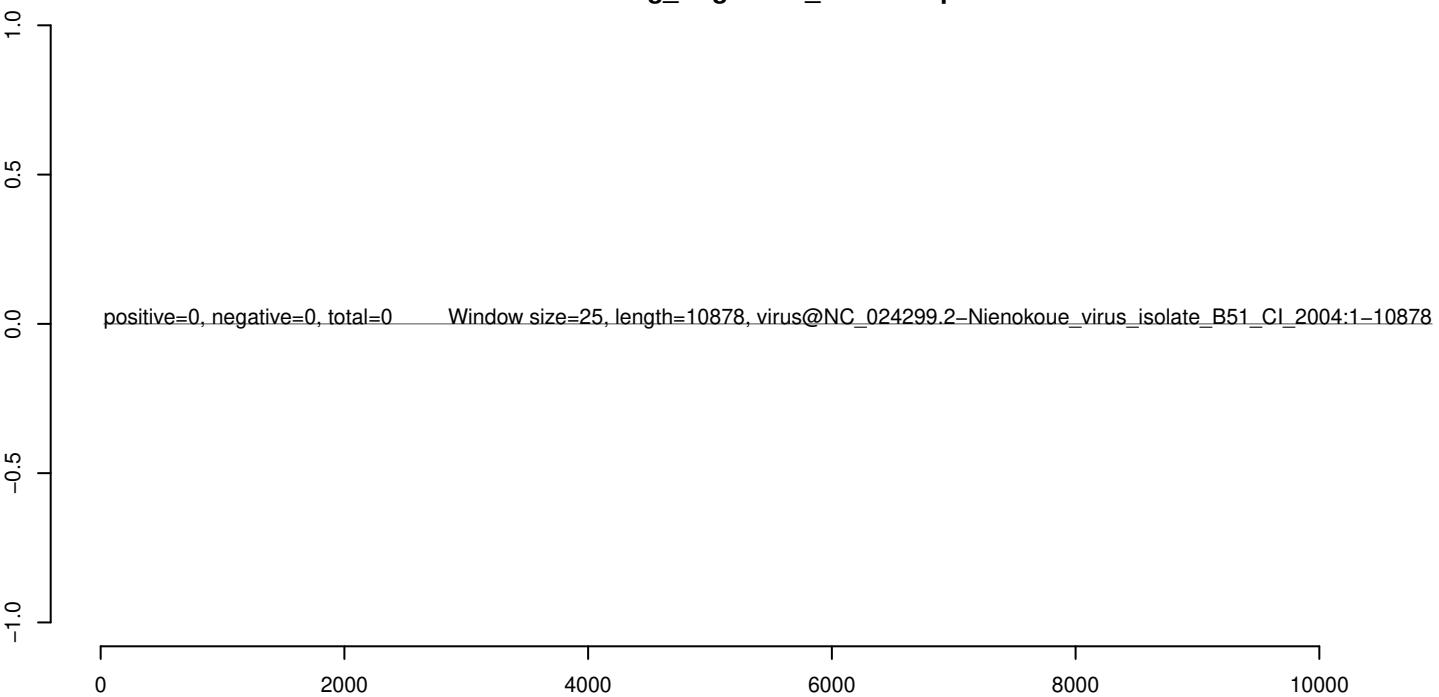
AeAeg_Aag2SINV_GFP.IP.18_23.rep



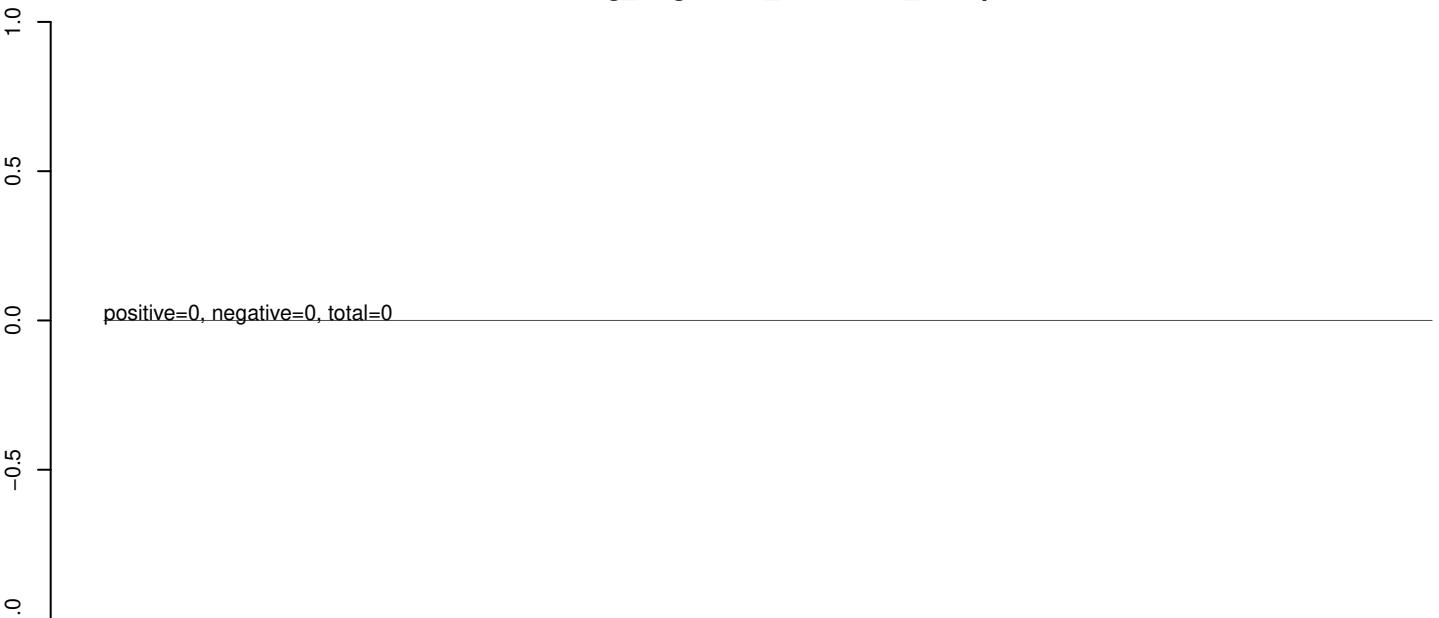
AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



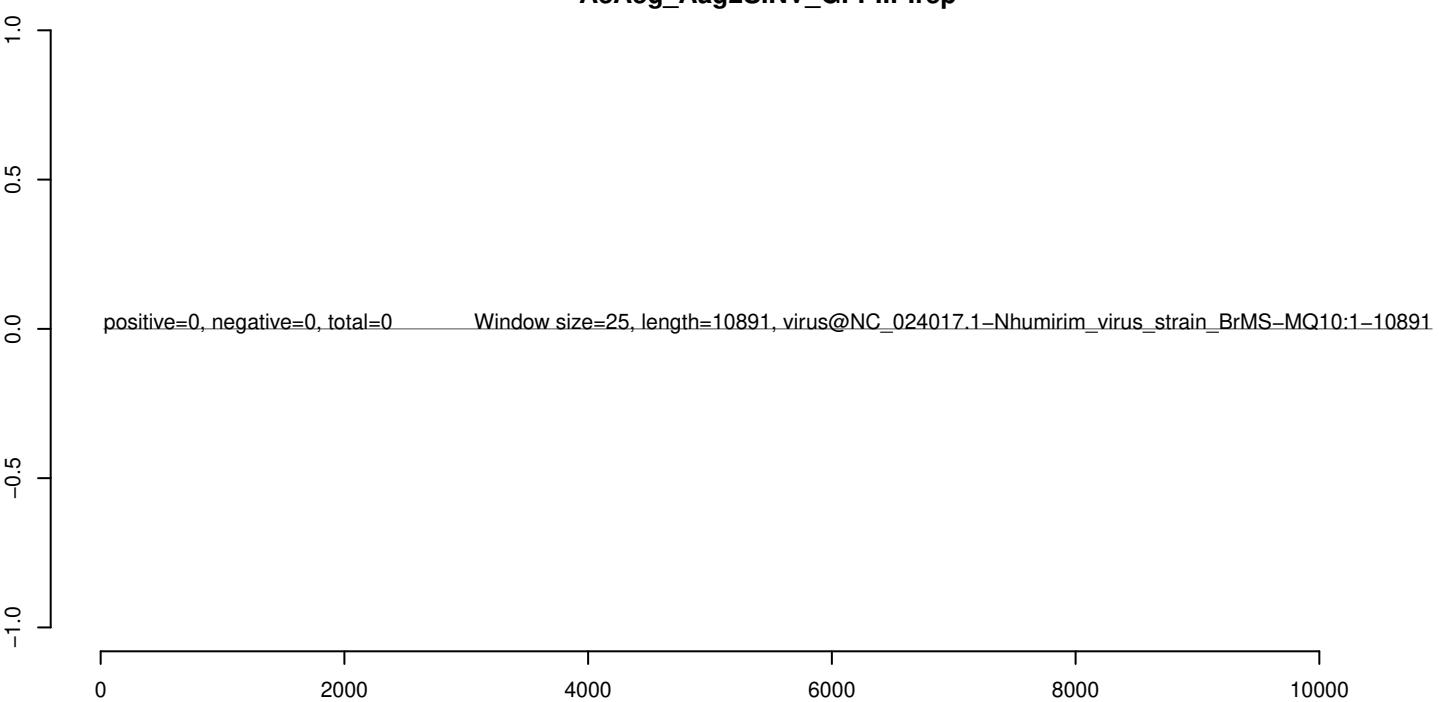
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=25, total=11624, virus@NC_023812.1-Madariaga_virus_strain_MADV_Cebus_apella_BRA_BEAN5122_1956:1-11624

AeAeg_Aag2SINV_GFP.IP.18_23.rep

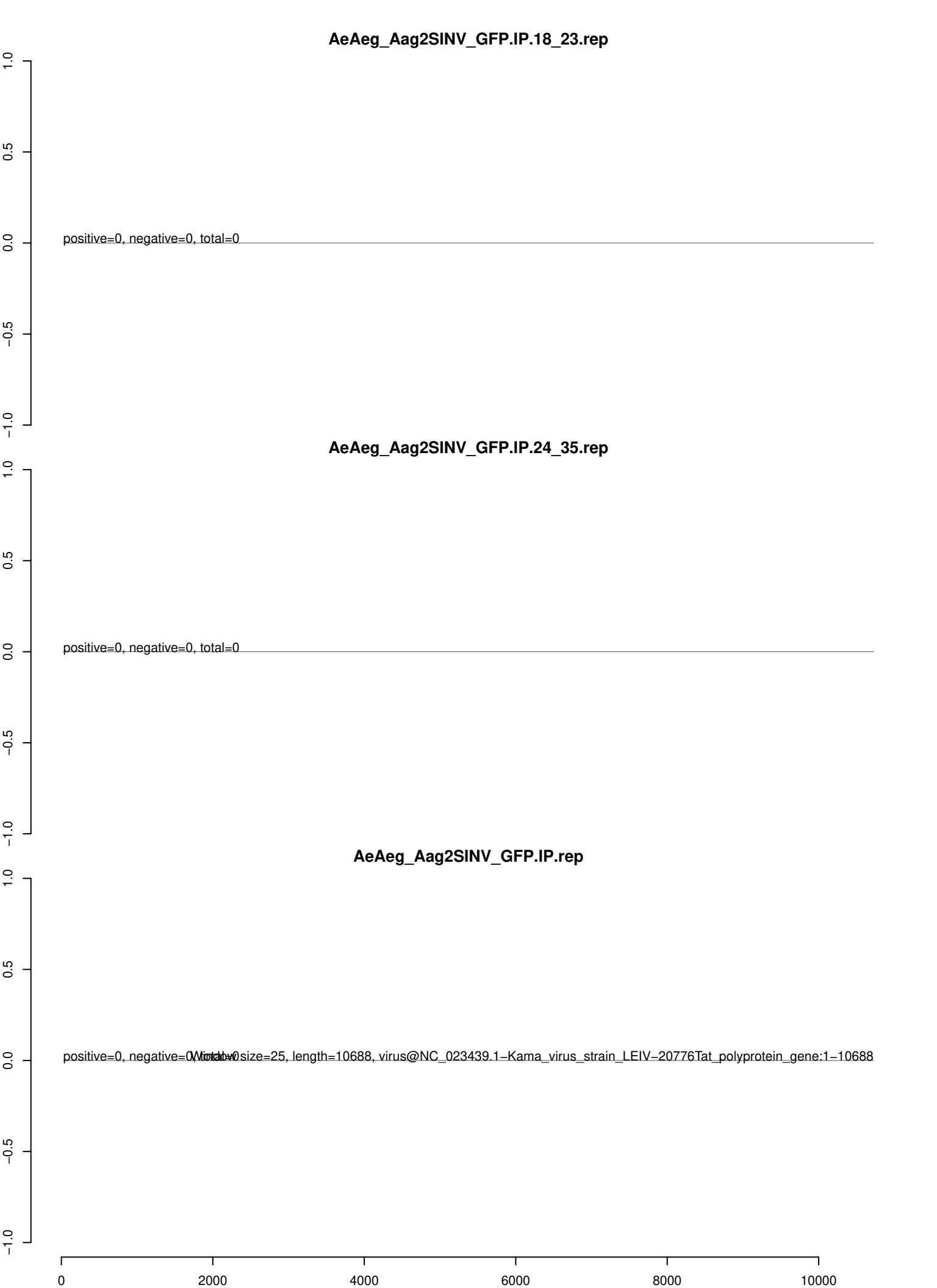
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0 size=25, length=10688, virus@NC_023439.1-Kama_virus_strain_LEIV-20776Tat_polyprotein_gene:1-10688



AeAeg_Aag2SINV_GFP.IP.18_23.rep

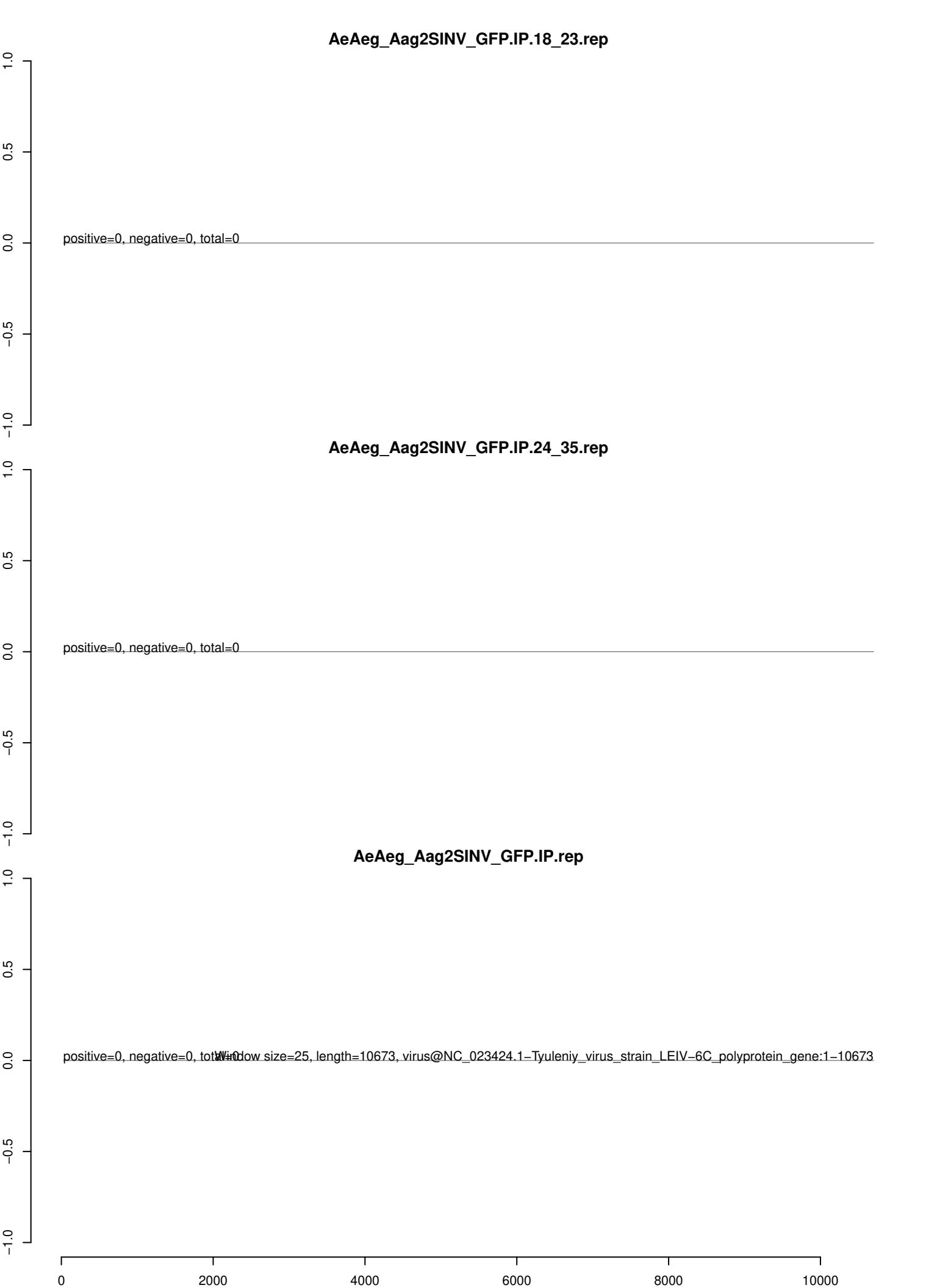
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

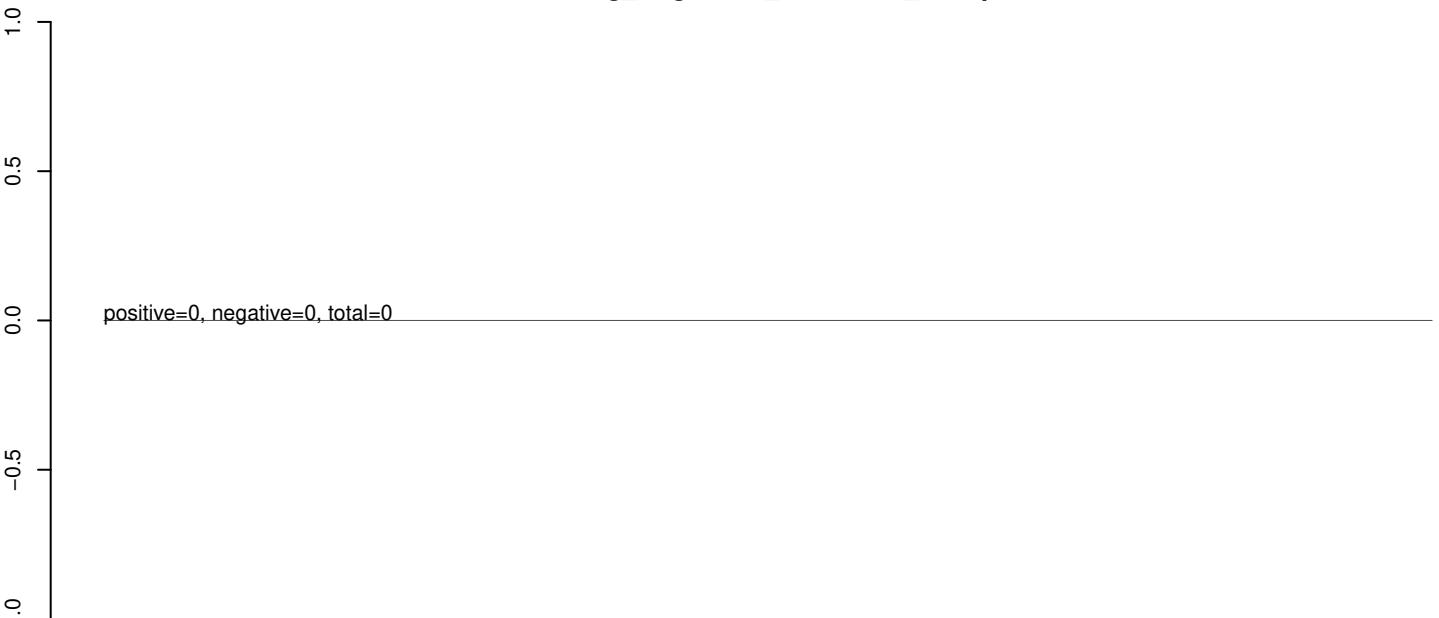
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

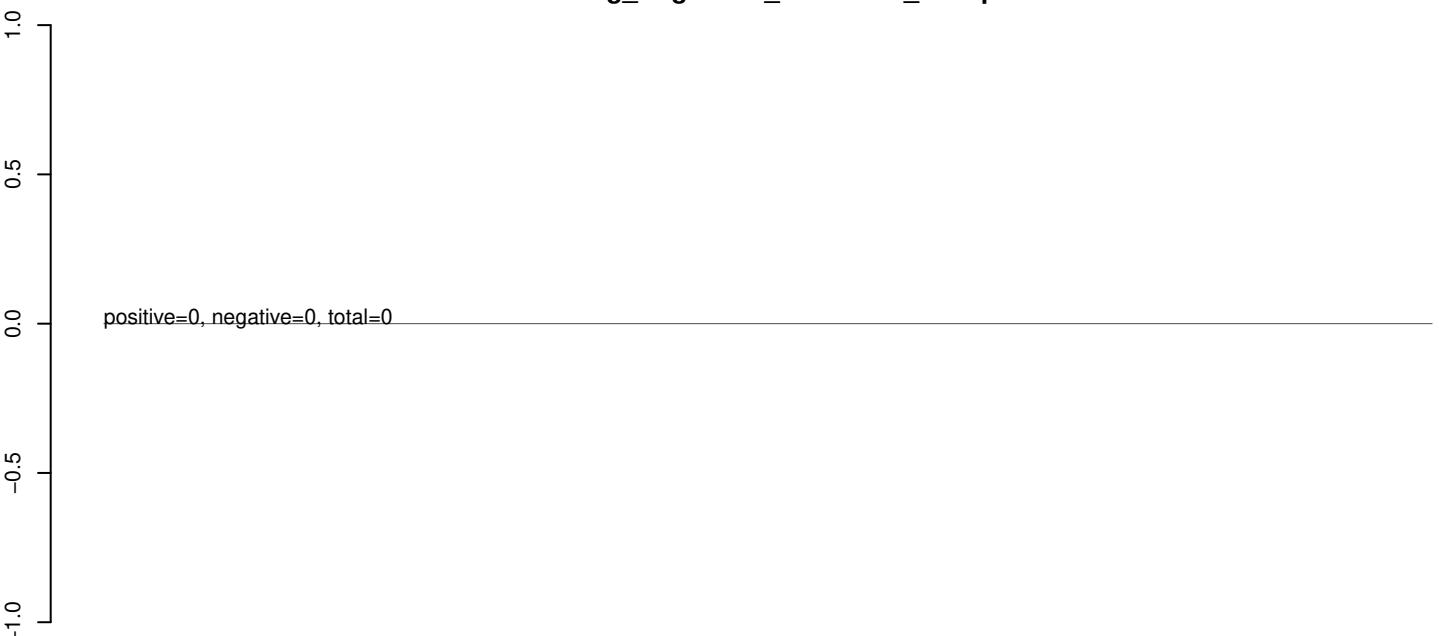
positive=0, negative=0, total=0, window size=25, length=10673, virus@NC_023424.1-Tyuleniy_virus_strain_LEIV-6C_polyprotein_gene:1-10673



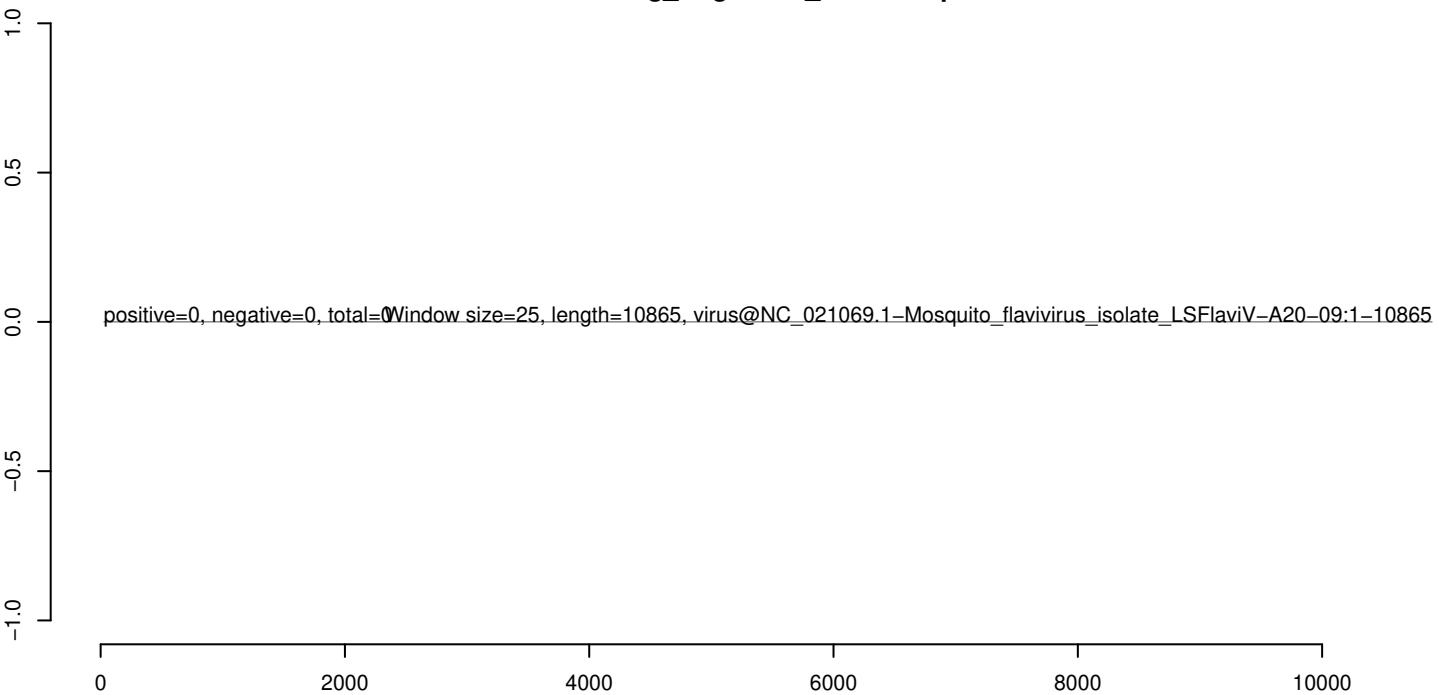
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

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

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window.size=25, length=10943, virus@NC_018705.3-Ntaya_virus:1-10943

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

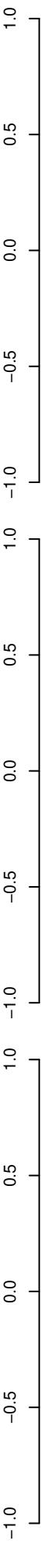
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

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



0

5000

10000

15000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

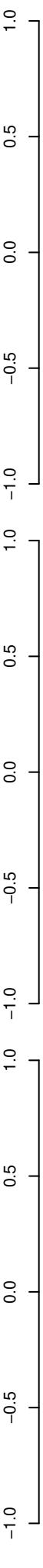
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

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



5000

10000

15000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window.size=25, length=10733, virus@NC_017086.1-Chaoyang_virus:1-10733

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=11688, virus@NC_016959.1-Ndumu_virus:1-11688

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=25, total=10990, virus@NC_015843.2_Tembusu_virus_flavivirus_polyprotein_flavivirus_polyprotein_gene_gene:1-10990

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

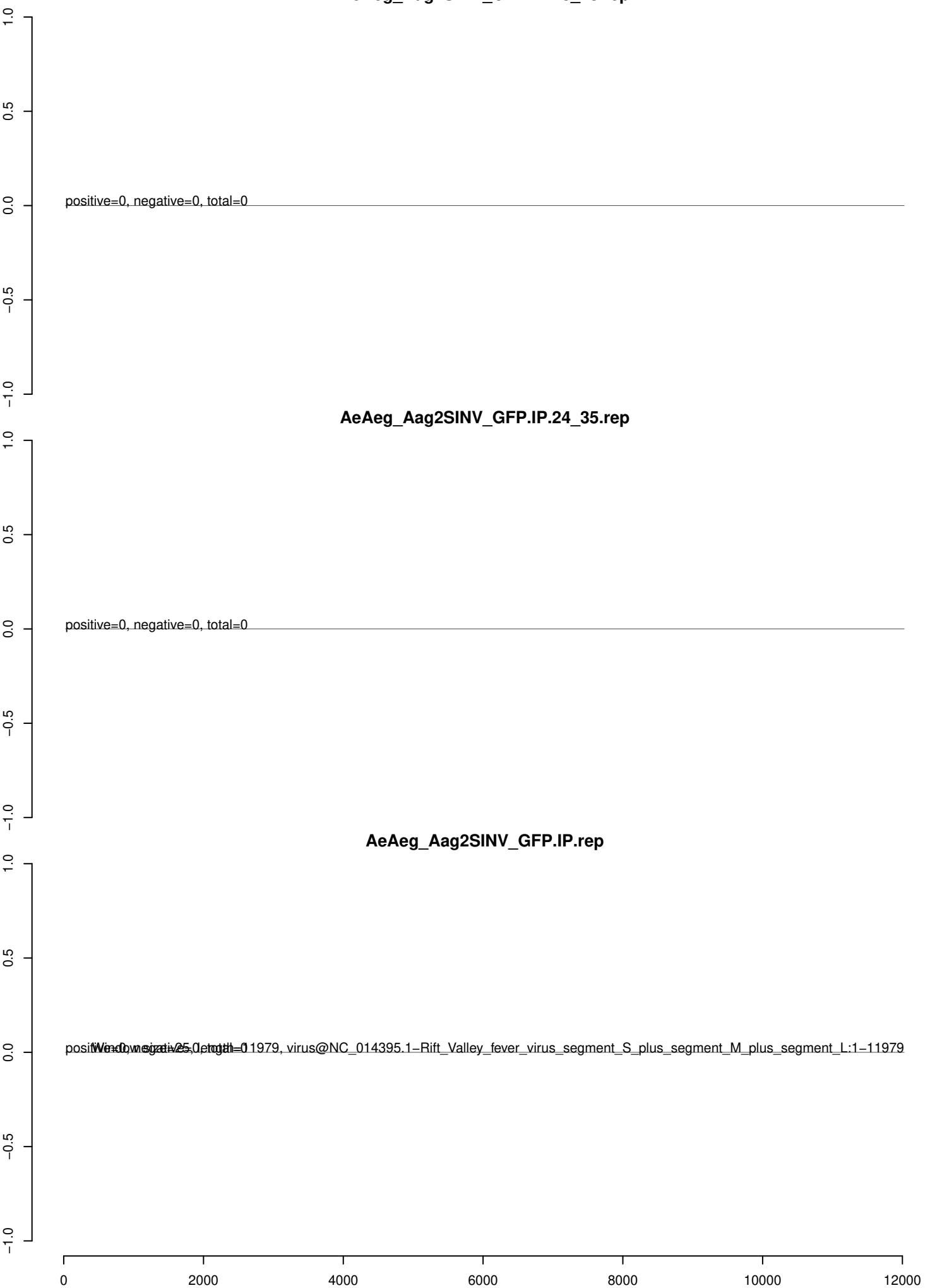
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=25, total=25, date=01/09/1979, virus@NC_014395.1-Rift_Valley_fever_virus_segment_S_plus_segment_M_plus_segment_L:1-11979



AeAeg_Aag2SINV_GFP.IP.18_23.rep

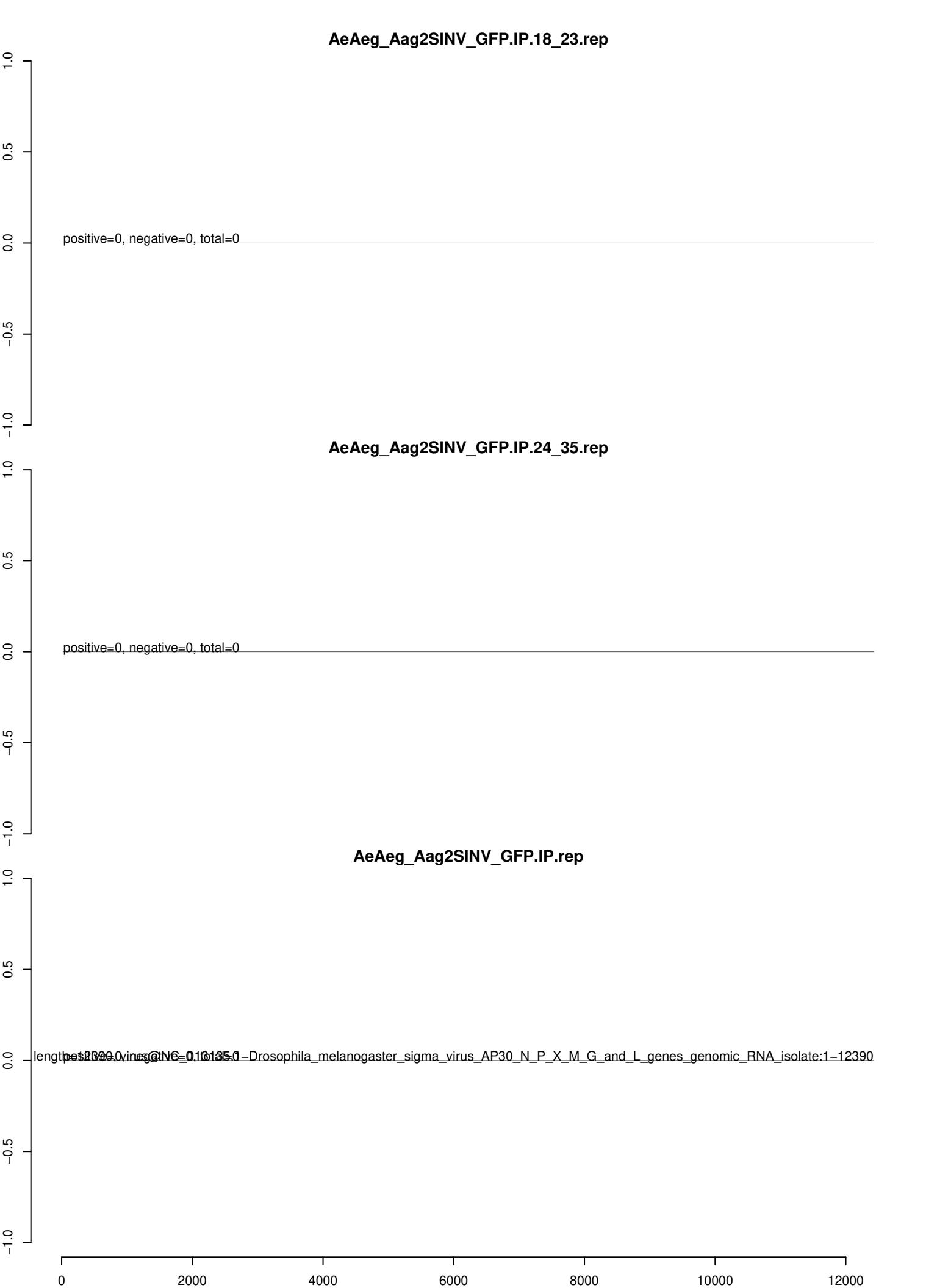
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

lengths@12390,virus@N6=0,1@12390-Drosophila_melanogaster_sigma_virus_AP30_N_P_X_M_G_and_L_genes_genomic_RNA_isolate:1-12390



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=4806, virus@NC_012958.1–Drosophila_A_virus:1–4806

0 1000 2000 3000 4000 5000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10814, virus@NC_012735.1-Wesselsbron_virus:1-10814

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10865, virus@NC_012671.1–Quang_Binh_virus:1–10865

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10941, virus@NC_012534.1-Bagaza_virus:1-10941

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window.size=25, length=10723, virus@NC_012533.1-Kedougou_virus:1-10723

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, width=25, length=10794, virus@NC_012532.1-Zika_virus_isolate_ZIKV_Monkey_Uganda_MR766_1947:1-10794

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=11029, virus@NC_009942.1-West_Nile_virus_lineage_1:1-11029

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

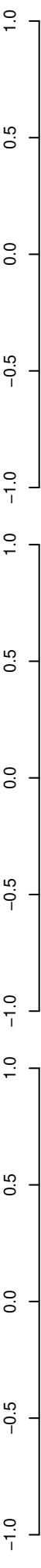
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10755, virus@NC_009028.2-llheus_virus:1-10755



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window.size=25, length=10815, virus@NC_009026.2-Bussuquara_virus:1-10815

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

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

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10510, virus@NC_008718.1-Entebbe_bat_virus:1-10510

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10837, virus@NC_008604.2-Culex_flavivirus:1-10837

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=12333, virus@NC_007919.3-Nora_virus:1-12333

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10940, virus@NC_007580.2-St_Louis_encephalitis_virus:1-10940

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=11440, virus@NC_007020.1-Tupaia_virus:1-11440

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

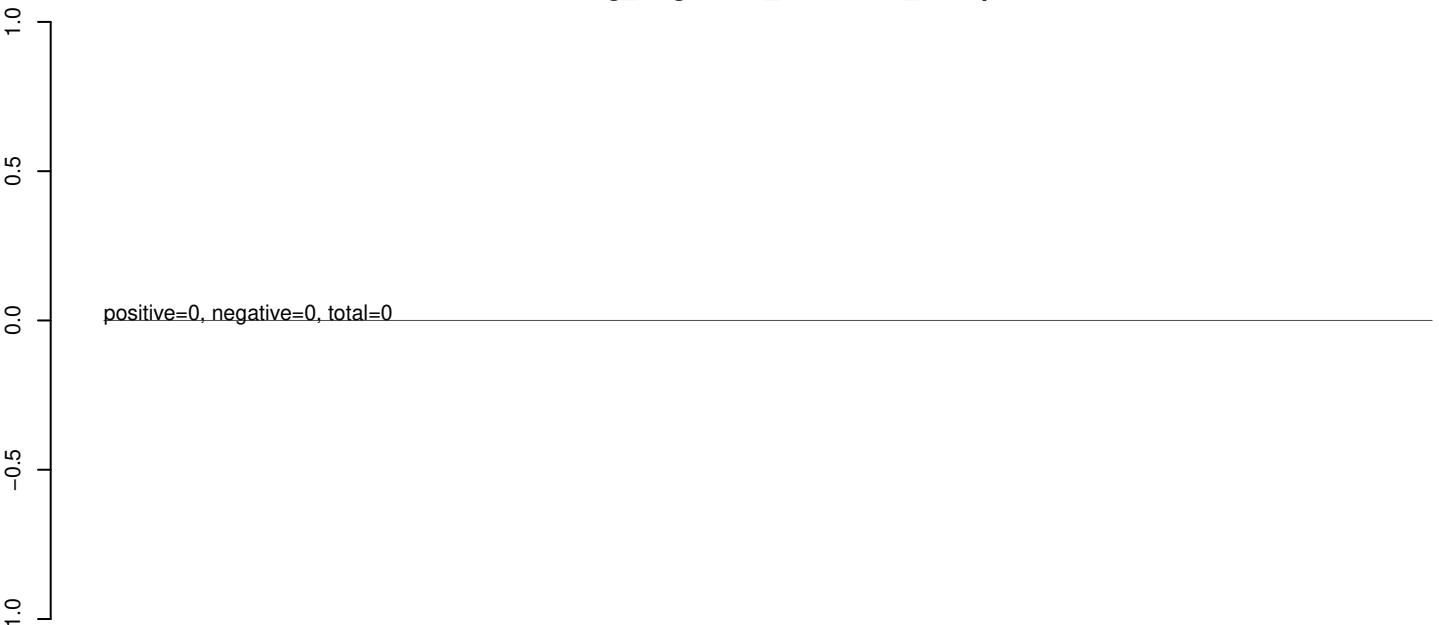
AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

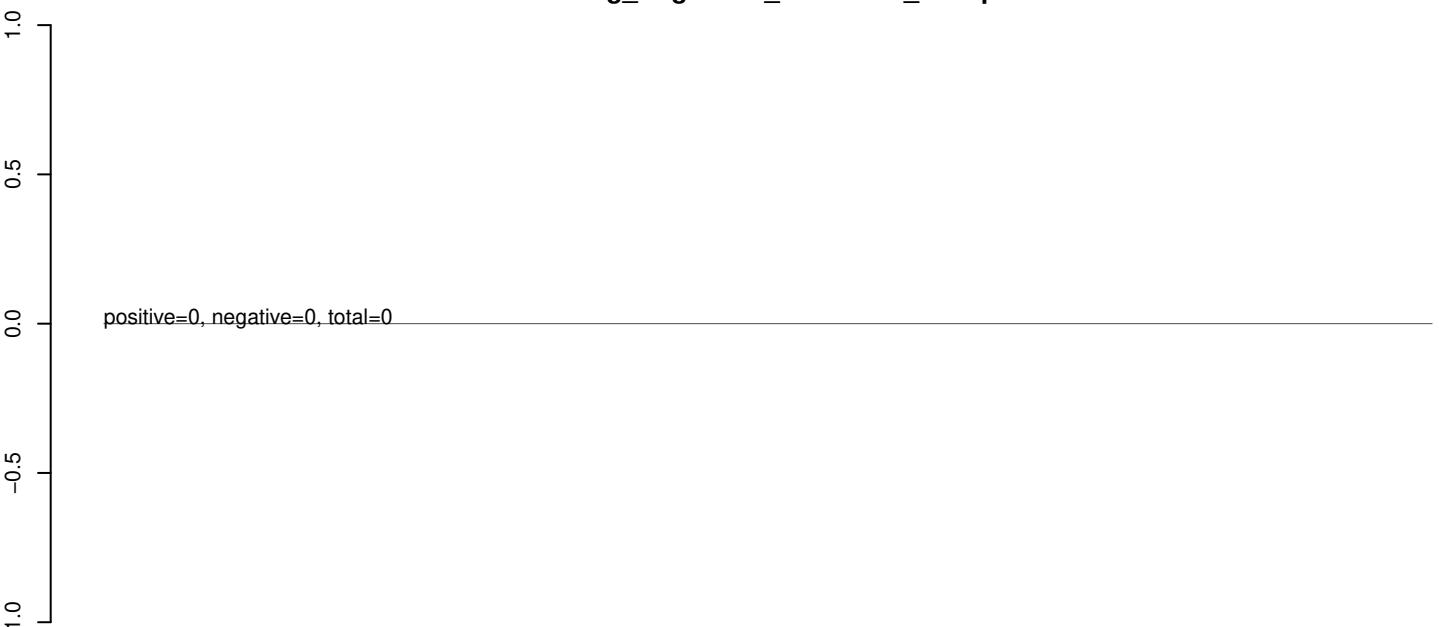
Window size=25, length=10653, virus@NC_006947.1-Karshi_virus:1-10653

0 2000 4000 6000 8000 10000

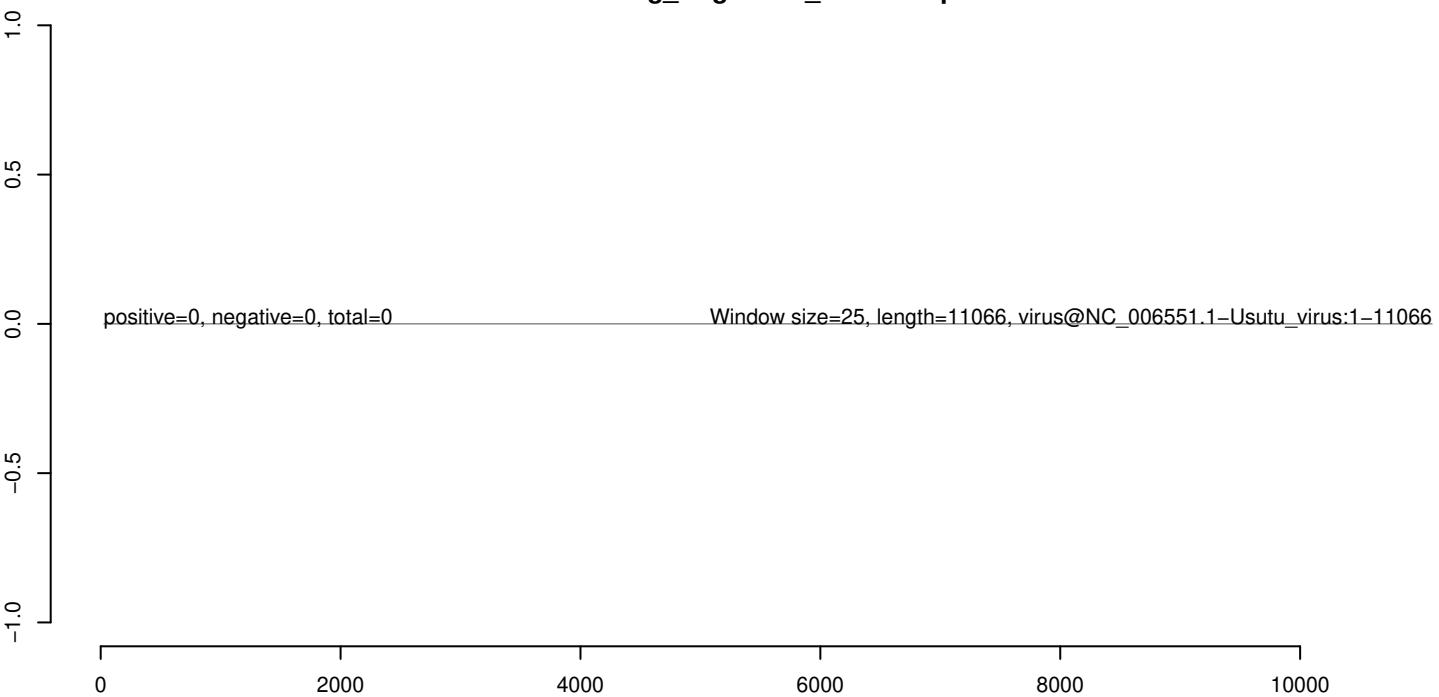
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

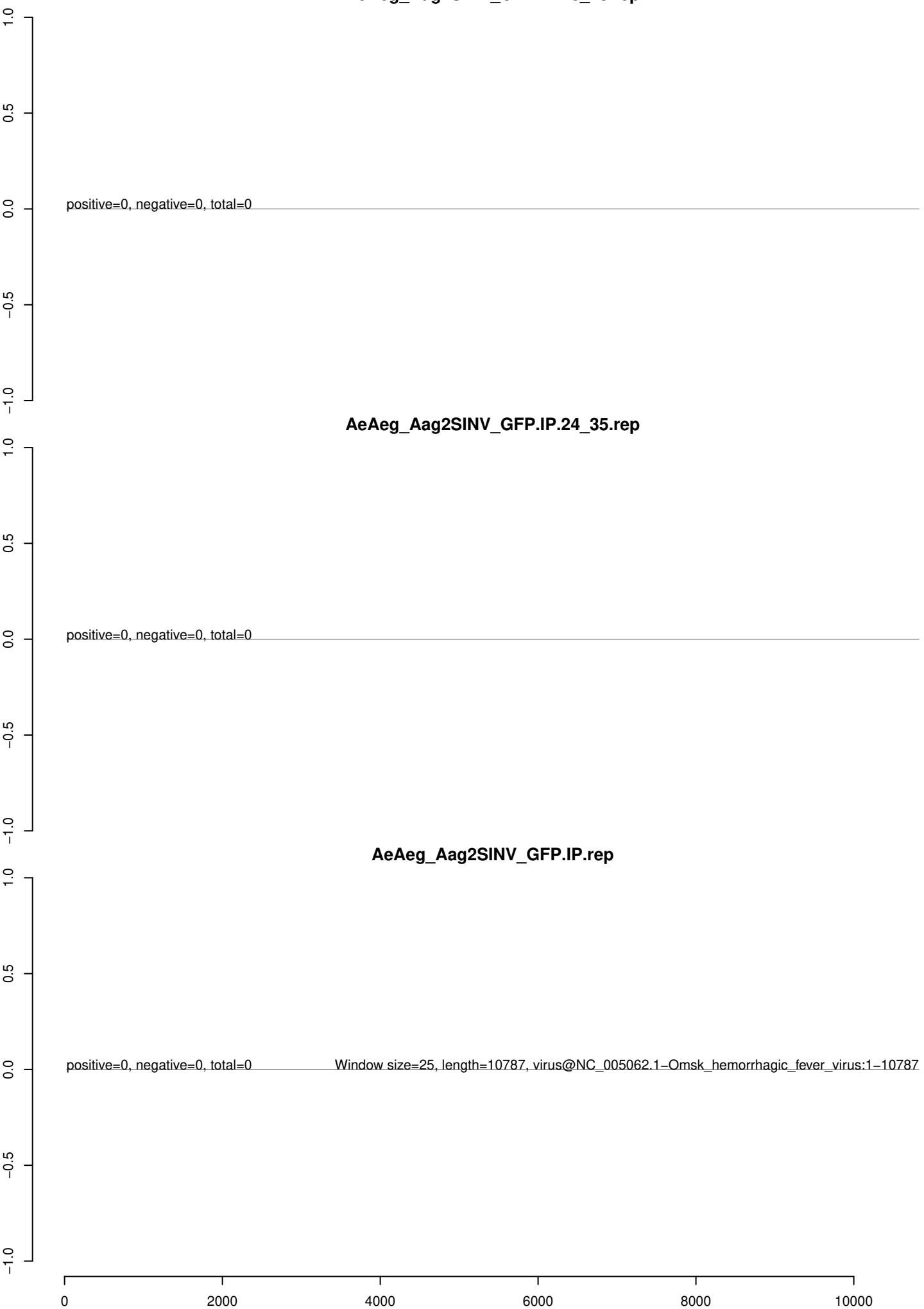
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

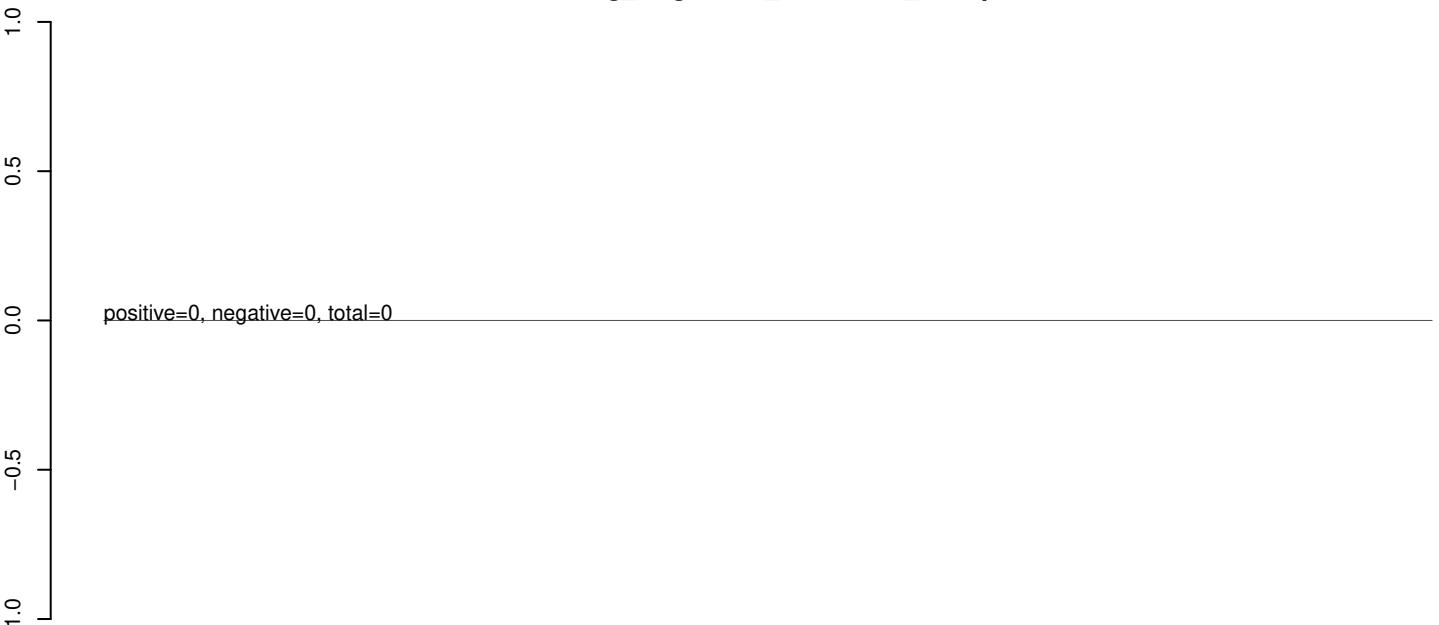
AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

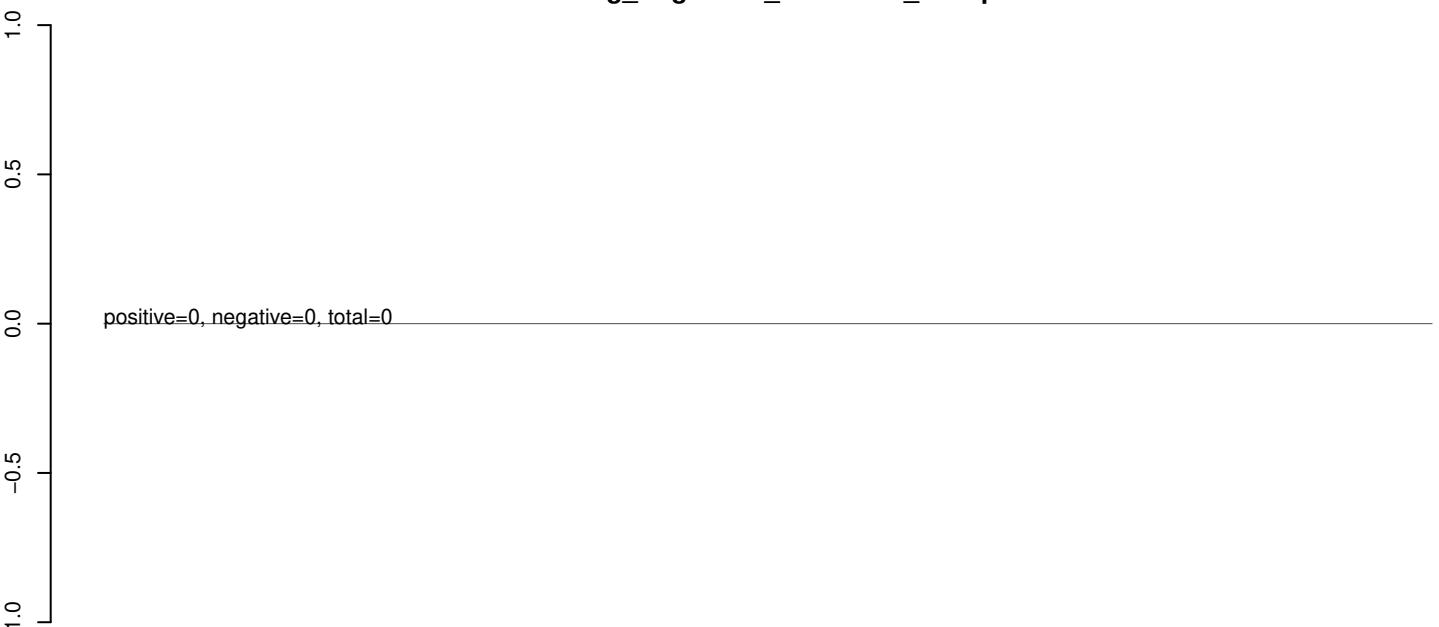
Window size=25, length=10787, virus@NC_005062.1-Omsk_hemorrhagic_fever_virus:1-10787



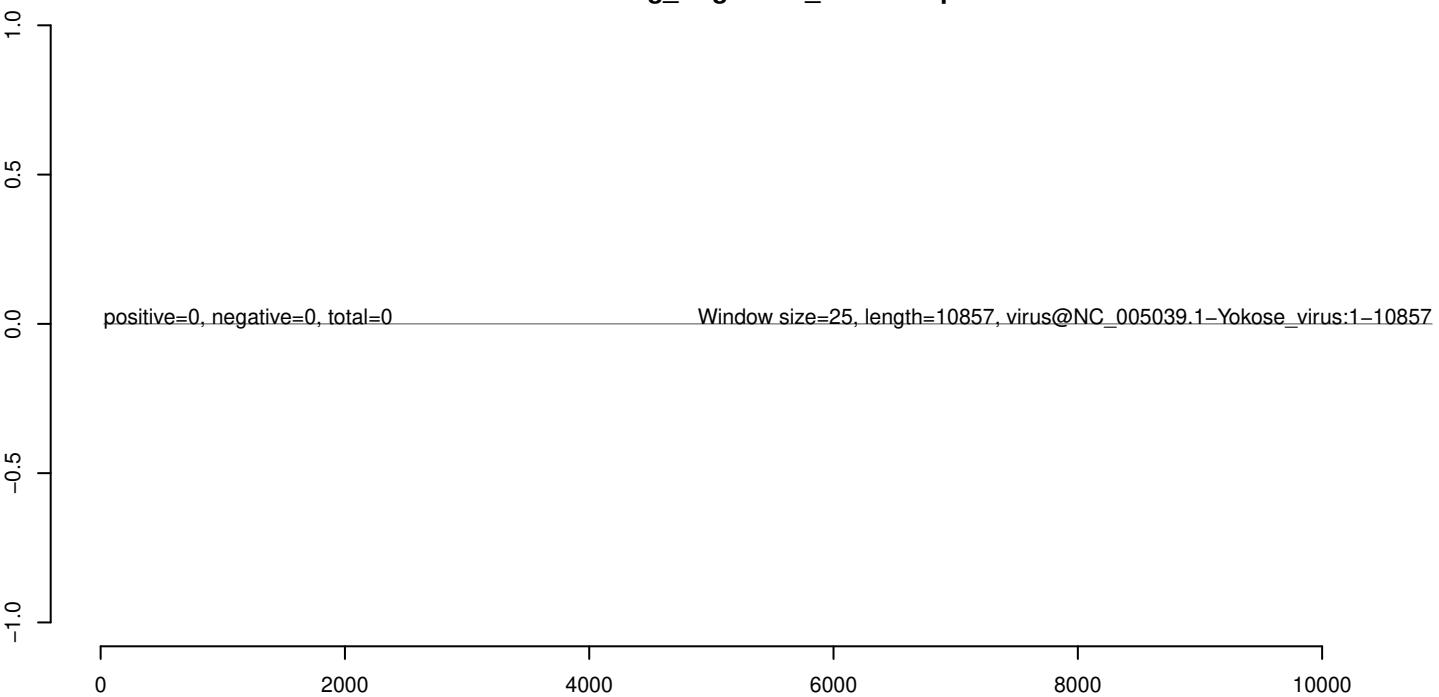
AeAeg_Aag2SINV_GFP.IP.18_23.rep



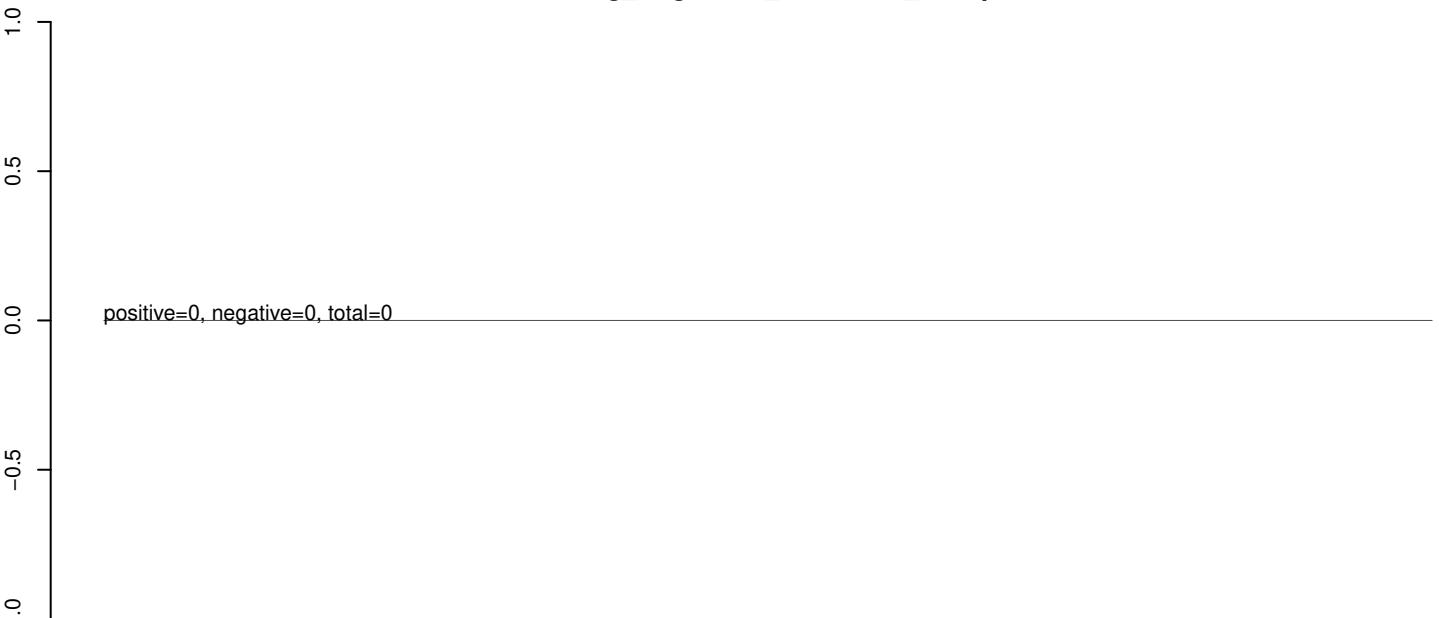
AeAeg_Aag2SINV_GFP.IP.24_35.rep



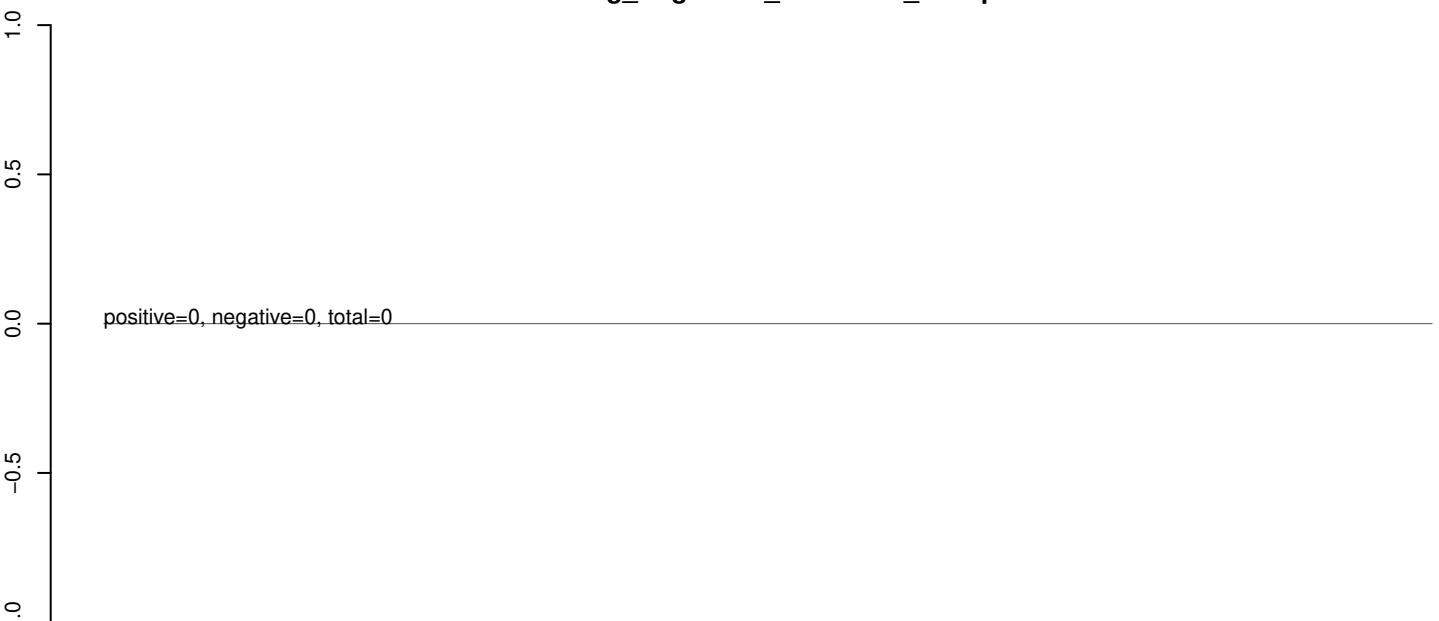
AeAeg_Aag2SINV_GFP.IP.rep



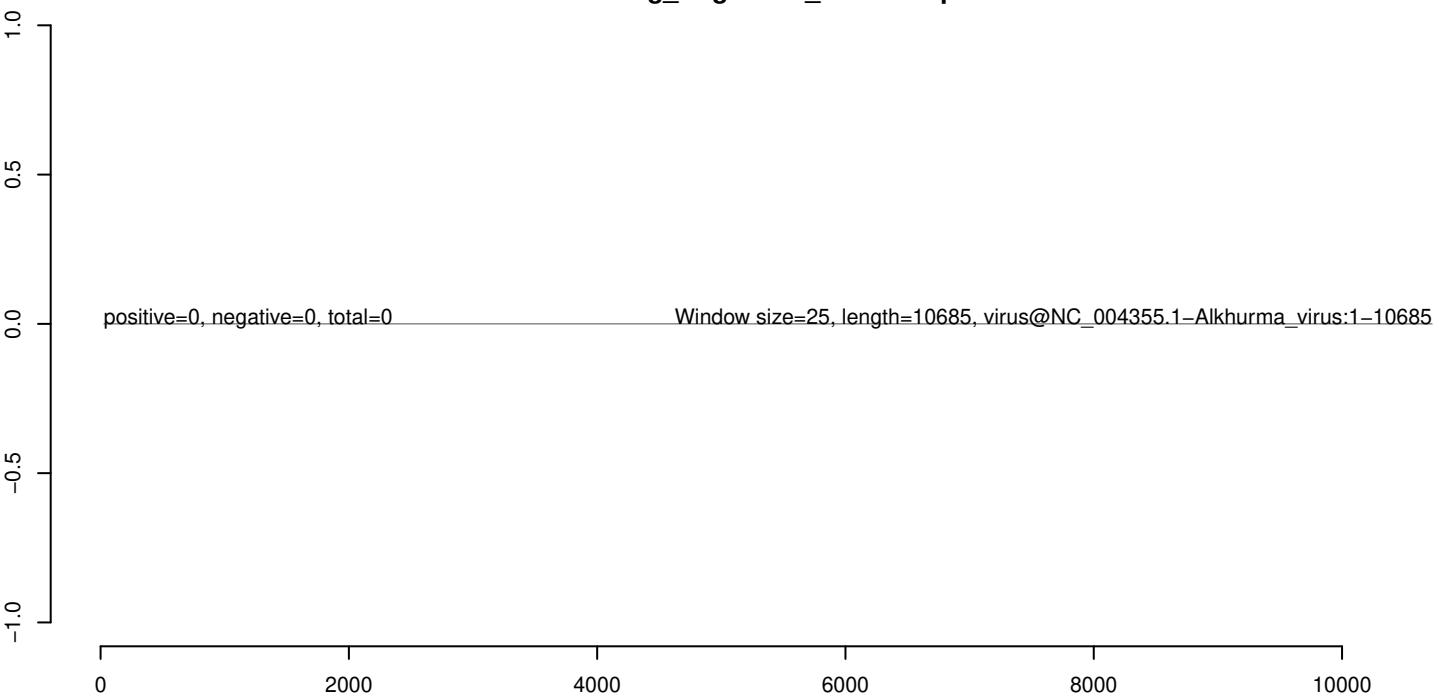
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=4507, virus@NC_004146.1-Flock_house_virus_RNA_1_and_RNA2:1-4507

0 1000 2000 3000 4000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

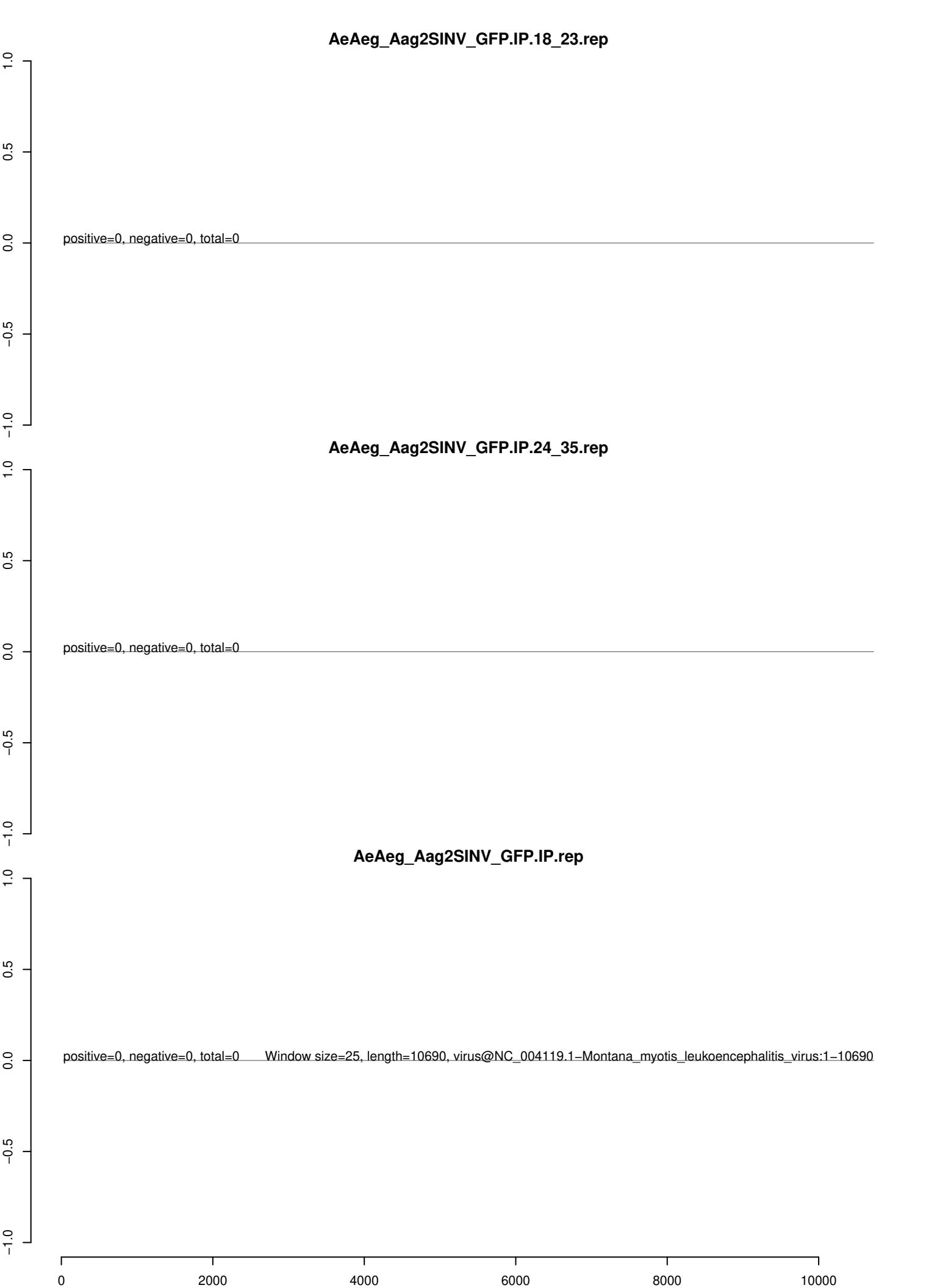
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0 Window size=25, length=10690, virus@NC_004119.1-Montana_myotis_leukoencephalitis_virus:1-10690



AeAeg_Aag2SINV_GFP.IP.18_23.rep

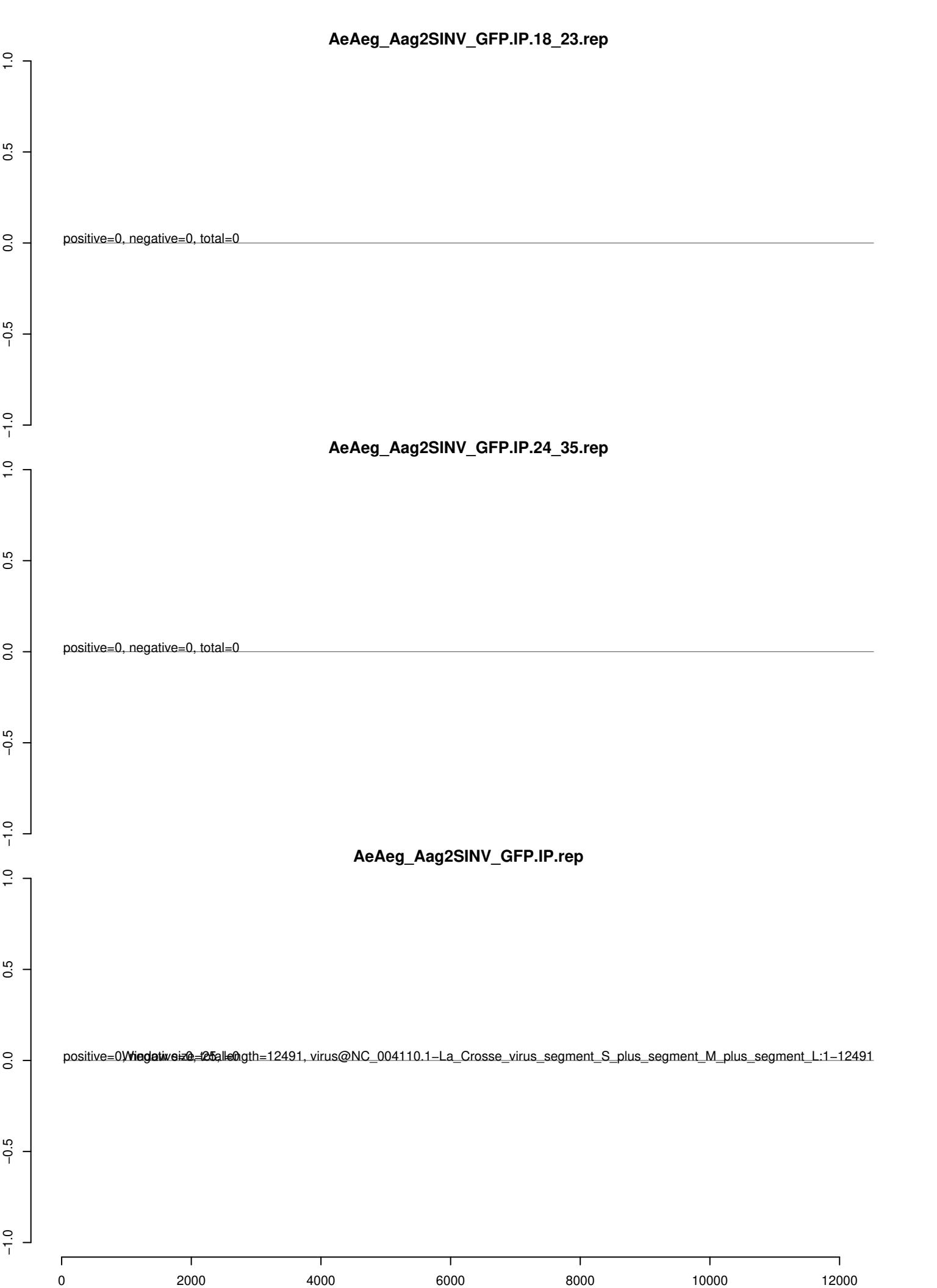
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=25, false=0, length=12491, virus@NC_004110.1-La_Crosse_virus_segment_S_plus_segment_M_plus_segment_L:1-12491



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

Vind positive=25, negative=0, total=0 @ NC_003924.1-Cricket_paralysis_virus_nonstructural_polyprotein_and_structural_polyprotein_genes:1-9185

0 2000 4000 6000 8000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0 Window size=25, length=11484, virus@NC_003908.1-Western_equine_encephalomyelitis_virus:1-11484

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10943, virus@NC_003690.1-Langat_virus:1-10943

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window.size=25, length=10839, virus@NC_003687.1-Powassan_virus:1-10839

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

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

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

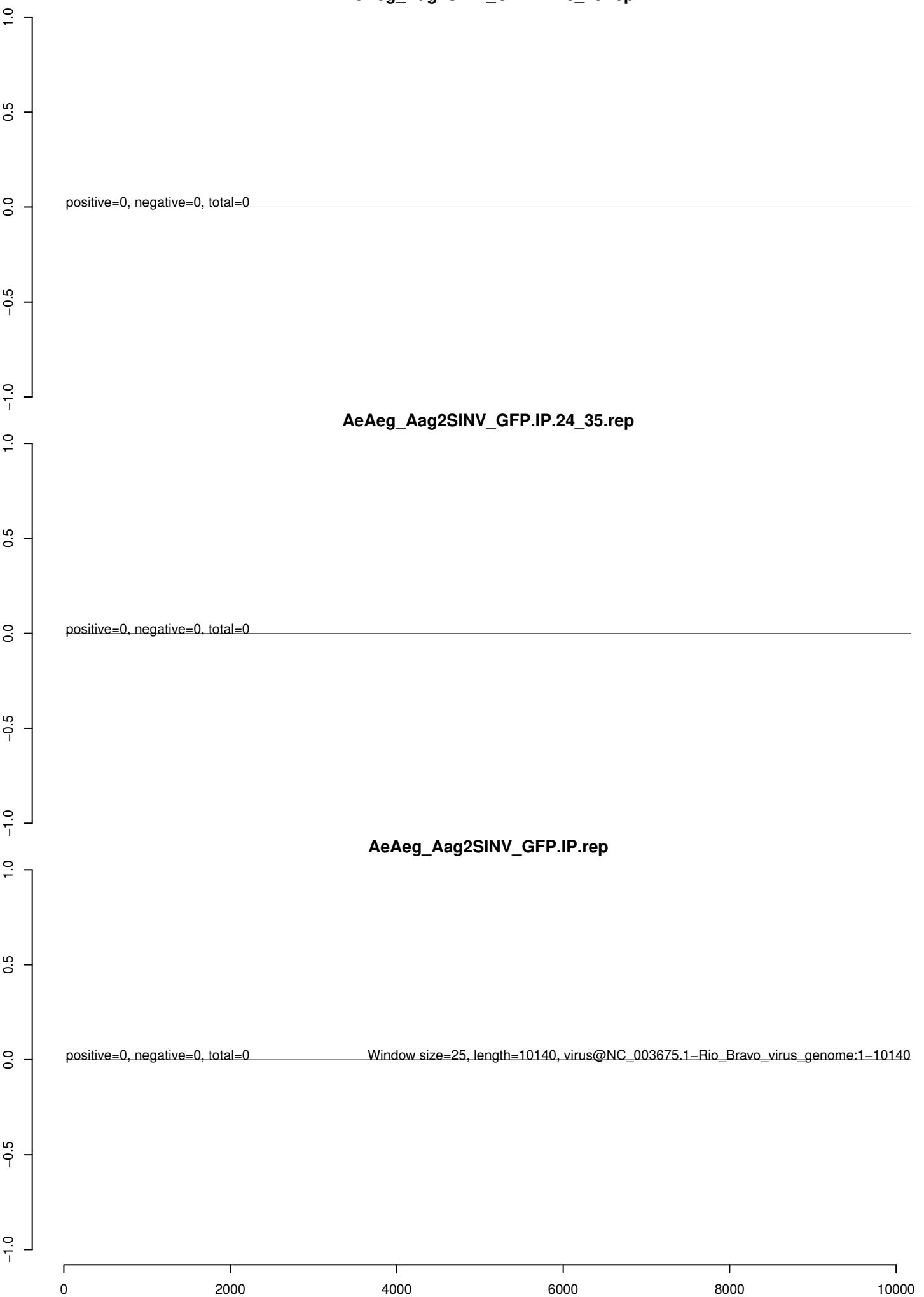
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10140, virus@NC_003675.1–Rio_Bravo_virus_genome:1–10140



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10600, virus@NC_003635.1-Modoc_virus:1-10600

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

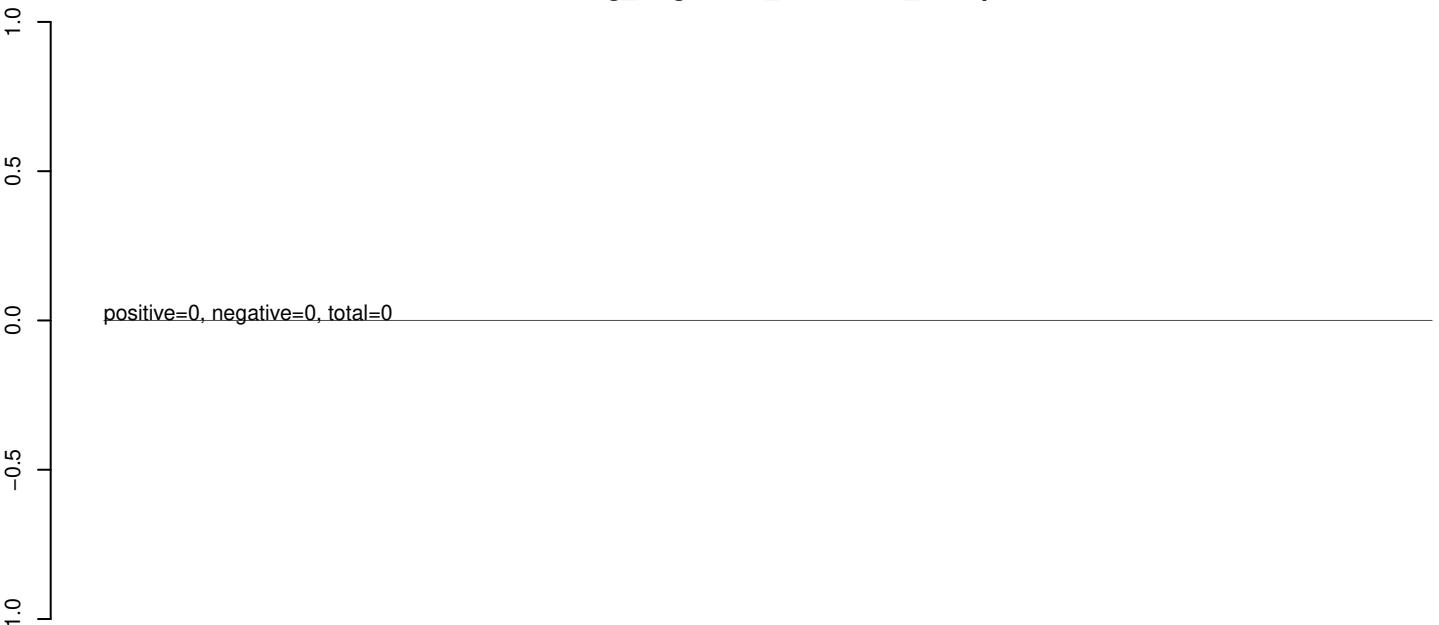
AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

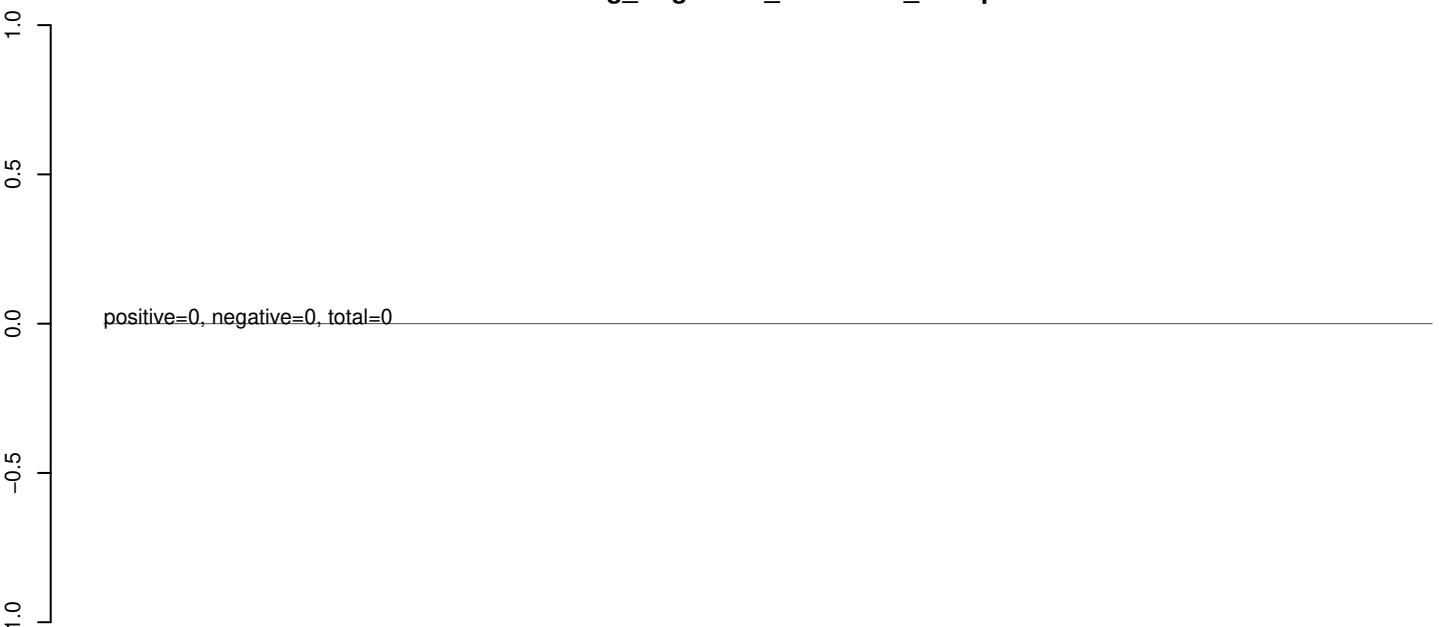
Window size=25, length=11442, virus@NC_003215.1-Semliki_forest_virus:1-11442

0 2000 4000 6000 8000 10000

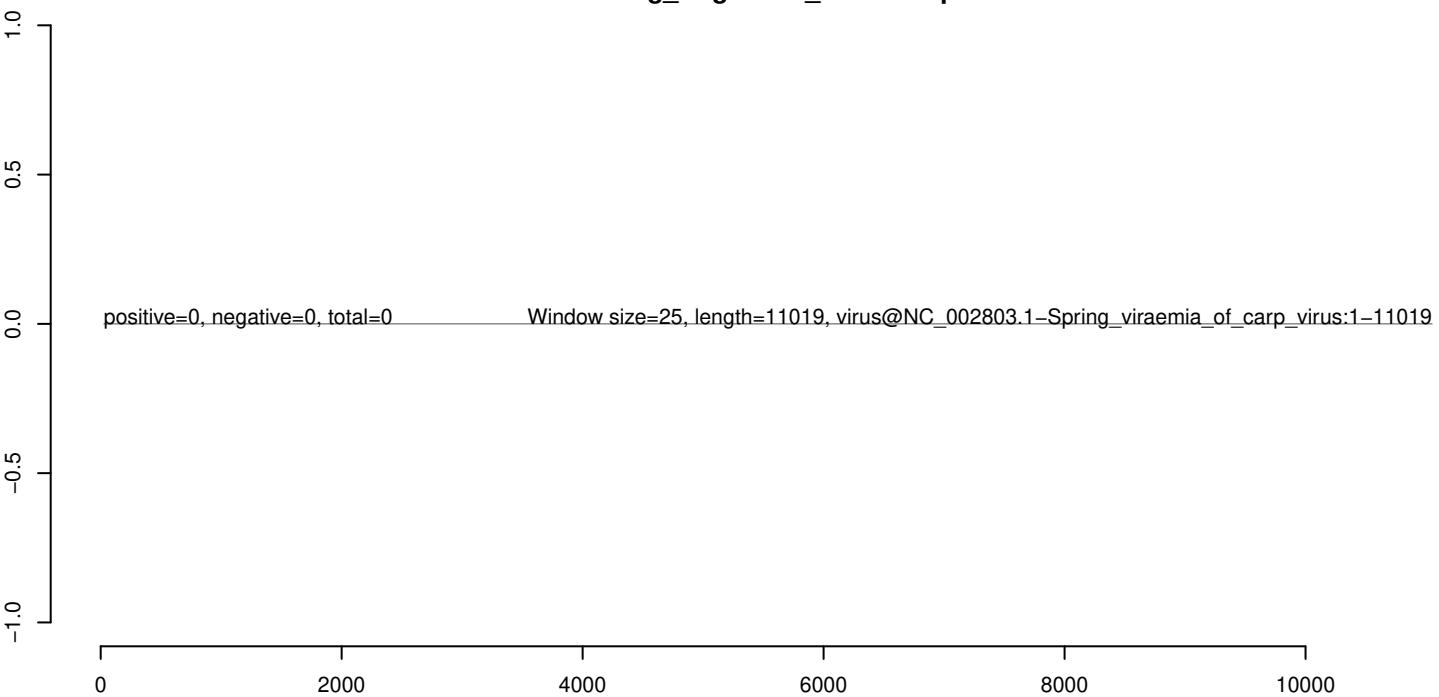
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window.size=25, length=10649, virus@NC_002640.1-Dengue_virus_4:1-10649

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10862, virus@NC_002031.1-Yellow_fever_virus:1-10862

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=25, total=25, virus@NC_001927.1-Bunyamwera_virus_segment_S_plus_segment_M_plus_segment_L:1-12294

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

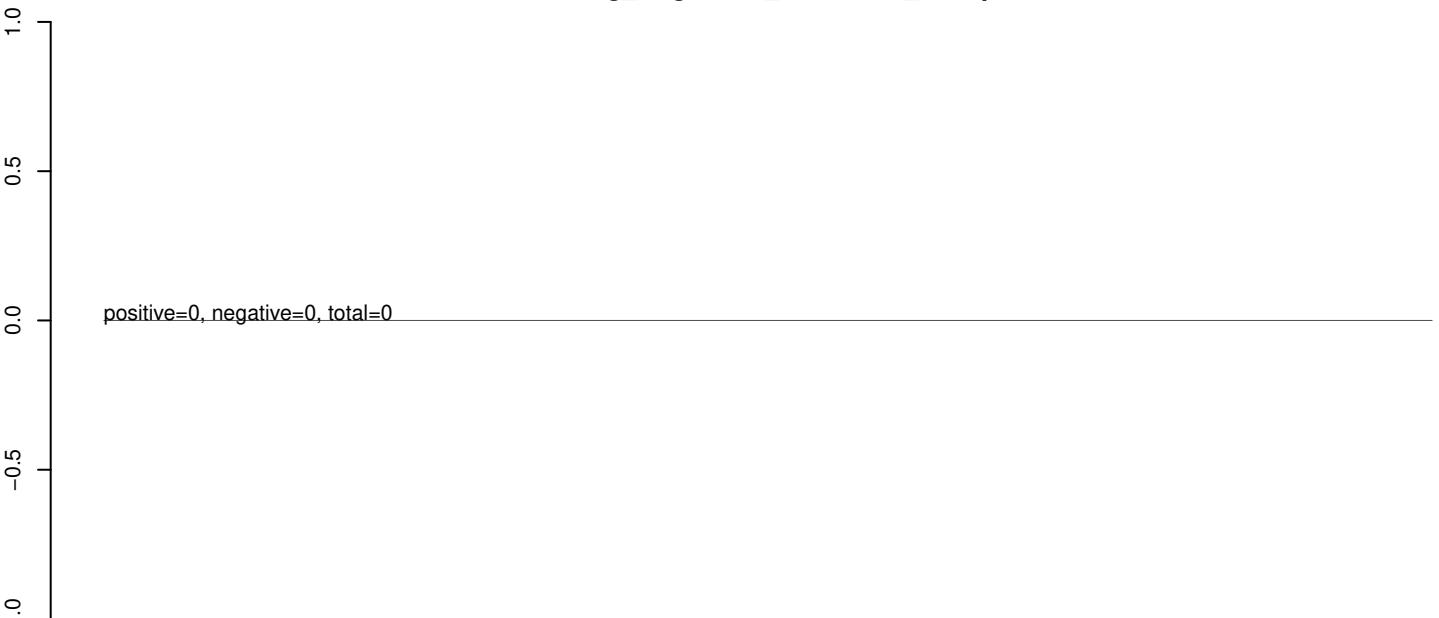
AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

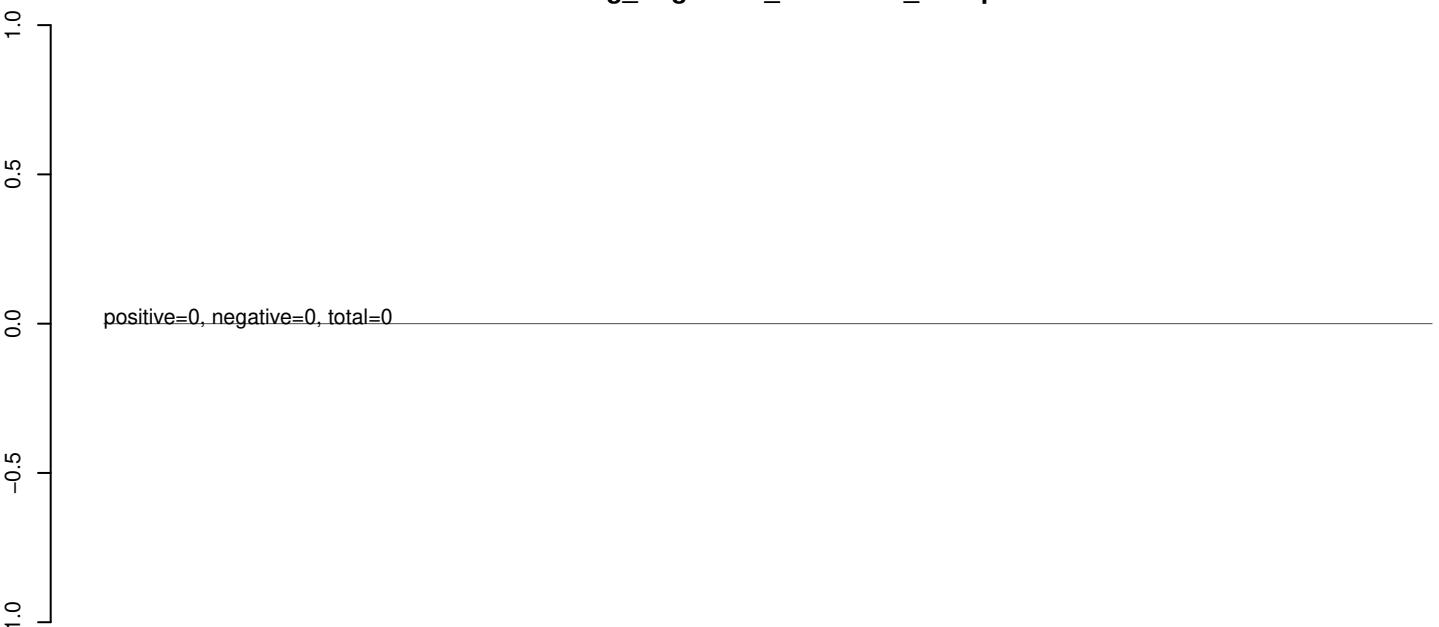
Window size=25, length=9264, virus@NC_001834.1-Drosophila_C_virus_strain_EB:1-9264

0 2000 4000 6000 8000

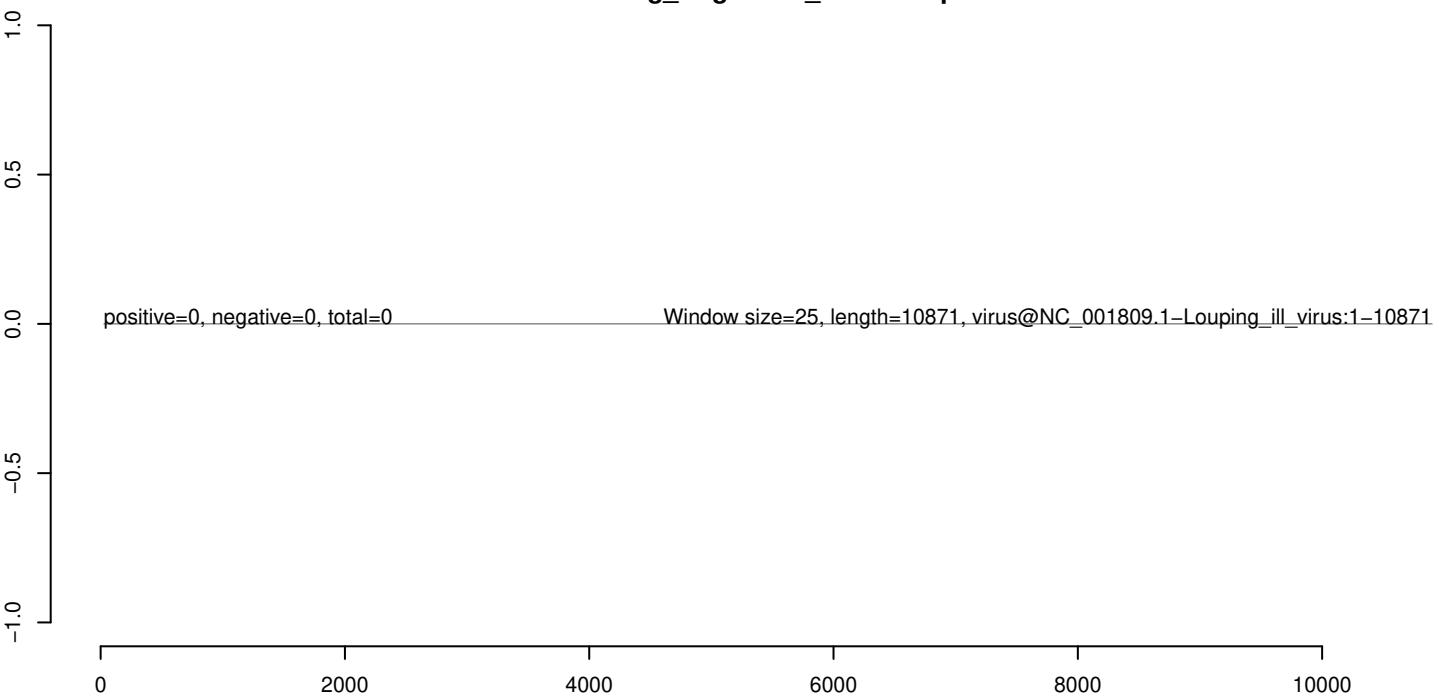
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

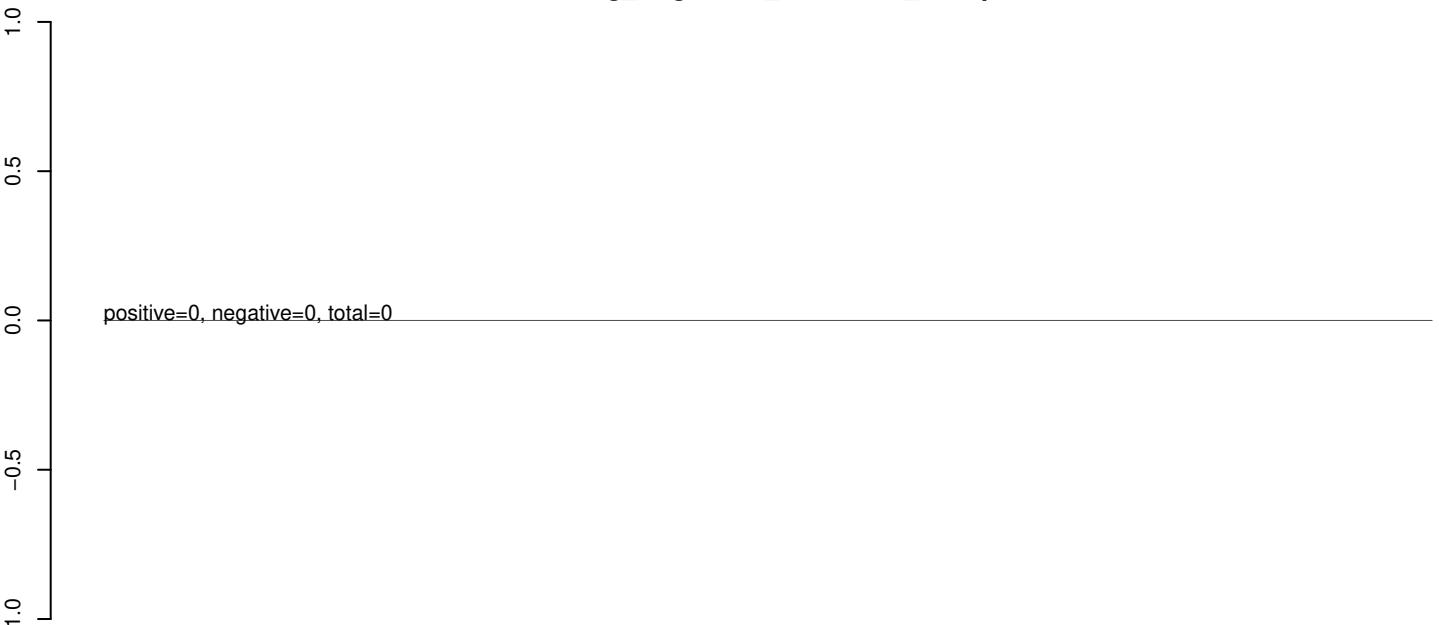
AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

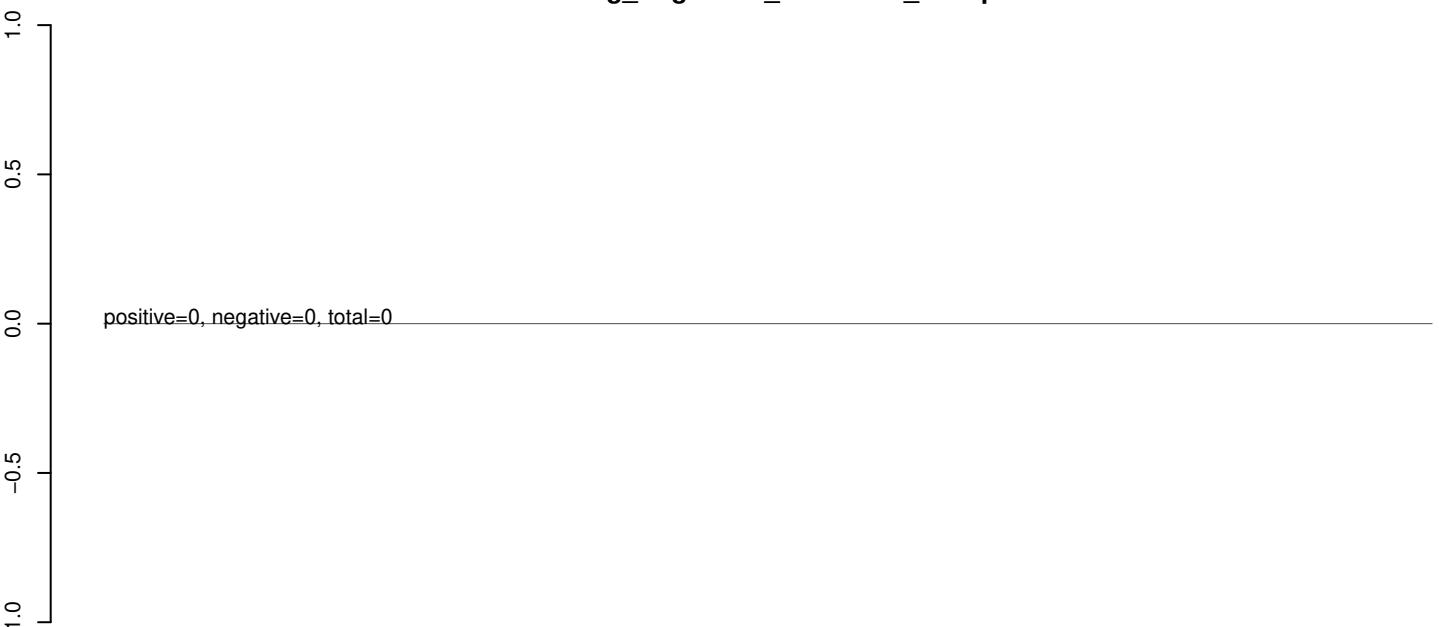
Window size=25, length=11141, virus@NC_001672.1-Tick-borne_encephalitis_virus:1-11141

0 2000 4000 6000 8000 10000

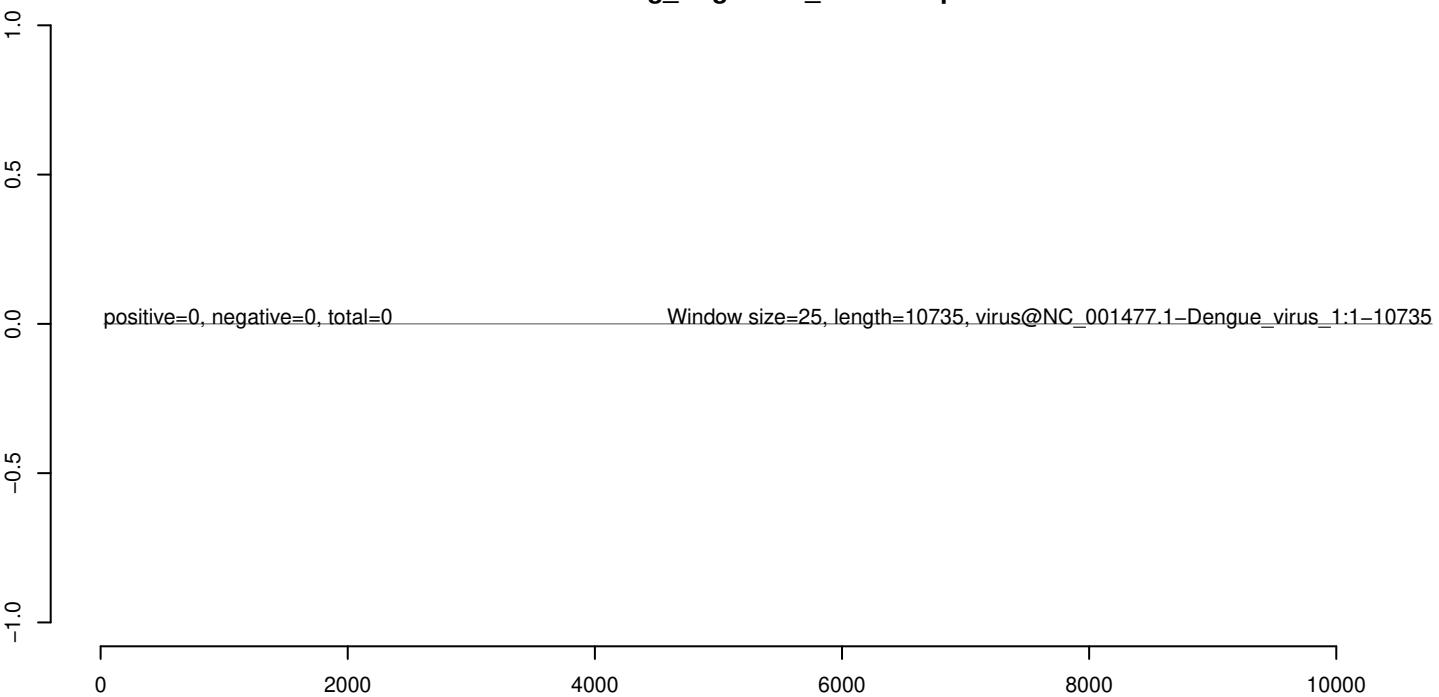
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window.size=25, length=10707, virus@NC_001475.2-Dengue_virus_3:1-10707

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window.size=25, length=10723, virus@NC_001474.2-Dengue_virus_2:1-10723

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window.size=25, length=10976, virus@NC_001437.1-Japanese_encephalitis_virus_genome:1-10976

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=4505, virus@NC_001411.2-Black_beetle_virus_RNA1_and_RNA2:1-4505

0 1000 2000 3000 4000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=11014, virus@NC_000943.1–Murray_Valley_encephalitis_virus:1–11014

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

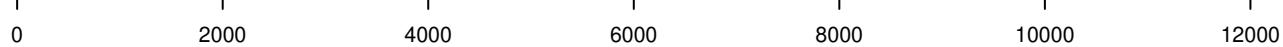
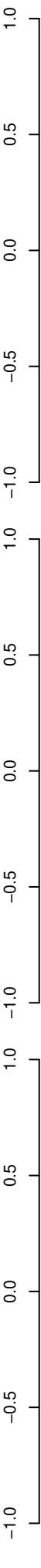
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=12916, virus@MH237595.1-Aedes_aegypti_anphevirus_strain_1a:1-12916



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0 Window size=25, length=3157, virus@MH188043.1-Culex_densovirus_strain_CDenv_Madera-2:1-3157

AeAeg_Aag2SINV_GFP.IP.18_23.rep

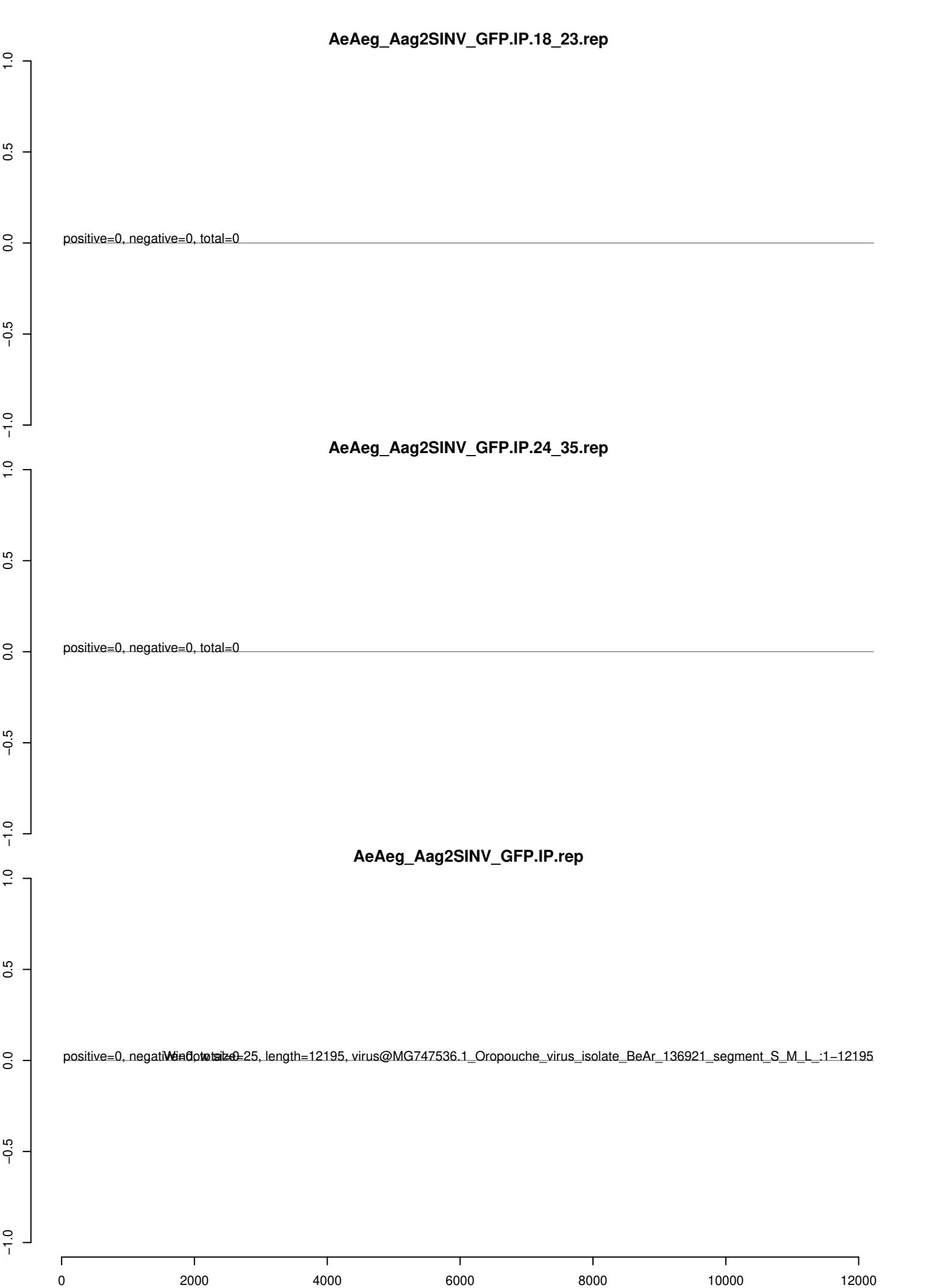
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, Window size=25, length=12195, virus@MG747536.1_Oropouche_virus_isolate_BeAr_136921_segment_S_M_L_1-12195



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, size=25, length=11664, virus@MG731558.1_Akabane_virus_isolate_HN10174_nucleoprotein_mRNA:1-11664

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

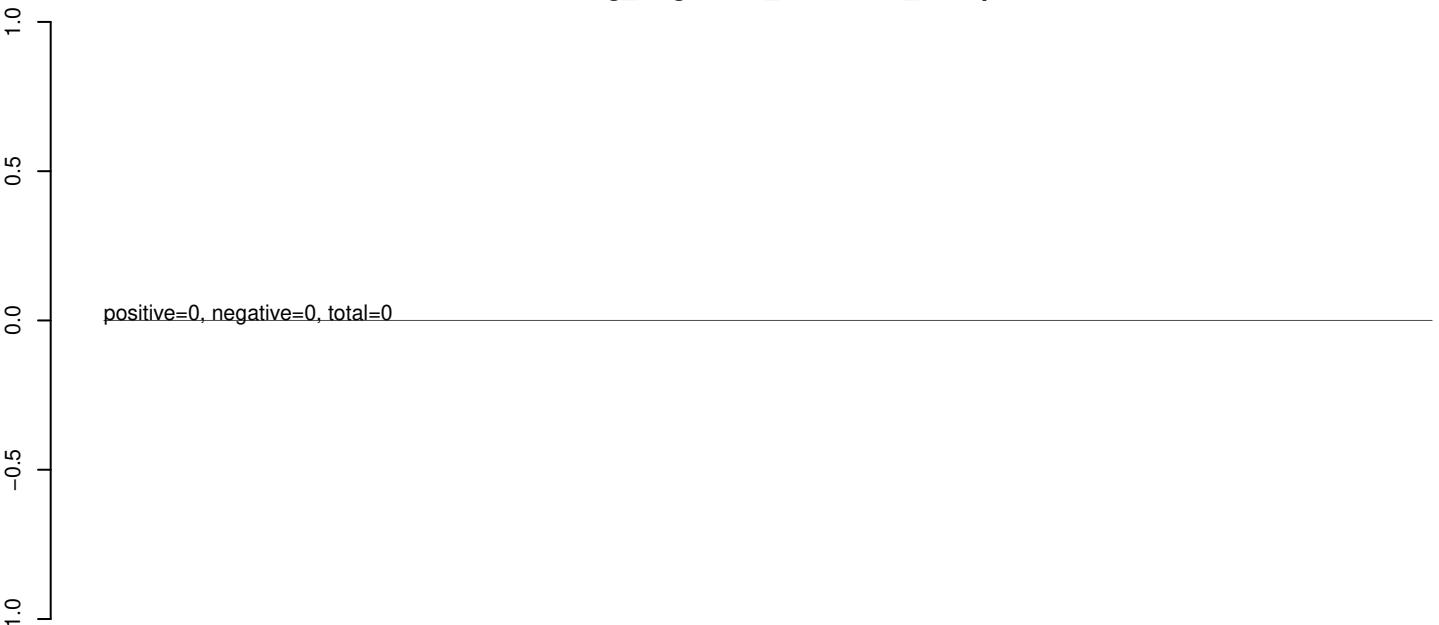
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

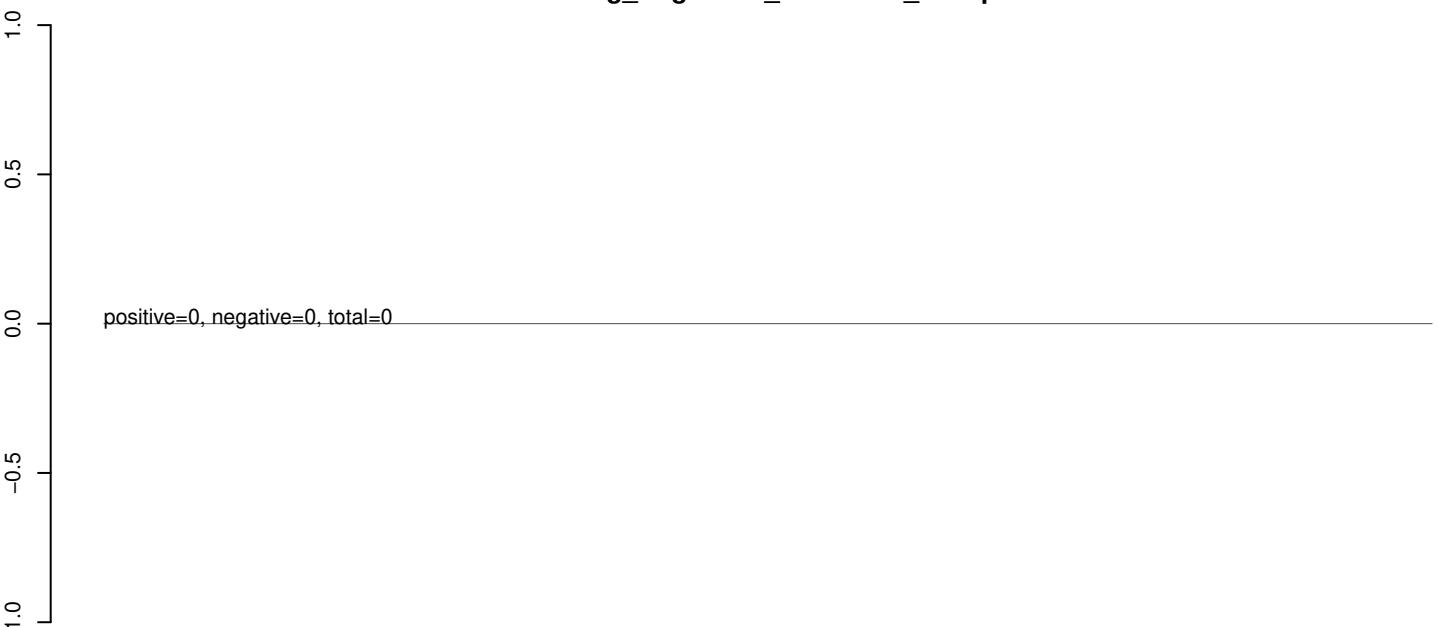
positive=0, negative=0, total=0, length=4245, virus@MF176386.1_Culex_luteo-like_virus_strain_mosWSX49277_segment_1_2:1-4245

0 1000 2000 3000 4000

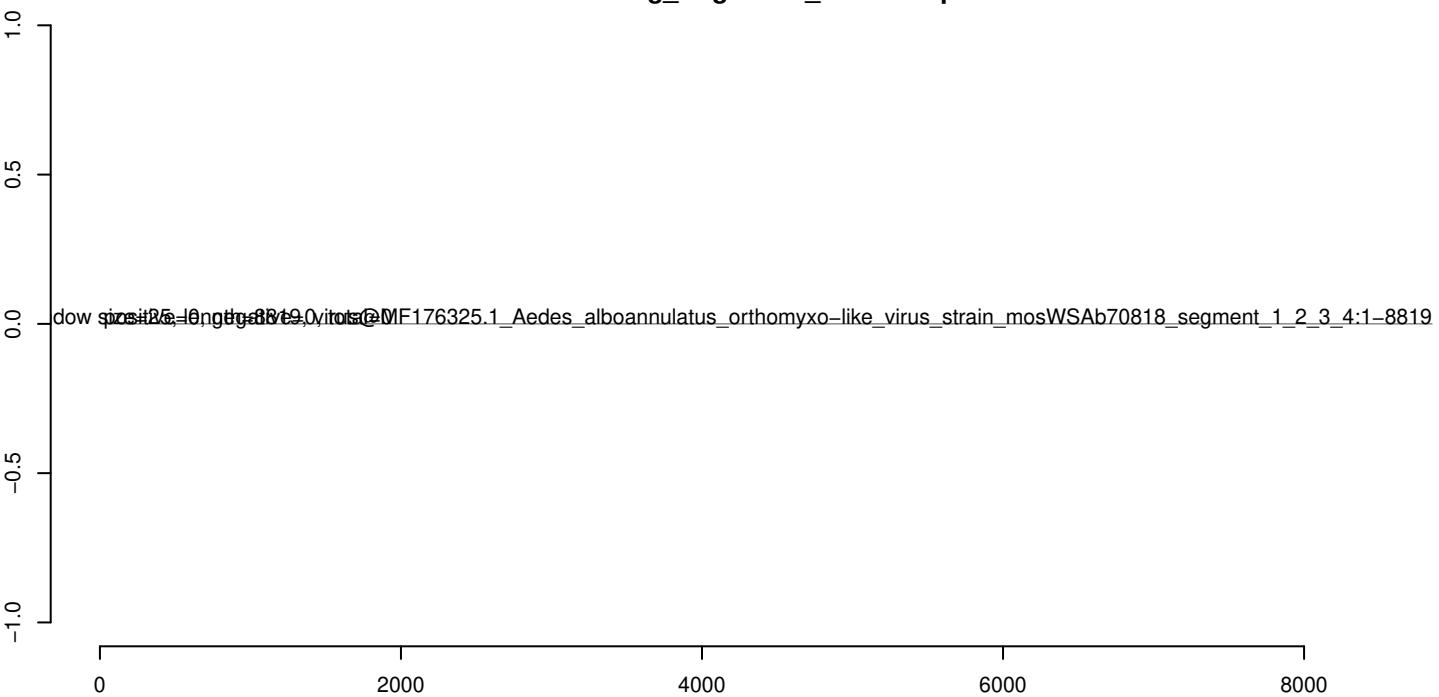
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=6, total=15, length=5982, virus@MF176317.1-Culex_mononega-like_virus_1_strain_mos191X45127_segment_2:1-5982

AeAeg_Aag2SINV_GFP.IP.18_23.rep

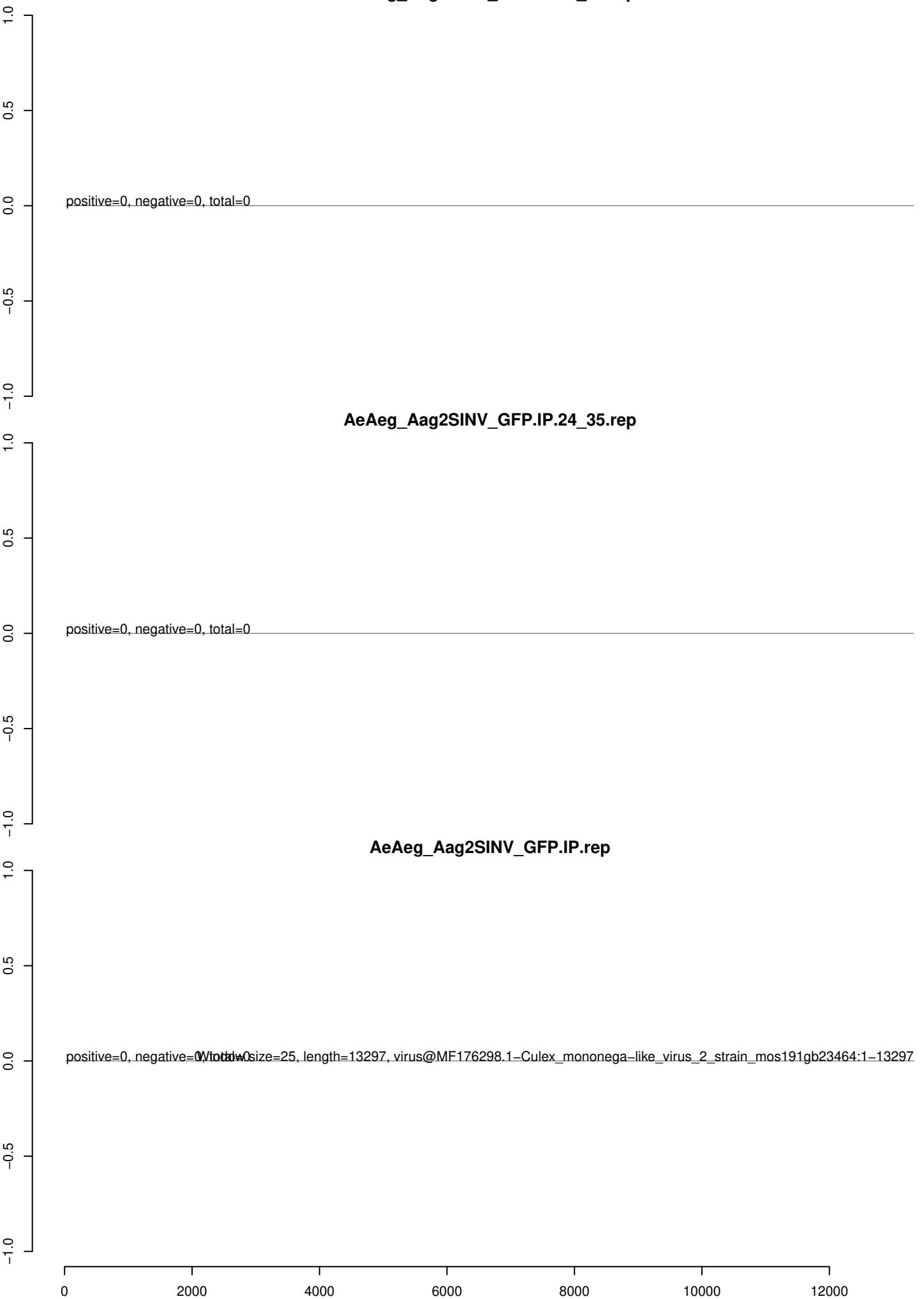
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, size=25, length=13297, virus@MF176298.1-Culex_mononega-like_virus_2_strain_mos191gb23464:1-13297



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, window size=25, length=7347, virus@MF176289.1_Wilkie_qin-like_virus_strain_mos182CP103635_segment_1_2:1-7347

0 2000 4000 6000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

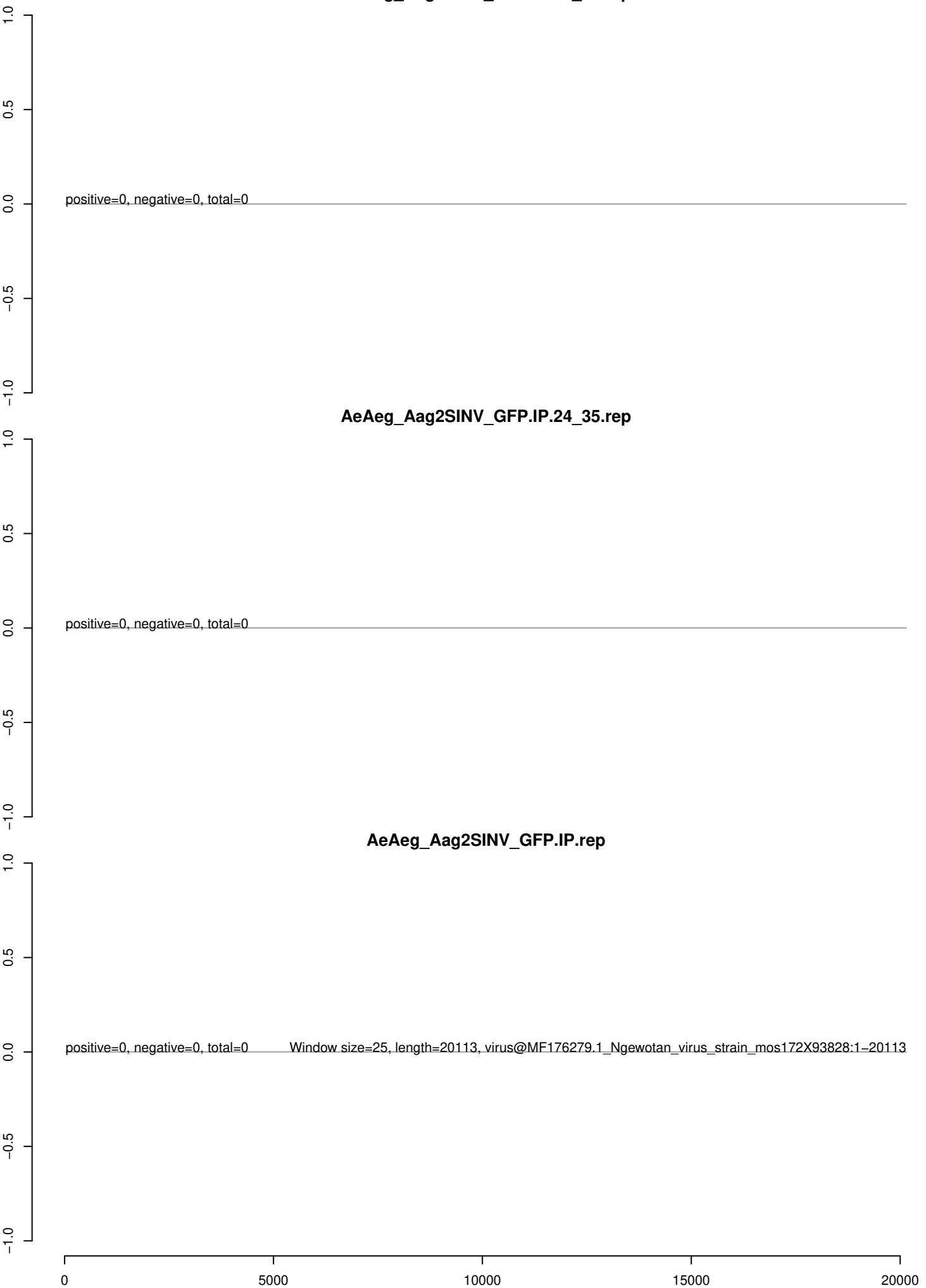
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=20113, virus@MF176279.1_Ngewotan_virus_strain_mos172X93828:1-20113



AeAeg_Aag2SINV_GFP.IP.18_23.rep

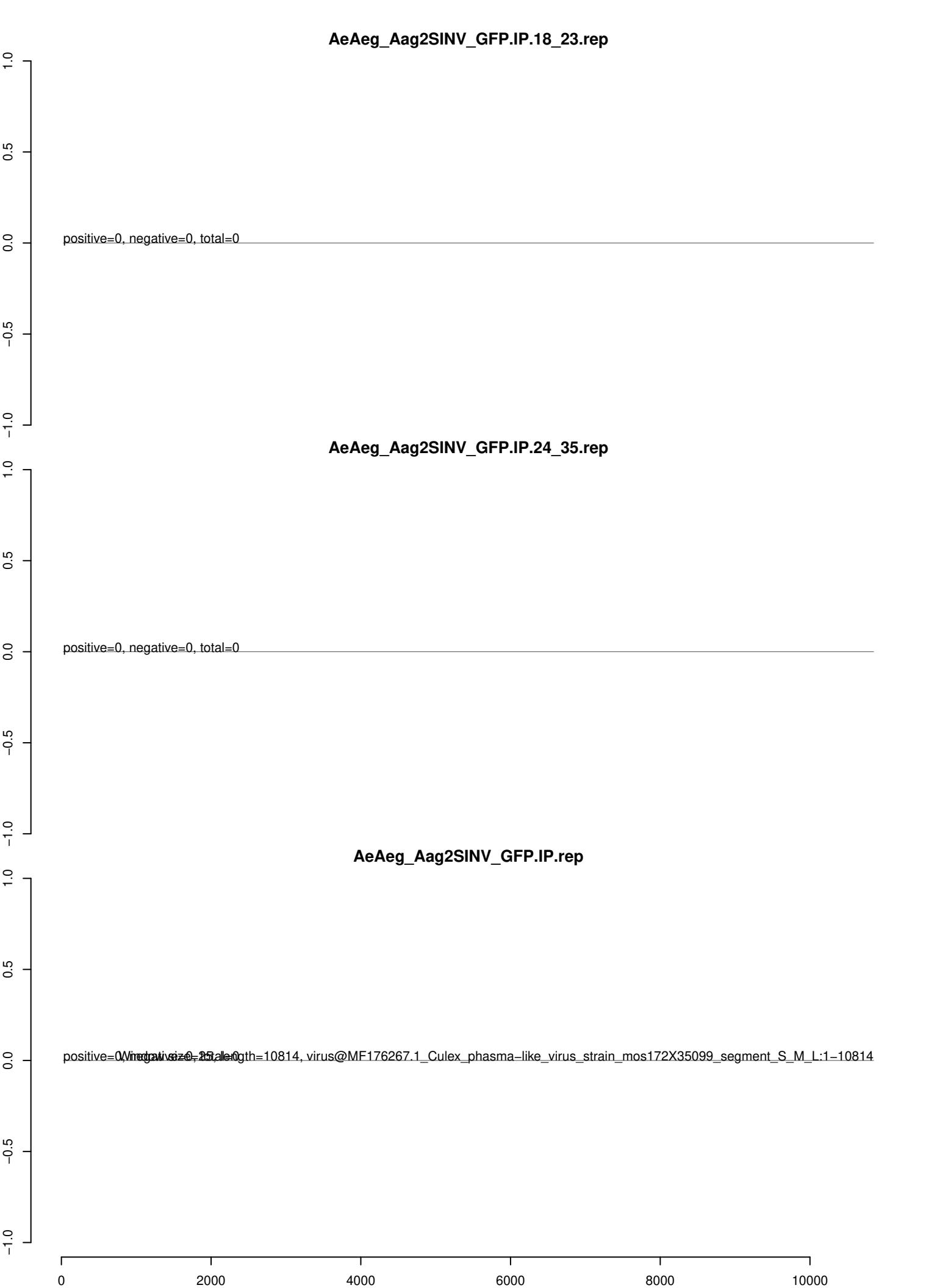
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=25, total=0, length=10814, virus@MF176267.1_Culex_phasma-like_virus_strain_mos172X35099_segment_S_M_L:1-10814





positive=0, negative=0, total=0

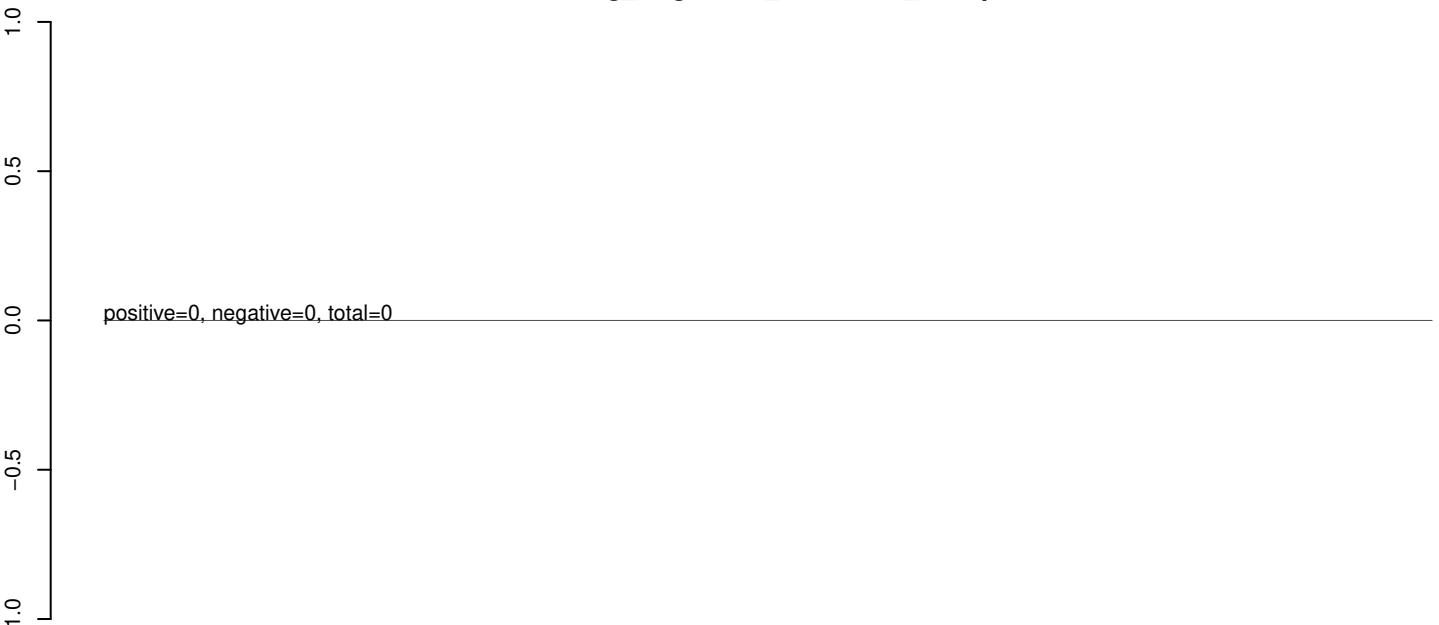


positive=0, negative=0, total=0

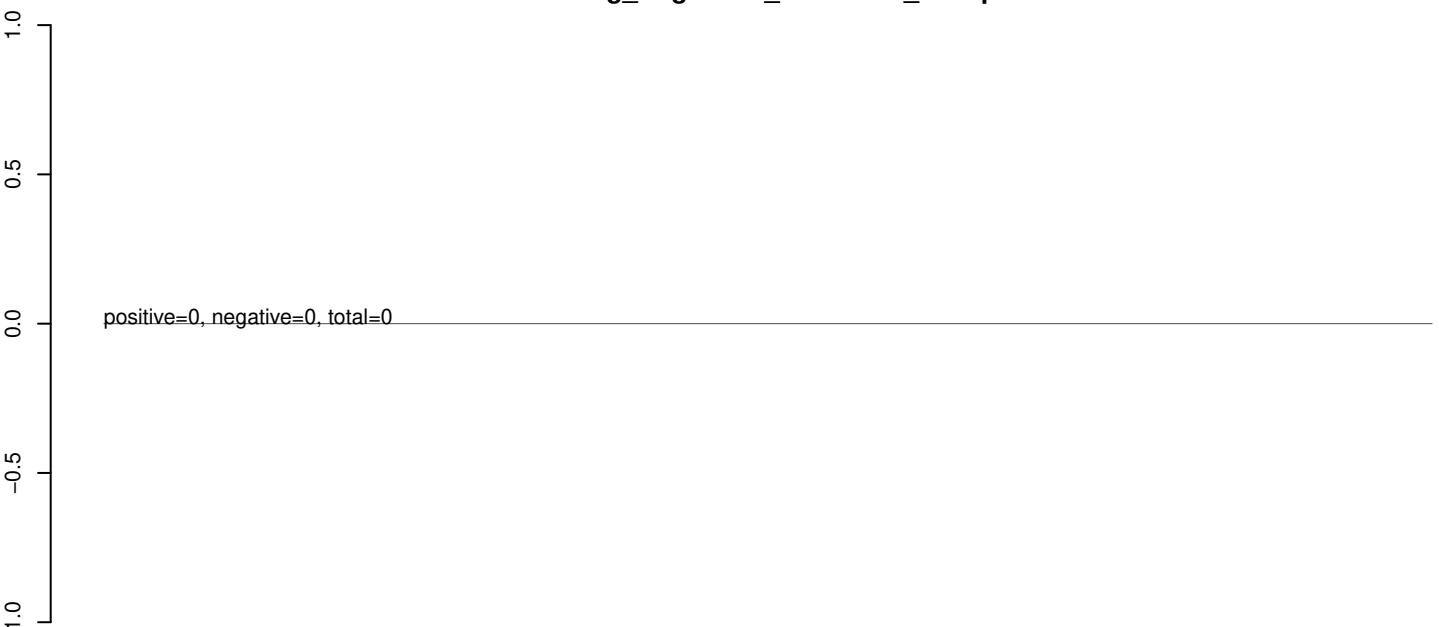


postWt65361#25-(erato)T427.virus@ME176248.1 Wuhan Mosquito Virus 6 strain mos172gb65361 segment 1 2 3 4 5:1-10427

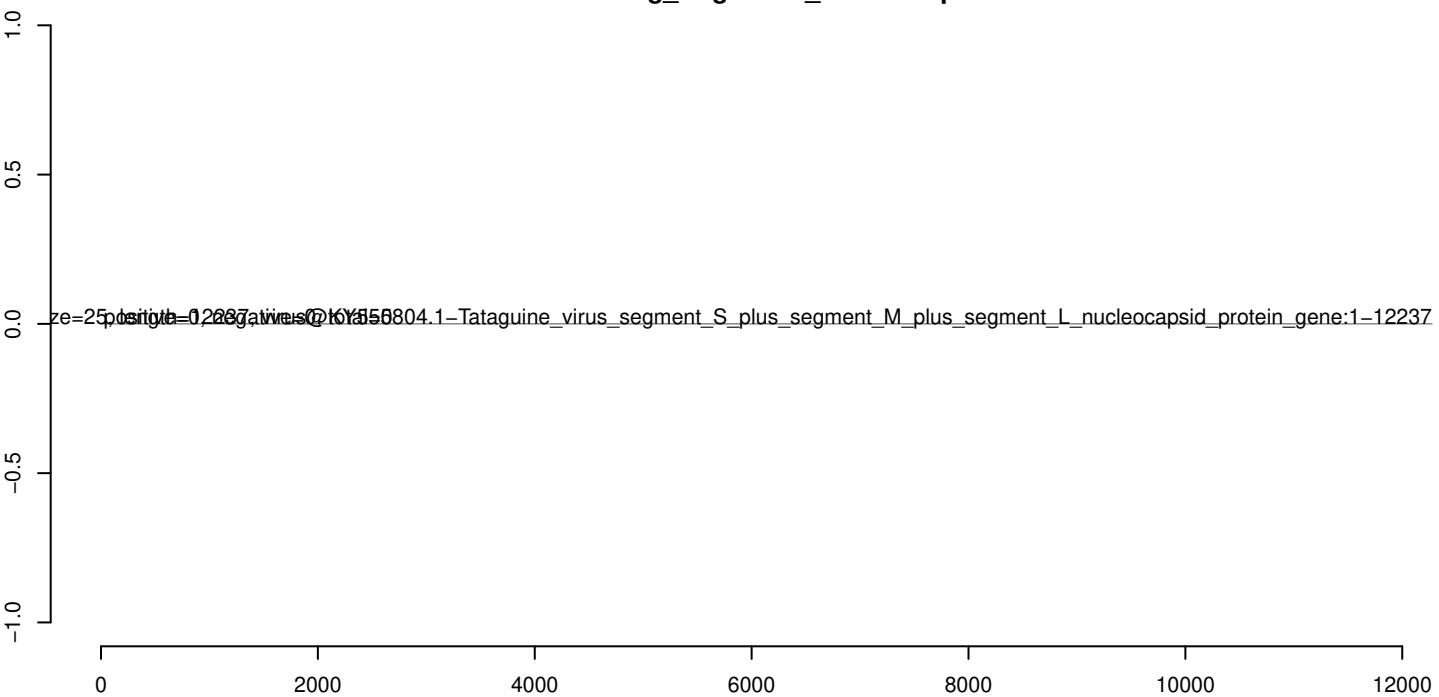
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

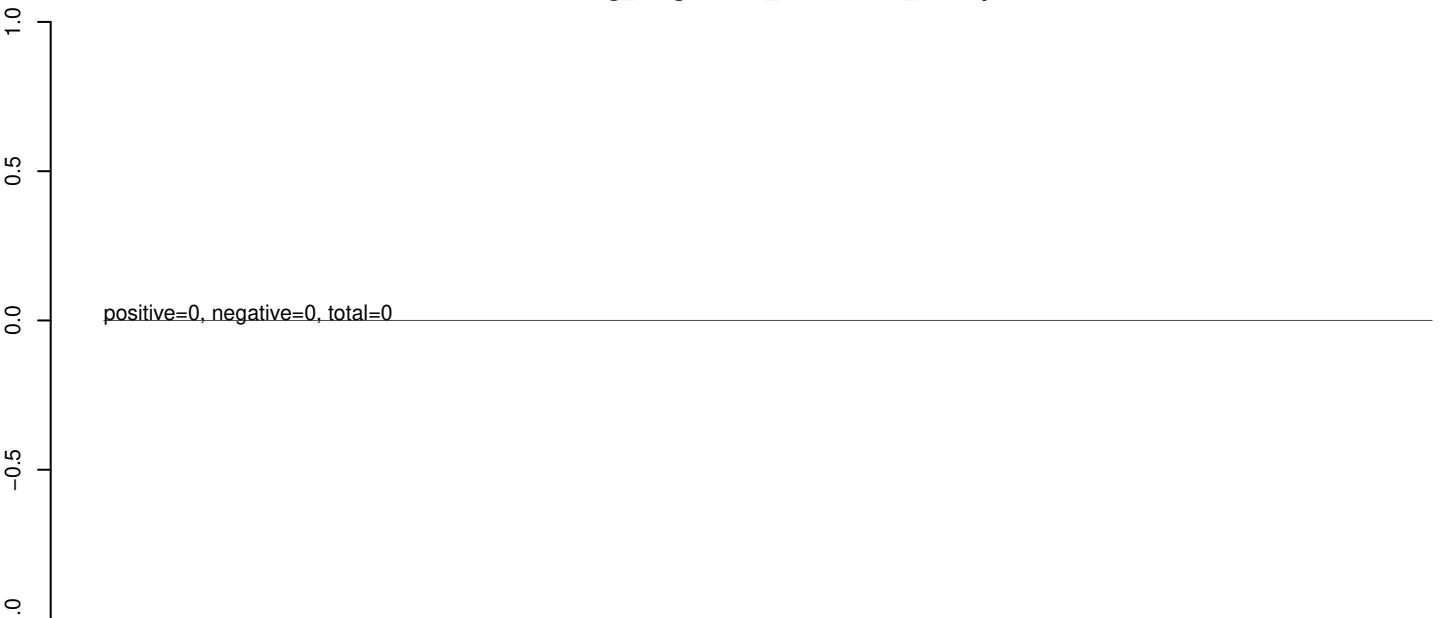
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

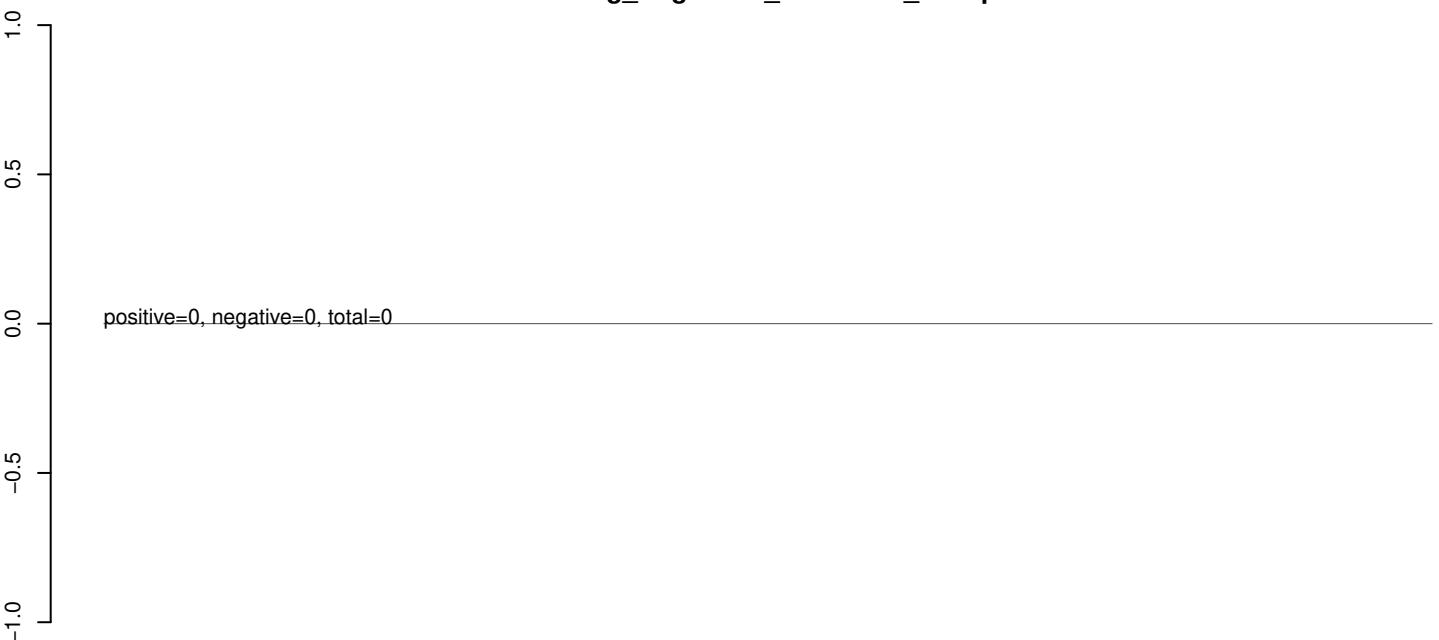
positive=0, negative=0, total=0, window_size=25, length=10749, virus@KX669682.1-Calbertado_virus_isolate_Cal_mos_2_polyprotein_gene:1-10749

0 2000 4000 6000 8000 10000

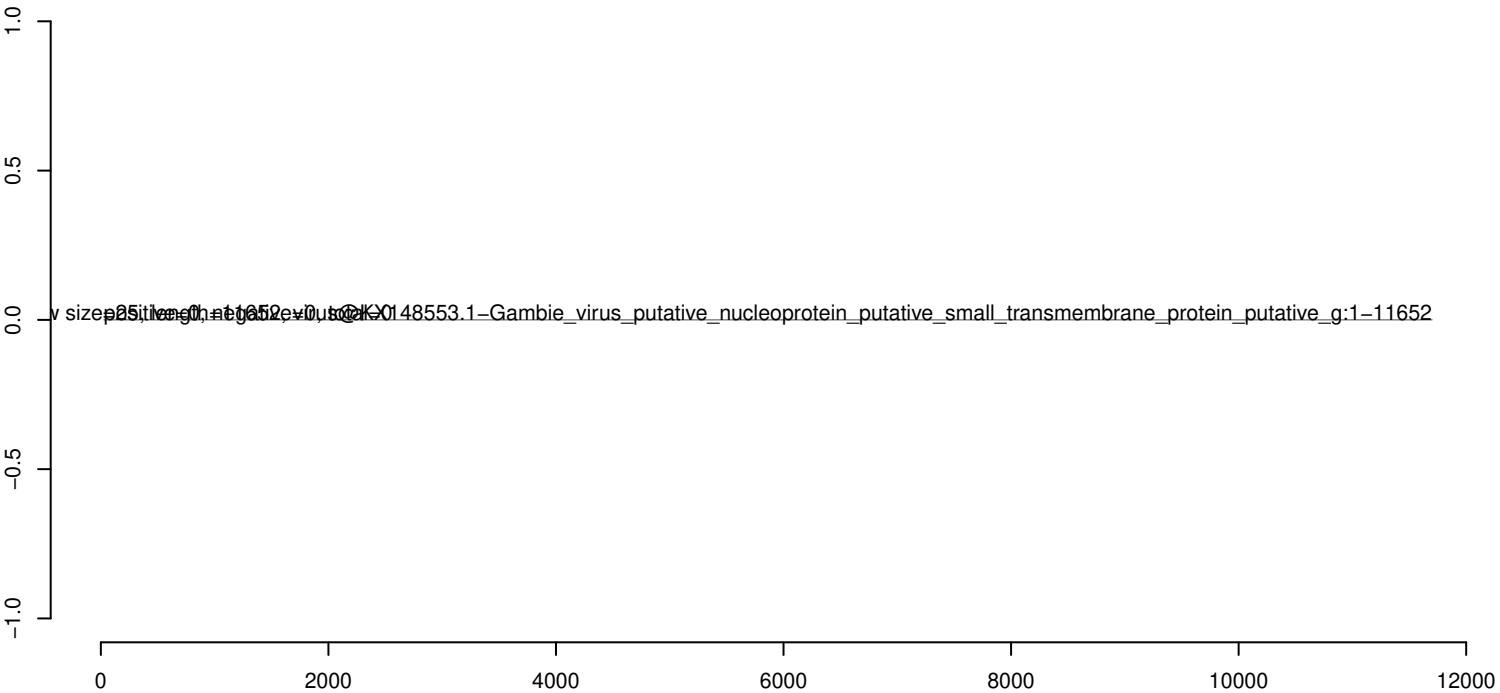
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=11798, virus@KU194360.1_Merida_virus_isolate_MERD-Mex07:1-11798

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=25, length=4780, virus@KU169879.1-Anopheles_cytopivirus_RNA-dependent_RNA_polymerase_gene:1-4780

0 1000 2000 3000 4000 5000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

e=25, positive=4102, negative=0, total=4402
1-Humaita-Tubiacanga_virus_isolate_Rio_segment_RNA-dependent_RNA_polymerase:1-4402

0 1000 2000 3000 4000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

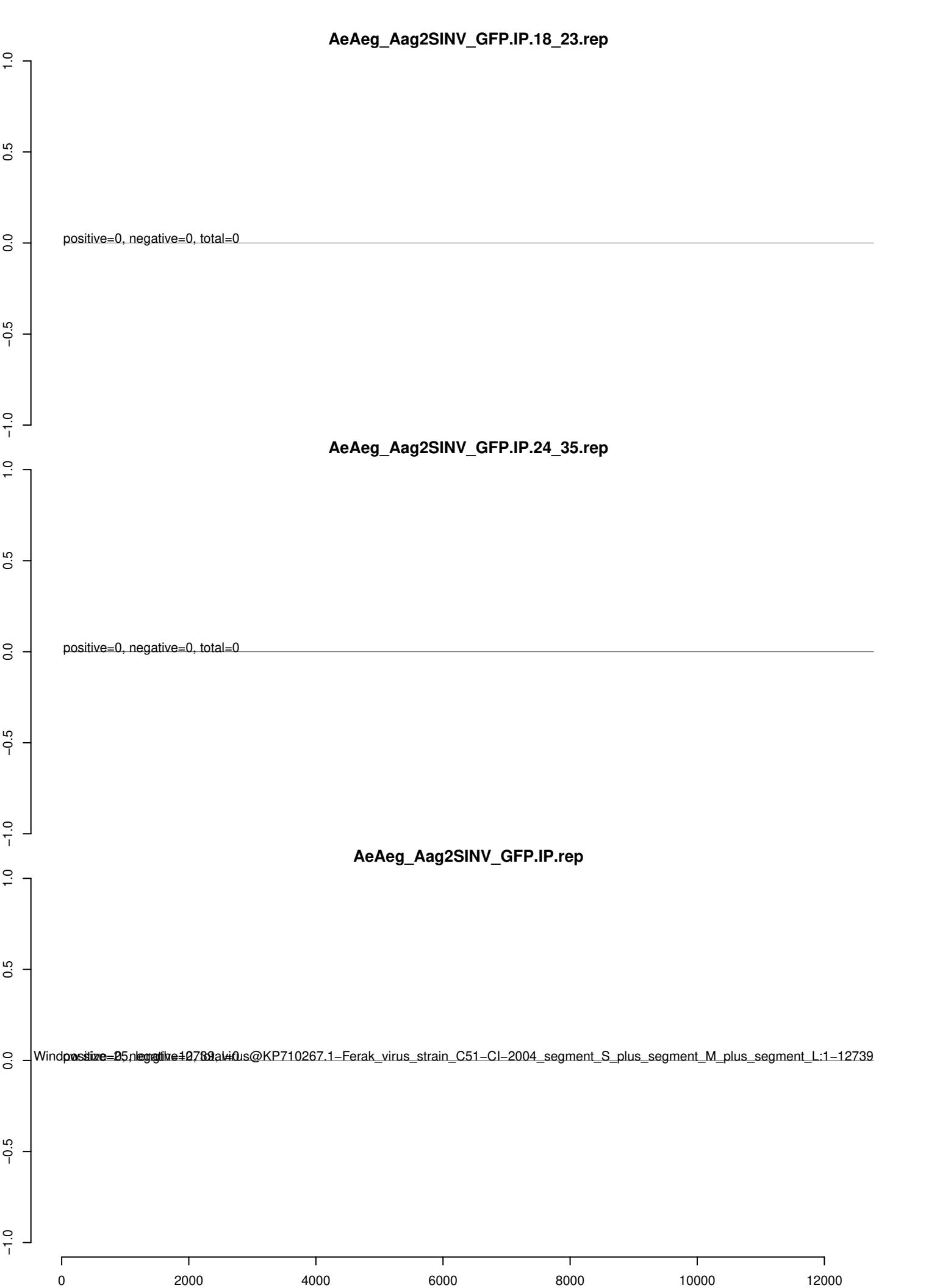
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

Wind positive=25 negative=10789 virus@KP710267.1-Ferak_virus_strain_C51-CI-2004_segment_S_plus_segment_M_plus_segment_L:1-12739



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10735, virus@KM225265.1-Torres_virus_strain_TS5273:1-10735

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10648, virus@KM225264.1-Bainyik_virus_strain_MK7979:1-10648

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window.size=25, length=10560, virus@KM225263.1–Stratford_virus_strain_C338:1–10560

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

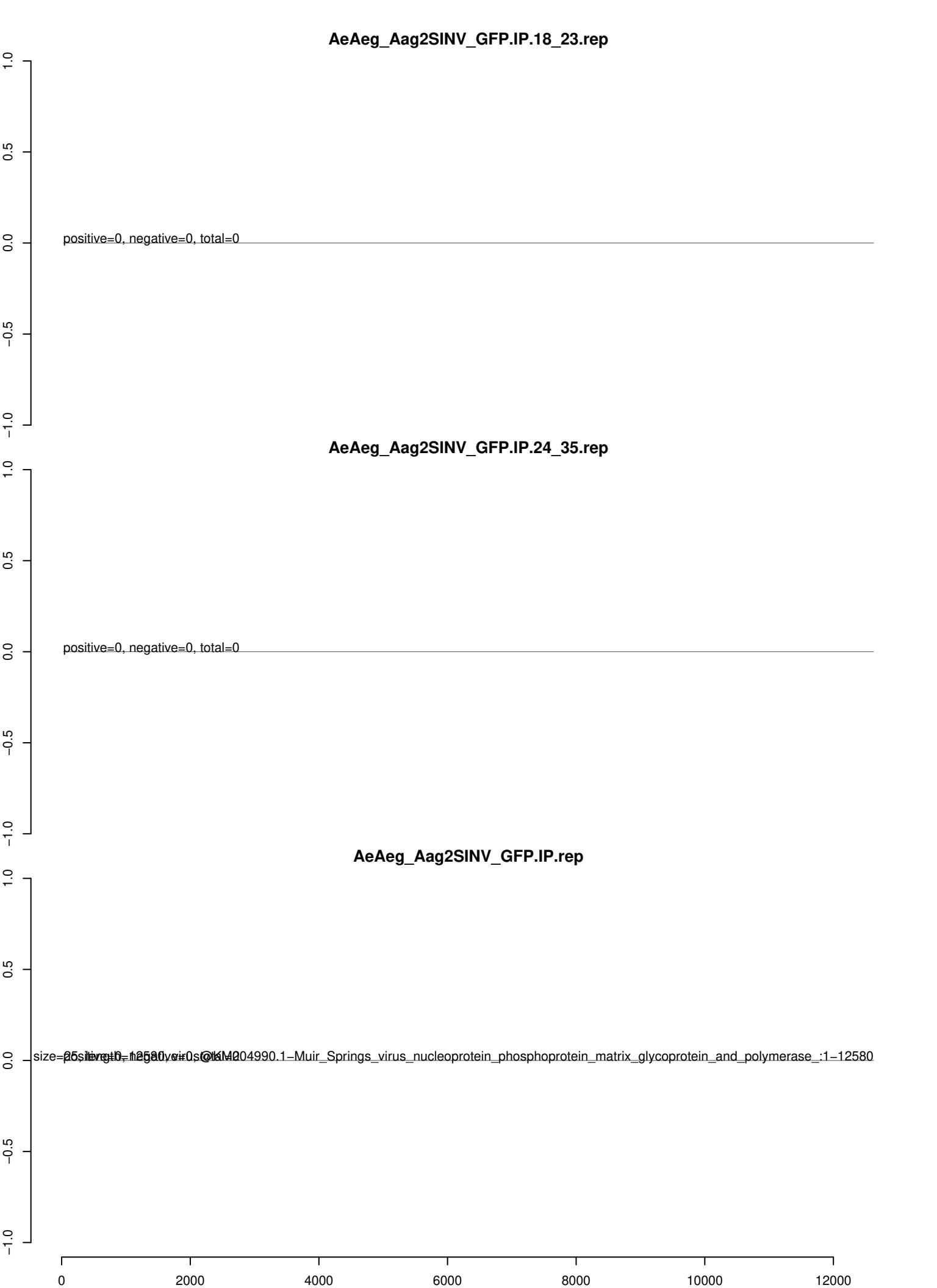
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

size=2511 length=12580 virion=0 score=1.004990.1 Muir_Springs_virus_nucleoprotein_phosphoprotein_matrix_glycoprotein_and_polymerase_1-12580



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

e=25, positive=26, negative=0, total=264.1-Anopheline-associated_C_virus_isolate_acc_1.1_segment_1_RNA-dependent_RNA_polymeras:1-3601

0 1000 2000 3000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

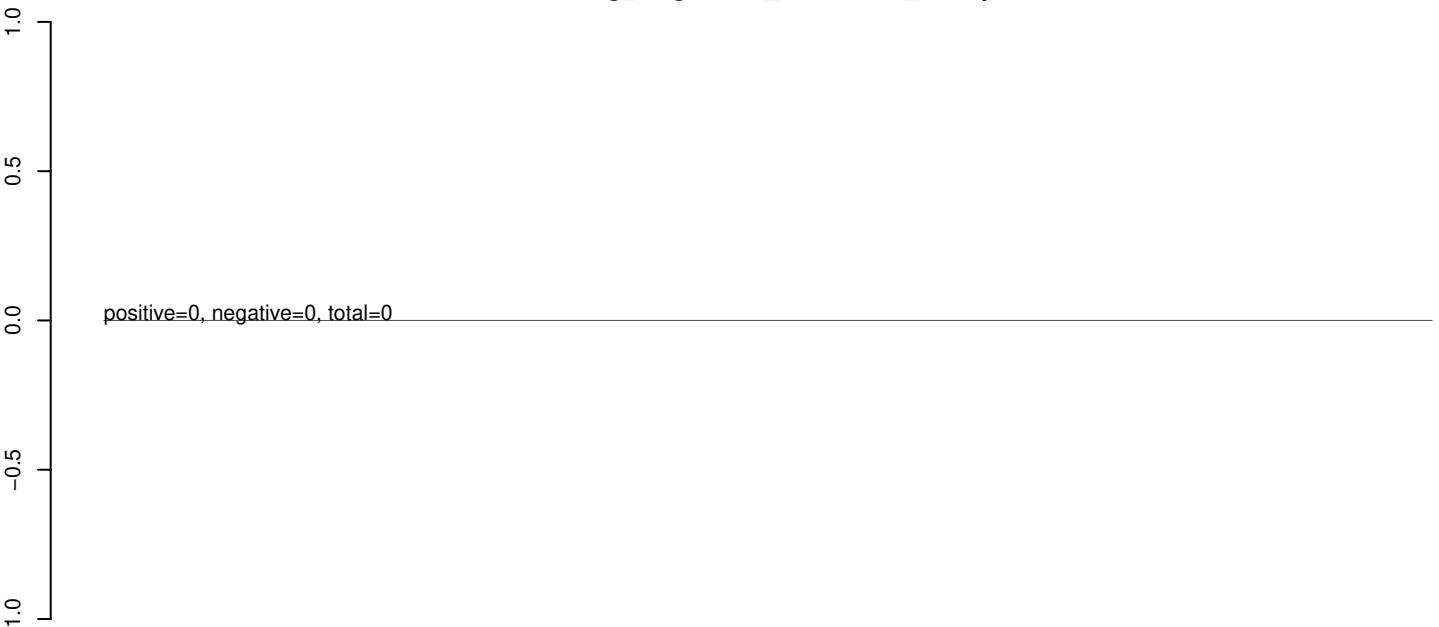
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

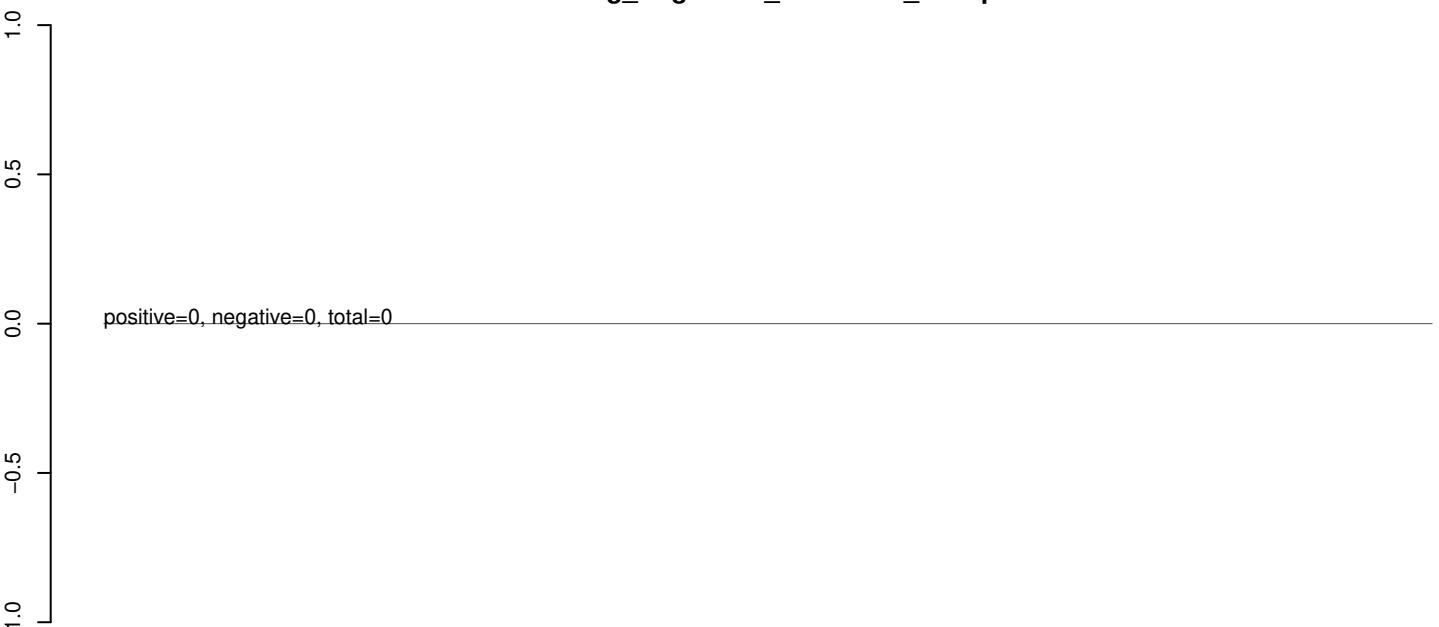
AeAeg_Aag2SINV_GFP.IP.rep

size=25 single=0,1595 unique=0,16020664.1_Flanders_virus_strain_M11750-TX-2005_U1_protein-U1_U2_protein-U2_and_U3_protein-U:1-1535

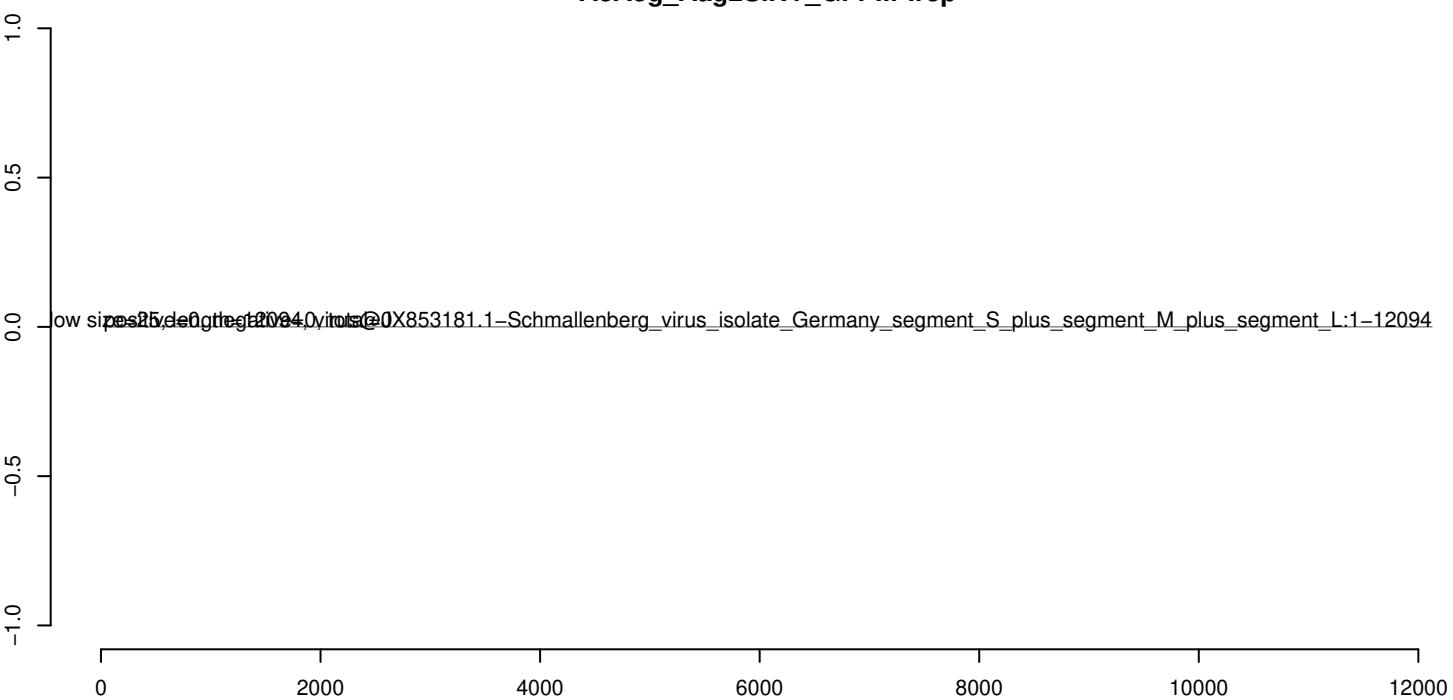
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

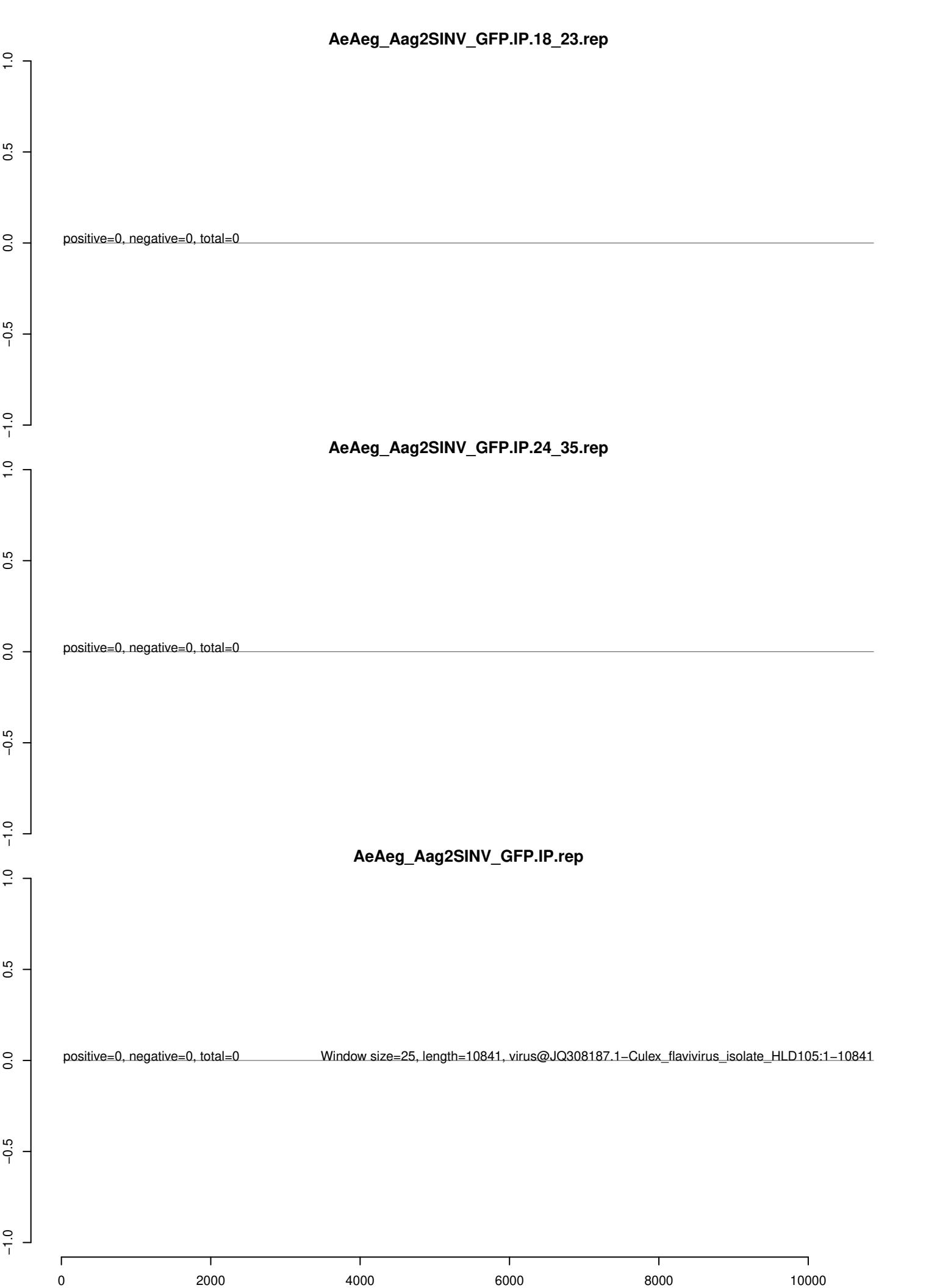
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

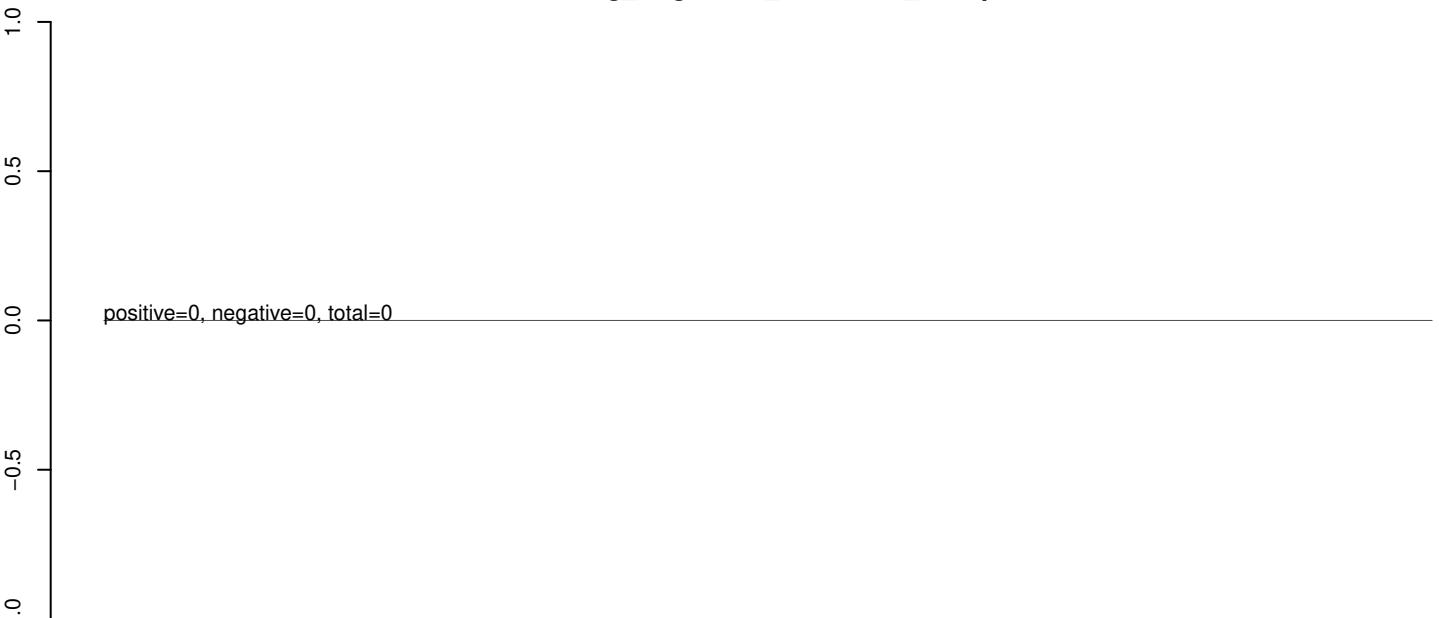
AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

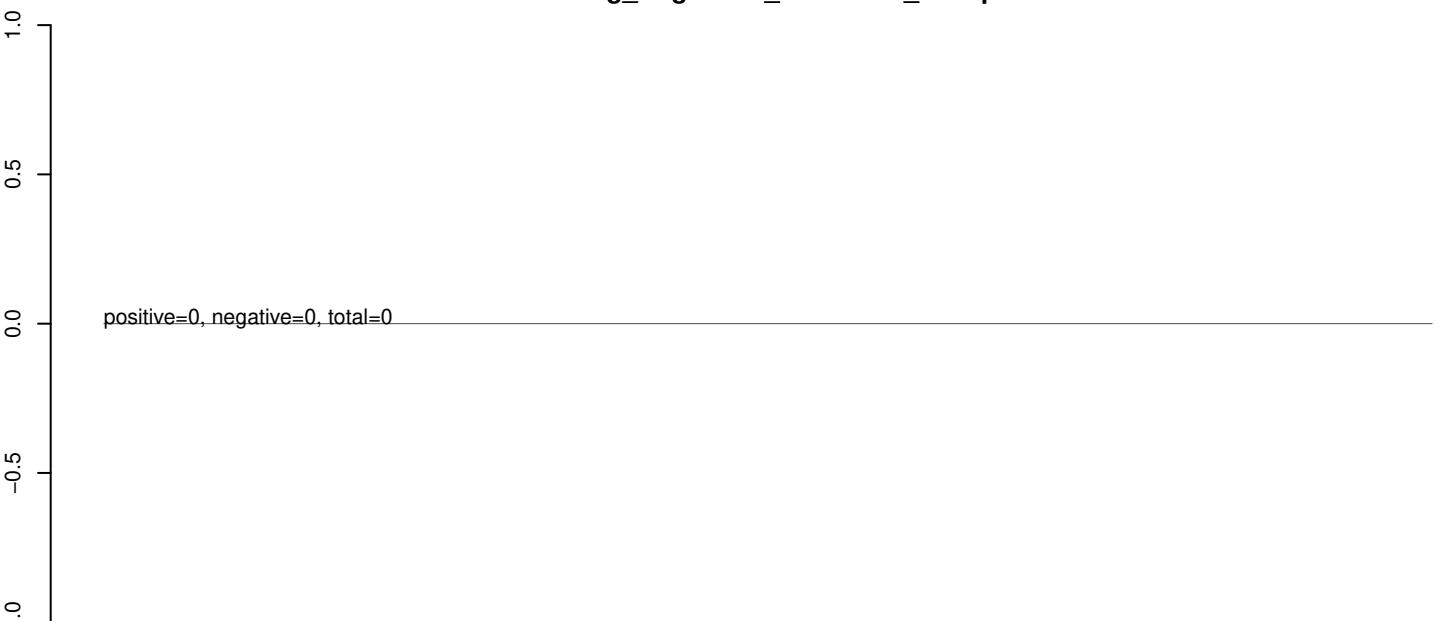
Window size=25, length=10841, virus@JQ308187.1–Culex_flavivirus_isolate_HLD105:1–10841



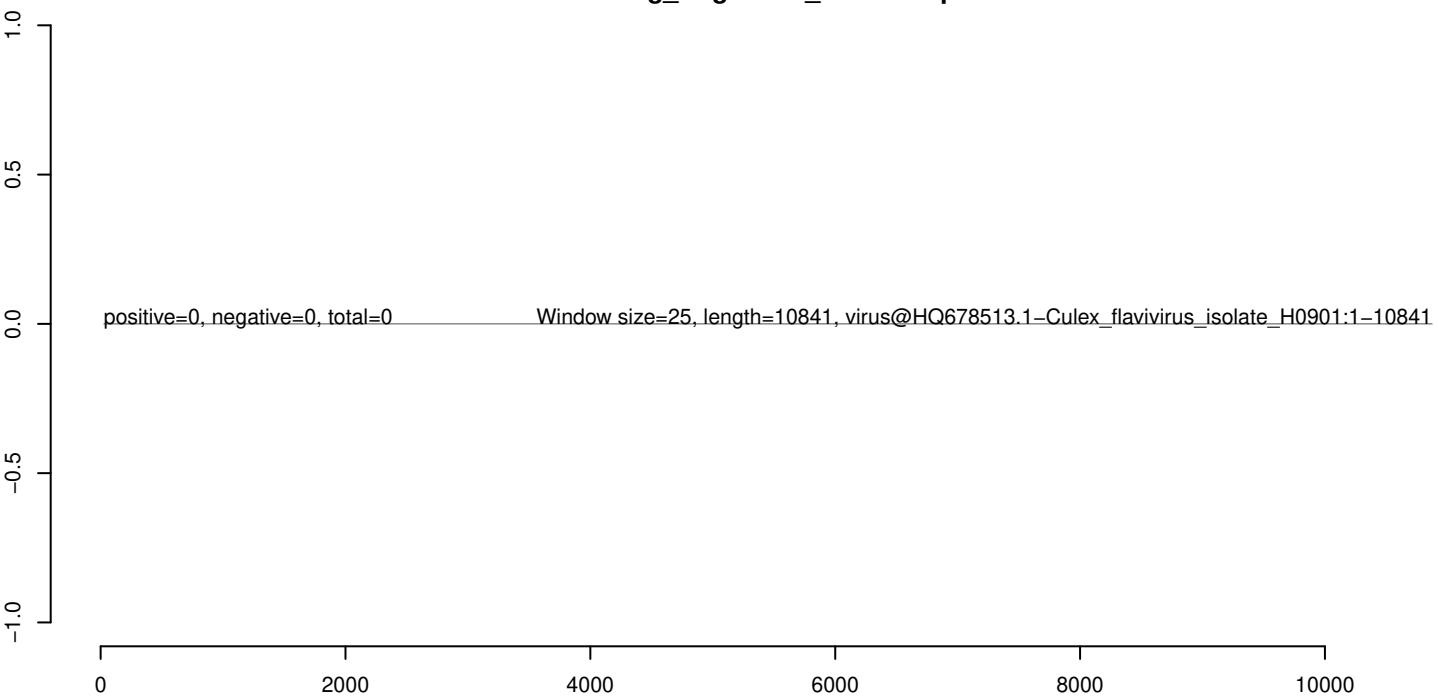
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

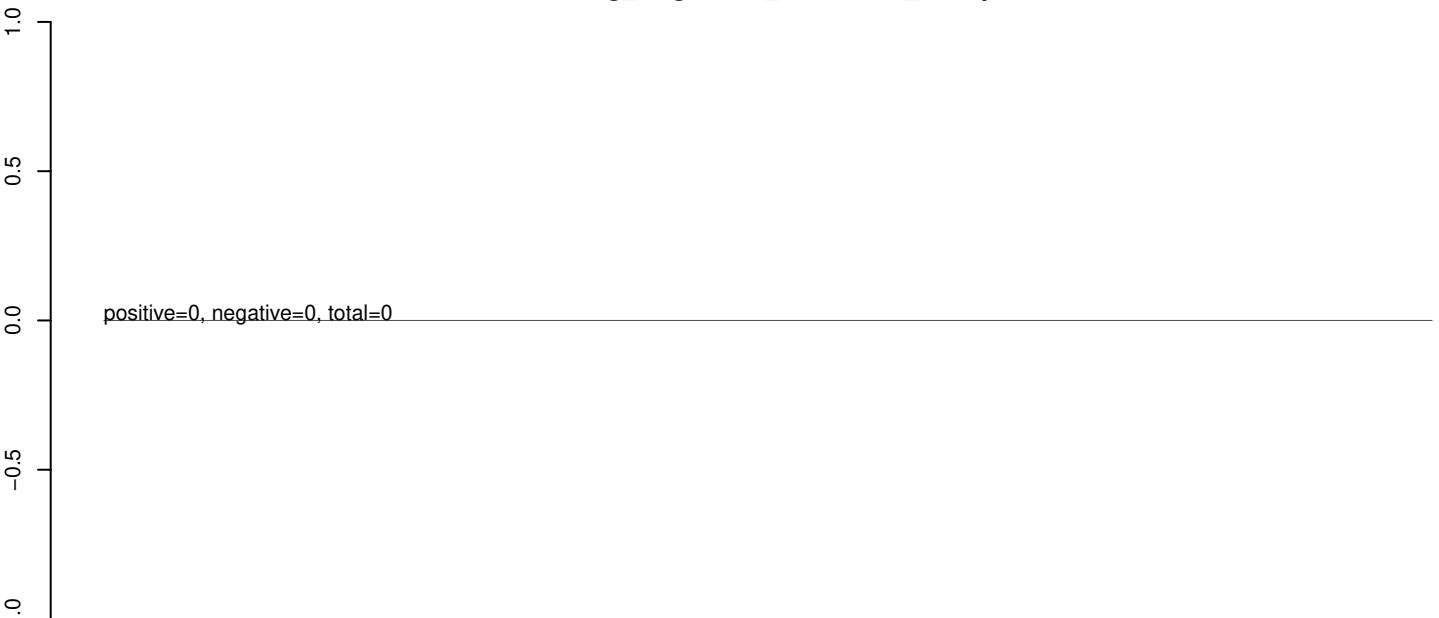
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

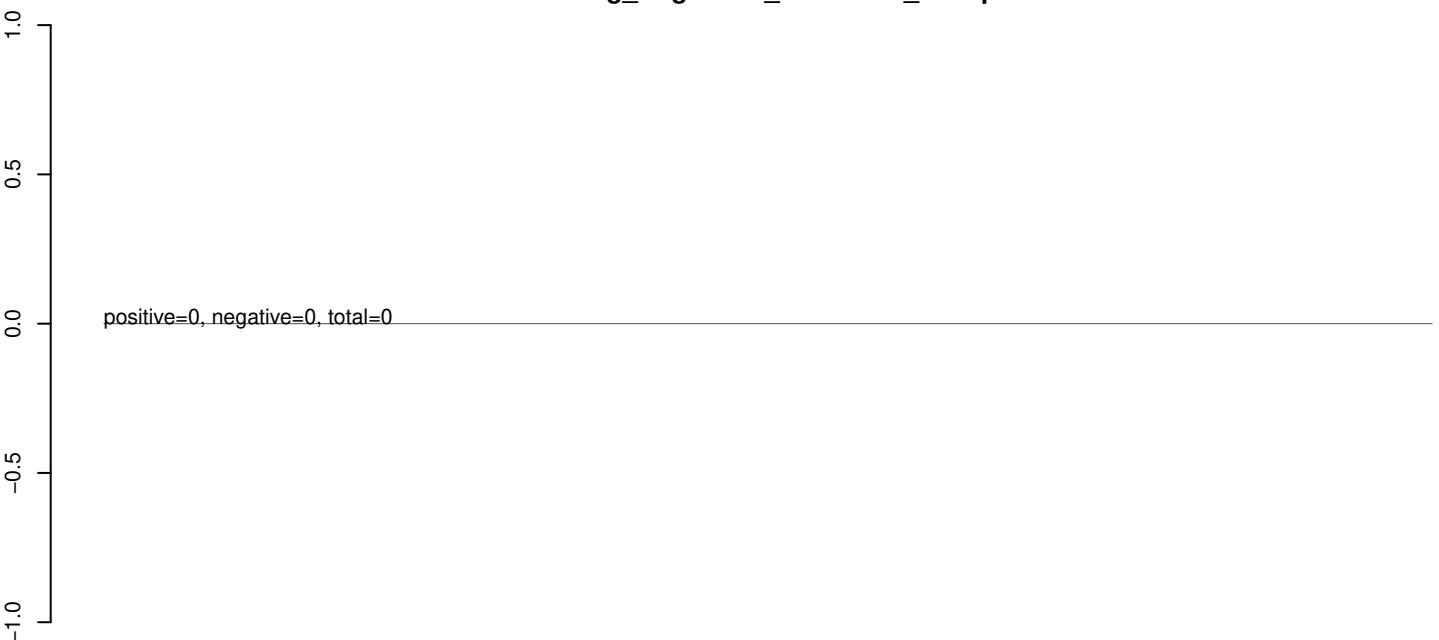
positive=0, negative=0, window size=25, length=10774, virus@HM055369.1-Kyasanur_forest_disease_virus_strain_KFD_P_9605:1-10774

0 2000 4000 6000 8000 10000

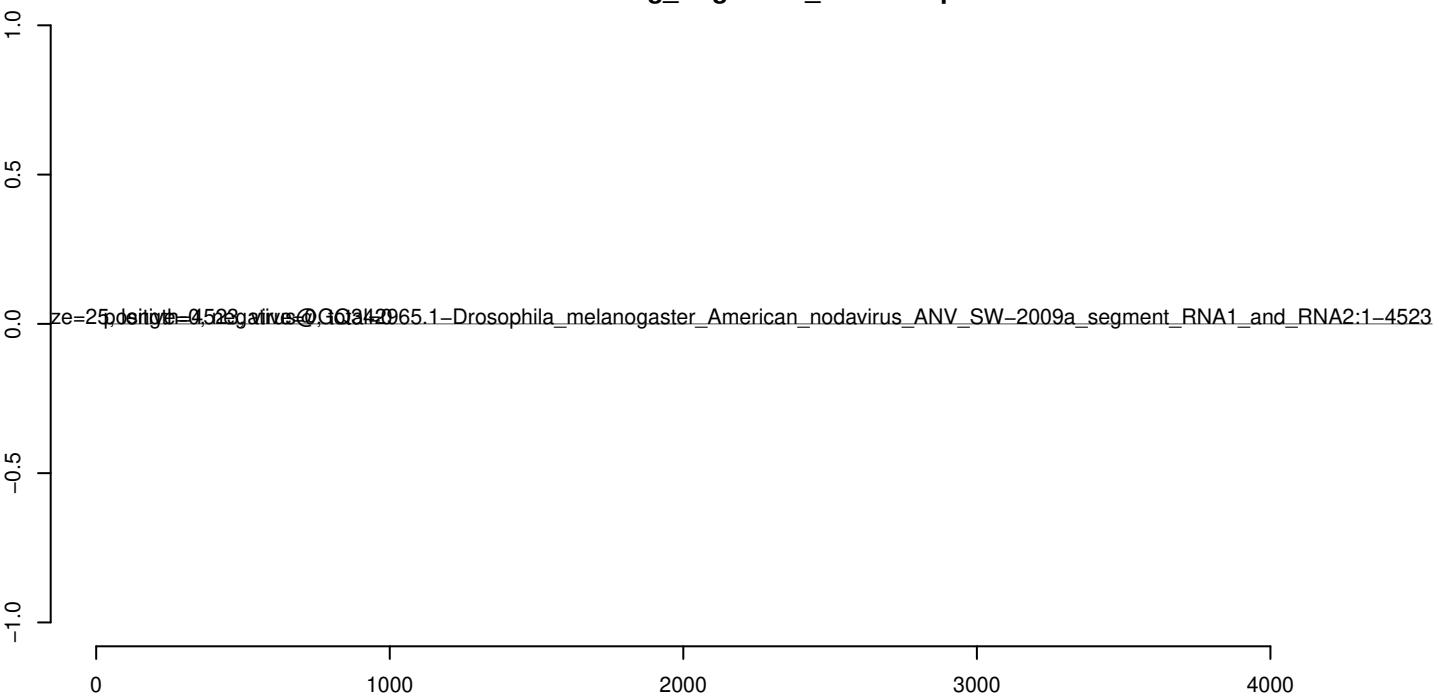
AeAeg_Aag2SINV_GFP.IP.18_23.rep



AeAeg_Aag2SINV_GFP.IP.24_35.rep



AeAeg_Aag2SINV_GFP.IP.rep



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10837, virus@FJ663034.1-Culex_flavivirus_isolate_Iowa07:1-10837

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

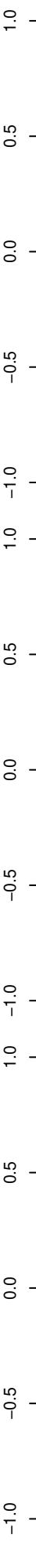
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10837, virus@FJ502995.1-Culex_flavivirus_strain_HOU24518:1-10837



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

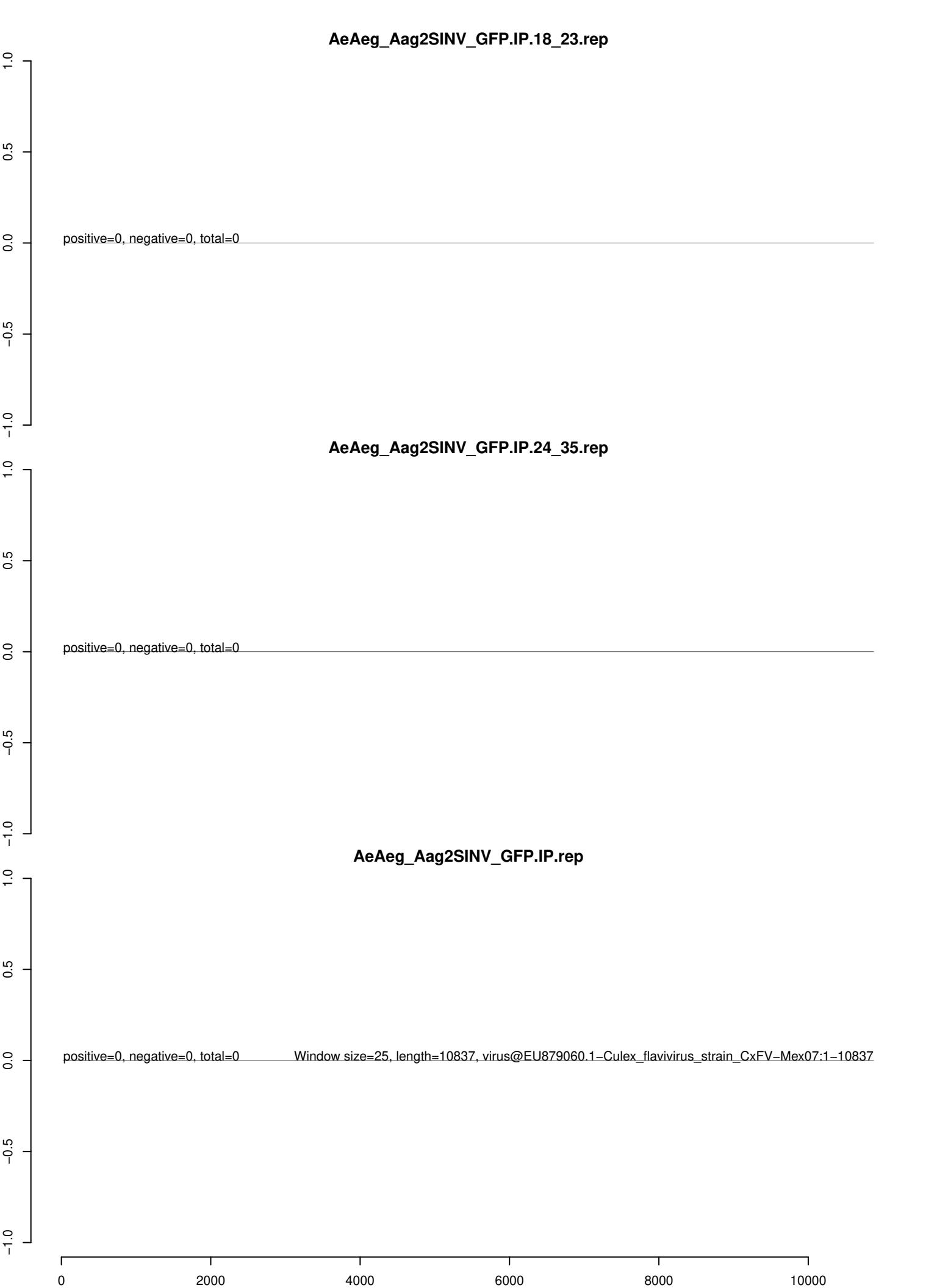
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10837, virus@EU879060.1-Culex_flavivirus_strain_CxFV-Mex07:1-10837



AeAeg_Aag2SINV_GFP.IP.18_23.rep

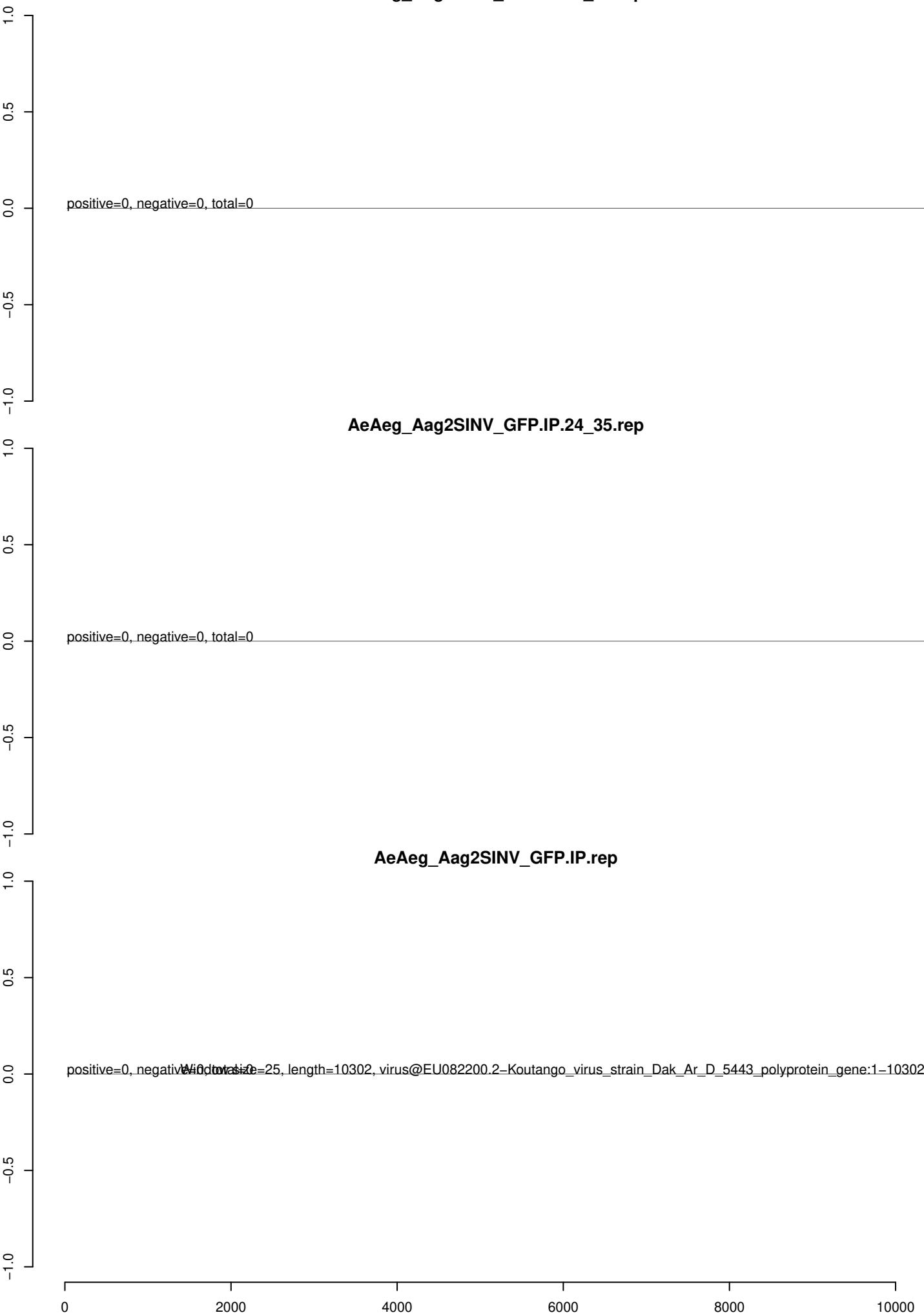
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, length=10302, virus@EU082200.2-Koutango_virus_strain_Dak_Ar_D_5443_polyprotein_gene:1-10302



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0, wsize=25, length=10254, virus@DQ235149.1-Royal_Farm_virus_from_Afghanistan_polyprotein_gene:1-10254

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10960, virus@AY898809.1-Alfuy_virus_strain_MRM3929:1-10960

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

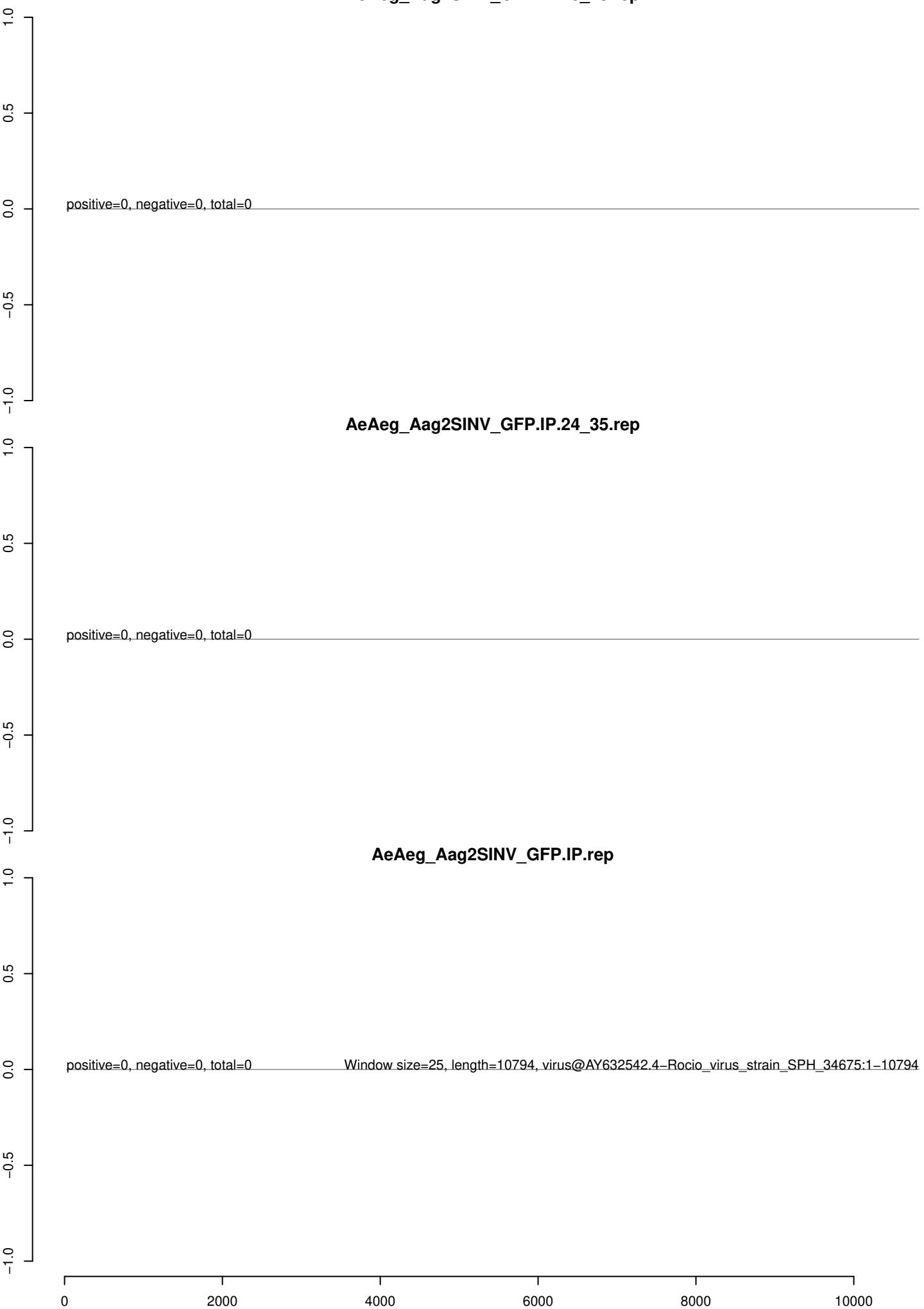
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10794, virus@AY632542.4–Rocio_virus_strain_SPH_34675:1–10794



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

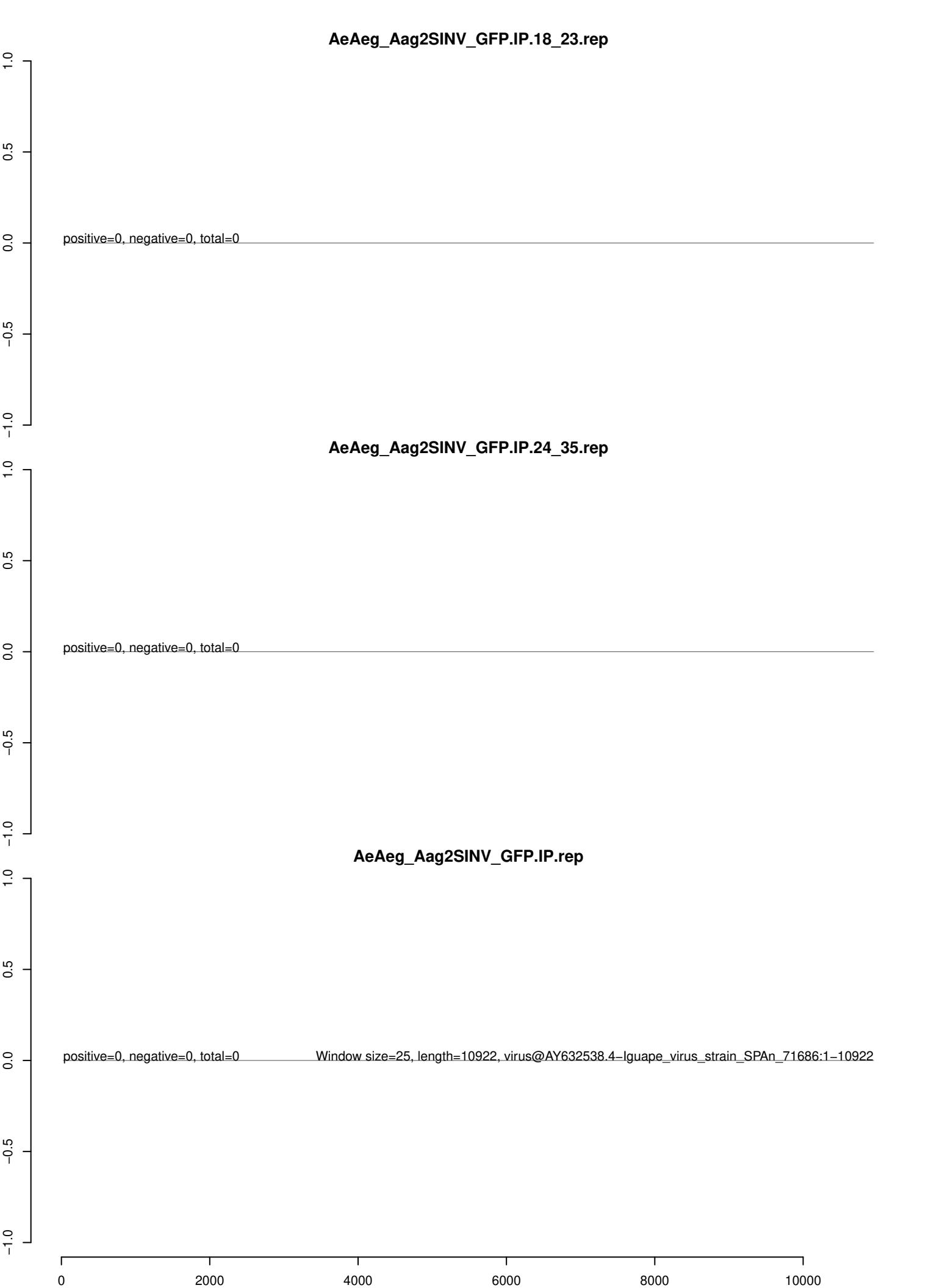
AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window size=25, length=10922, virus@AY632538.4—Iguape_virus_strain_SPAn_71686:1–10922



AeAeg_Aag2SINV_GFP.IP.18_23.rep

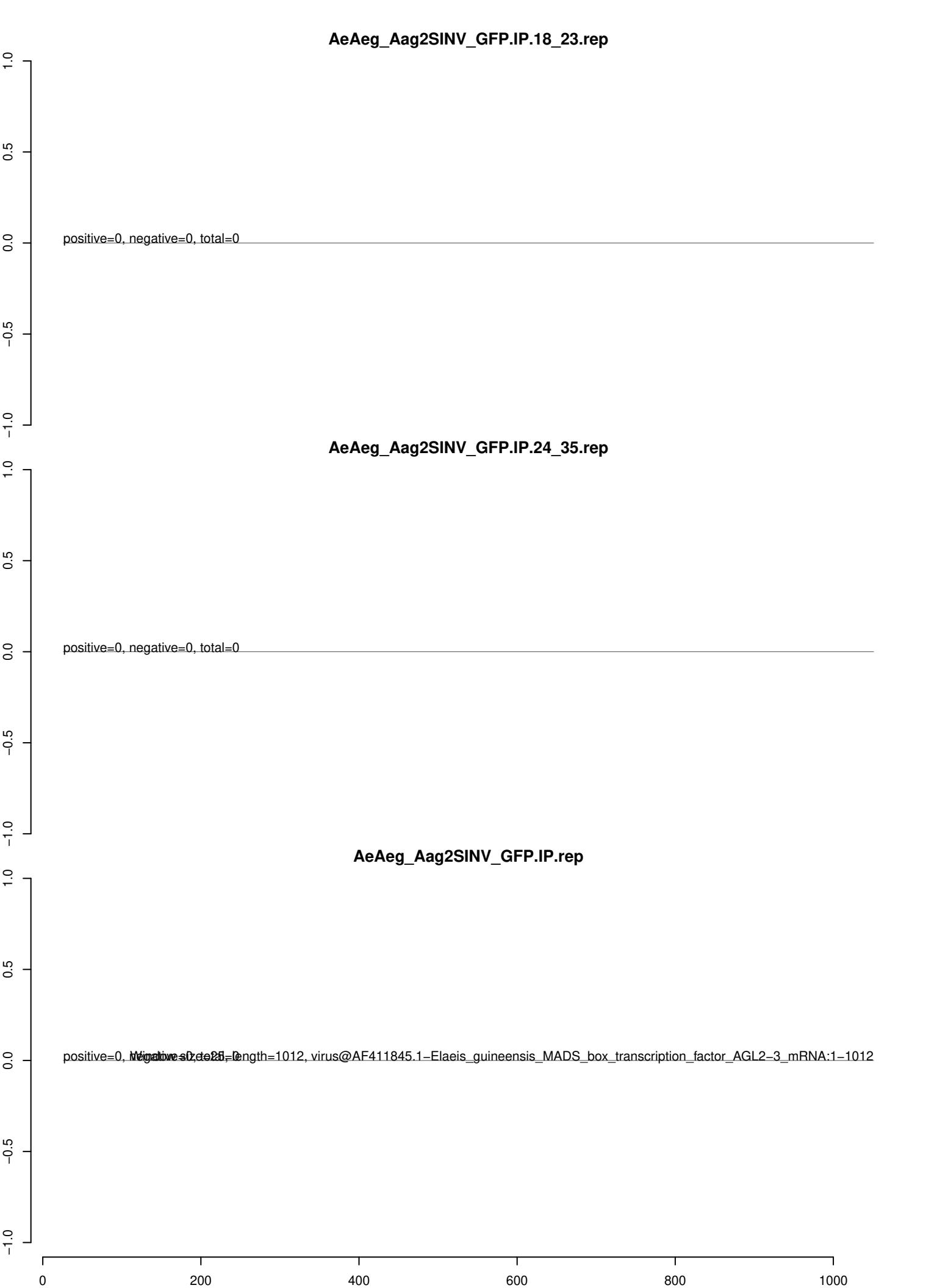
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

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



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=20, length=972, virus@AF411844.1-Elaeis_guineensis_MADS_box_transcription_factor_AGL2-2_mRNA:1-972

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

Window.size=25, length=11494, virus@AF075258.1–Rio_Negro_virus_strain_AG80–663:1–11494

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

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

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0

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

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0 Window size=25, length=11391, virus@AF075253.1-Mucambo_virus_strain_Mucambo_BeAn_8:1-11391

0 2000 4000 6000 8000 10000

AeAeg_Aag2SINV_GFP.IP.18_23.rep

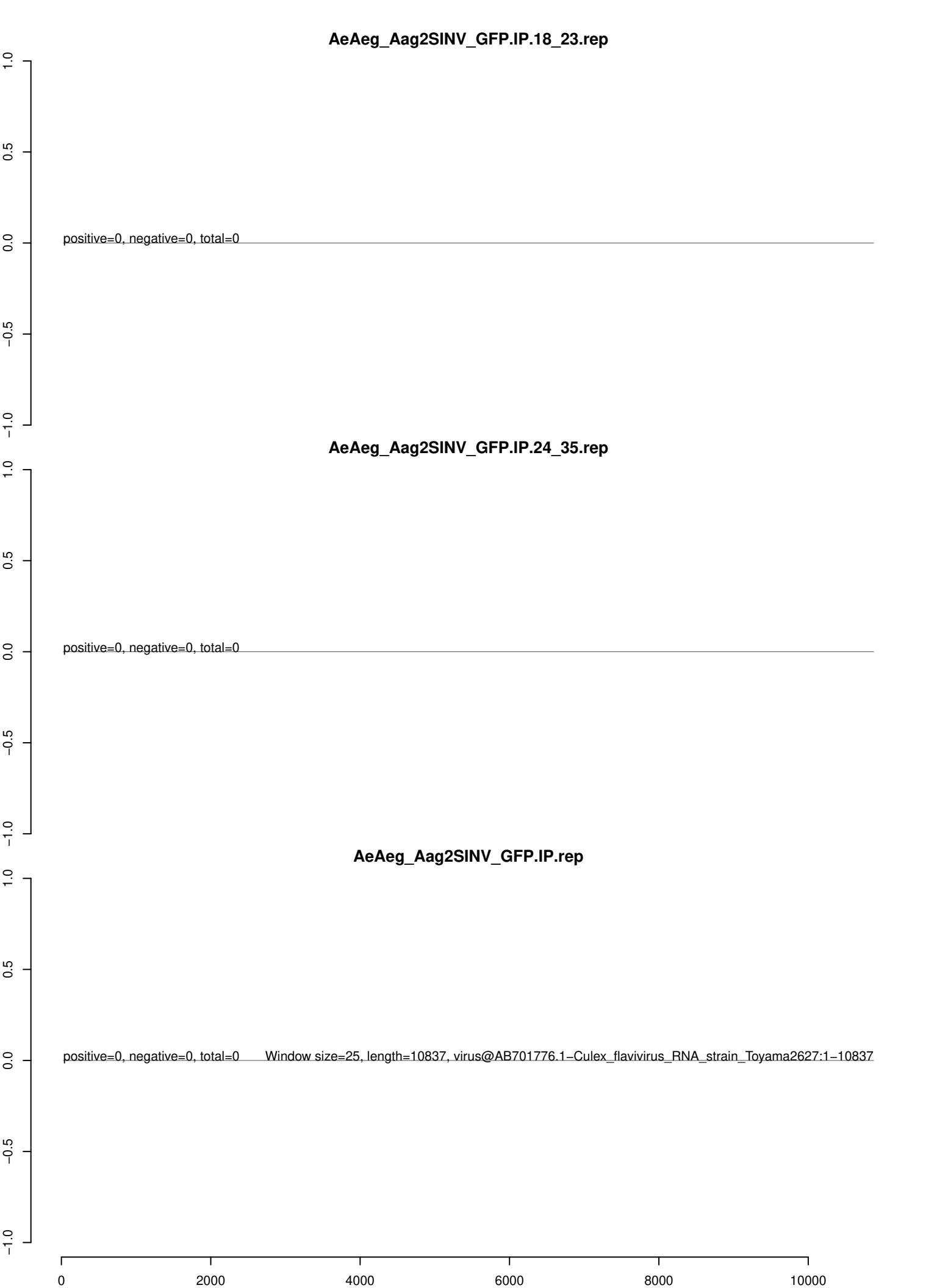
positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0 Window size=25, length=10837, virus@AB701776.1-Culex_flavivirus_RNA_strain_Toyama2627:1-10837



AeAeg_Aag2SINV_GFP.IP.18_23.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.24_35.rep

positive=0, negative=0, total=0

AeAeg_Aag2SINV_GFP.IP.rep

positive=0, negative=0, total=0 Window size=25, length=10837, virus@AB262759.2-Culex_flavivirus_genomic_RNA_strain_Tokyo:1-10837

