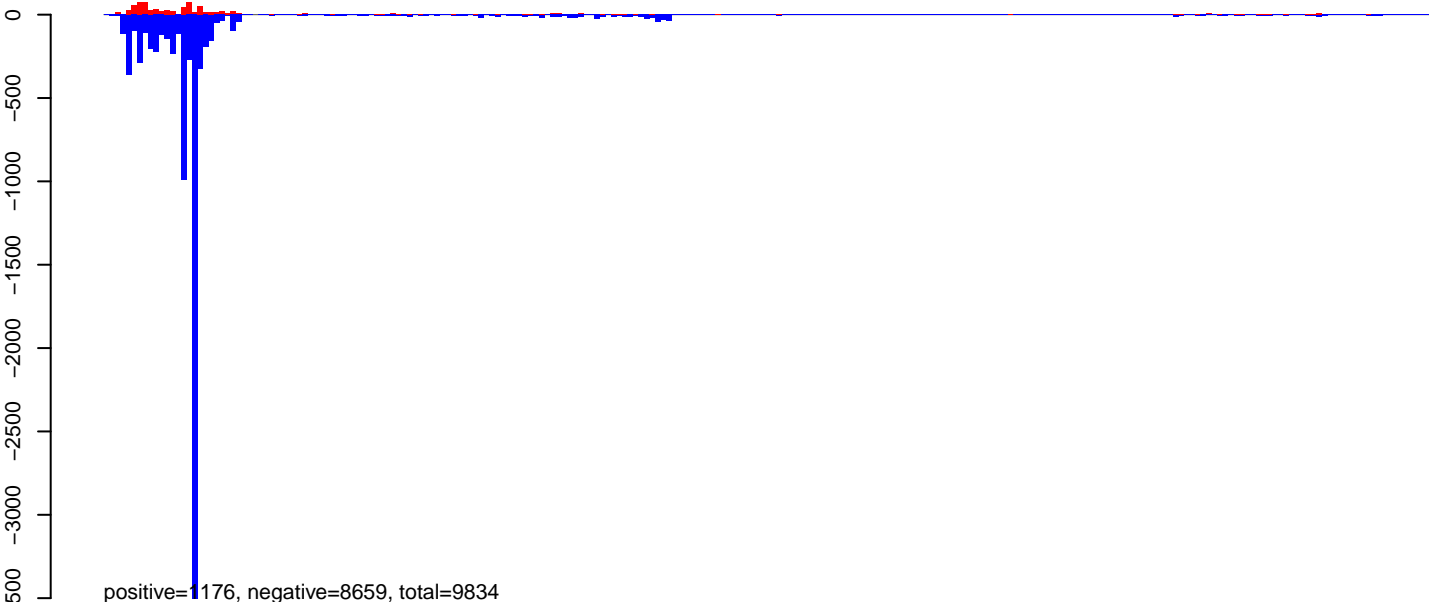


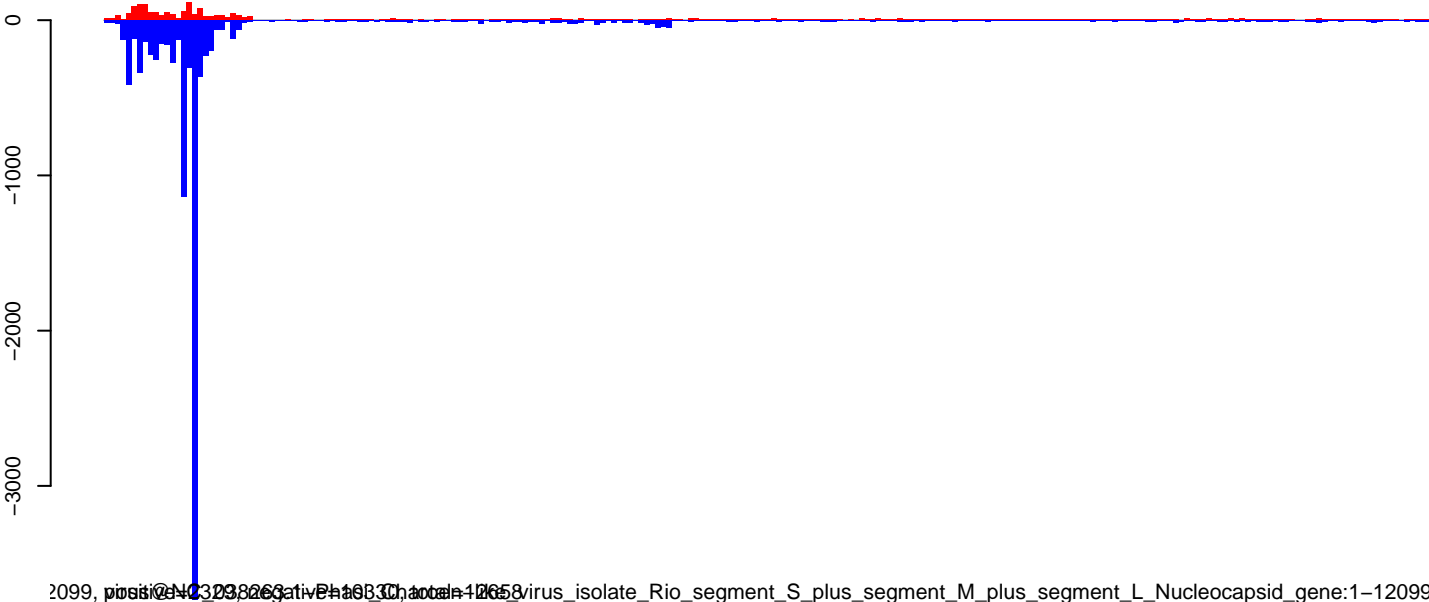
AeAeg_Aag2_DENV1.18_23.rep



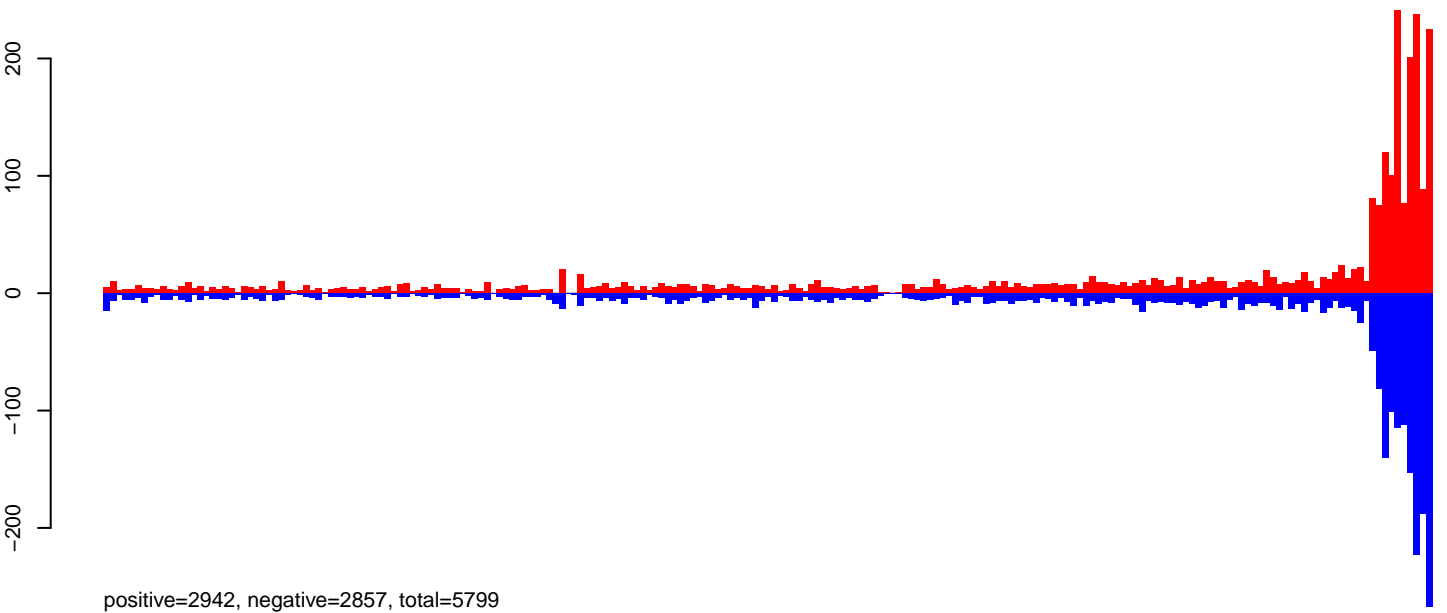
AeAeg_Aag2_DENV1.24_35.rep



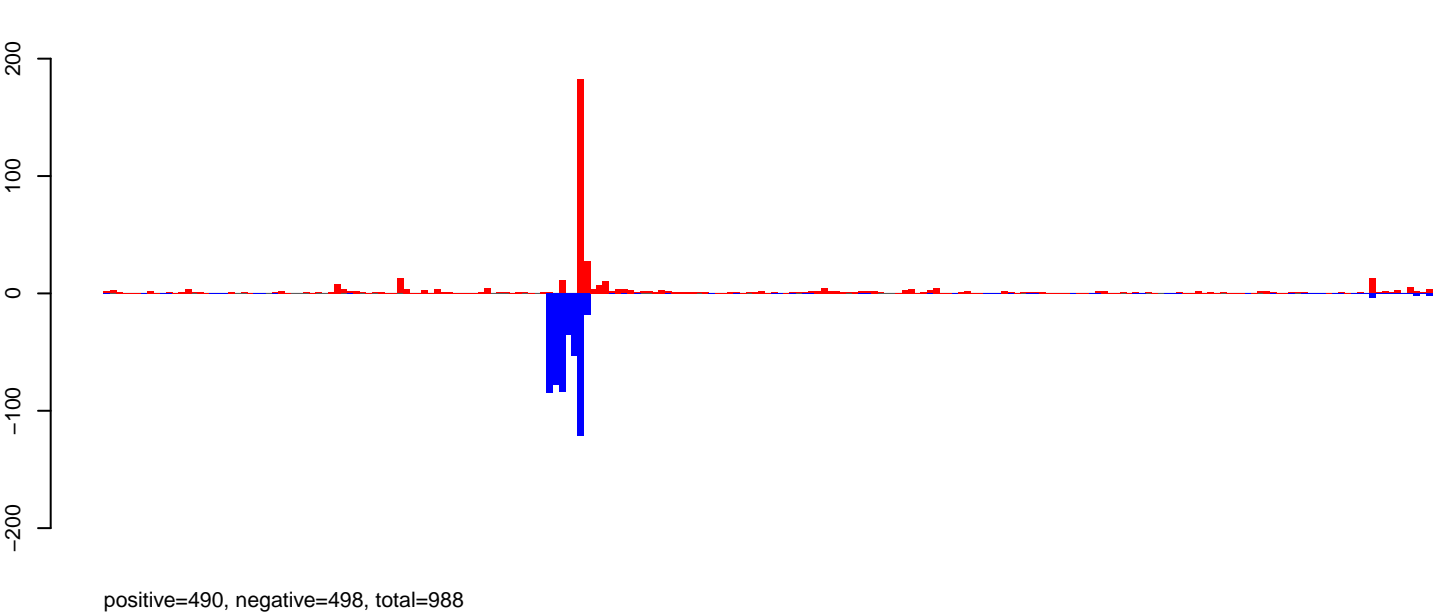
AeAeg_Aag2_DENV1.rep



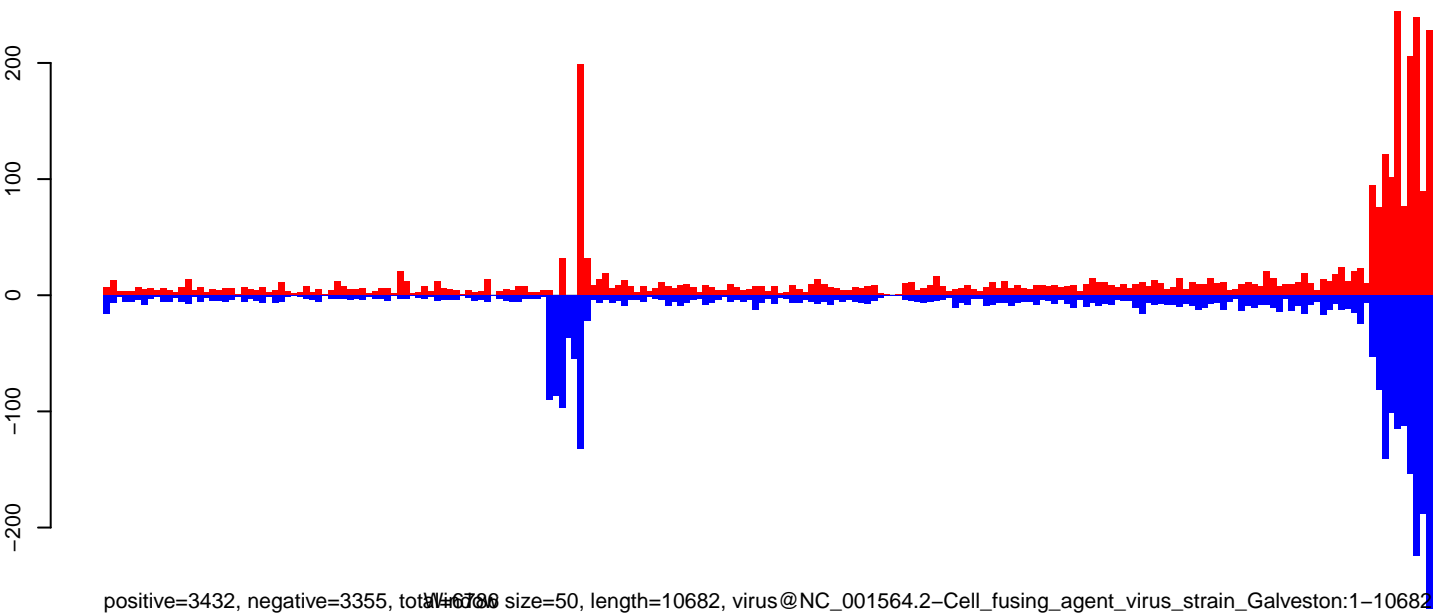
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

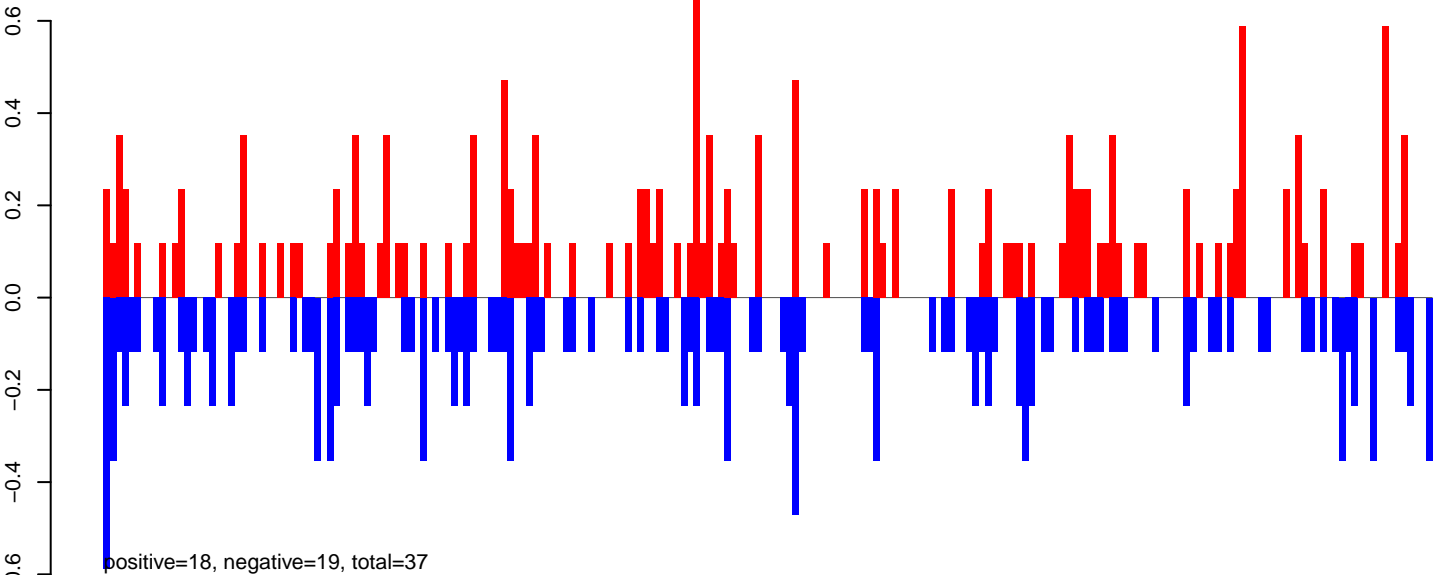


AeAeg_Aag2_DENV1.rep

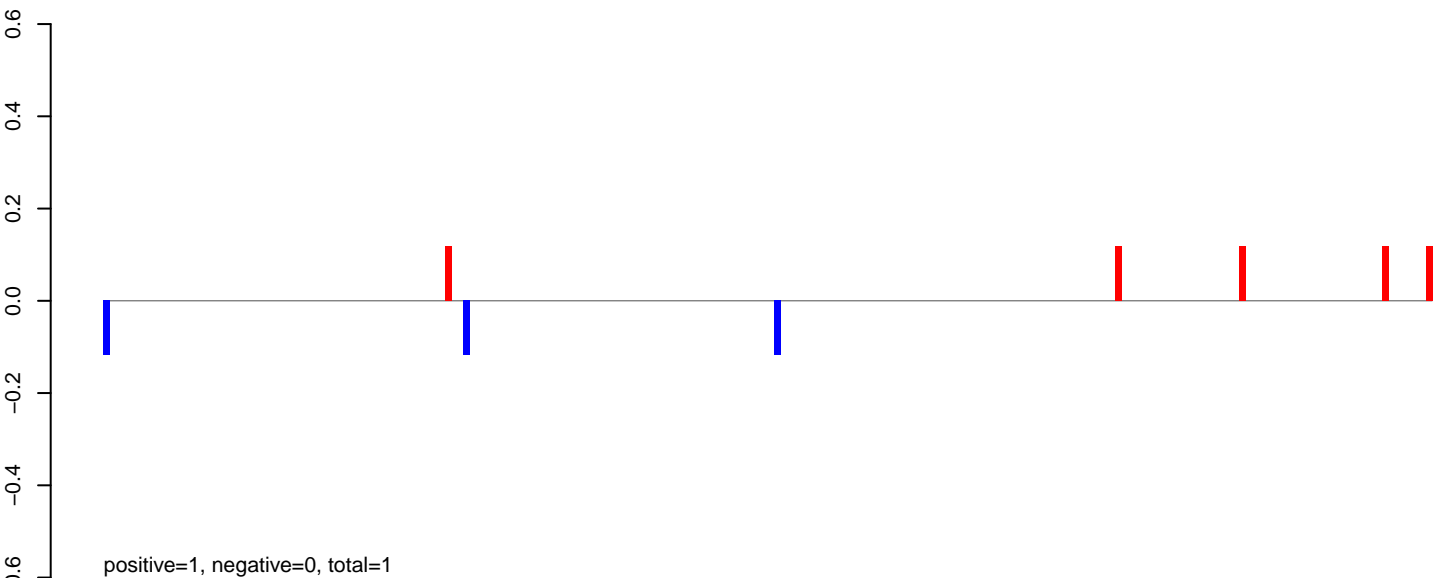


size=50, length=10682, virus@NC_001564.2-Cell_fusing_agent_virus_strain_Galveston:1-10682

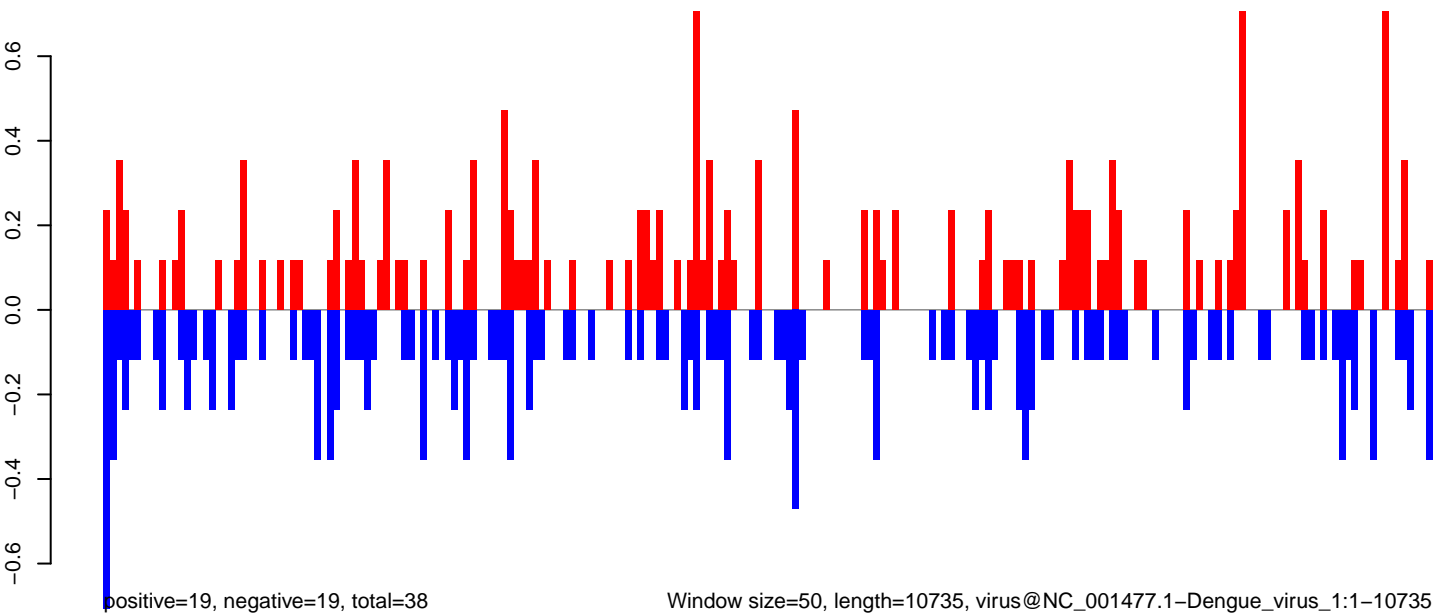
AeAeg_Aag2_DENV1.18_23.rep



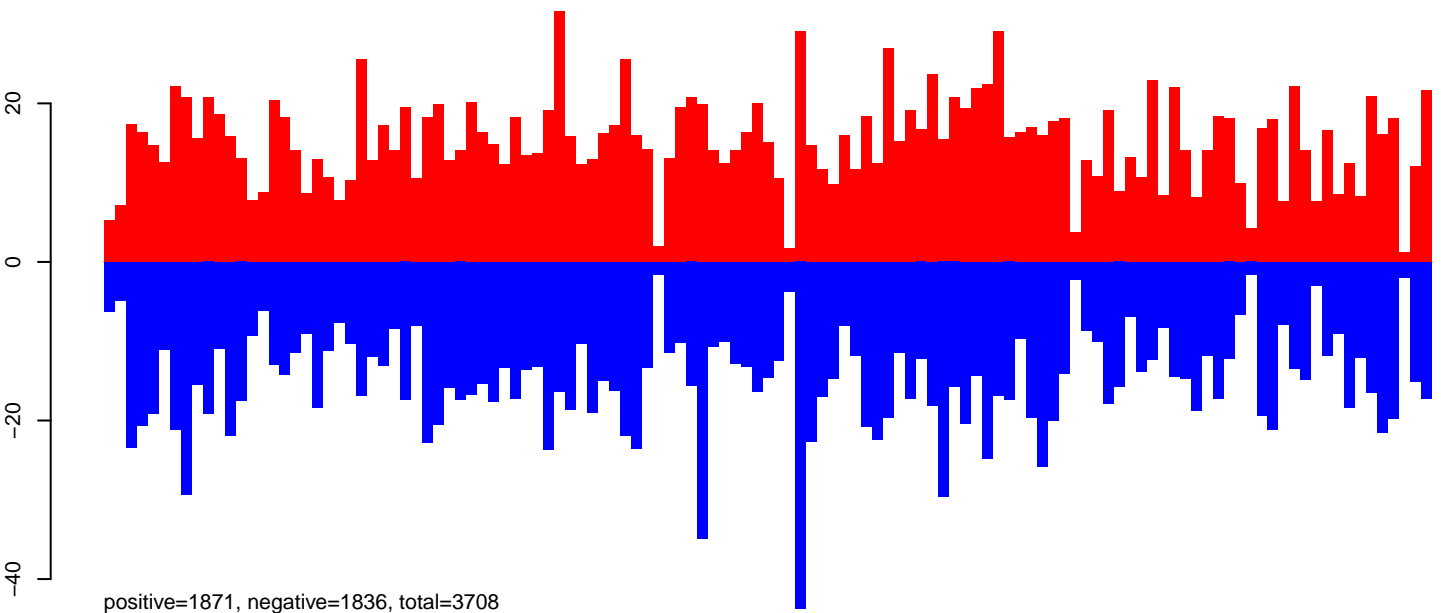
AeAeg_Aag2_DENV1.24_35.rep



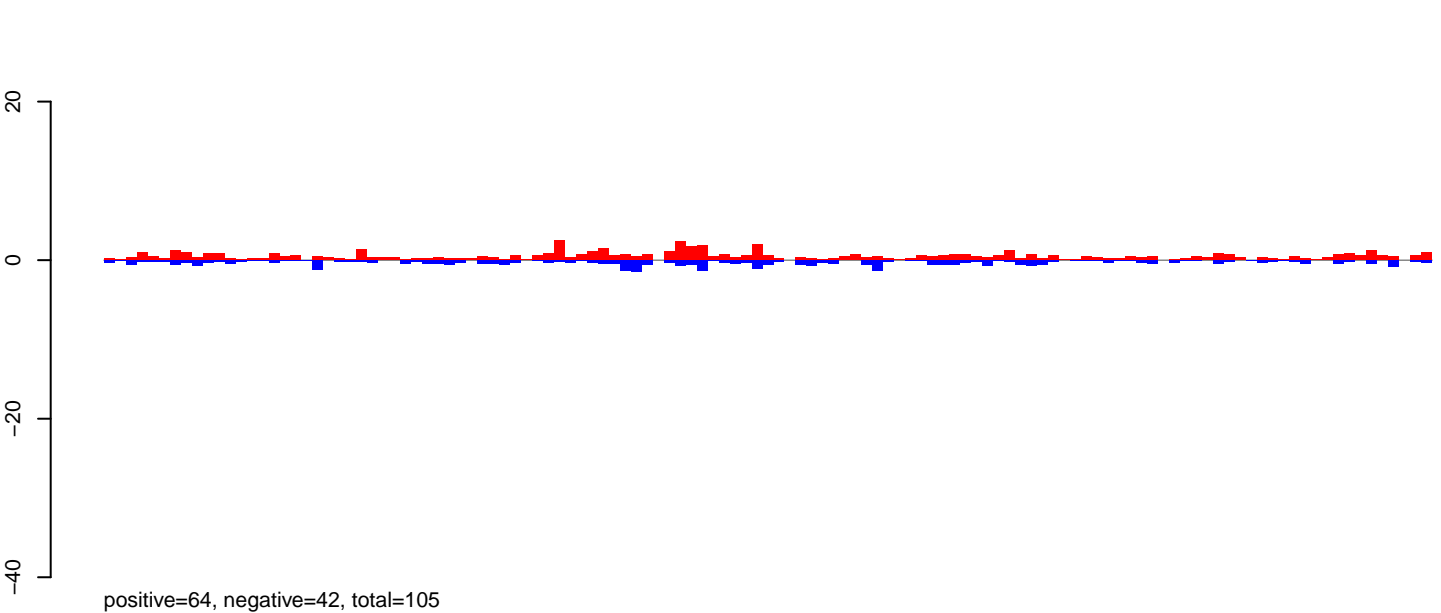
AeAeg_Aag2_DENV1.rep



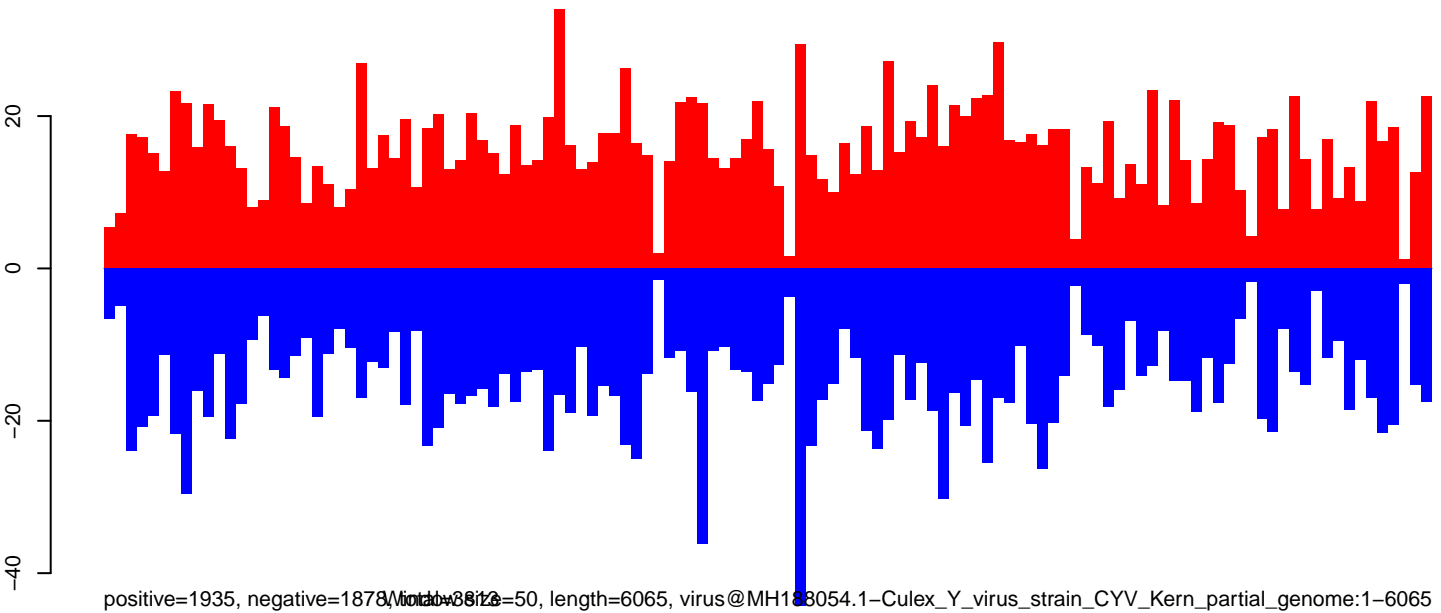
AeAeg_Aag2_DENV1.18_23.rep



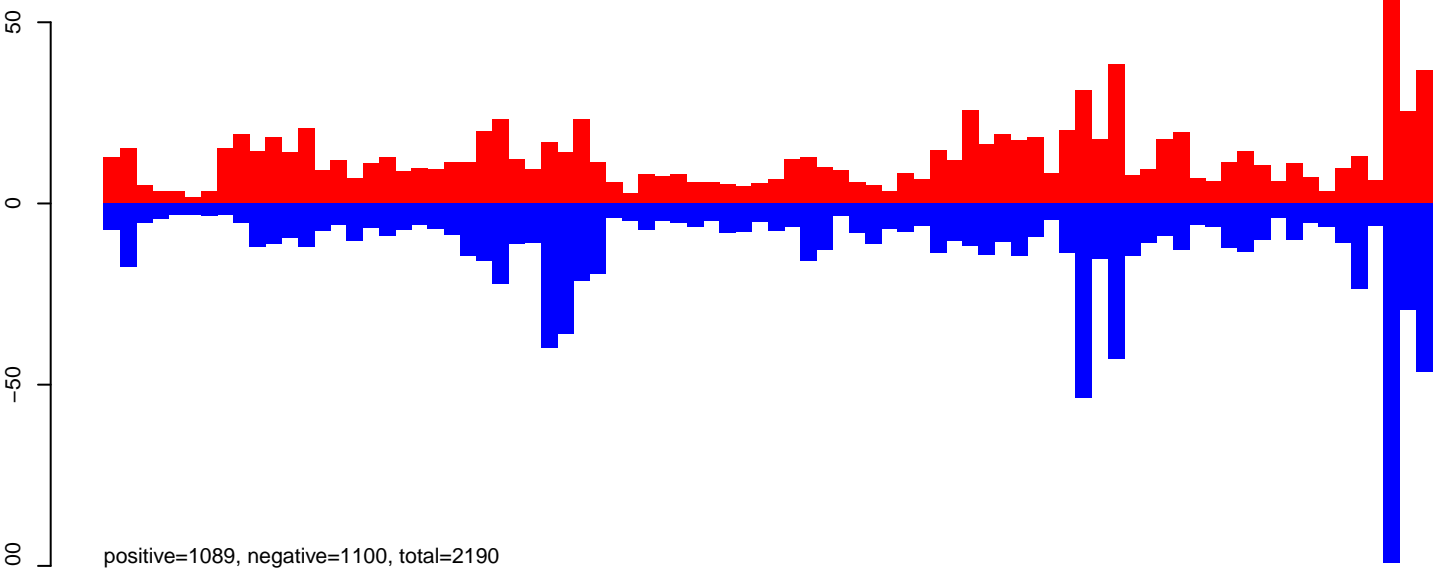
AeAeg_Aag2_DENV1.24_35.rep



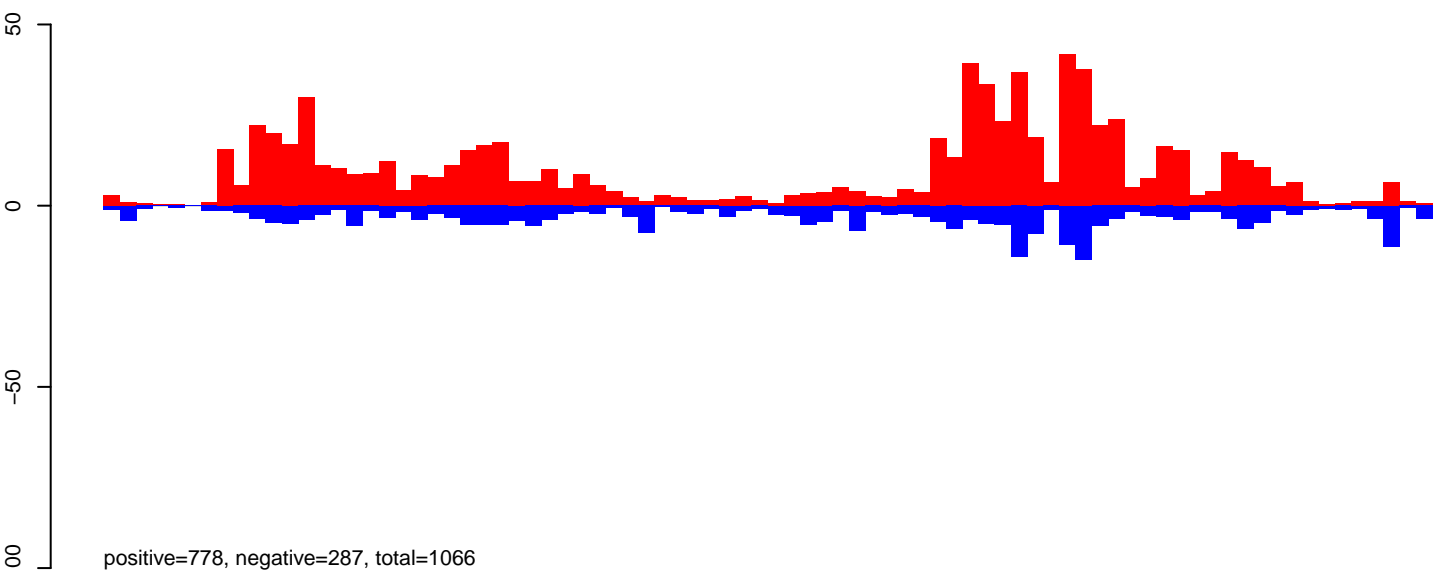
AeAeg_Aag2_DENV1.rep



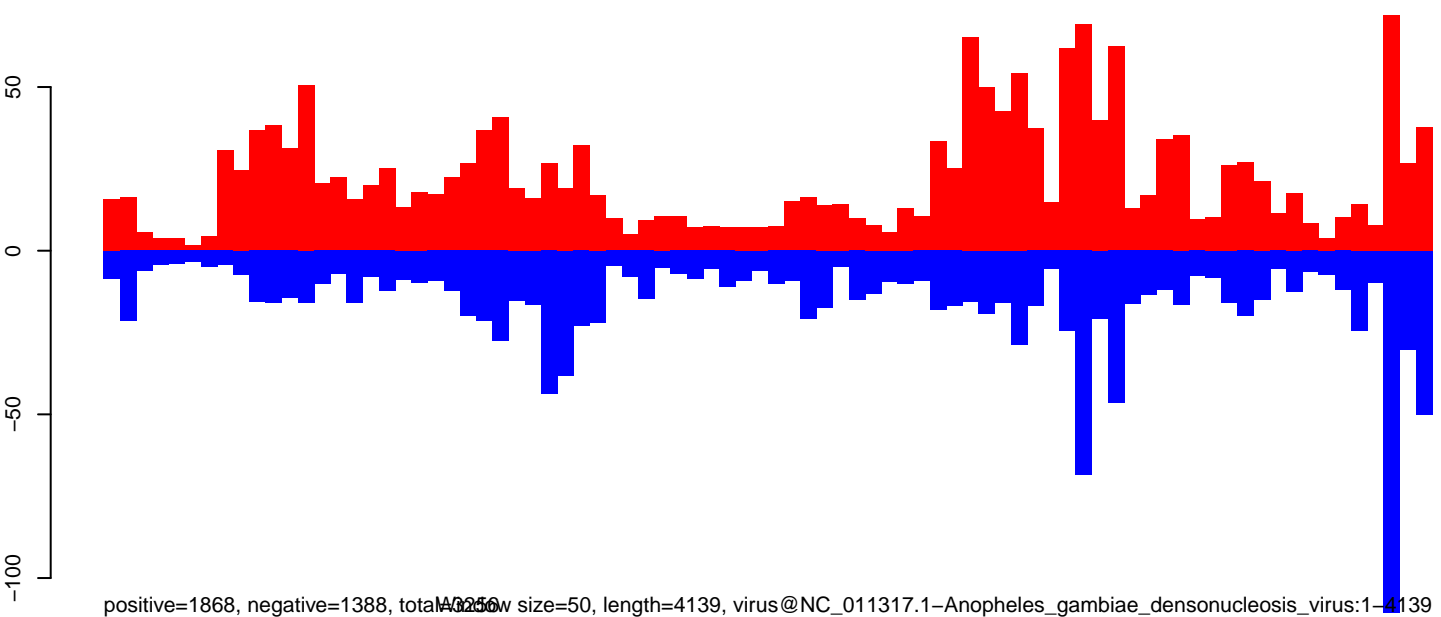
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

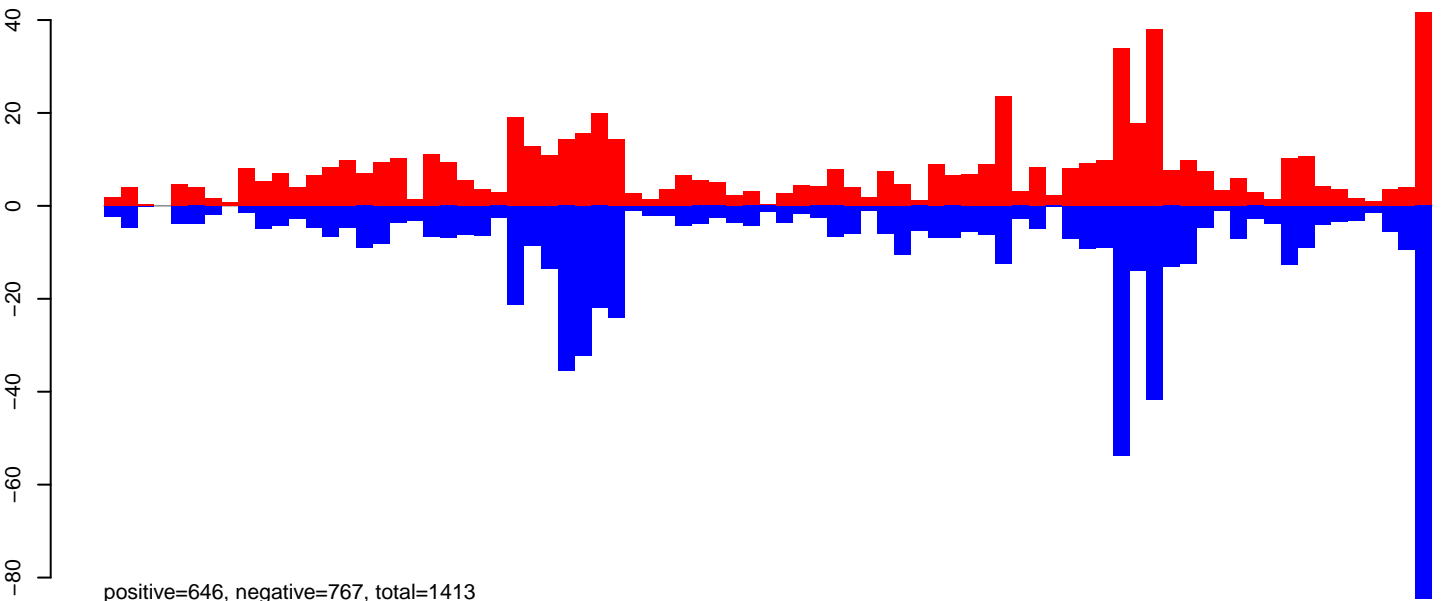


AeAeg_Aag2_DENV1.rep

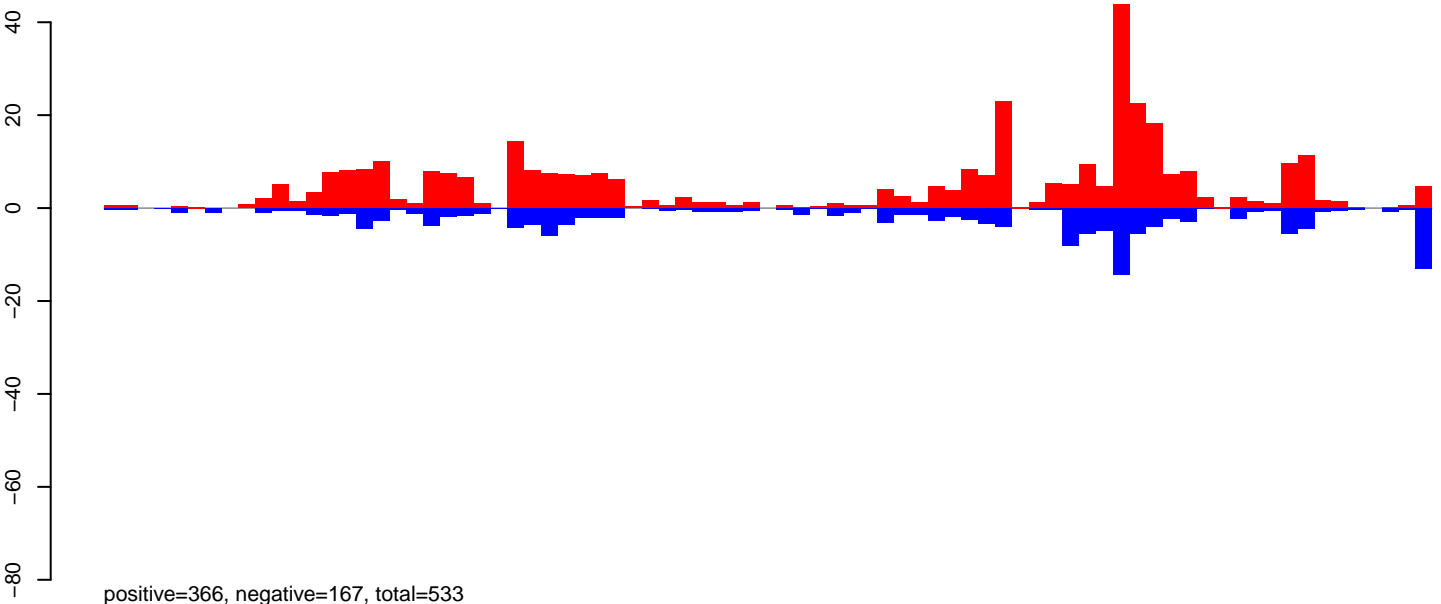


Window size=50, length=4139, virus@NC_011317.1-Anopheles_gambiae_denonucleosis_virus:1-4139

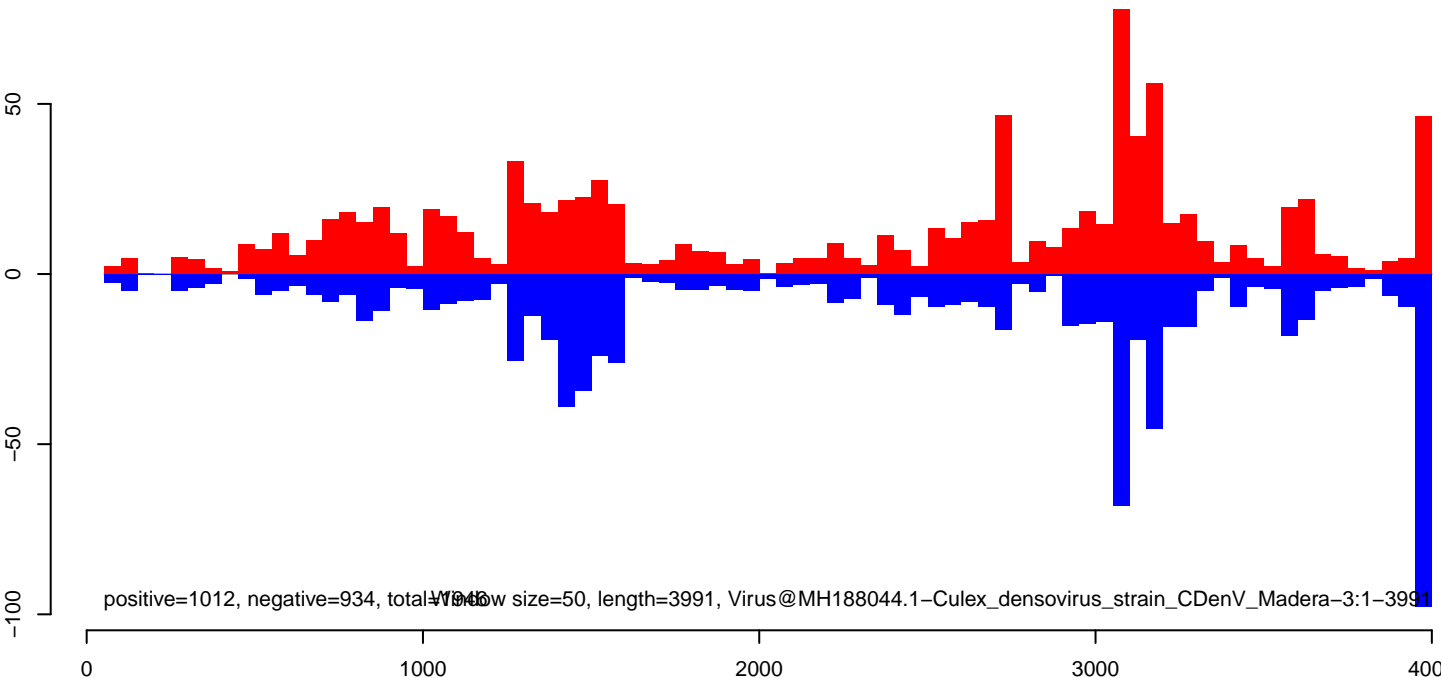
AeAeg_Aag2_DENV1.18_23.rep



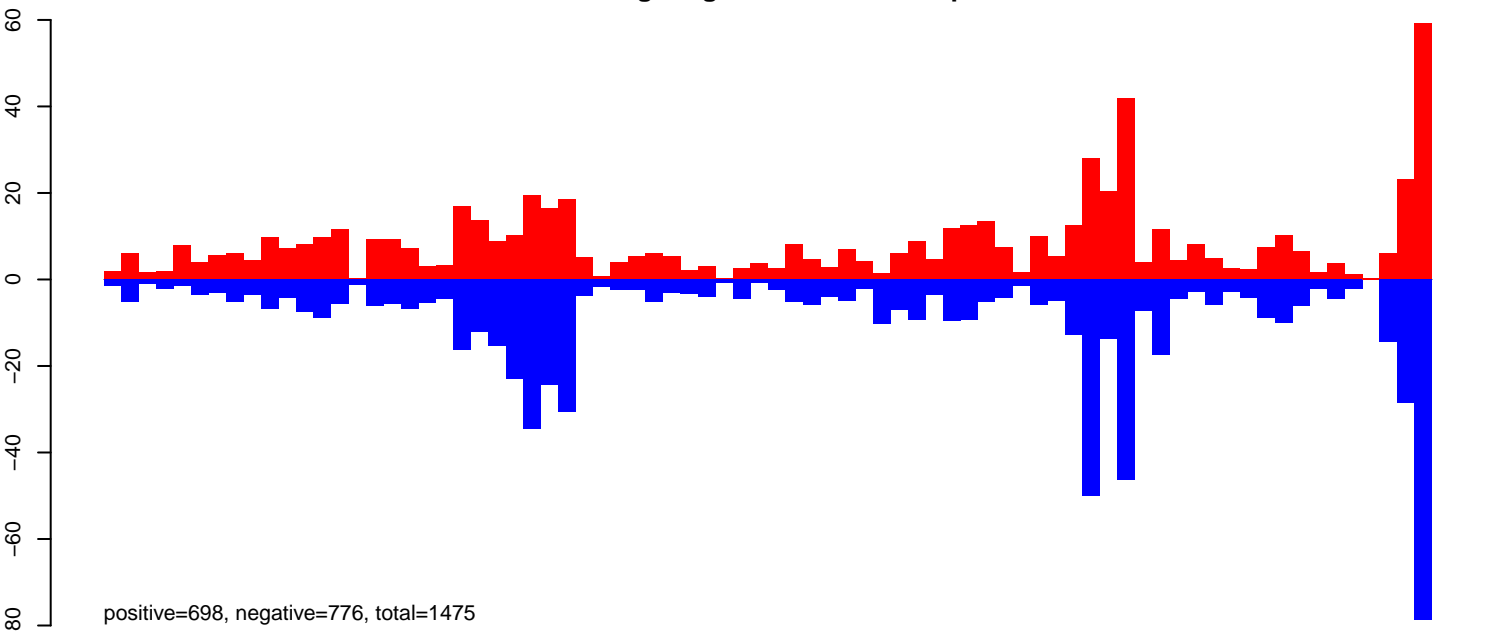
AeAeg_Aag2_DENV1.24_35.rep



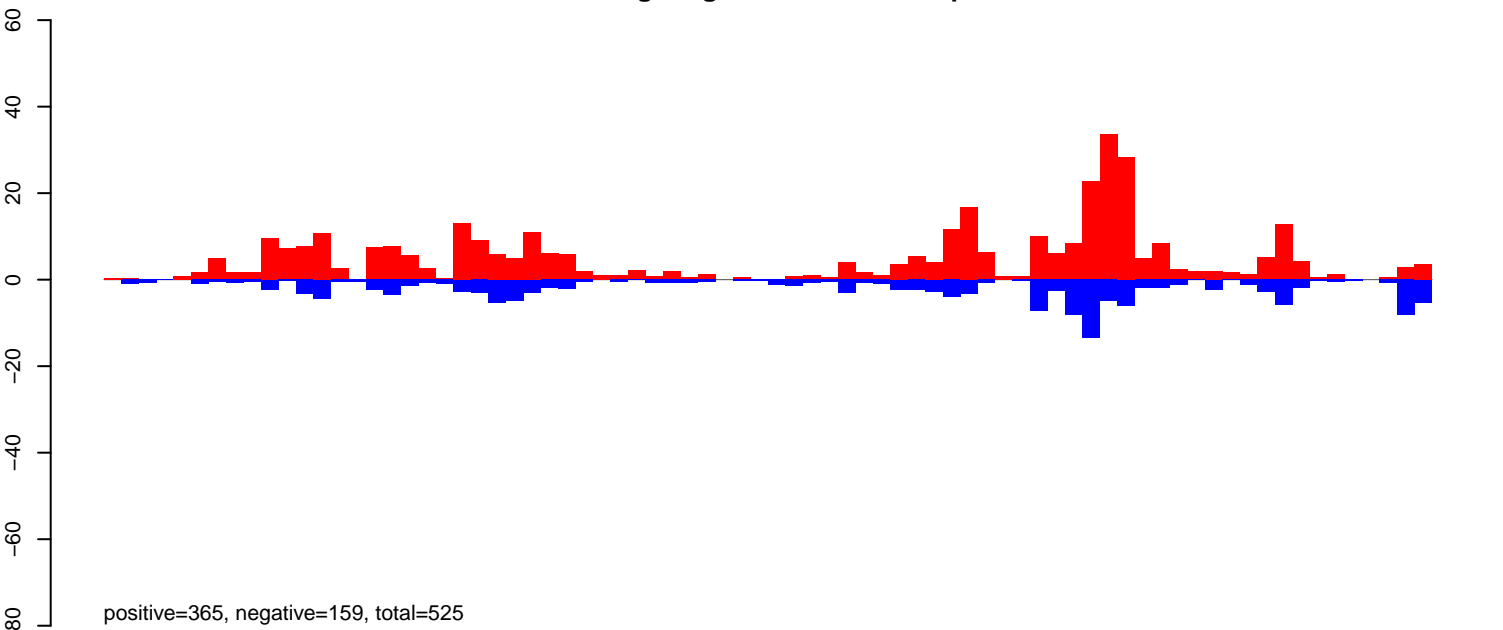
AeAeg_Aag2_DENV1.rep



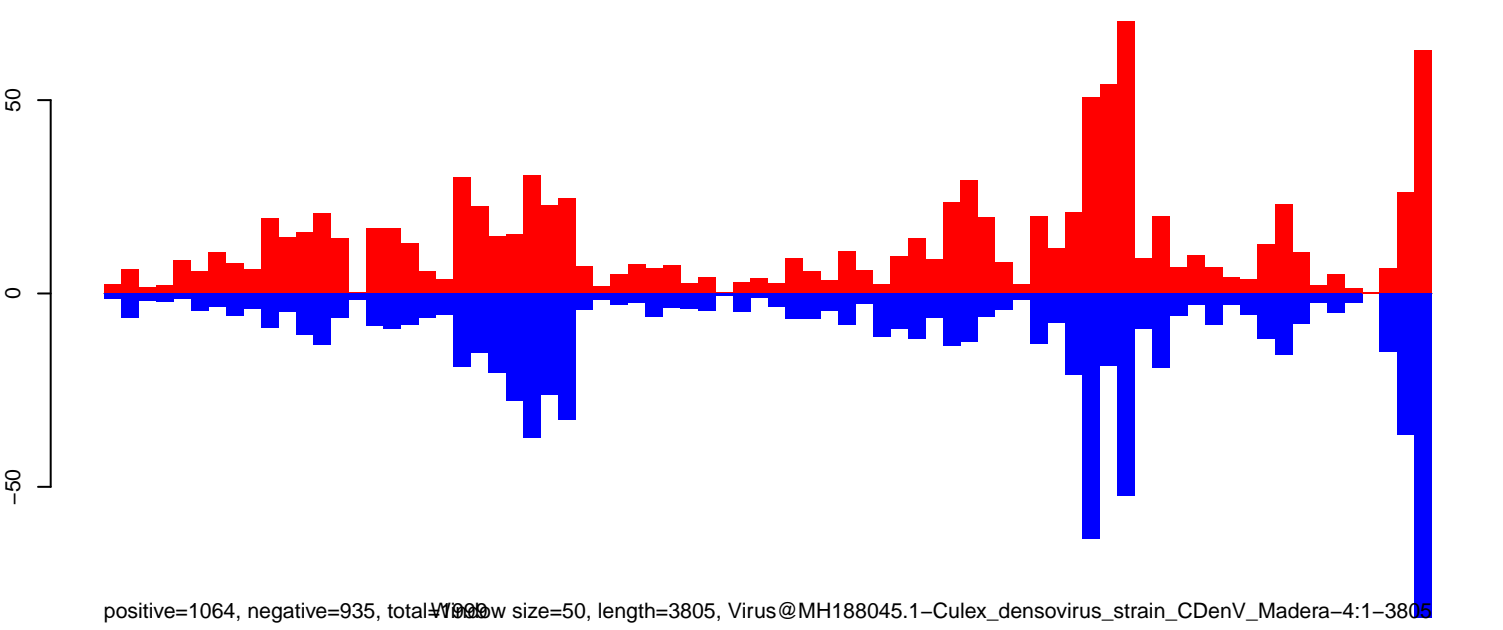
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

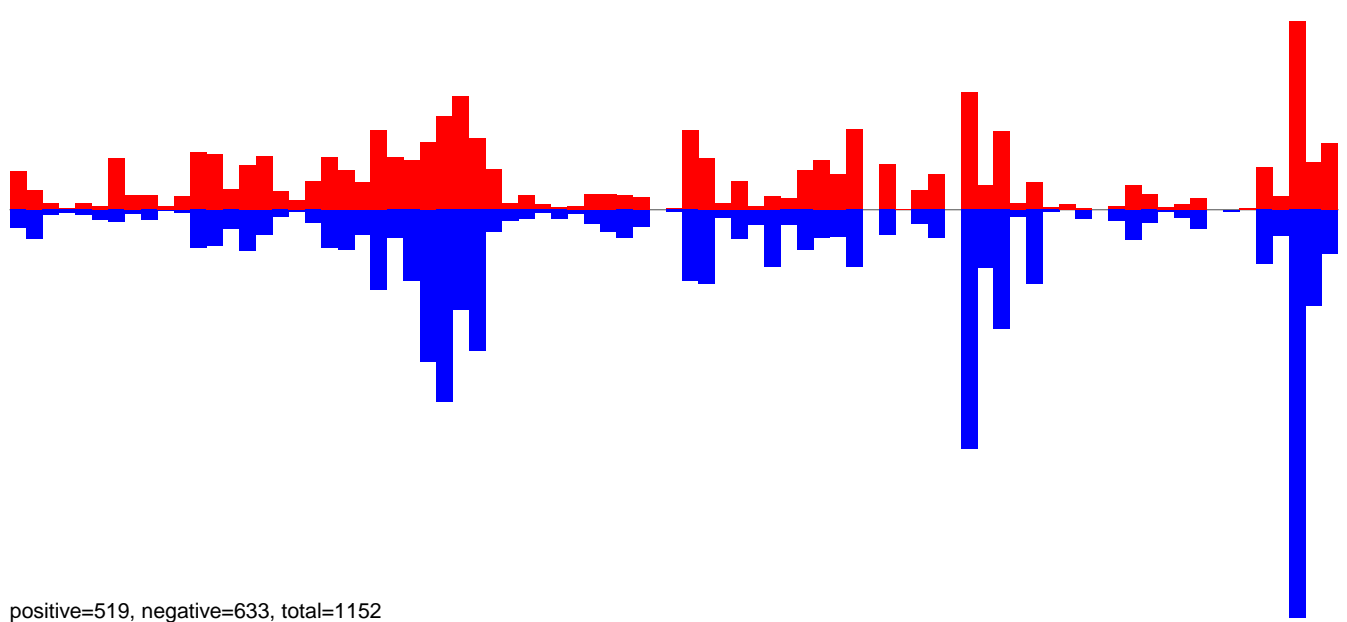


AeAeg_Aag2_DENV1.rep

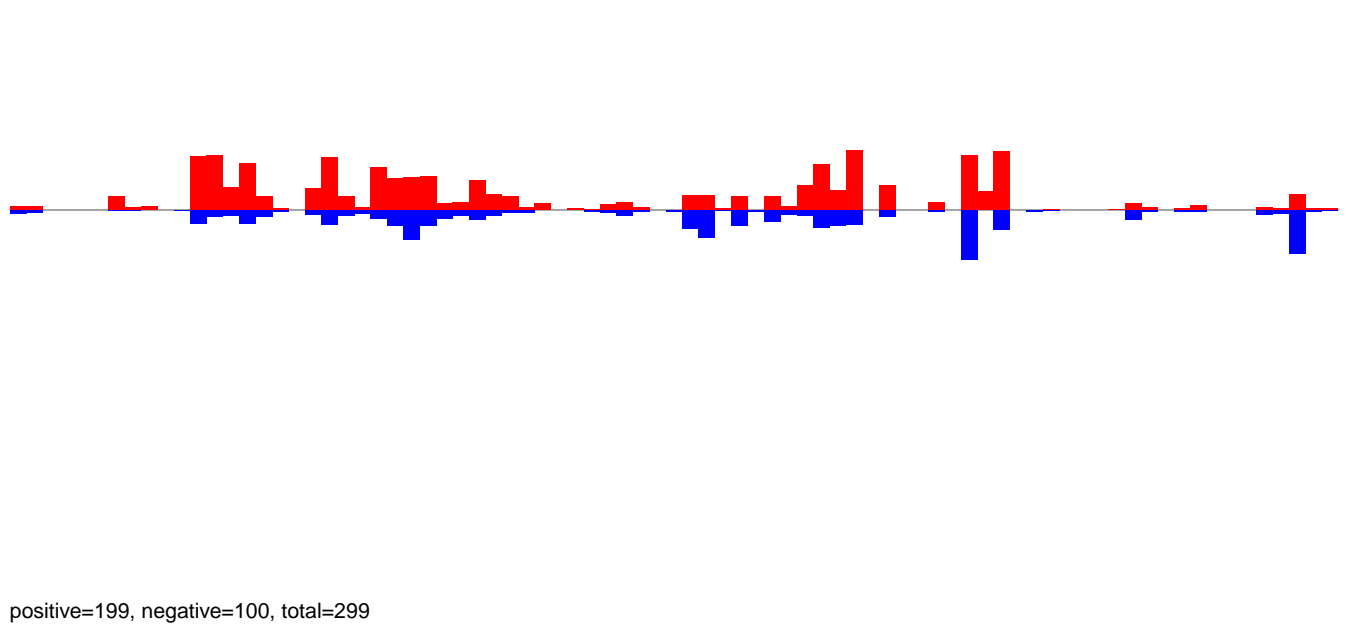


Window size=50, length=3805, Virus@MH188045.1-Culex_densovirus_strain_CDenV_Madera-4:1-3805

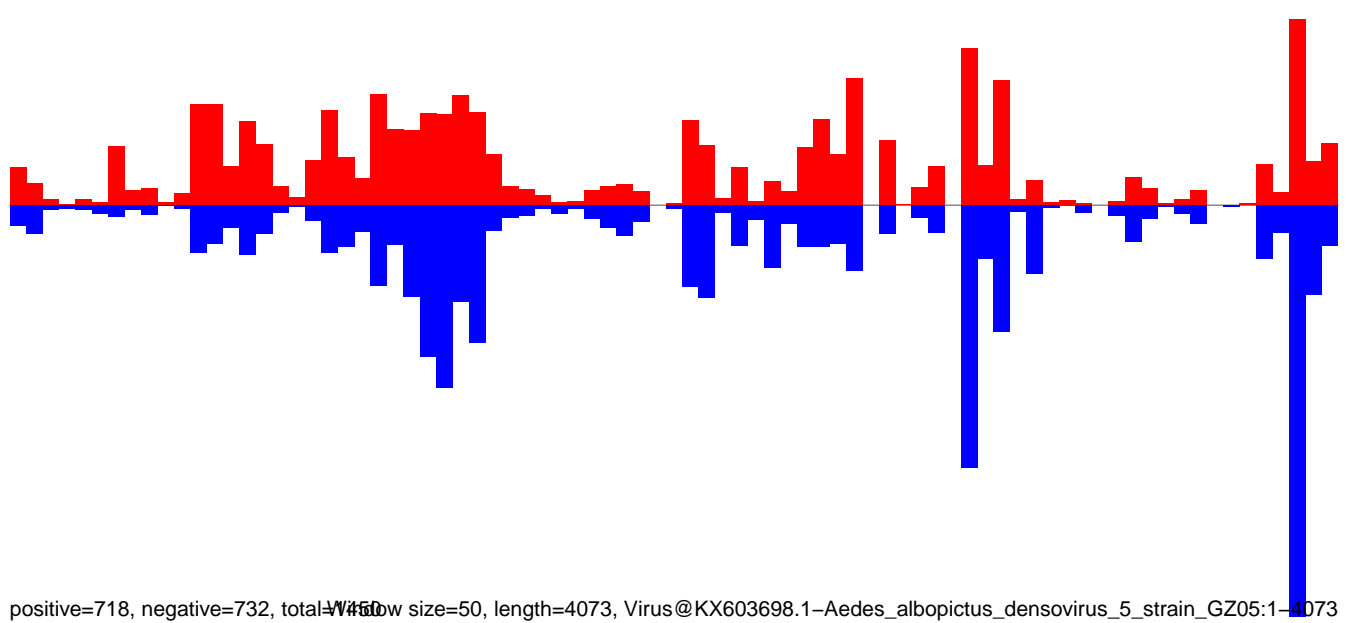
AeAeg_Aag2_DENV1.18_23.rep



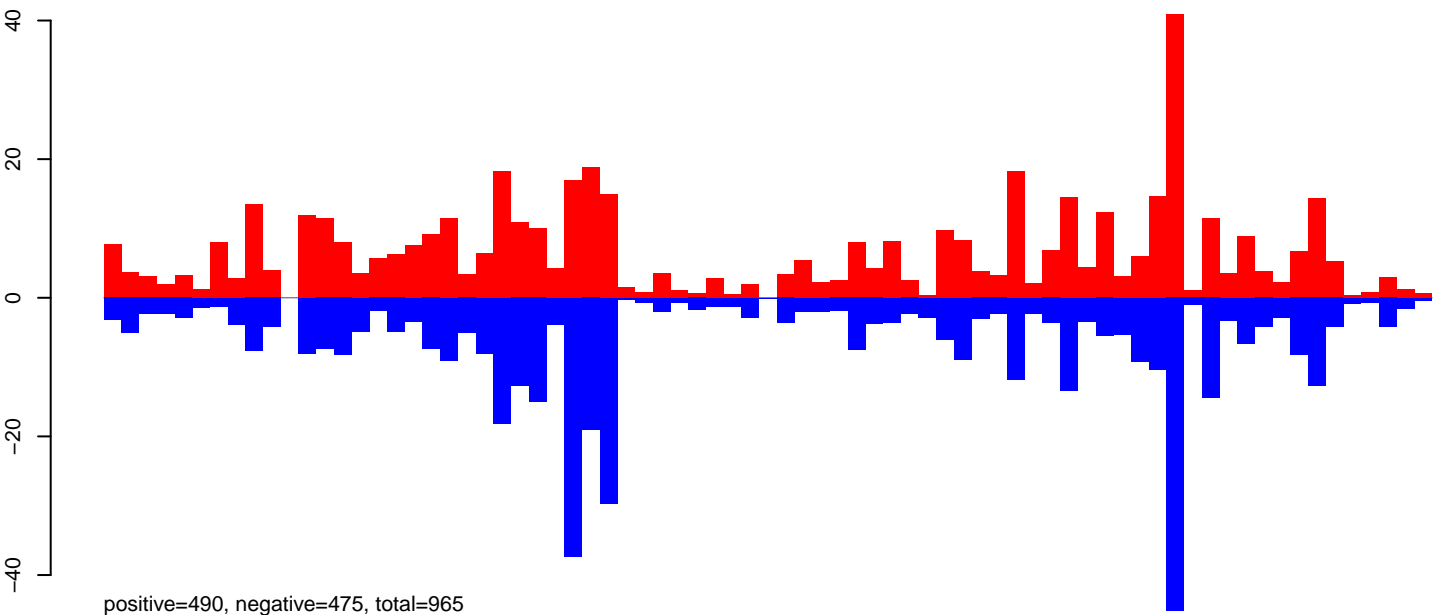
AeAeg_Aag2_DENV1.24_35.rep



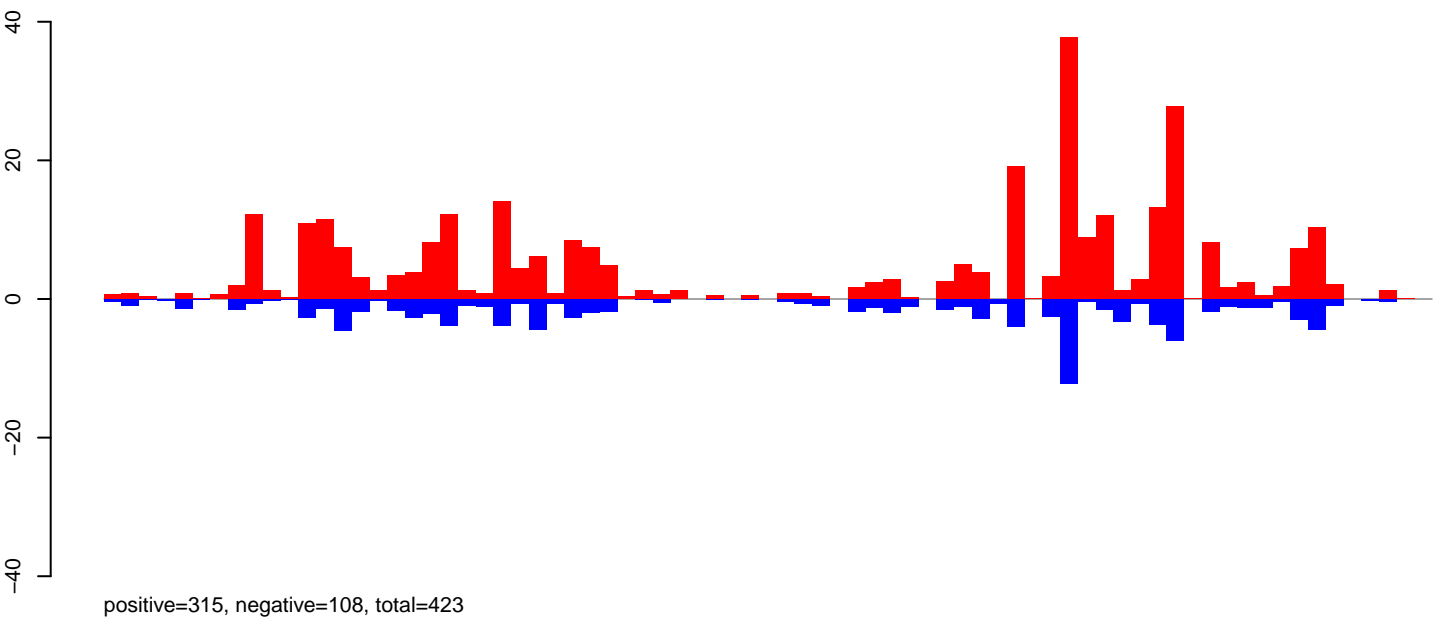
AeAeg_Aag2_DENV1.rep



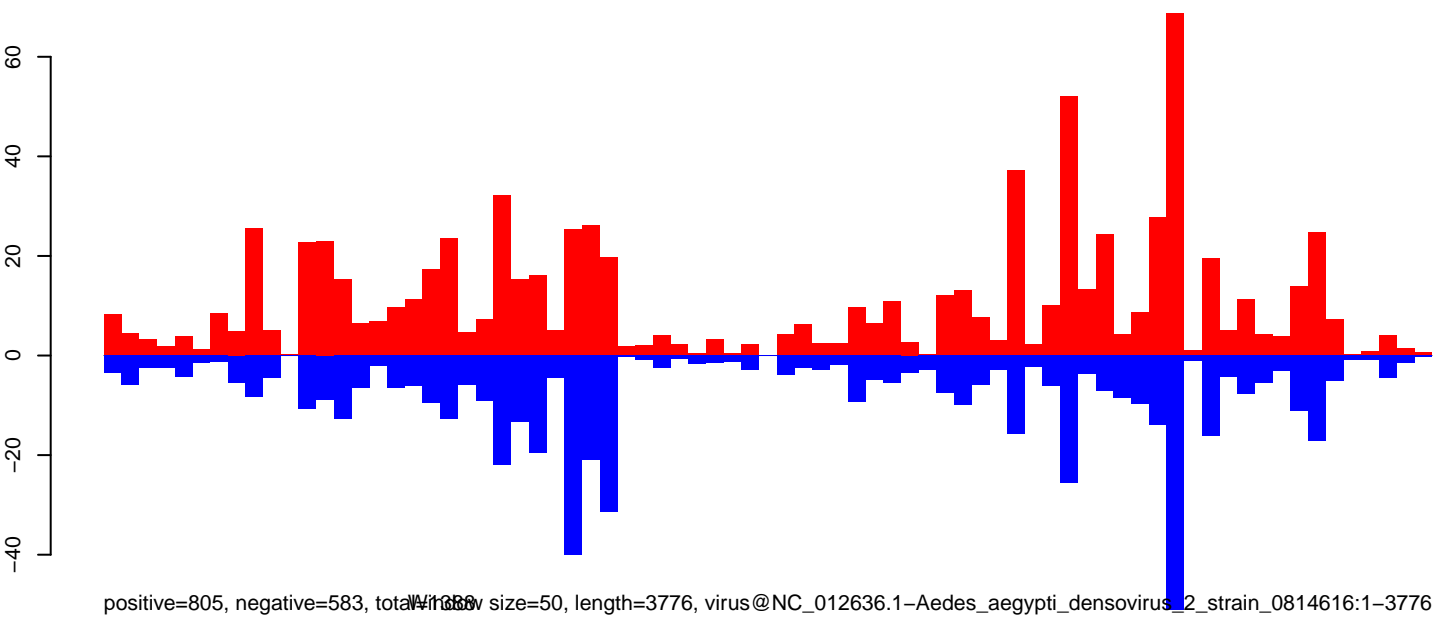
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

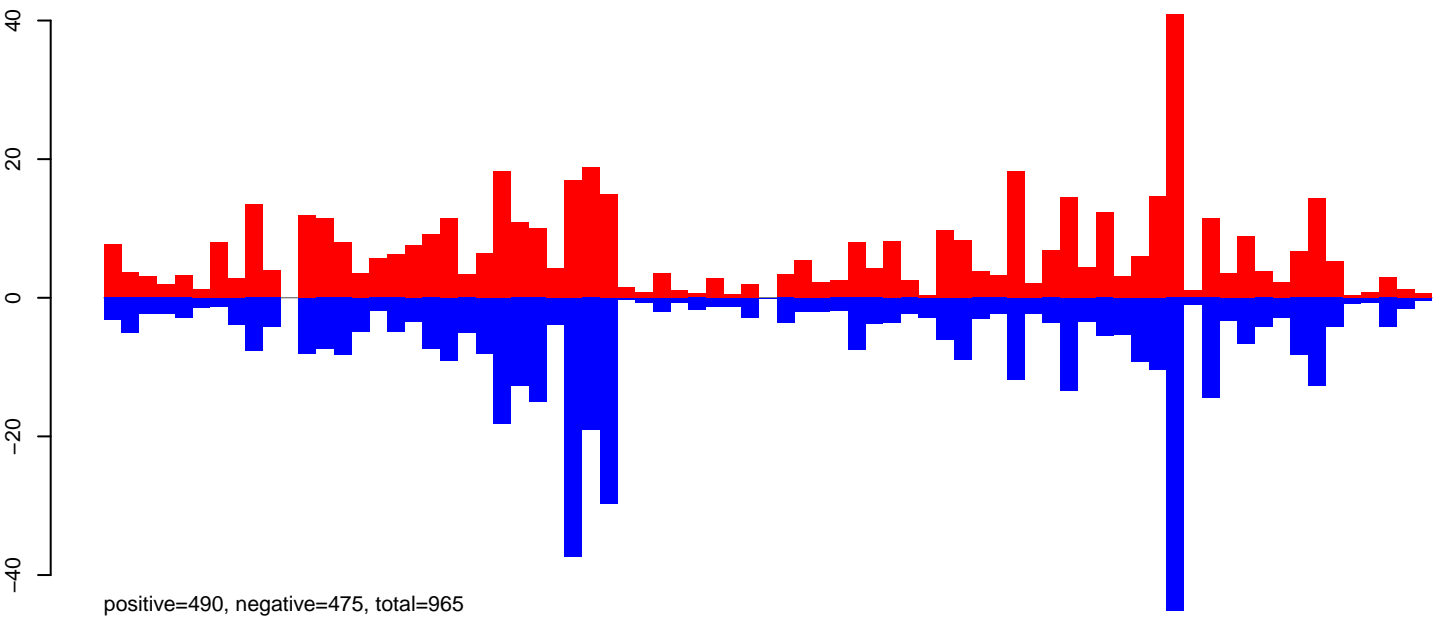


AeAeg_Aag2_DENV1.rep

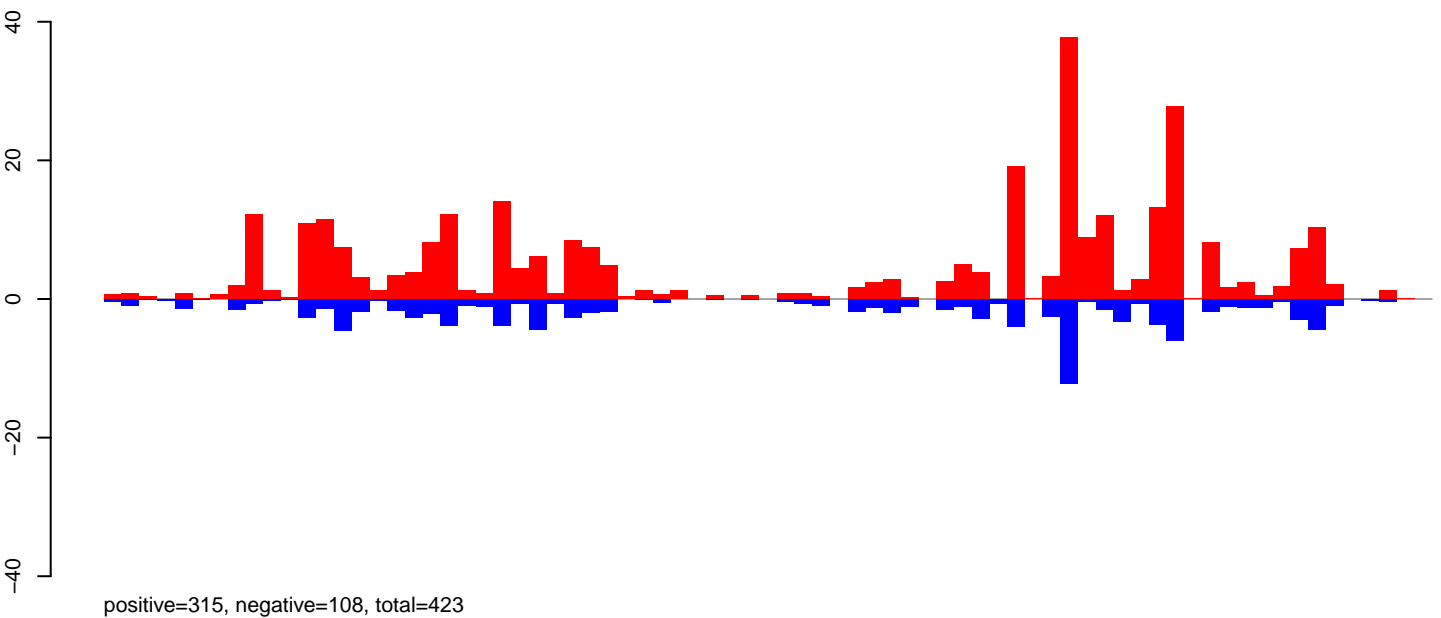


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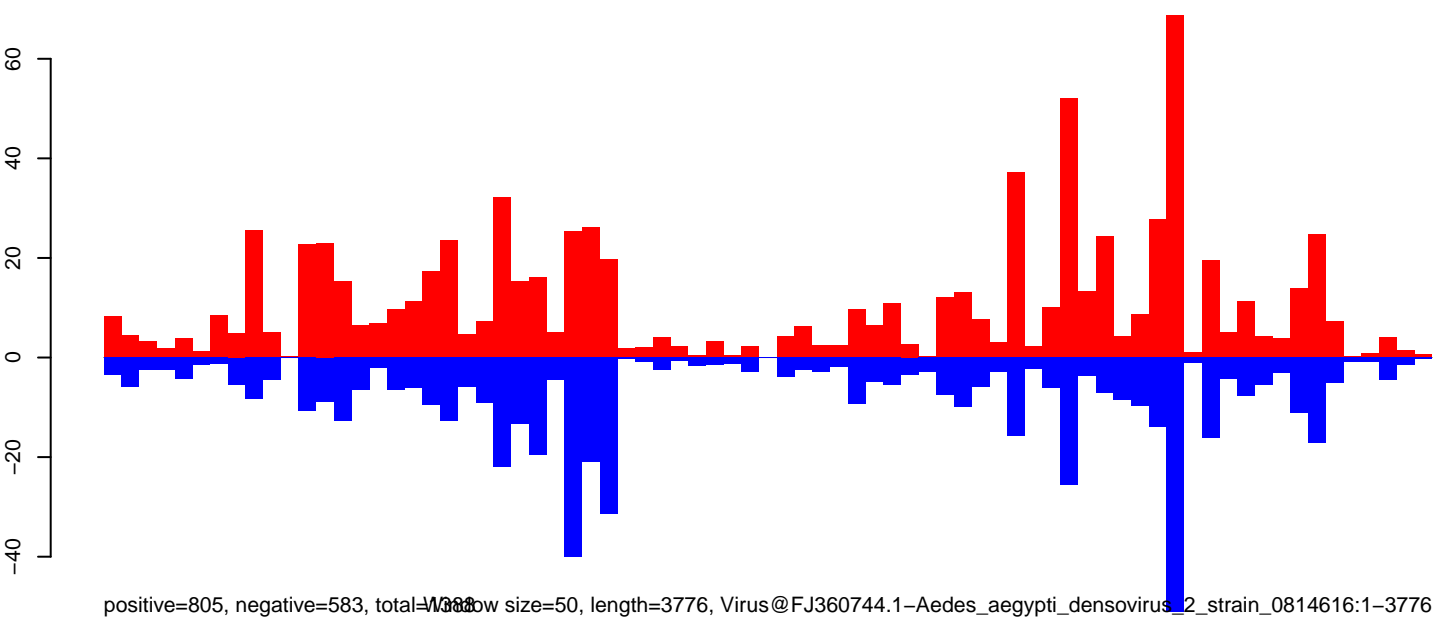
AeAeg_Aag2_DENV1.18_23.rep



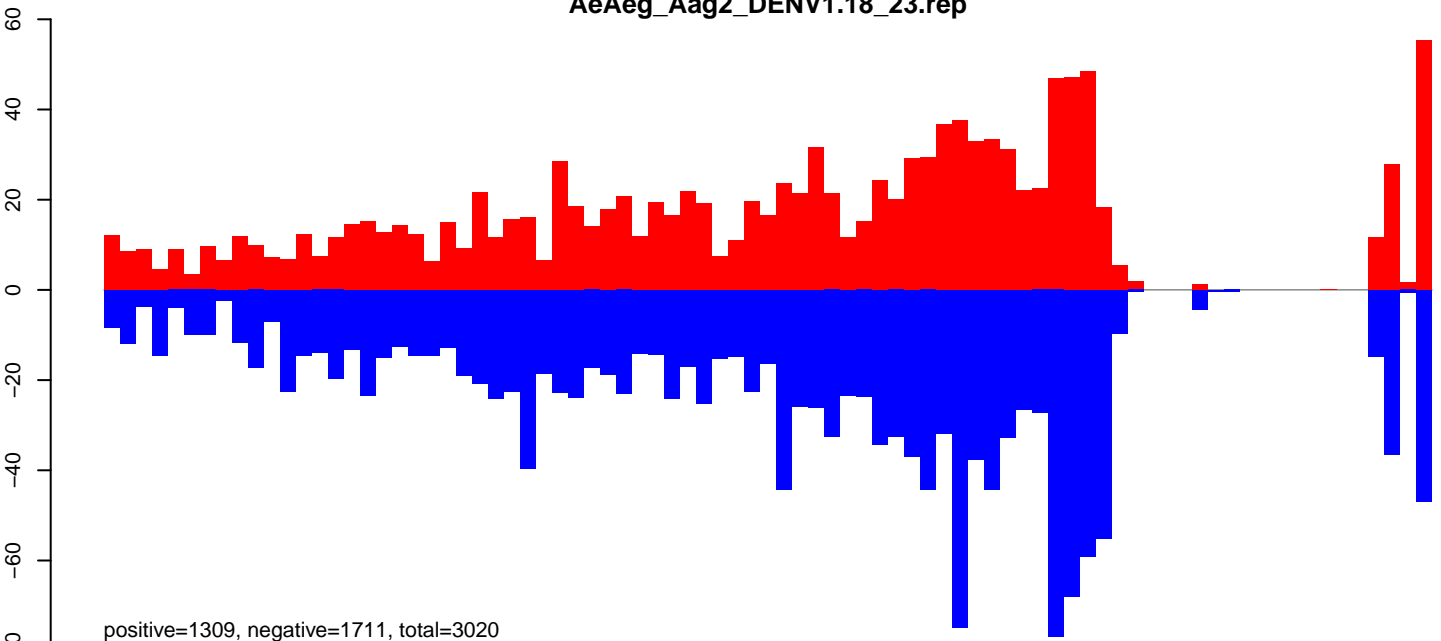
AeAeg_Aag2_DENV1.24_35.rep



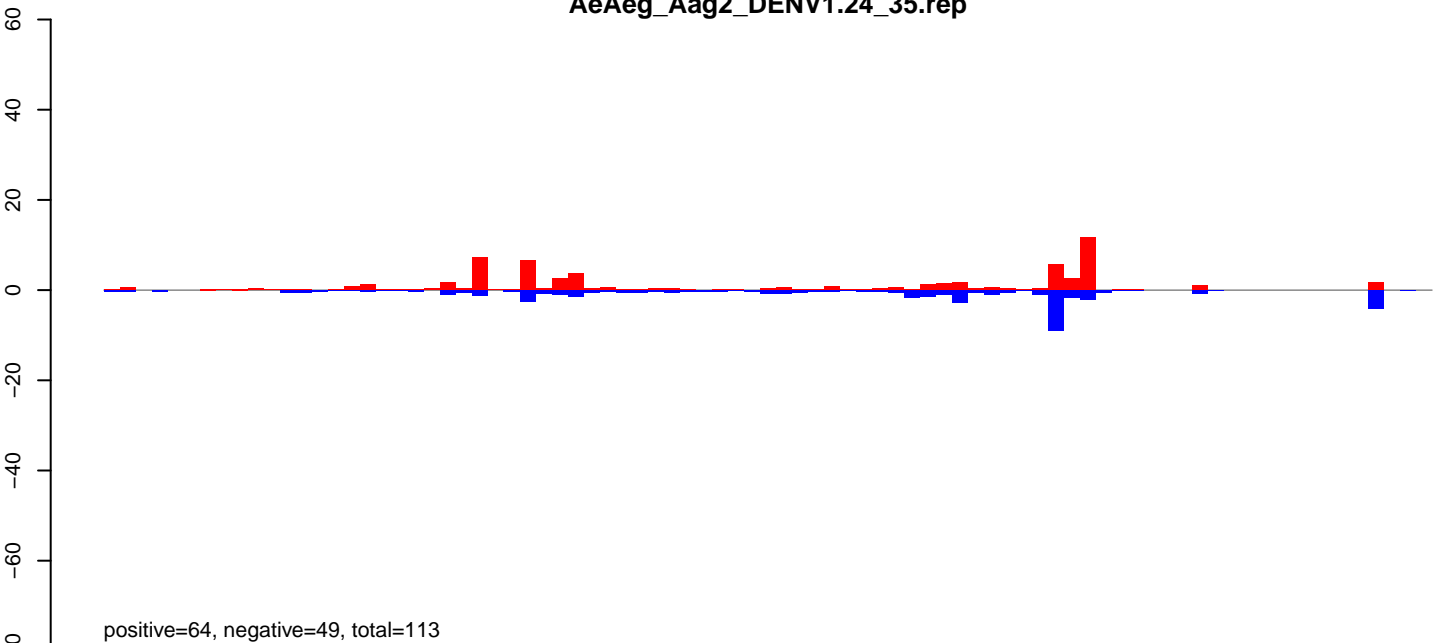
AeAeg_Aag2_DENV1.rep



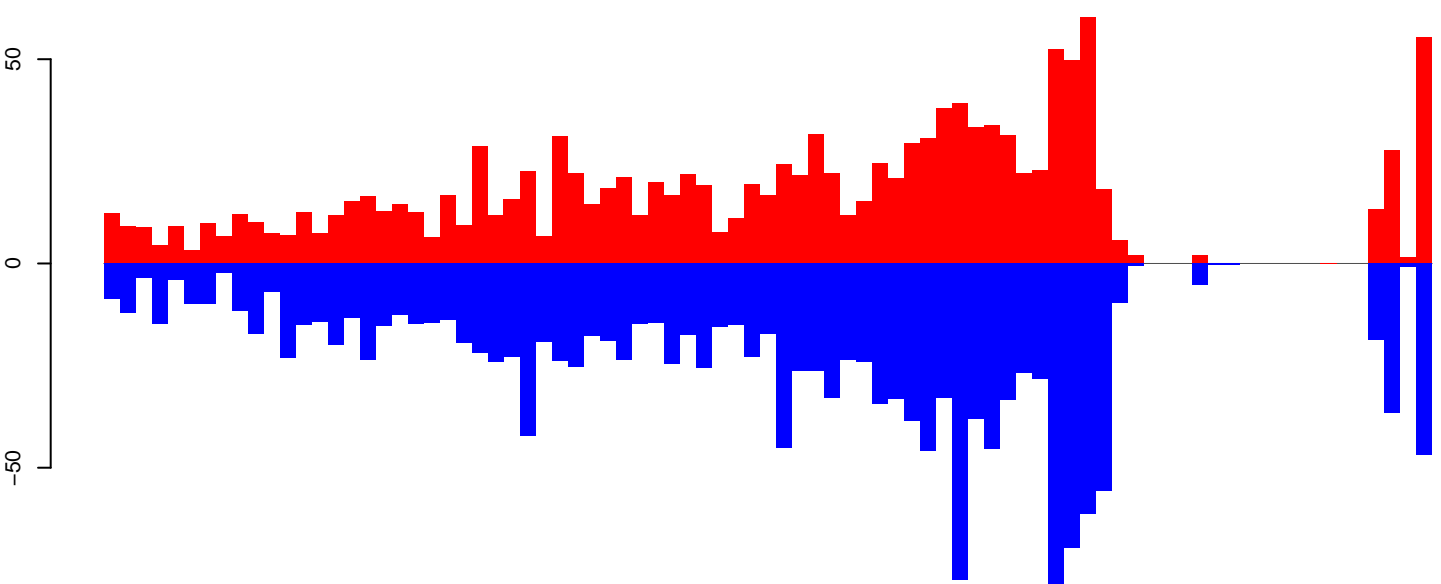
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



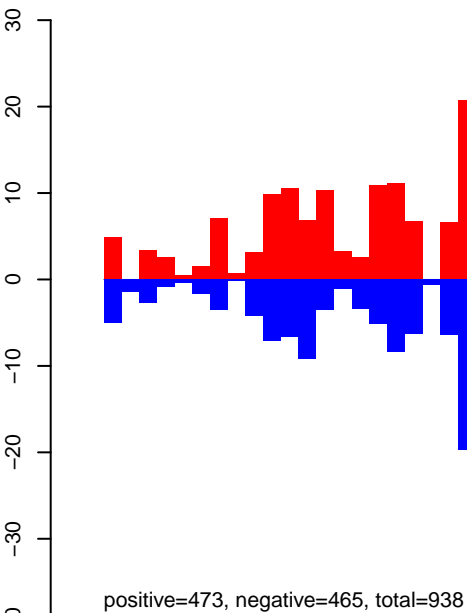
AeAeg_Aag2_DENV1.rep



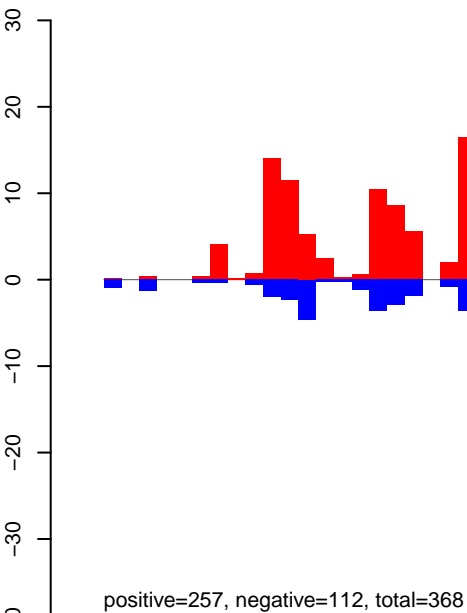
Window size=50, length=4176, Virus@X74945.1-Aedes albopictus_densovirus_2_DNA:1-4176

0 1000 2000 3000 4000

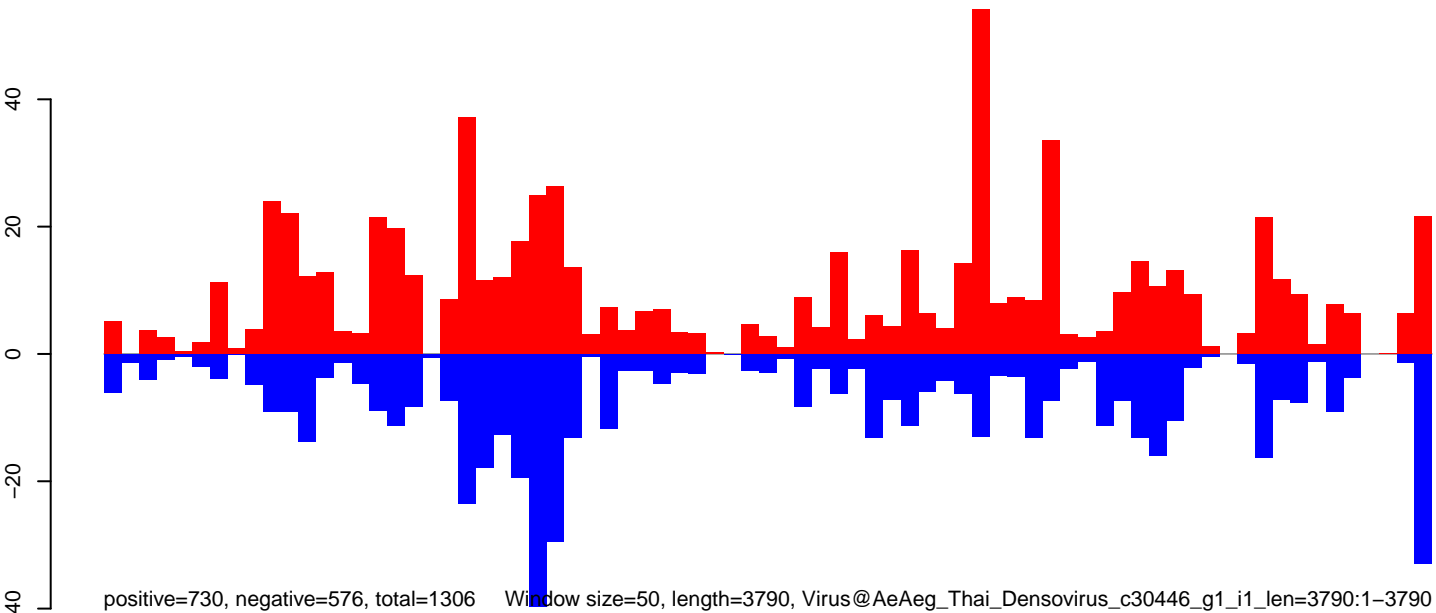
AeAeg_Aag2_DENV1.18_23.rep



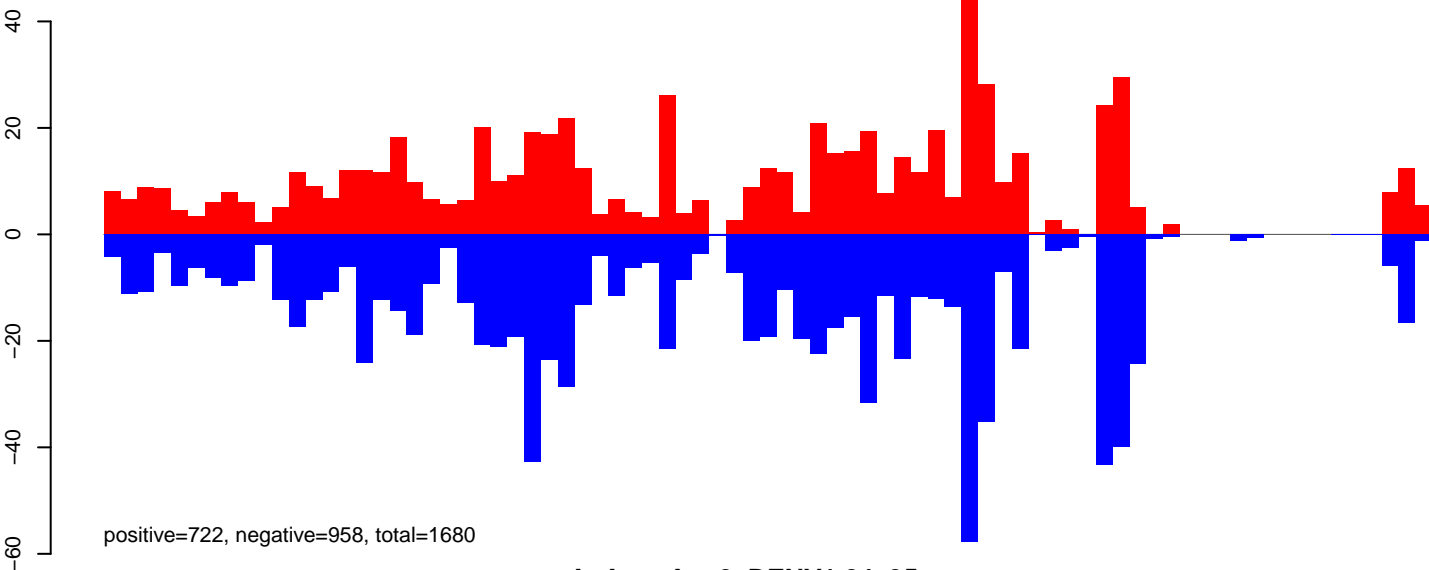
AeAeg_Aag2_DENV1.24_35.rep



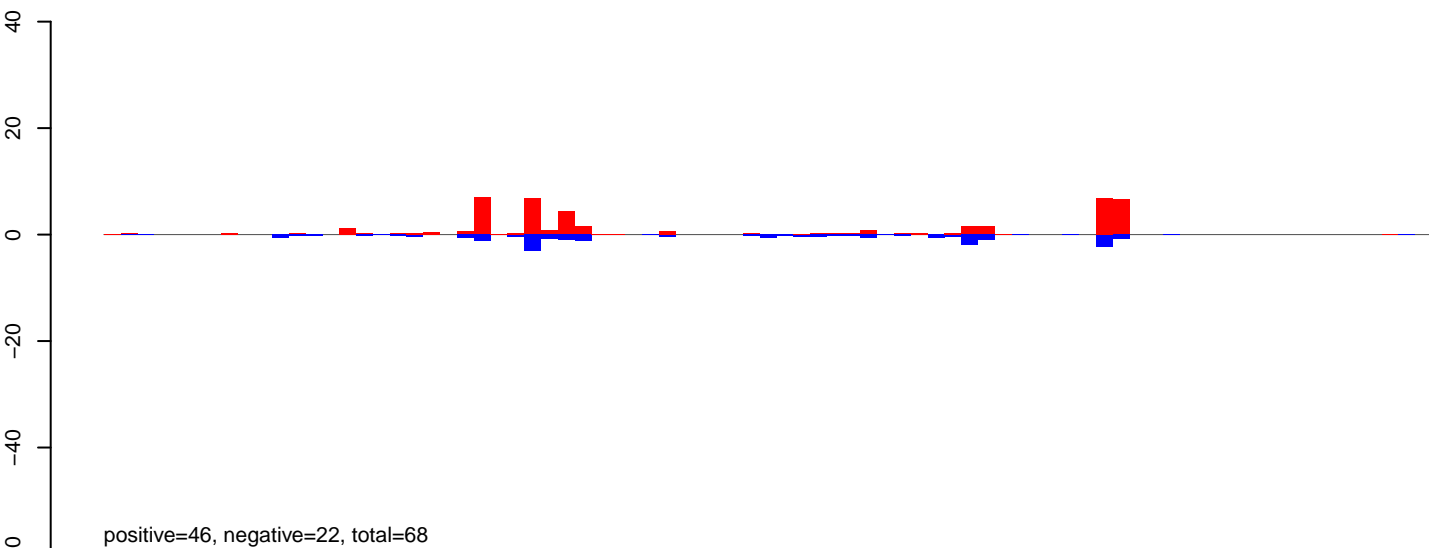
AeAeg_Aag2_DENV1.rep



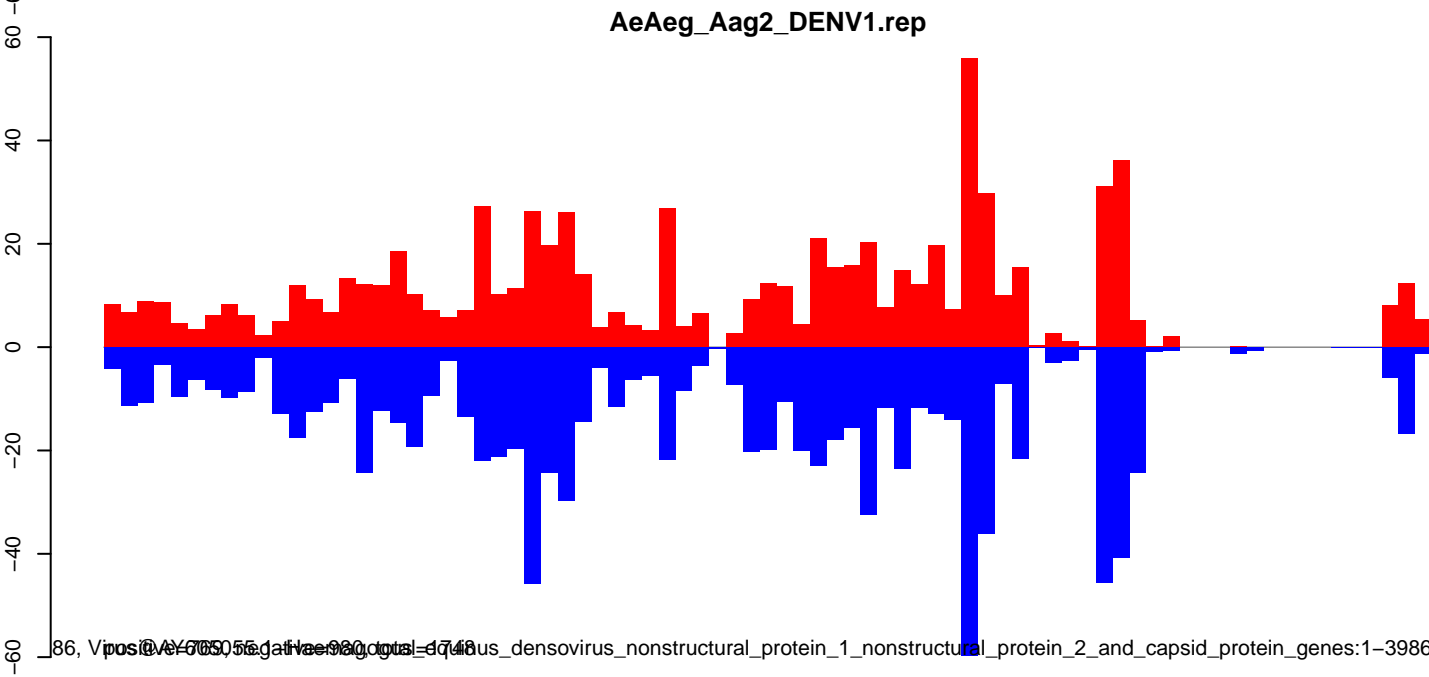
AeAeg_Aag2_DENV1.18_23.rep



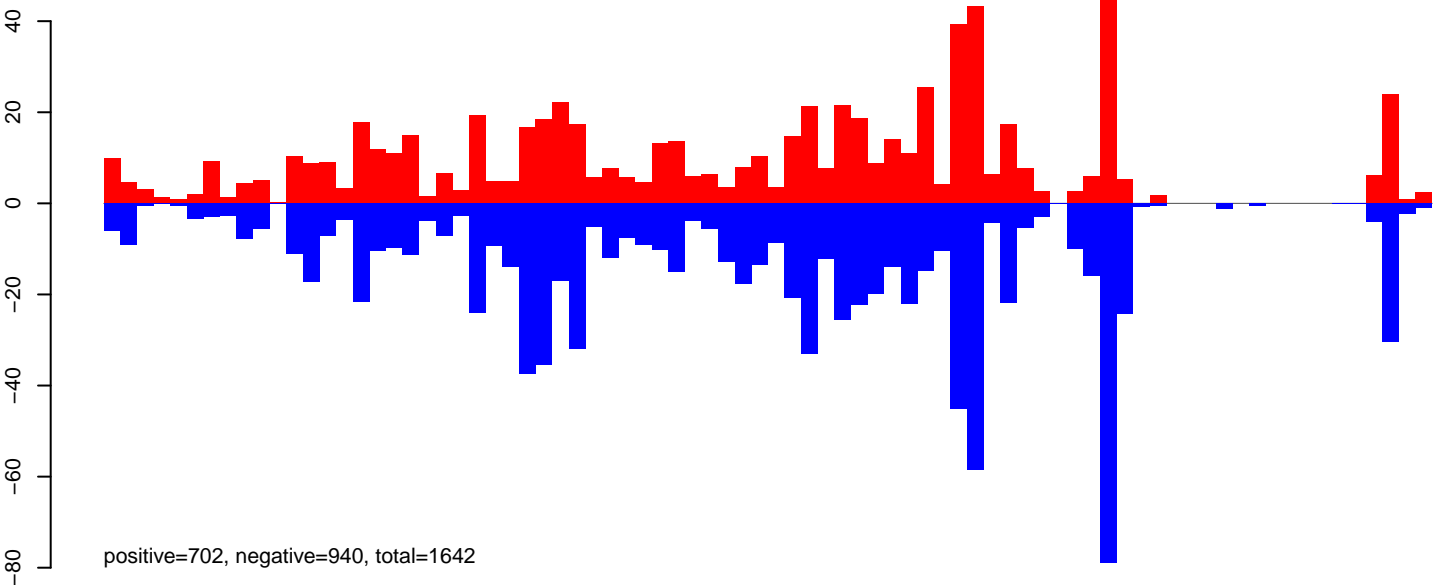
AeAeg_Aag2_DENV1.24_35.rep



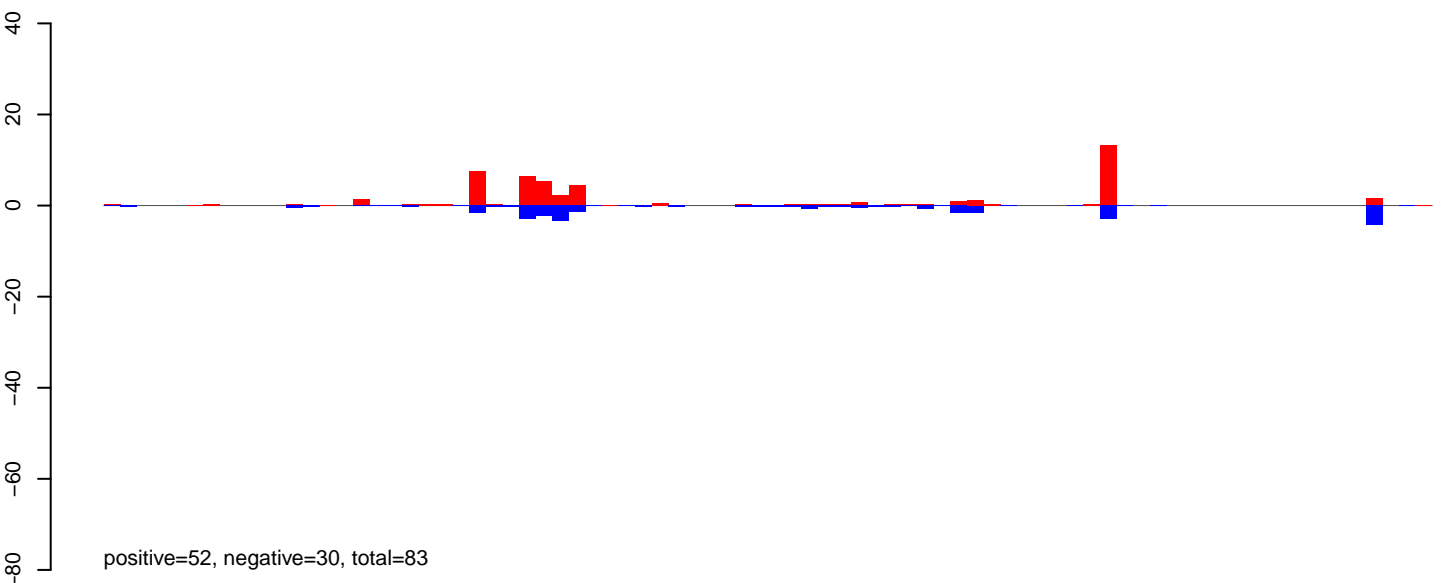
AeAeg_Aag2_DENV1.rep



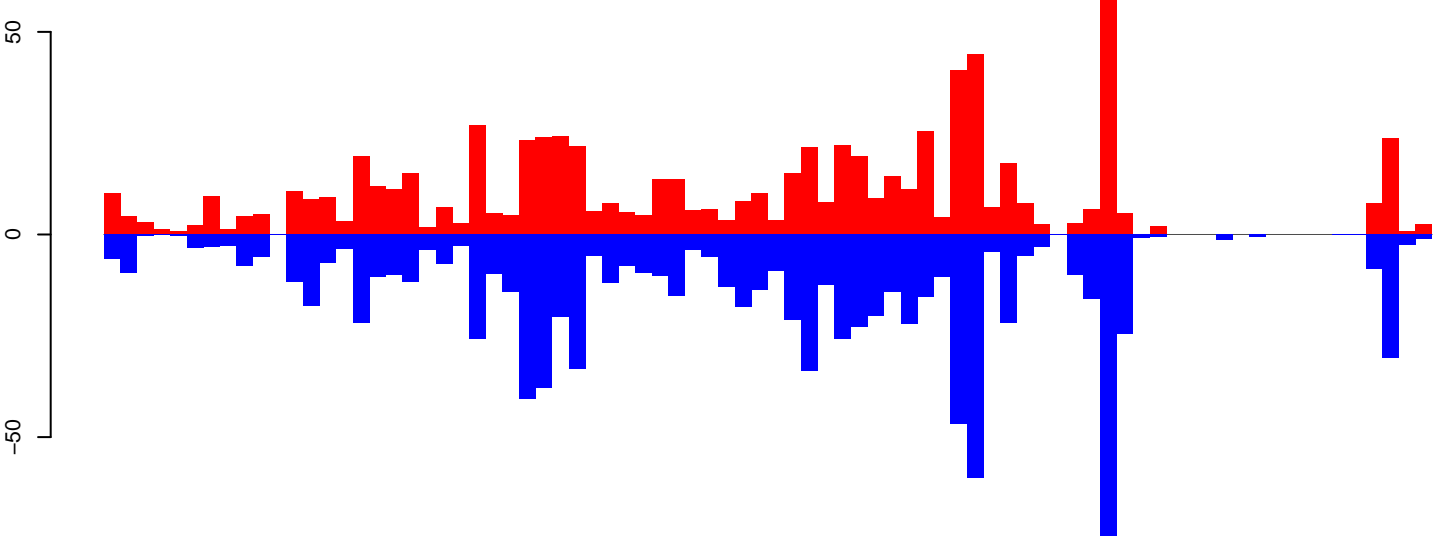
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



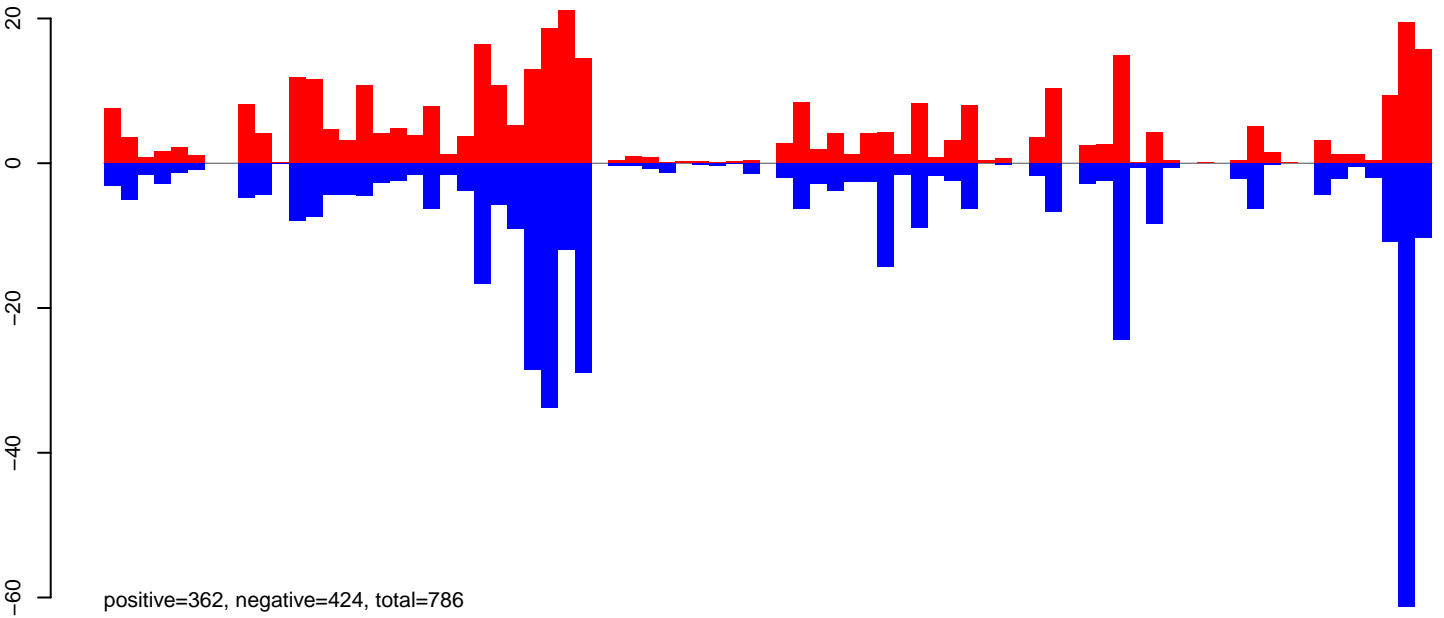
AeAeg_Aag2_DENV1.rep



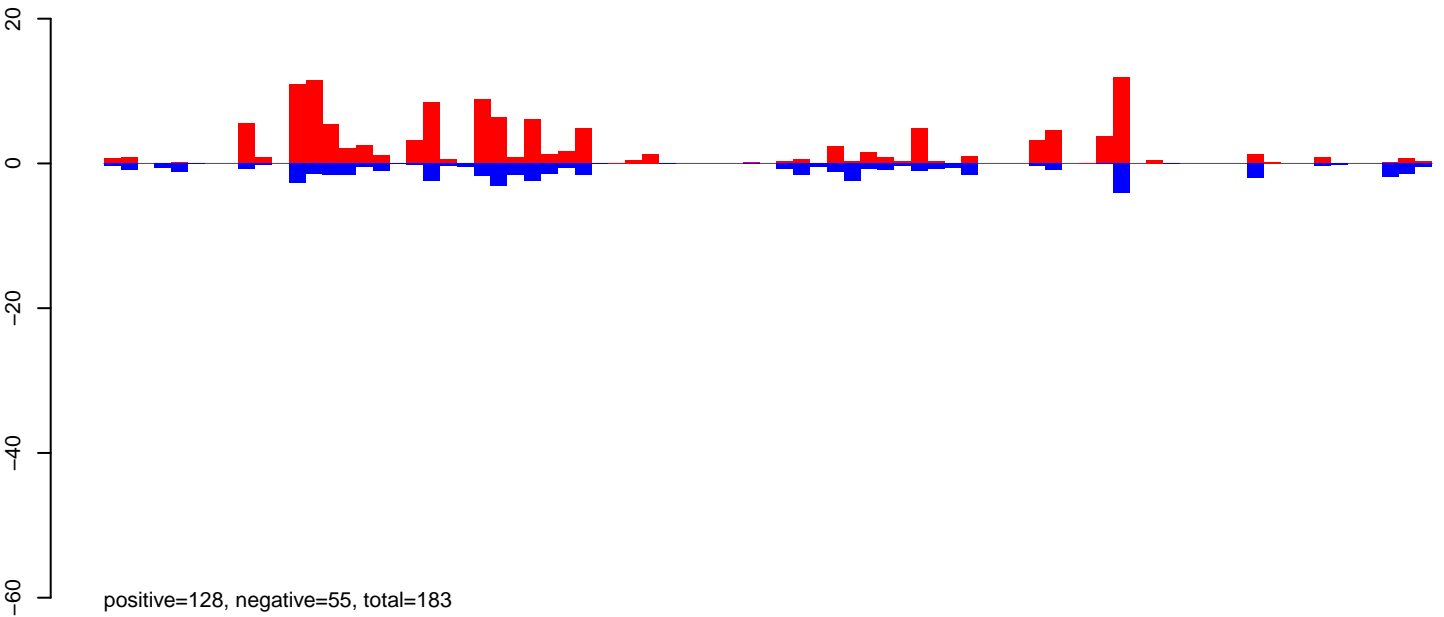
AY310871-754_region0751061724_densovirus_non-structural_protein_1_non-structural_protein_2_and_capsid_protein_genes:1-4006

0 1000 2000 3000 4000

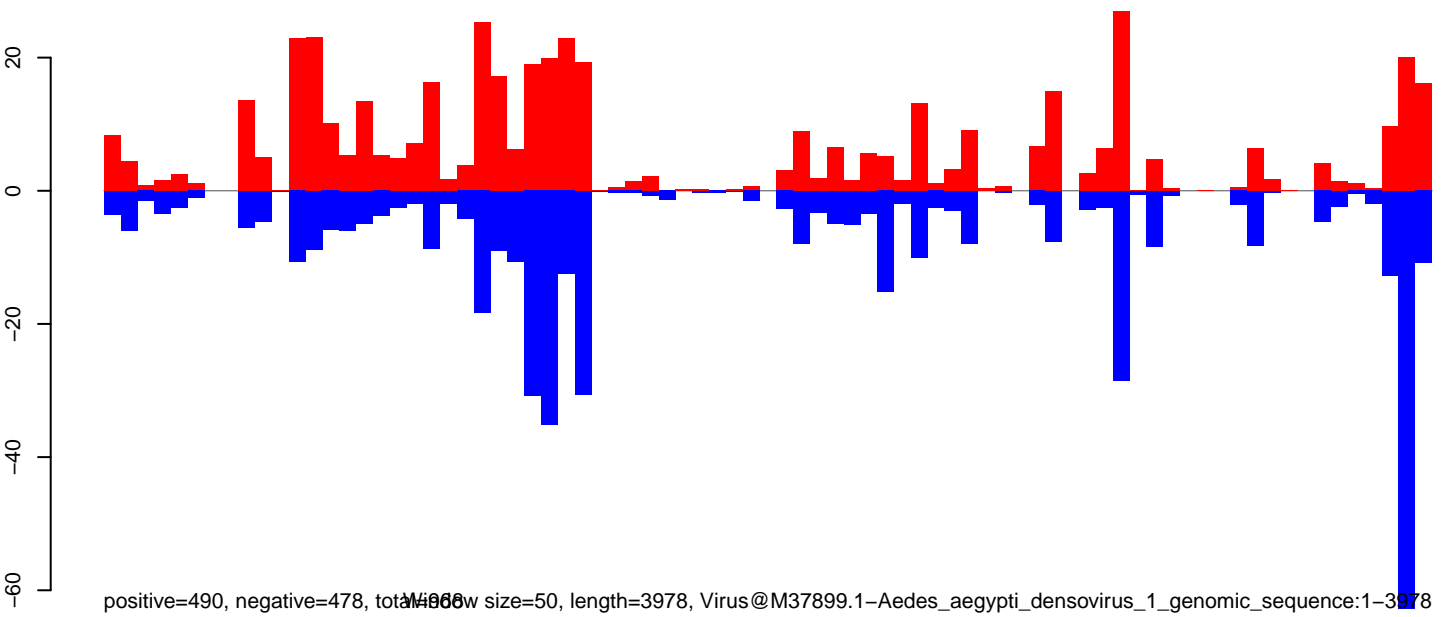
AeAeg_Aag2_DENV1.18_23.rep



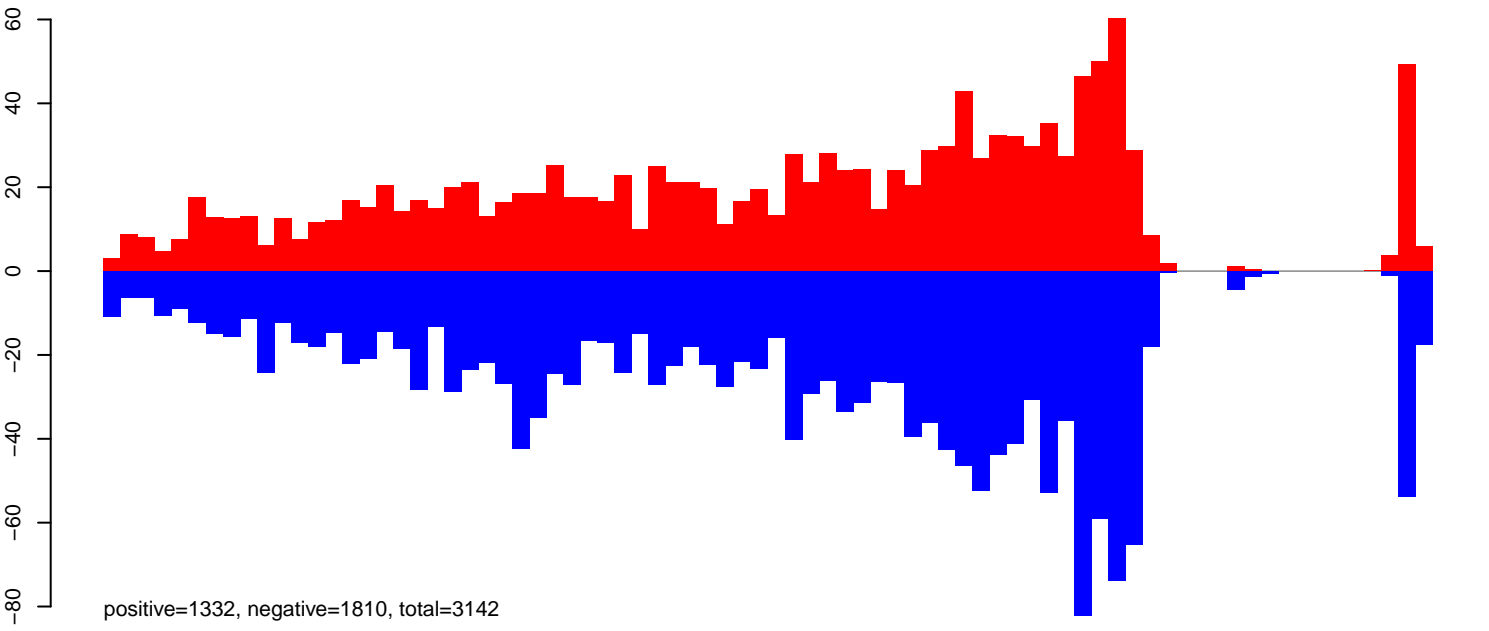
AeAeg_Aag2_DENV1.24_35.rep



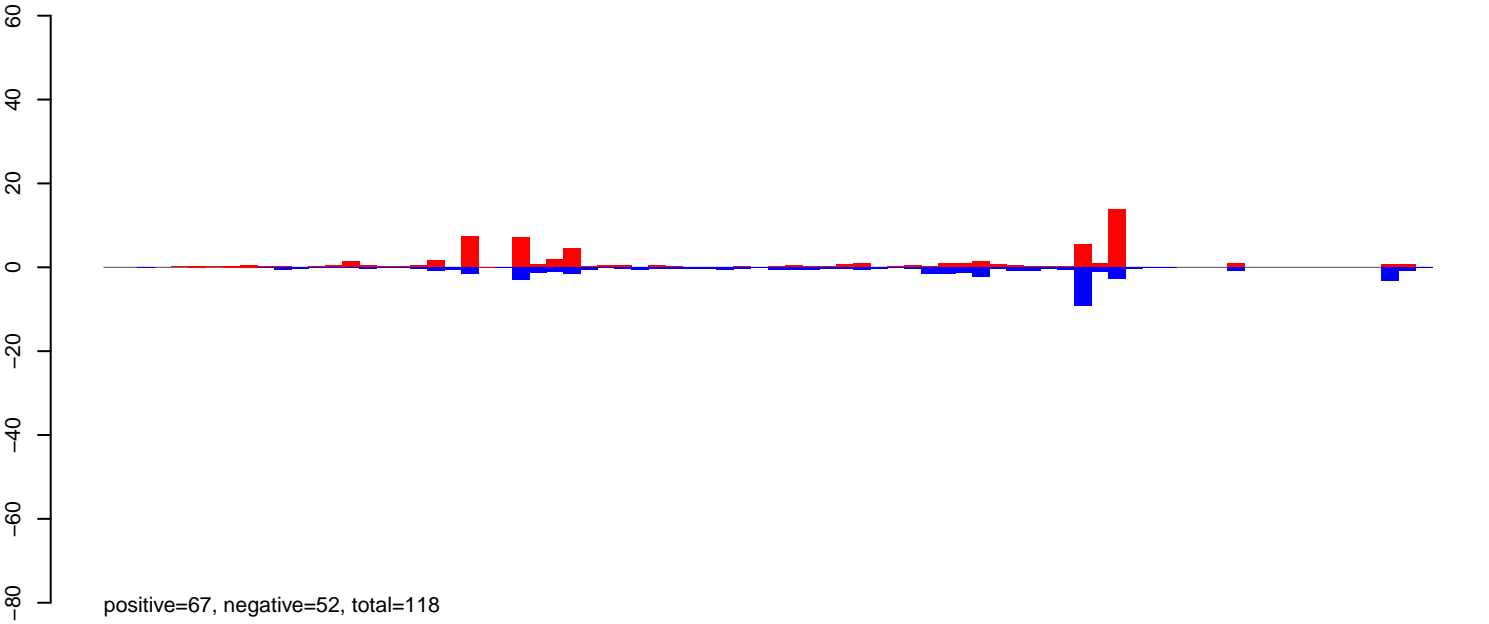
AeAeg_Aag2_DENV1.rep



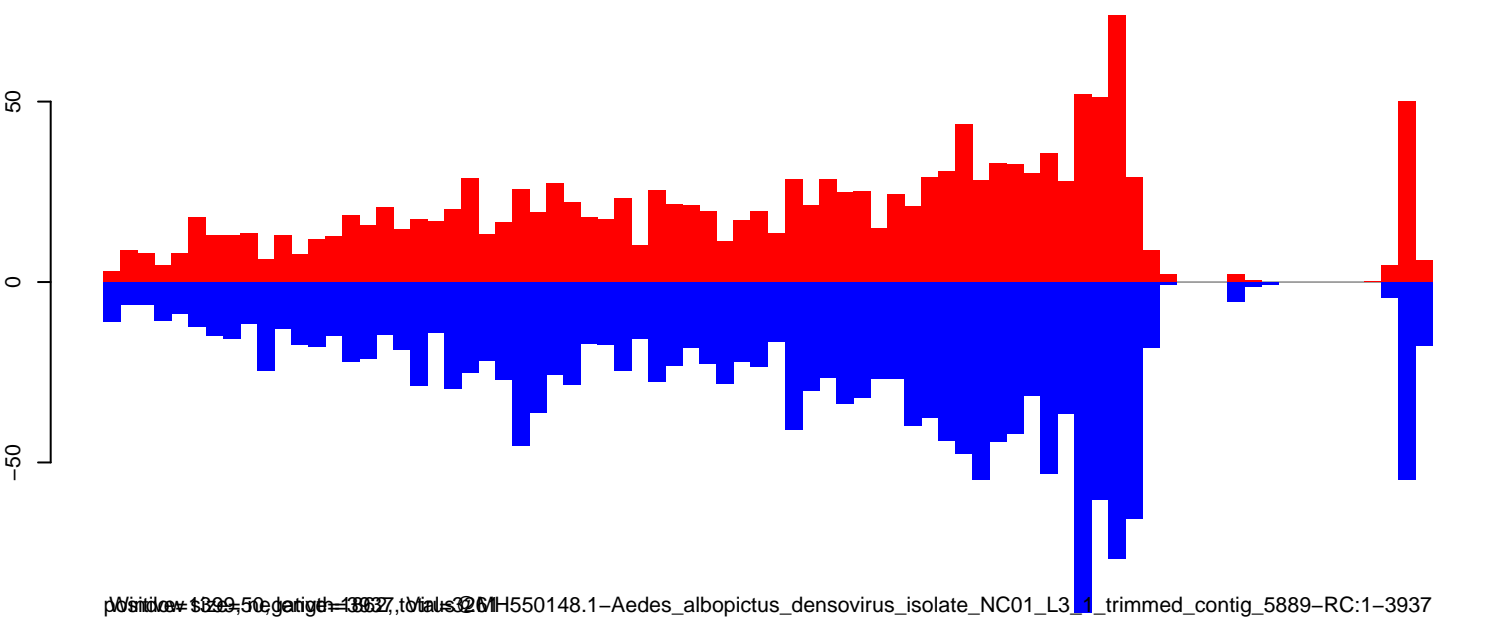
AeAeg_Aag2_DENV1.18_23.rep



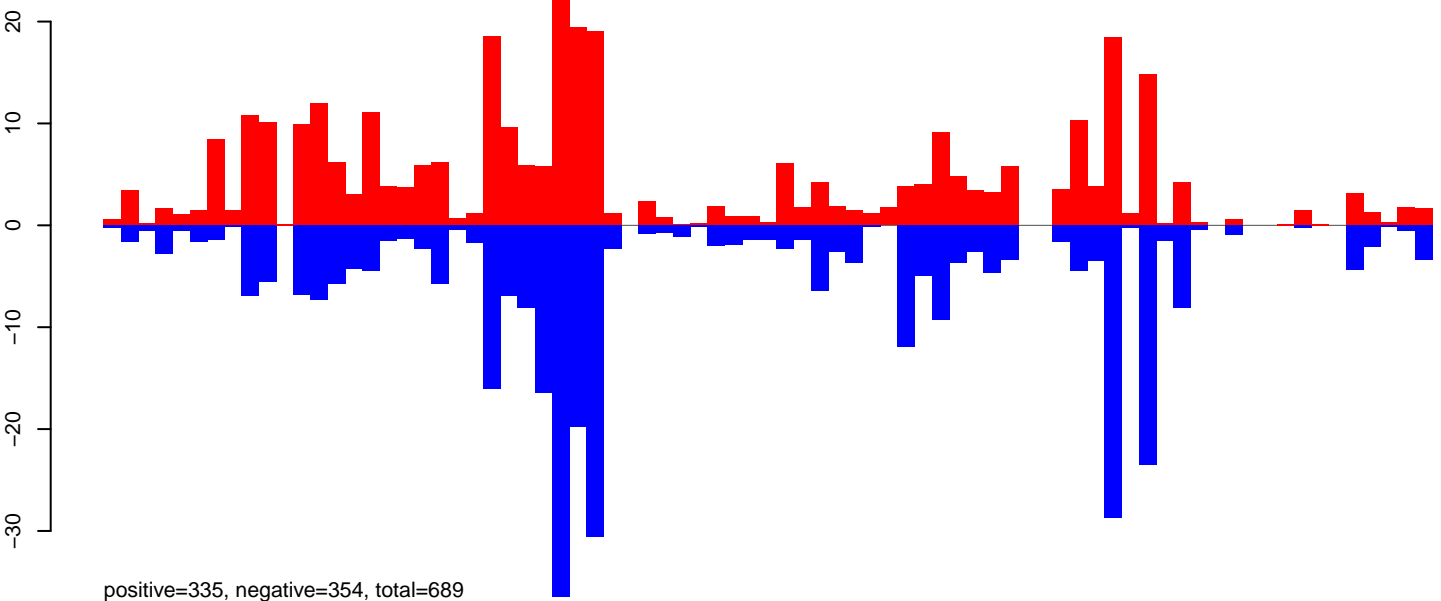
AeAeg_Aag2_DENV1.24_35.rep



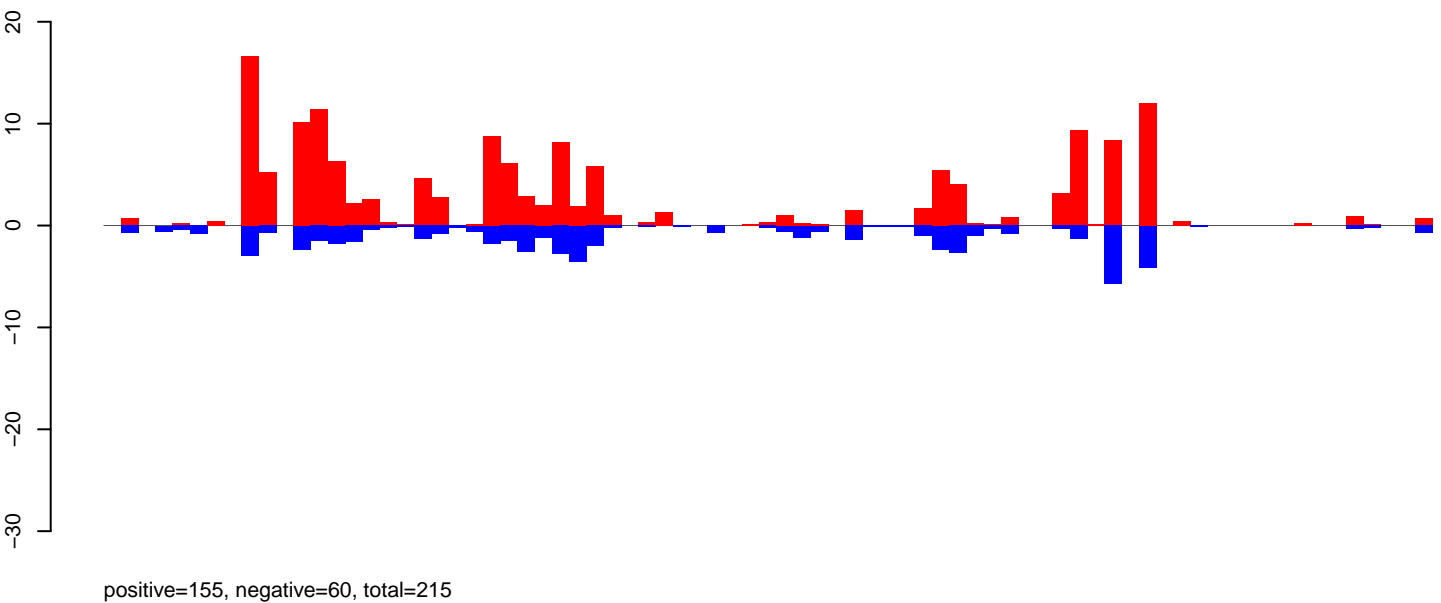
AeAeg_Aag2_DENV1.rep



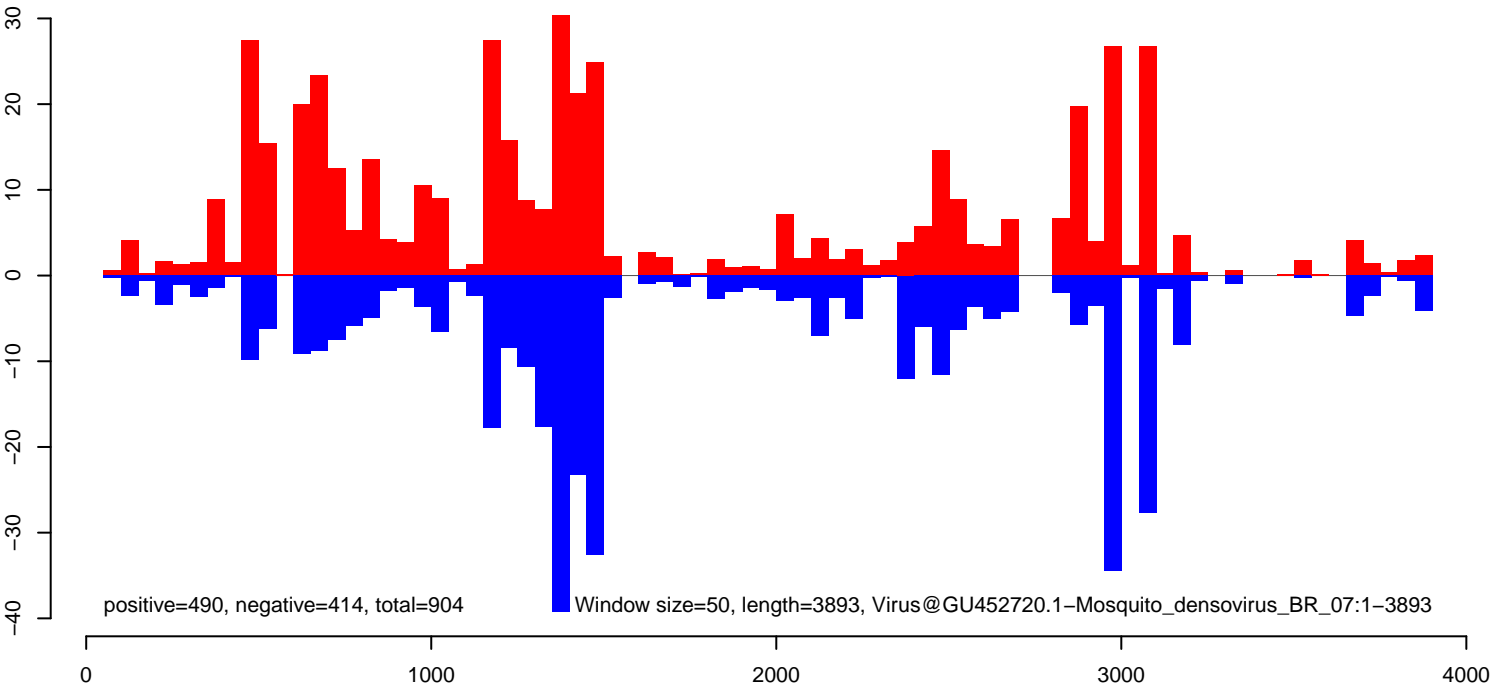
AeAeg_Aag2_DENV1.18_23.rep



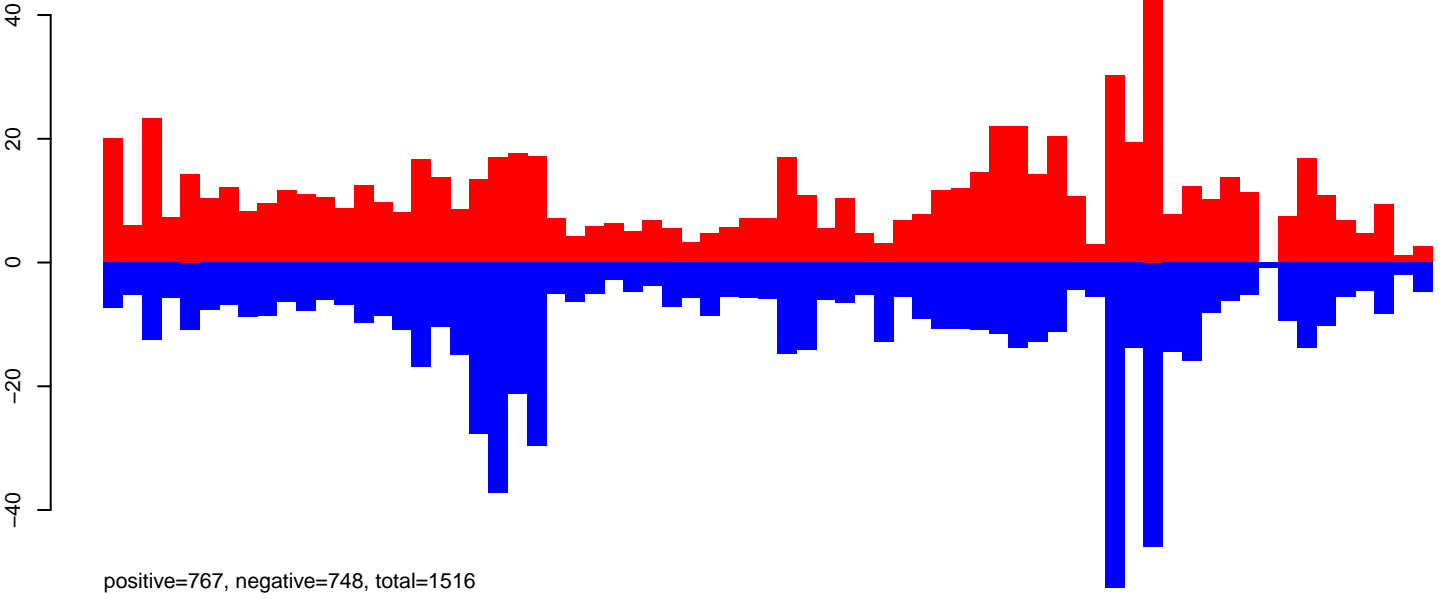
AeAeg_Aag2_DENV1.24_35.rep



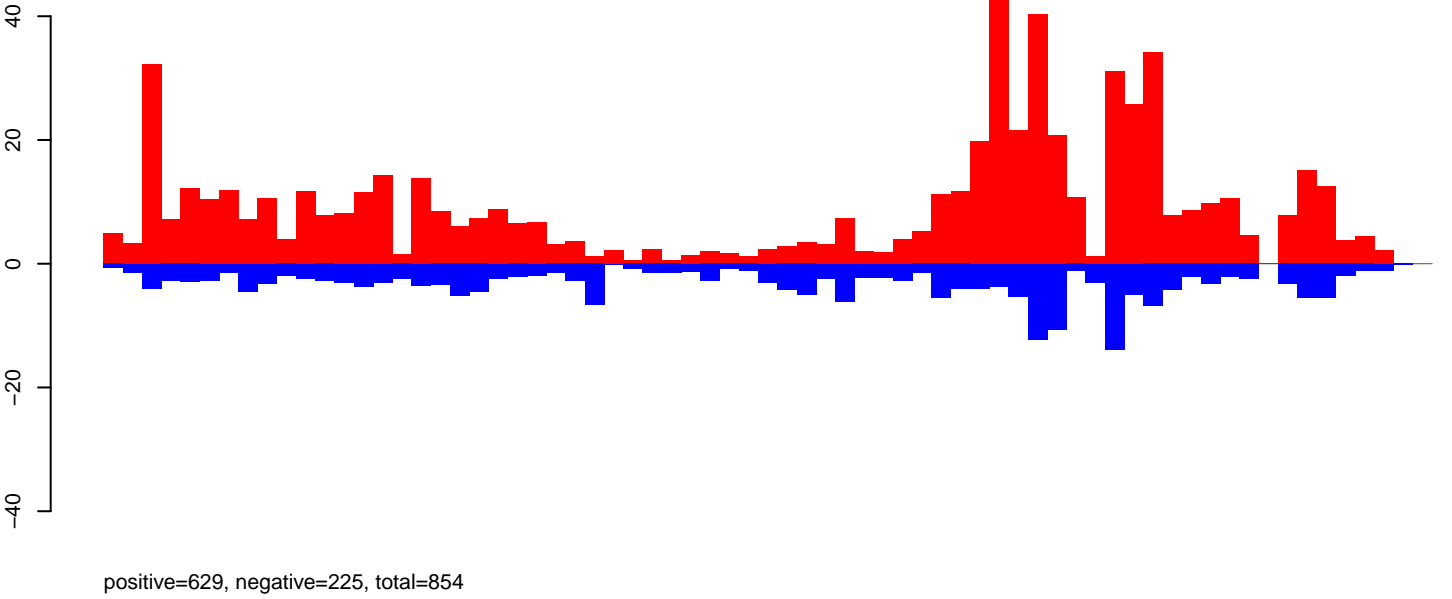
AeAeg_Aag2_DENV1.rep



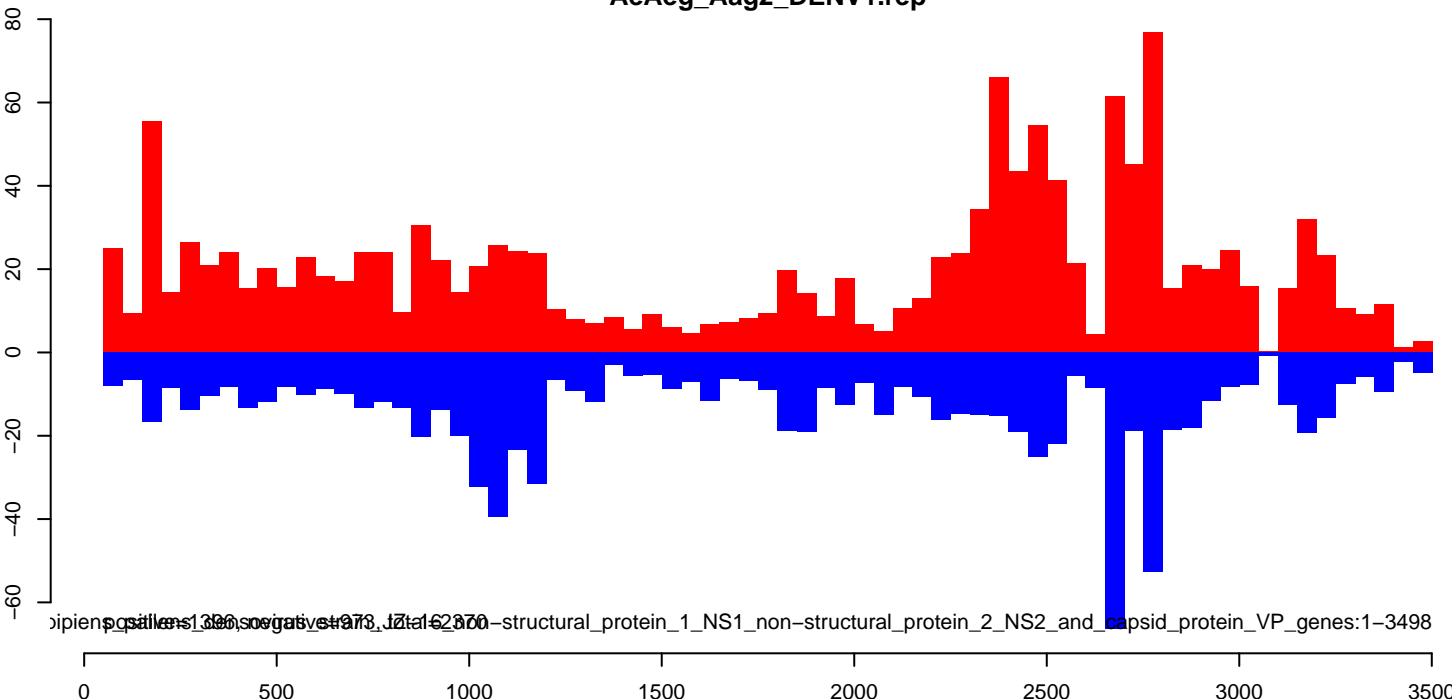
AeAeg_Aag2_DENV1.18_23.rep



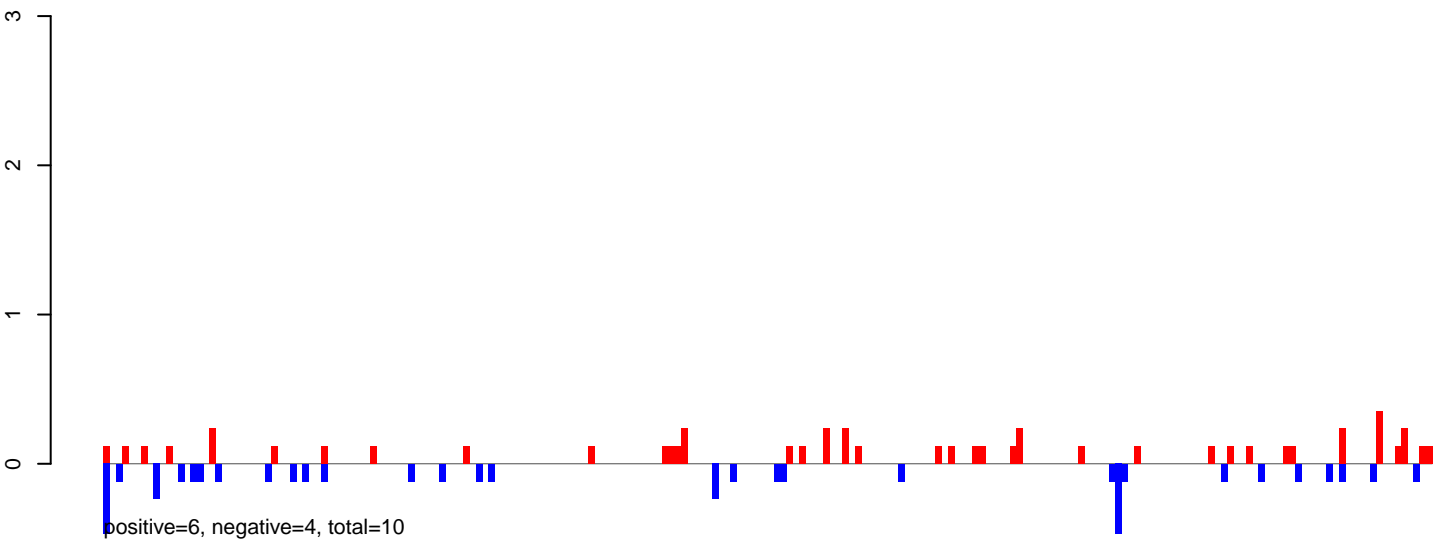
AeAeg_Aag2_DENV1.24_35.rep



AeAeg_Aag2_DENV1.rep



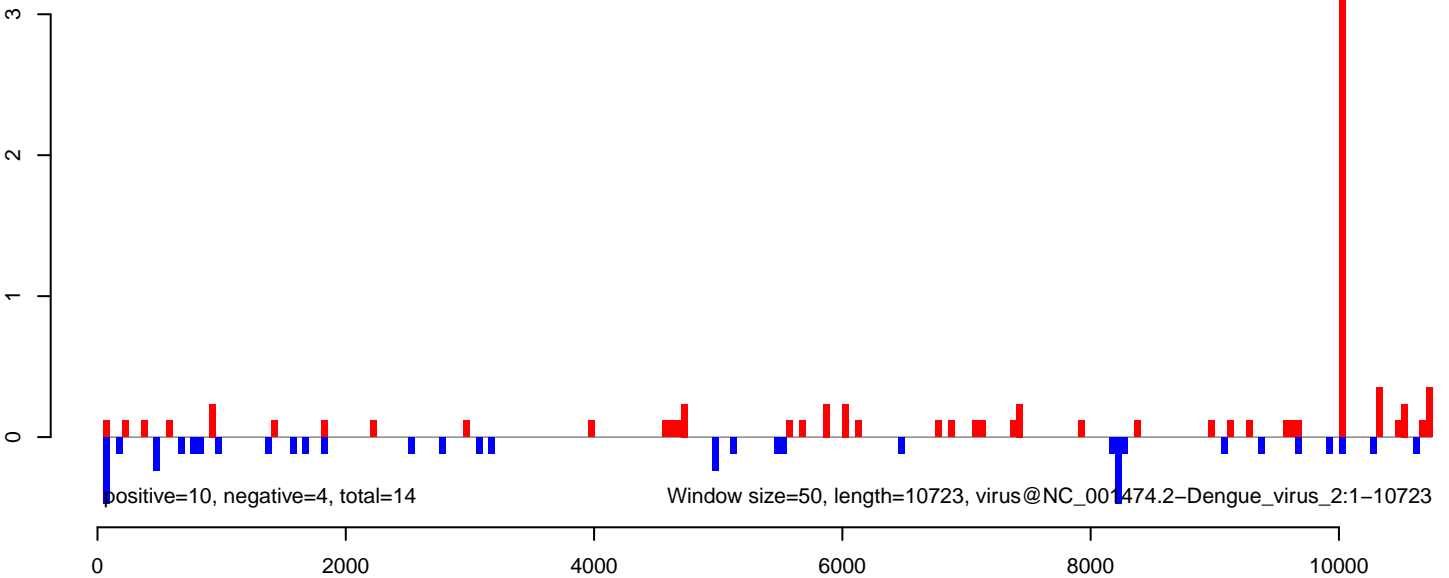
AeAeg_Aag2_DENV1.18_23.rep



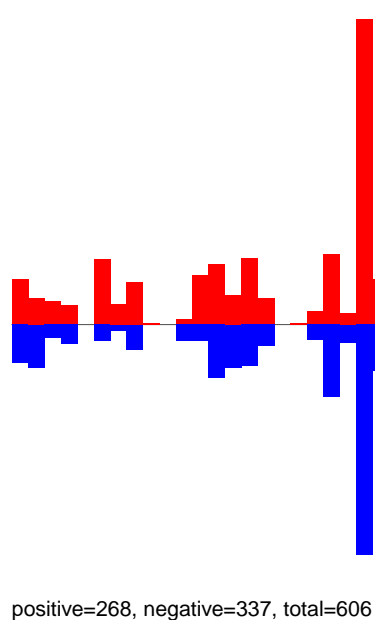
AeAeg_Aag2_DENV1.24_35.rep



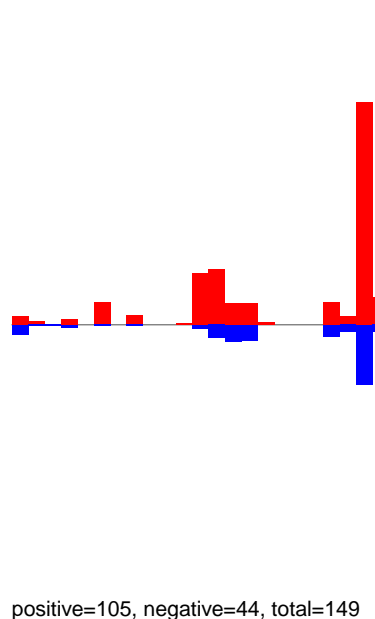
AeAeg_Aag2_DENV1.rep



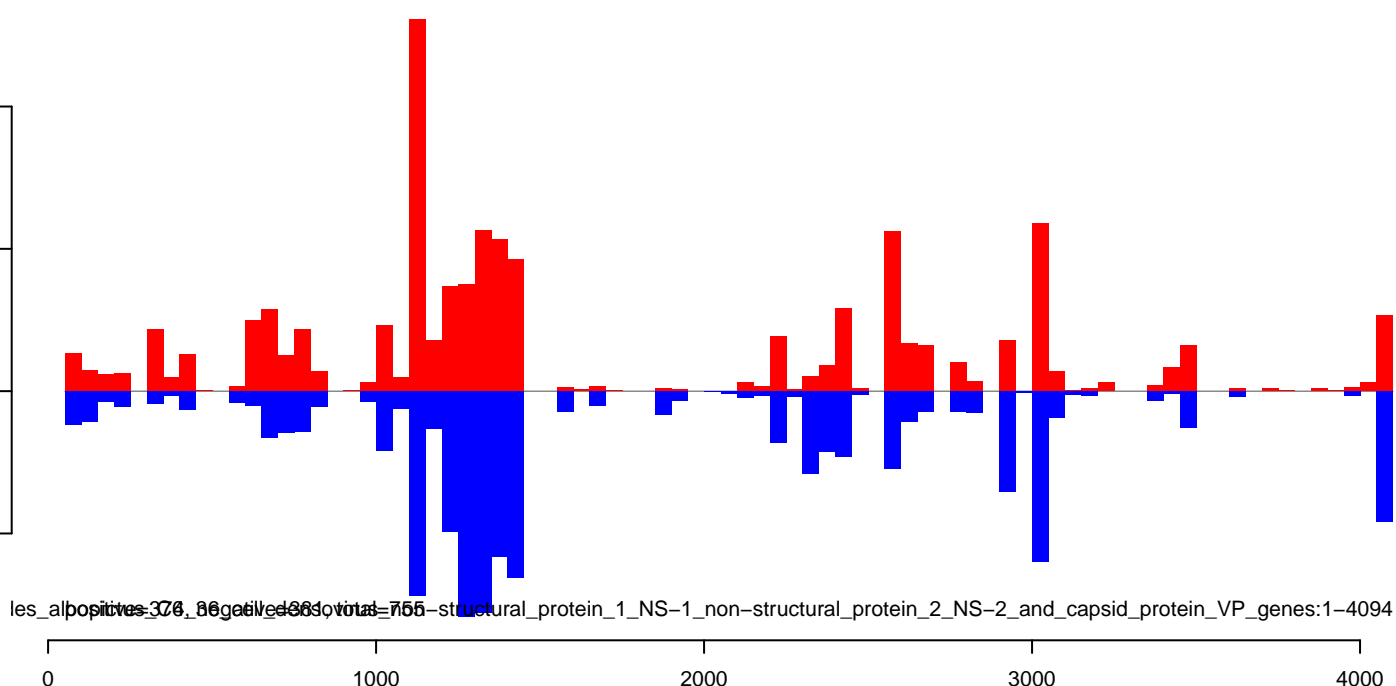
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



AeAeg_Aag2_DENV1.rep

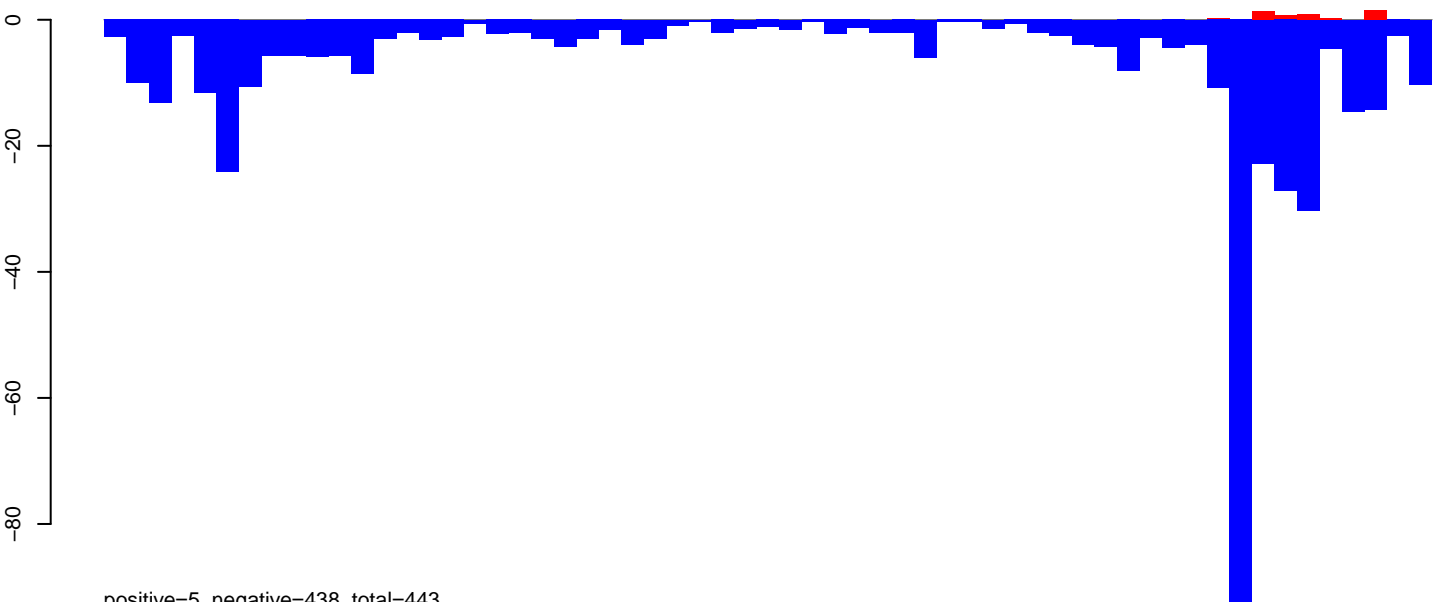


AeAeg_Aag2_DENV1.18_23.rep



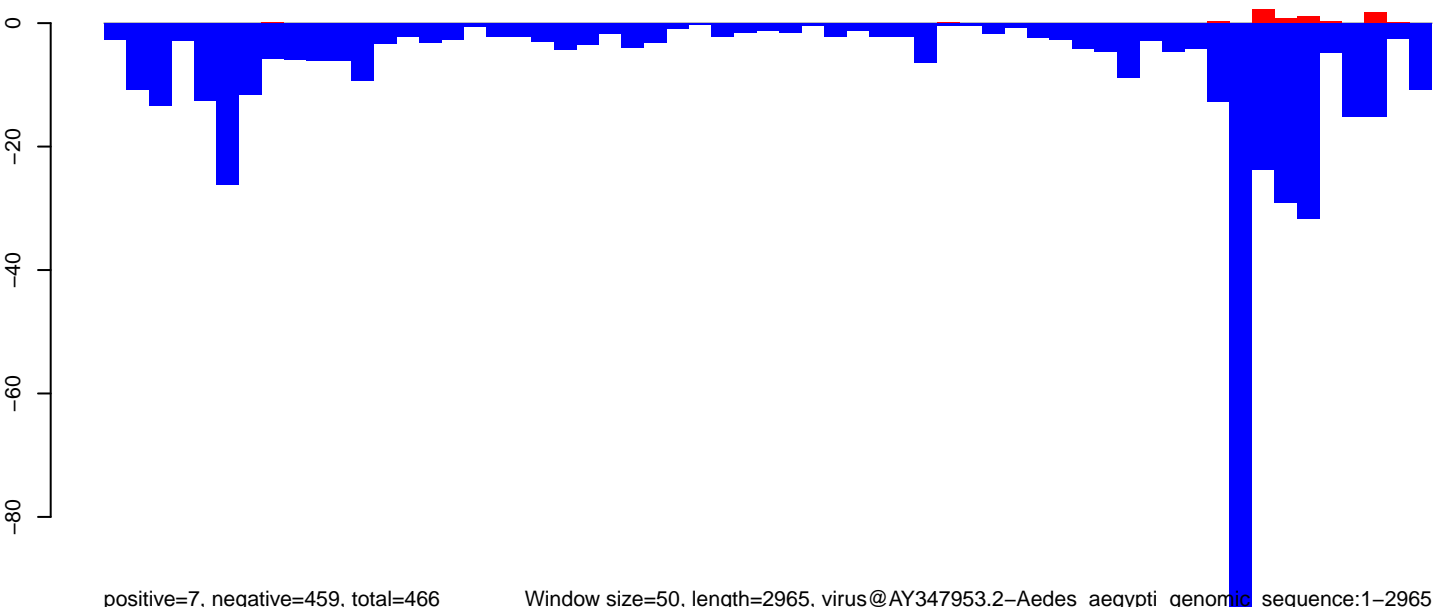
positive=2, negative=21, total=23

AeAeg_Aag2_DENV1.24_35.rep



positive=5, negative=438, total=443

AeAeg_Aag2_DENV1.rep

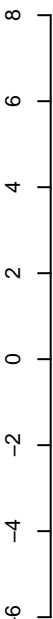


positive=7, negative=459, total=466

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

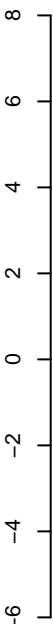
0 500 1000 1500 2000 2500 3000

AeAeg_Aag2_DENV1.18_23.rep



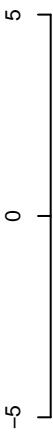
positive=17, negative=17, total=34

AeAeg_Aag2_DENV1.24_35.rep



positive=17, negative=3, total=20

AeAeg_Aag2_DENV1.rep

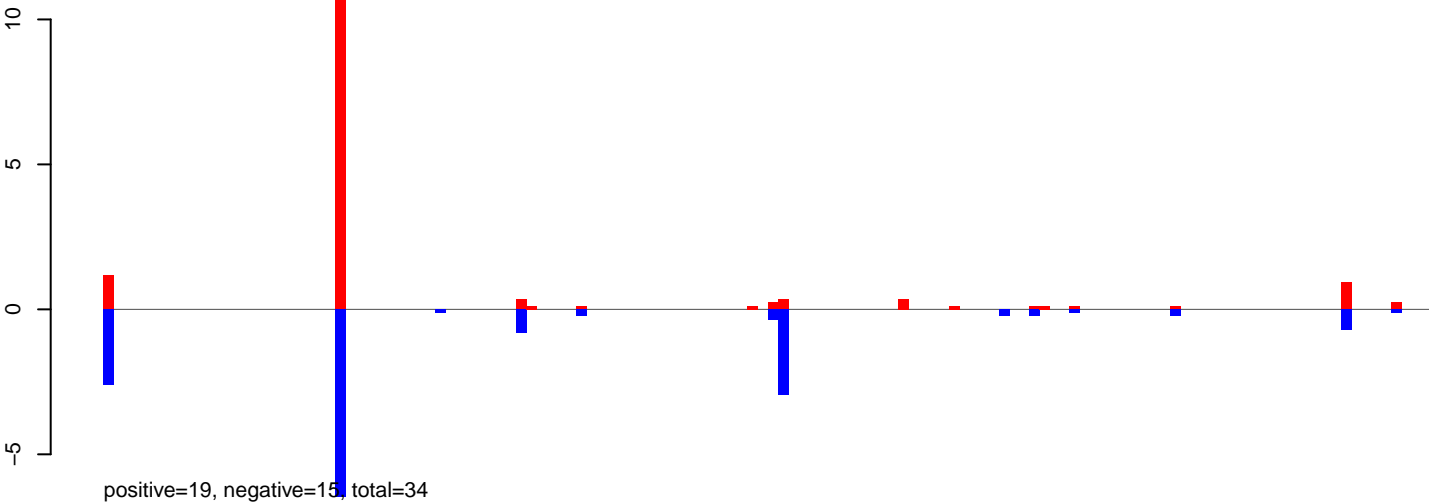


positive=33, negative=21, total=54

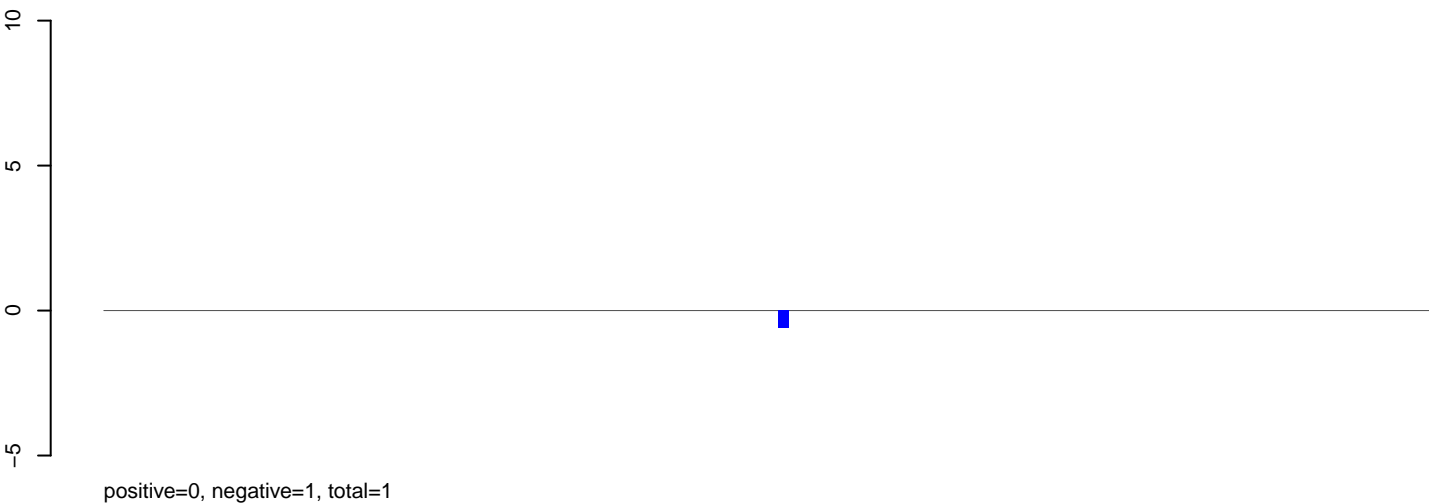
Window size=50, length=11375, virus@NC_005064.1-Kamiti_River_virus:1-11375



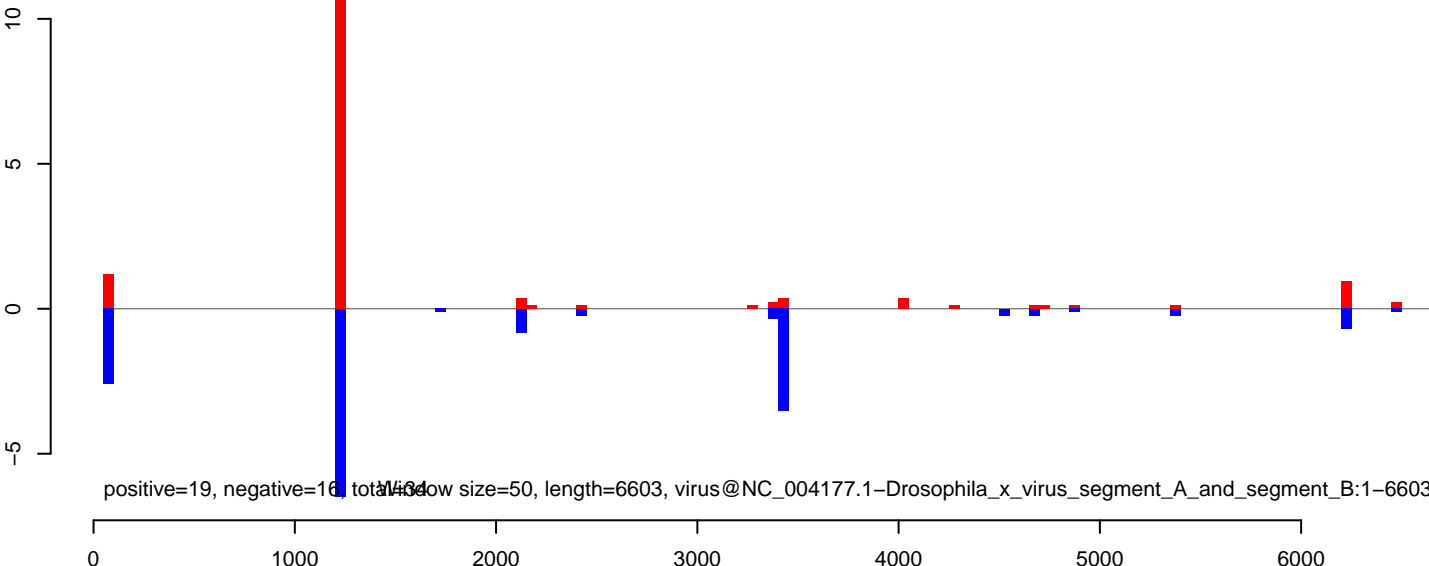
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



AeAeg_Aag2_DENV1.rep



AeAeg_Aag2_DENV1.18_23.rep

20
0
-20

positive=91, negative=67, total=158

AeAeg_Aag2_DENV1.24_35.rep

20
0
-20

positive=6, negative=1, total=7

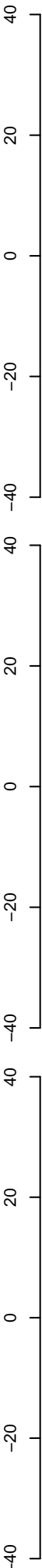
AeAeg_Aag2_DENV1.rep

20
0
-20

positive=97, negative=68, total=165, length=11079, virus@KC181923.1-Aedes_flavivirus_strain_AEFV-SPFLD-MO-2011-MP6:1-11079

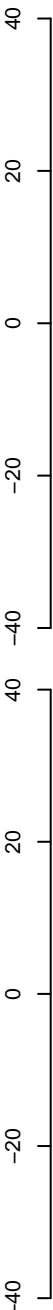
0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV1.18_23.rep



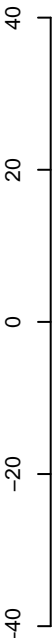
positive=90, negative=66, total=156

AeAeg_Aag2_DENV1.24_35.rep



positive=6, negative=1, total=7

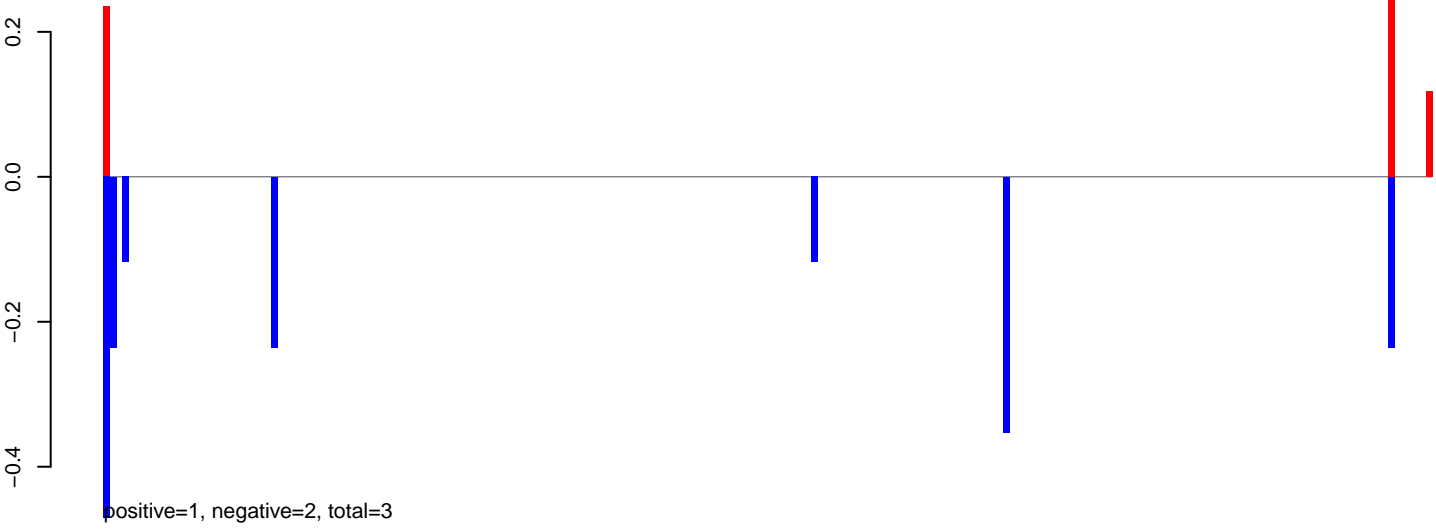
AeAeg_Aag2_DENV1.rep



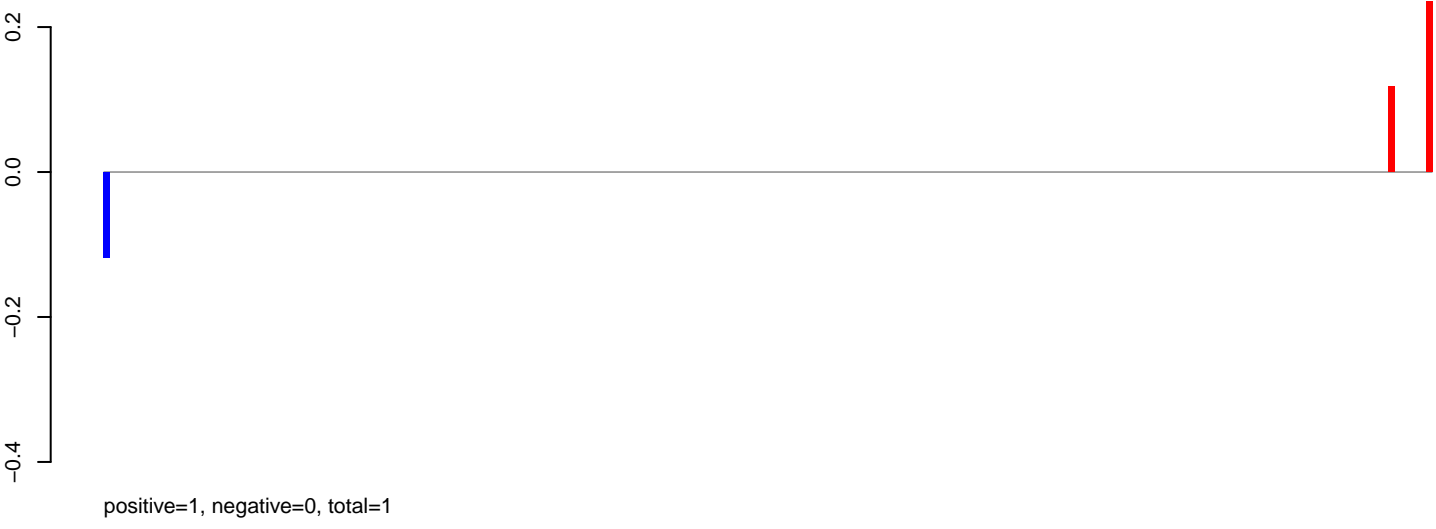
positive=96, negative=67, total=163, window size=50, length=11064, virus@NC_012932.1-Aedes_flavivirus_genomic_RNA_strain_Narita-21:1-11064



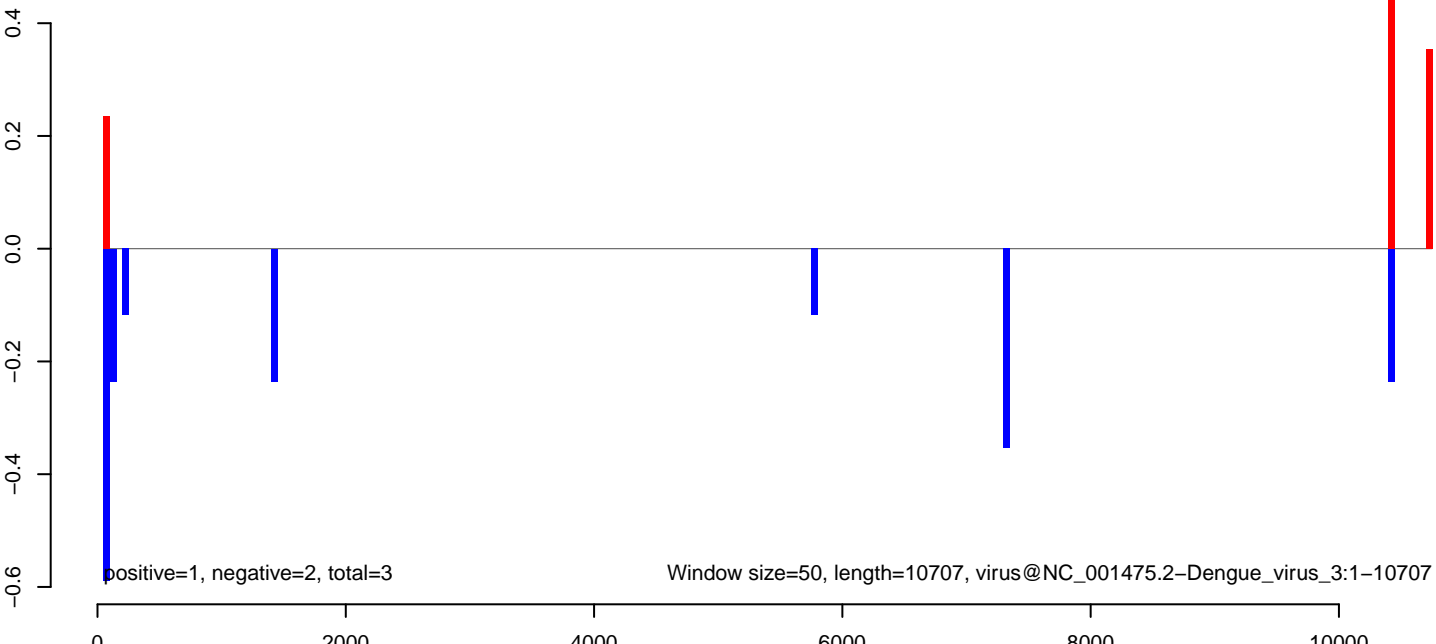
AeAeg_Aag2_DENV1.18_23.rep



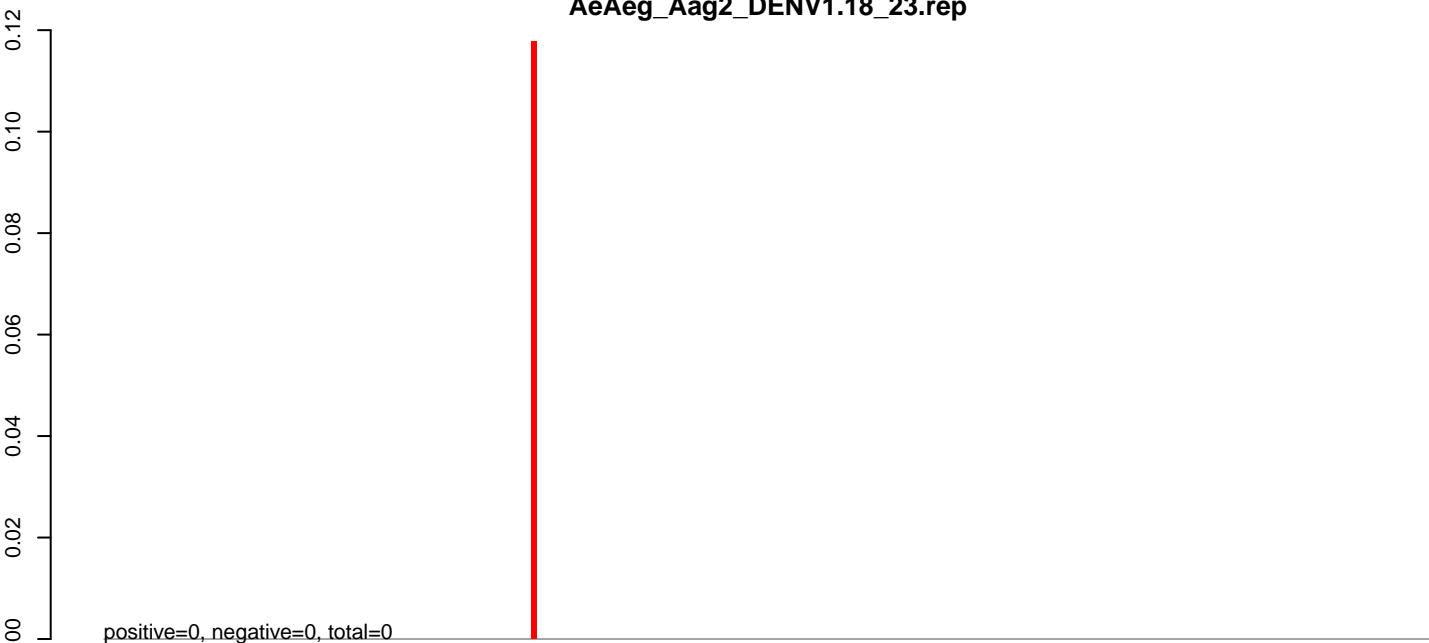
AeAeg_Aag2_DENV1.24_35.rep



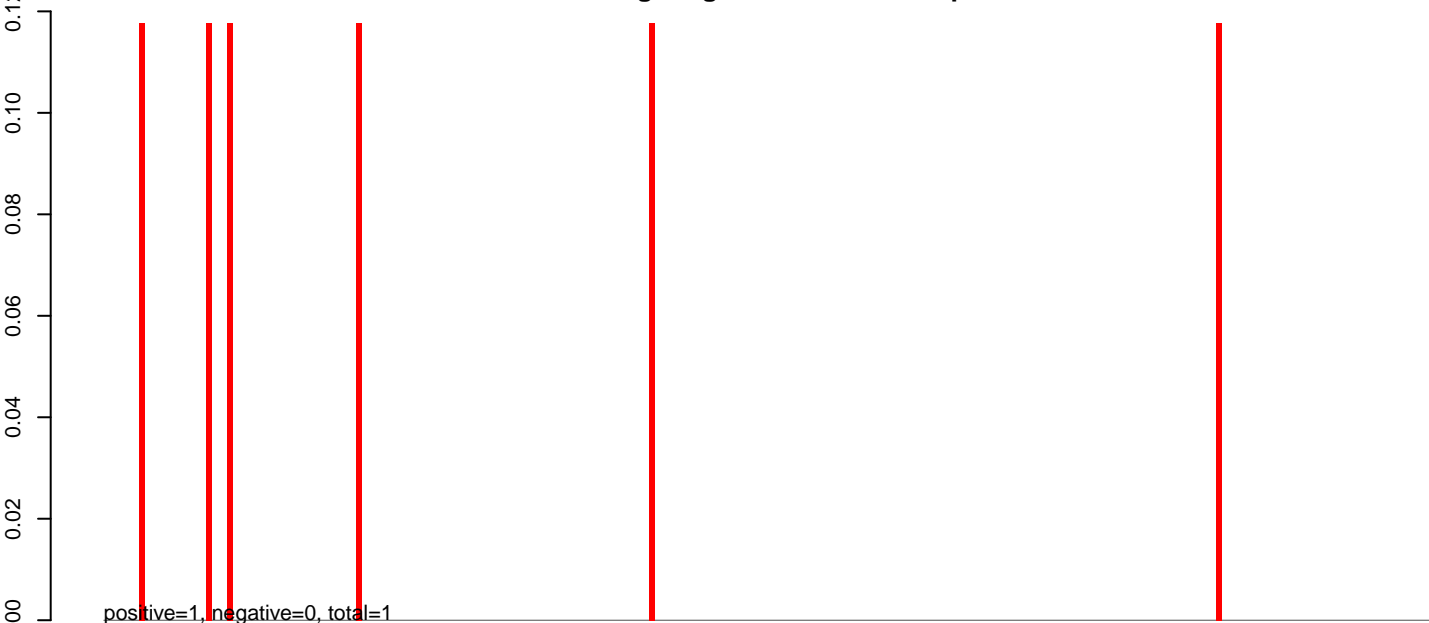
AeAeg_Aag2_DENV1.rep



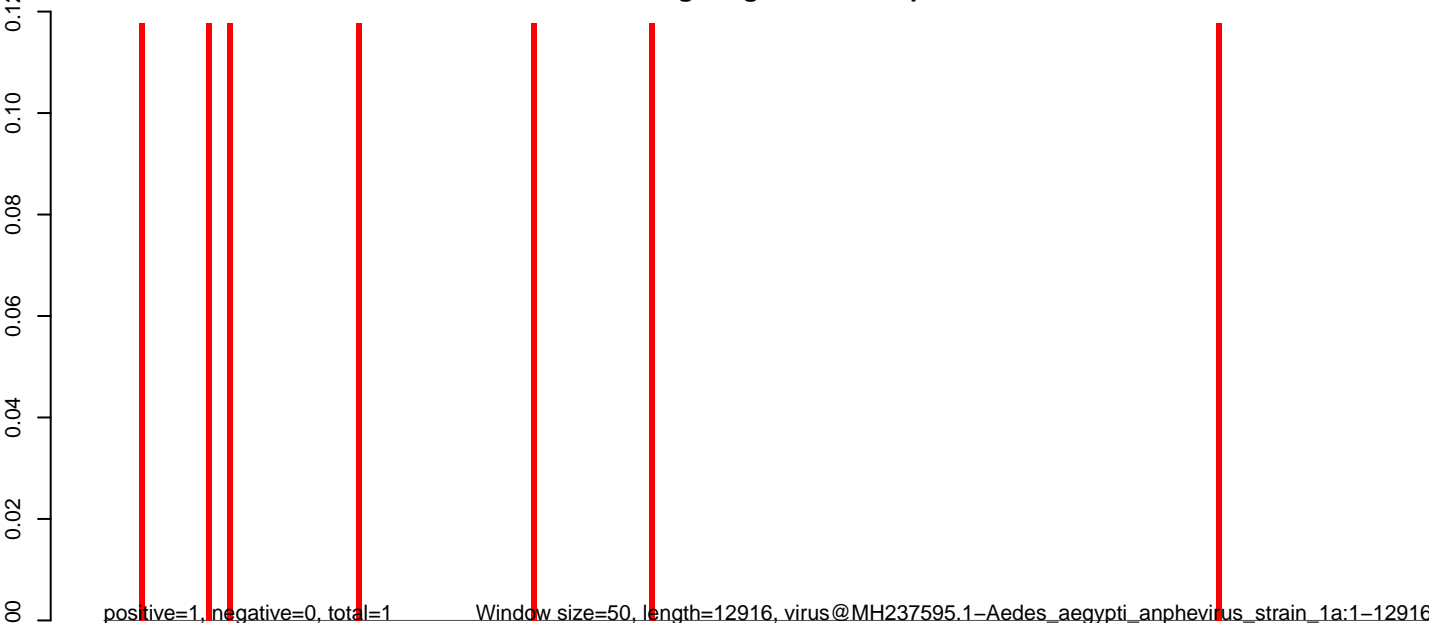
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

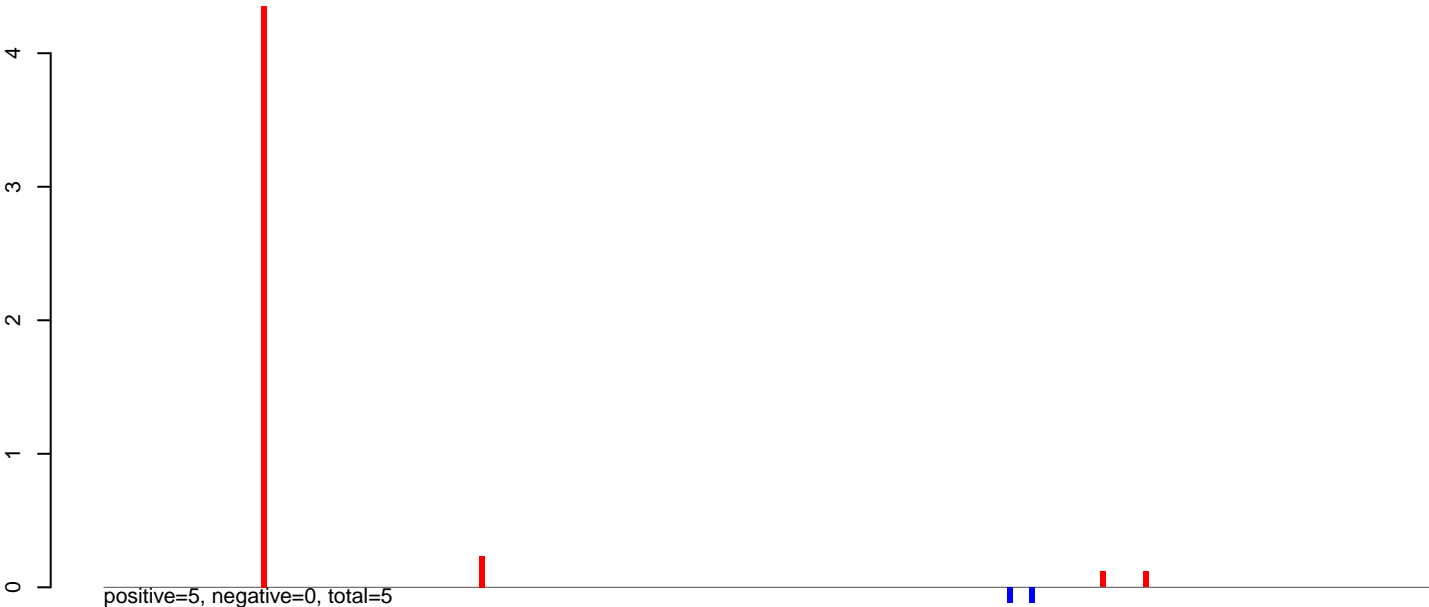


AeAeg_Aag2_DENV1.rep

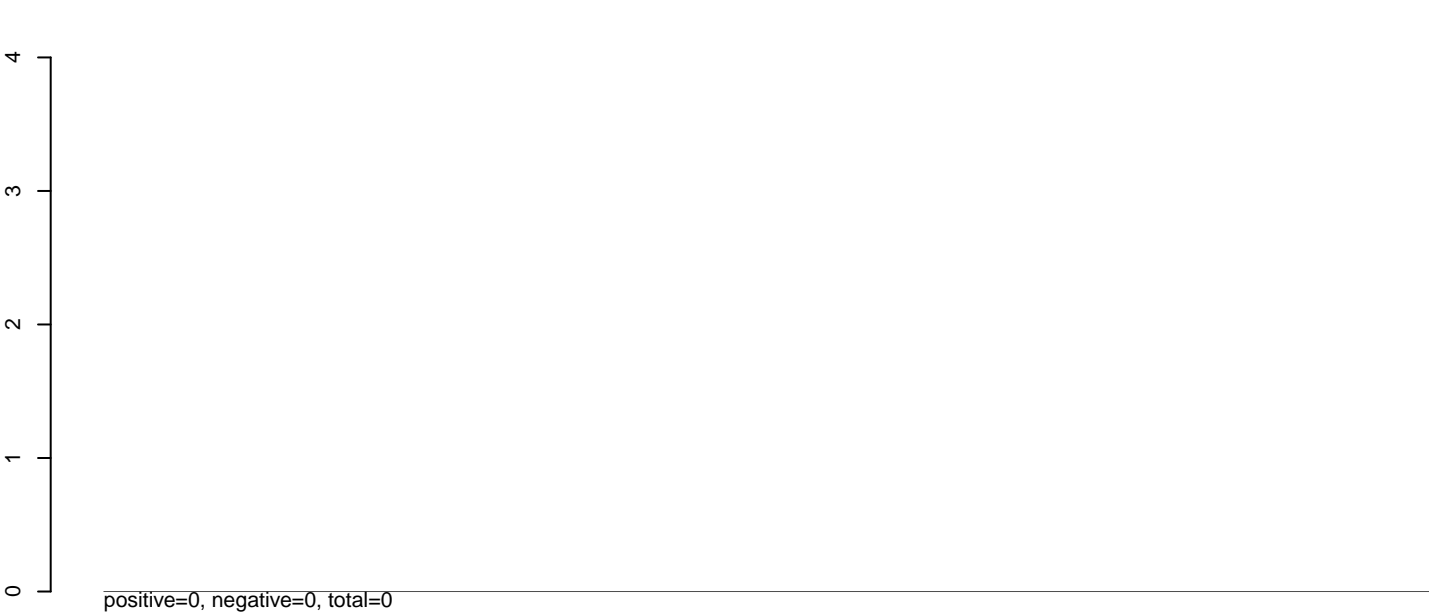


0 2000 4000 6000 8000 10000 12000

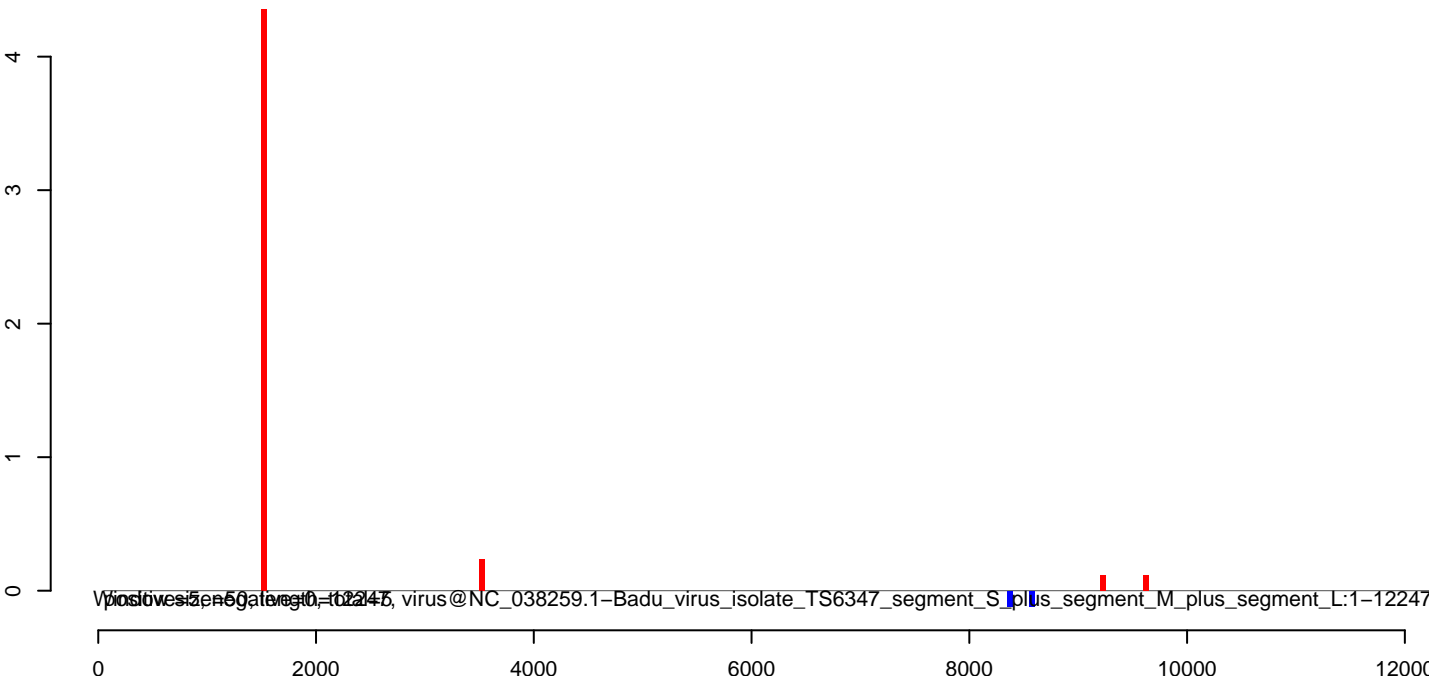
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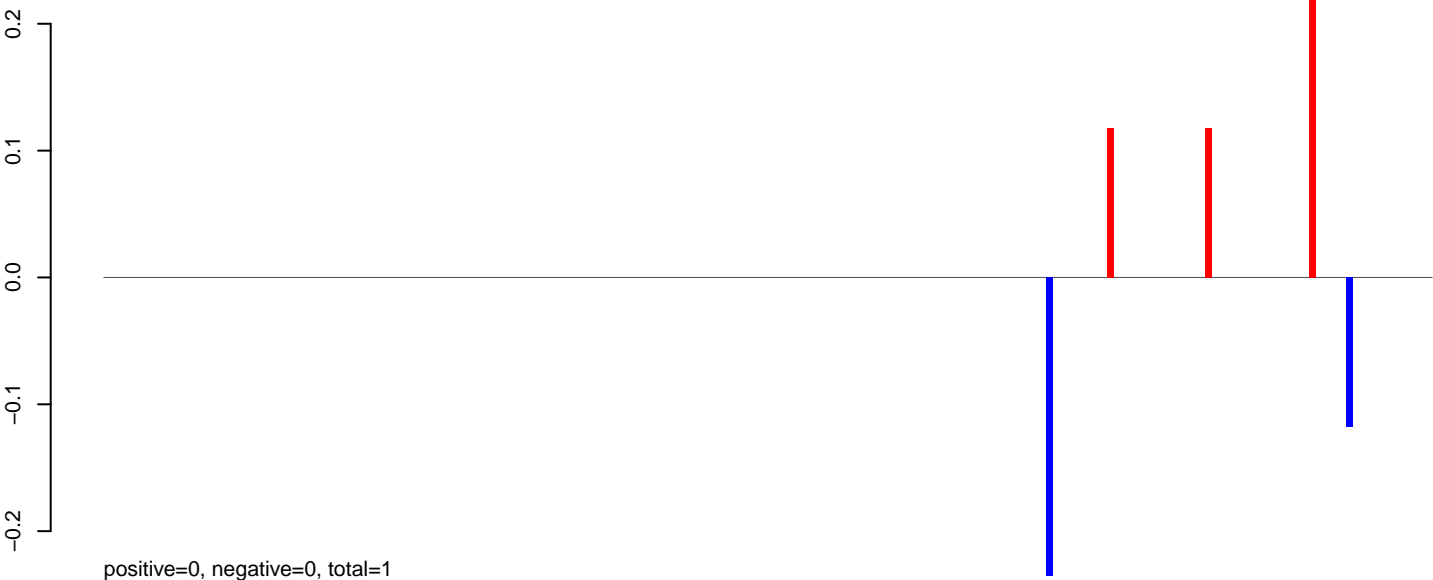
AeAeg_Aag2_DENV1.24_35.rep



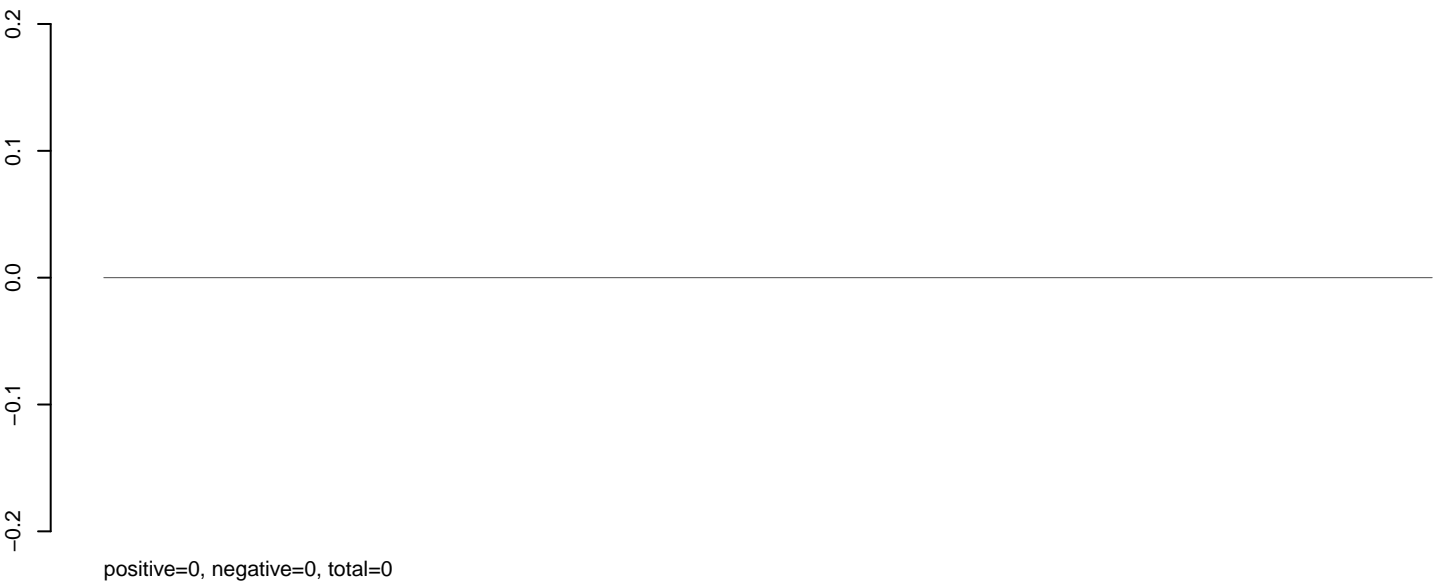
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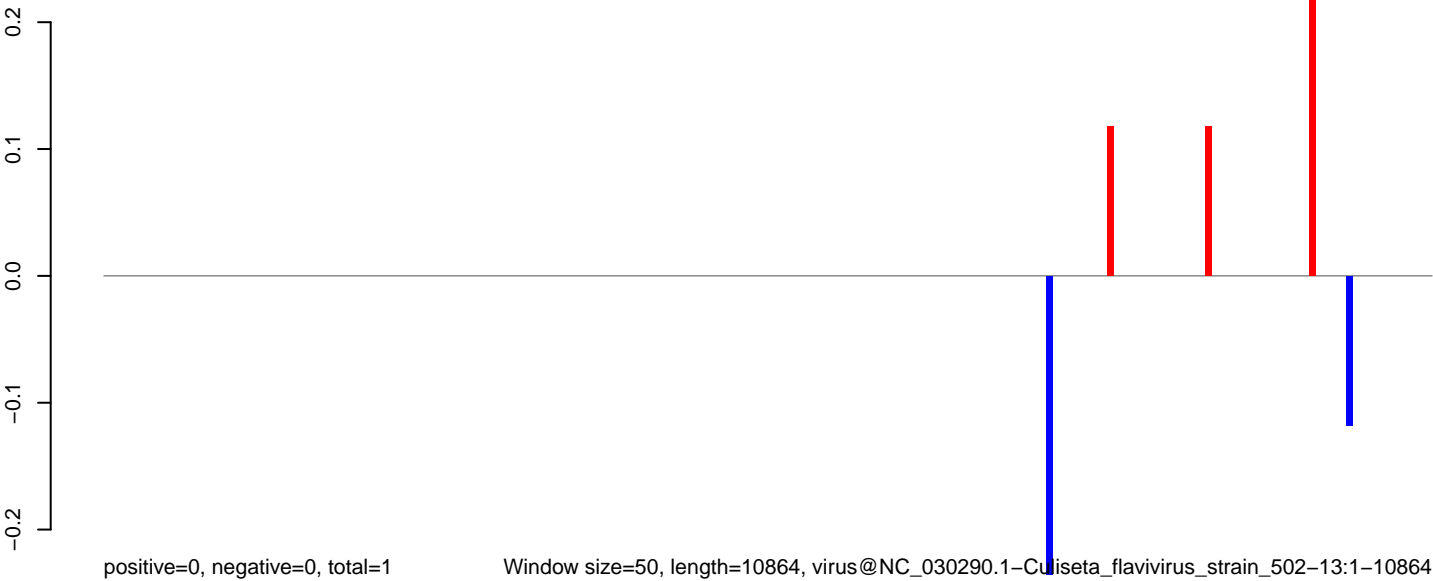
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AeAeg_Aag2_DENV1.24_35.rep



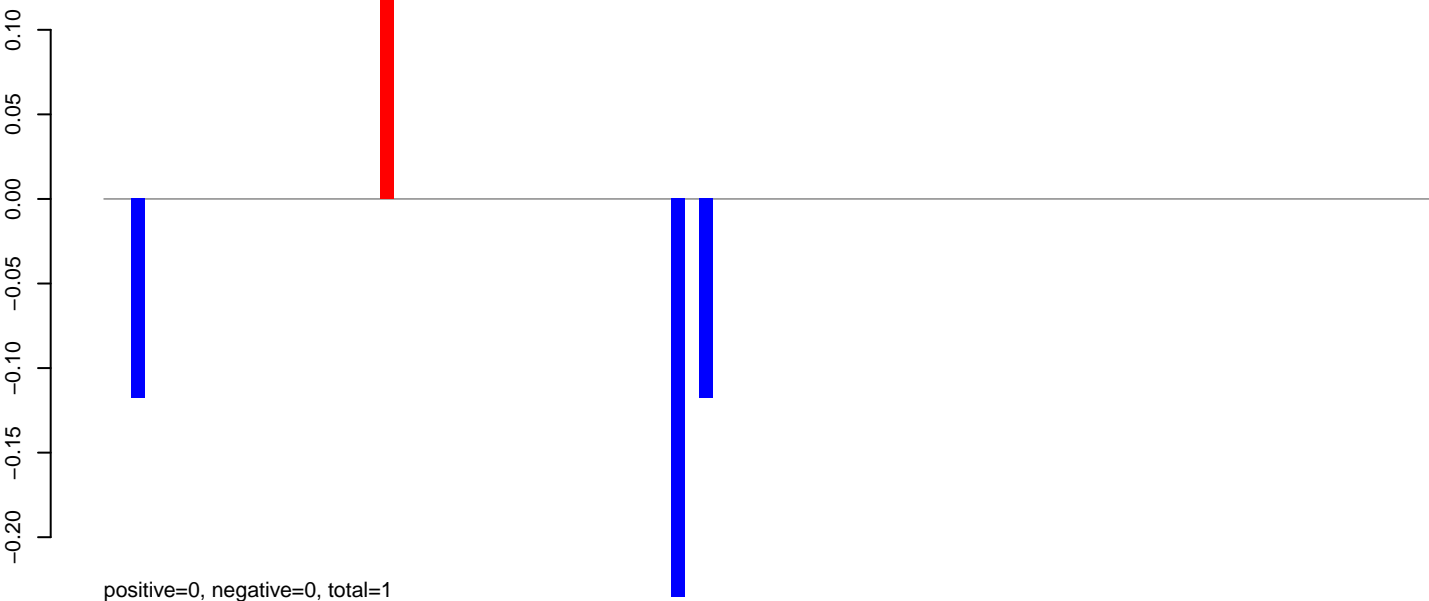
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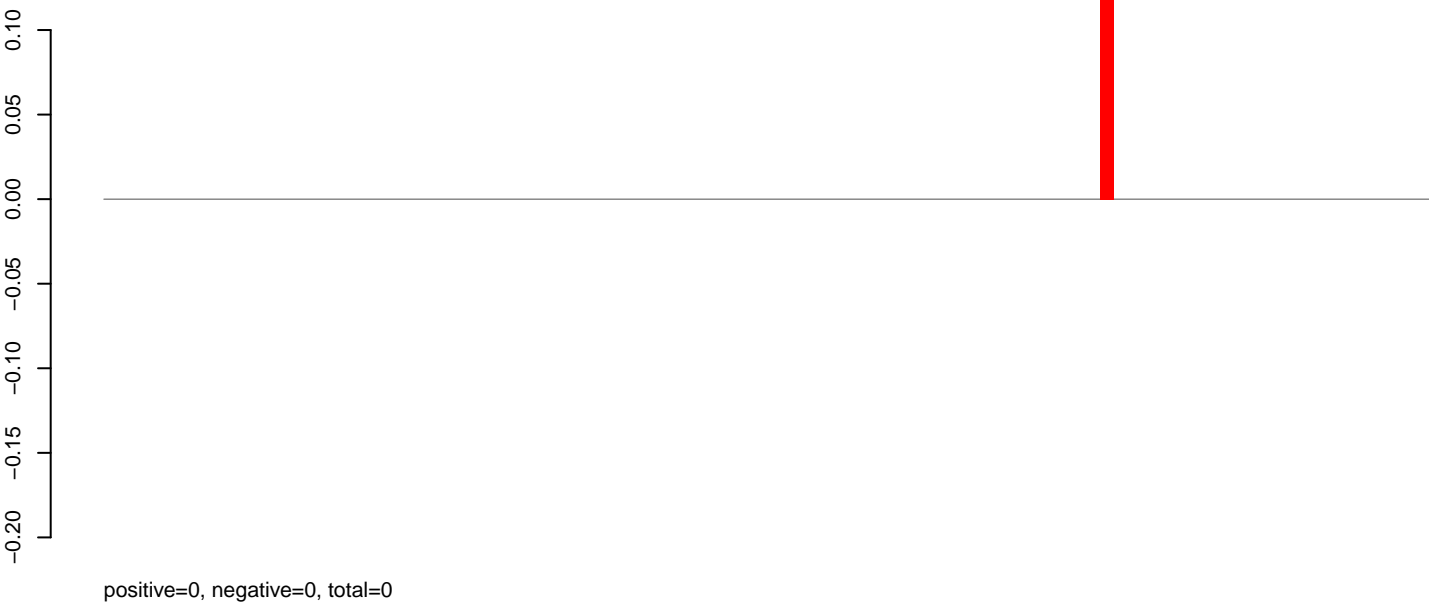
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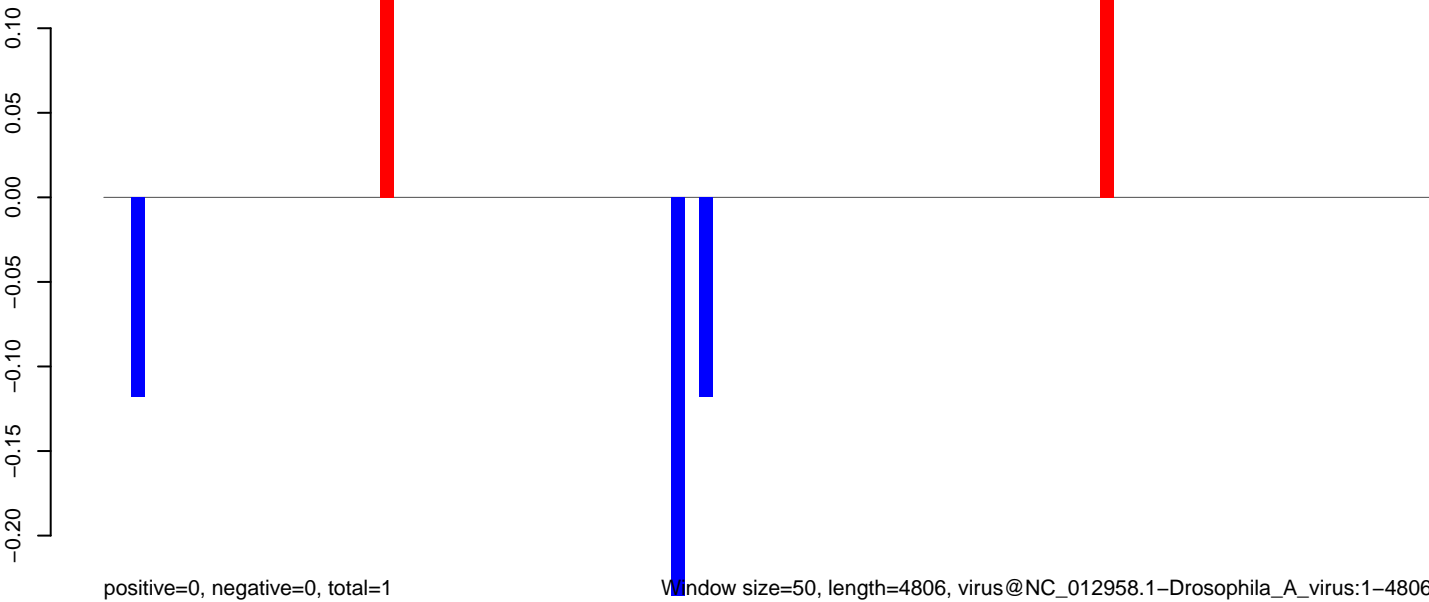
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



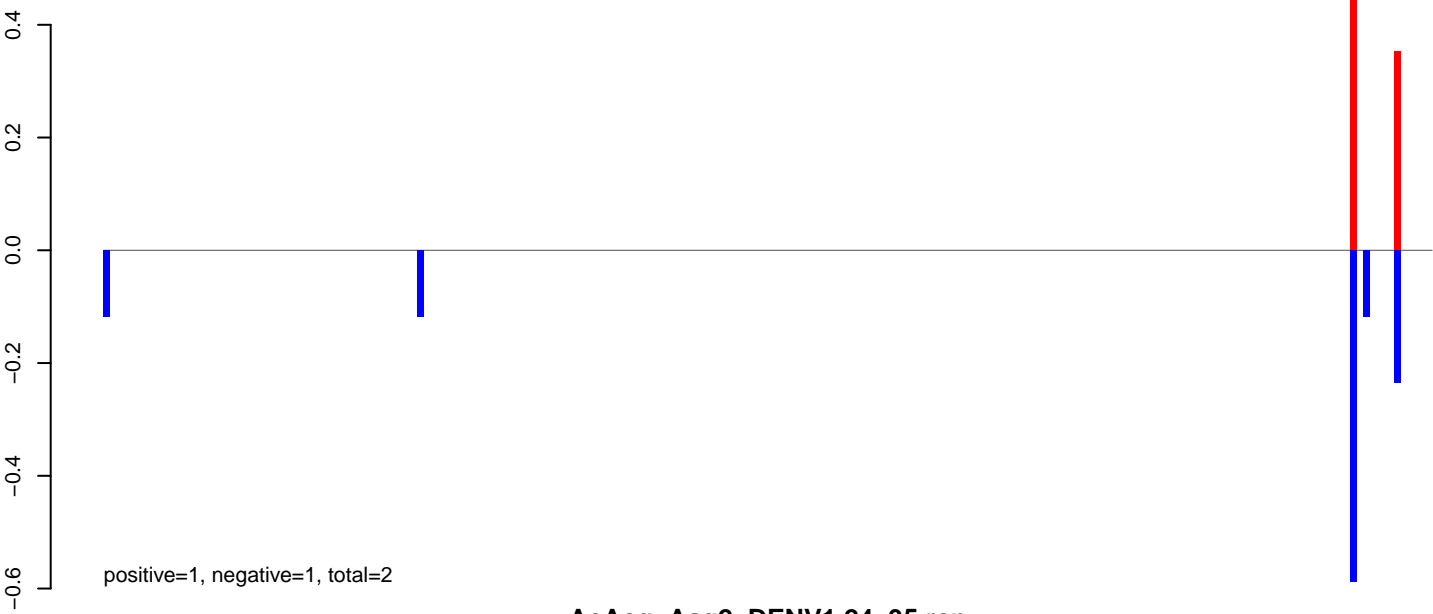
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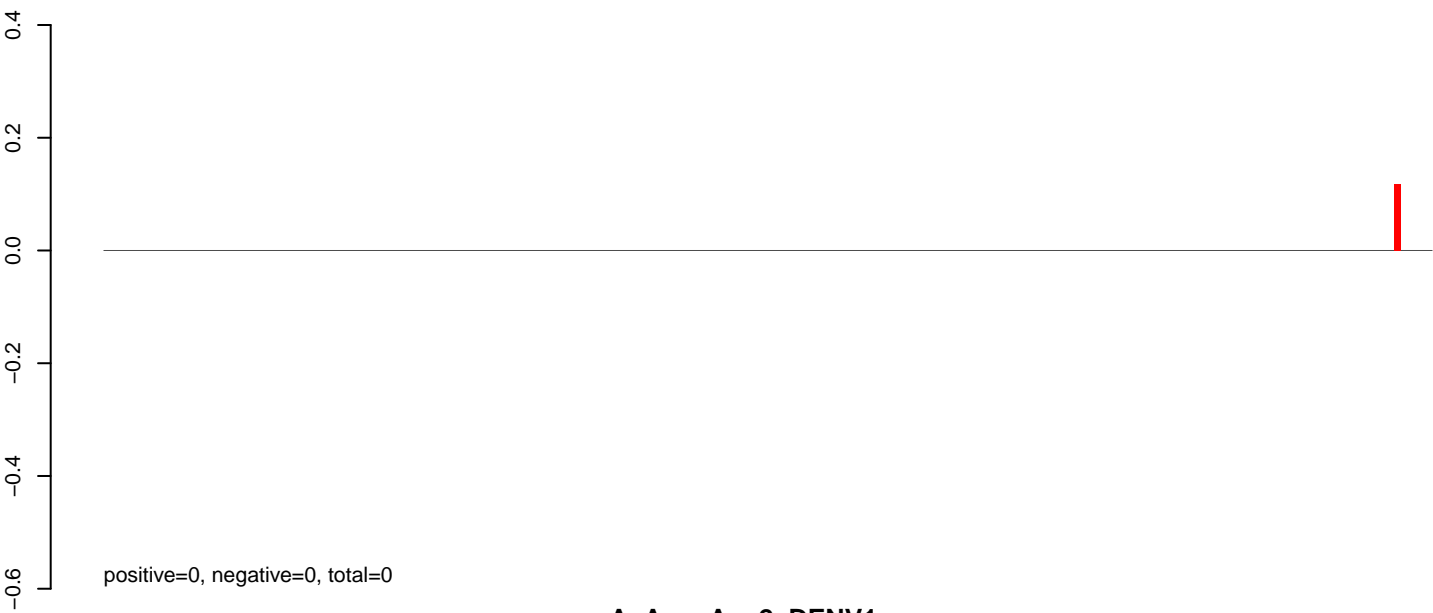
Window size=50, length=4806, virus@NC_012958.1-Drosophila_A_virus:1-4806

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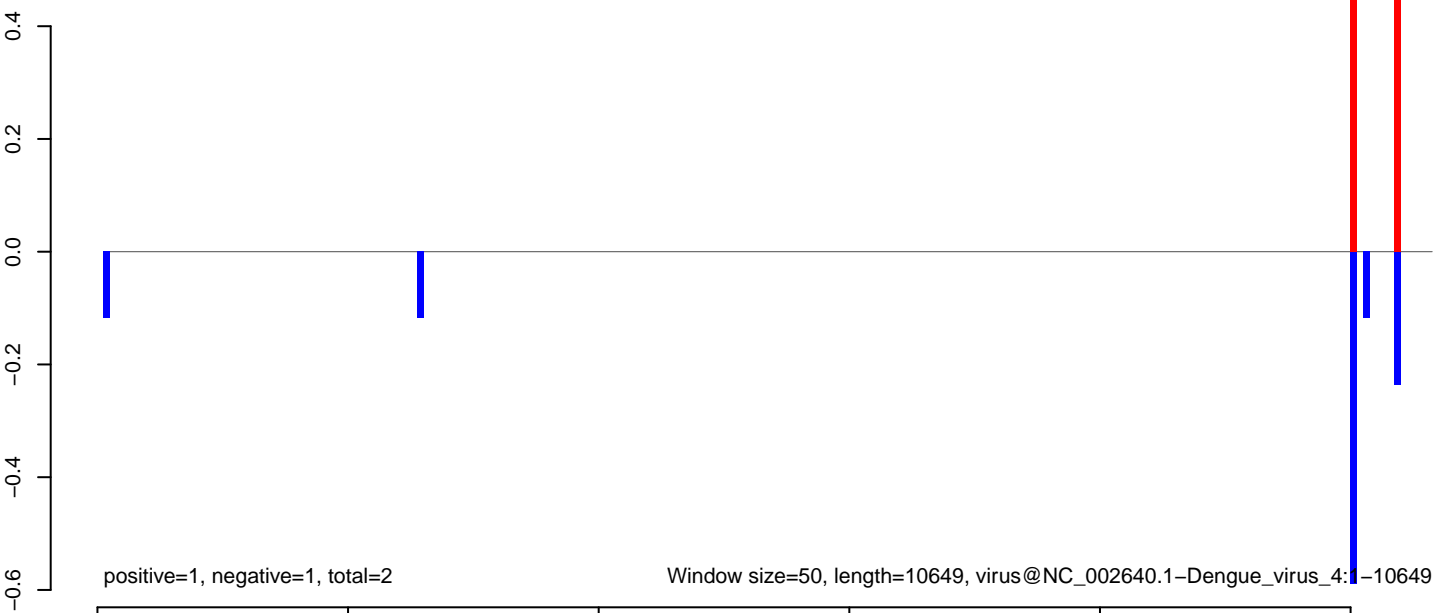
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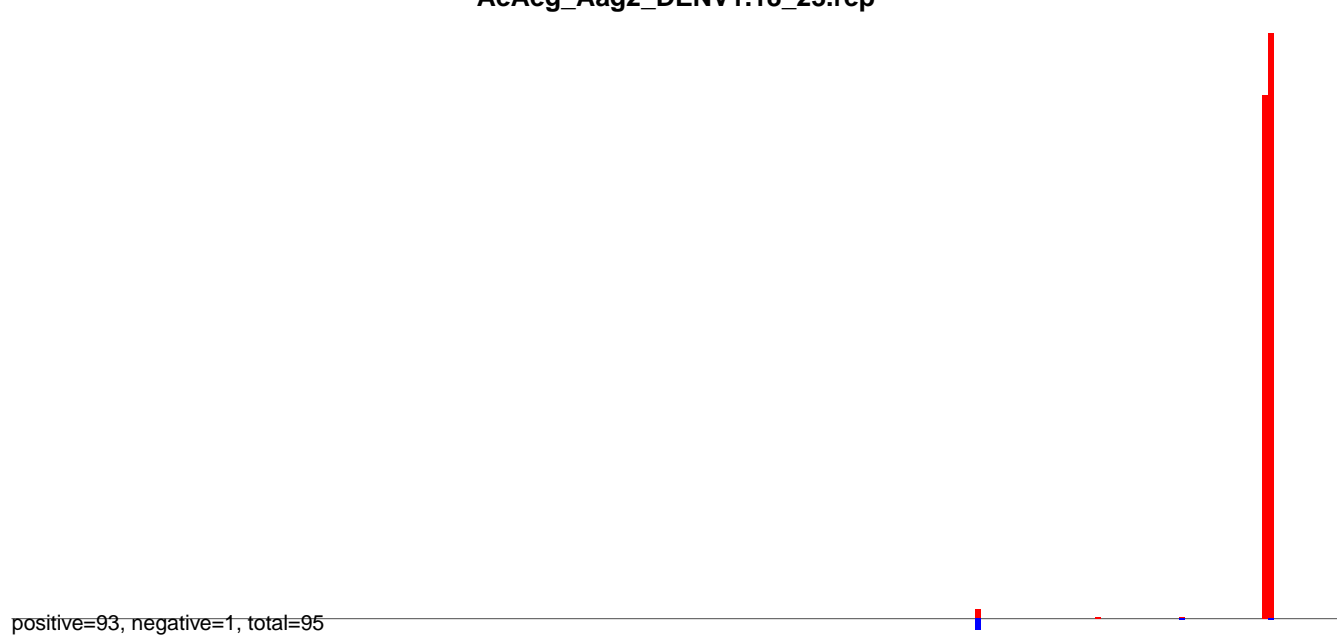
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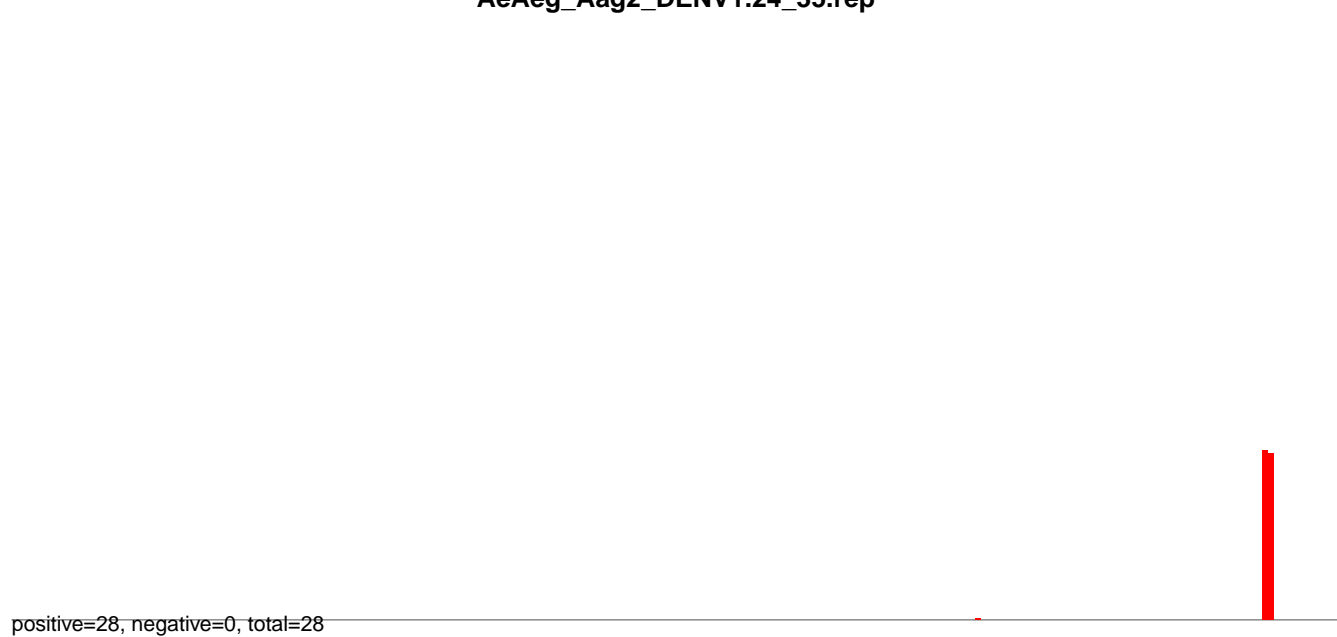
AeAeg_Aag2_DENV1.rep



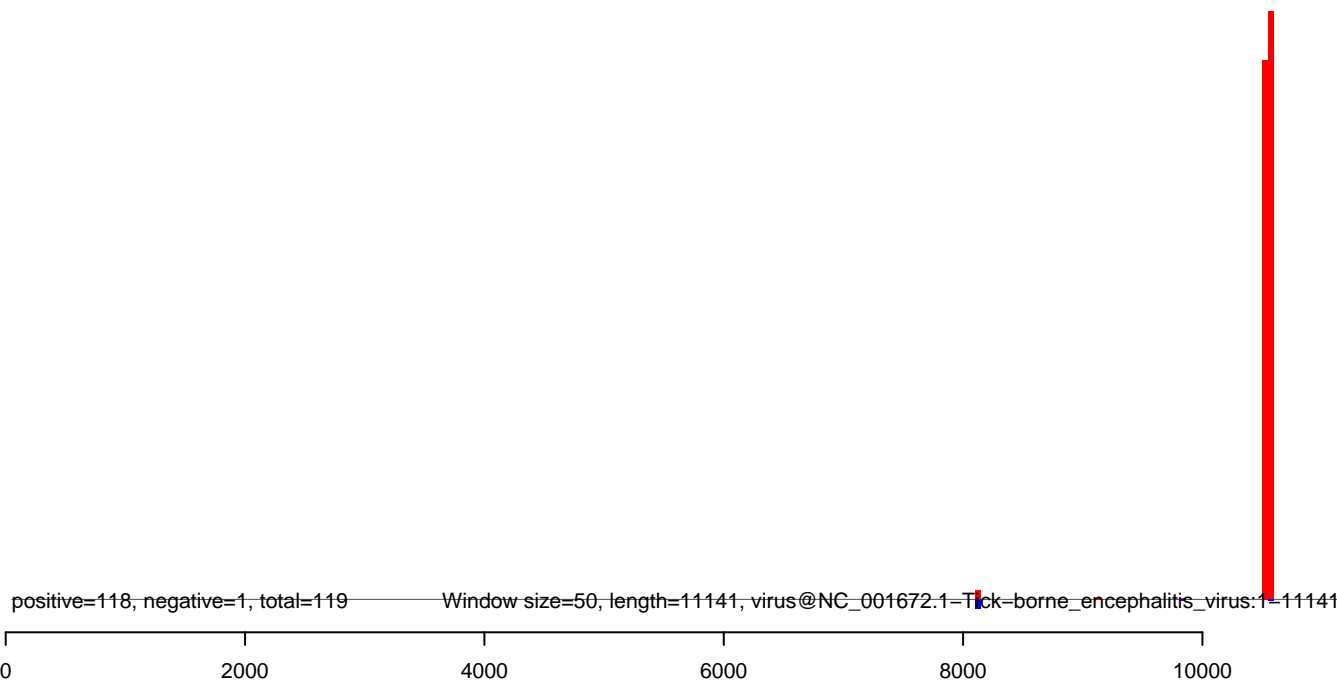
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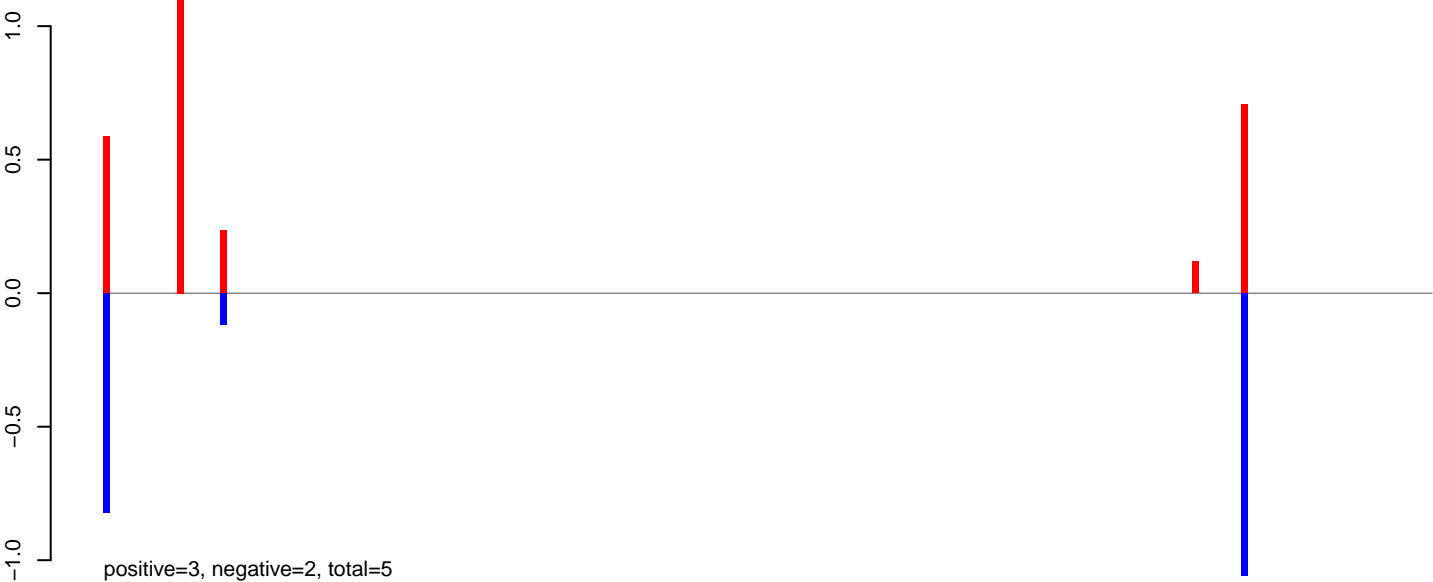
AeAeg_Aag2_DENV1.24_35.rep



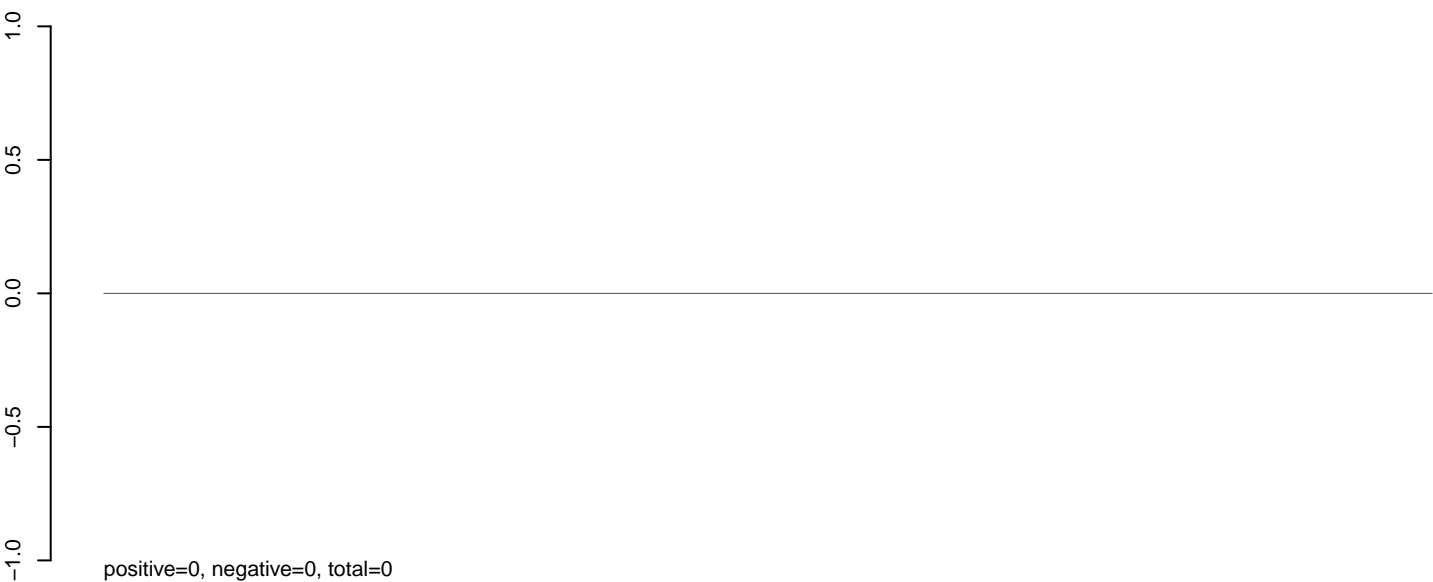
AeAeg_Aag2_DENV1.rep



AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



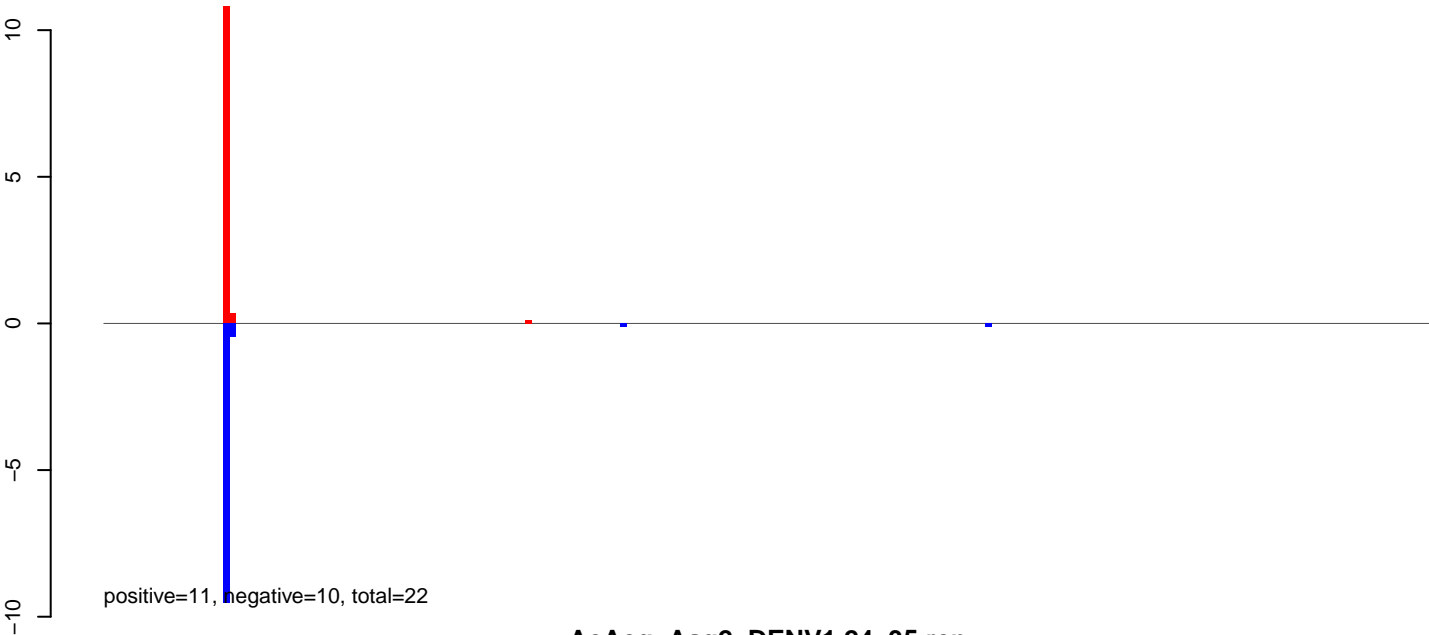
AeAeg_Aag2_DENV1.rep



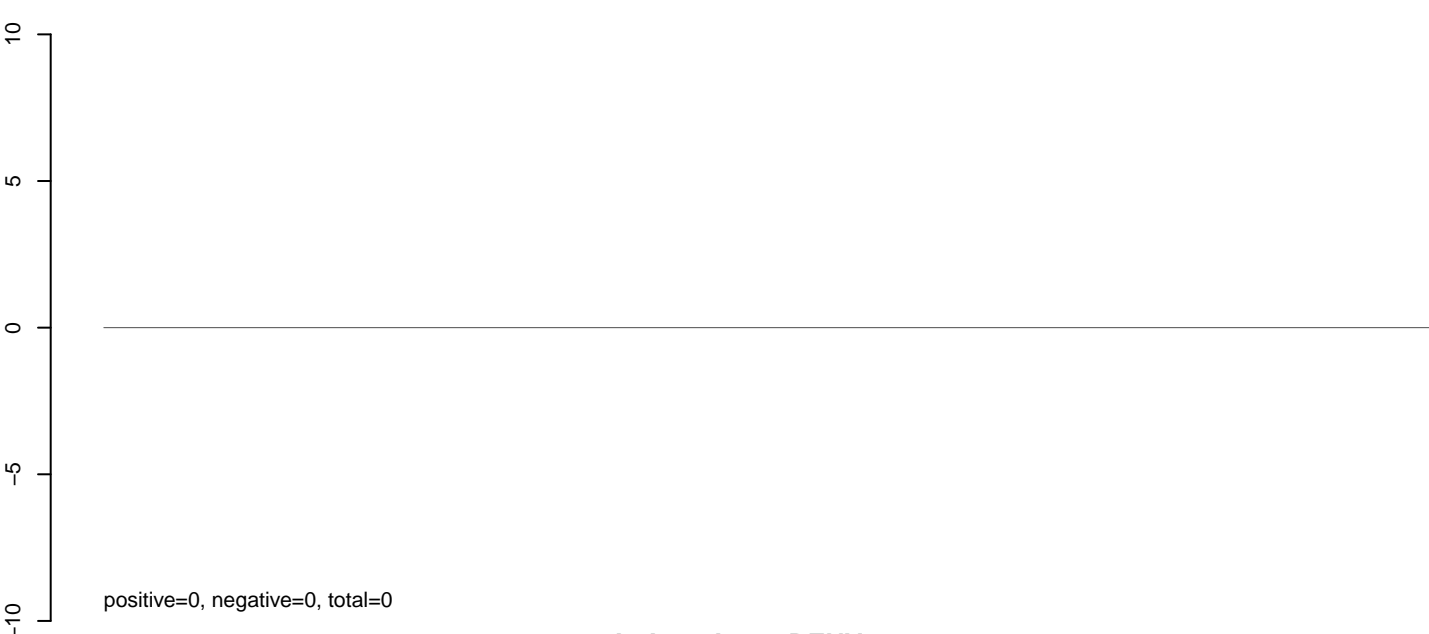
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0 2000 4000 6000 8000 10000

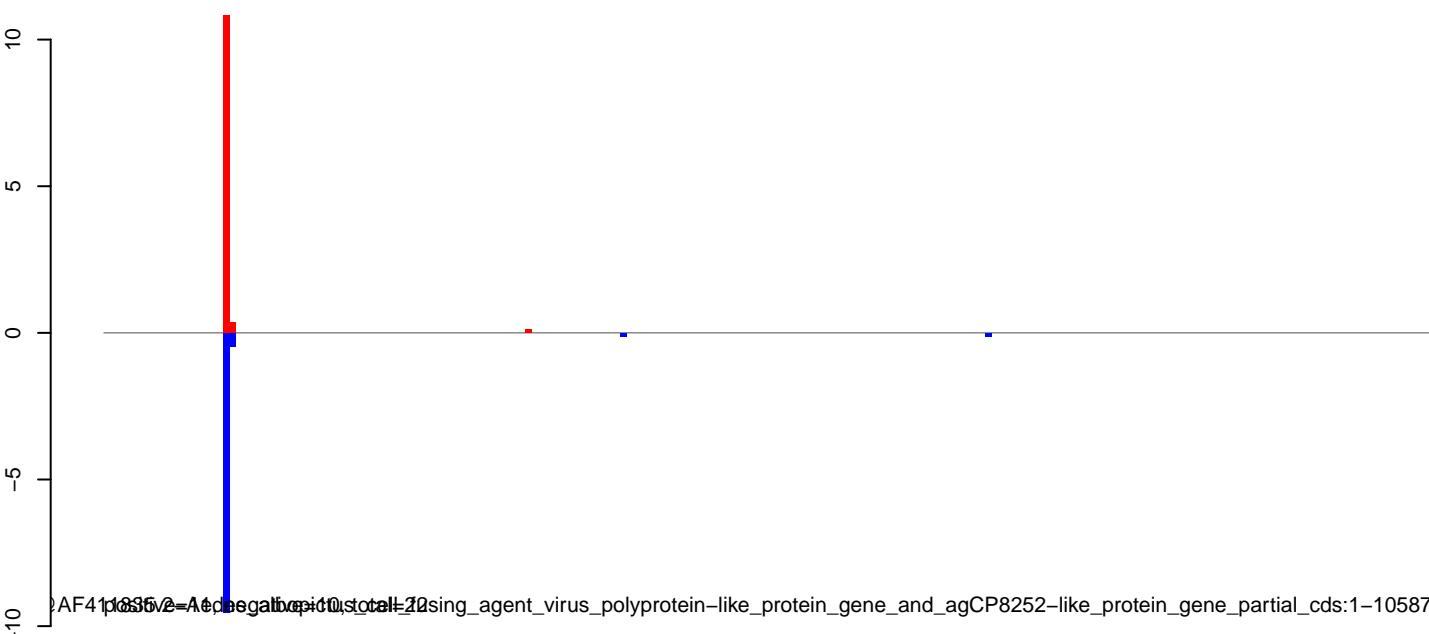
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

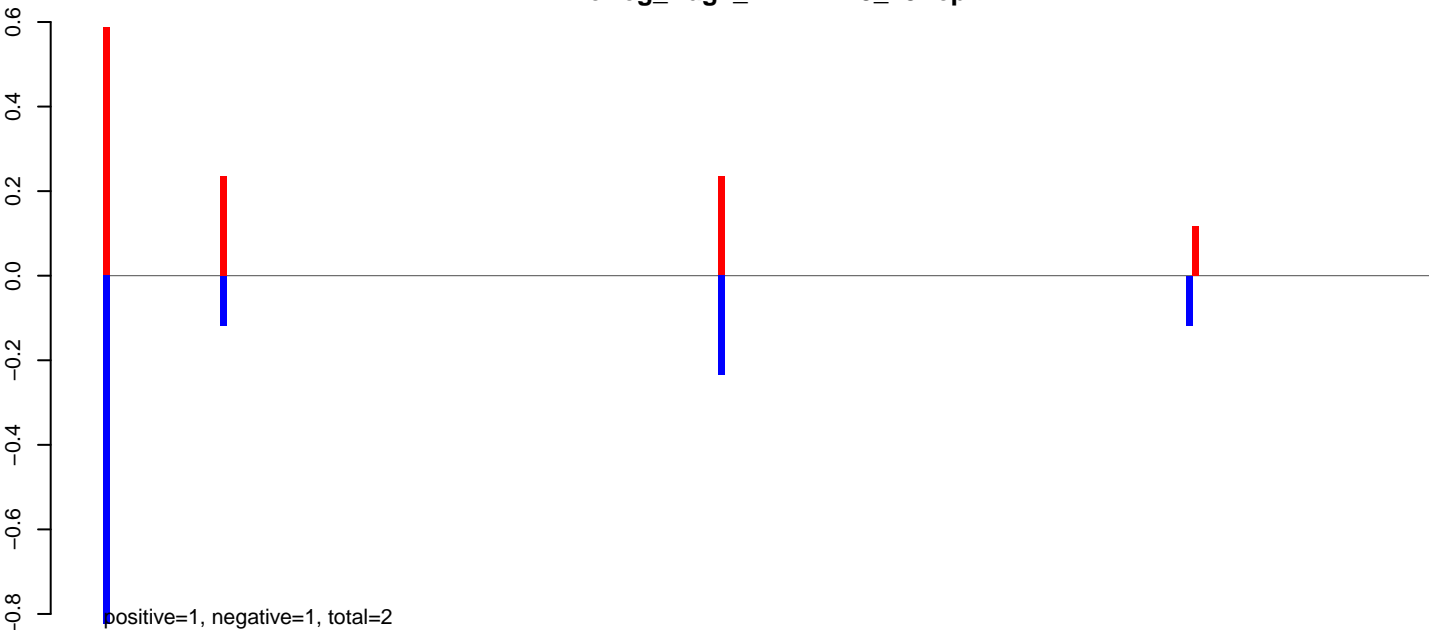


AeAeg_Aag2_DENV1.rep

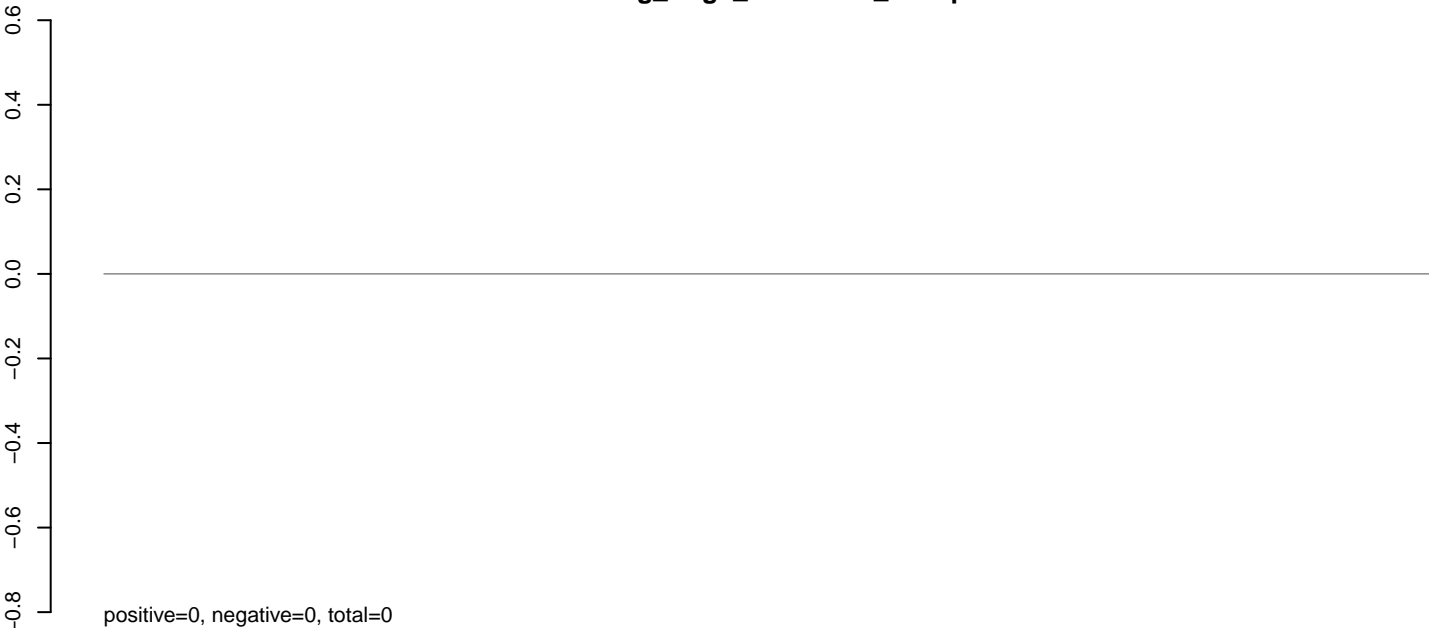


AF416352-Aedes albopictus CP8252-like protein gene partial cds:1-10587

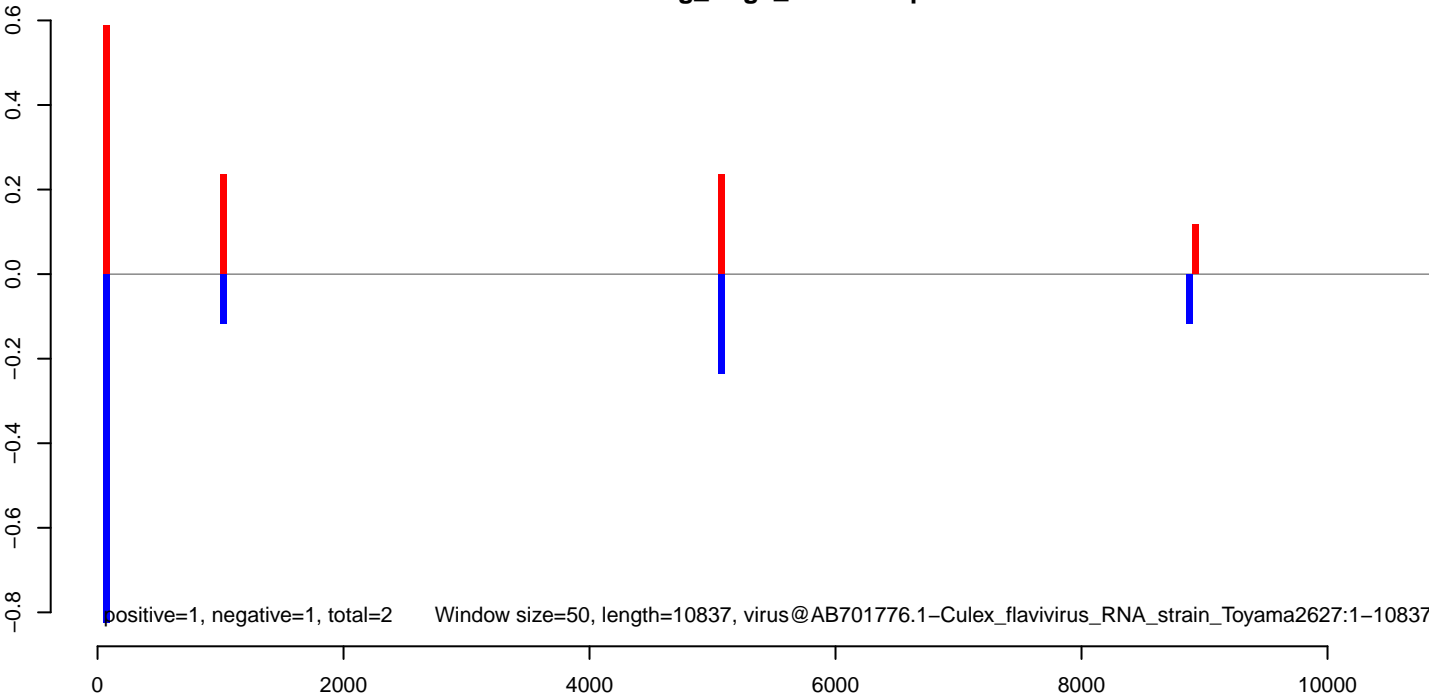
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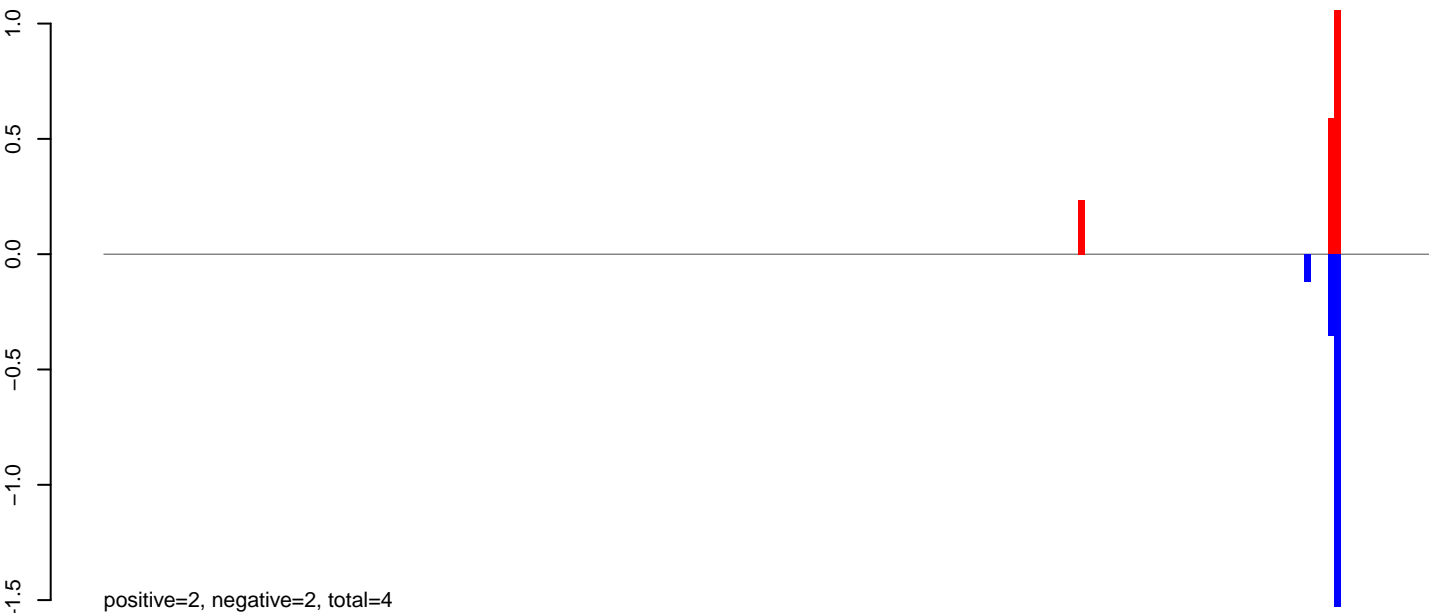
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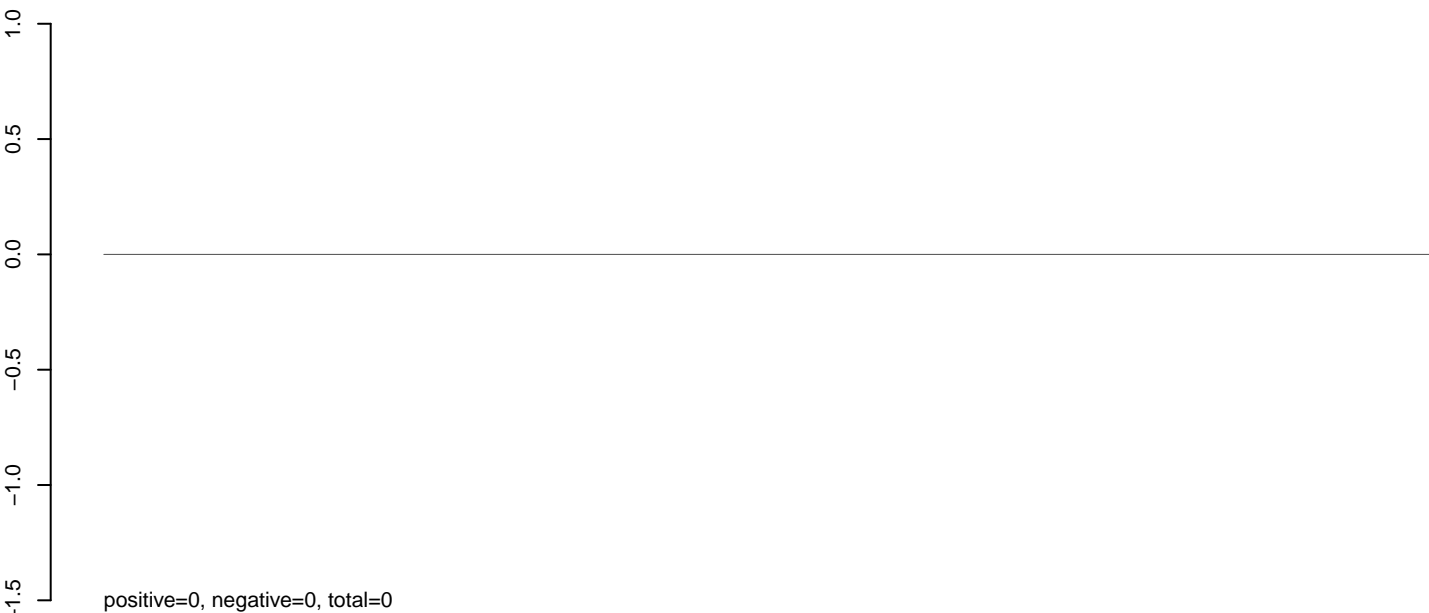
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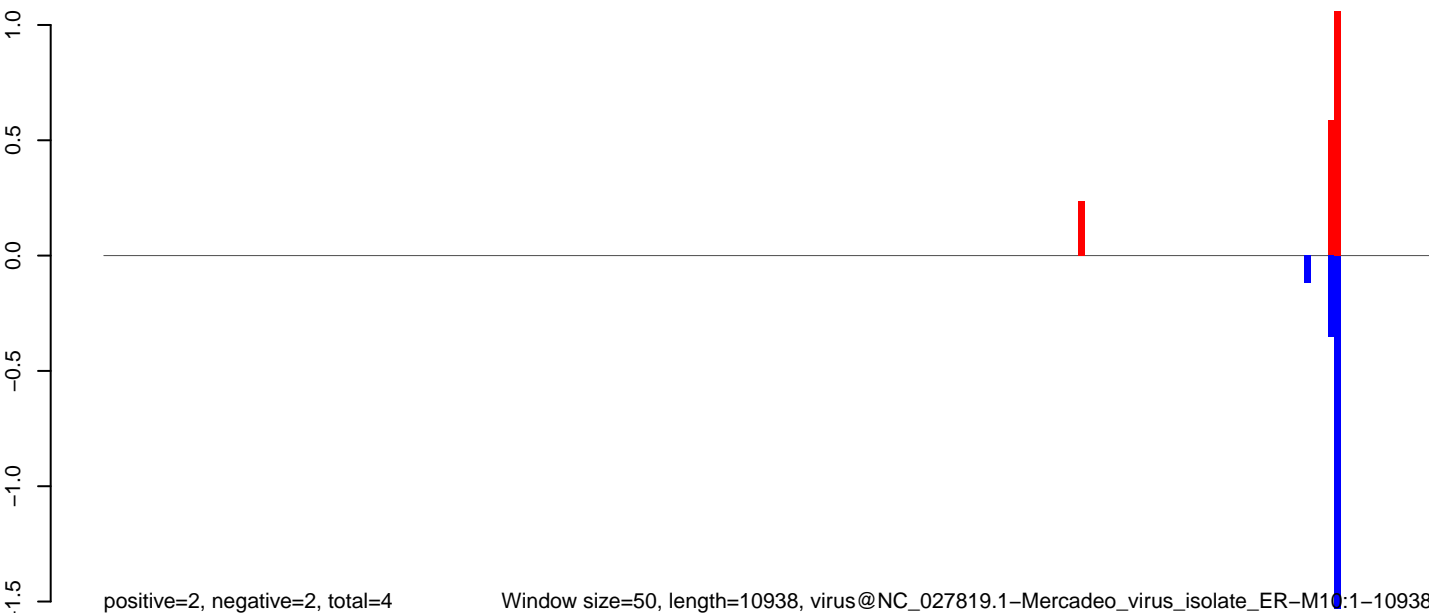
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AeAeg_Aag2_DENV1.24_35.rep



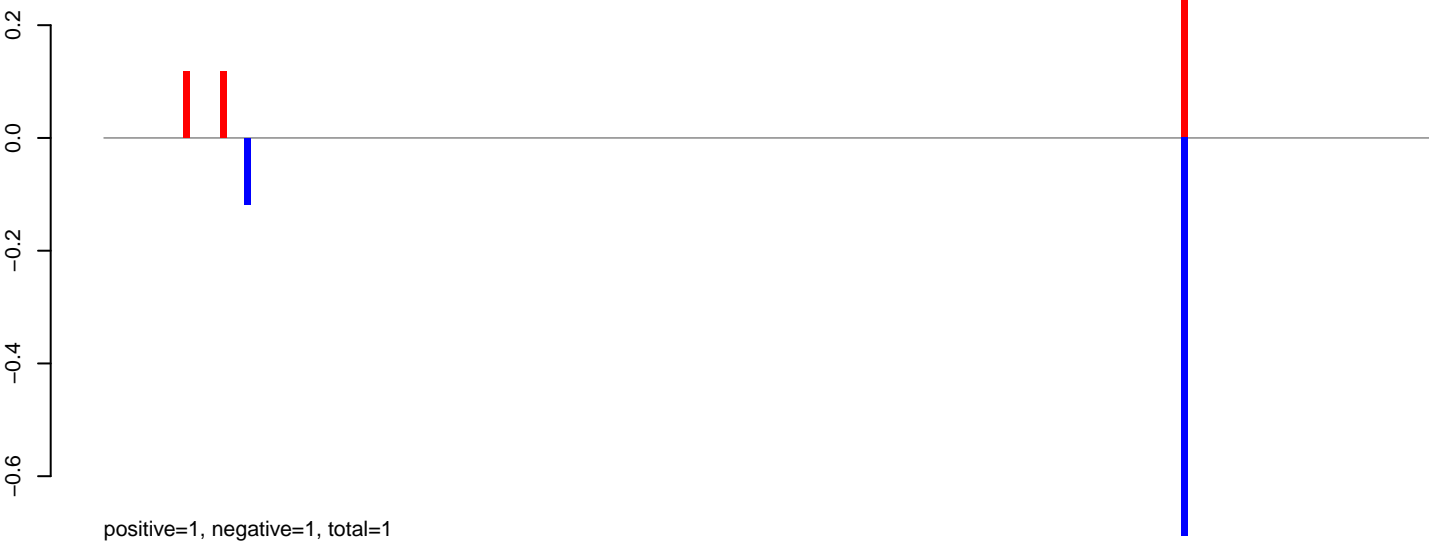
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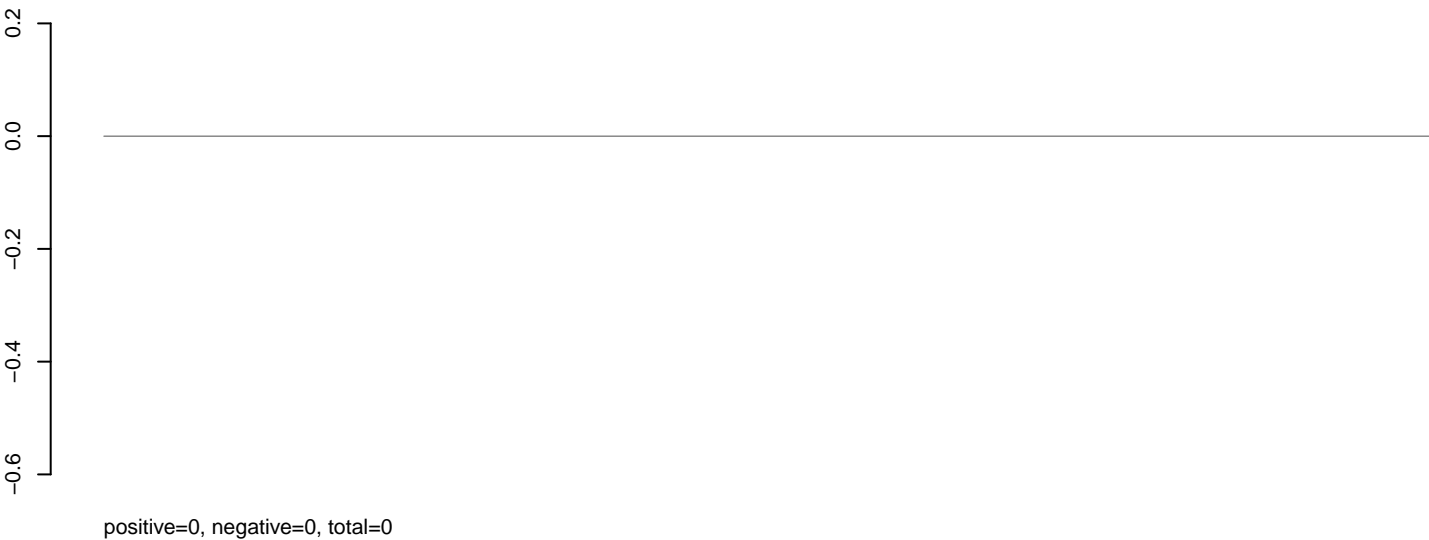
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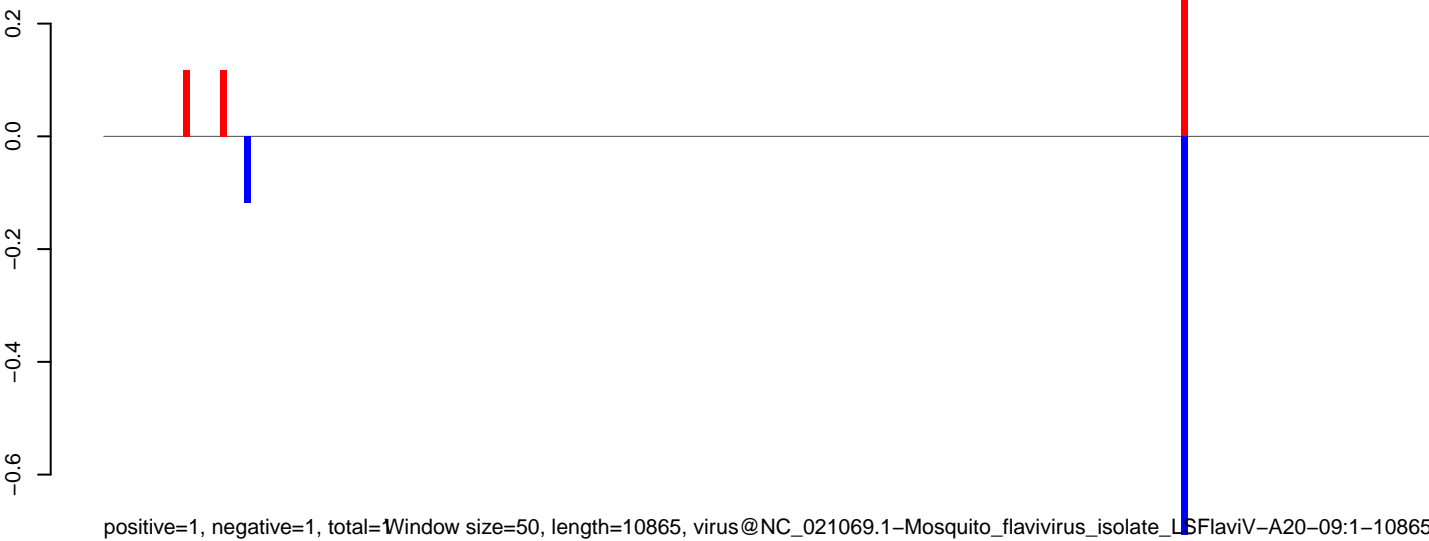
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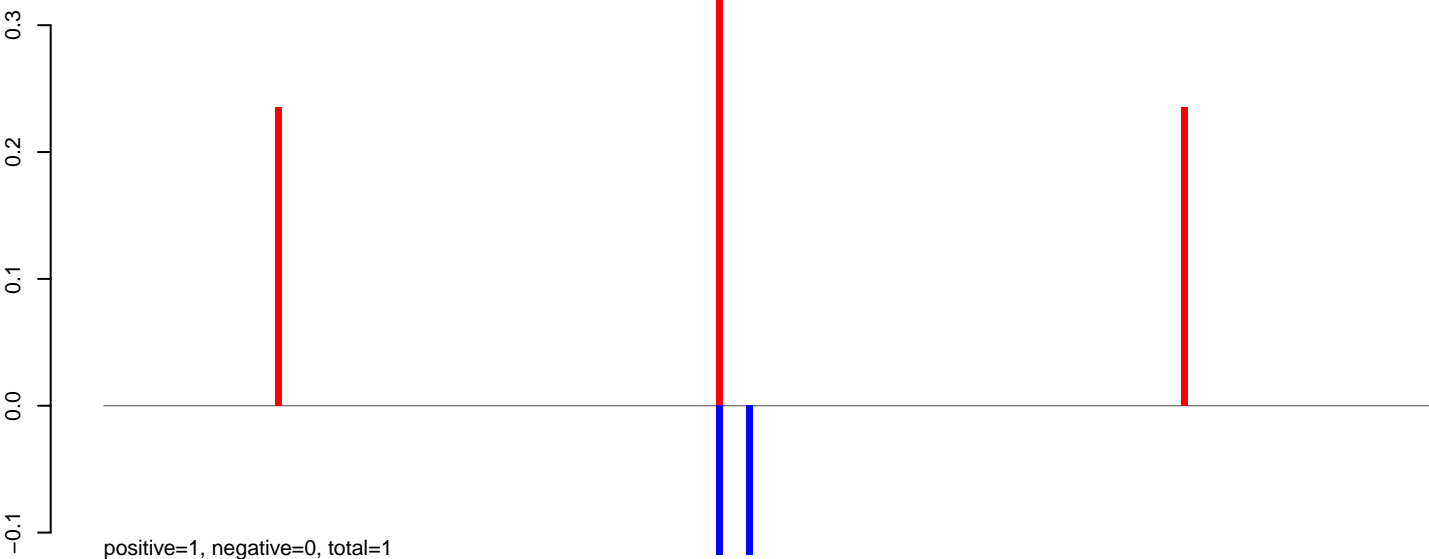
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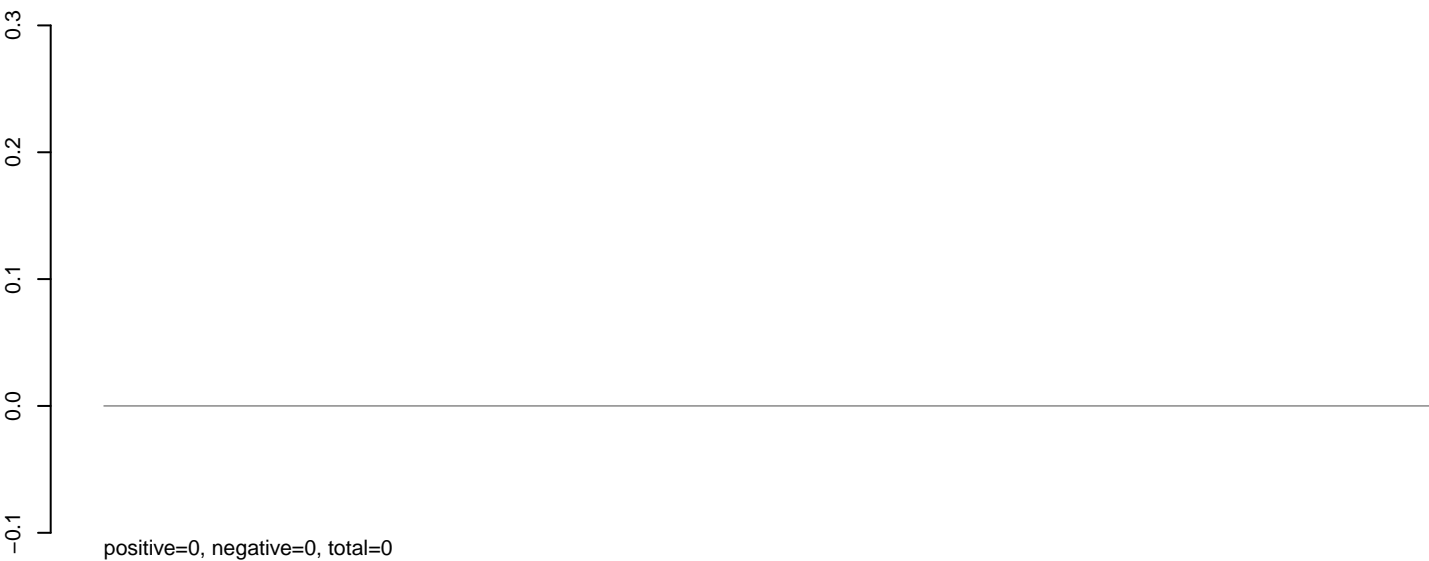
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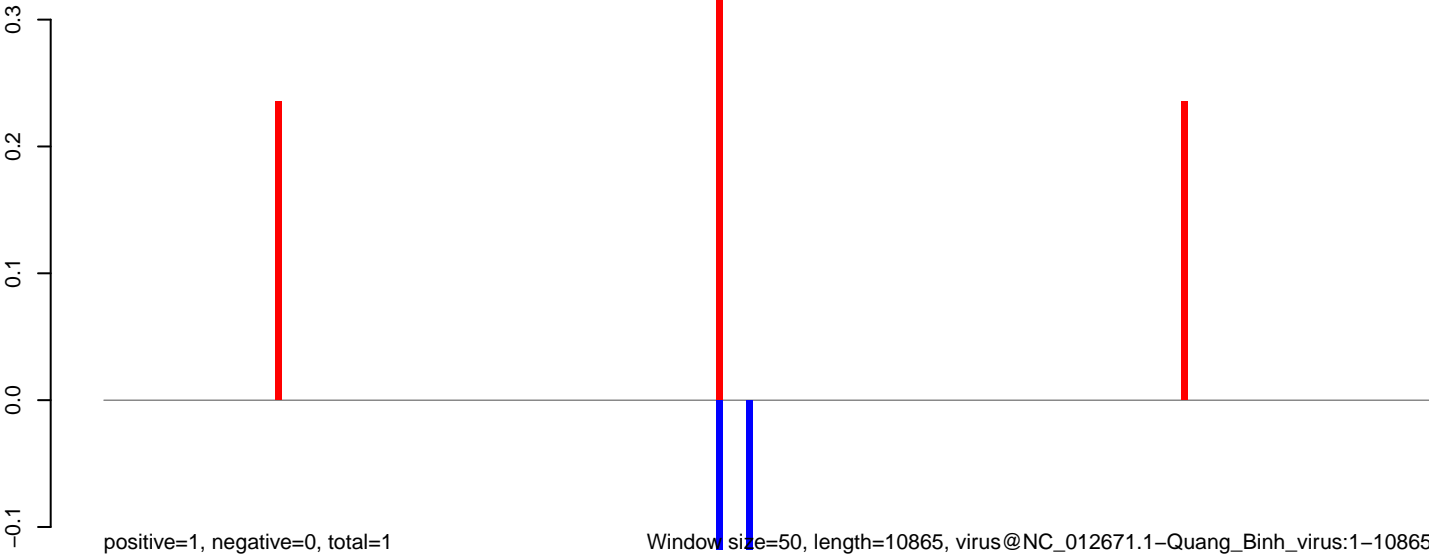
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AeAeg_Aag2_DENV1.24_35.rep

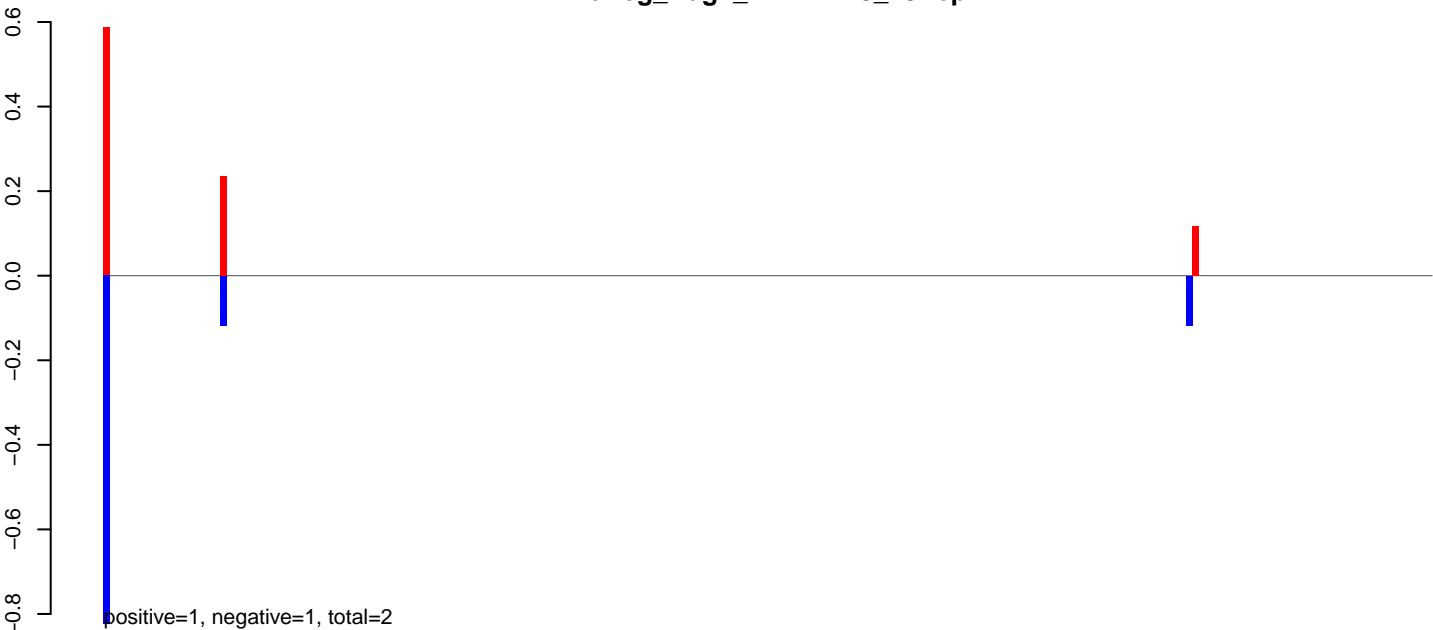


AeAeg_Aag2_DENV1.rep

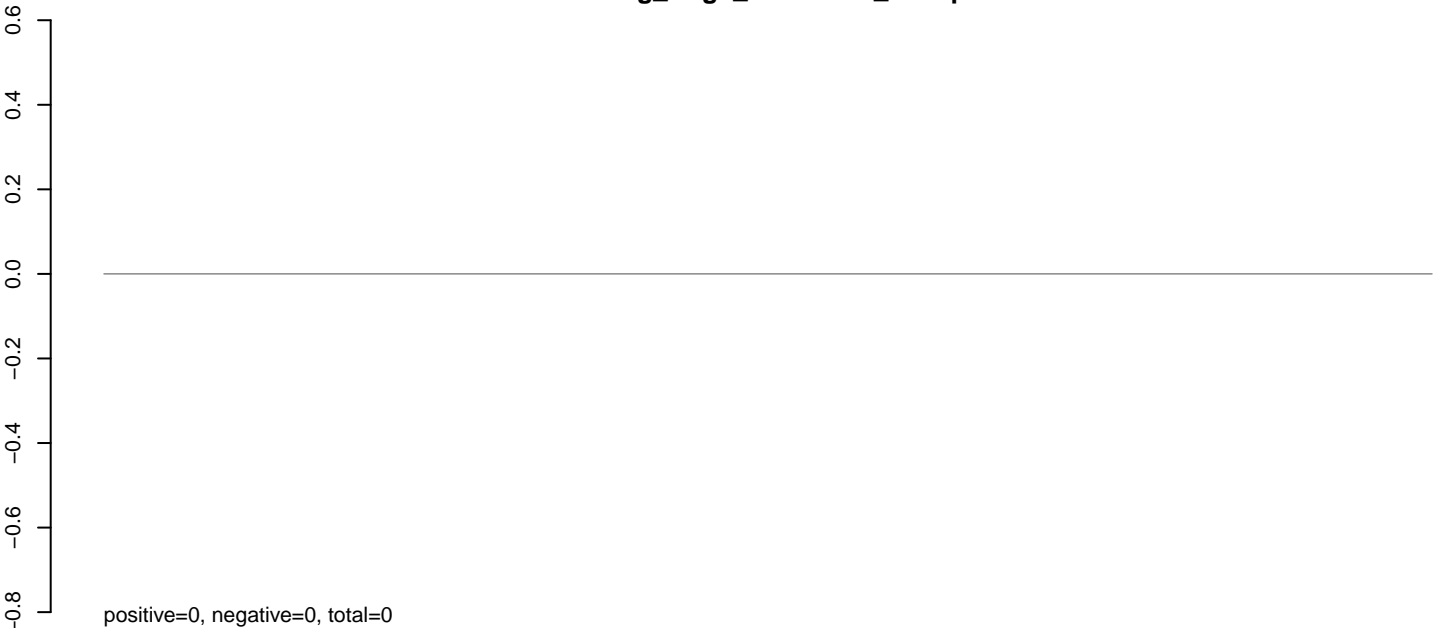


0 2000 4000 6000 8000 10000

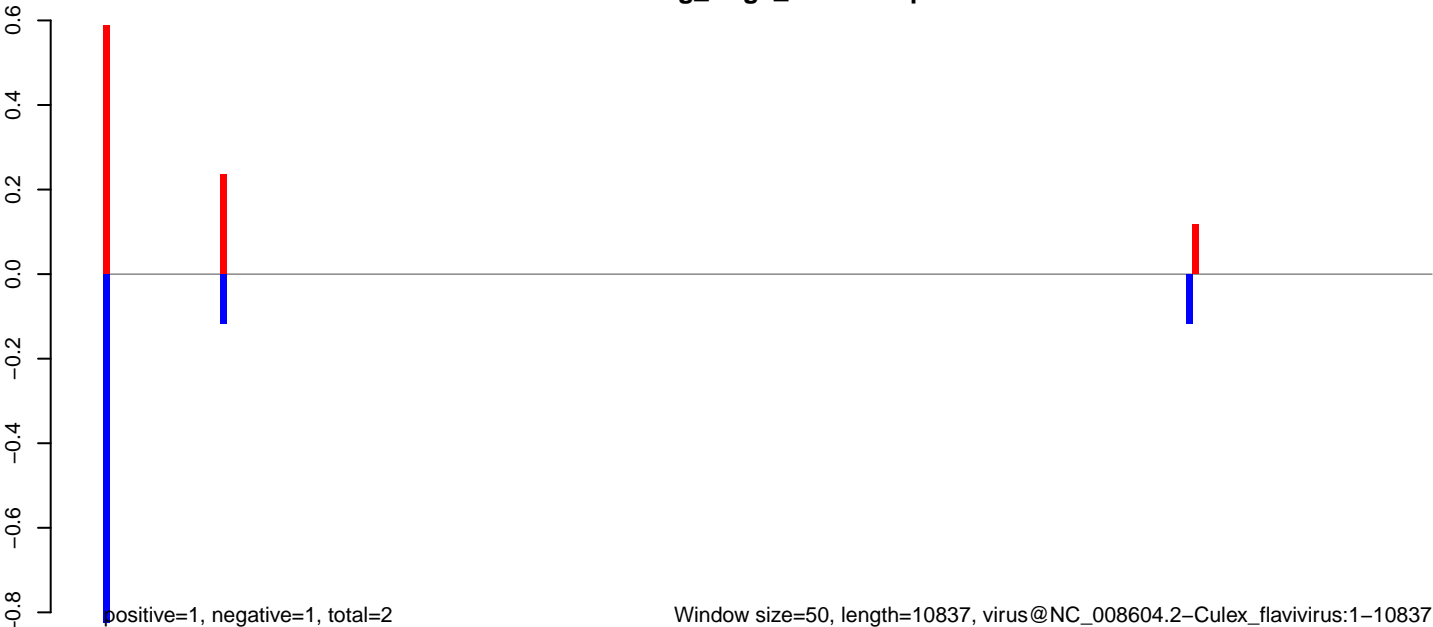
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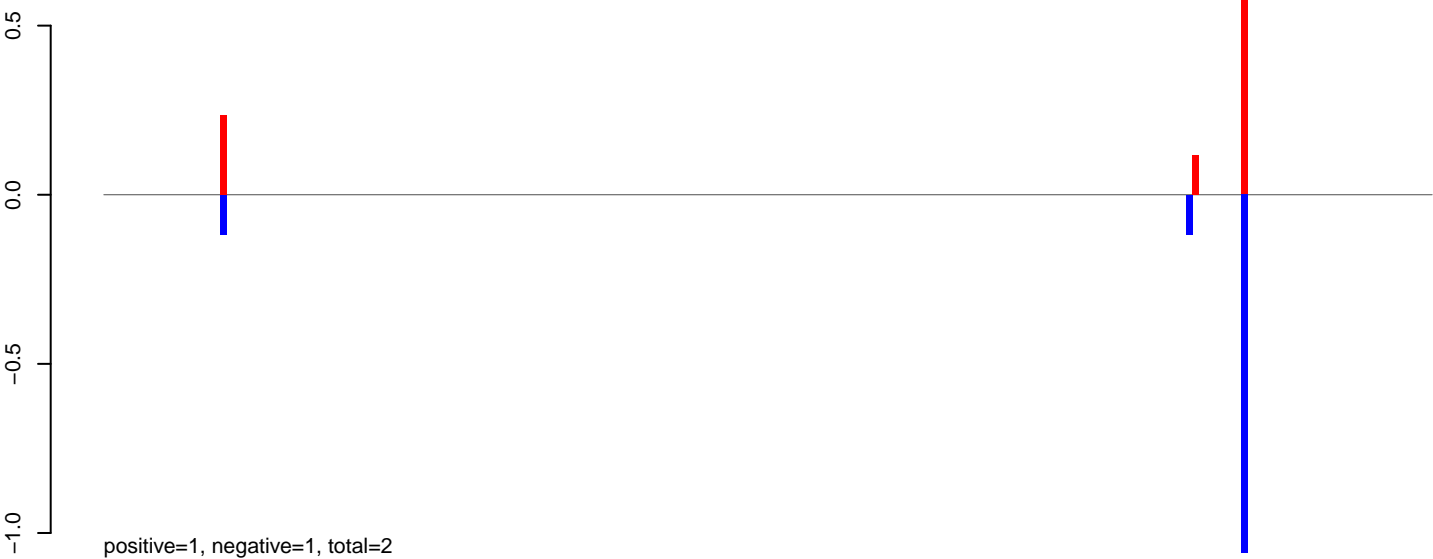
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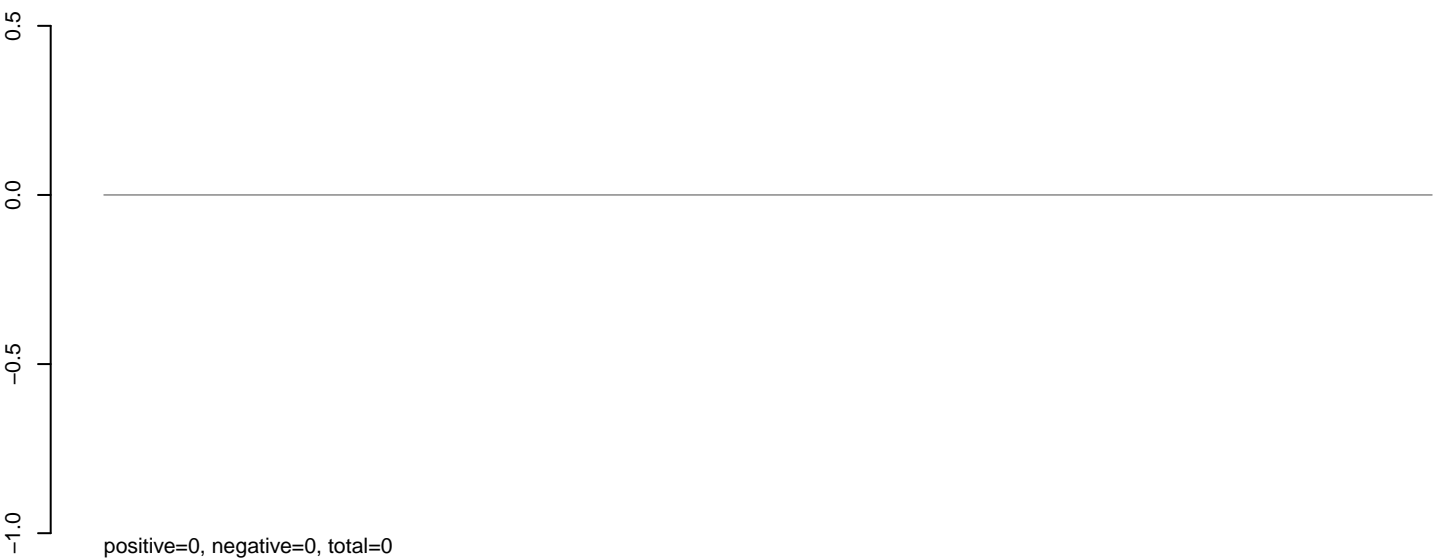
AeAeg_Aag2_DENV1.rep



AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



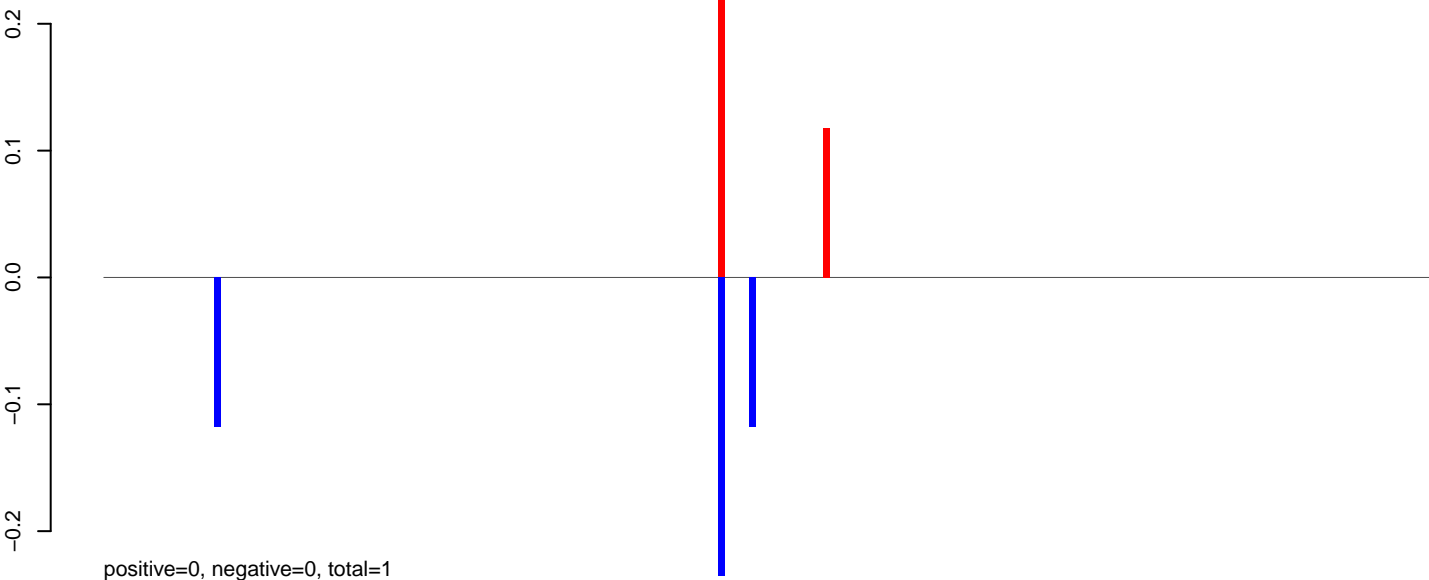
AeAeg_Aag2_DENV1.rep



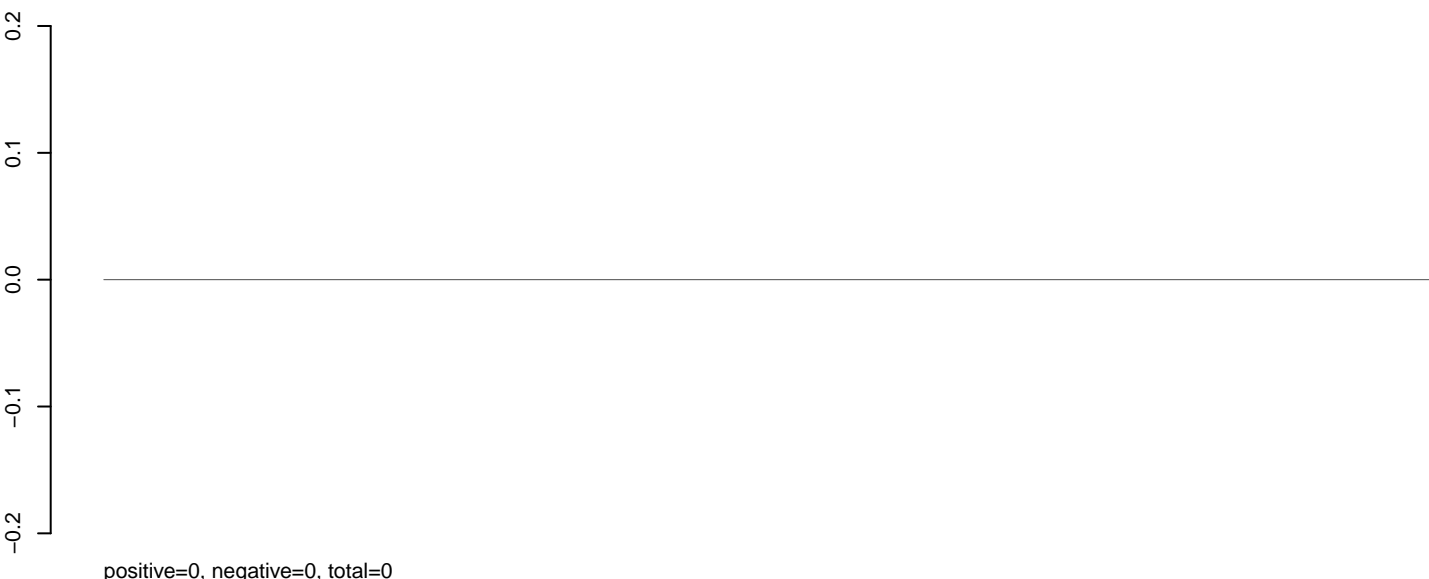
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0 2000 4000 6000 8000 10000

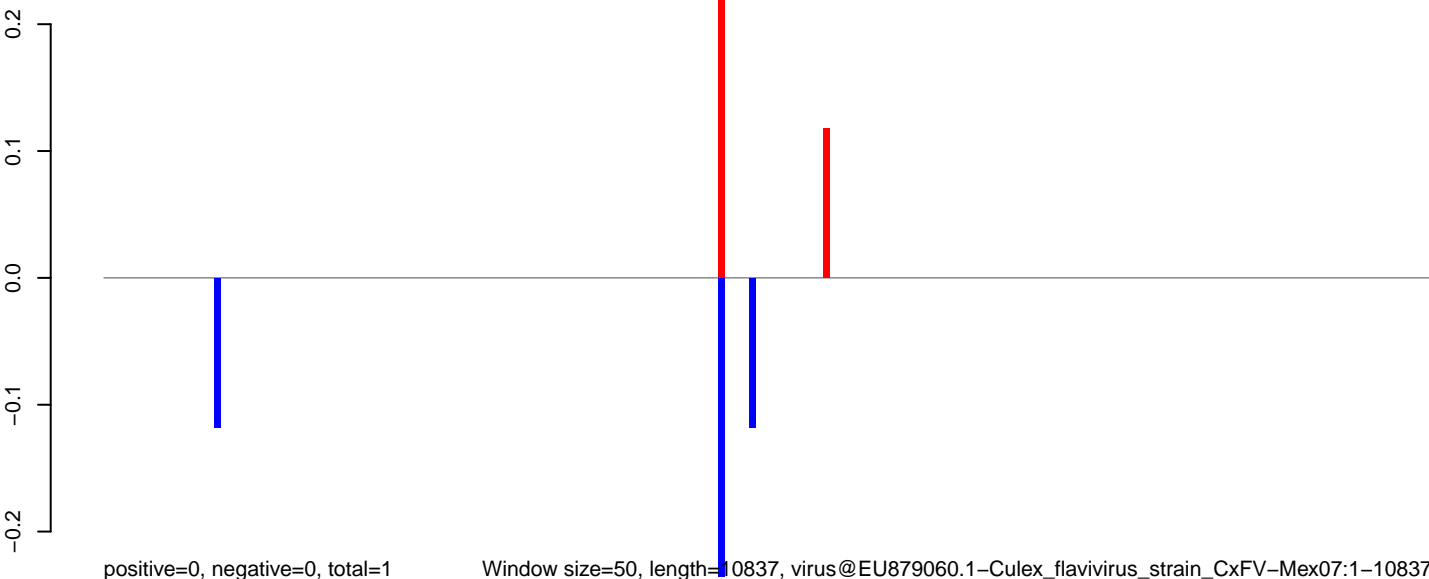
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AeAeg_Aag2_DENV1.24_35.rep



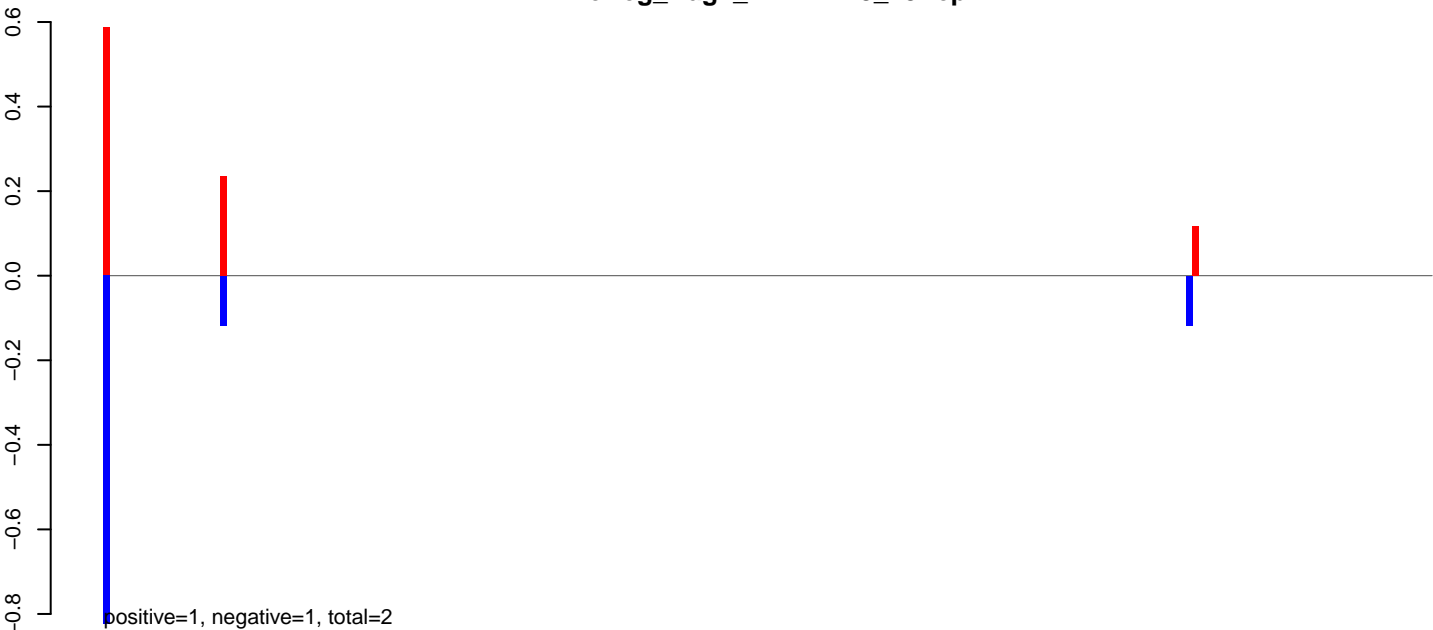
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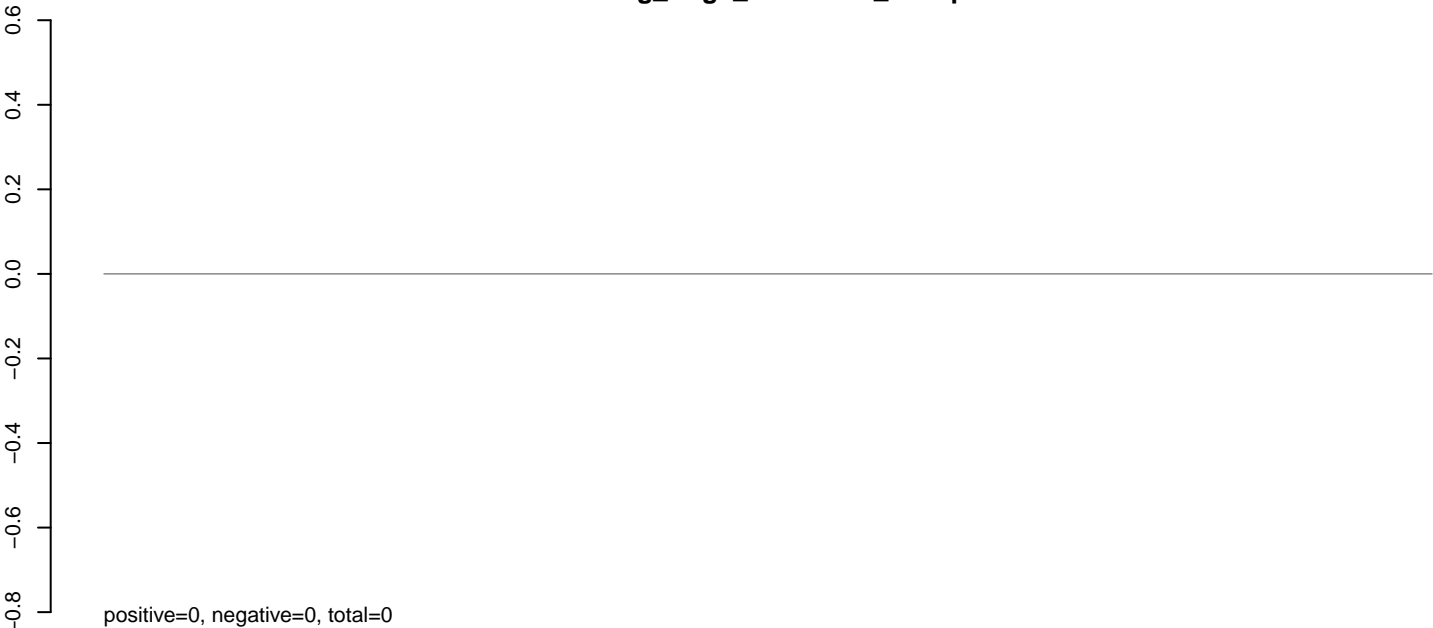
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0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV1.18_23.rep



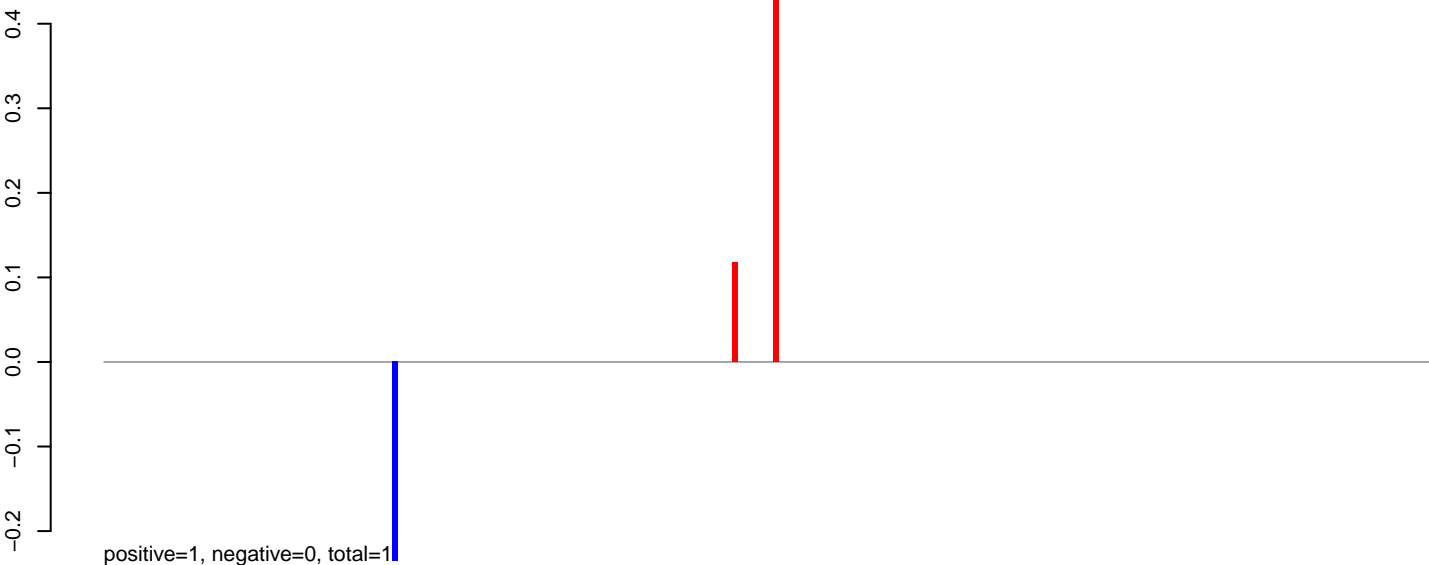
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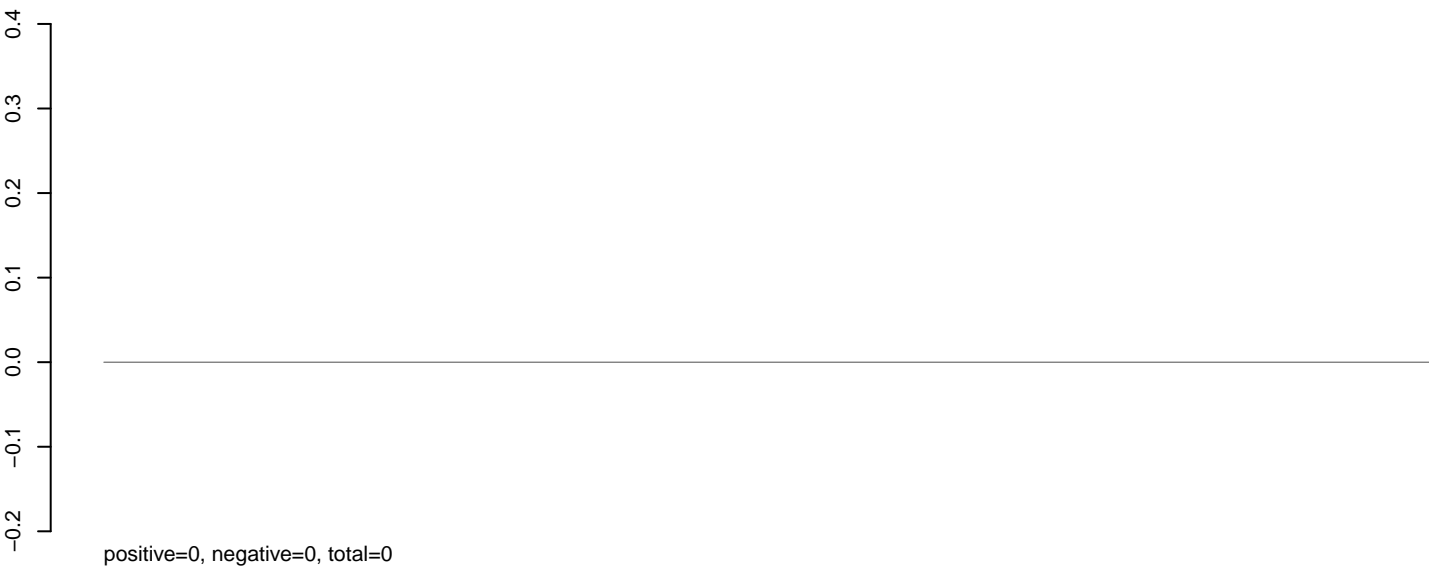
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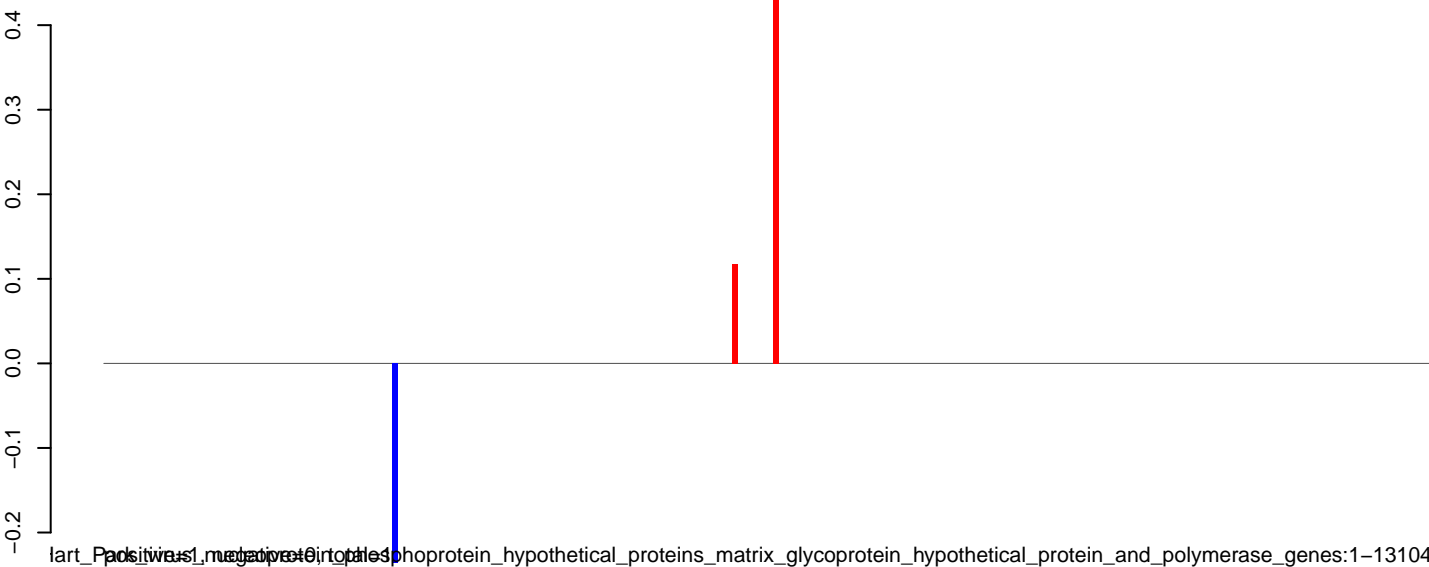
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



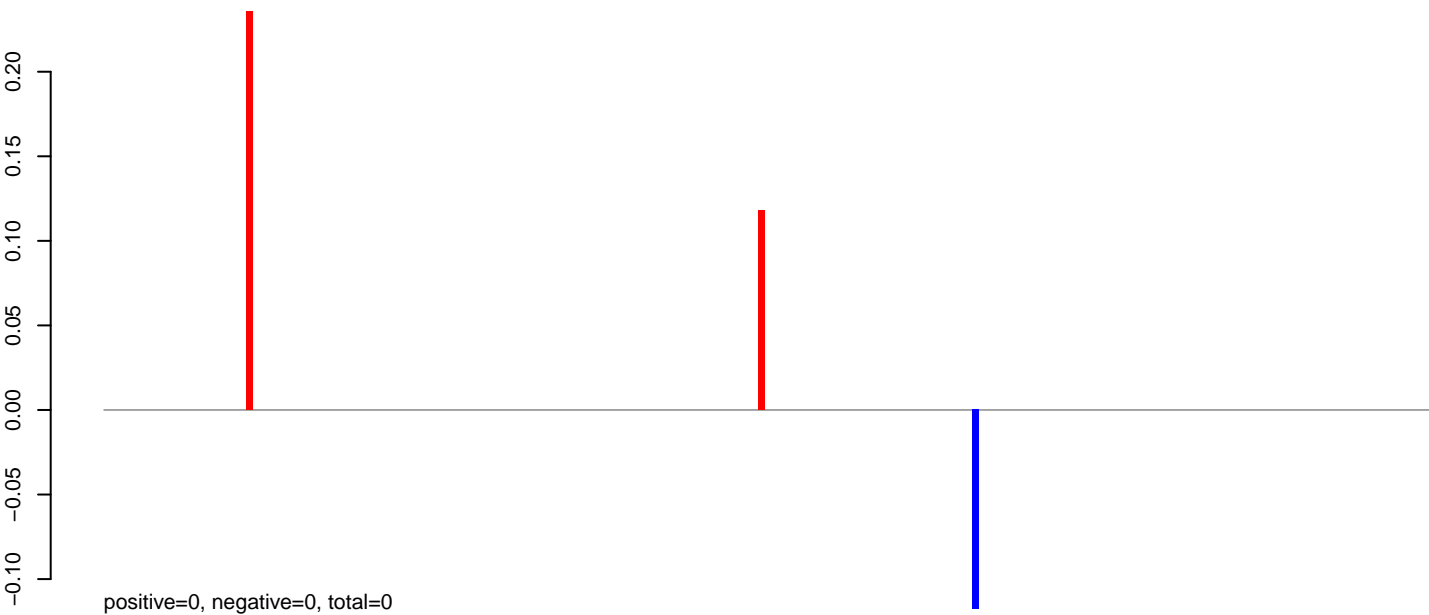
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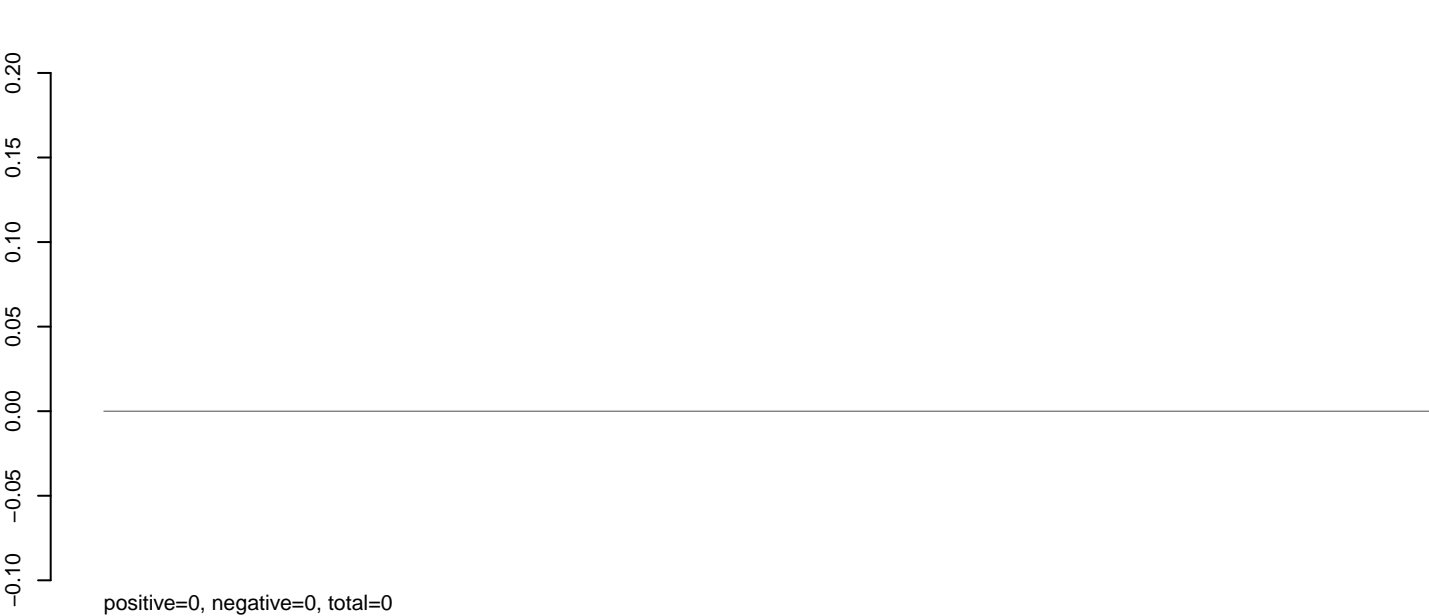
part_positives=1, negative=0, total=1 phosphoprotein_hypothetical_proteins_matrix_glycoprotein_hypothetical_protein_and_polymerase_genes:1-13104

0 2000 4000 6000 8000 10000 12000

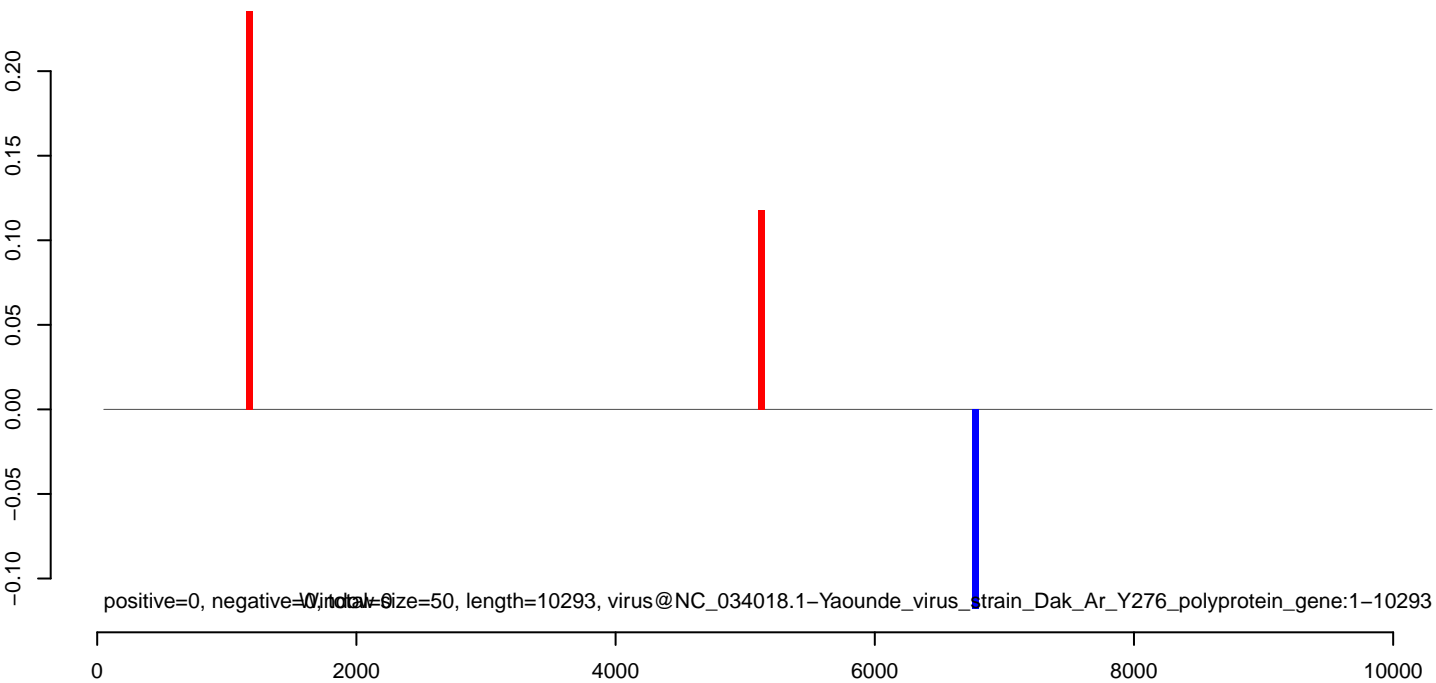
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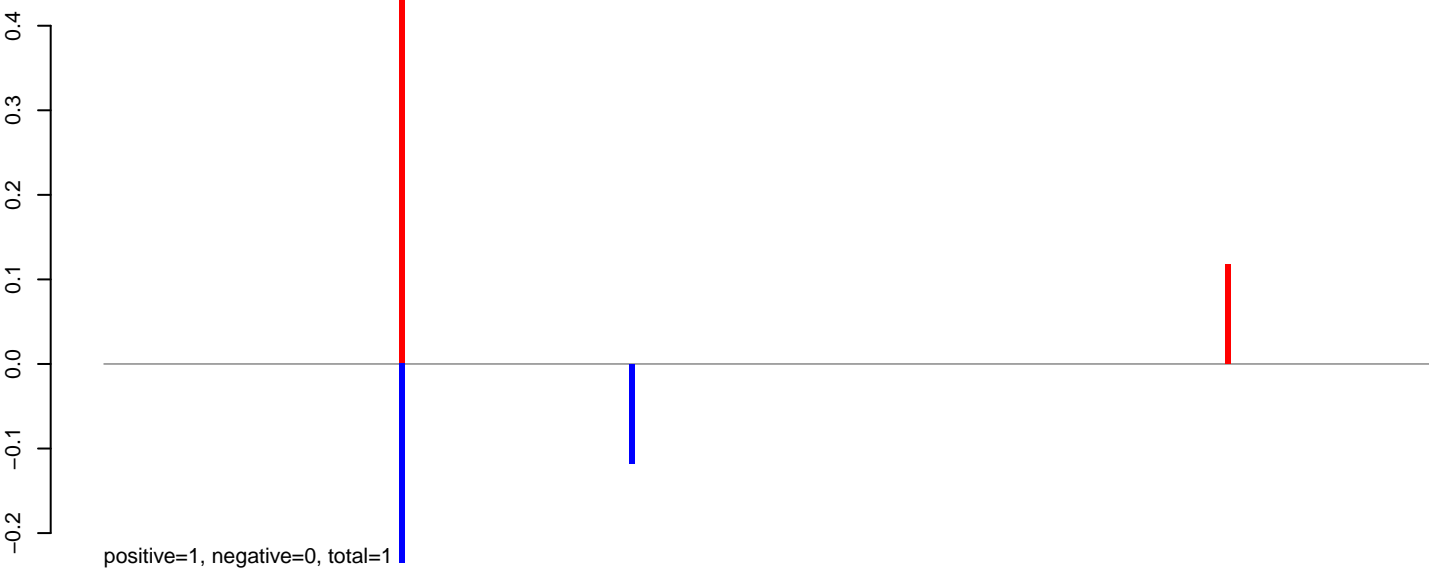
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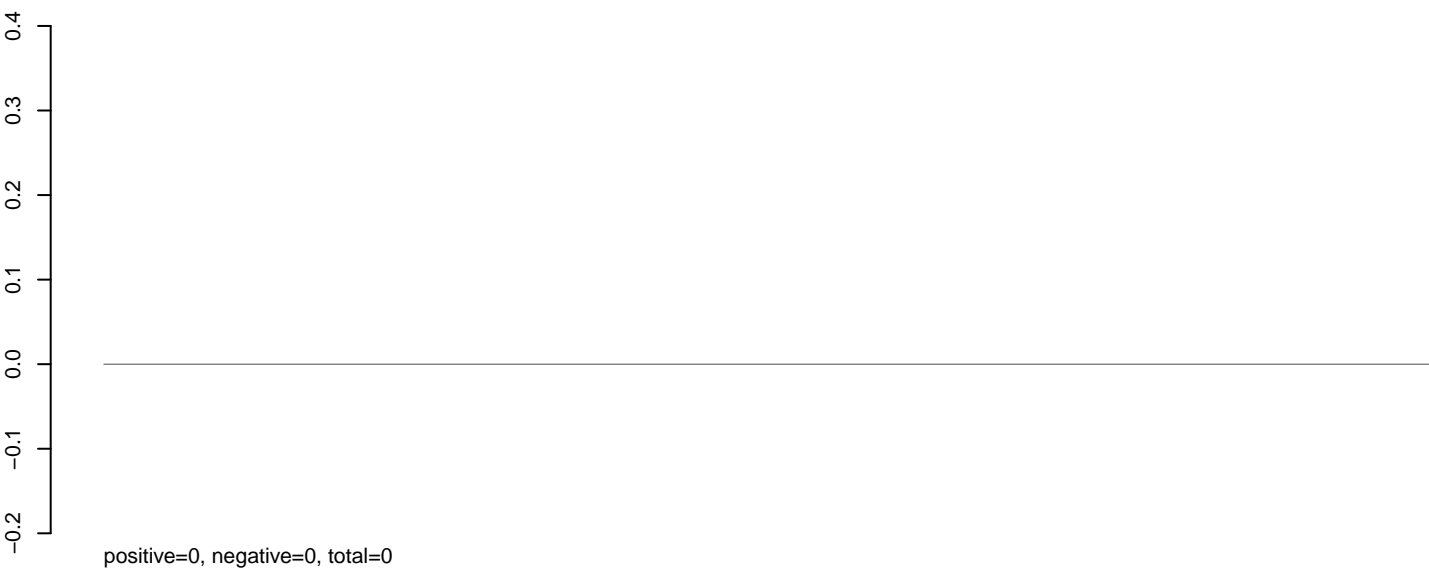
AeAeg_Aag2_DENV1.rep



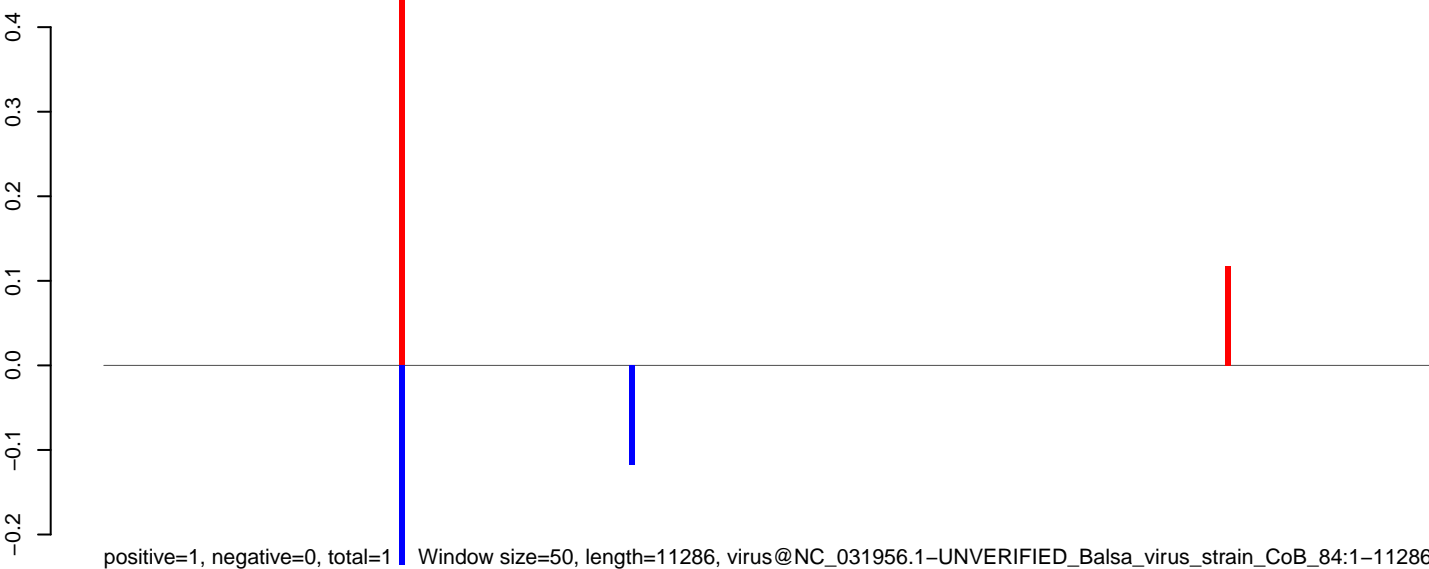
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



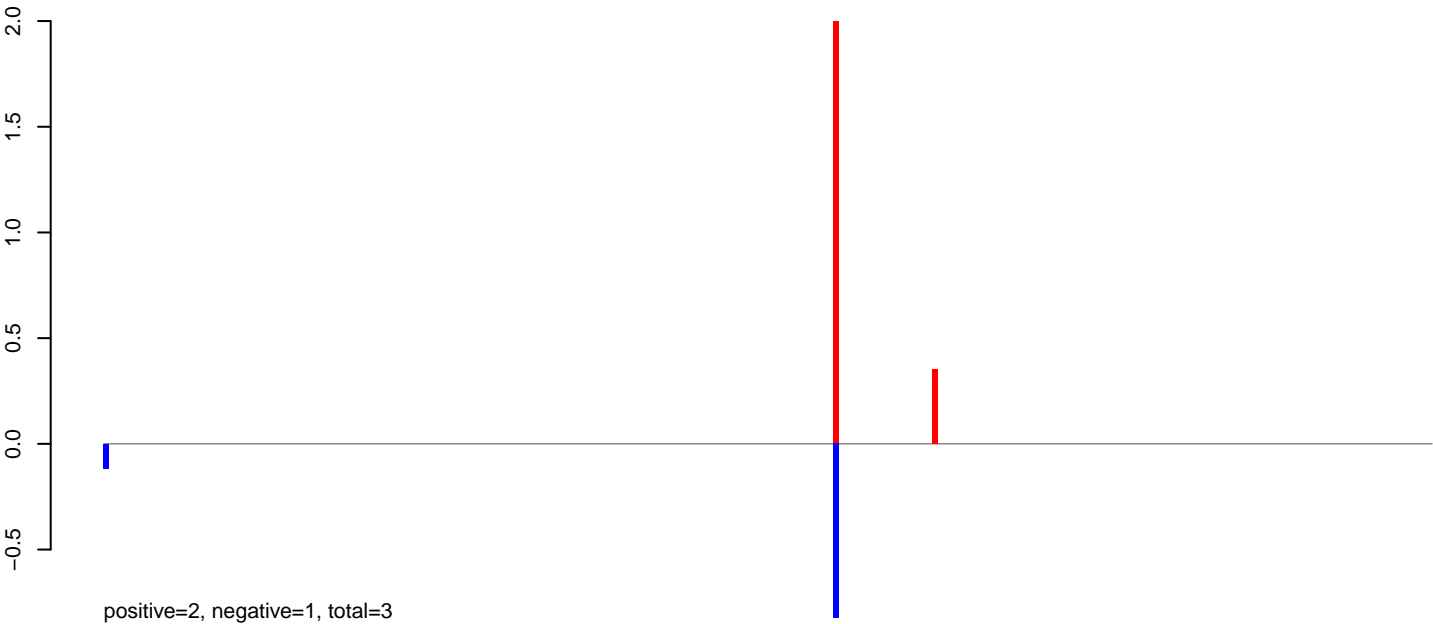
AeAeg_Aag2_DENV1.rep



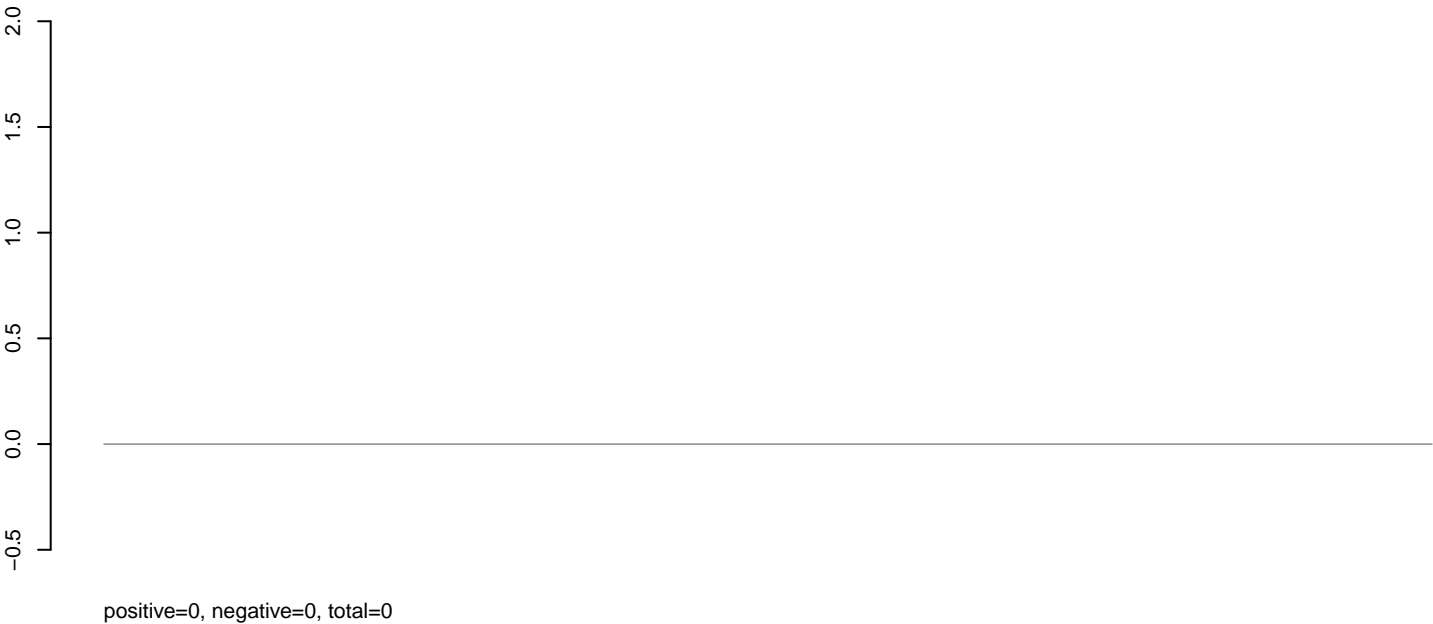
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0 2000 4000 6000 8000 10000

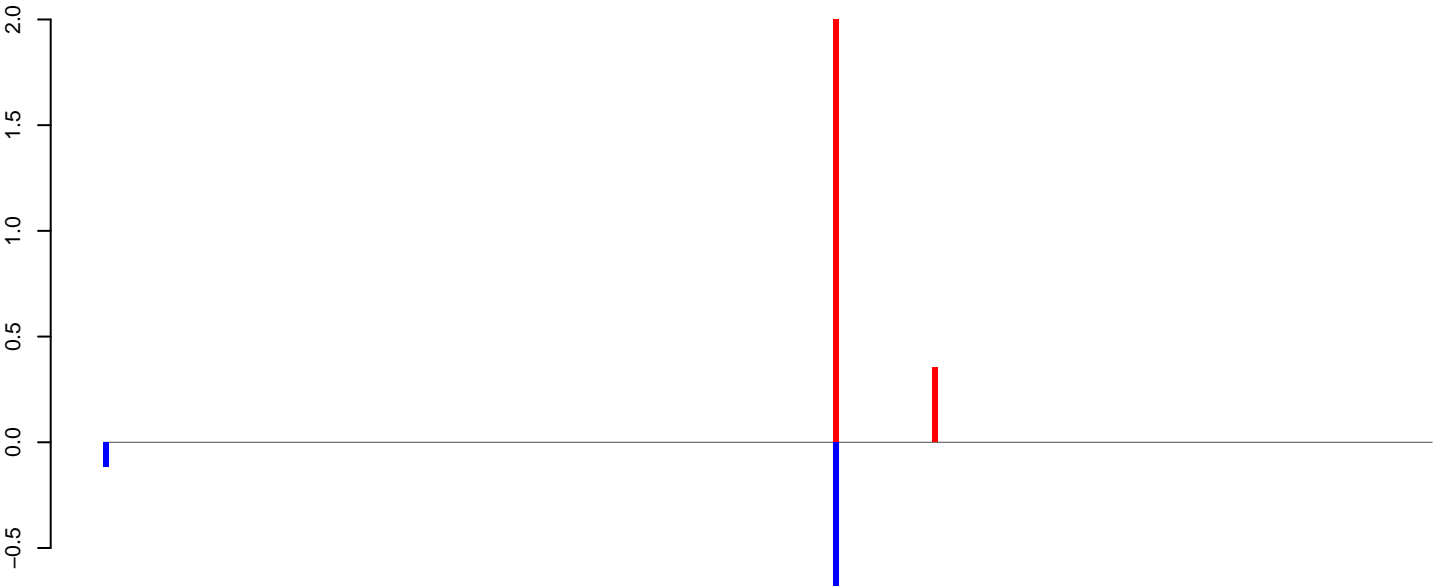
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



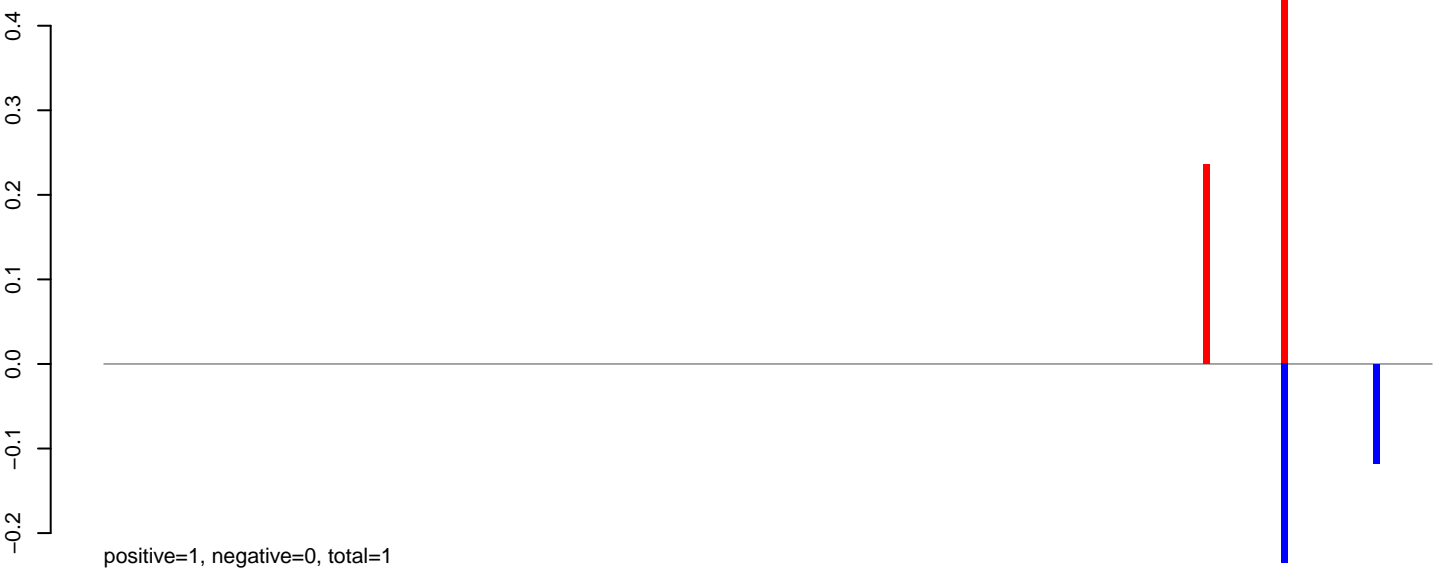
AeAeg_Aag2_DENV1.rep



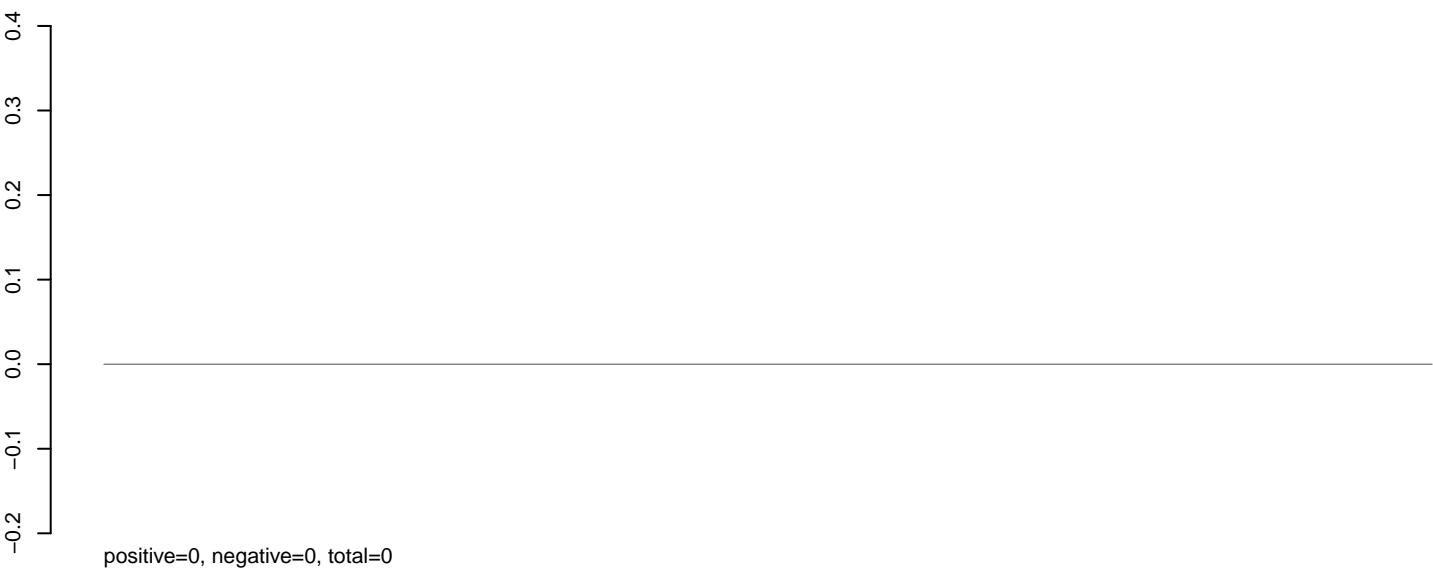
ncmp110seq21megaise1rainak31-6_ORF1_ORF1_ORF2_ORF2_glycoprotein_G_and_RNA-dependent_RNA_polymerase_L_genes:1-12774

0 2000 4000 6000 8000 10000 12000

AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



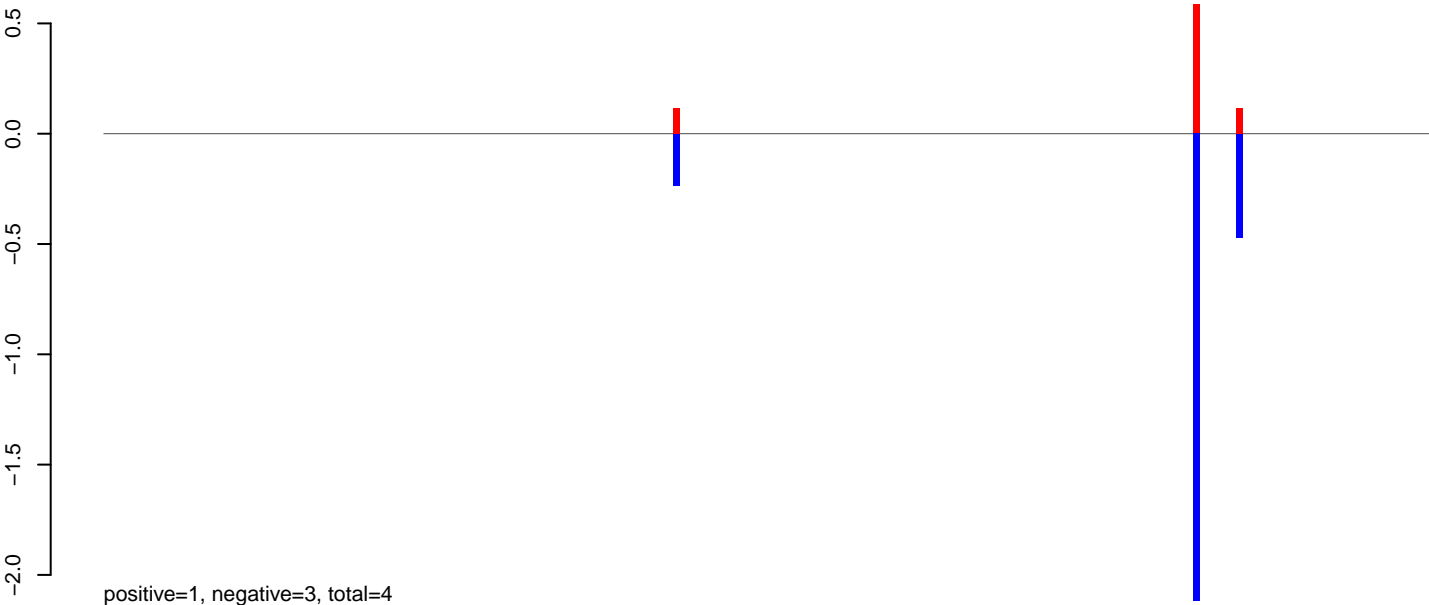
AeAeg_Aag2_DENV1.rep



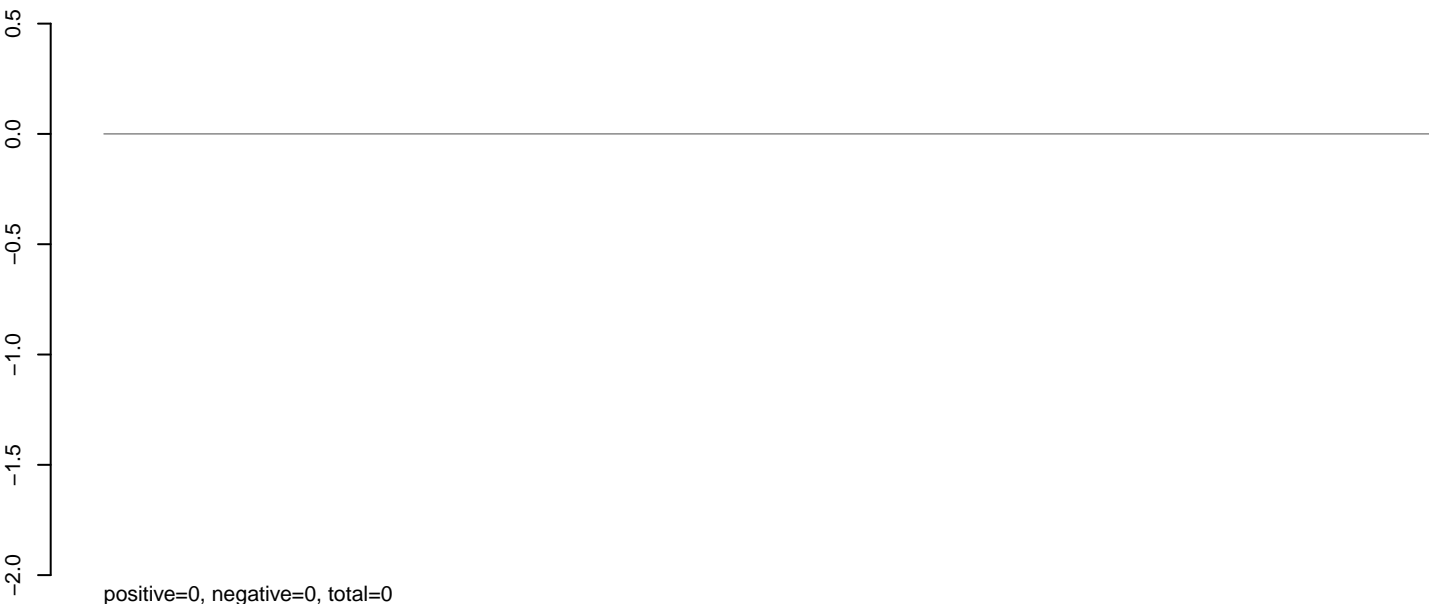
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0 2000 4000 6000 8000 10000

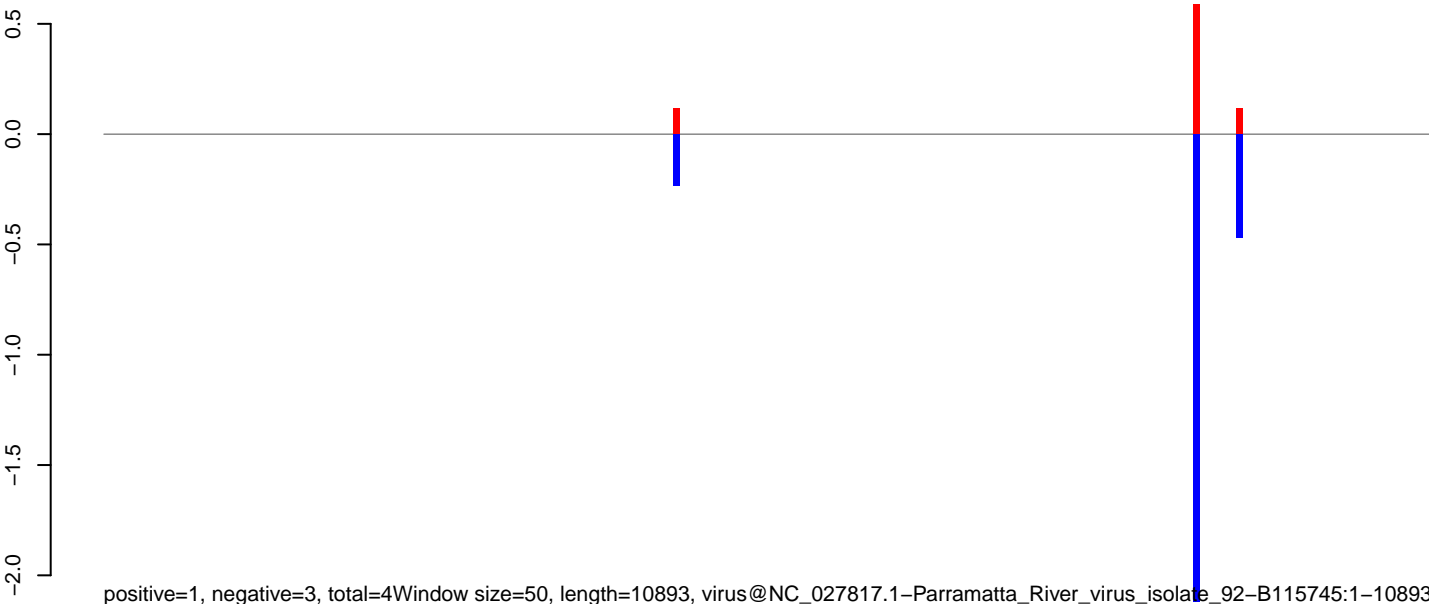
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



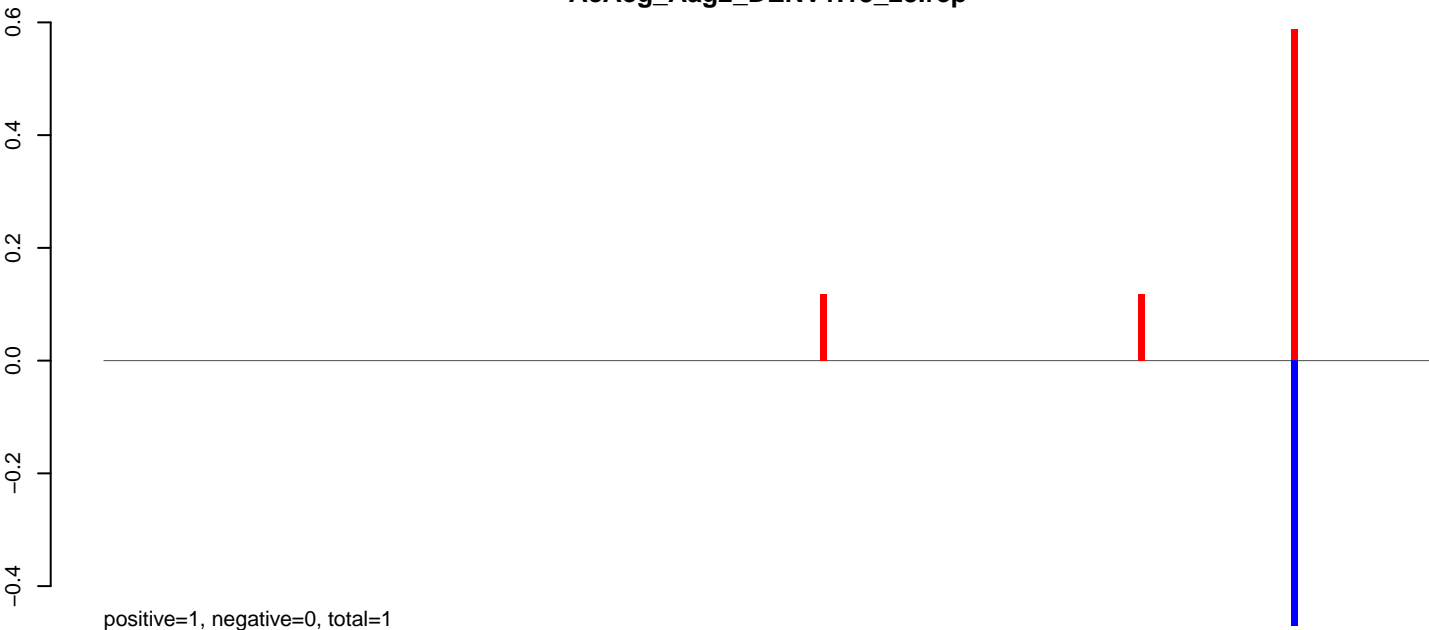
AeAeg_Aag2_DENV1.rep



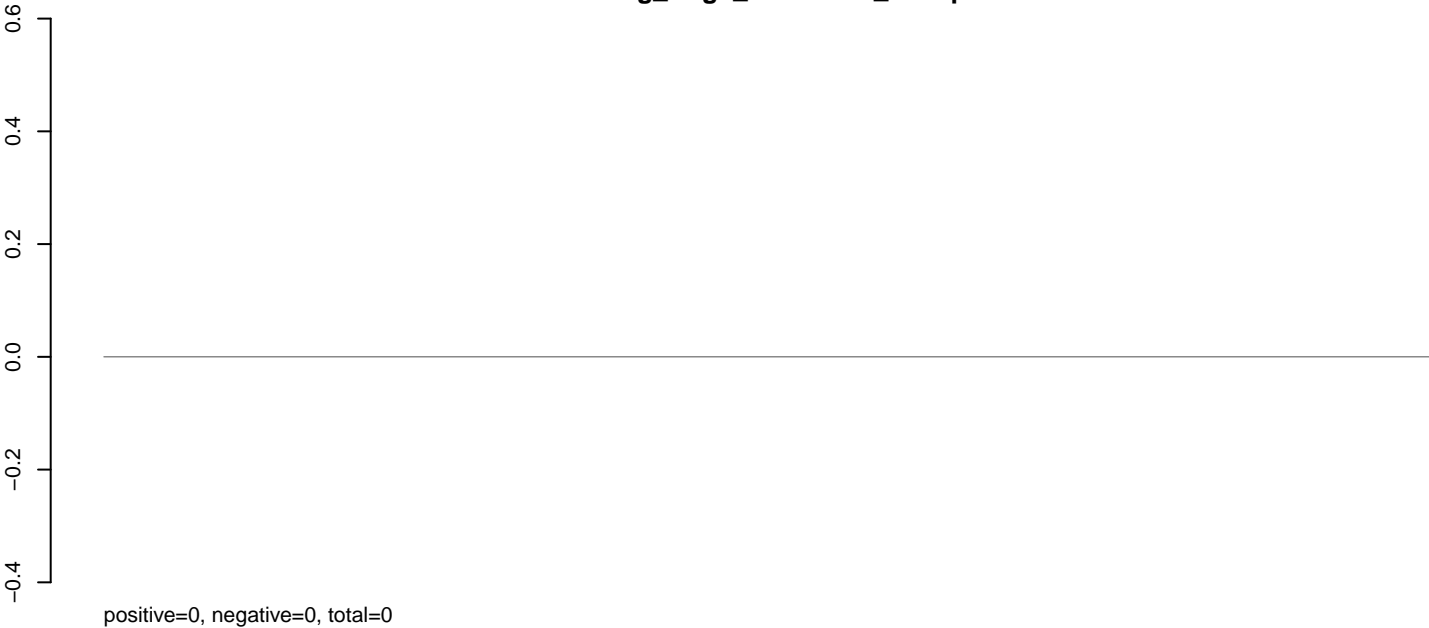
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0 2000 4000 6000 8000 10000

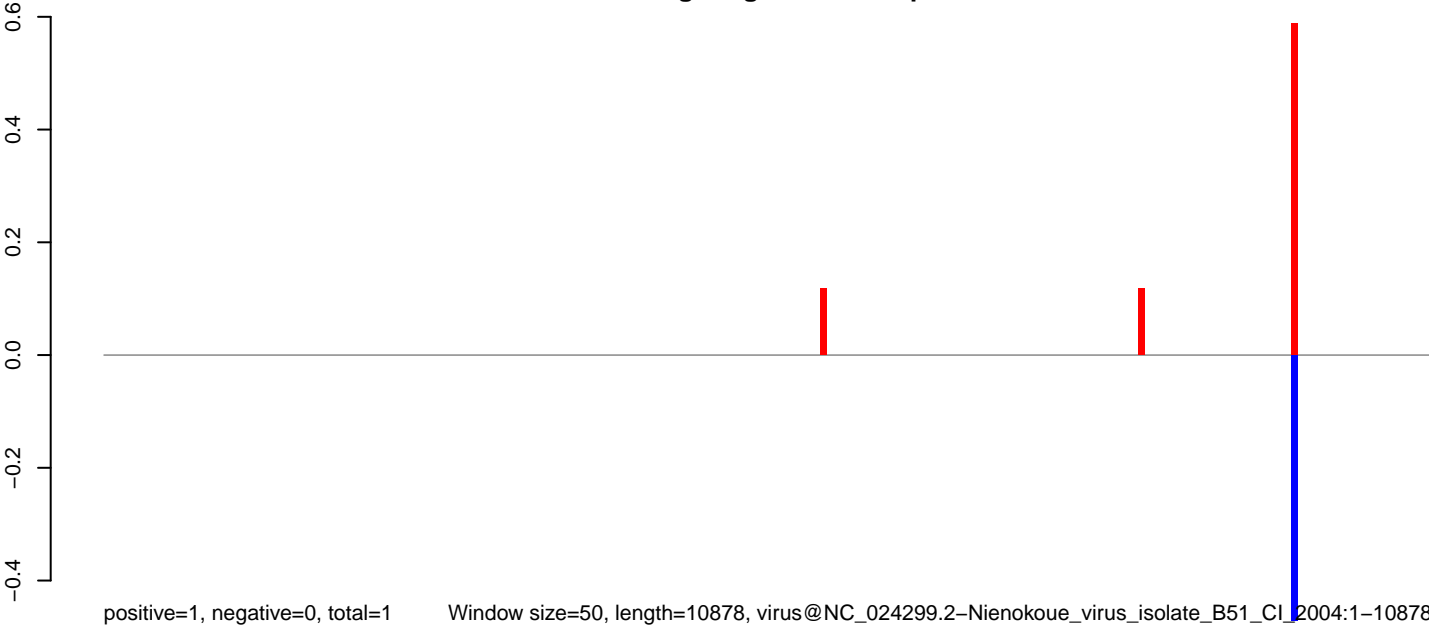
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



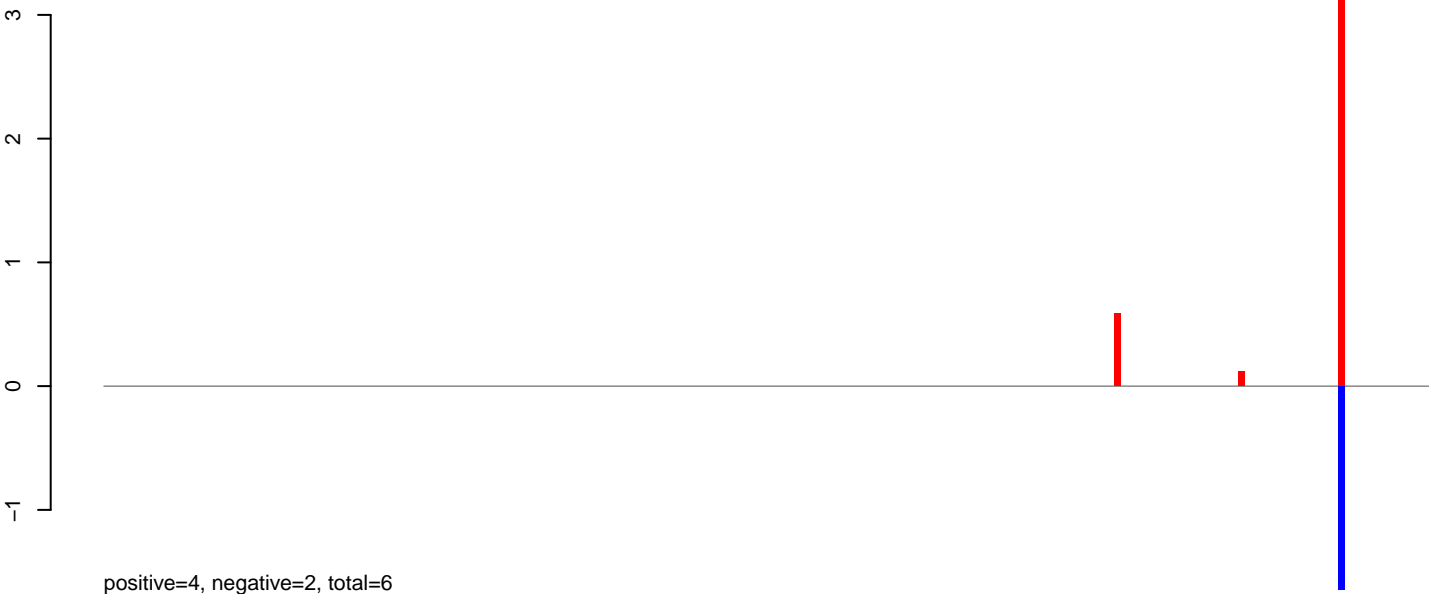
AeAeg_Aag2_DENV1.rep



Window size=50, length=10878, virus@NC_024299.2-Nienokoue_virus_isolate_B51_CI_2004:1-10878

0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV1.18_23.rep



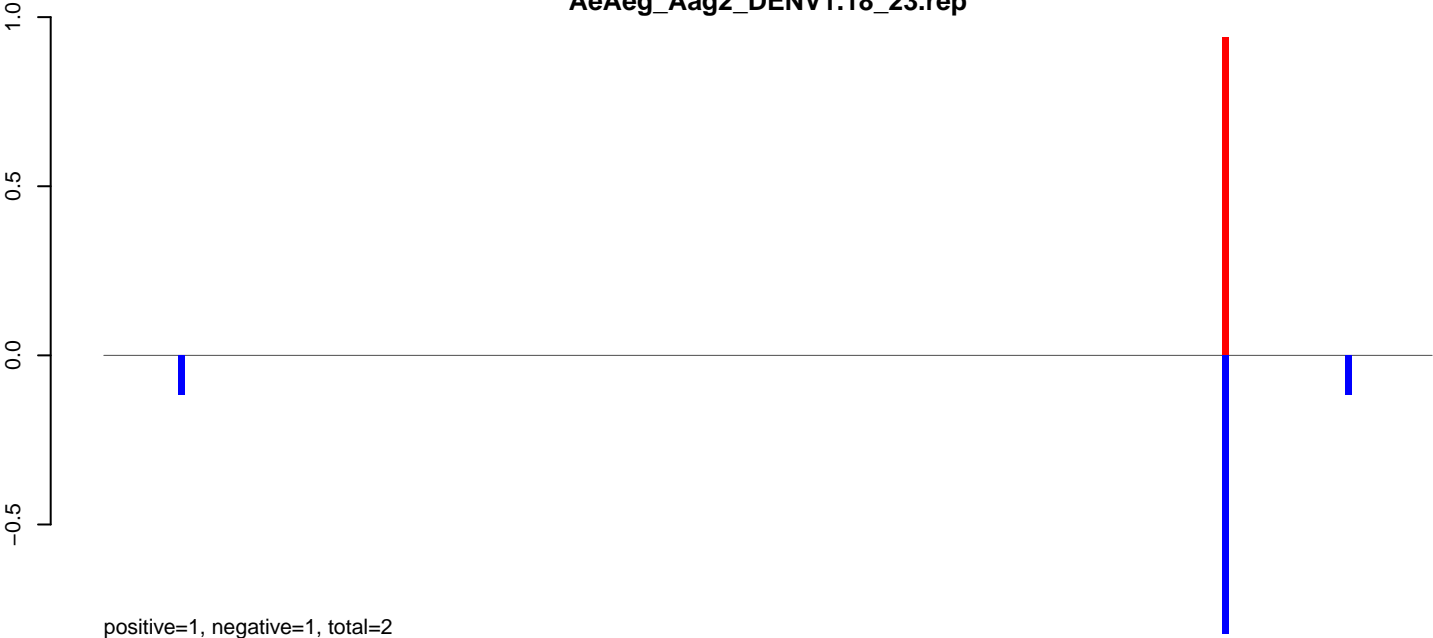
AeAeg_Aag2_DENV1.24_35.rep



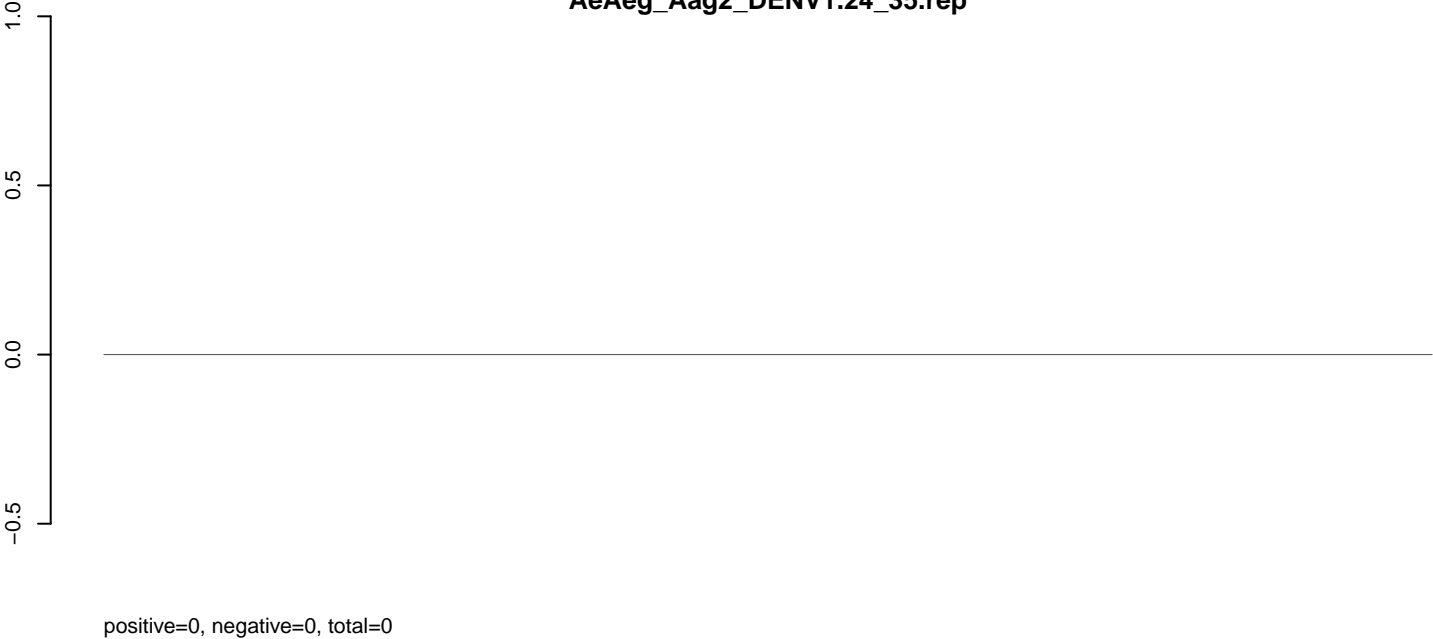
AeAeg_Aag2_DENV1.rep



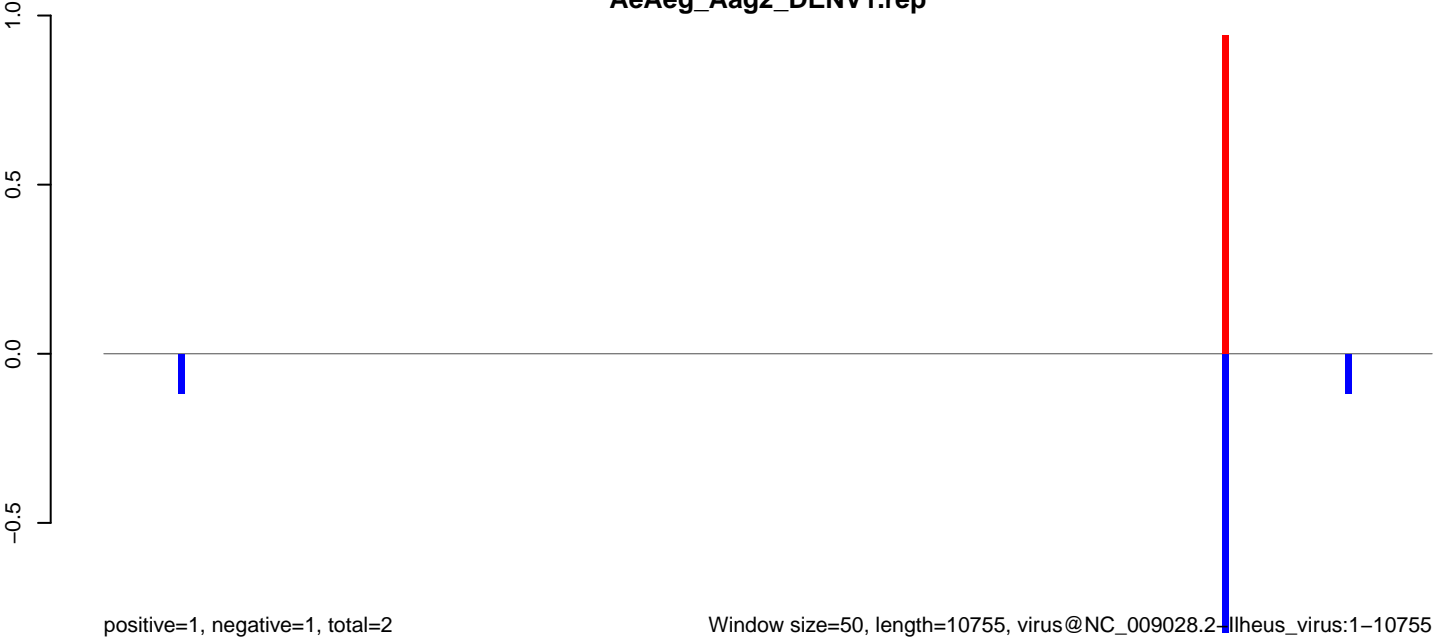
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



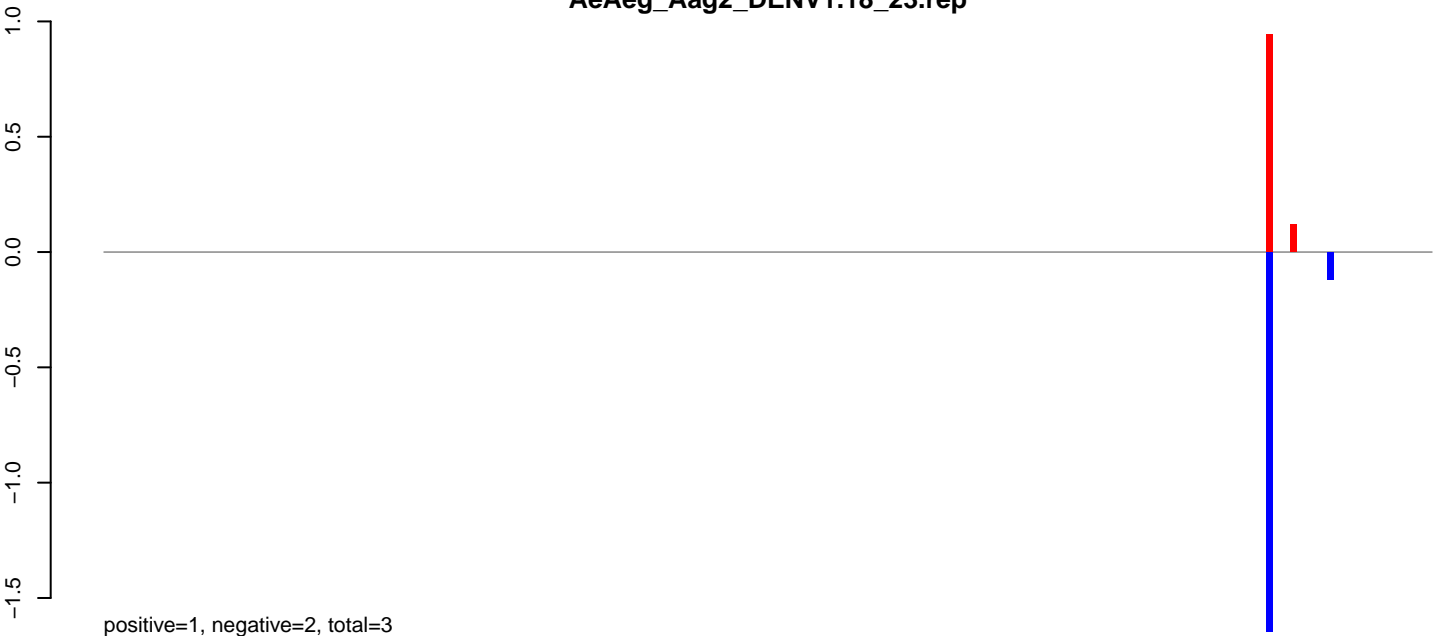
AeAeg_Aag2_DENV1.rep



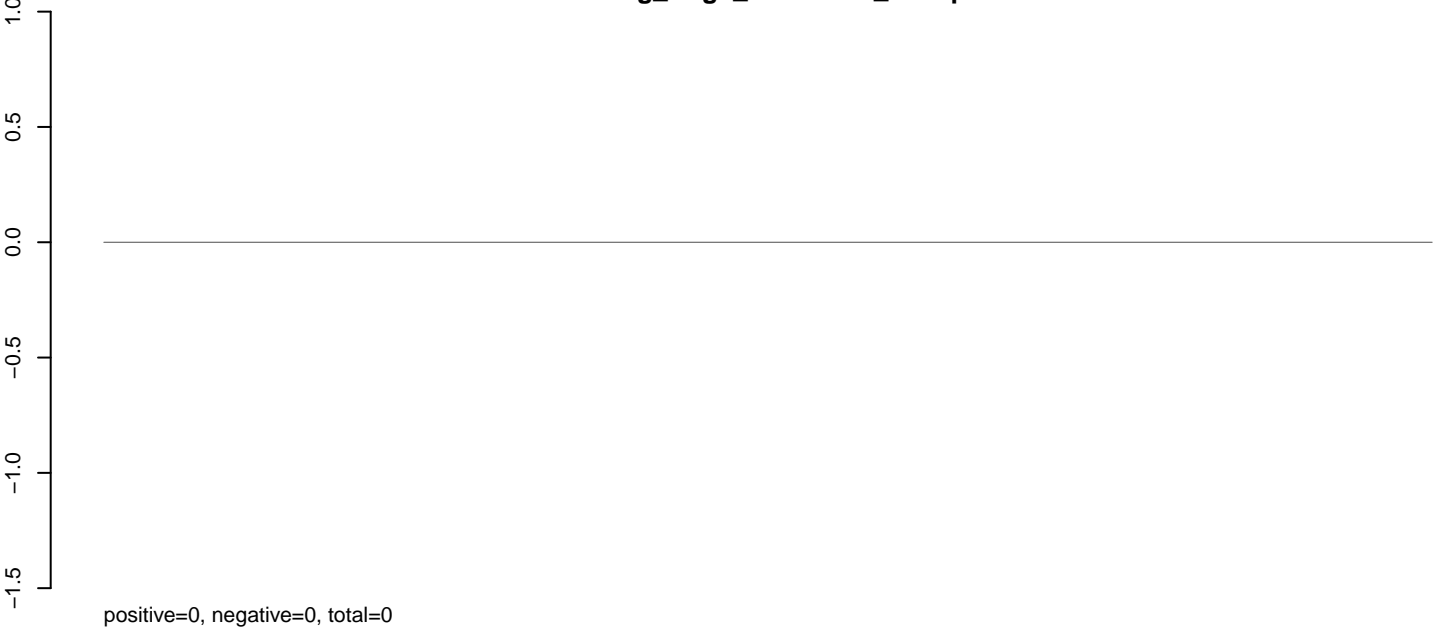
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0 2000 4000 6000 8000 10000

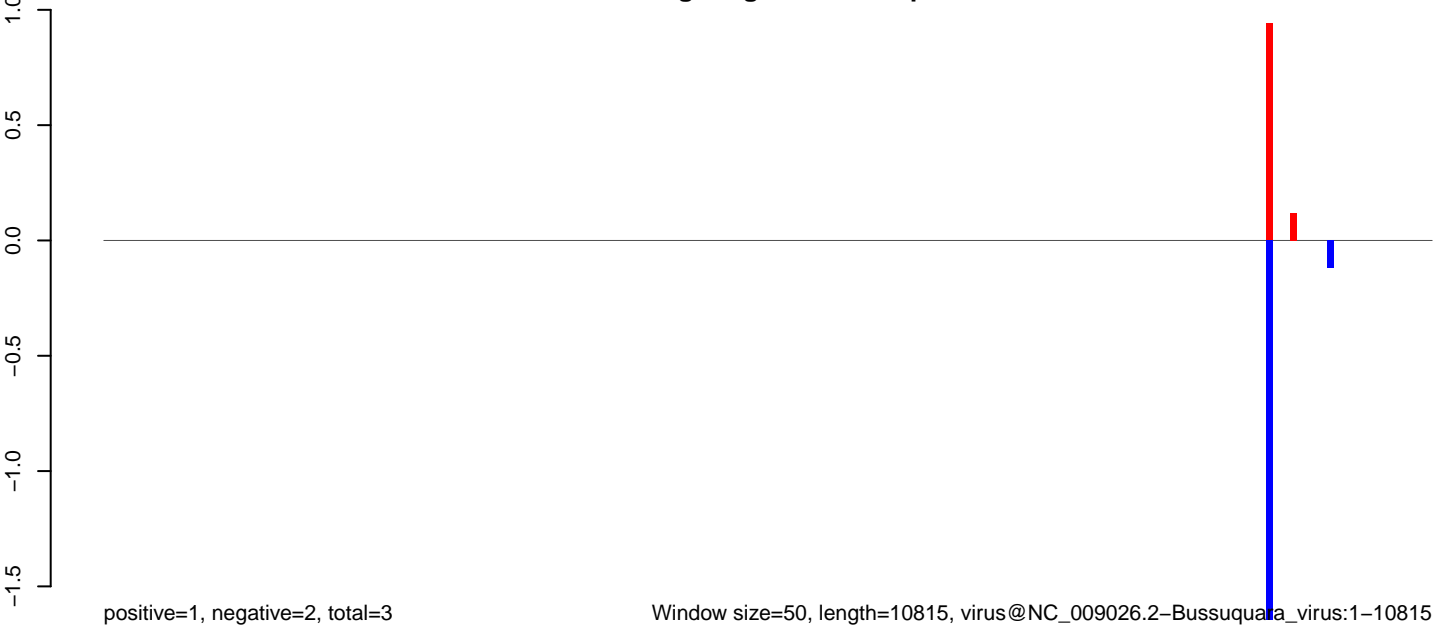
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



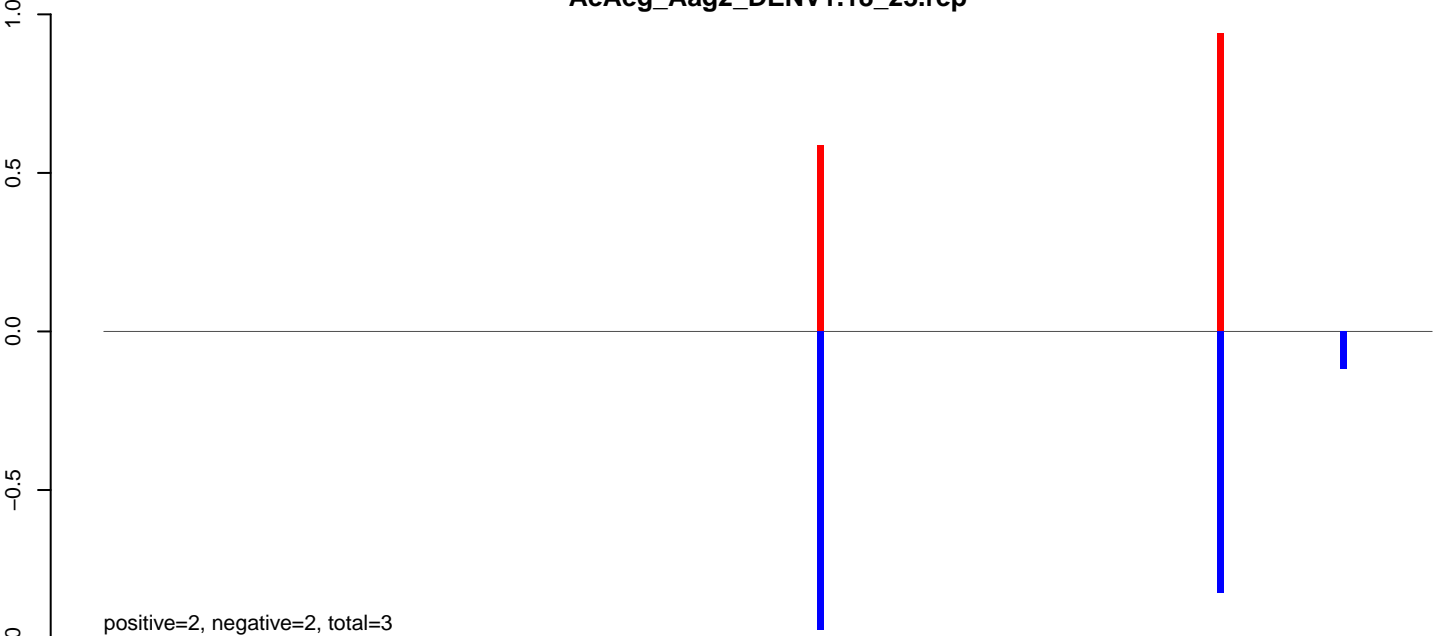
AeAeg_Aag2_DENV1.rep



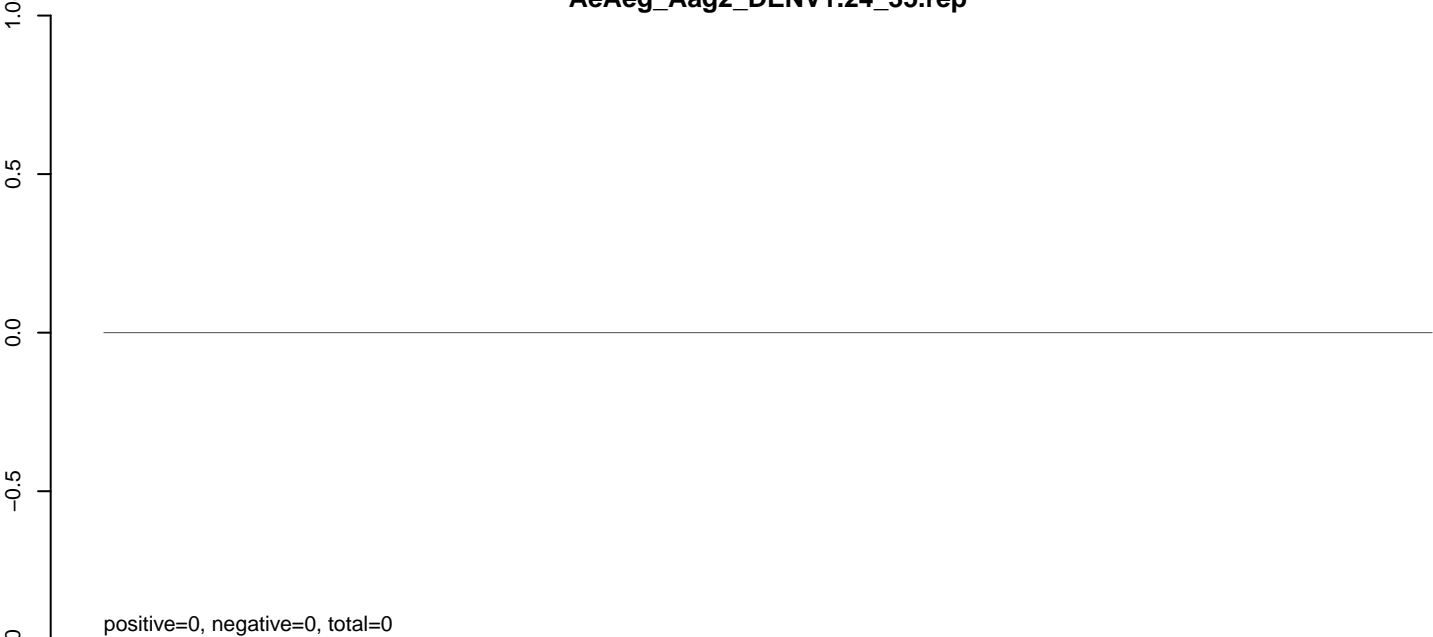
Window size=50, length=10815, virus@NC_009026.2-Bussuquara_virus:1-10815

0 2000 4000 6000 8000 10000

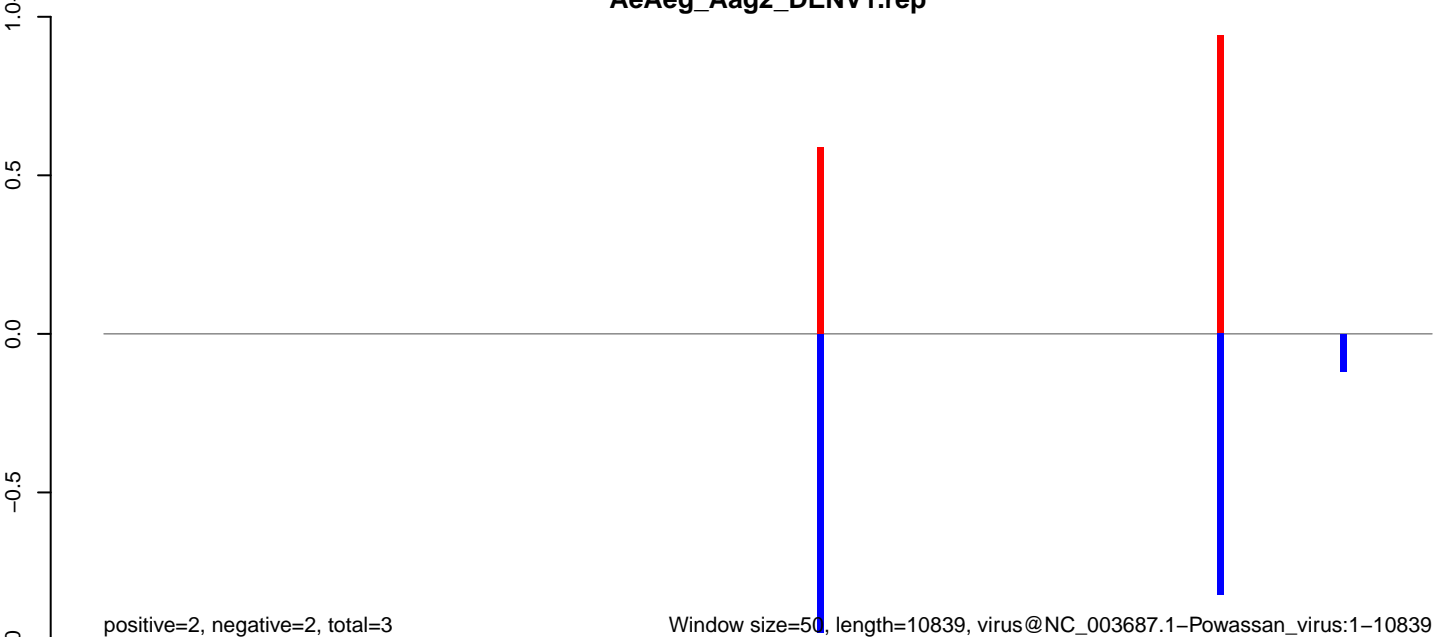
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

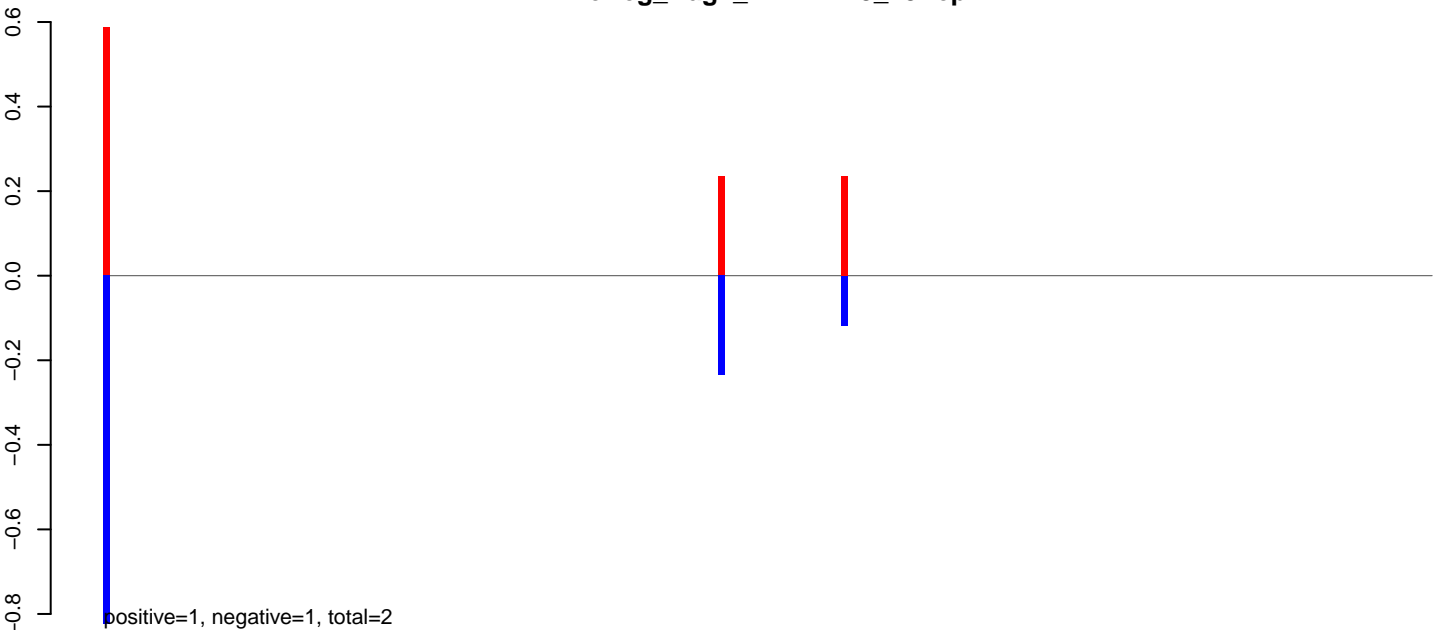


AeAeg_Aag2_DENV1.rep

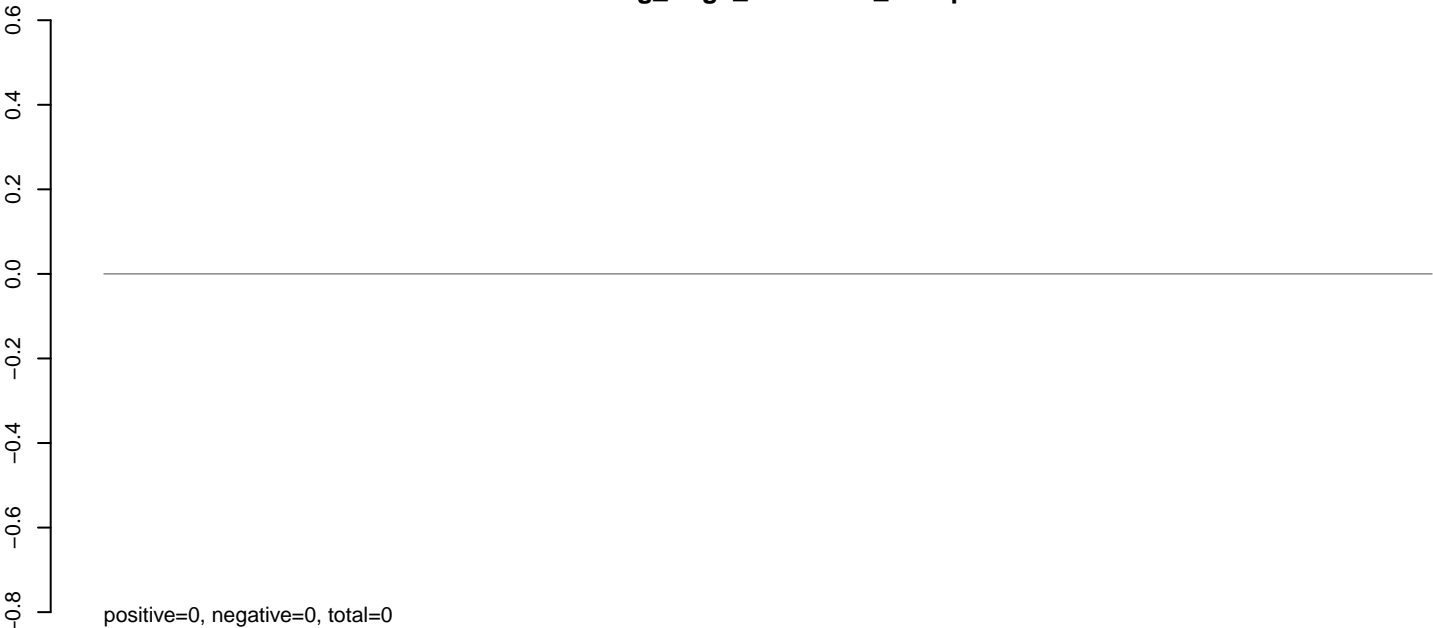


Window size=50, length=10839, virus@NC_003687.1-Powassan_virus:1-10839

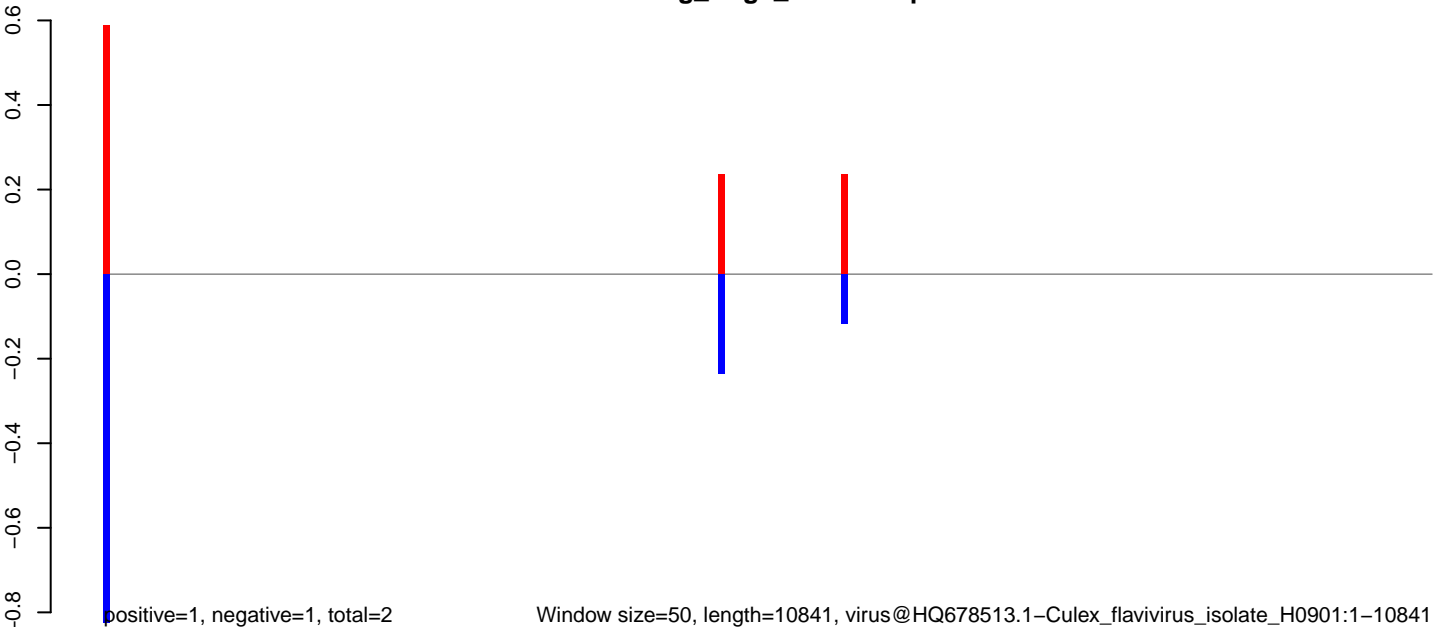
AeAeg_Aag2_DENV1.18_23.rep



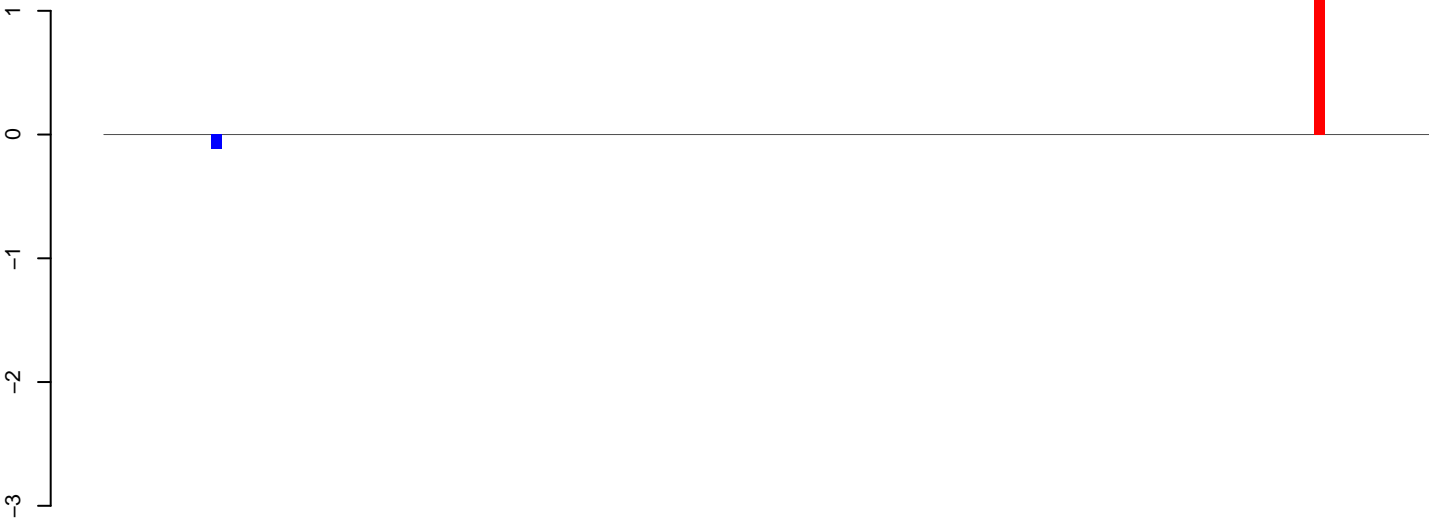
AeAeg_Aag2_DENV1.24_35.rep



AeAeg_Aag2_DENV1.rep

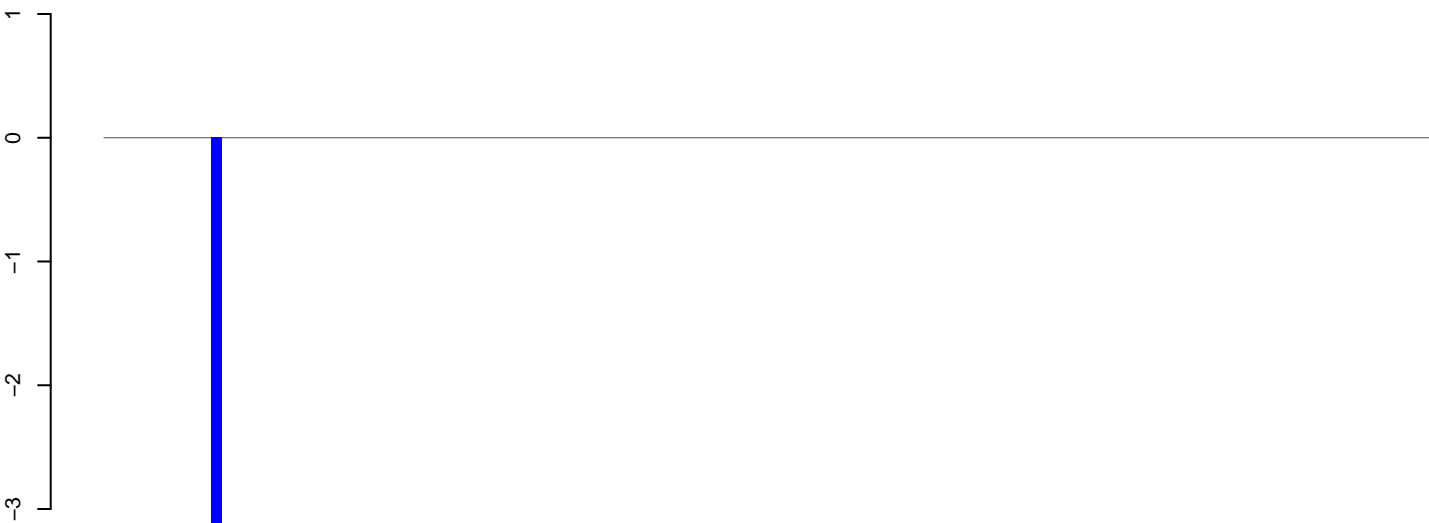


AeAeg_Aag2_DENV1.18_23.rep



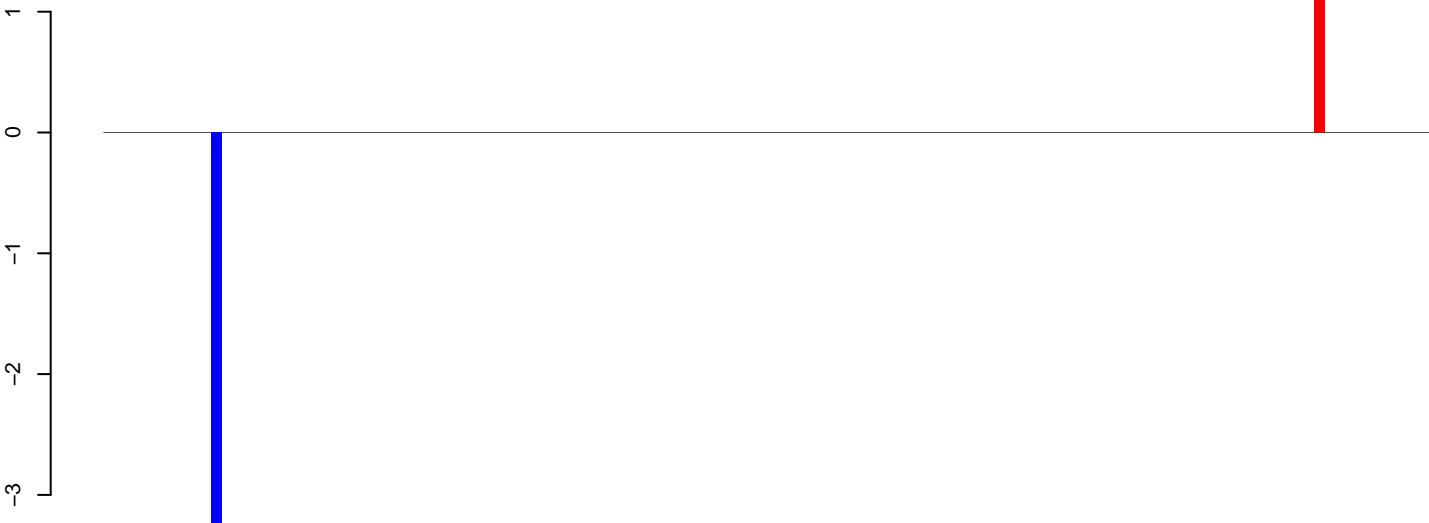
positive=1, negative=0, total=1

AeAeg_Aag2_DENV1.24_35.rep

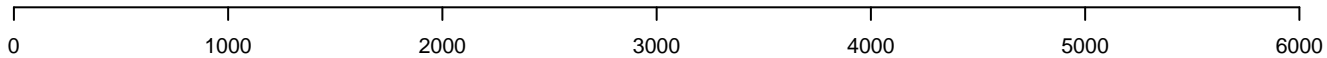


positive=0, negative=4, total=4

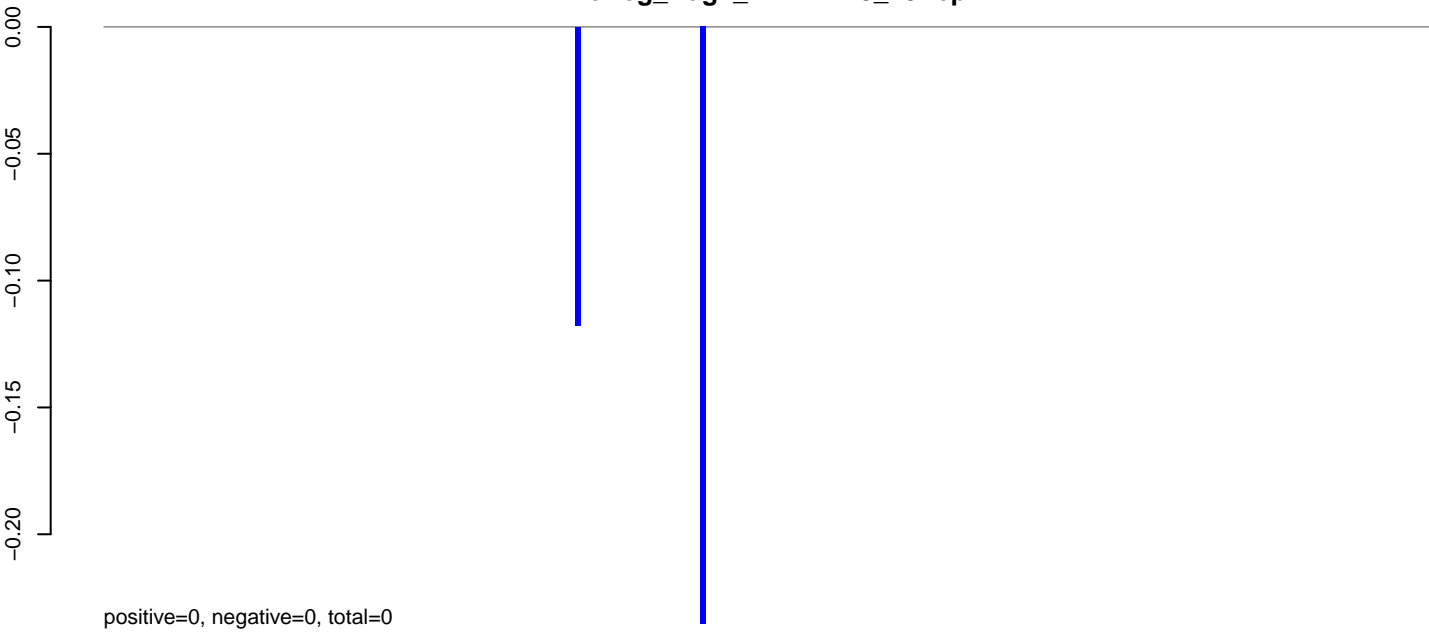
AeAeg_Aag2_DENV1.rep



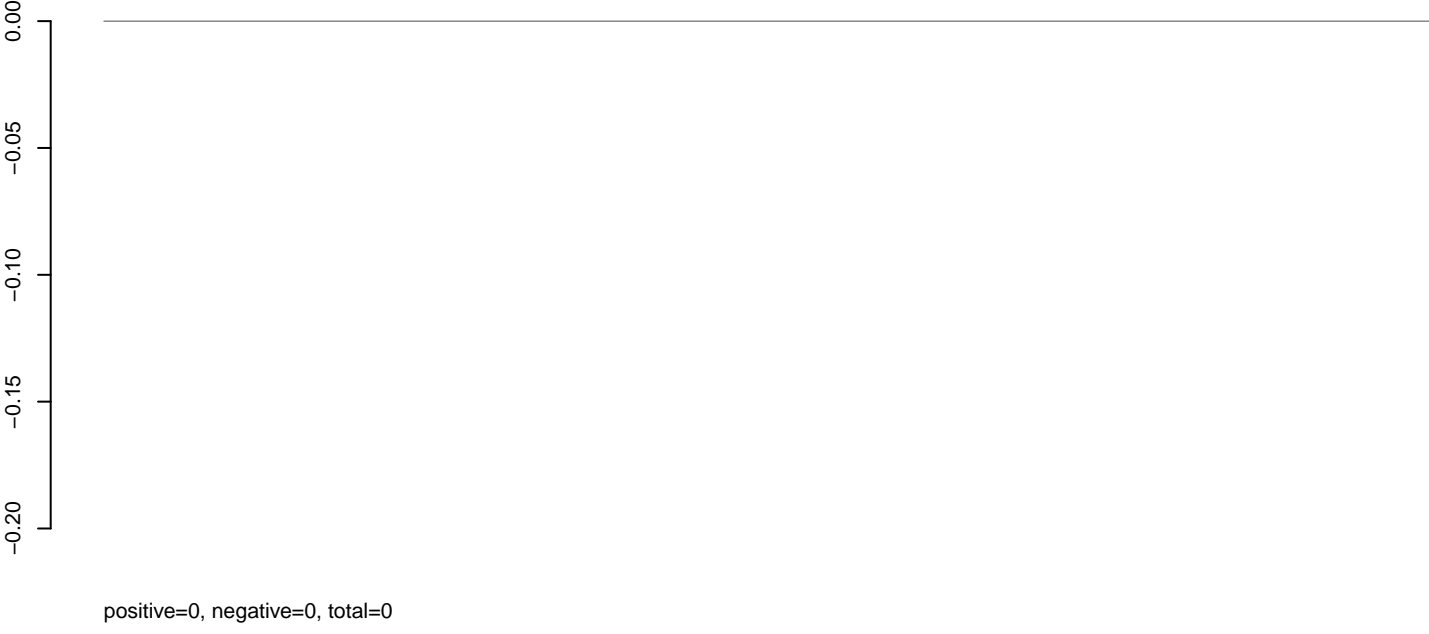
positive=1, negative=4, total=5, window size=50, length=6203, virus@NC_035674.1-Australian_Anopheles_totivirus_isolate_AATV_150840:1-6203



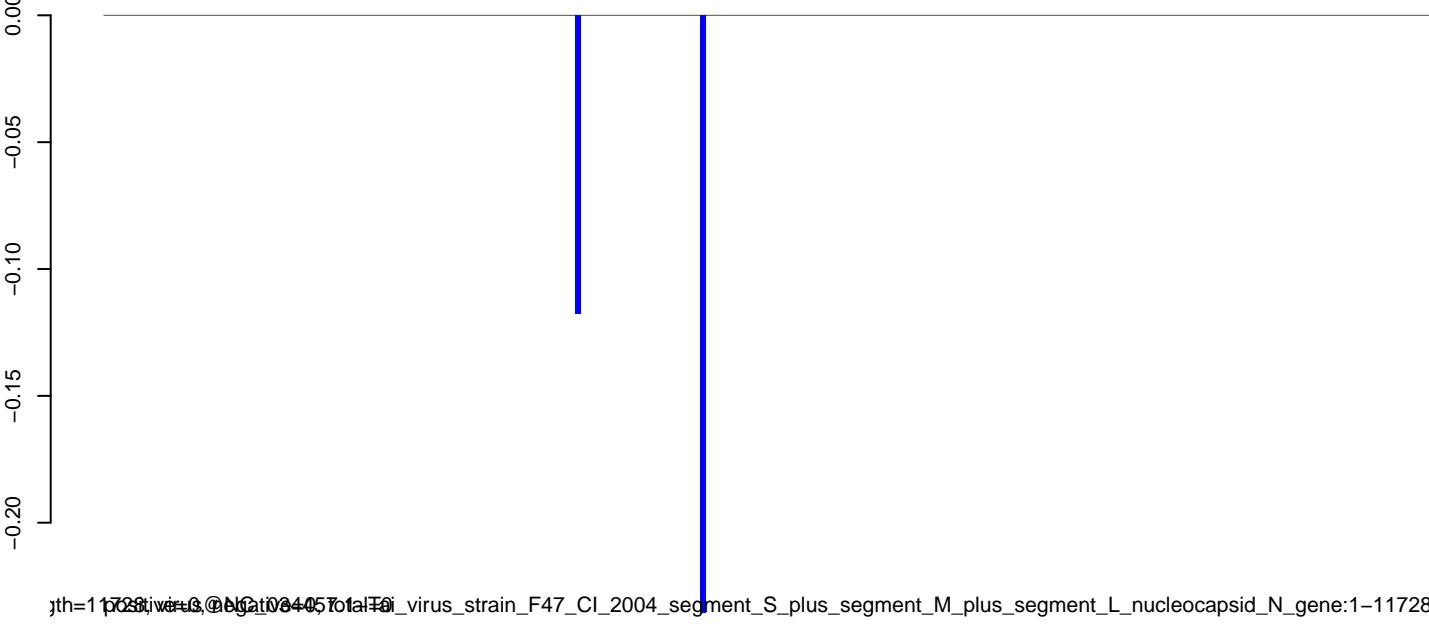
AeAeg_Aag2_DENV1.18_23.rep



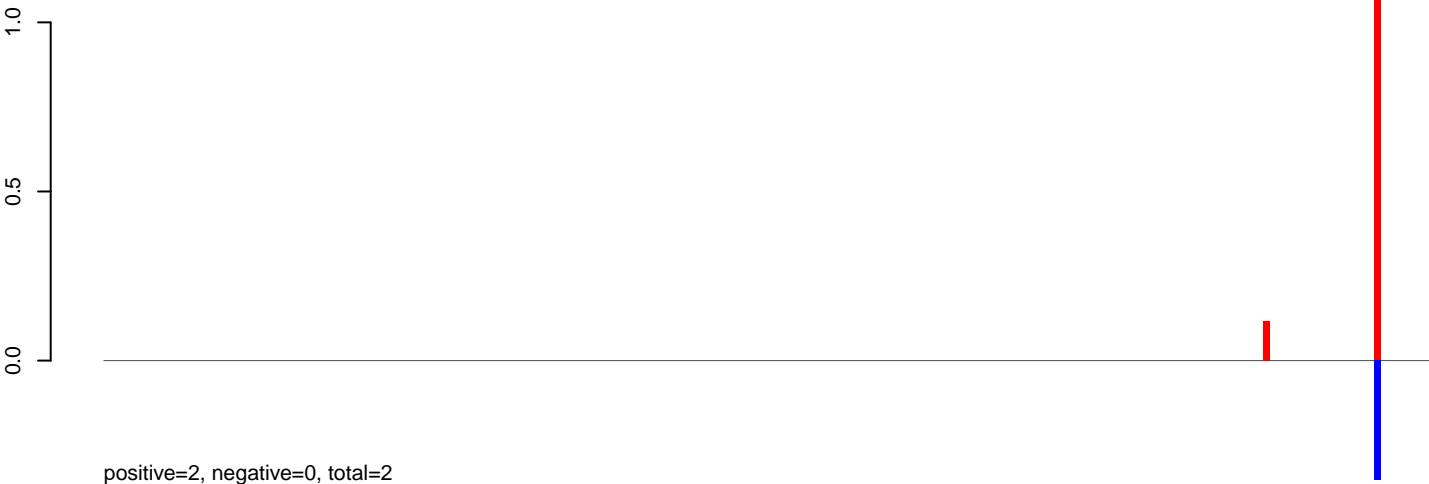
AeAeg_Aag2_DENV1.24_35.rep



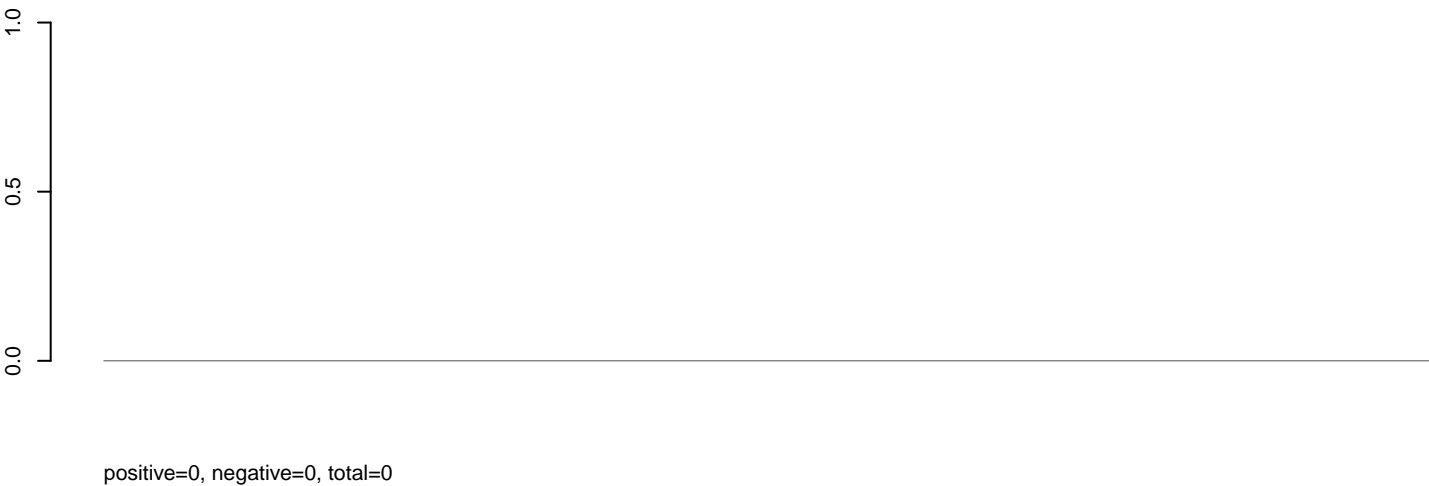
AeAeg_Aag2_DENV1.rep



AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



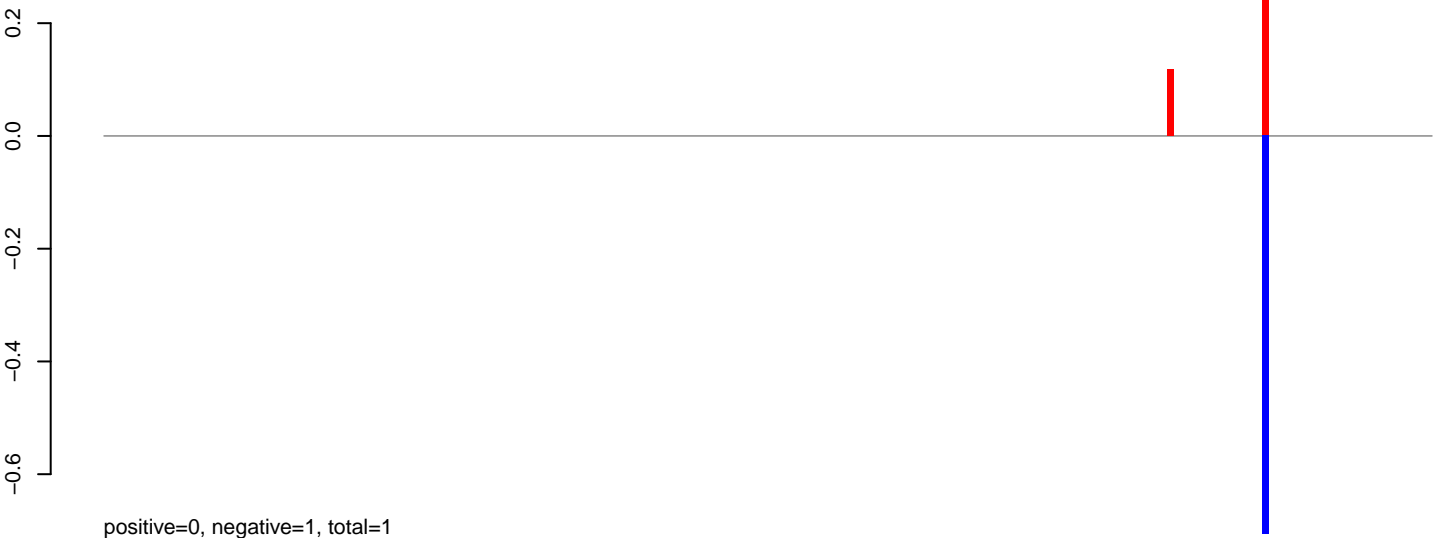
AeAeg_Aag2_DENV1.rep



Window size=50, length=10251, virus@NC_033723.1-Gadgets_Gully_virus_from_Australia_polyprotein_gene:1-10251

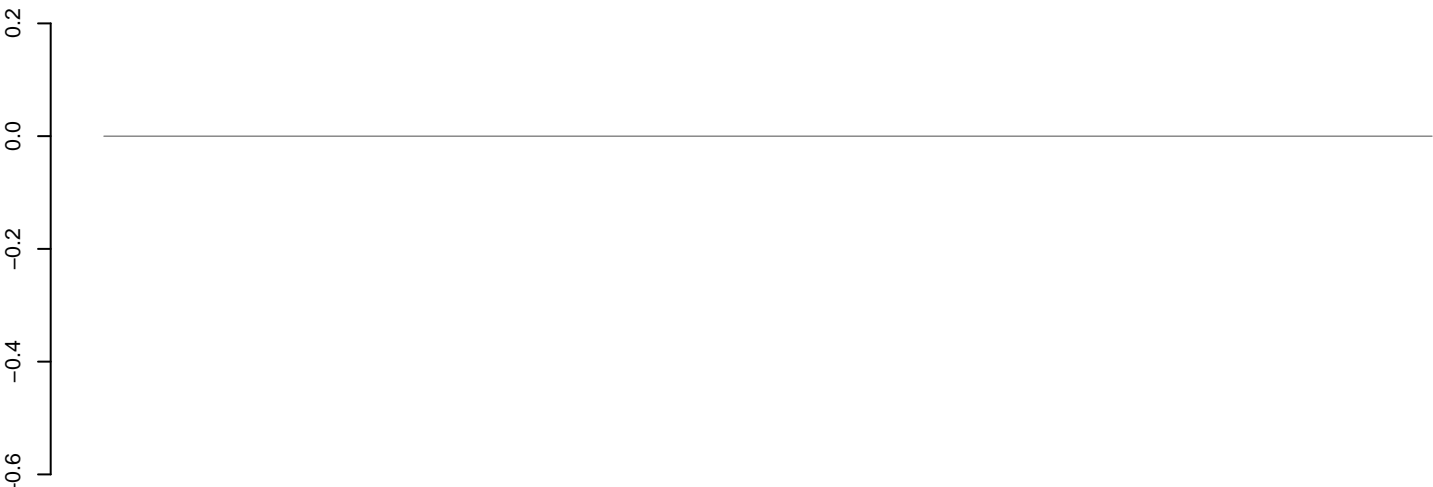
0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV1.18_23.rep



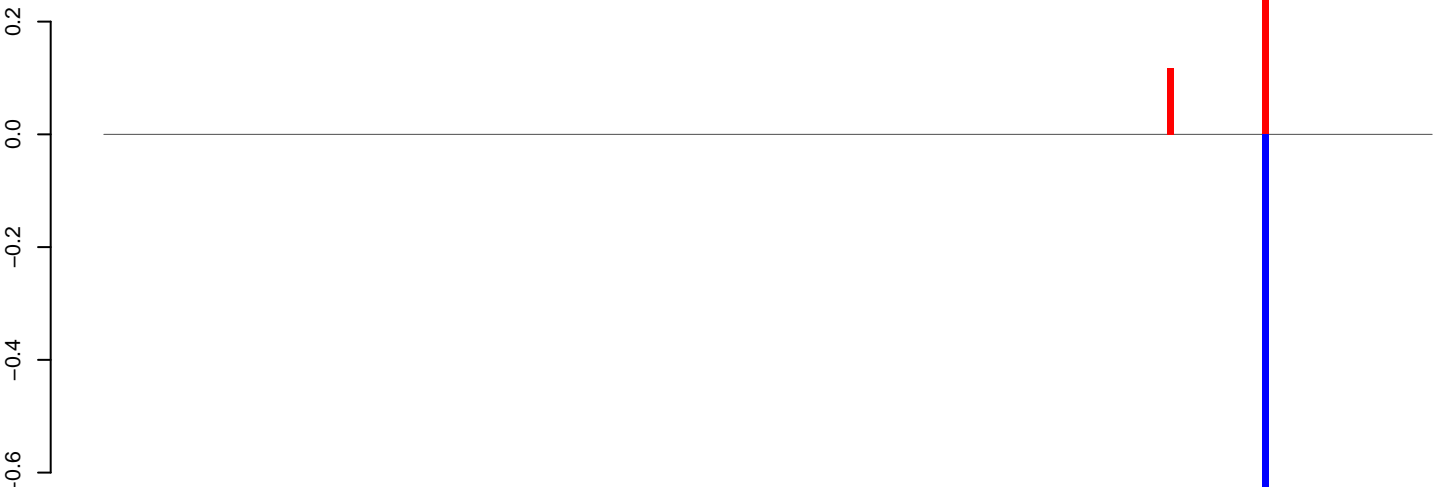
positive=0, negative=1, total=1

AeAeg_Aag2_DENV1.24_35.rep



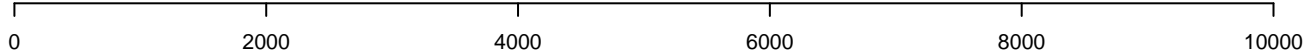
positive=0, negative=0, total=0

AeAeg_Aag2_DENV1.rep

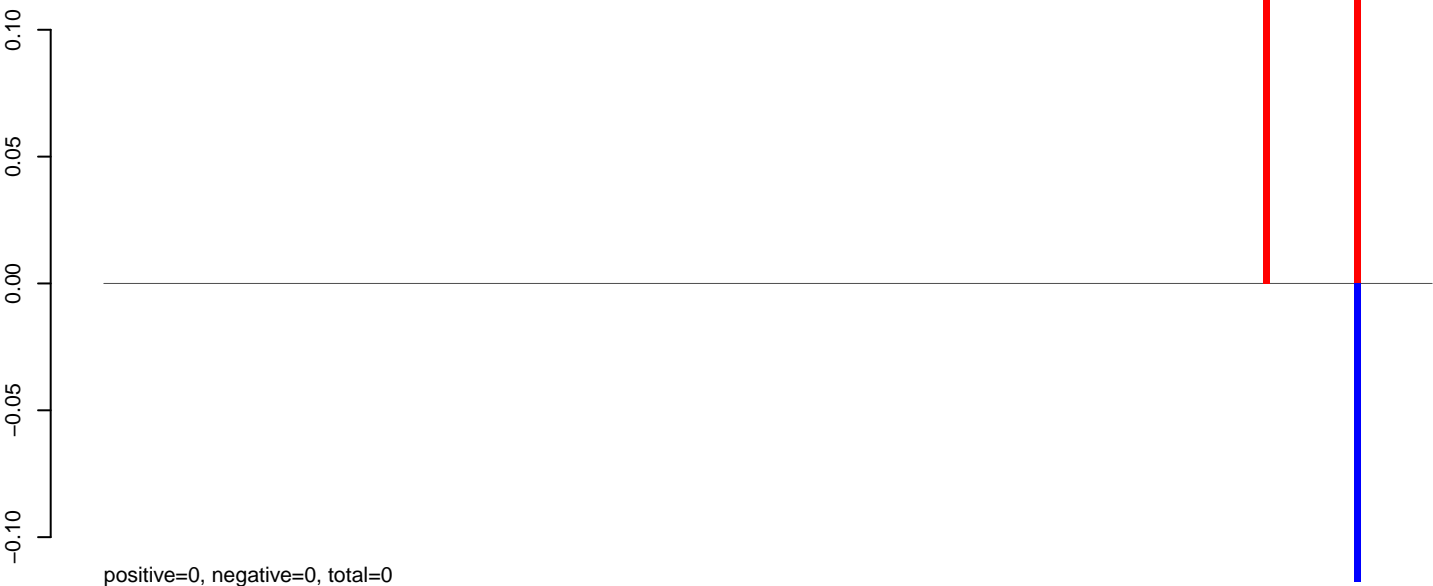


positive=0, negative=1, total=1

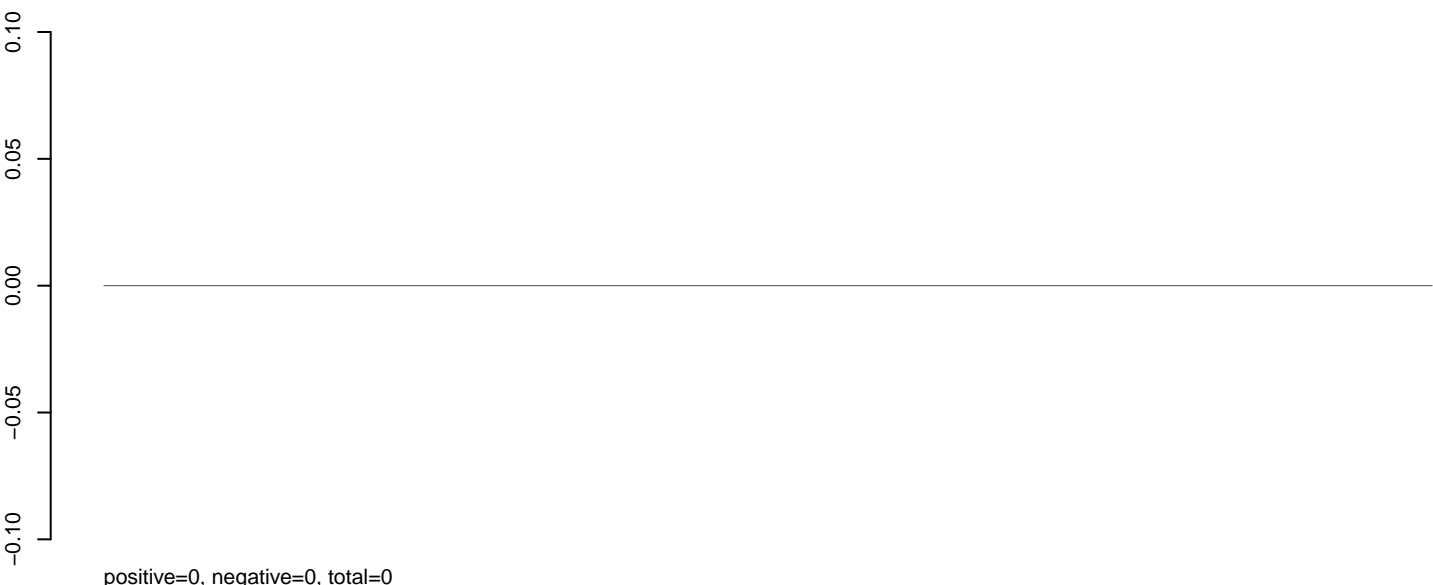
Window size=50, length=10588, virus@NC_031327.1-Anopheles_flavivirus_variant_1:1-10588



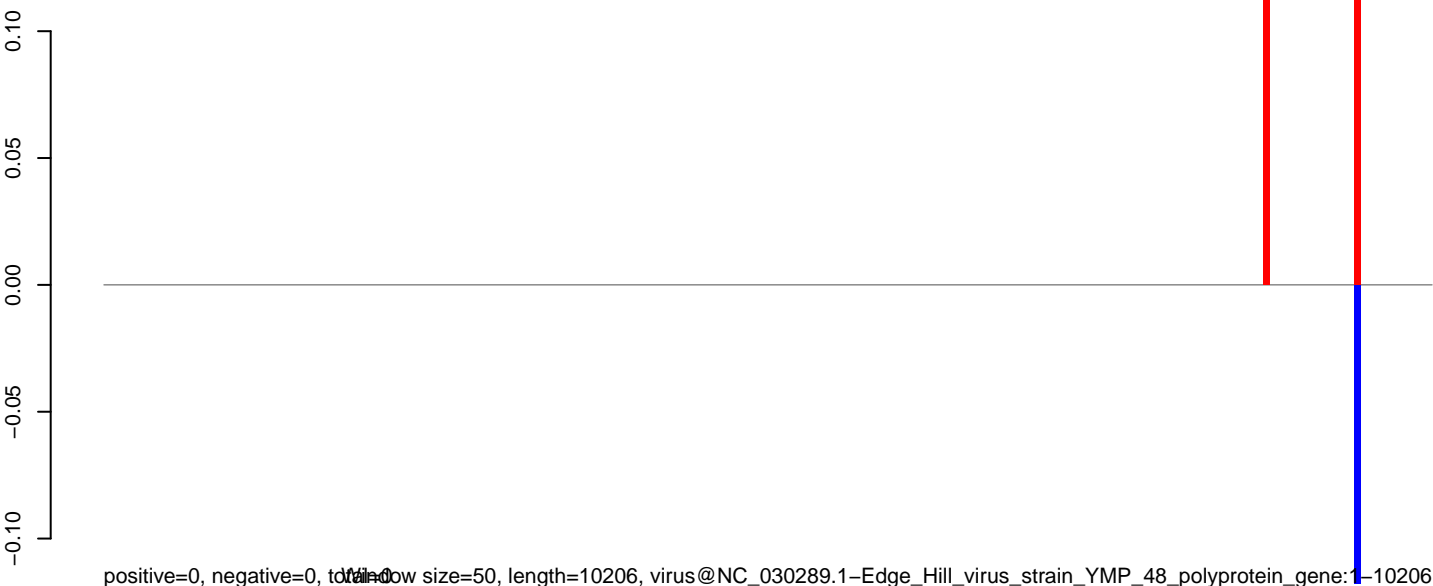
AeAeg_Aag2_DENV1.18_23.rep



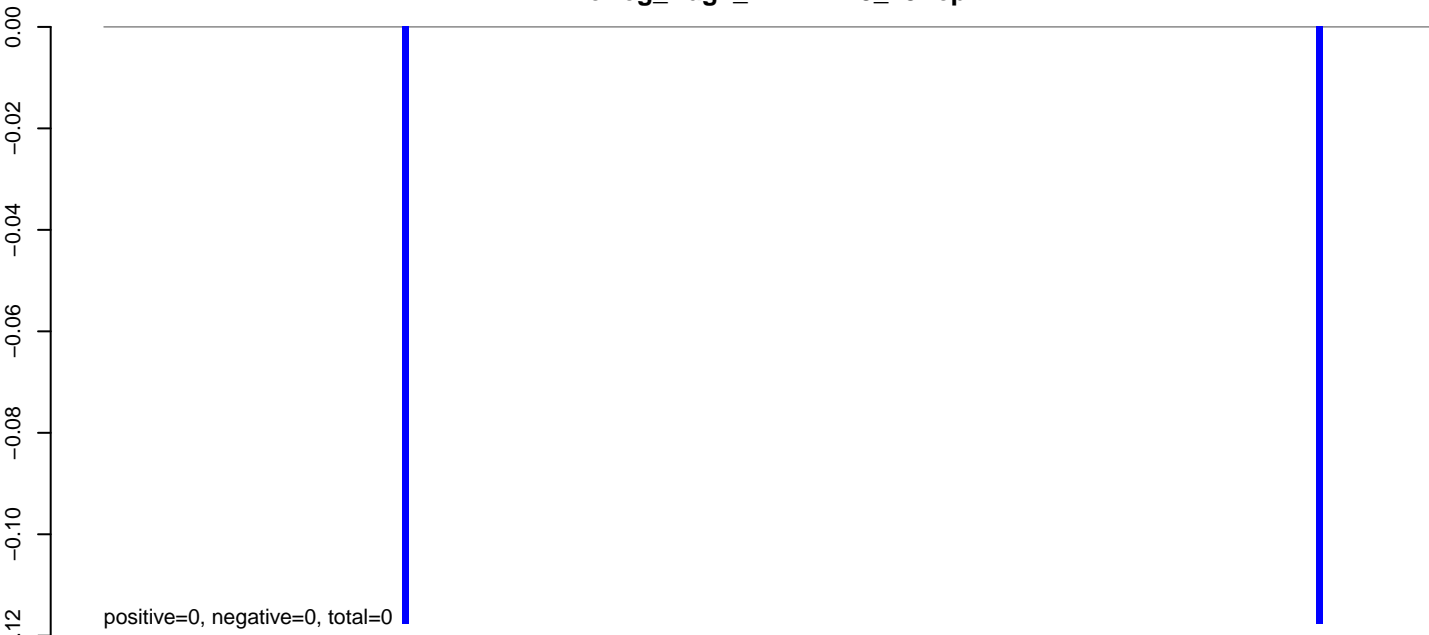
AeAeg_Aag2_DENV1.24_35.rep



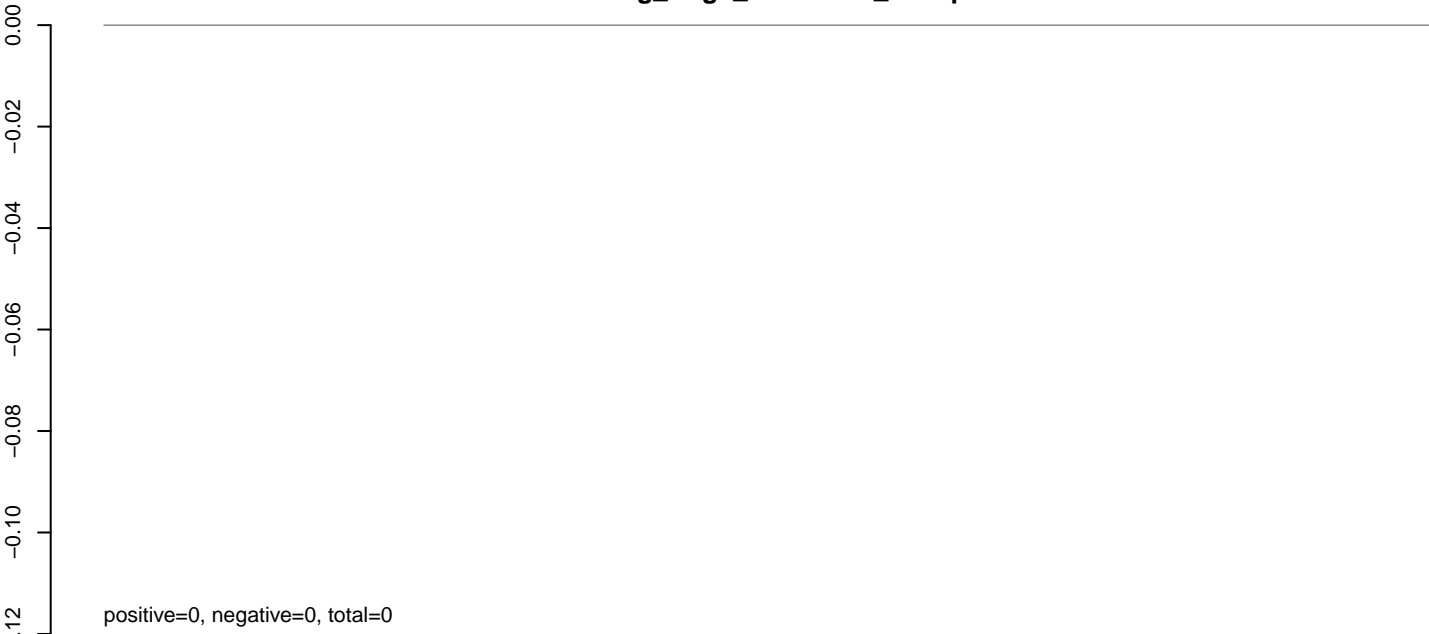
AeAeg_Aag2_DENV1.rep



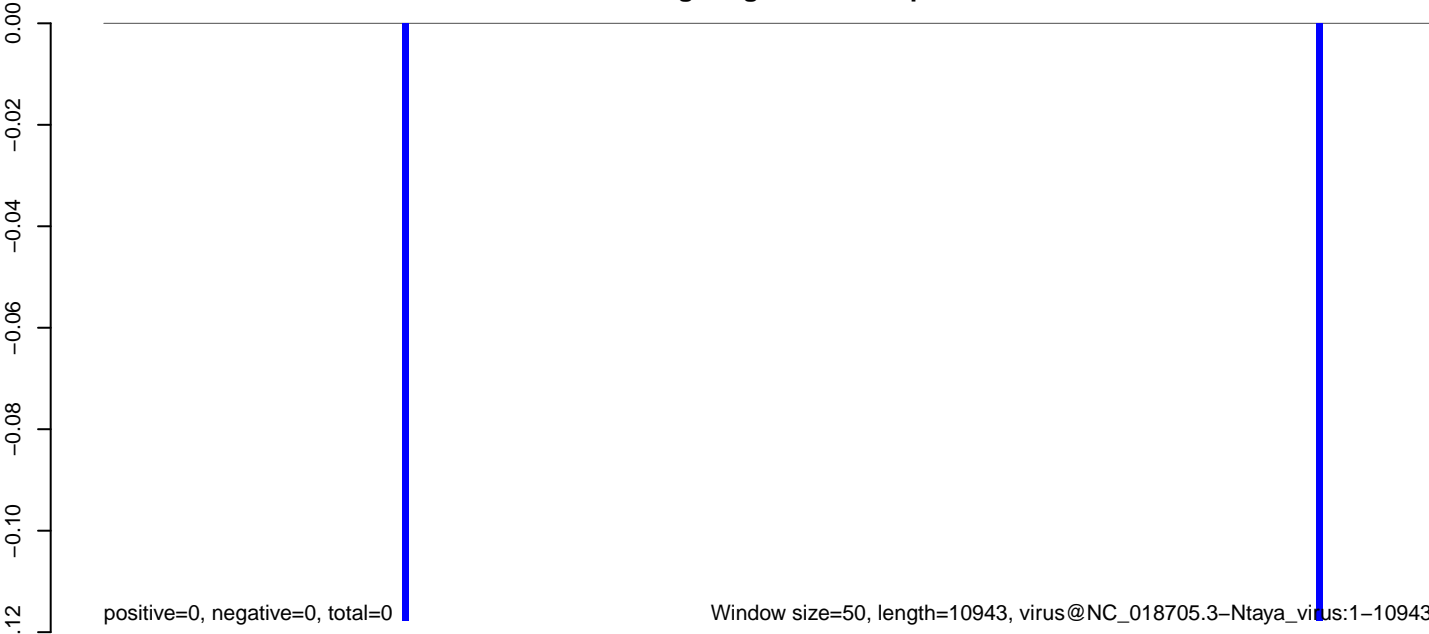
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

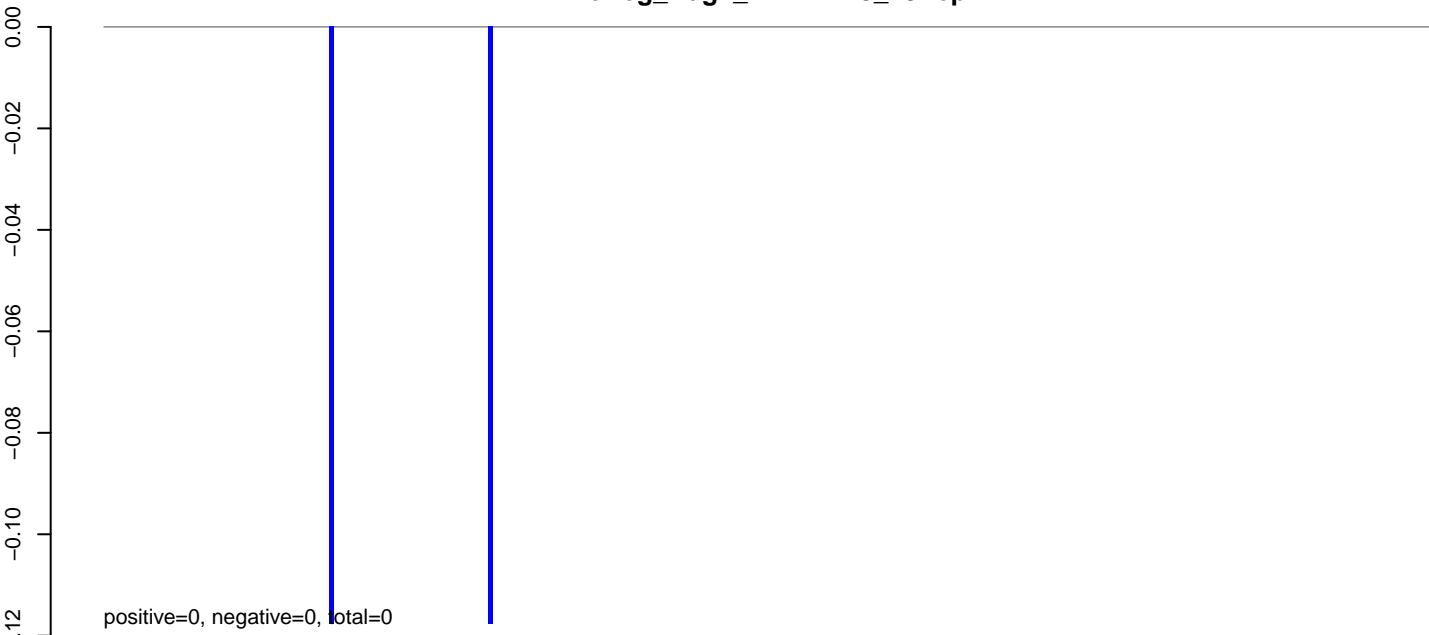


AeAeg_Aag2_DENV1.rep

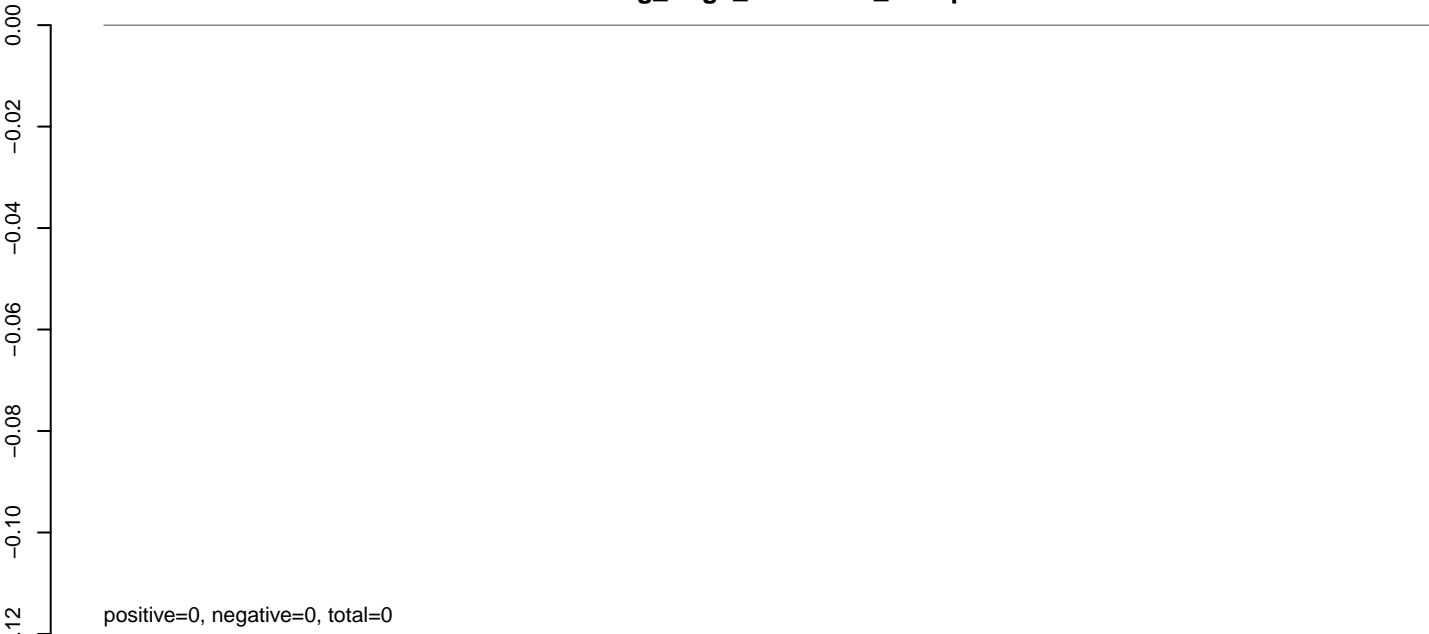


0 2000 4000 6000 8000 10000

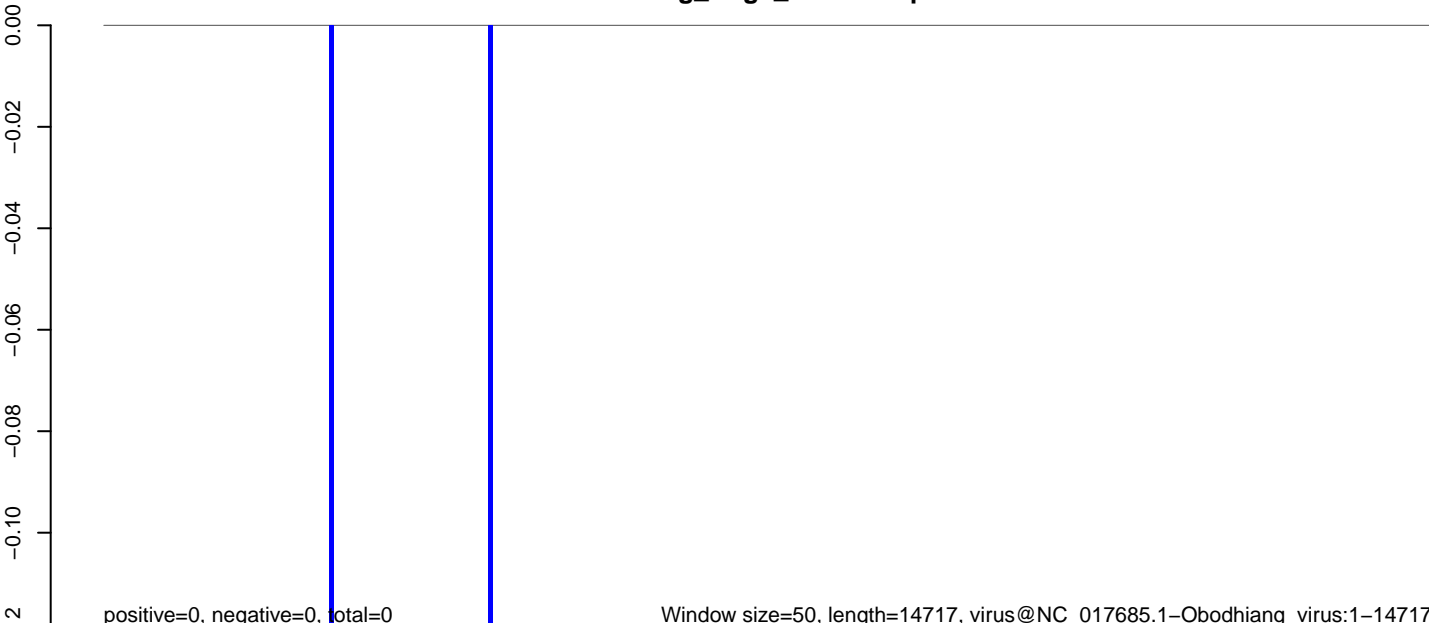
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

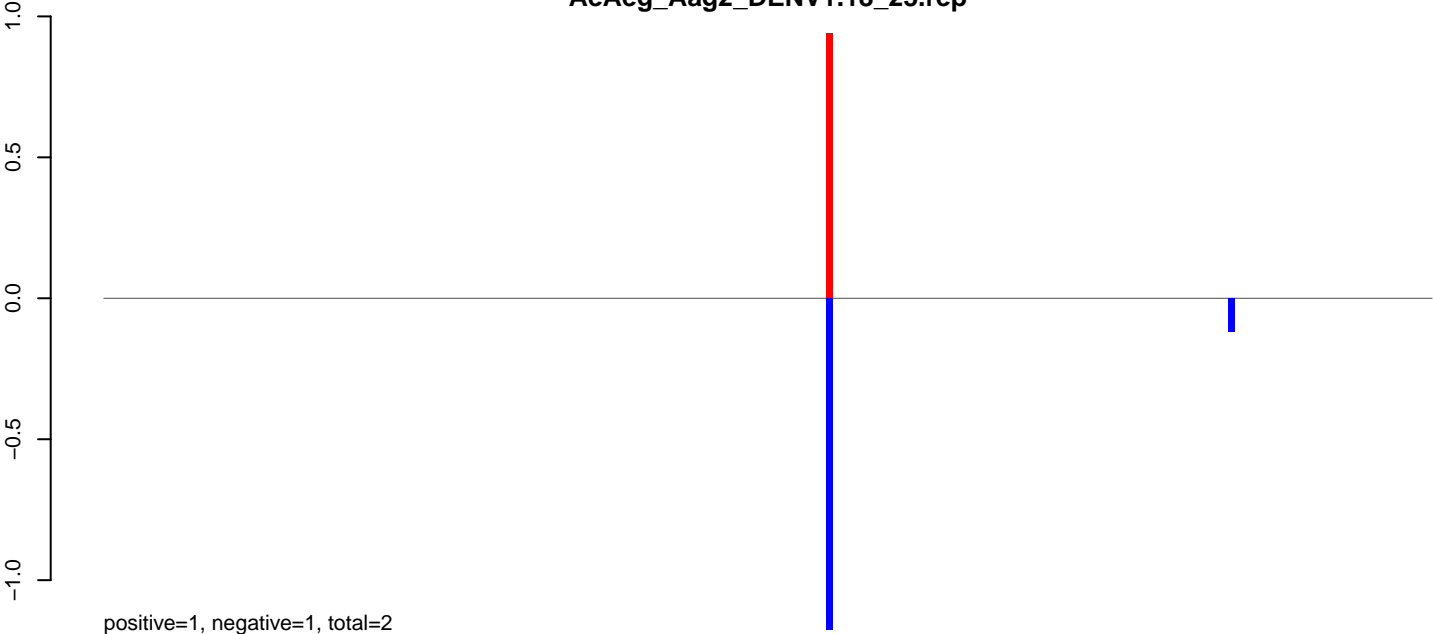


AeAeg_Aag2_DENV1.rep



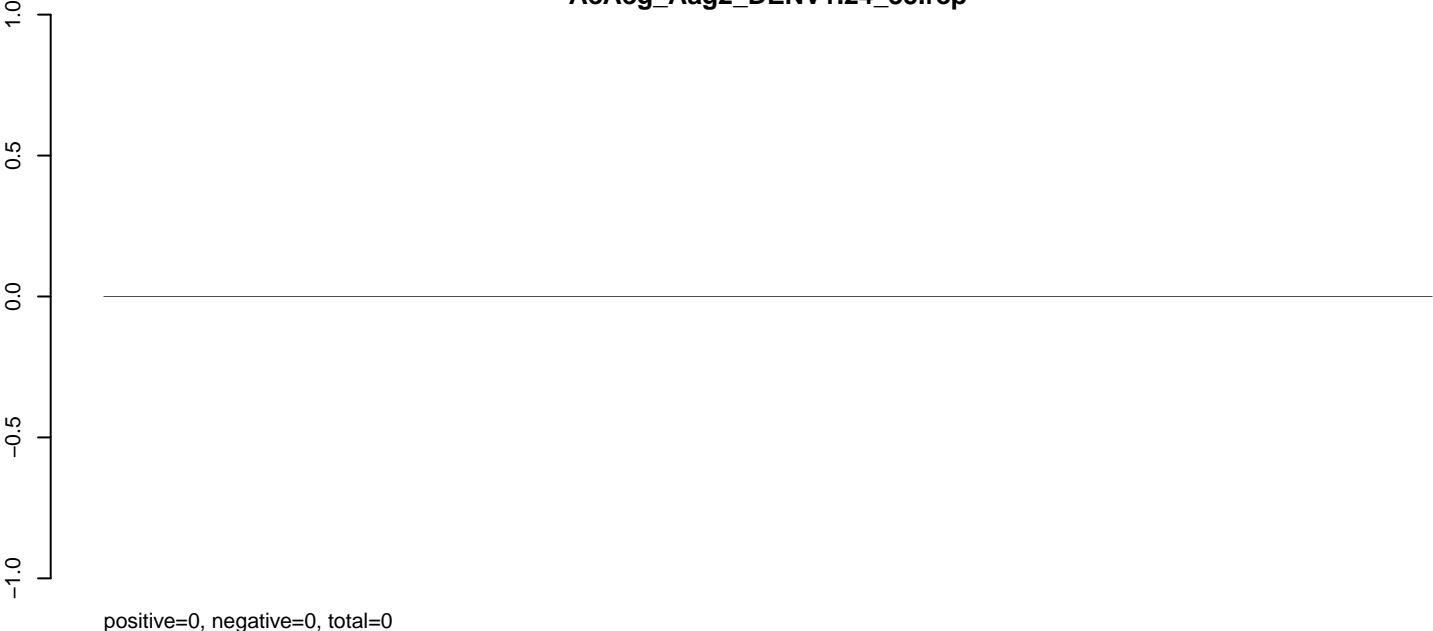
Window size=50, length=14717, virus@NC_017685.1-Obodhiang_virus:1-14717

AeAeg_Aag2_DENV1.18_23.rep



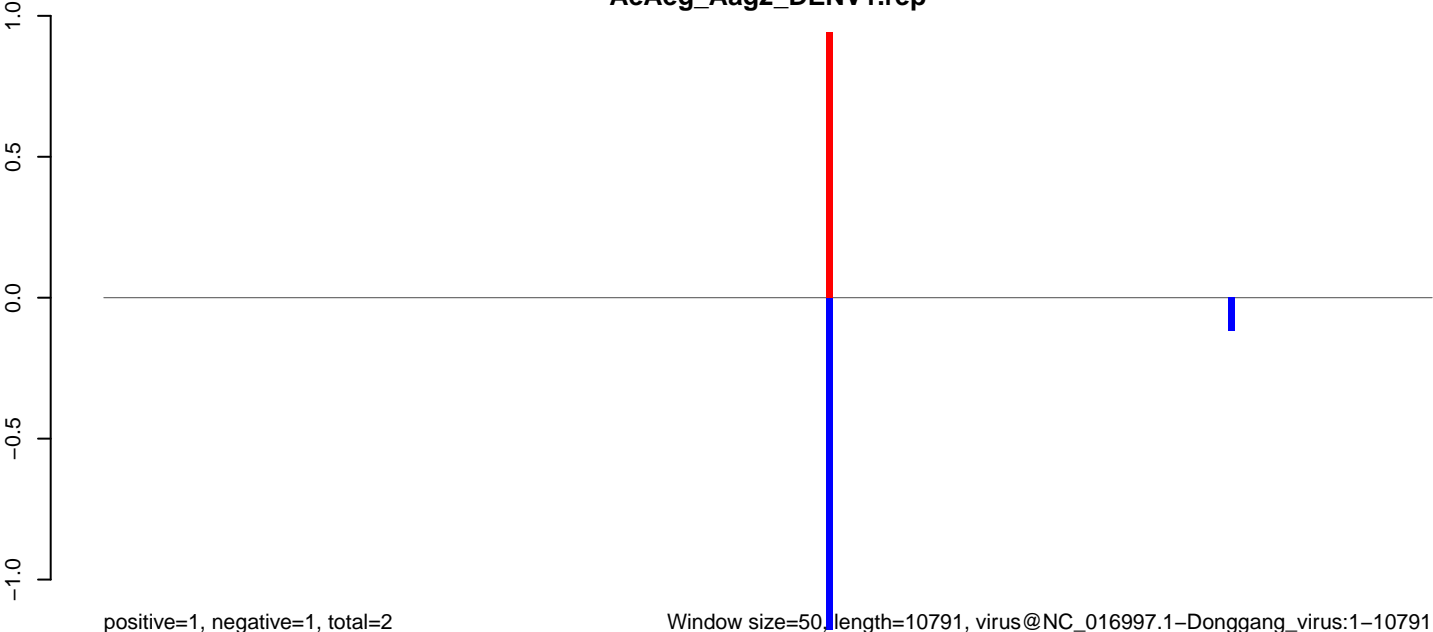
positive=1, negative=1, total=2

AeAeg_Aag2_DENV1.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_DENV1.rep

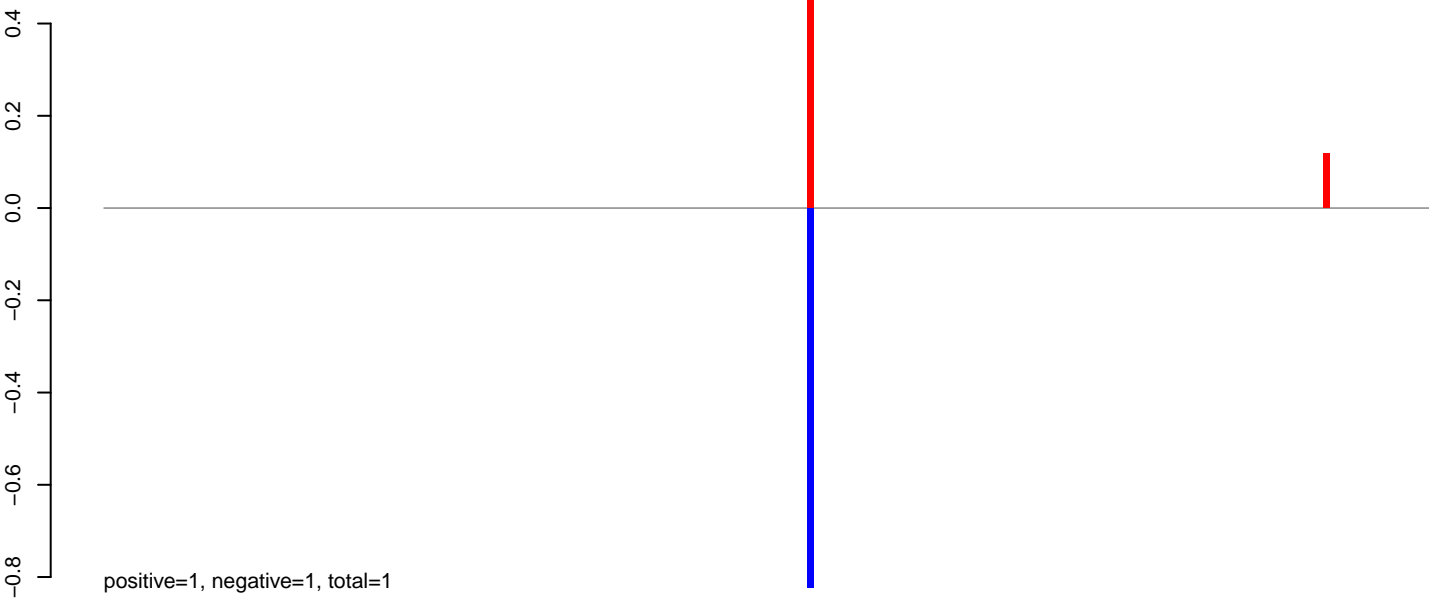


positive=1, negative=1, total=2

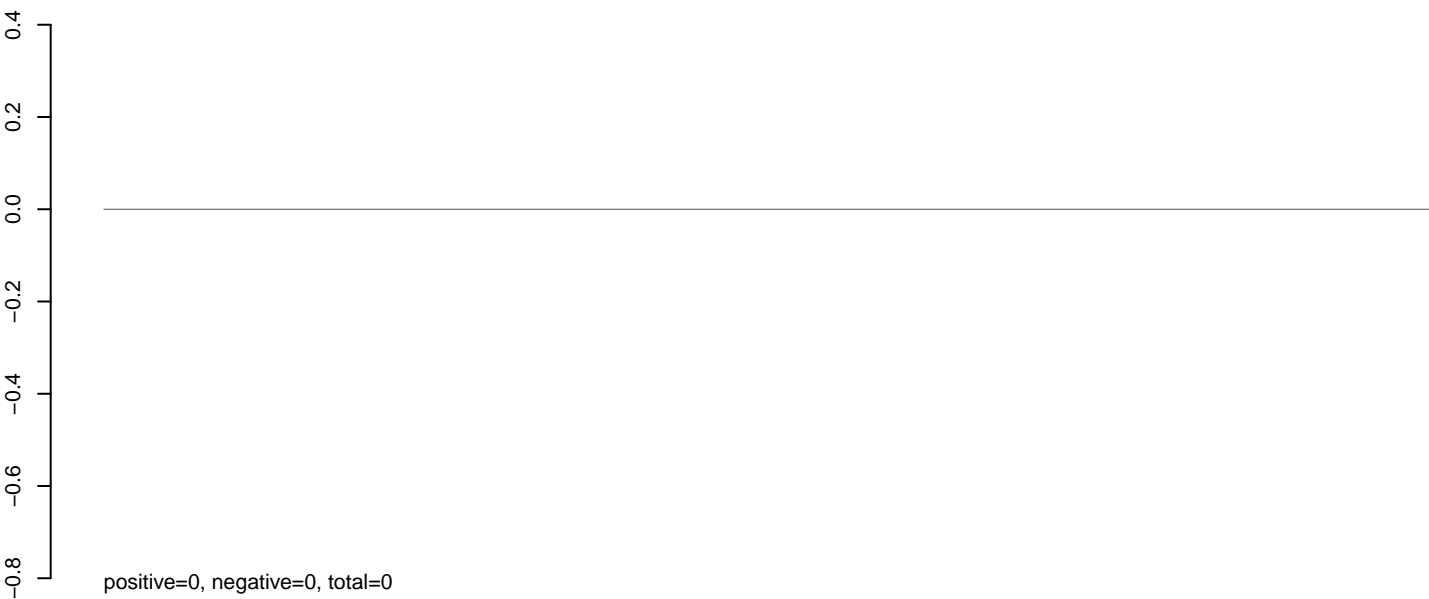
Window size=50, length=10791, virus@NC_016997.1-Donggang_virus:1-10791

0 2000 4000 6000 8000 10000

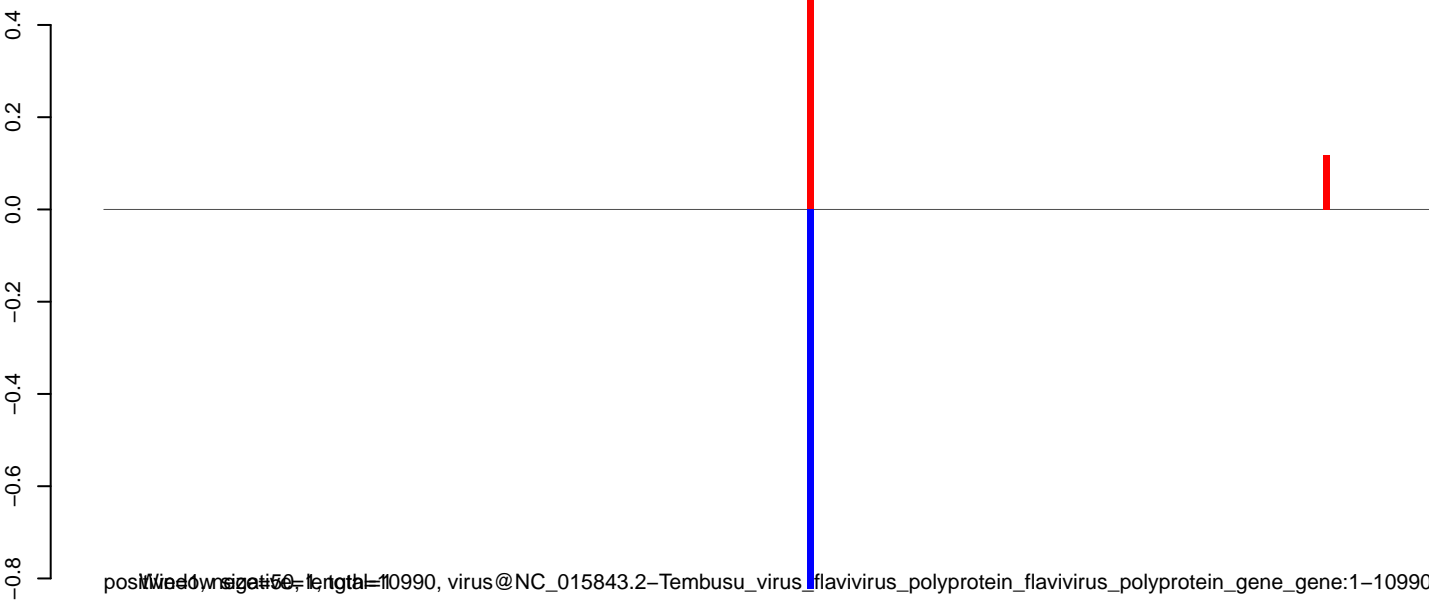
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

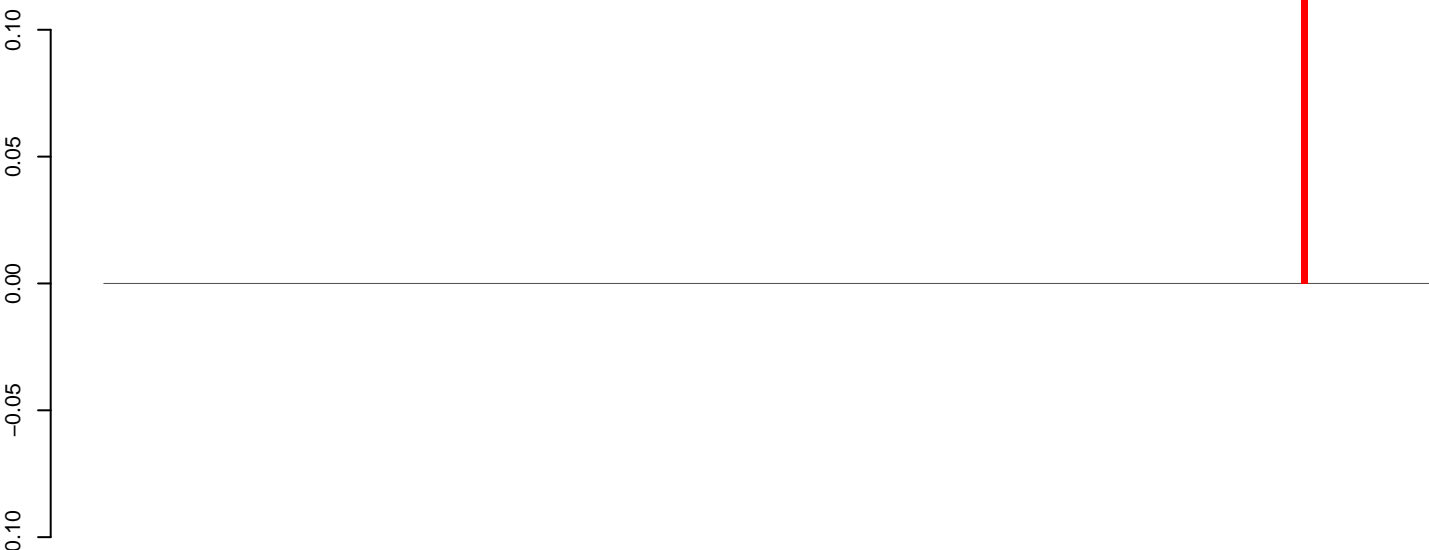


AeAeg_Aag2_DENV1.rep



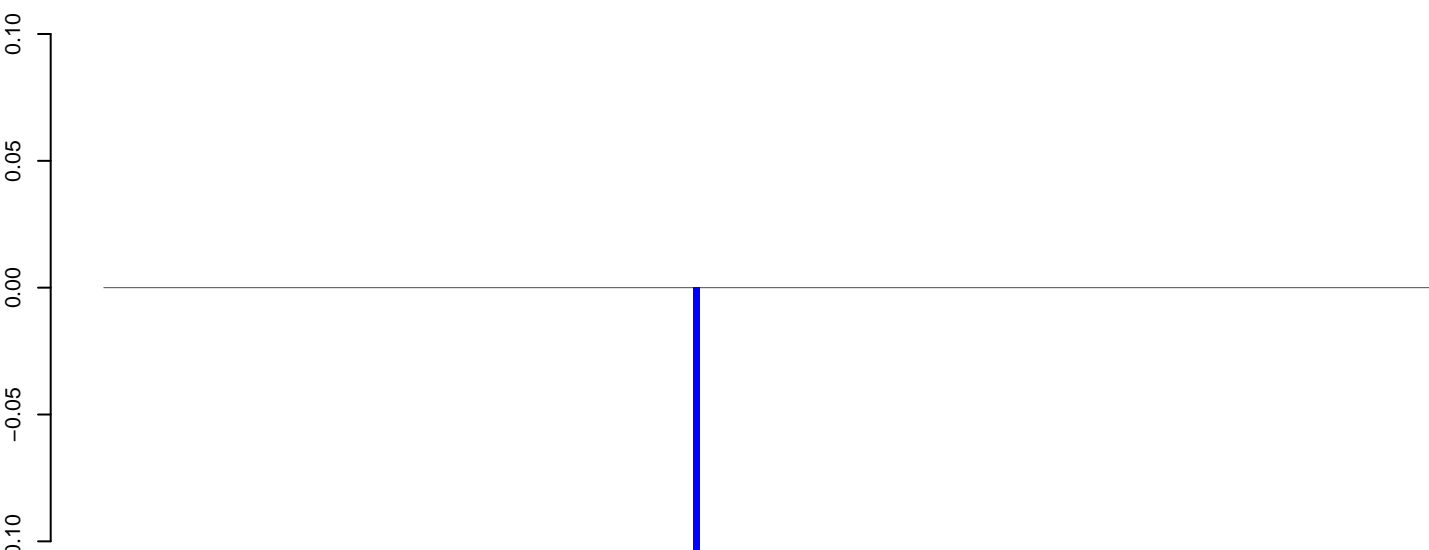
Windows size = 50, length = 10990, virus@NC_015843.2-Tembusu_virus_flavivirus_polyprotein_flavivirus_polyprotein_gene_gene:1-10990

AeAeg_Aag2_DENV1.18_23.rep



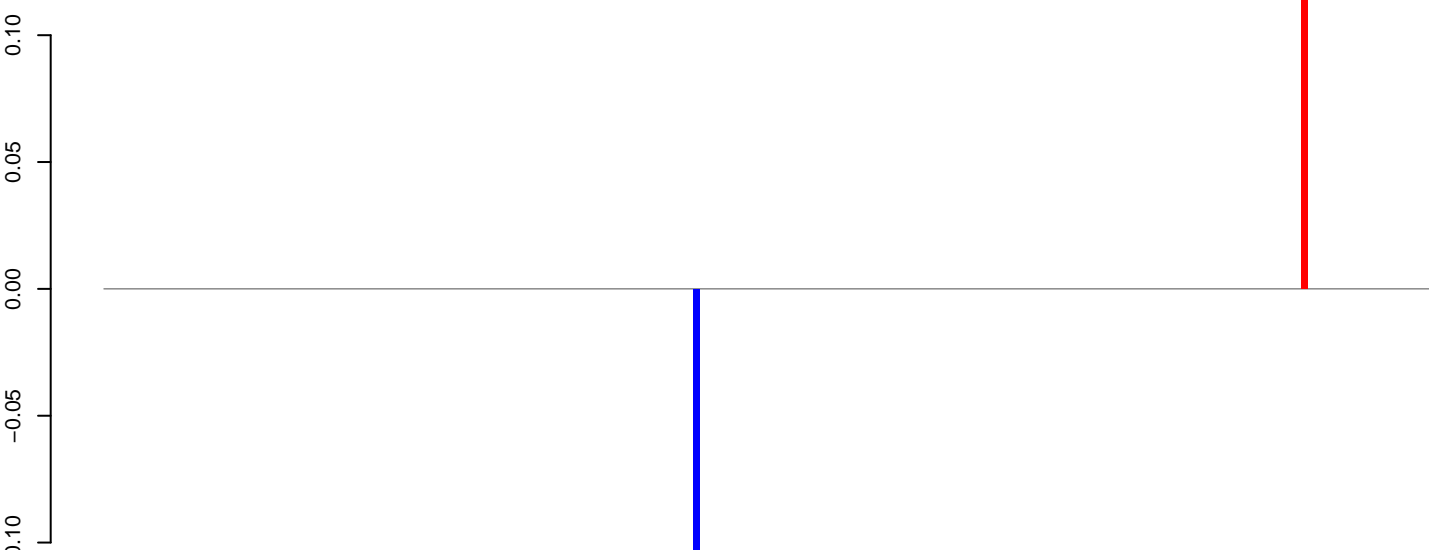
positive=0, negative=0, total=0

AeAeg_Aag2_DENV1.24_35.rep



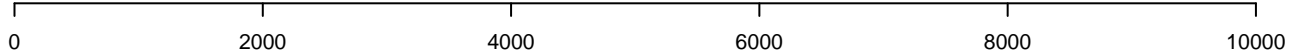
positive=0, negative=0, total=0

AeAeg_Aag2_DENV1.rep

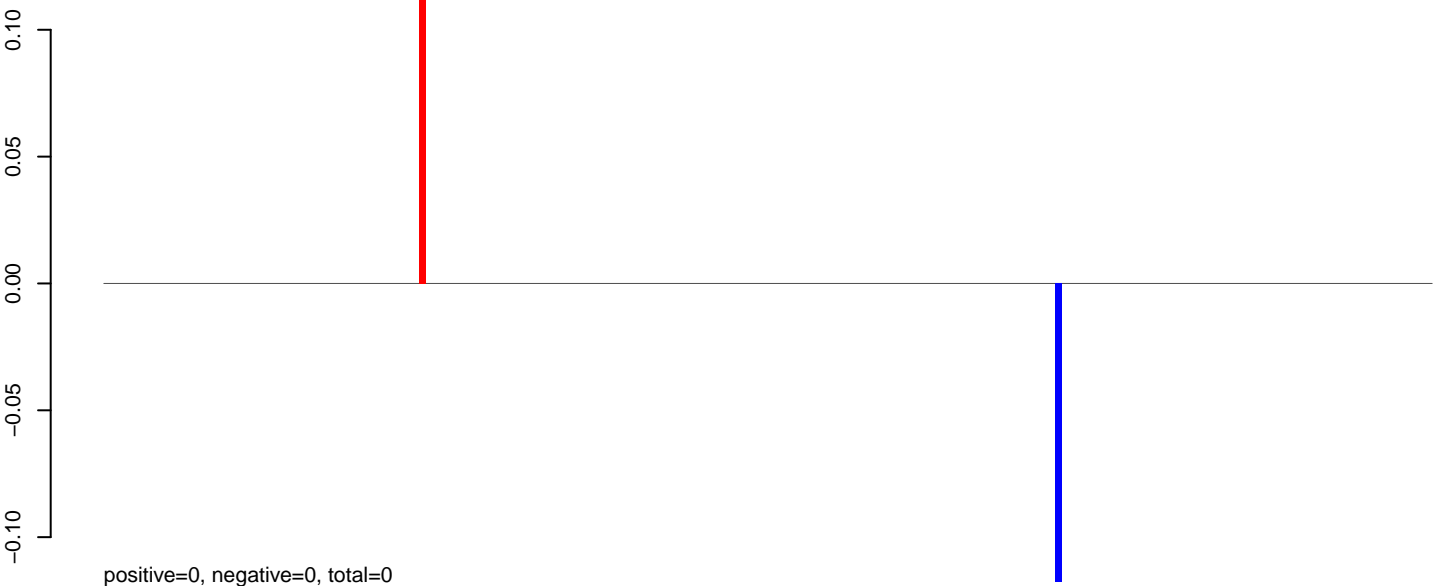


positive=0, negative=0, total=0

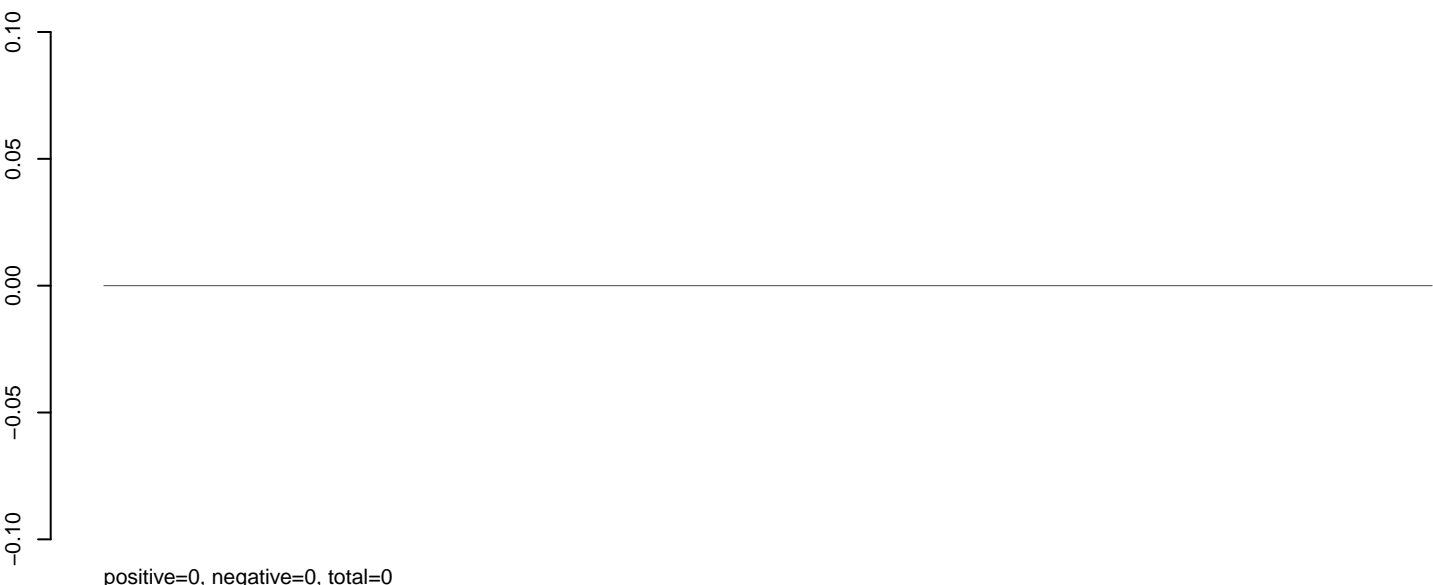
Window size=50, length=10723, virus@NC_012533.1-Kedougou_virus:1-10723



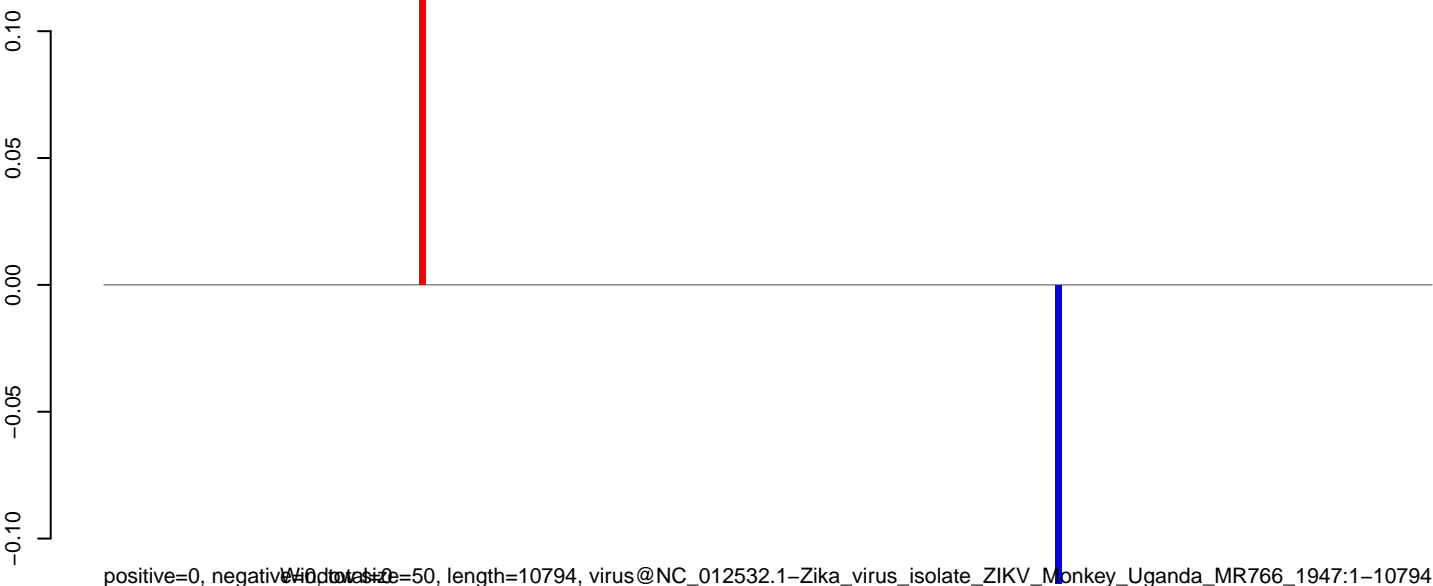
AeAeg_Aag2_DENV1.18_23.rep



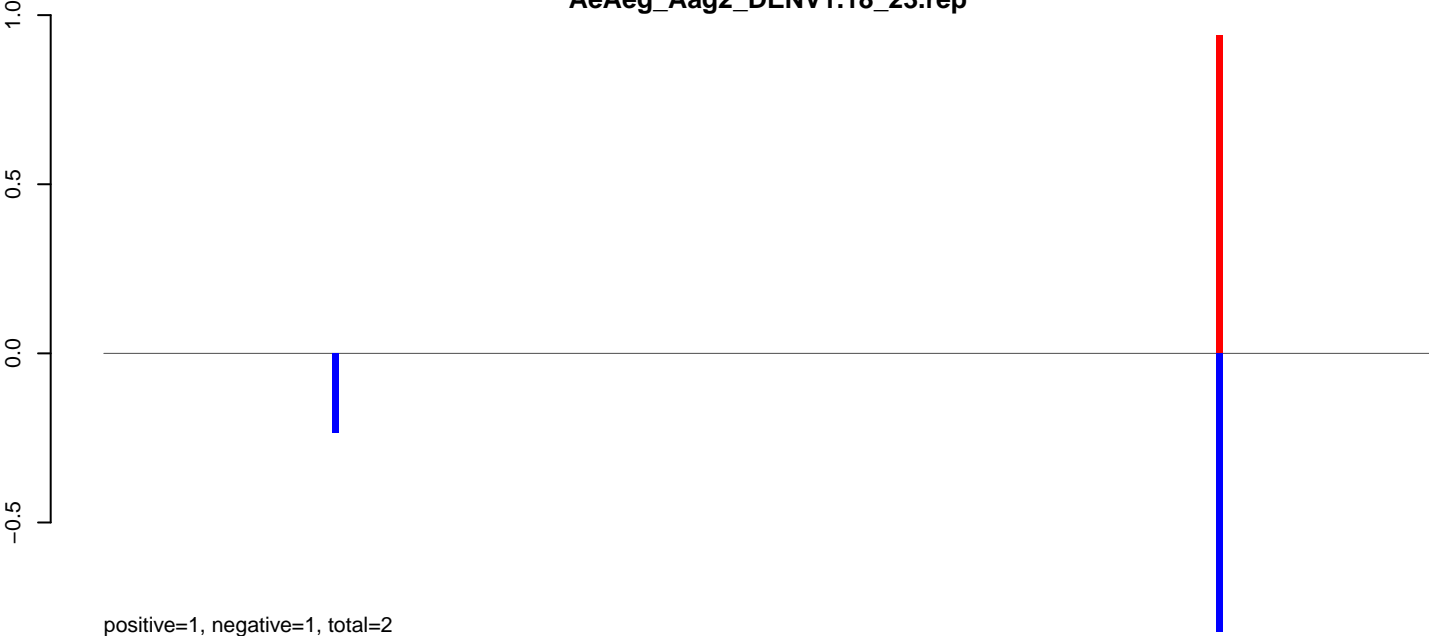
AeAeg_Aag2_DENV1.24_35.rep



AeAeg_Aag2_DENV1.rep

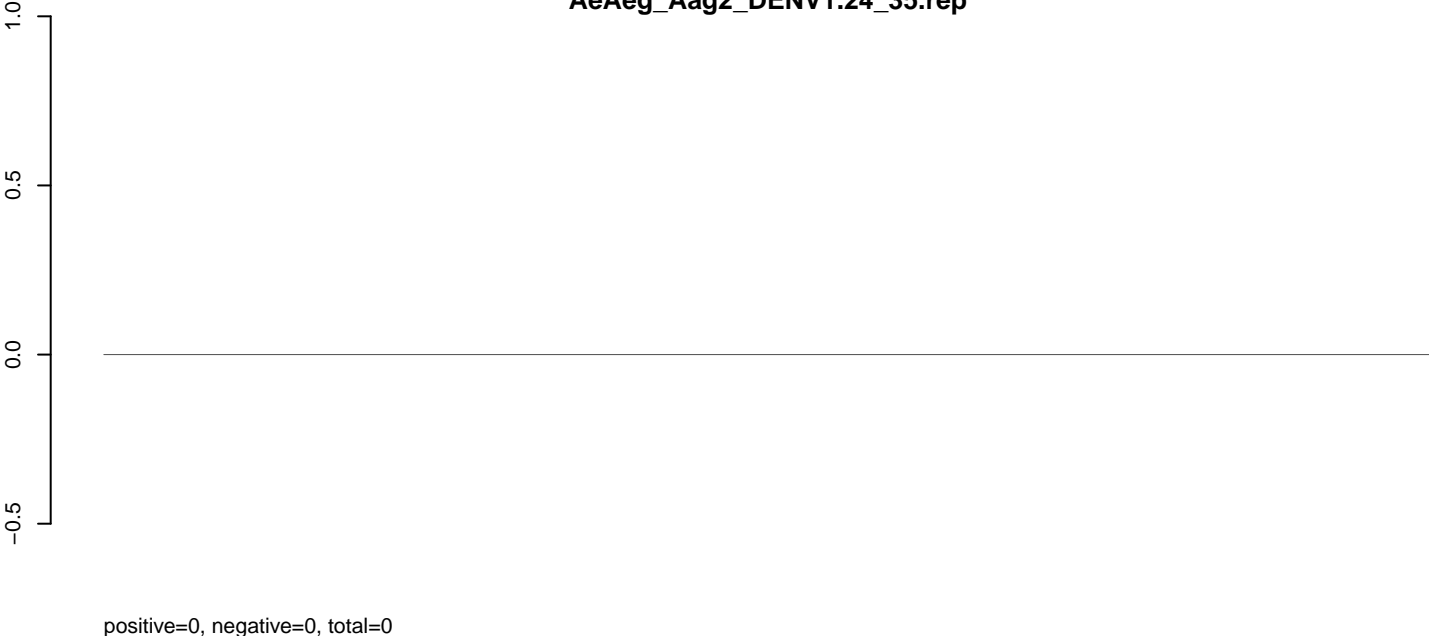


AeAeg_Aag2_DENV1.18_23.rep



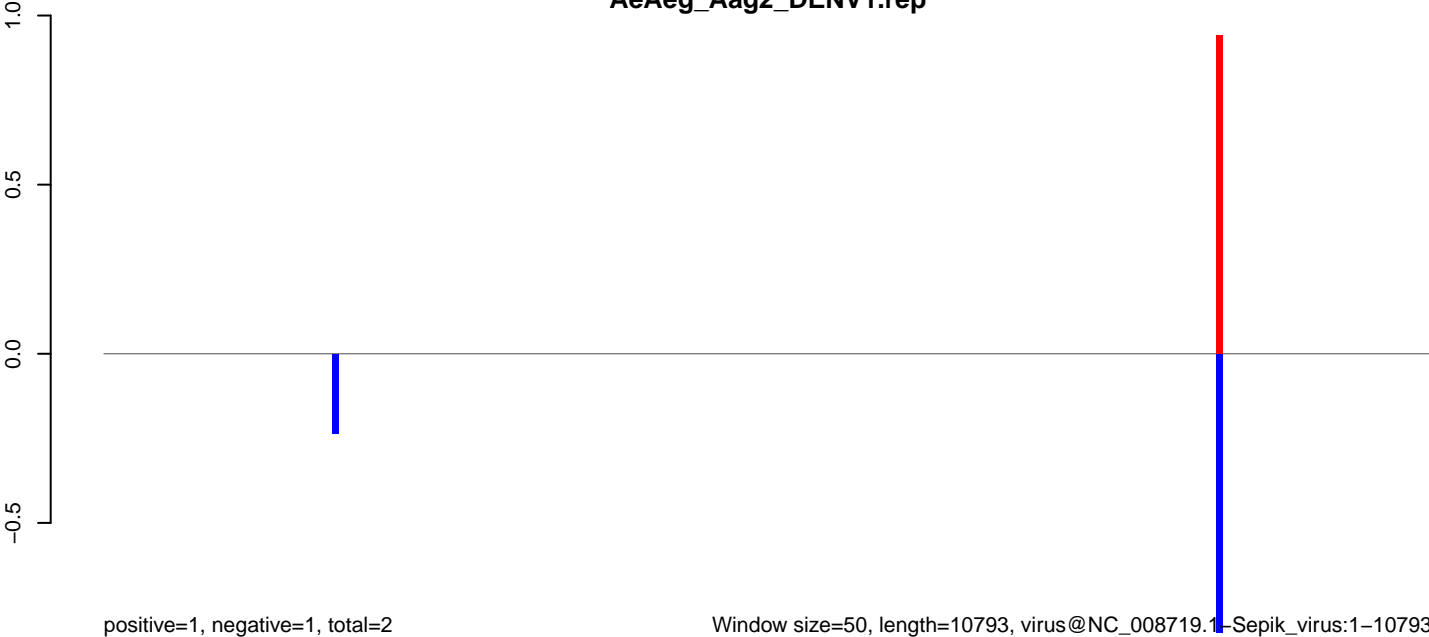
positive=1, negative=1, total=2

AeAeg_Aag2_DENV1.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2_DENV1.rep



positive=1, negative=1, total=2

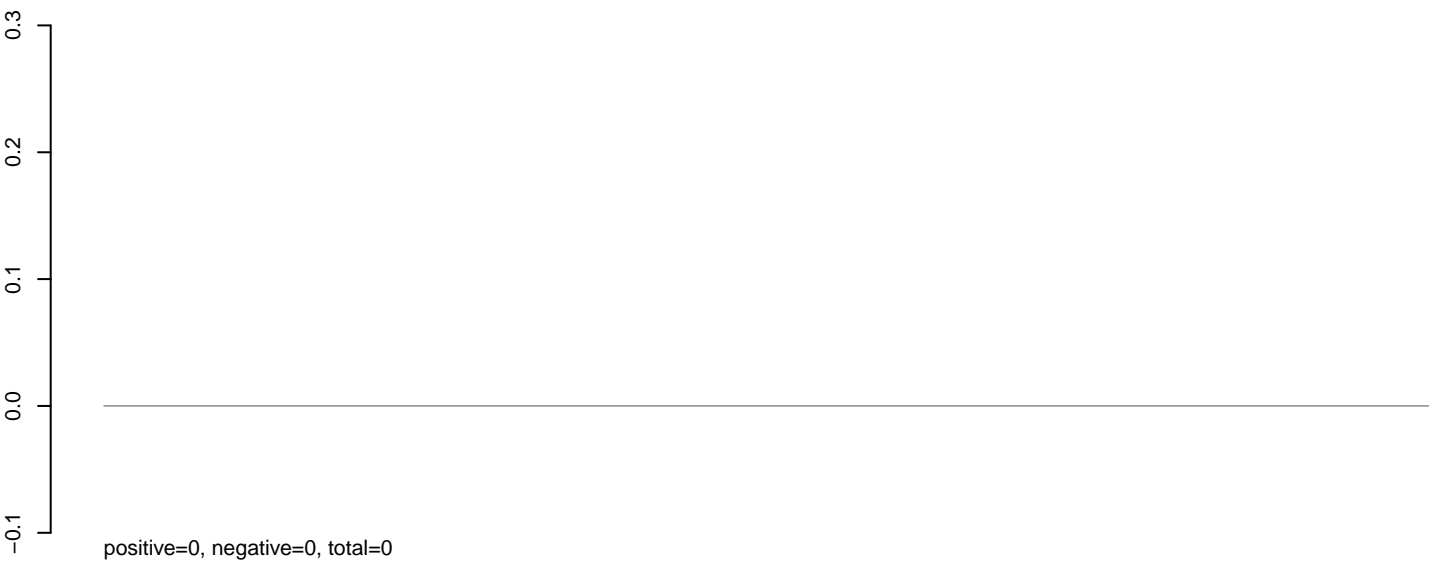
Window size=50, length=10793, virus@NC_008719.1-Sepik_virus:1-10793

0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

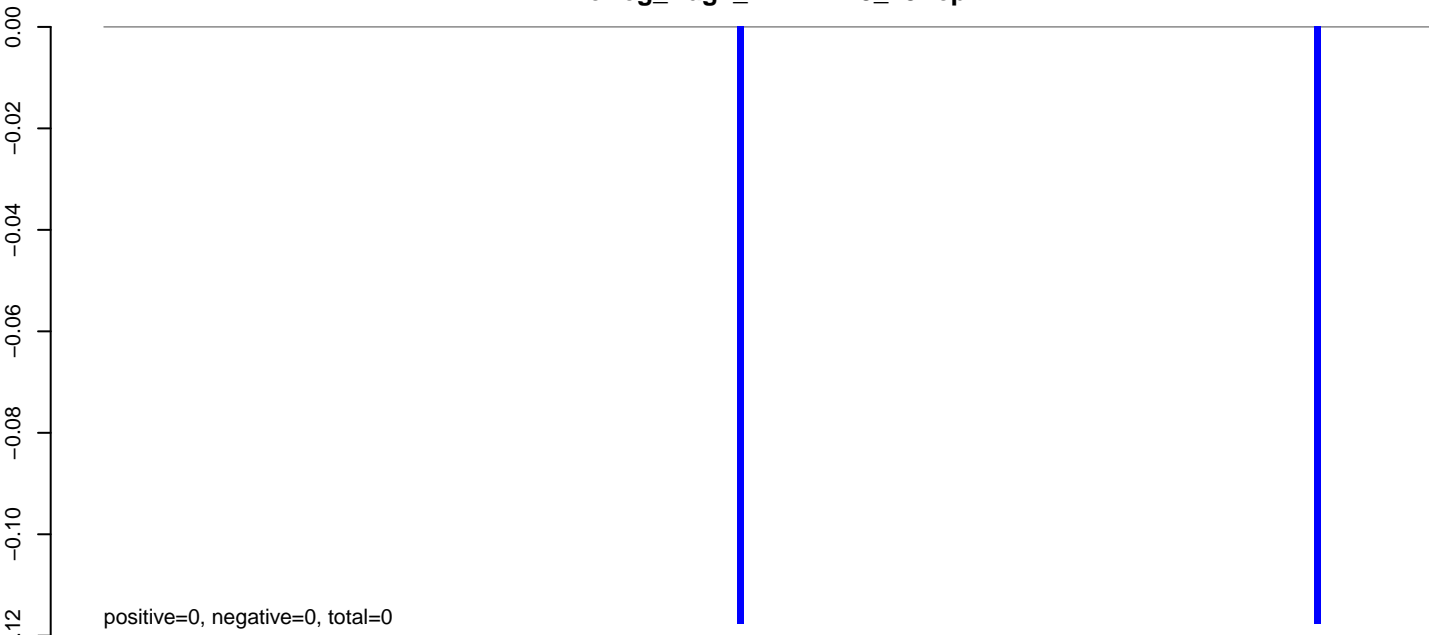


AeAeg_Aag2_DENV1.rep

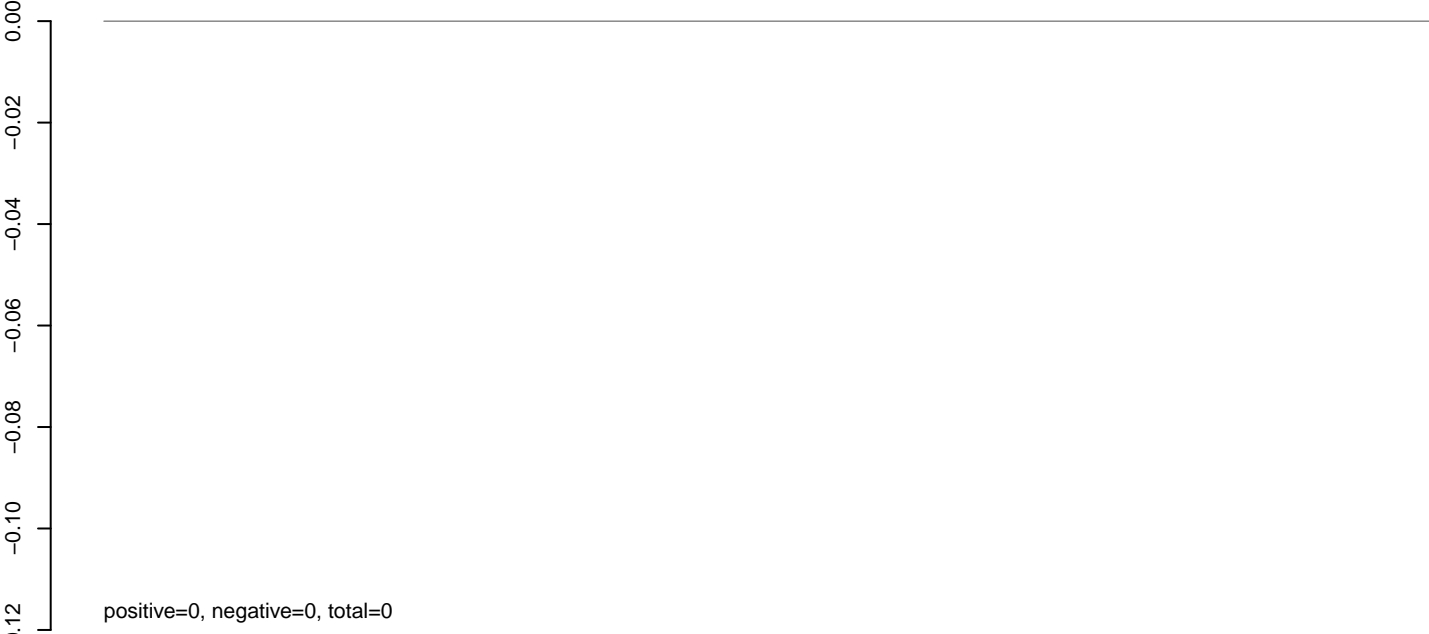


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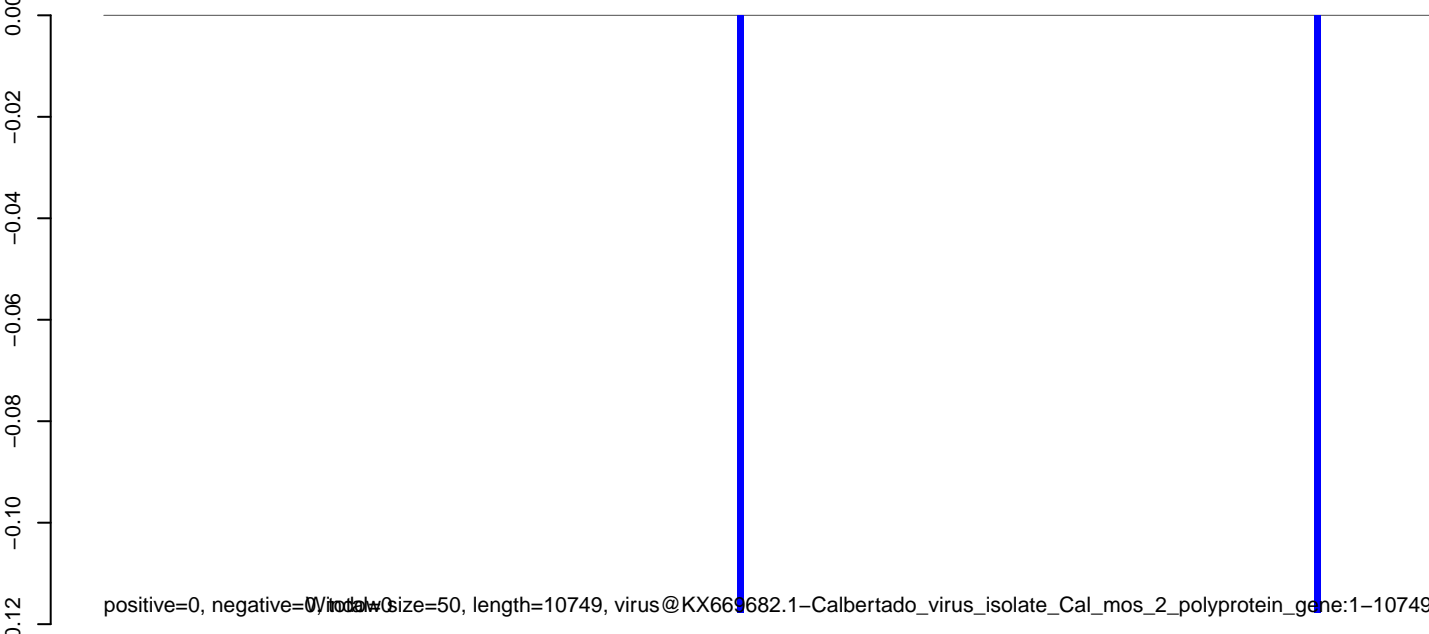
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



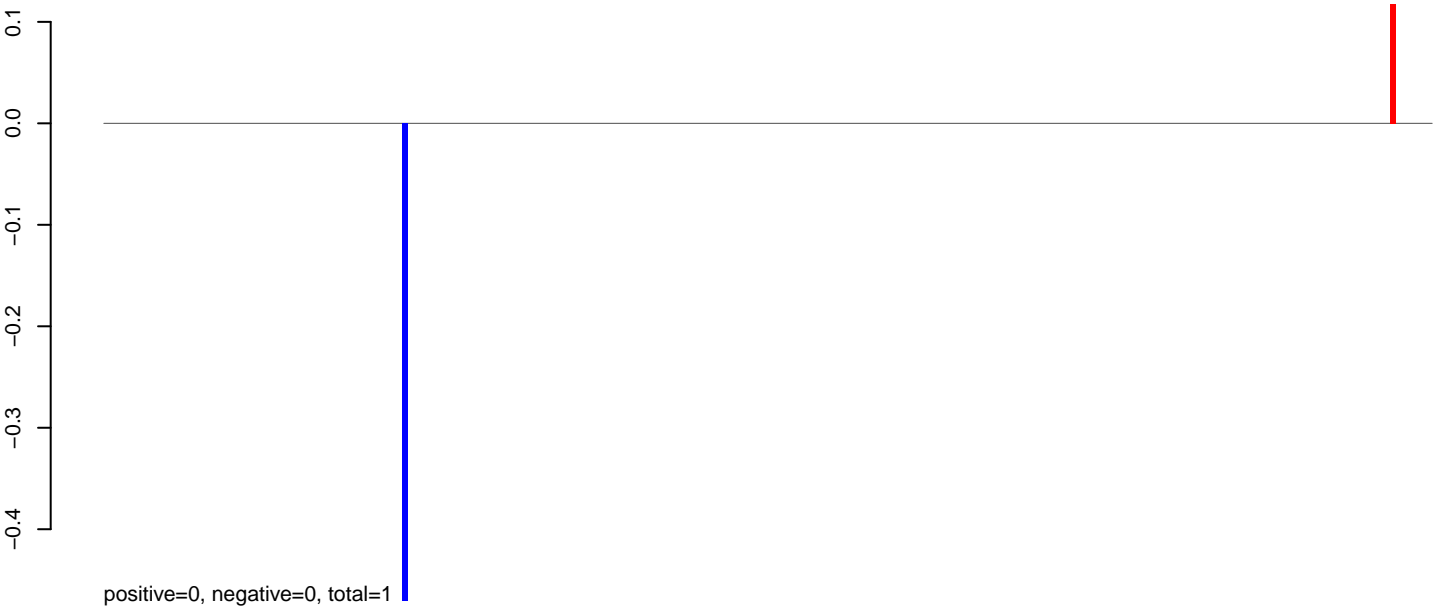
AeAeg_Aag2_DENV1.rep



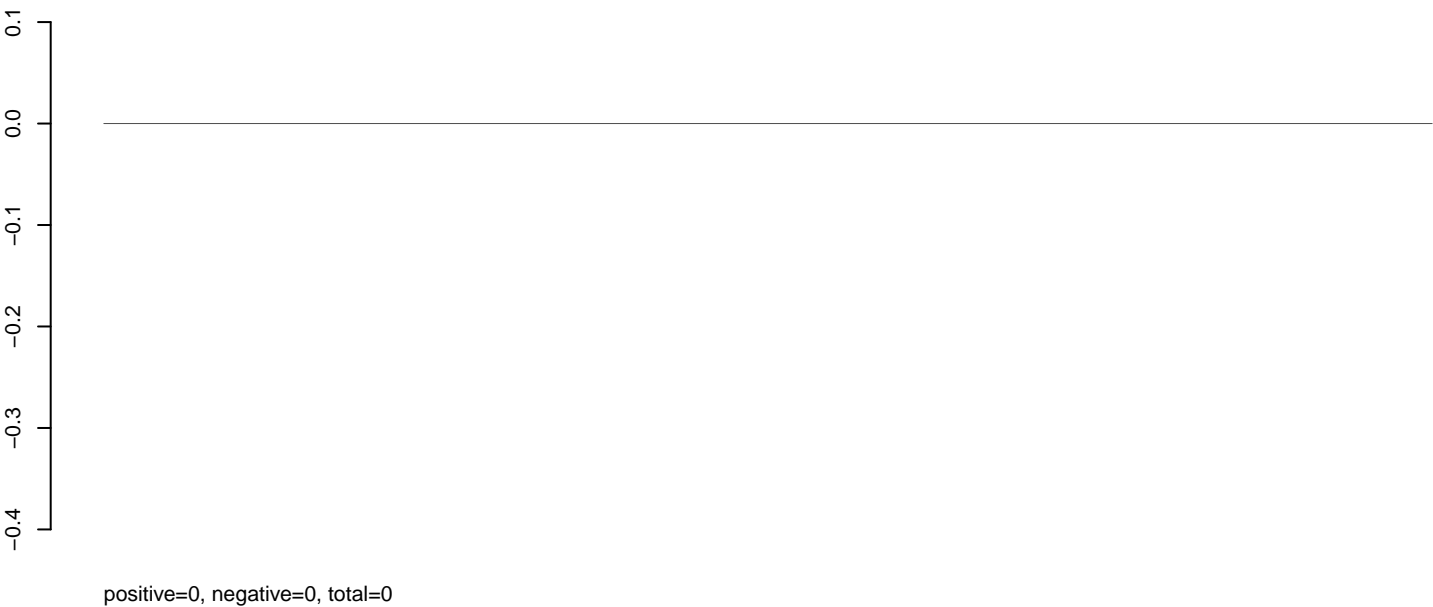
Window size=50, length=10749, virus@KX661682.1-Calbertado_virus_isolate_Cal_mos_2_polyprotein_gene:1-10749

0 2000 4000 6000 8000 10000

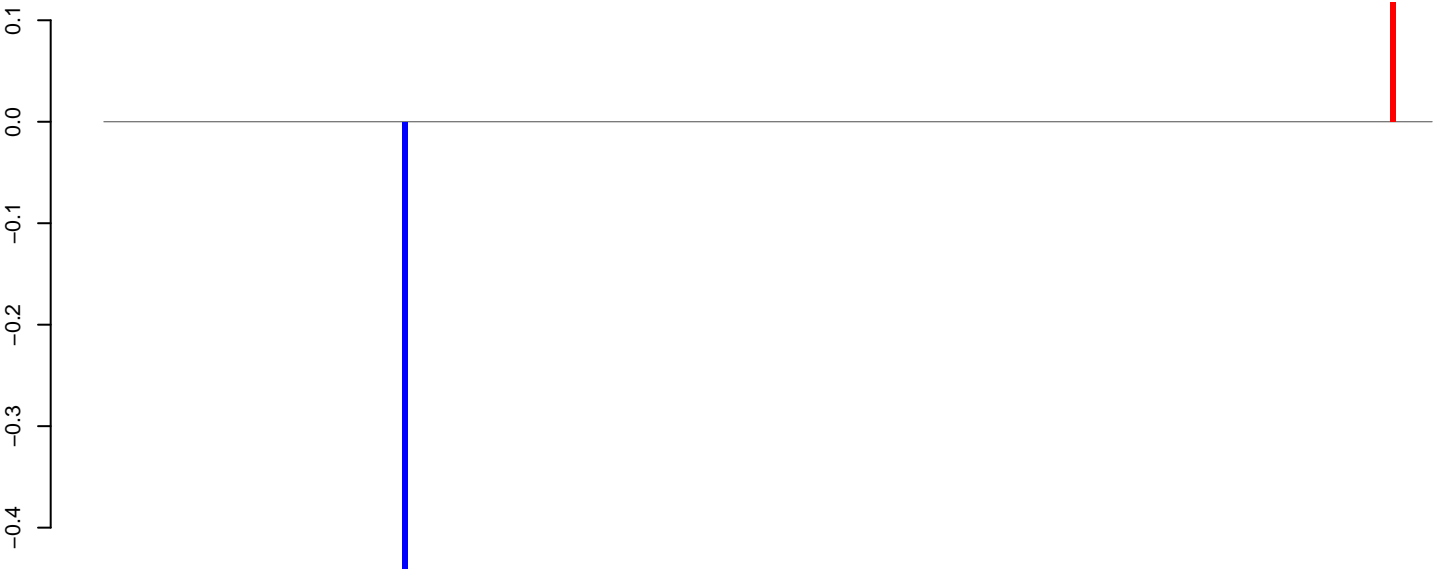
AeAeg_Aag2_DENV1.18_23.rep

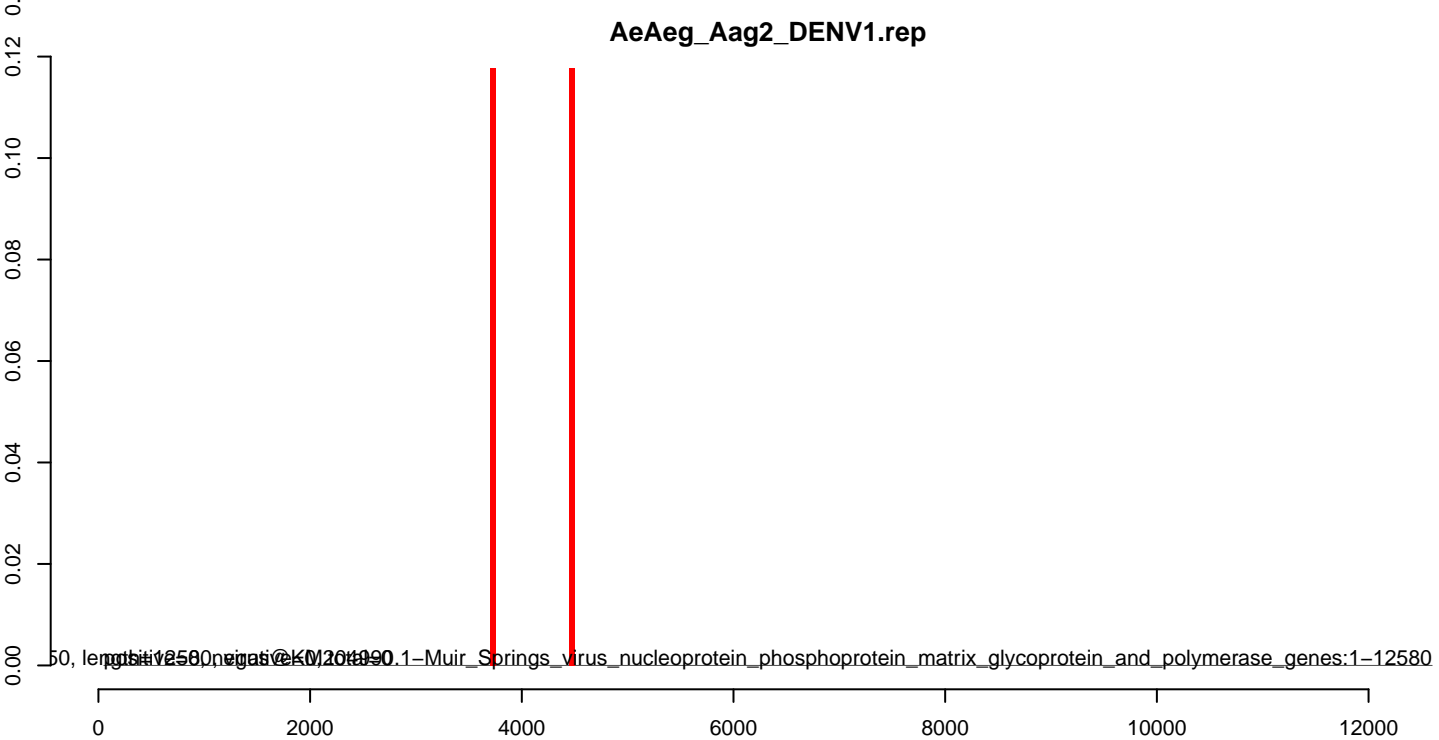
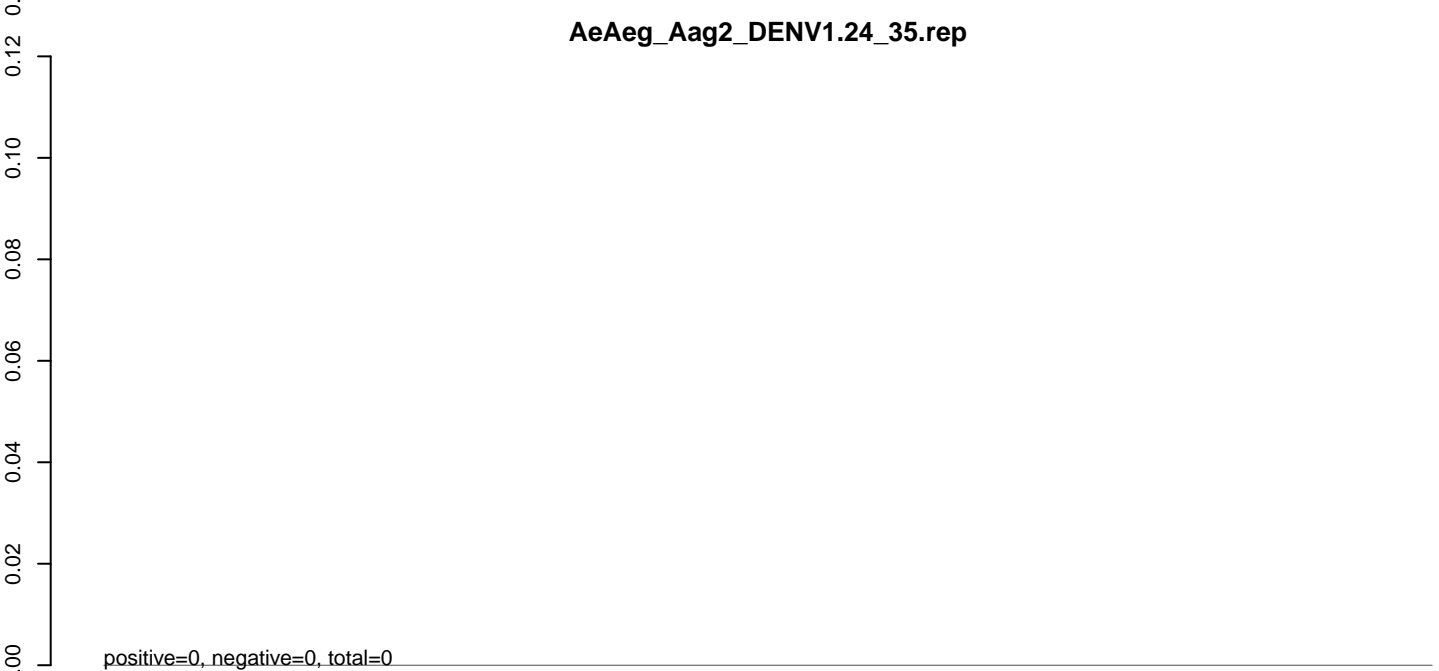
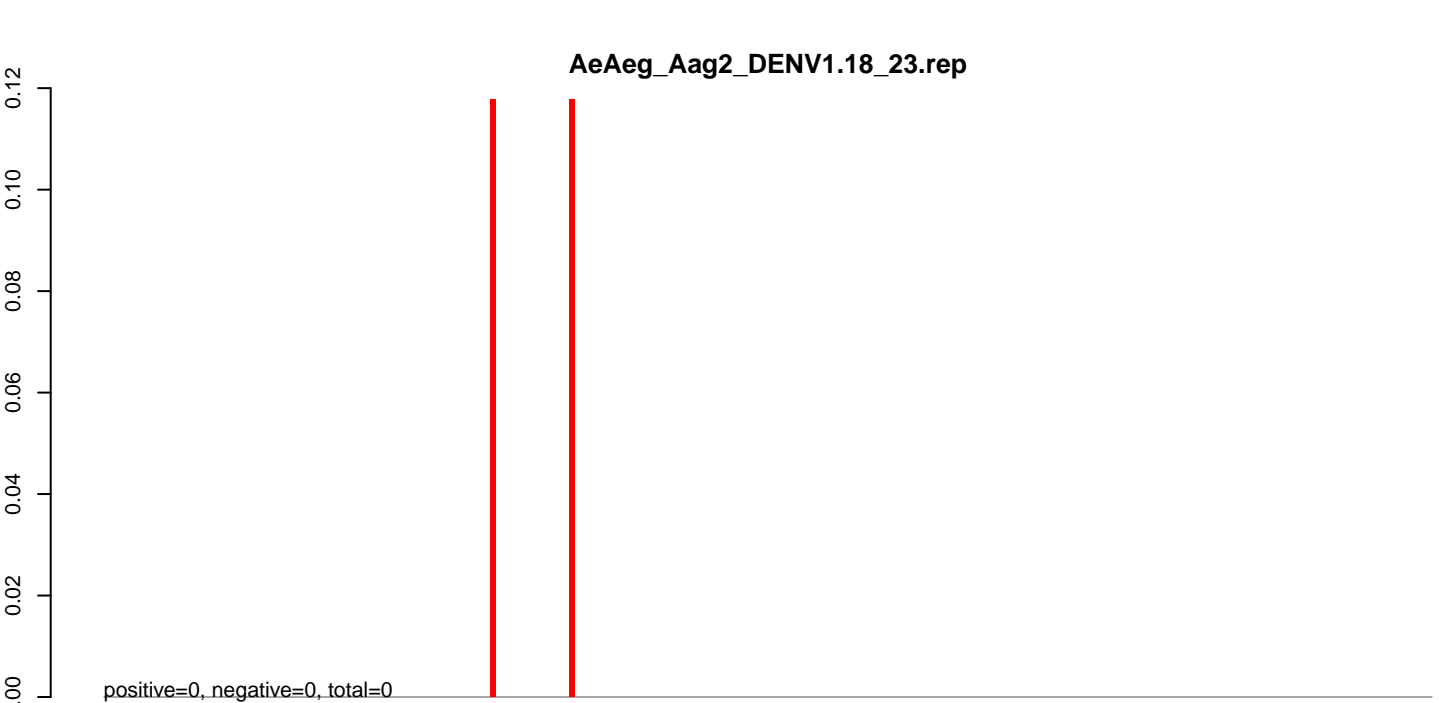


AeAeg_Aag2_DENV1.24_35.rep

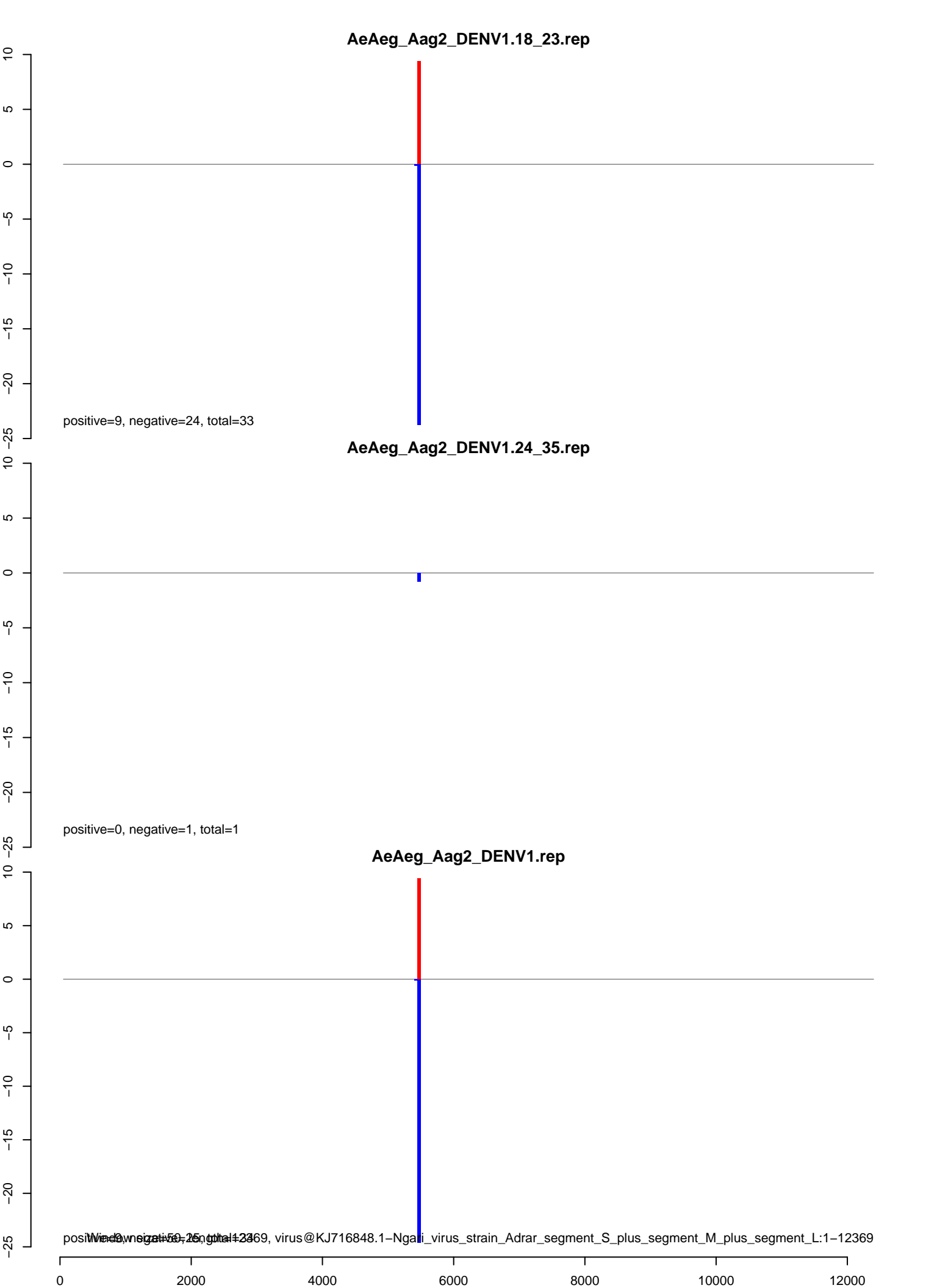


AeAeg_Aag2_DENV1.rep





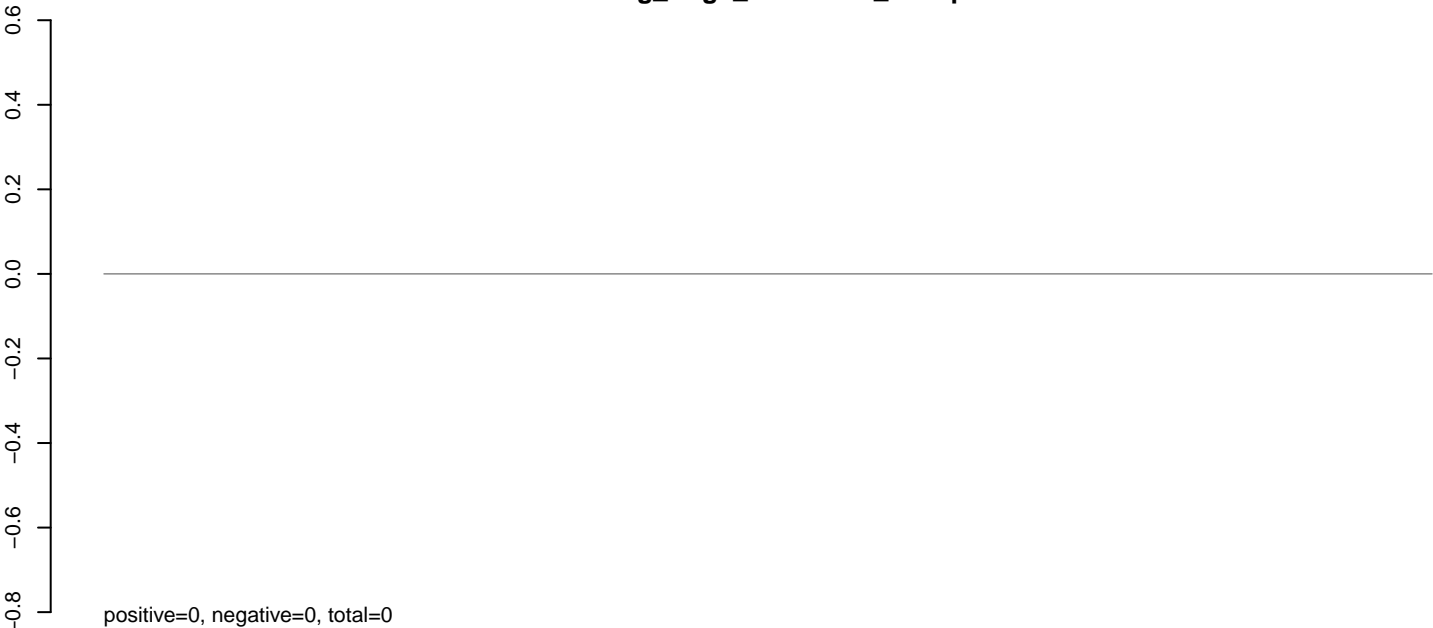
50, length=12580, negative=0, total=0, 1-Muir_Springs_virus_nucleoprotein_phosphoprotein_matrix_glycoprotein_and_polymerase_genes:1-12580



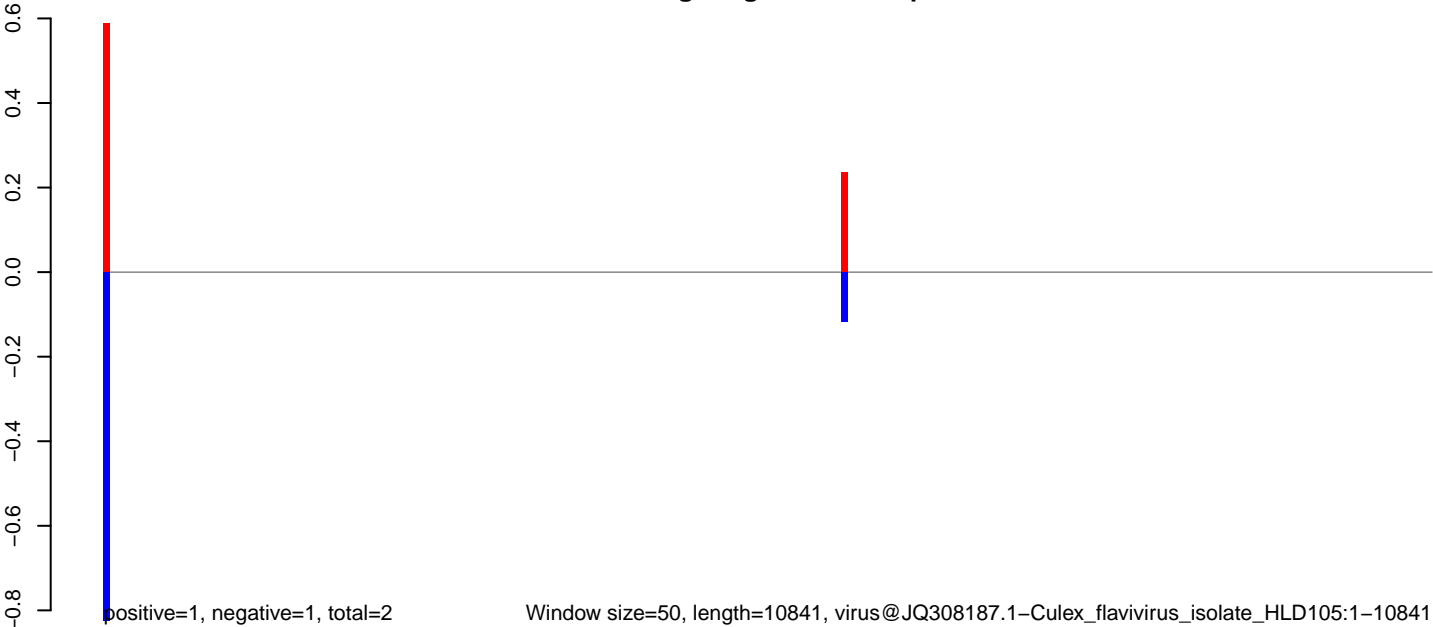
AeAeg_Aag2_DENV1.18_23.rep



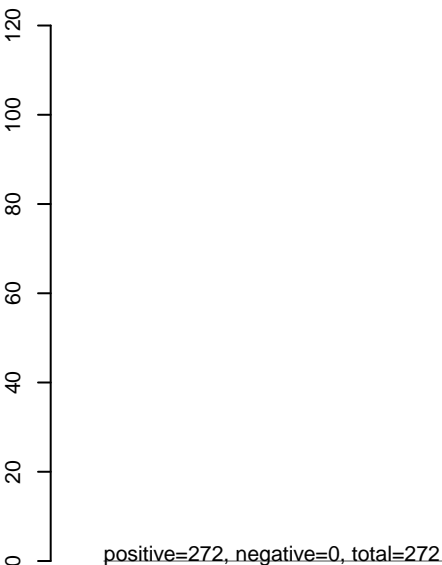
AeAeg_Aag2_DENV1.24_35.rep



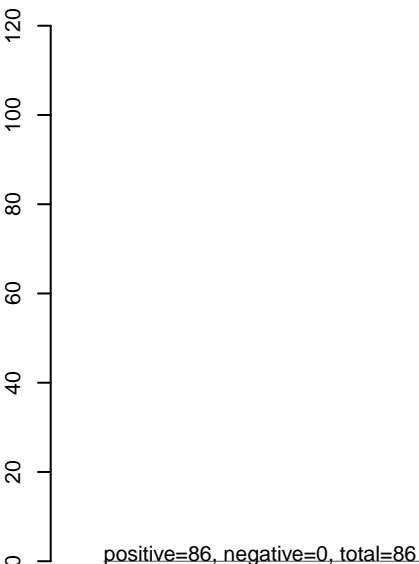
AeAeg_Aag2_DENV1.rep



AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

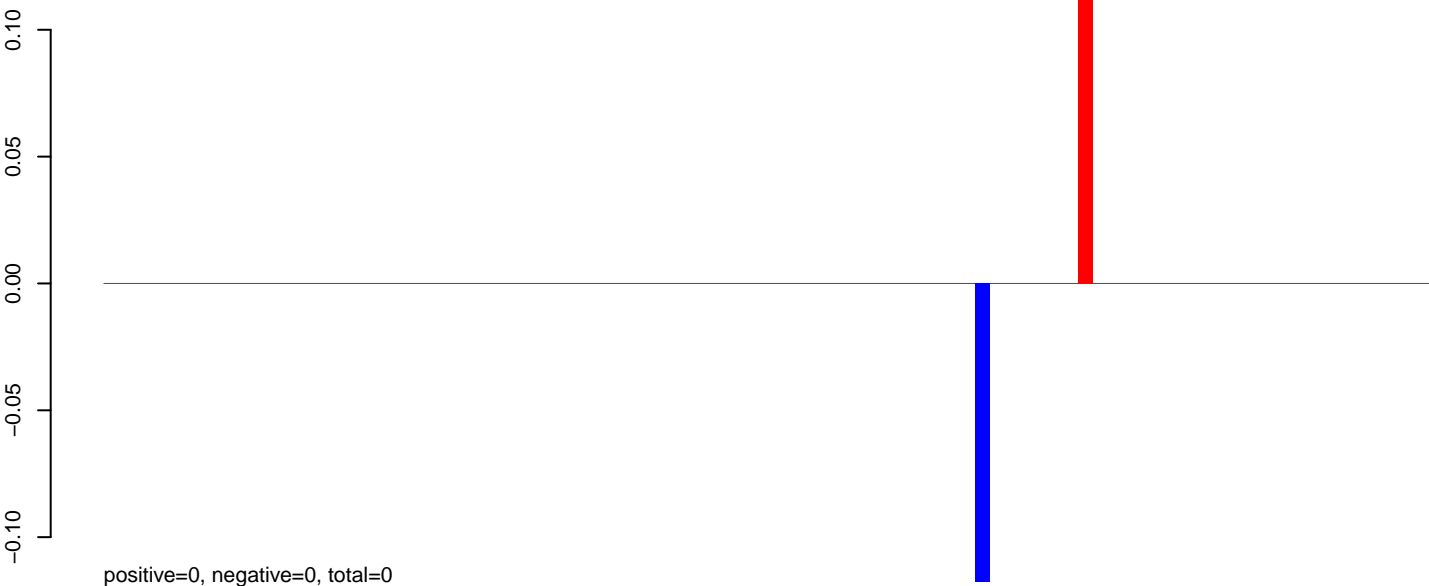


AeAeg_Aag2_DENV1.rep

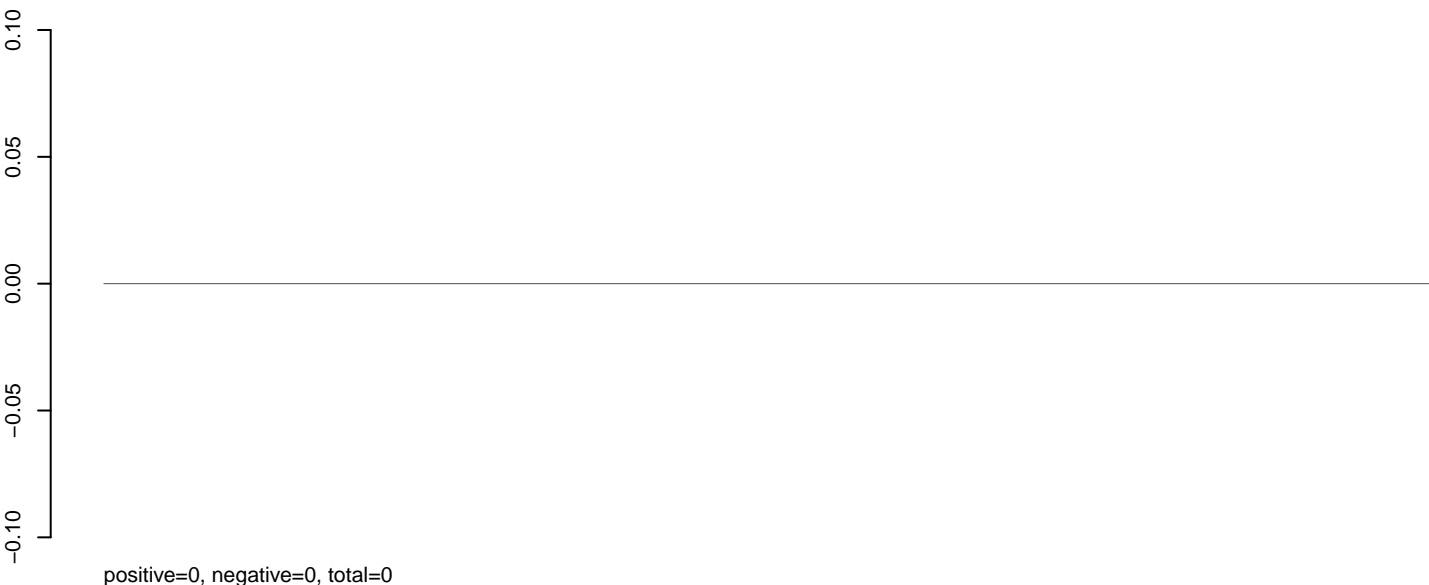


0 2000 4000 6000 8000 10000 12000

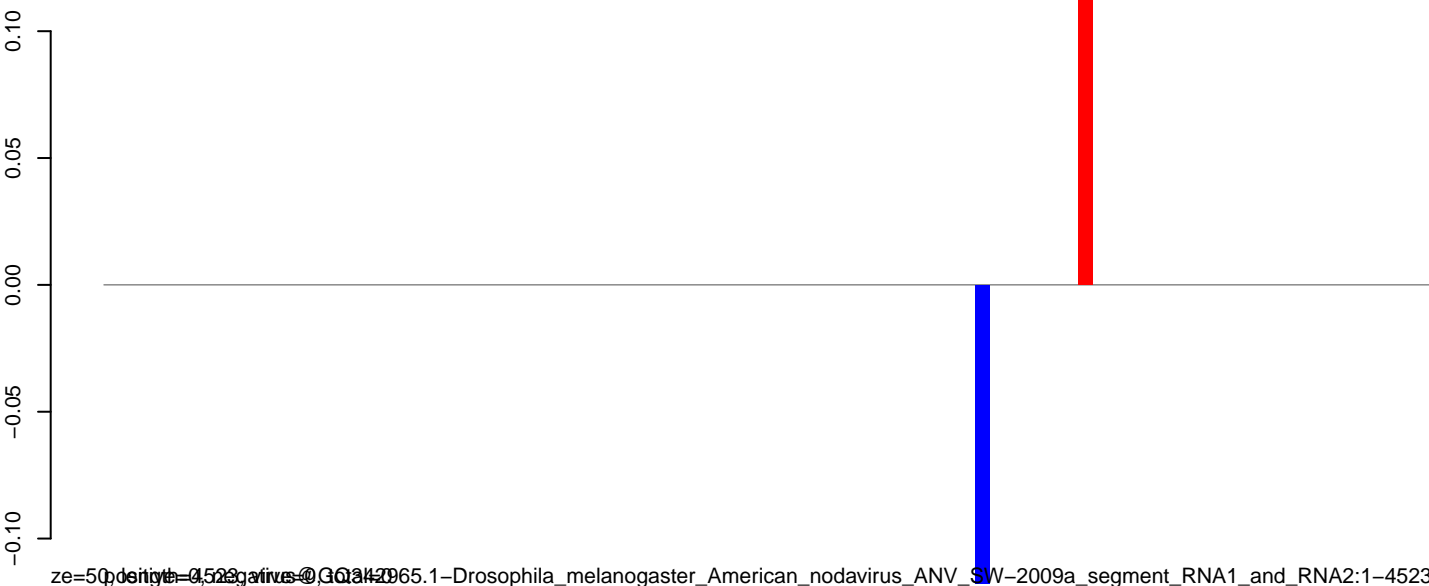
AeAeg_Aag2_DENV1.18_23.rep



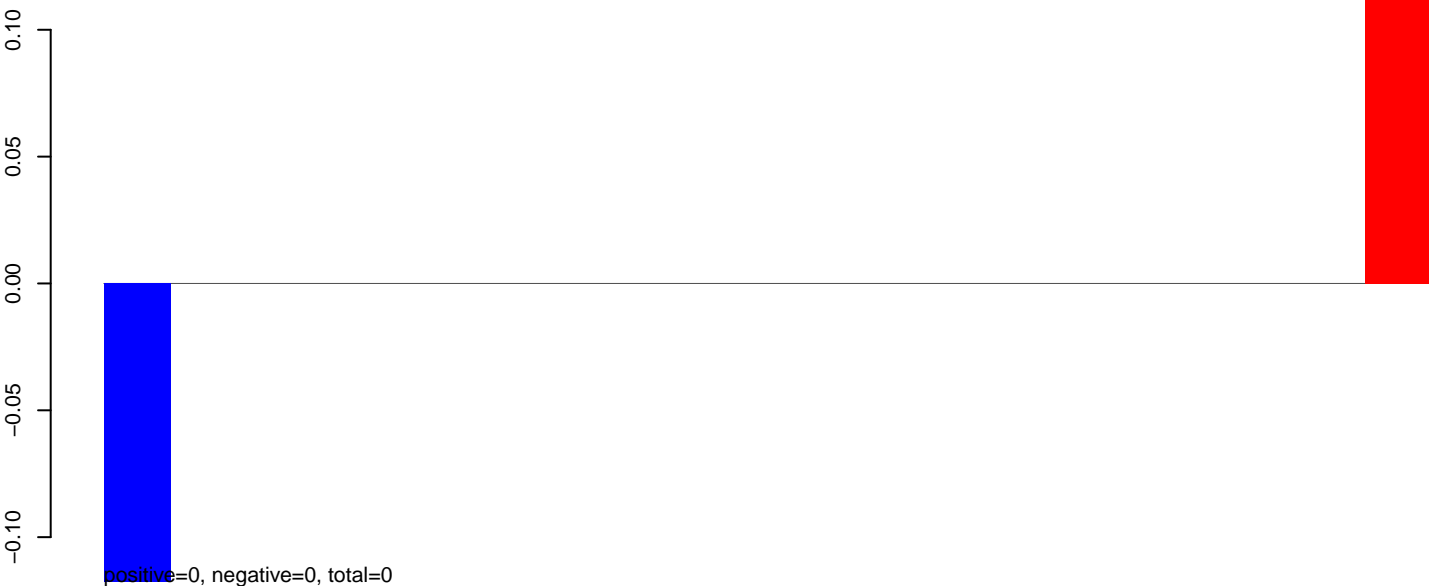
AeAeg_Aag2_DENV1.24_35.rep



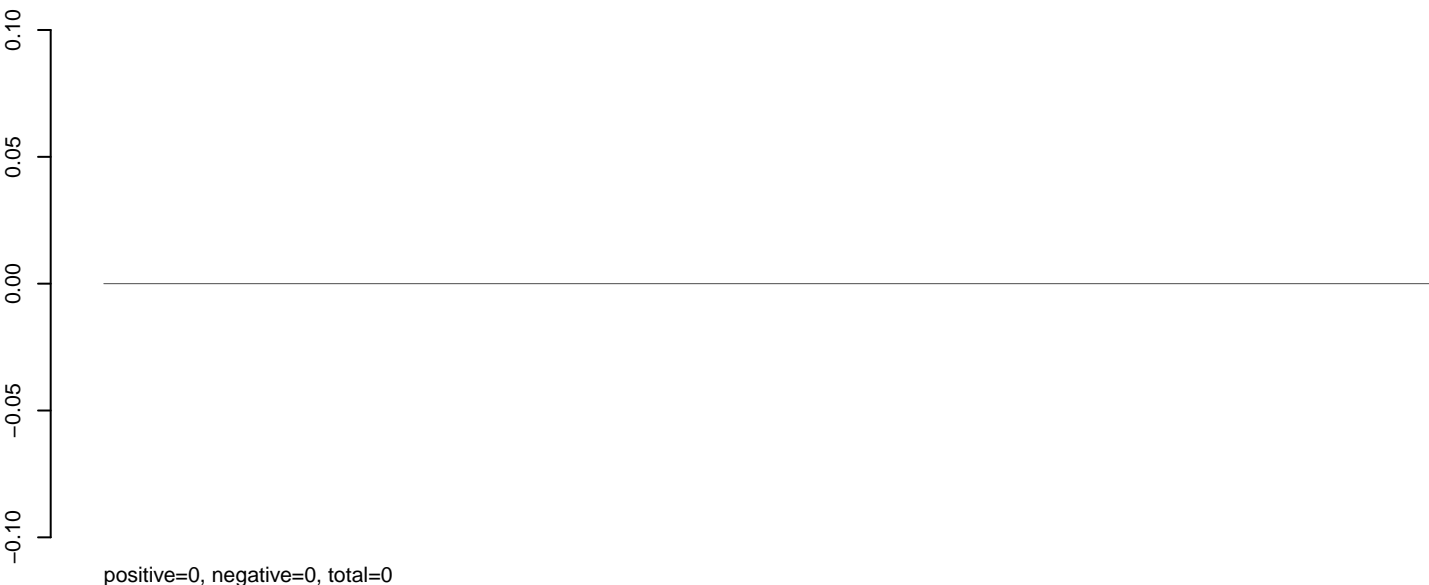
AeAeg_Aag2_DENV1.rep



AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



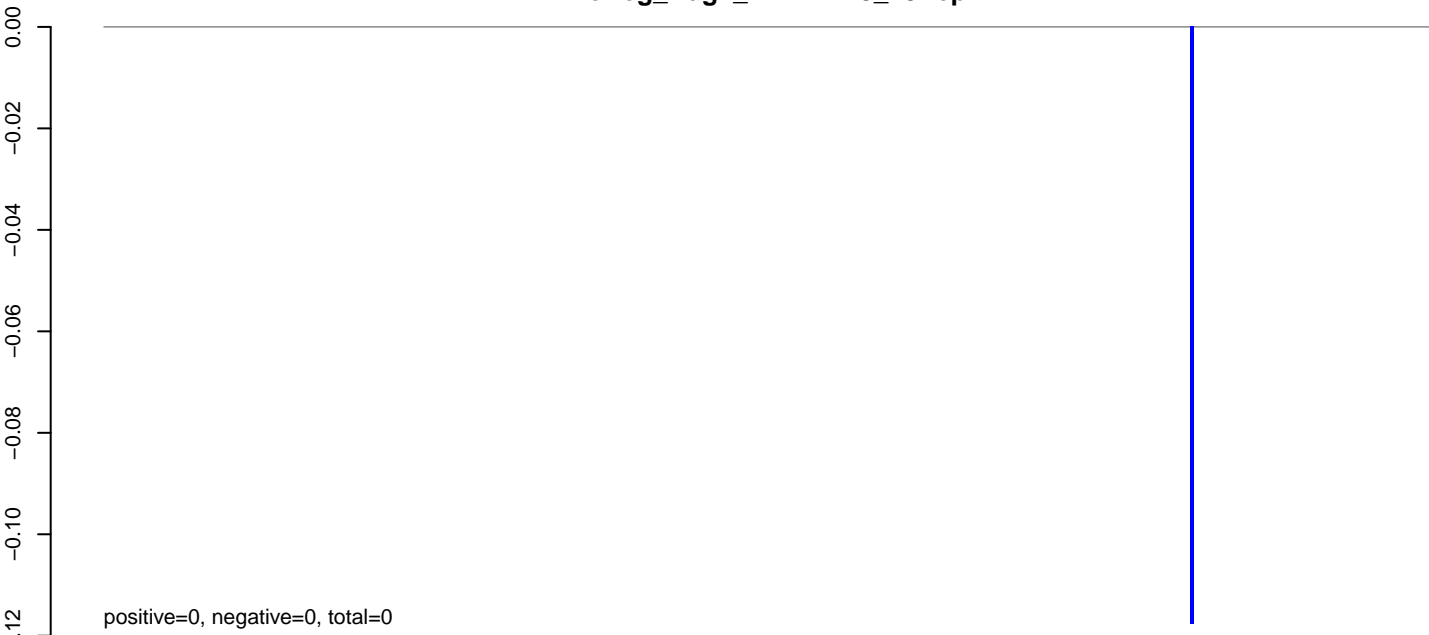
AeAeg_Aag2_DENV1.rep



positive=0, negative=0, total=0

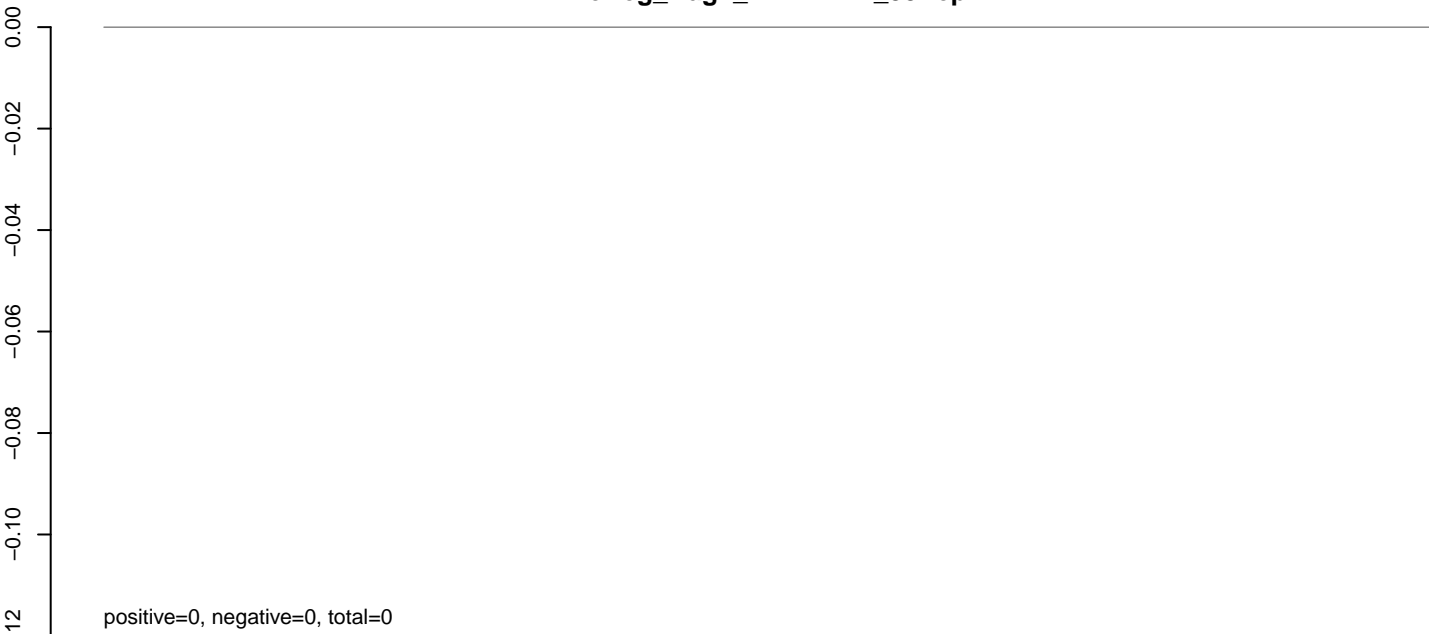
200 400 600 800 1000

AeAeg_Aag2_DENV1.18_23.rep



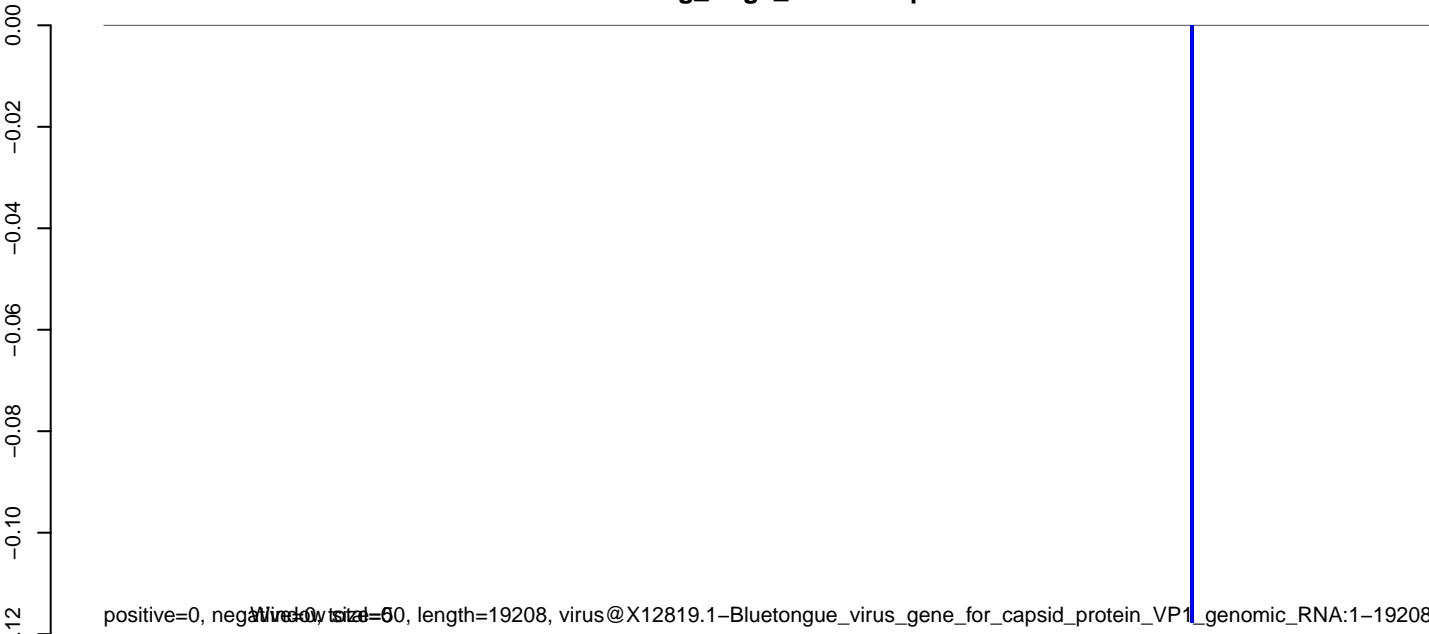
positive=0, negative=0, total=0

AeAeg_Aag2_DENV1.24_35.rep



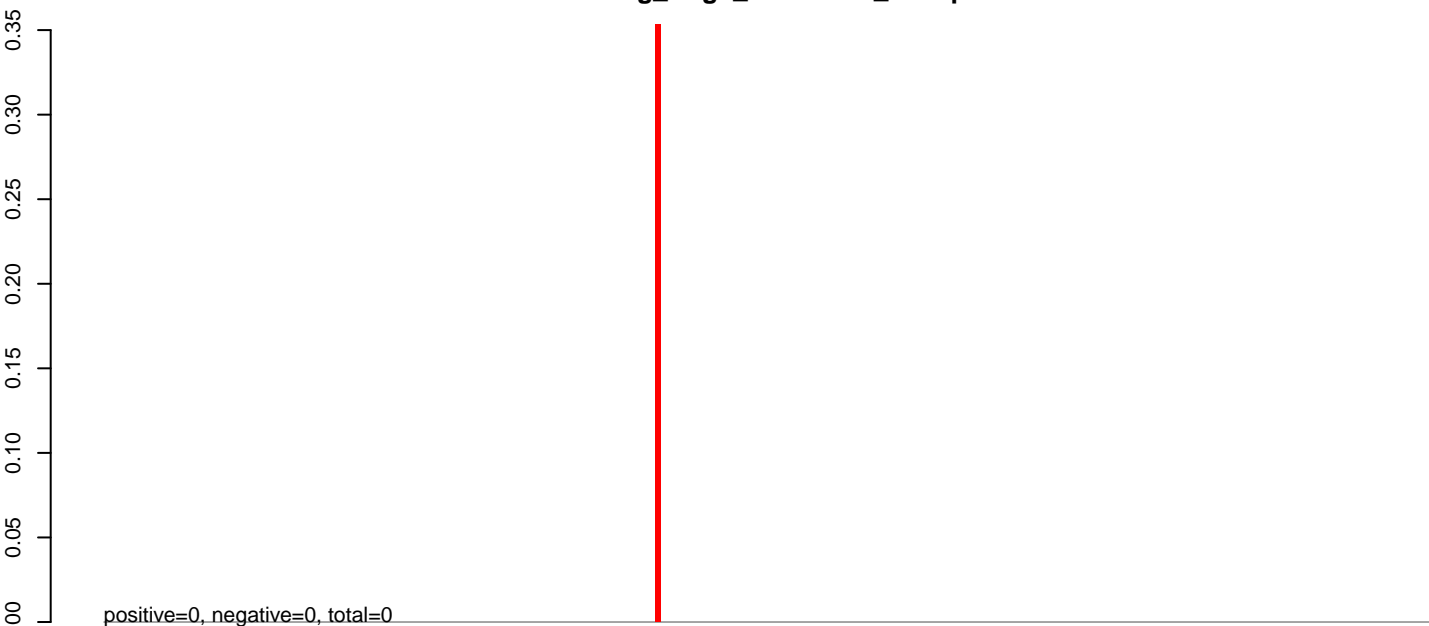
positive=0, negative=0, total=0

AeAeg_Aag2_DENV1.rep

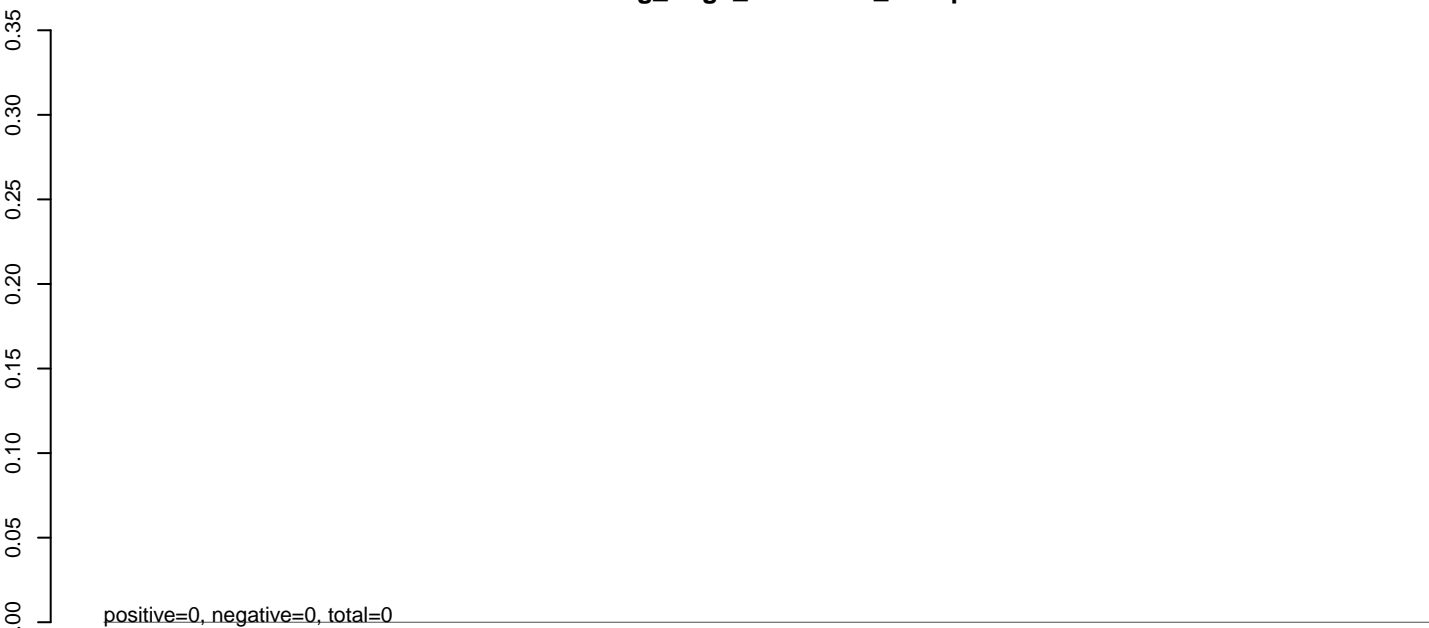


positive=0, negative=0, total=0, size=60, length=19208, virus@X12819.1-Bluetongue_virus_gene_for_capsid_protein_VP1_genomic_RNA:1-19208

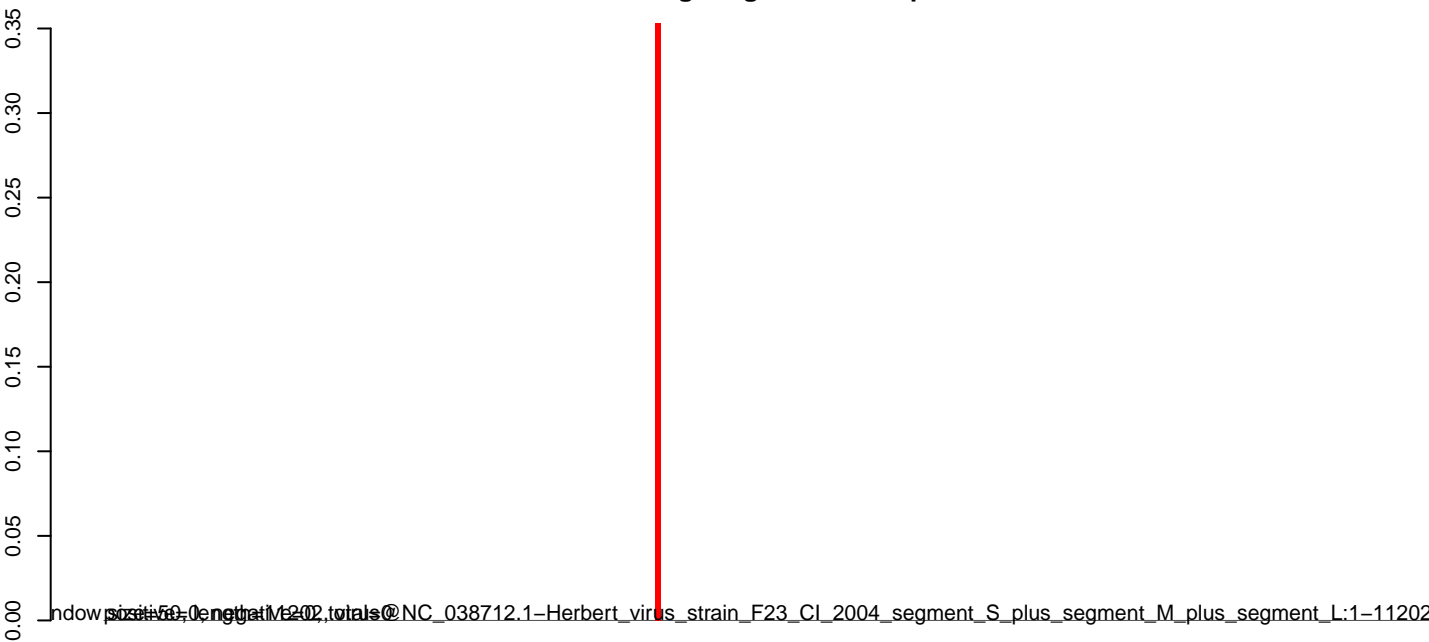
AeAeg_Aag2_DENV1.18_23.rep



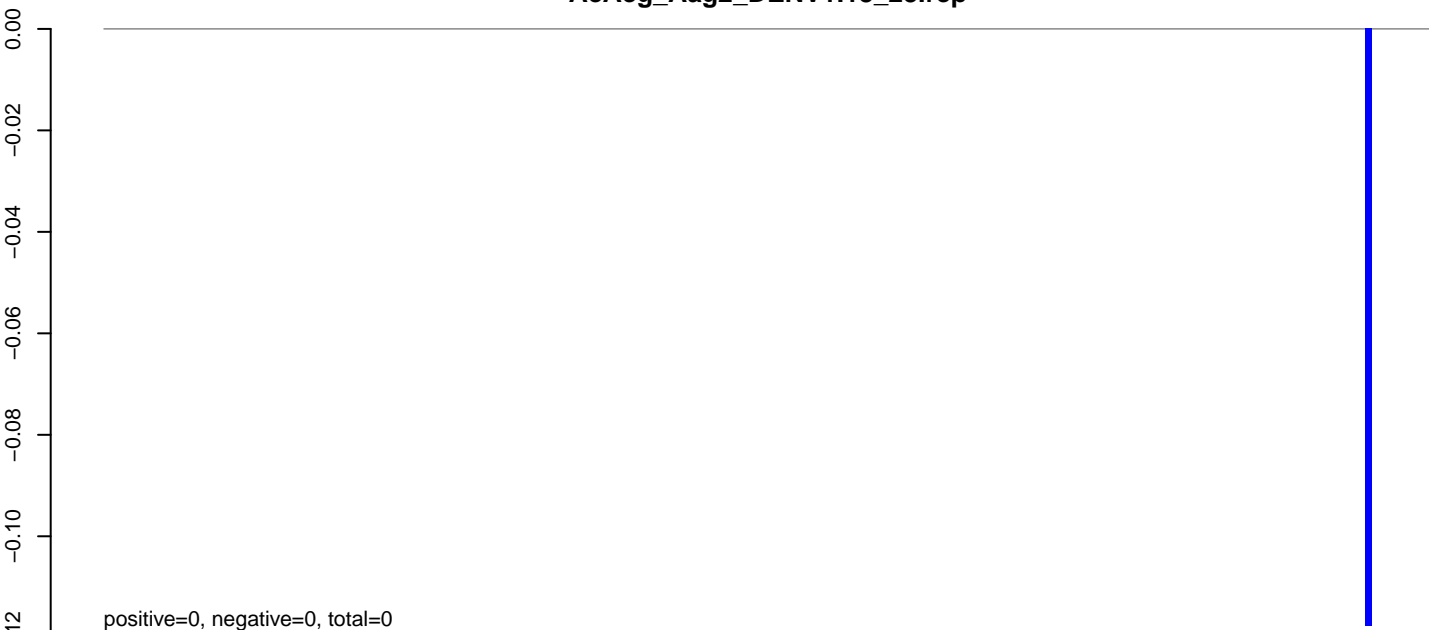
AeAeg_Aag2_DENV1.24_35.rep



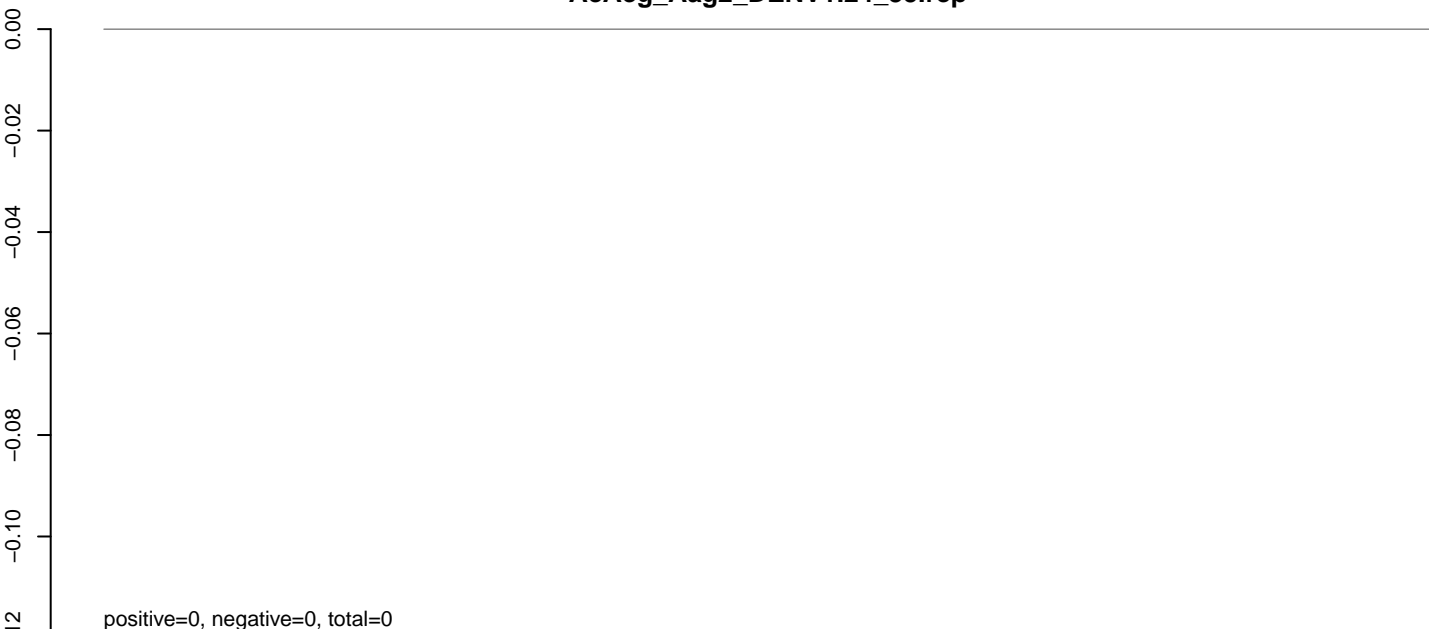
AeAeg_Aag2_DENV1.rep



AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



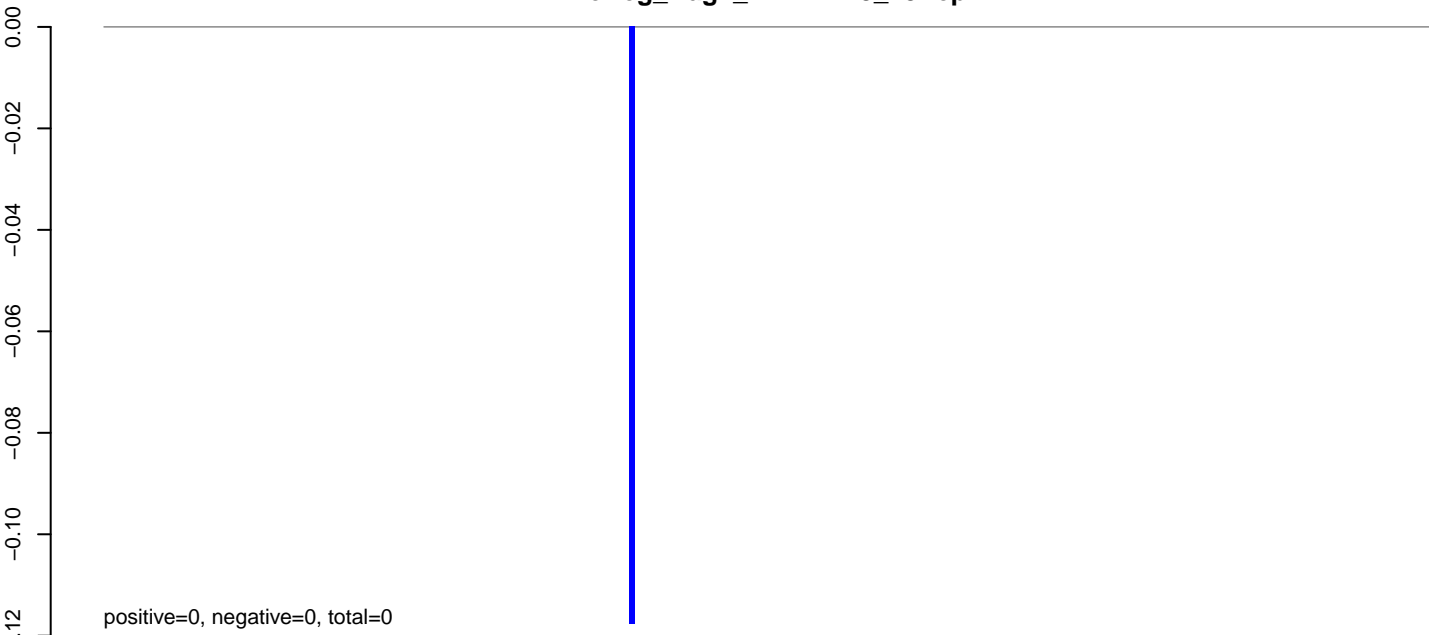
AeAeg_Aag2_DENV1.rep



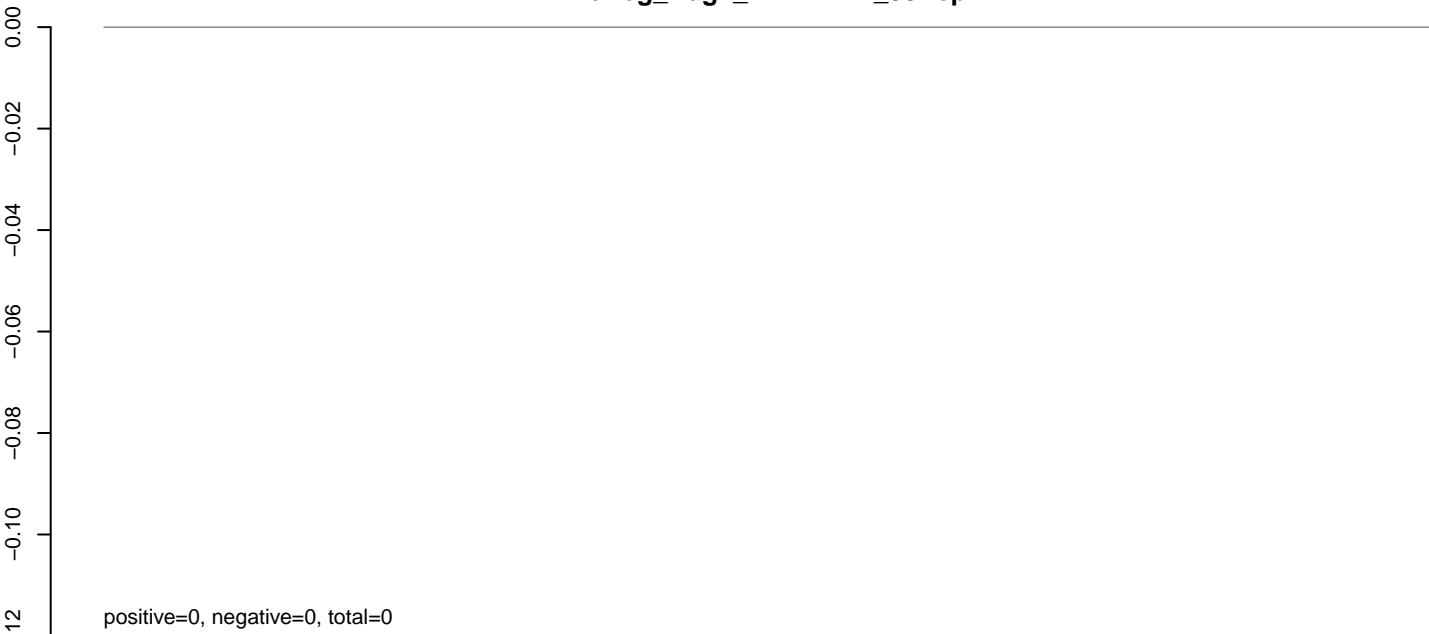
50, length=10968, gene=NC_024639.1-Nkolbisson_virus_nucleoprotein_phosphoprotein_matrix_glycoprotein_and_polymerase_genes:1-10968

0 2000 4000 6000 8000 10000

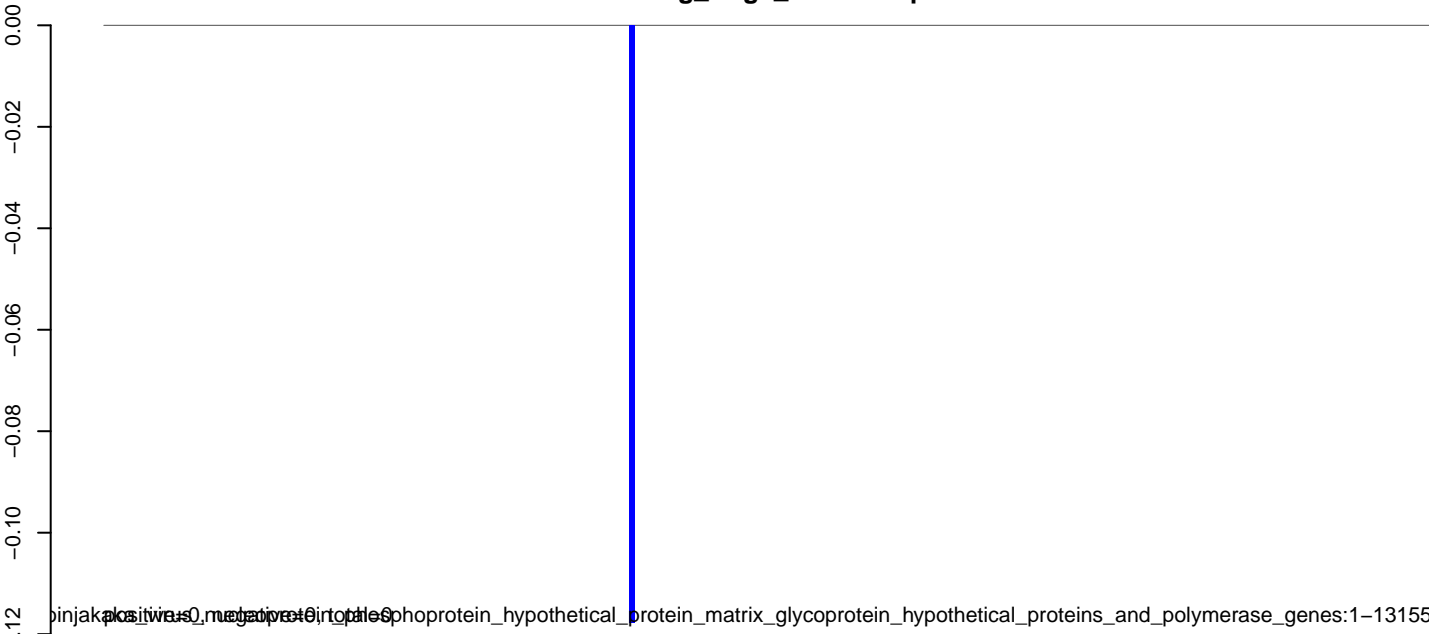
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

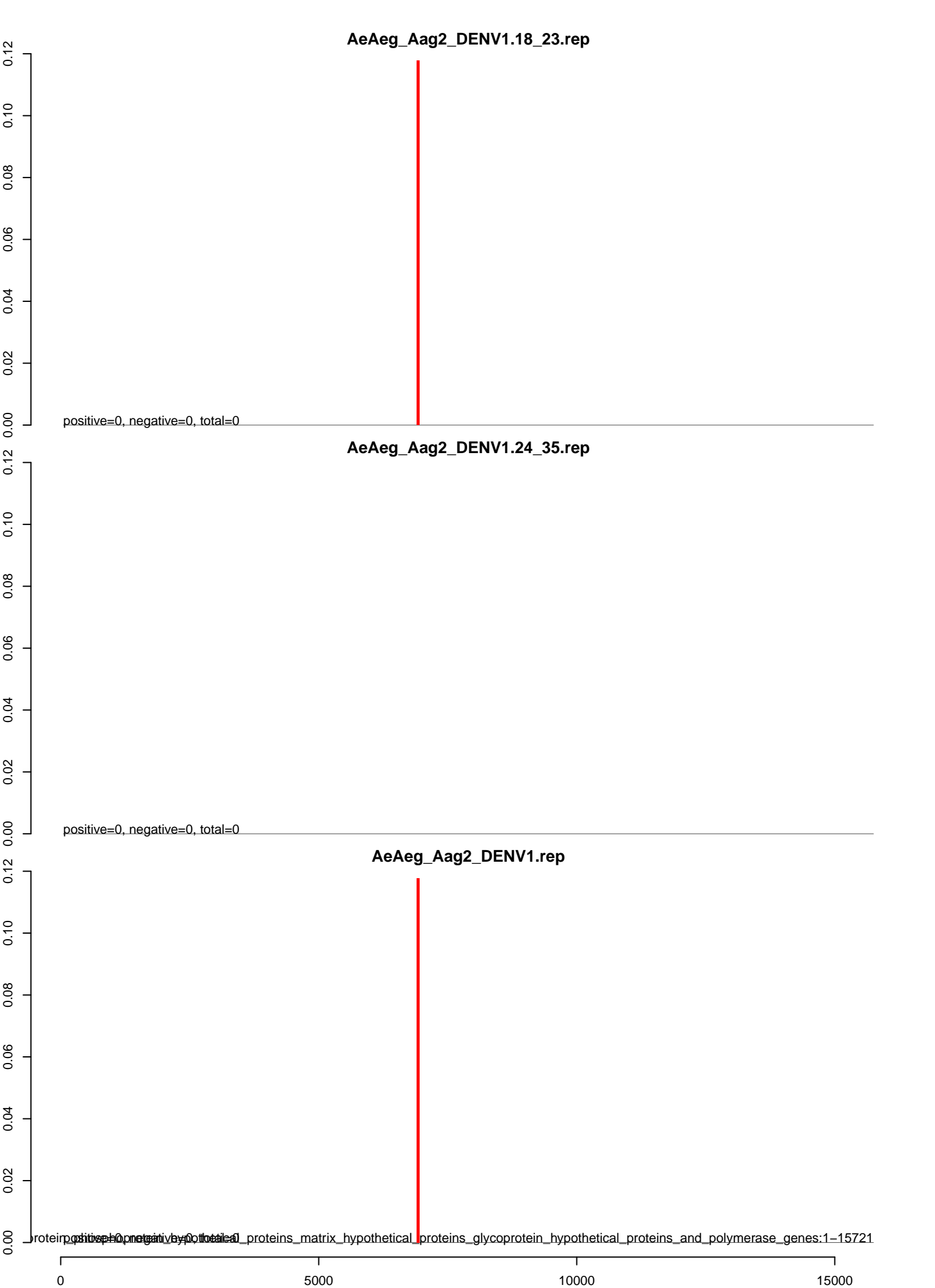


AeAeg_Aag2_DENV1.rep

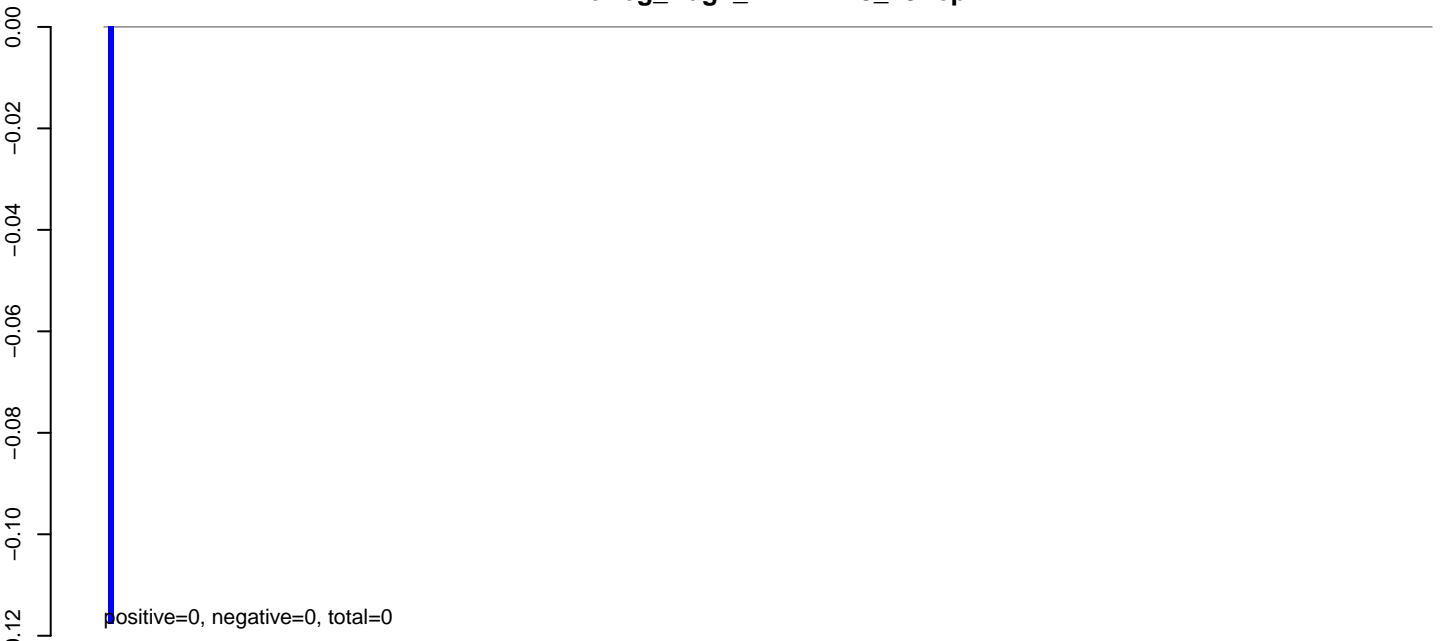


binjakas... phosphoprotein_hypothetical_protein_matrix_glycoprotein_hypothetical_proteins_and_polymerase_genes:1-13155

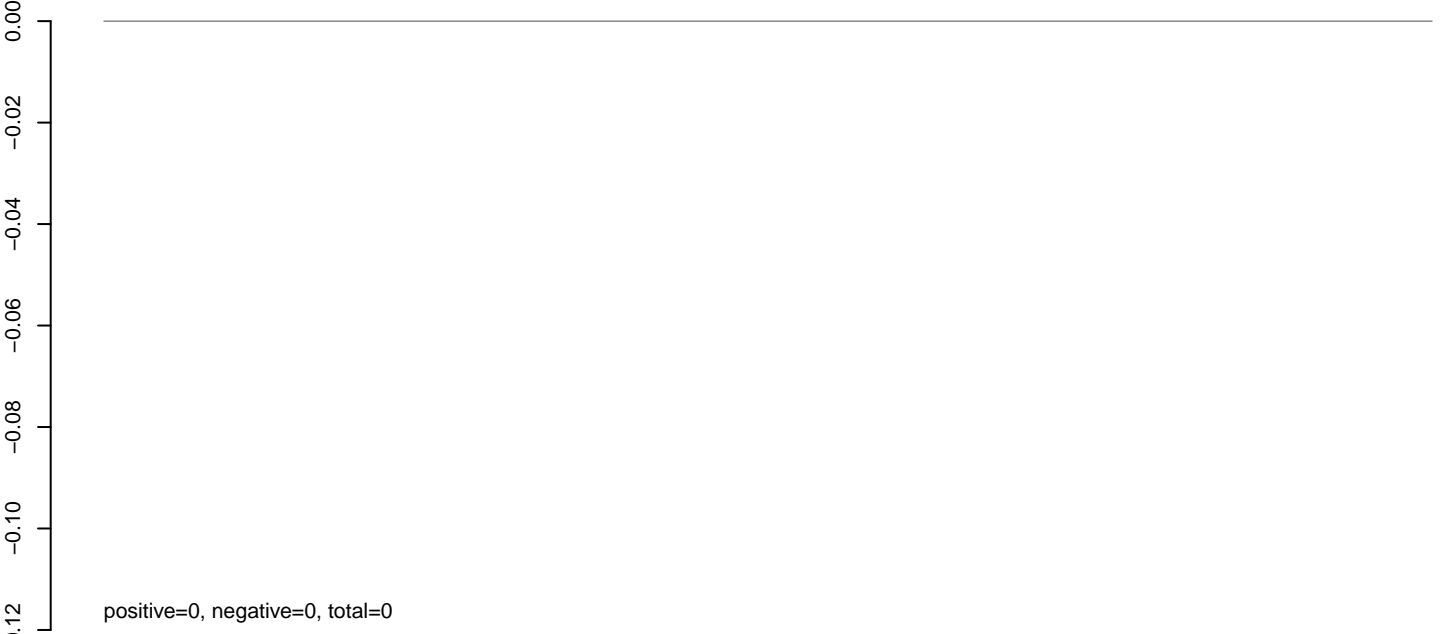
0 2000 4000 6000 8000 10000 12000



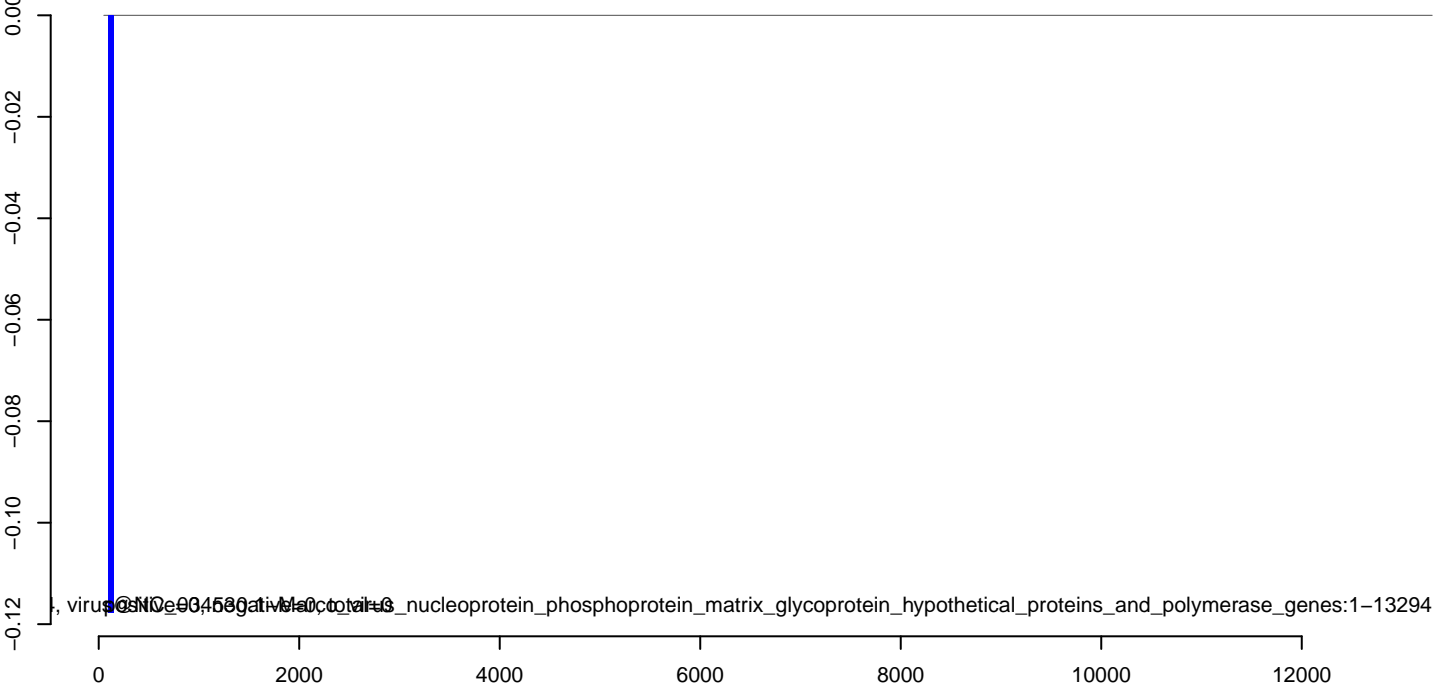
AeAeg_Aag2_DENV1.18_23.rep



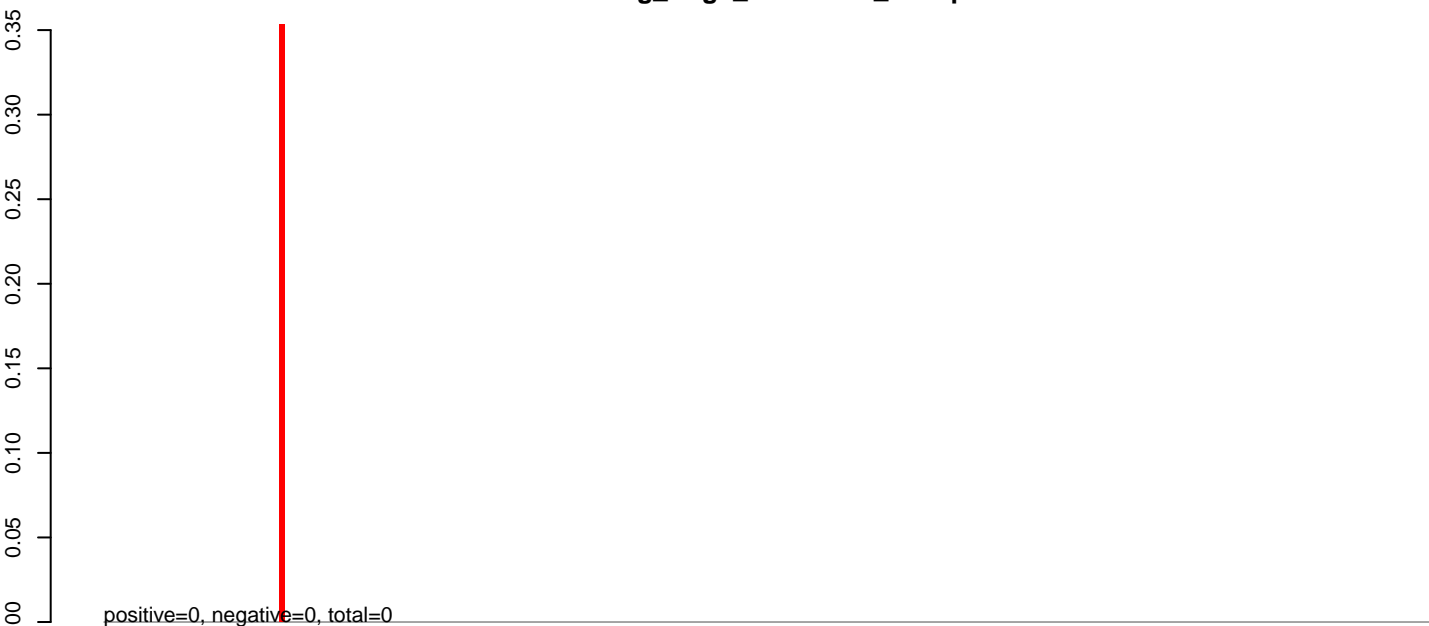
AeAeg_Aag2_DENV1.24_35.rep



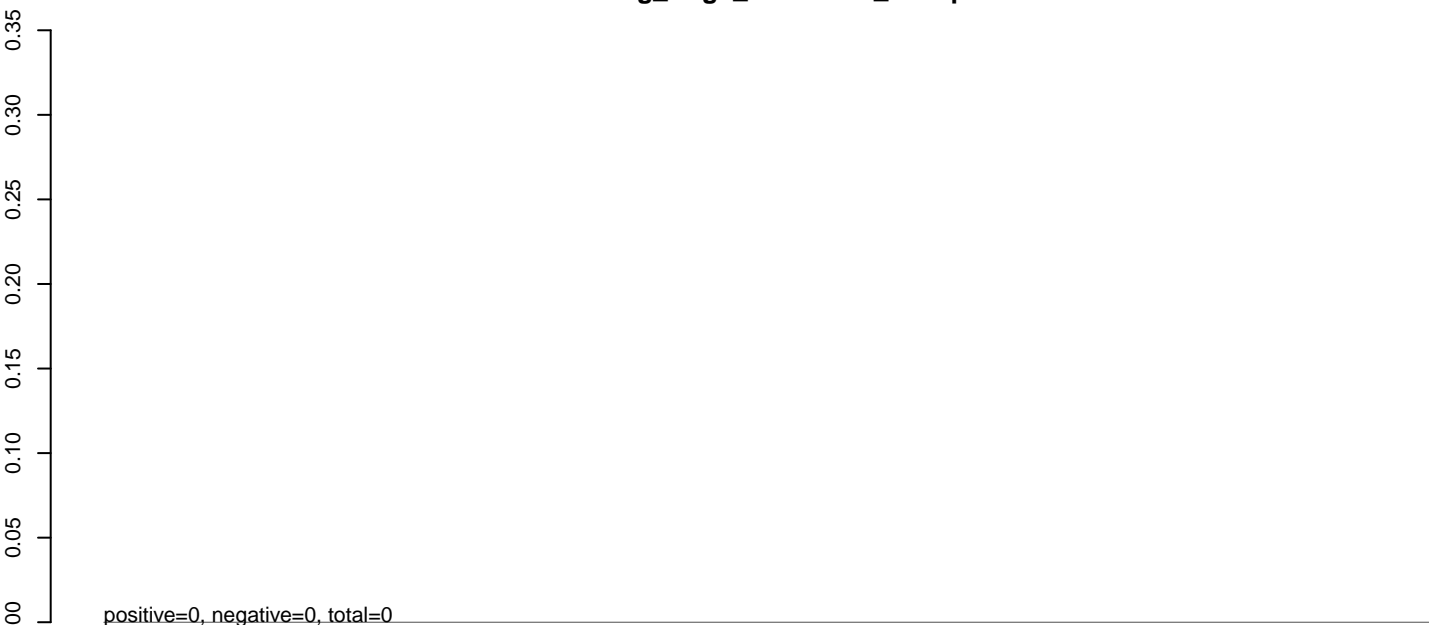
AeAeg_Aag2_DENV1.rep



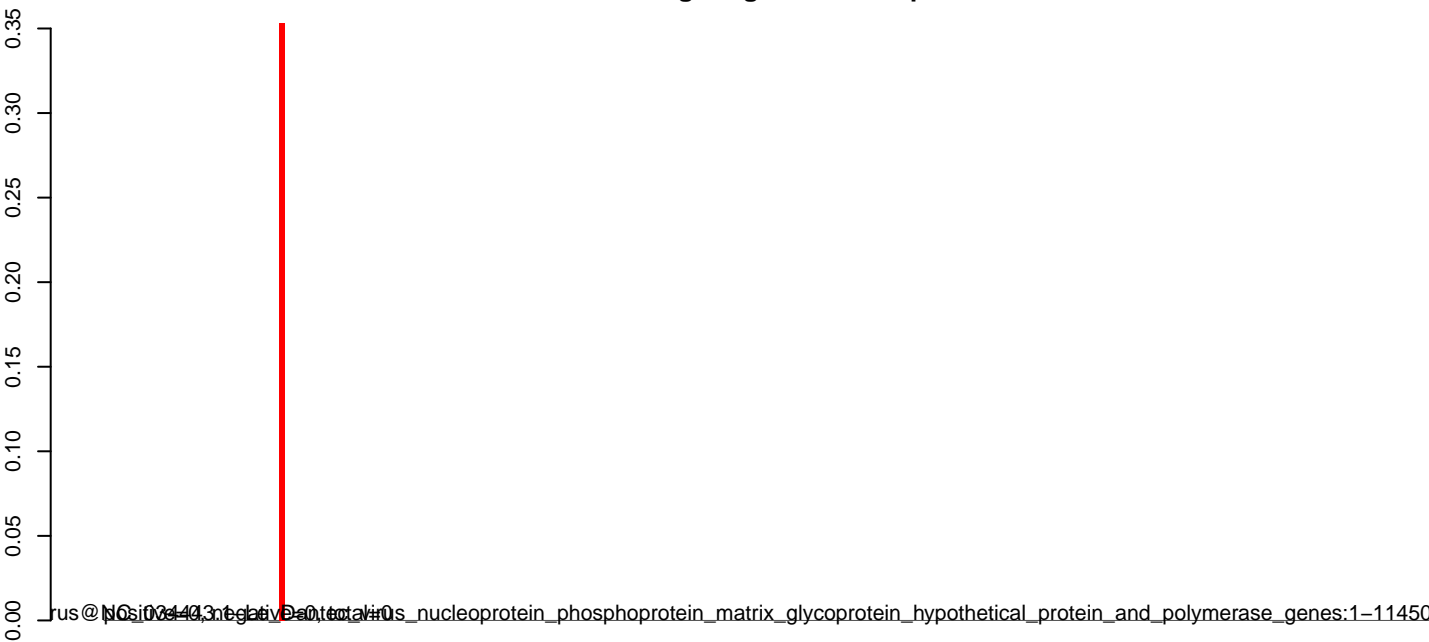
AeAeg_Aag2_DENV1.18_23.rep



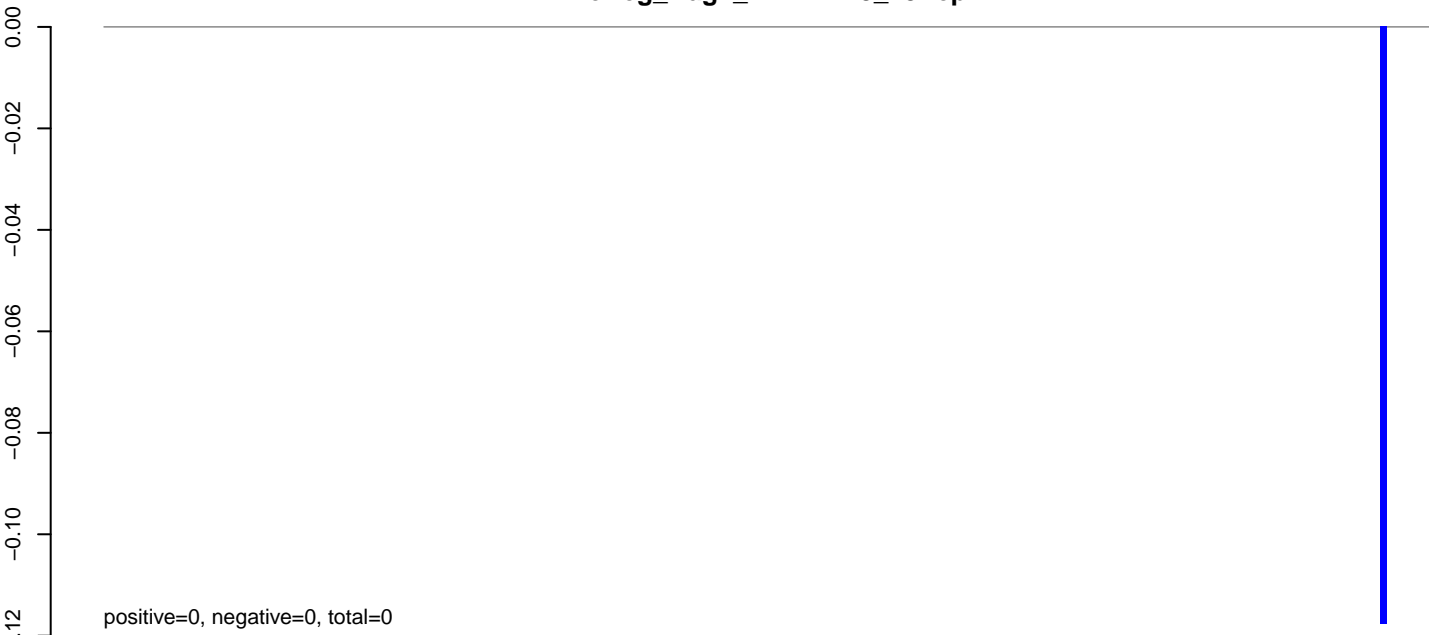
AeAeg_Aag2_DENV1.24_35.rep



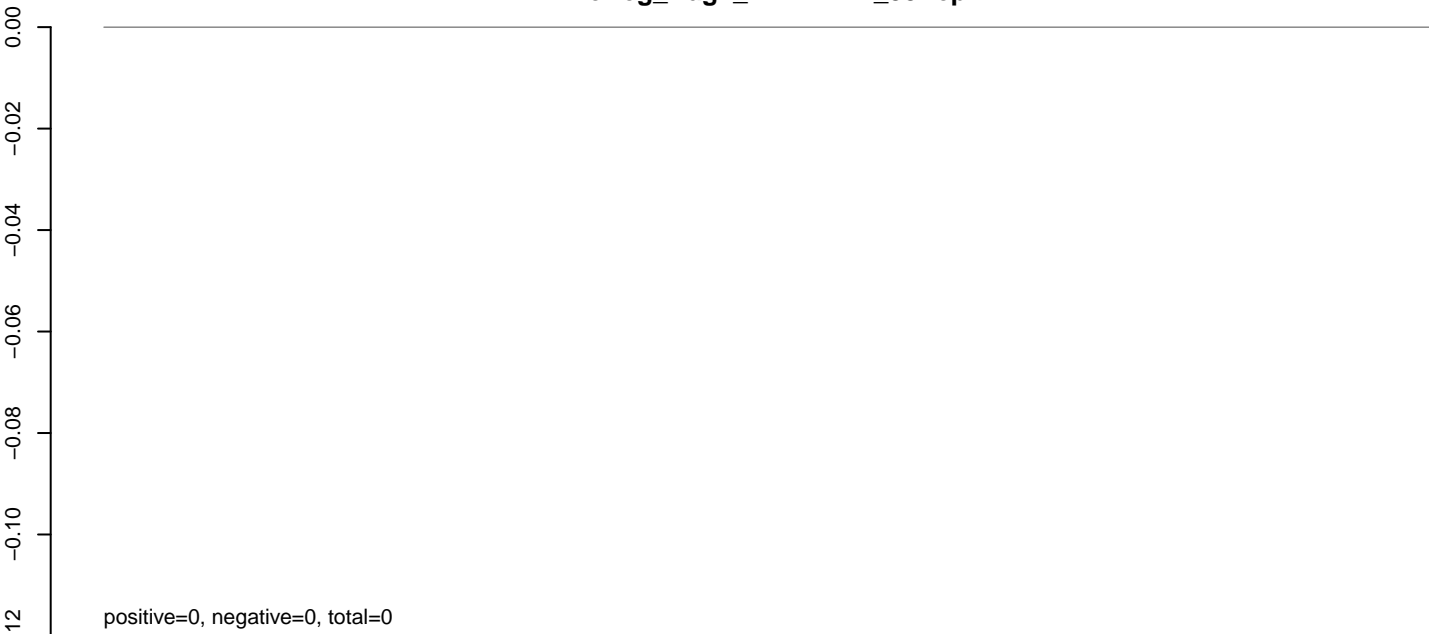
AeAeg_Aag2_DENV1.rep



AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

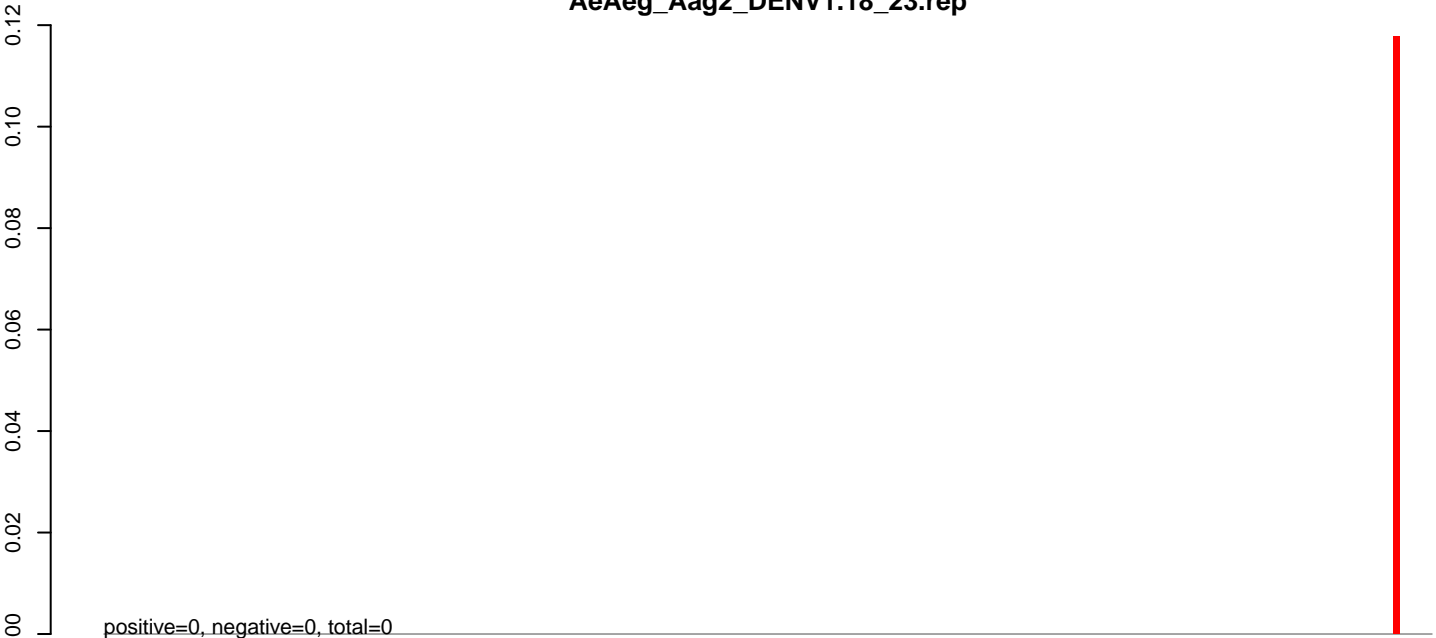


AeAeg_Aag2_DENV1.rep

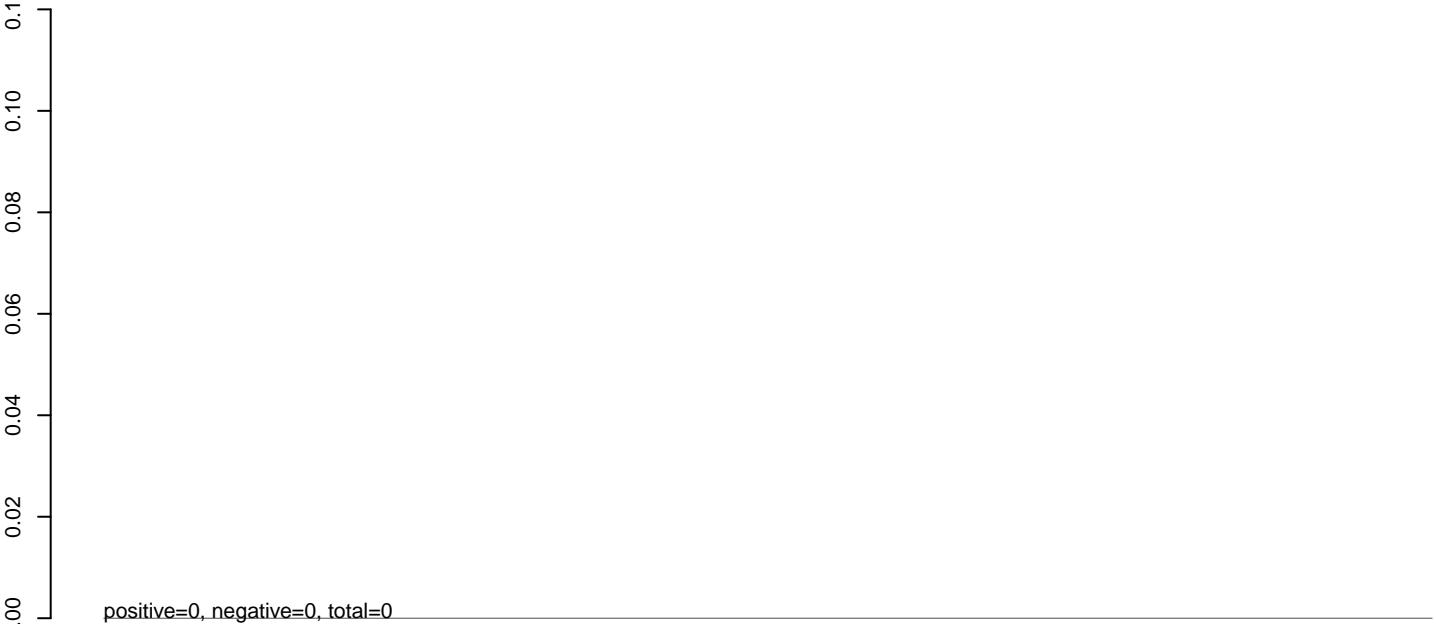


Window size=50, length=10173, virus@NC_033699.1-Jugra_virus_strain_P-9-314_polyprotein_gene:1-10173

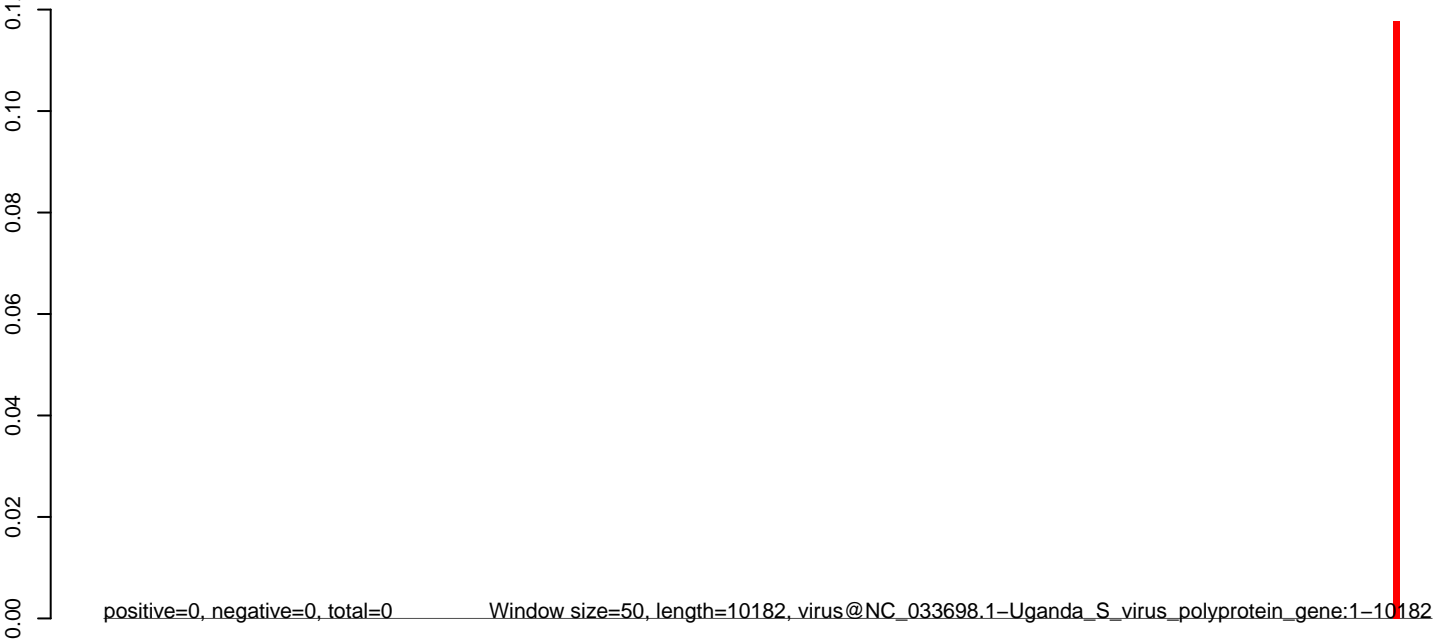
AeAeg_Aag2_DENV1.18_23.rep



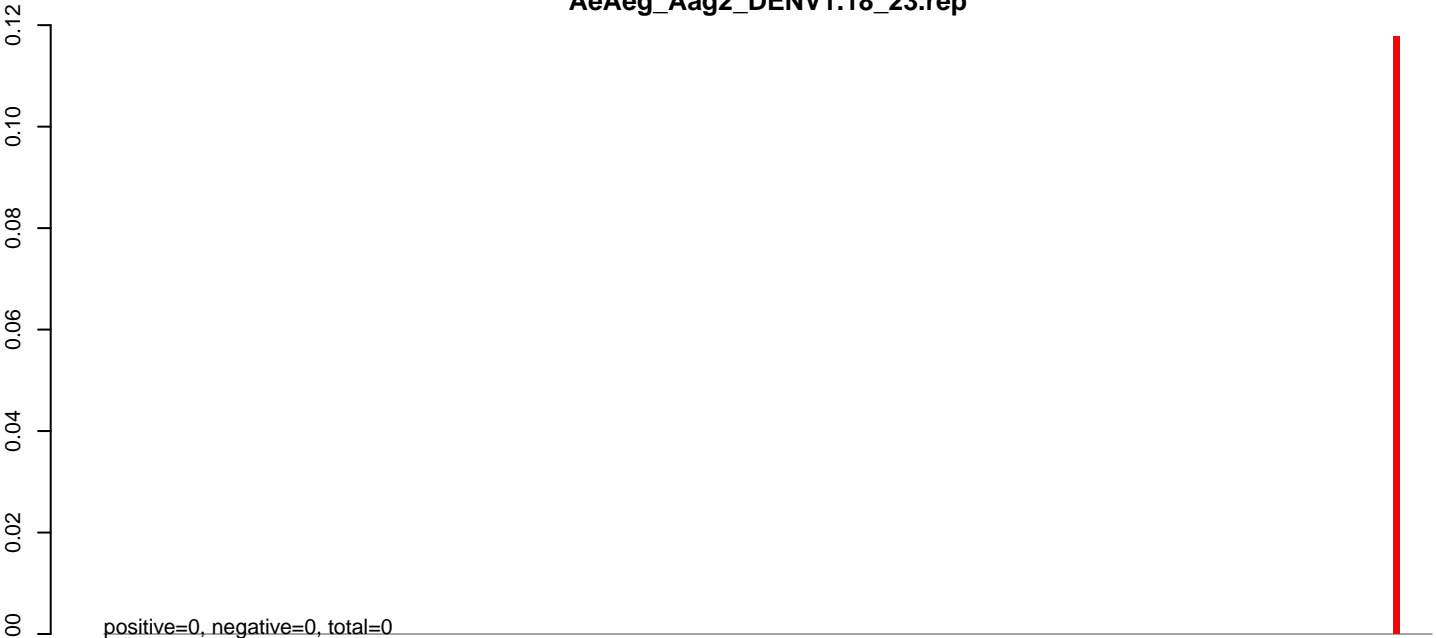
AeAeg_Aag2_DENV1.24_35.rep



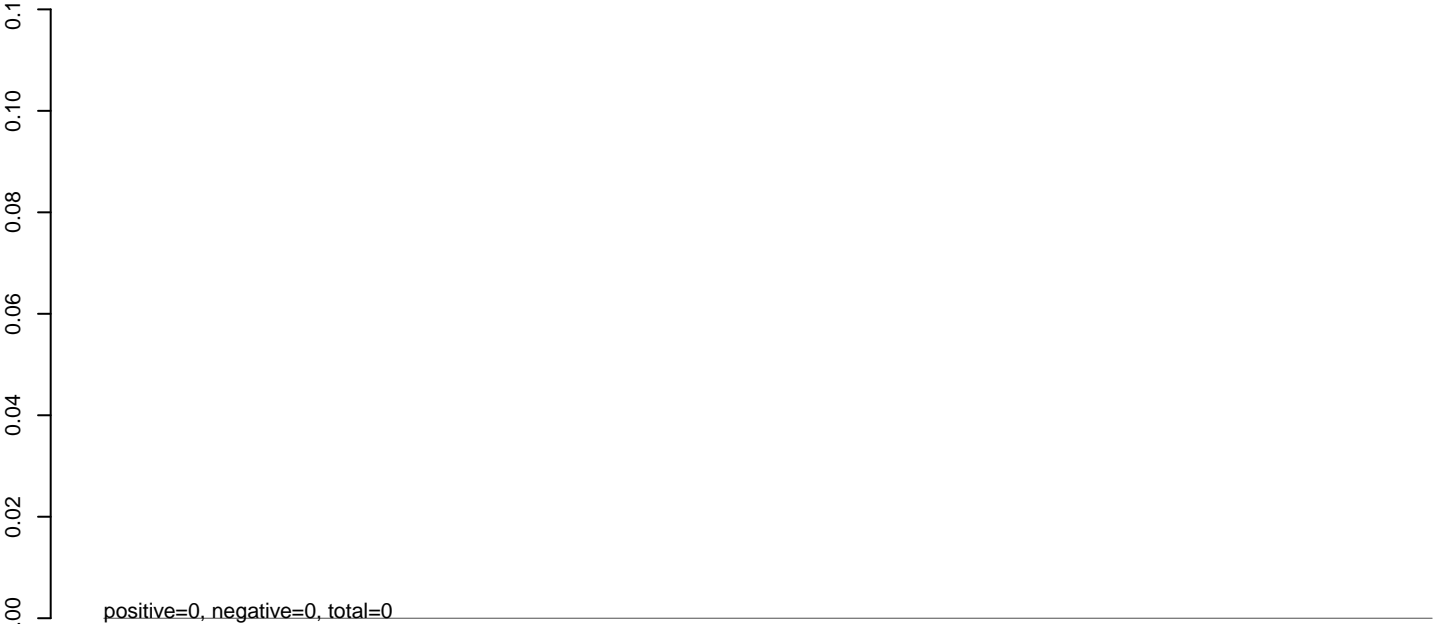
AeAeg_Aag2_DENV1.rep



AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

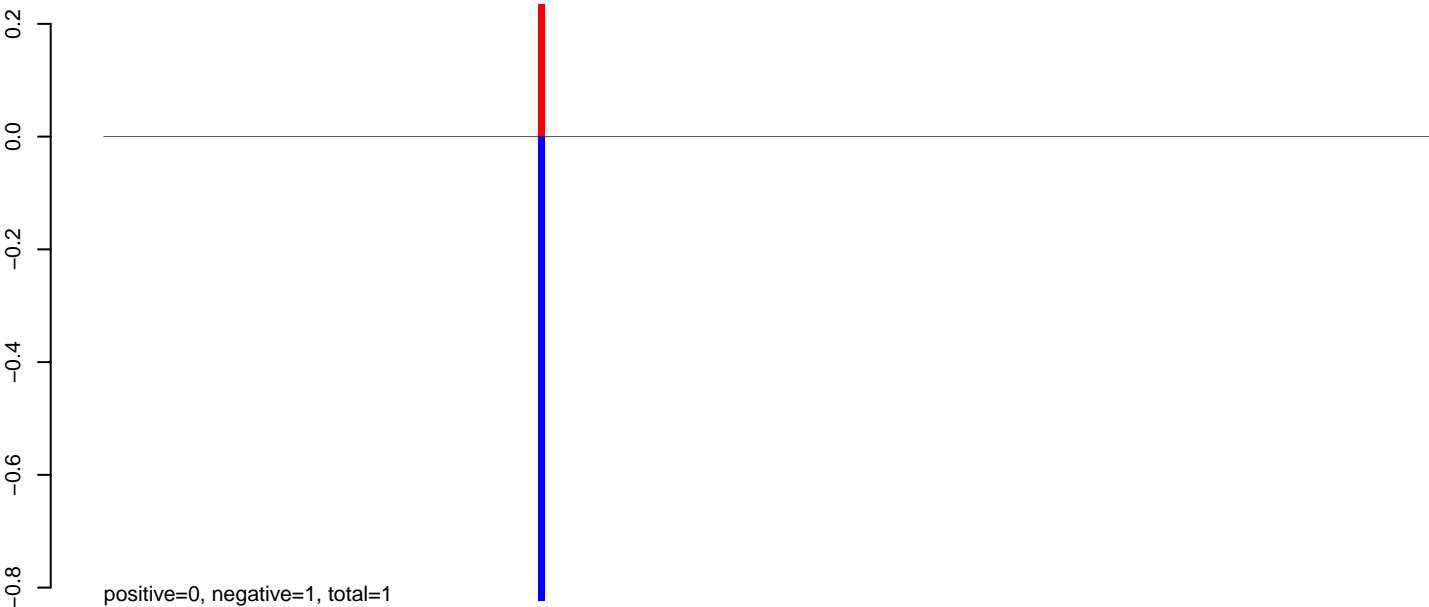


AeAeg_Aag2_DENV1.rep

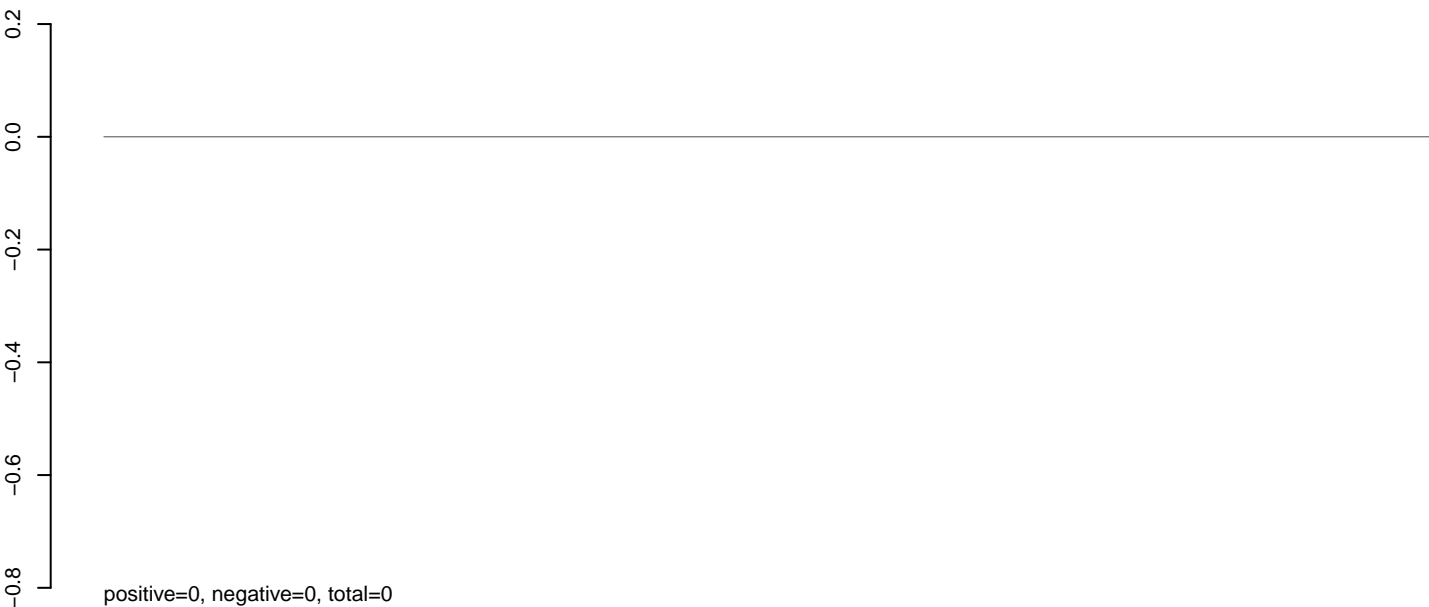


0 2000 4000 6000 8000 10000

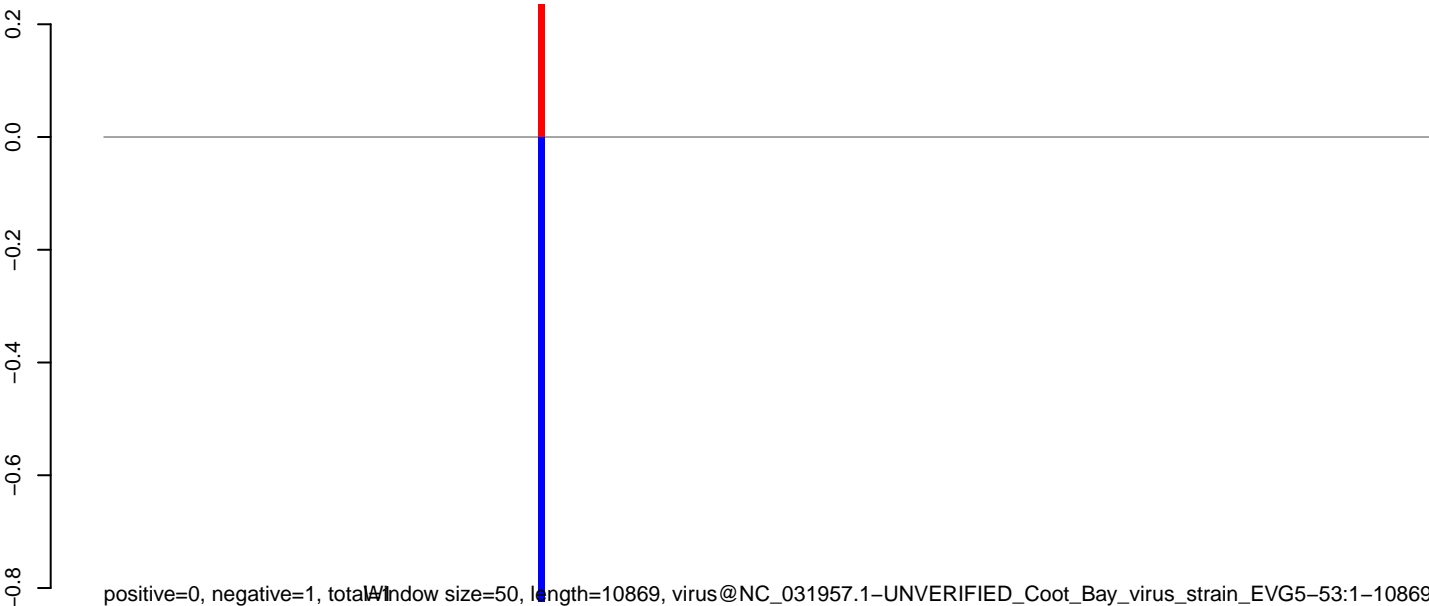
AeAeg_Aag2_DENV1.18_23.rep



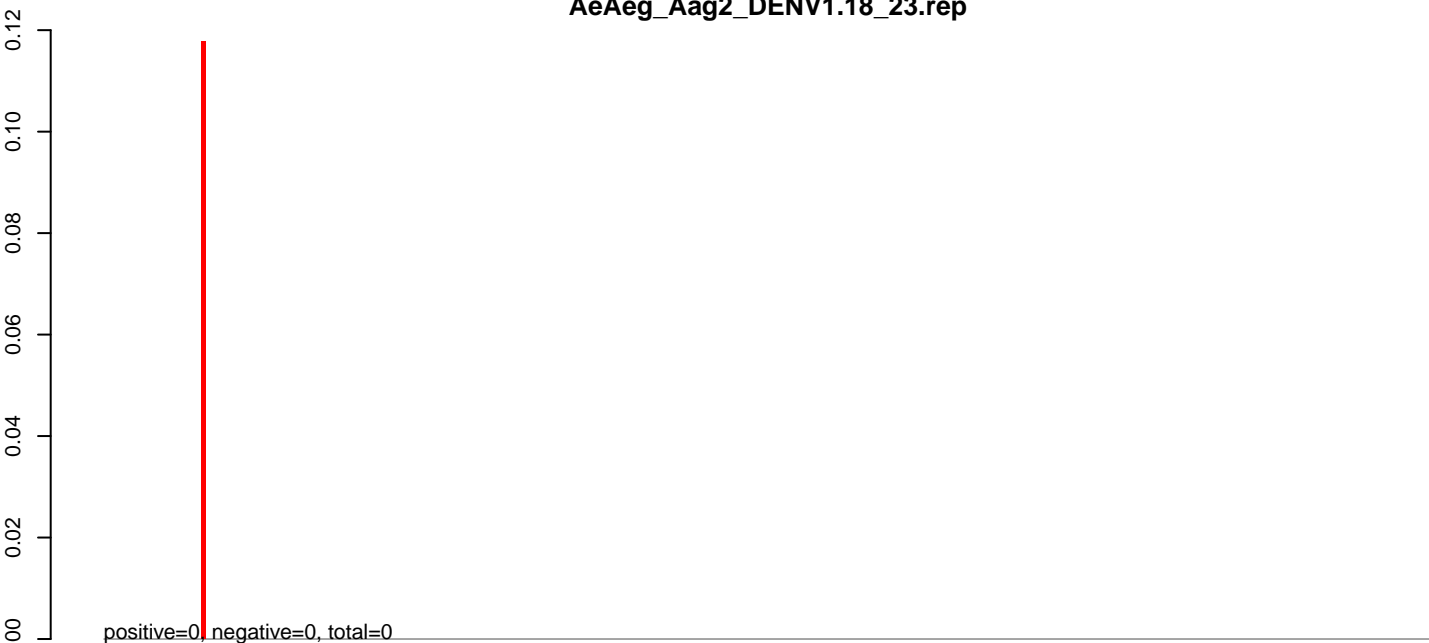
AeAeg_Aag2_DENV1.24_35.rep



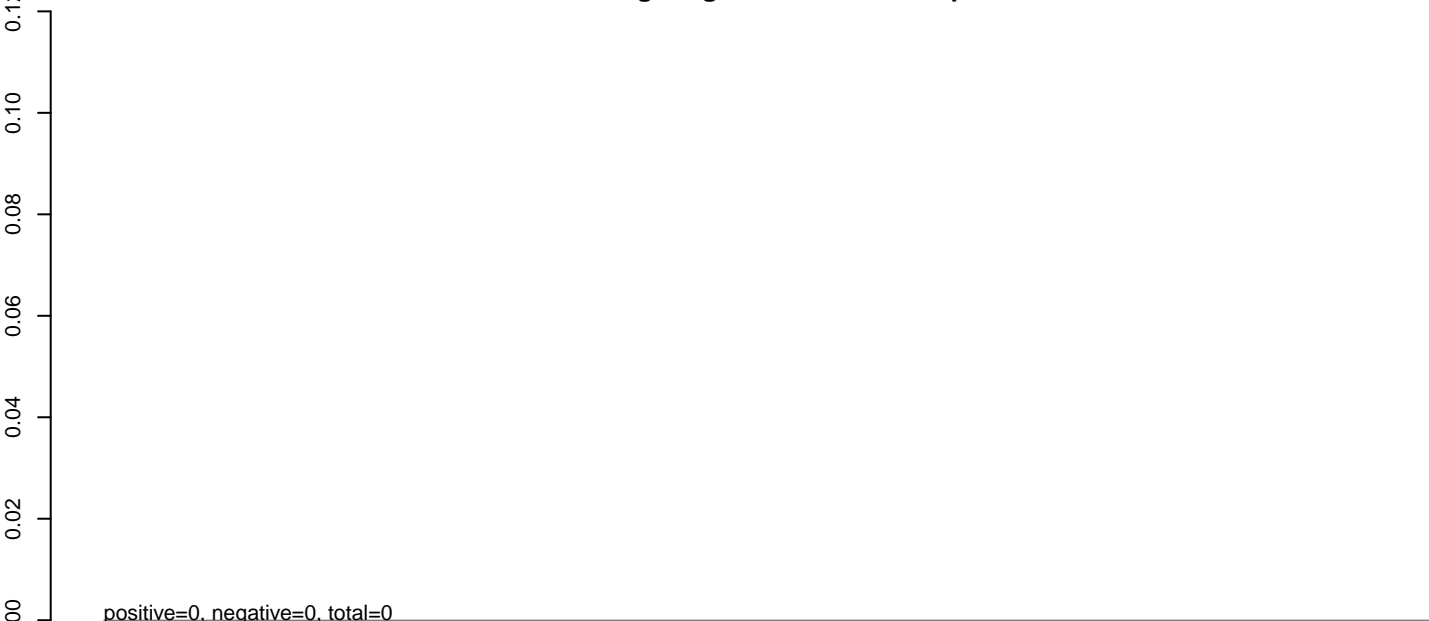
AeAeg_Aag2_DENV1.rep



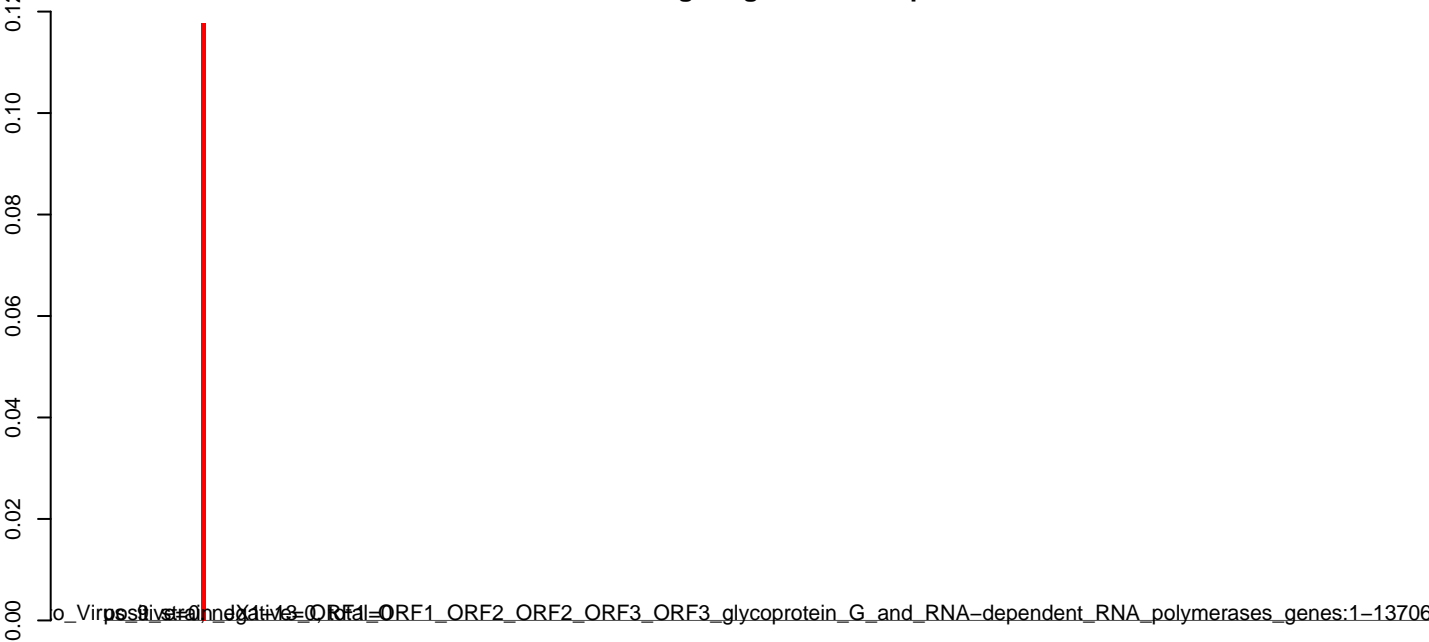
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



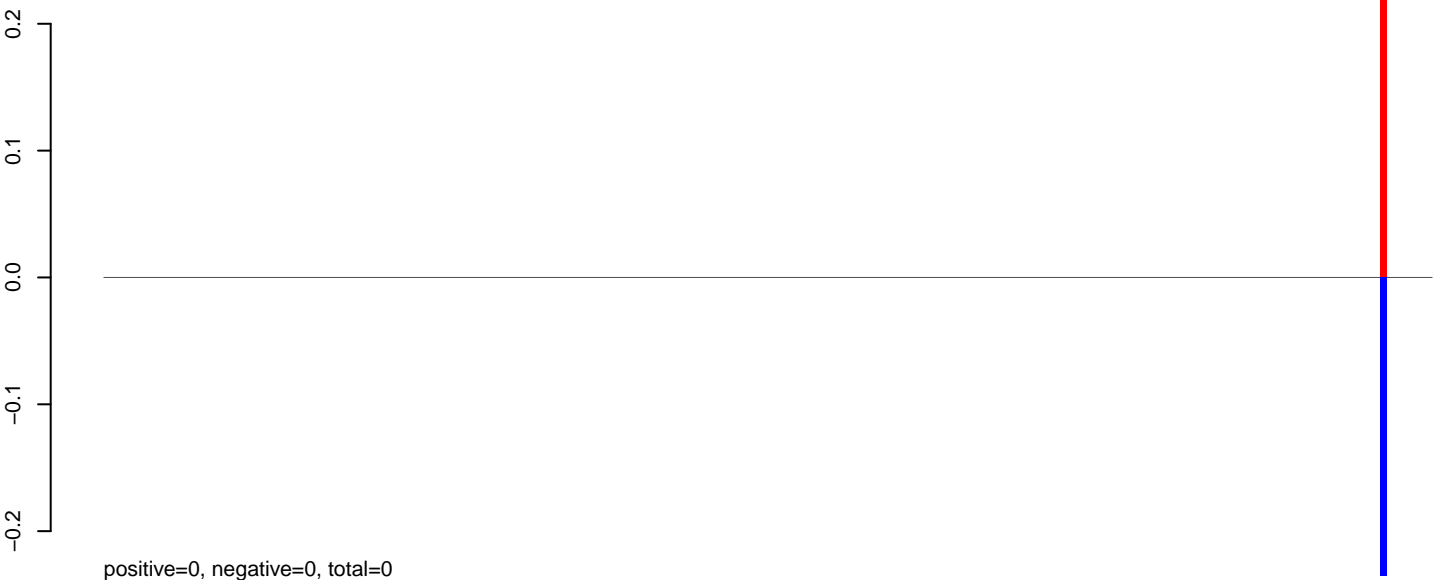
AeAeg_Aag2_DENV1.rep



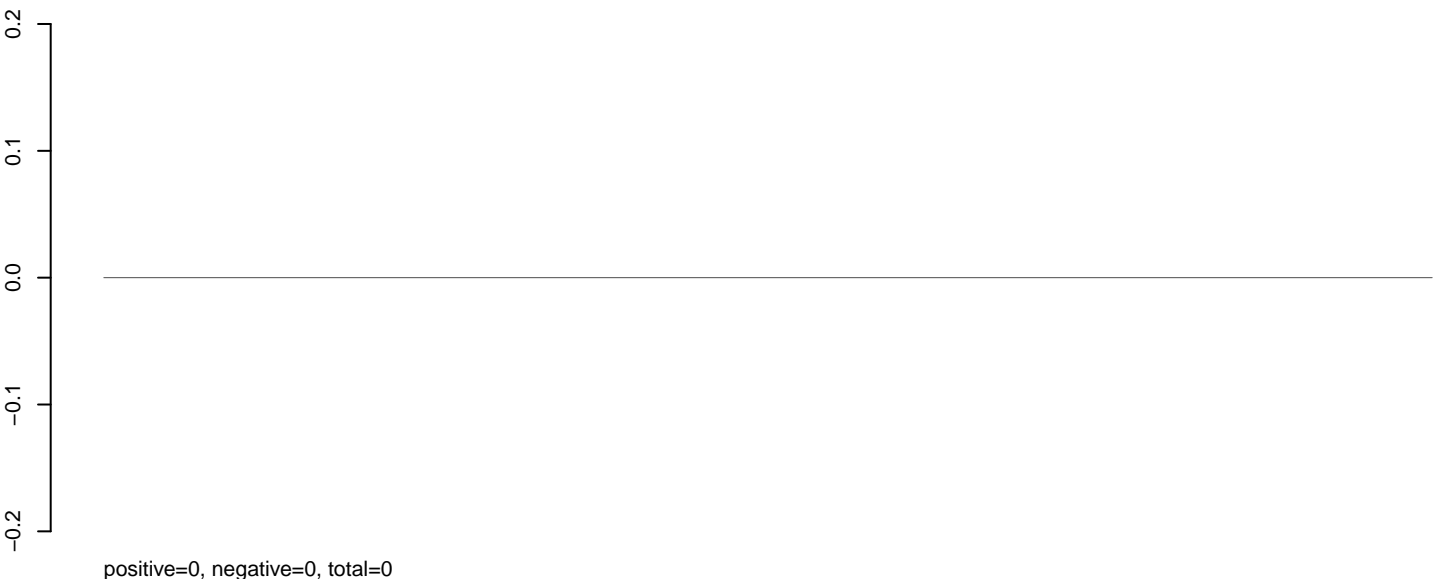
o_Virus_13-ORF1-ORF1-ORF2-ORF2-ORF3-ORF3-glycoprotein_G_and_RNA-dependent_RNA_polymerases_genes:1-13706

0 2000 4000 6000 8000 10000 12000 14000

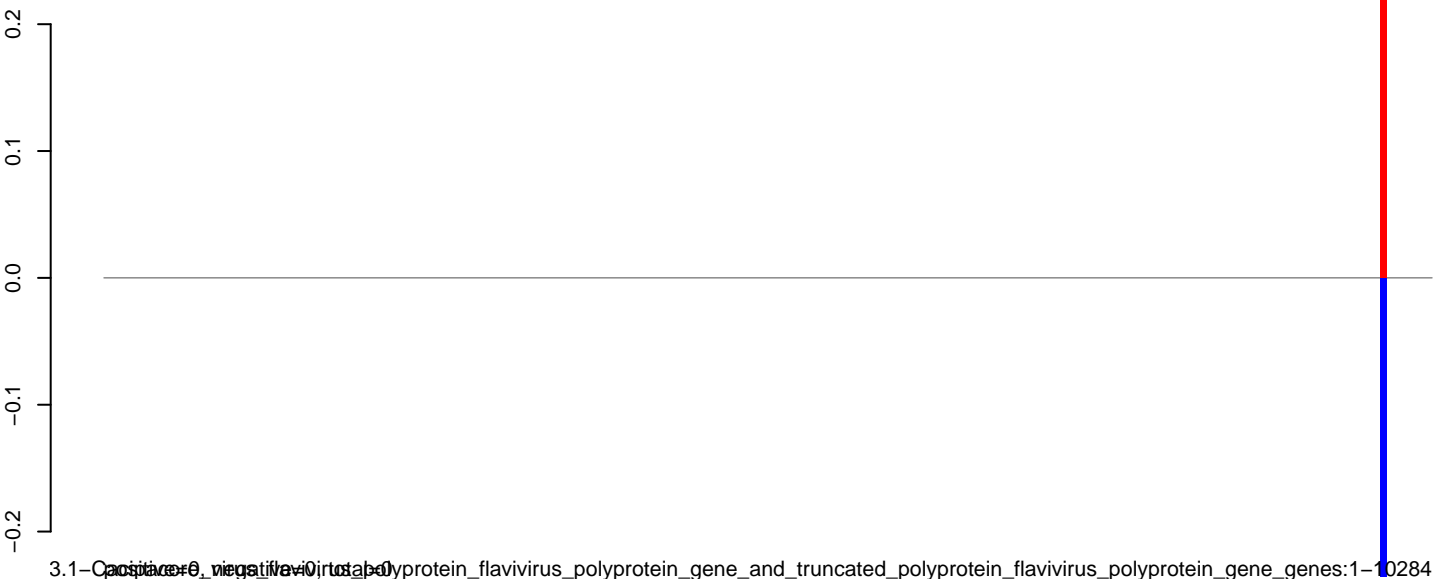
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



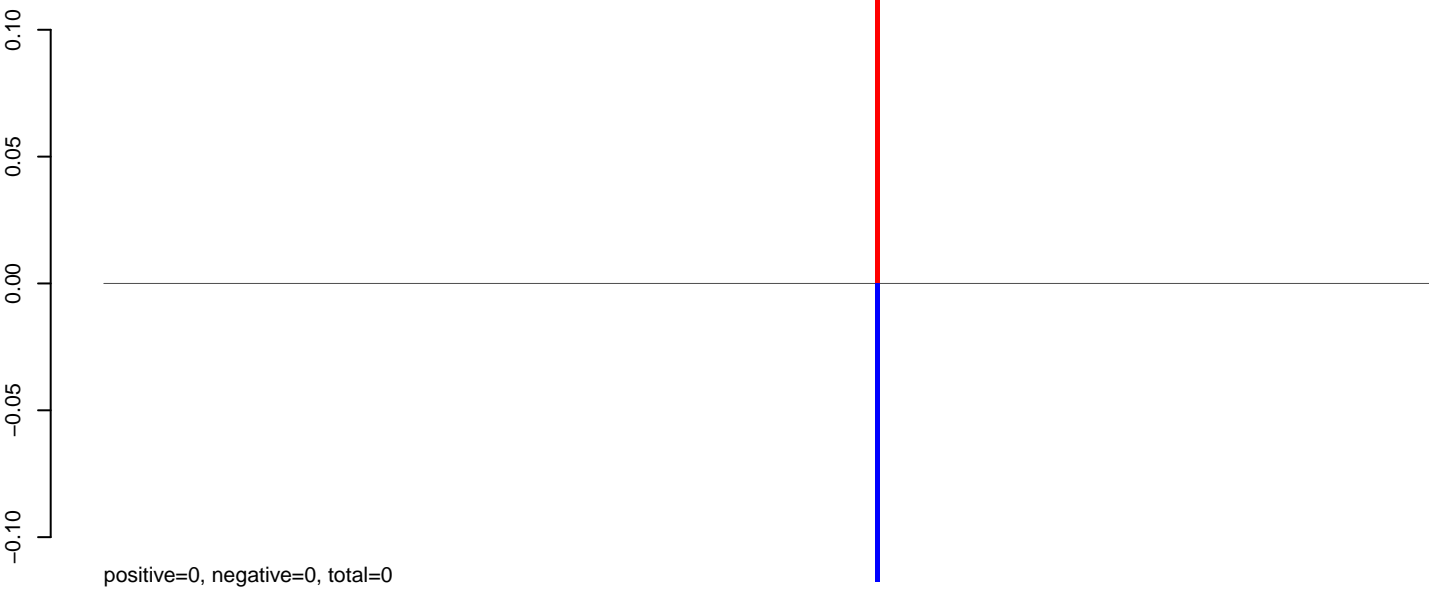
AeAeg_Aag2_DENV1.rep



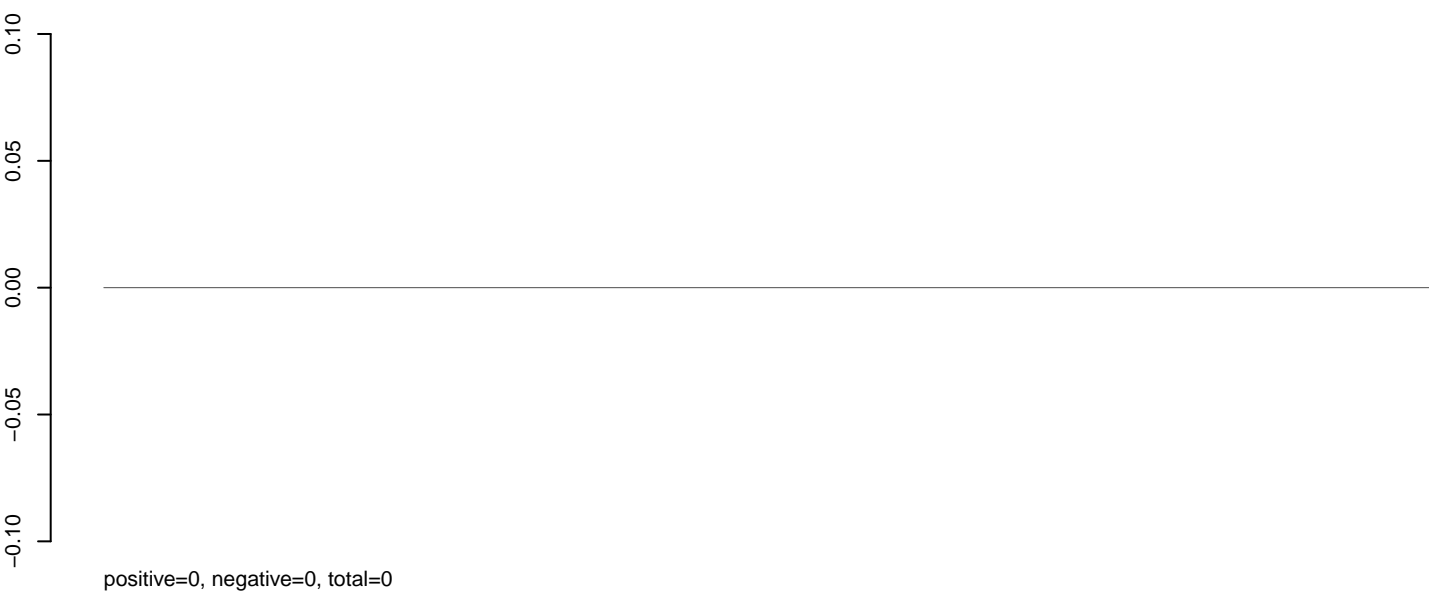
3.1-Opisano_0_negativo_0_total_0 polyprotein_flavivirus_polyprotein_gene_and_truncated_polyprotein_flavivirus_polyprotein_gene_genes:1-10284

0 2000 4000 6000 8000 10000

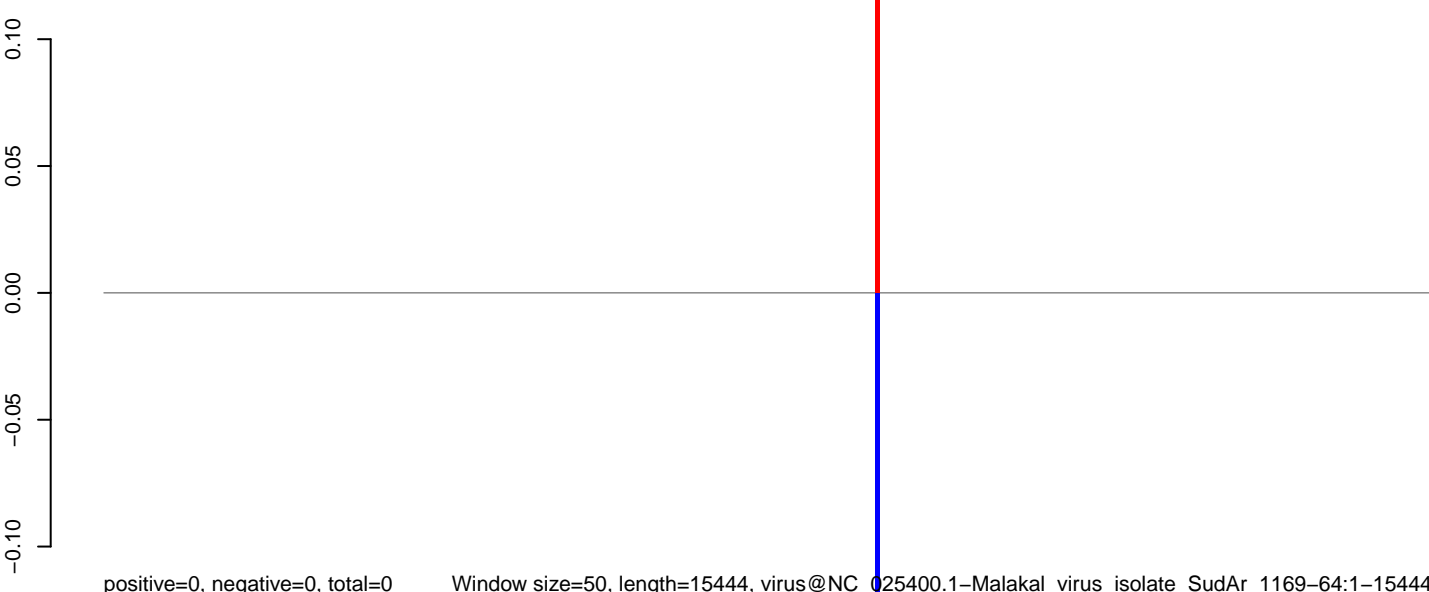
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



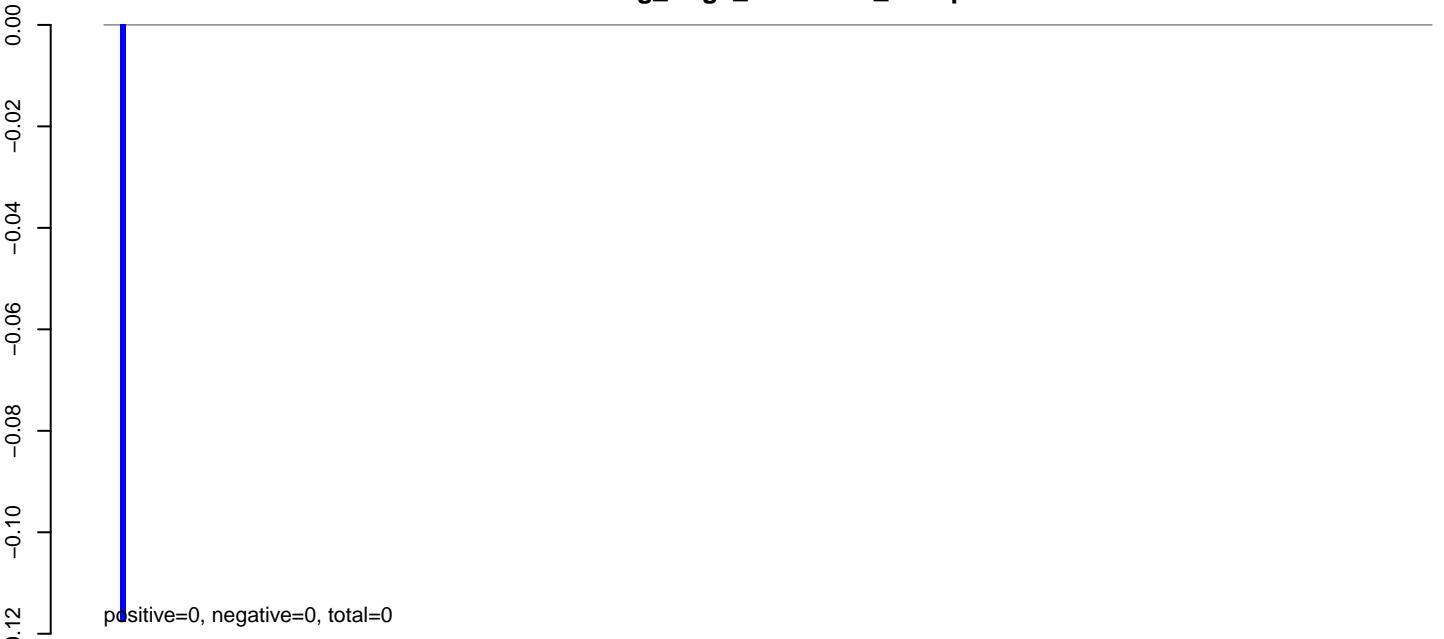
AeAeg_Aag2_DENV1.rep



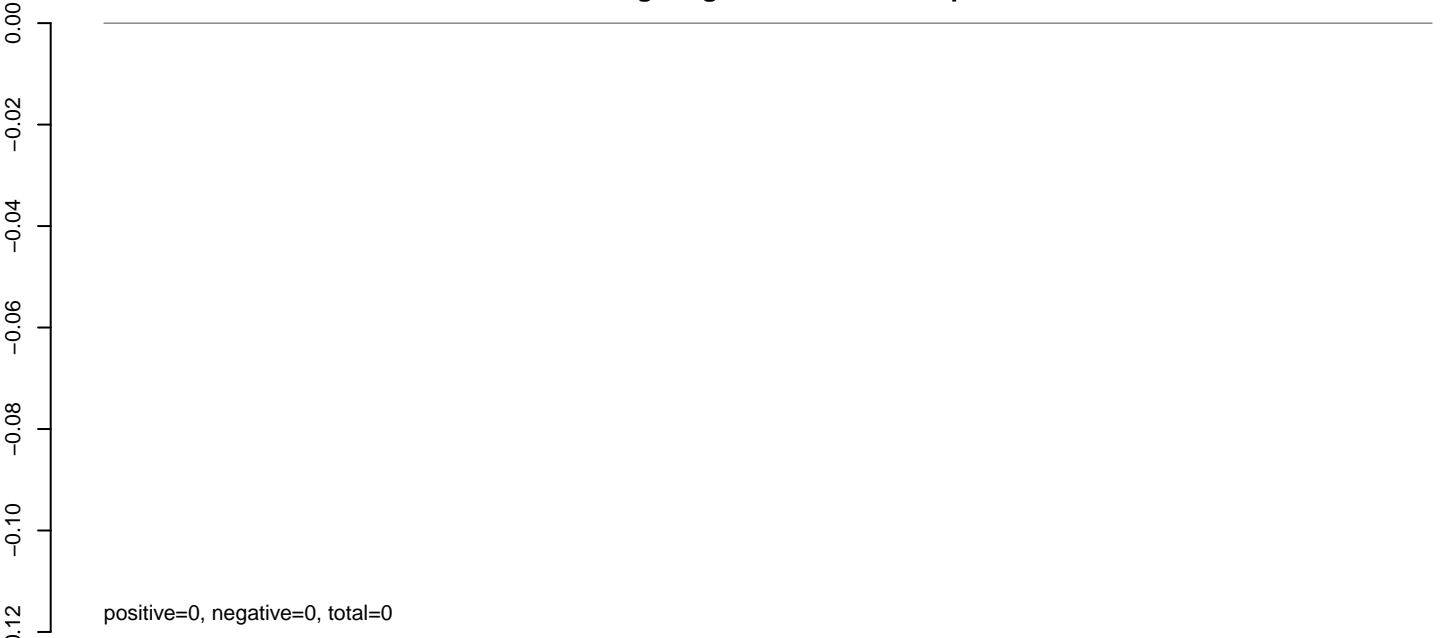
Window size=50, length=15444, virus@NC_025400.1-Malaka1_virus_isolate_SudAr_1169-64:1-15444

0 5000 10000 15000

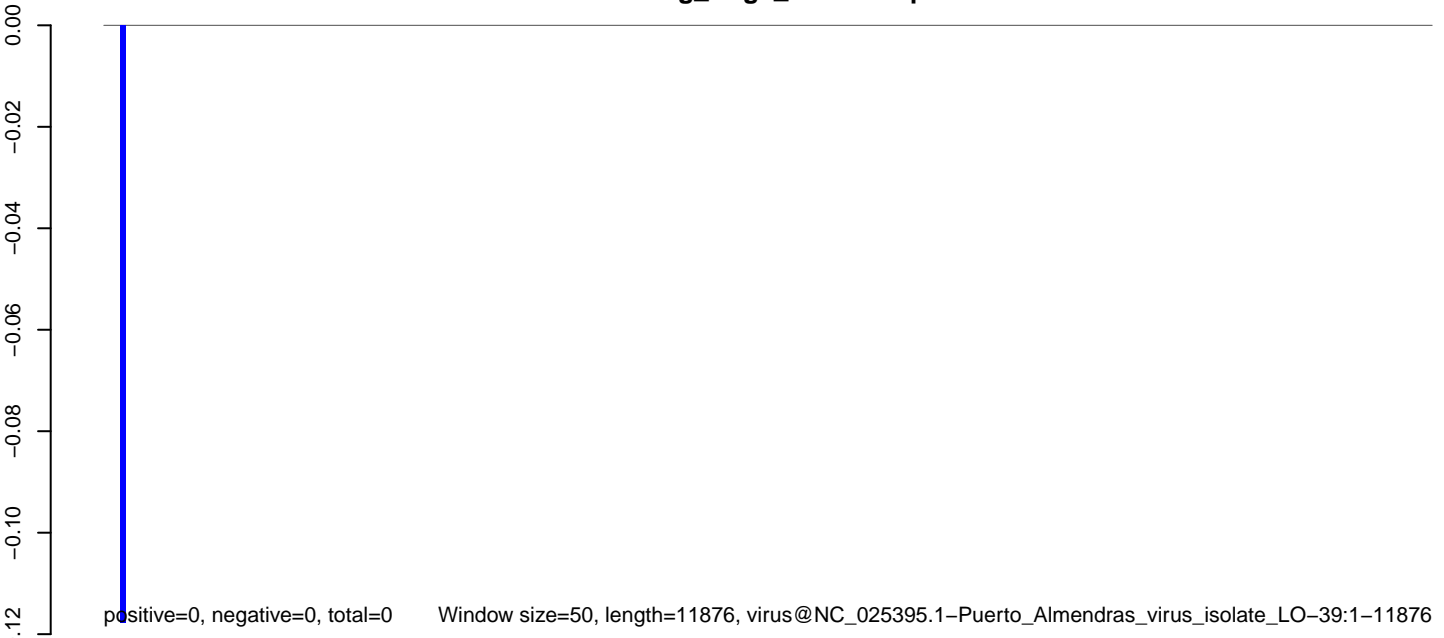
AeAeg_Aag2_DENV1.18_23.rep



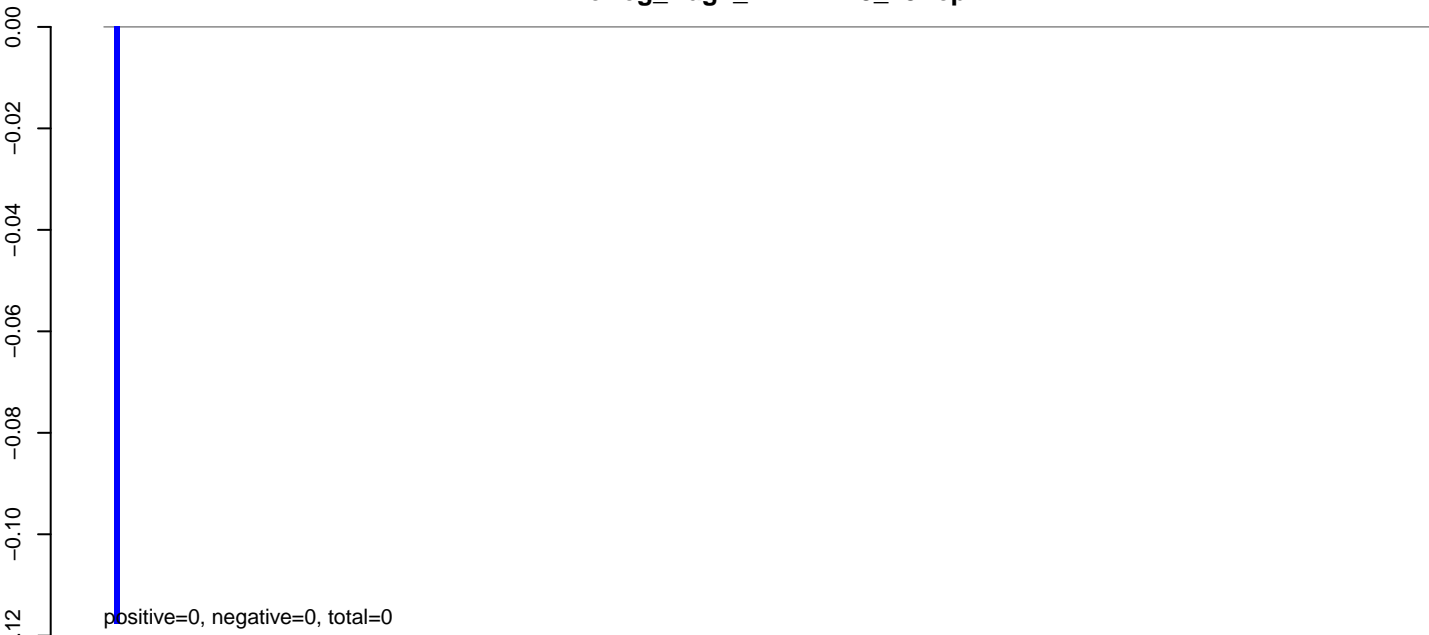
AeAeg_Aag2_DENV1.24_35.rep



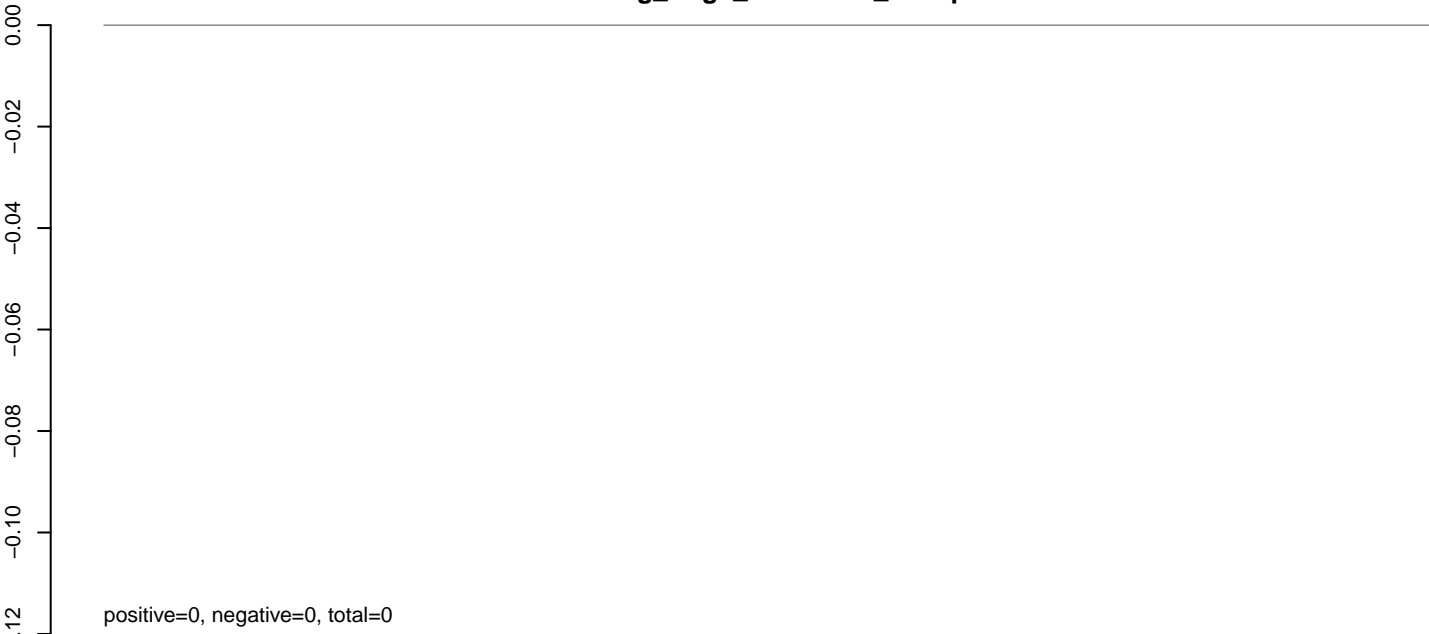
AeAeg_Aag2_DENV1.rep



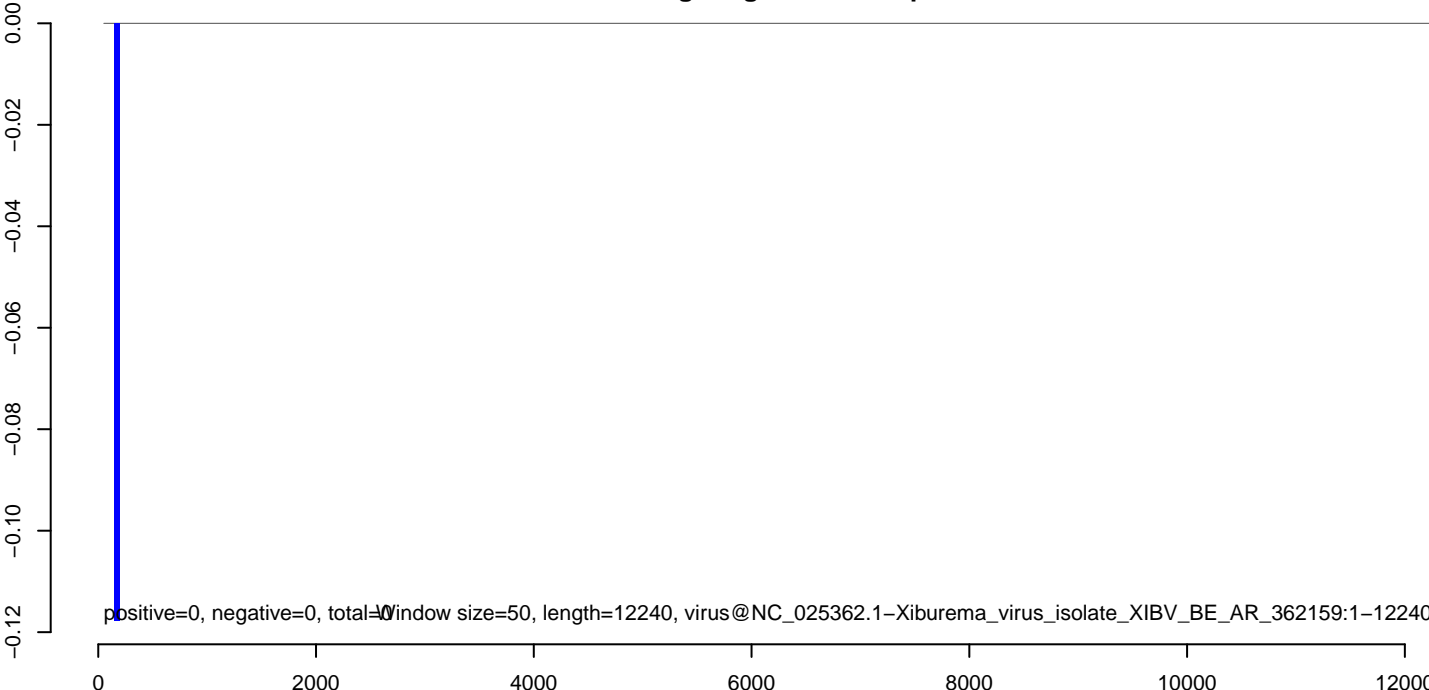
AeAeg_Aag2_DENV1.18_23.rep



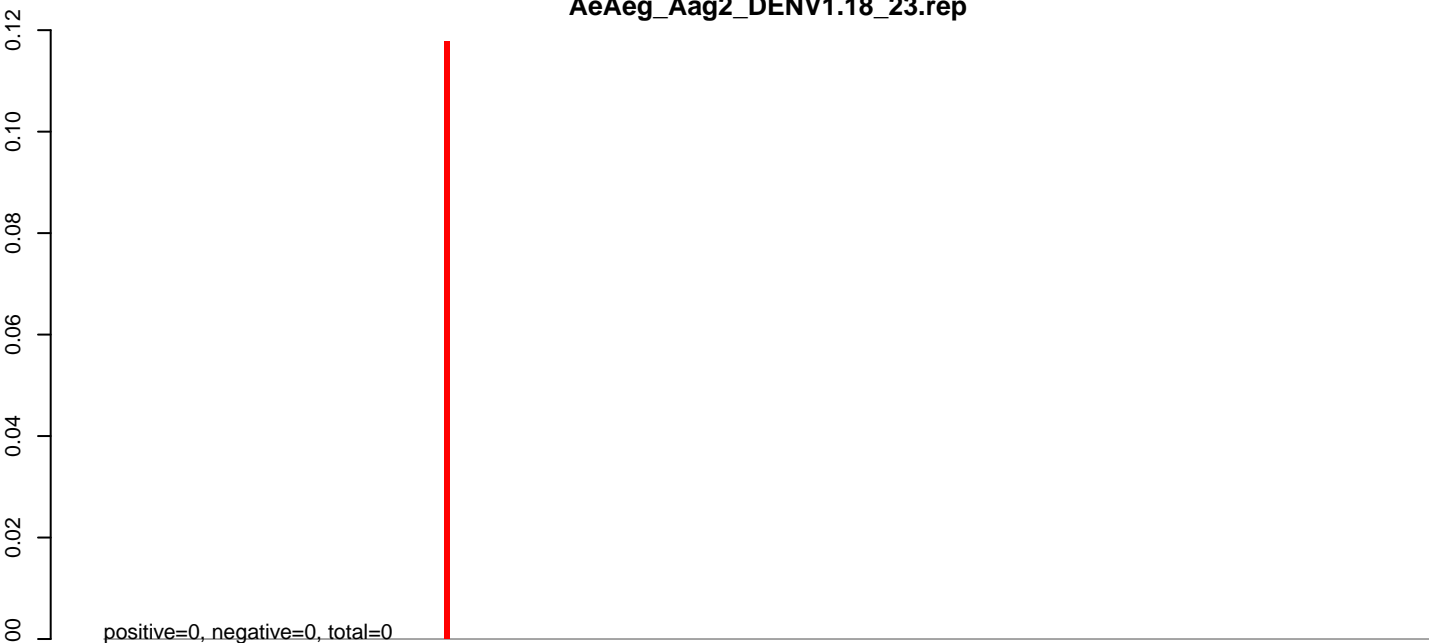
AeAeg_Aag2_DENV1.24_35.rep



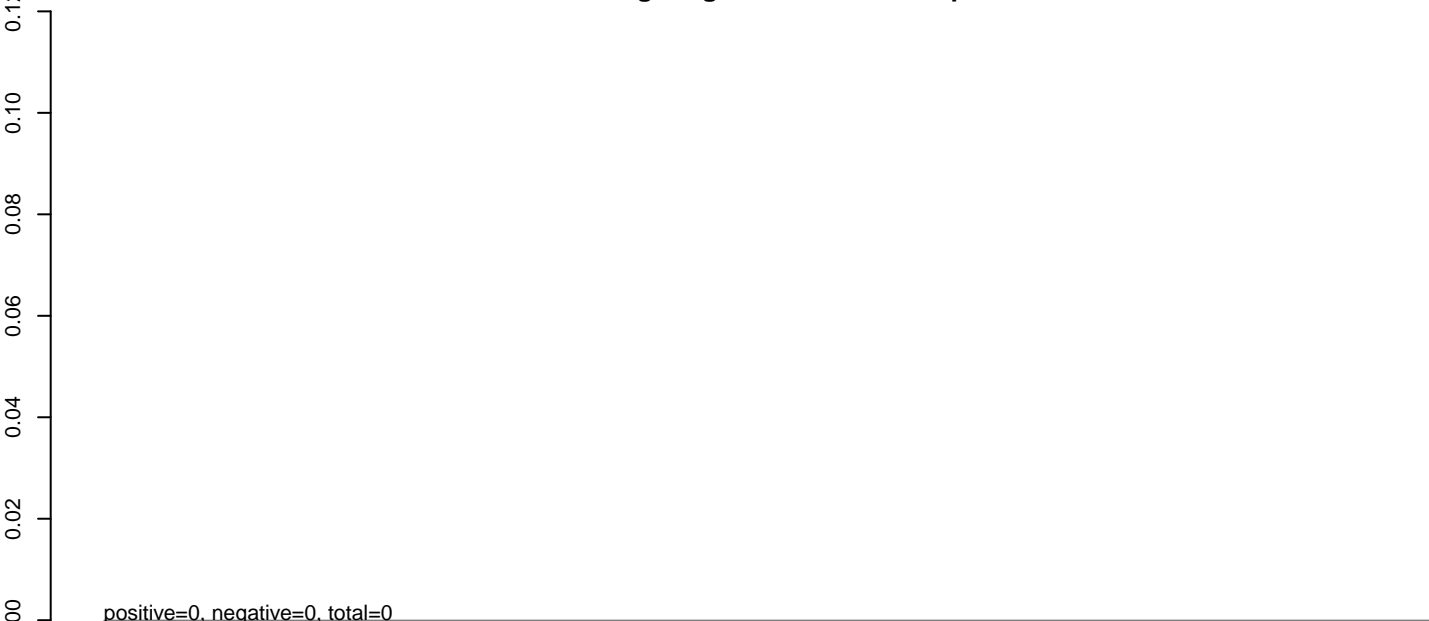
AeAeg_Aag2_DENV1.rep



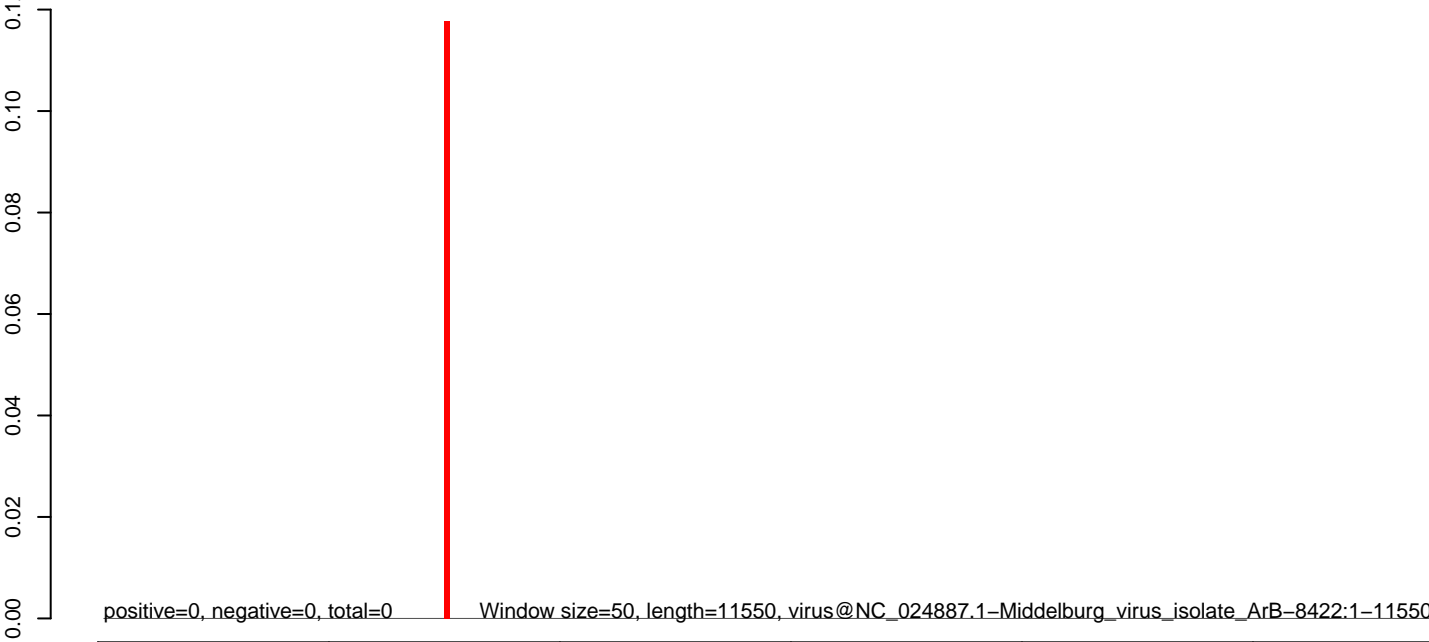
AeAeg_Aag2_DENV1.18_23.rep



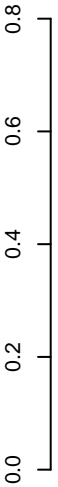
AeAeg_Aag2_DENV1.24_35.rep



AeAeg_Aag2_DENV1.rep

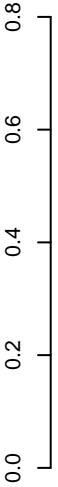


AeAeg_Aag2_DENV1.18_23.rep



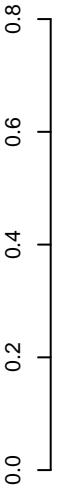
positive=2, negative=0, total=2

AeAeg_Aag2_DENV1.24_35.rep

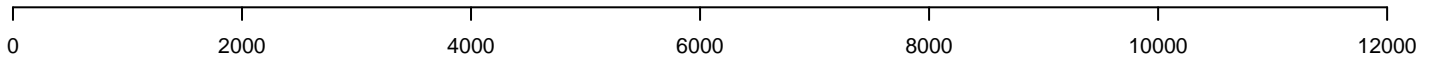


positive=0, negative=0, total=0

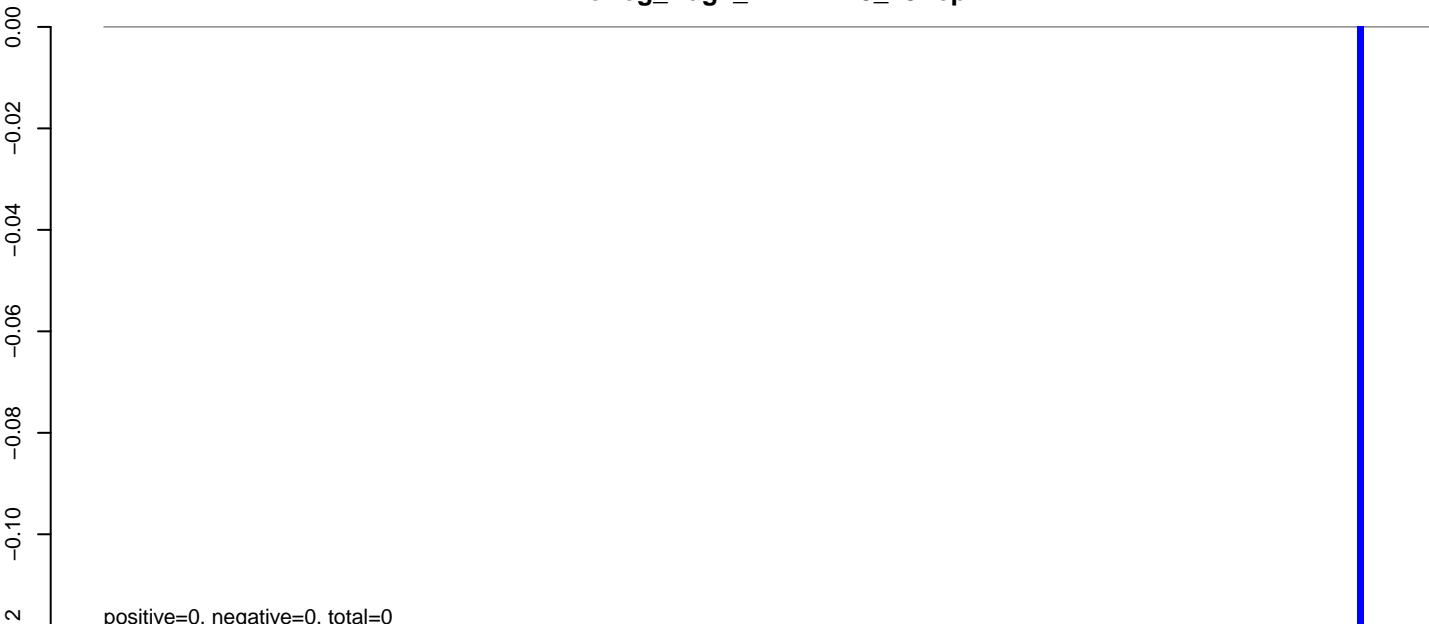
AeAeg_Aag2_DENV1.rep



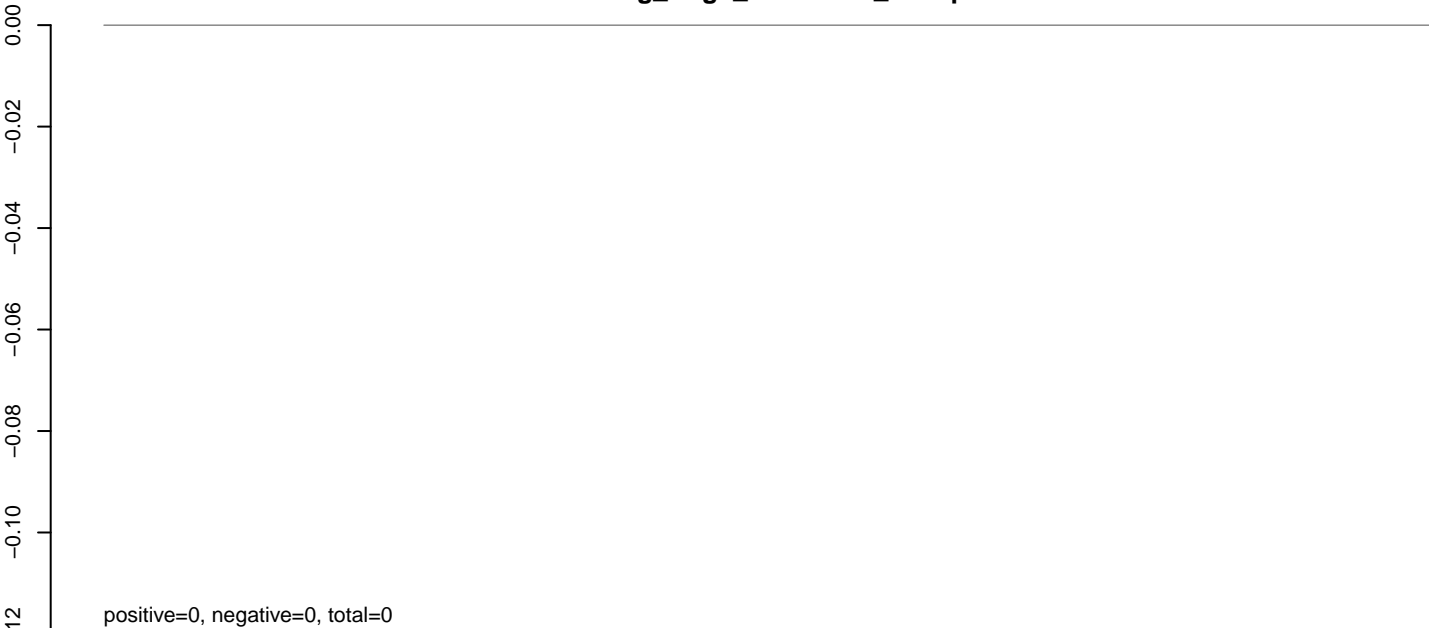
positive=2, negative=50, total=52, virus@NC_023812.1-Madariaga_virus_strain_MADV_Cebus_apella_BRA_BEAN5122_1956:1-11624



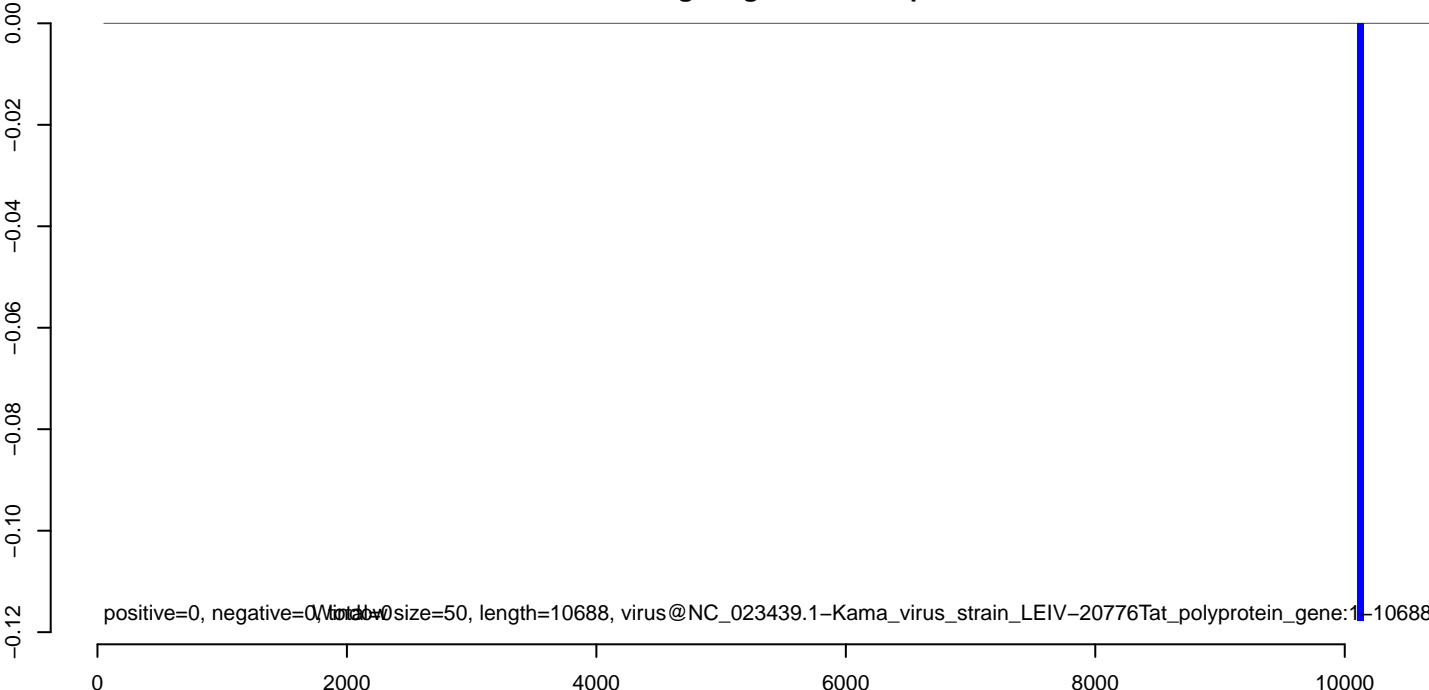
AeAeg_Aag2_DENV1.18_23.rep



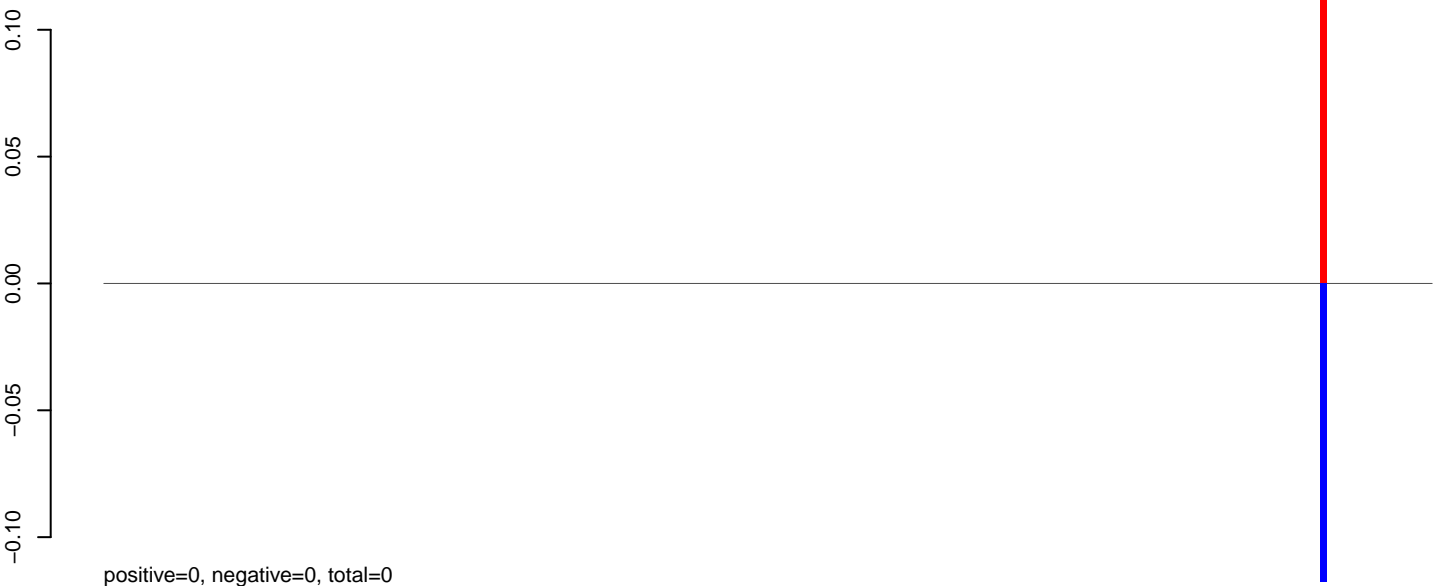
AeAeg_Aag2_DENV1.24_35.rep



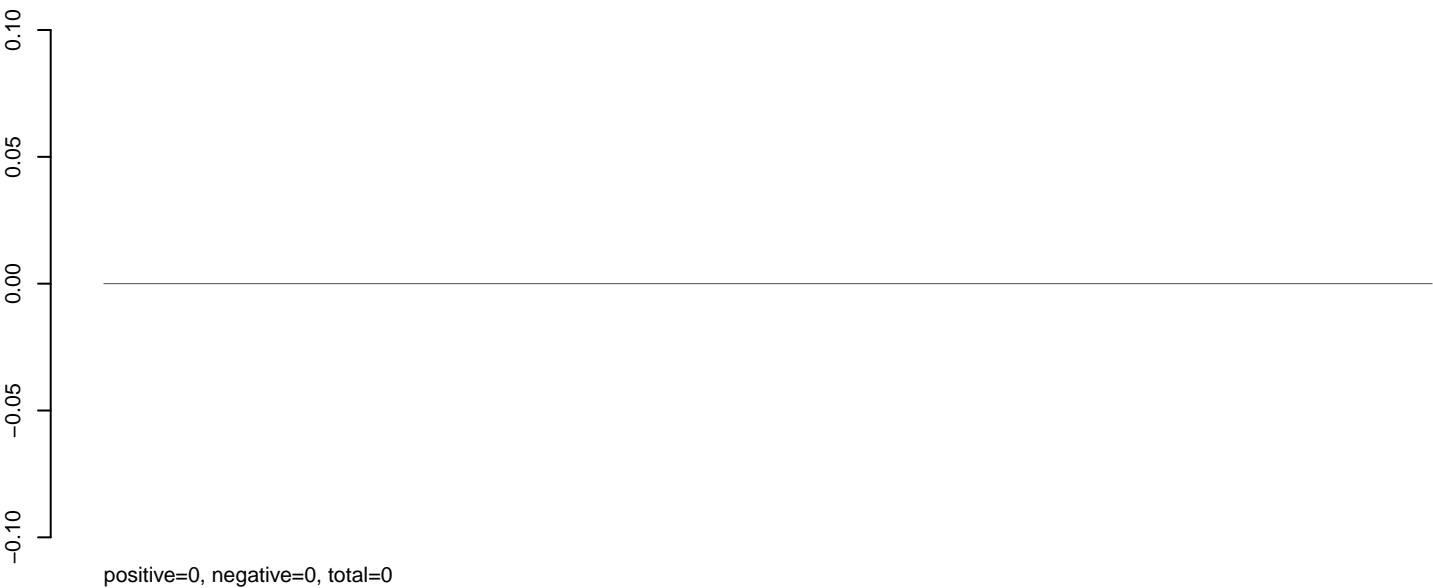
AeAeg_Aag2_DENV1.rep



AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



AeAeg_Aag2_DENV1.rep



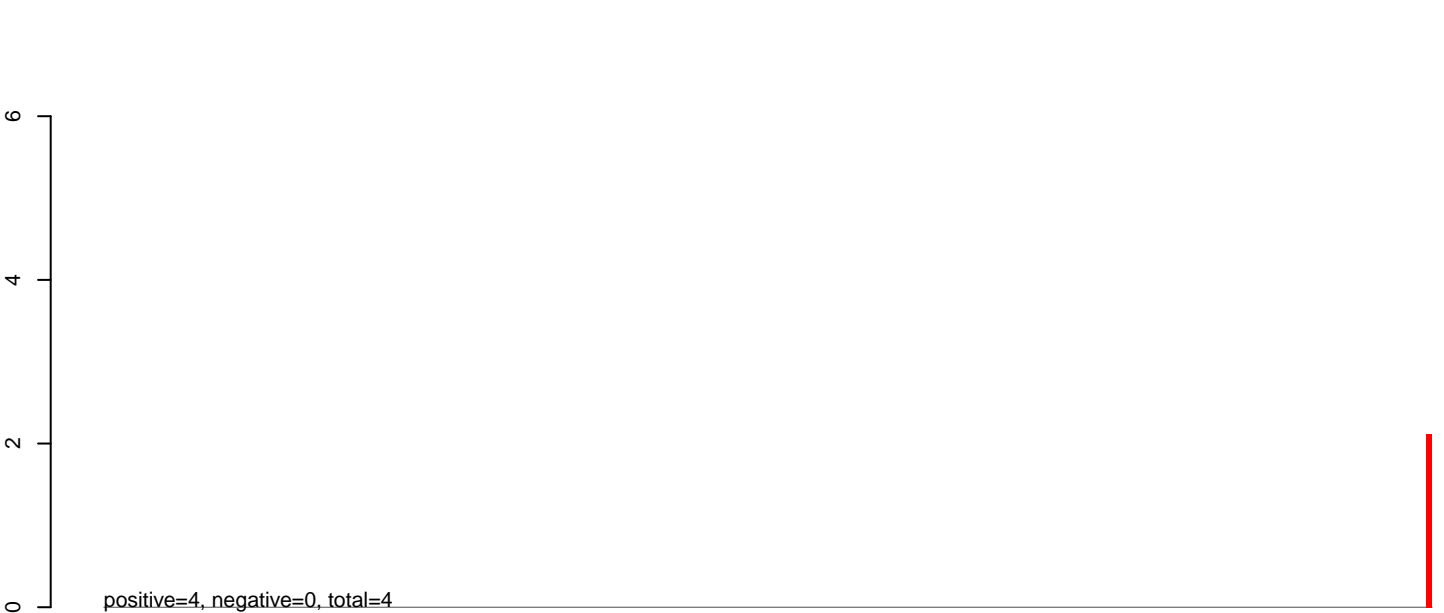
Window size=50, length=10733, virus@NC_017086.1-Chaoyang_virus:1-10733

0 2000 4000 6000 8000 10000

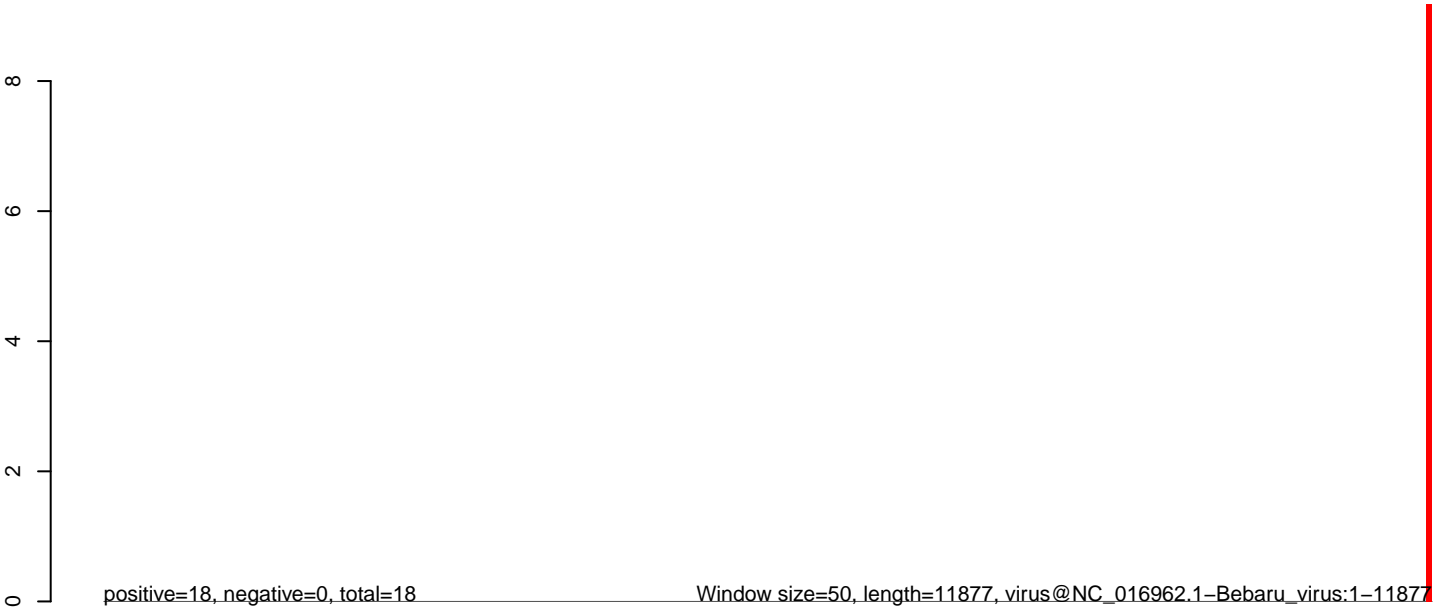
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

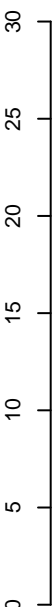


AeAeg_Aag2_DENV1.rep



0 2000 4000 6000 8000 10000 12000

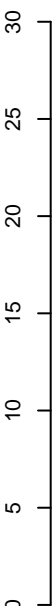
AeAeg_Aag2_DENV1.18_23.rep



positive=61, negative=0, total=61



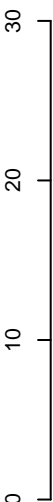
AeAeg_Aag2_DENV1.24_35.rep



positive=15, negative=0, total=15

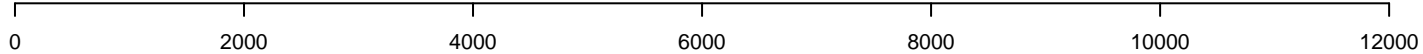


AeAeg_Aag2_DENV1.rep

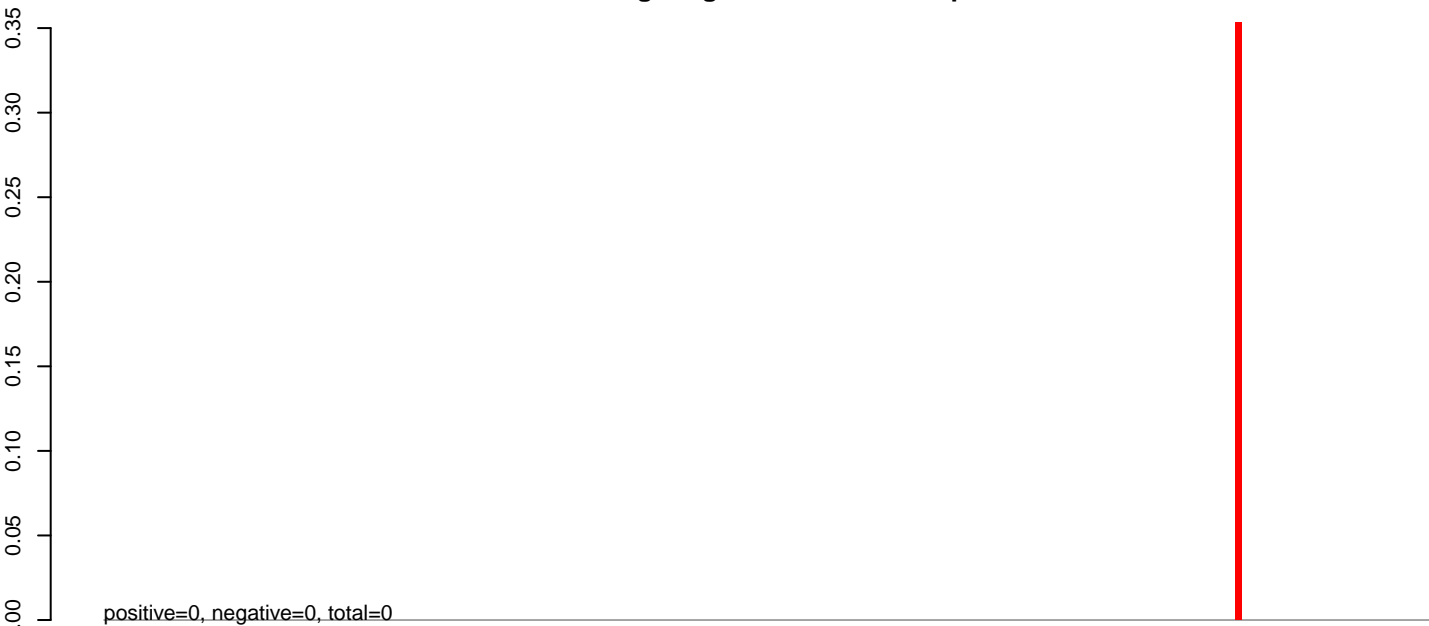


positive=75, negative=0, total=75

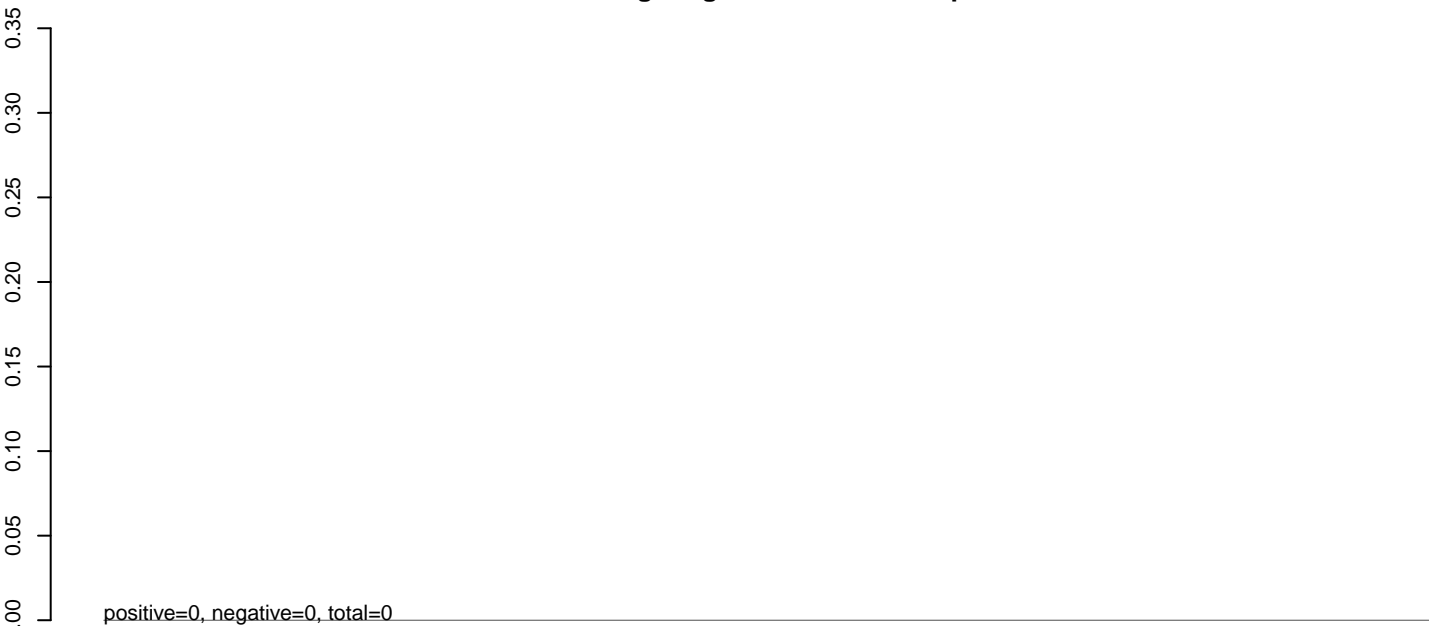
Window size=50, length=11616, virus@NC_016961.1-Whataroa_virus:1-11616



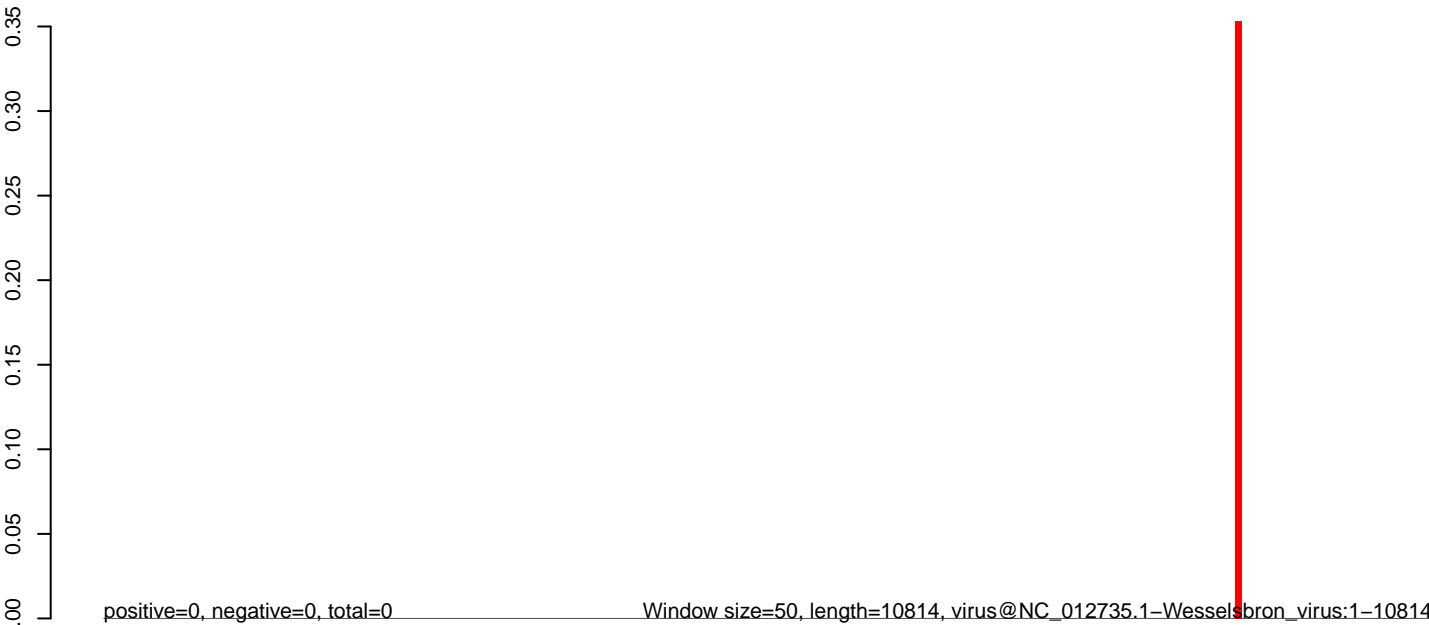
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

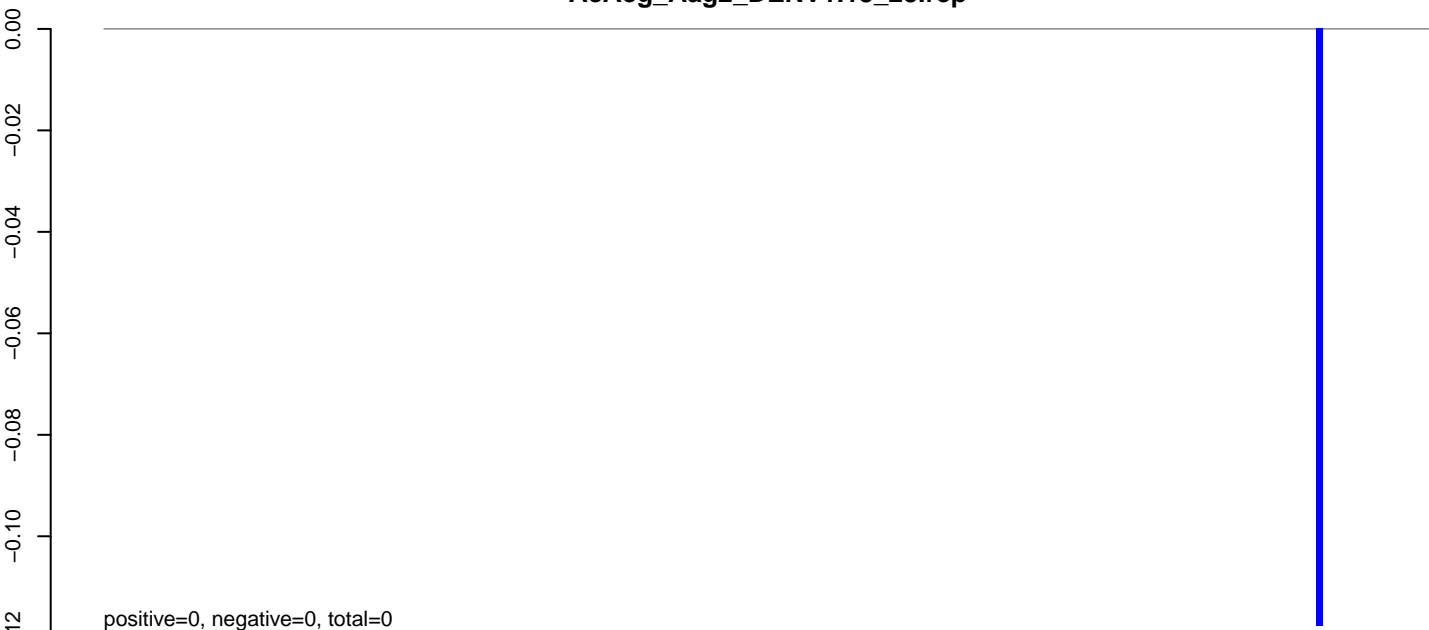


AeAeg_Aag2_DENV1.rep

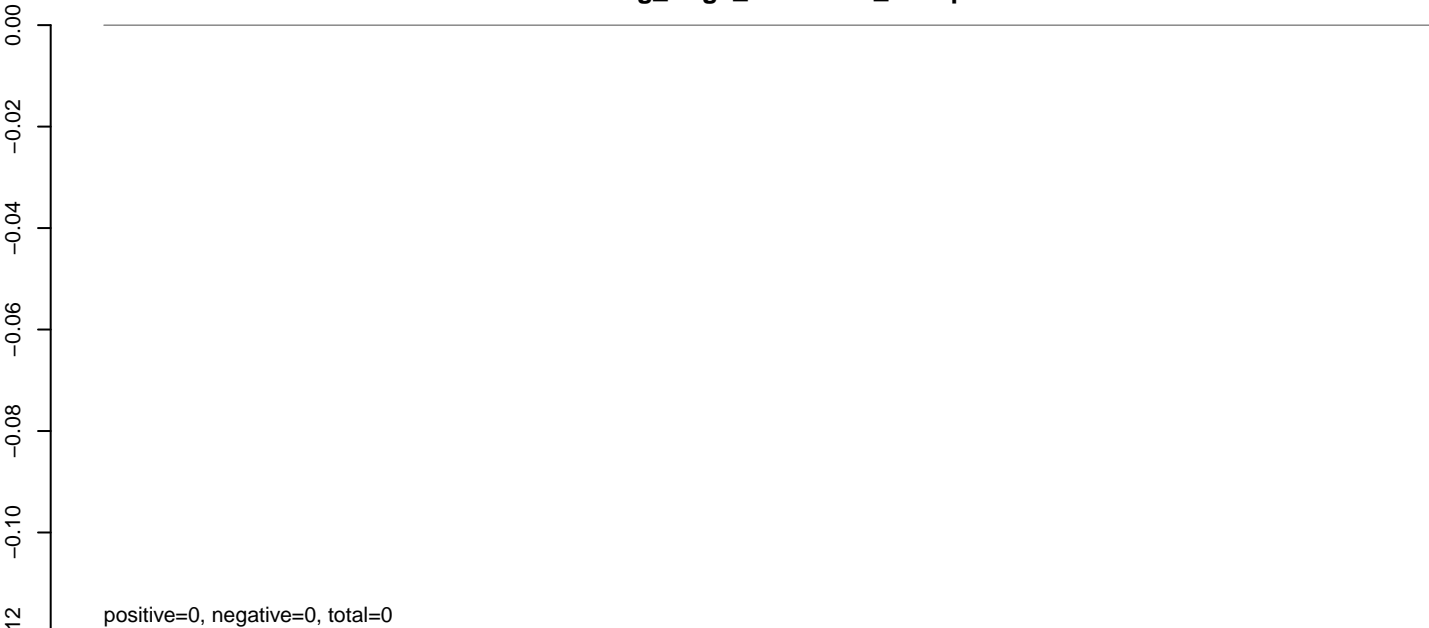


0 2000 4000 6000 8000 10000

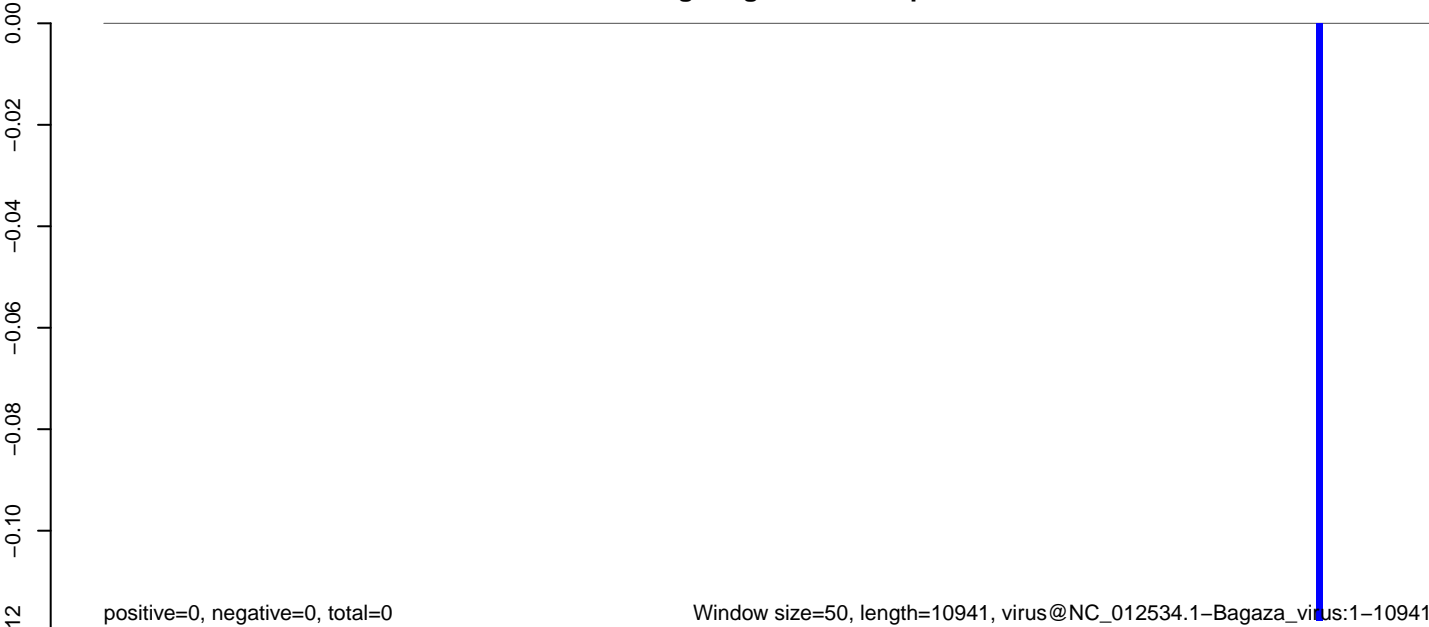
AeAeg_Aag2_DENV1.18_23.rep



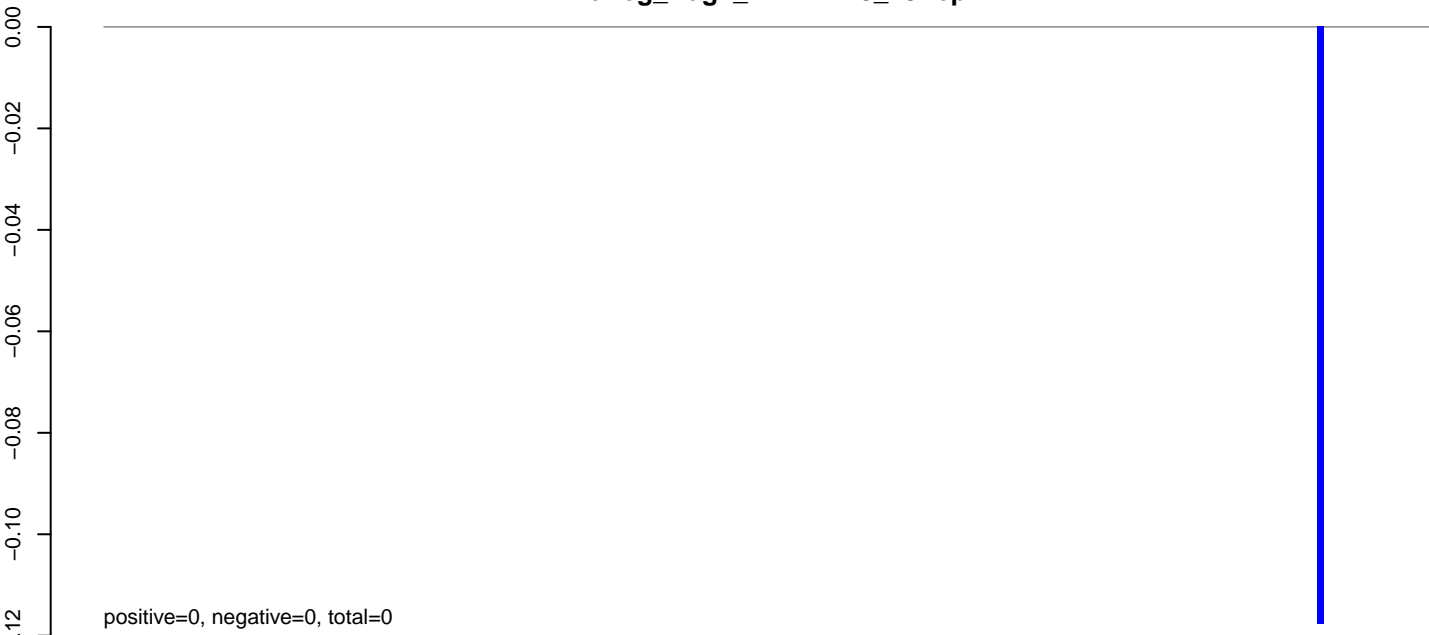
AeAeg_Aag2_DENV1.24_35.rep



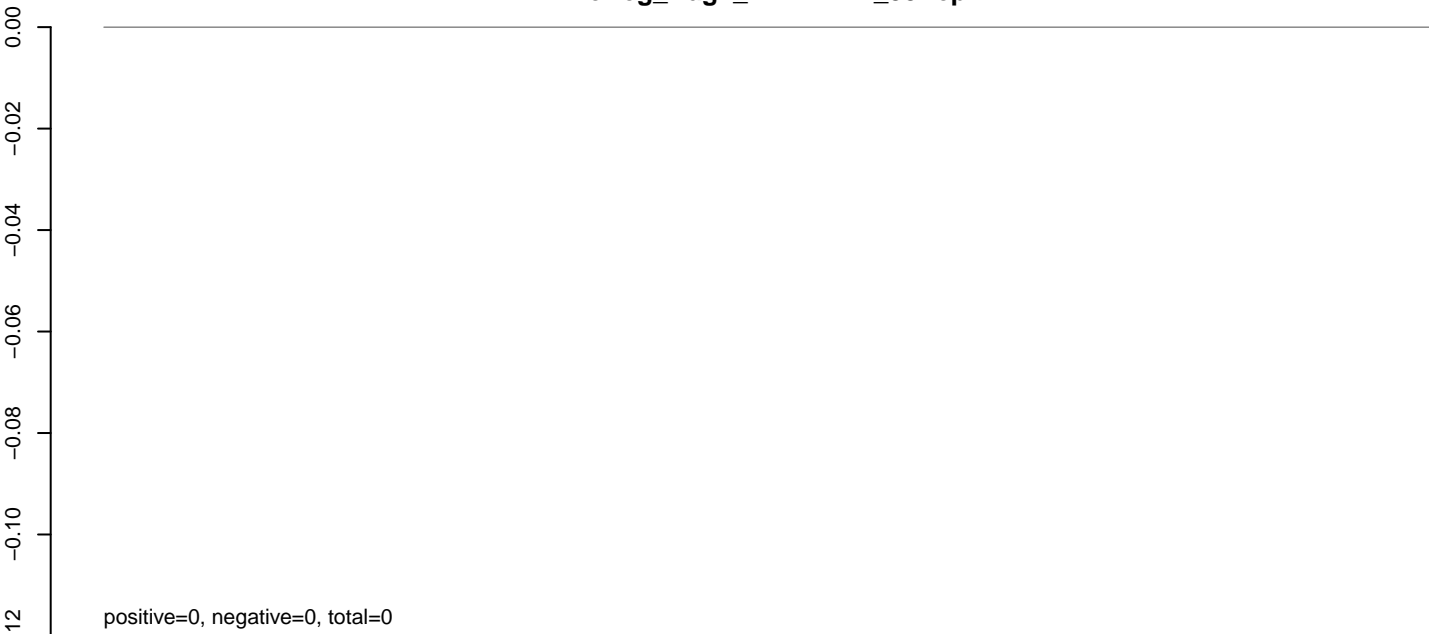
AeAeg_Aag2_DENV1.rep



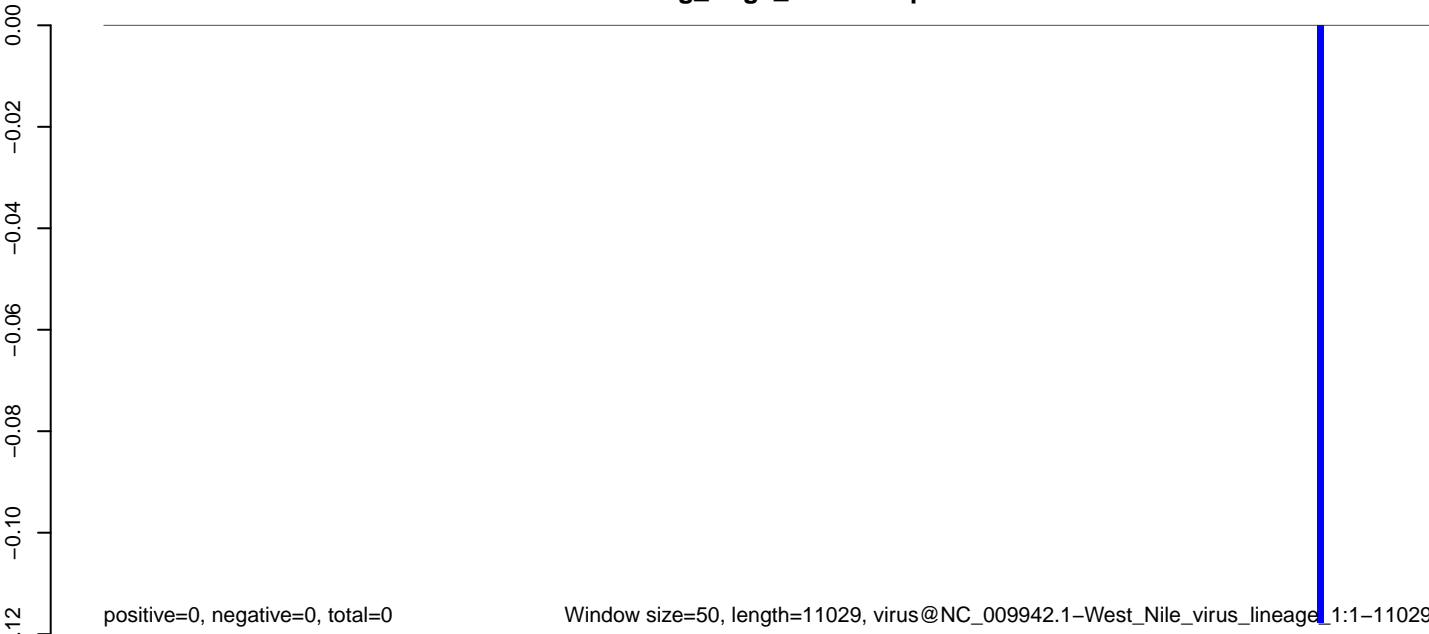
AeAeg_Aag2_DENV1.18_23.rep



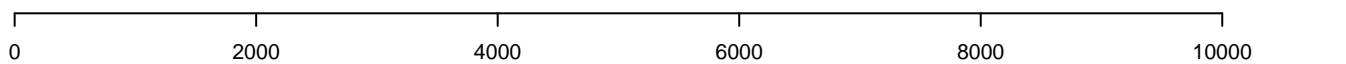
AeAeg_Aag2_DENV1.24_35.rep



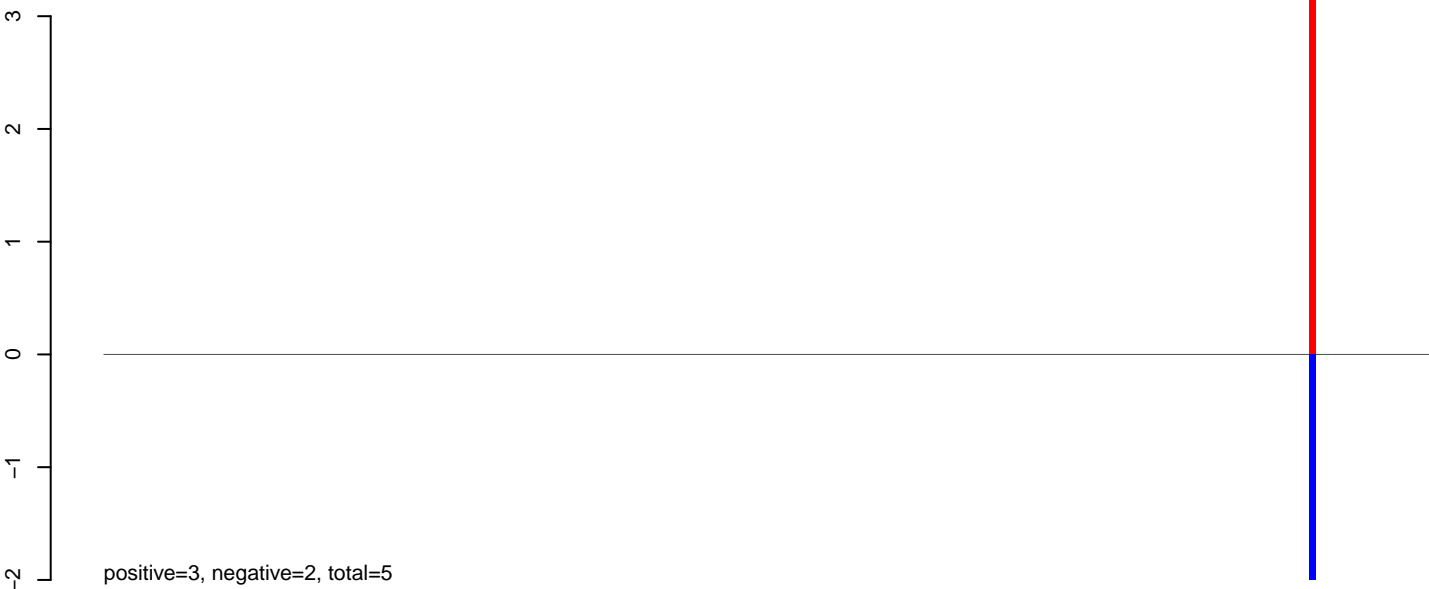
AeAeg_Aag2_DENV1.rep



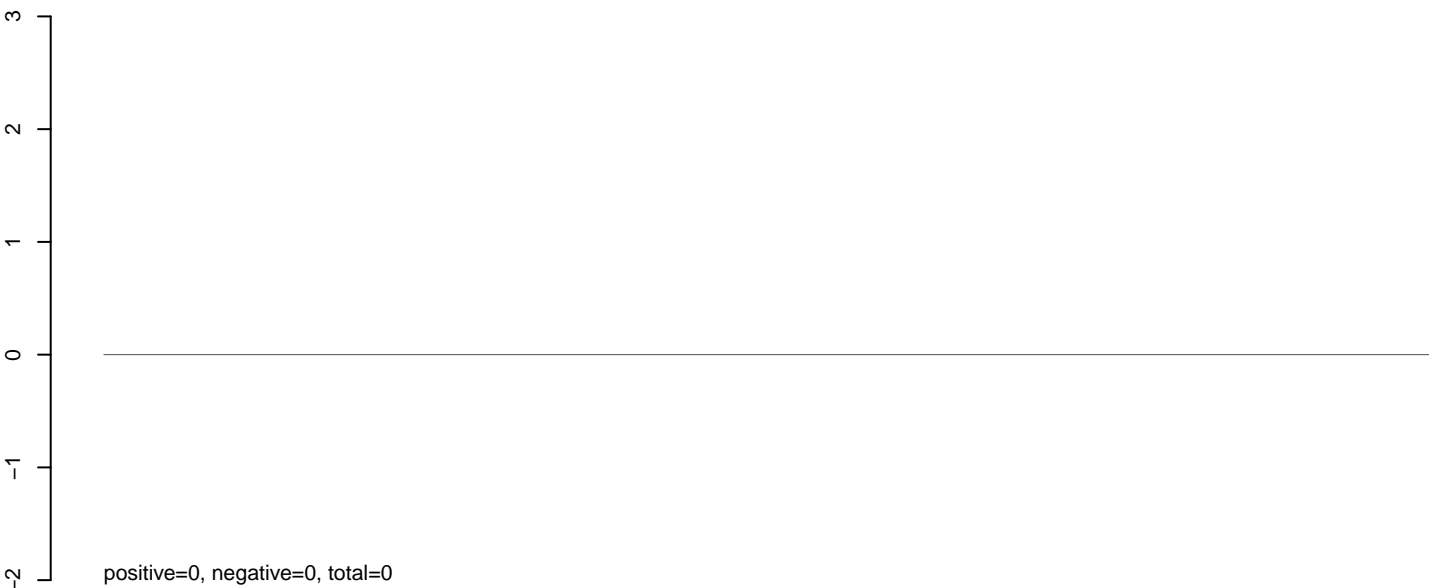
Window size=50, length=11029, virus@NC_009942.1-West_Nile_virus_lineage_1:1-11029



AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

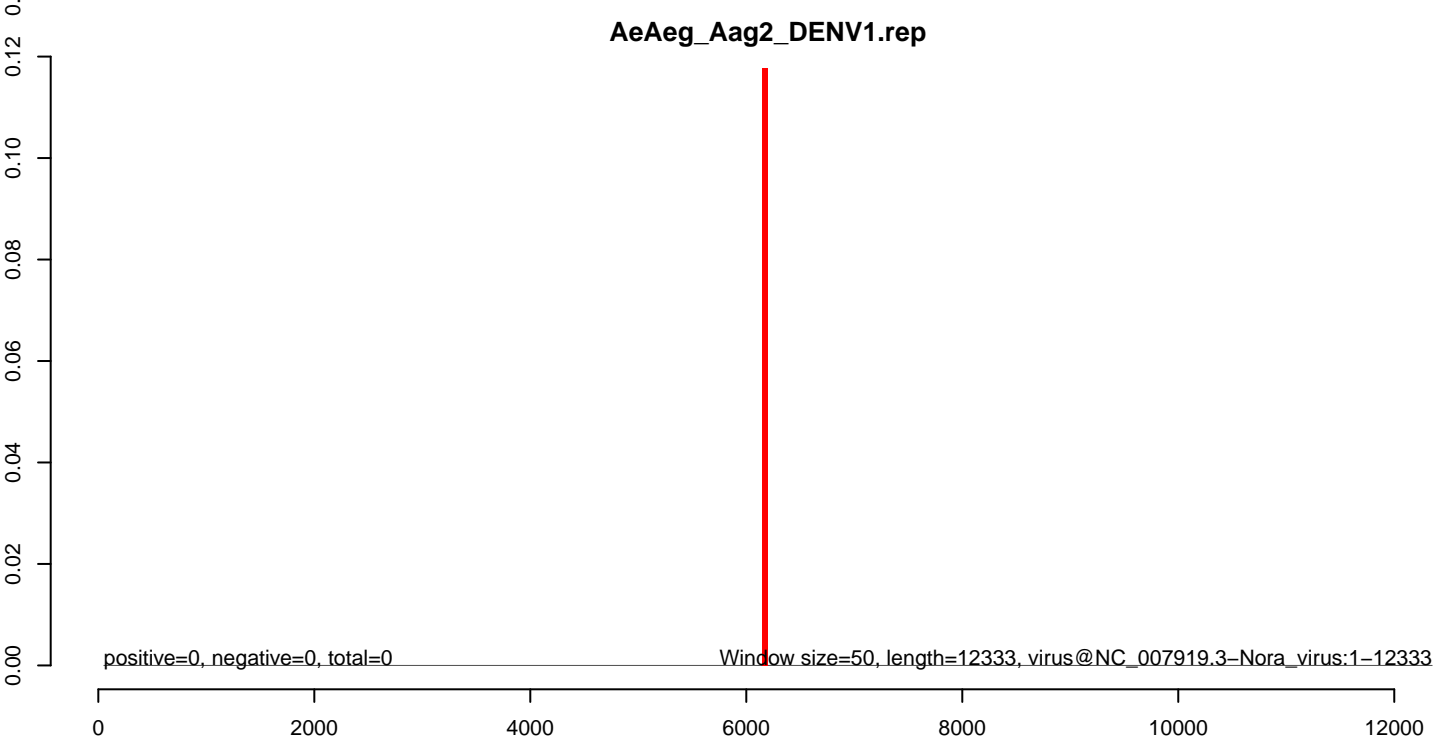
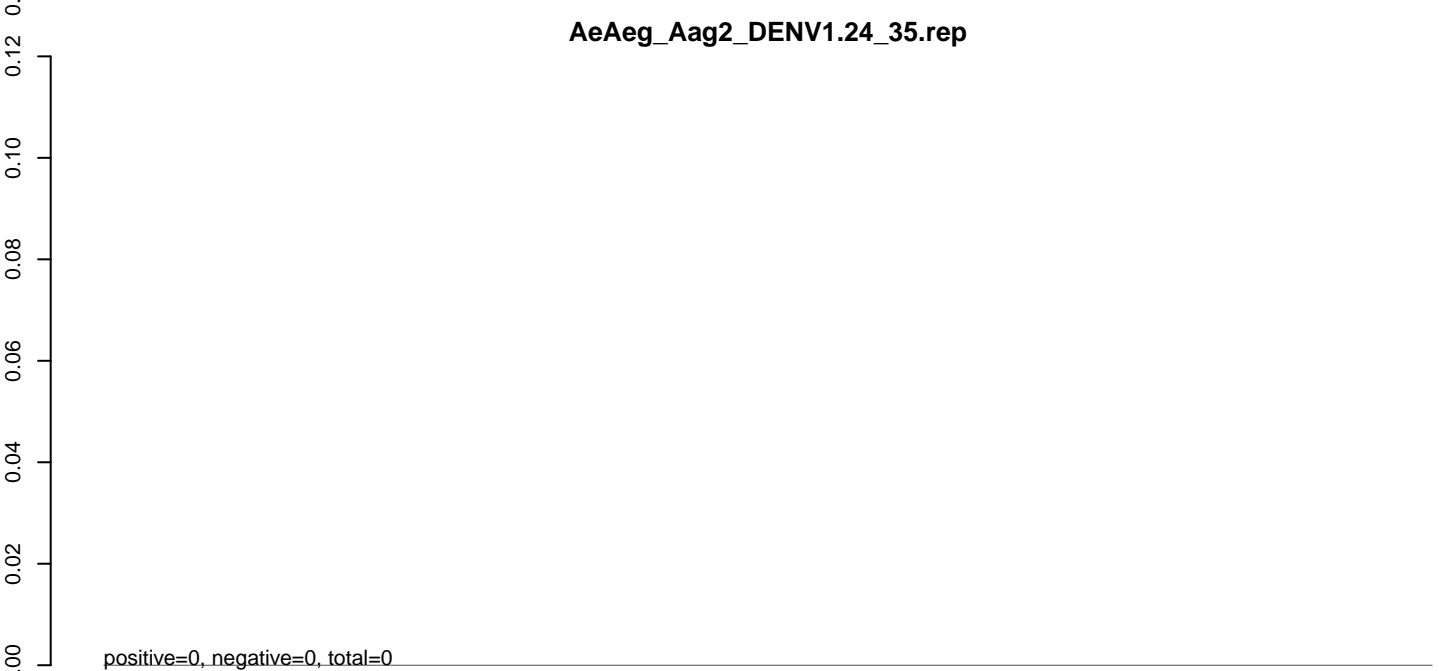
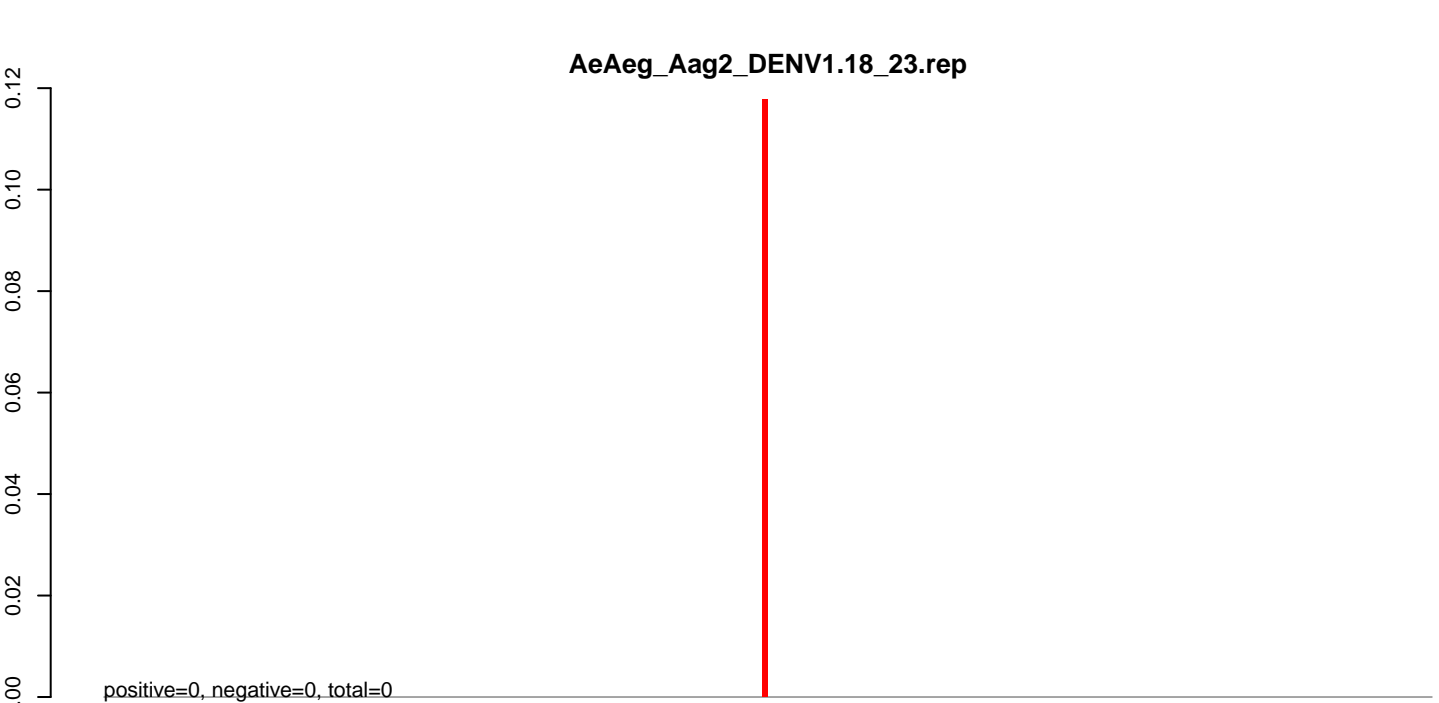


AeAeg_Aag2_DENV1.rep

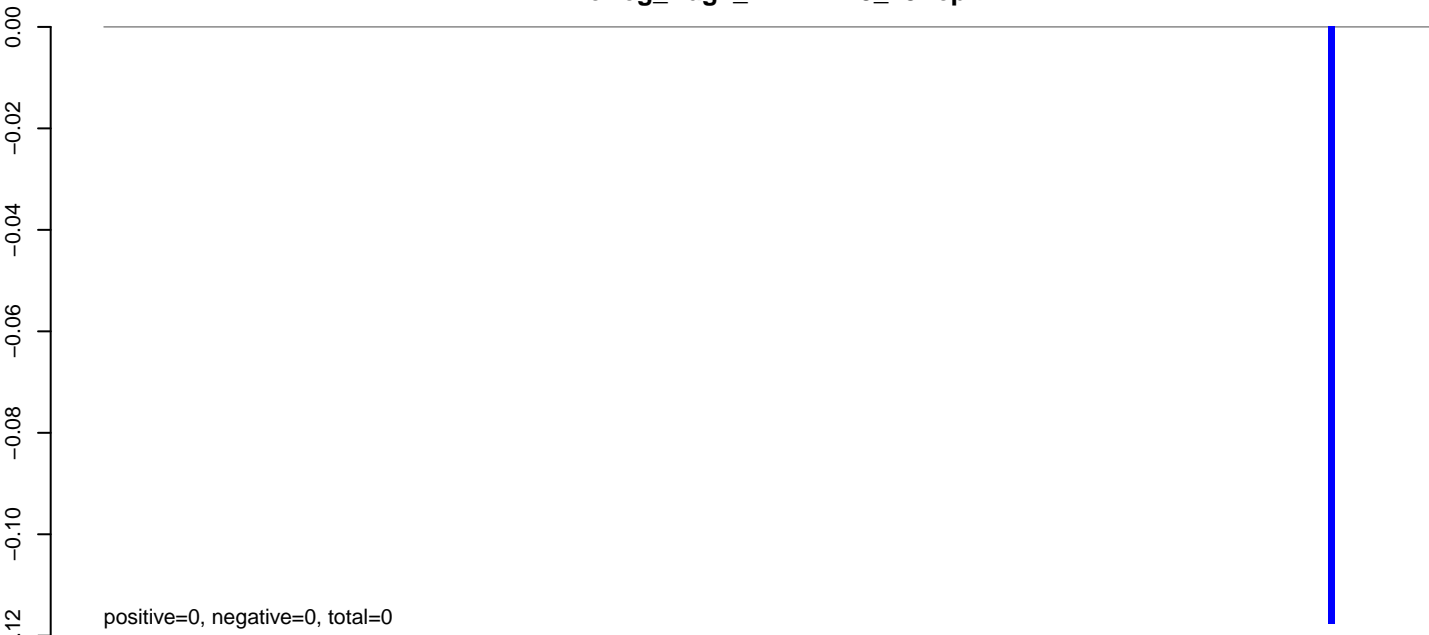


Window size=50, length=10874, virus@NC_009029.2-Kokobera_virus:1-10874

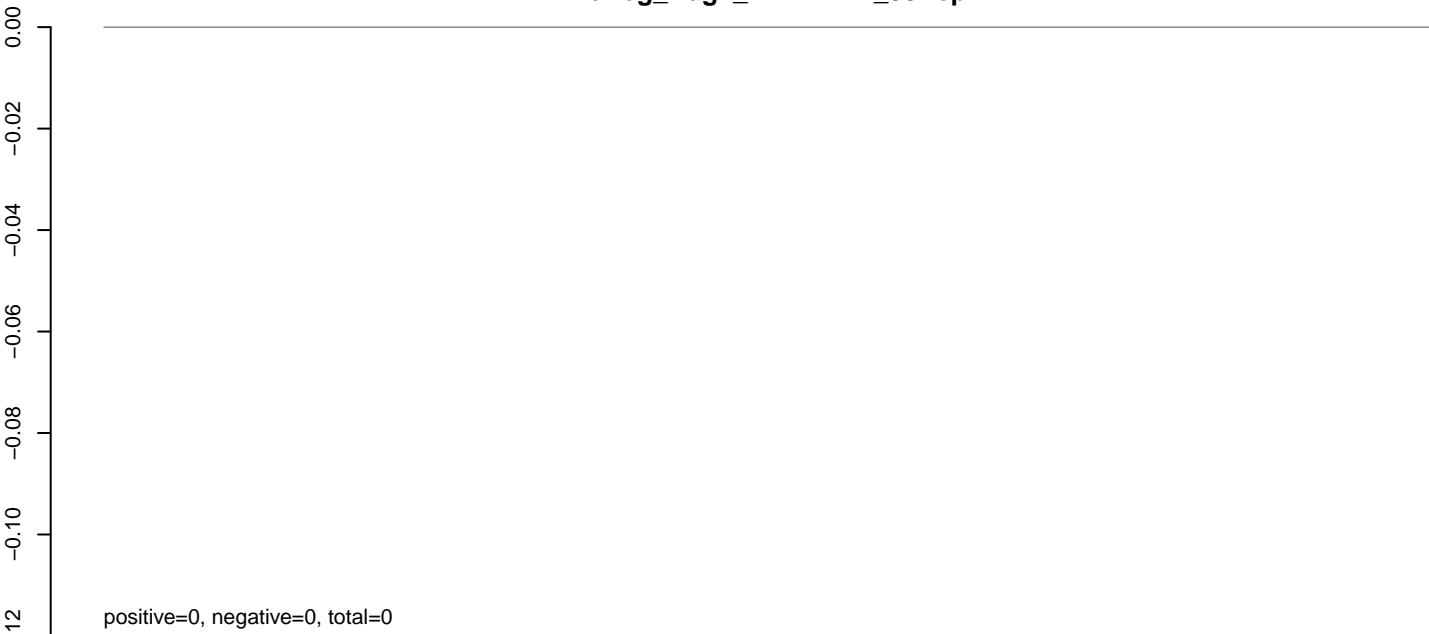
0 2000 4000 6000 8000 10000



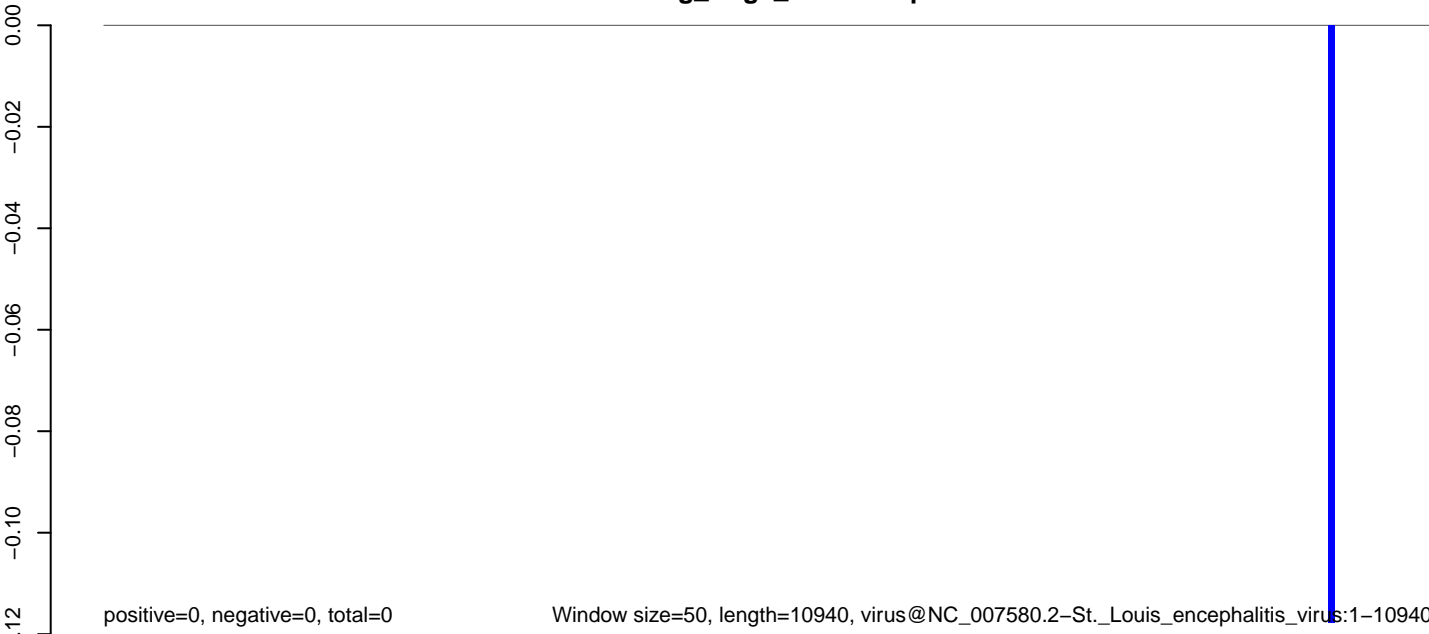
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



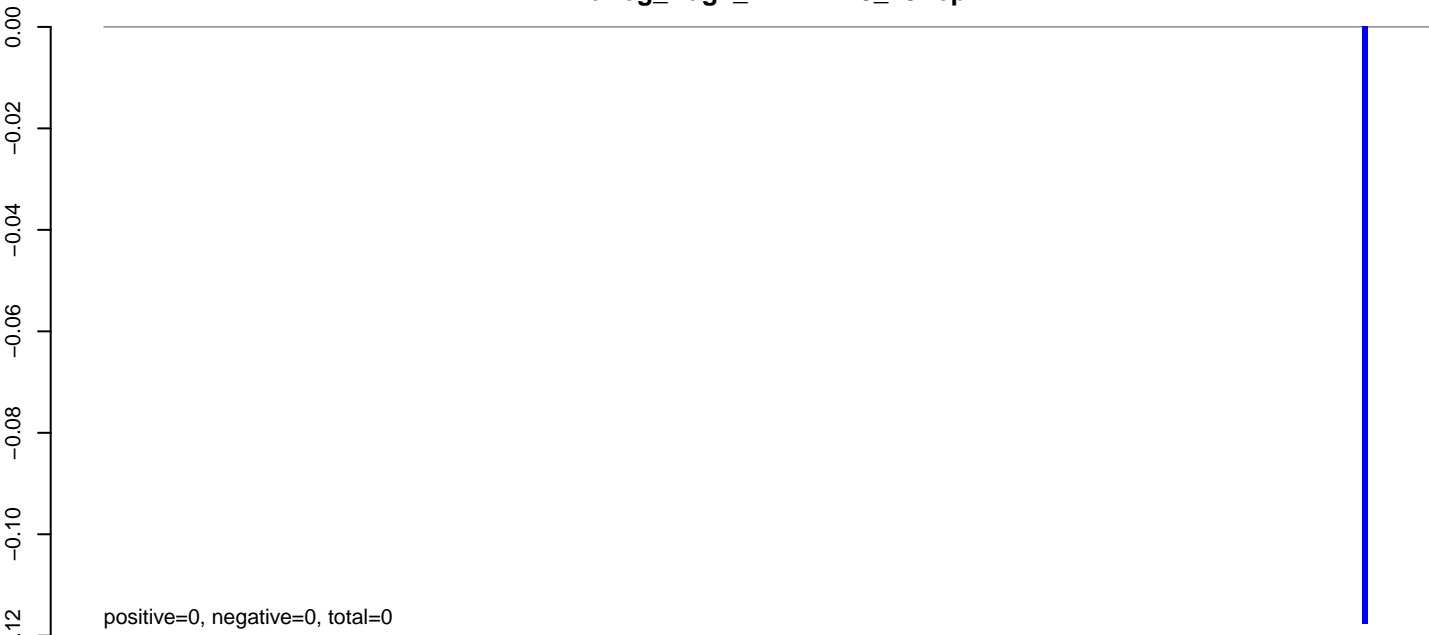
AeAeg_Aag2_DENV1.rep



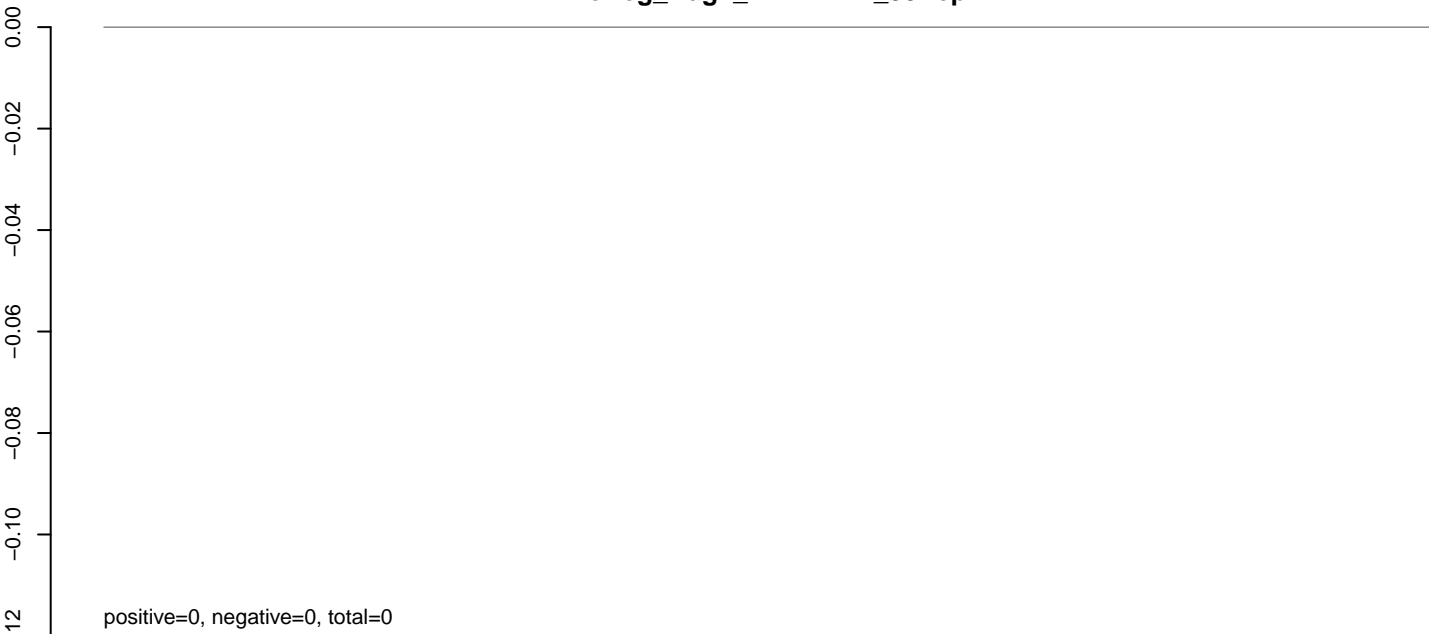
Window size=50, length=10940, virus@NC_007580.2-St._Louis_encephalitis_virus:1-10940

0 2000 4000 6000 8000 10000

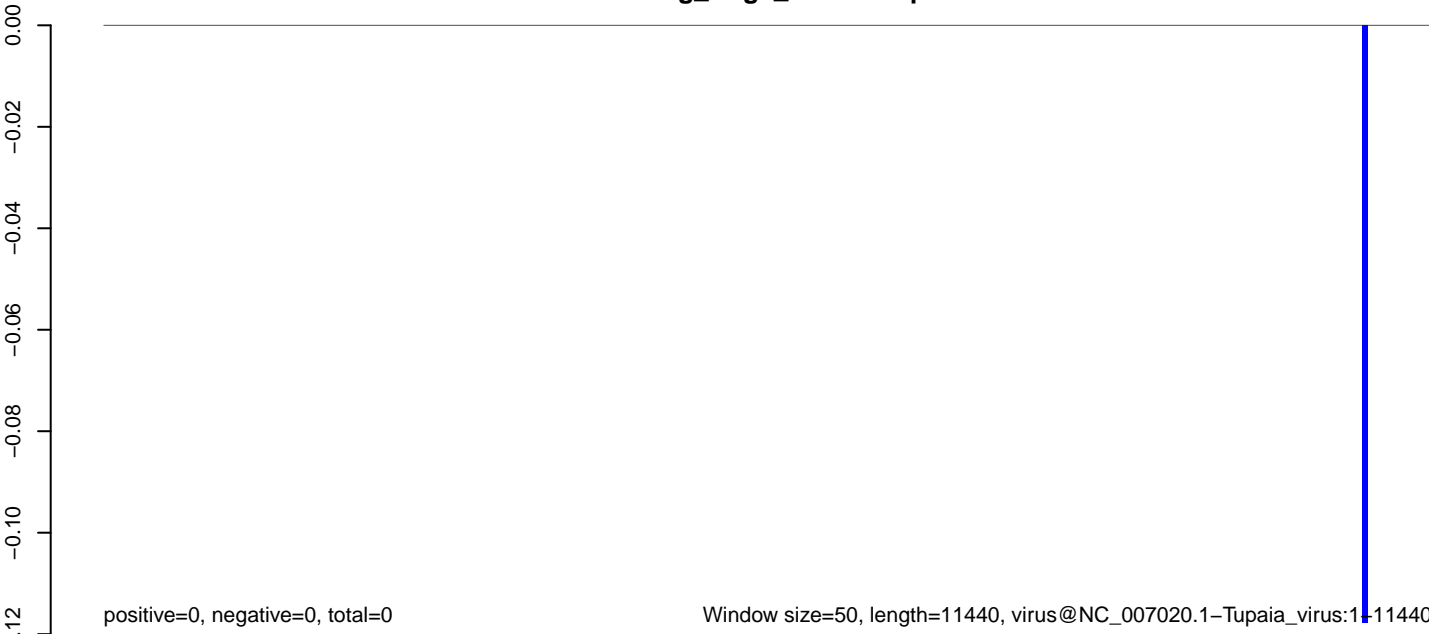
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



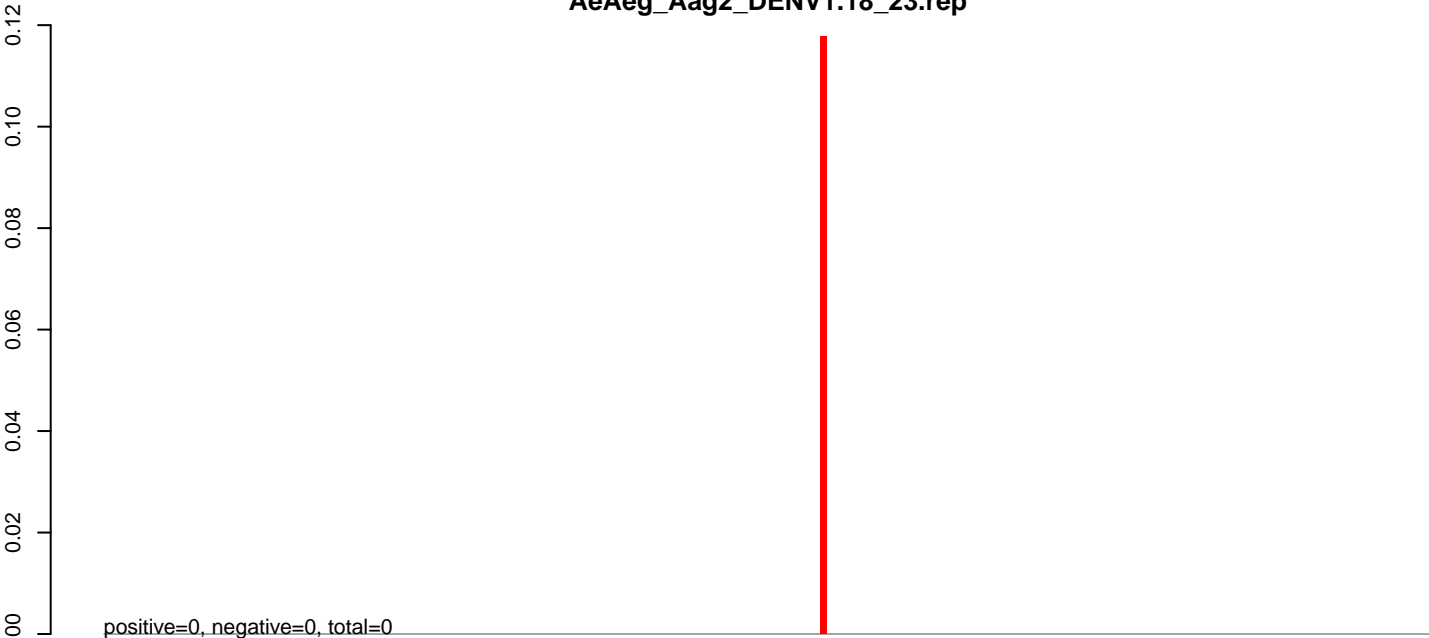
AeAeg_Aag2_DENV1.rep



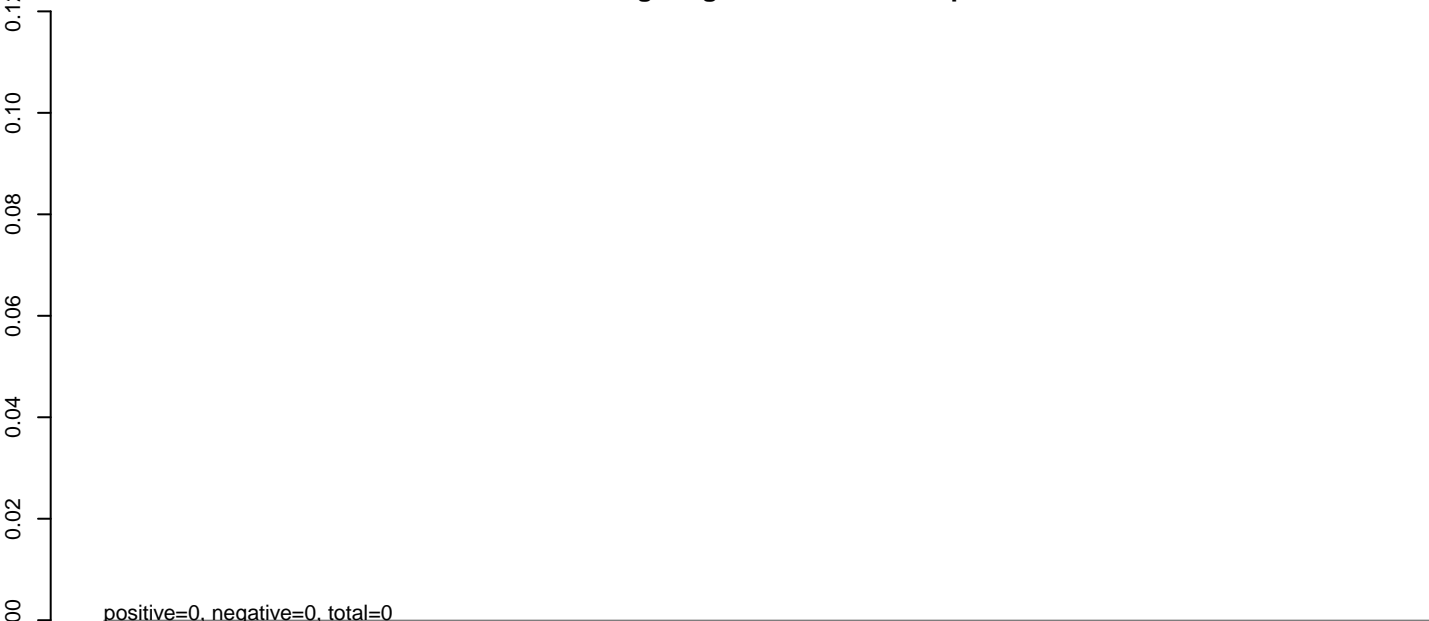
Window size=50, length=11440, virus@NC_007020.1-Tupaia_virus:1-11440

0 2000 4000 6000 8000 10000

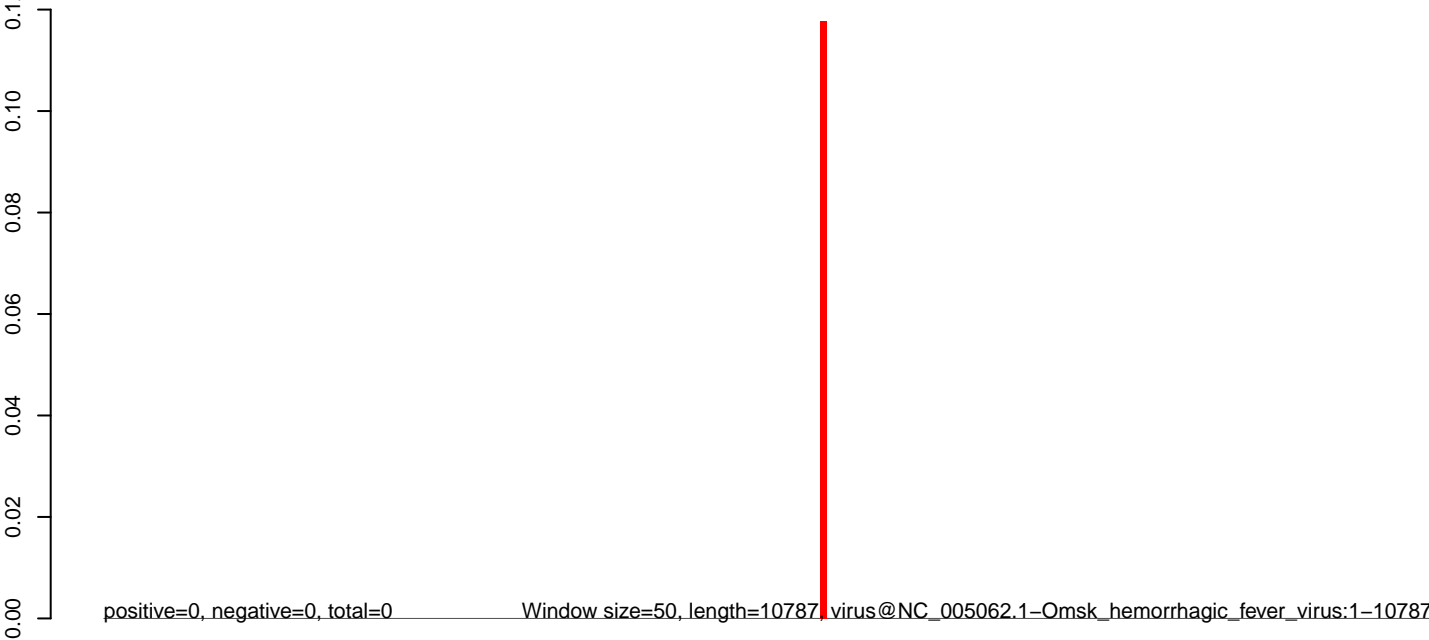
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



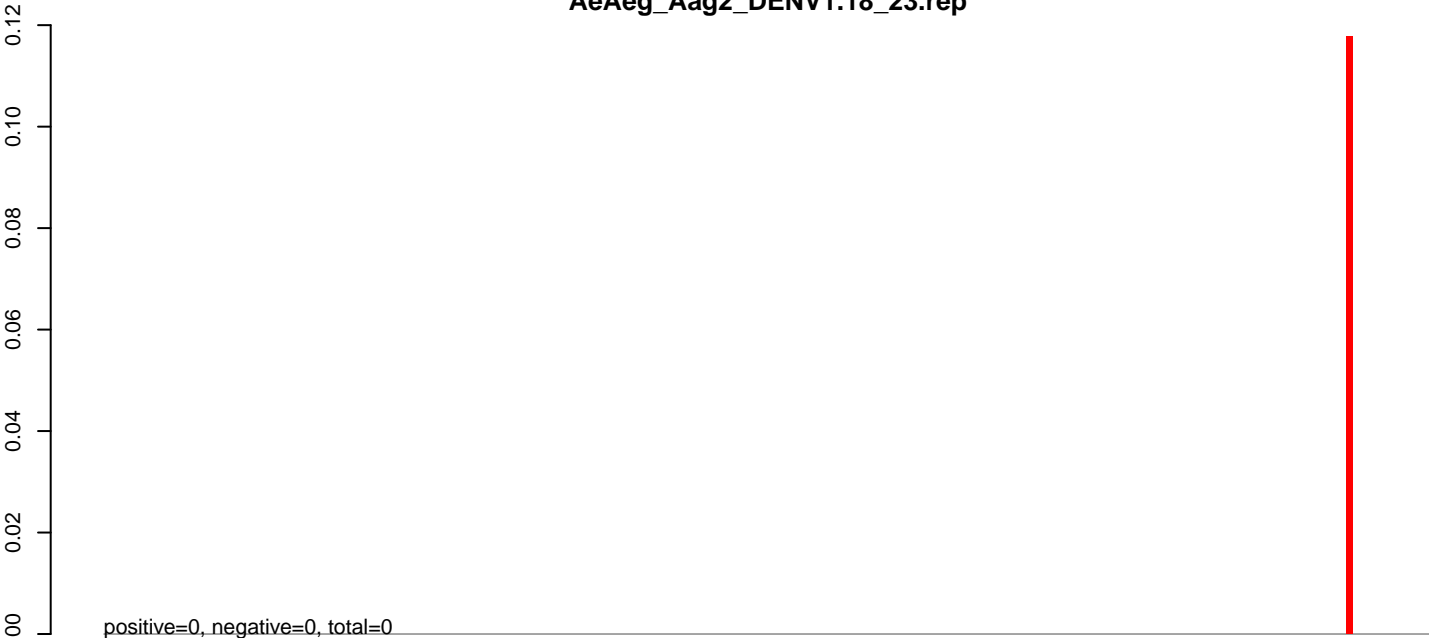
AeAeg_Aag2_DENV1.rep



0 2000 4000 6000 8000 10000

Window size=50, length=10787, virus@NC_005062.1-Omsk_hemorrhagic_fever_virus:1-10787

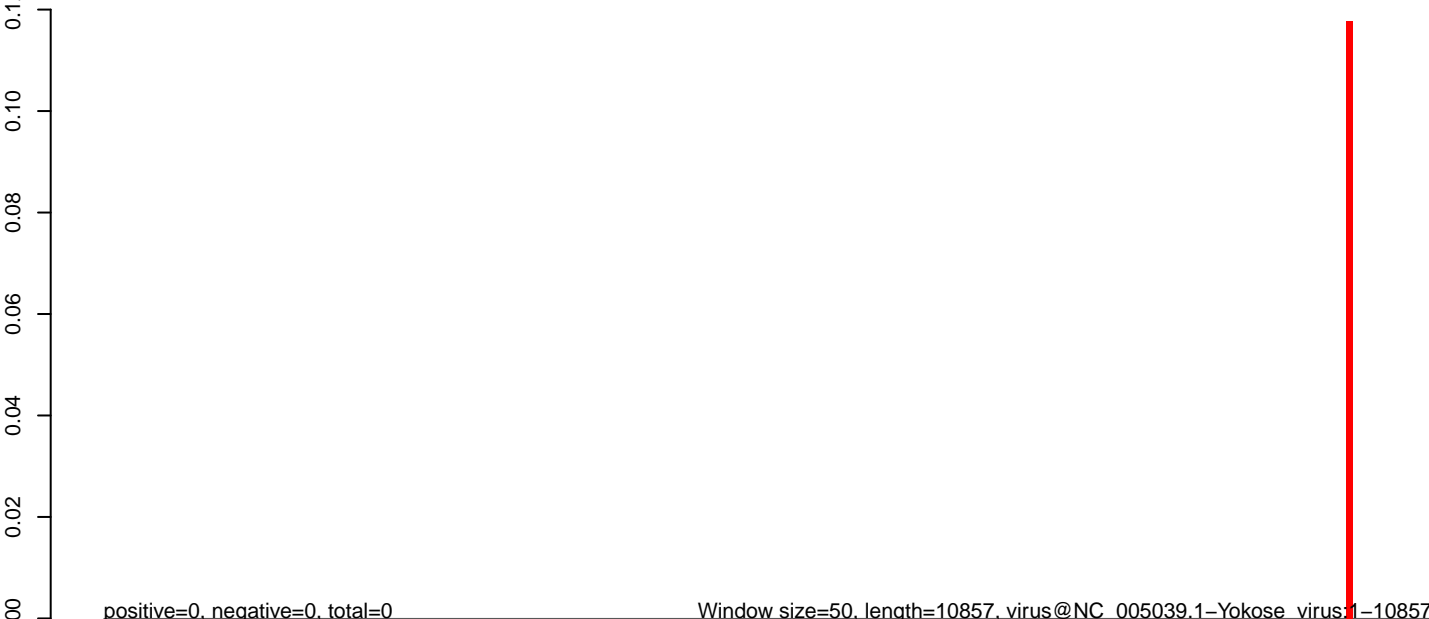
AeAeg_Aag2_DENV1.18_23.rep



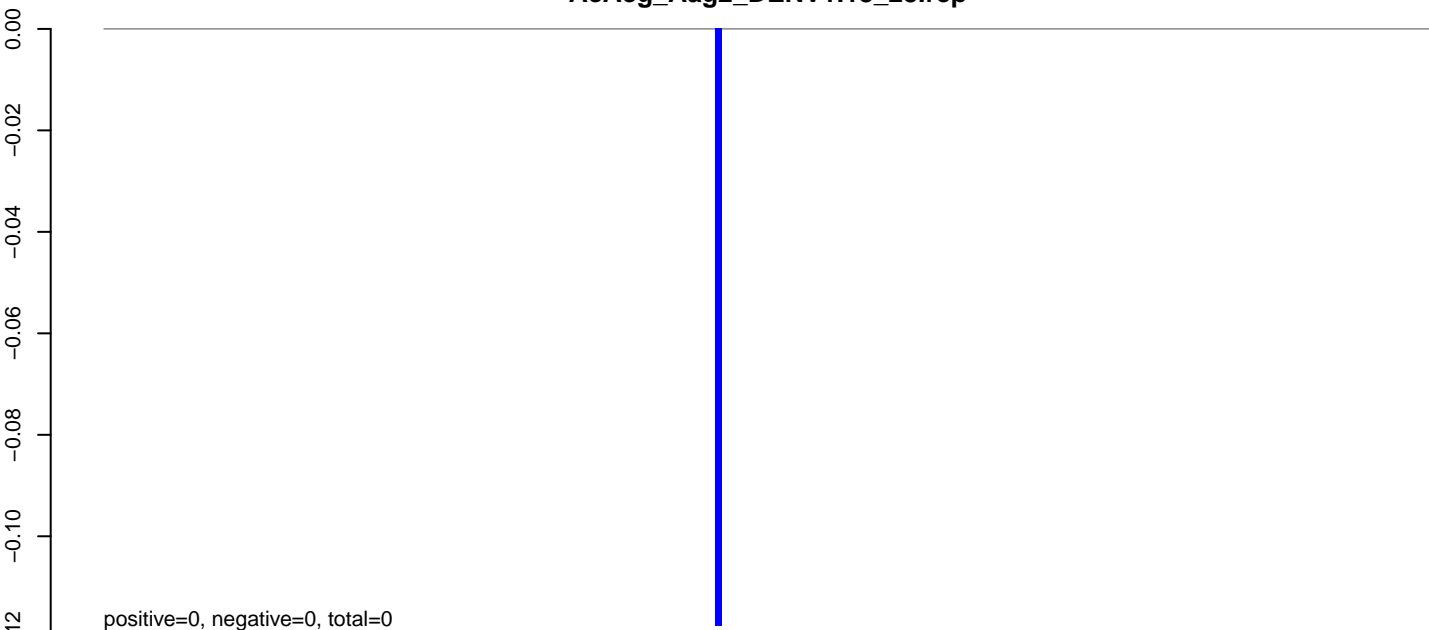
AeAeg_Aag2_DENV1.24_35.rep



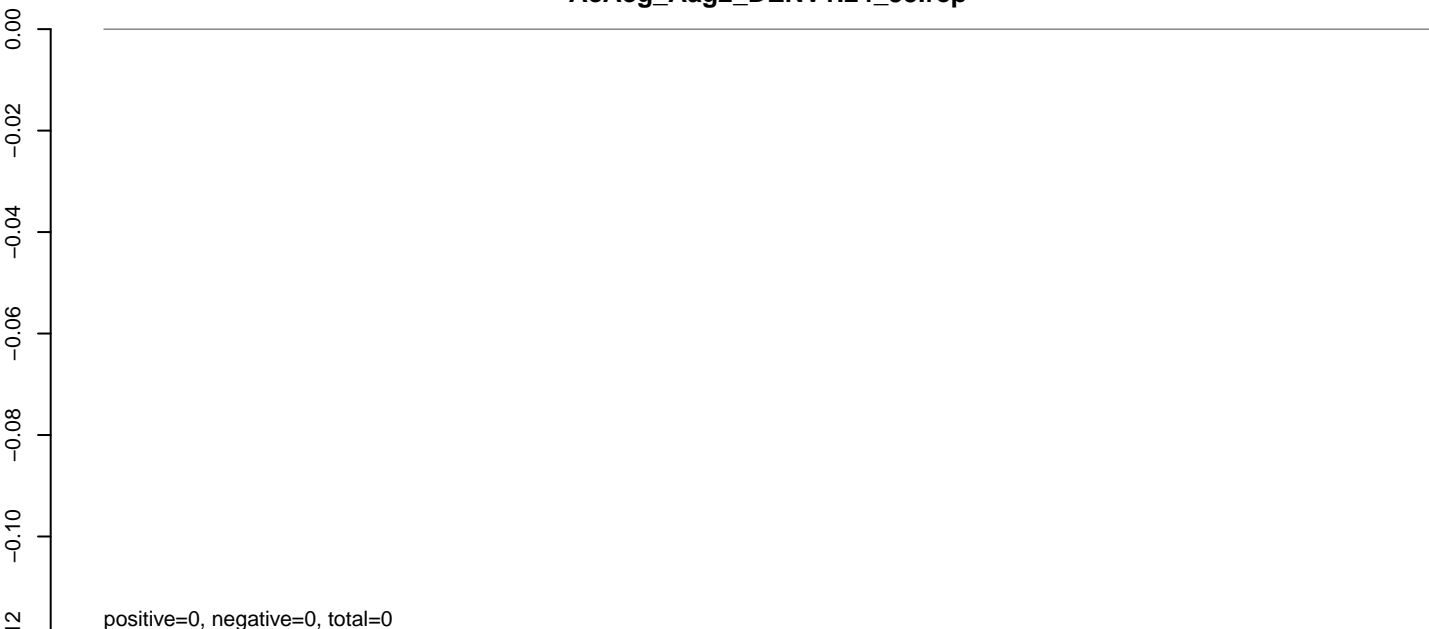
AeAeg_Aag2_DENV1.rep



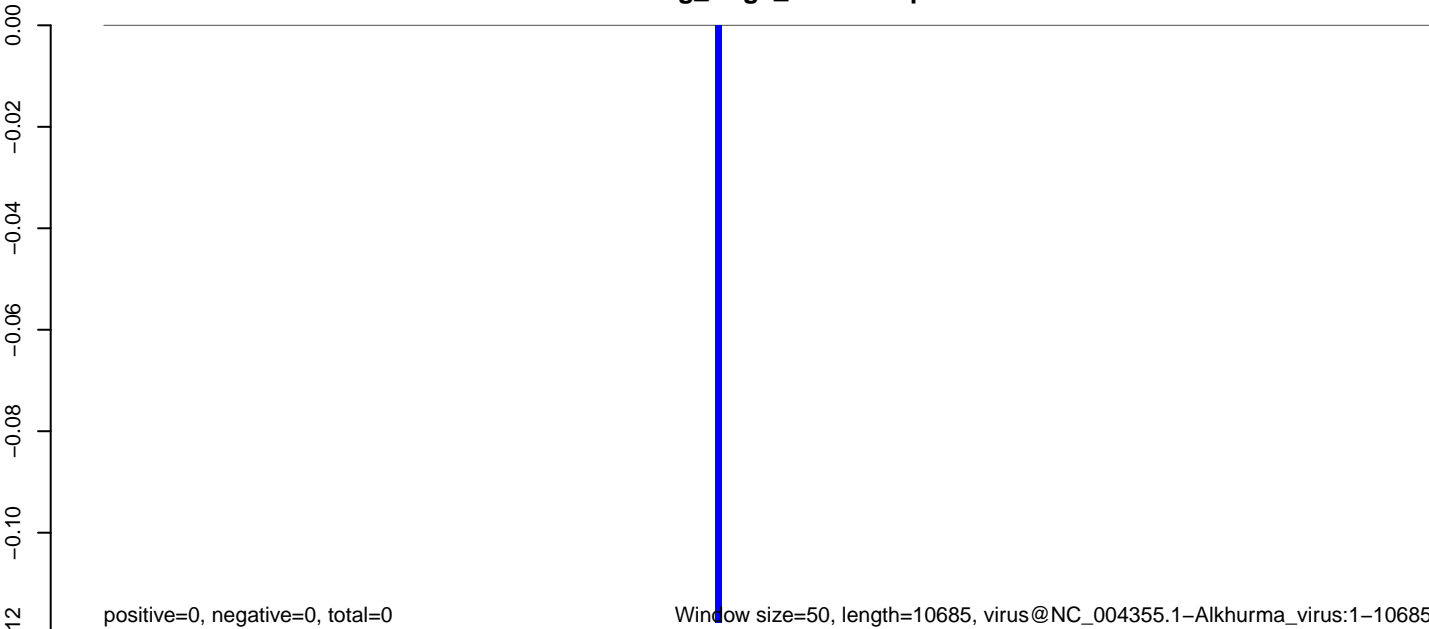
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

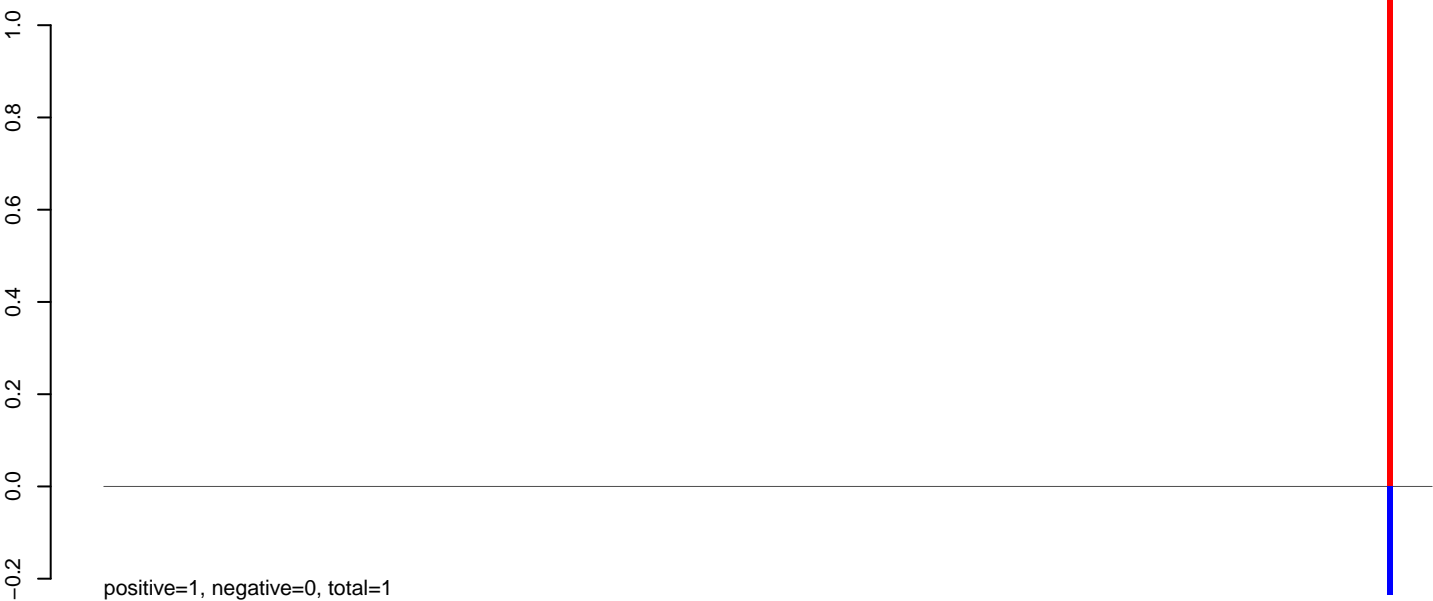


AeAeg_Aag2_DENV1.rep

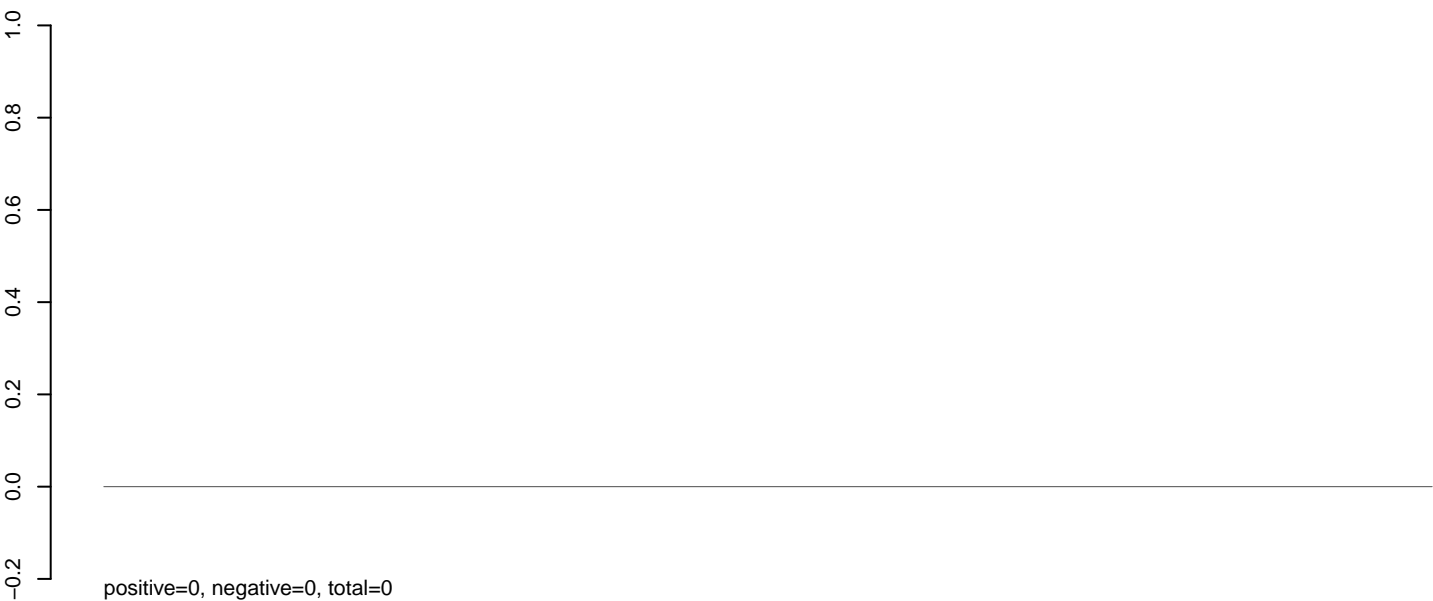


0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV1.18_23.rep



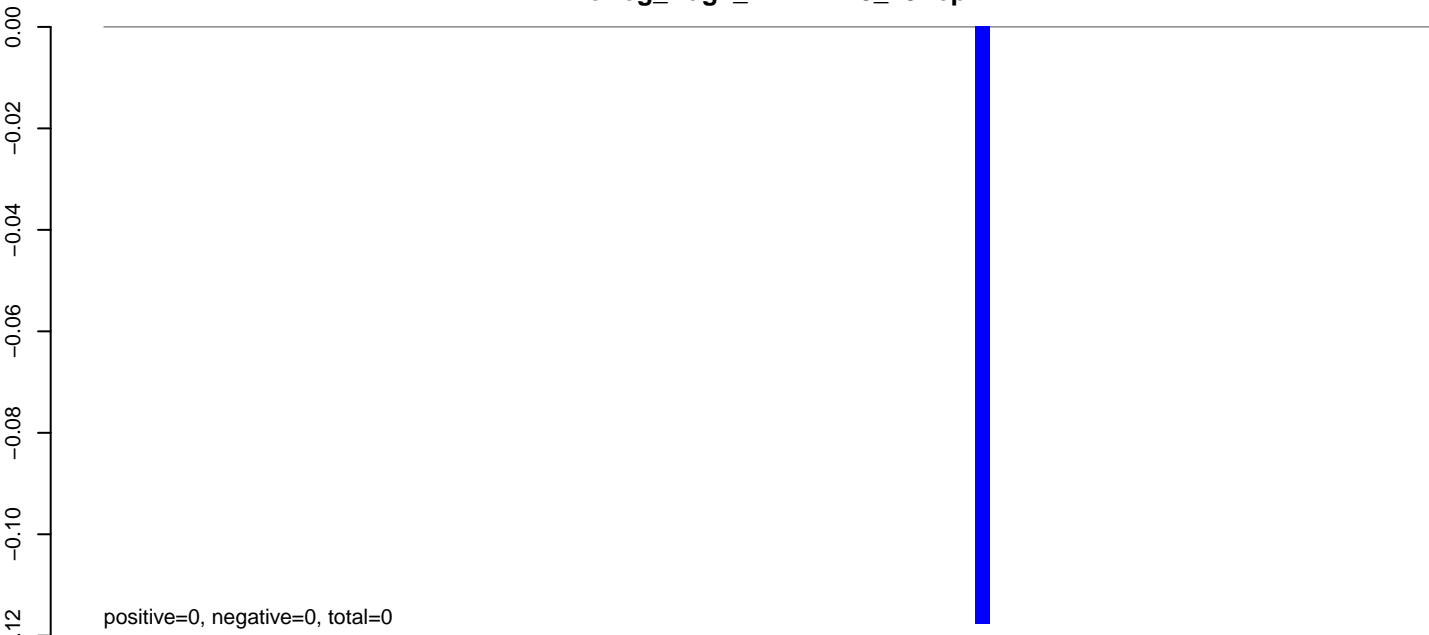
AeAeg_Aag2_DENV1.24_35.rep



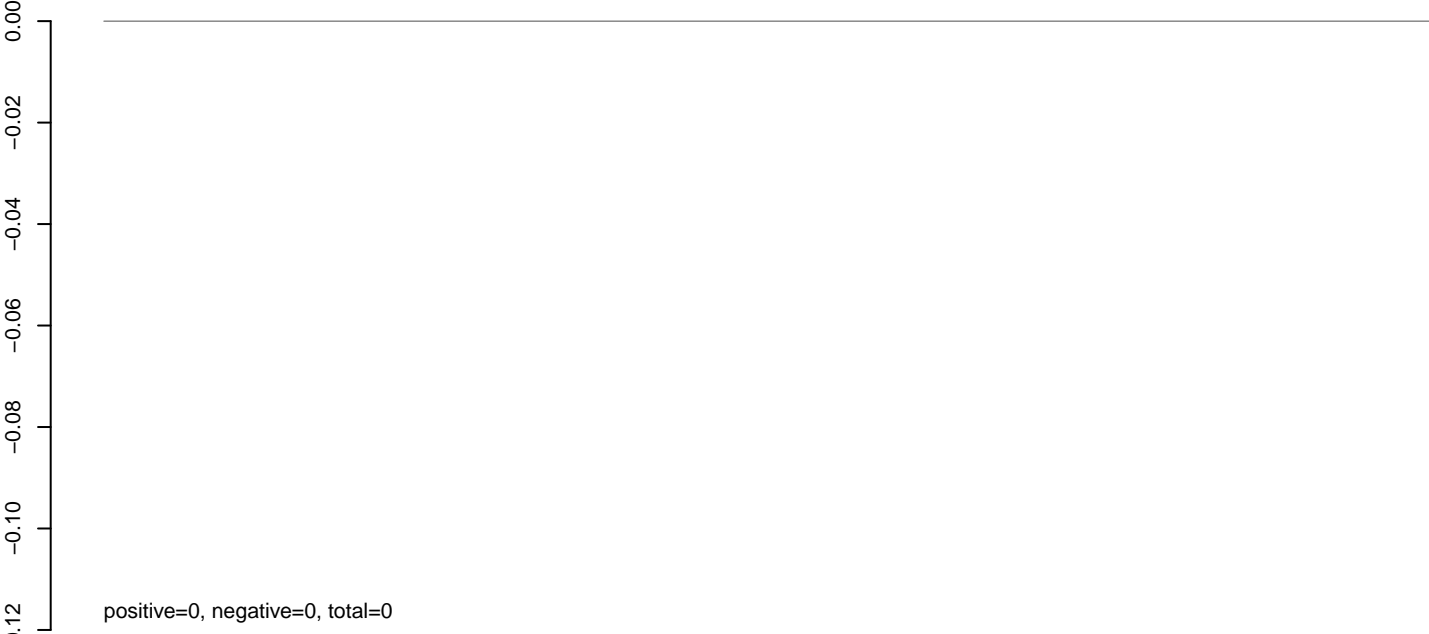
AeAeg_Aag2_DENV1.rep



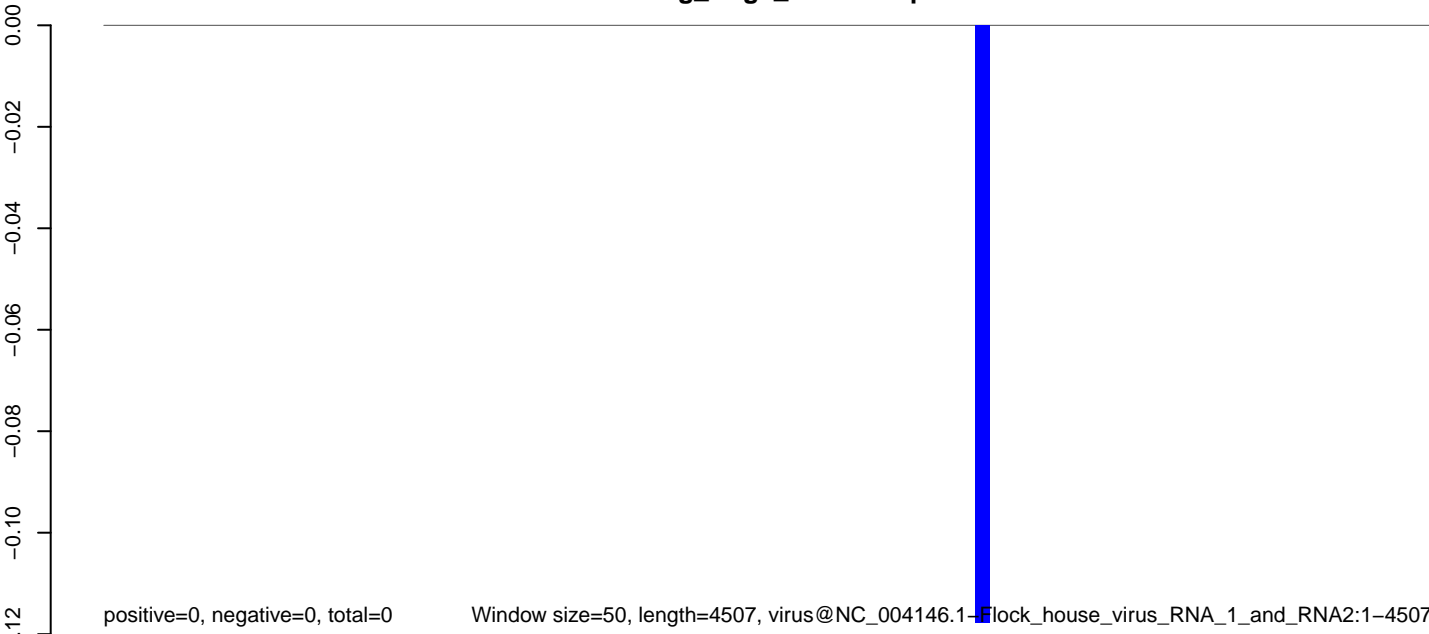
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



AeAeg_Aag2_DENV1.rep



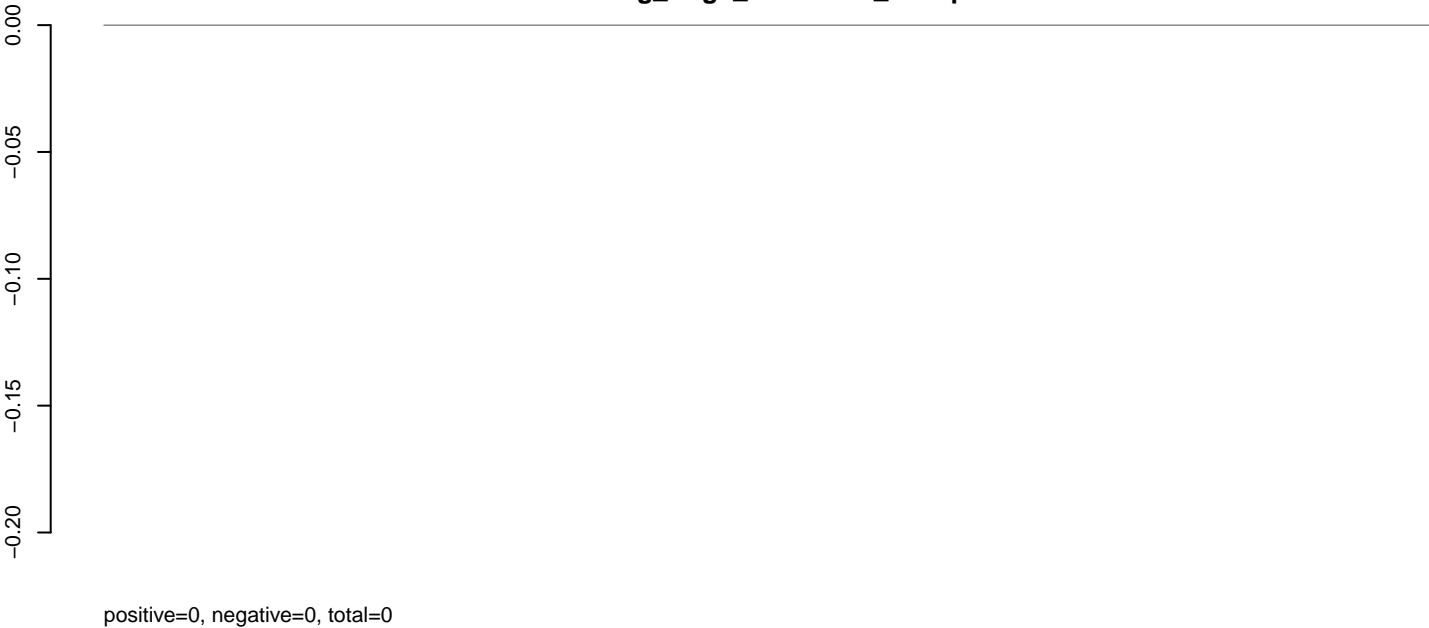
Window size=50, length=4507, virus@NC_004146.1-Flock_house_virus_RNA_1_and_RNA2:1-4507

0 1000 2000 3000 4000

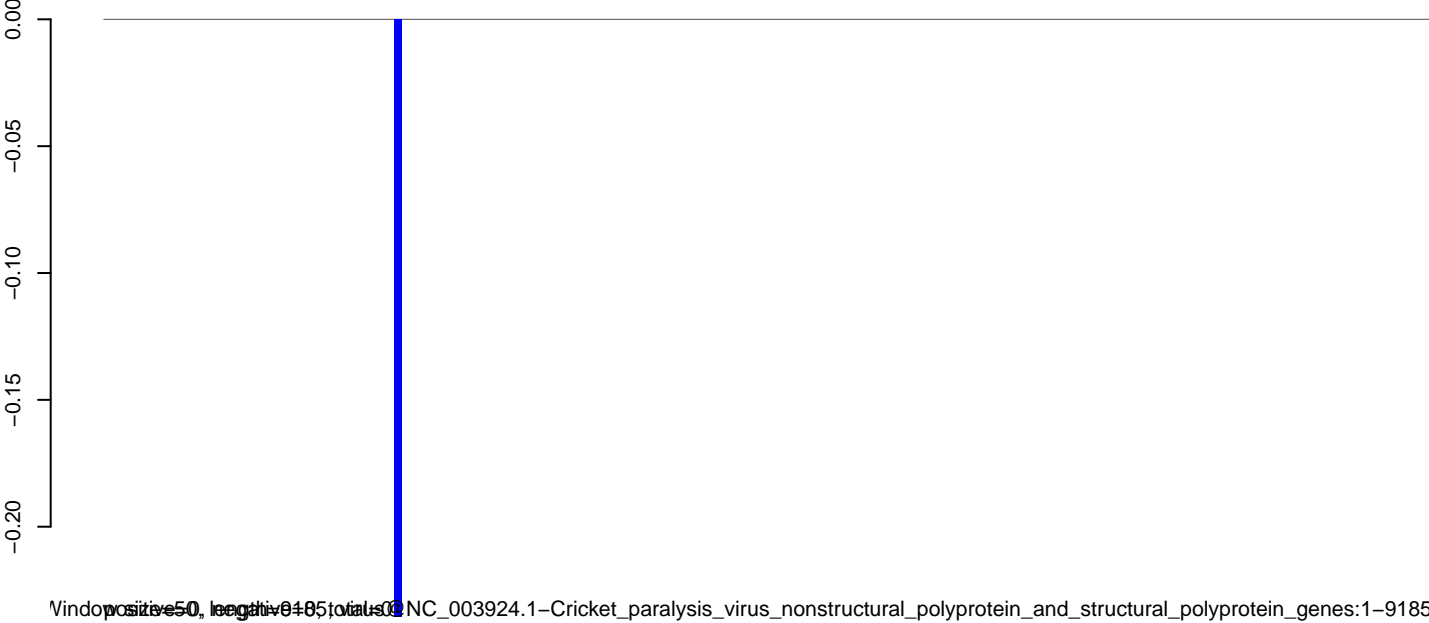
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



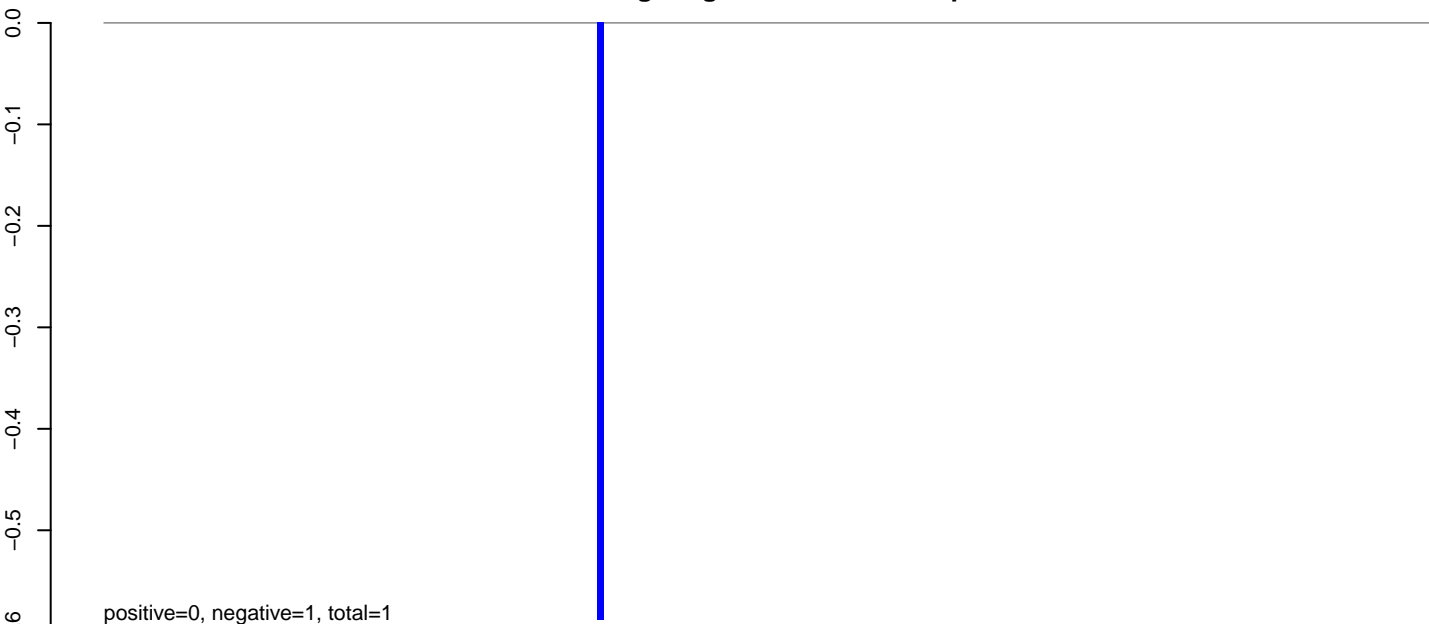
AeAeg_Aag2_DENV1.rep



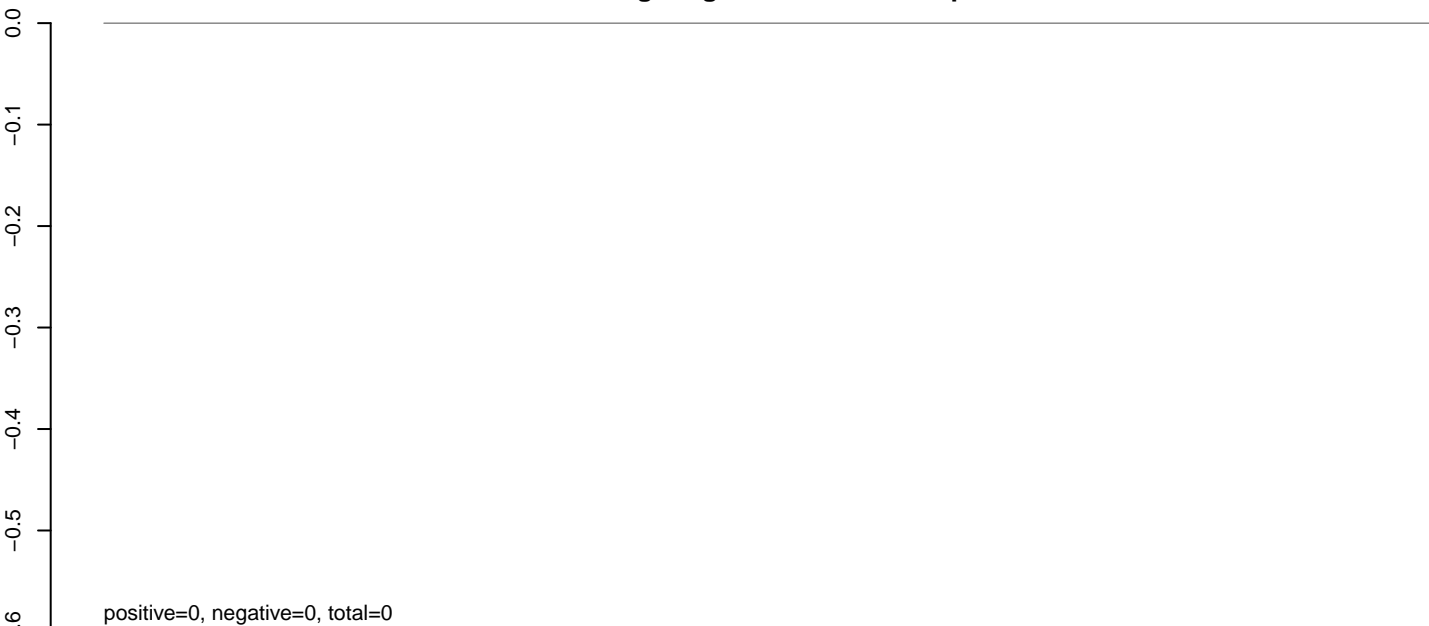
Vindon positive=50, negative=185, total=235 NC_003924.1-Cricket_paralysis_virus_nonstructural_polyprotein_and_structural_polyprotein_genes:1-9185

0 2000 4000 6000 8000

AeAeg_Aag2_DENV1.18_23.rep



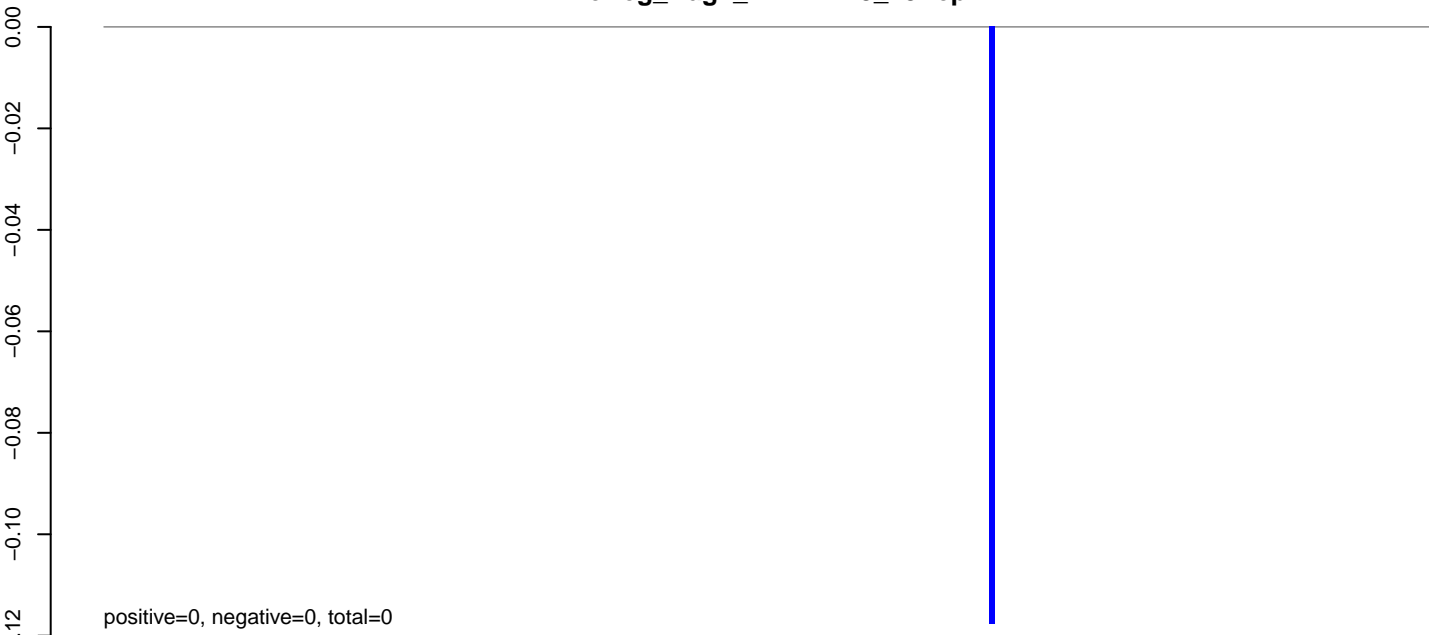
AeAeg_Aag2_DENV1.24_35.rep



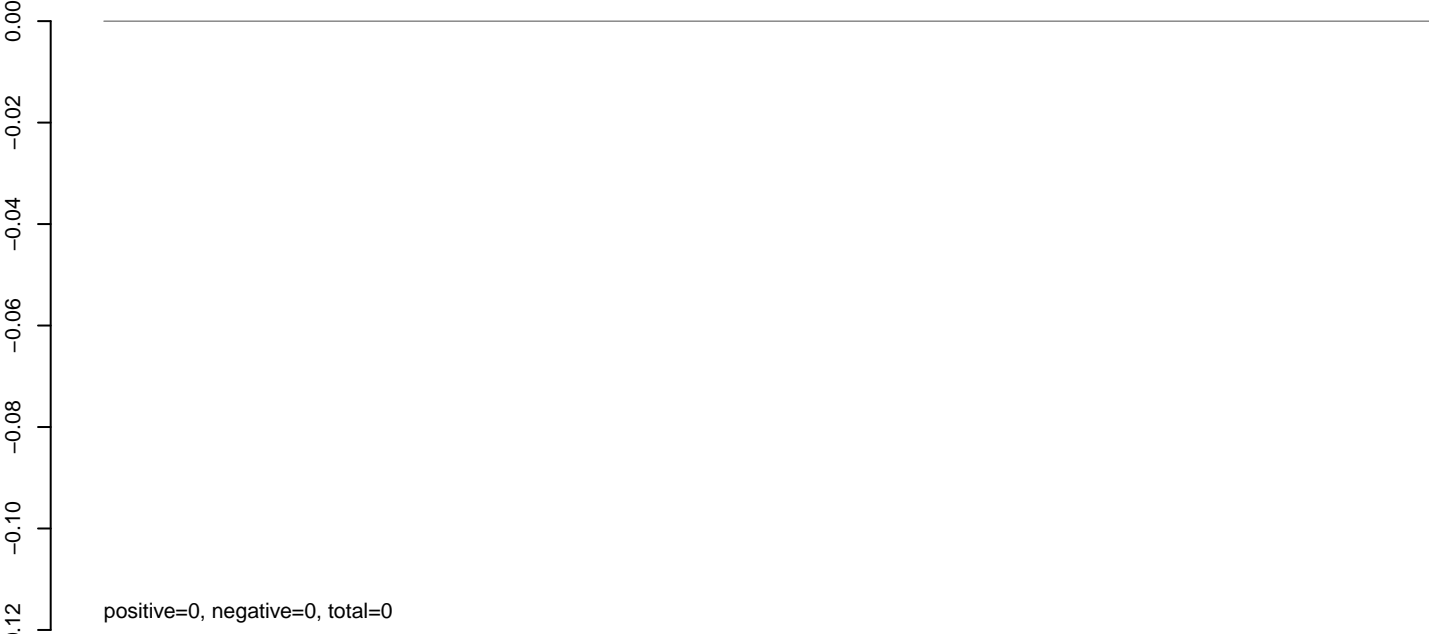
AeAeg_Aag2_DENV1.rep



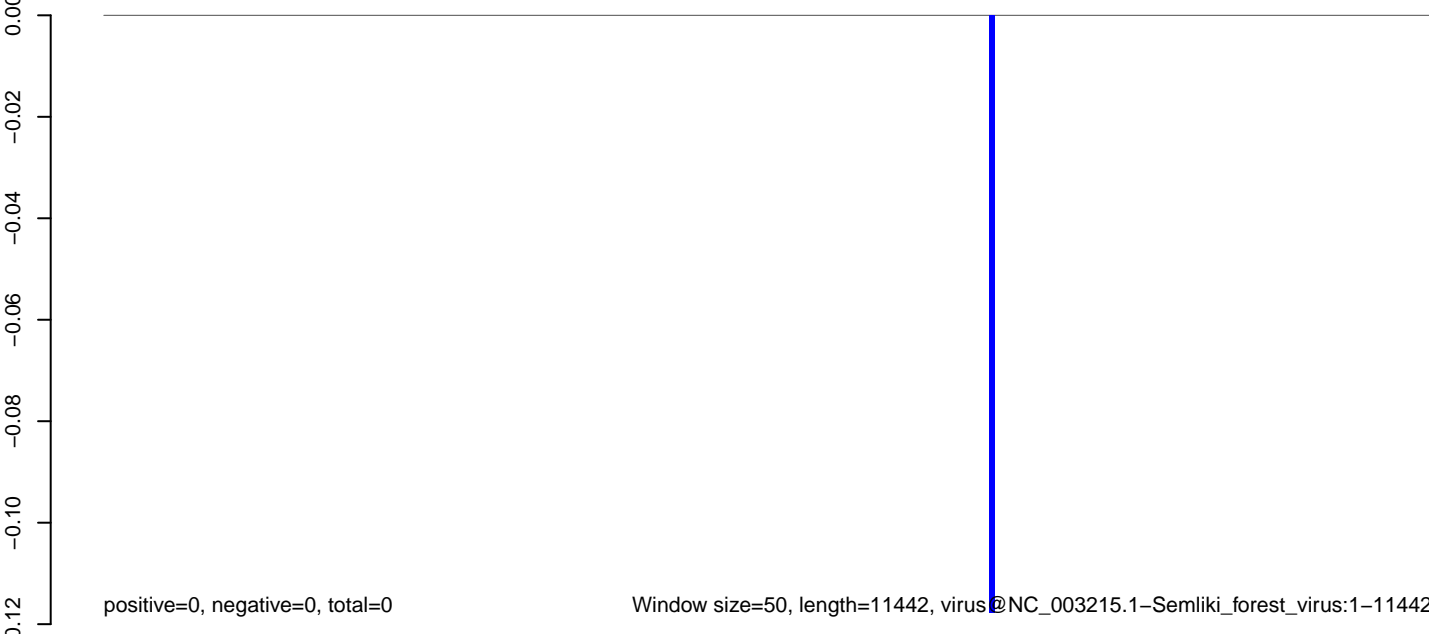
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

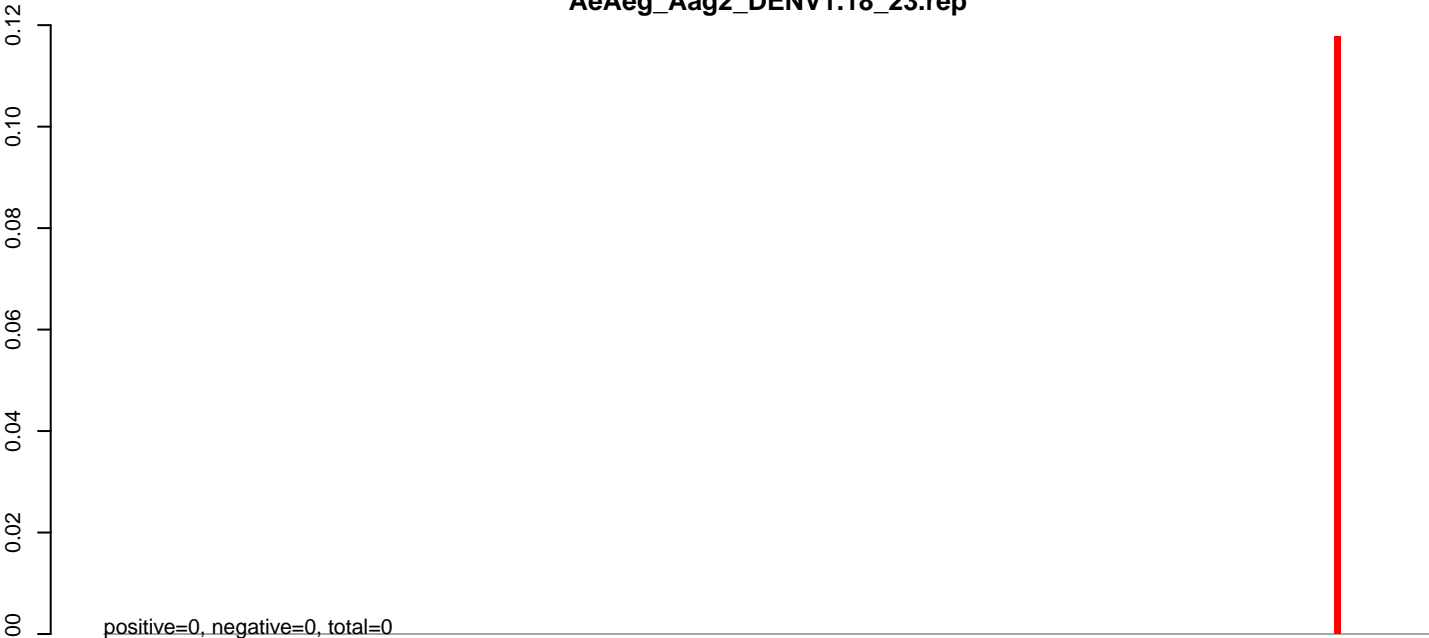


AeAeg_Aag2_DENV1.rep

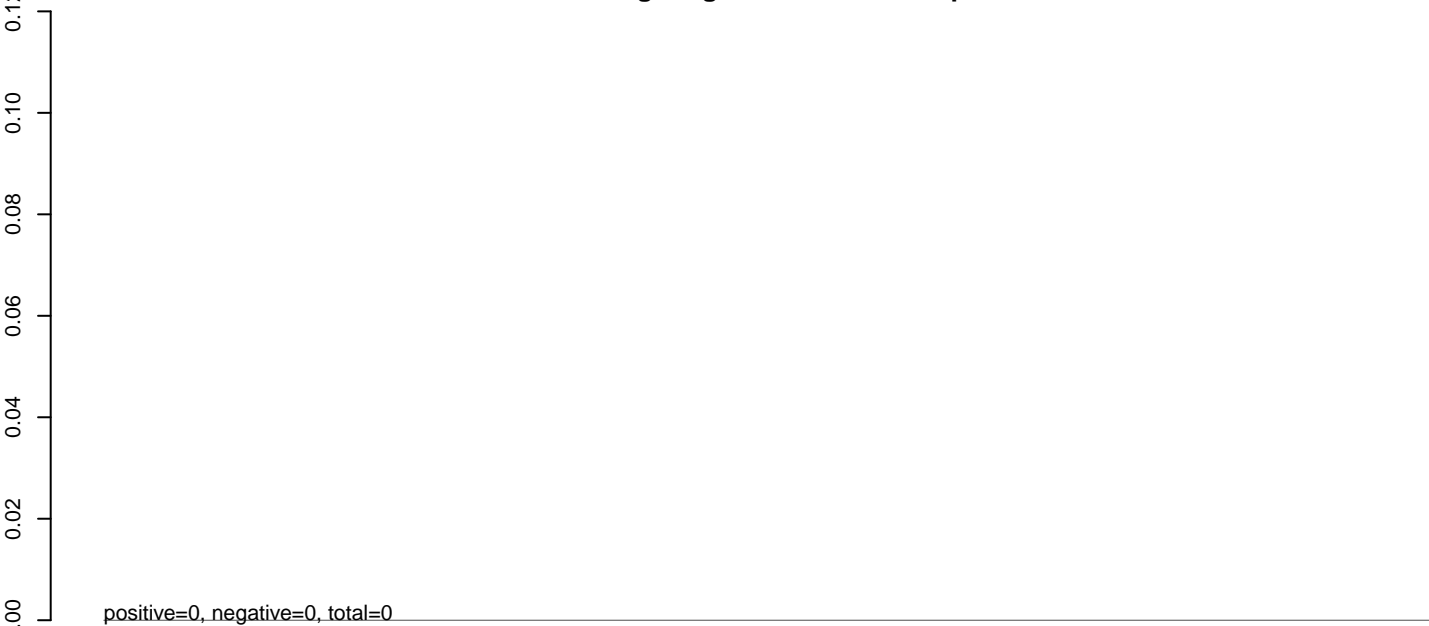


0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

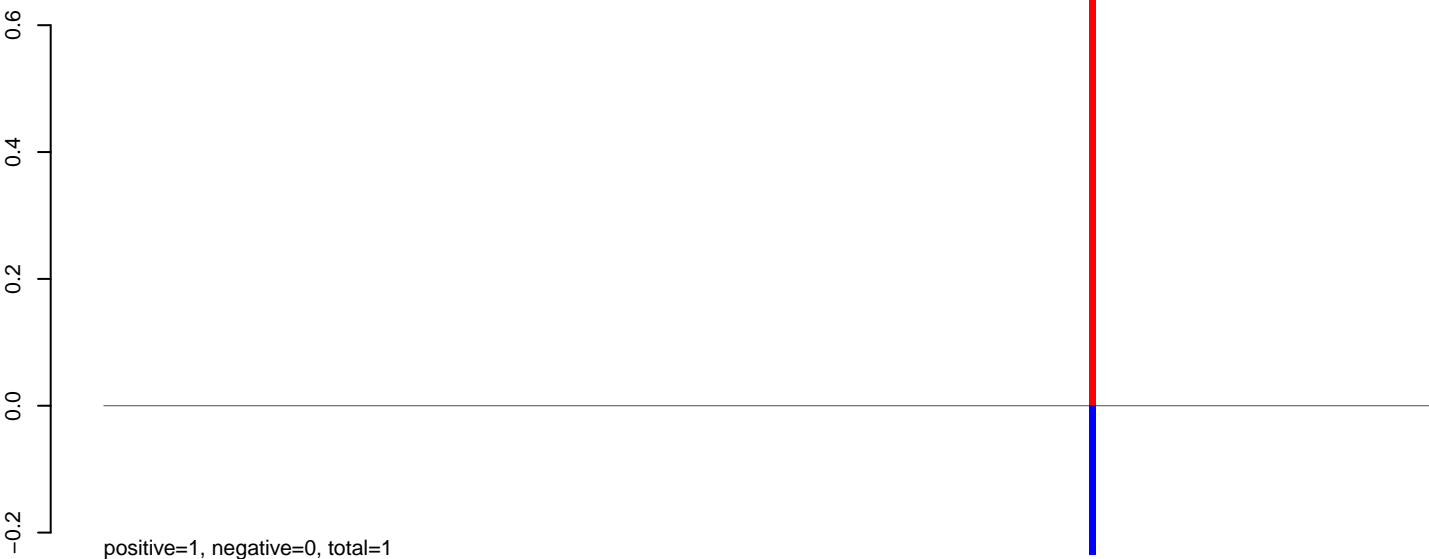


AeAeg_Aag2_DENV1.rep

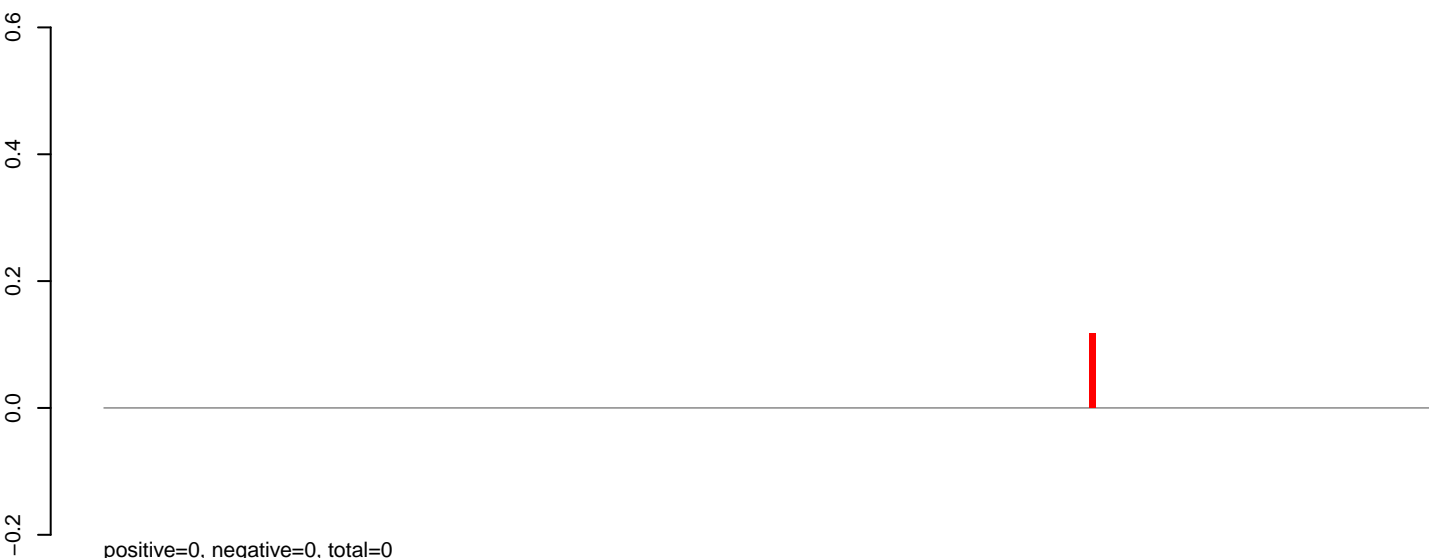


0 2000 4000 6000 8000 10000

AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

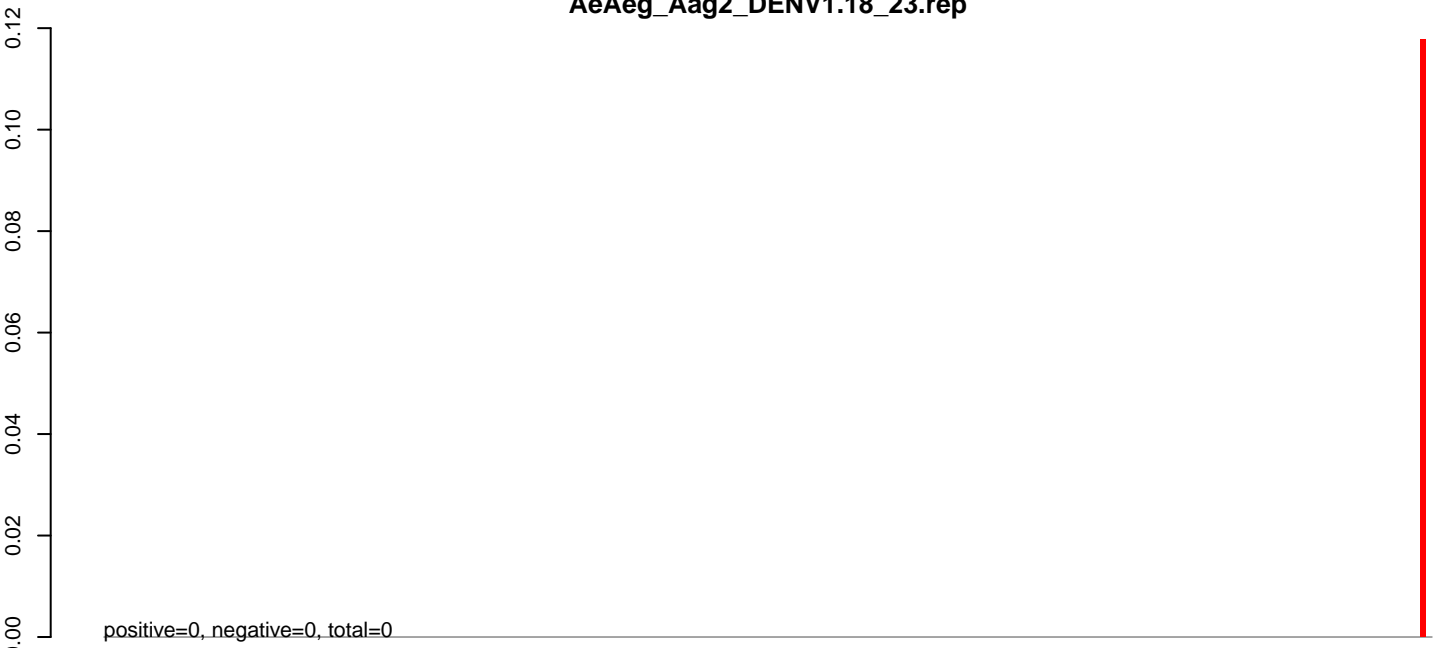


AeAeg_Aag2_DENV1.rep

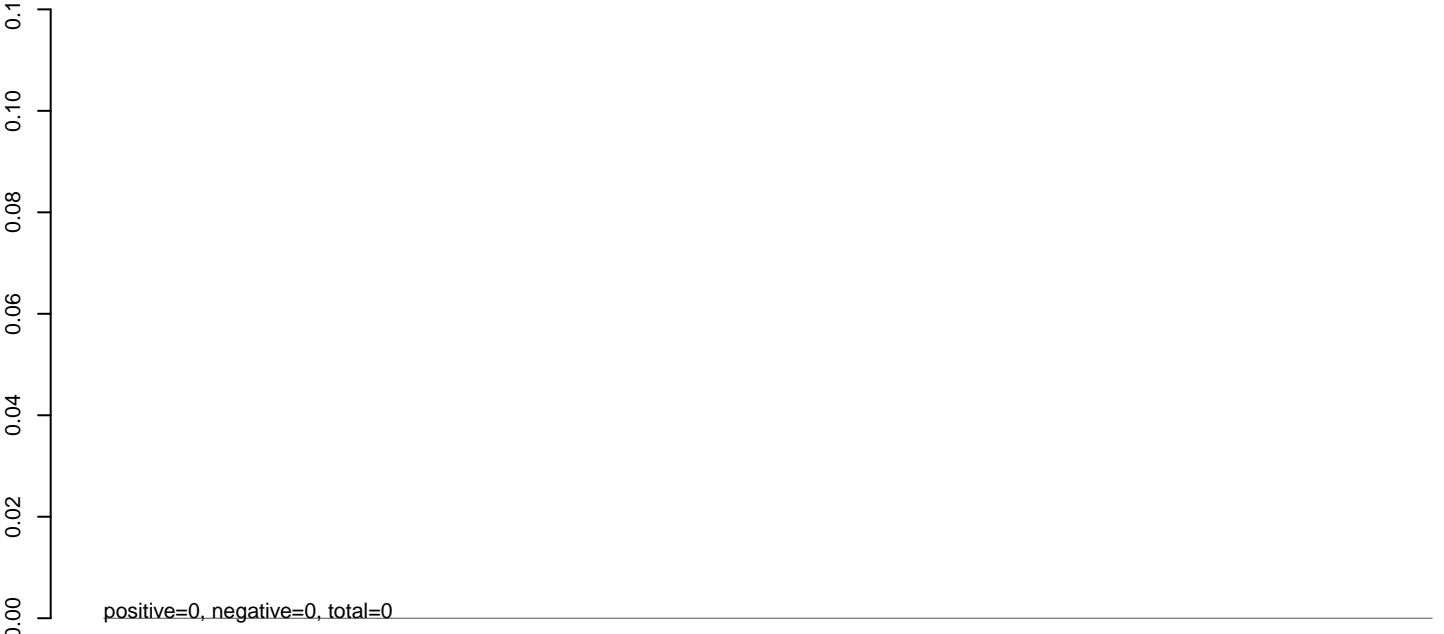


0 2000 4000 6000 8000 10000

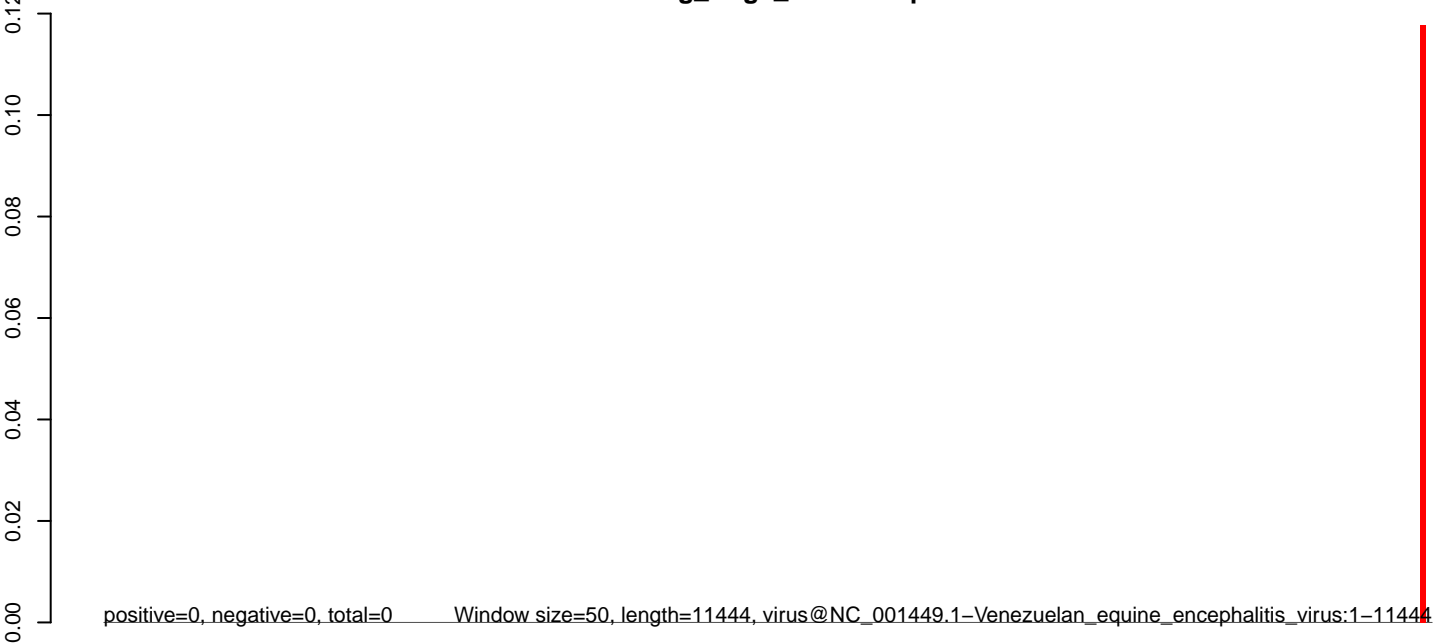
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

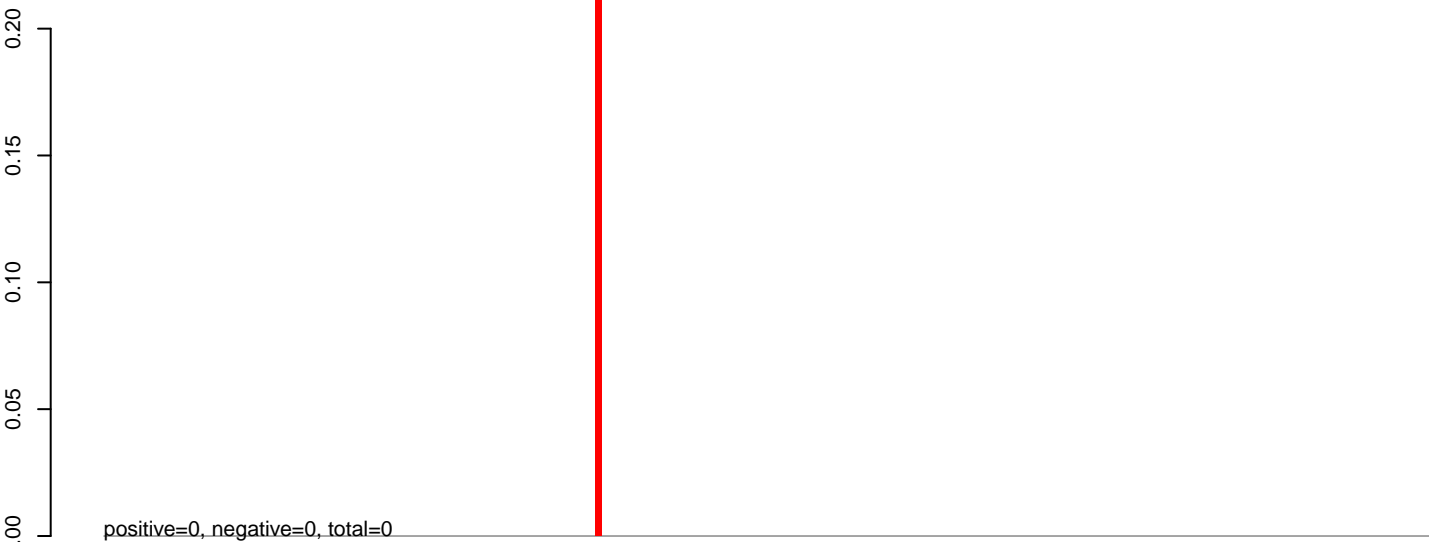


AeAeg_Aag2_DENV1.rep

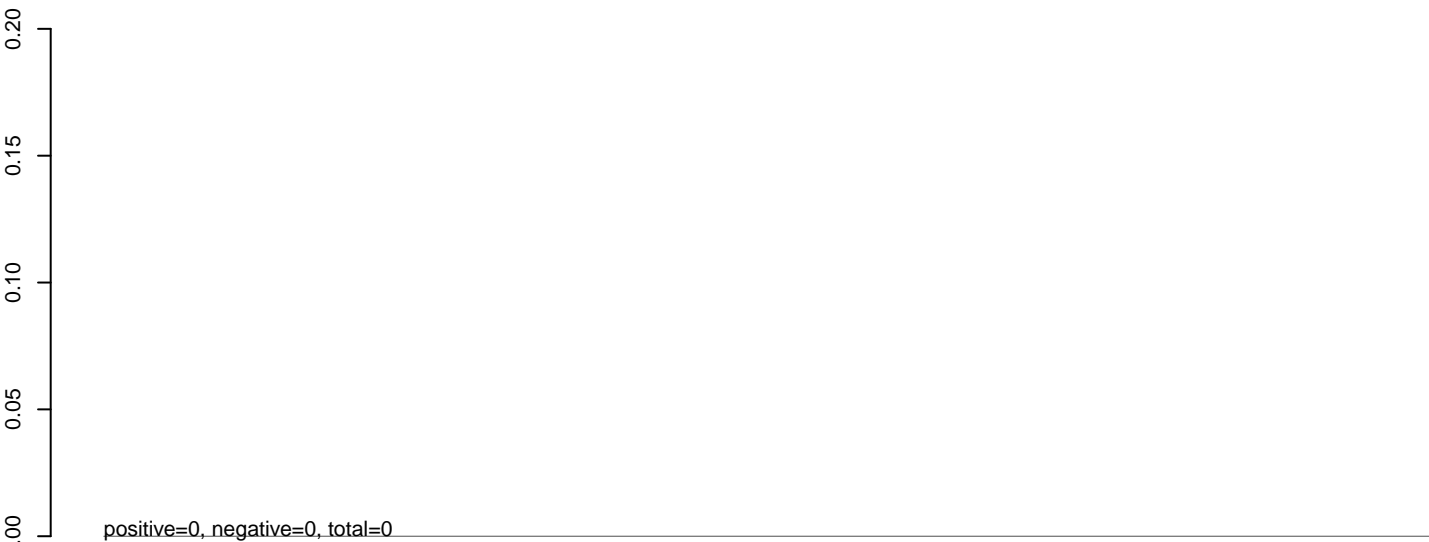


0 2000 4000 6000 8000 10000

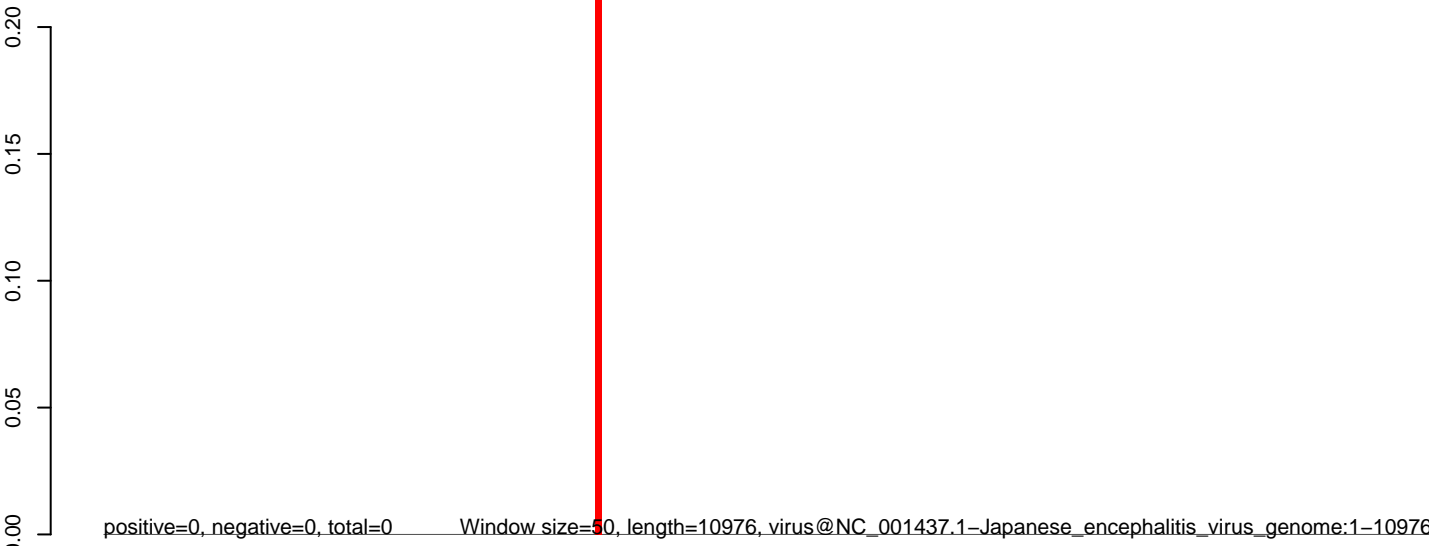
AeAeg_Aag2_DENV1.18_23.rep



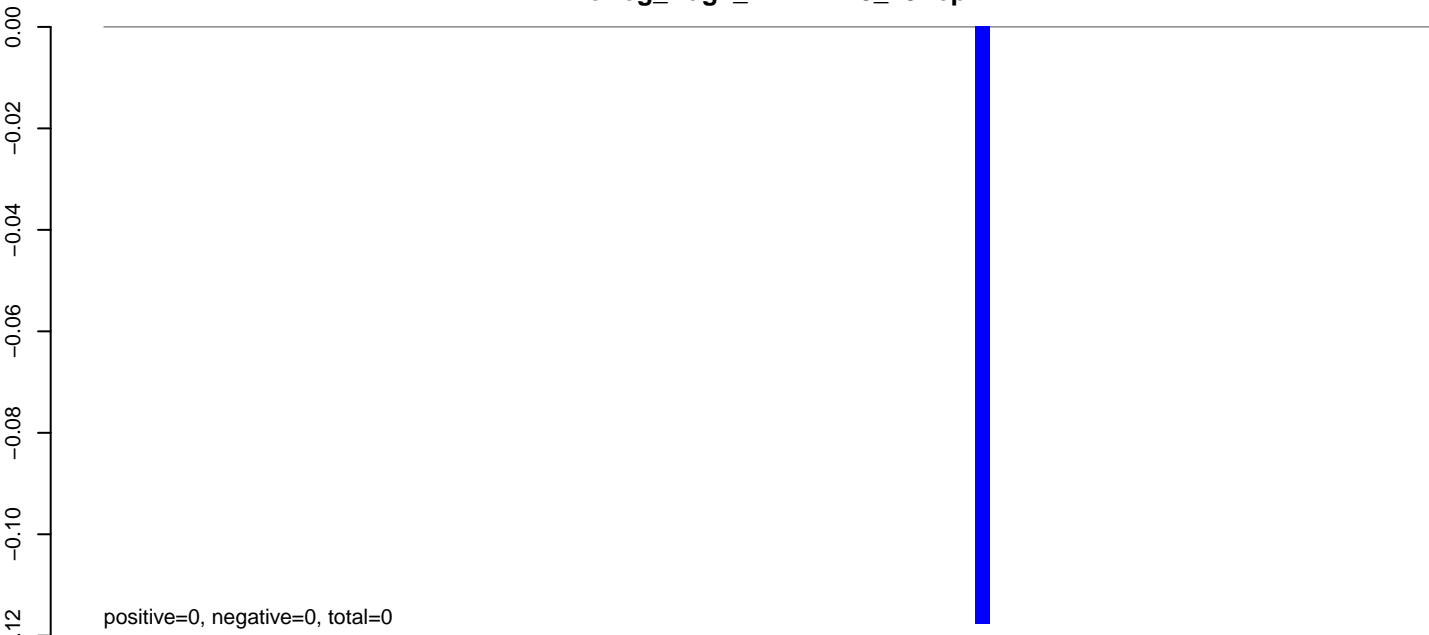
AeAeg_Aag2_DENV1.24_35.rep



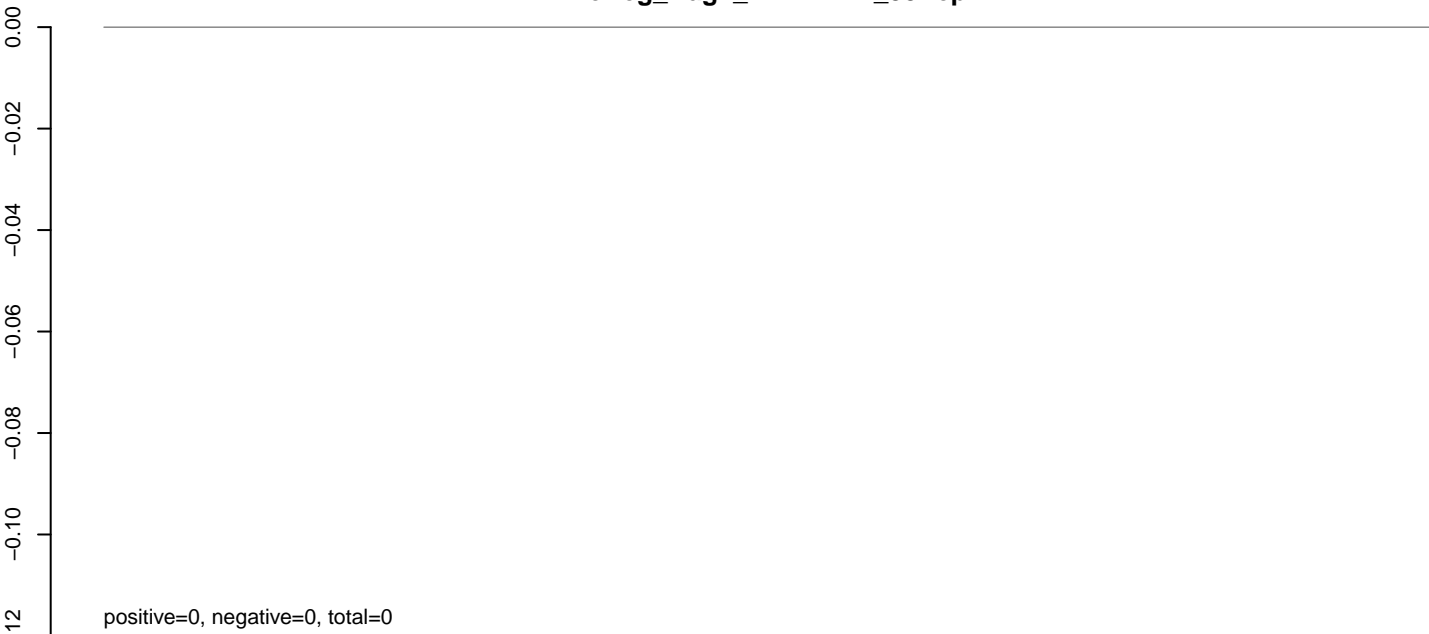
AeAeg_Aag2_DENV1.rep



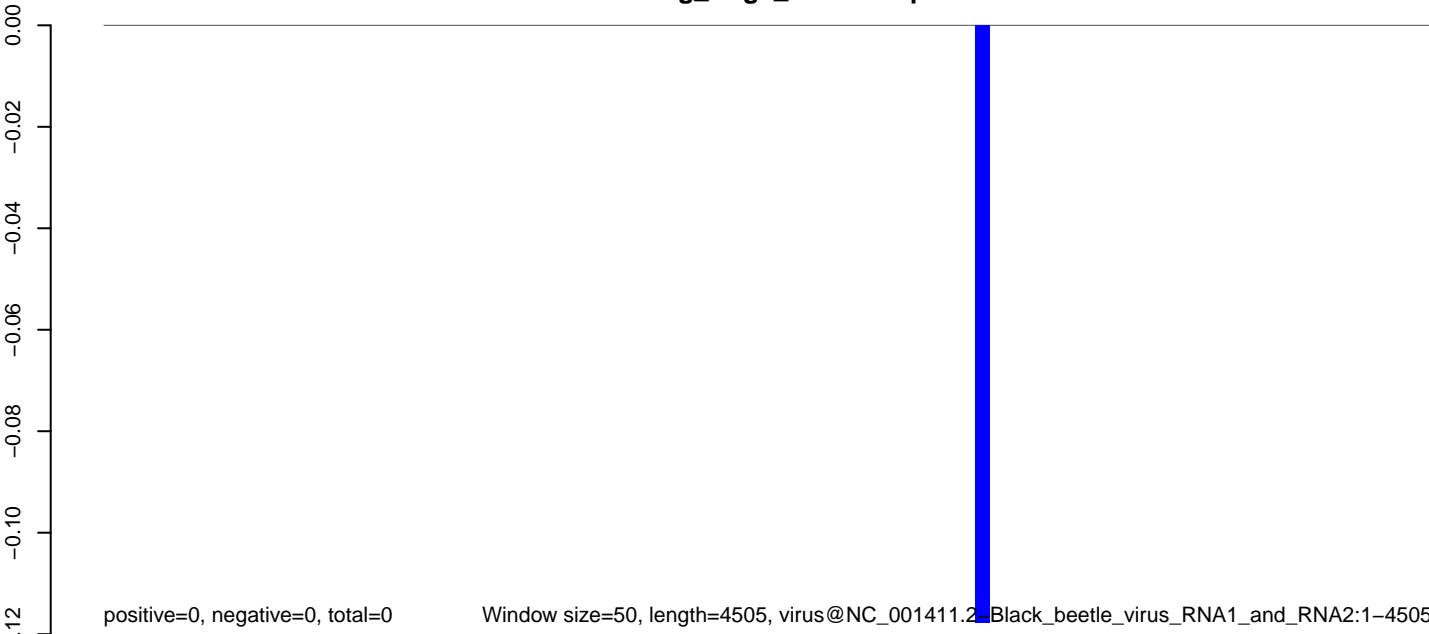
AeAeg_Aag2_DENV1.18_23.rep



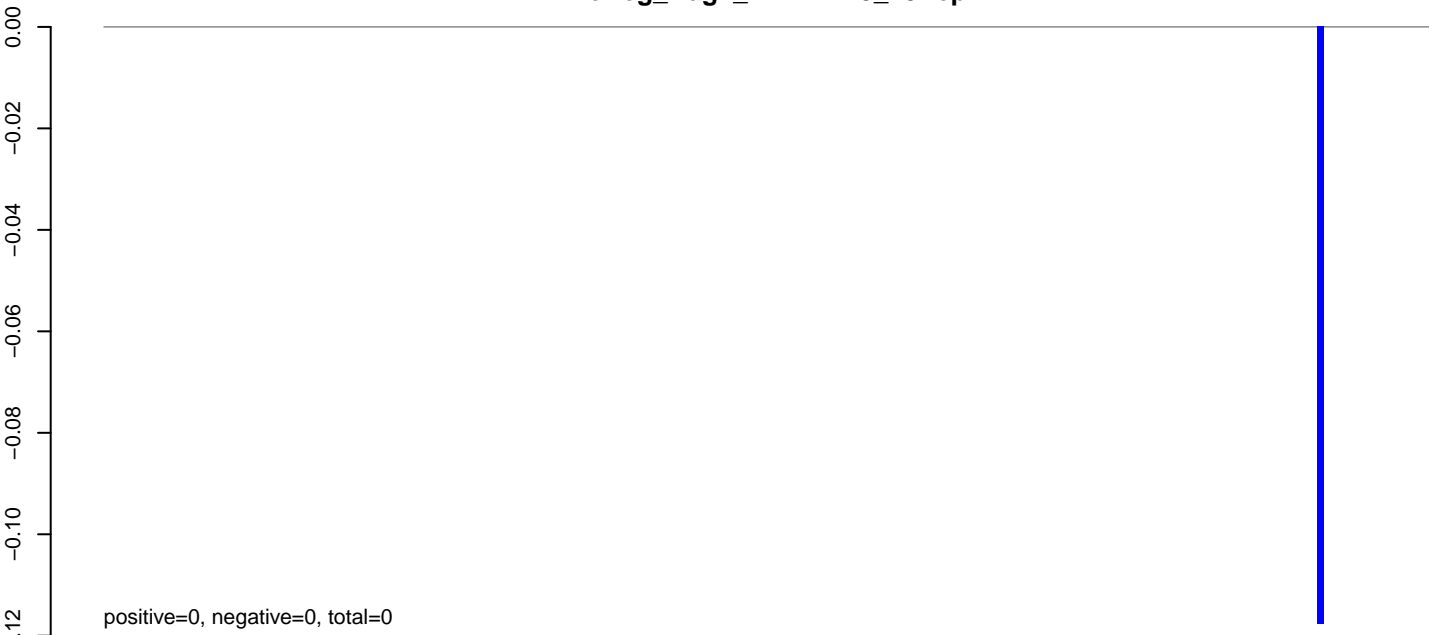
AeAeg_Aag2_DENV1.24_35.rep



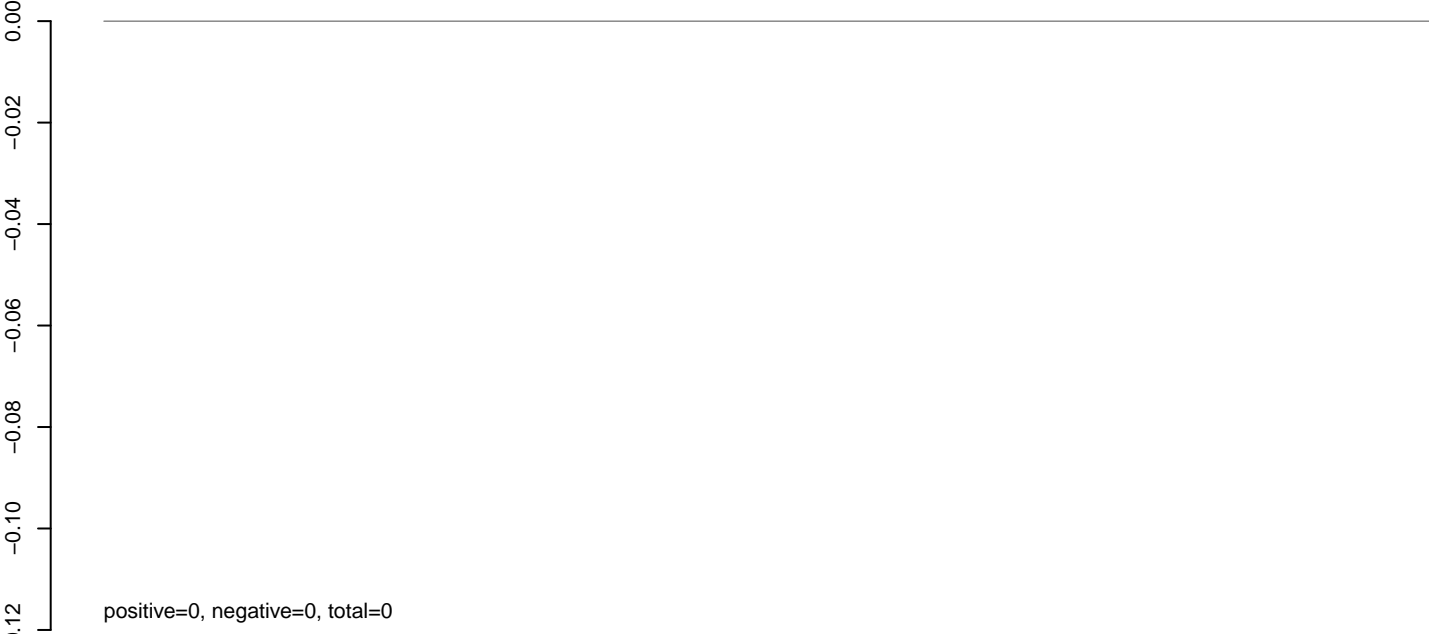
AeAeg_Aag2_DENV1.rep



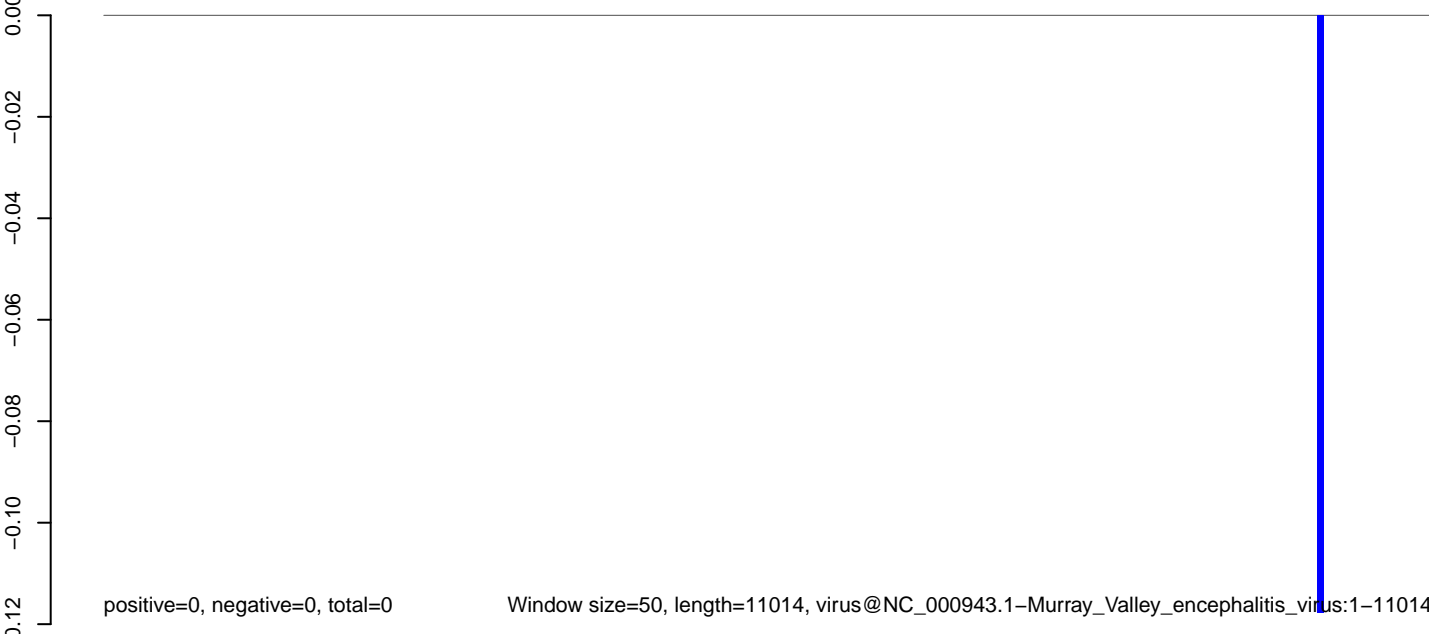
AeAeg_Aag2_DENV1.18_23.rep



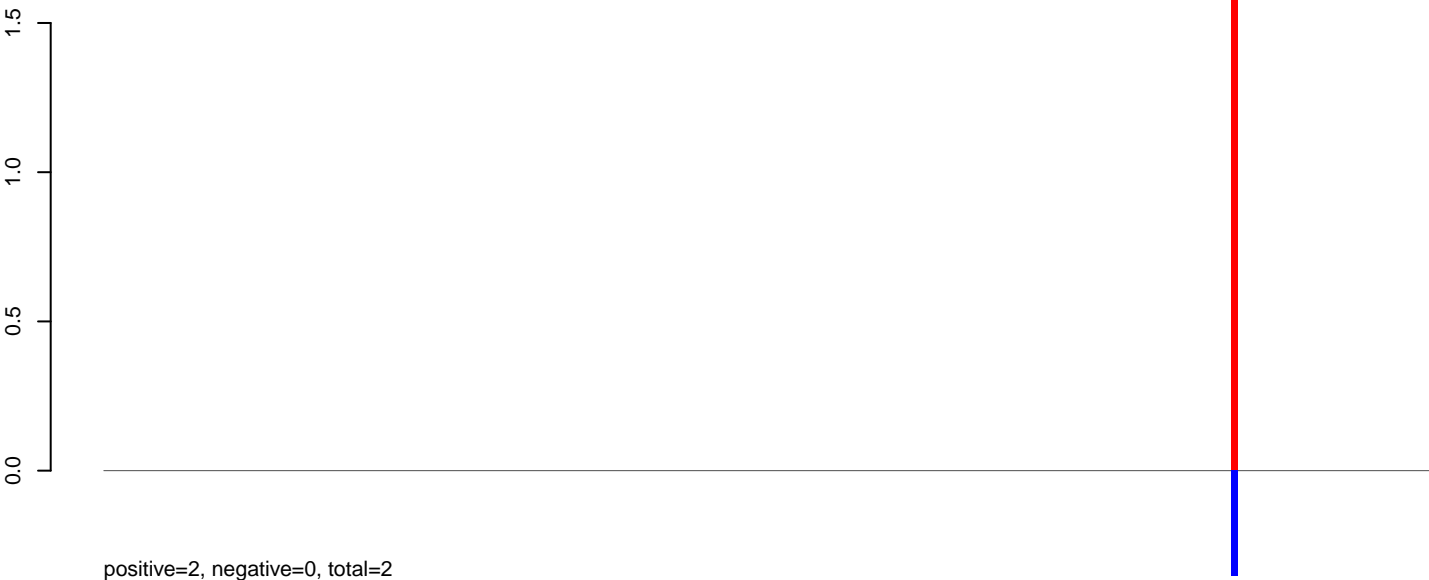
AeAeg_Aag2_DENV1.24_35.rep



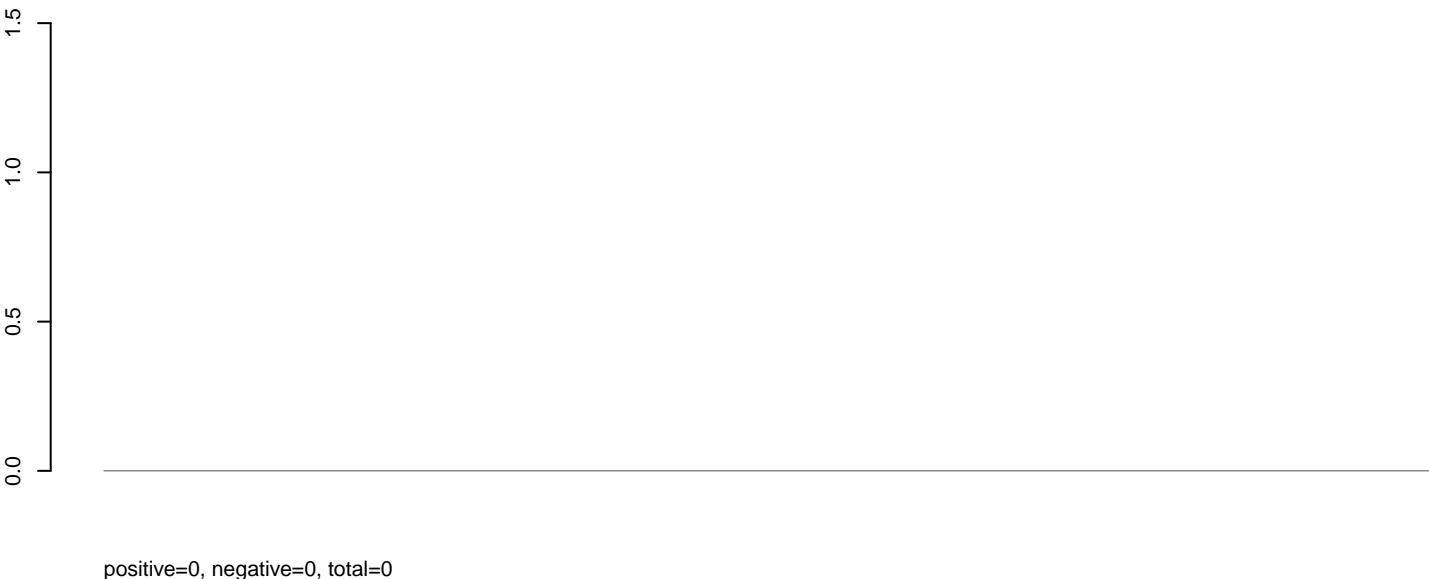
AeAeg_Aag2_DENV1.rep



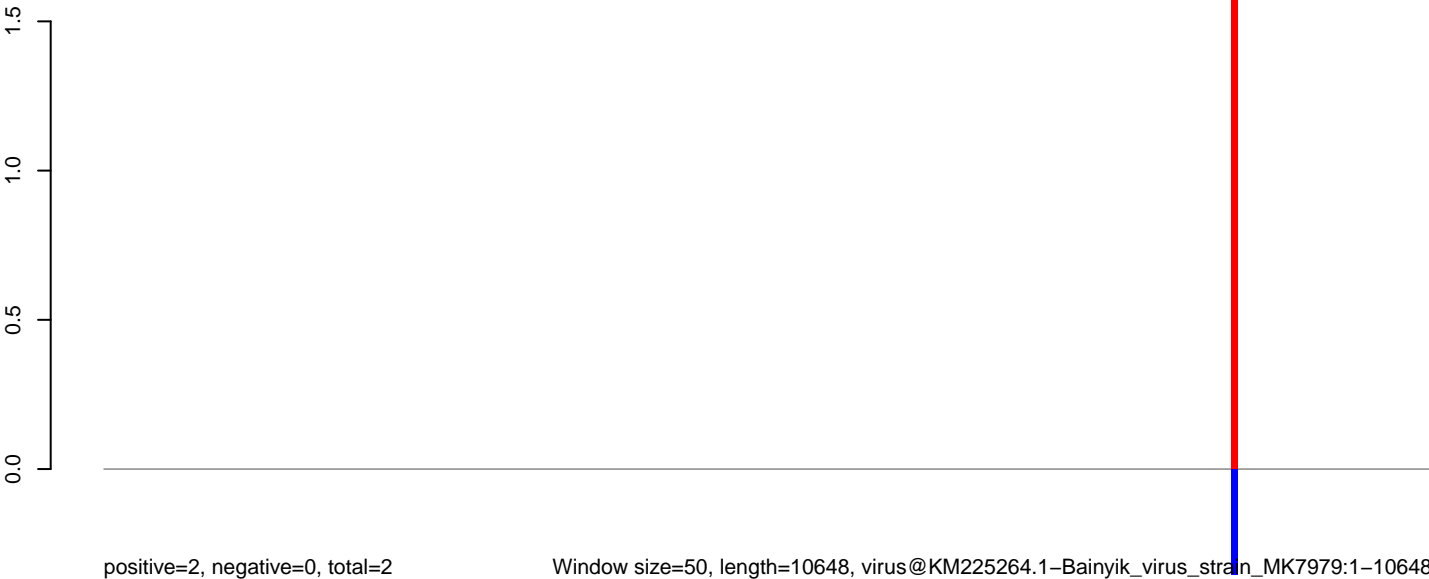
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



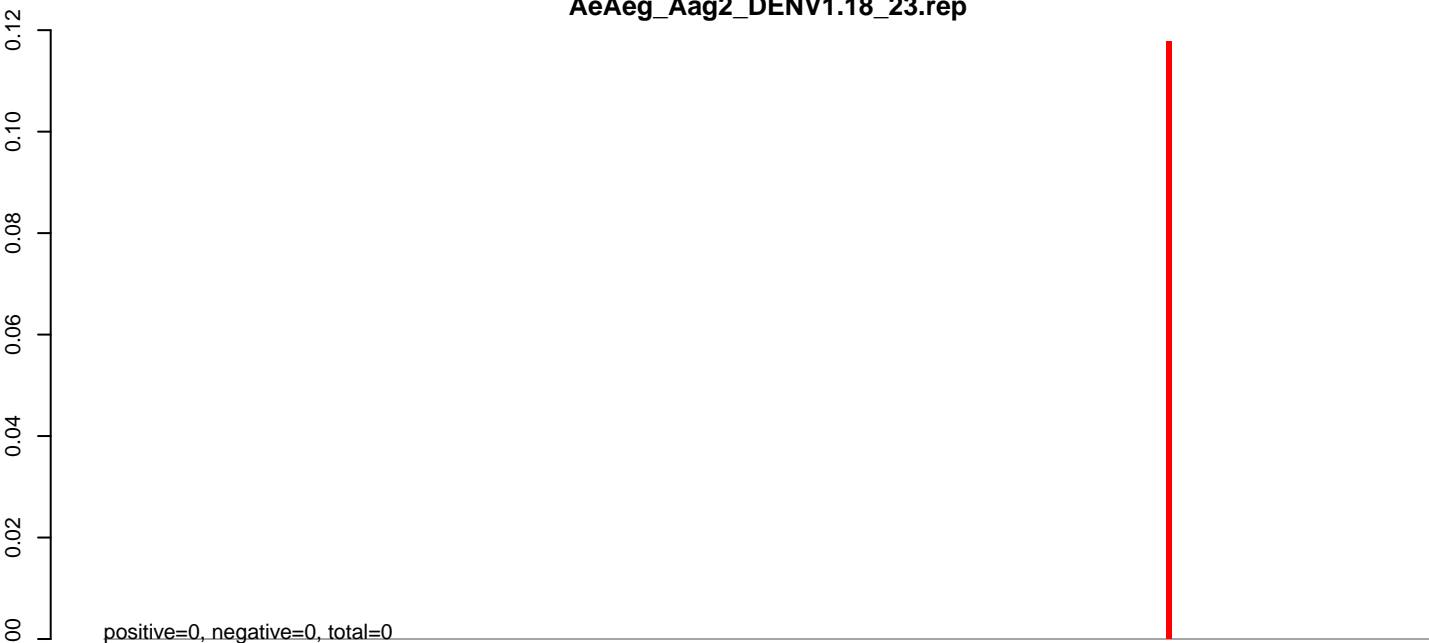
AeAeg_Aag2_DENV1.rep



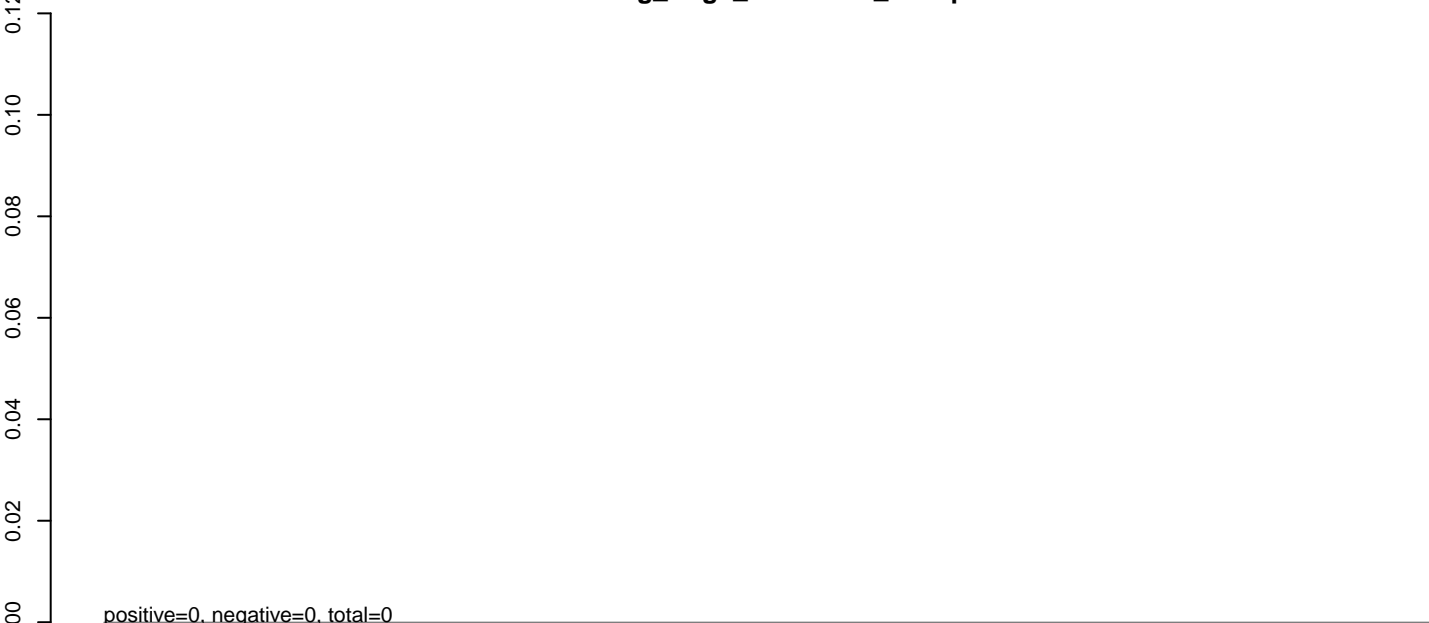
Window size=50, length=10648, virus@KM225264.1-Bainyik_virus_strain_MK7979:1-10648

0 2000 4000 6000 8000 10000

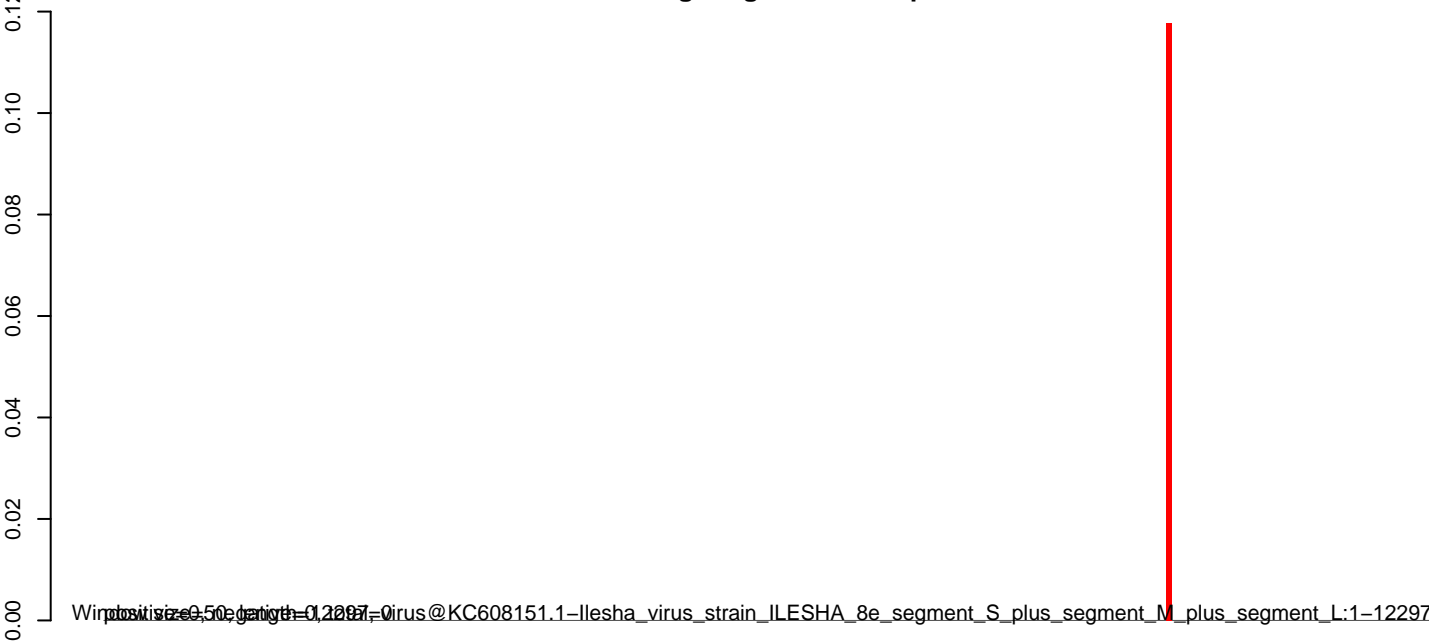
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



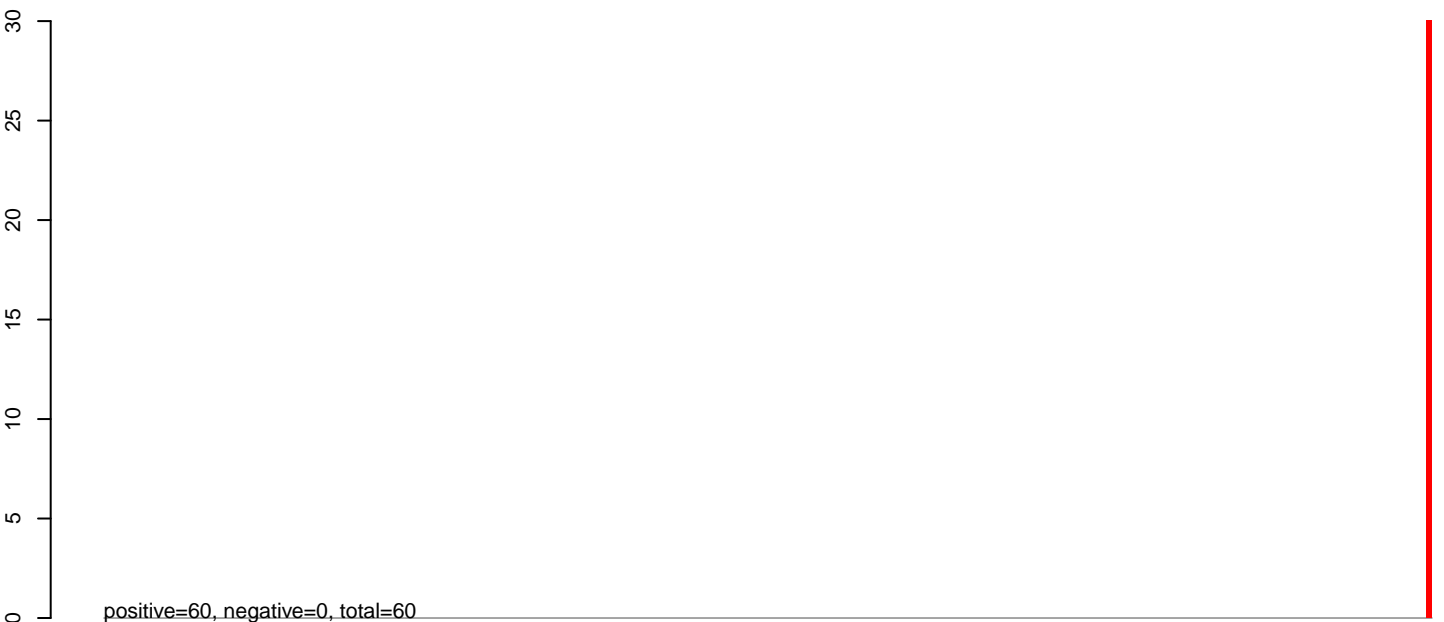
AeAeg_Aag2_DENV1.rep



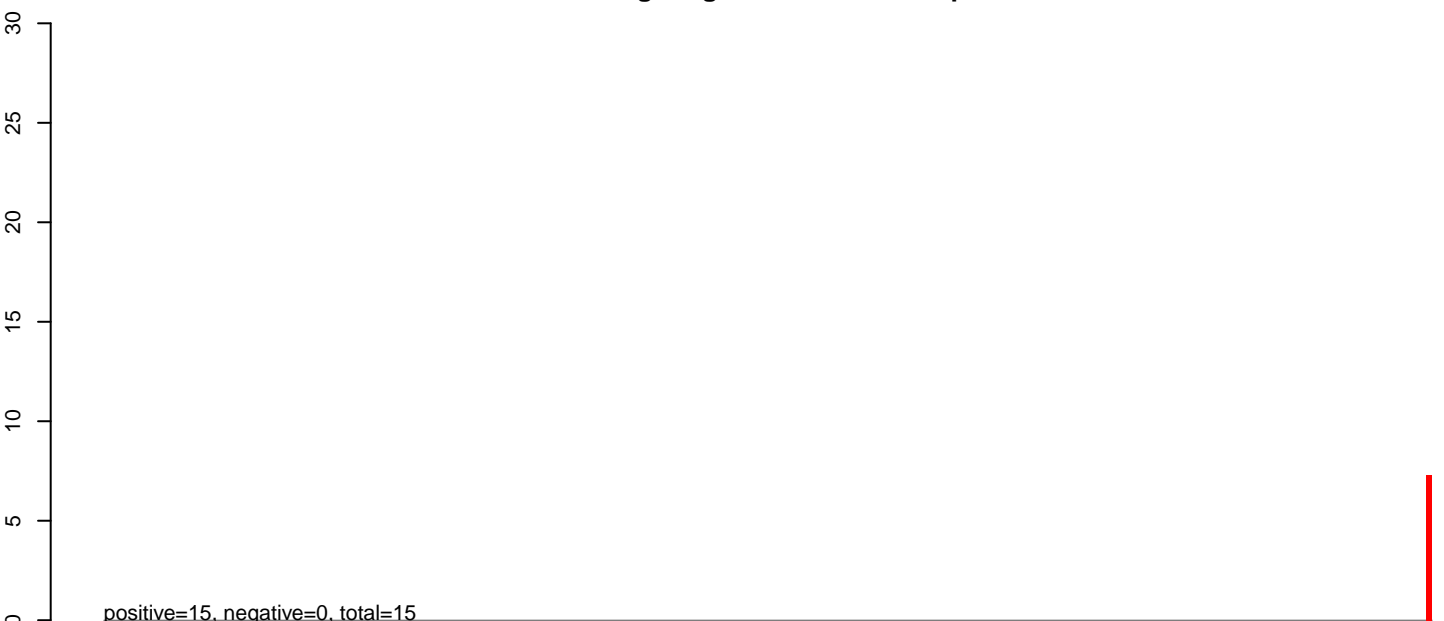
W:\msi\size_50\gaugh=0_2021-01-29-01\virus@KC608151.1-Ilesha_virus_strain_ILESHA_8e_segment_S_plus_segment_M_plus_segment_L:1-12297

0 2000 4000 6000 8000 10000 12000

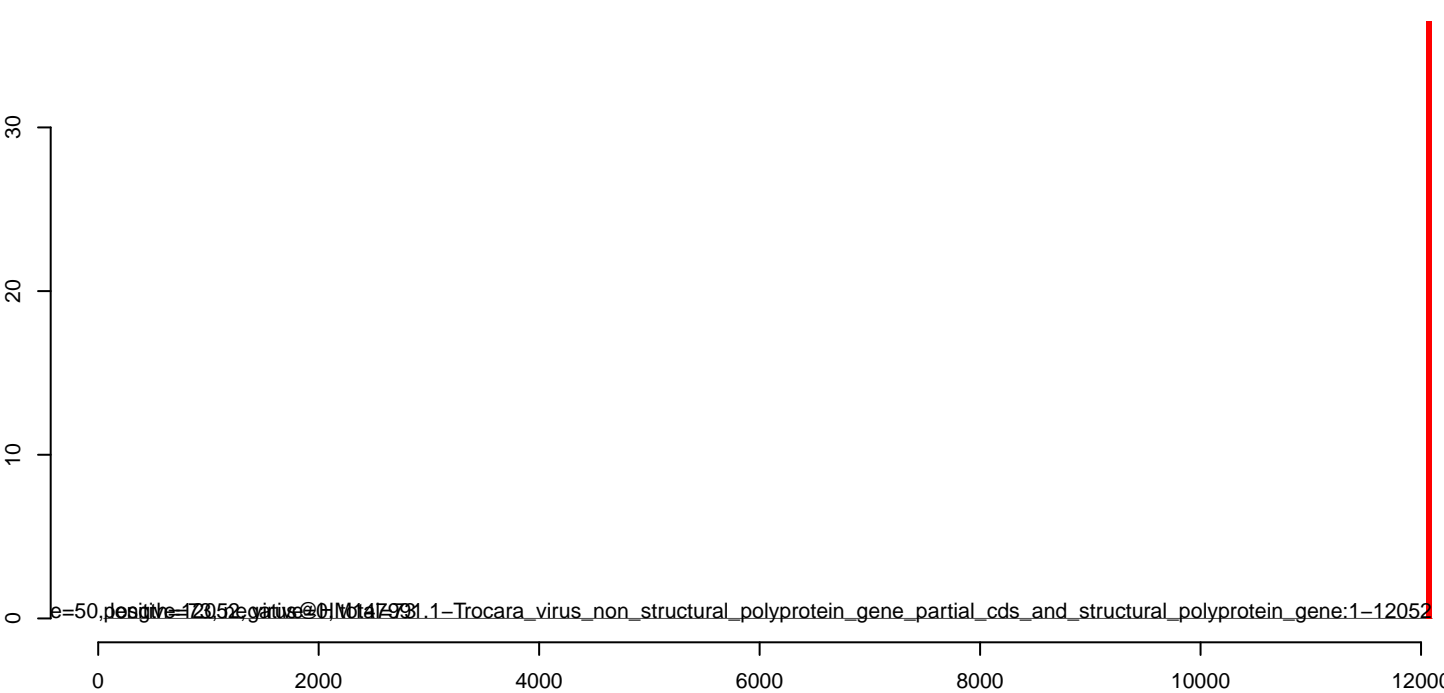
AeAeg_Aag2_DENV1.18_23.rep



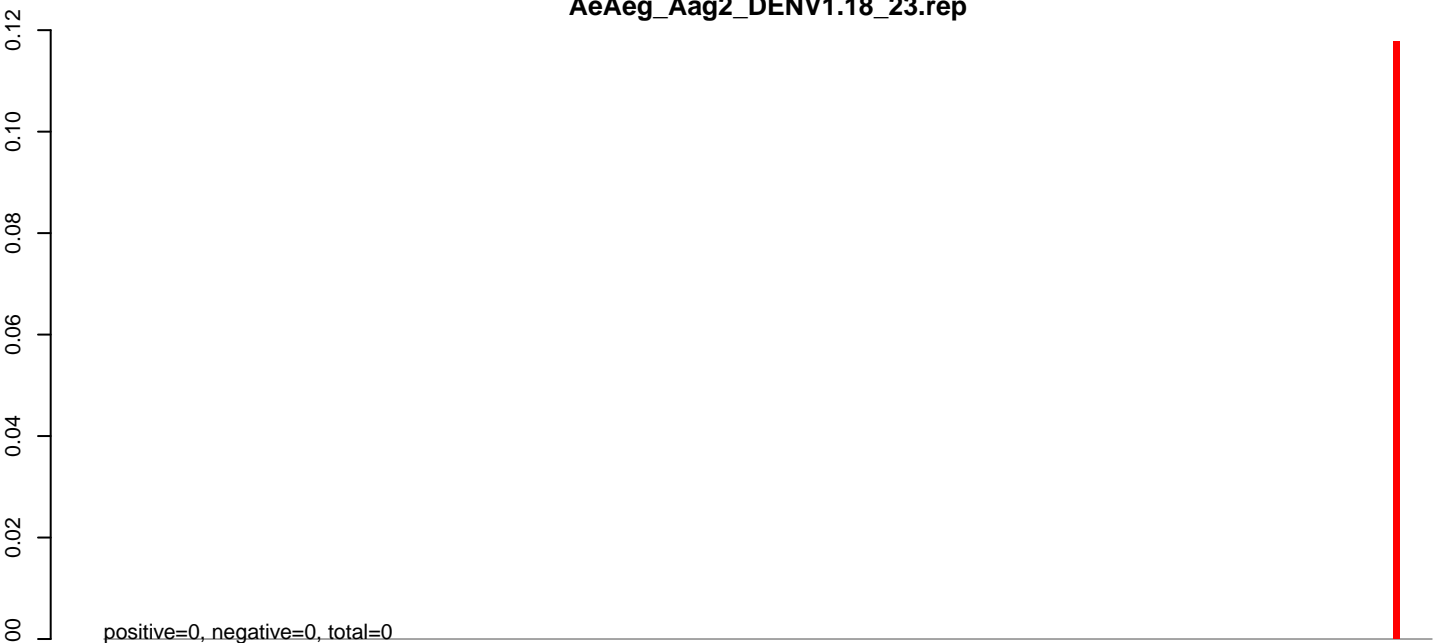
AeAeg_Aag2_DENV1.24_35.rep



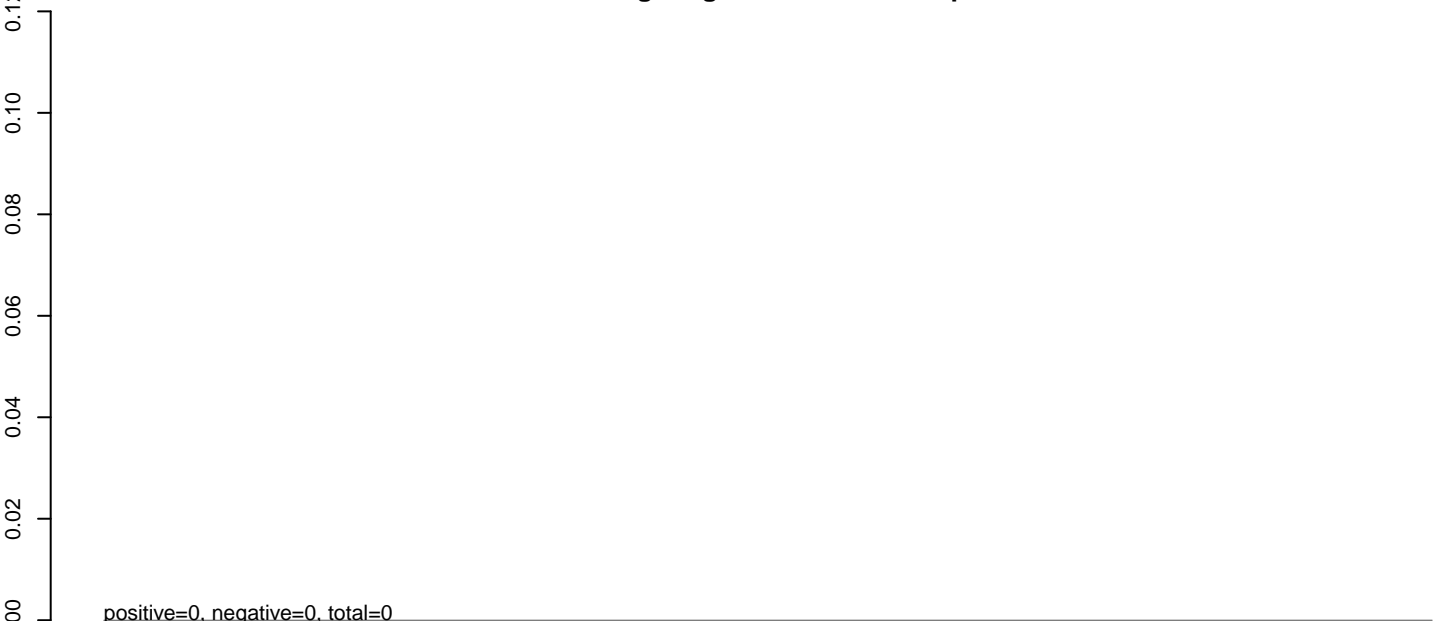
AeAeg_Aag2_DENV1.rep



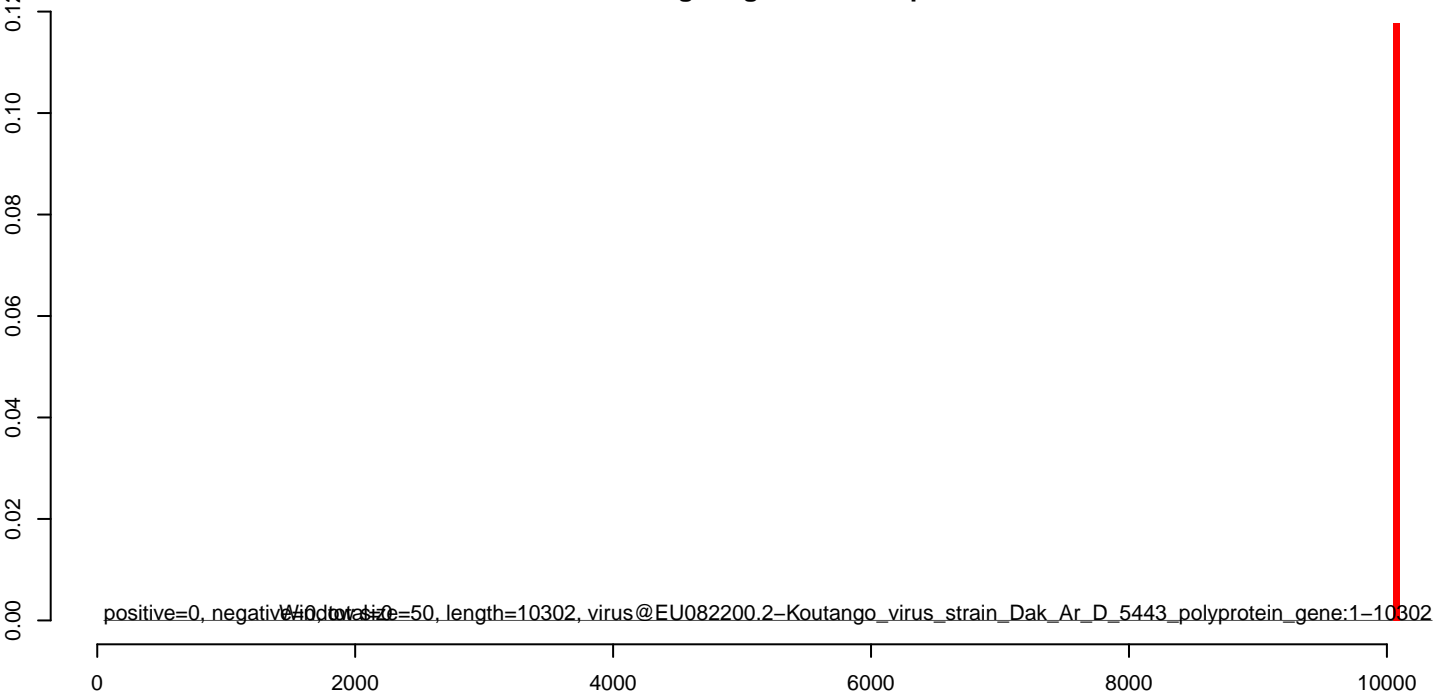
AeAeg_Aag2_DENV1.18_23.rep



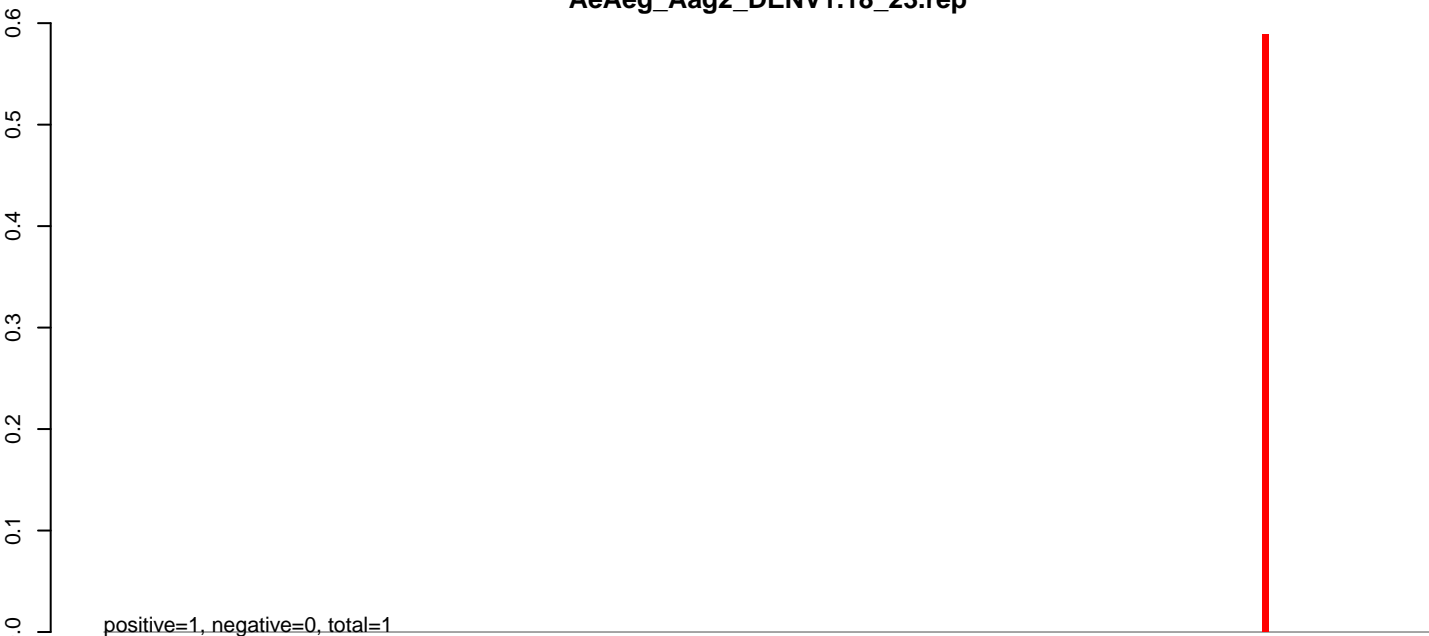
AeAeg_Aag2_DENV1.24_35.rep



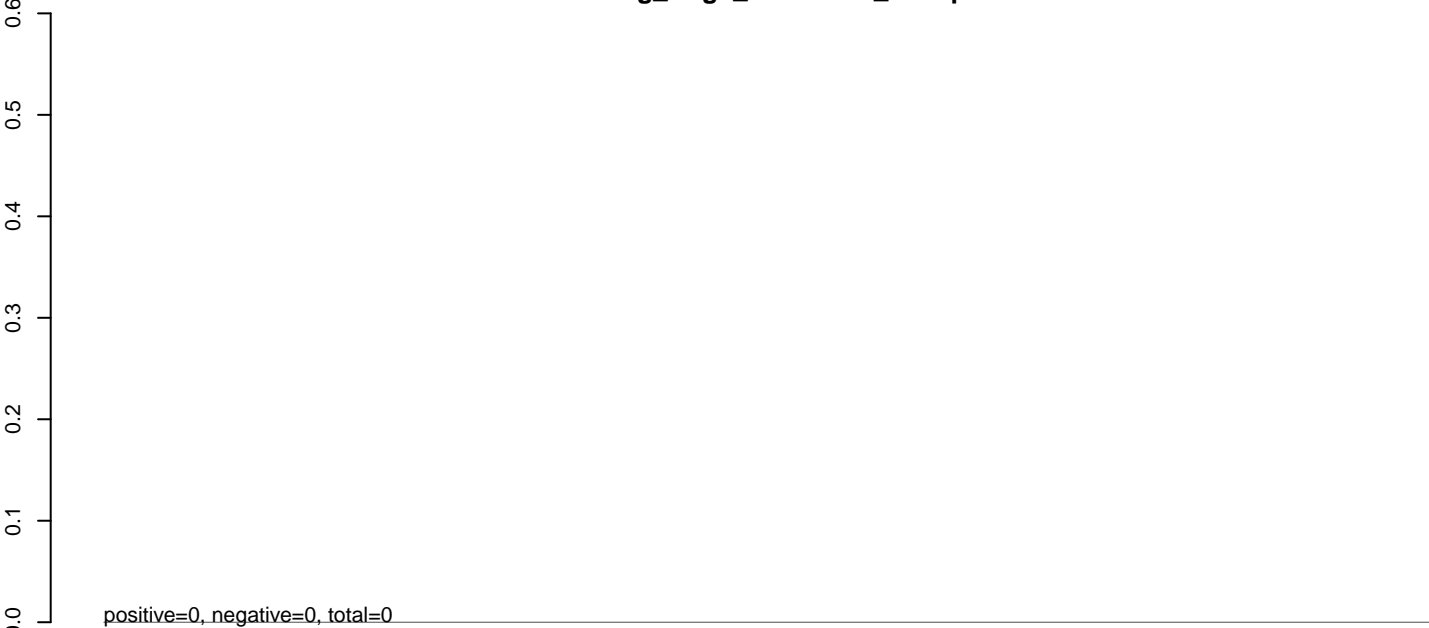
AeAeg_Aag2_DENV1.rep



AeAeg_Aag2_DENV1.18_23.rep



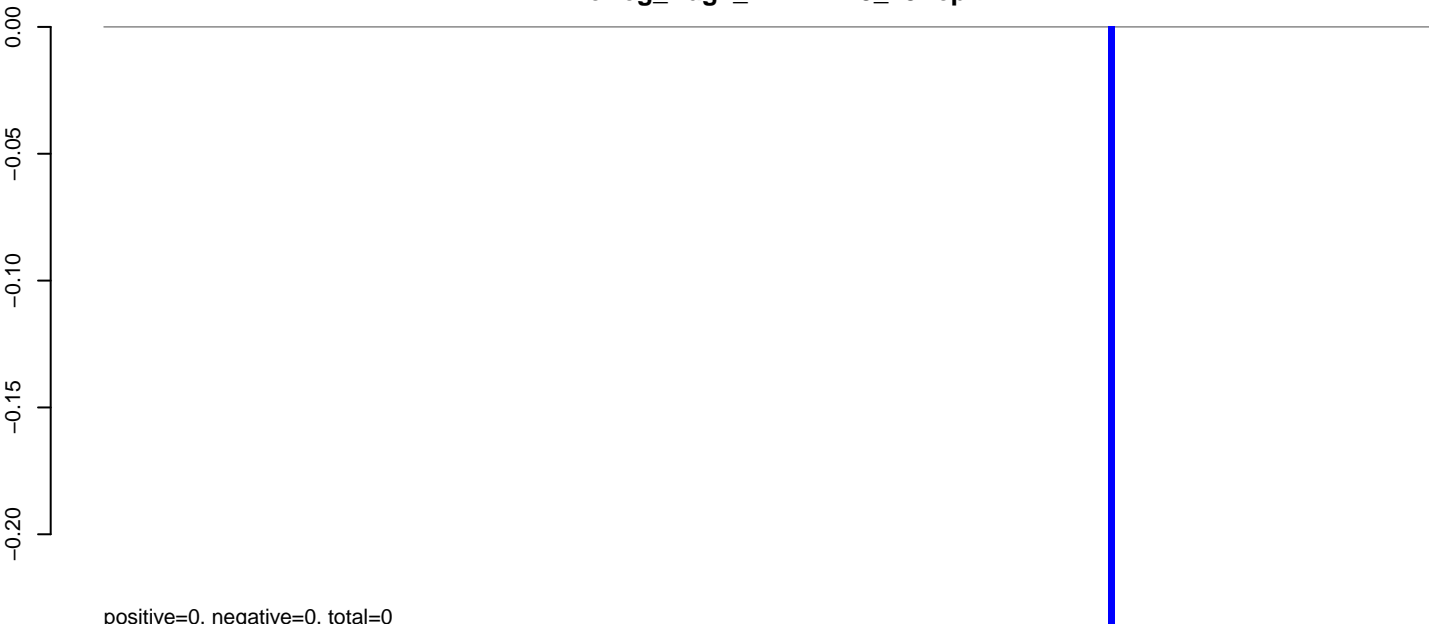
AeAeg_Aag2_DENV1.24_35.rep



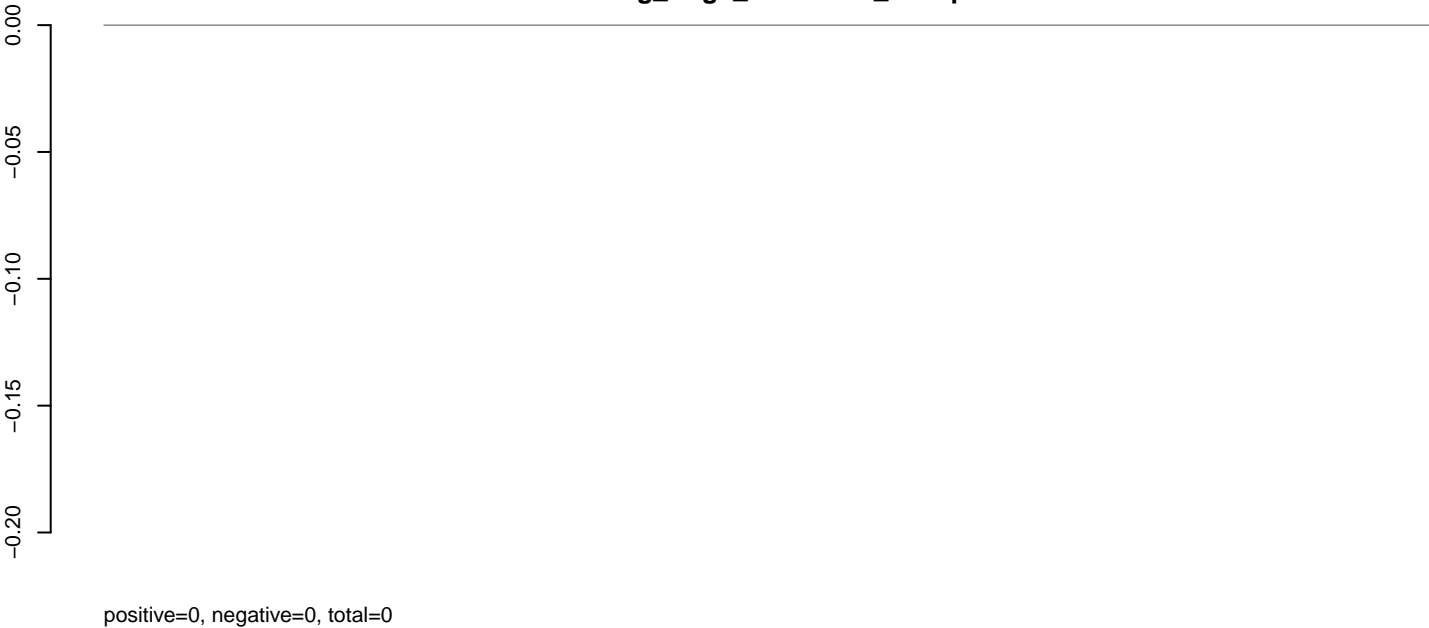
AeAeg_Aag2_DENV1.rep



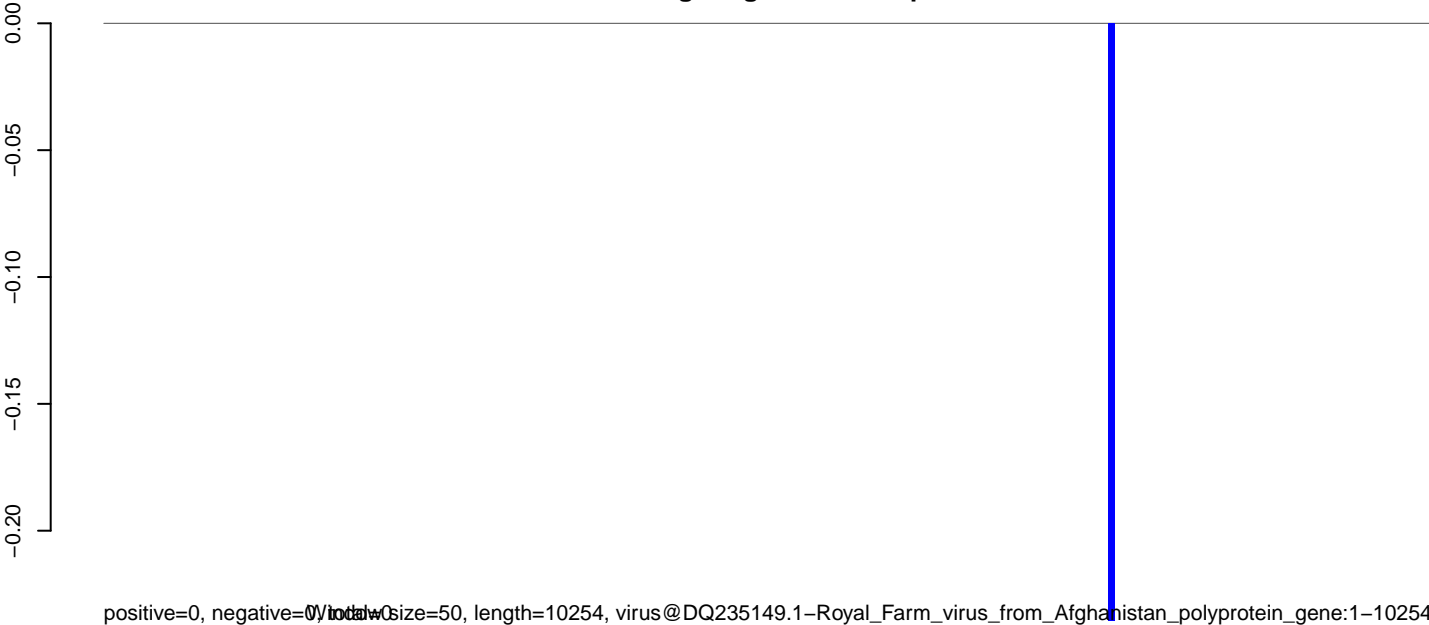
AeAeg_Aag2_DENV1.18_23.rep



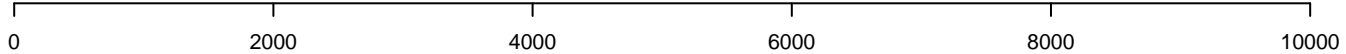
AeAeg_Aag2_DENV1.24_35.rep



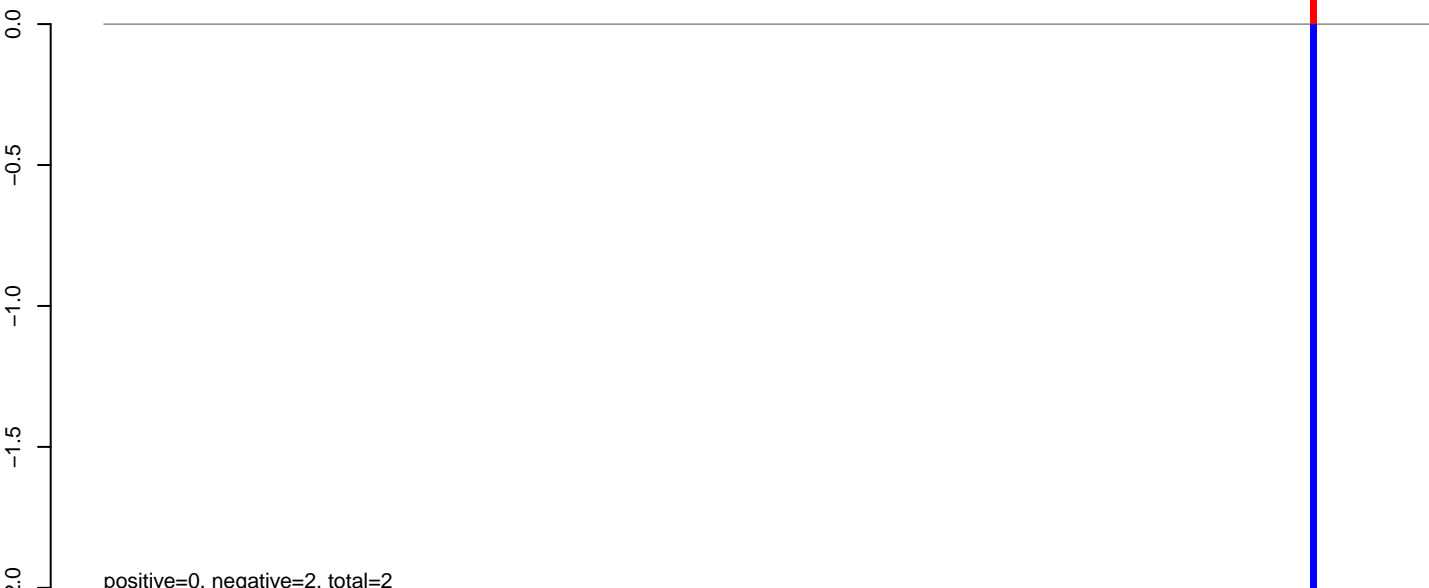
AeAeg_Aag2_DENV1.rep



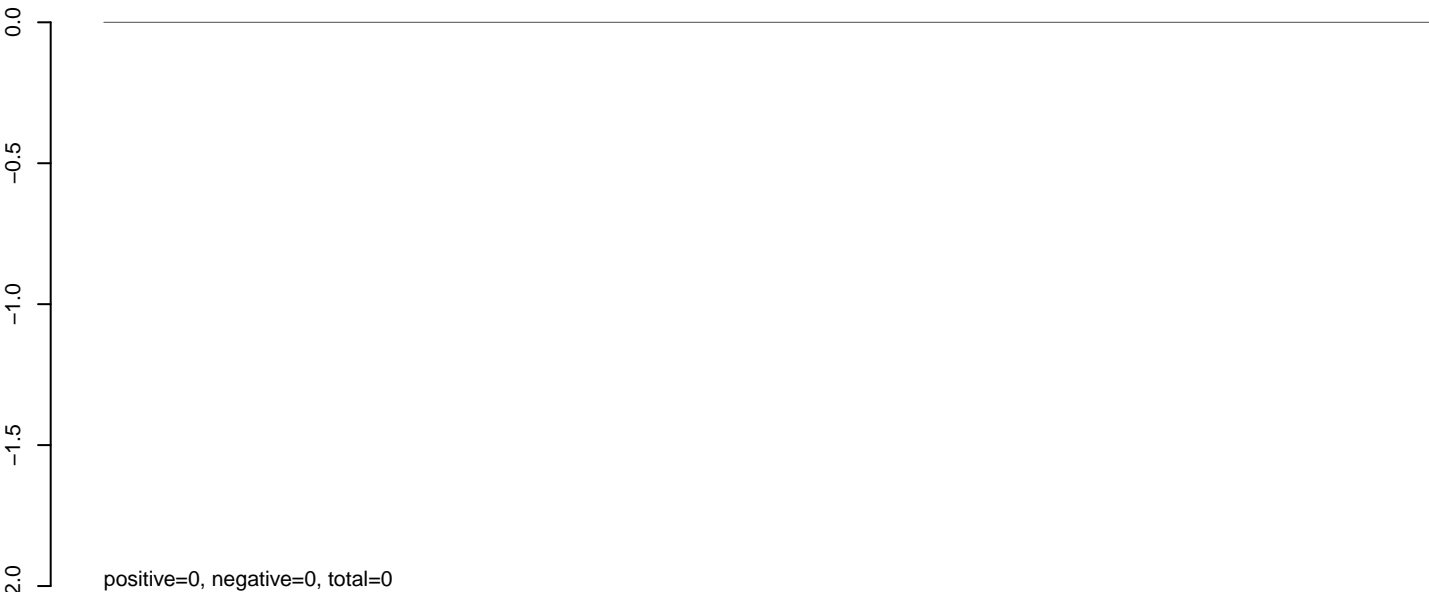
Window size=50, length=10254, virus@DQ235149.1-Royal_Farm_virus_from_Afghanistan_polyprotein_gene:1-10254



AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



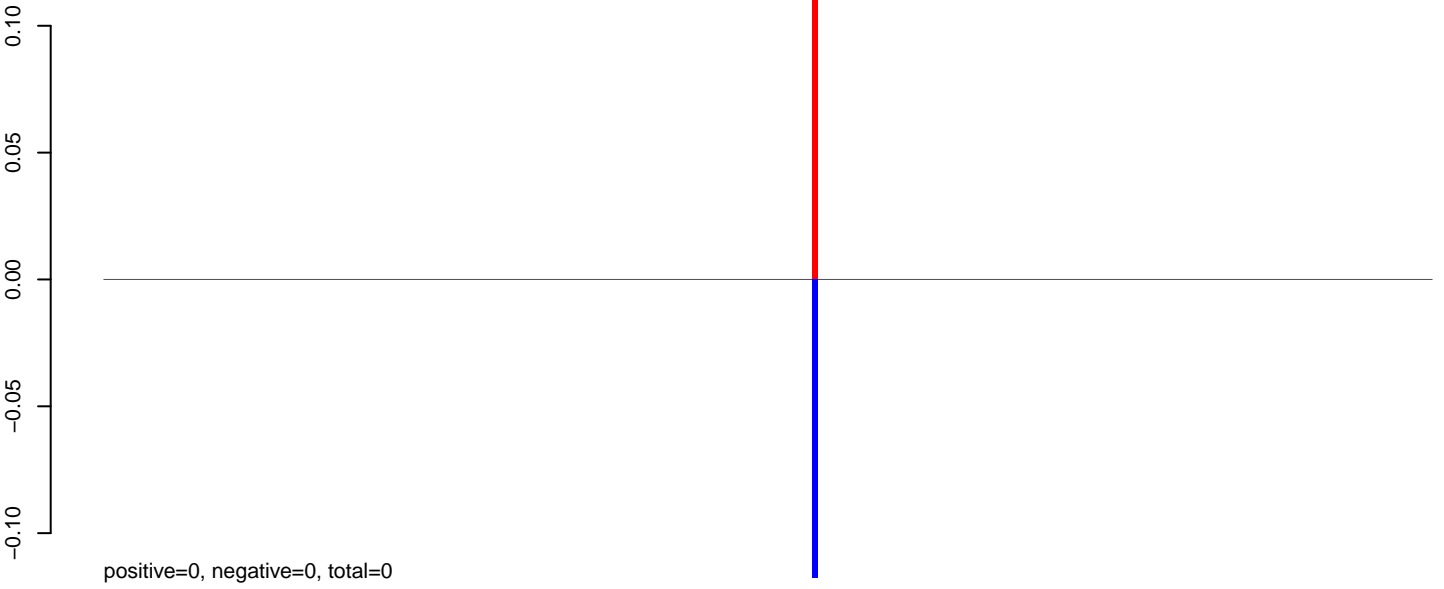
AeAeg_Aag2_DENV1.rep



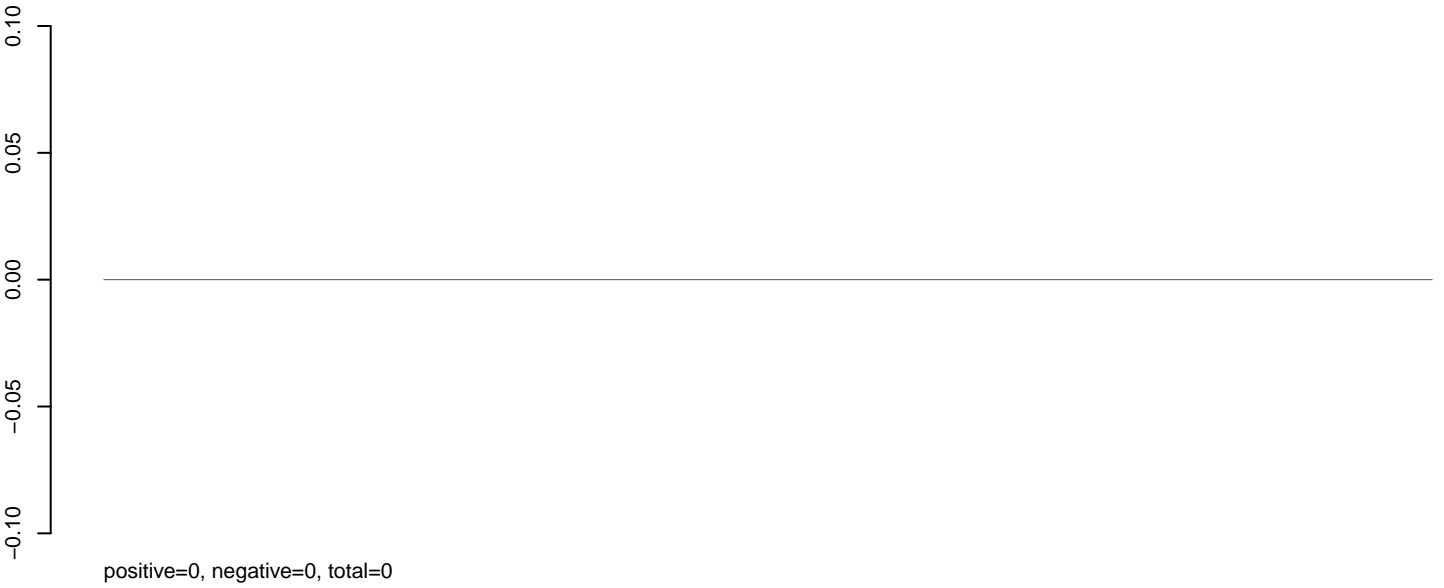
Window size=50, length=10922, virus@AY632538.4-Iguape_virus_strain_SPAn_71686:1-10922

0 2000 4000 6000 8000 10000

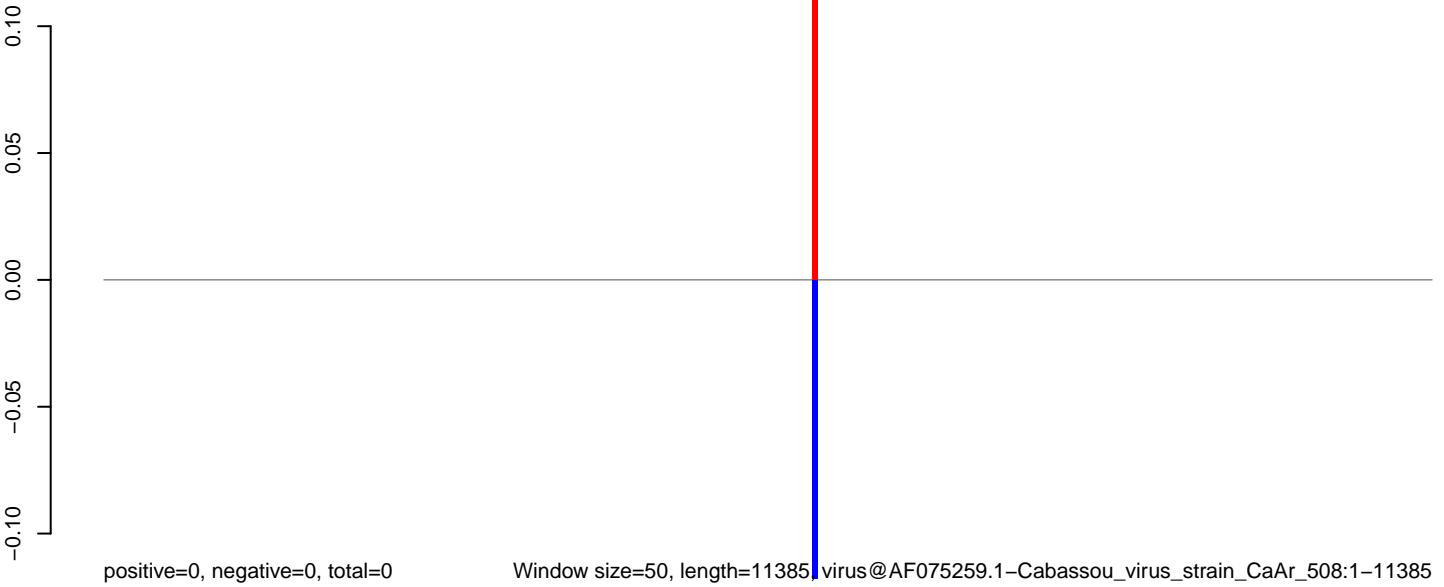
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



AeAeg_Aag2_DENV1.rep



0 2000 4000 6000 8000 10000

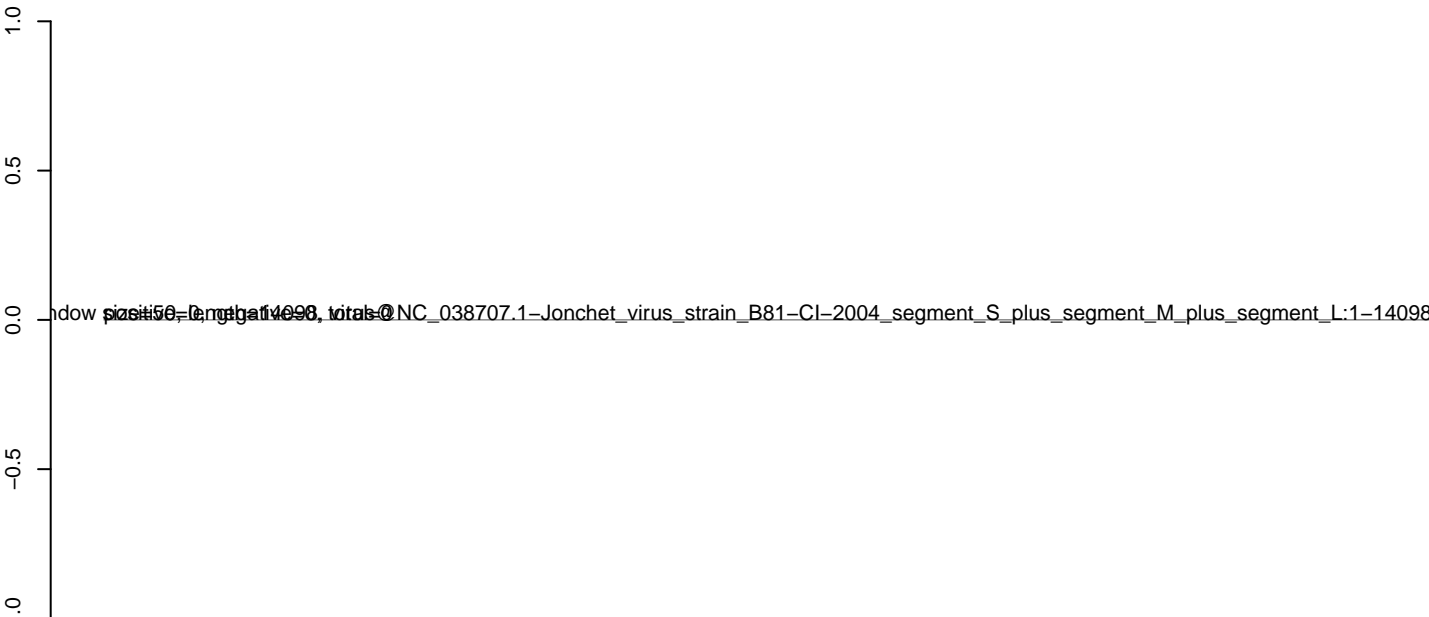
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep

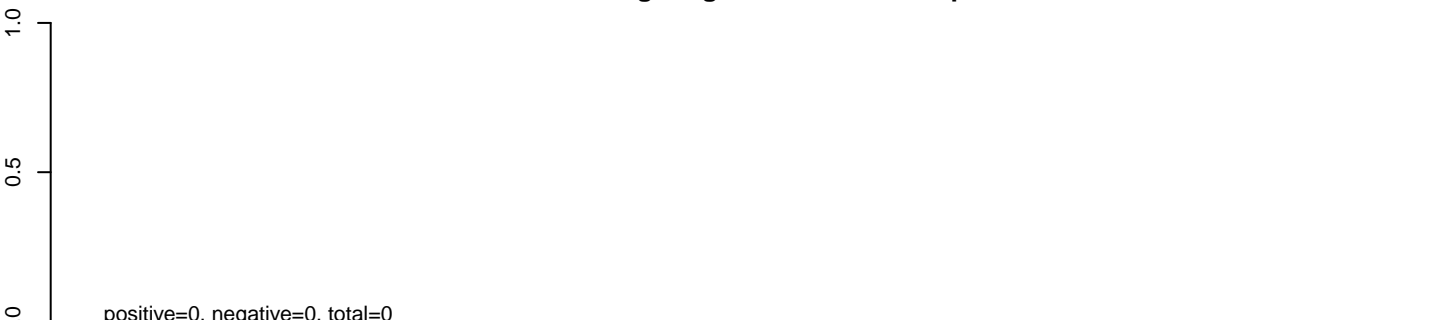


AeAeg_Aag2_DENV1.rep

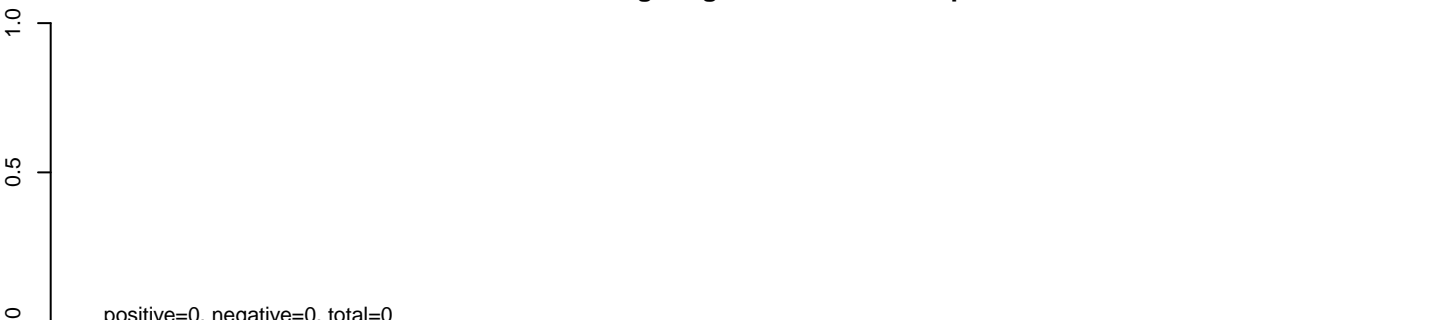


0 2000 4000 6000 8000 10000 12000 14000

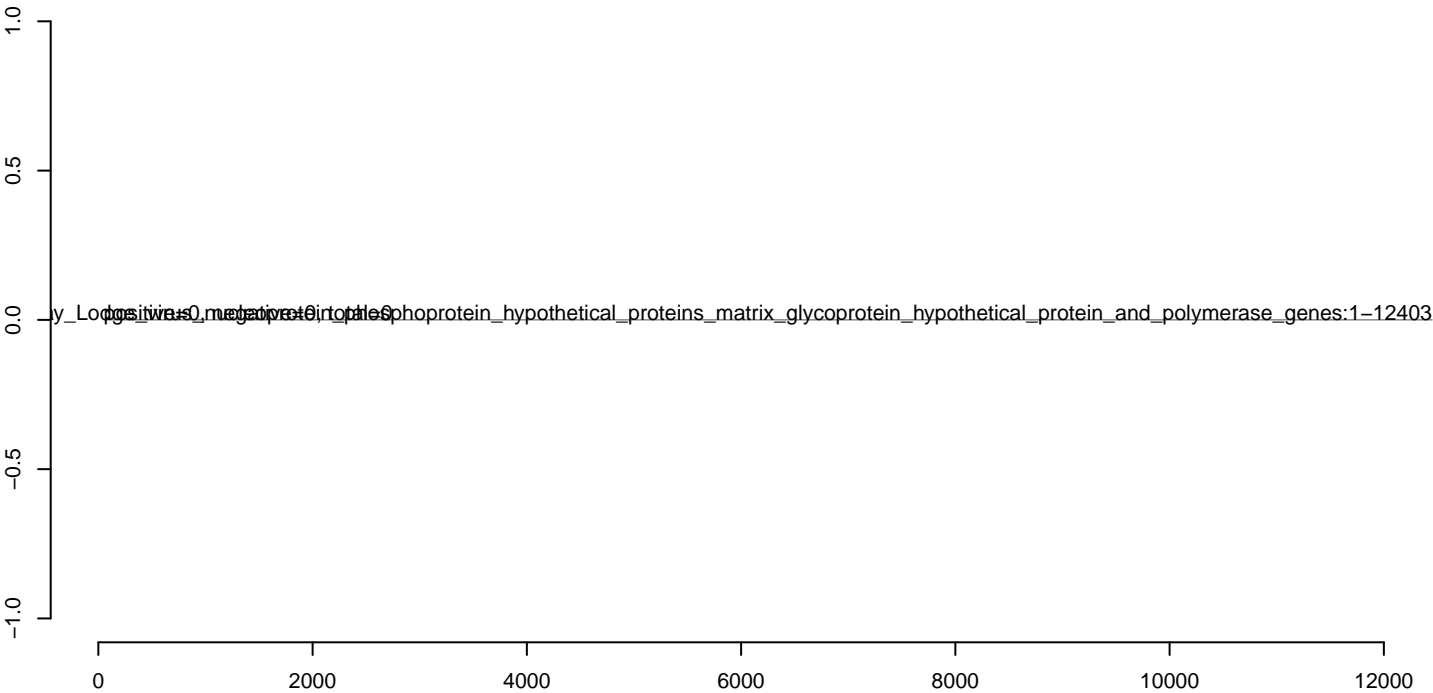
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



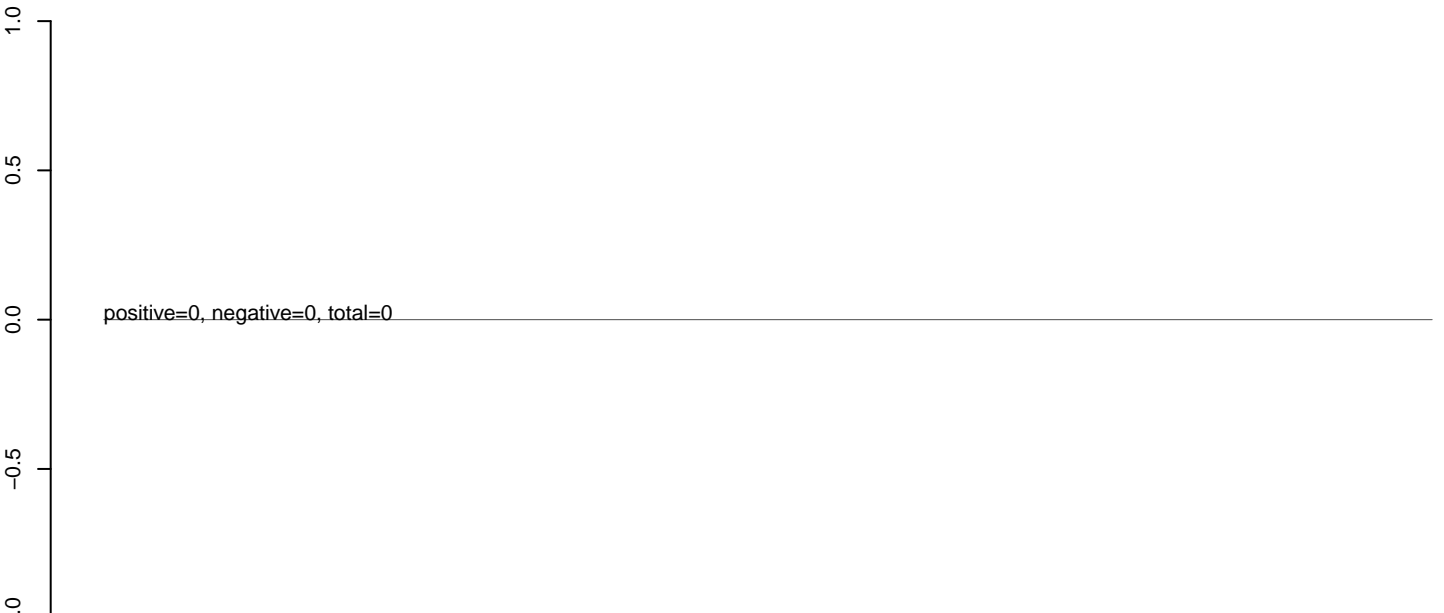
AeAeg_Aag2_DENV1.rep



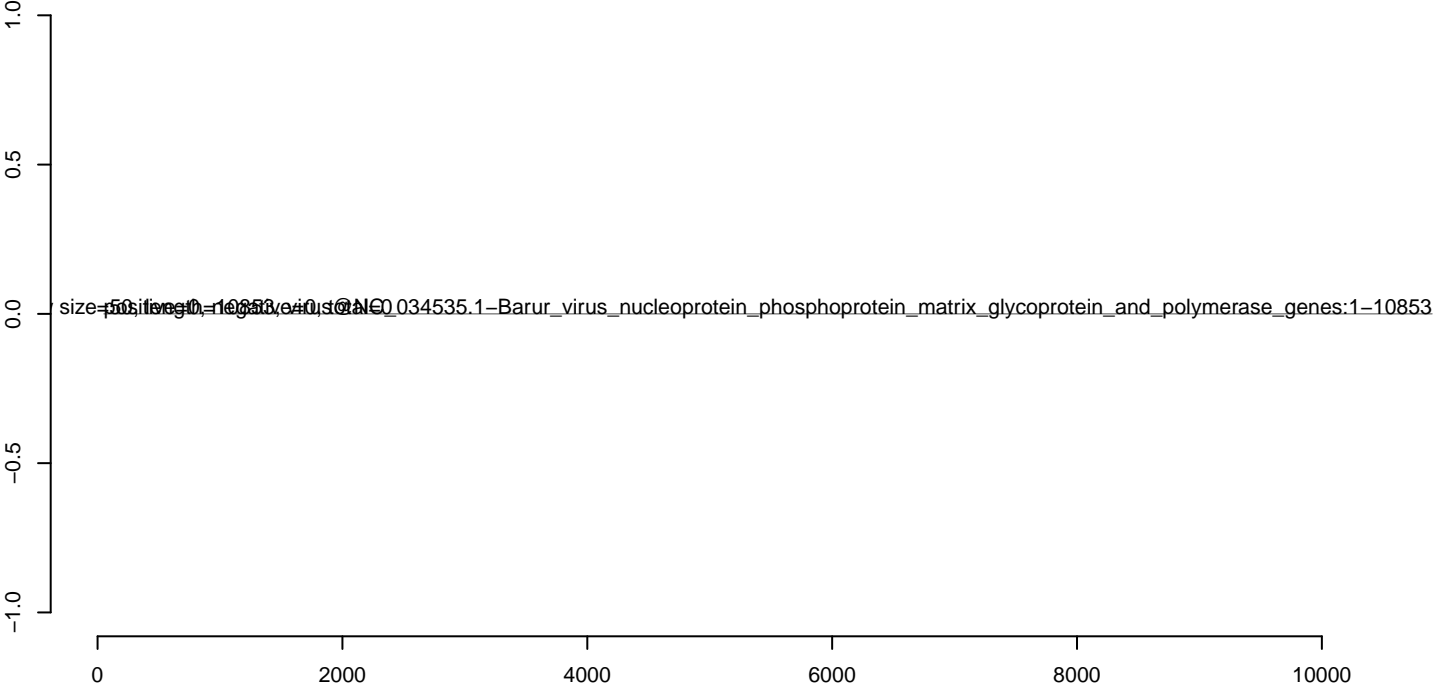
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



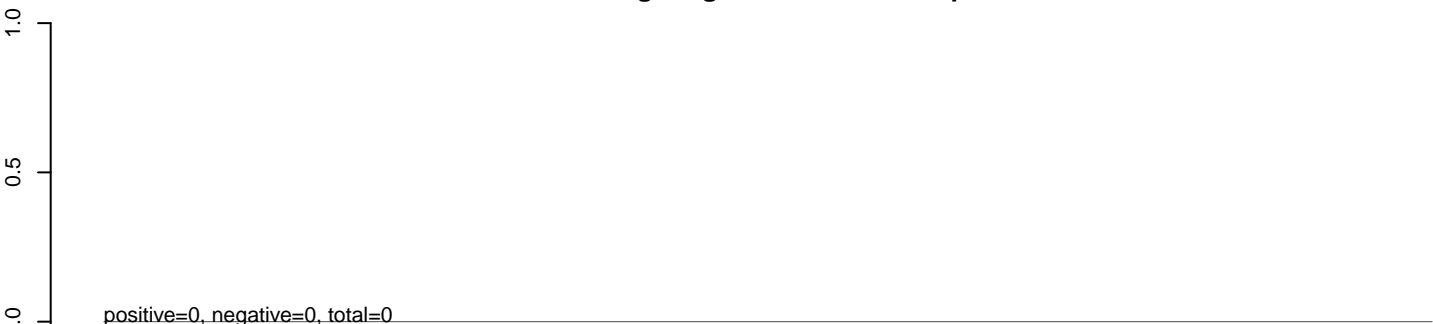
AeAeg_Aag2_DENV1.rep



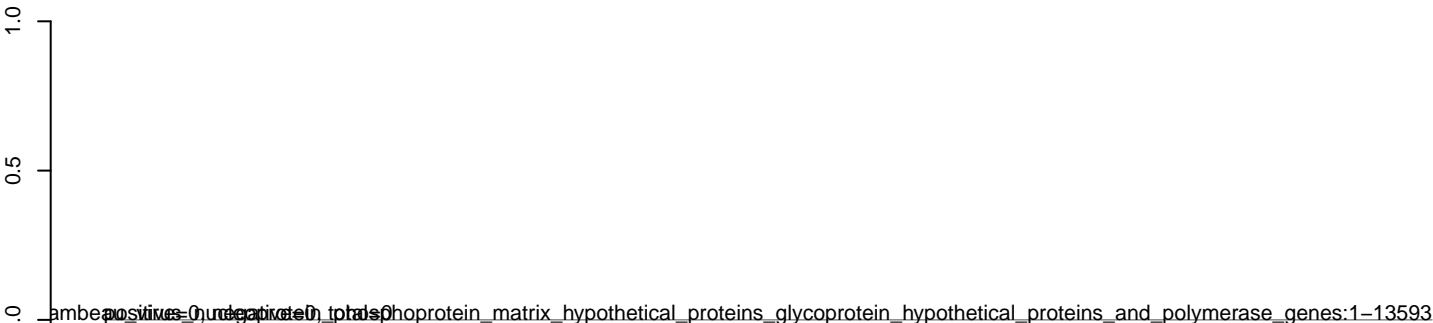
AeAeg_Aag2_DENV1.18_23.rep



AeAeg_Aag2_DENV1.24_35.rep



AeAeg_Aag2_DENV1.rep



0 2000 4000 6000 8000 10000 12000 14000

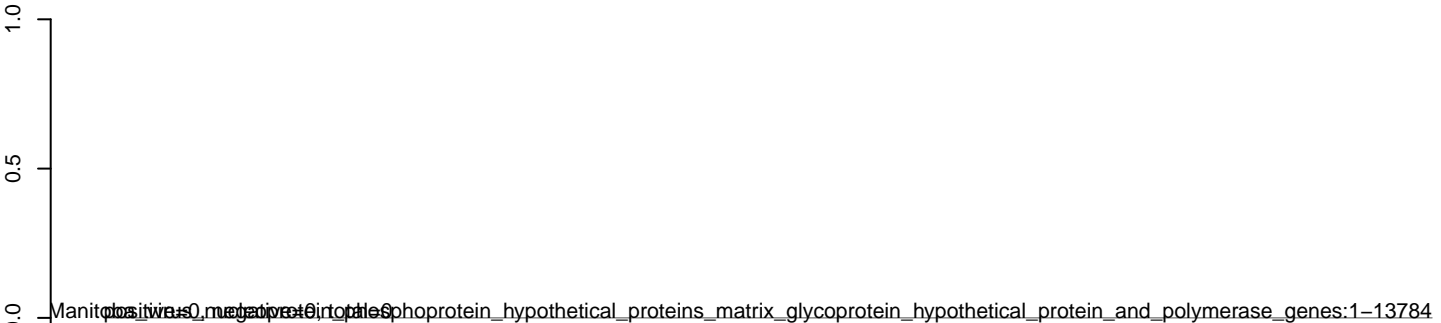
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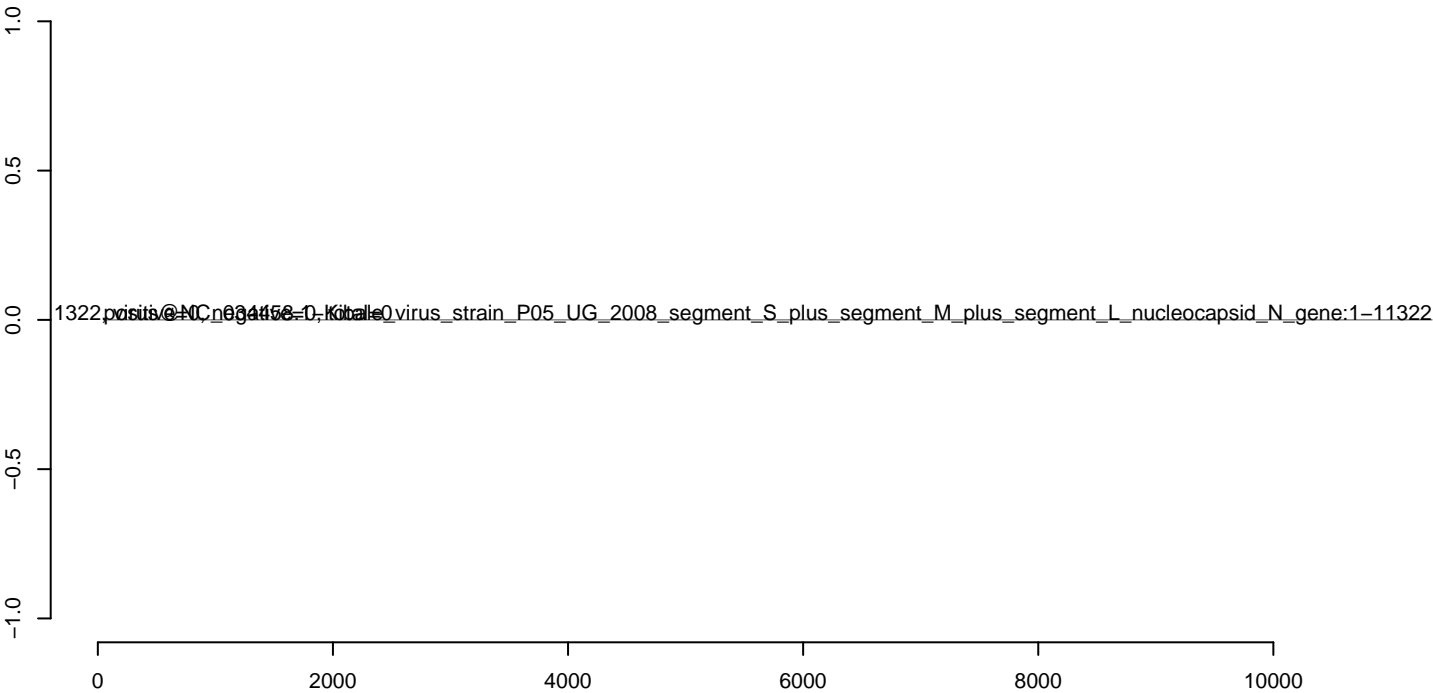
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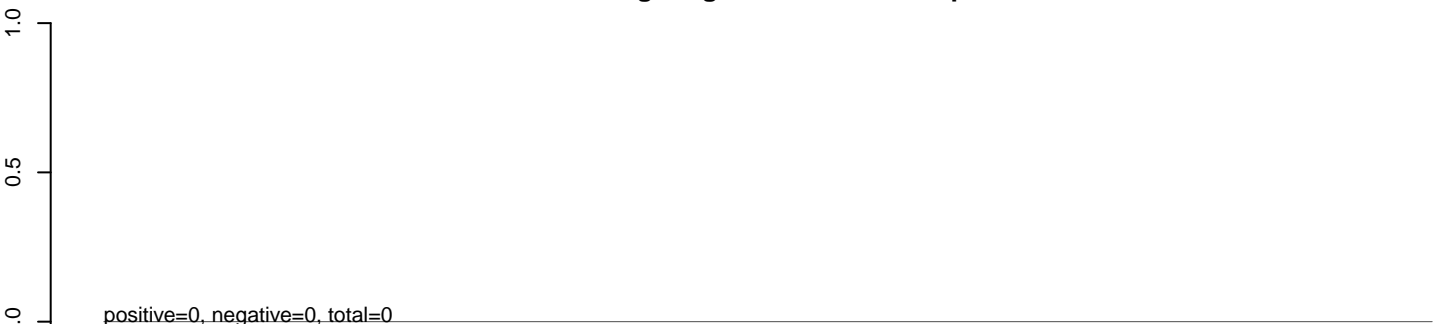
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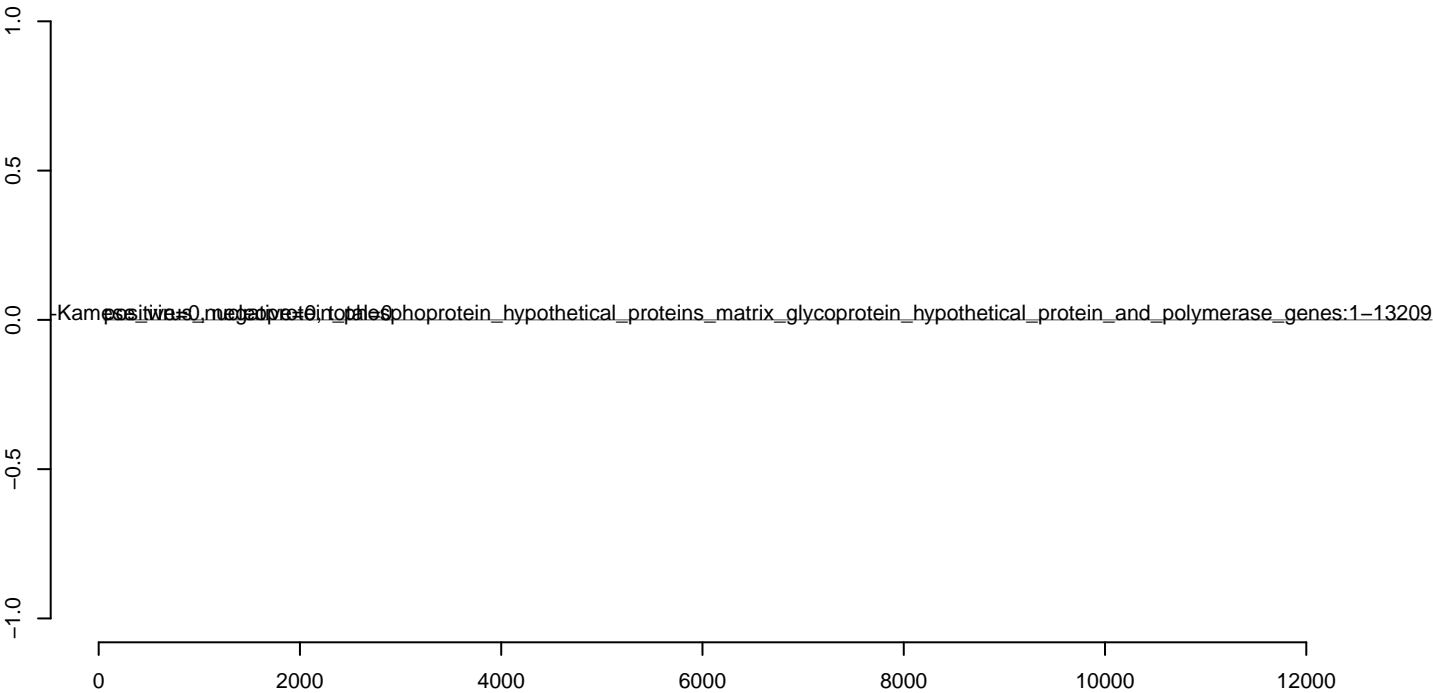
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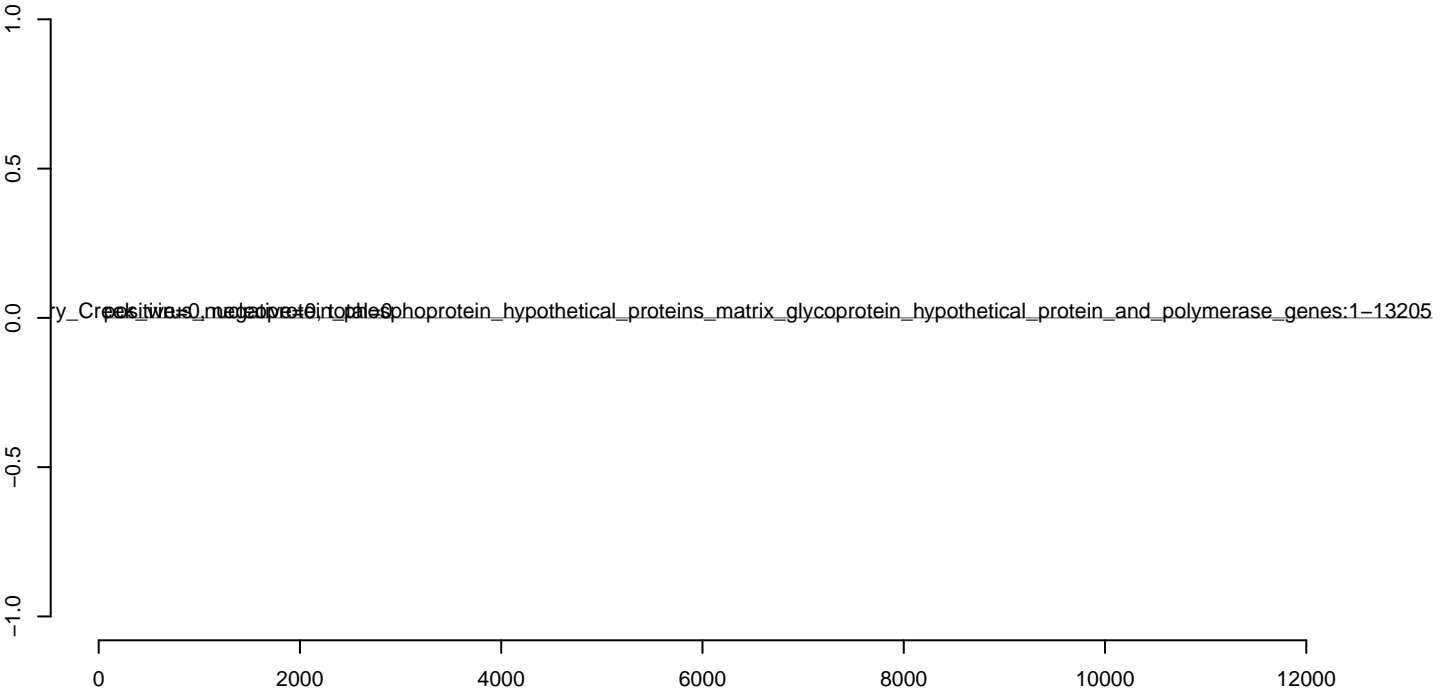
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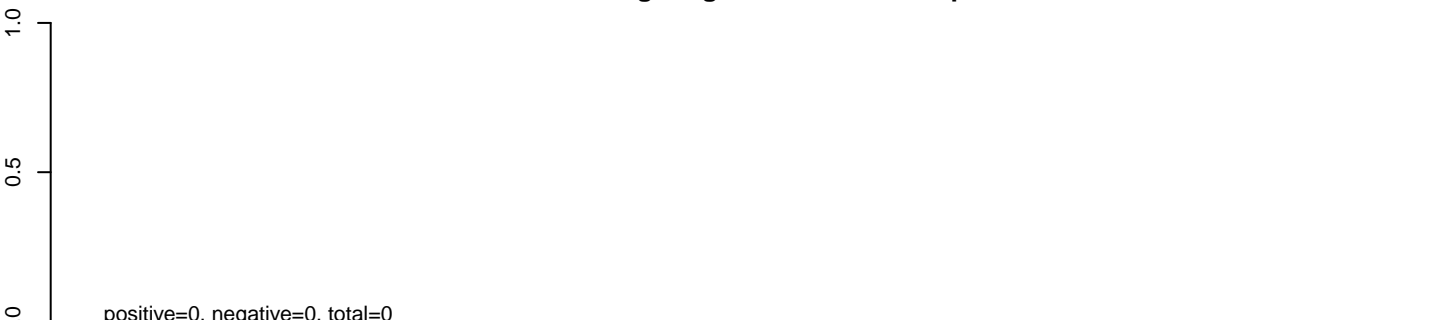
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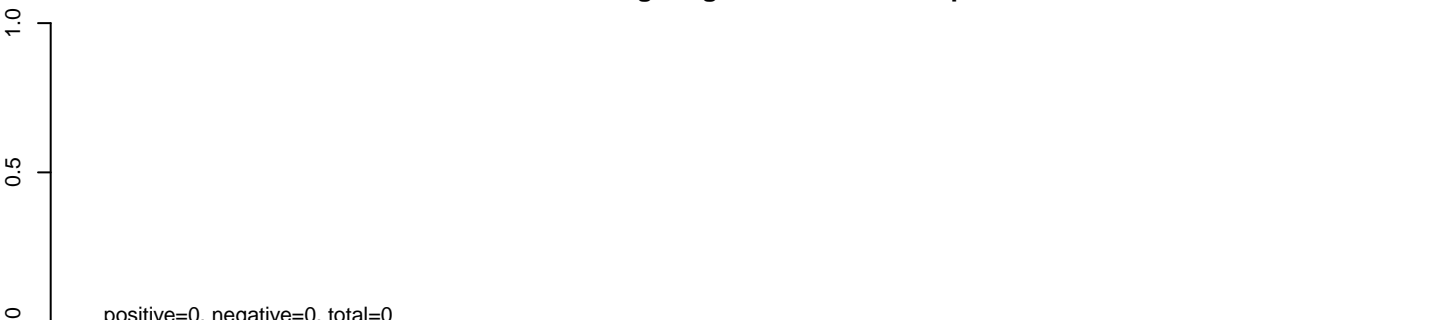
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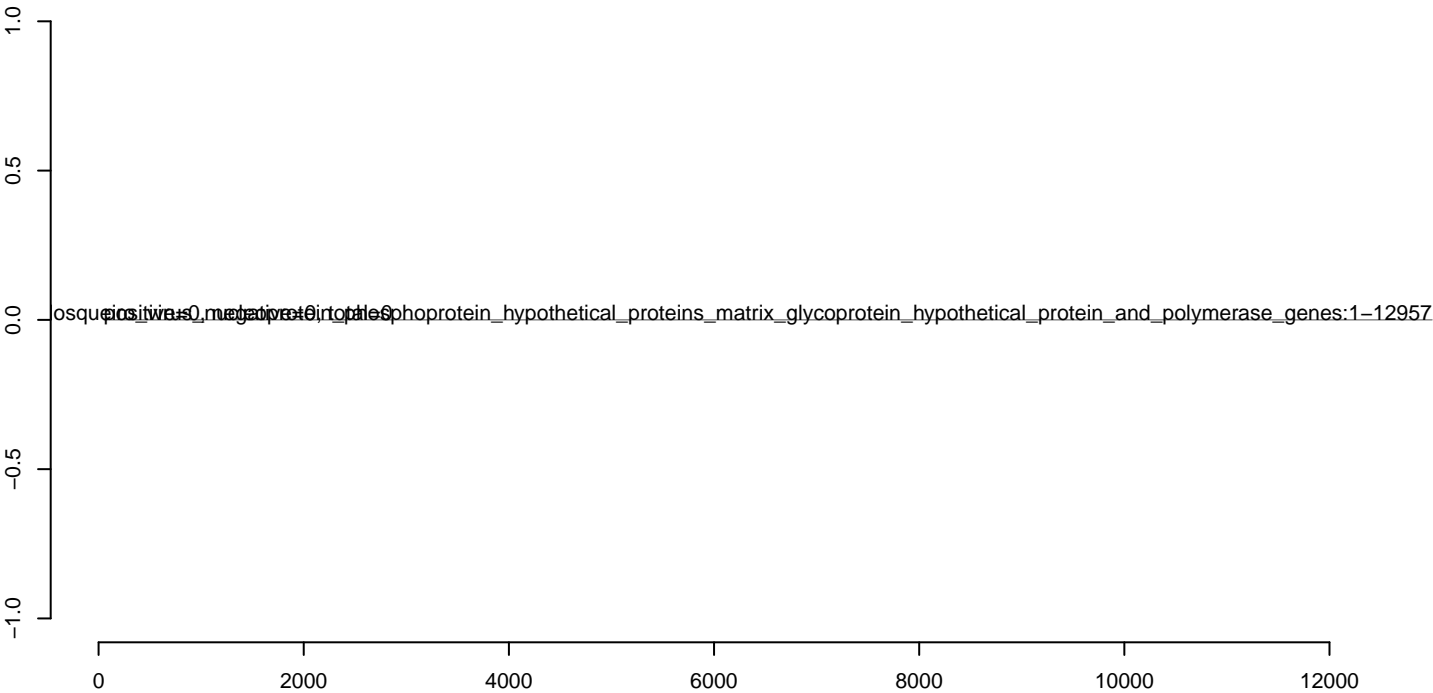
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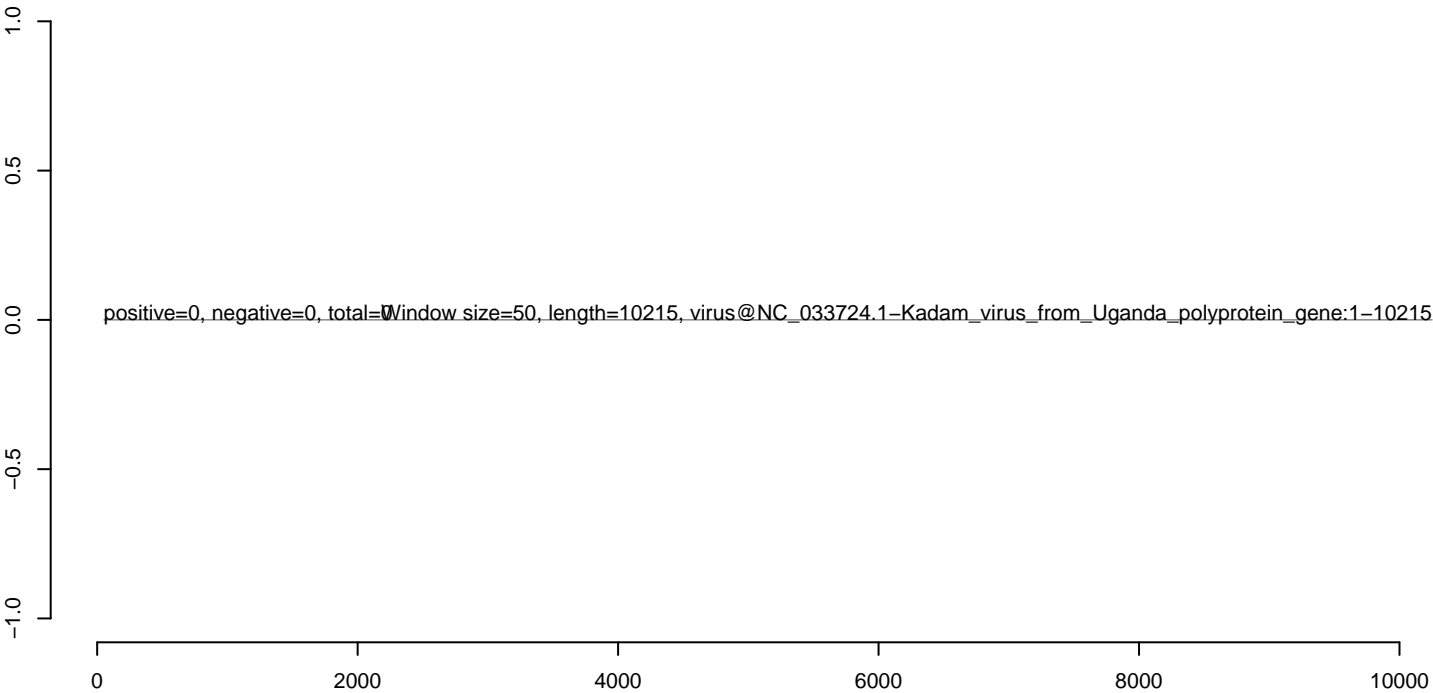
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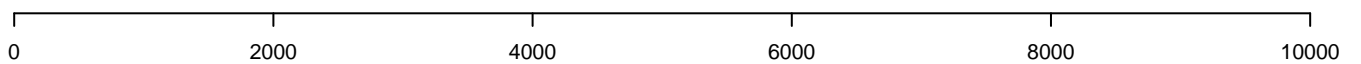
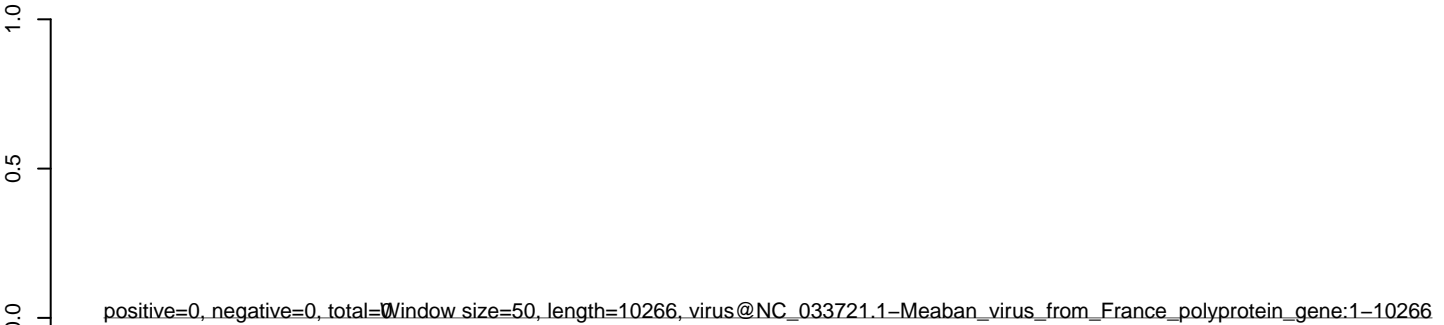
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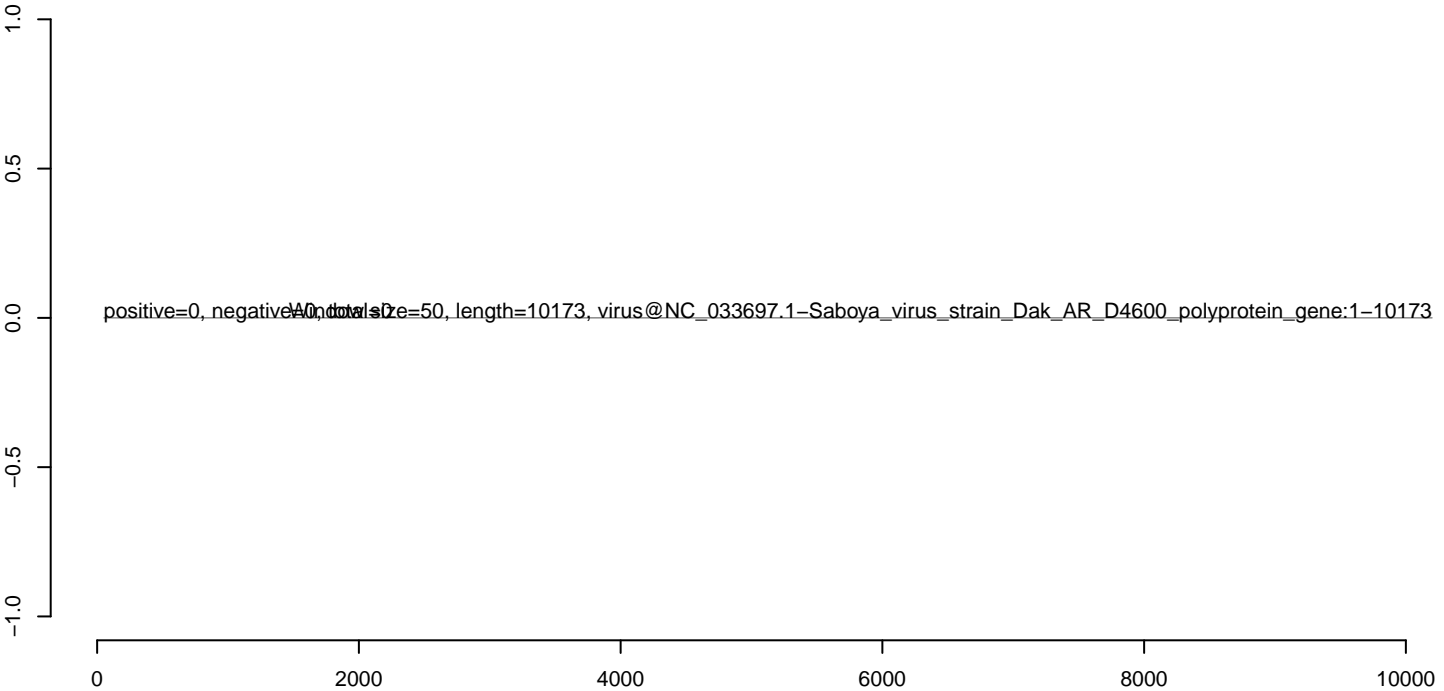
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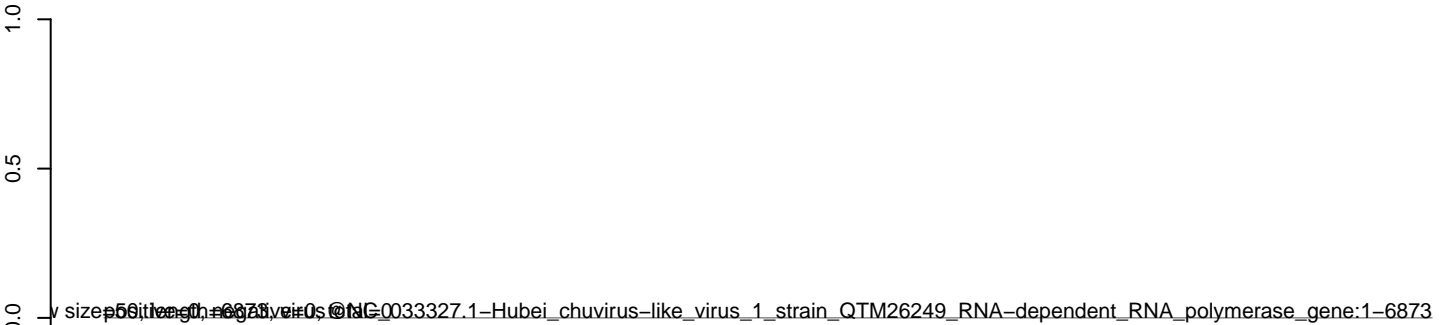
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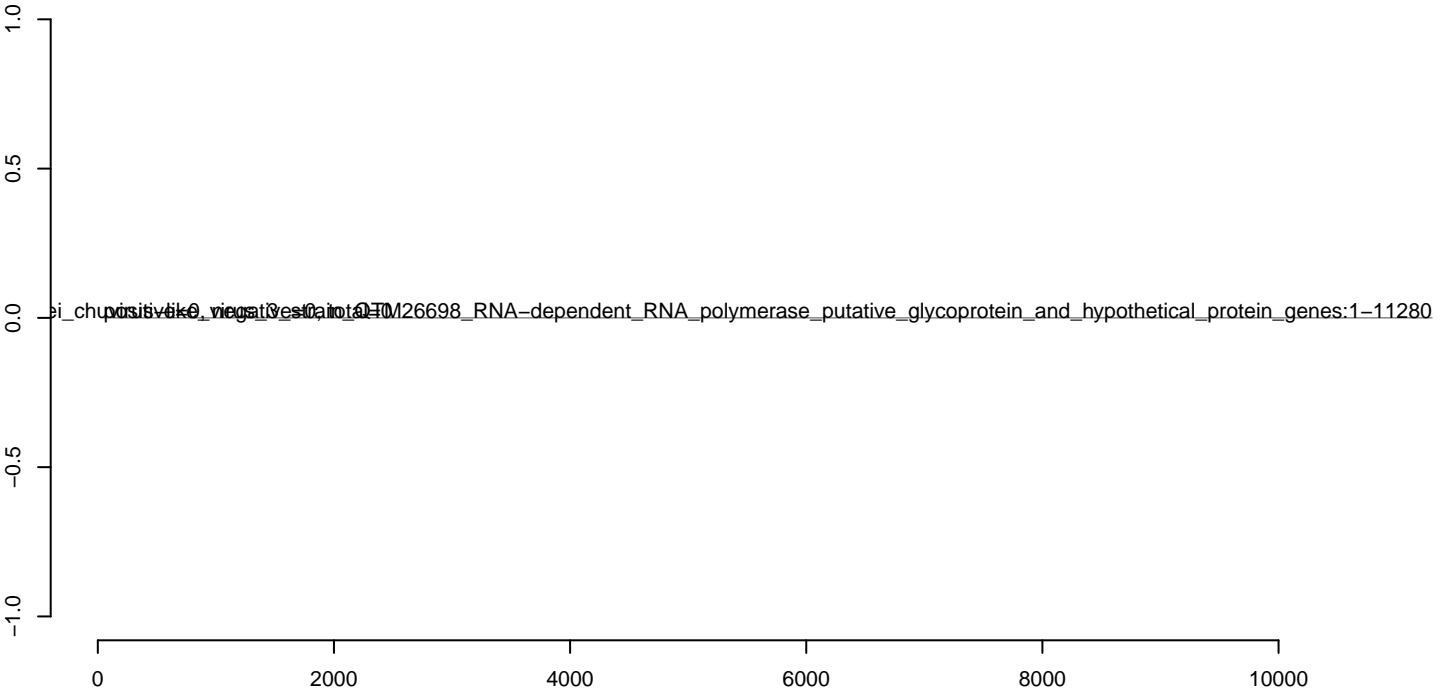
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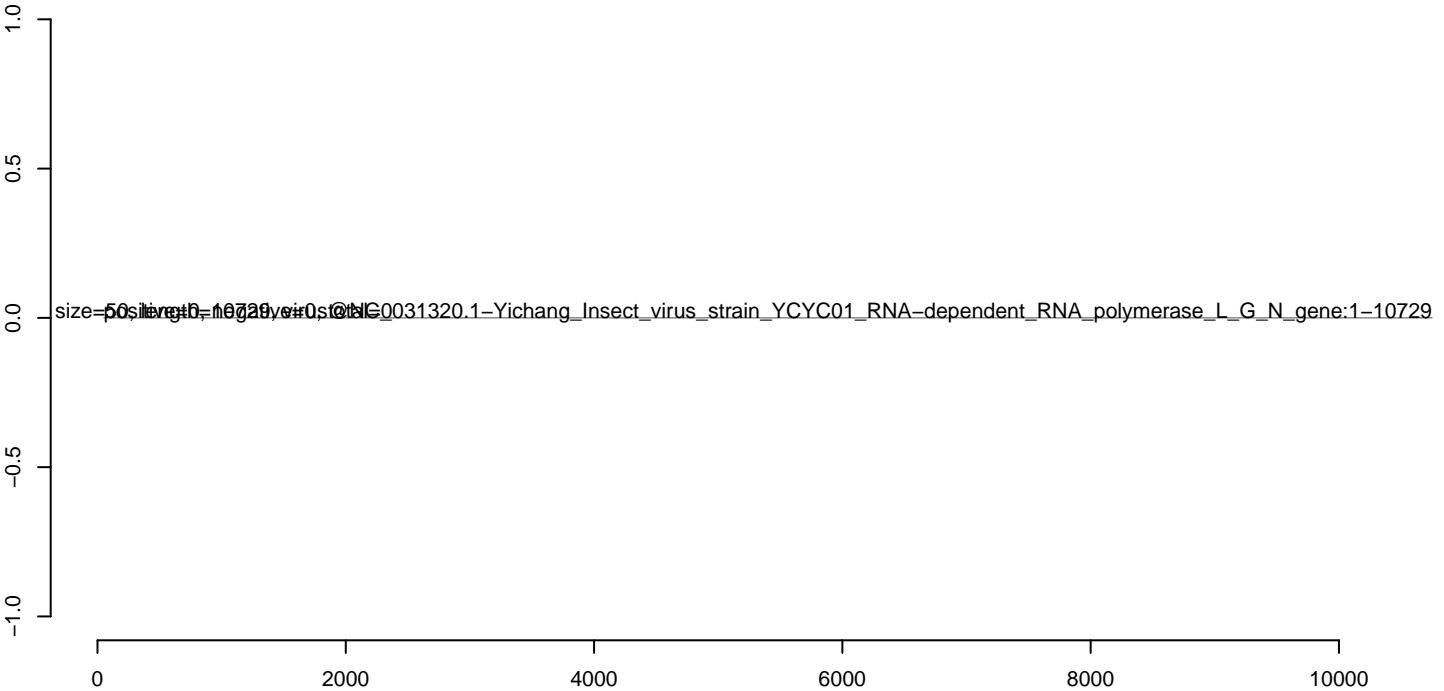
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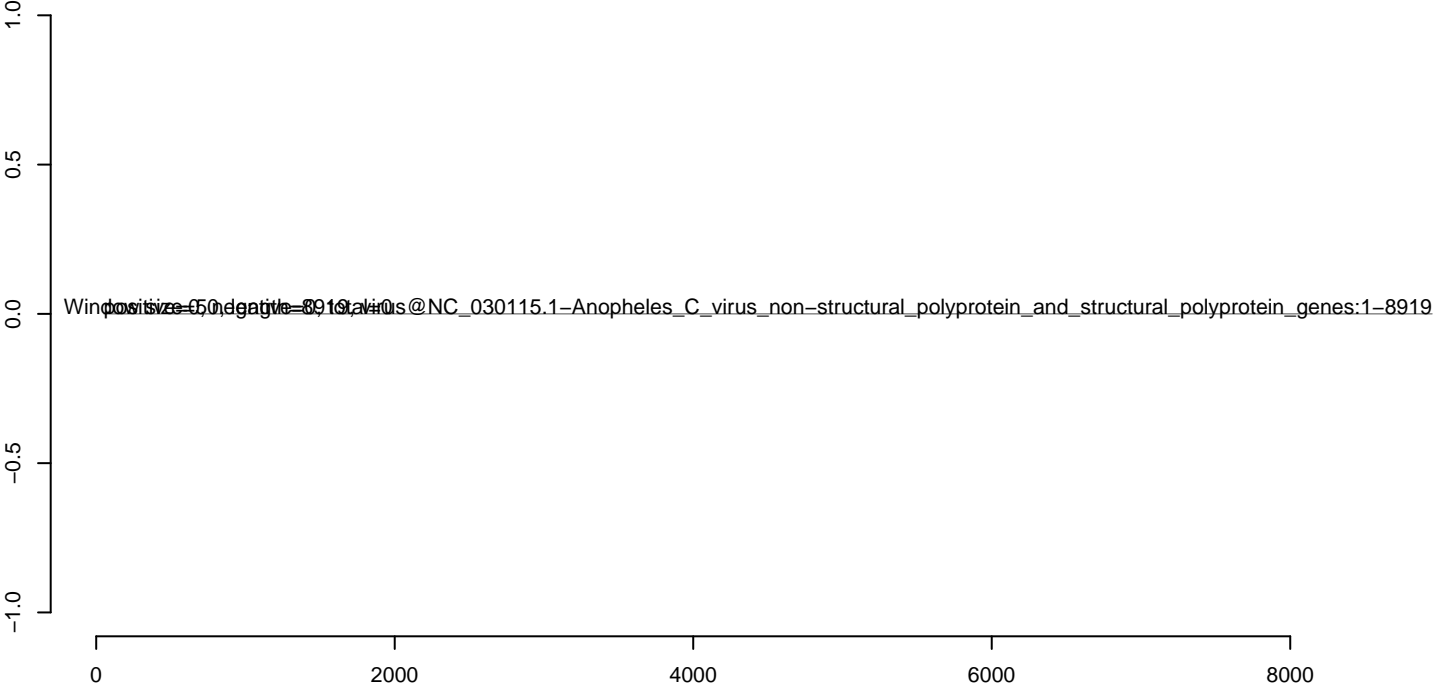
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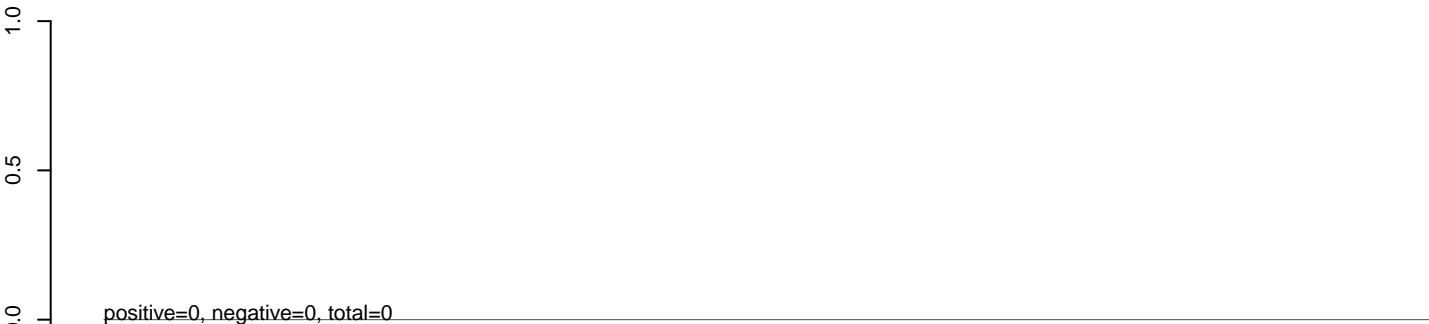
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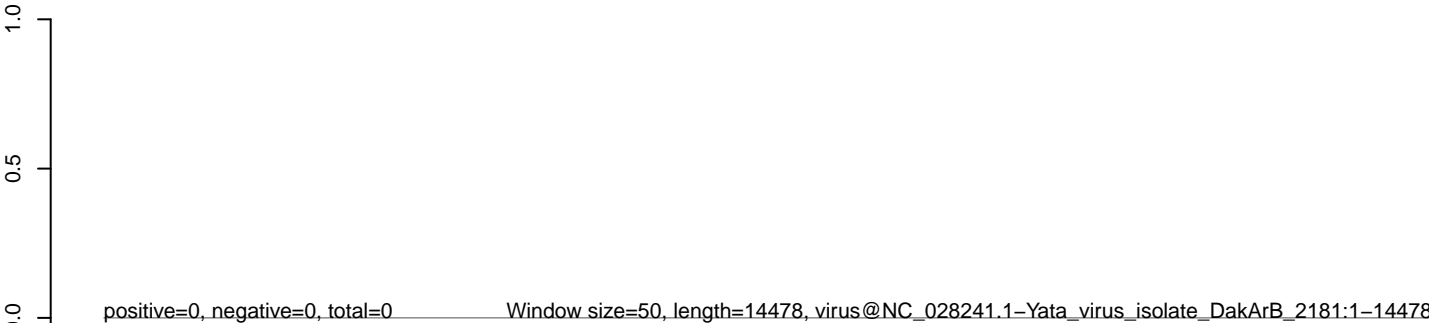
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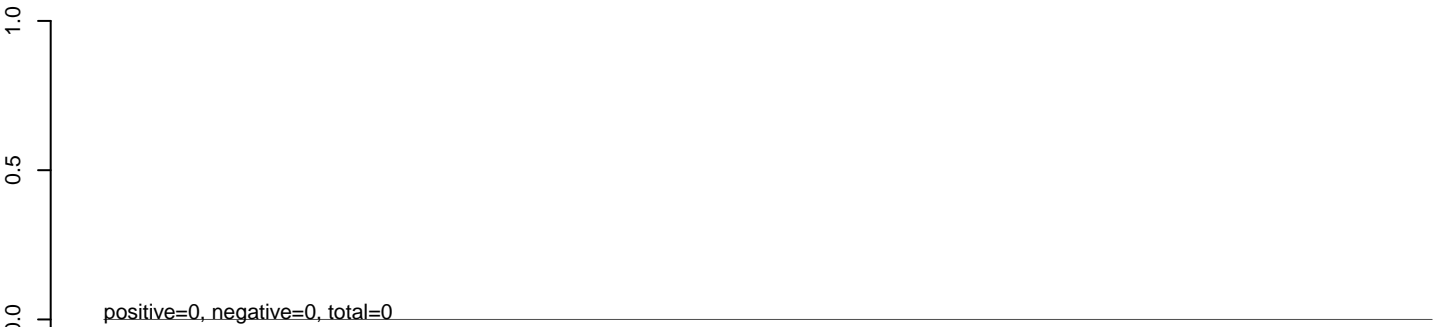
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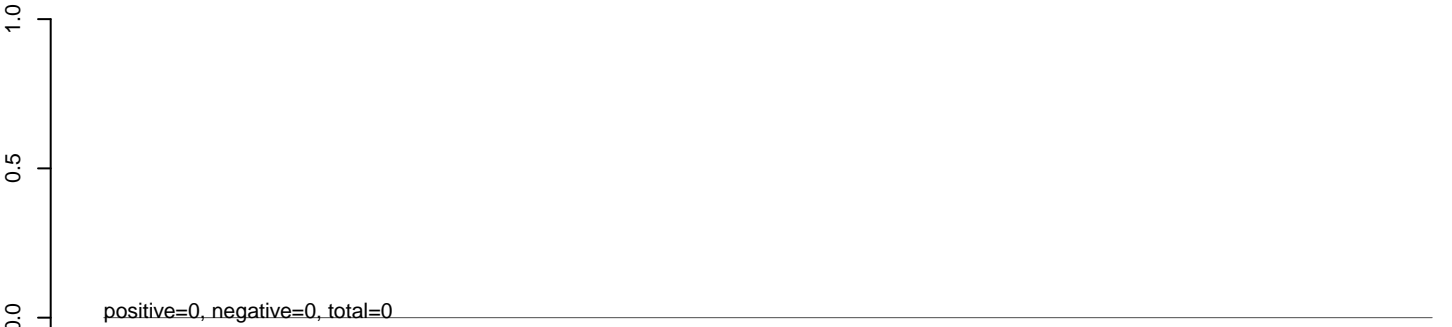
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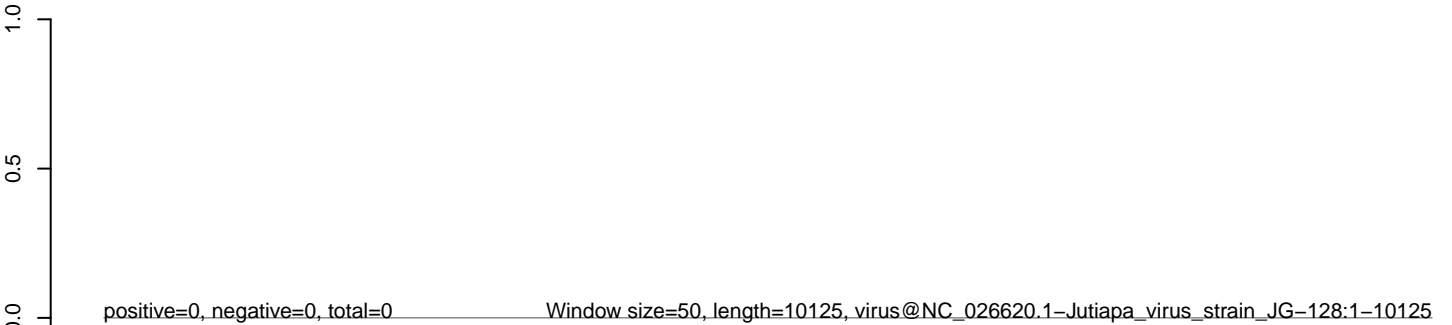
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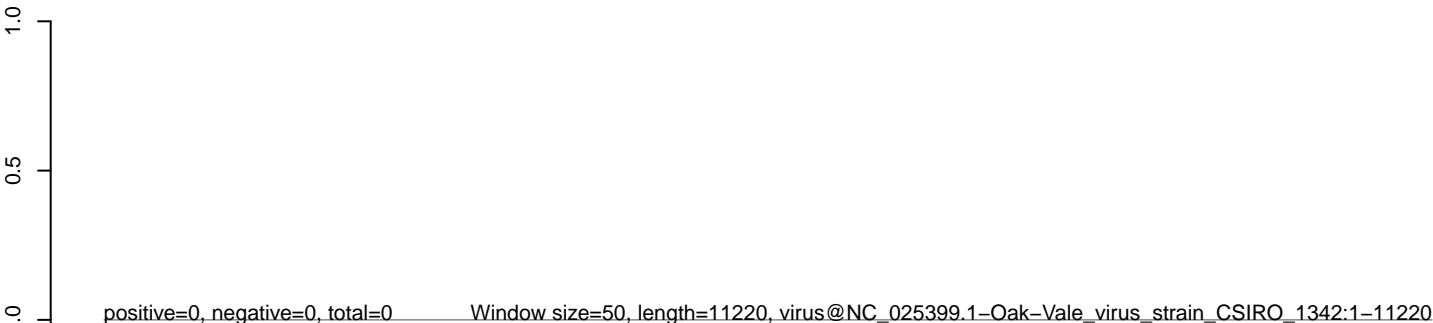
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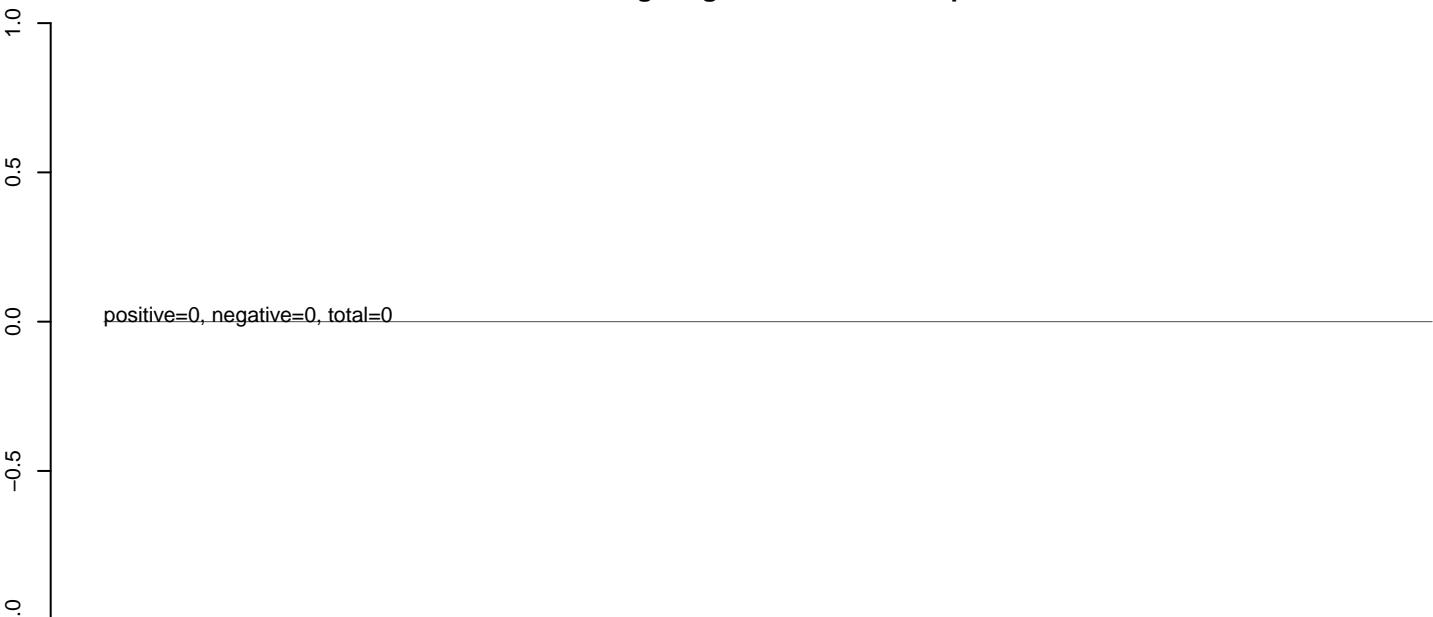


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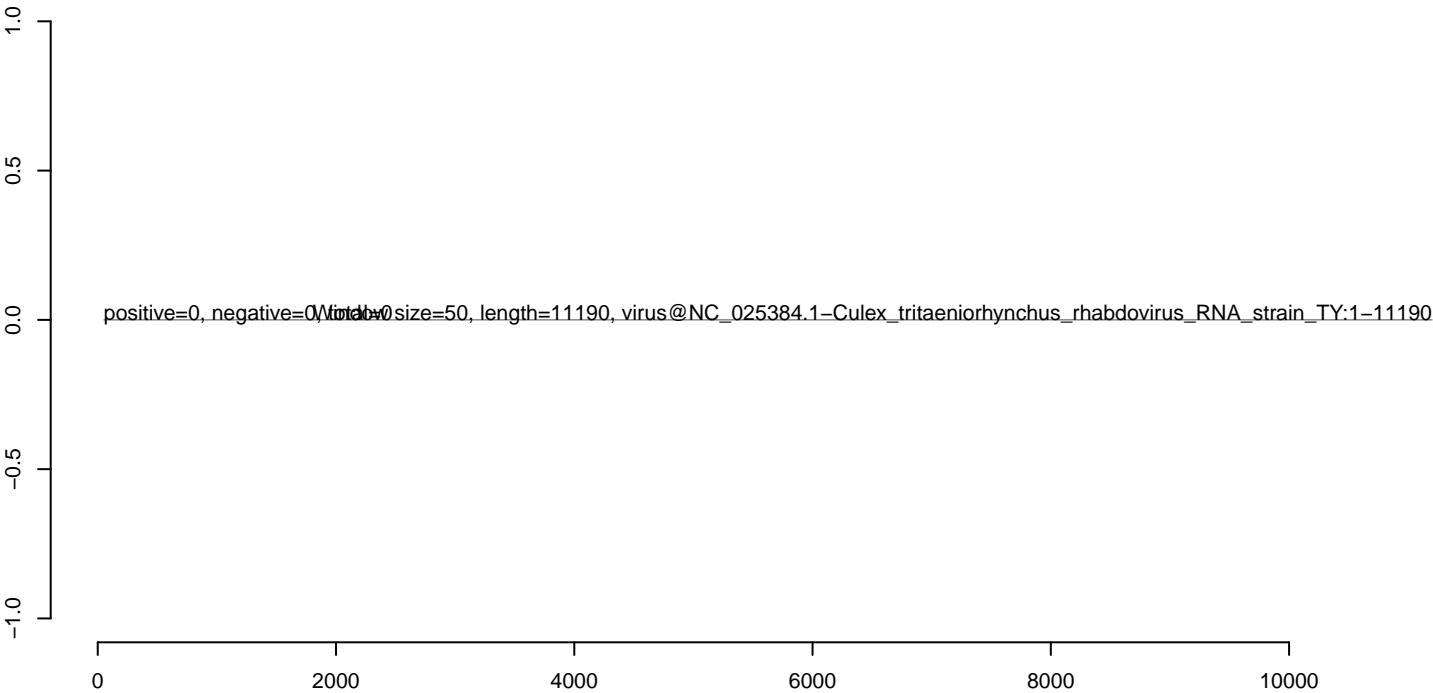
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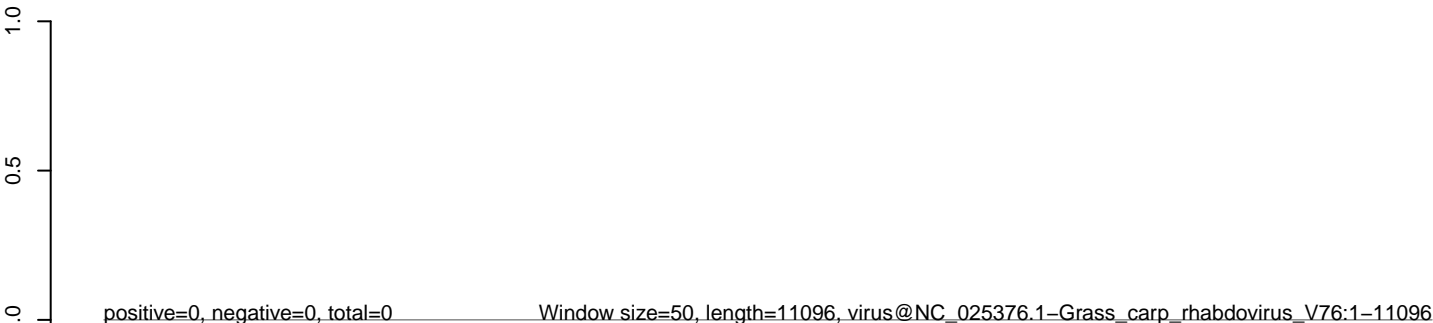
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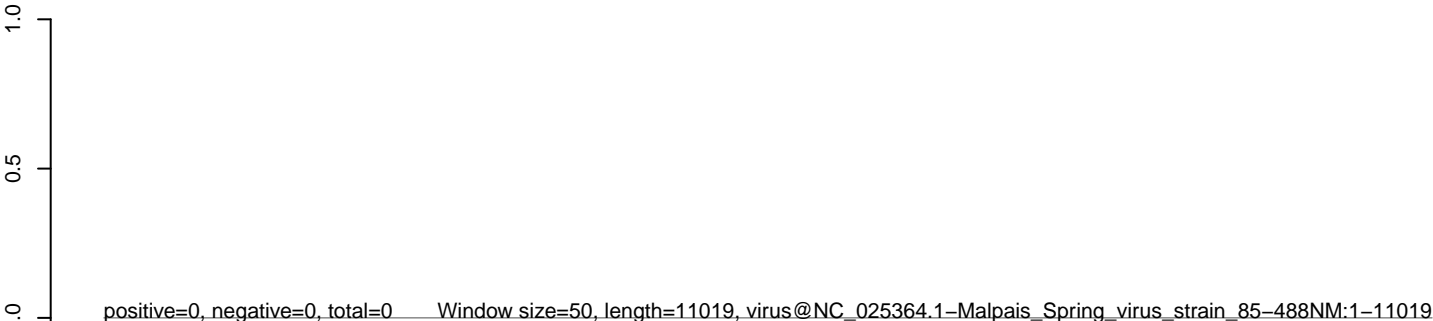
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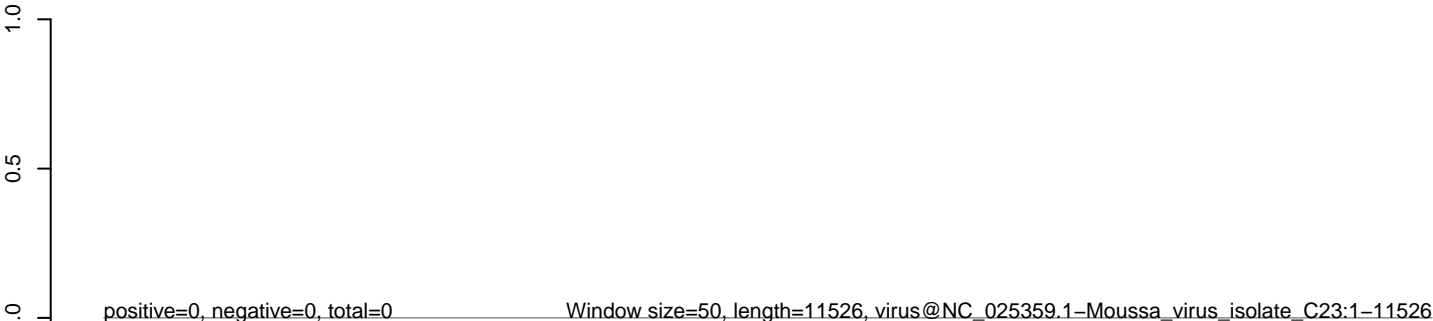
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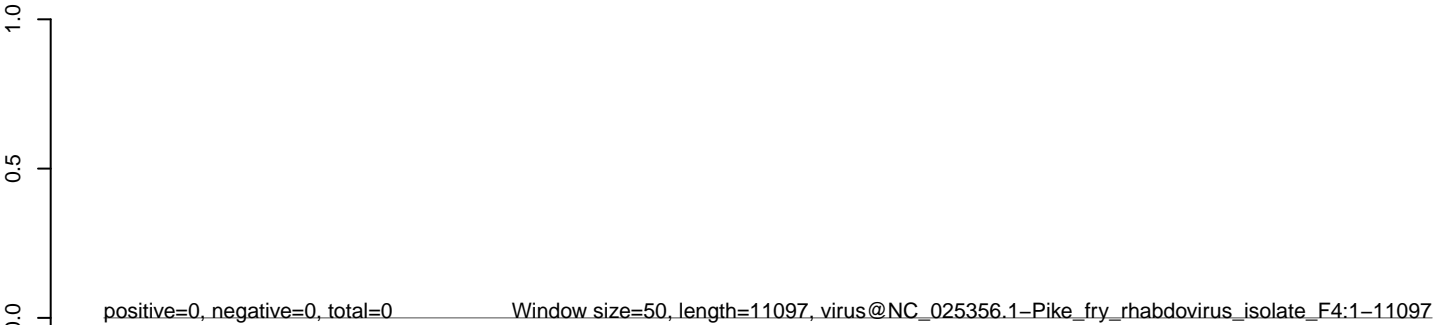
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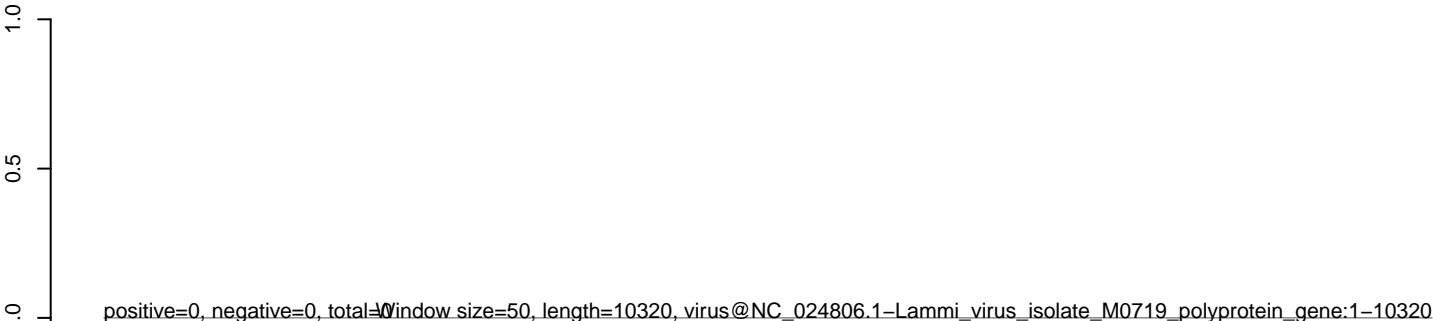
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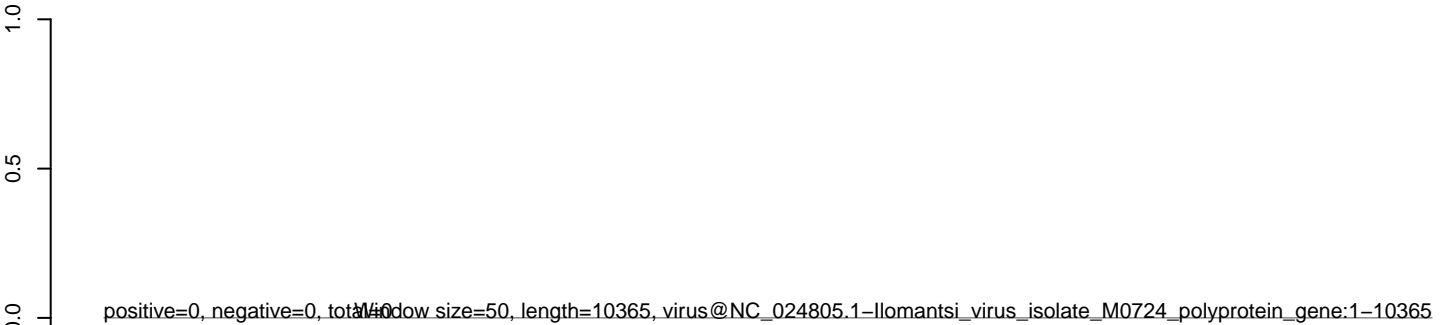
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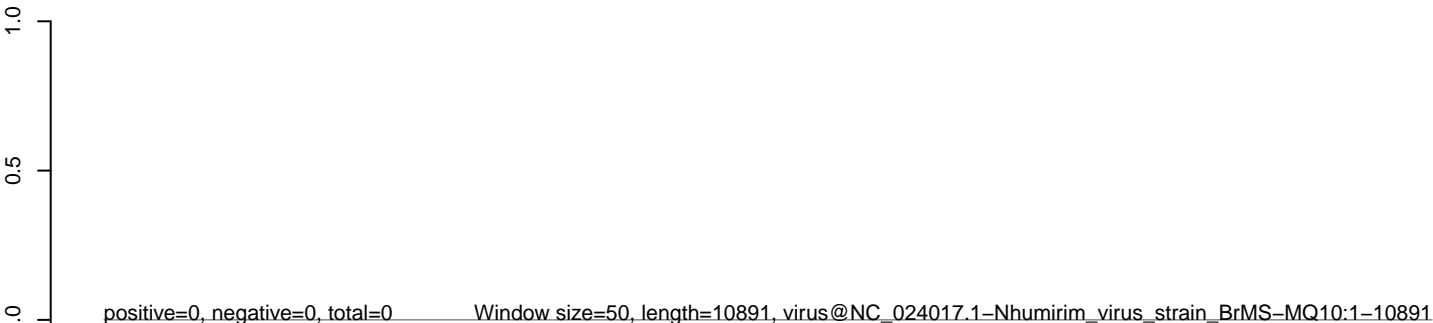
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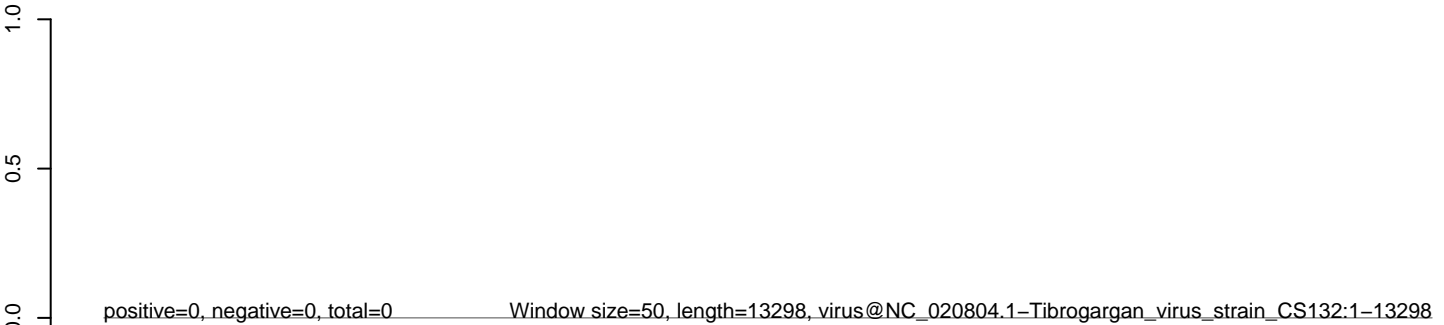
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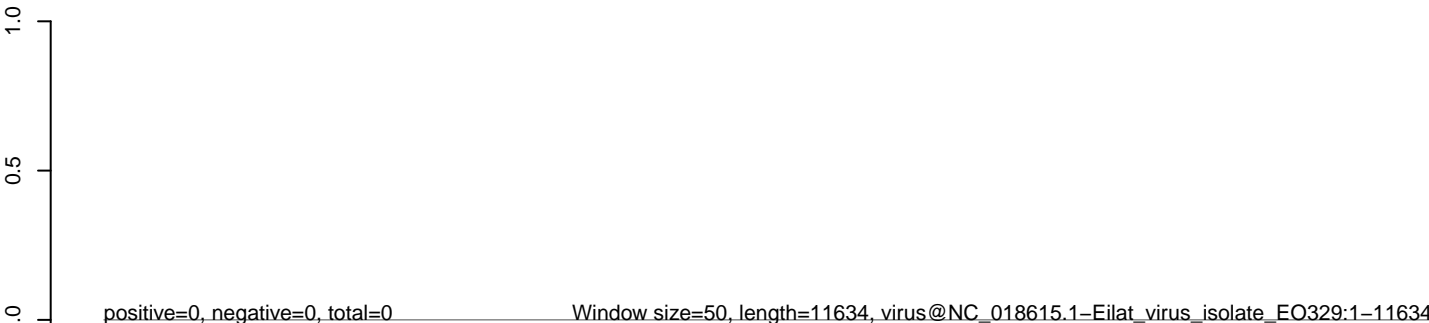
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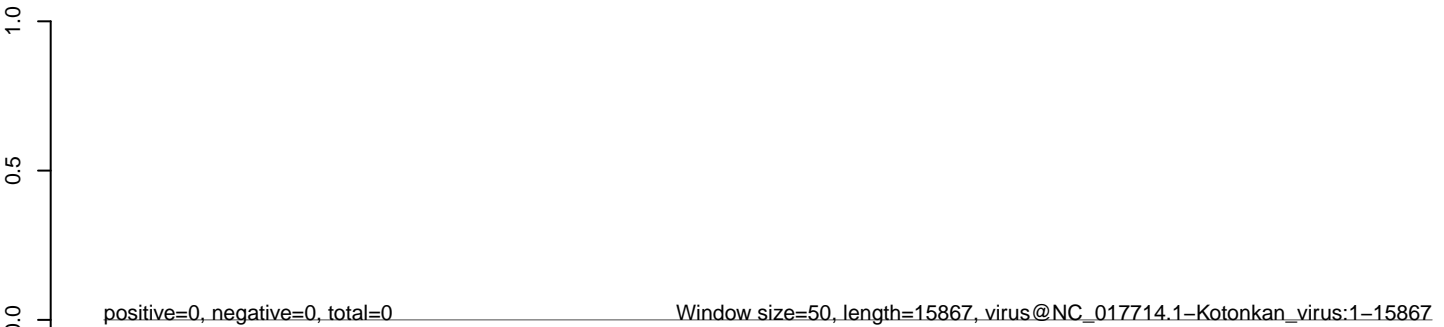
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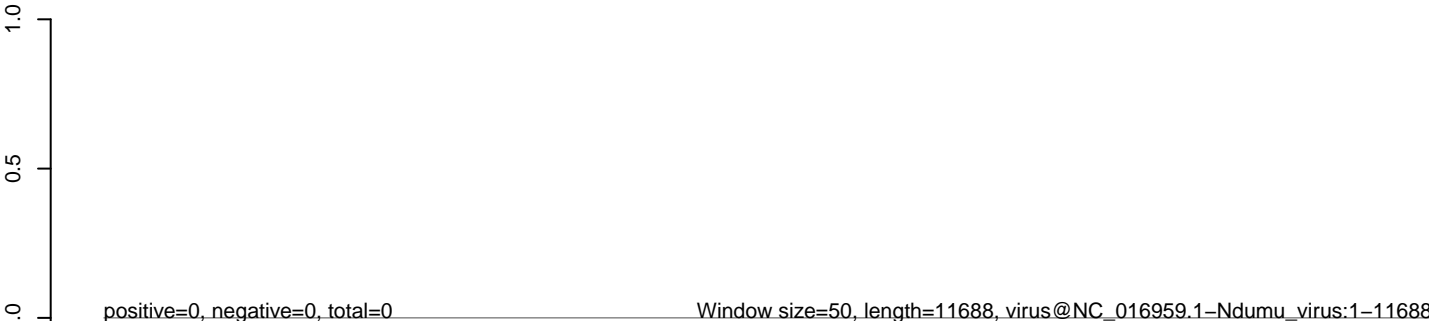
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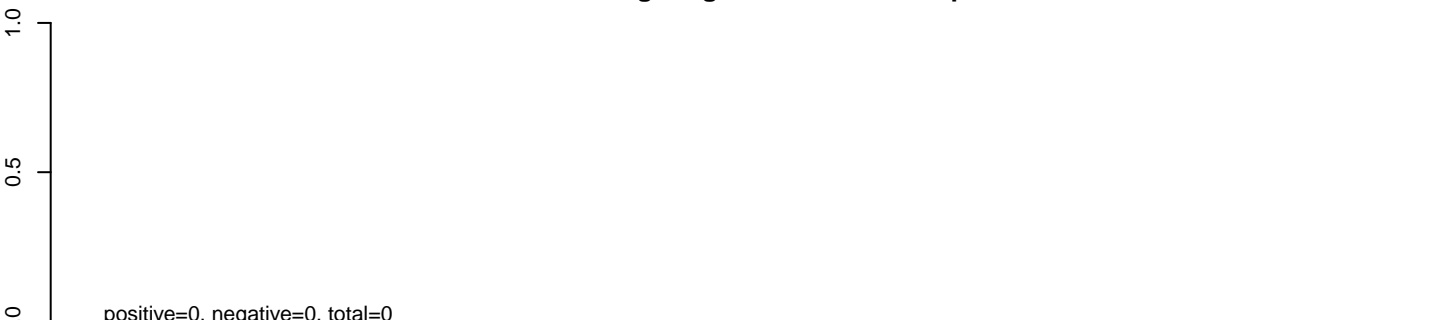


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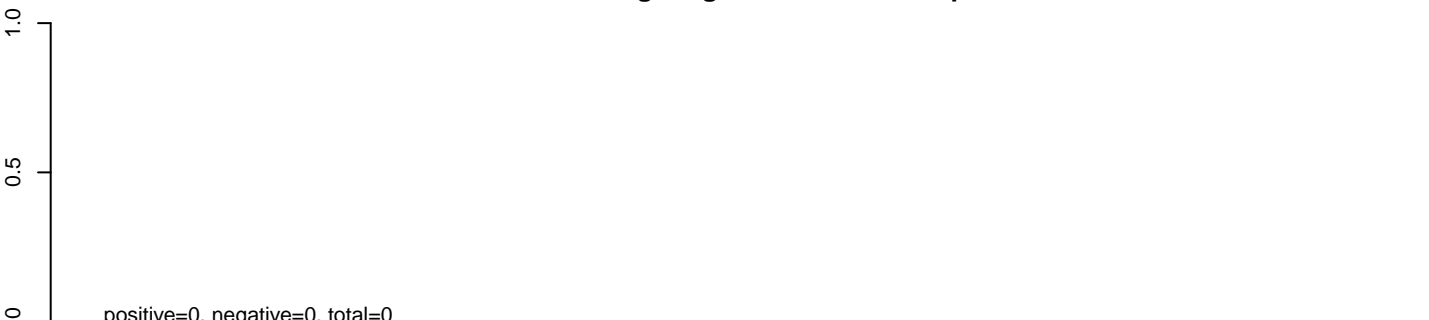


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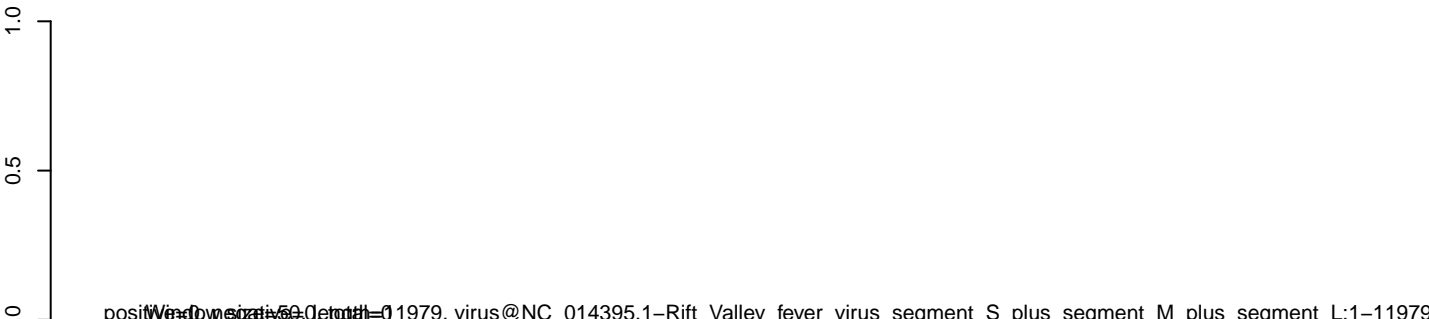
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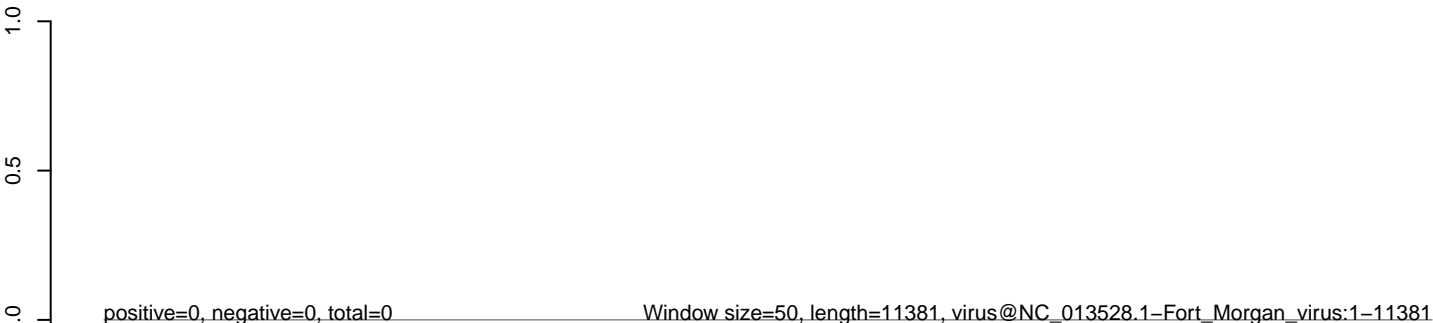
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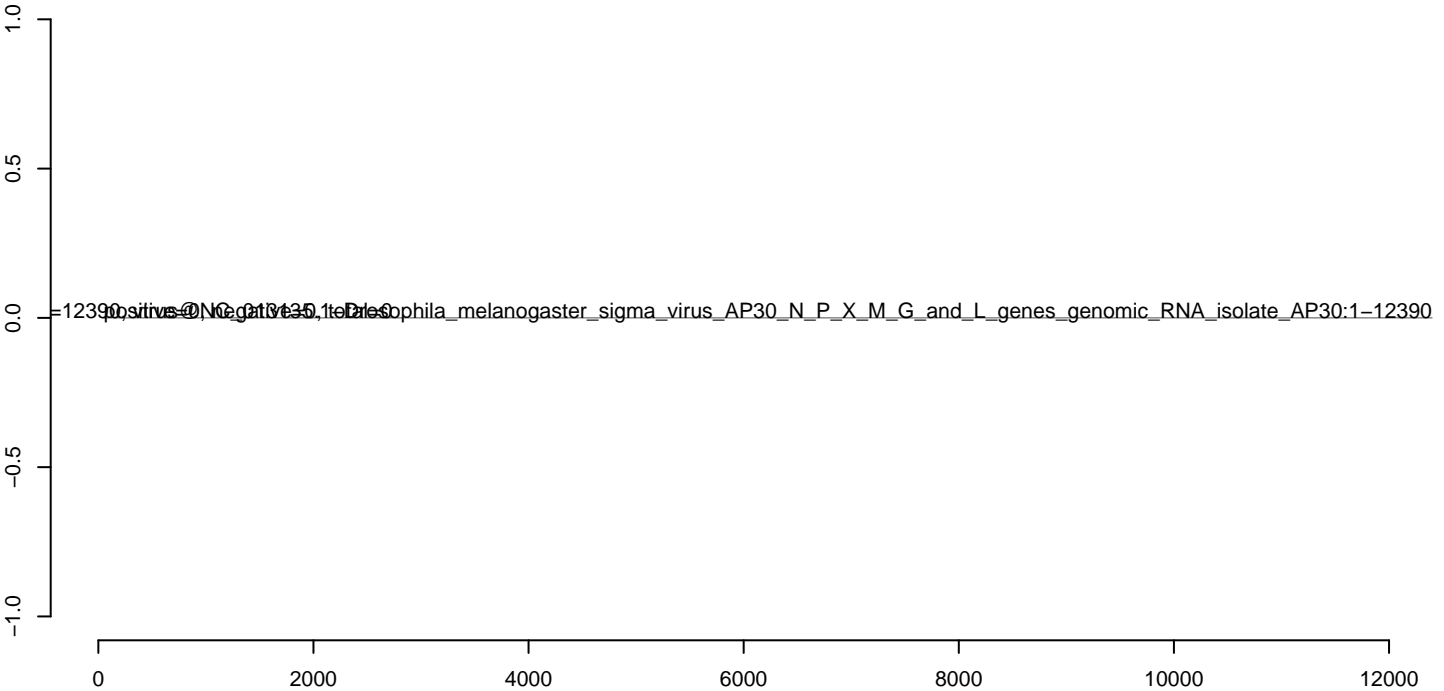
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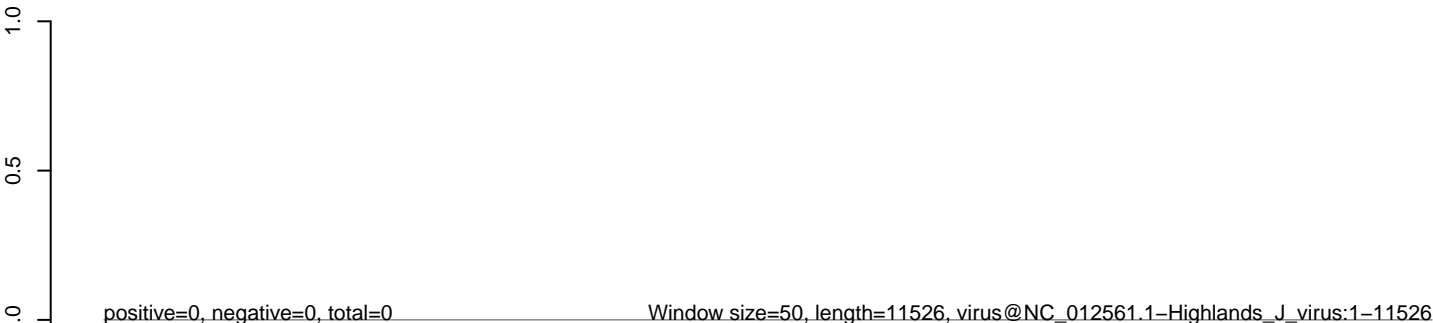
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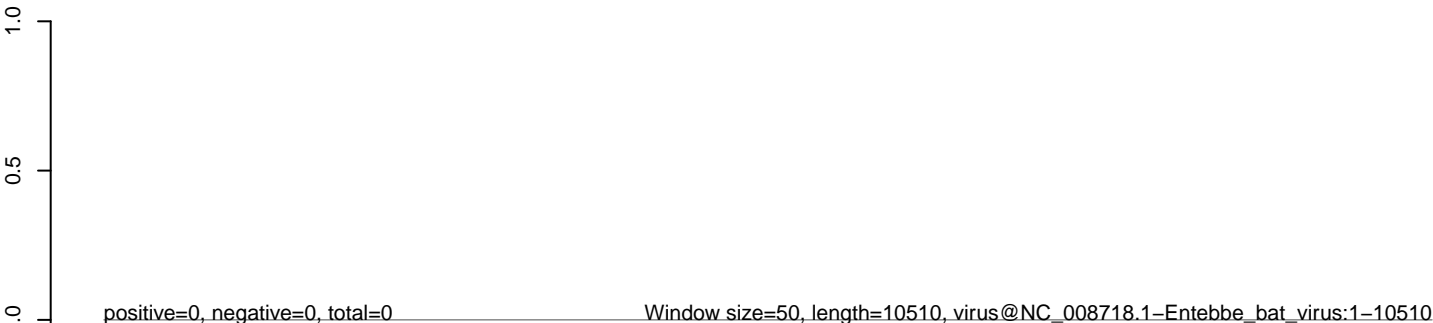
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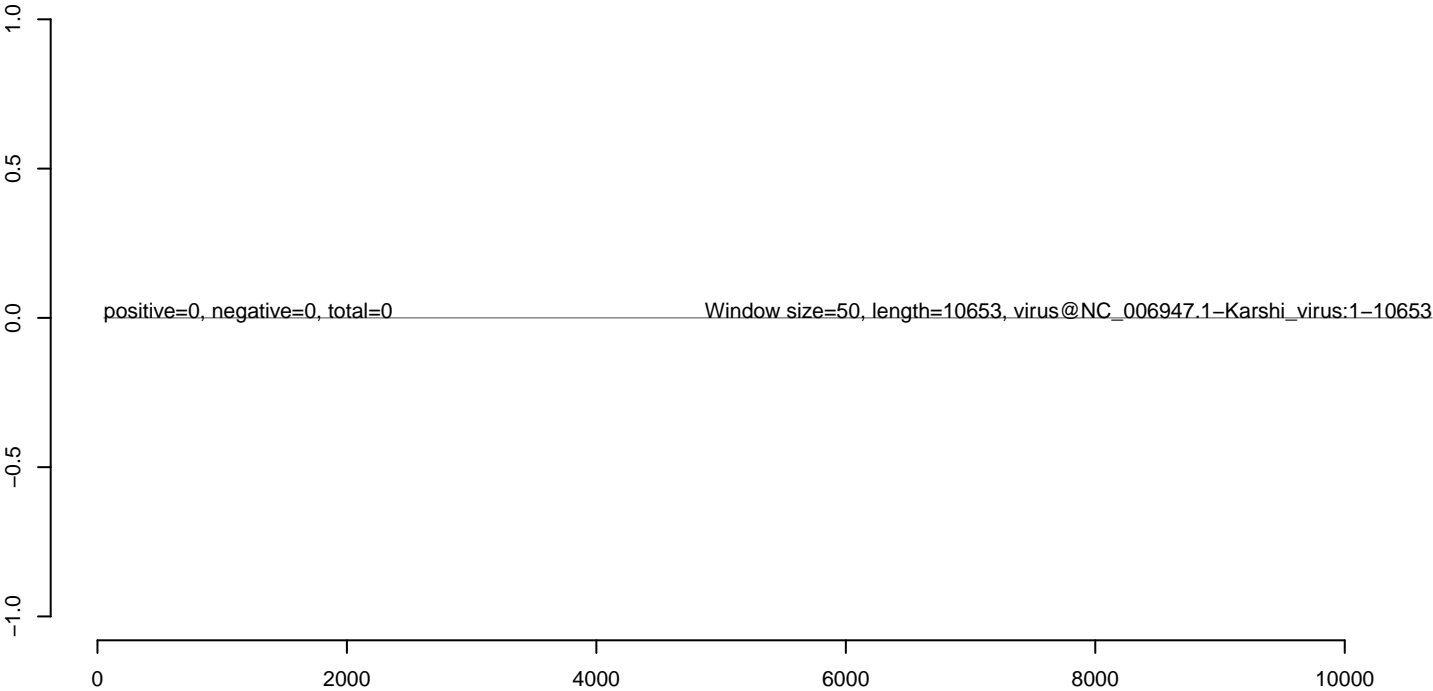
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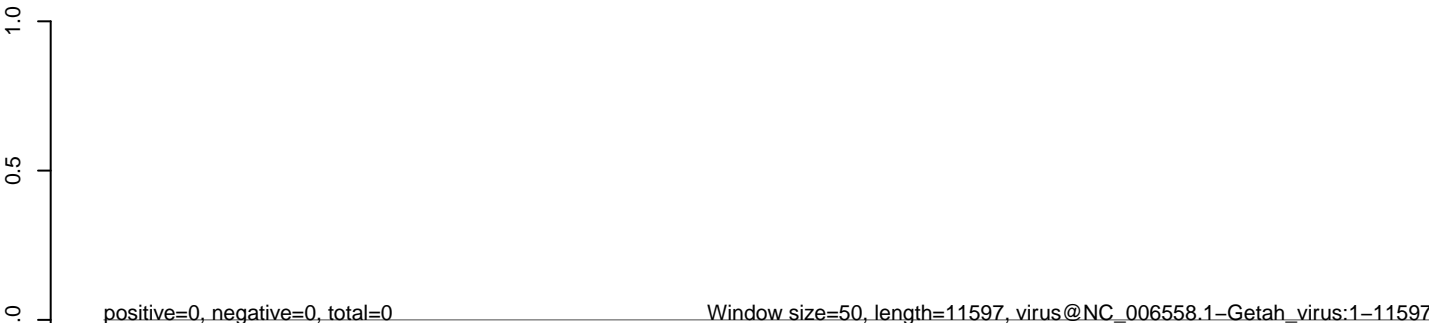
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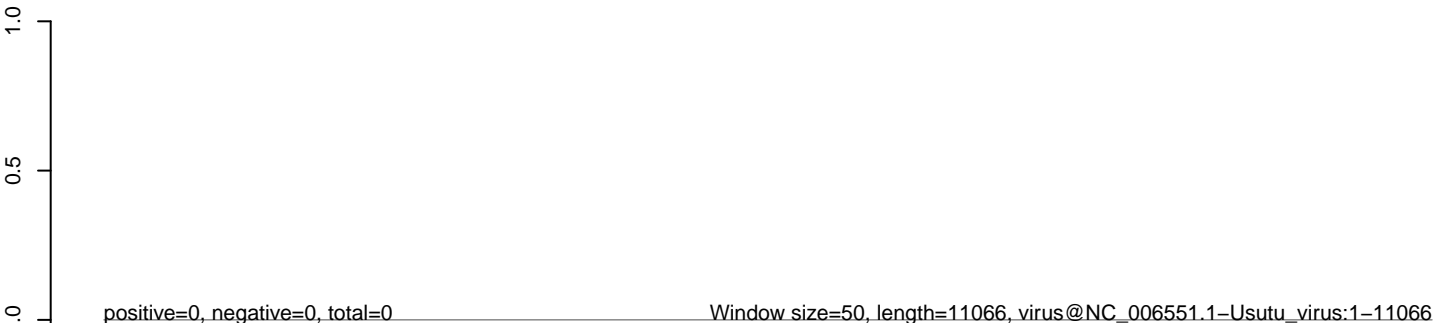
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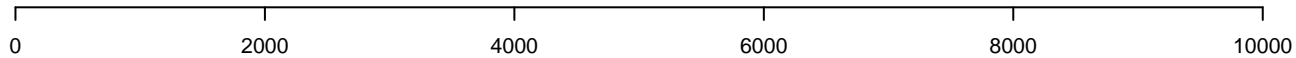
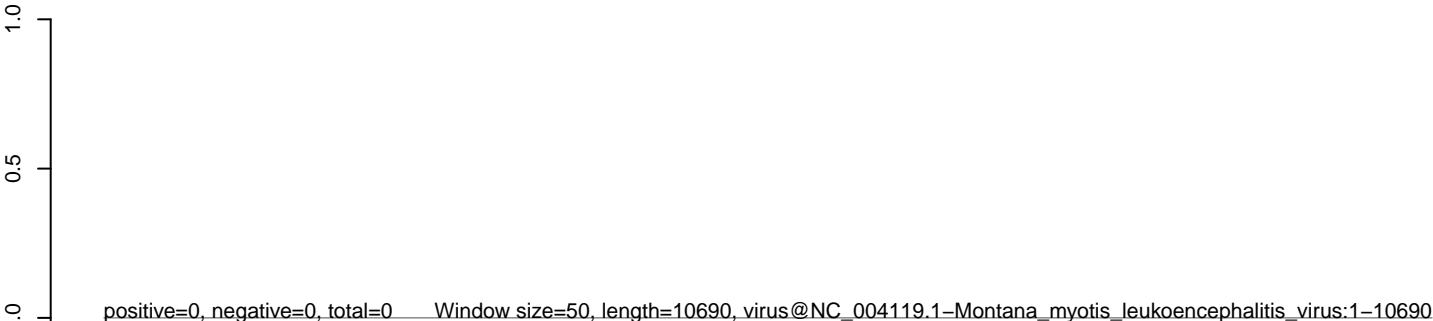
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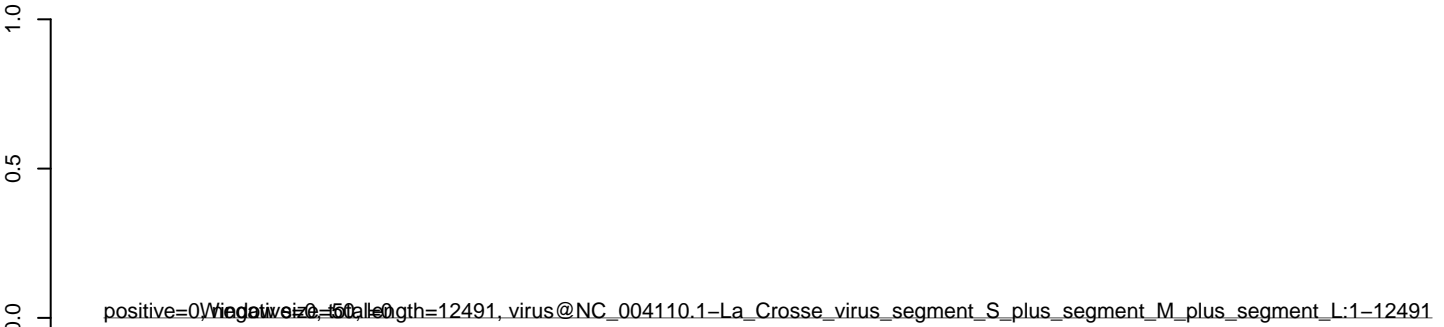
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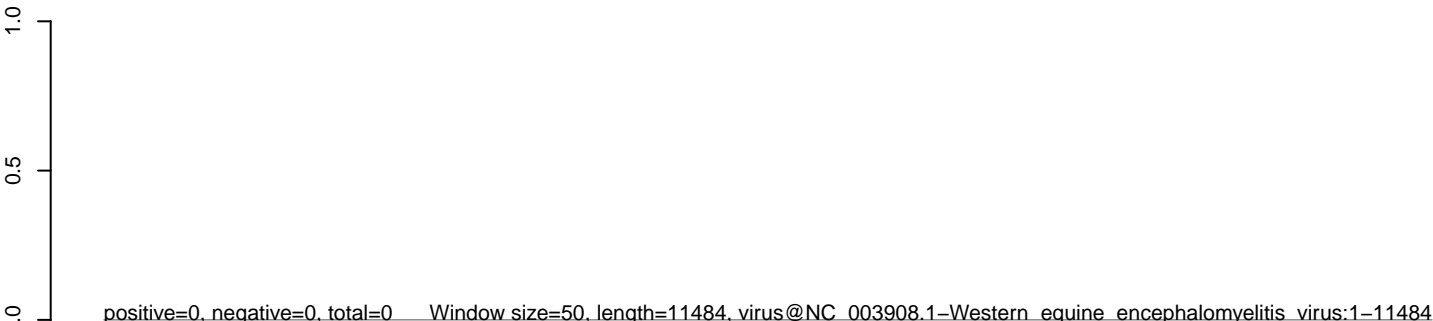
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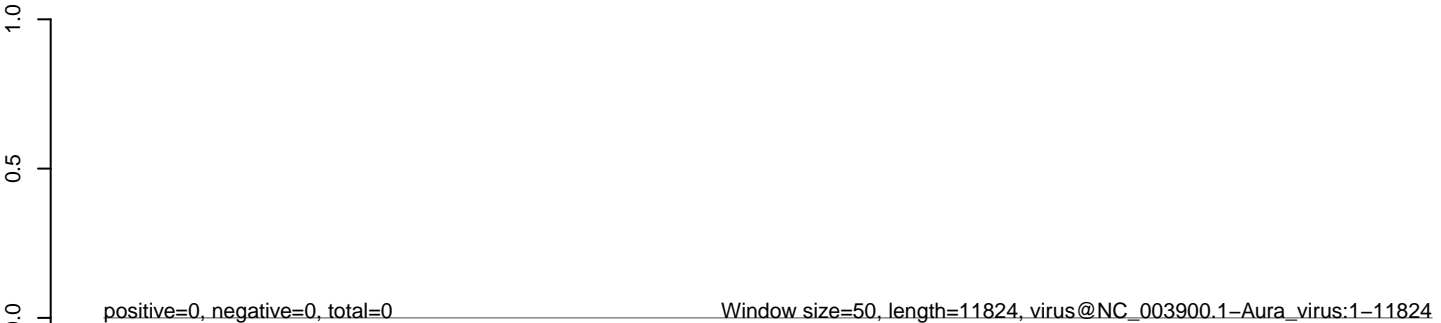
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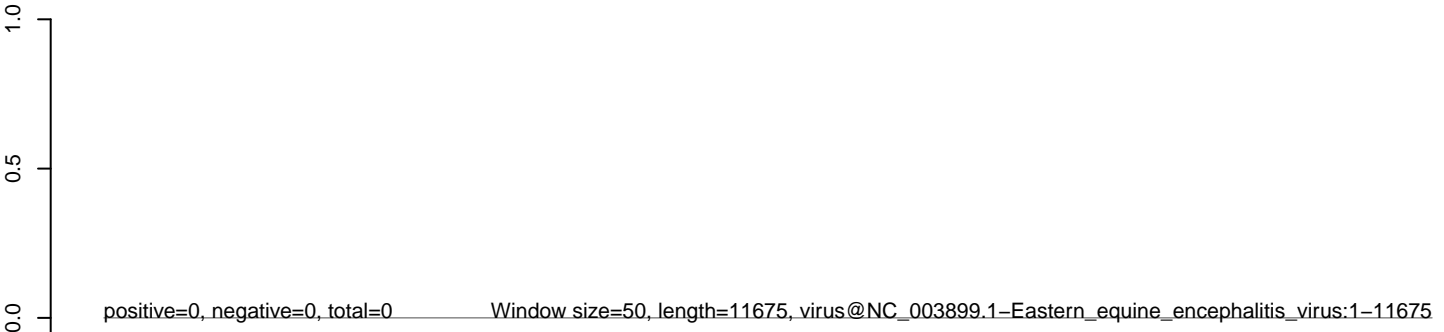
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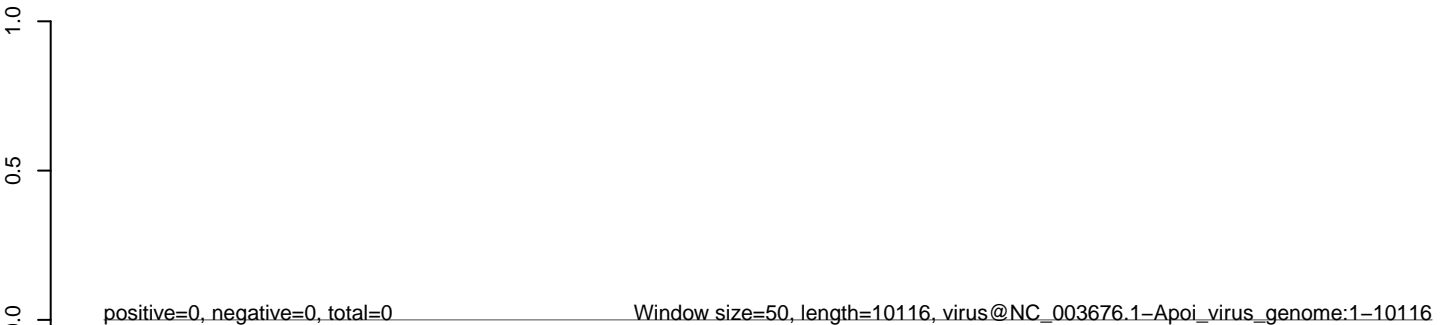
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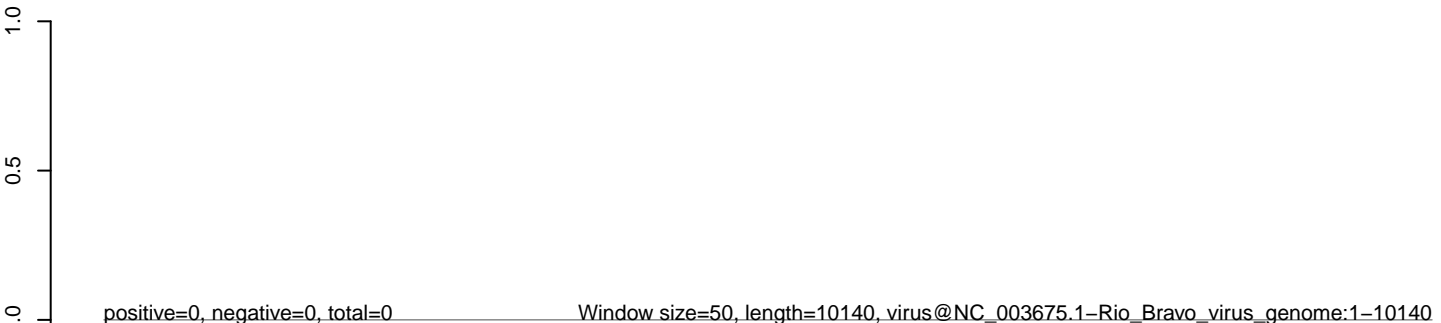
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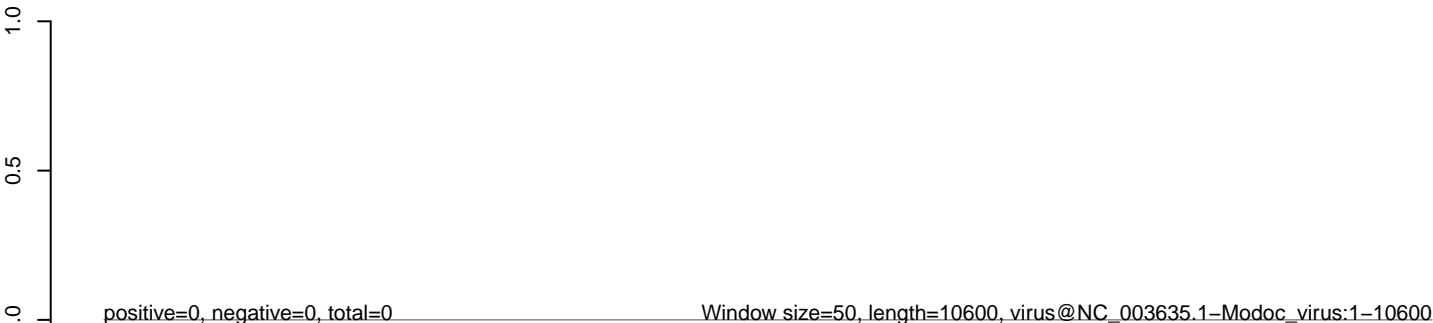
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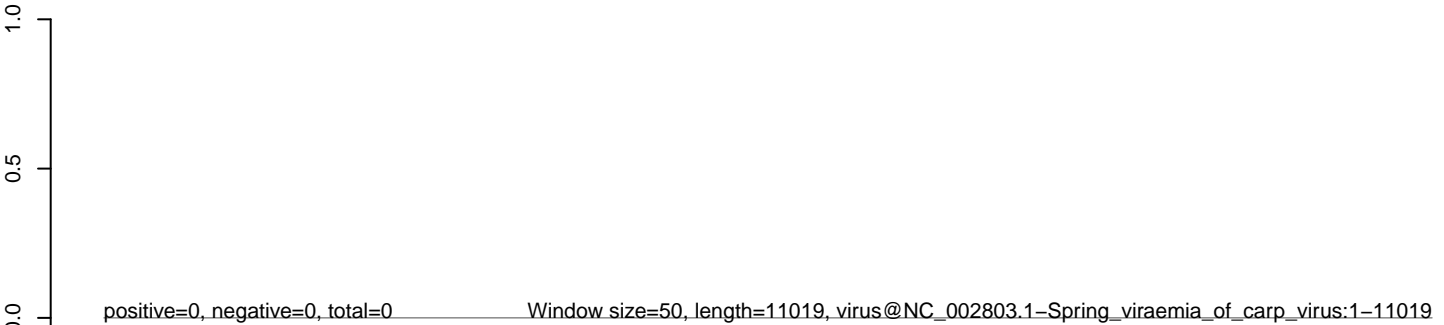
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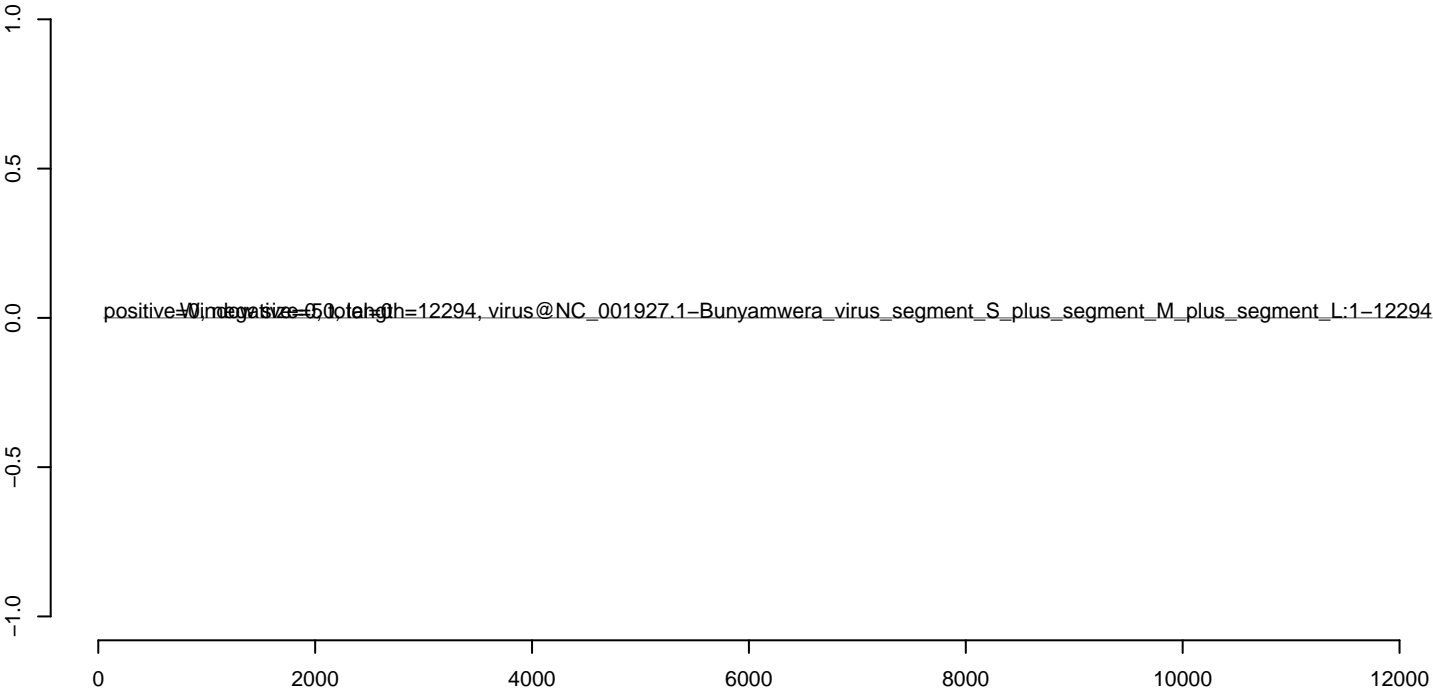
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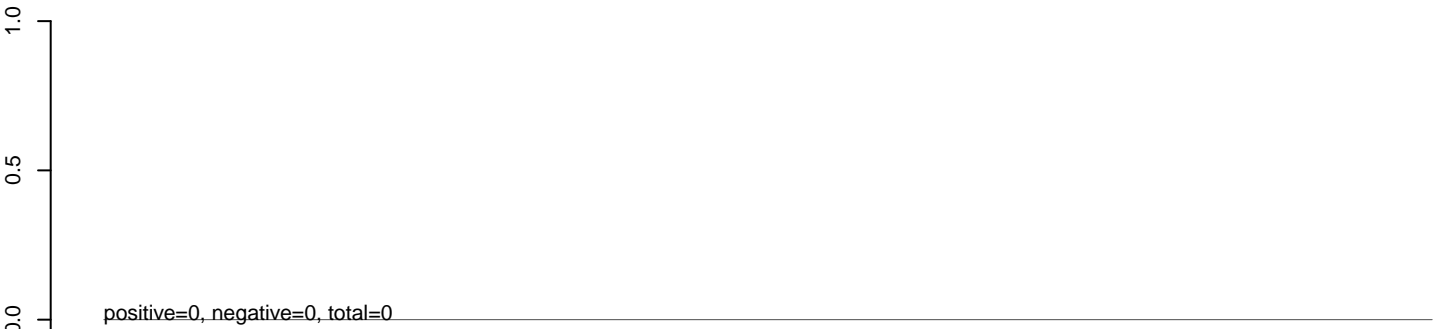
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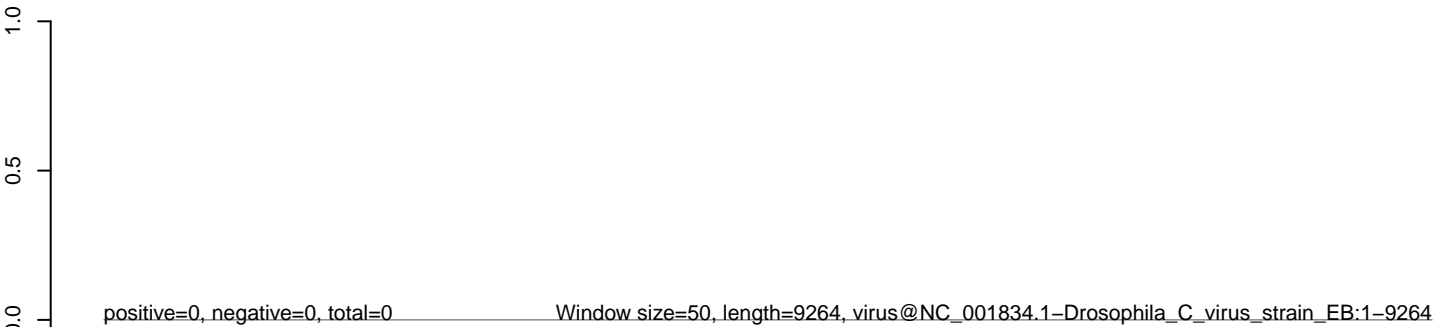
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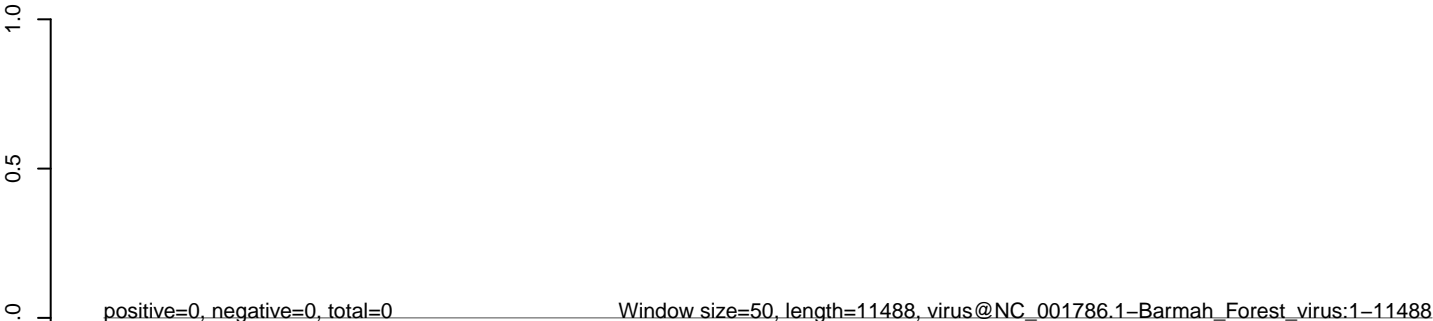
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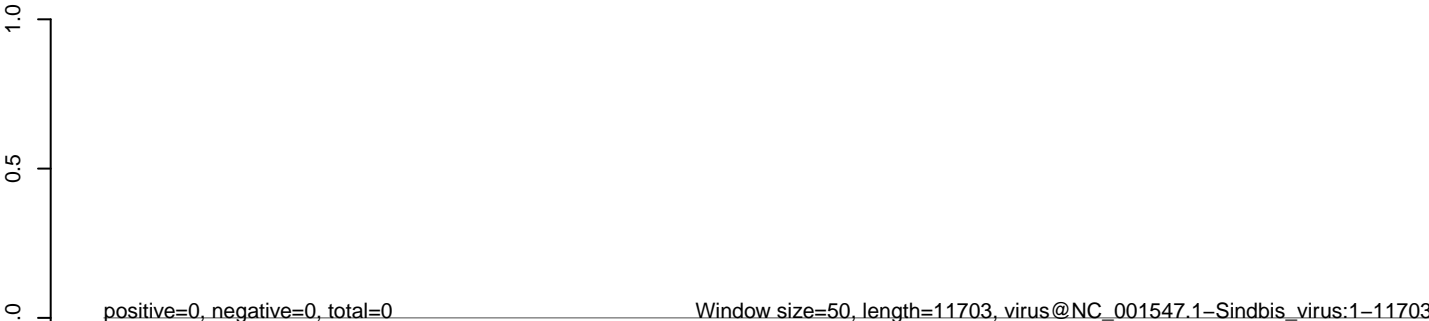
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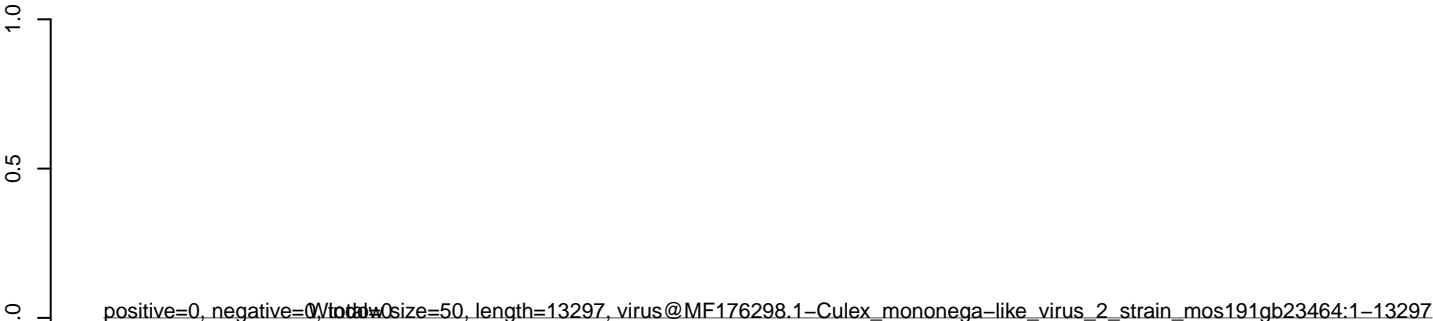
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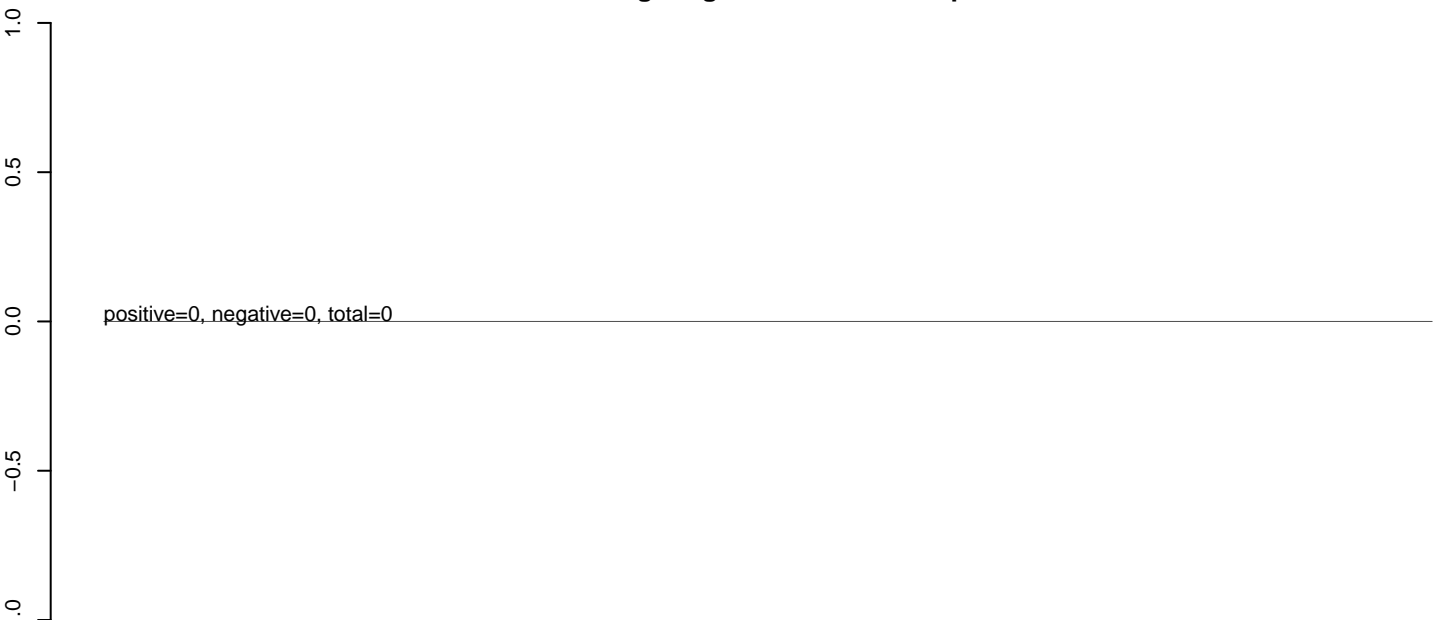


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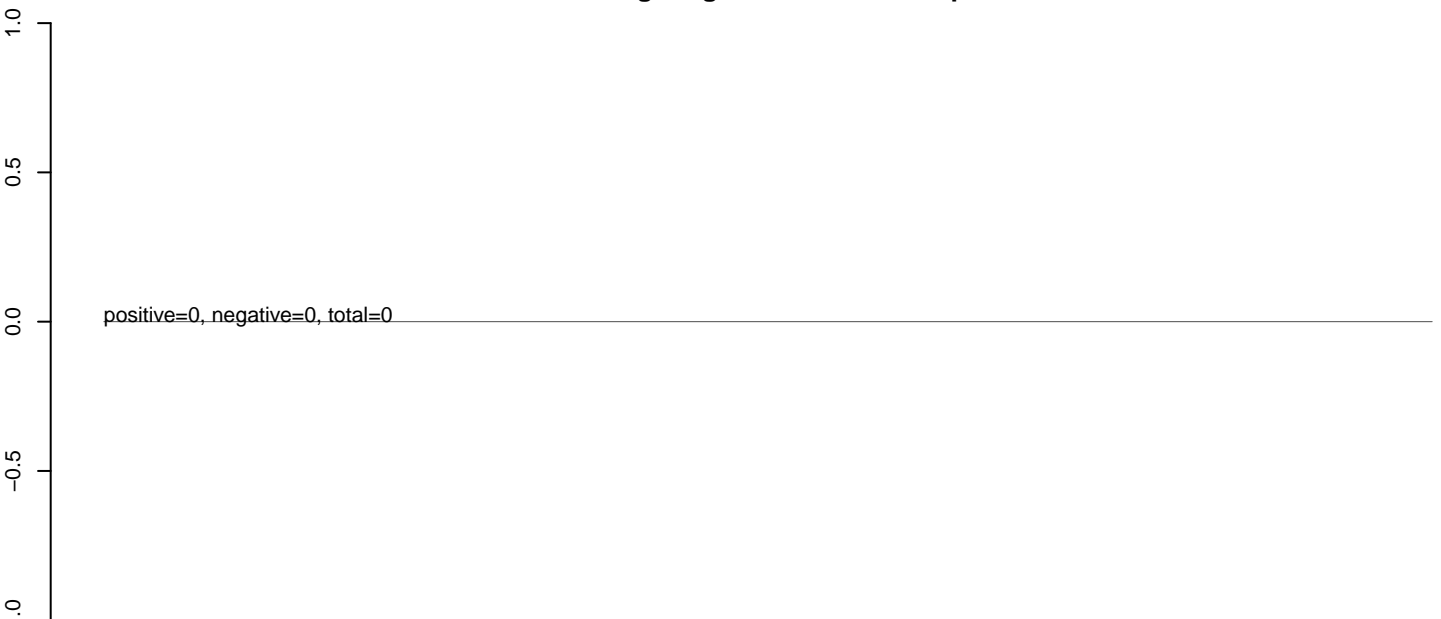


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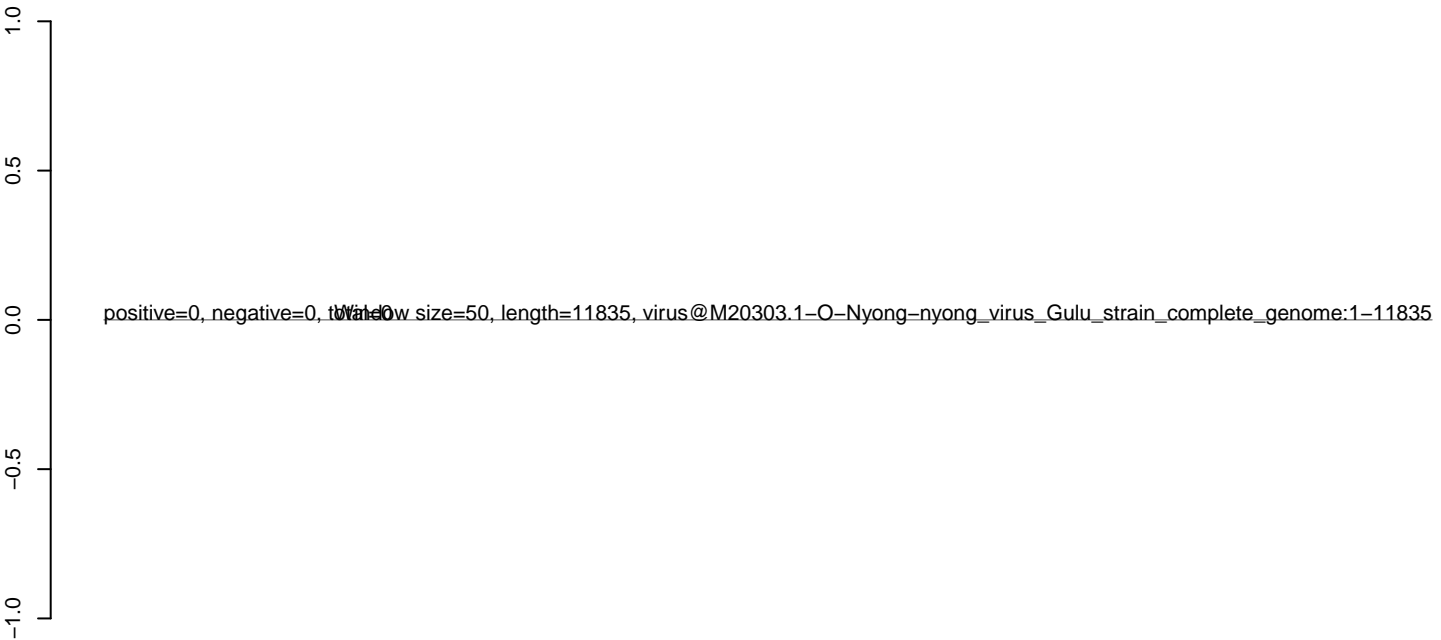
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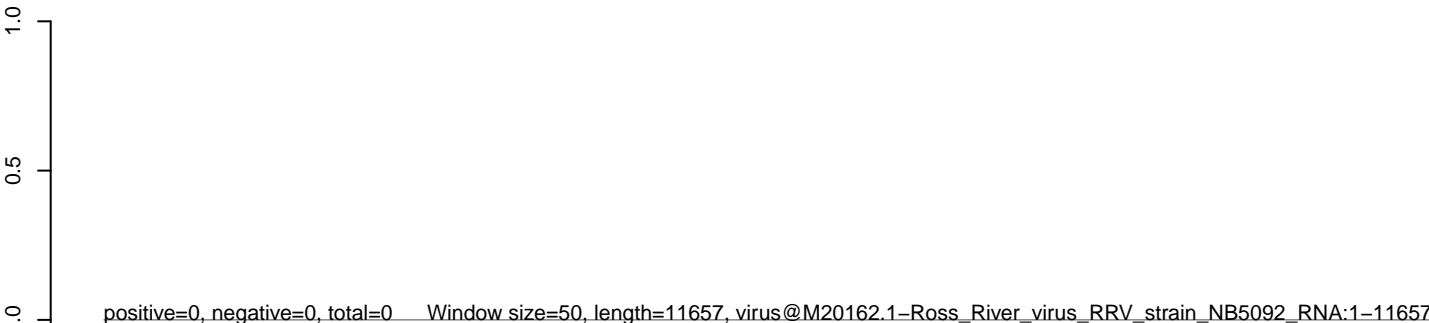
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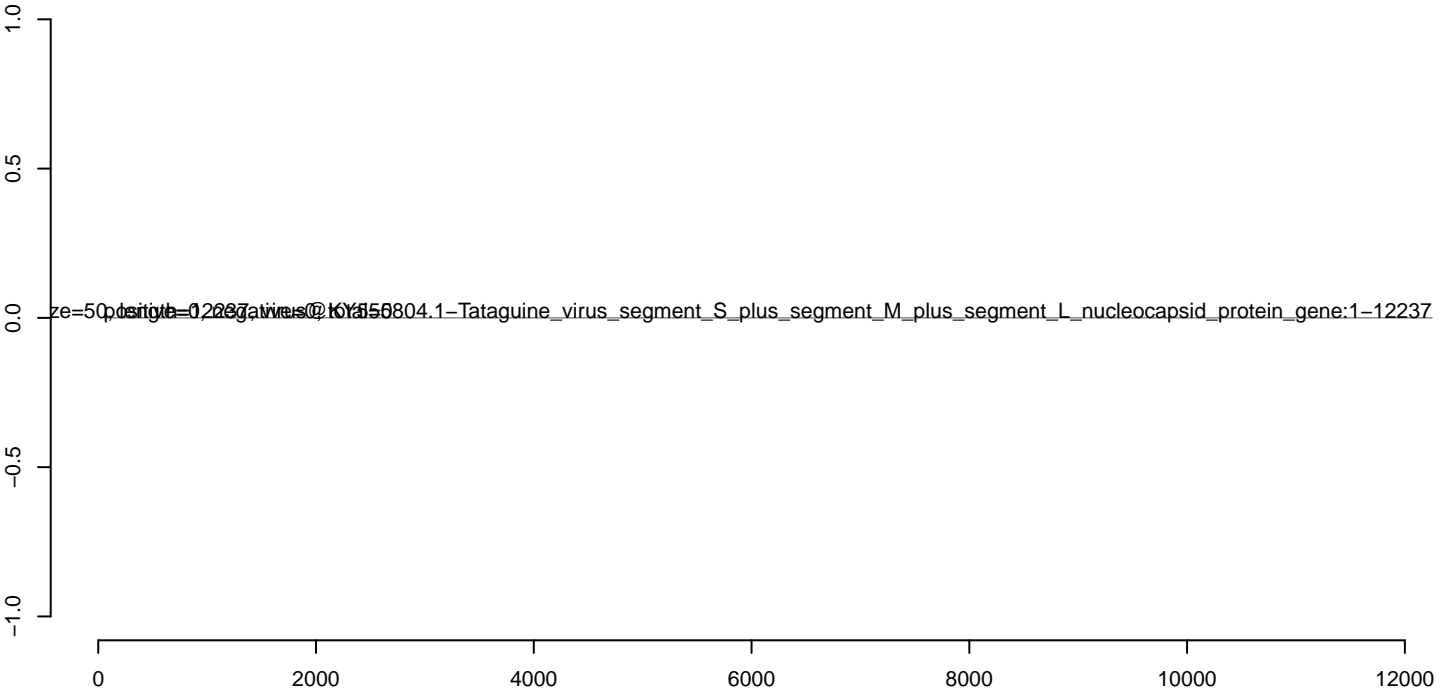
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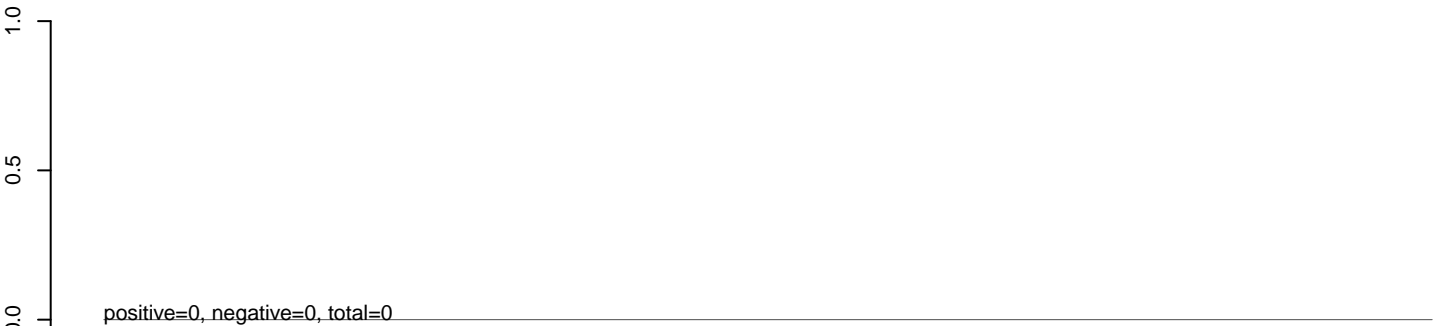
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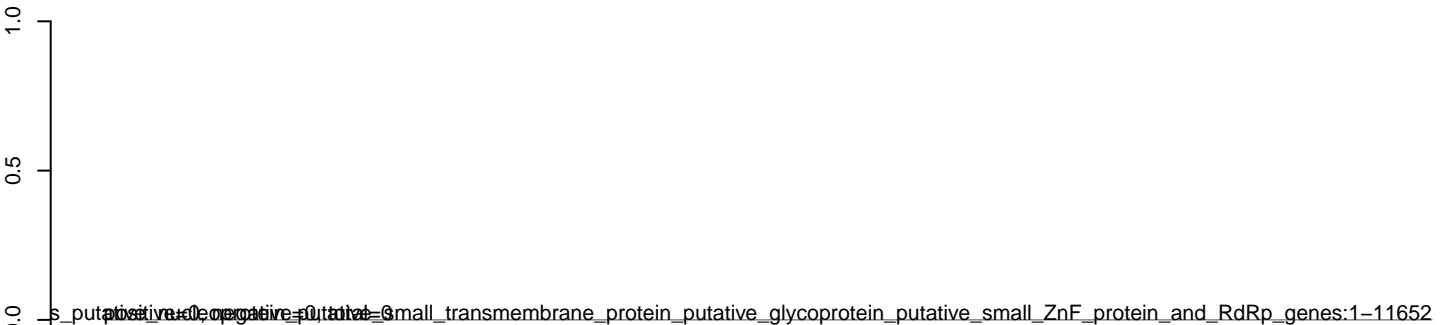
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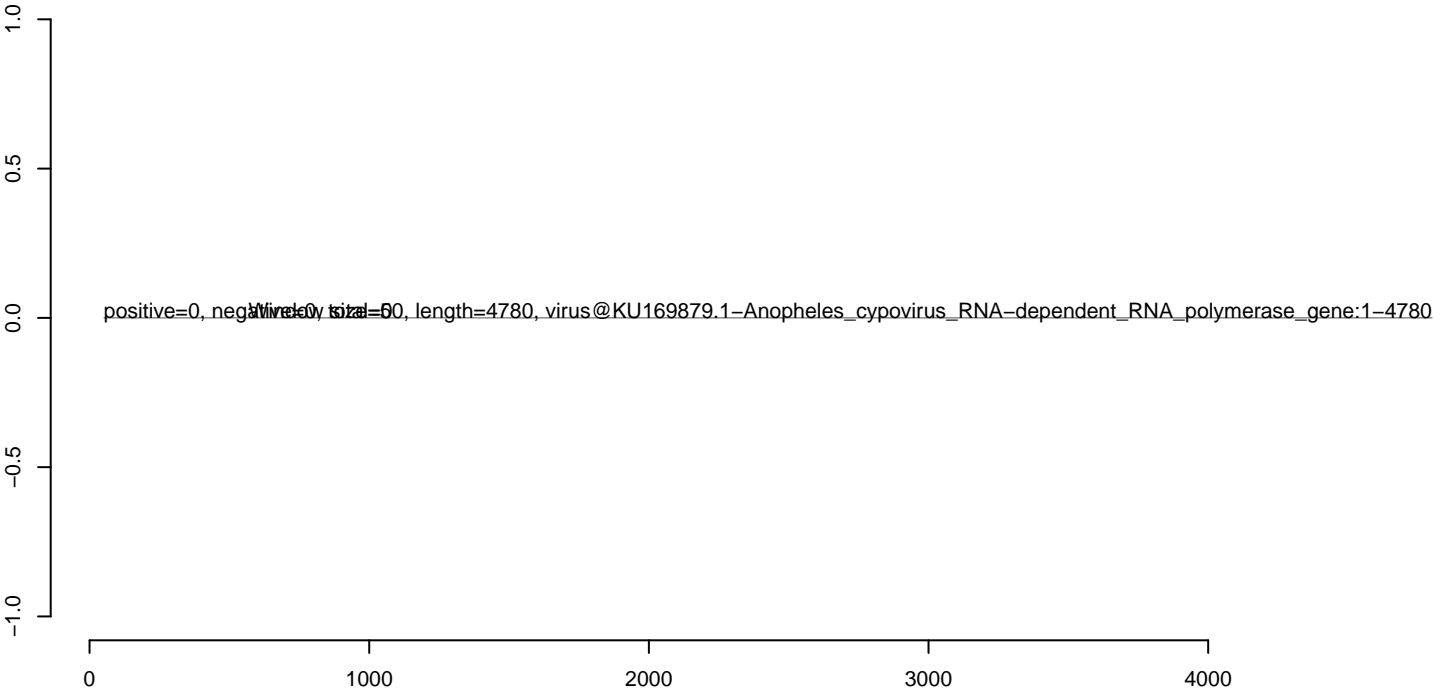
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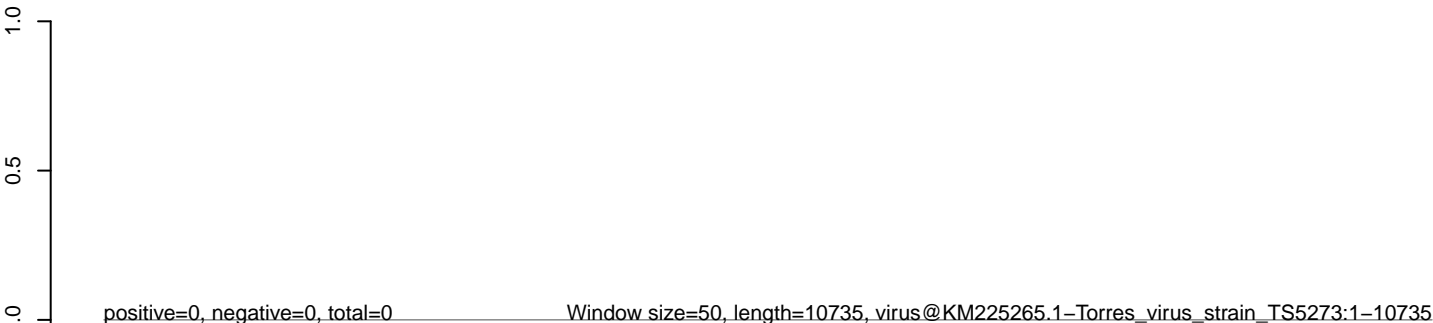
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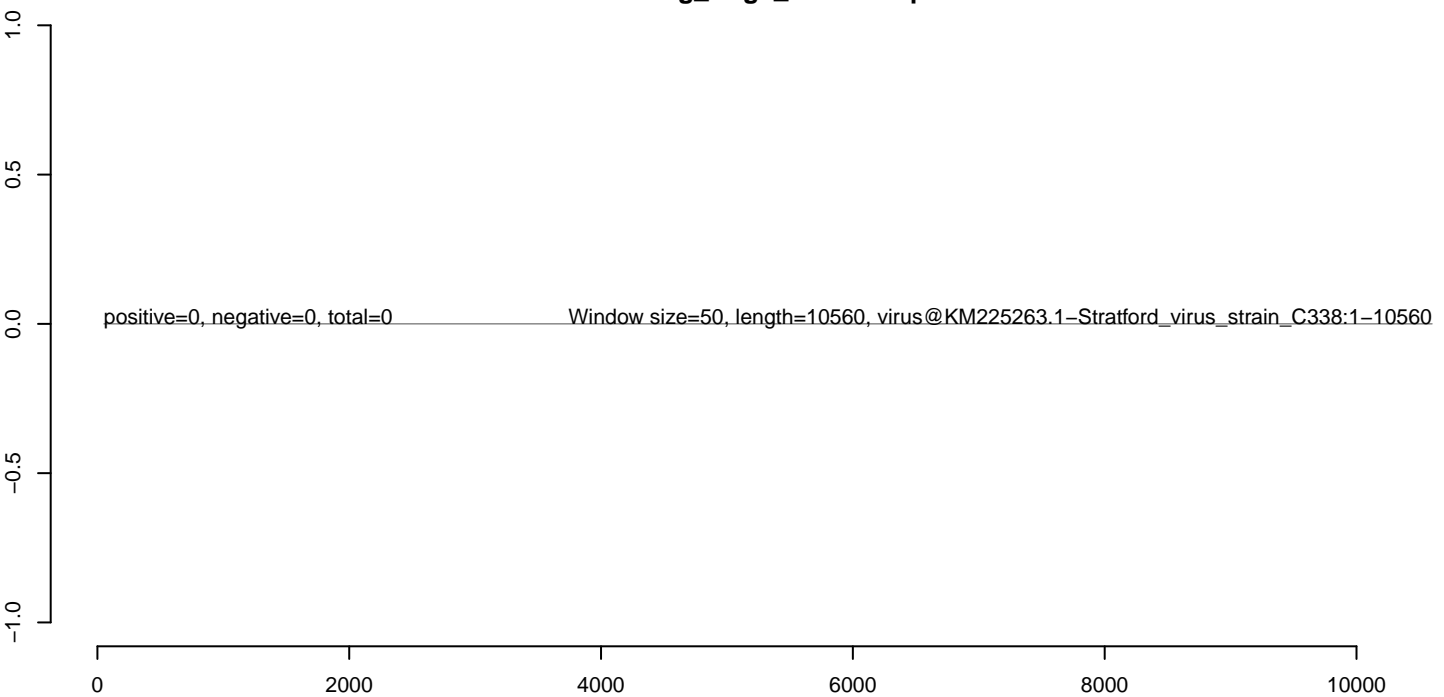
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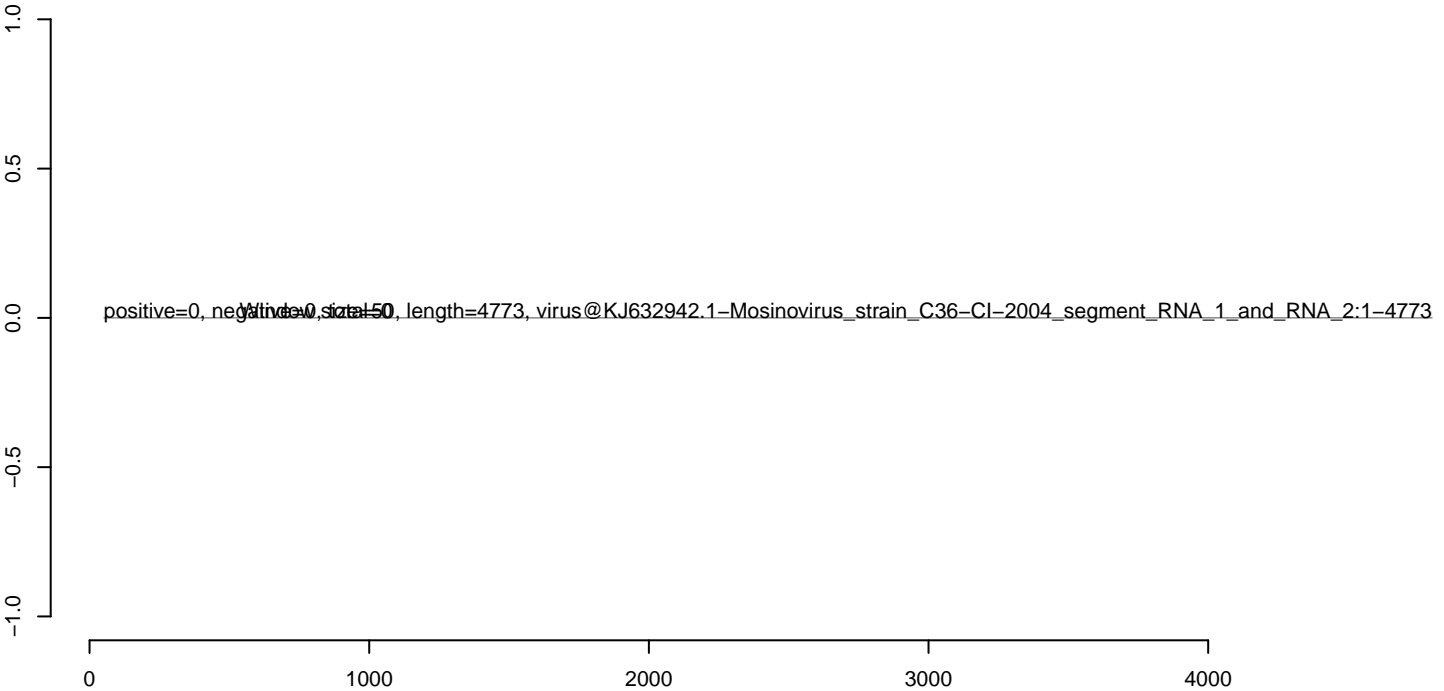
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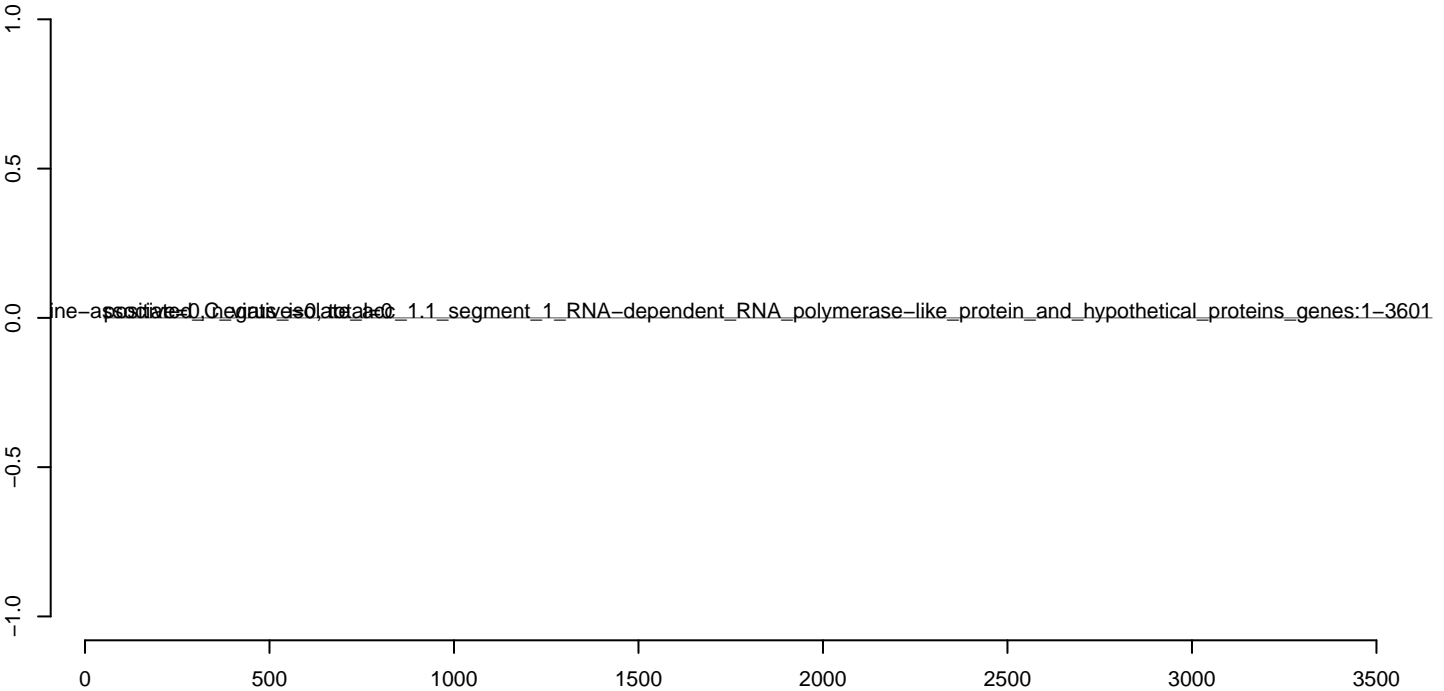
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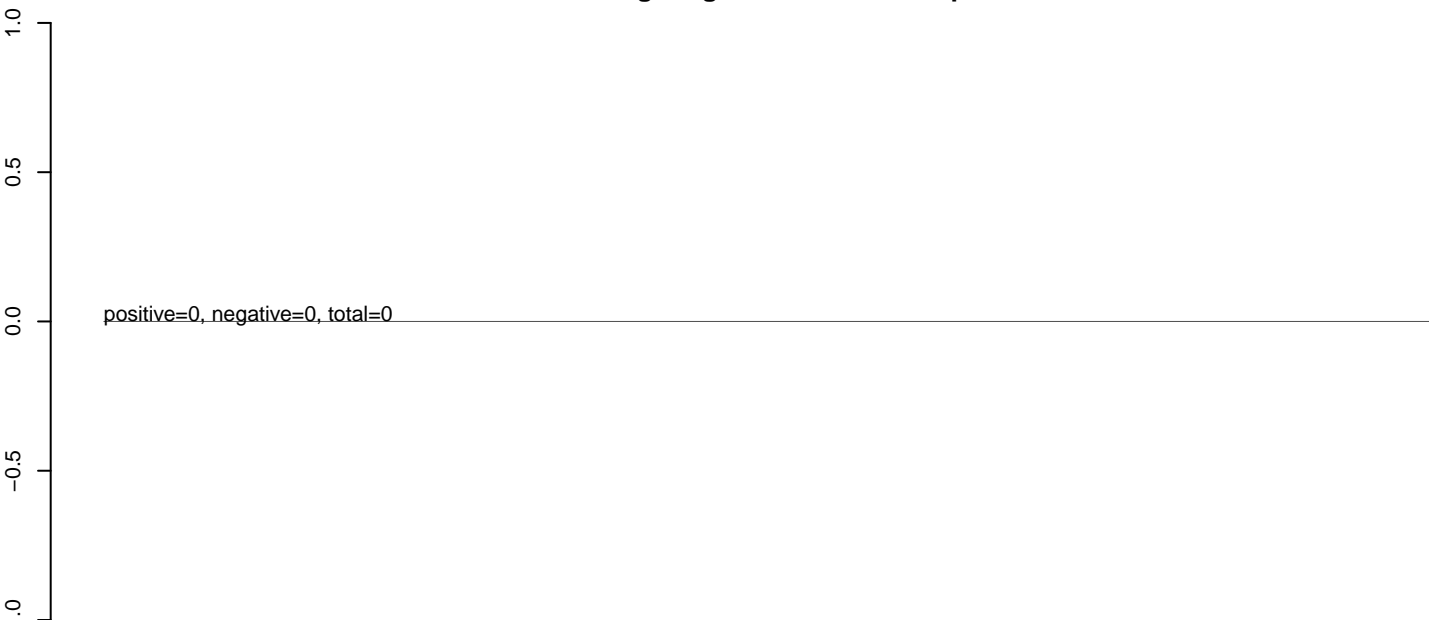
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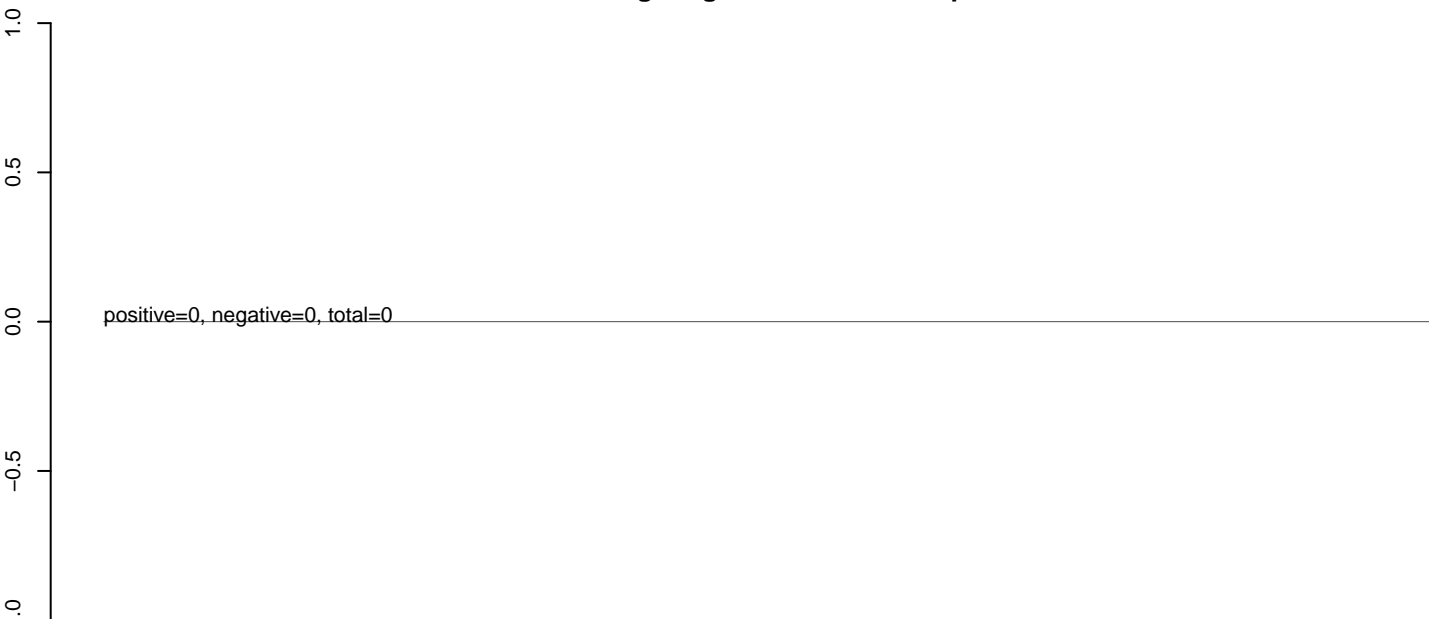
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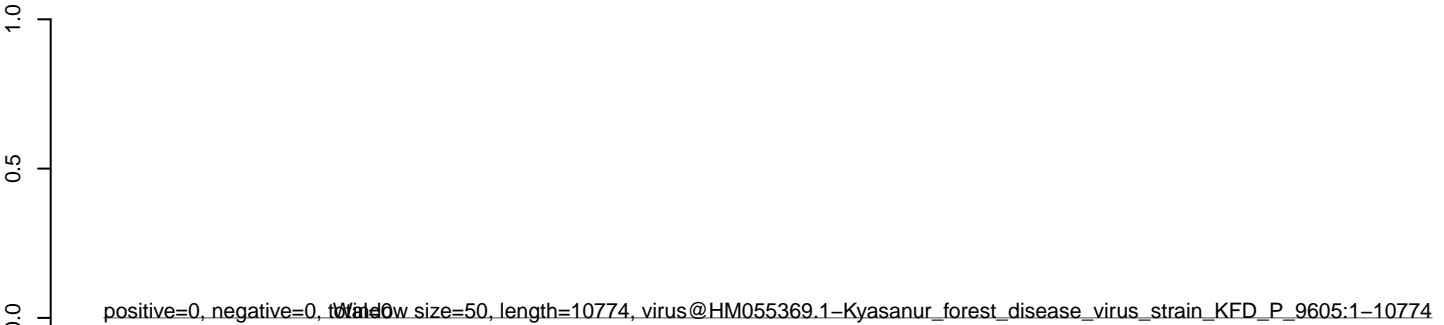
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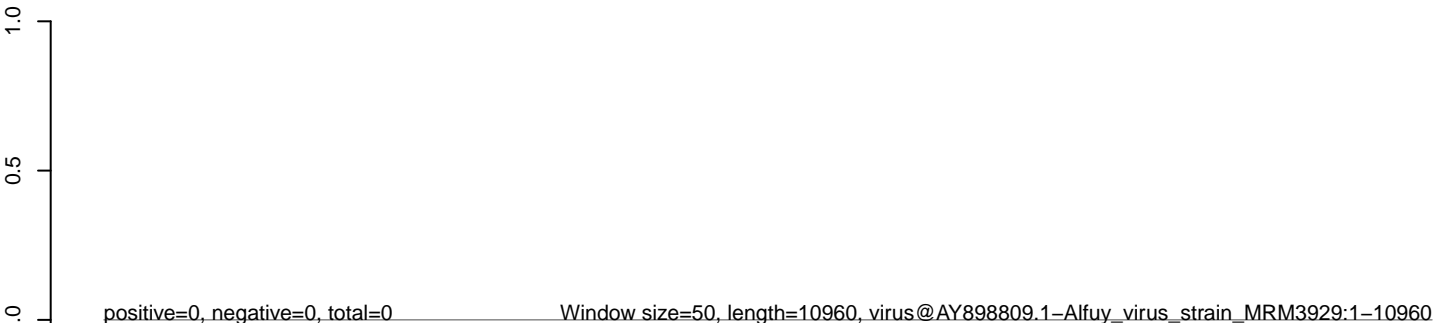
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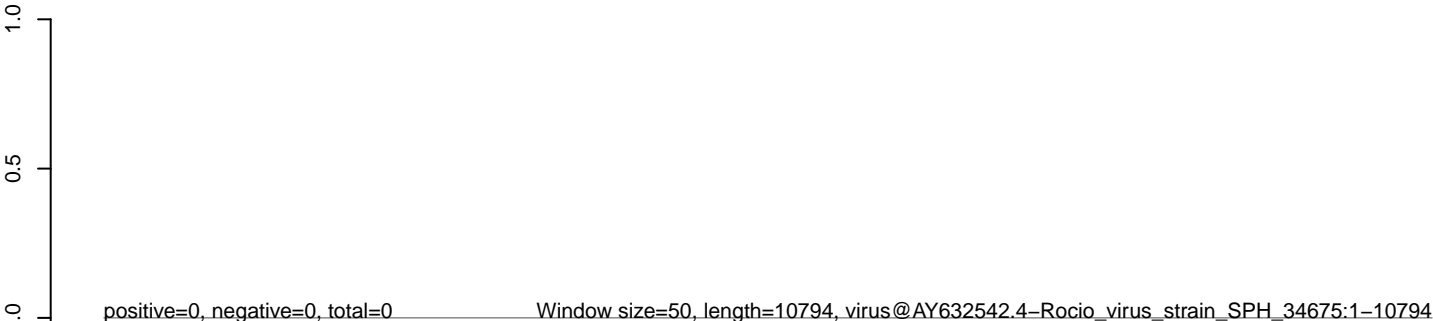
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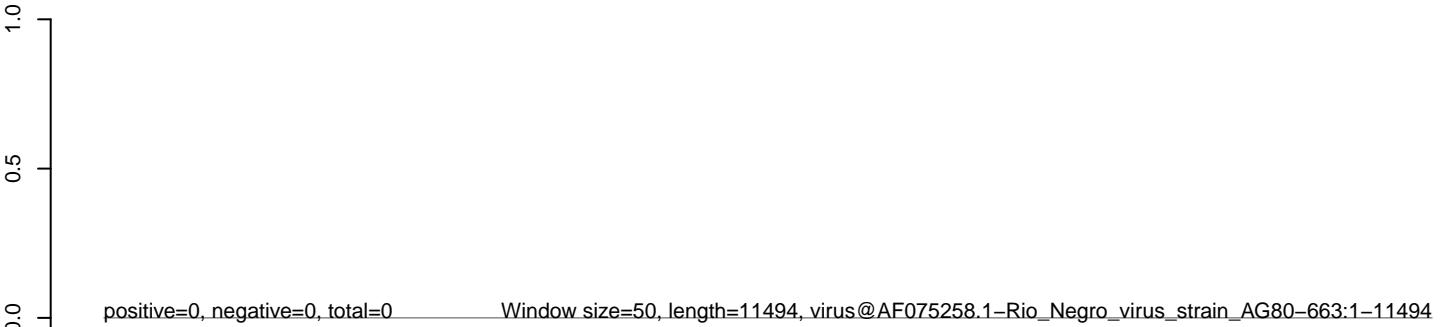
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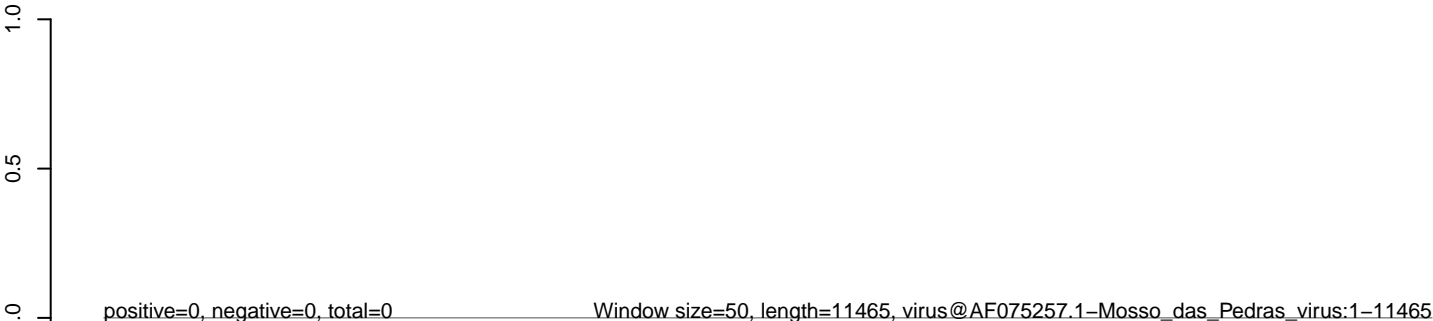
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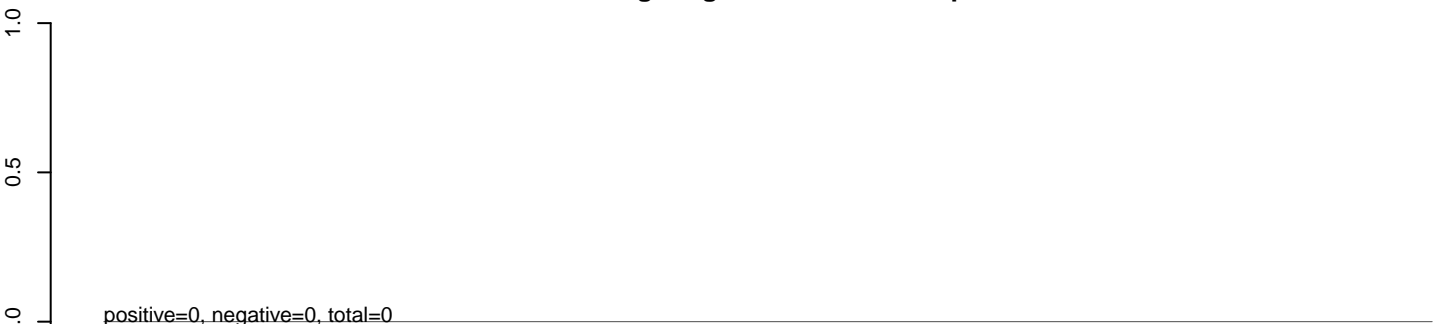


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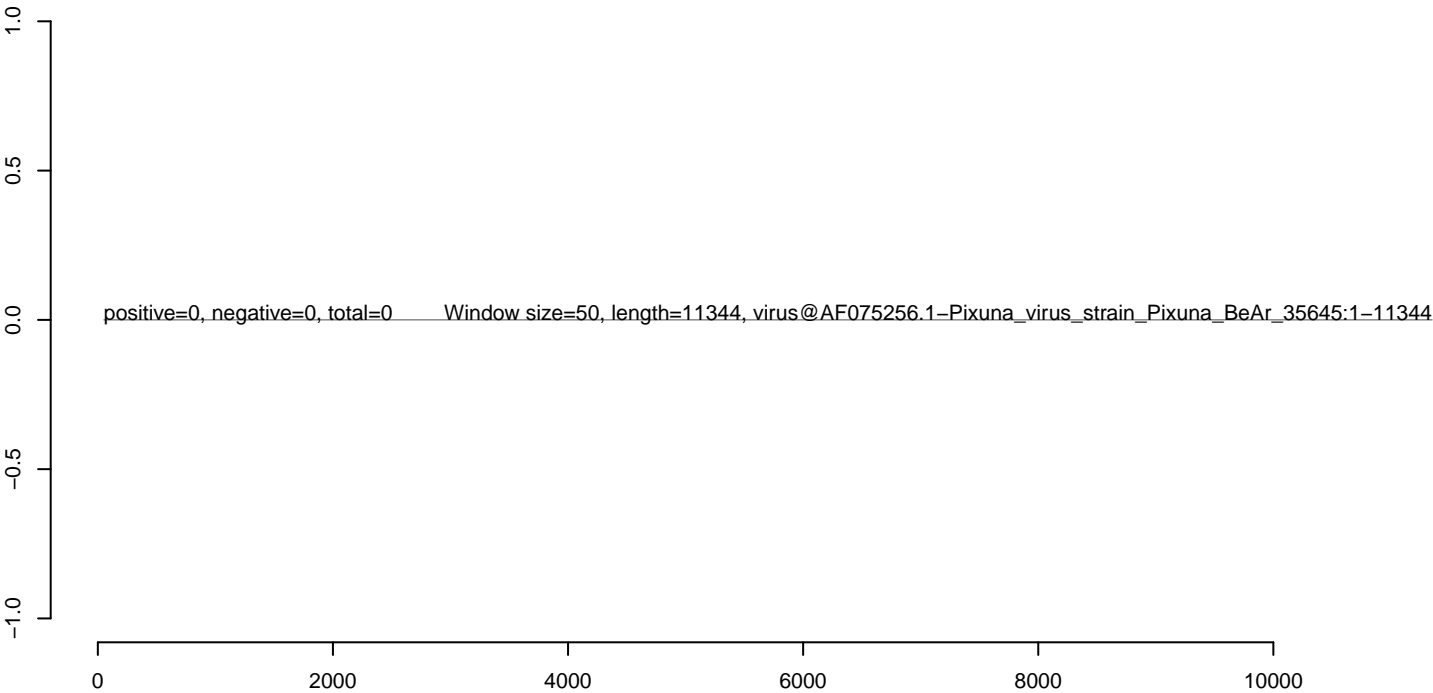
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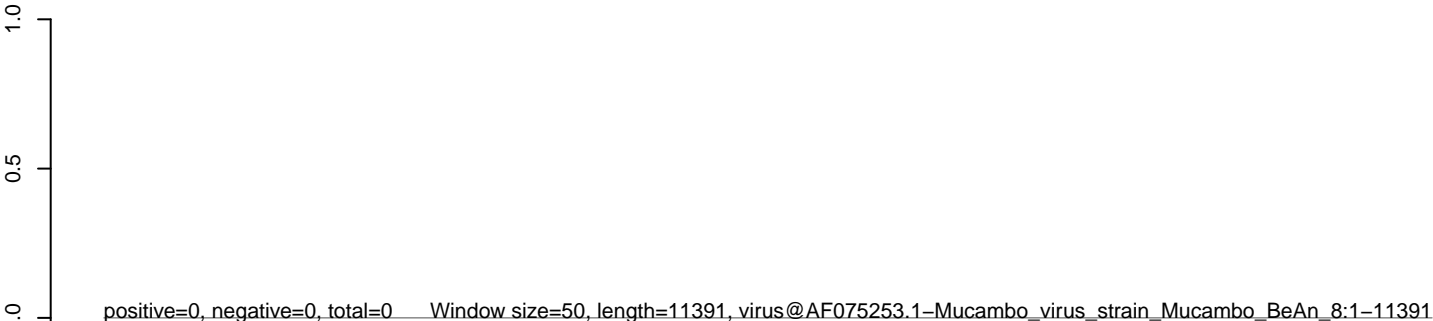
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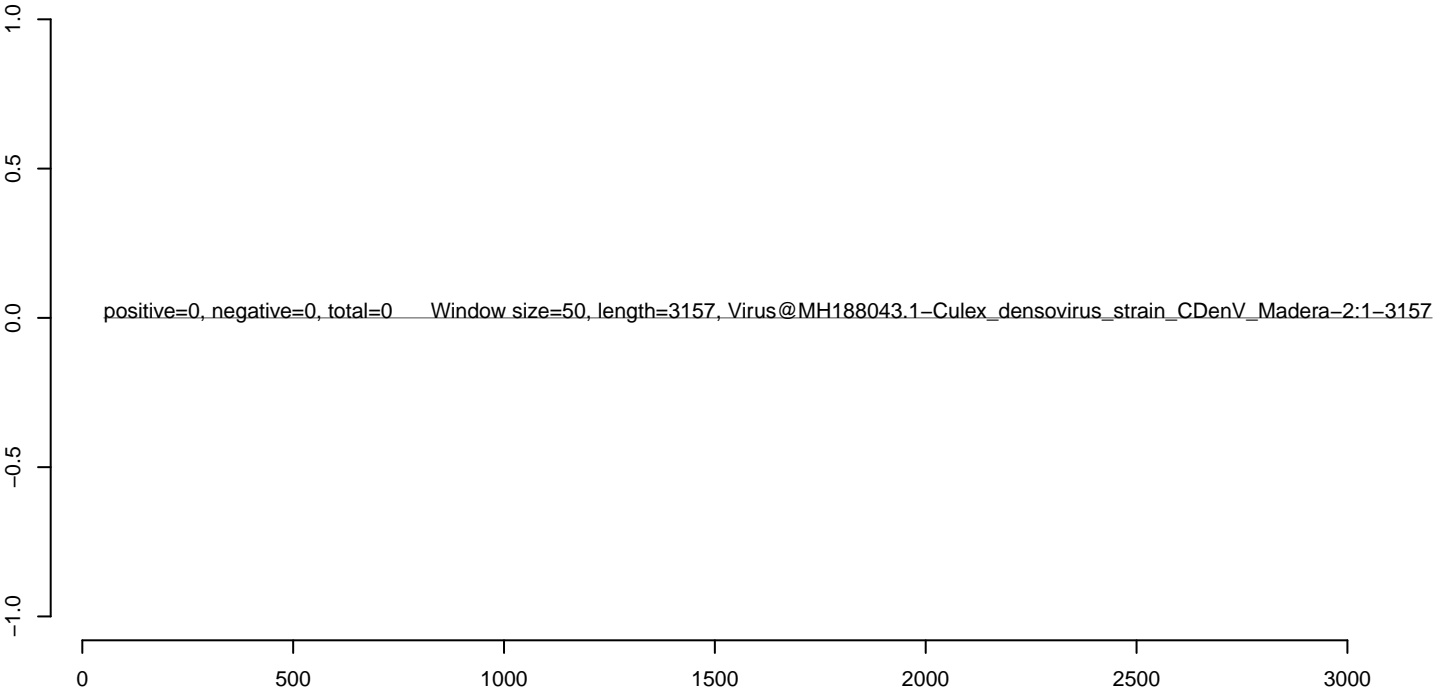
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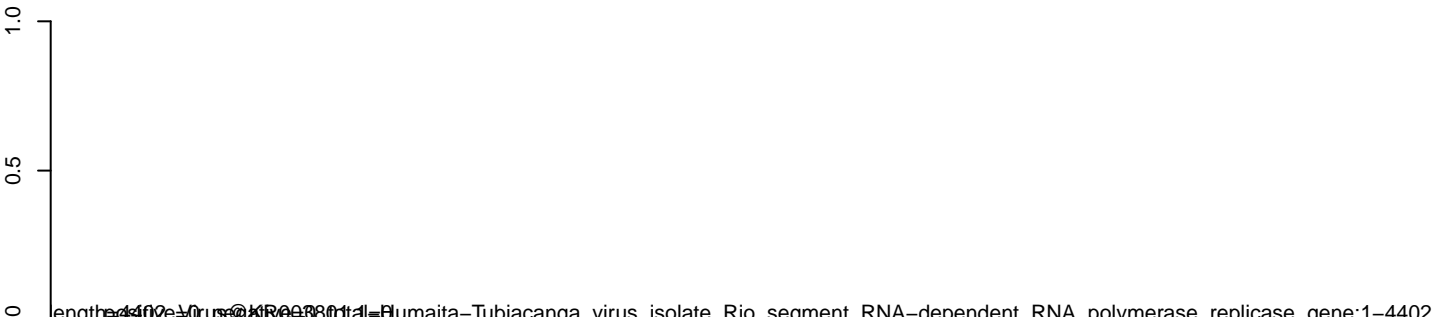
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