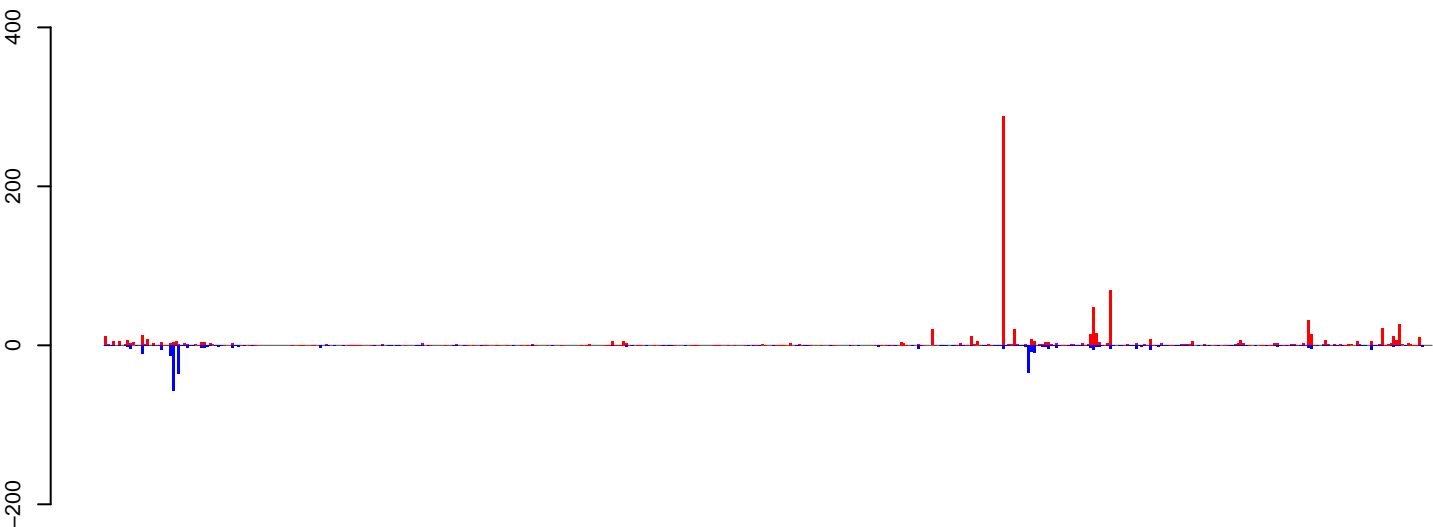
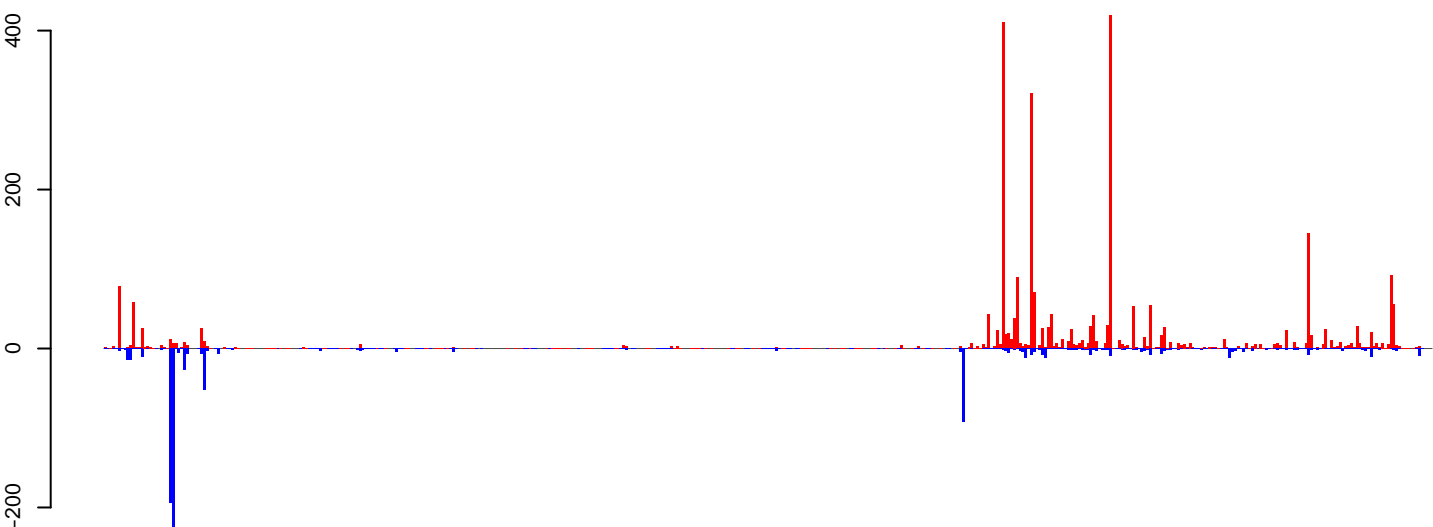


AeAlbo_C636_6dpi_SINV1.18_23.rep



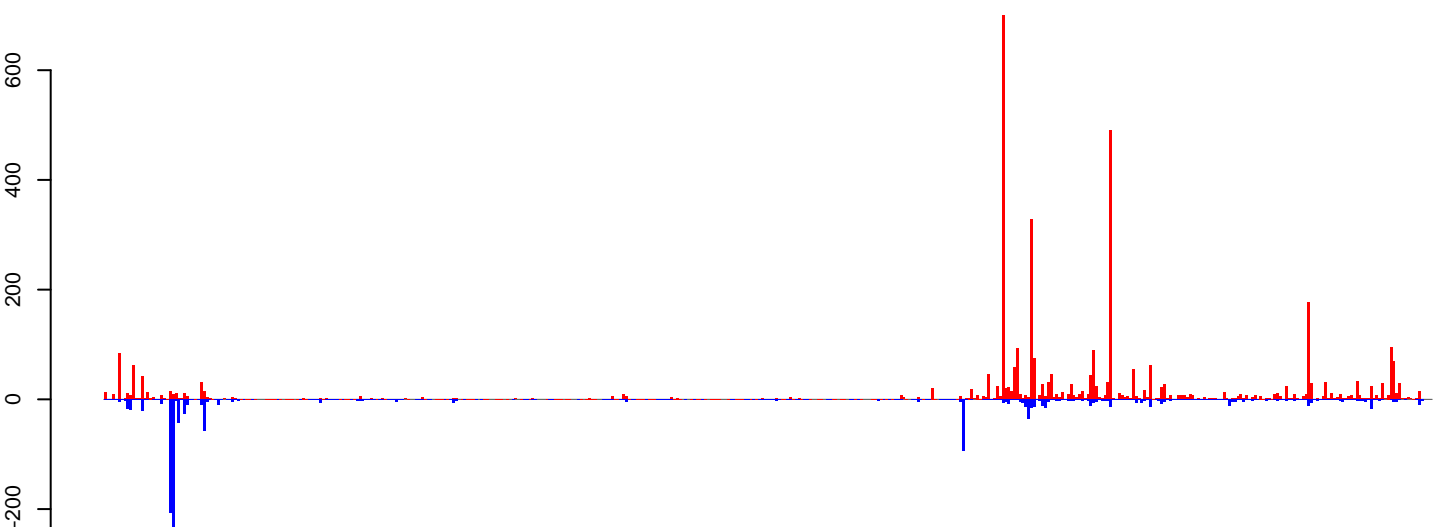
positive=955, negative=336, total=1291

AeAlbo_C636_6dpi_SINV1.24_35.rep



positive=3037, negative=1105, total=4142

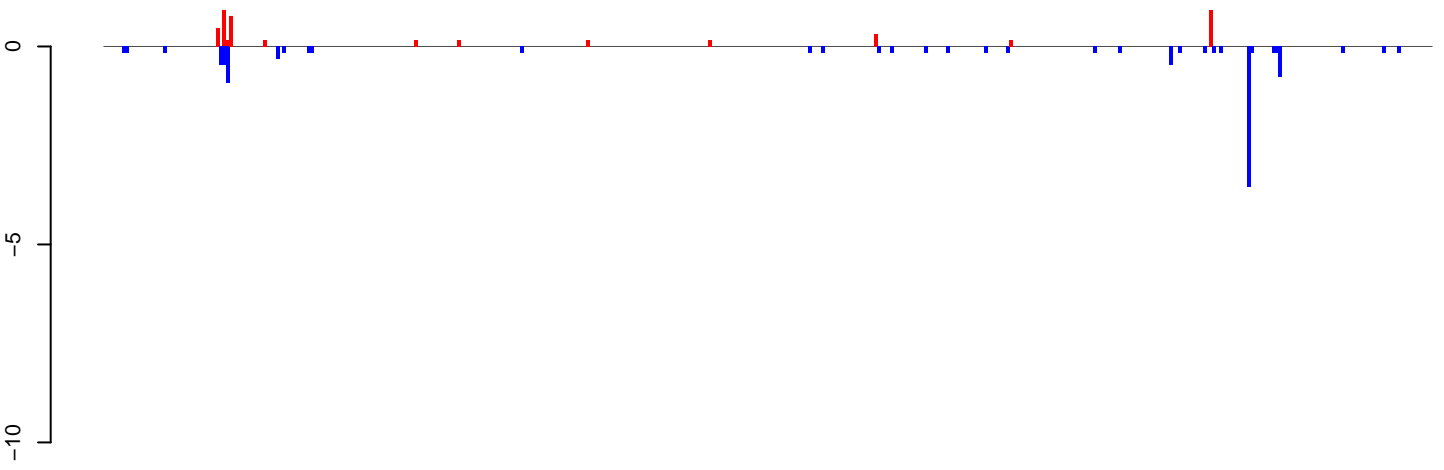
AeAlbo_C636_6dpi_SINV1.rep



positive=4075, negative=1471, total=5546

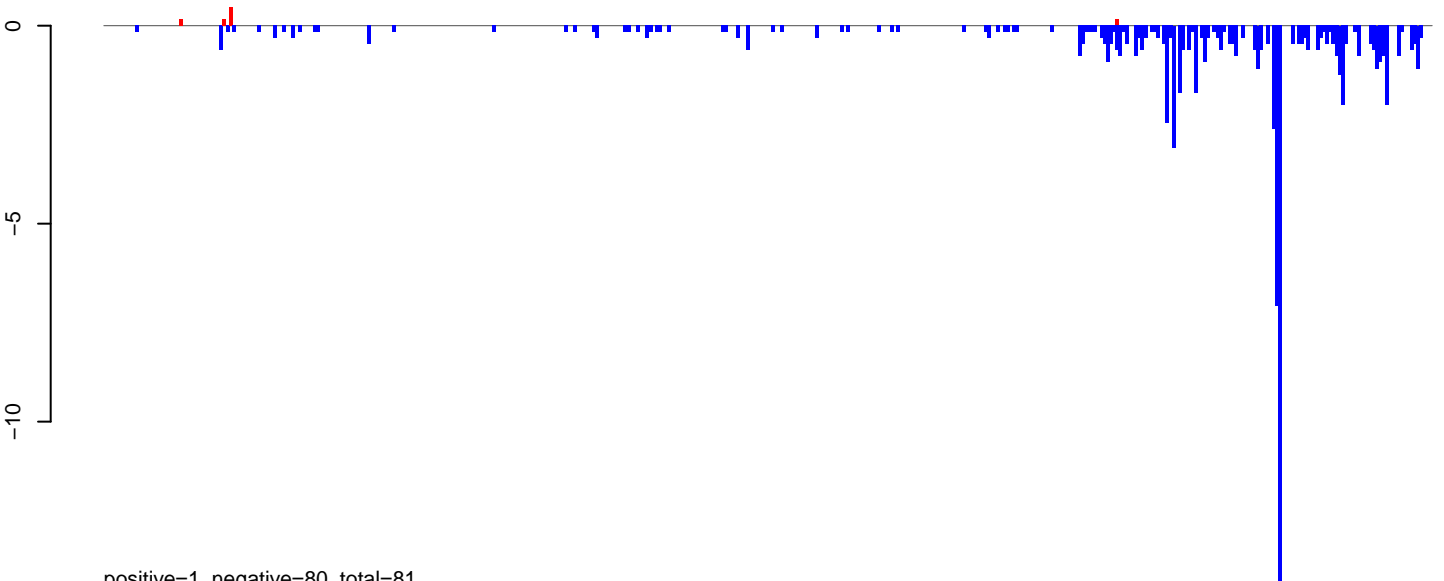
Window size=25, length=11703, virus@NC_001547.1-Sindbis_virus:1-11703

AeAlbo_C636_6dpi_SINV1.18_23.rep



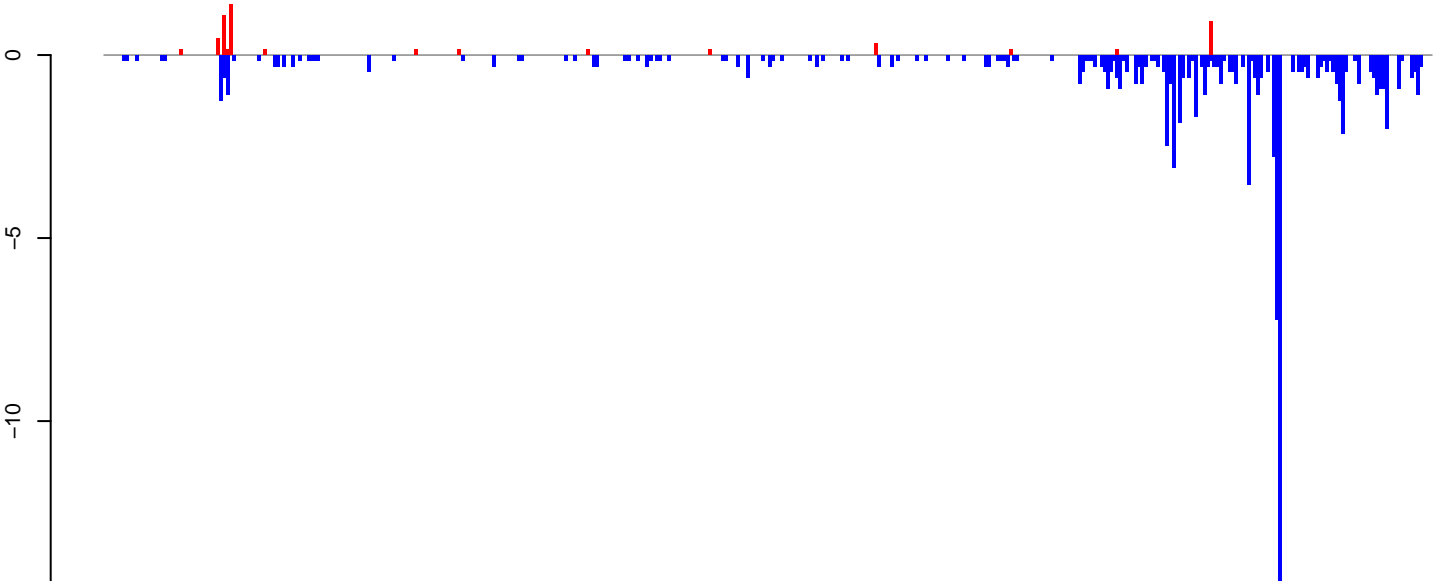
positive=4, negative=11, total=16

AeAlbo_C636_6dpi_SINV1.24_35.rep



positive=1, negative=80, total=81

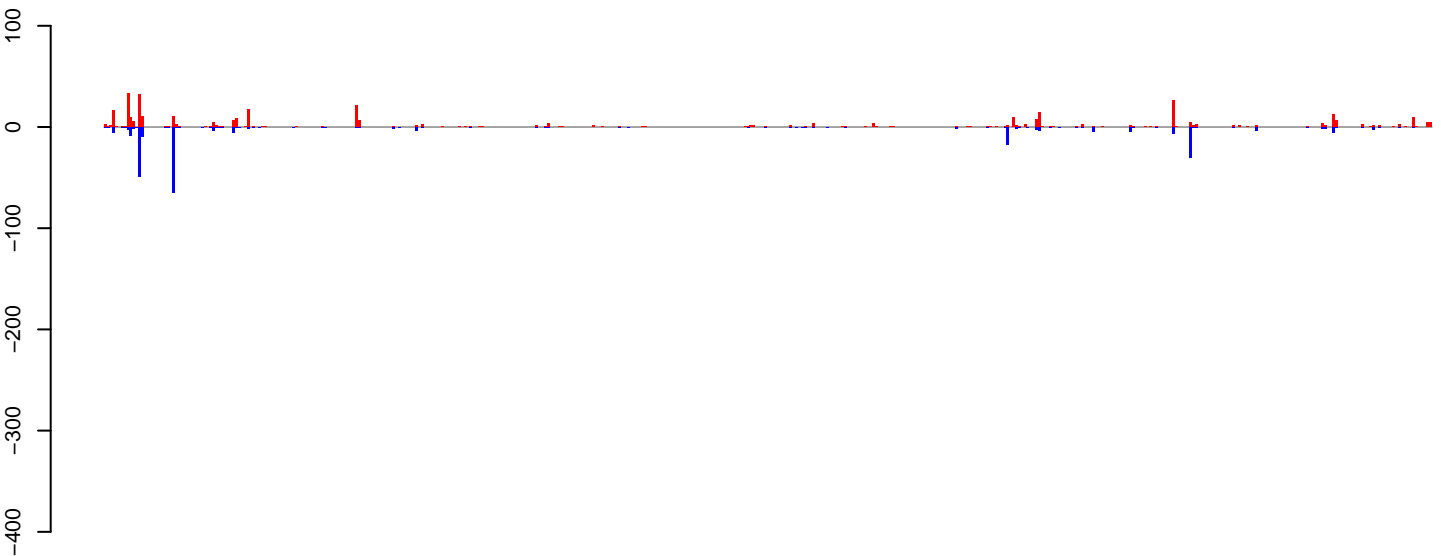
AeAlbo_C636_6dpi_SINV1.rep



size=25, length=10587, @AF499835.2-Aedes_albopictus_cell_fusing_agent_virus_polyprotein-like_protein_gene_and_agCP8252:1-10587

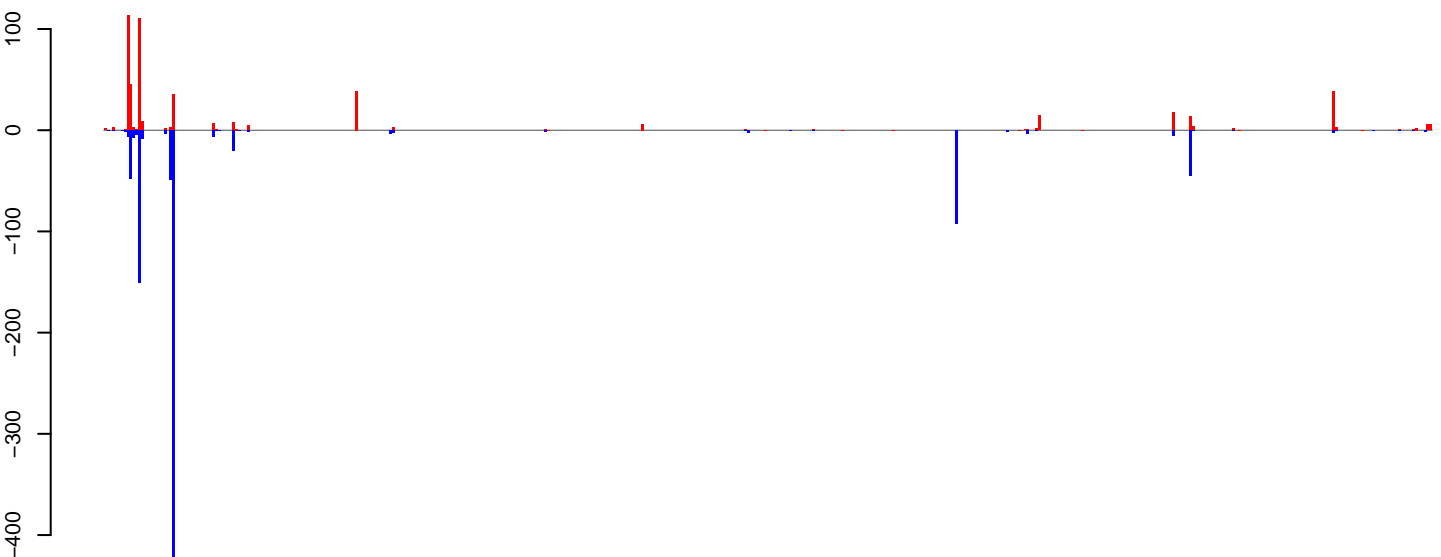
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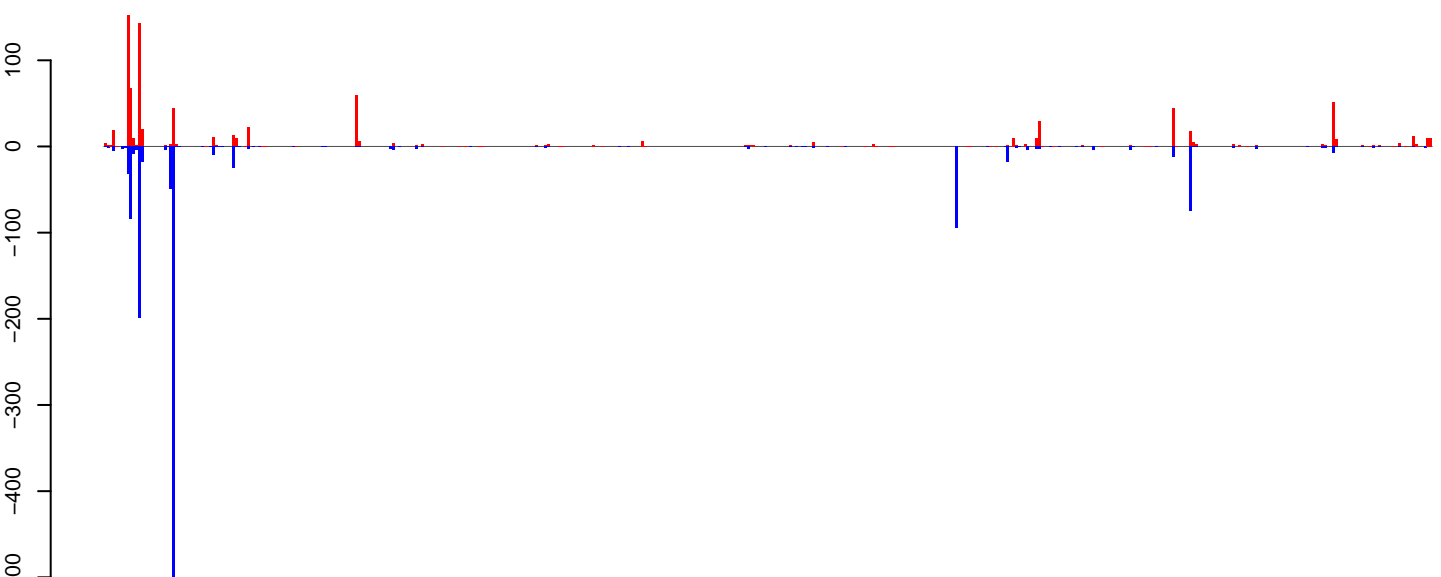
positive=363, negative=261, total=624

AeAlbo_C636_6dpi_SINV1.24_35.rep



positive=513, negative=949, total=1462

AeAlbo_C636_6dpi_SINV1.rep

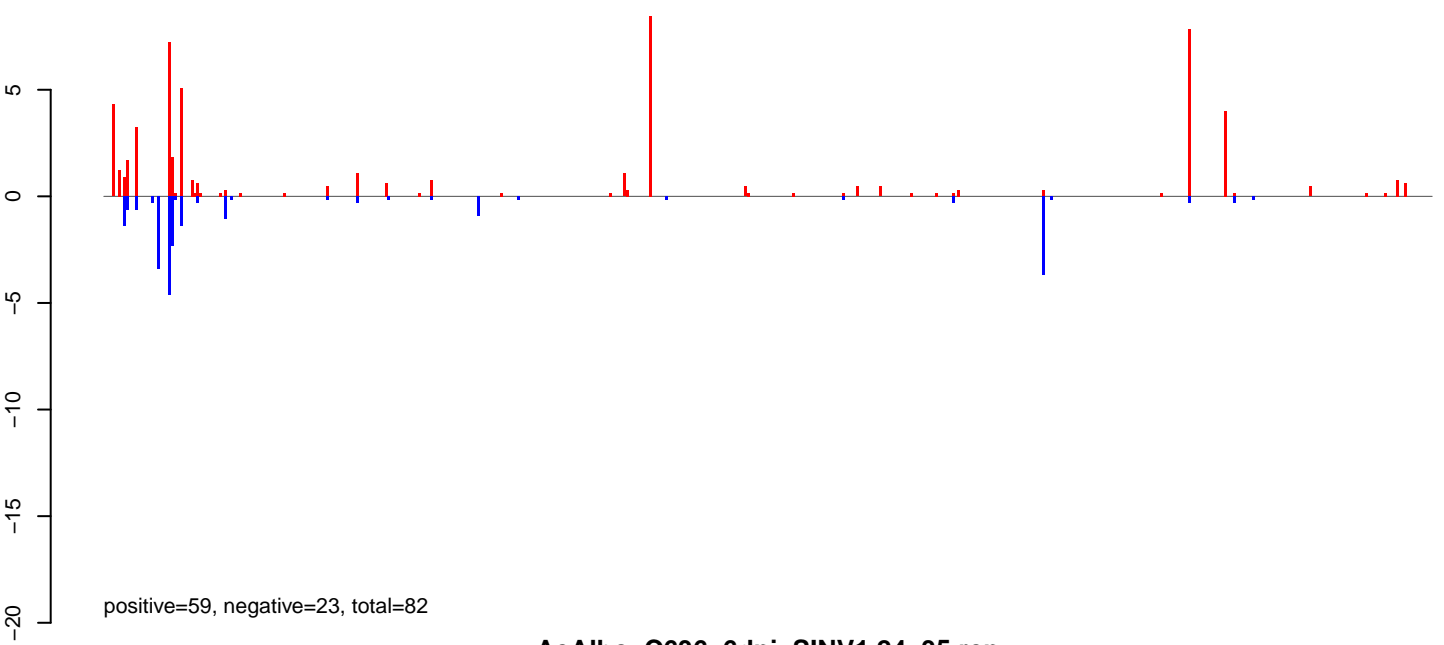


positive=899, negative=1264, total=2163

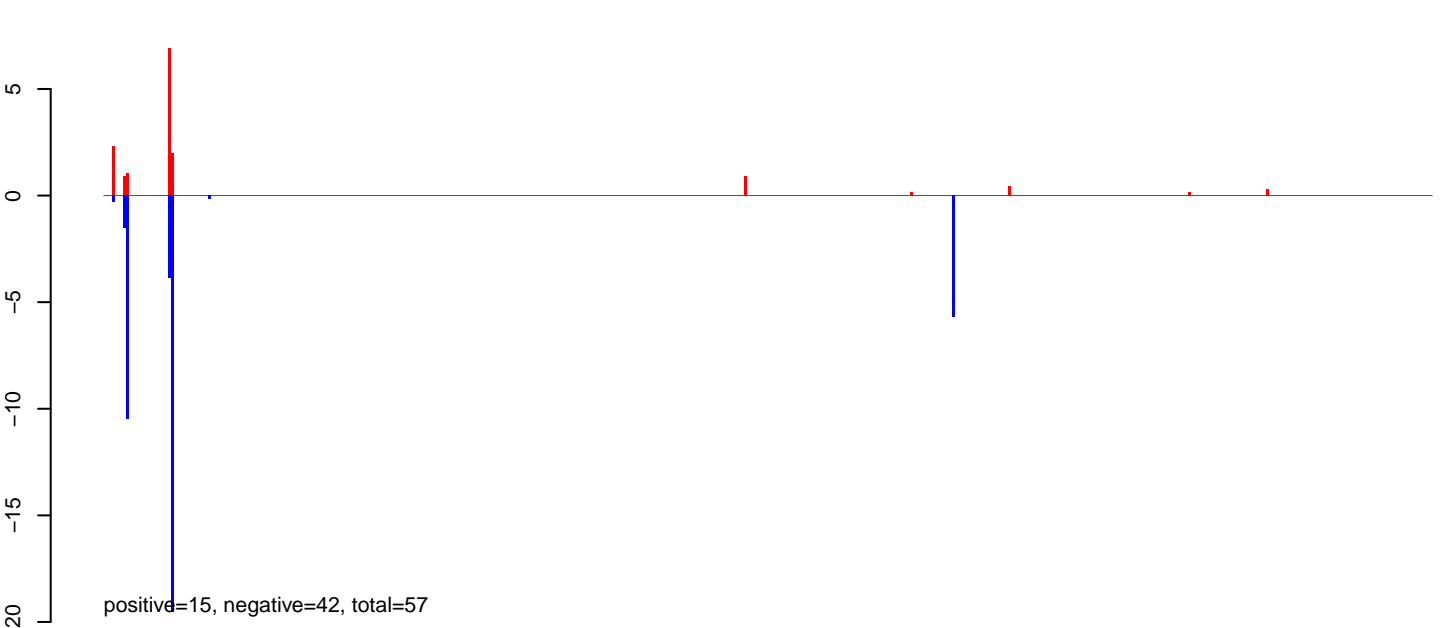
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0 2000 4000 6000 8000 10000 12000

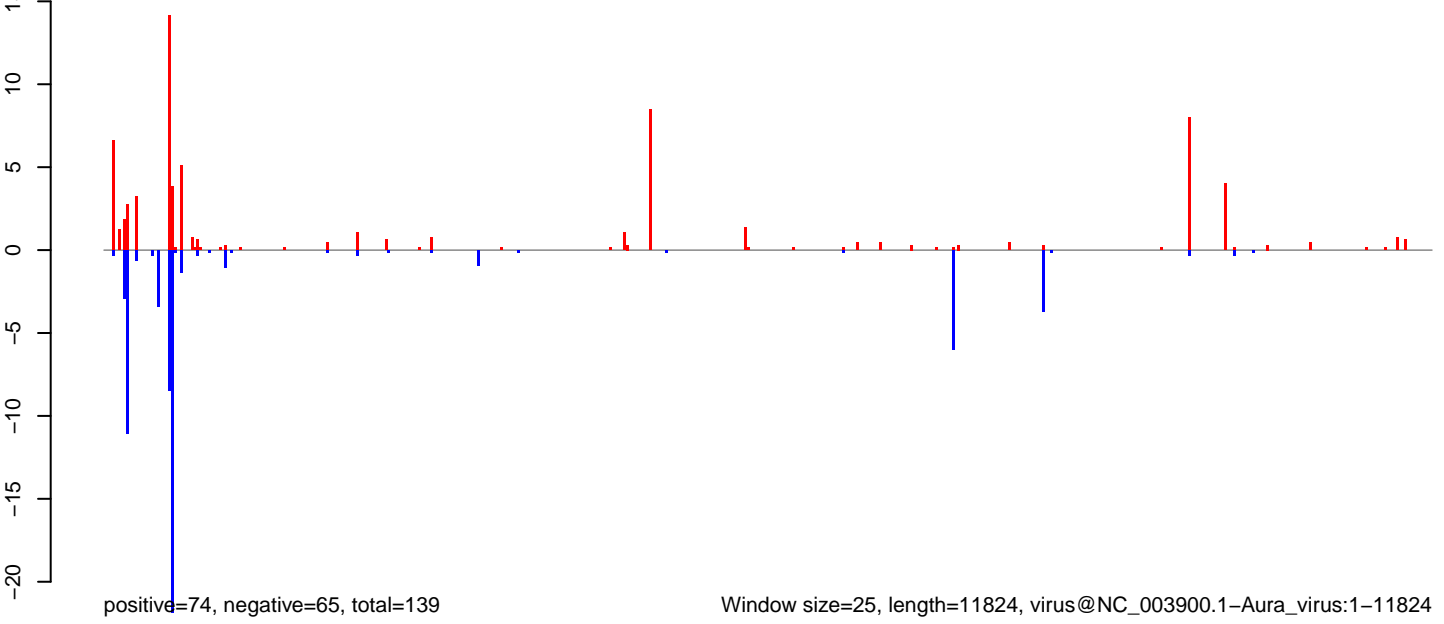
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



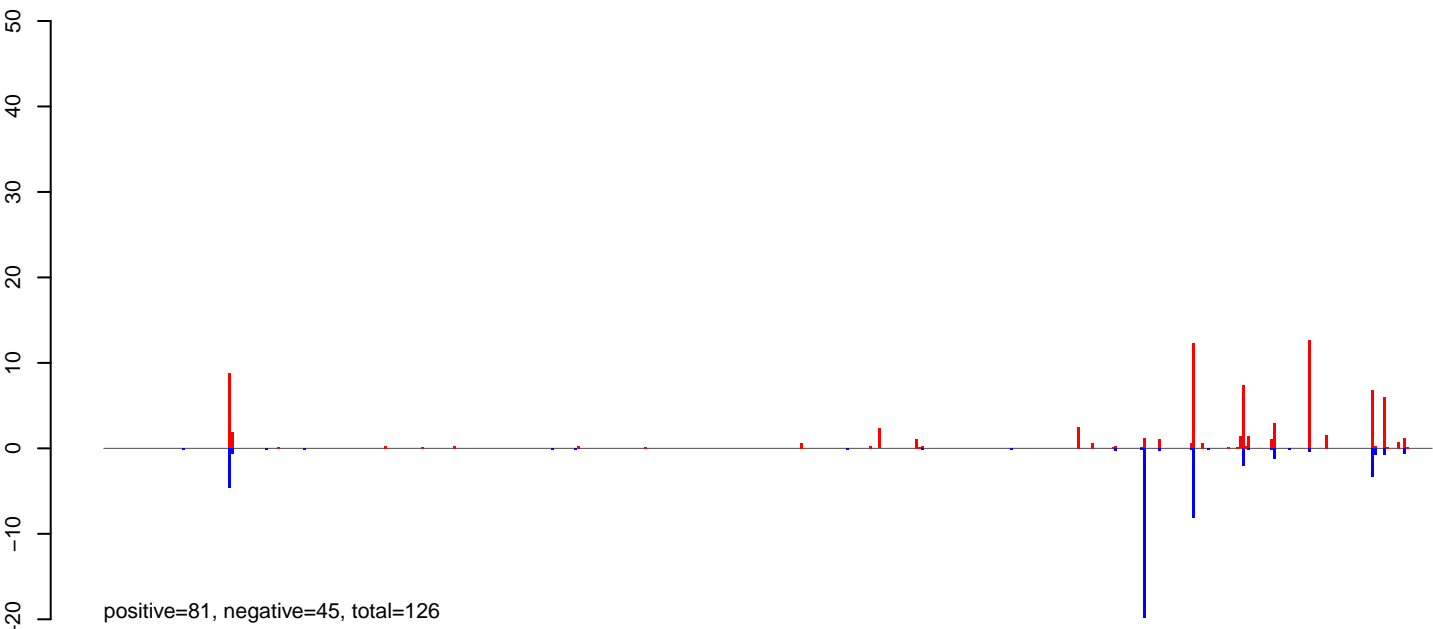
AeAlbo_C636_6dpi_SINV1.rep



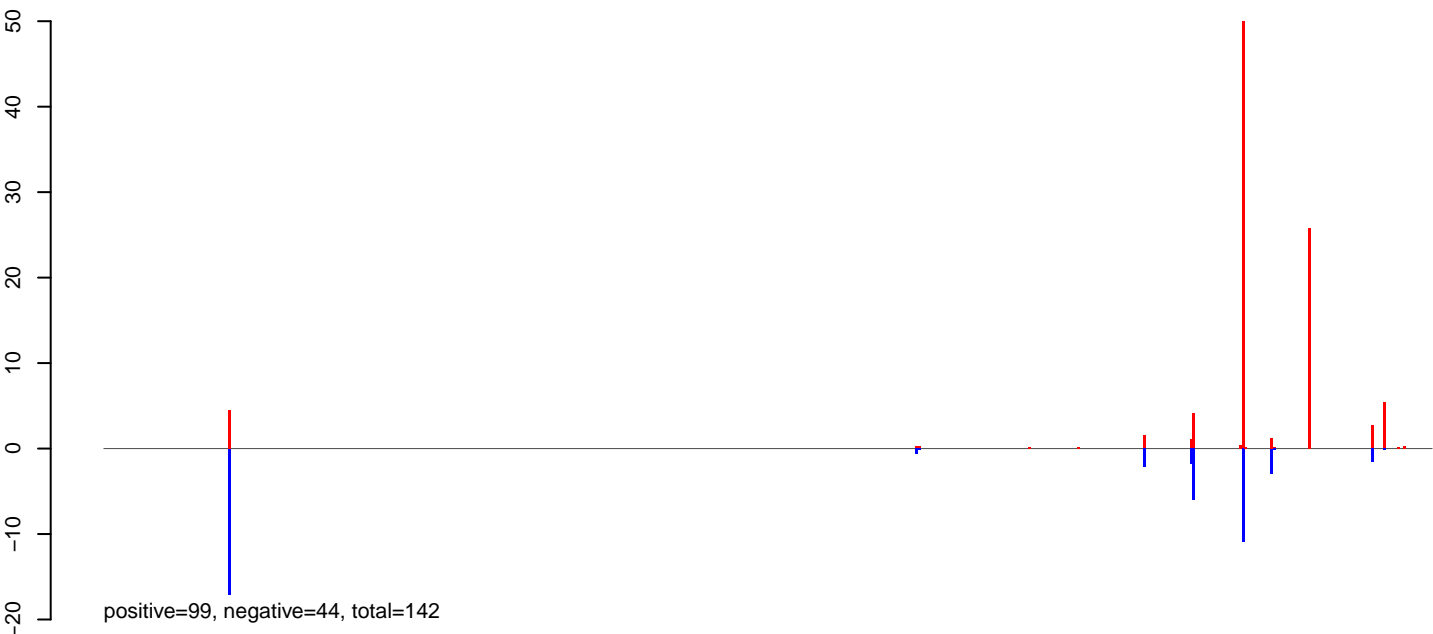
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0 2000 4000 6000 8000 10000 12000

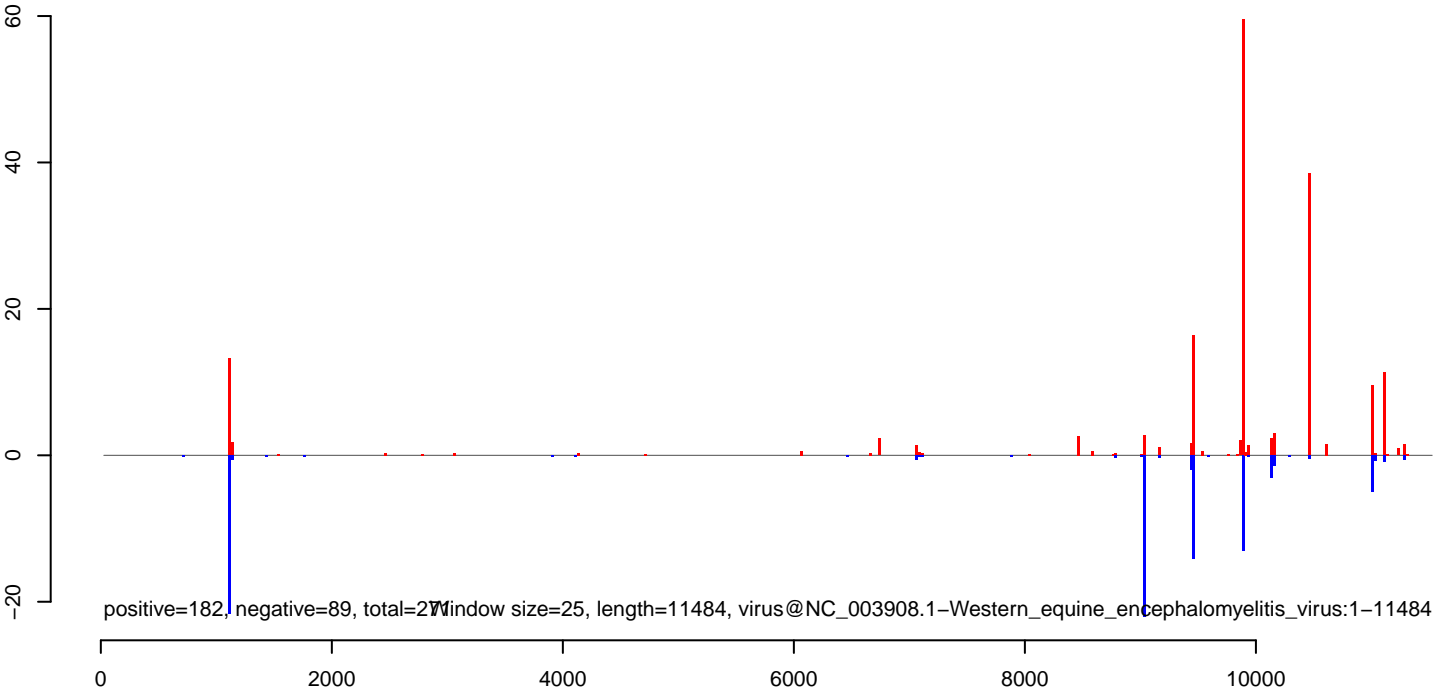
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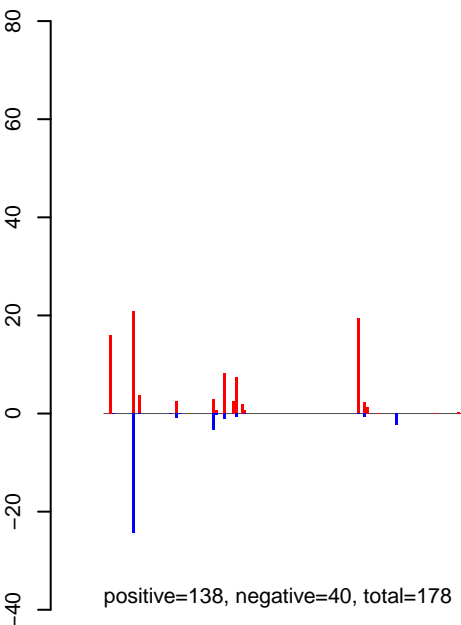
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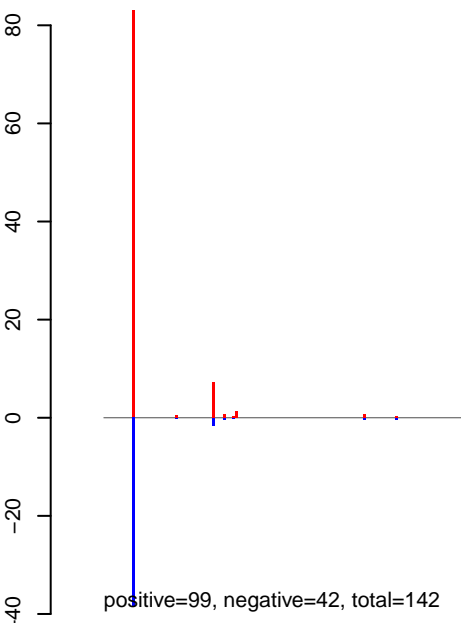
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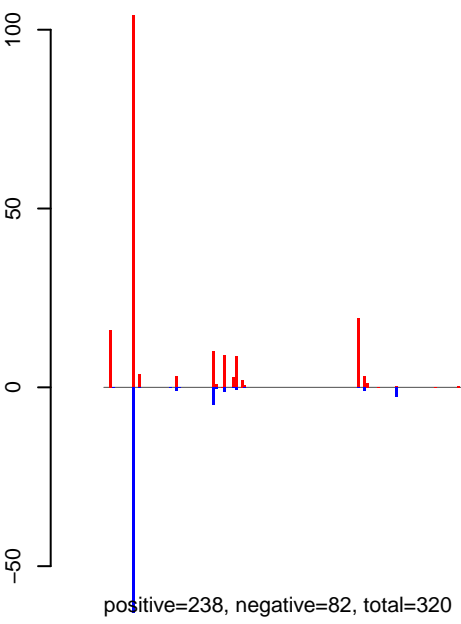
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



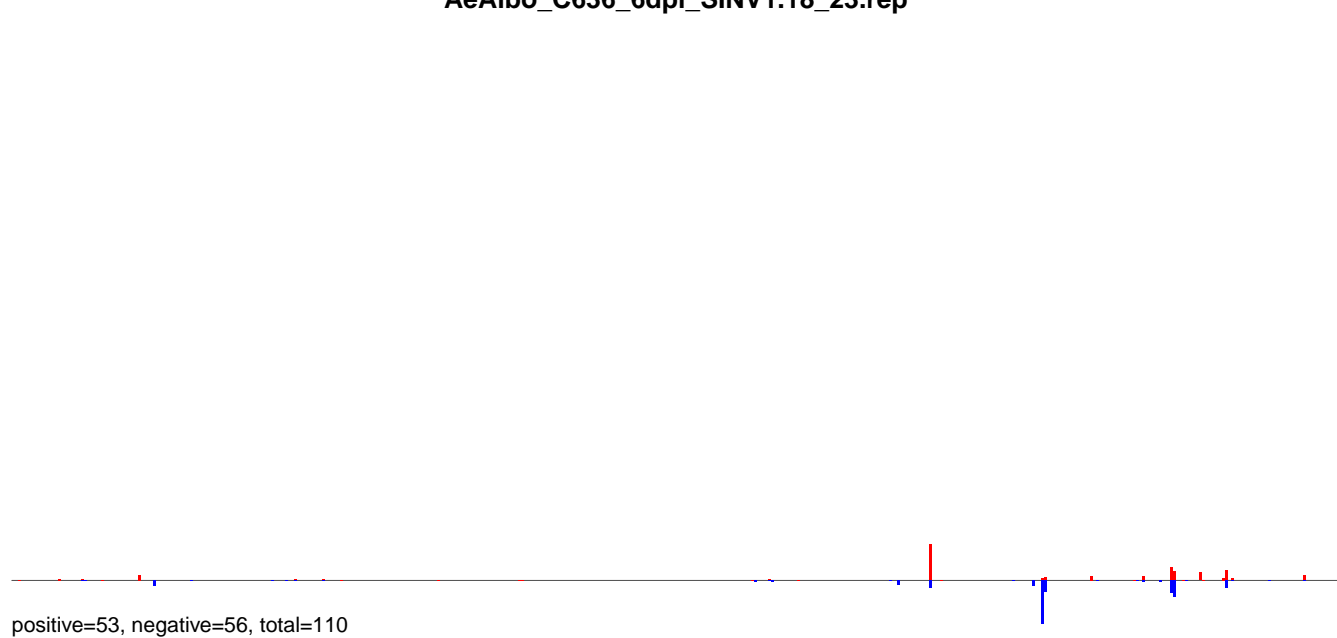
AeAlbo_C636_6dpi_SINV1.rep



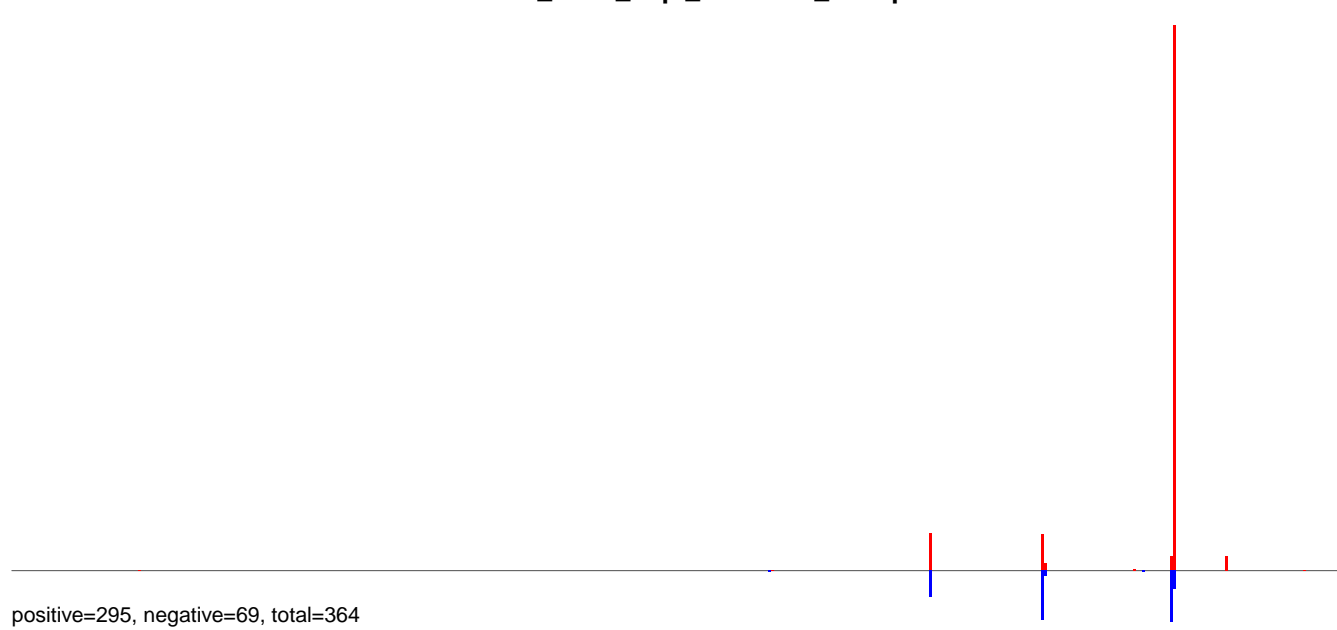
Window size=25, length=11634, virus@NC_018615.1-Eilat_virus_isolate_EO329:1-11634

0 2000 4000 6000 8000 10000 12000

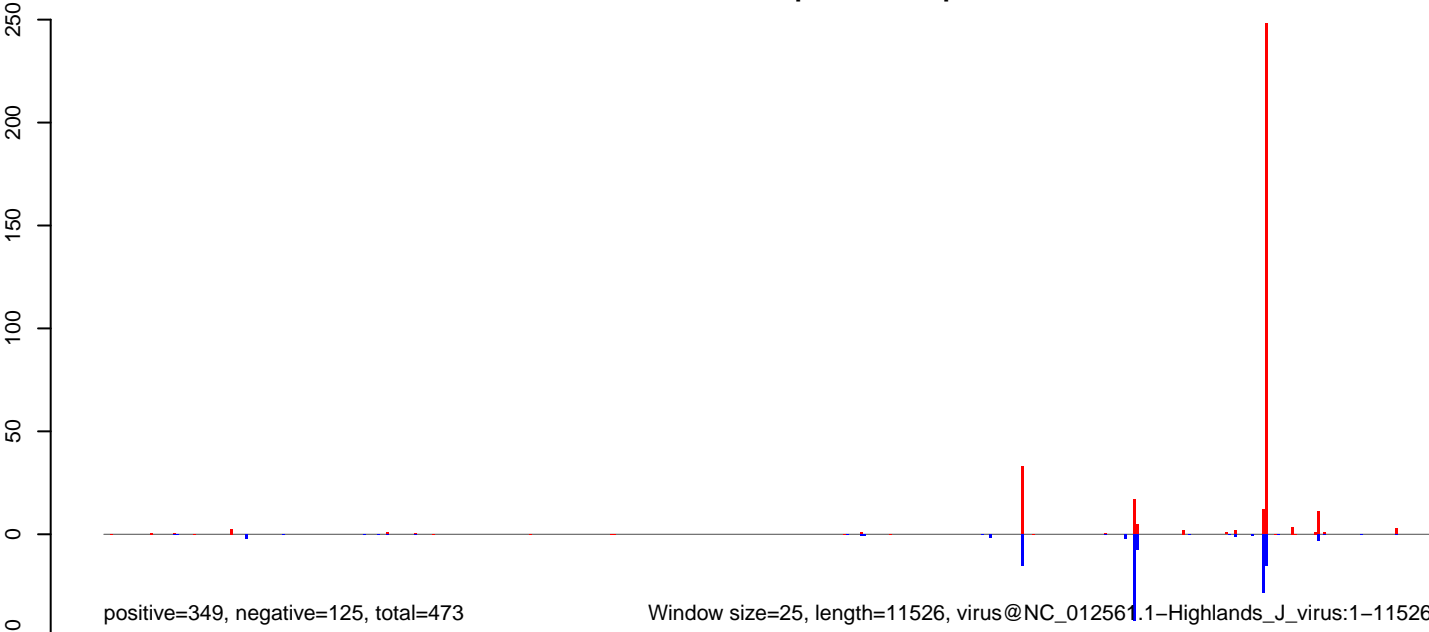
AeAlbo_C636_6dpi_SINV1.18_23.rep



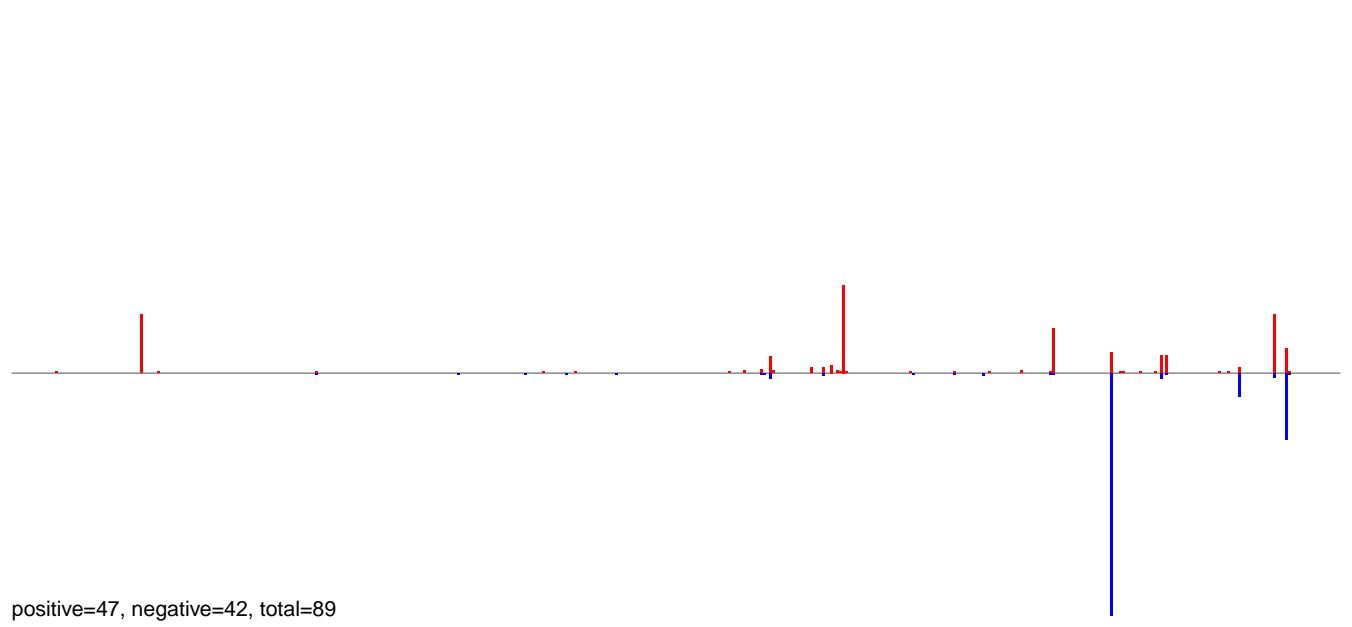
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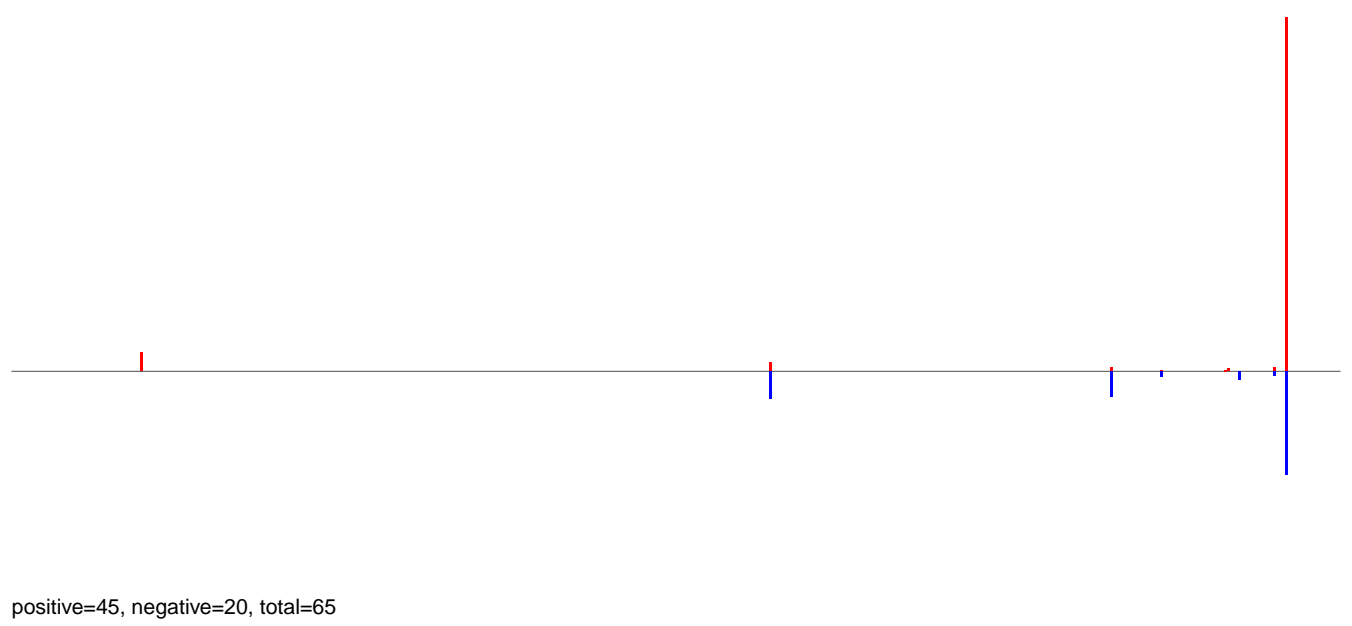
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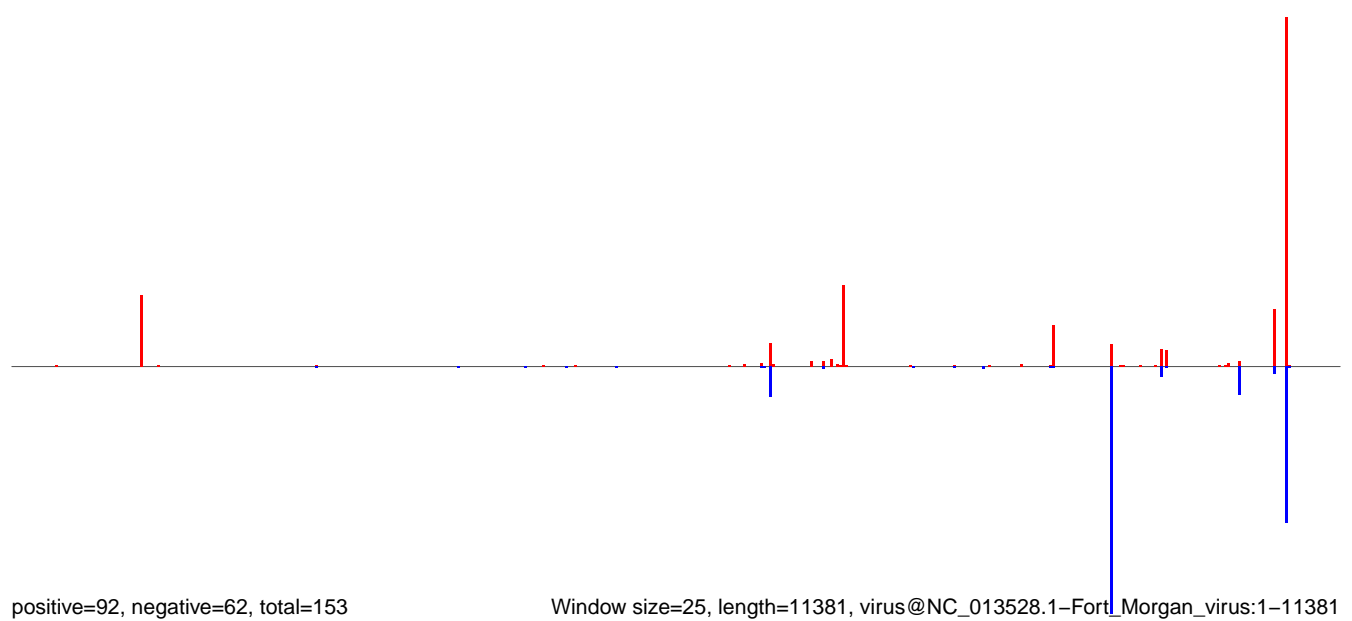
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AeAlbo_C636_6dpi_SINV1.24_35.rep



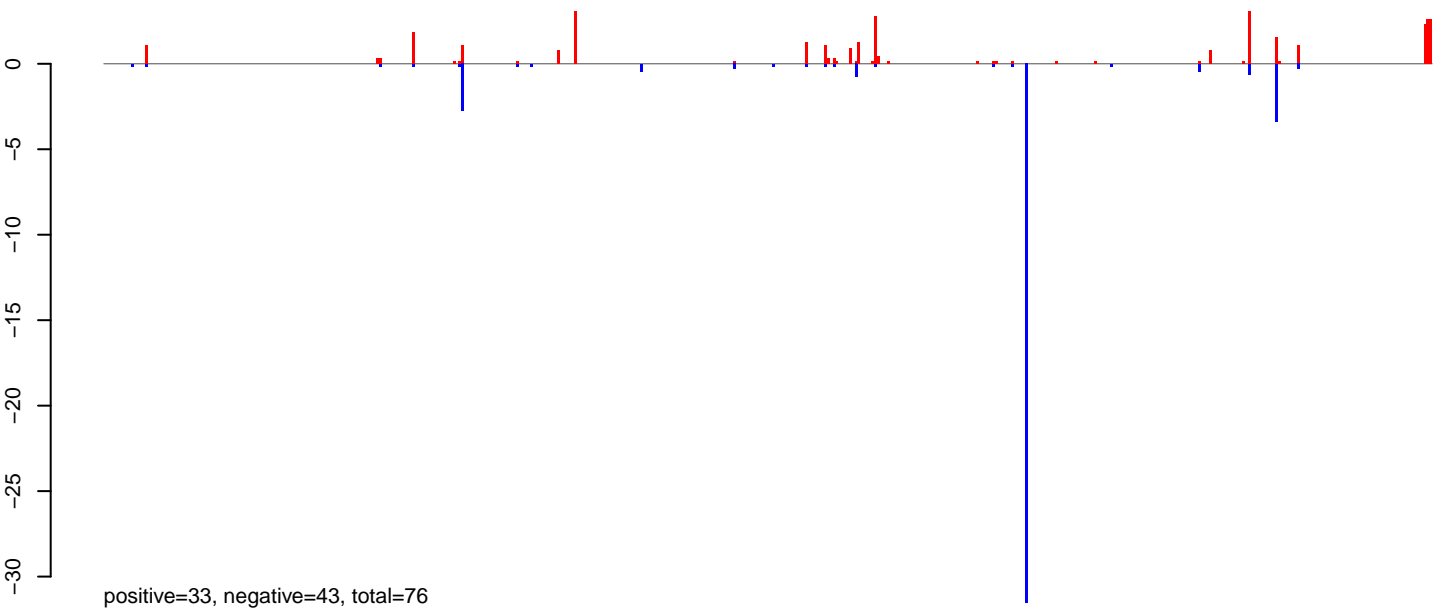
AeAlbo_C636_6dpi_SINV1.rep



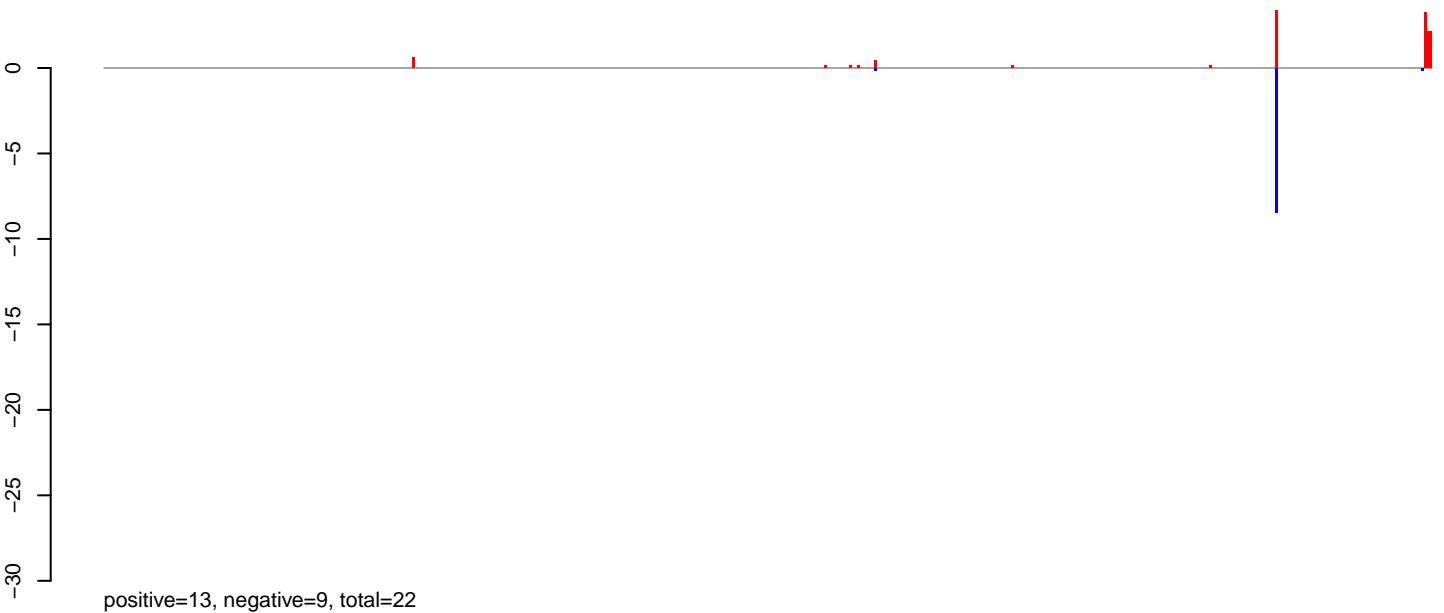
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0 2000 4000 6000 8000 10000

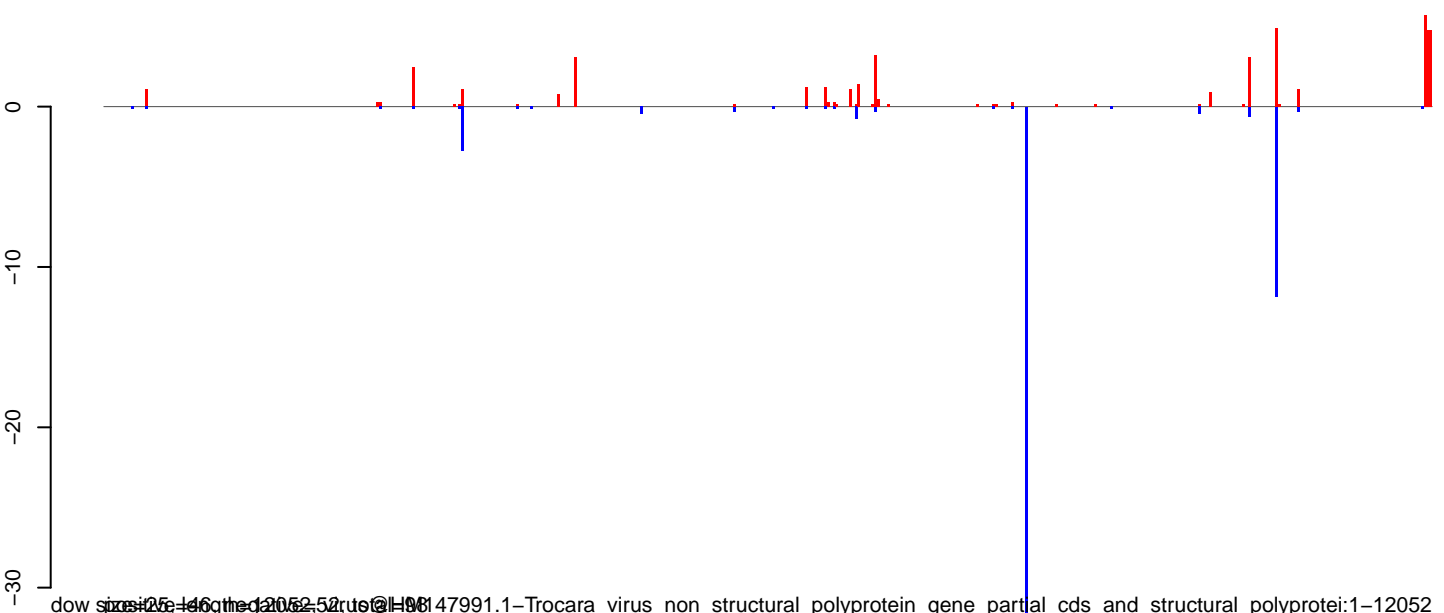
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



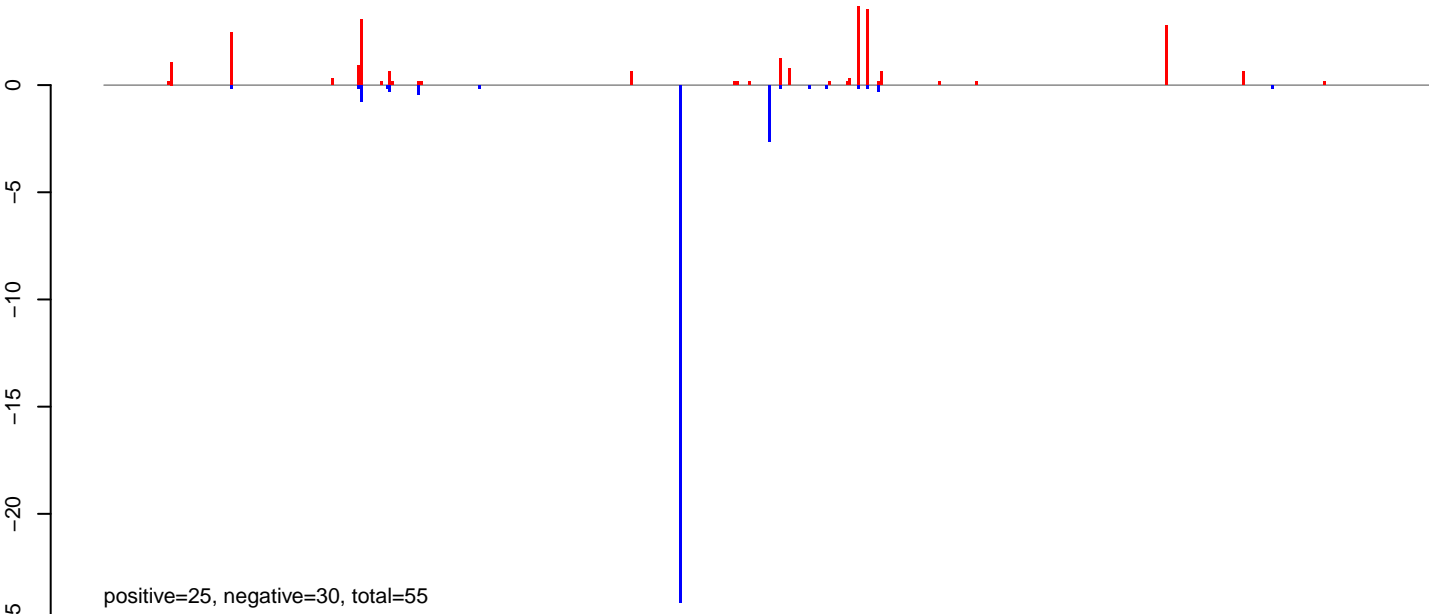
AeAlbo_C636_6dpi_SINV1.rep



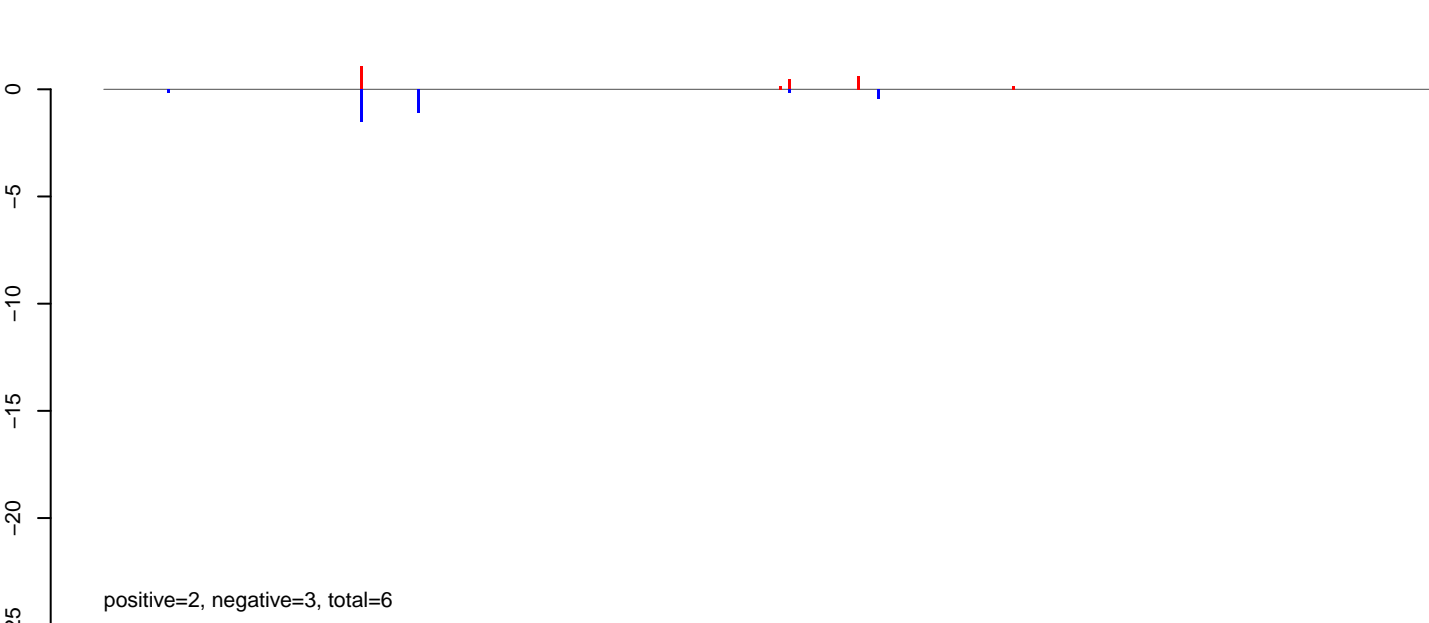
downloaded from the JGI database on 01/11/2016 14:06:52 UTC. ID: 47991.1-Trocaravirus_non_structural_polyprotein_gene_partial_cds_and_structural_polyprotei:1-12052

0 2000 4000 6000 8000 10000 12000

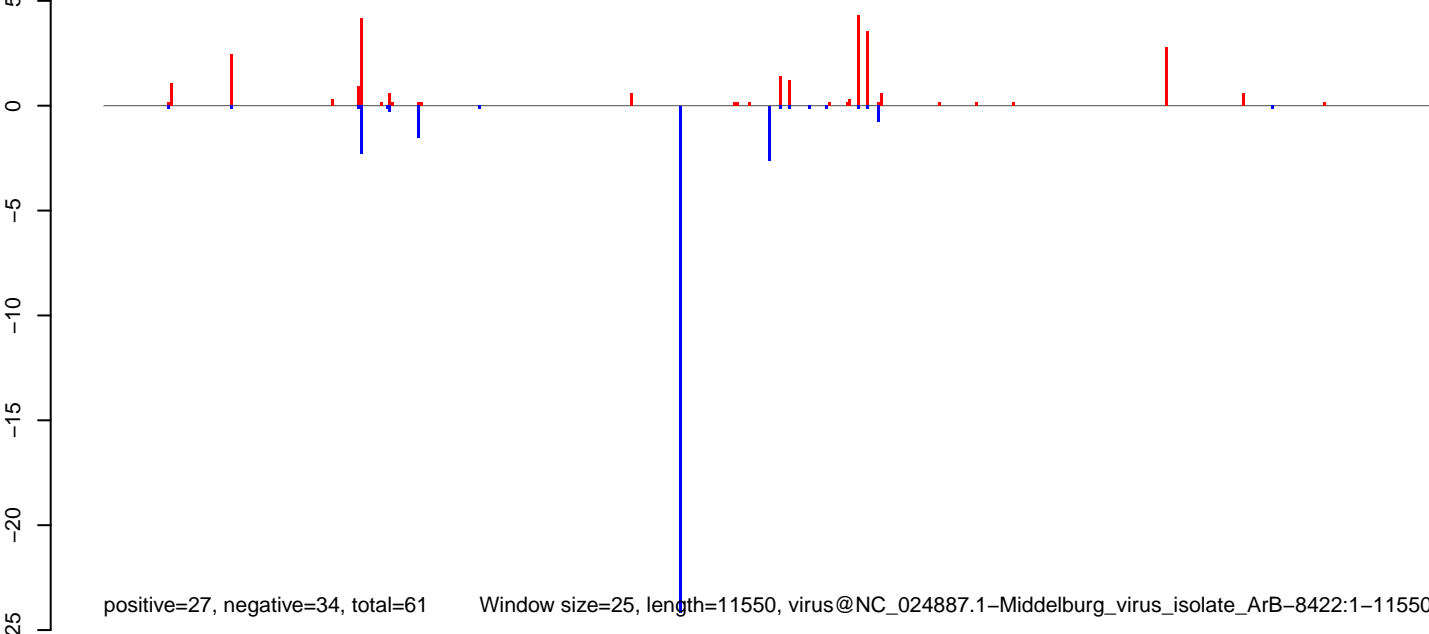
AeAlbo_C636_6dpi_SINV1.18_23.rep



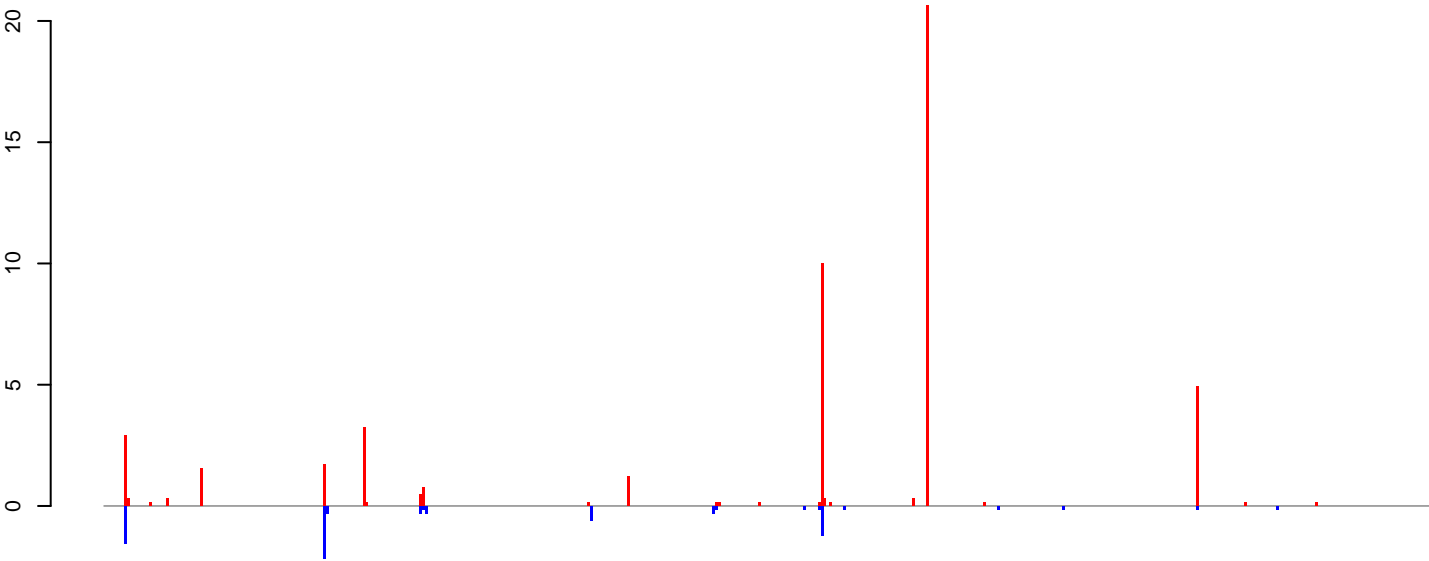
AeAlbo_C636_6dpi_SINV1.24_35.rep



AeAlbo_C636_6dpi_SINV1.rep

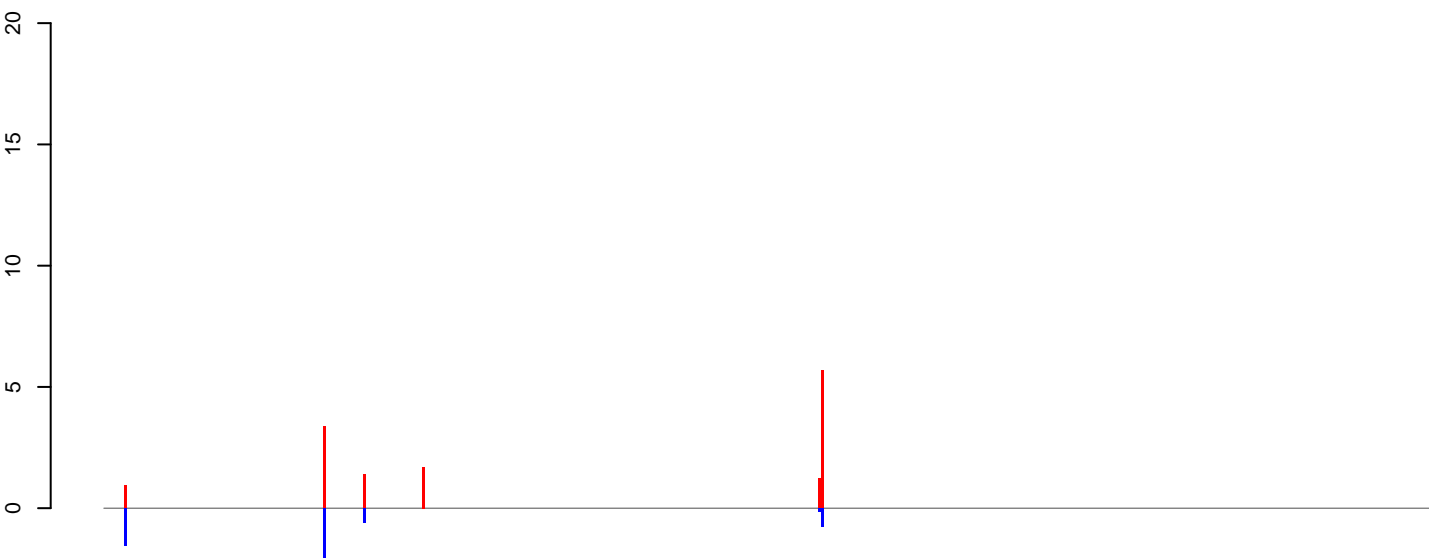


AeAlbo_C636_6dpi_SINV1.18_23.rep



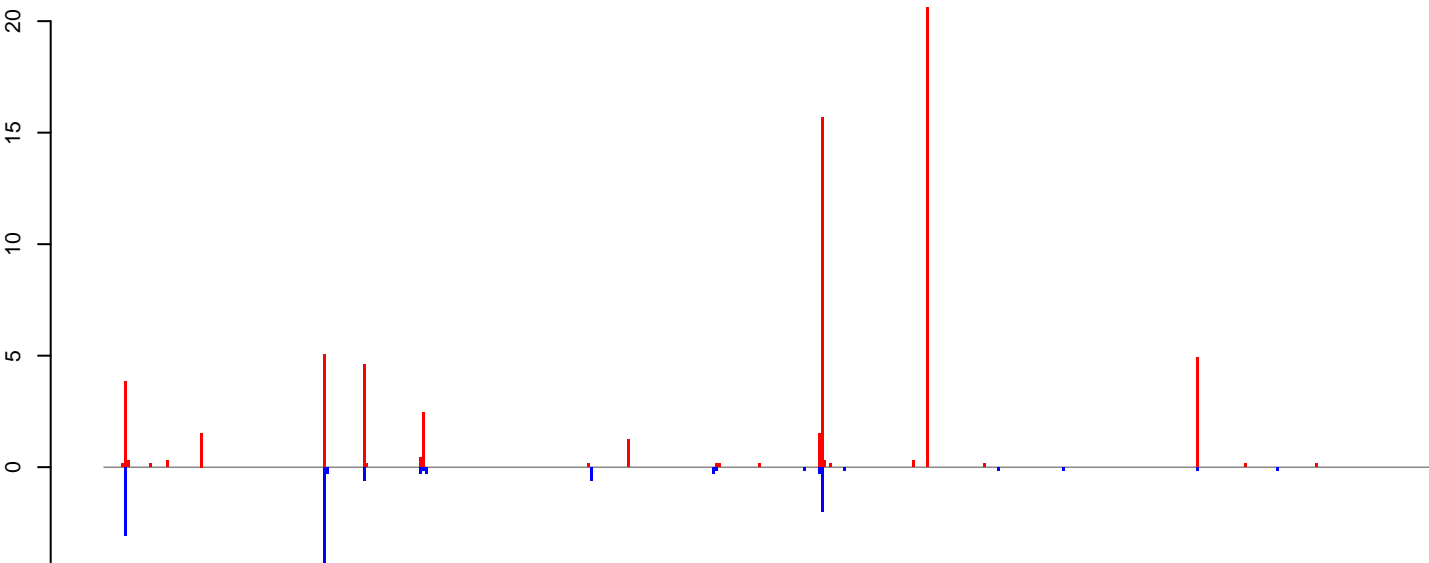
positive=50, negative=8, total=58

AeAlbo_C636_6dpi_SINV1.24_35.rep



positive=14, negative=7, total=21

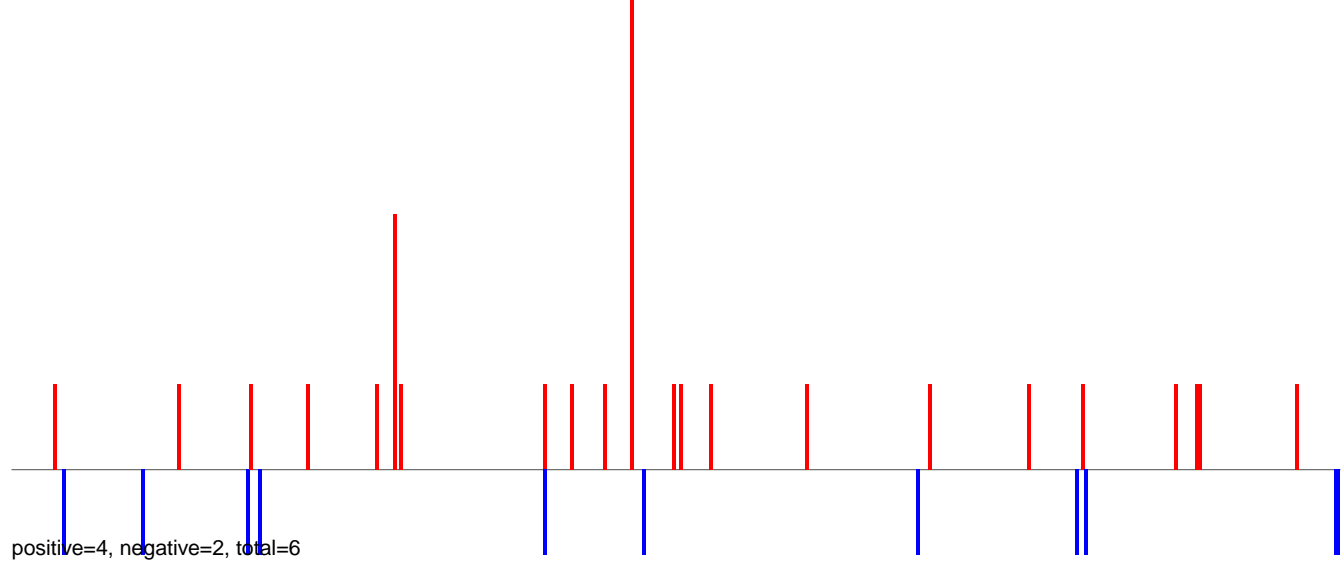
AeAlbo_C636_6dpi_SINV1.rep



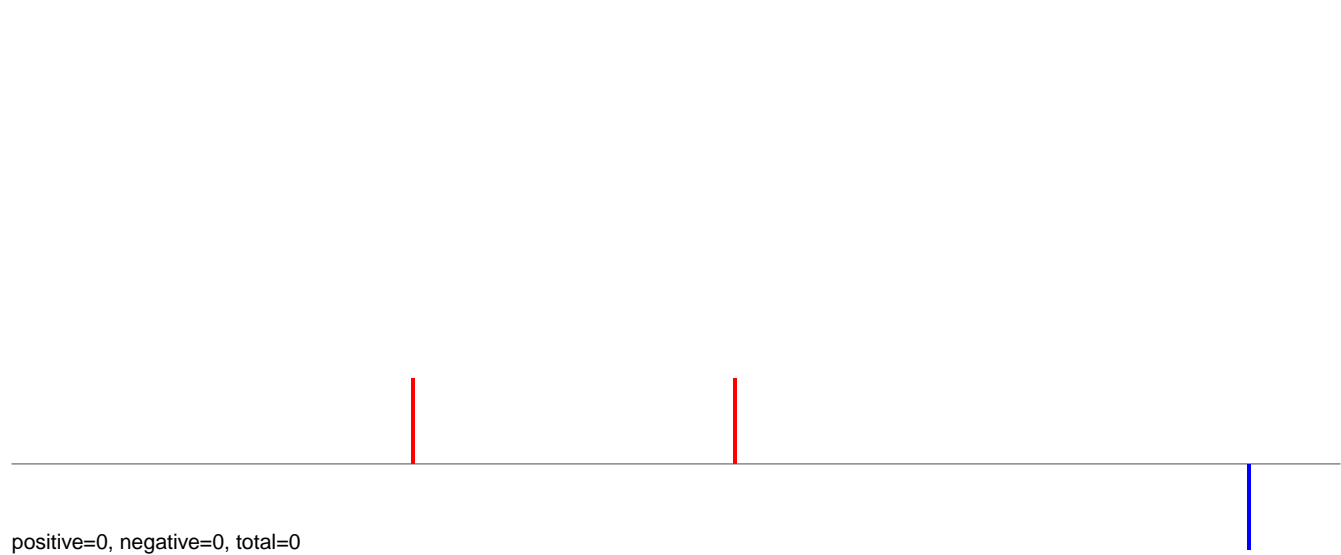
positive=65, negative=15, total=80Window size=25, length=11657, virus@M20162.1-Ross_River_virus_RRV_strain_NB5092_RNA:1-11657

0 2000 4000 6000 8000 10000 12000

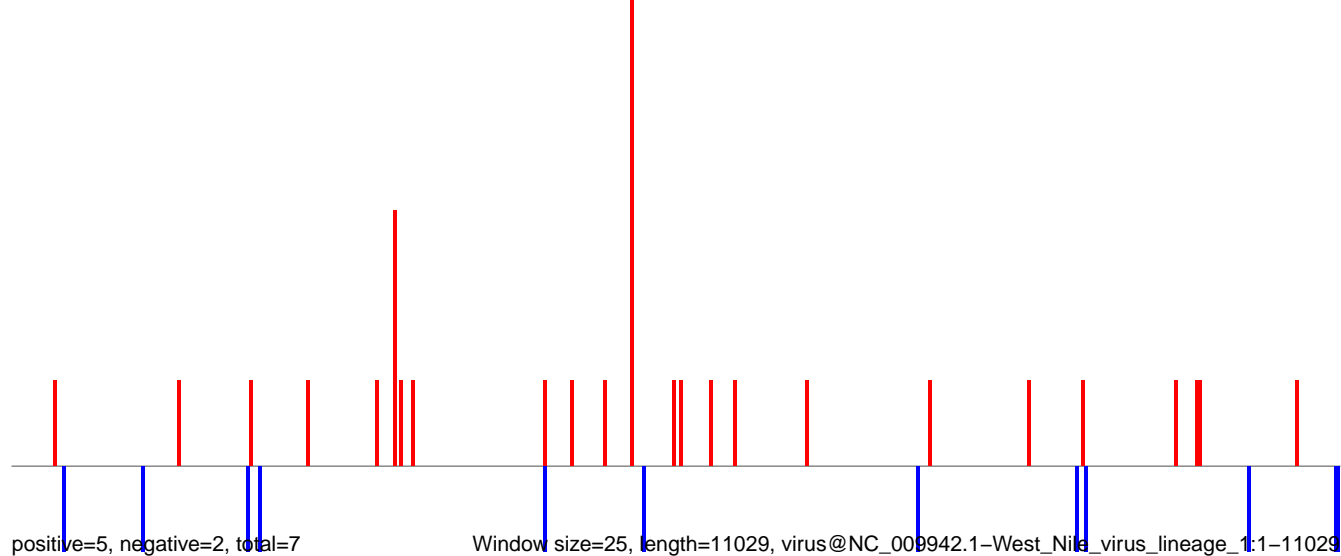
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AeAlbo_C636_6dpi_SINV1.24_35.rep



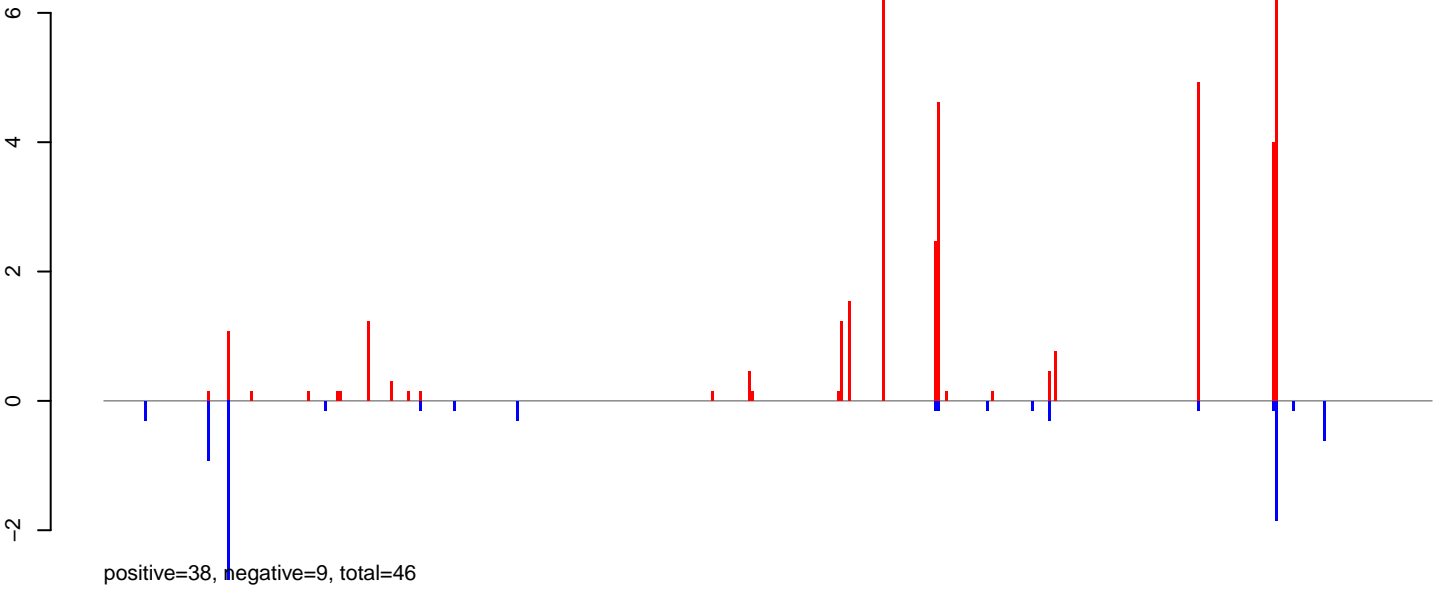
AeAlbo_C636_6dpi_SINV1.rep



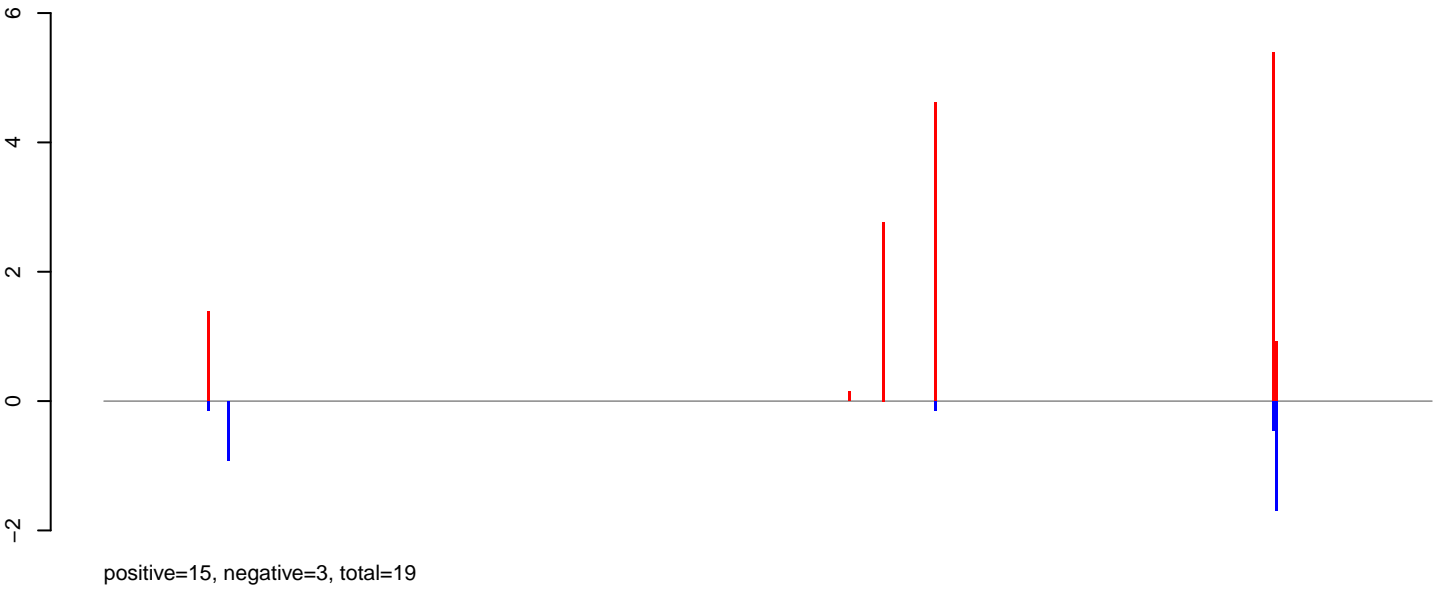
Window size=25, length=11029, virus@NC_009942.1-West_Nile_virus_lineage_1:1-11029

0 2000 4000 6000 8000 10000

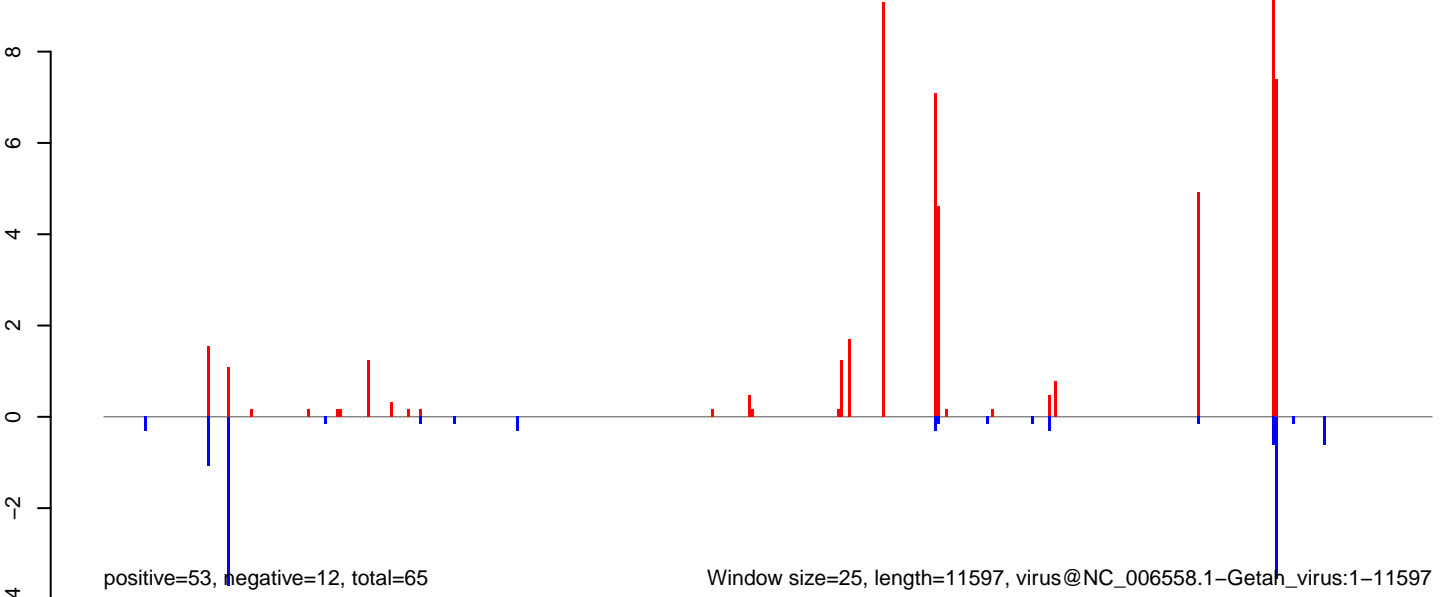
AeAlbo_C636_6dpi_SINV1.18_23.rep



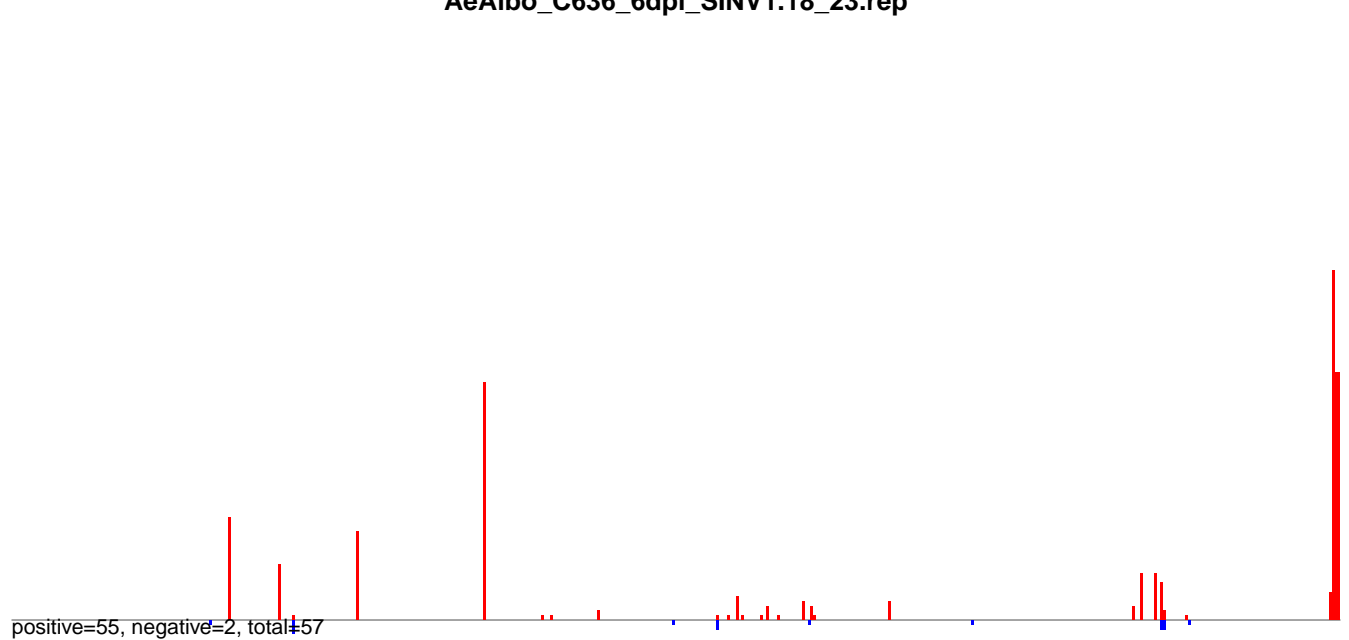
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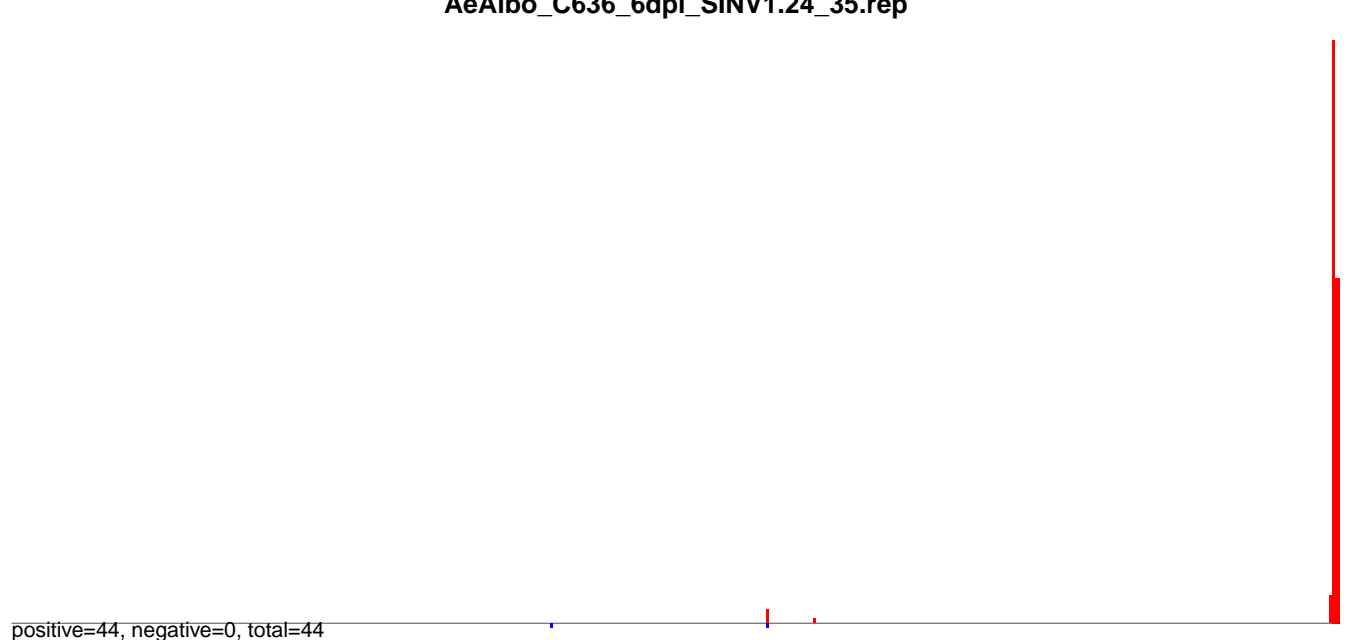
AeAlbo_C636_6dpi_SINV1.rep



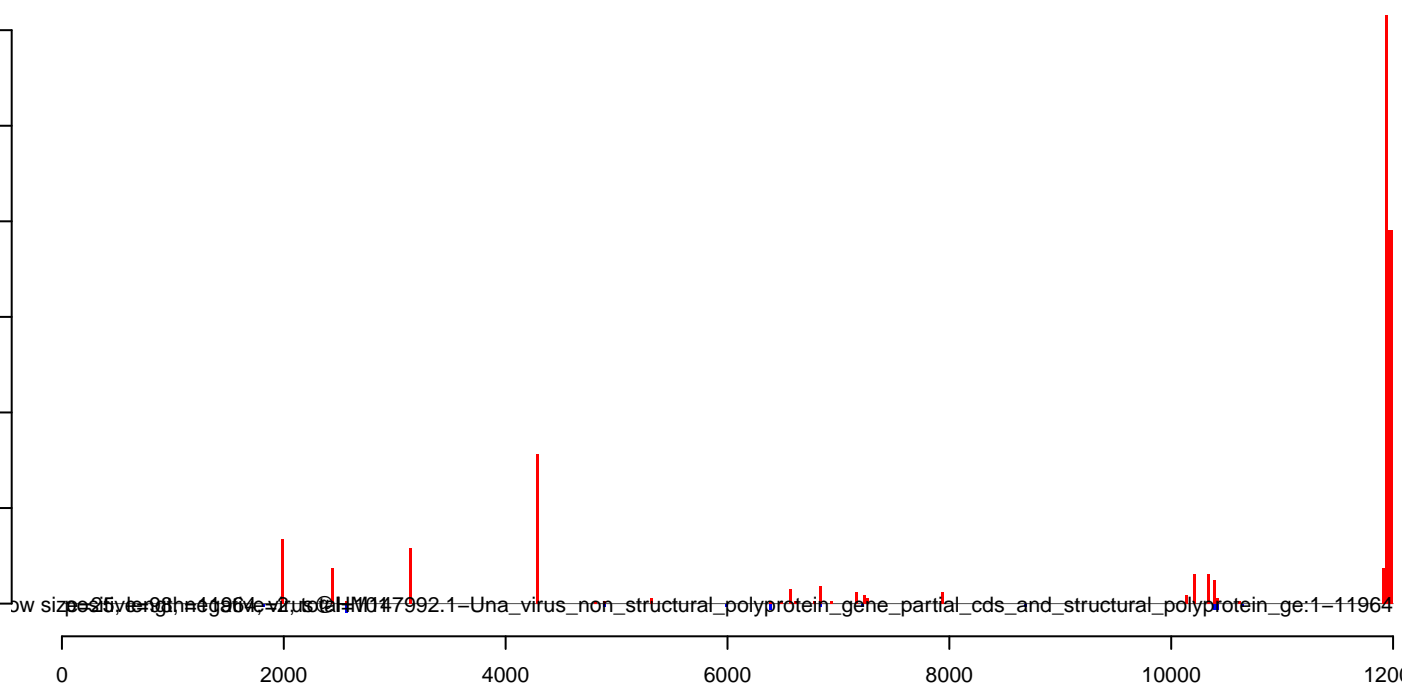
AeAlbo_C636_6dpi_SINV1.18_23.rep



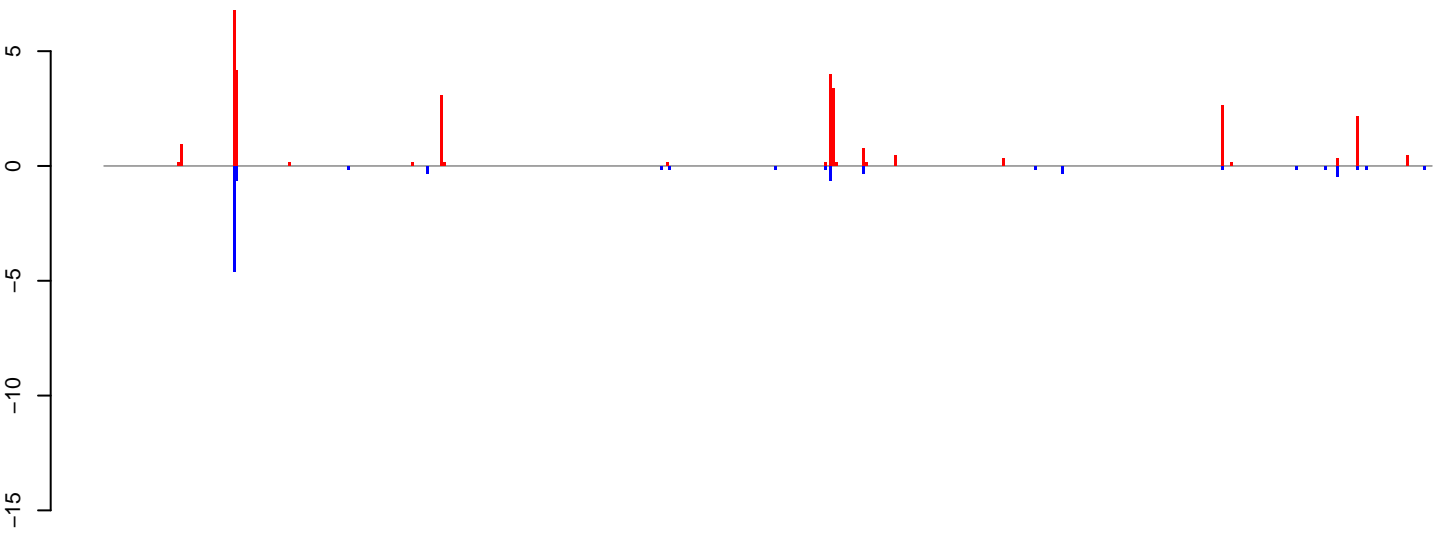
AeAlbo_C636_6dpi_SINV1.24_35.rep



AeAlbo_C636_6dpi_SINV1.rep



AeAlbo_C636_6dpi_SINV1.18_23.rep



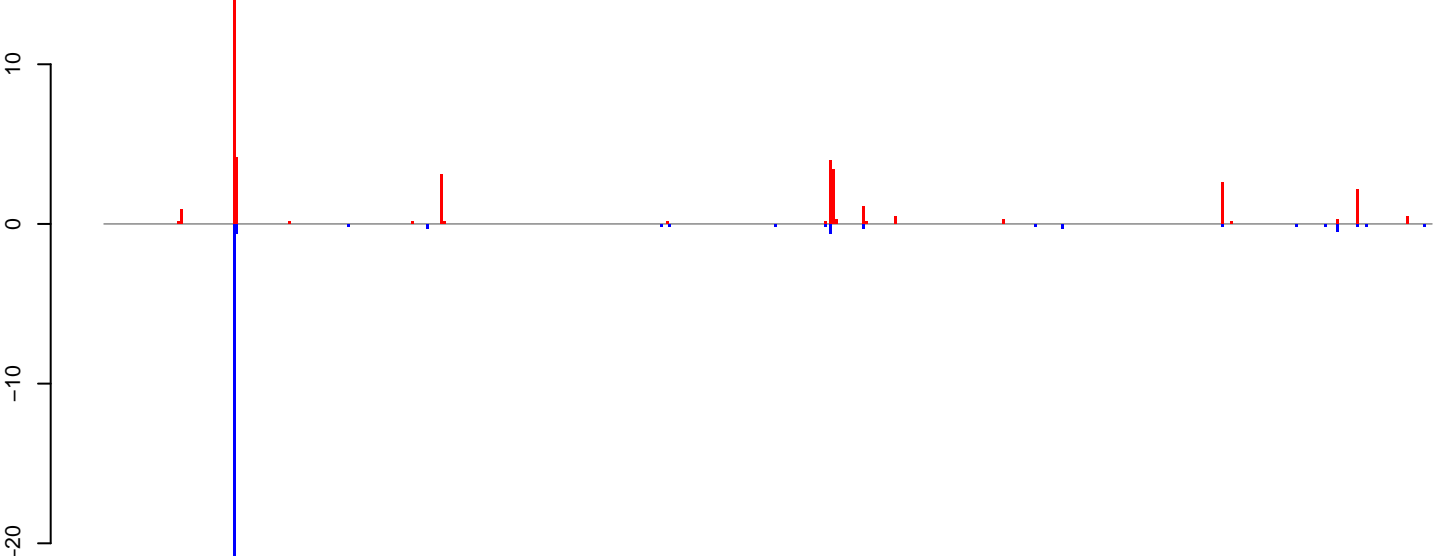
positive=31, negative=9, total=40

AeAlbo_C636_6dpi_SINV1.24_35.rep



positive=8, negative=19, total=26

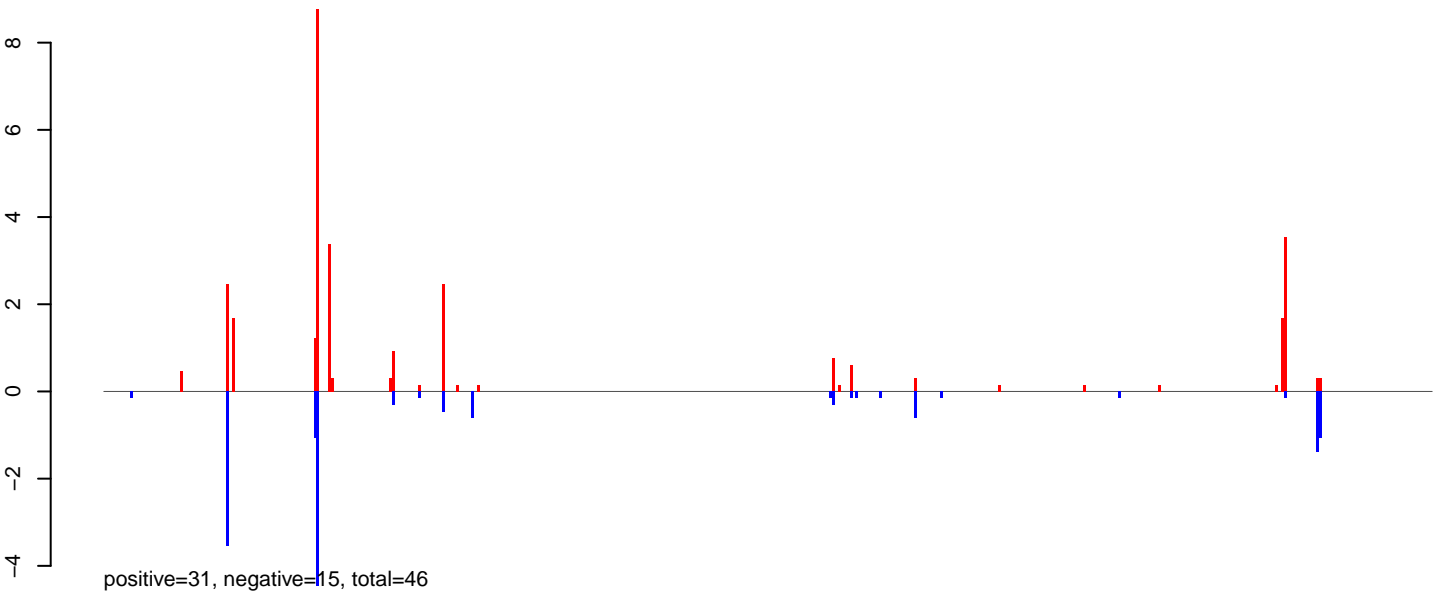
AeAlbo_C636_6dpi_SINV1.rep



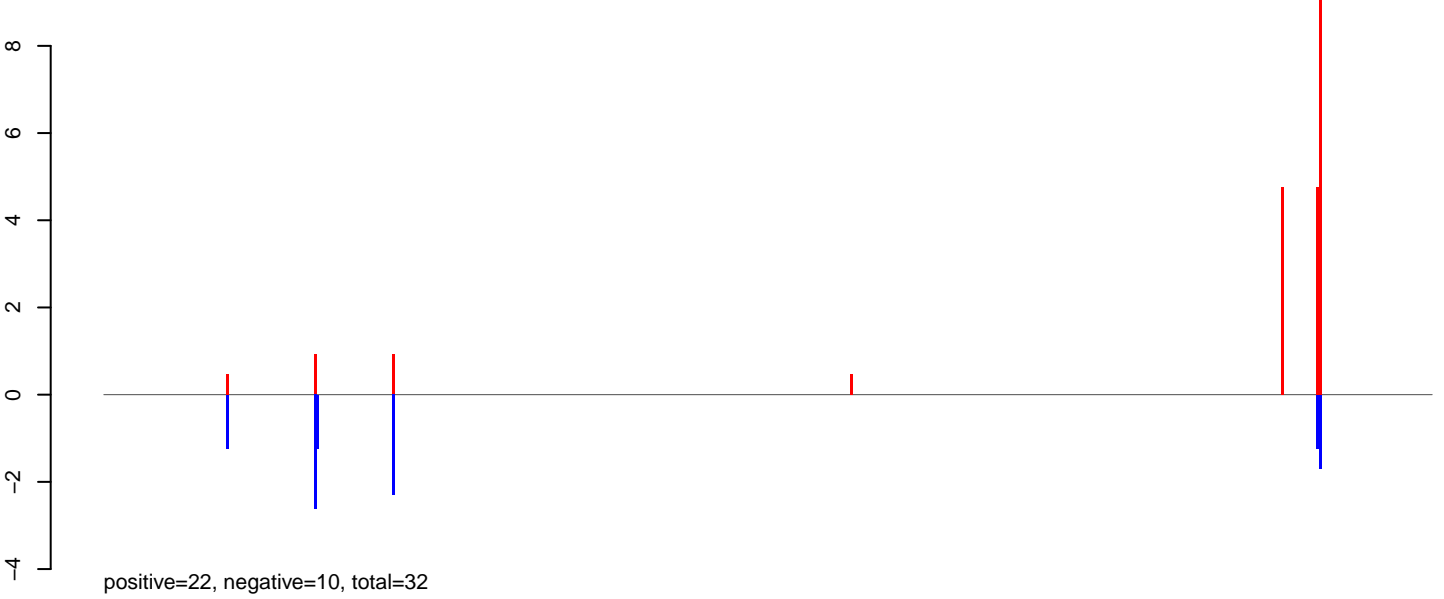
positive=38, negative=28, total=66 Window size=25, length=11344, virus@AF075256.1-Pixuna_virus_strain_Pixuna_BeAr_35645:1-11344

0 2000 4000 6000 8000 10000

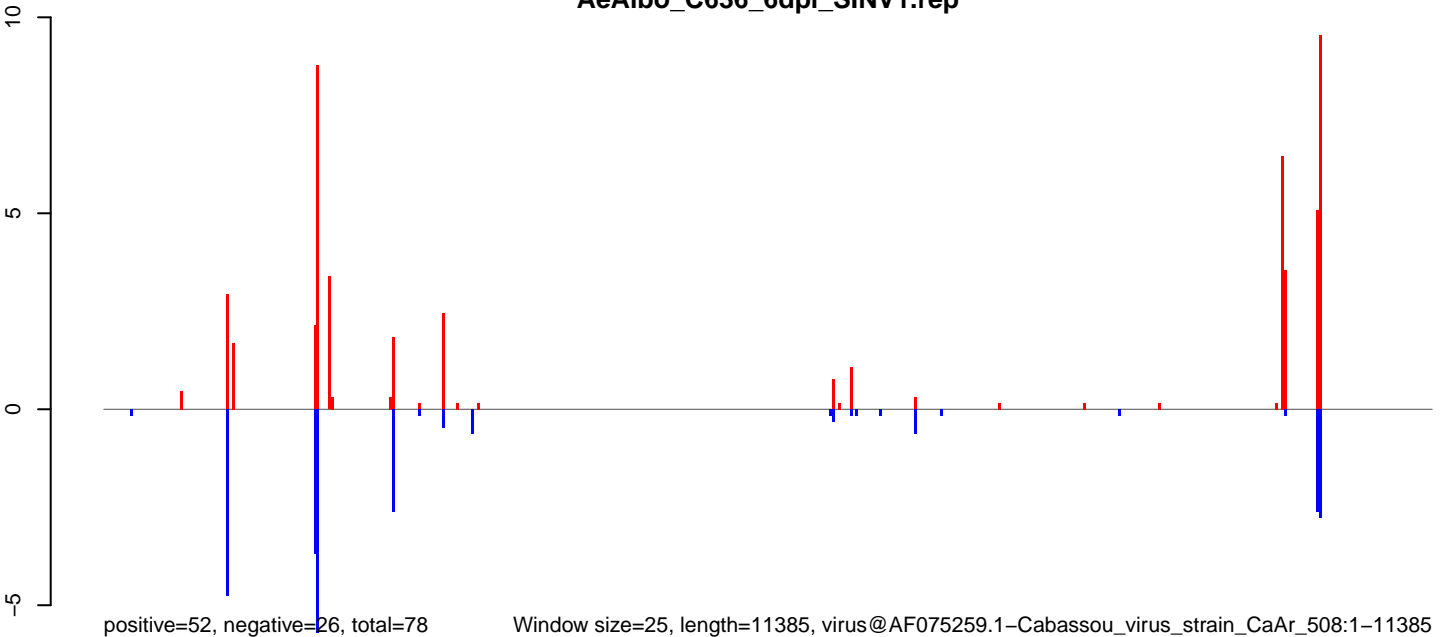
AeAlbo_C636_6dpi_SINV1.18_23.rep



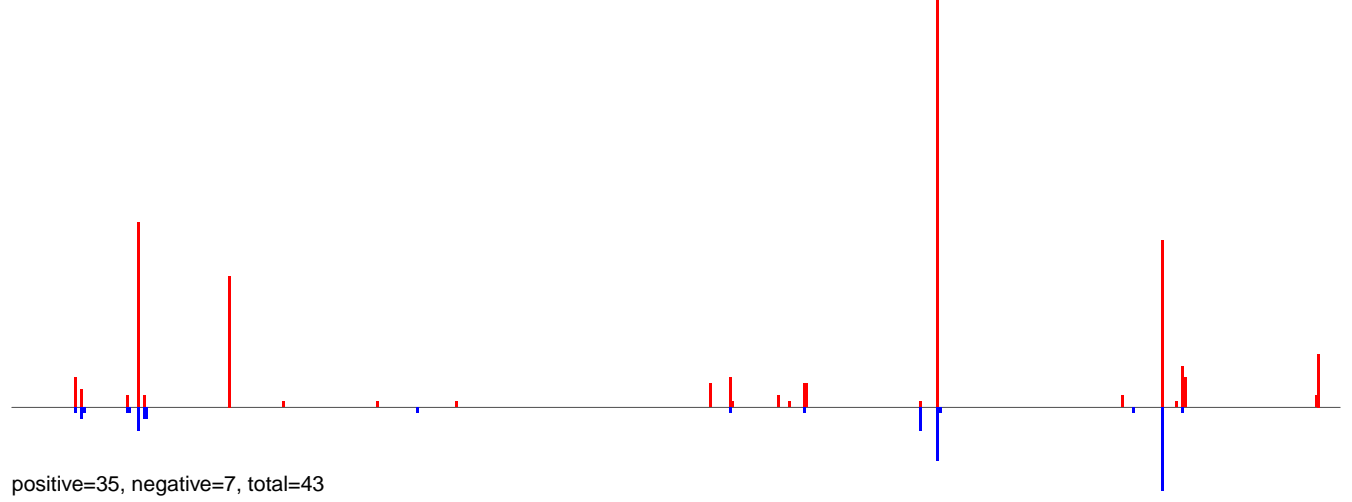
AeAlbo_C636_6dpi_SINV1.24_35.rep



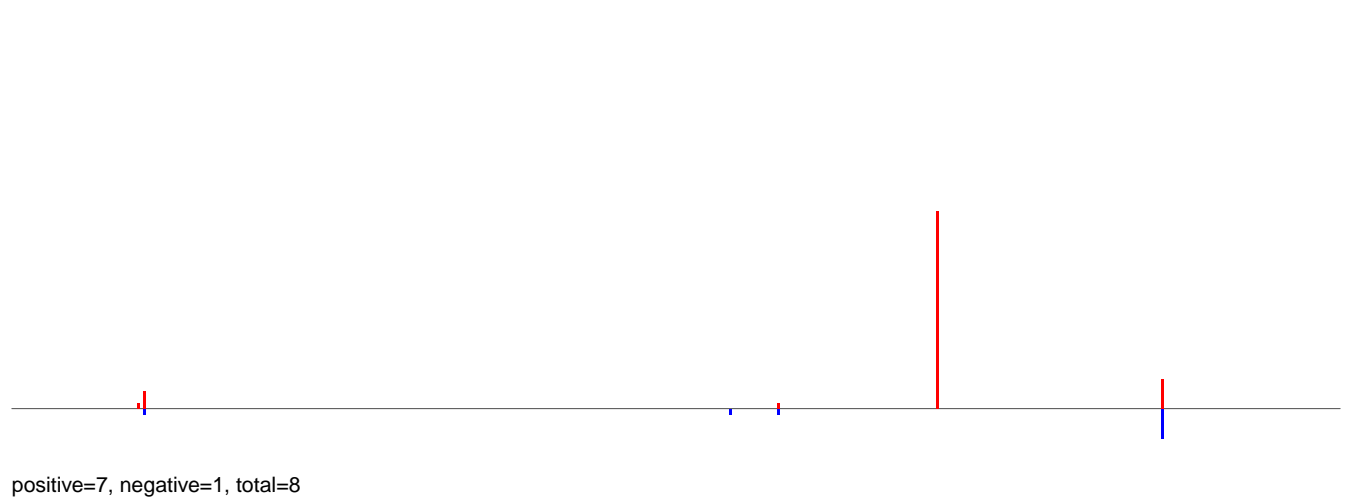
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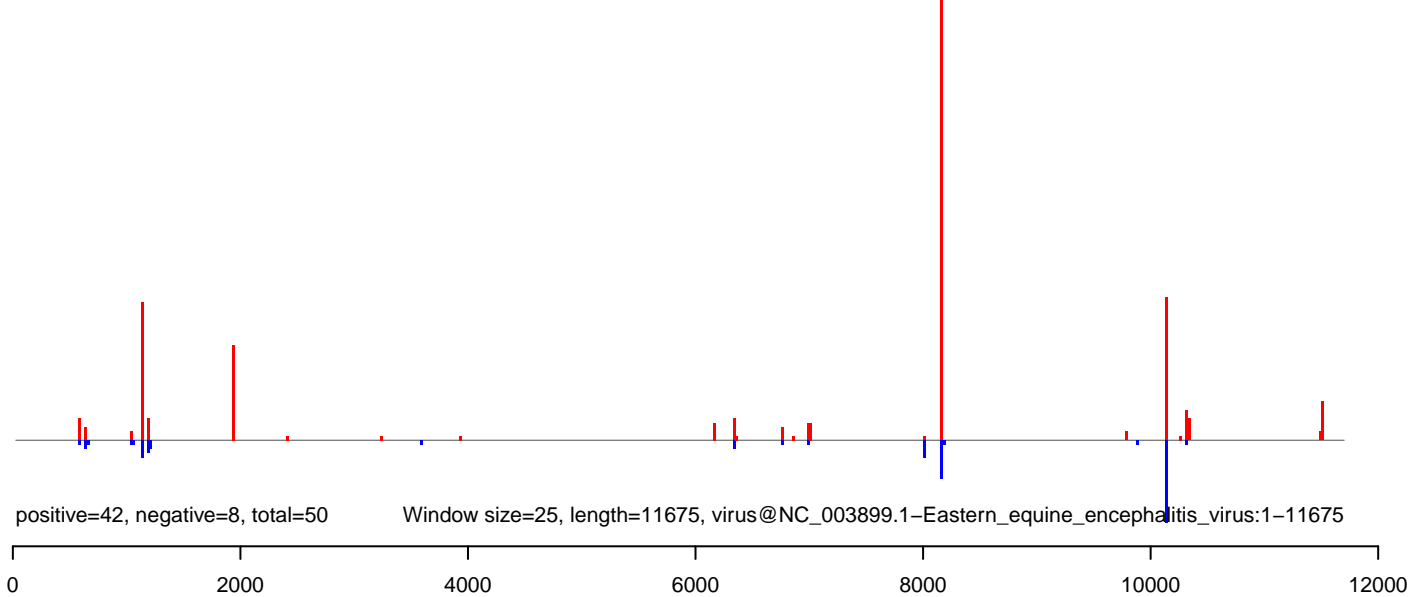
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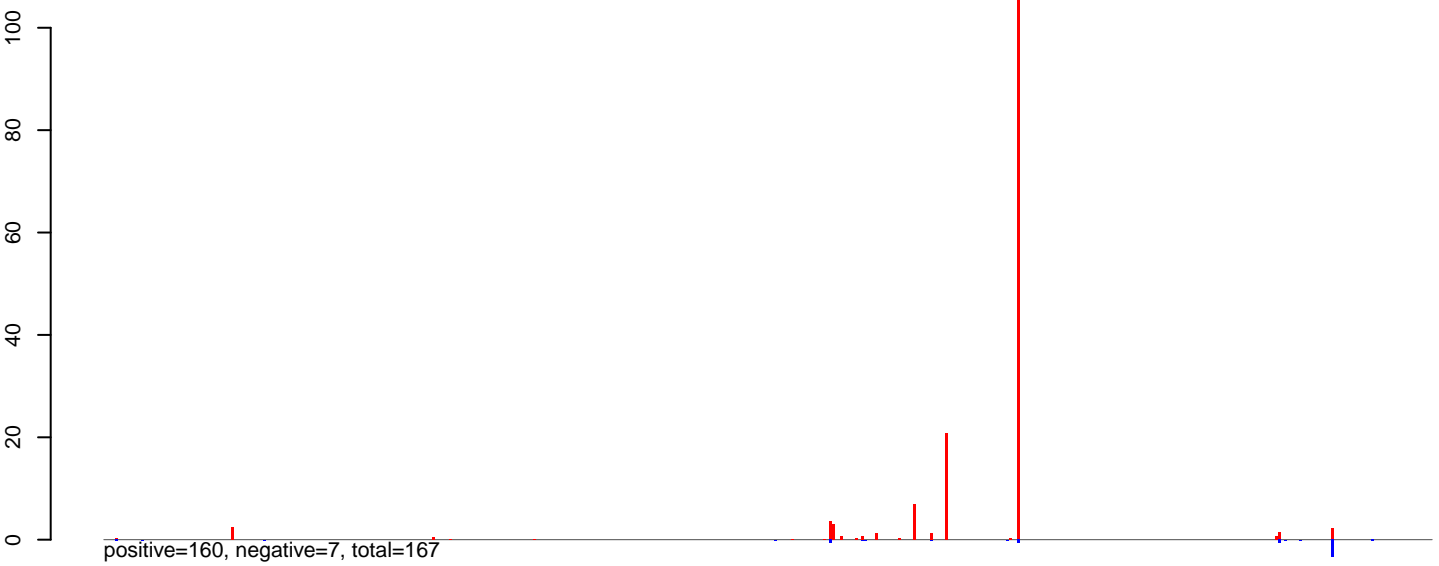
AeAlbo_C636_6dpi_SINV1.24_35.rep



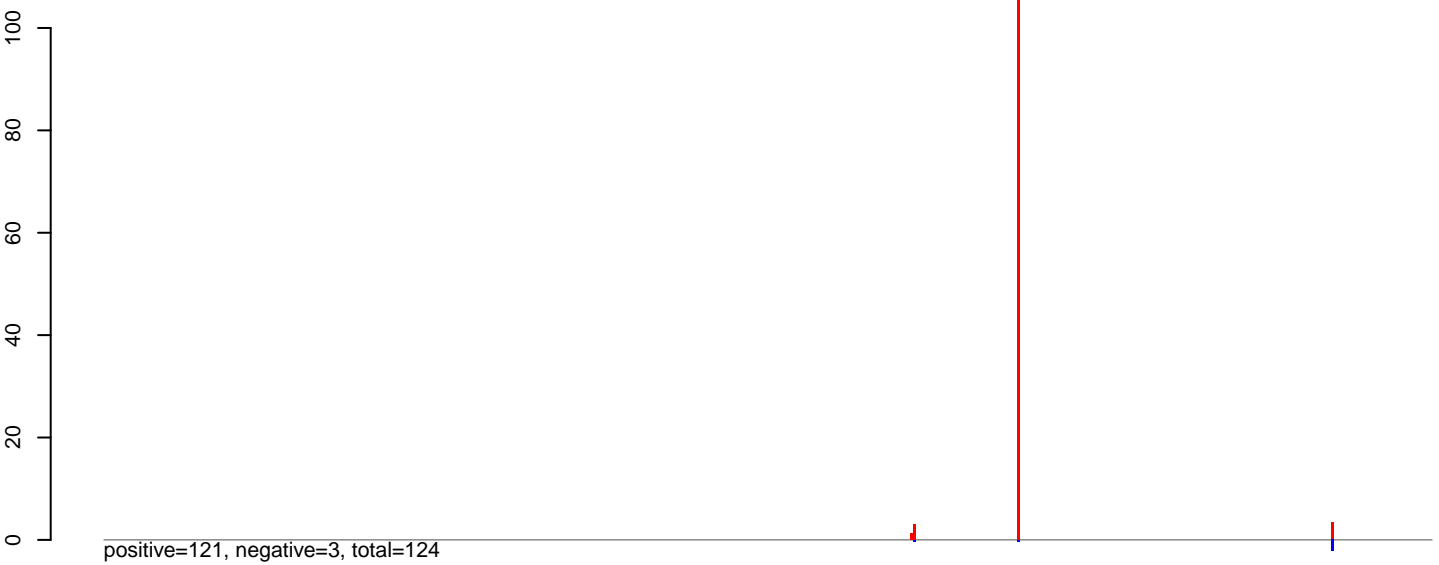
AeAlbo_C636_6dpi_SINV1.rep



AeAlbo_C636_6dpi_SINV1.18_23.rep



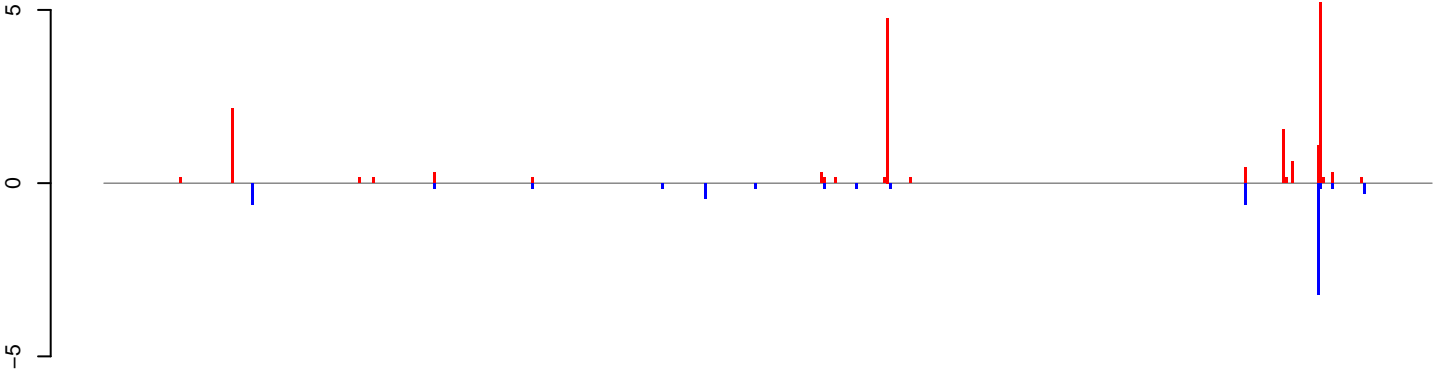
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AeAlbo_C636_6dpi_SINV1.rep

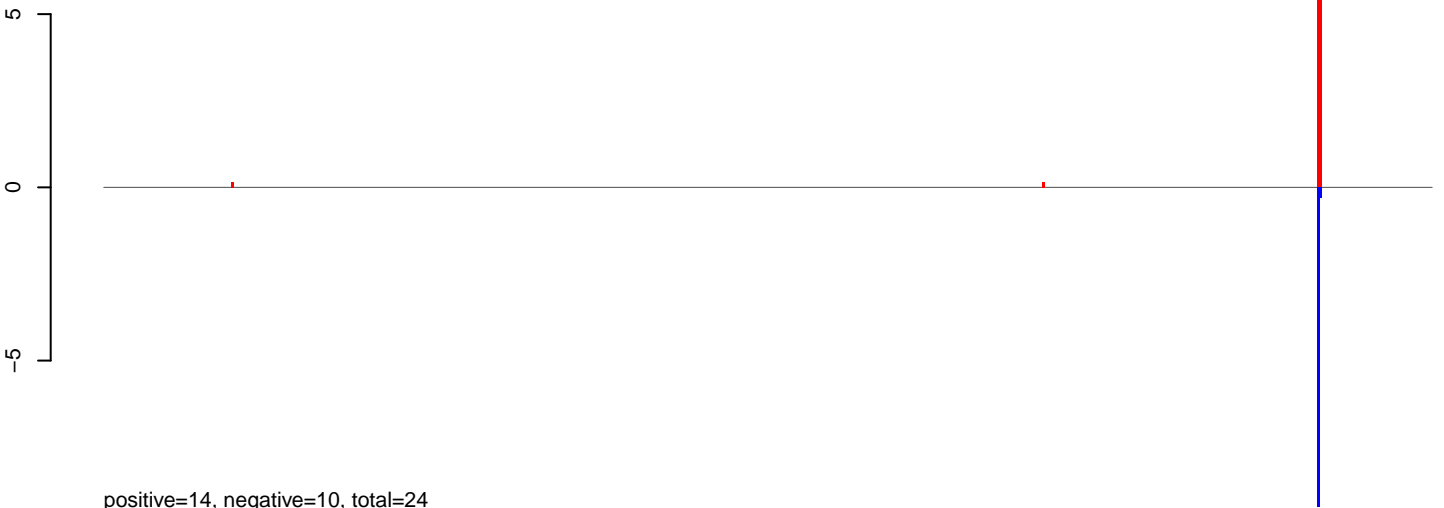


AeAlbo_C636_6dpi_SINV1.18_23.rep



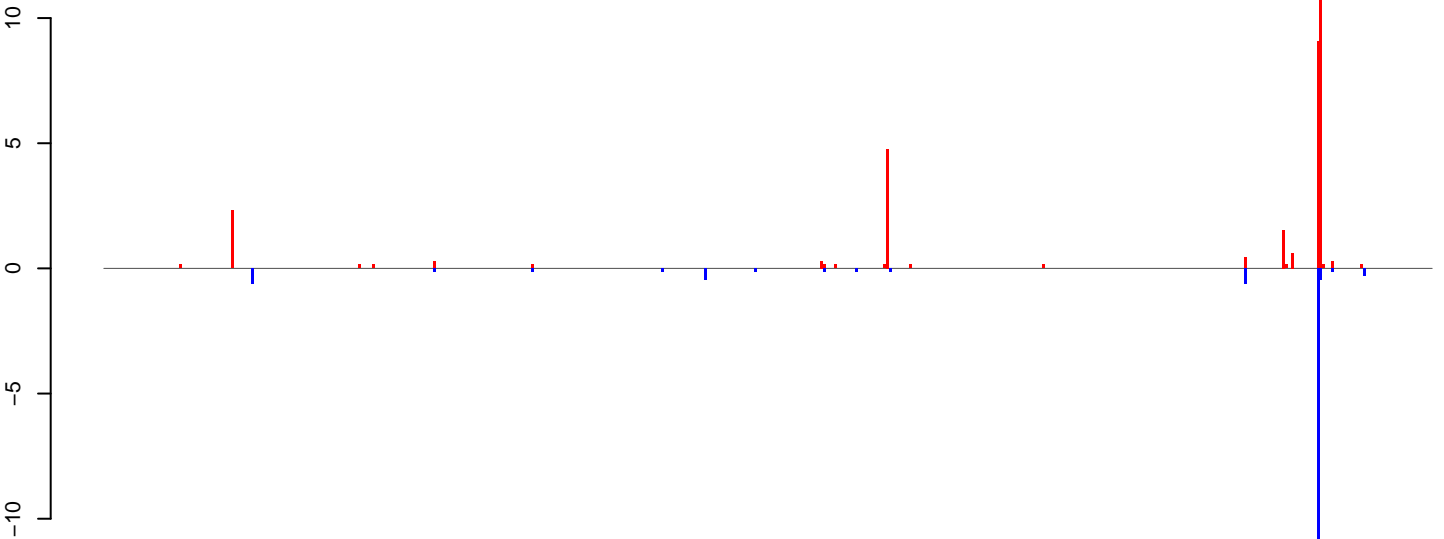
positive=18, negative=7, total=25

AeAlbo_C636_6dpi_SINV1.24_35.rep



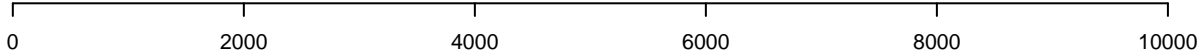
positive=14, negative=10, total=24

AeAlbo_C636_6dpi_SINV1.rep

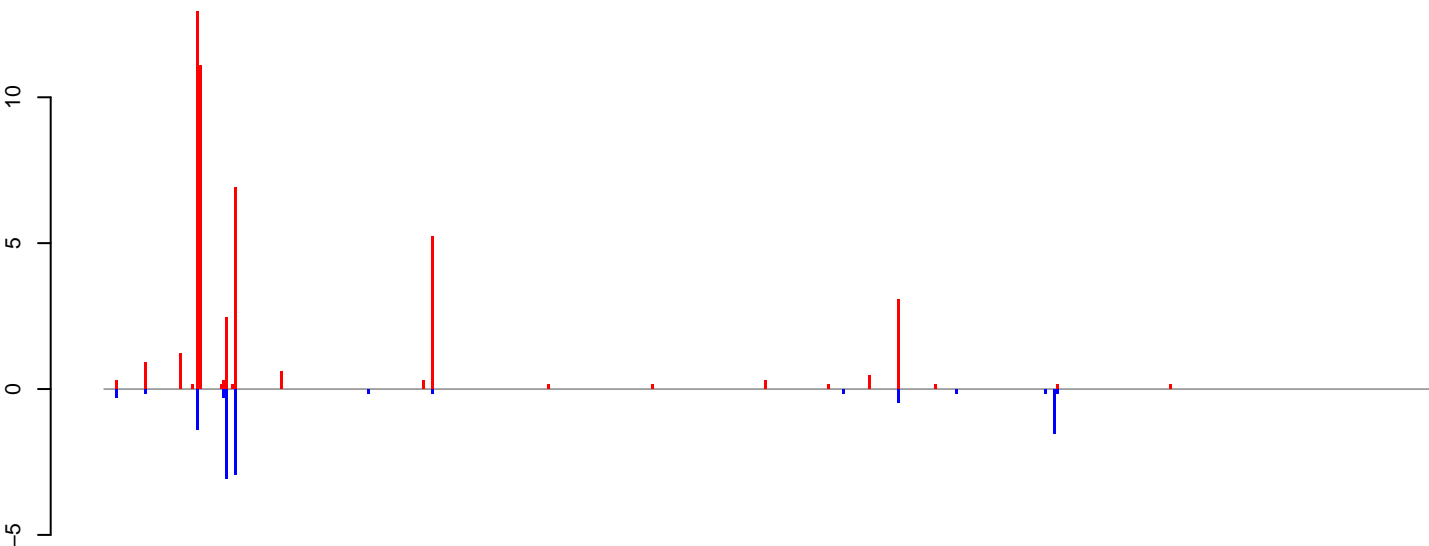


positive=33, negative=16, total=49

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

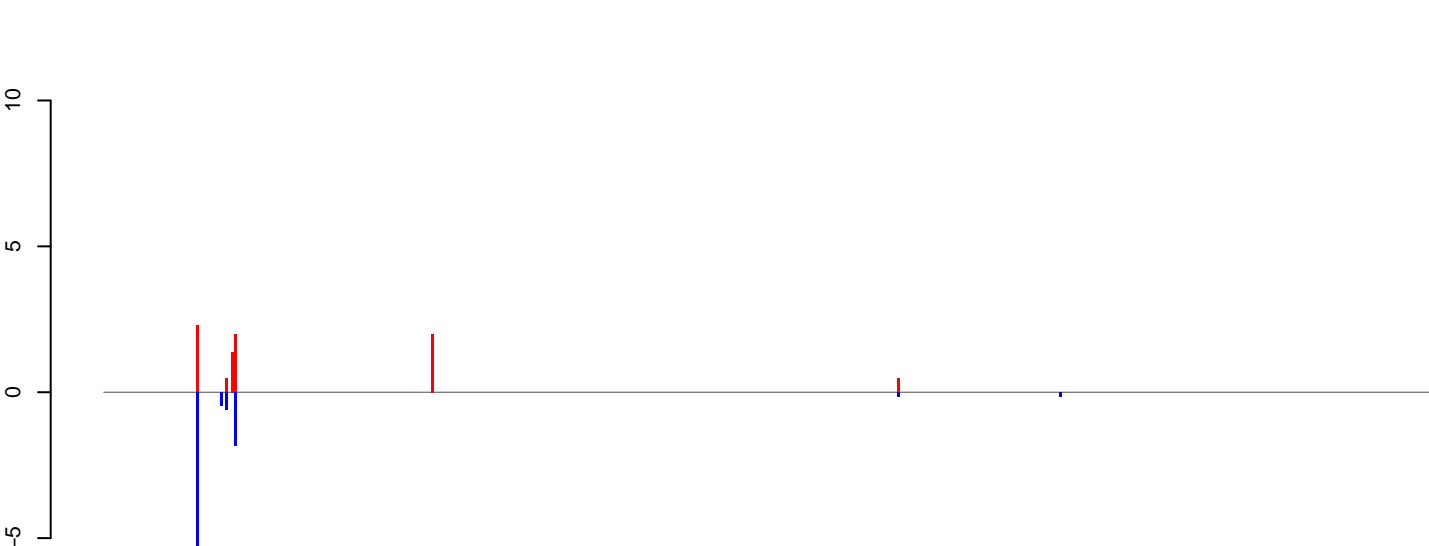


AeAlbo_C636_6dpi_SINV1.18_23.rep



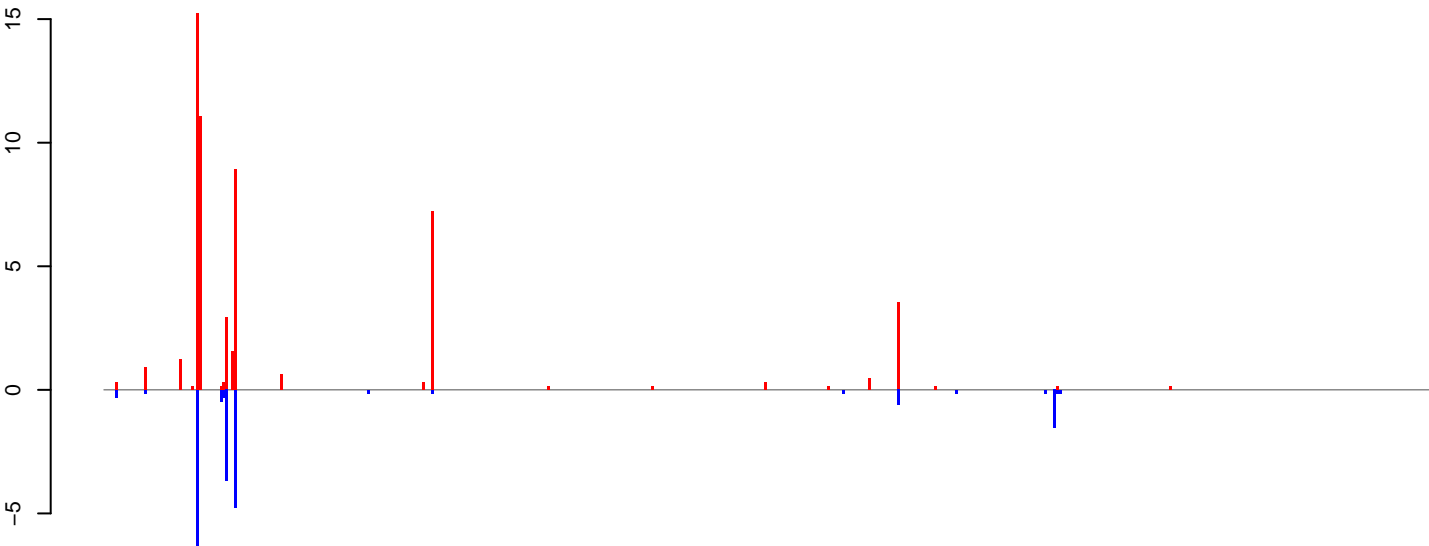
positive=48, negative=11, total=59

AeAlbo_C636_6dpi_SINV1.24_35.rep



positive=9, negative=11, total=19

AeAlbo_C636_6dpi_SINV1.rep

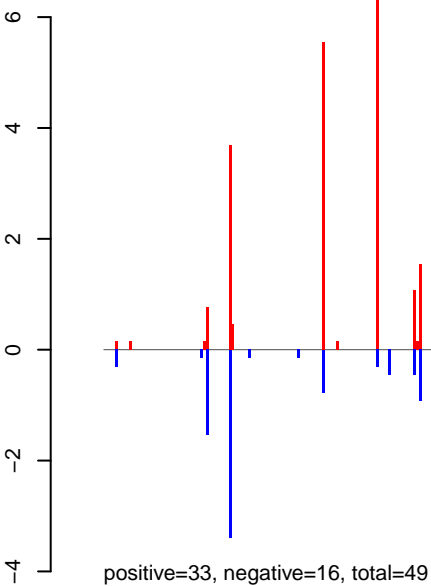


positive=56, negative=22, total=78

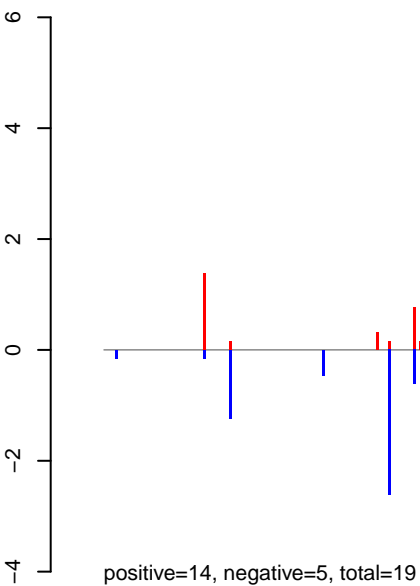
Window size=25, length=11465, virus@AF075257.1-Mosso_das_Pedras_virus:1-11465

0 2000 4000 6000 8000 10000

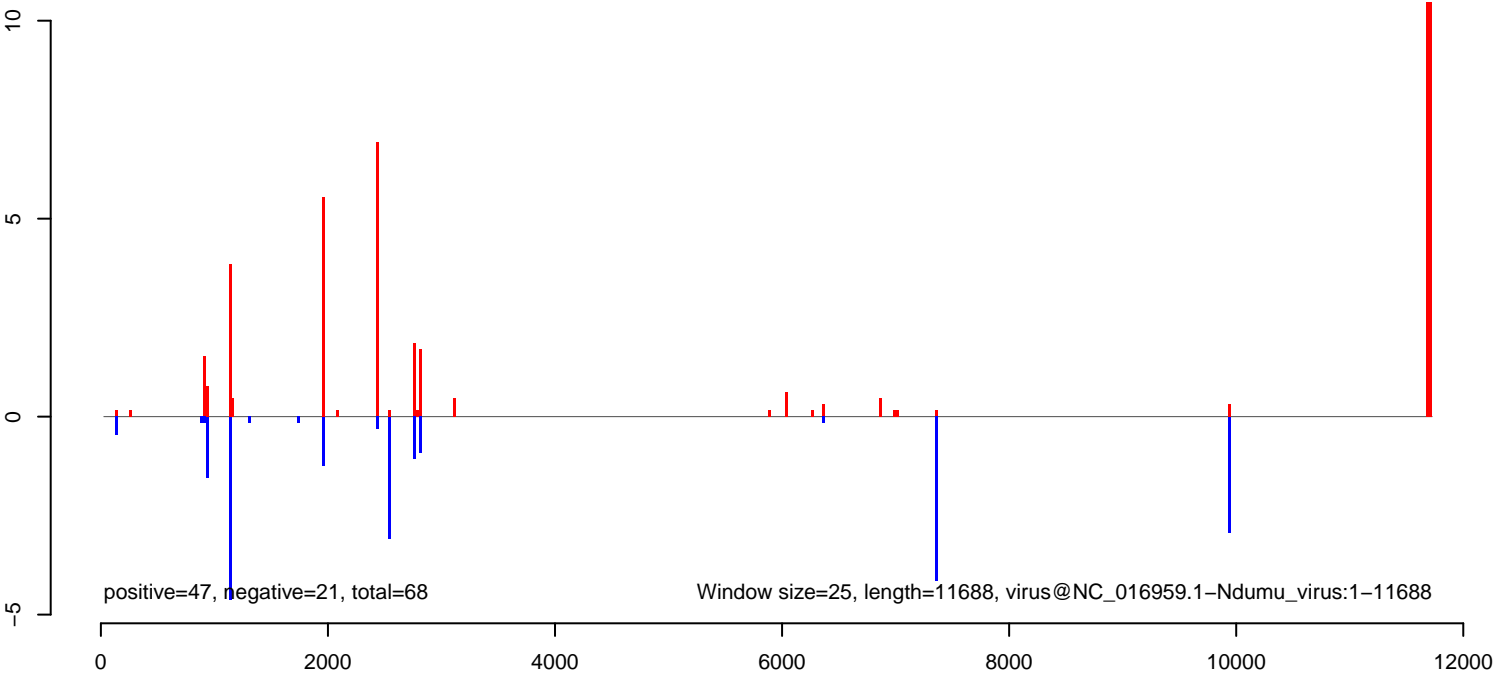
AeAlbo_C636_6dpi_SINV1.18_23.rep



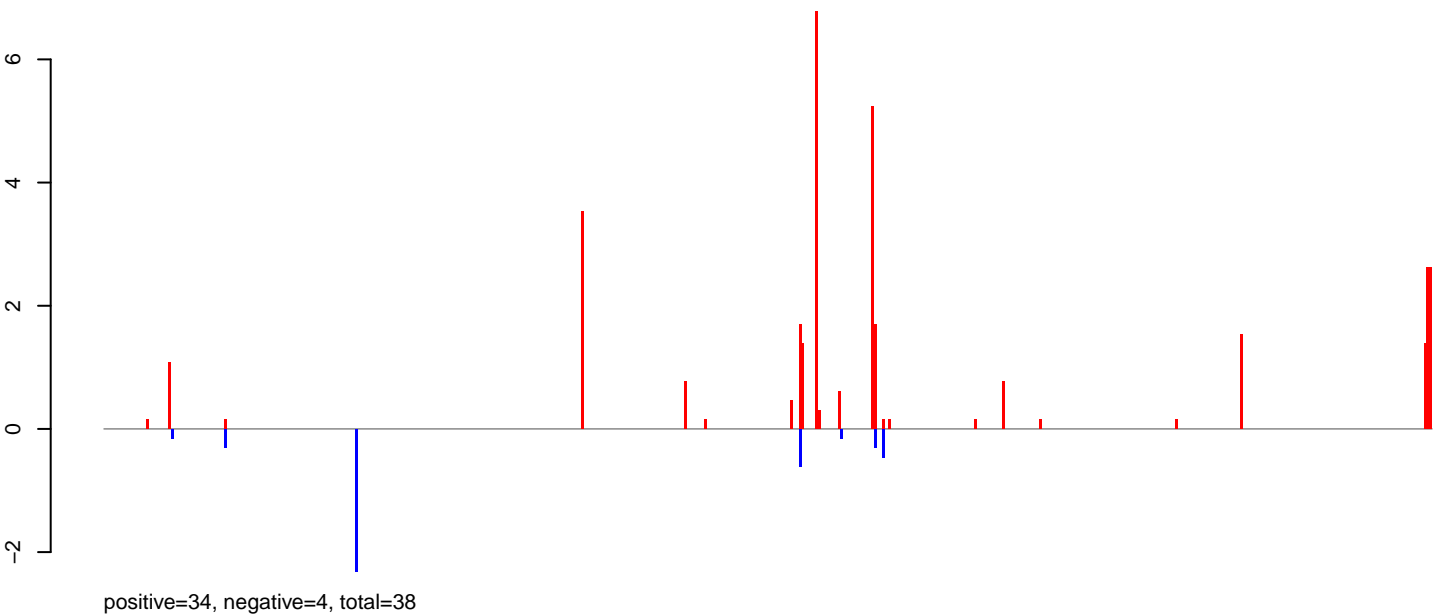
AeAlbo_C636_6dpi_SINV1.24_35.rep



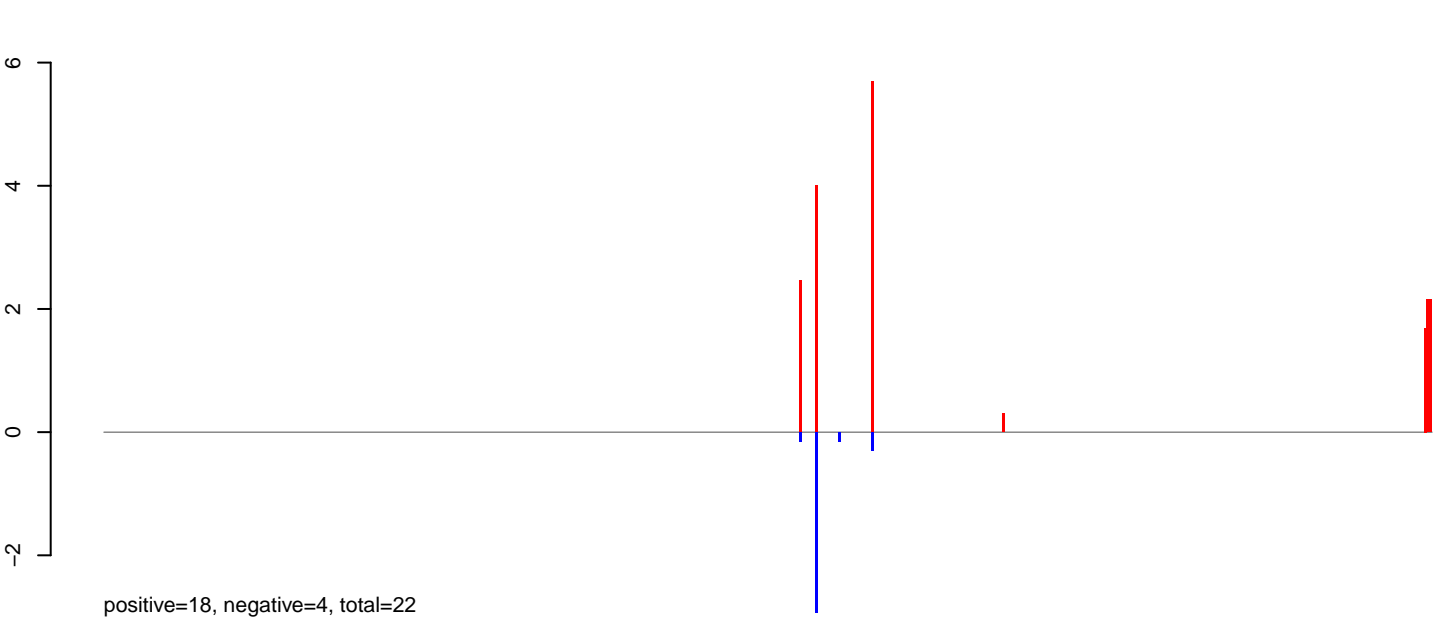
AeAlbo_C636_6dpi_SINV1.rep



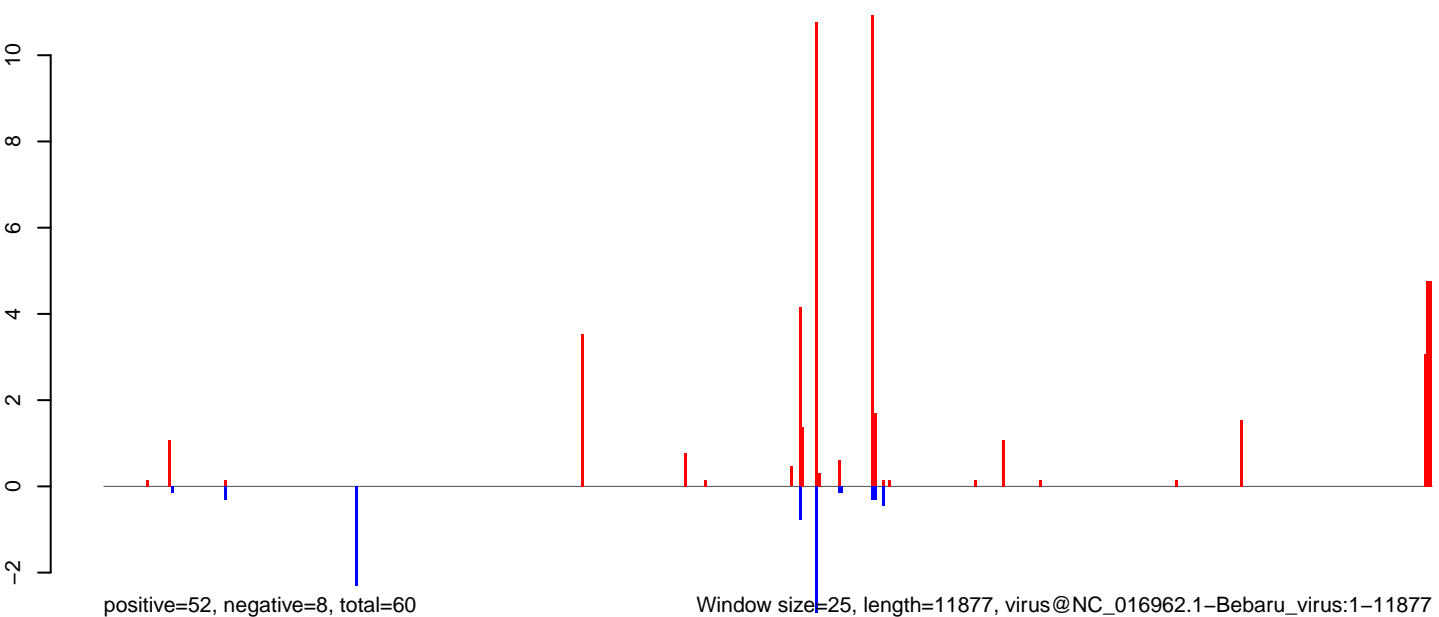
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep

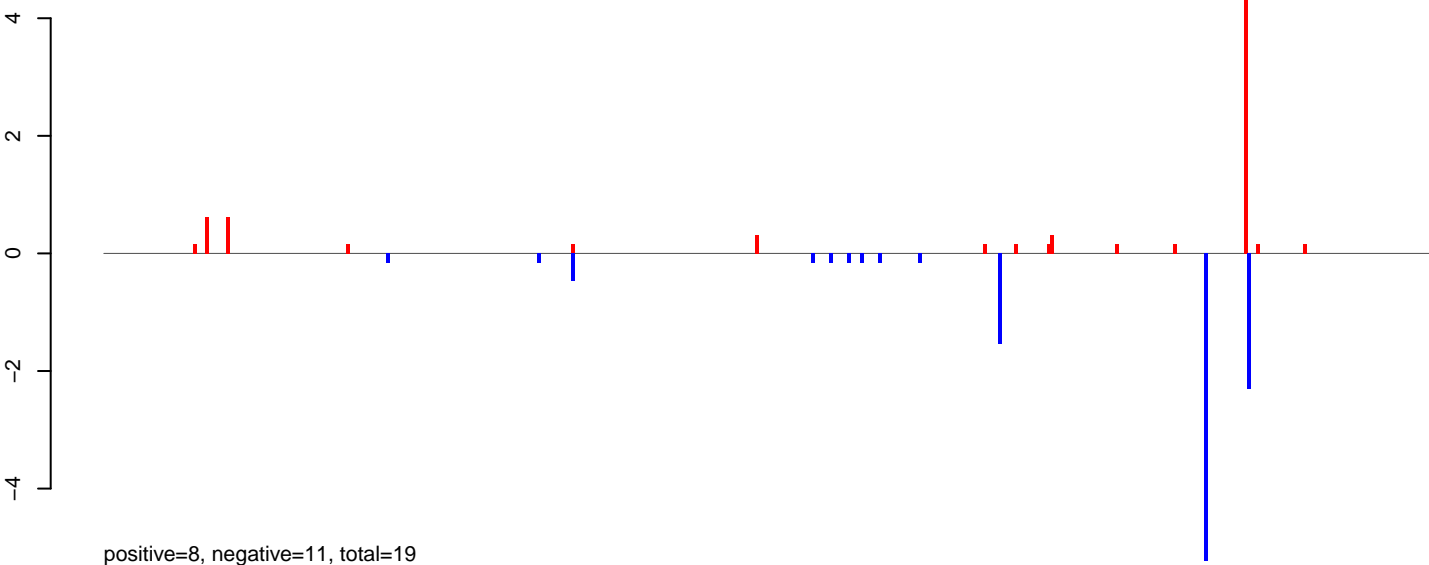


AeAlbo_C636_6dpi_SINV1.rep

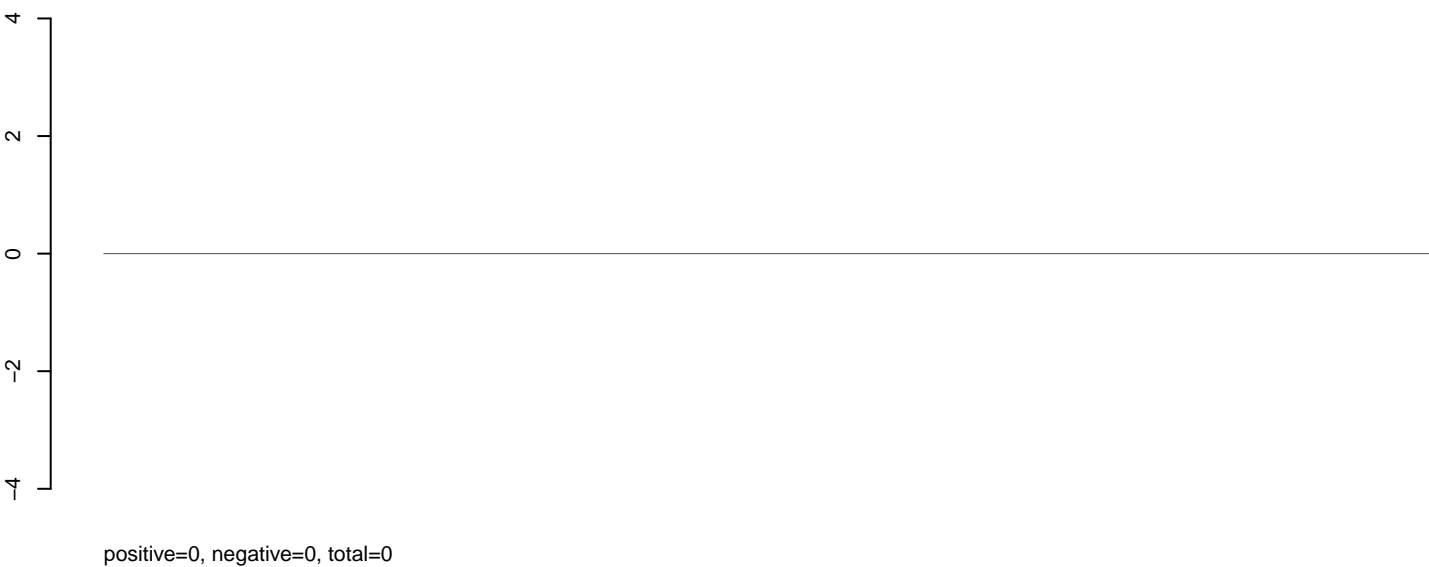


0 2000 4000 6000 8000 10000 12000

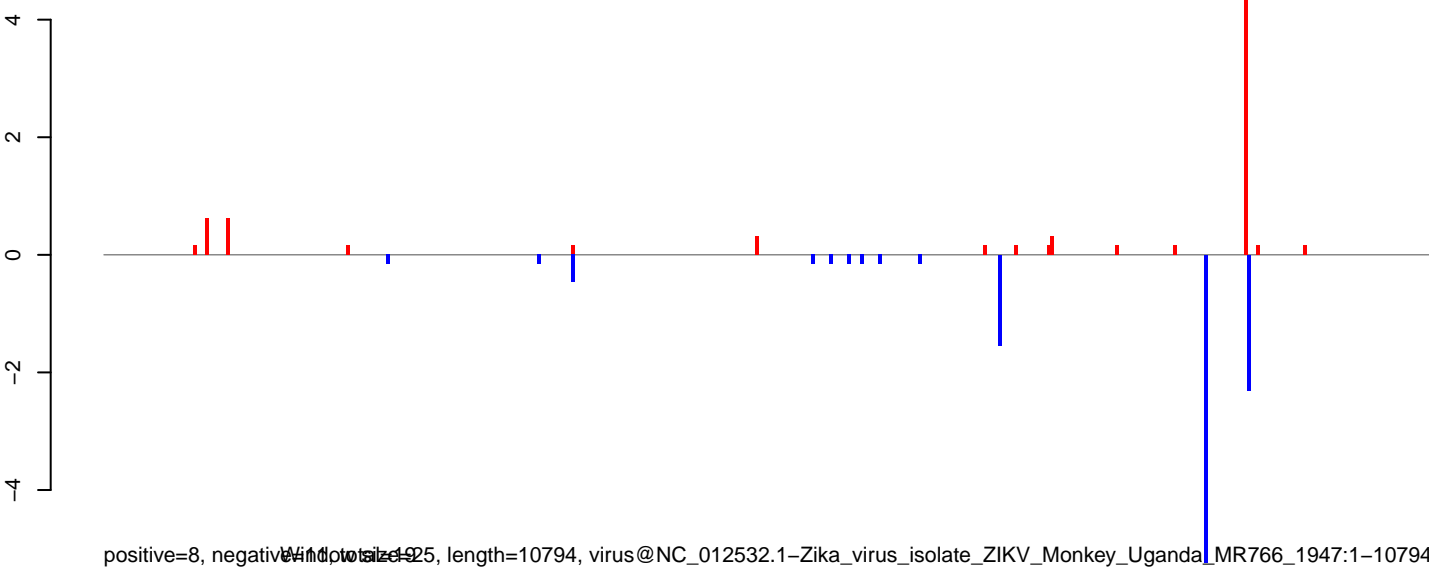
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



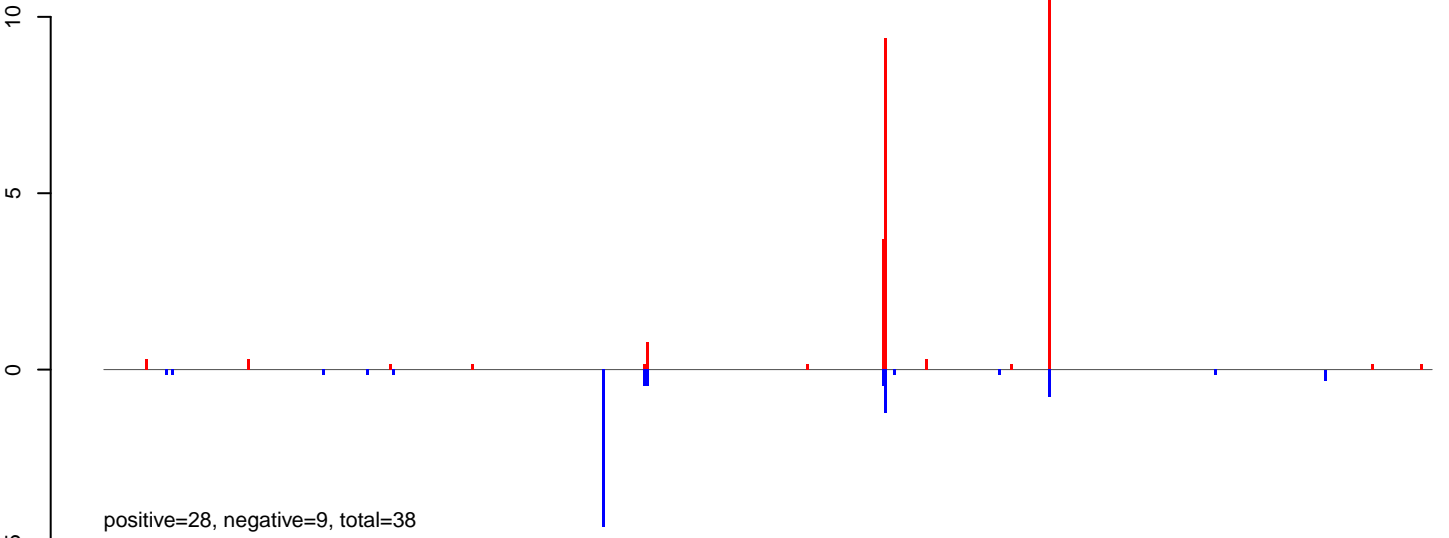
AeAlbo_C636_6dpi_SINV1.rep



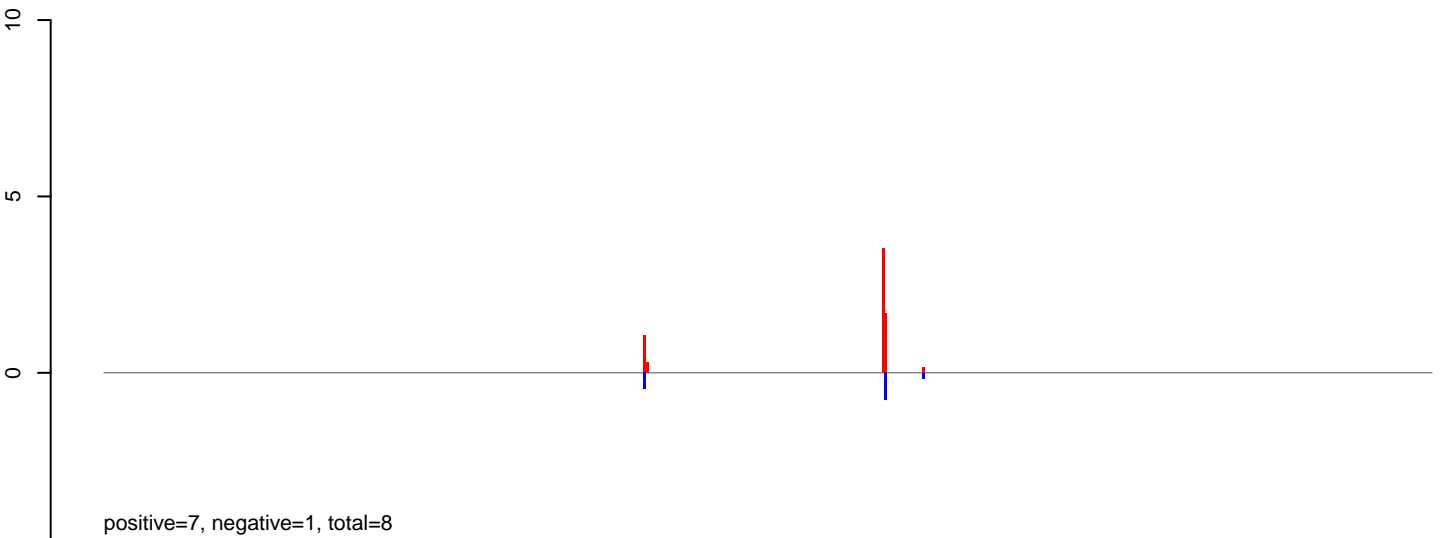
Window size=25, length=10794, virus@NC_012532.1-Zika_virus_isolate_ZIKV_Monkey_Uganda_MR766_1947:1-10794

0 2000 4000 6000 8000 10000

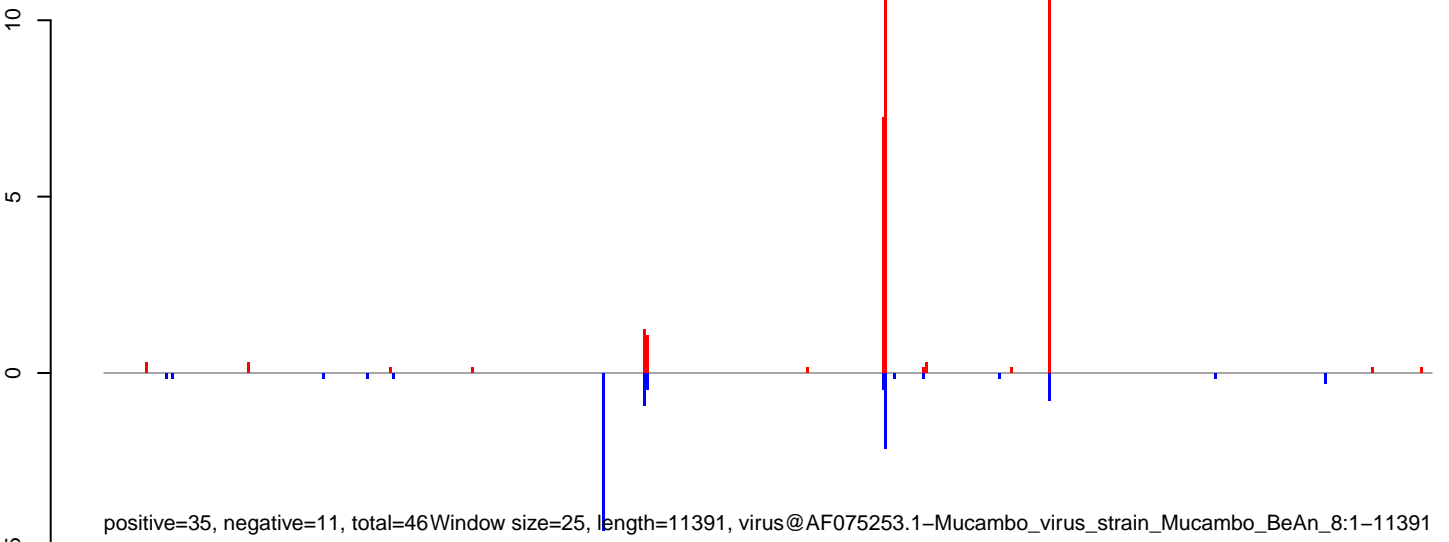
AeAlbo_C636_6dpi_SINV1.18_23.rep



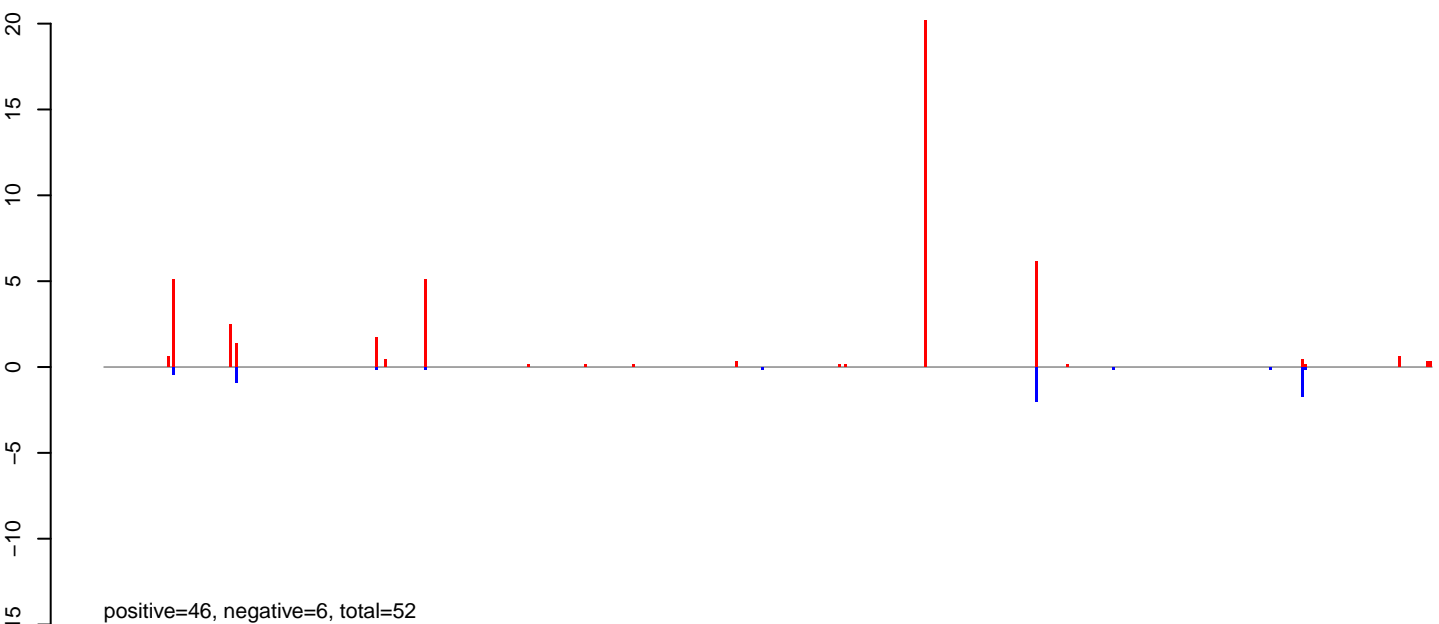
AeAlbo_C636_6dpi_SINV1.24_35.rep



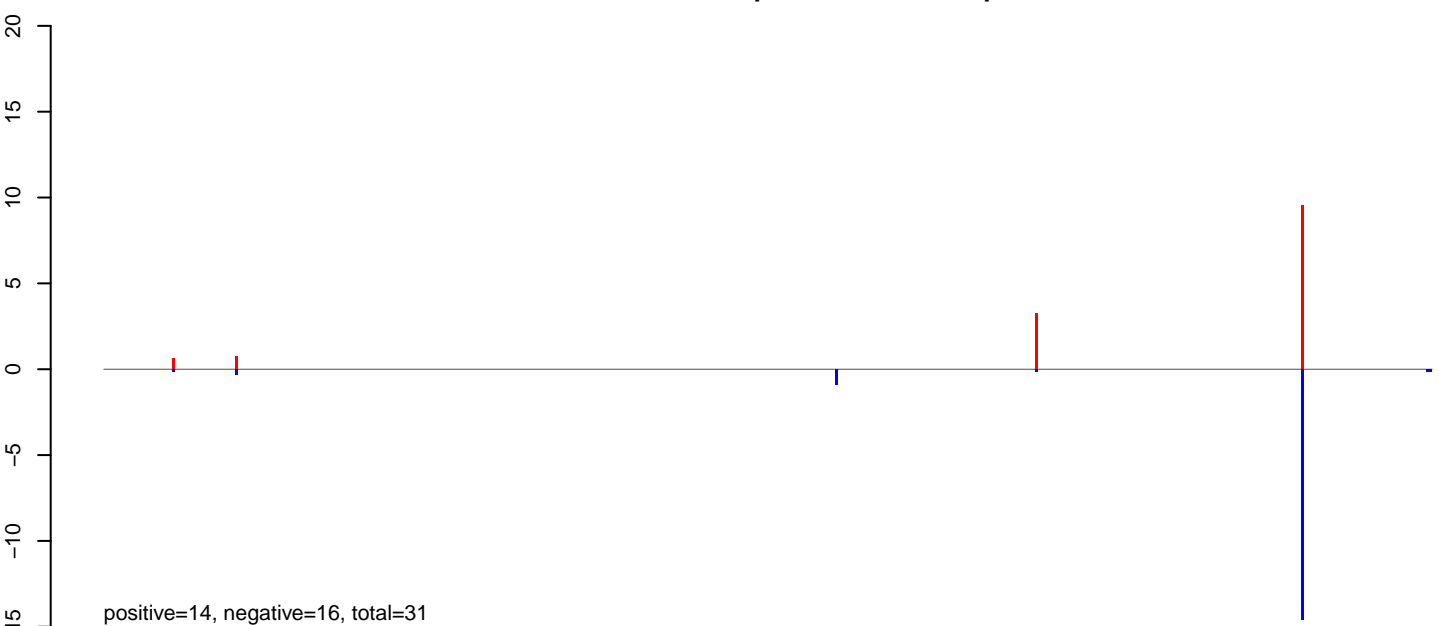
AeAlbo_C636_6dpi_SINV1.rep



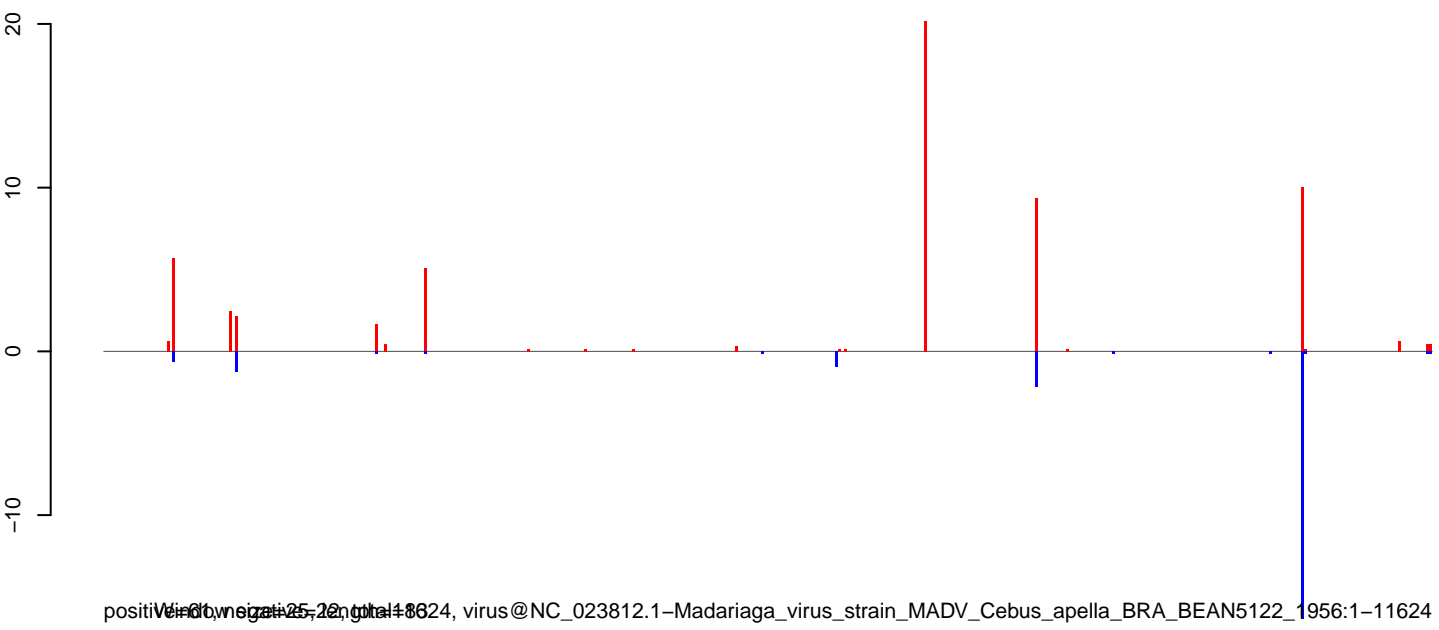
AeAlbo_C636_6dpi_SINV1.18_23.rep



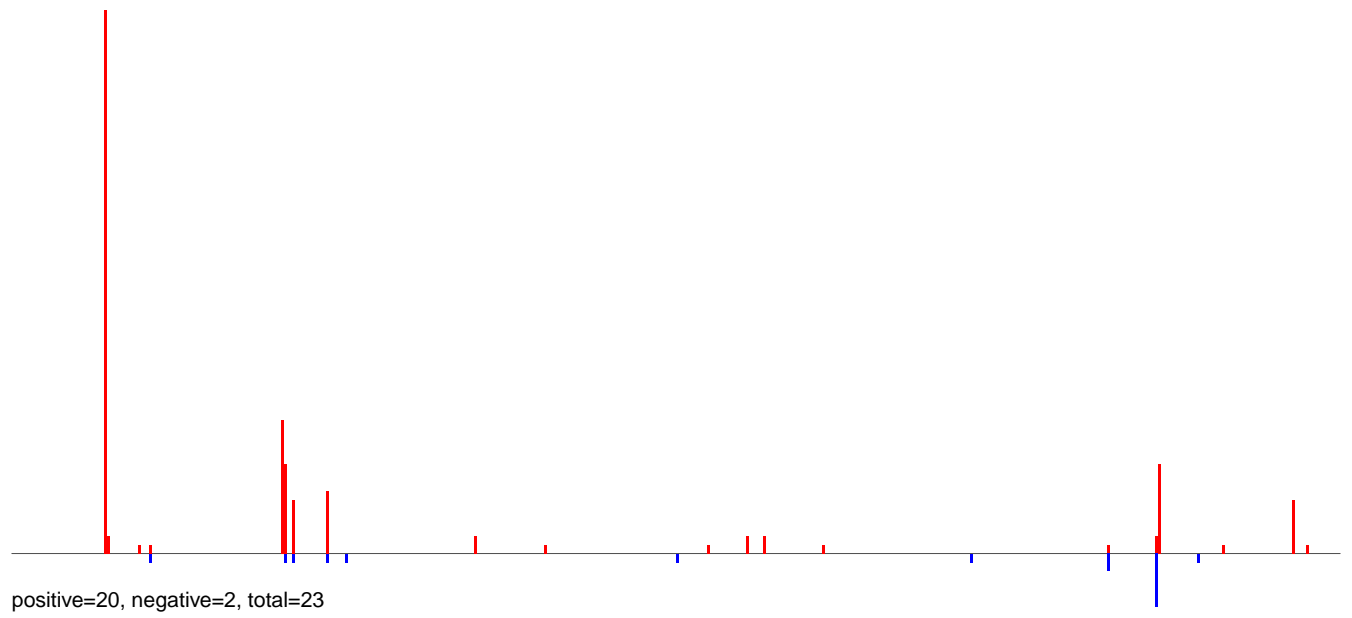
AeAlbo_C636_6dpi_SINV1.24_35.rep



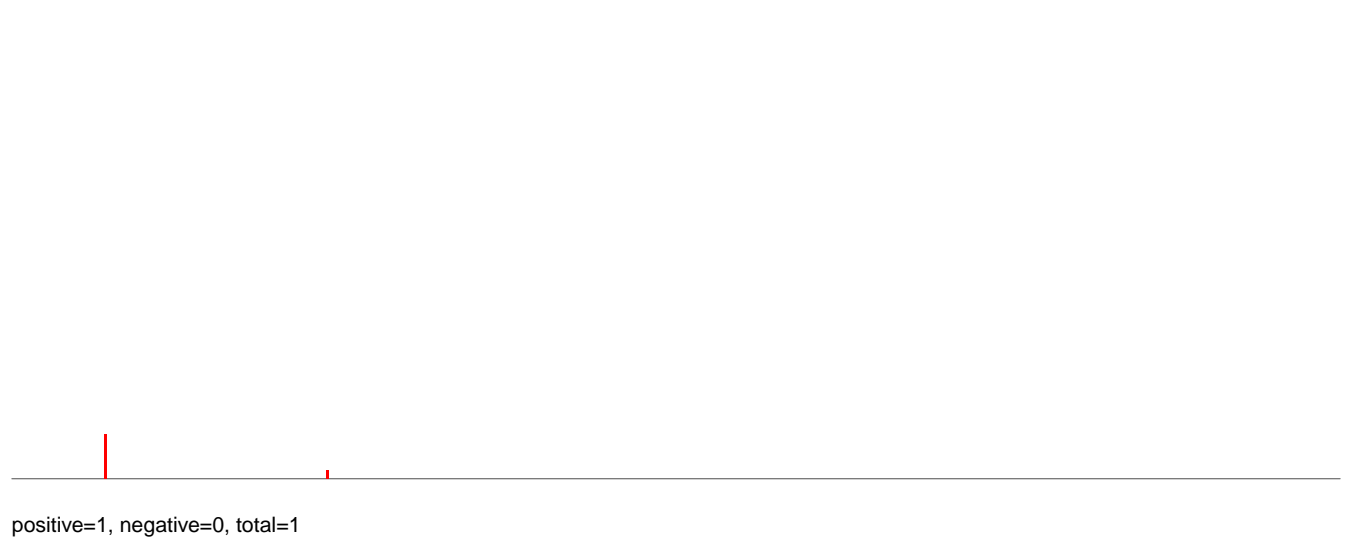
AeAlbo_C636_6dpi_SINV1.rep



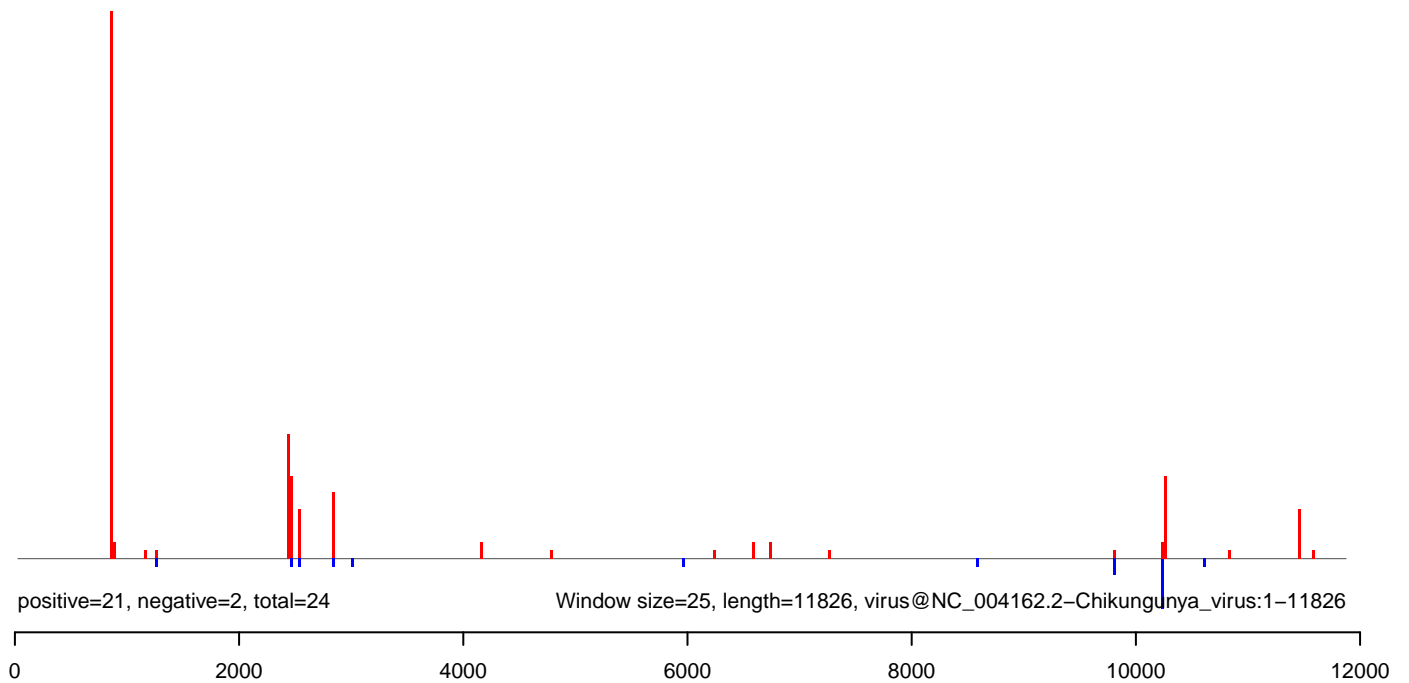
AeAlbo_C636_6dpi_SINV1.18_23.rep



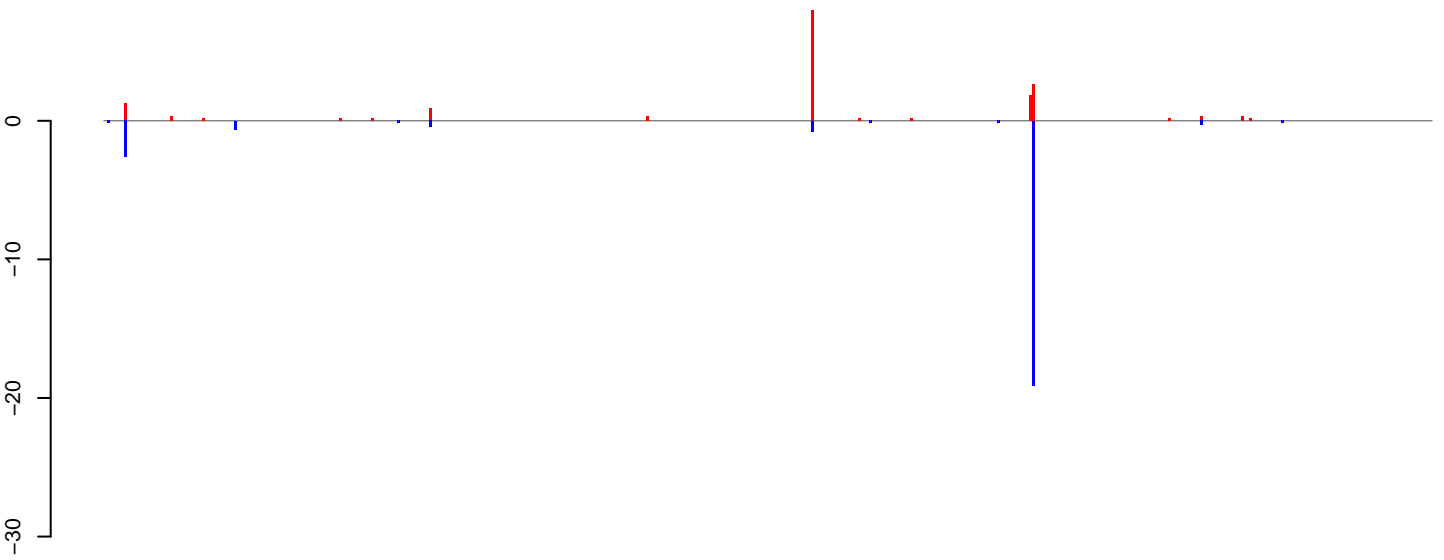
AeAlbo_C636_6dpi_SINV1.24_35.rep



AeAlbo_C636_6dpi_SINV1.rep



AeAlbo_C636_6dpi_SINV1.18_23.rep



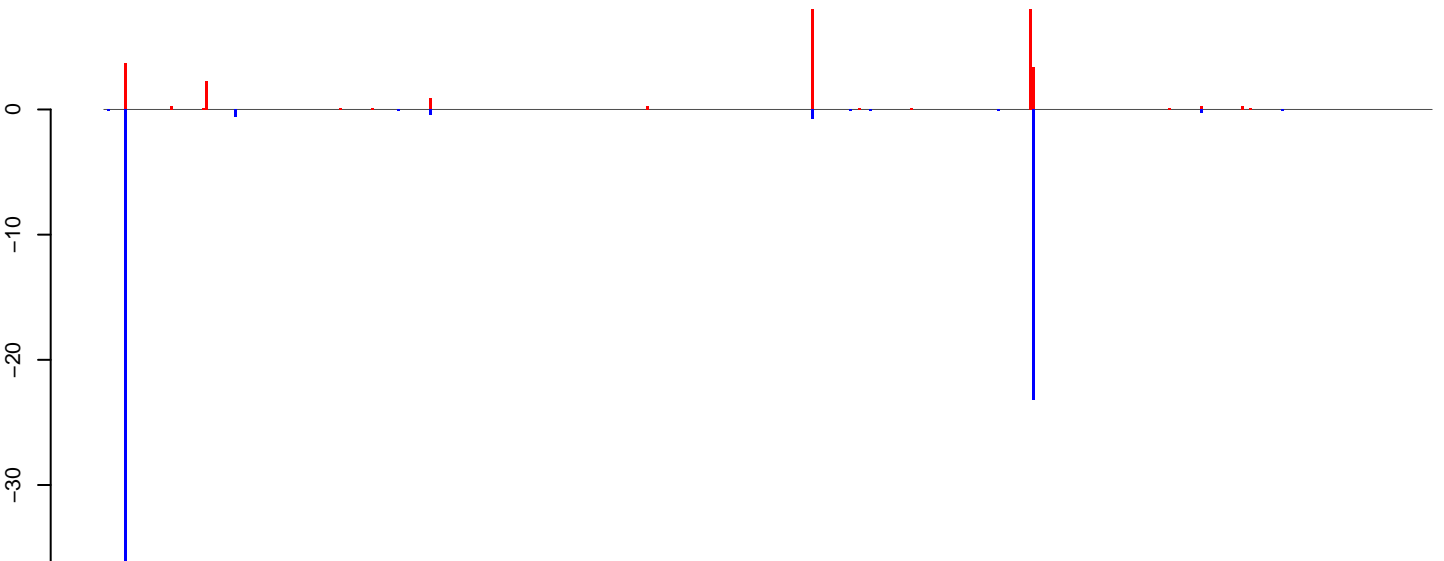
positive=17, negative=25, total=42

AeAlbo_C636_6dpi_SINV1.24_35.rep



positive=12, negative=39, total=51

AeAlbo_C636_6dpi_SINV1.rep

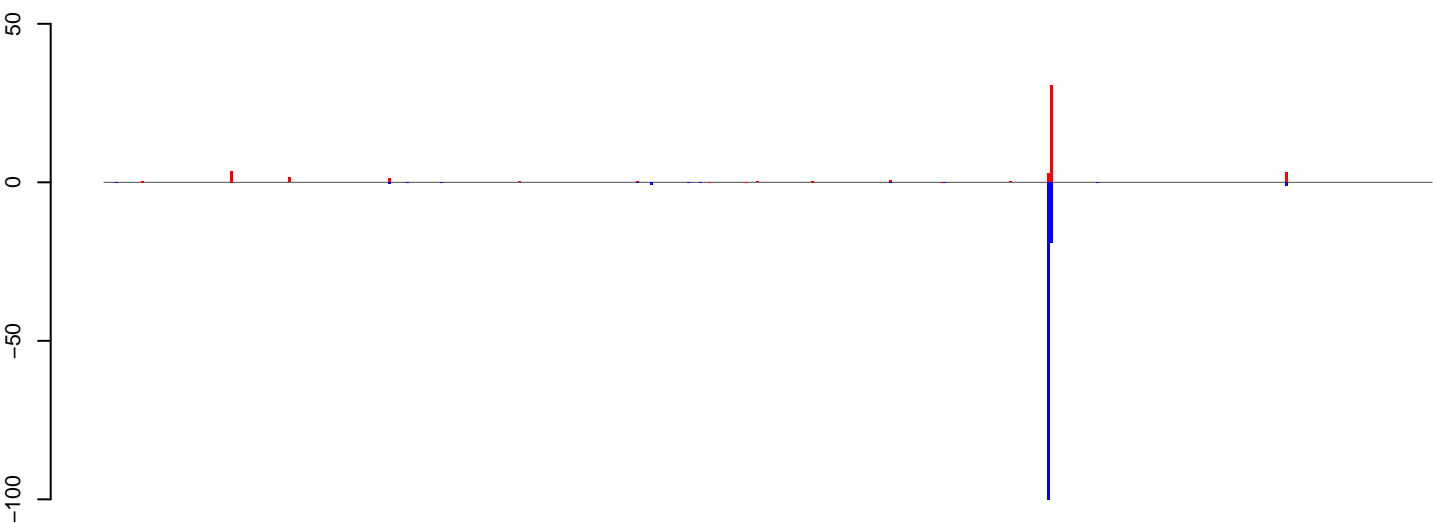


positive=29, negative=66, total=95

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

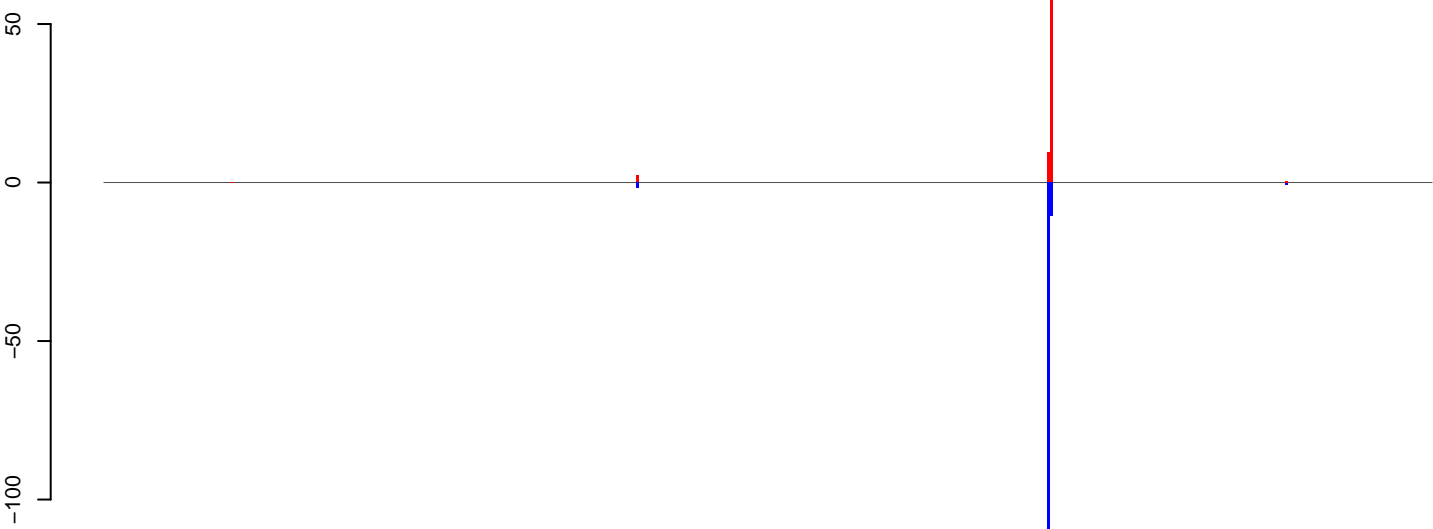
0 2000 4000 6000 8000 10000

AeAlbo_C636_6dpi_SINV1.18_23.rep



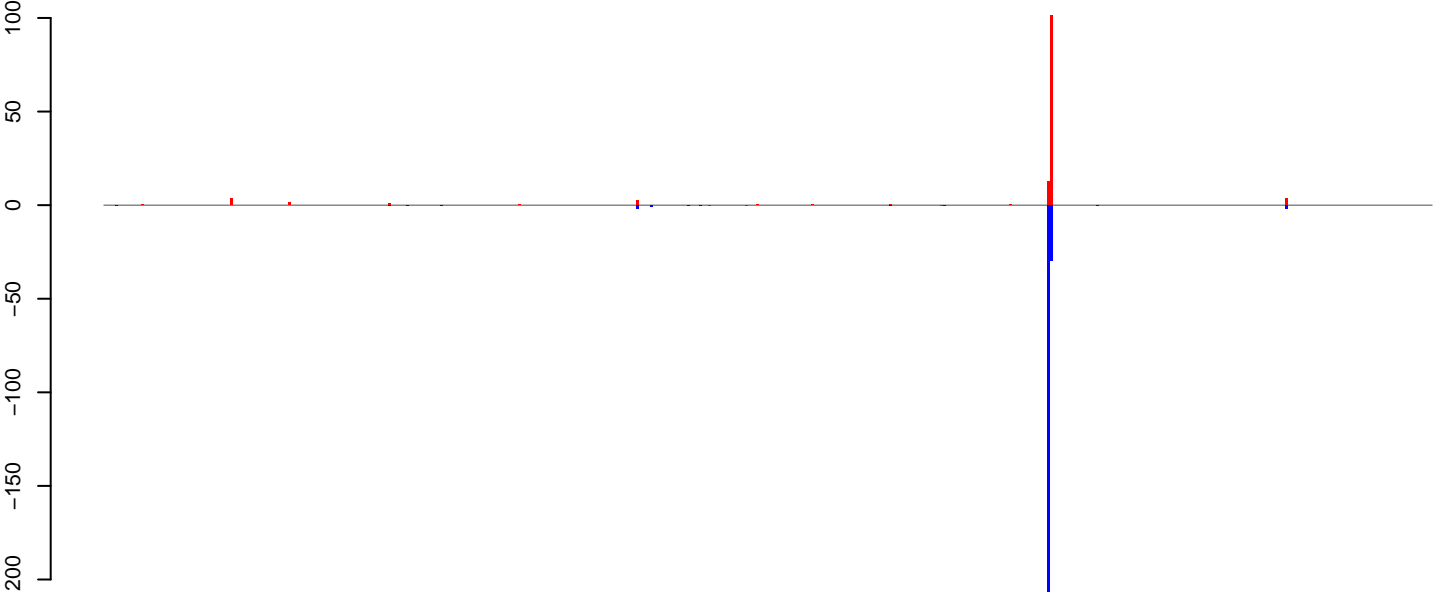
positive=46, negative=123, total=170

AeAlbo_C636_6dpi_SINV1.24_35.rep



positive=84, negative=130, total=214

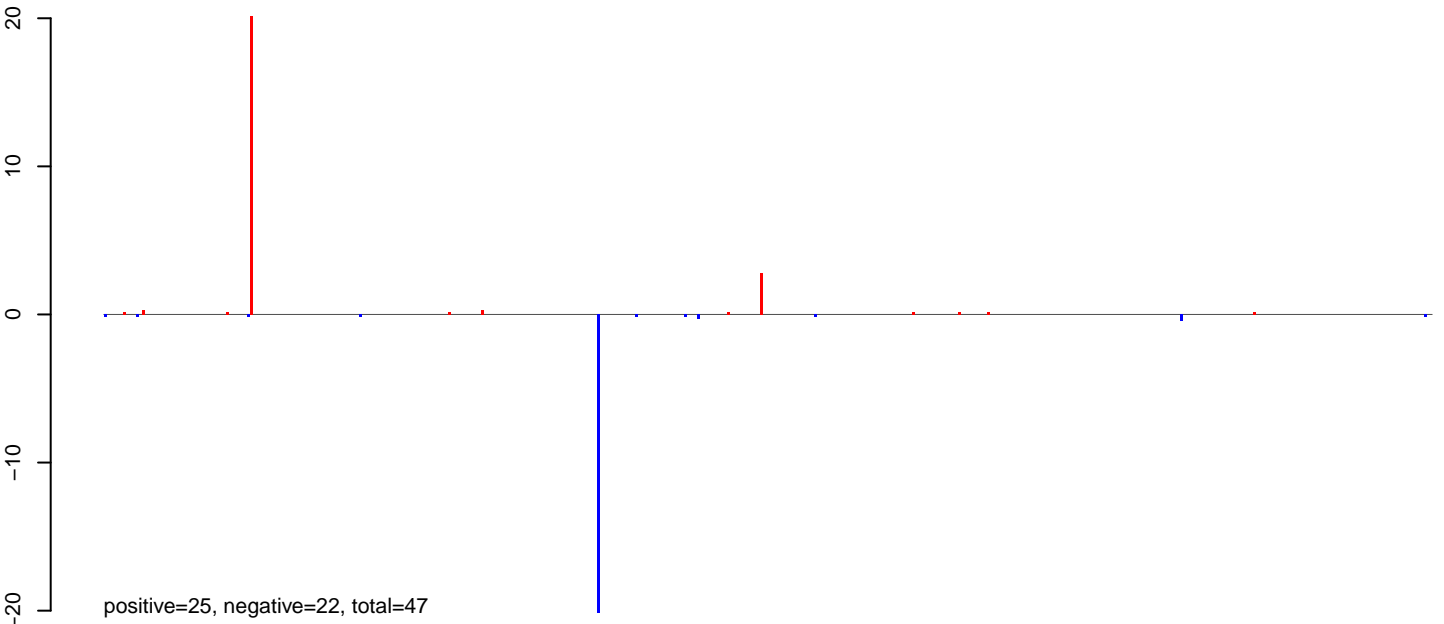
AeAlbo_C636_6dpi_SINV1.rep



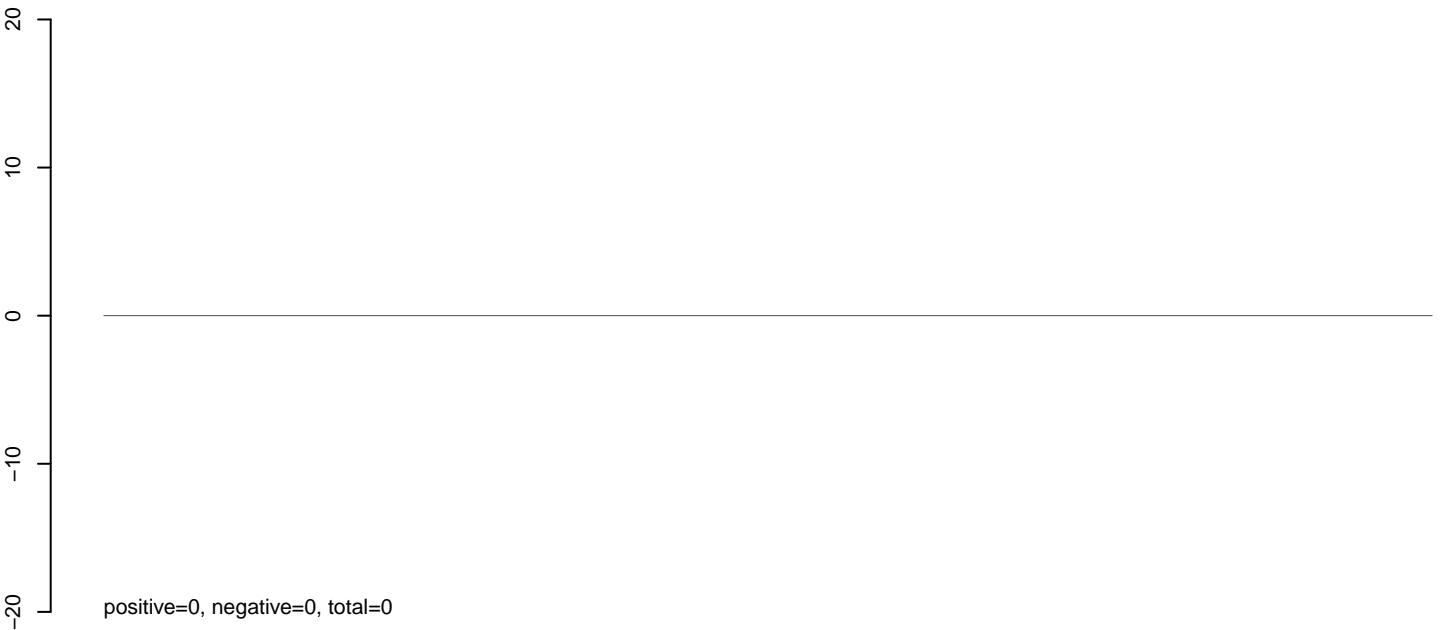
positive=130, negative=254, total=384, window size=25, length=11530, virus@AF075254.1-Tonate_virus_strain_Tonate_CaAn_410d:1-11530

0 2000 4000 6000 8000 10000 12000

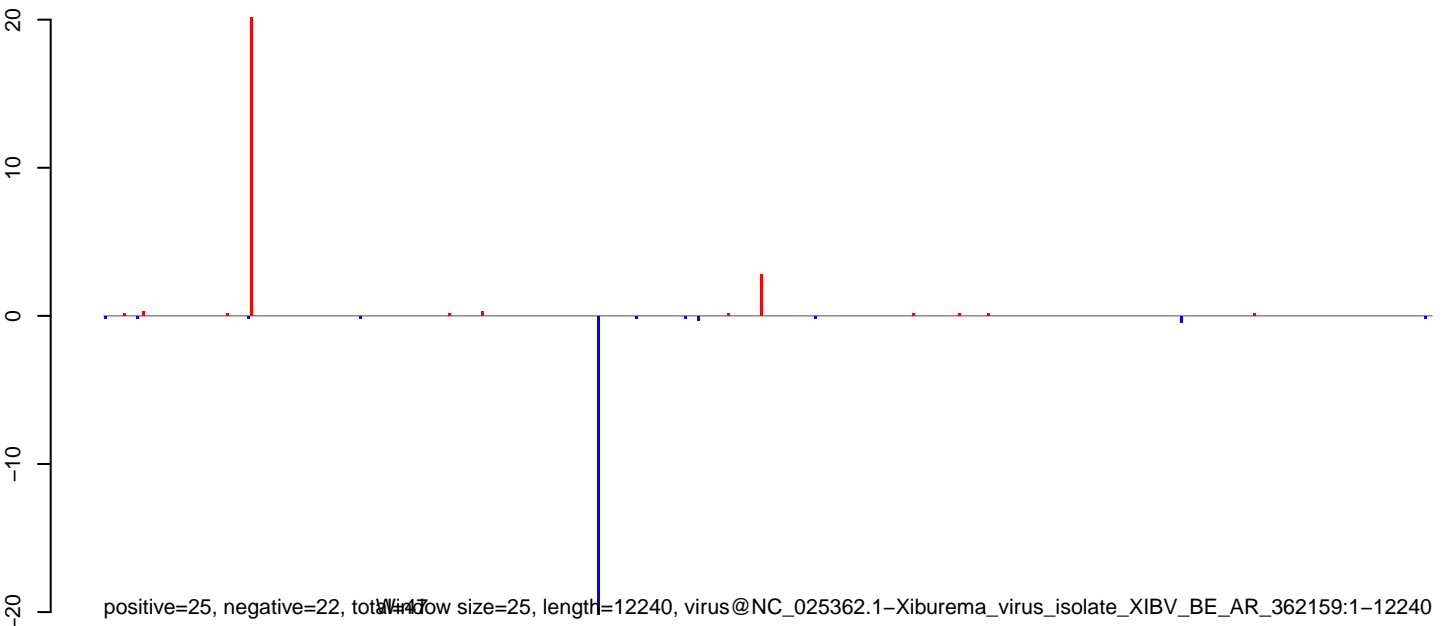
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



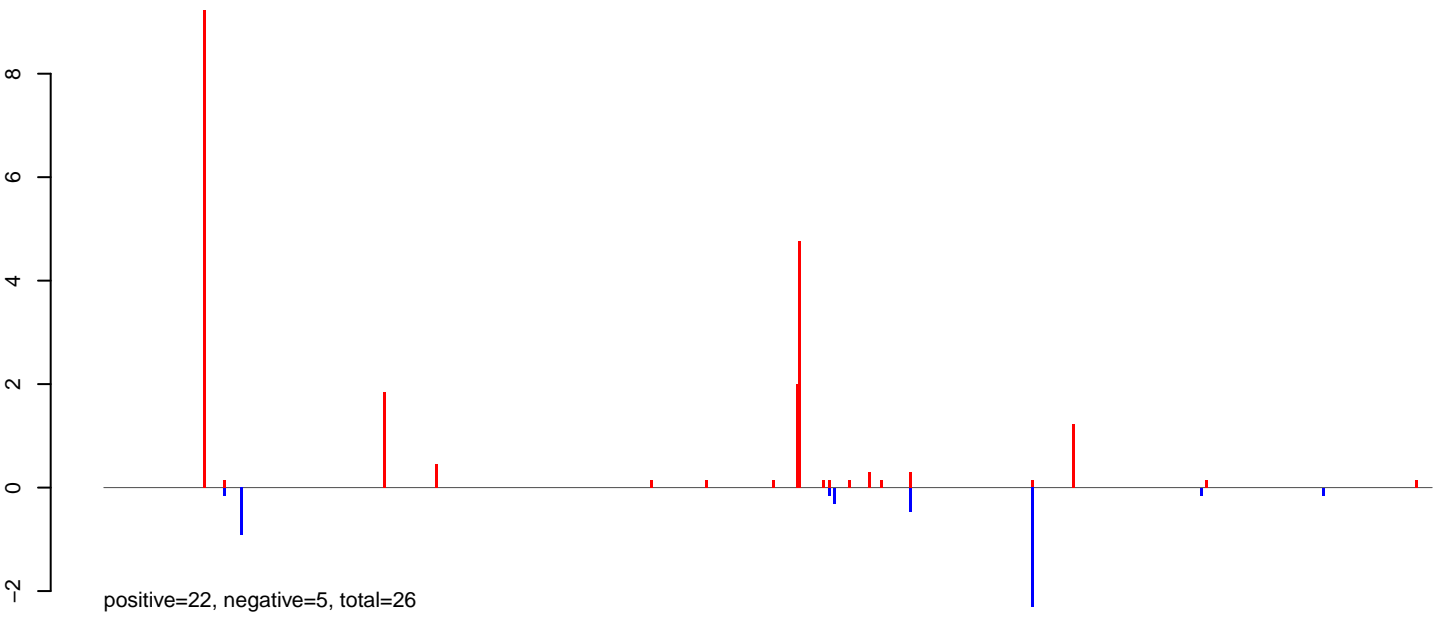
AeAlbo_C636_6dpi_SINV1.rep



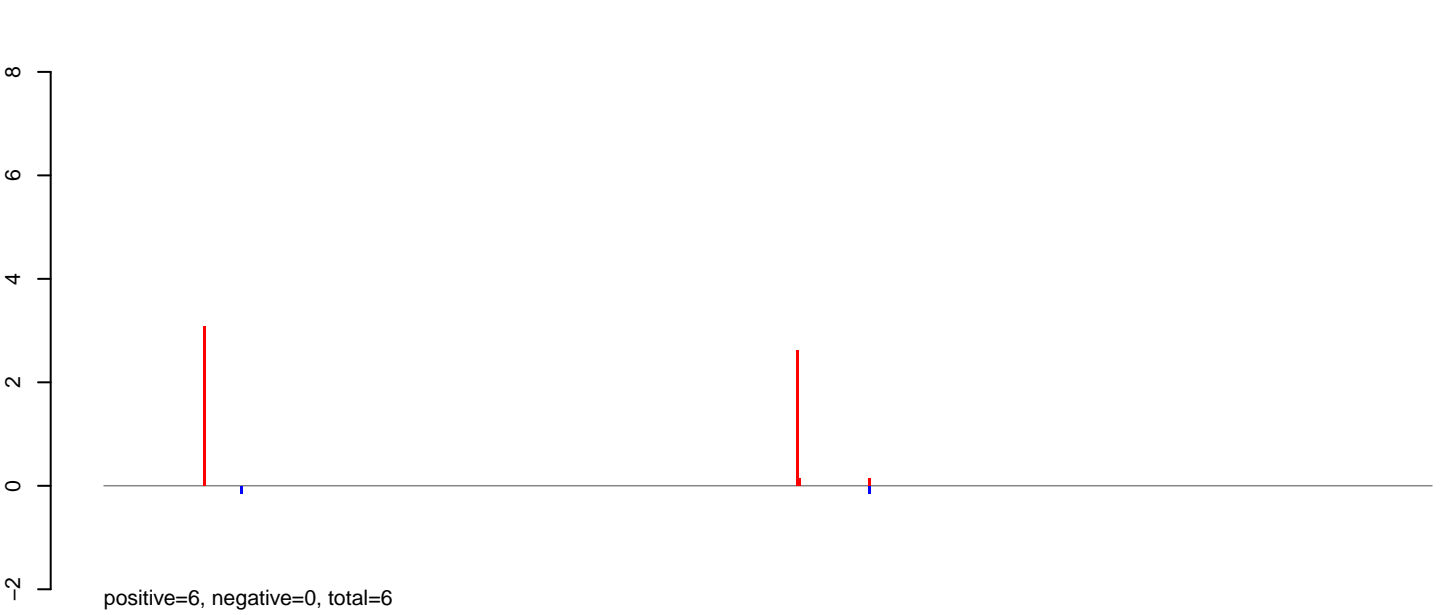
Window size=25, length=12240, virus@NC_025362.1-Xiburema_virus_isolate_XIBV_BE_AR_362159:1-12240

0 2000 4000 6000 8000 10000 12000

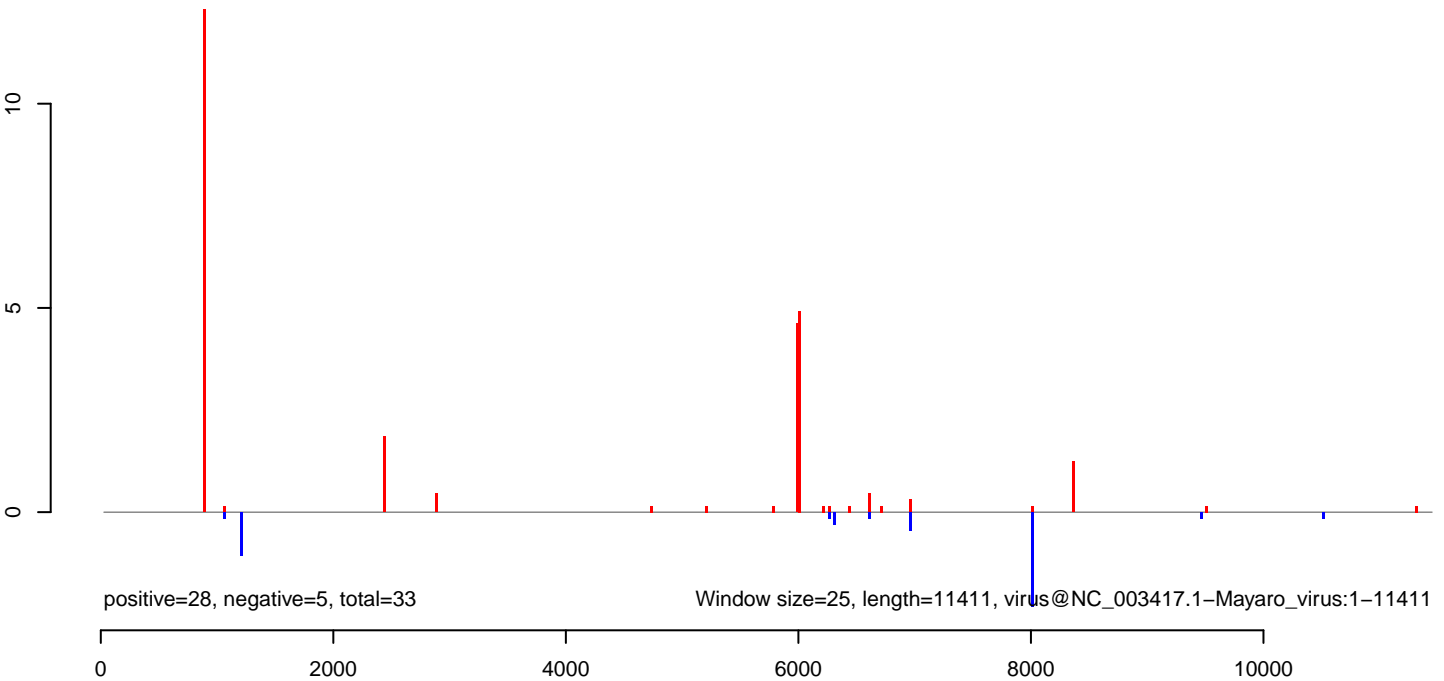
AeAlbo_C636_6dpi_SINV1.18_23.rep



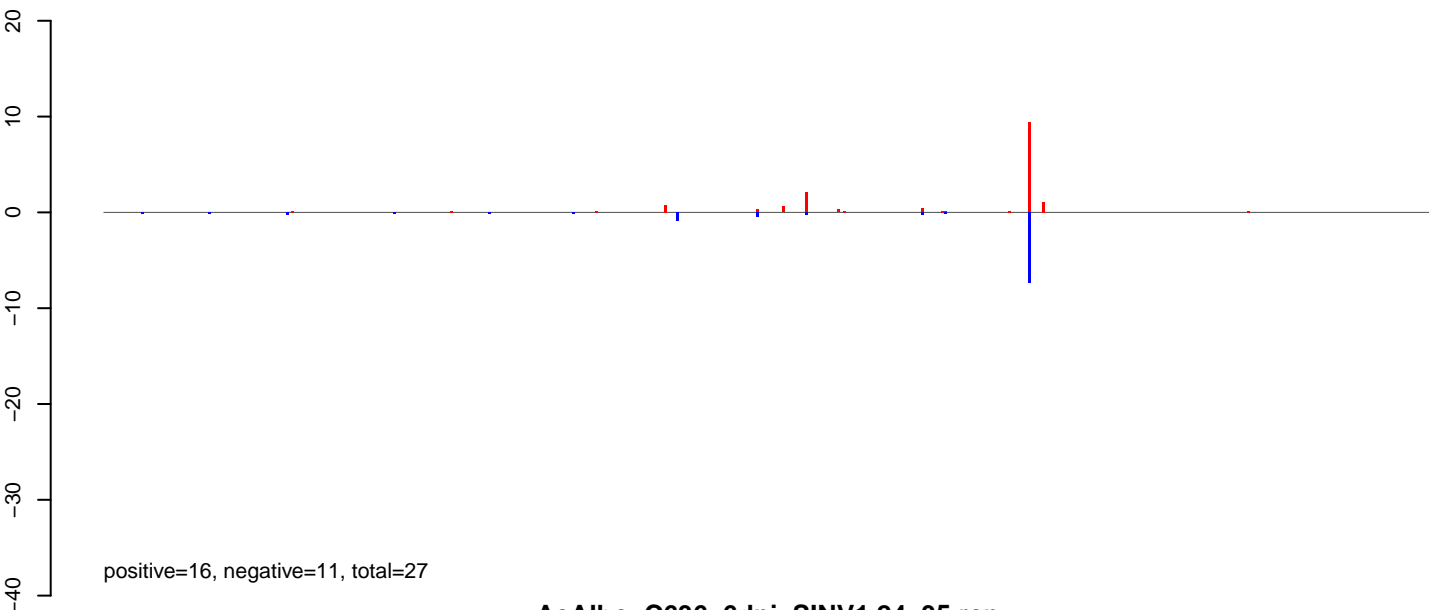
AeAlbo_C636_6dpi_SINV1.24_35.rep



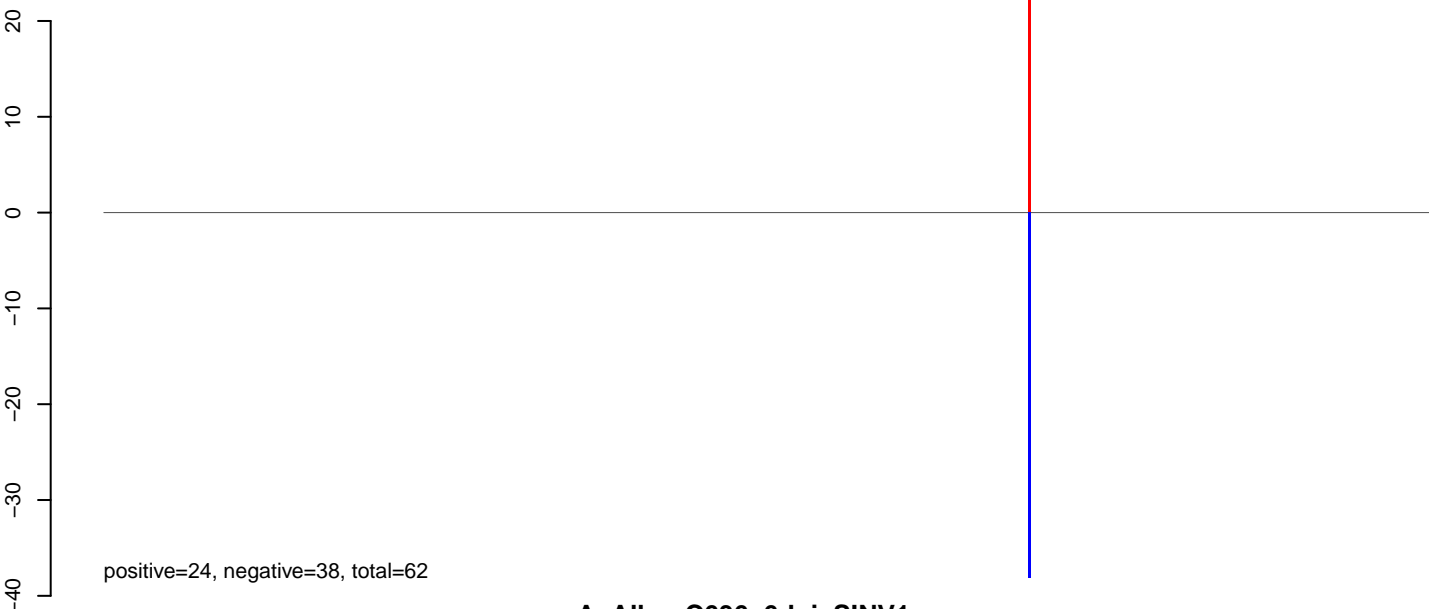
AeAlbo_C636_6dpi_SINV1.rep



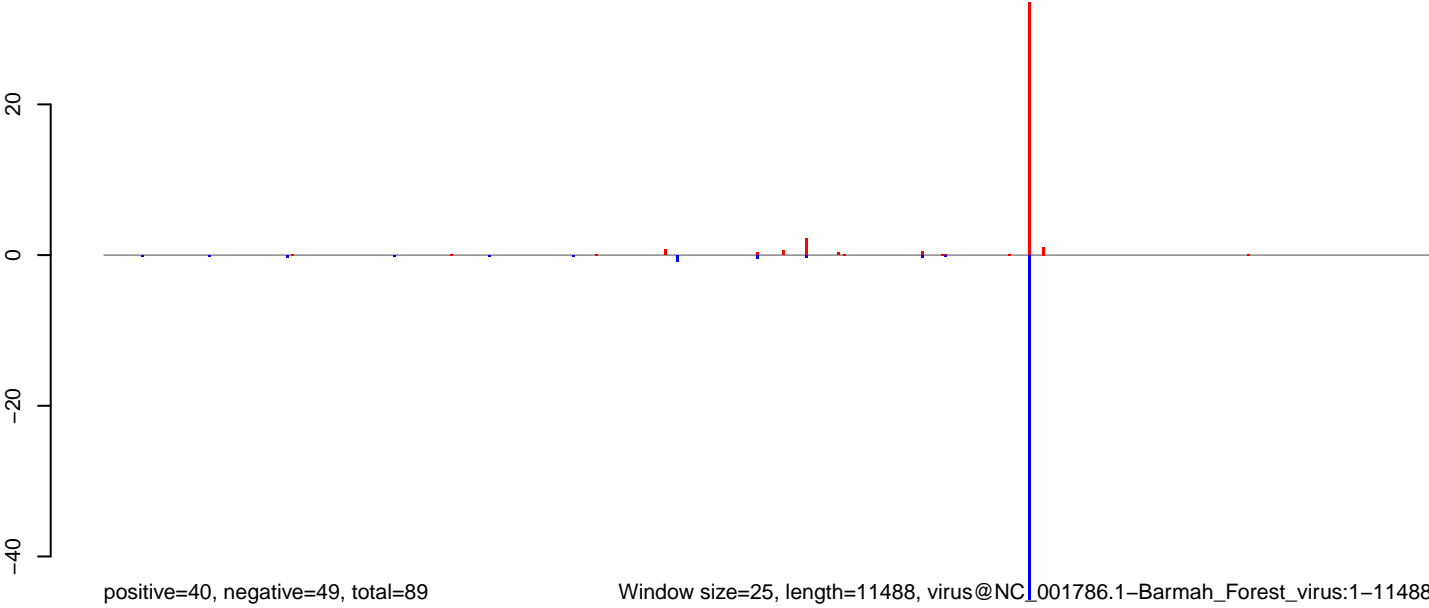
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



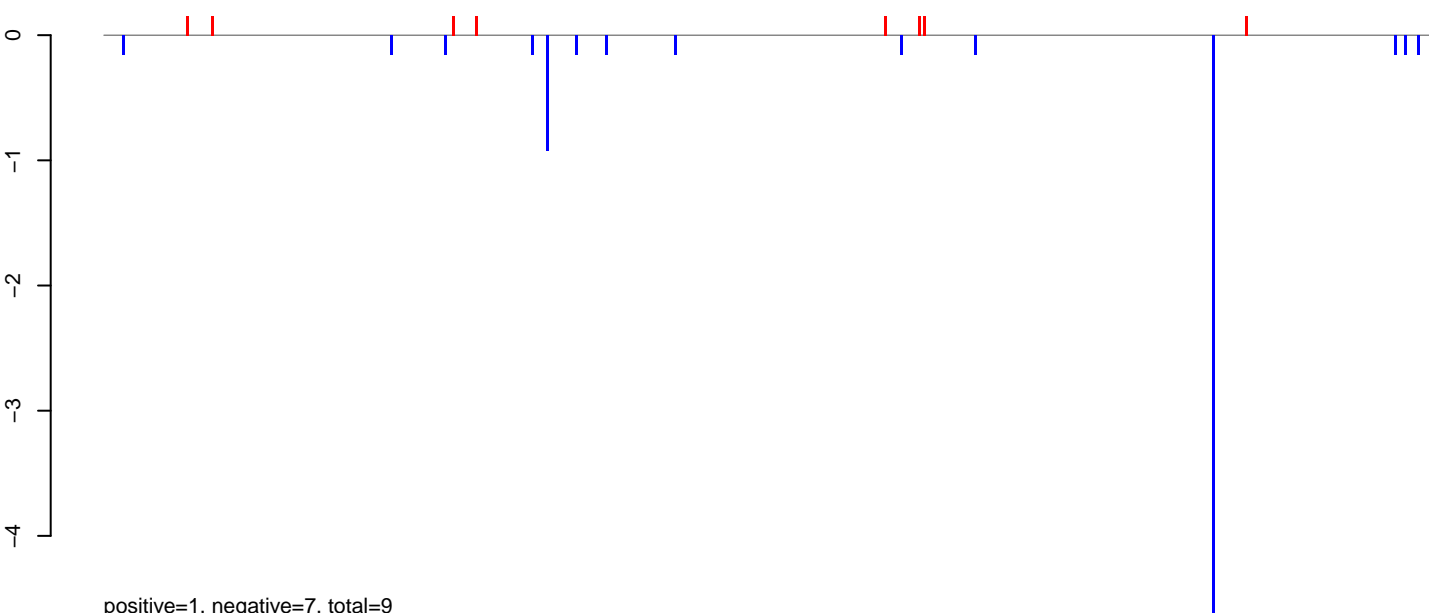
AeAlbo_C636_6dpi_SINV1.rep



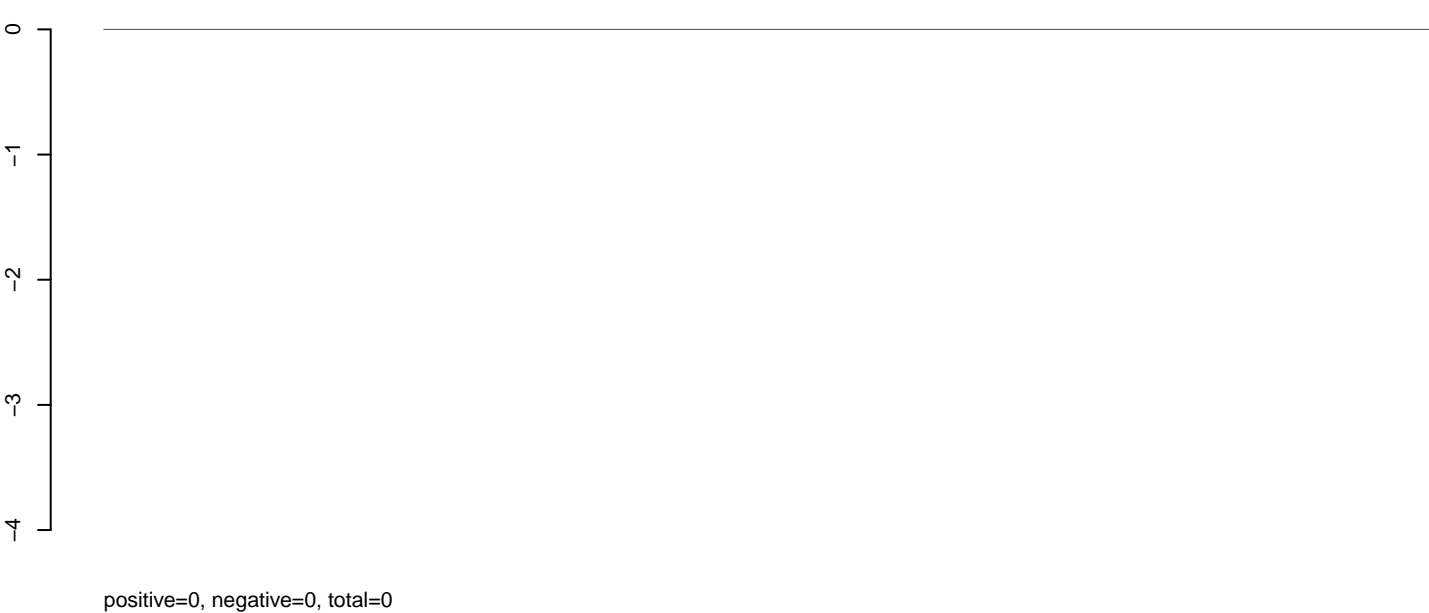
Window size=25, length=11488, virus@NC_001786.1-Barmah_Forest_virus:1-11488

0 2000 4000 6000 8000 10000

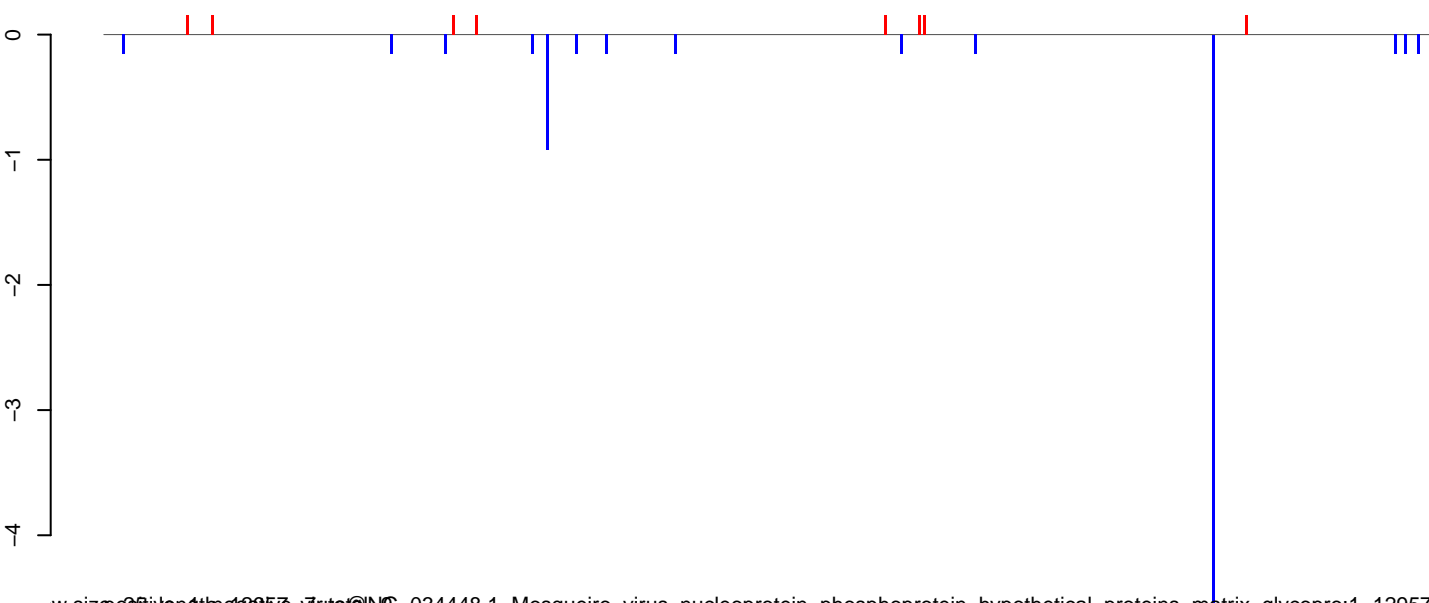
AeAlbo_C636_6dpi_SINV1.18_23.rep



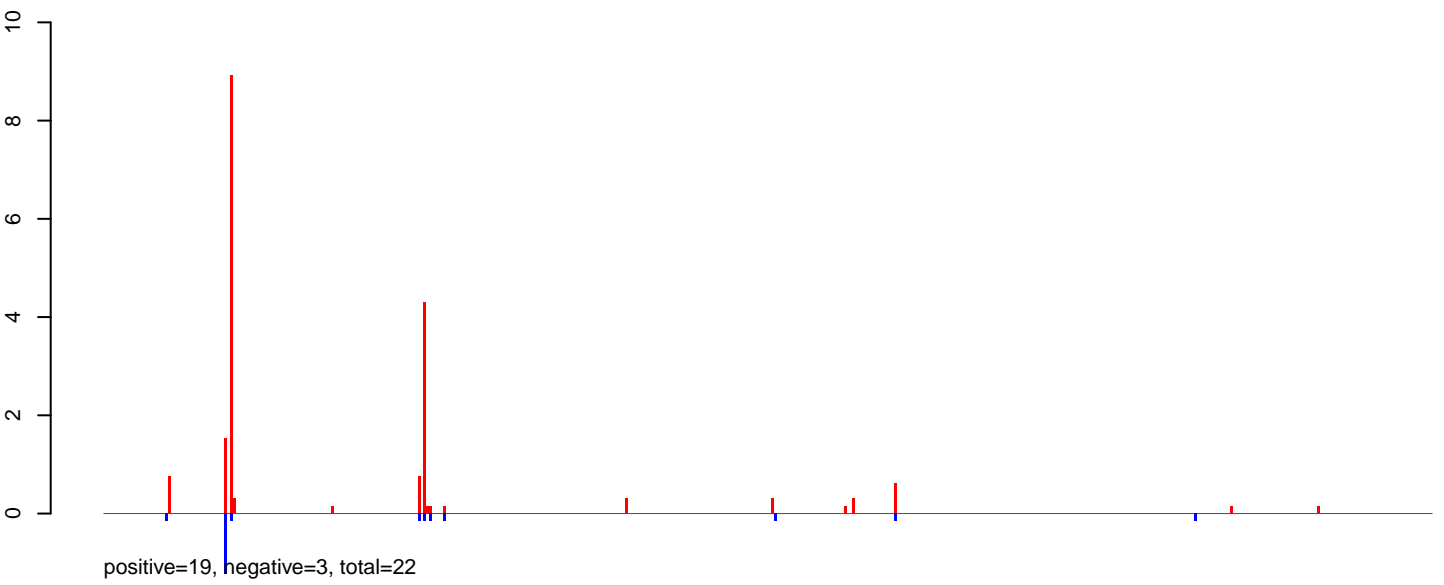
AeAlbo_C636_6dpi_SINV1.24_35.rep



AeAlbo_C636_6dpi_SINV1.rep



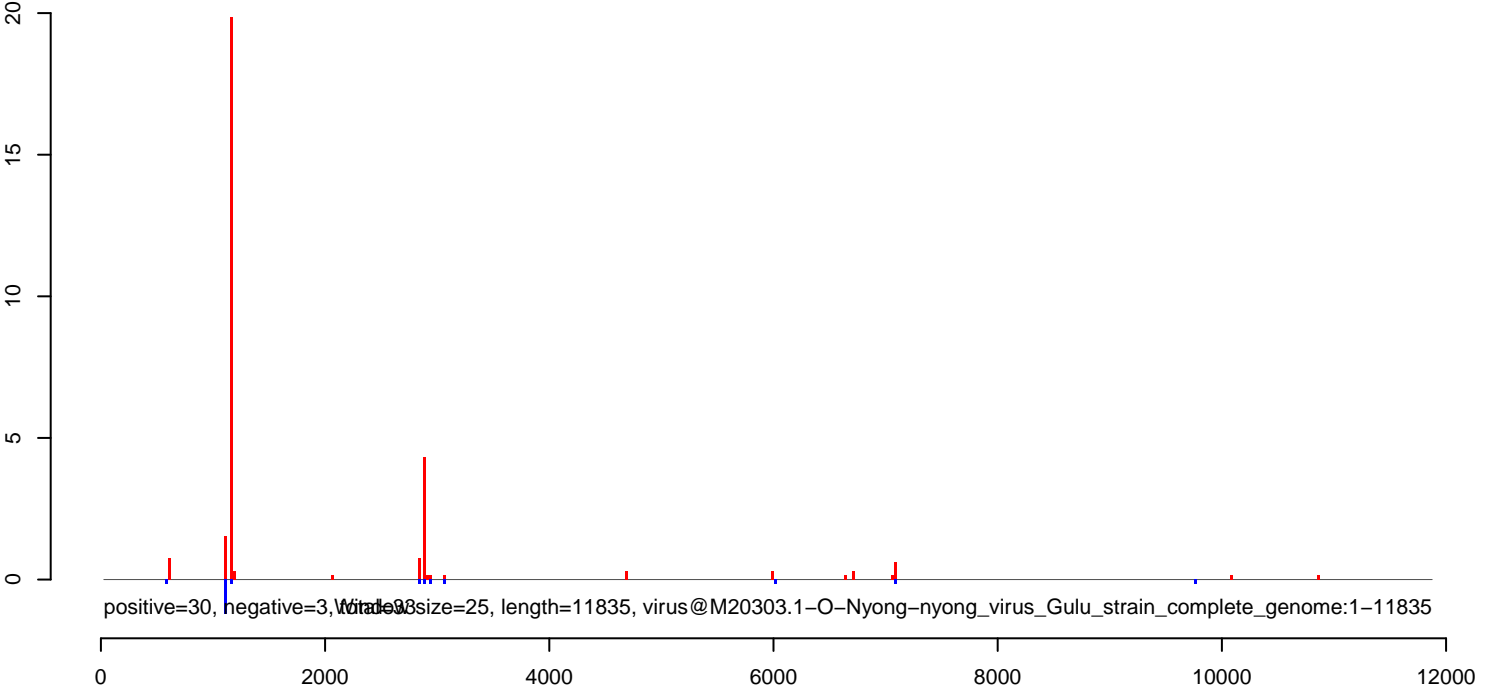
AeAlbo_C636_6dpi_SINV1.18_23.rep



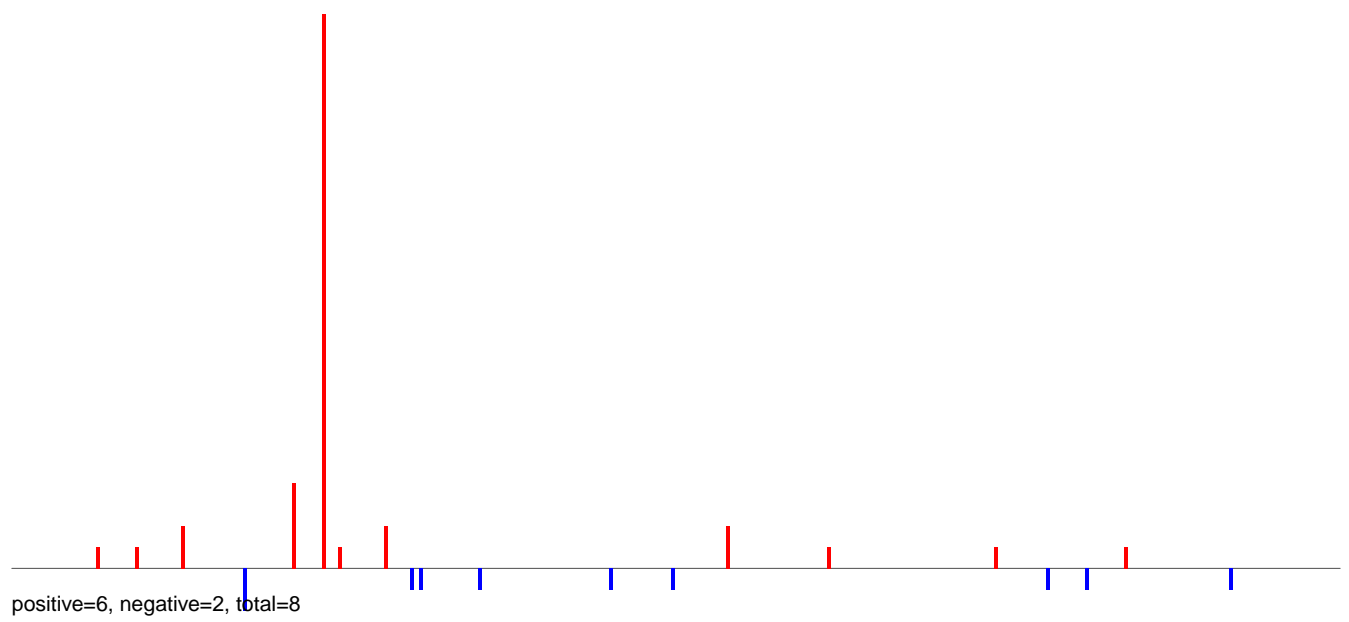
AeAlbo_C636_6dpi_SINV1.24_35.rep



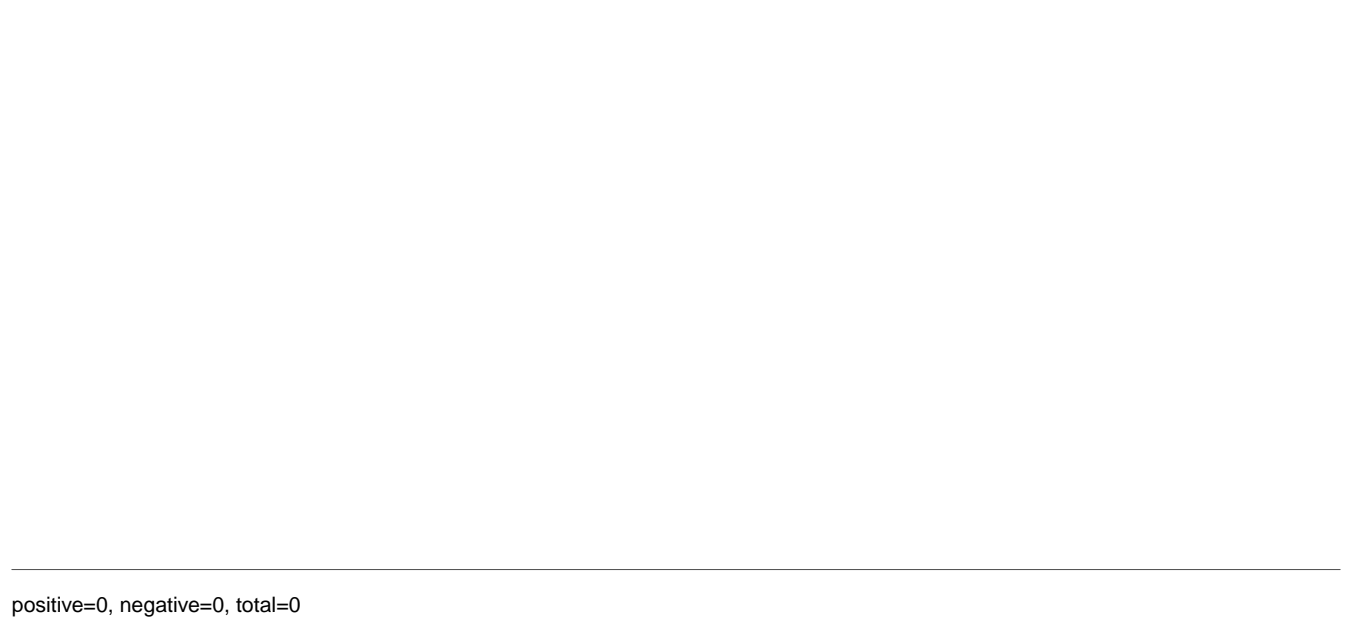
AeAlbo_C636_6dpi_SINV1.rep



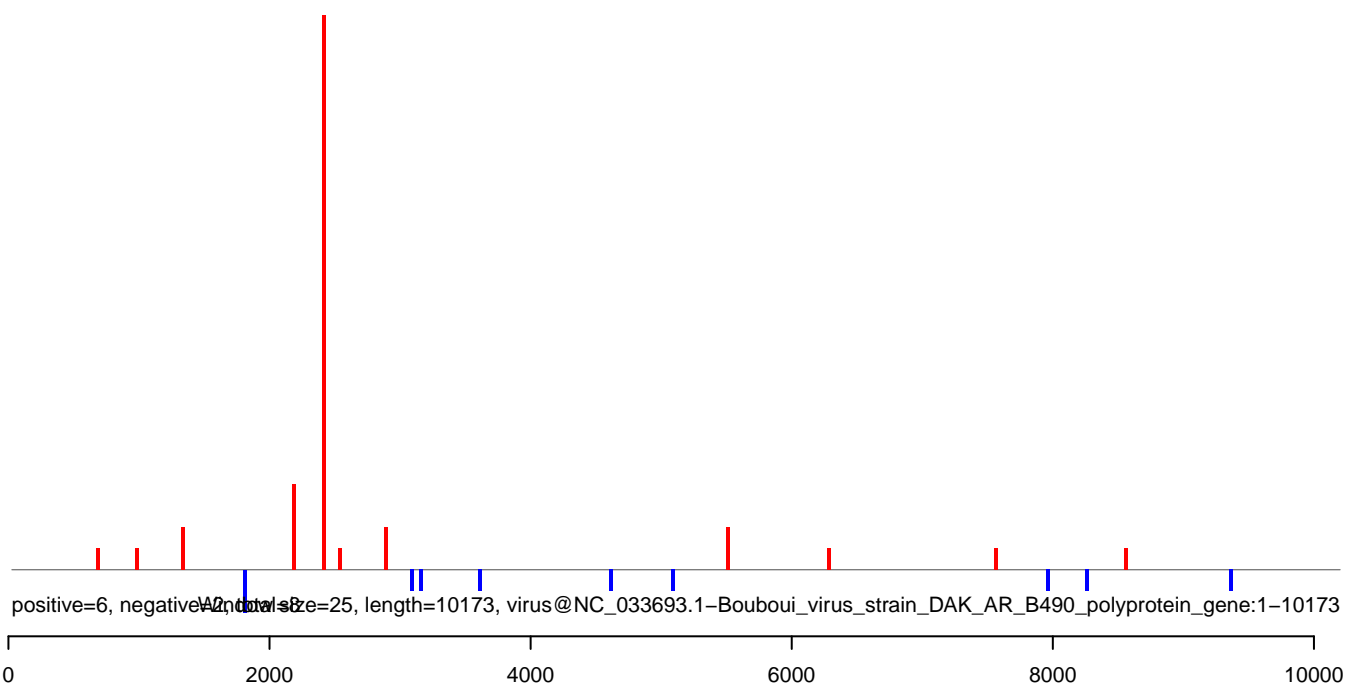
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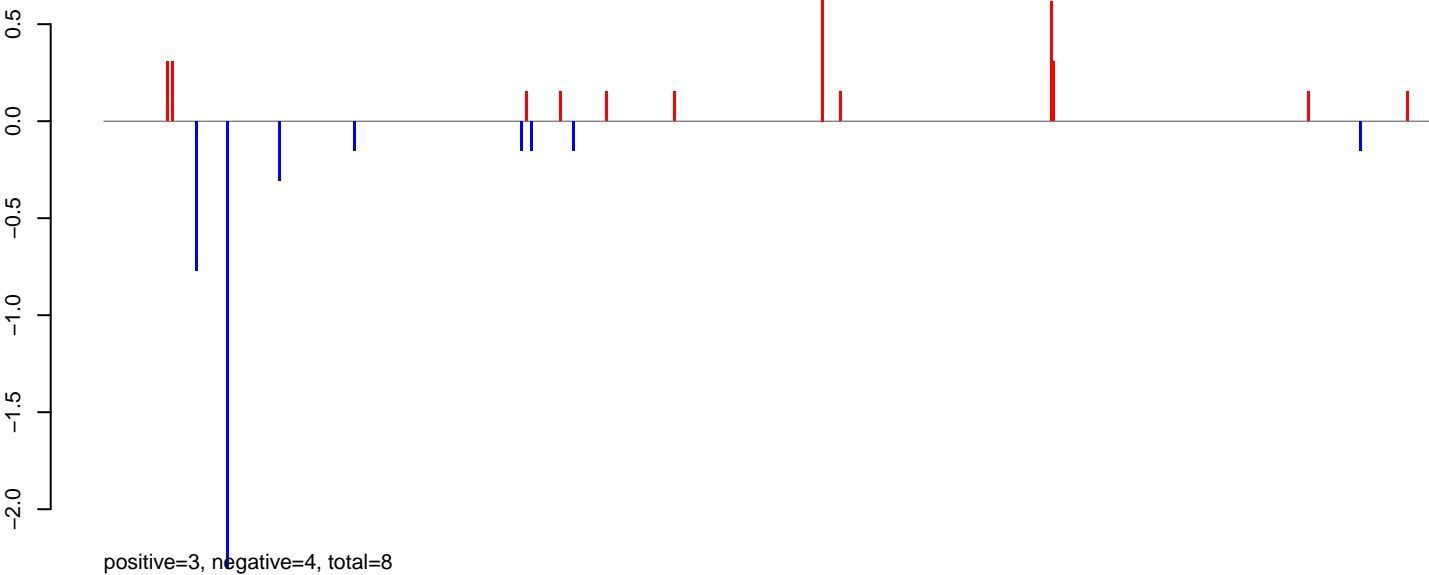
AeAlbo_C636_6dpi_SINV1.24_35.rep



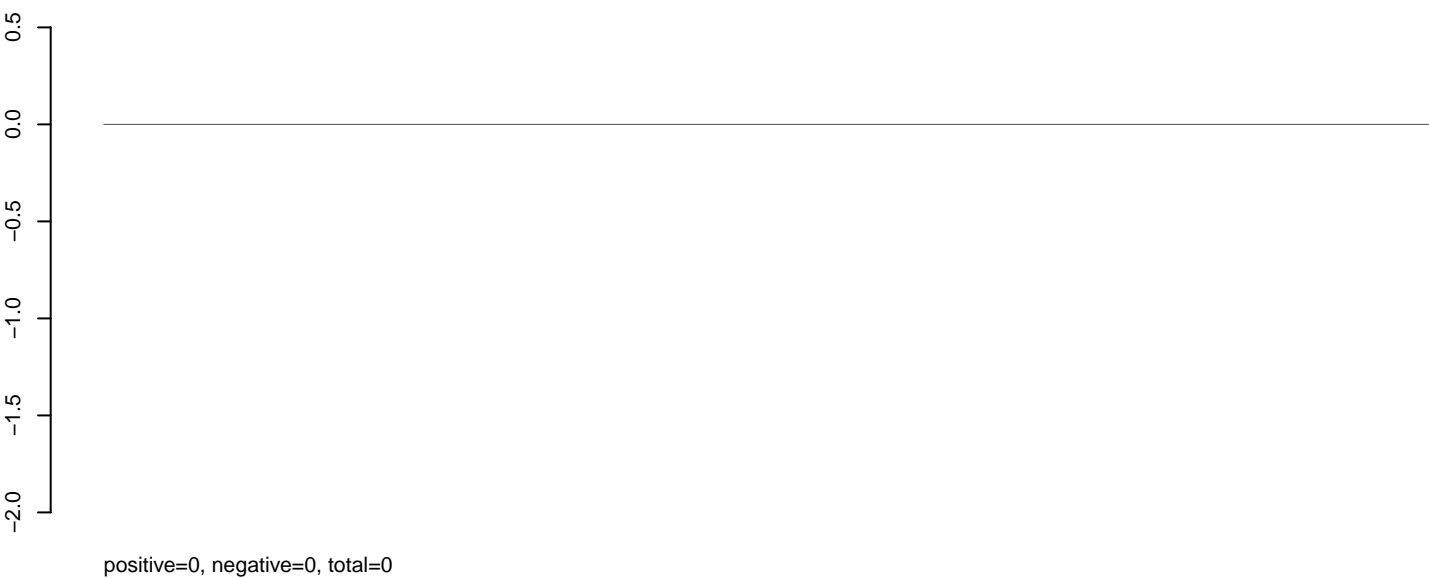
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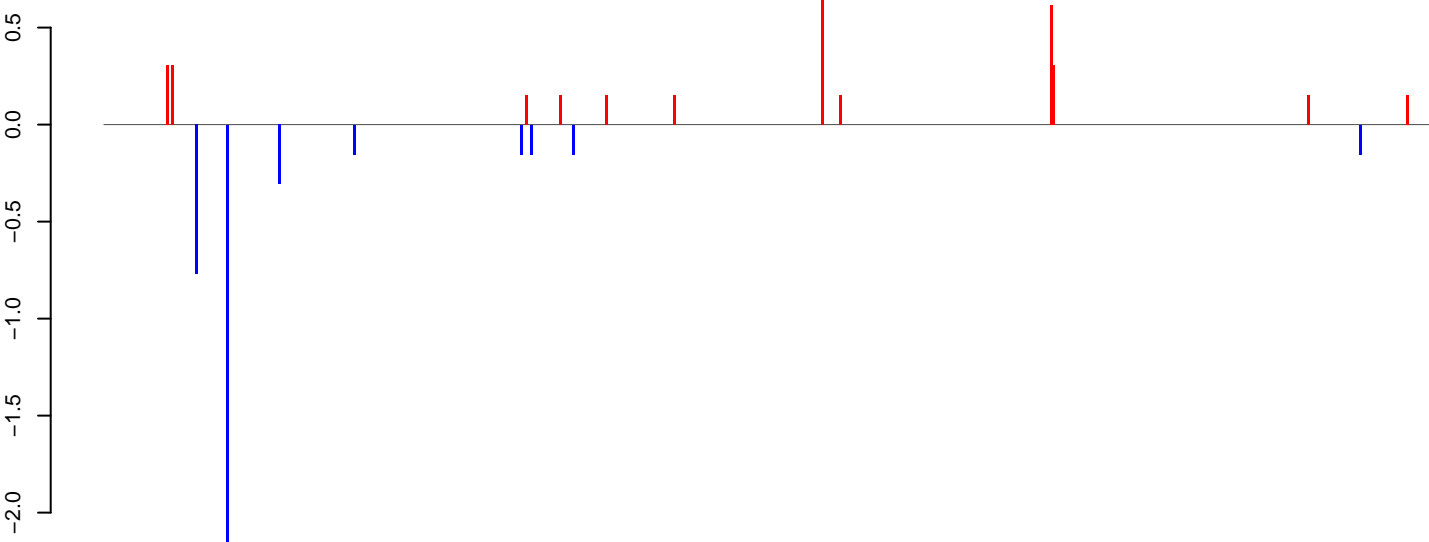
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



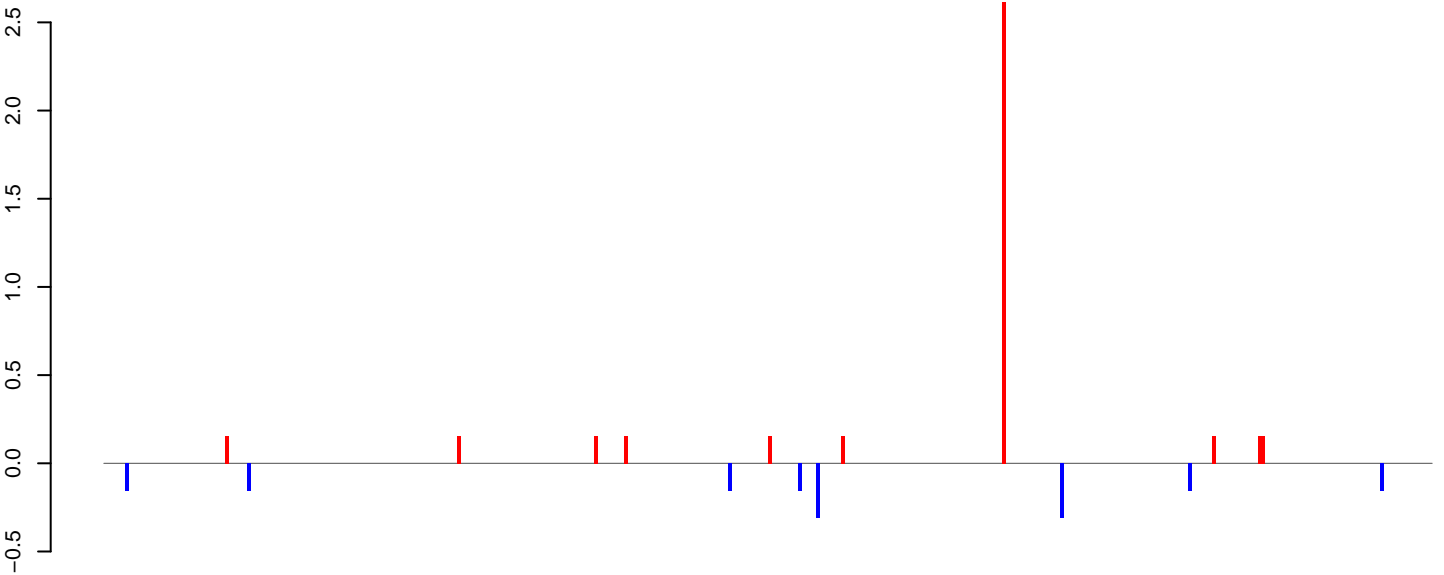
AeAlbo_C636_6dpi_SINV1.rep



length=12774, coverage=0.942, total=8, Xincheng_Mosquito_Virus_strain_XC1-6_ORF1_ORF1_ORF2_ORF2_glycoprotein_G_and_RNA-de:1-12774

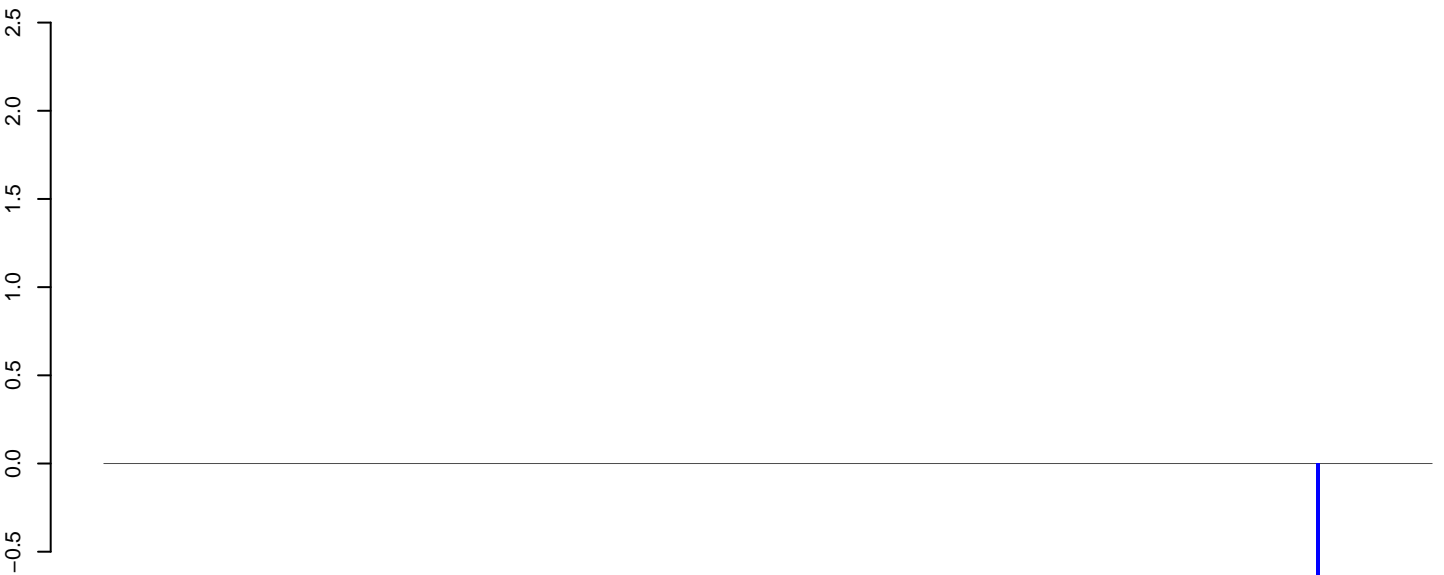
0 2000 4000 6000 8000 10000 12000

AeAlbo_C636_6dpi_SINV1.18_23.rep



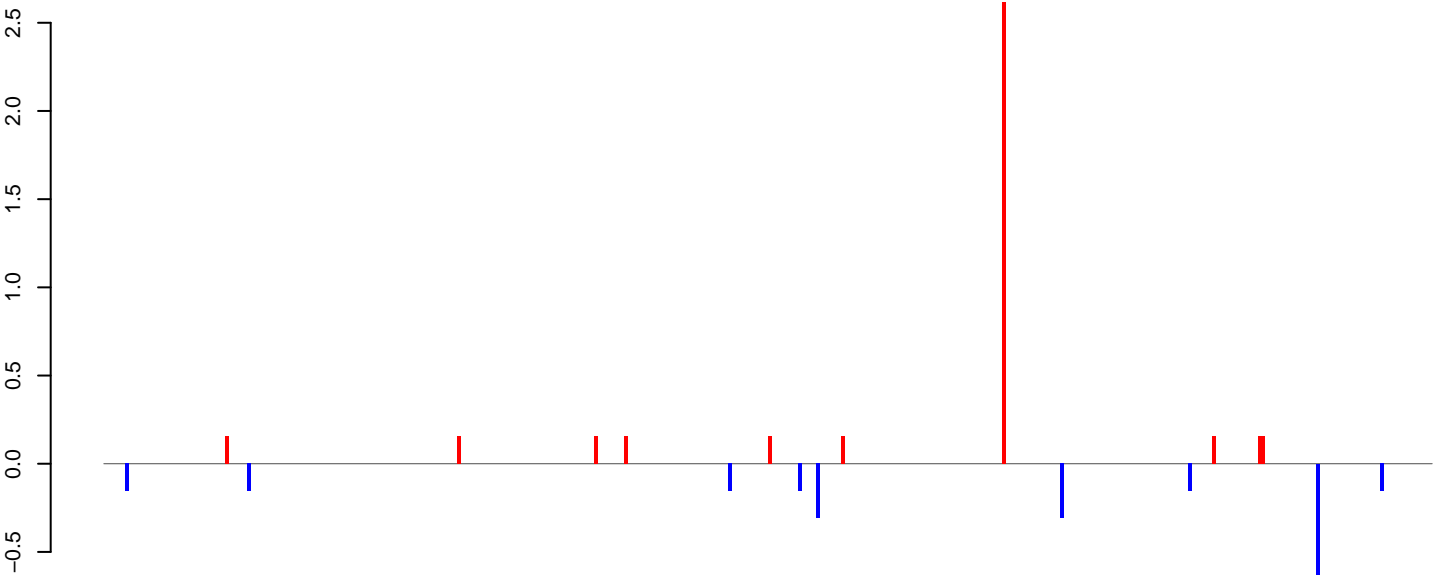
positive=4, negative=2, total=6

AeAlbo_C636_6dpi_SINV1.24_35.rep



positive=0, negative=1, total=1

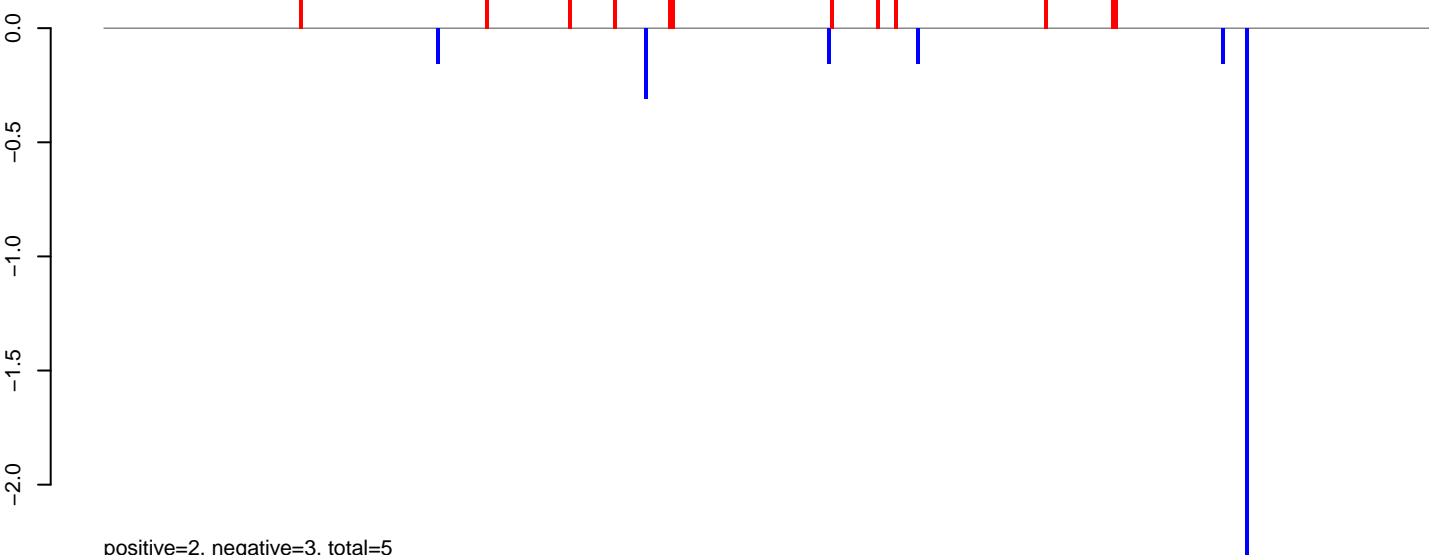
AeAlbo_C636_6dpi_SINV1.rep



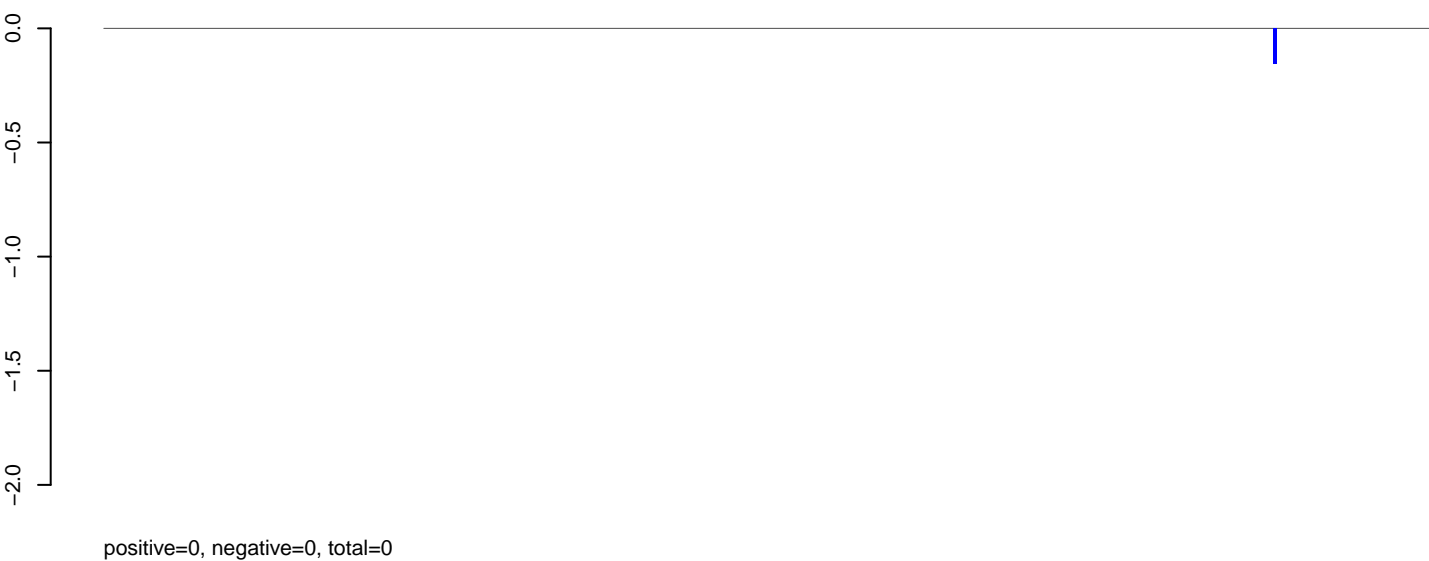
positive=4, negative=2, total=6 Window size=25, length=10893, virus@NC_027817.1-Parramatta_River_virus_isolate_92-B115745:1-10893

0 2000 4000 6000 8000 10000

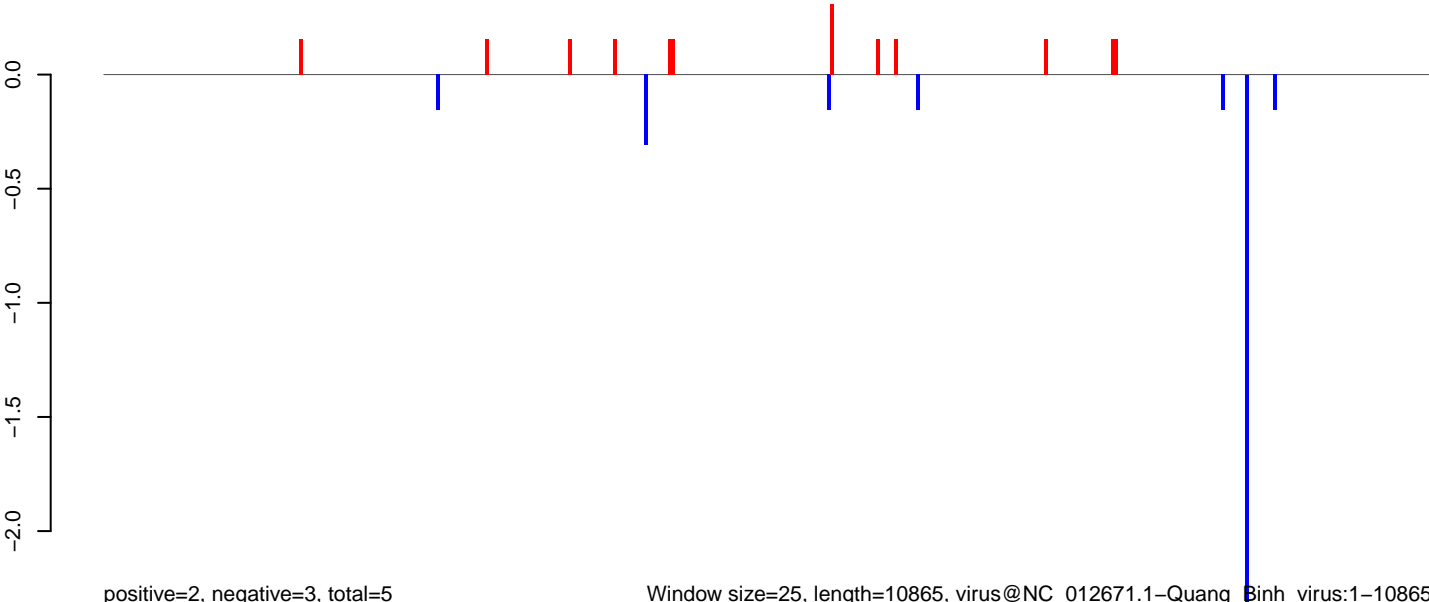
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



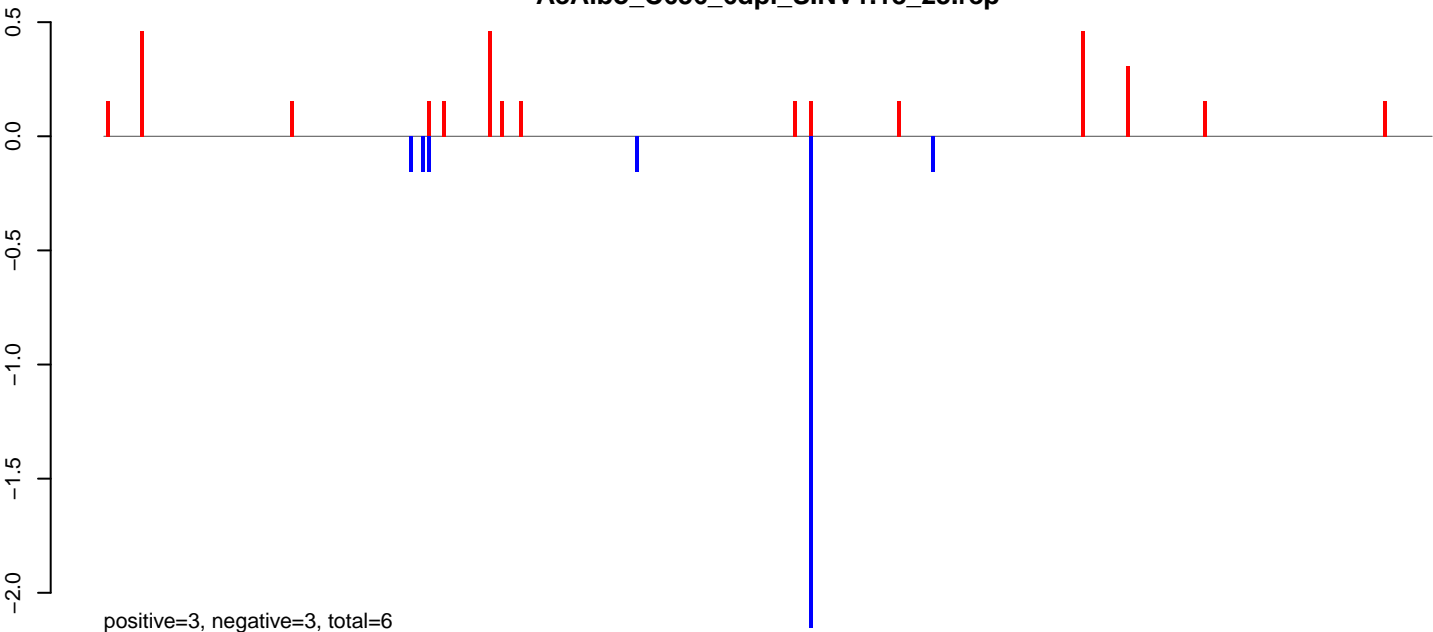
AeAlbo_C636_6dpi_SINV1.rep



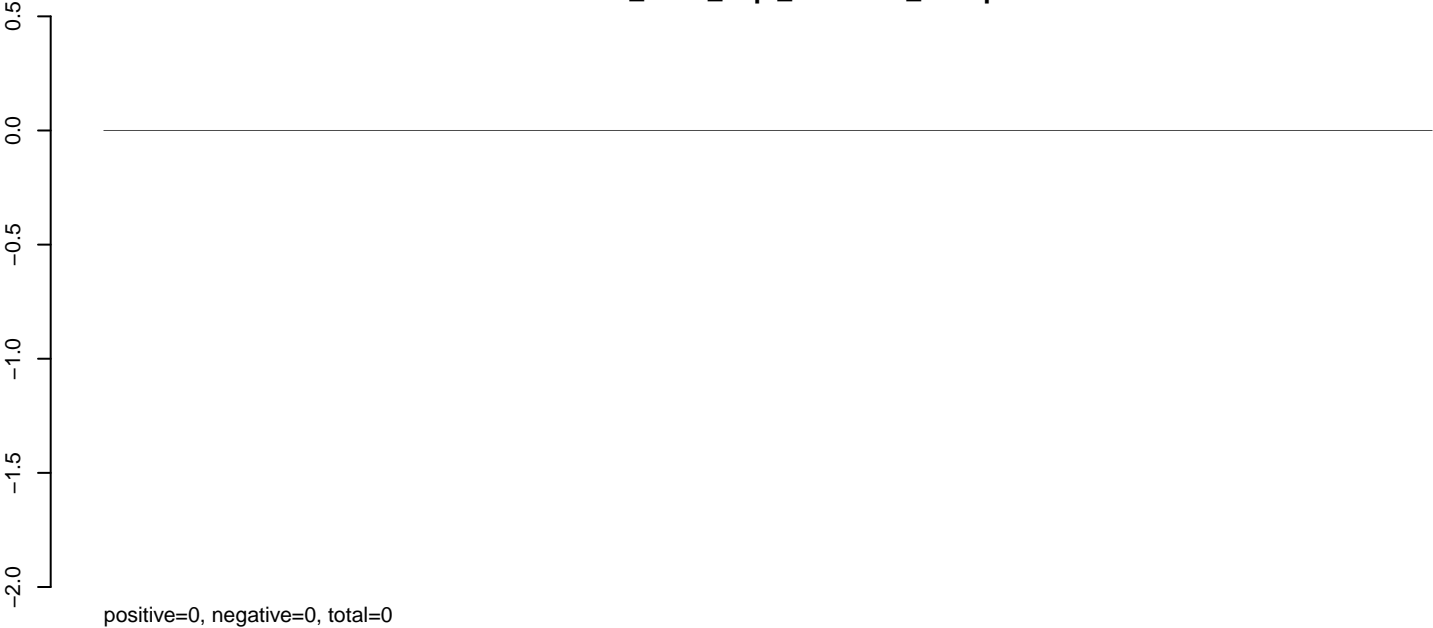
Window size=25, length=10865, virus@NC_012671.1-Quang_Binh_virus:1-10865

0 2000 4000 6000 8000 10000

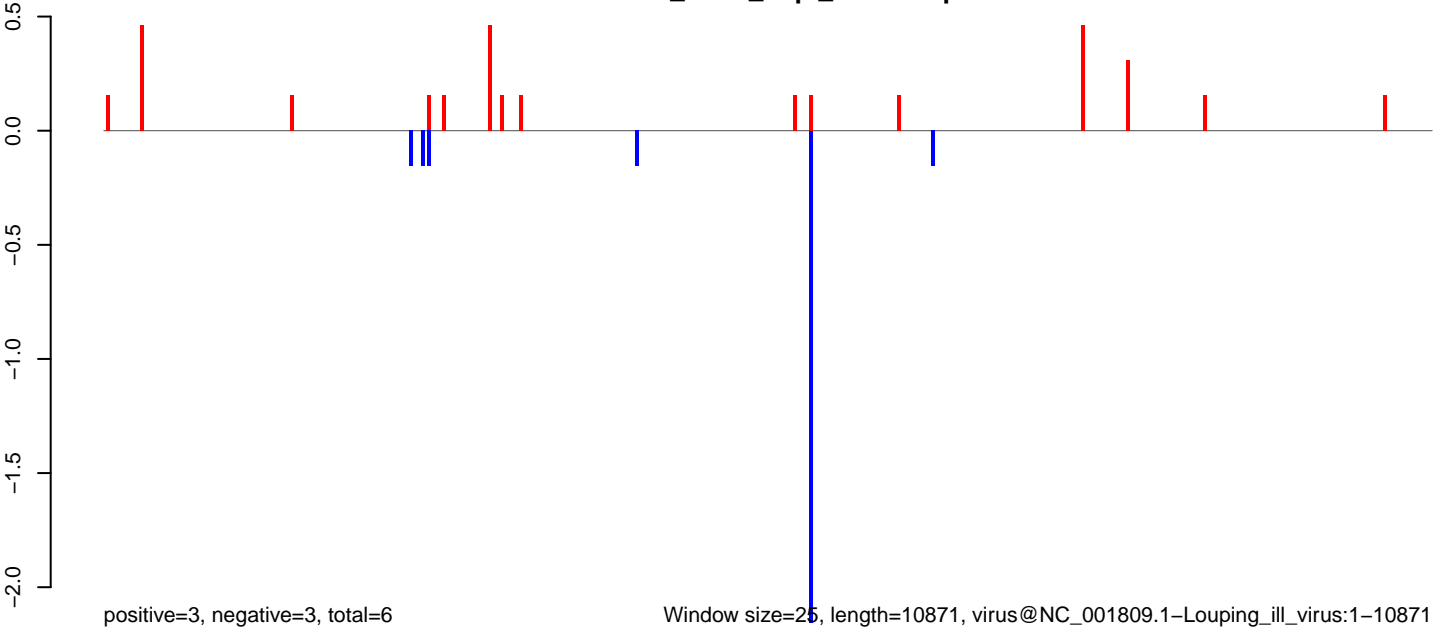
AeAlbo_C636_6dpi_SINV1.18_23.rep



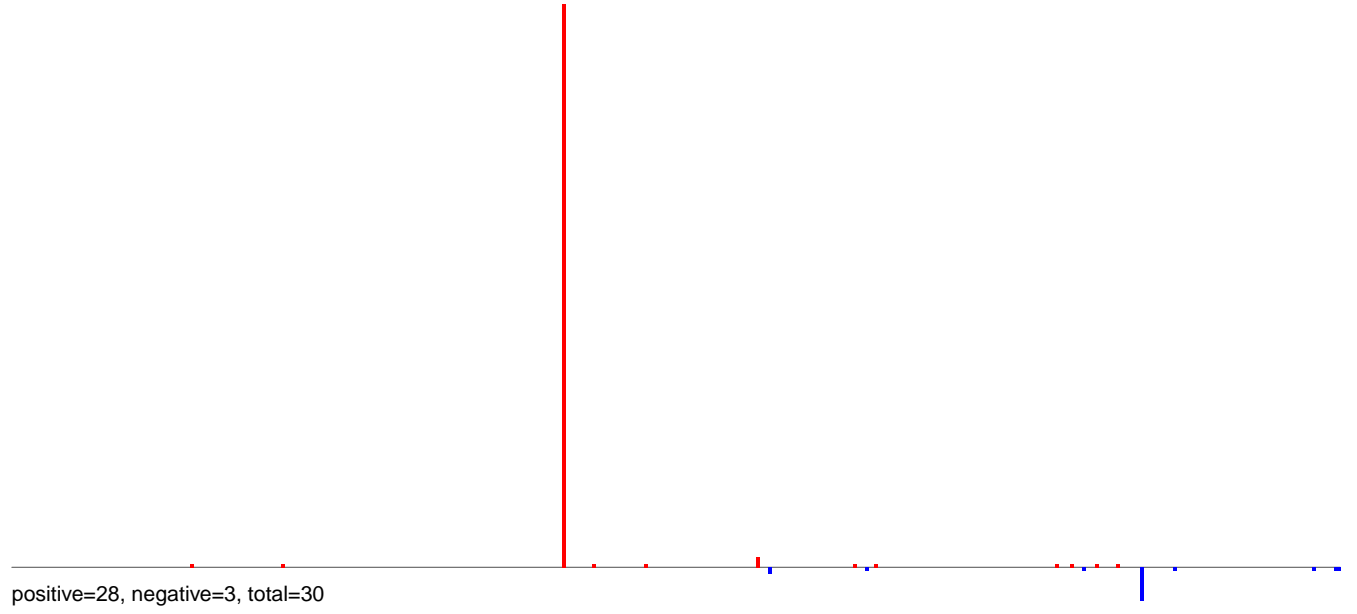
AeAlbo_C636_6dpi_SINV1.24_35.rep



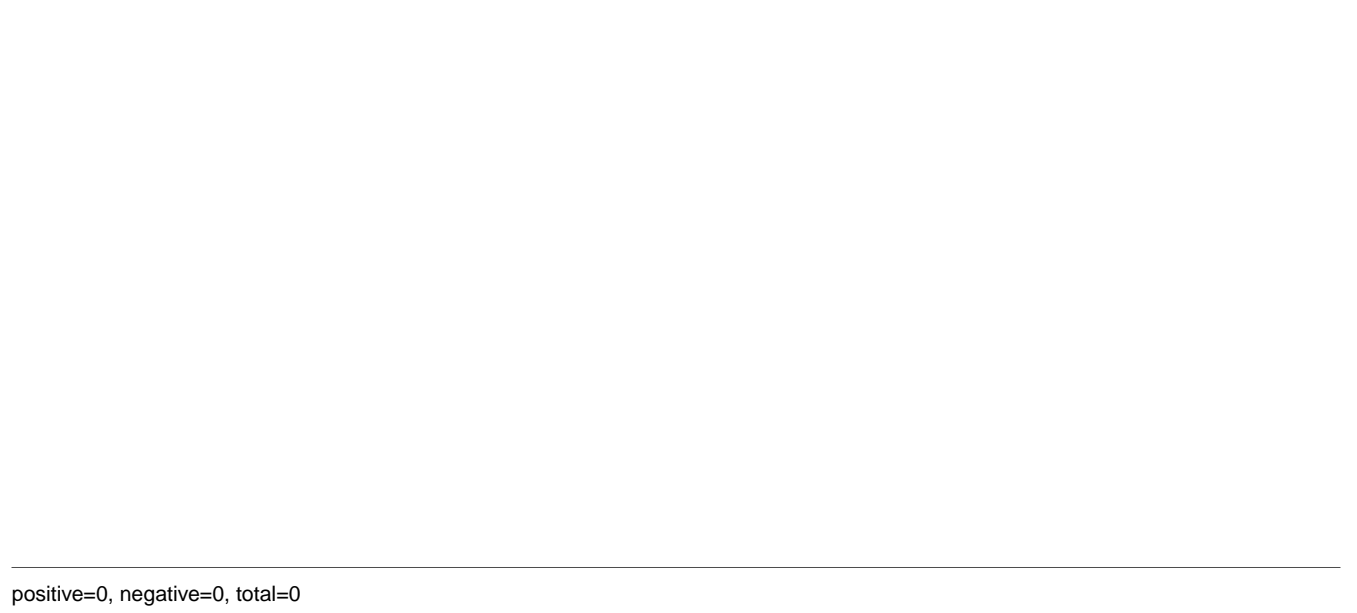
AeAlbo_C636_6dpi_SINV1.rep



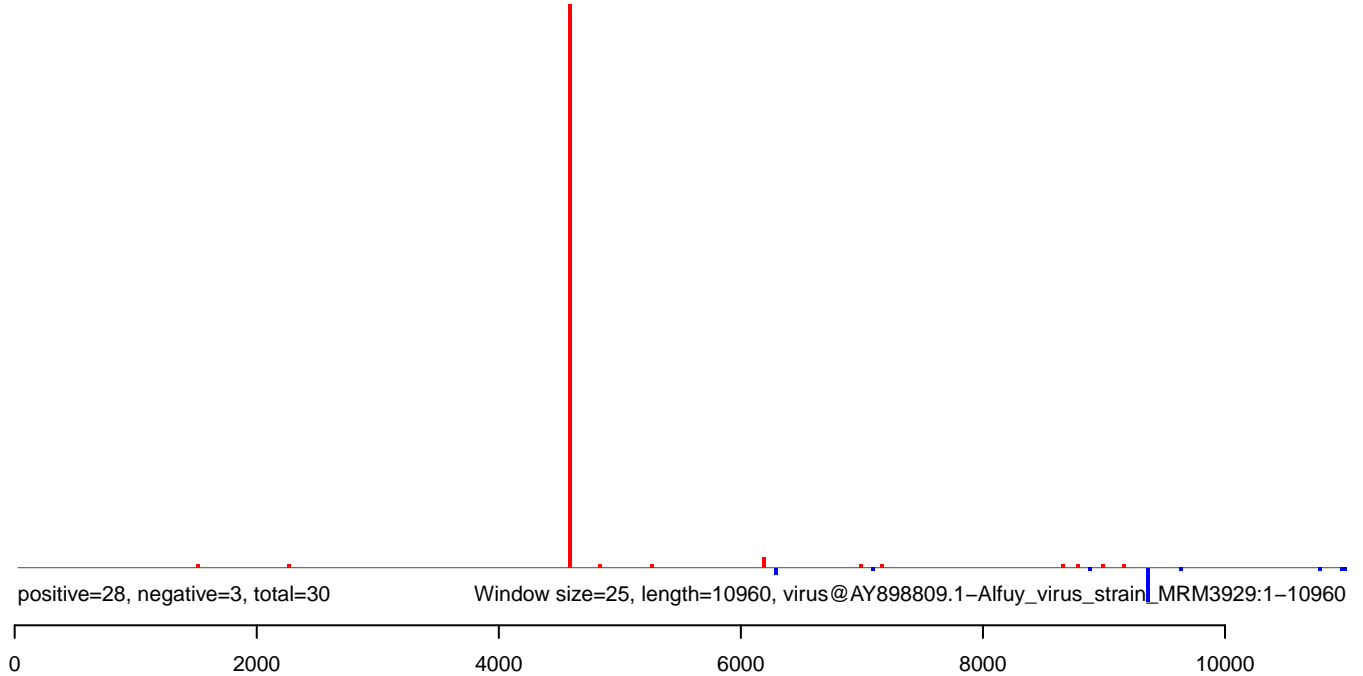
AeAlbo_C636_6dpi_SINV1.18_23.rep



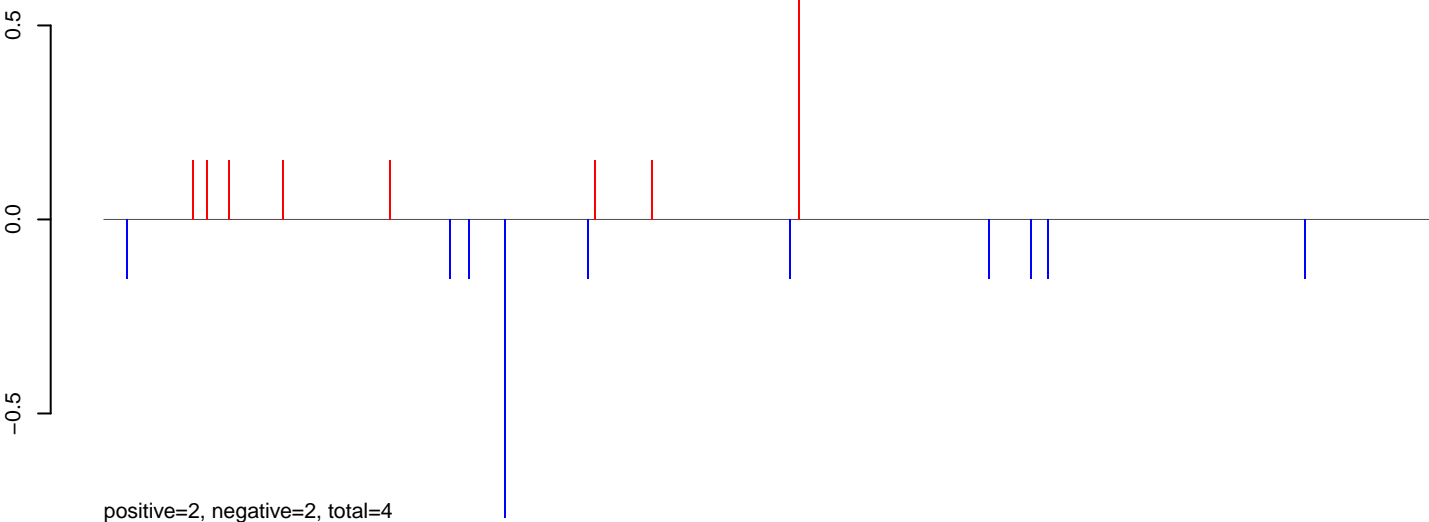
AeAlbo_C636_6dpi_SINV1.24_35.rep



AeAlbo_C636_6dpi_SINV1.rep



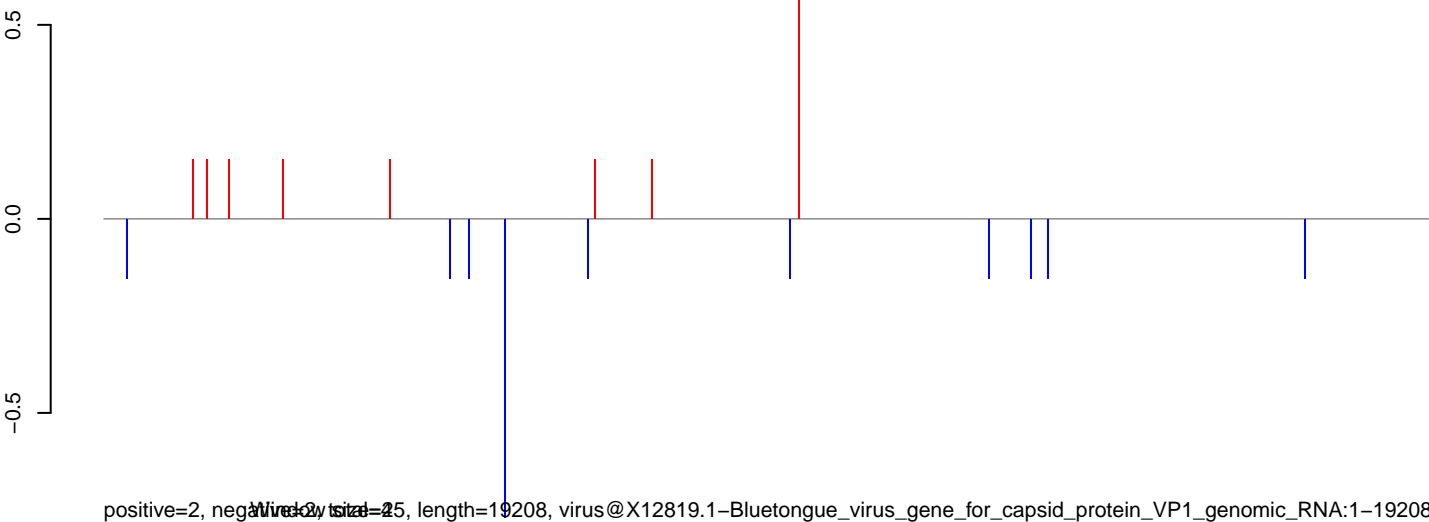
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



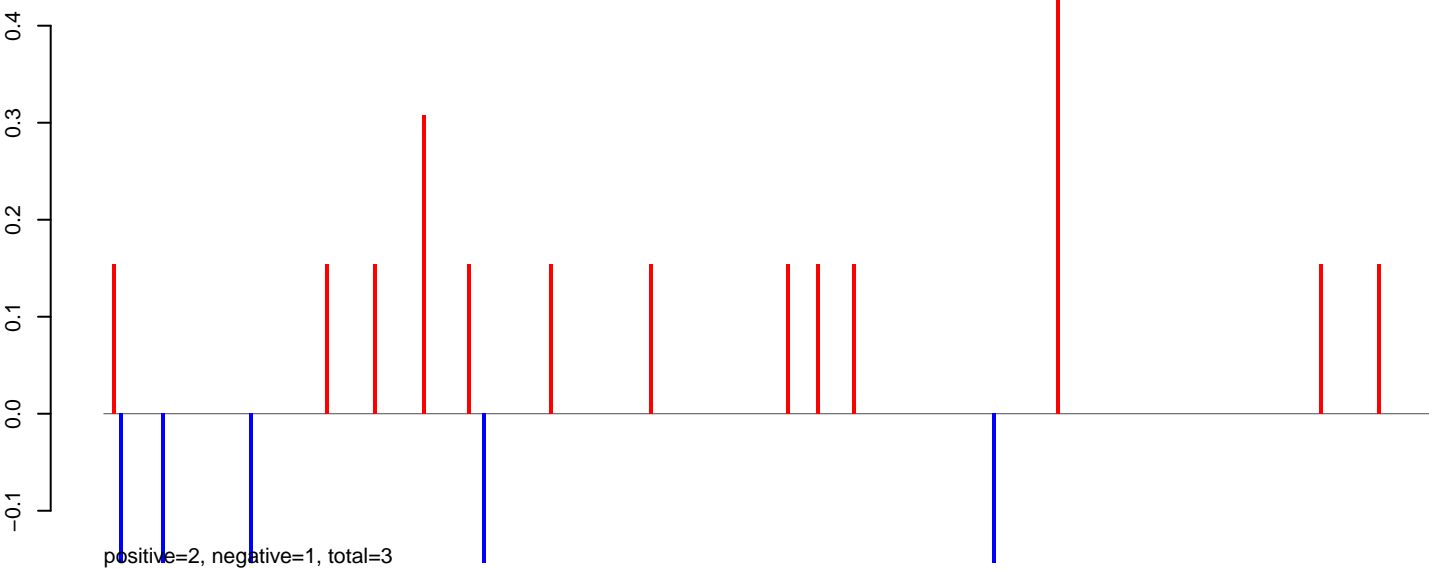
AeAlbo_C636_6dpi_SINV1.rep



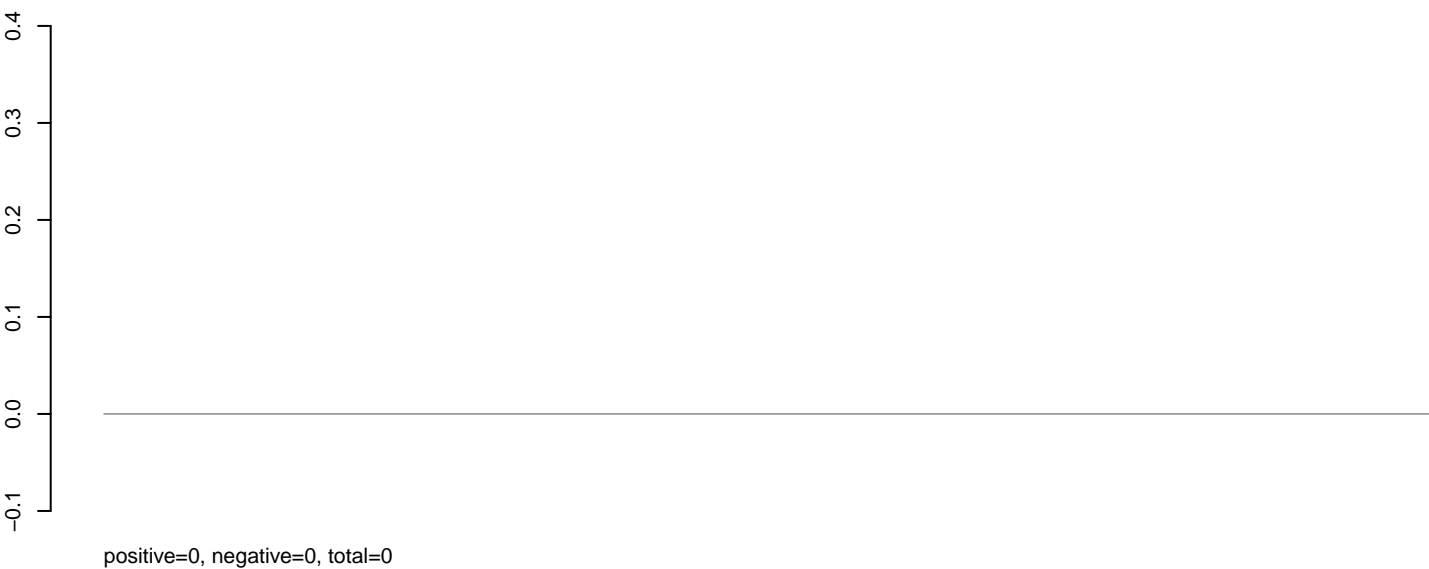
positive=2, negative=2, total=4, virus@X12819.1-Bluetongue_virus_gene_for_capsid_protein_VP1_genomic_RNA:1-19208

0 5000 10000 15000 20000

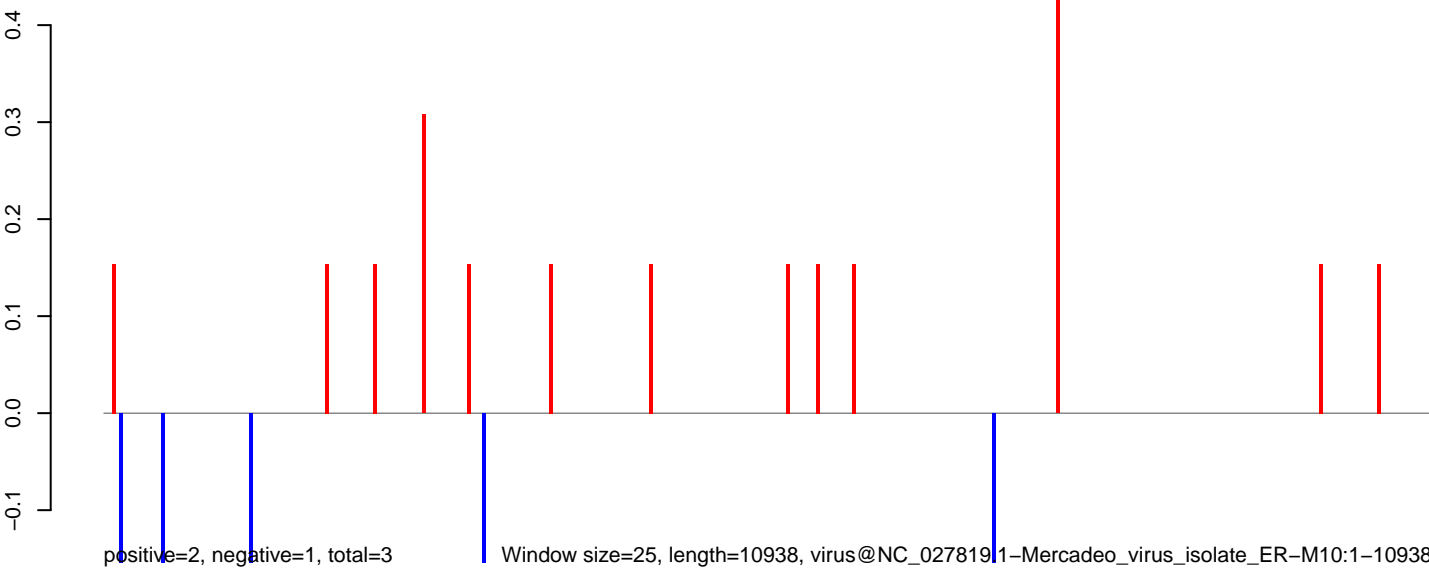
AeAlbo_C636_6dpi_SINV1.18_23.rep



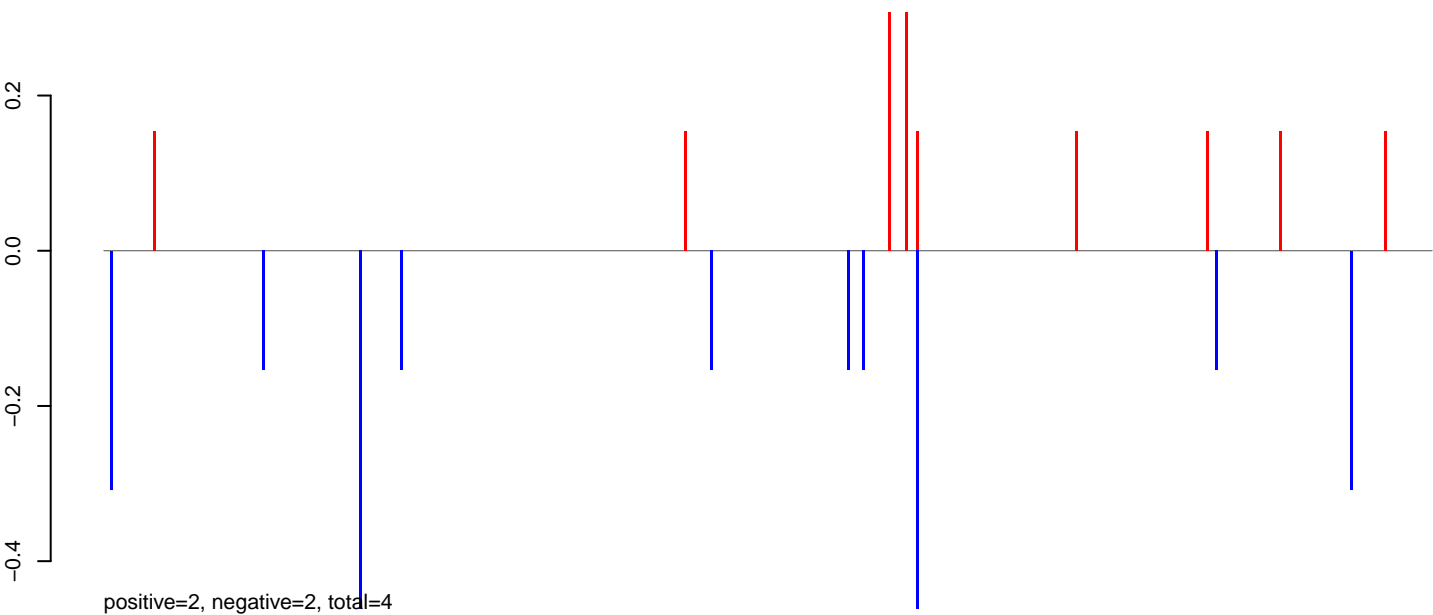
AeAlbo_C636_6dpi_SINV1.24_35.rep



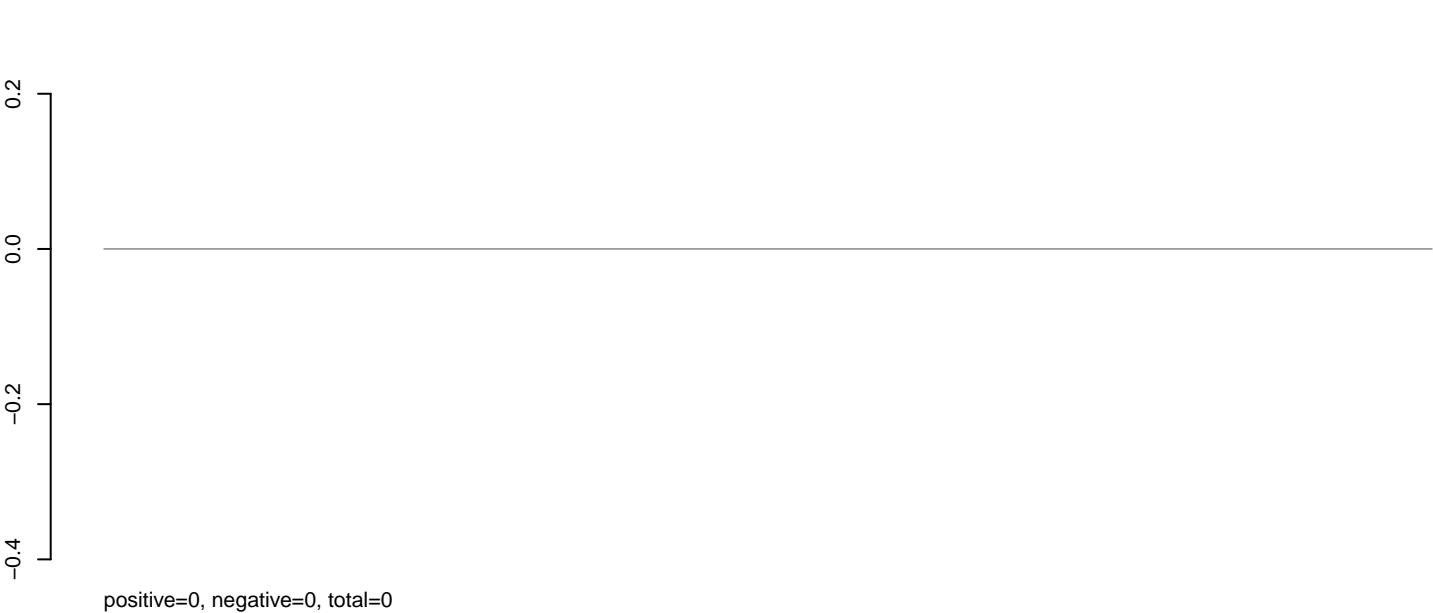
AeAlbo_C636_6dpi_SINV1.rep



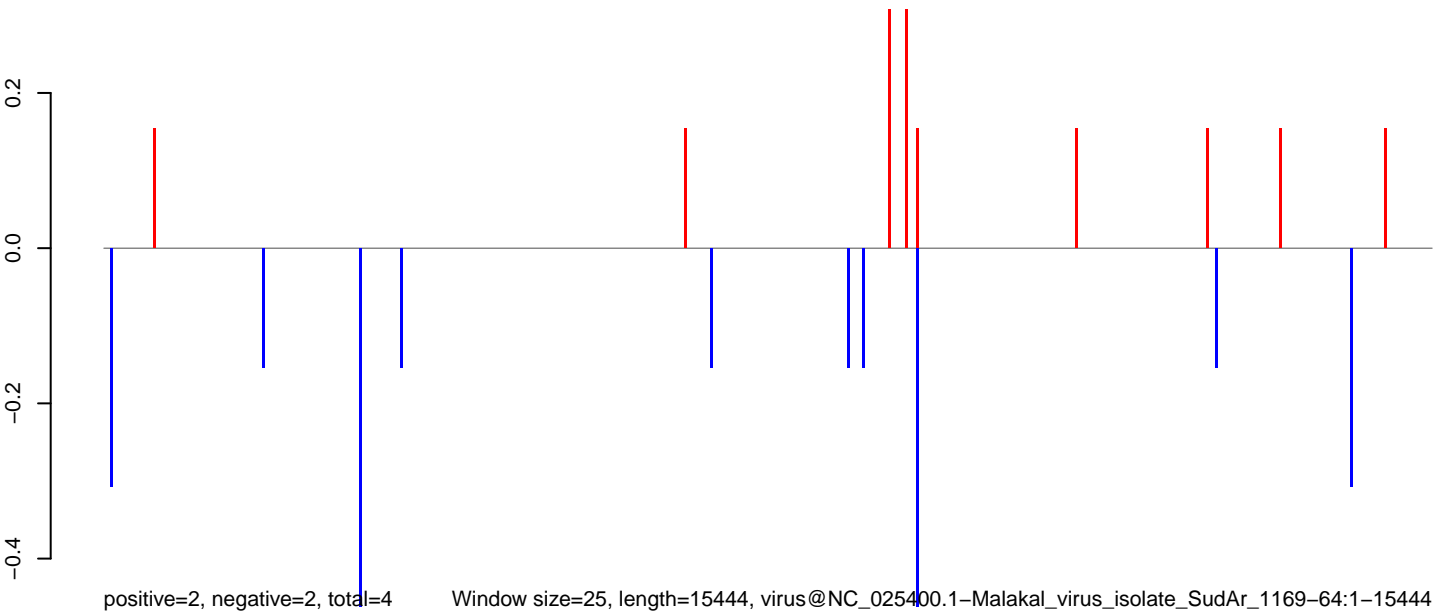
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep

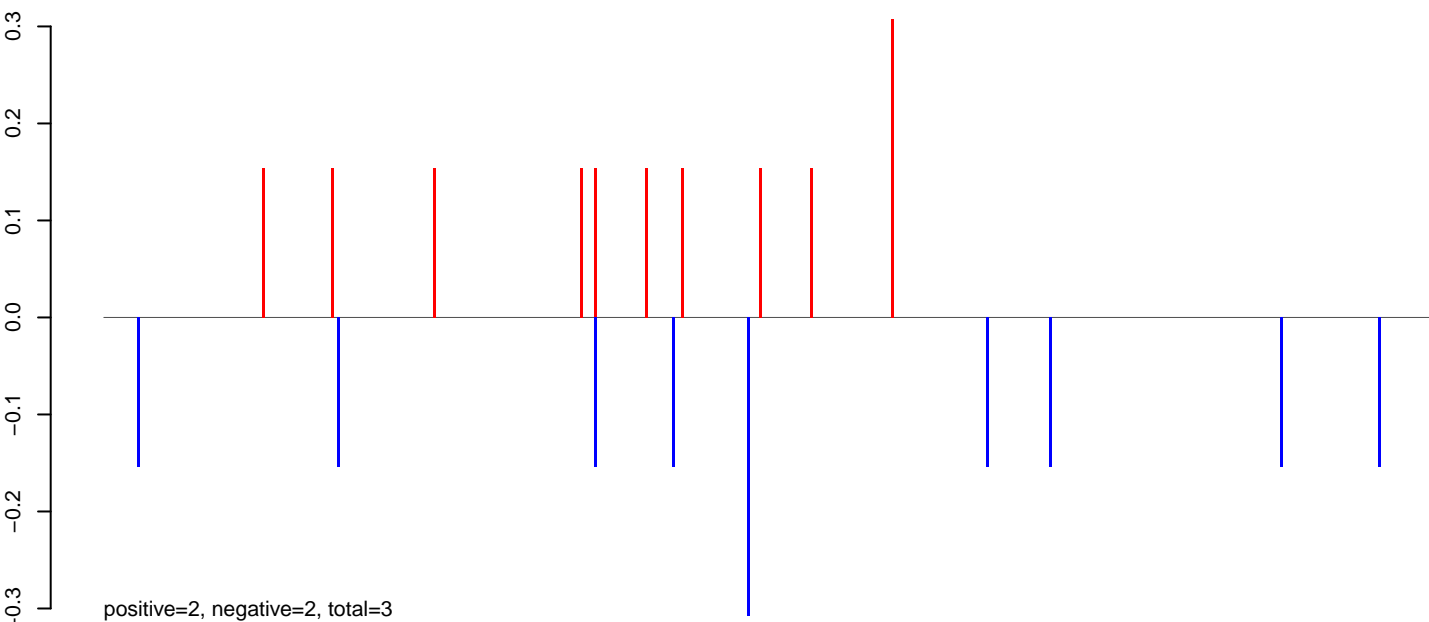


AeAlbo_C636_6dpi_SINV1.rep

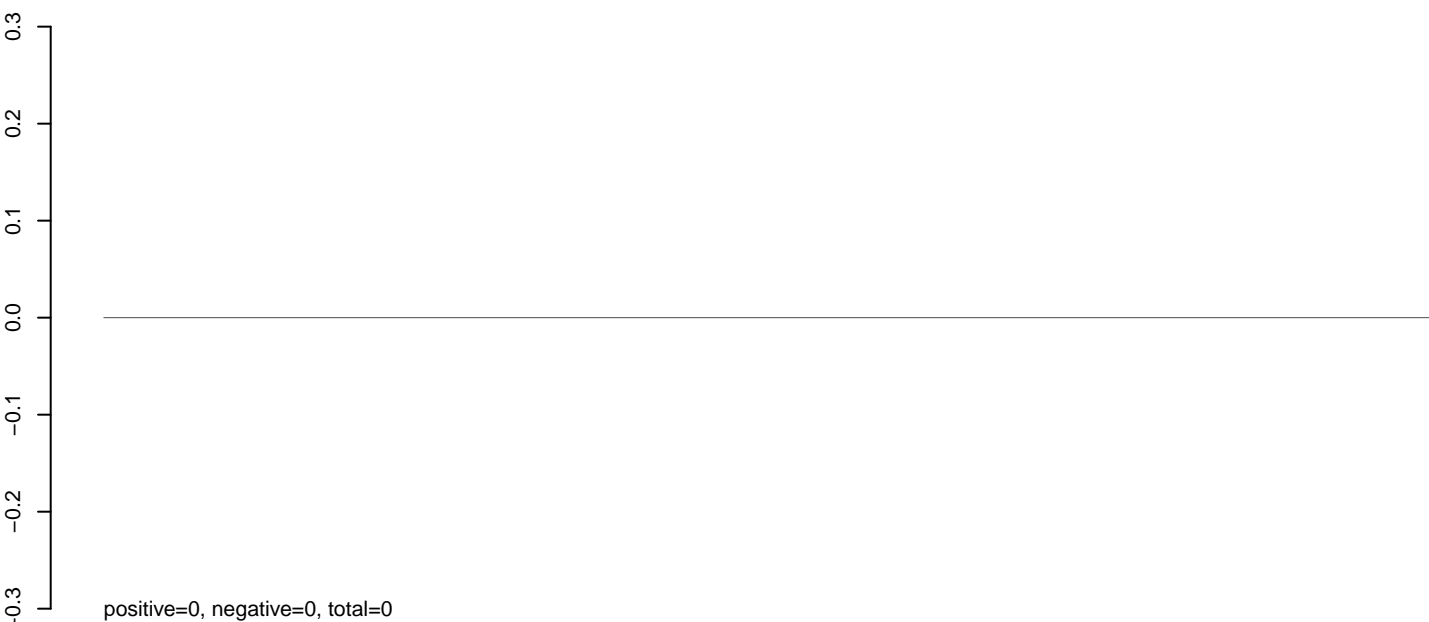


0 5000 10000 15000

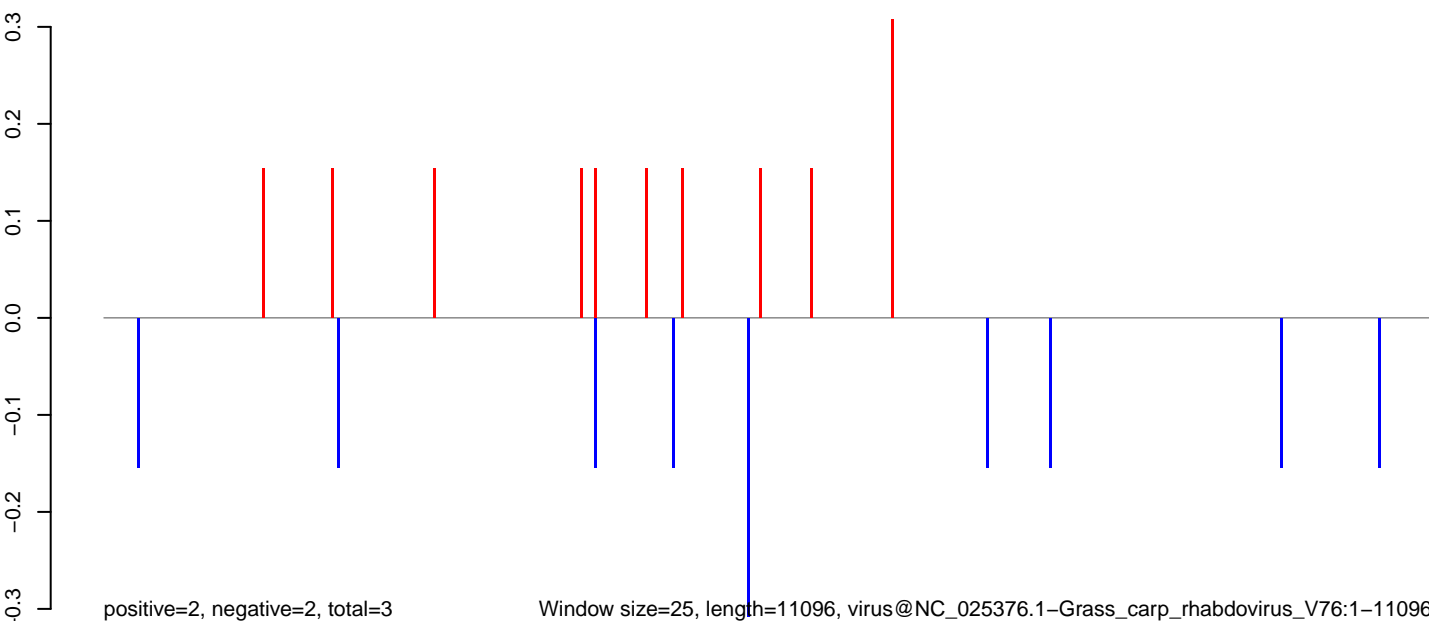
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



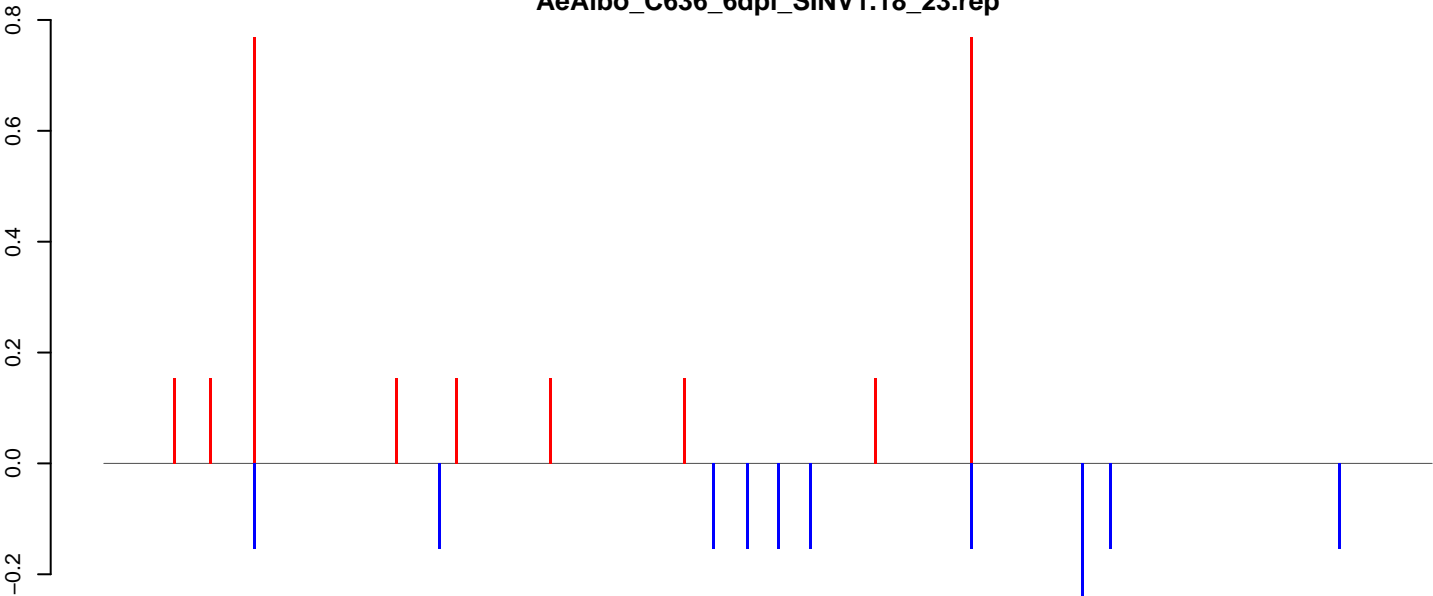
AeAlbo_C636_6dpi_SINV1.rep



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

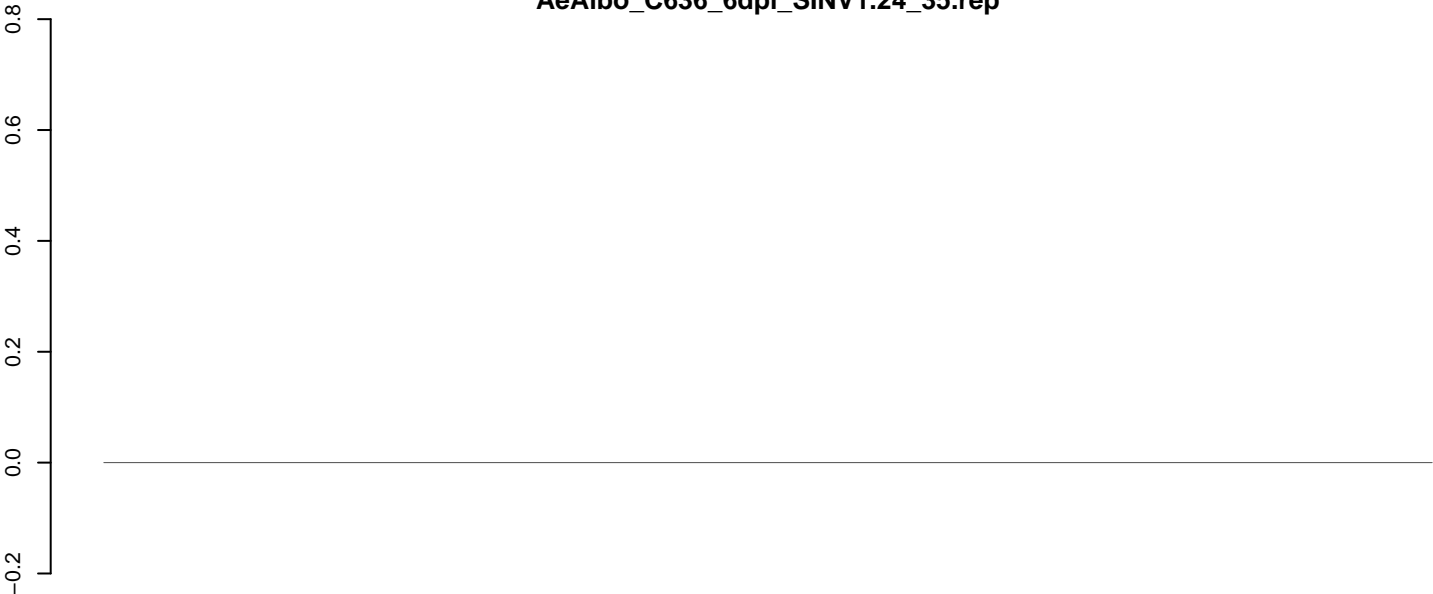
0 2000 4000 6000 8000 10000

AeAlbo_C636_6dpi_SINV1.18_23.rep



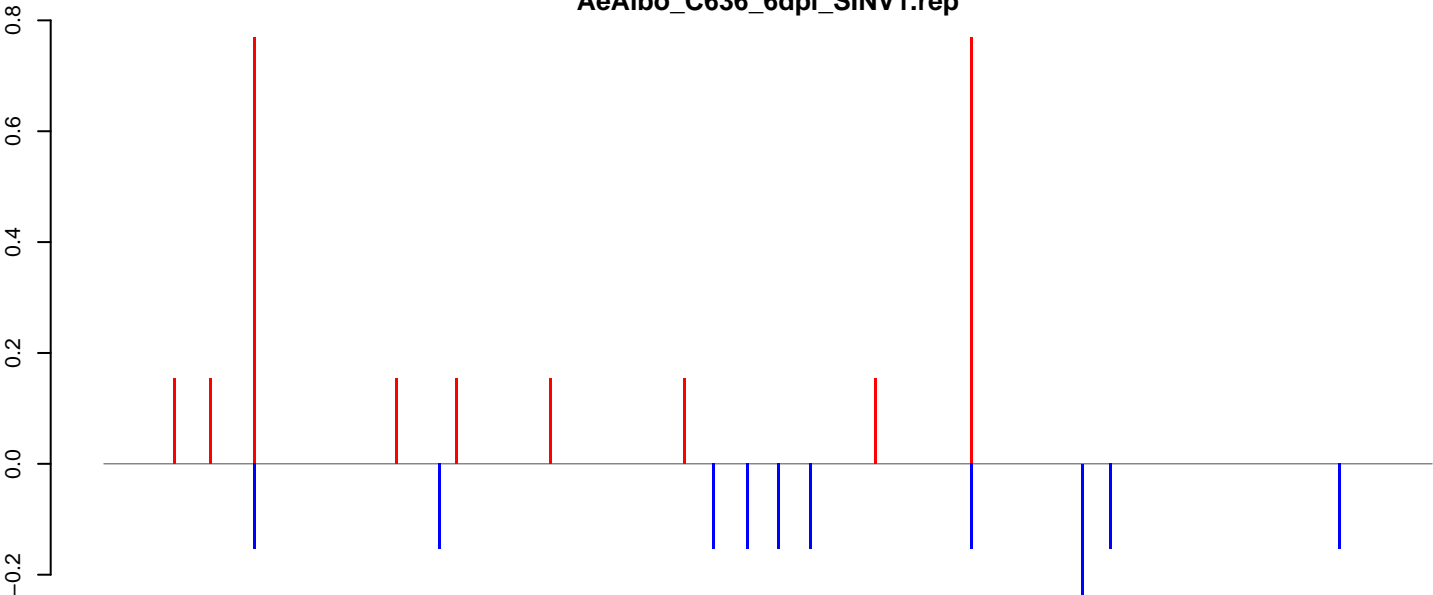
positive=3, negative=2, total=4

AeAlbo_C636_6dpi_SINV1.24_35.rep



positive=0, negative=0, total=0

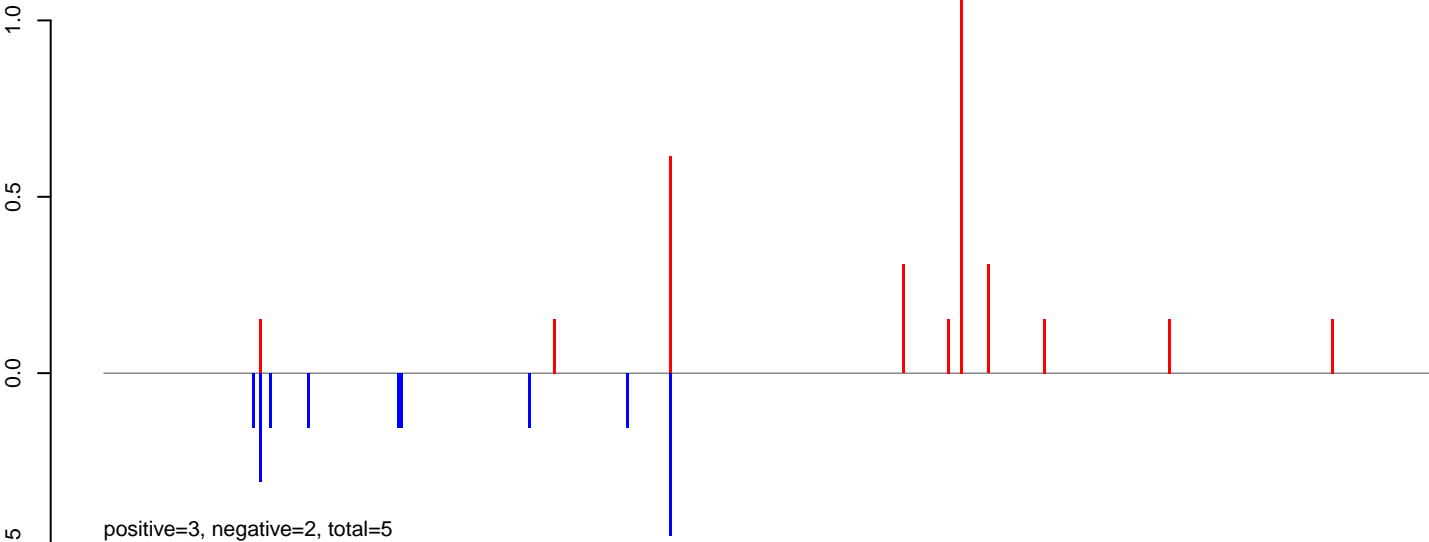
AeAlbo_C636_6dpi_SINV1.rep



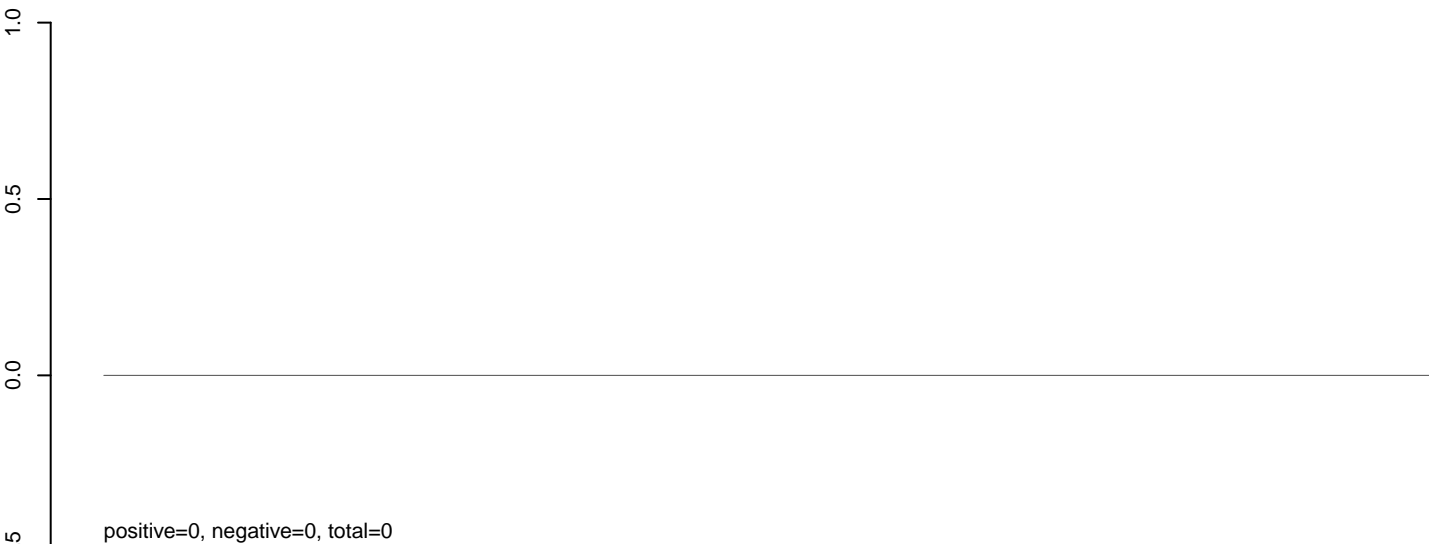
bow size=25, length=13784, interval=2, use=4 NC_034531.1-Manitoba_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycoprot:1-13784

0 2000 4000 6000 8000 10000 12000 14000

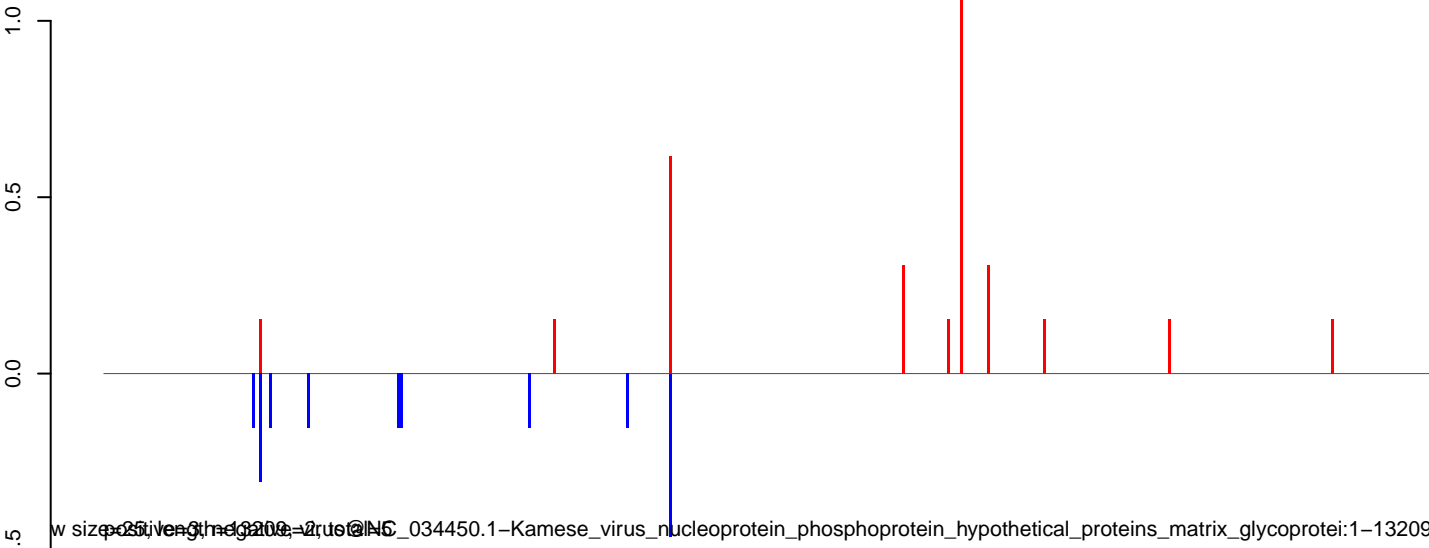
AeAlbo_C636_6dpi_SINV1.18_23.rep



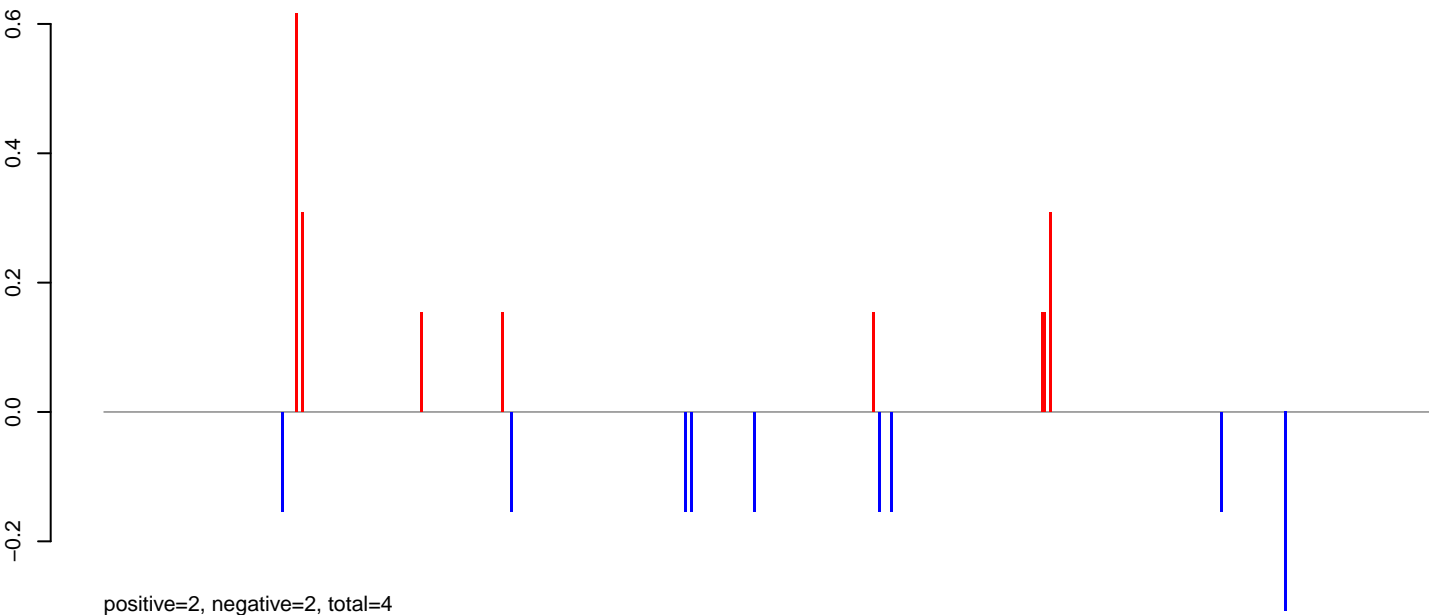
AeAlbo_C636_6dpi_SINV1.24_35.rep



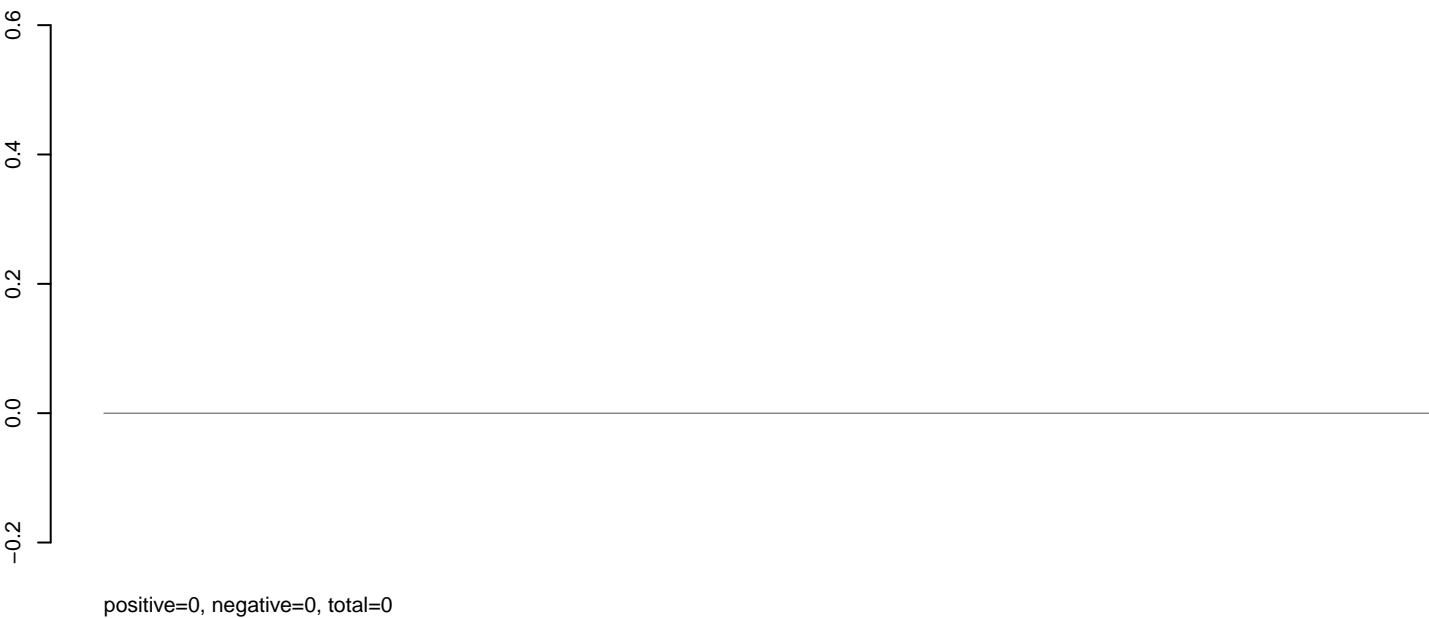
AeAlbo_C636_6dpi_SINV1.rep



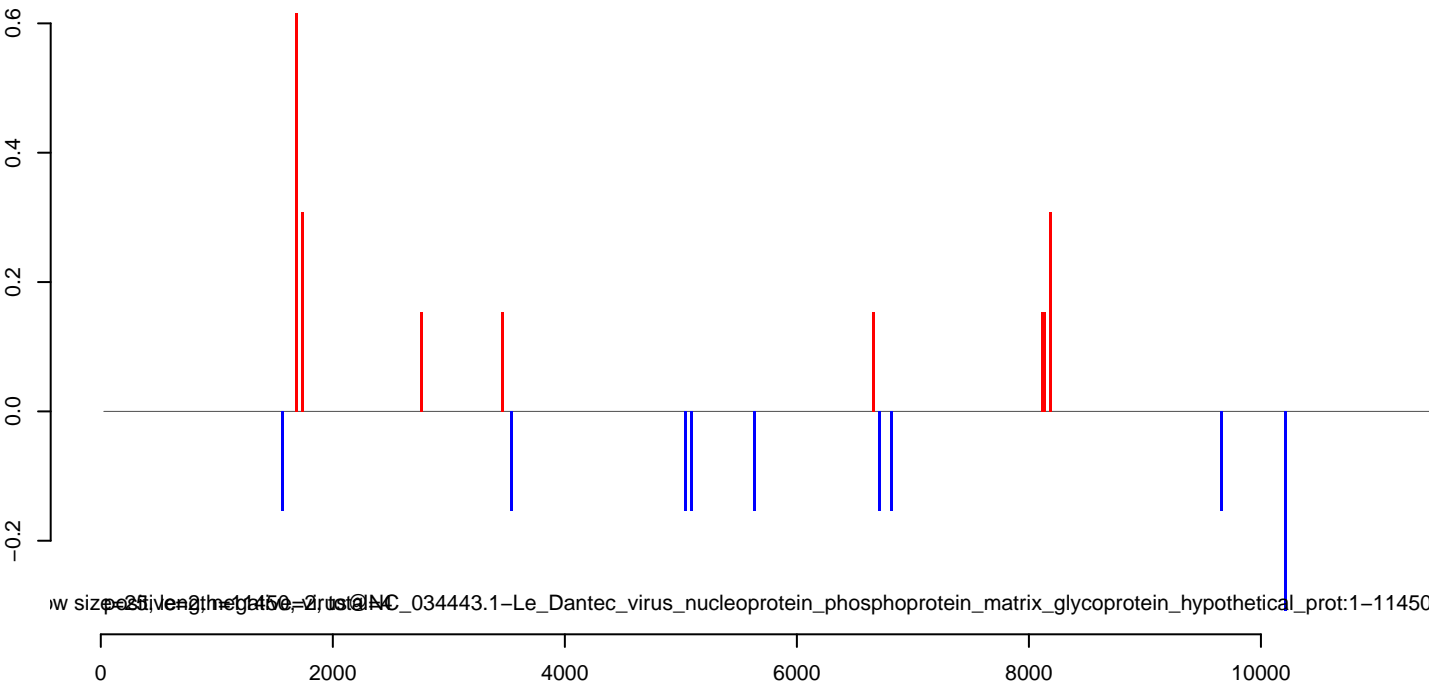
AeAlbo_C636_6dpi_SINV1.18_23.rep



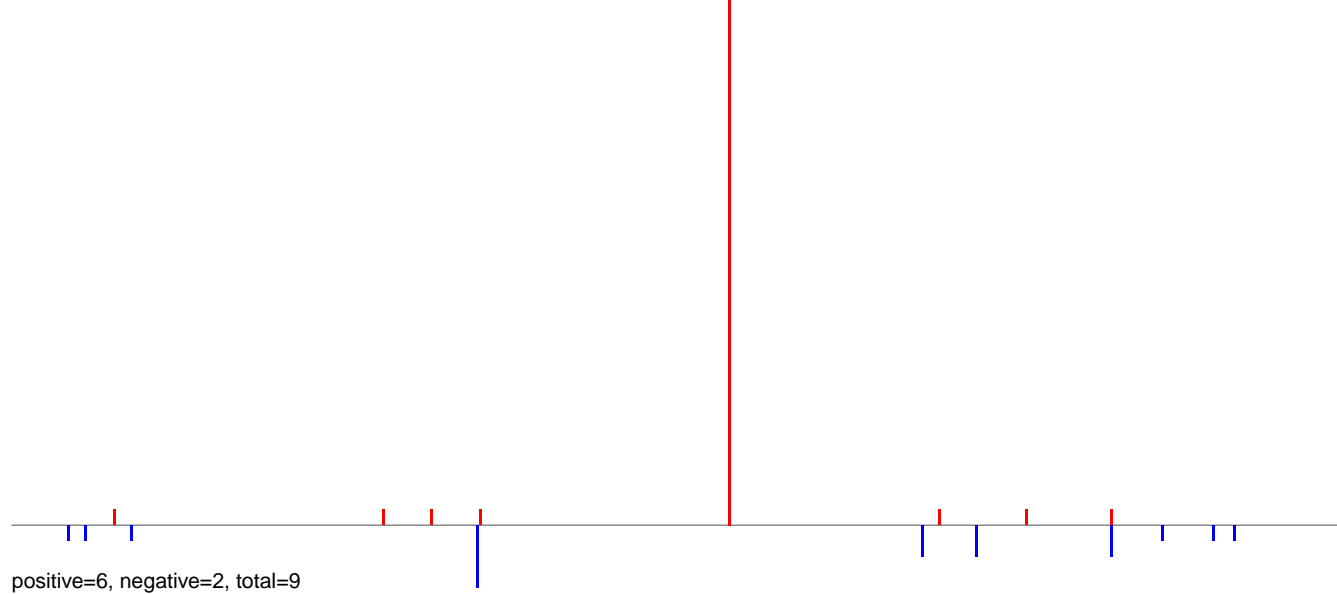
AeAlbo_C636_6dpi_SINV1.24_35.rep



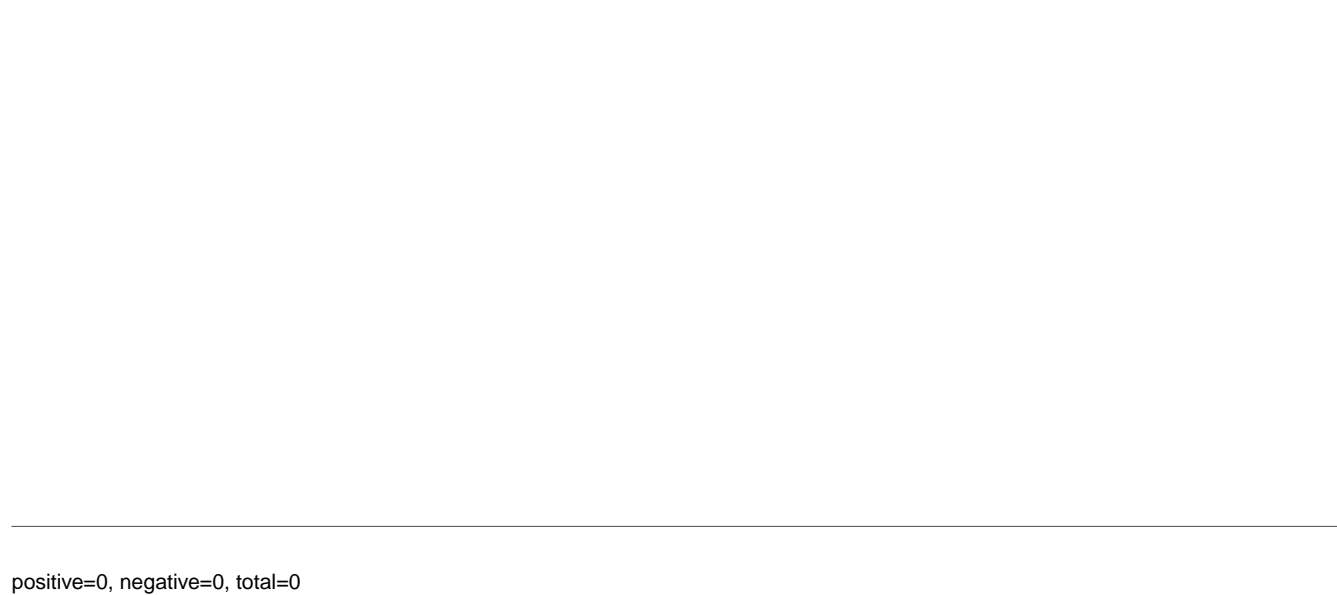
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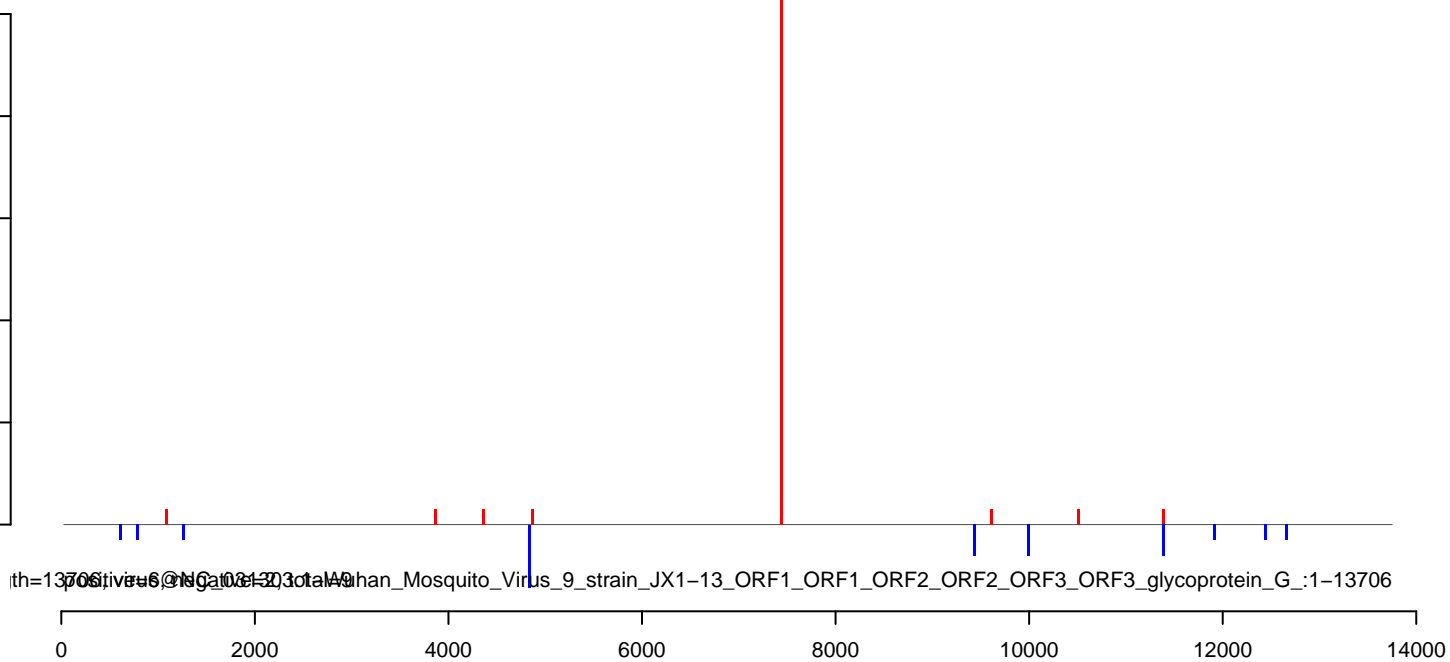
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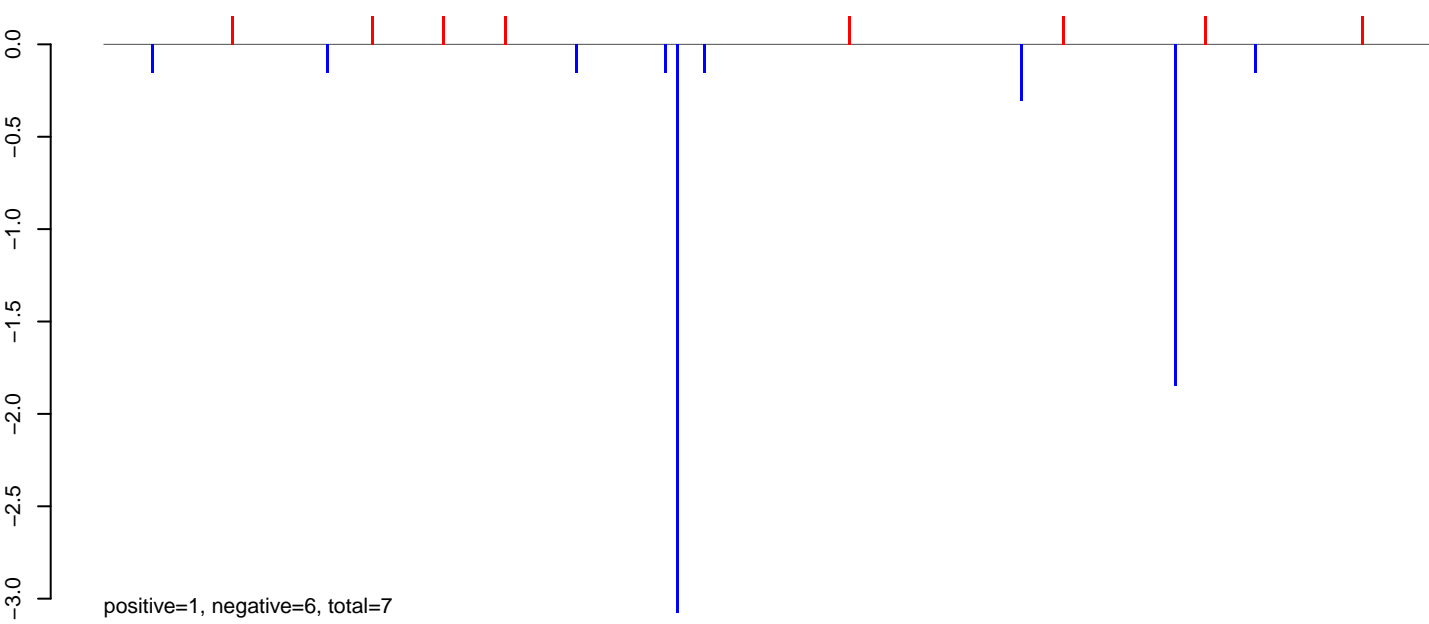
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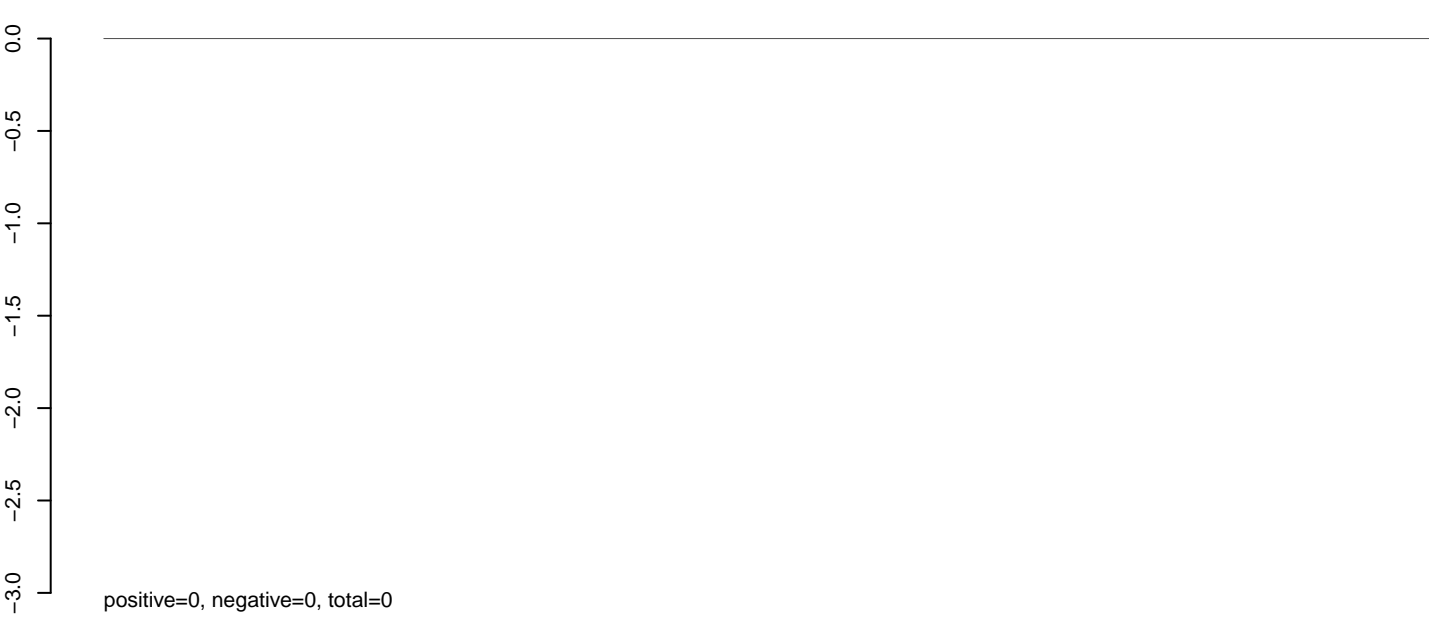
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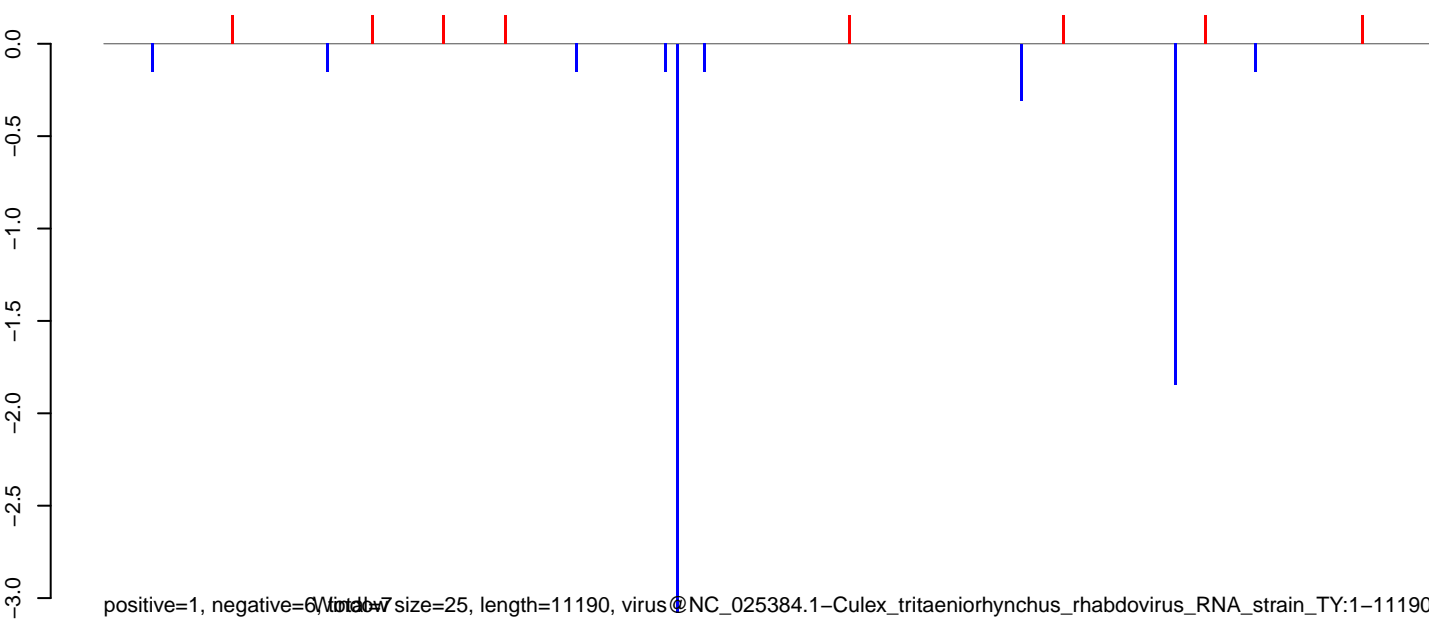
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AeAlbo_C636_6dpi_SINV1.24_35.rep



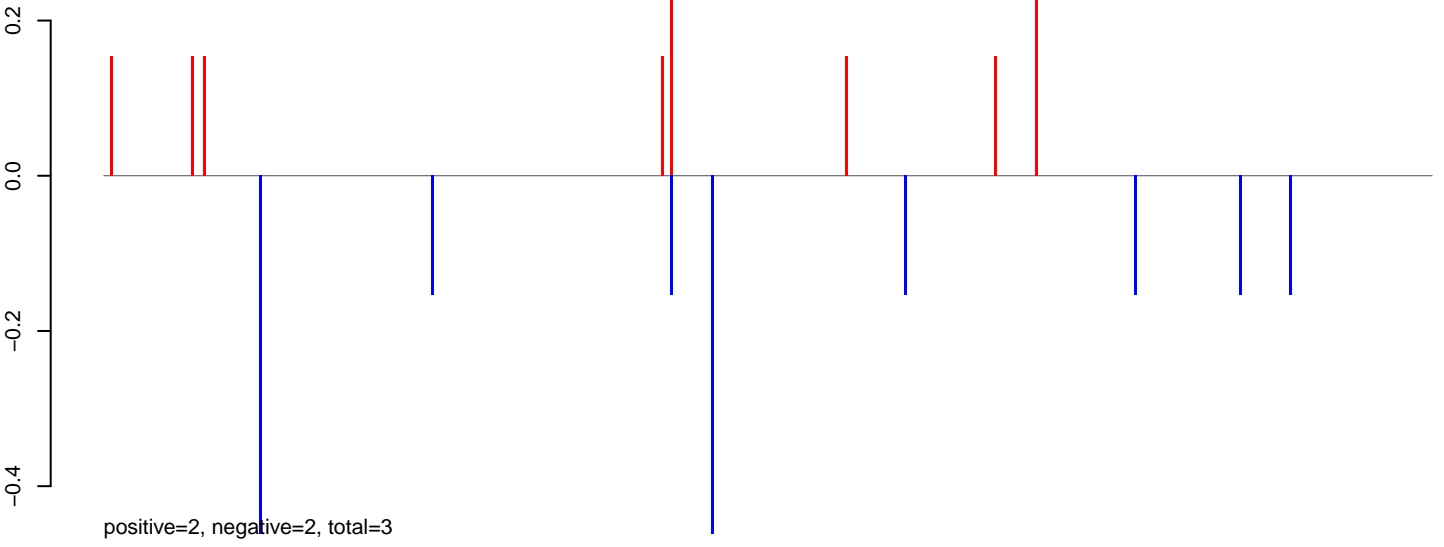
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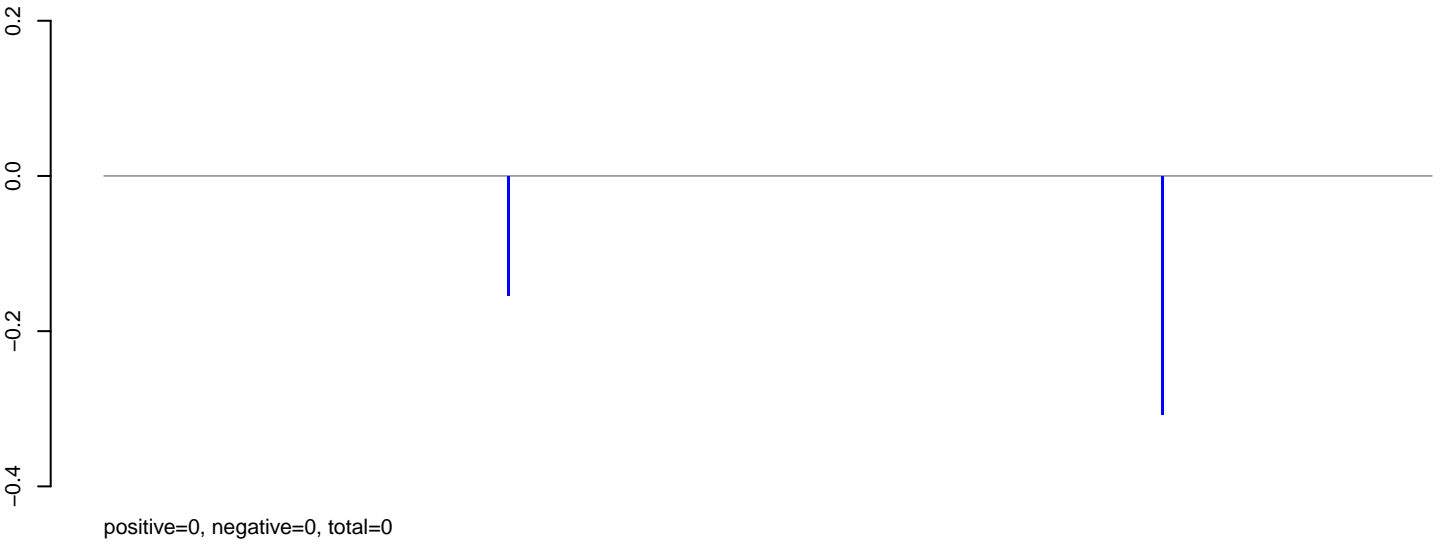
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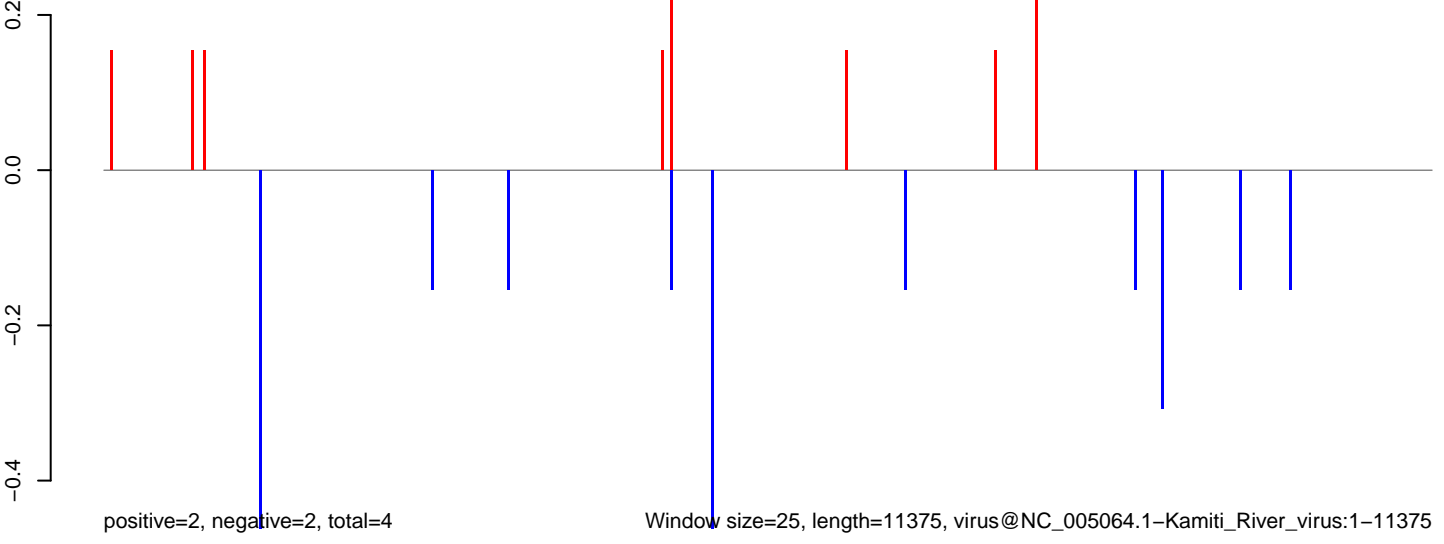
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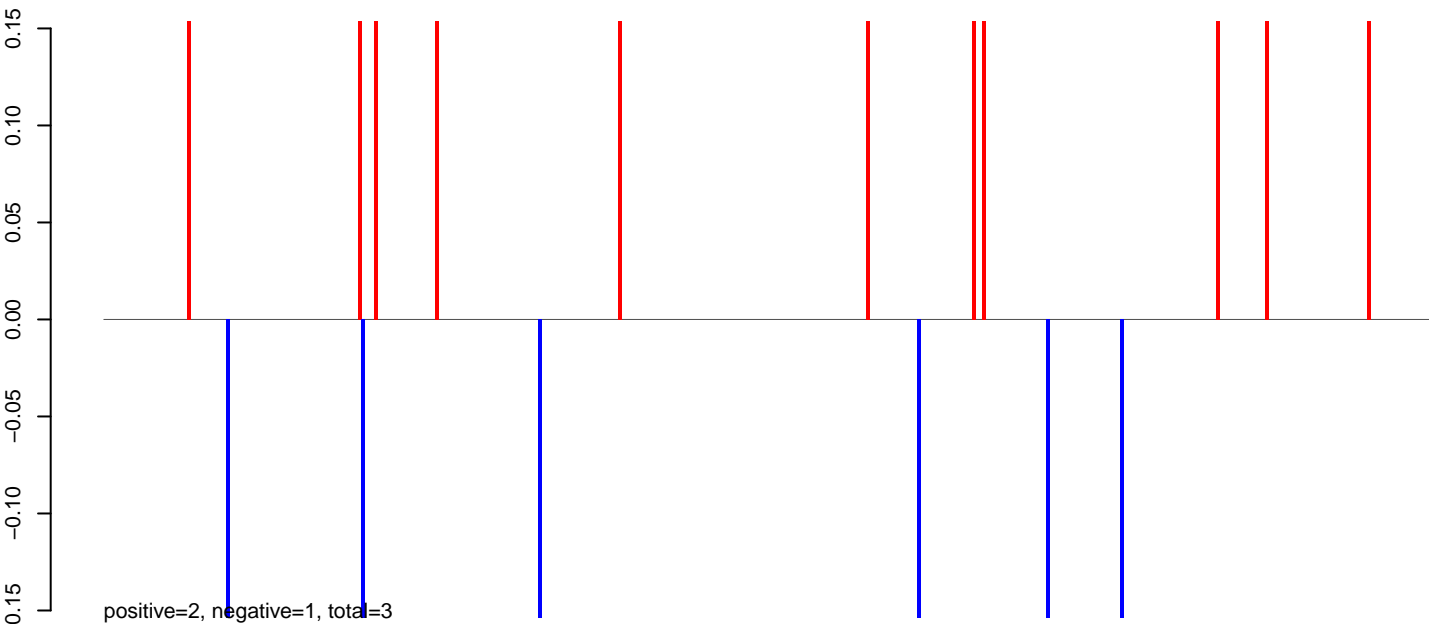


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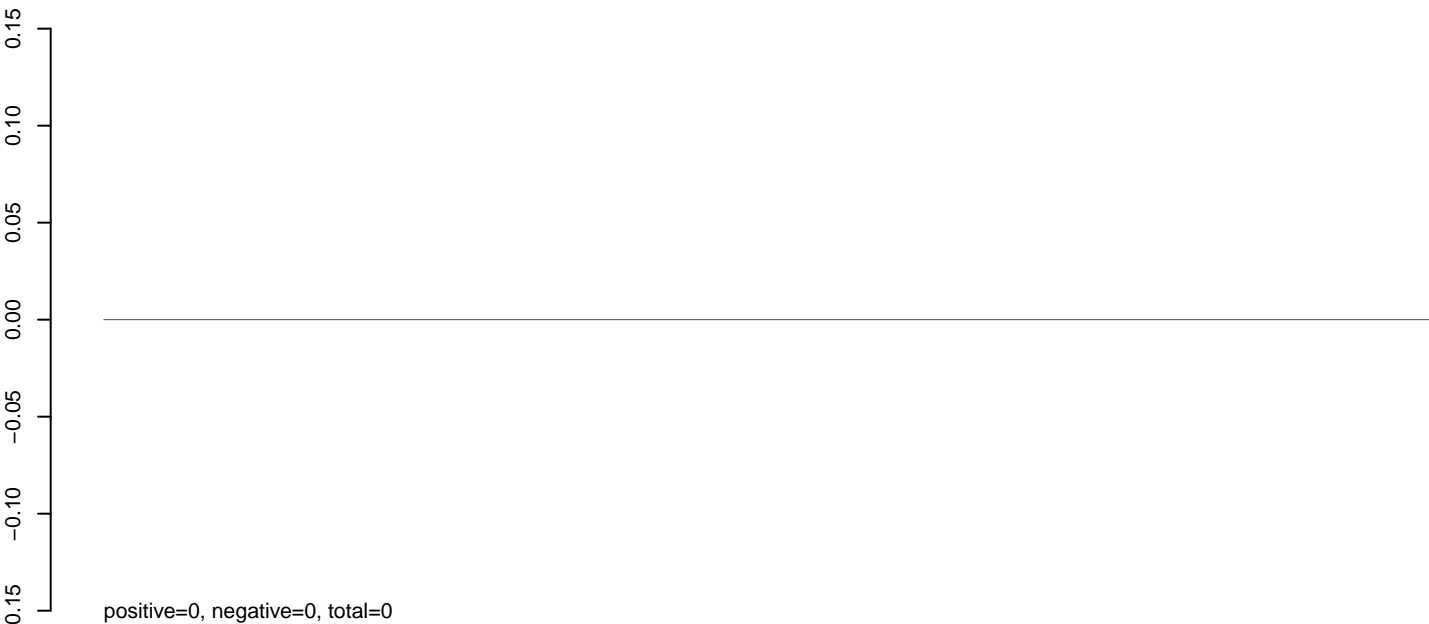


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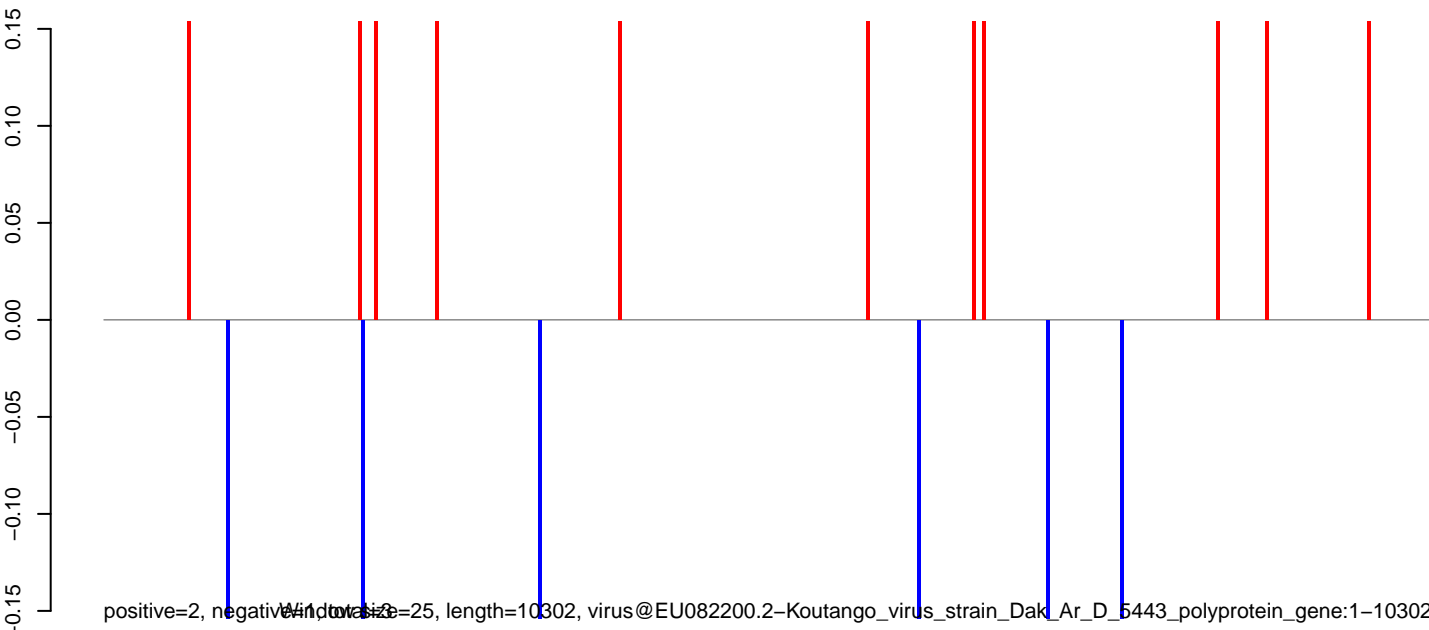
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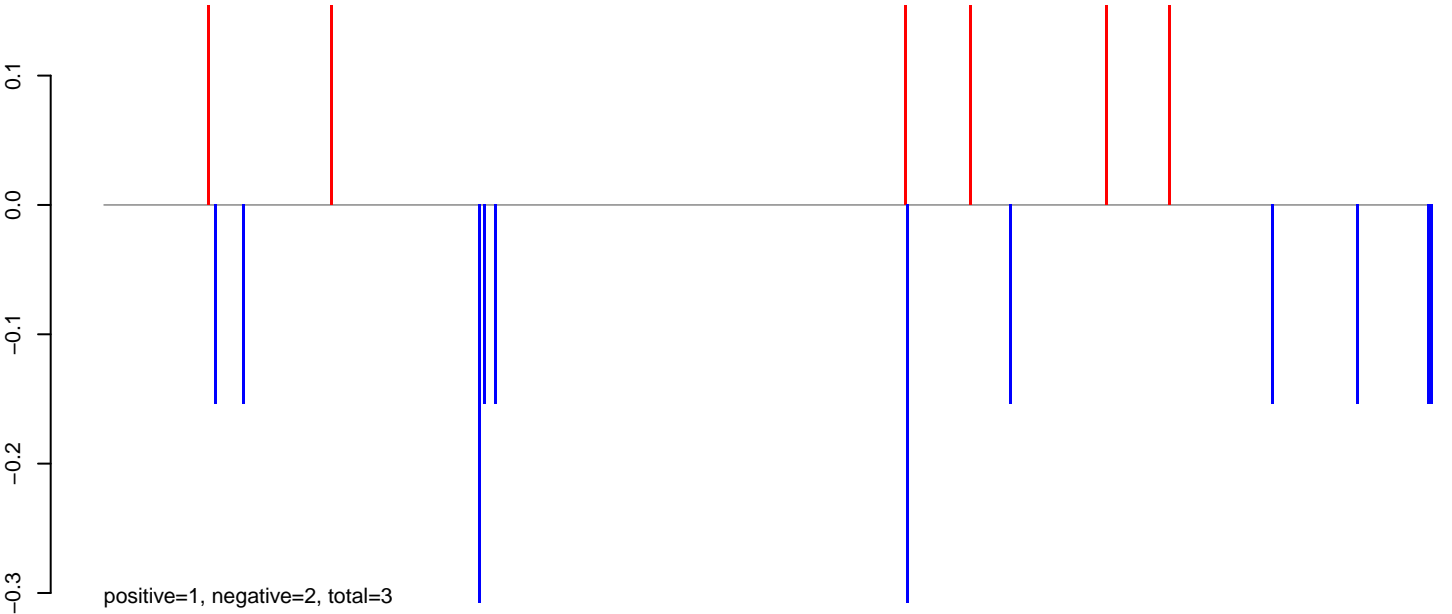


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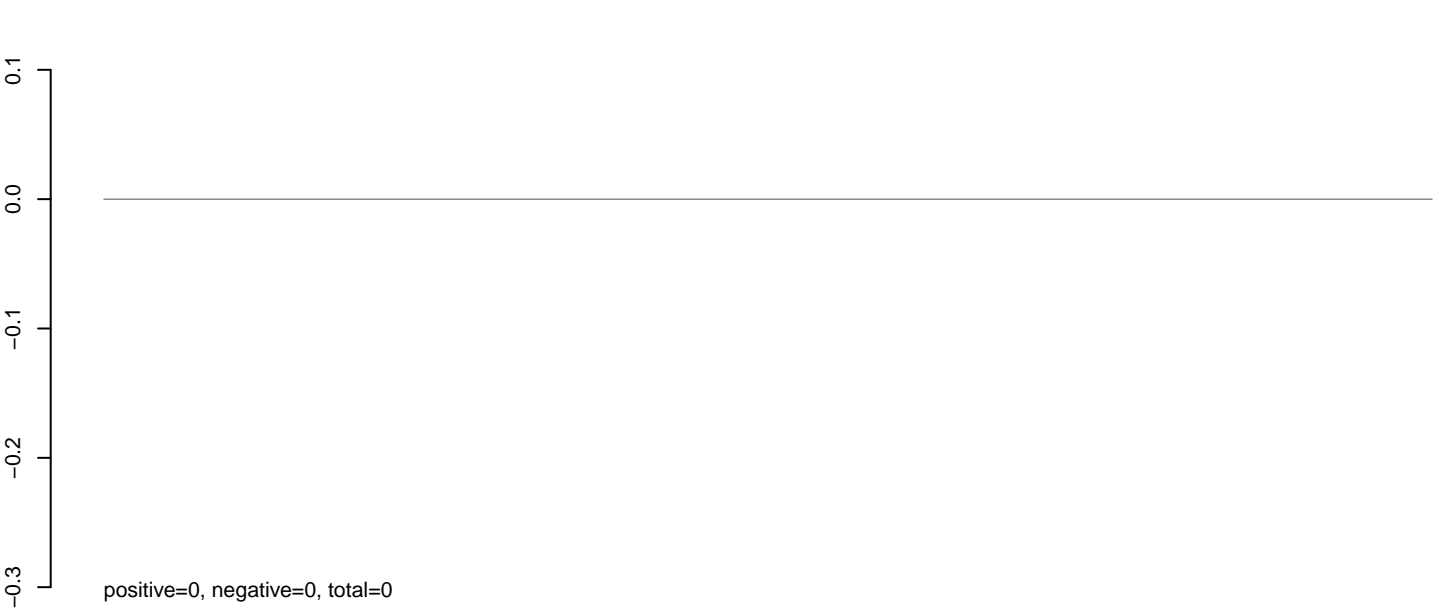


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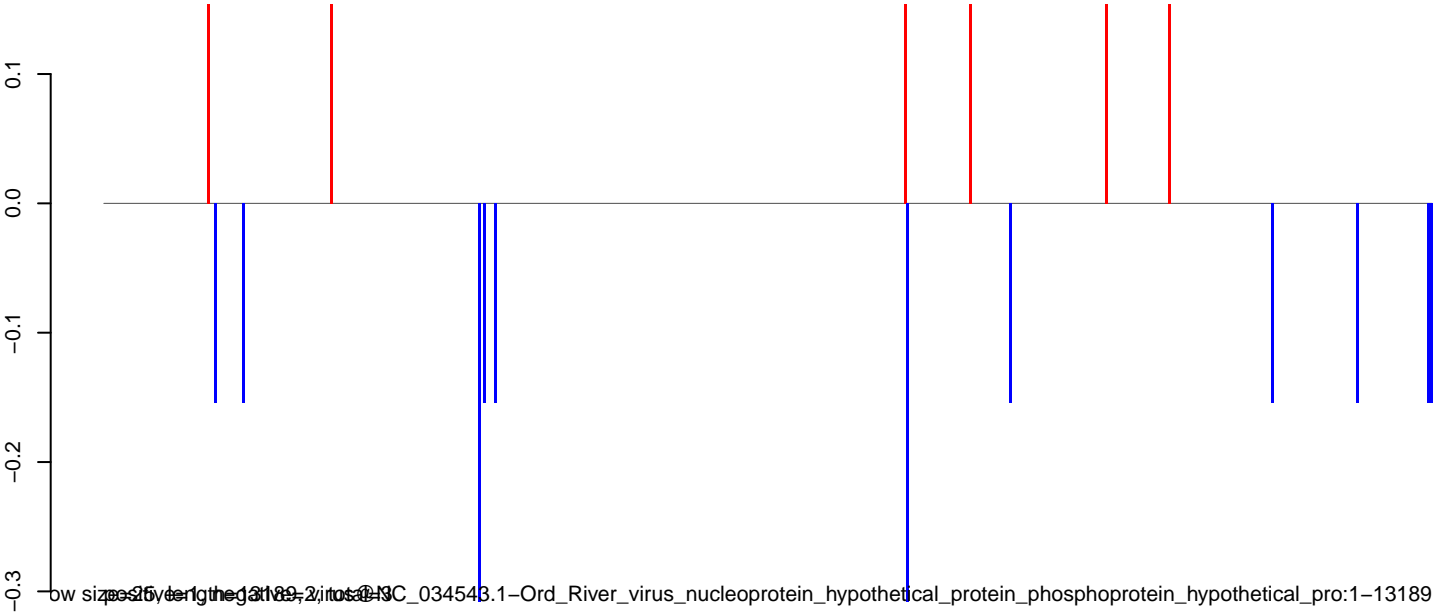
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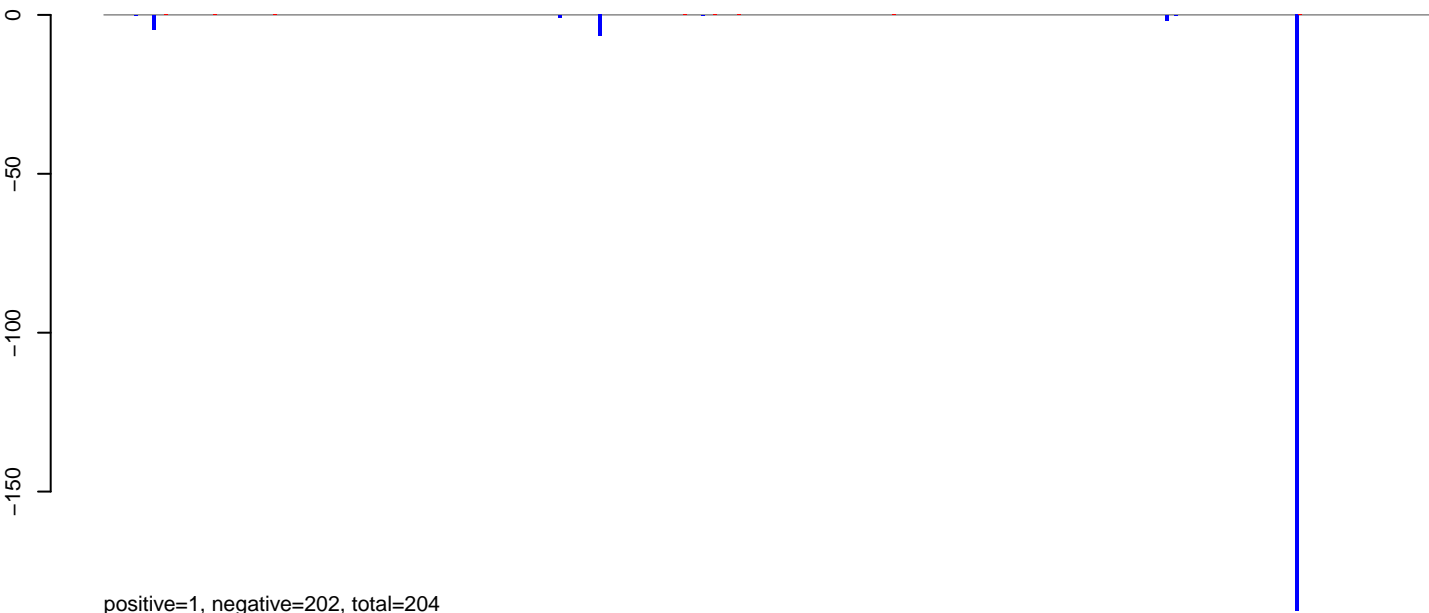
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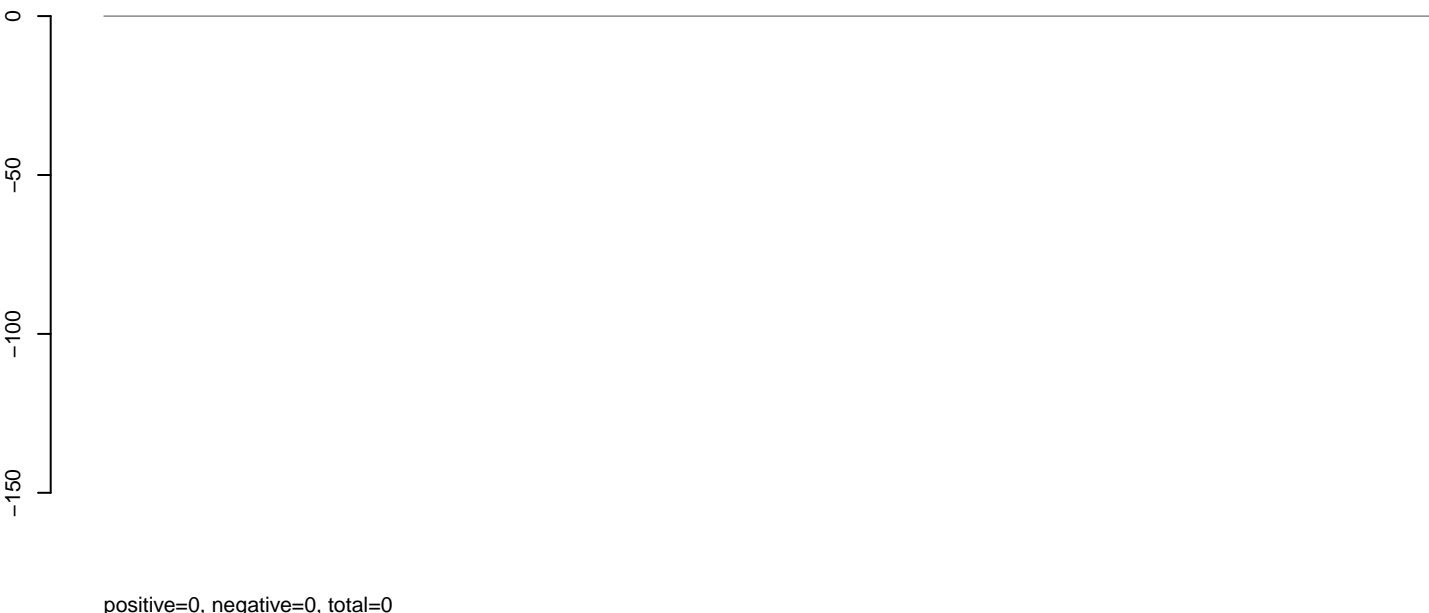
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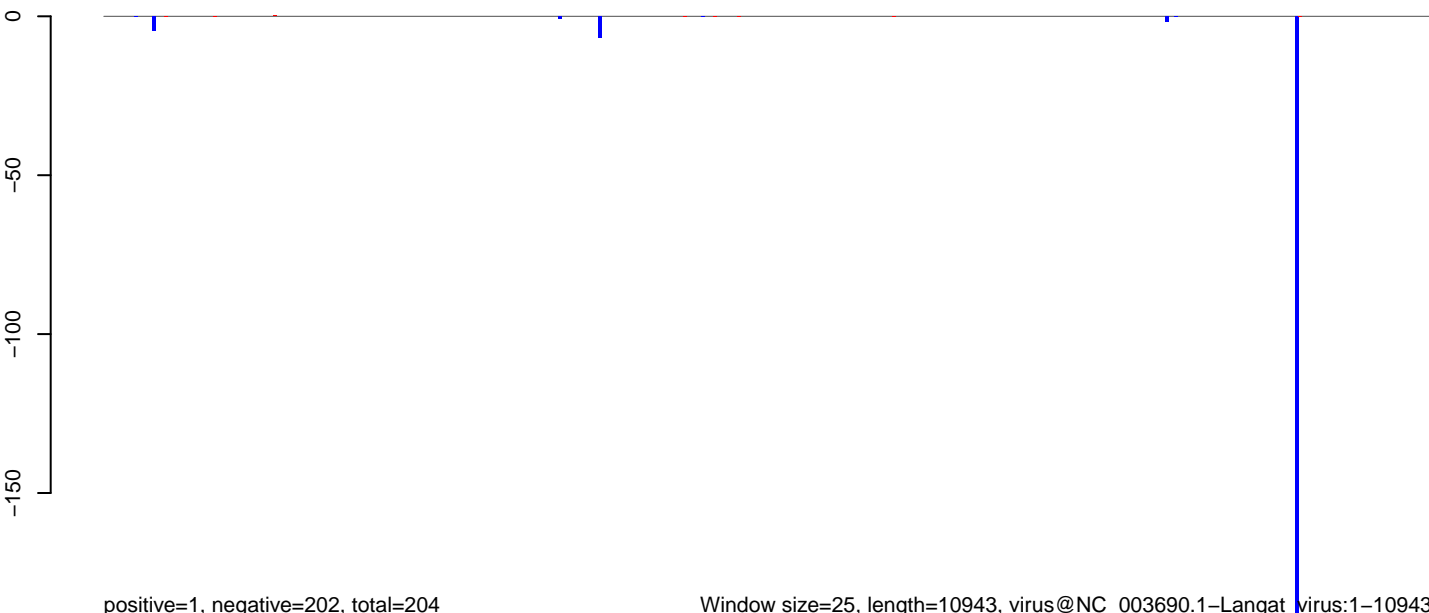
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AeAlbo_C636_6dpi_SINV1.24_35.rep



positive=0, negative=0, total=0

AeAlbo_C636_6dpi_SINV1.rep

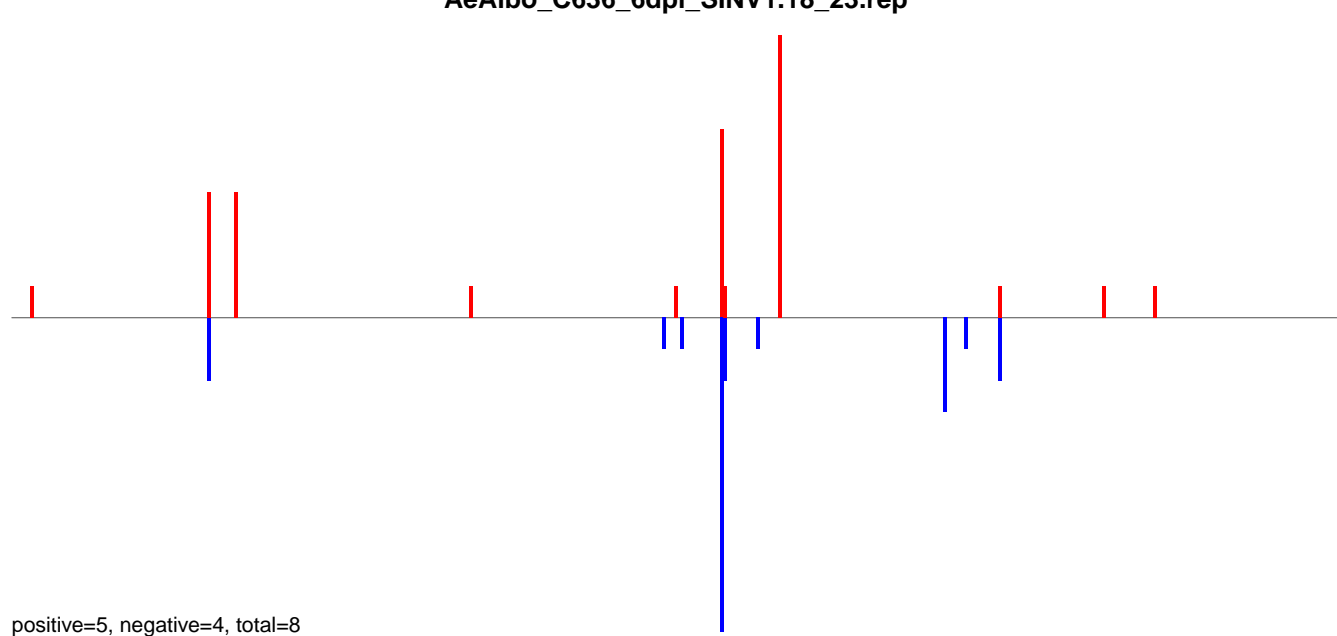


positive=1, negative=202, total=204

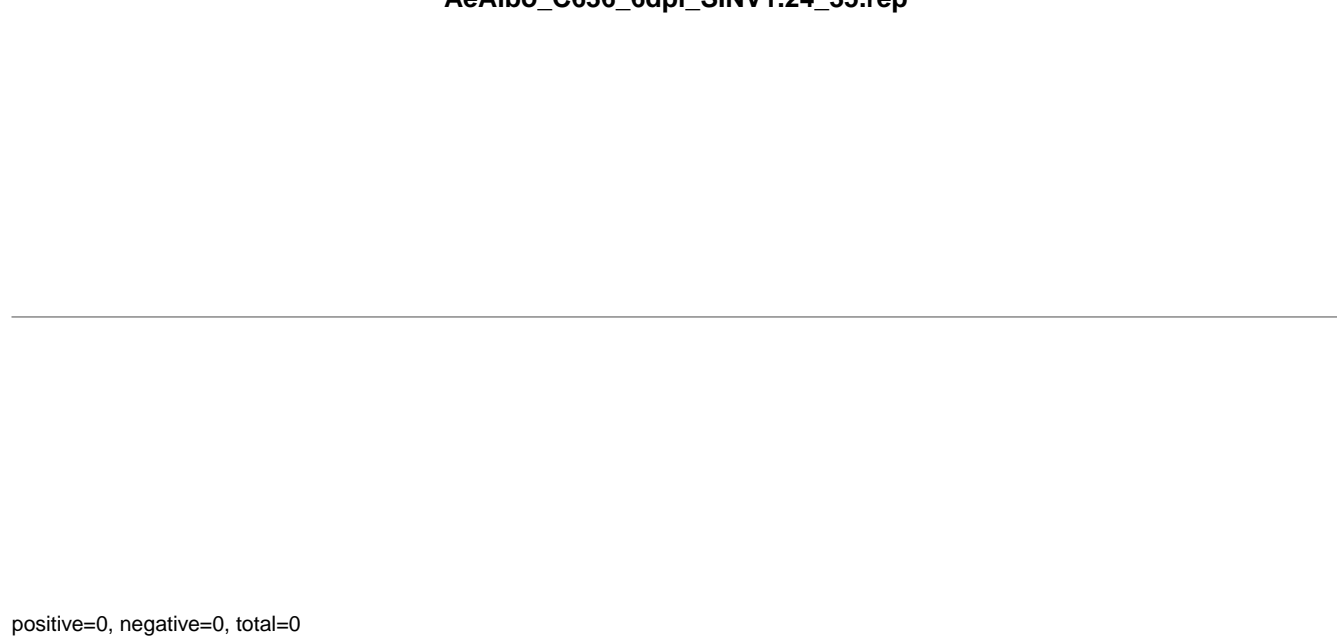
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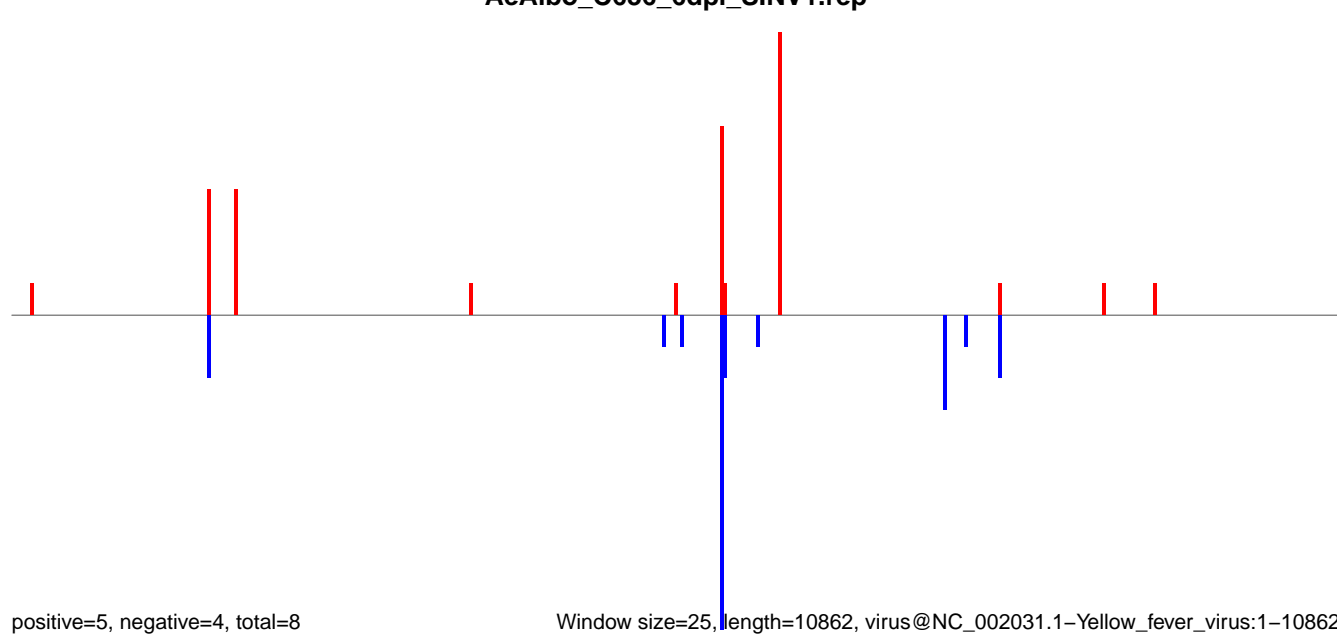
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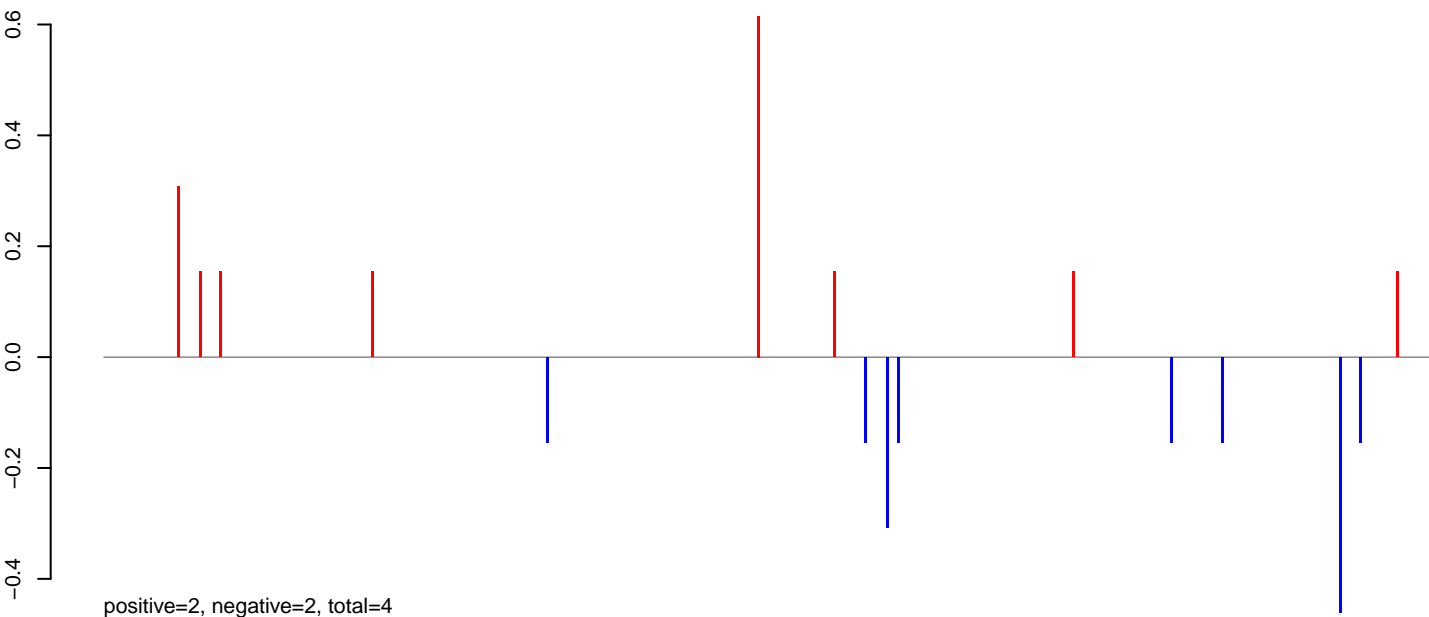


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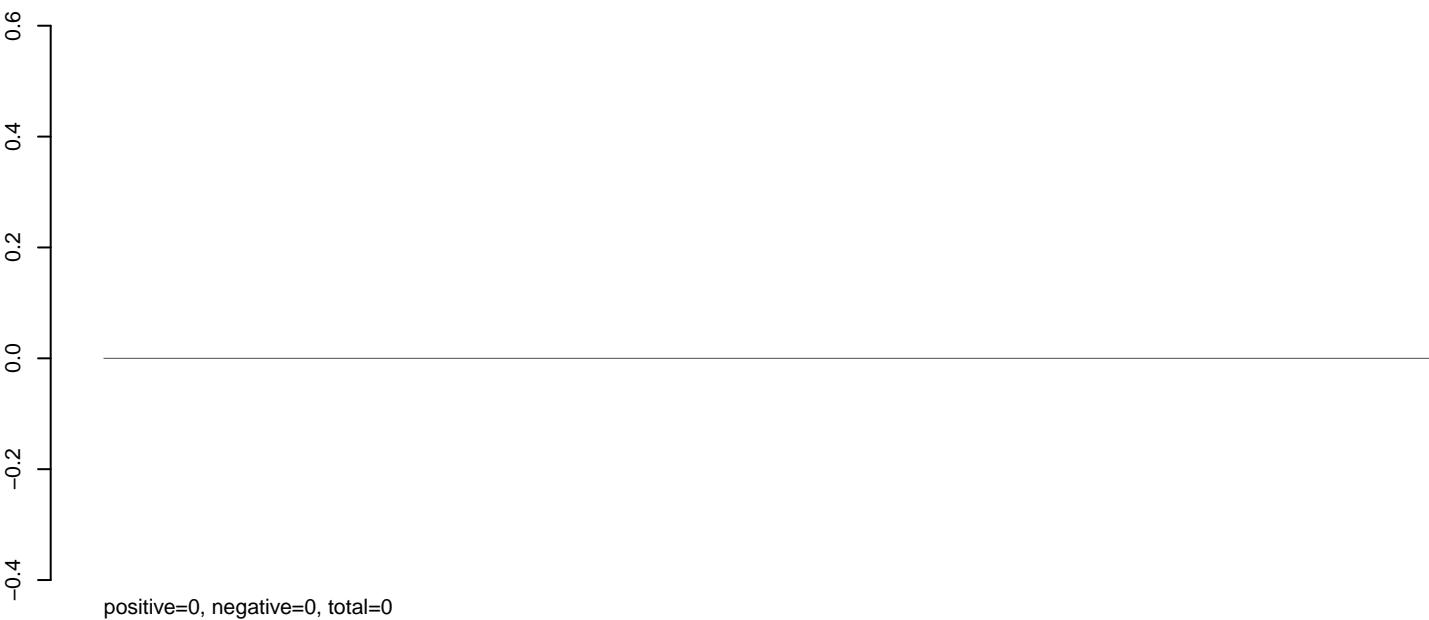


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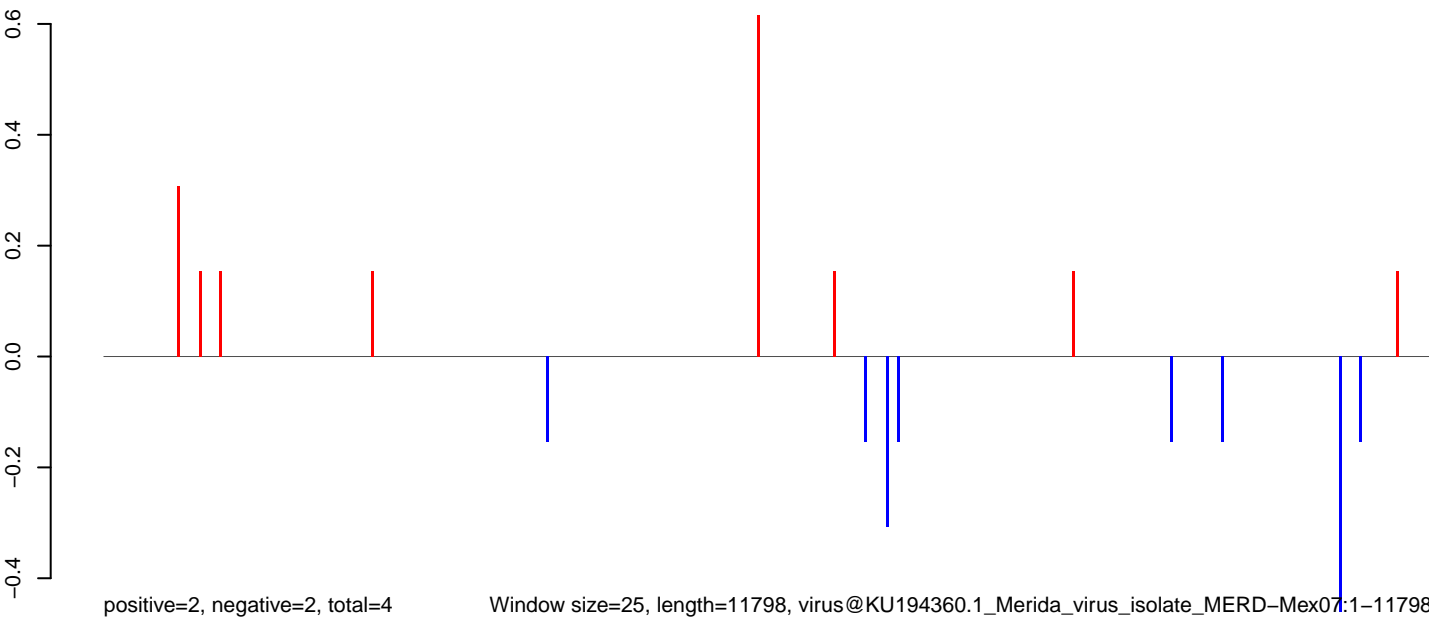
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AeAlbo_C636_6dpi_SINV1.24_35.rep



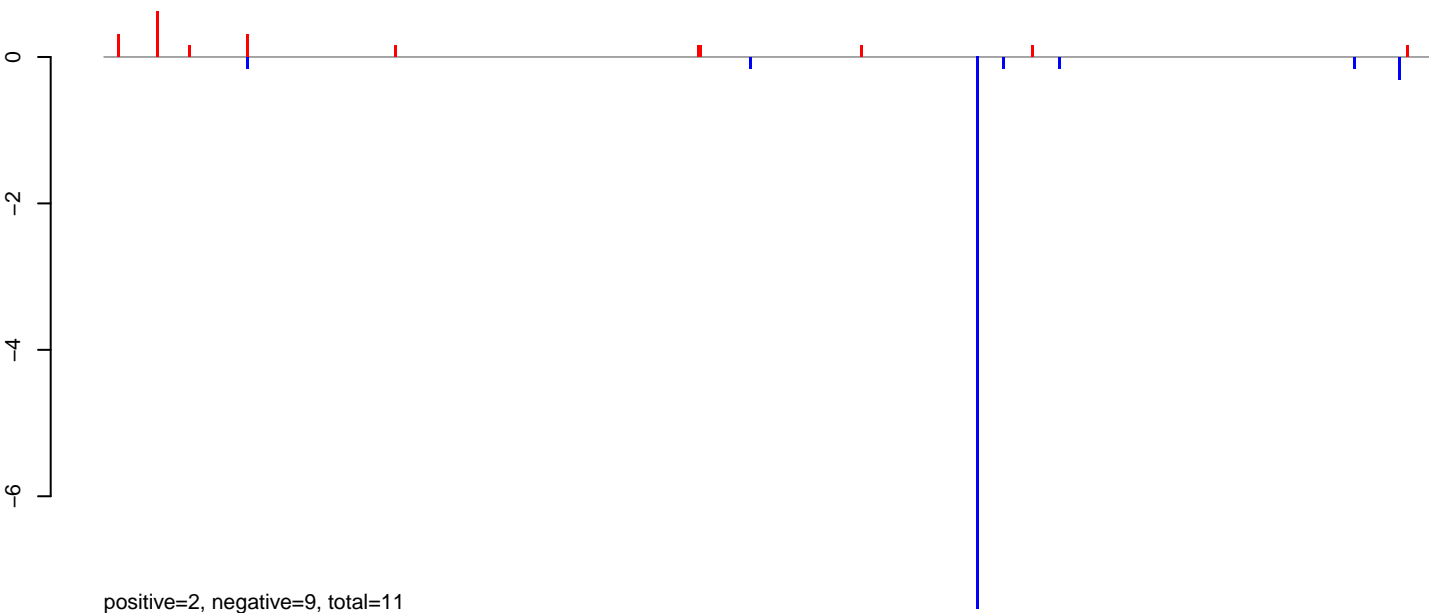
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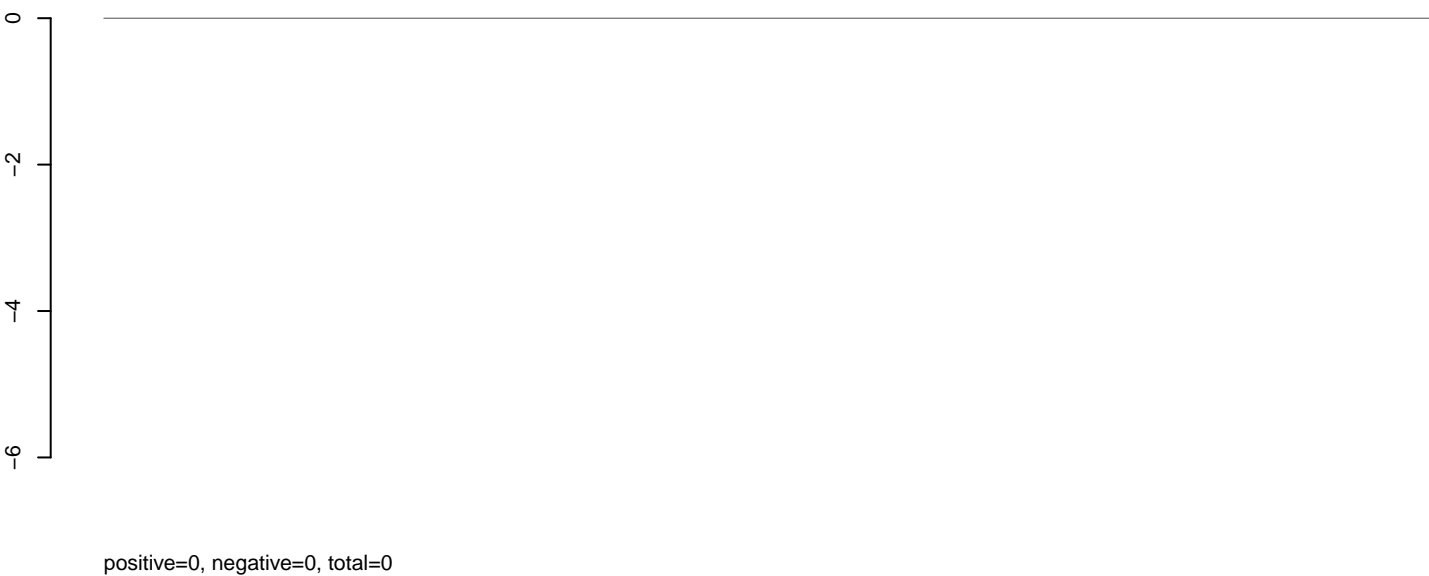
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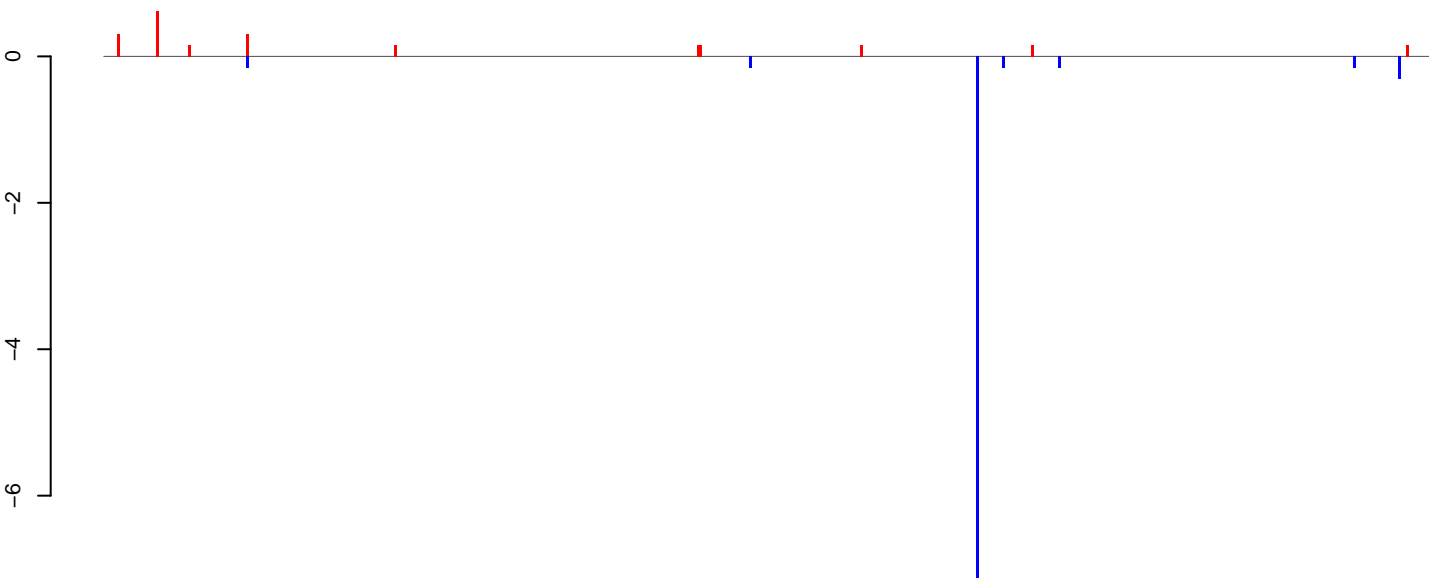
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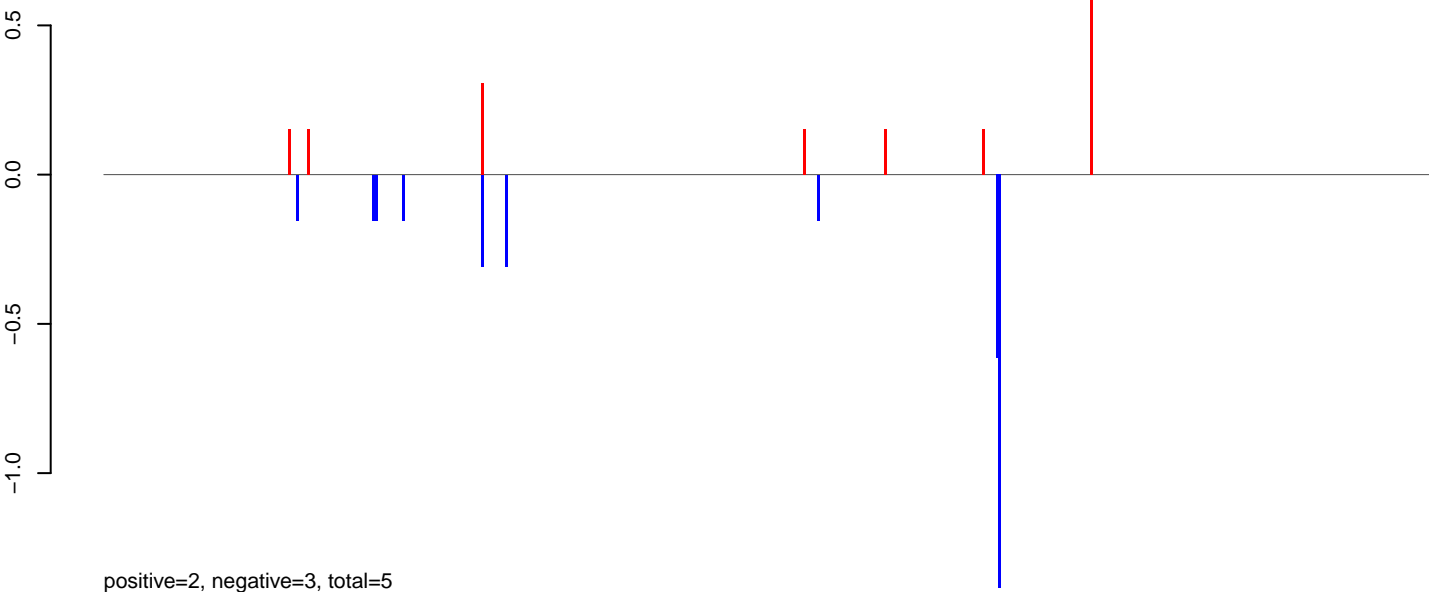
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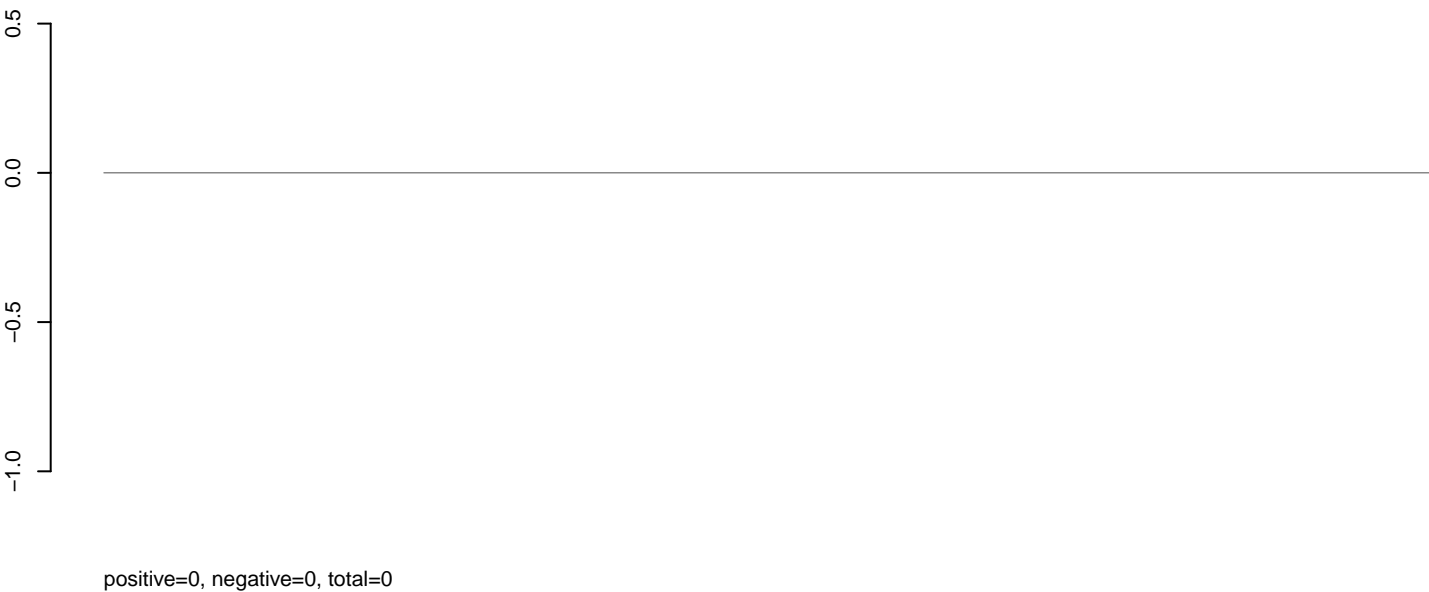
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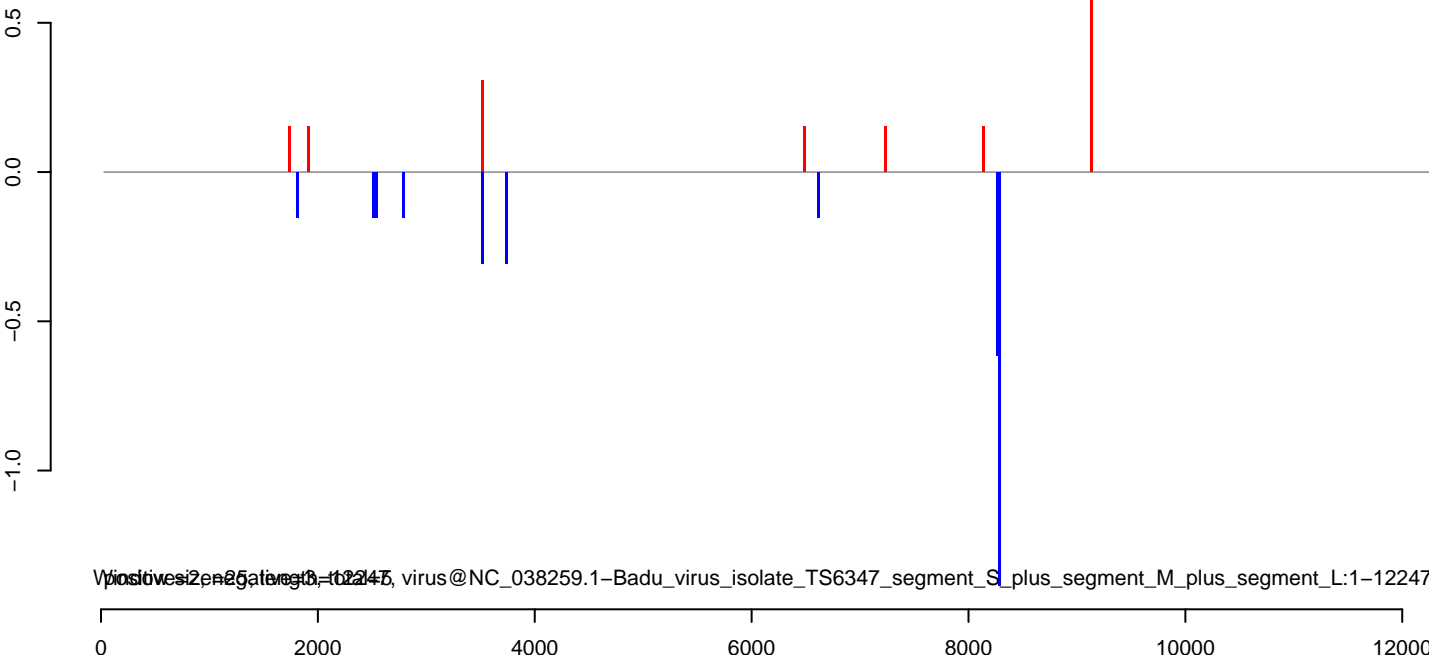
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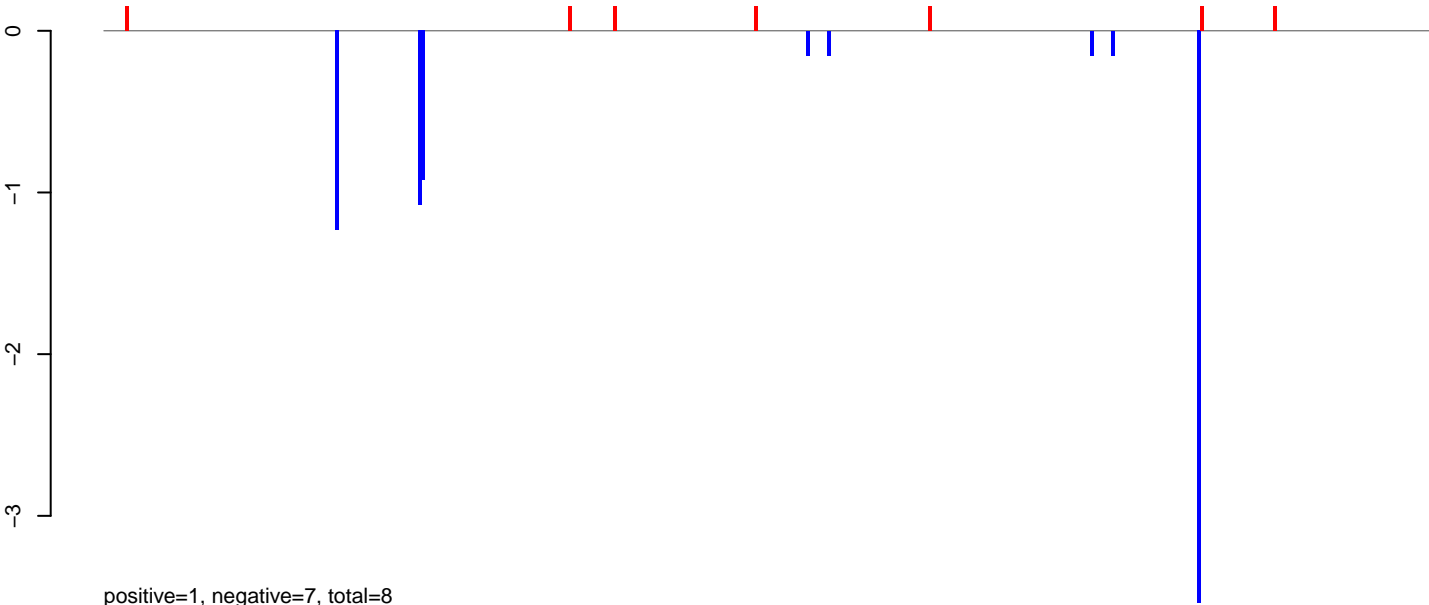
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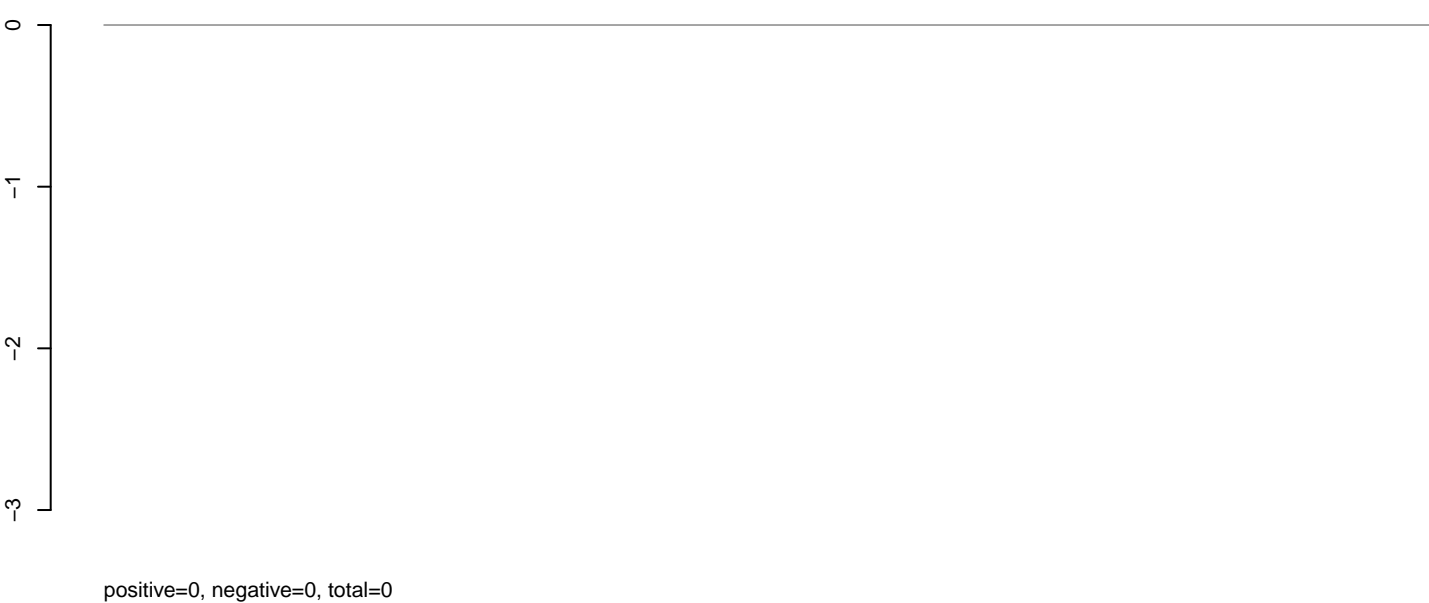
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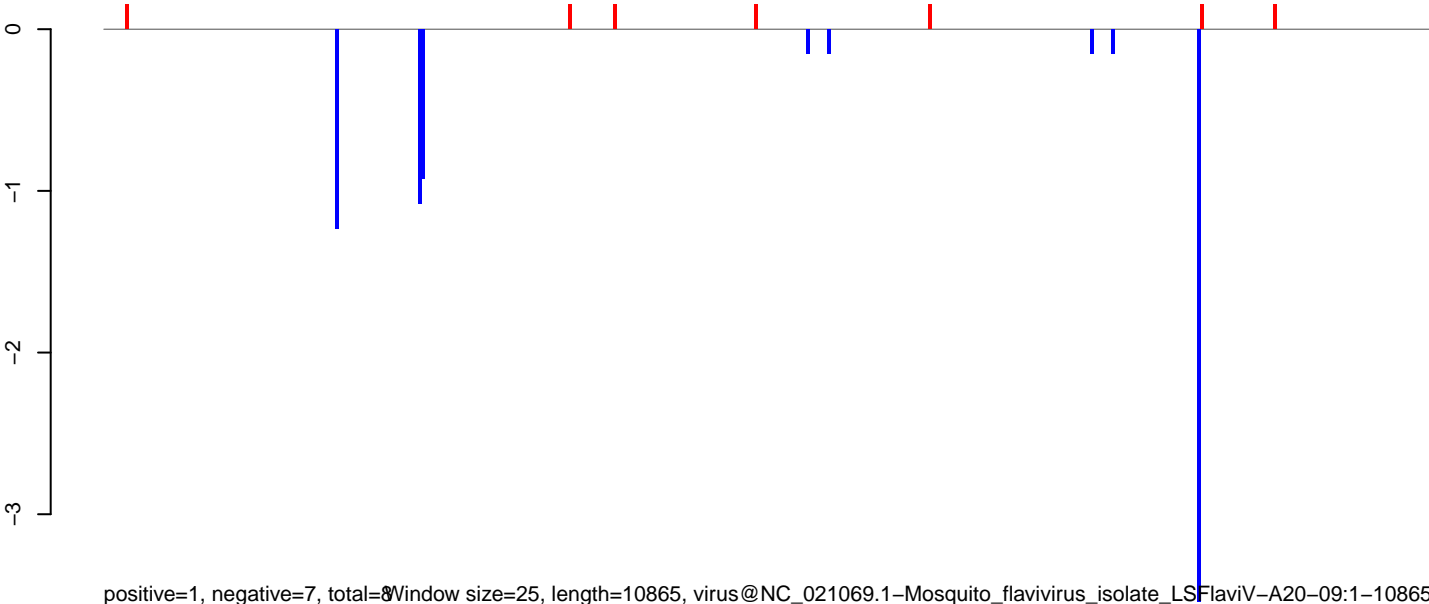
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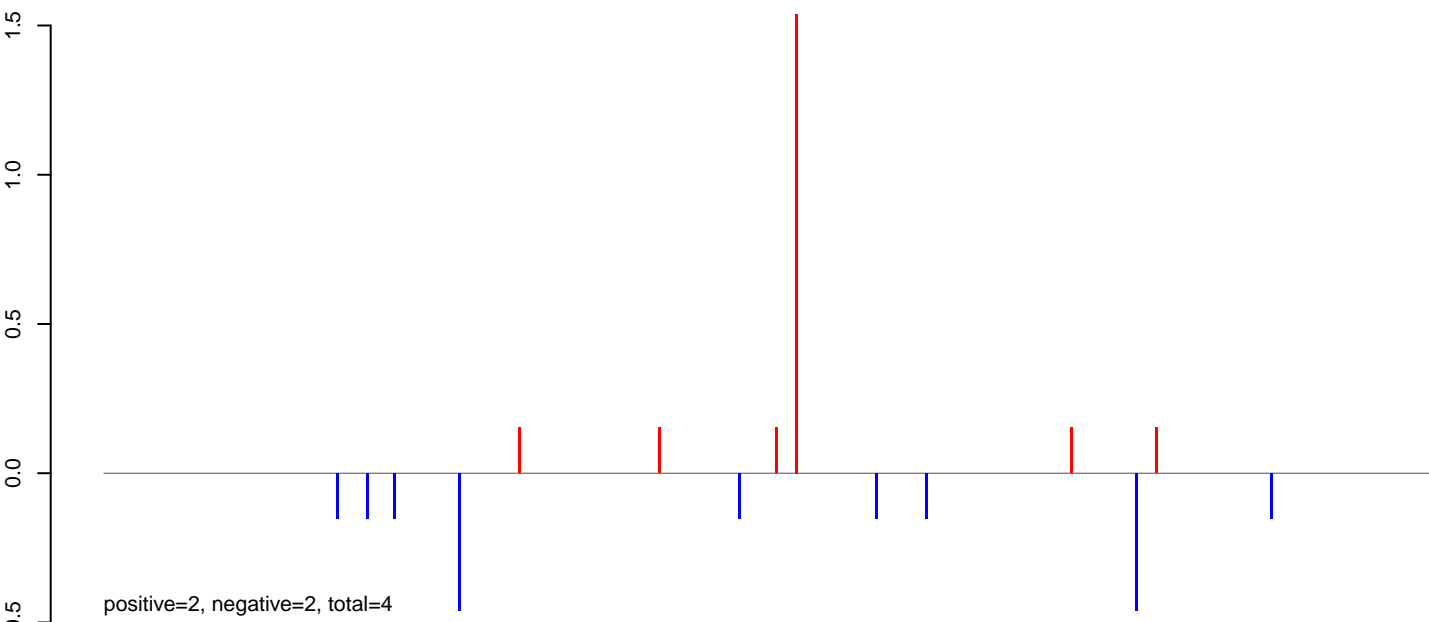
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AeAlbo_C636_6dpi_SINV1.rep



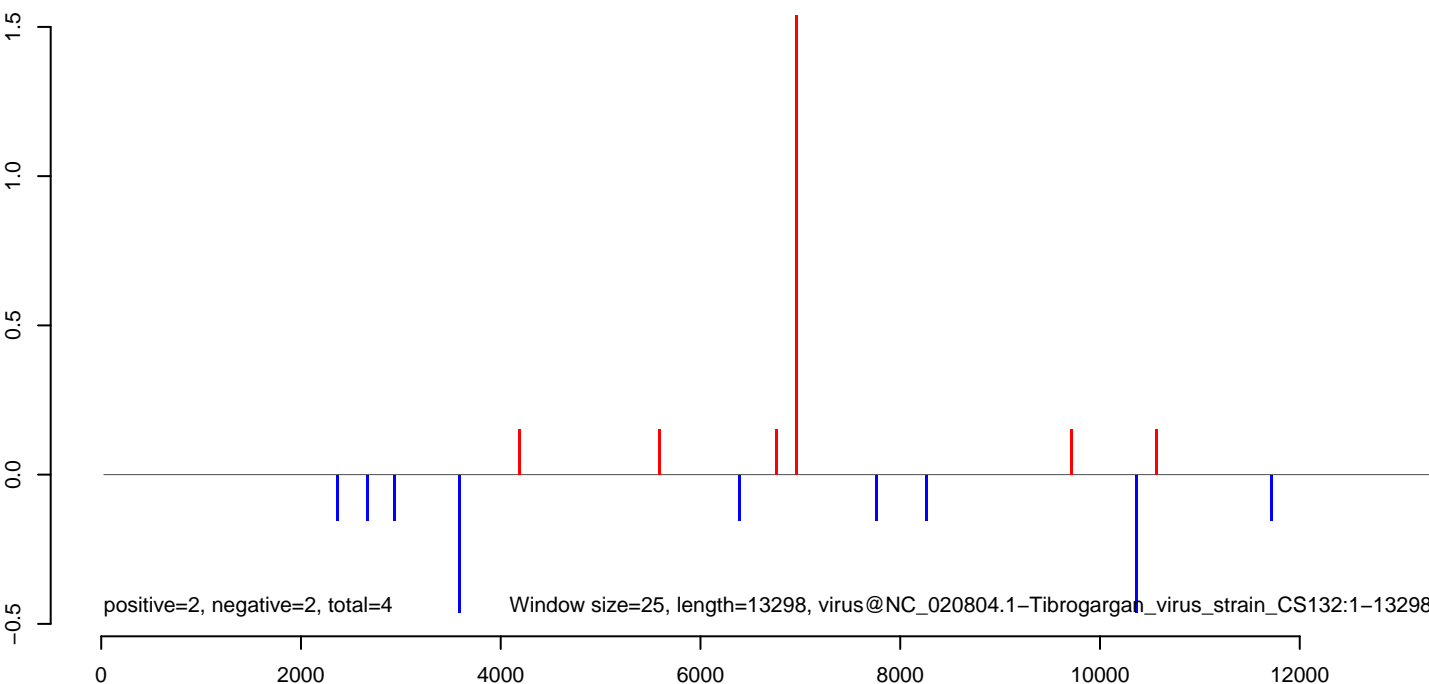
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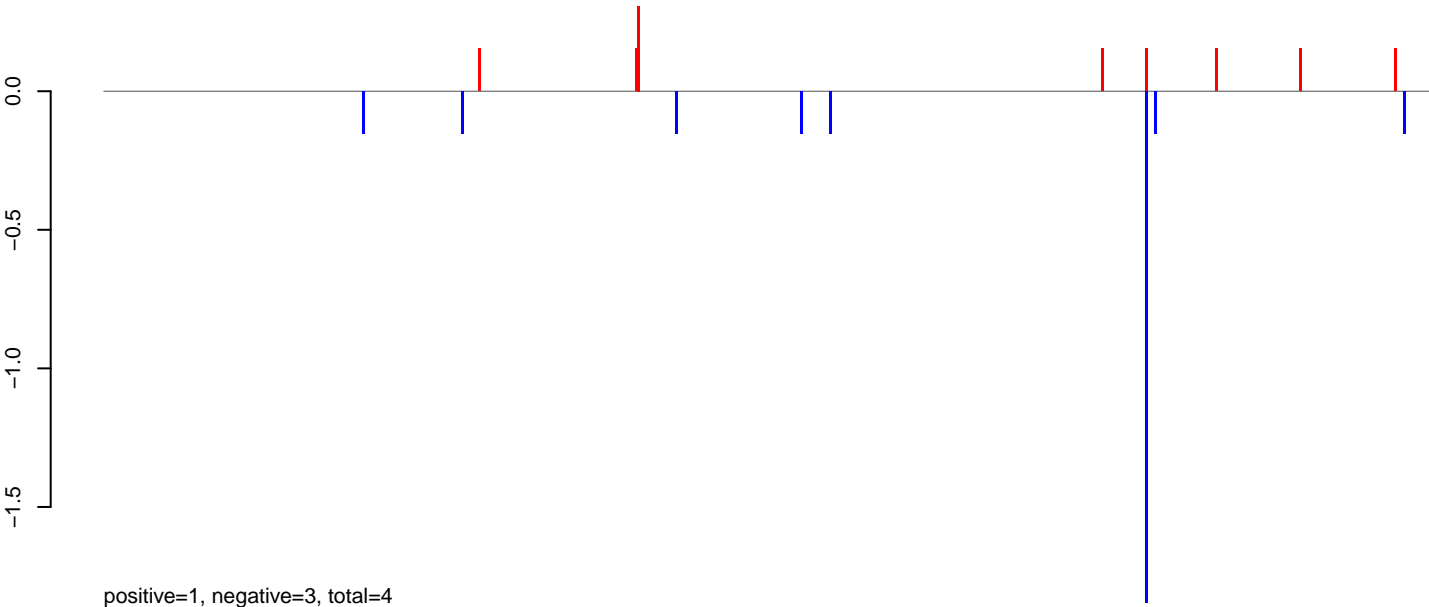
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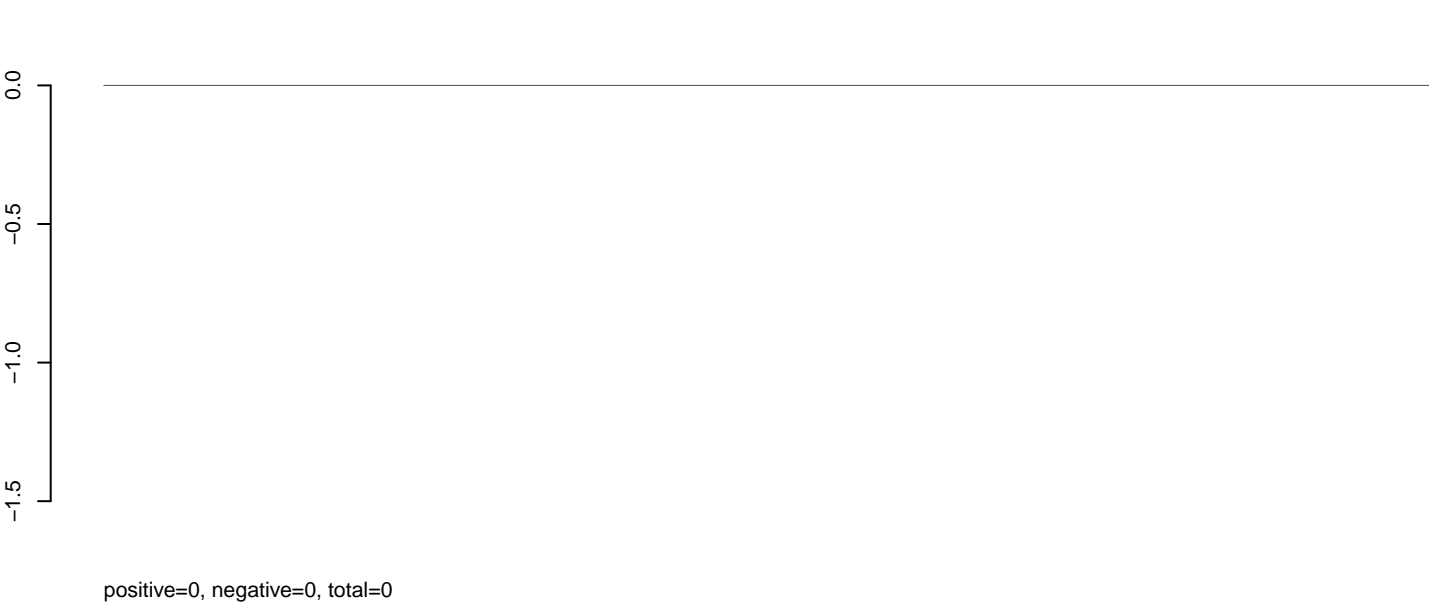
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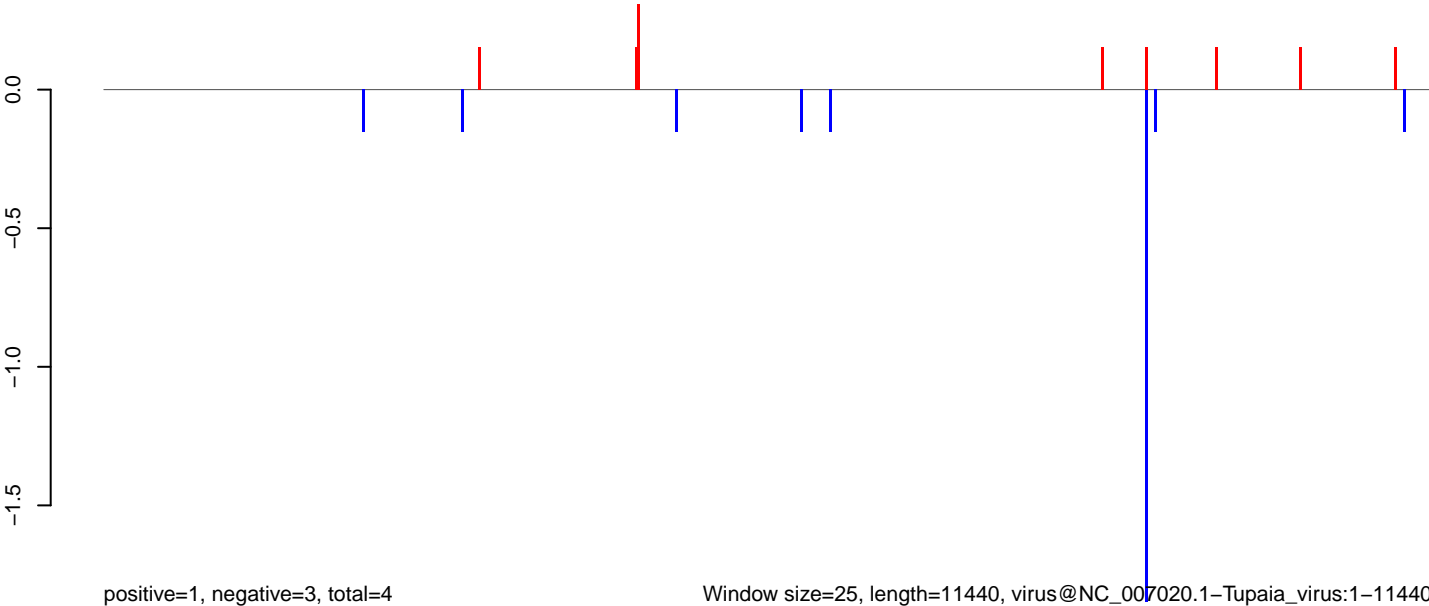
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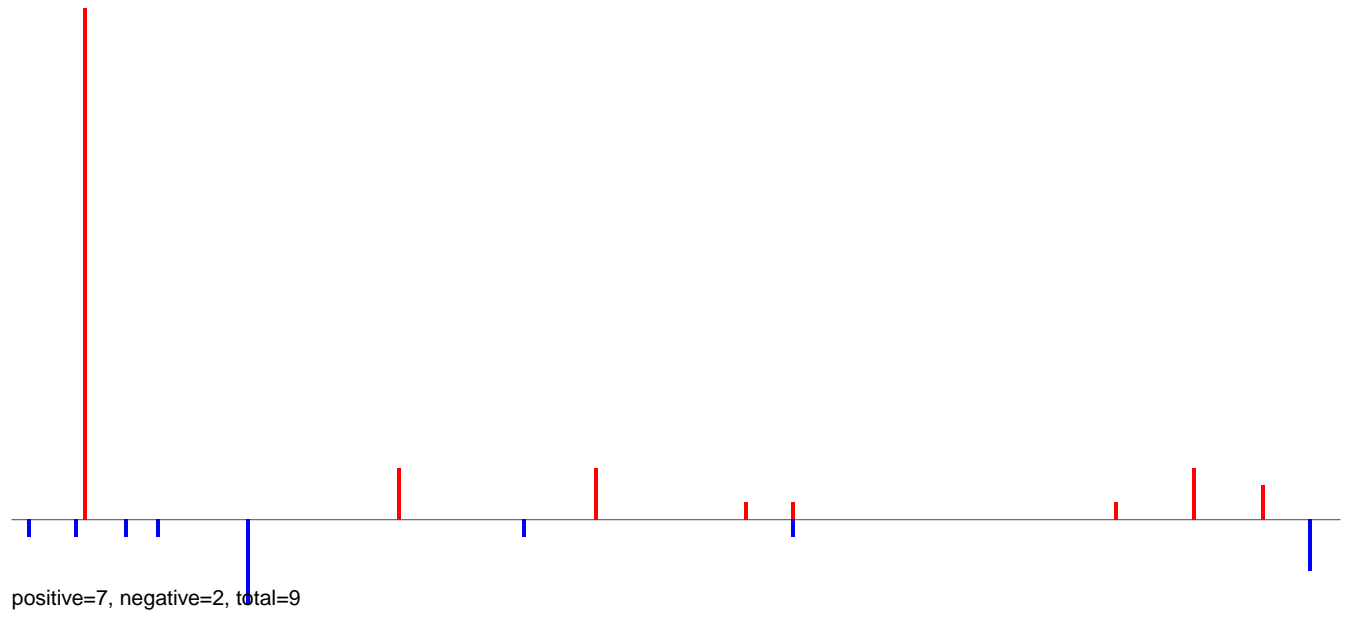
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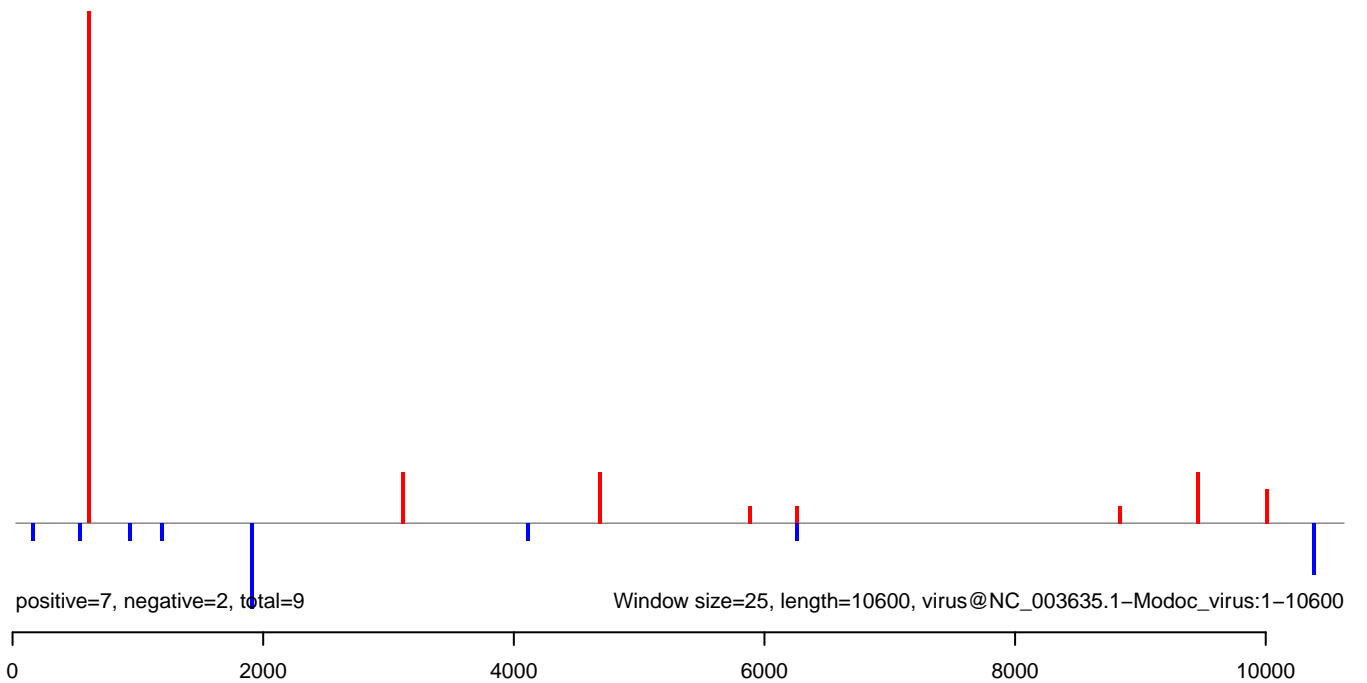
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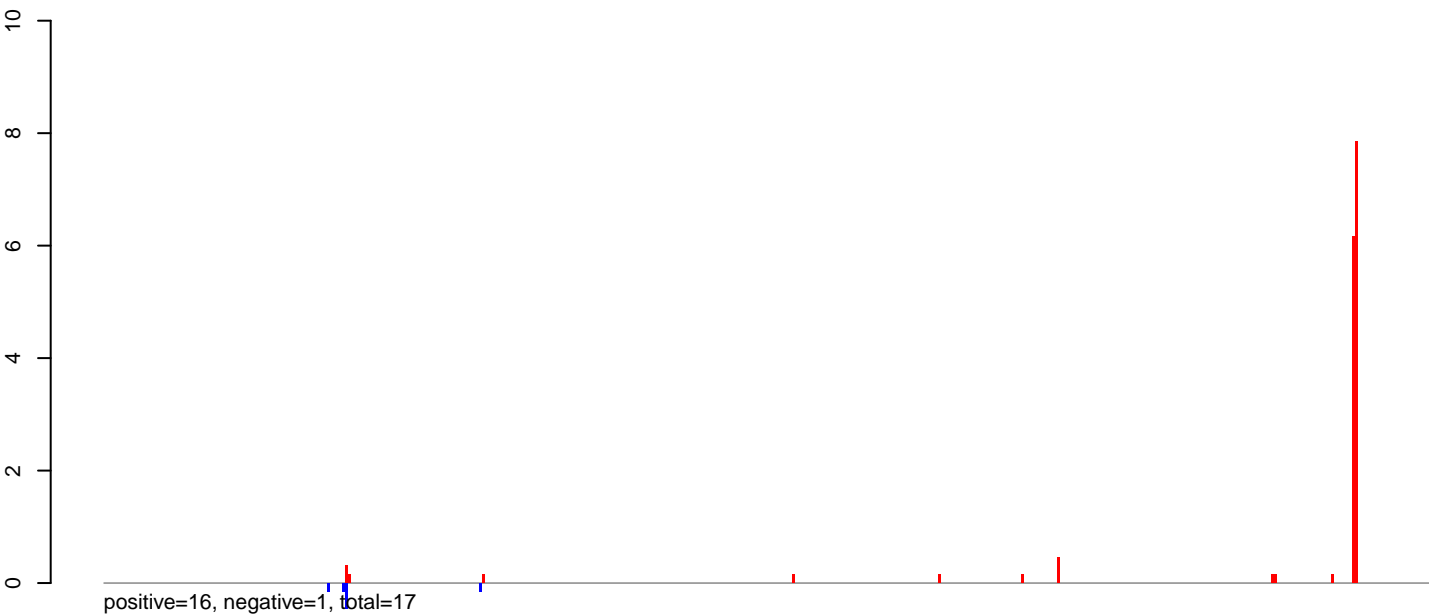
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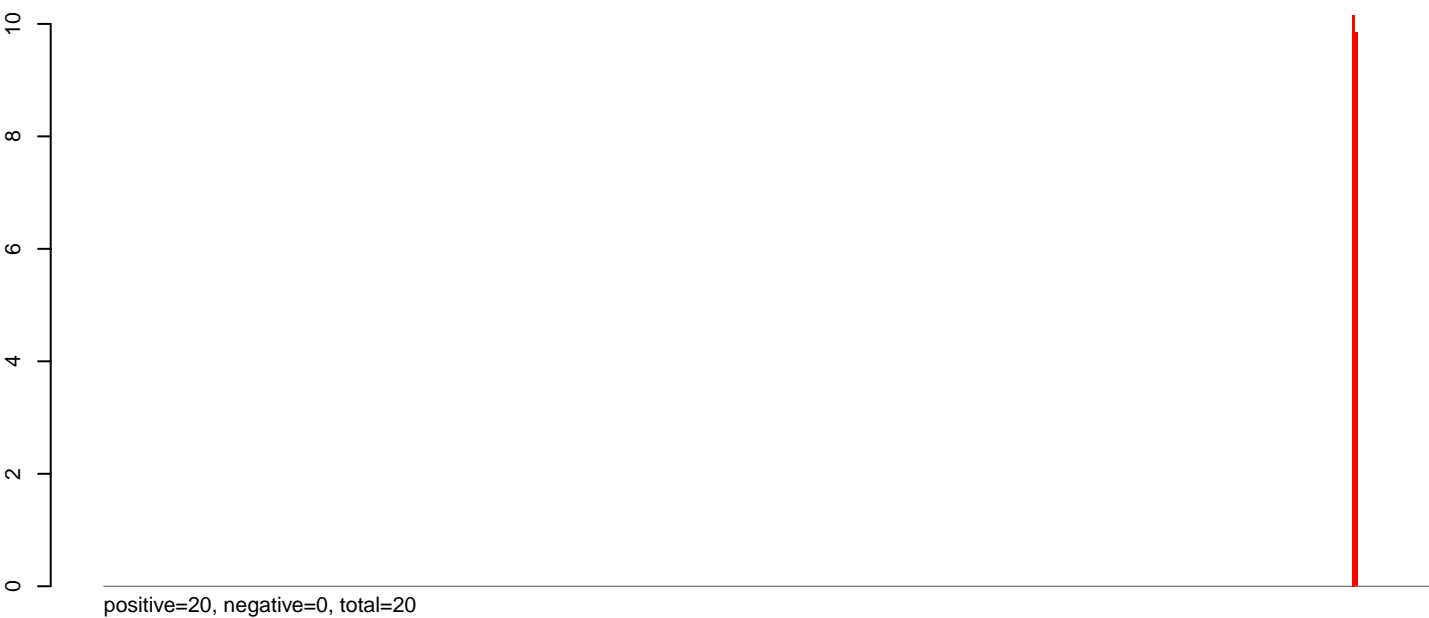
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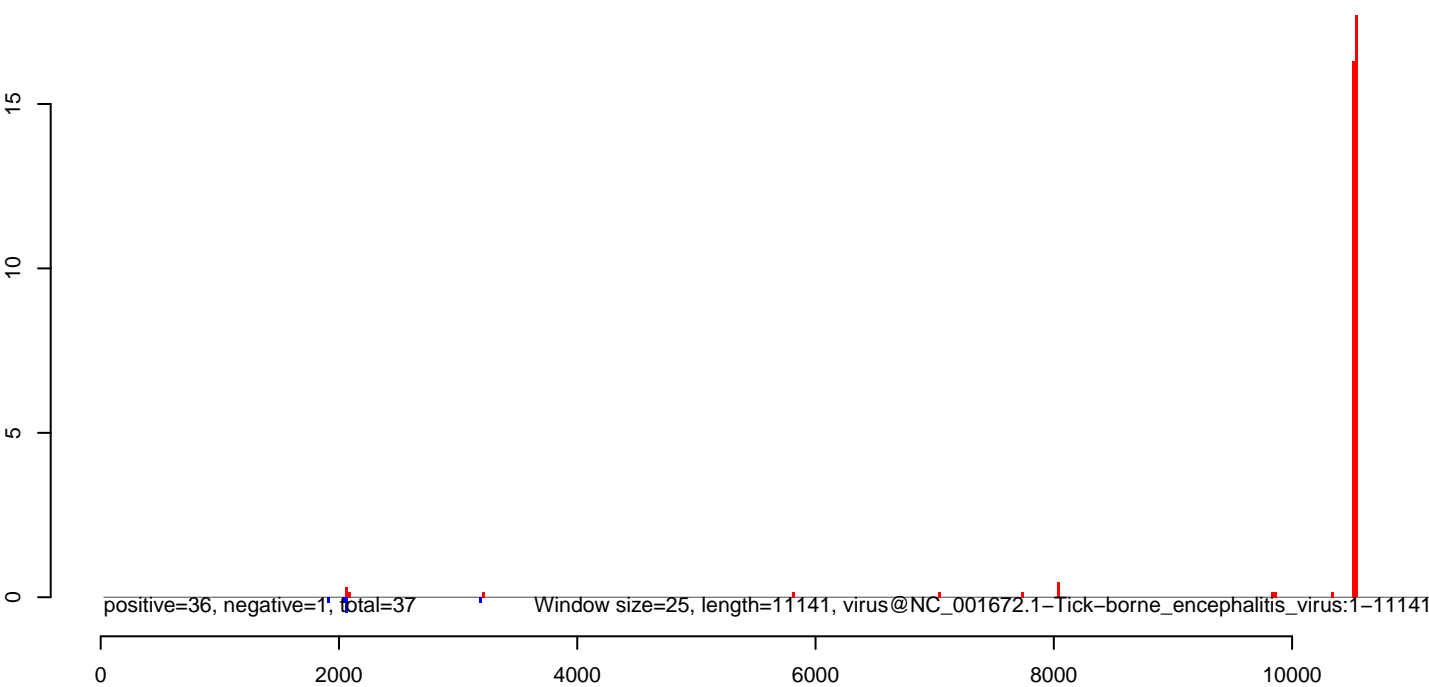
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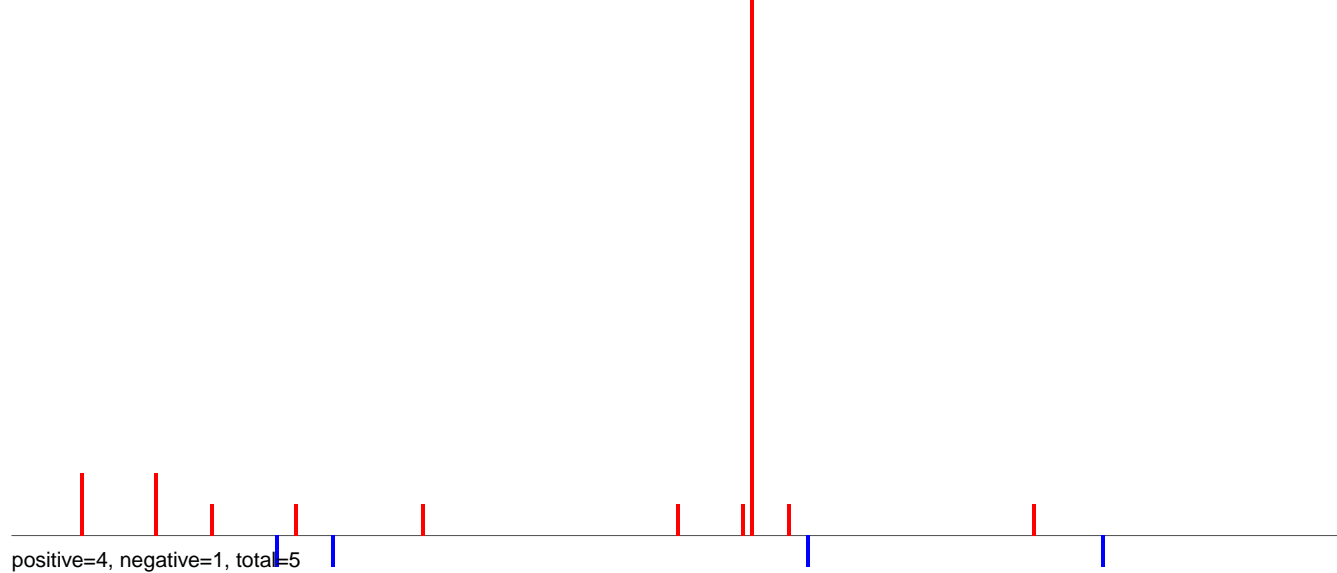
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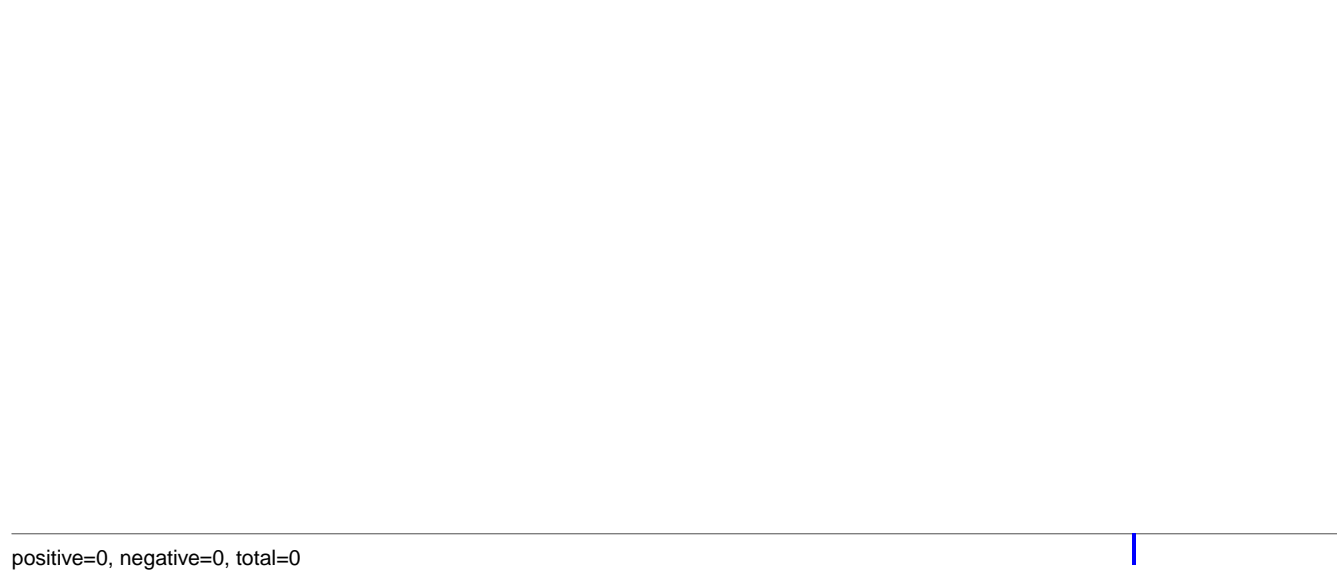
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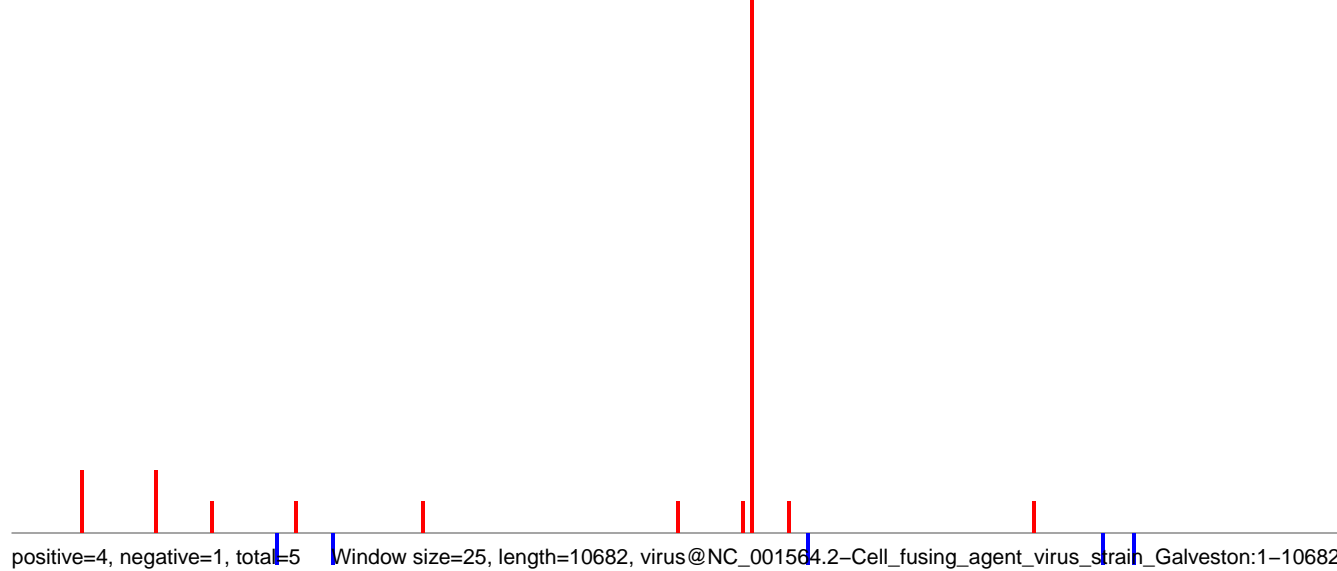
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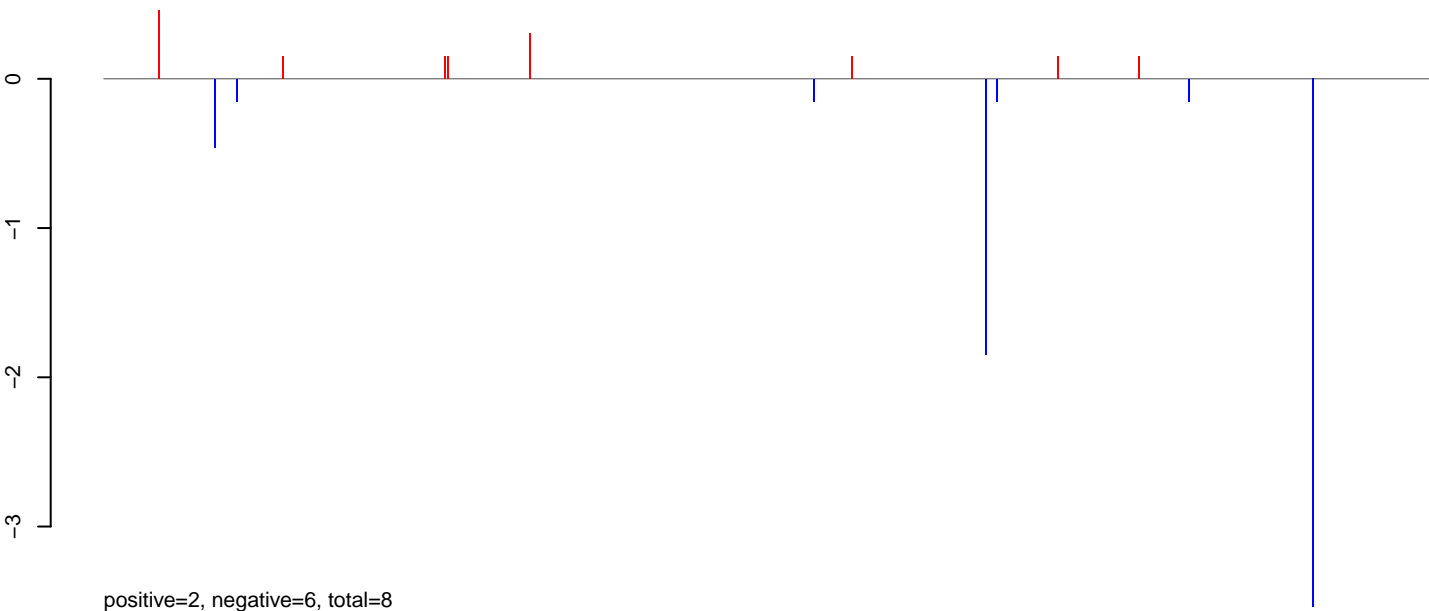


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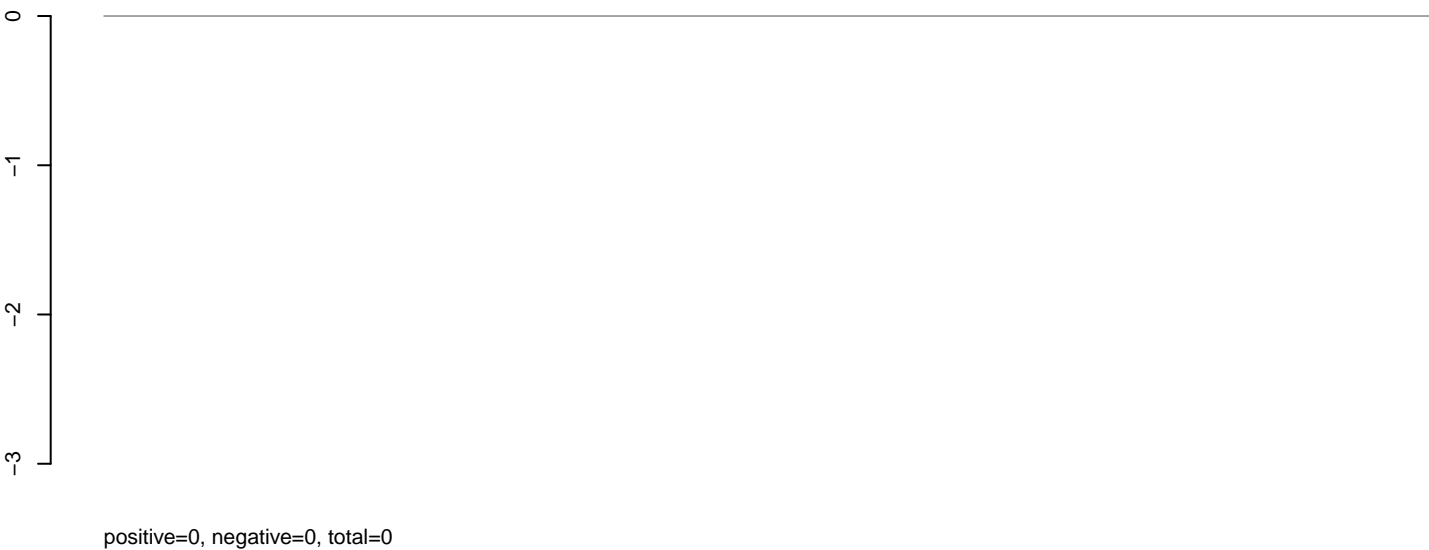


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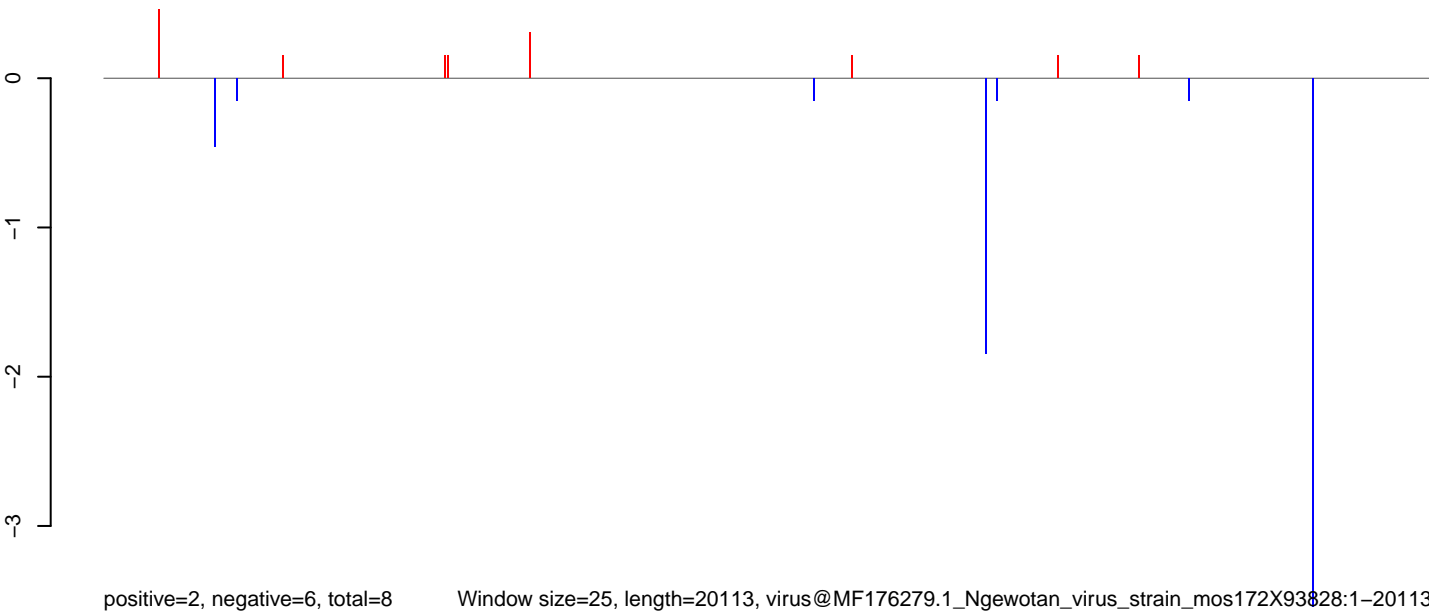
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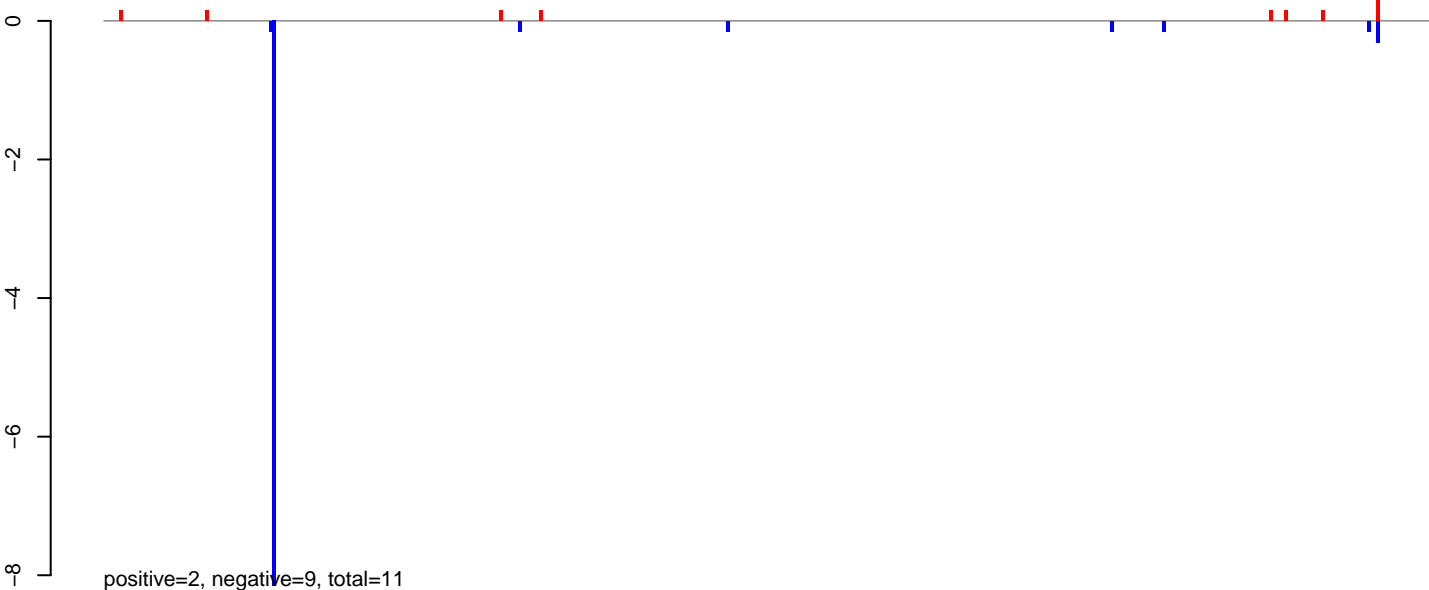


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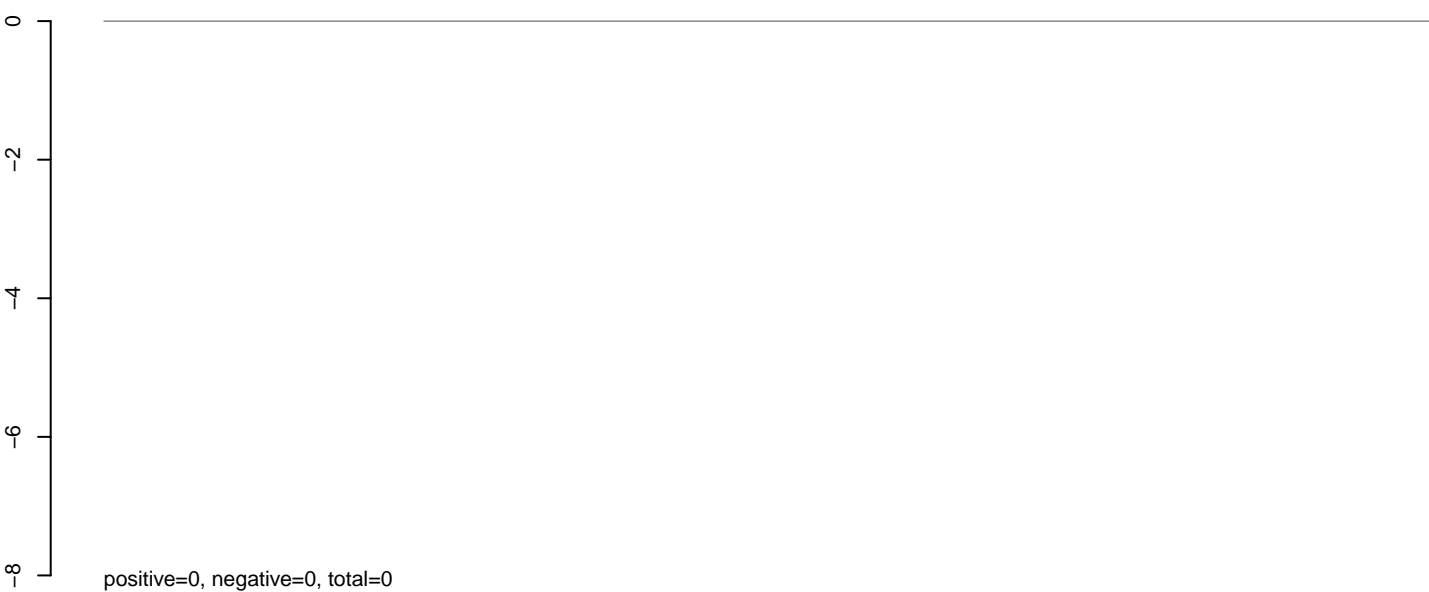


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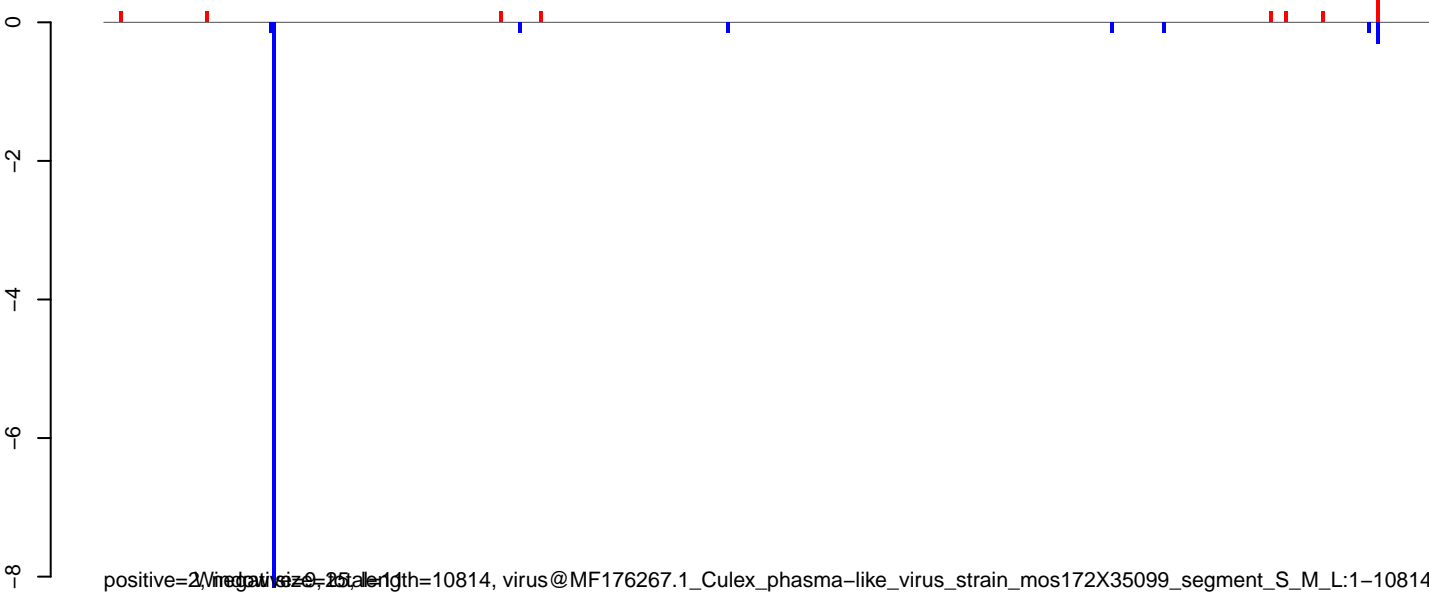
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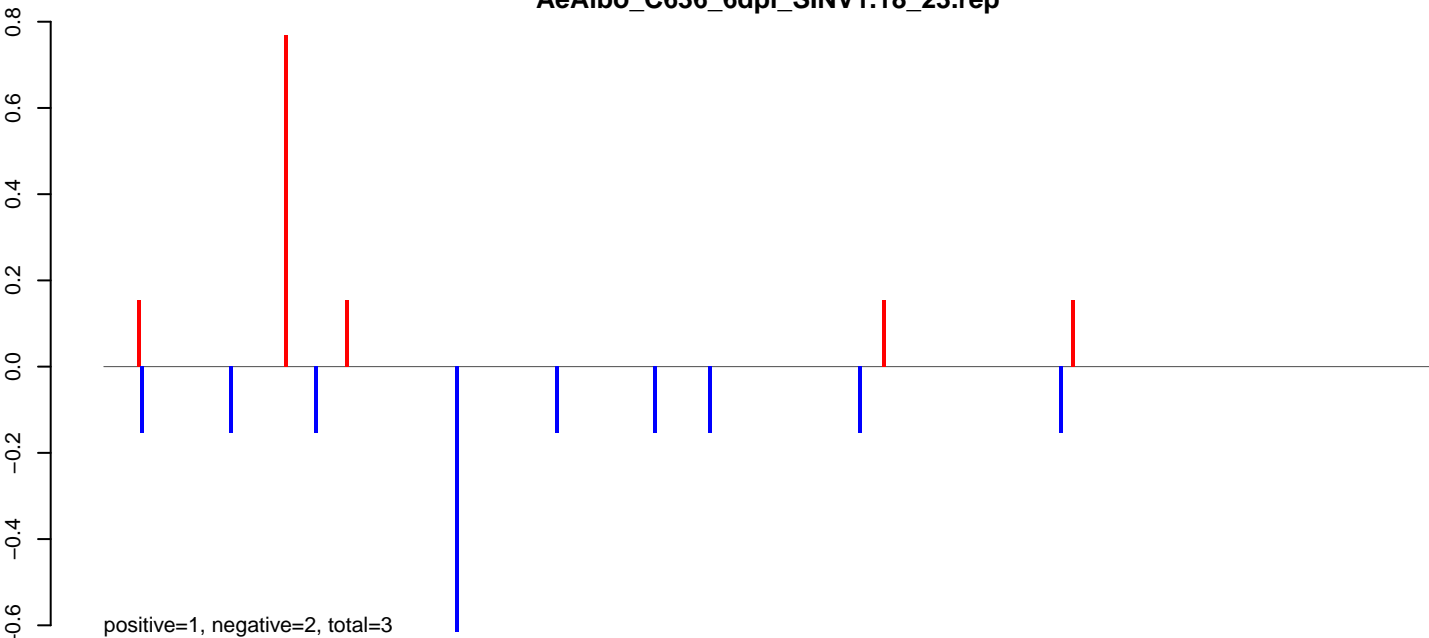
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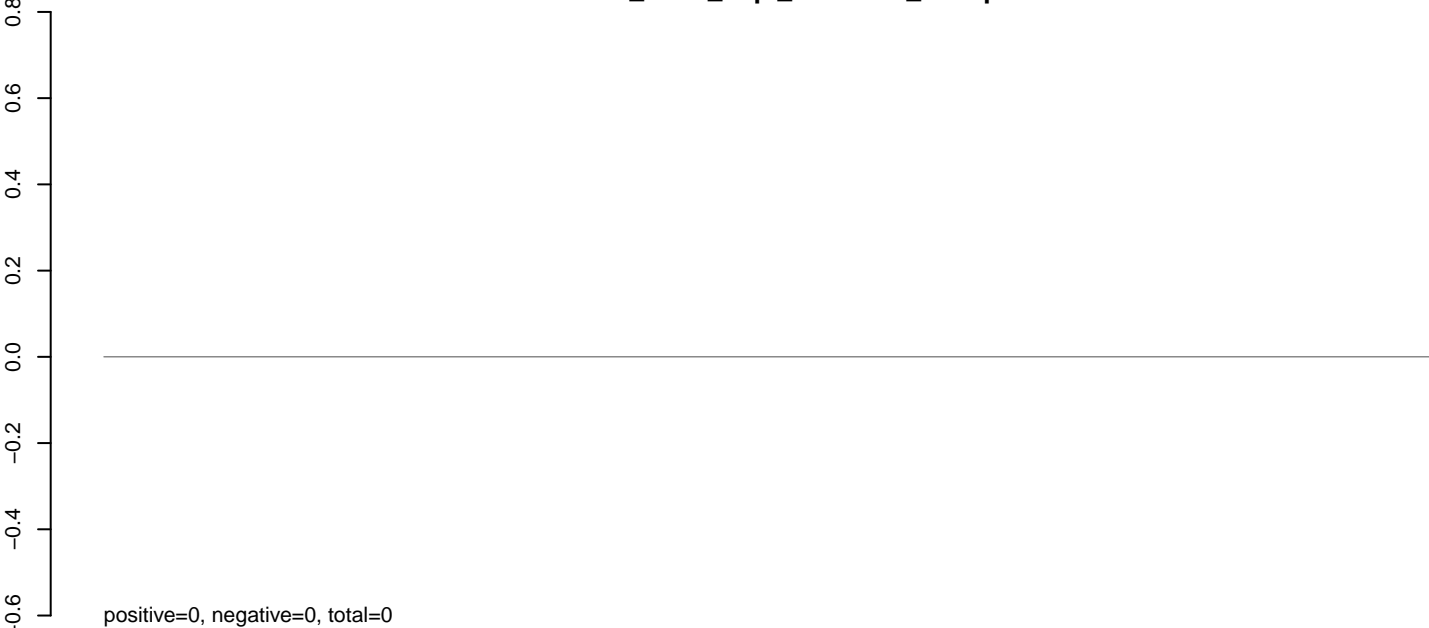
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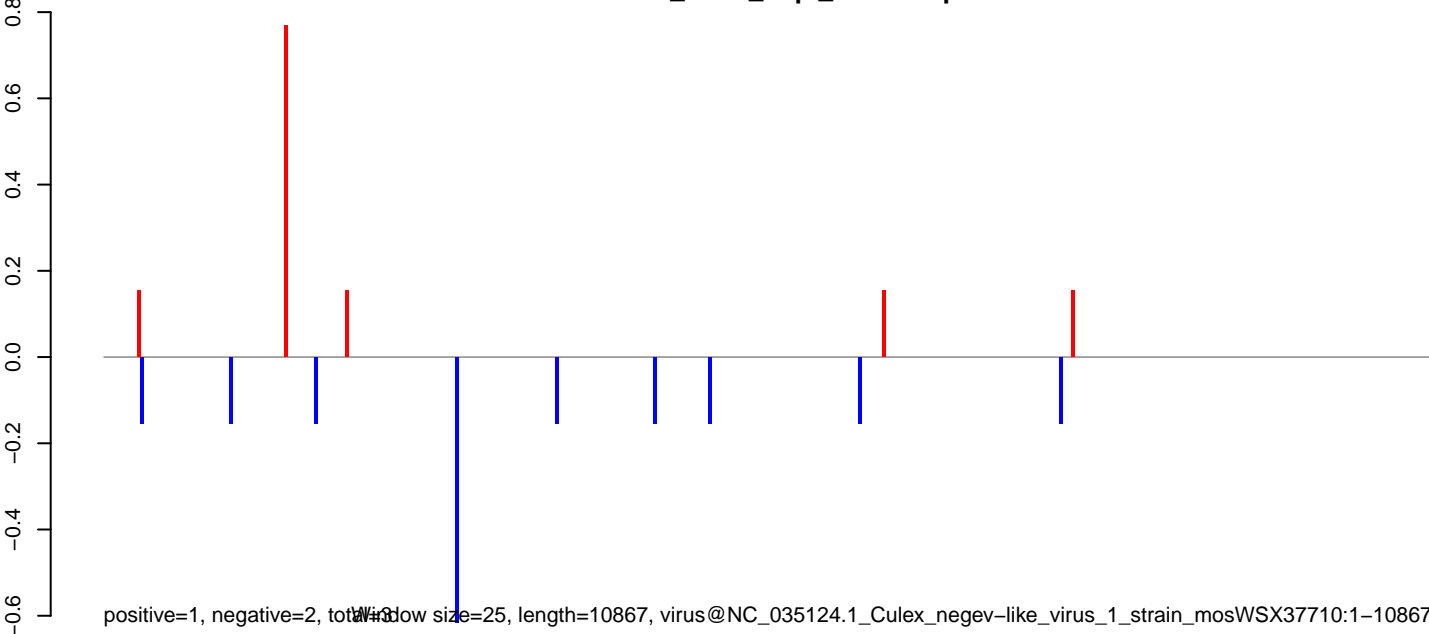
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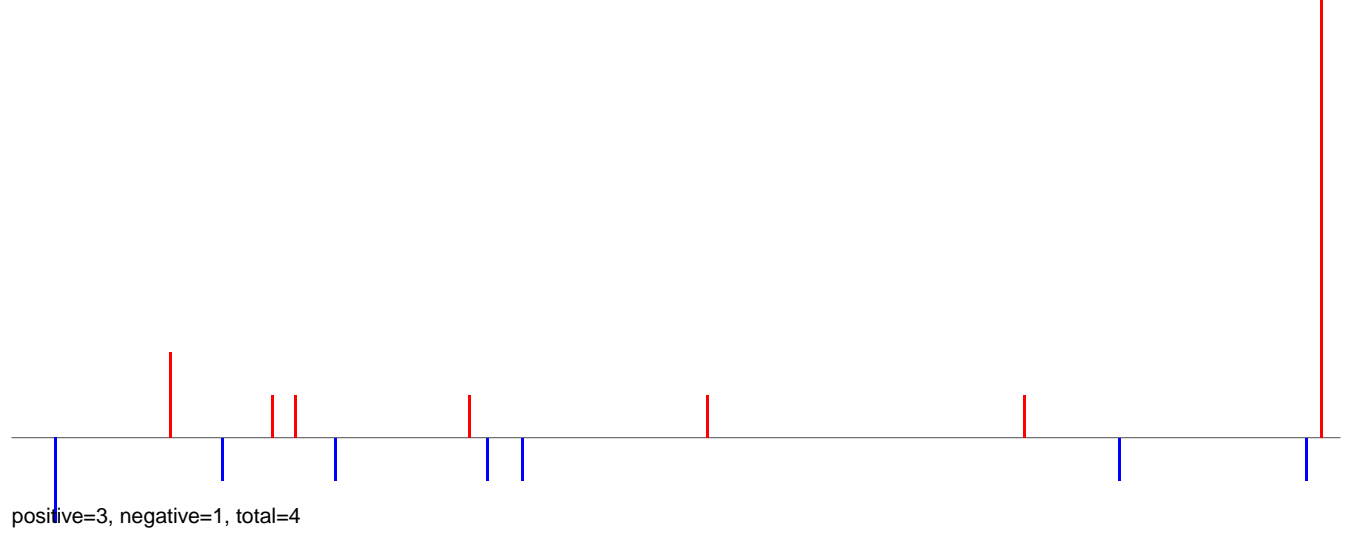
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AeAlbo_C636_6dpi_SINV1.rep



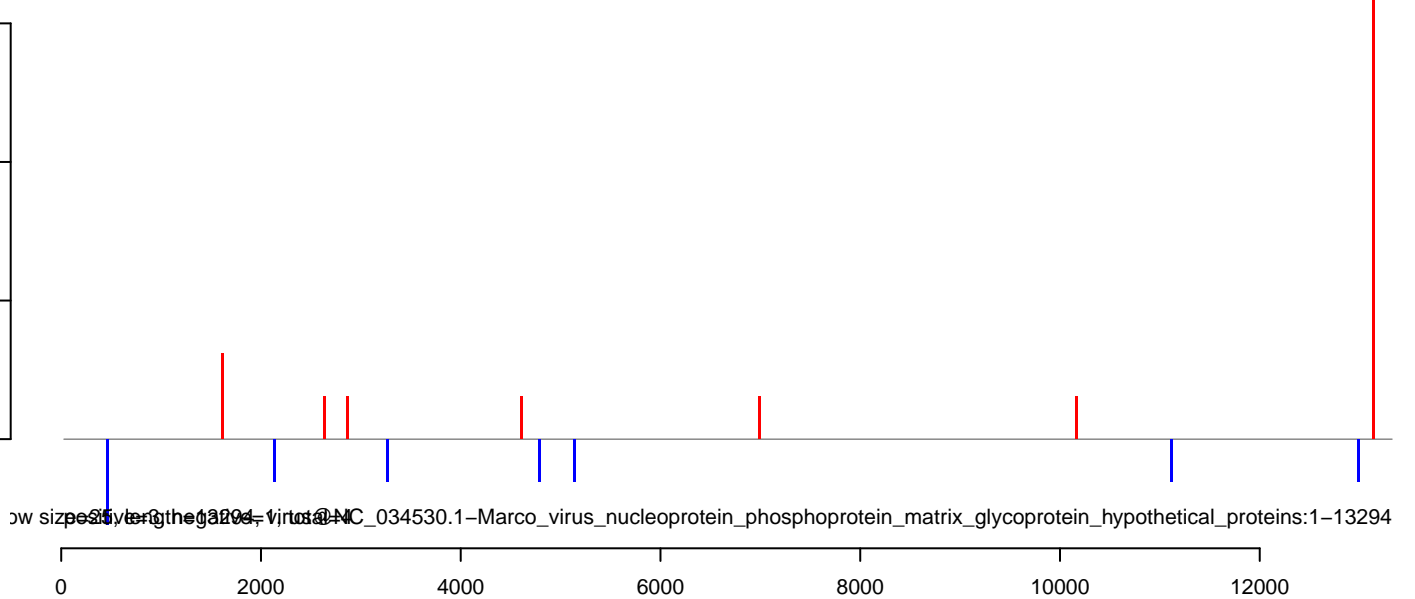
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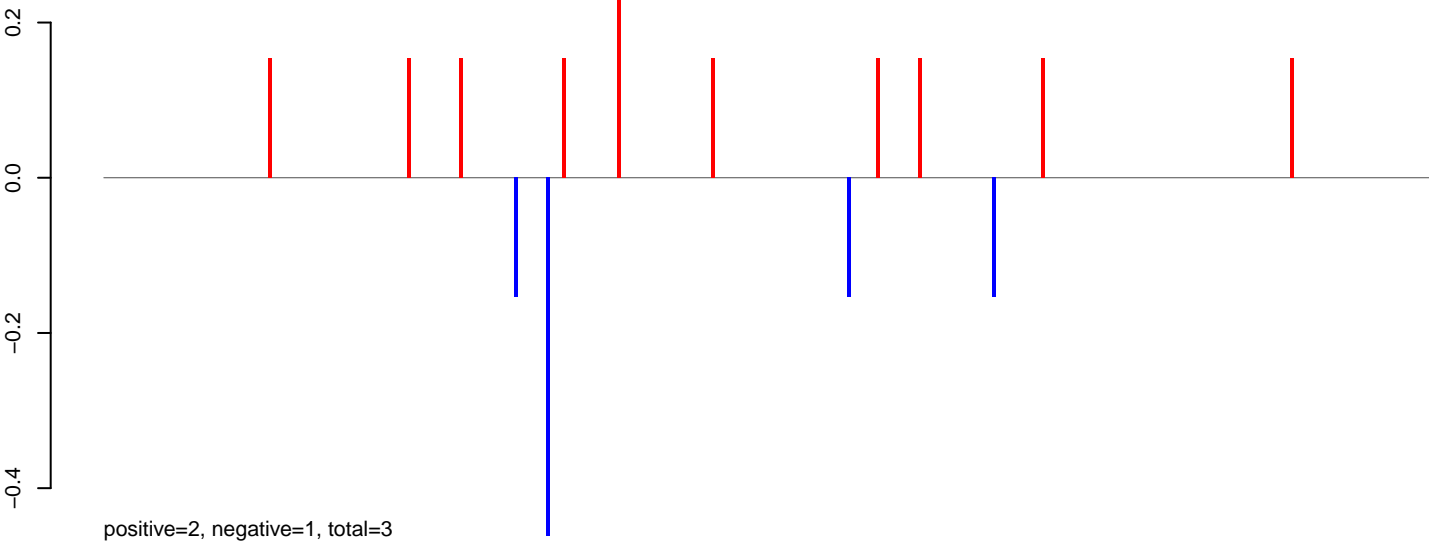
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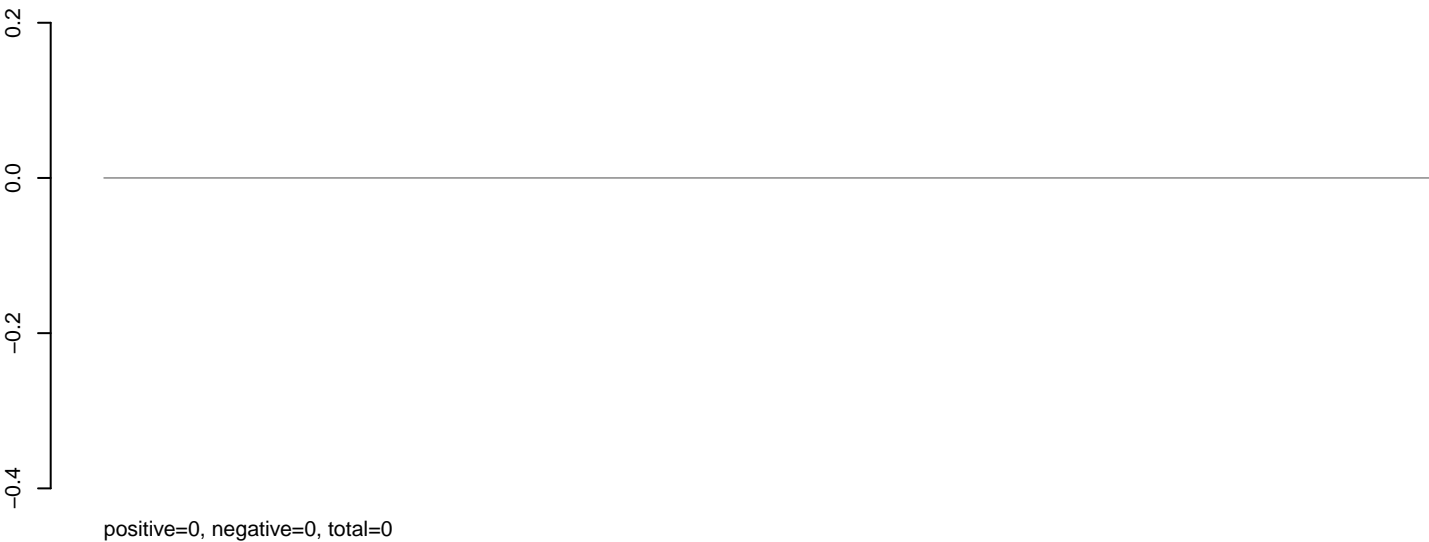
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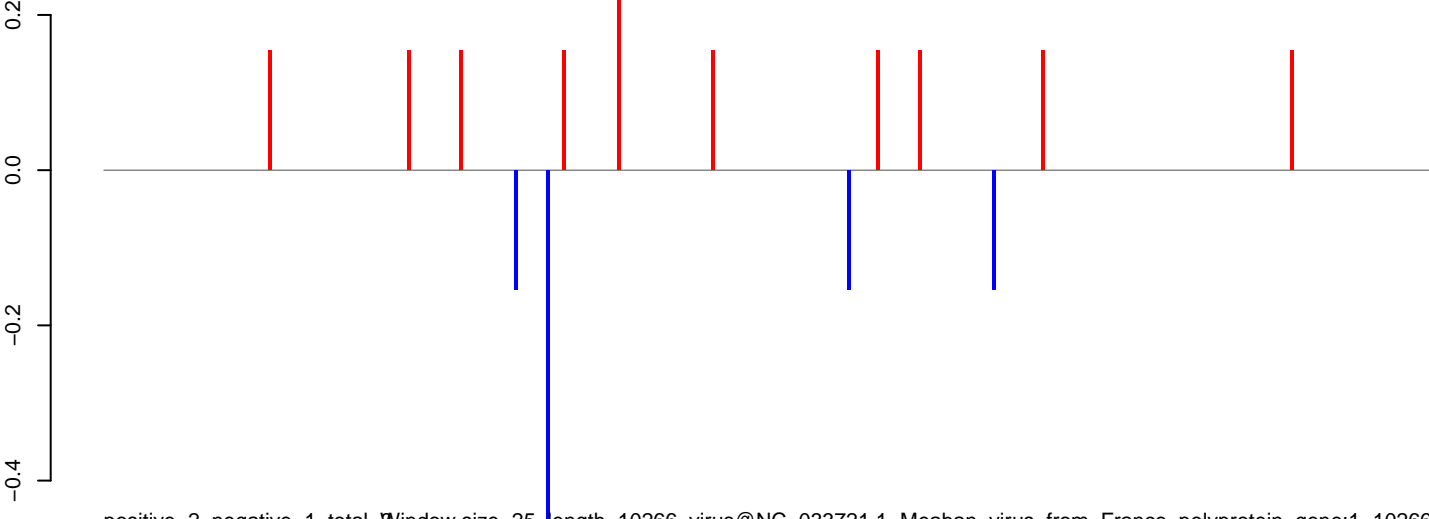
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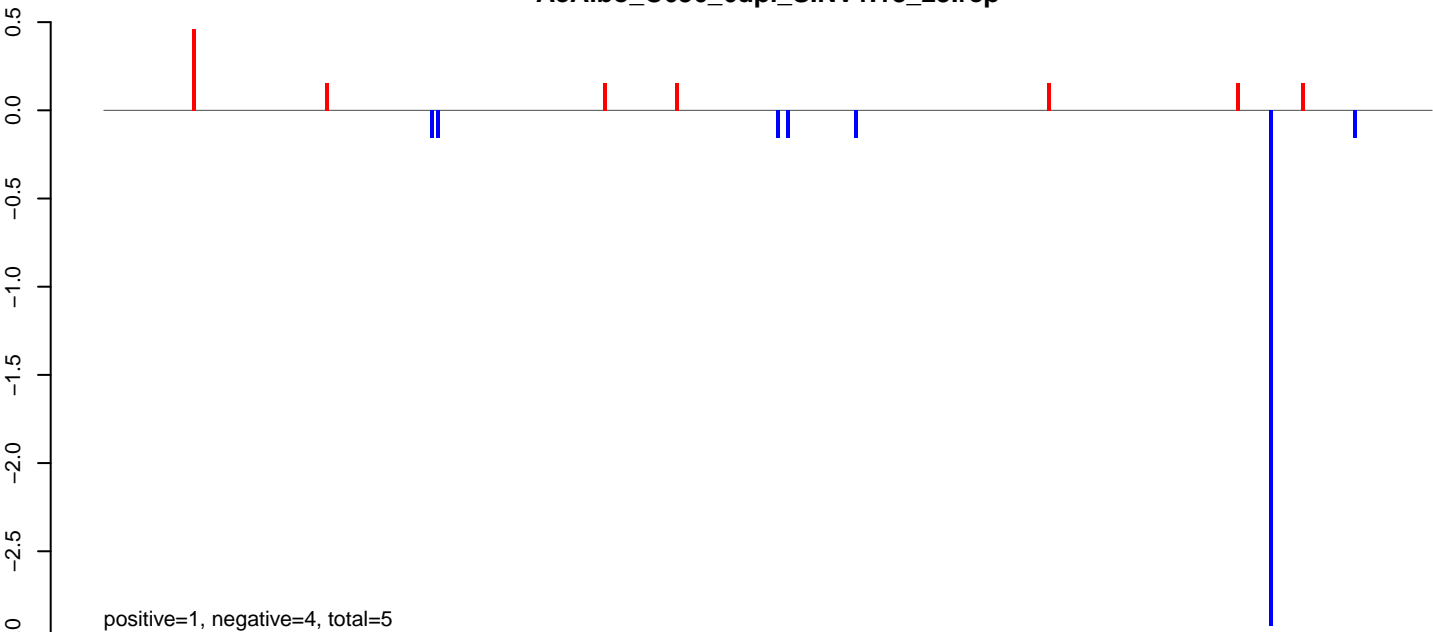
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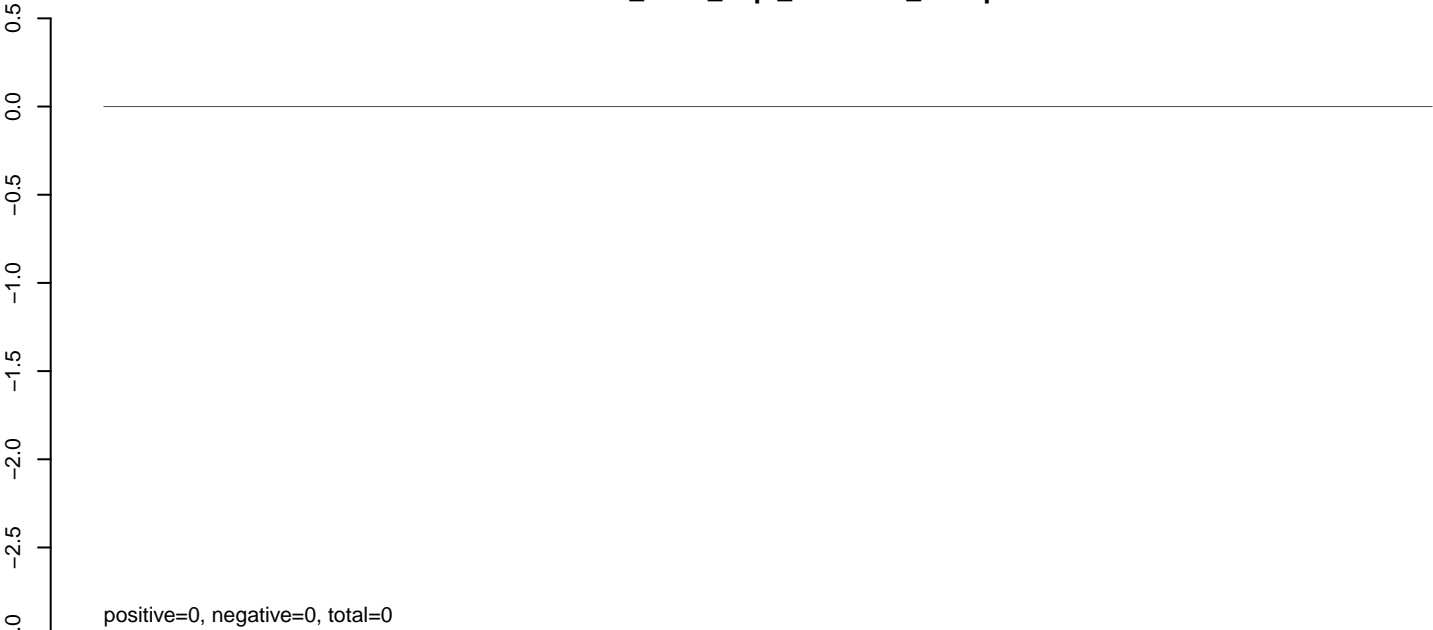
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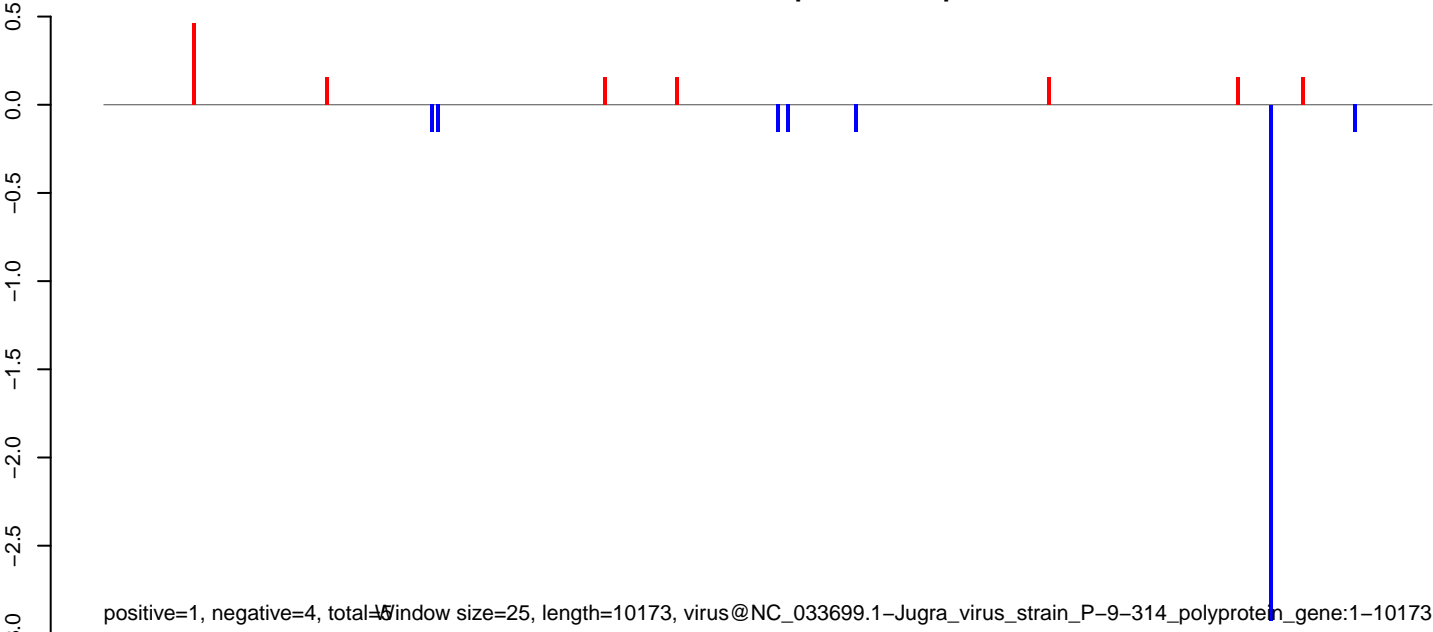
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AeAlbo_C636_6dpi_SINV1.24_35.rep



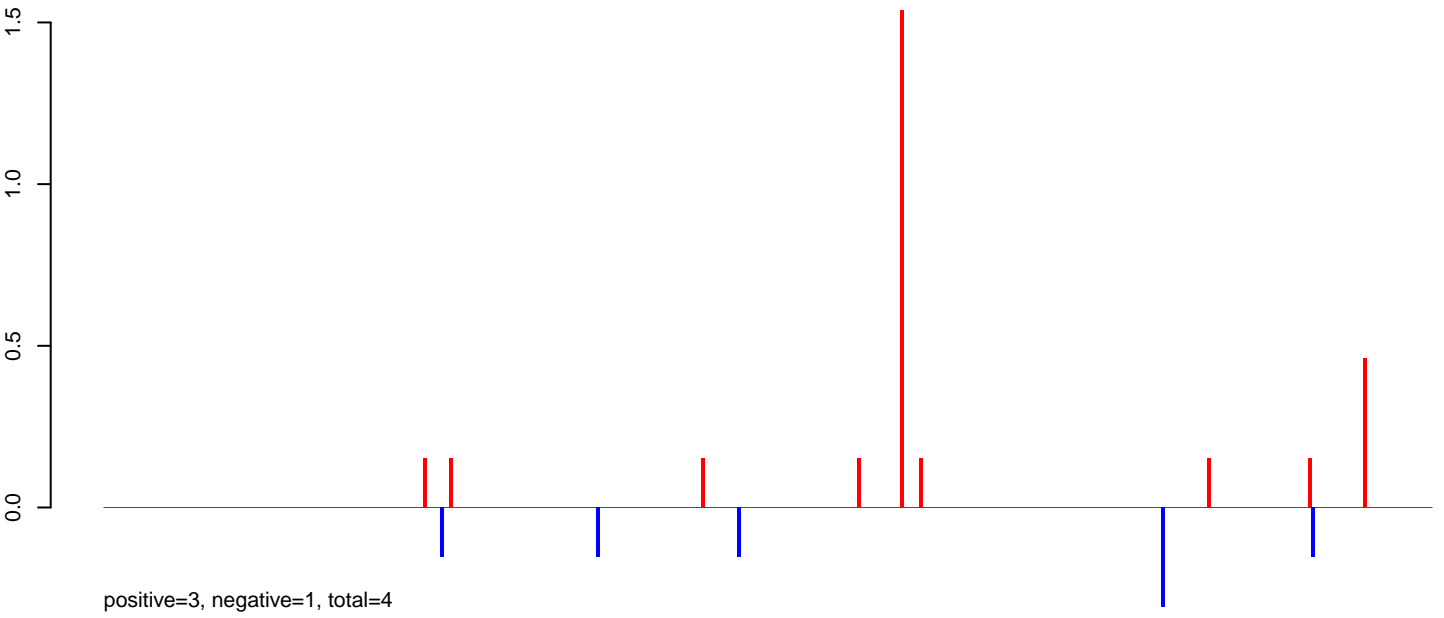
AeAlbo_C636_6dpi_SINV1.rep



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0 2000 4000 6000 8000 10000

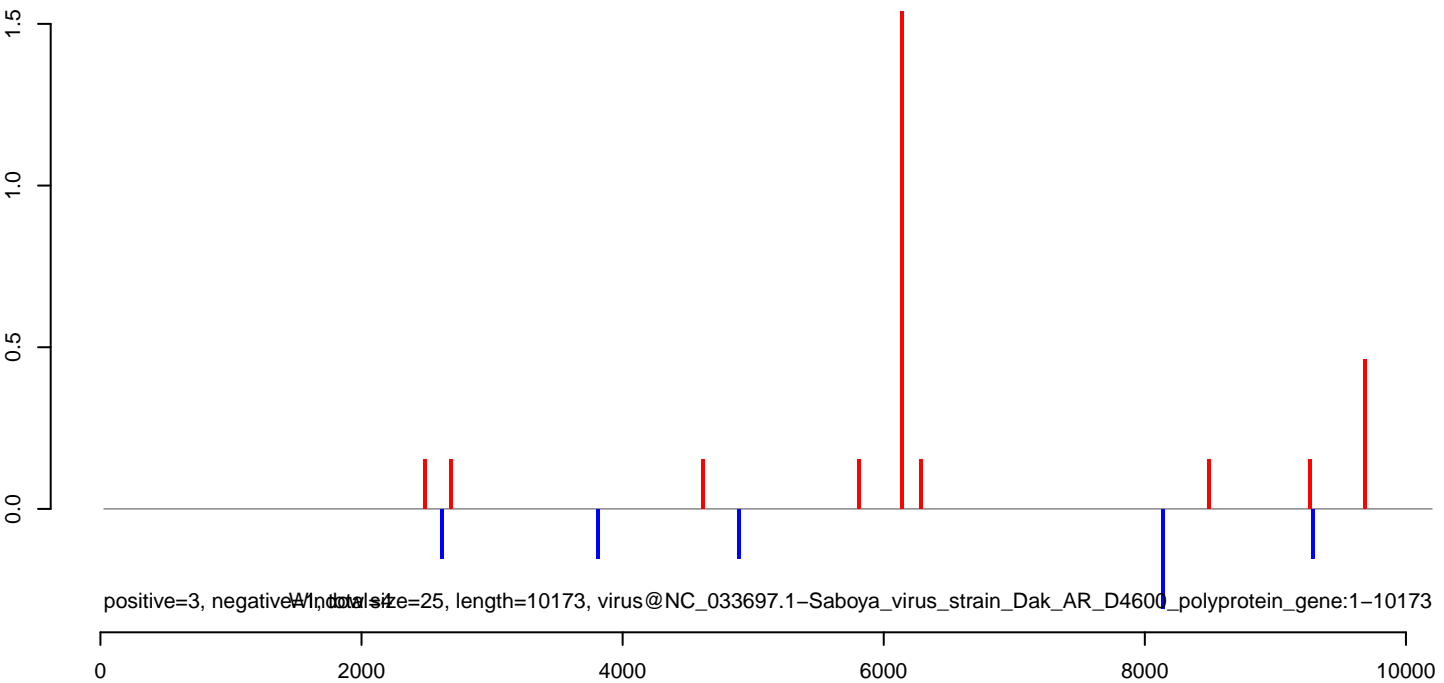
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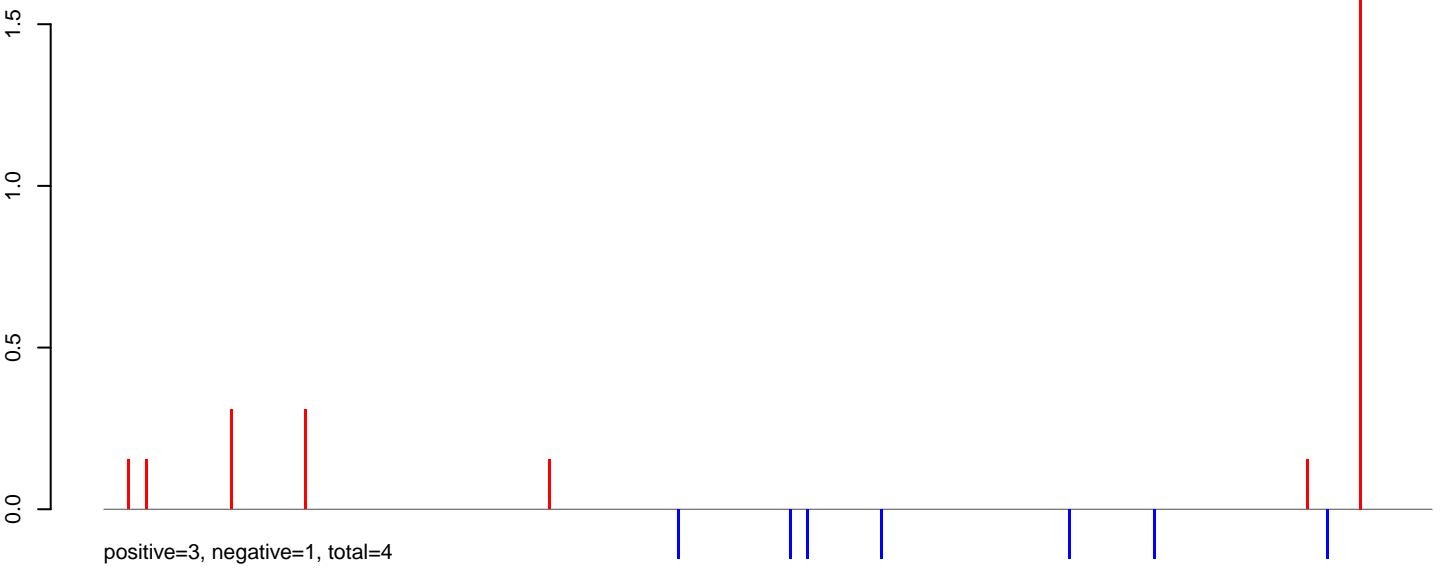
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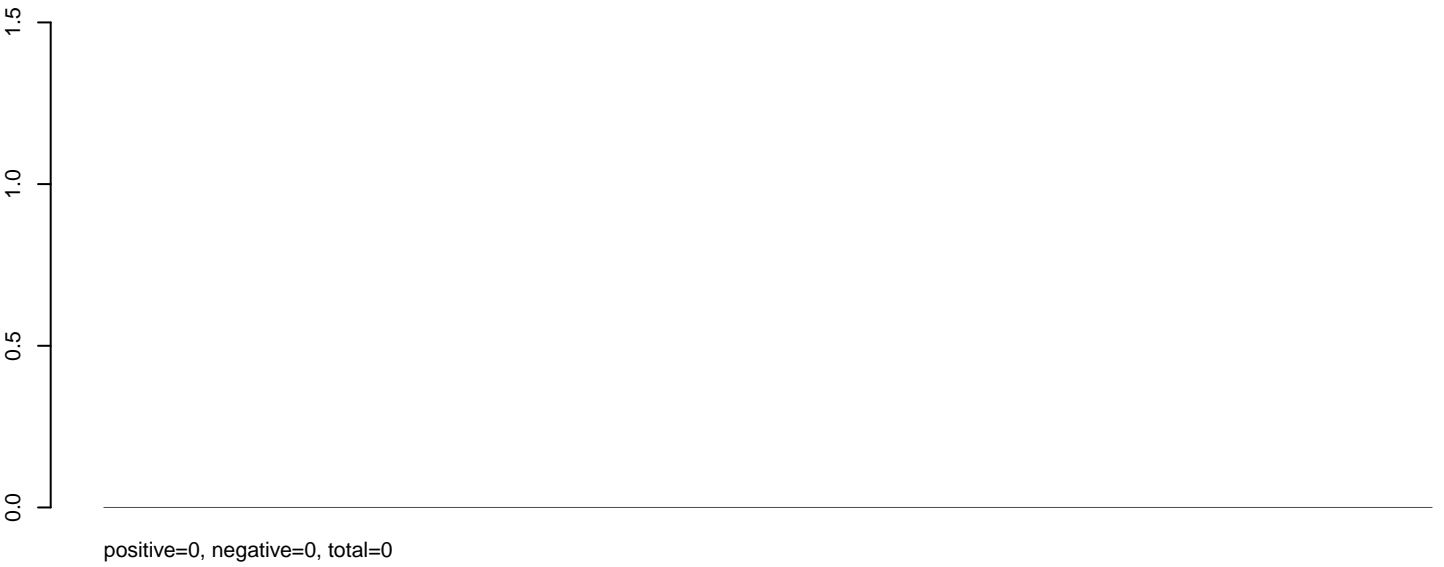
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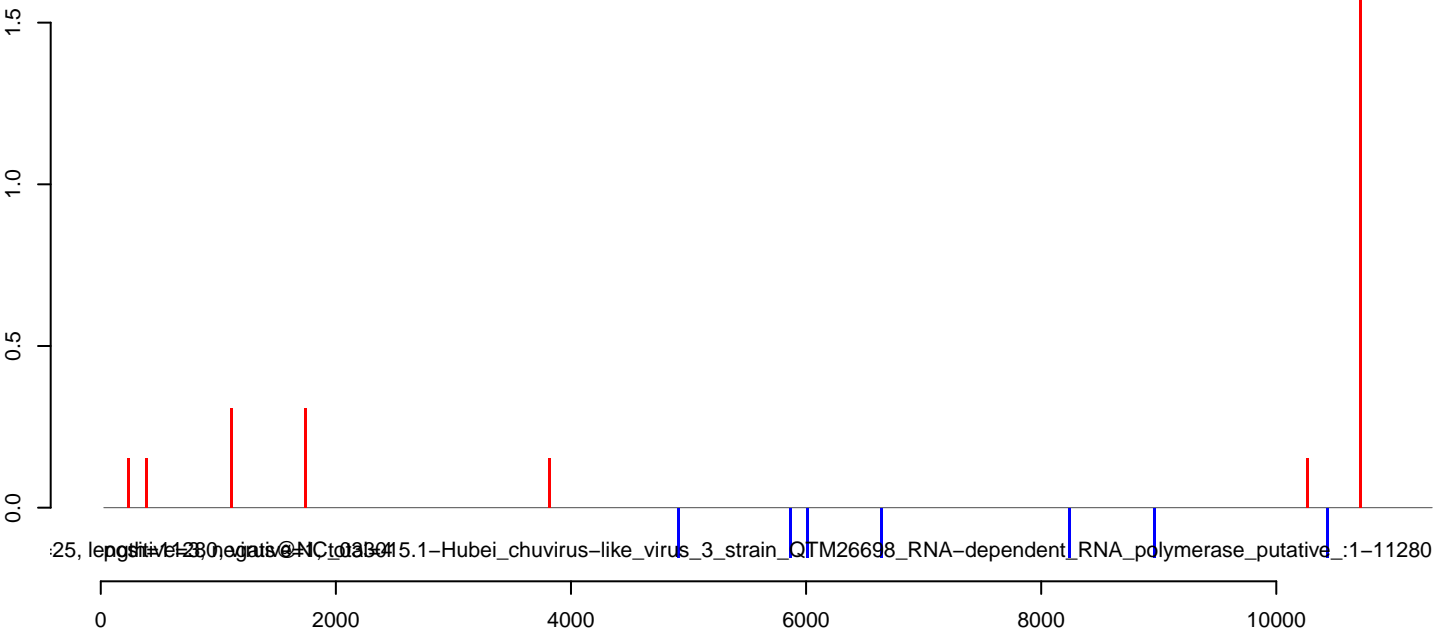
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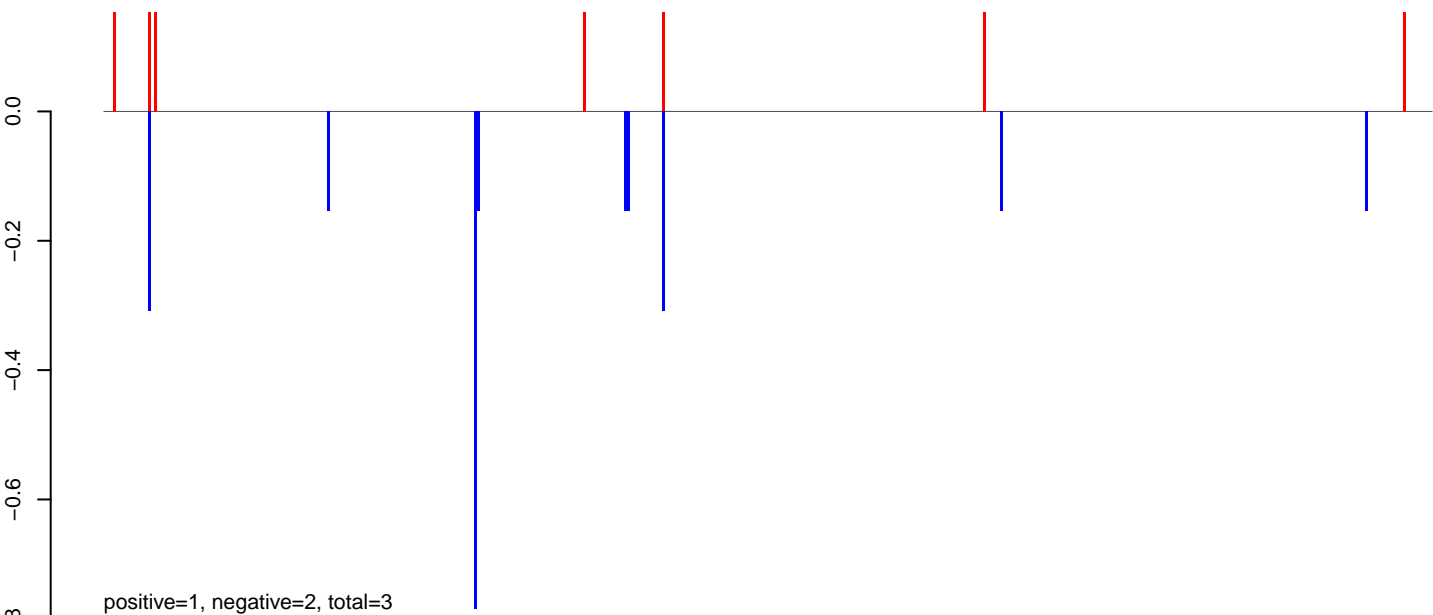
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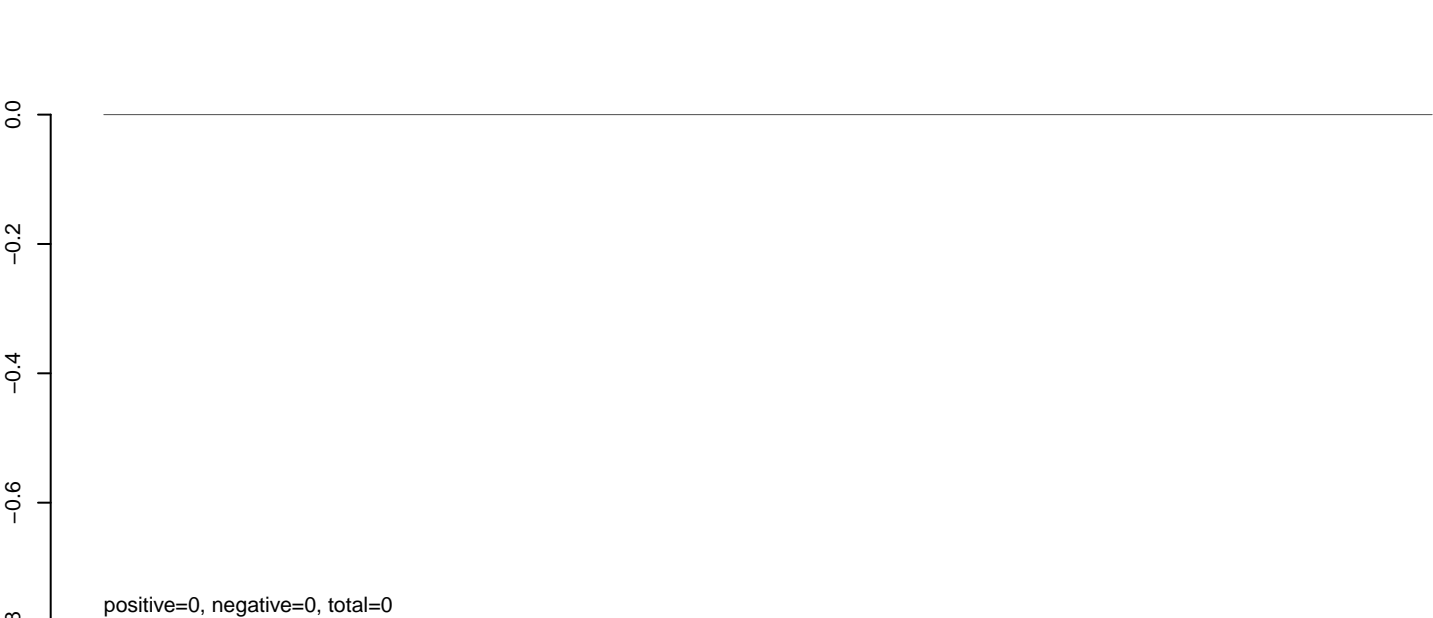
AeAlbo_C636_6dpi_SINV1.rep



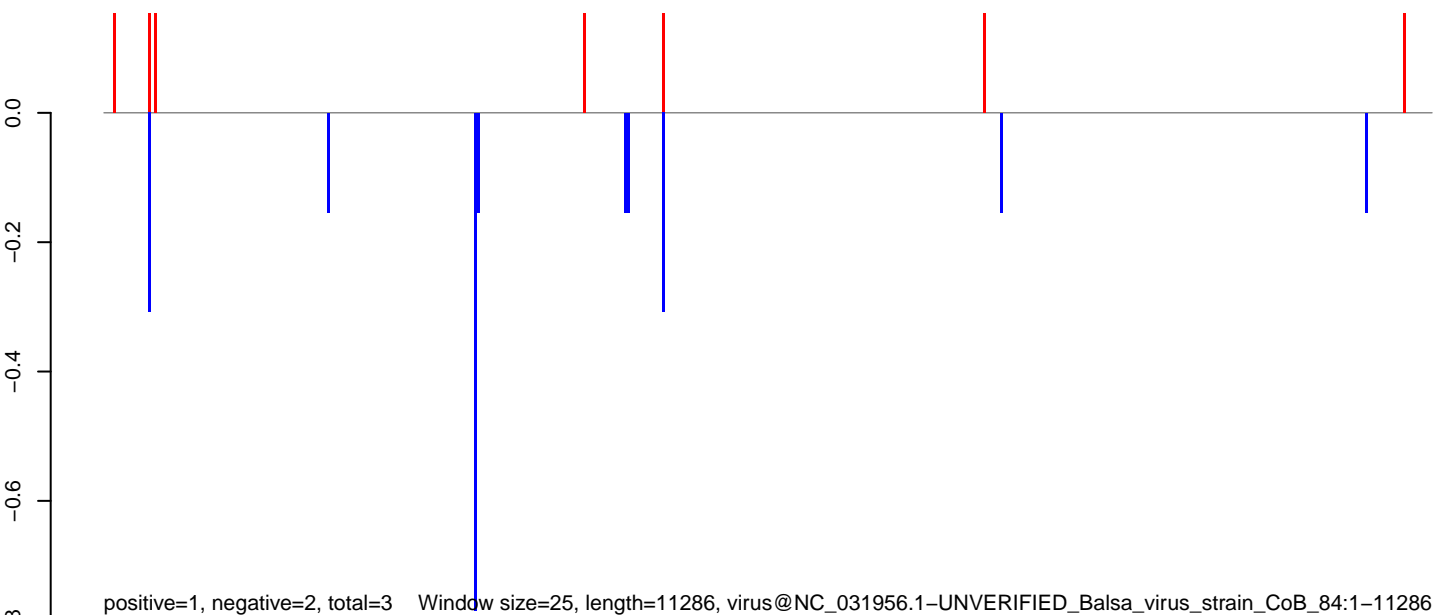
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep

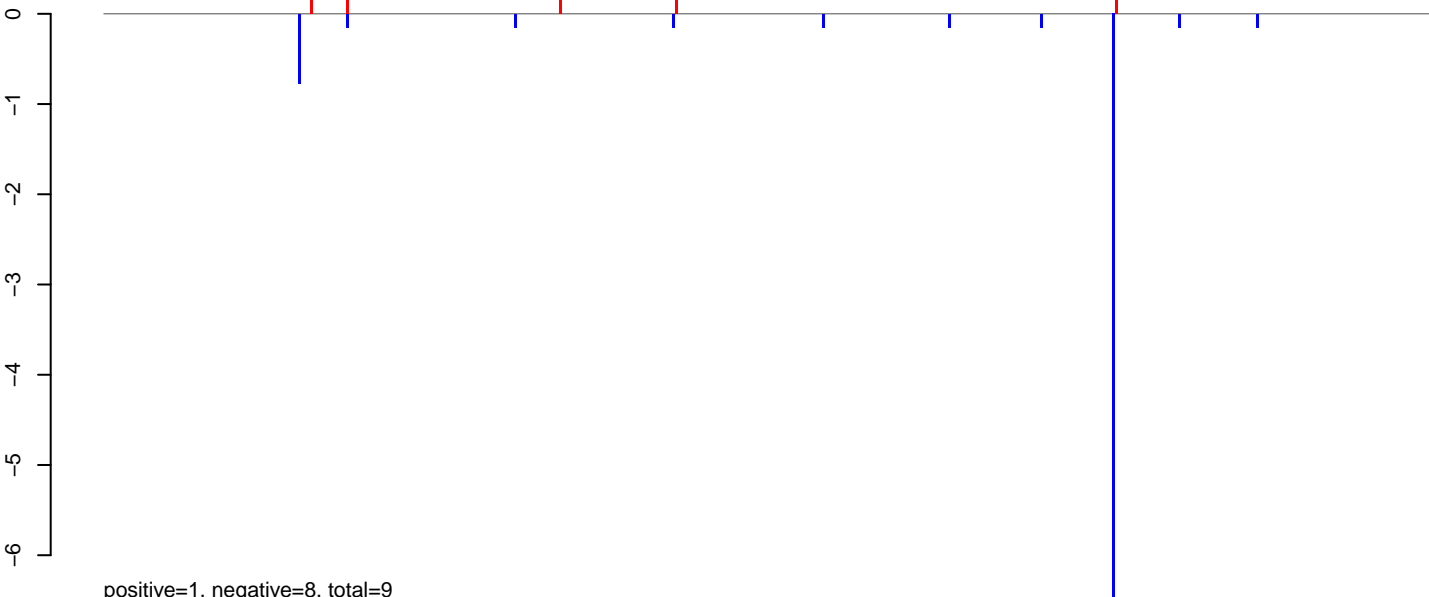


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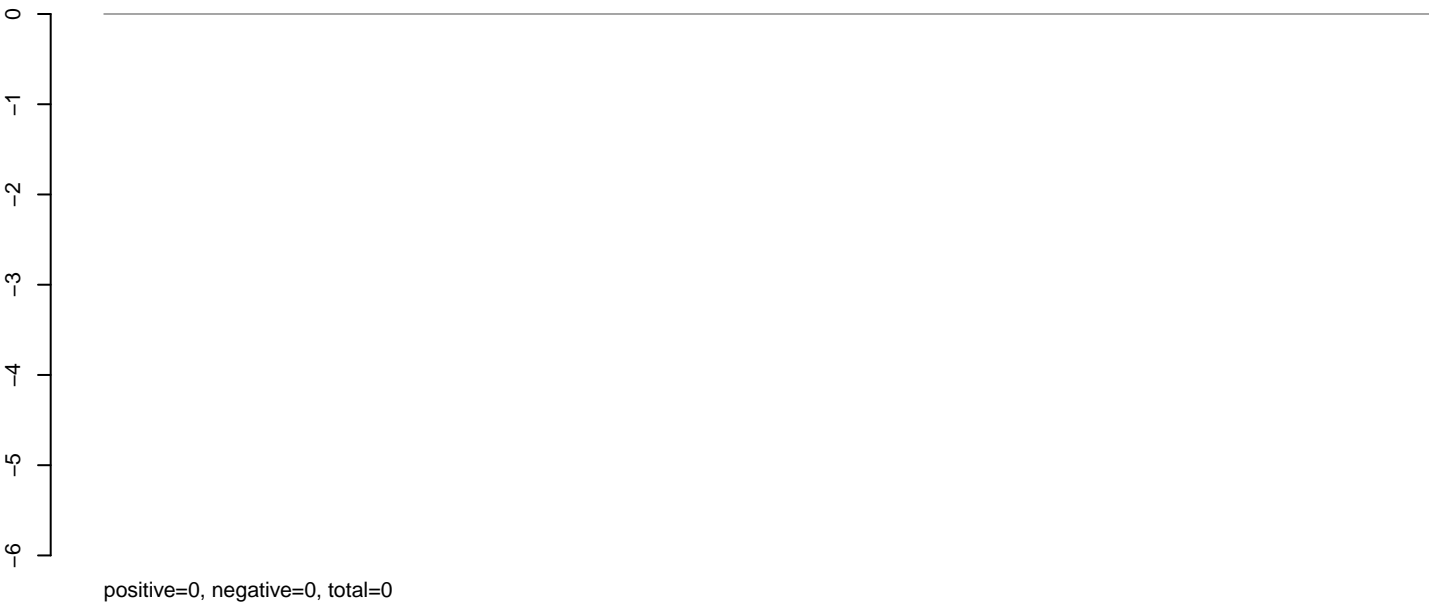


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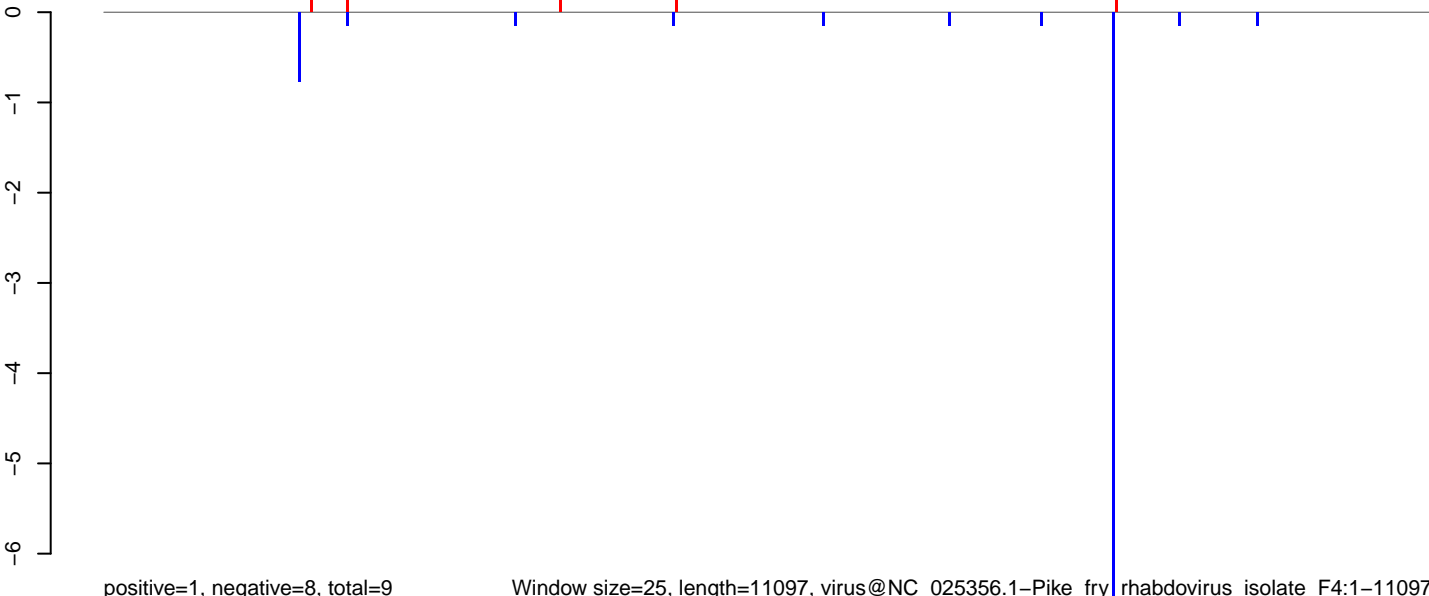
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



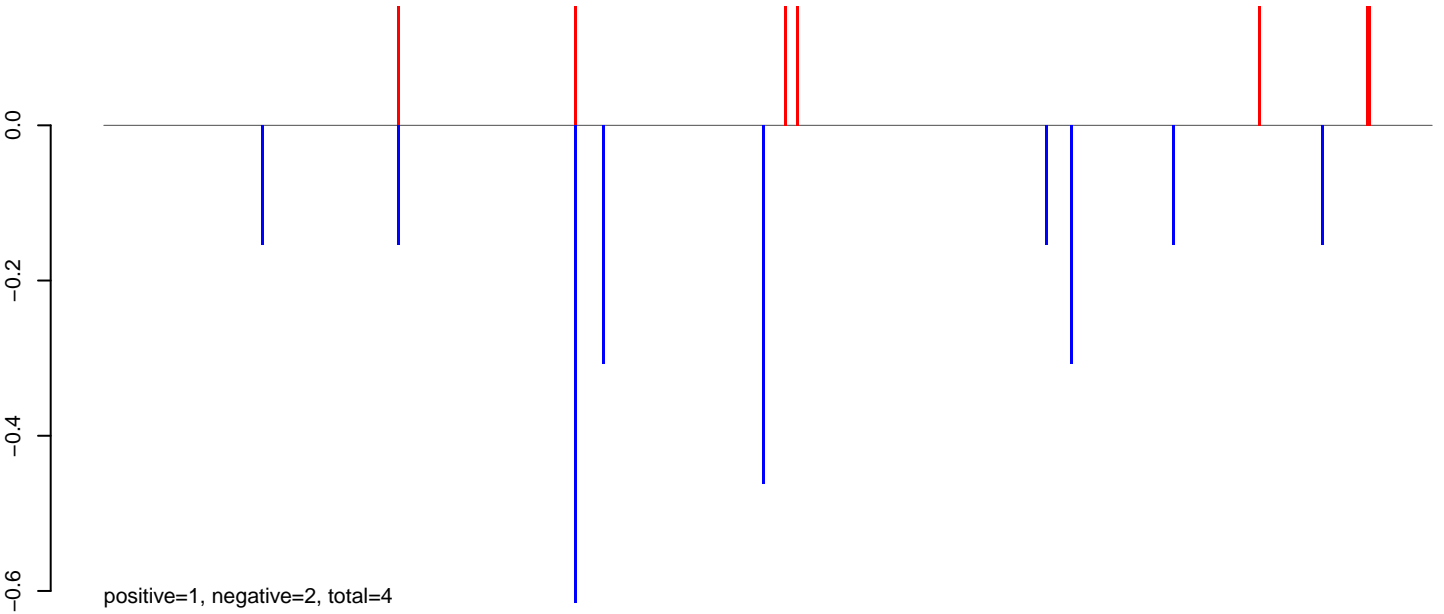
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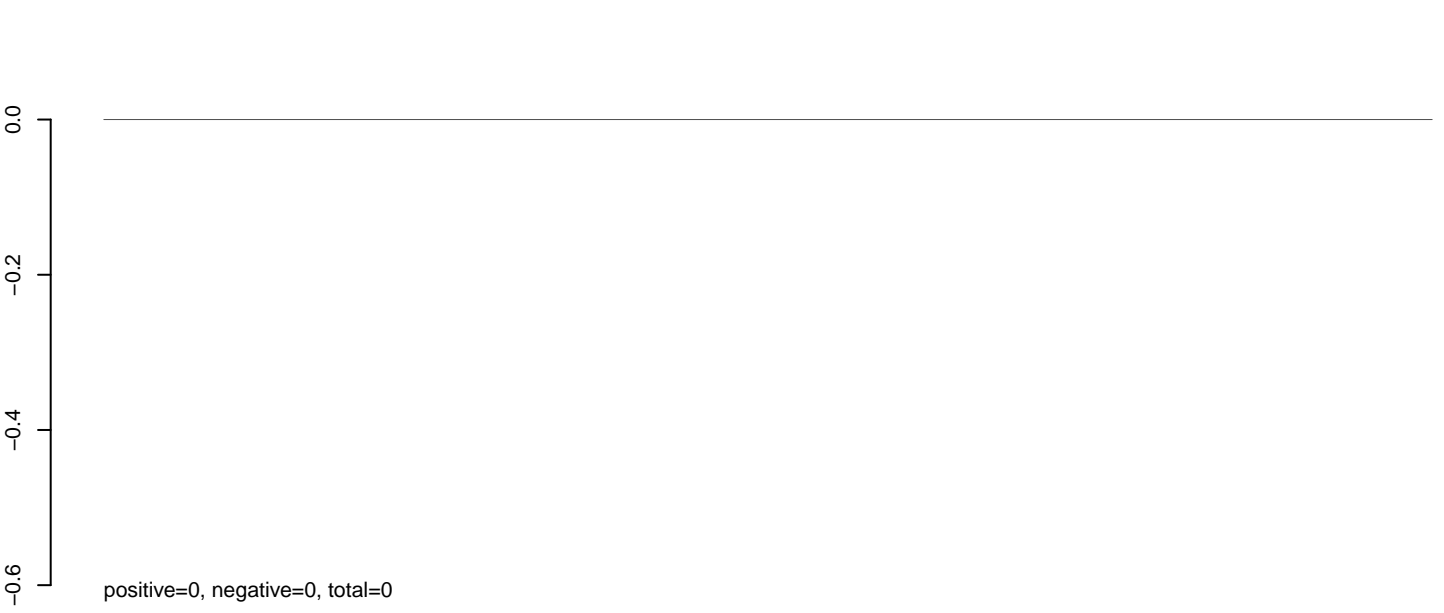
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0 2000 4000 6000 8000 10000

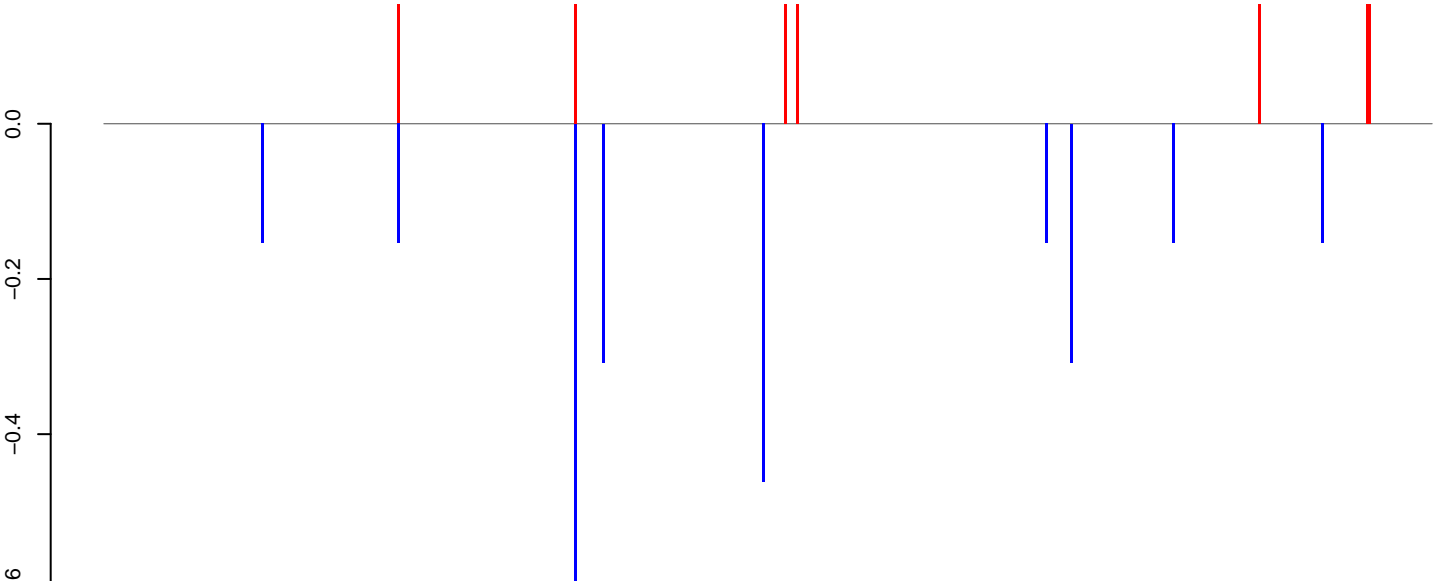
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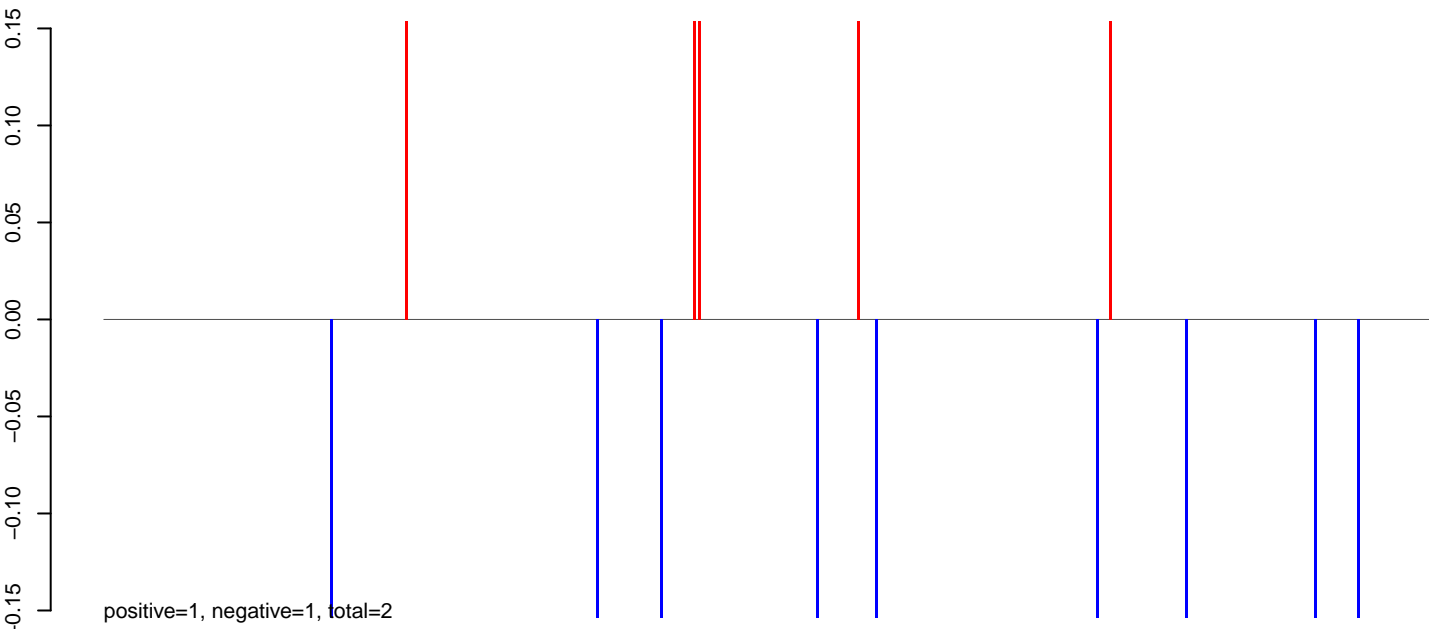
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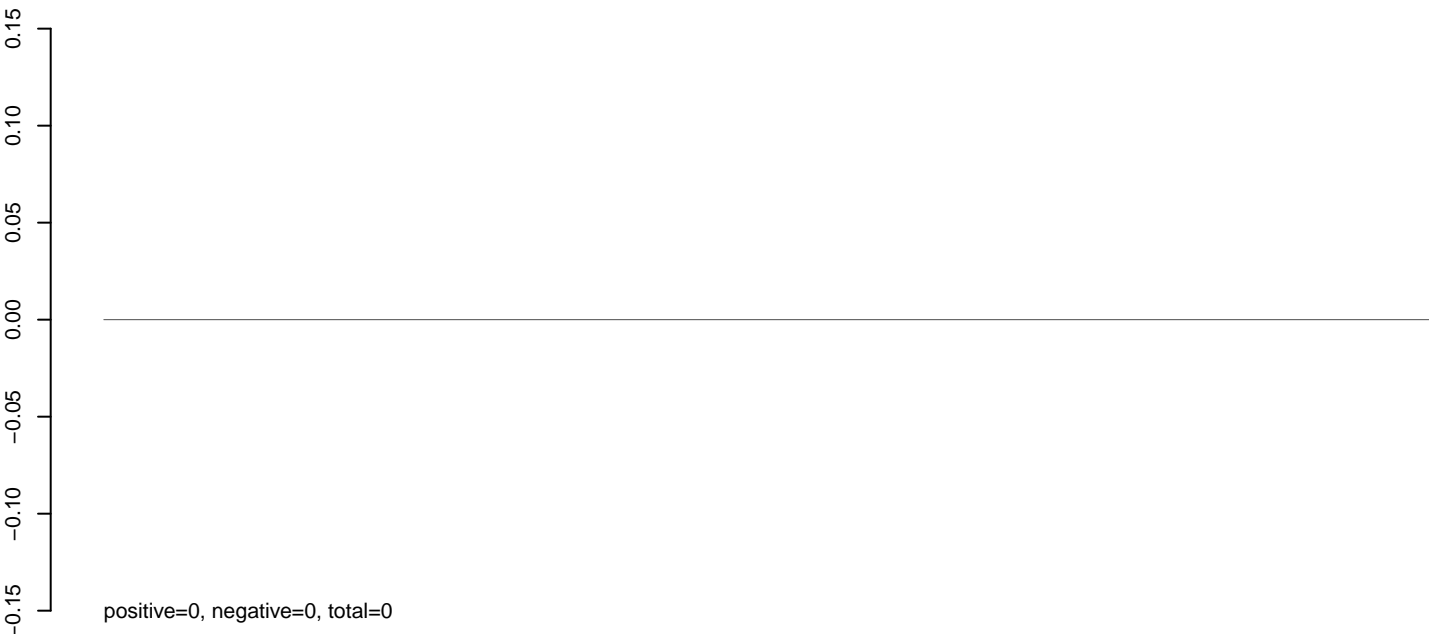
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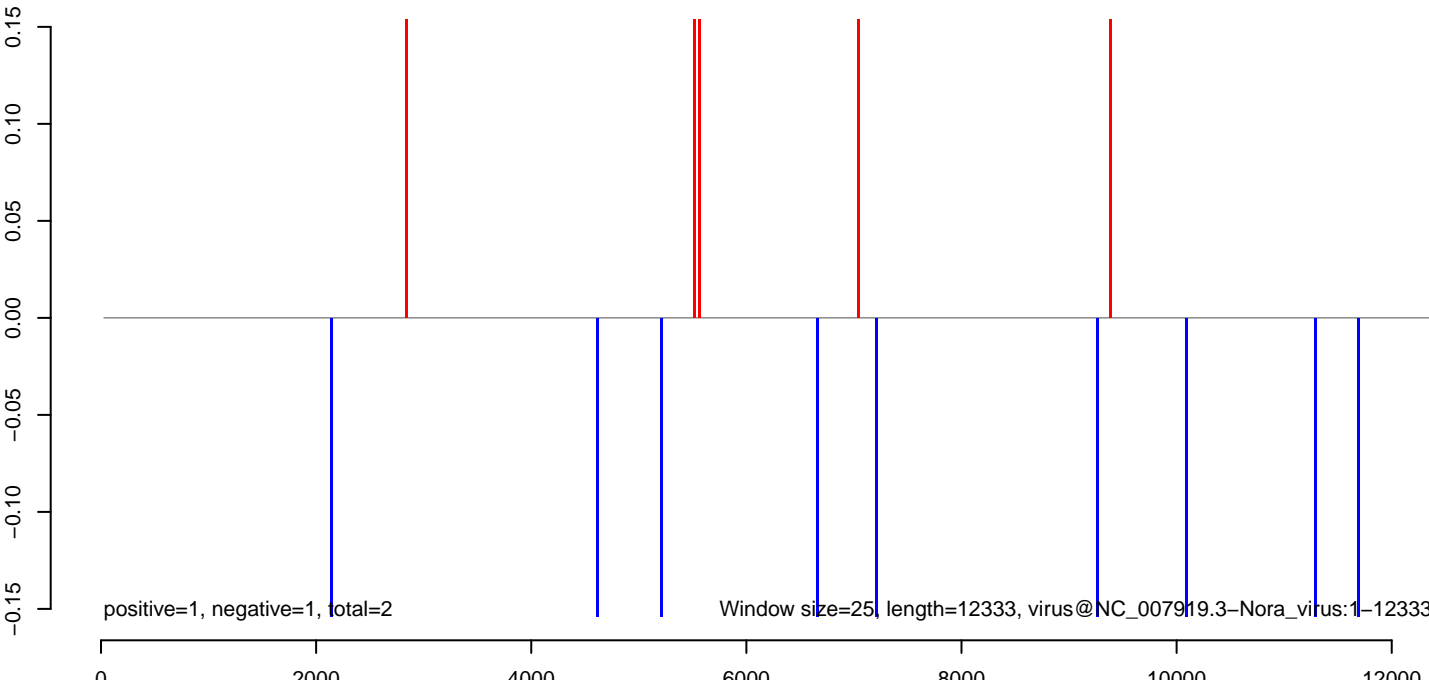
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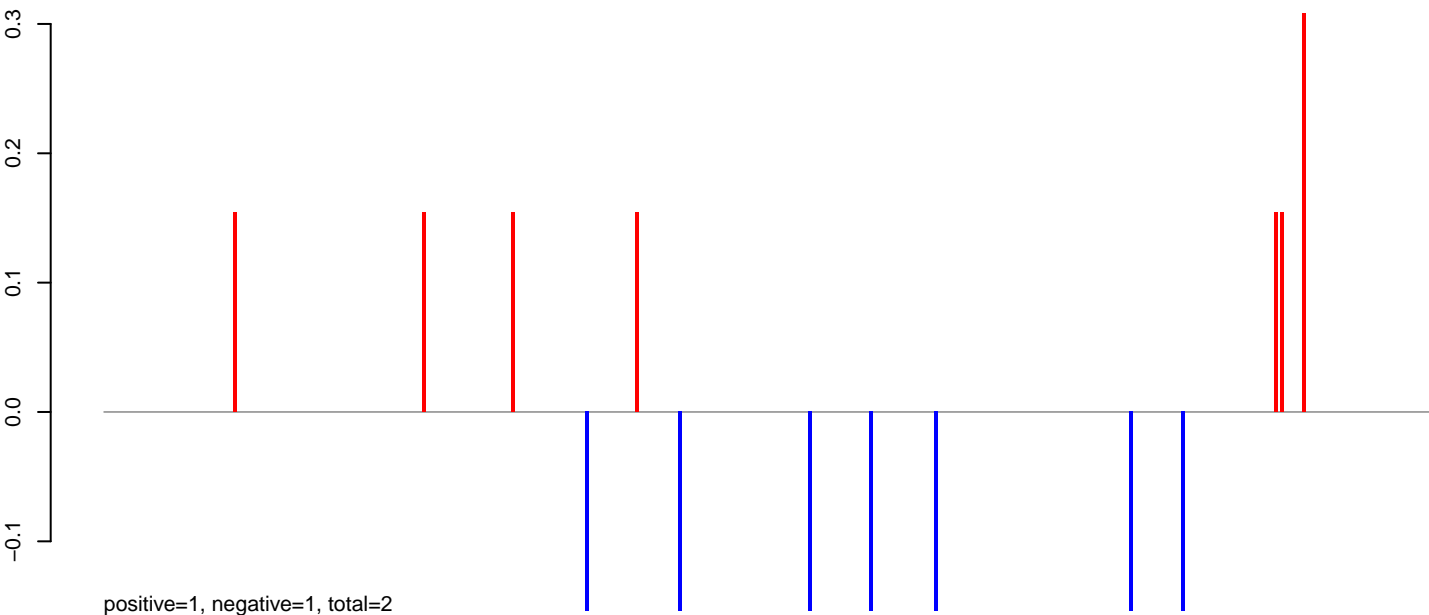
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AeAlbo_C636_6dpi_SINV1.rep



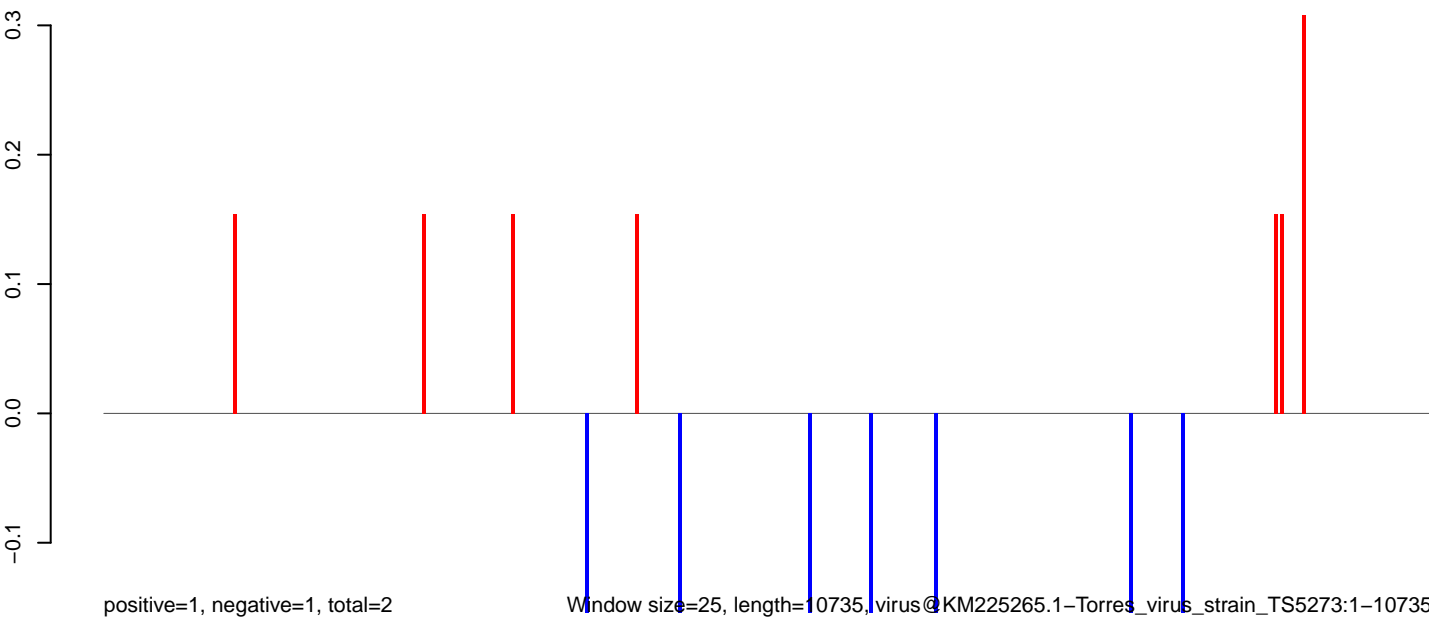
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



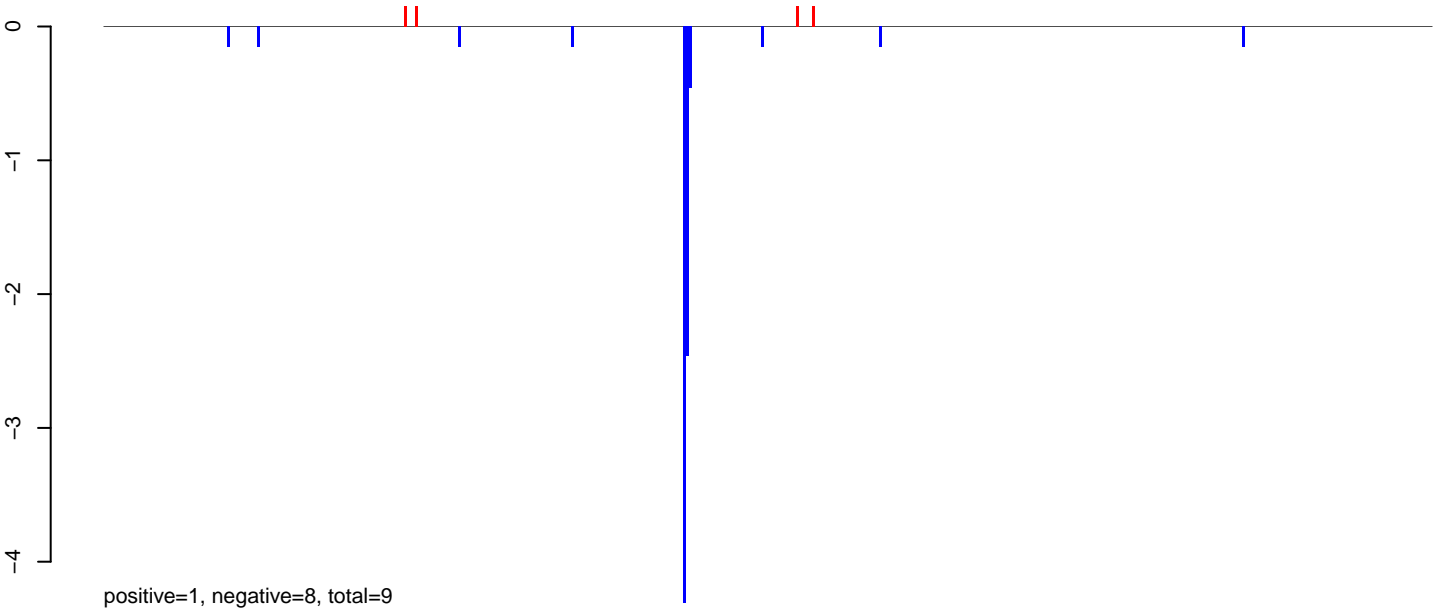
AeAlbo_C636_6dpi_SINV1.rep



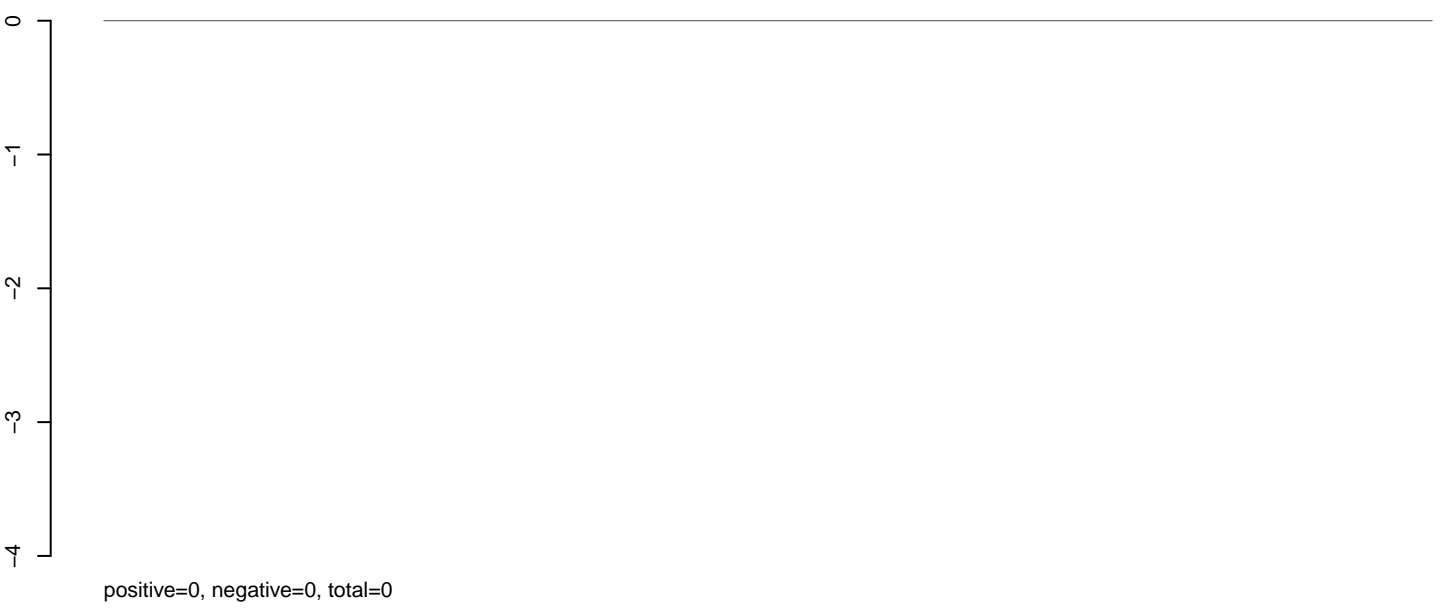
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0 2000 4000 6000 8000 10000

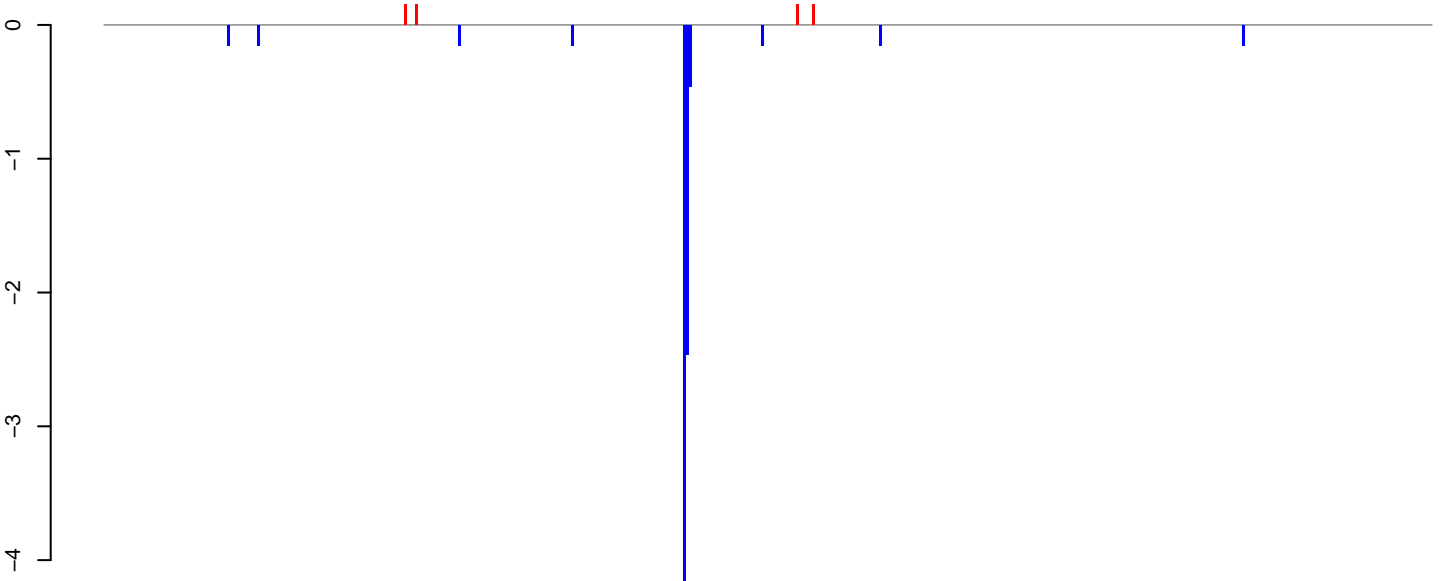
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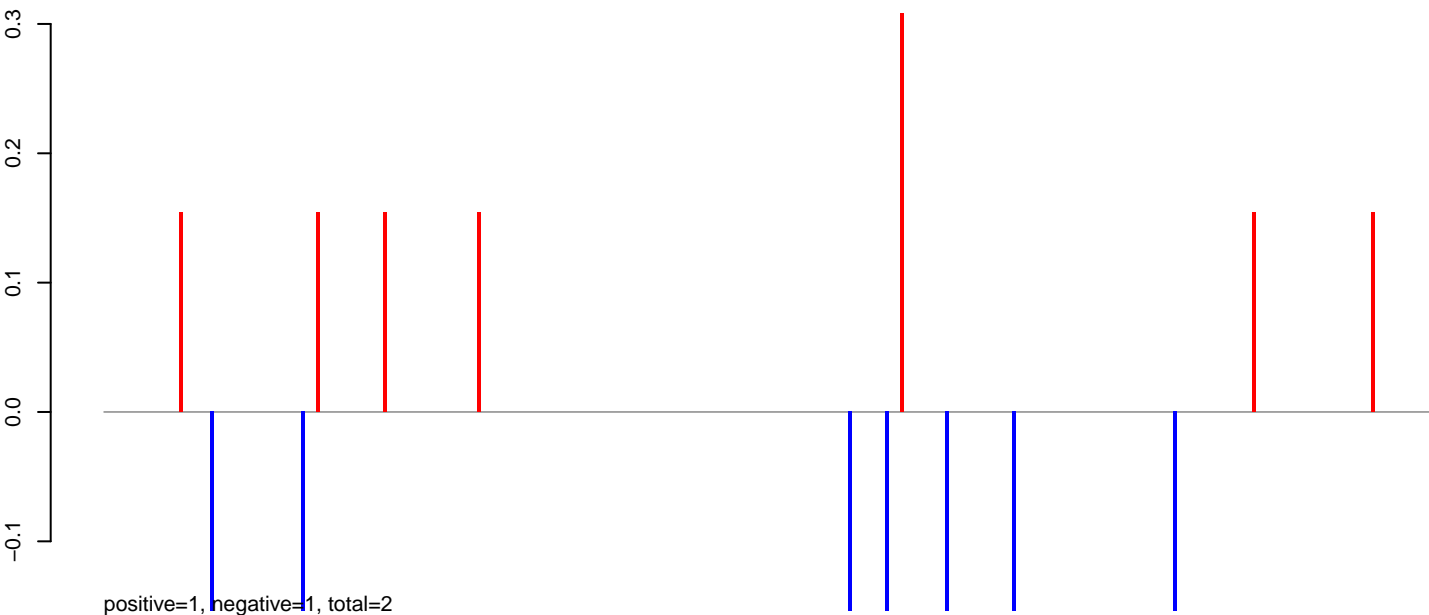
AeAlbo_C636_6dpi_SINV1.24_35.rep



AeAlbo_C636_6dpi_SINV1.rep



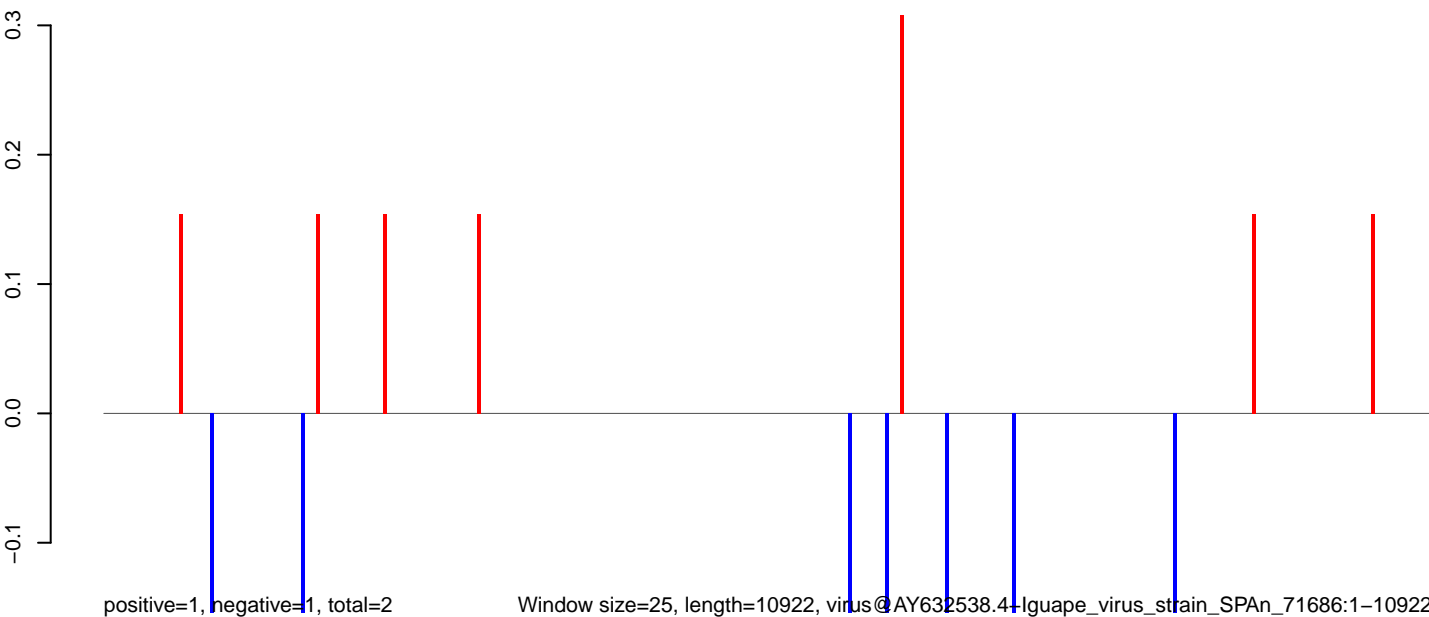
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



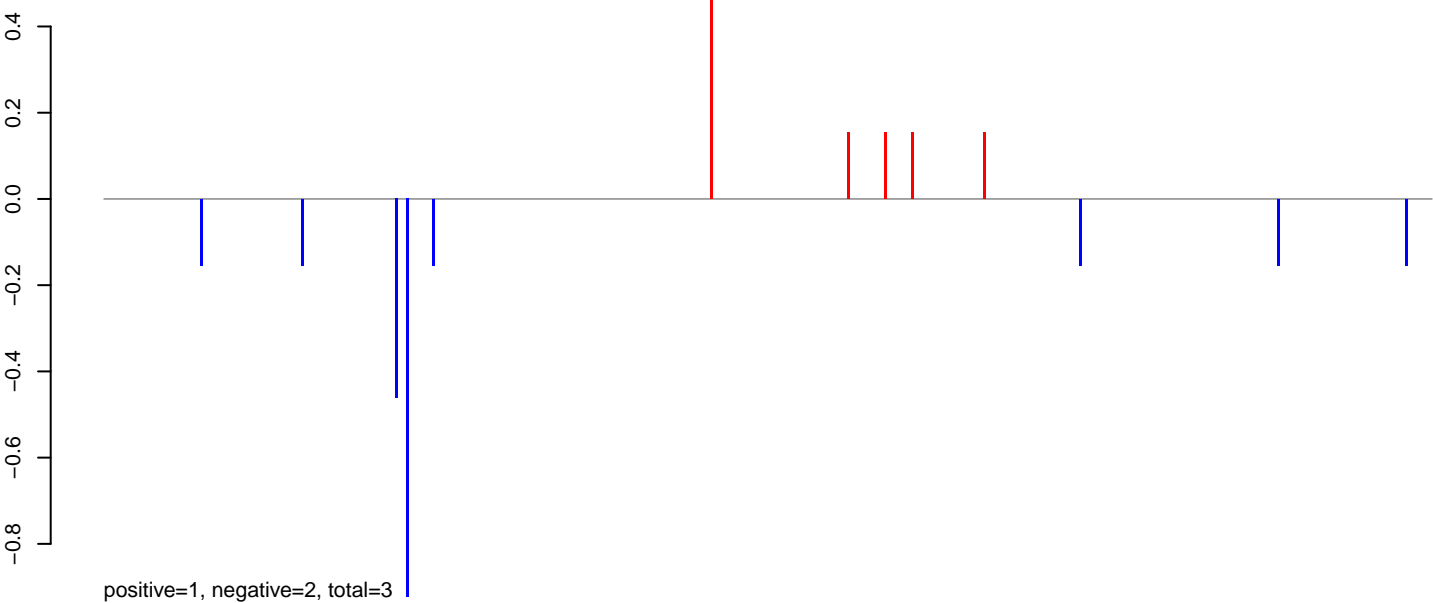
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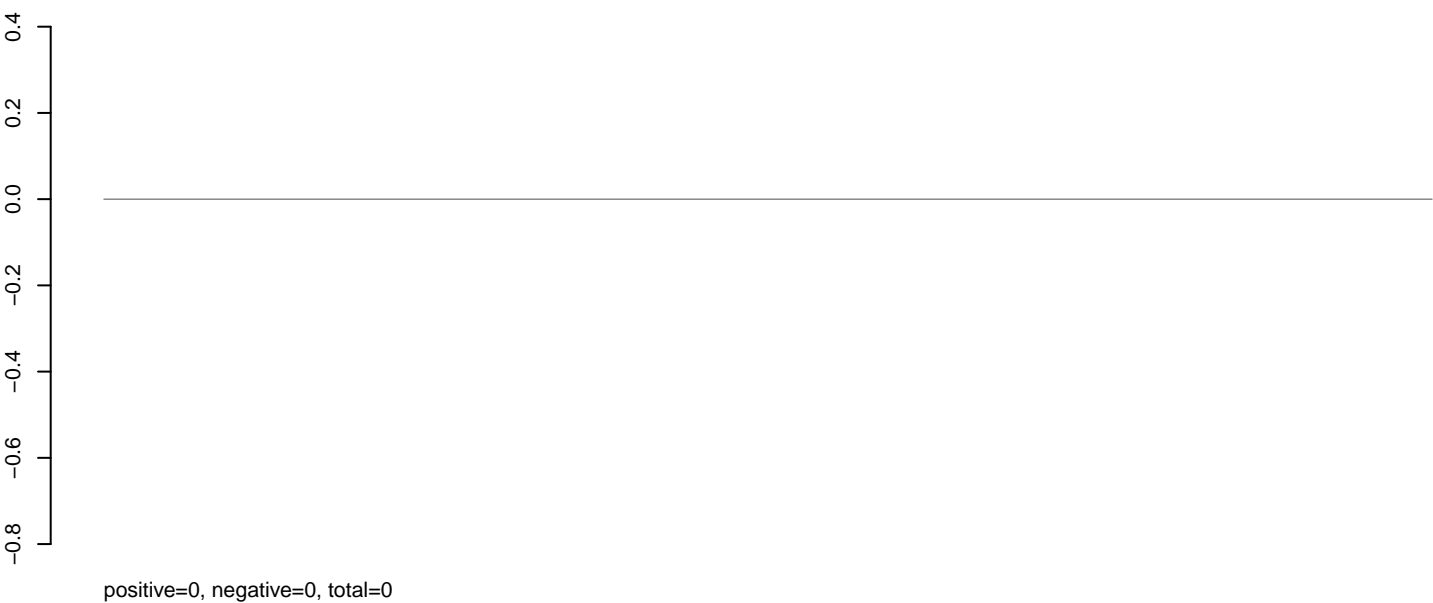
Window size=25, length=10922, virus@AY632538.4-Iguape_virus_strain_SPAn_71686:1-10922

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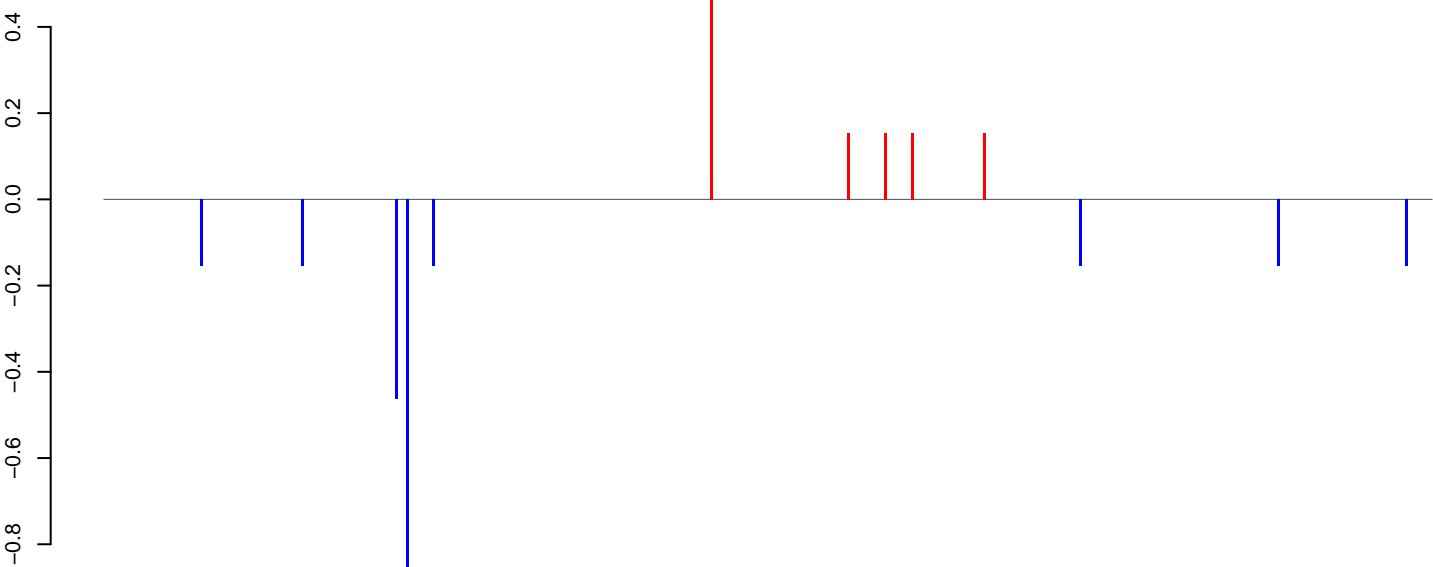
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



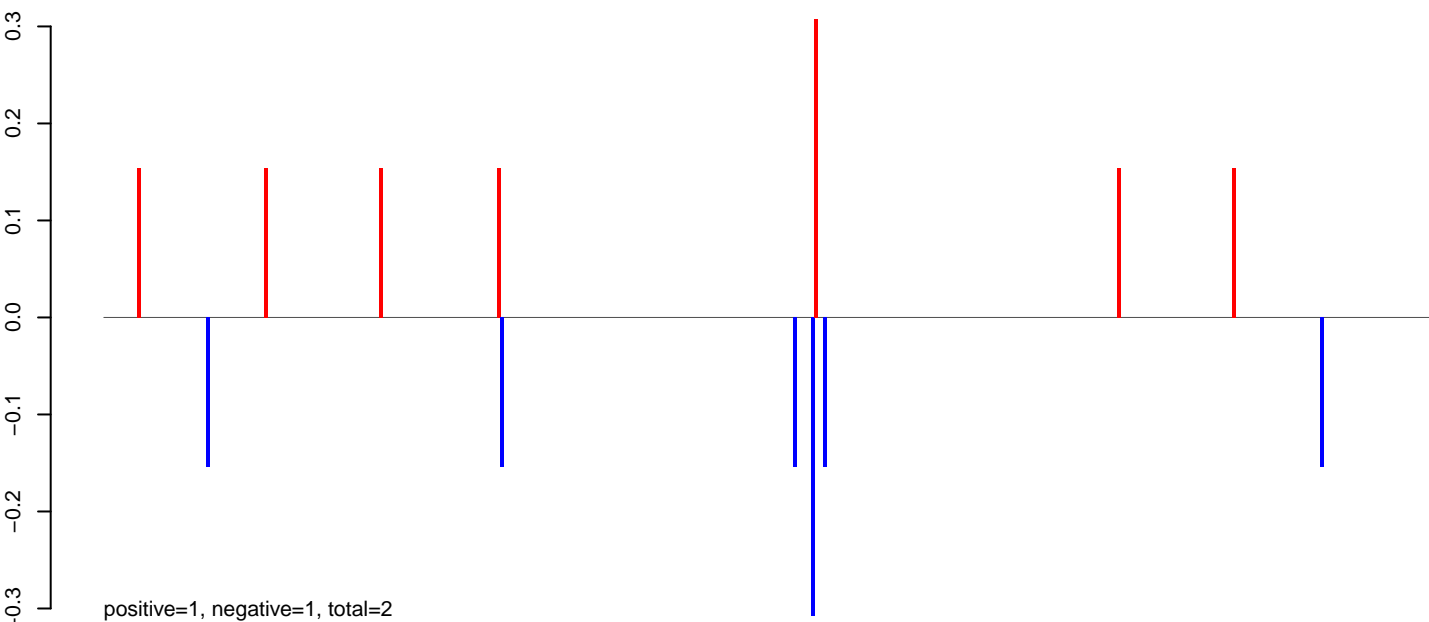
AeAlbo_C636_6dpi_SINV1.rep



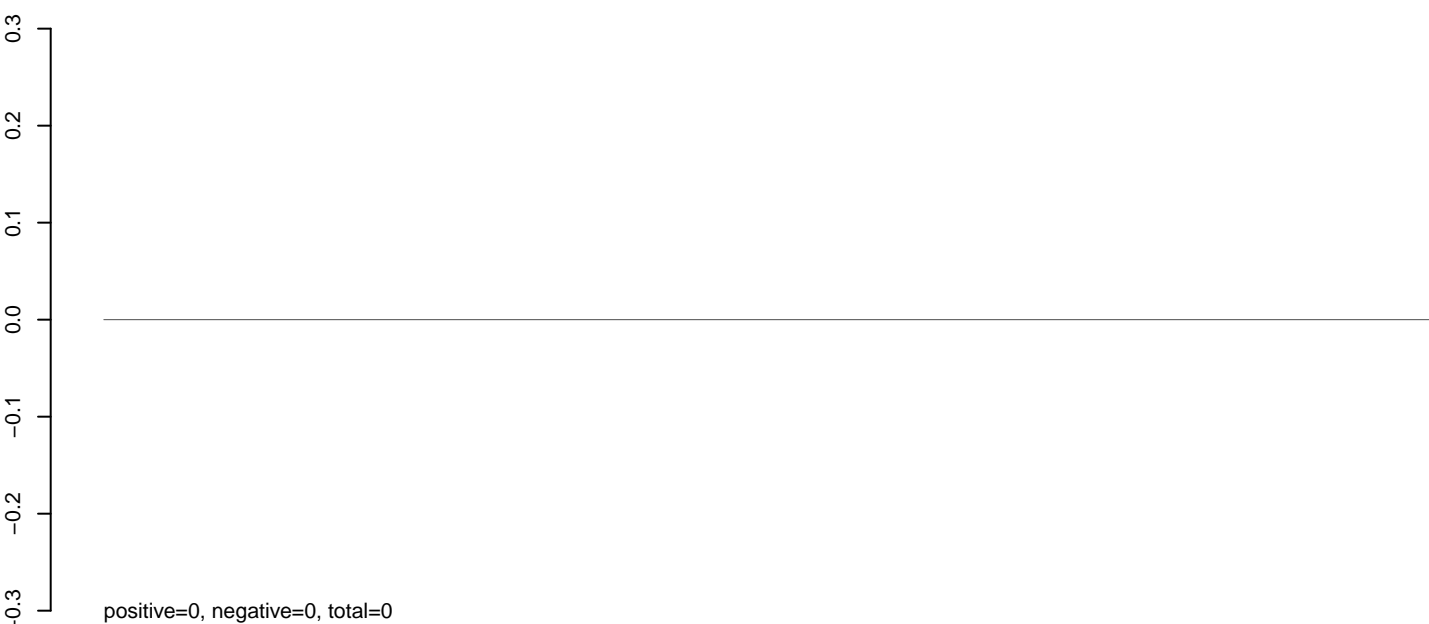
v sizes=25, length=12400, use=0, NC_034541.1-Gray_Lodge_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopr:1-12403

0 2000 4000 6000 8000 10000 12000

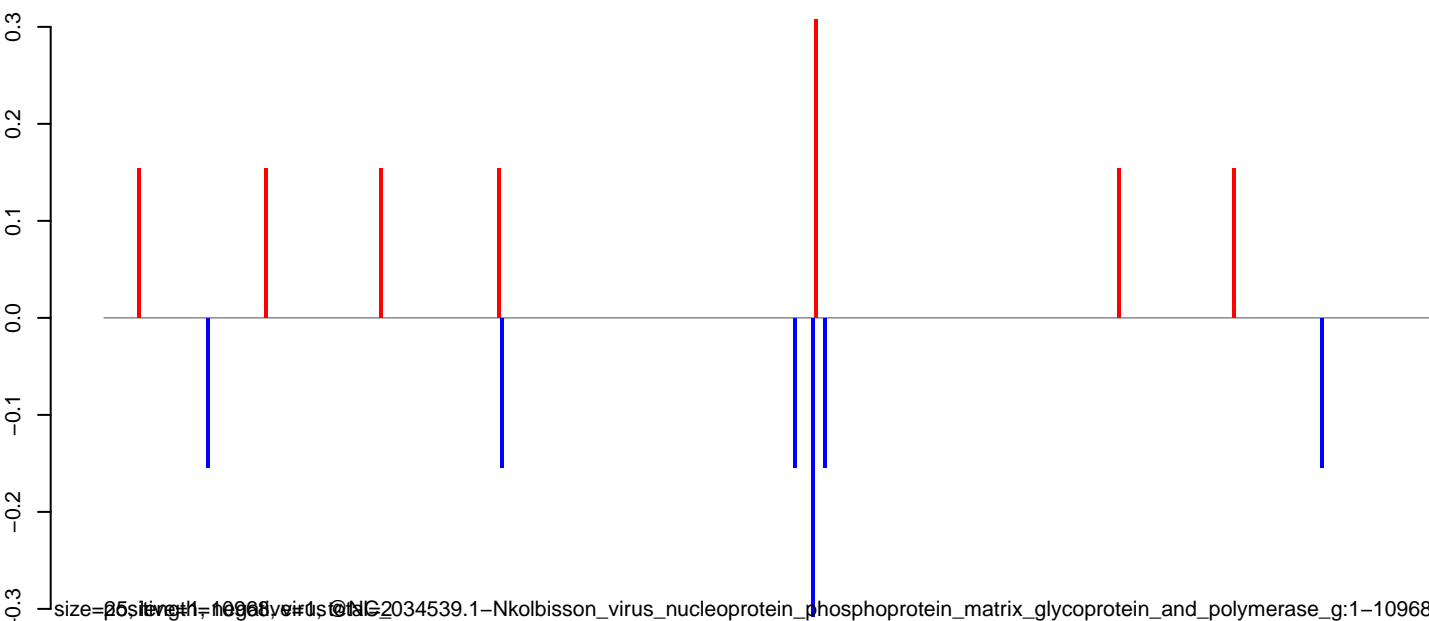
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep

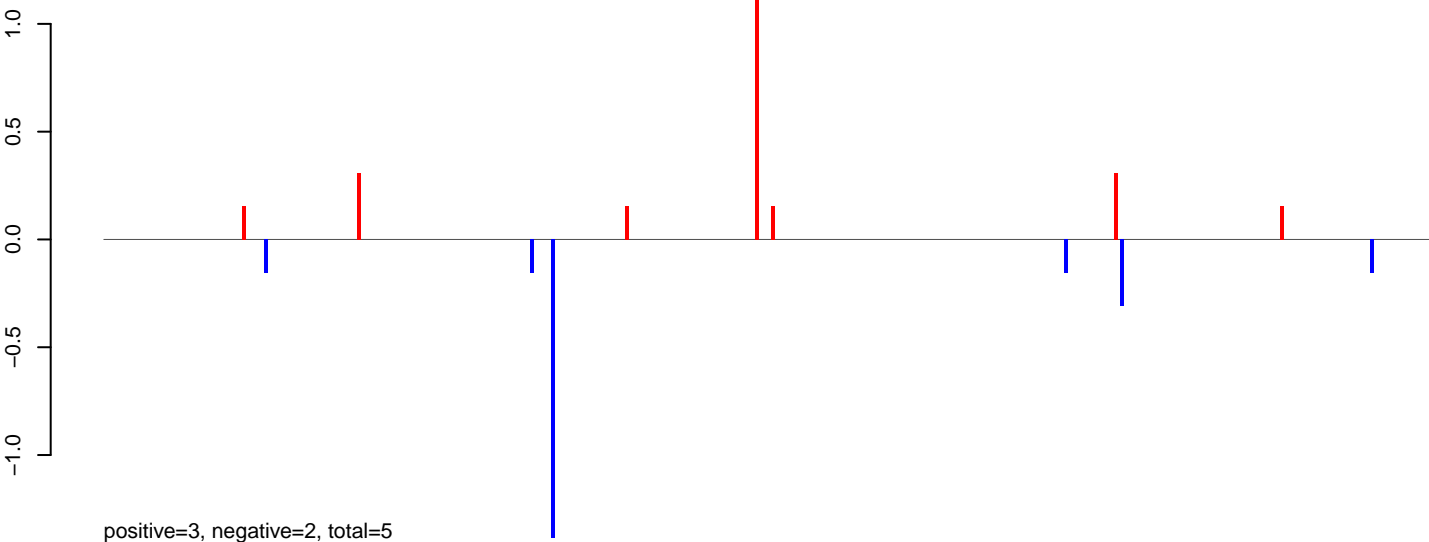


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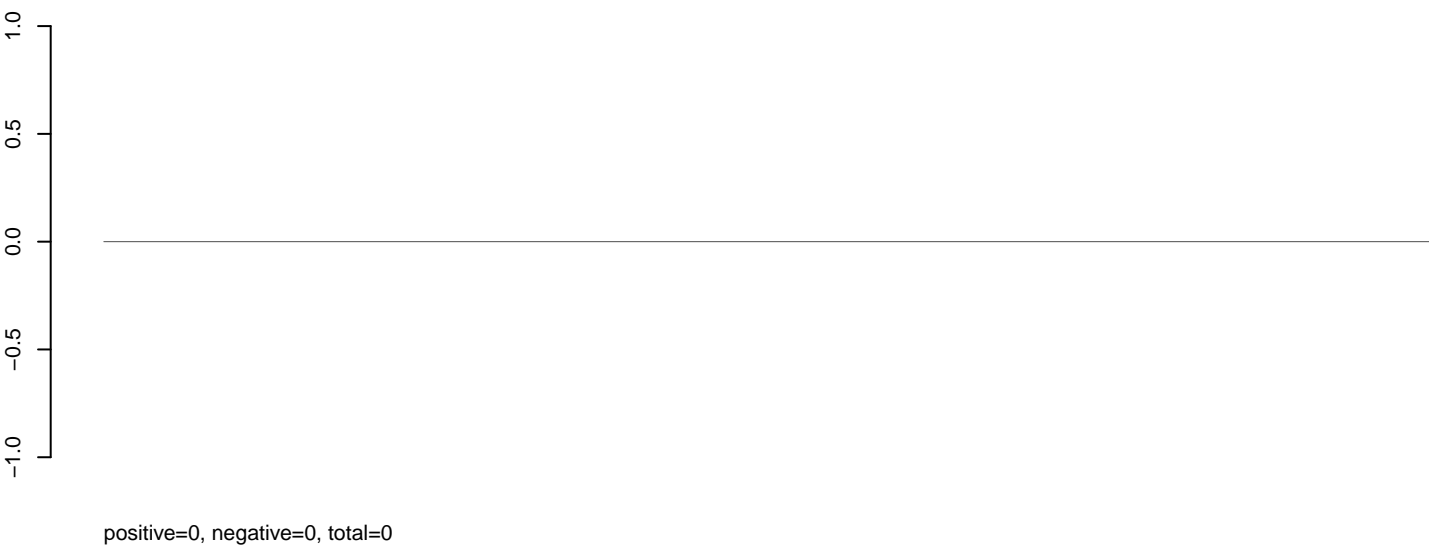


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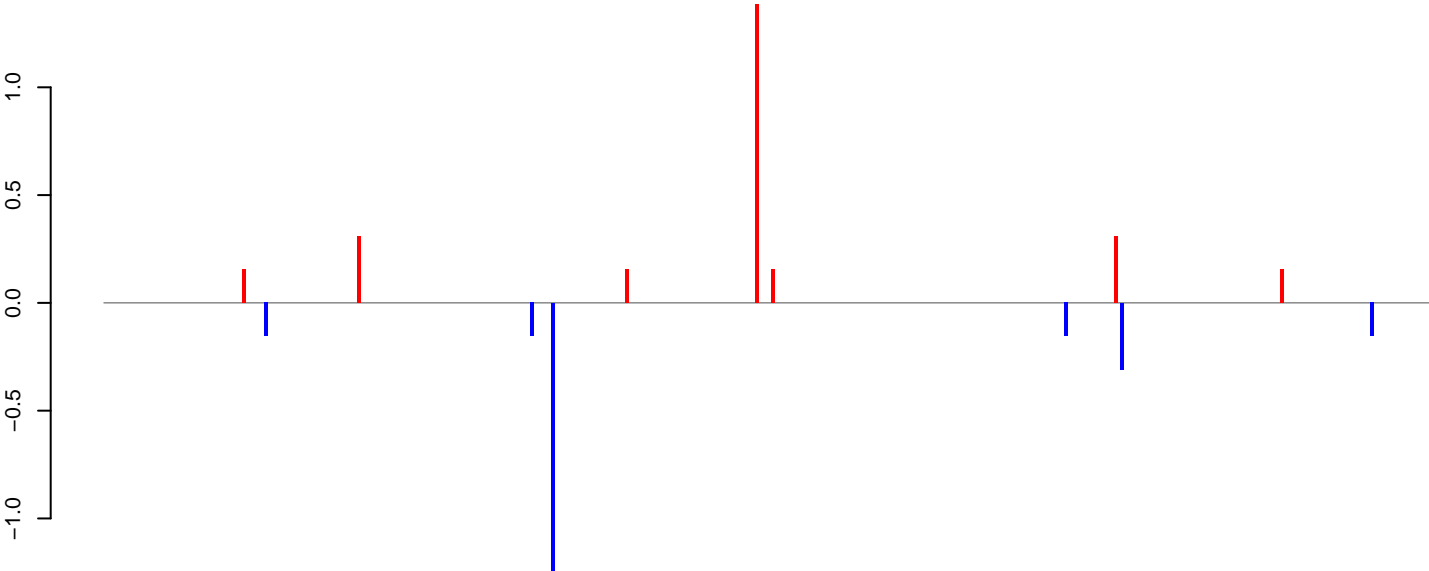
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AeAlbo_C636_6dpi_SINV1.24_35.rep



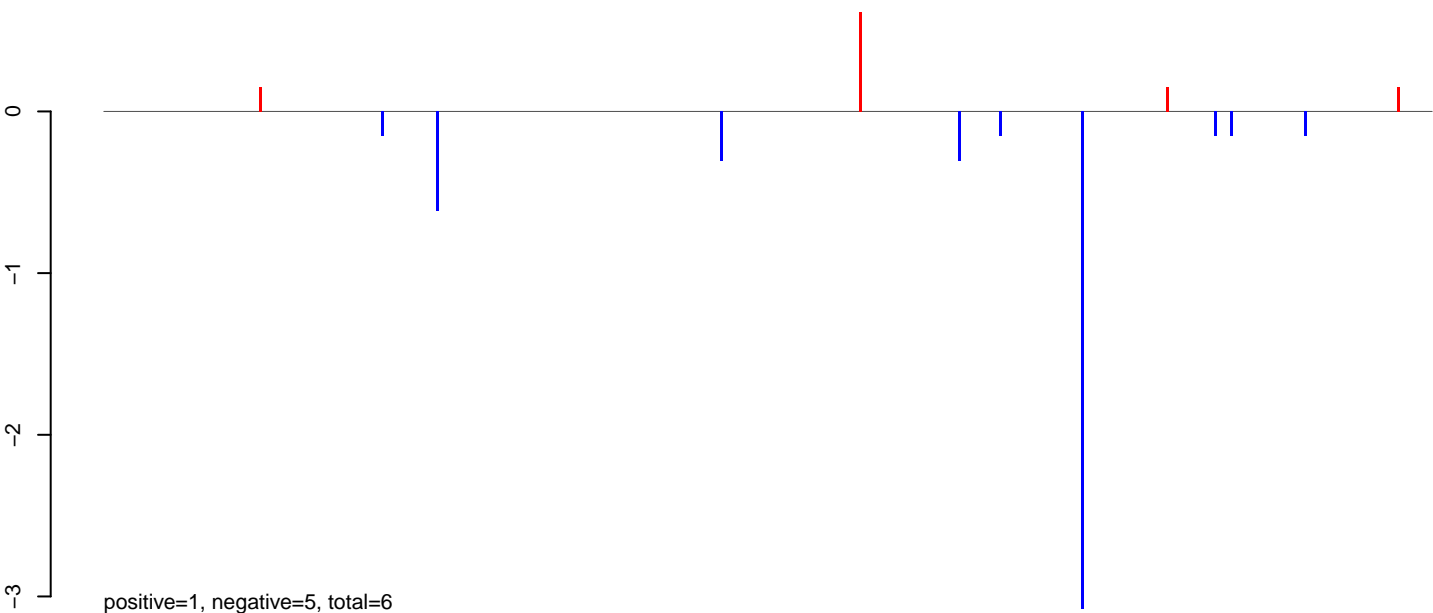
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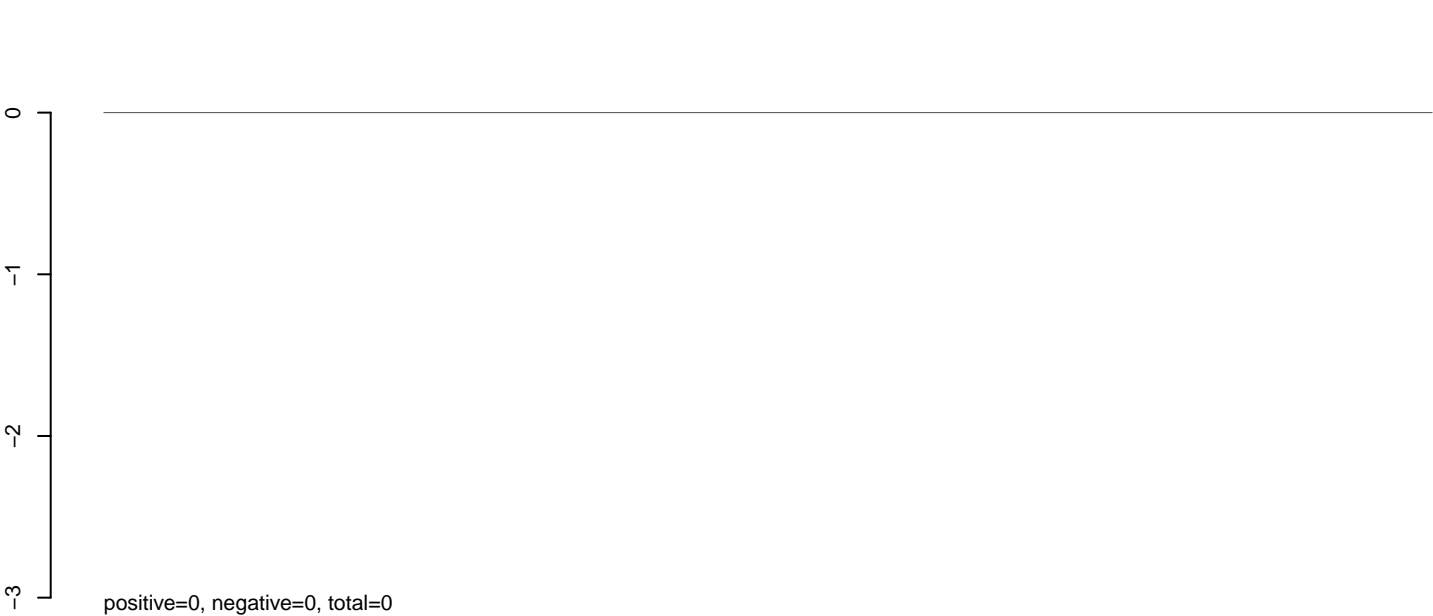
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0 2000 4000 6000 8000 10000

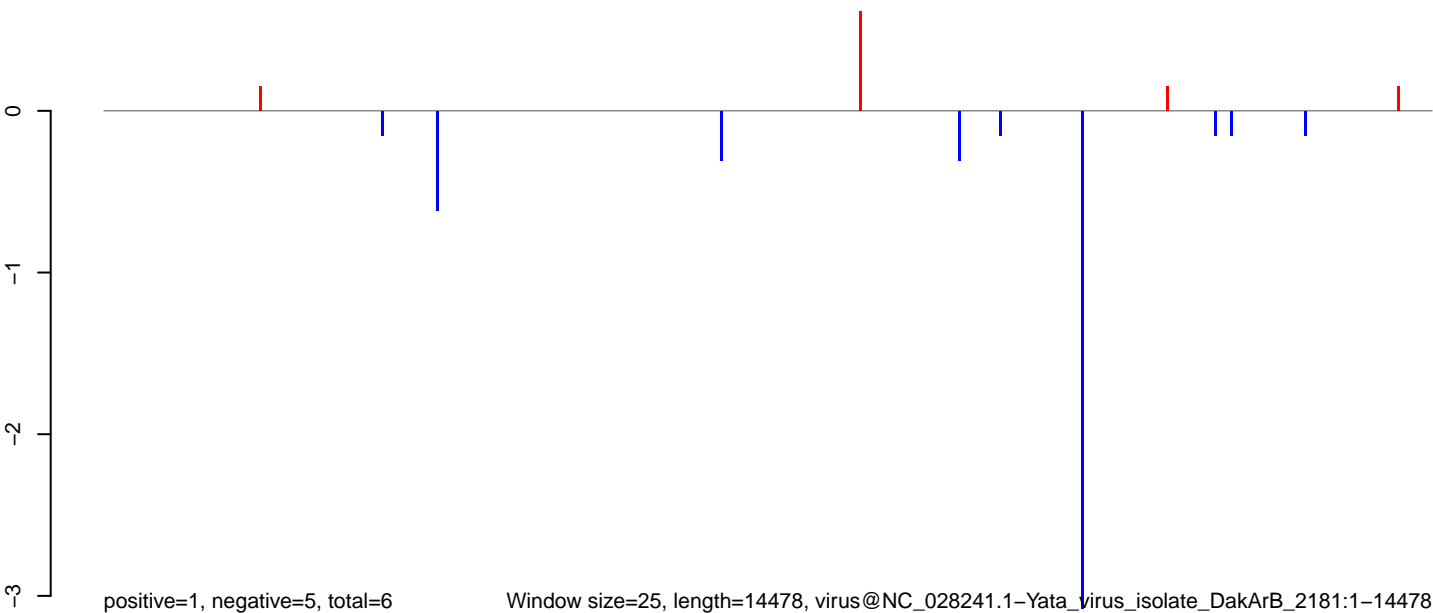
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep

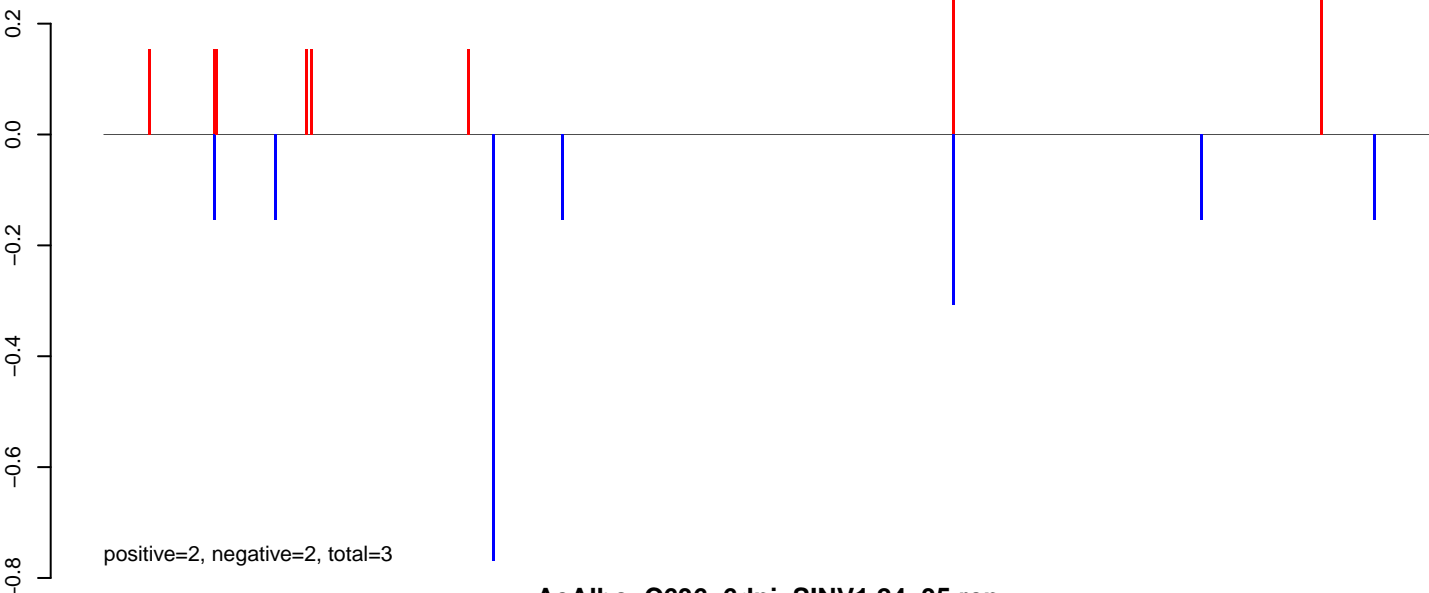


AeAlbo_C636_6dpi_SINV1.rep

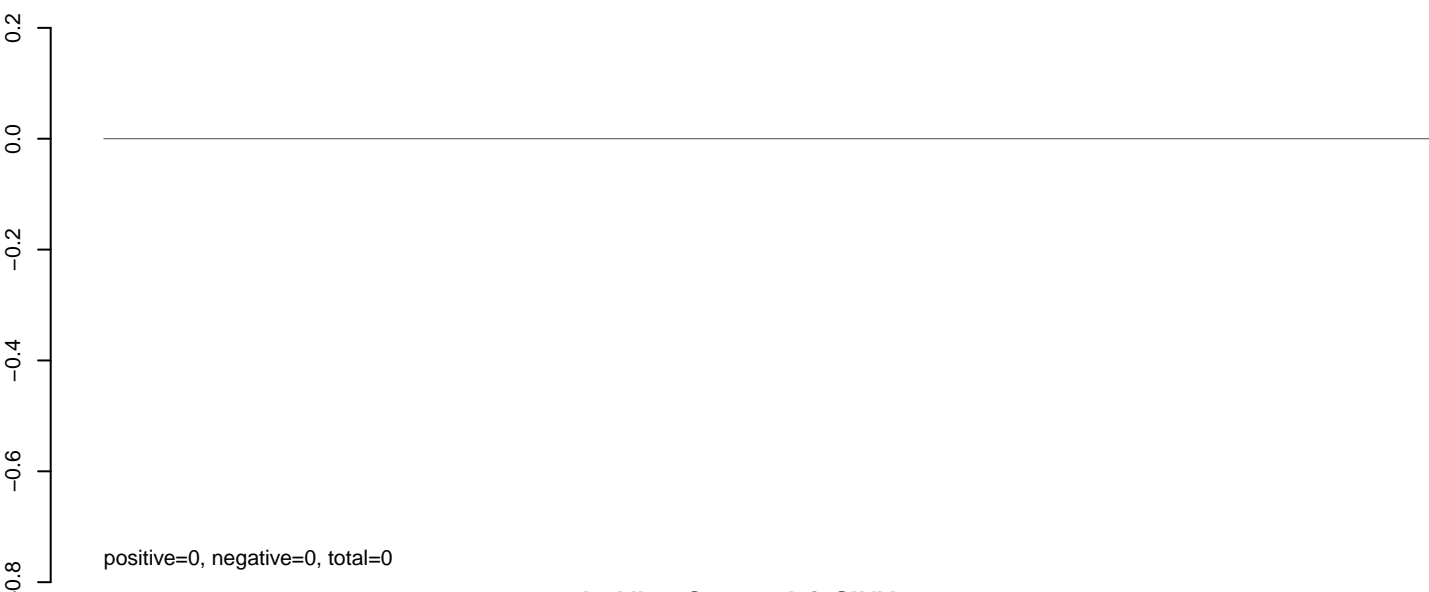


Window size=25, length=14478, virus@NC_028241.1-Yata_virus_isolate_DakArB_2181:1-14478

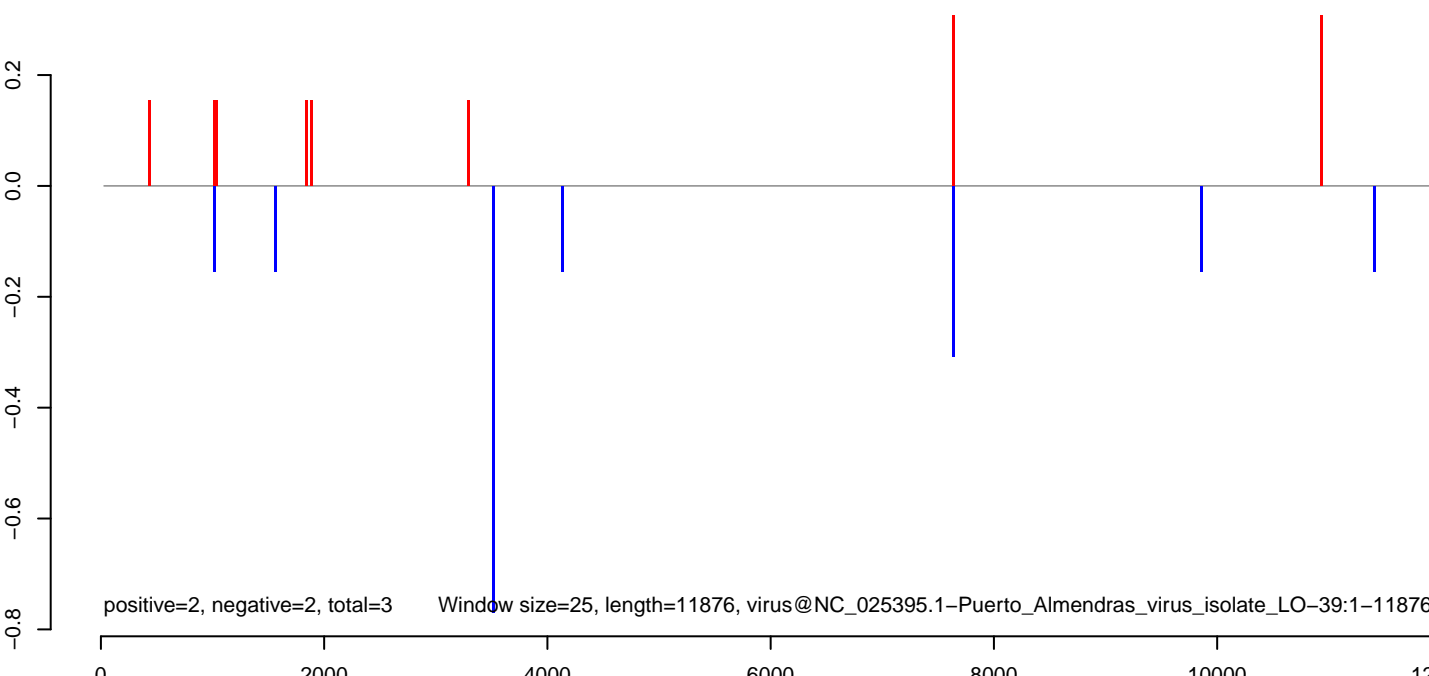
AeAlbo_C636_6dpi_SINV1.18_23.rep



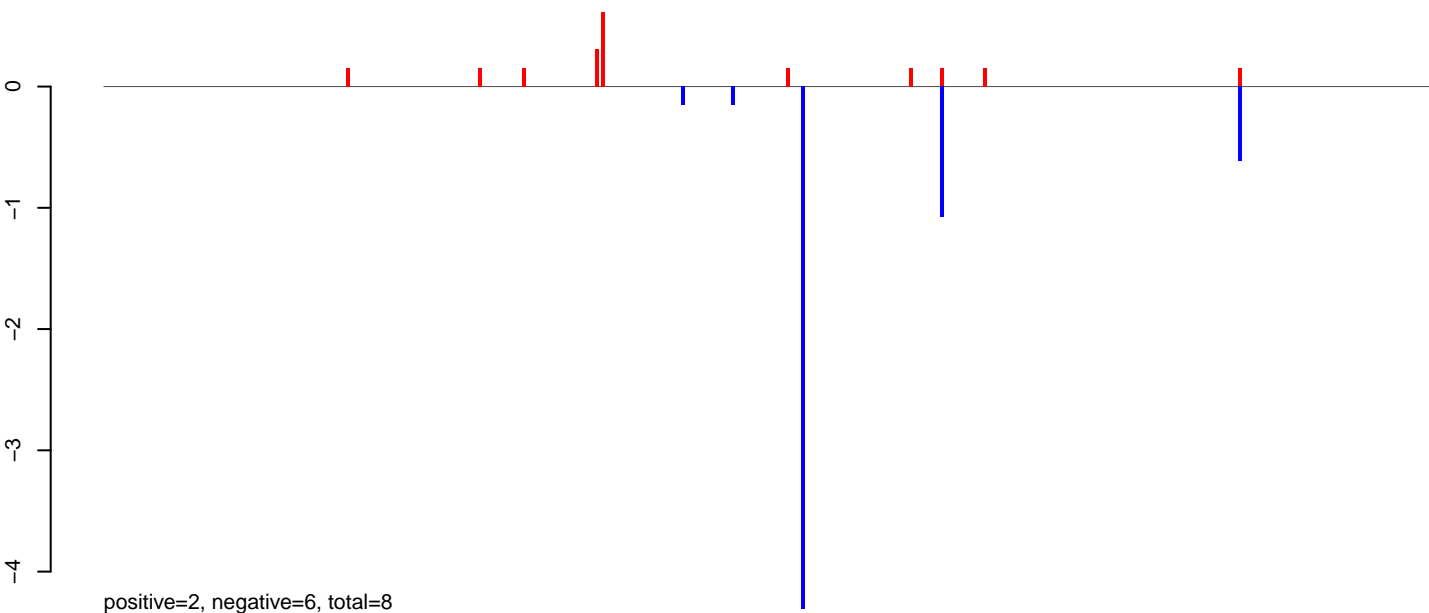
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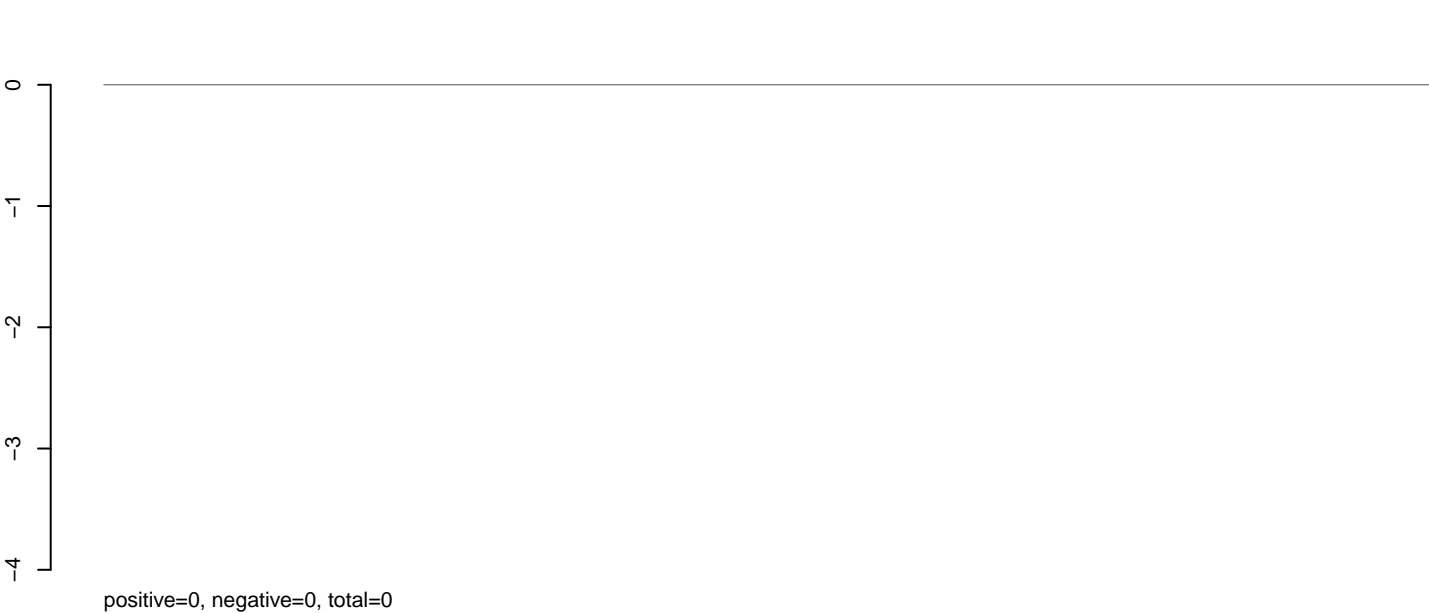
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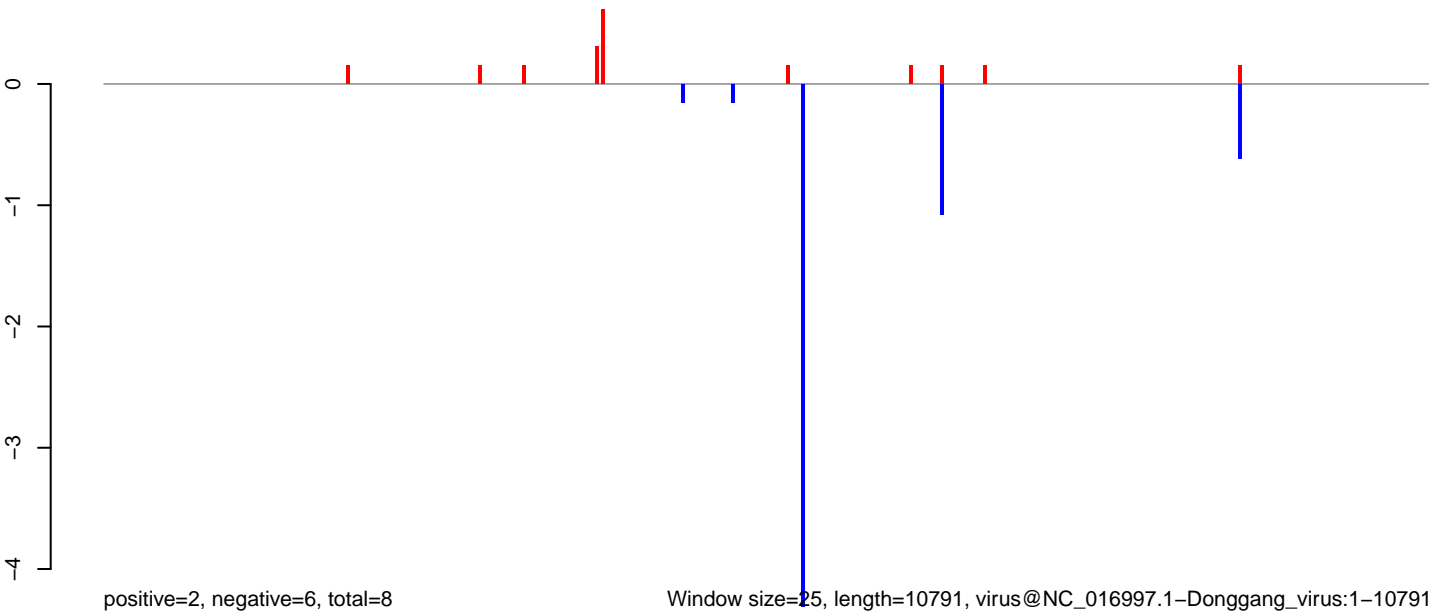
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



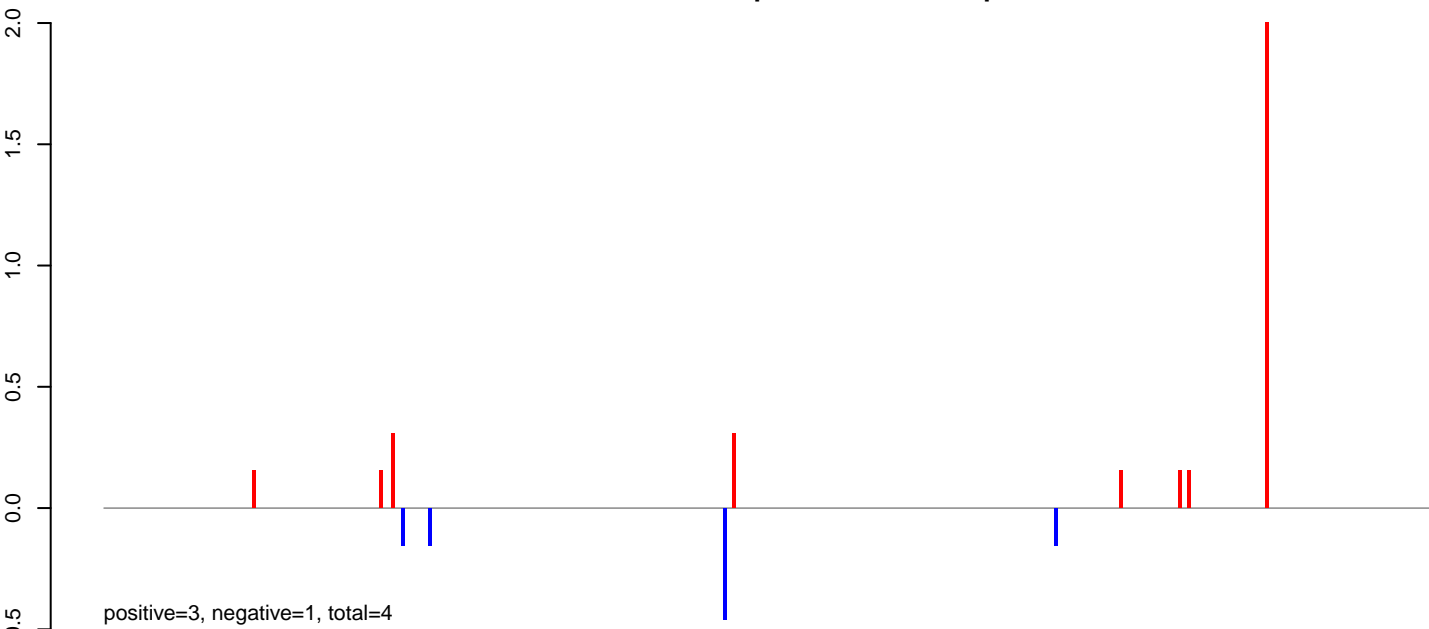
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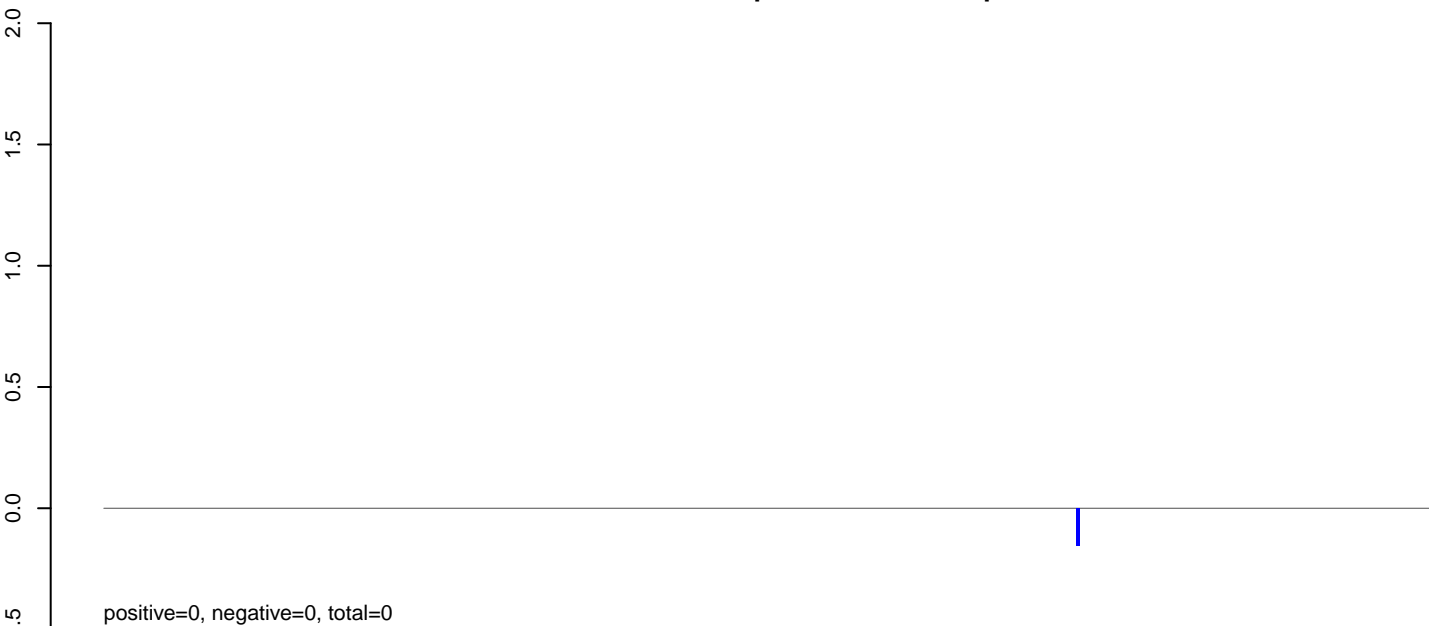
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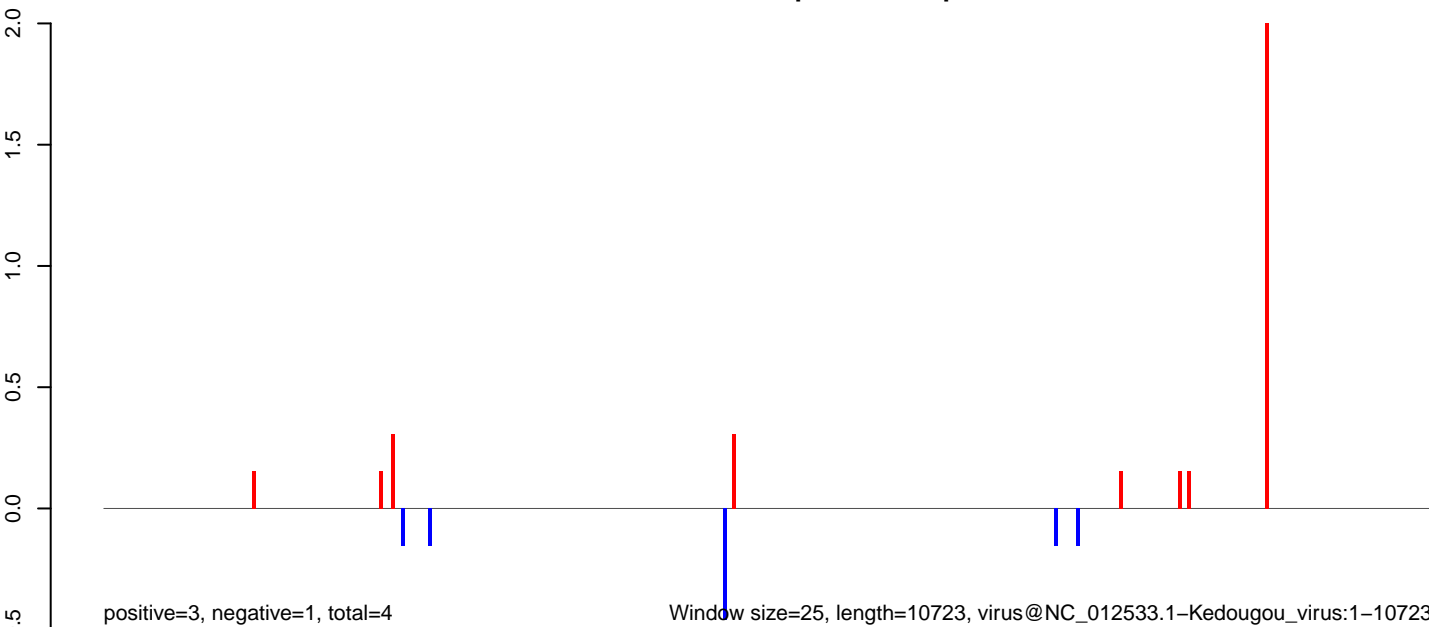
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AeAlbo_C636_6dpi_SINV1.24_35.rep

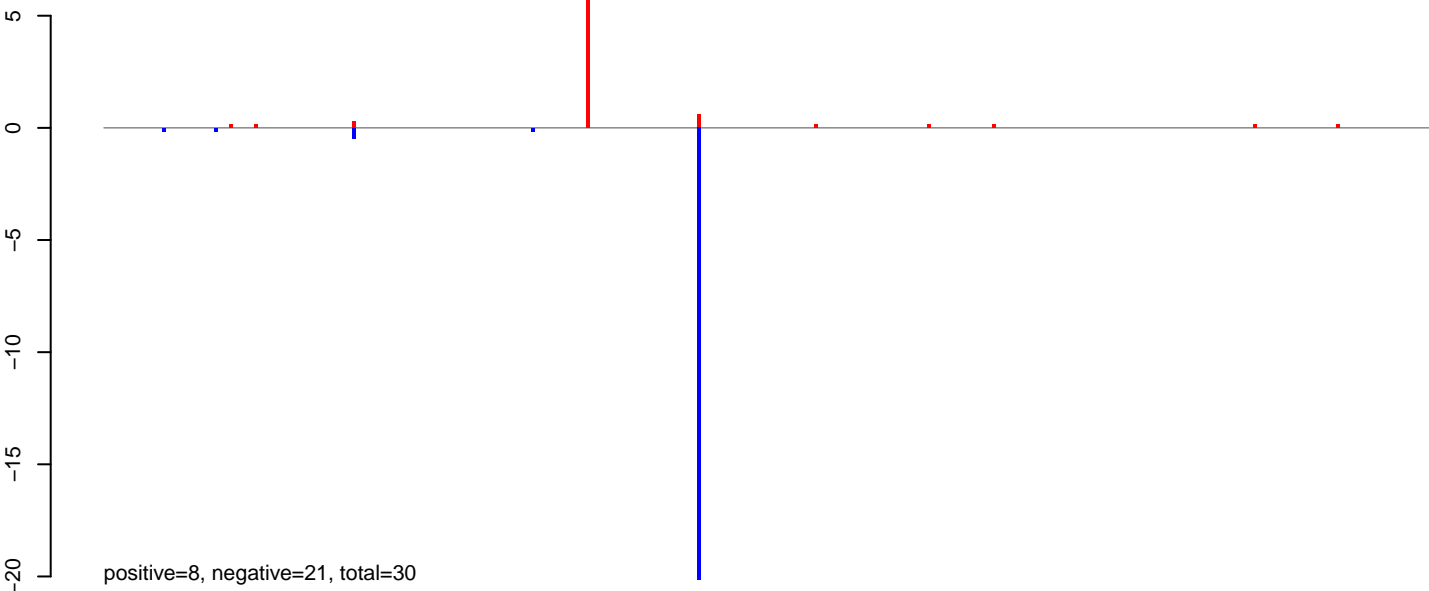


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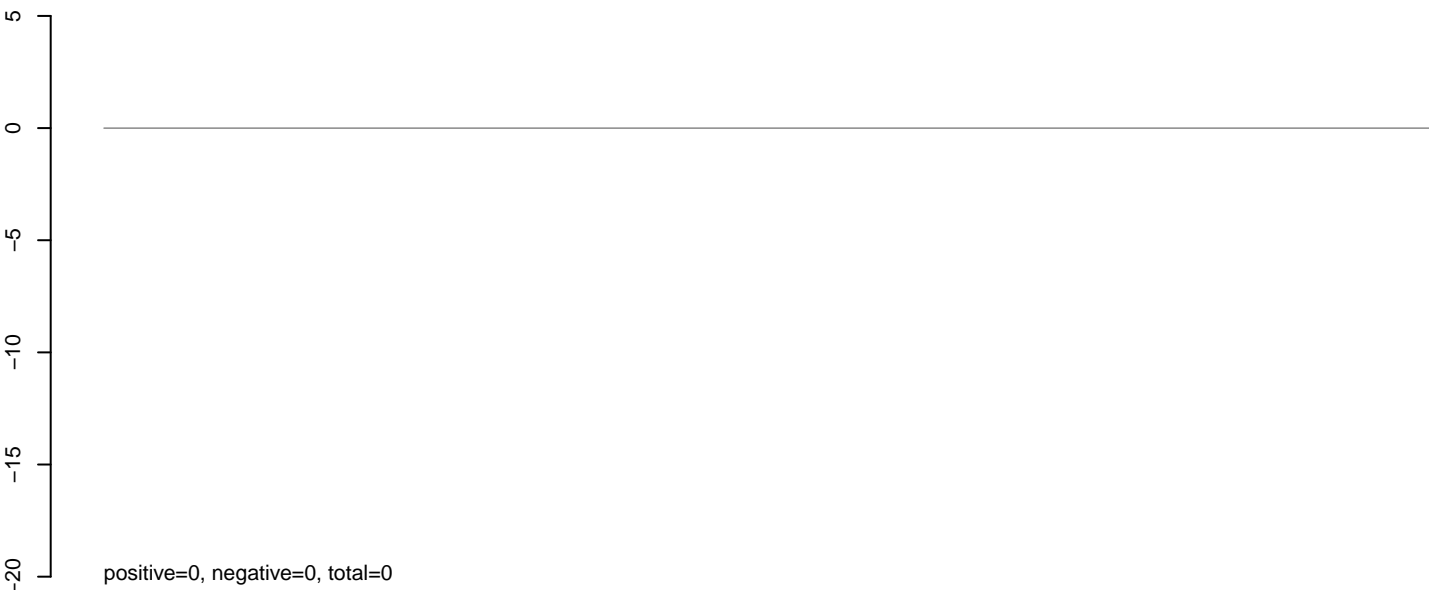


0 2000 4000 6000 8000 10000

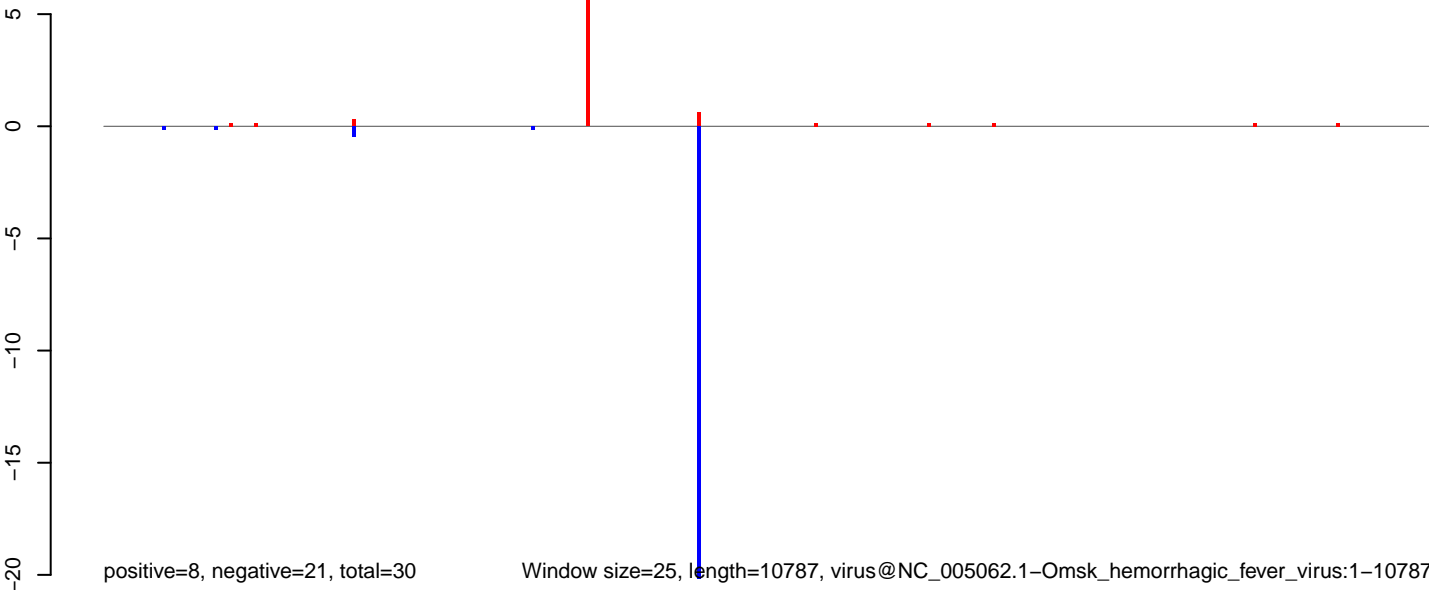
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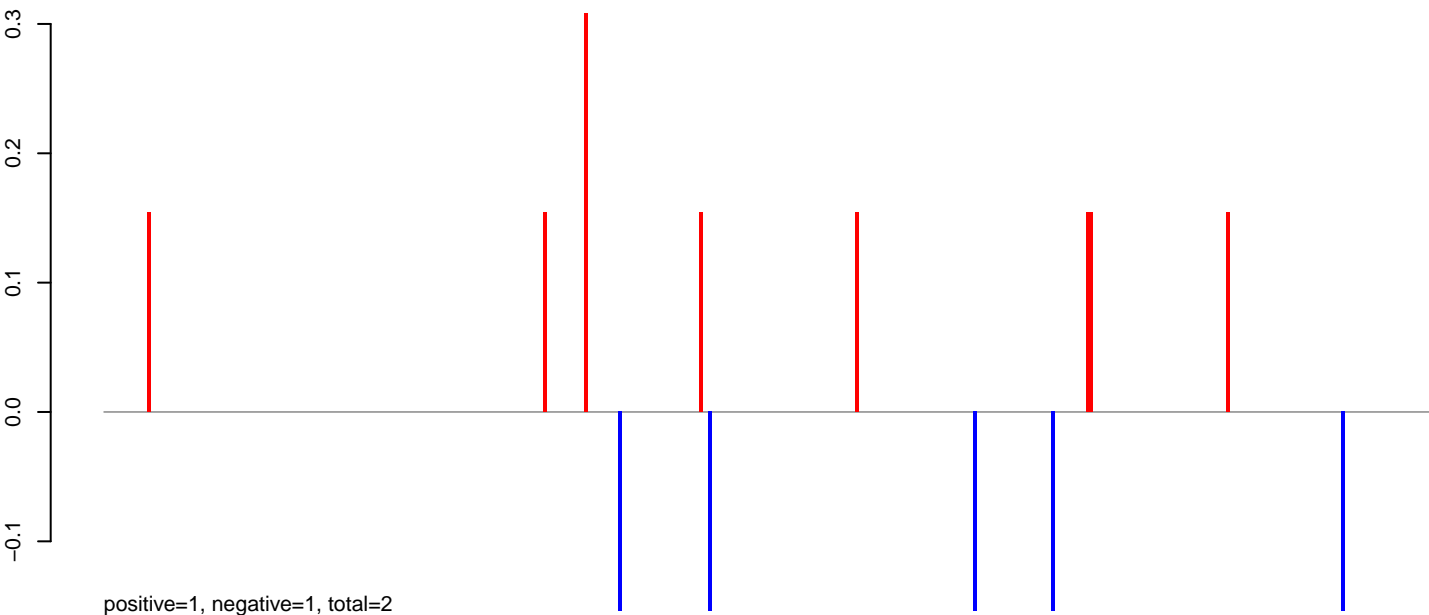
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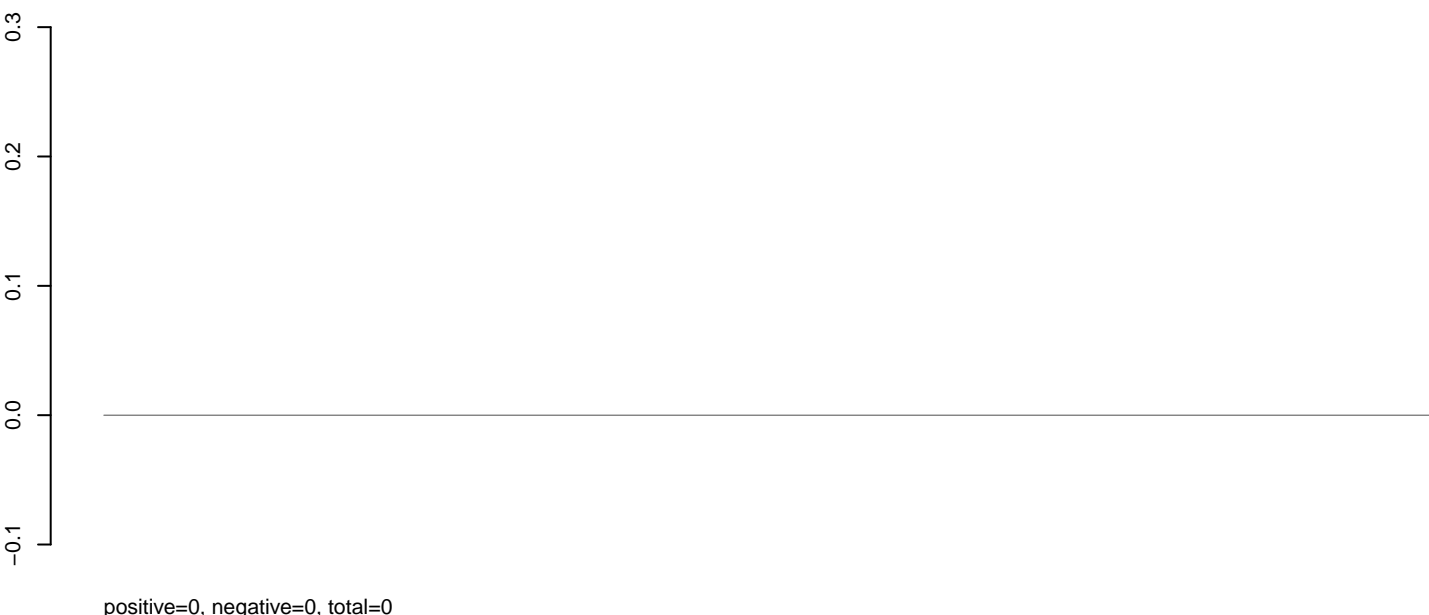
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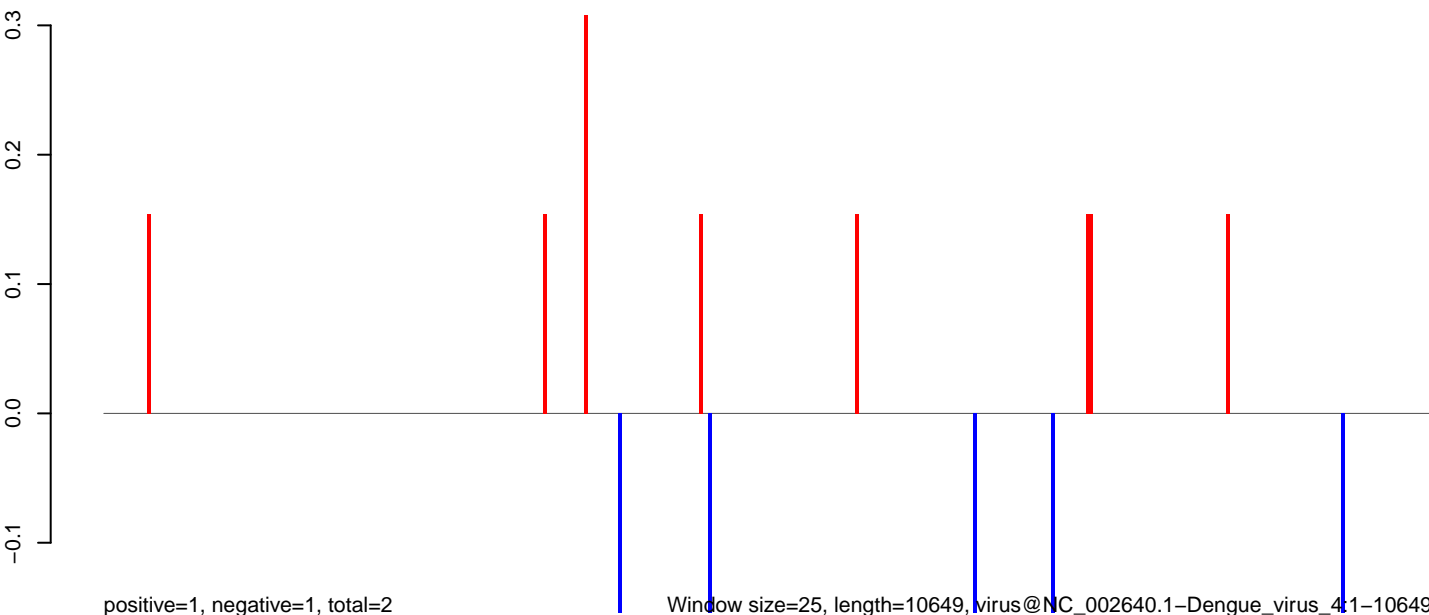
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AeAlbo_C636_6dpi_SINV1.24_35.rep



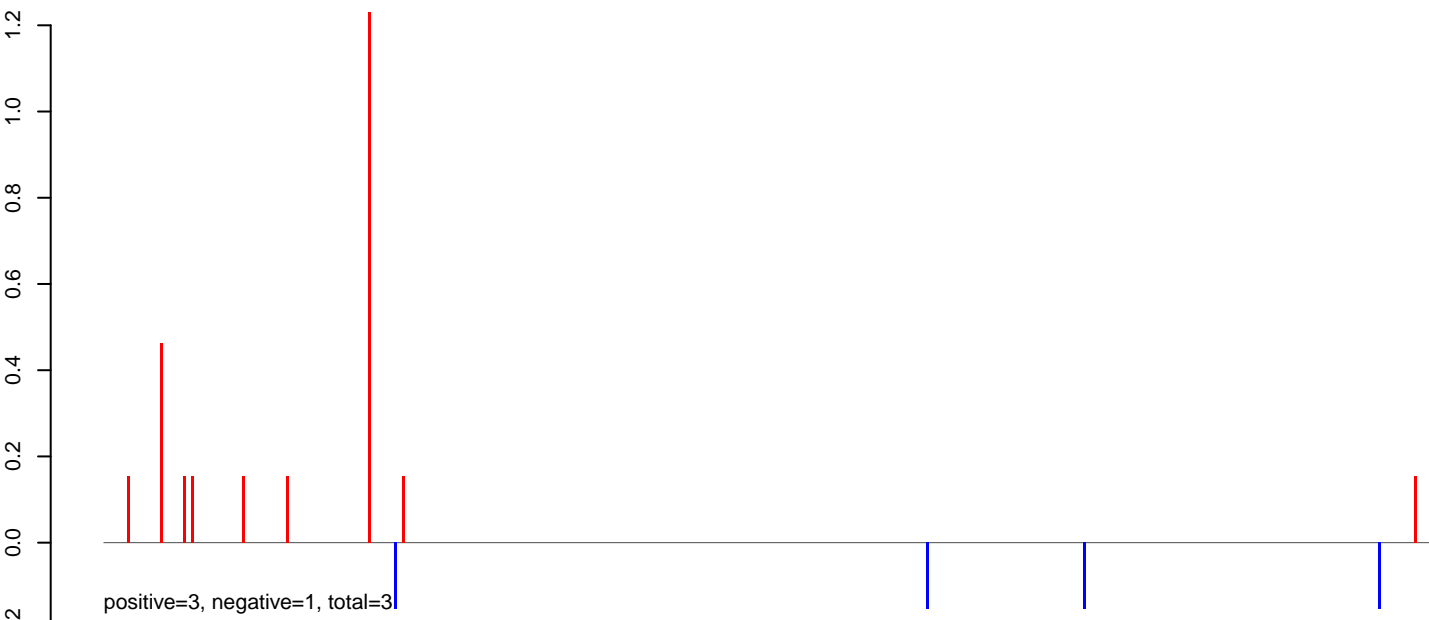
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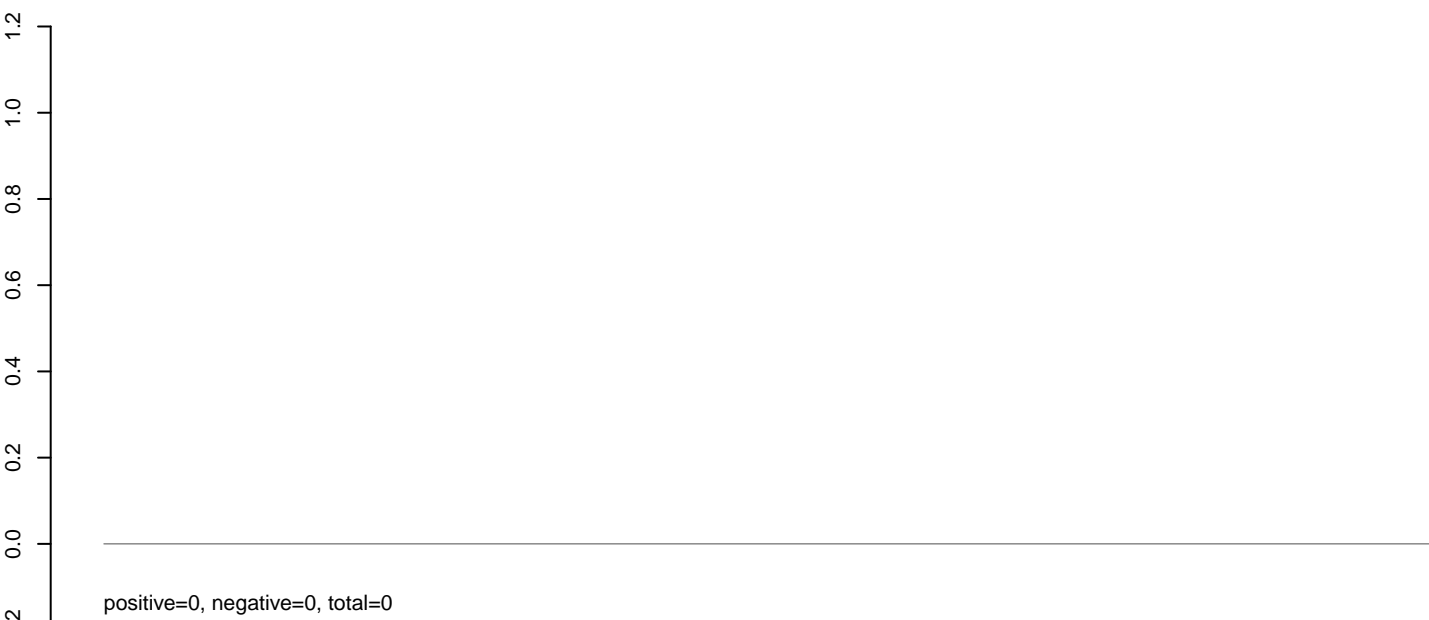
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0 2000 4000 6000 8000 10000

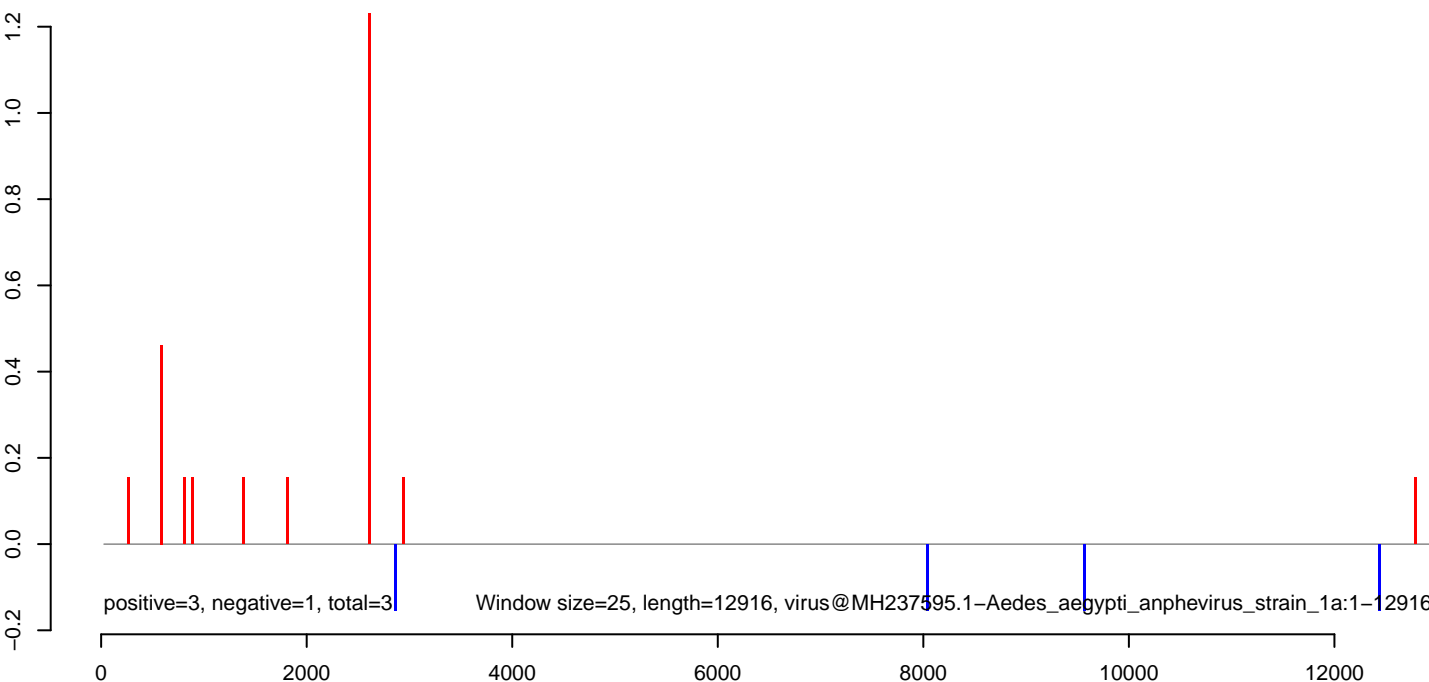
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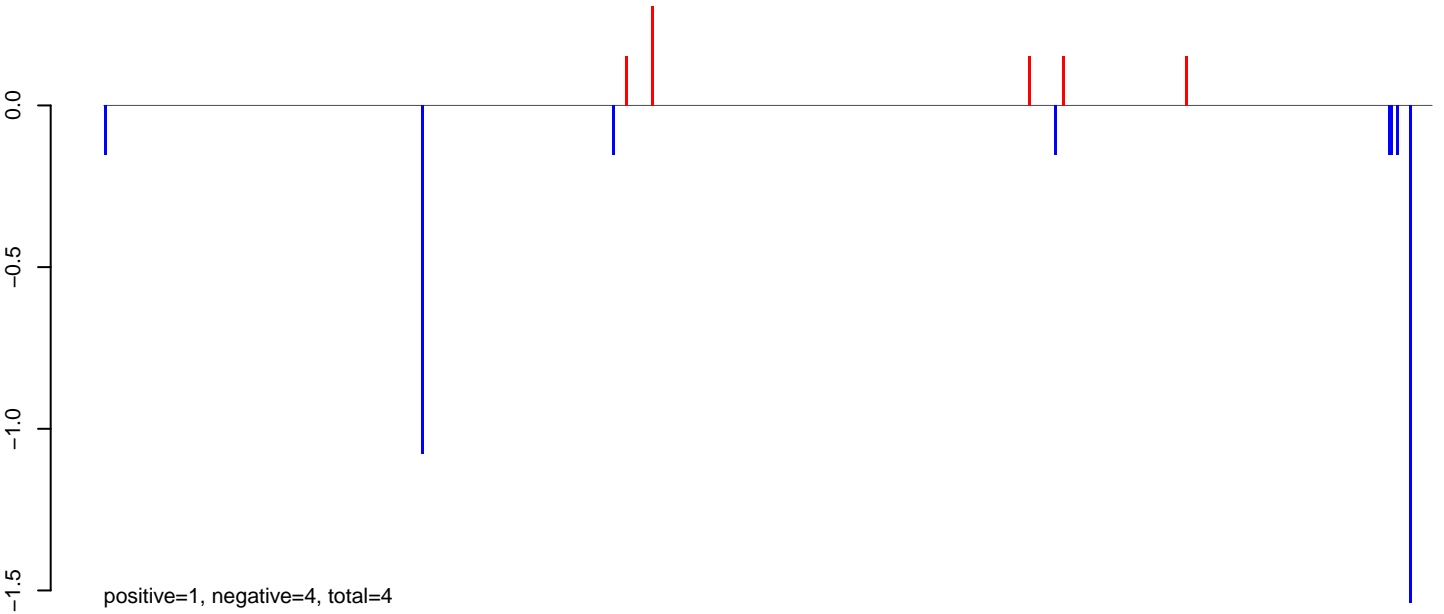
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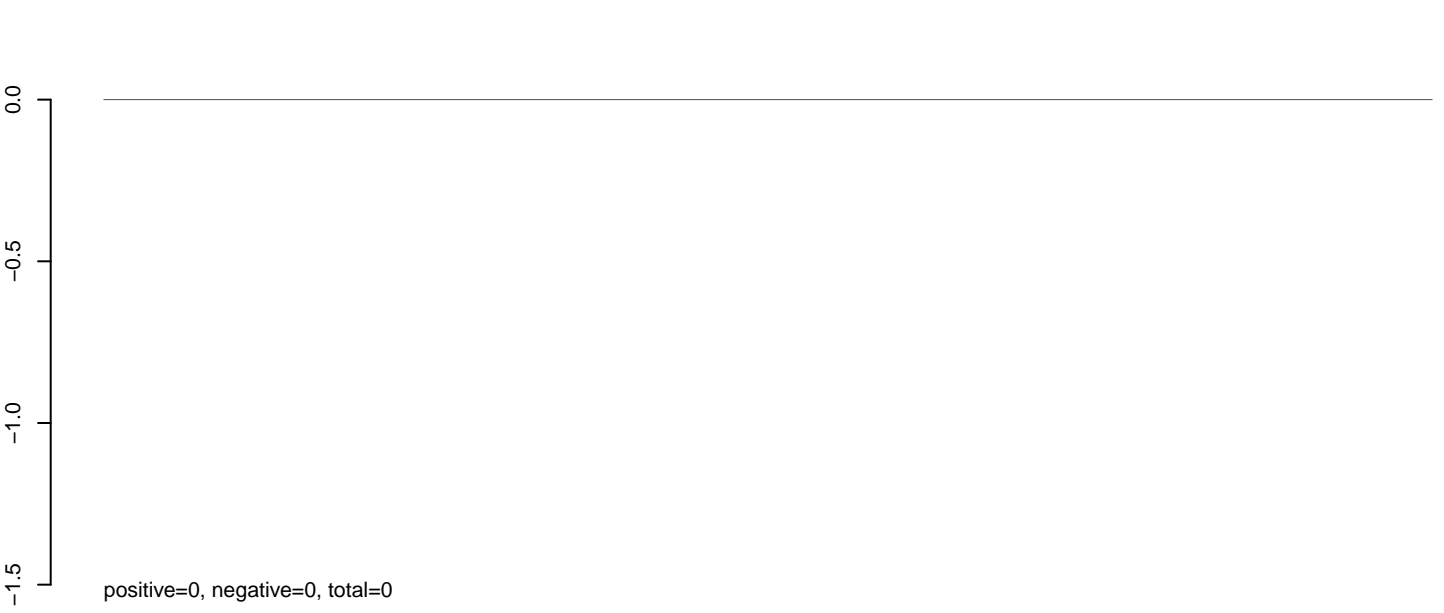
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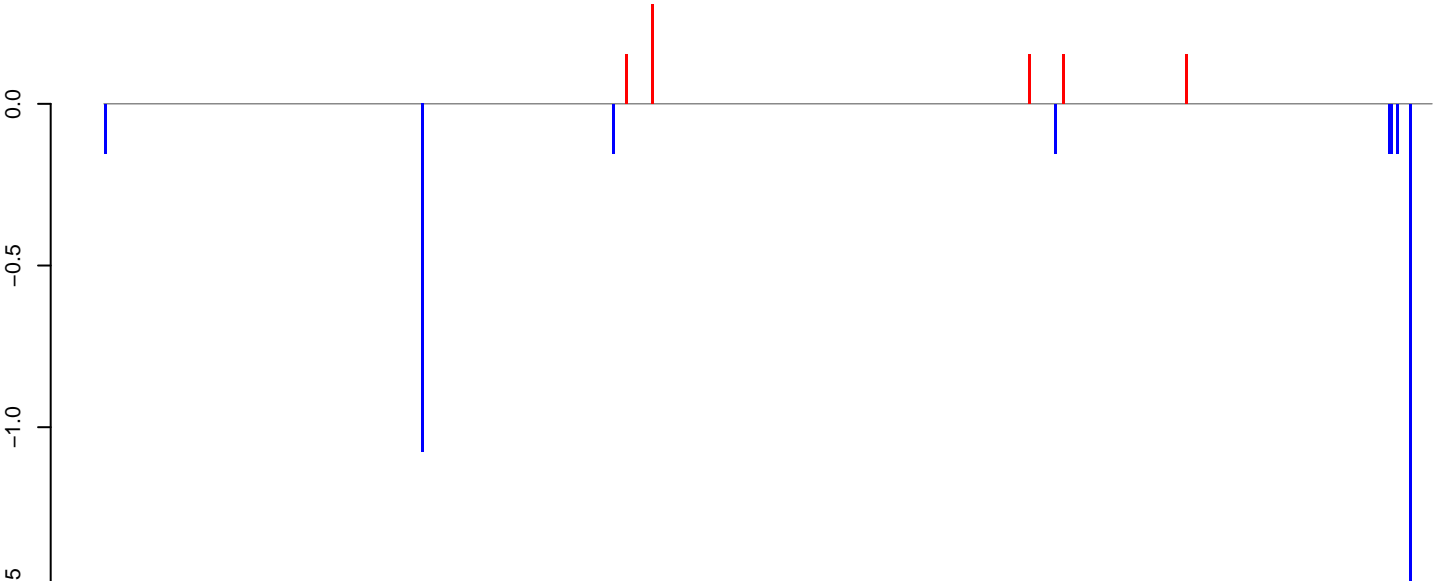
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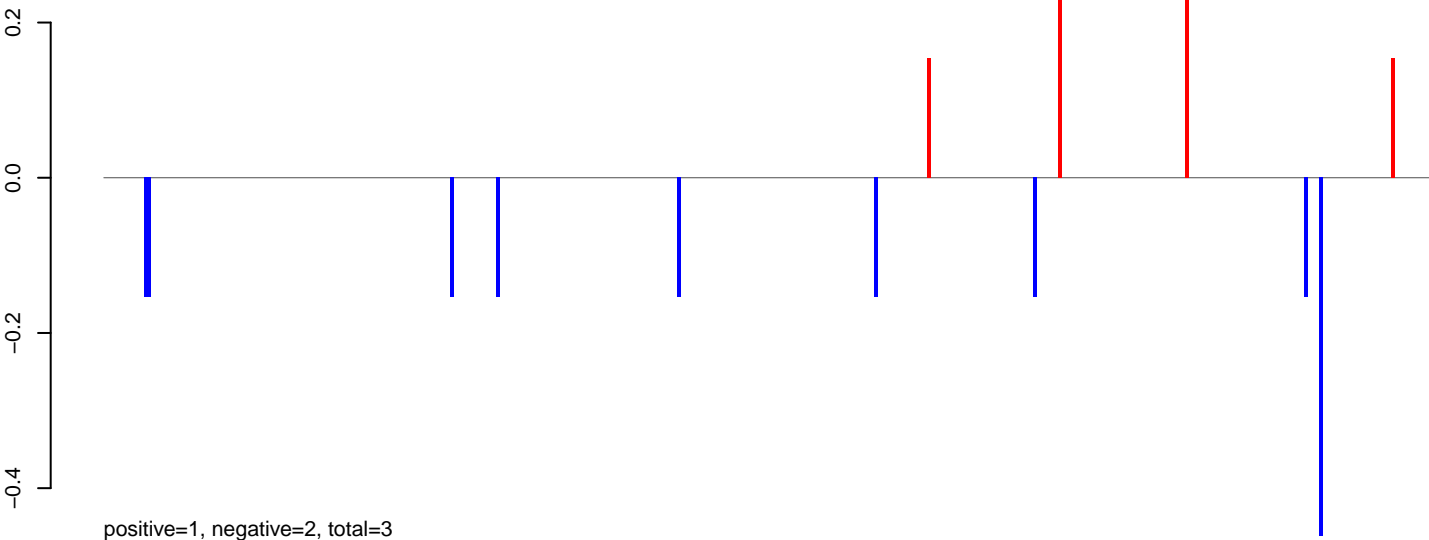
AeAlbo_C636_6dpi_SINV1.24_35.rep



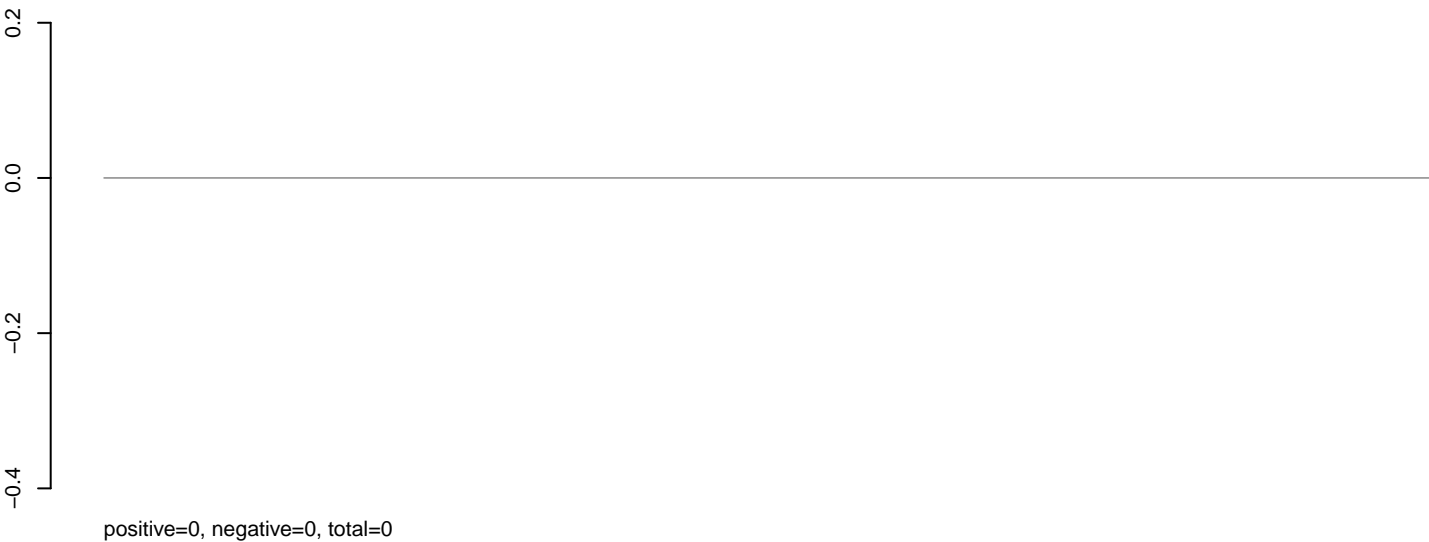
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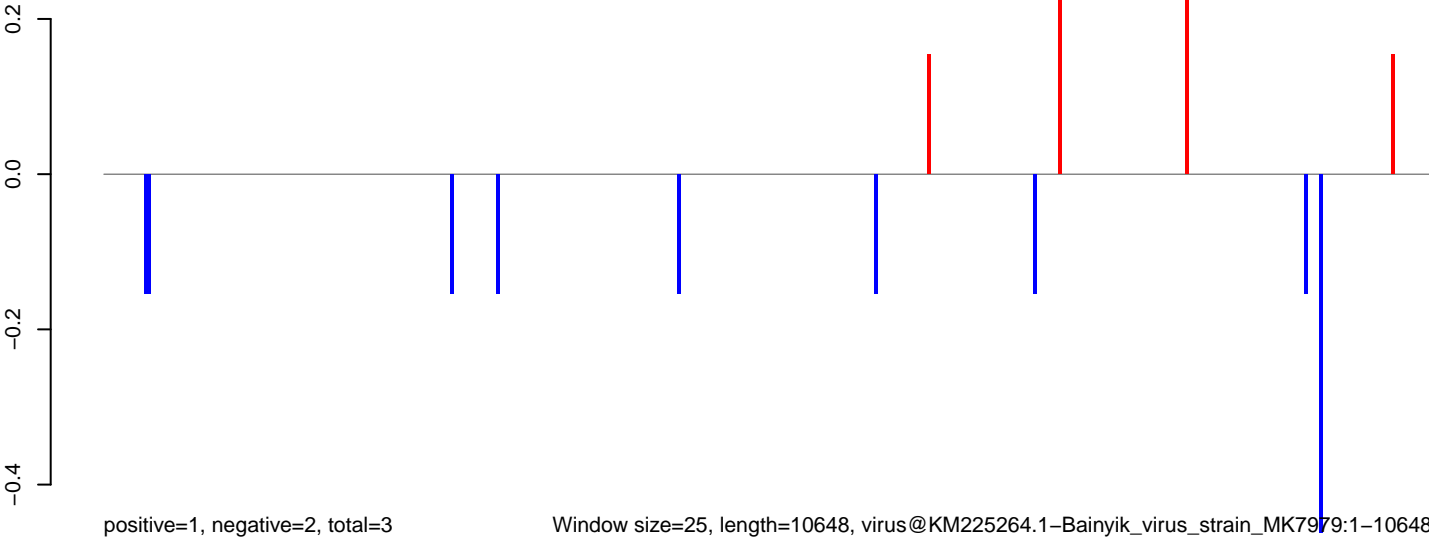
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AeAlbo_C636_6dpi_SINV1.24_35.rep



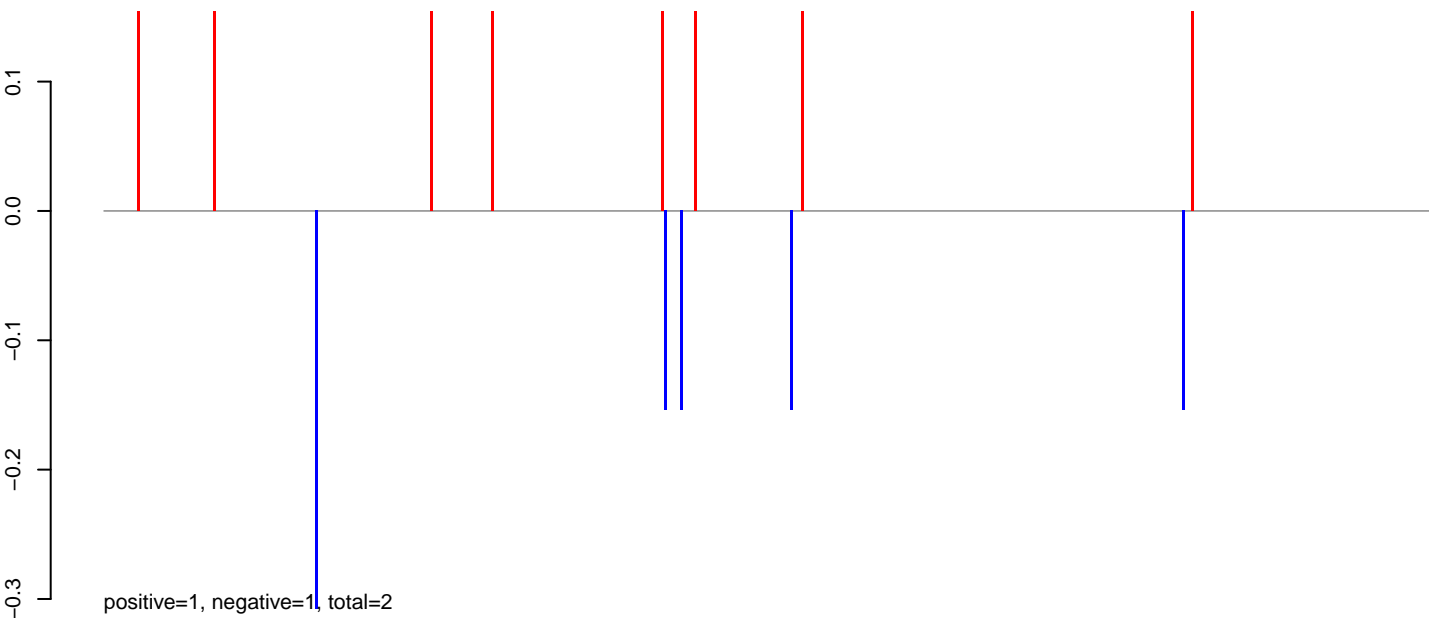
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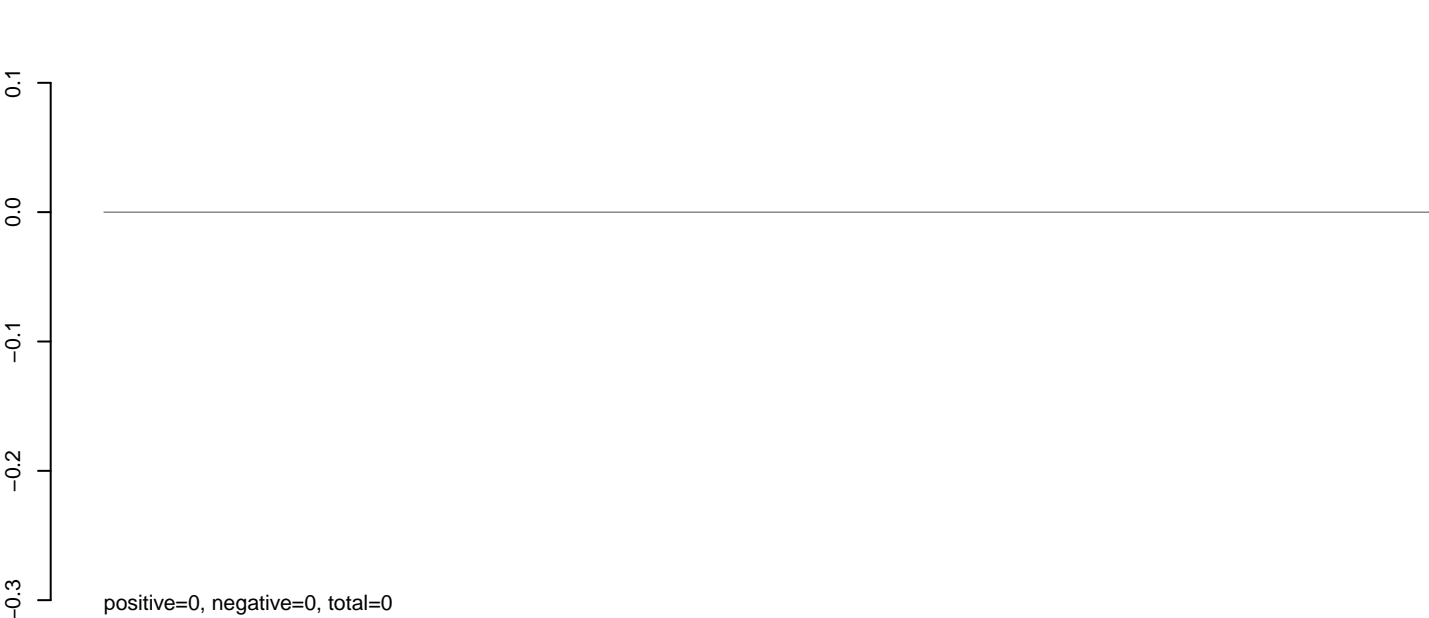
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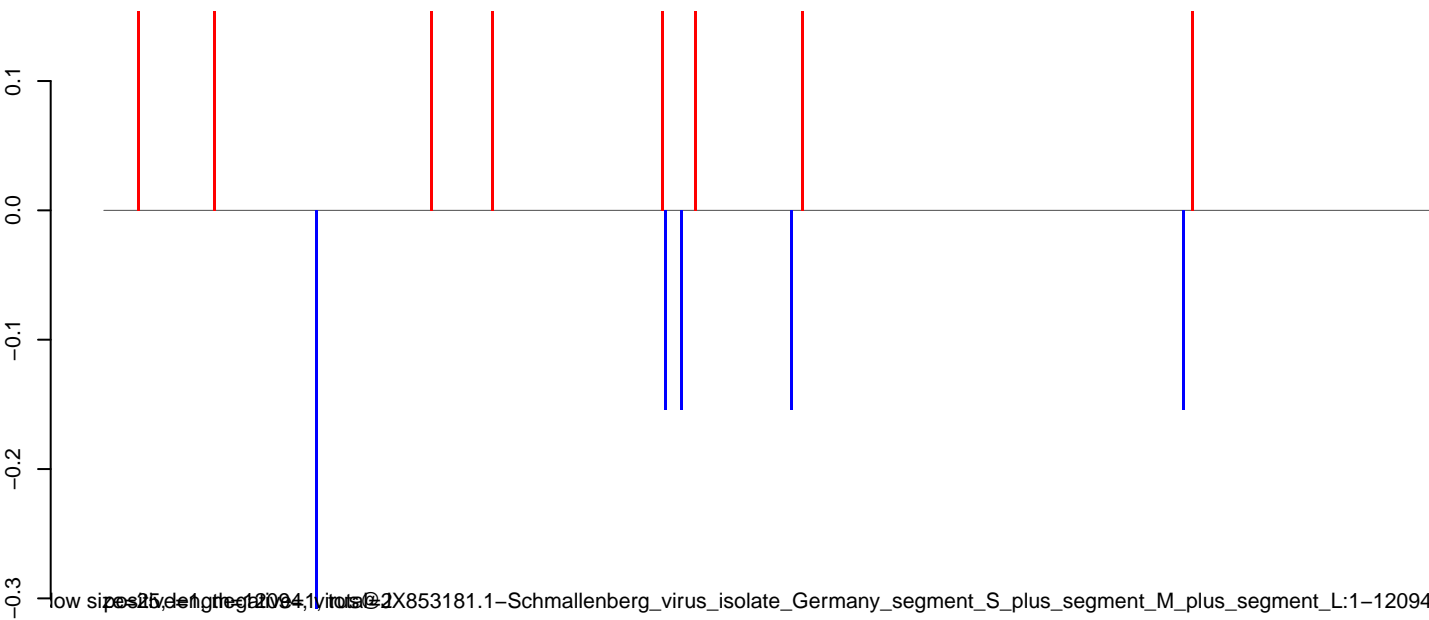
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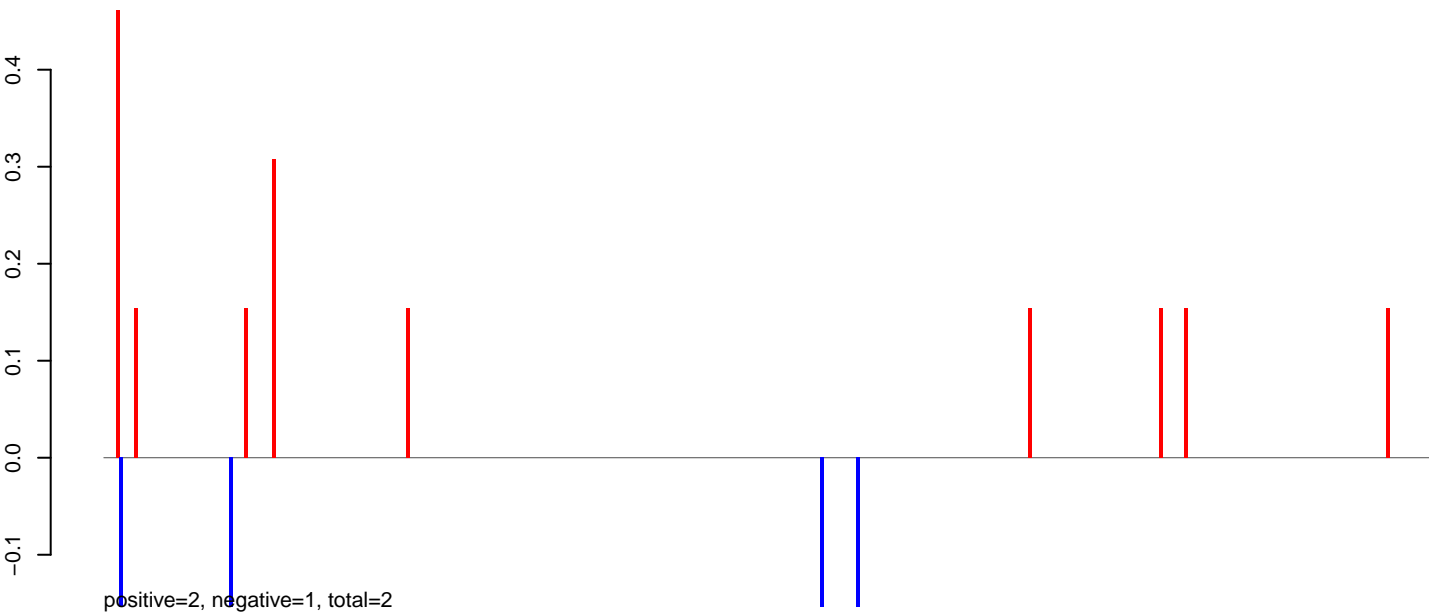
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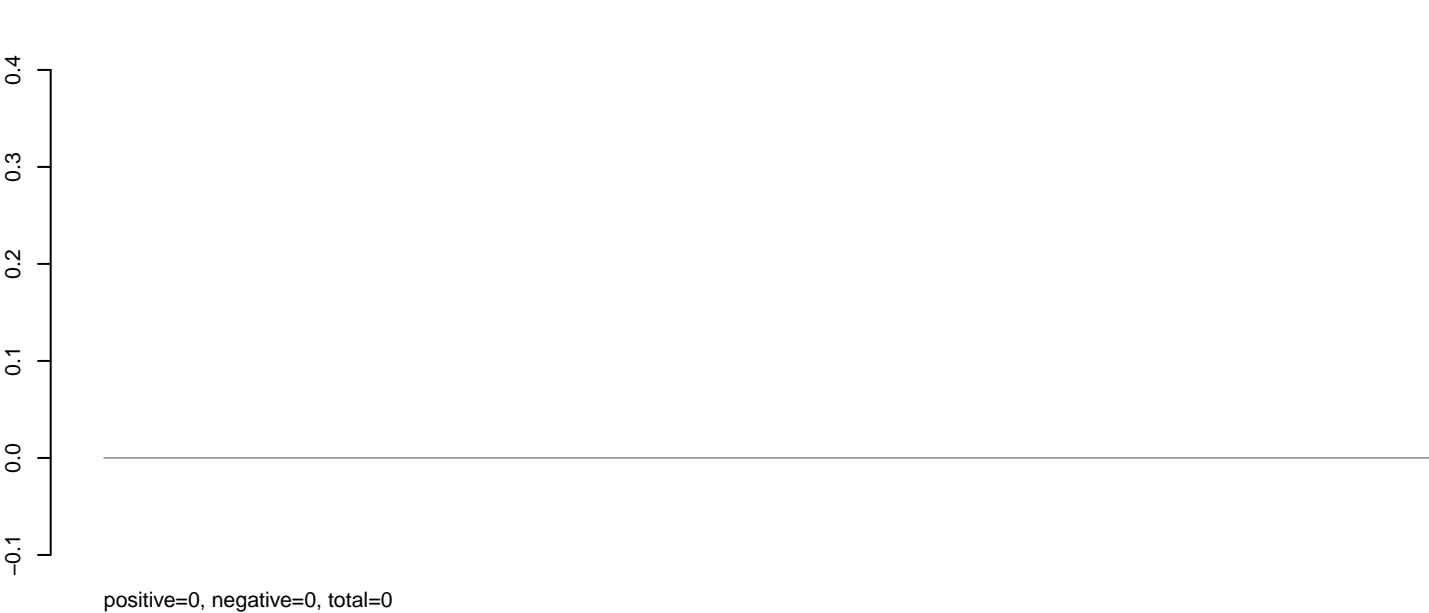
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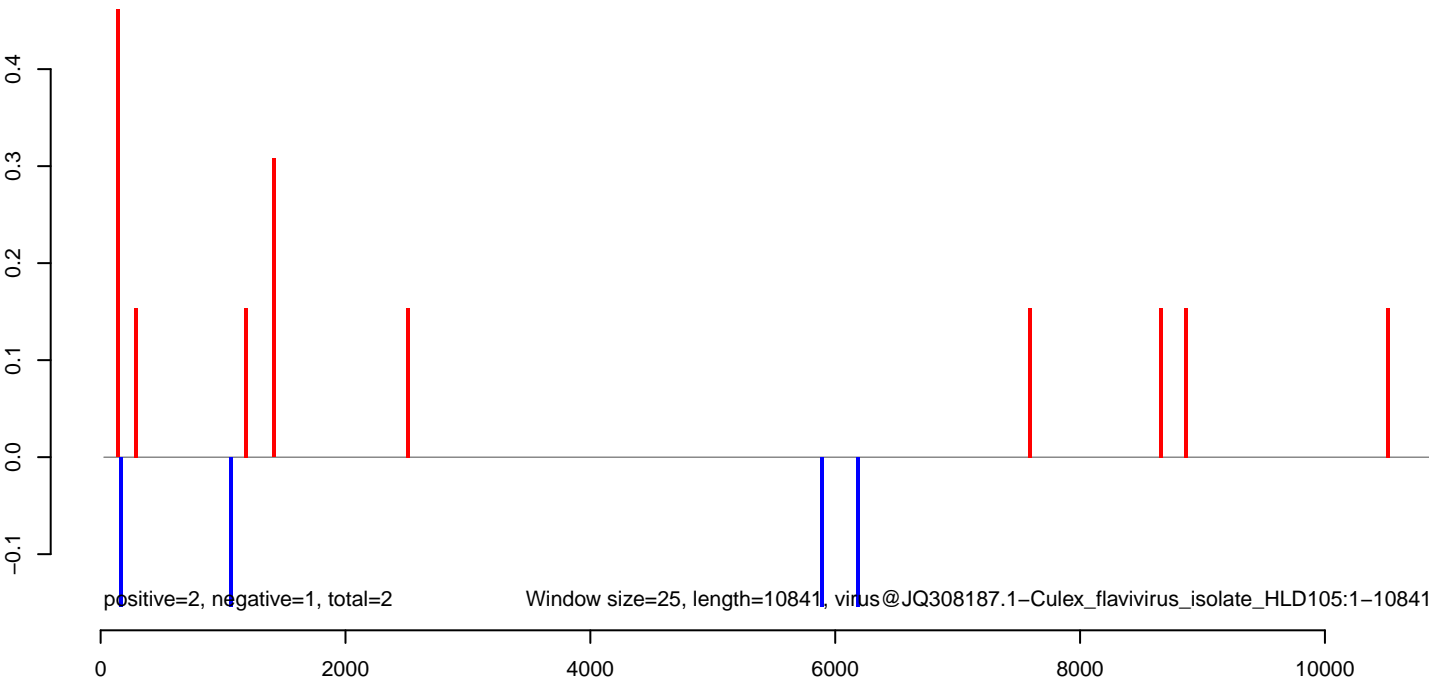
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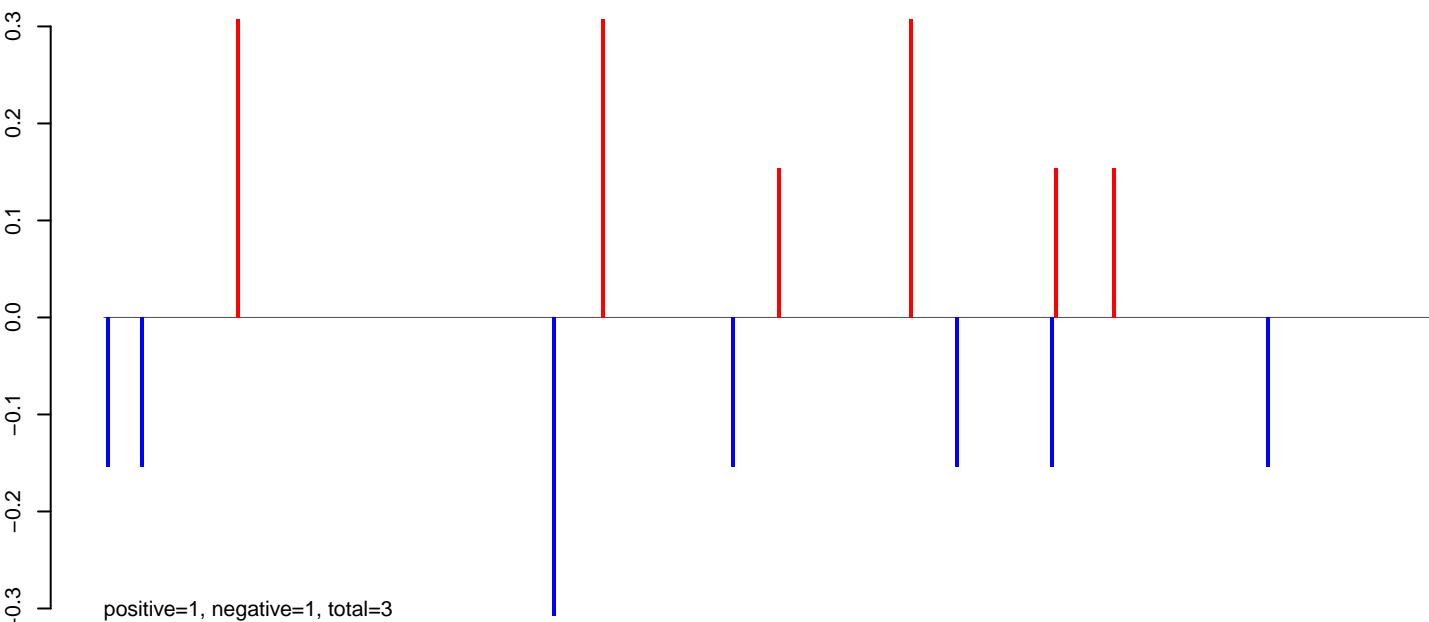
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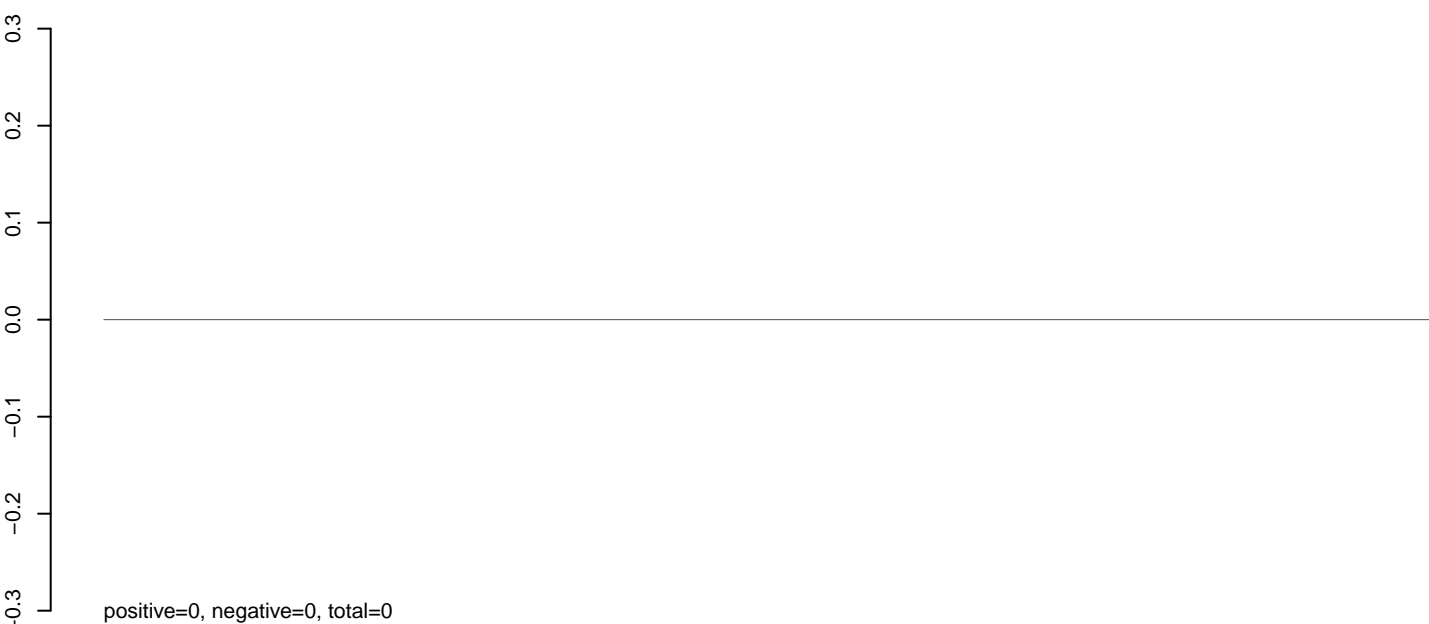
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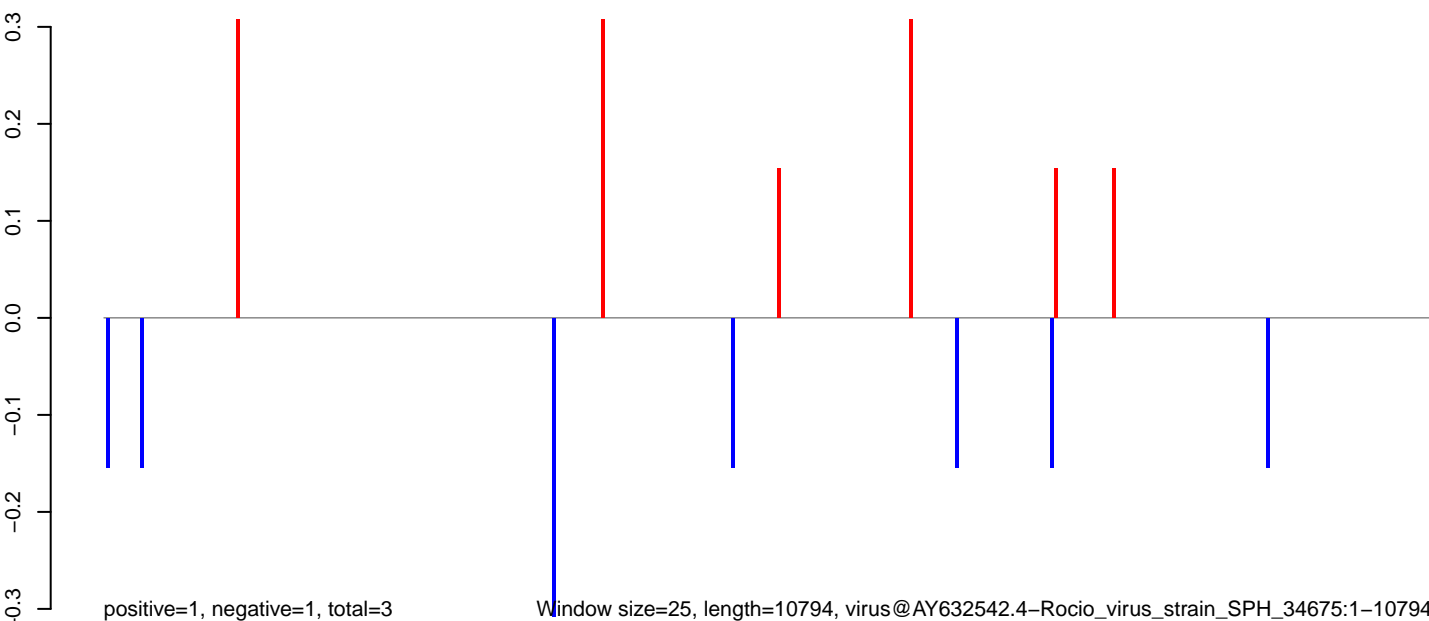
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AeAlbo_C636_6dpi_SINV1.24_35.rep



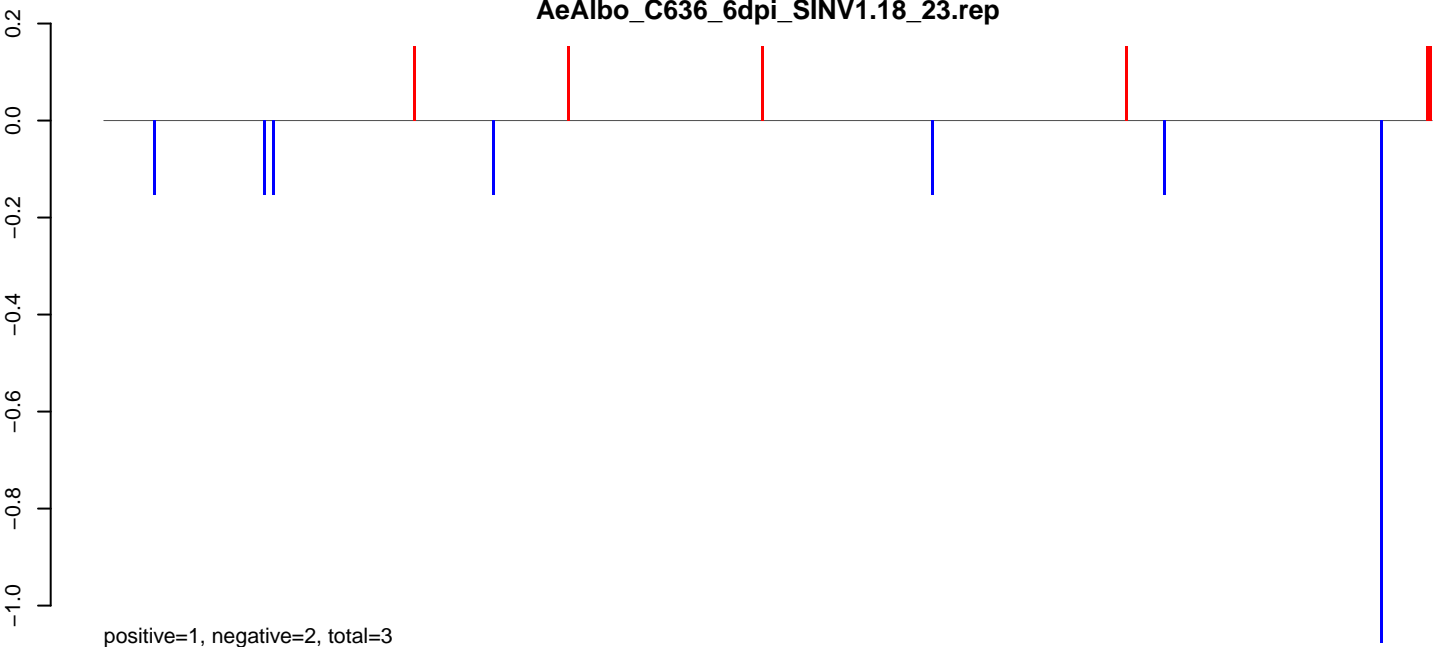
AeAlbo_C636_6dpi_SINV1.rep



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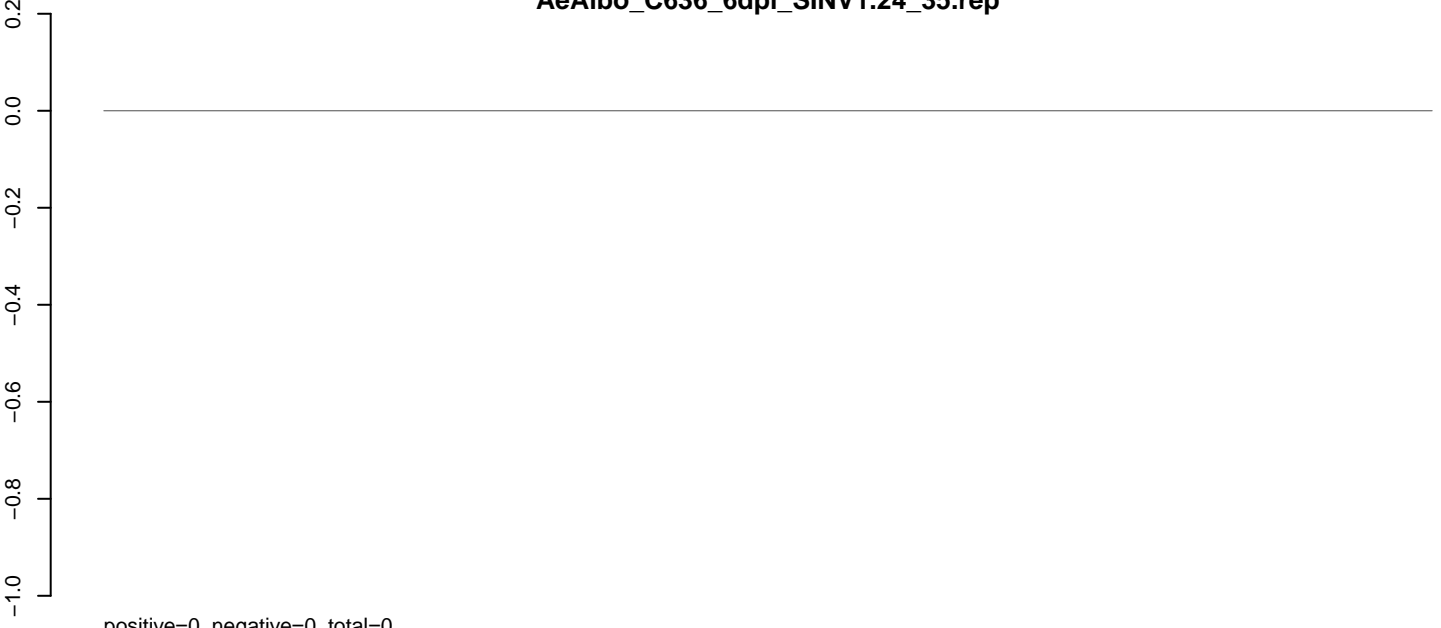
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AeAlbo_C636_6dpi_SINV1.18_23.rep



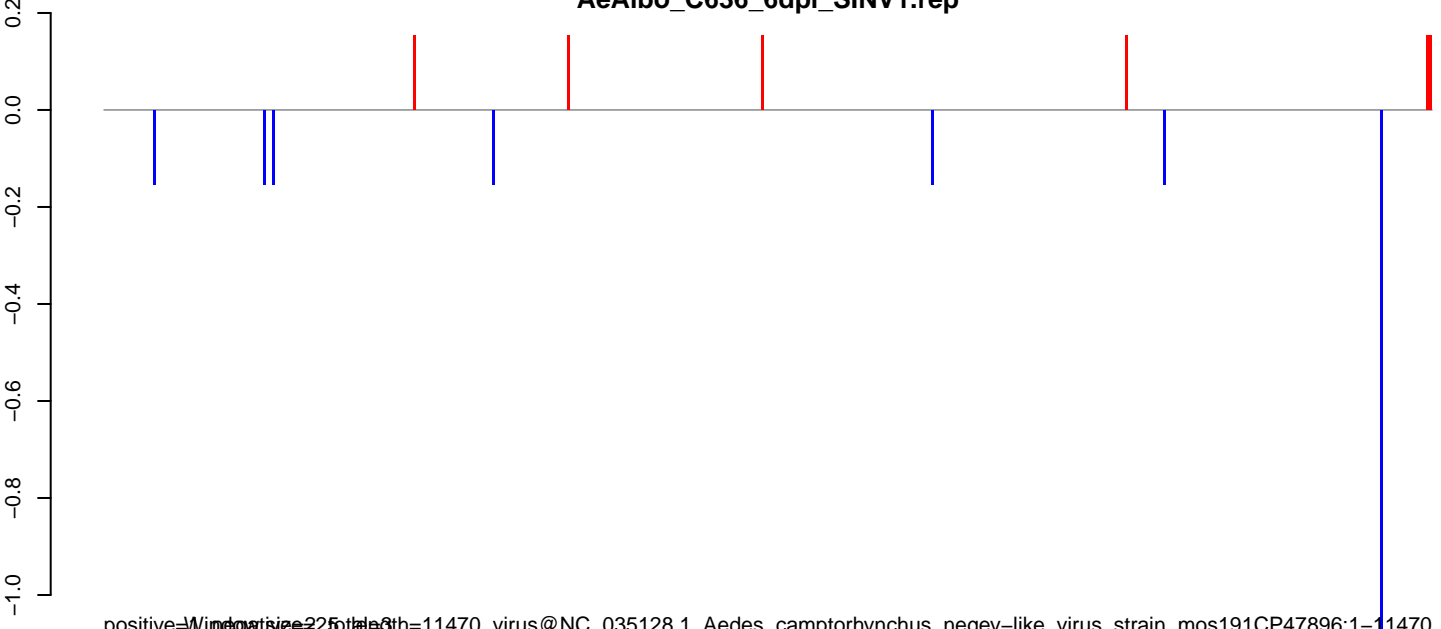
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AeAlbo_C636_6dpi_SINV1.24_35.rep



positive=0, negative=0, total=0

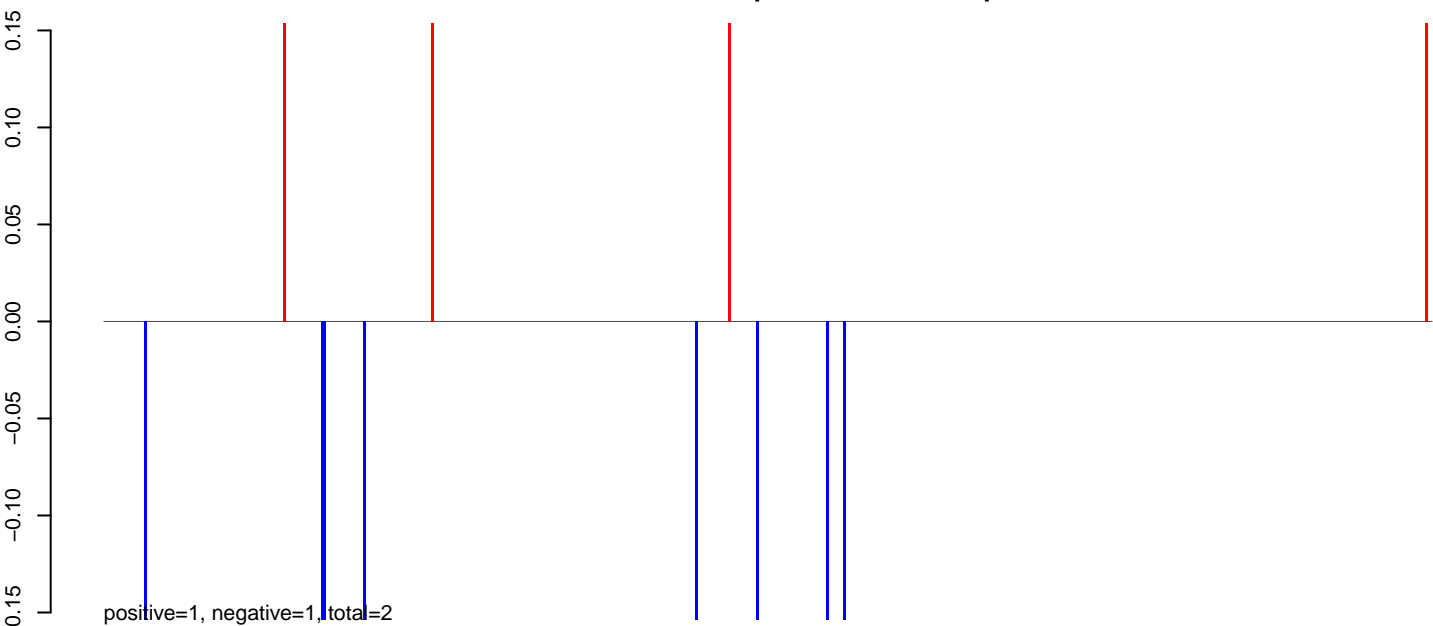
AeAlbo_C636_6dpi_SINV1.rep



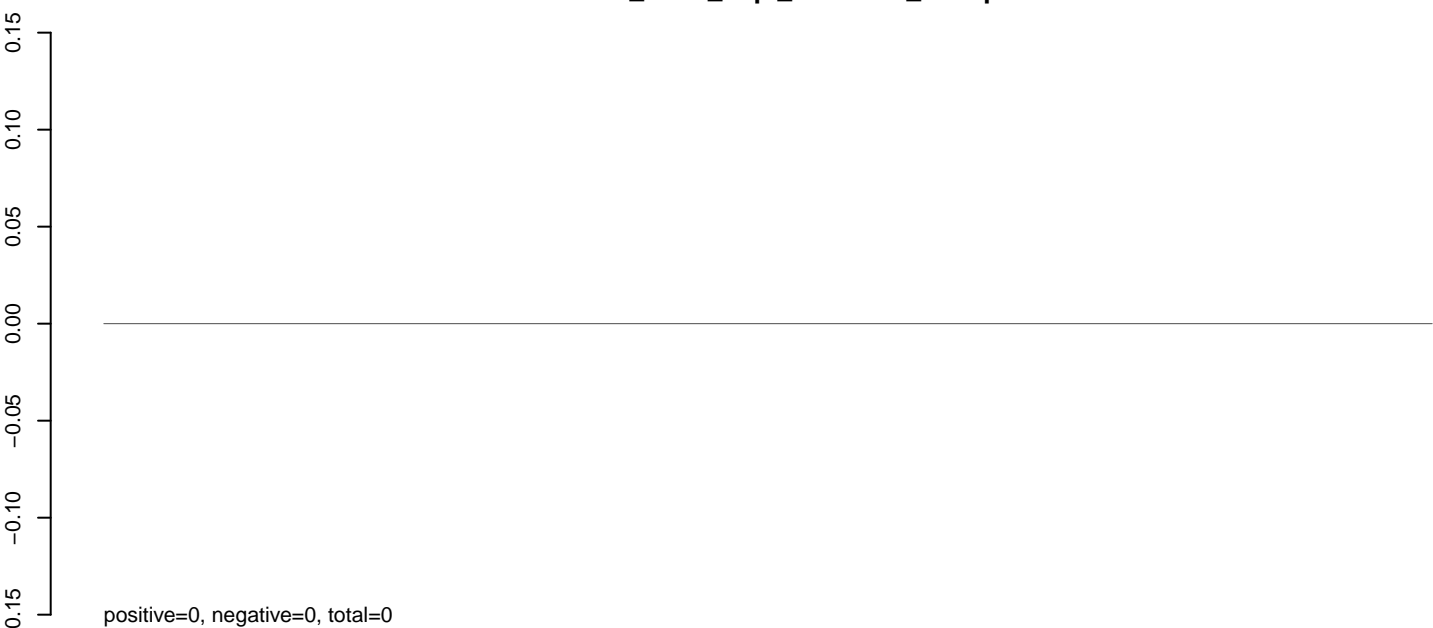
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0 2000 4000 6000 8000 10000

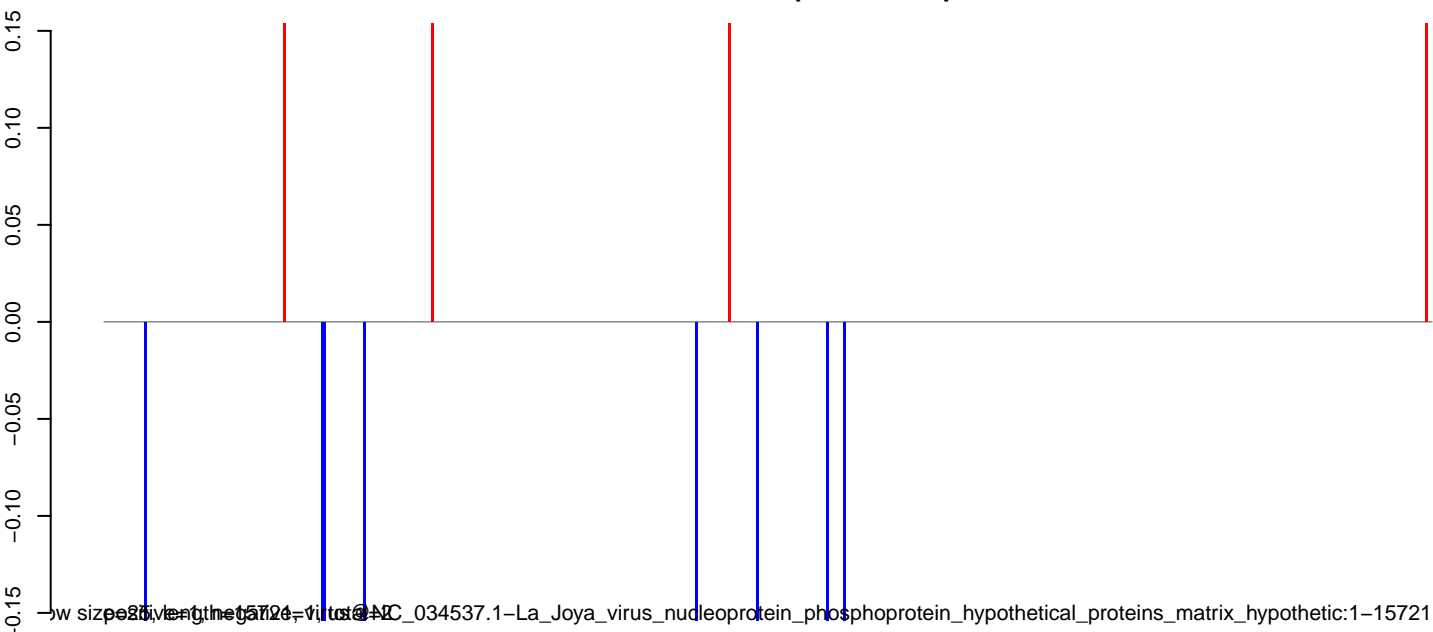
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



AeAlbo_C636_6dpi_SINV1.rep



hw size positive length 5724 -virus@AC_034537.1-La_Joya_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_hypothetic:1-15721

0 5000 10000 15000

AeAlbo_C636_6dpi_SINV1.18_23.rep

0.2
0.0
-0.2
-0.4
-0.6

positive=1, negative=2, total=3

AeAlbo_C636_6dpi_SINV1.24_35.rep

0.2
0.0
-0.2
-0.4
-0.6

positive=0, negative=0, total=0

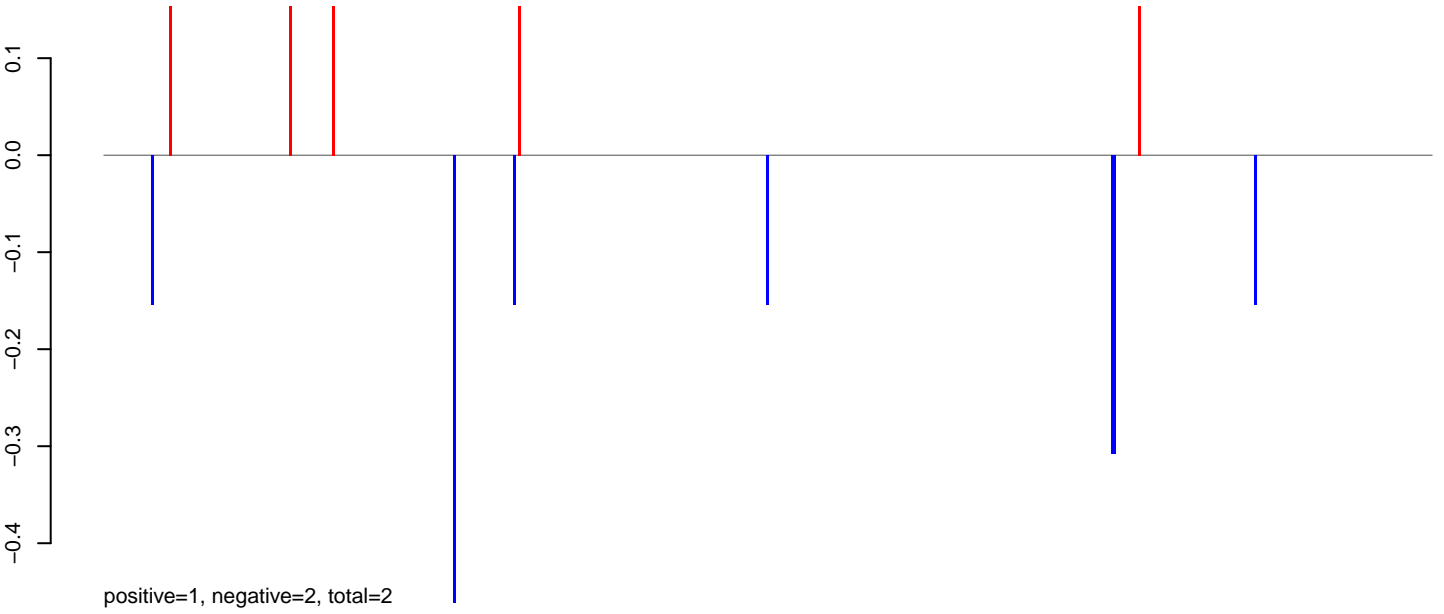
AeAlbo_C636_6dpi_SINV1.rep

0.2
0.0
-0.2
-0.4
-0.6

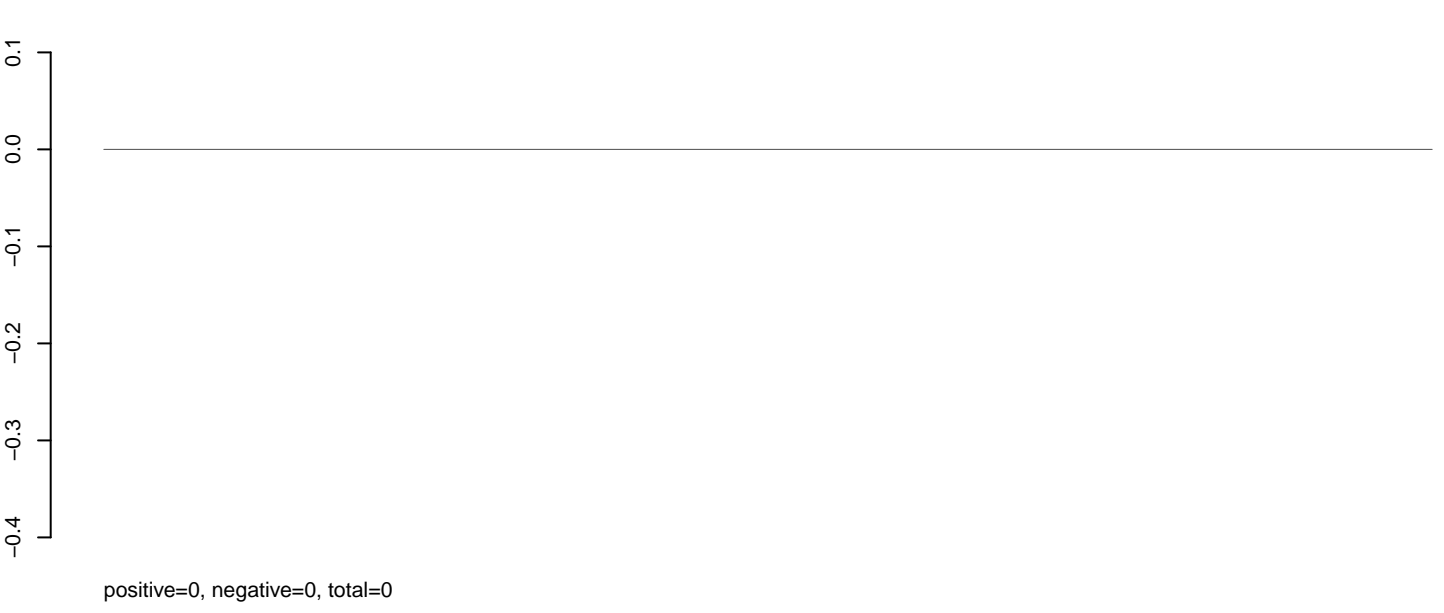
size=25, length=13593, version=2, source=C:\3034534.1-Rochambeau_virus_nucleoprotein_phosphoprotein_matrix_hypothetical_proteins_glycopr:1-13593

0 2000 4000 6000 8000 10000 12000 14000

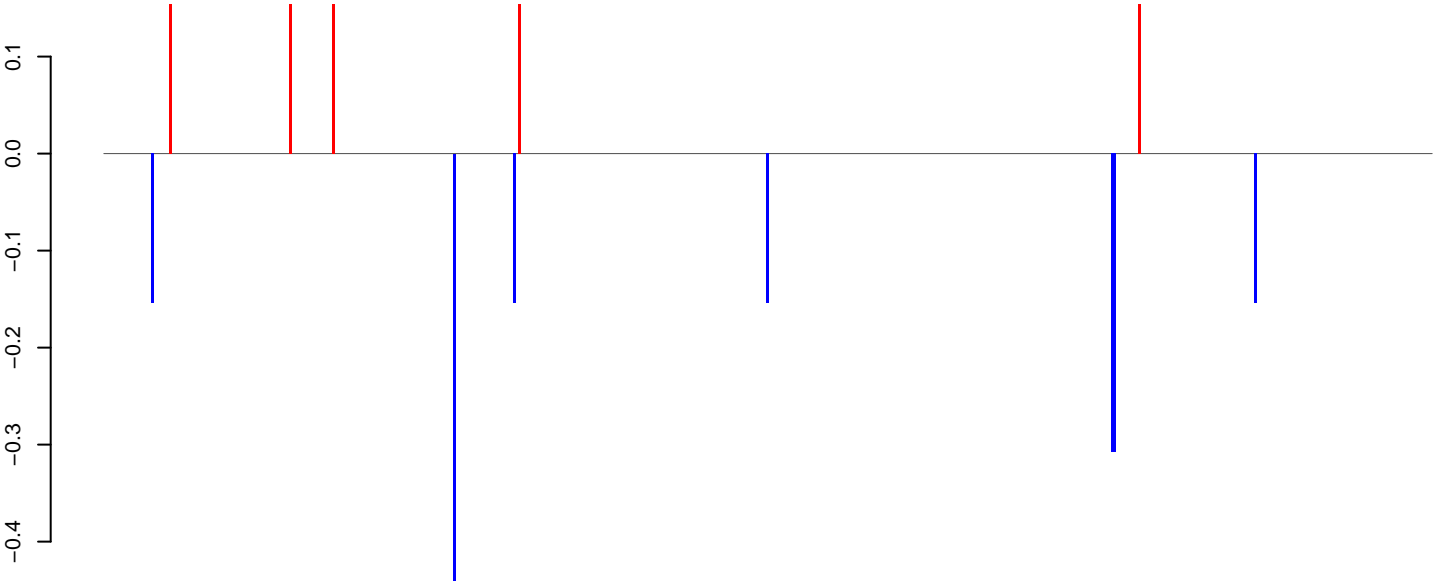
AeAlbo_C636_6dpi_SINV1.18_23.rep



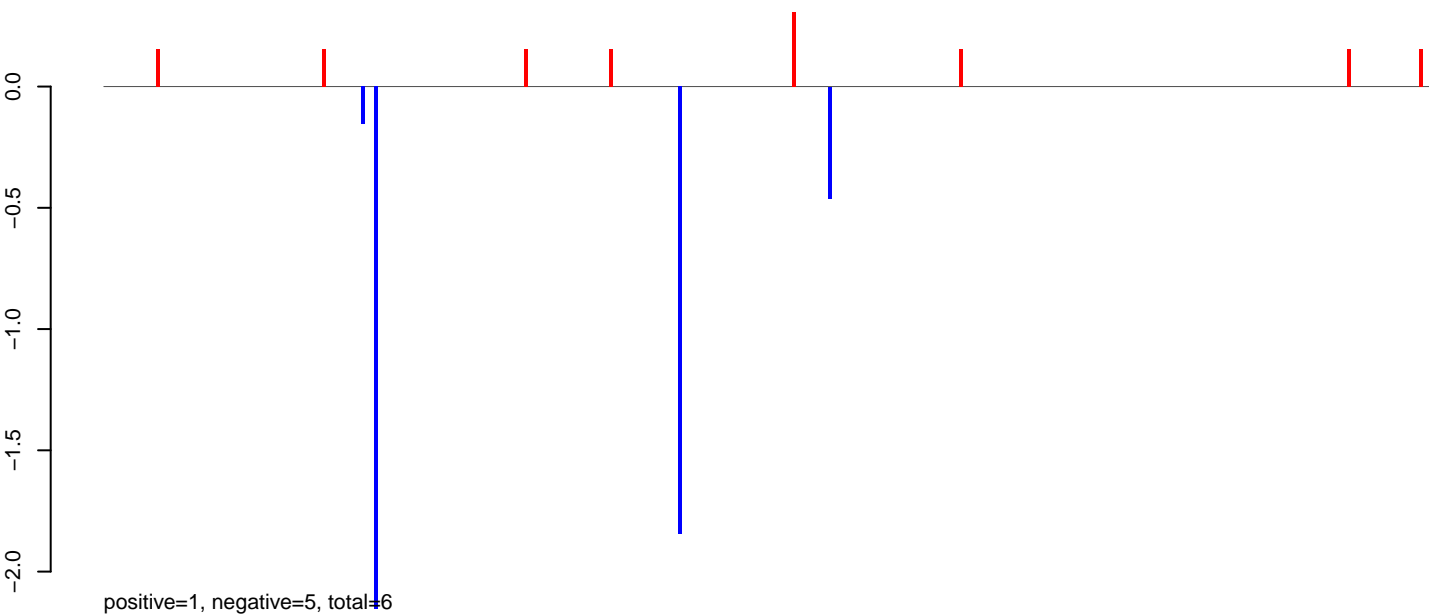
AeAlbo_C636_6dpi_SINV1.24_35.rep



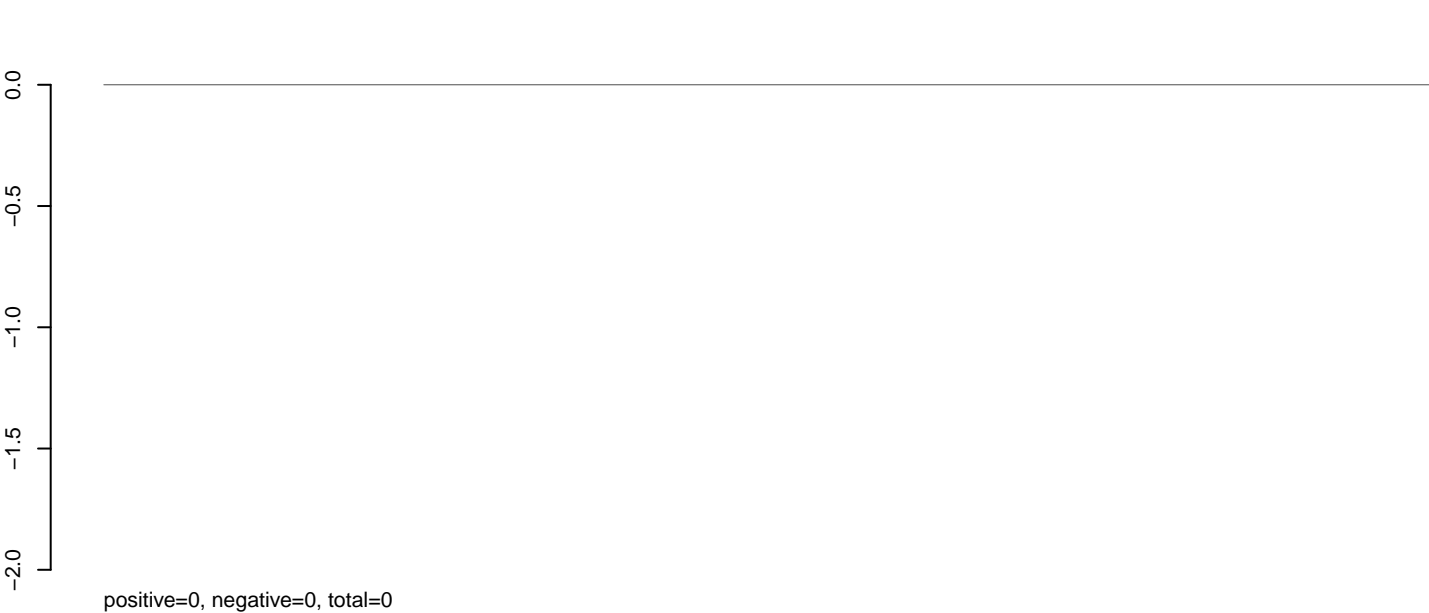
AeAlbo_C636_6dpi_SINV1.rep



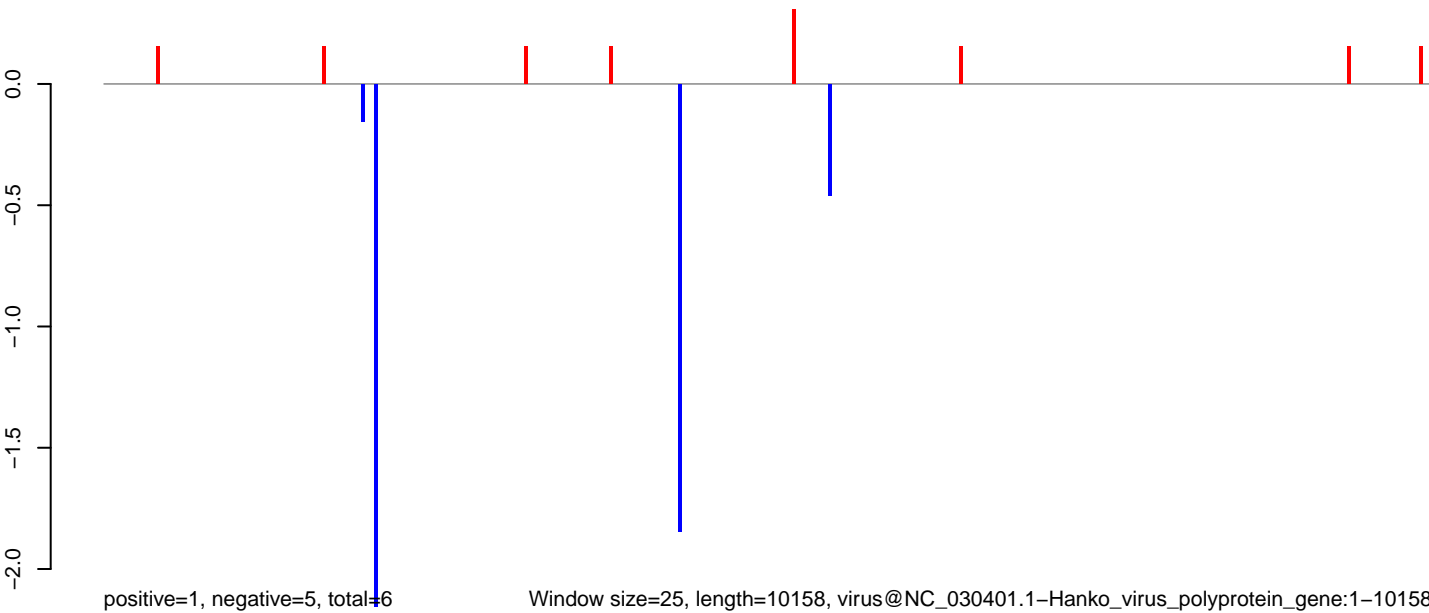
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep

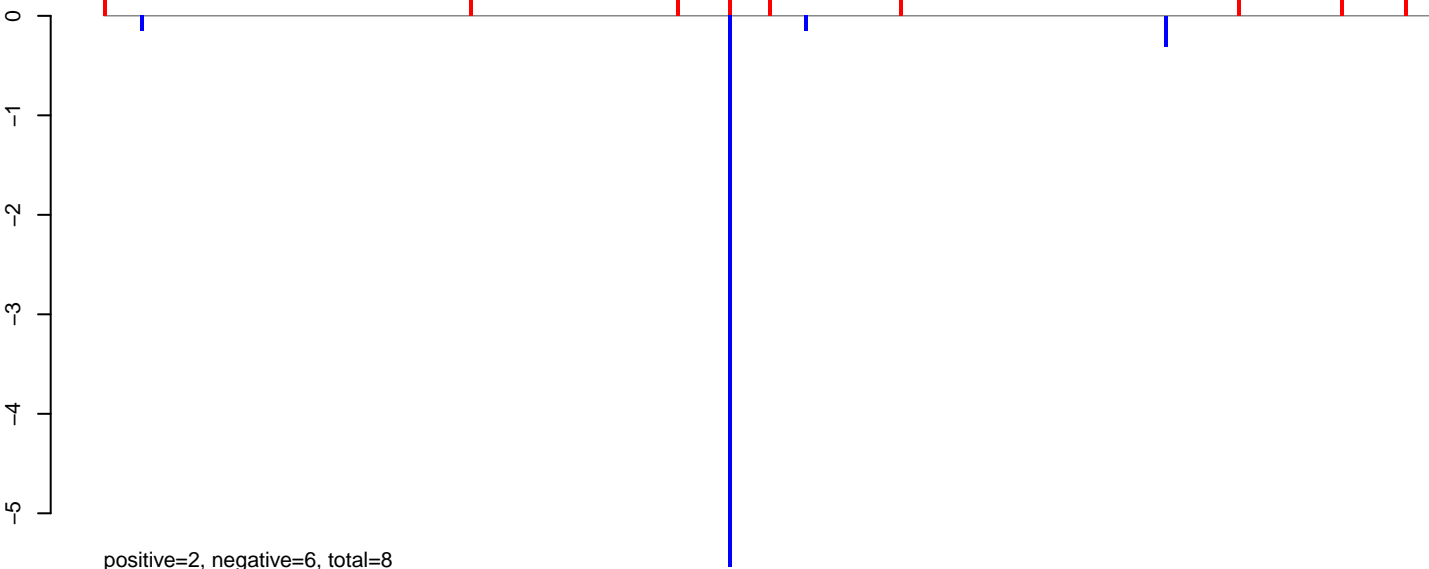


AeAlbo_C636_6dpi_SINV1.rep

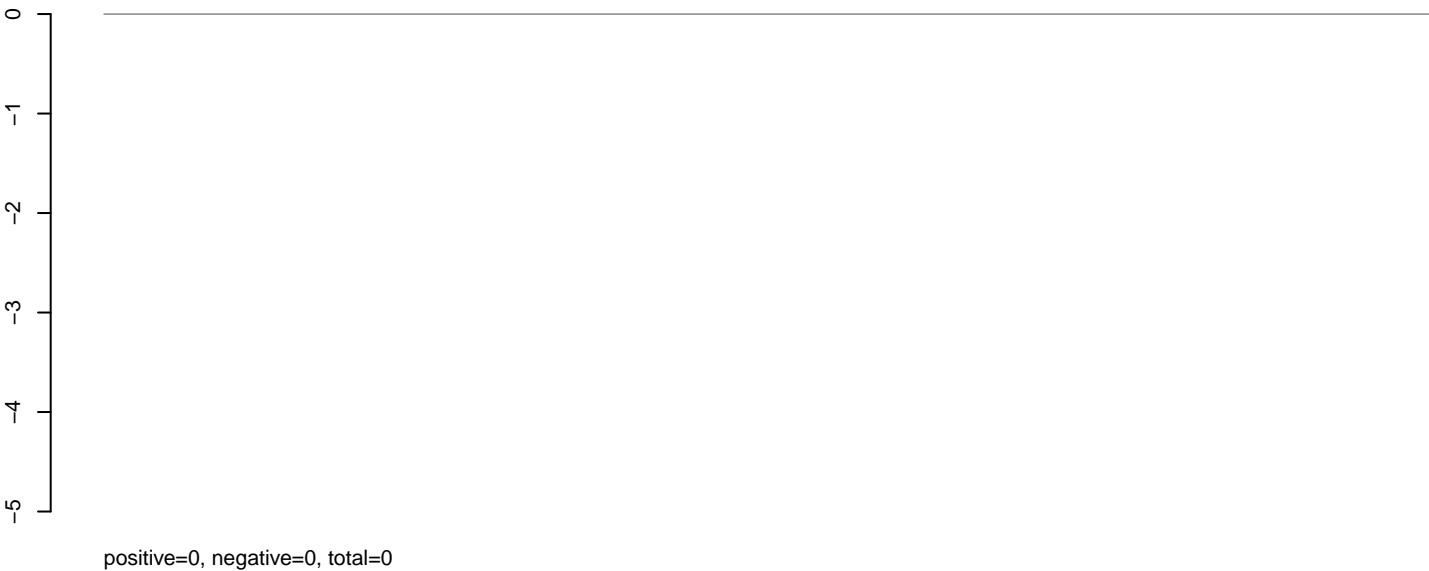


Window size=25, length=10158, virus@NC_030401.1-Hanko_virus_polyprotein_gene:1-10158

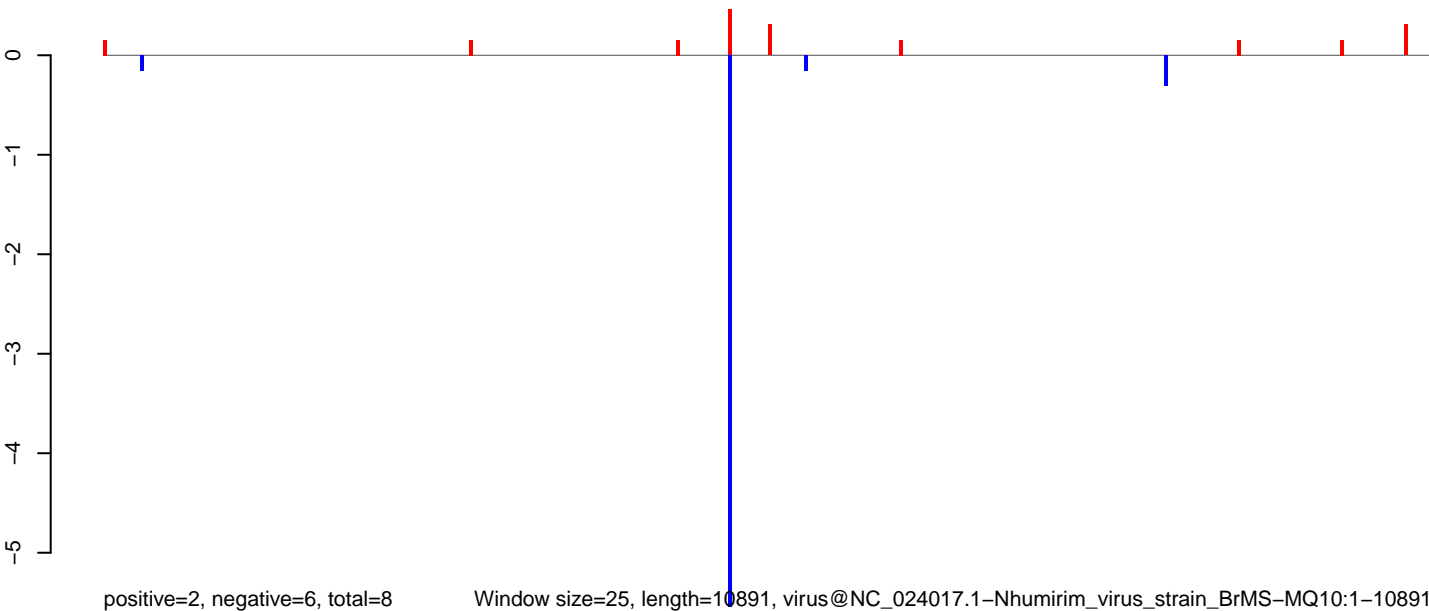
AeAlbo_C636_6dpi_SINV1.18_23.rep



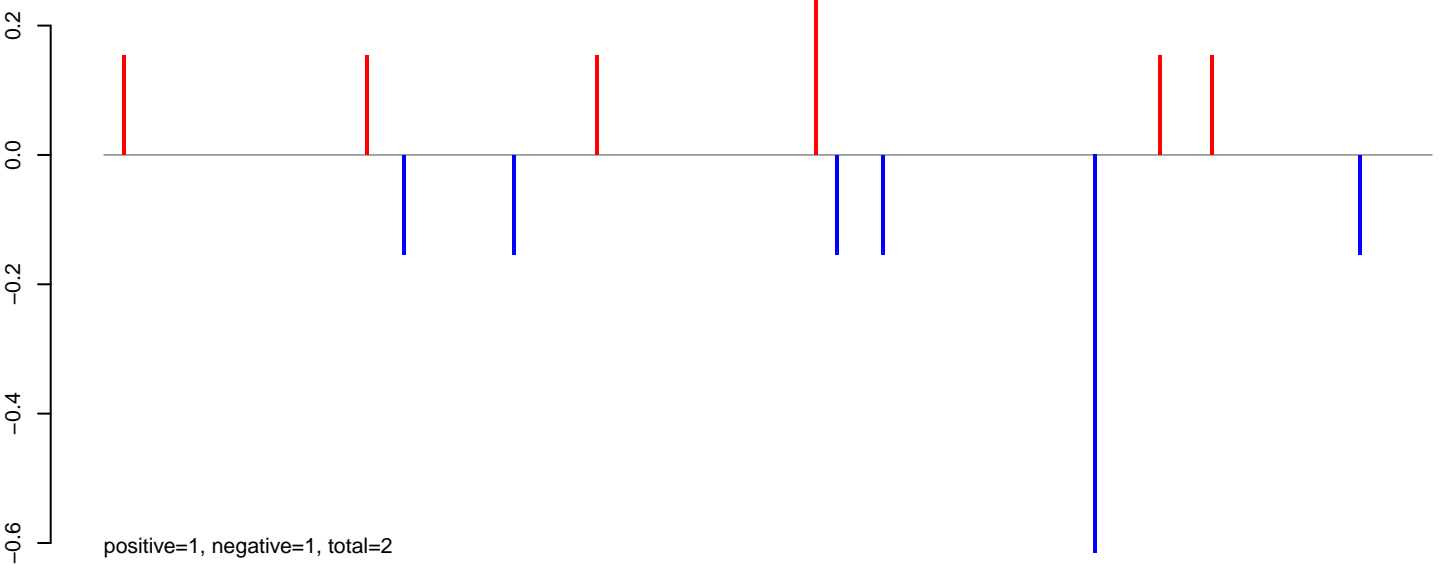
AeAlbo_C636_6dpi_SINV1.24_35.rep



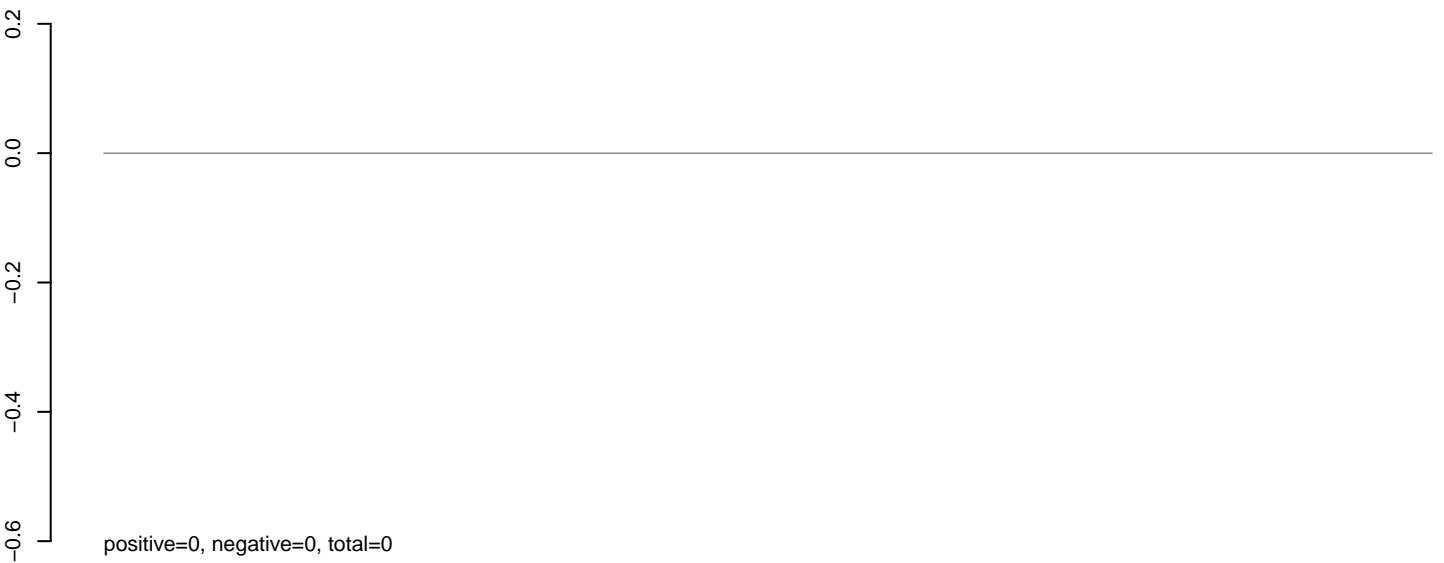
AeAlbo_C636_6dpi_SINV1.rep



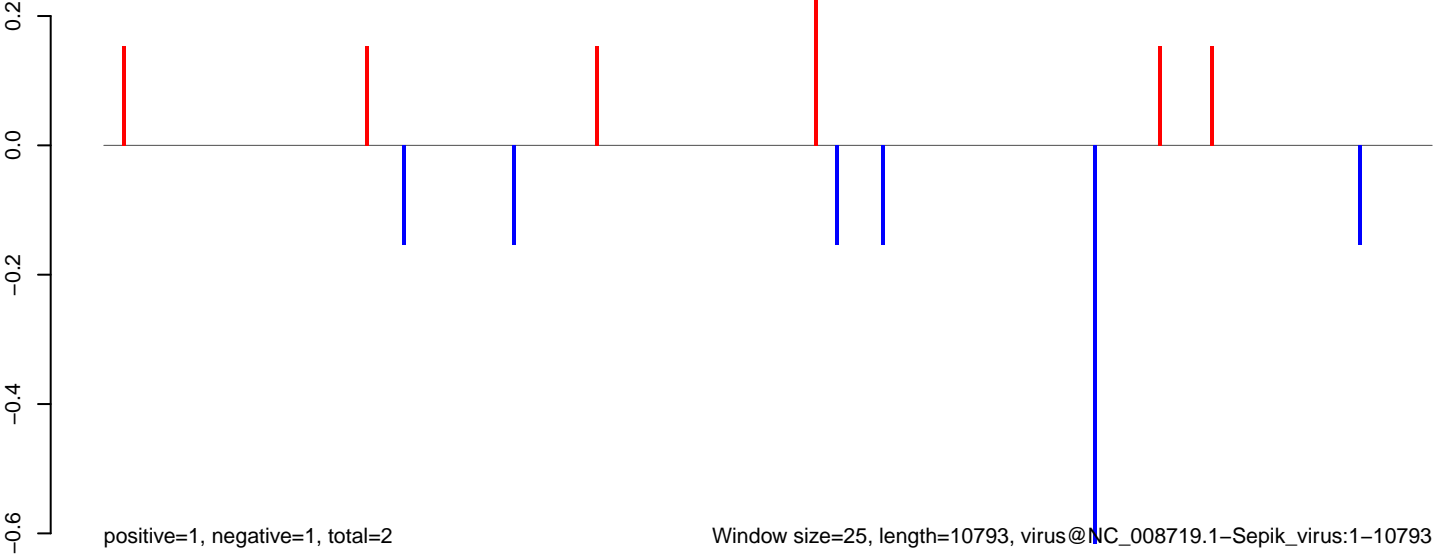
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



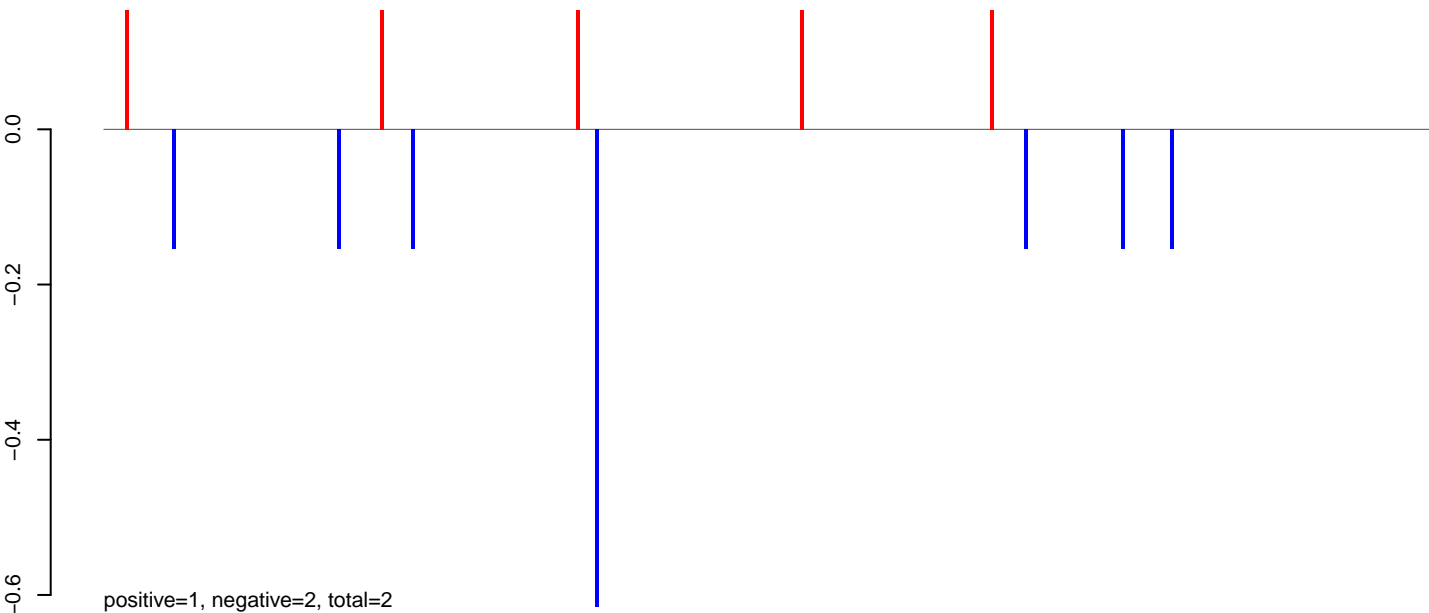
AeAlbo_C636_6dpi_SINV1.rep



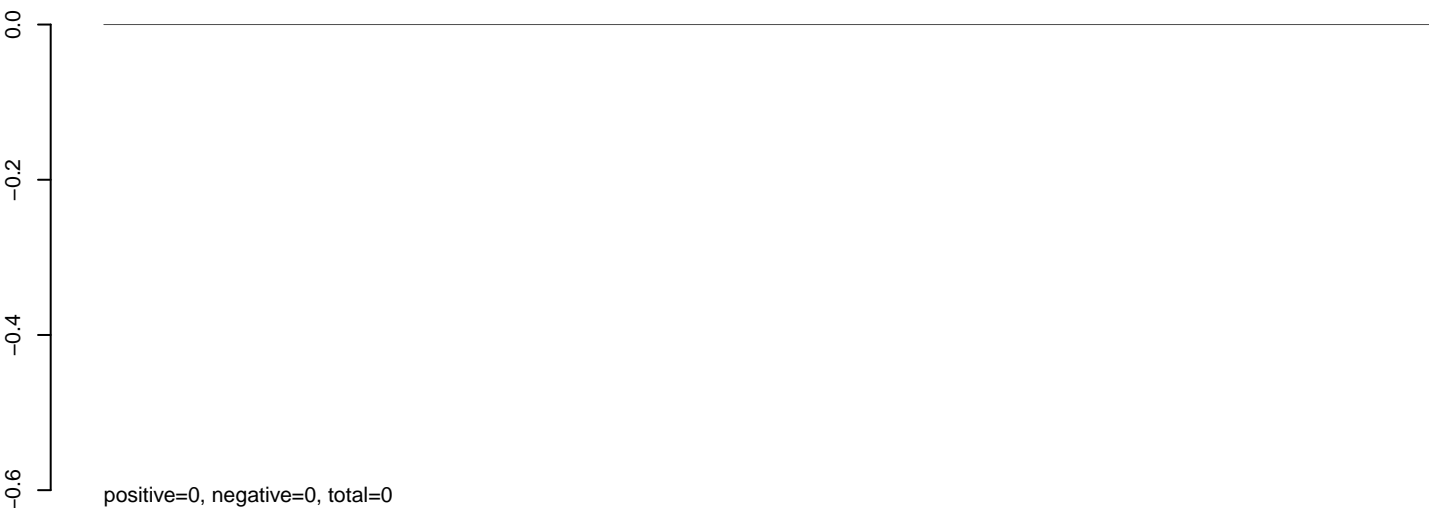
Window size=25, length=10793, virus@NC_008719.1-Sepik_virus:1-10793

0 2000 4000 6000 8000 10000

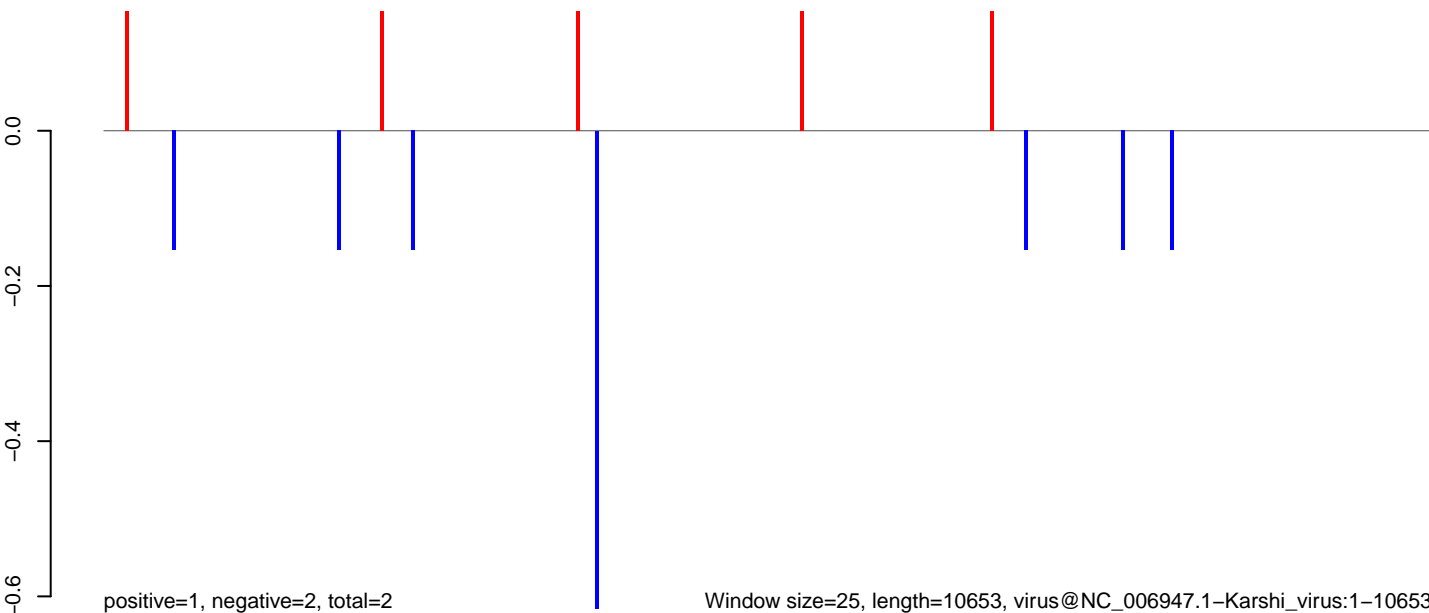
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



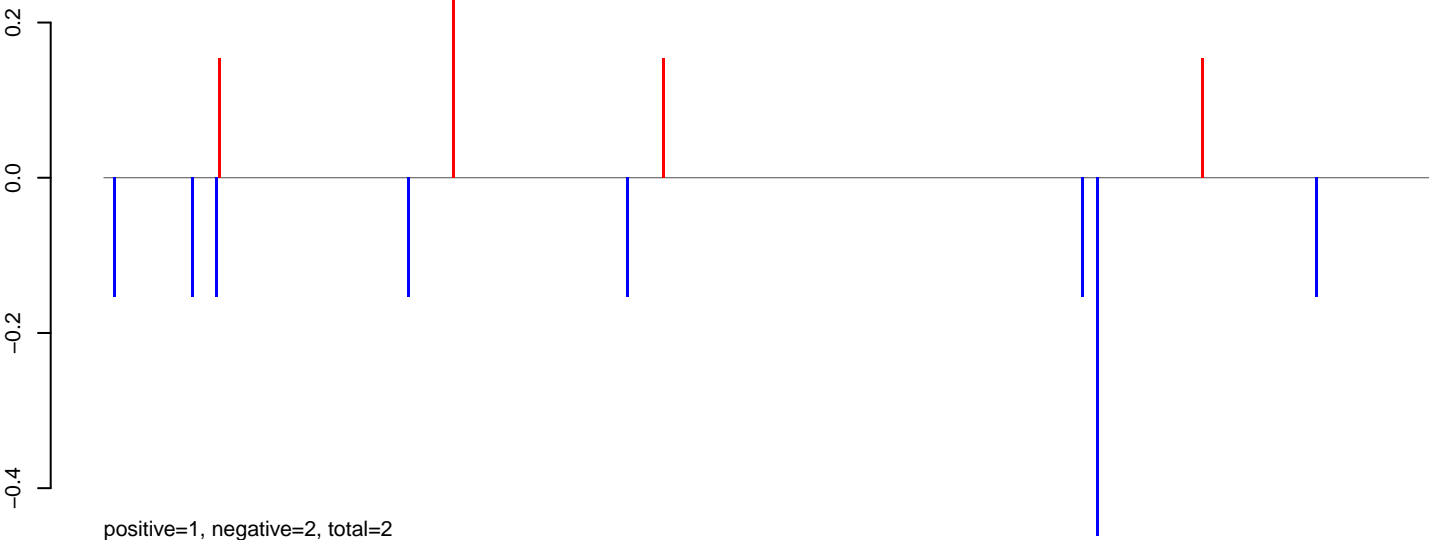
AeAlbo_C636_6dpi_SINV1.rep



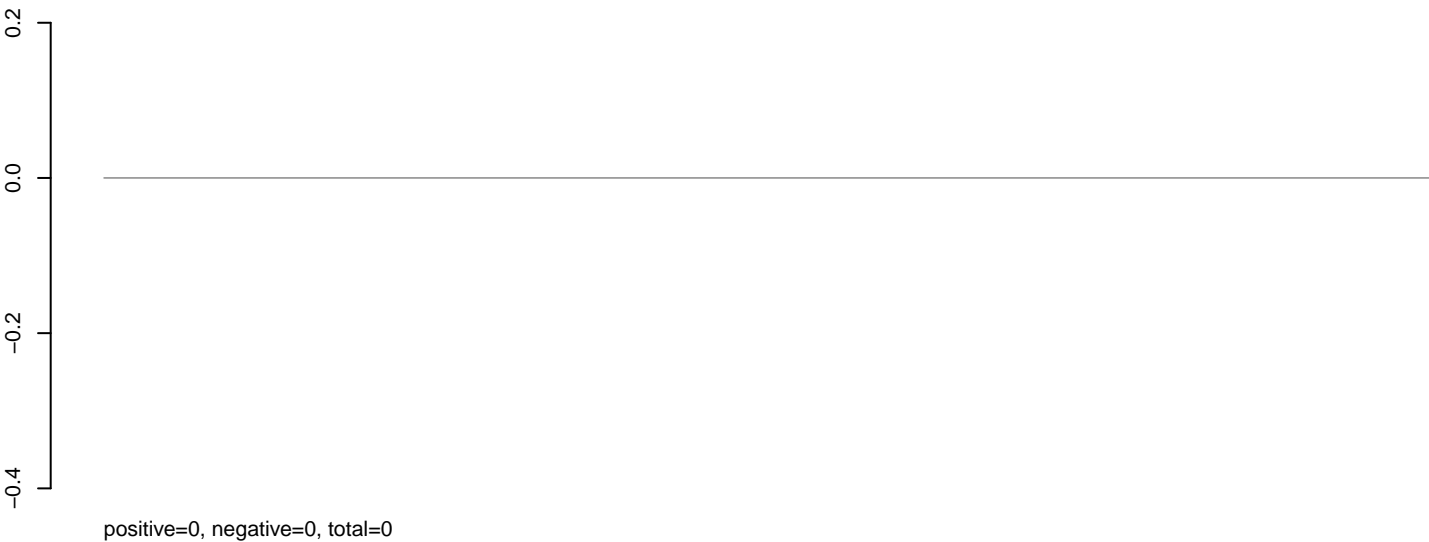
Window size=25, length=10653, virus@NC_006947.1-Karshi_virus:1-10653

0 2000 4000 6000 8000 10000

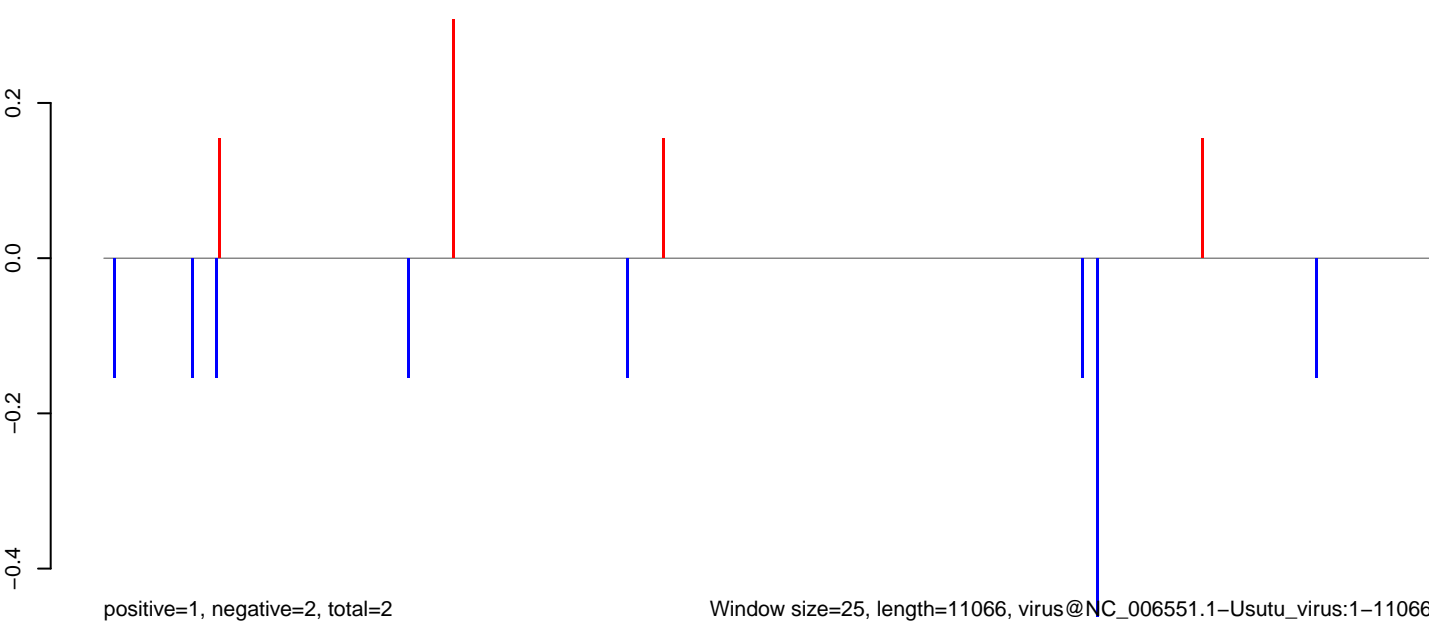
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep

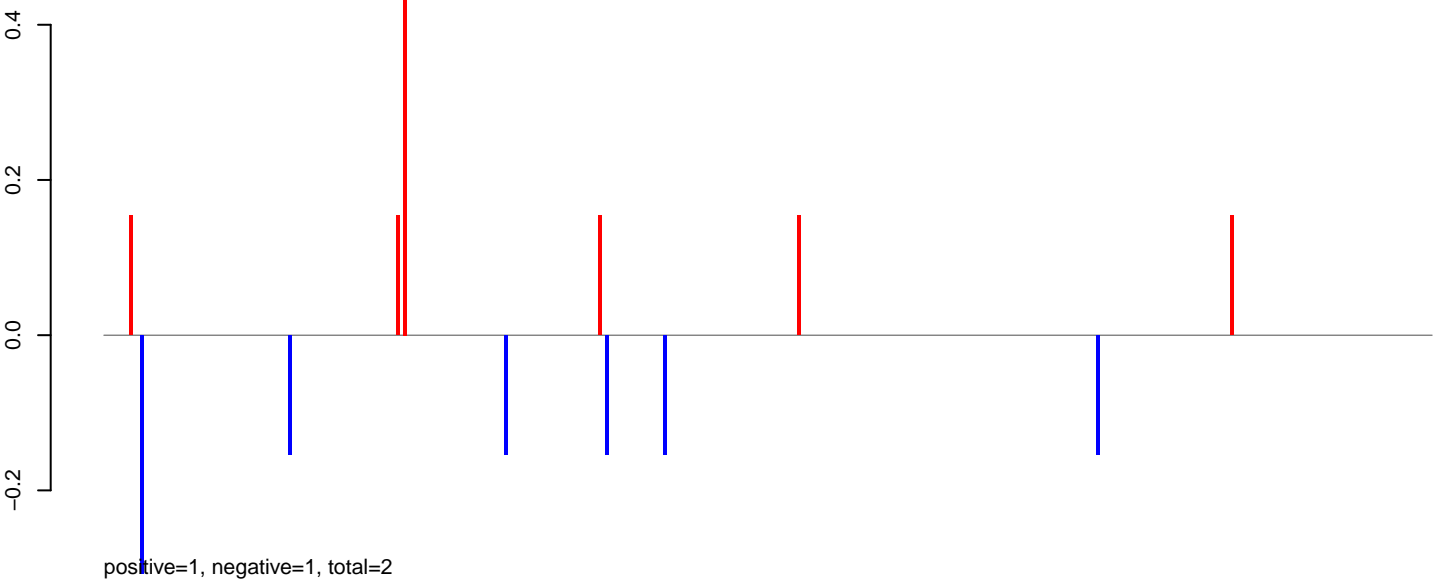


AeAlbo_C636_6dpi_SINV1.rep



Window size=25, length=11066, virus@NC_006551.1-Uсутu_virus:1-11066

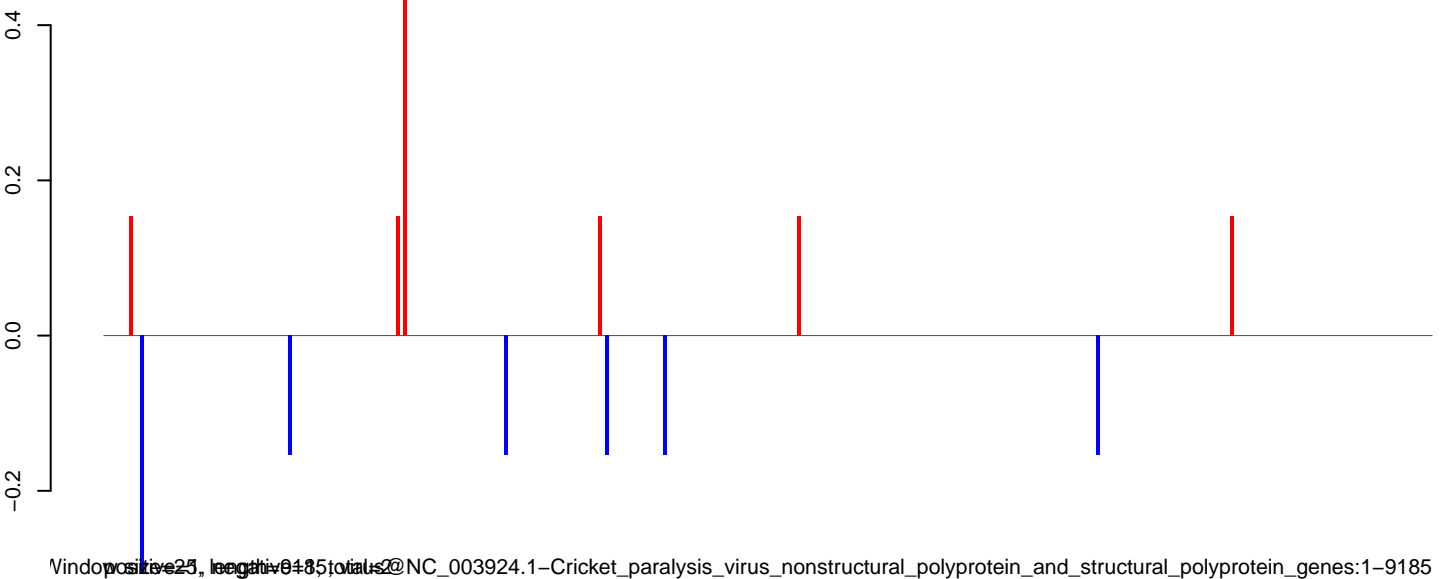
AeAlbo_C636_6dpi_SINV1.18_23.rep



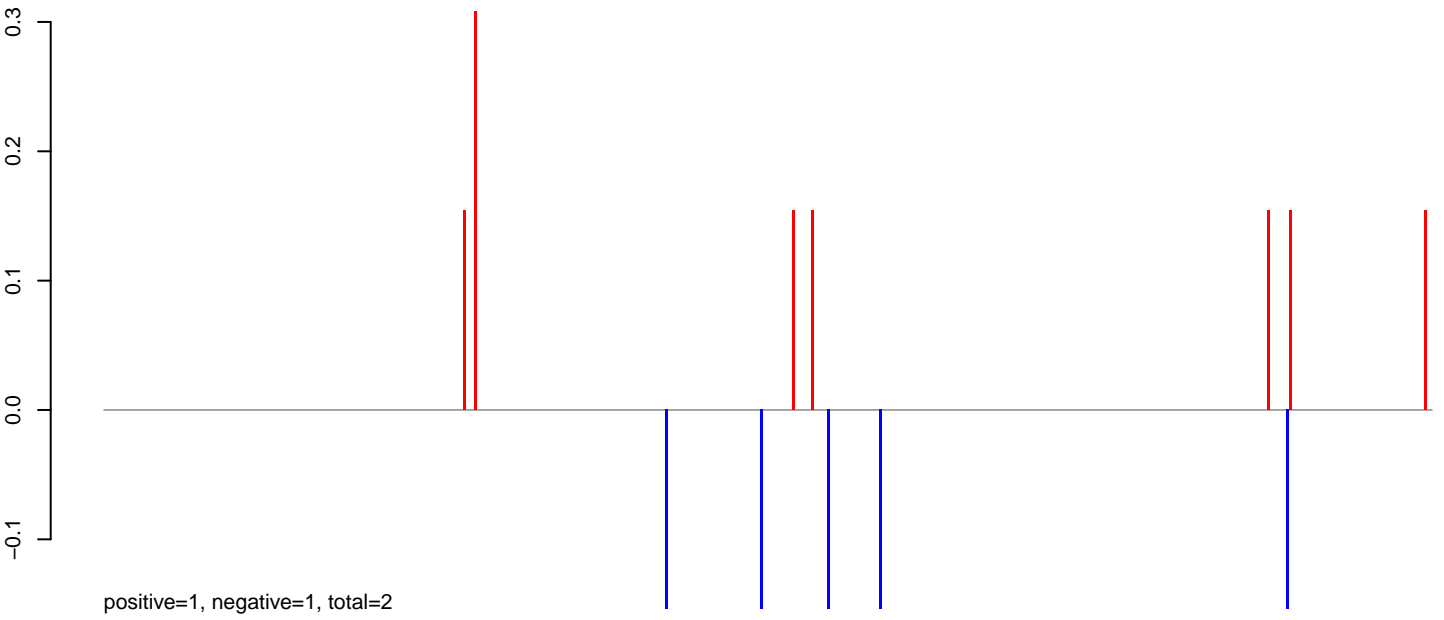
AeAlbo_C636_6dpi_SINV1.24_35.rep



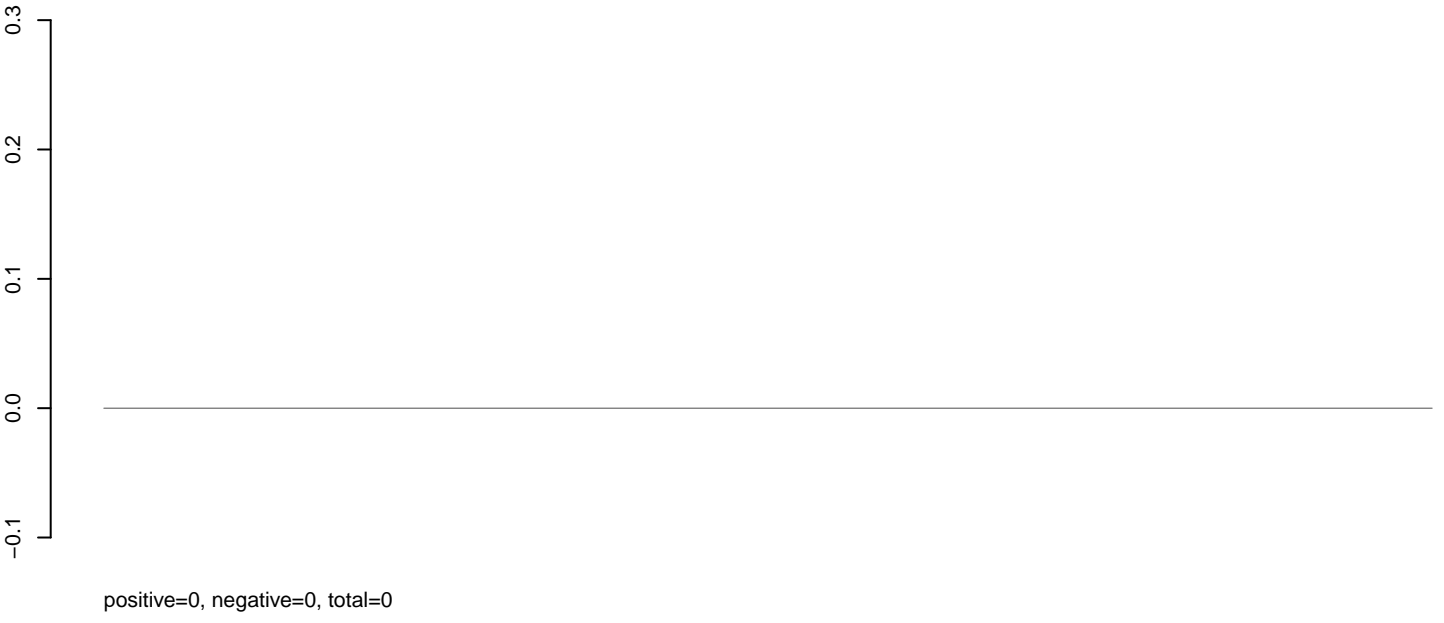
AeAlbo_C636_6dpi_SINV1.rep



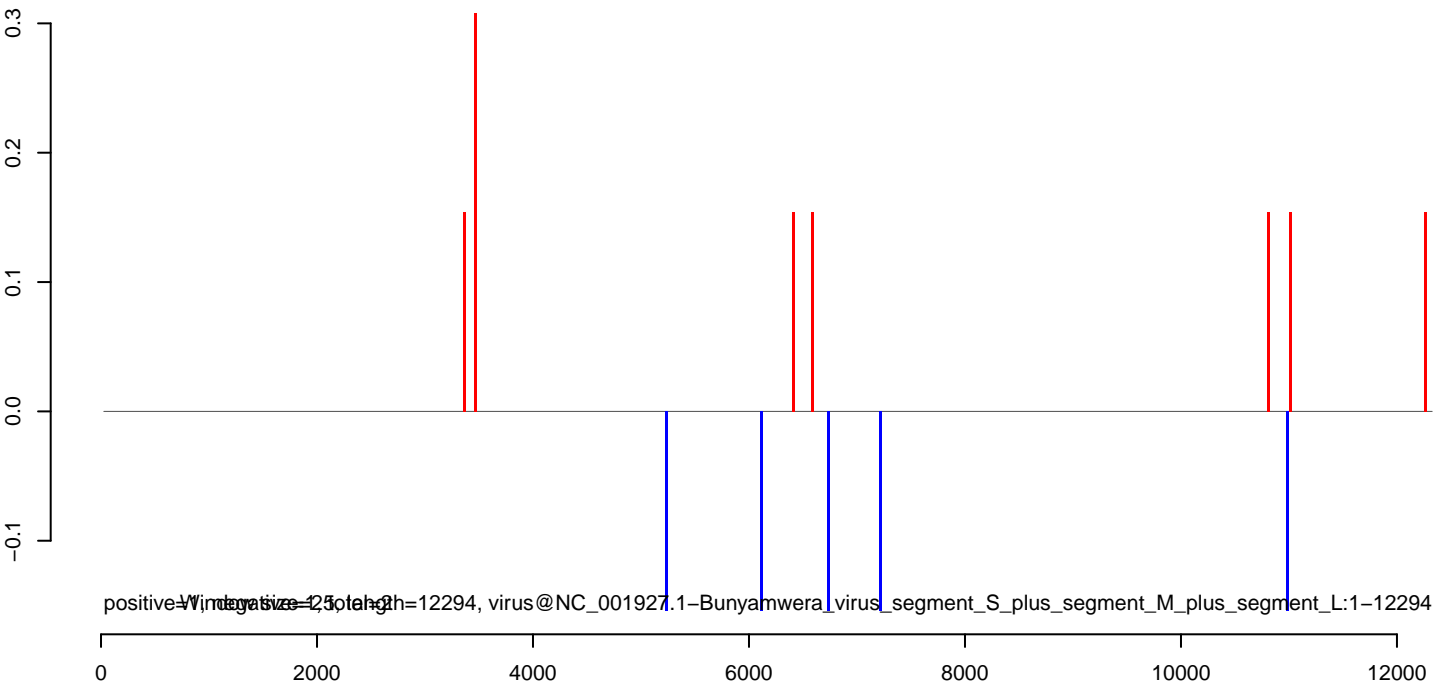
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep

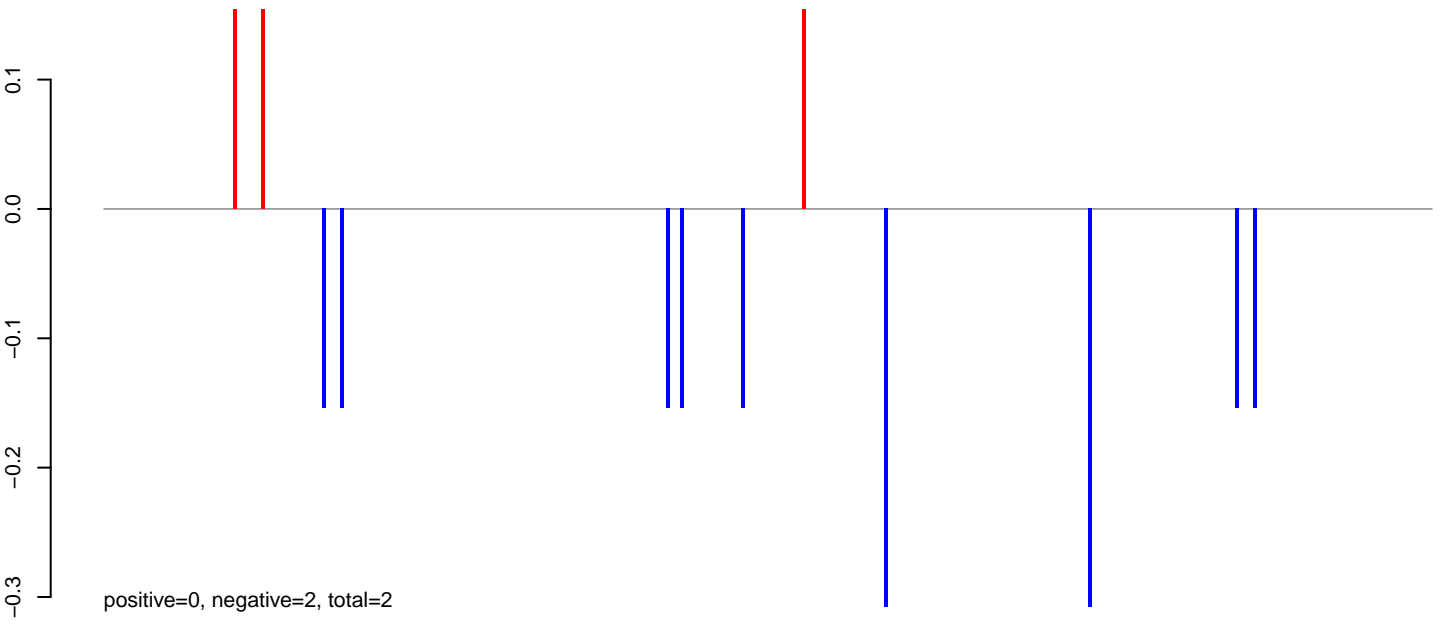


AeAlbo_C636_6dpi_SINV1.rep

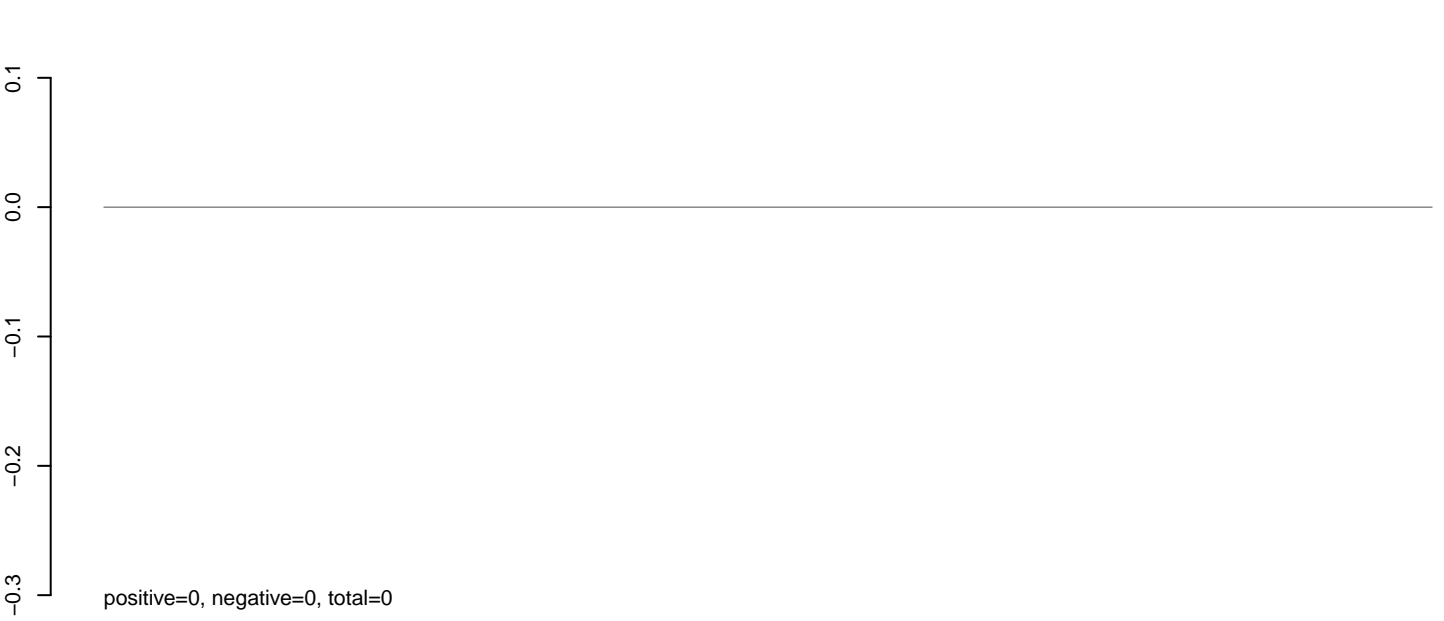


positive=1, negative=1, total=2
Window size=25, step=1, length=12294, virus@NC_001927.1-Bunyamwera_virus_segment_S_plus_segment_M_plus_segment_L:1-12294

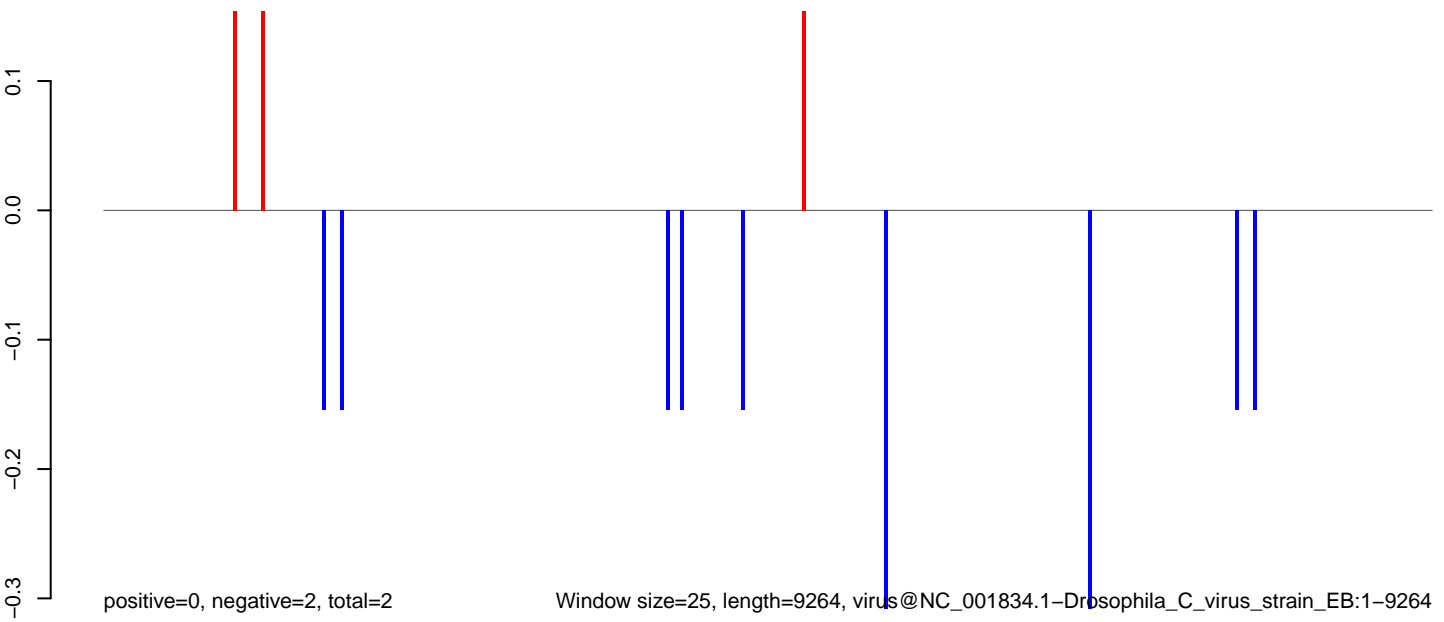
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



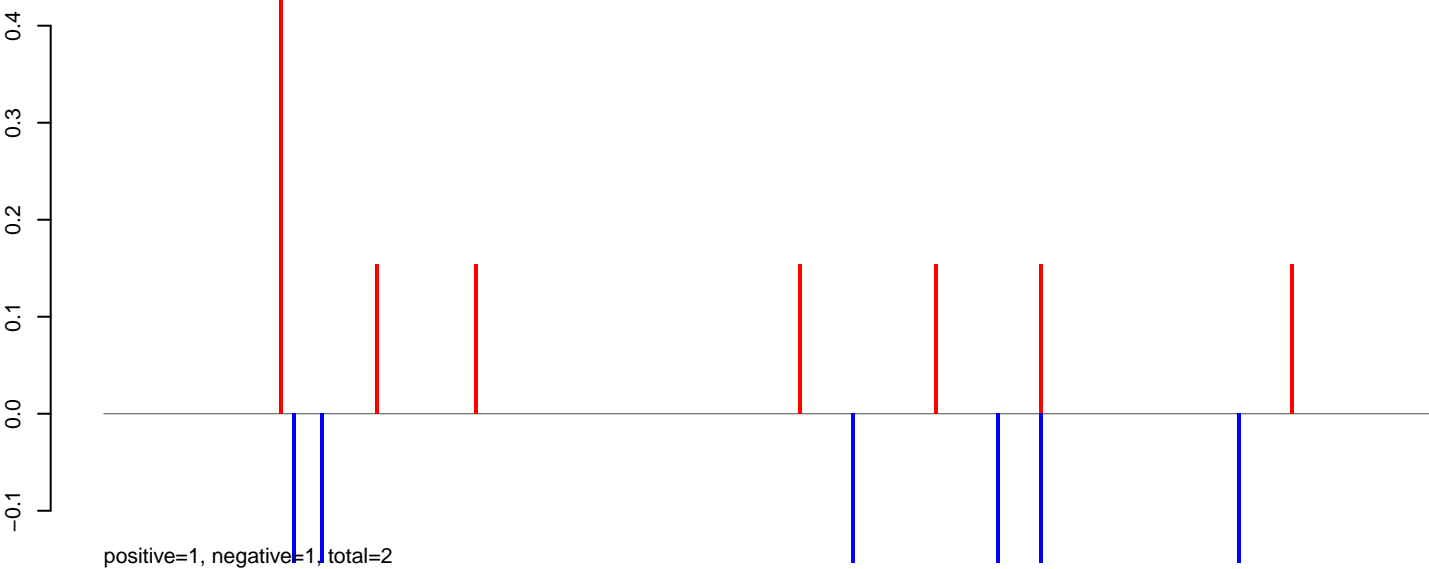
AeAlbo_C636_6dpi_SINV1.rep



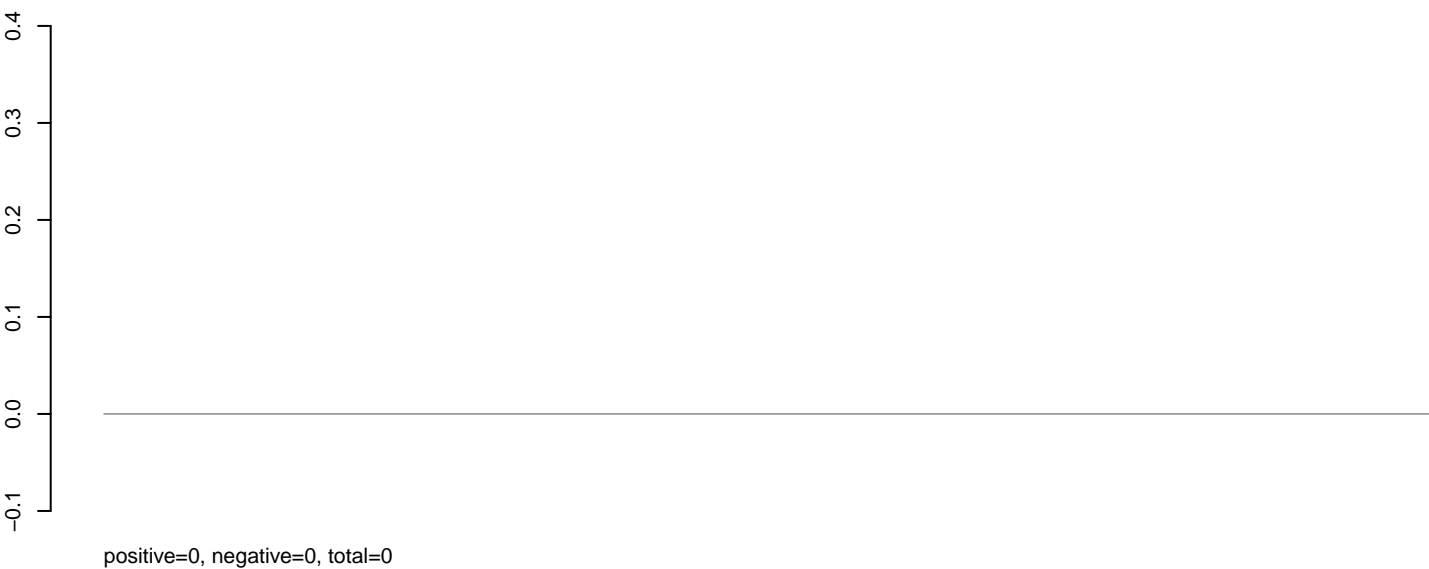
Window size=25, length=9264, virus@NC_001834.1-Drosophila_C_virus_strain_EB:1-9264

0 2000 4000 6000 8000

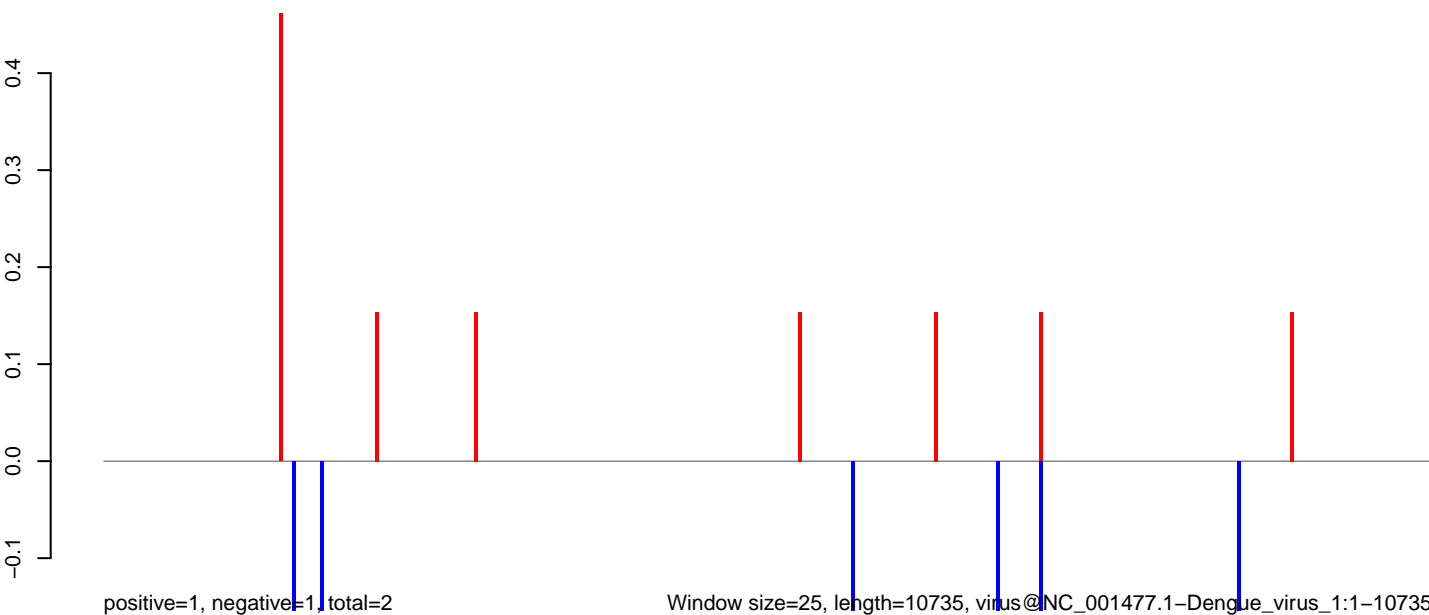
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



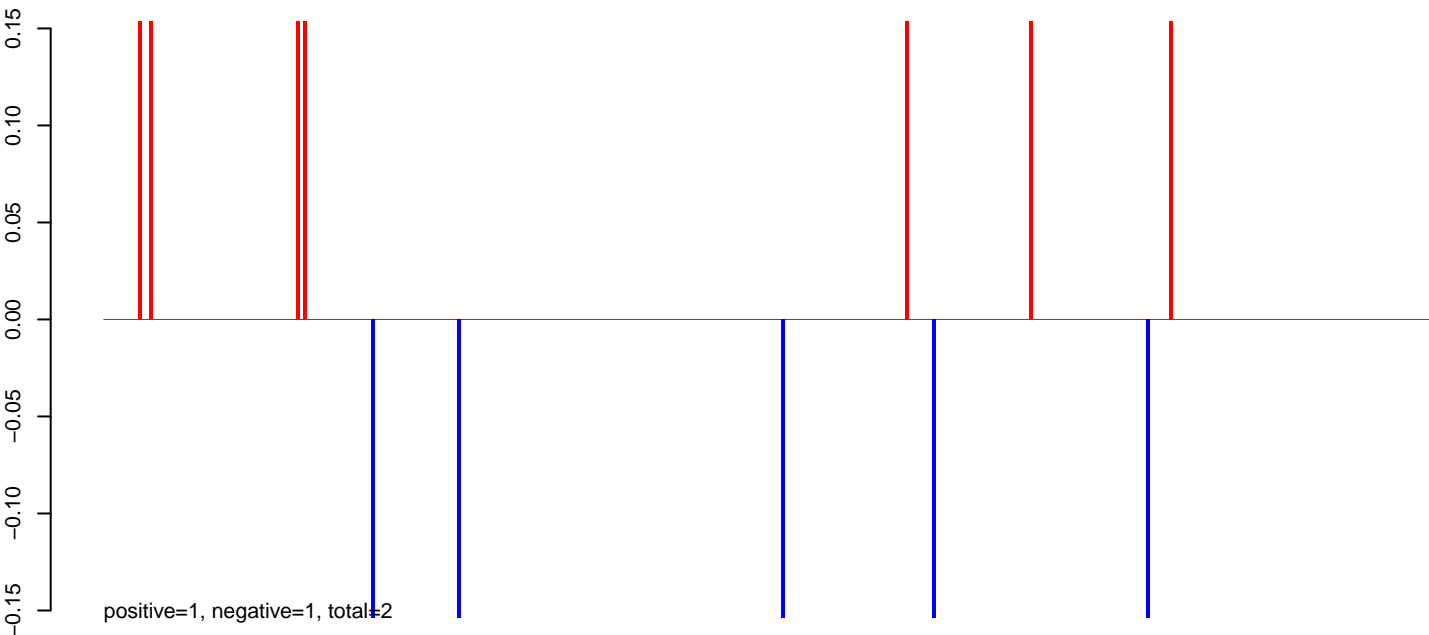
AeAlbo_C636_6dpi_SINV1.rep



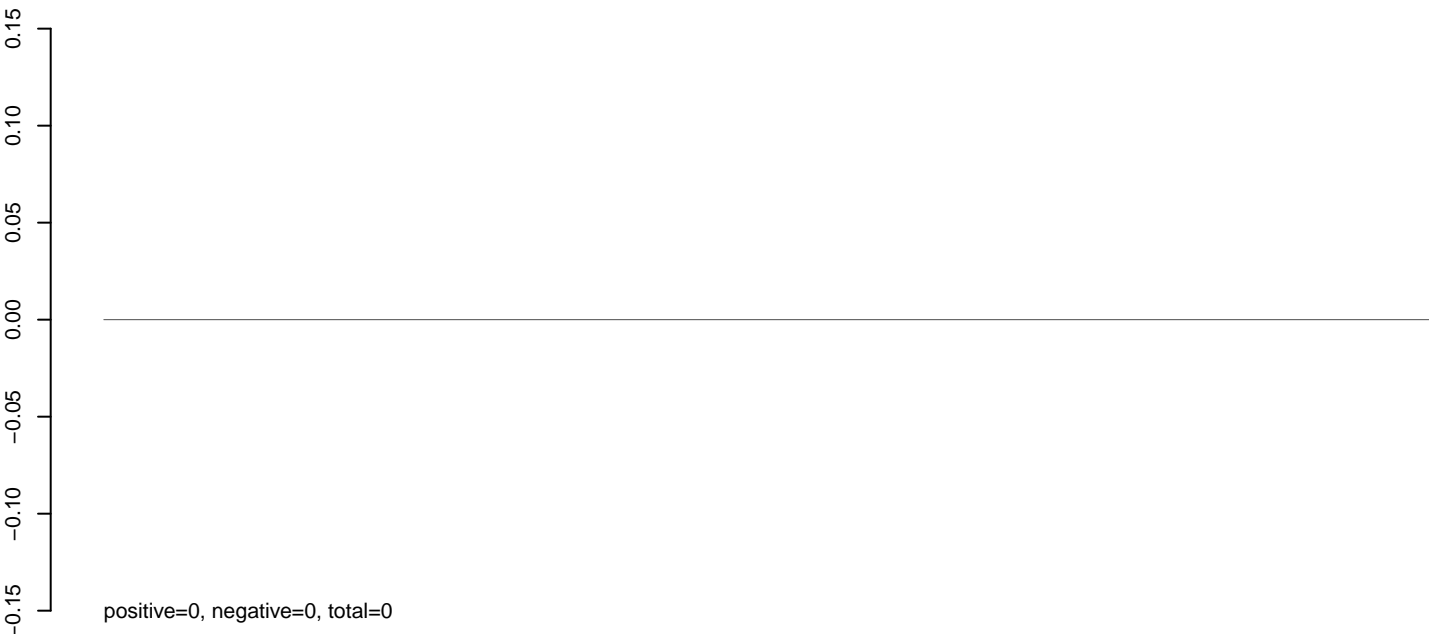
Window size=25, length=10735, virus@NC_001477.1-Dengue_virus_1:1-10735

0 2000 4000 6000 8000 10000

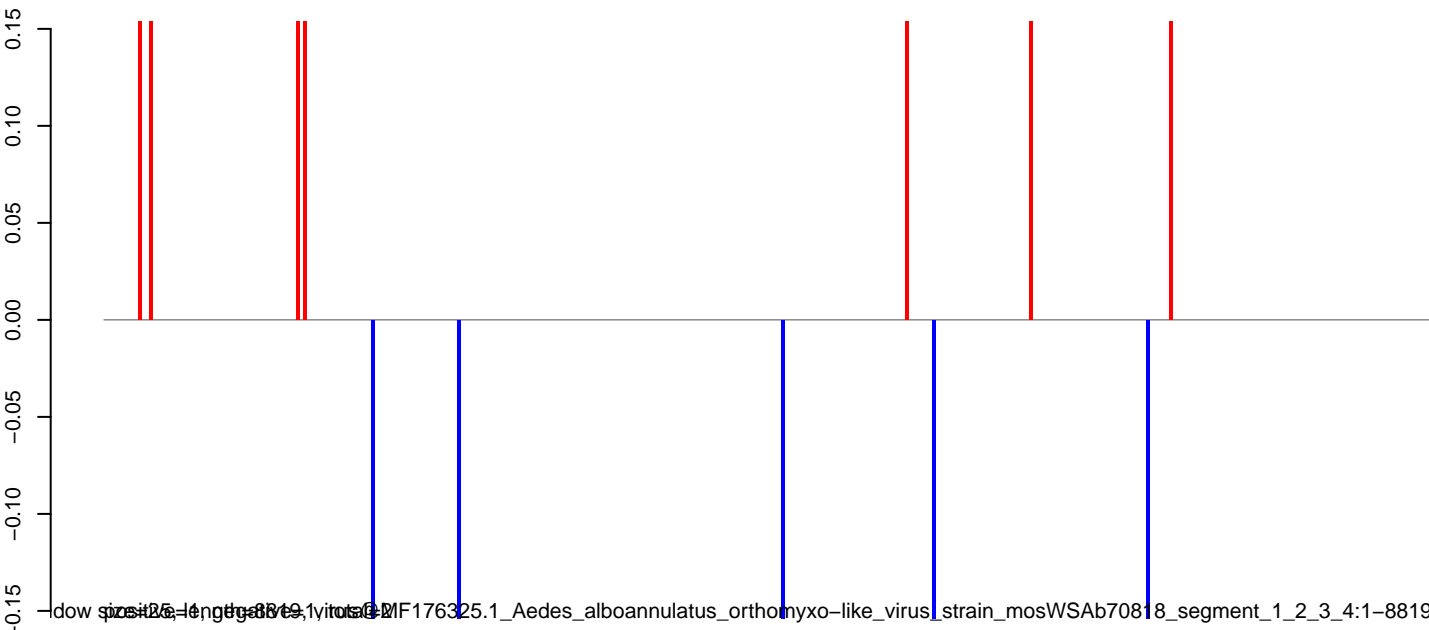
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



AeAlbo_C636_6dpi_SINV1.rep



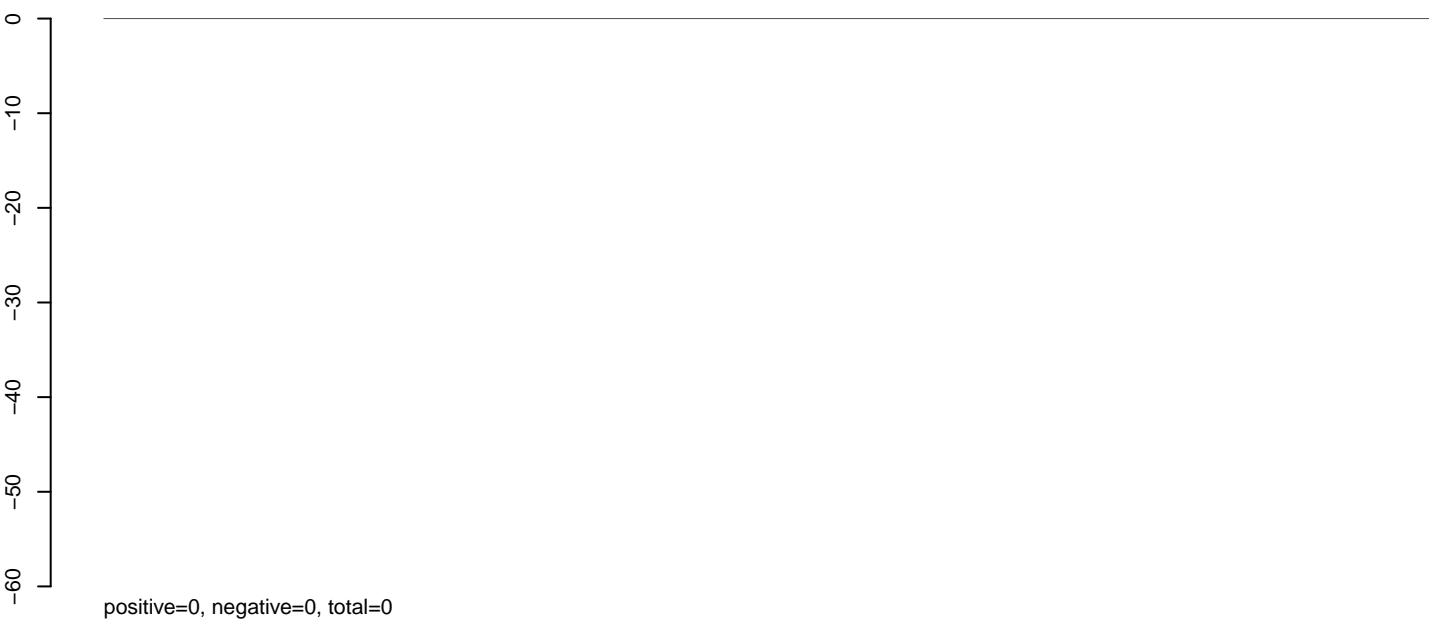
downloaded from https://www.ncbi.nlm.nih.gov/trace/blast/blast.cgi?db=GenBank&term=AF176325.1_Aedes_alboannulatus_orthomyxo-like_virus_strain_mosWSAb70818_segment_1_2_3_4:1-8819

0 2000 4000 6000 8000

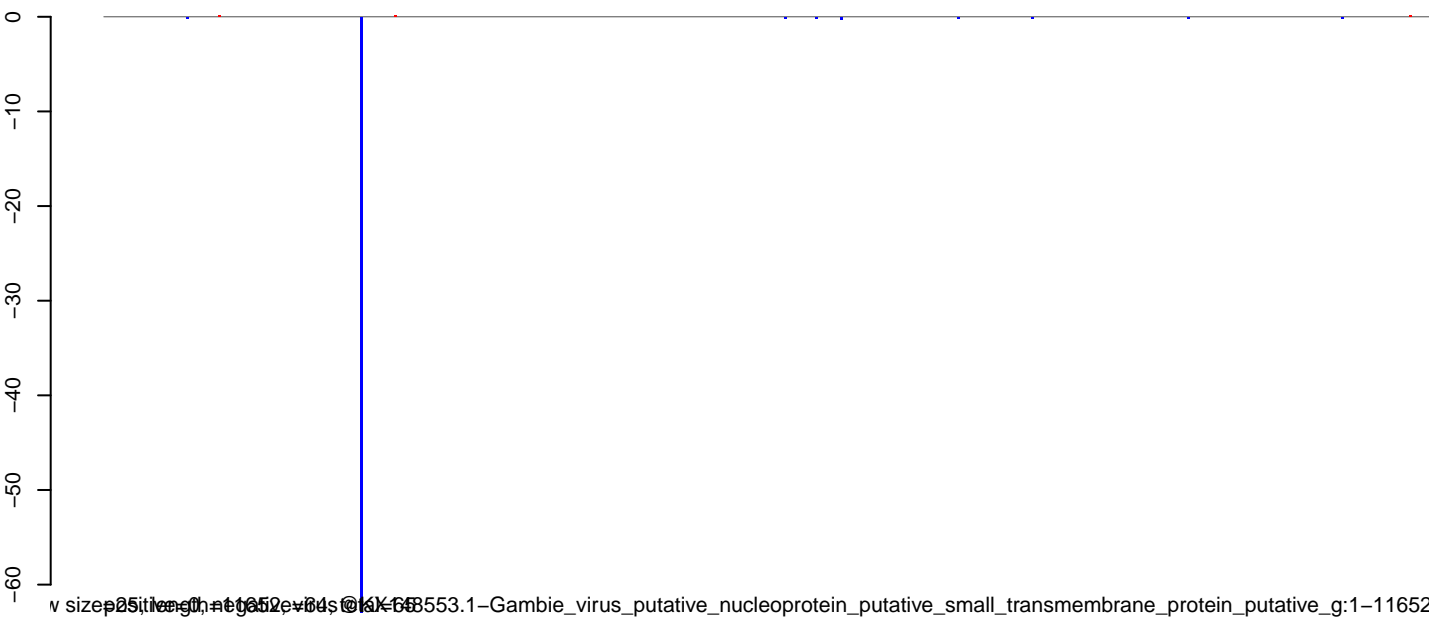
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



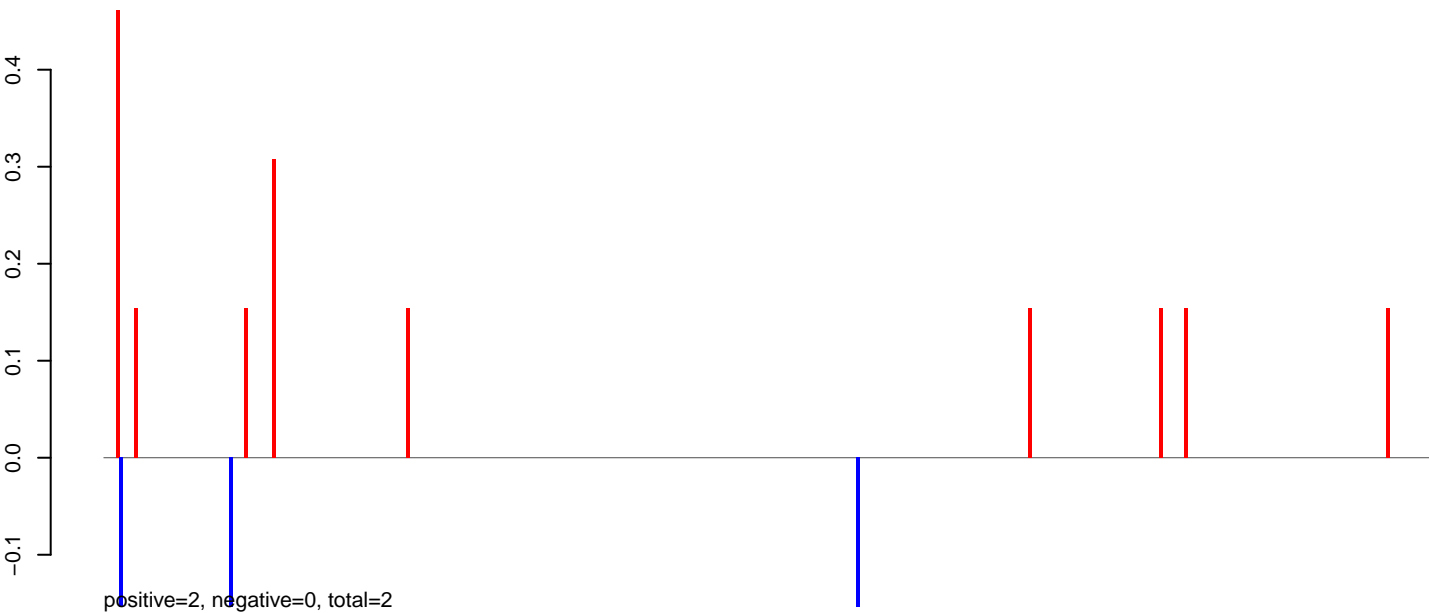
AeAlbo_C636_6dpi_SINV1.rep



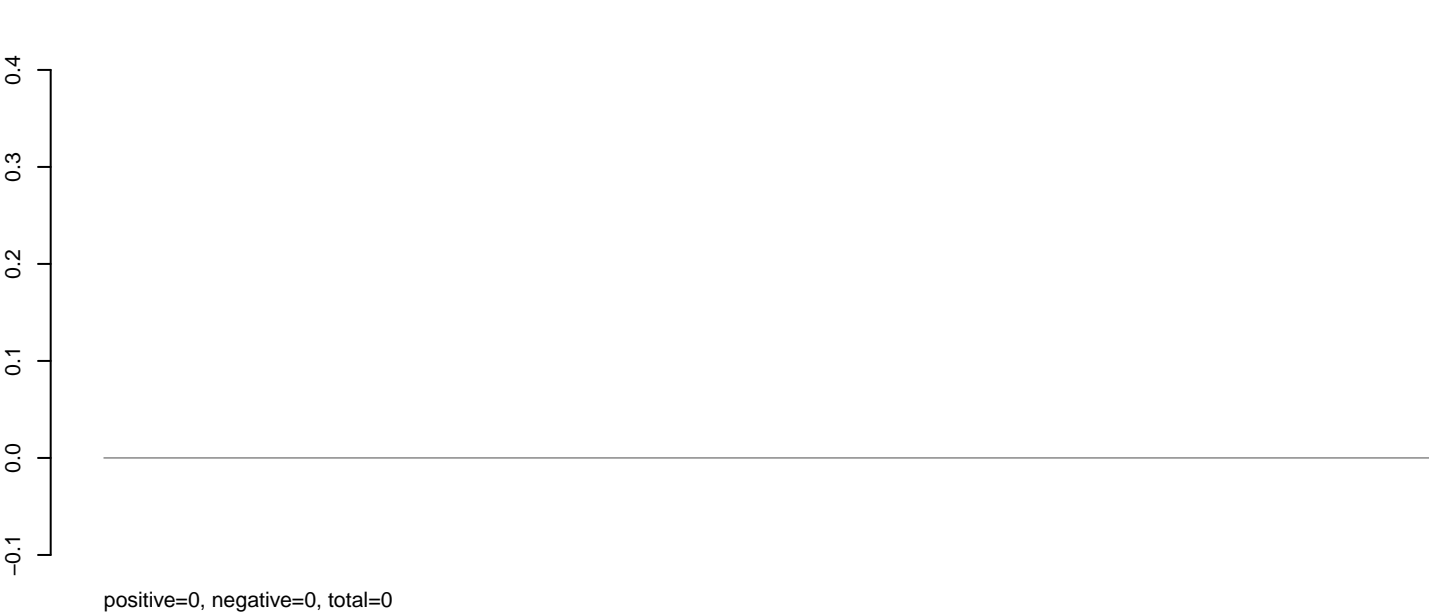
v size=25, length=11652, nt=65, cov=0.64, @1X-65 553.1-Gambie_virus_putative_nucleoprotein_putative_small_transmembrane_protein_putative_g:1-11652

0 2000 4000 6000 8000 10000 12000

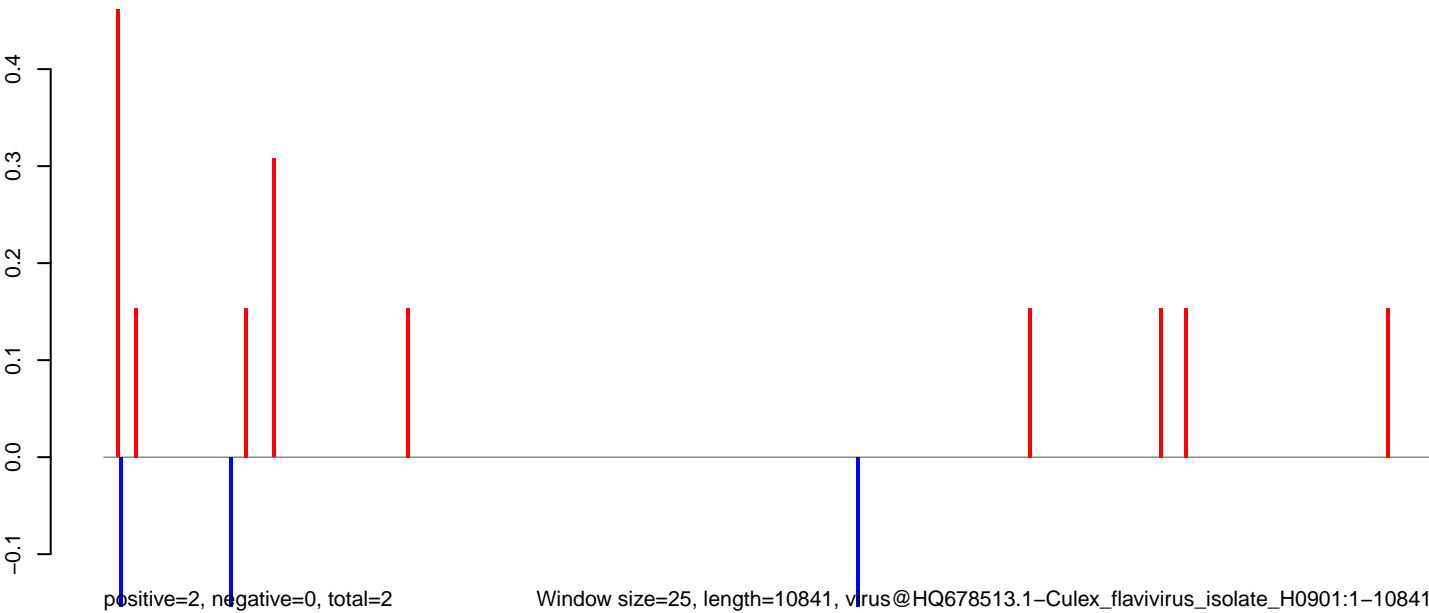
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep

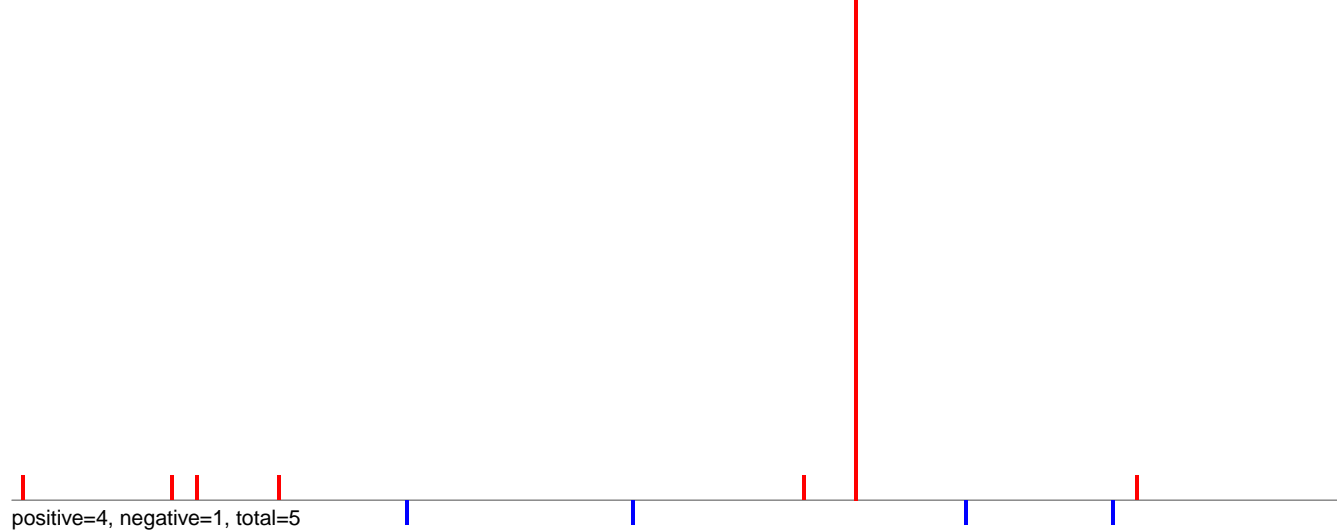


AeAlbo_C636_6dpi_SINV1.rep

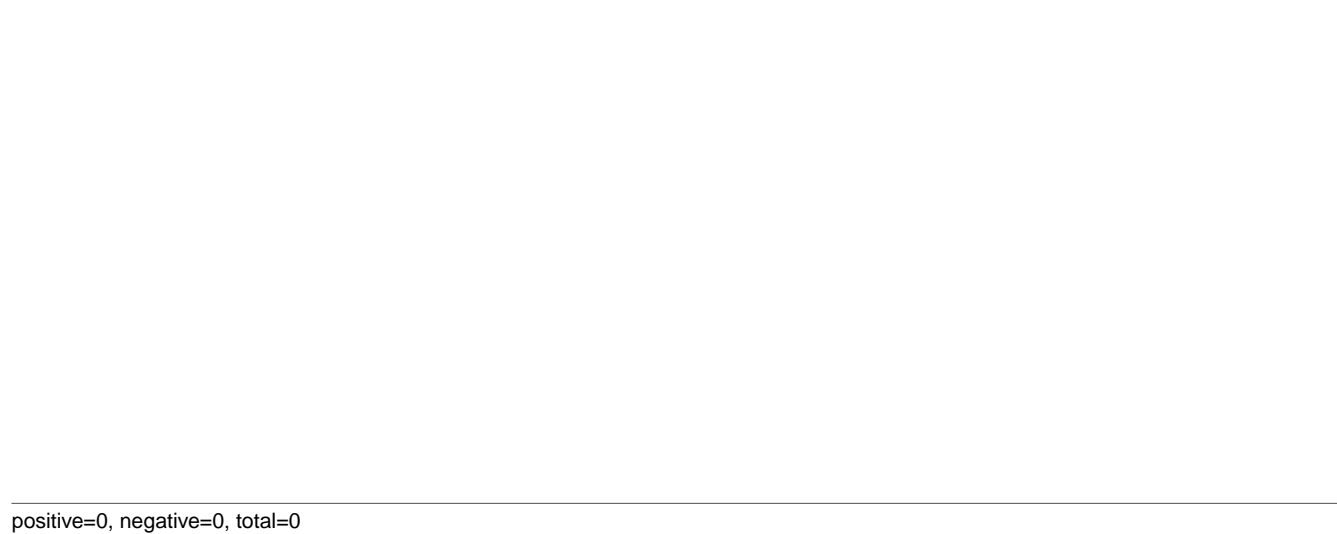


Window size=25, length=10841, virus@HQ678513.1-Culex_flavivirus_isolate_H0901:1-10841

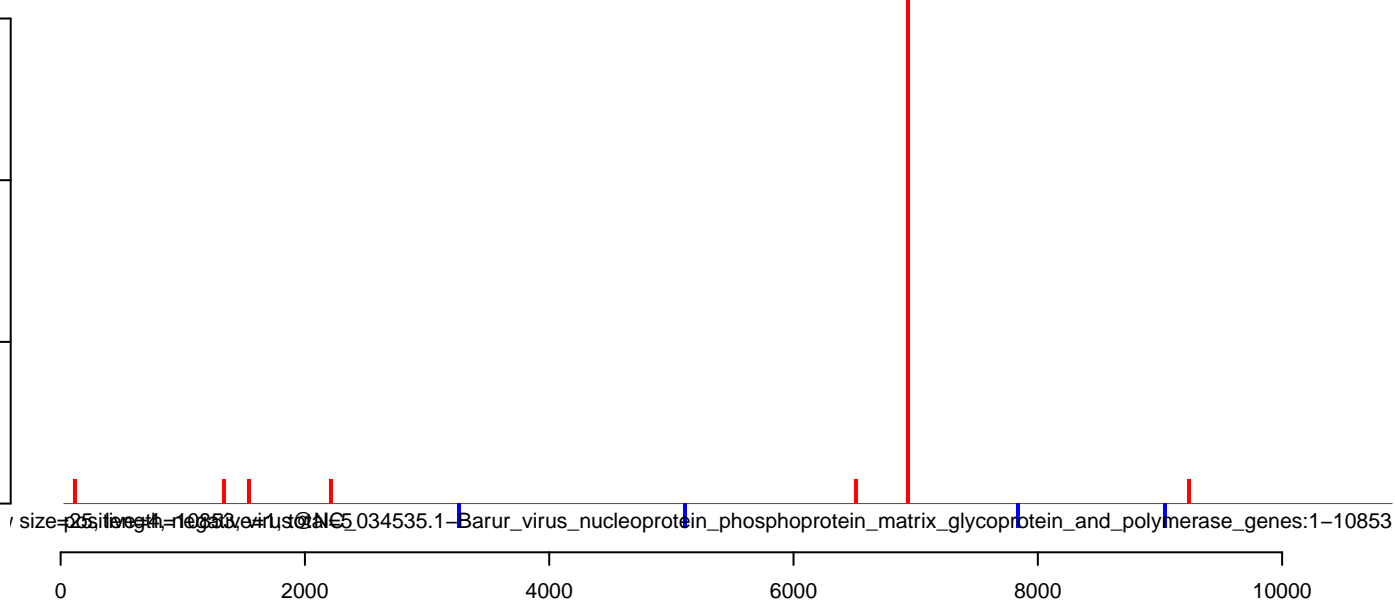
AeAlbo_C636_6dpi_SINV1.18_23.rep



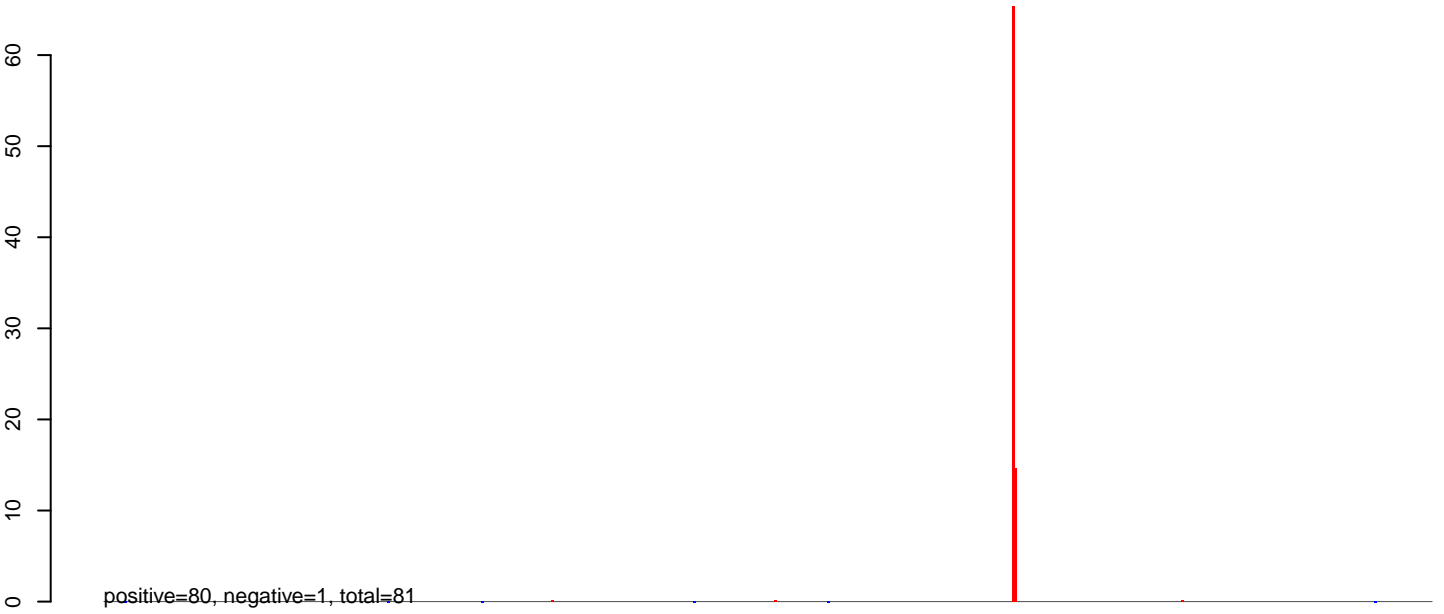
AeAlbo_C636_6dpi_SINV1.24_35.rep



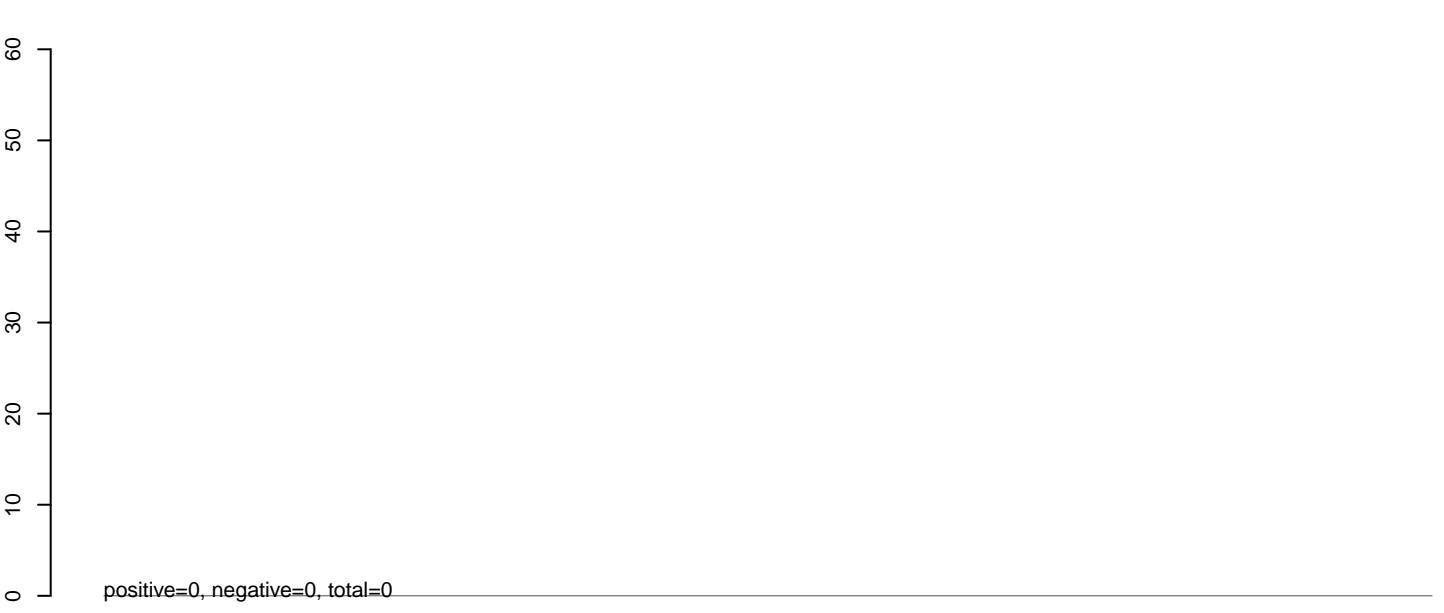
AeAlbo_C636_6dpi_SINV1.rep



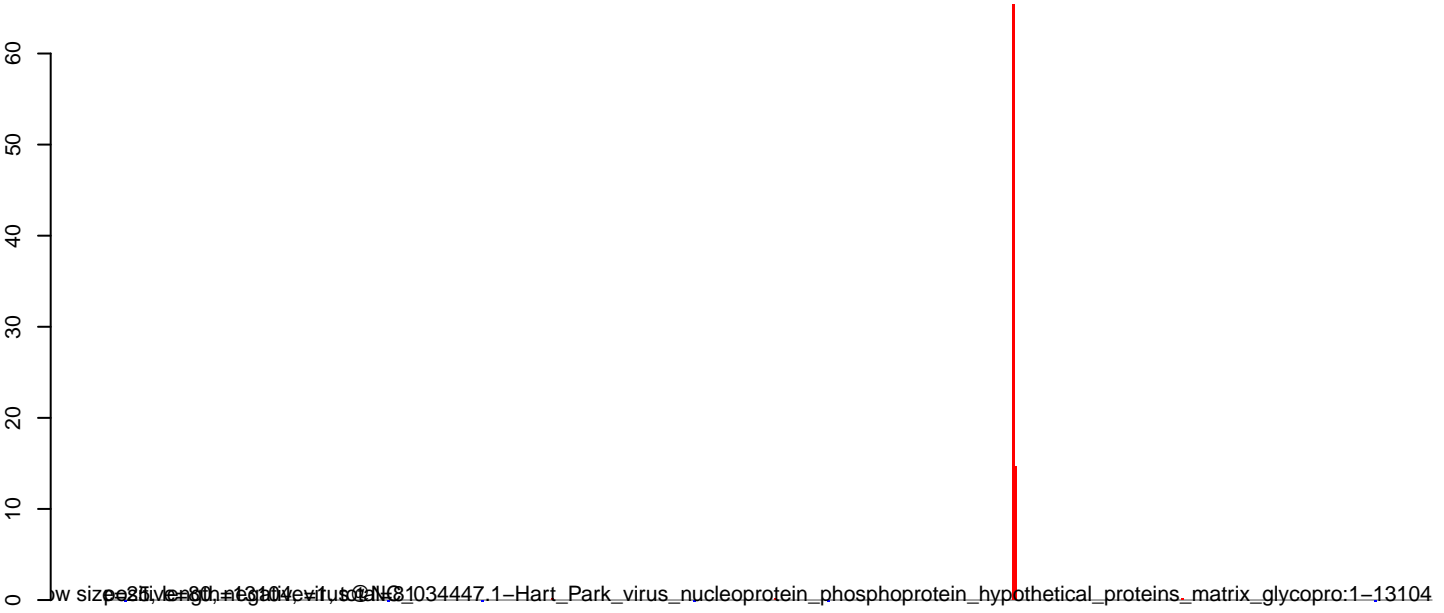
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



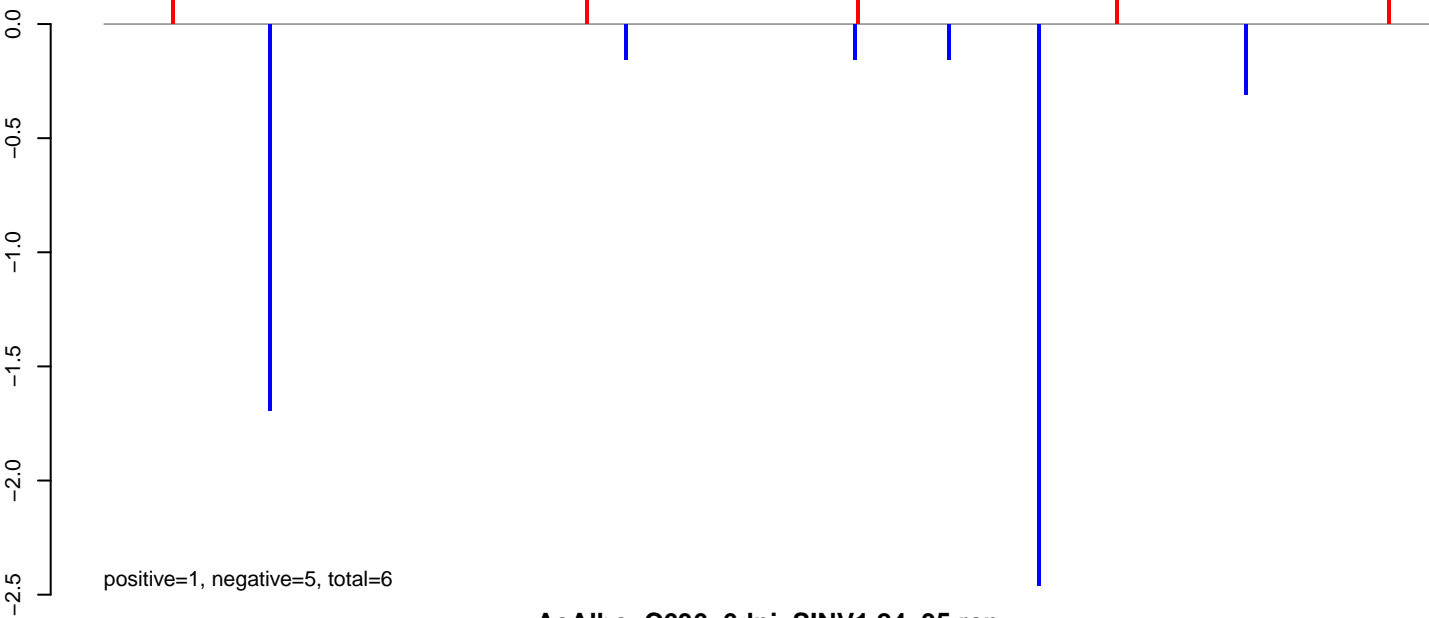
AeAlbo_C636_6dpi_SINV1.rep



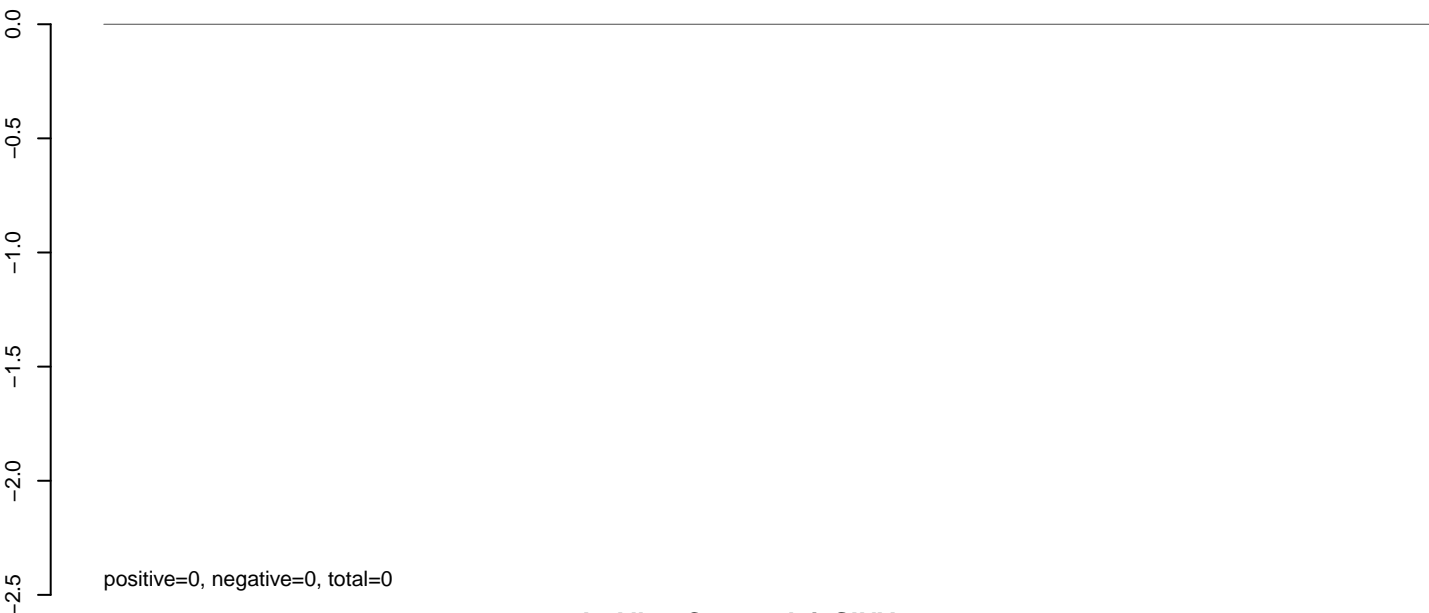
low size positive=80, negative=1, total=81 virus@NC_01034447.1-Hart_Park_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopro:1-13104

0 2000 4000 6000 8000 10000 12000

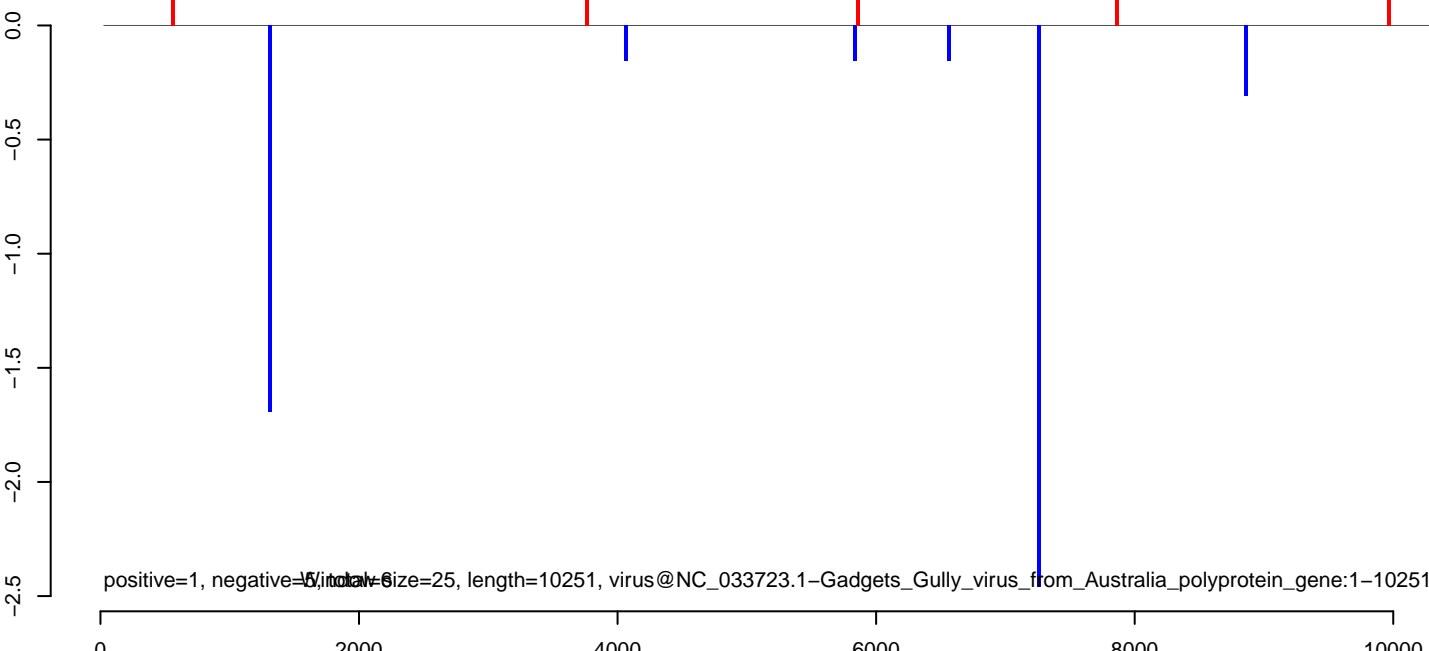
AeAlbo_C636_6dpi_SINV1.18_23.rep



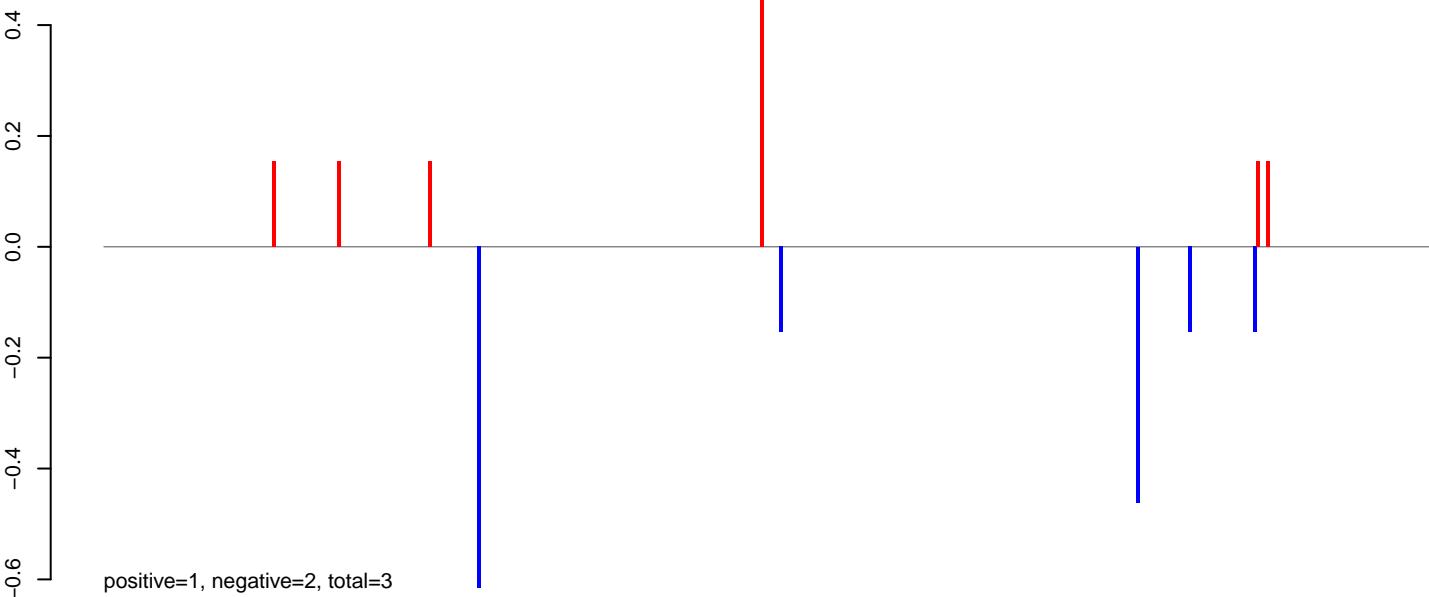
AeAlbo_C636_6dpi_SINV1.24_35.rep



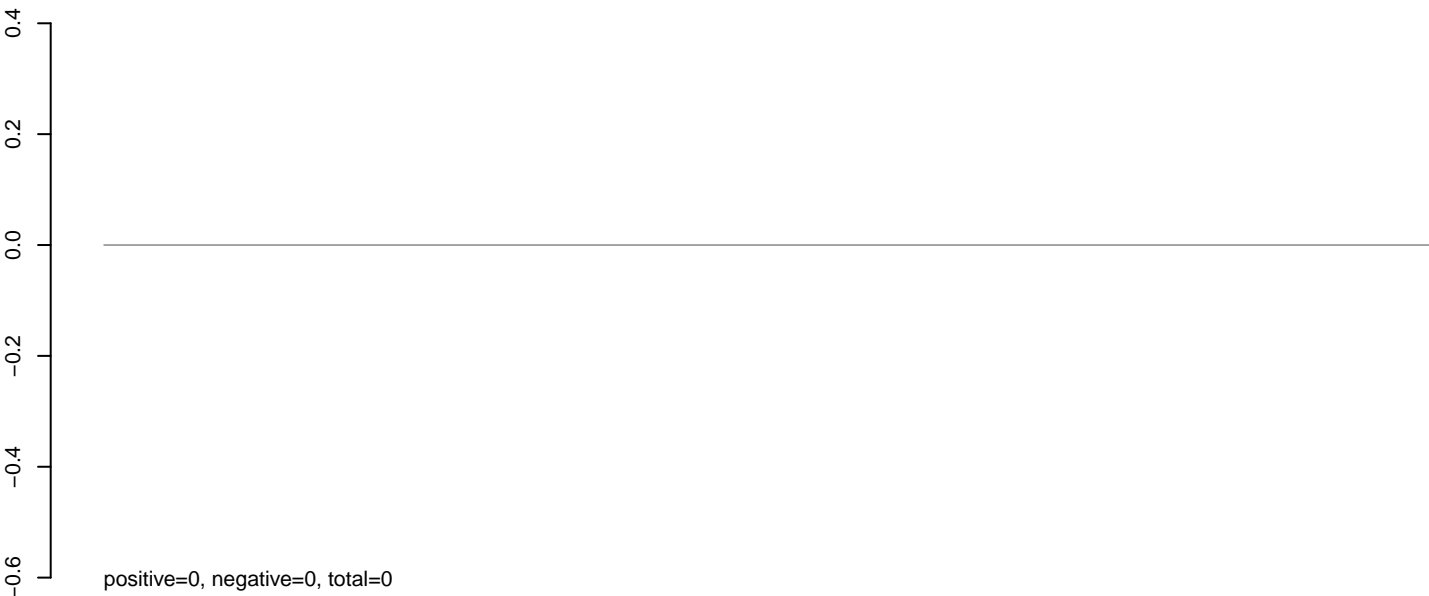
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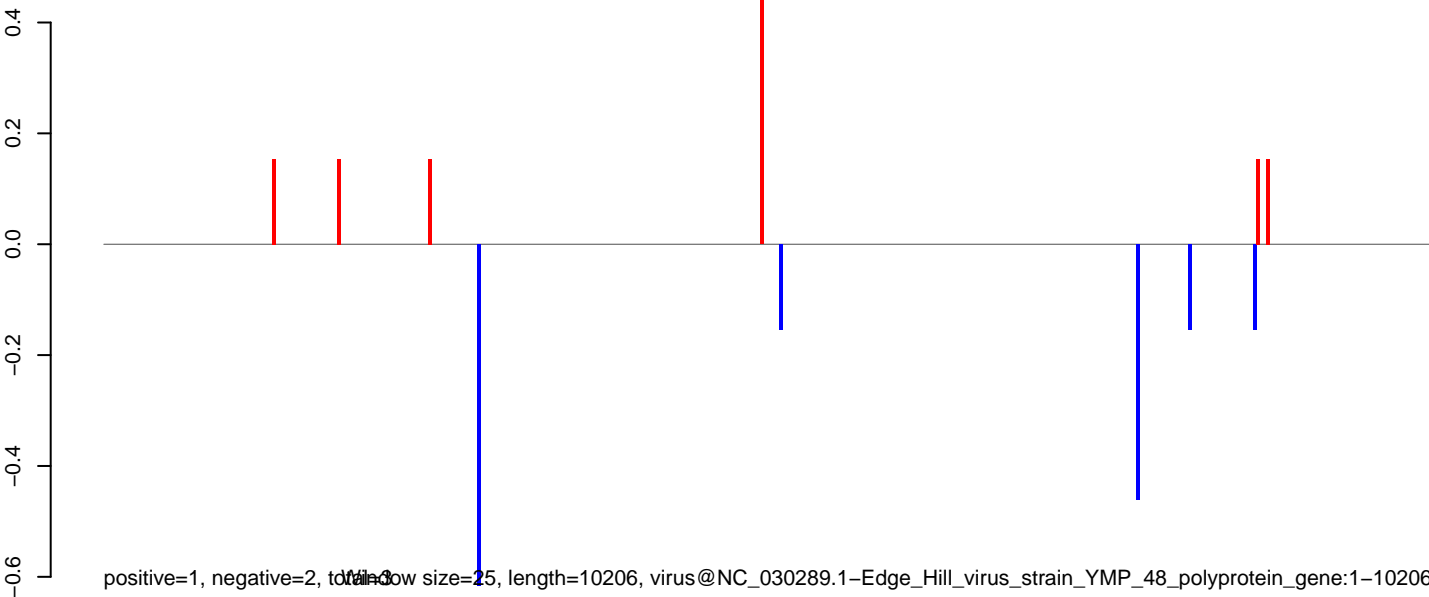
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



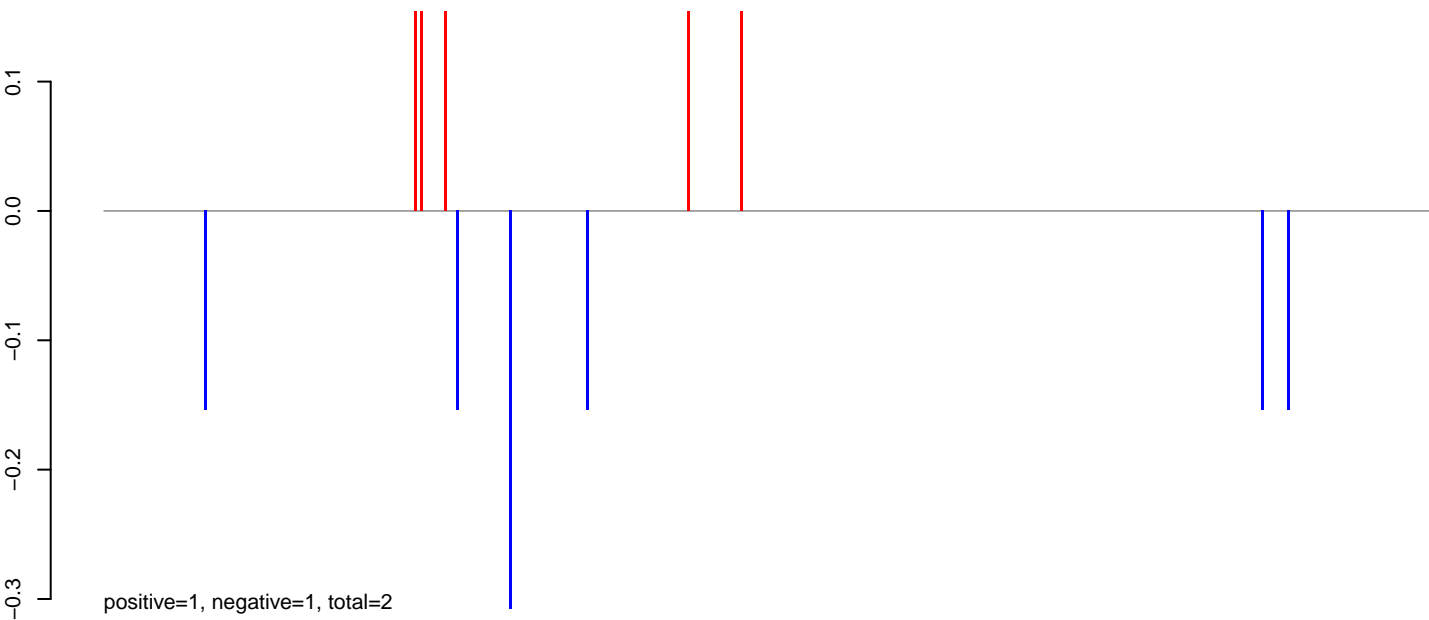
AeAlbo_C636_6dpi_SINV1.rep



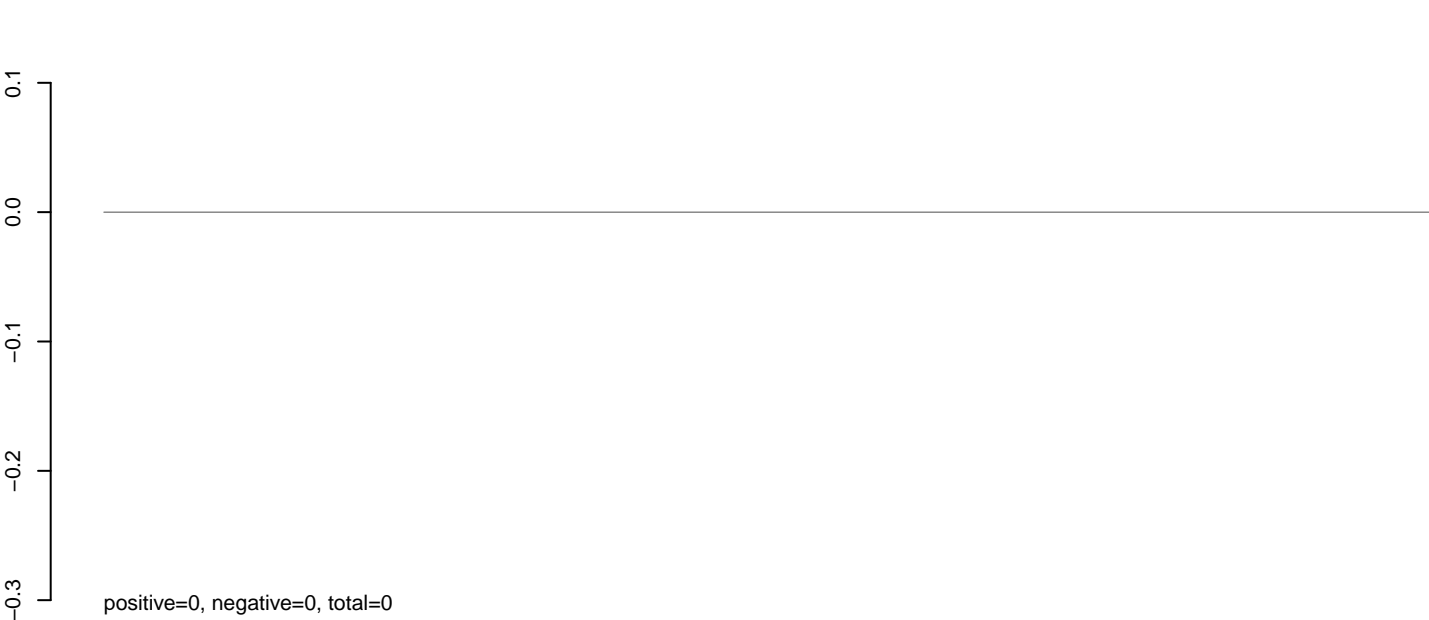
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0 2000 4000 6000 8000 10000

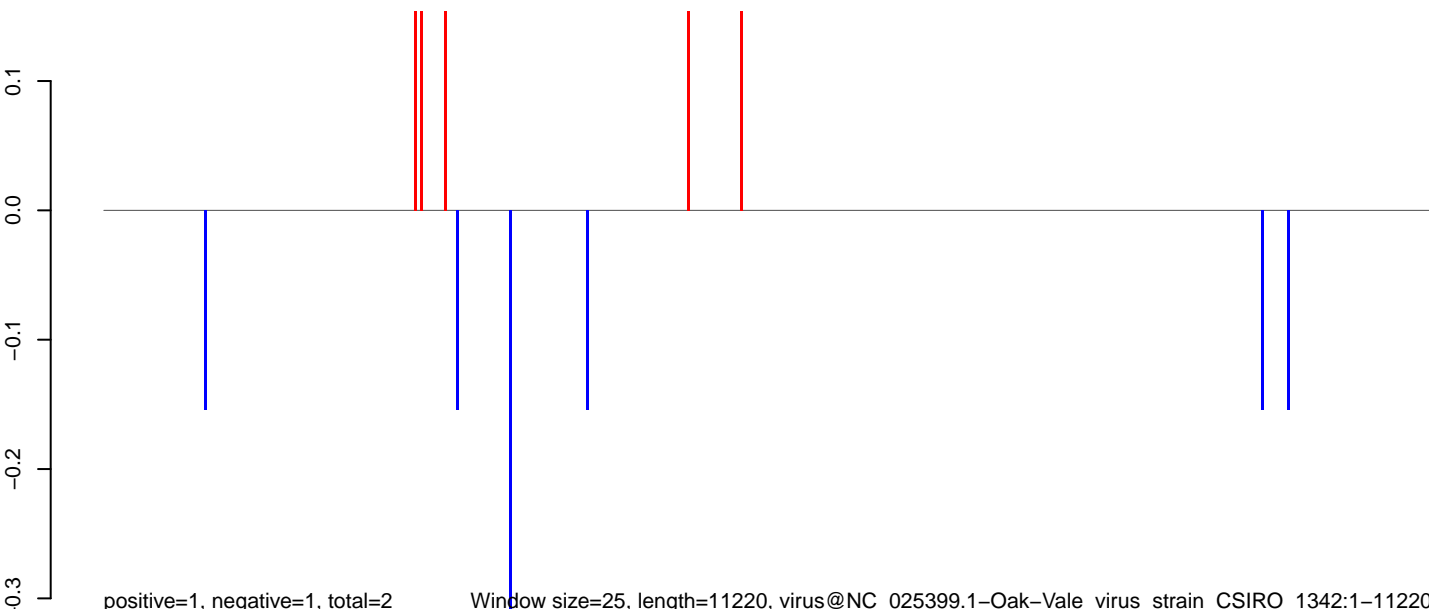
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



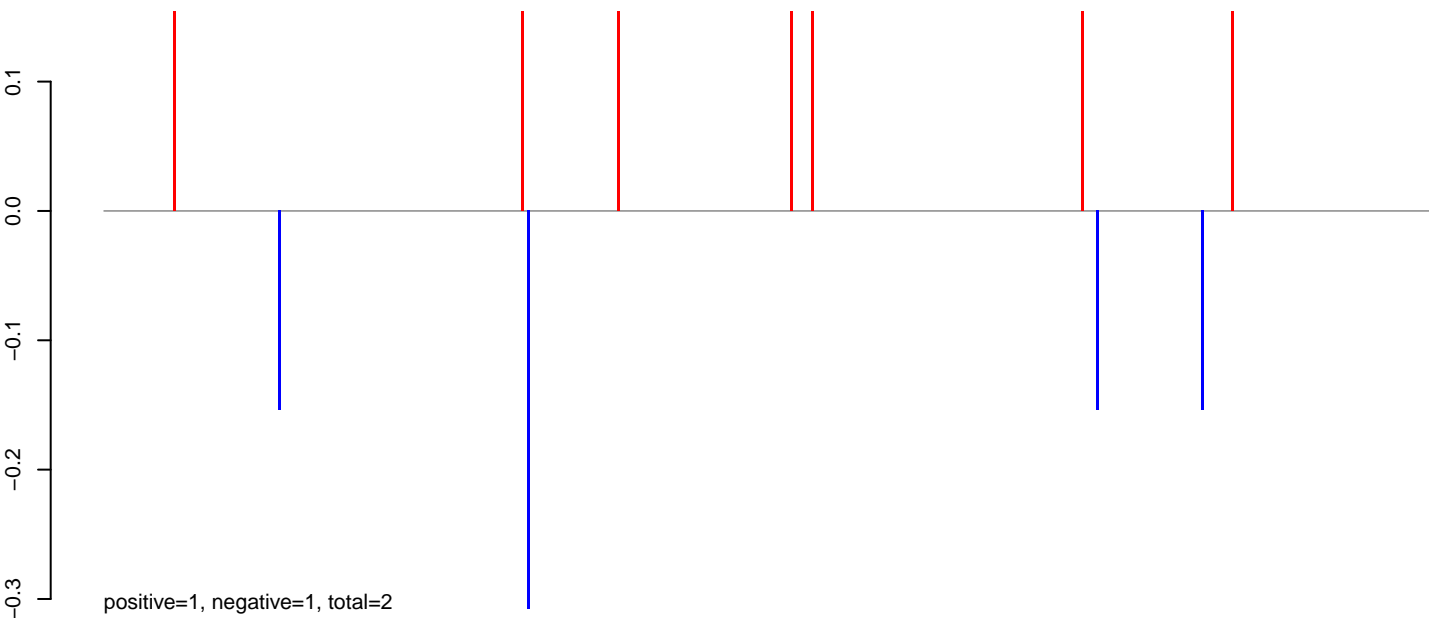
AeAlbo_C636_6dpi_SINV1.rep



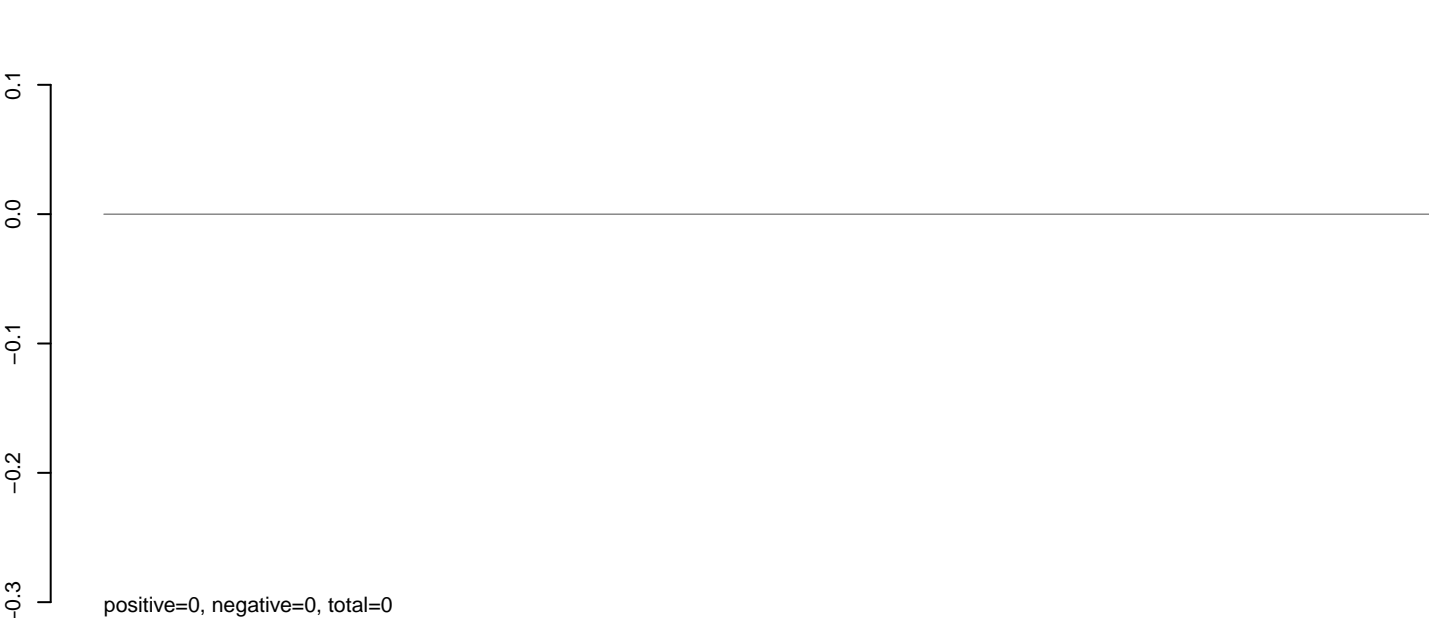
Window size=25, length=11220, virus@NC_025399.1-Oak-Vale_virus_strain_CSIRO_1342:1-11220

0 2000 4000 6000 8000 10000

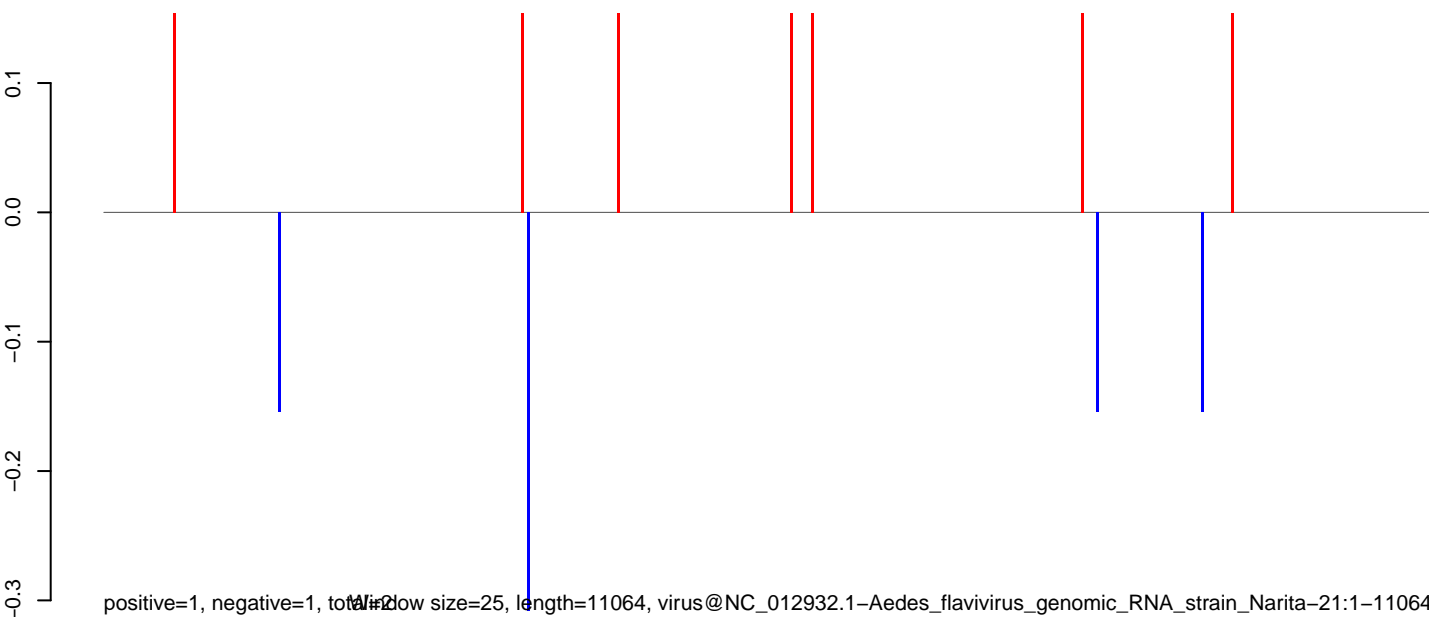
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



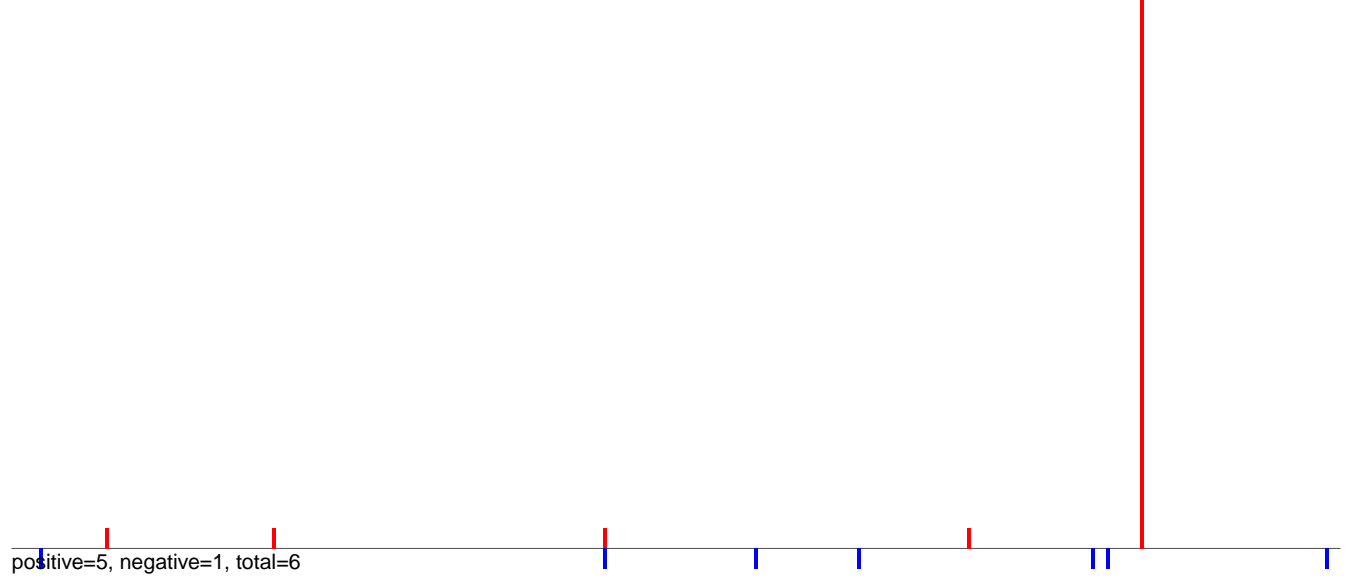
AeAlbo_C636_6dpi_SINV1.rep



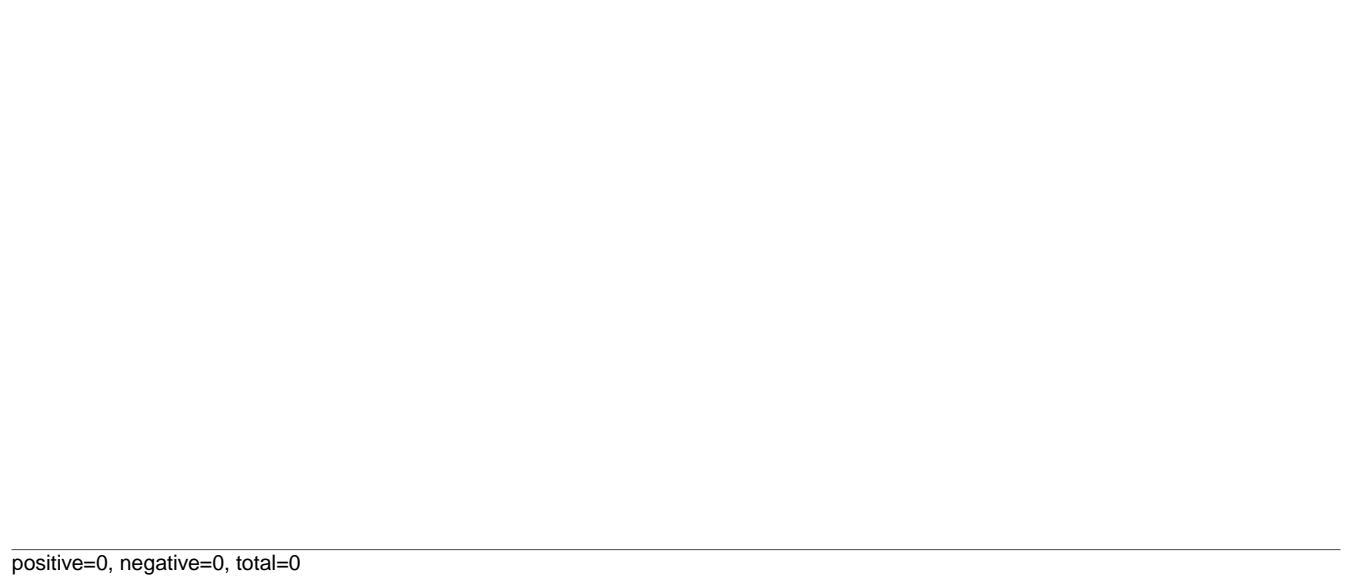
total size=25, length=11064, virus@NC_012932.1-Aedes_flavivirus_genomic_RNA_strain_Narita-21:1-11064

0 2000 4000 6000 8000 10000

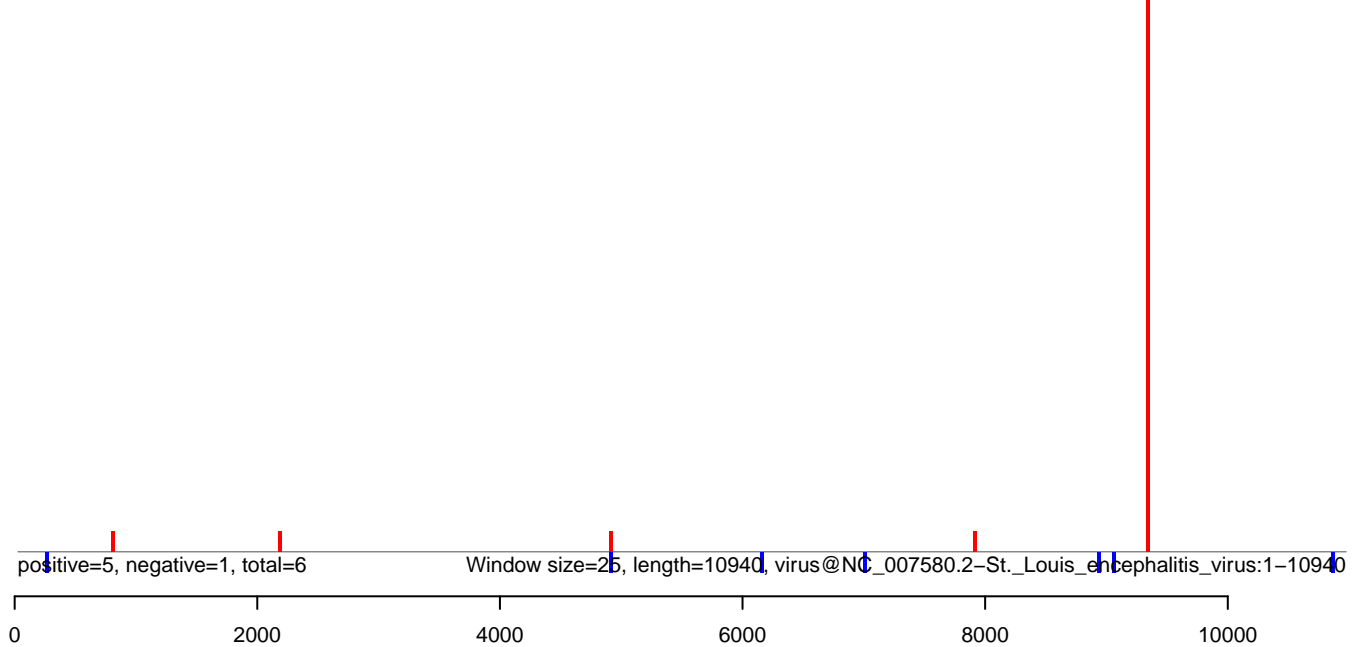
AeAlbo_C636_6dpi_SINV1.18_23.rep



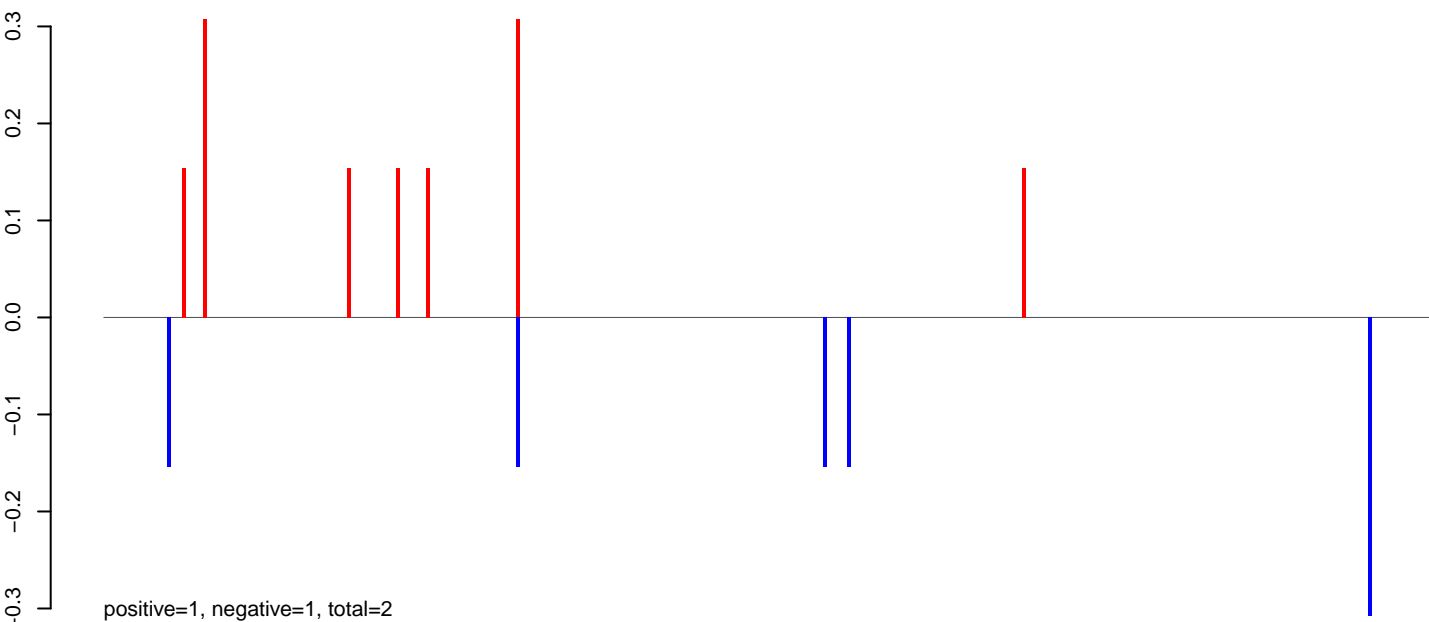
AeAlbo_C636_6dpi_SINV1.24_35.rep



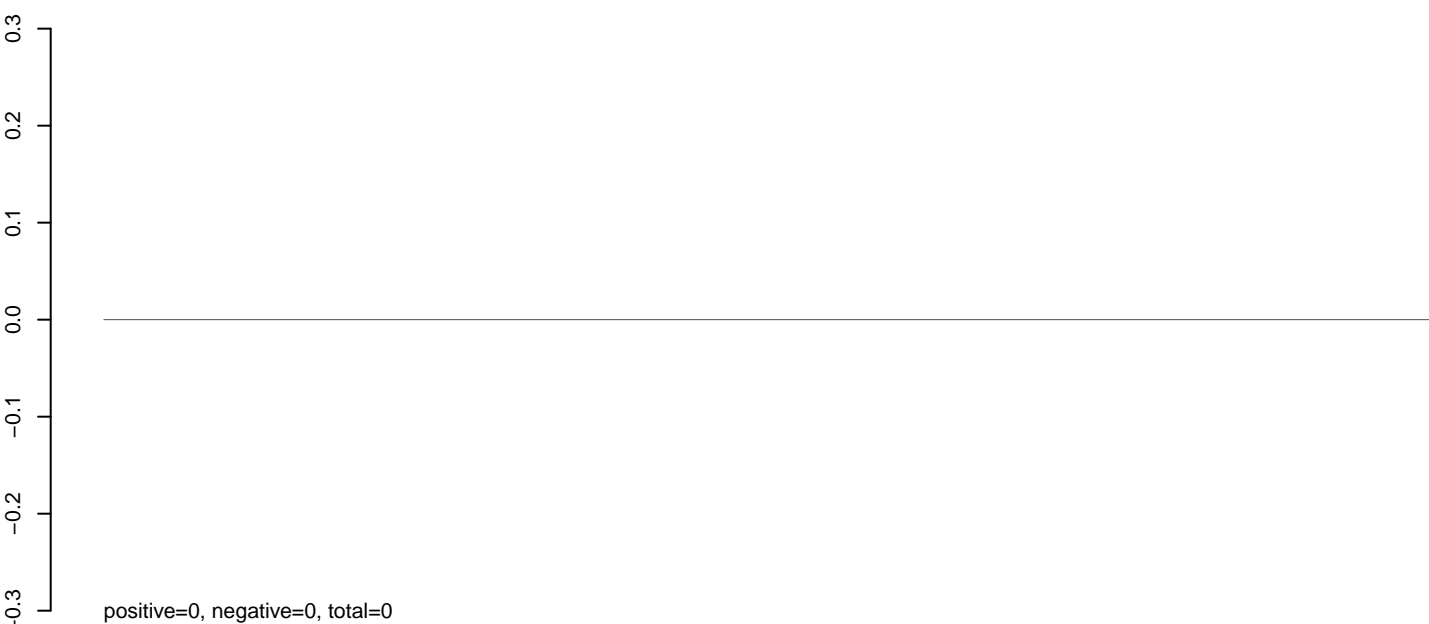
AeAlbo_C636_6dpi_SINV1.rep



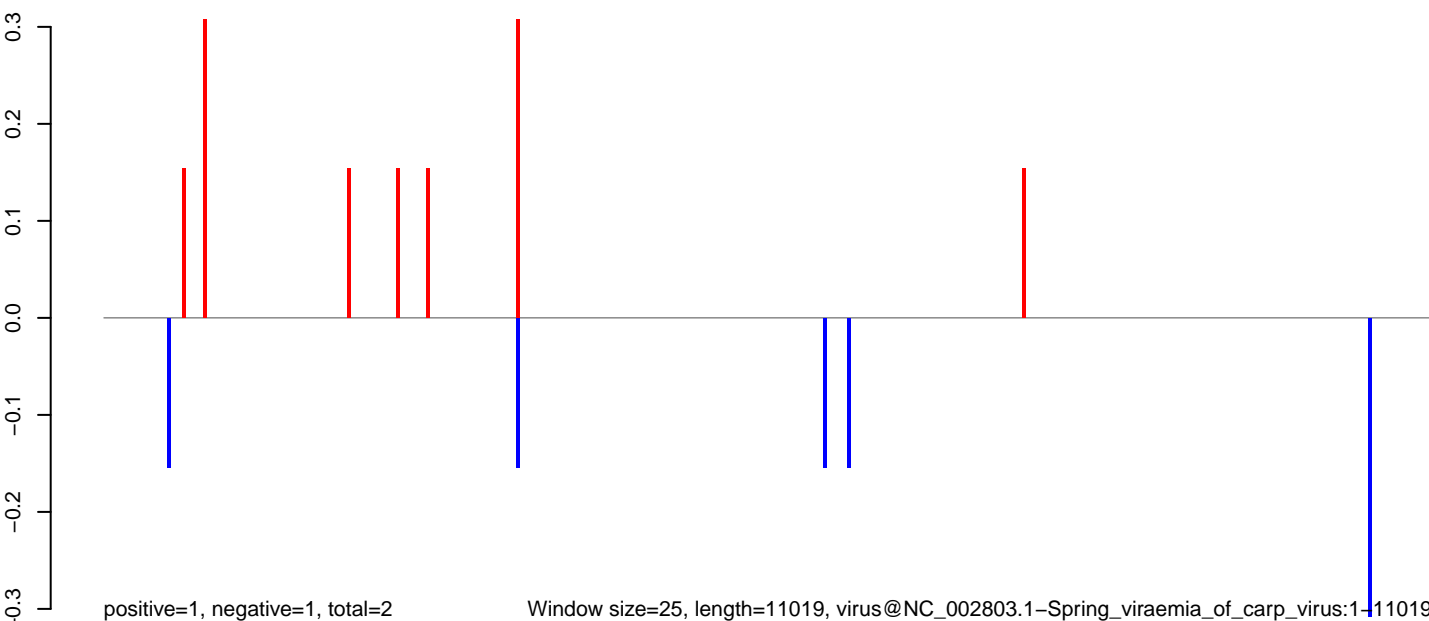
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



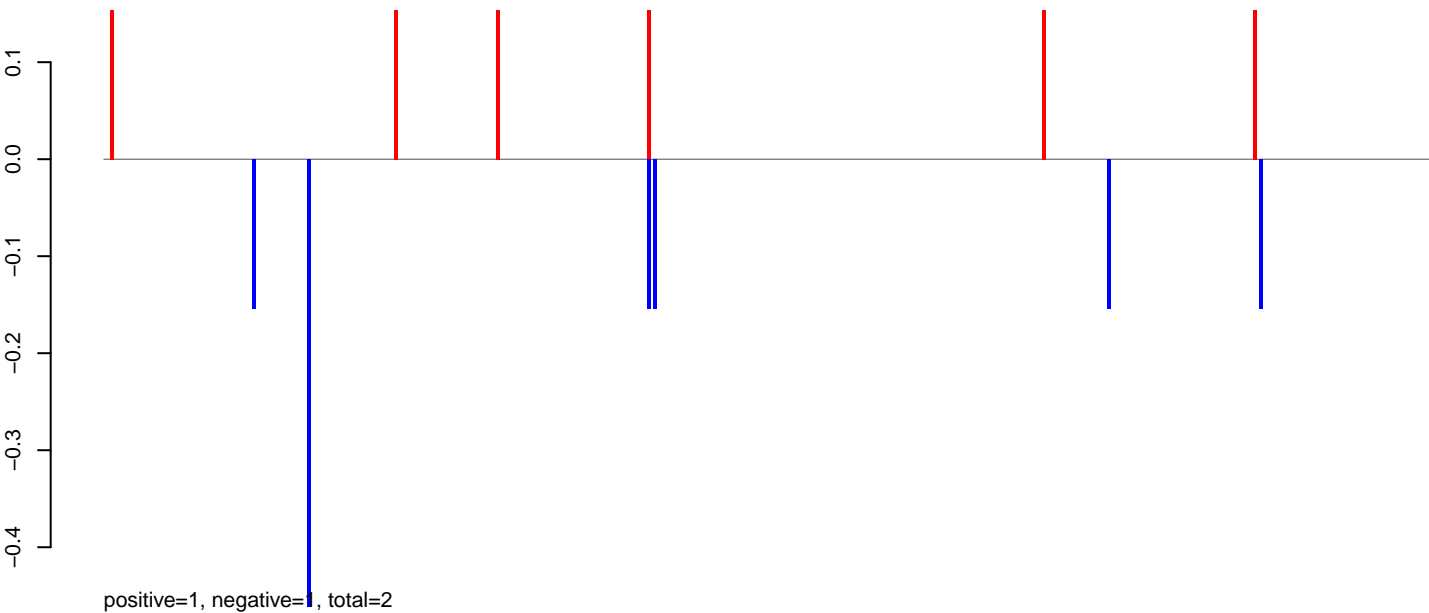
AeAlbo_C636_6dpi_SINV1.rep



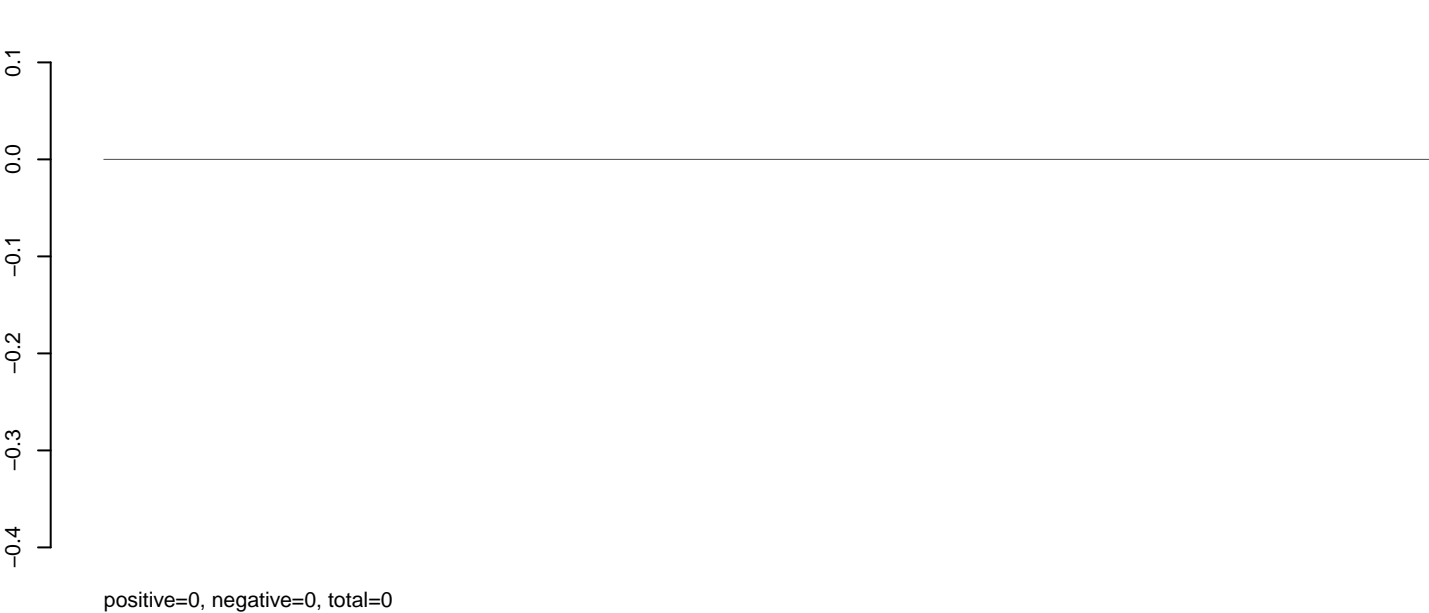
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0 2000 4000 6000 8000 10000

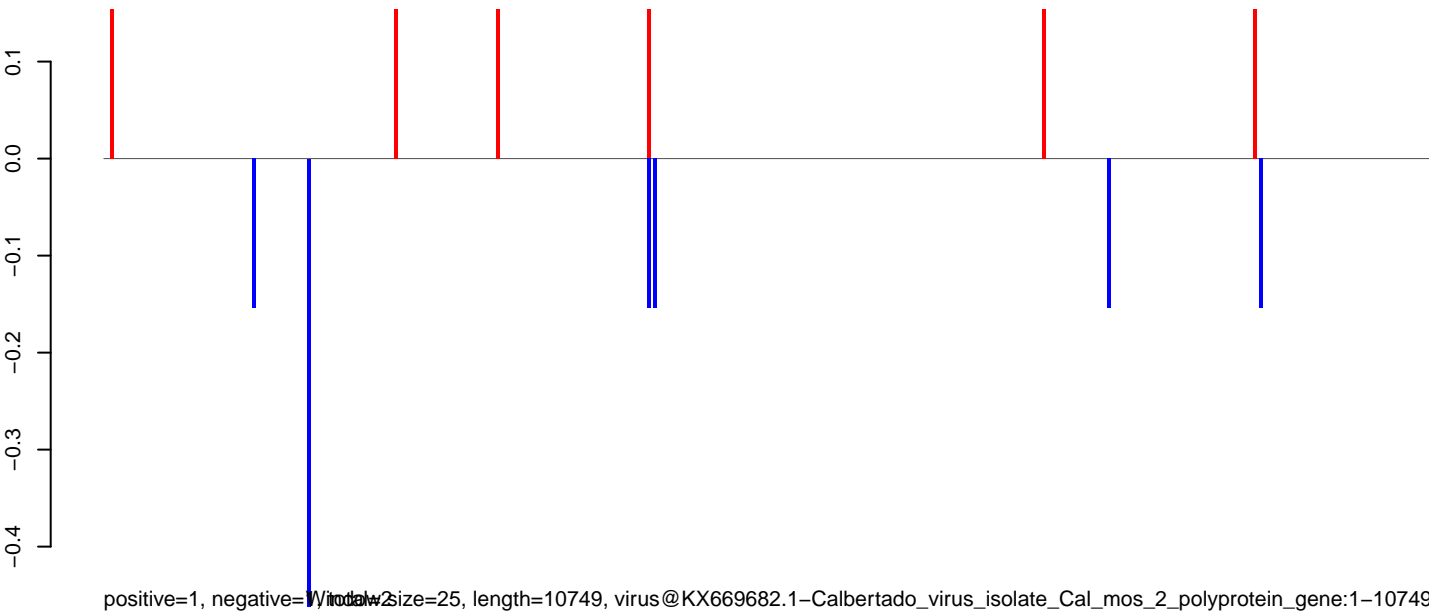
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



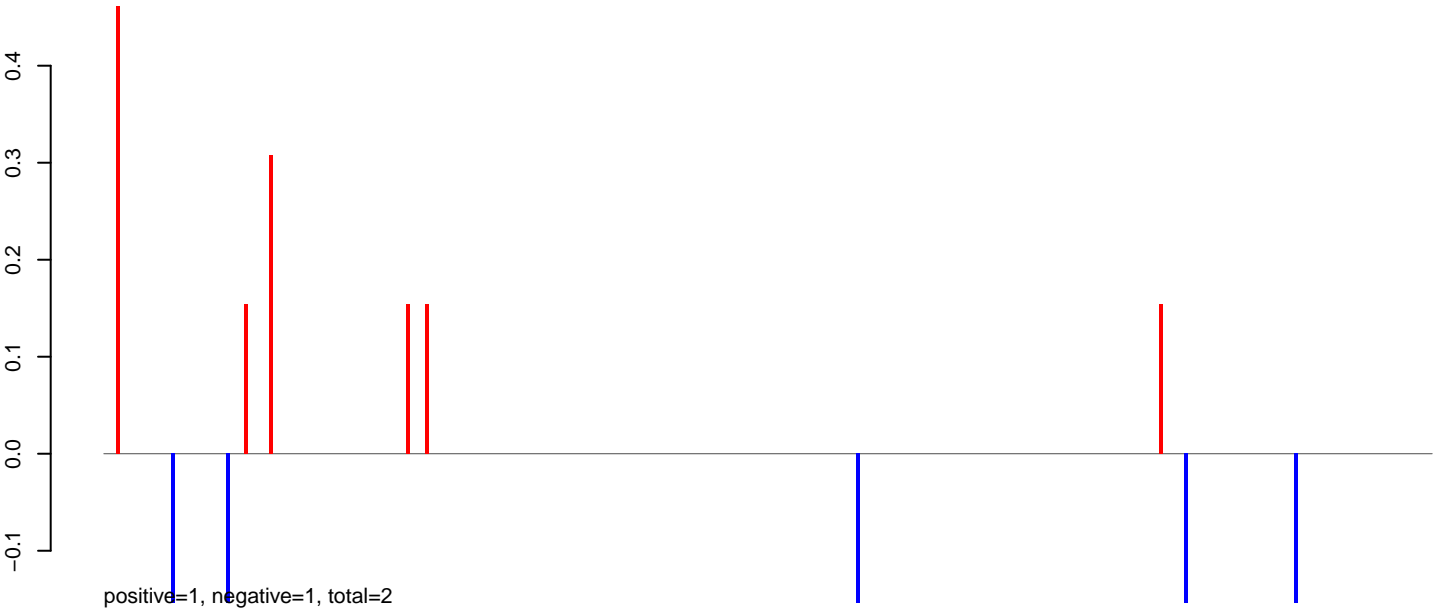
AeAlbo_C636_6dpi_SINV1.rep



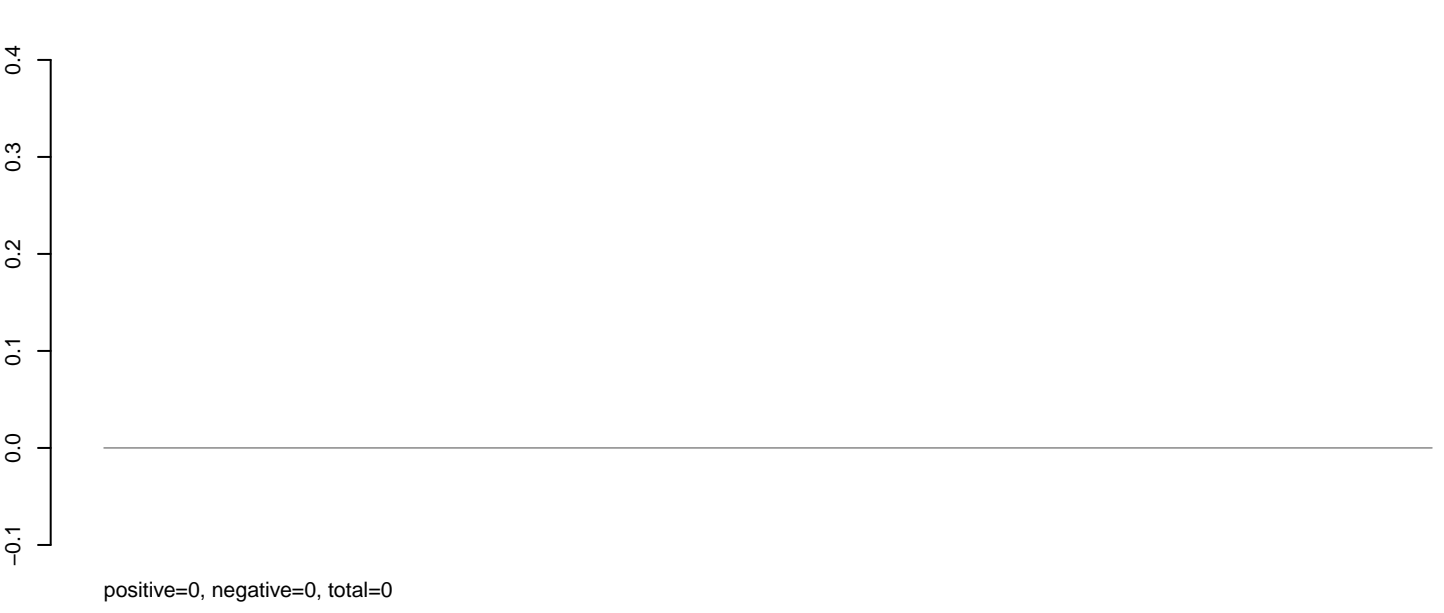
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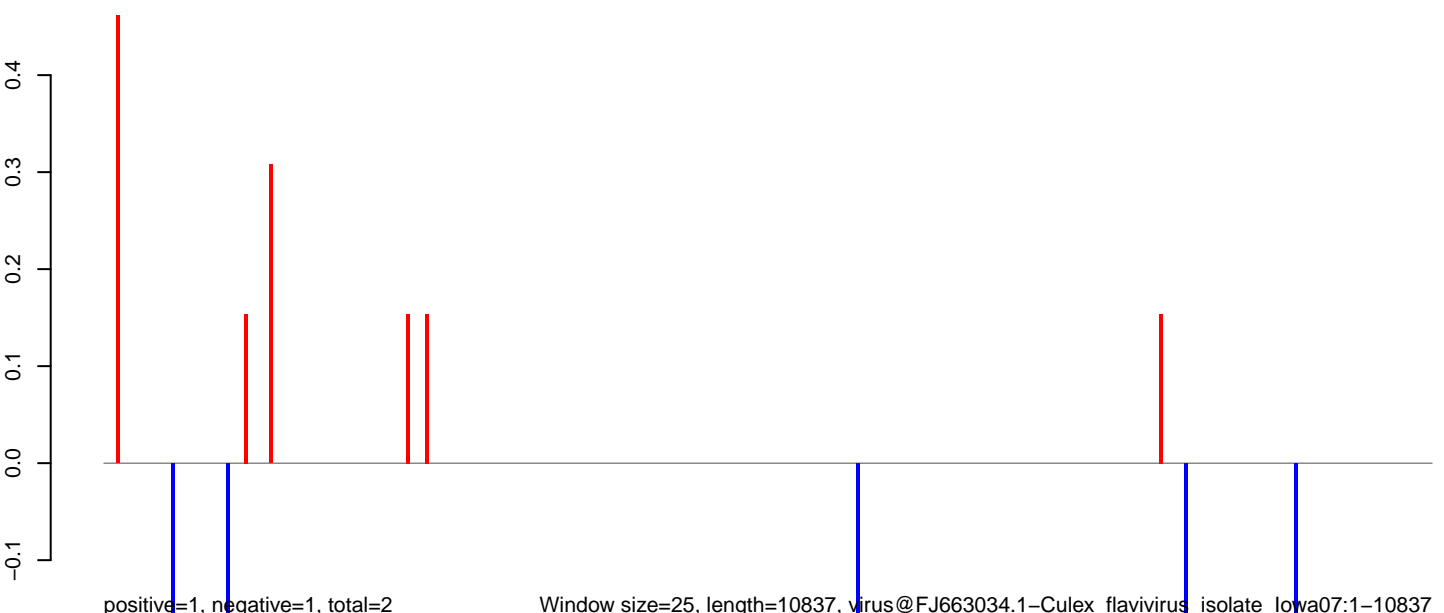
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AeAlbo_C636_6dpi_SINV1.24_35.rep

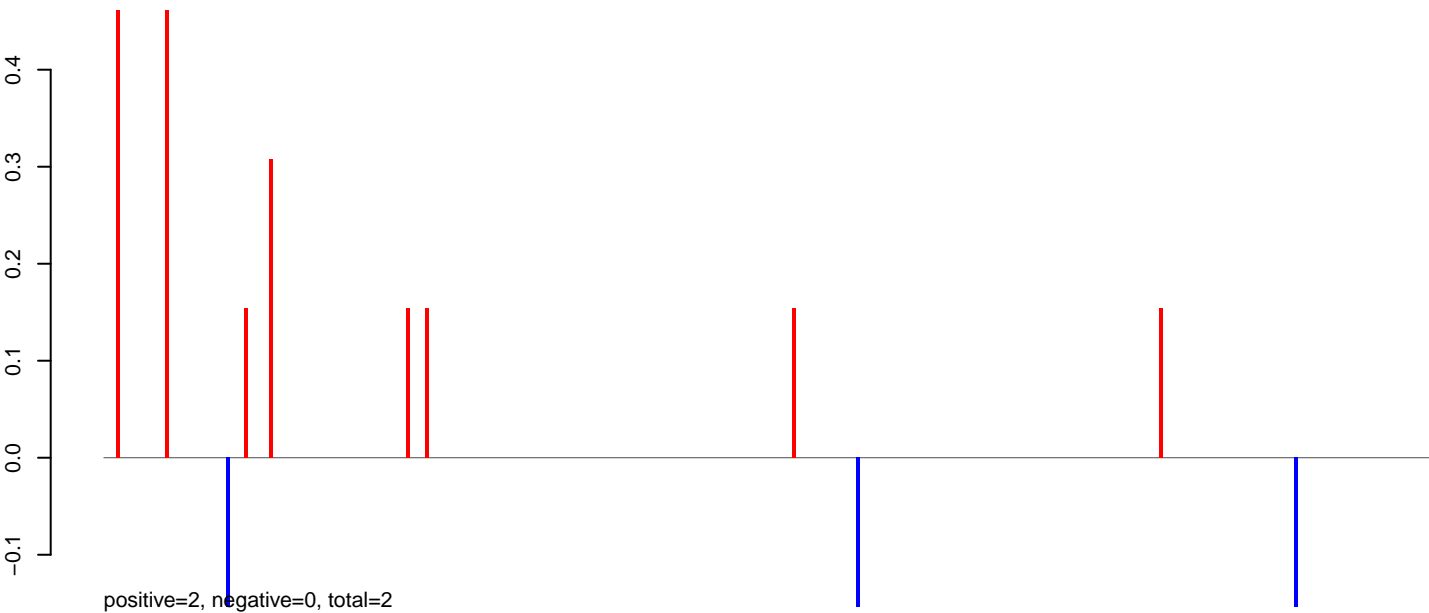


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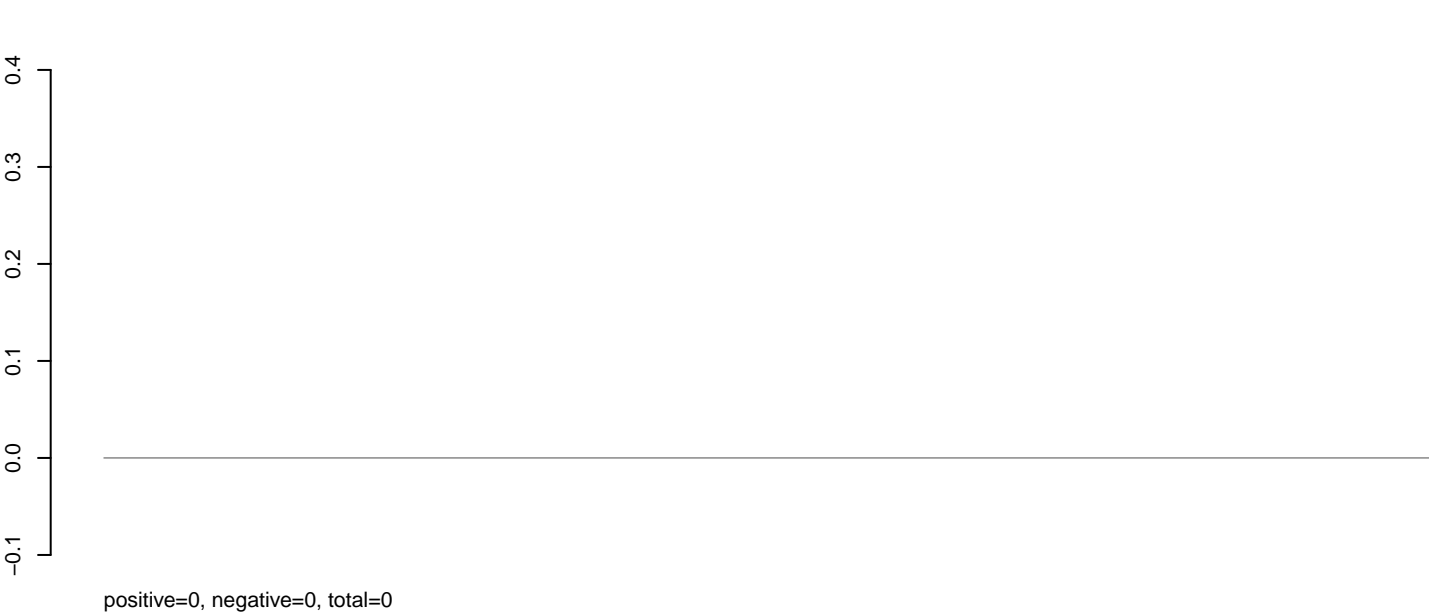


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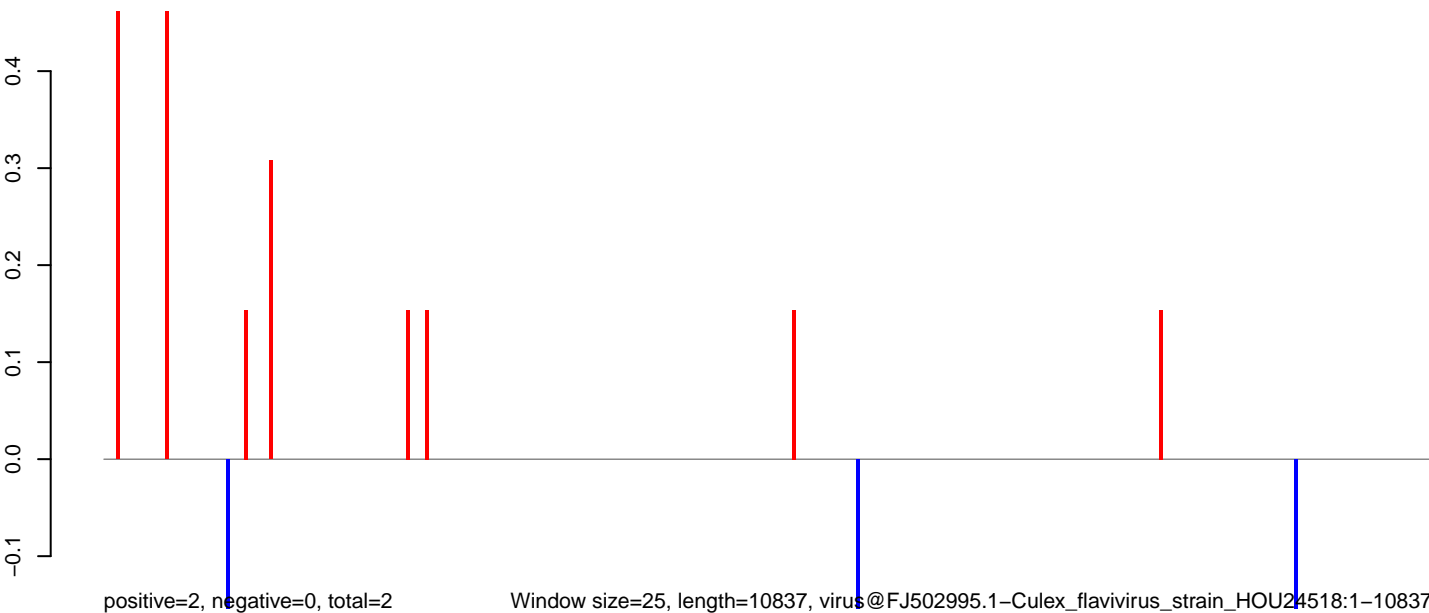
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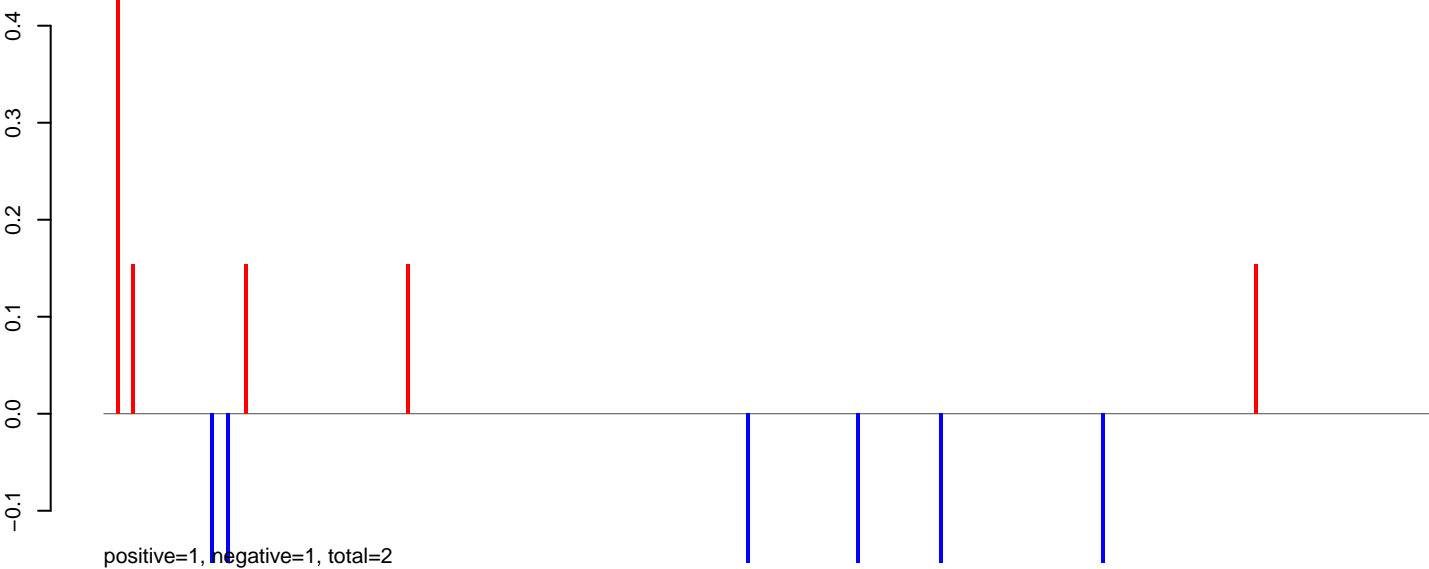
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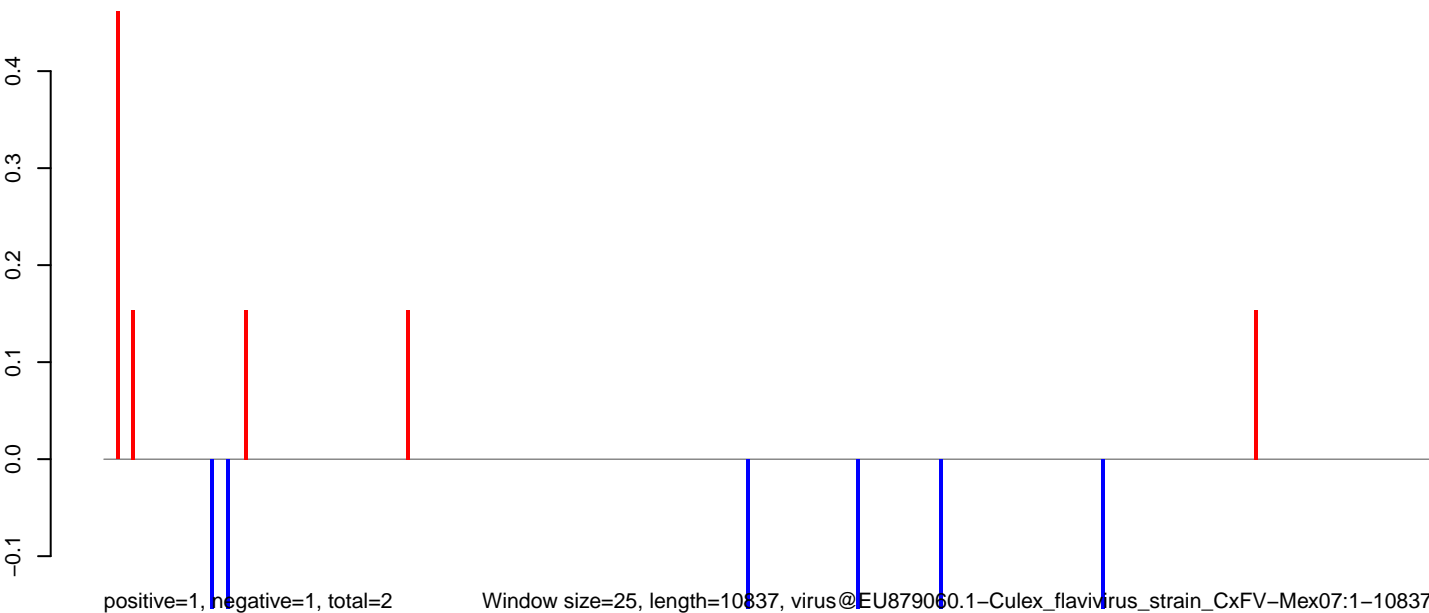
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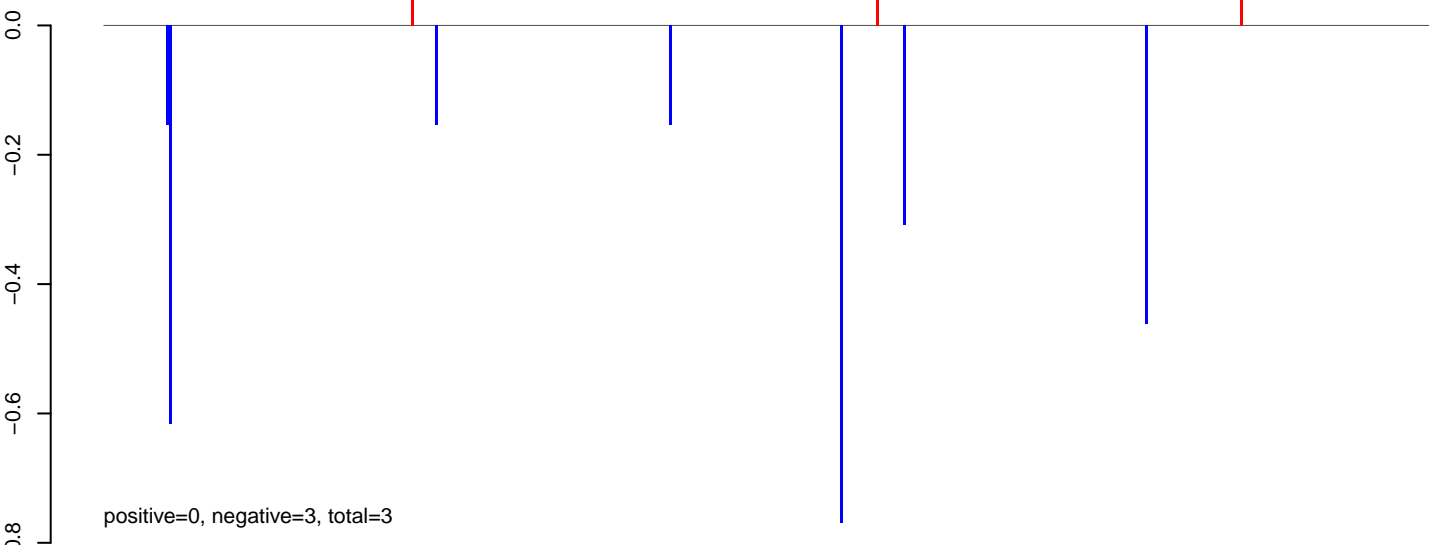
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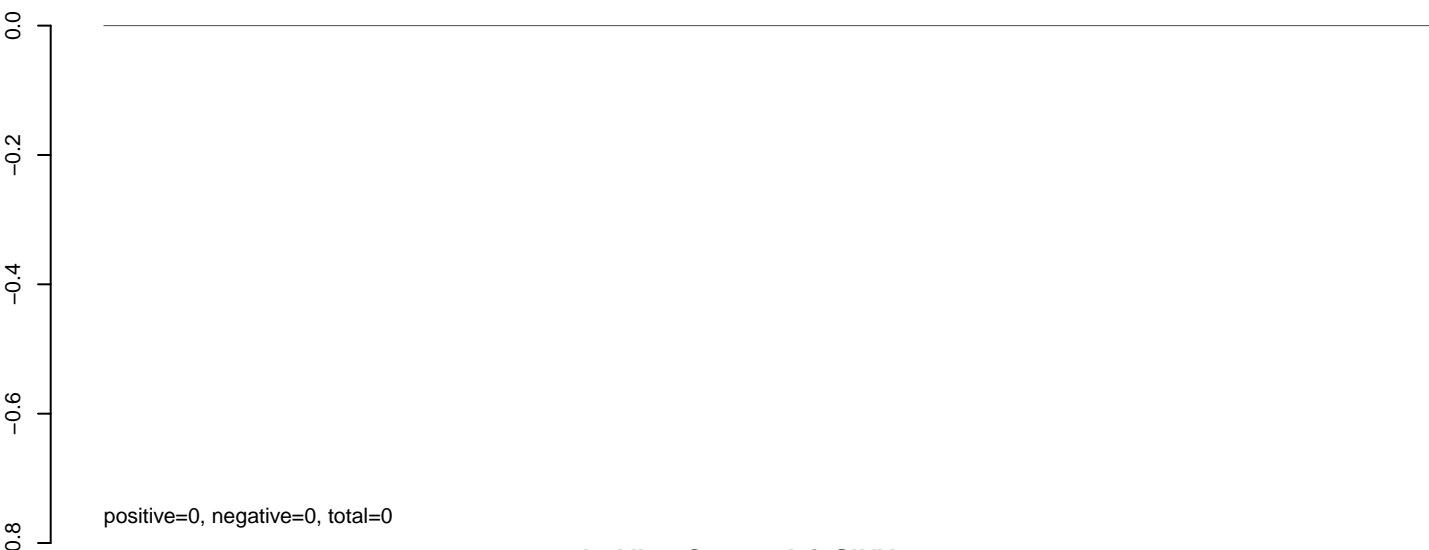
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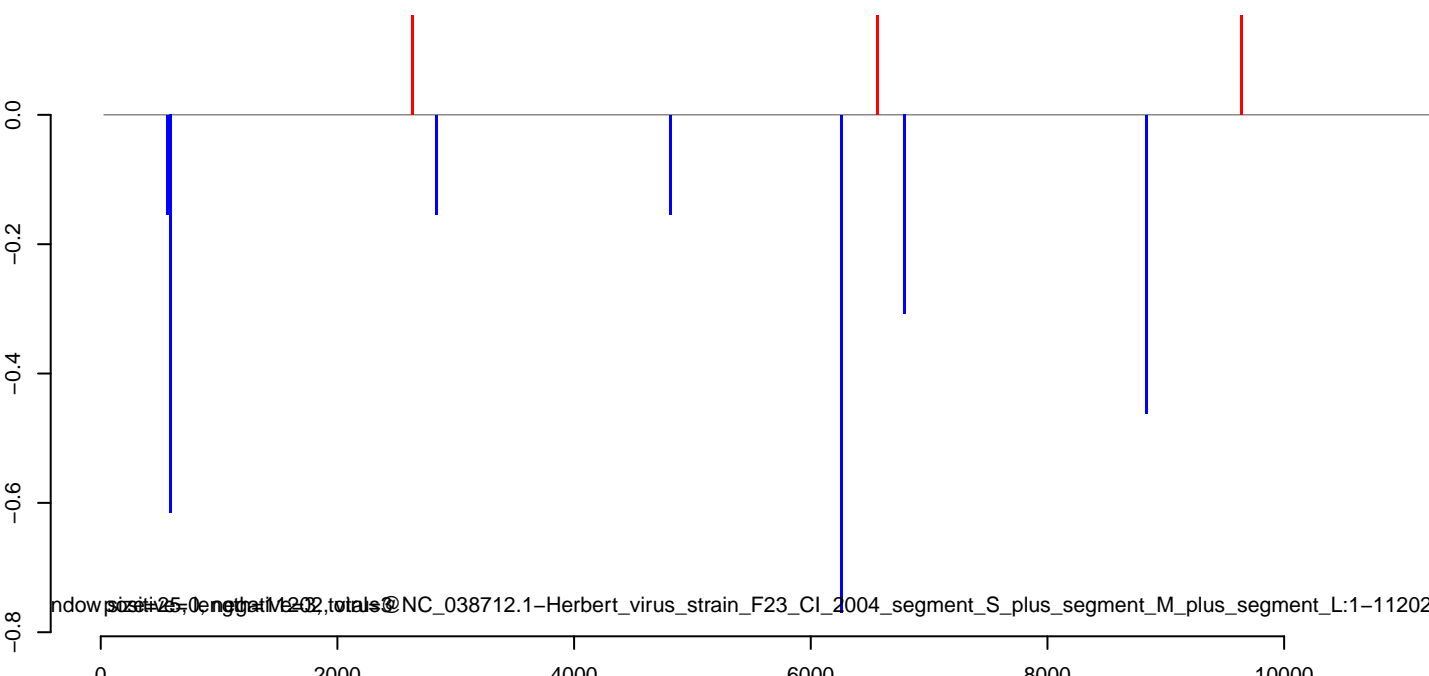
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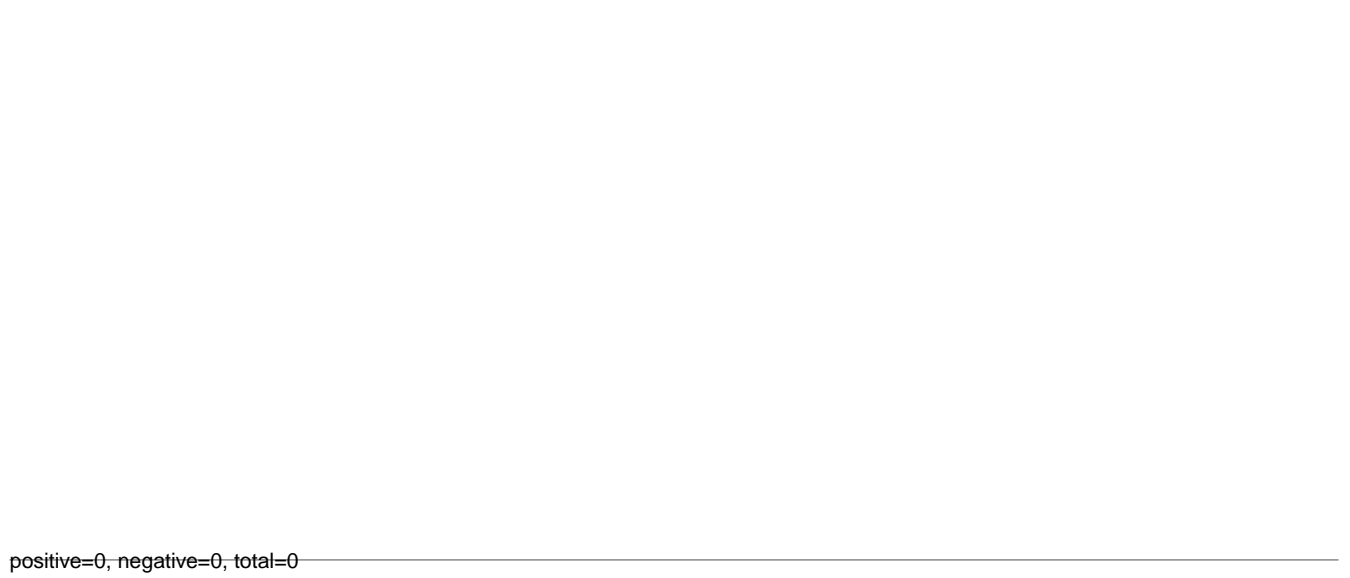
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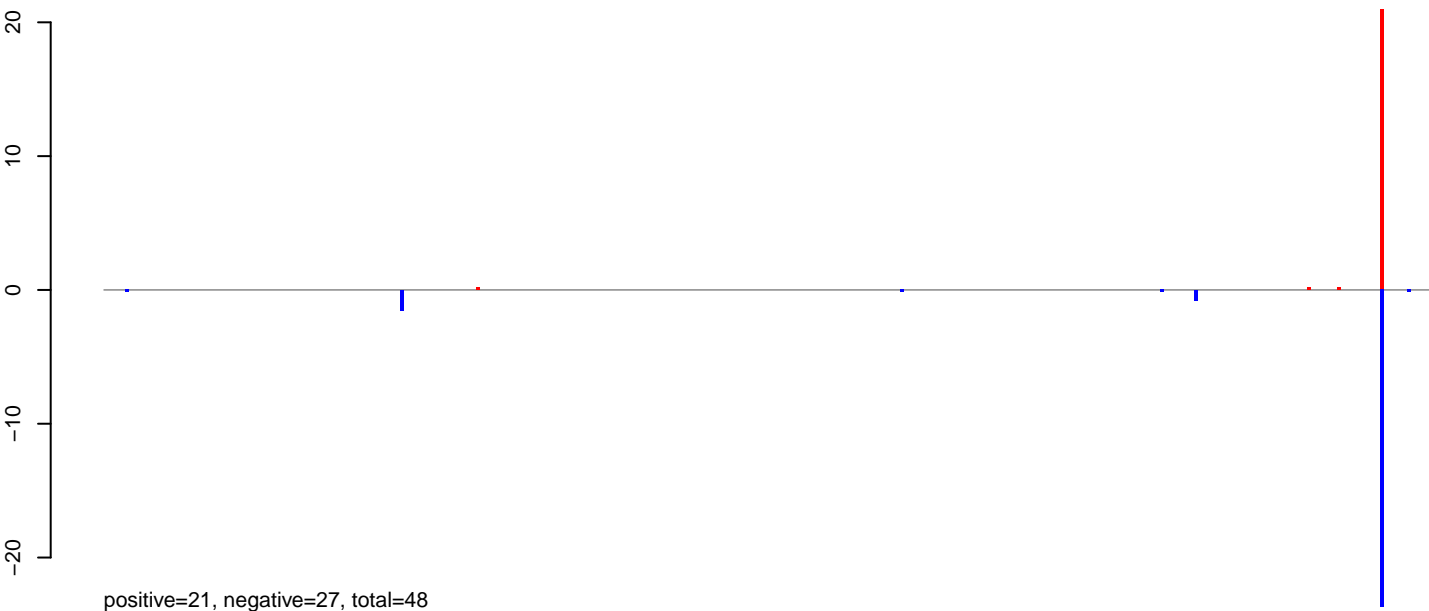
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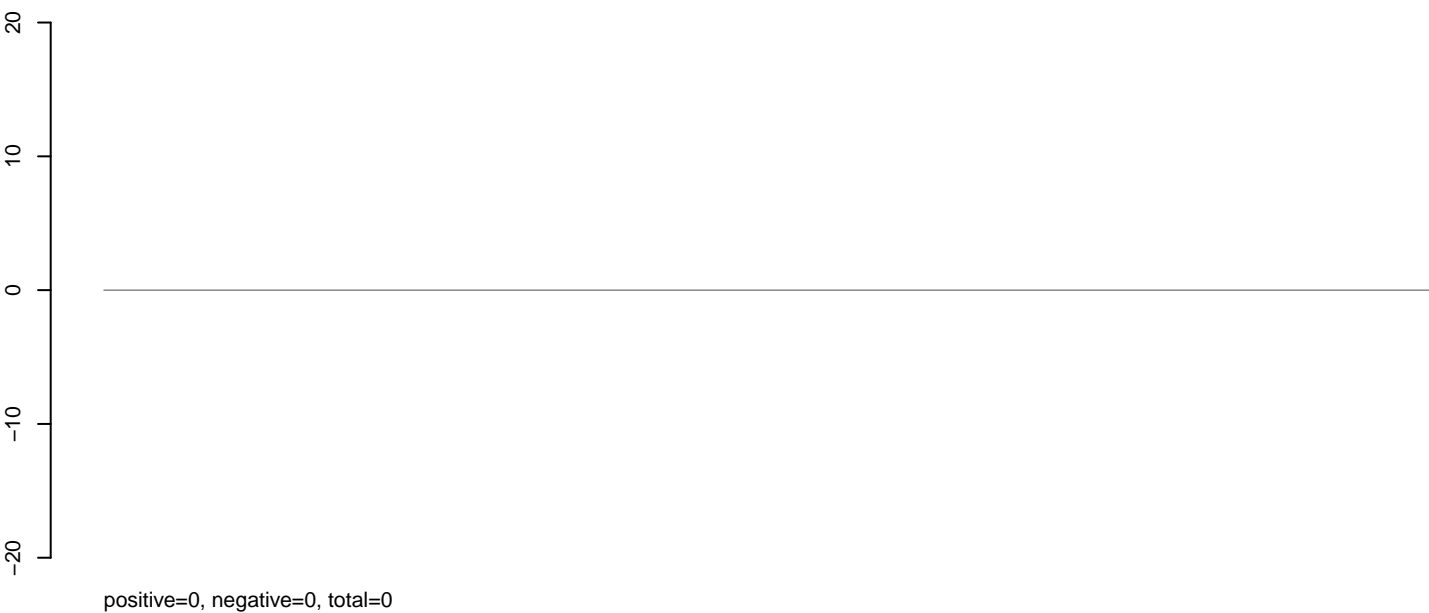
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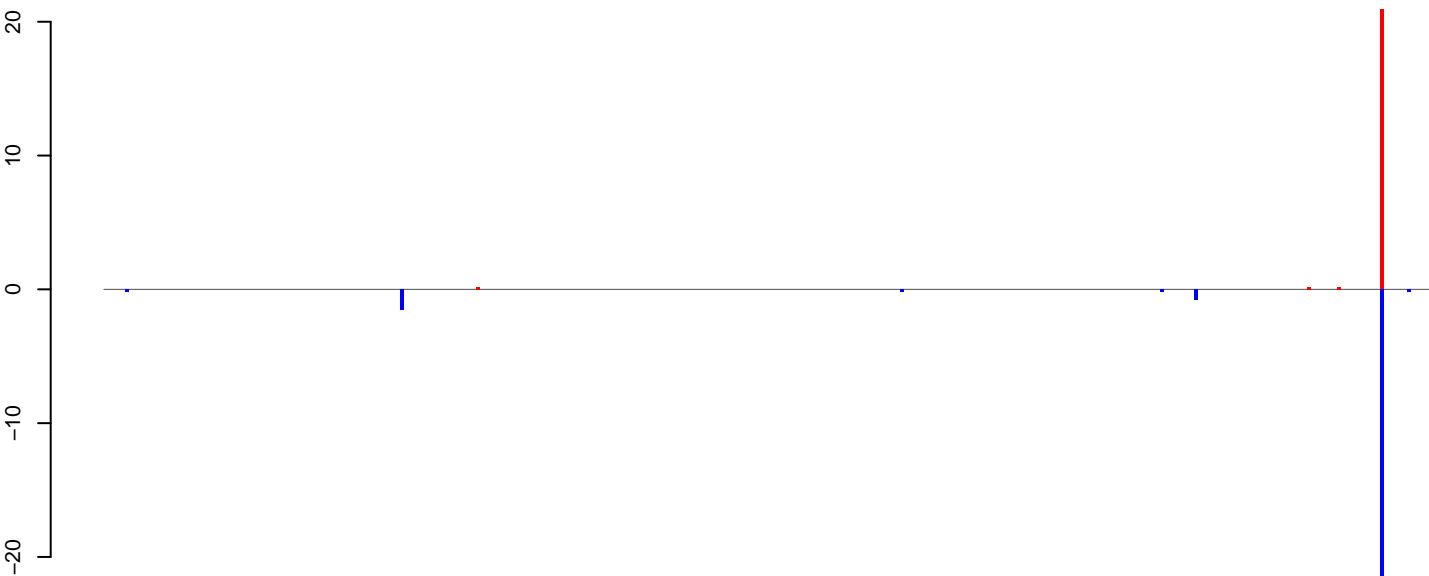
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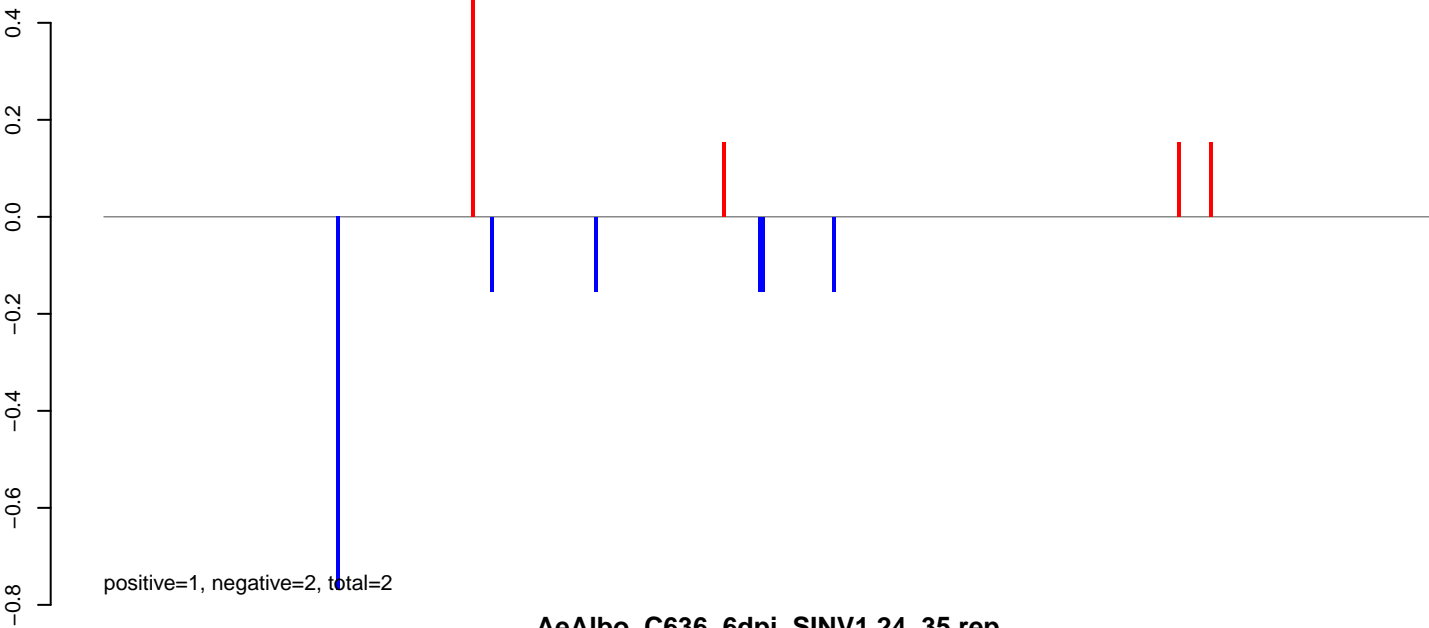
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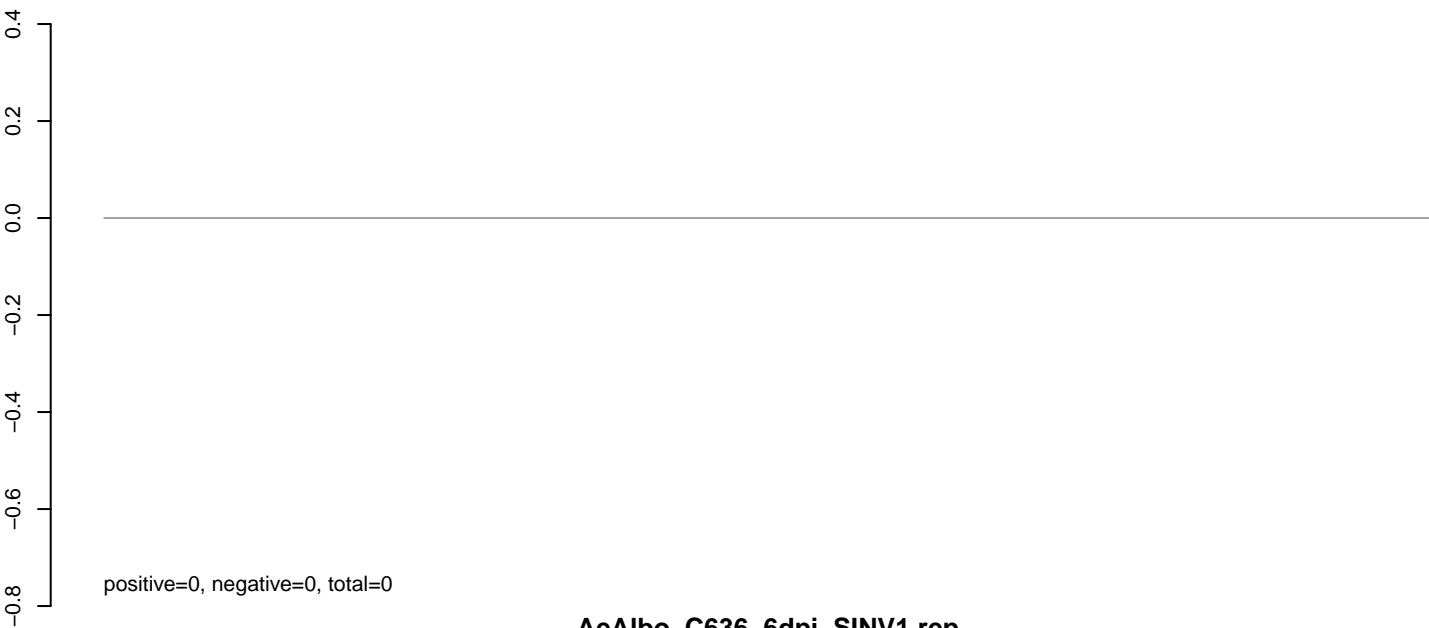
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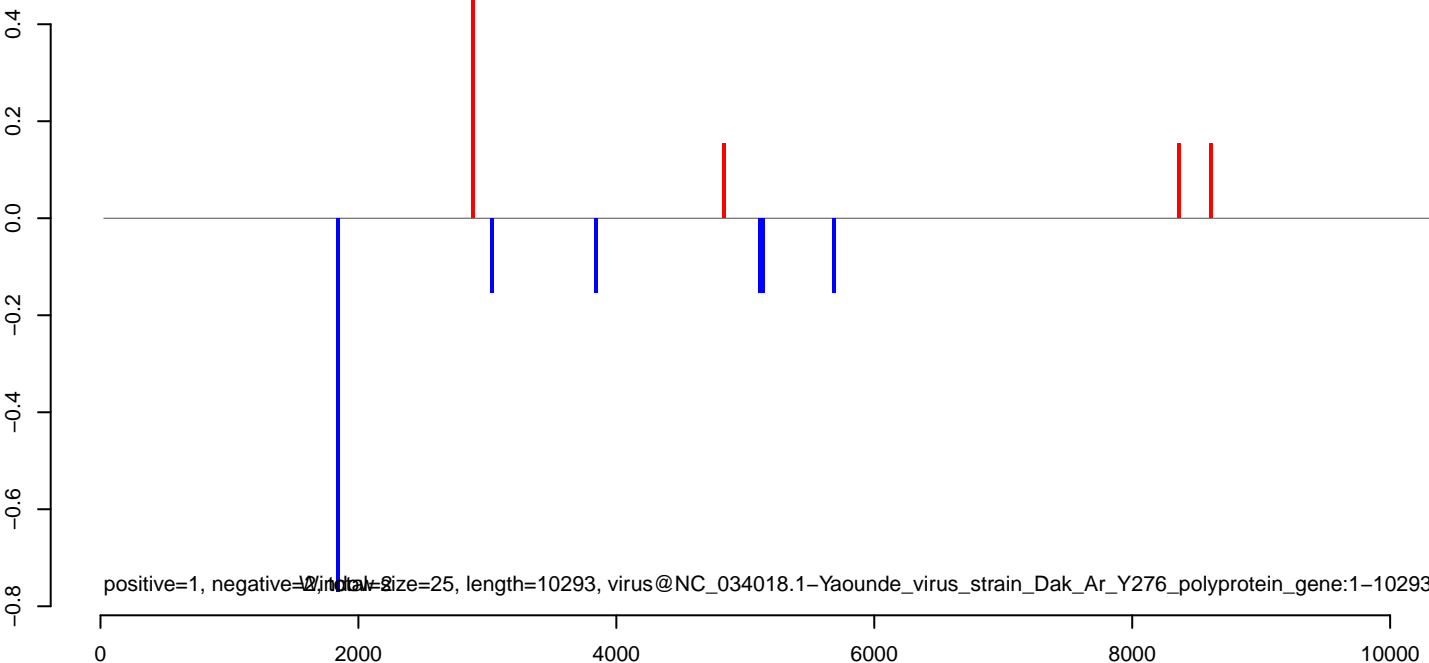
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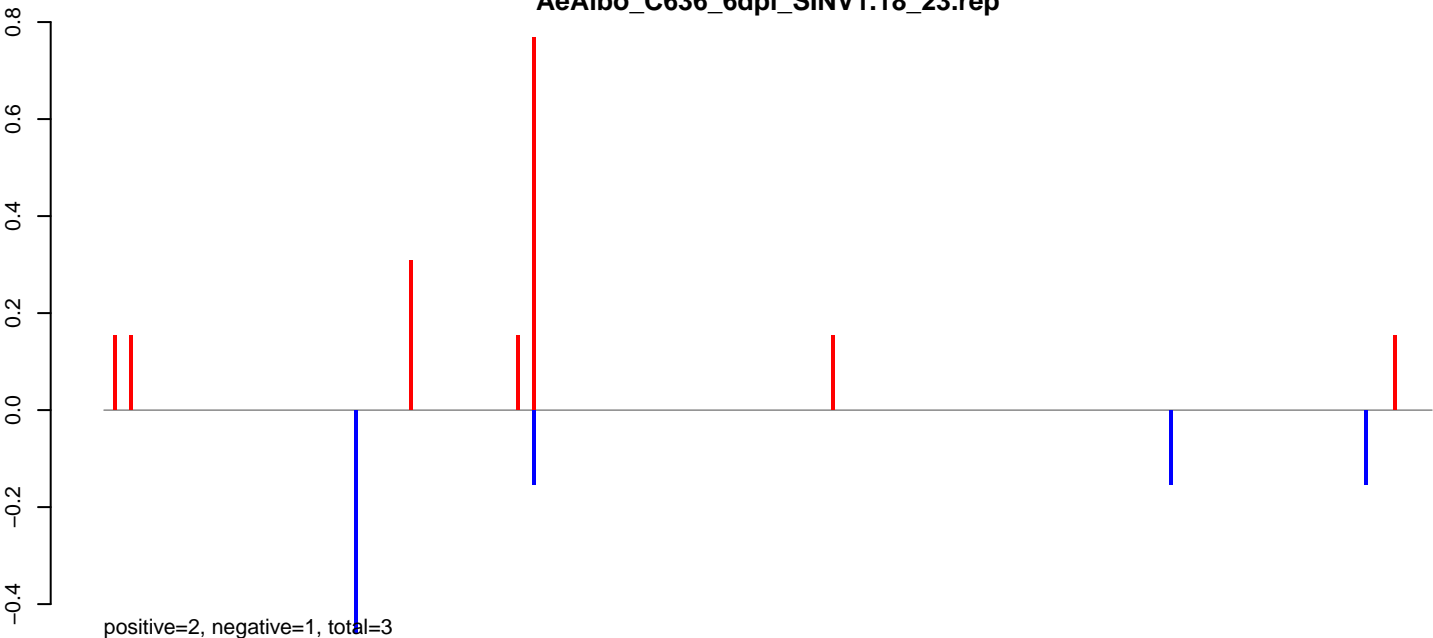
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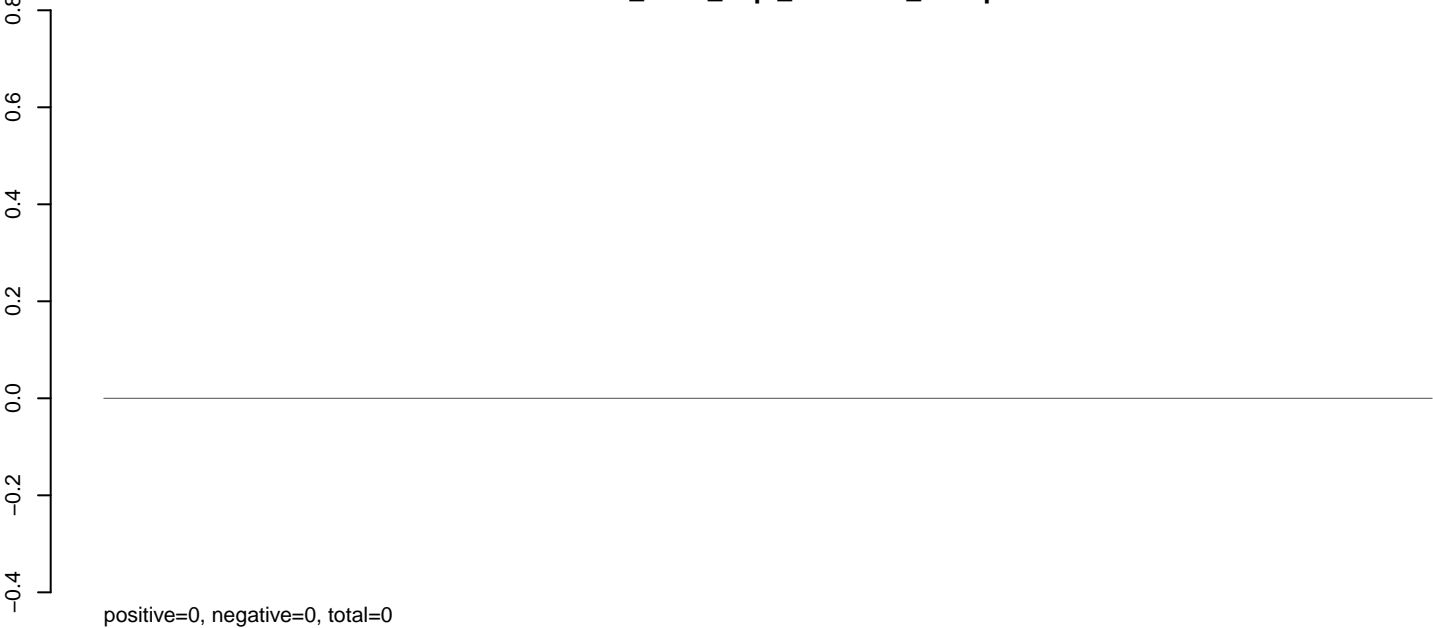
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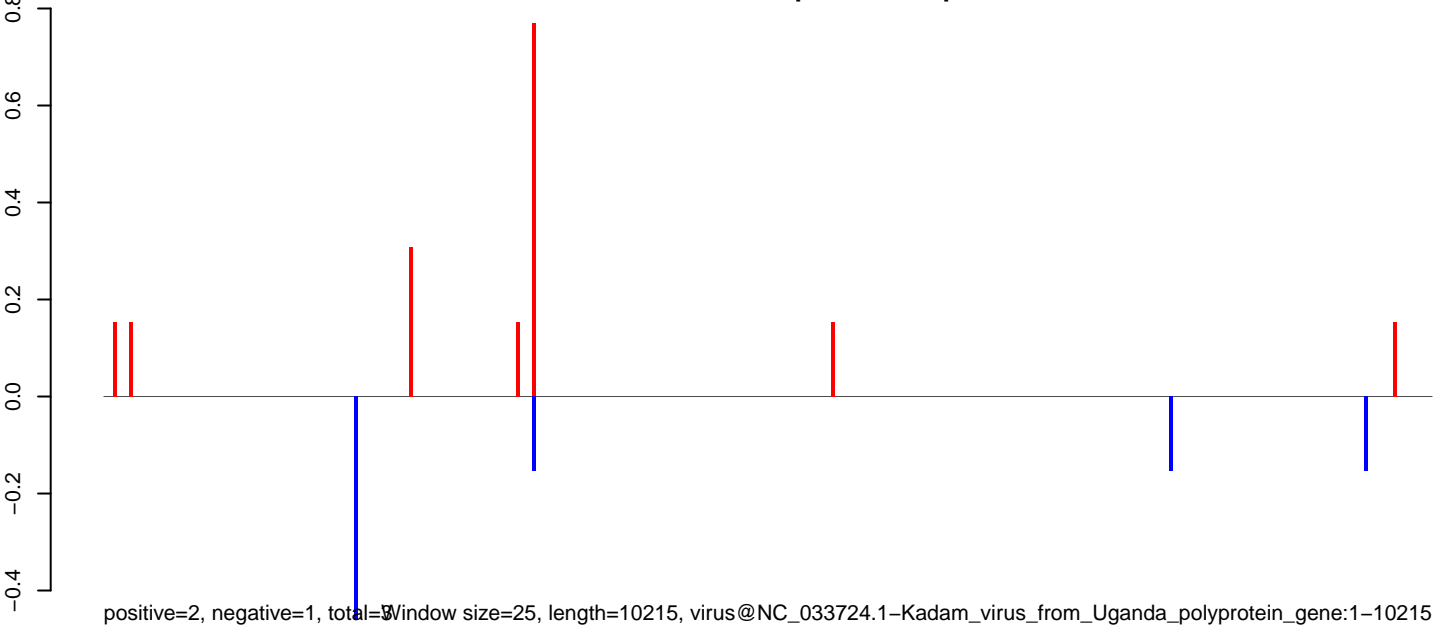
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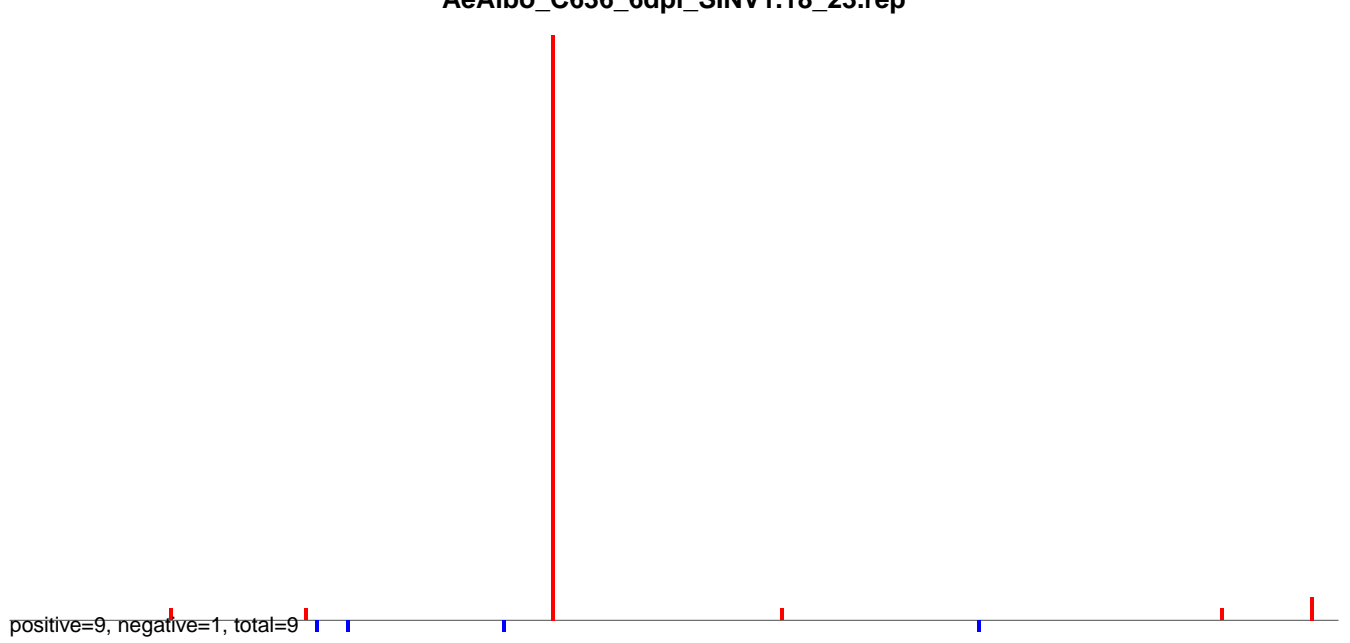
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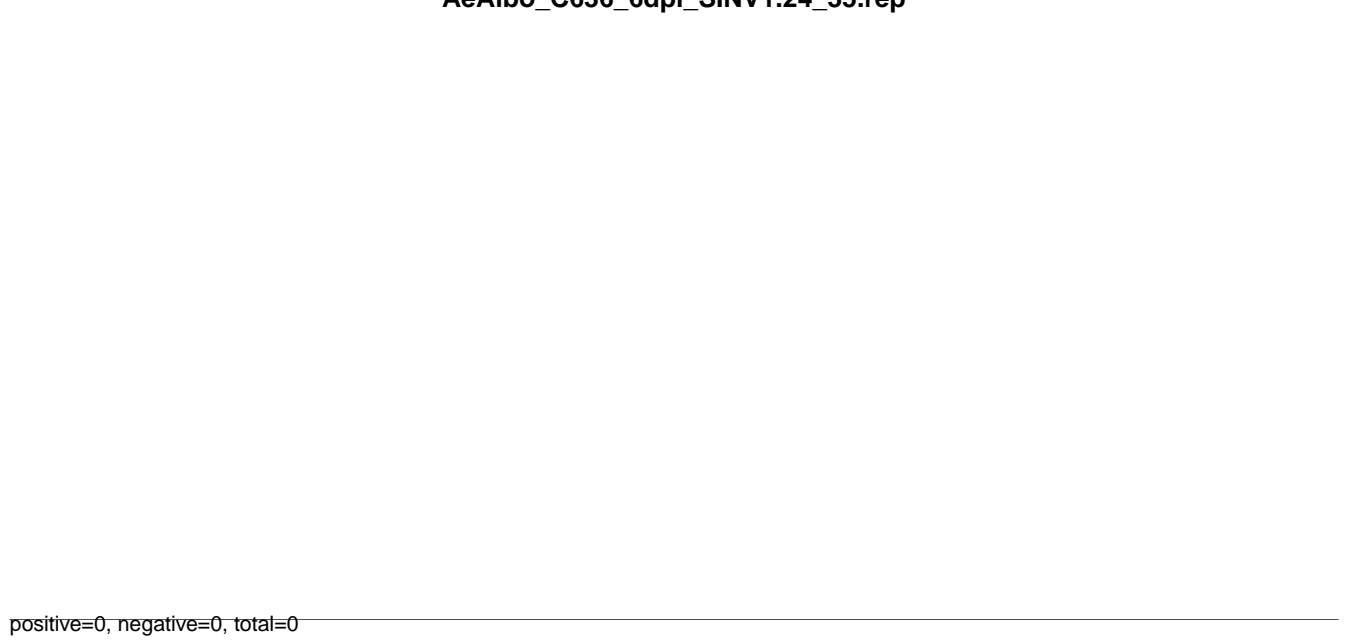
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AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep

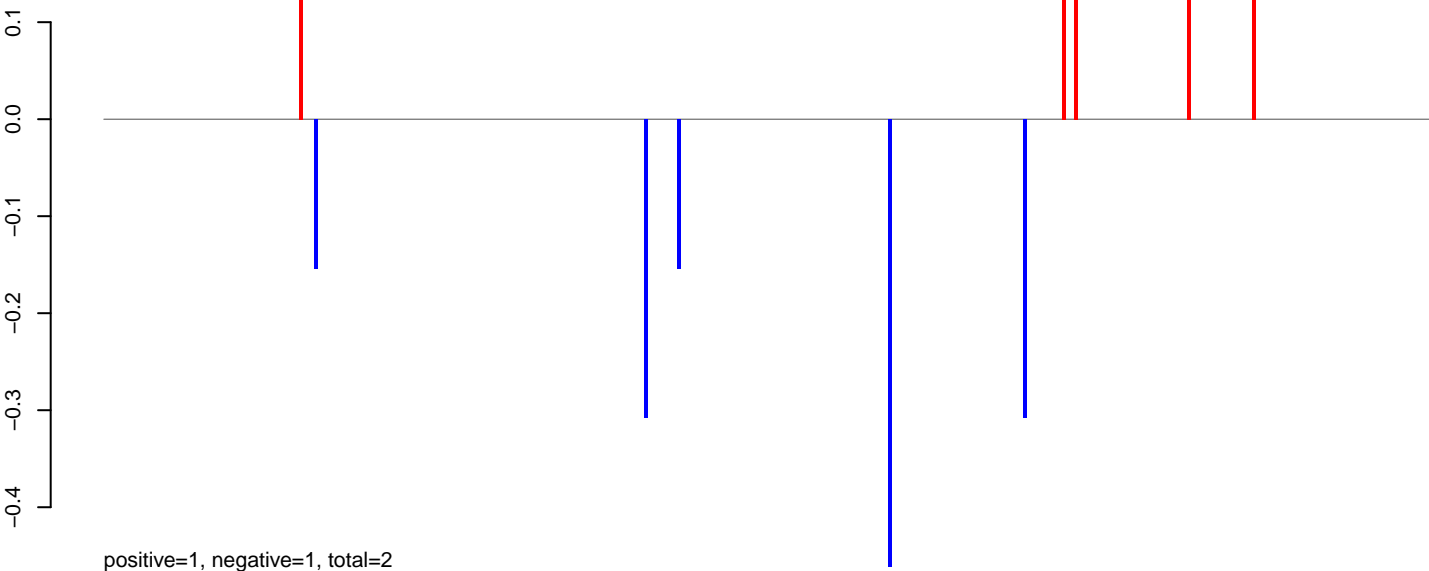


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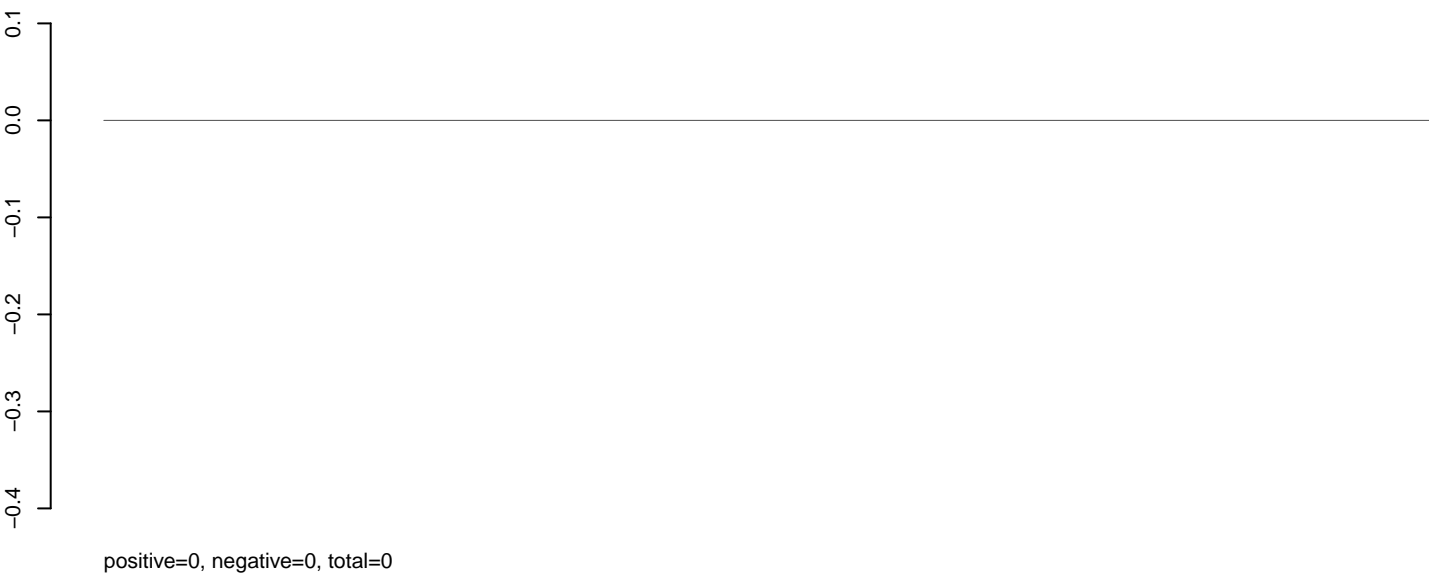


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AeAlbo_C636_6dpi_SINV1.24_35.rep
AeAlbo_C636_6dpi_SINV1.rep
virus@NC_033716.1_Zhejiang_mosquito_virus_1_strain_mosZJ35407_hypothetical_protein_gene:1-9558

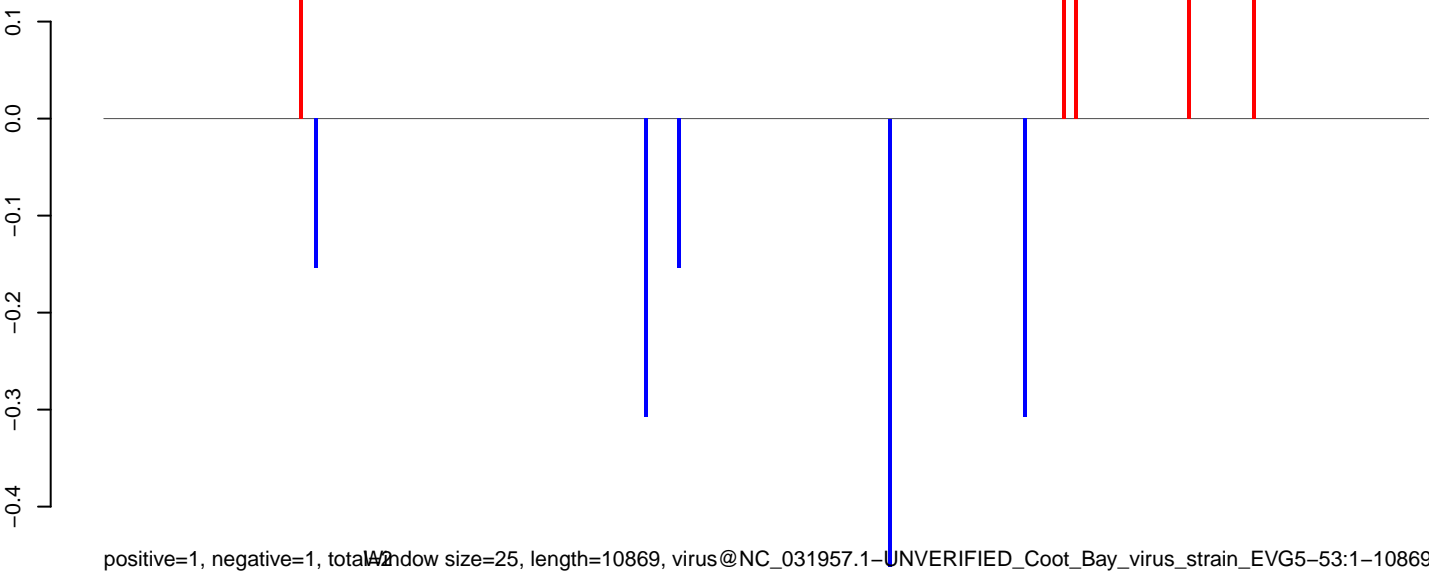
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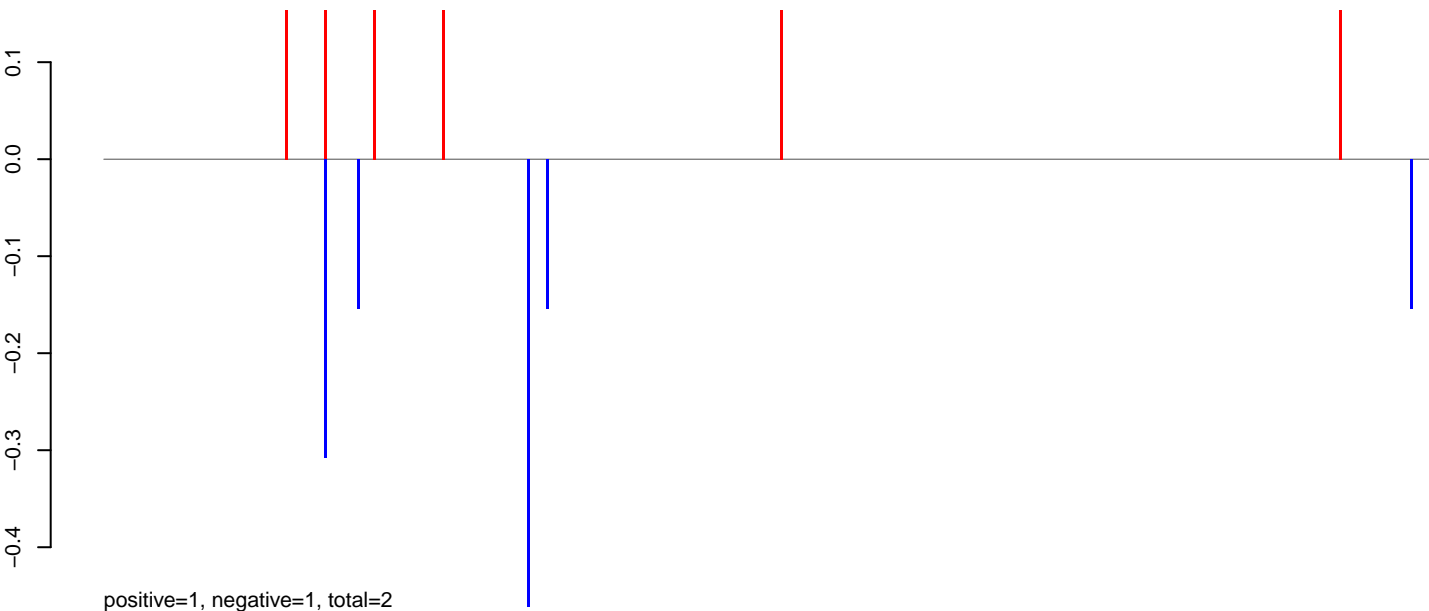
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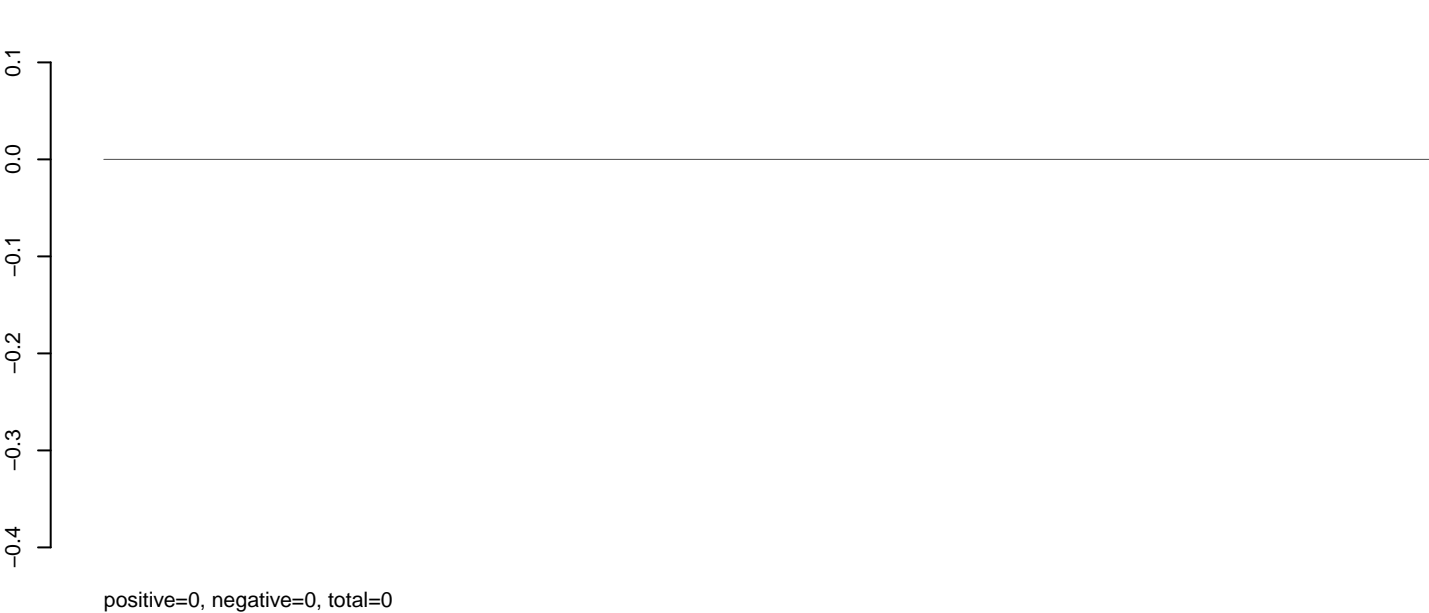
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0 2000 4000 6000 8000 10000

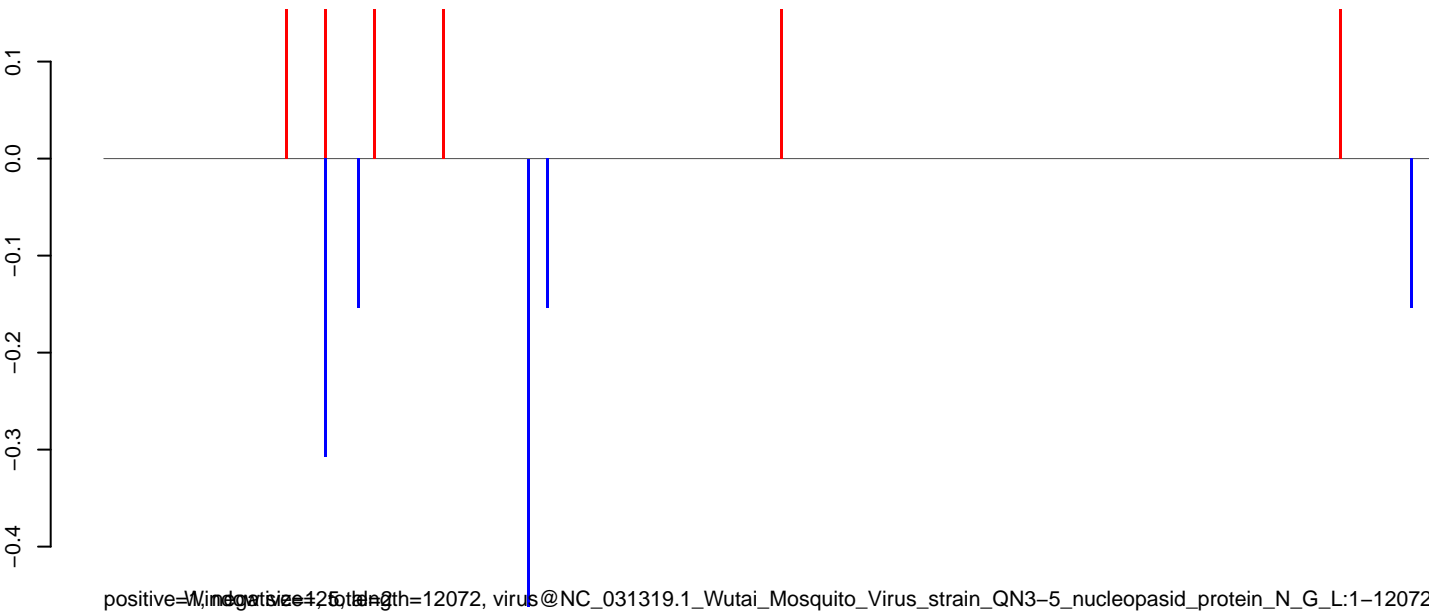
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AeAlbo_C636_6dpi_SINV1.24_35.rep



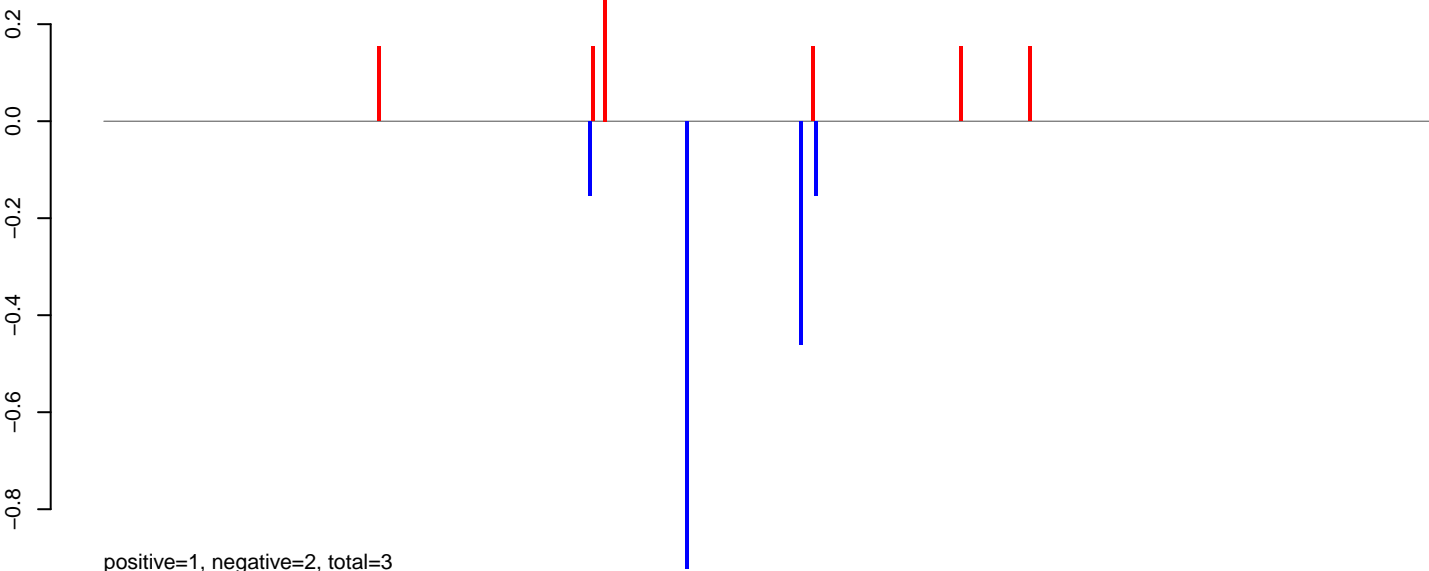
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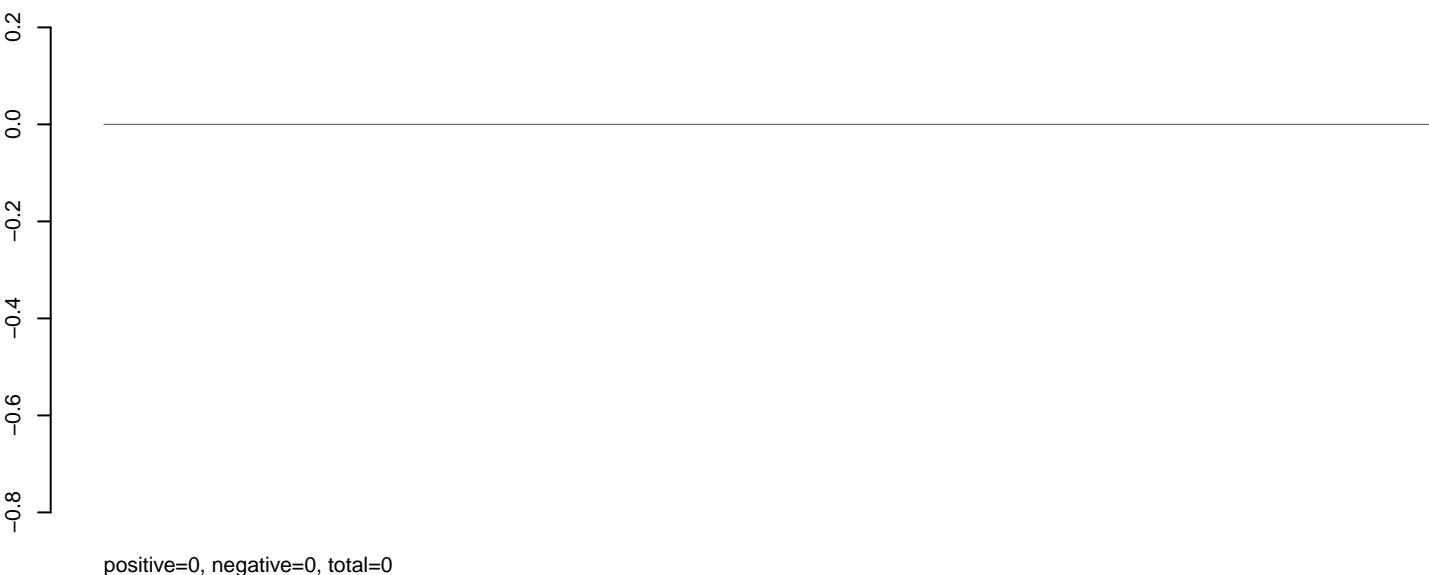
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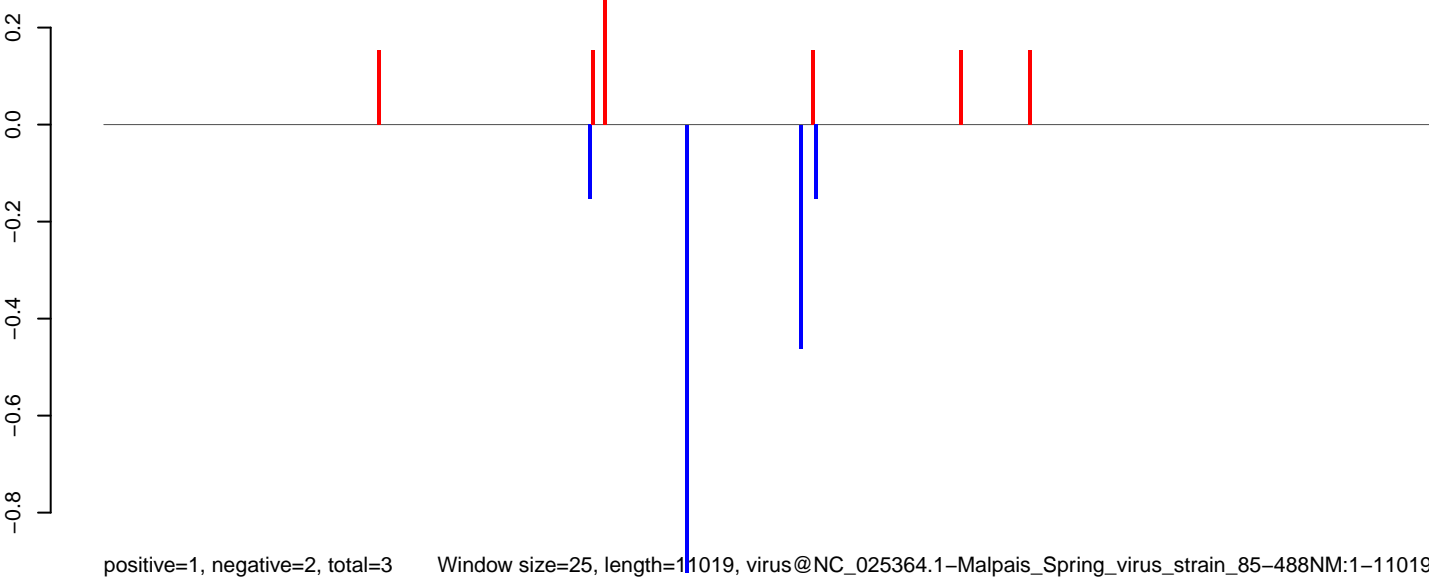
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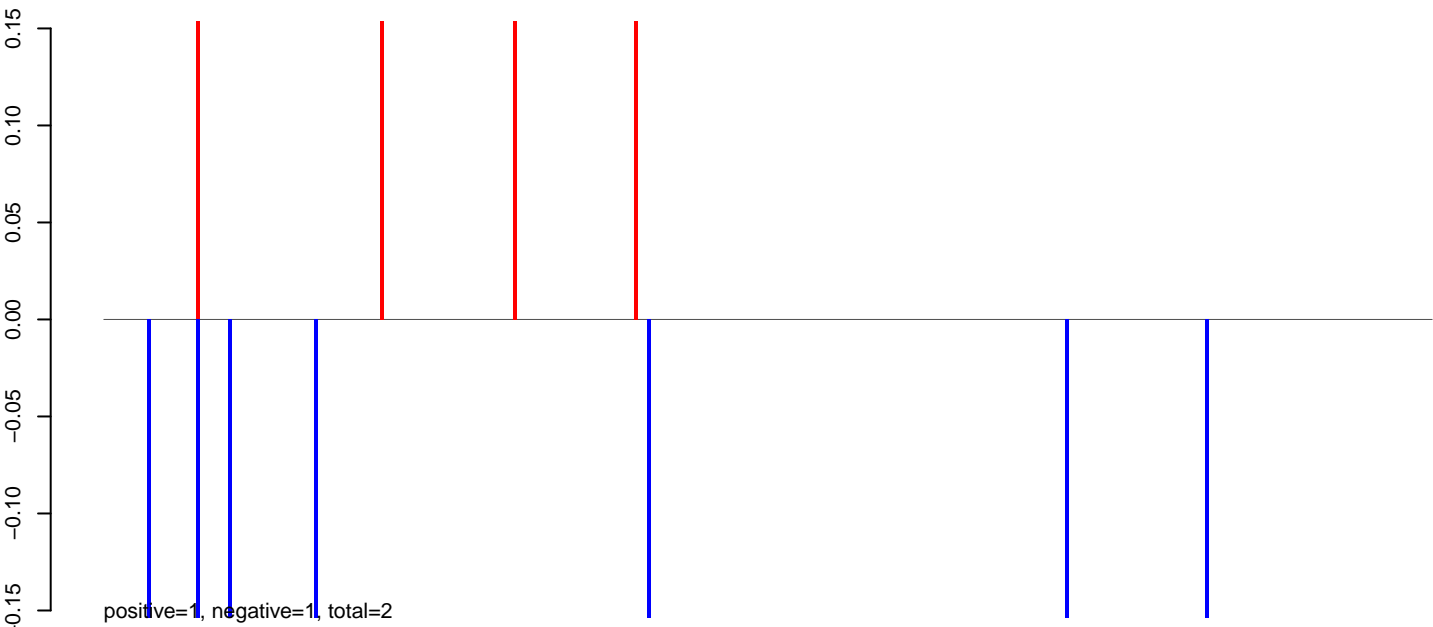
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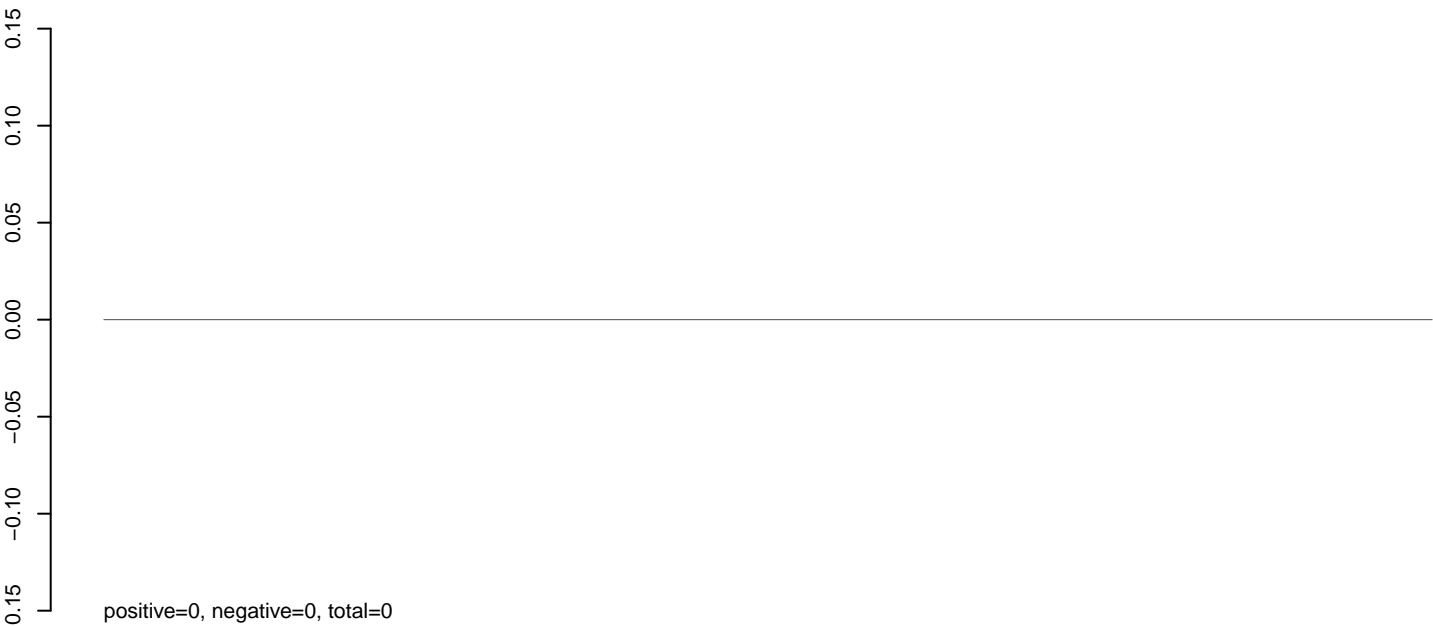
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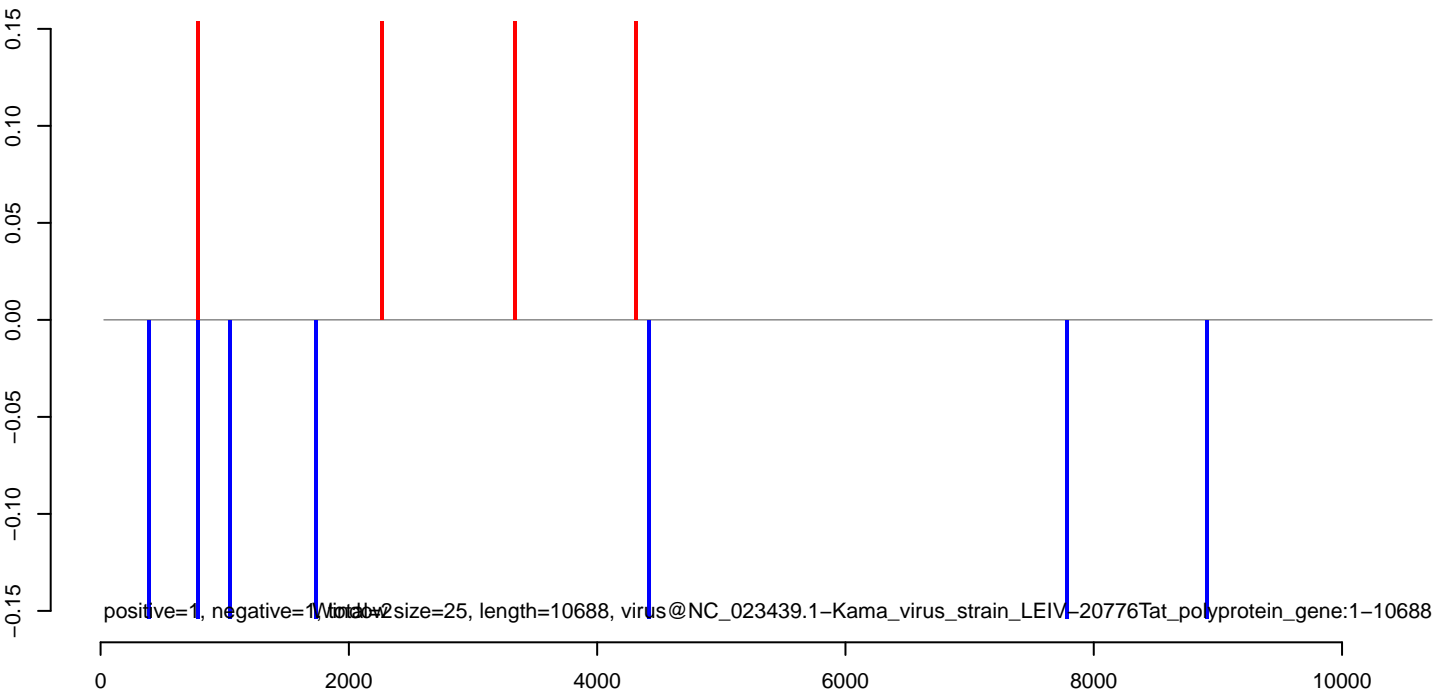
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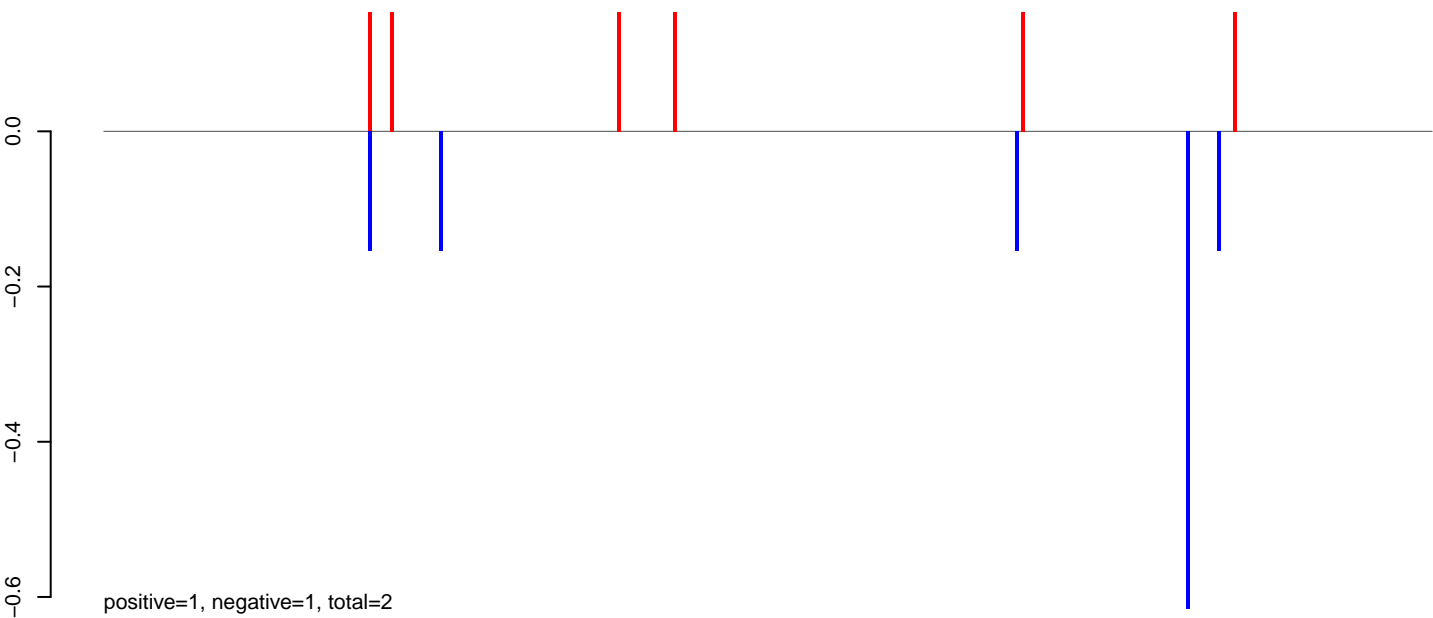
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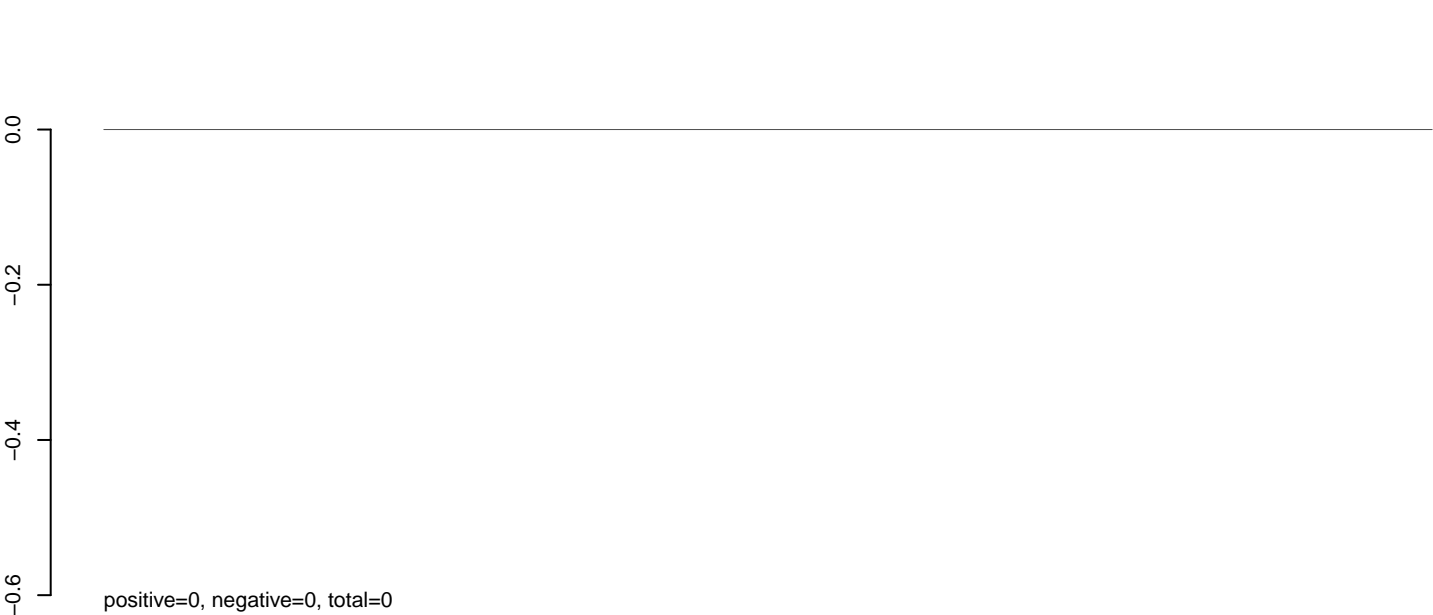
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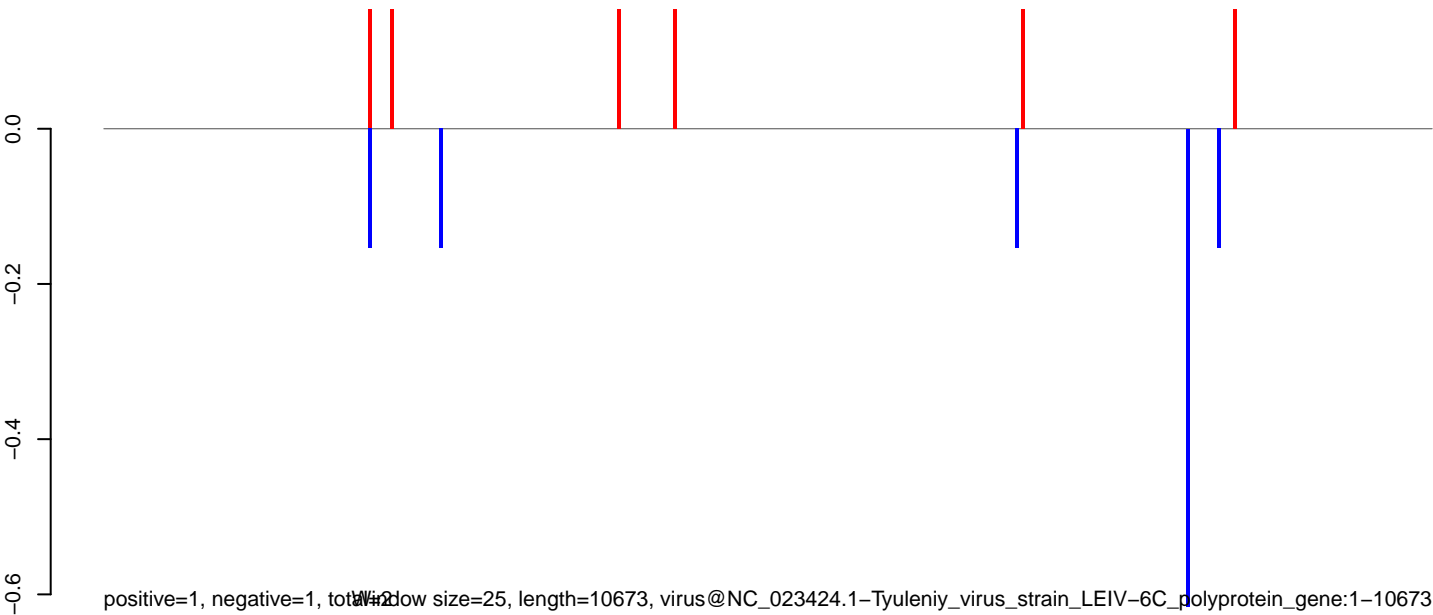
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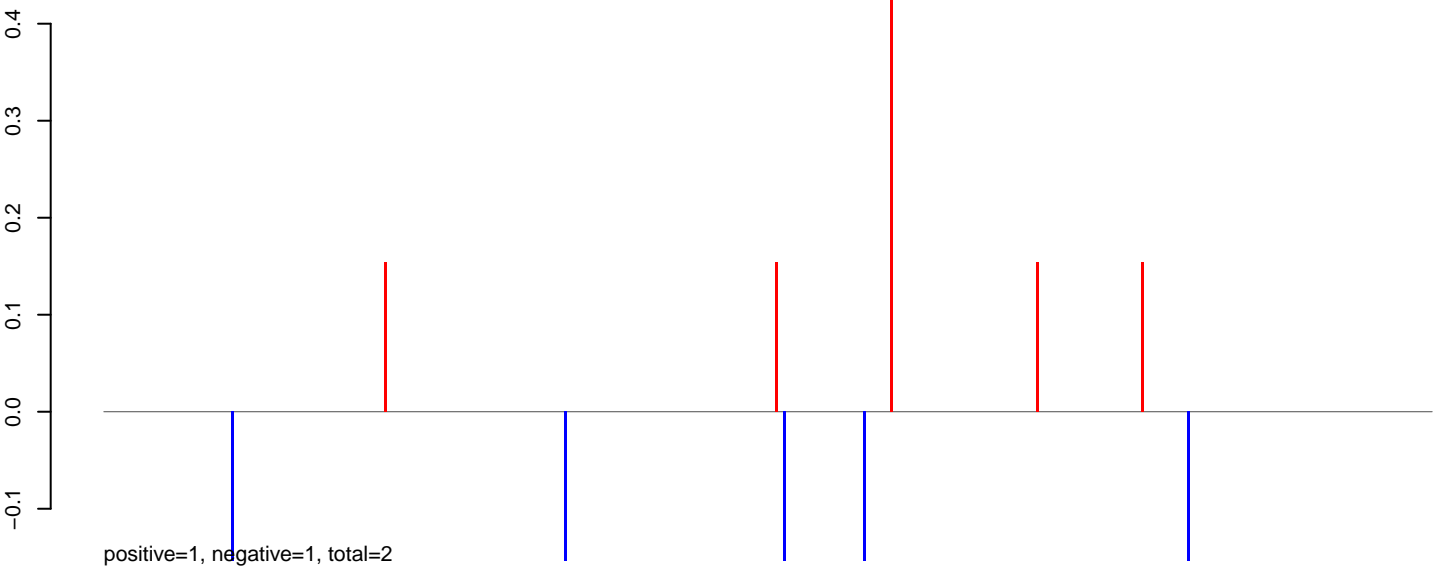
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W42 slow size=25, length=10673, virus@NC_023424.1-Tyuleniy_virus_strain_LEIV-6C_polyprotein_gene:1-10673

0 2000 4000 6000 8000 10000

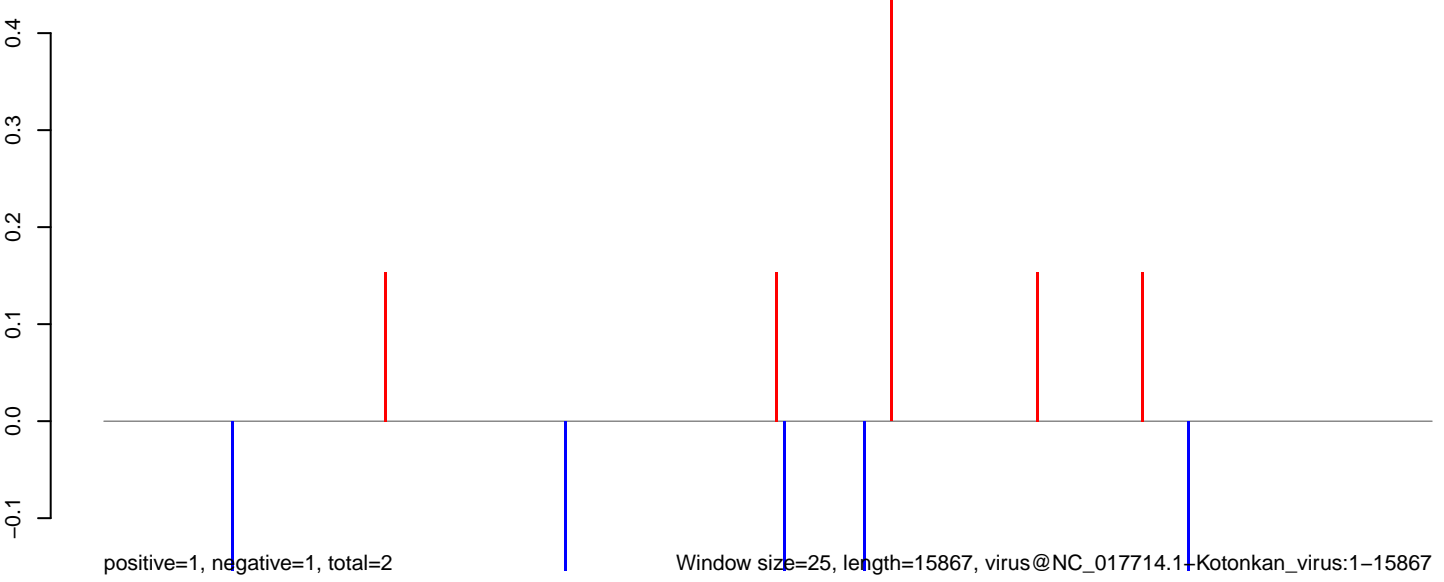
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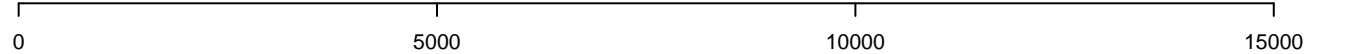
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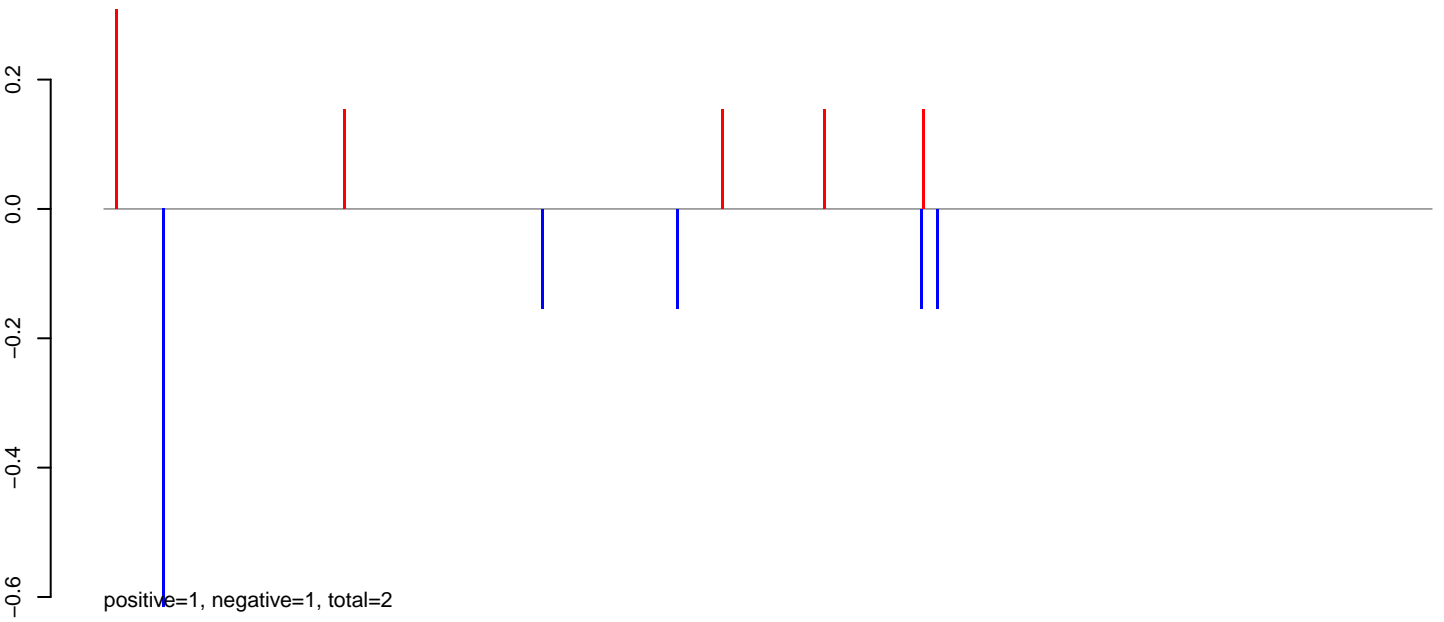
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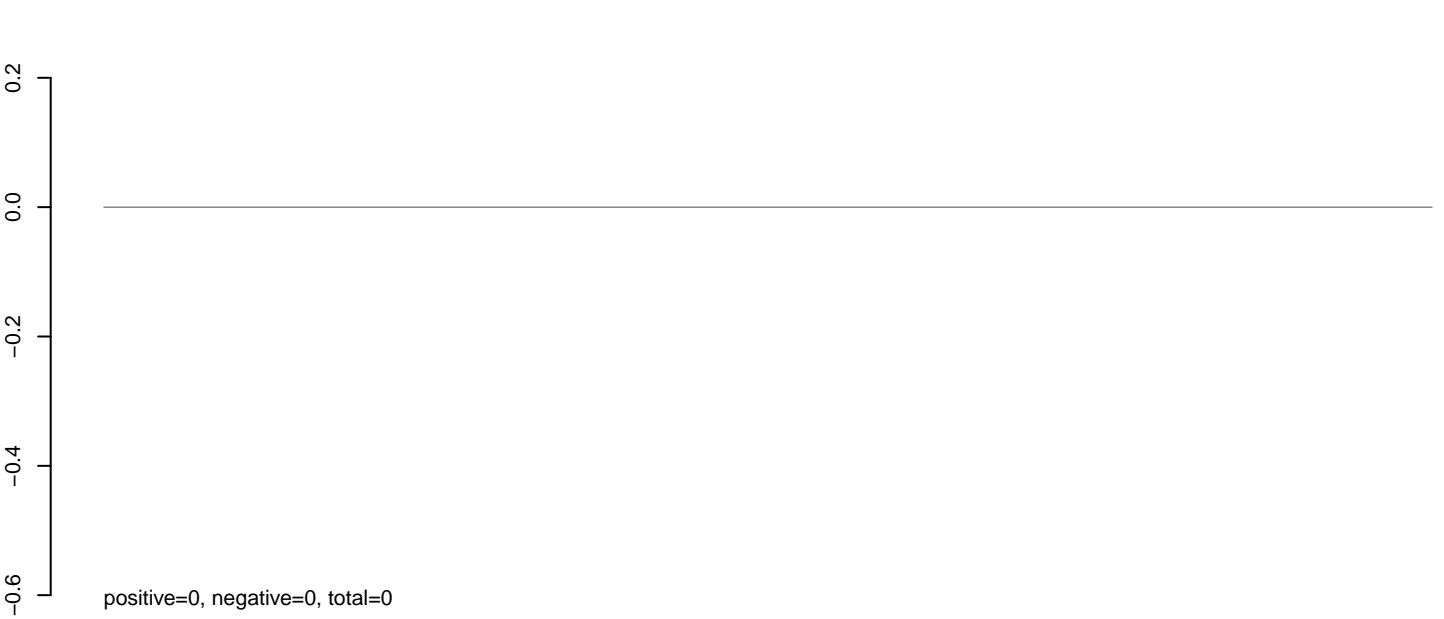
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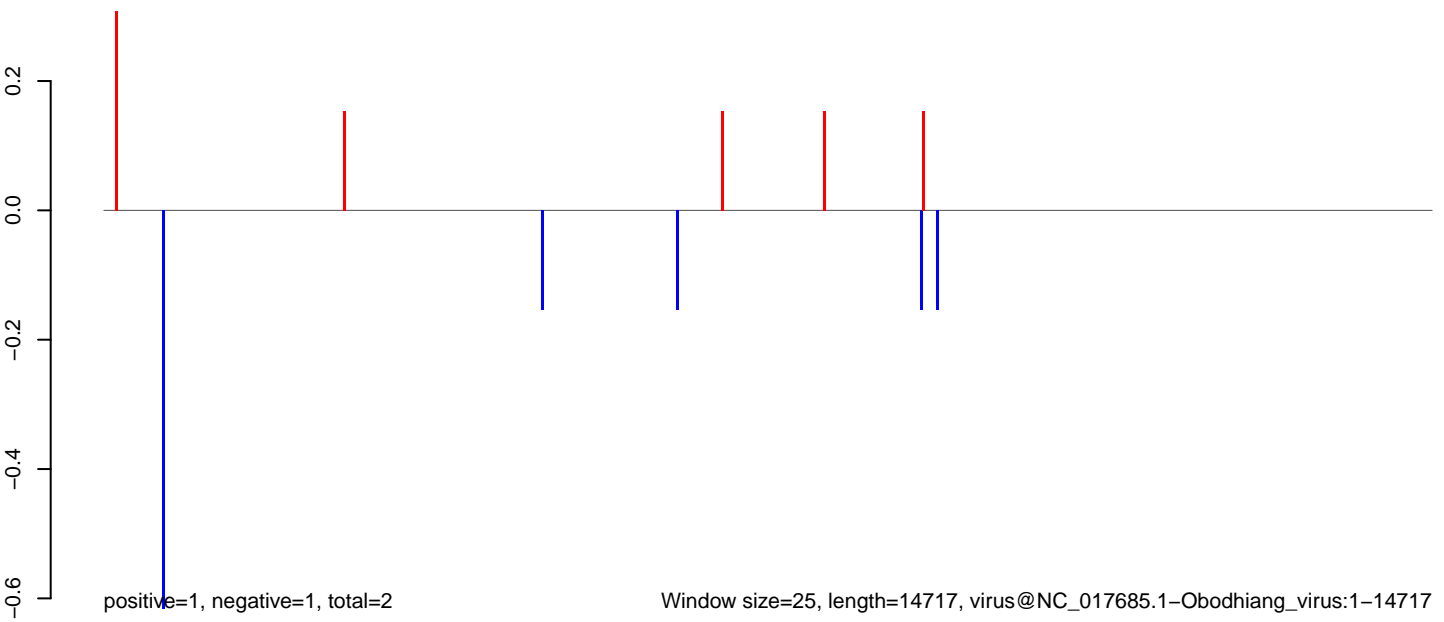
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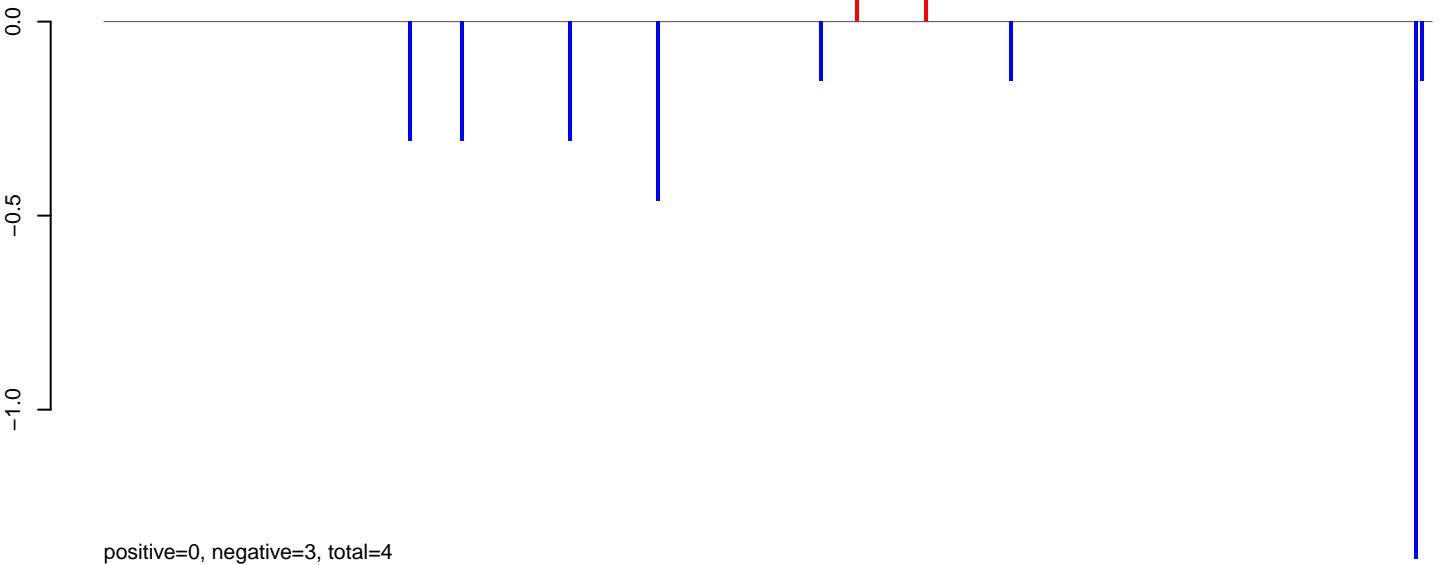


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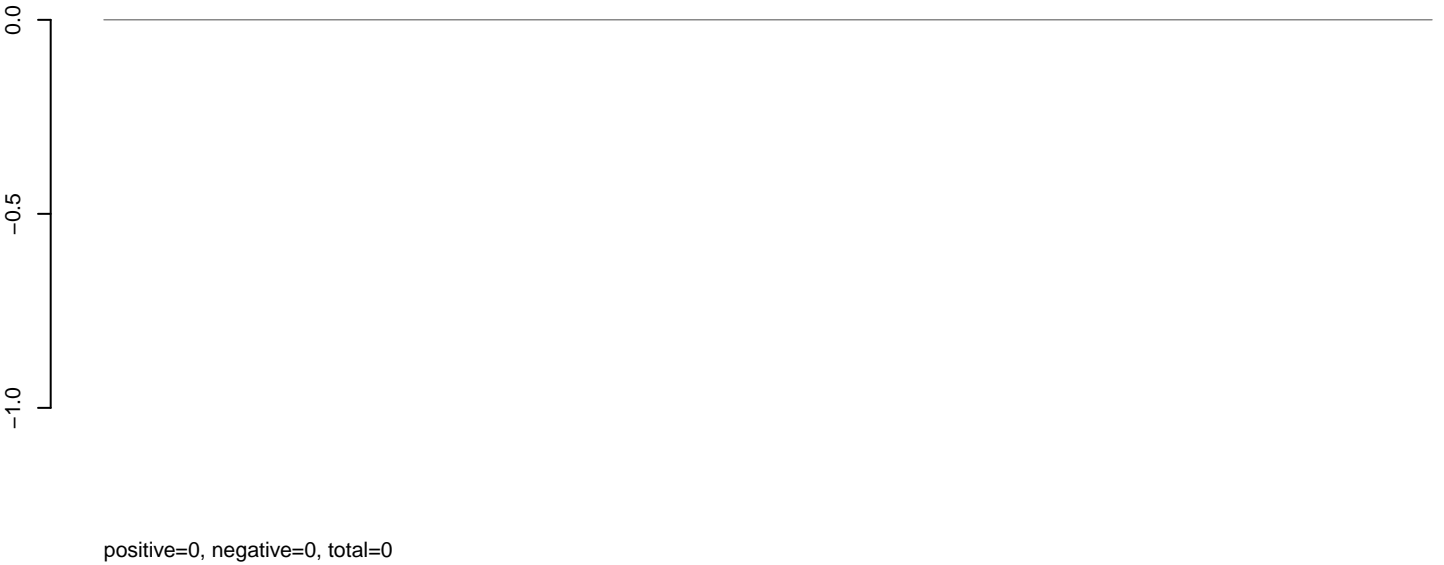


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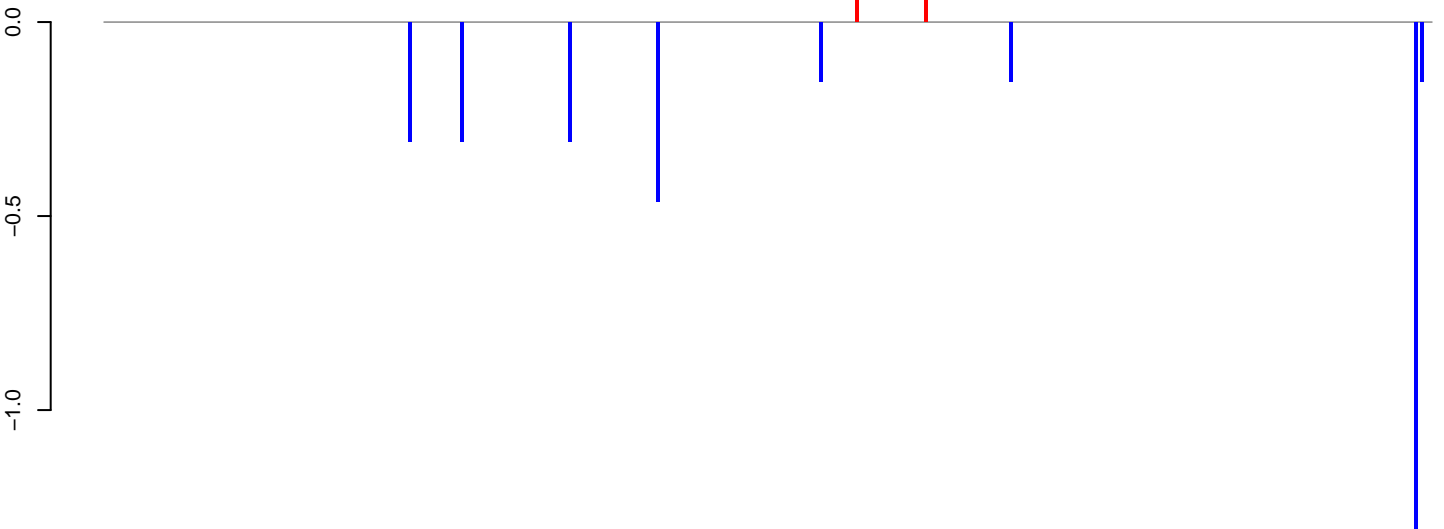
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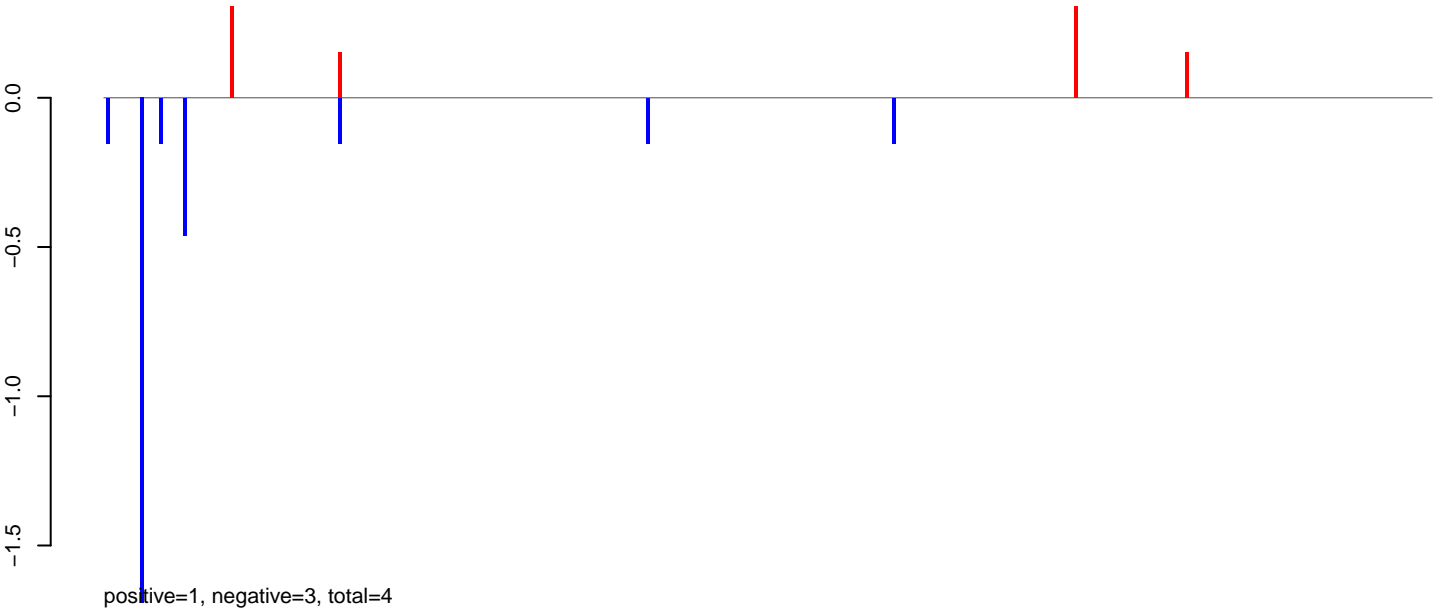
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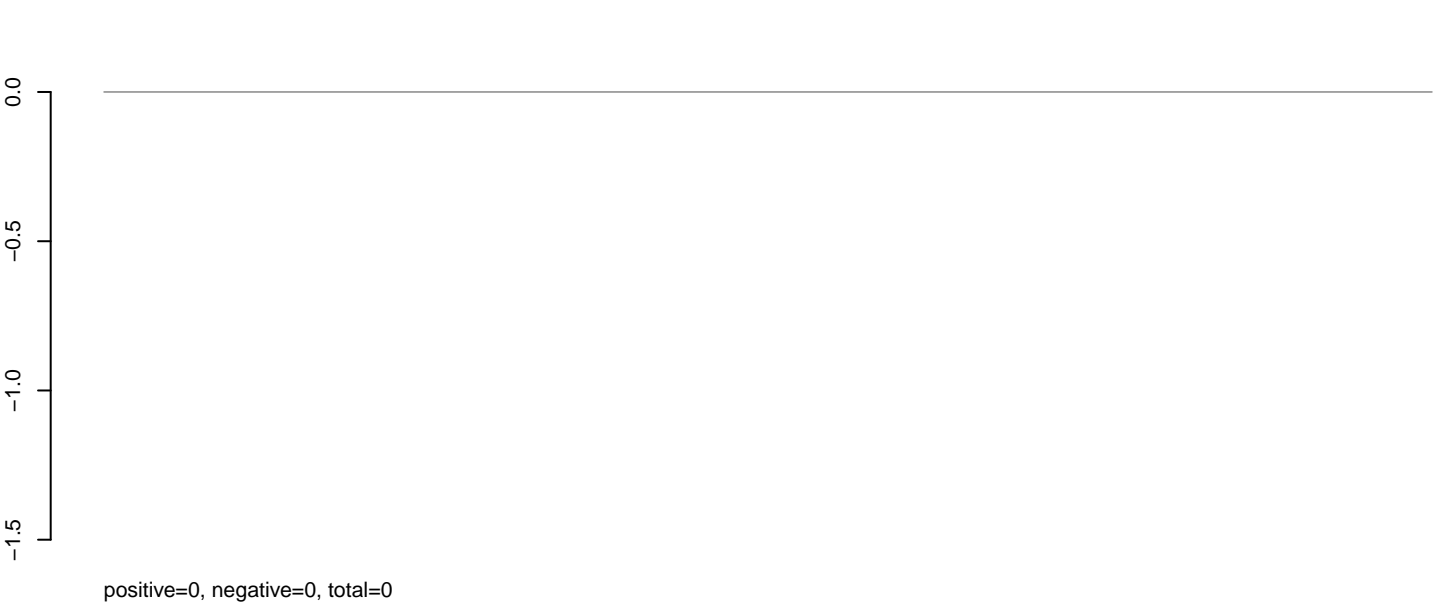
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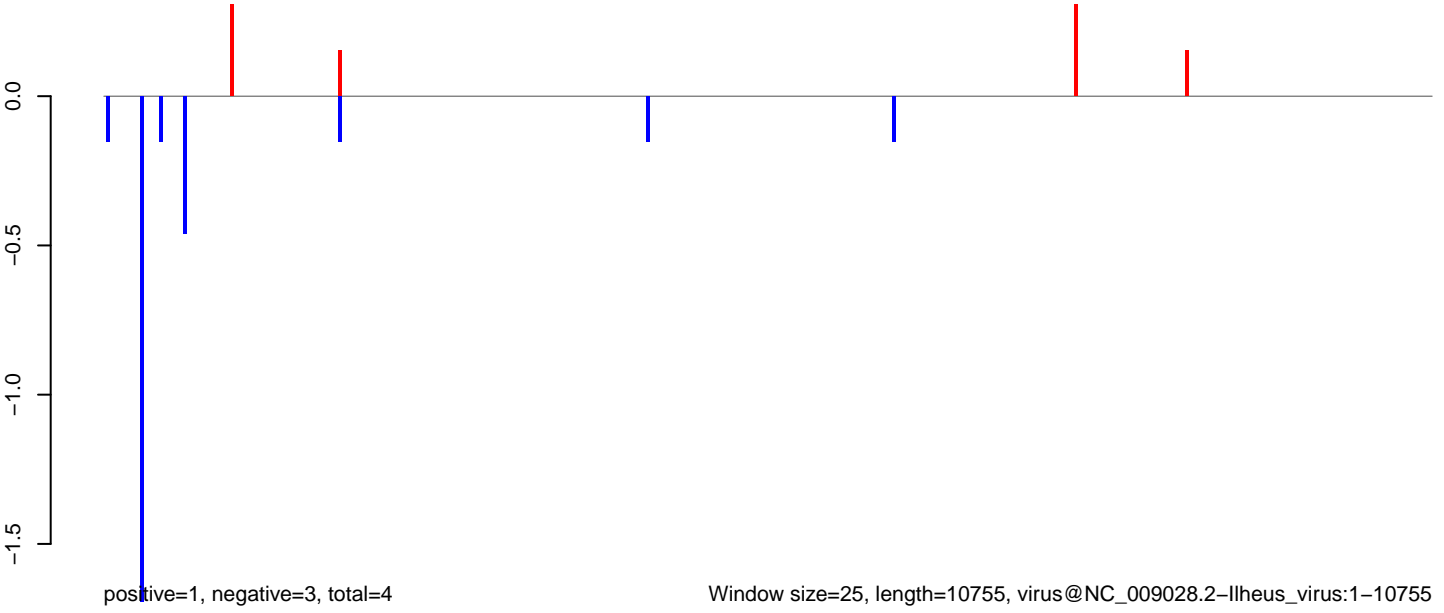
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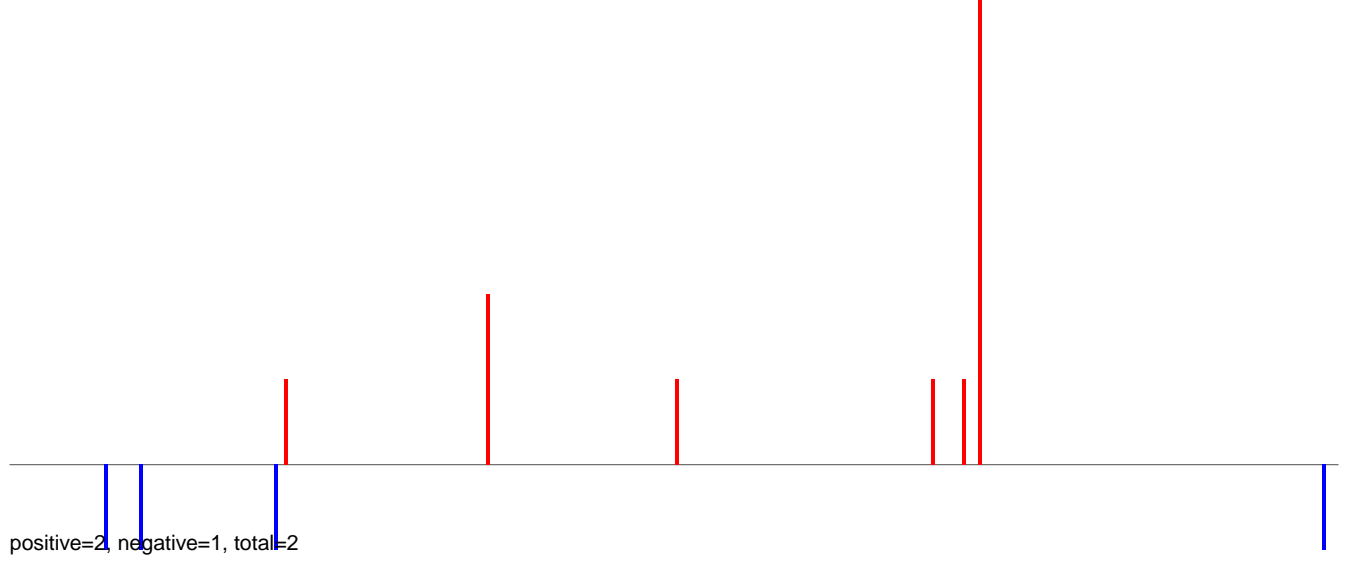
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AeAlbo_C636_6dpi_SINV1.rep



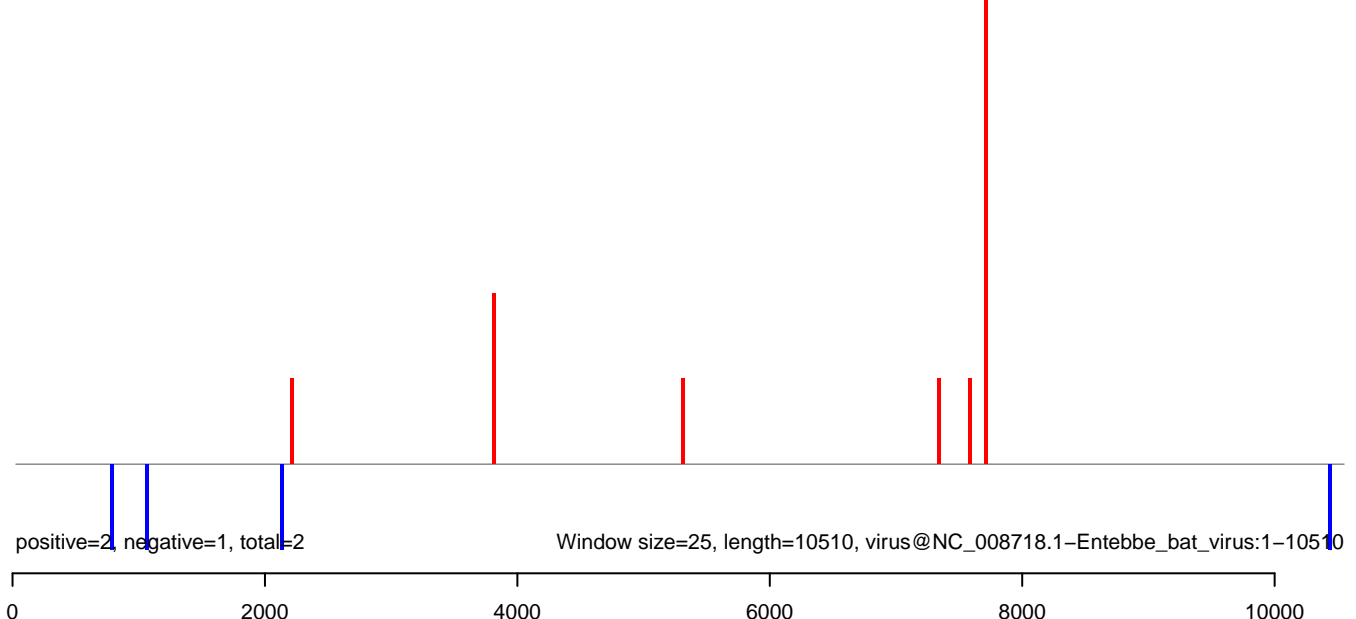
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AeAlbo_C636_6dpi_SINV1.24_35.rep



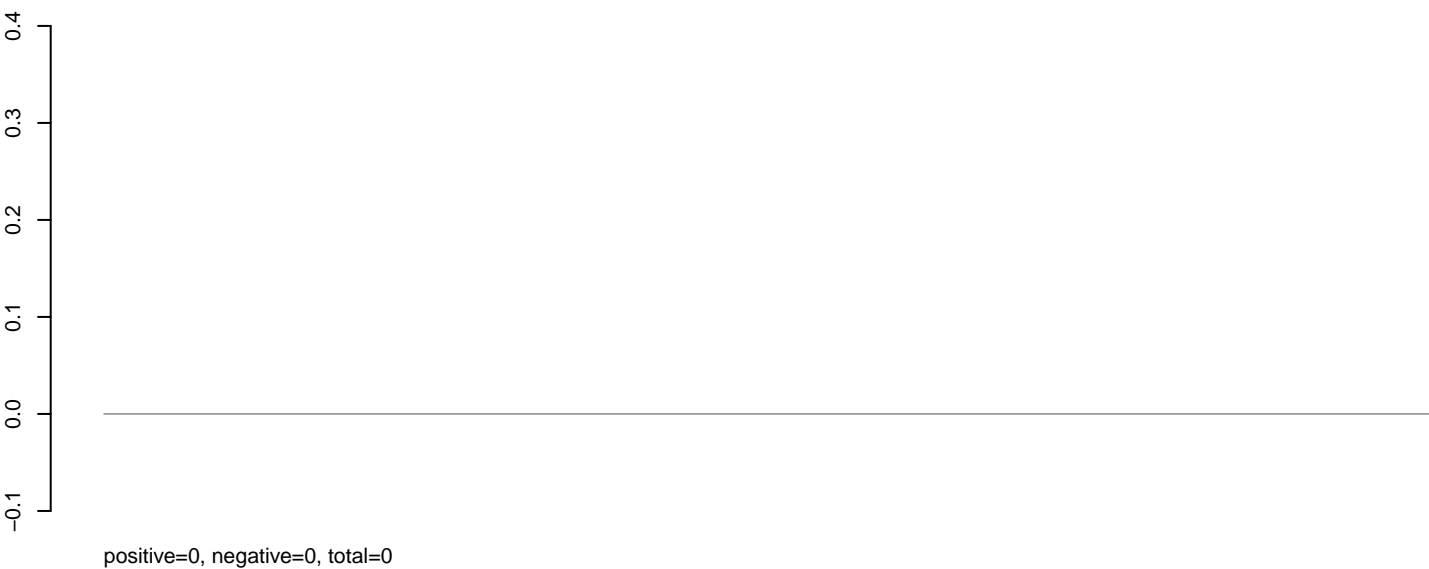
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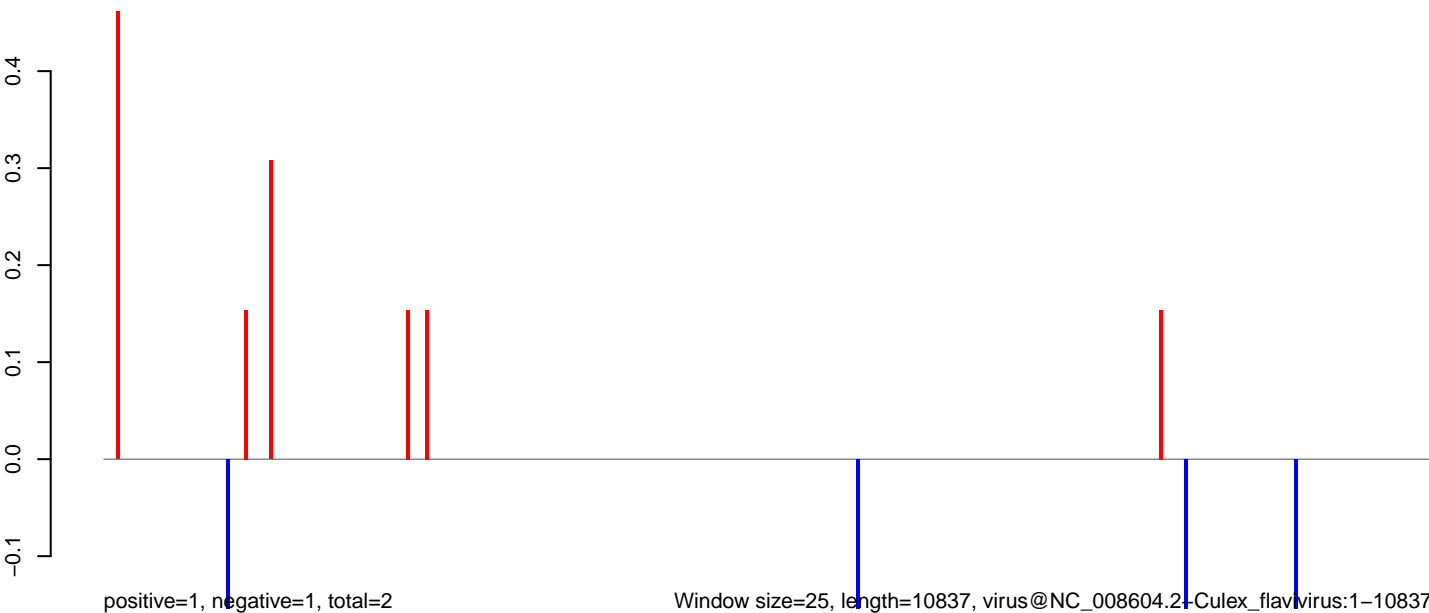
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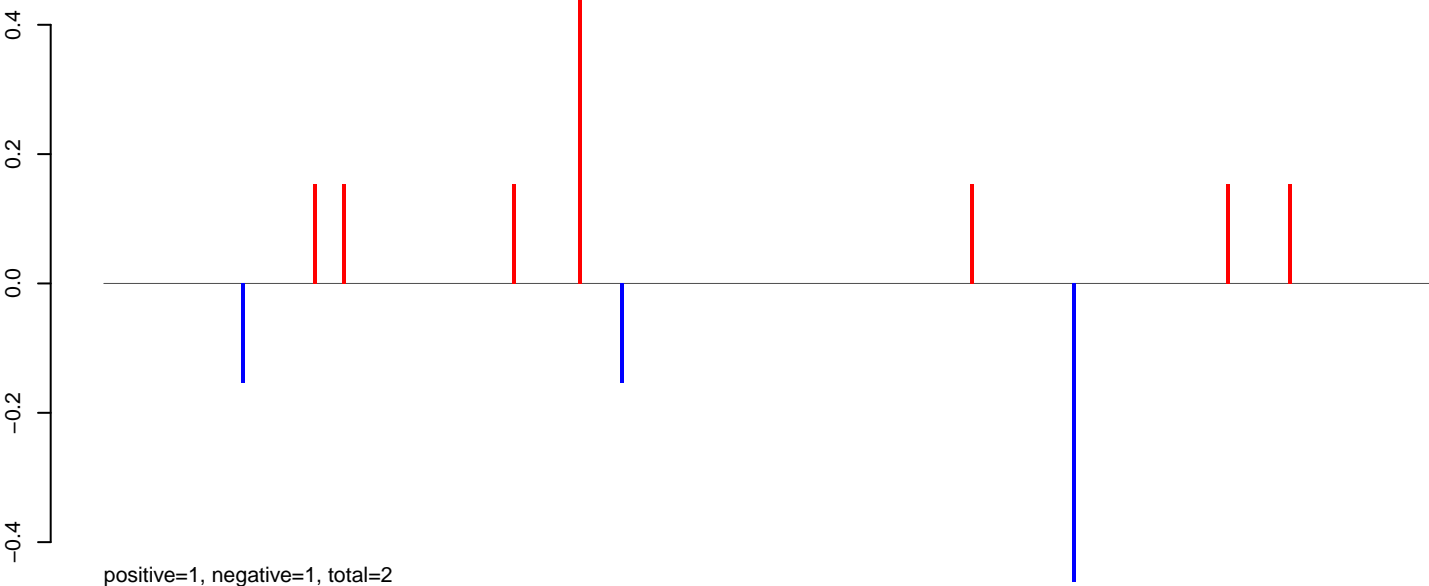
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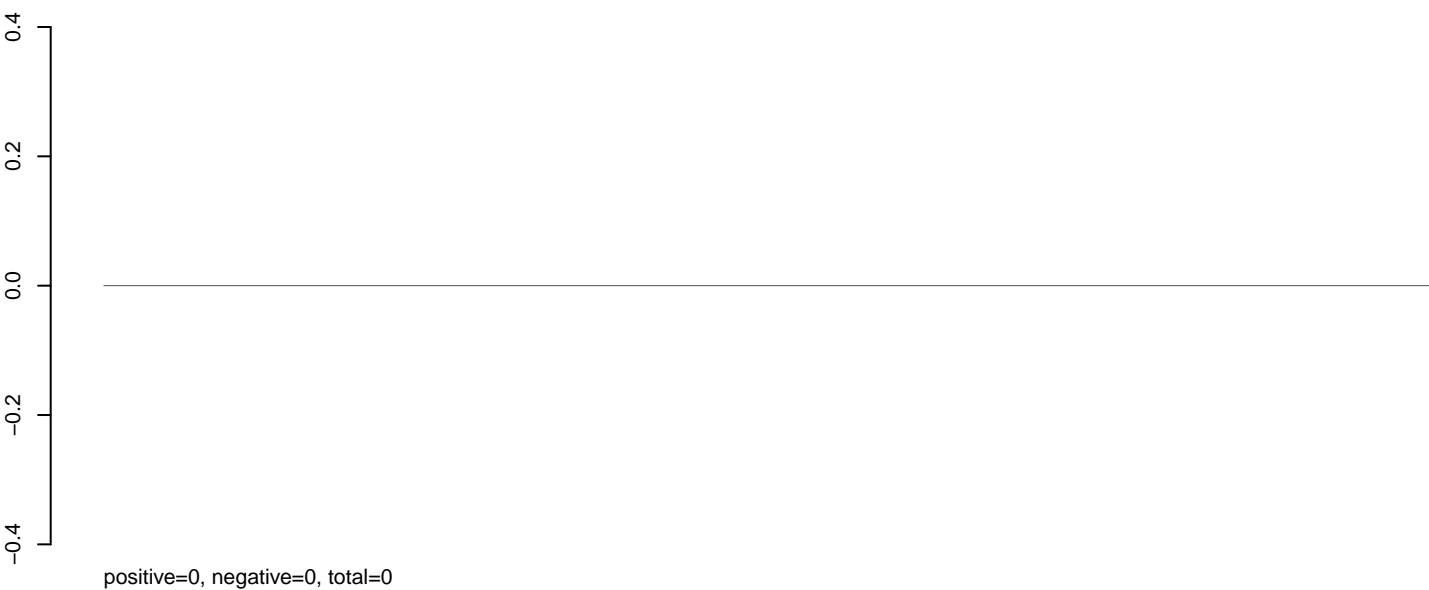
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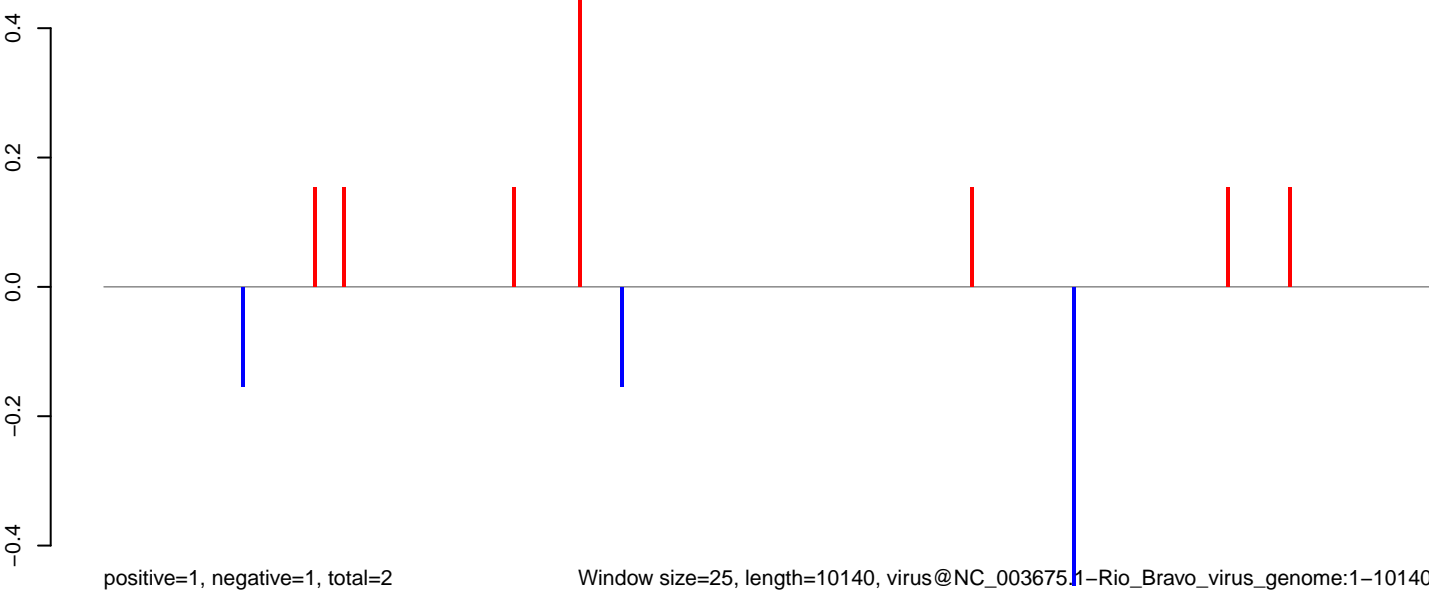
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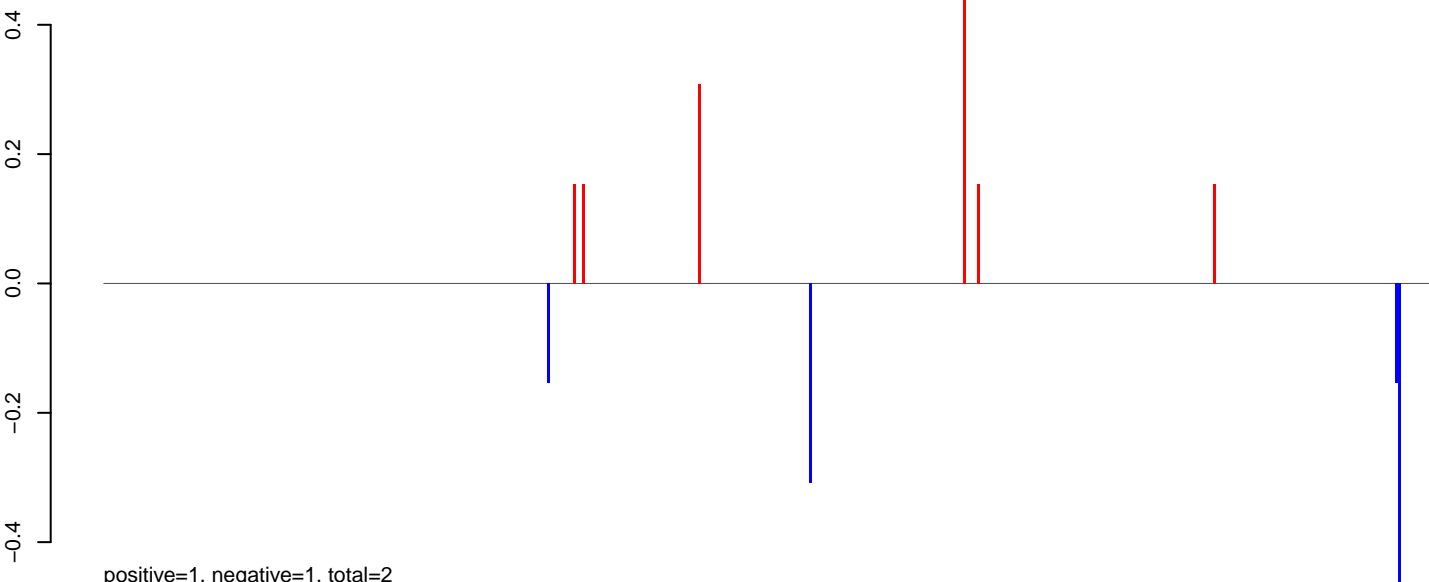
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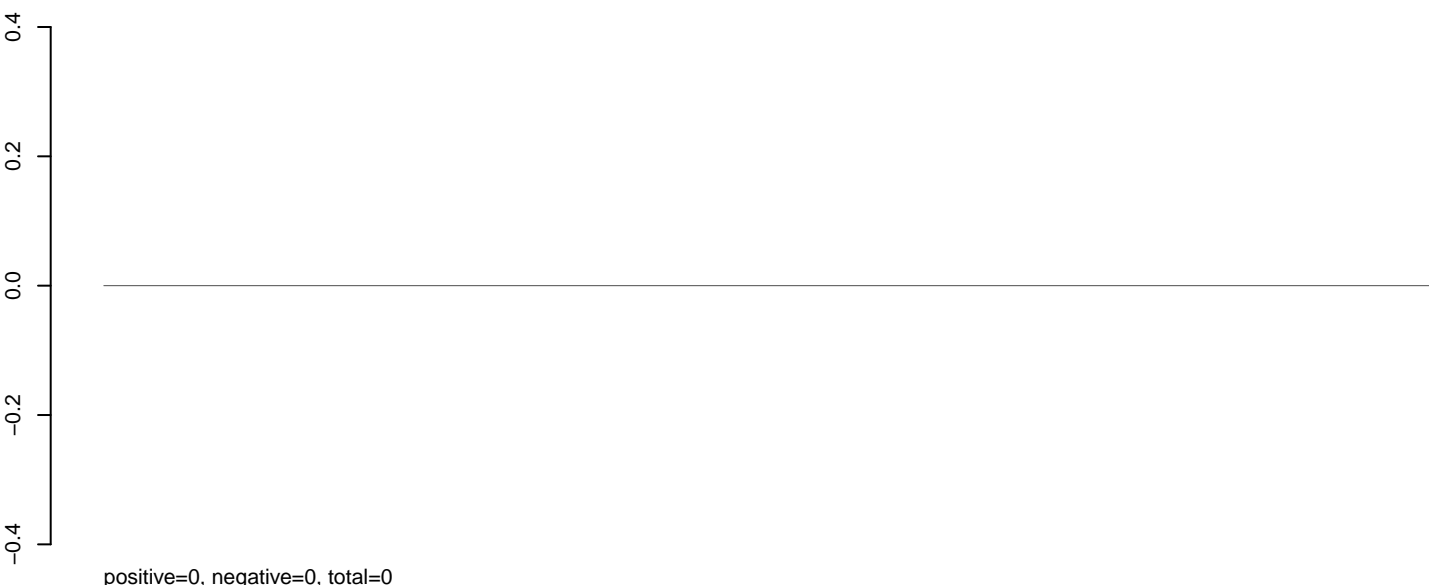
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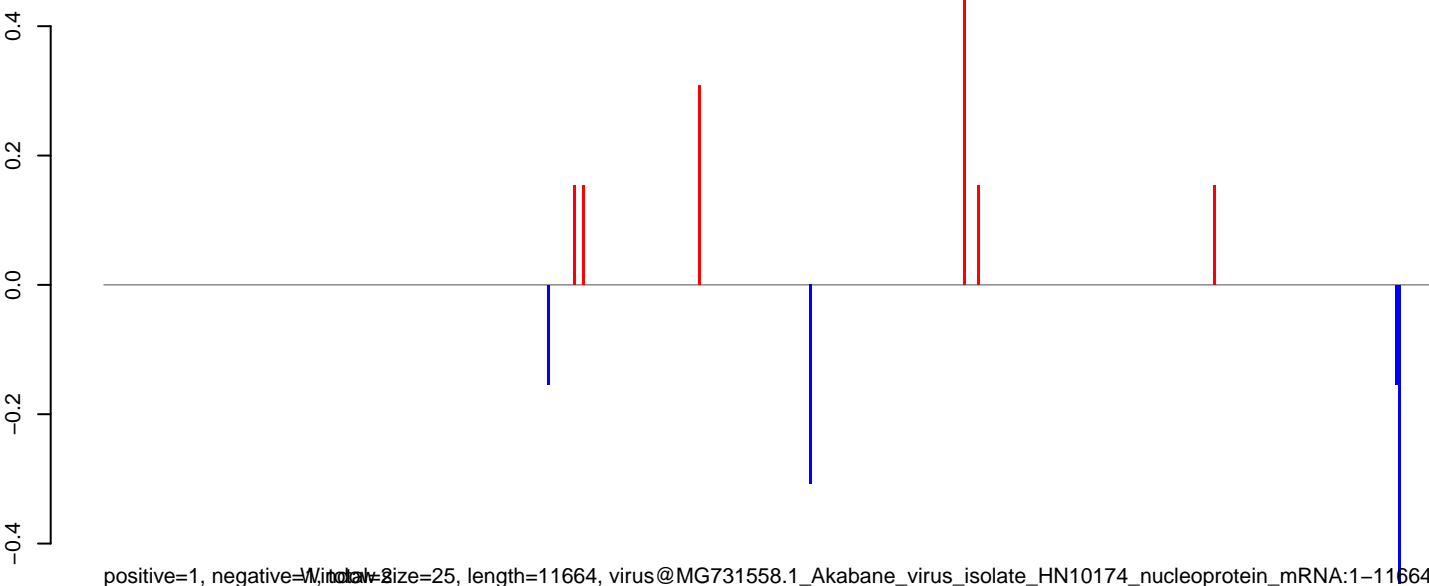
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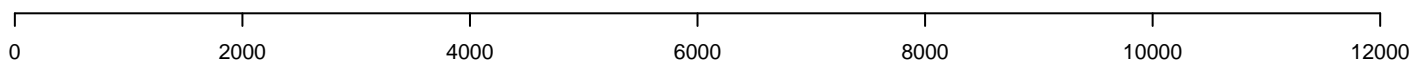
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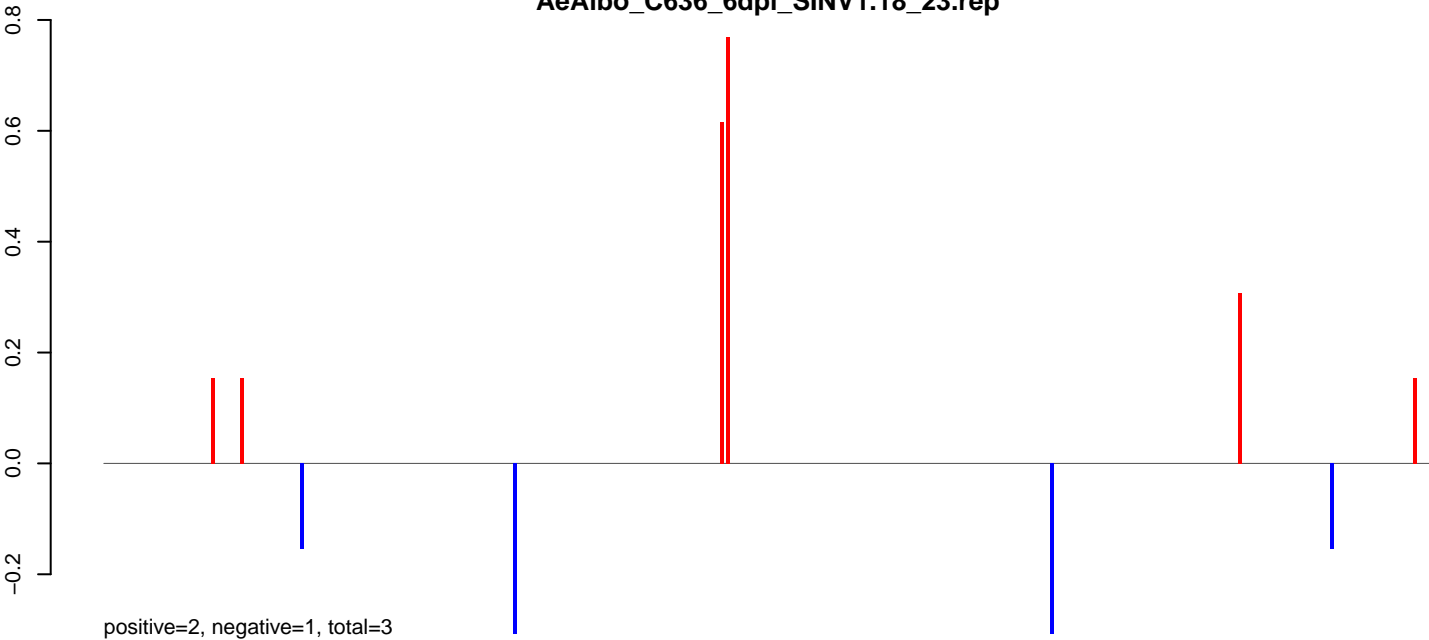
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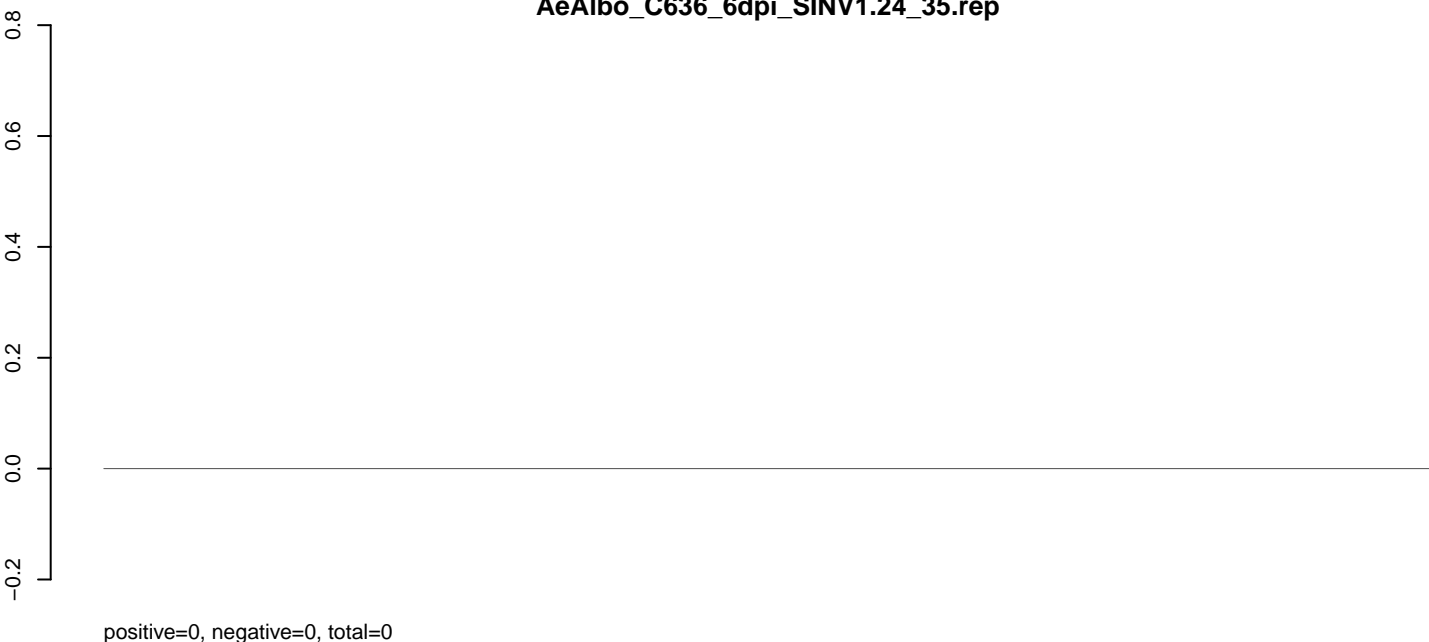
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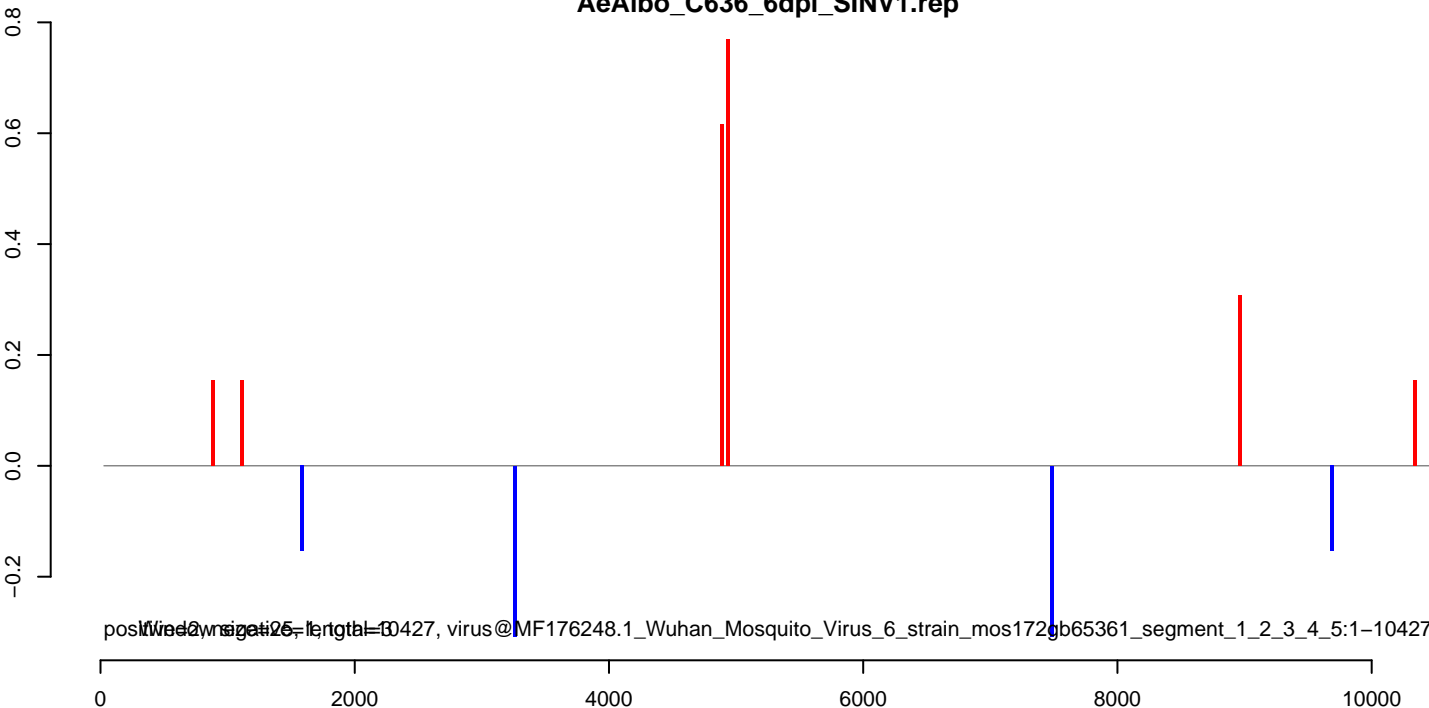
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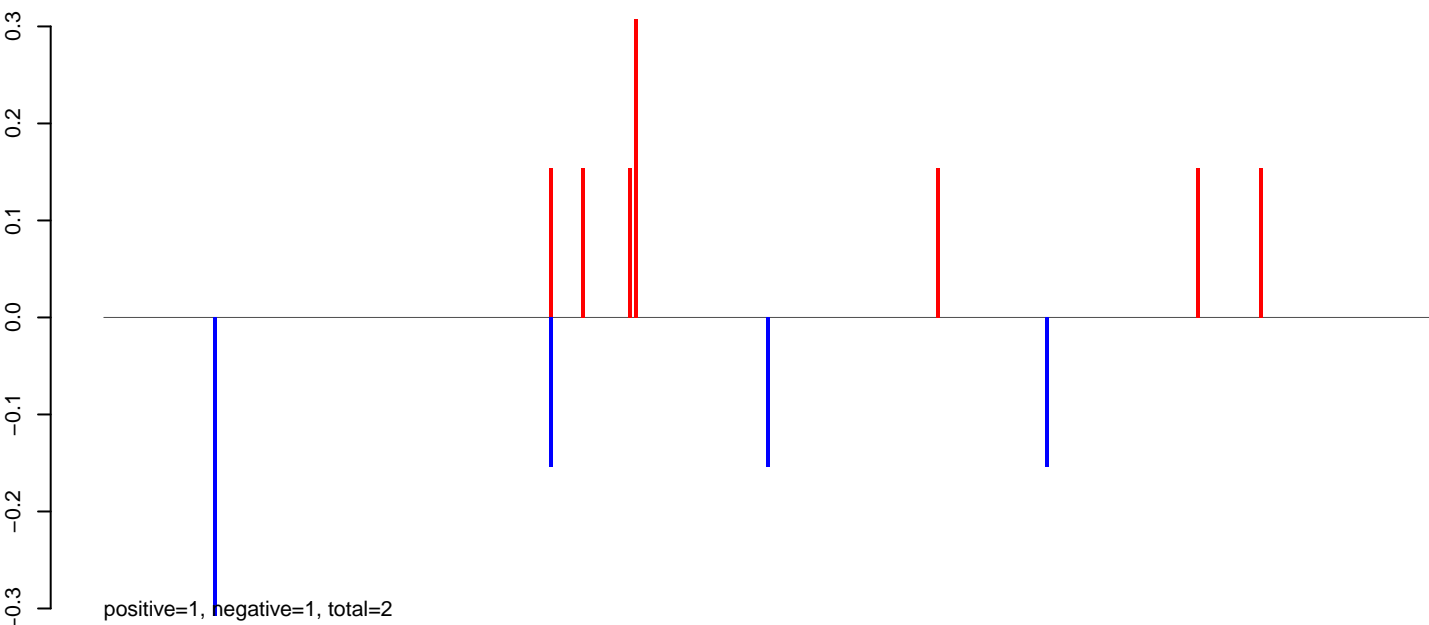
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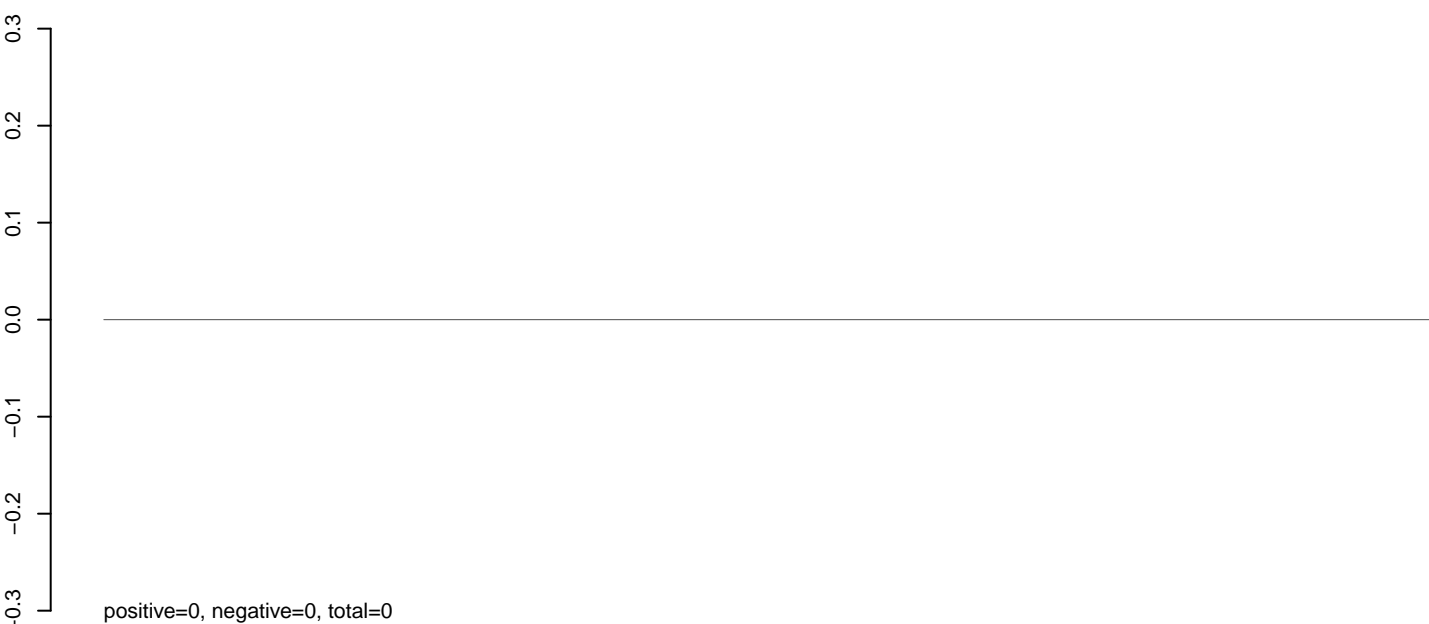
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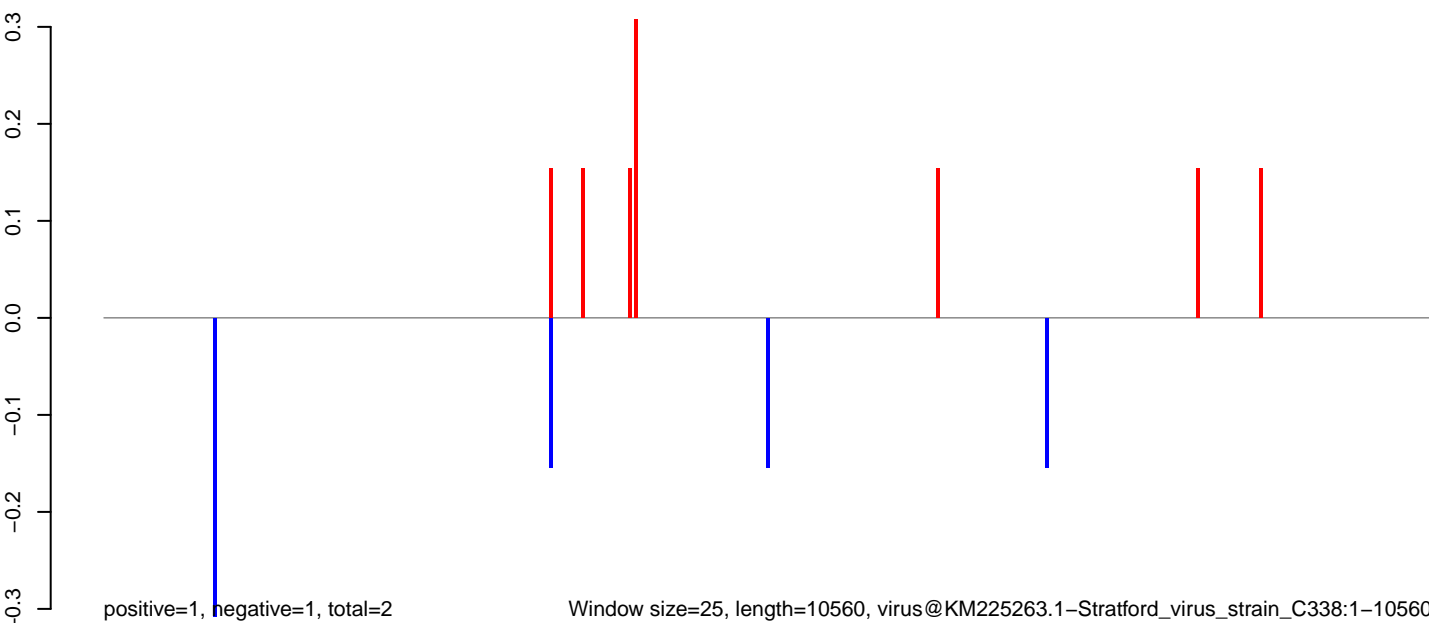
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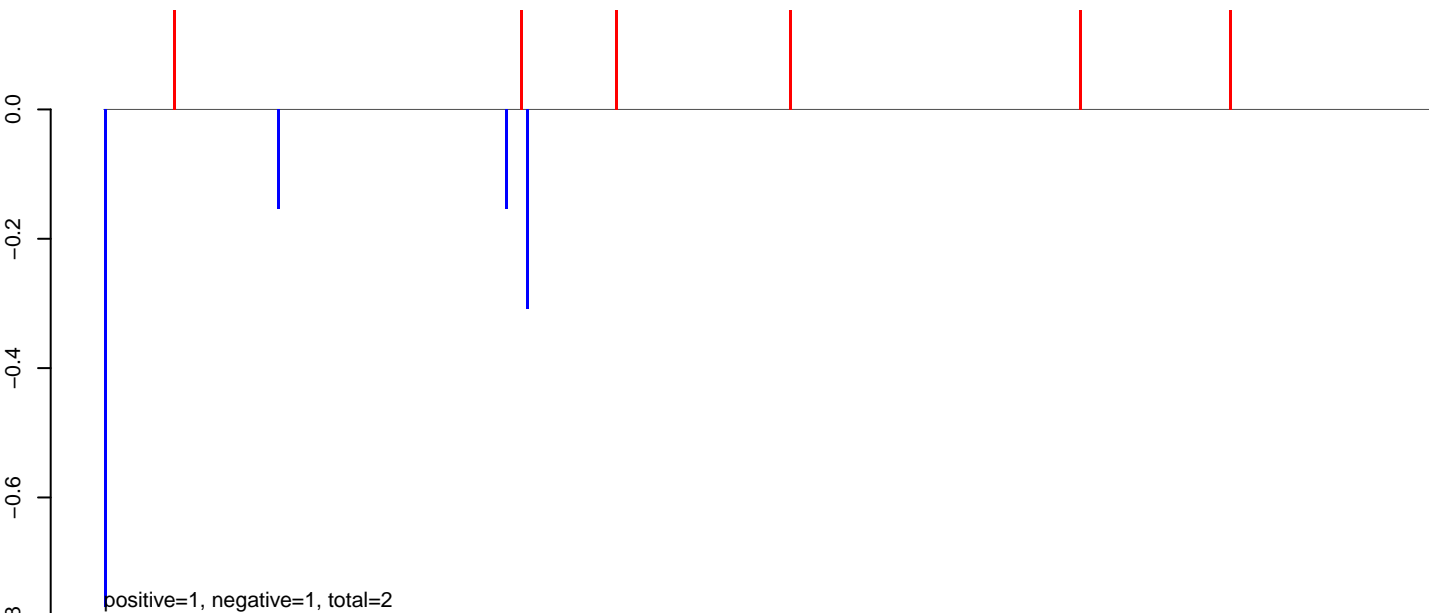
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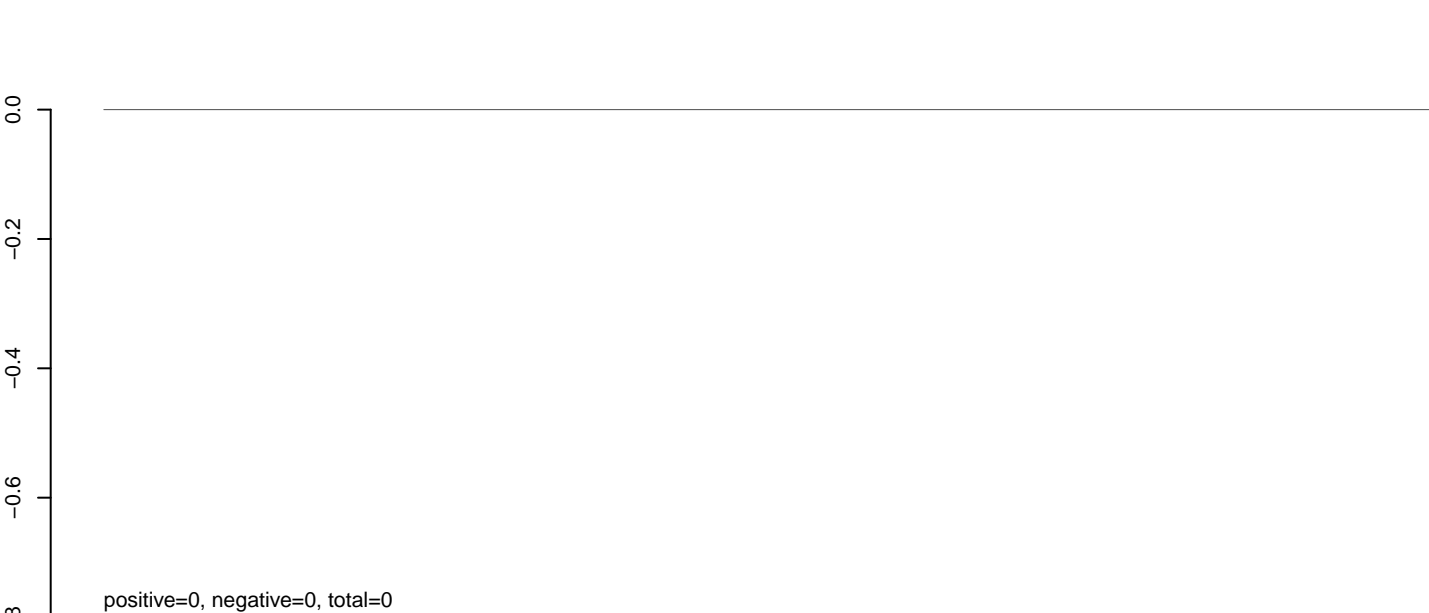
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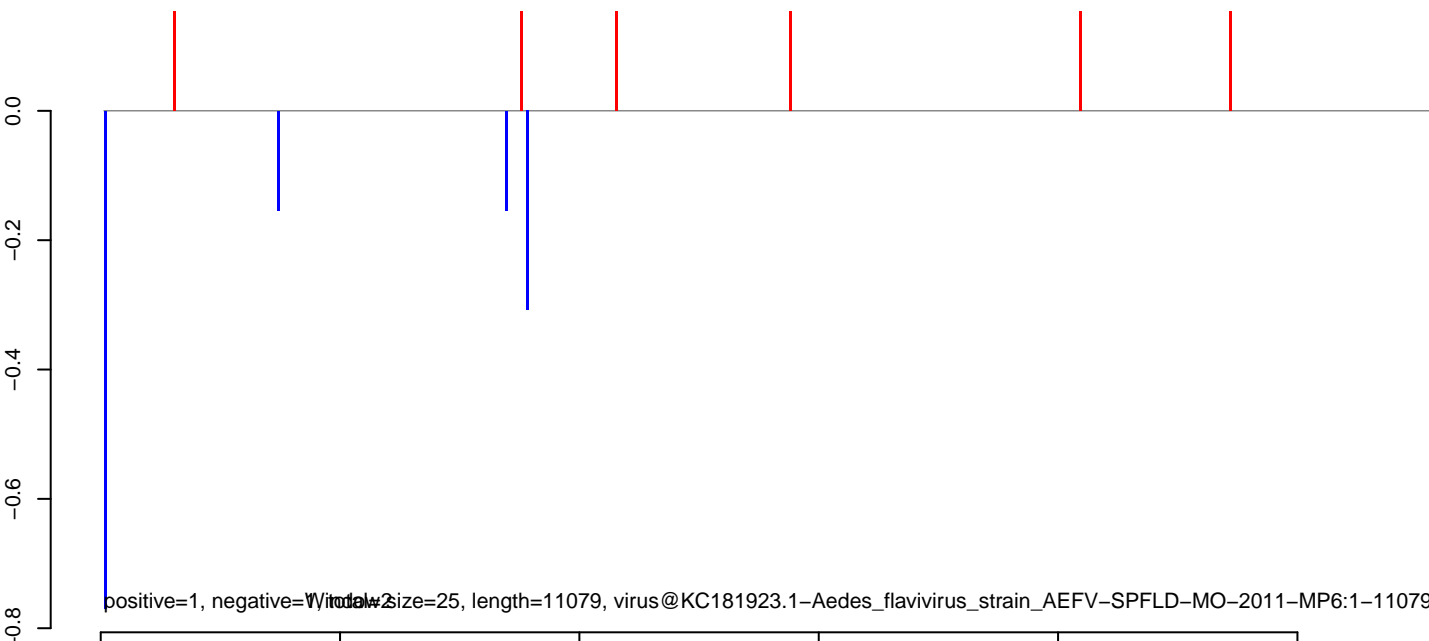
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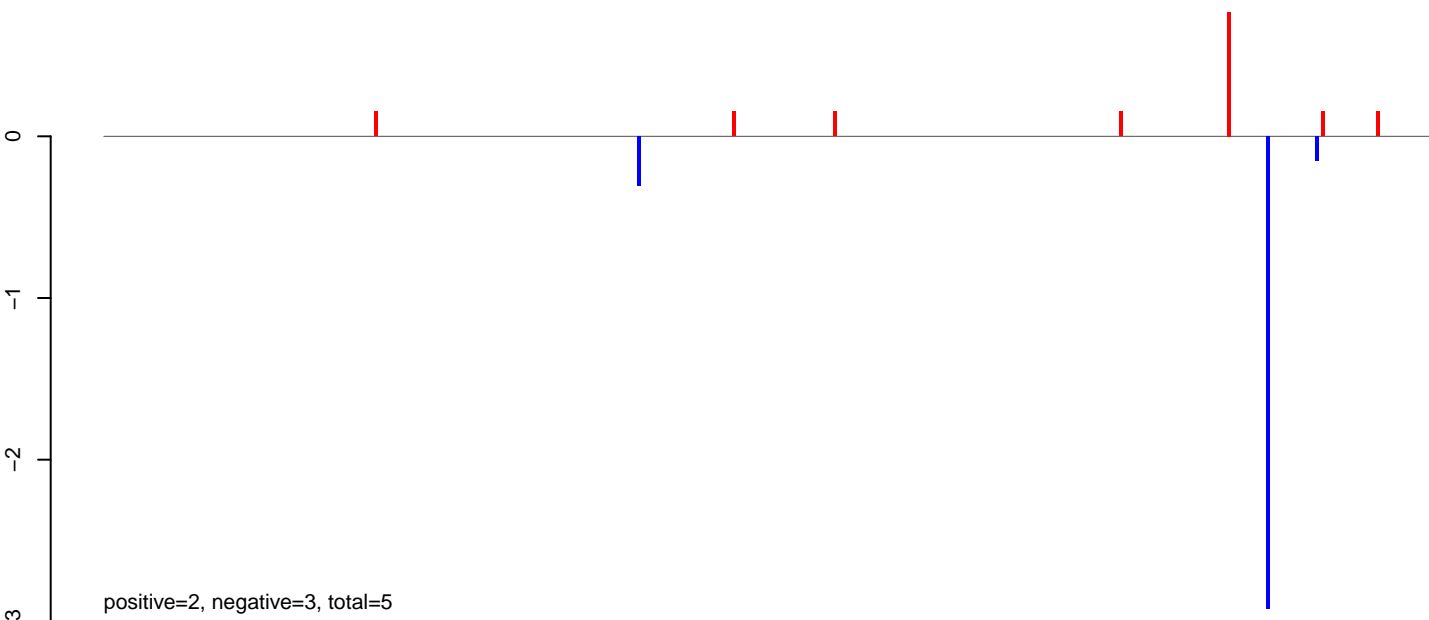
AeAlbo_C636_6dpi_SINV1.24_35.rep



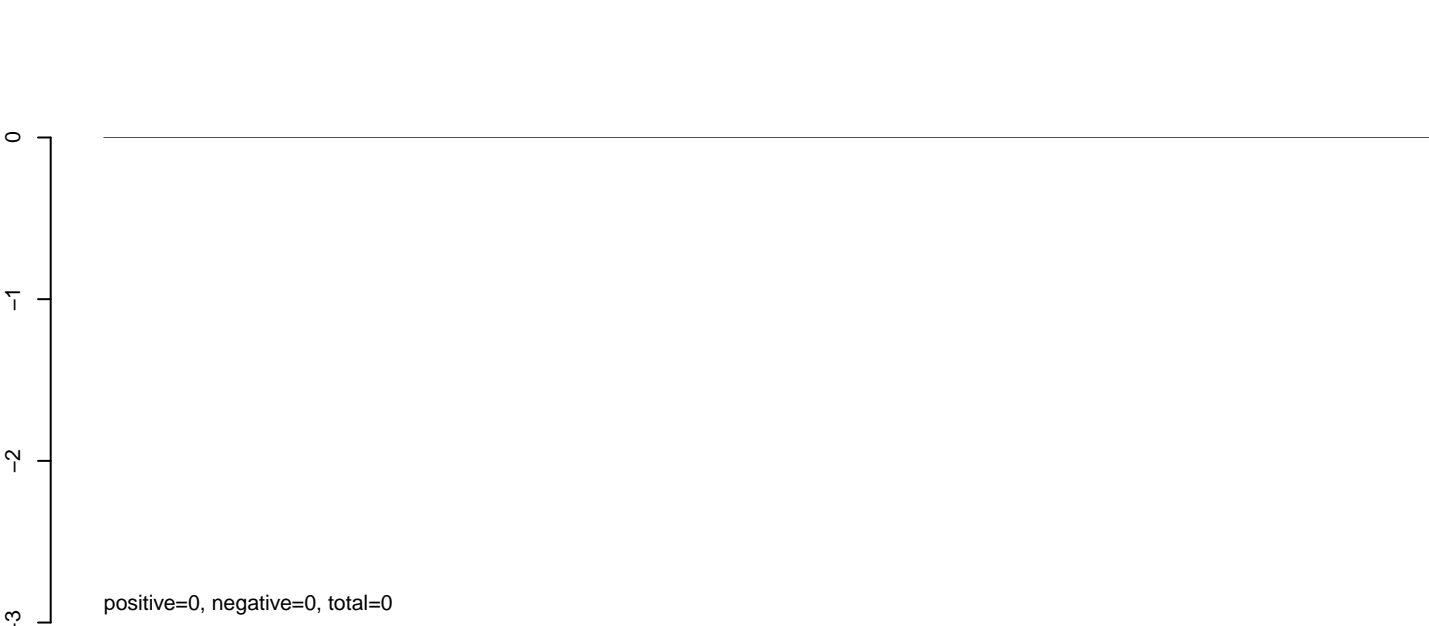
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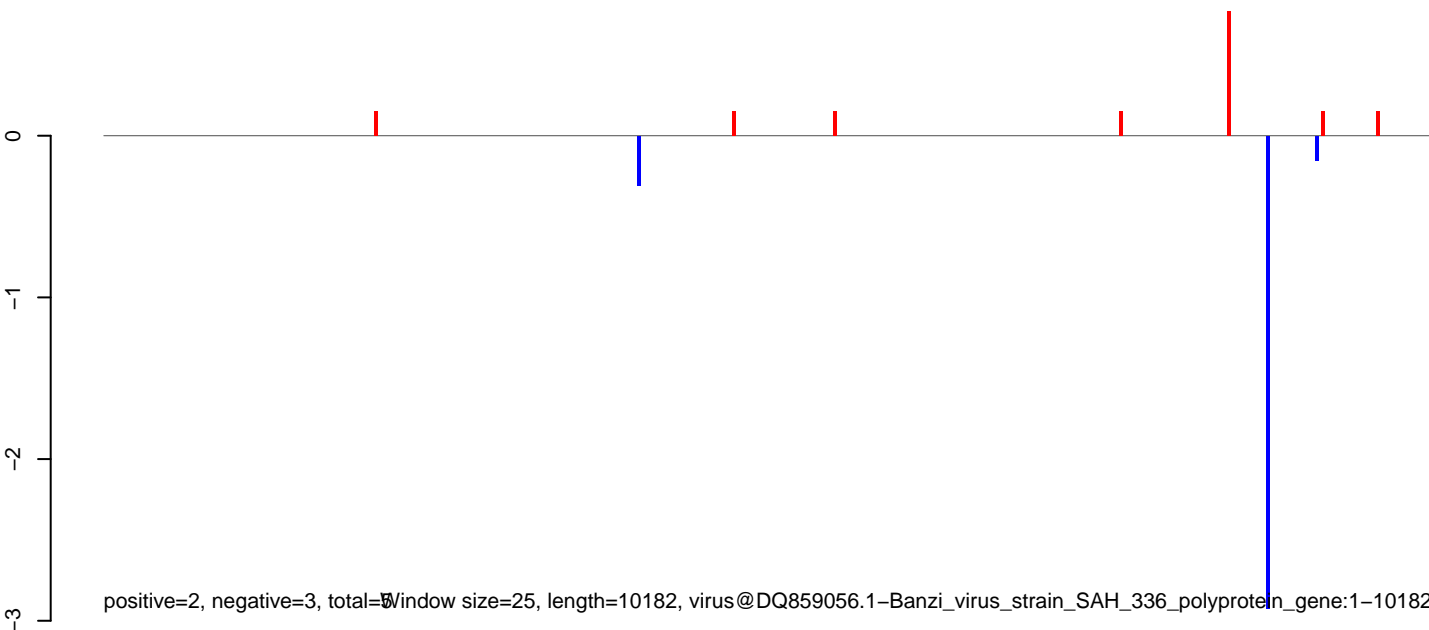
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AeAlbo_C636_6dpi_SINV1.24_35.rep



AeAlbo_C636_6dpi_SINV1.rep



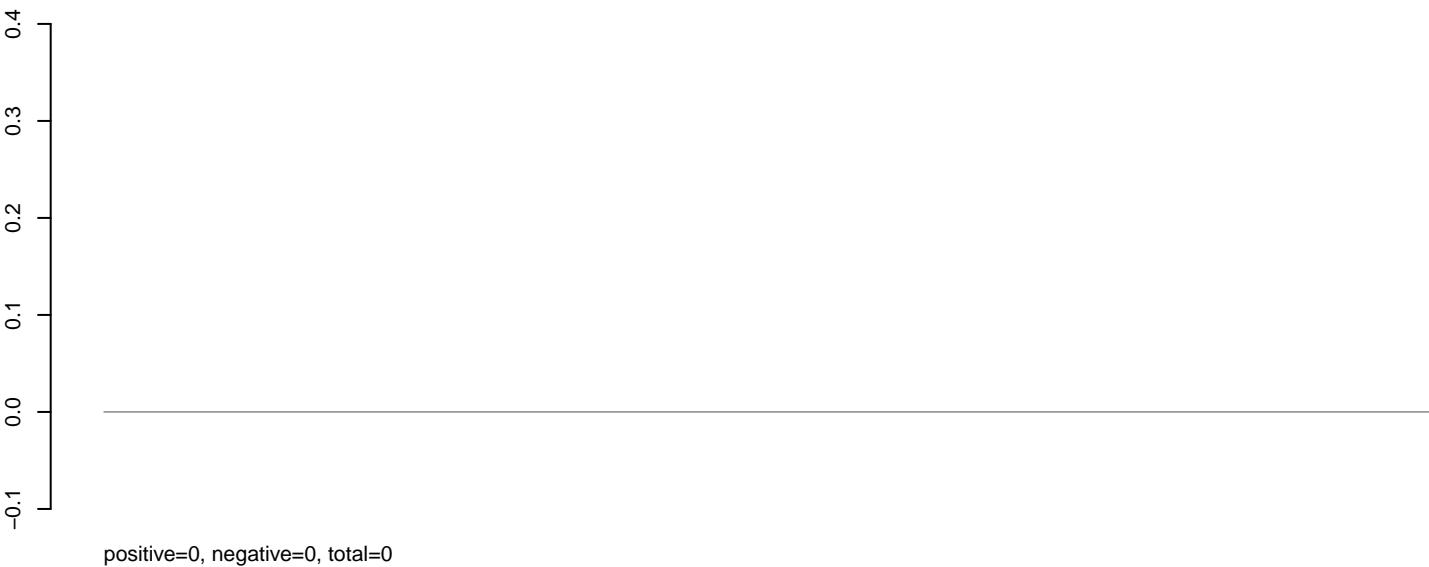
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0 2000 4000 6000 8000 10000

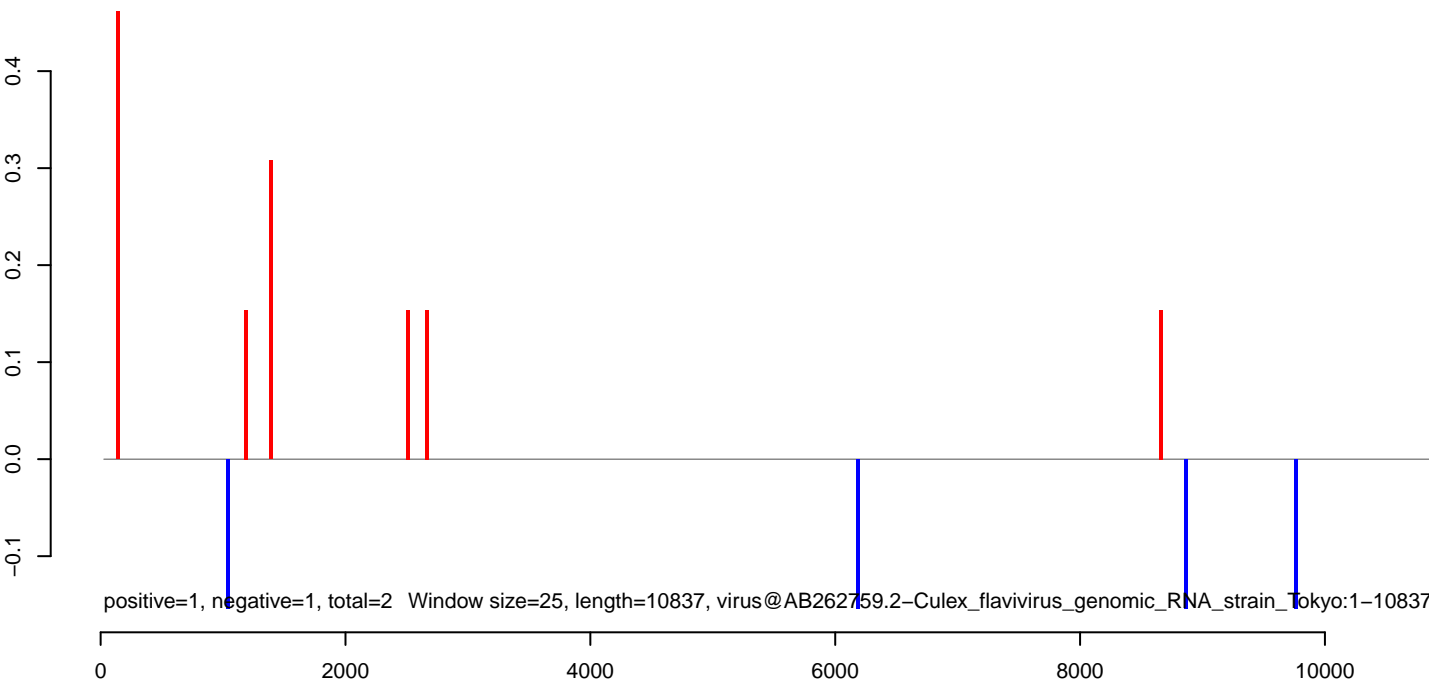
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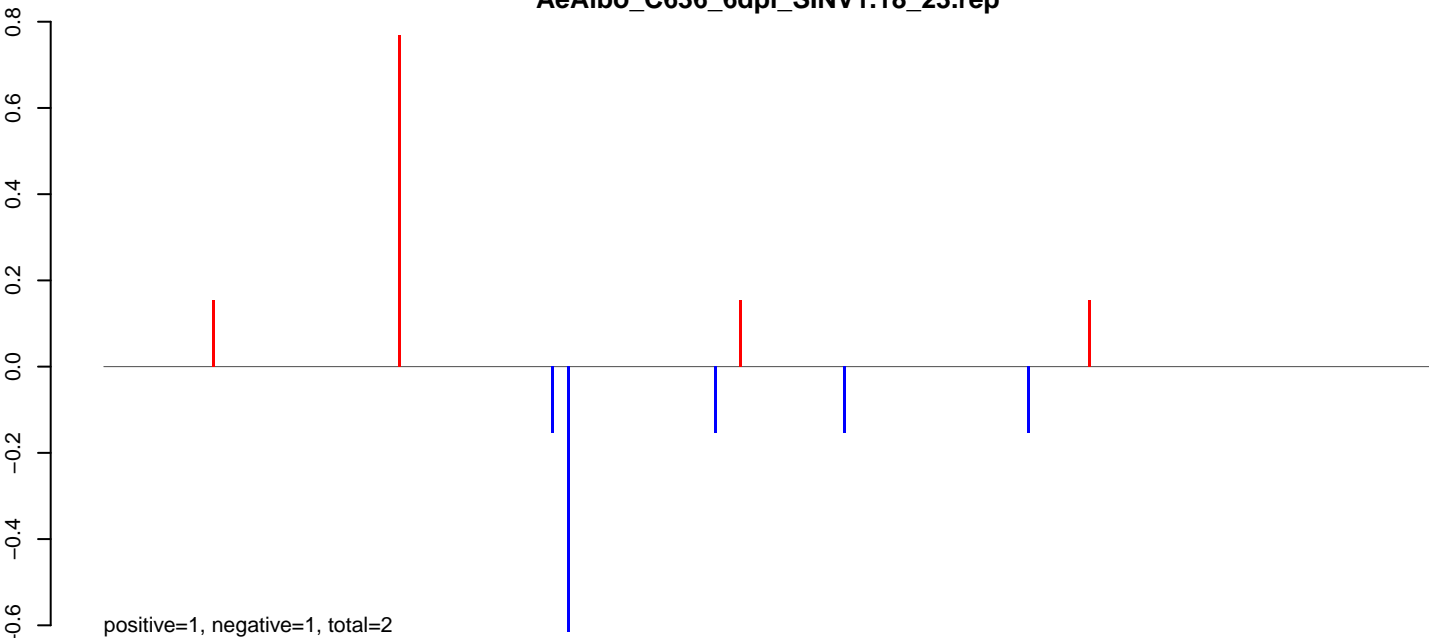
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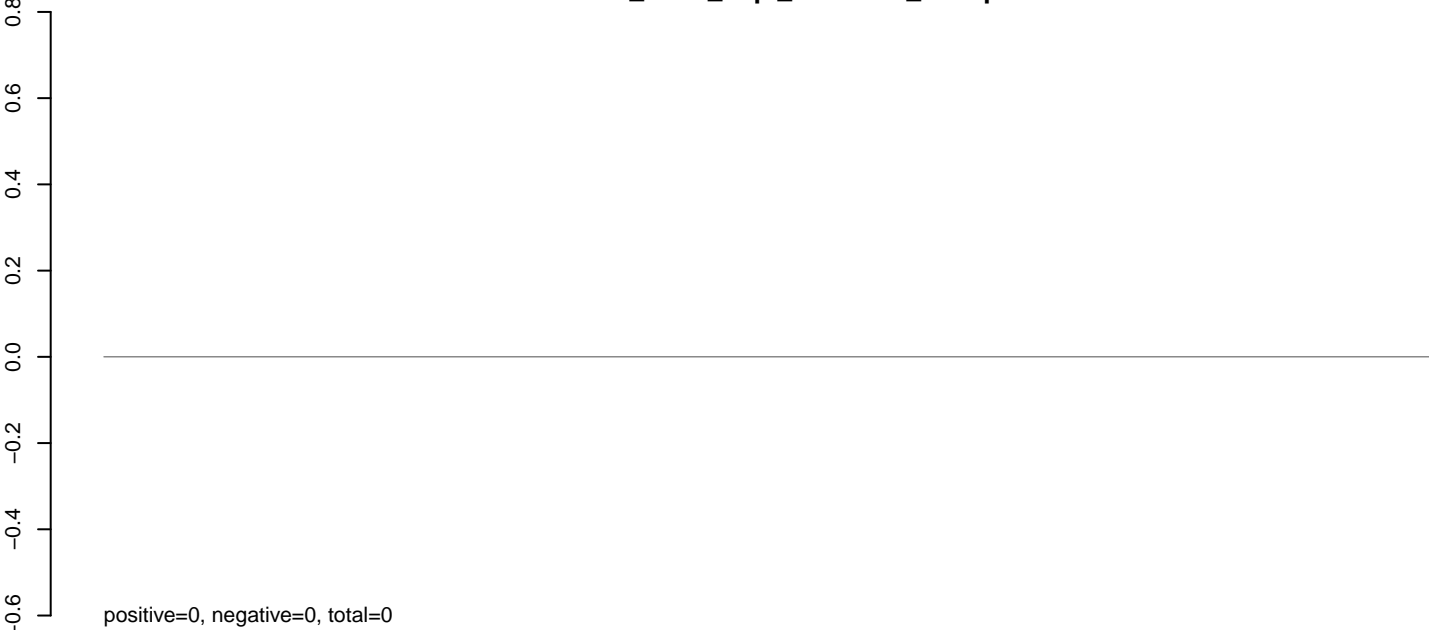
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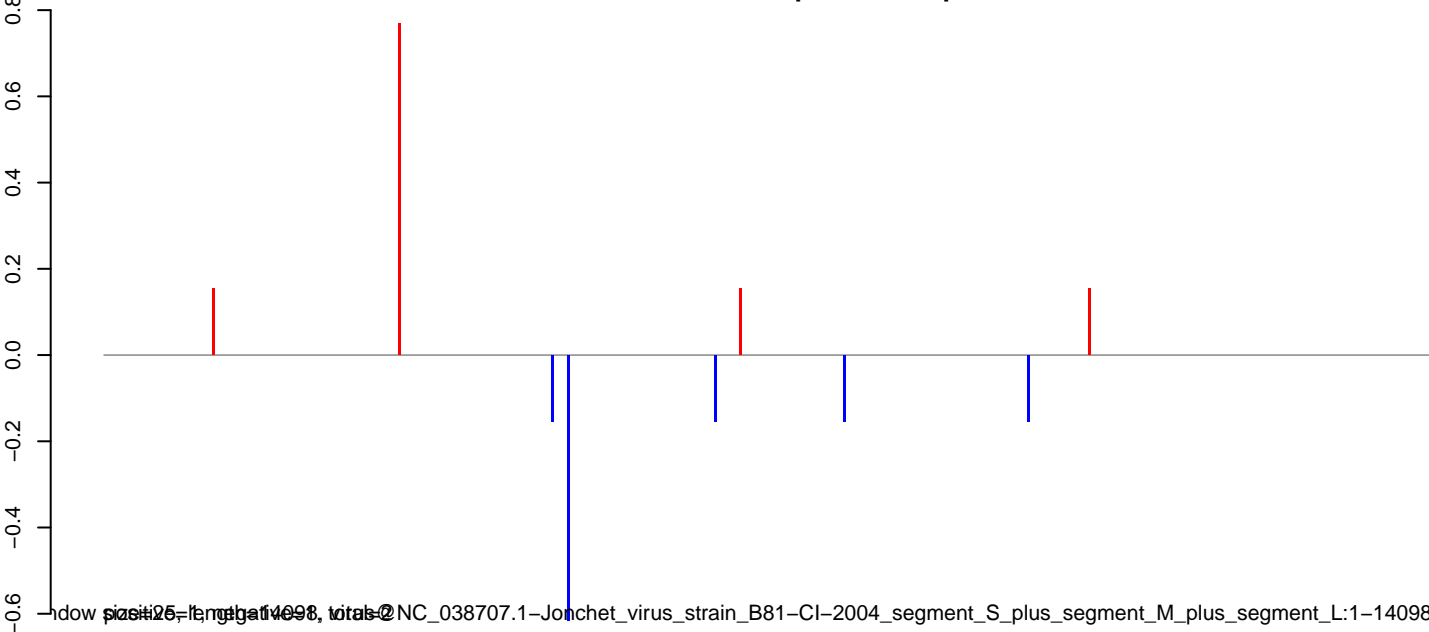
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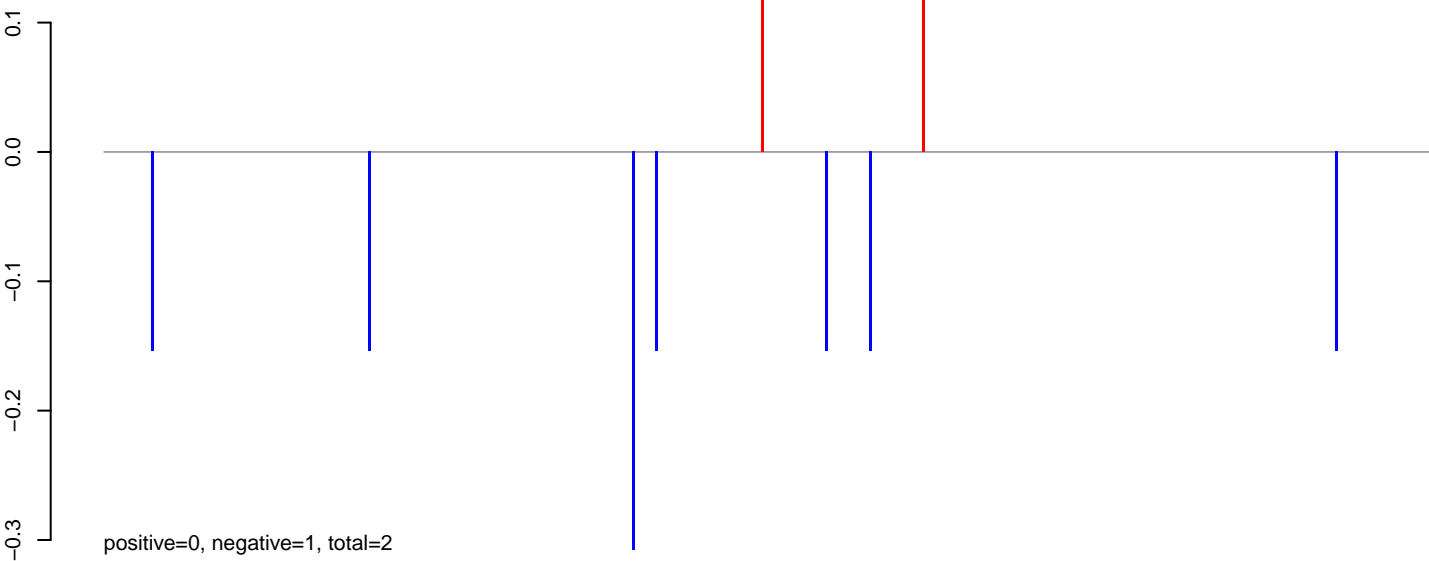
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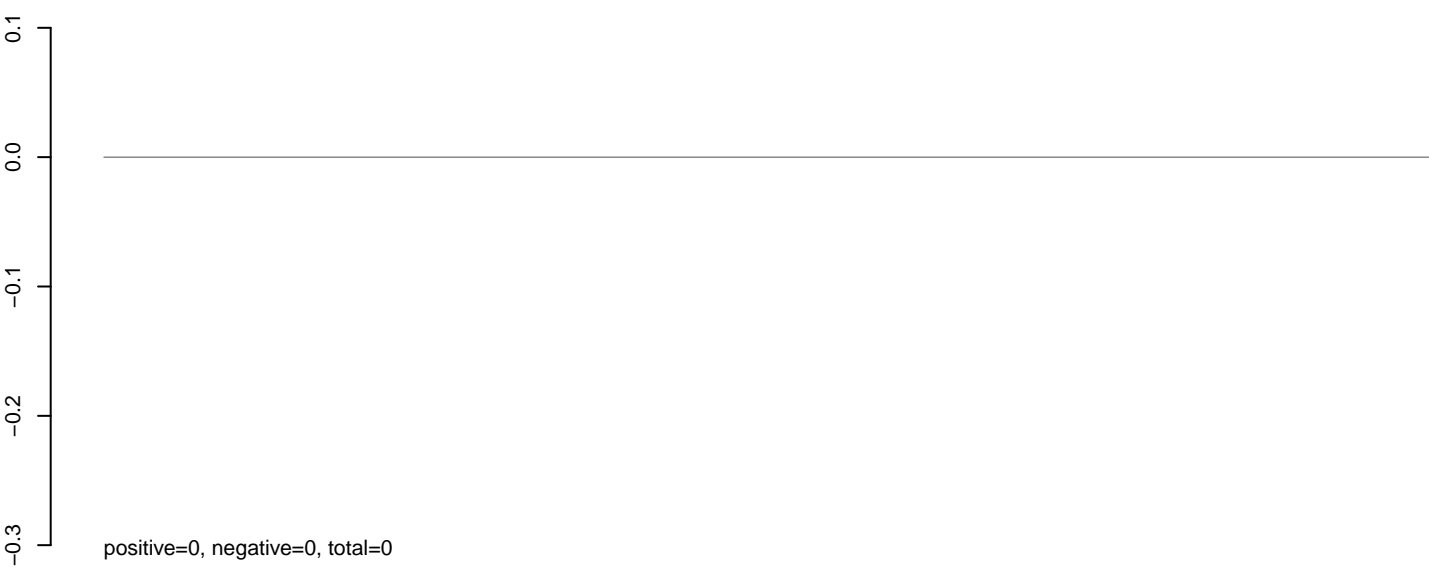
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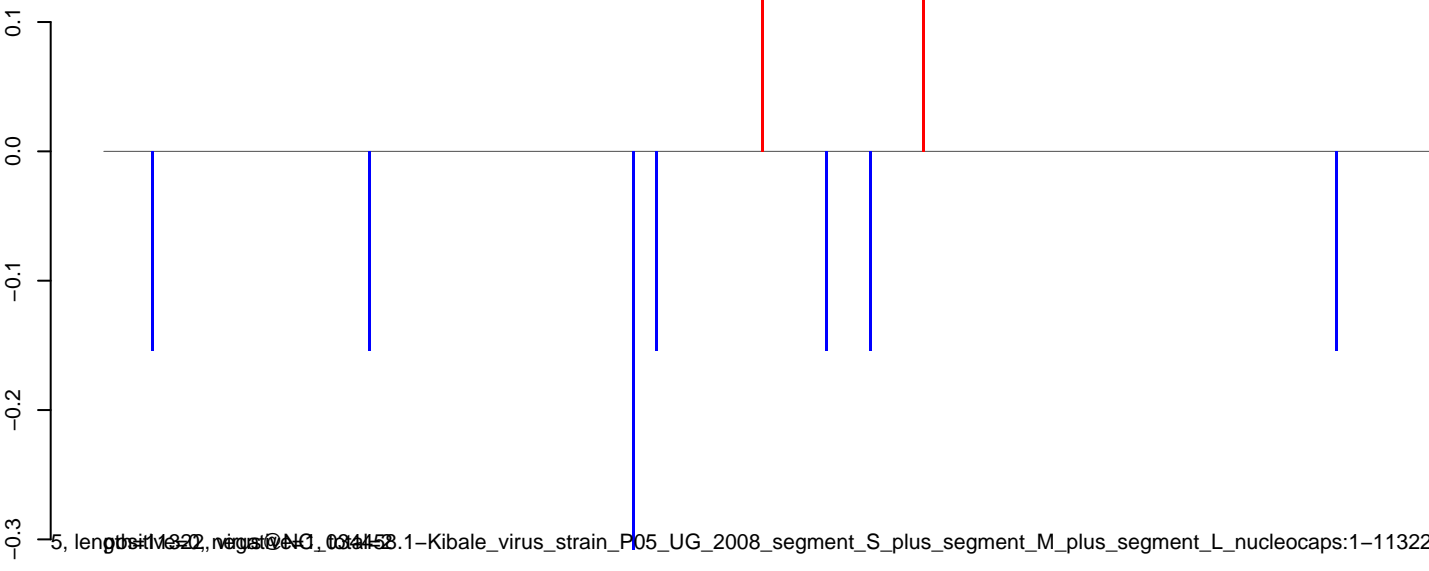
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AeAlbo_C636_6dpi_SINV1.24_35.rep

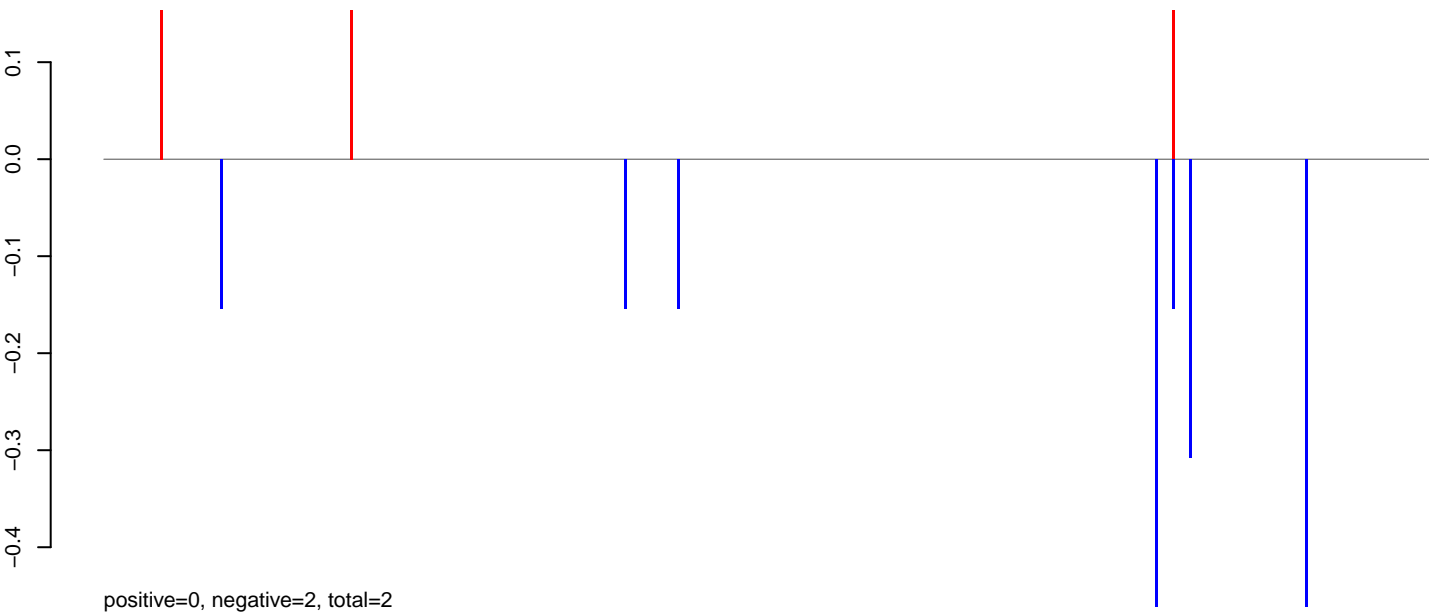


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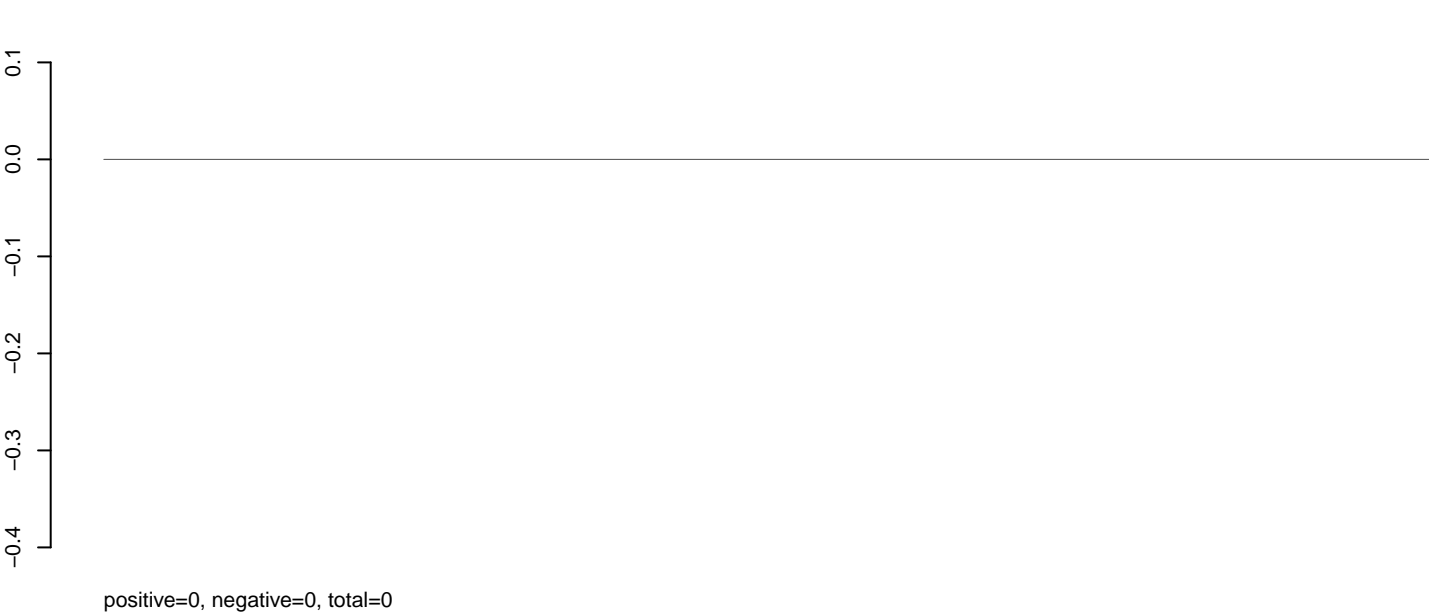


5, length=11322, negative=0, total=0, 1-Kibale_virus_strain_P05_UG_2008_segment_S_plus_segment_M_plus_segment_L_nucleocaps:1-11322

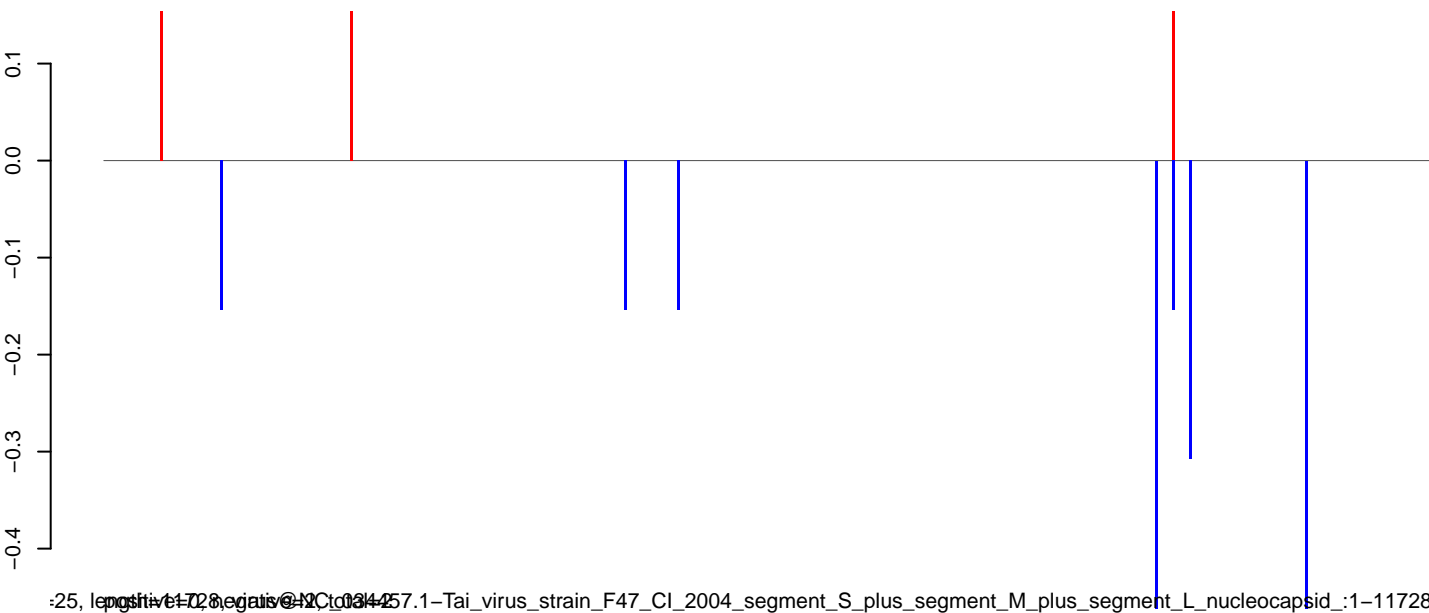
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AeAlbo_C636_6dpi_SINV1.24_35.rep



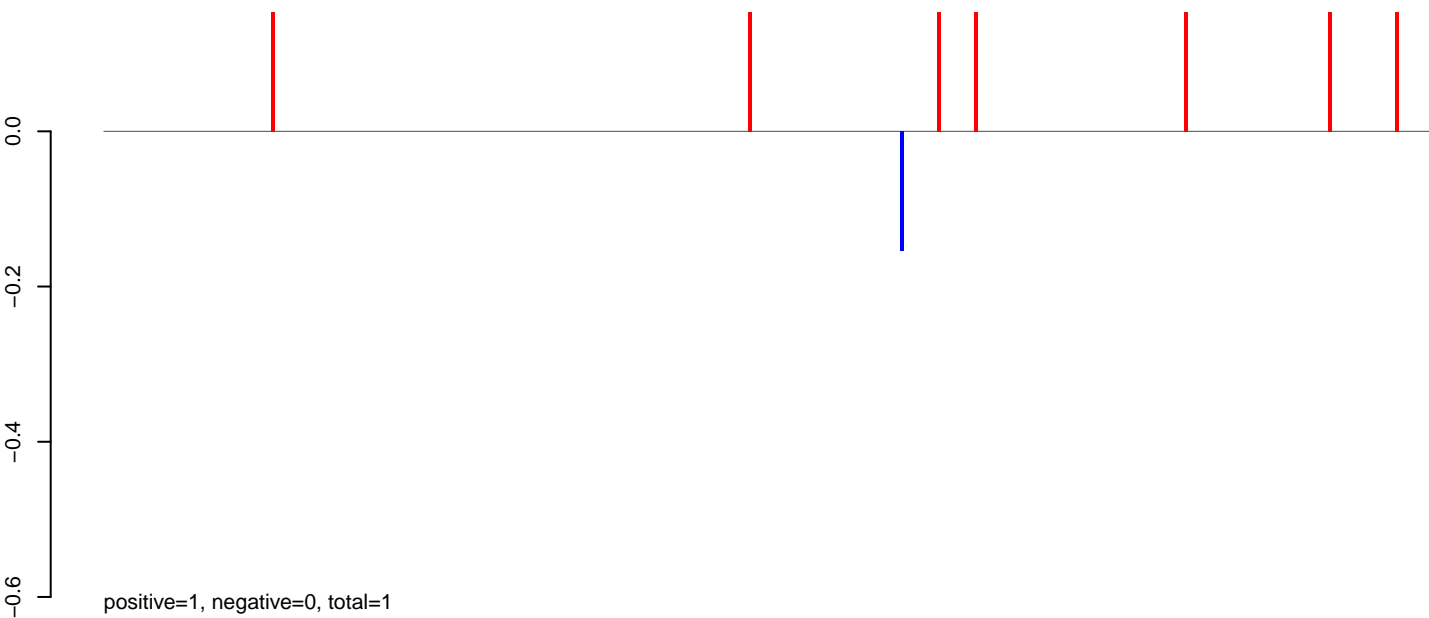
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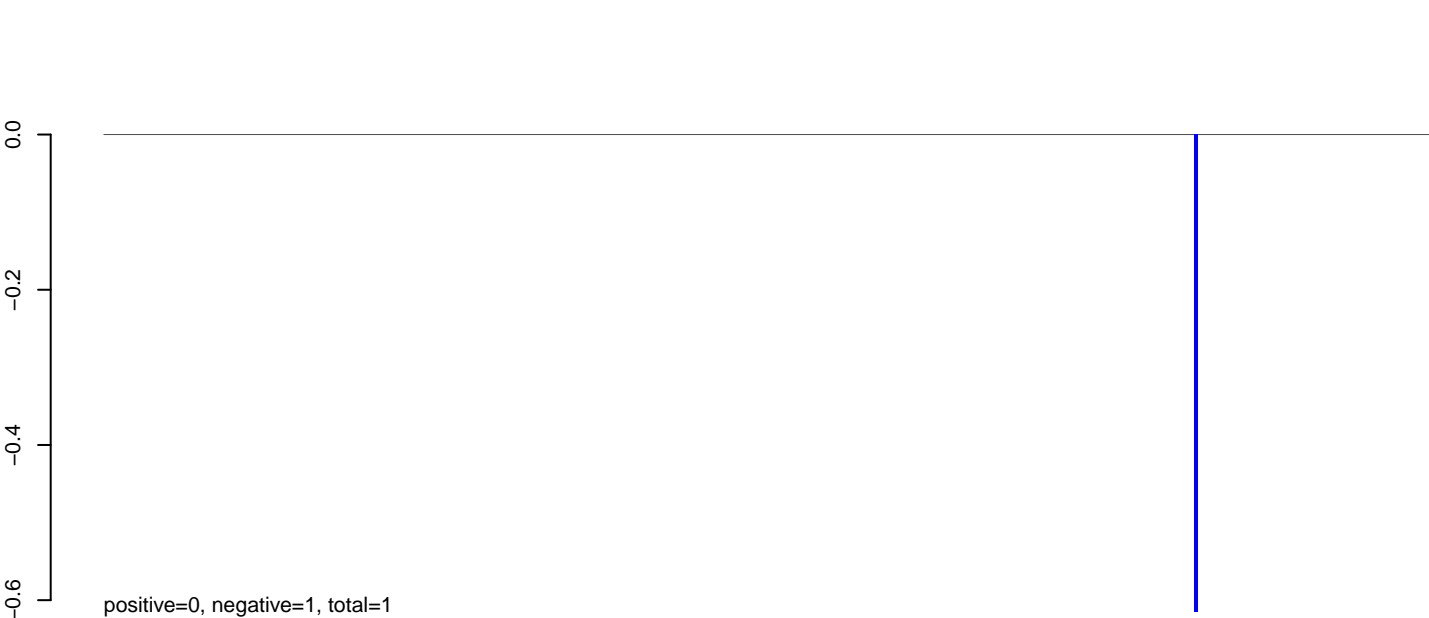
25, length=11728, negative=5, total=8
103445-1-Tai_virus_strain_F47_CI_2004_segment_S_plus_segment_M_plus_segment_L_nucleocapsid_:1-11728

0 2000 4000 6000 8000 10000 12000

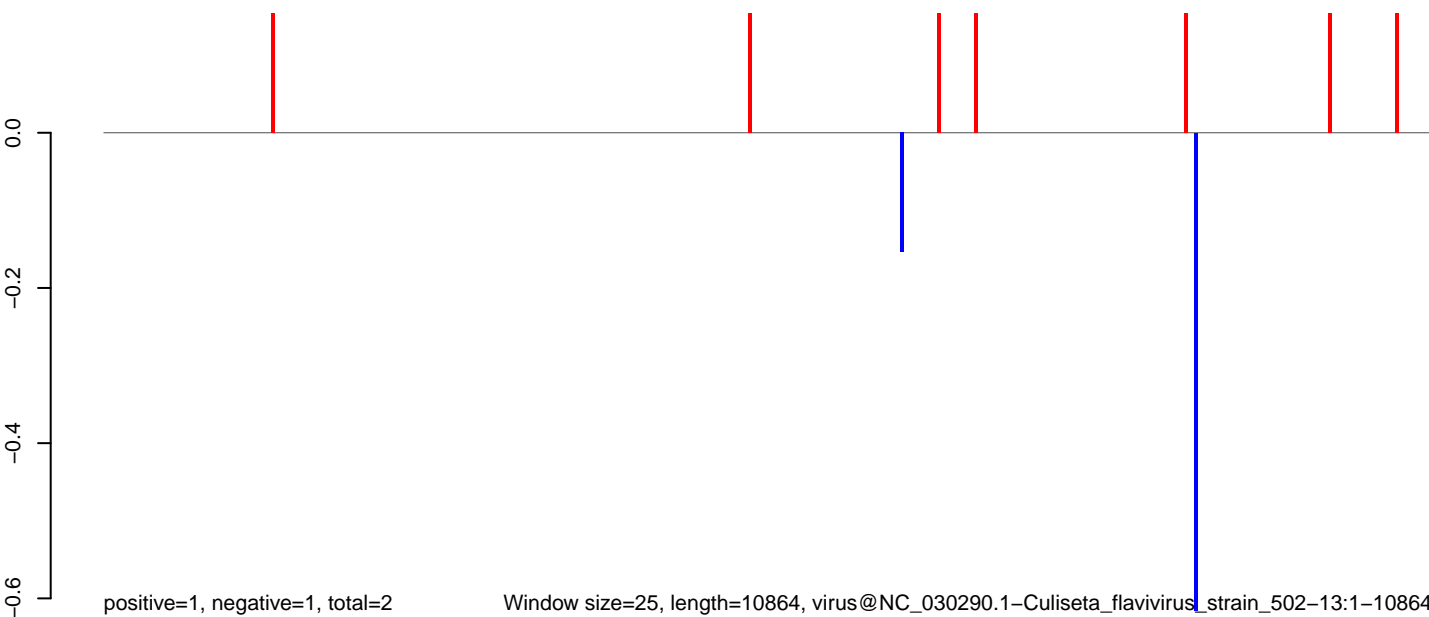
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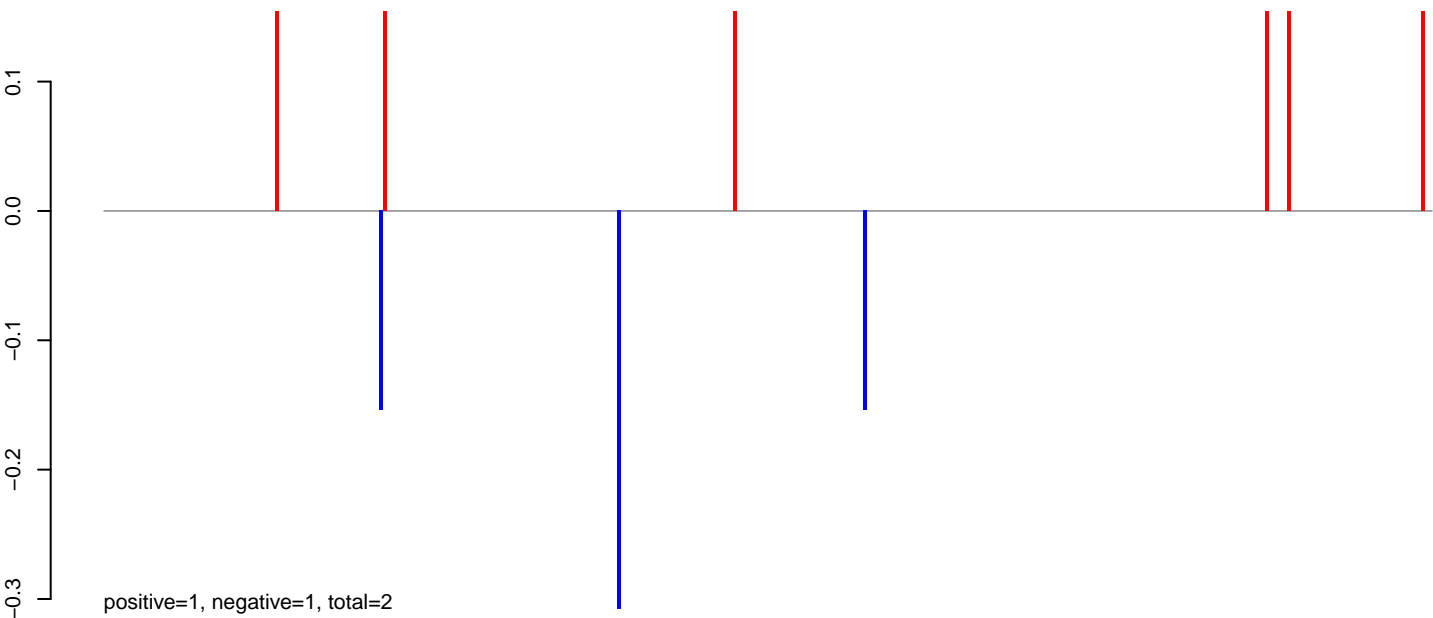
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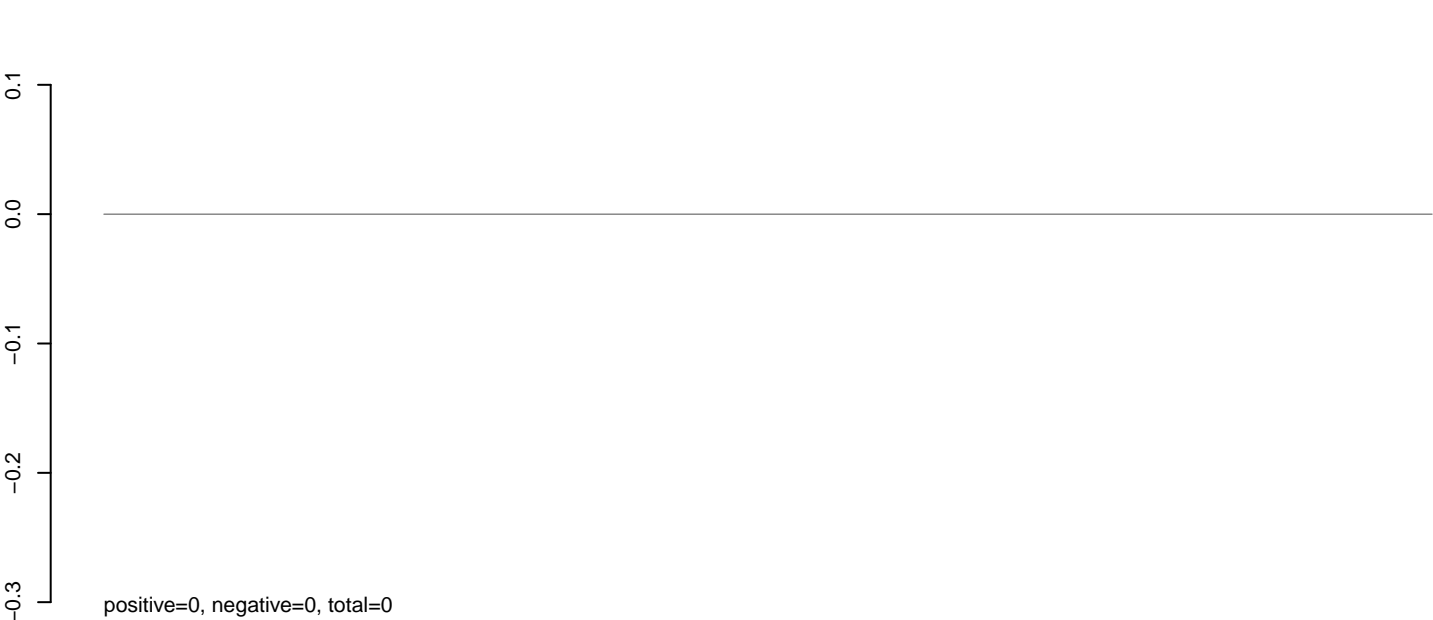
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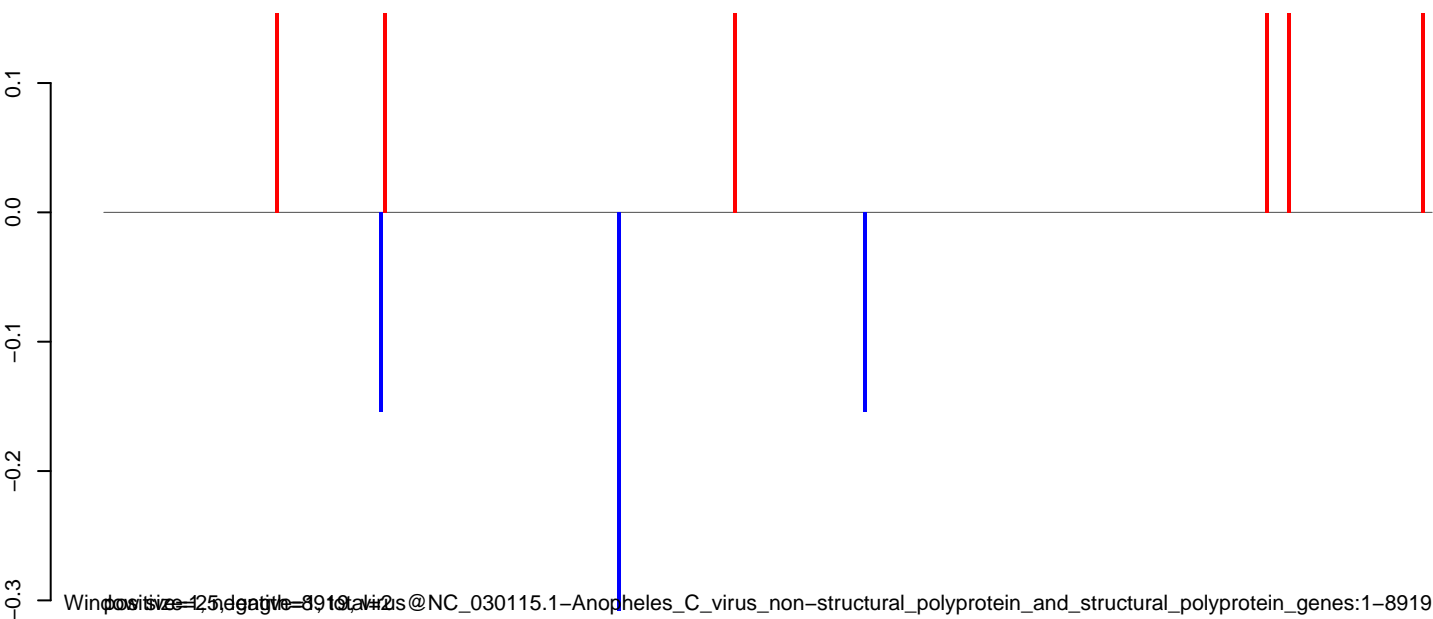
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AeAlbo_C636_6dpi_SINV1.24_35.rep



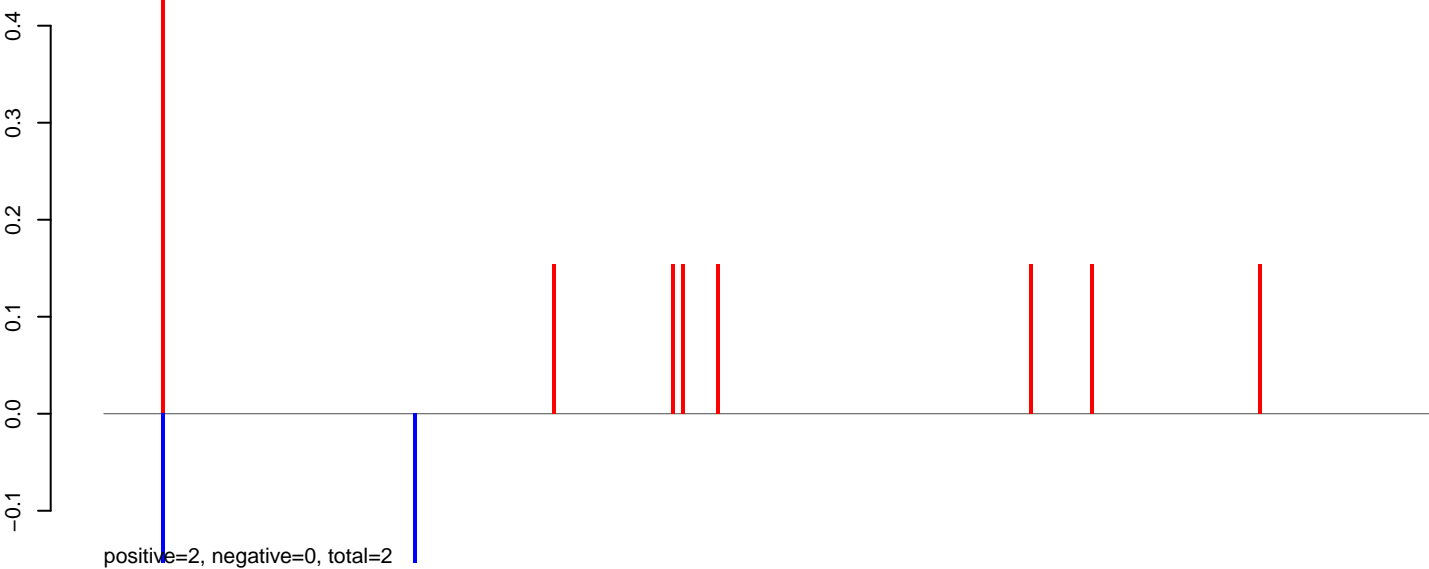
AeAlbo_C636_6dpi_SINV1.rep



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0 2000 4000 6000 8000

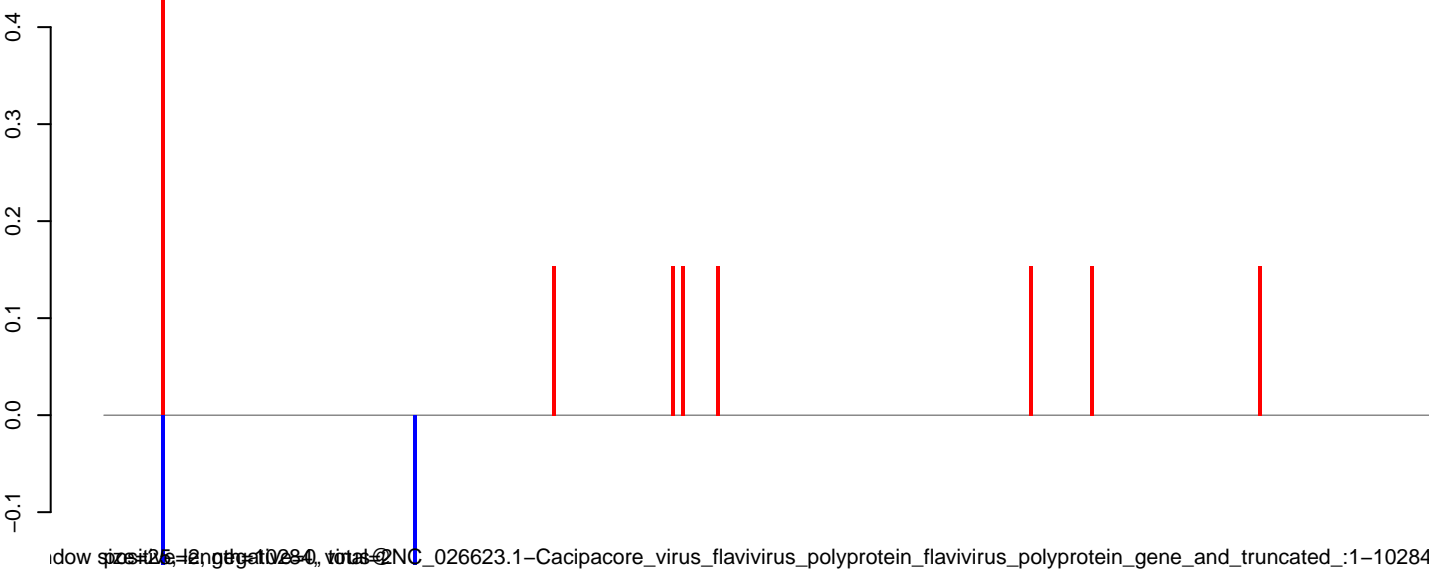
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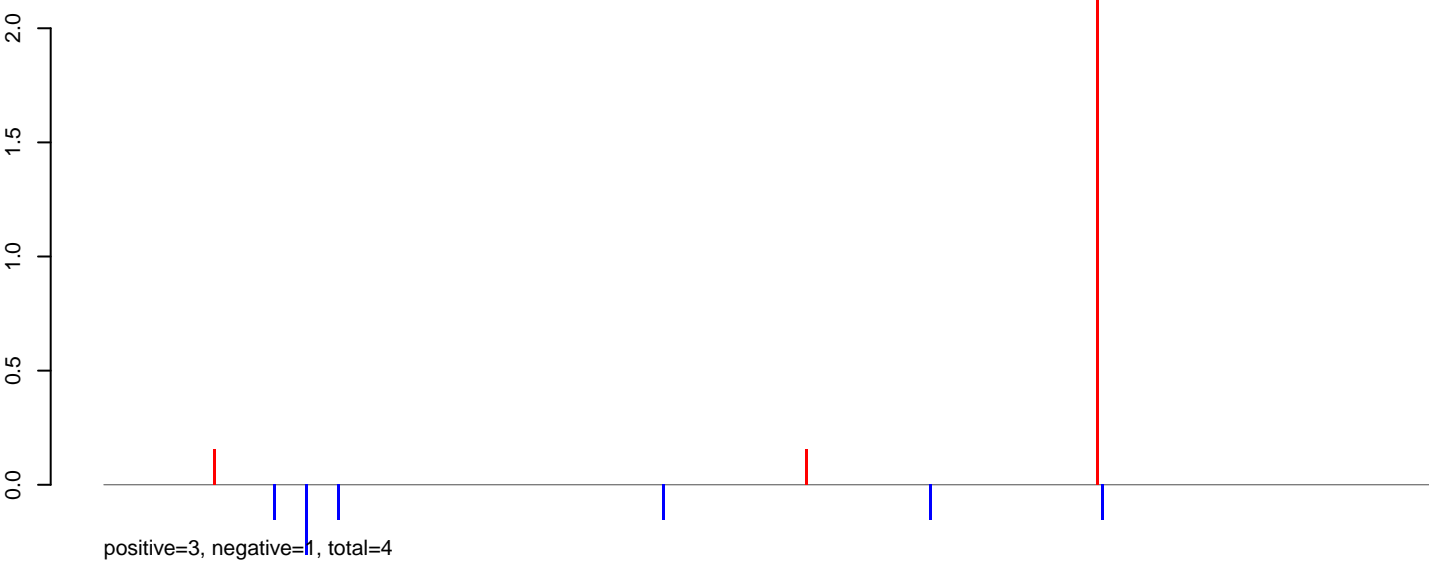
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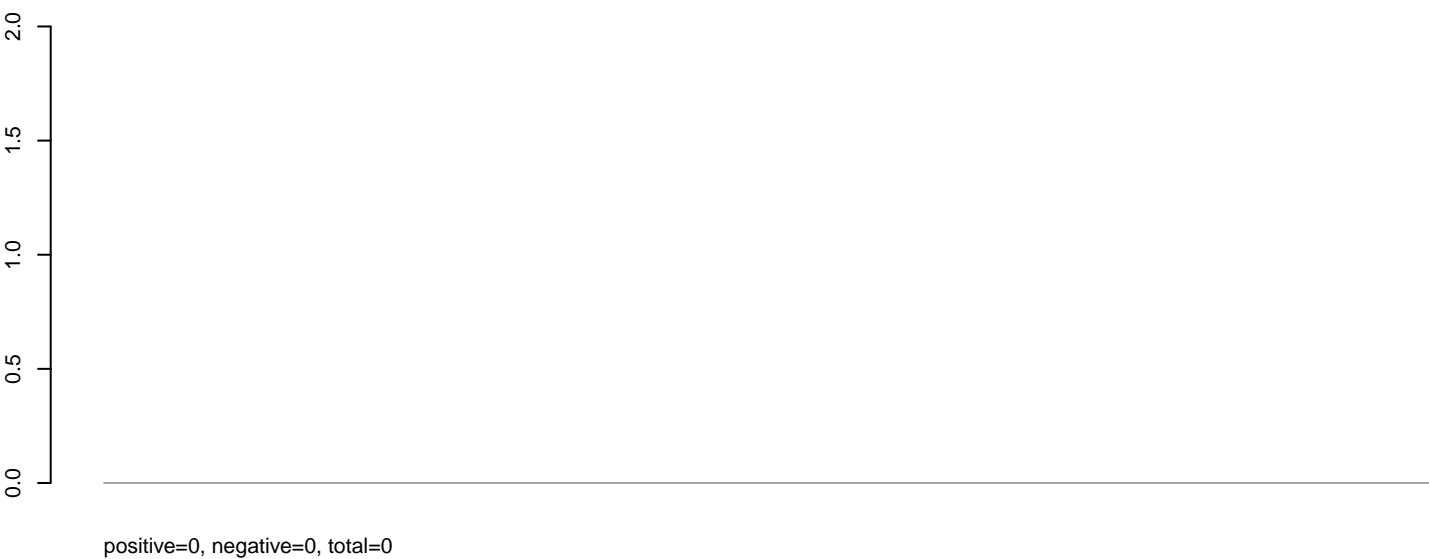
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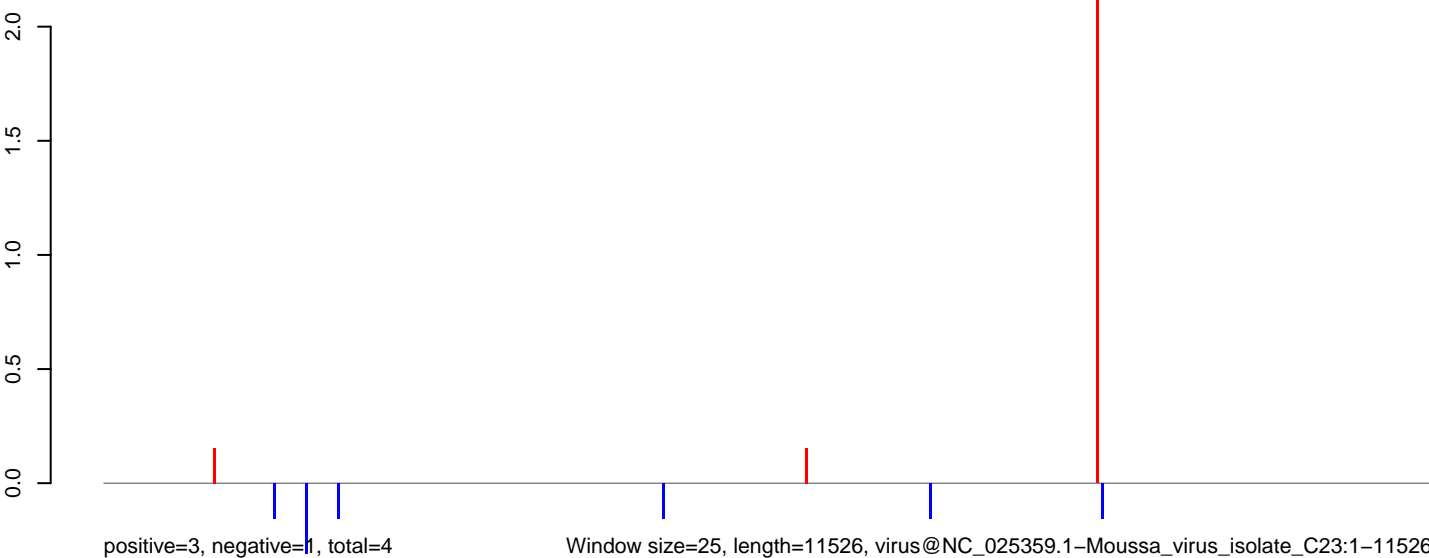
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AeAlbo_C636_6dpi_SINV1.24_35.rep

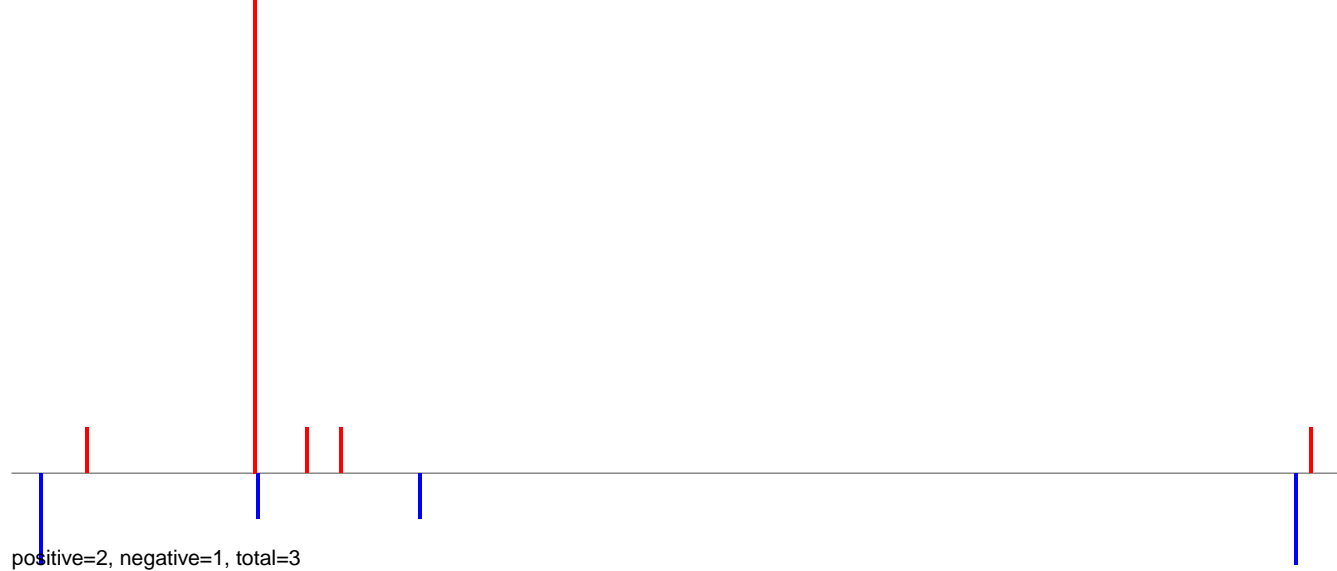


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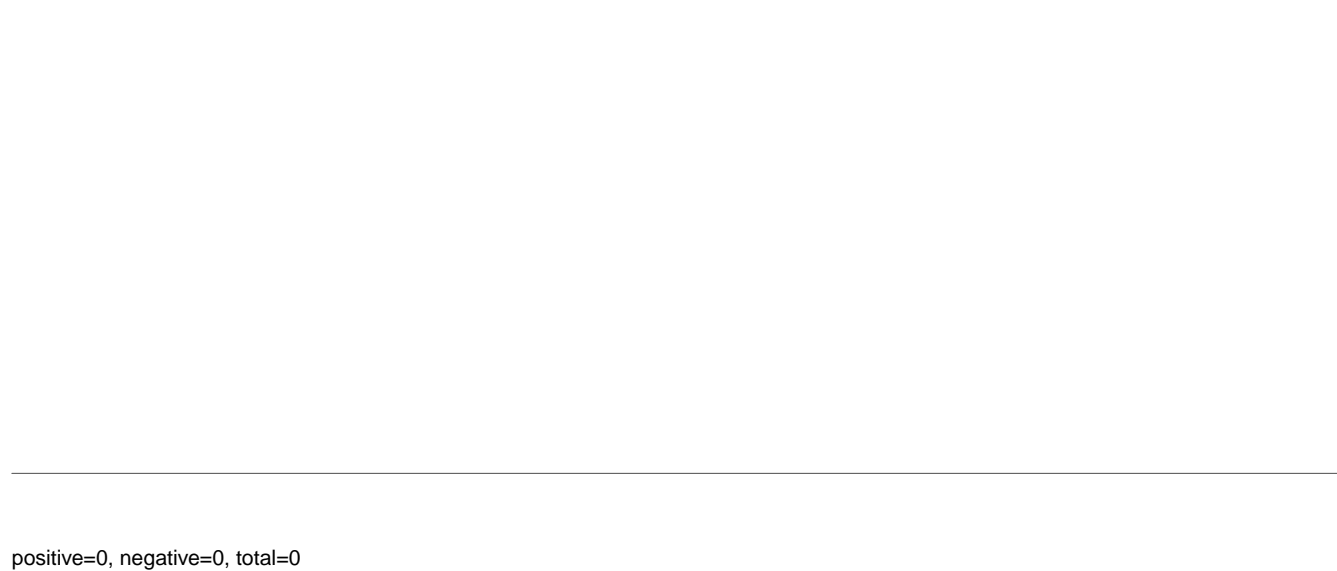


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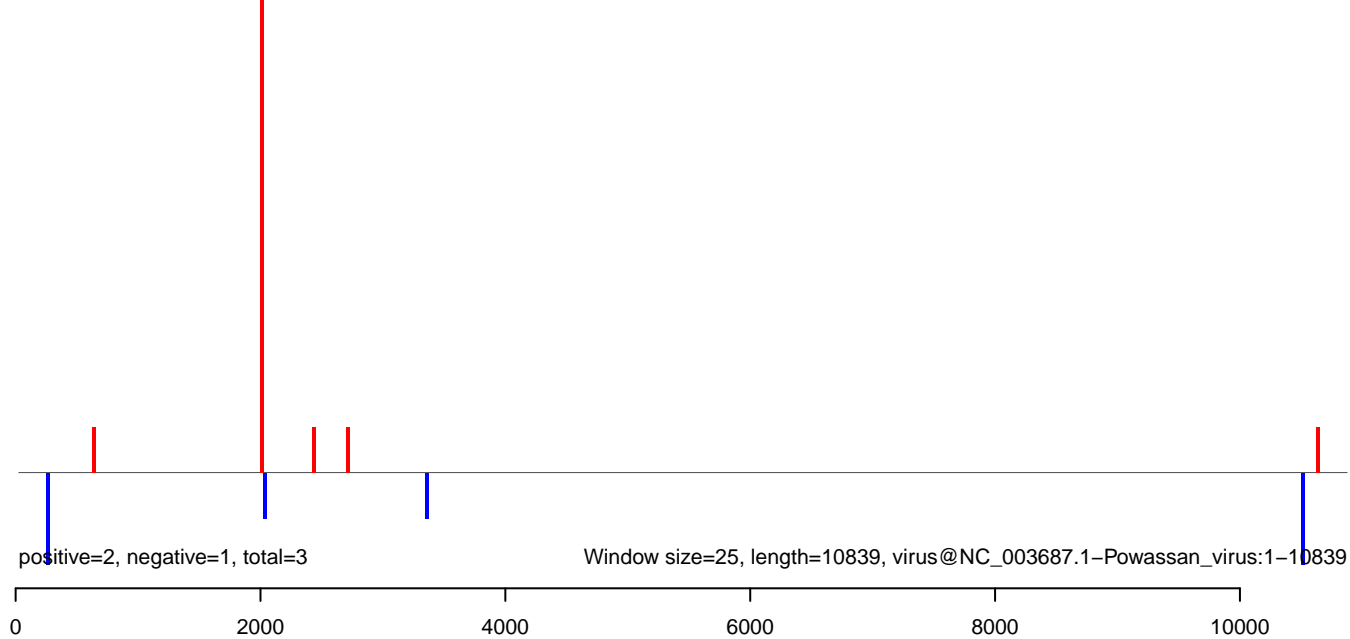
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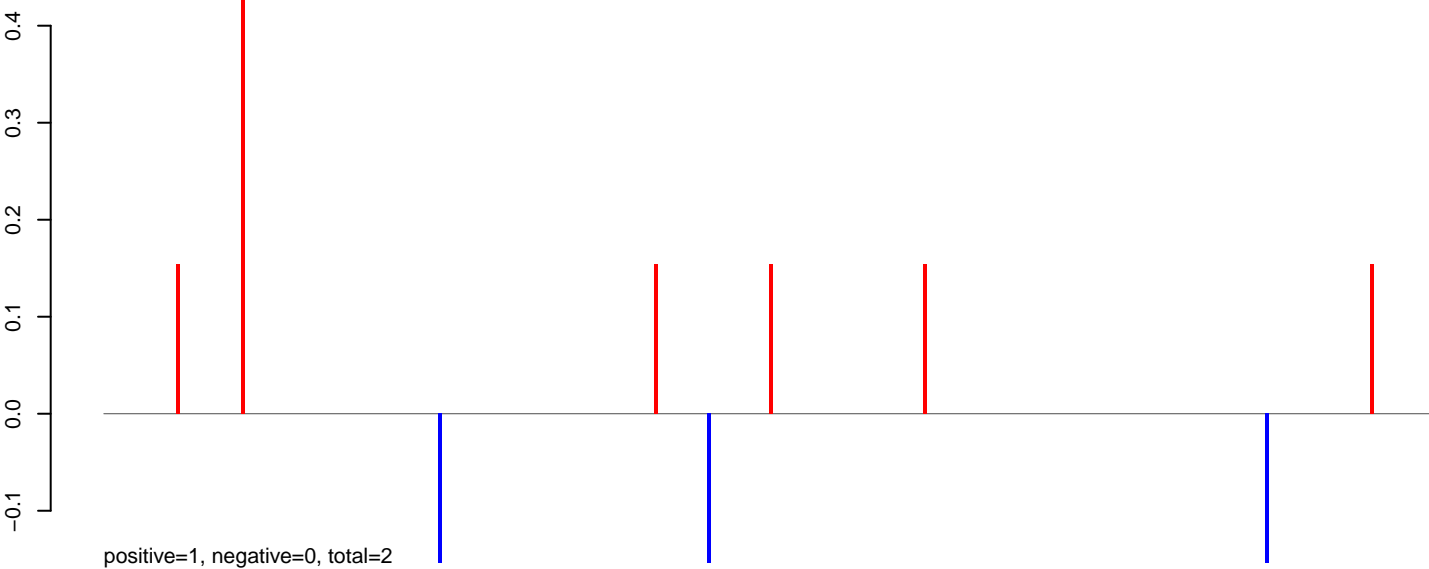
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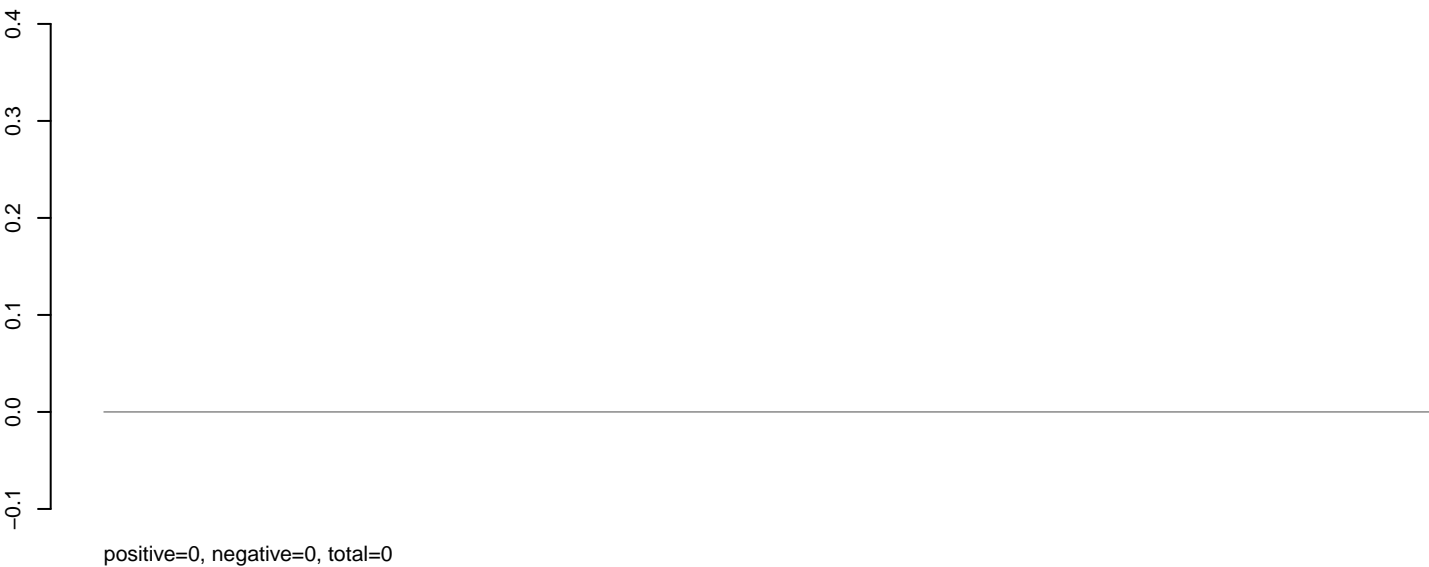
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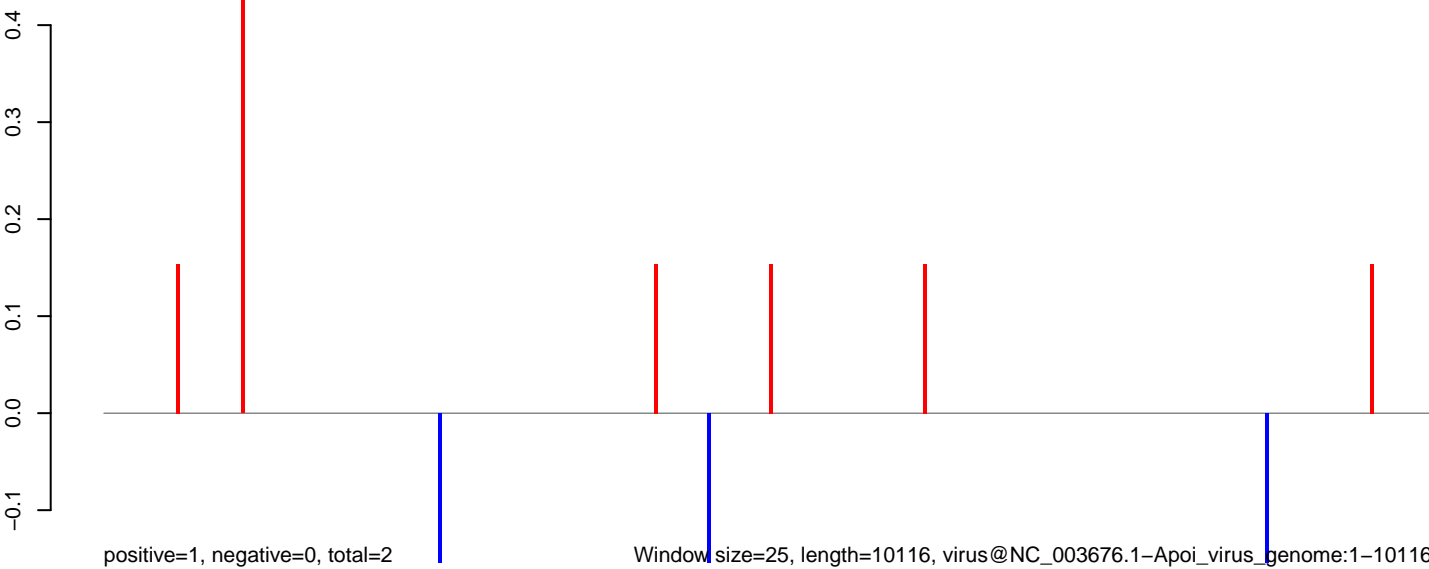
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AeAlbo_C636_6dpi_SINV1.24_35.rep

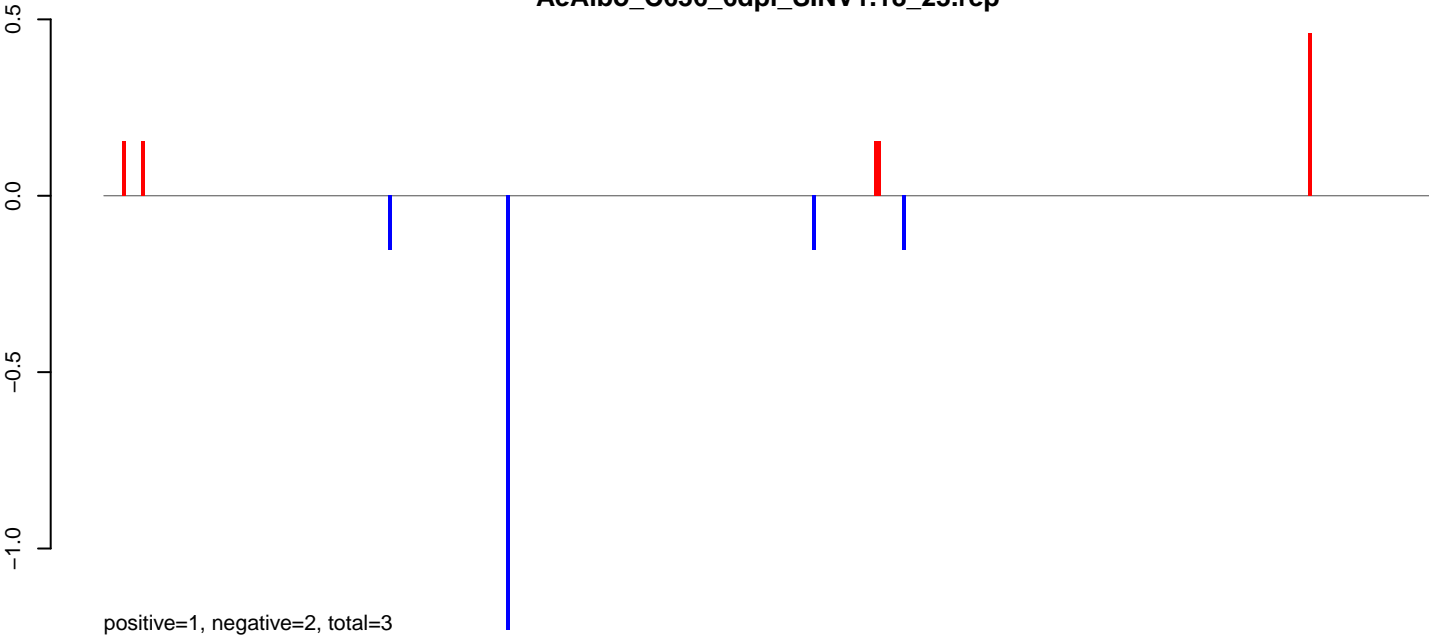


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0 2000 4000 6000 8000 10000

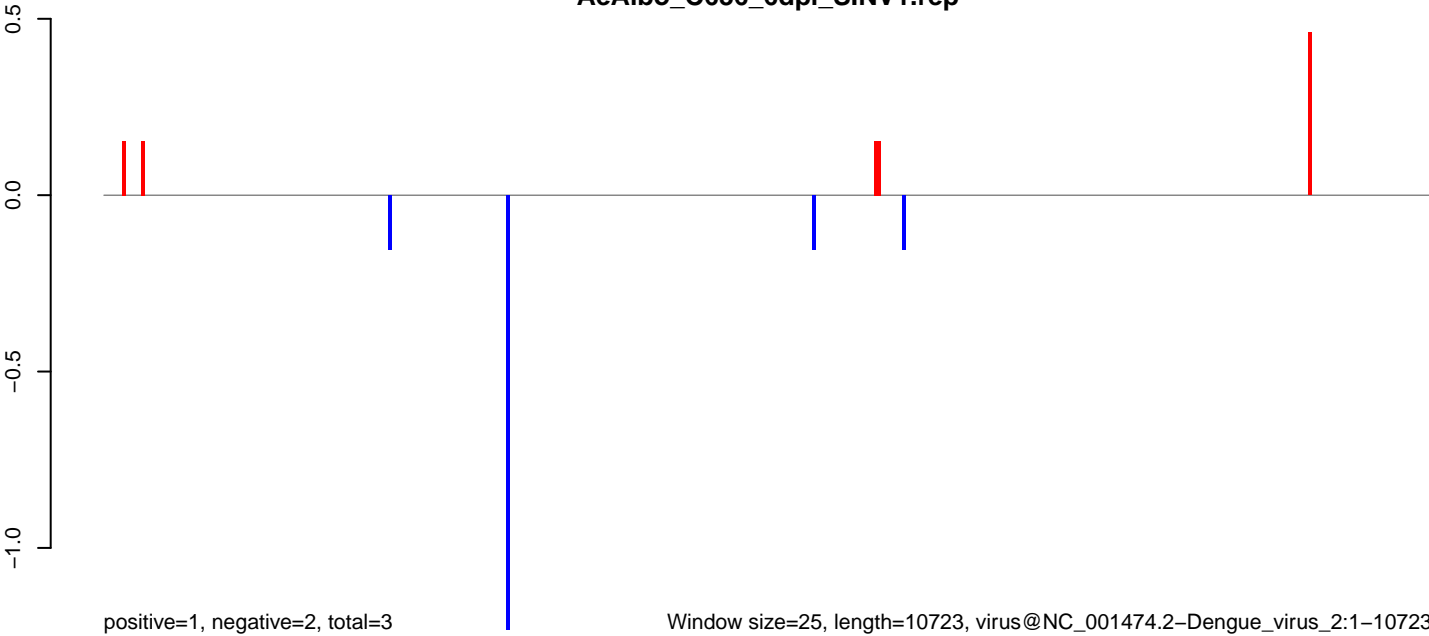
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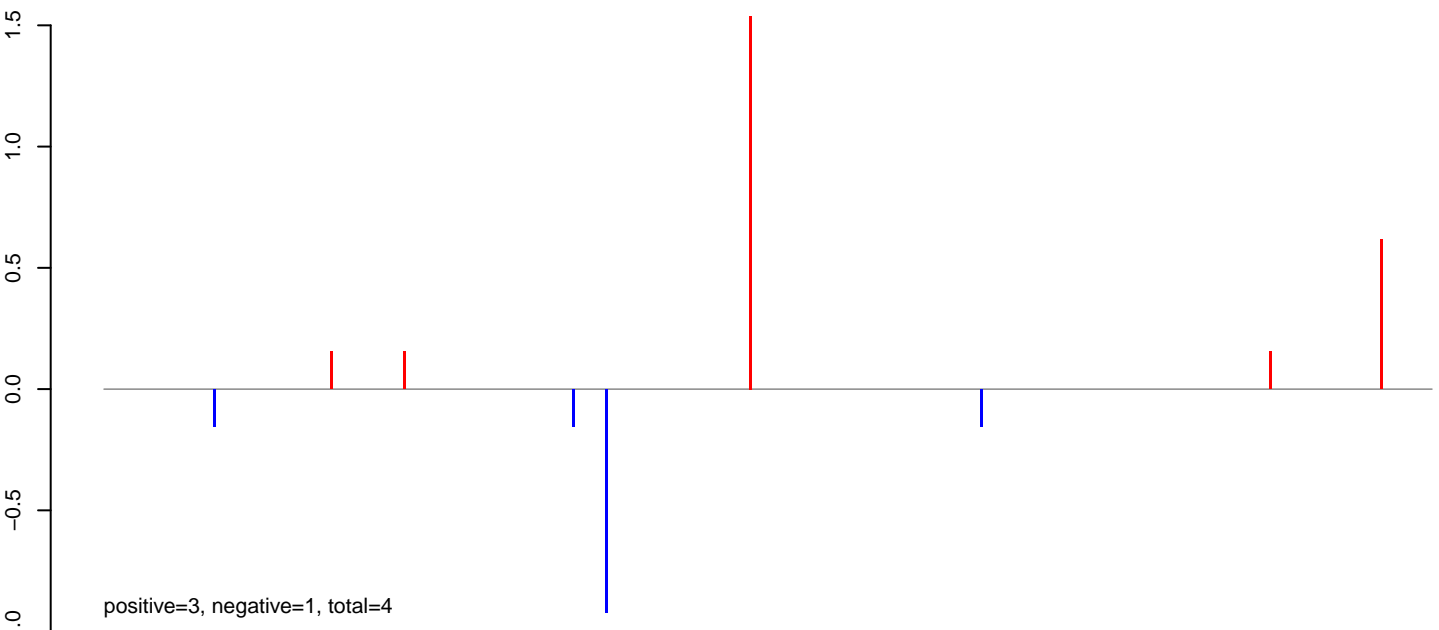
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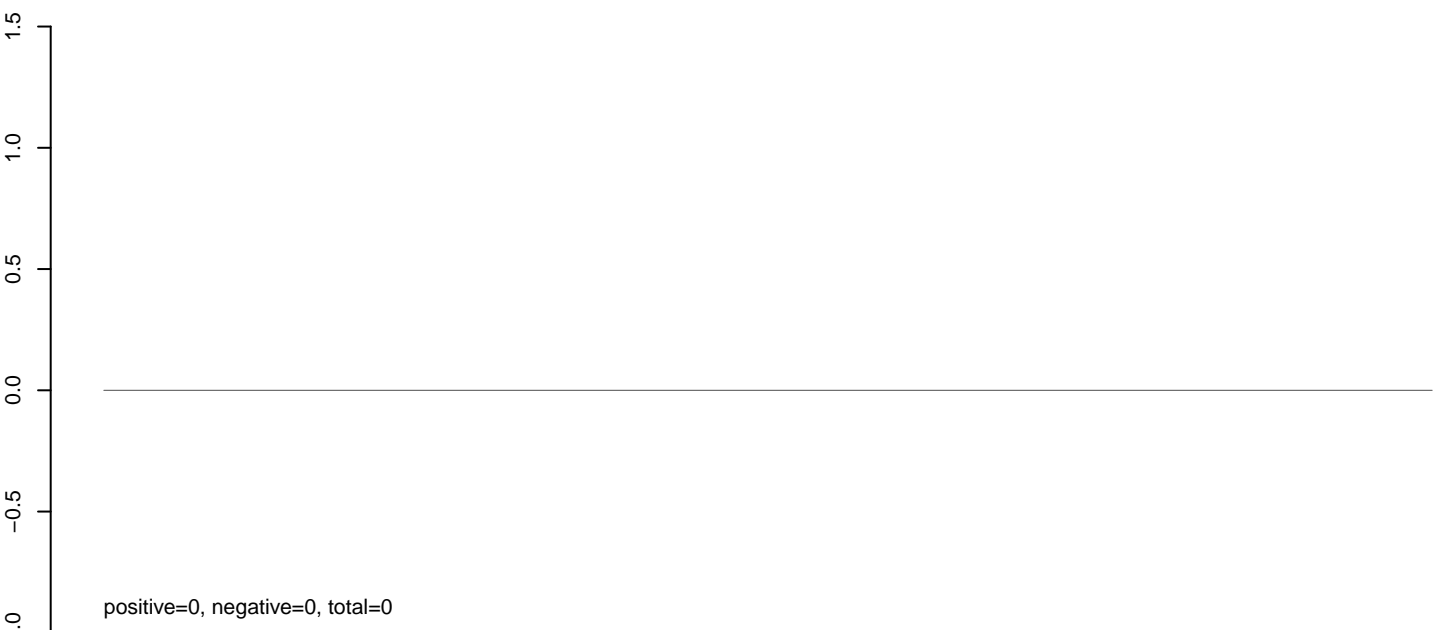
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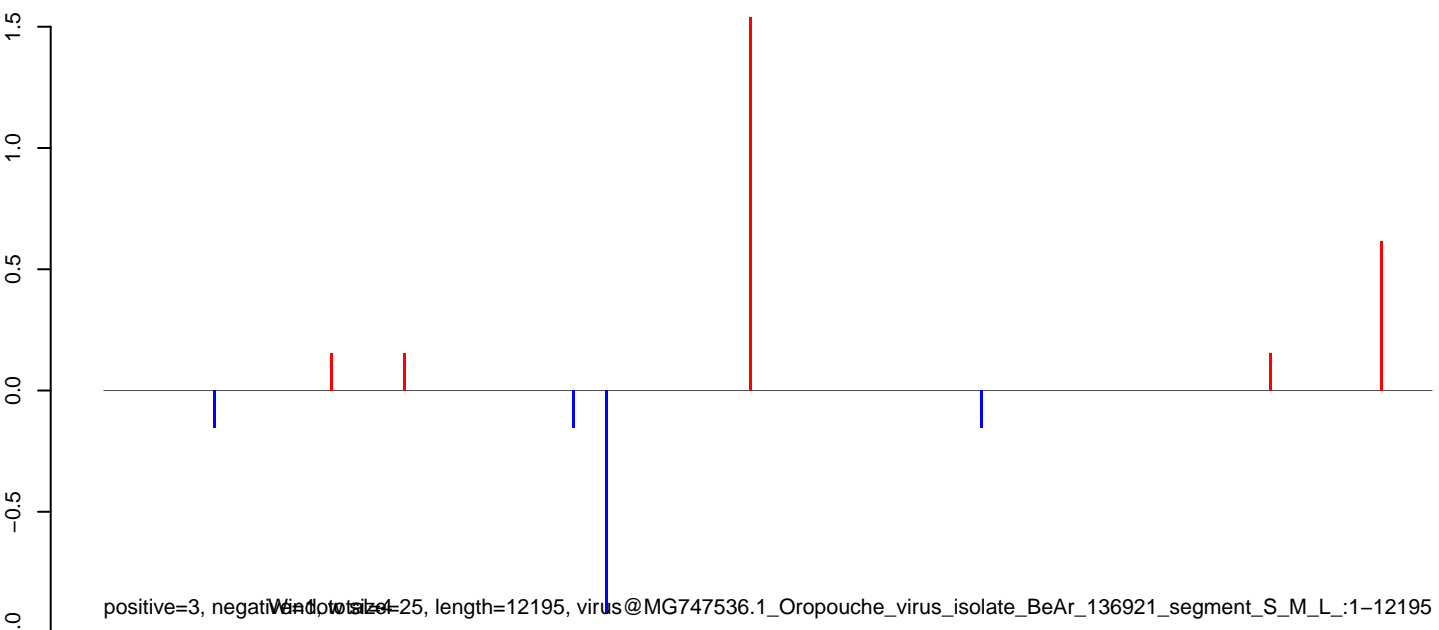
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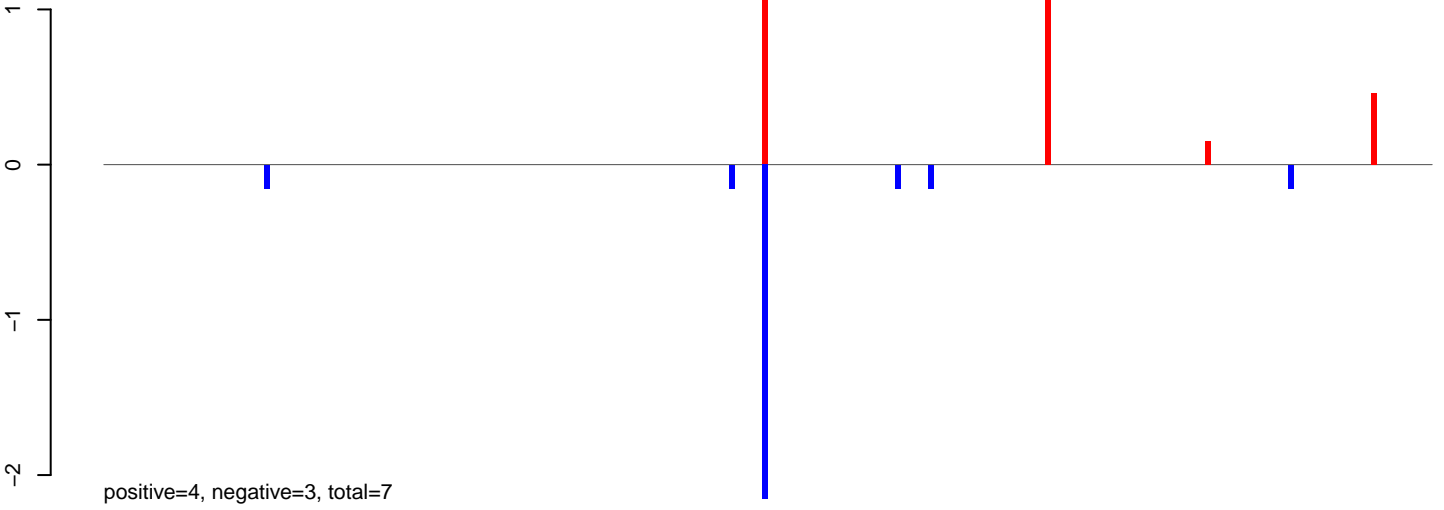
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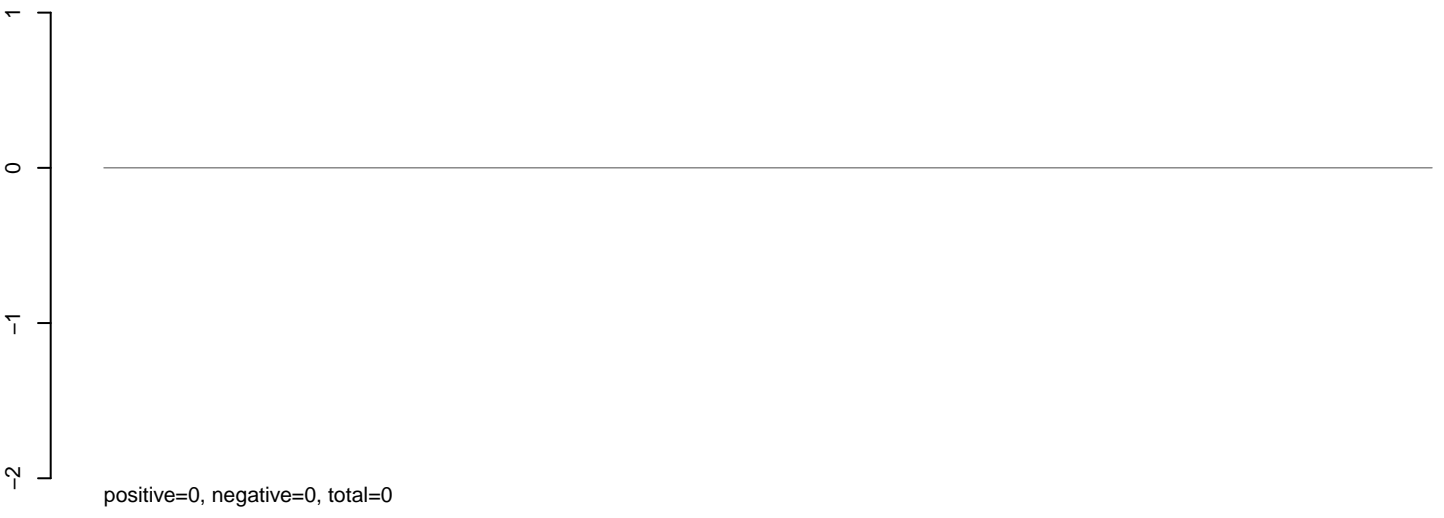
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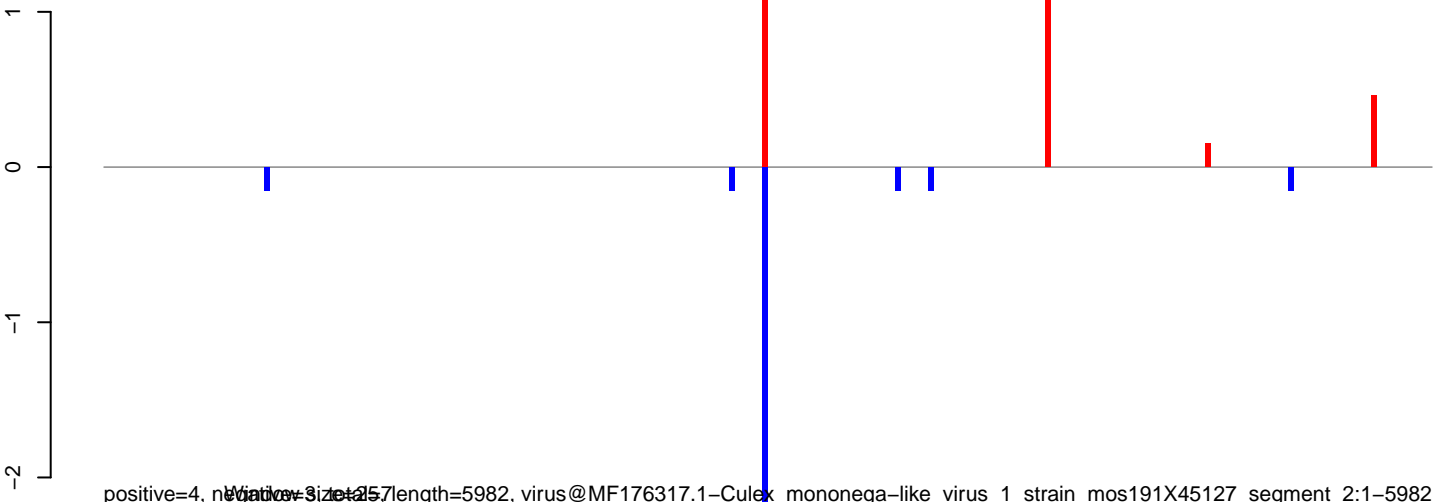
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AeAlbo_C636_6dpi_SINV1.24_35.rep



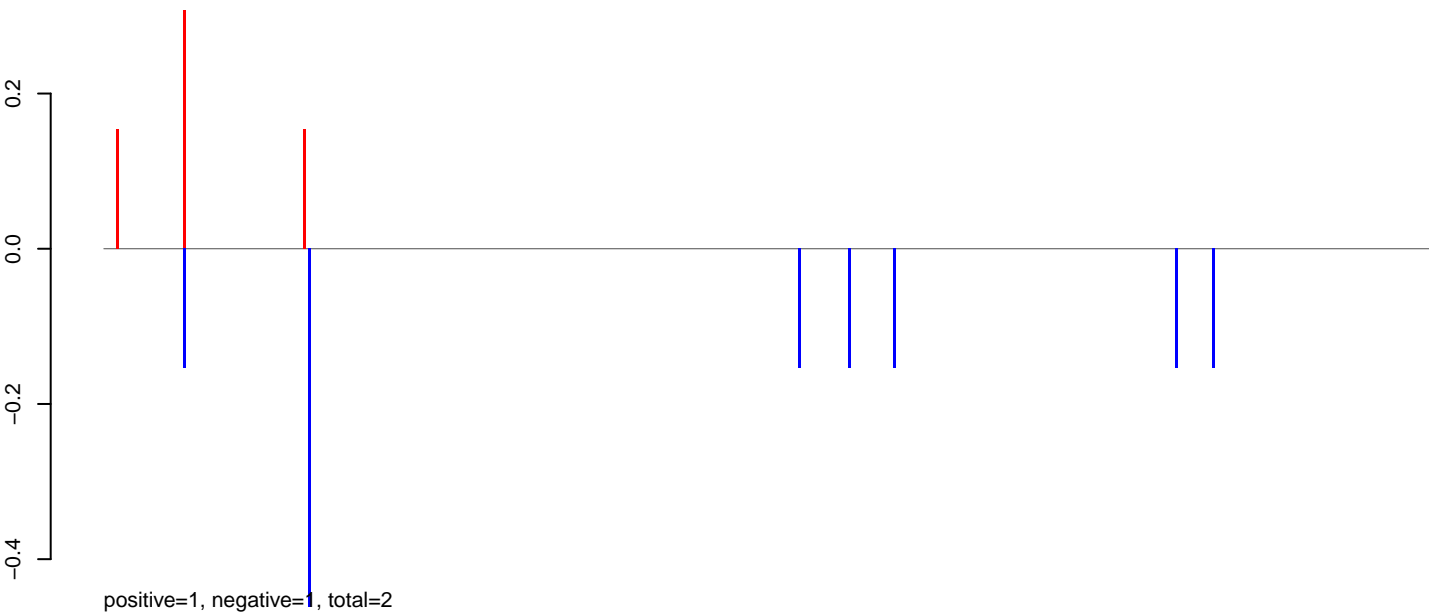
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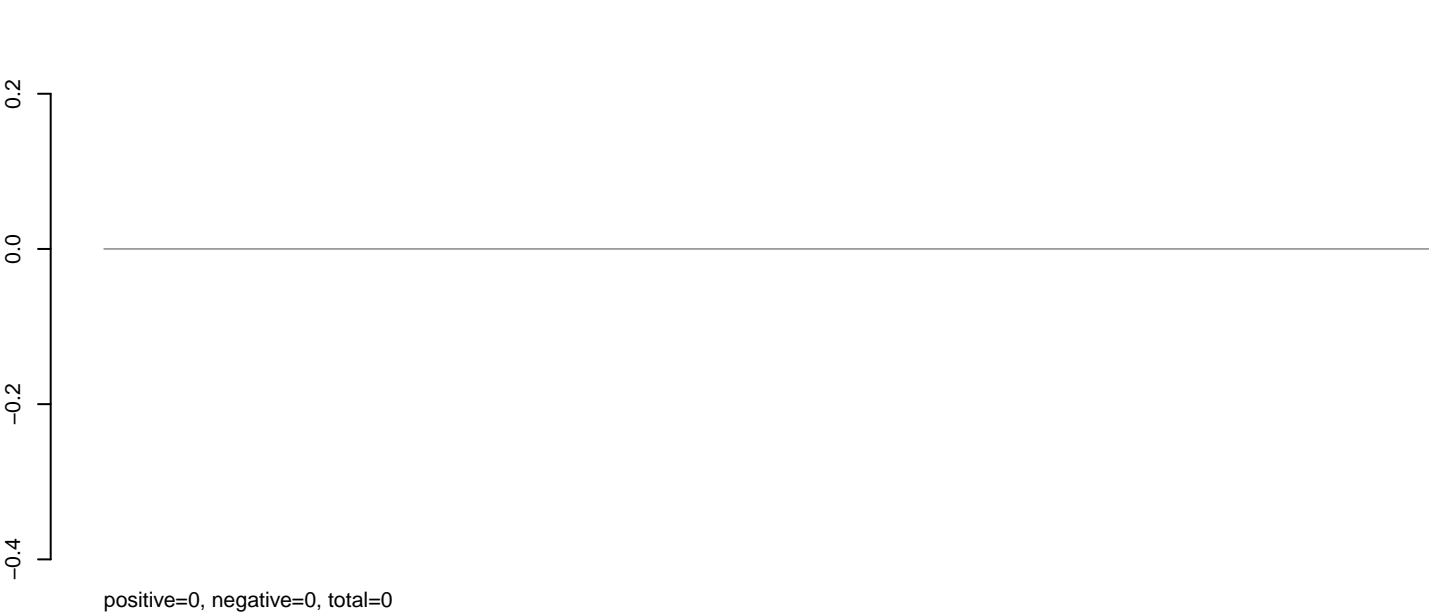
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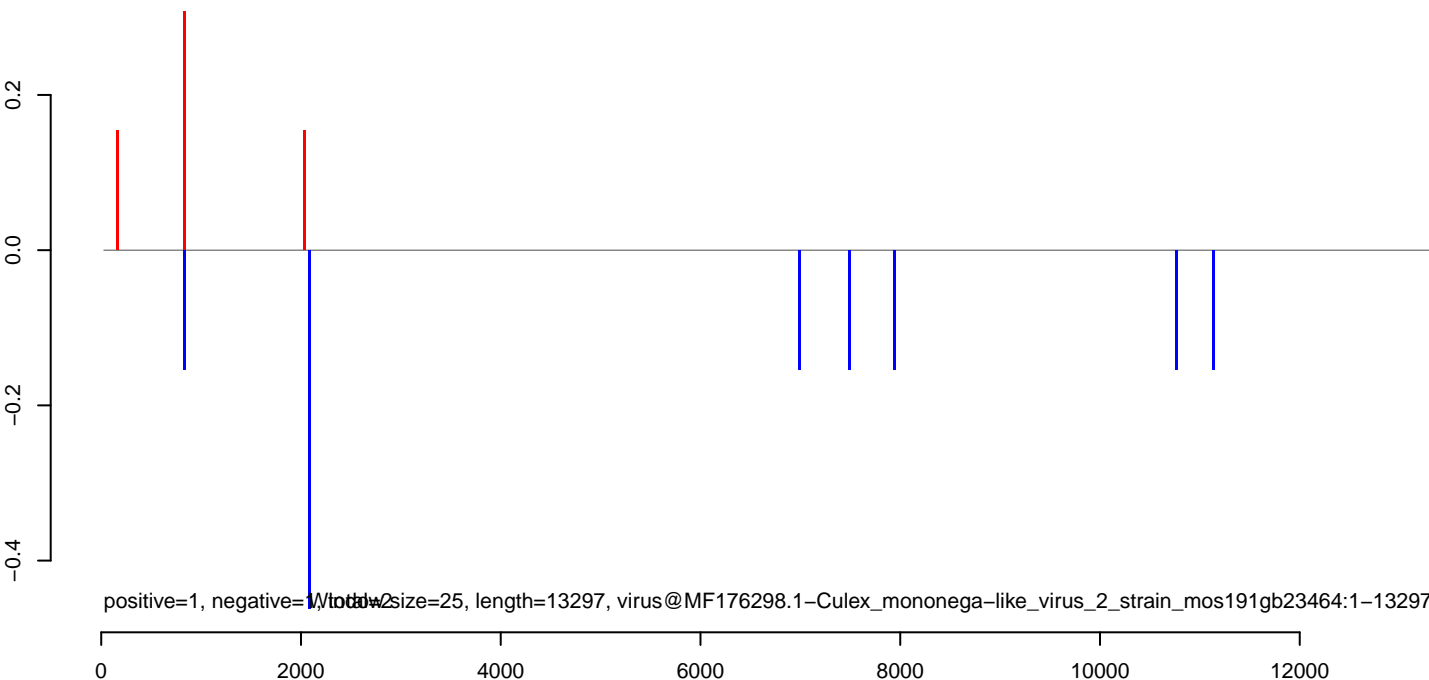
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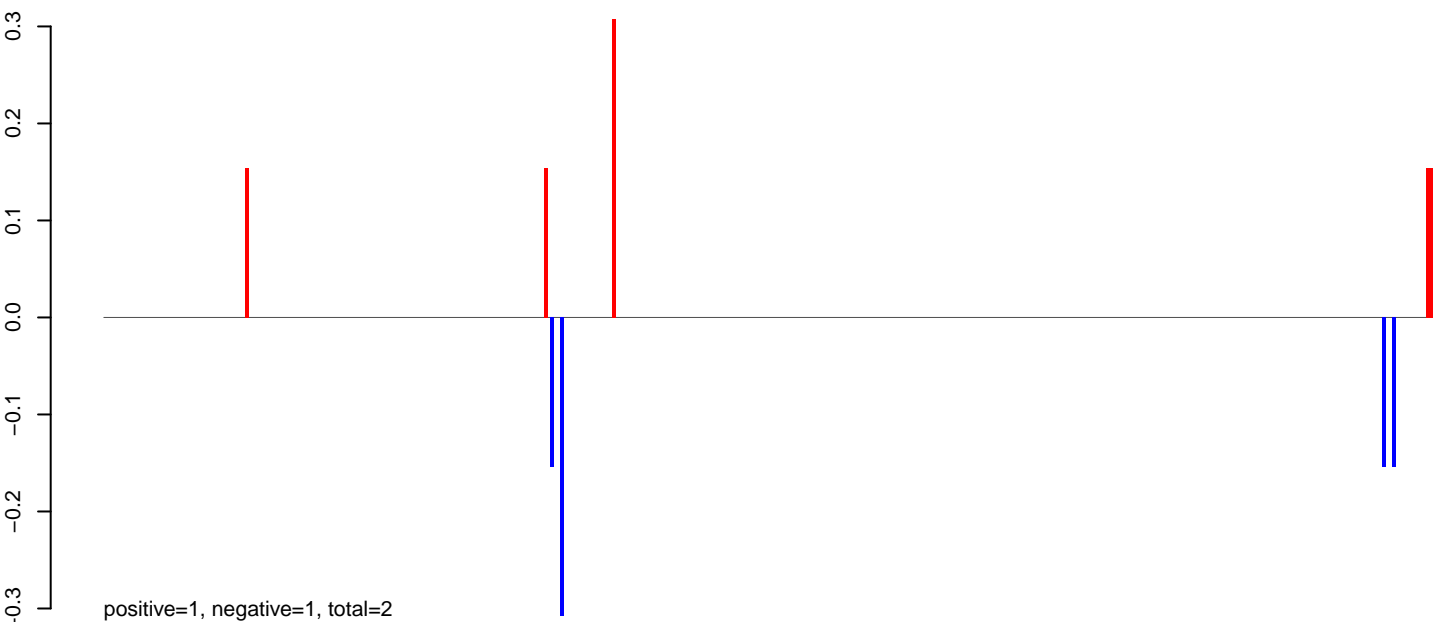
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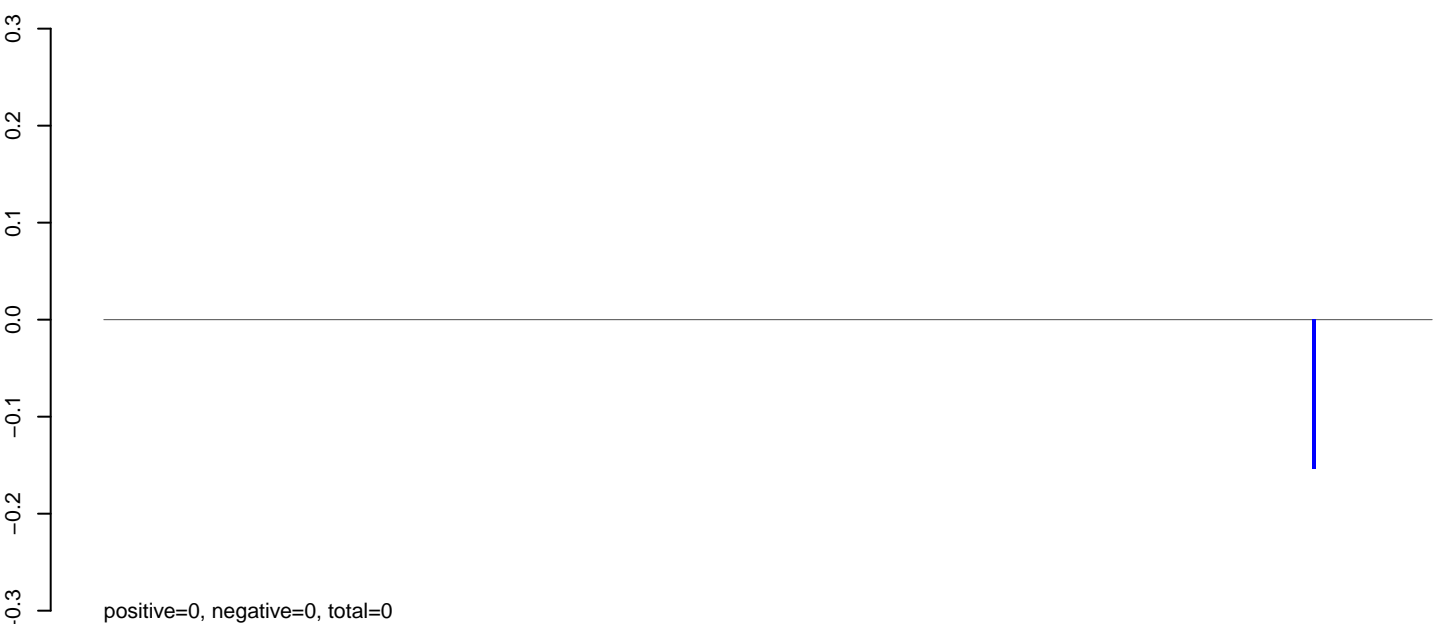
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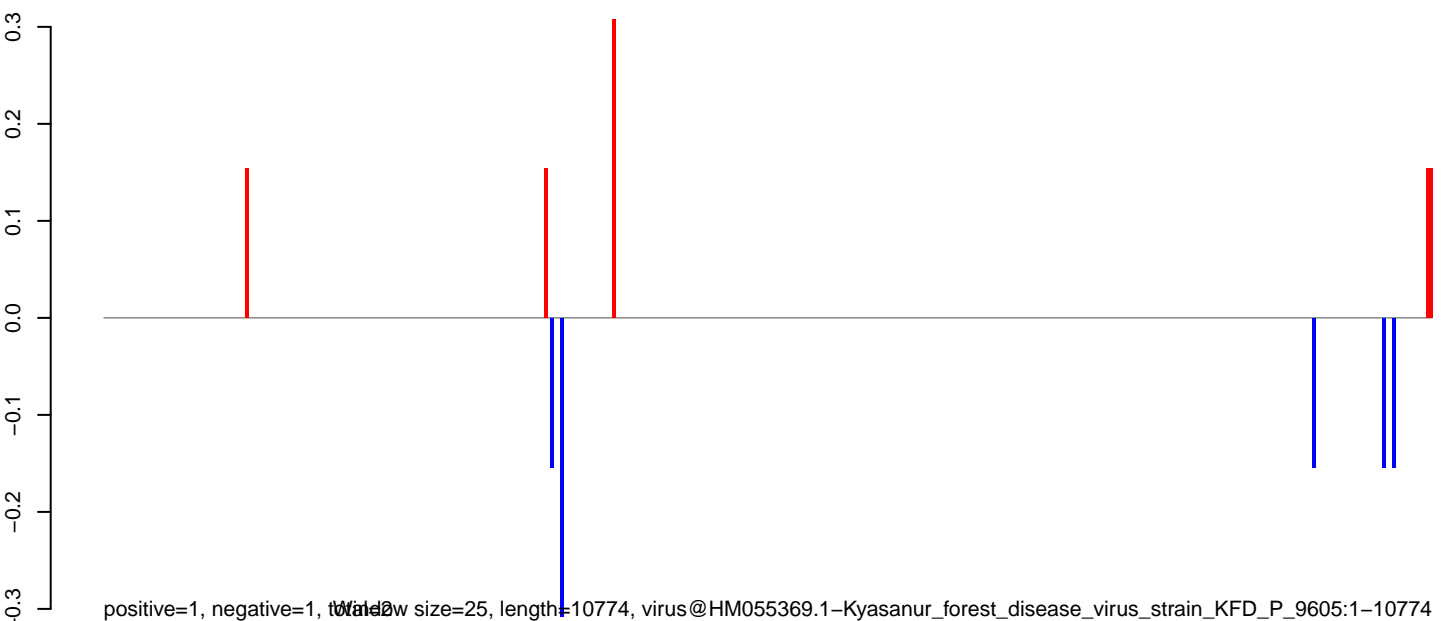
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AeAlbo_C636_6dpi_SINV1.24_35.rep



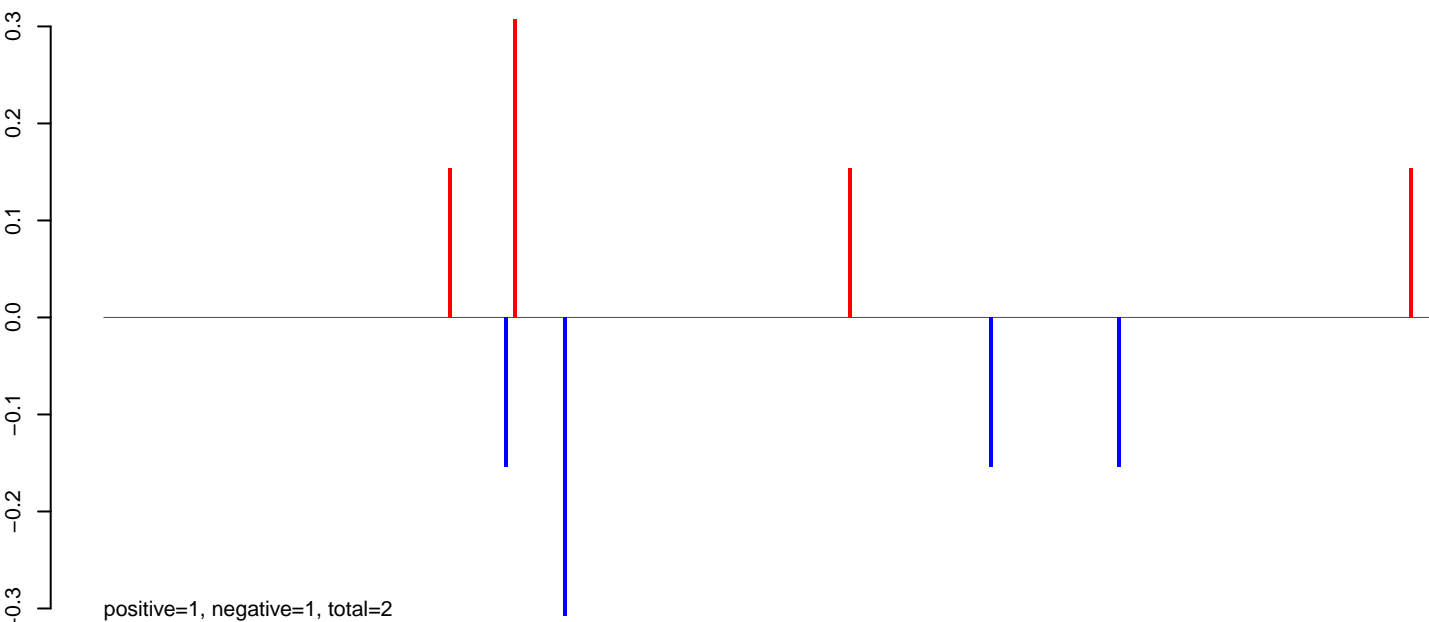
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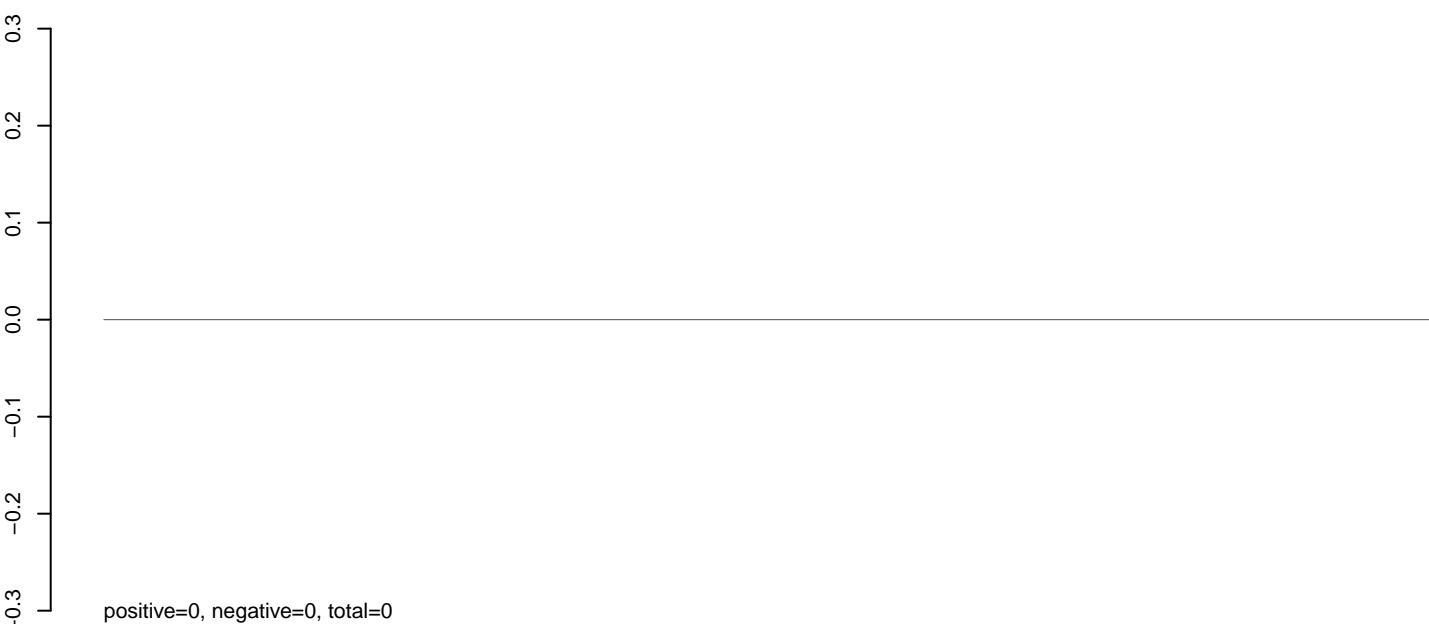
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0 2000 4000 6000 8000 10000

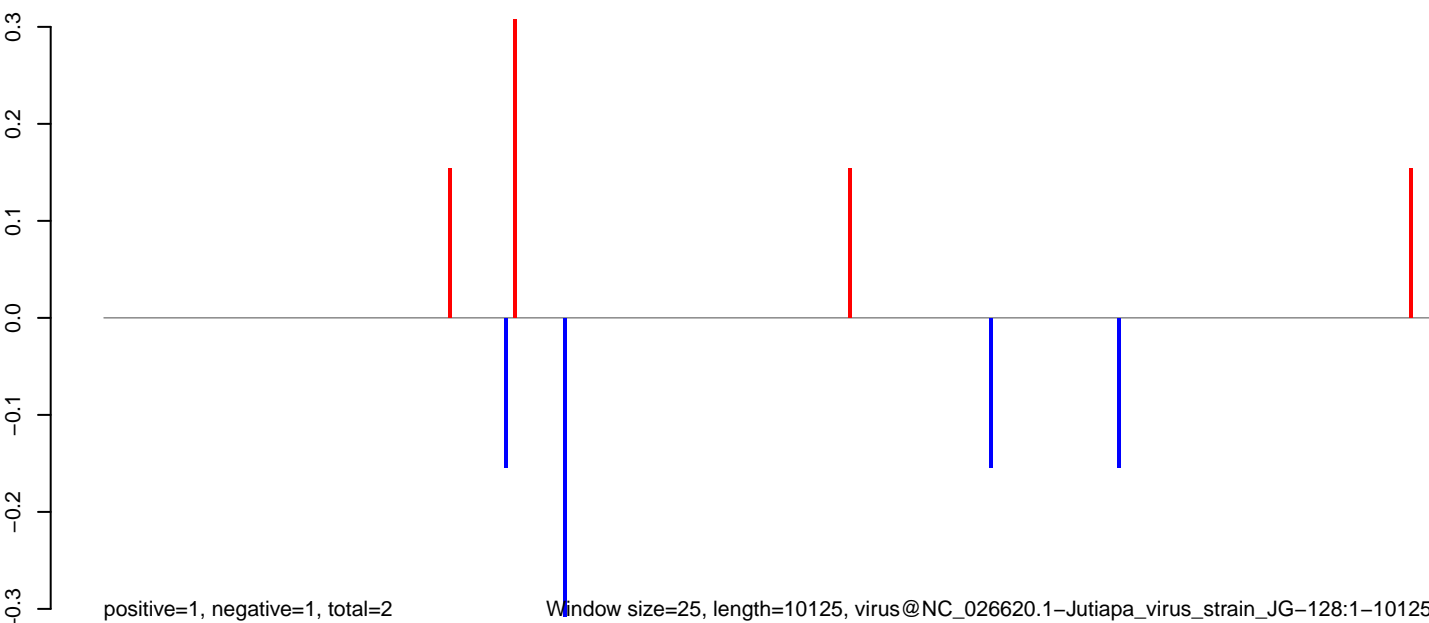
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AeAlbo_C636_6dpi_SINV1.24_35.rep



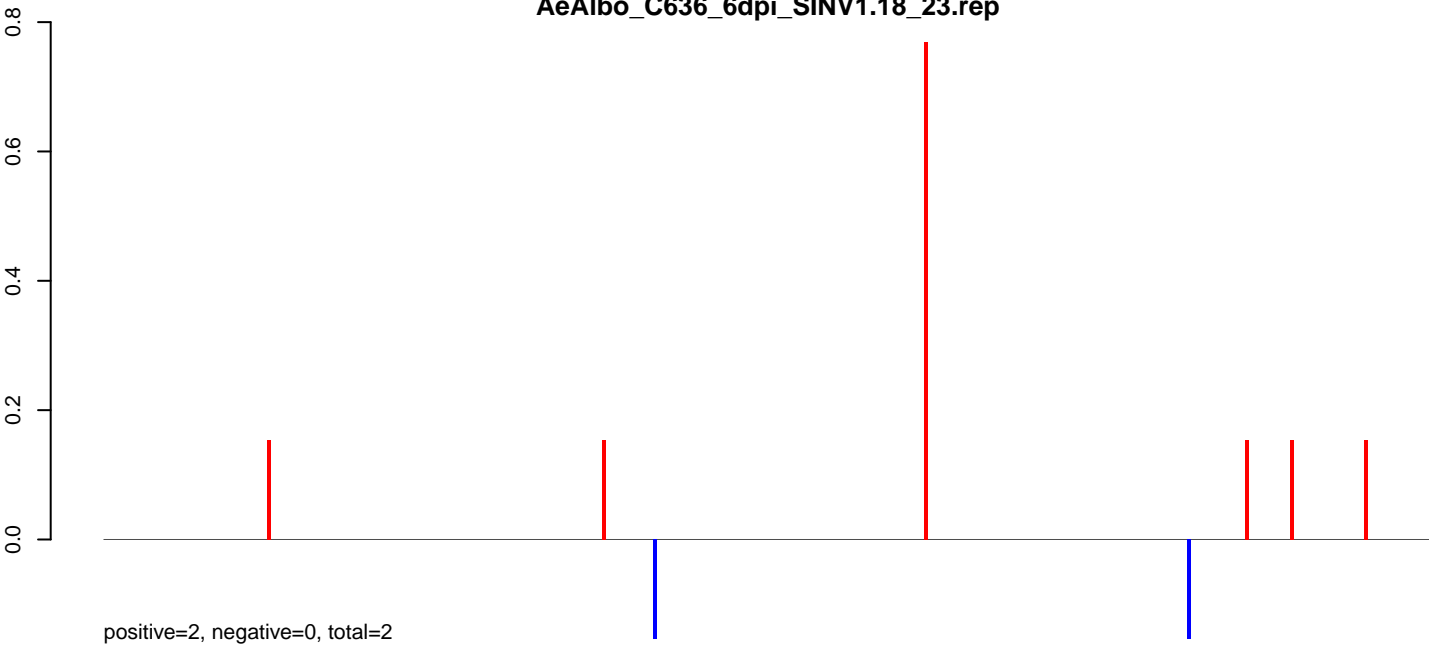
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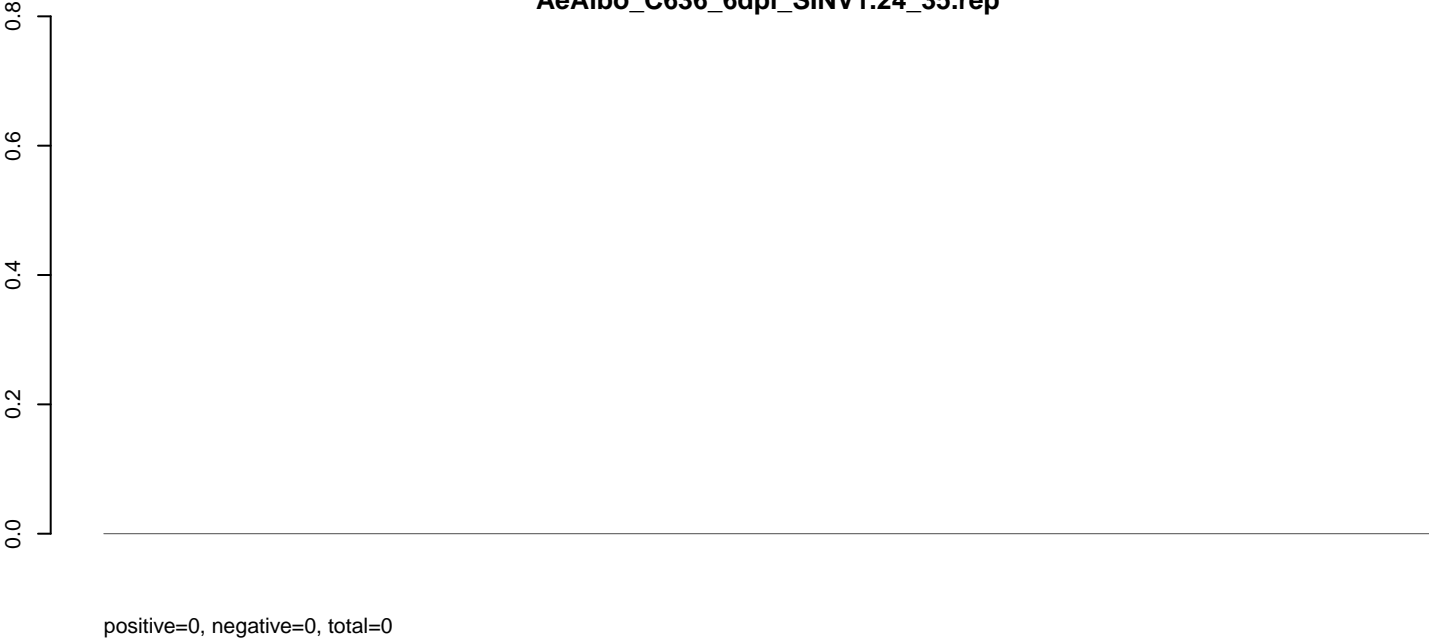
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0 2000 4000 6000 8000 10000

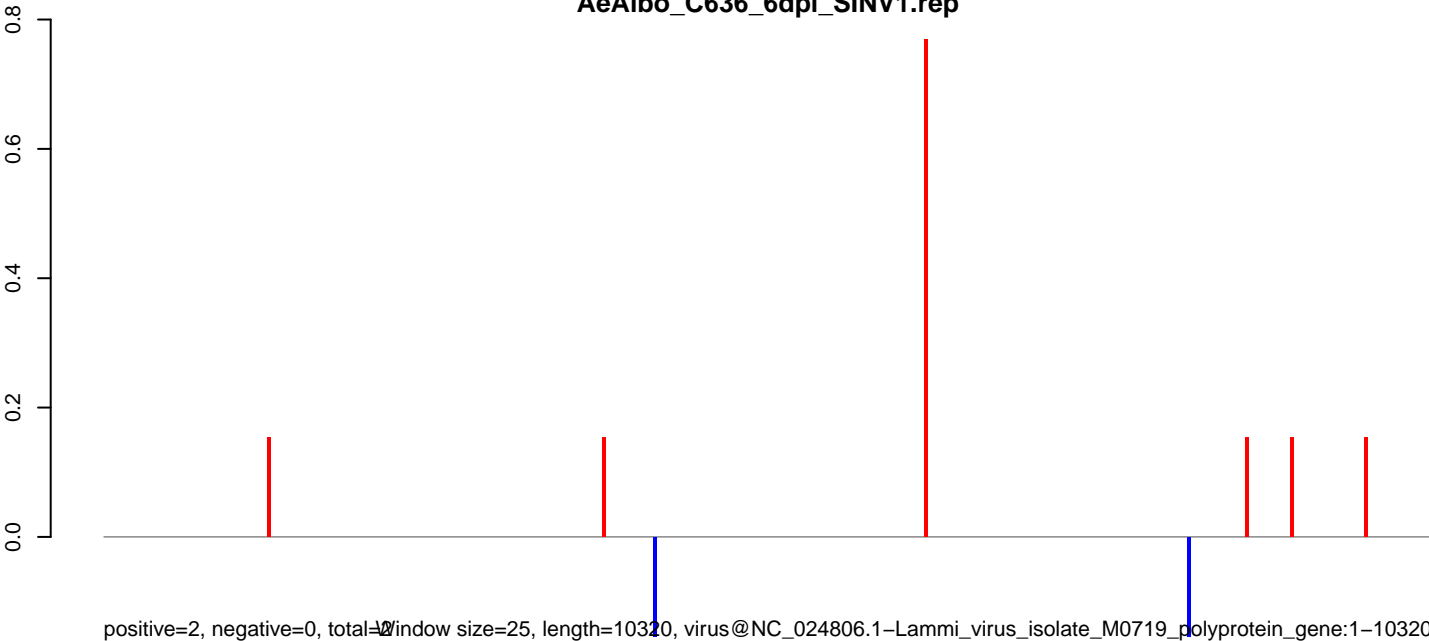
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AeAlbo_C636_6dpi_SINV1.24_35.rep



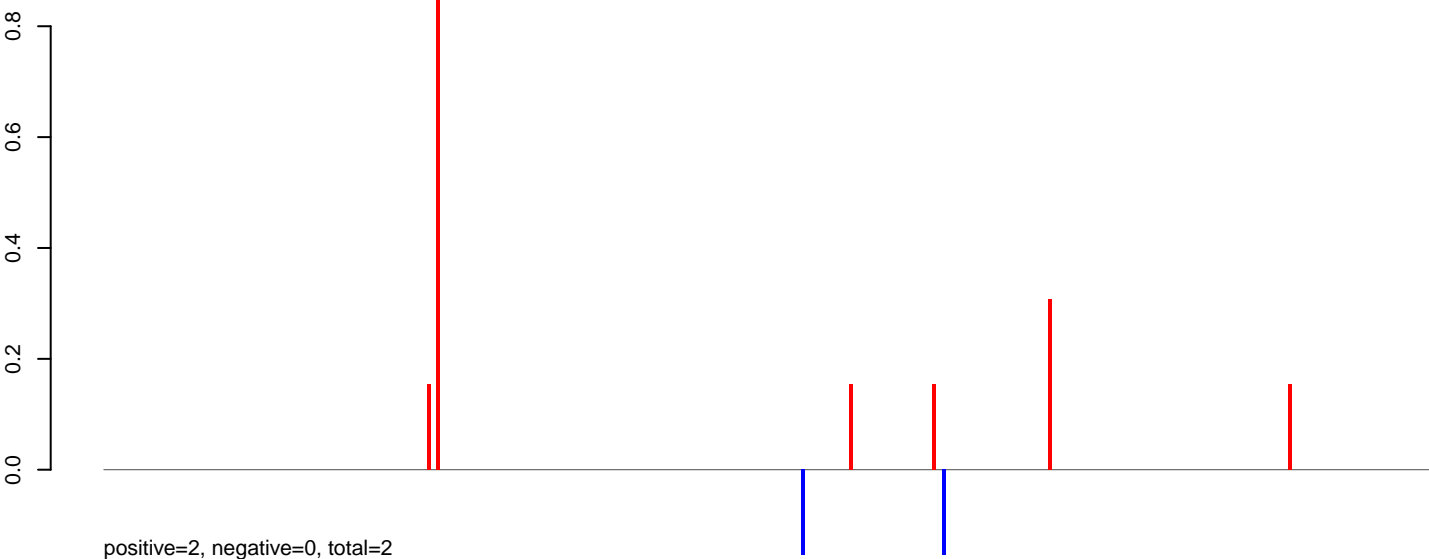
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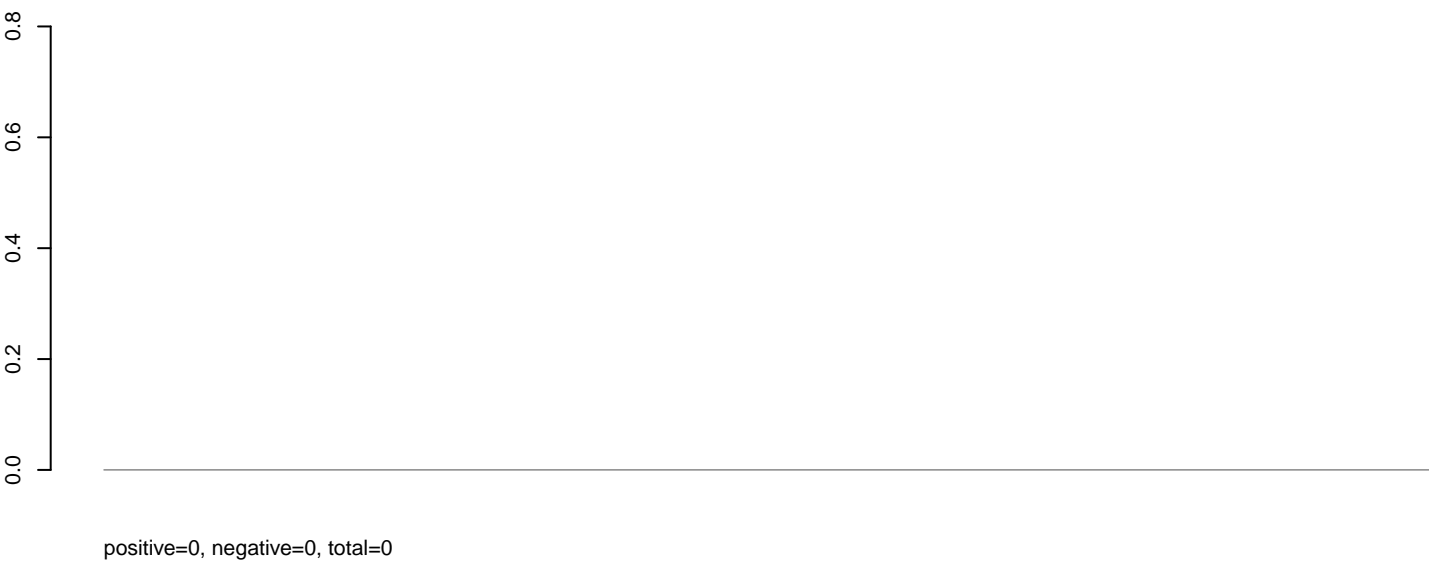
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0 2000 4000 6000 8000 10000

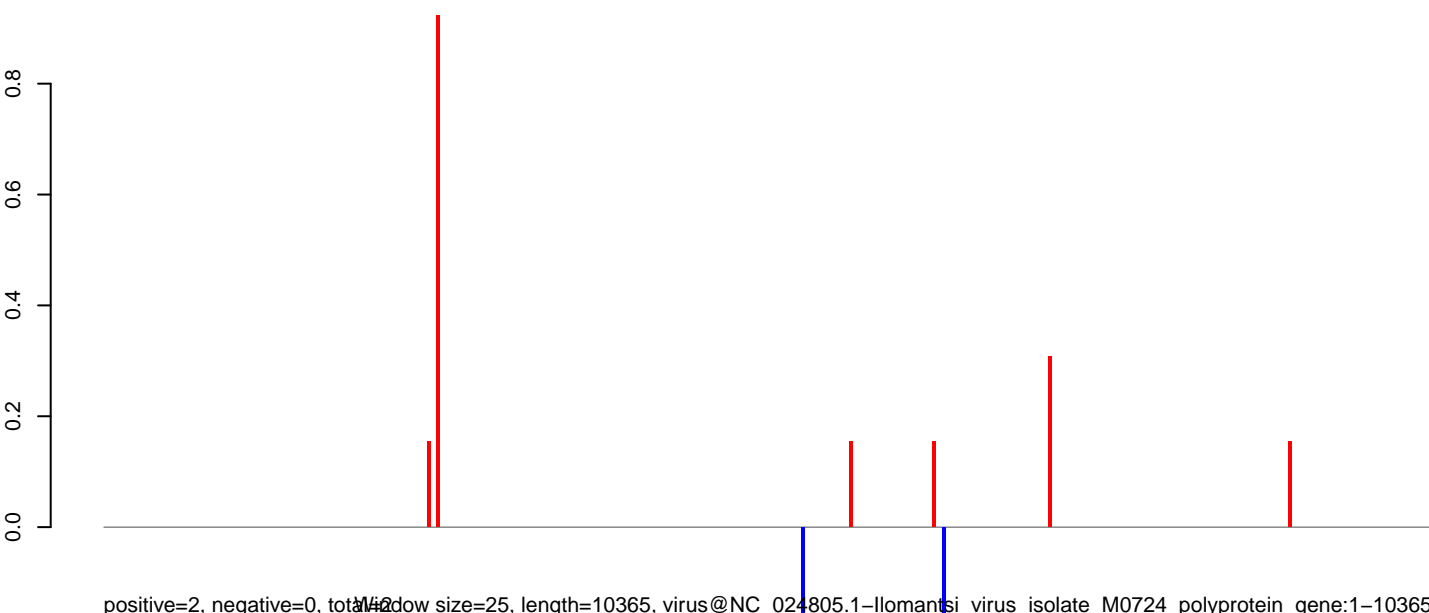
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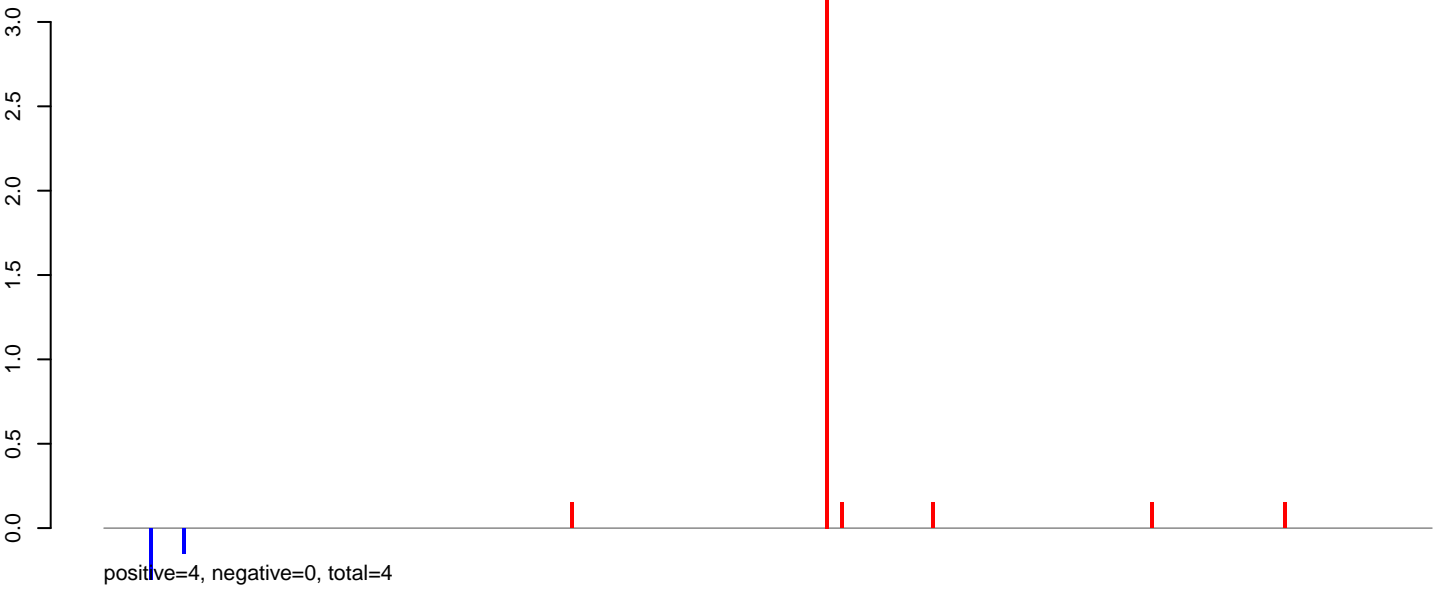
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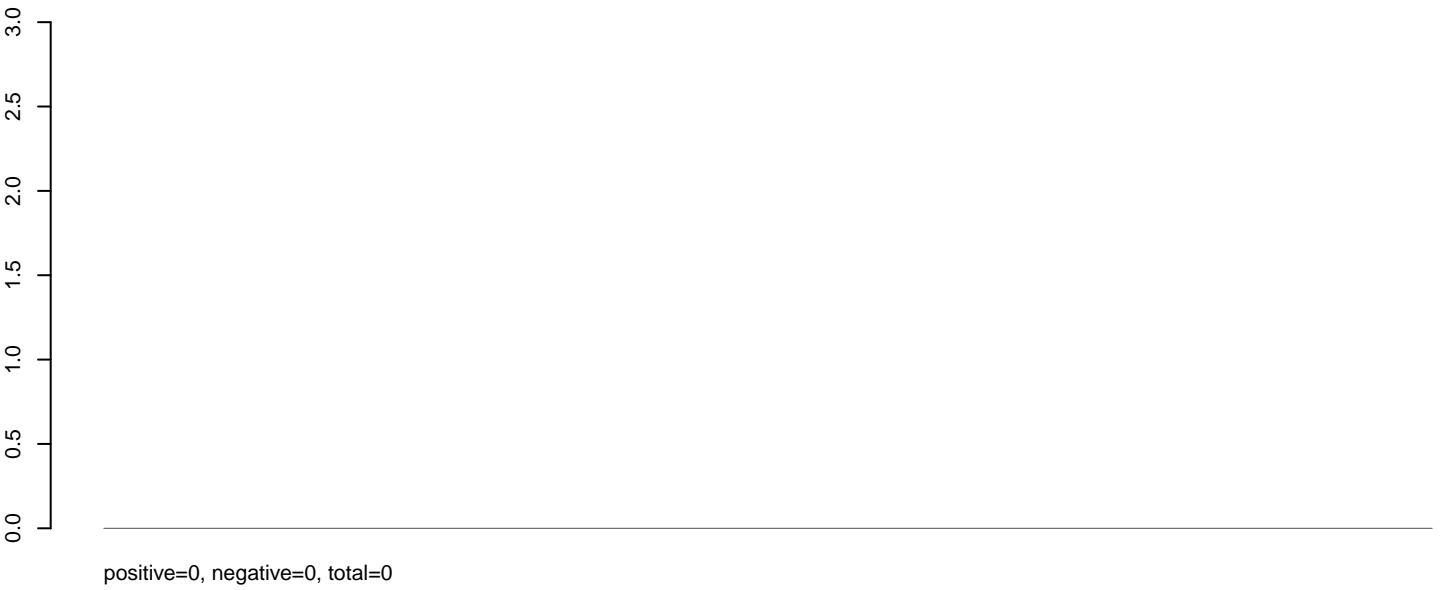
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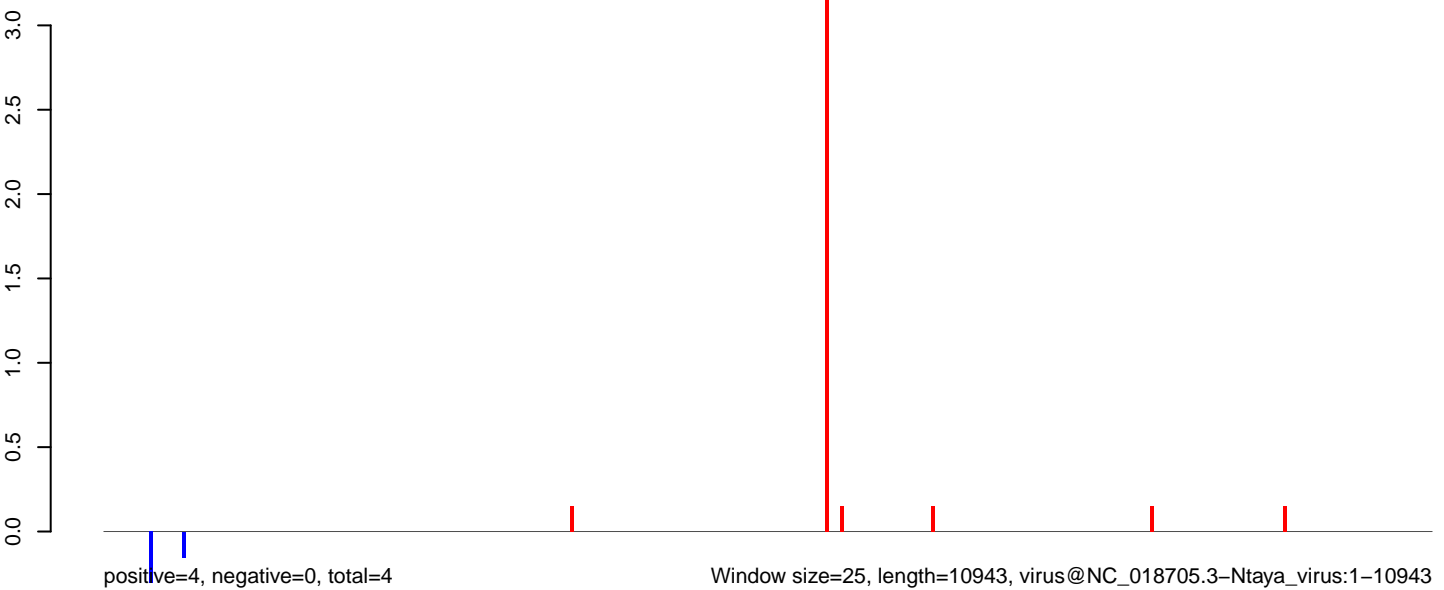
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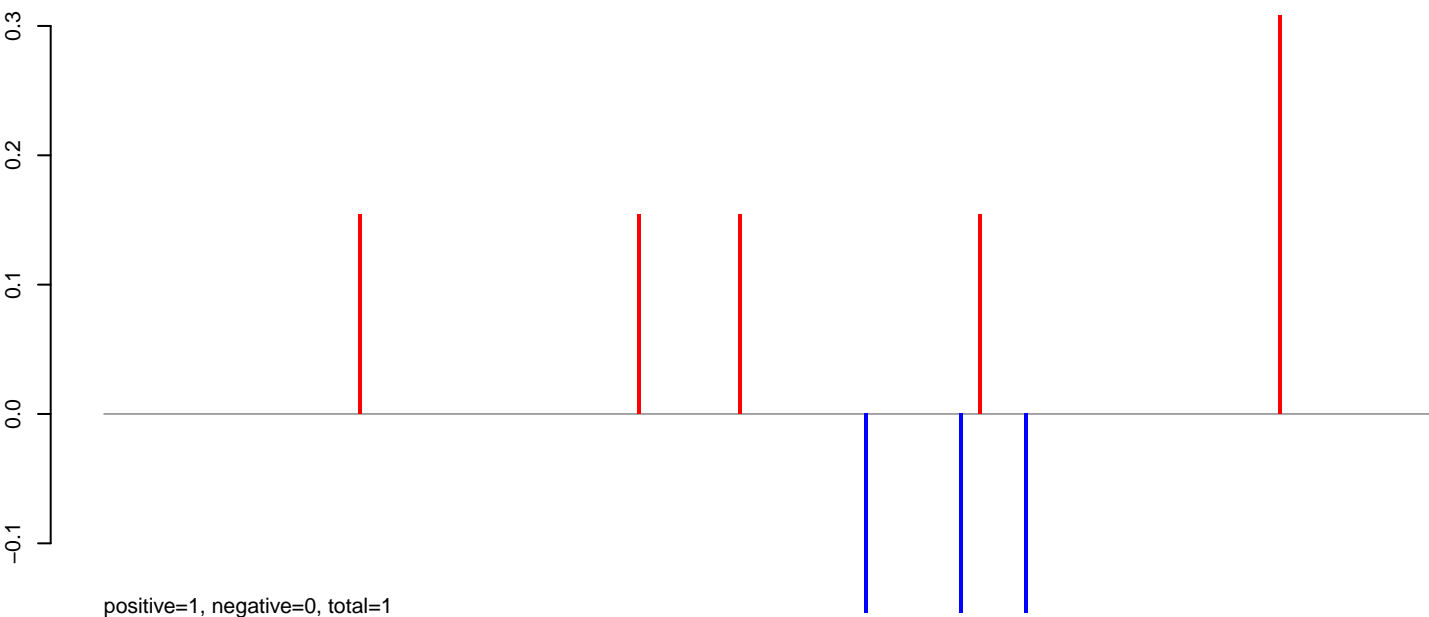
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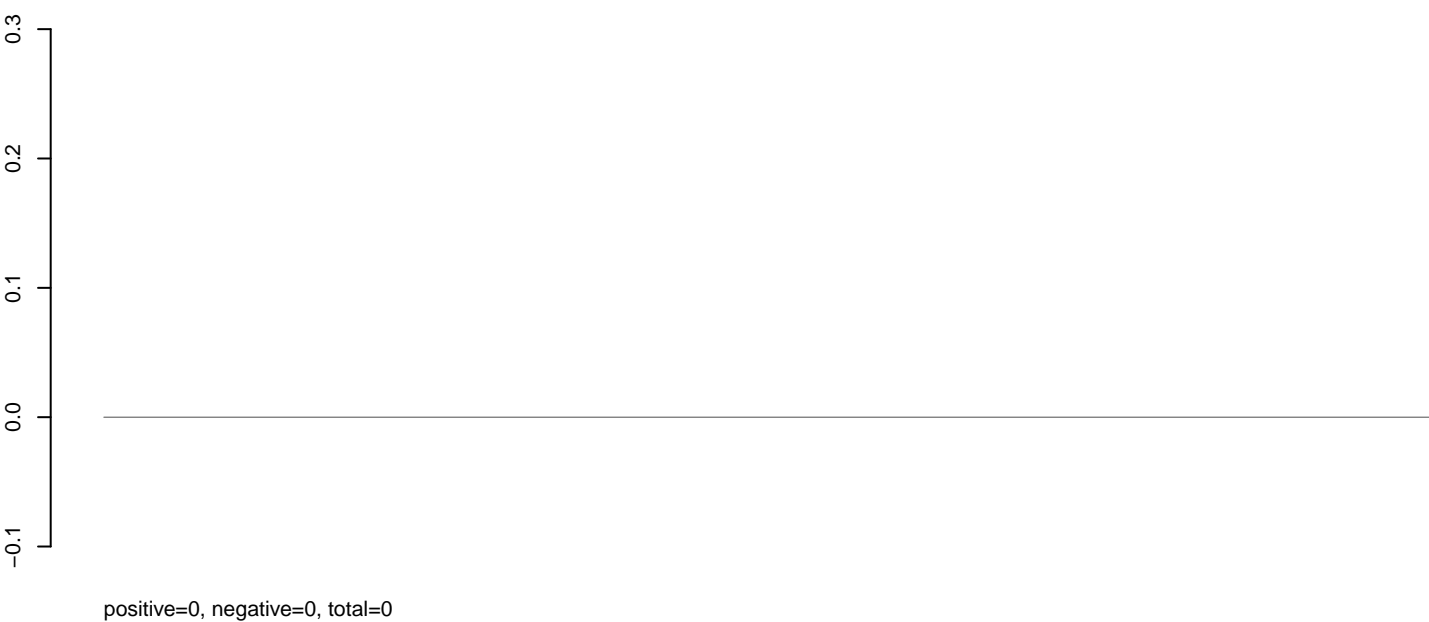
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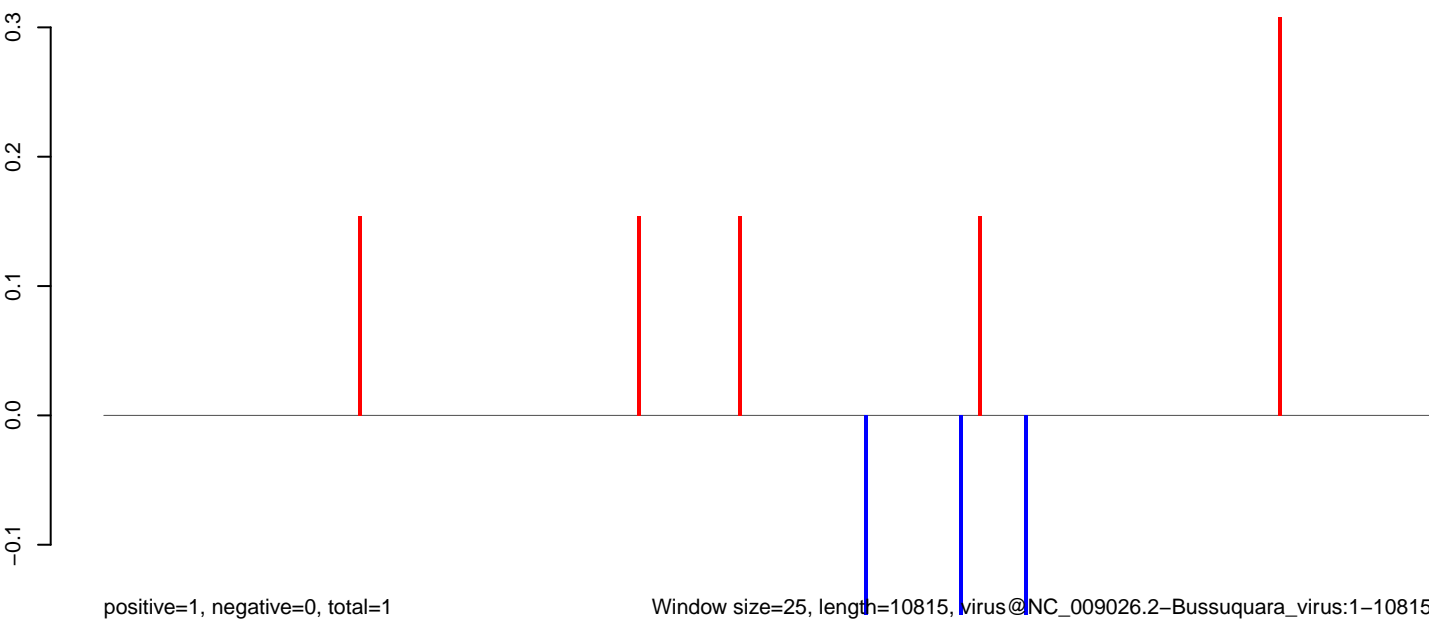
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



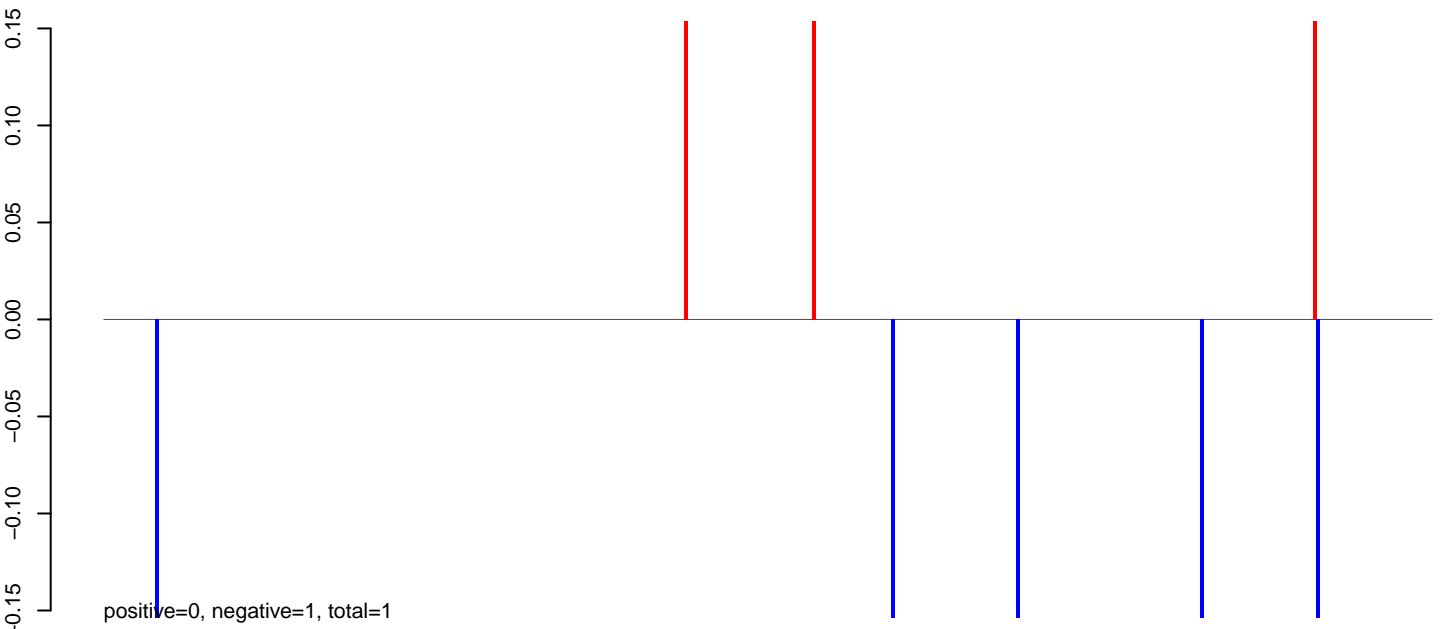
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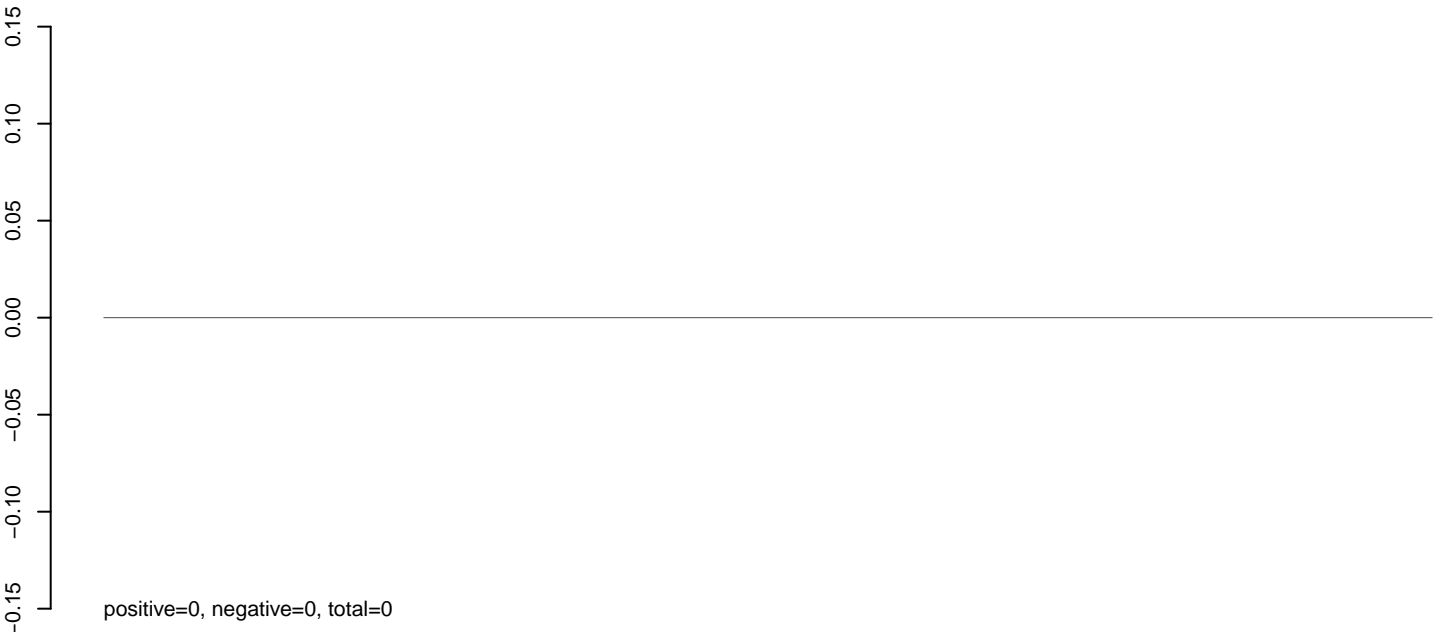
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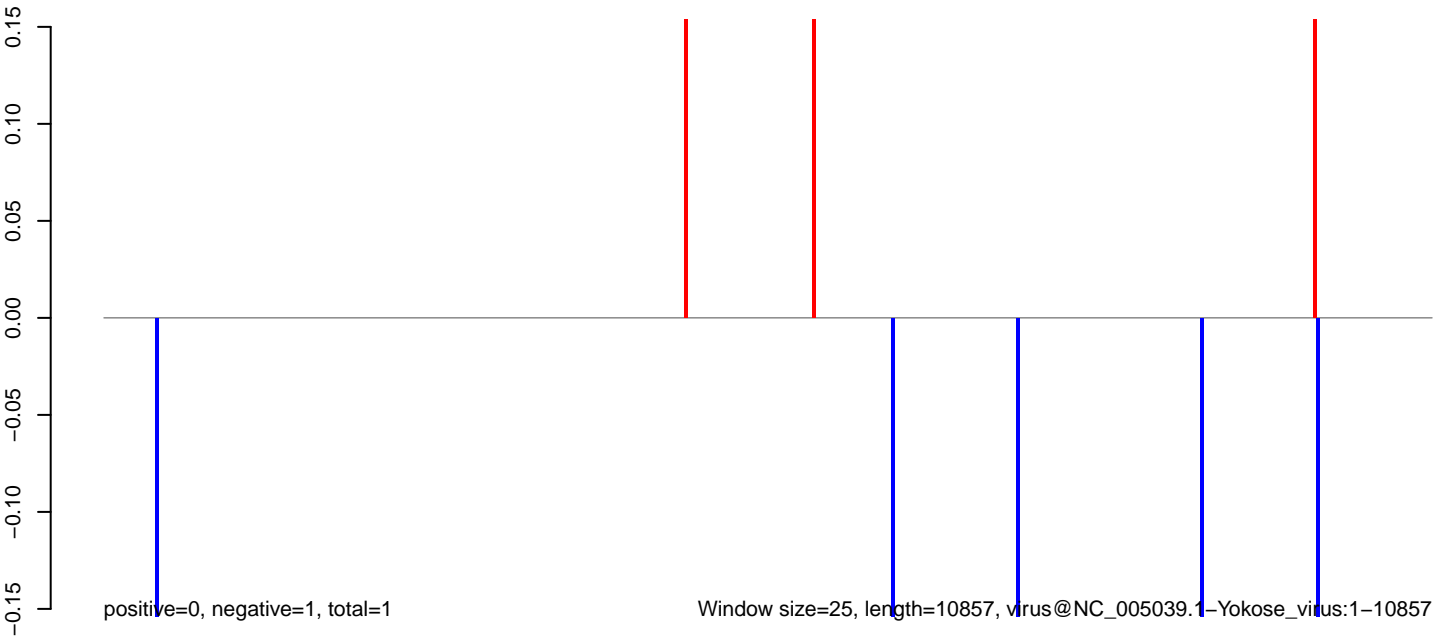
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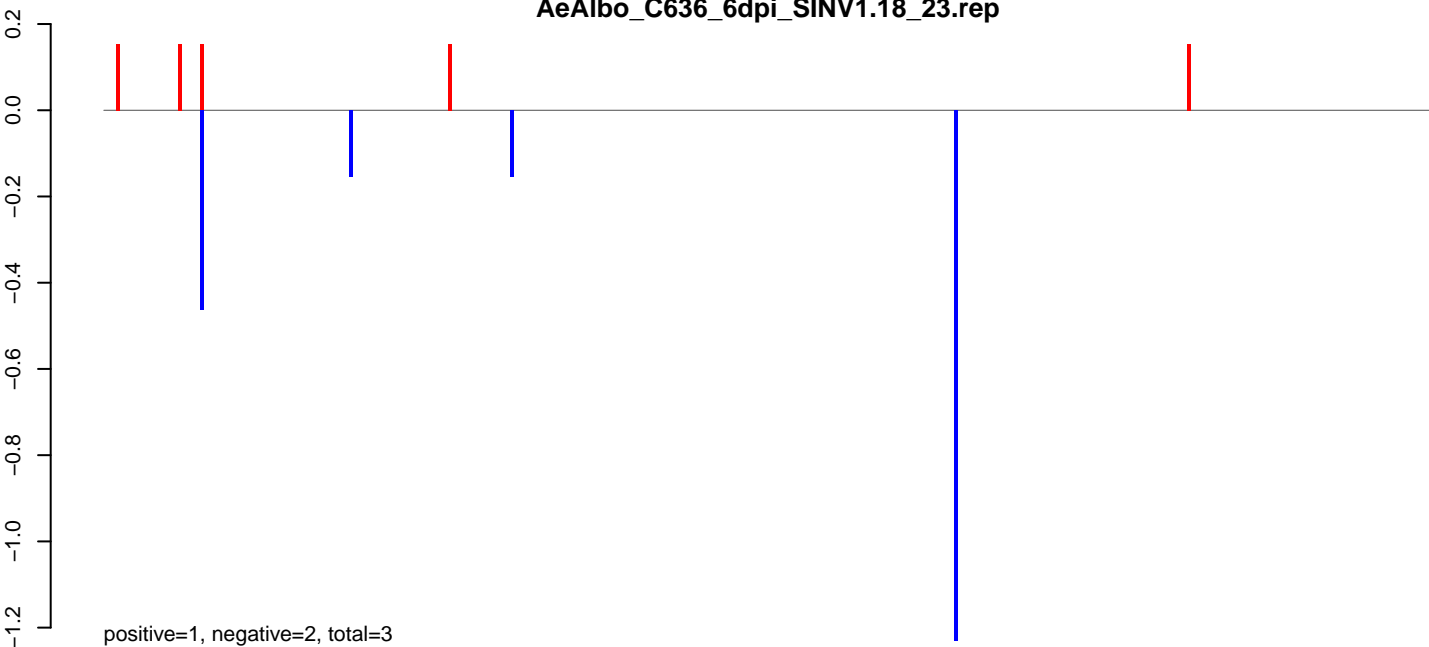
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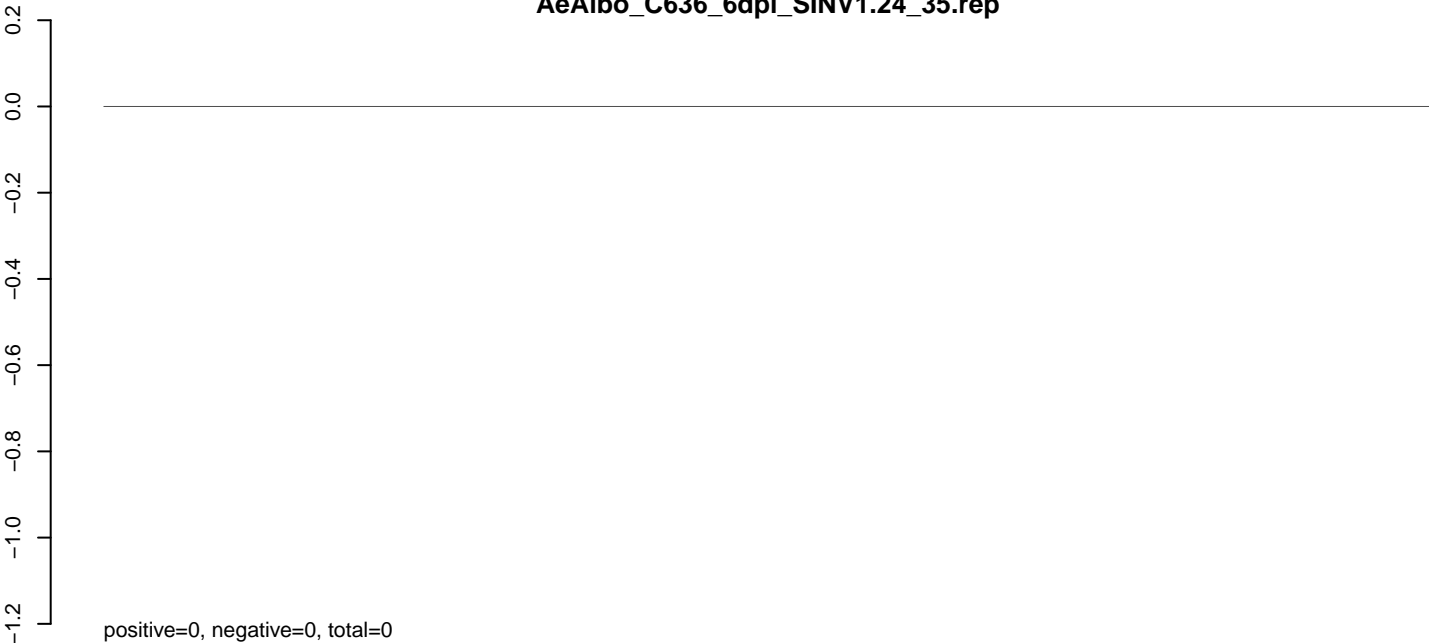
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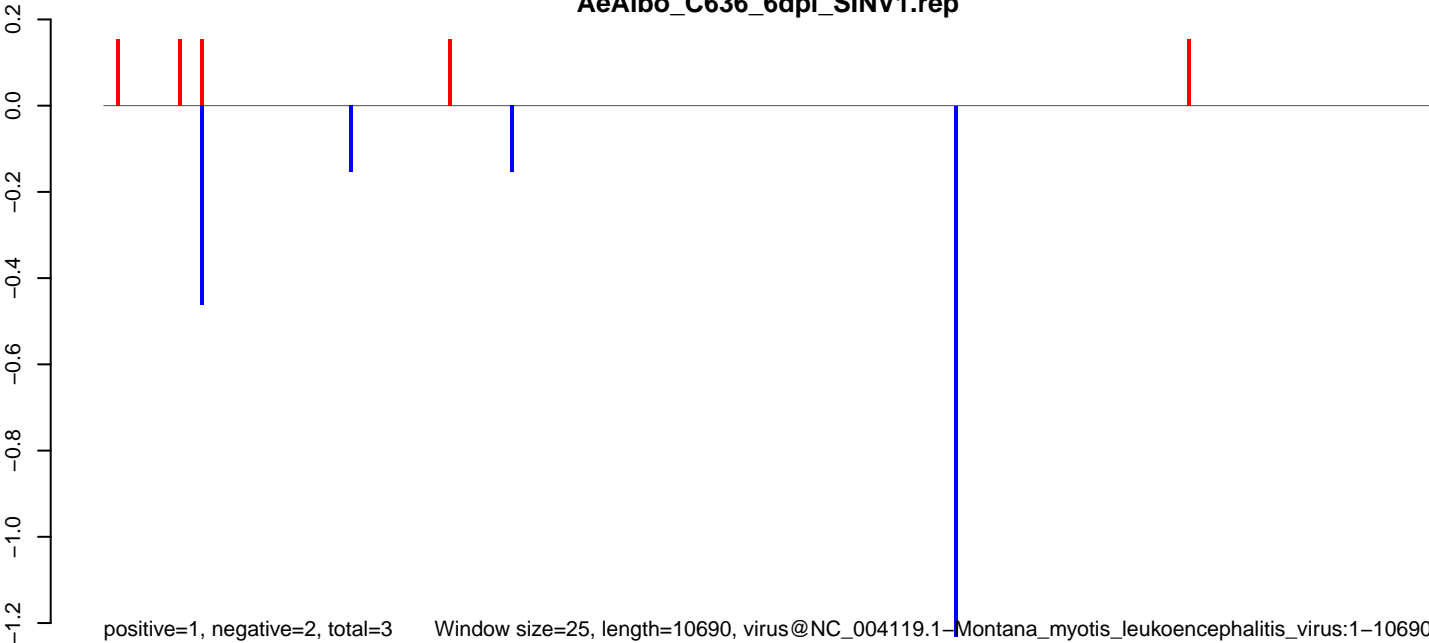
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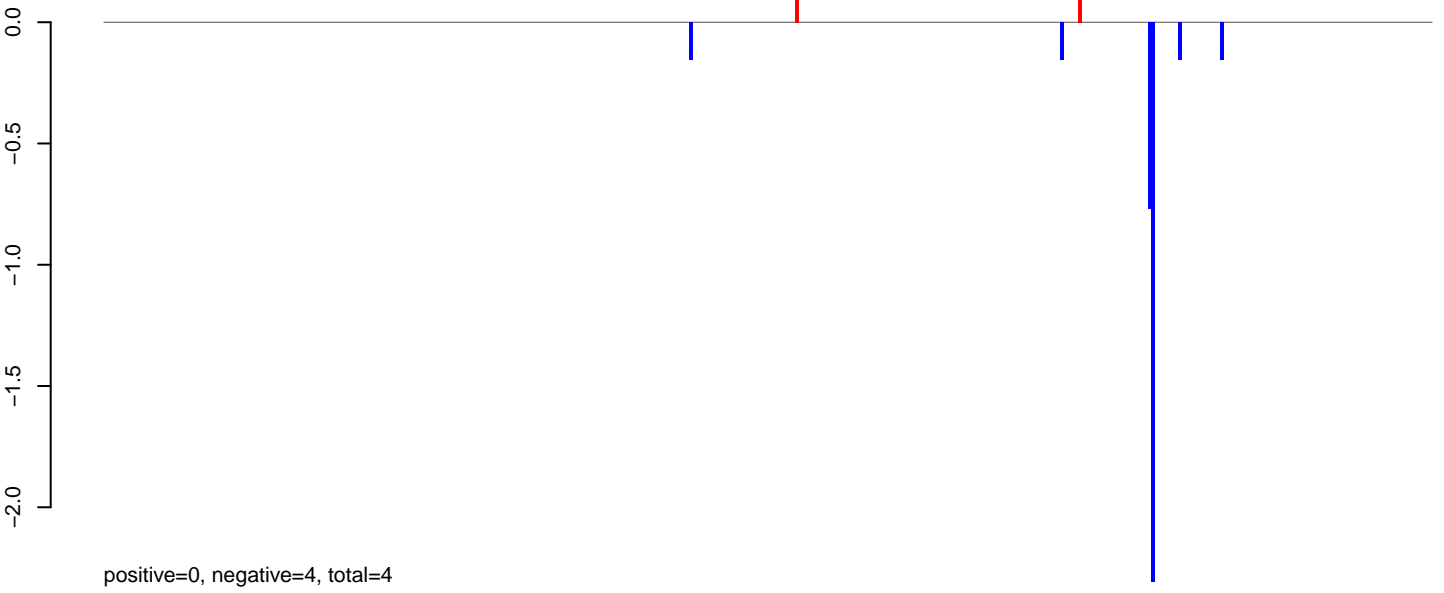
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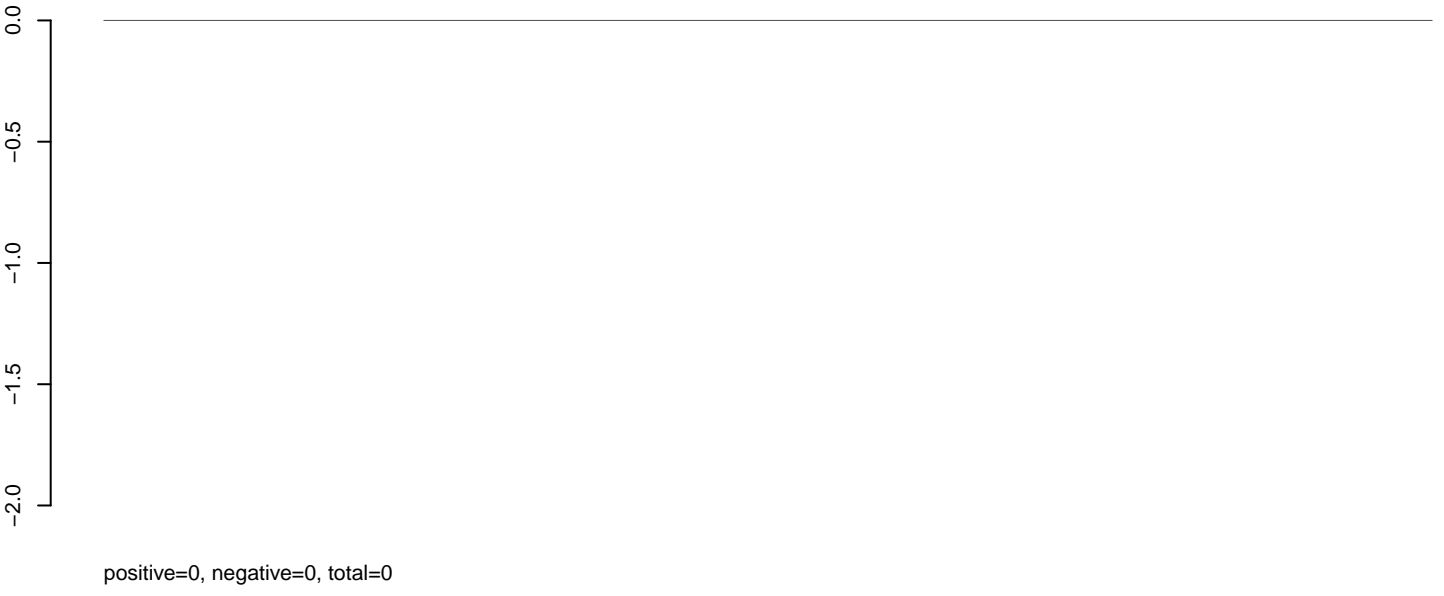
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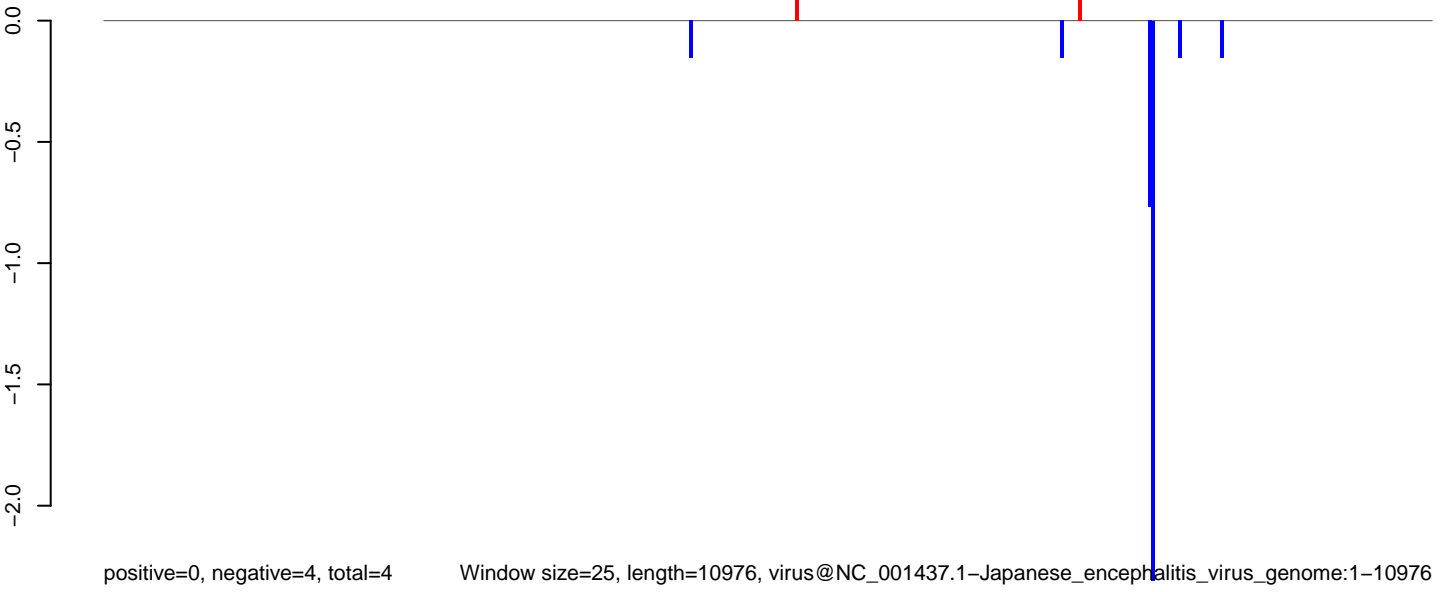
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AeAlbo_C636_6dpi_SINV1.24_35.rep



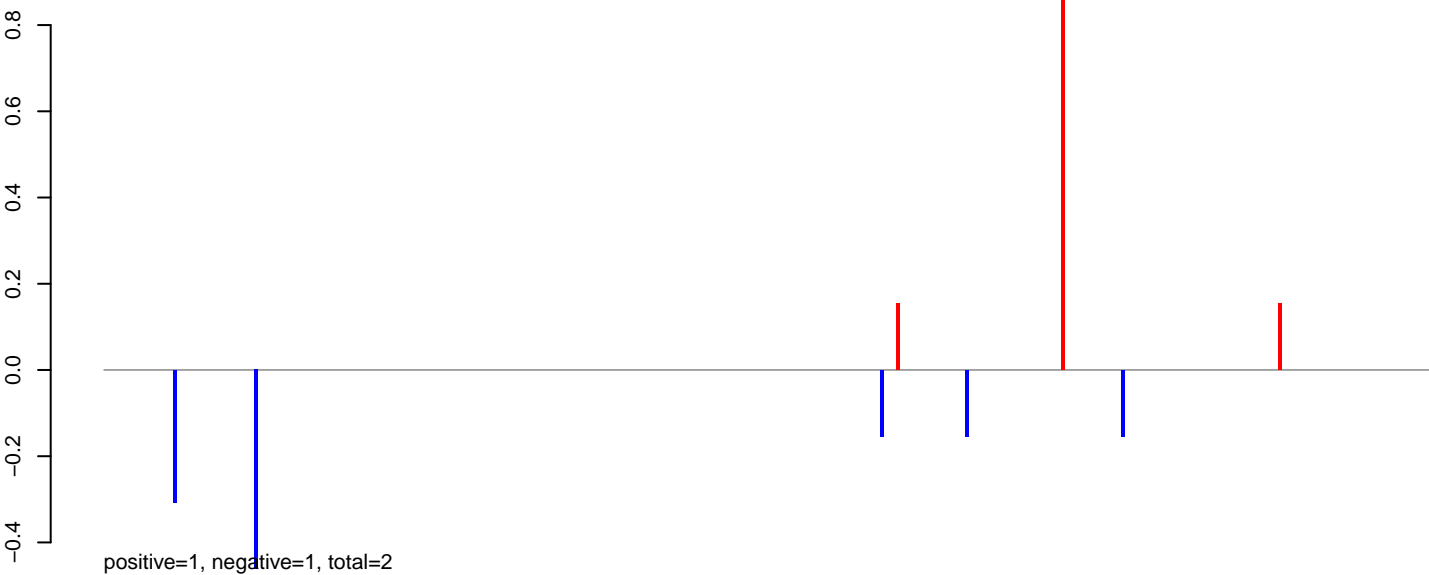
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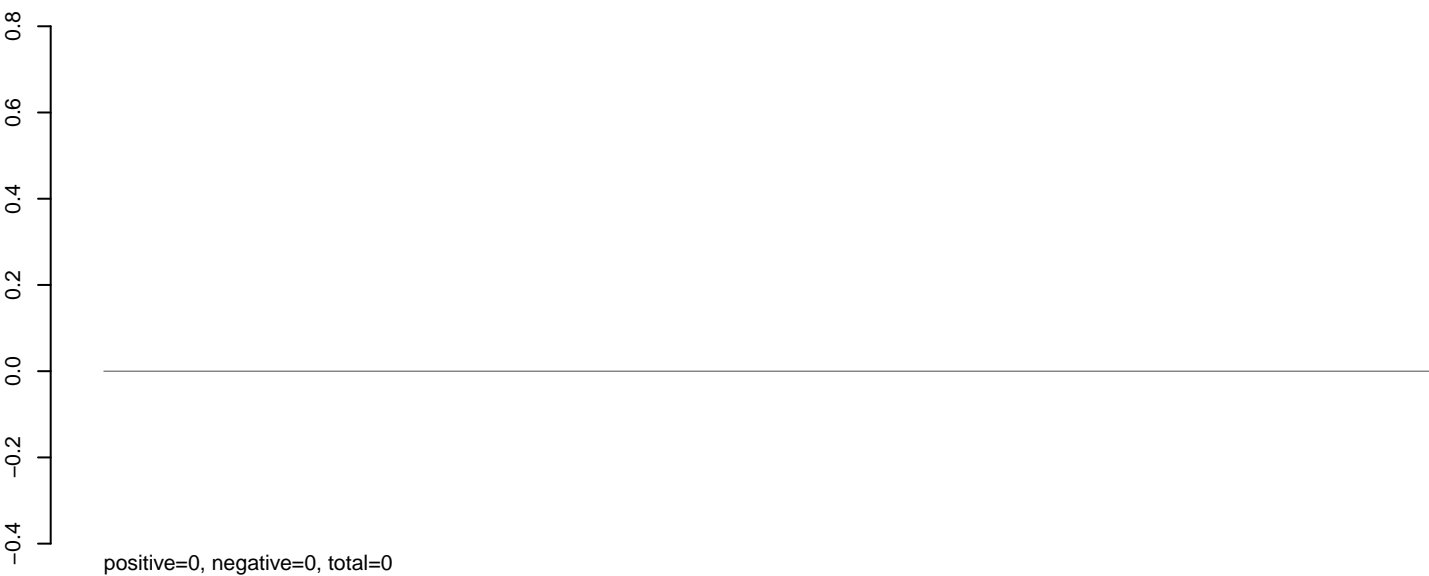
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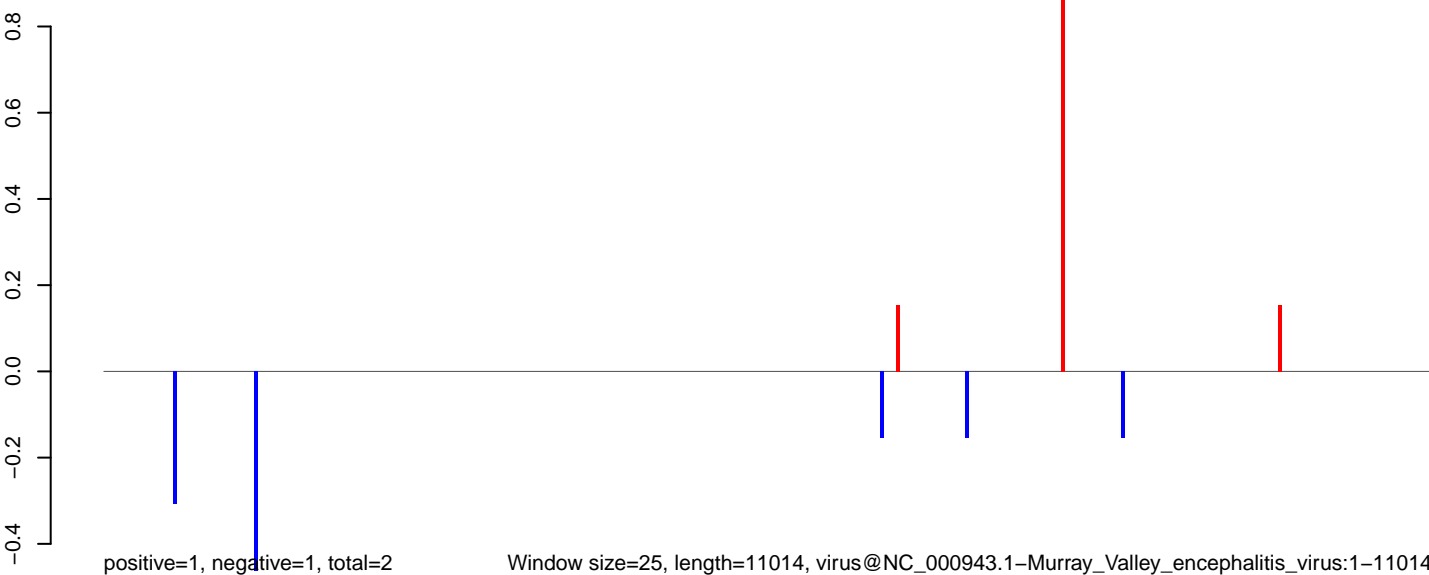
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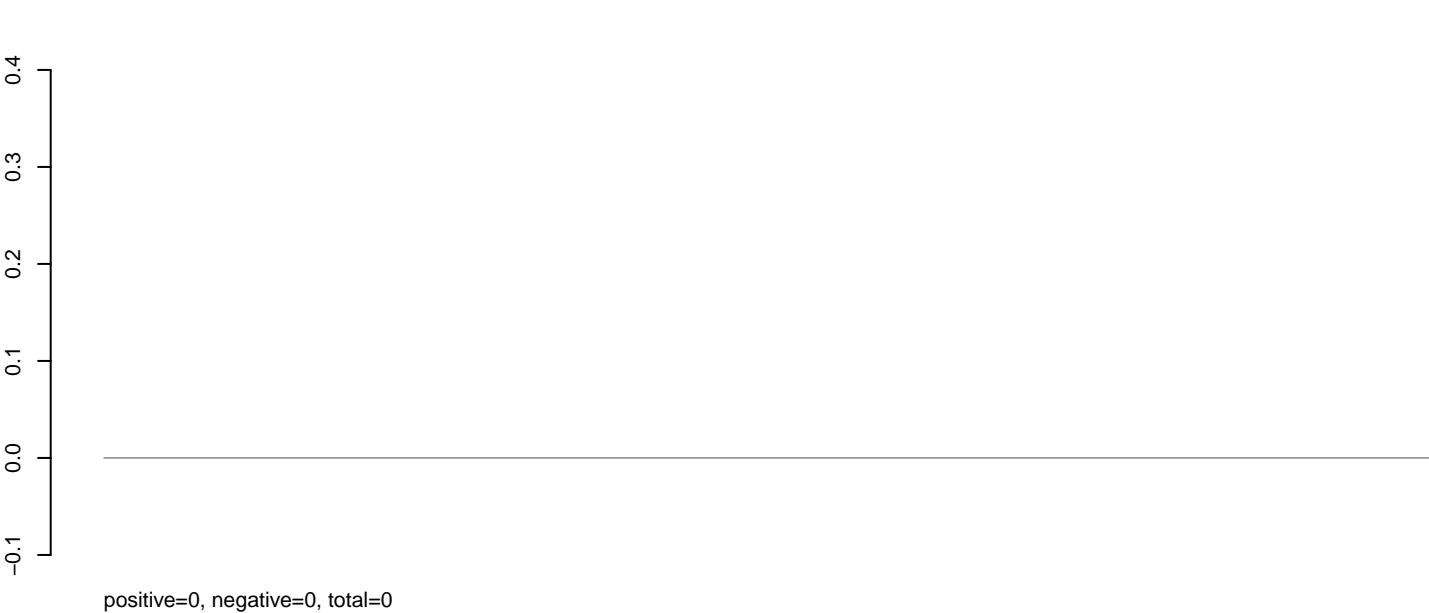
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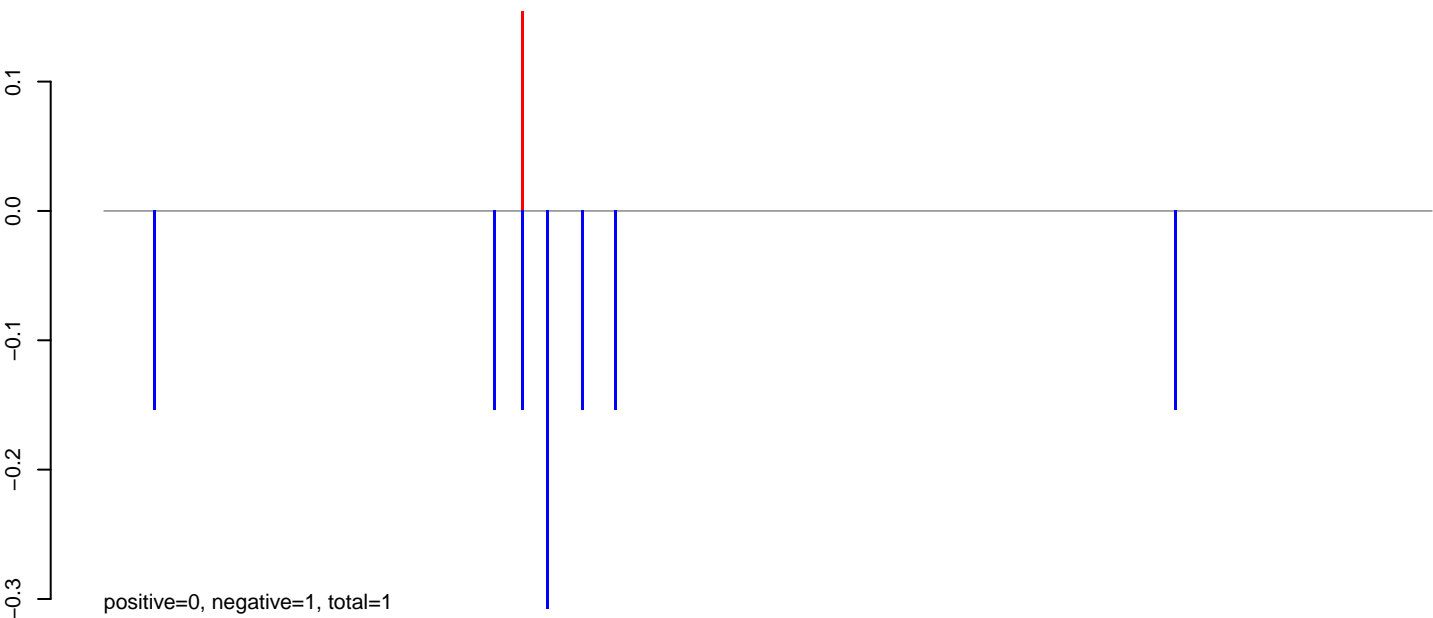
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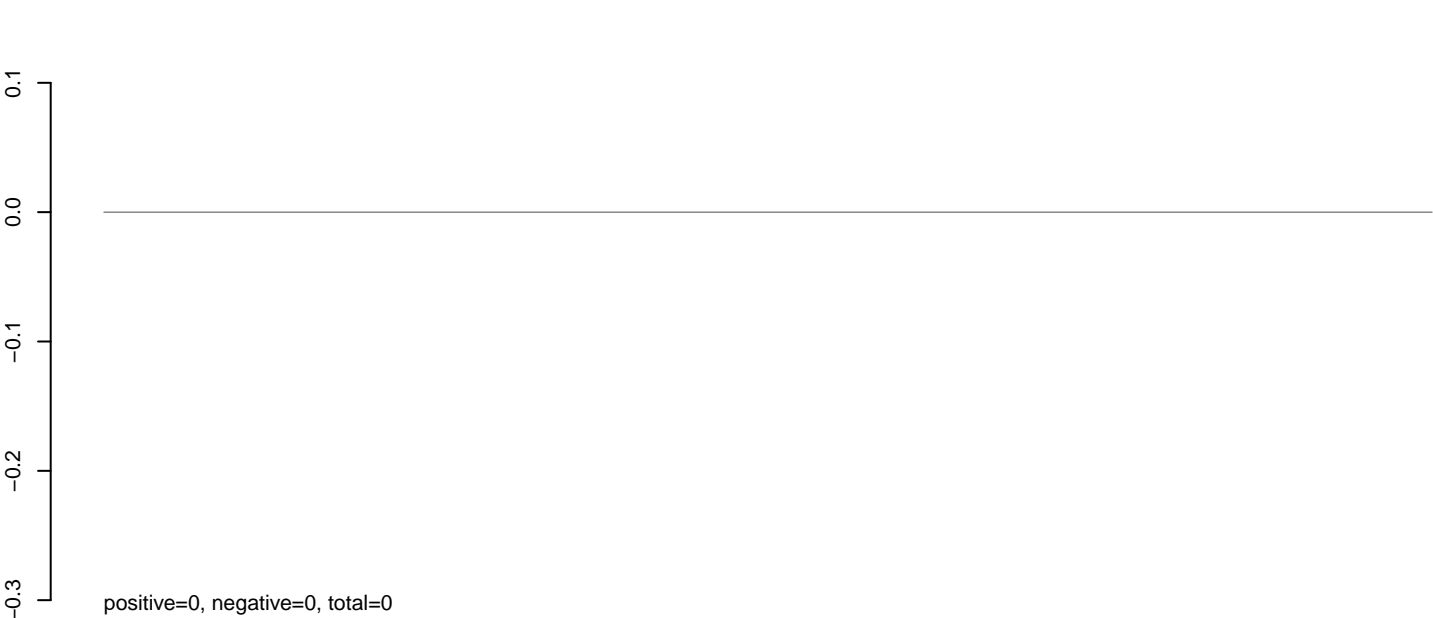
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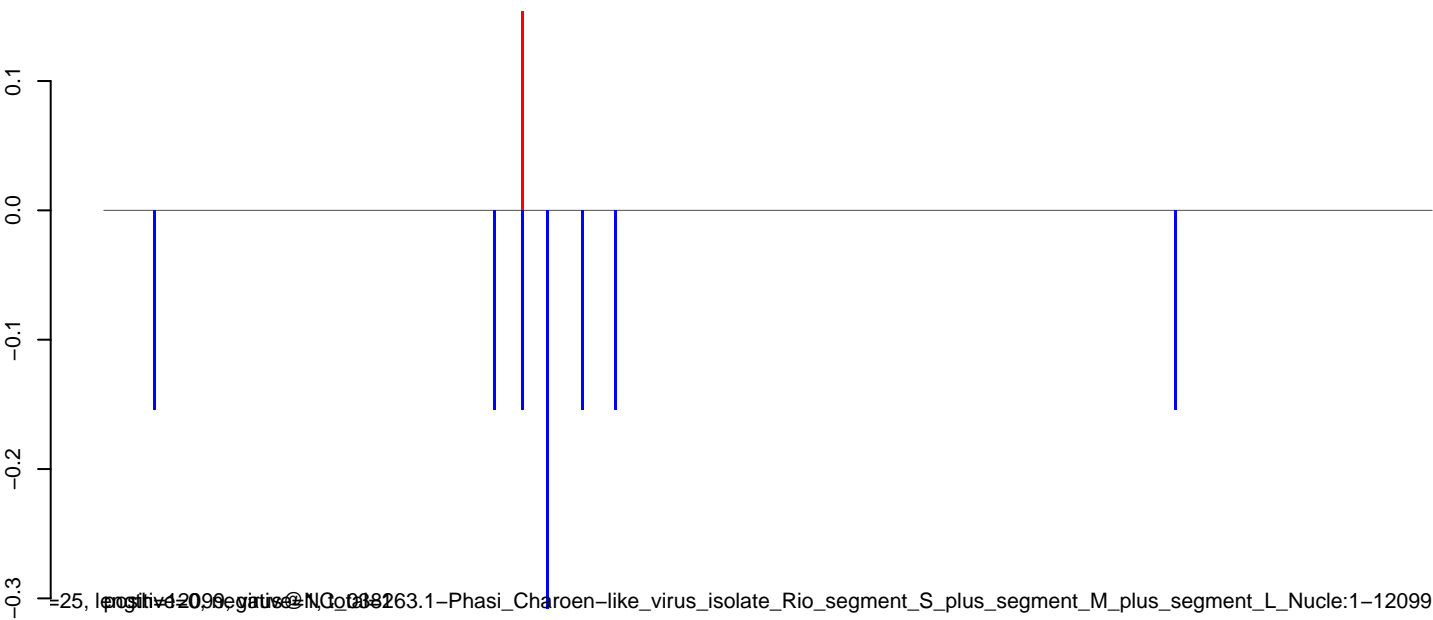
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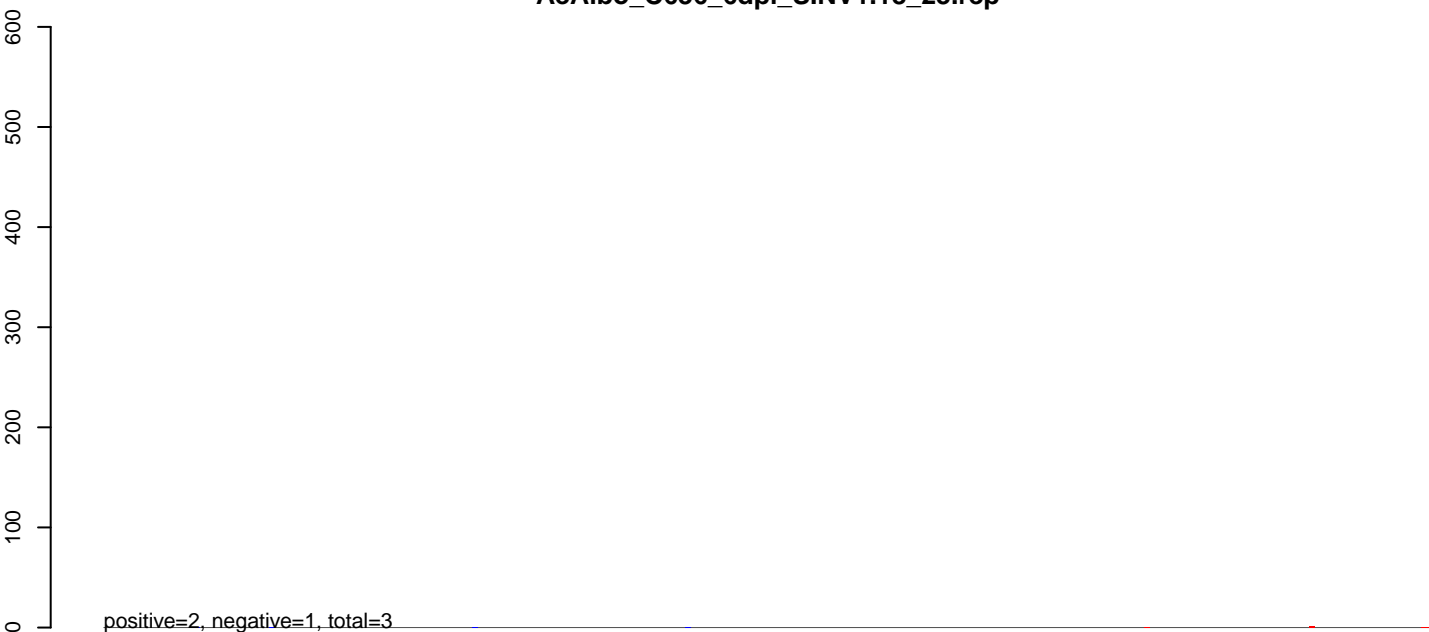
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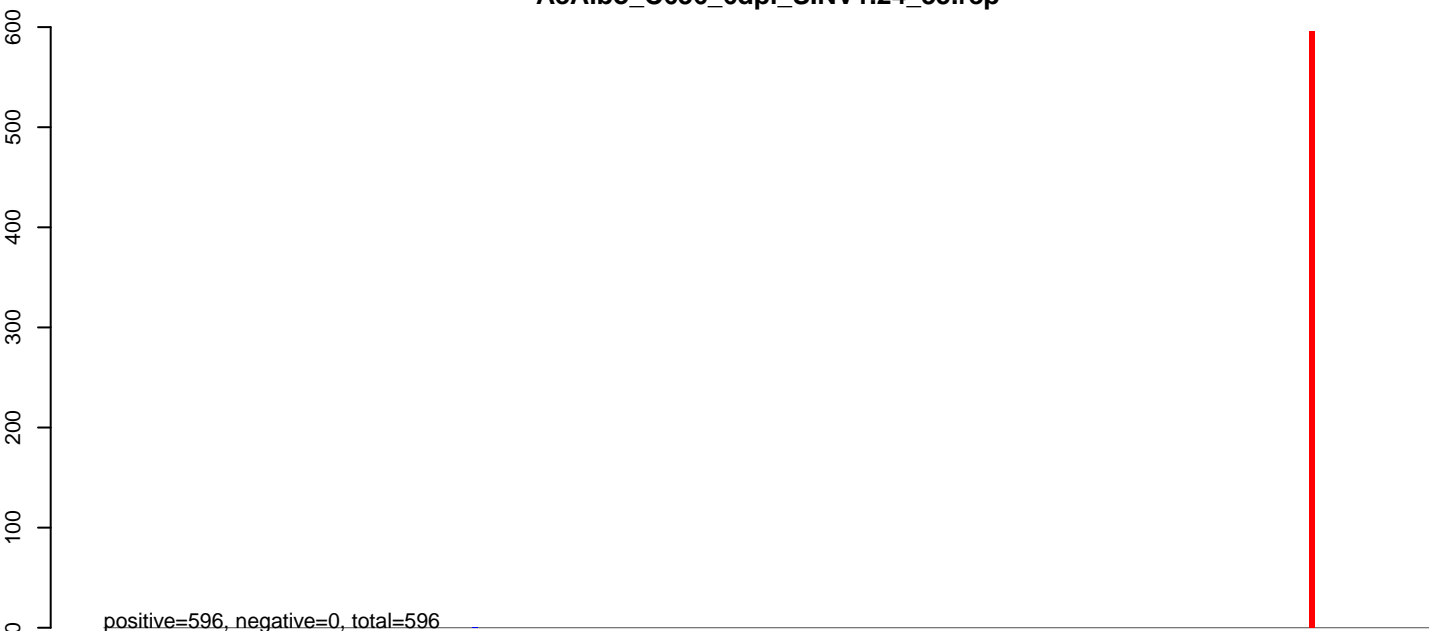
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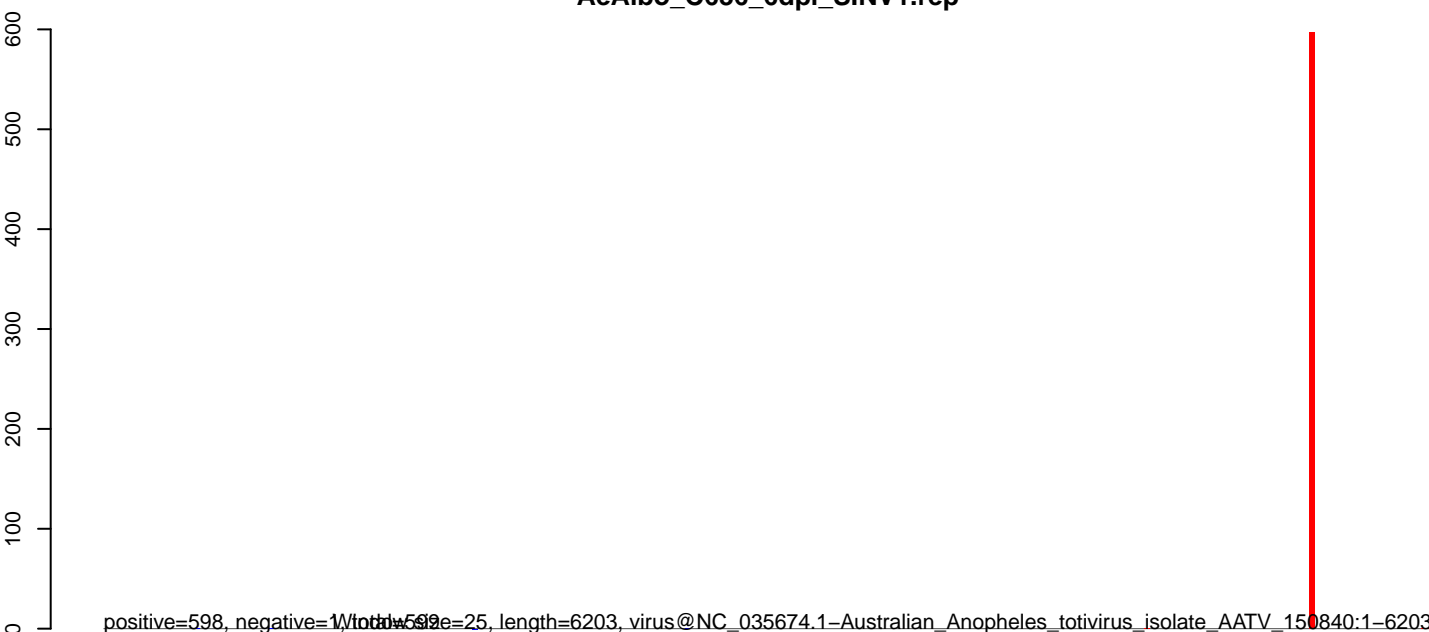
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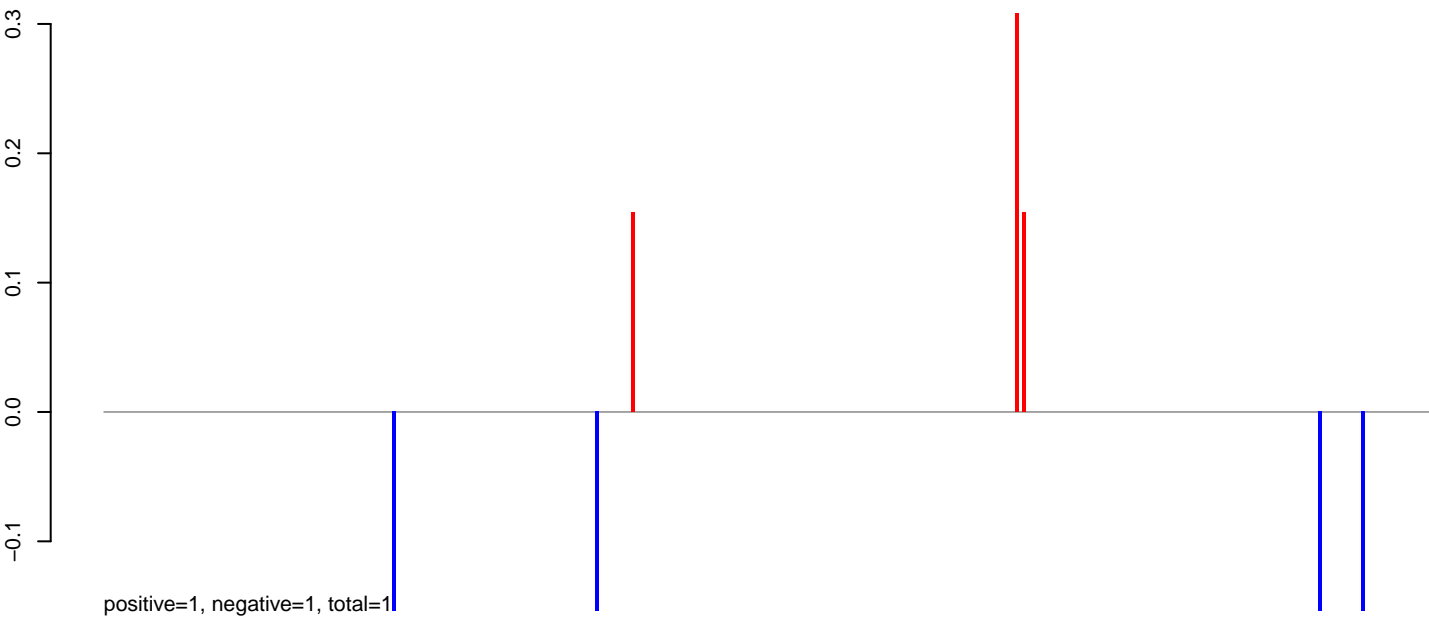
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0 1000 2000 3000 4000 5000 6000

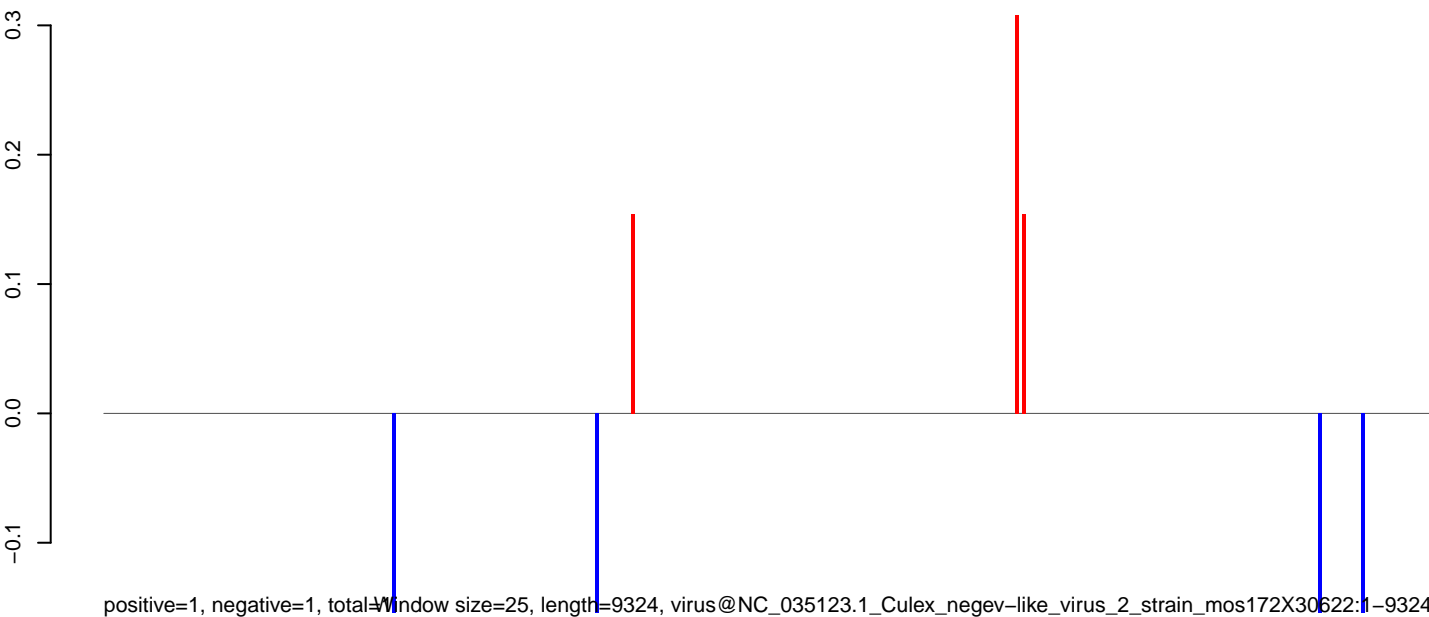
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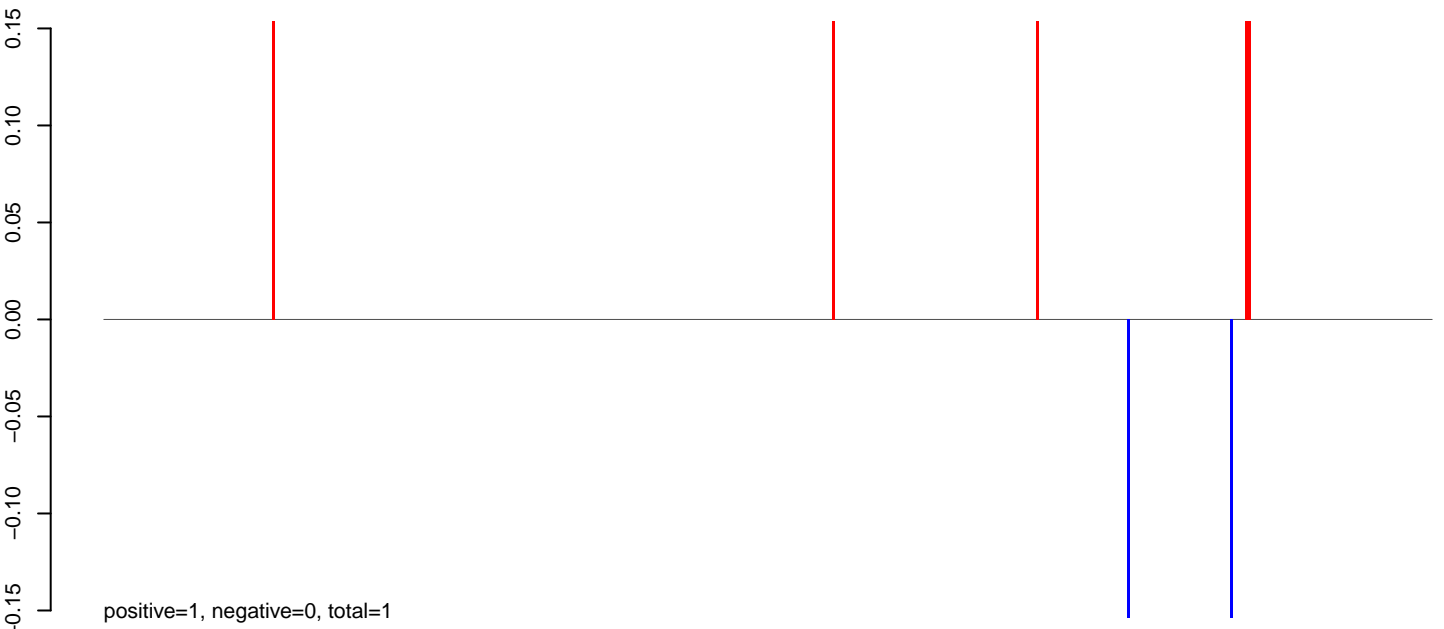


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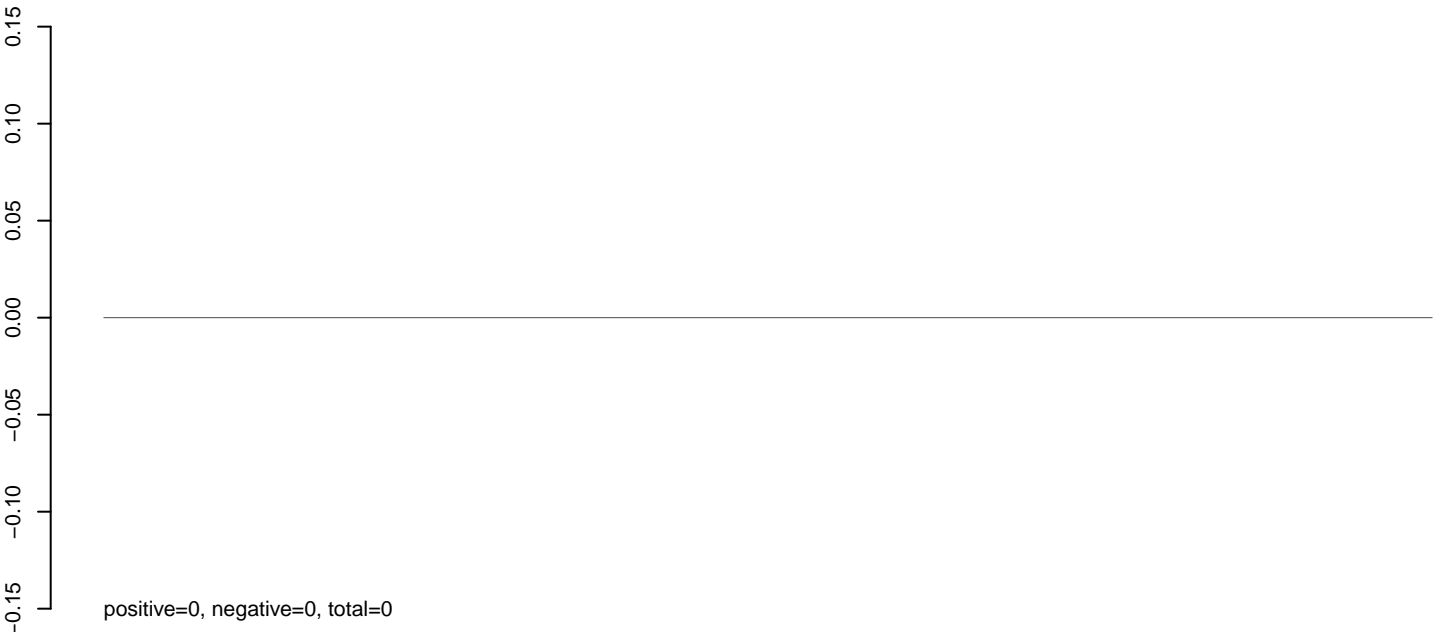


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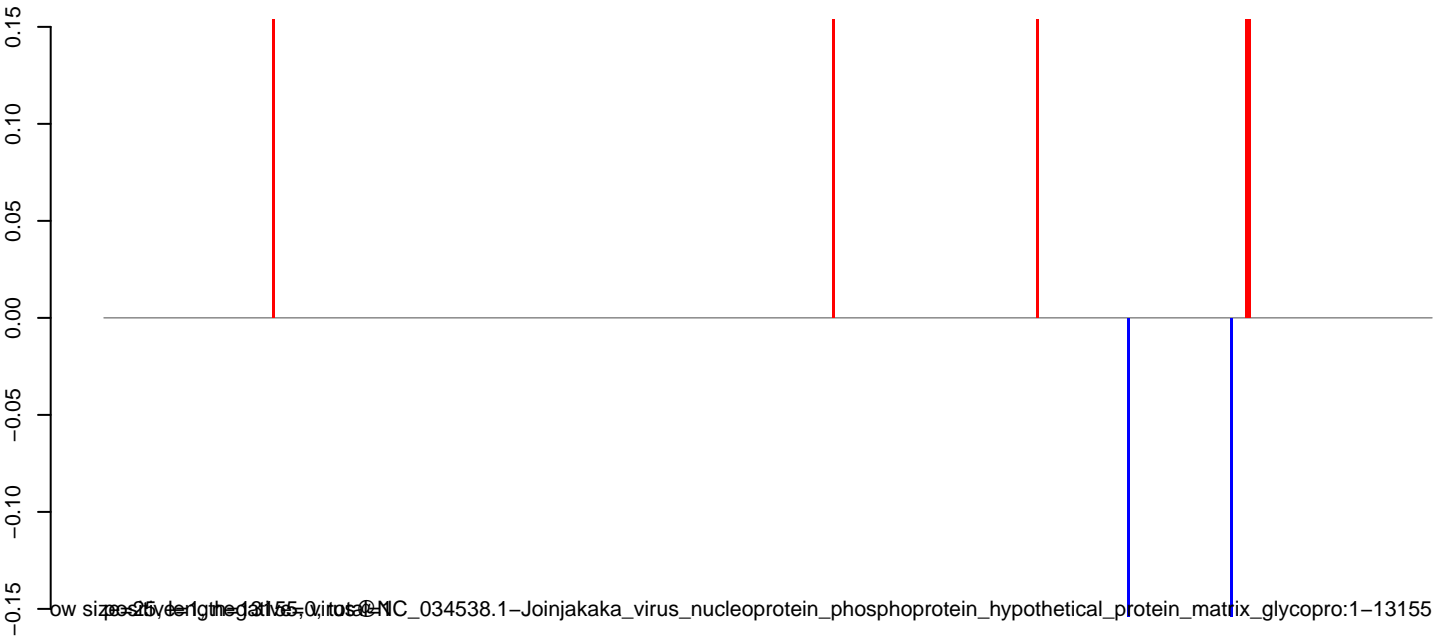
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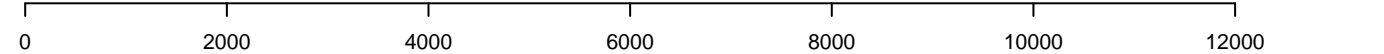
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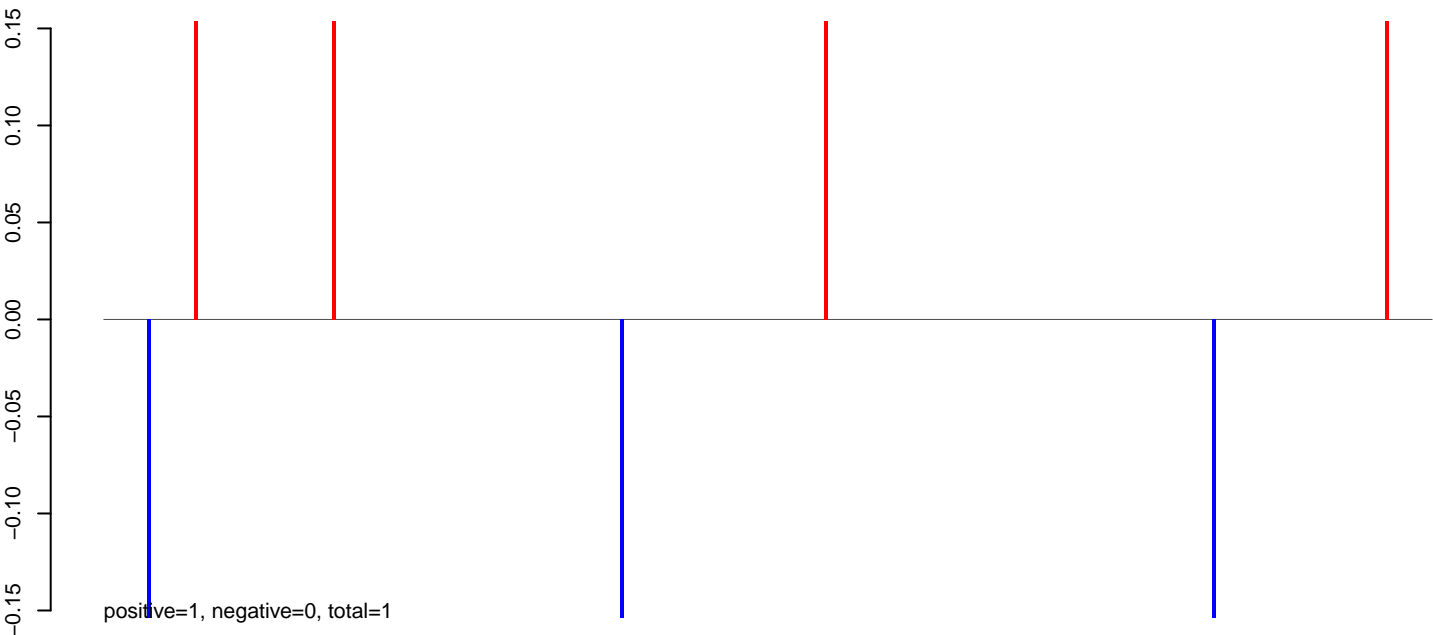
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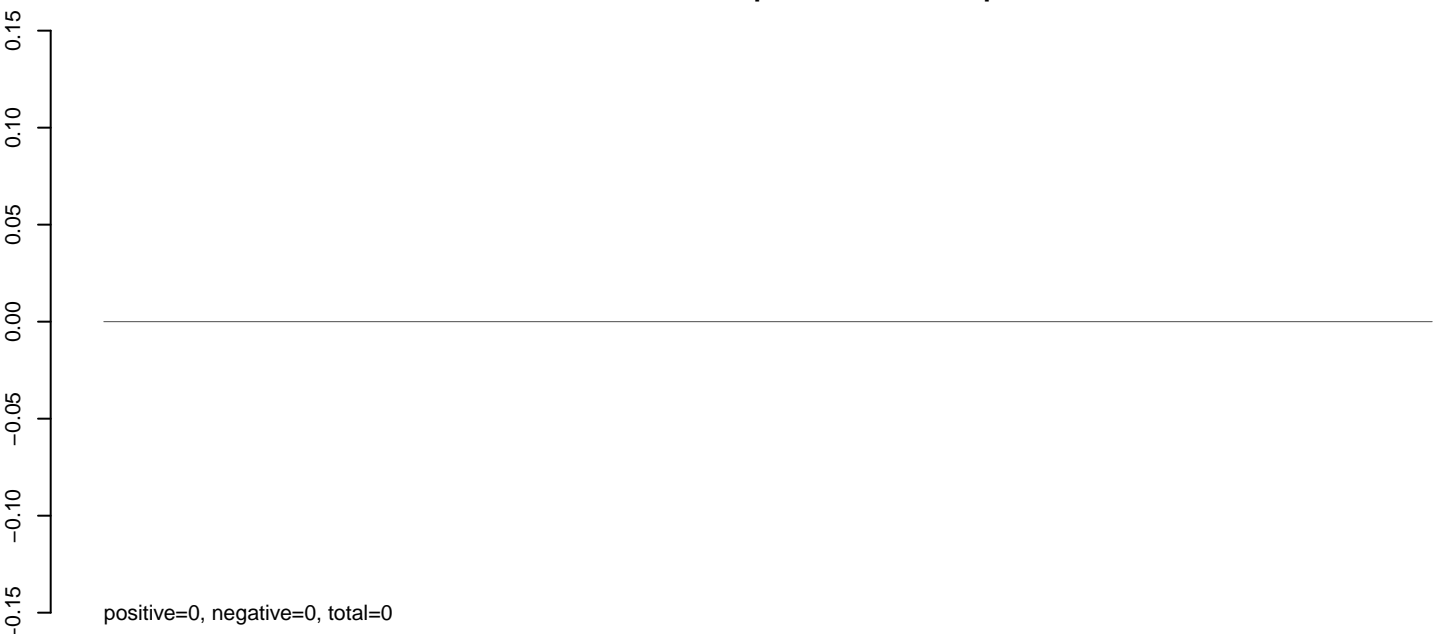
low size of the genome, it is a NC_034538.1-Joinjakaka_virus_nucleoprotein_phosphoprotein_hypothetical_protein_matrix_glycopro:1-13155



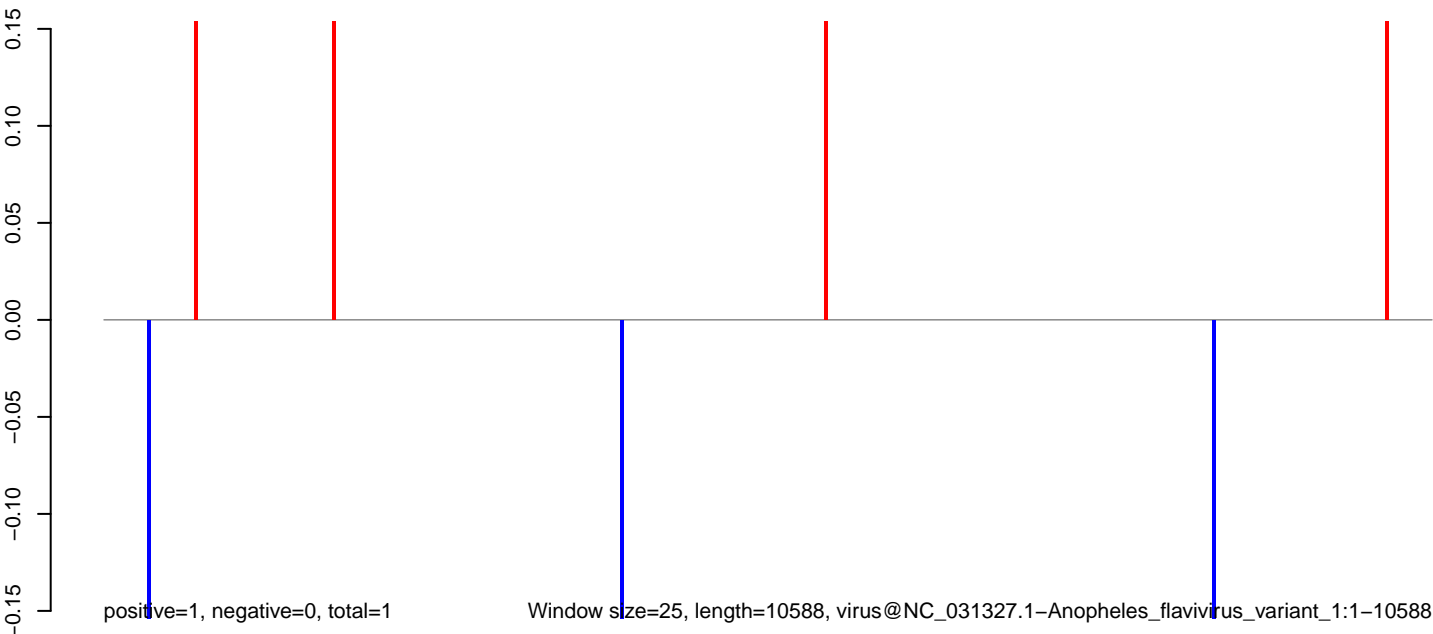
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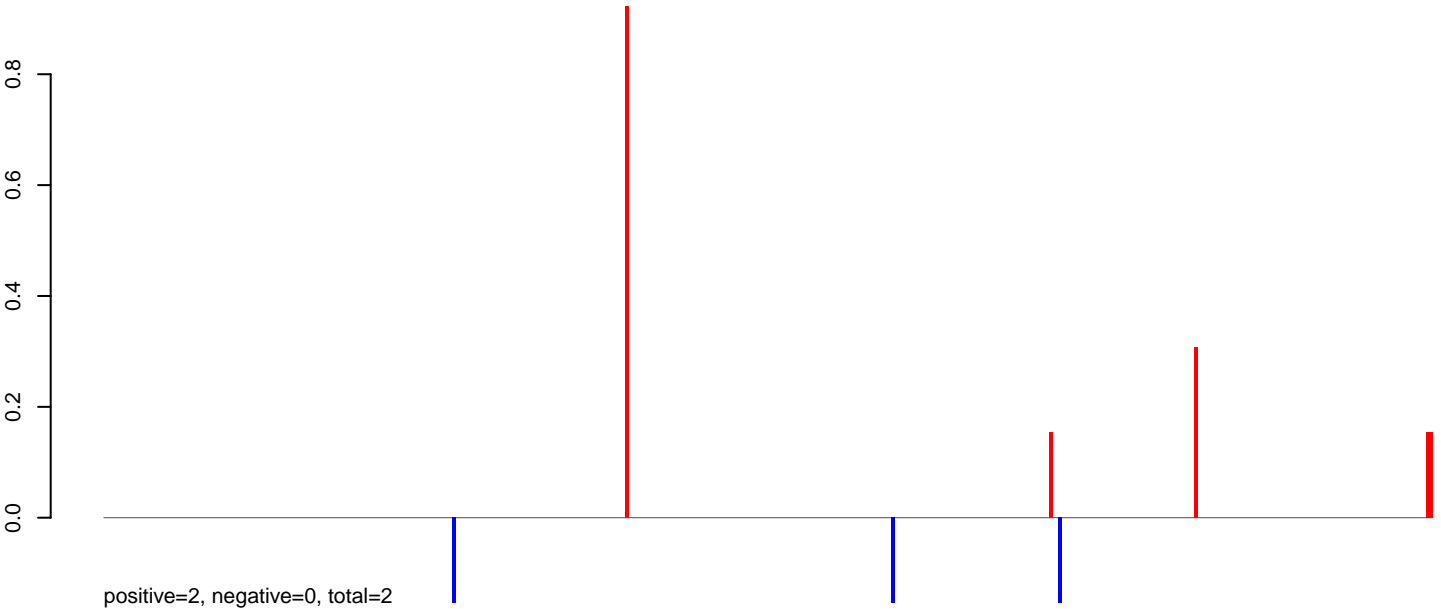


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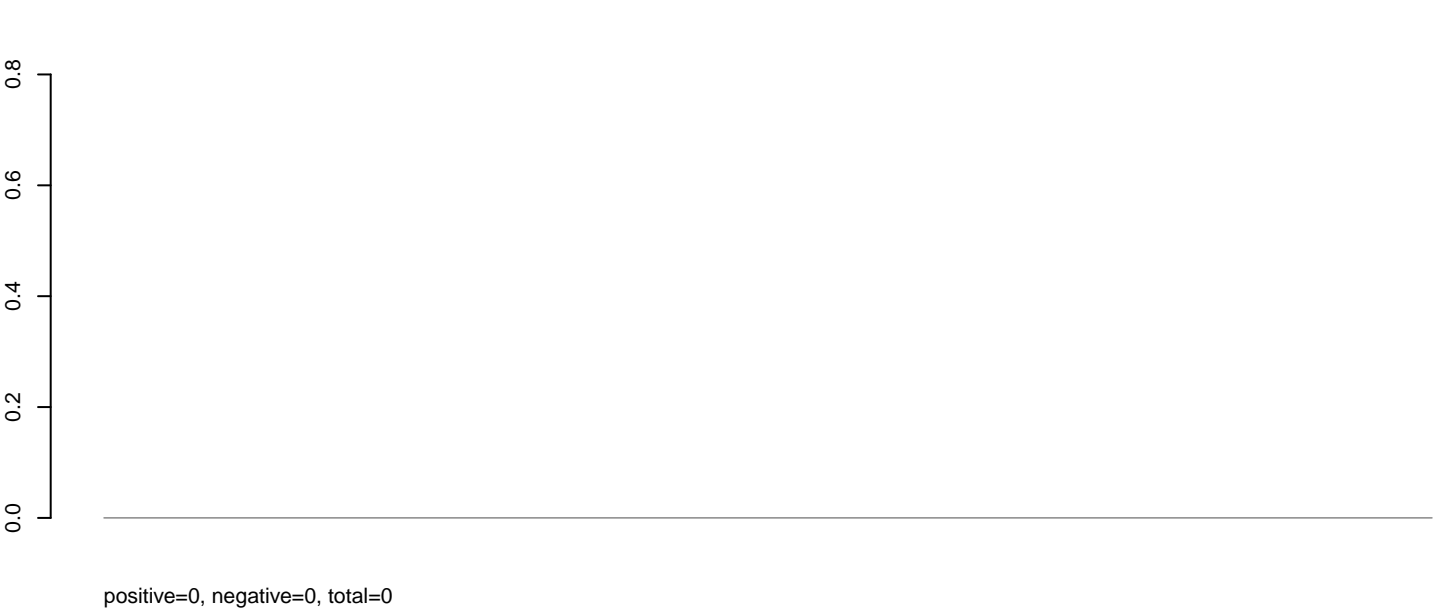


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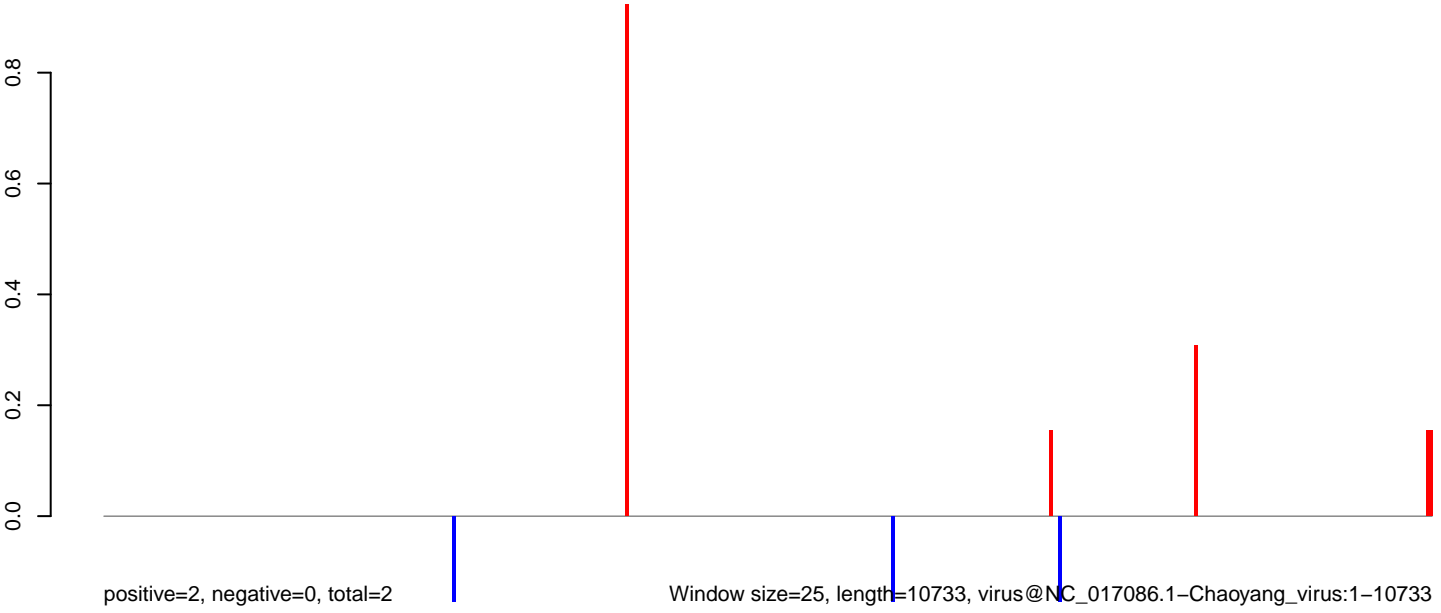
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AeAlbo_C636_6dpi_SINV1.24_35.rep

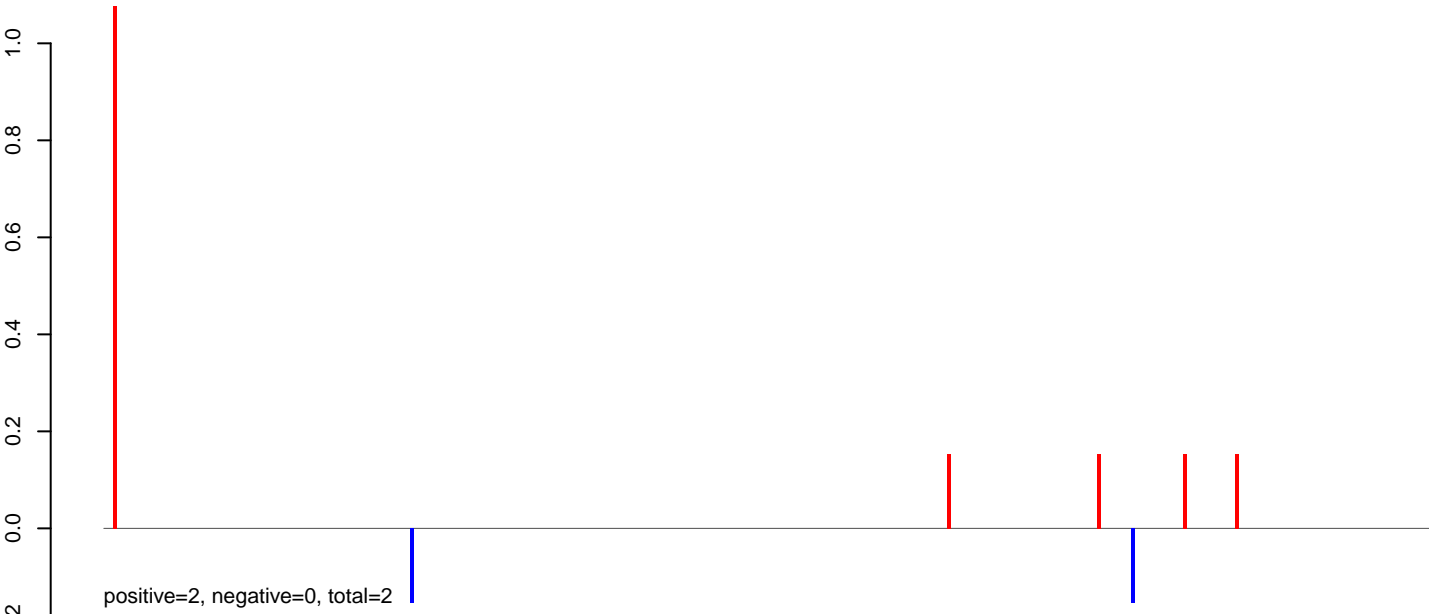


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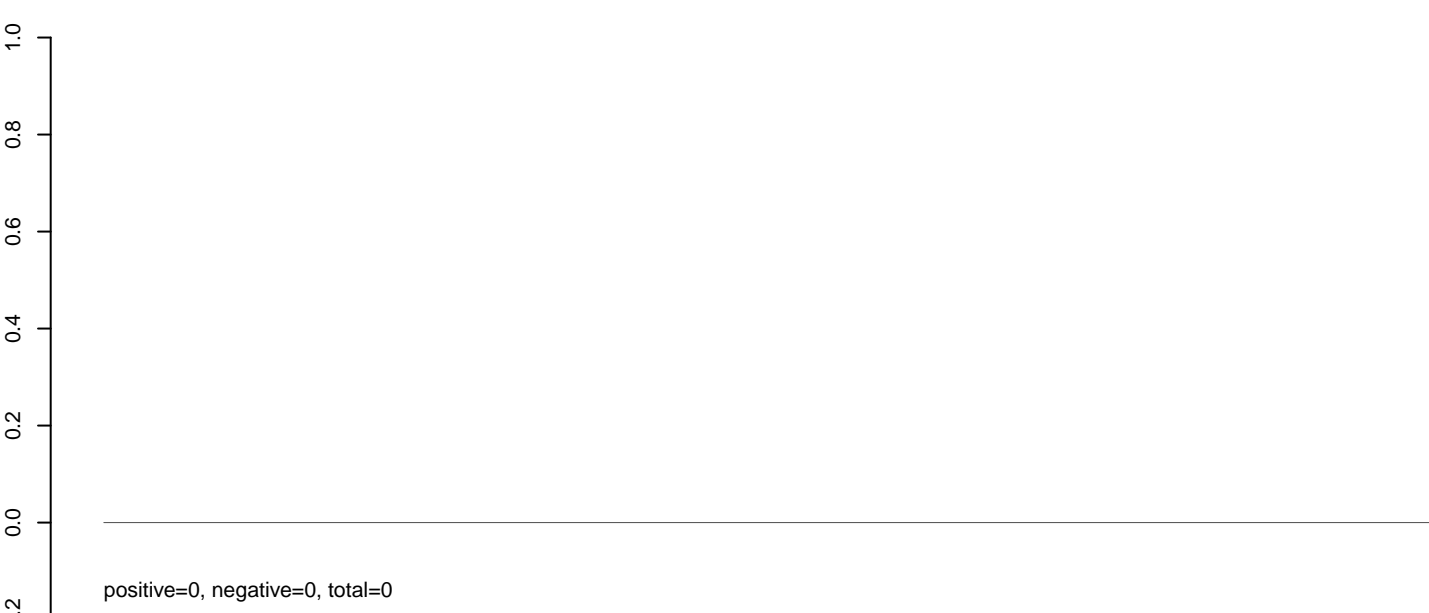


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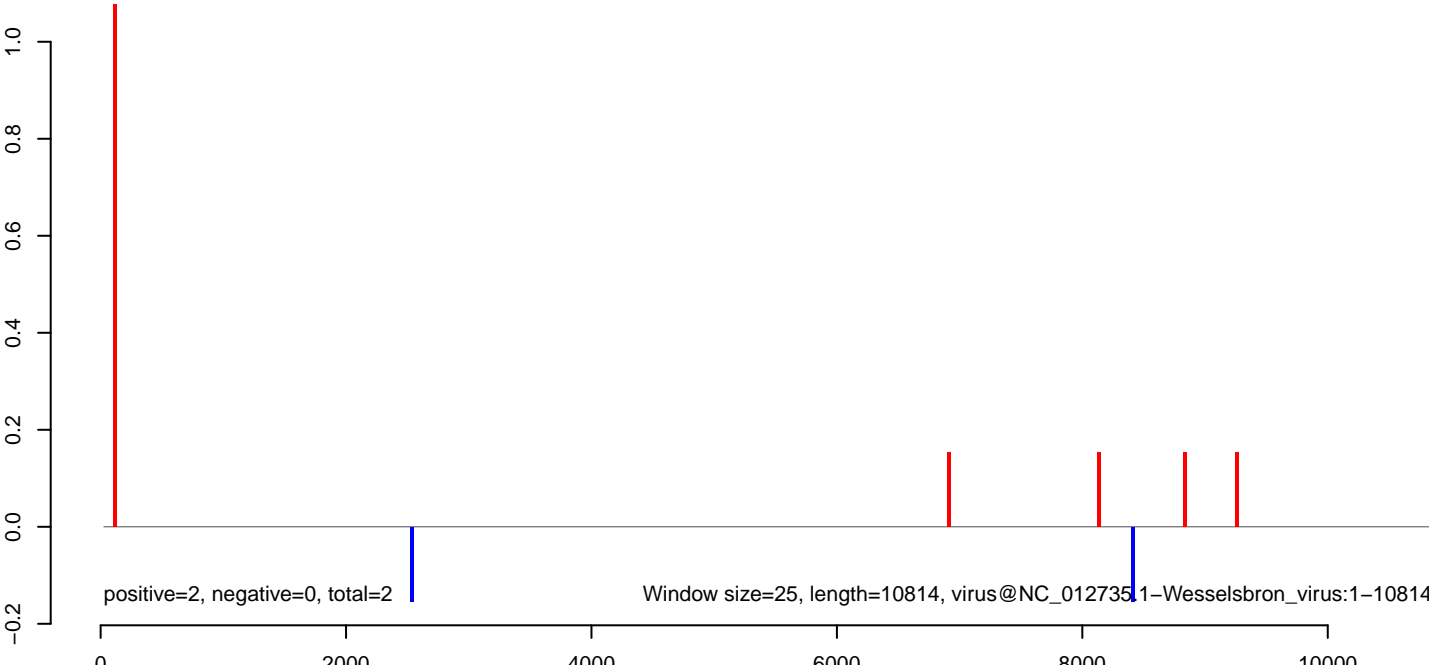
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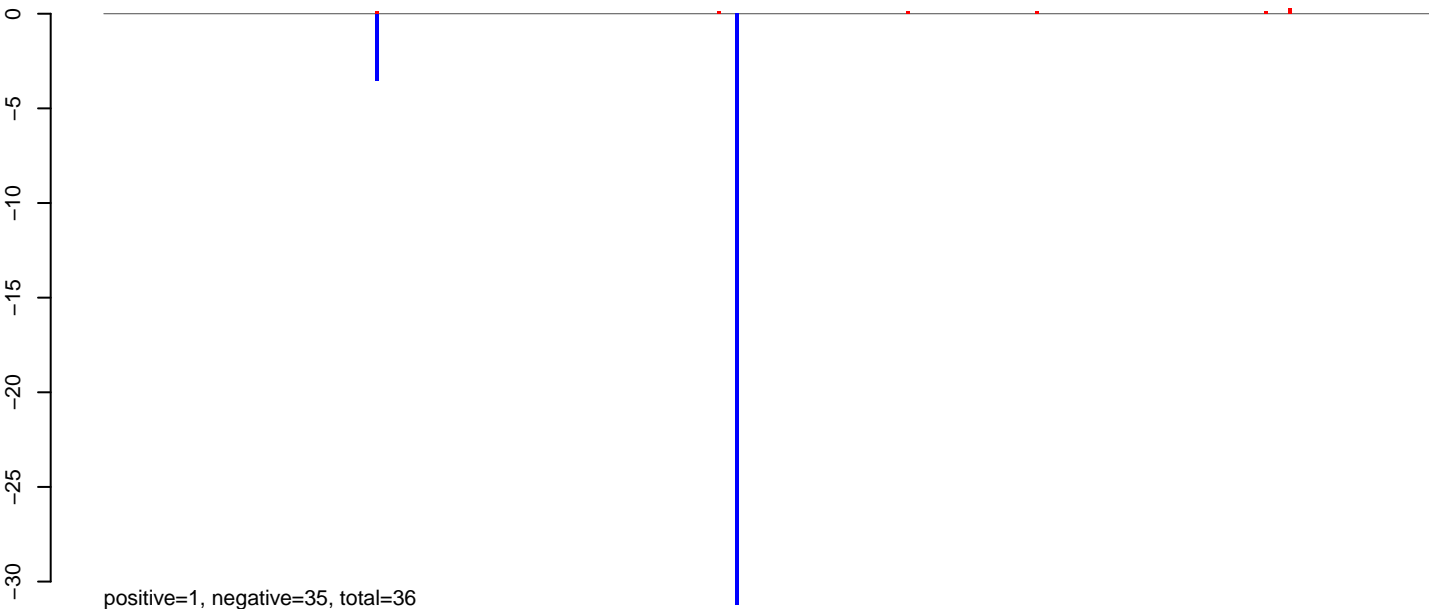
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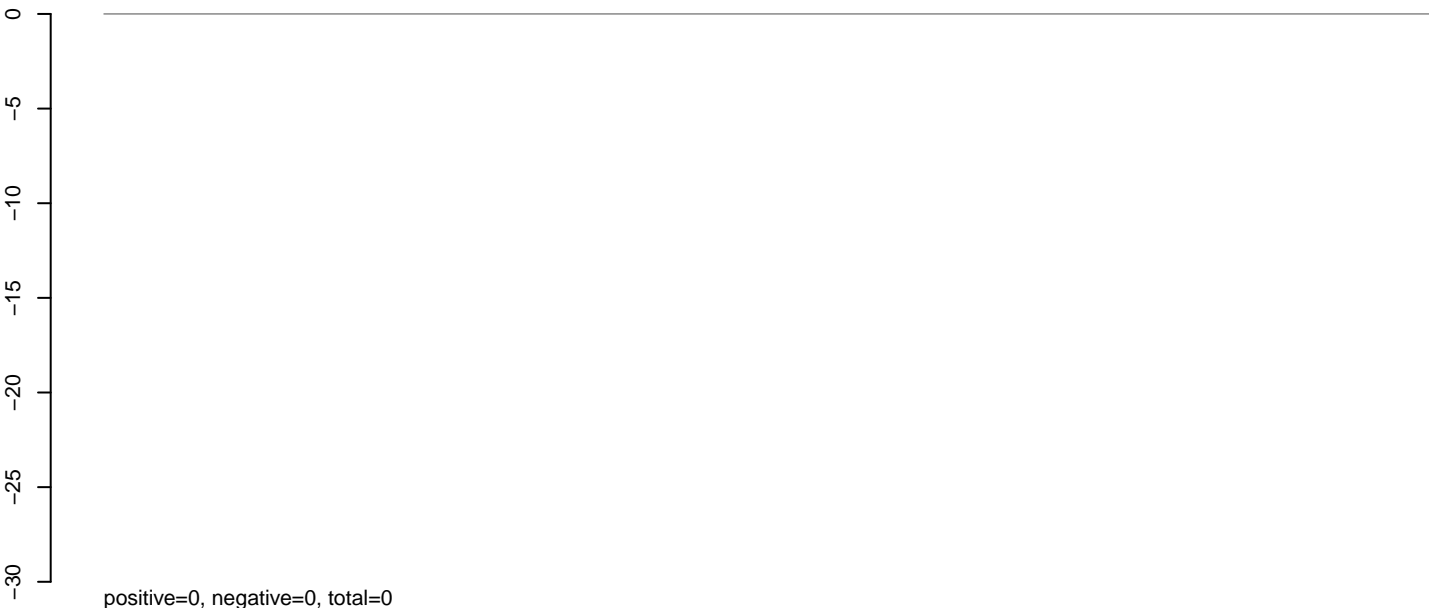
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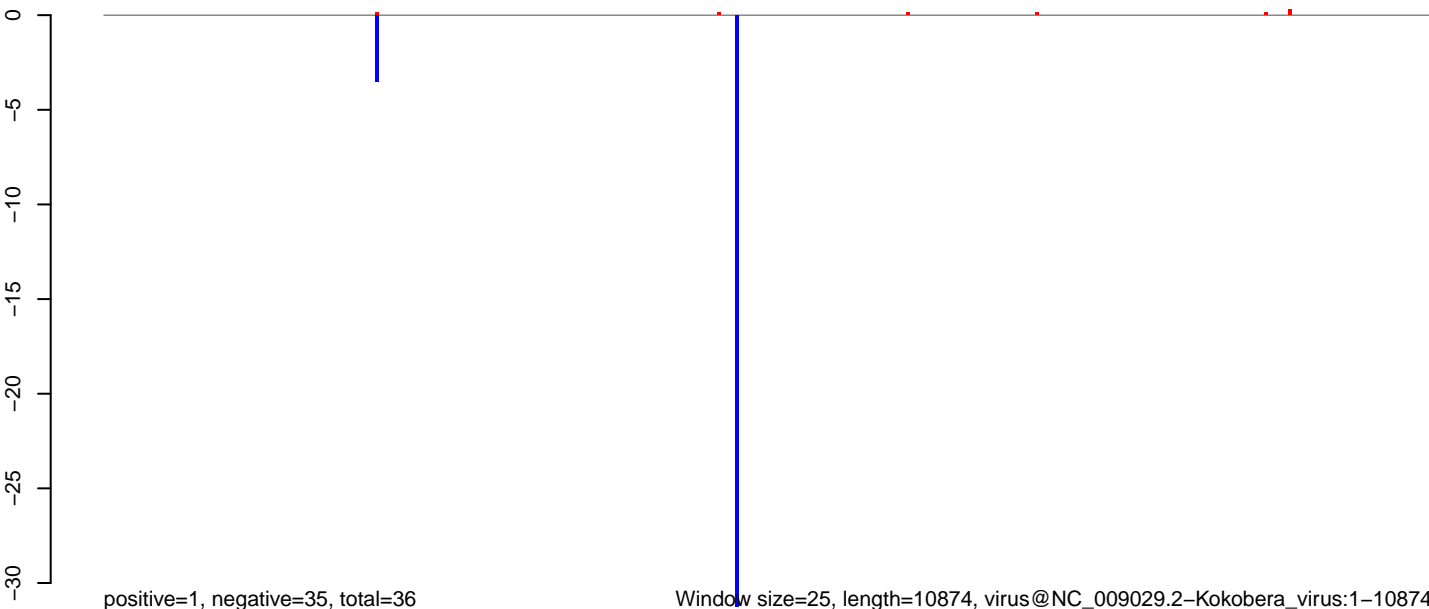
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AeAlbo_C636_6dpi_SINV1.24_35.rep



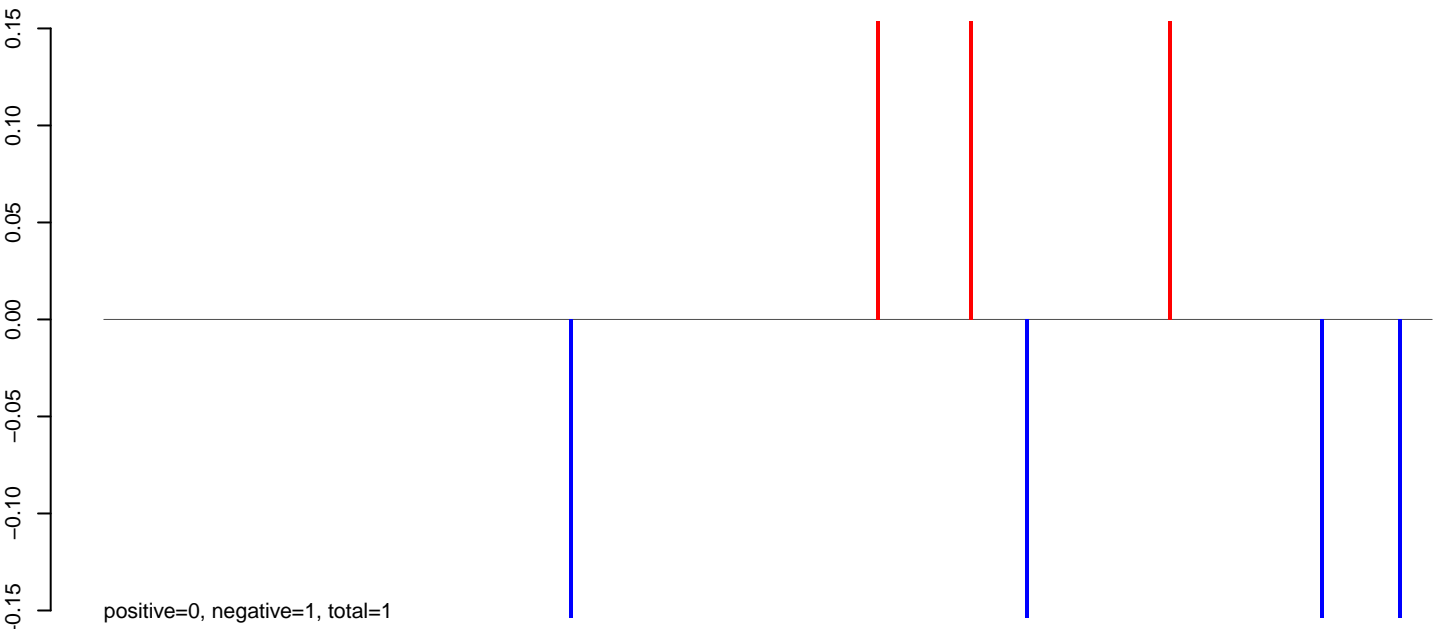
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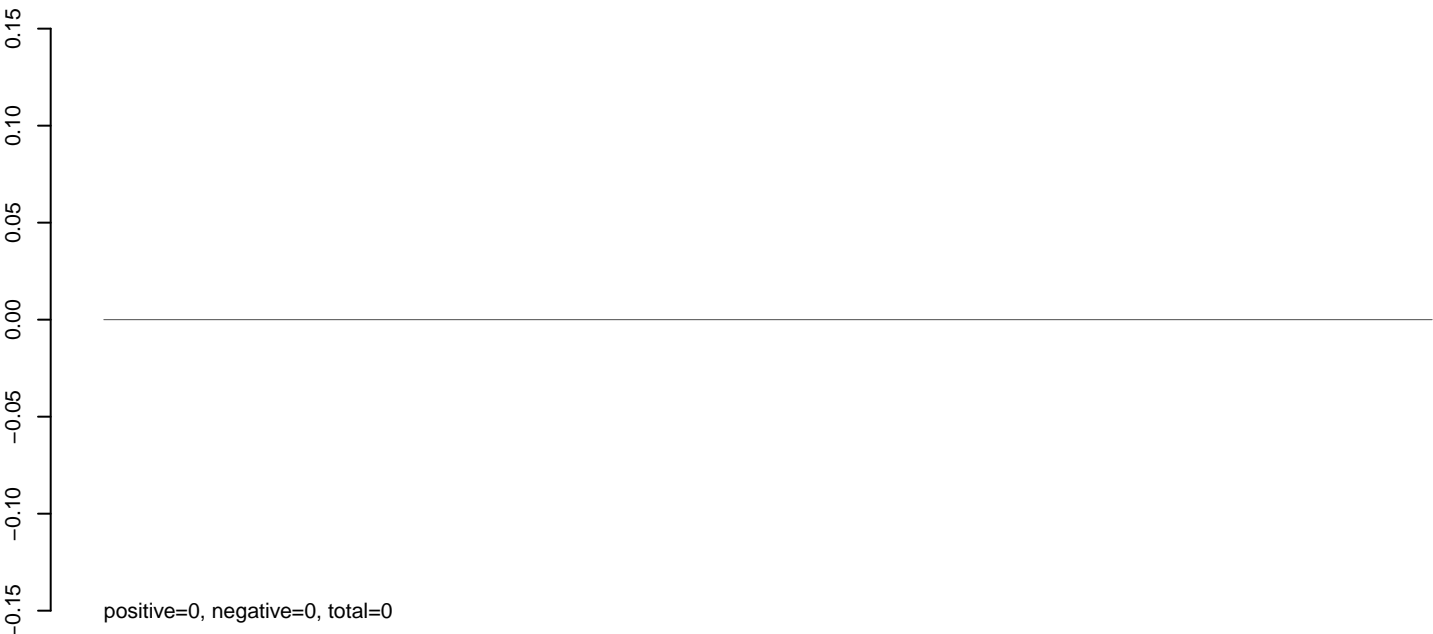
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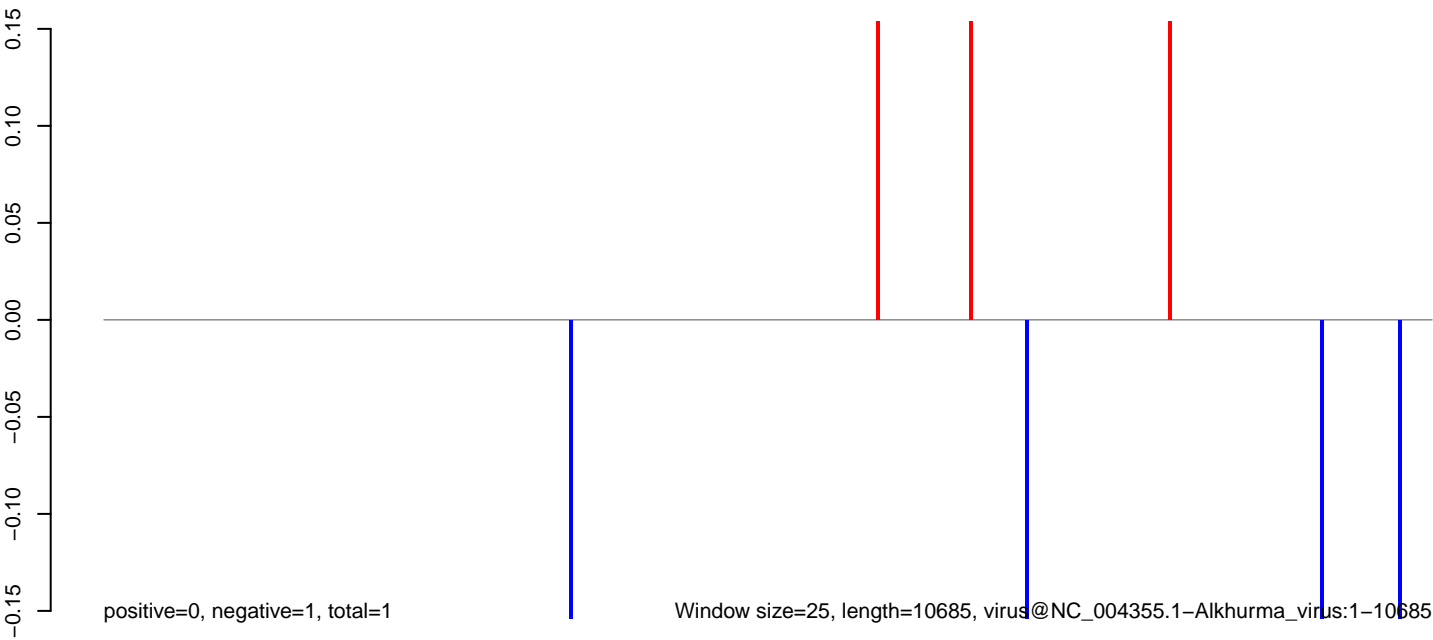
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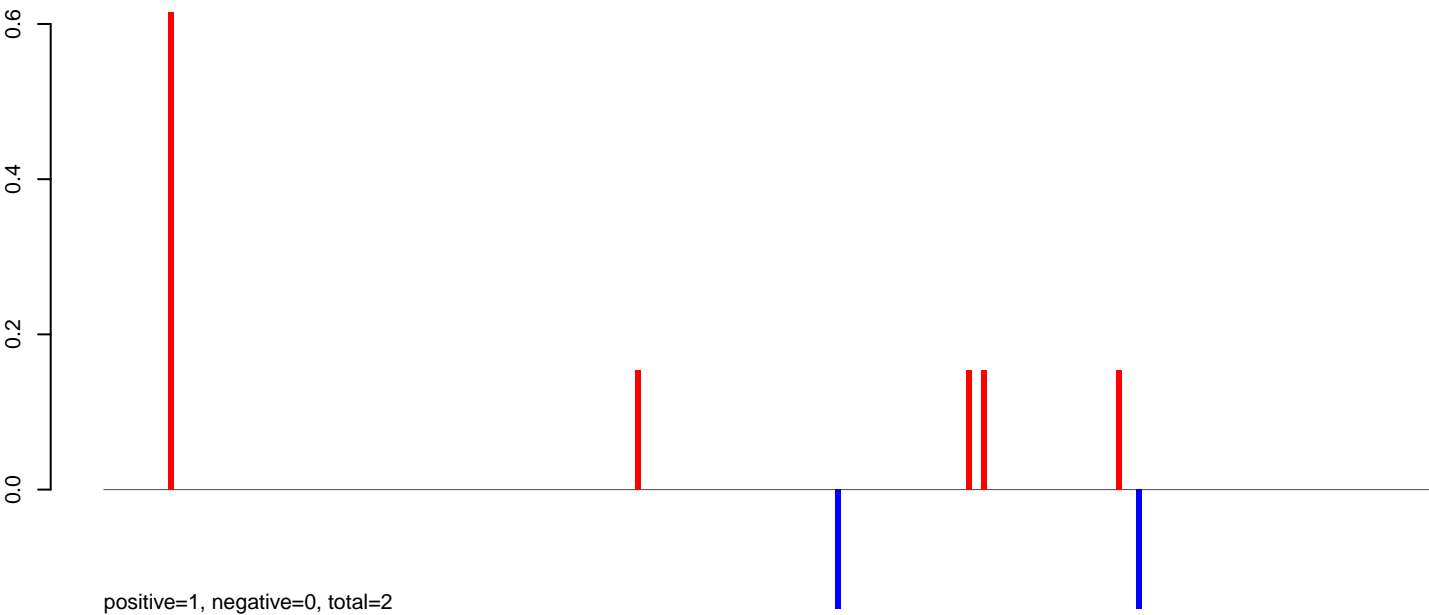
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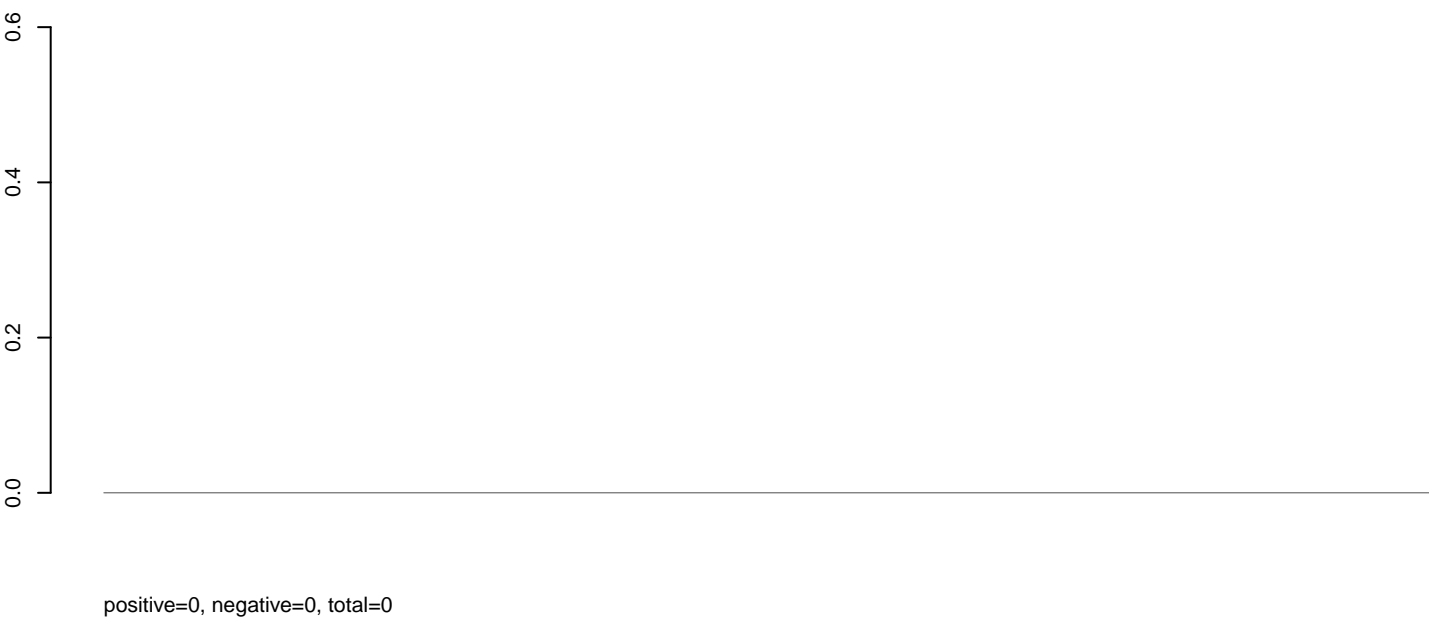
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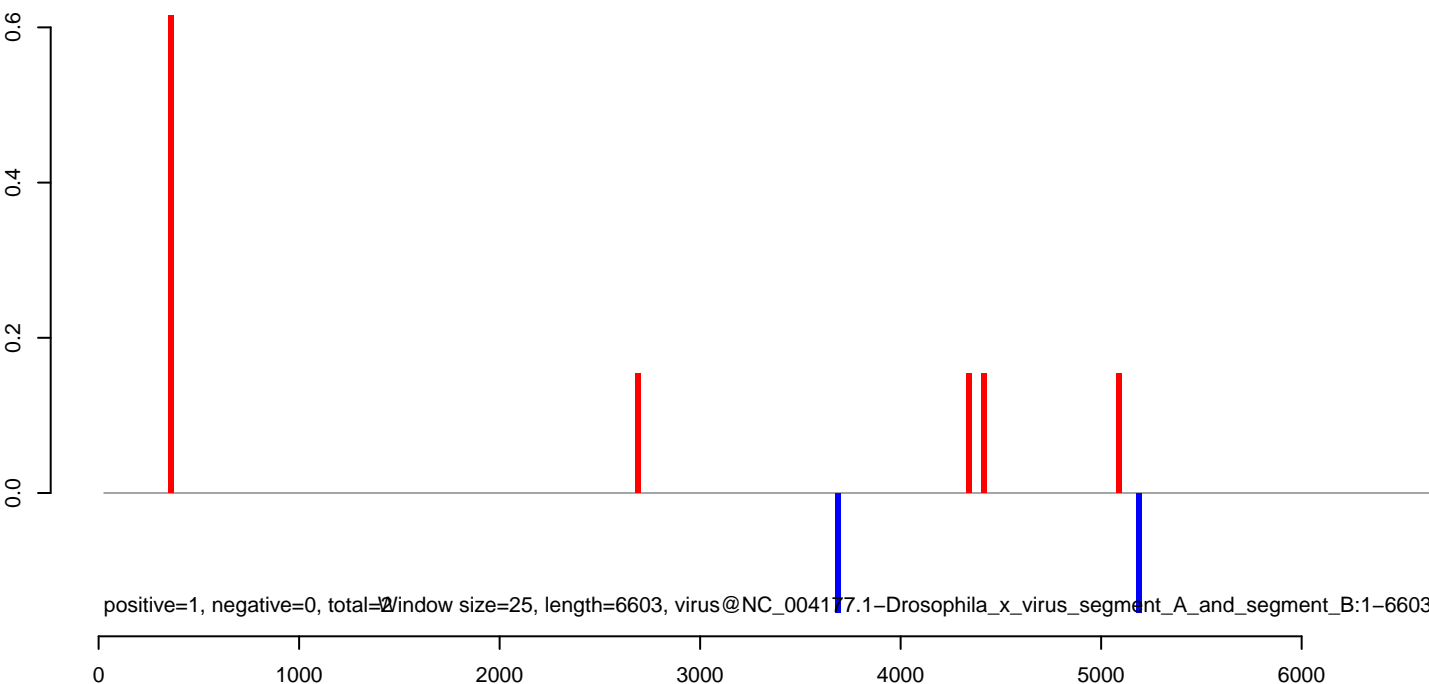
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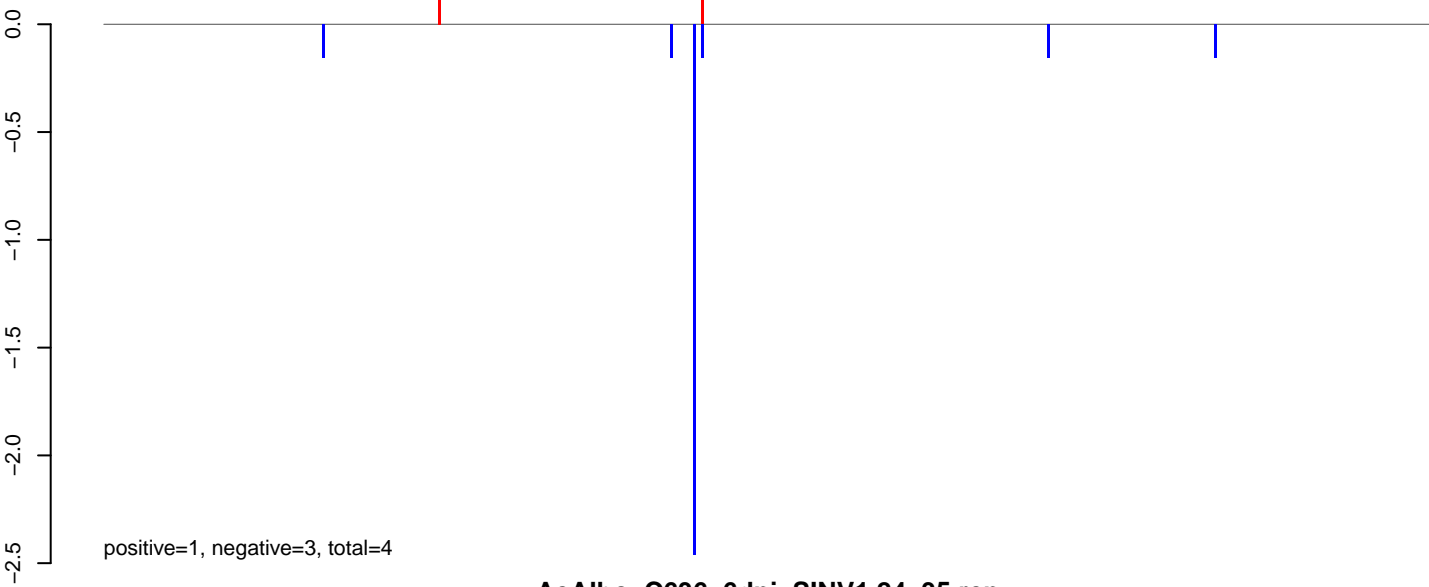
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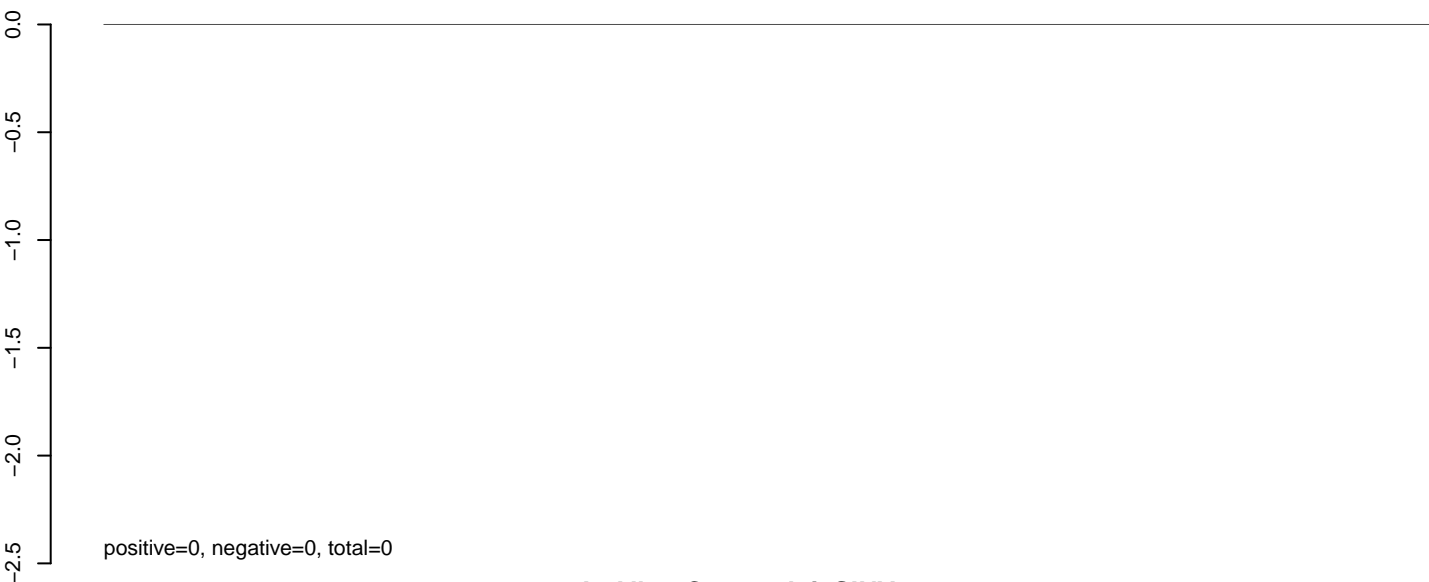
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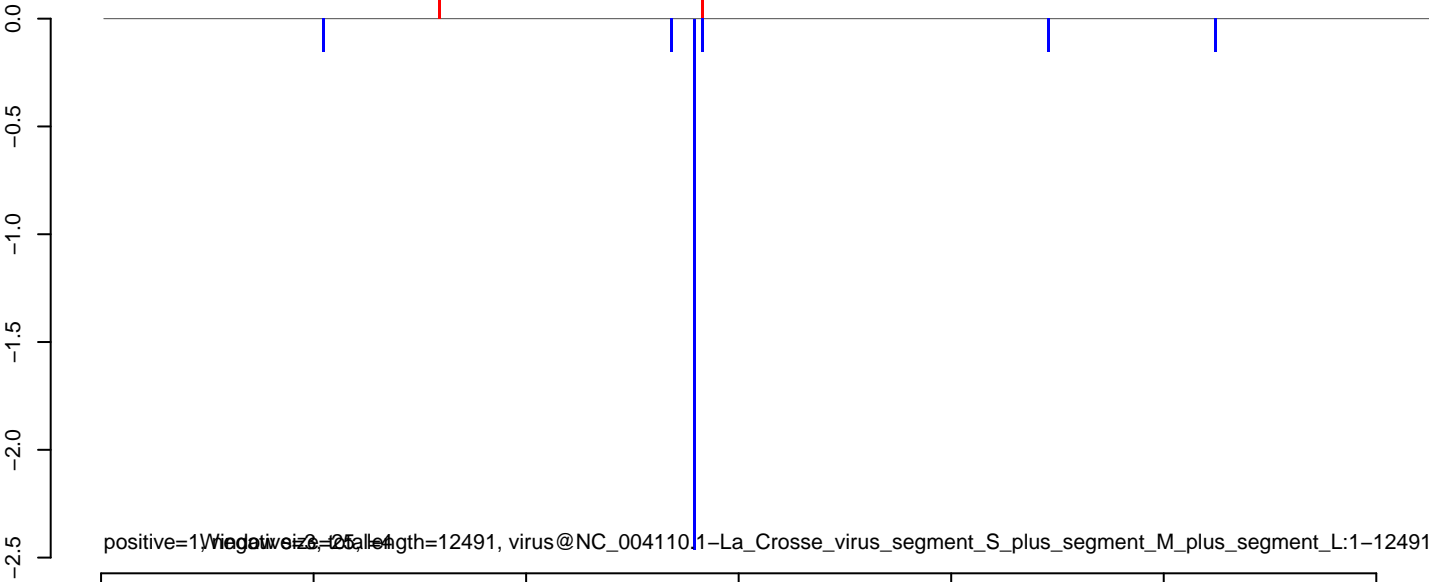
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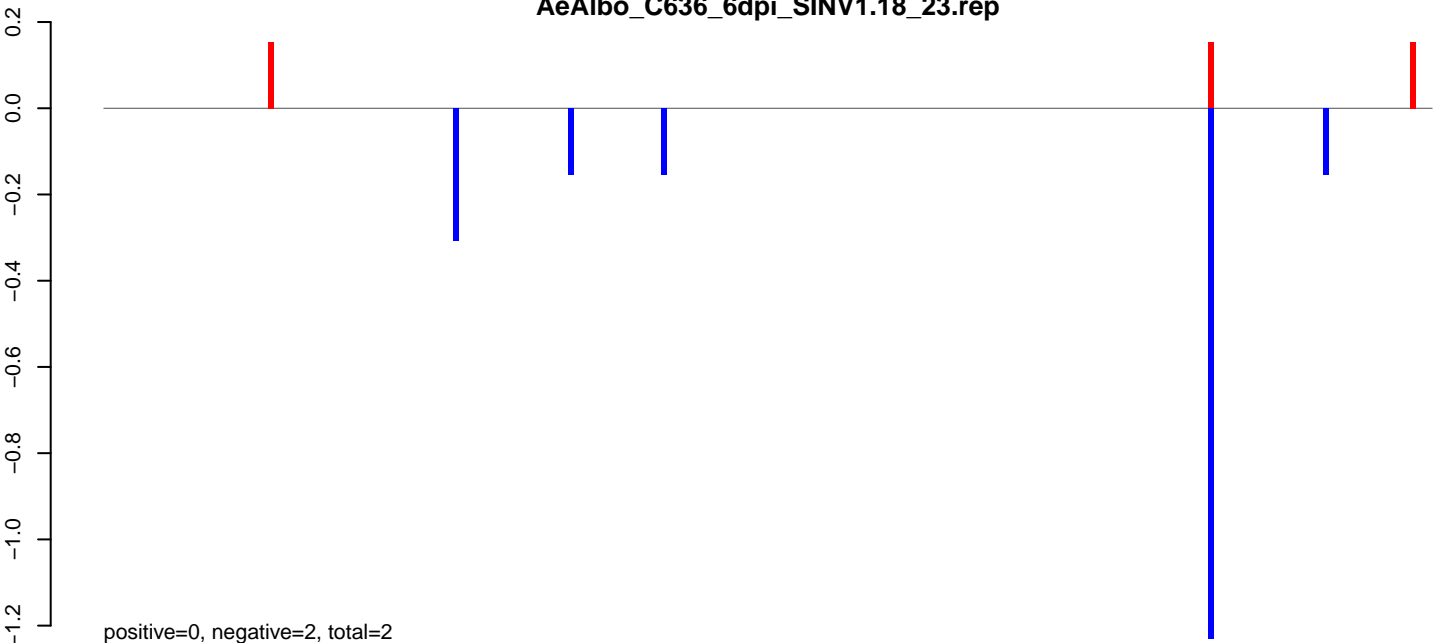
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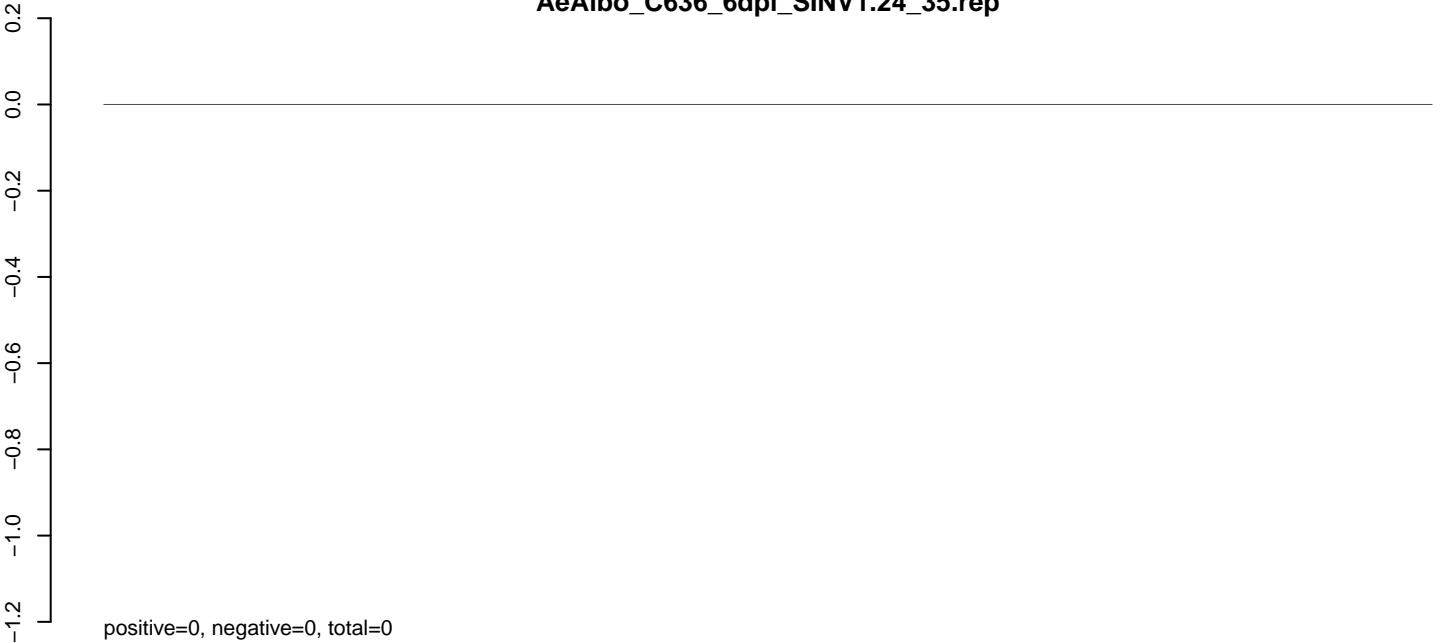
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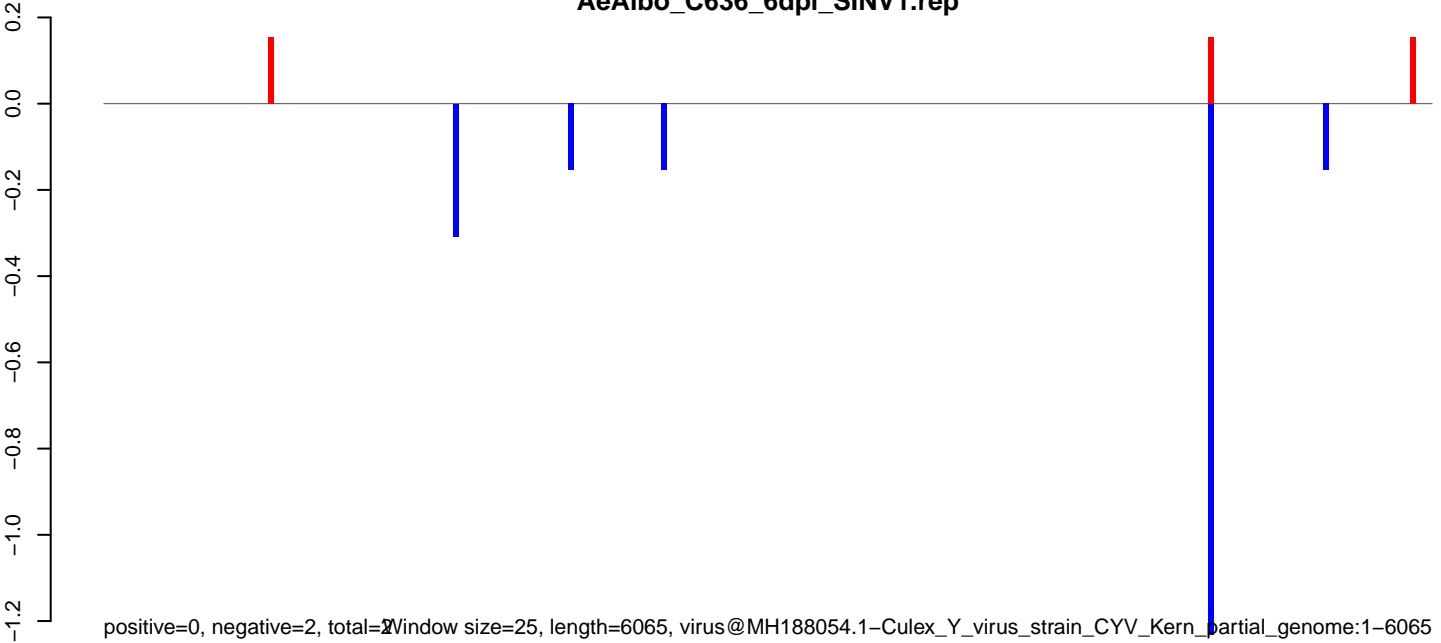
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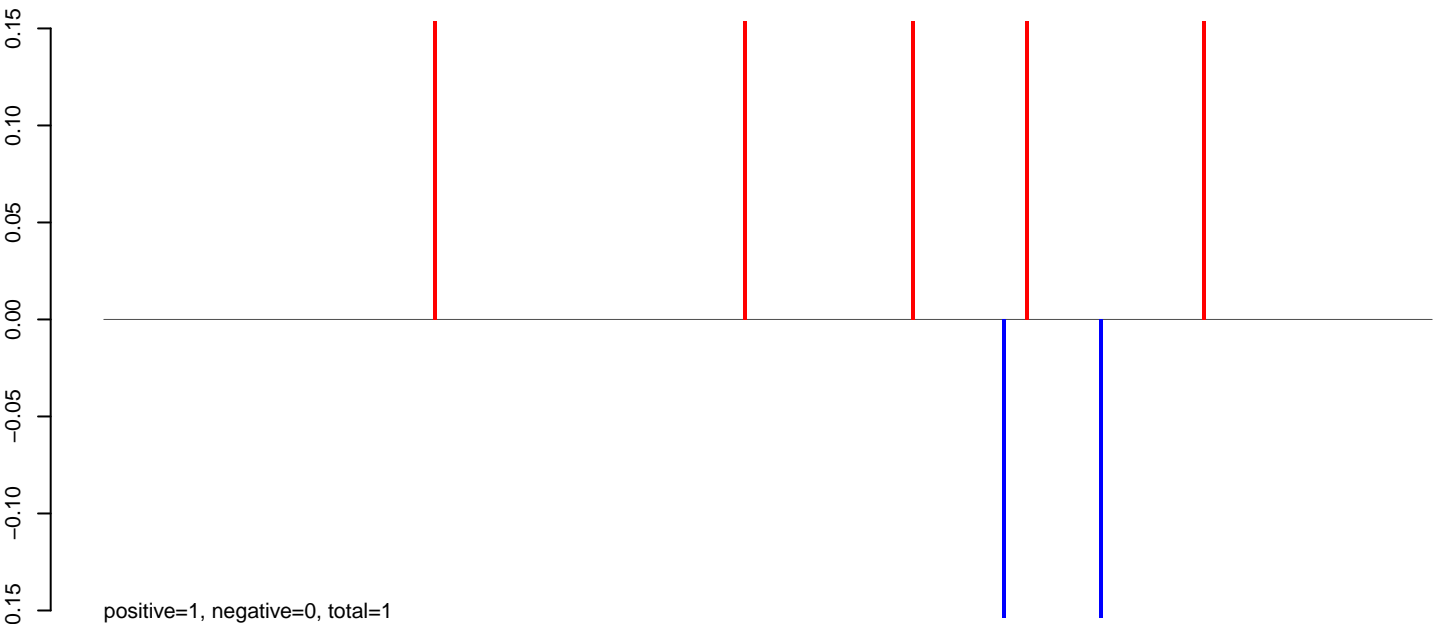
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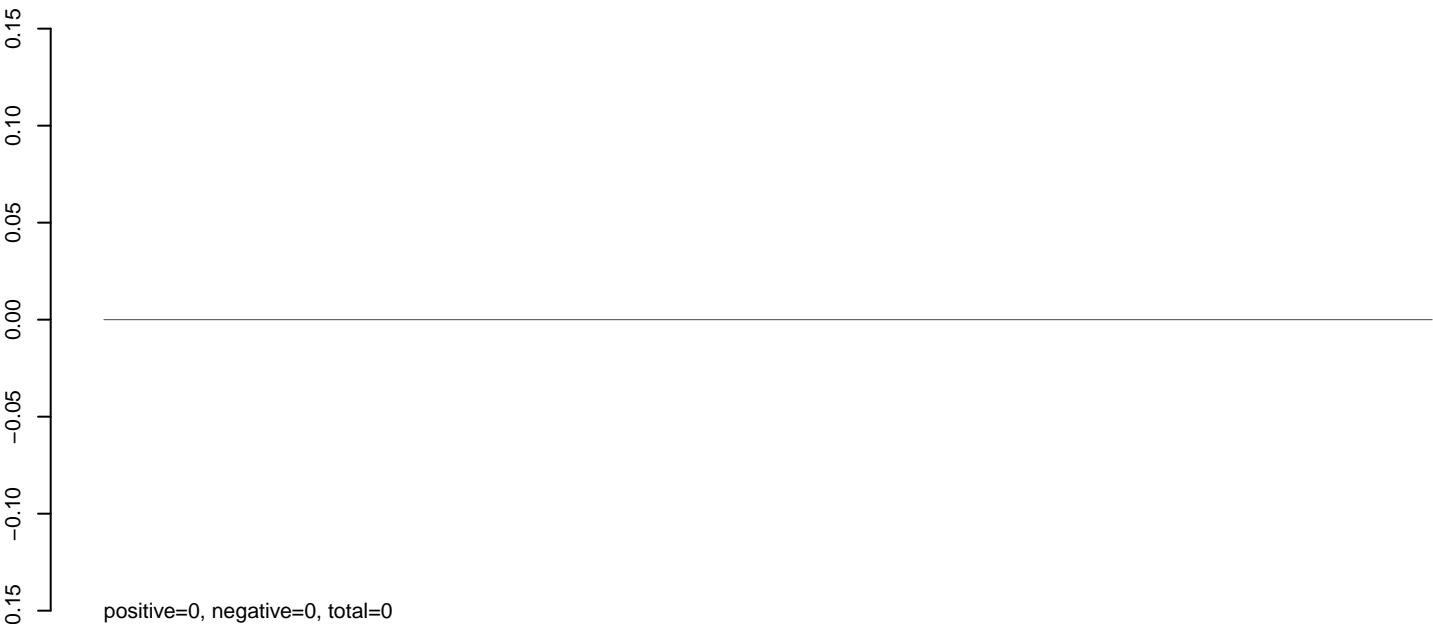
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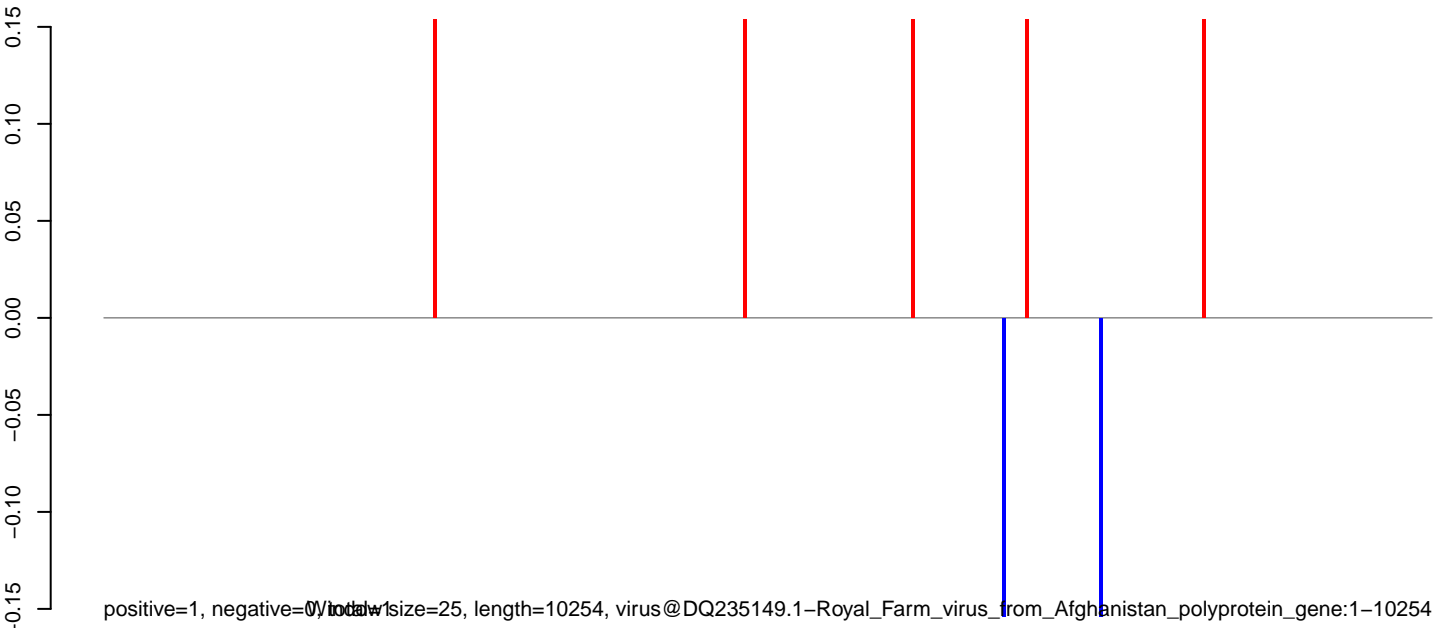
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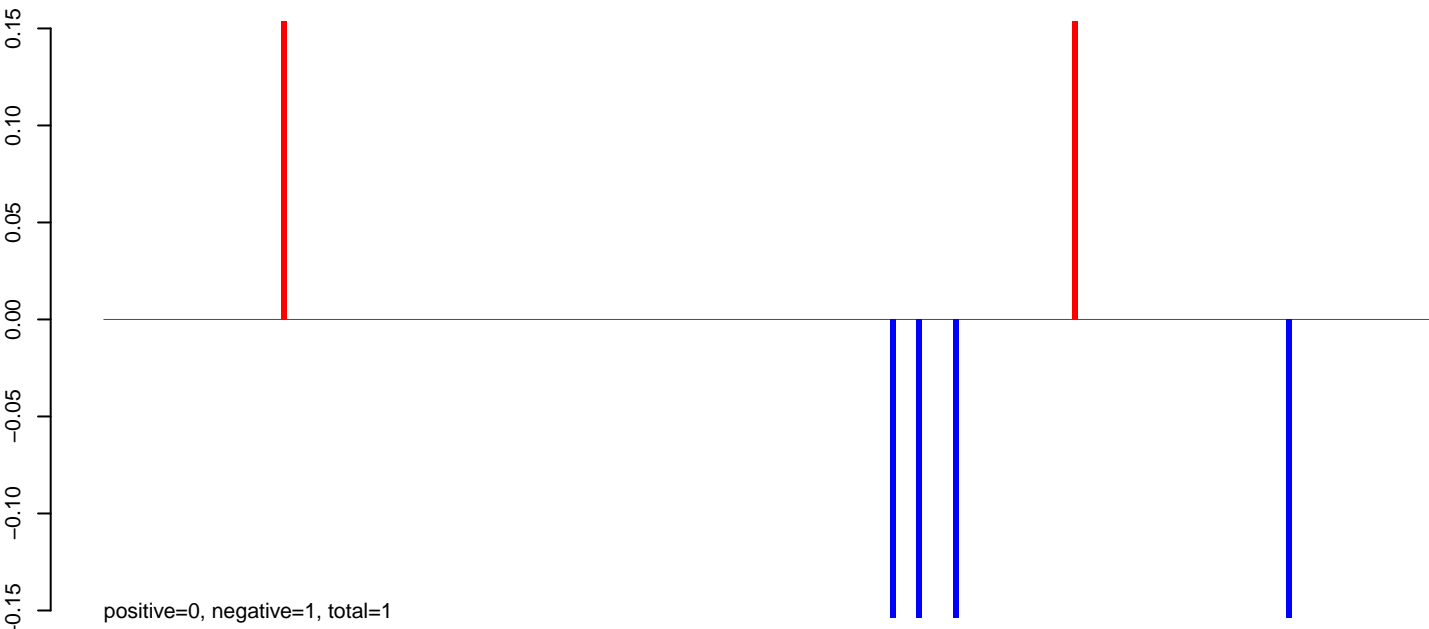
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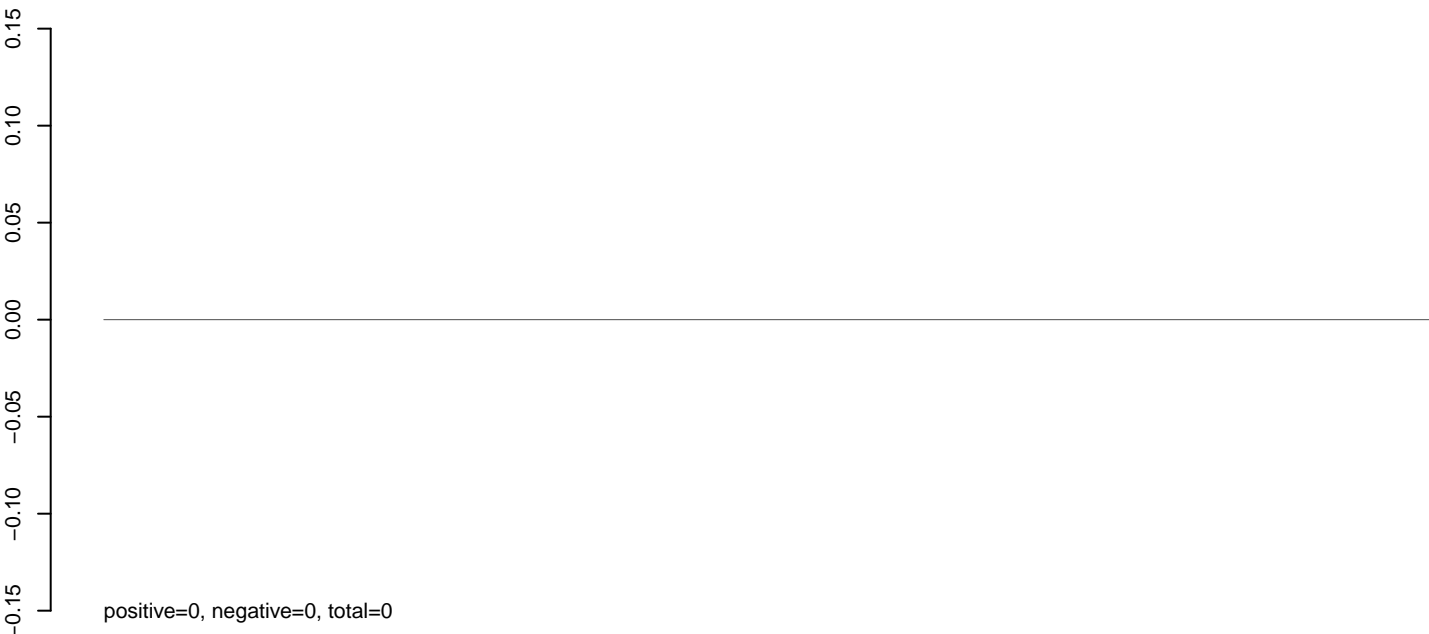
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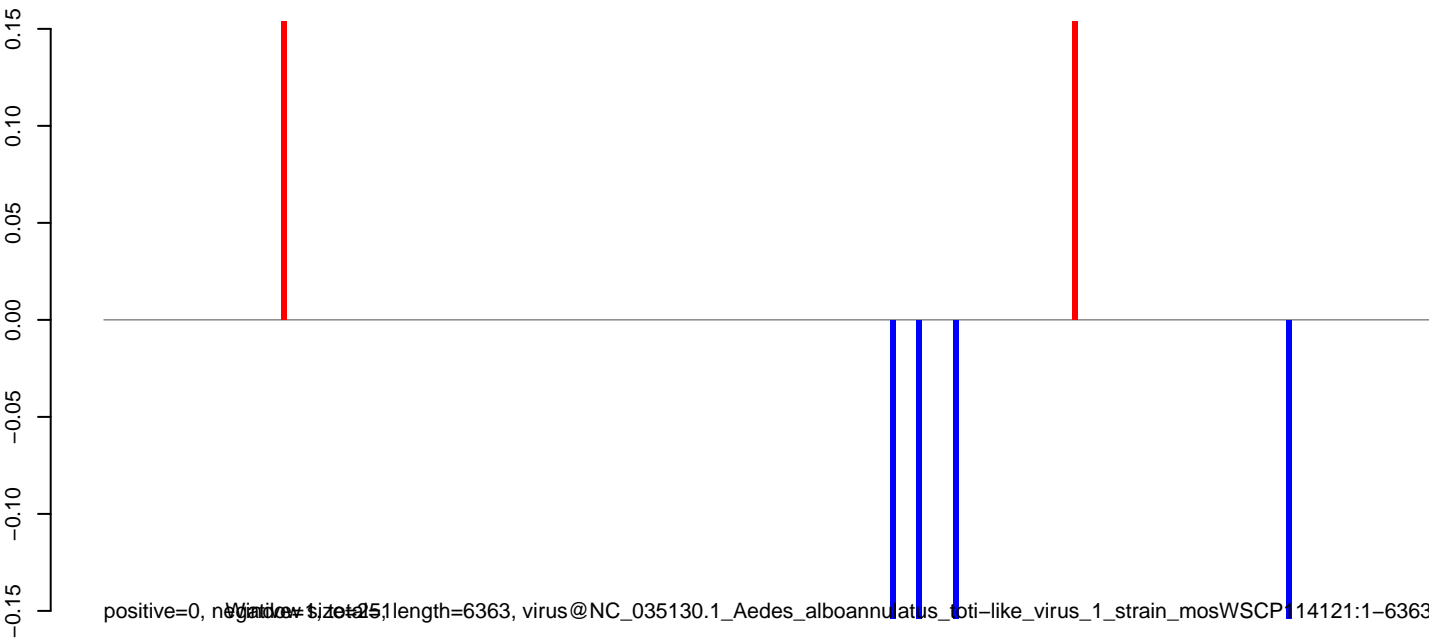
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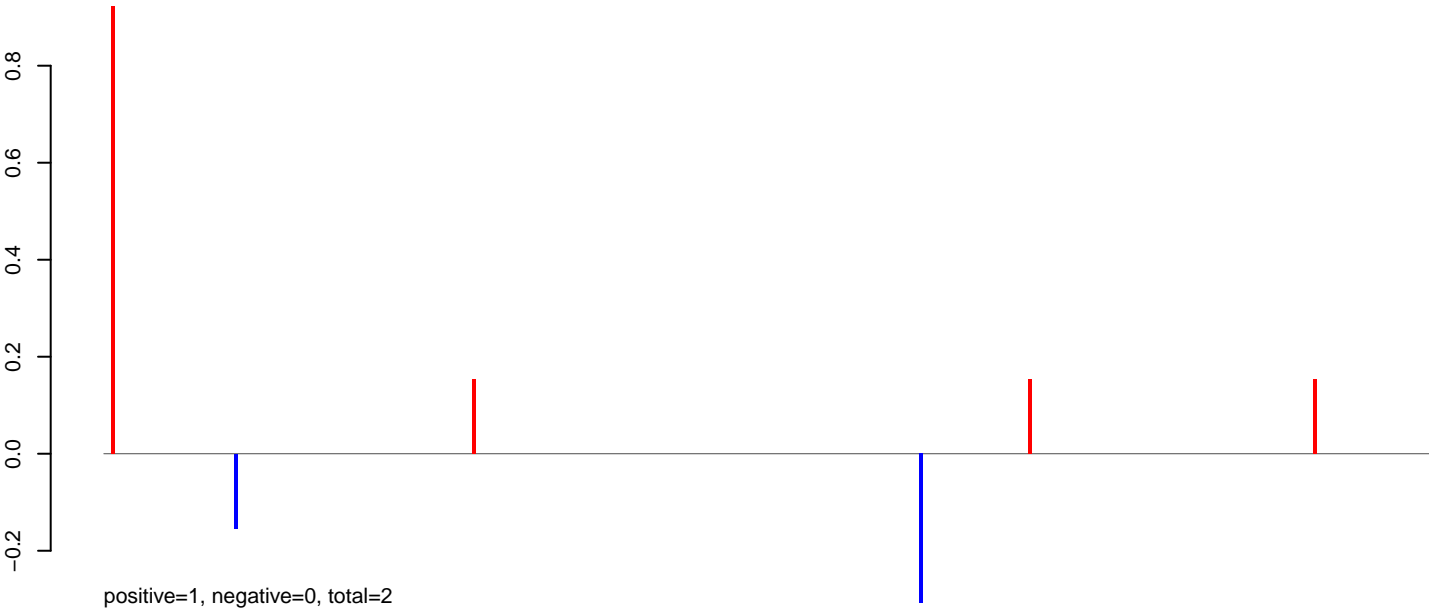
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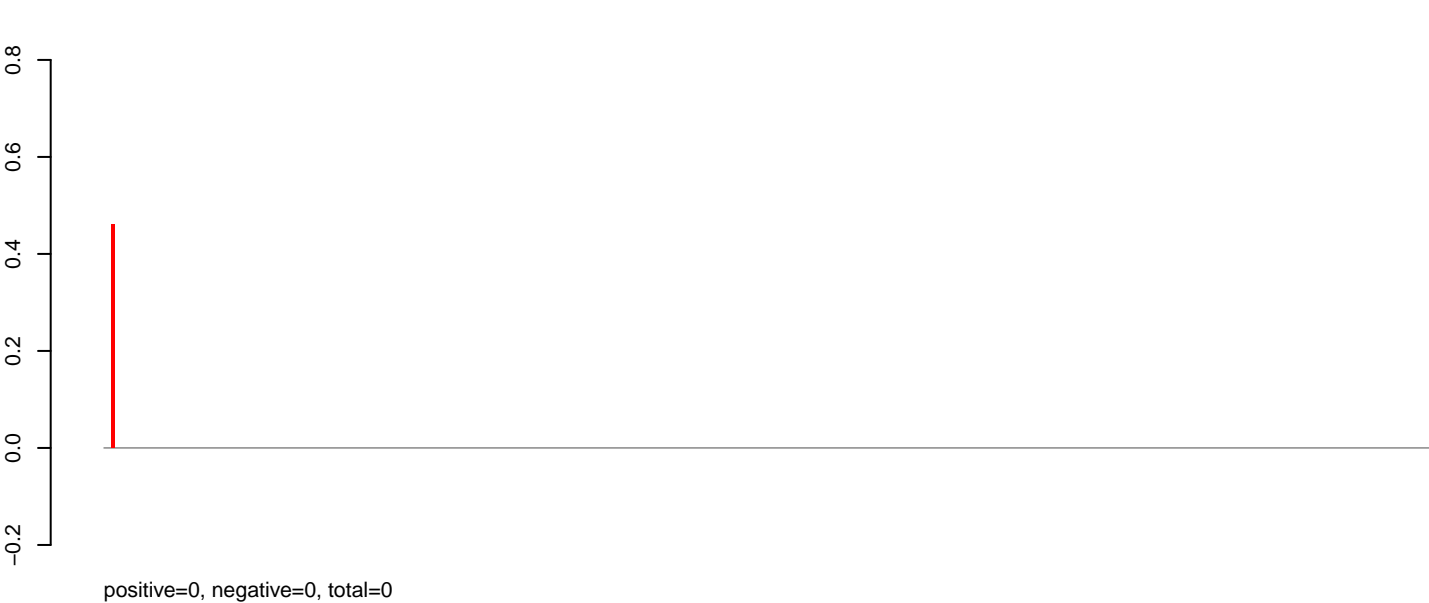
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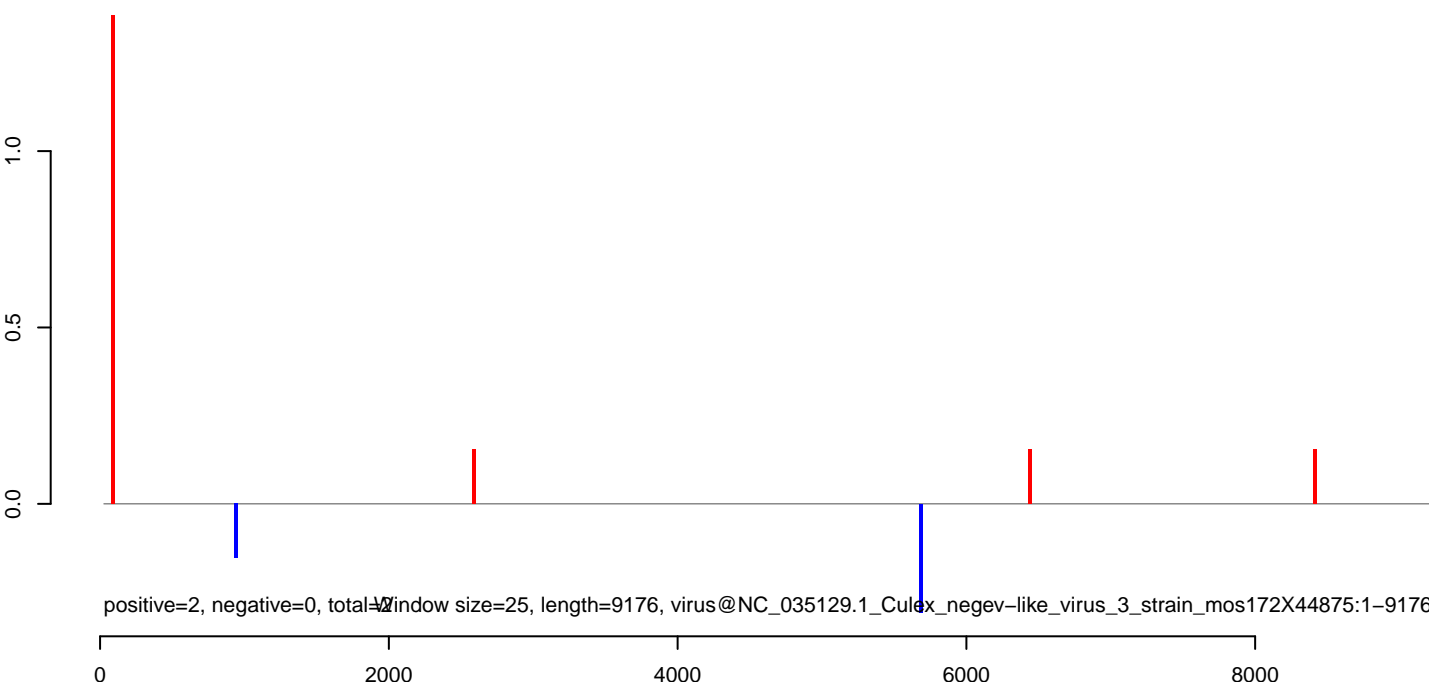
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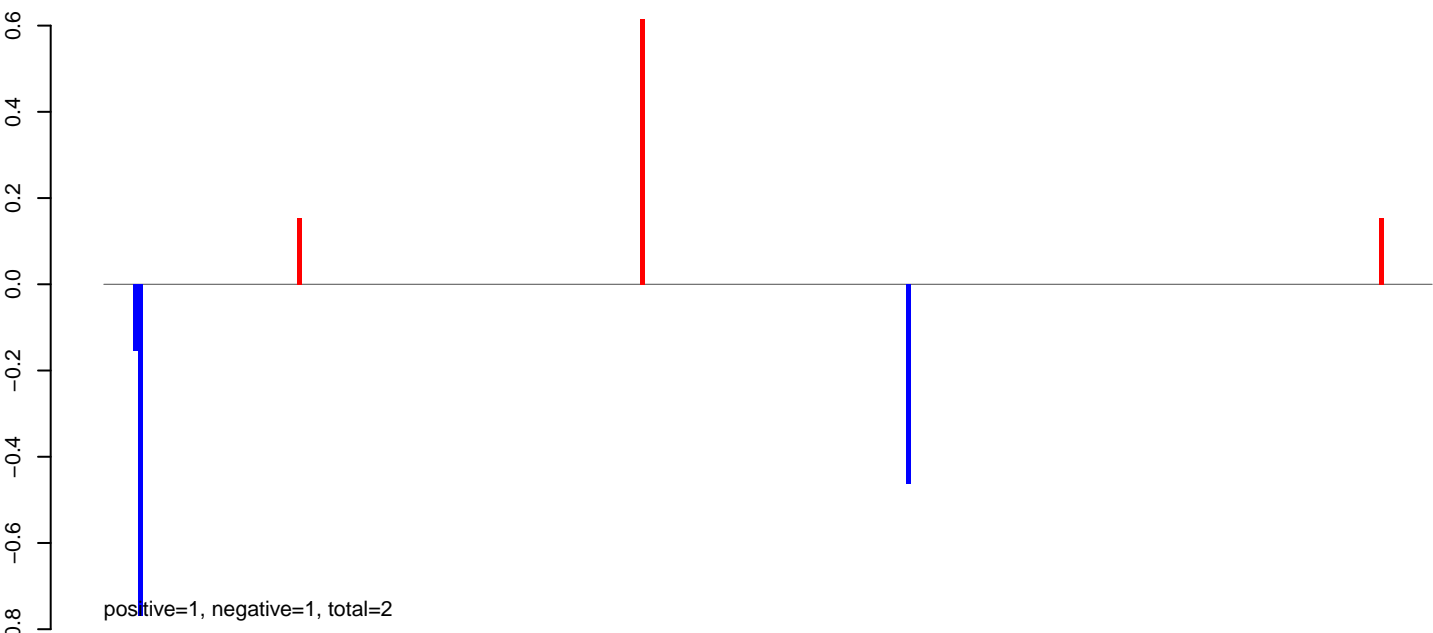
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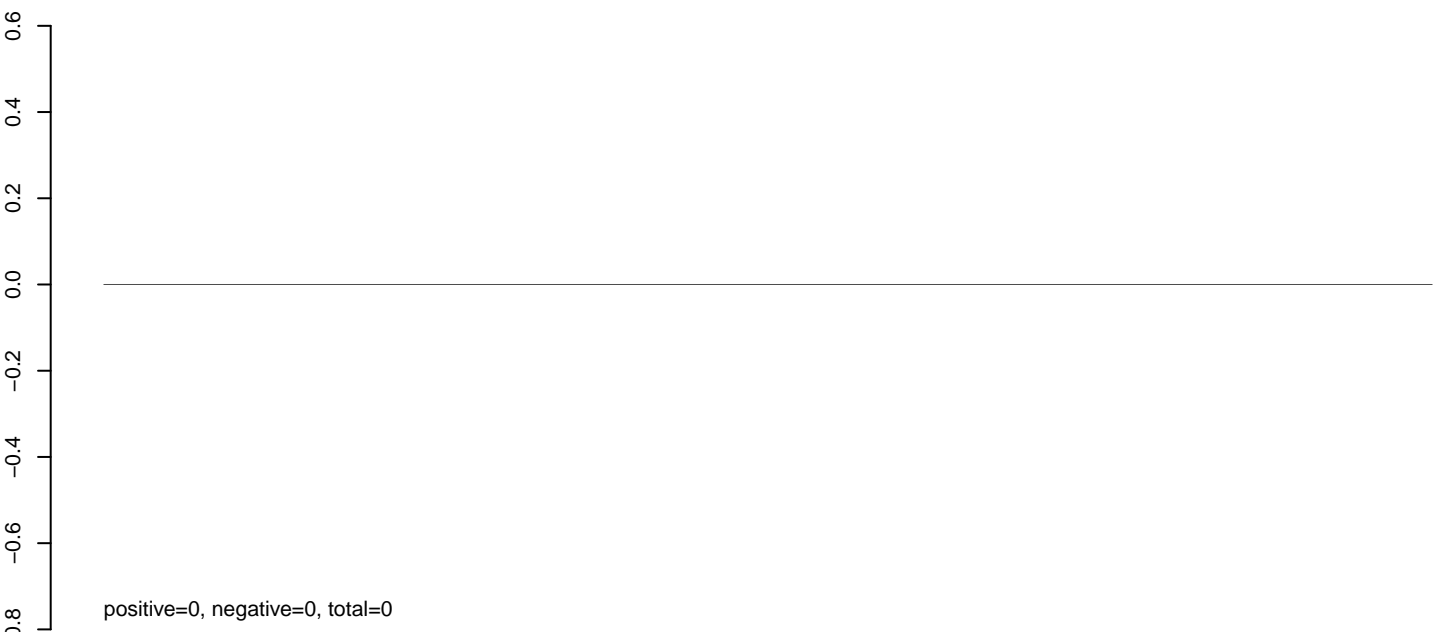
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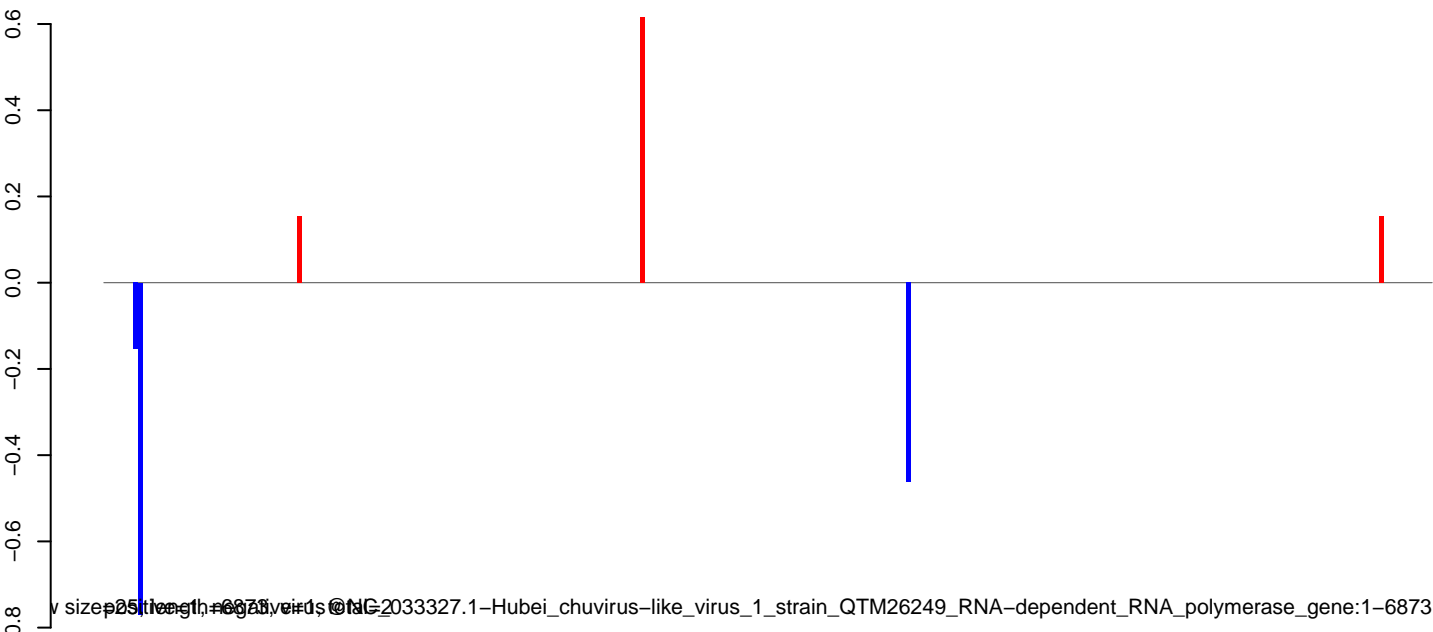
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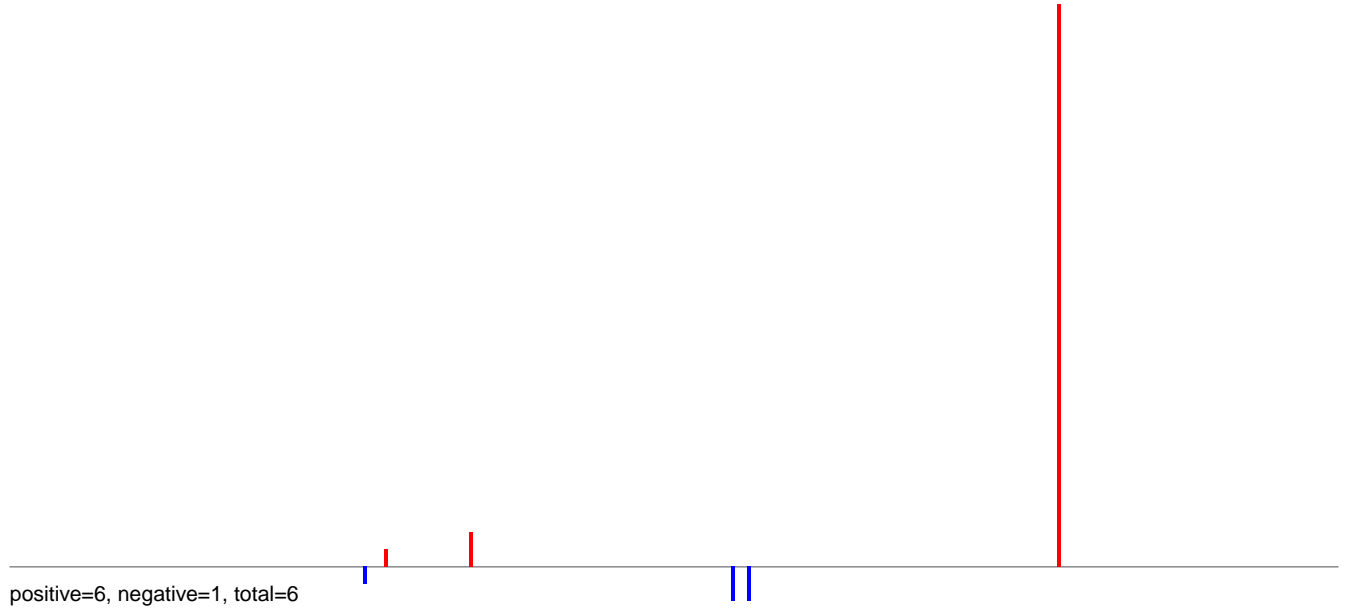
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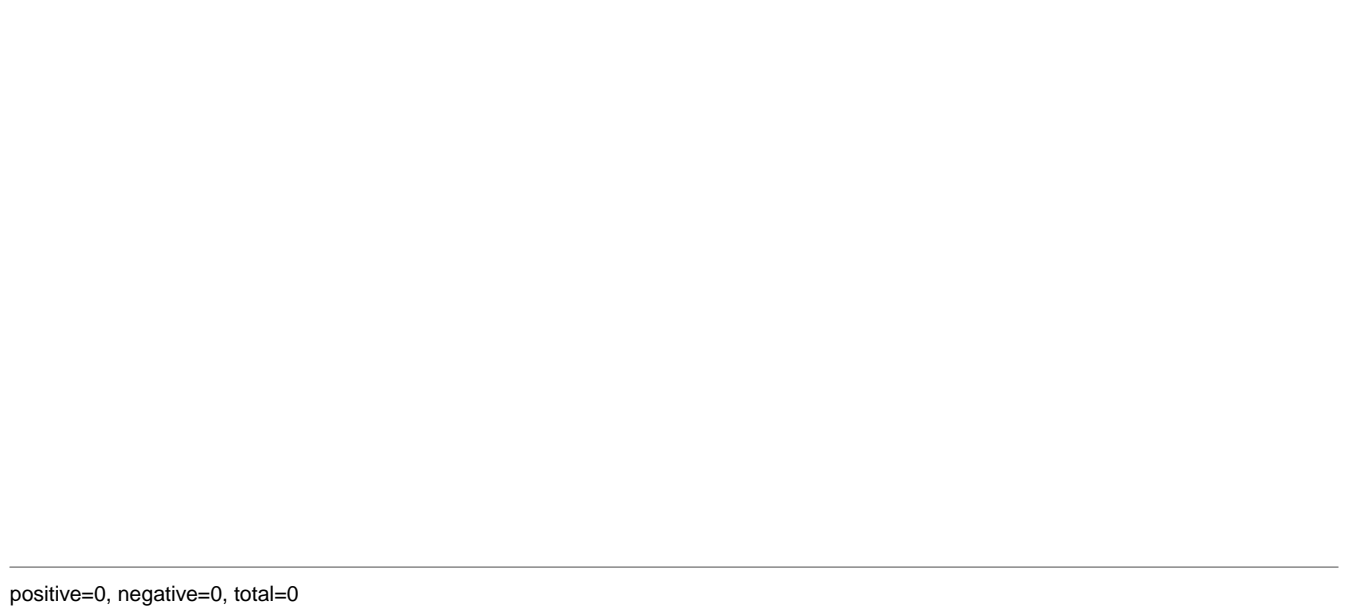
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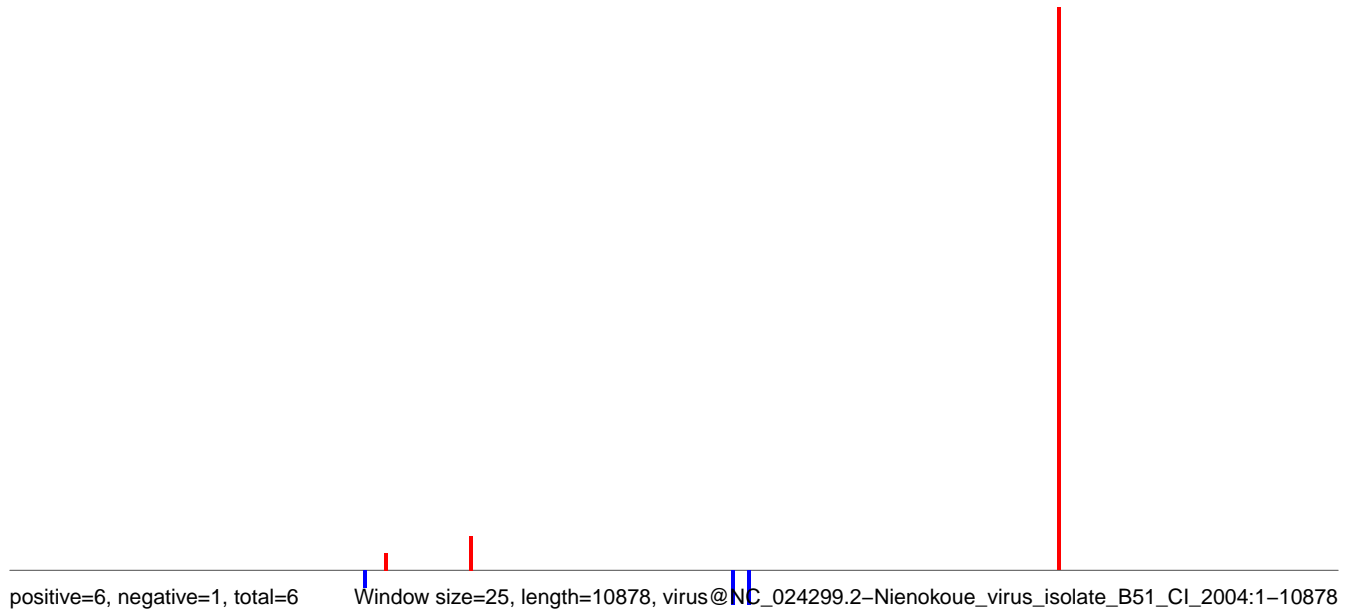
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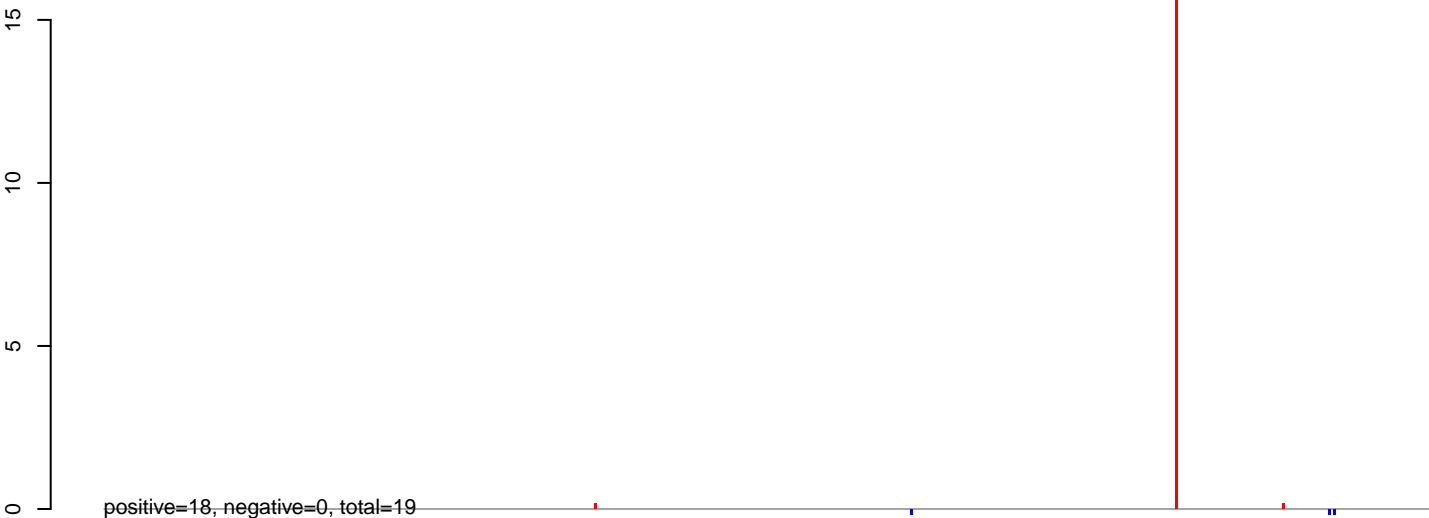


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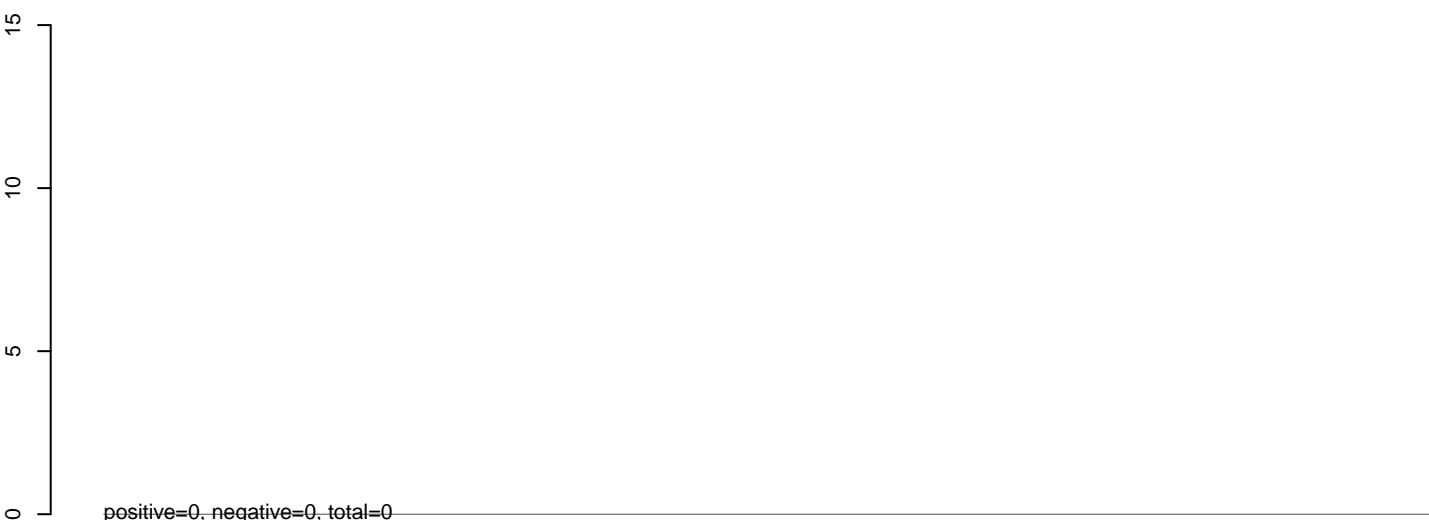


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AeAlbo_C636_6dpi_SINV1.18_23.rep



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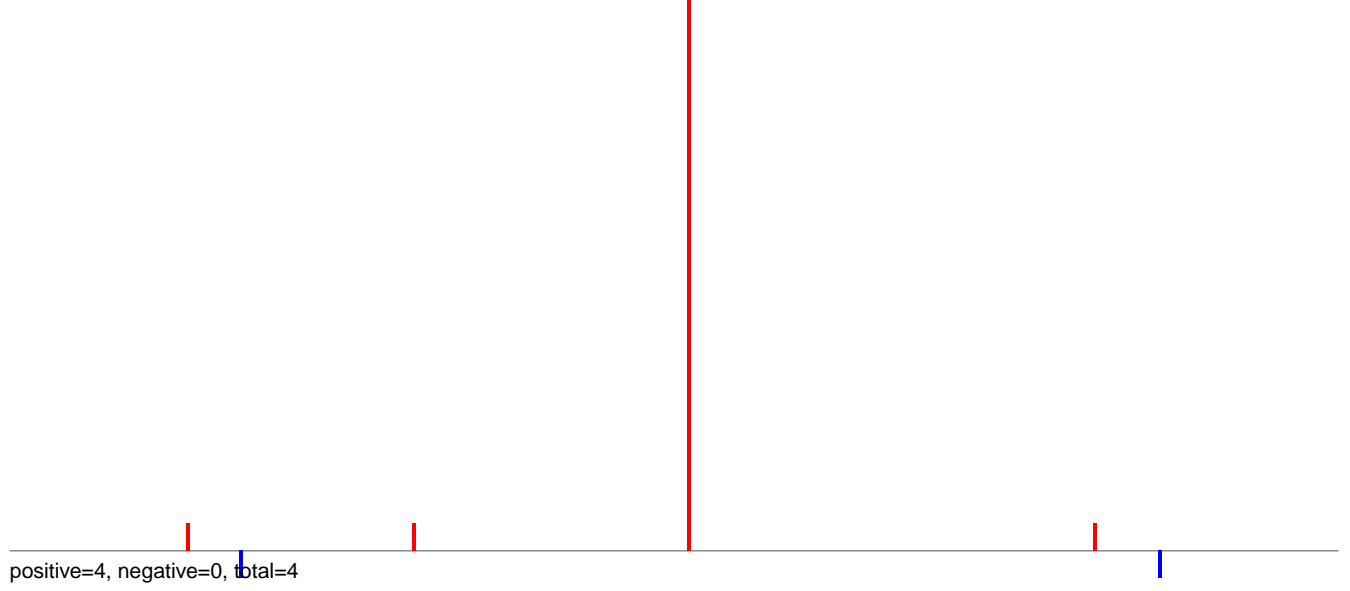
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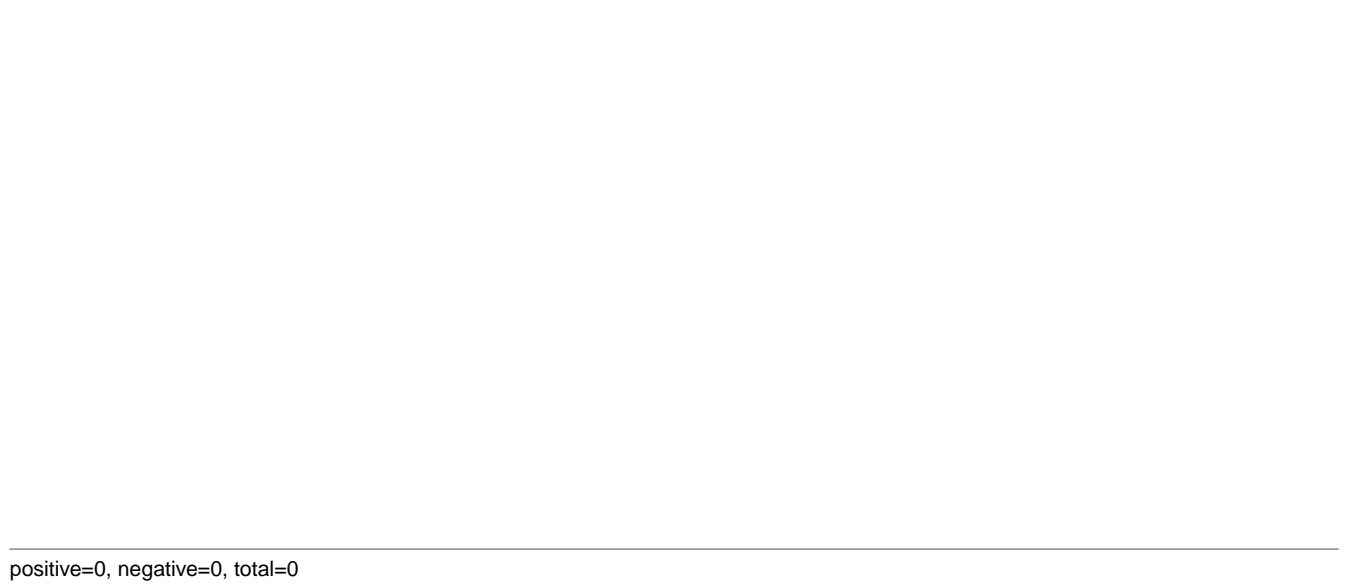
lengths: 2390, 19, 90, 3, 135, 449
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0 2000 4000 6000 8000 10000 12000

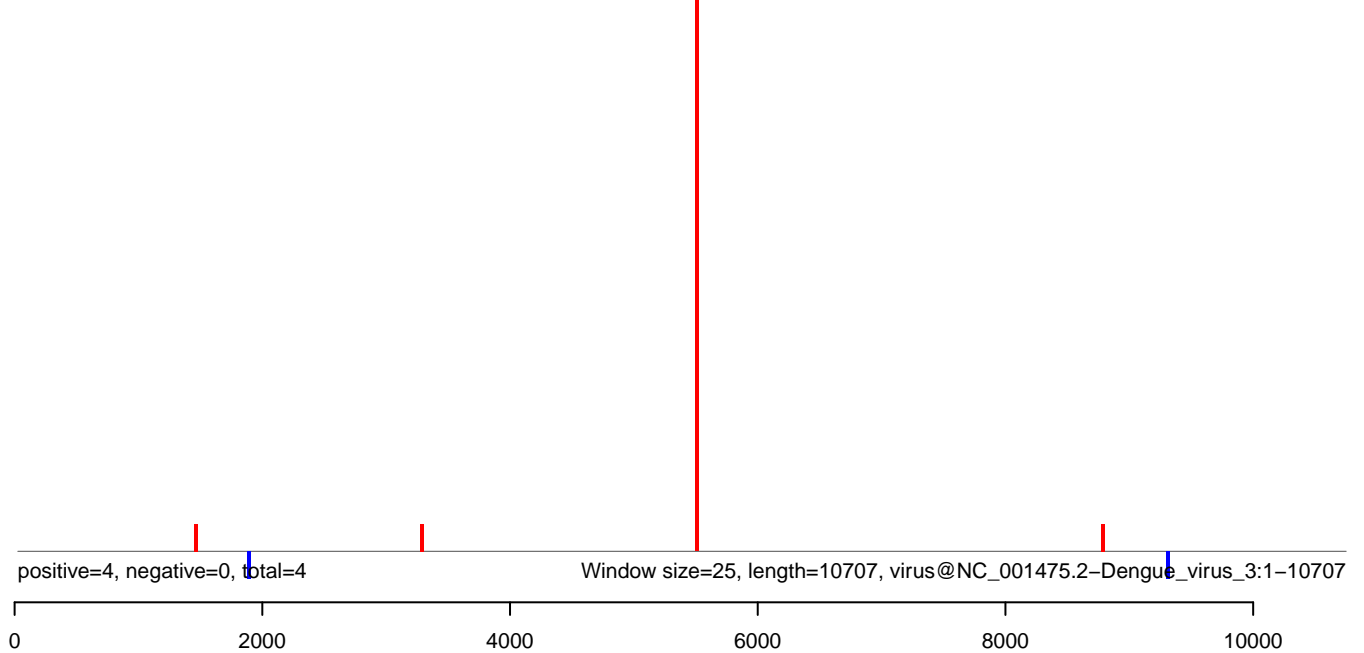
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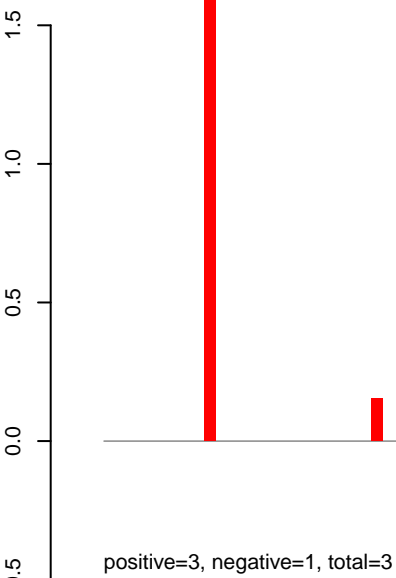
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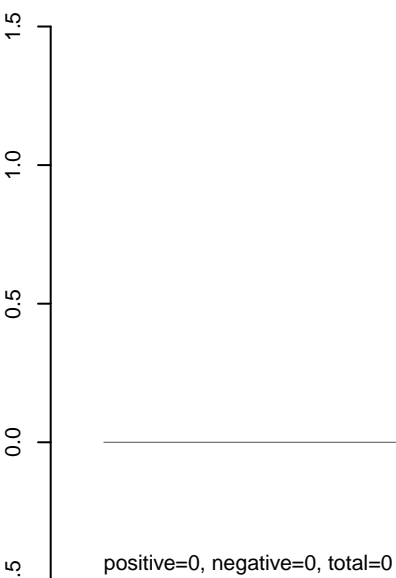
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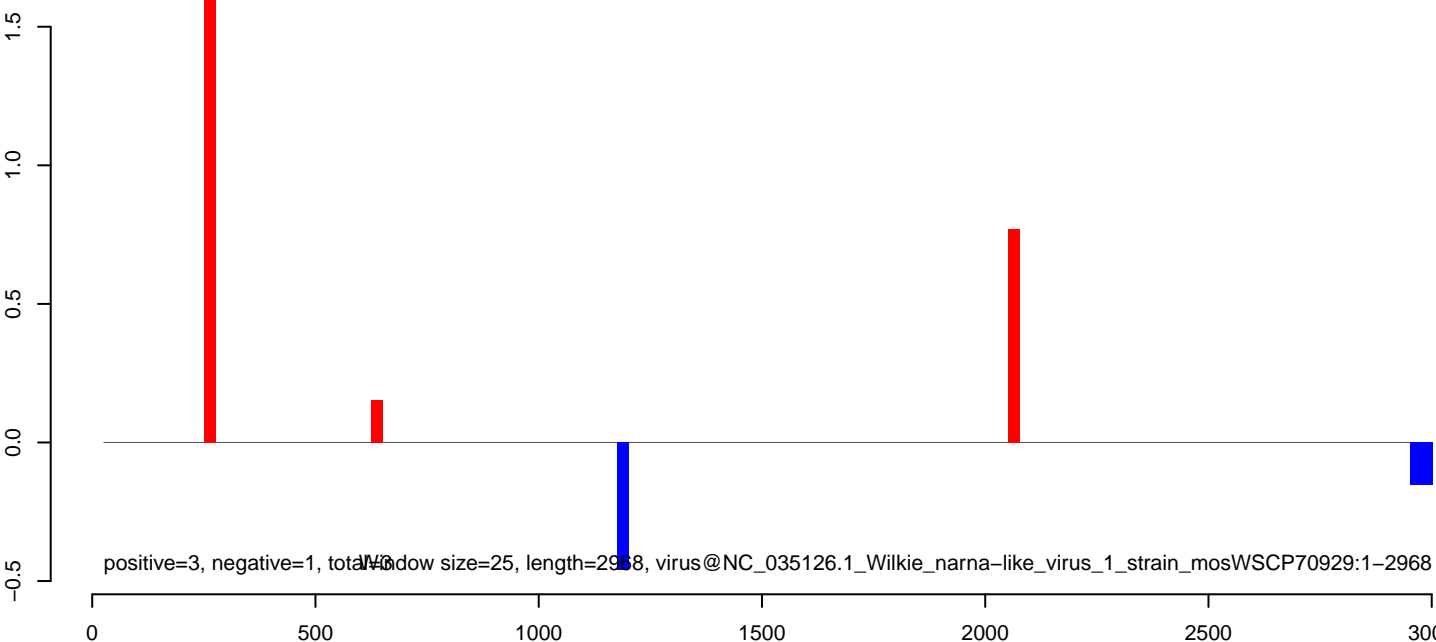
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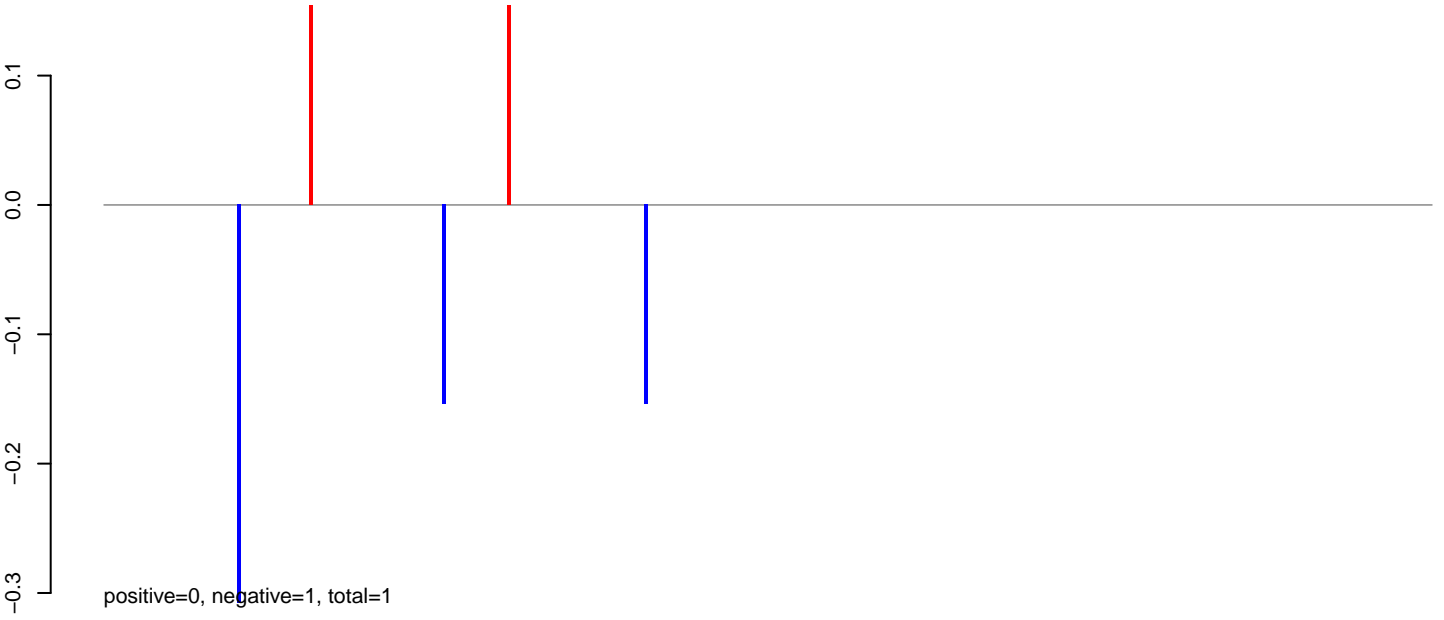
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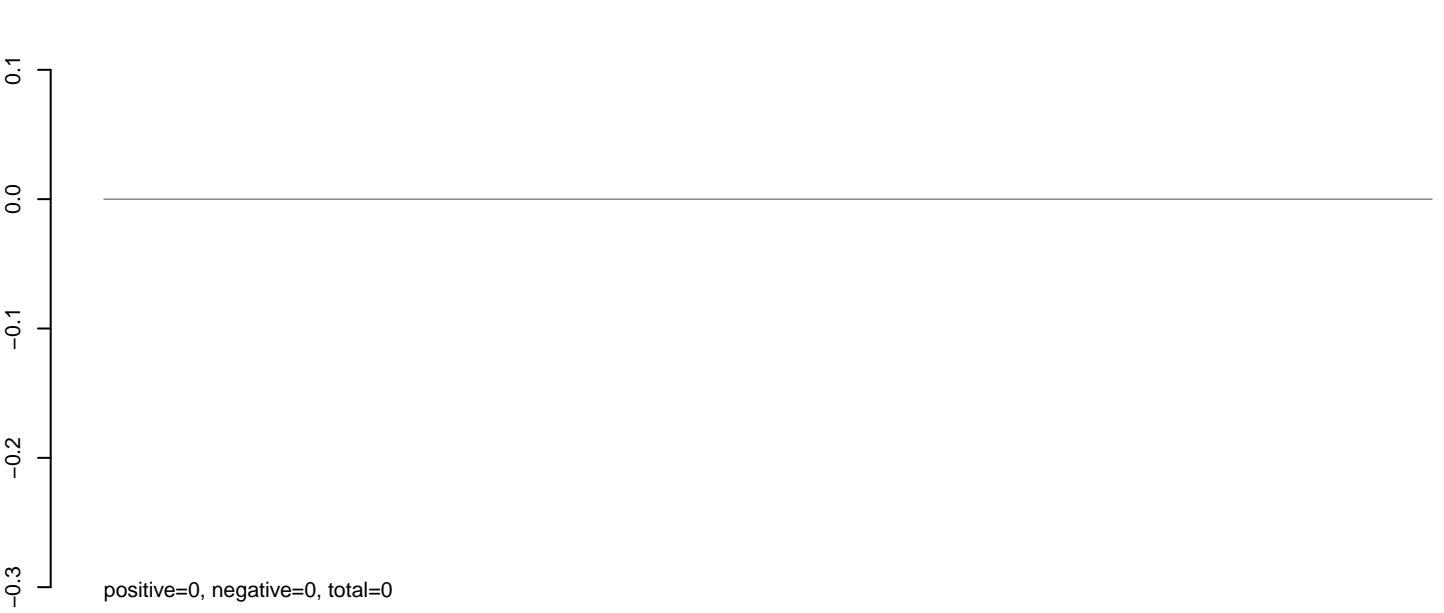
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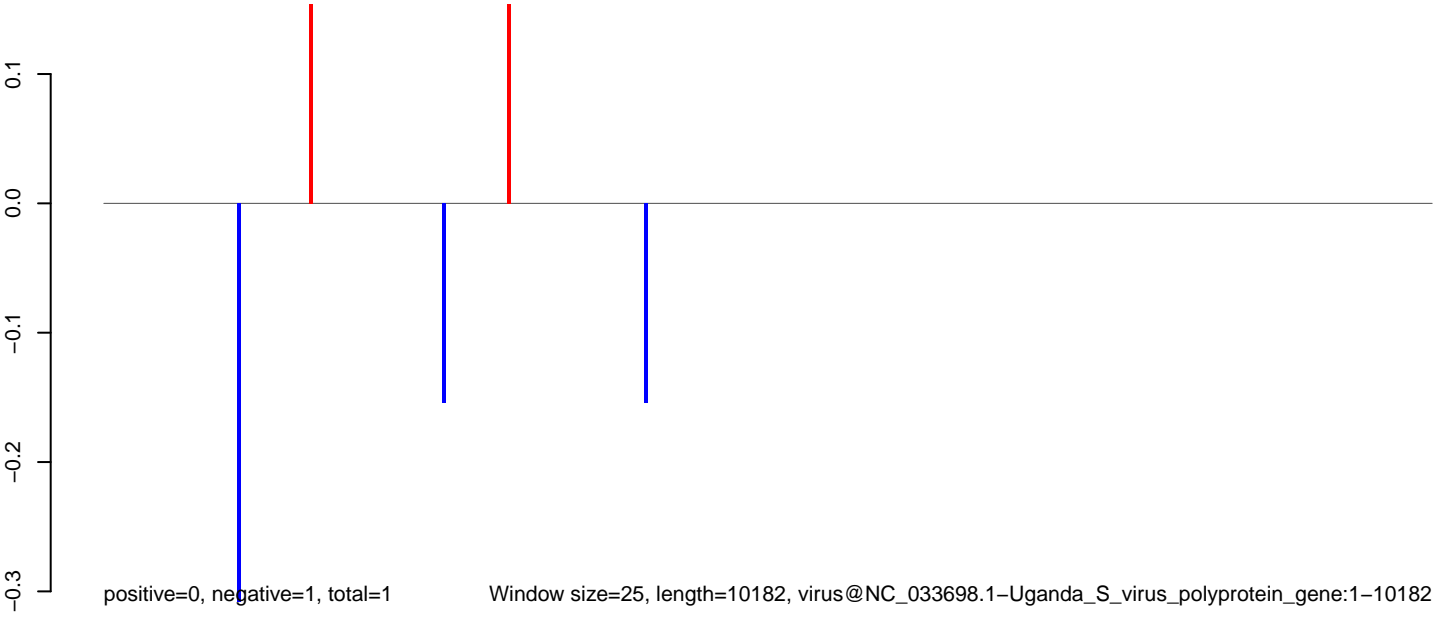
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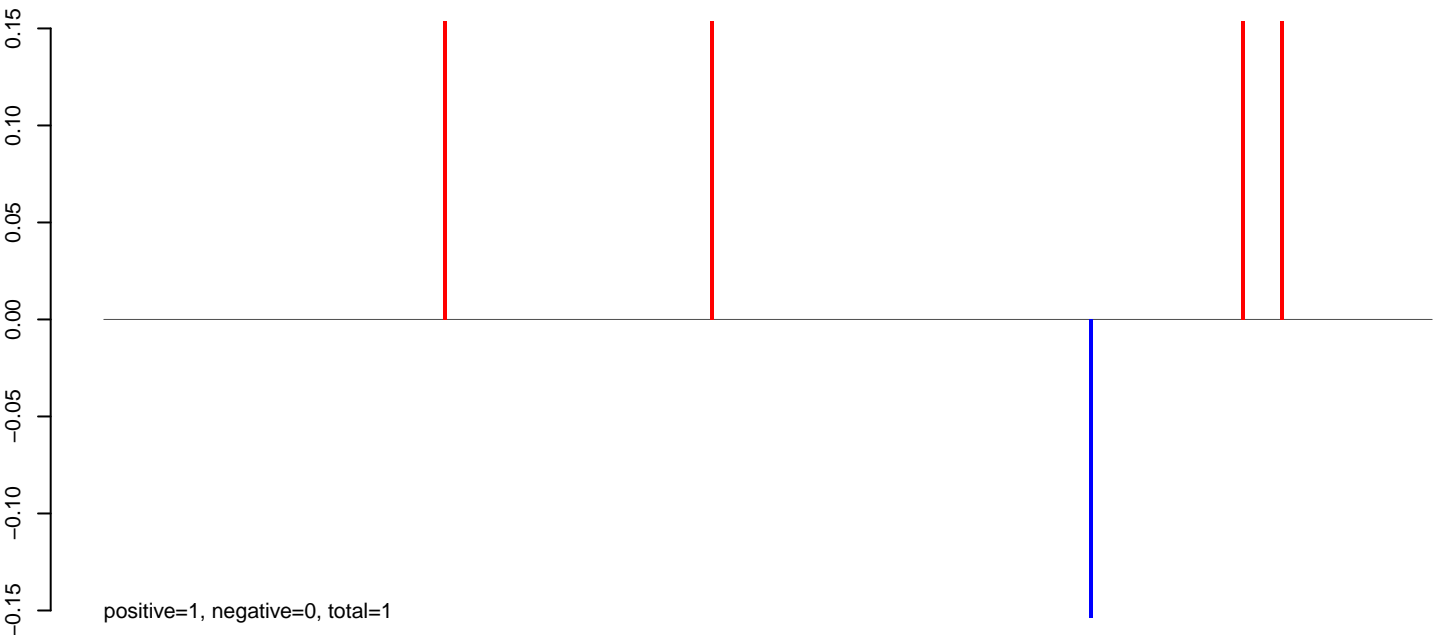
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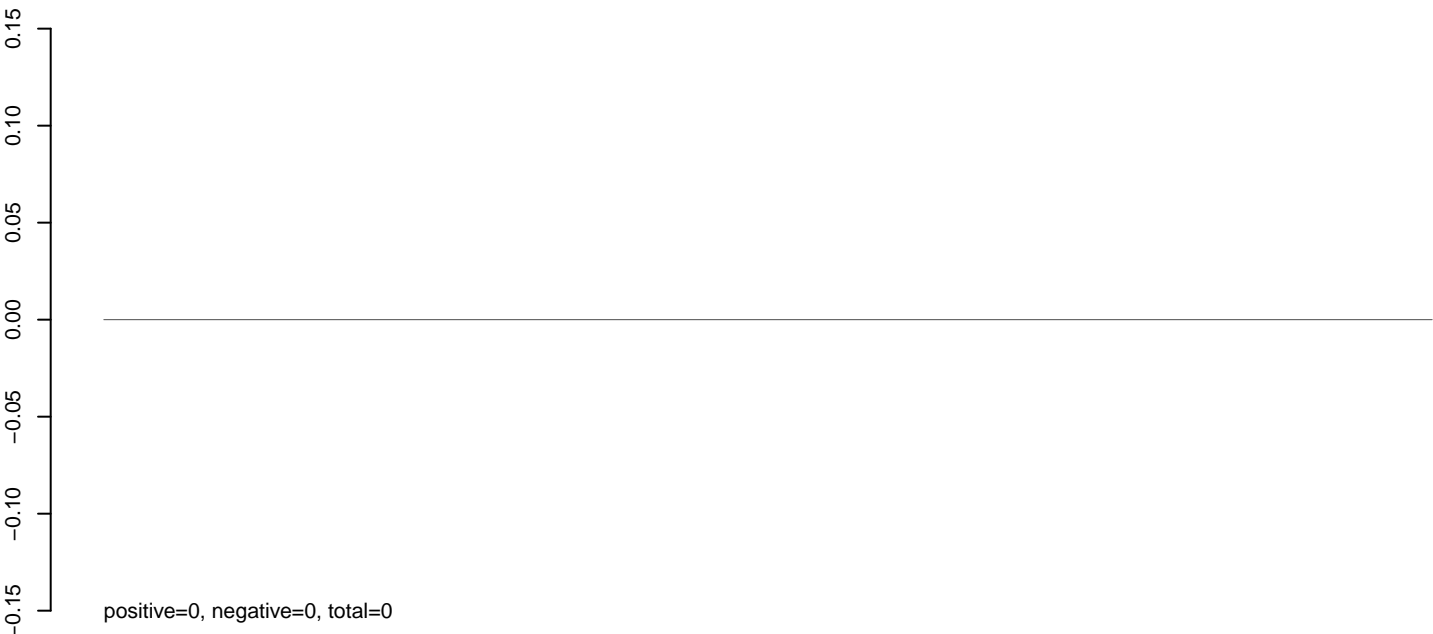
Window size=25, length=10182, virus@NC_033698.1-Uganda_S_virus_polyprotein_gene:1-10182

0 2000 4000 6000 8000 10000

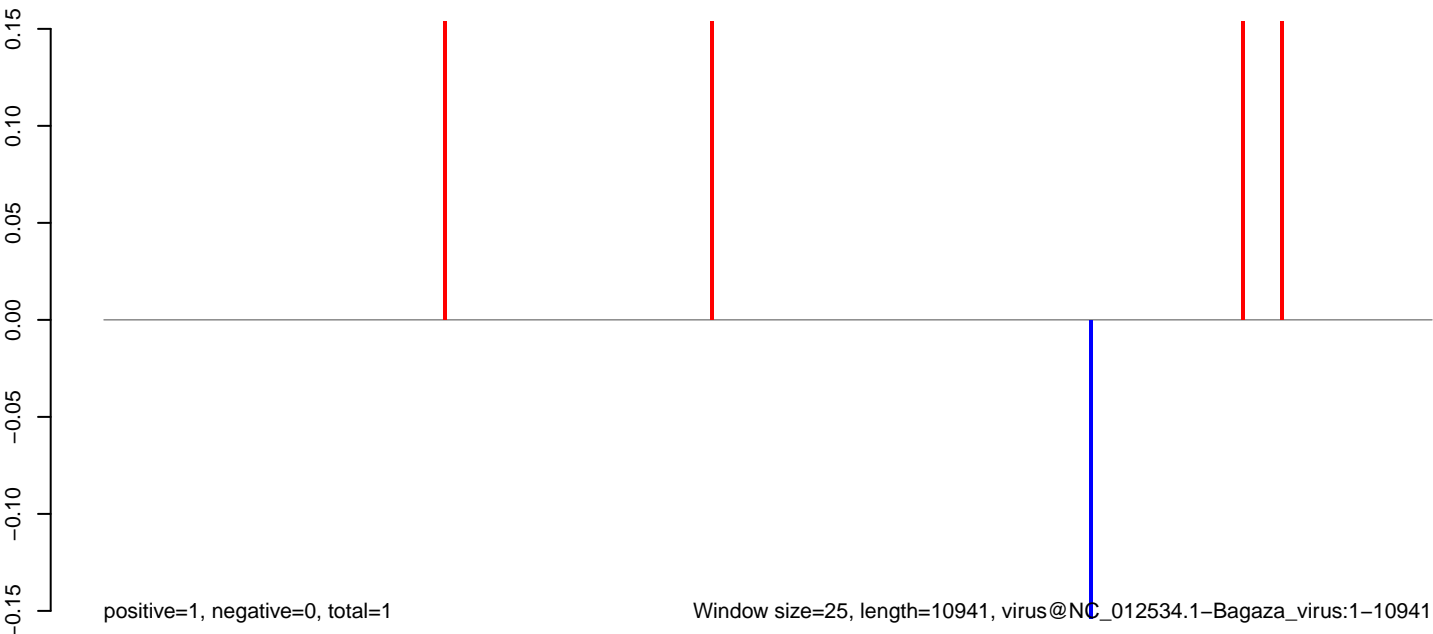
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



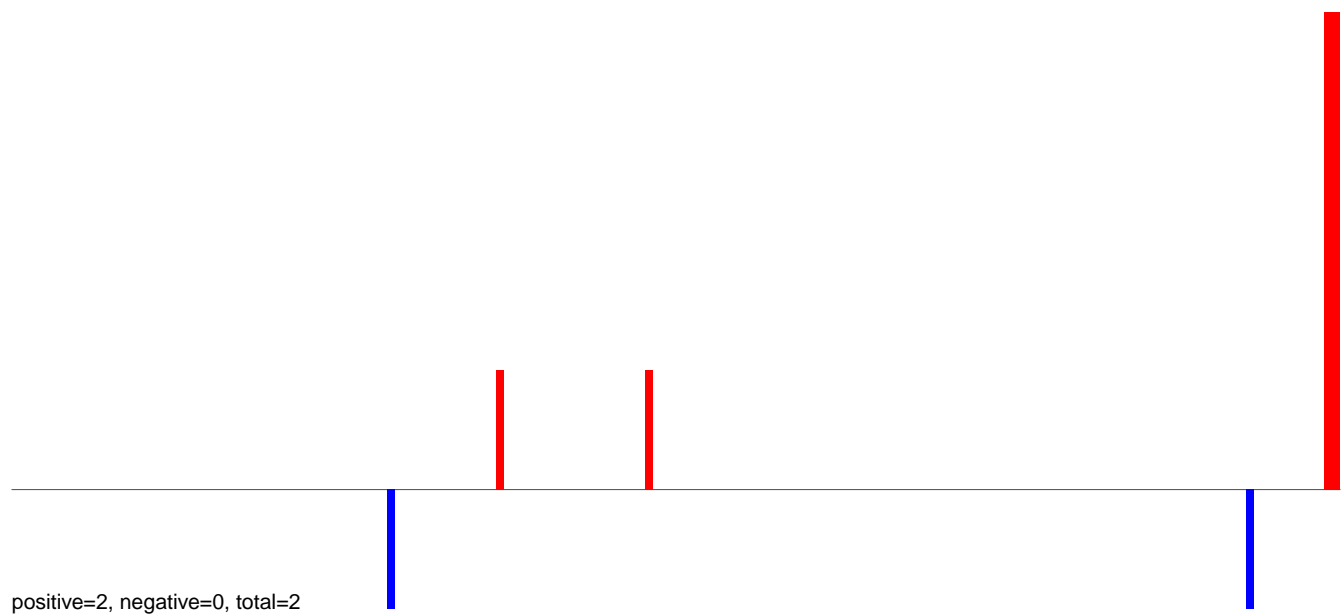
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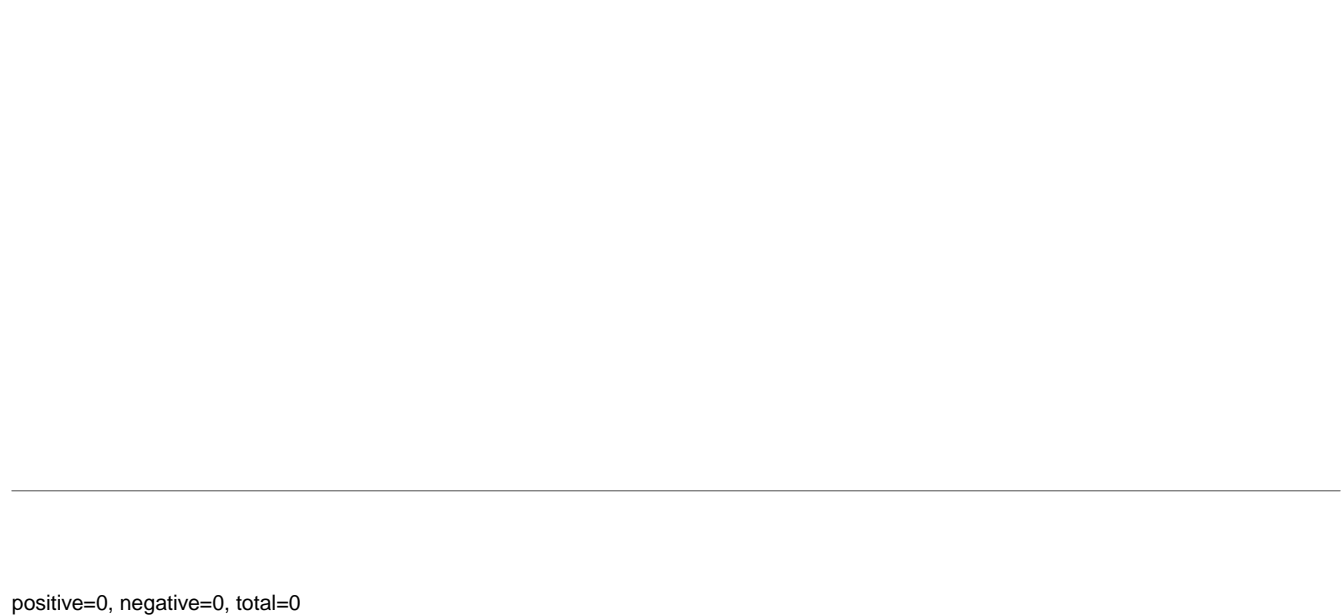
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0 2000 4000 6000 8000 10000

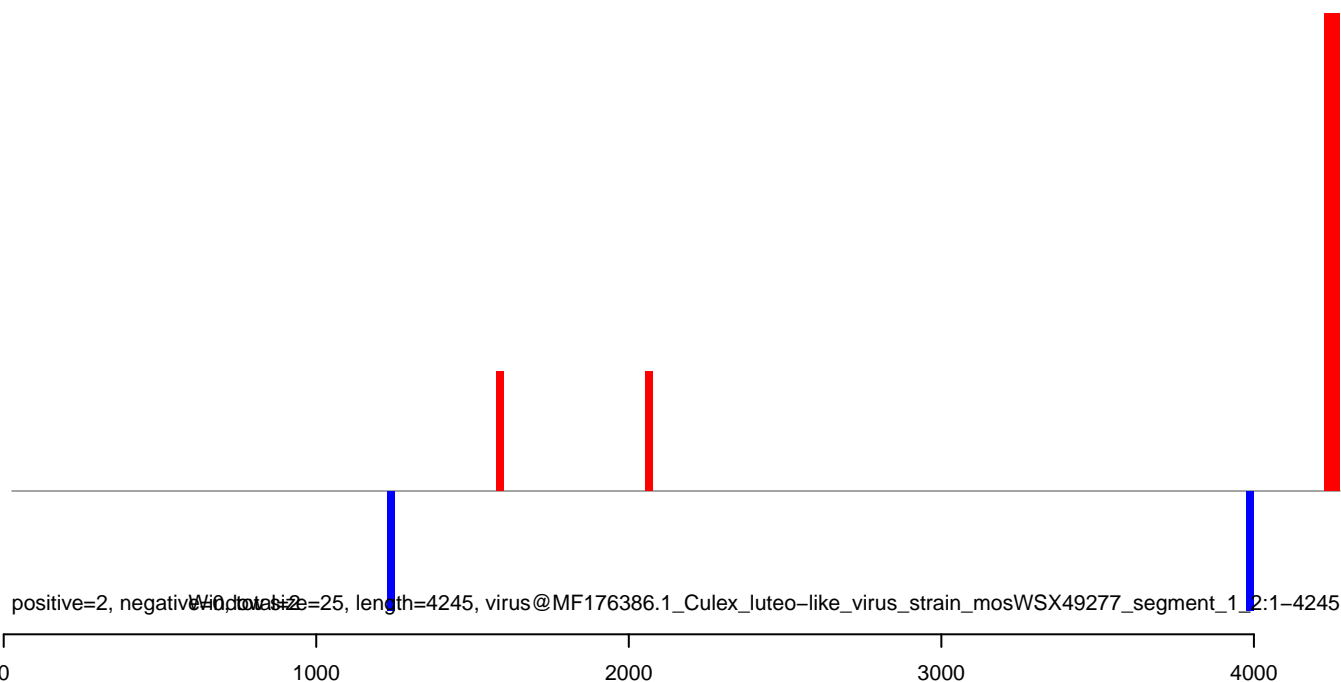
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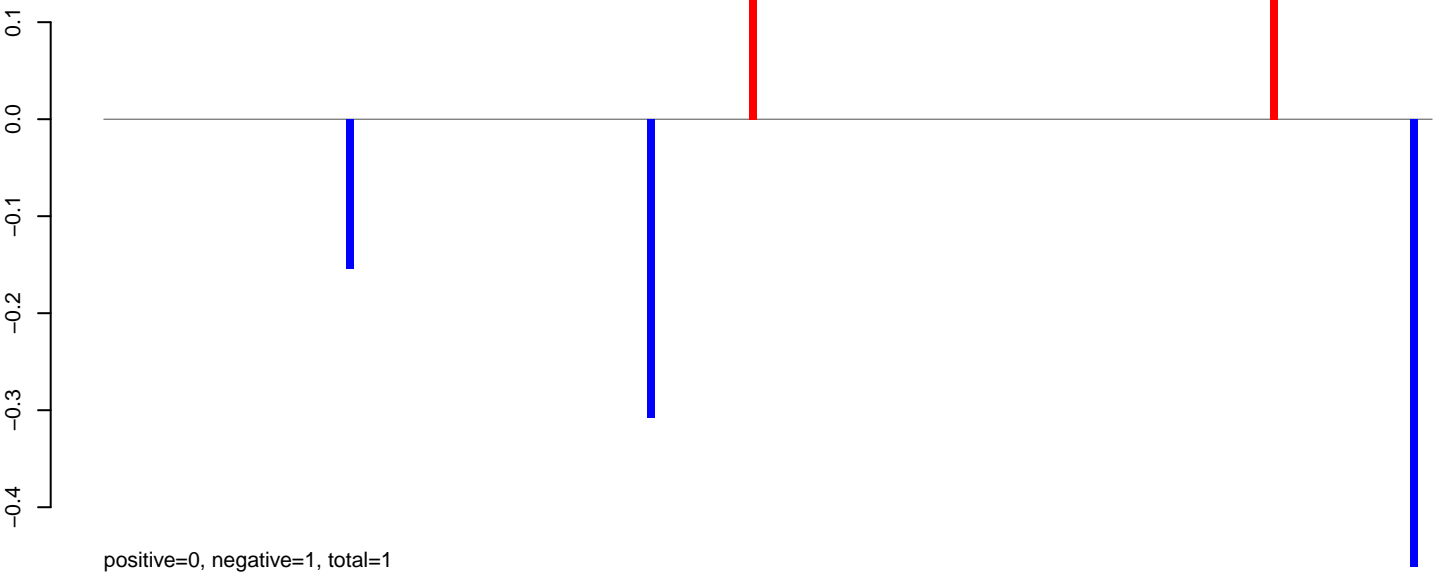
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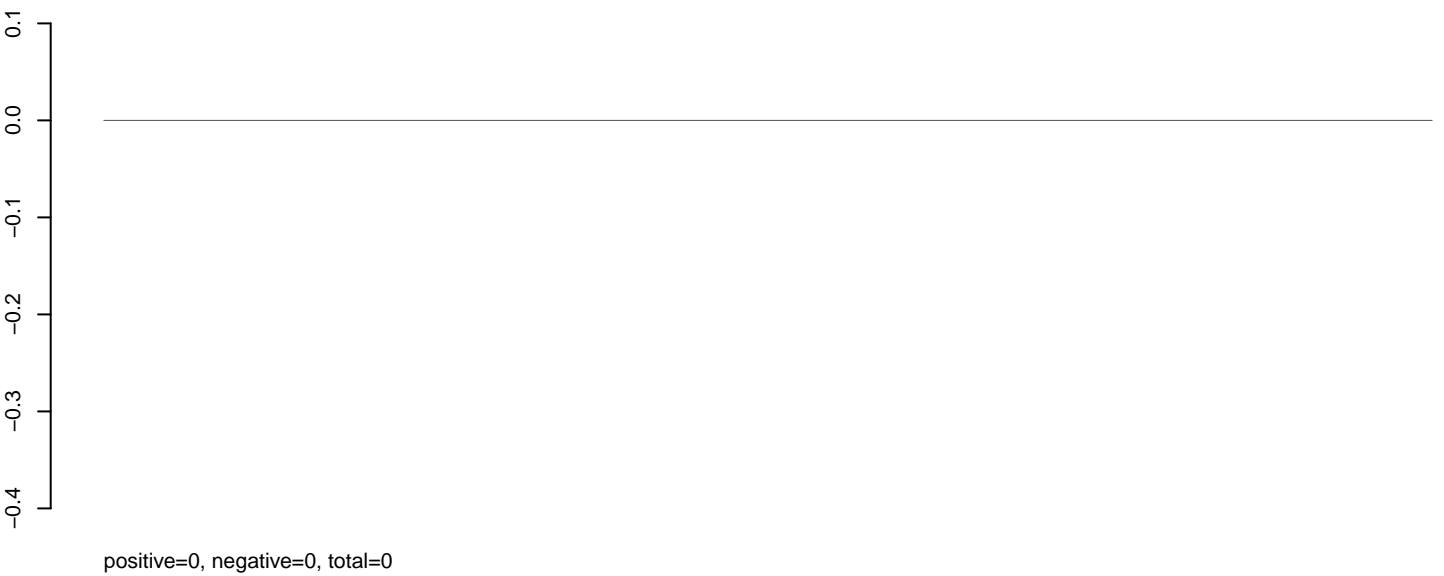
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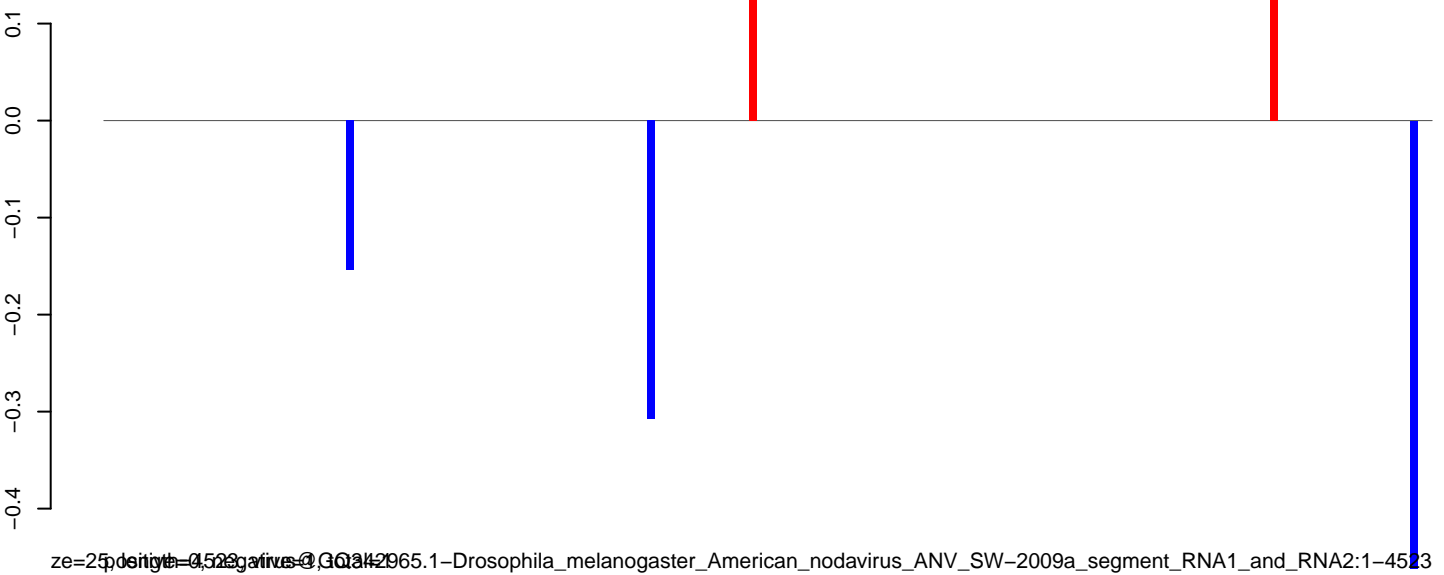
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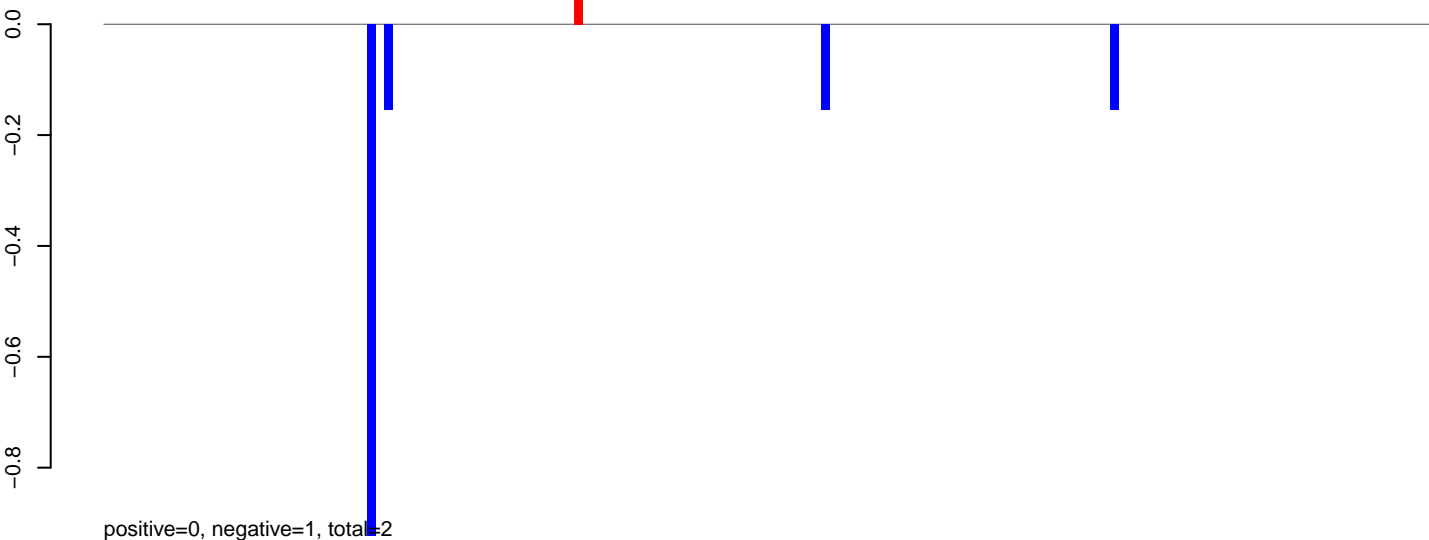
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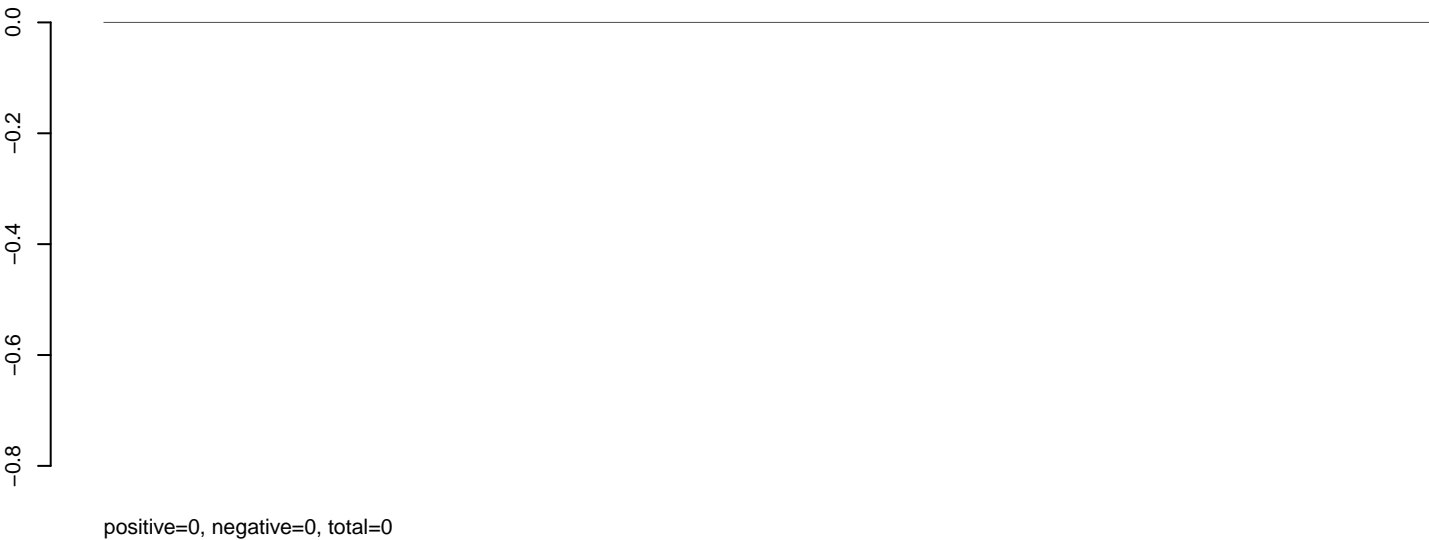
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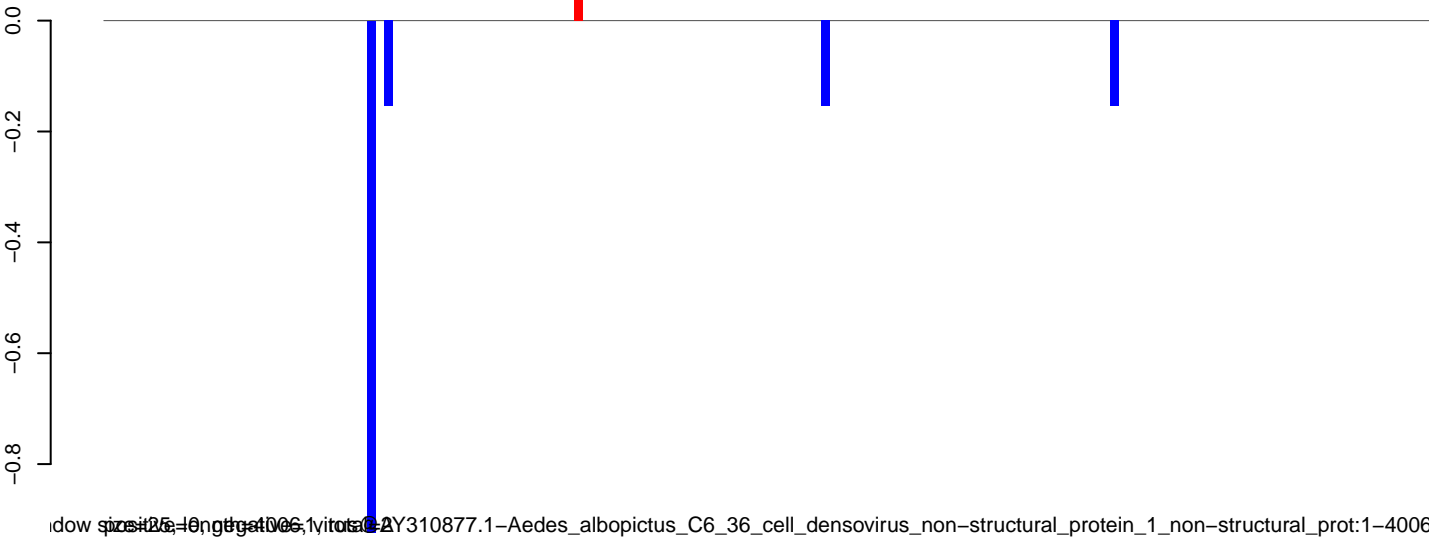
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AeAlbo_C636_6dpi_SINV1.24_35.rep



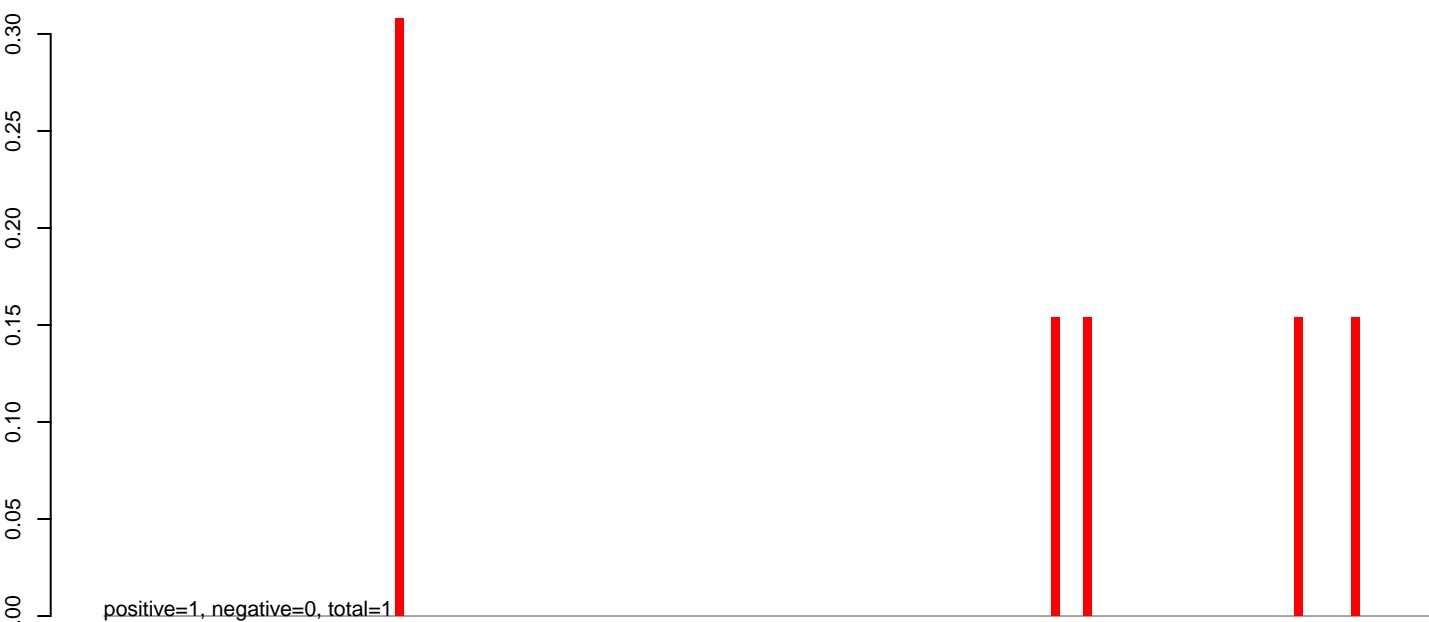
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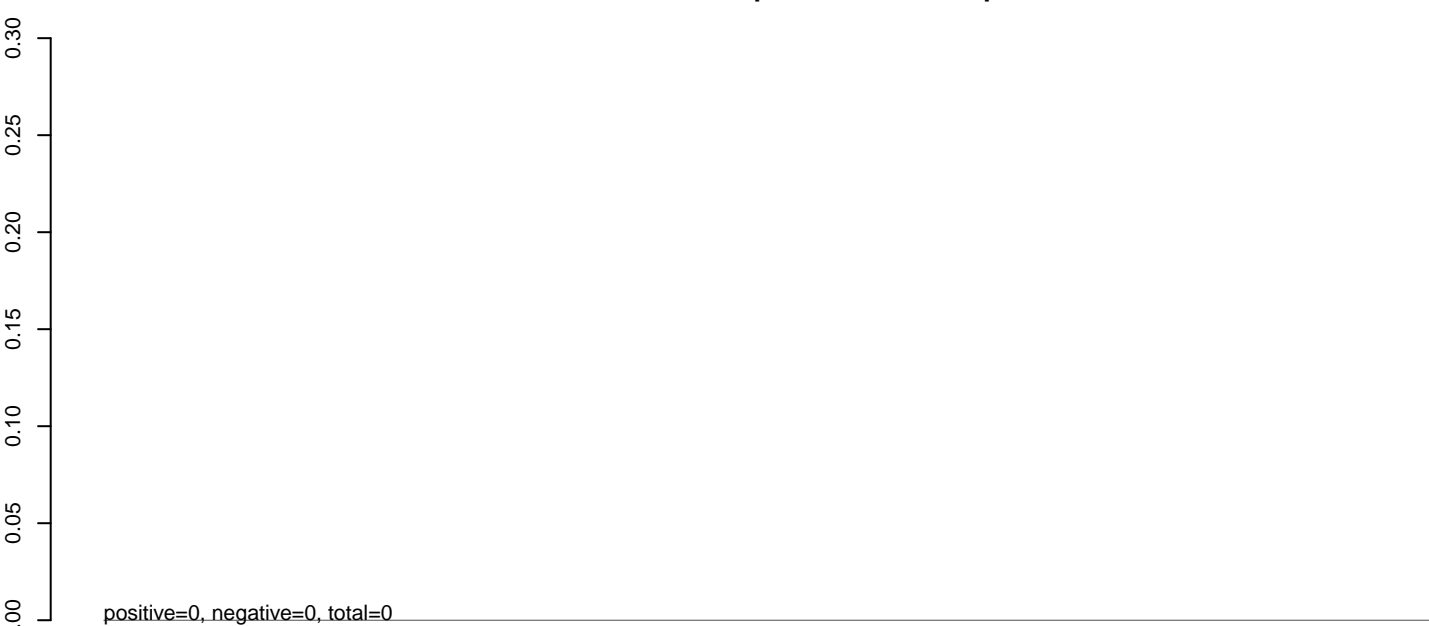
downloaded from <https://www.ncbi.nlm.nih.gov/trace.ncbi.nlm.nih.gov/projects/genome/assembly/blast/> on 2025-10-06, 11:06:21, total=2, AY310877.1-Aedes albopictus_C6_36_cell_densovirus_non-structural_protein_1_non-structural_prot:1-4006

0 1000 2000 3000 4000

AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep

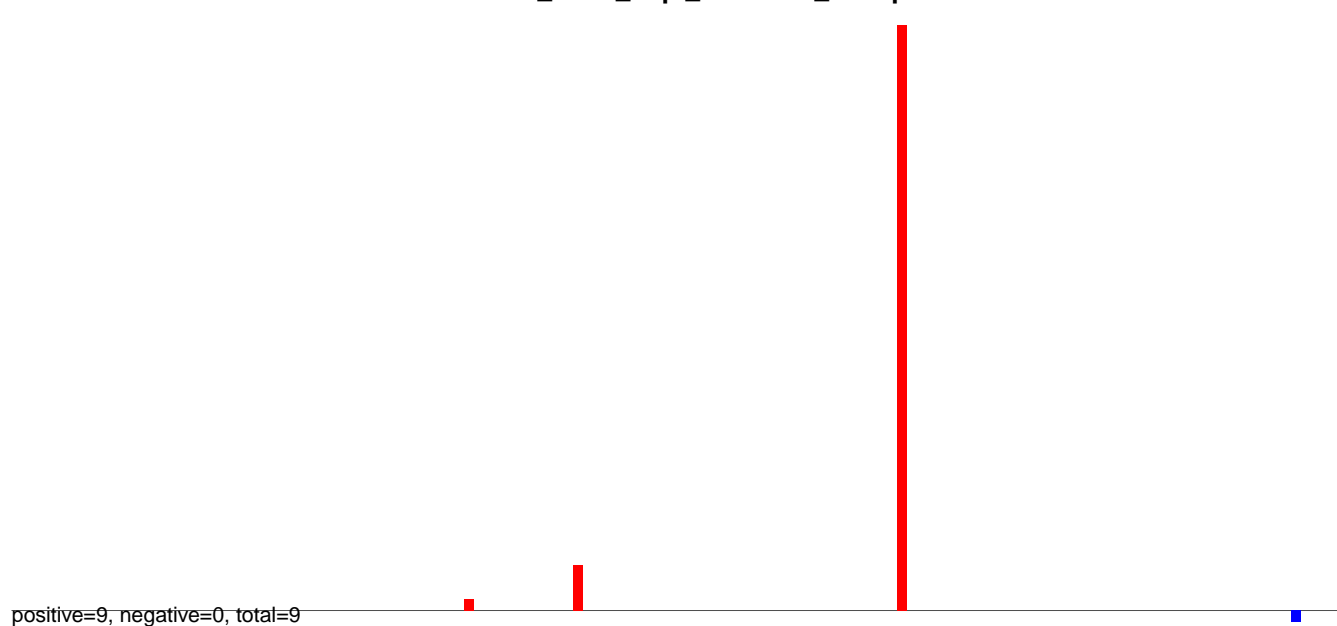


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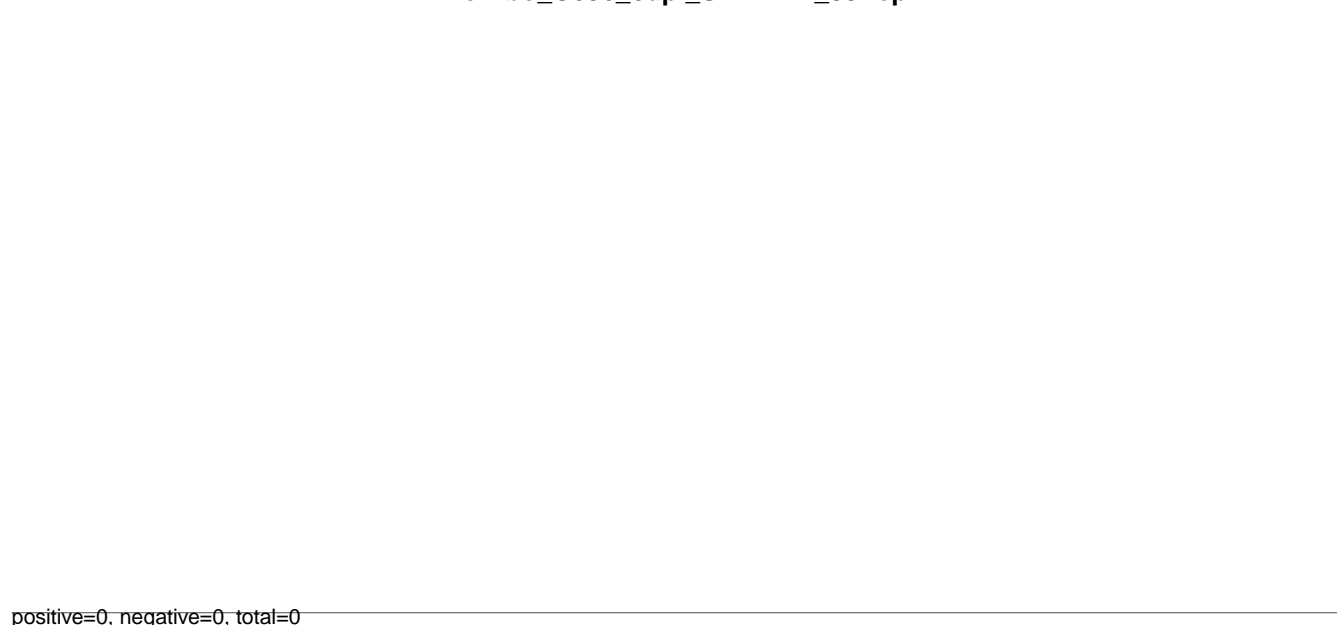


bow size 35, len 4094, the 100% is @A1095351.2-Aedes_albopictus_C6_36_cell_densovirus_non-structural_protein_1_NS-1_non-structural_1-4094

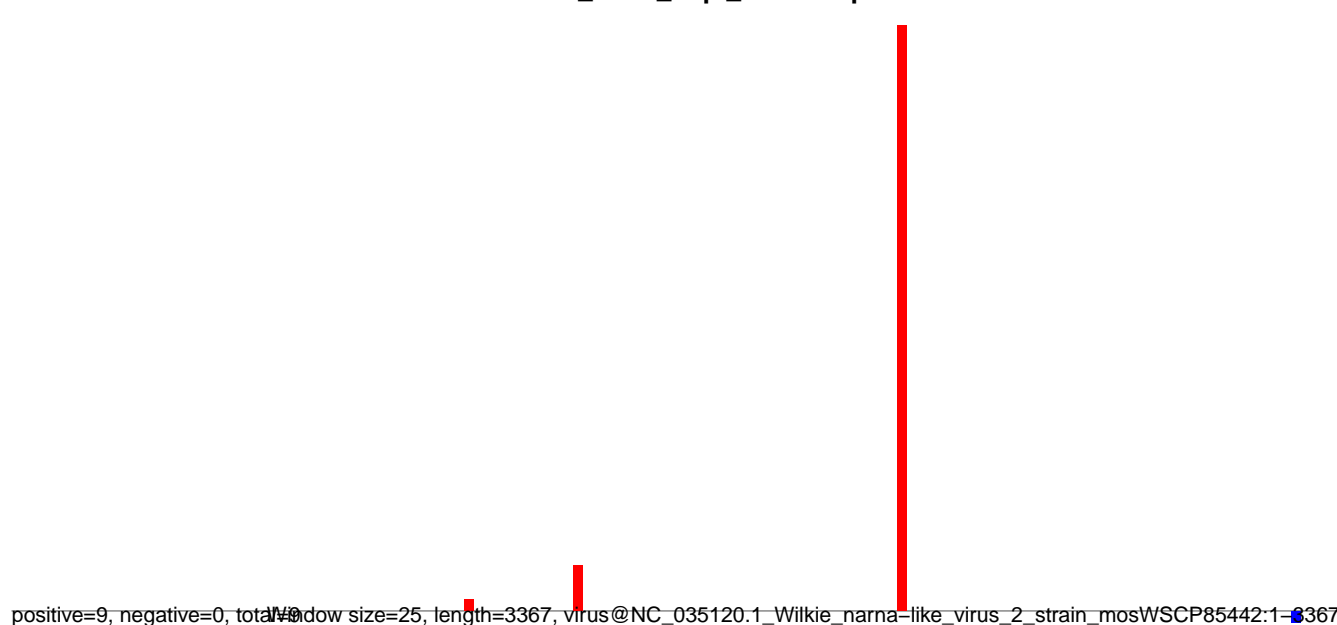
AeAlbo_C636_6dpi_SINV1.18_23.rep



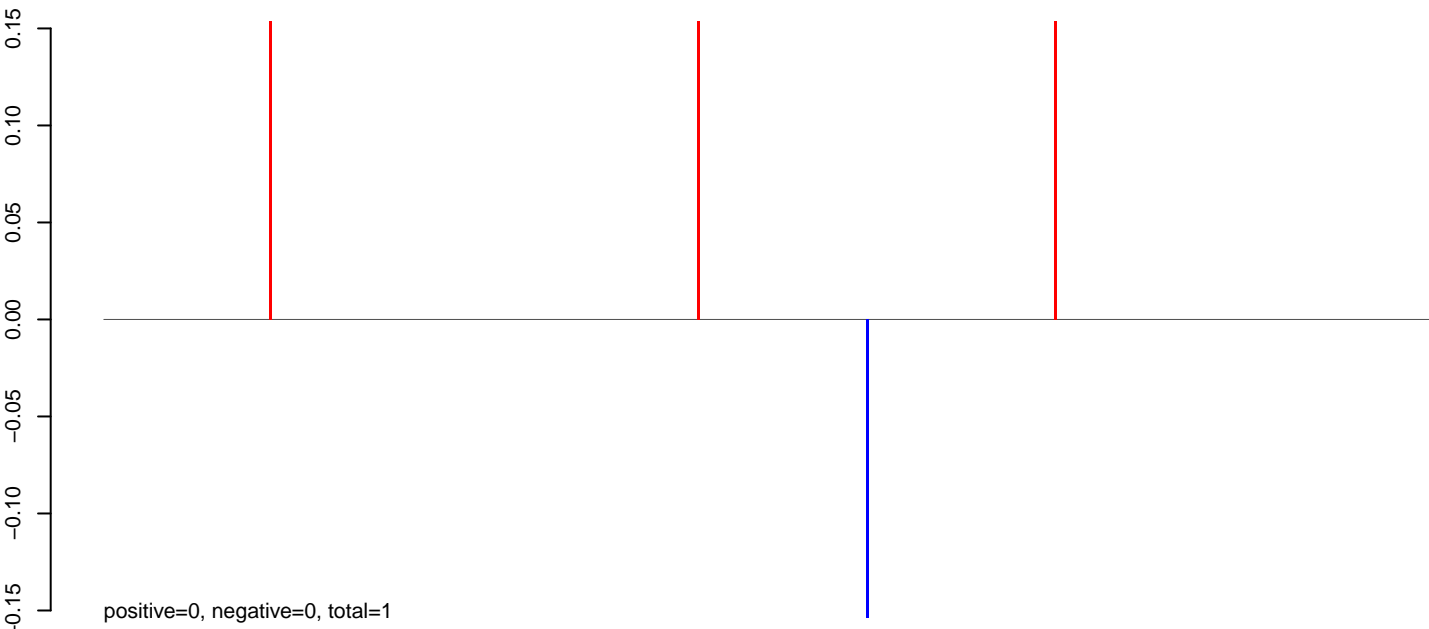
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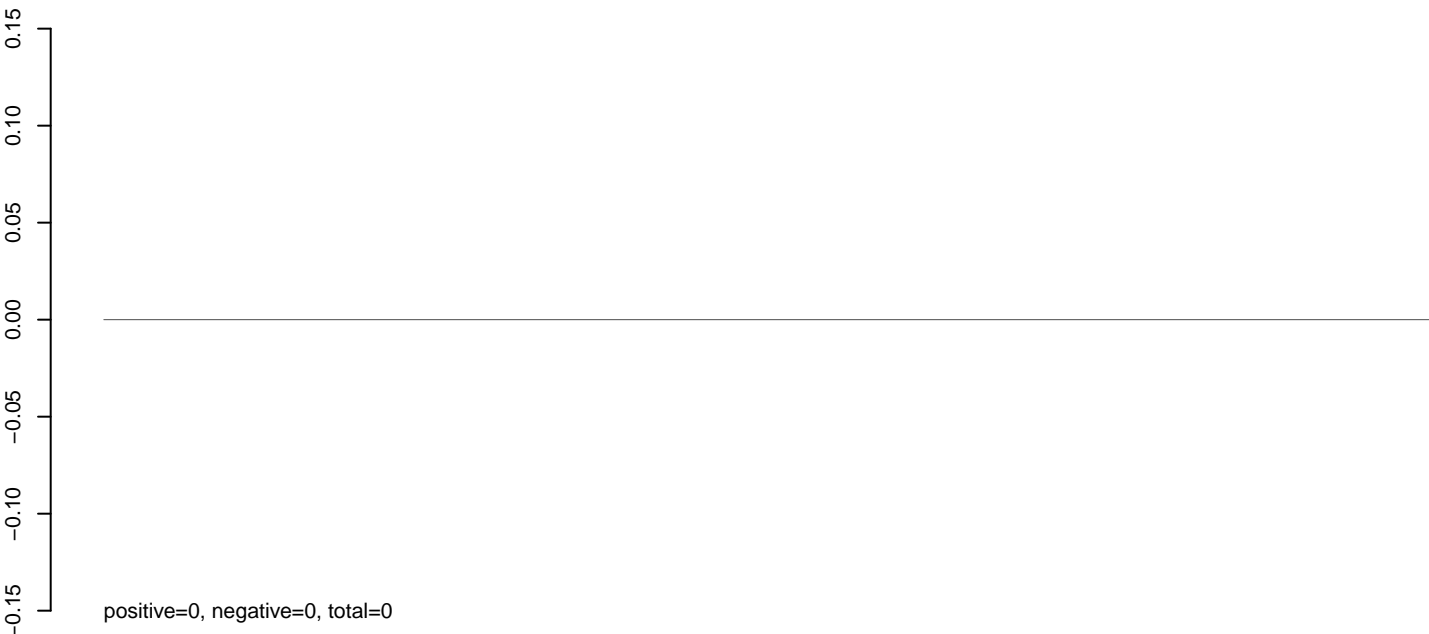
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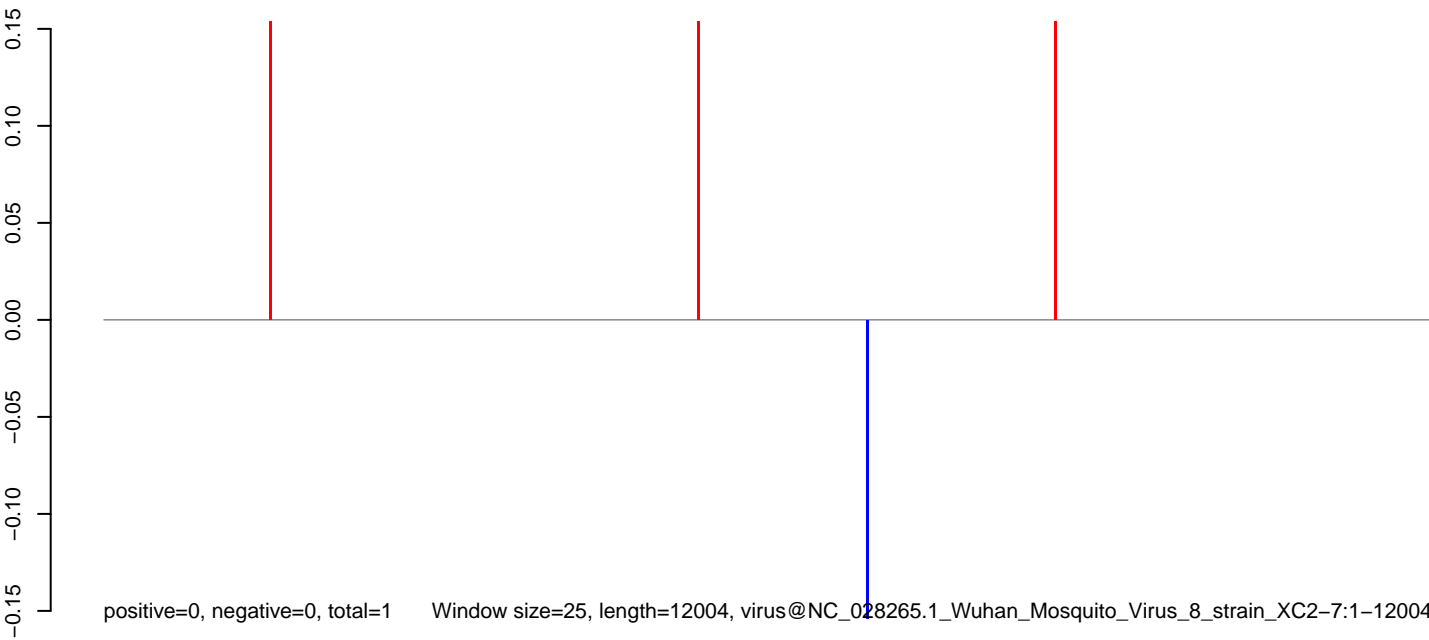
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AeAlbo_C636_6dpi_SINV1.24_35.rep



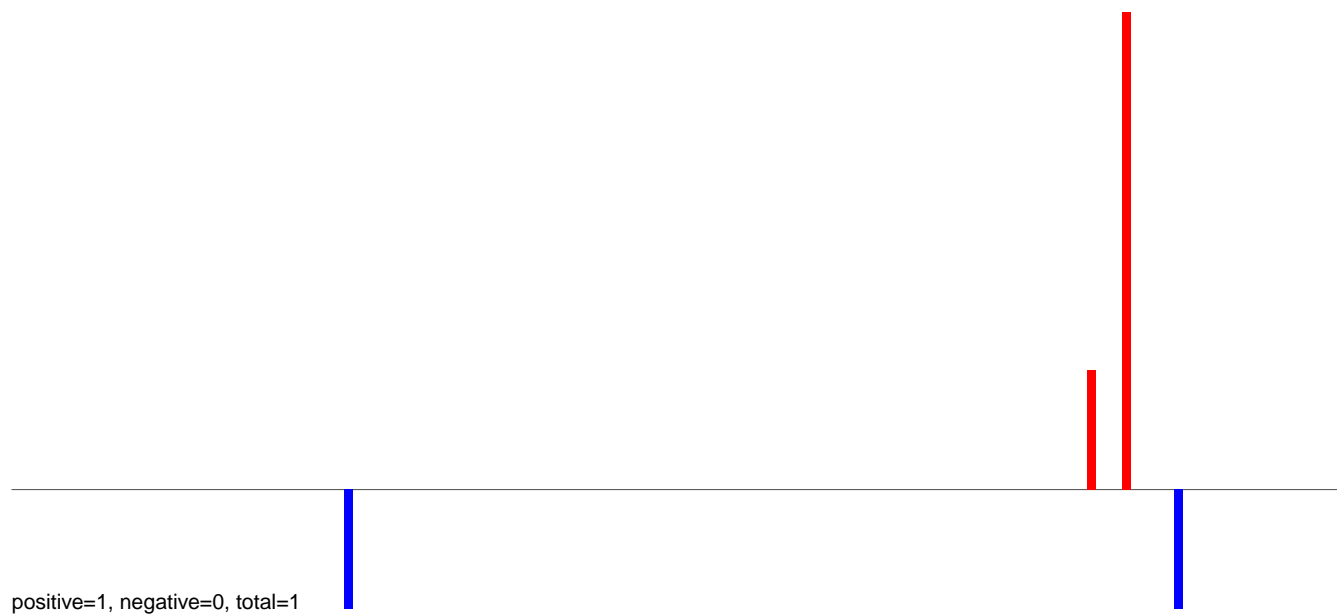
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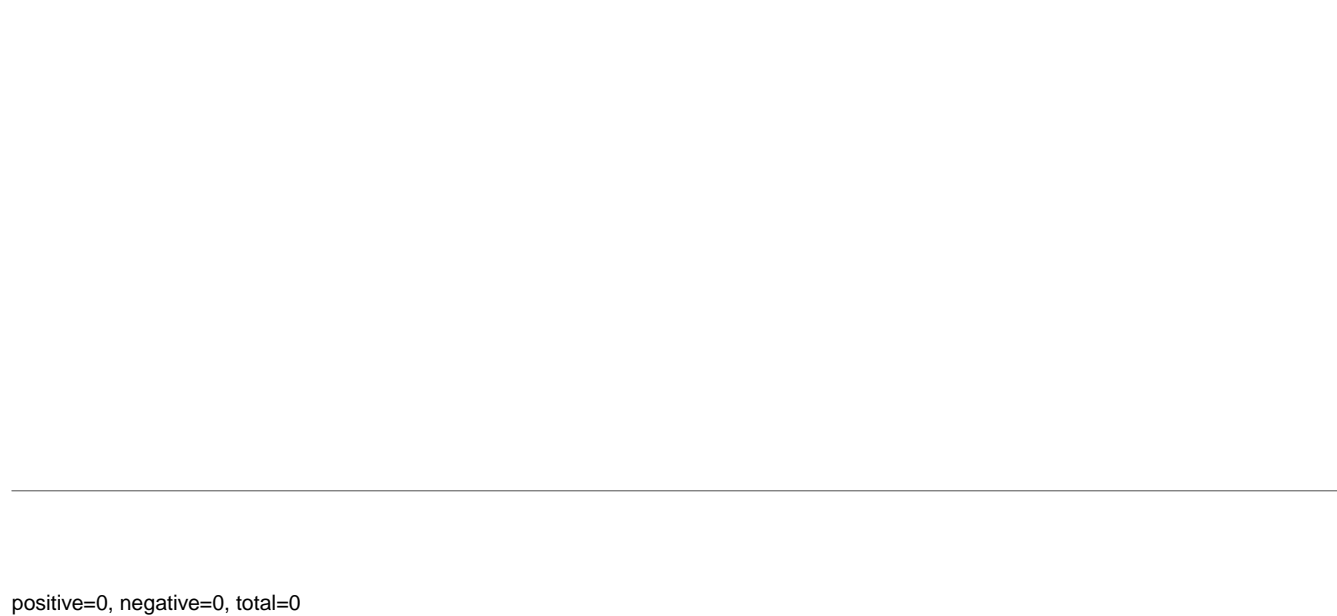
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0 2000 4000 6000 8000 10000 12000

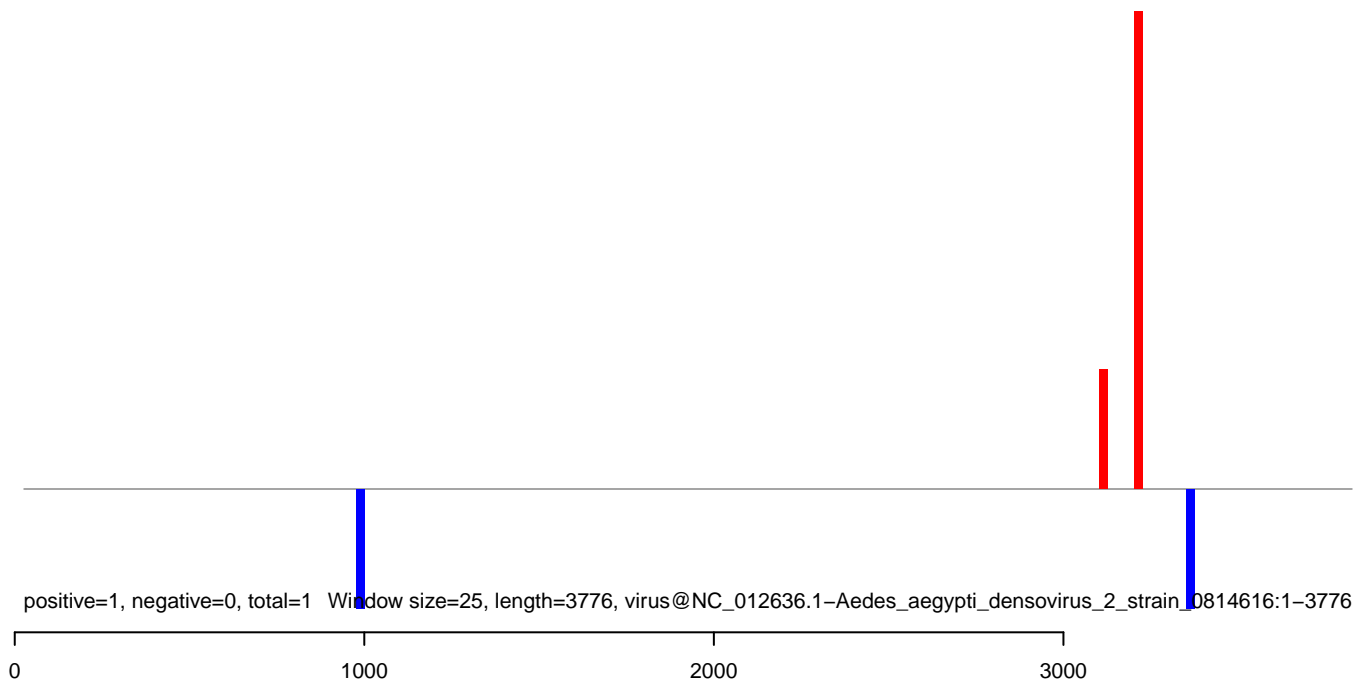
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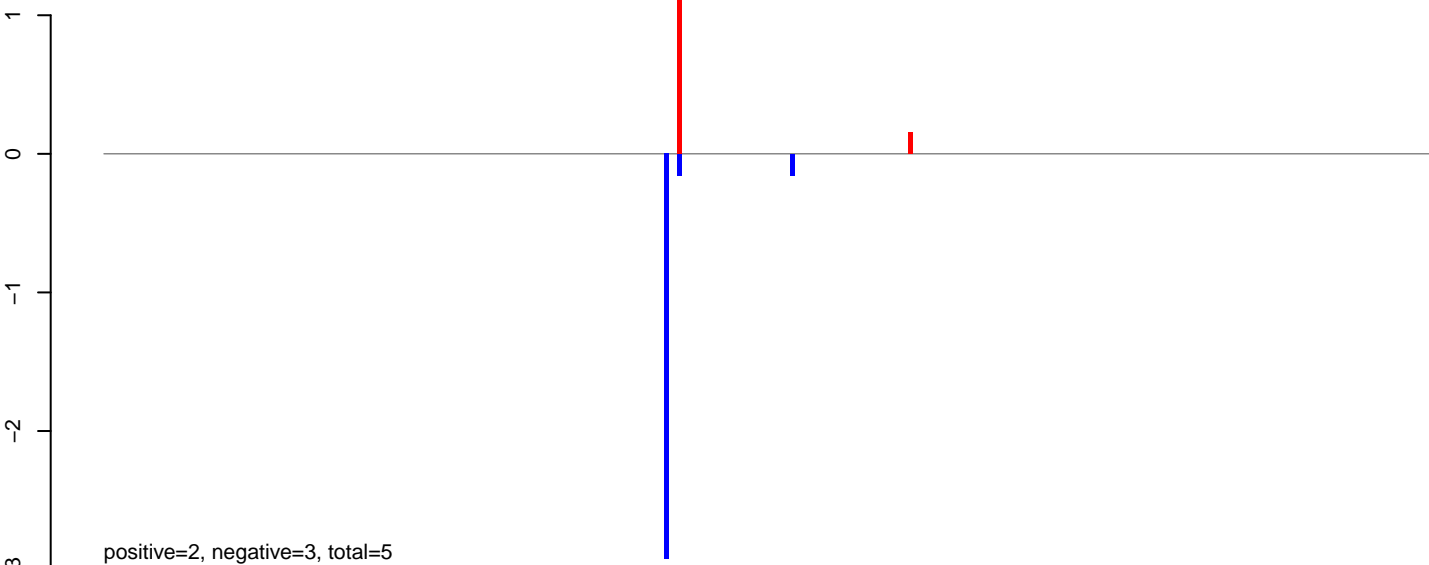
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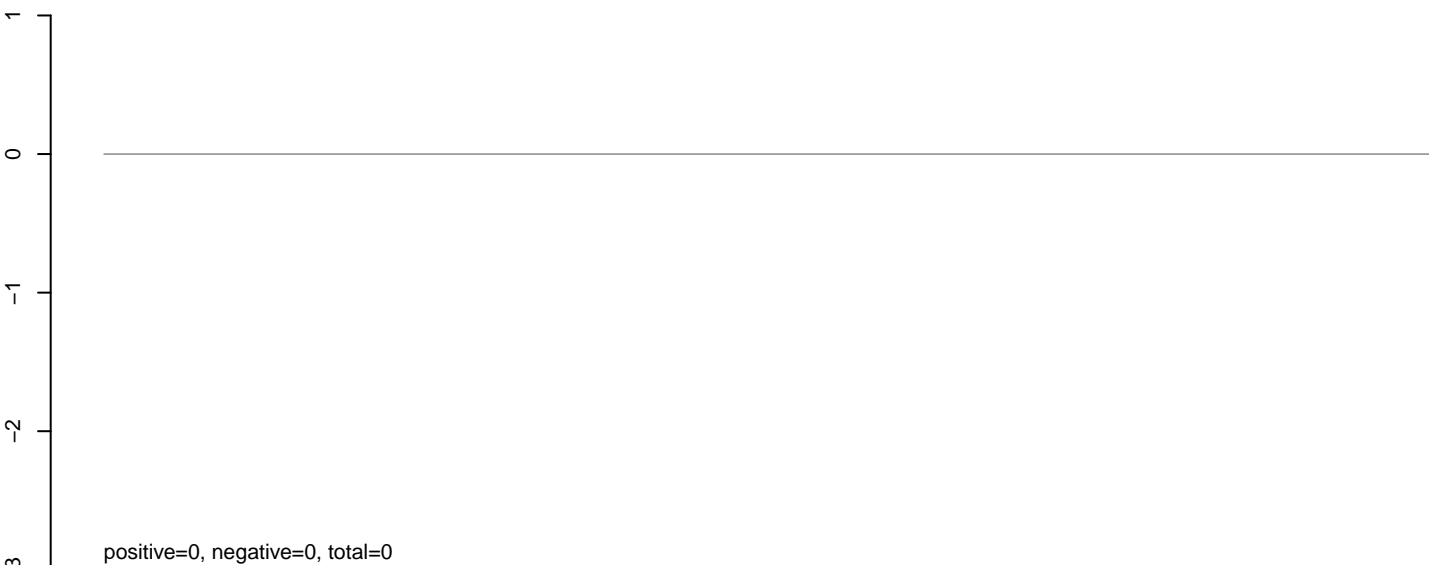
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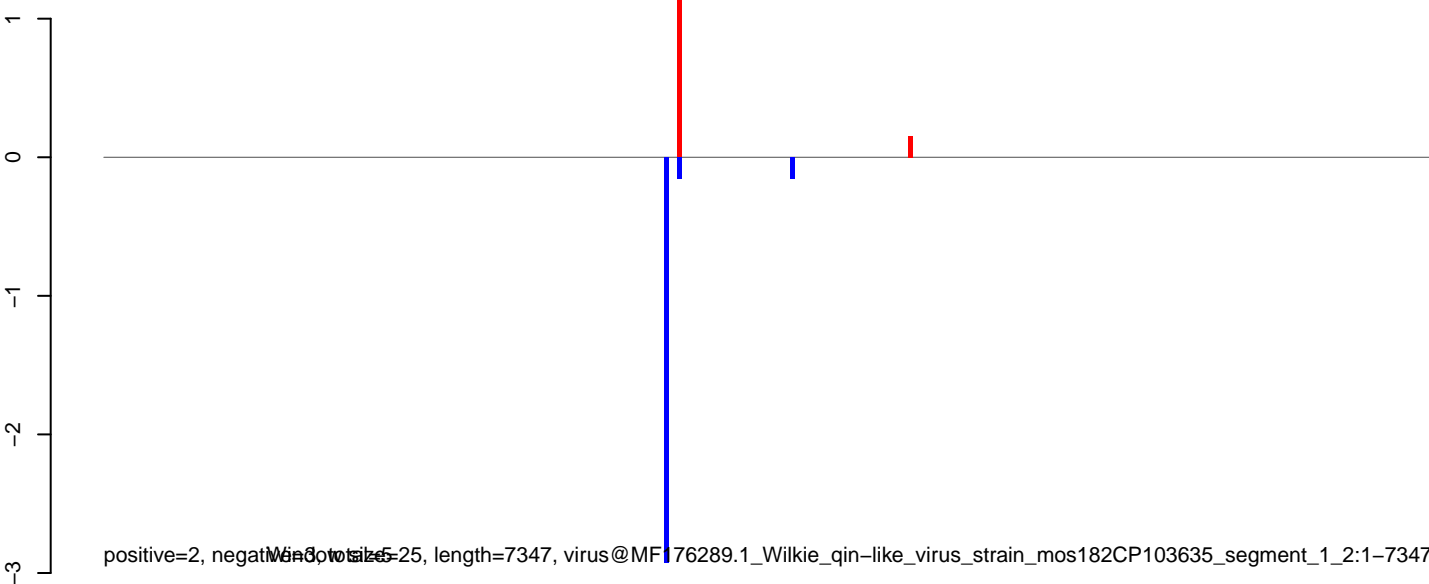
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AeAlbo_C636_6dpi_SINV1.24_35.rep



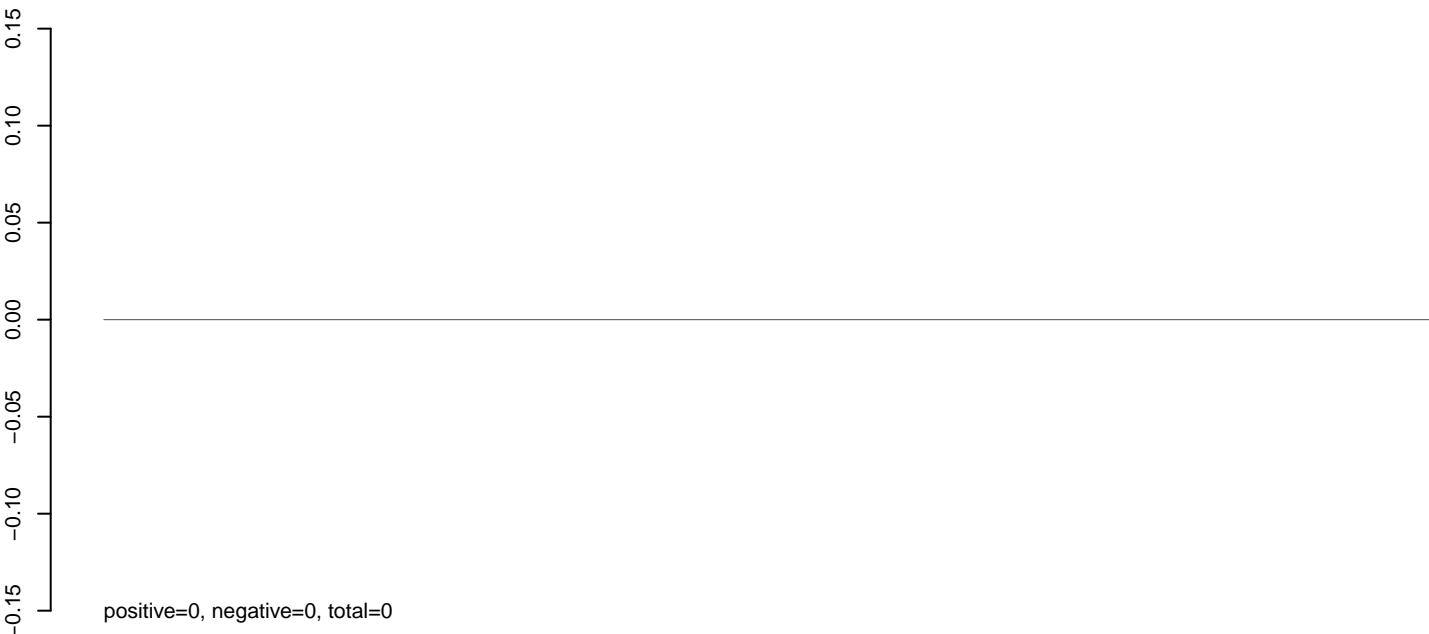
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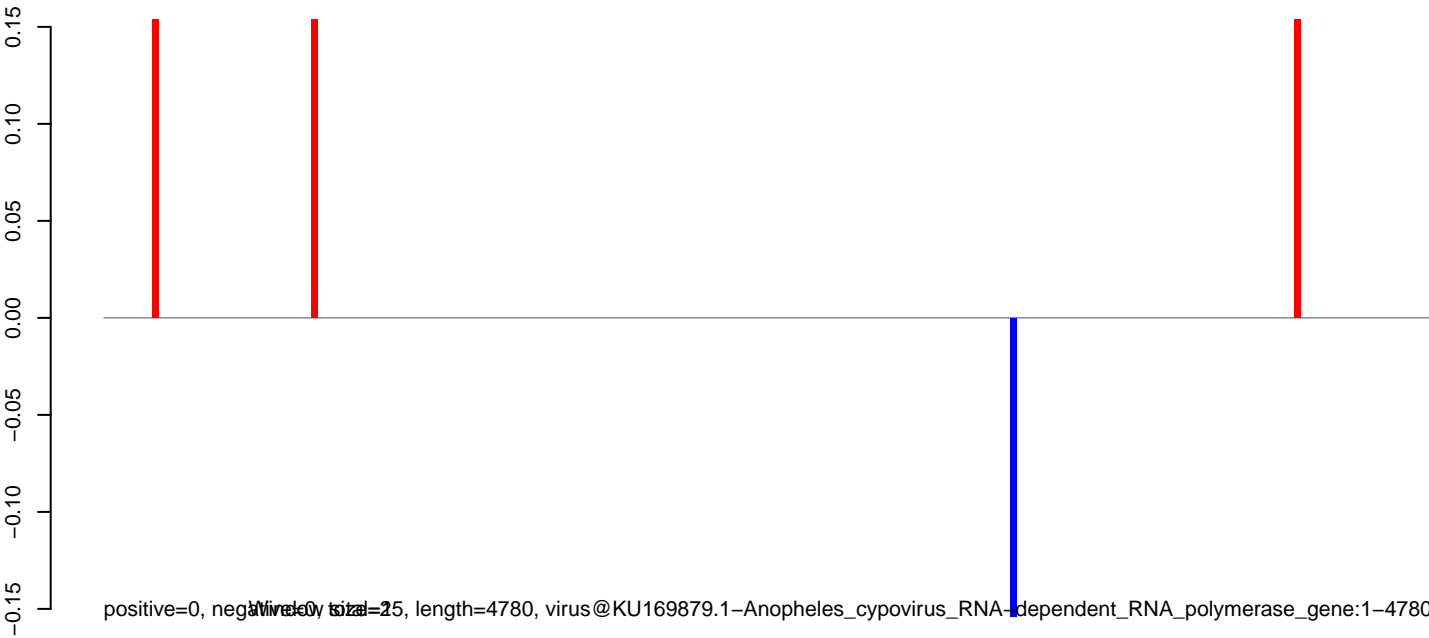
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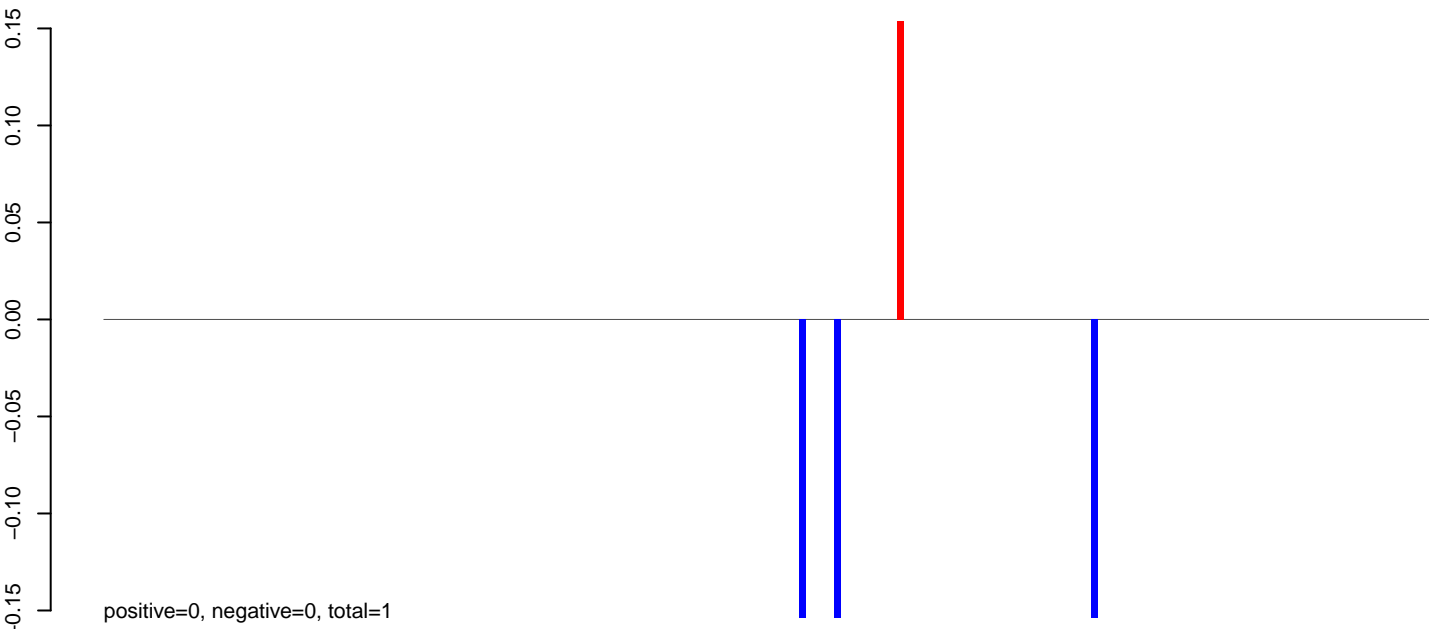
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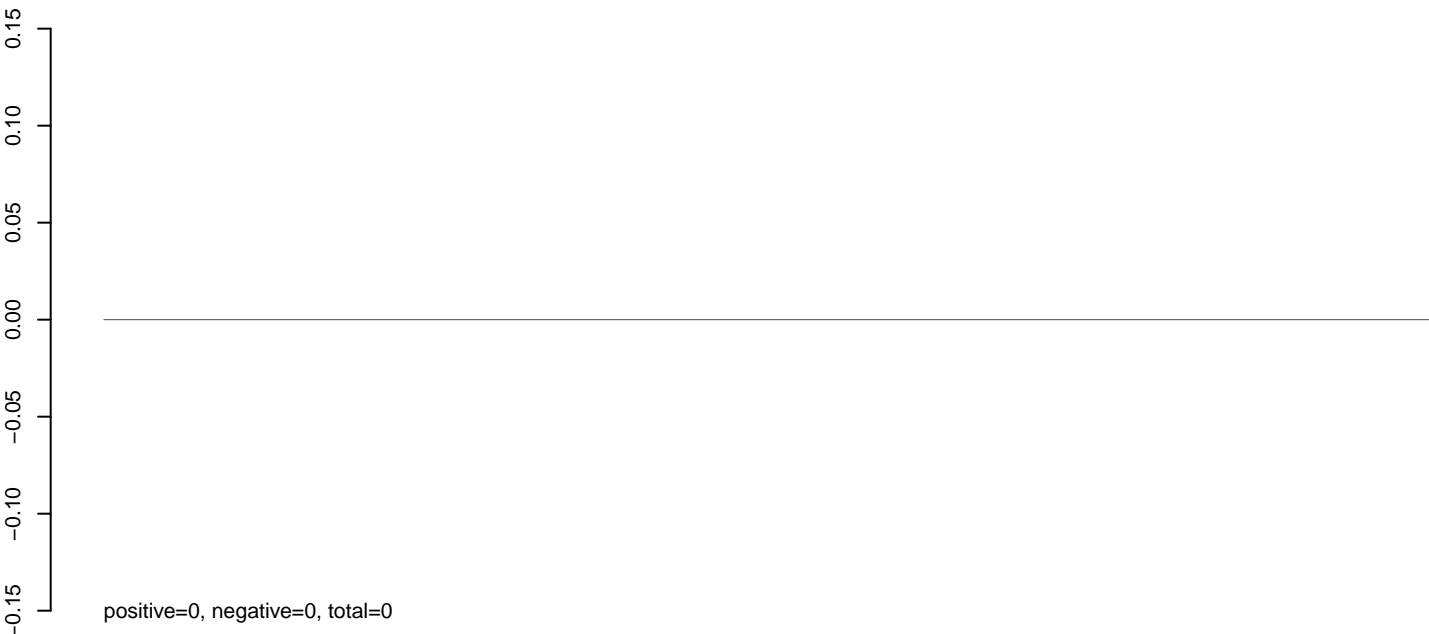
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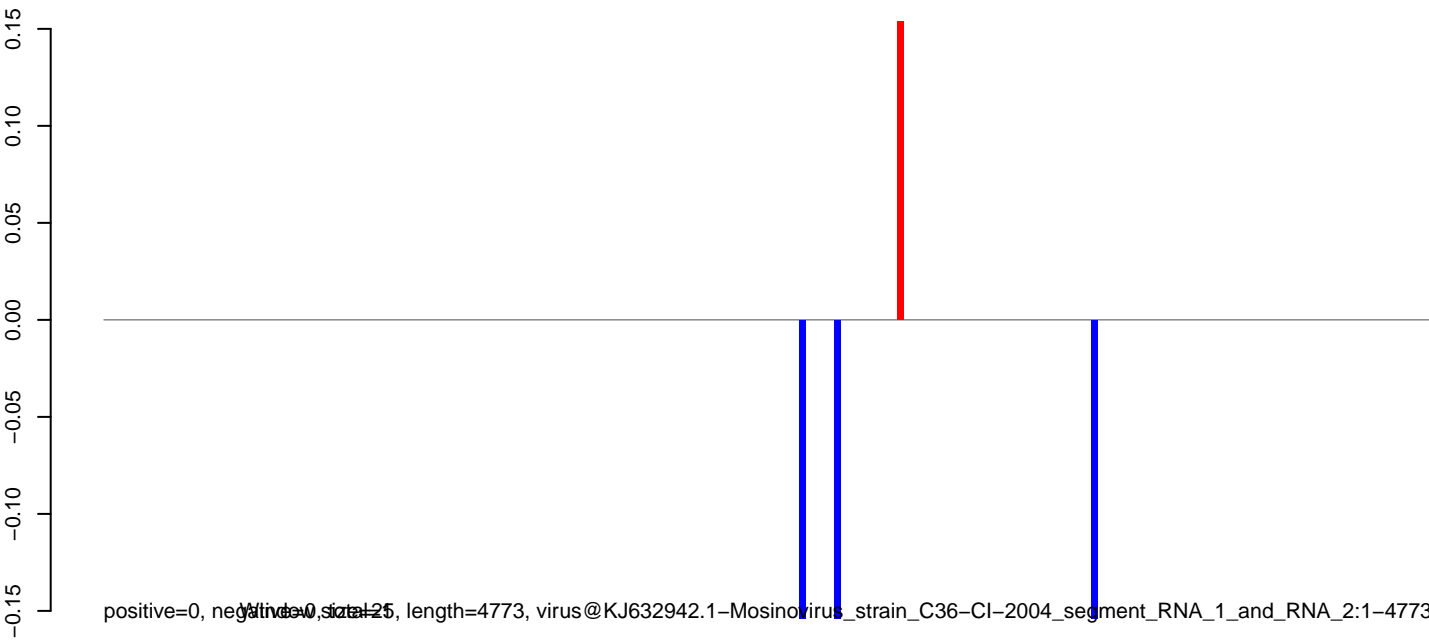
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



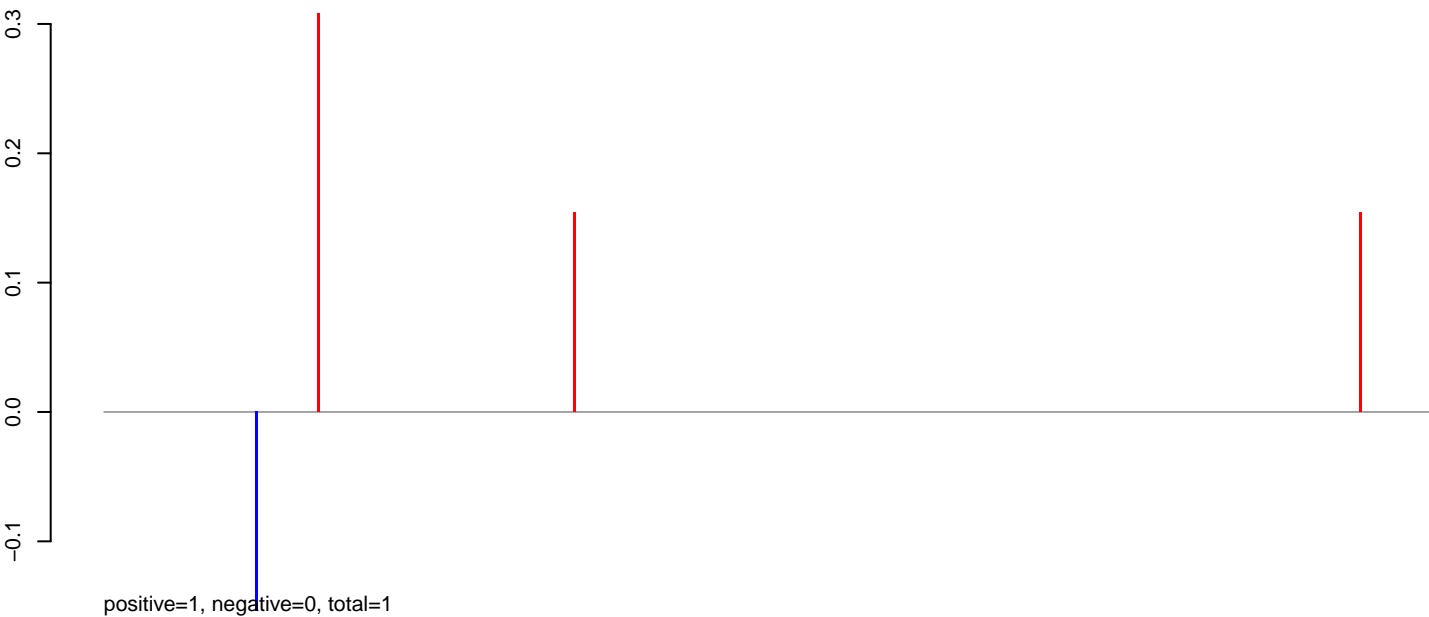
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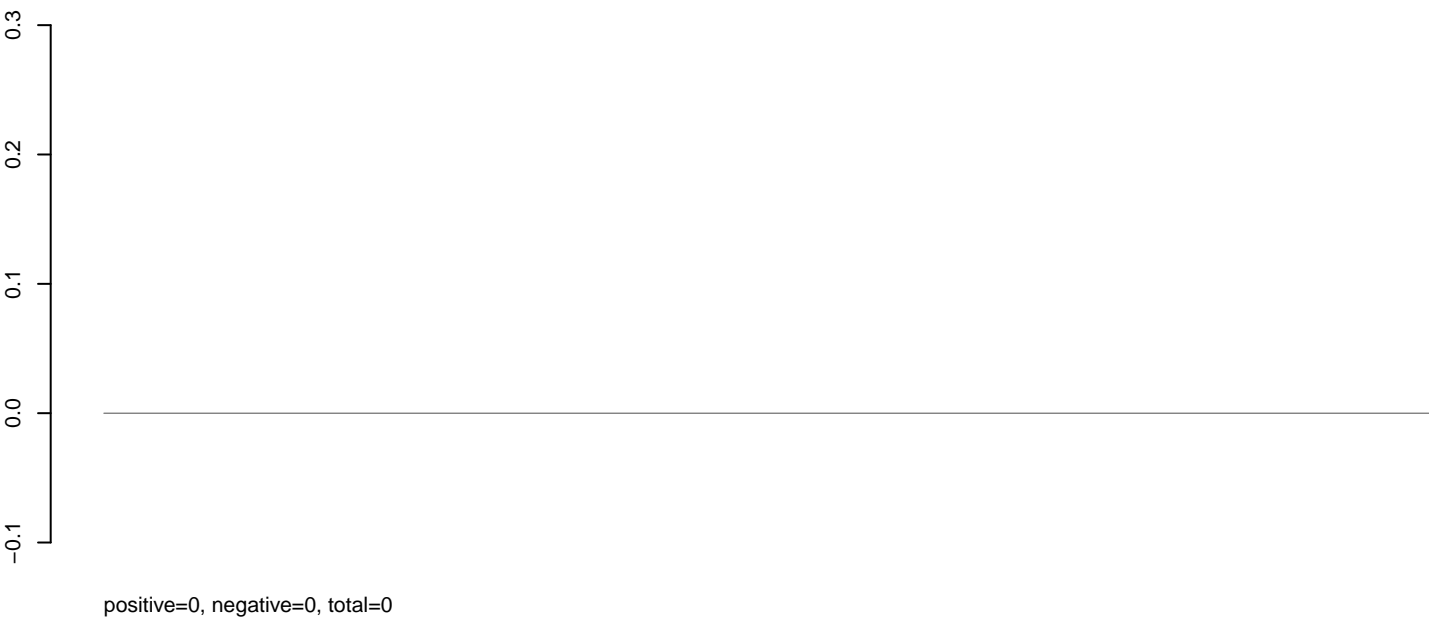
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0 1000 2000 3000 4000

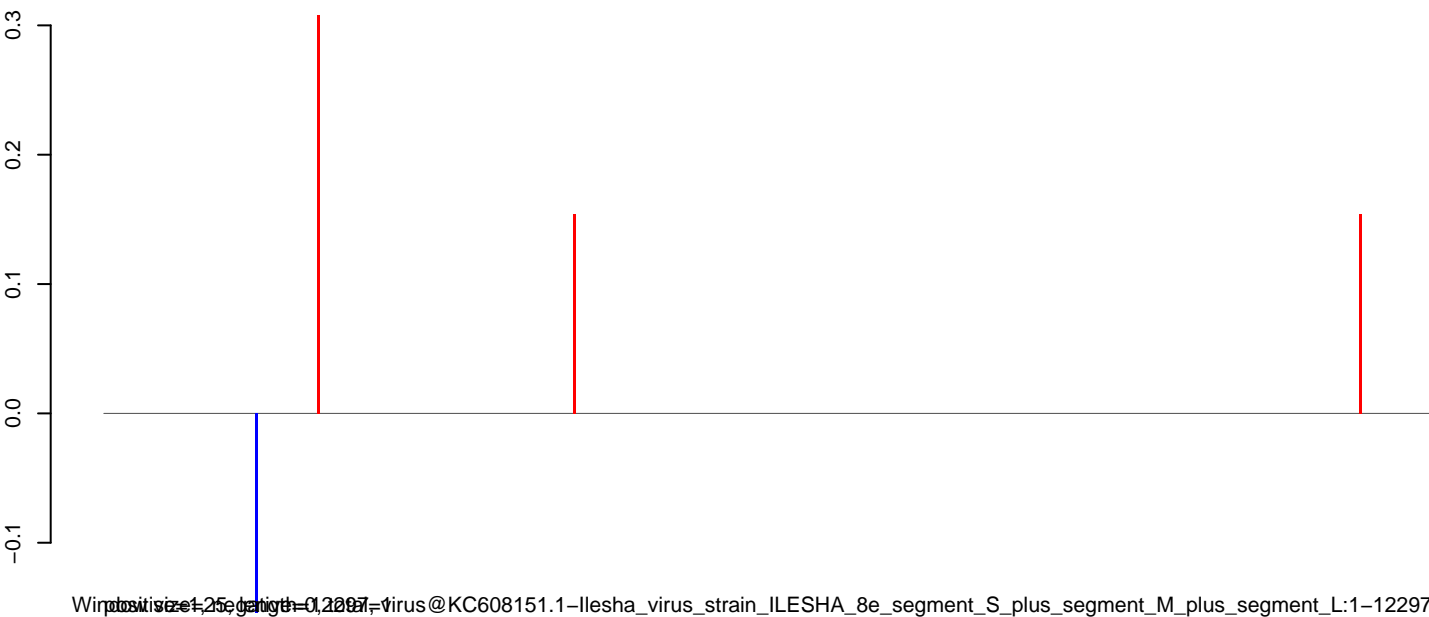
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



AeAlbo_C636_6dpi_SINV1.rep

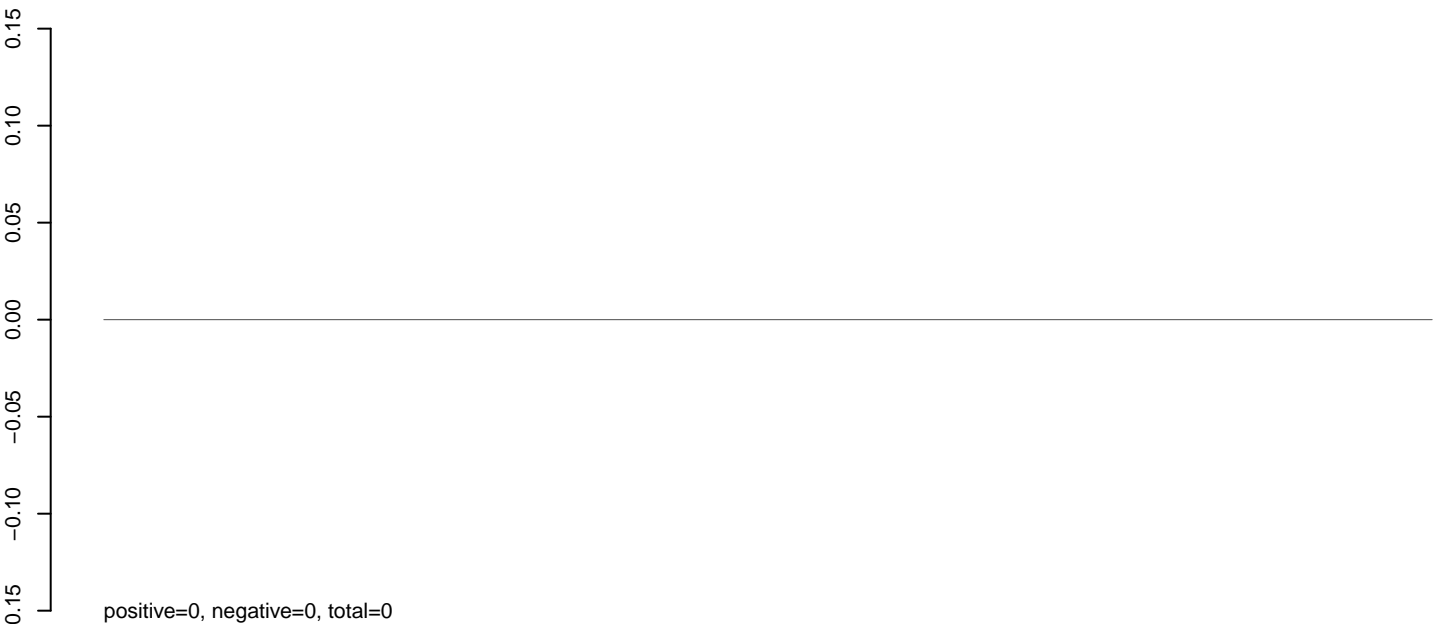


AeAlbo_C636_6dpi_SINV1.18_23.rep



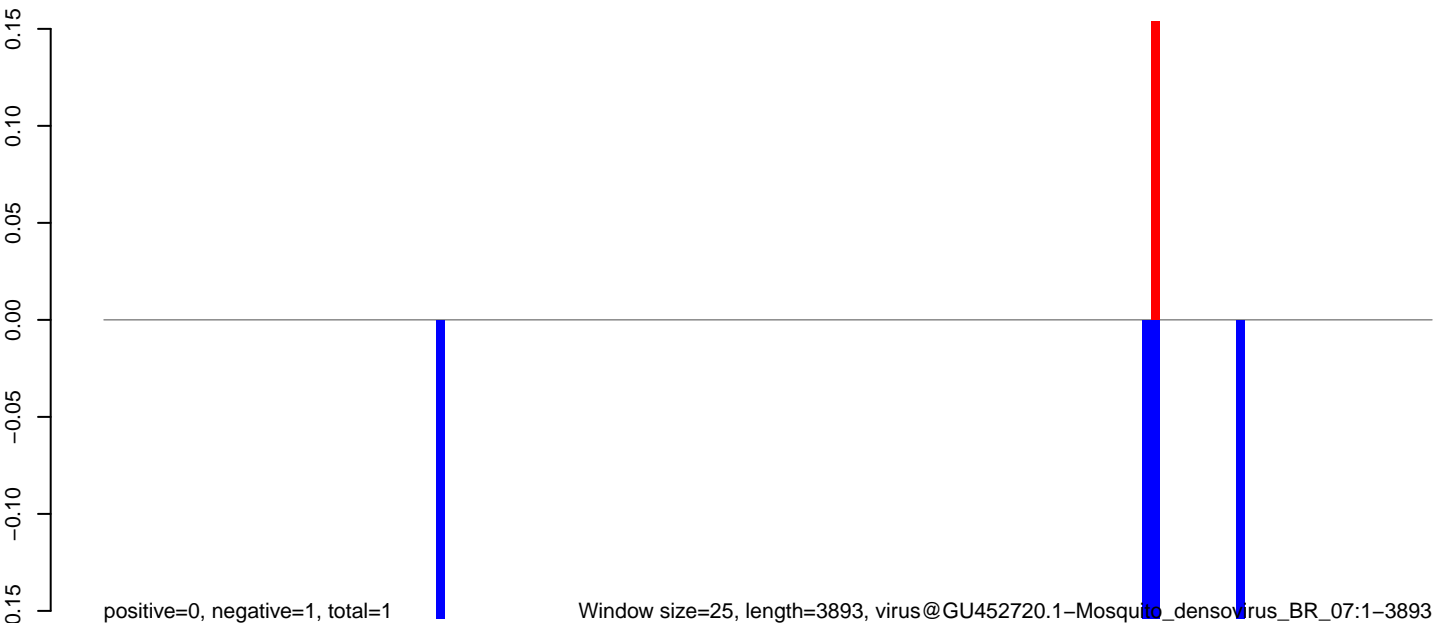
positive=0, negative=1, total=1

AeAlbo_C636_6dpi_SINV1.24_35.rep



positive=0, negative=0, total=0

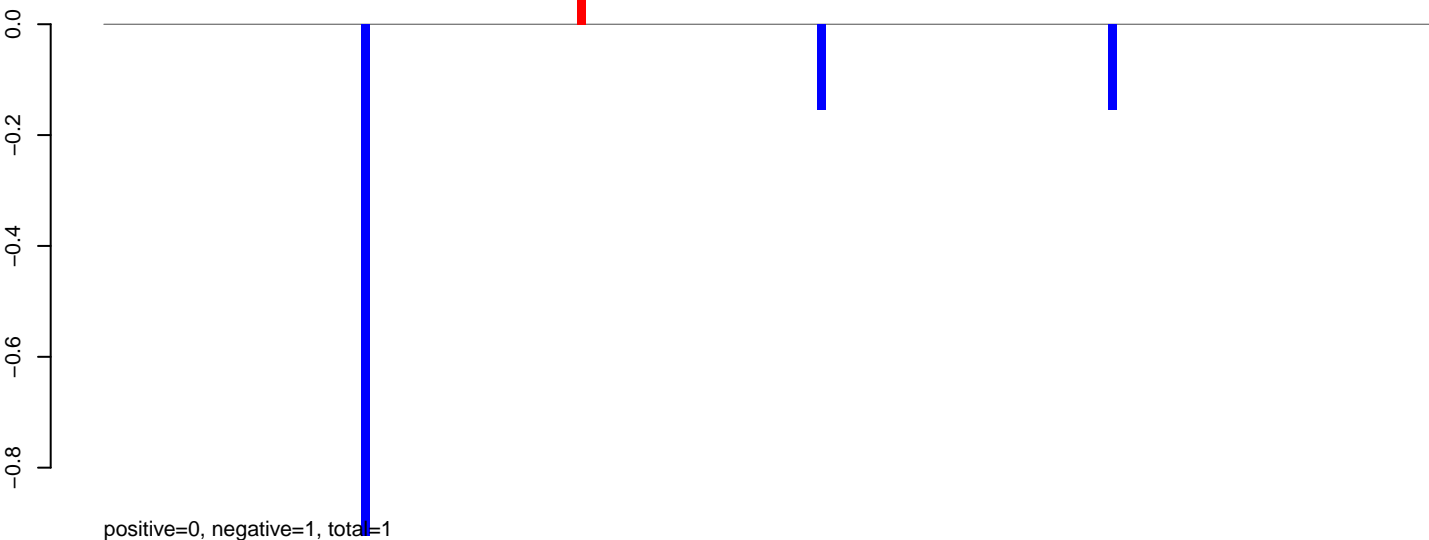
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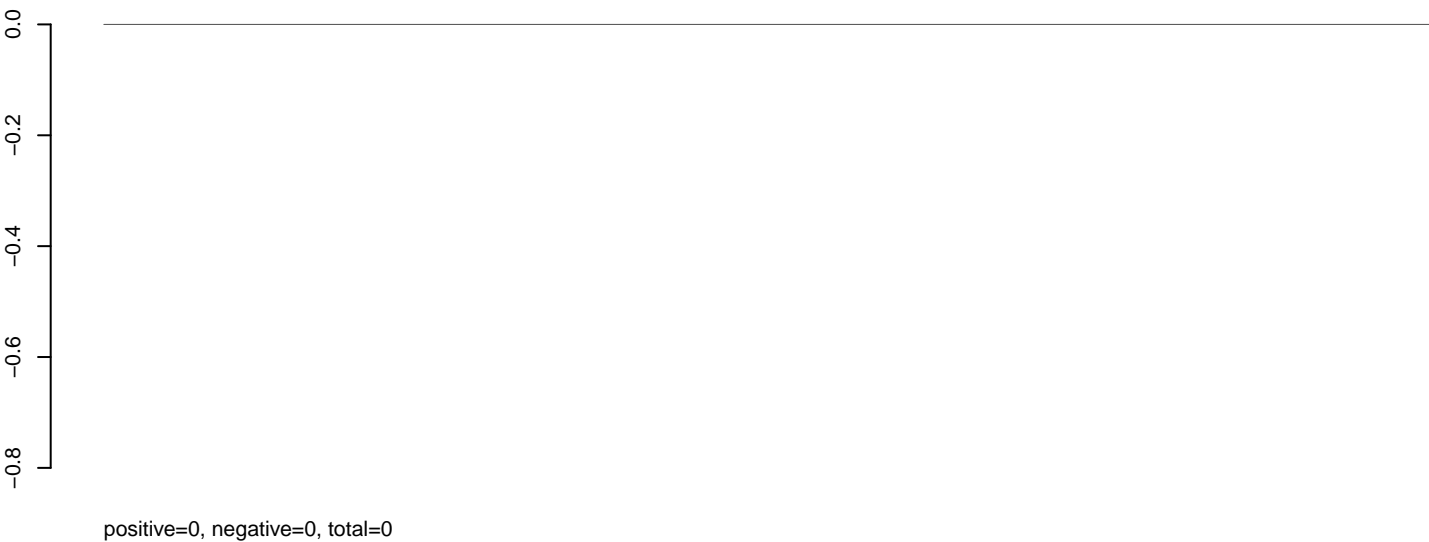
positive=0, negative=1, total=1

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

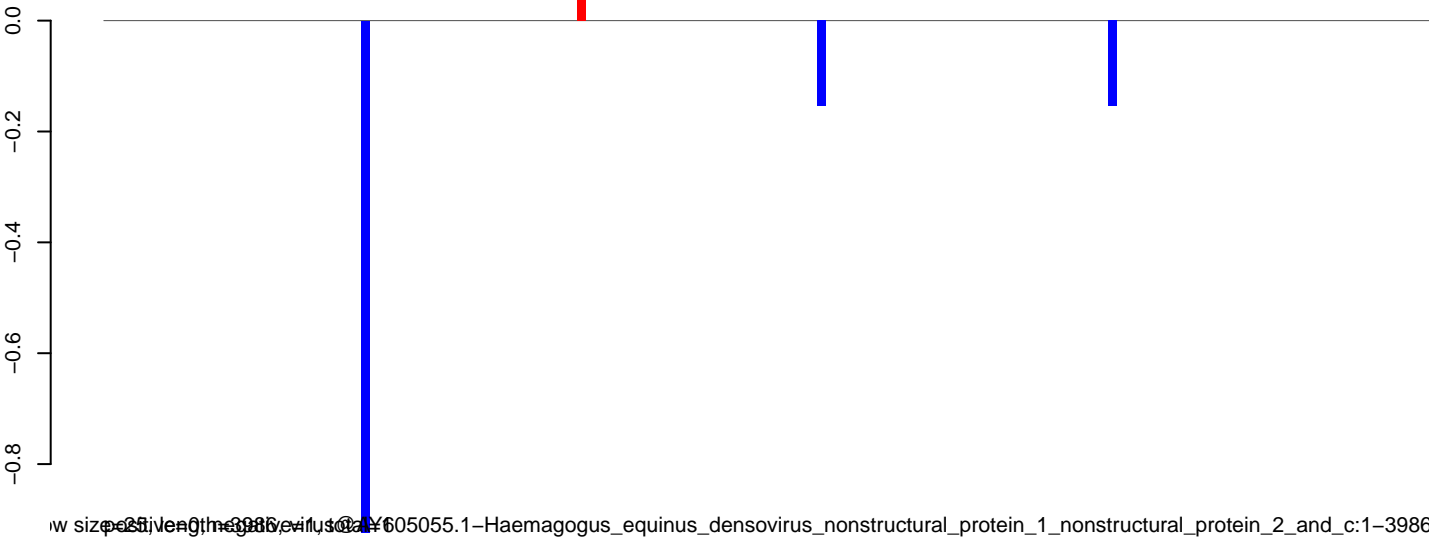
AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



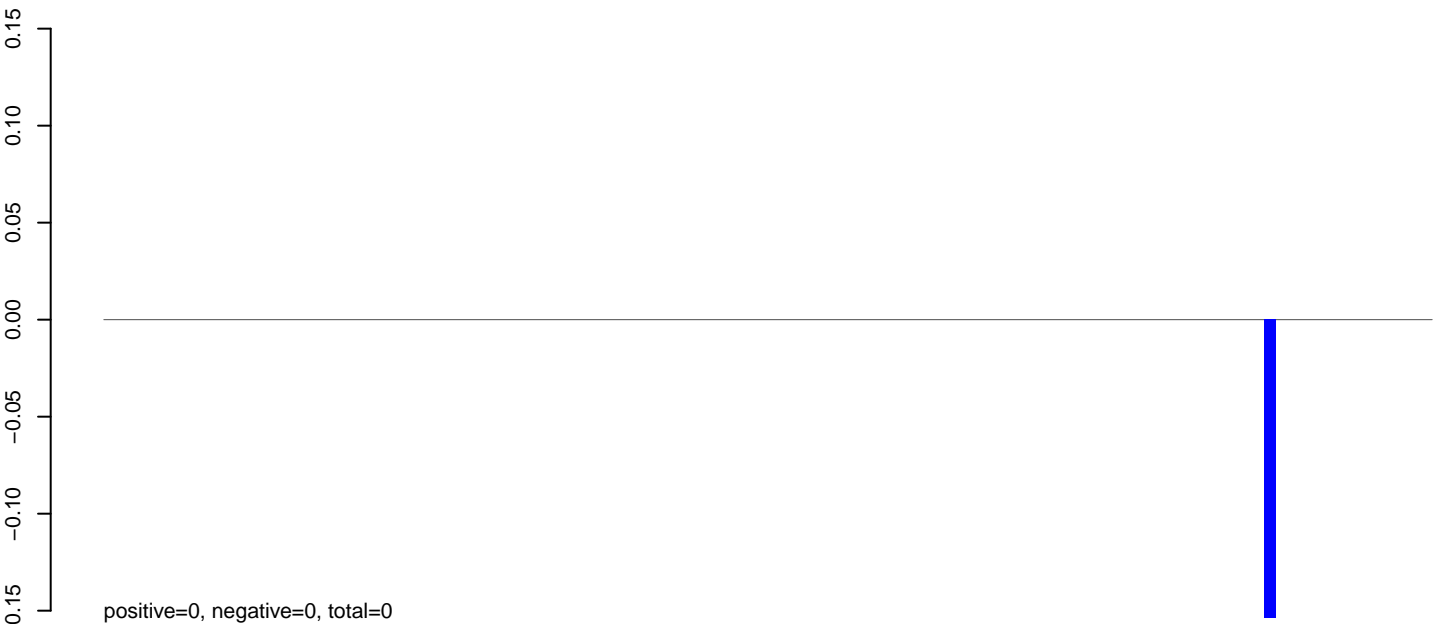
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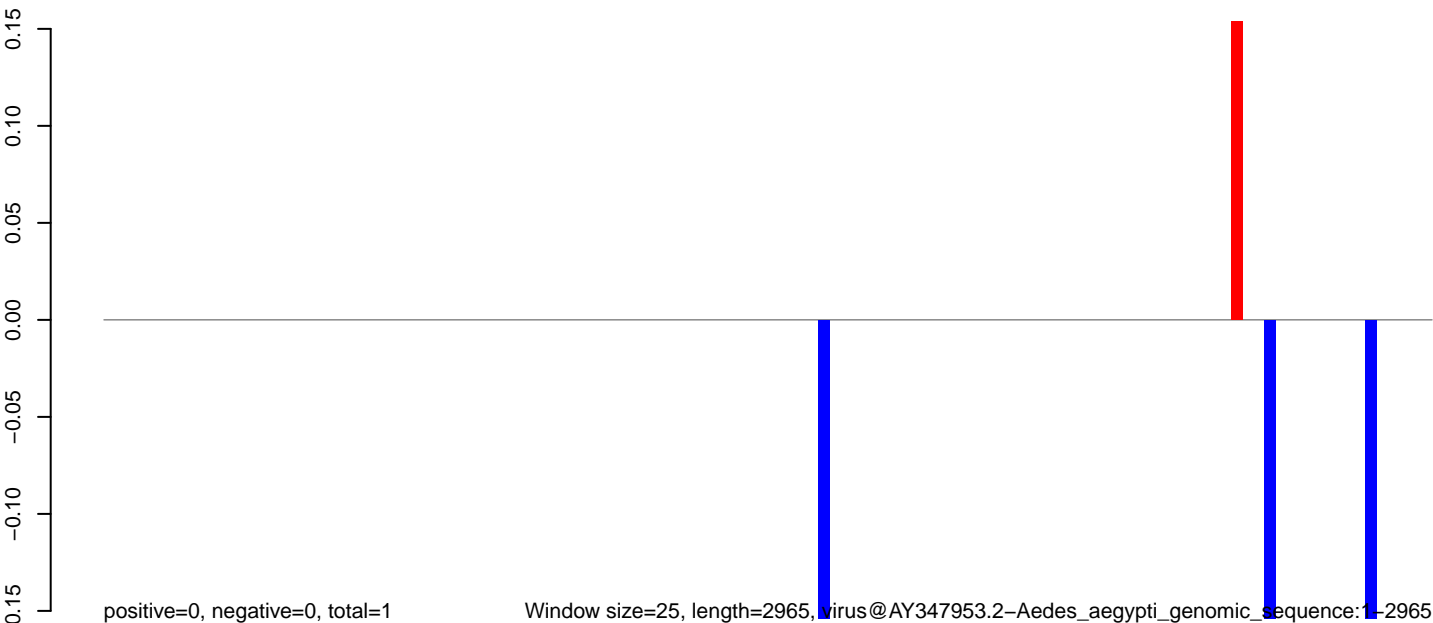
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AeAlbo_C636_6dpi_SINV1.24_35.rep



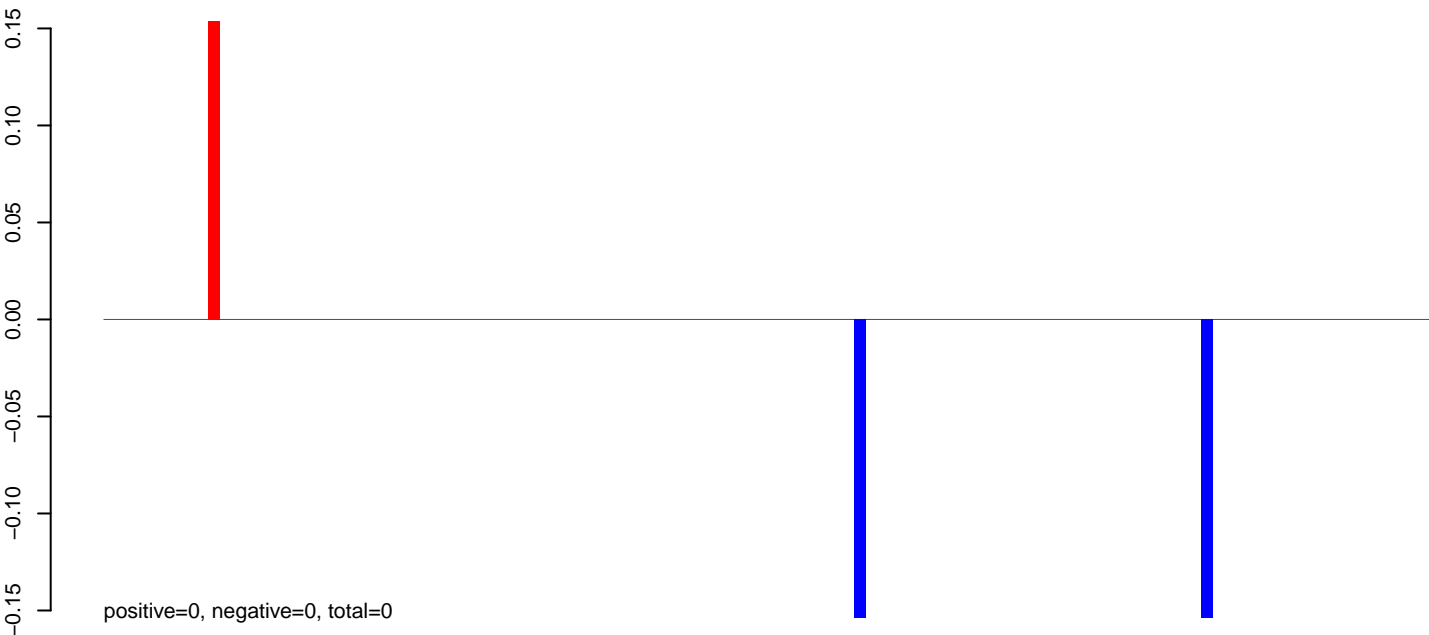
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Window size=25, length=2965, virus@AY347953.2-Aedes_aegypti_genomic_sequence:1-2965

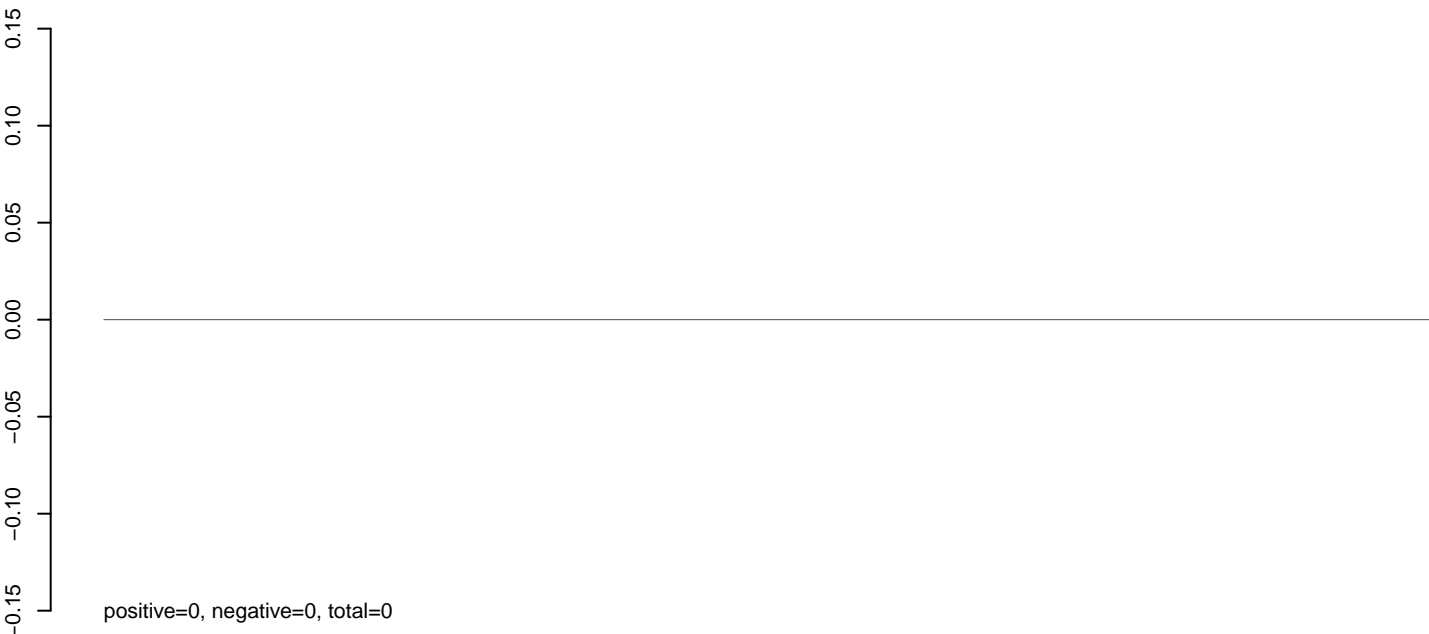
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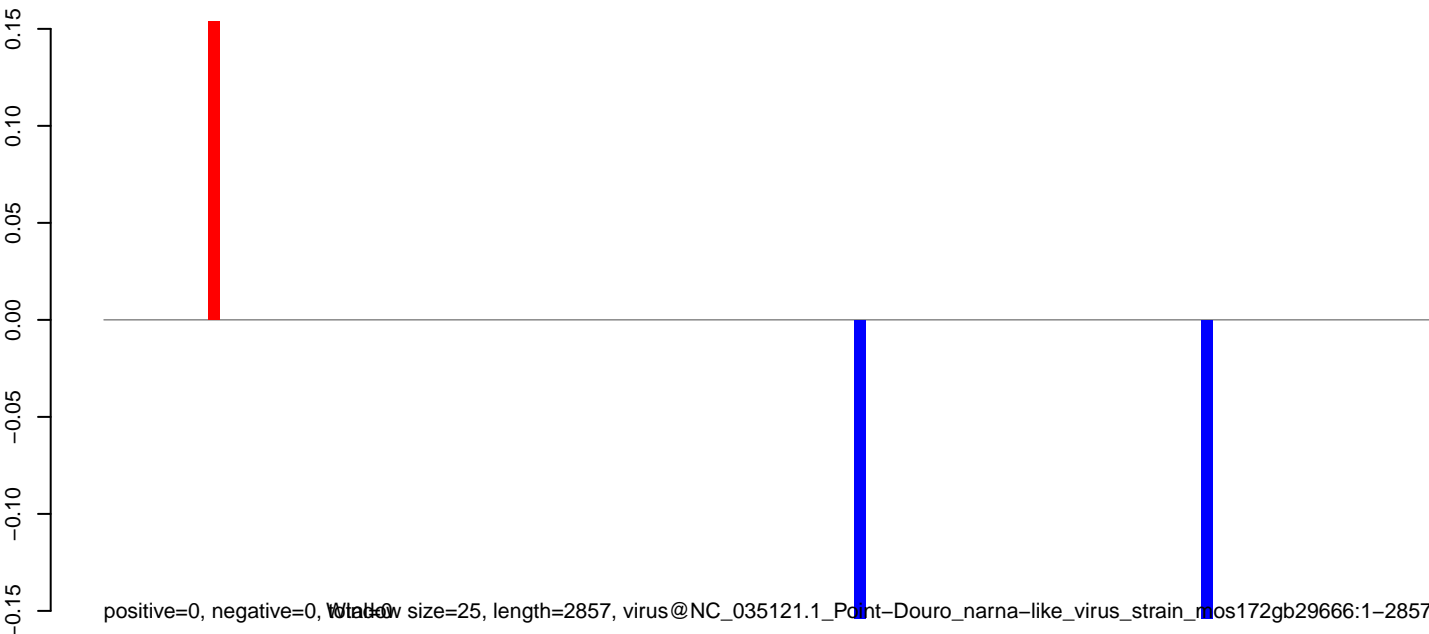
positive=0, negative=0, total=0

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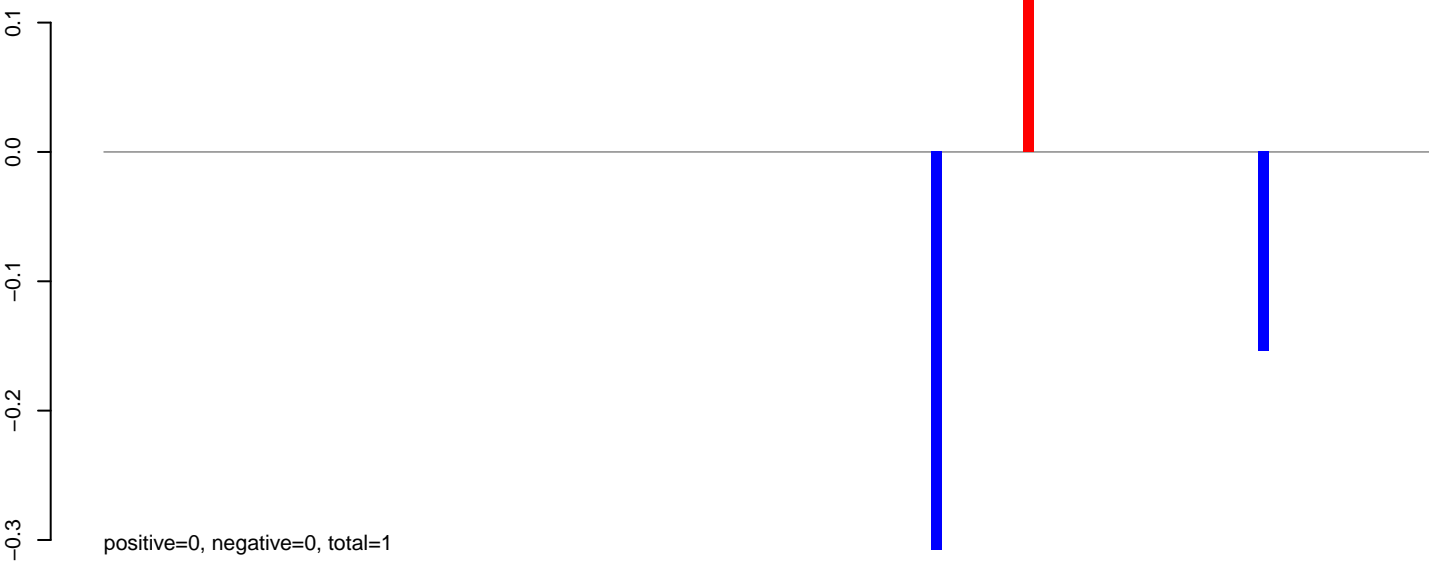
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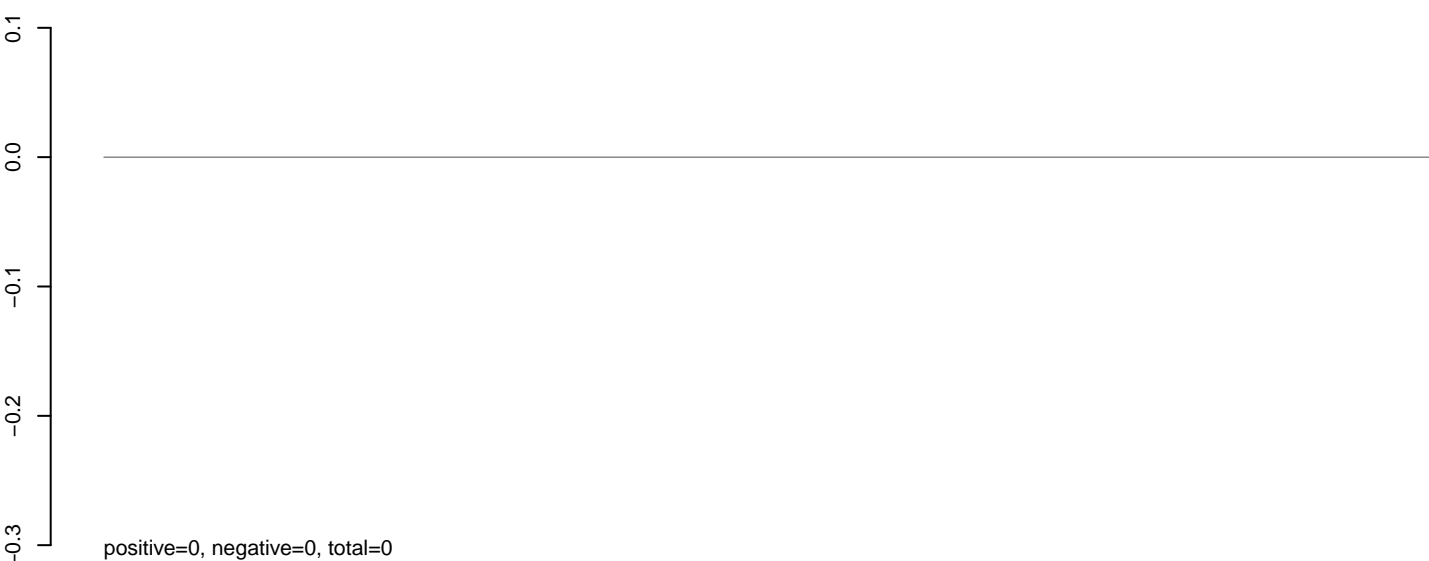


positive=0, negative=0, window size=25, length=2857, virus@NC_035121.1_Point-Douro_narna-like_virus_strain_mos172gb29666:1-2857

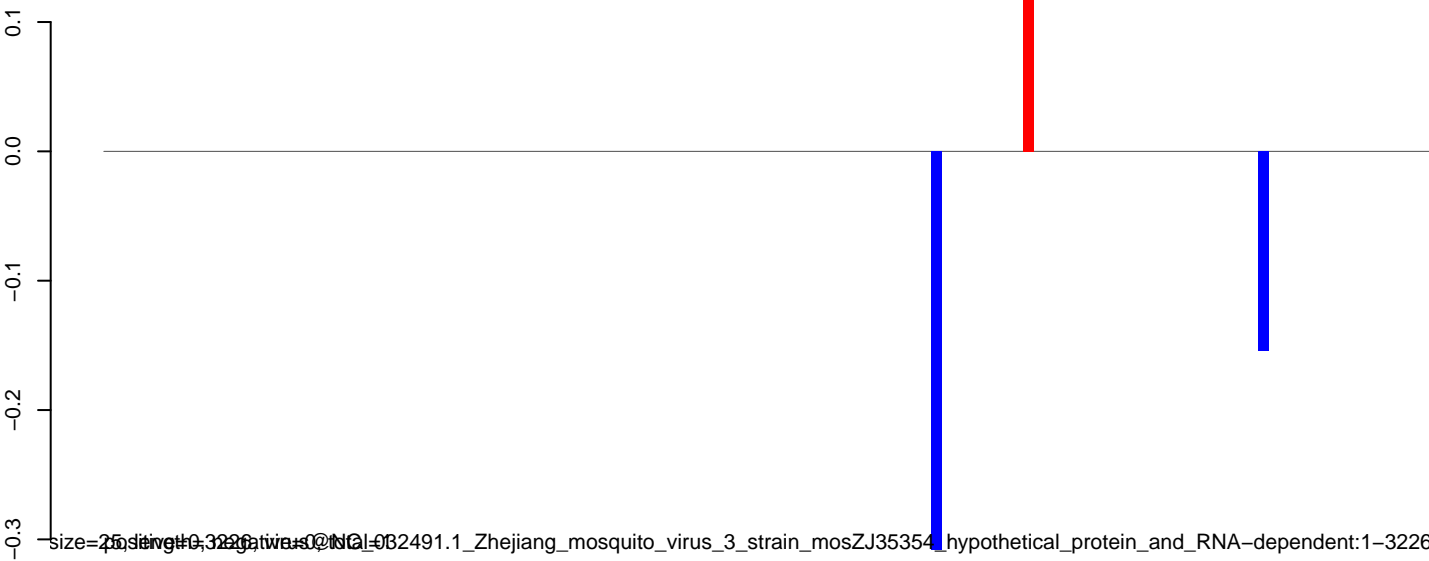
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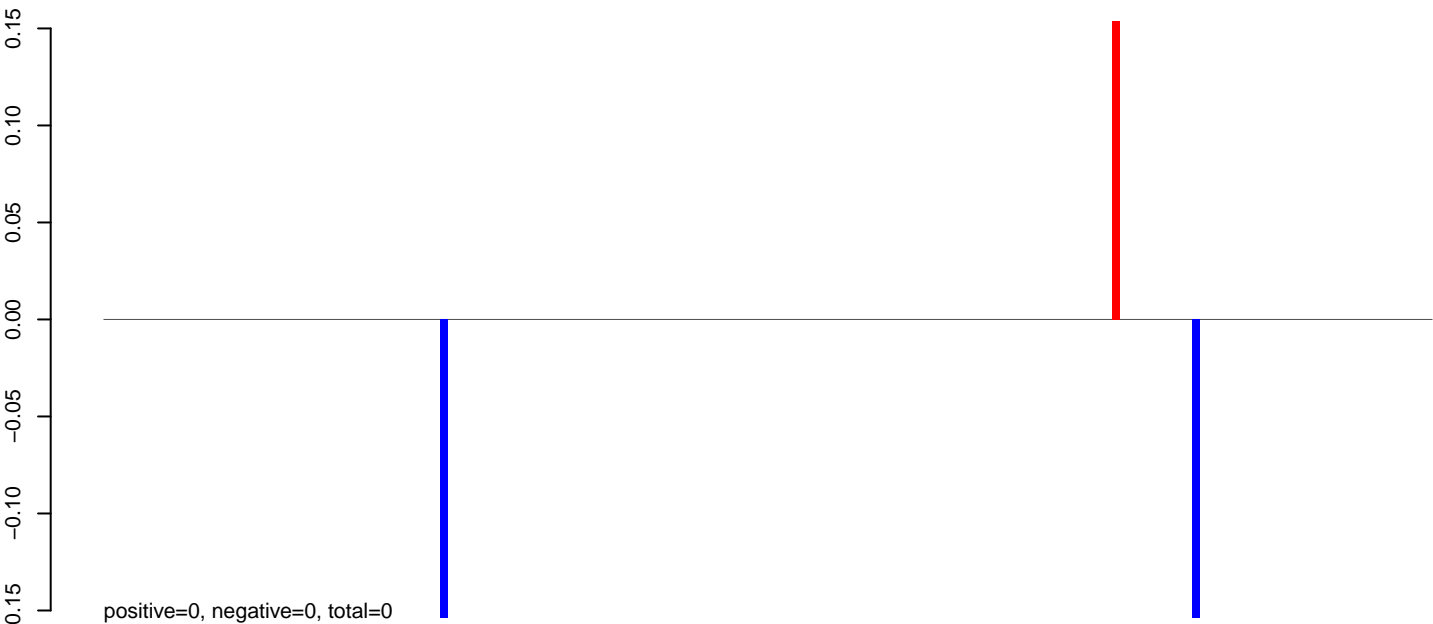
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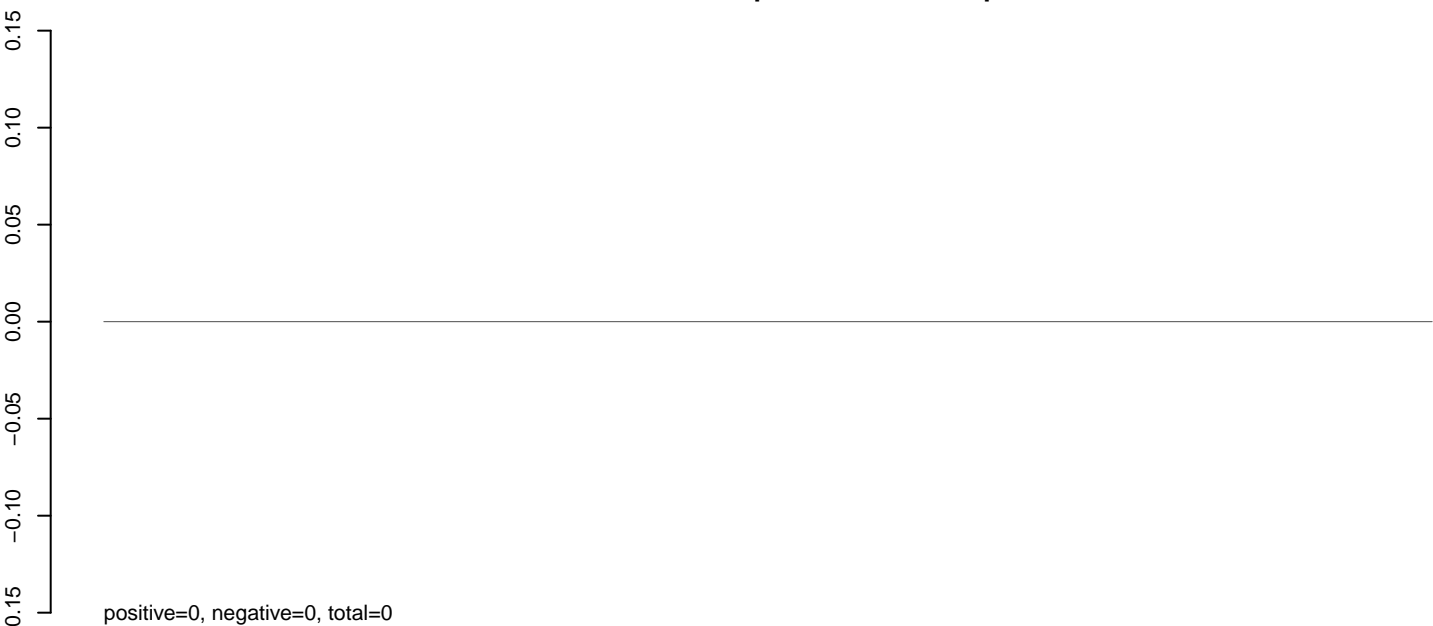
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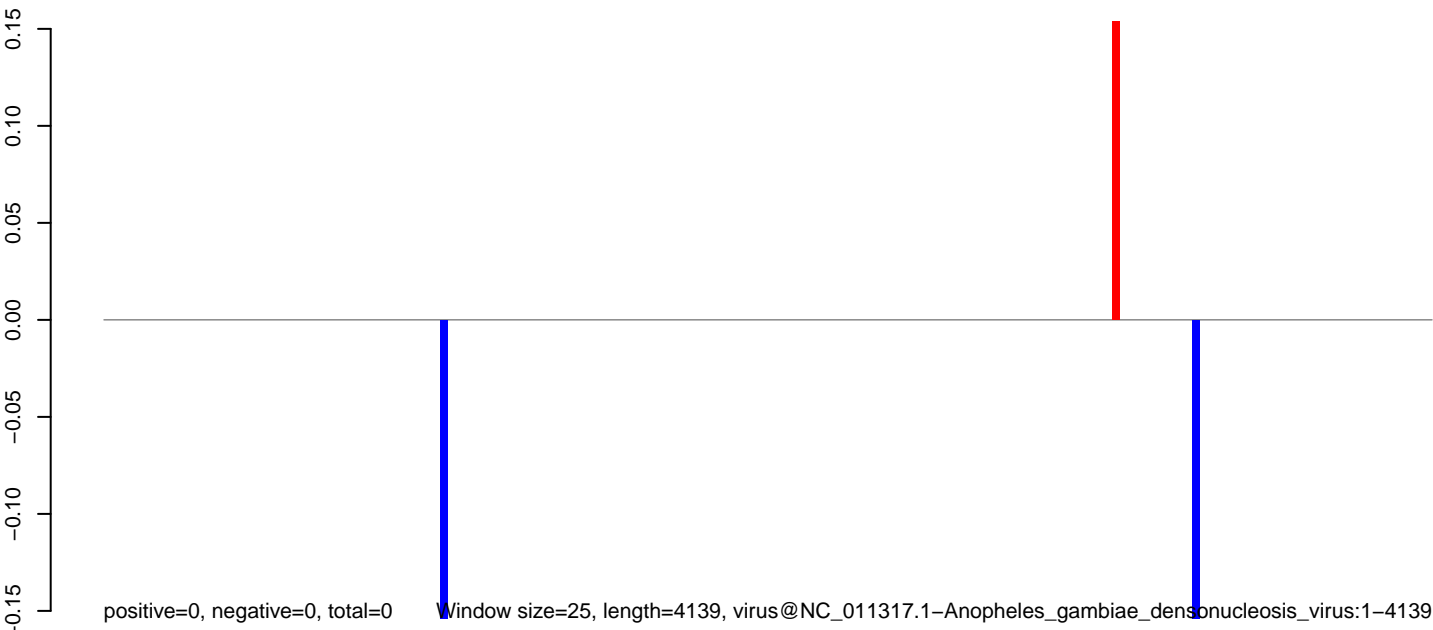
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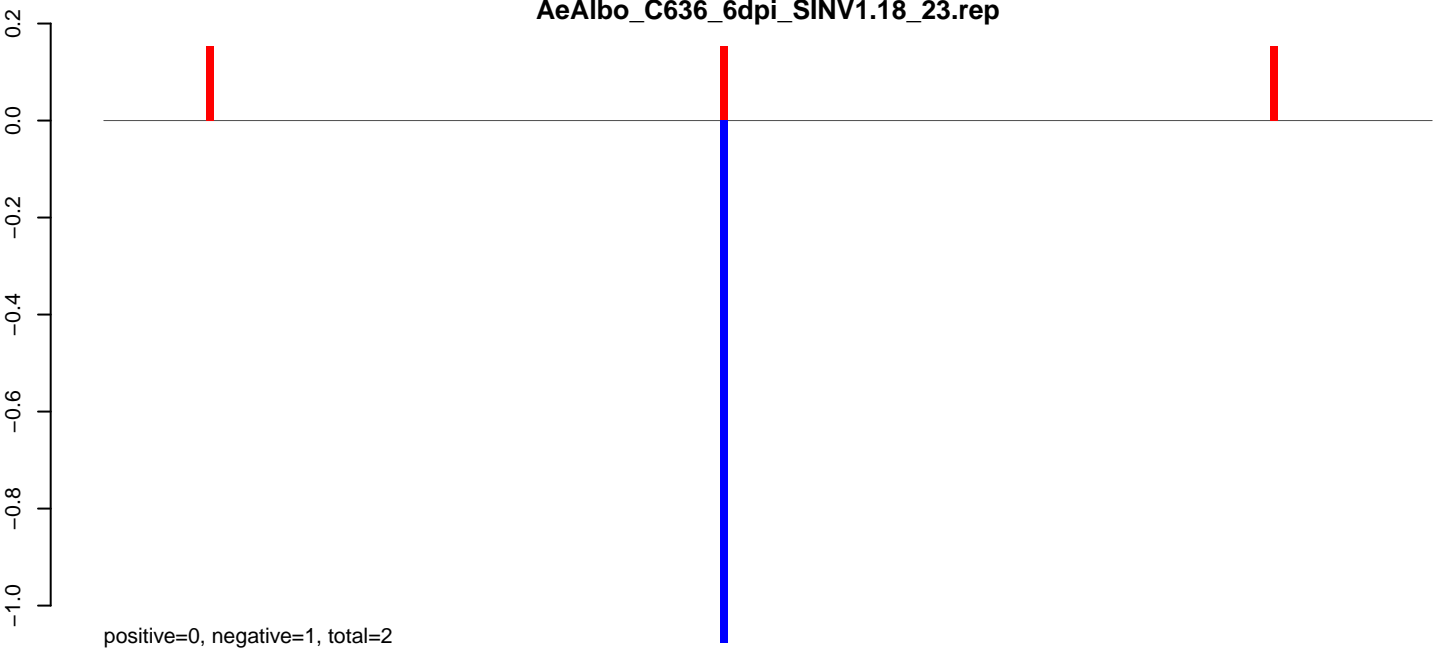
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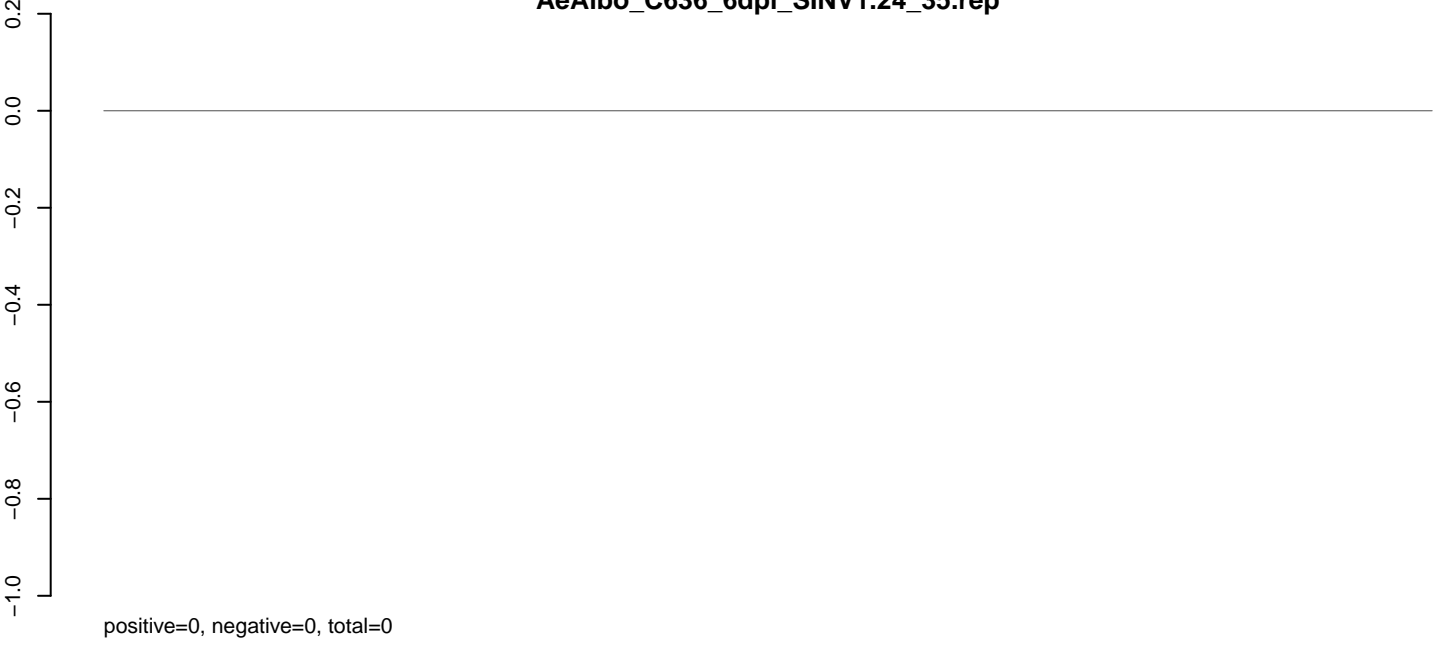
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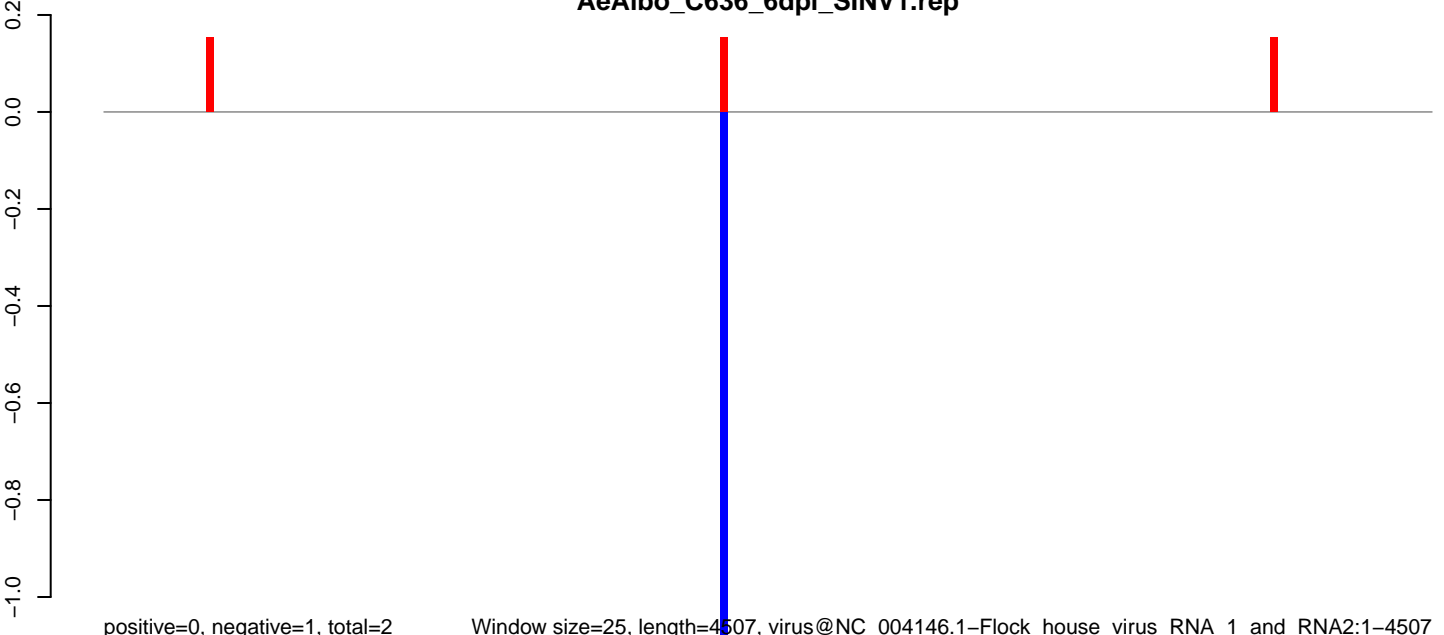
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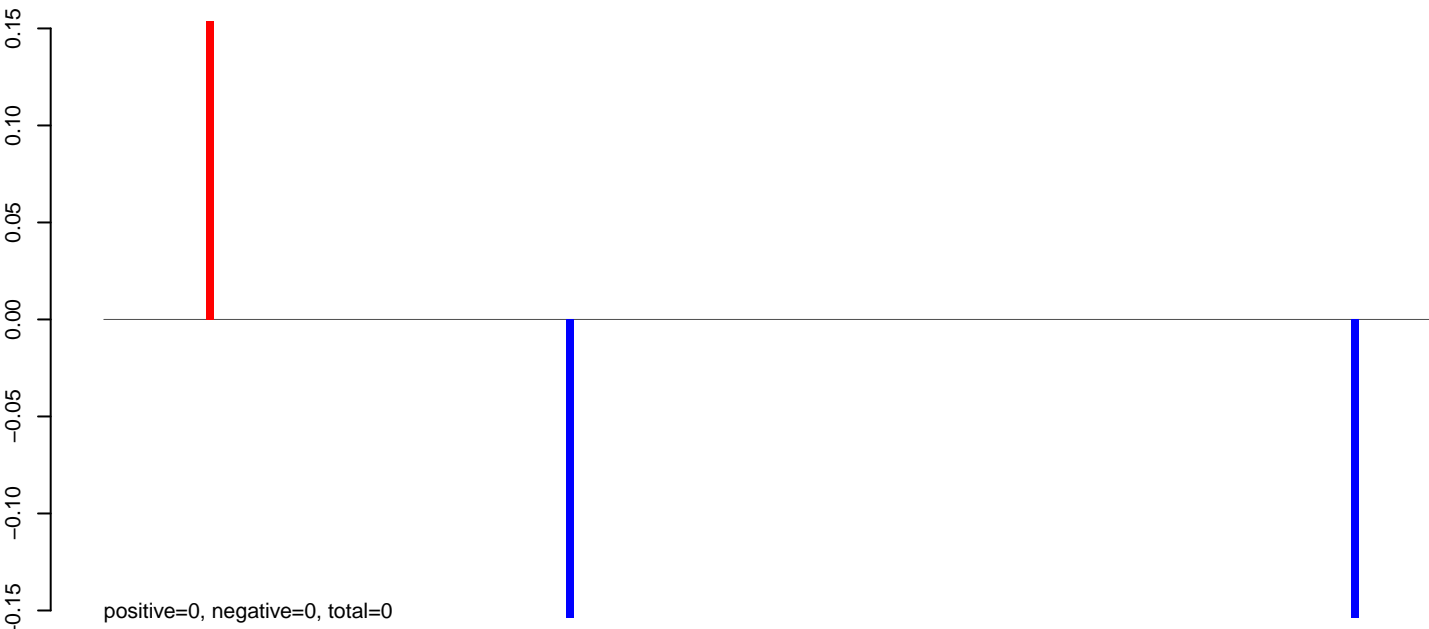
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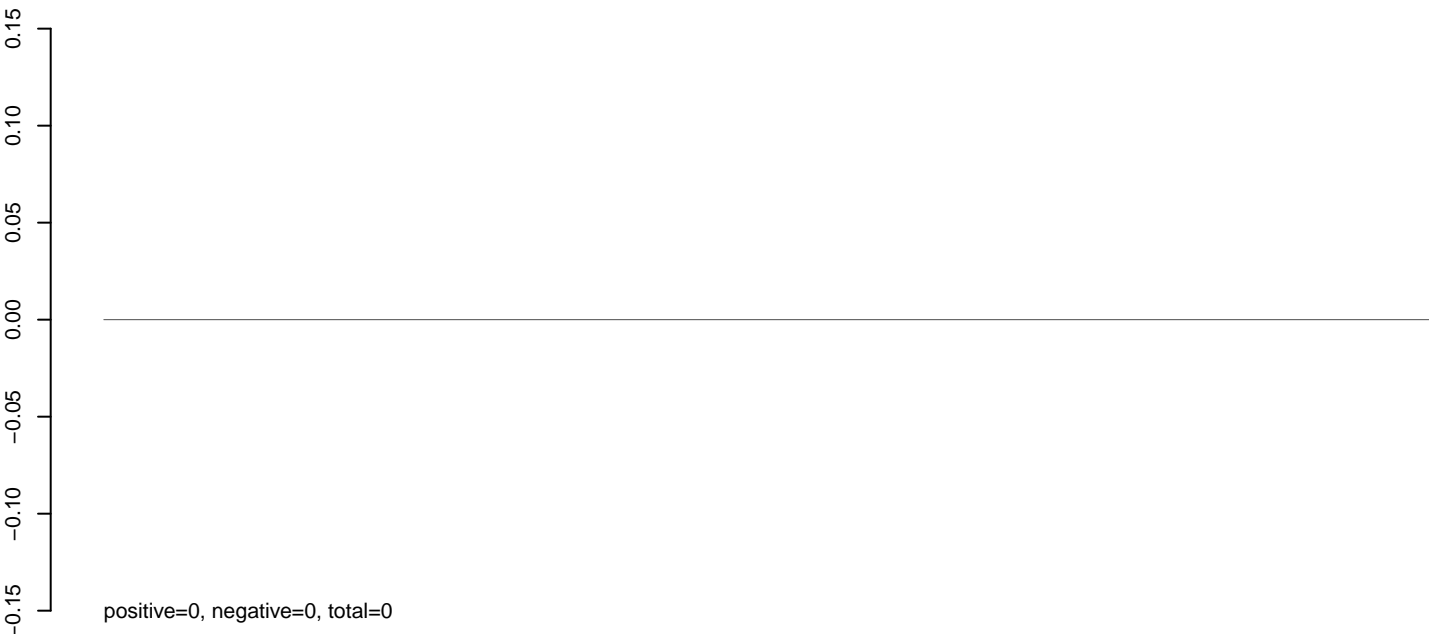
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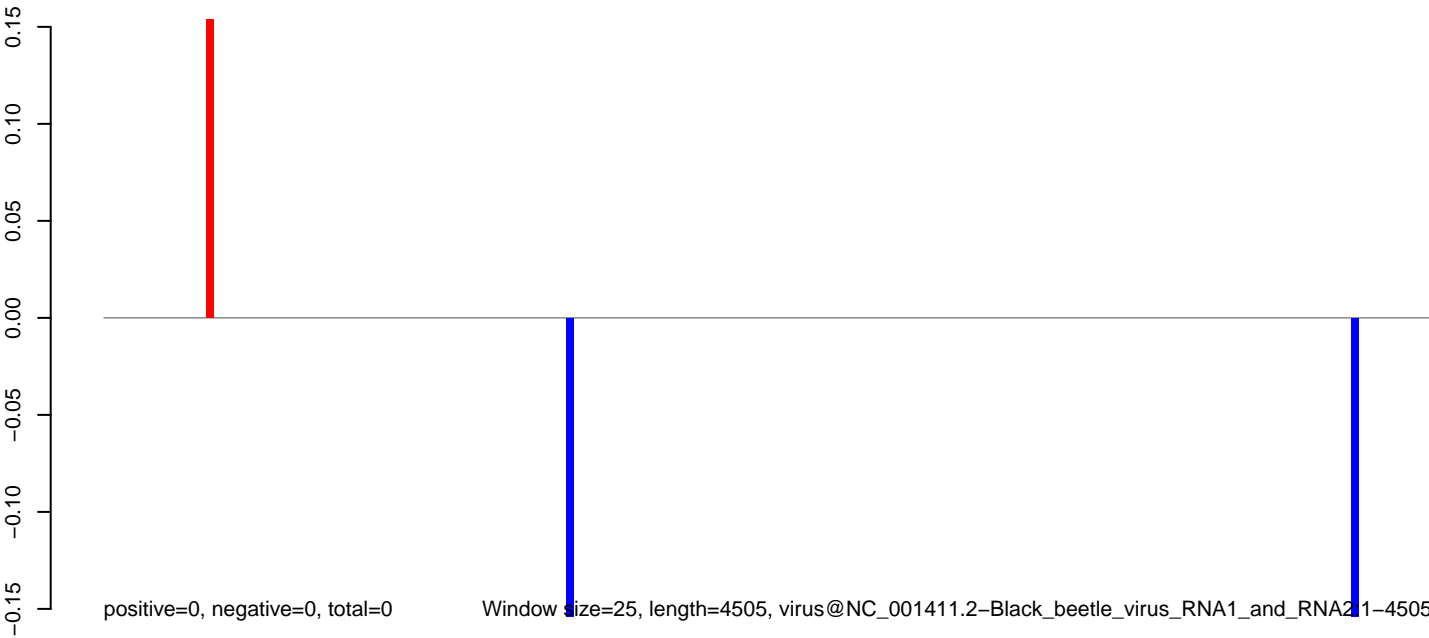
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AeAlbo_C636_6dpi_SINV1.24_35.rep

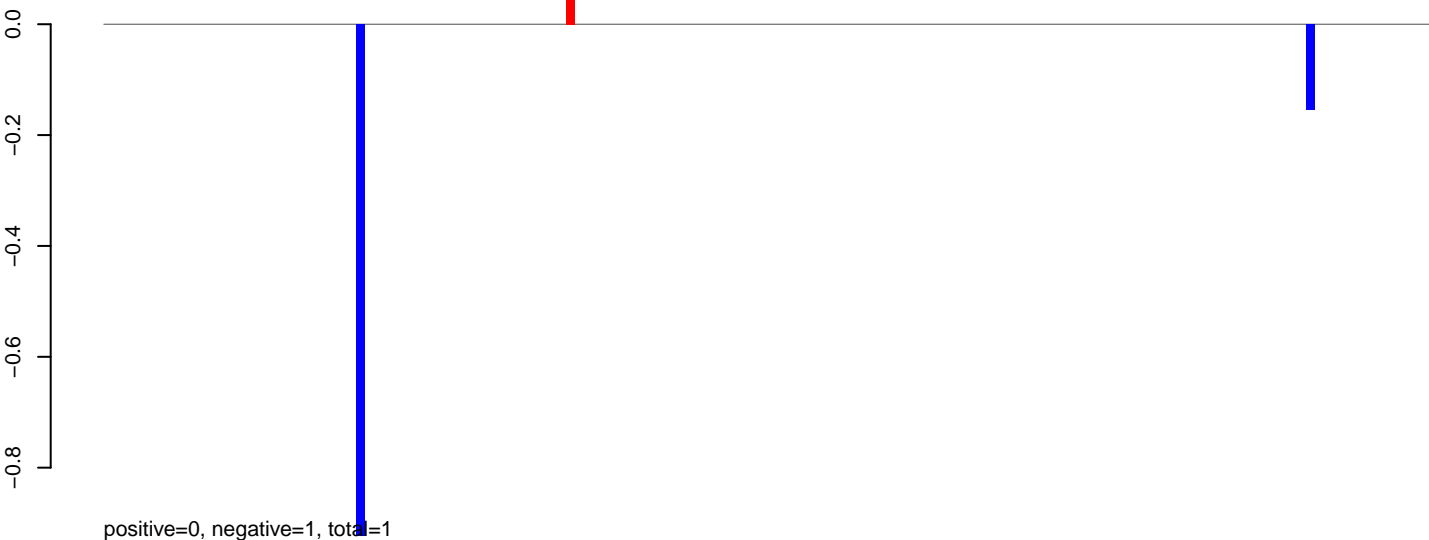


AeAlbo_C636_6dpi_SINV1.rep

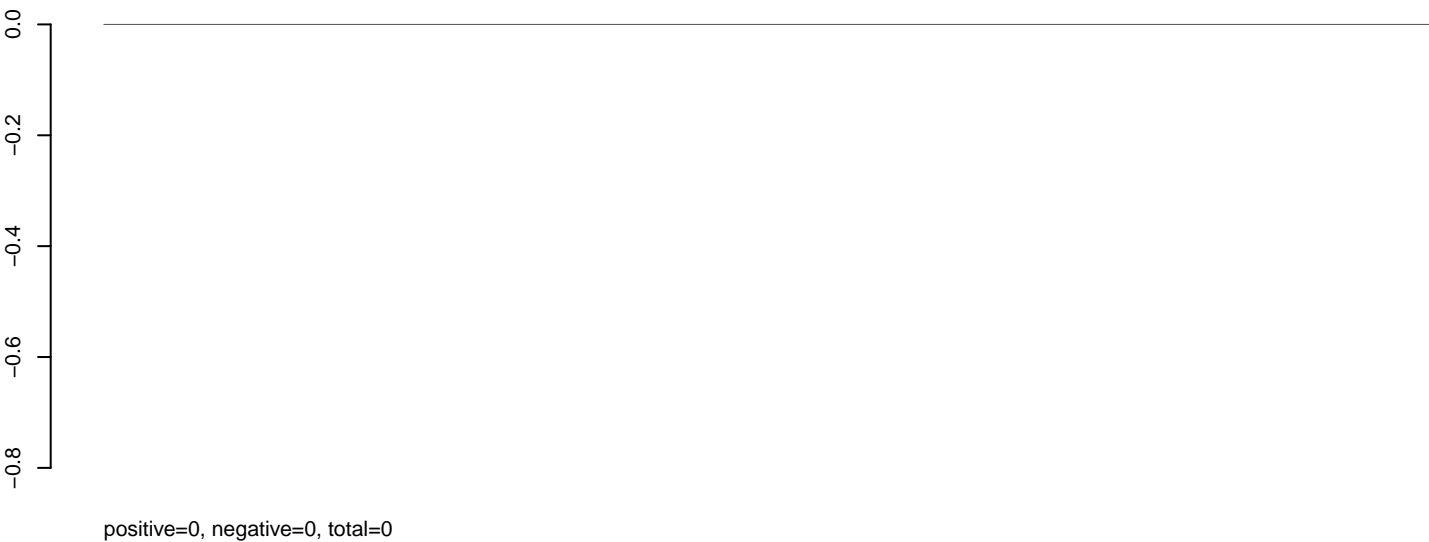


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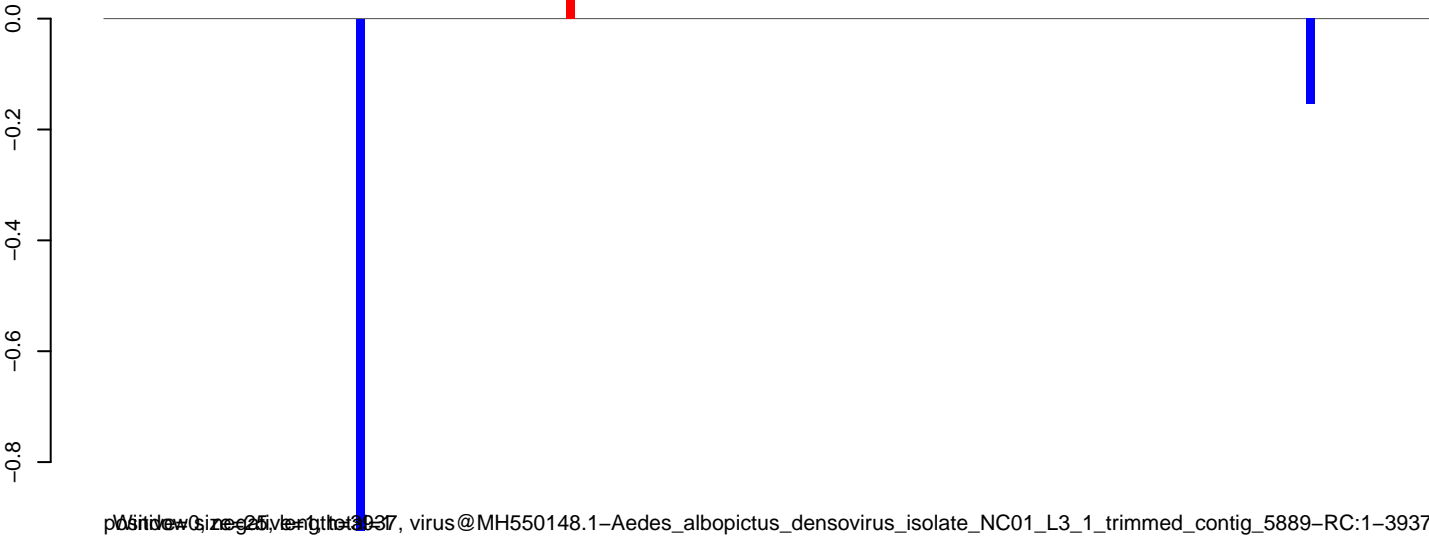
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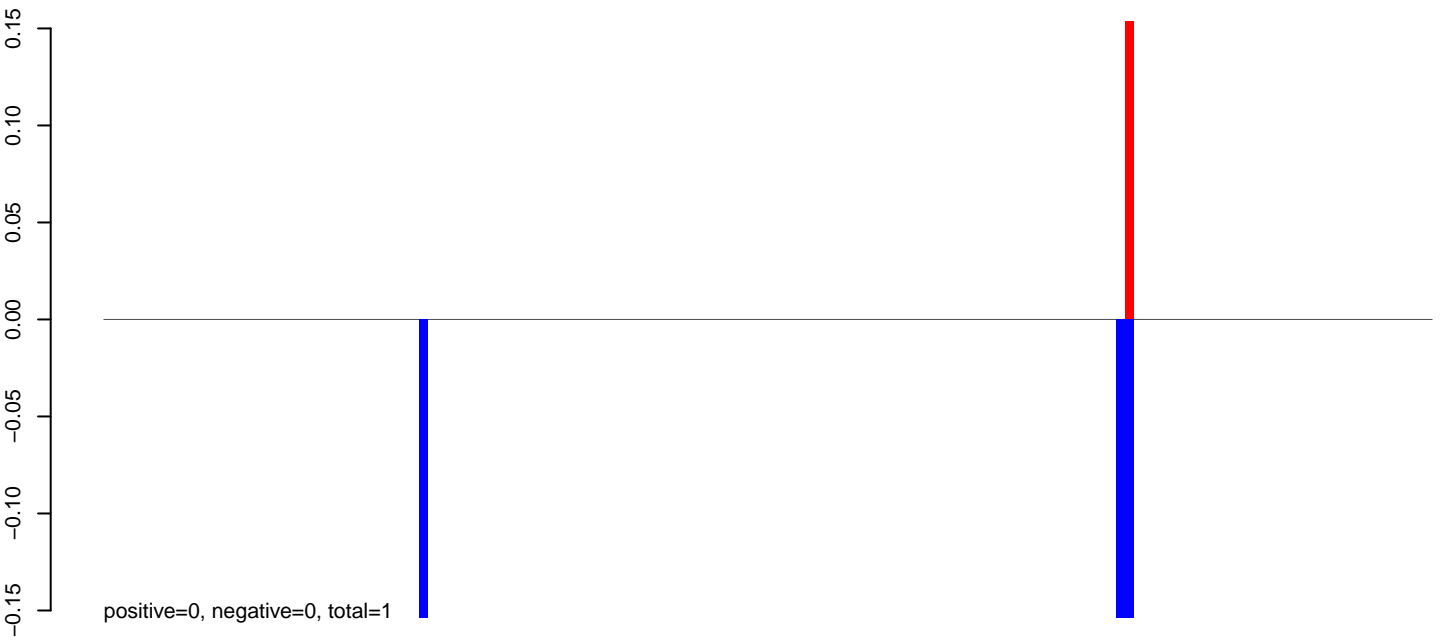
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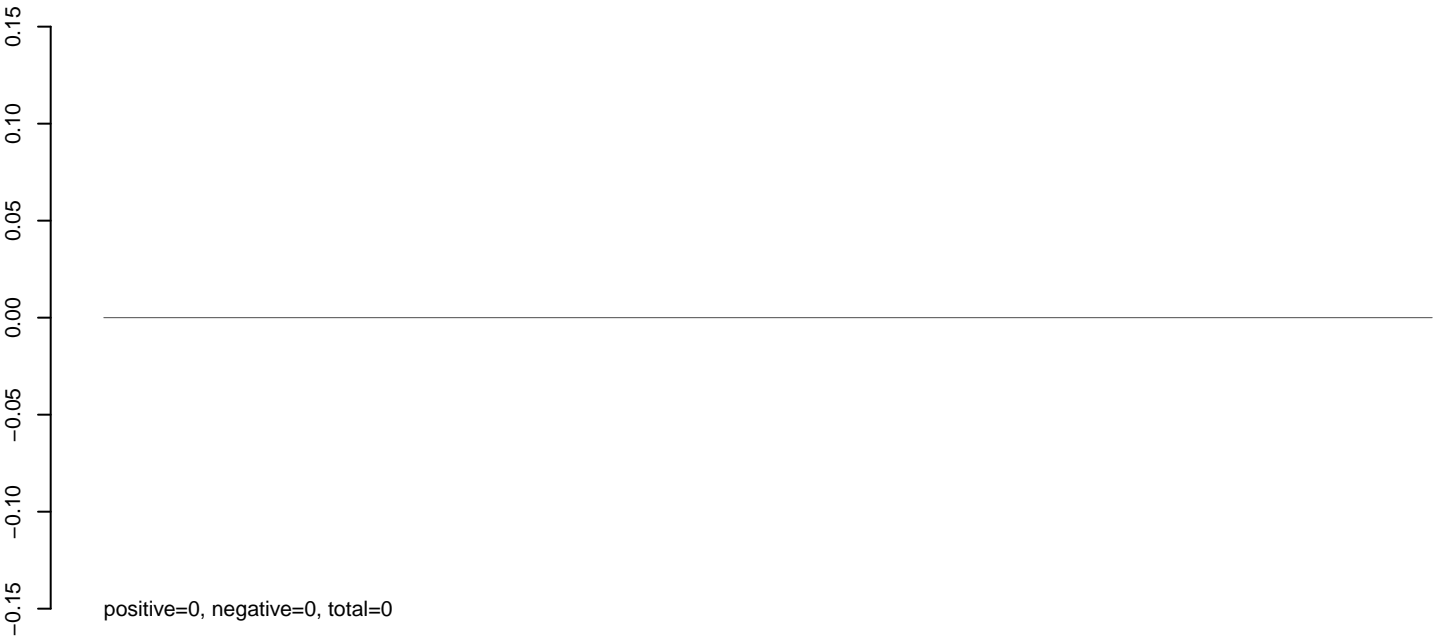
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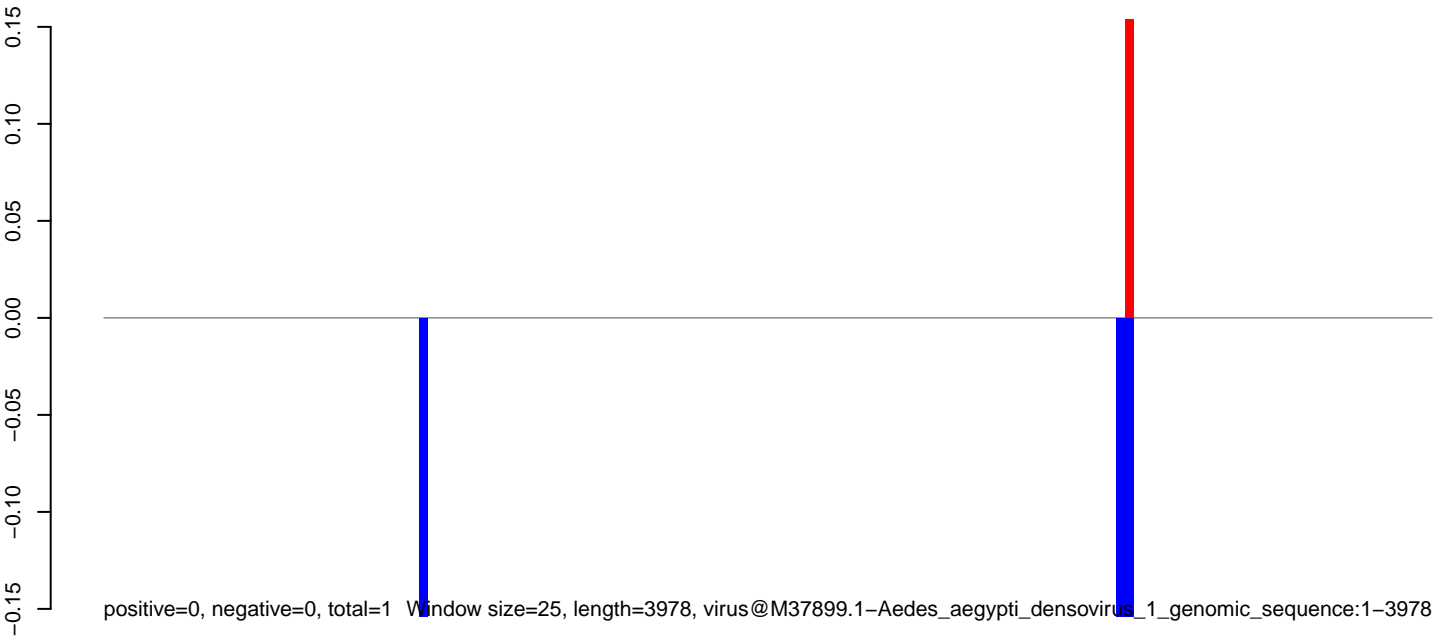
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AeAlbo_C636_6dpi_SINV1.24_35.rep

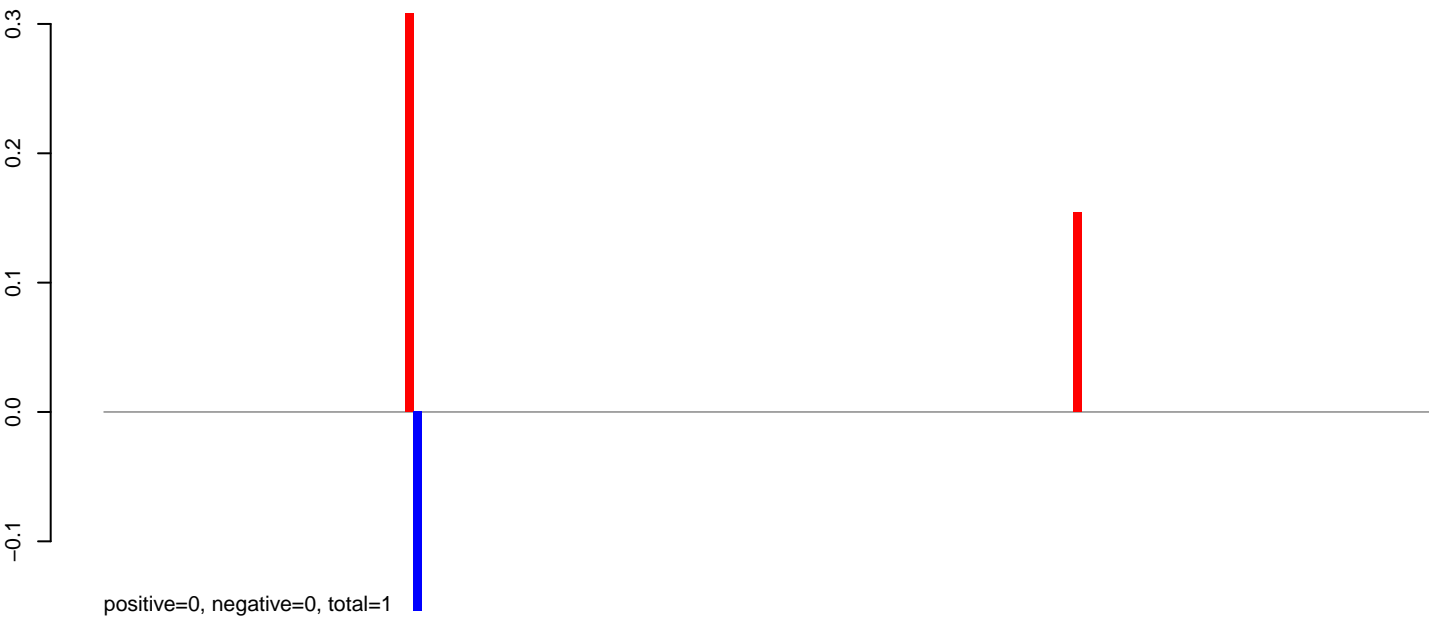


AeAlbo_C636_6dpi_SINV1.rep



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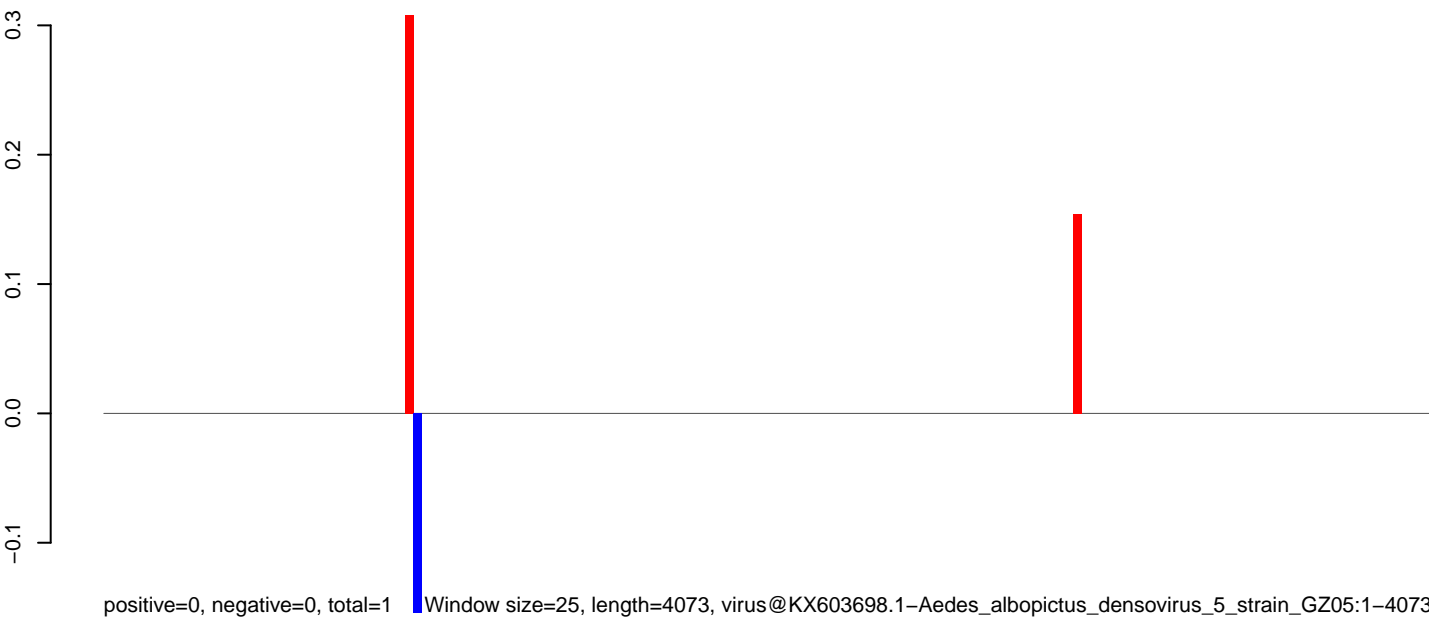
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AeAlbo_C636_6dpi_SINV1.24_35.rep



AeAlbo_C636_6dpi_SINV1.rep



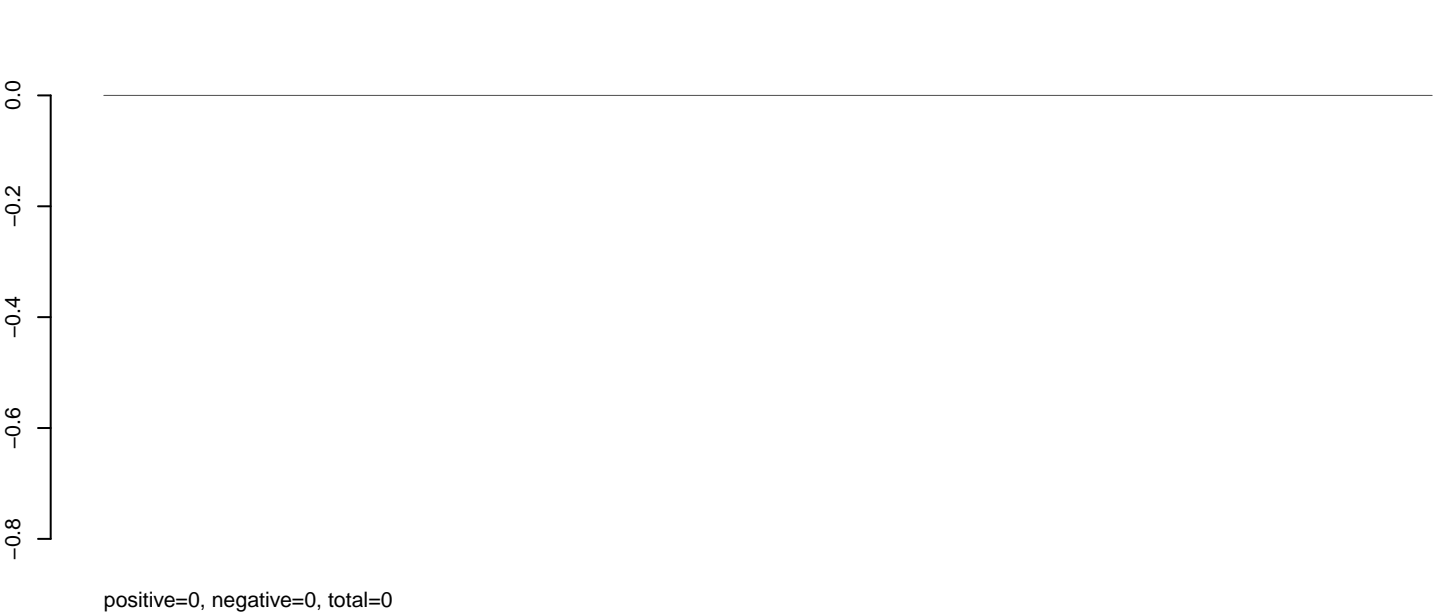
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AeAlbo_C636_6dpi_SINV1.18_23.rep



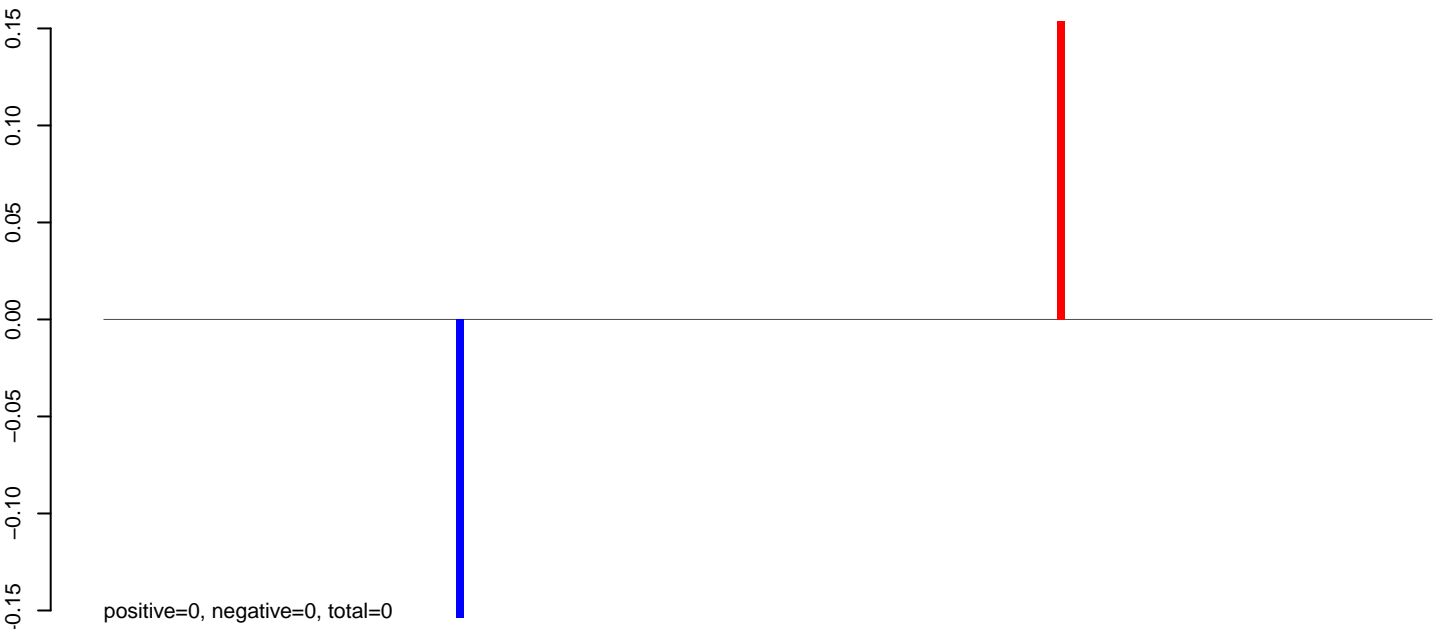
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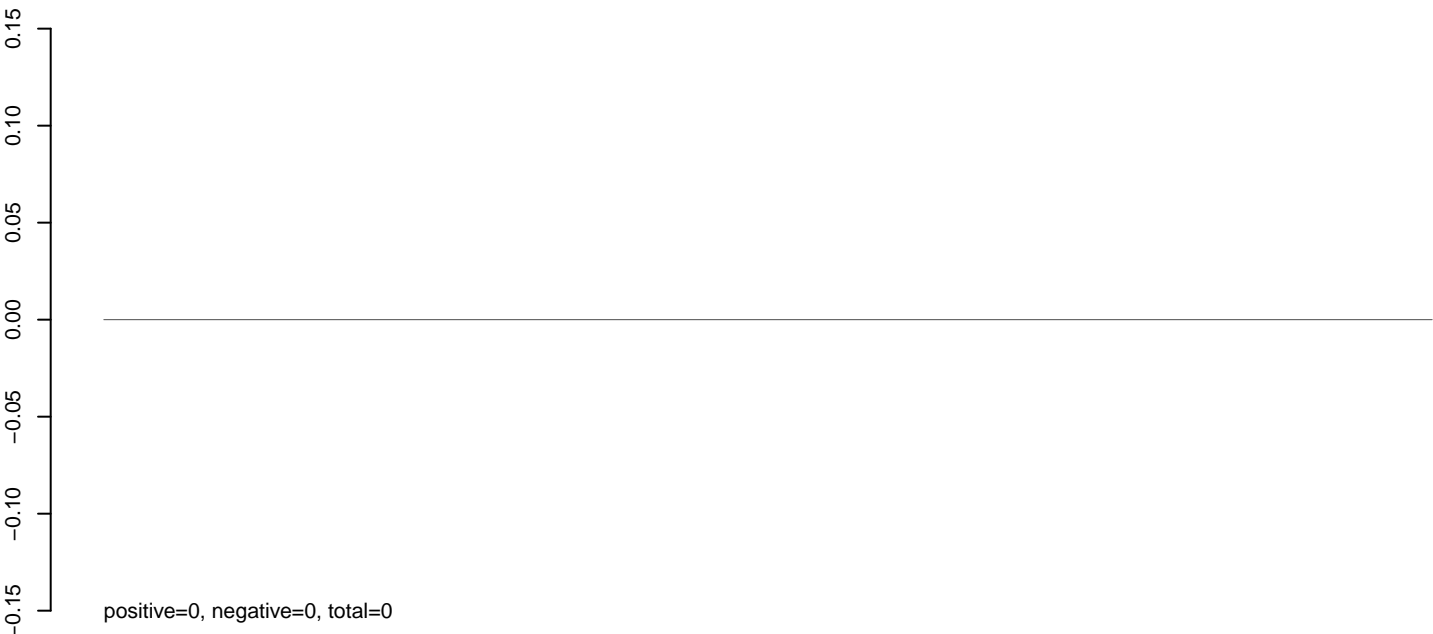
AeAlbo_C636_6dpi_SINV1.rep



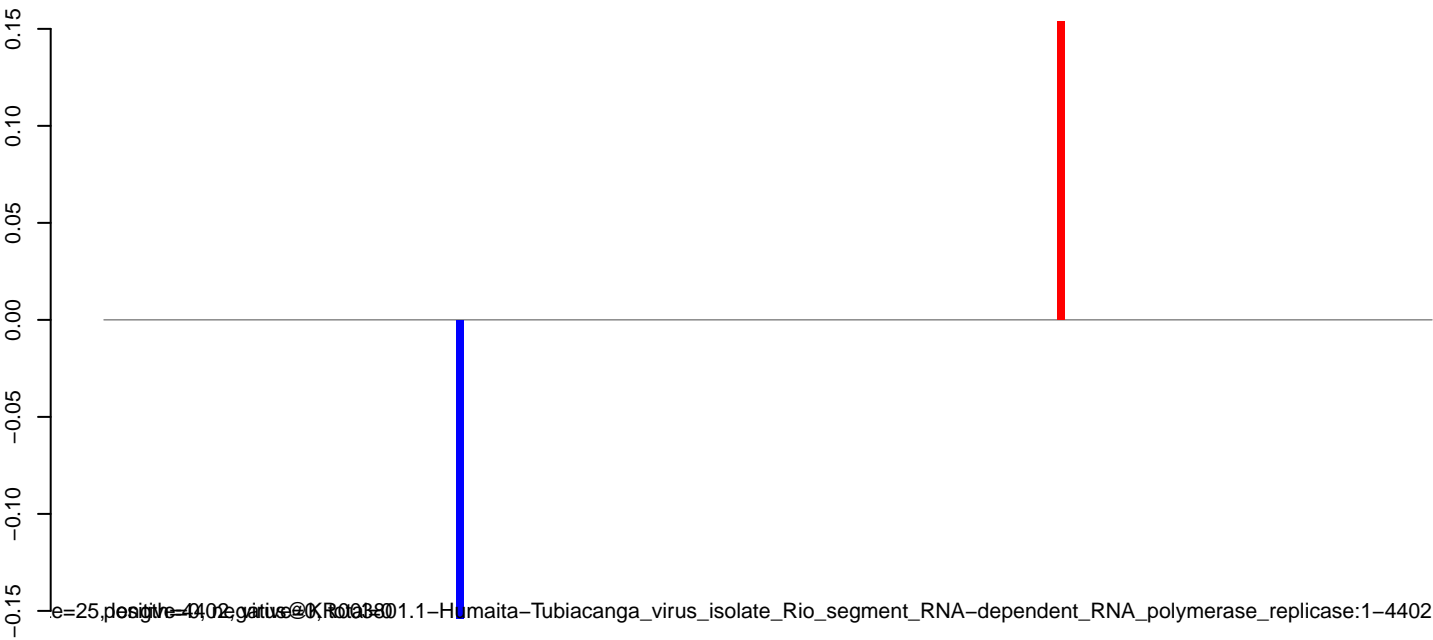
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AeAlbo_C636_6dpi_SINV1.24_35.rep



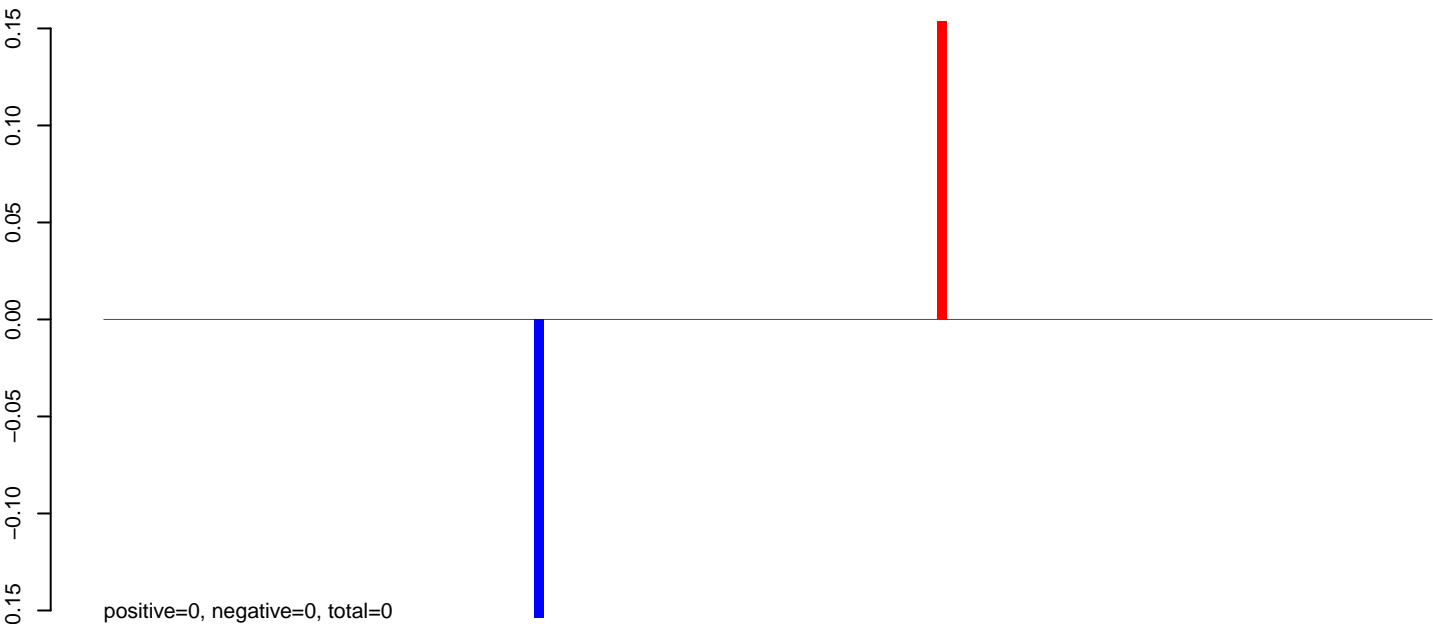
AeAlbo_C636_6dpi_SINV1.rep



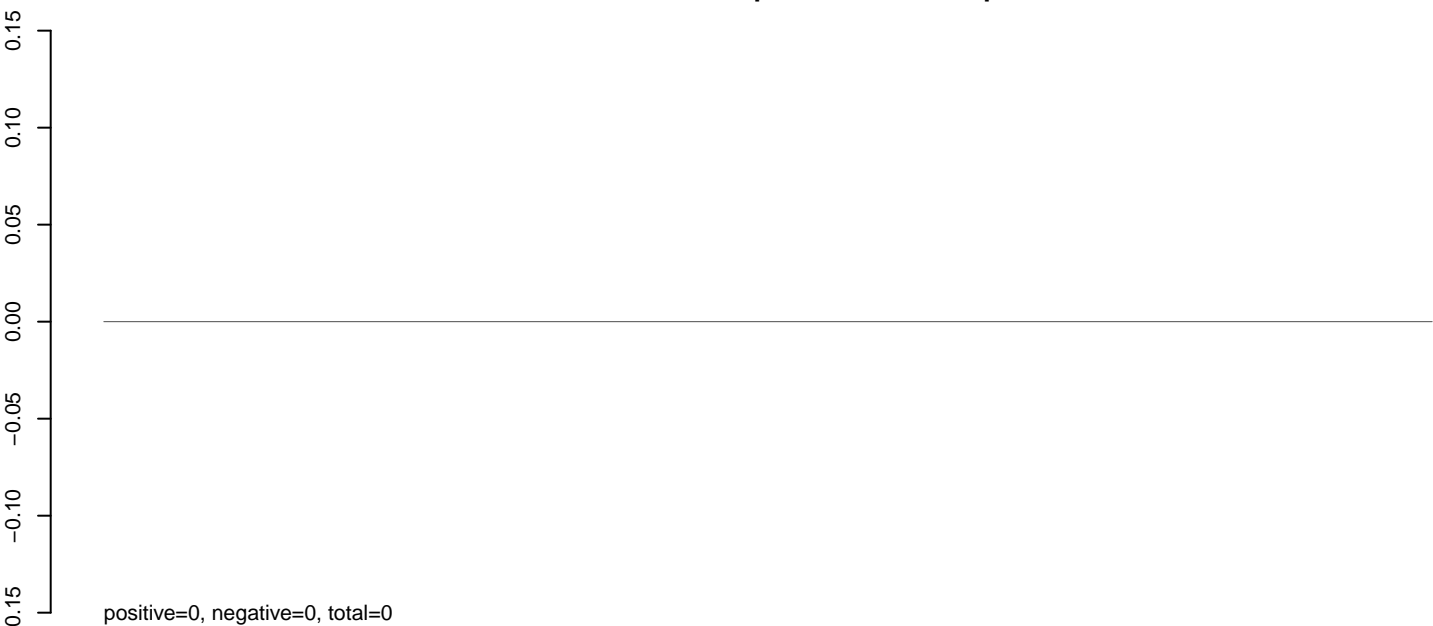
e=25, positive=0, negative=0, total=0, KR038011.1-Himaita-Tubiacanga_virus_isolate_Rio_segment_RNA-dependent_RNA_polymerase_replicase:1-4402

0 1000 2000 3000 4000

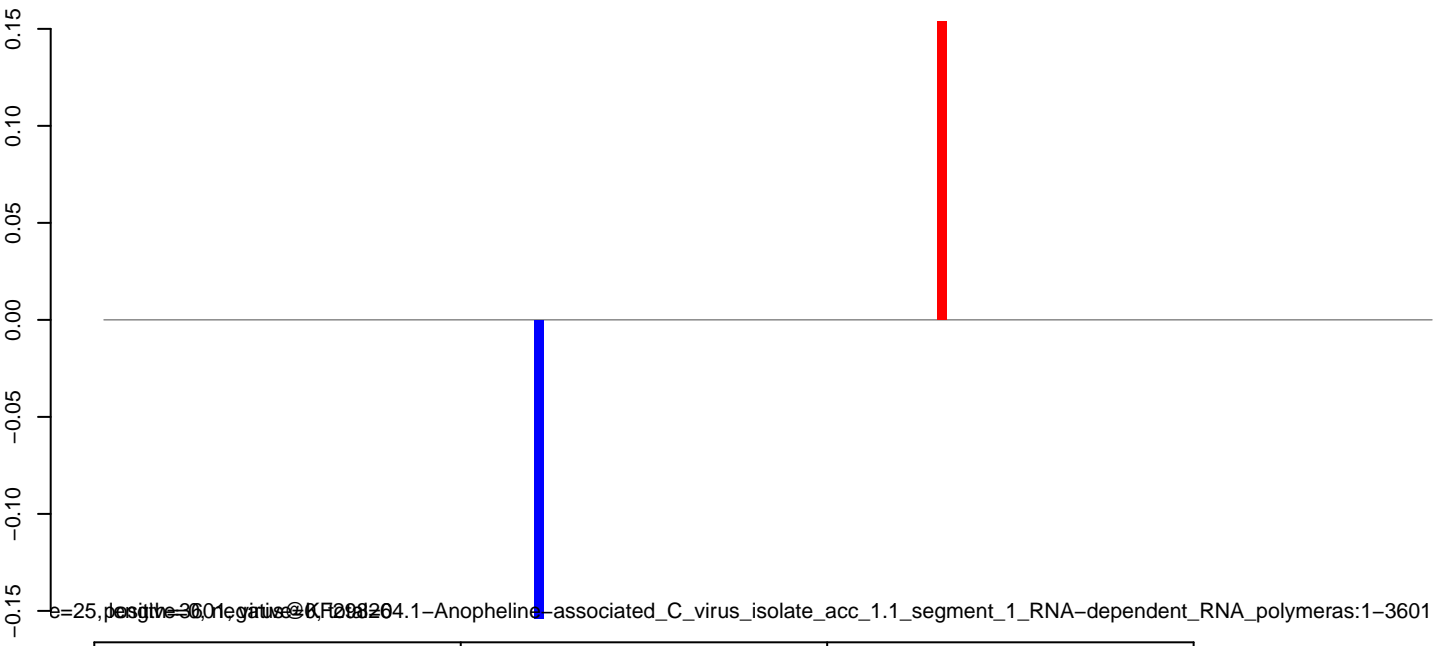
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AeAlbo_C636_6dpi_SINV1.24_35.rep



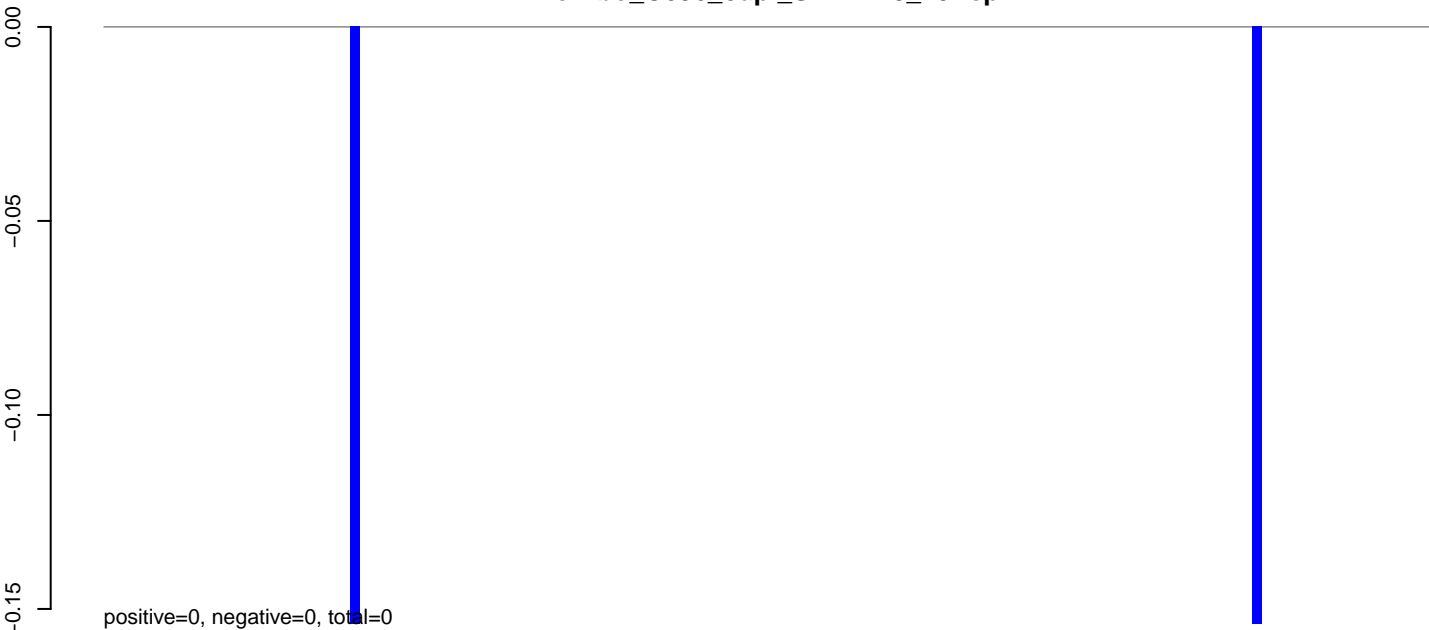
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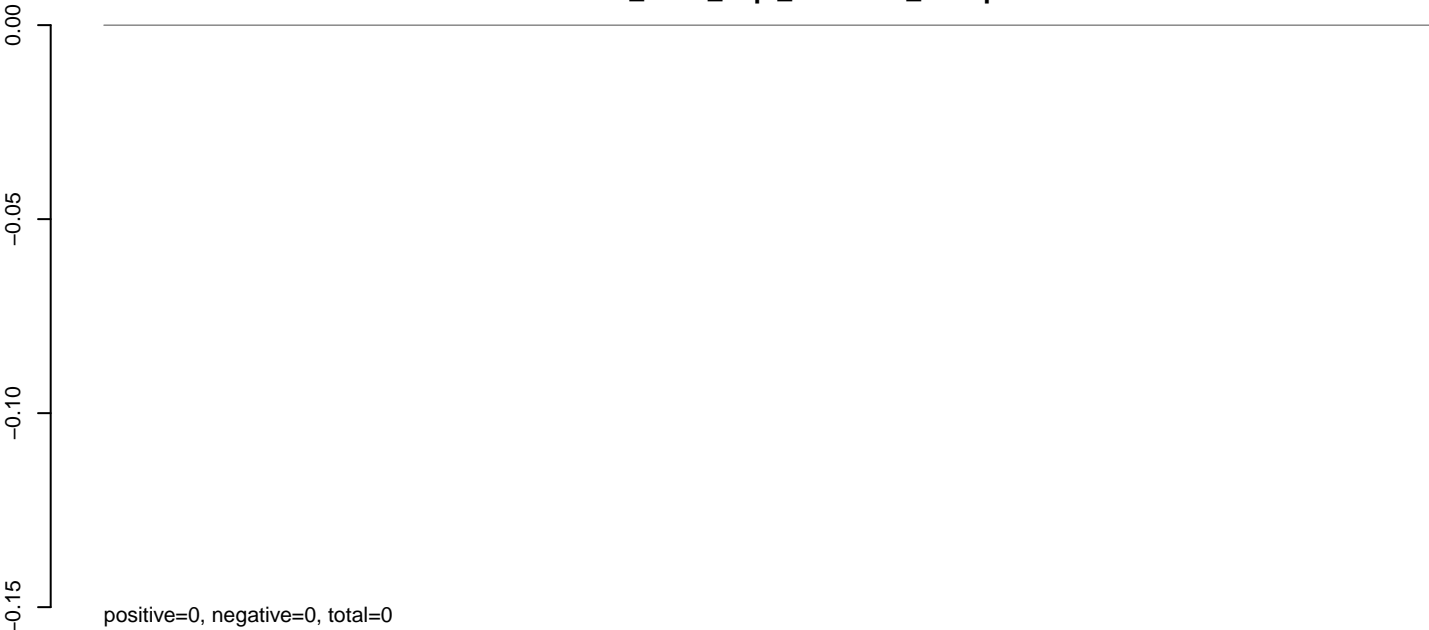
g=25, positive=0, negative=0, total=0, R=204.1 - Anopheline-associated_C_virus_isolate_acc.1.1_segment_1_RNA-dependent_RNA_polymerase:1-3601

0 1000 2000 3000

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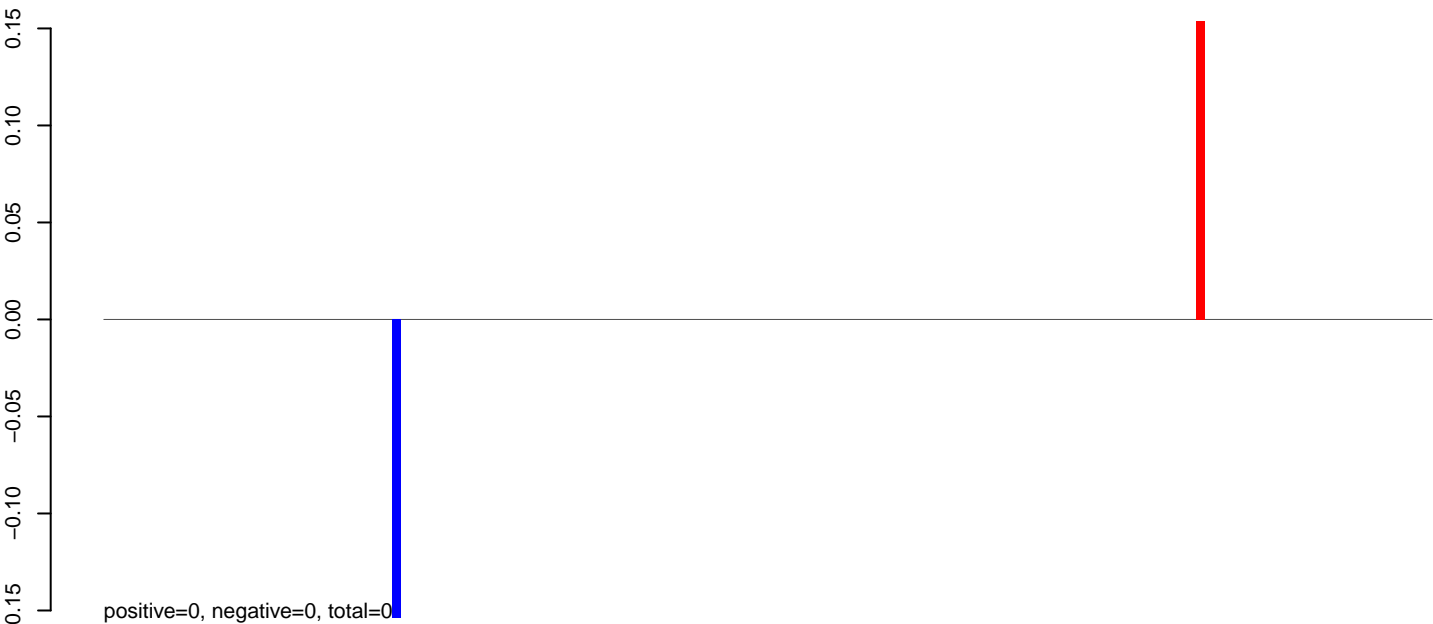
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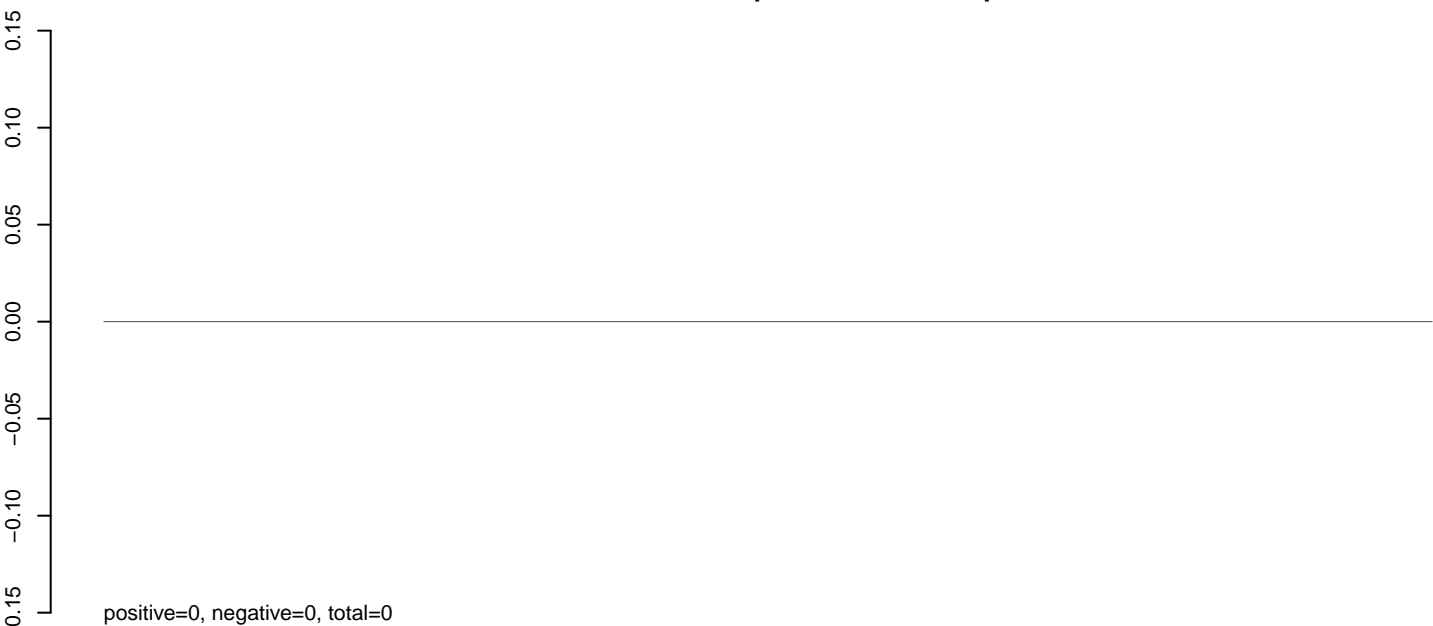
579756.1-Culex pipiens pallens densovirus strain JZ-16_non-structural_protein_1_NS1_non-stru:1-3498

0 500 1000 1500 2000 2500 3000 3500

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AeAlbo_C636_6dpi_SINV1.24_35.rep



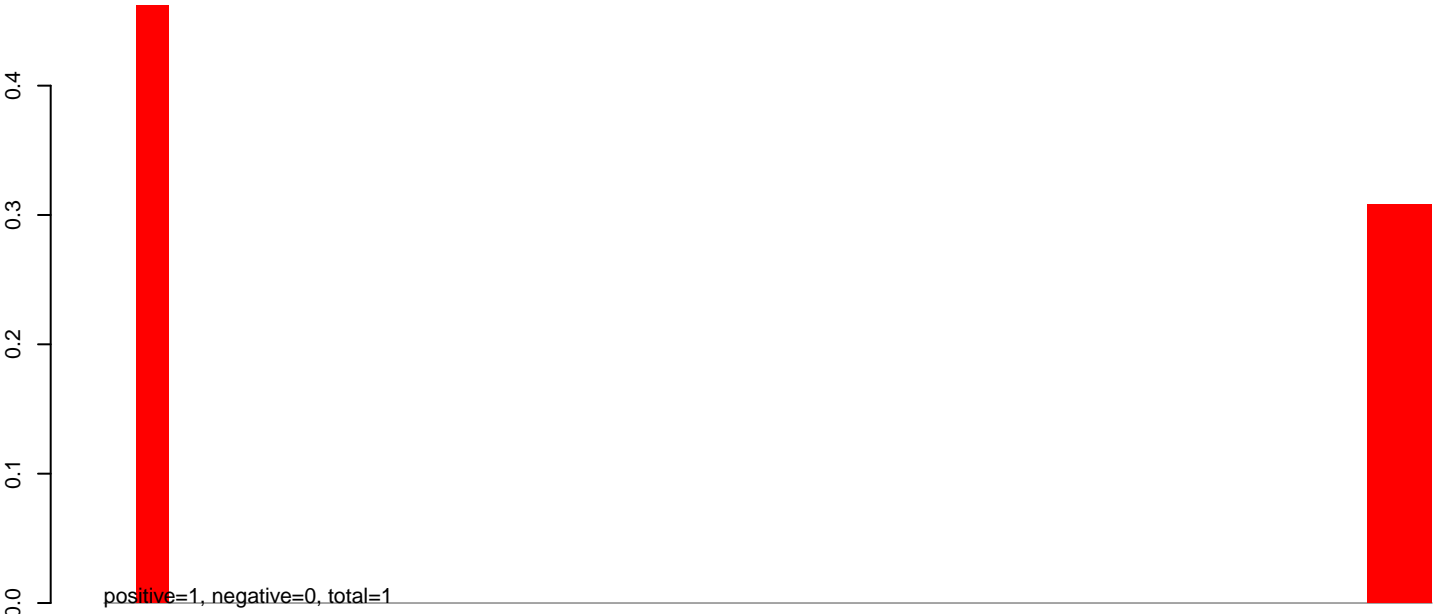
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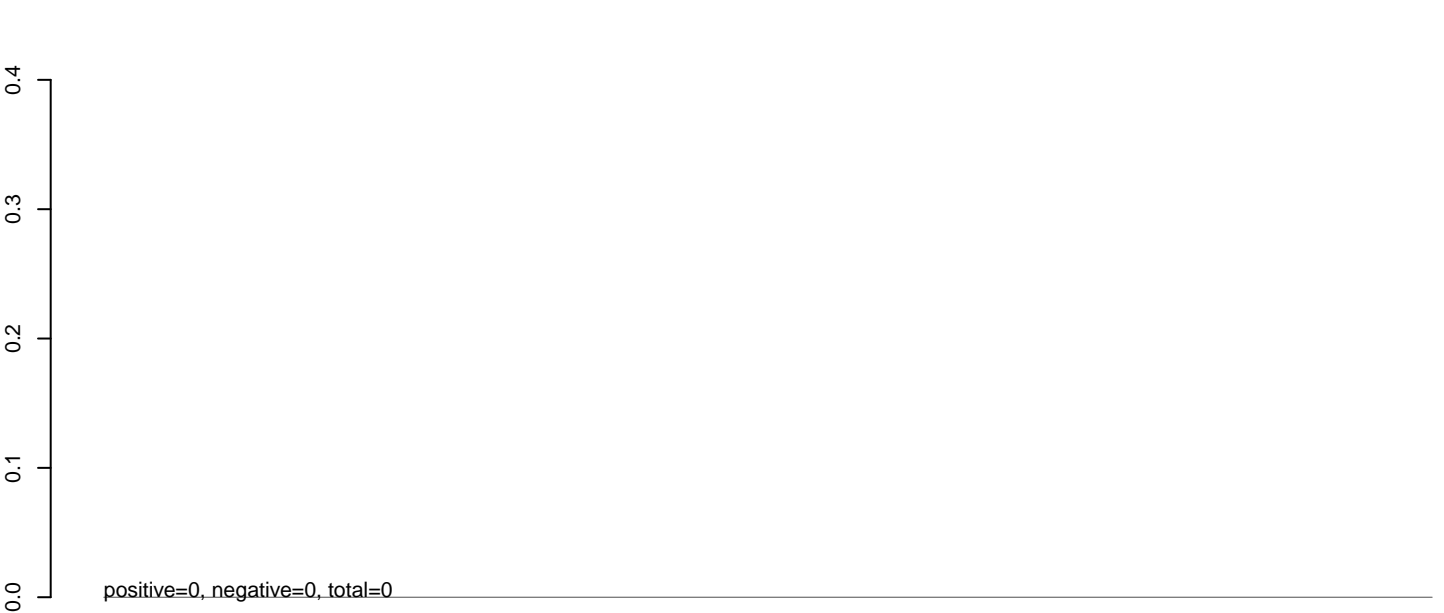
Window size=25, length=3790, virus@AeAeg_Thai_Densovirus_c30446_g1_i1_len=3790:1-3790

0 1000 2000 3000

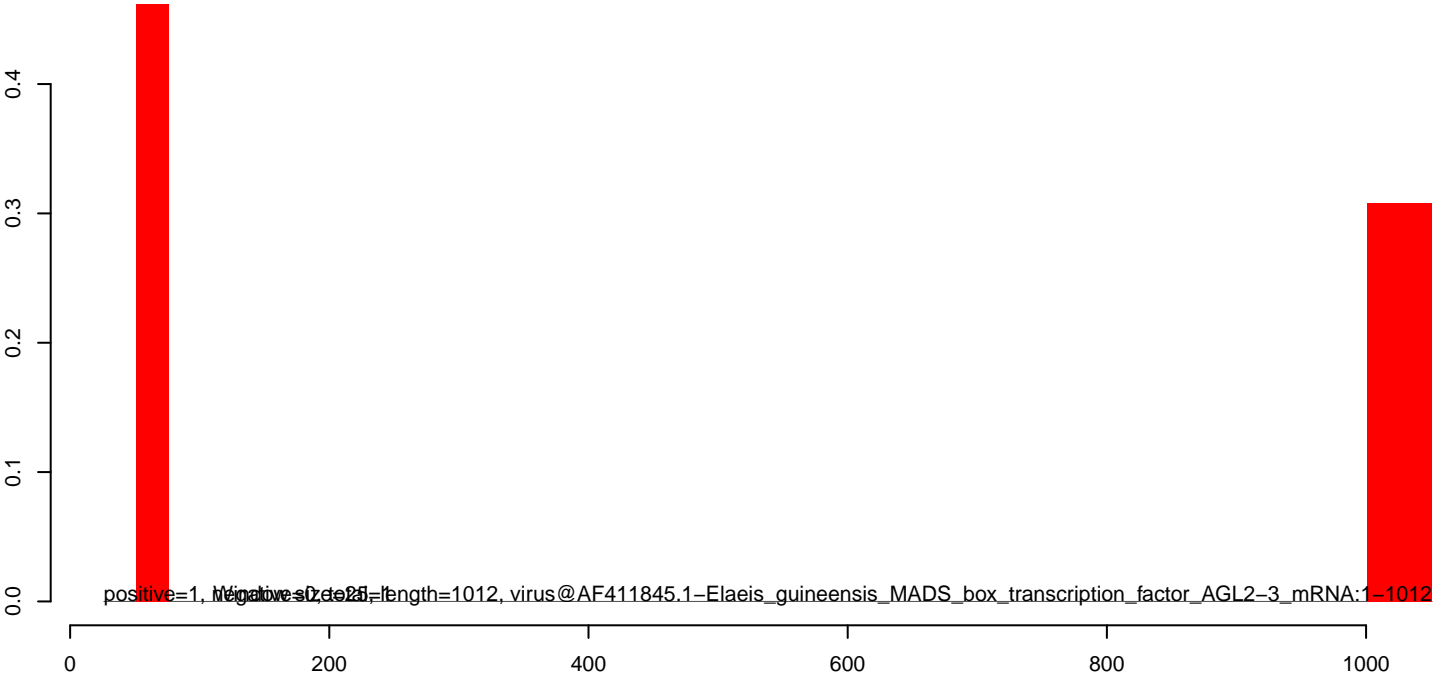
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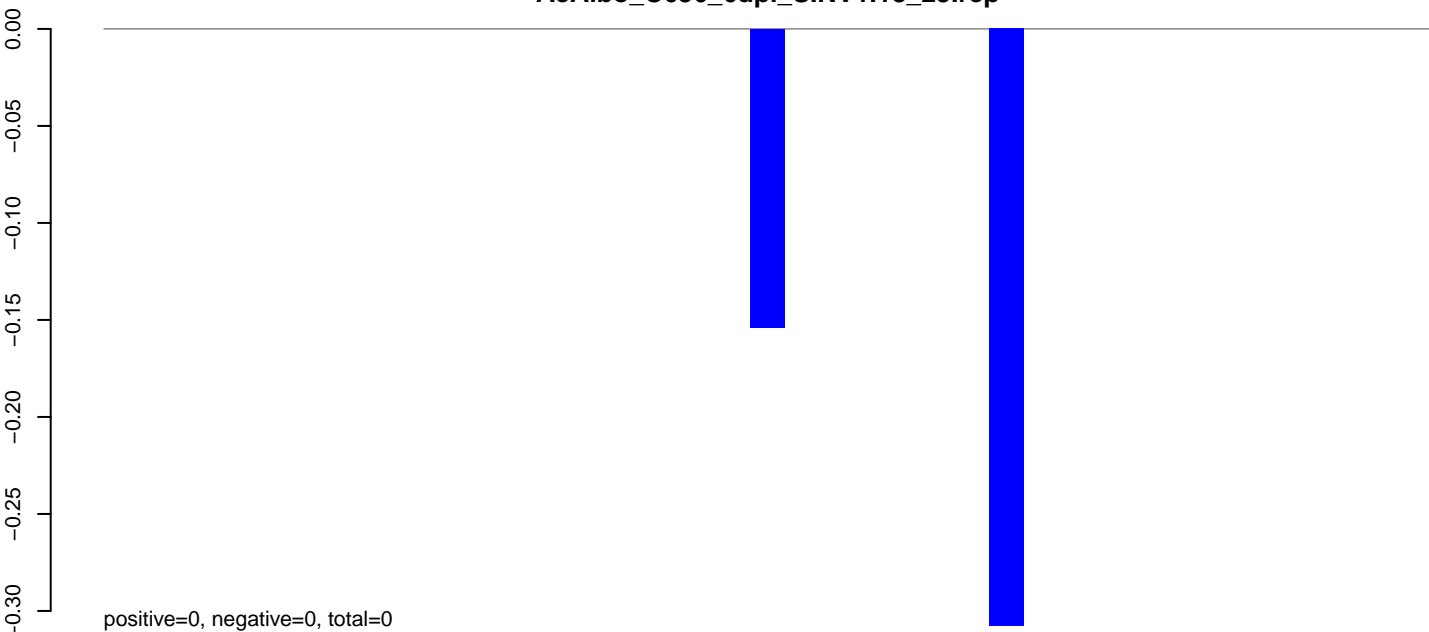
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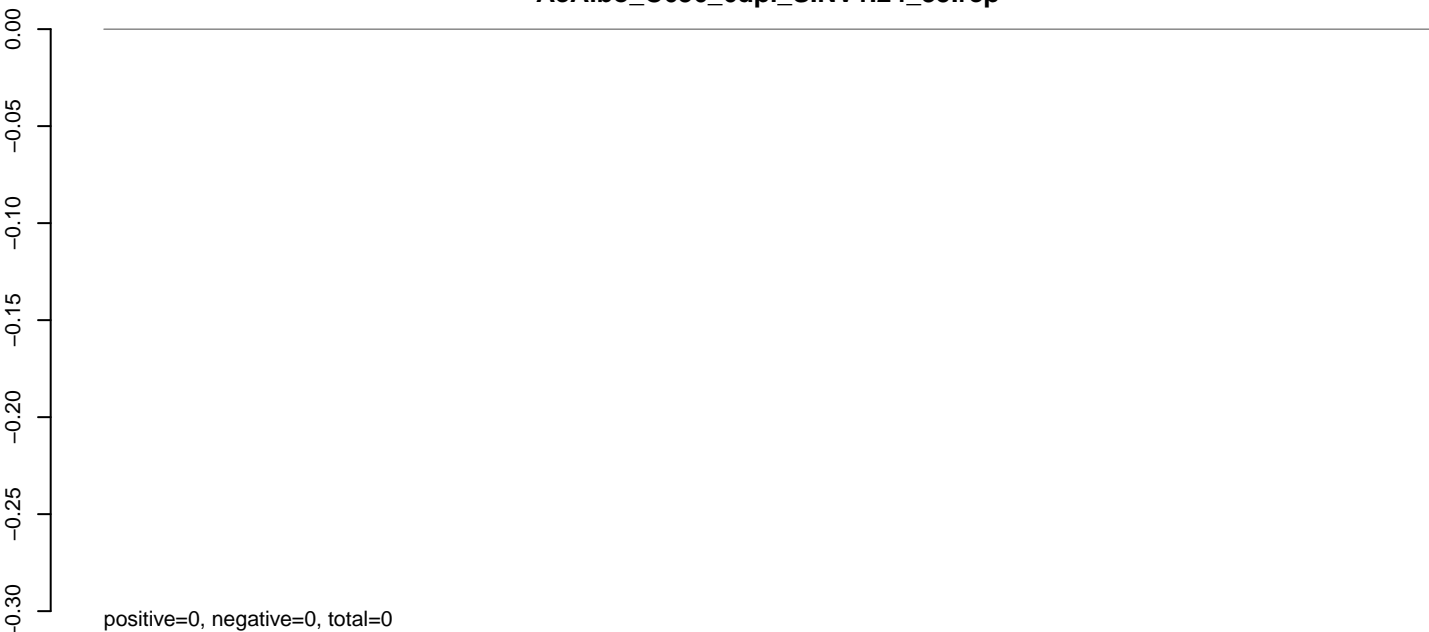
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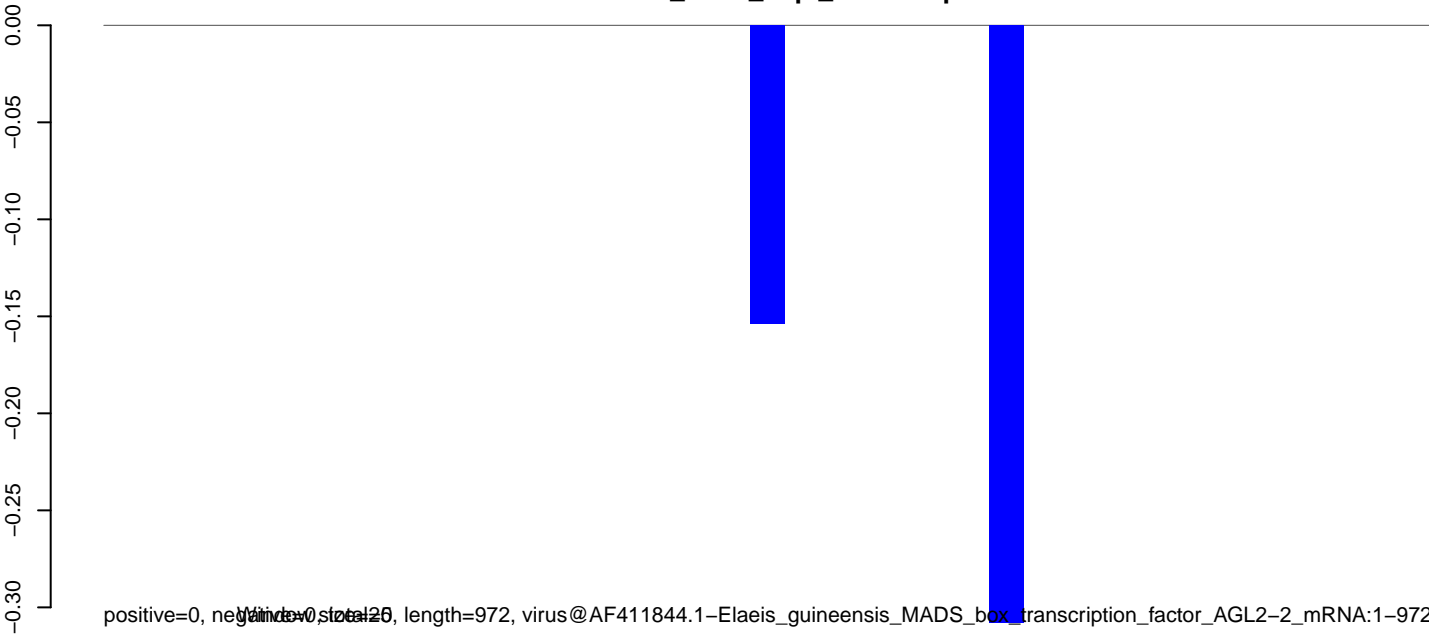
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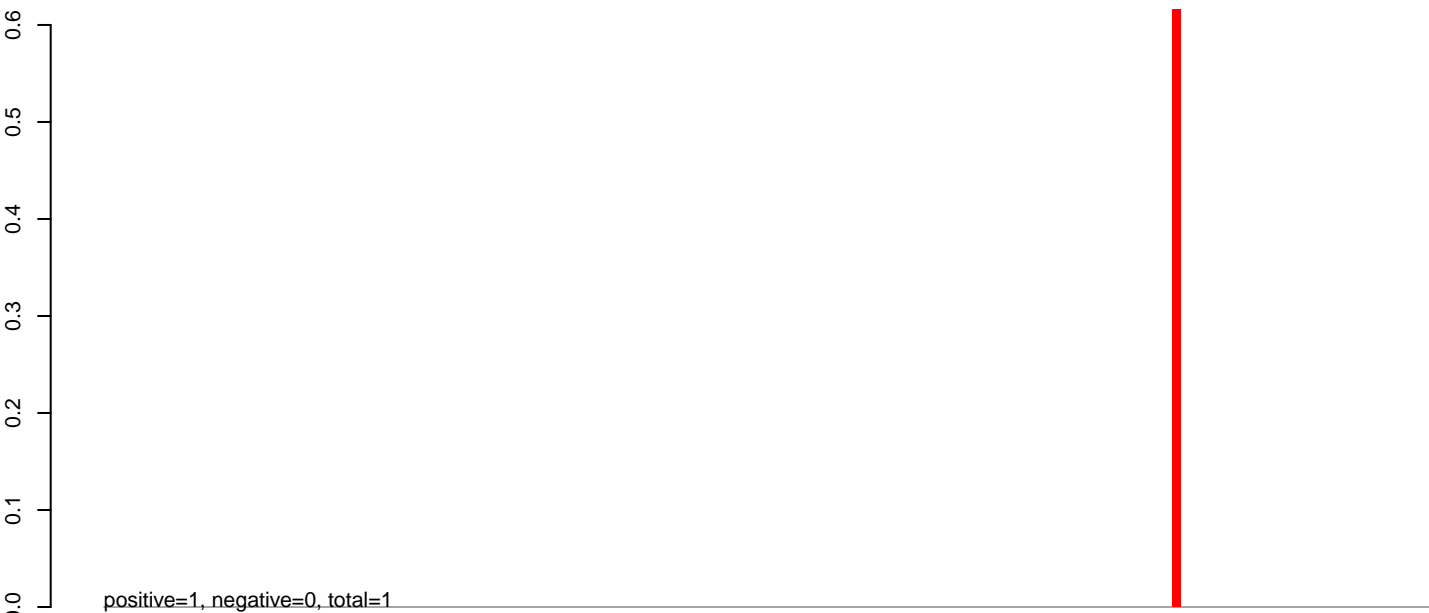
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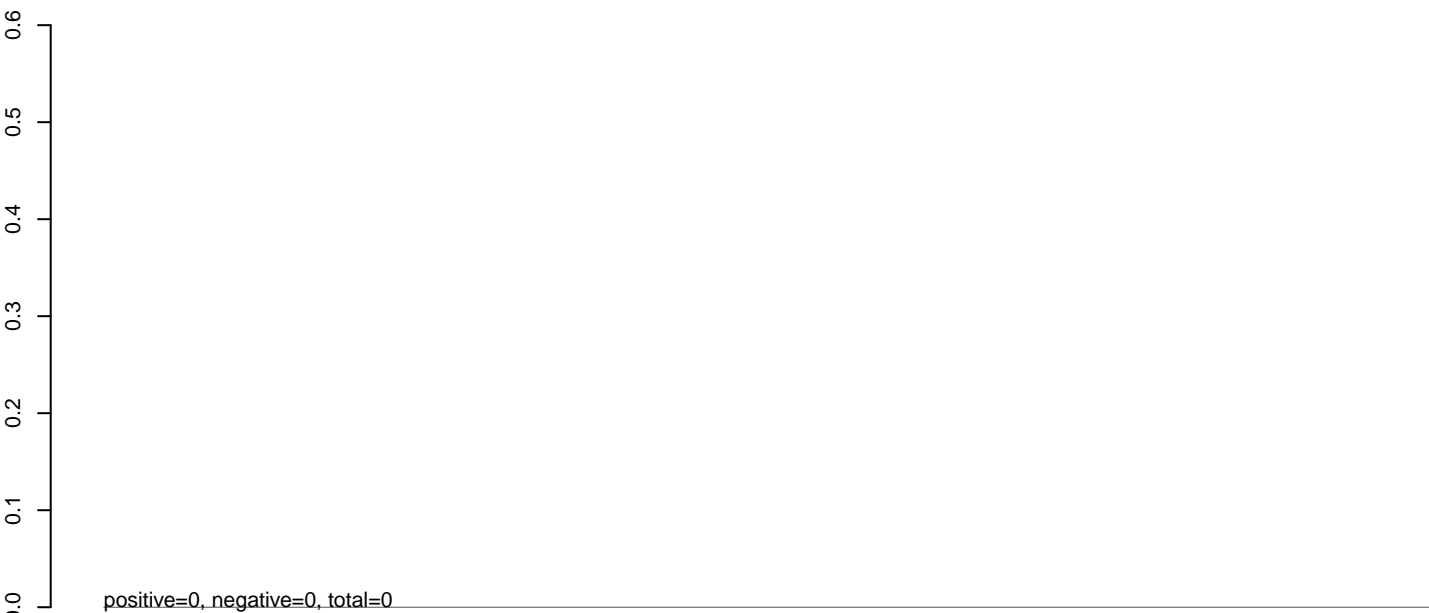
positive=0, negative=0, total=0, window_size=26, length=972, virus@AF411844.1-Elaeis_guineensis_MADS_box_transcription_factor_AGL2-2_mRNA:1-972

0 200 400 600 800 1000

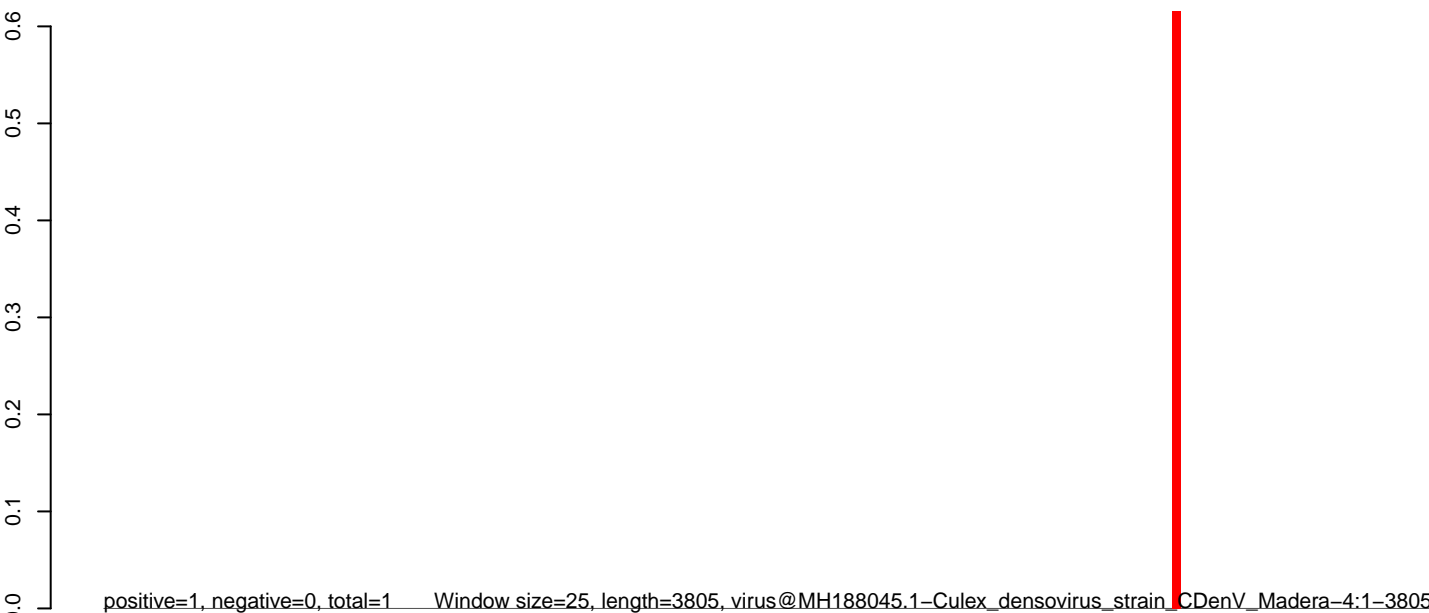
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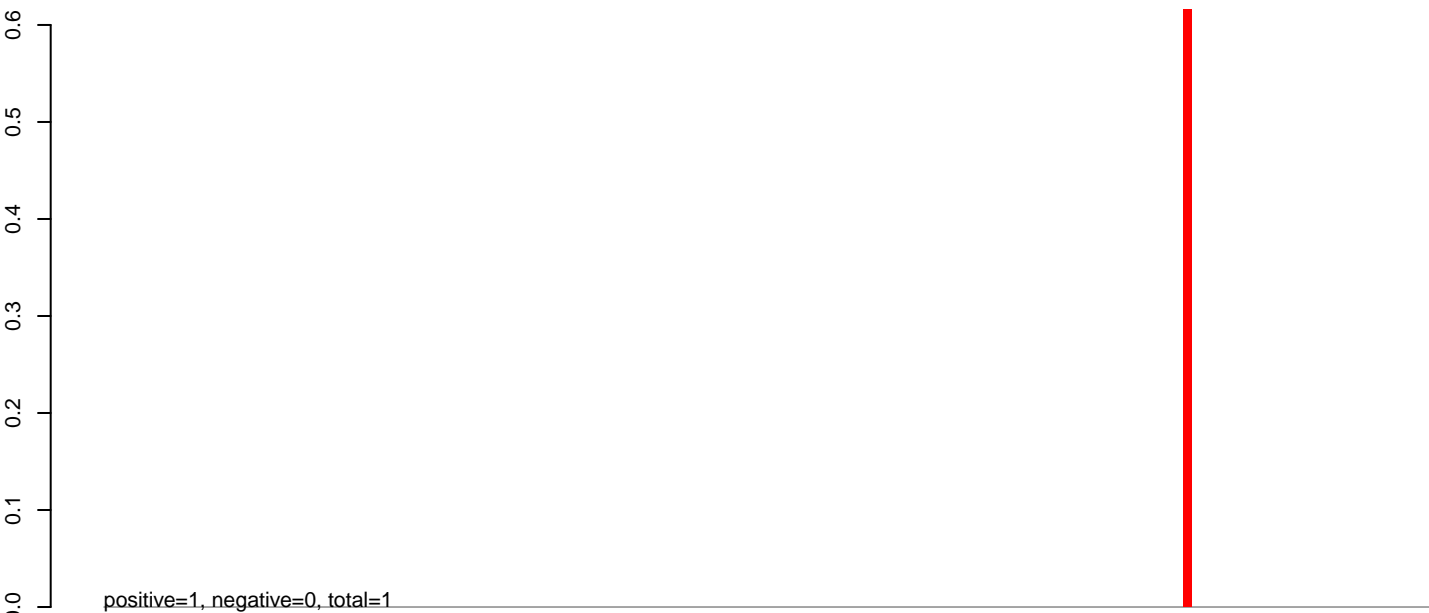


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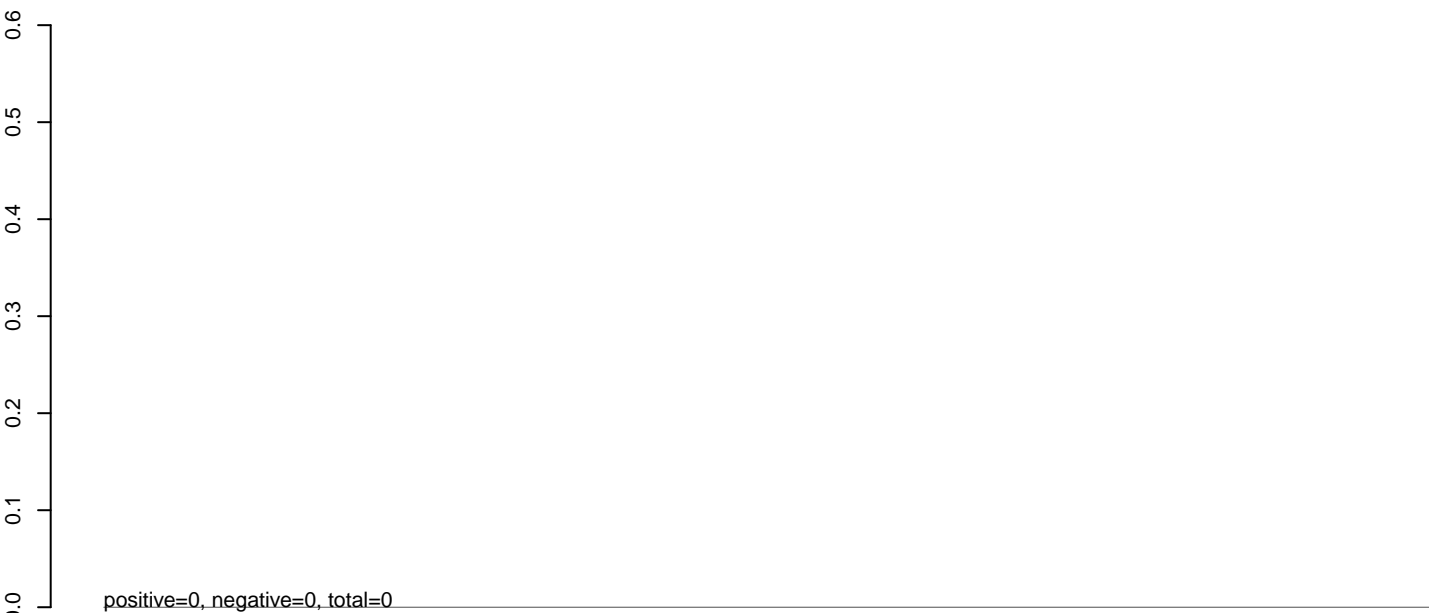


0 1000 2000 3000 4000

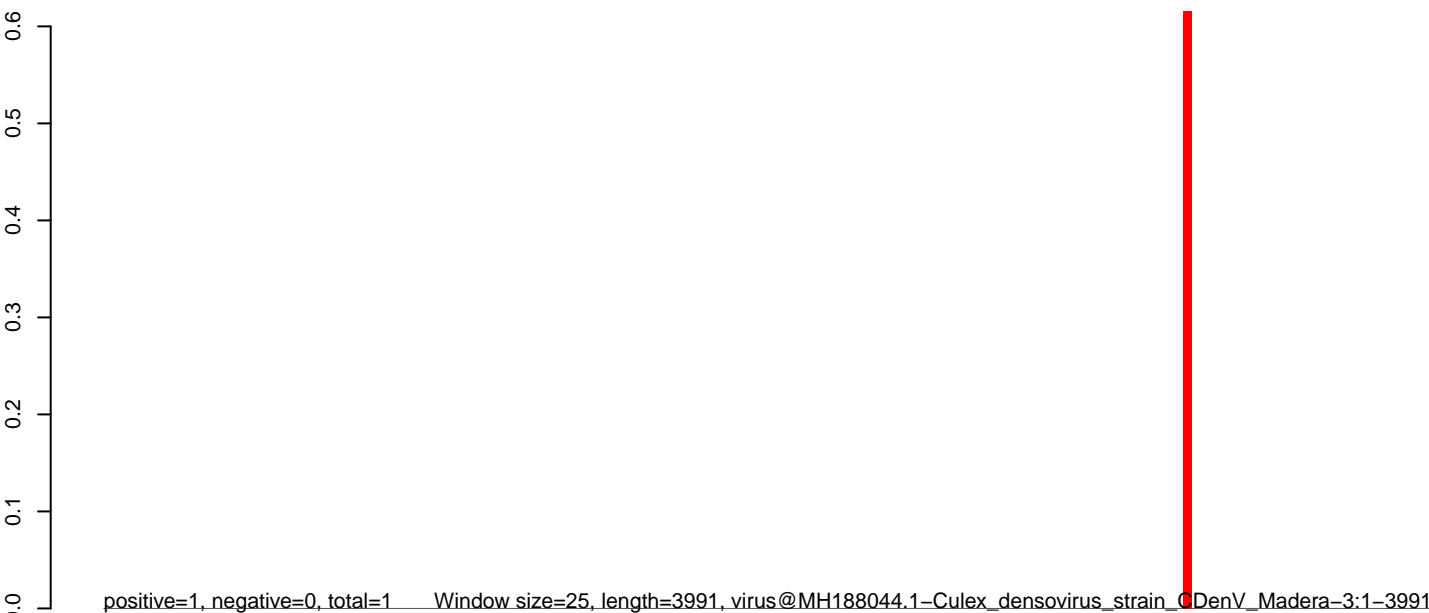
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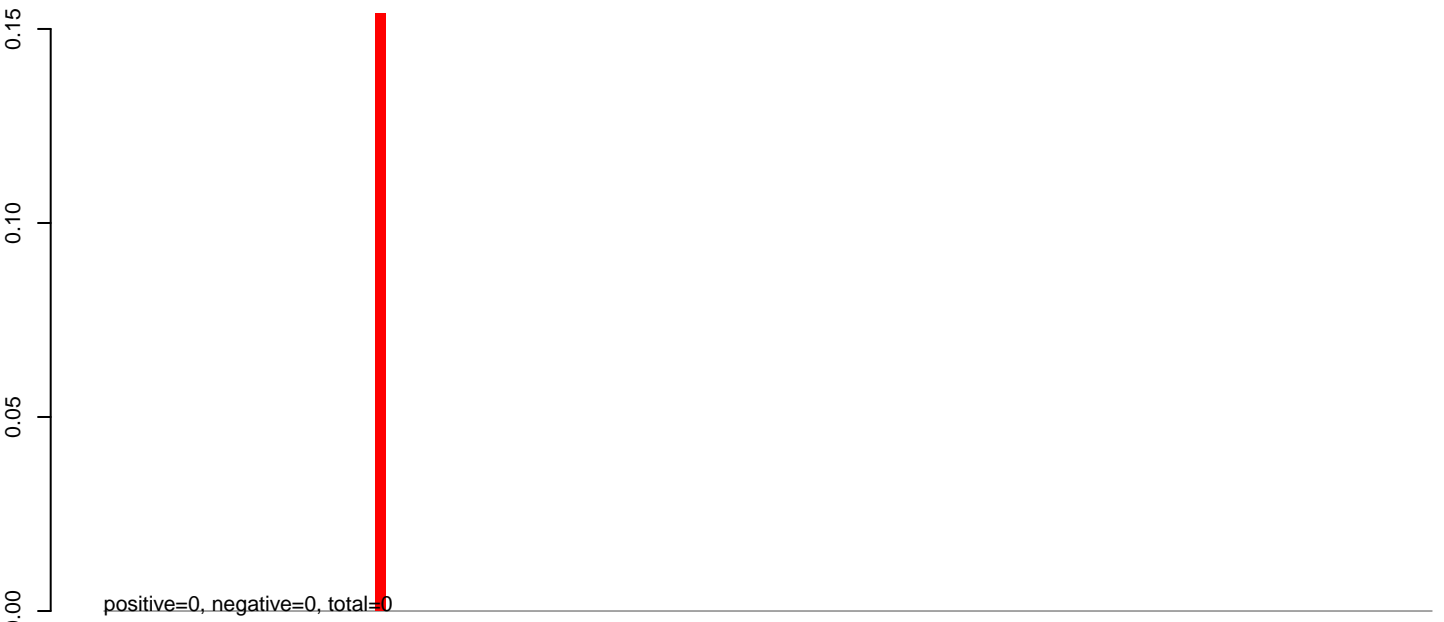
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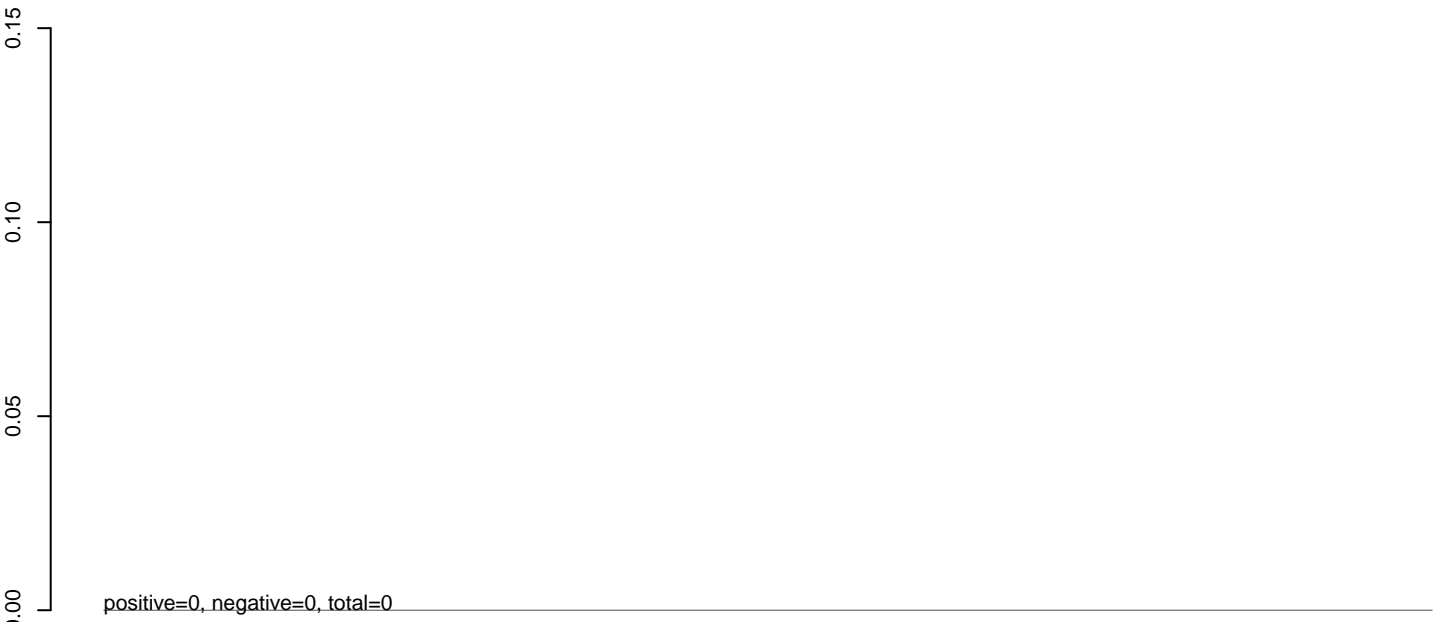
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AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



AeAlbo_C636_6dpi_SINV1.rep



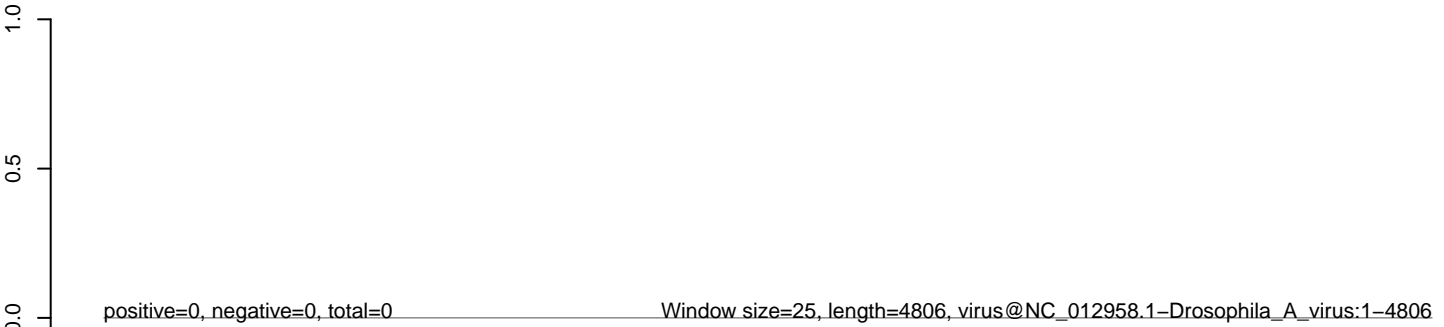
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AeAlbo_C636_6dpi_SINV1.rep



AeAlbo_C636_6dpi_SINV1.18_23.rep



AeAlbo_C636_6dpi_SINV1.24_35.rep



AeAlbo_C636_6dpi_SINV1.rep

