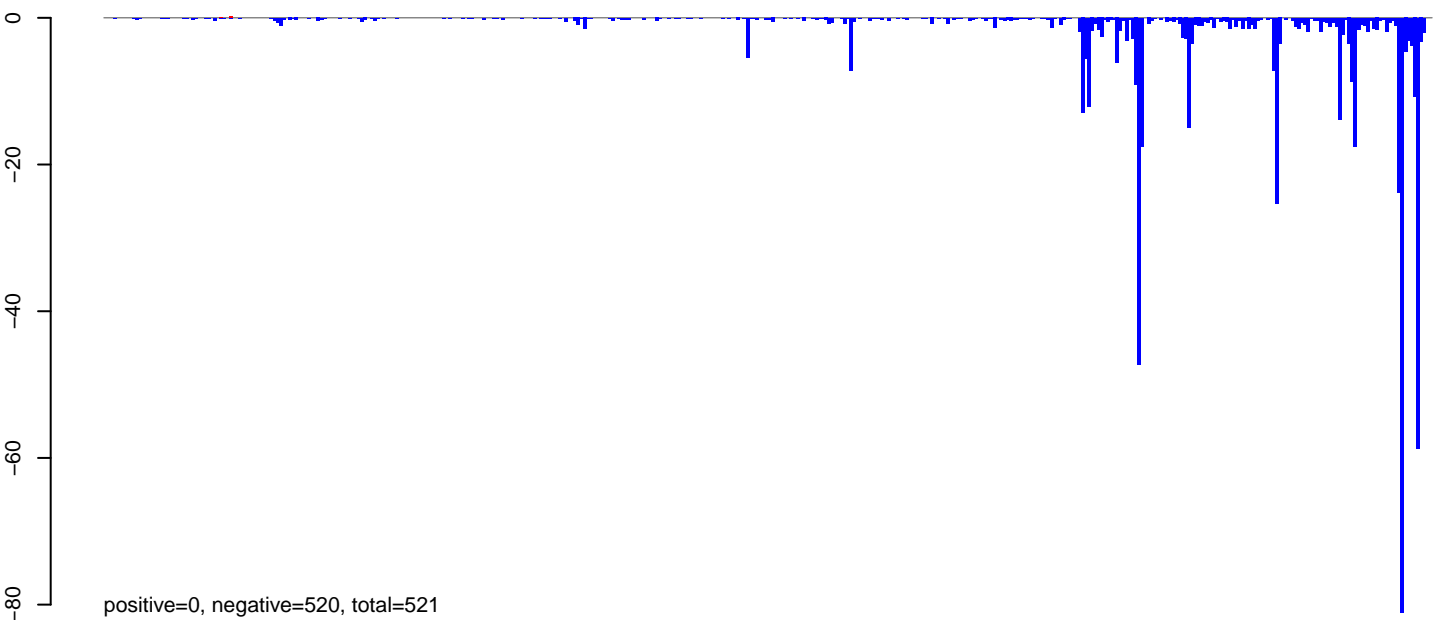


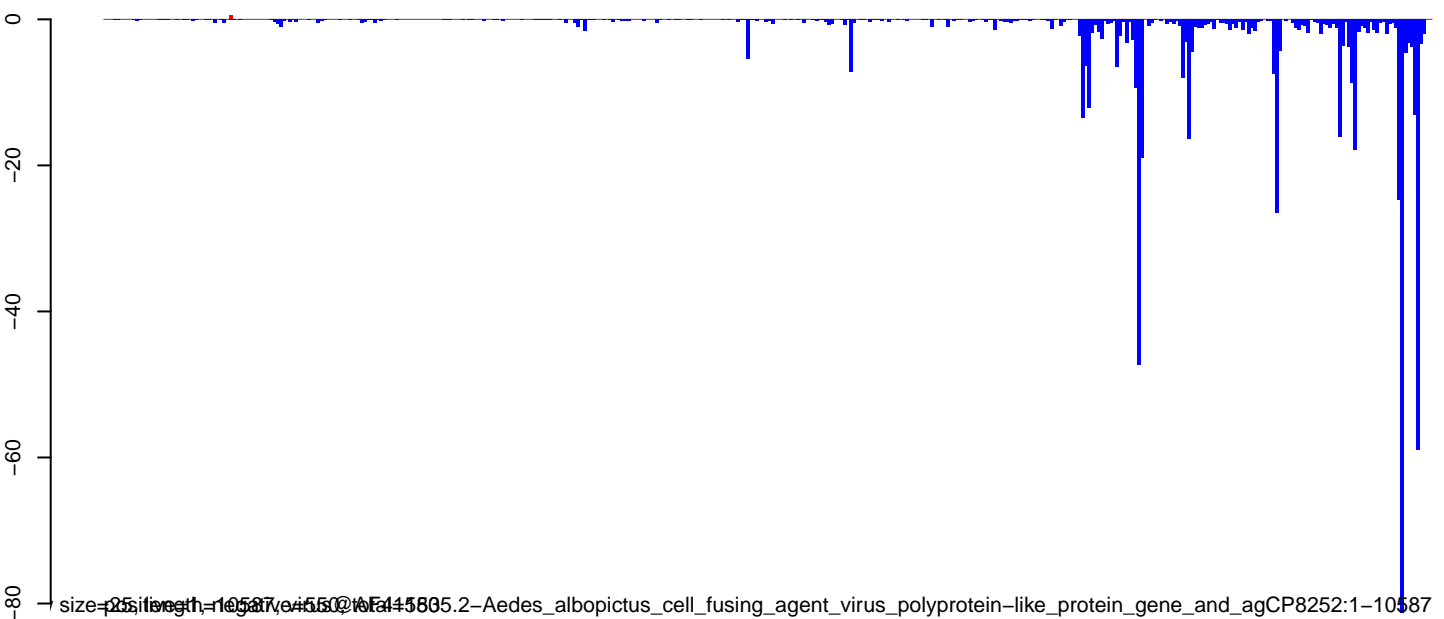
AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep

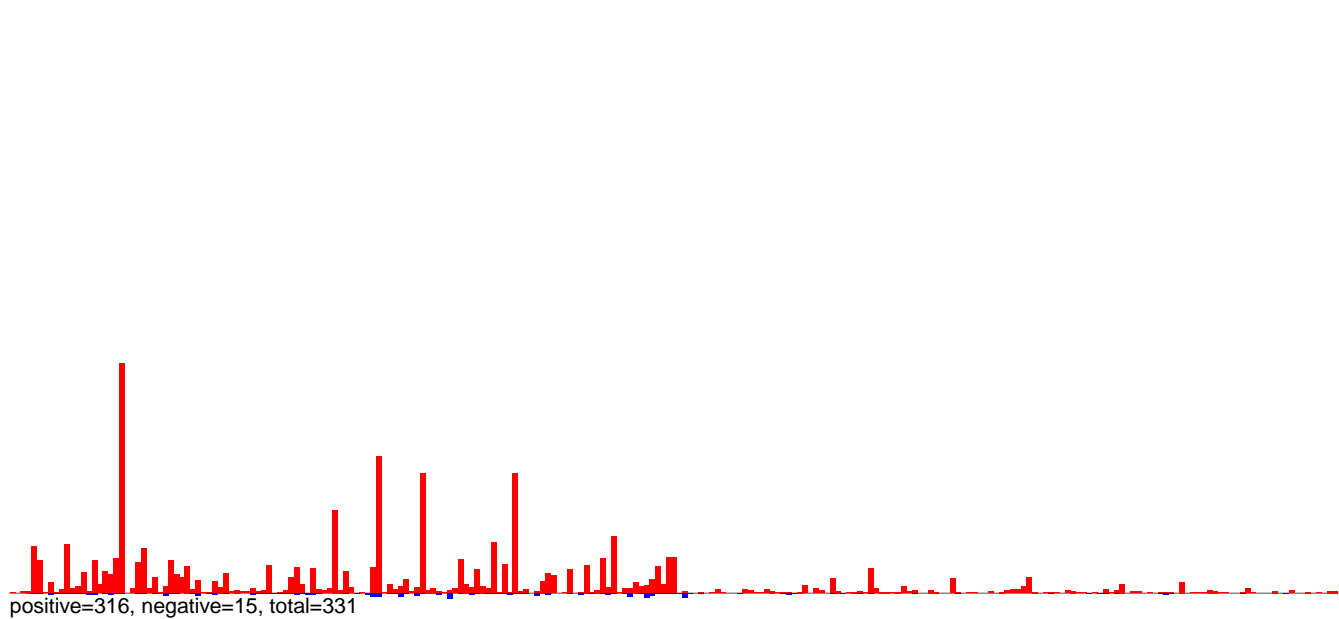


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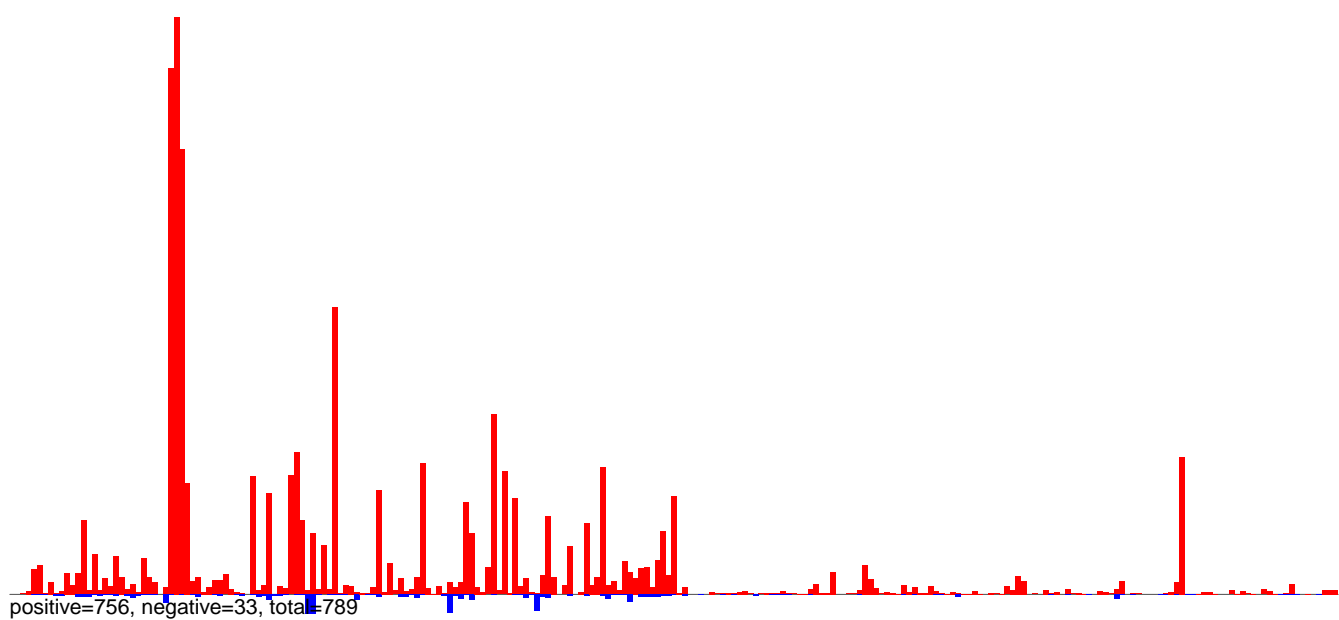


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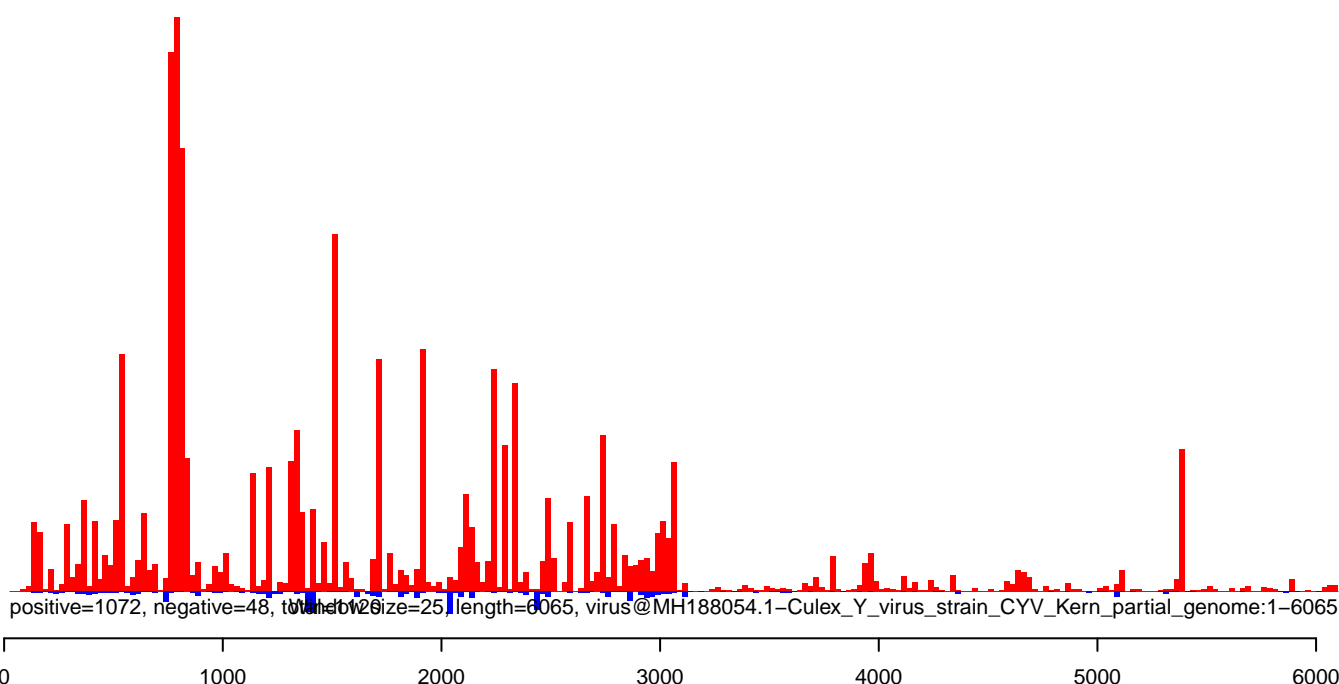
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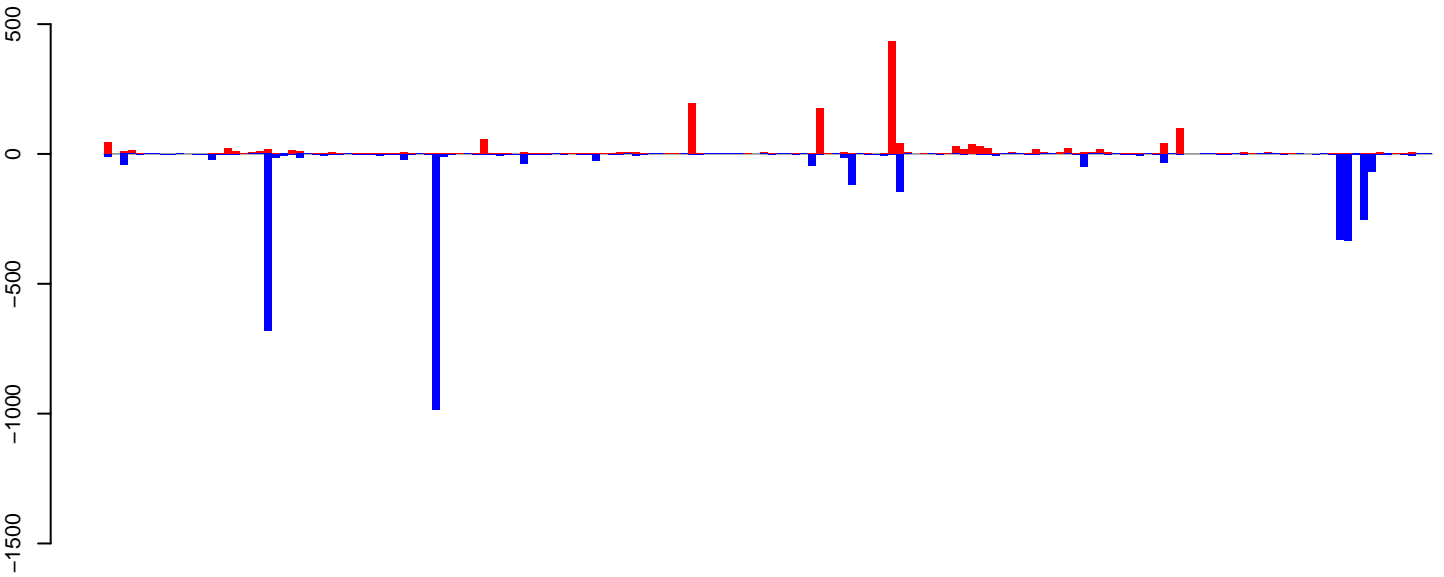
AeAlbo_C636_B2_FHV_r54q.24_35.rep



AeAlbo_C636_B2_FHV_r54q.rep

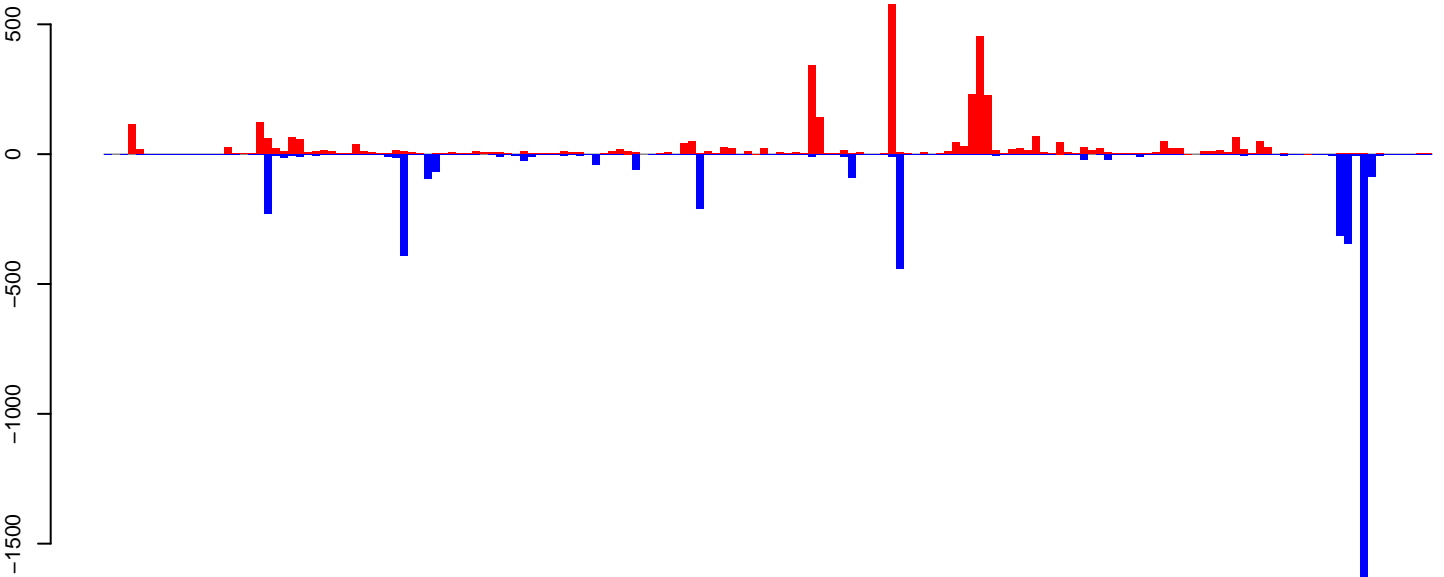


AeAlbo_C636_B2_FHV_r54q.18_23.rep



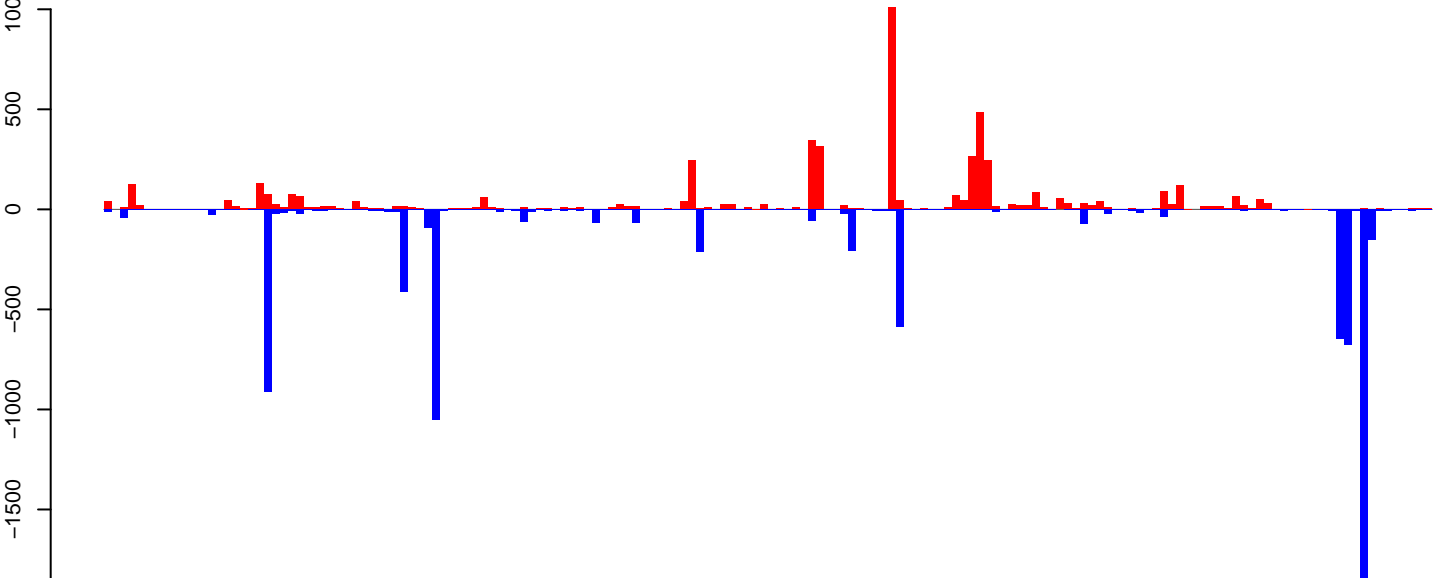
positive=1596, negative=3388, total=4984

AeAlbo_C636_B2_FHV_r54q.24_35.rep



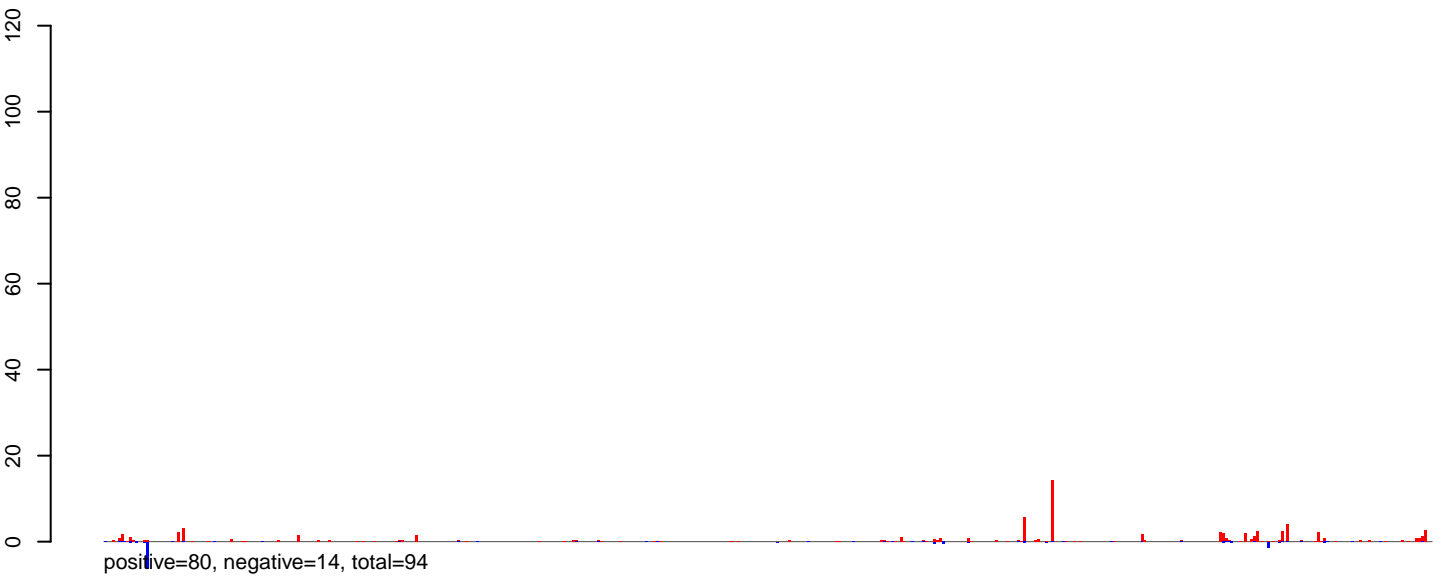
positive=3766, negative=4451, total=8217

AeAlbo_C636_B2_FHV_r54q.rep

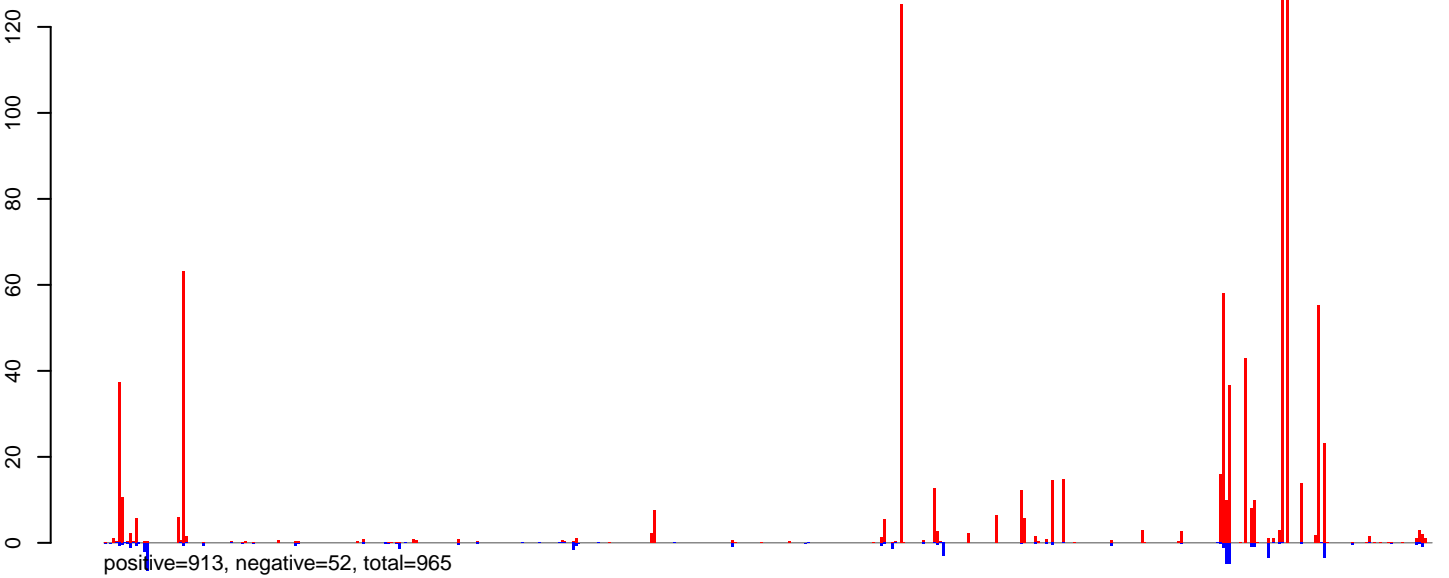


positive=5361, negative=7839, total=13199, size=25, length=4139, virus@NC_011317.1-Anopheles_gambiae_densovirus_1-4139

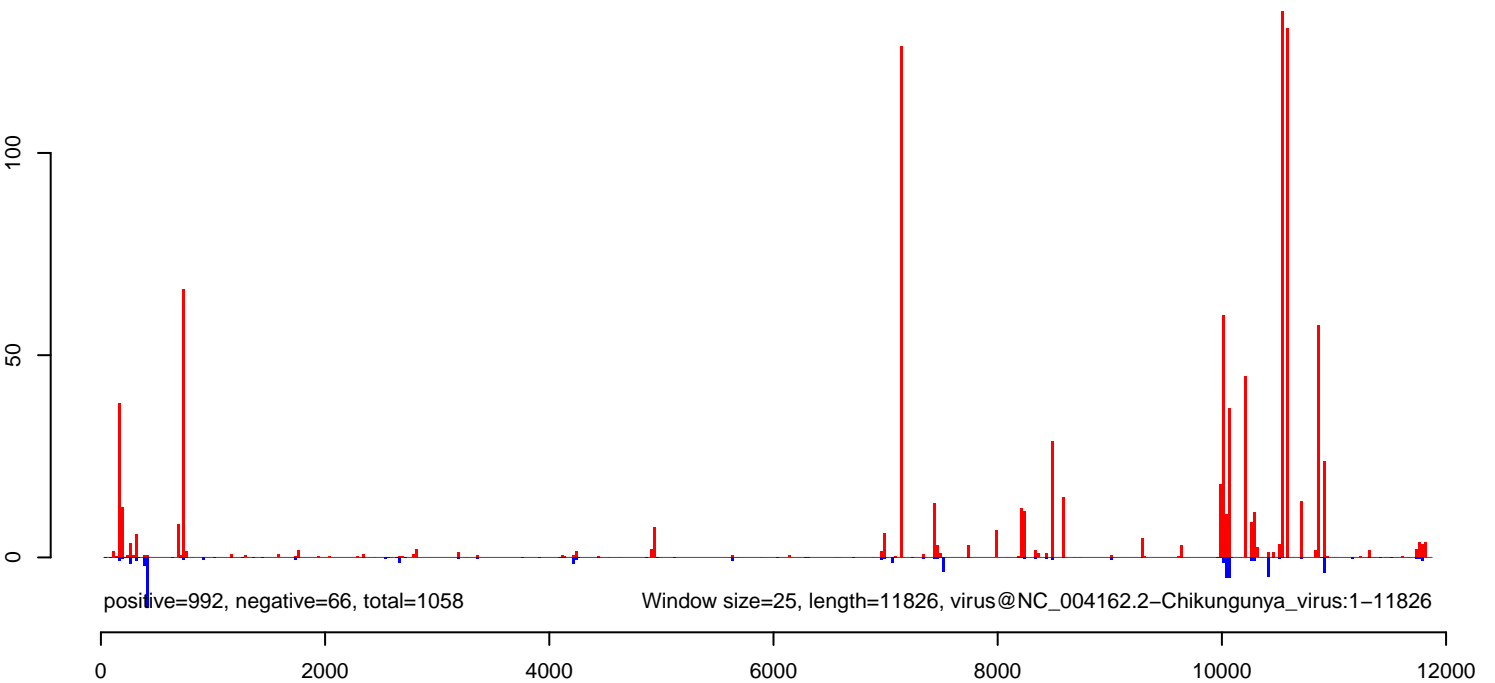
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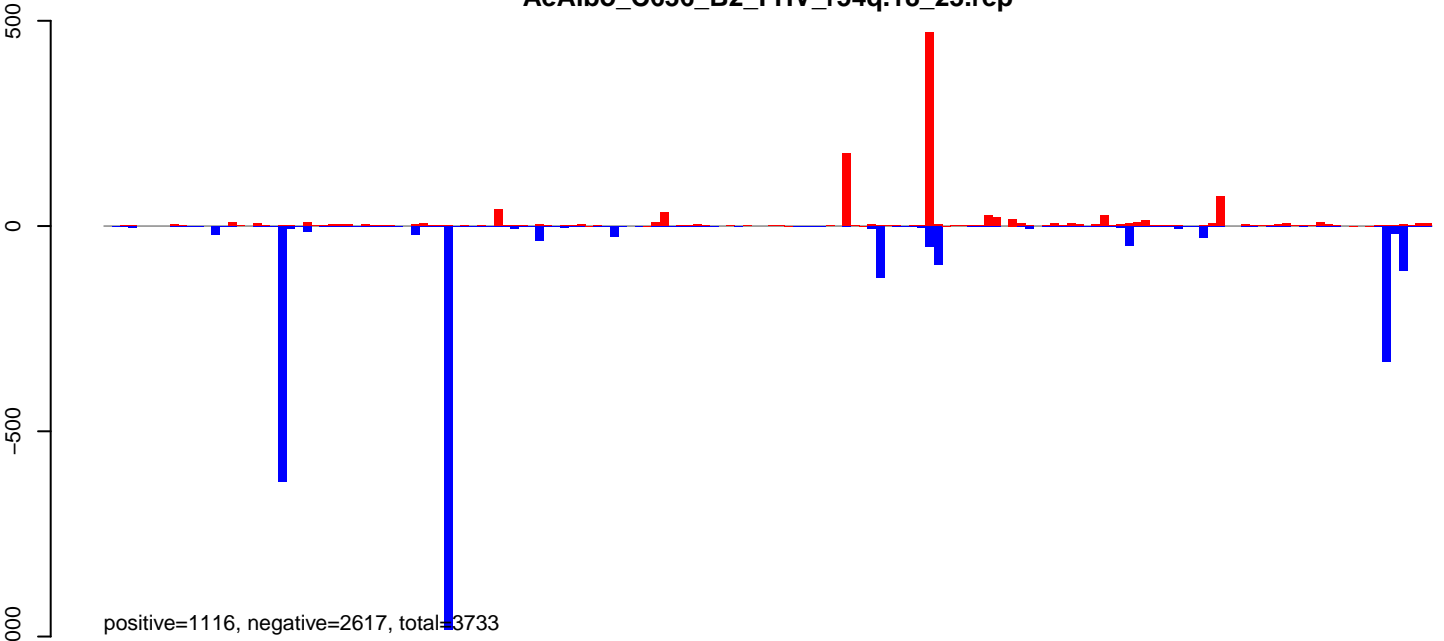
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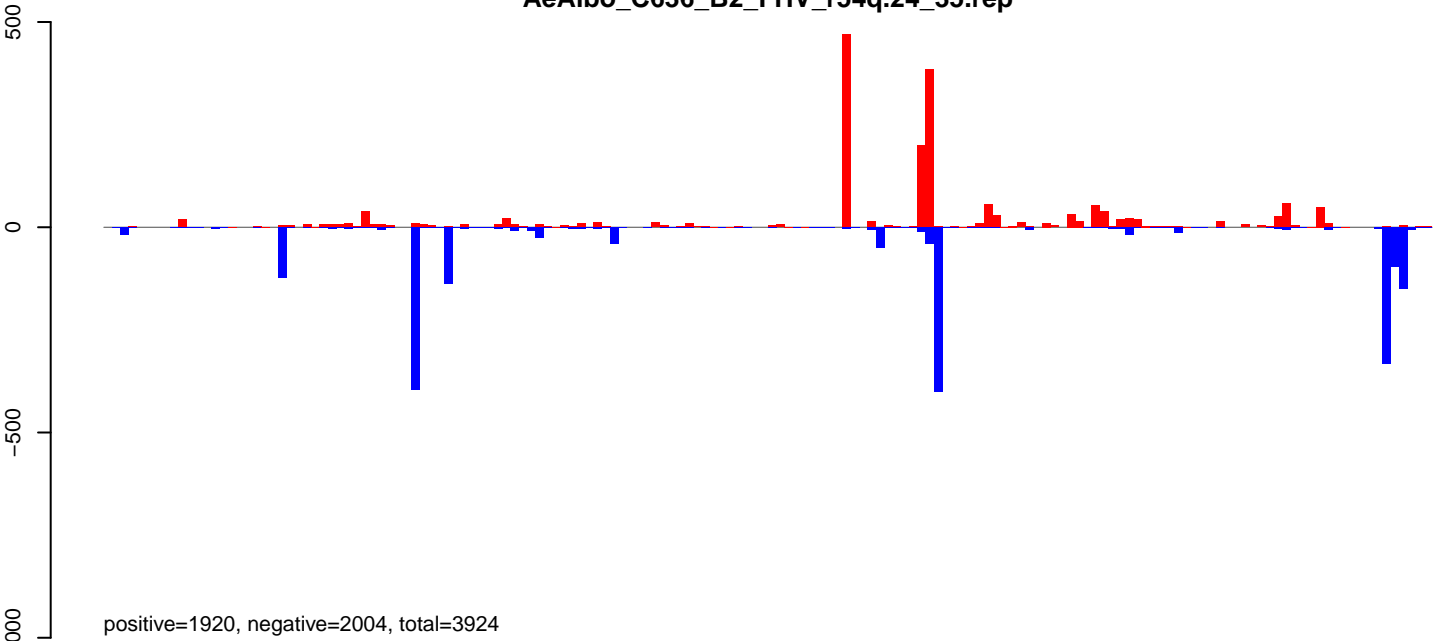
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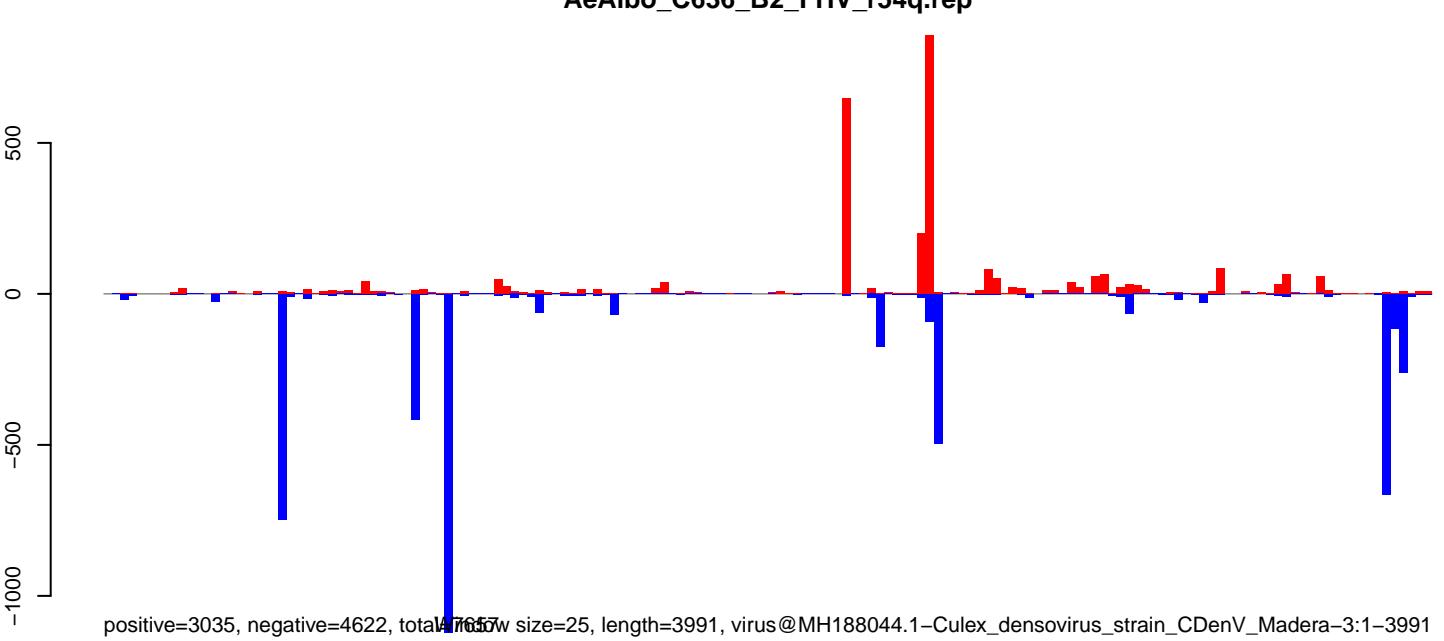
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AeAlbo_C636_B2_FHV_r54q.24_35.rep

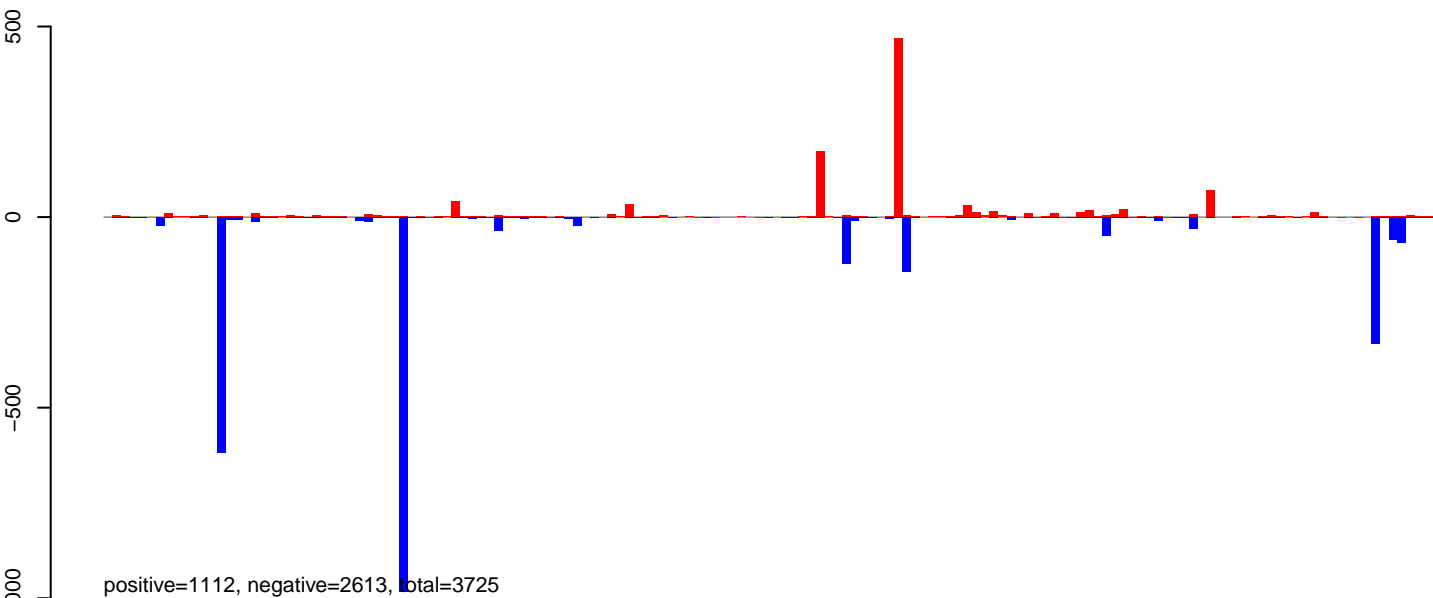


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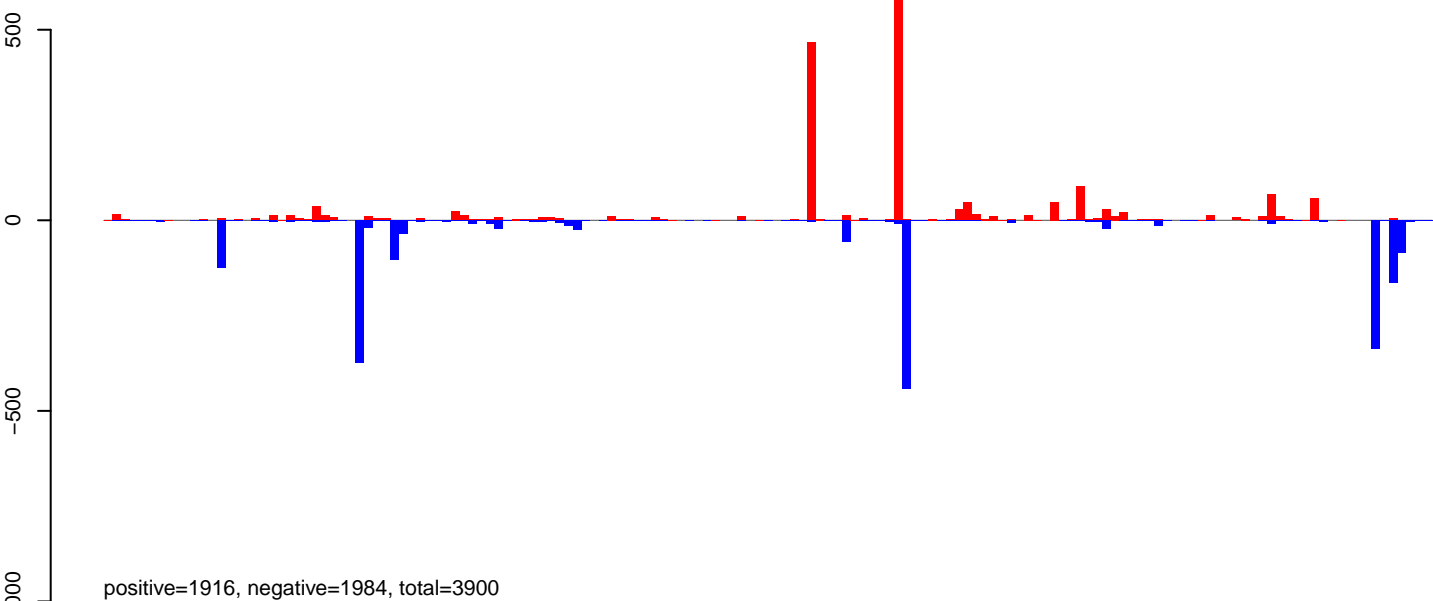


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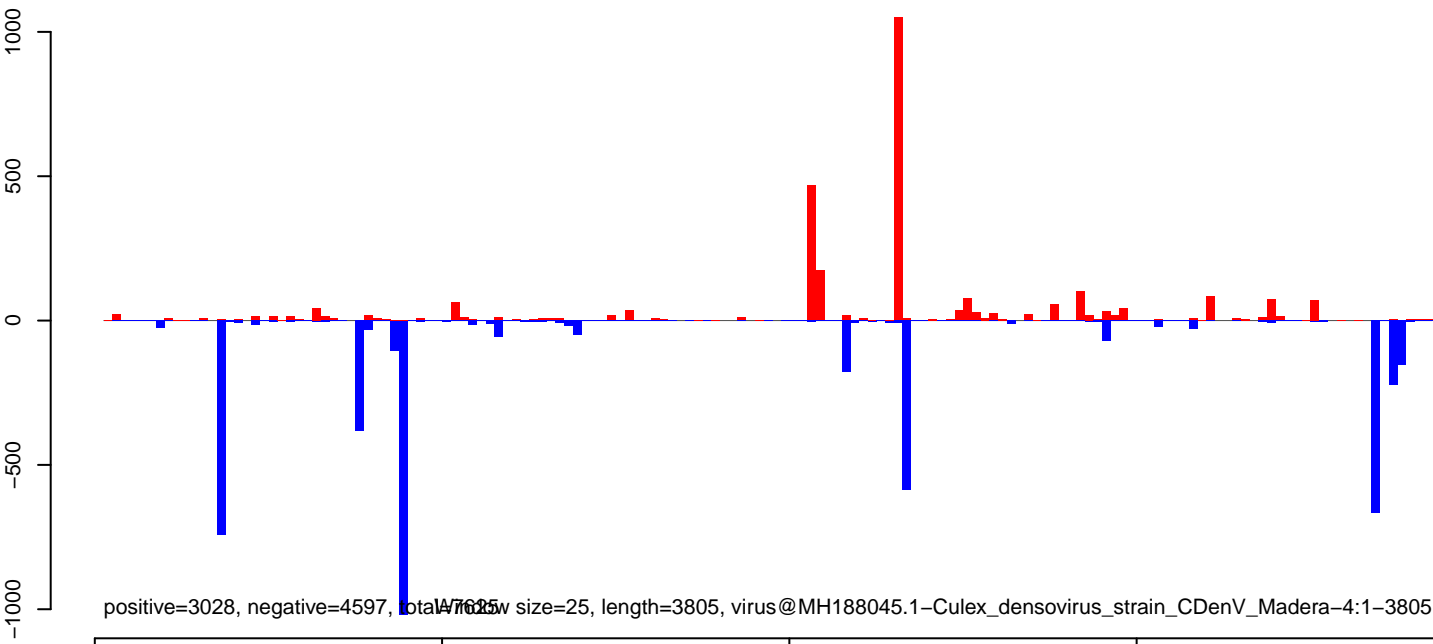
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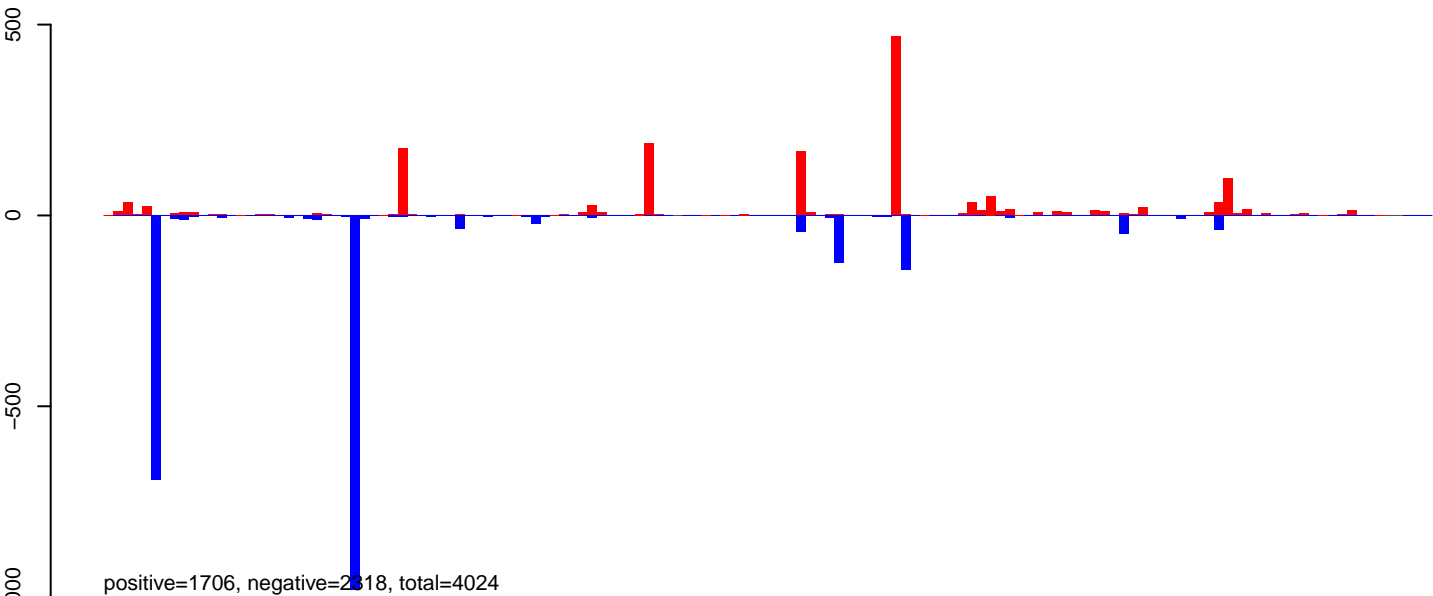
AeAlbo_C636_B2_FHV_r54q.24_35.rep



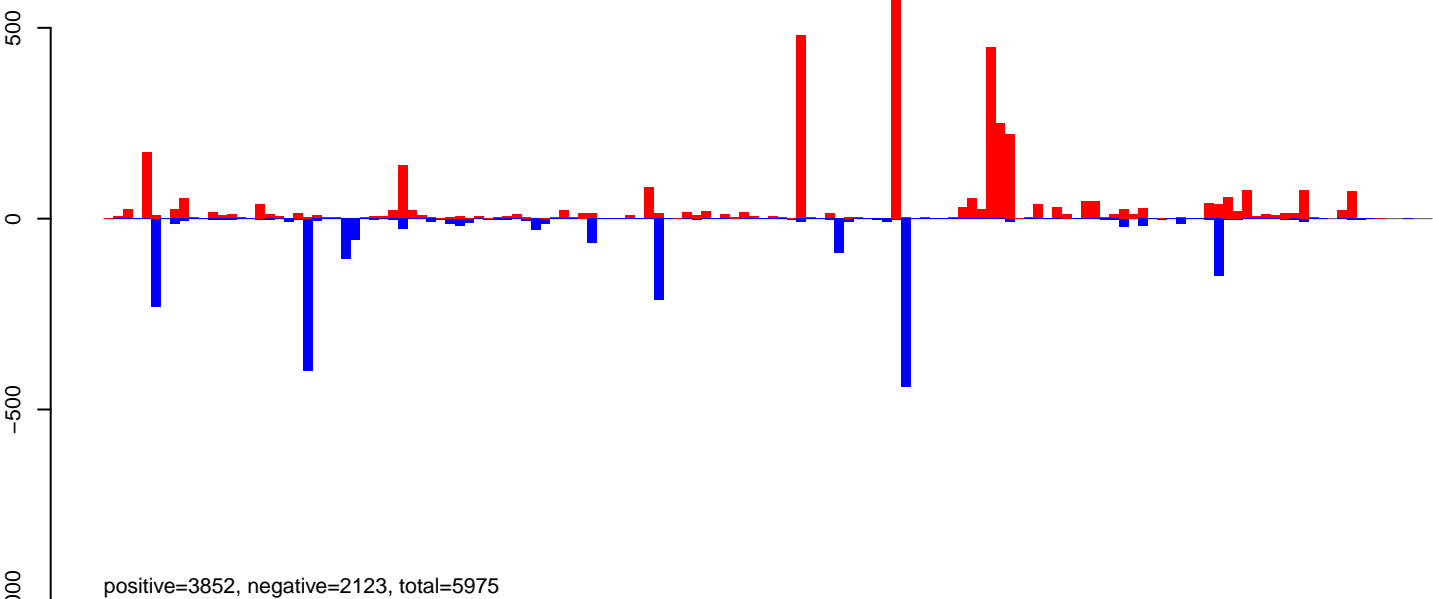
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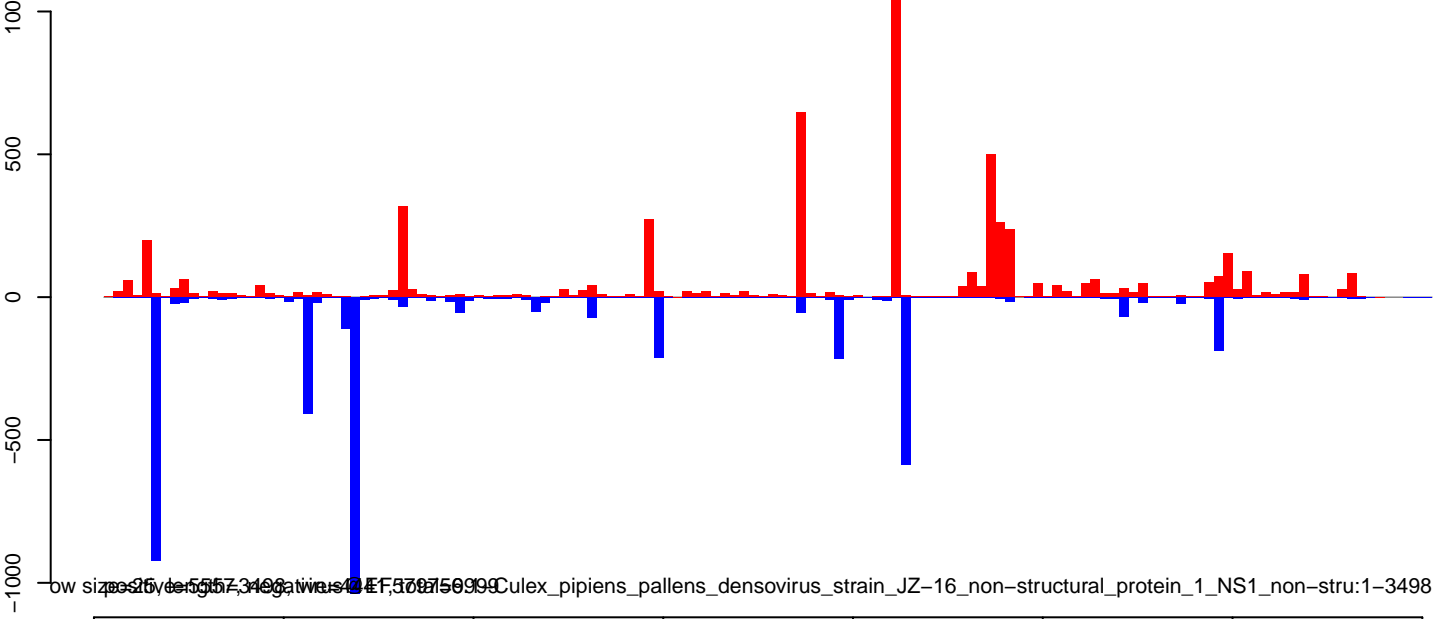
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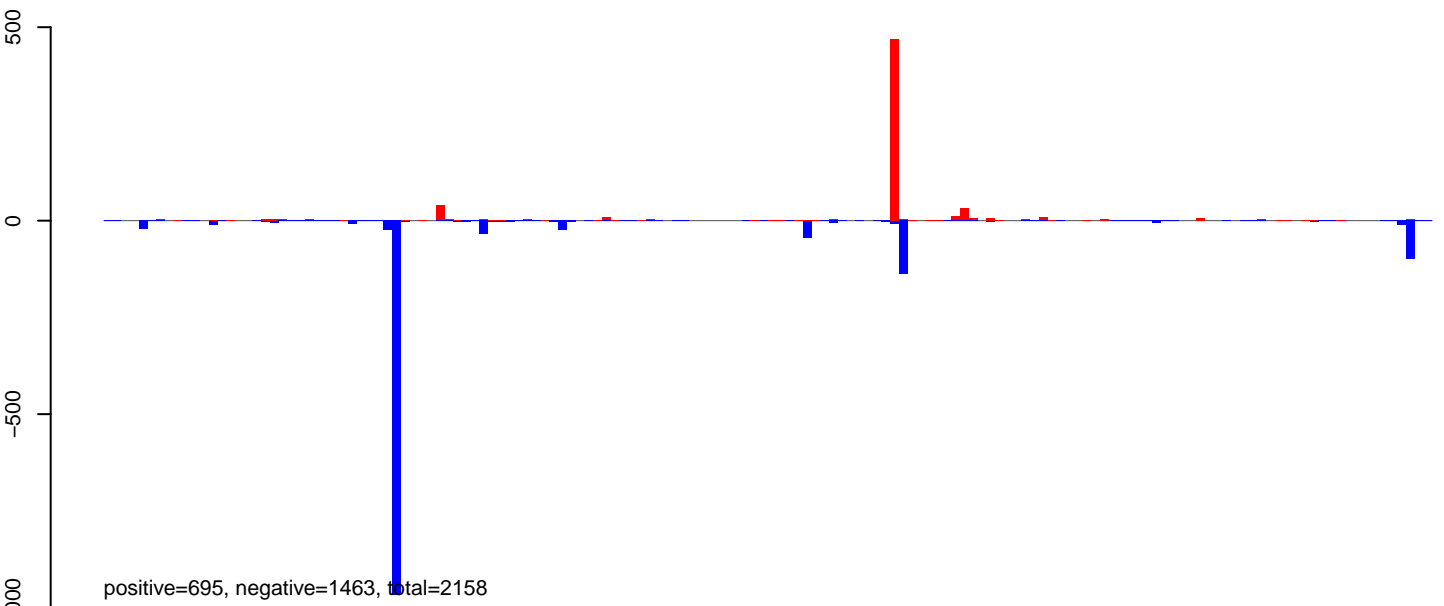
AeAlbo_C636_B2_FHV_r54q.24_35.rep



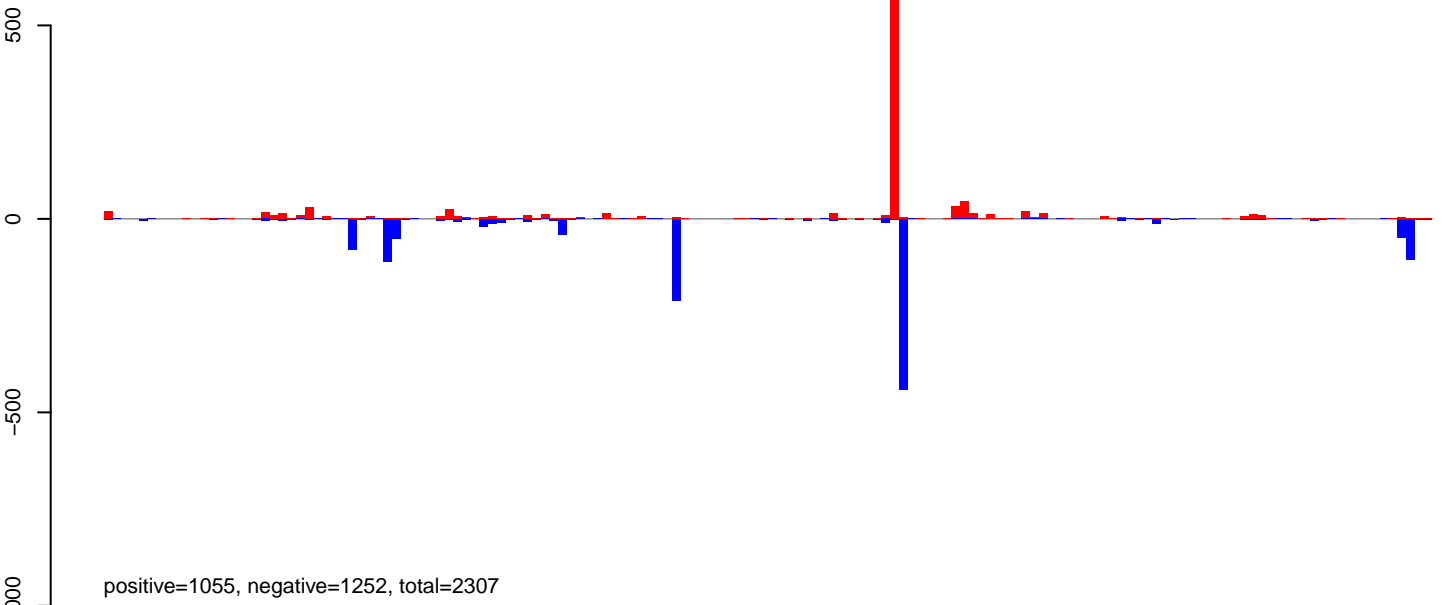
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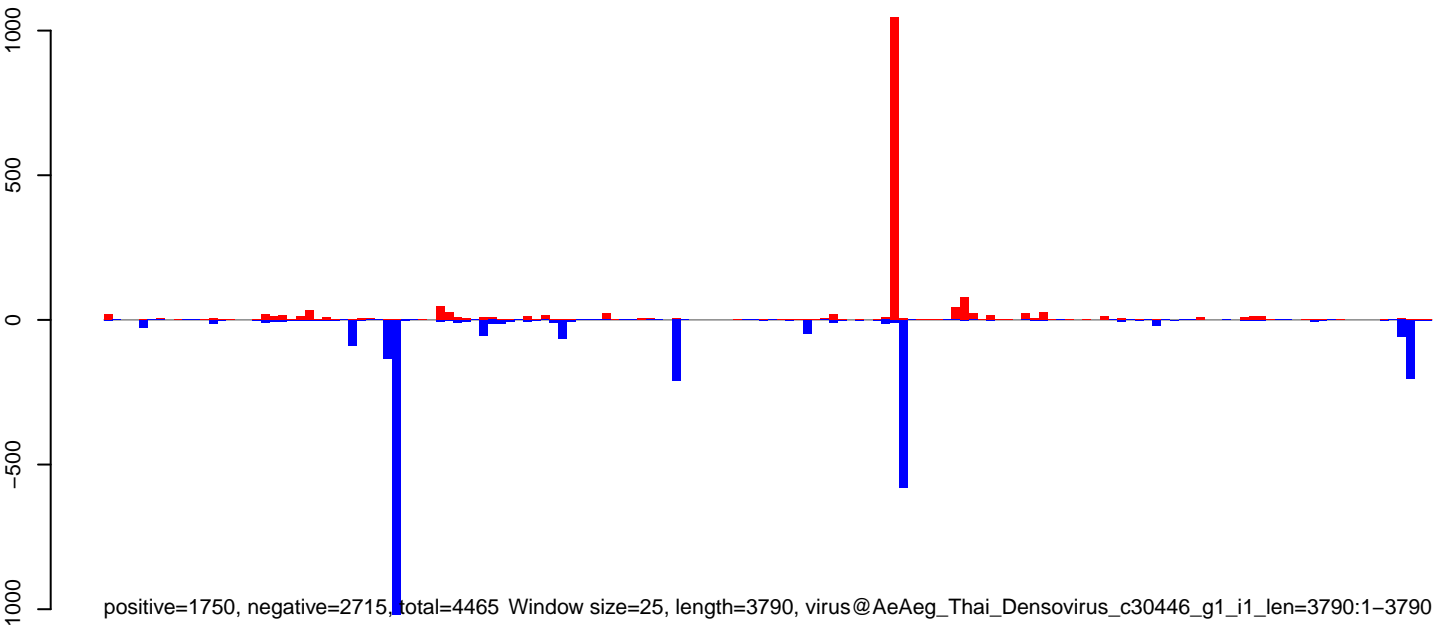
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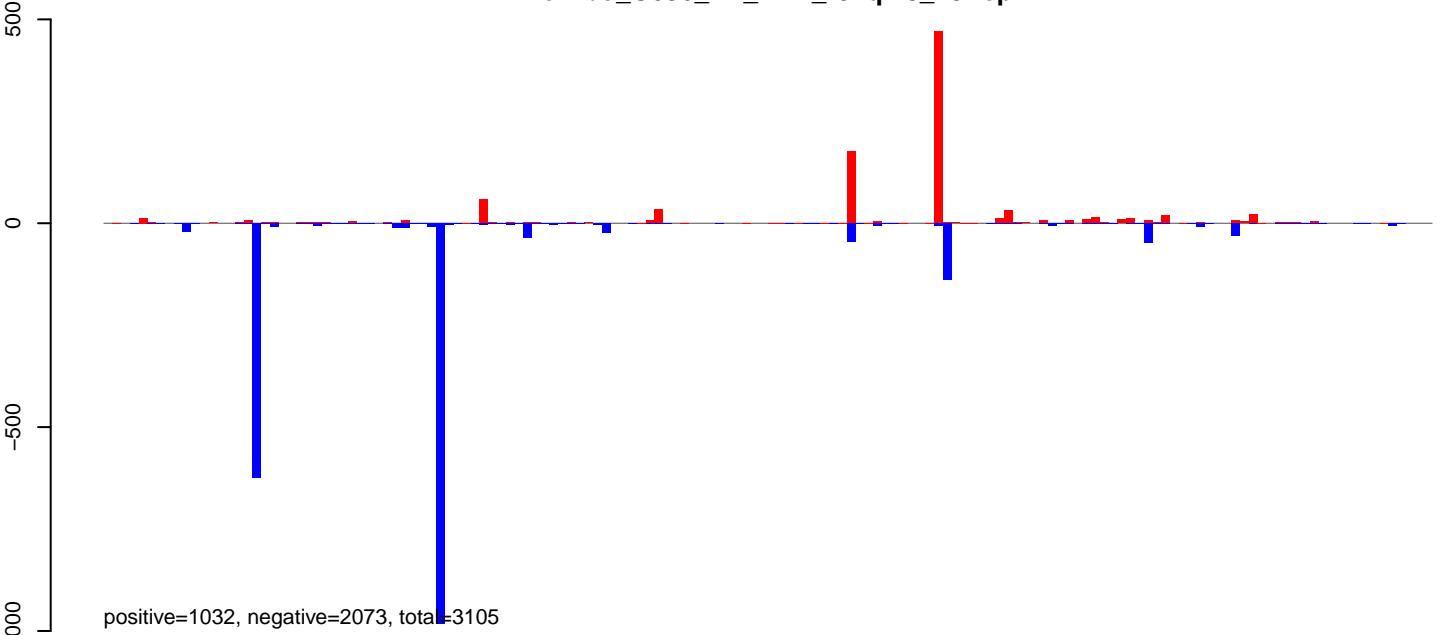
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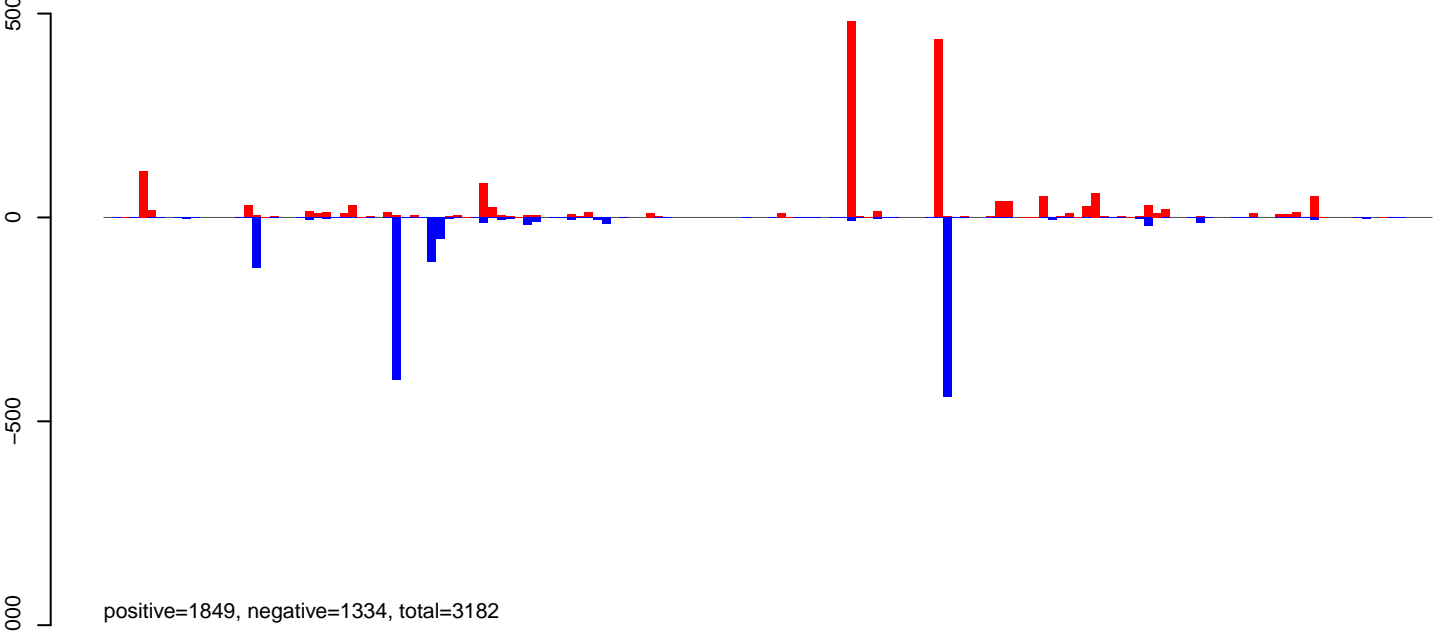
AeAlbo_C636_B2_FHV_r54q.rep



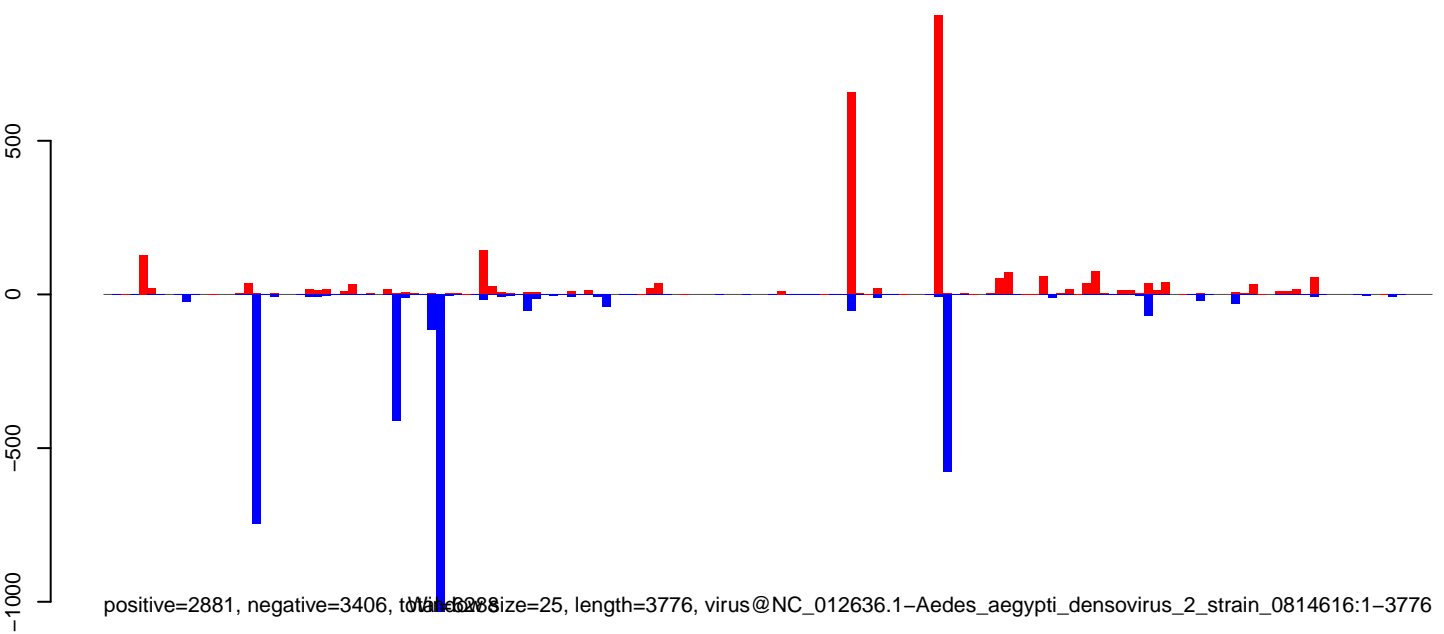
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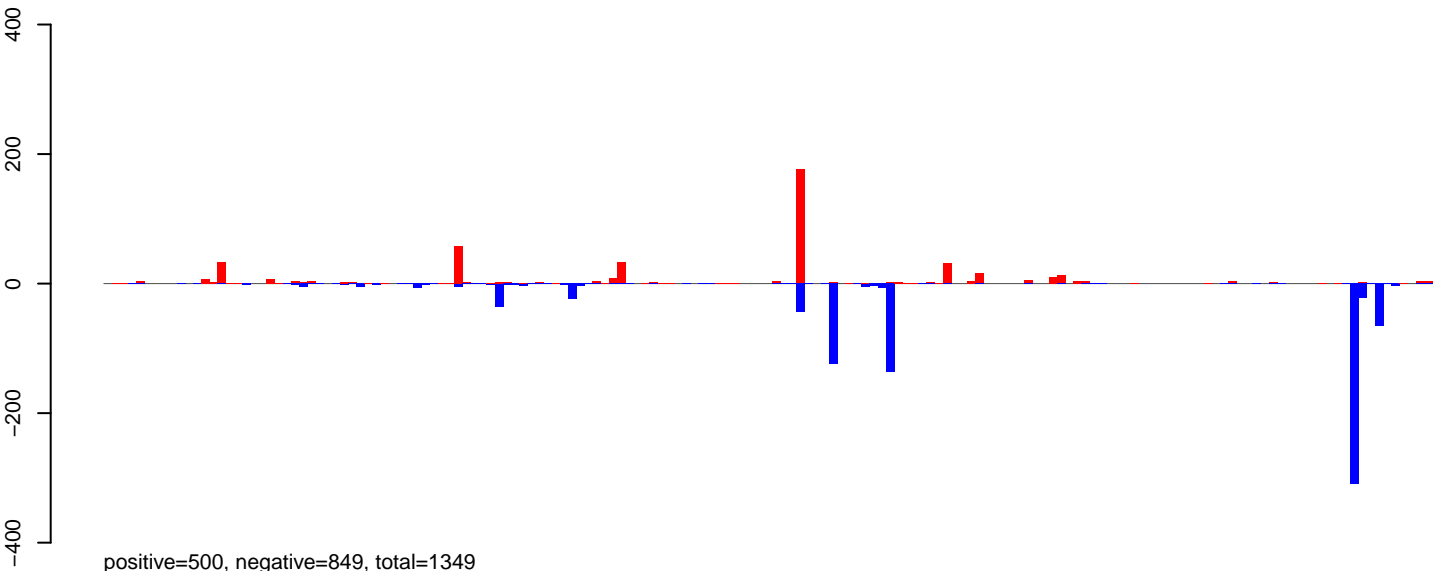
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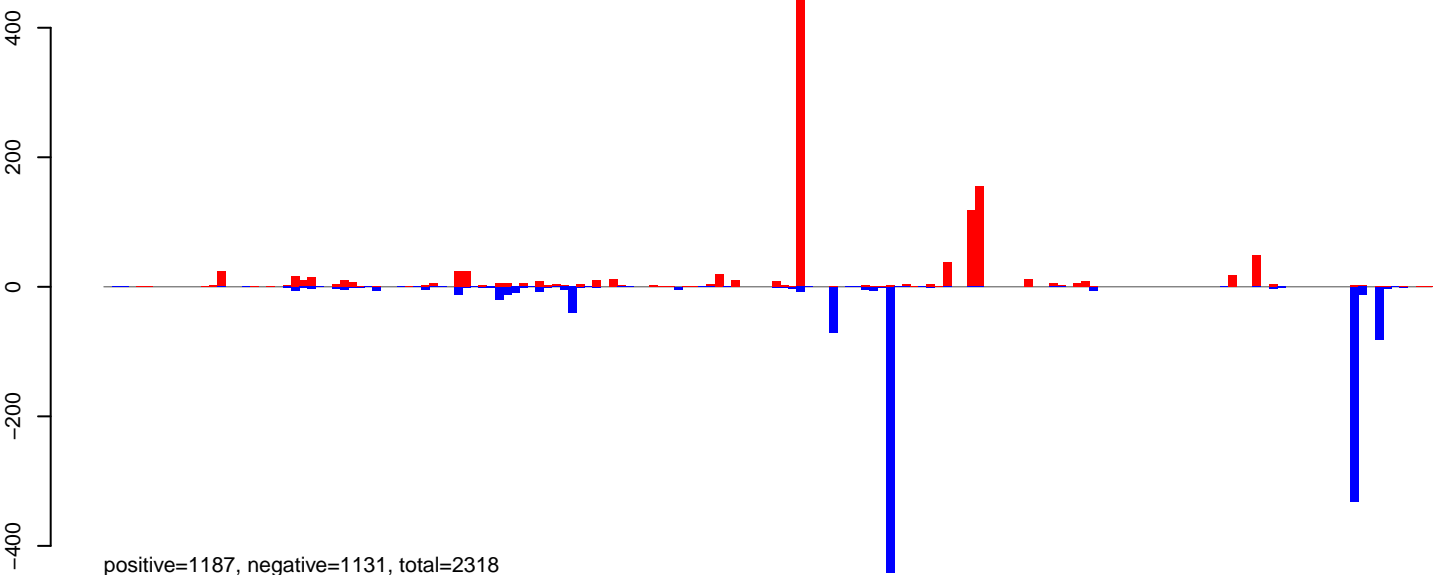
AeAlbo_C636_B2_FHV_r54q.rep



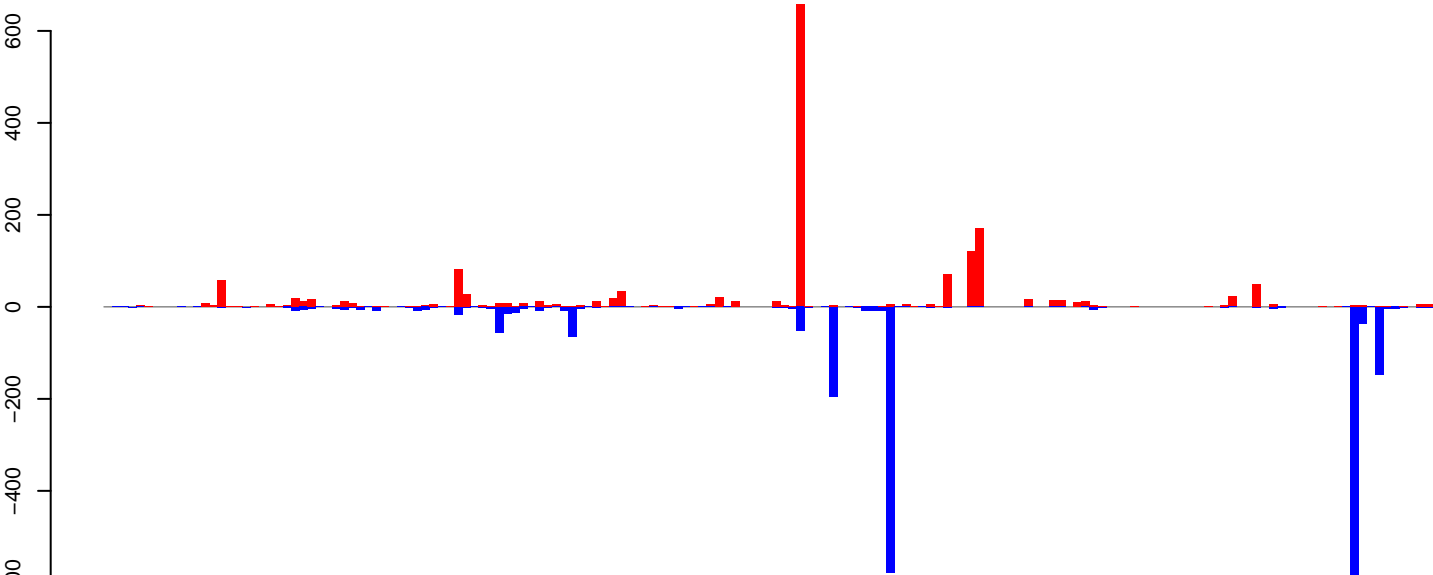
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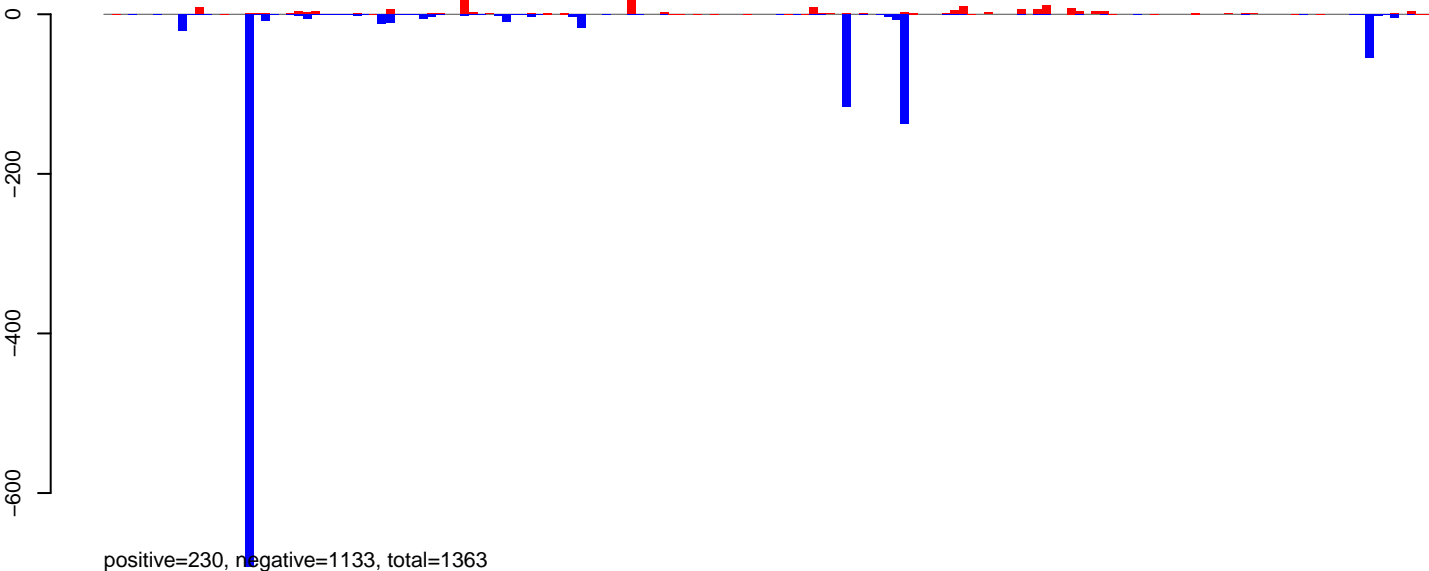
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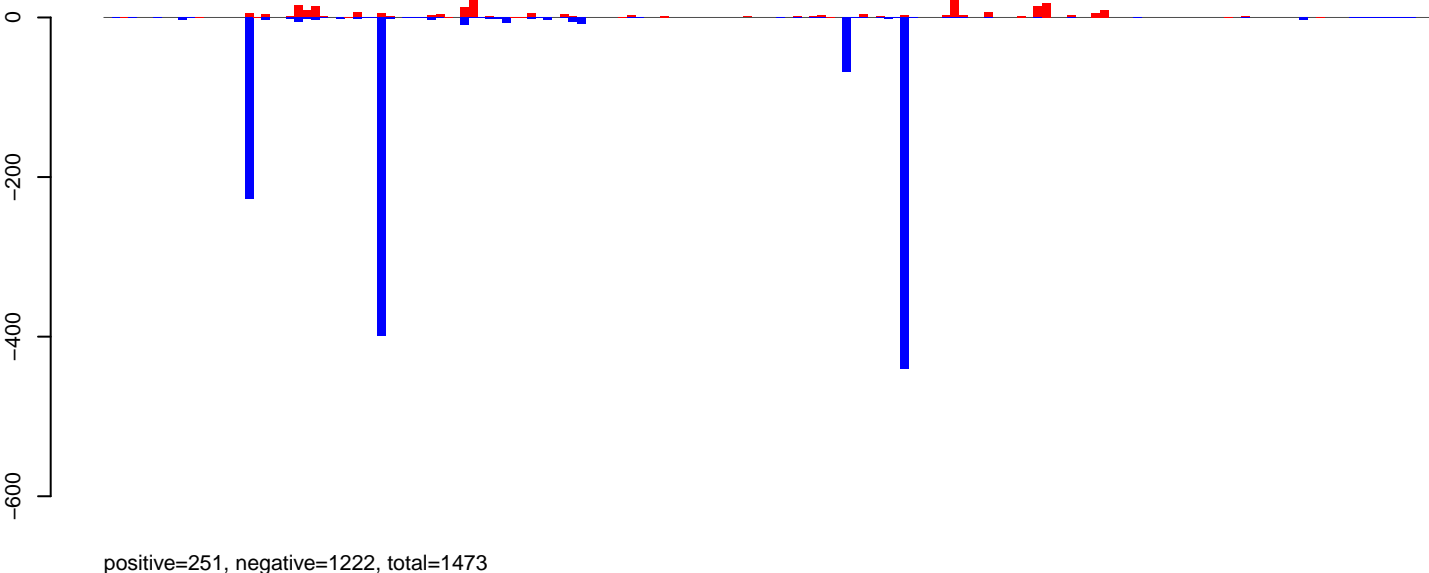
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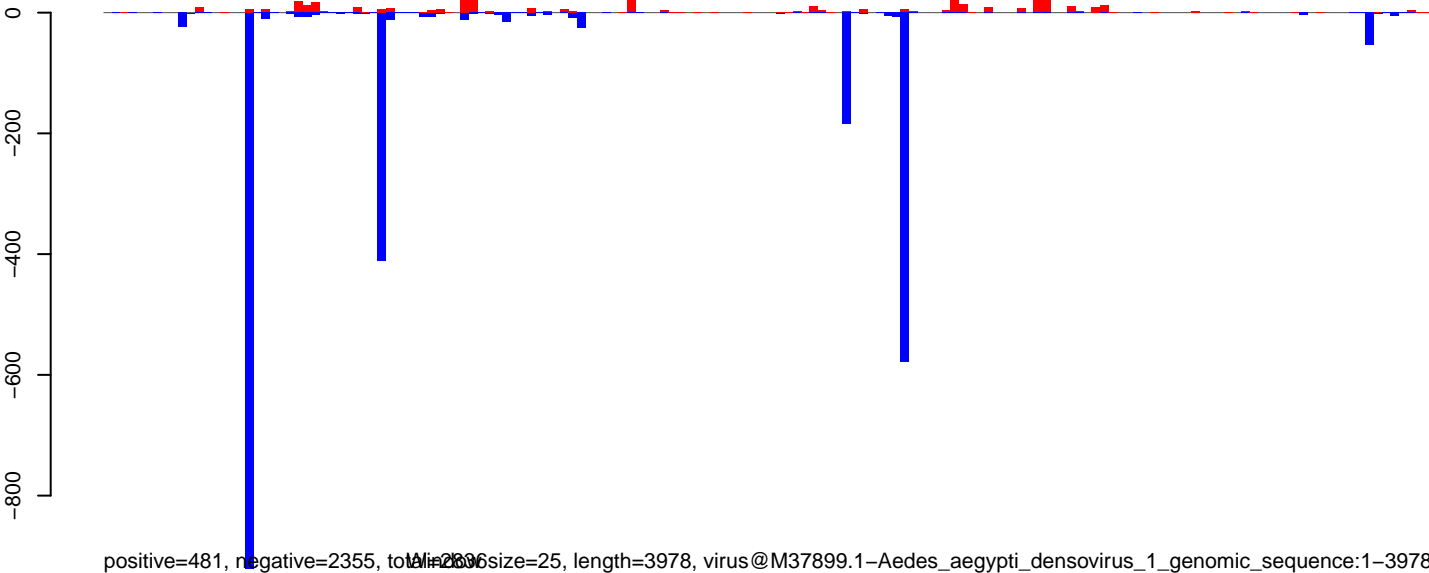
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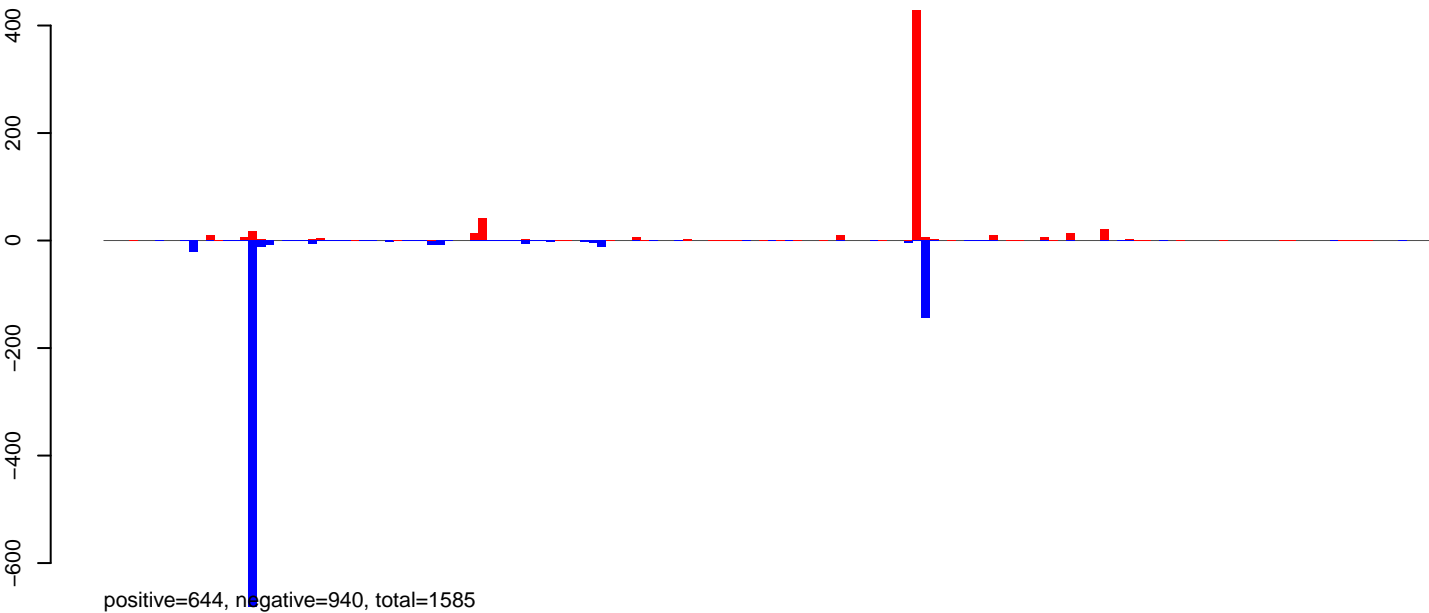
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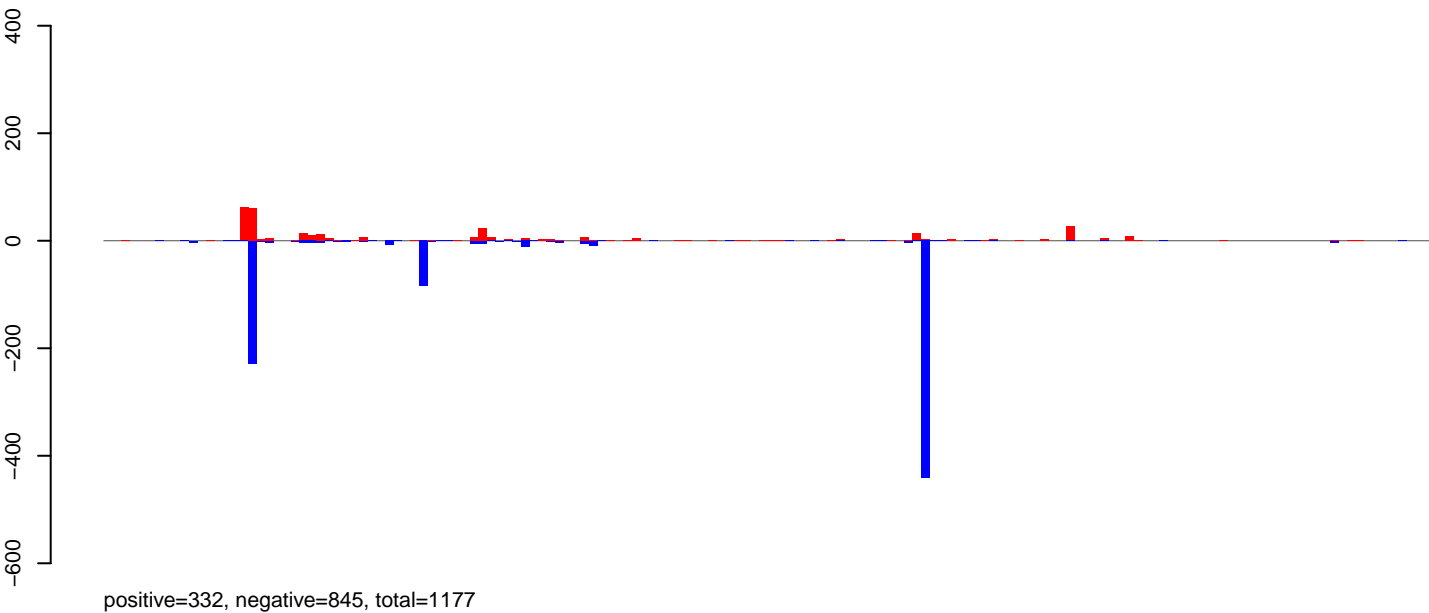
AeAlbo_C636_B2_FHV_r54q.rep



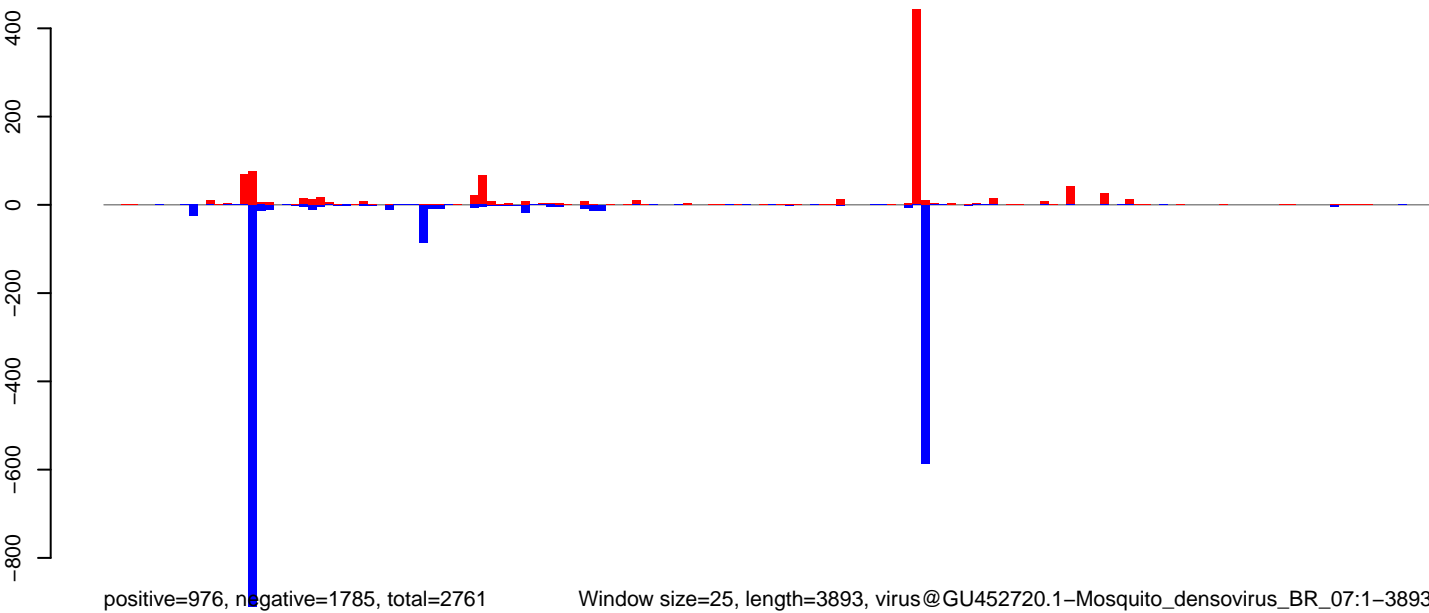
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AeAlbo_C636_B2_FHV_r54q.24_35.rep

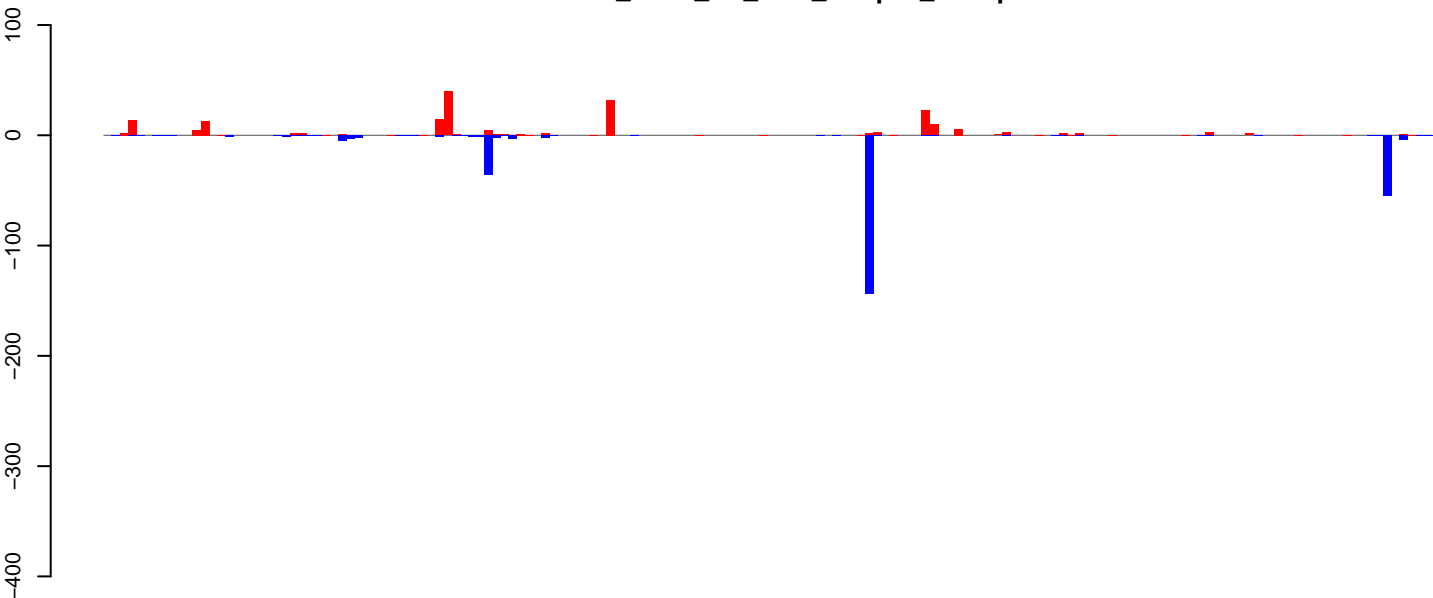


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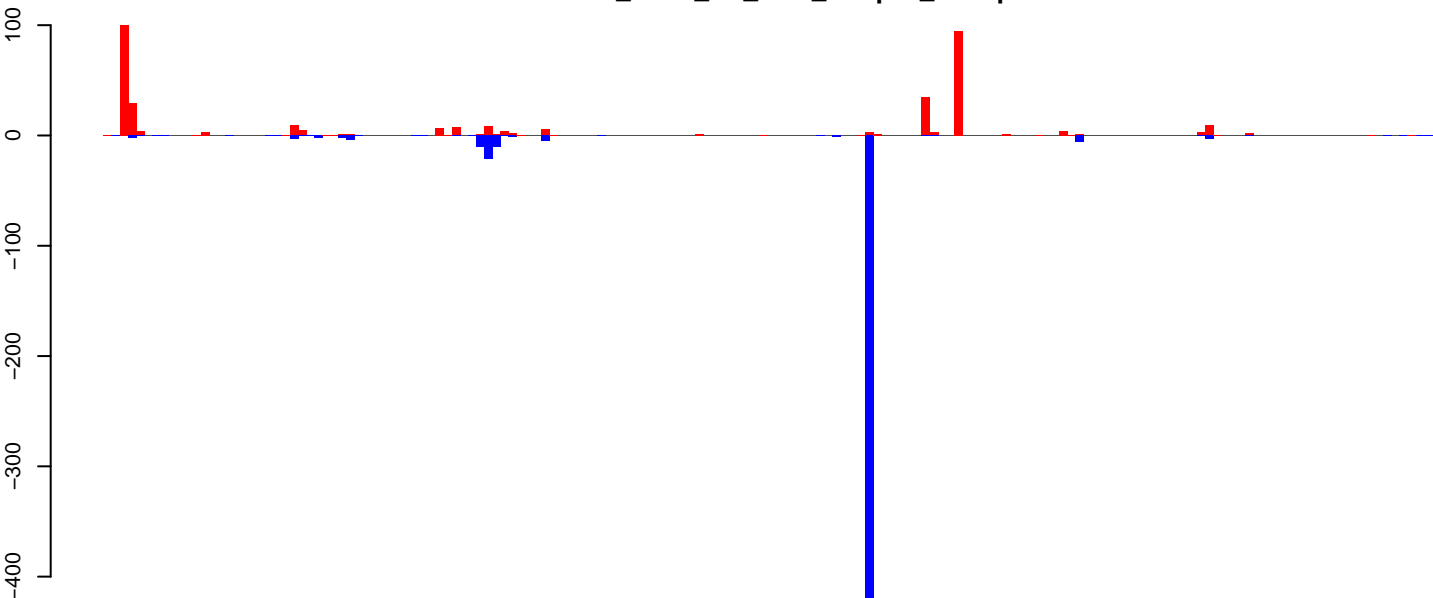
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AeAlbo_C636_B2_FHV_r54q.18_23.rep



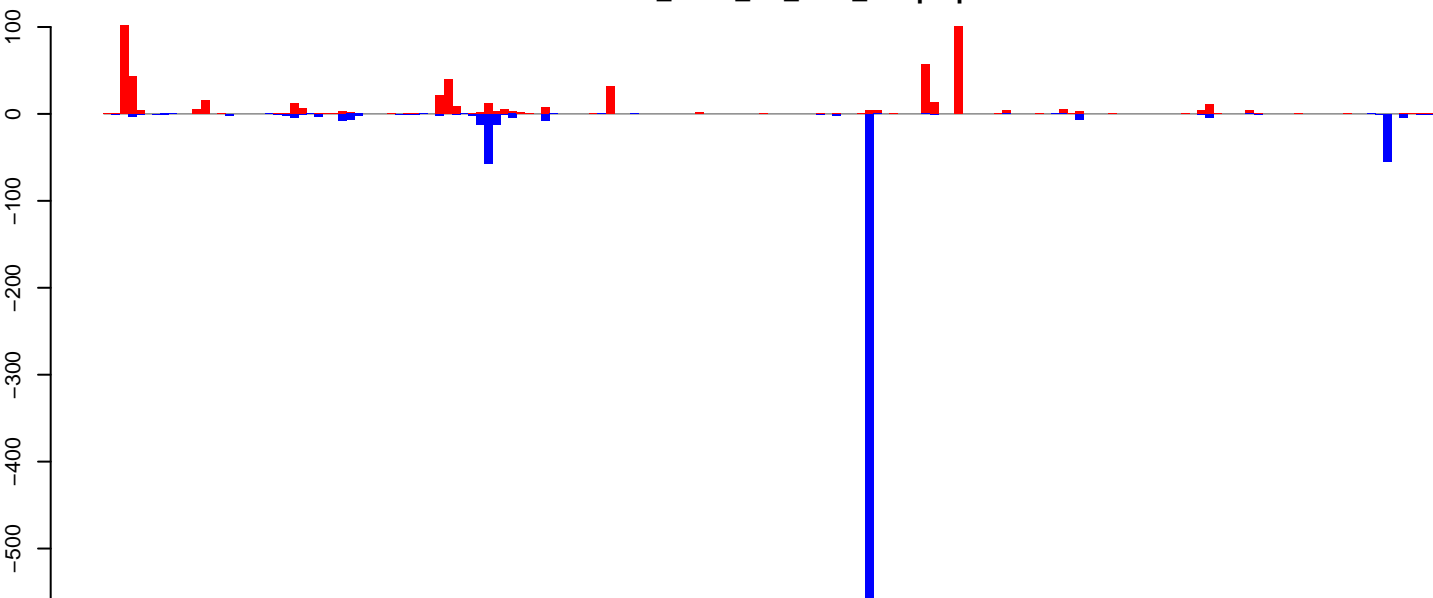
positive=194, negative=275, total=468

AeAlbo_C636_B2_FHV_r54q.24_35.rep



positive=358, negative=521, total=879

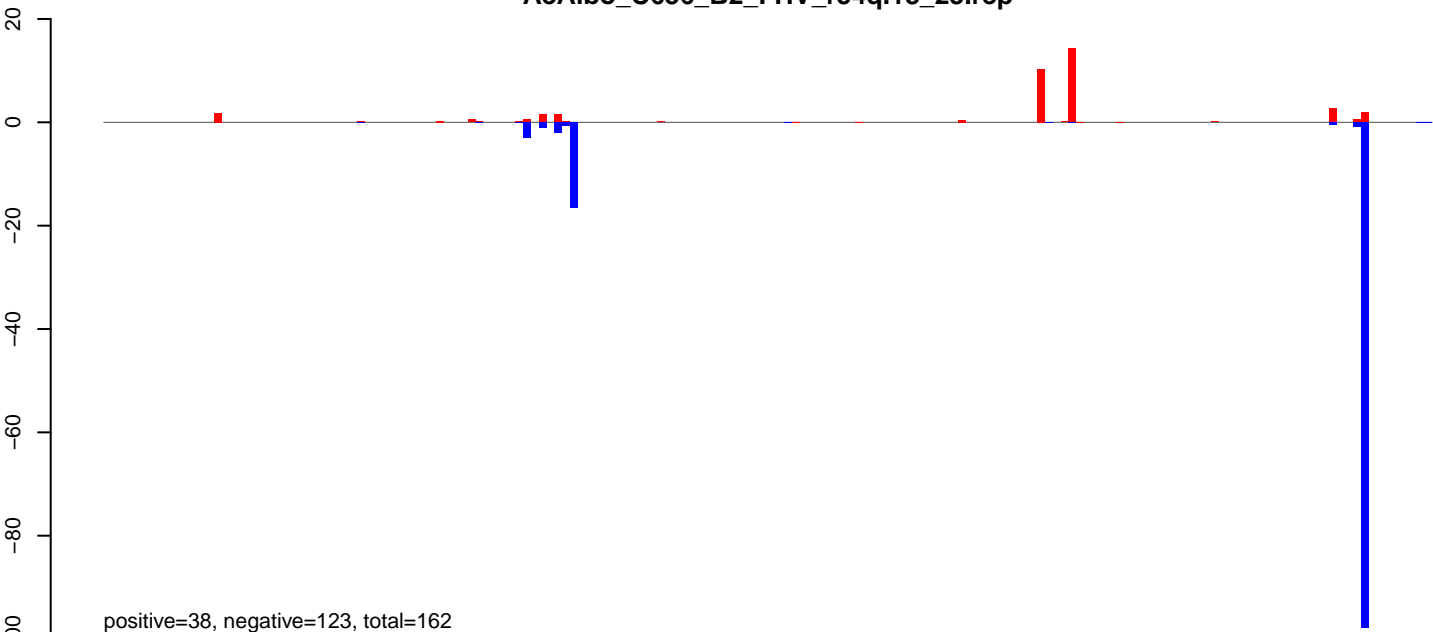
AeAlbo_C636_B2_FHV_r54q.rep



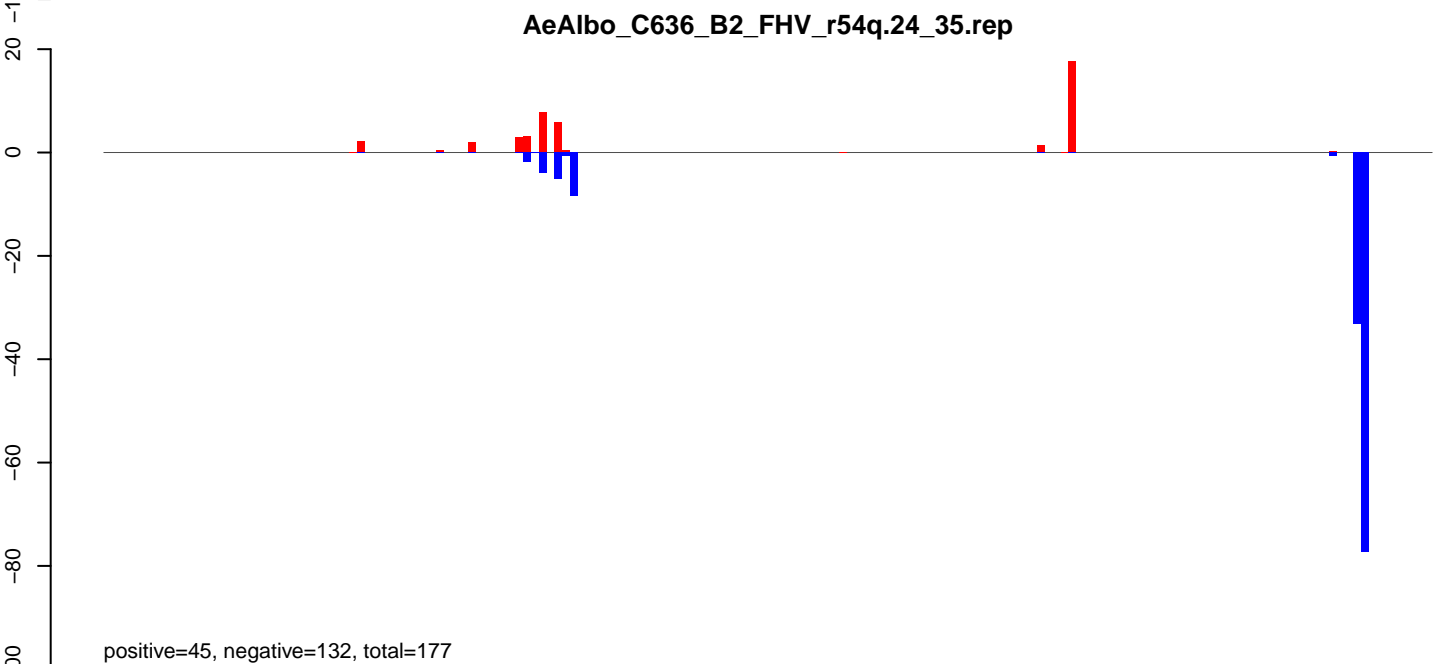
bow size=512, length=4094, positive=756, negative=1053, total=1809 - Aedes_albopictus_C6_36_cell_densovirus_non-structural_protein_1_NS-1_non-structural:1-4094

0 1000 2000 3000 4000

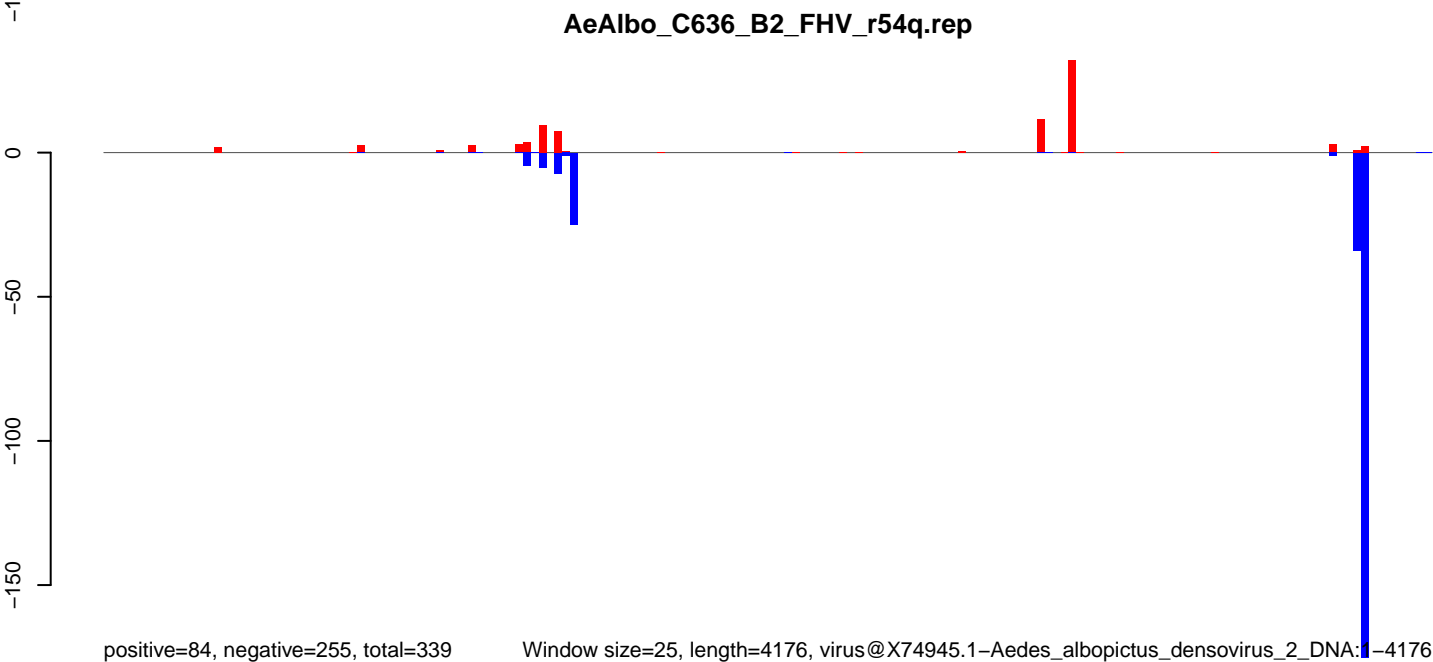
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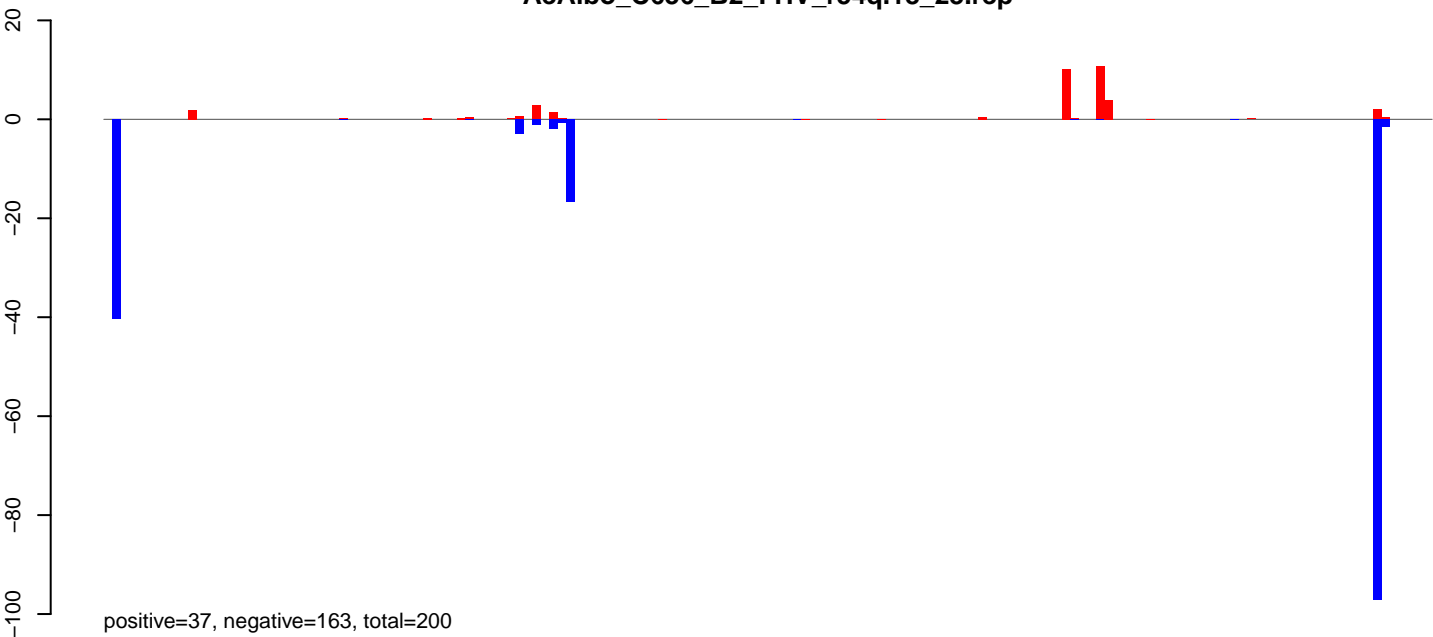
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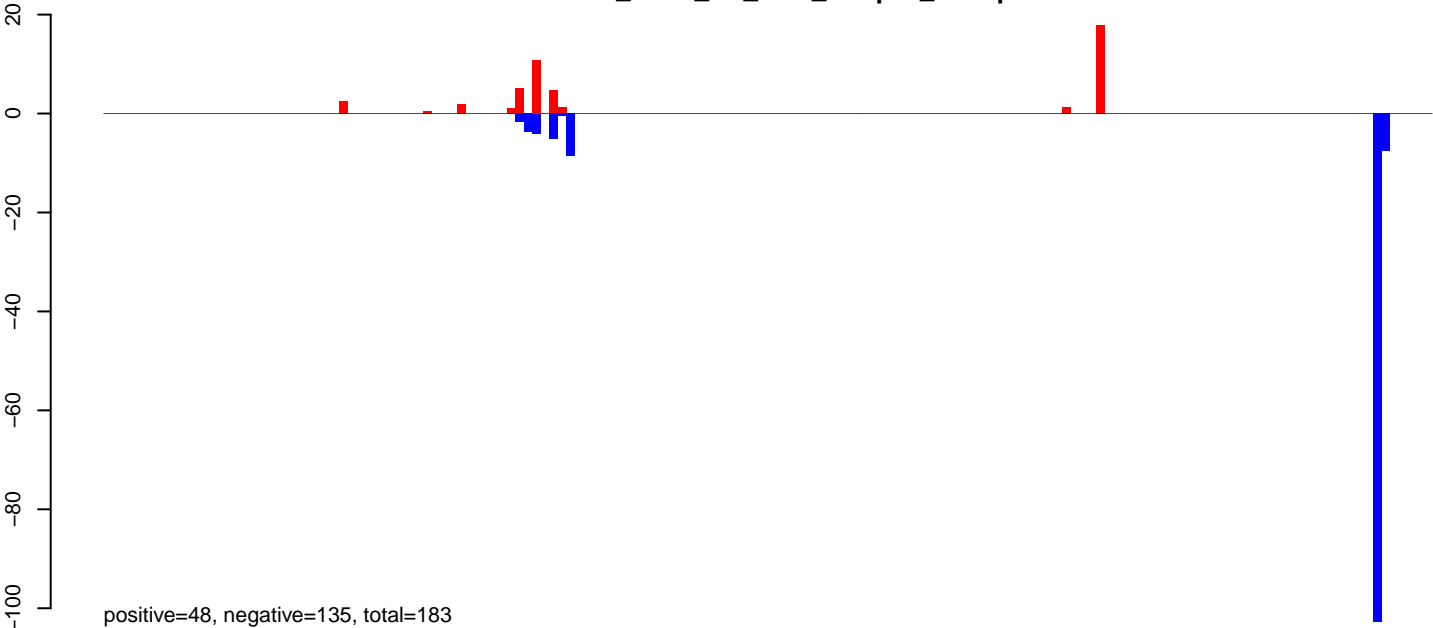
AeAlbo_C636_B2_FHV_r54q.rep



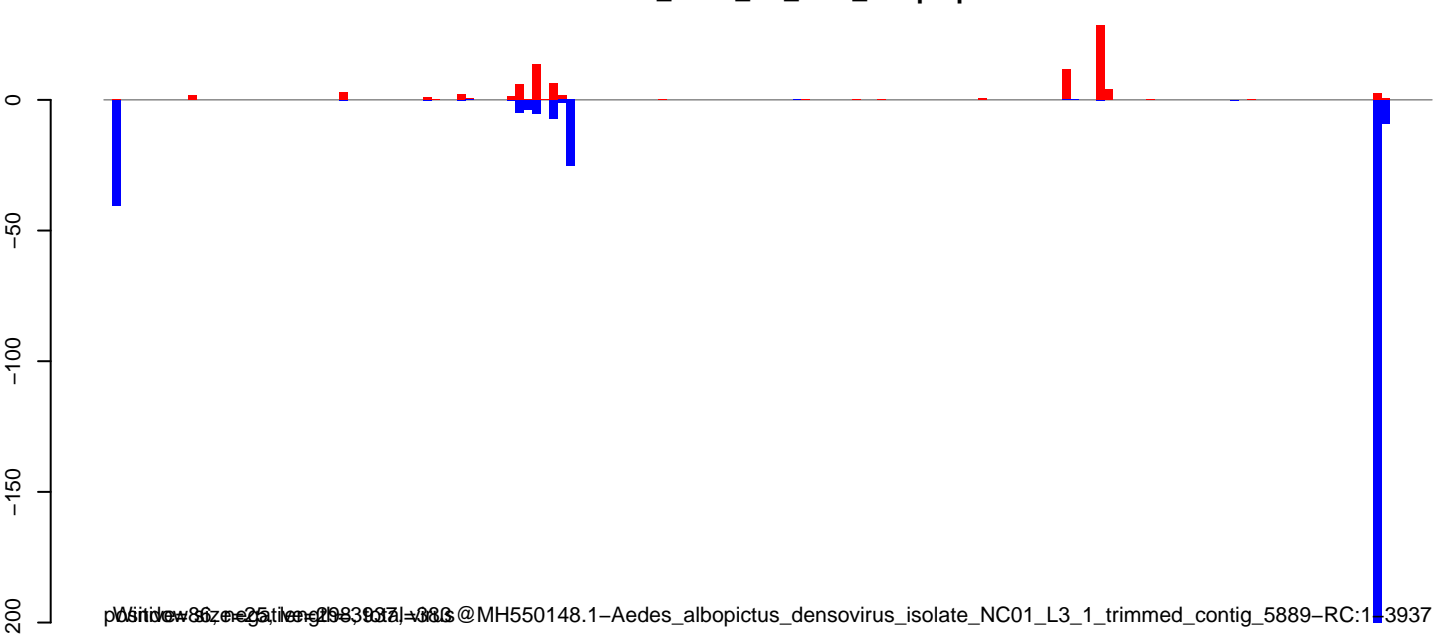
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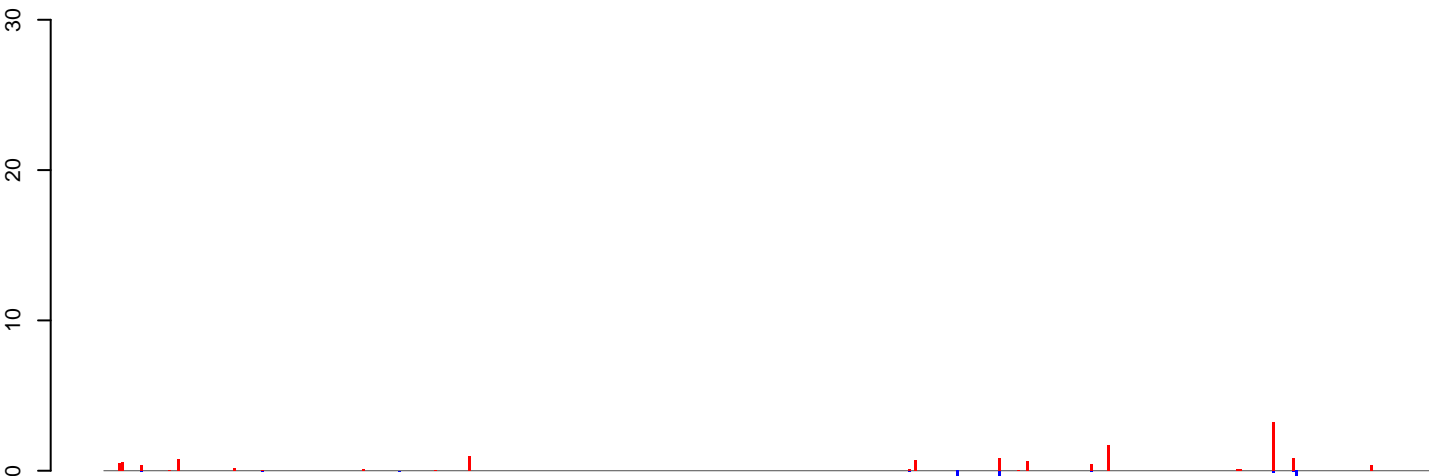
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AeAlbo_C636_B2_FHV_r54q.rep

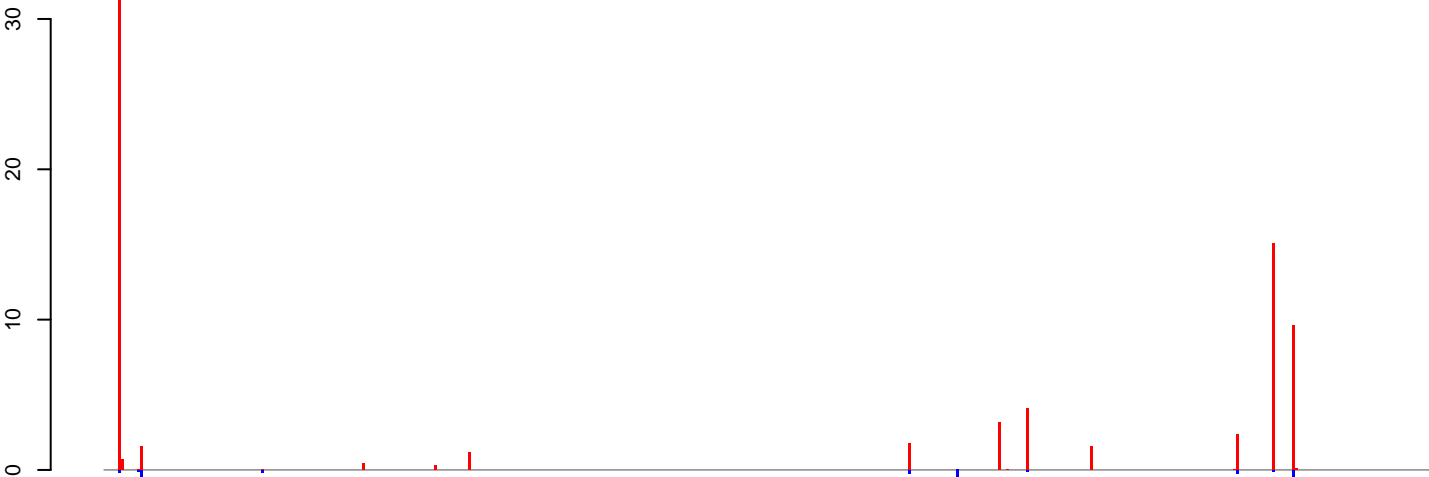


AeAlbo_C636_B2_FHV_r54q.18_23.rep



positive=13, negative=2, total=15

AeAlbo_C636_B2_FHV_r54q.24_35.rep



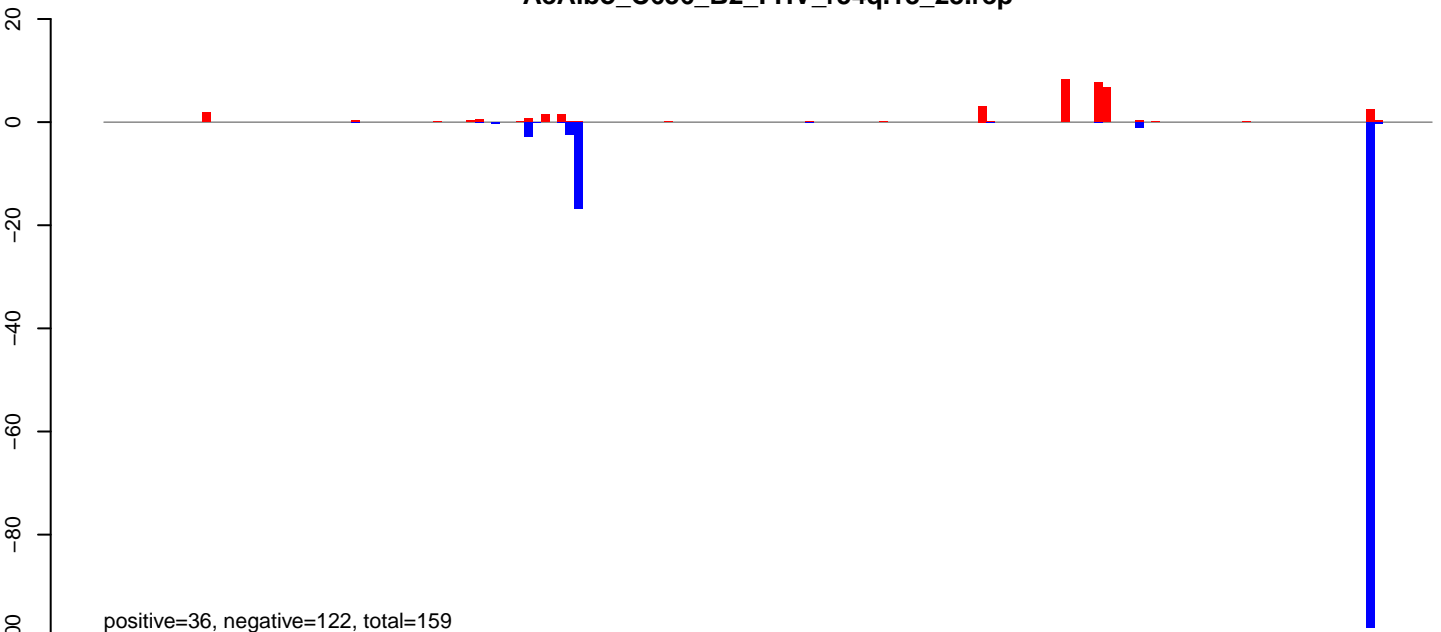
positive=77, negative=8, total=85

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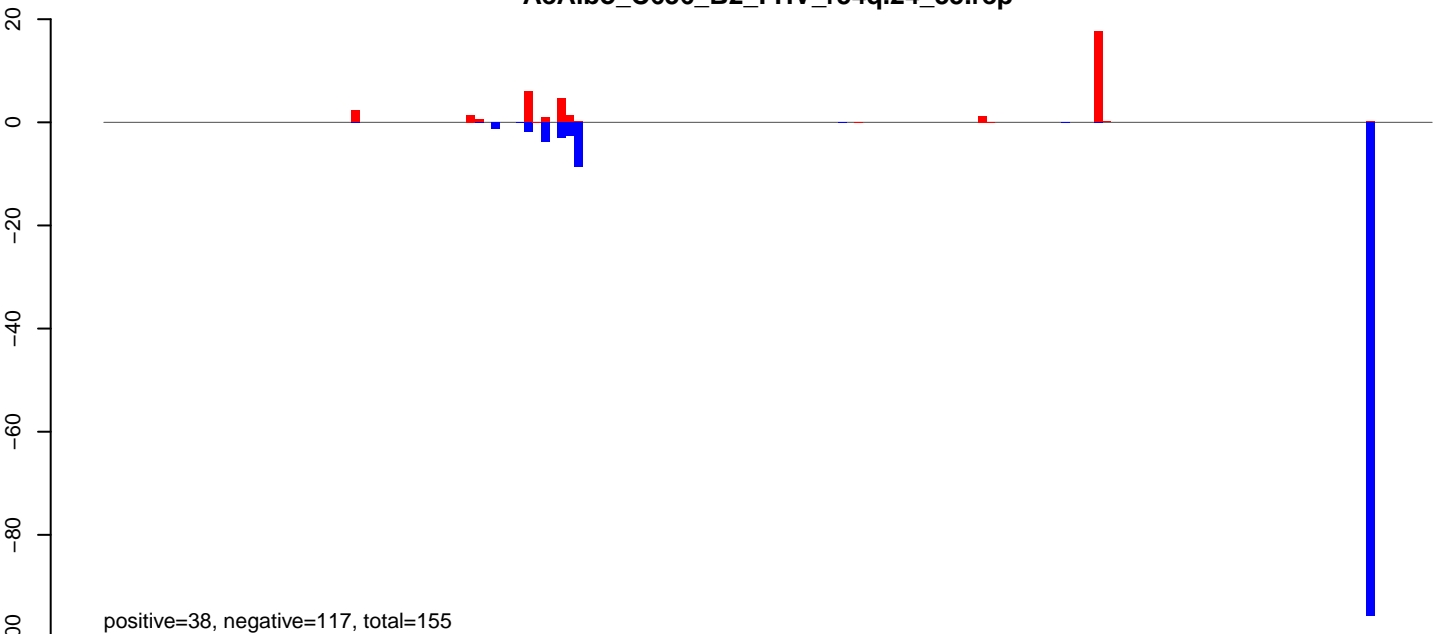


positive=90, negative=10, total=100, window size=25, length=11835, virus@M20303.1-O-Nyong-nyong_virus_Gulu_strain_complete_genome:1-11835

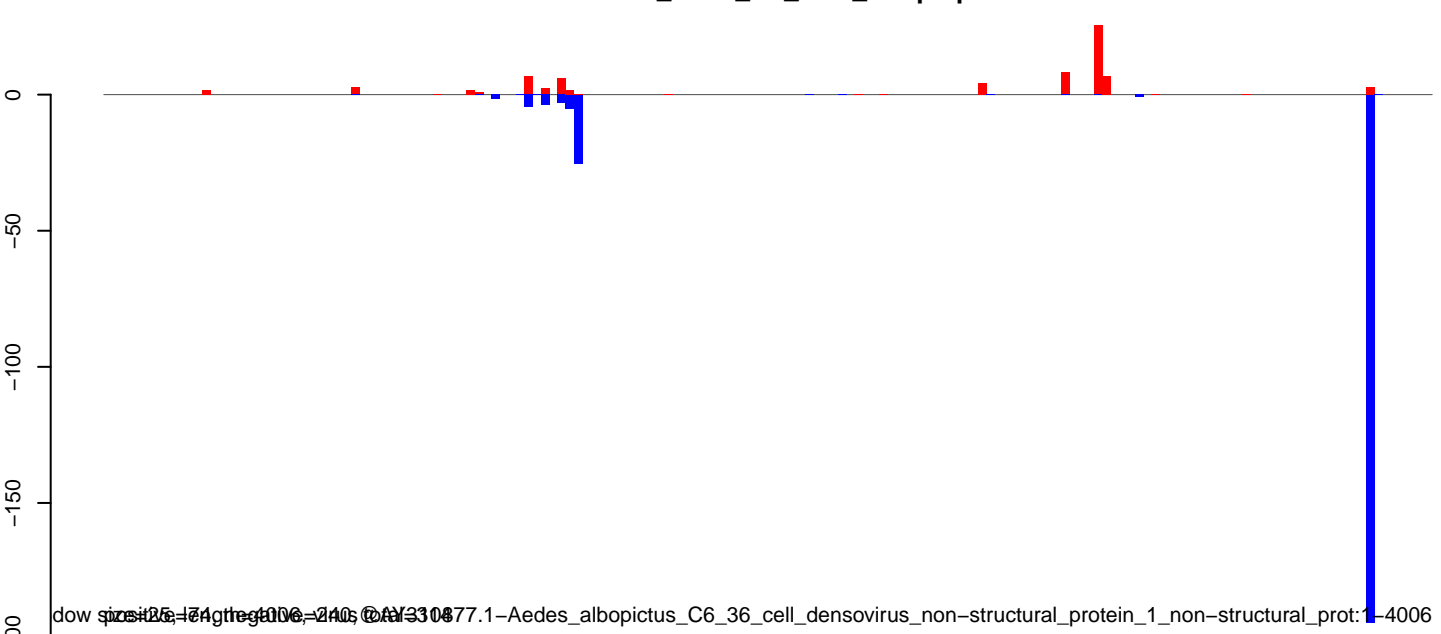
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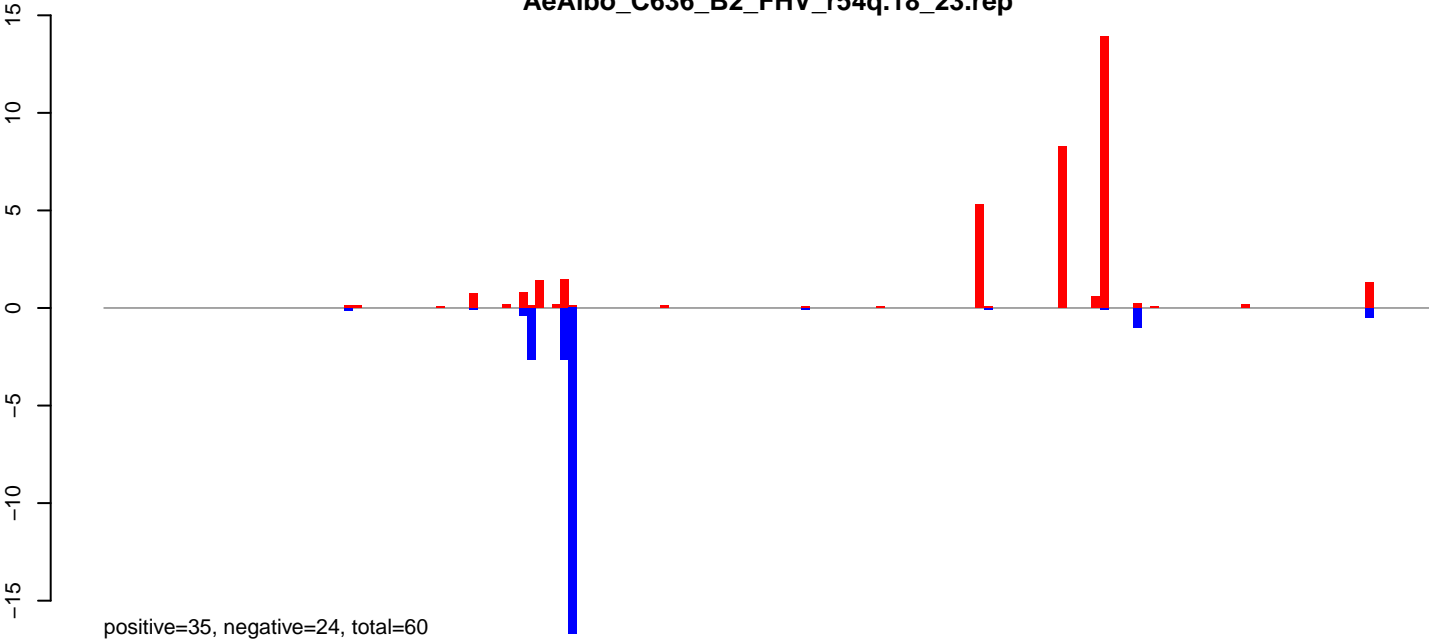
AeAlbo_C636_B2_FHV_r54q.24_35.rep



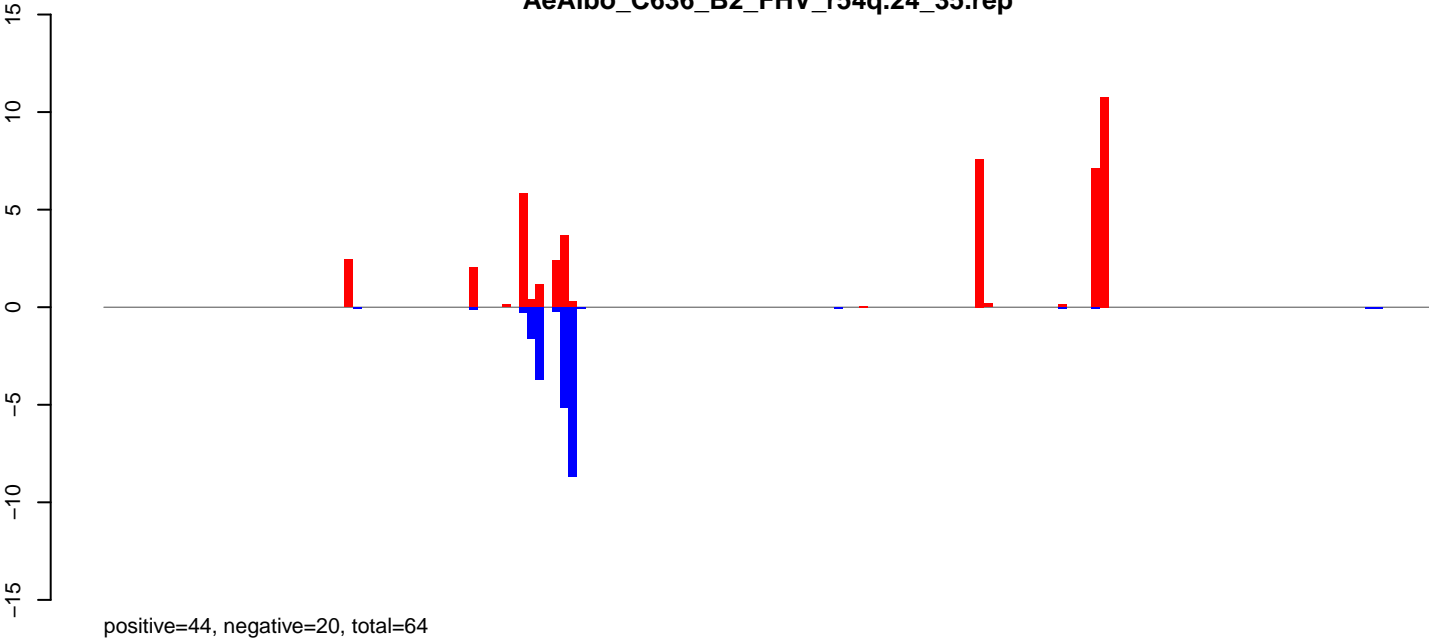
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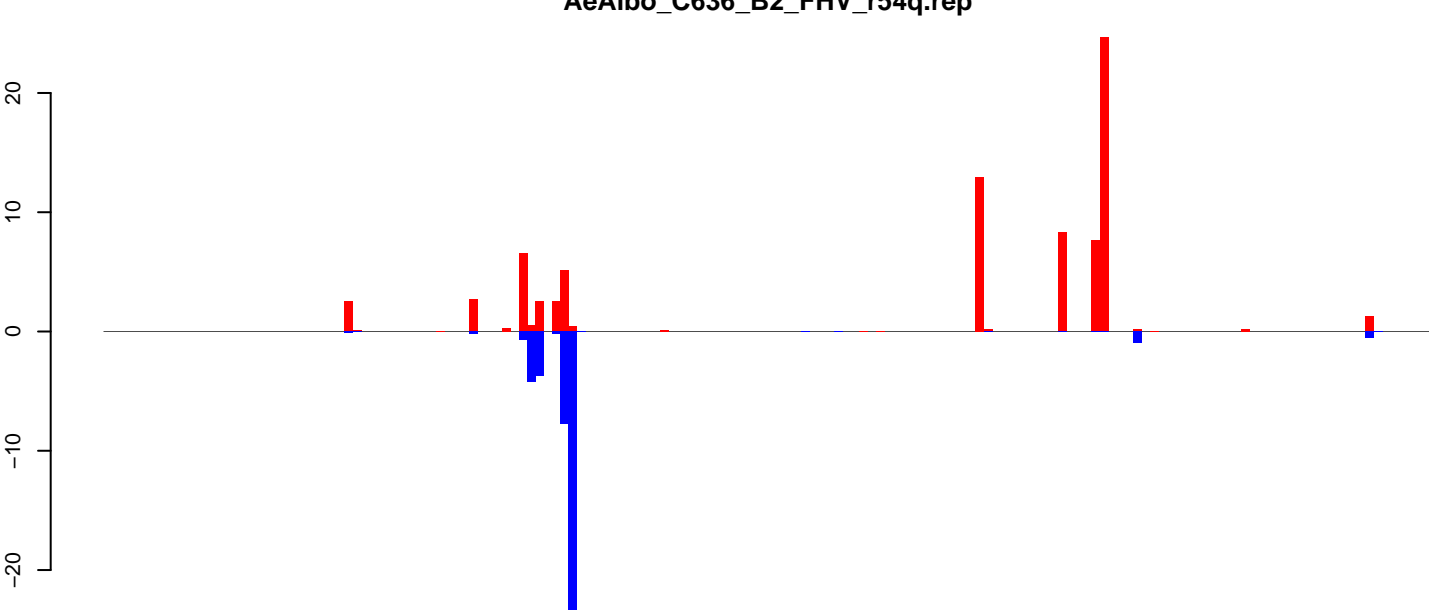
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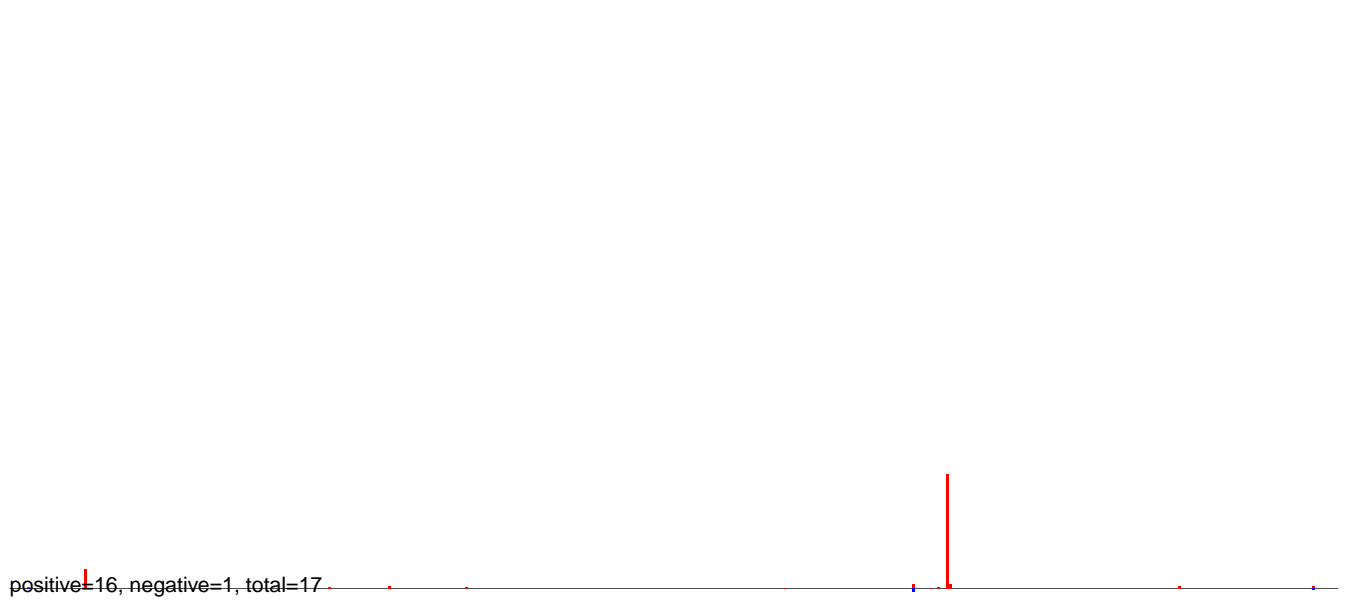
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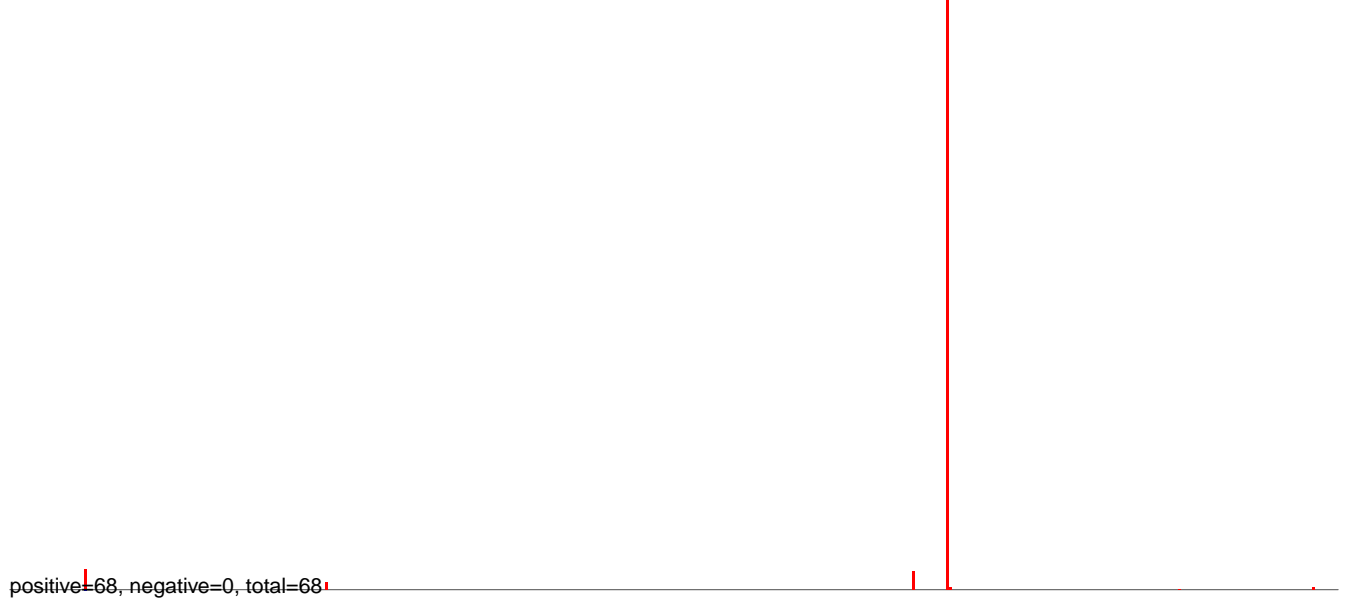
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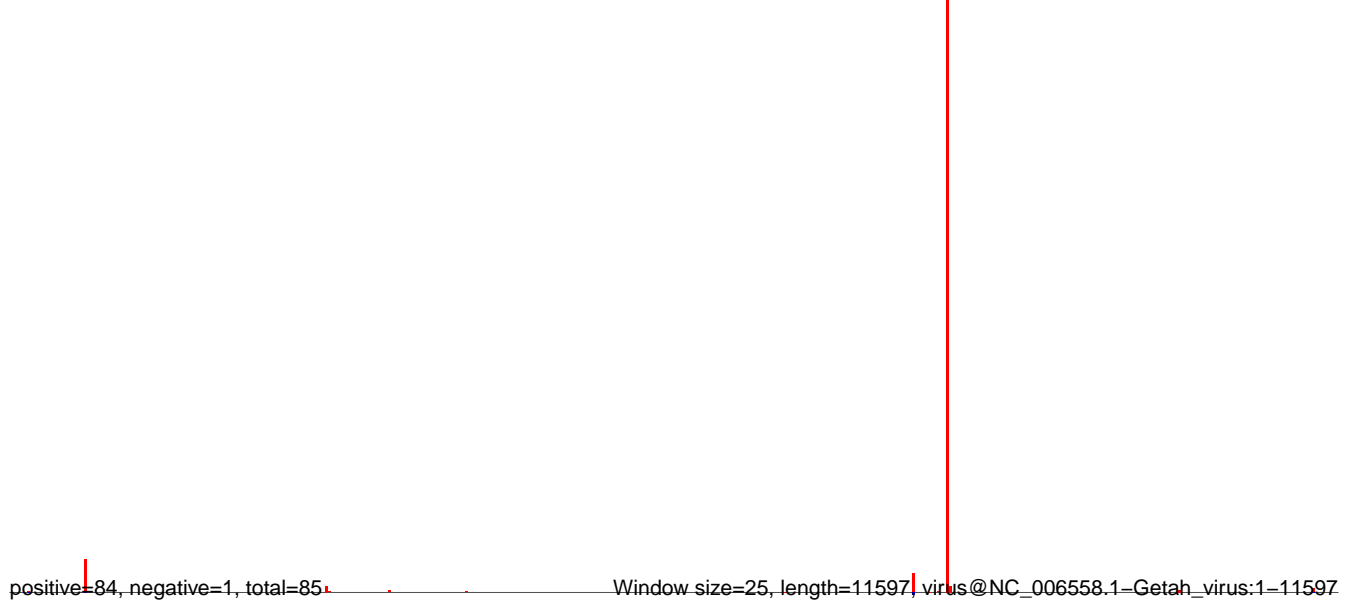
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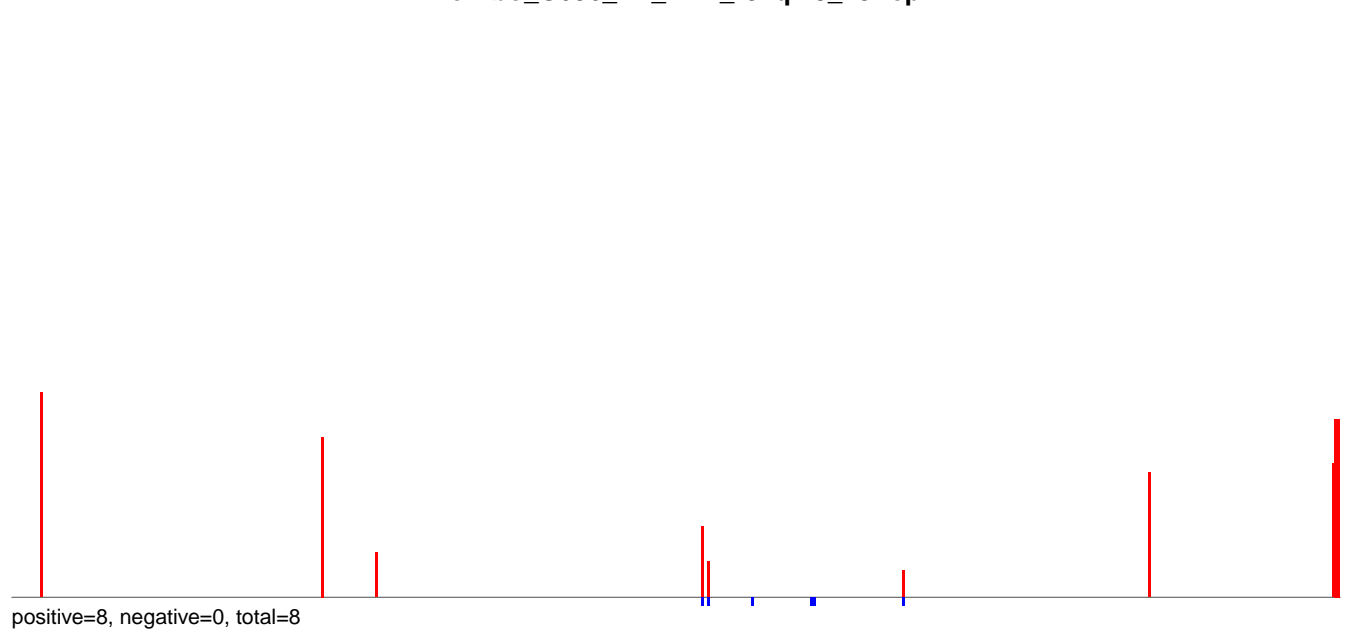
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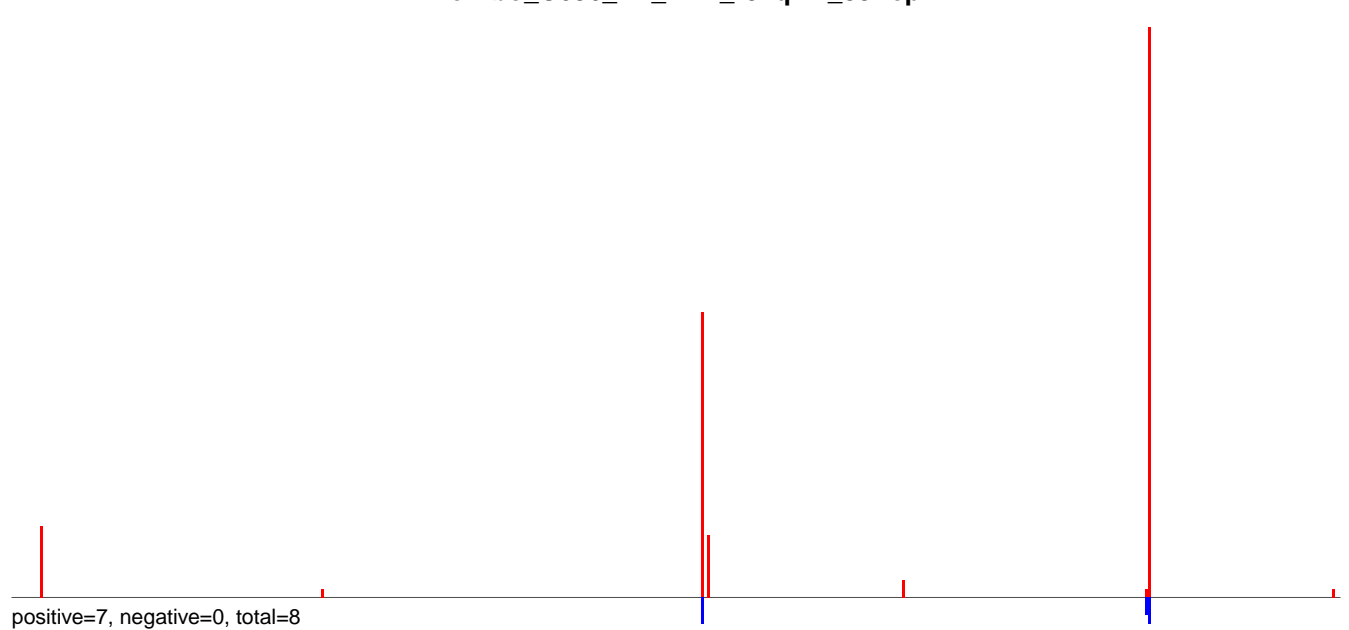
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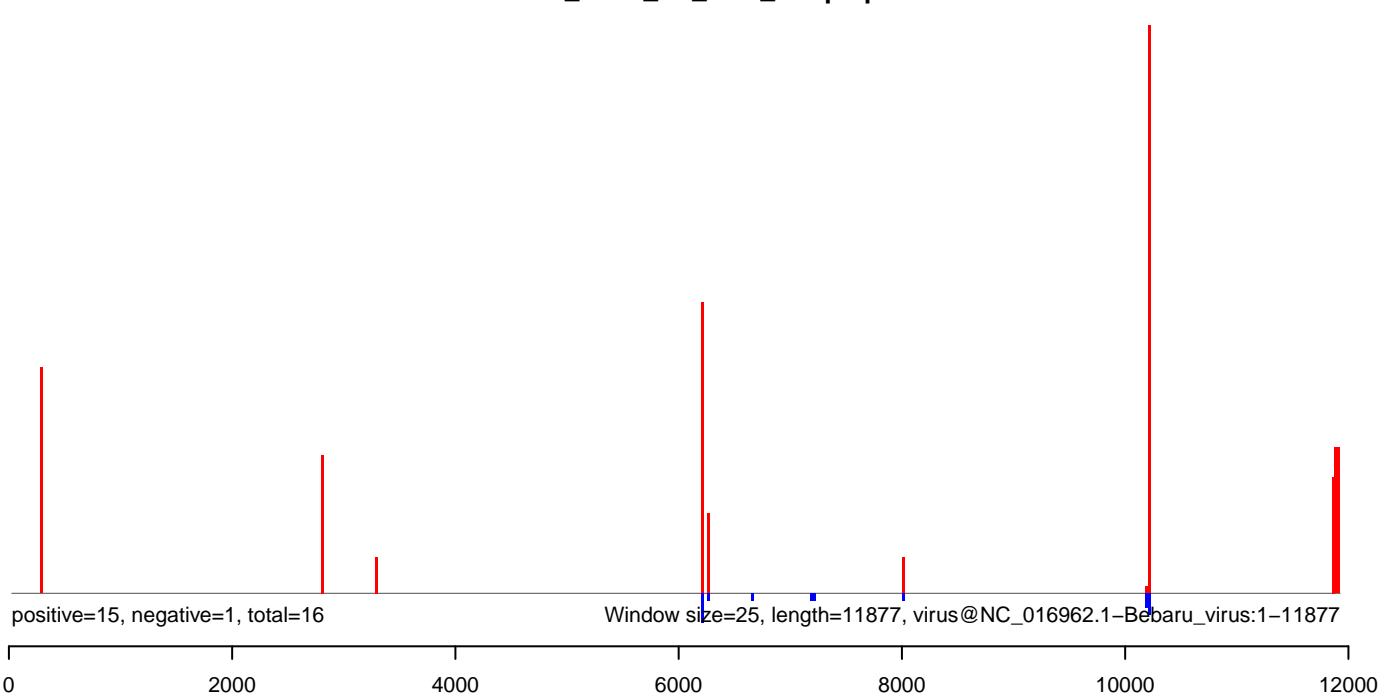
AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep



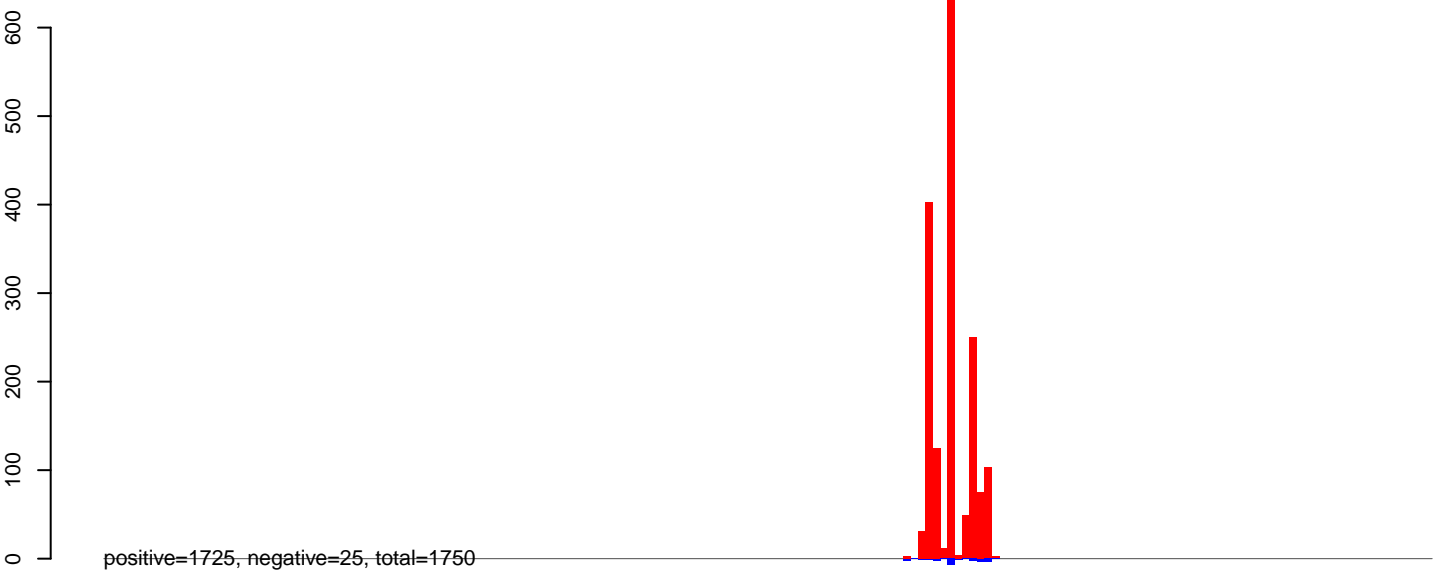
AeAlbo_C636_B2_FHV_r54q.rep



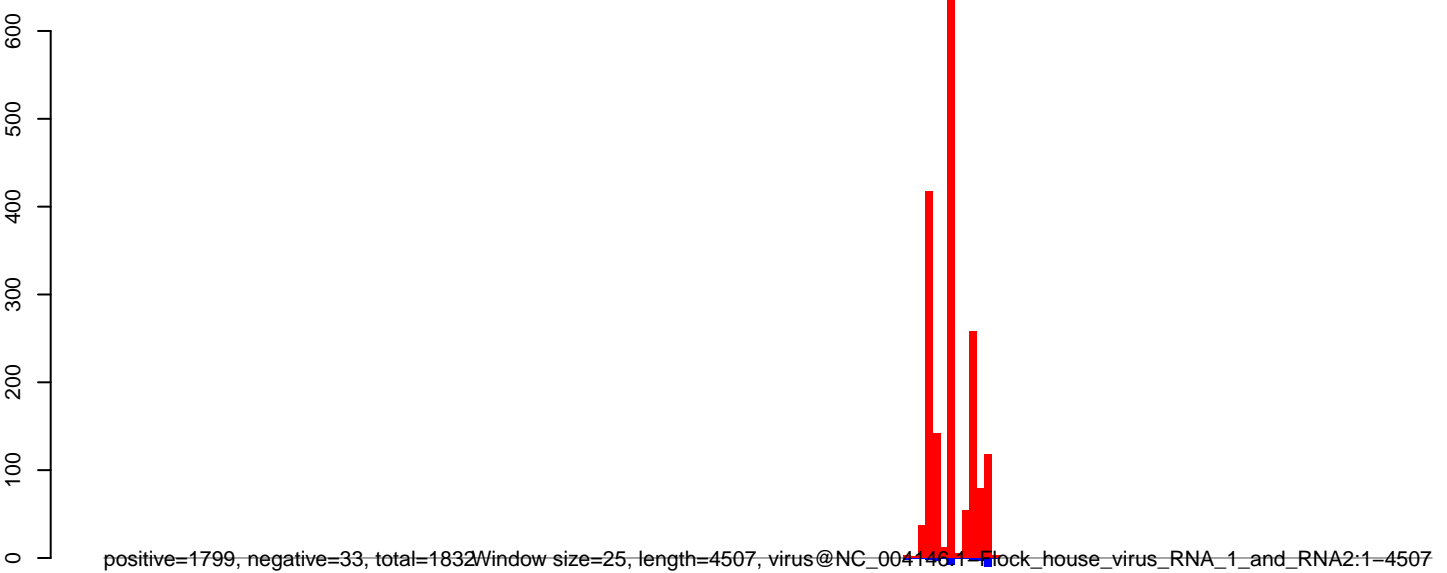
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AeAlbo_C636_B2_FHV_r54q.24_35.rep



AeAlbo_C636_B2_FHV_r54q.rep

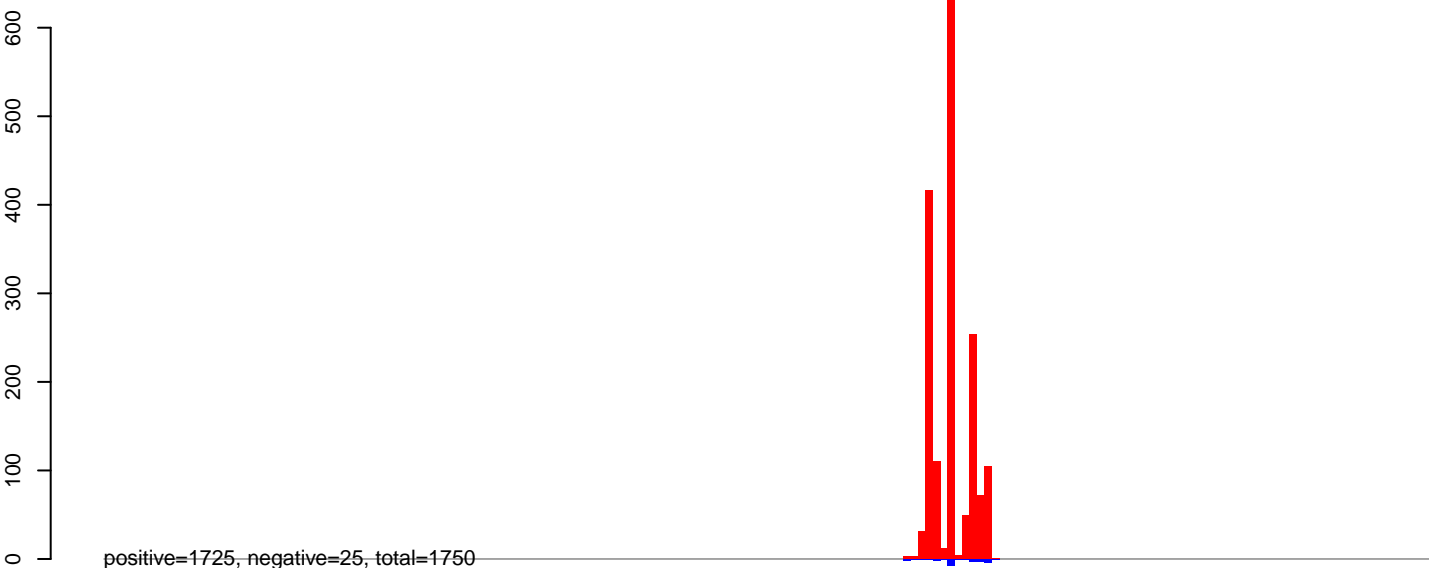


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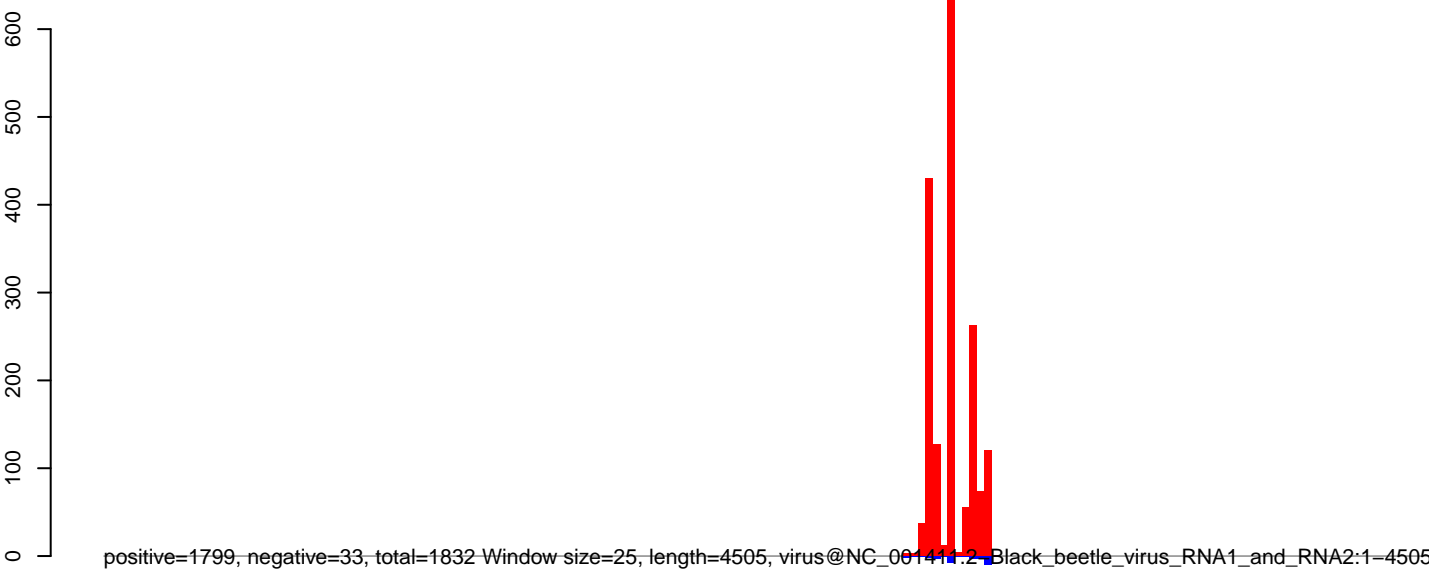
AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep



AeAlbo_C636_B2_FHV_r54q.rep



0 1000 2000 3000 4000

AeAlbo_C636_B2_FHV_r54q.18_23.rep

positive=40, negative=6, total=46

AeAlbo_C636_B2_FHV_r54q.24_35.rep

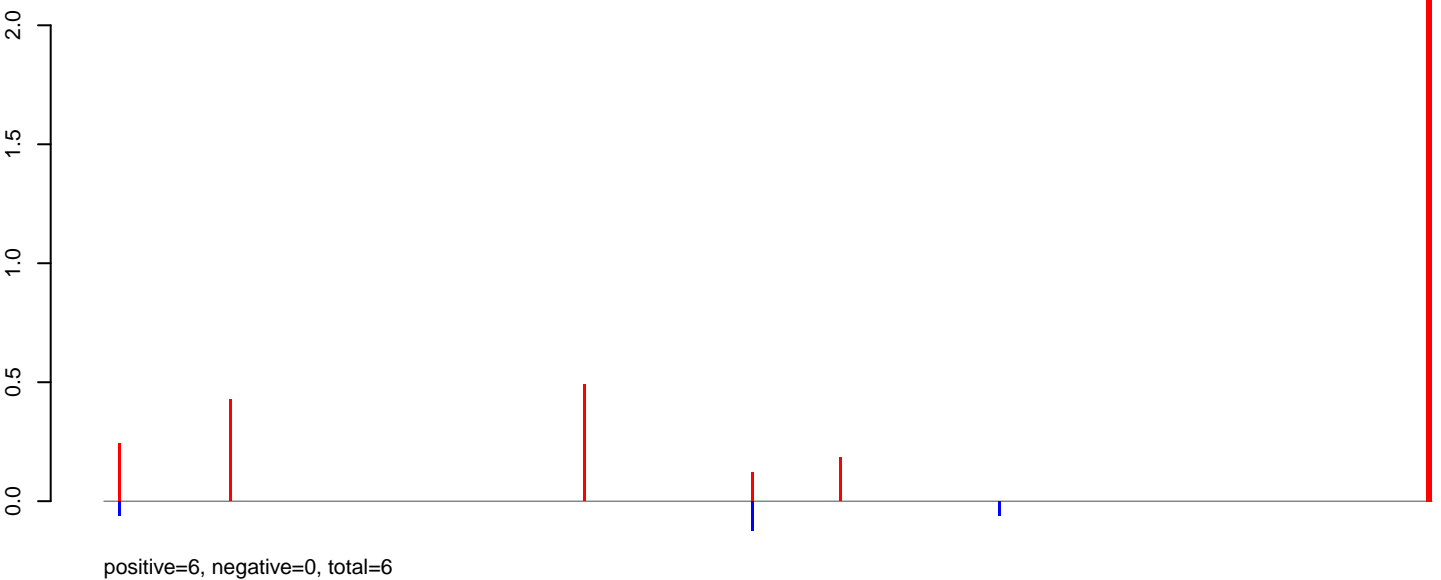
positive=356, negative=10, total=367

AeAlbo_C636_B2_FHV_r54q.rep

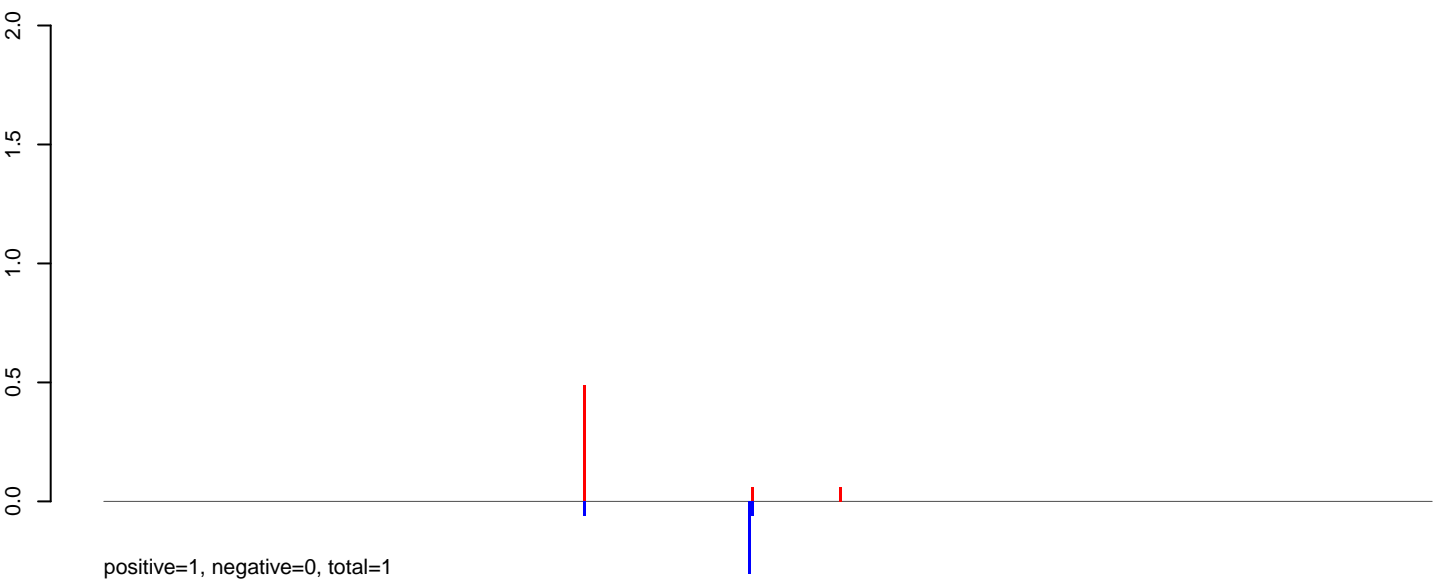
ze=25 positive=352, negative=10, total=362 Drosophila_melanogaster_American_nodavirus_ANV_SW-2009a_segment_RNA1_and_RNA2:1-4523

0 1000 2000 3000 4000

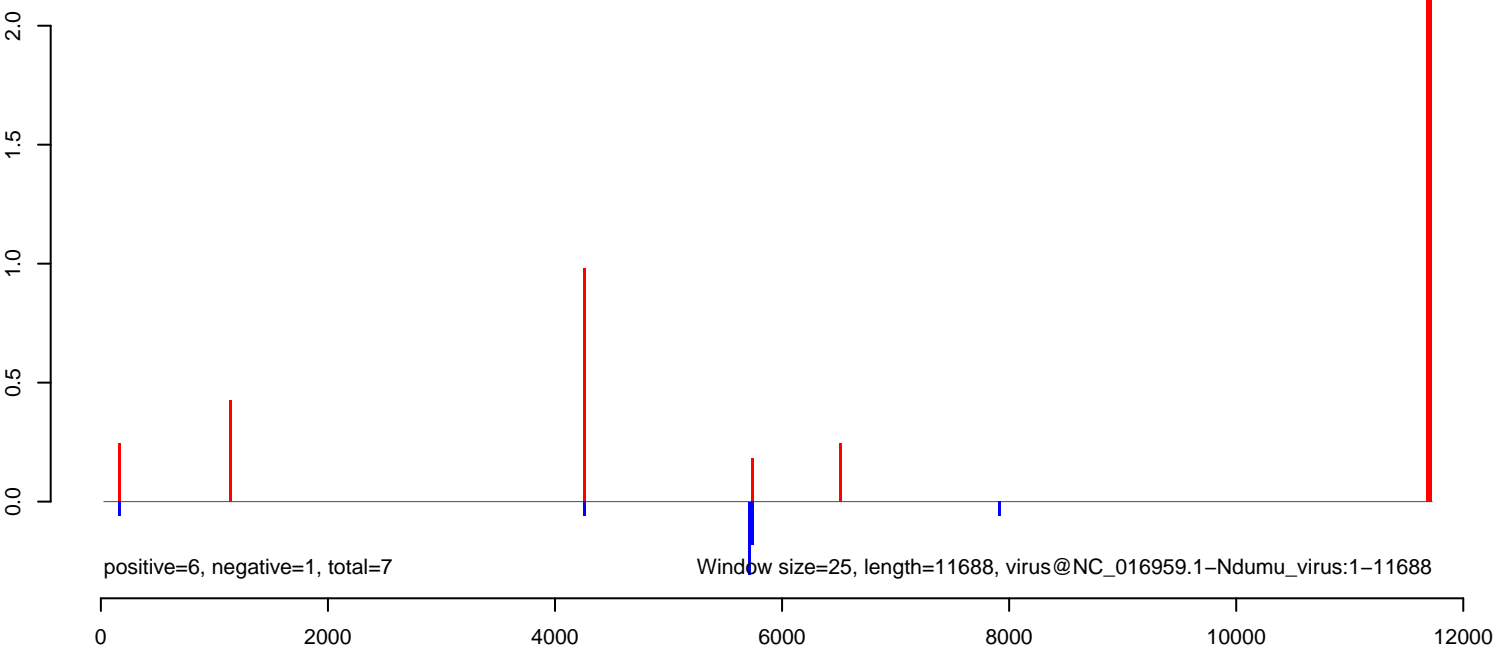
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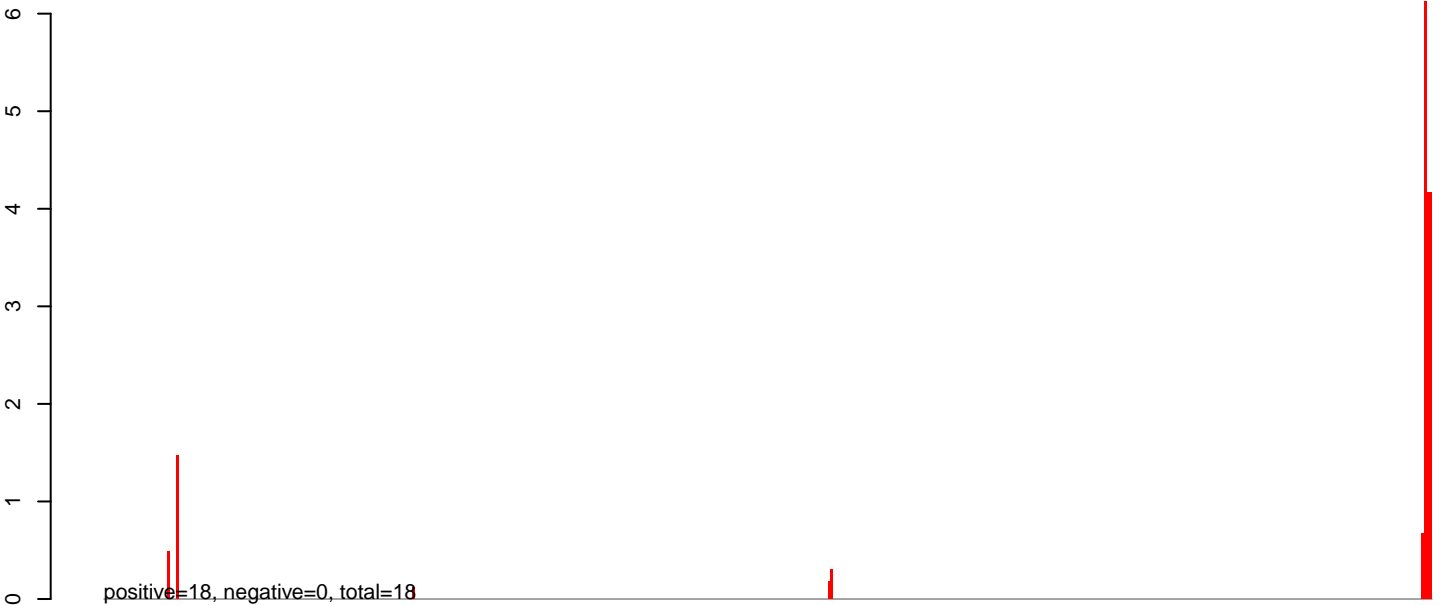
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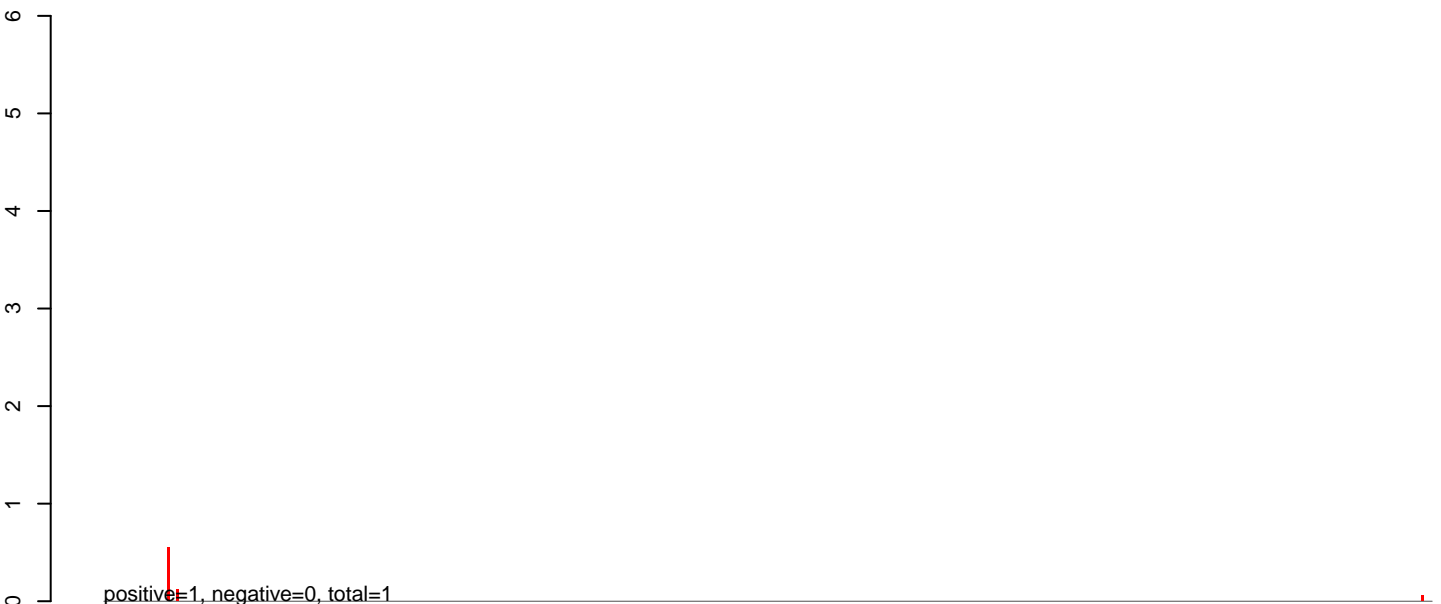
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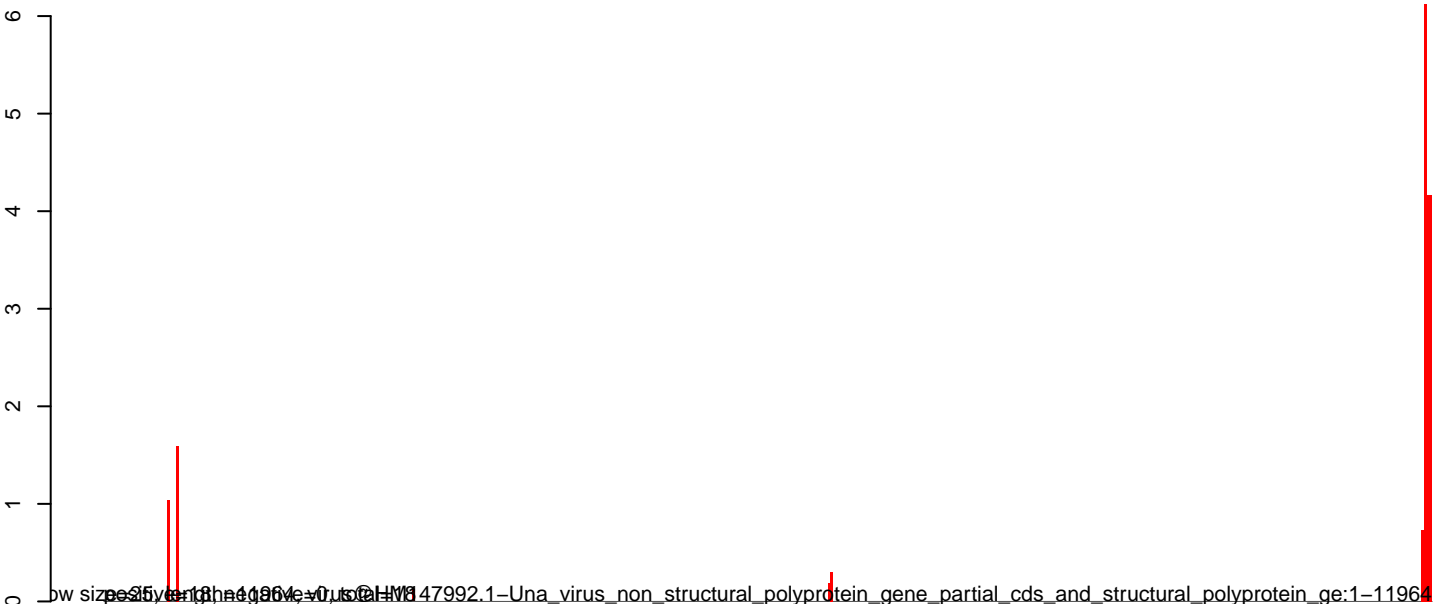
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AeAlbo_C636_B2_FHV_r54q.24_35.rep



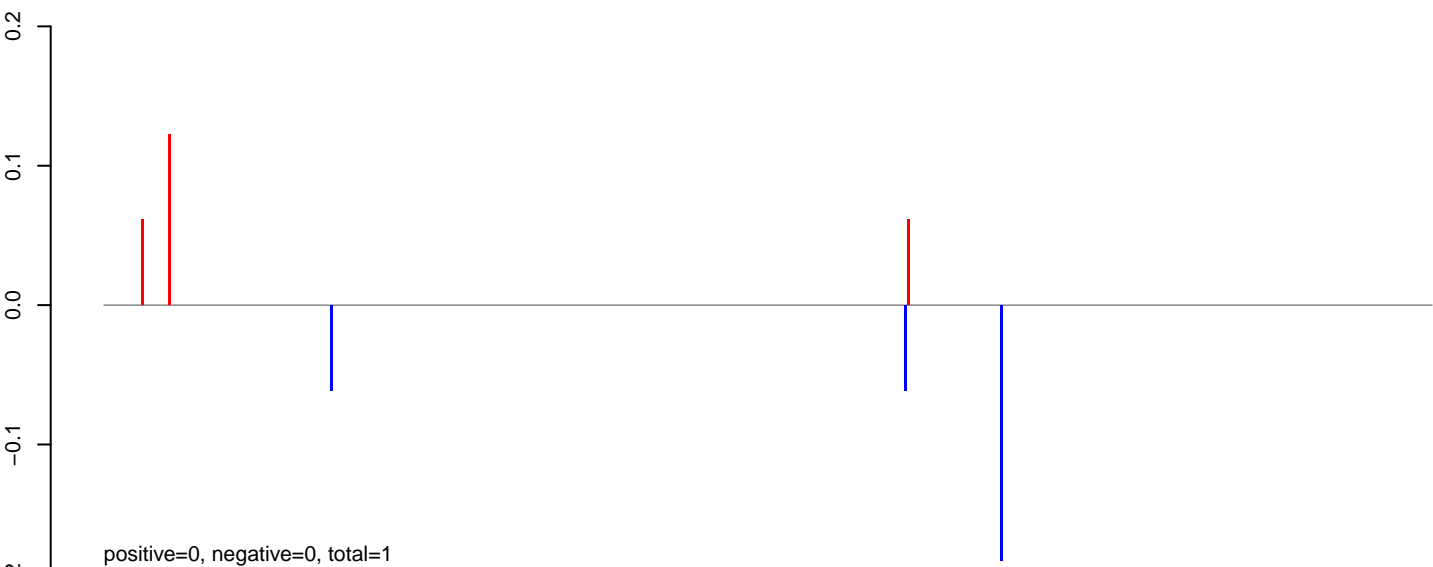
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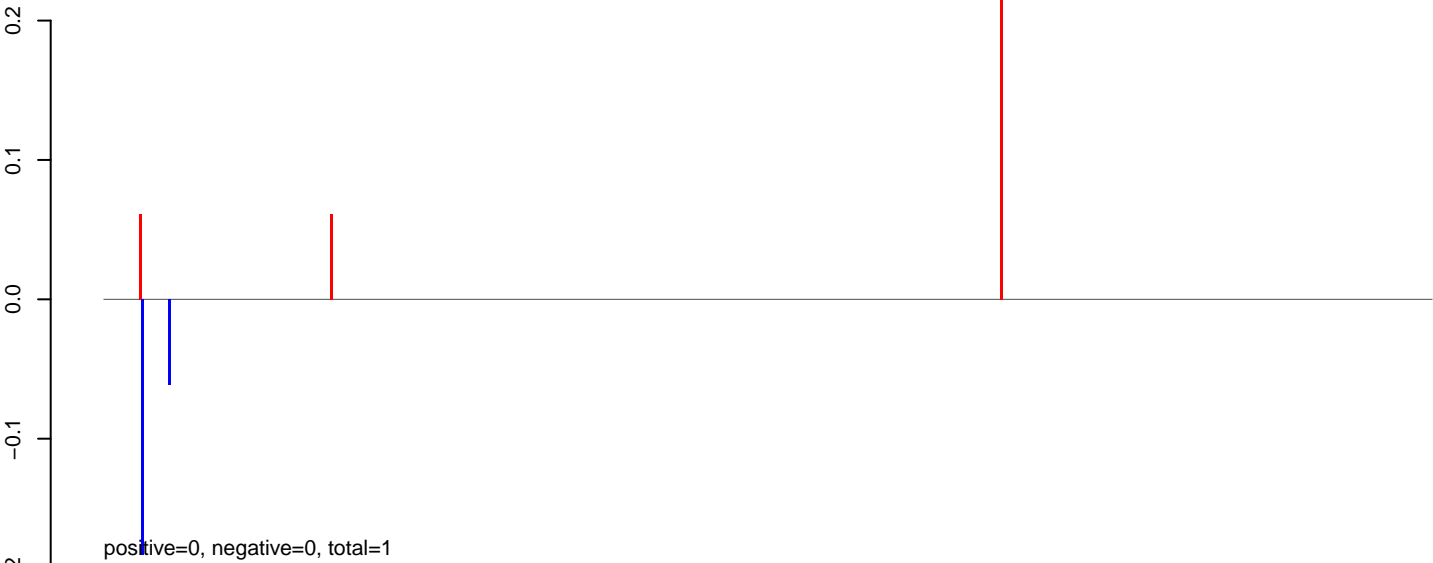
bow size=35, length=11964, n=0, total=1
1-Una_virus_non_structural_polyprotein_gene_partial_cds_and_structural_polyprotein_ge:1-11964

0 2000 4000 6000 8000 10000 12000

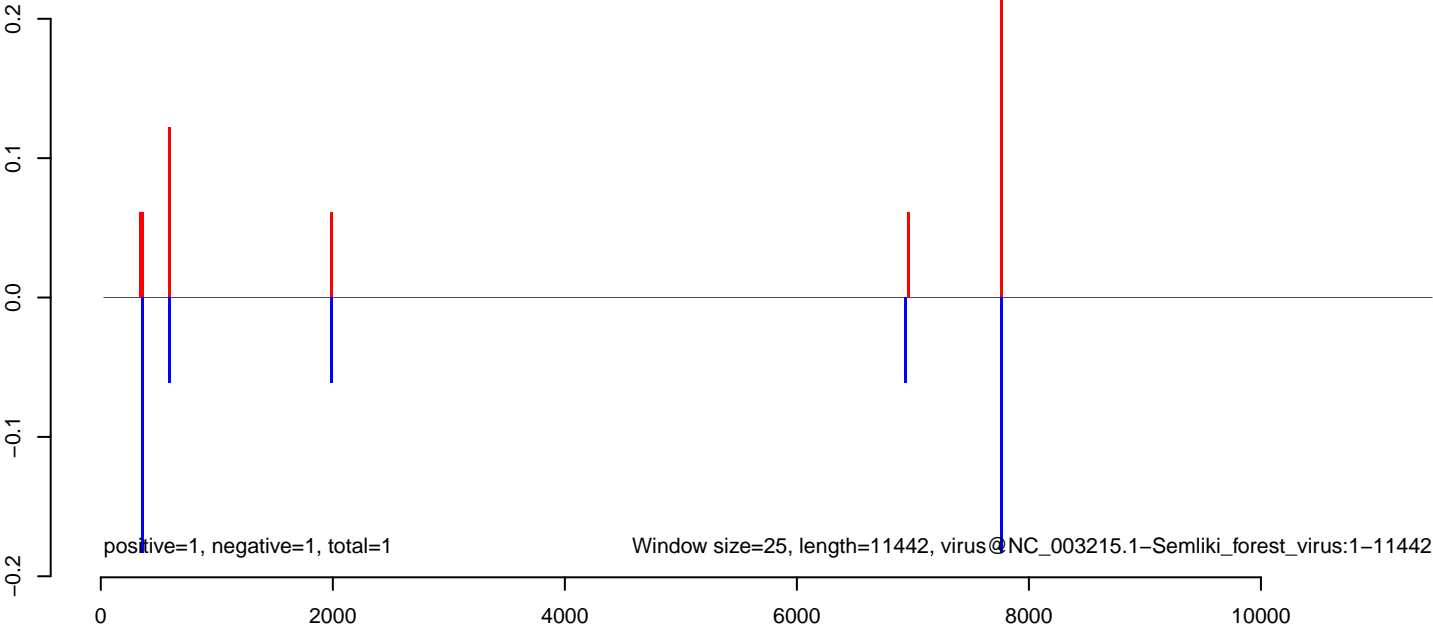
AeAlbo_C636_B2_FHV_r54q.18_23.rep



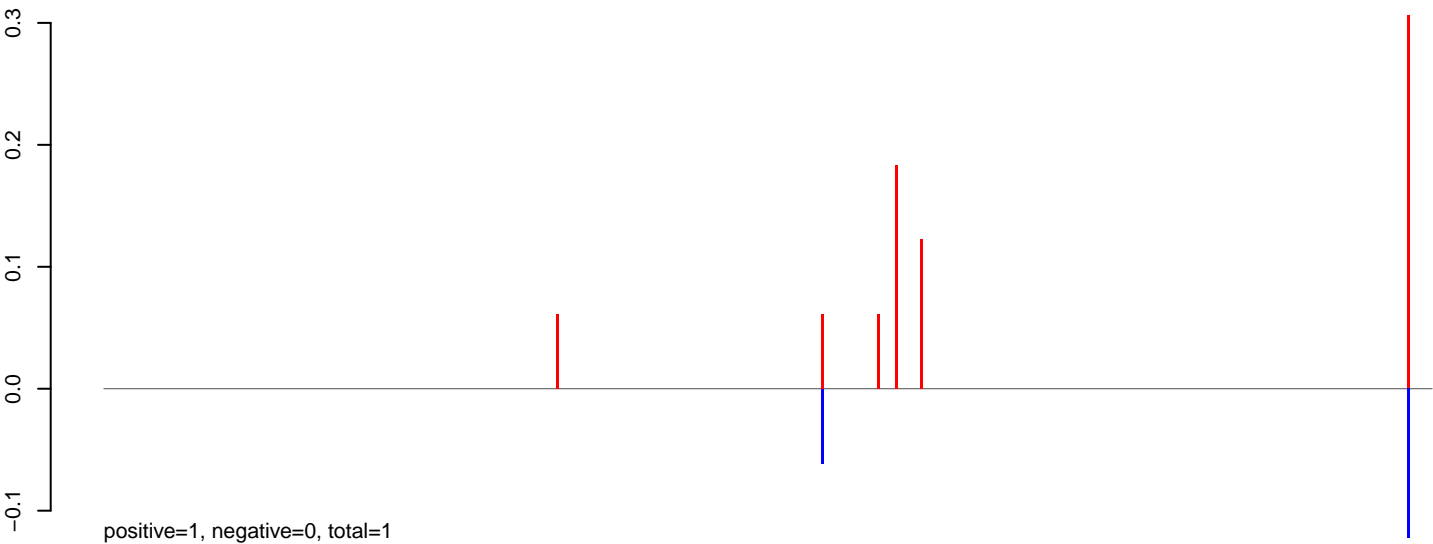
AeAlbo_C636_B2_FHV_r54q.24_35.rep



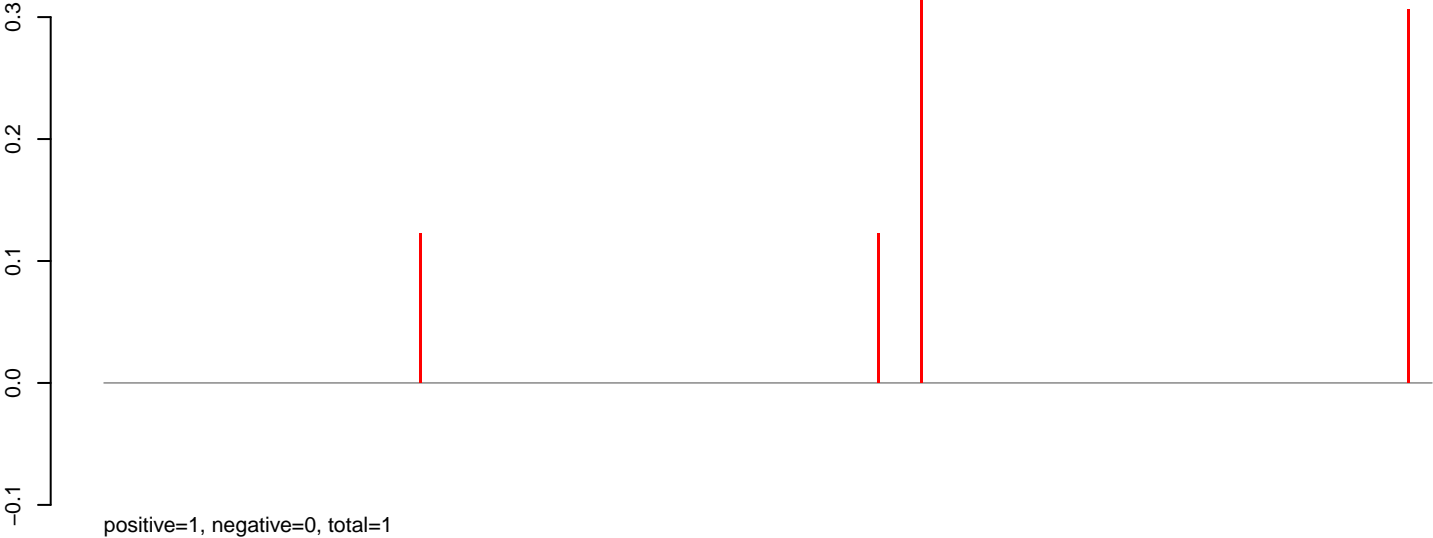
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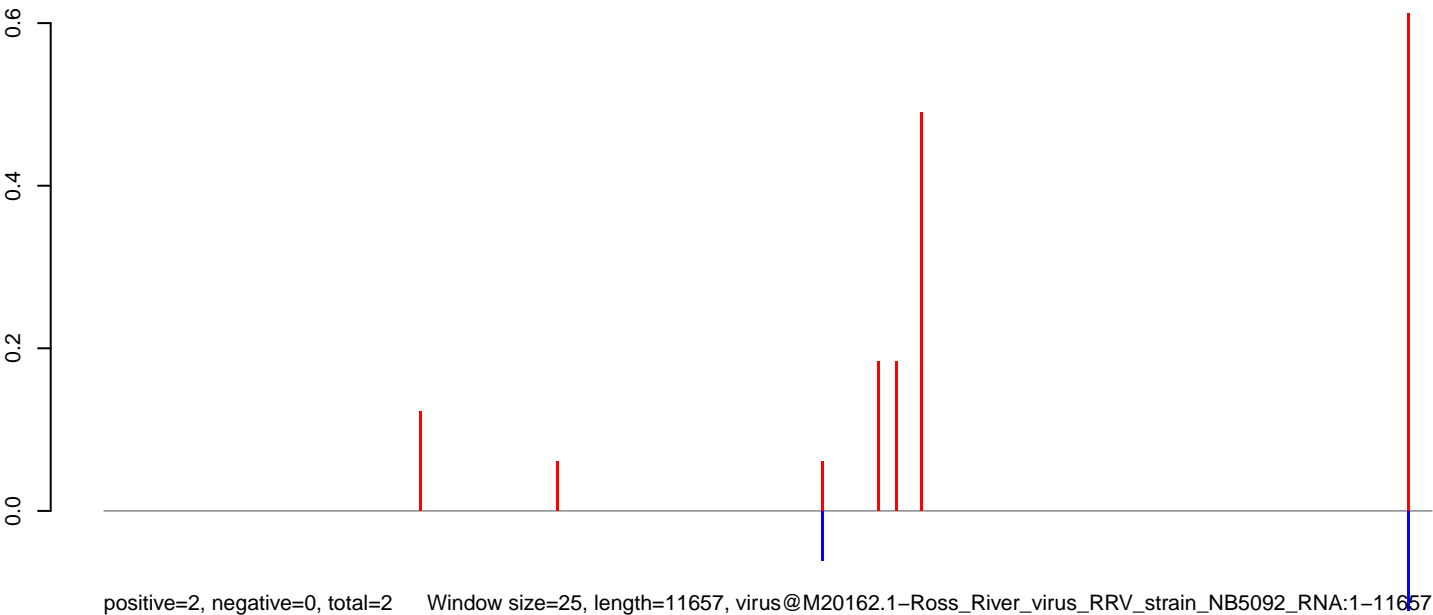
AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep



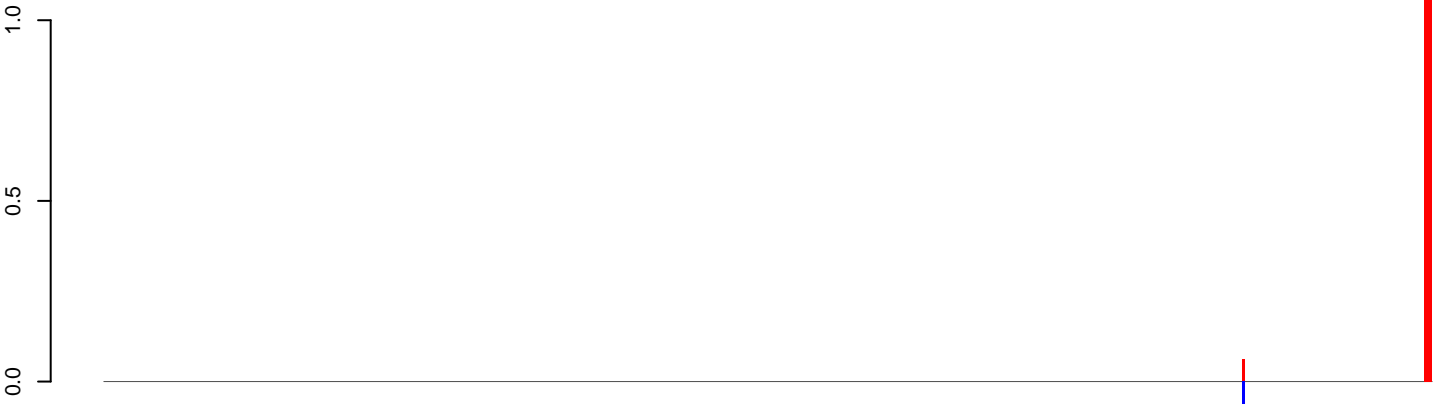
AeAlbo_C636_B2_FHV_r54q.rep



Window size=25, length=11657, virus@M20162.1-Ross_River_virus_RRV_strain_NB5092_RNA:1-11657

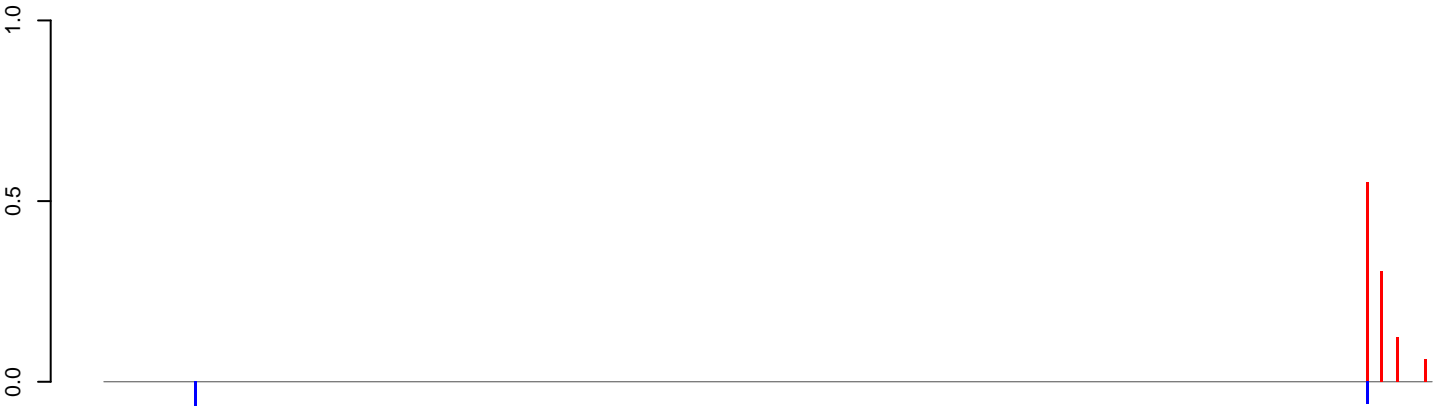
0 2000 4000 6000 8000 10000 12000

AeAlbo_C636_B2_FHV_r54q.18_23.rep



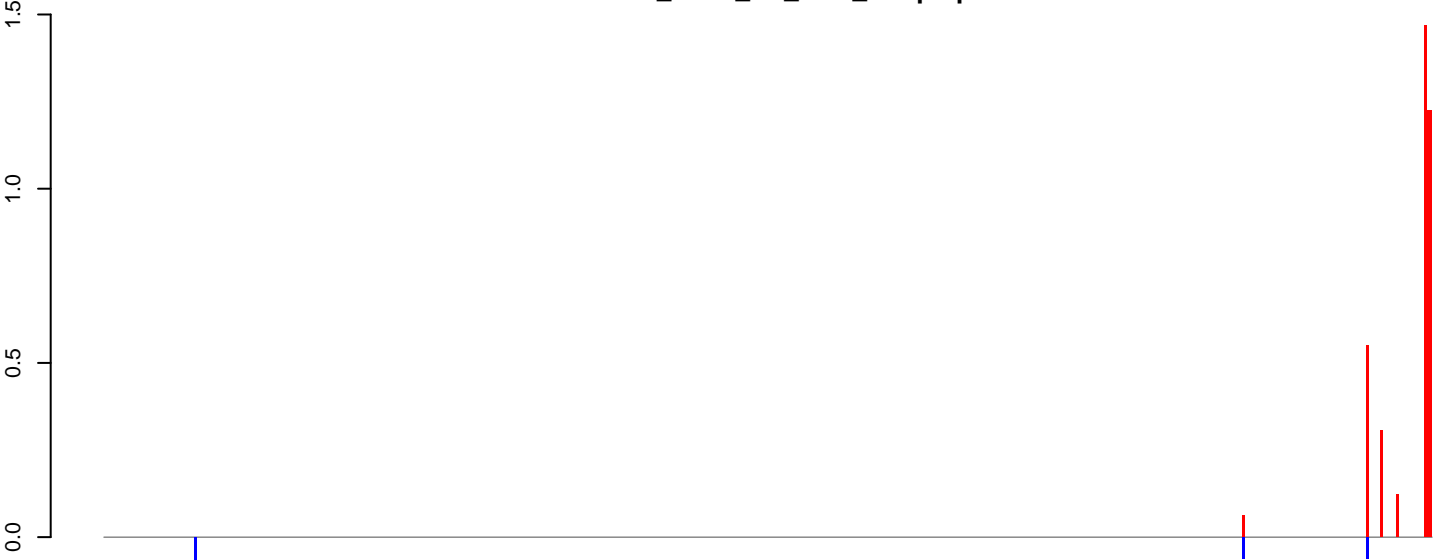
positive=4, negative=0, total=4

AeAlbo_C636_B2_FHV_r54q.24_35.rep



positive=1, negative=0, total=1

AeAlbo_C636_B2_FHV_r54q.rep



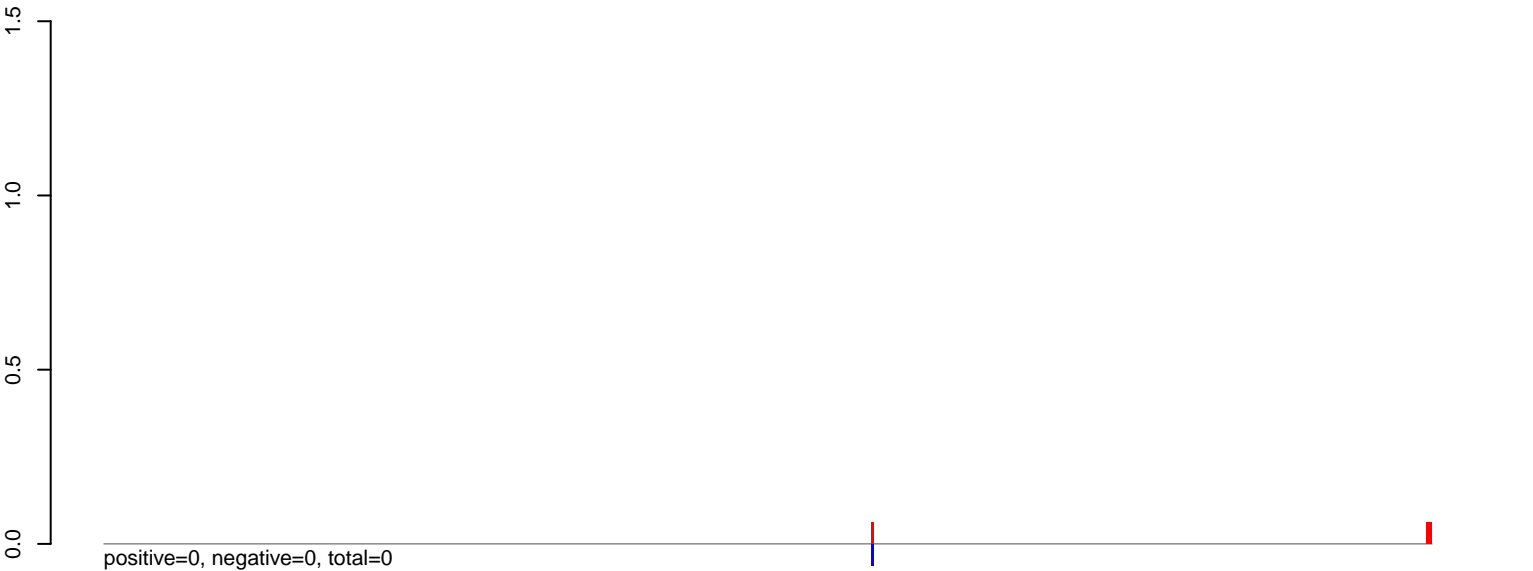
downloaded from <https://www.ncbi.nlm.nih.gov/trace/viral/147991.1/> Trocara_virus_non_structural_polyprotein_gene_partial_cds_and_structural_polyprotei:1-12052

0 2000 4000 6000 8000 10000 12000

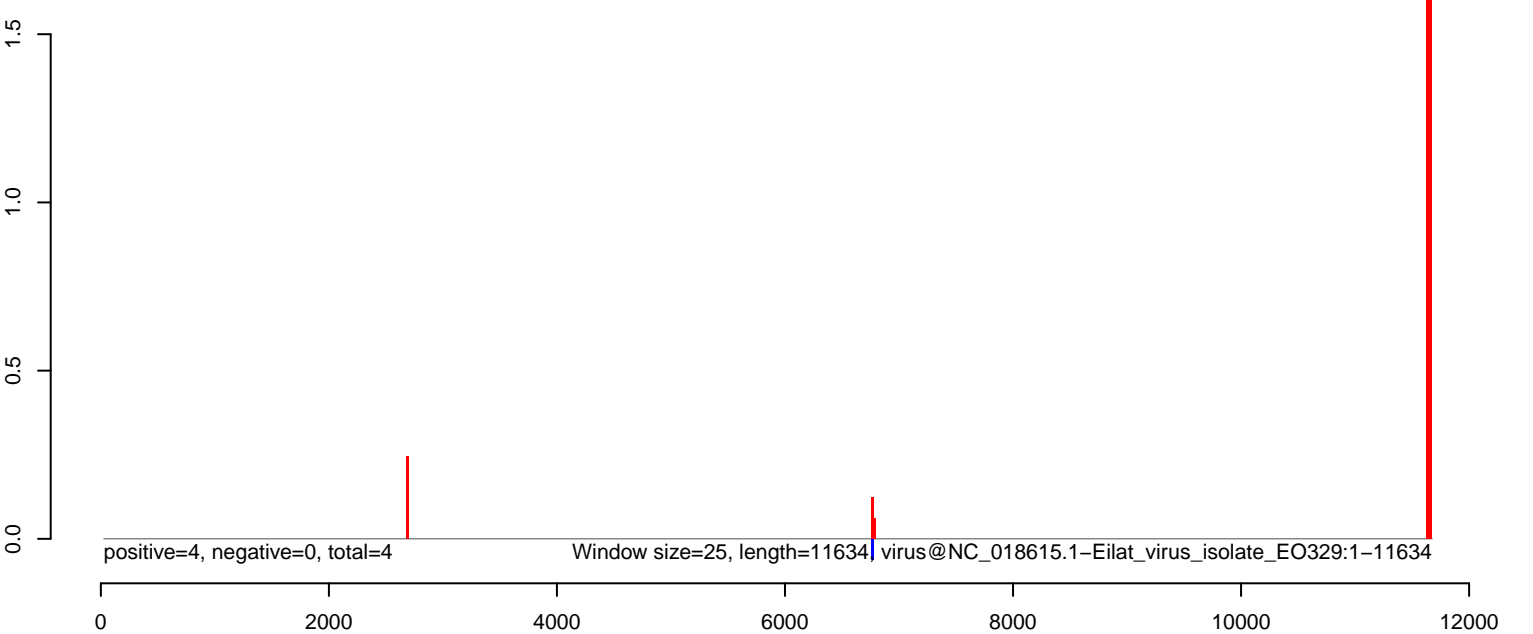
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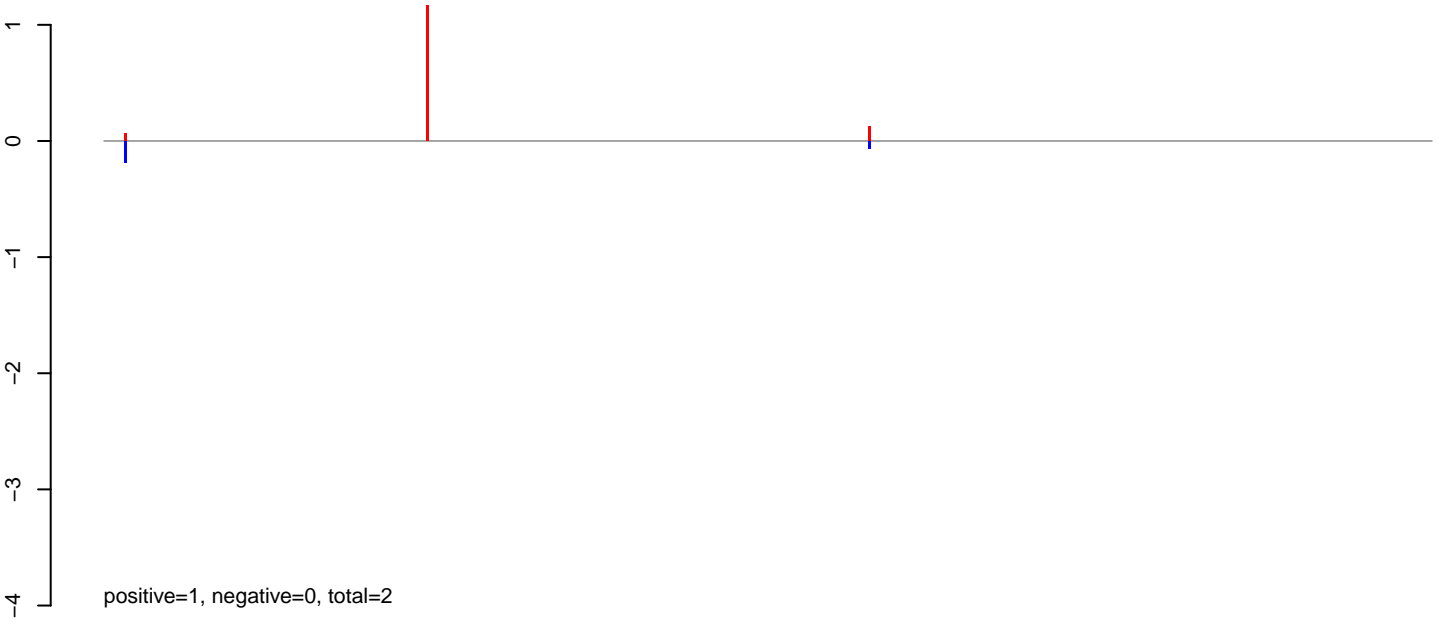
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AeAlbo_C636_B2_FHV_r54q.rep



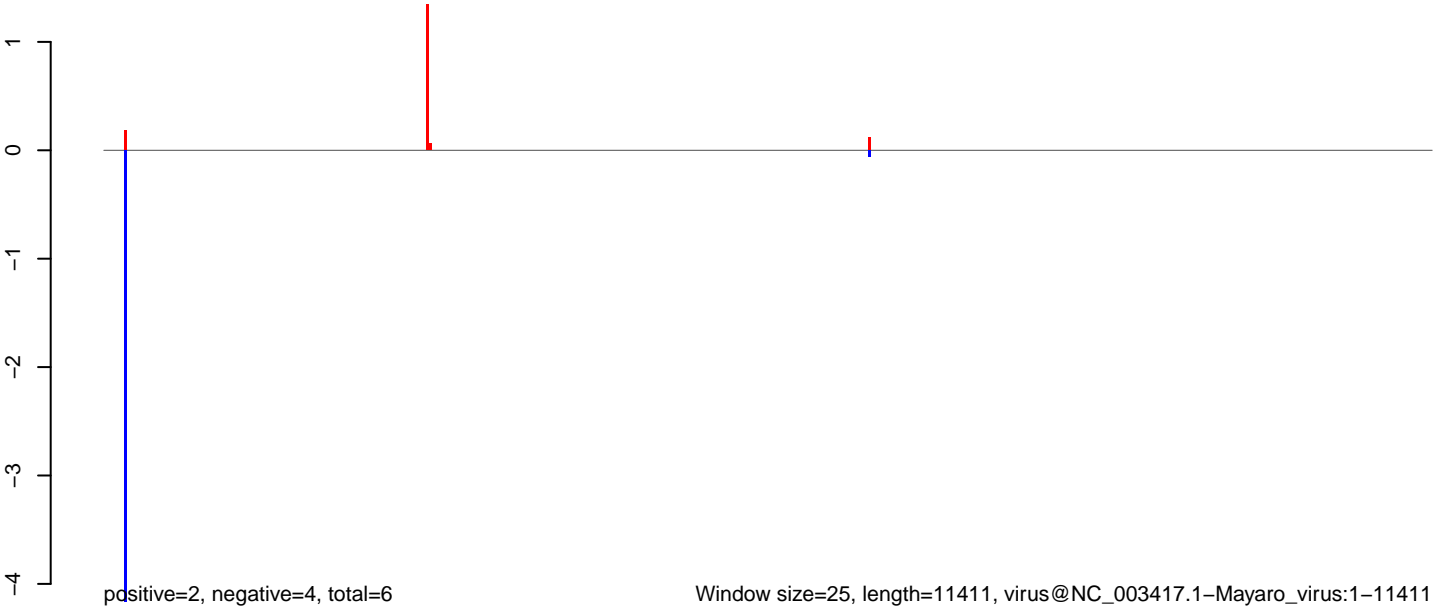
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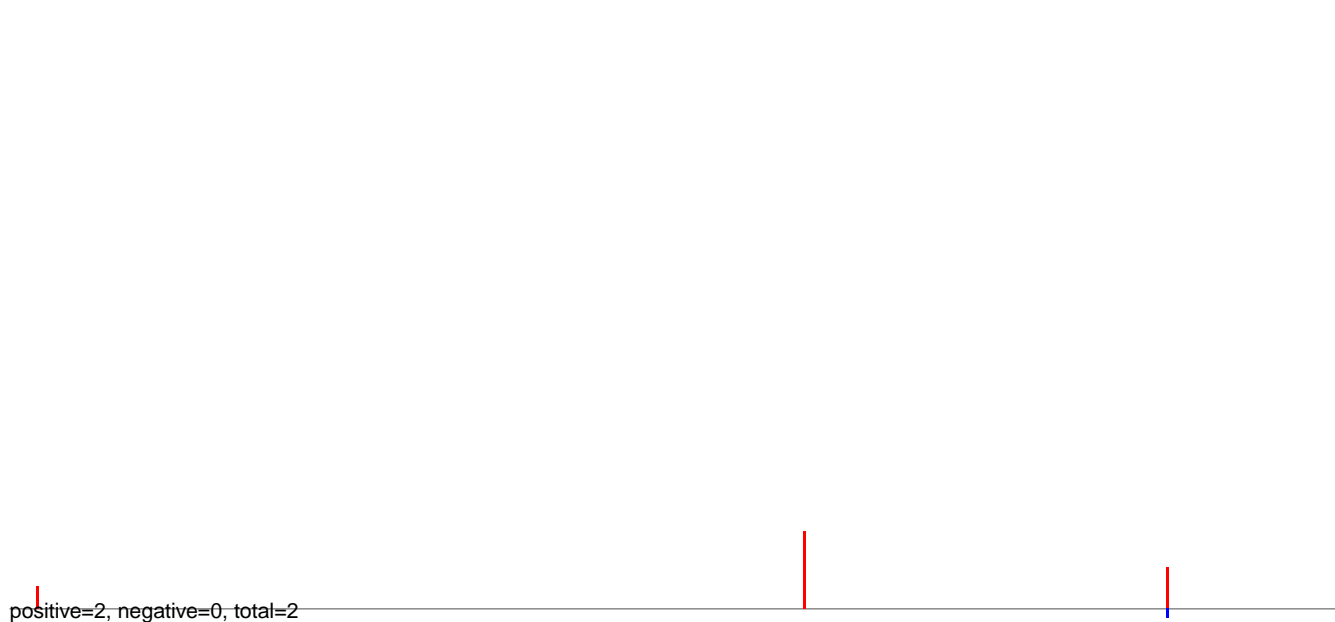
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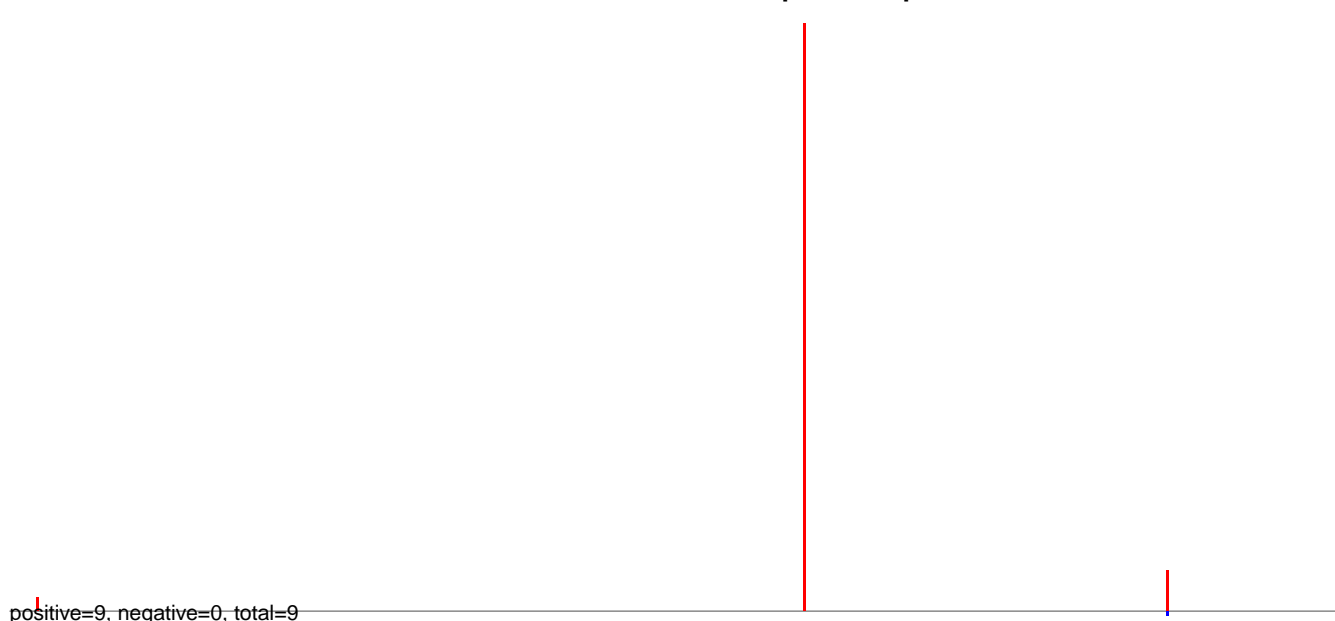
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AeAlbo_C636_B2_FHV_r54q.18_23.rep



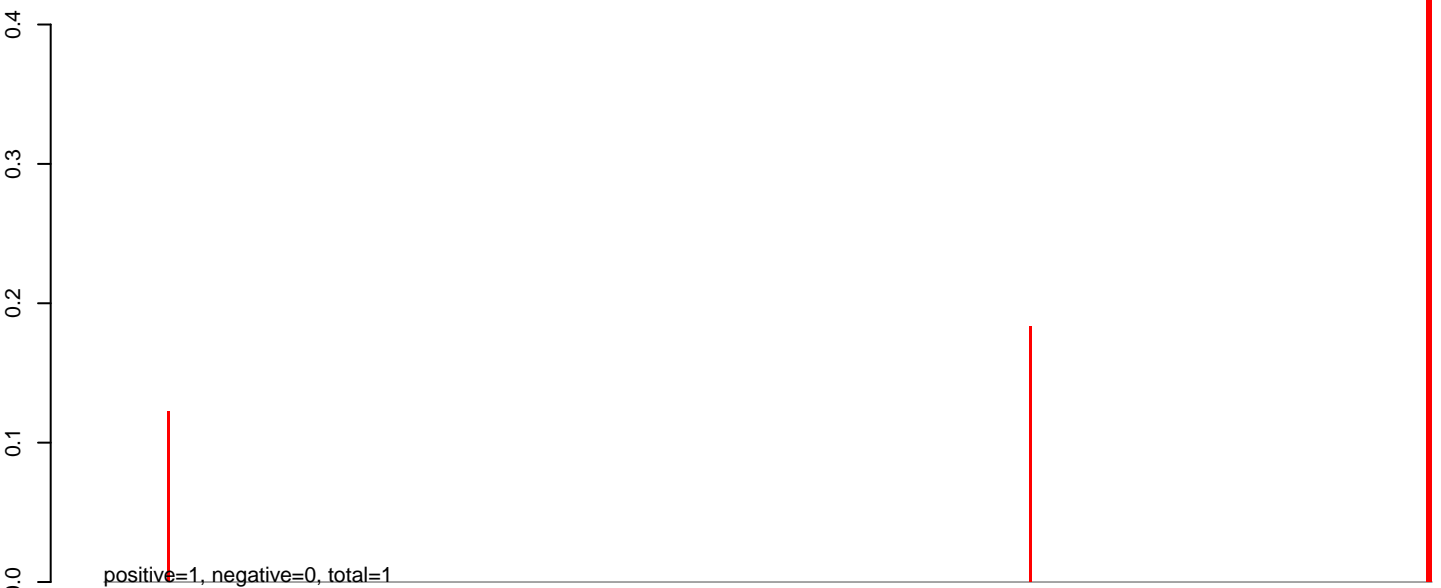
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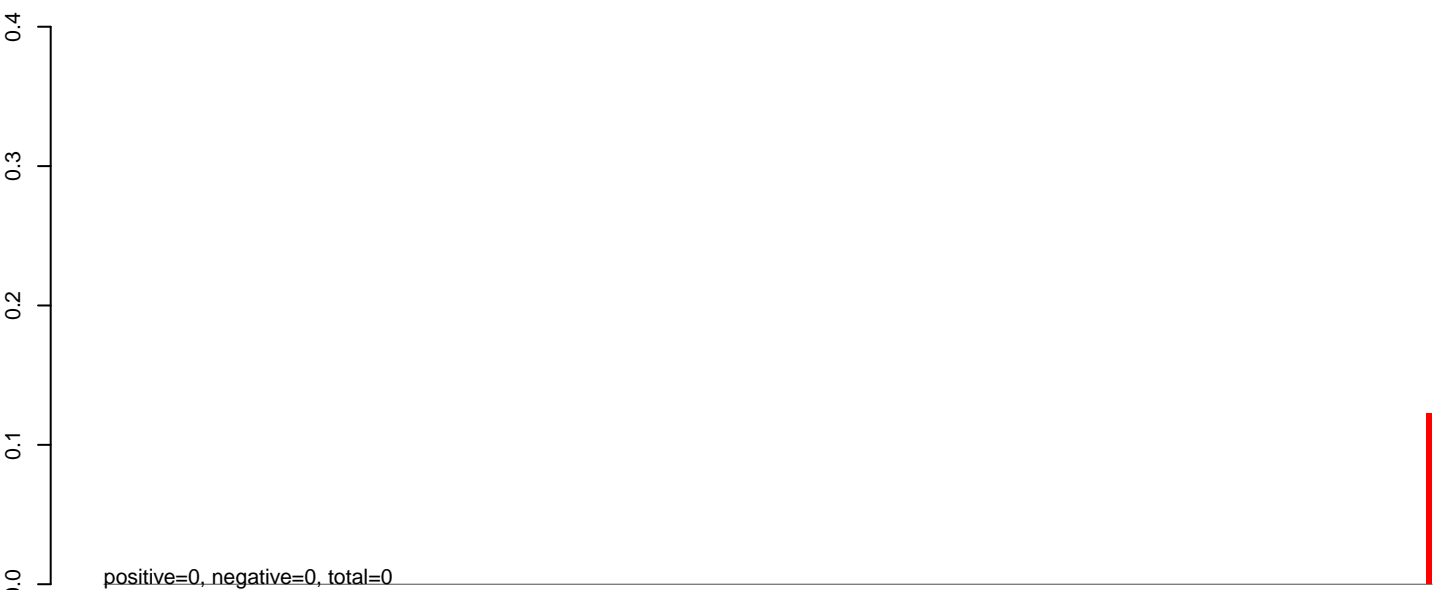
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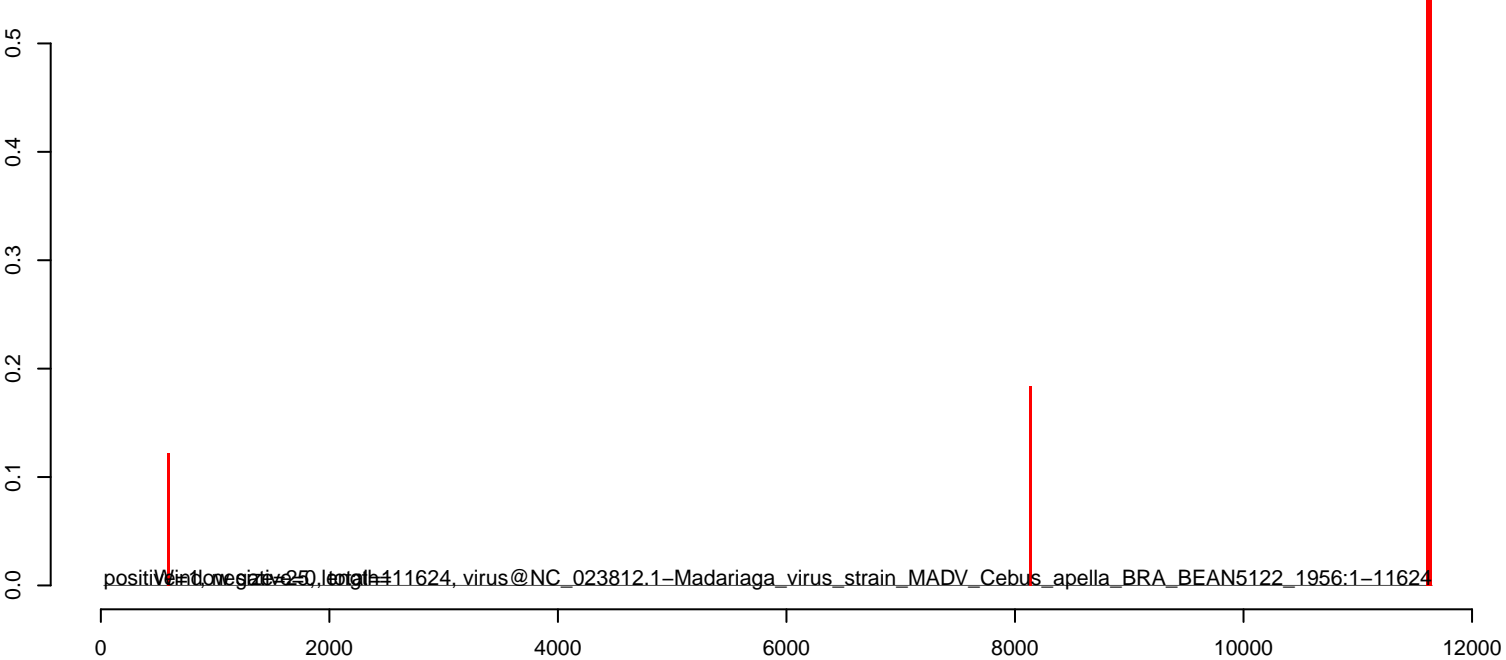
AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep



AeAlbo_C636_B2_FHV_r54q.rep



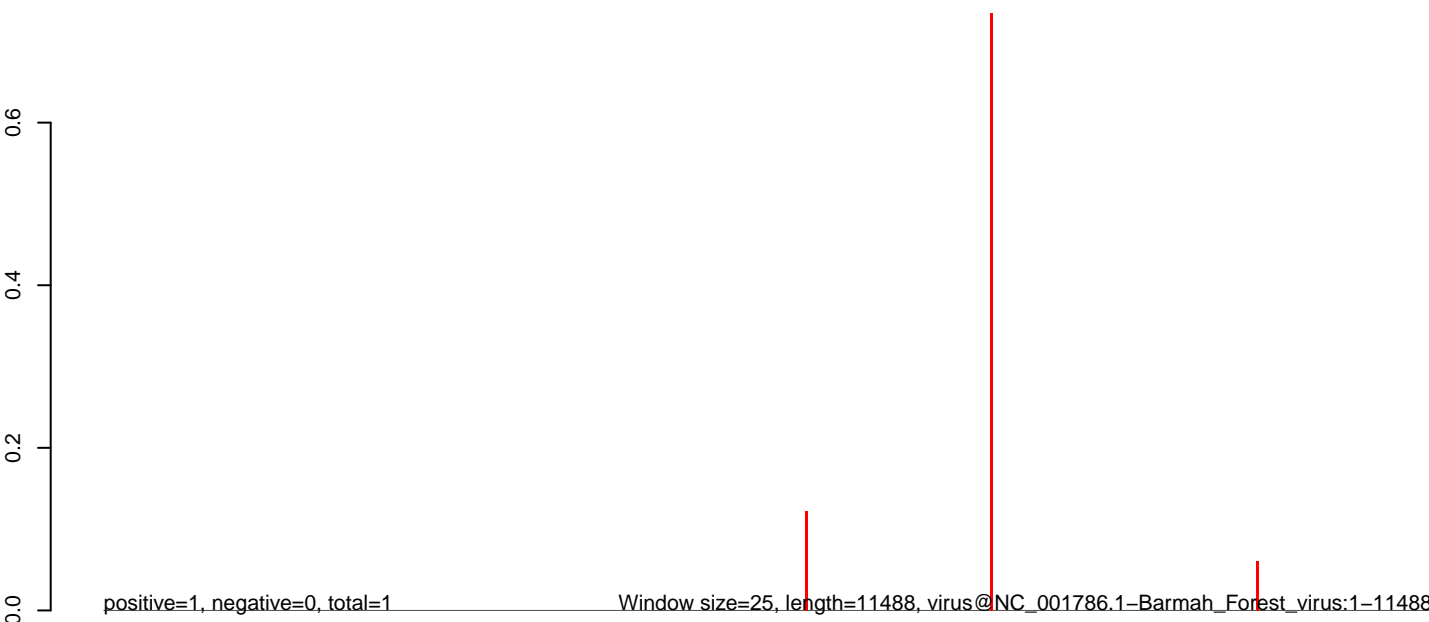
AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep



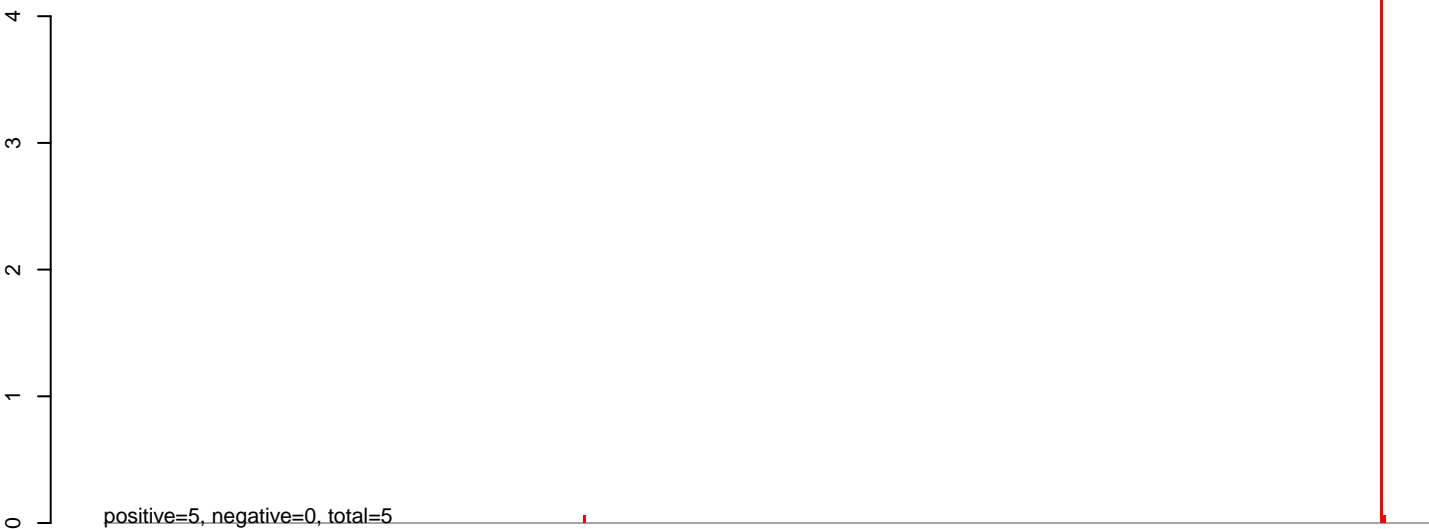
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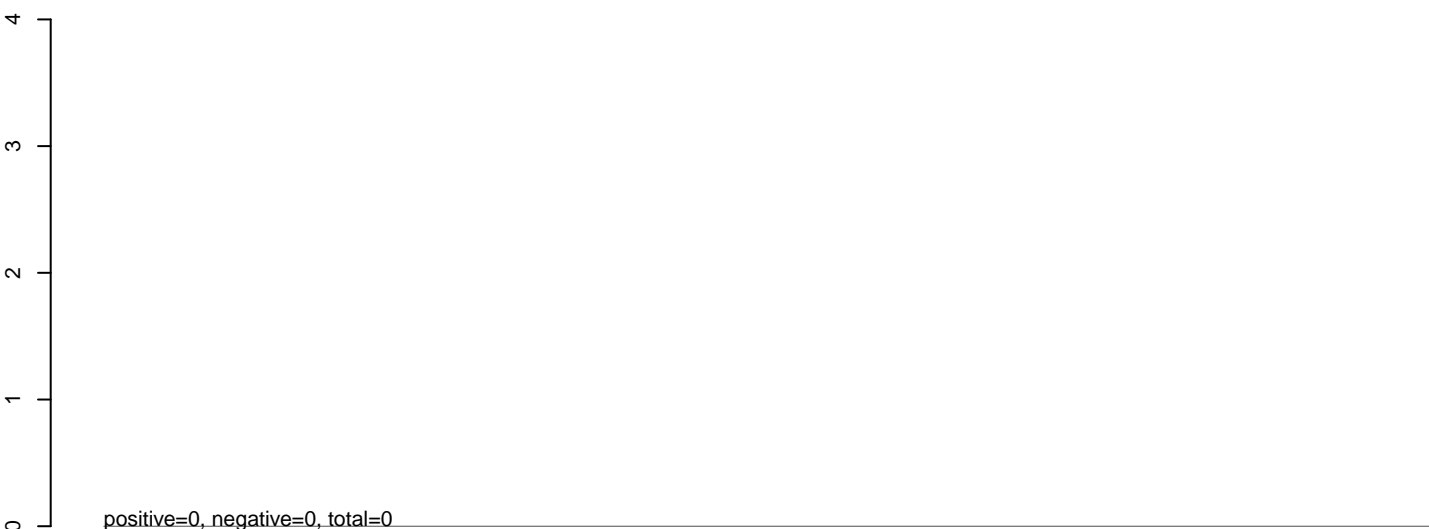
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0 2000 4000 6000 8000 10000

AeAlbo_C636_B2_FHV_r54q.18_23.rep



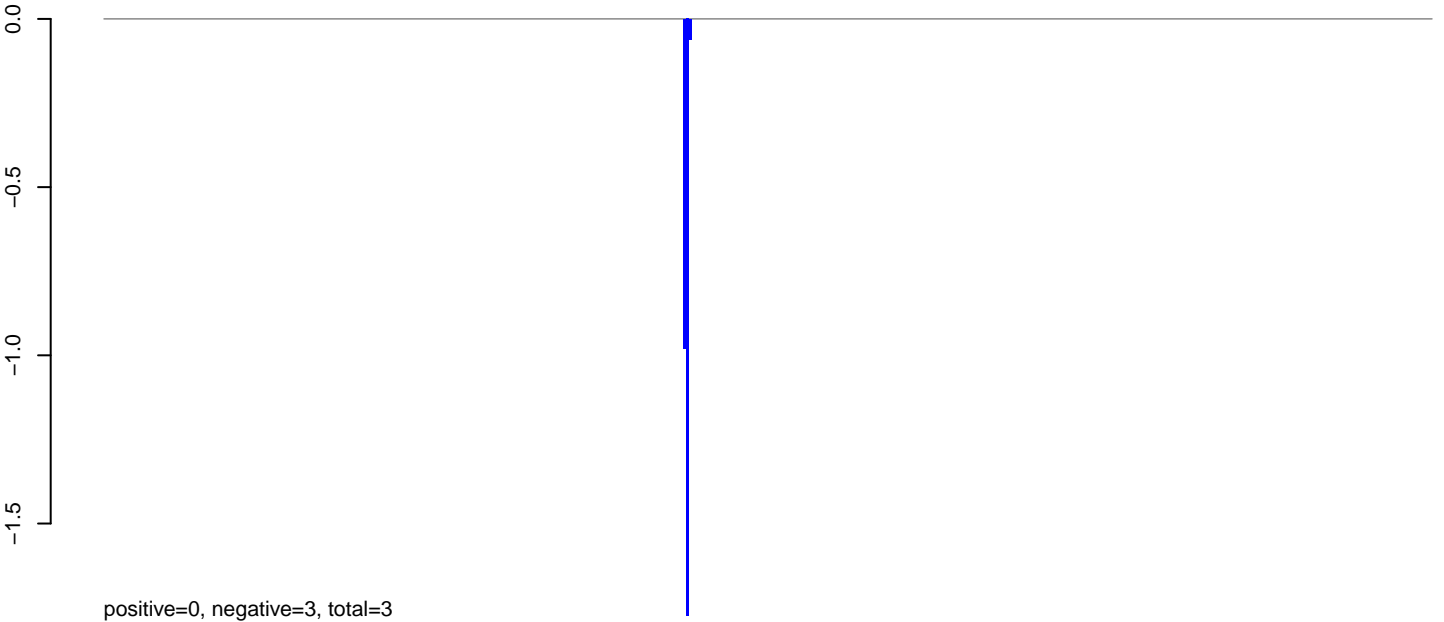
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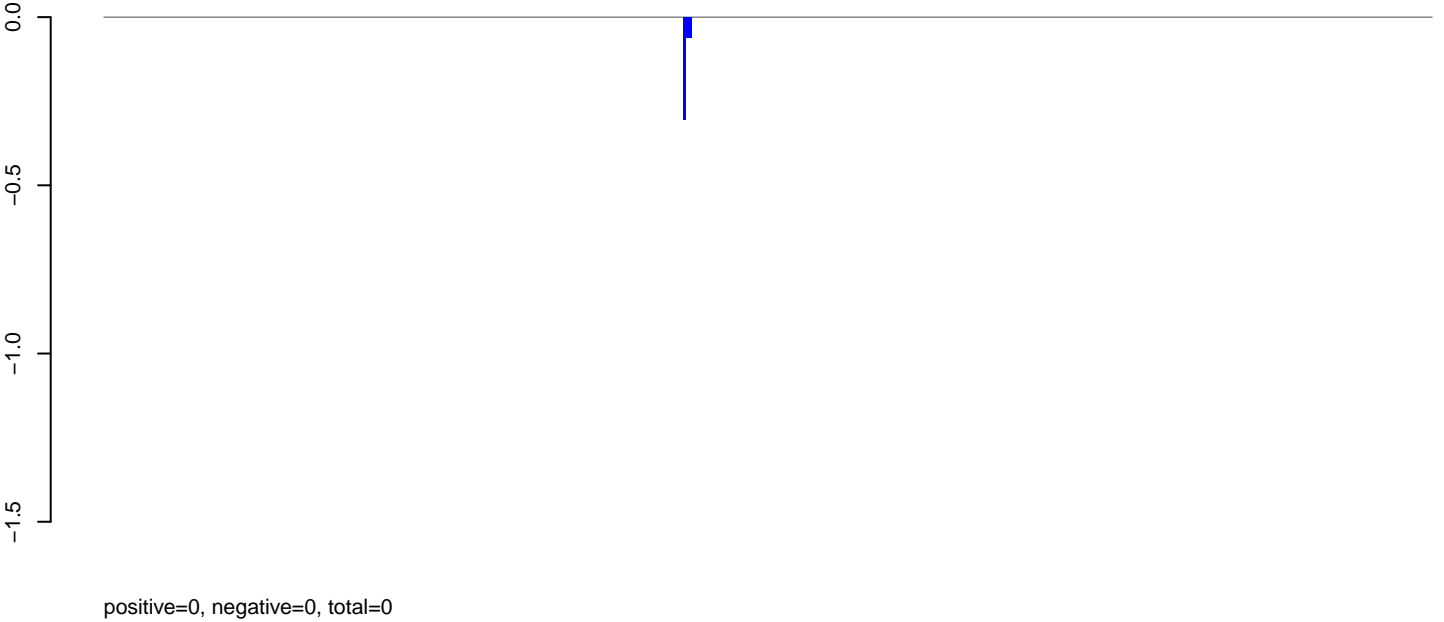
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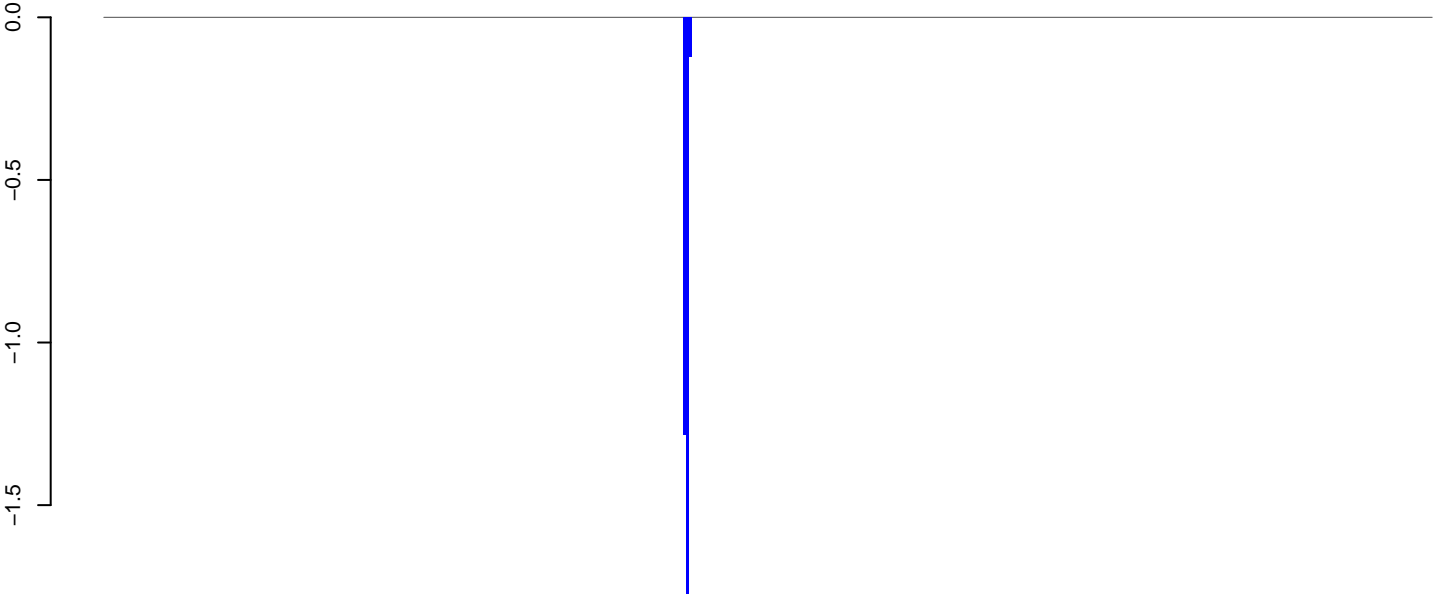
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AeAlbo_C636_B2_FHV_r54q.24_35.rep

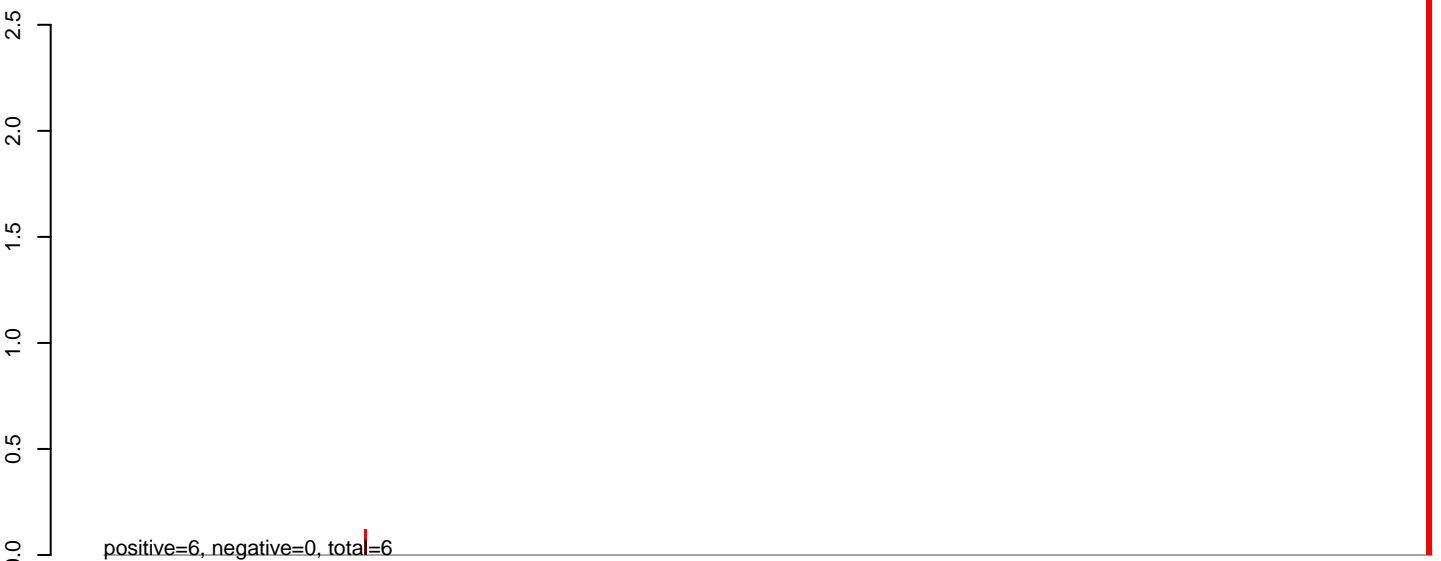


AeAlbo_C636_B2_FHV_r54q.rep

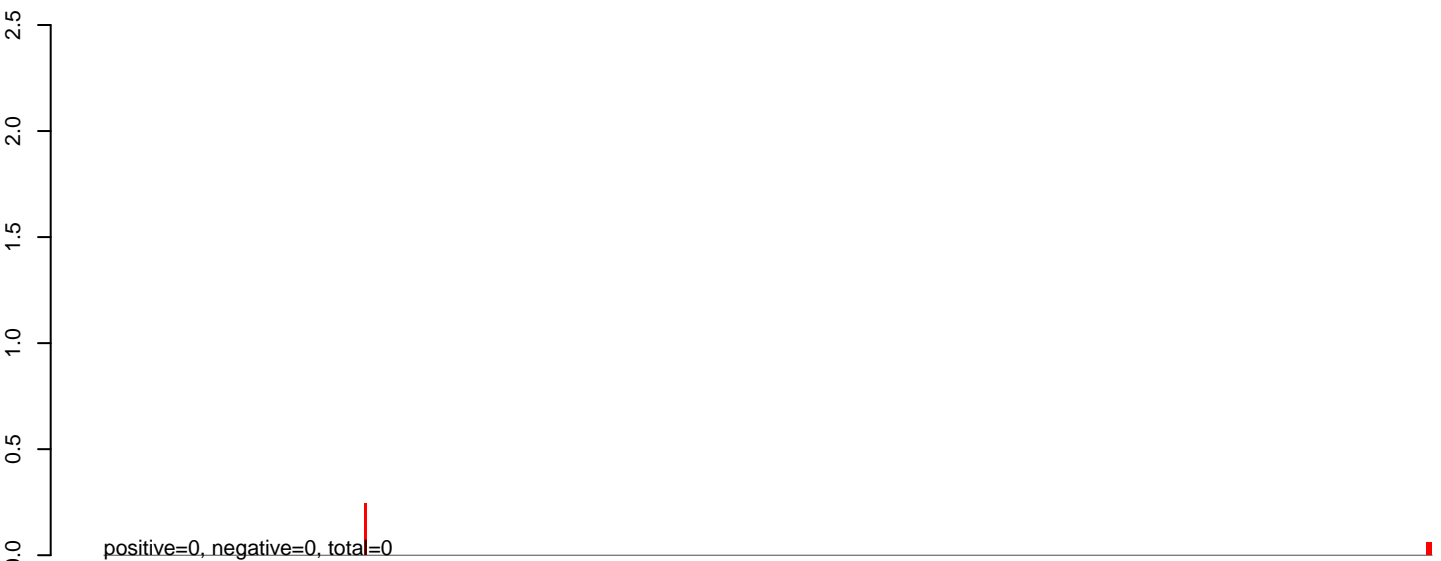


0 2000 4000 6000 8000 10000 12000

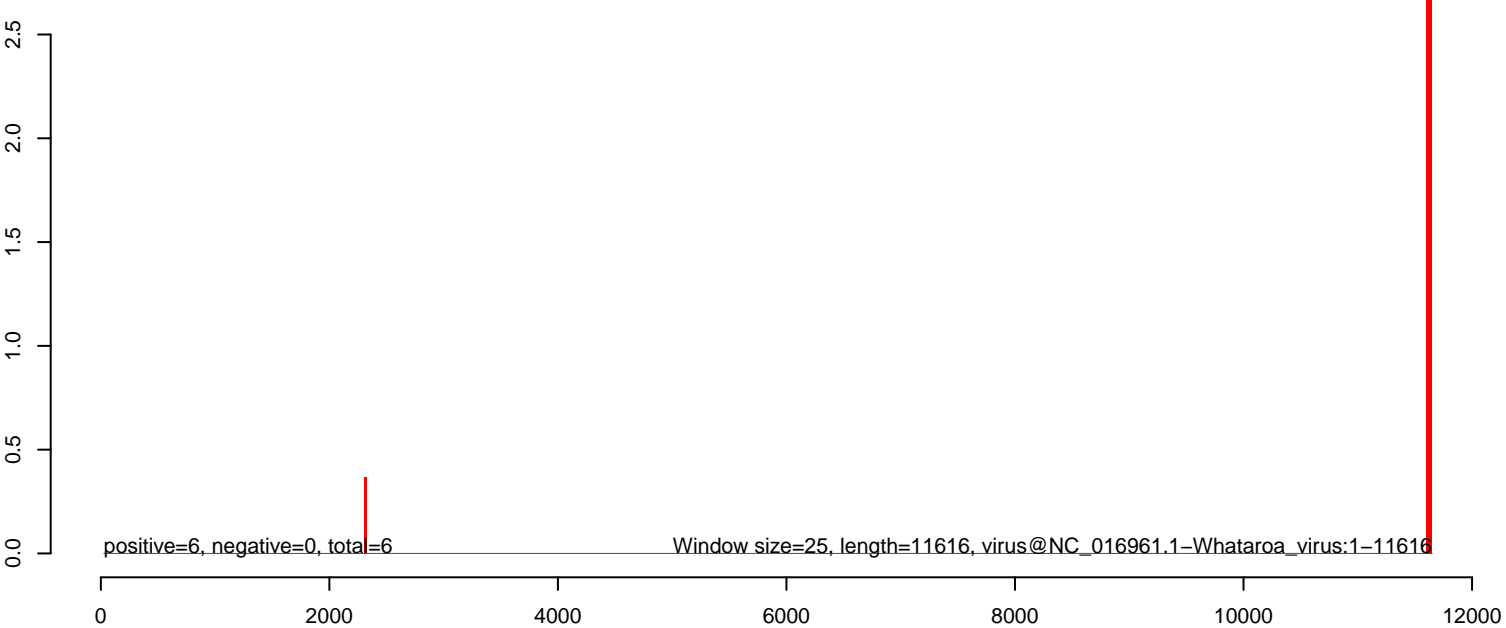
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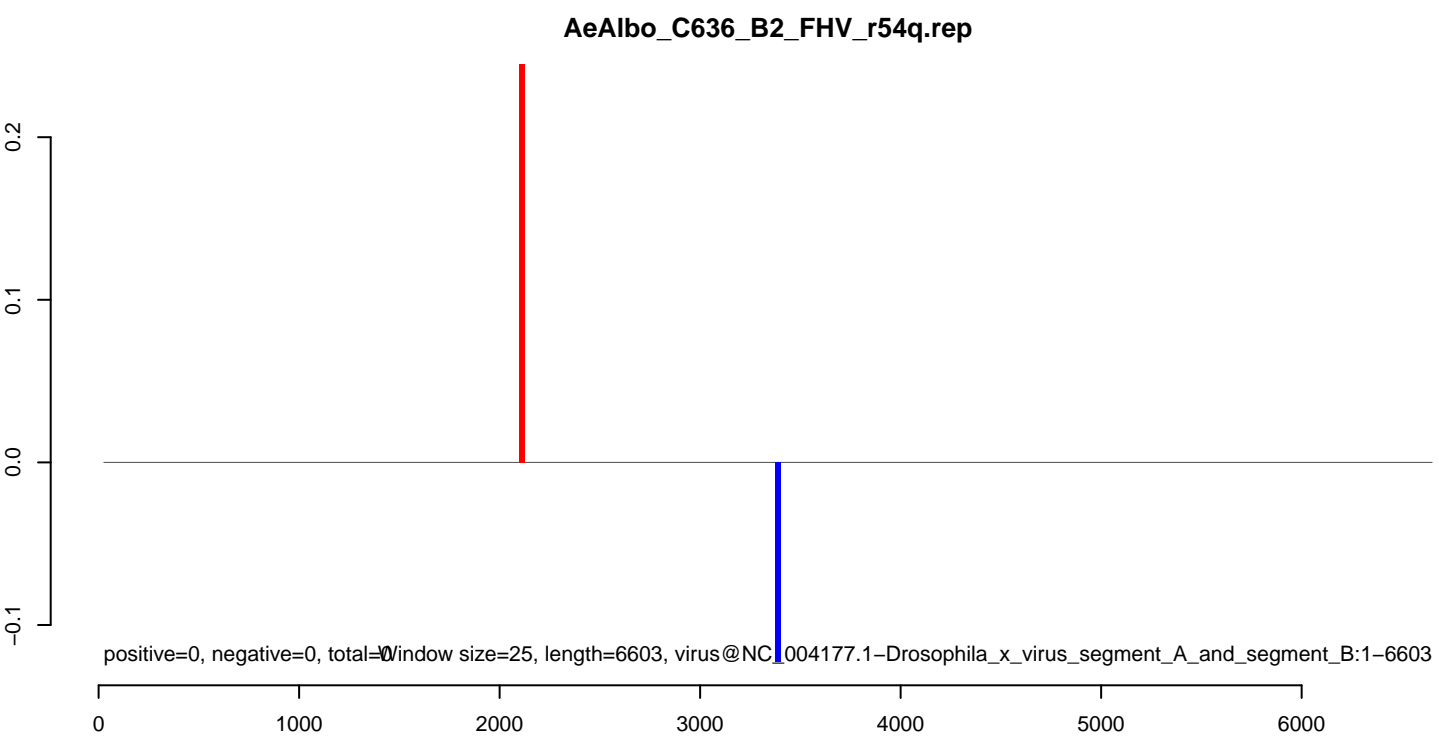
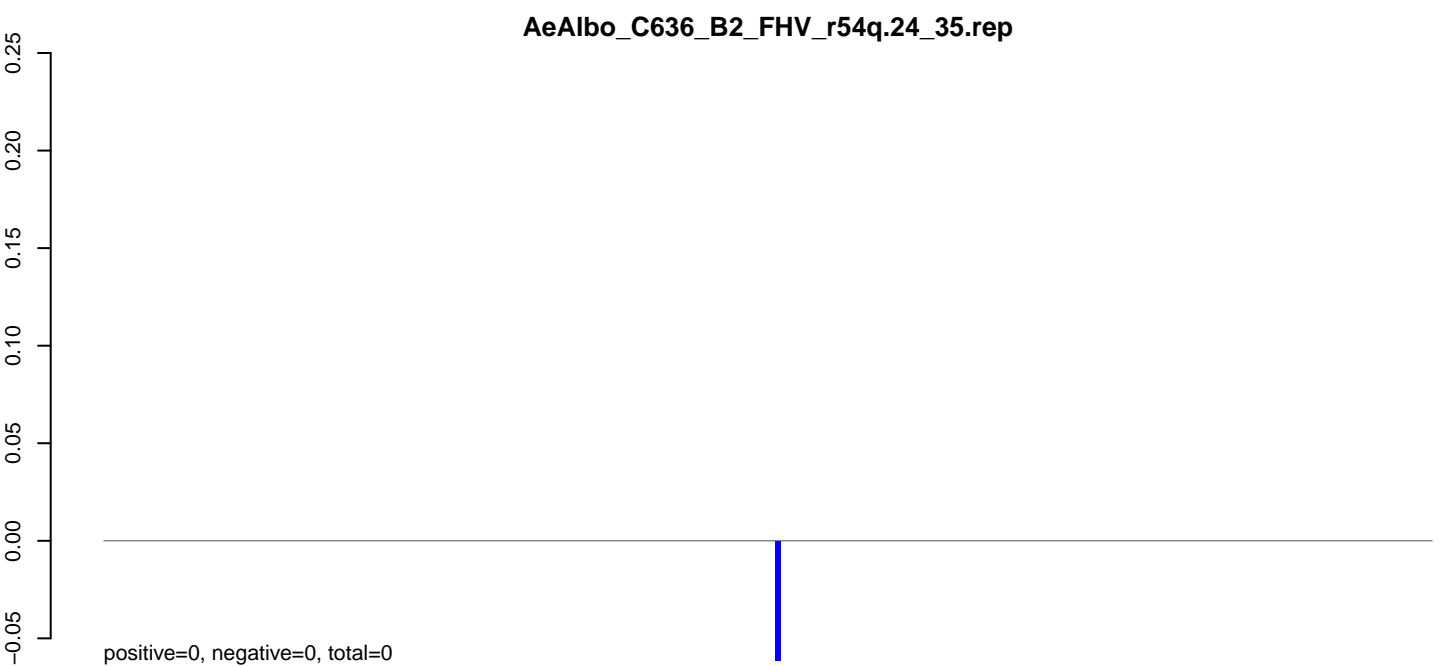
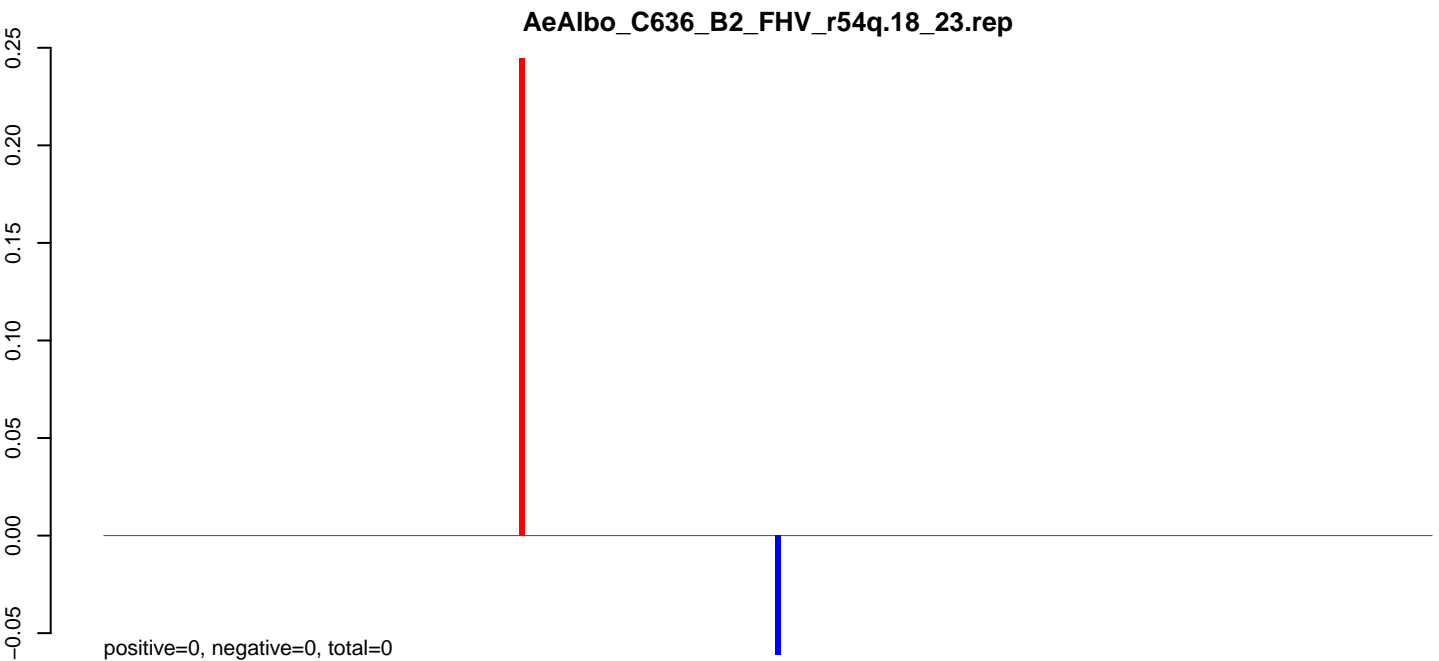


AeAlbo_C636_B2_FHV_r54q.24_35.rep

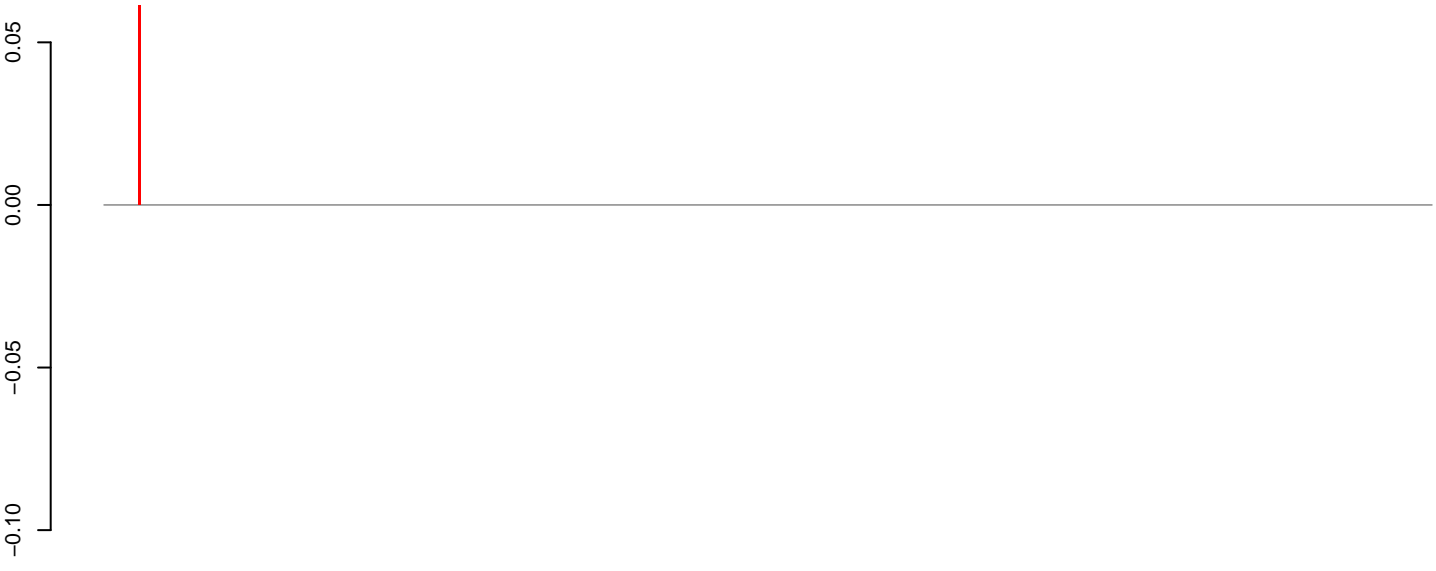


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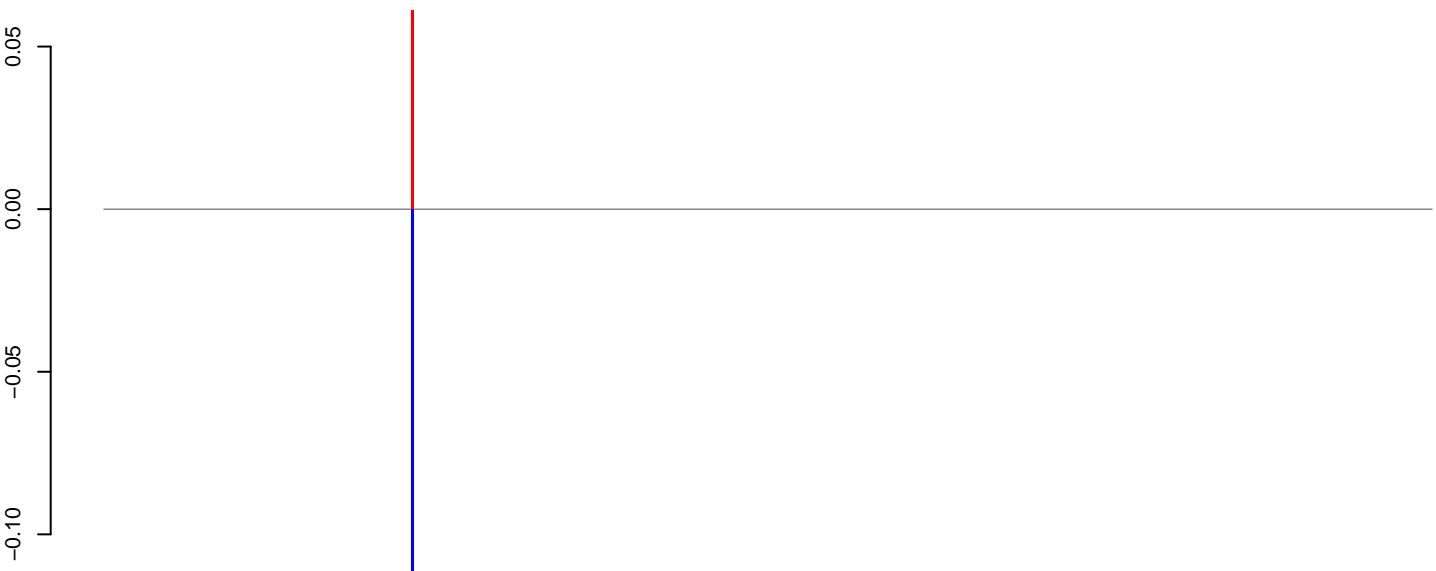


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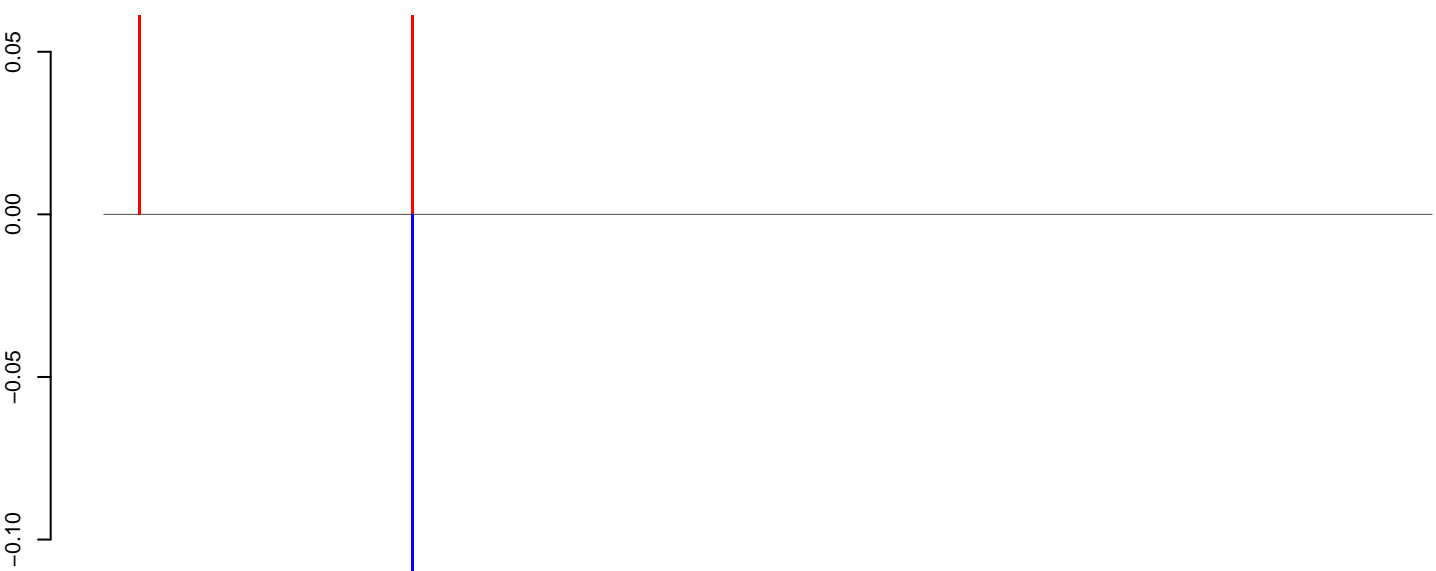
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AeAlbo_C636_B2_FHV_r54q.24_35.rep



positive=0, negative=0, total=0

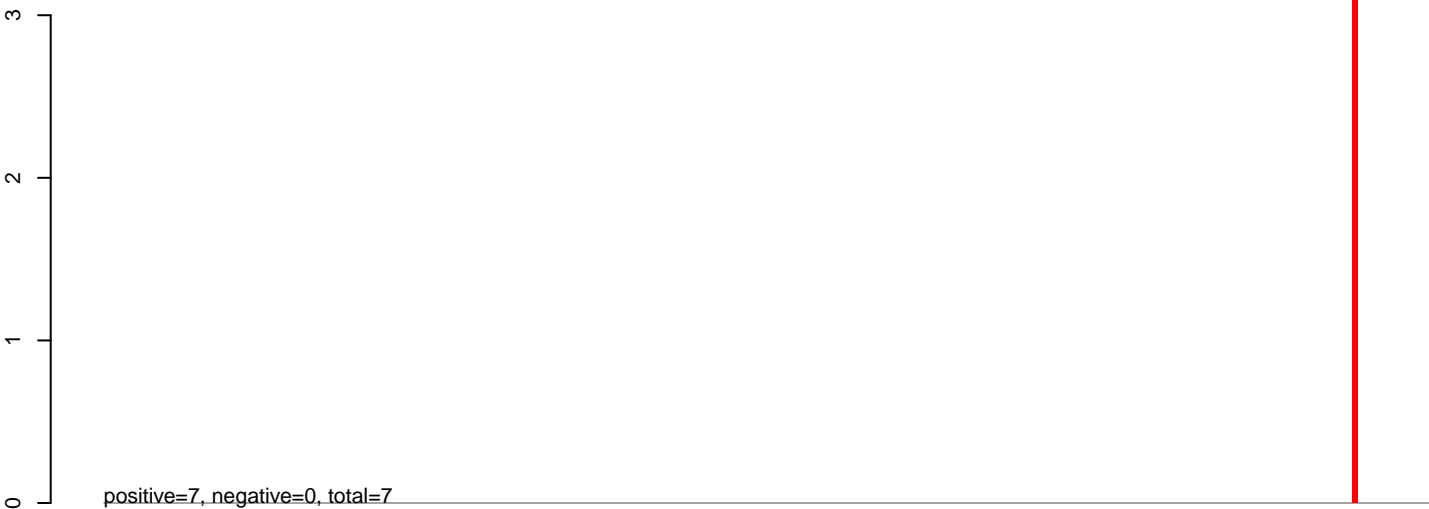
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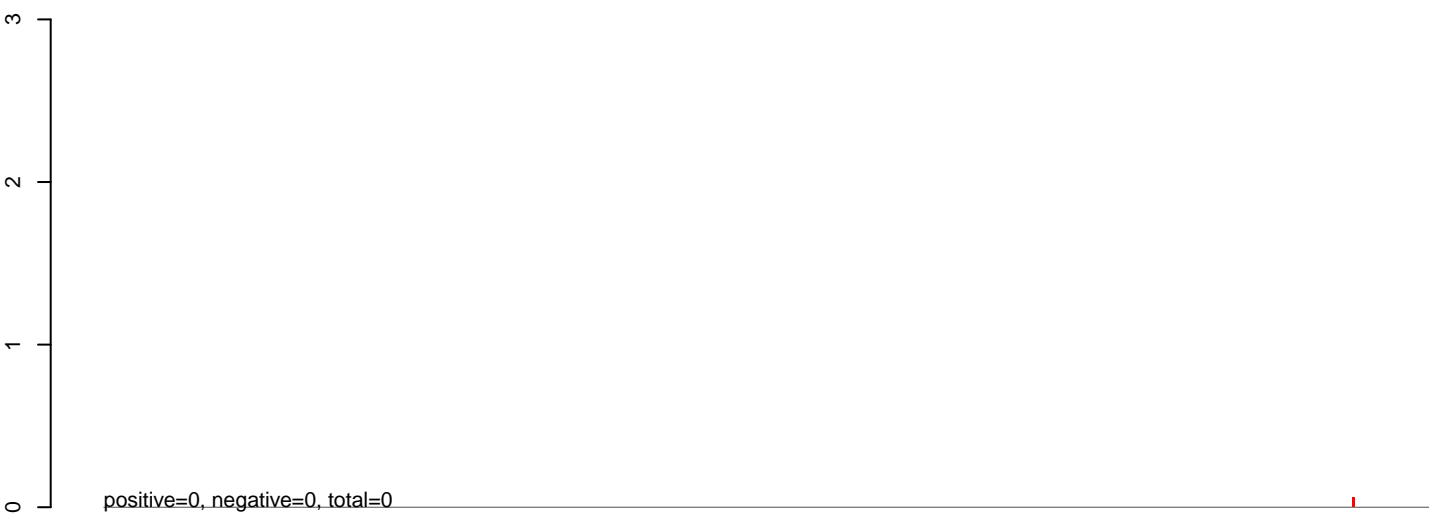
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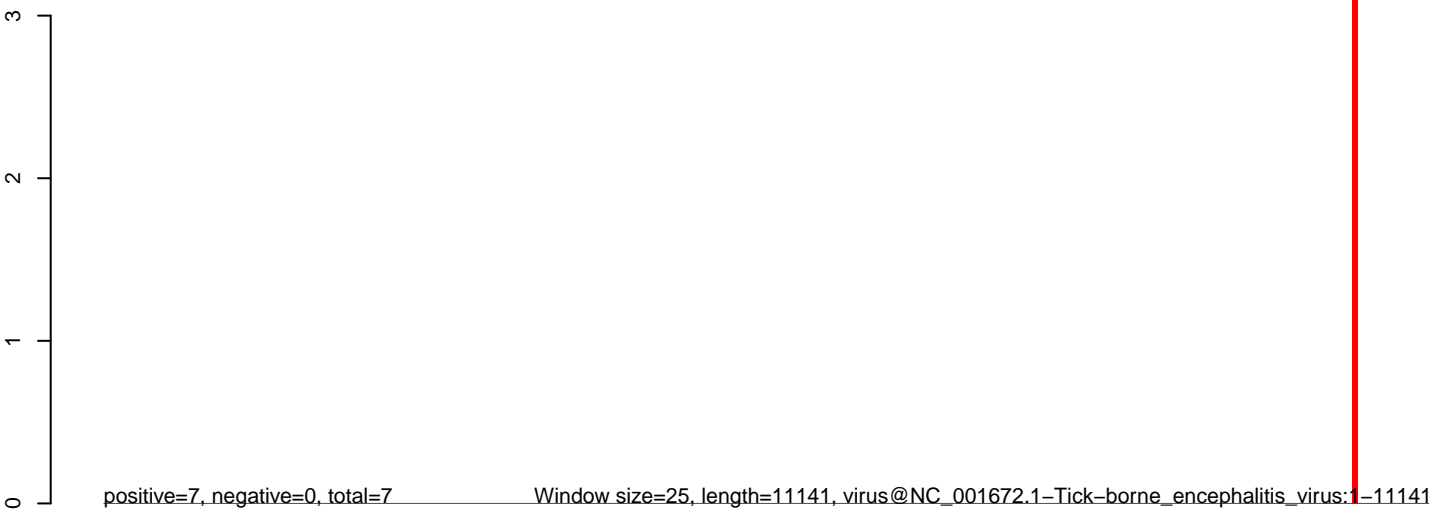
AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep

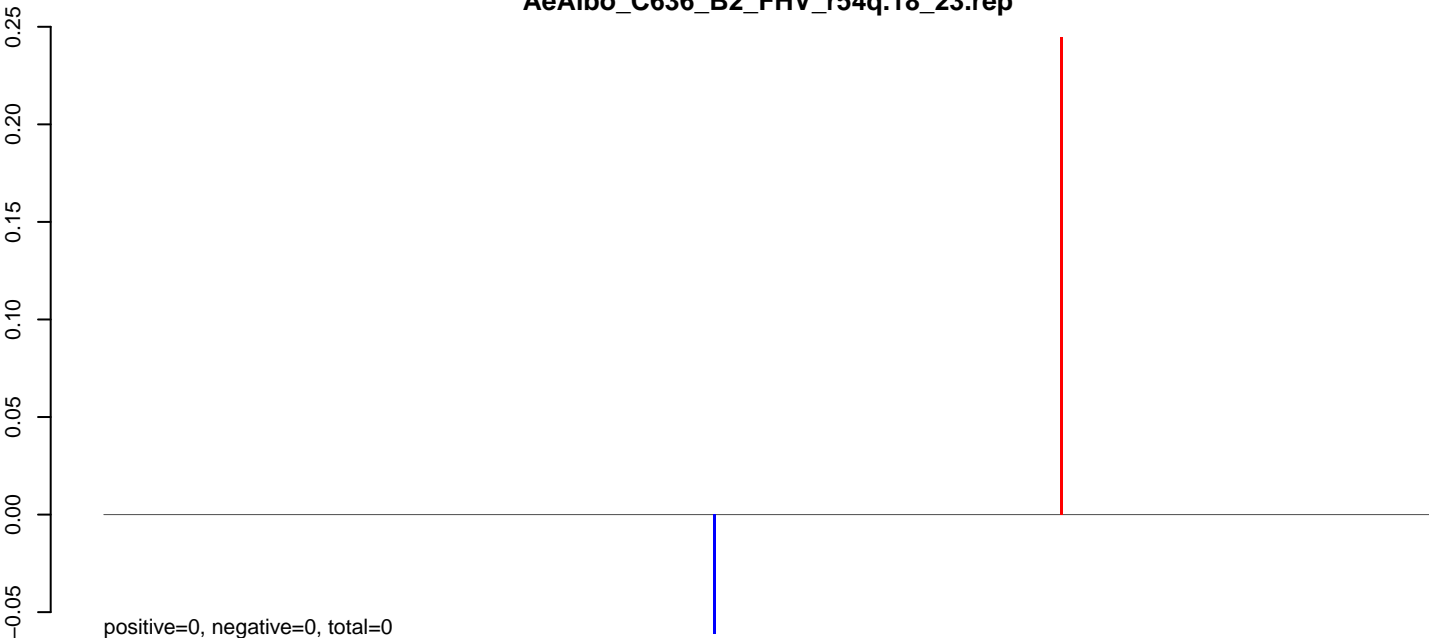


AeAlbo_C636_B2_FHV_r54q.rep



0 2000 4000 6000 8000 10000

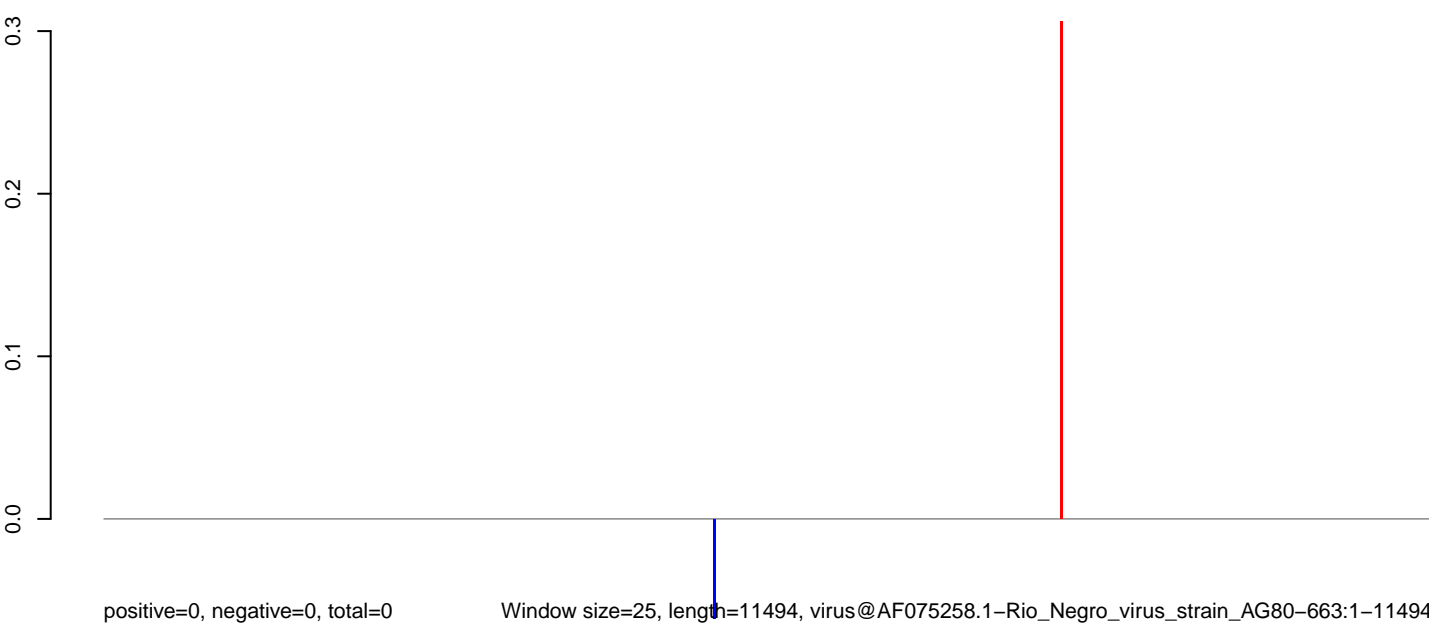
AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep



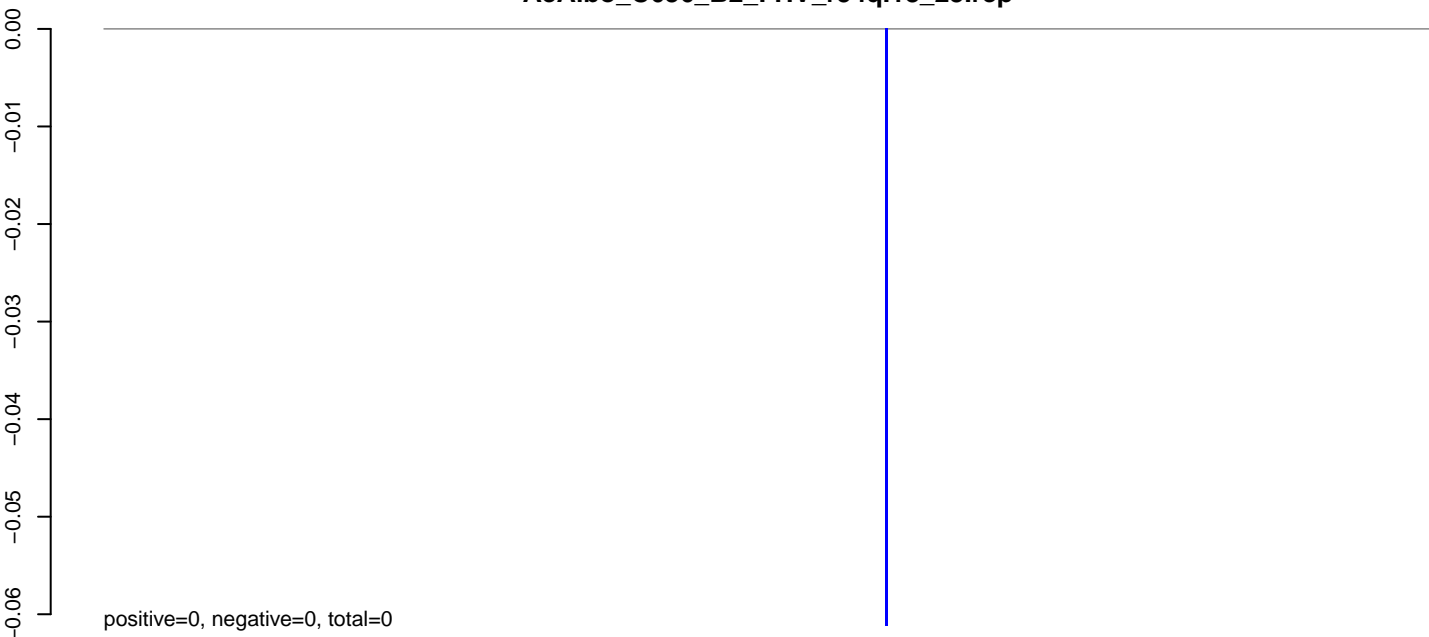
AeAlbo_C636_B2_FHV_r54q.rep



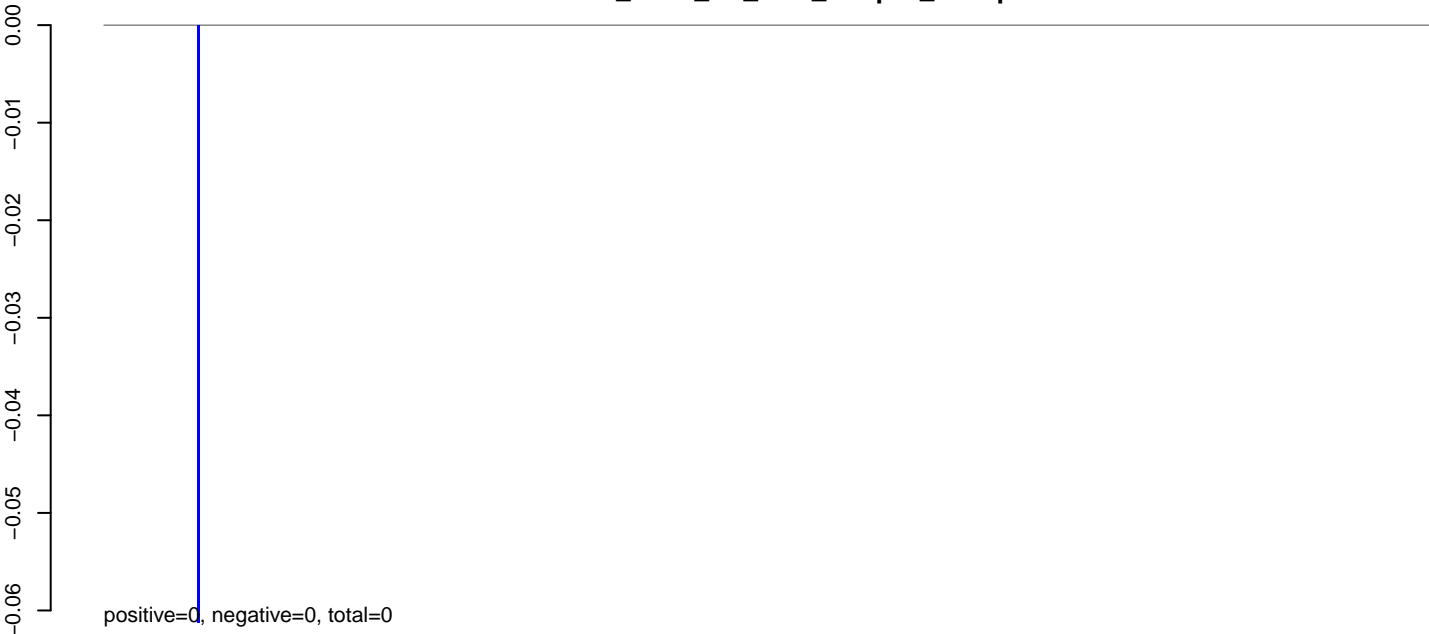
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0 2000 4000 6000 8000 10000

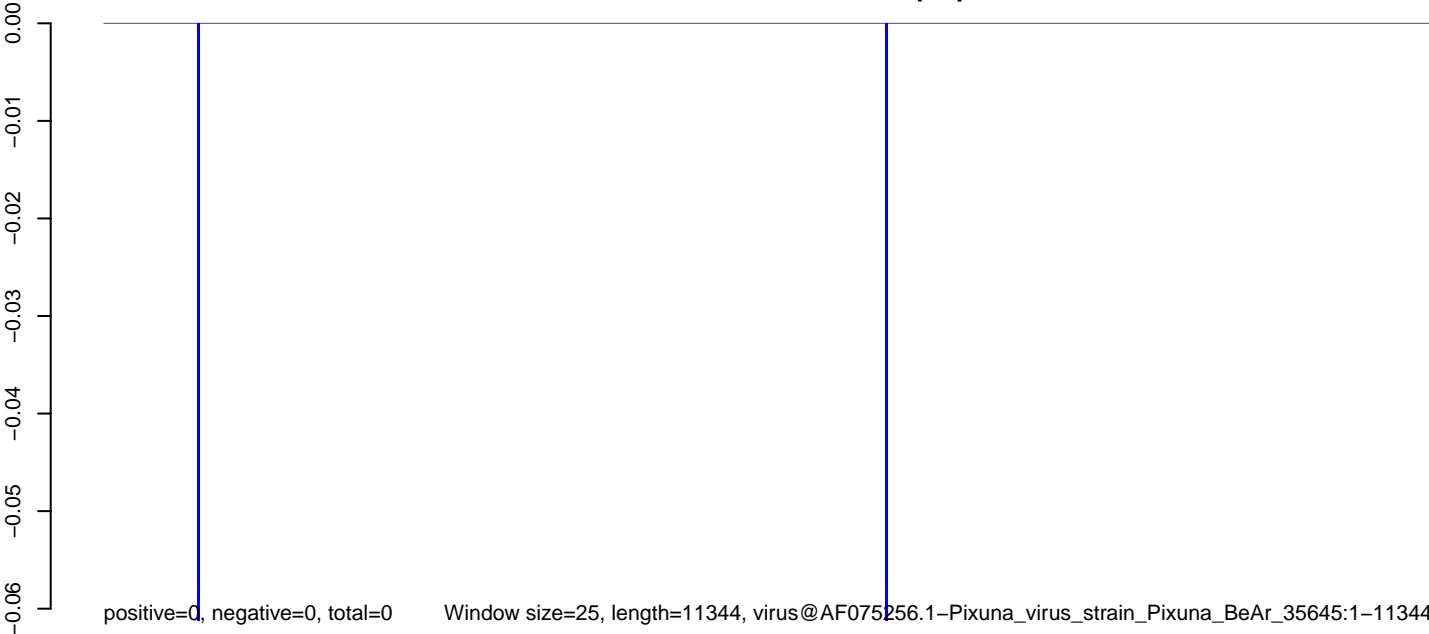
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AeAlbo_C636_B2_FHV_r54q.24_35.rep



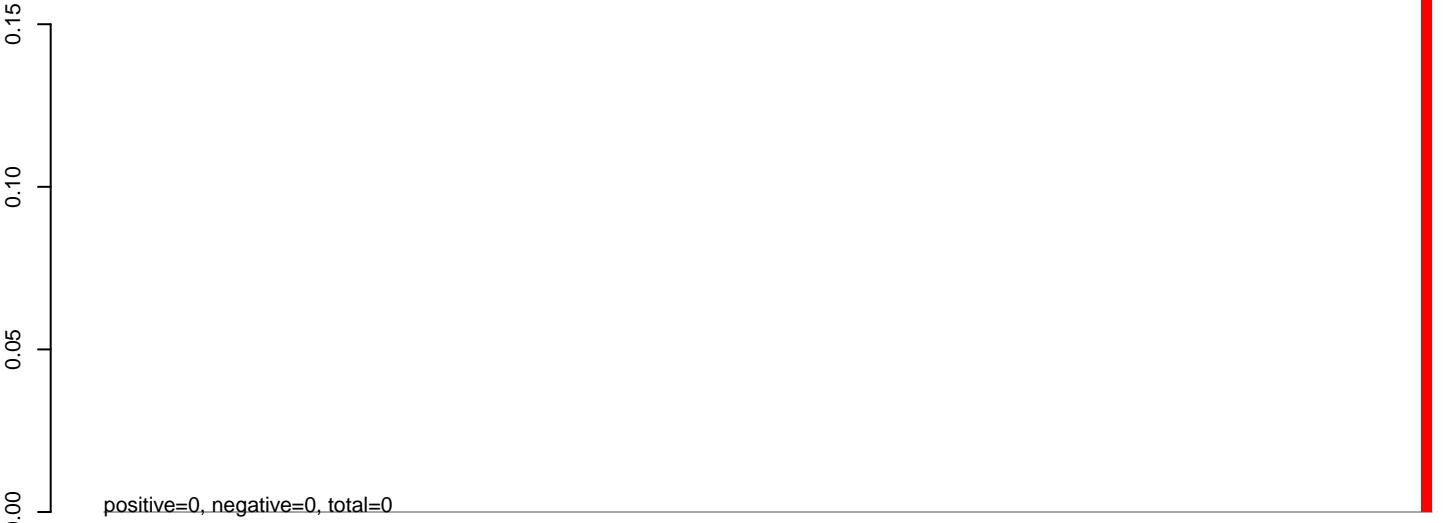
AeAlbo_C636_B2_FHV_r54q.rep



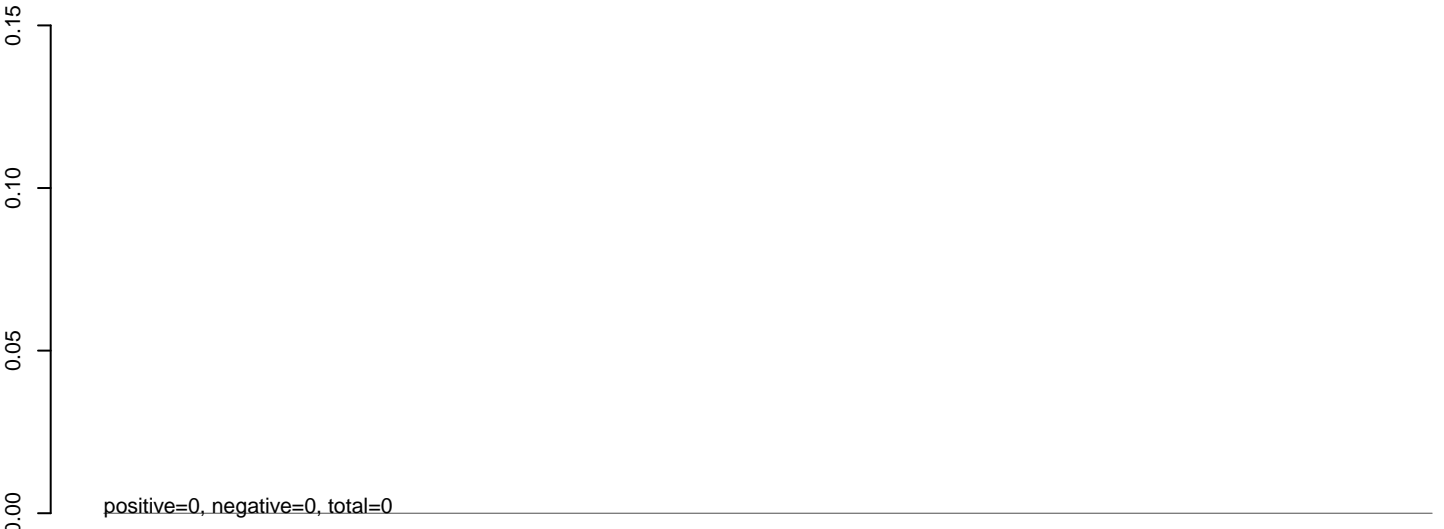
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0 2000 4000 6000 8000 10000

AeAlbo_C636_B2_FHV_r54q.18_23.rep



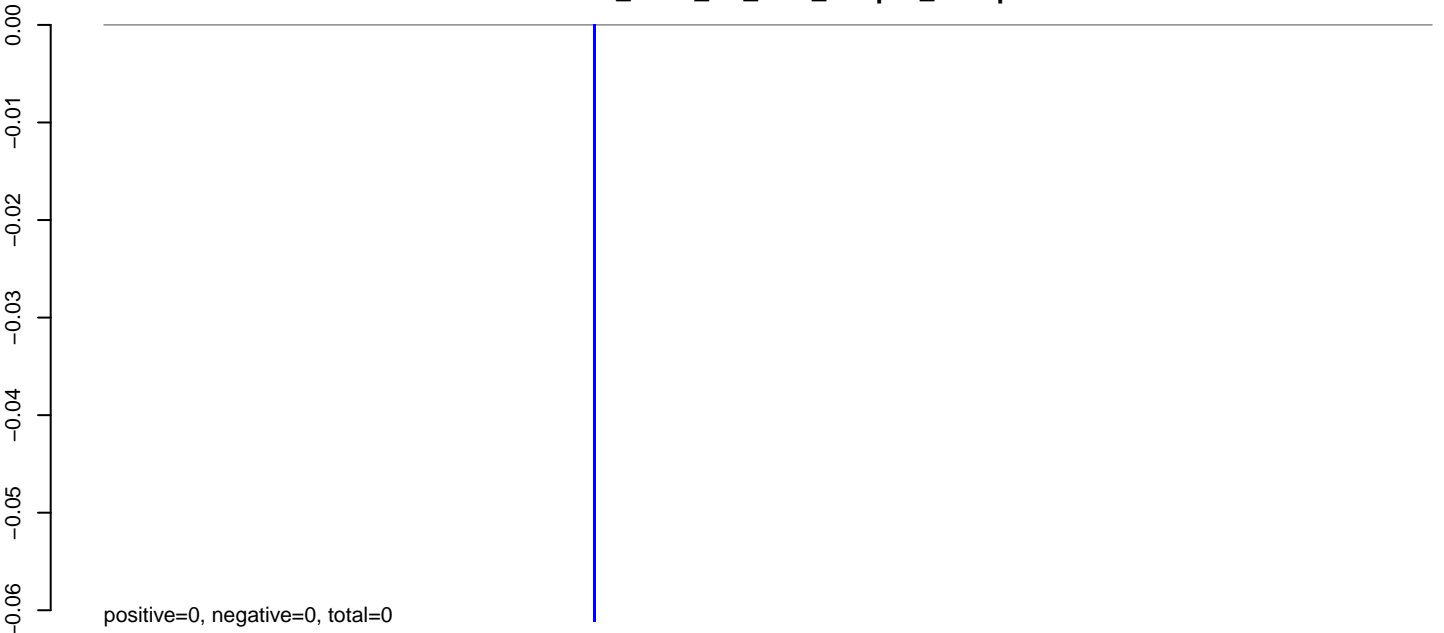
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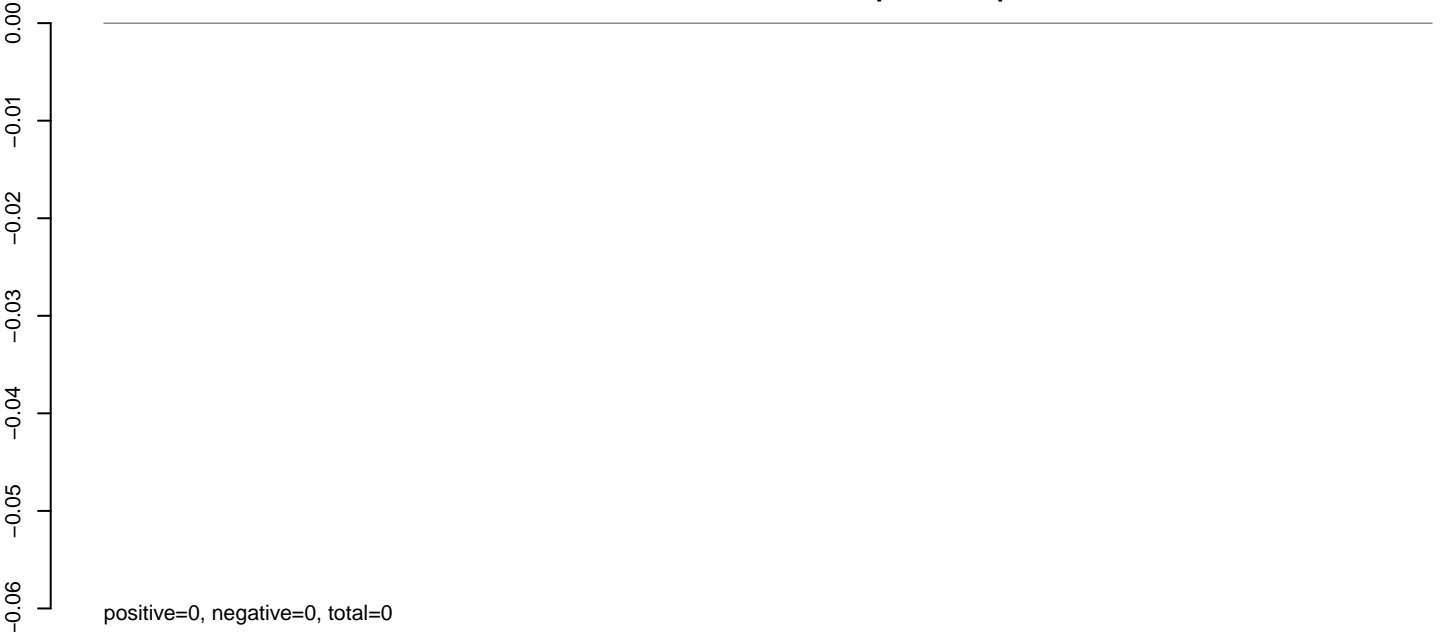
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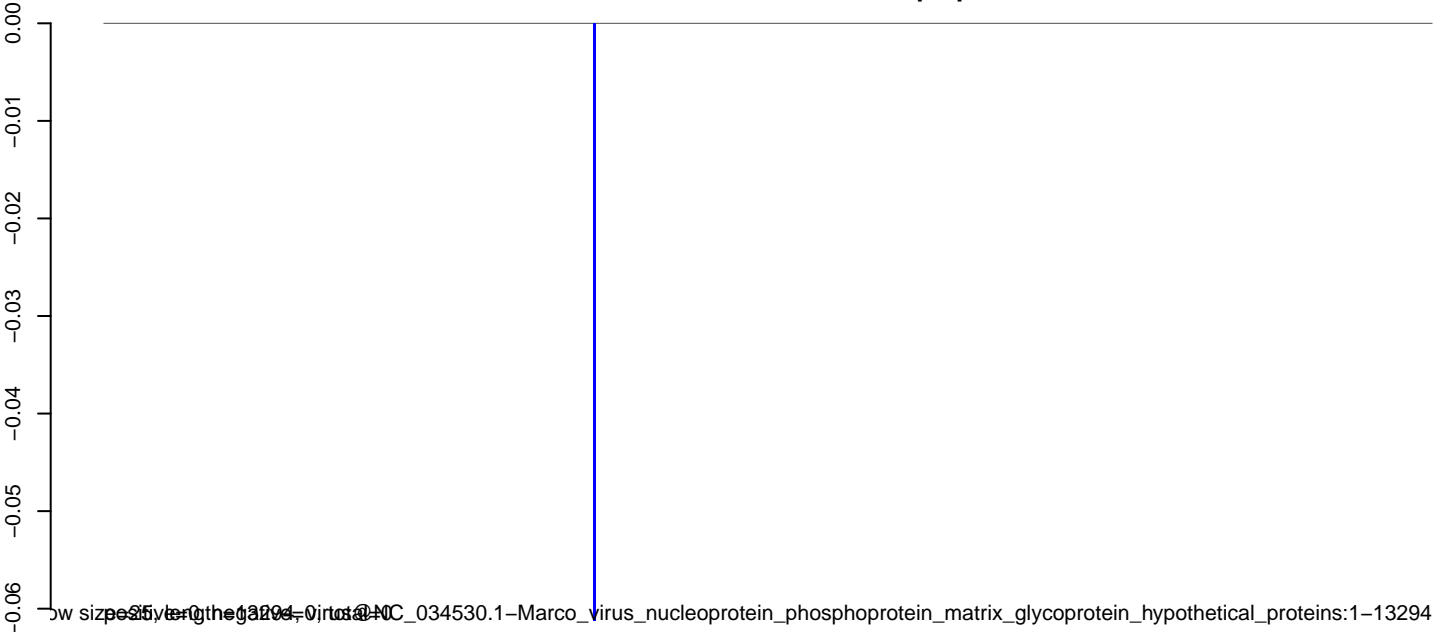
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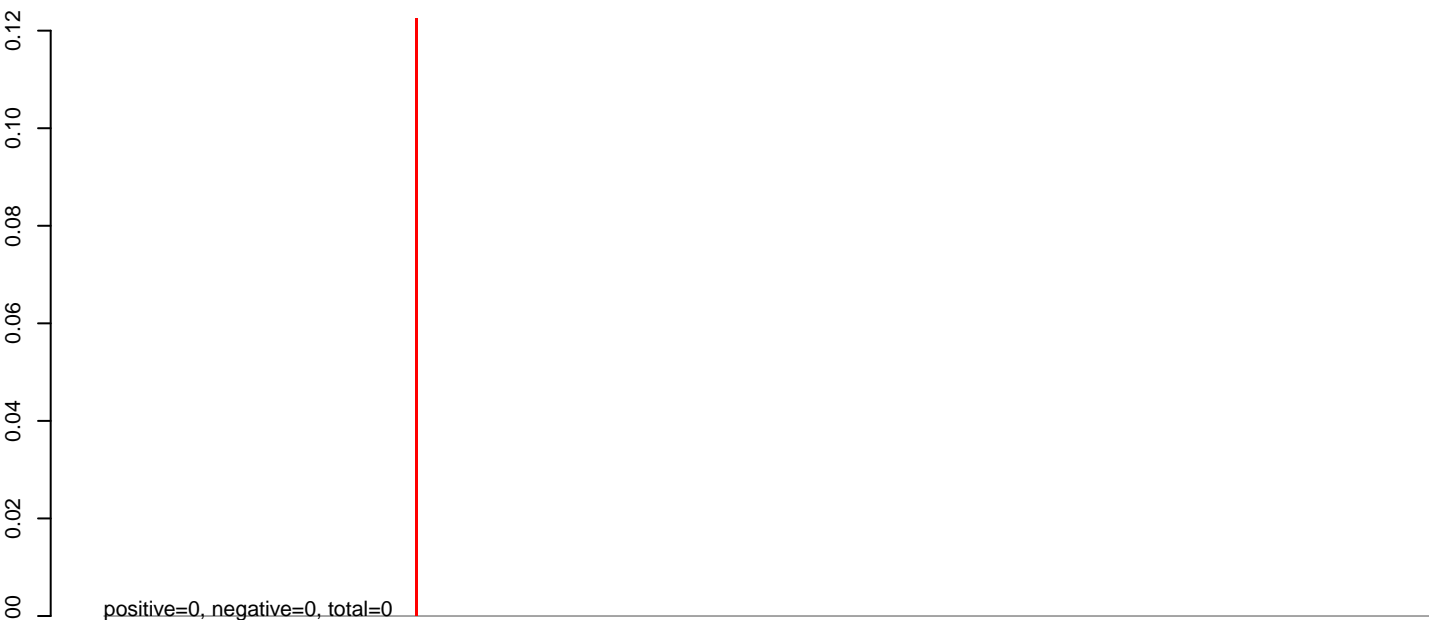
AeAlbo_C636_B2_FHV_r54q.24_35.rep



AeAlbo_C636_B2_FHV_r54q.rep



AeAlbo_C636_B2_FHV_r54q.18_23.rep



positive=0, negative=0, total=0

AeAlbo_C636_B2_FHV_r54q.24_35.rep



positive=0, negative=0, total=0

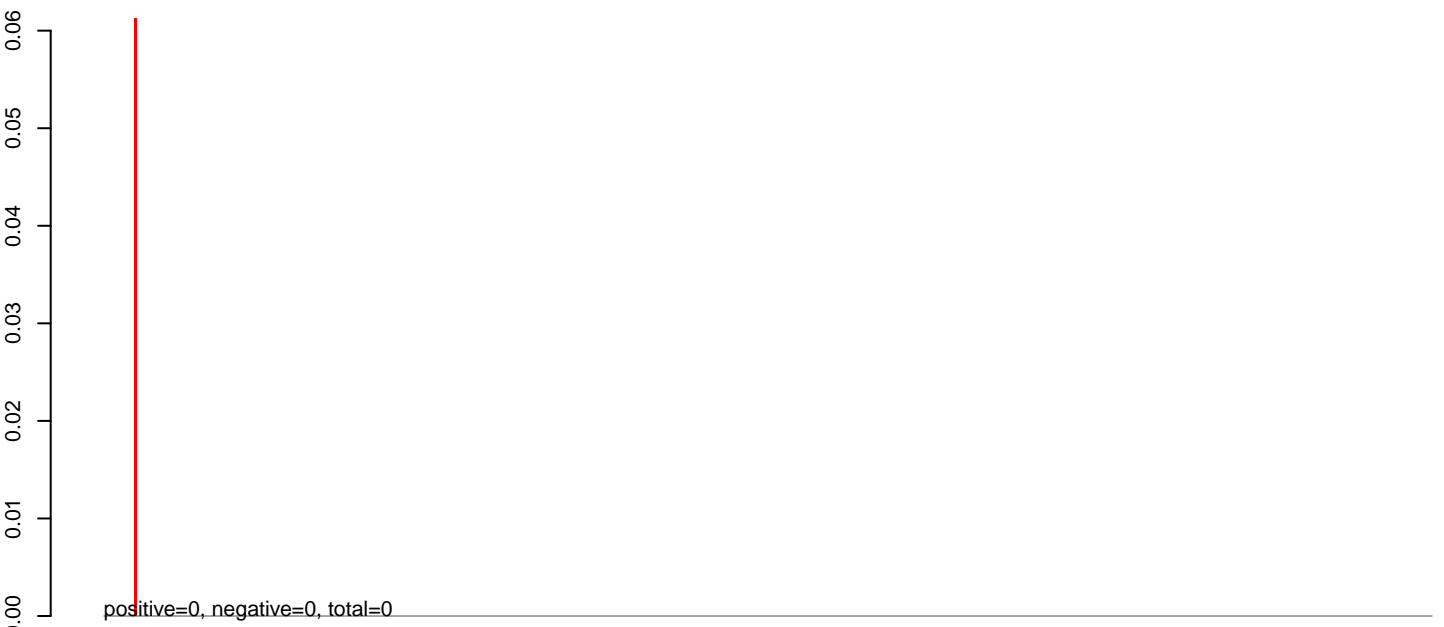
AeAlbo_C636_B2_FHV_r54q.rep



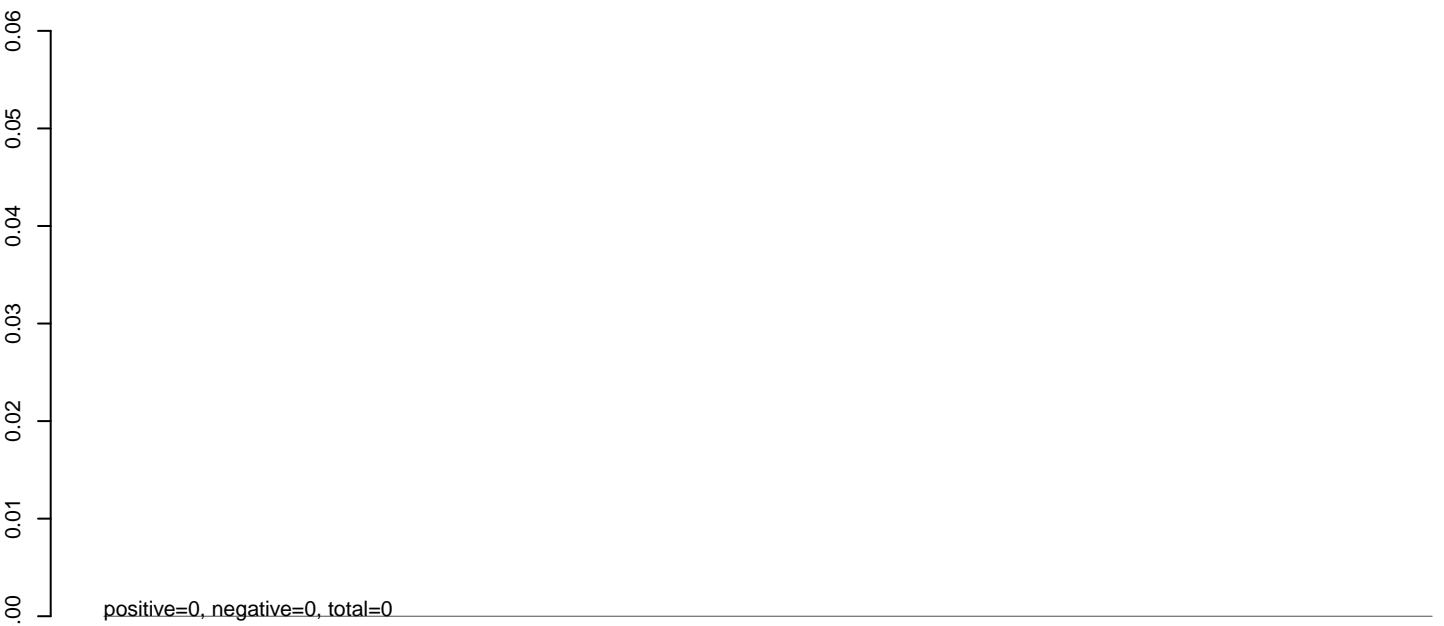
25, length=11728, negative=0, total=11728
1-Tai_virus_strain_F47_CI_2004_segment_S_plus_segment_M_plus_segment_L_nucleocapsid :1-11728

0 2000 4000 6000 8000 10000 12000

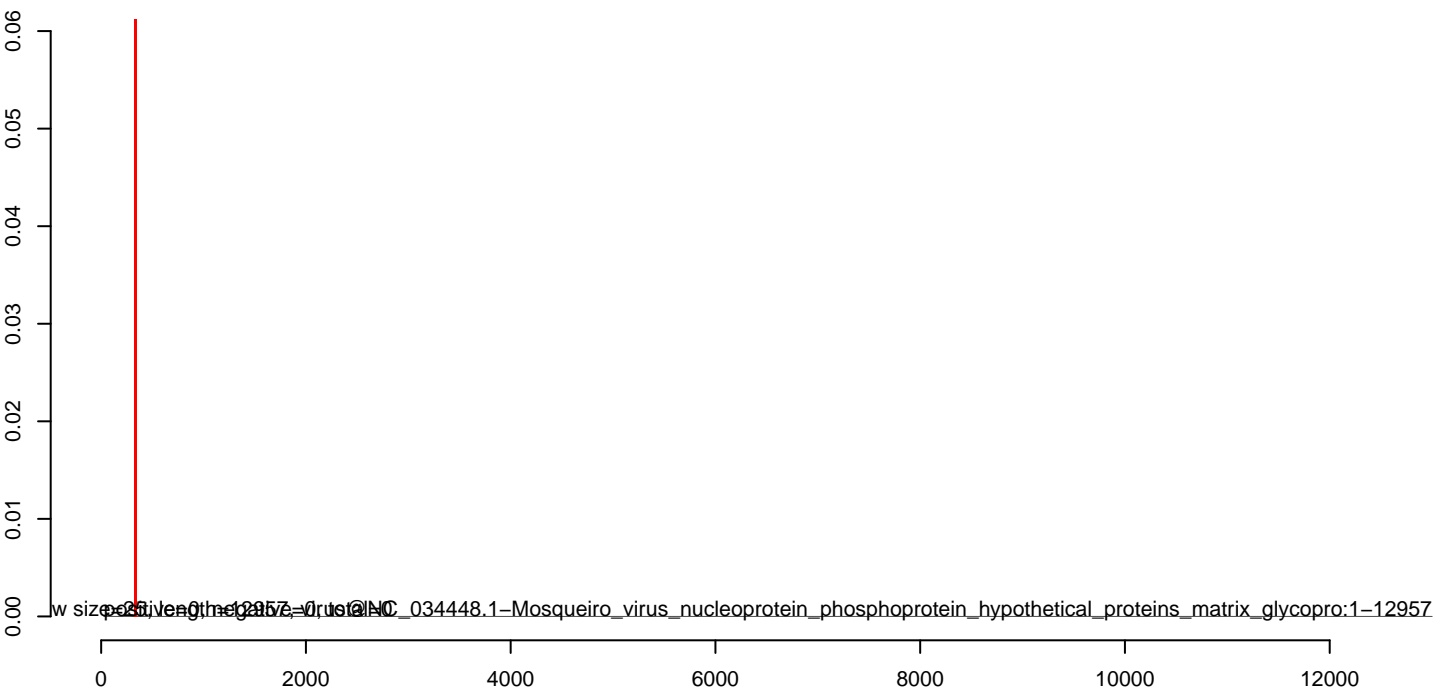
AeAlbo_C636_B2_FHV_r54q.18_23.rep



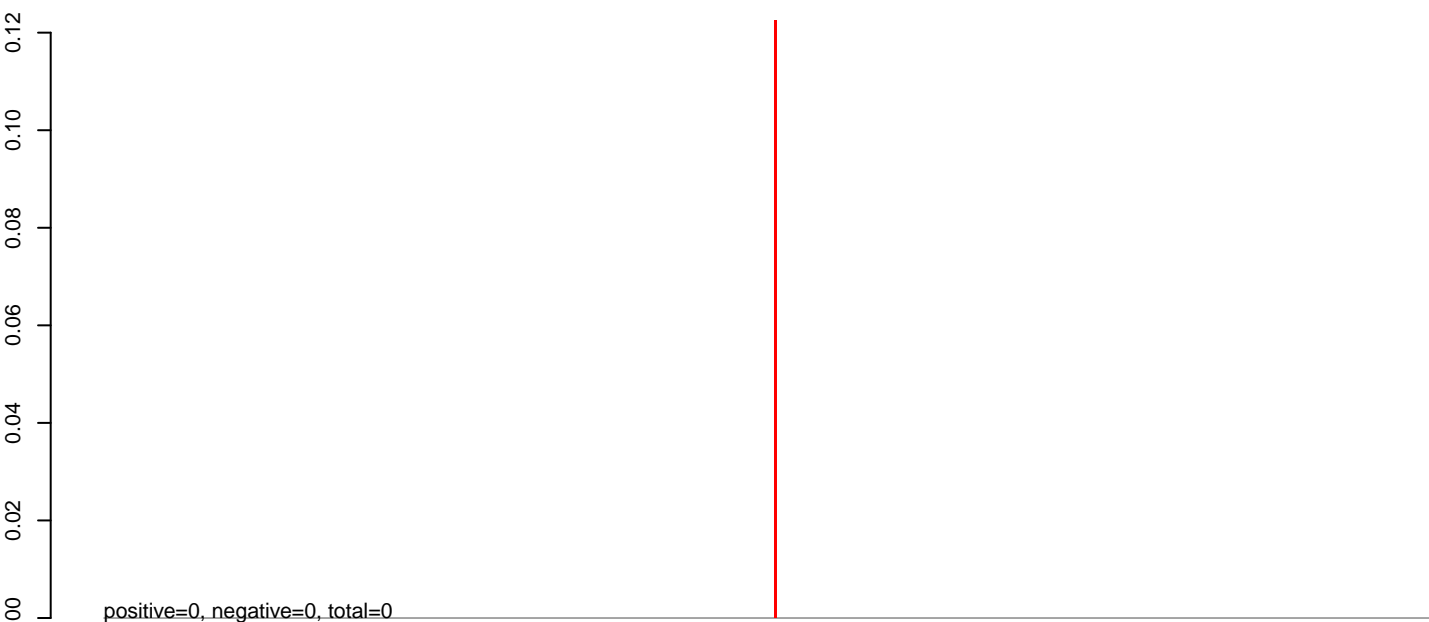
AeAlbo_C636_B2_FHV_r54q.24_35.rep



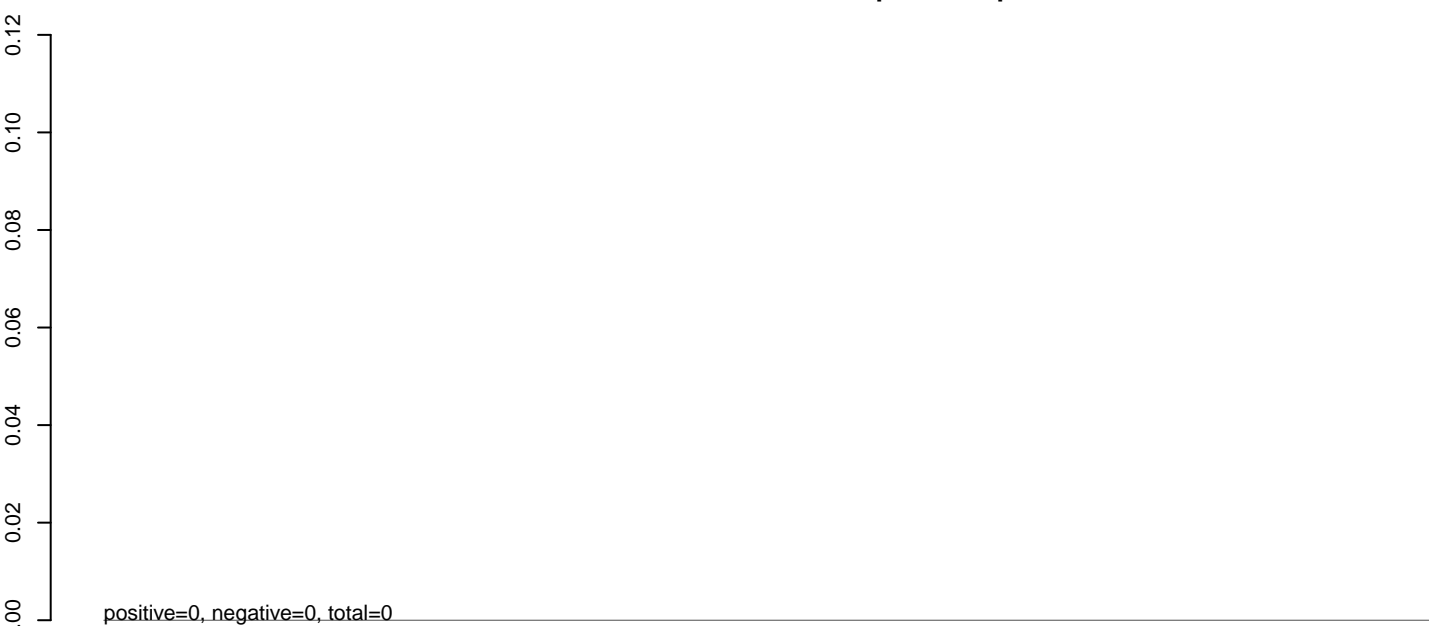
AeAlbo_C636_B2_FHV_r54q.rep



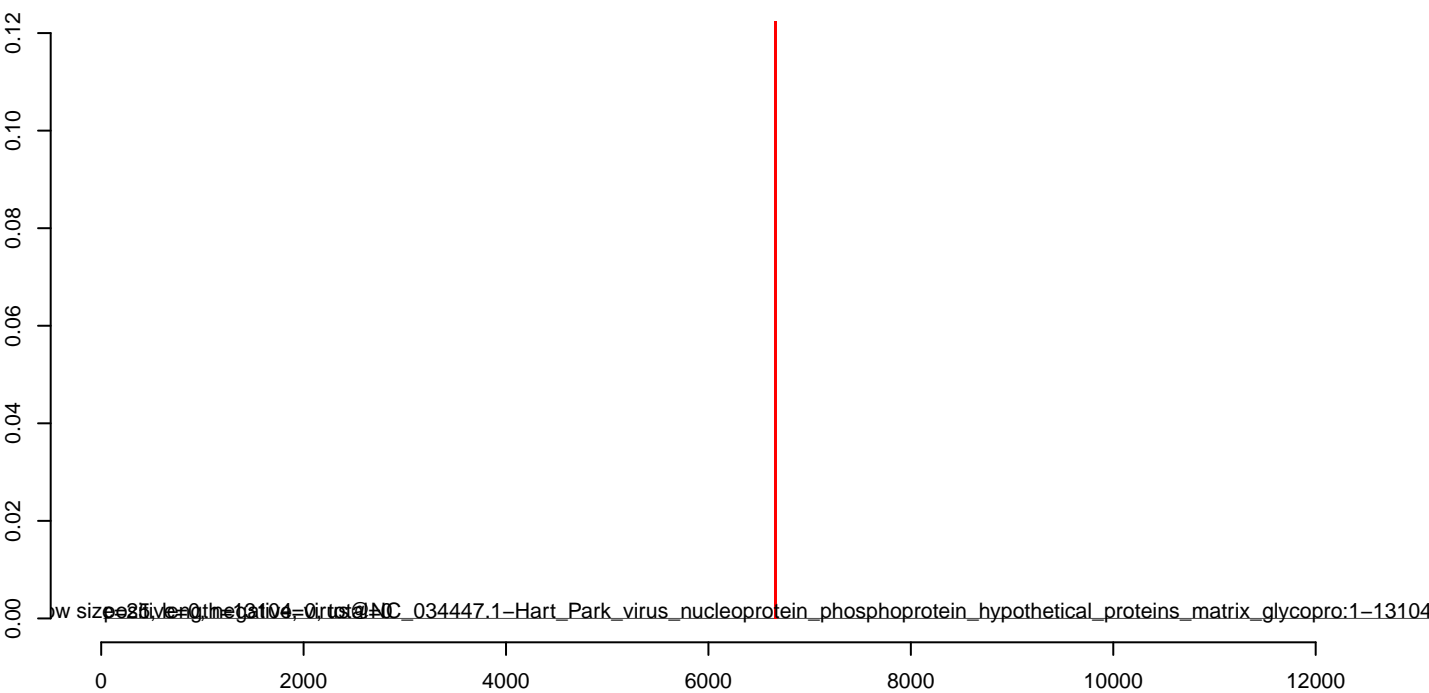
AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep

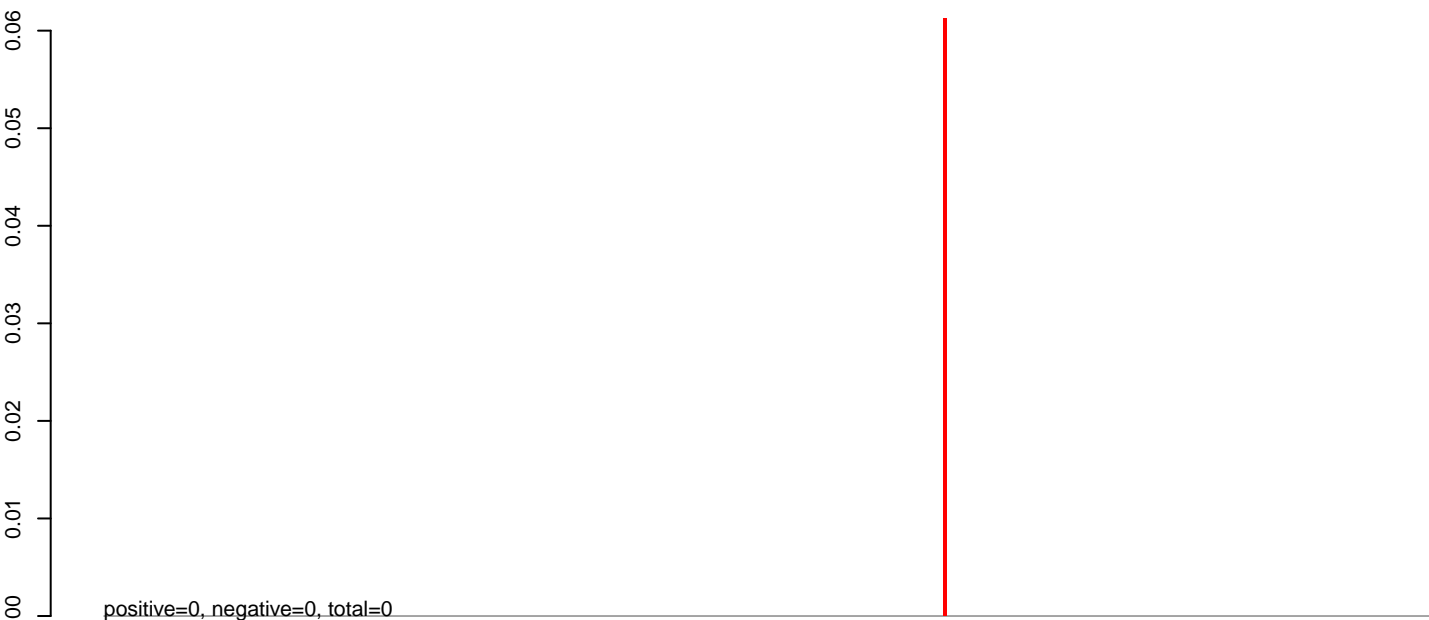


AeAlbo_C636_B2_FHV_r54q.rep

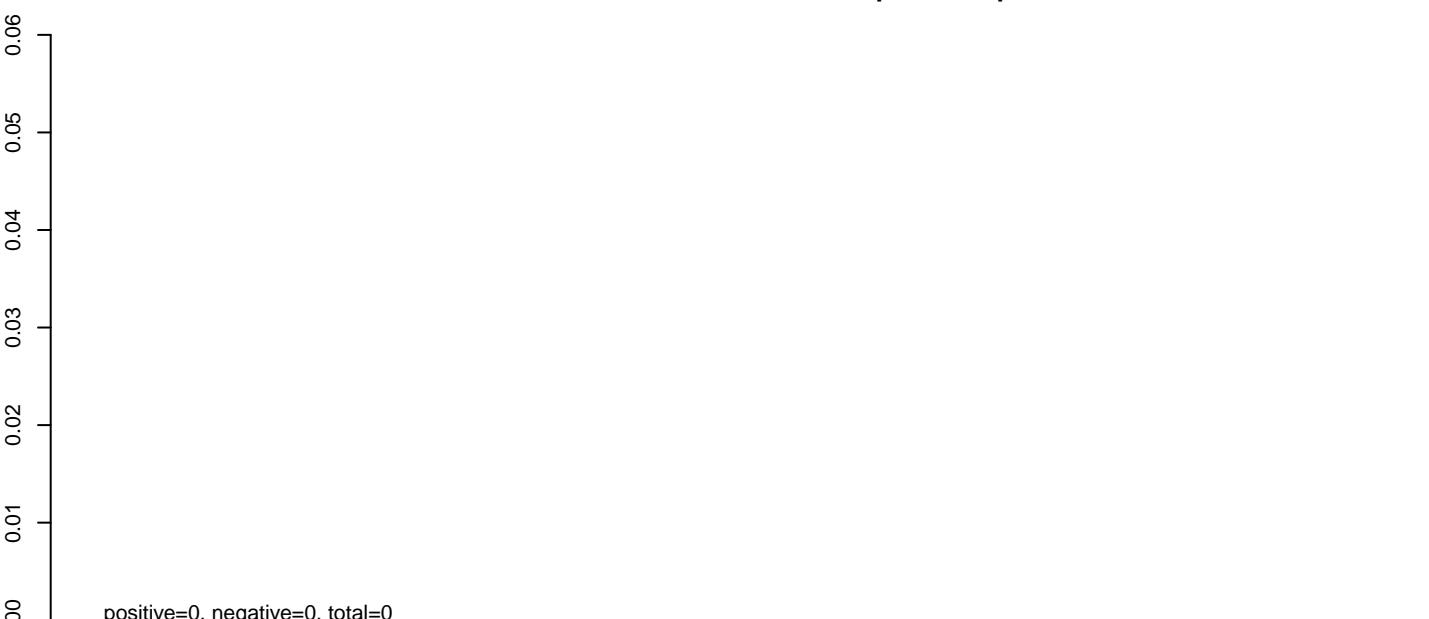


034447.1-Hart_Park_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycopro:1-13104

AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep



AeAlbo_C636_B2_FHV_r54q.rep



Window size=25, length=10869, virus@NC_031957.1-UNVERIFIED_Coot_Bay_virus_strain_EVG5-53:1-10869

0 2000 4000 6000 8000 10000

AeAlbo_C636_B2_FHV_r54q.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10
-0.12

positive=0, negative=0, total=0

AeAlbo_C636_B2_FHV_r54q.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10
-0.12

positive=0, negative=0, total=0

AeAlbo_C636_B2_FHV_r54q.rep

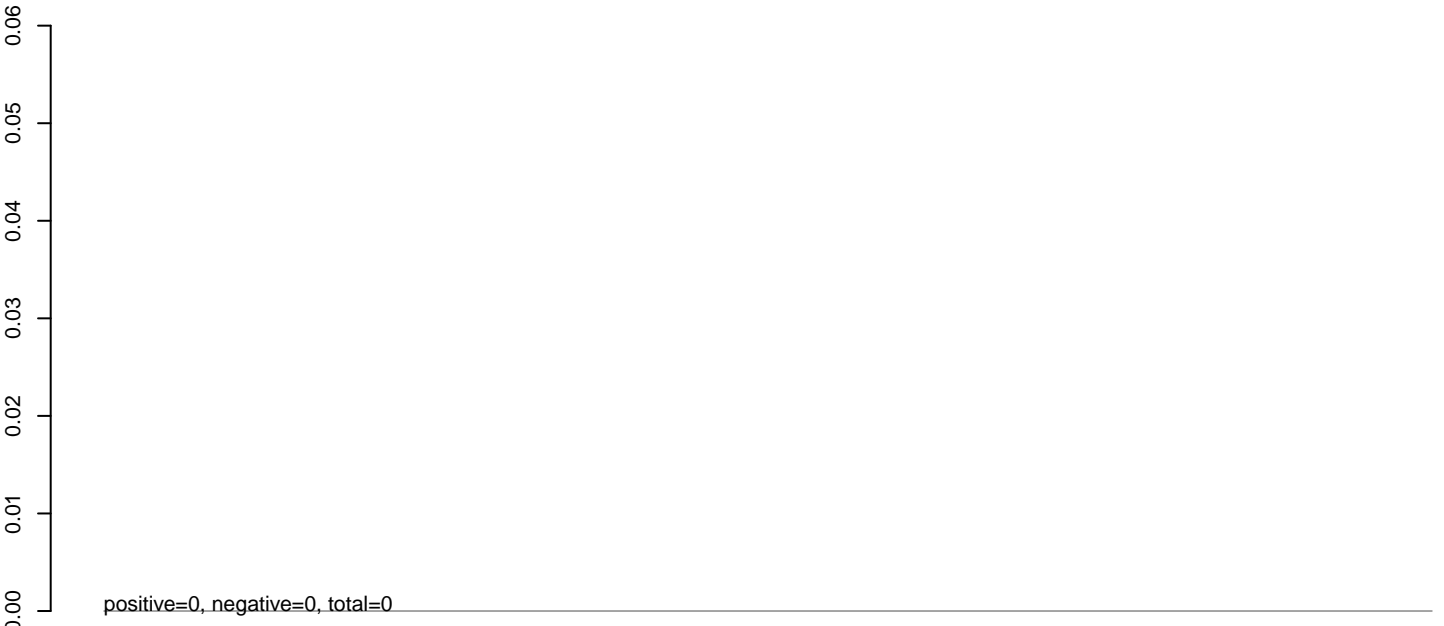
0.00
-0.02
-0.04
-0.06
-0.08
-0.10
-0.12

positive=0, negative=0, total=0

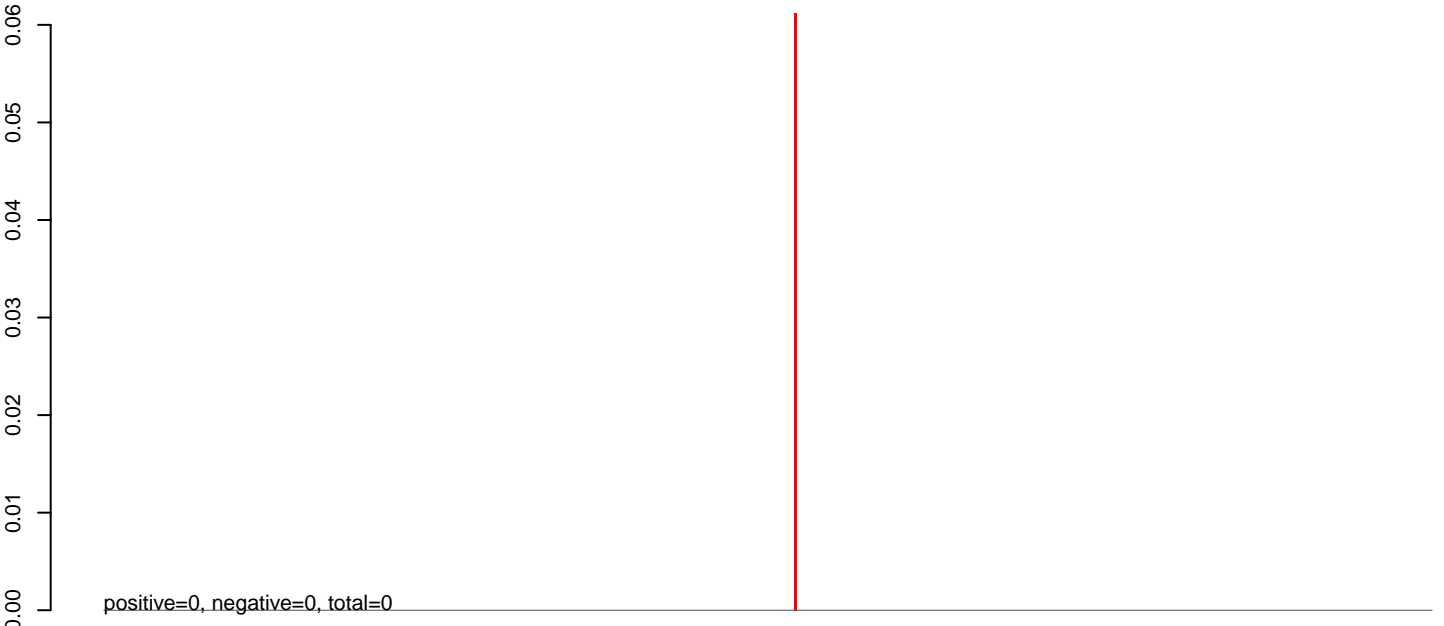
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0 2000 4000 6000 8000 10000

AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep



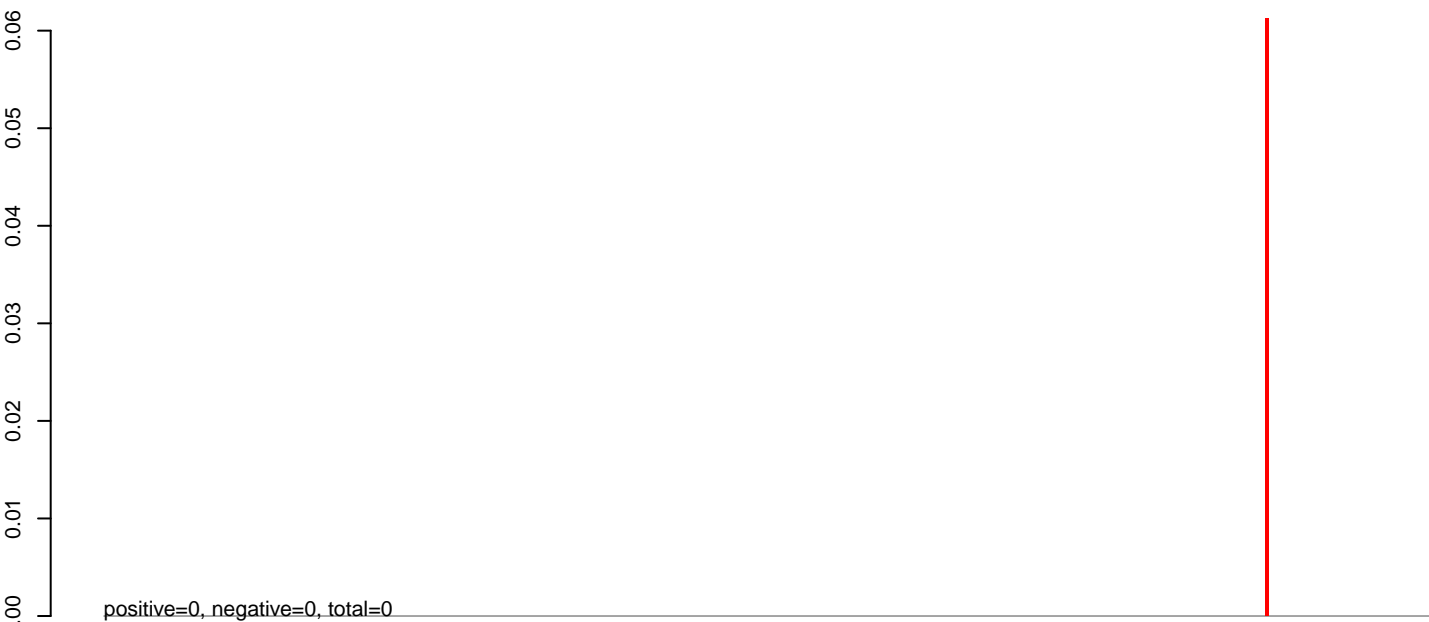
AeAlbo_C636_B2_FHV_r54q.rep



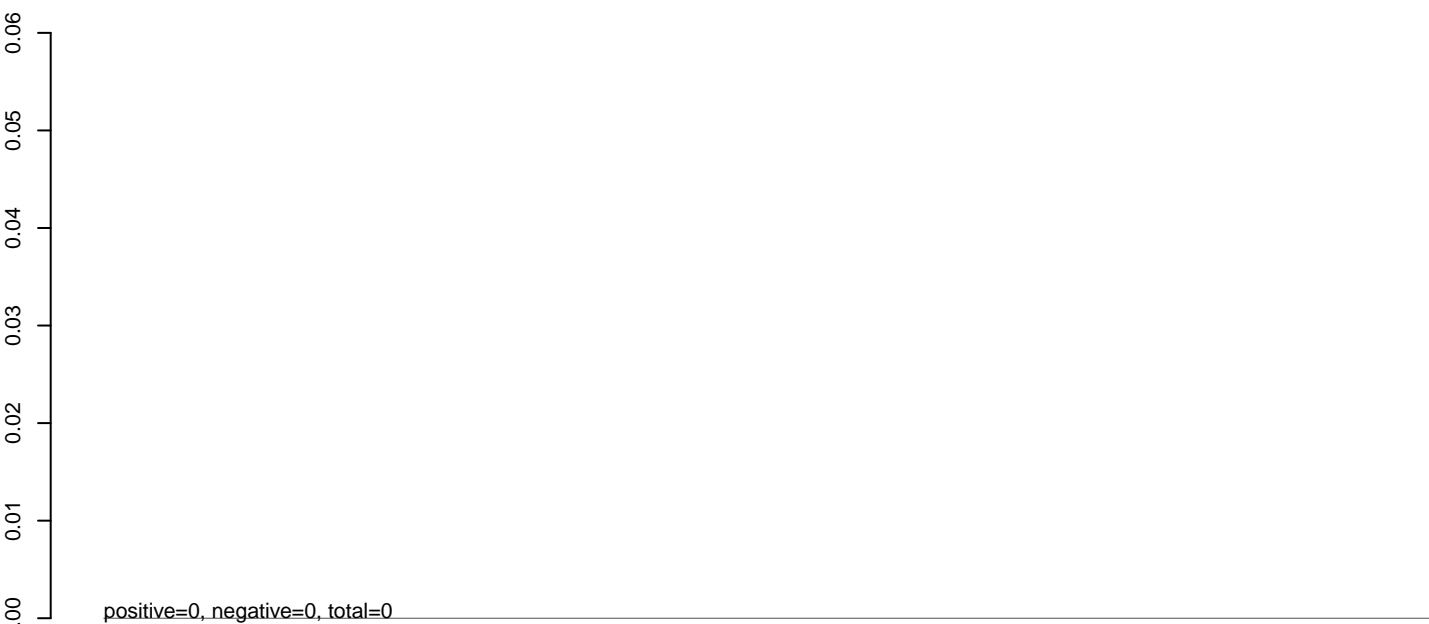
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0 2000 4000 6000 8000 10000 12000

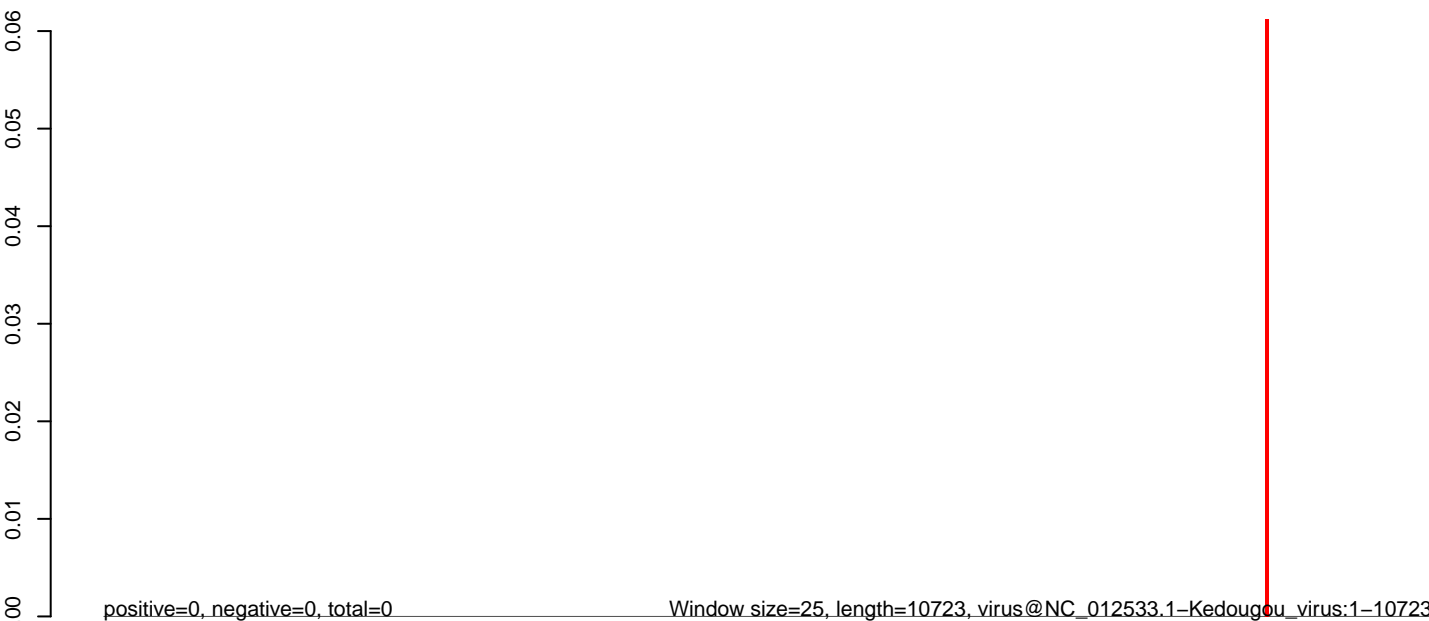
AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep

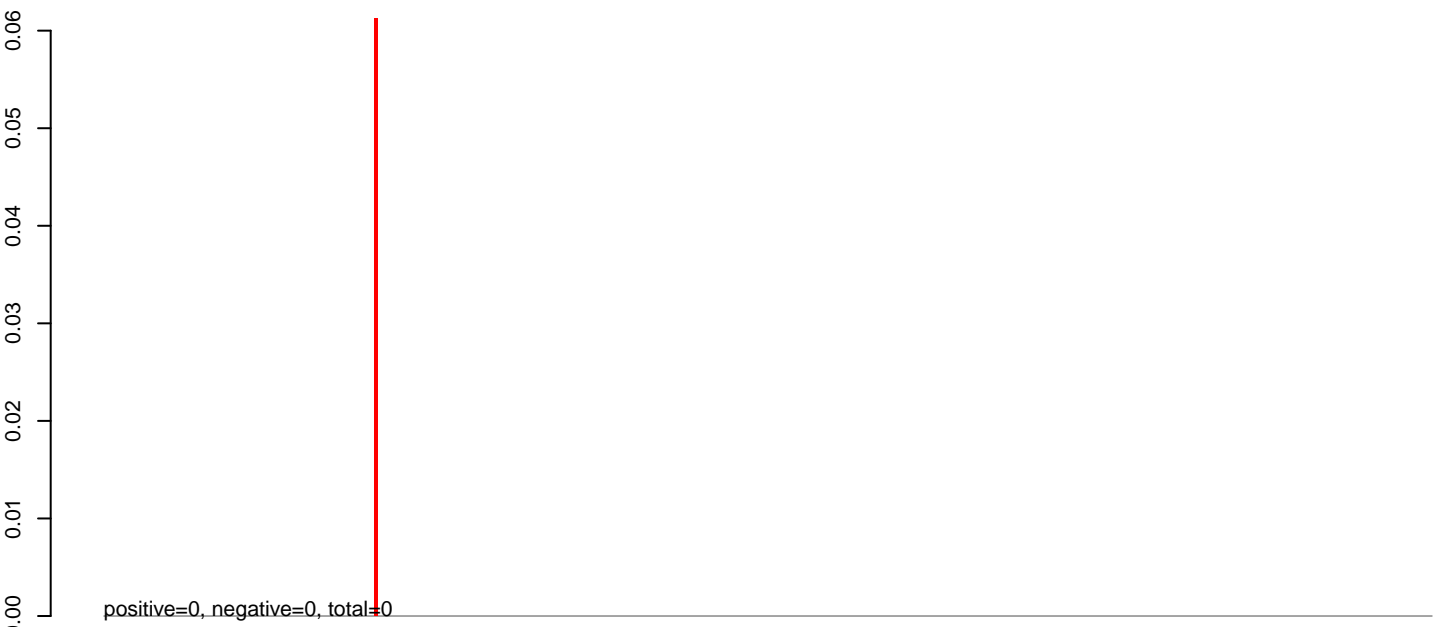


AeAlbo_C636_B2_FHV_r54q.rep

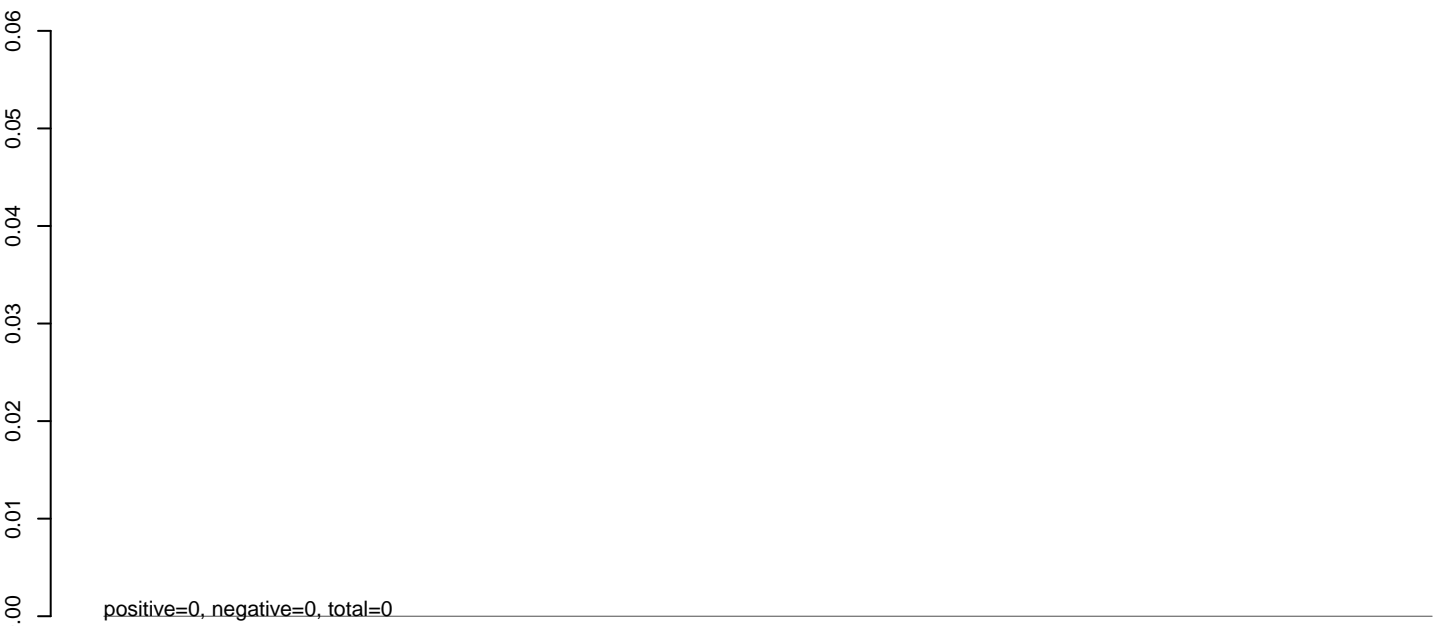


0 2000 4000 6000 8000 10000

AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep



AeAlbo_C636_B2_FHV_r54q.rep



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

0 2000 4000 6000 8000 10000

AeAlbo_C636_B2_FHV_r54q.18_23.rep

0.00
-0.01
-0.02
-0.03
-0.04
-0.05
-0.06

positive=0, negative=0, total=0

AeAlbo_C636_B2_FHV_r54q.24_35.rep

0.00
-0.01
-0.02
-0.03
-0.04
-0.05
-0.06

positive=0, negative=0, total=0

AeAlbo_C636_B2_FHV_r54q.rep

0.00
-0.01
-0.02
-0.03
-0.04
-0.05
-0.06

positive=0, negative=0, total=0

Window size=25, length=11375, virus@NC_005064.1-Kamiti_River_virus:1-11375

0 2000 4000 6000 8000 10000

AeAlbo_C636_B2_FHV_r54q.18_23.rep

0.00
-0.01
-0.02
-0.03
-0.04
-0.05
-0.06

positive=0, negative=0, total=0

AeAlbo_C636_B2_FHV_r54q.24_35.rep

0.00
-0.01
-0.02
-0.03
-0.04
-0.05
-0.06

positive=0, negative=0, total=0

AeAlbo_C636_B2_FHV_r54q.rep

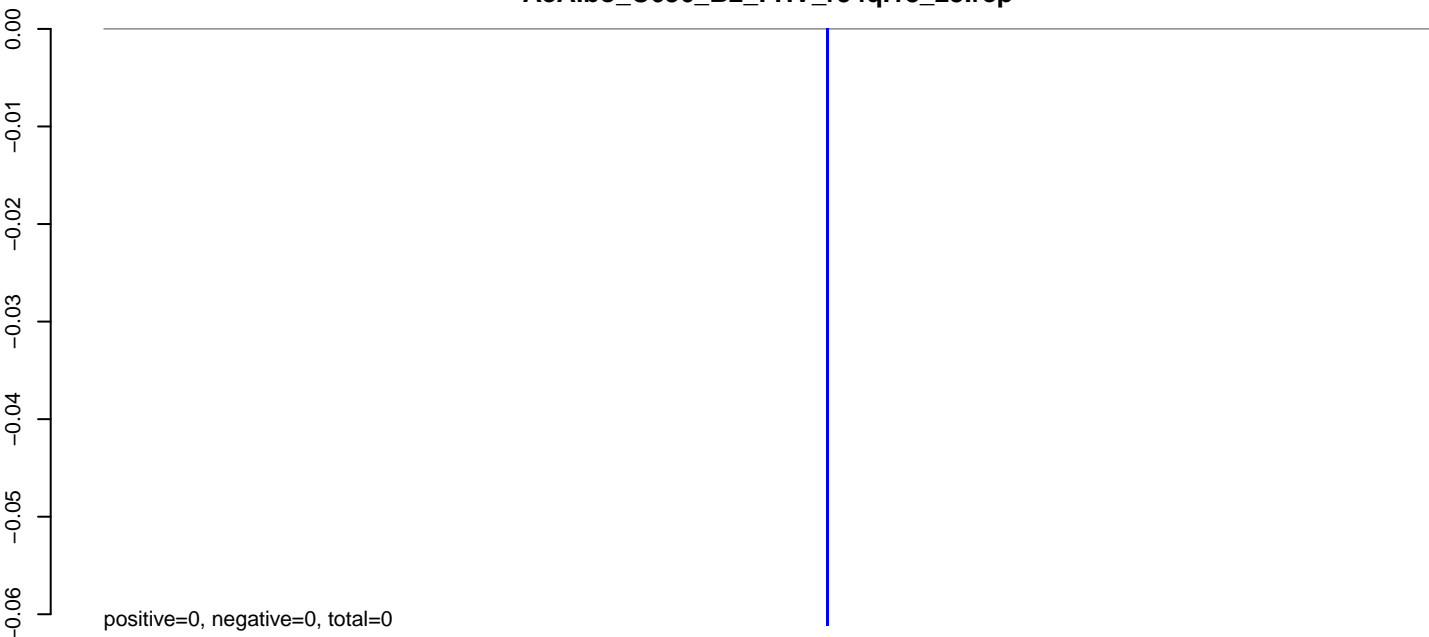
0.00
-0.01
-0.02
-0.03
-0.04
-0.05
-0.06

positive=0, negative=0, total=0

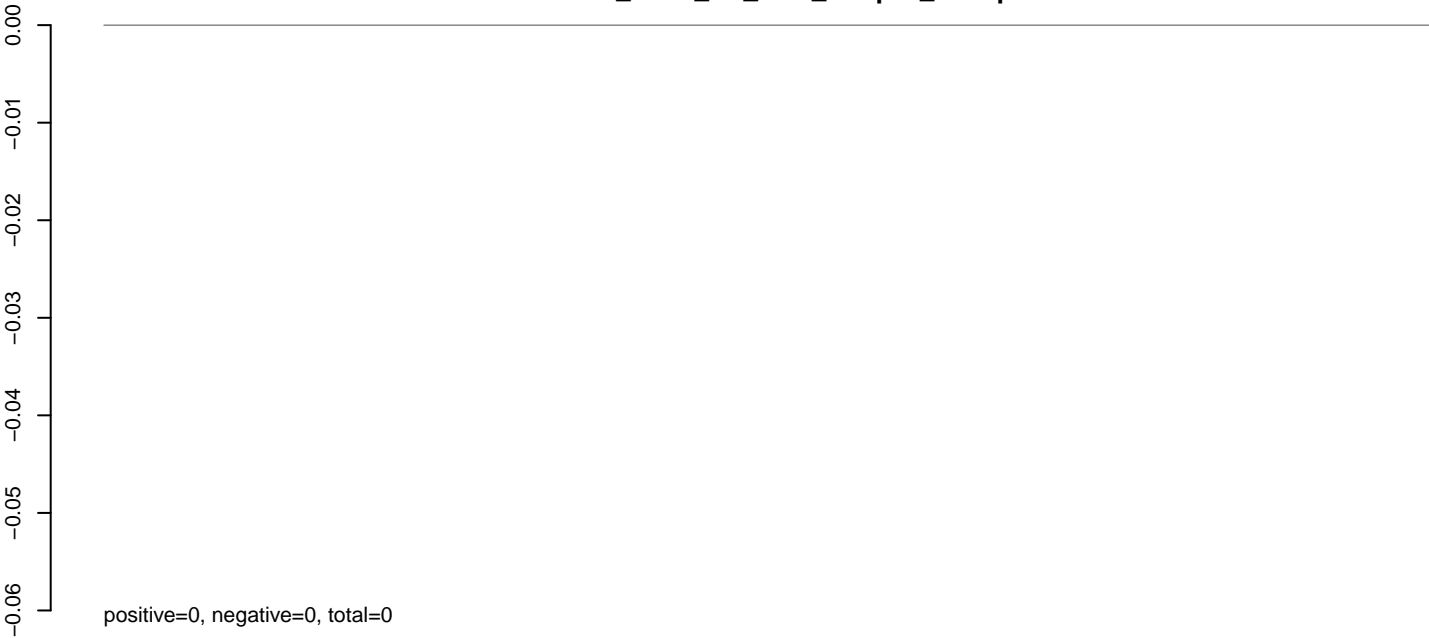
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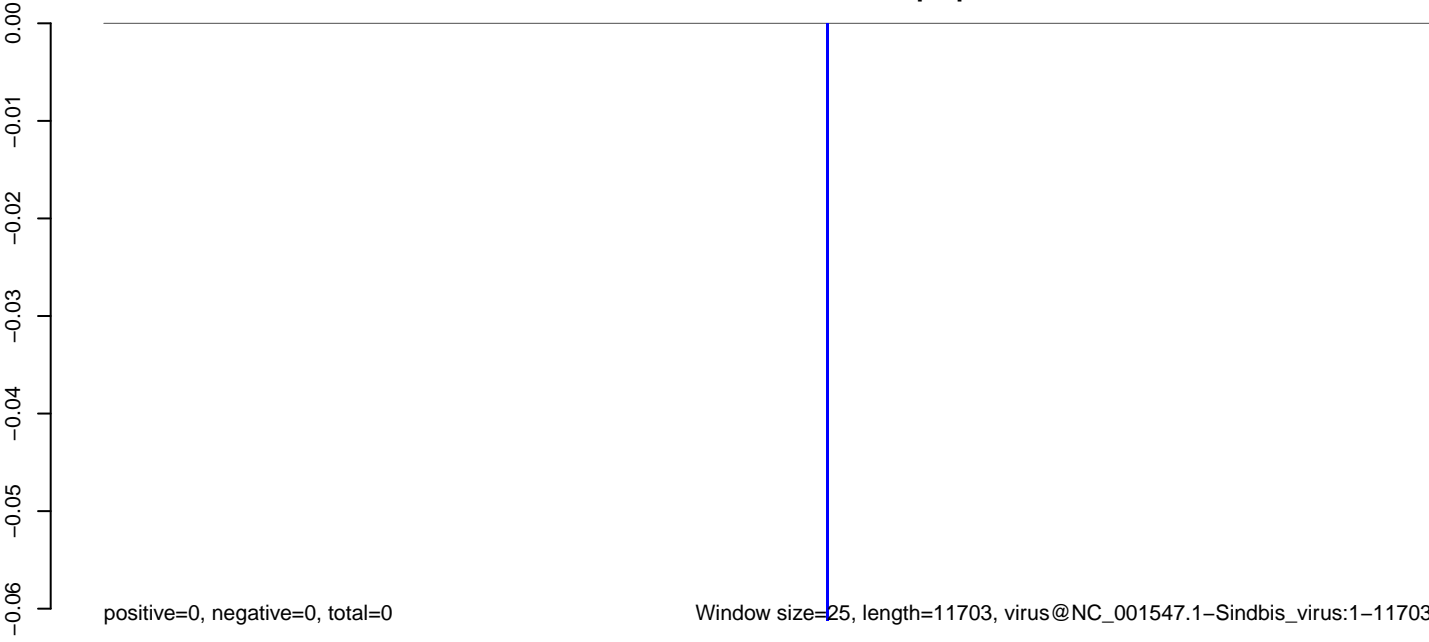
AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep



AeAlbo_C636_B2_FHV_r54q.rep



0 2000 4000 6000 8000 10000 12000

AeAlbo_C636_B2_FHV_r54q.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10
-0.12

positive=0, negative=0, total=0

AeAlbo_C636_B2_FHV_r54q.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10
-0.12

positive=0, negative=0, total=0

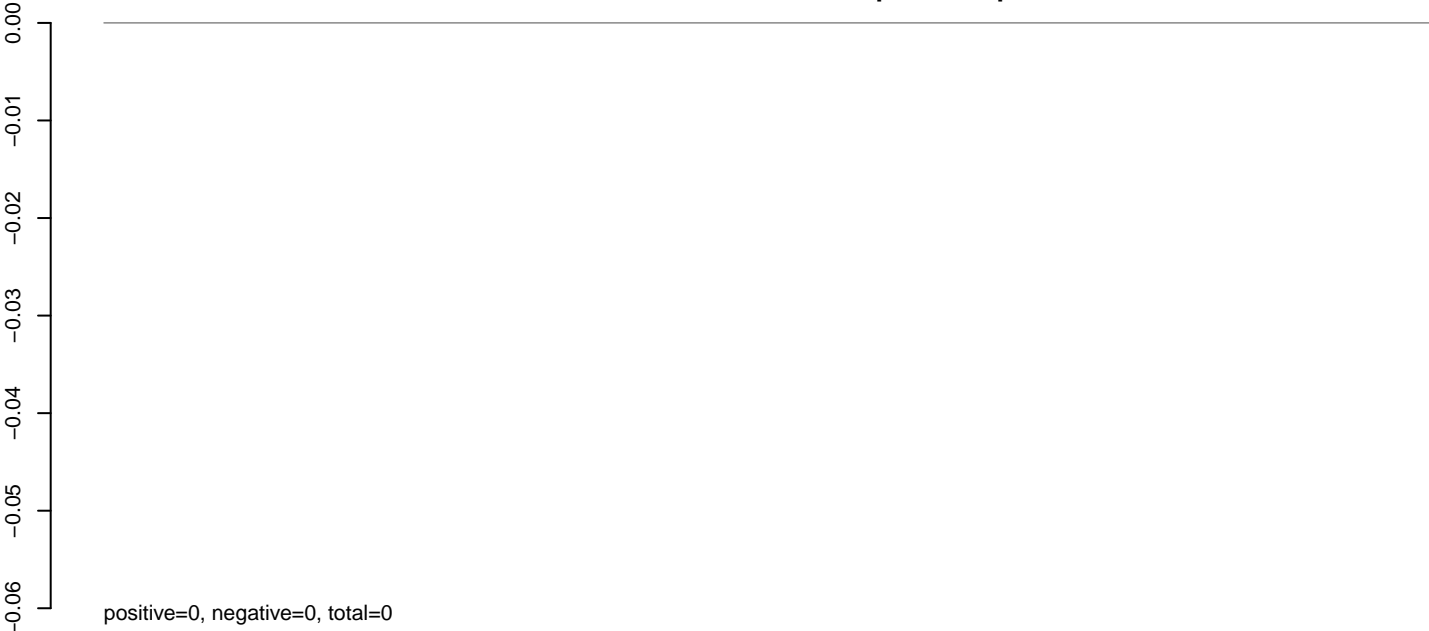
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0.00
-0.02
-0.04
-0.06
-0.08
-0.10
-0.12

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

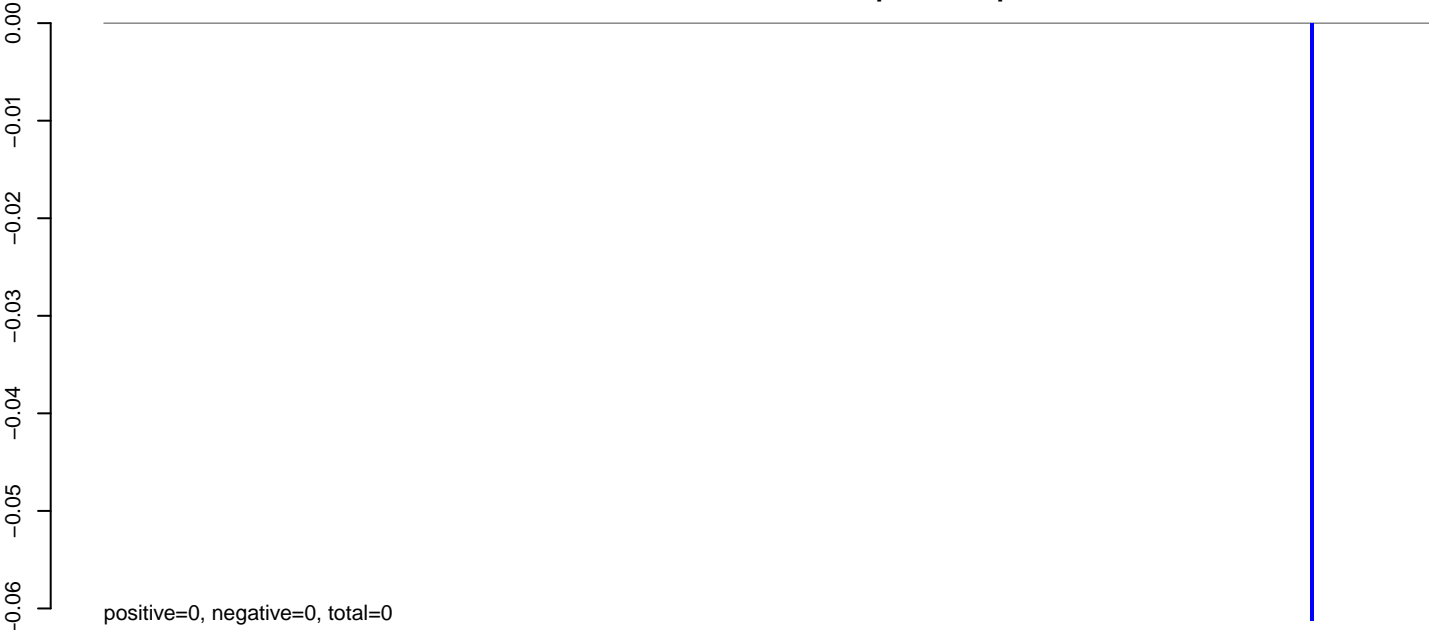
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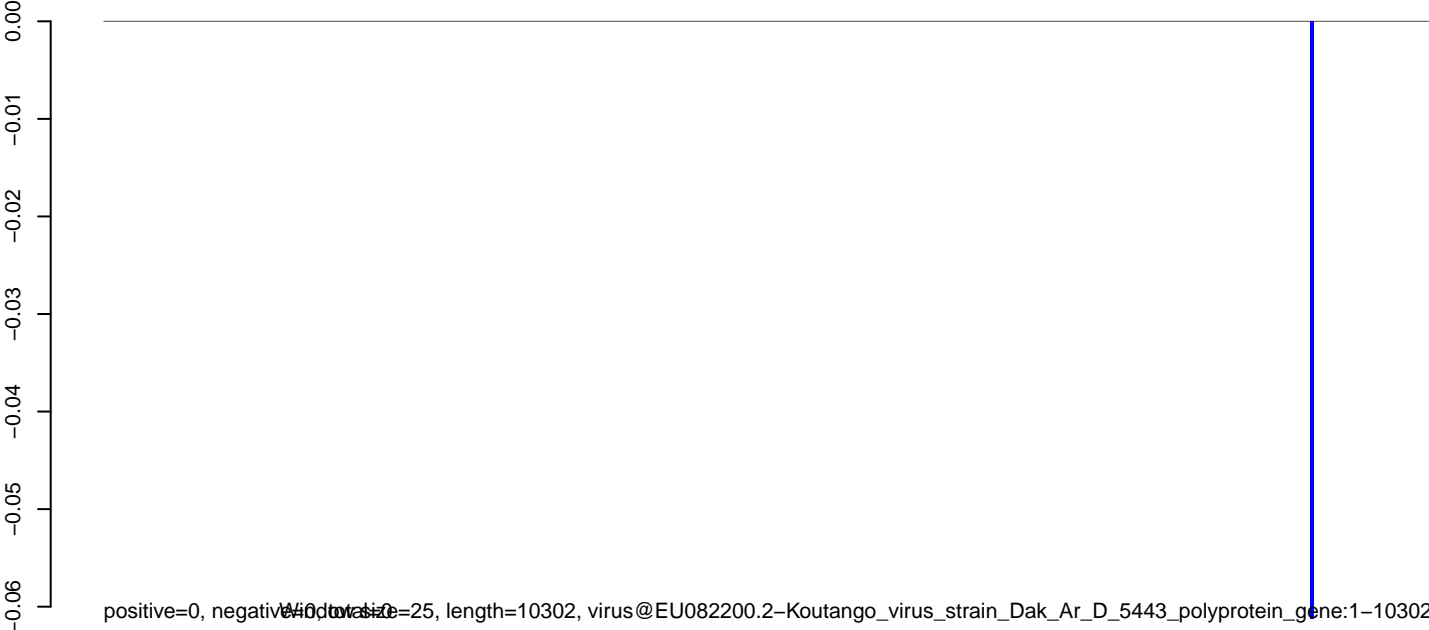
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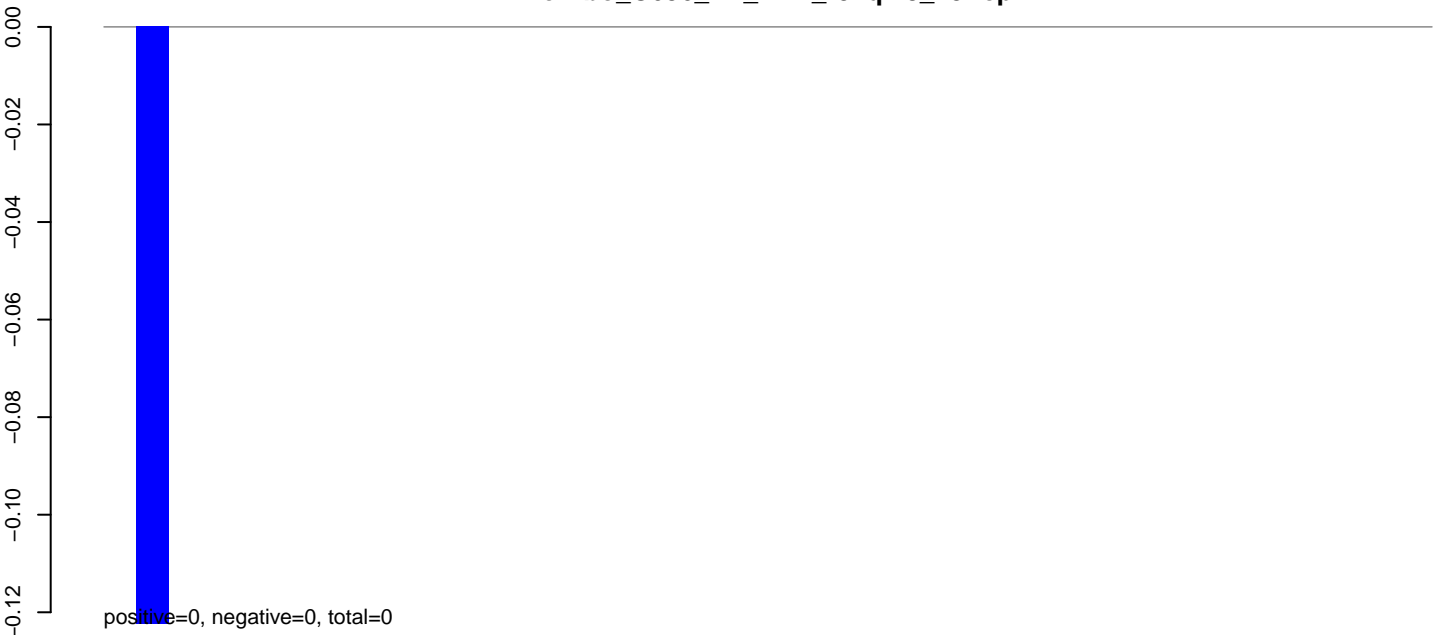
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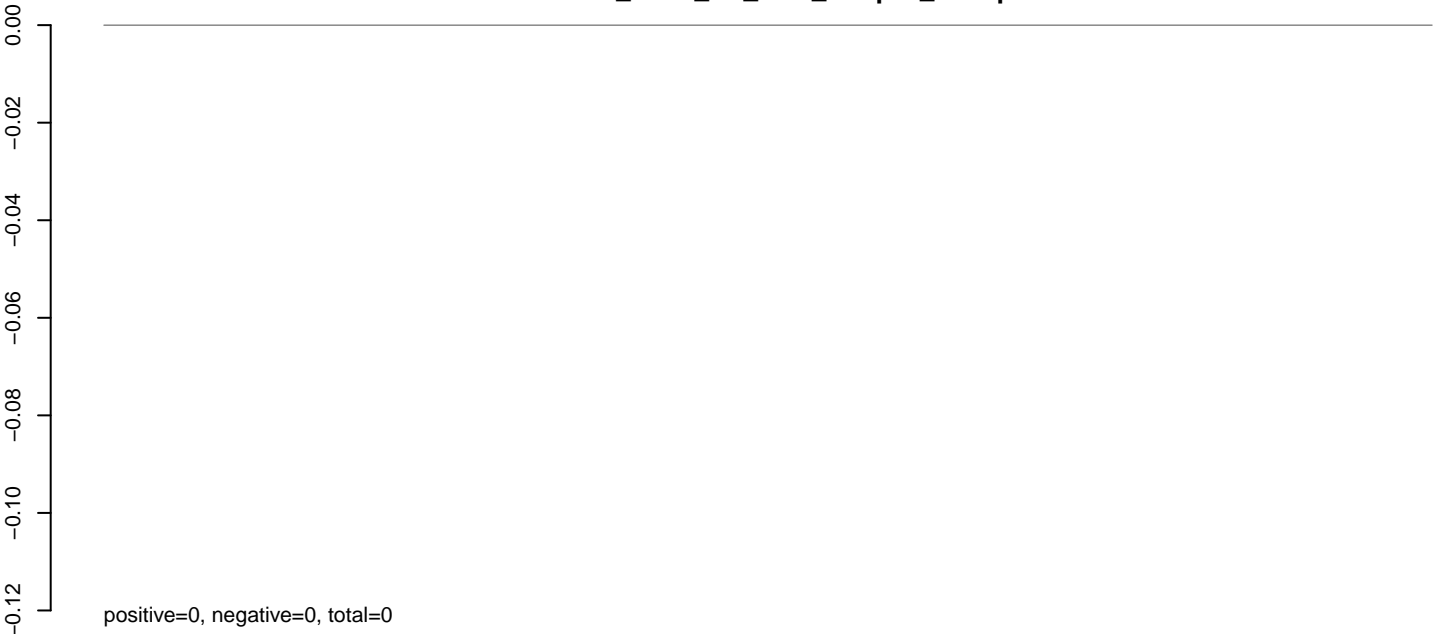


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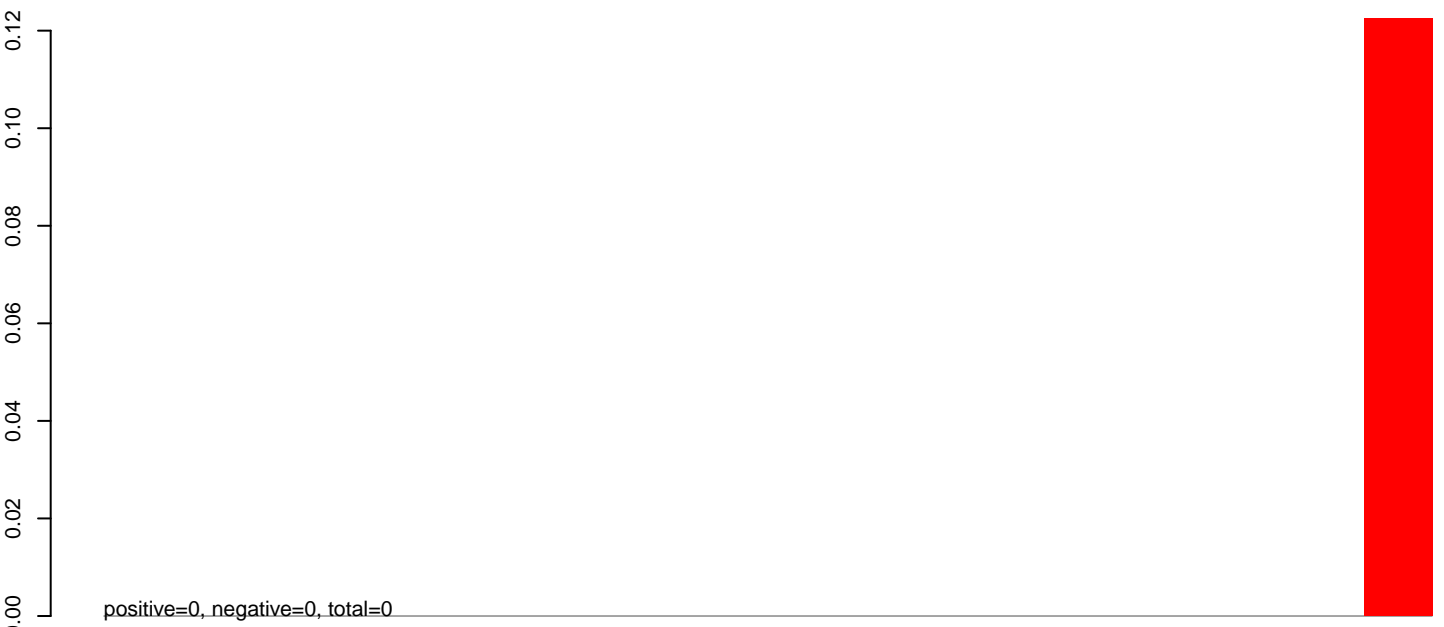
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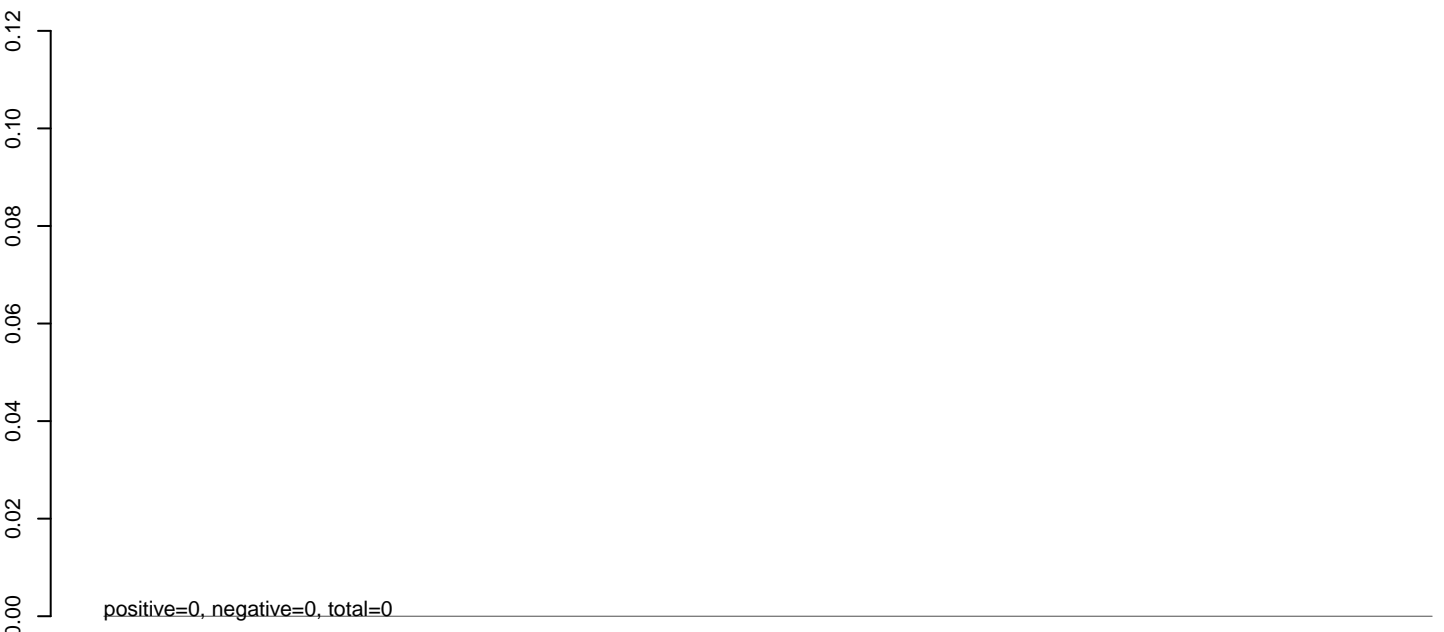
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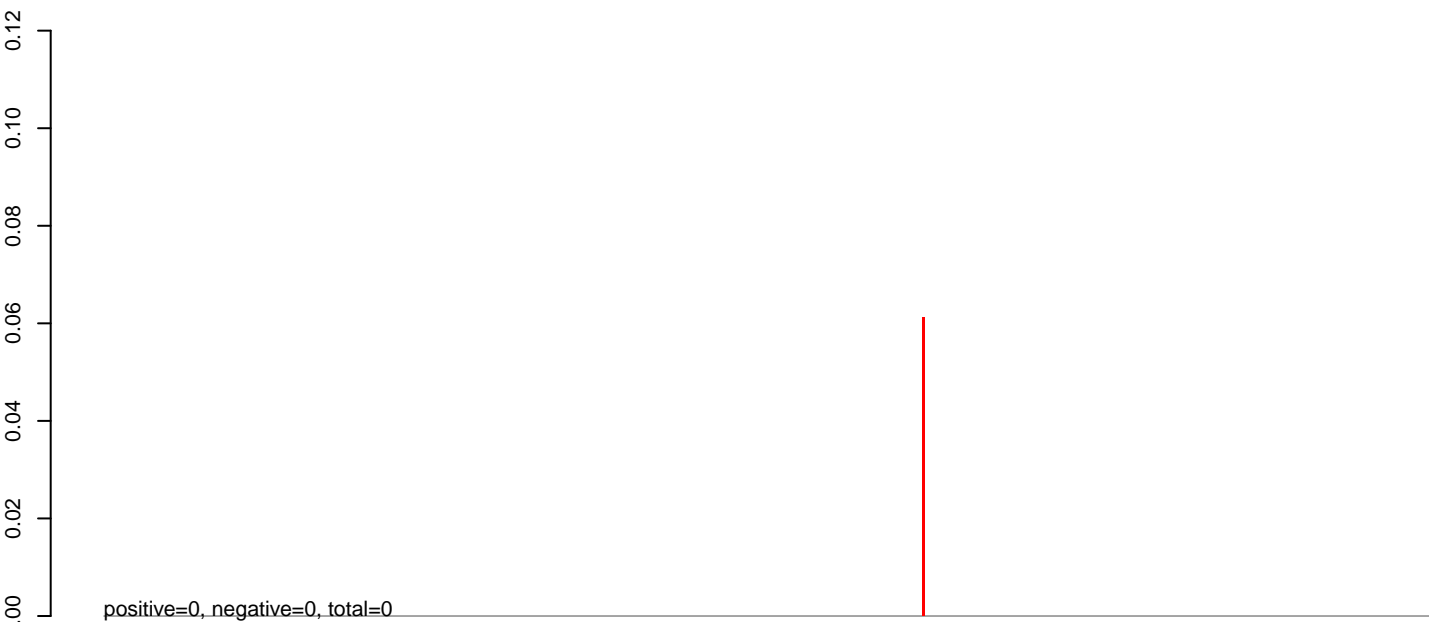


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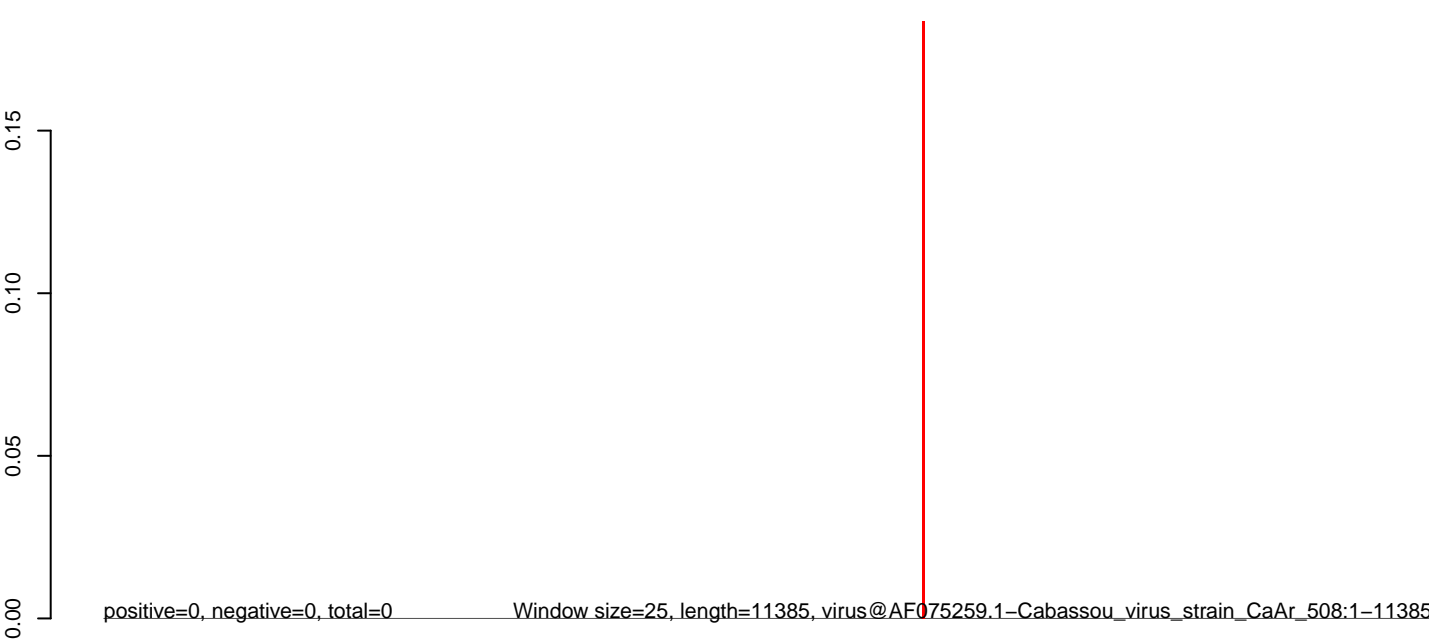
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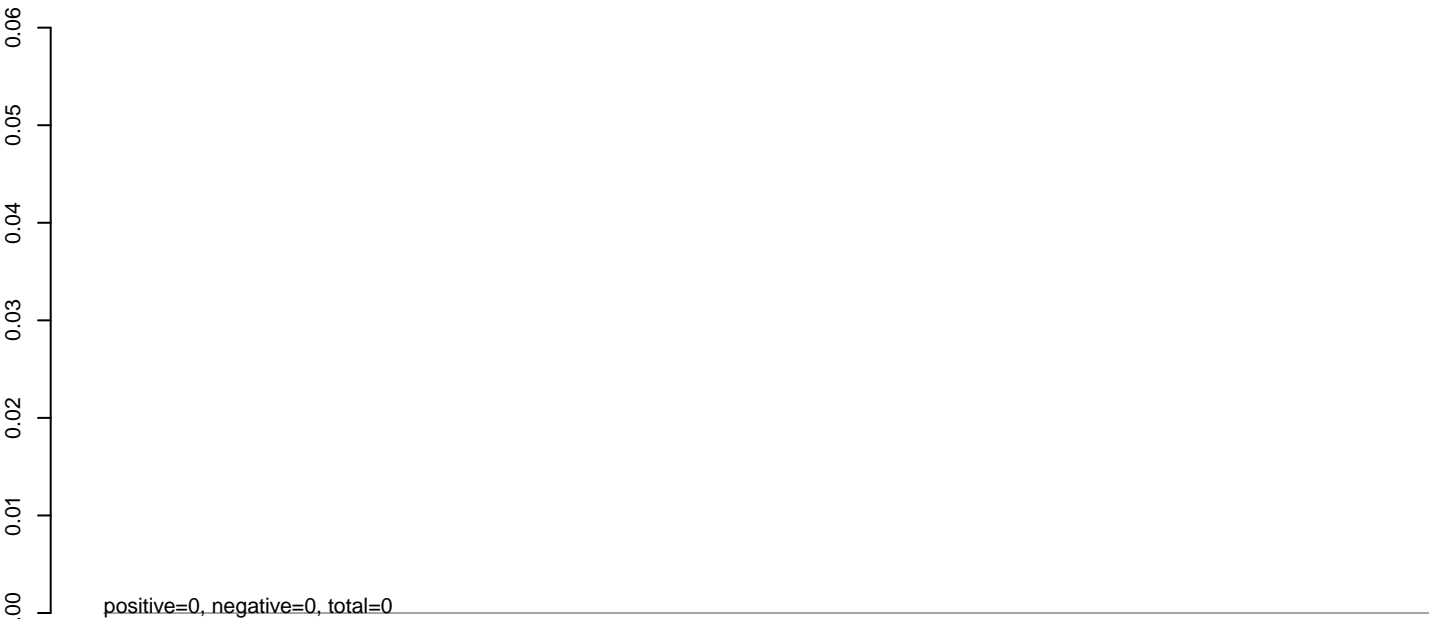


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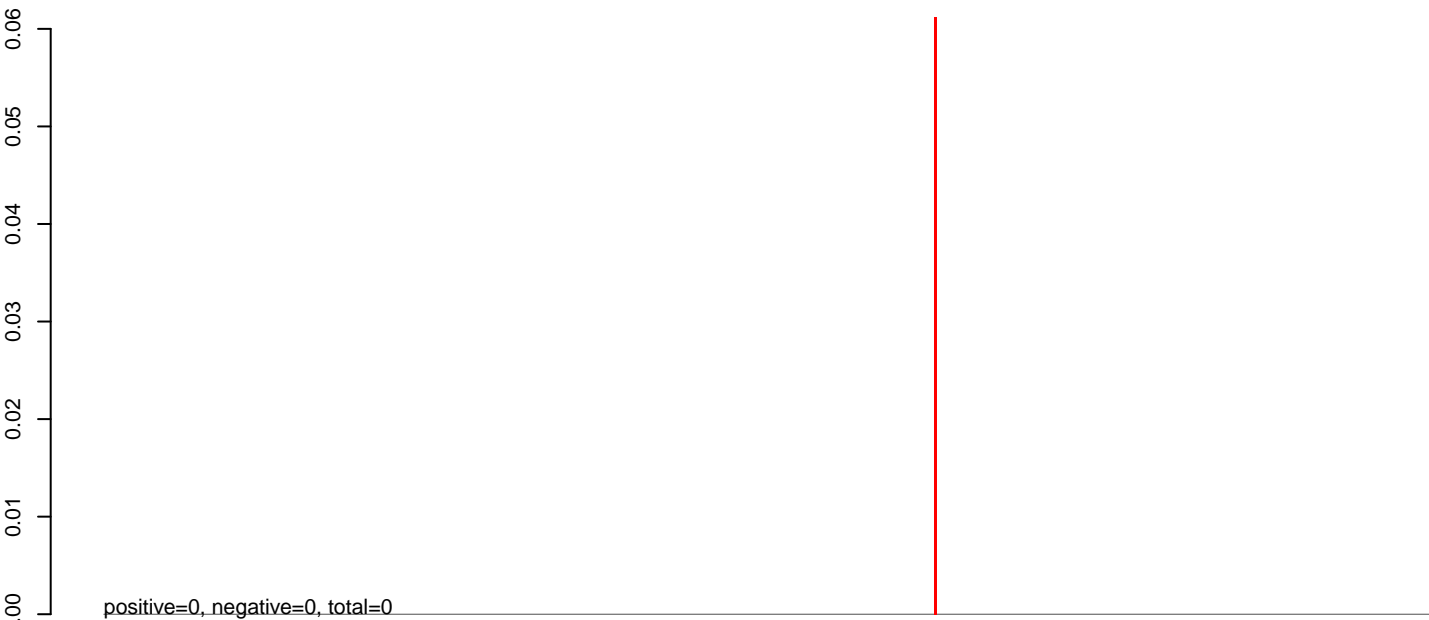


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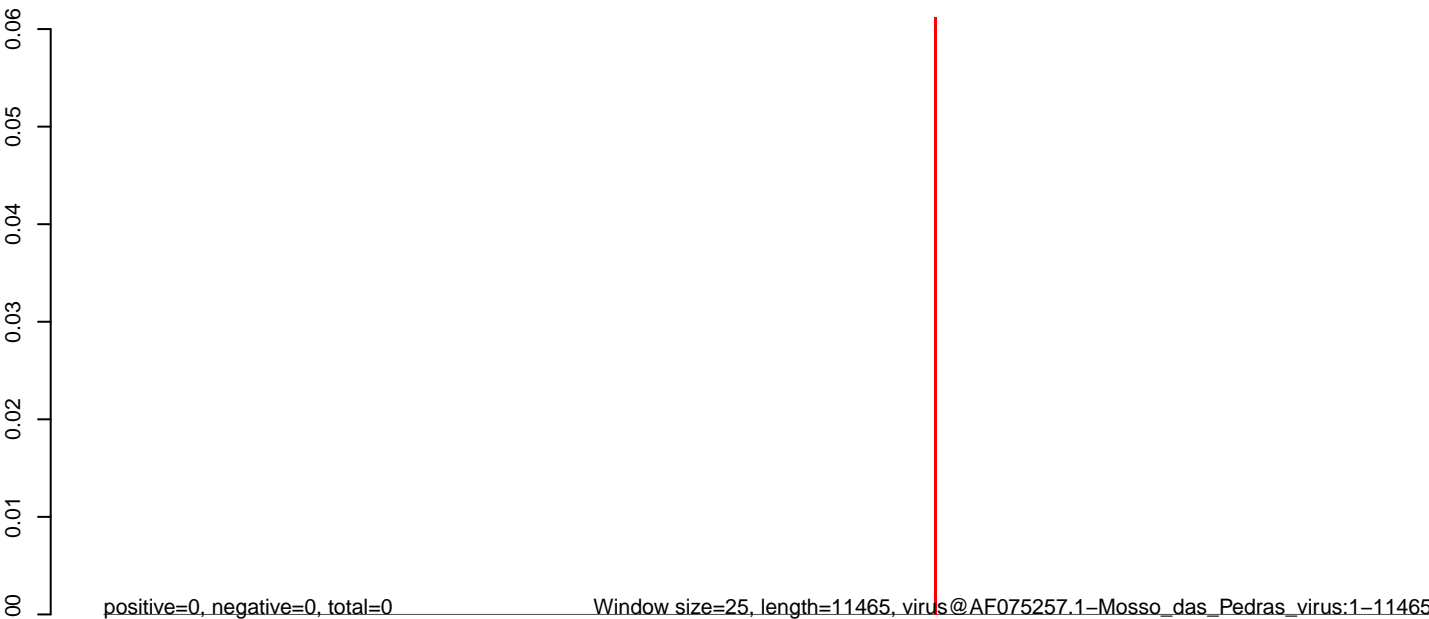
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AeAlbo_C636_B2_FHV_r54q.24_35.rep



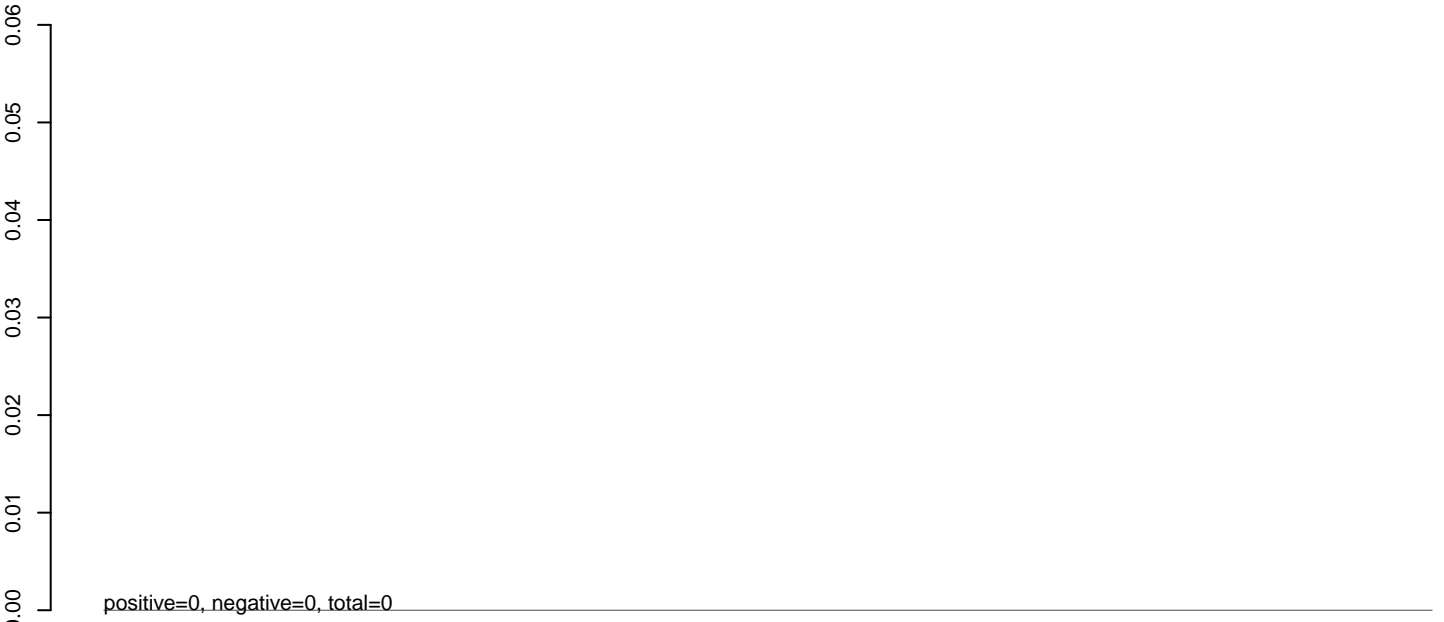
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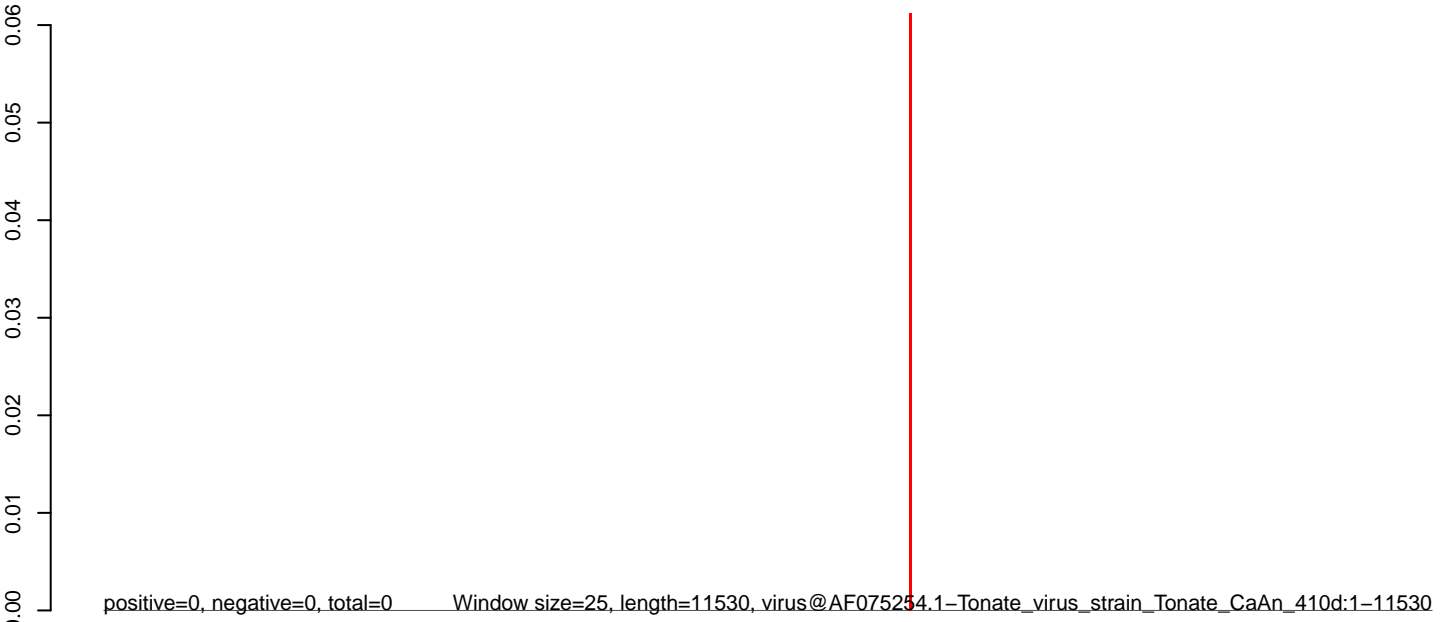
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AeAlbo_C636_B2_FHV_r54q.24_35.rep



AeAlbo_C636_B2_FHV_r54q.rep



0 2000 4000 6000 8000 10000 12000

AeAlbo_C636_B2_FHV_r54q.18_23.rep



AeAlbo_C636_B2_FHV_r54q.24_35.rep



AeAlbo_C636_B2_FHV_r54q.rep



0 5000 10000 15000 20000

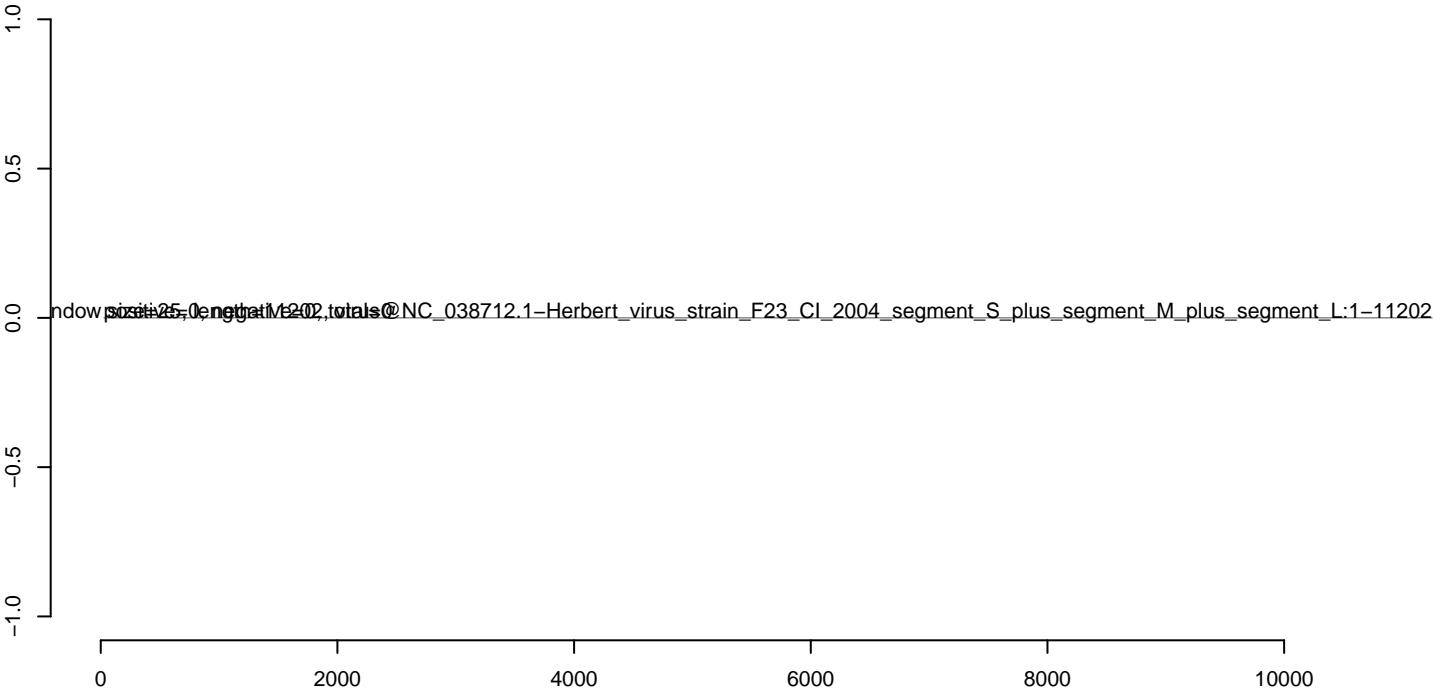
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AeAlbo_C636_B2_FHV_r54q.24_35.rep



AeAlbo_C636_B2_FHV_r54q.rep



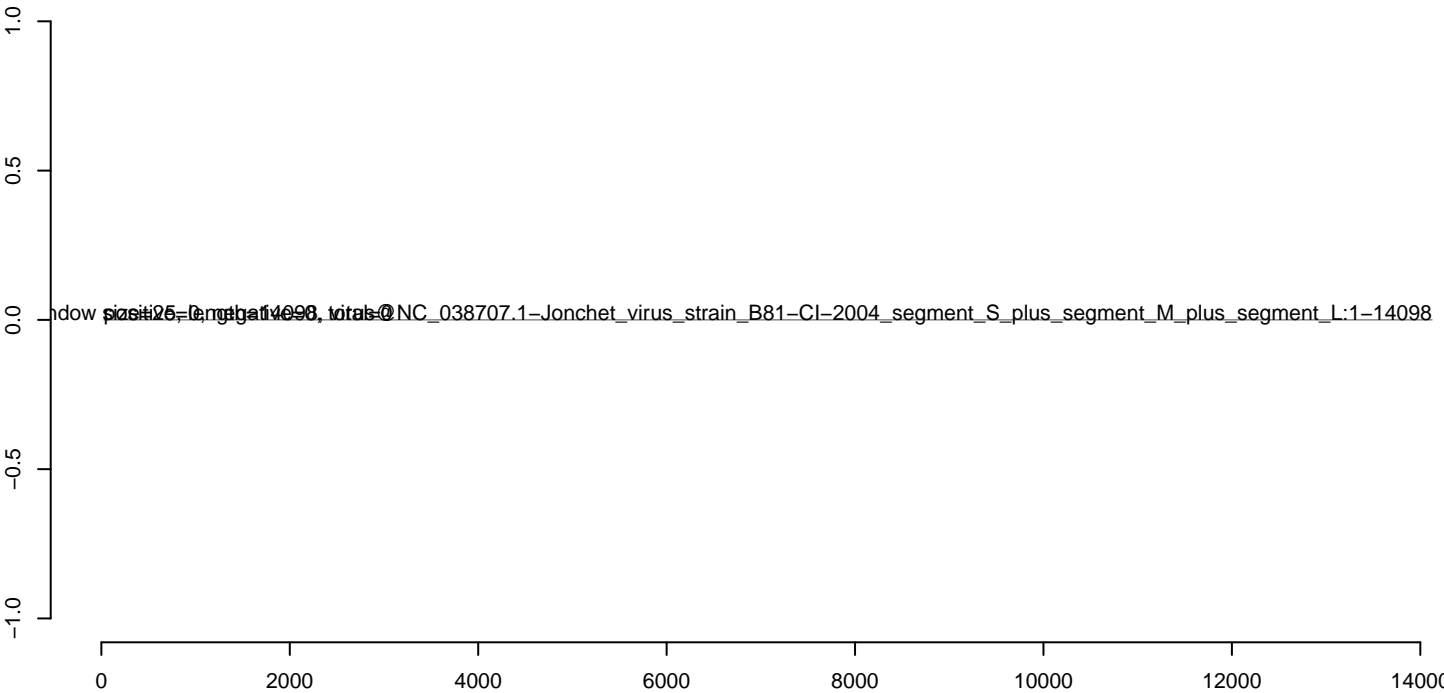
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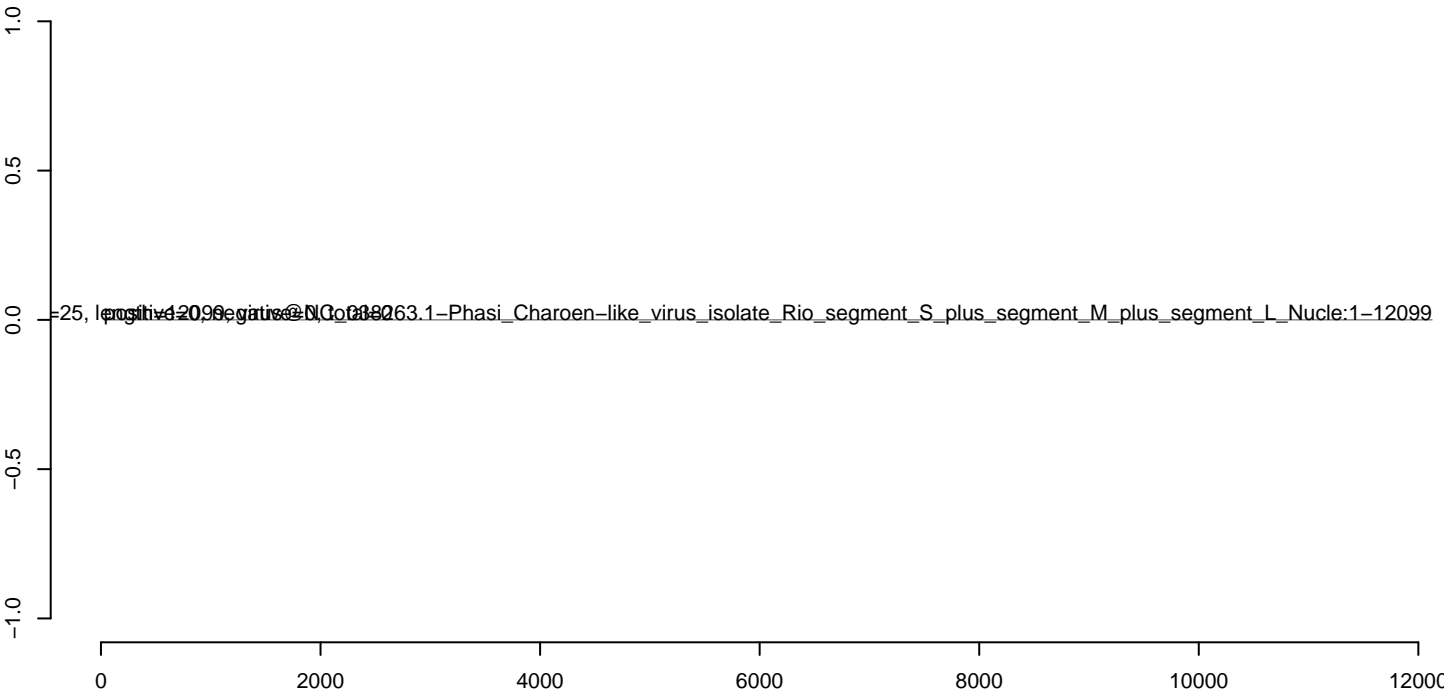
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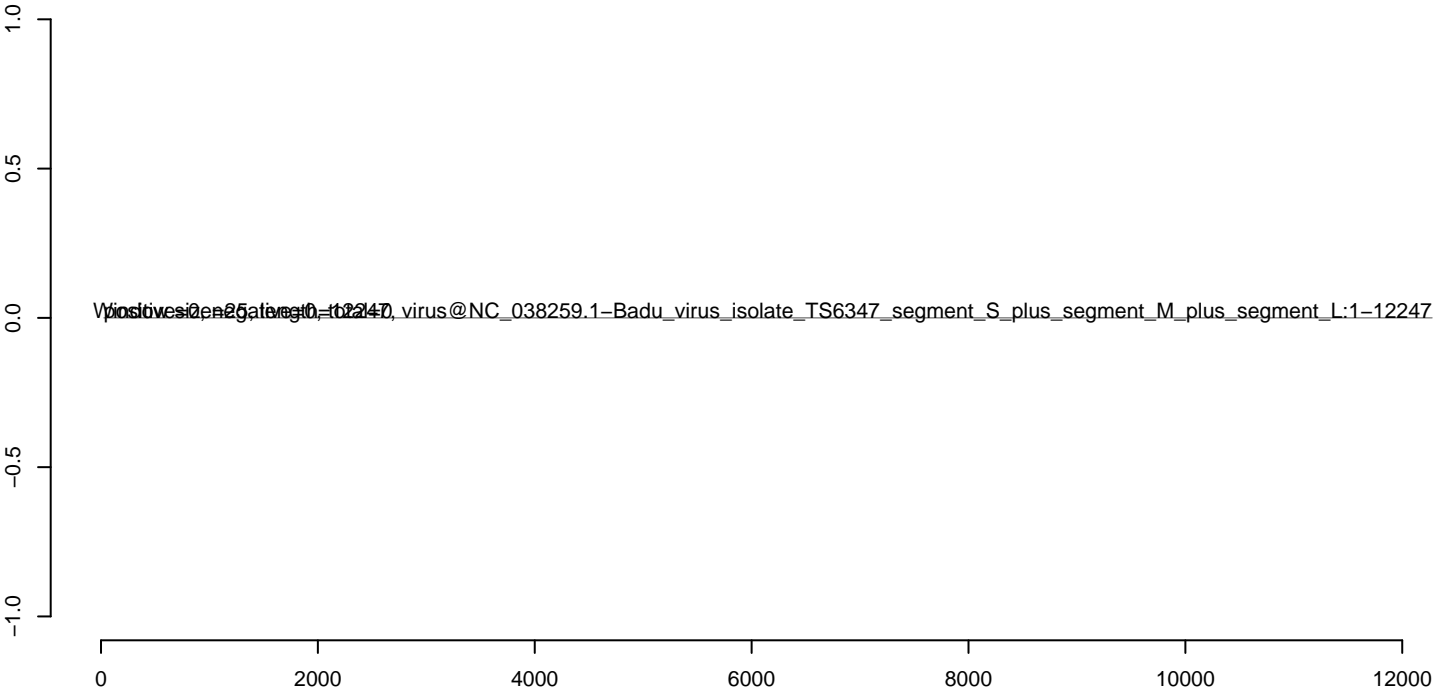
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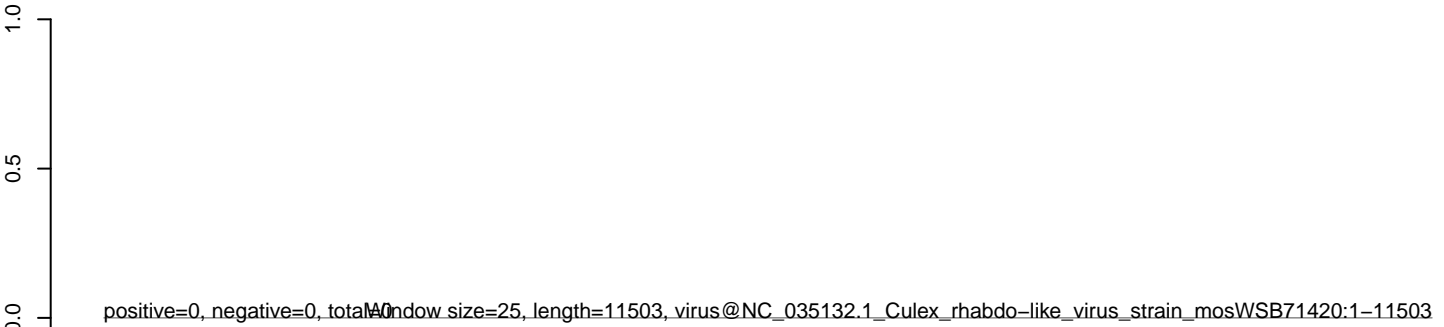
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AeAlbo_C636_B2_FHV_r54q.rep



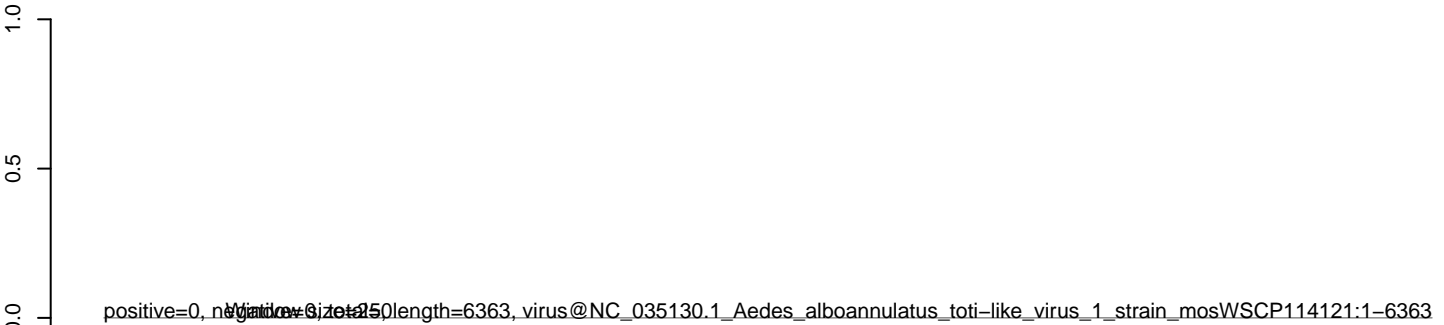
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AeAlbo_C636_B2_FHV_r54q.rep



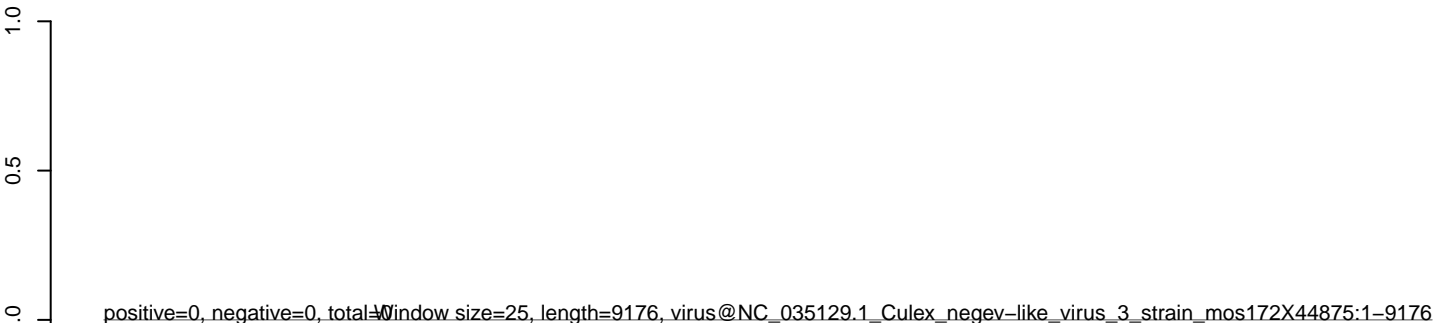
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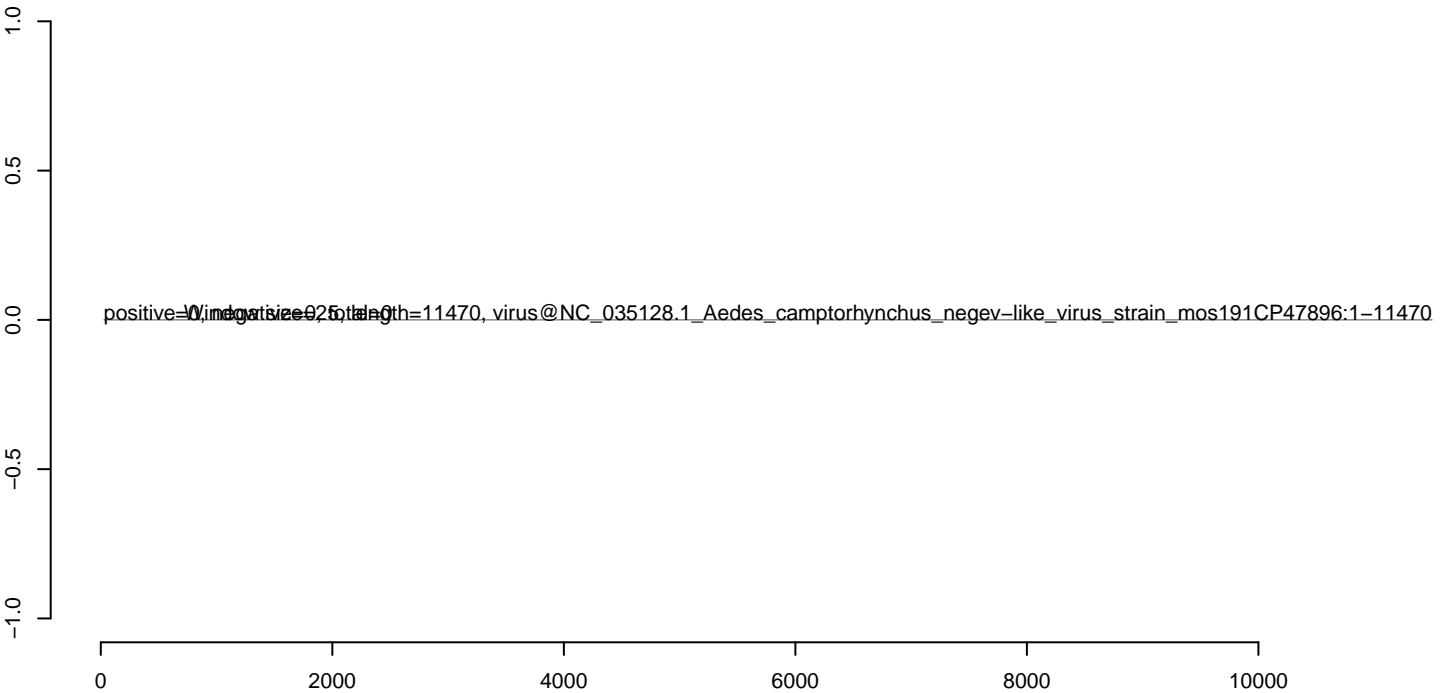
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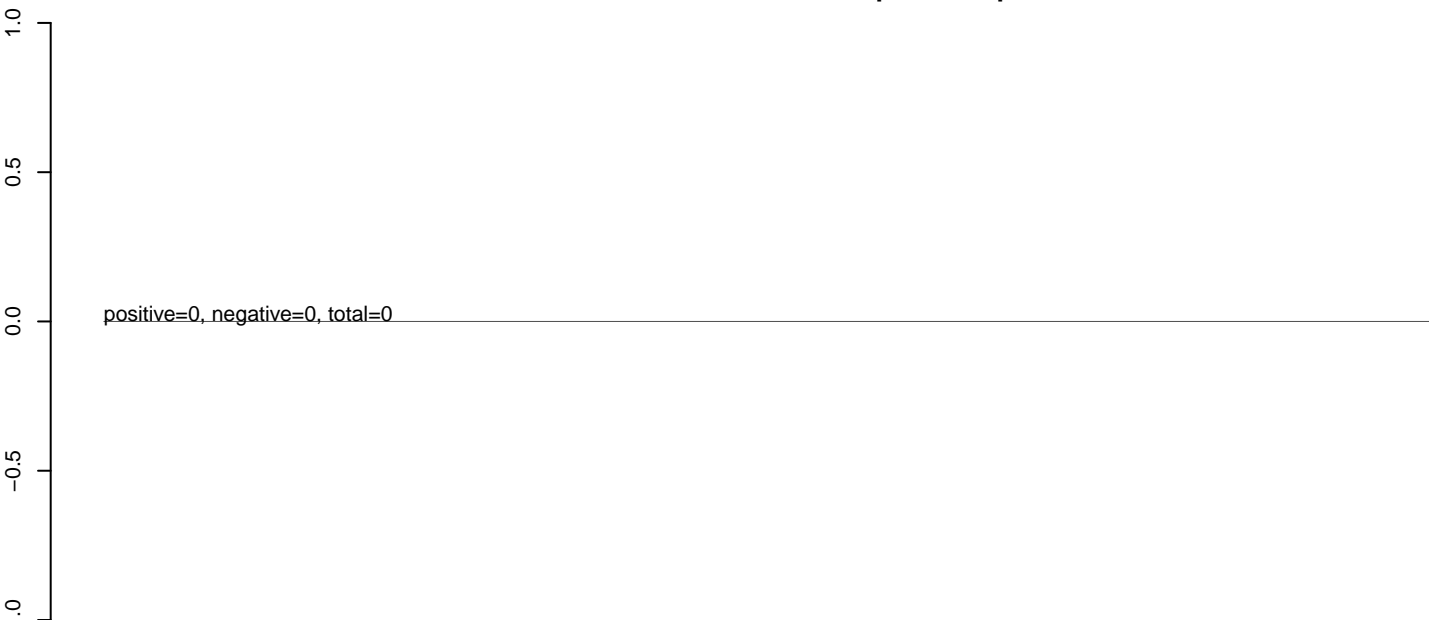
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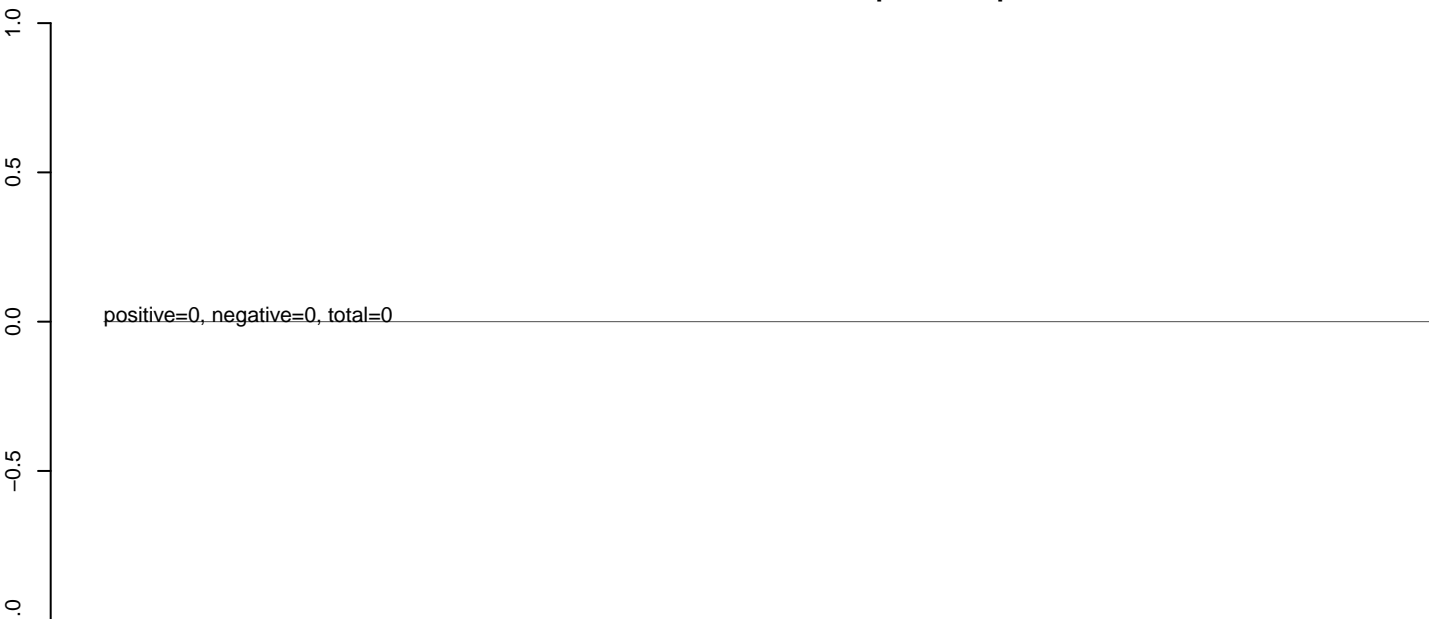
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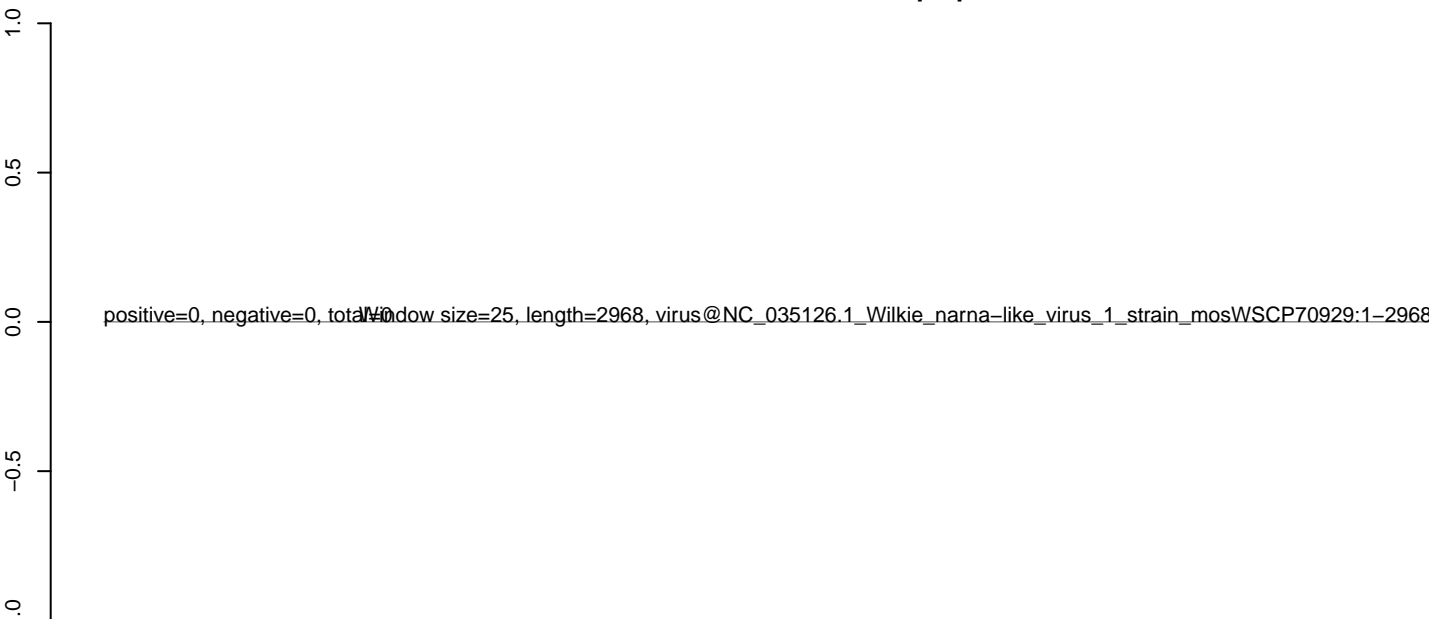
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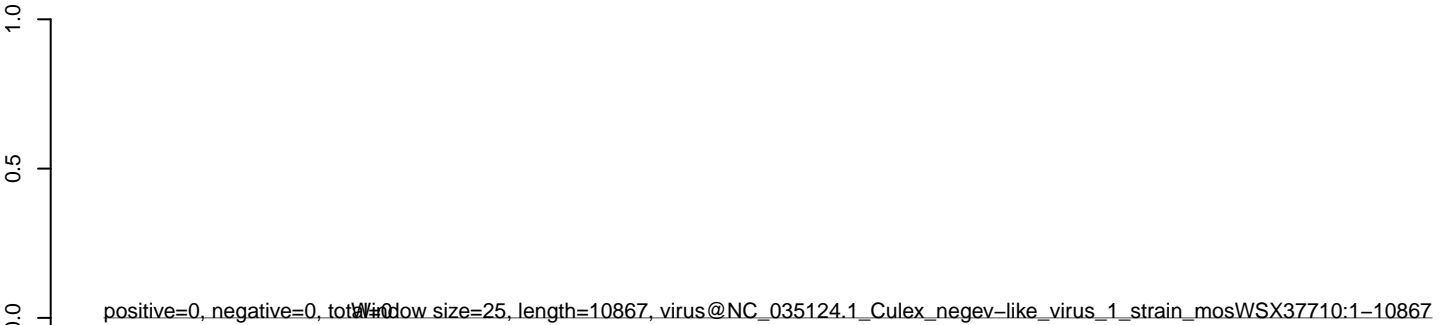
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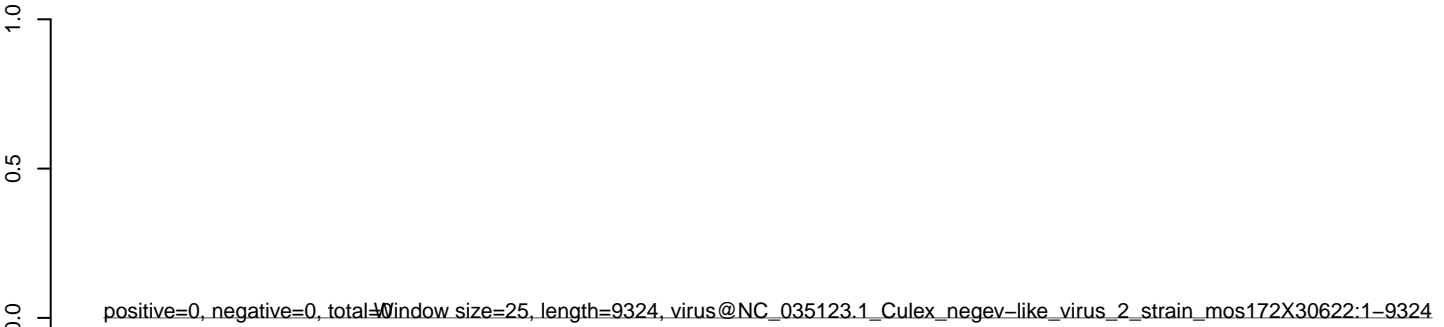
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AeAlbo_C636_B2_FHV_r54q.rep



0 2000 4000 6000 8000

AeAlbo_C636_B2_FHV_r54q.18_23.rep



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AeAlbo_C636_B2_FHV_r54q.rep



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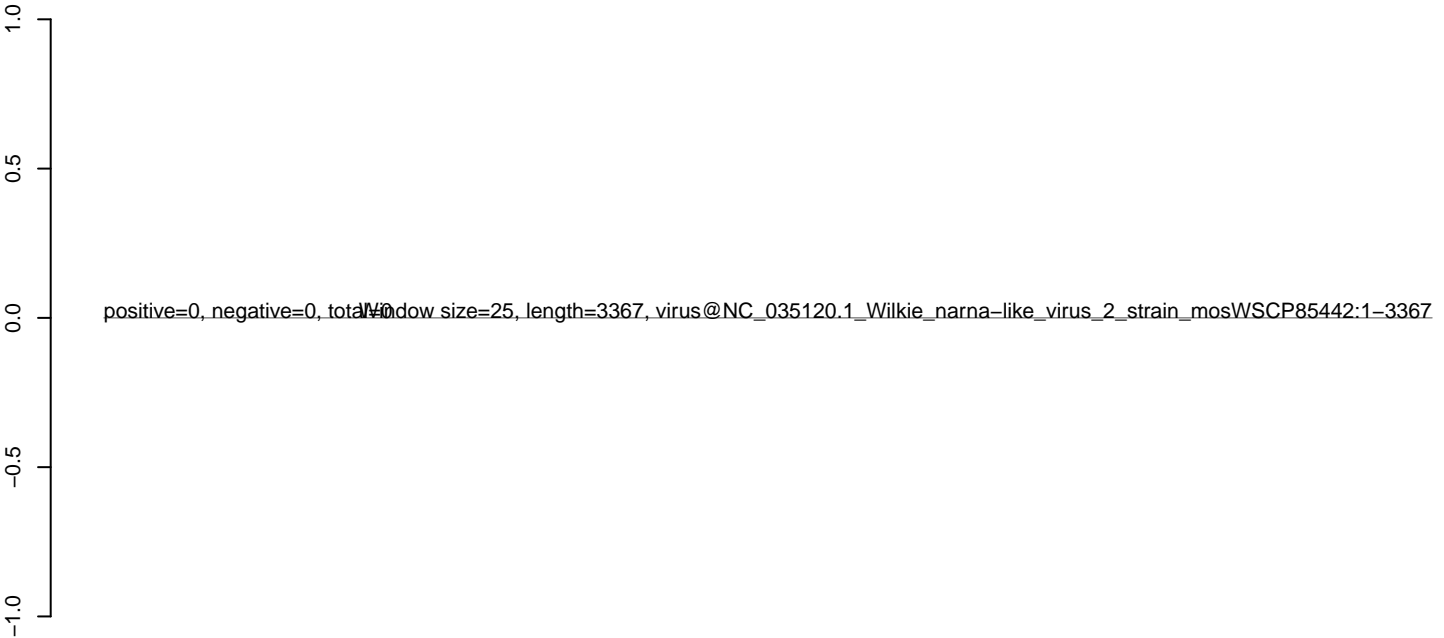
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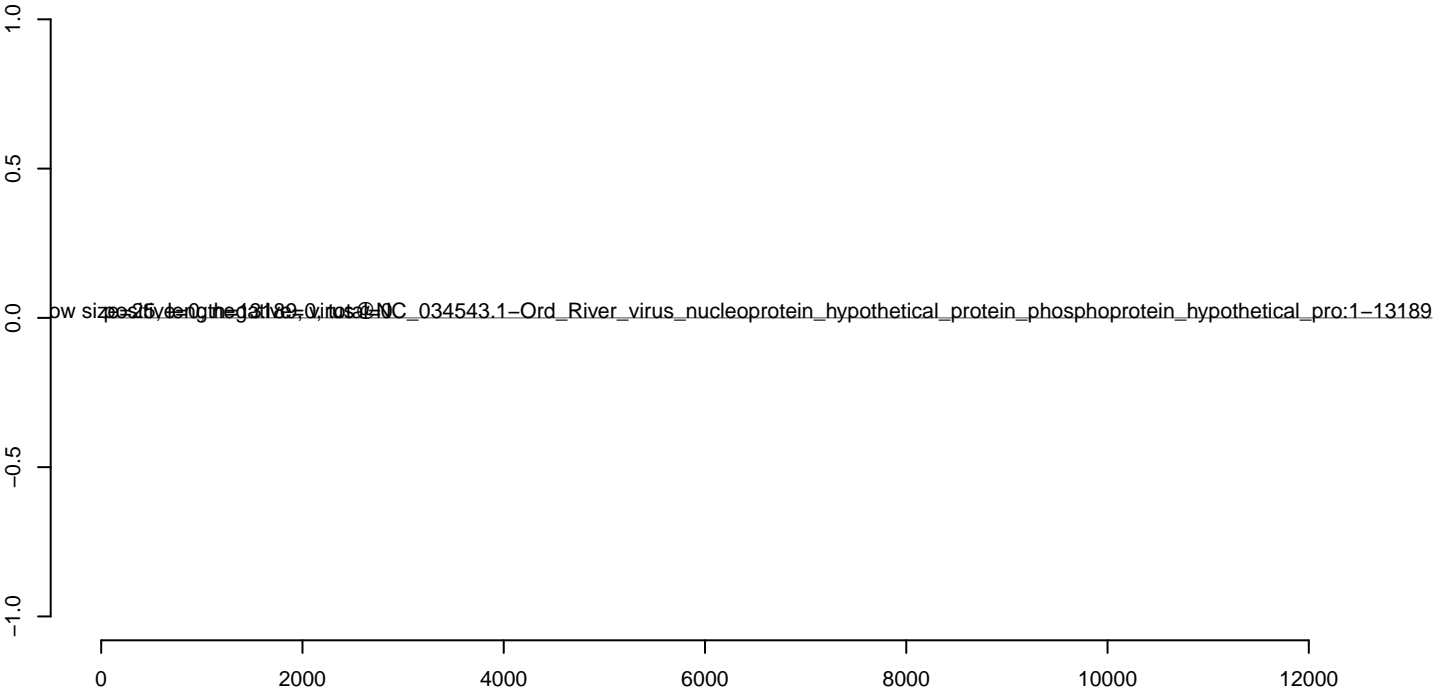
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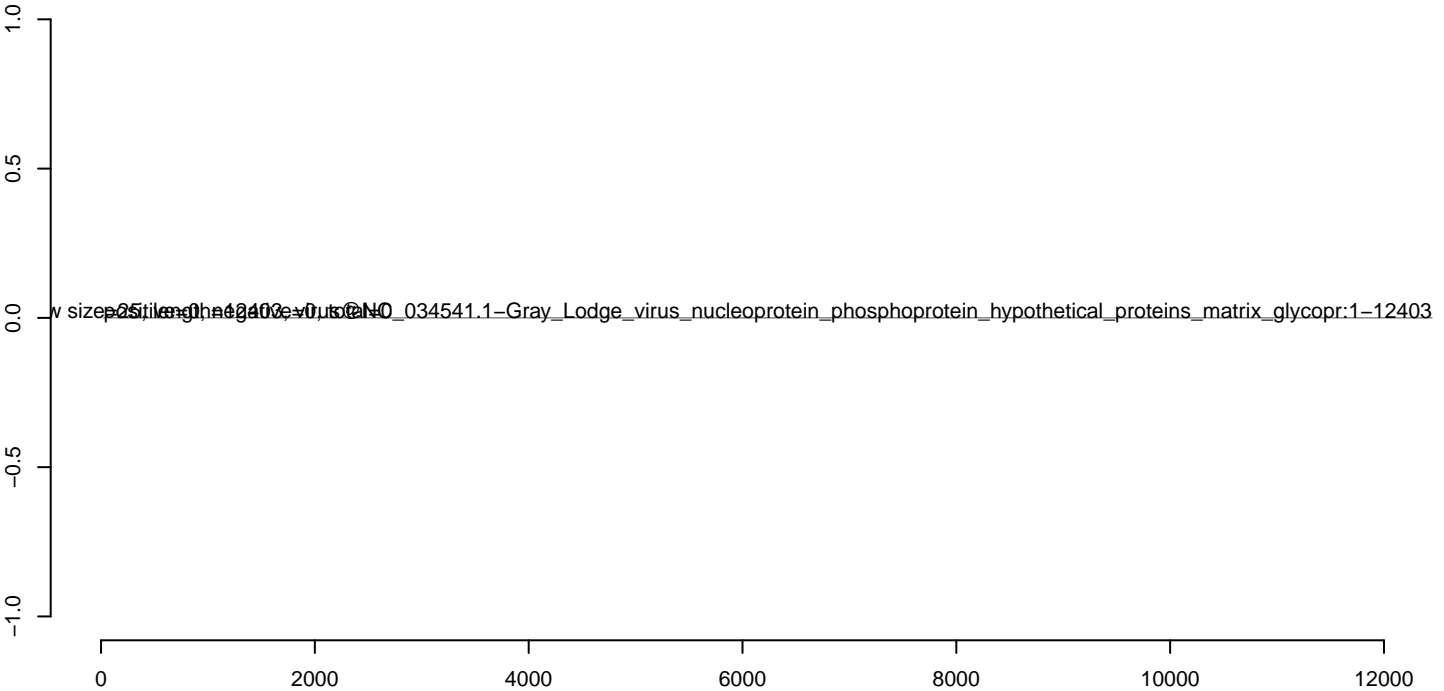
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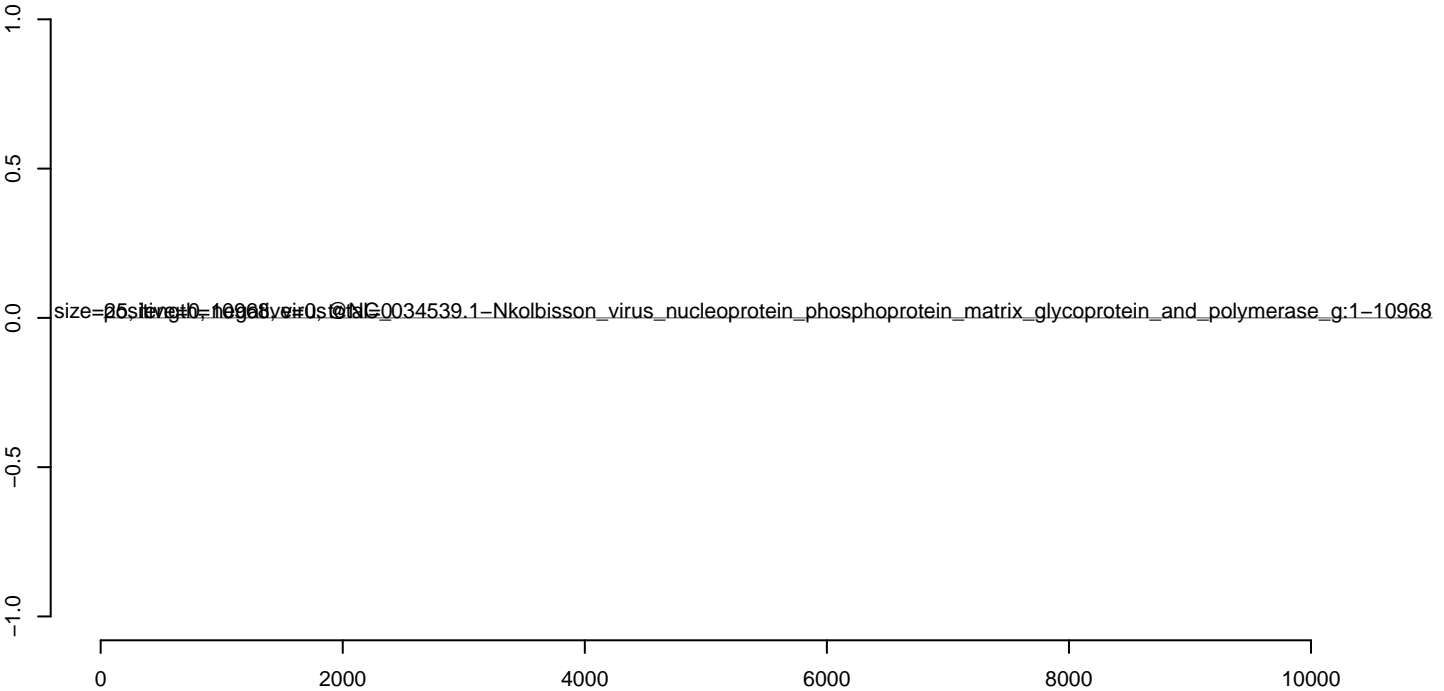
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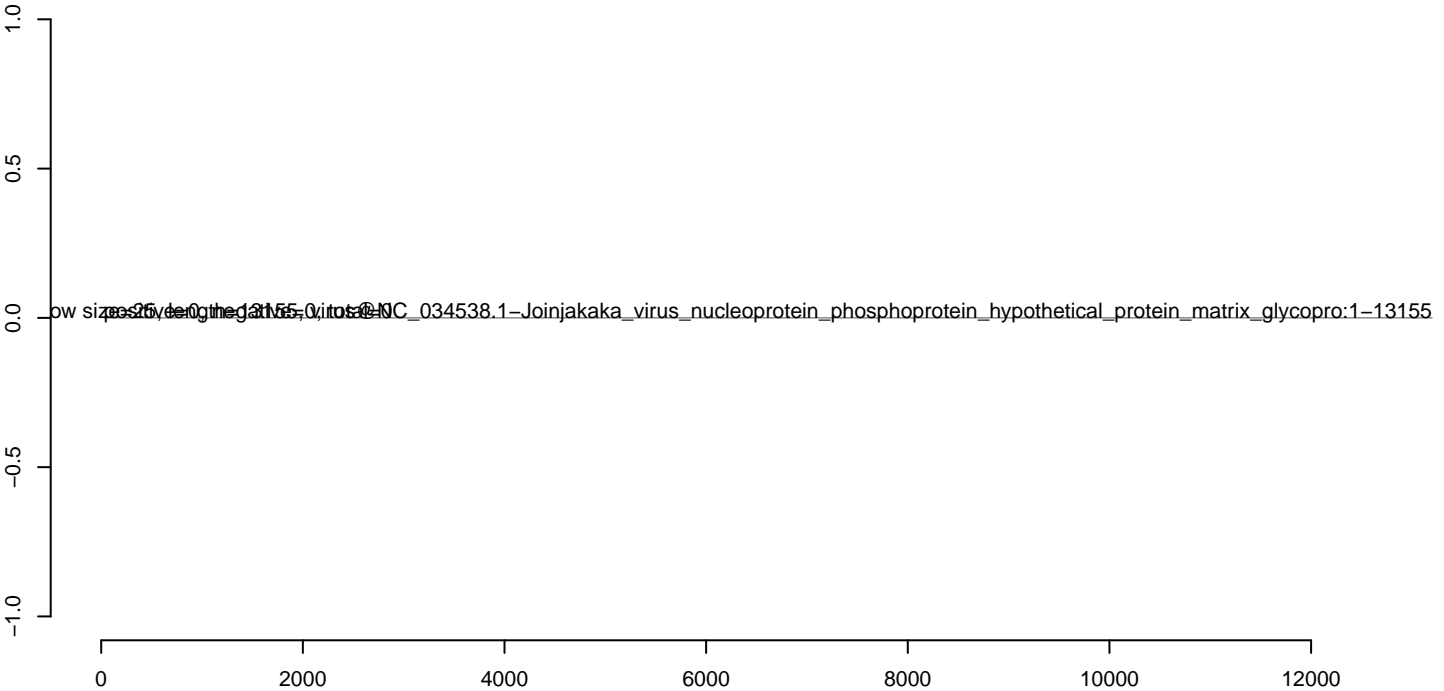
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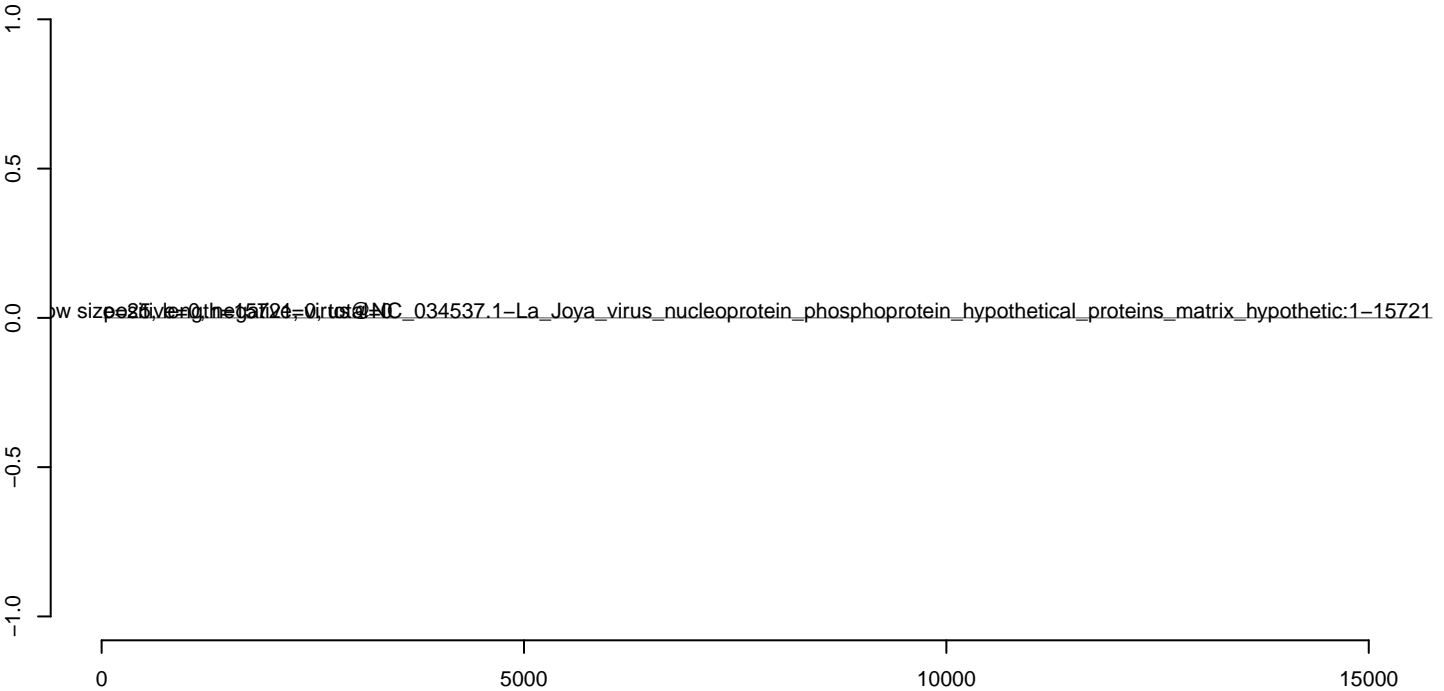
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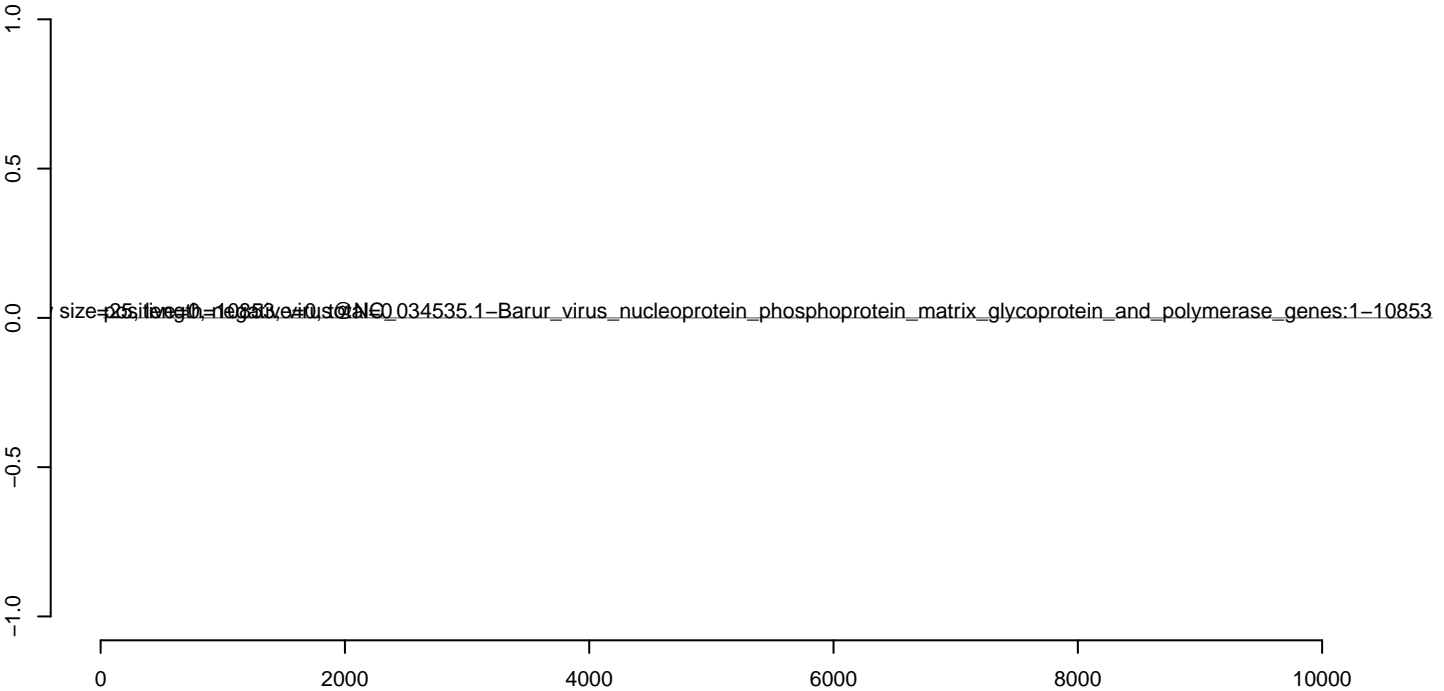
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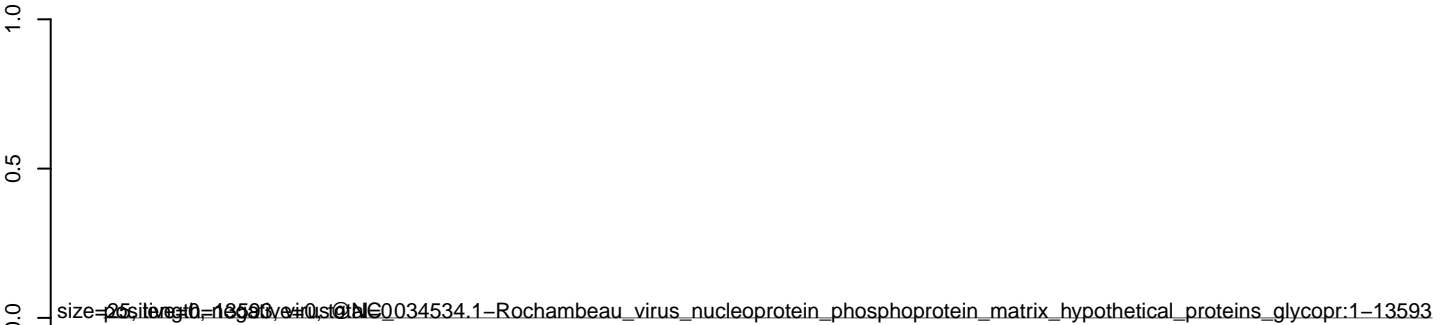
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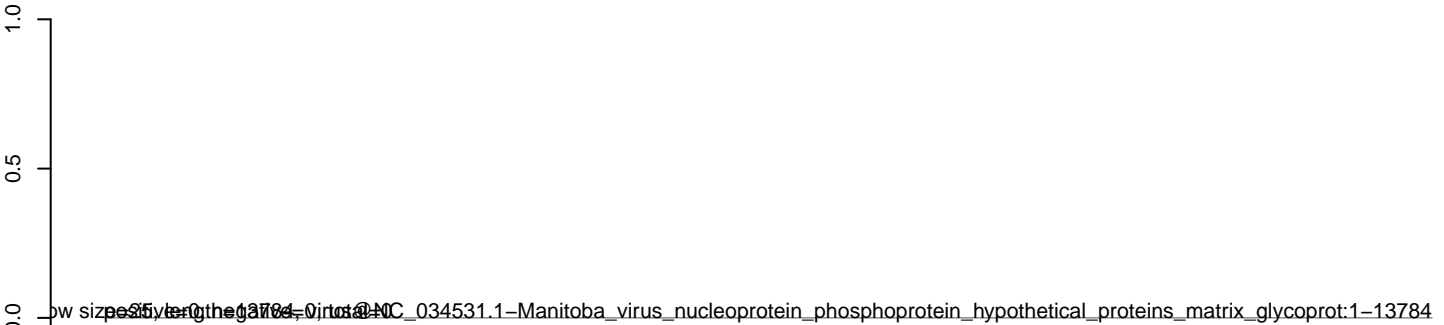
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AeAlbo_C636_B2_FHV_r54q.rep



low size: 51704-0 into @NC_034531.1-Manitoba_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycoprot:1-13784

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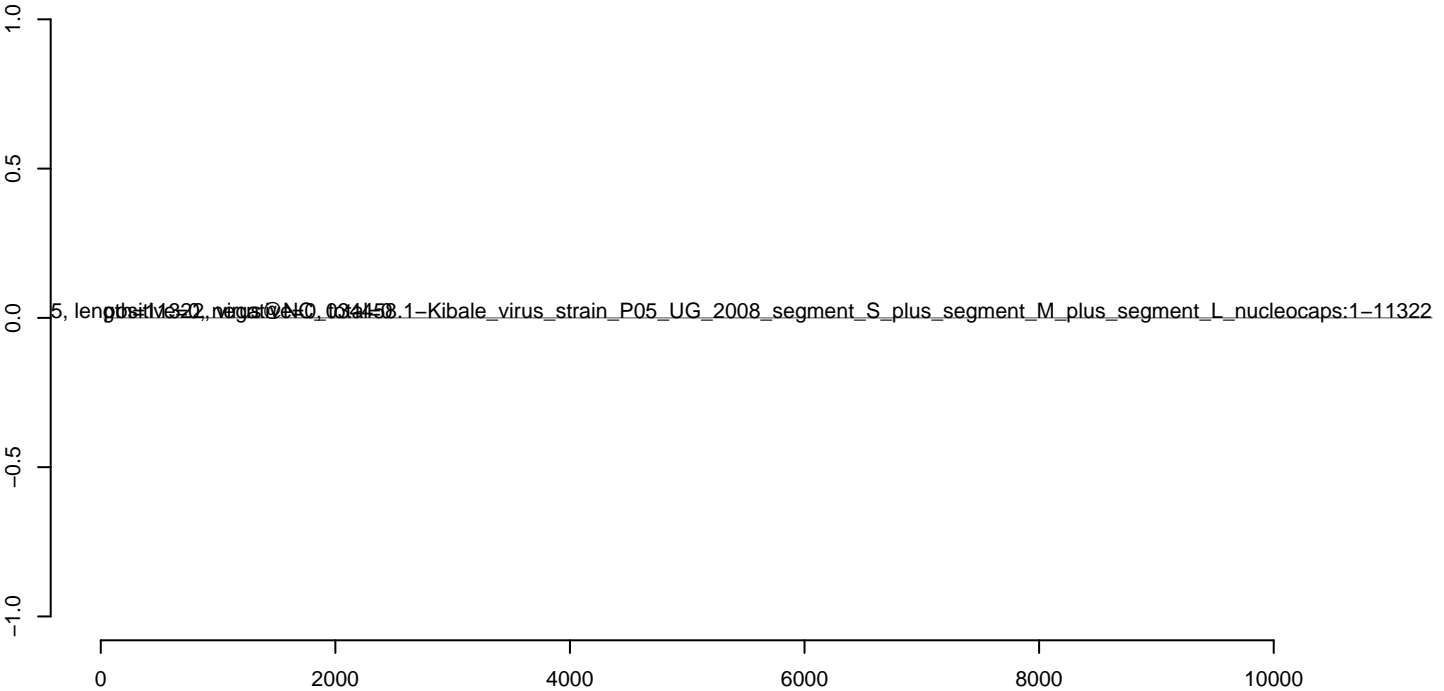
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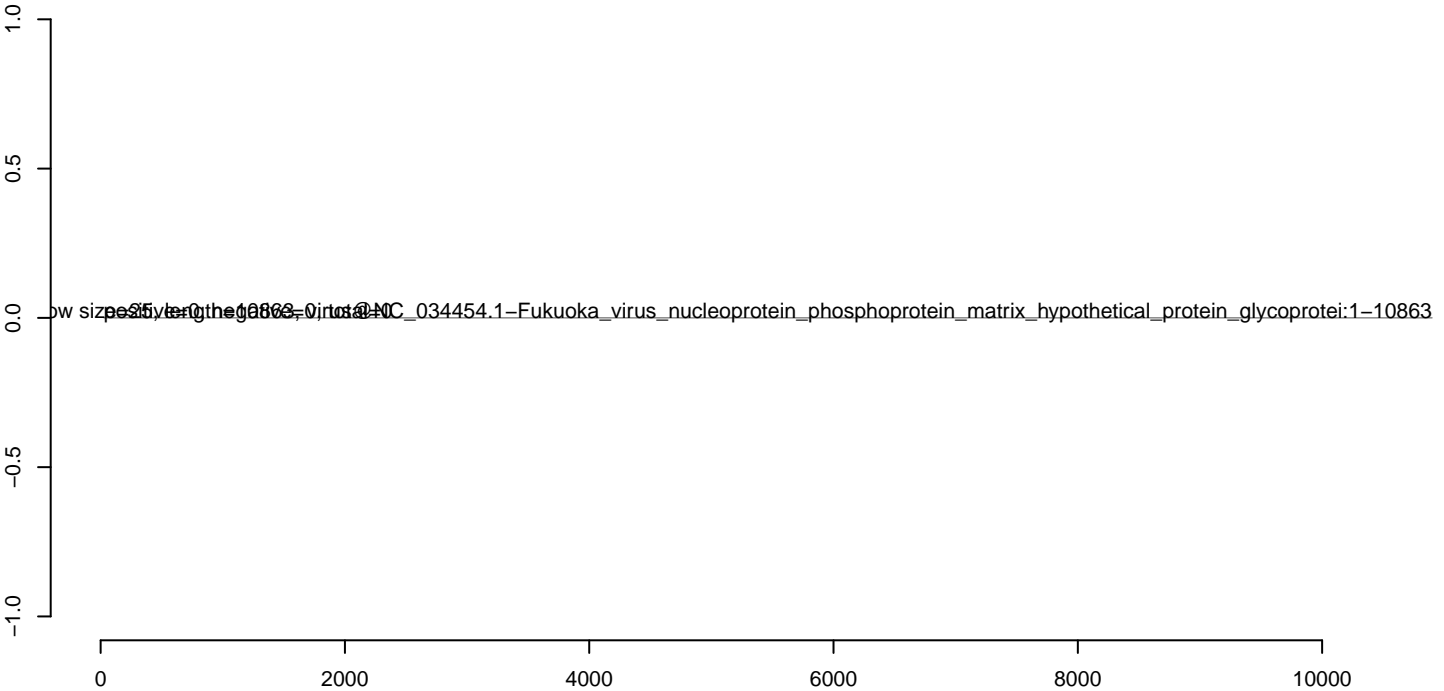
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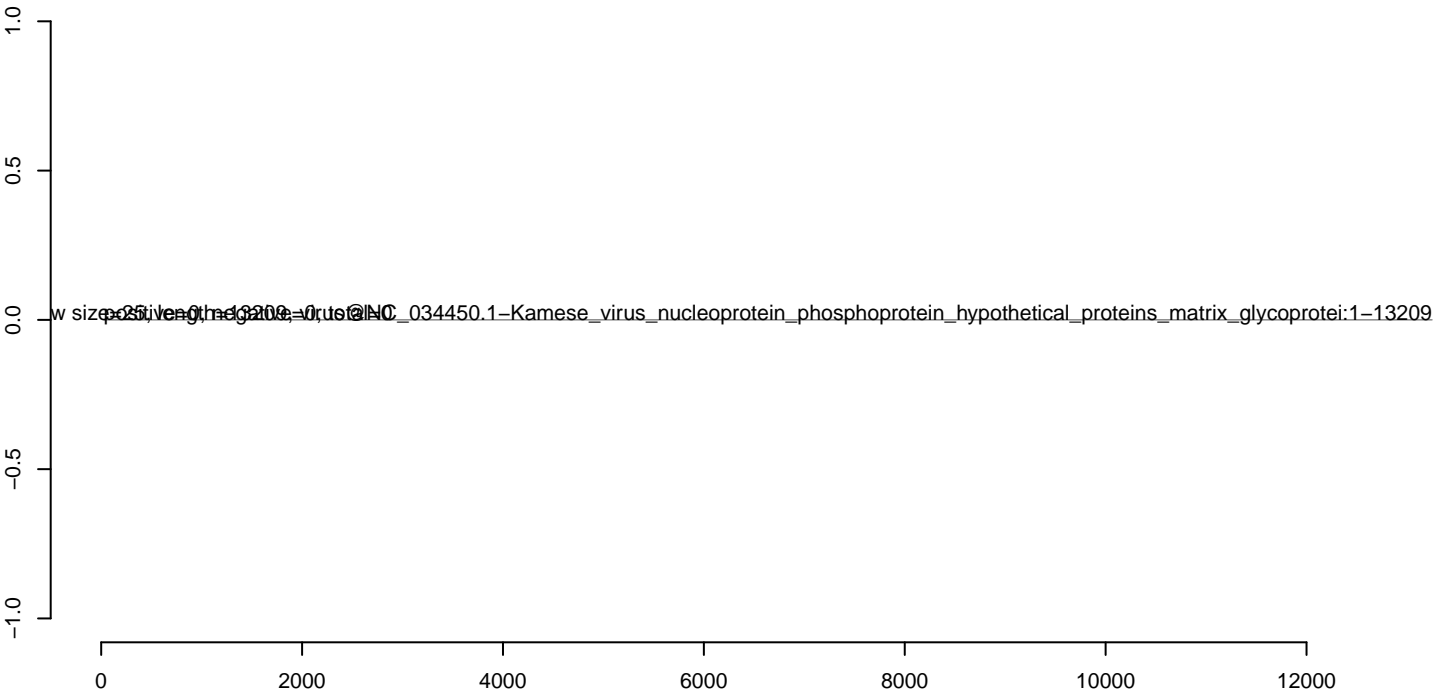
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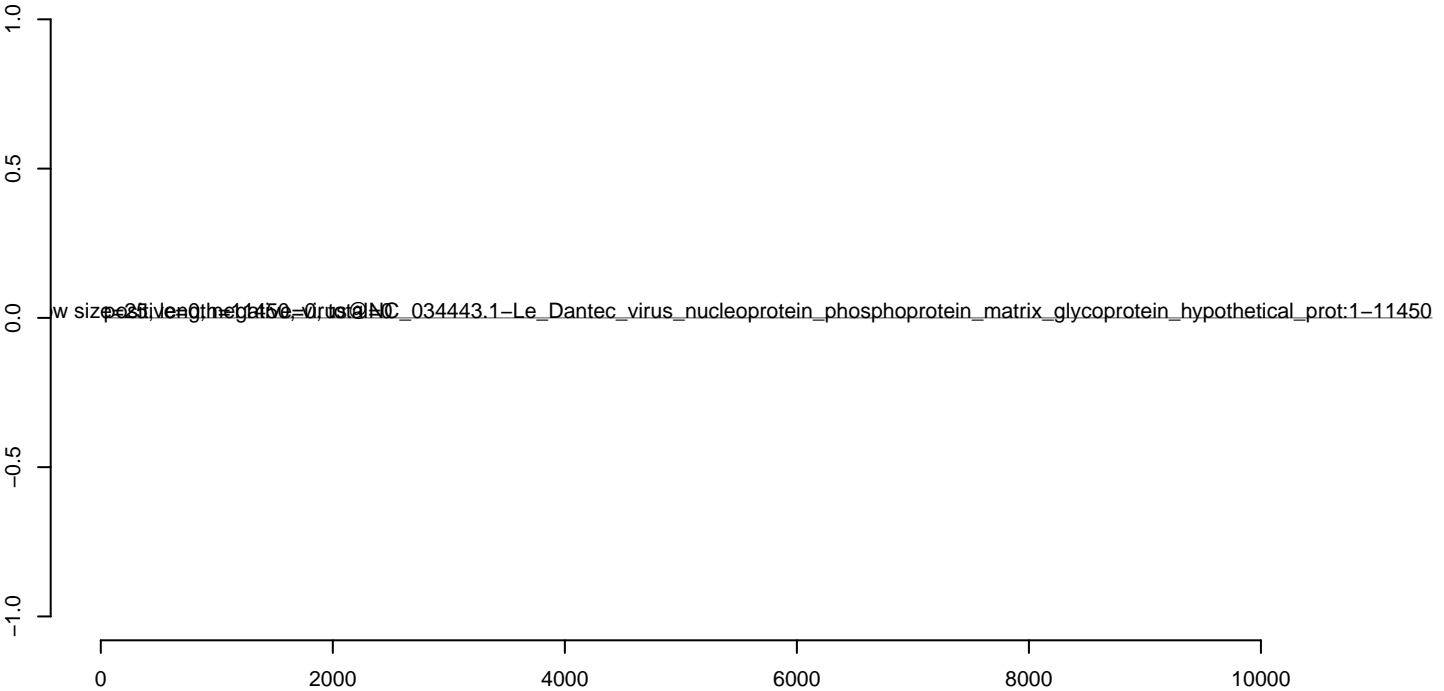
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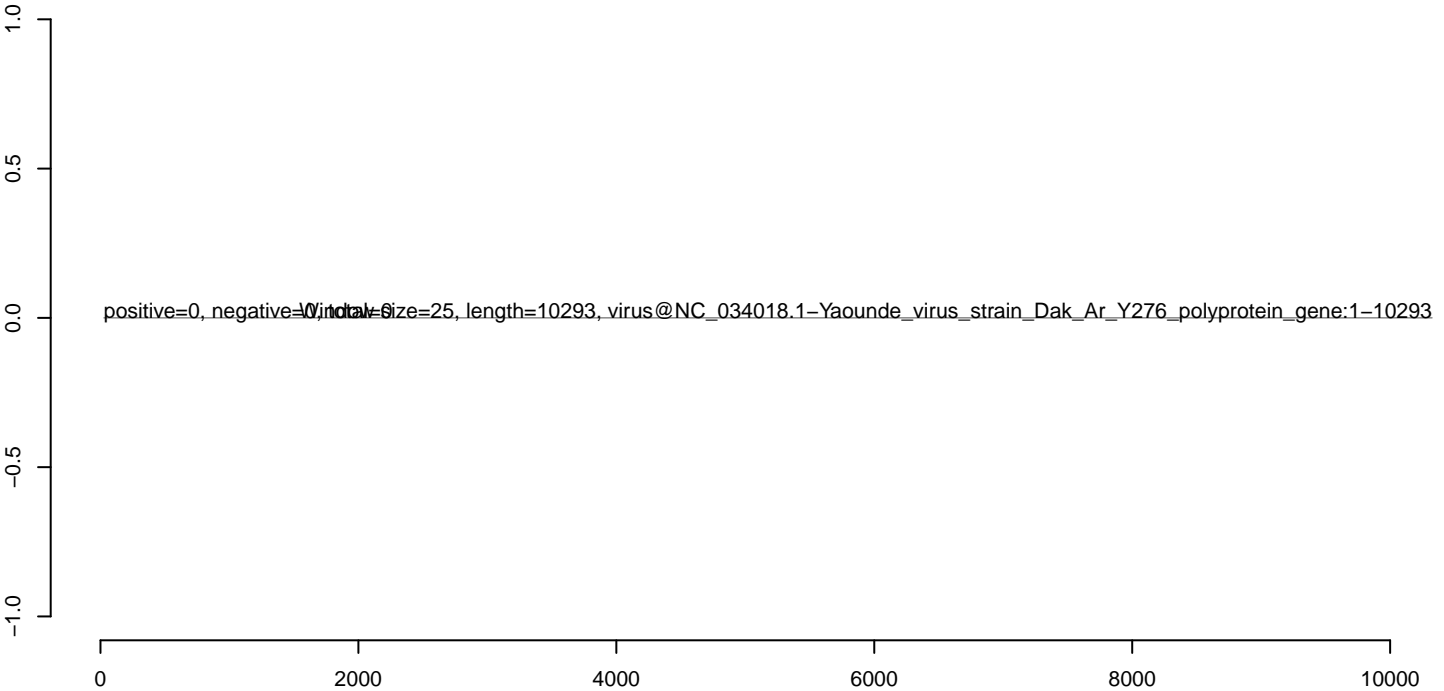
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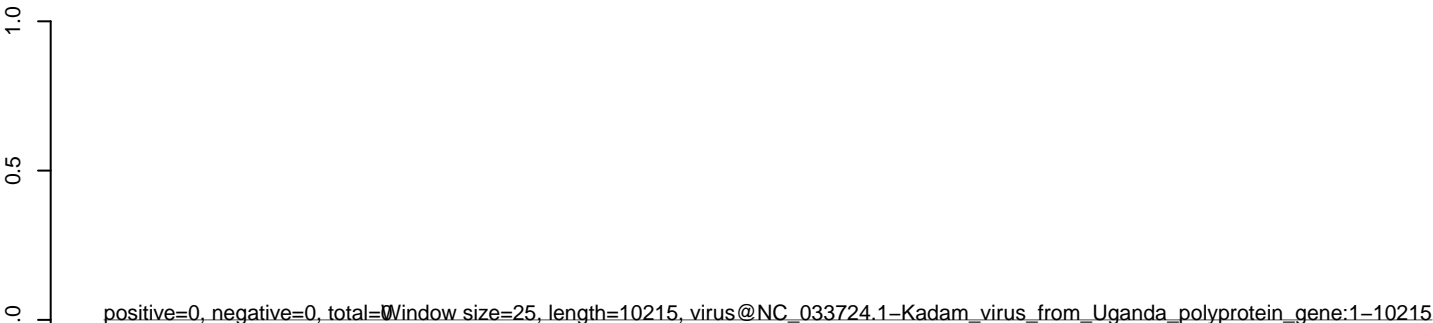
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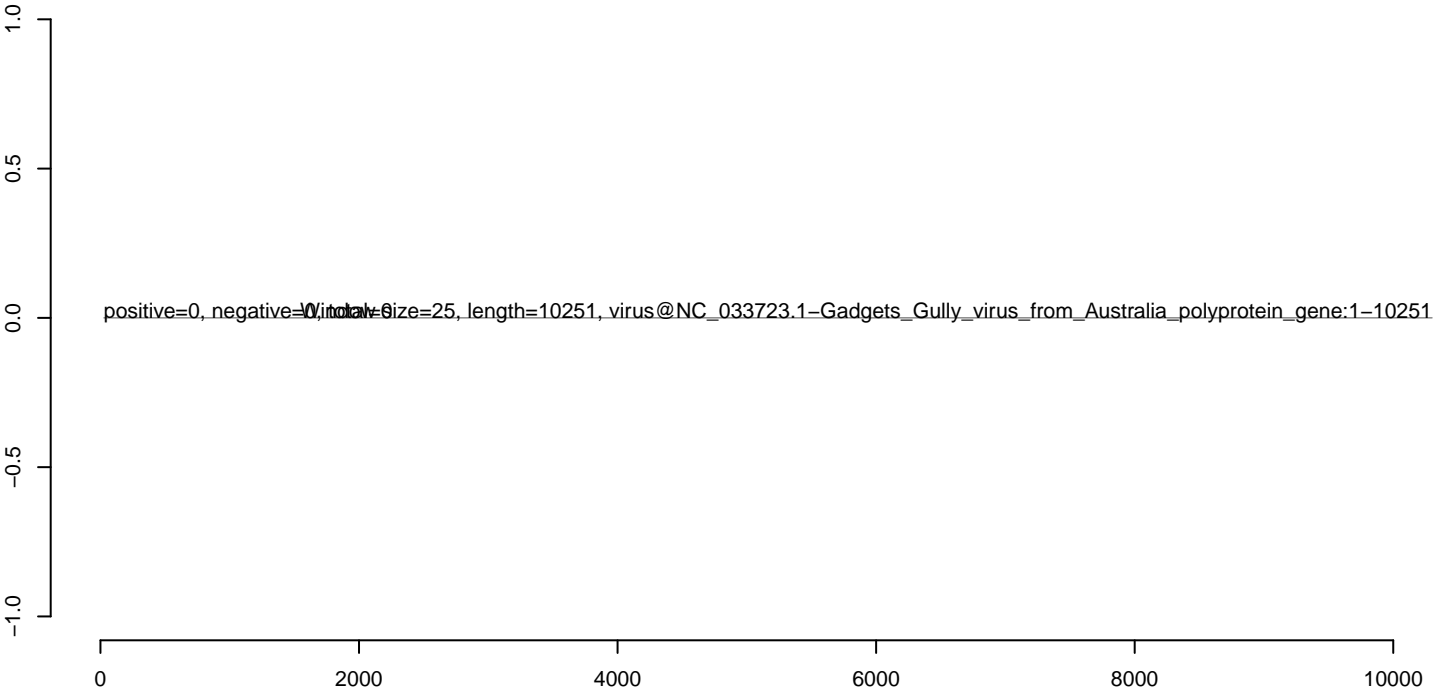
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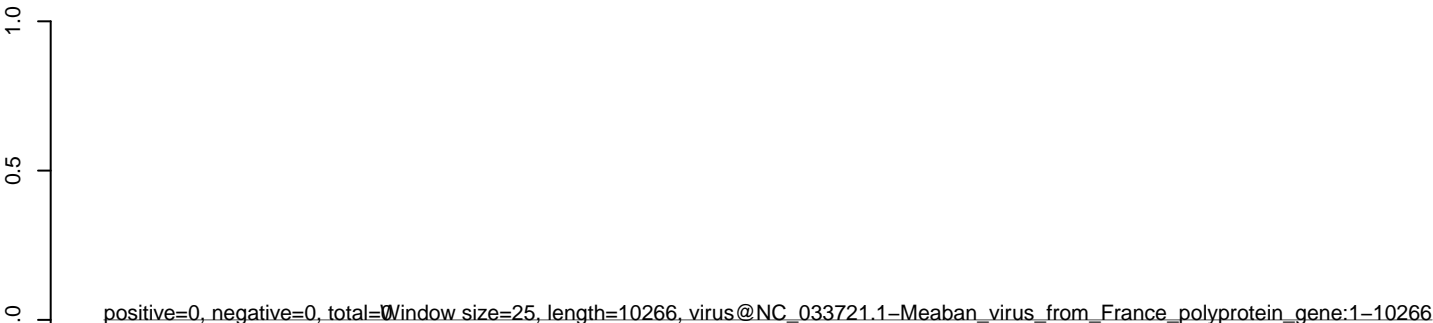
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AeAlbo_C636_B2_FHV_r54q.rep



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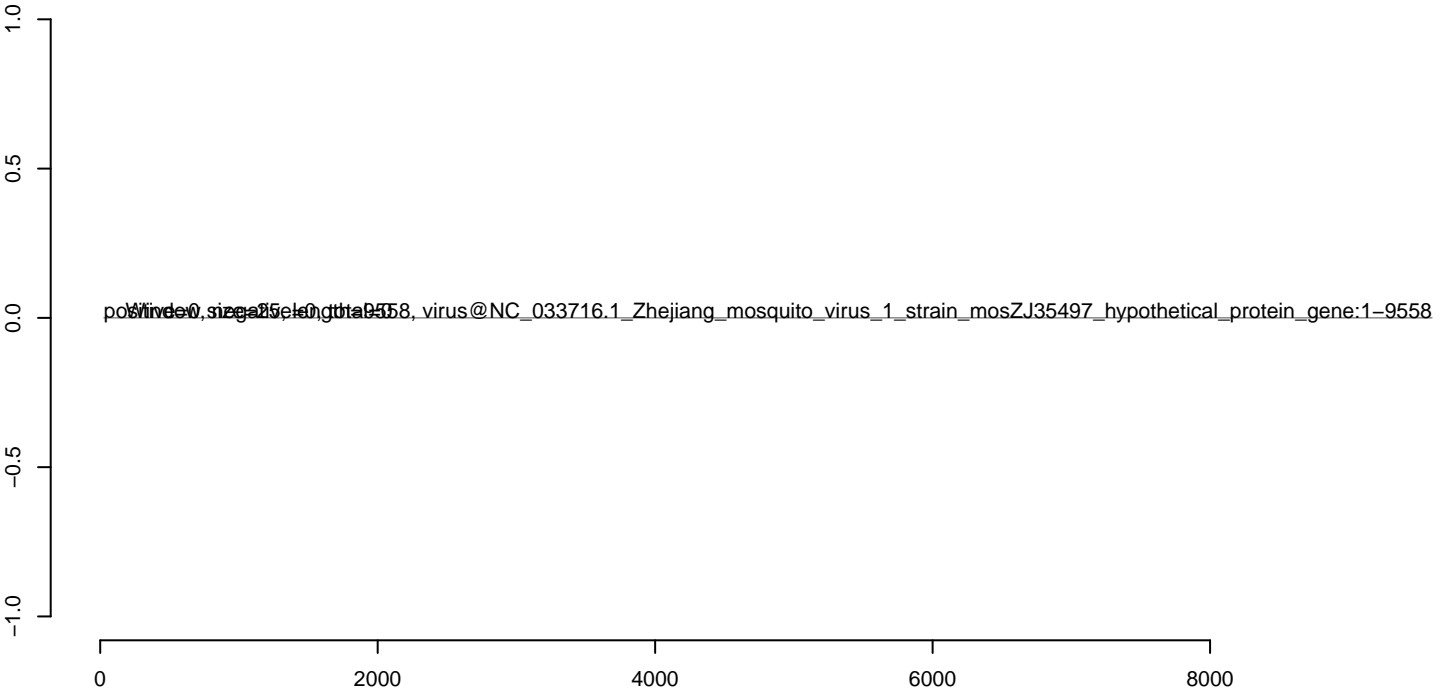
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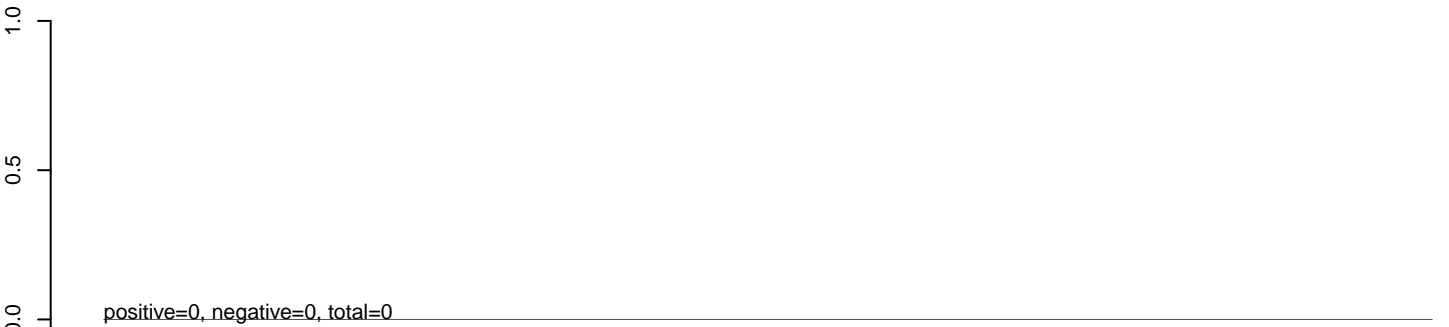
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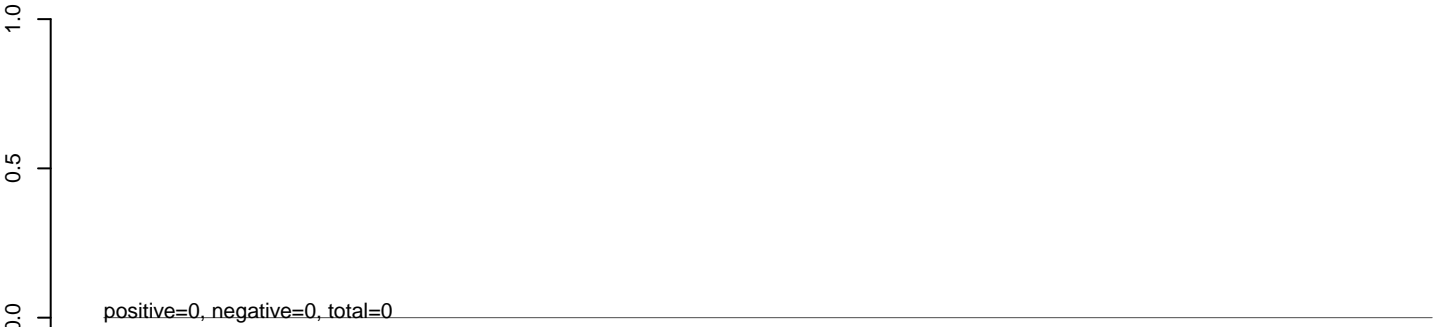
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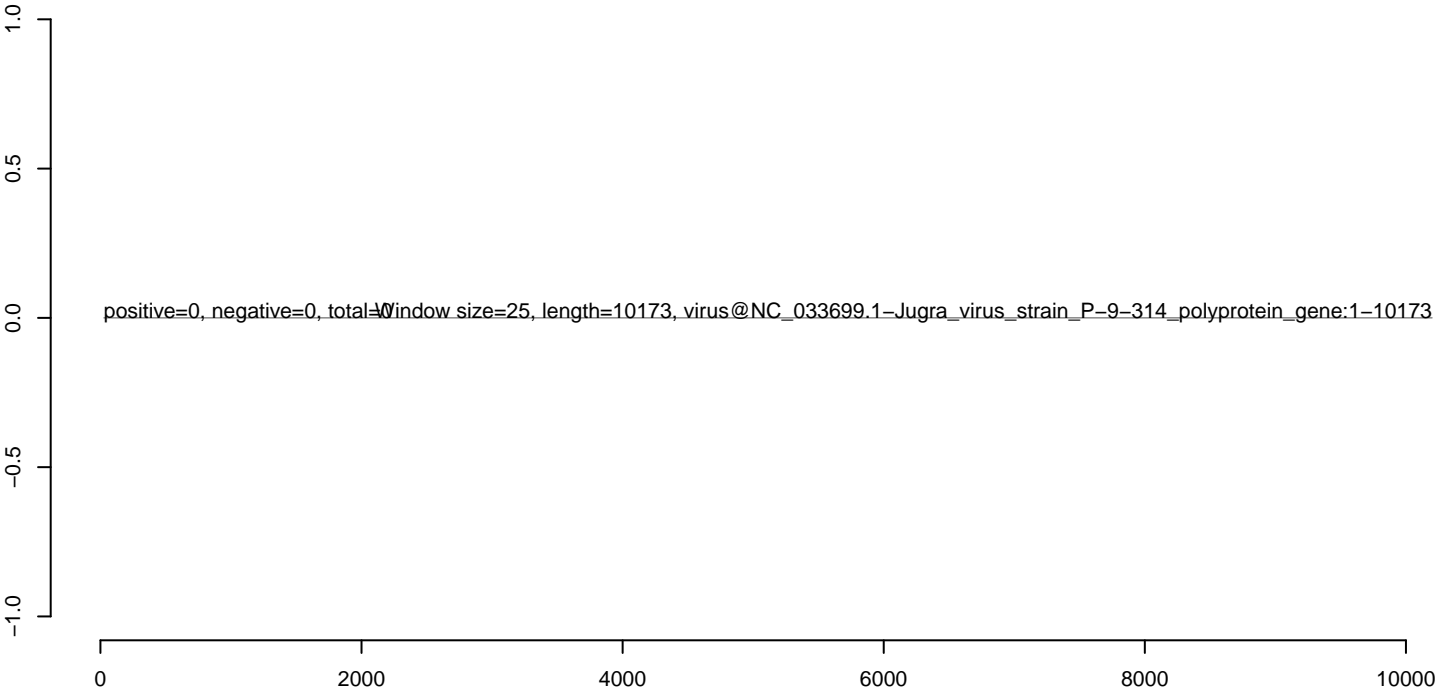
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AeAlbo_C636_B2_FHV_r54q.24_35.rep



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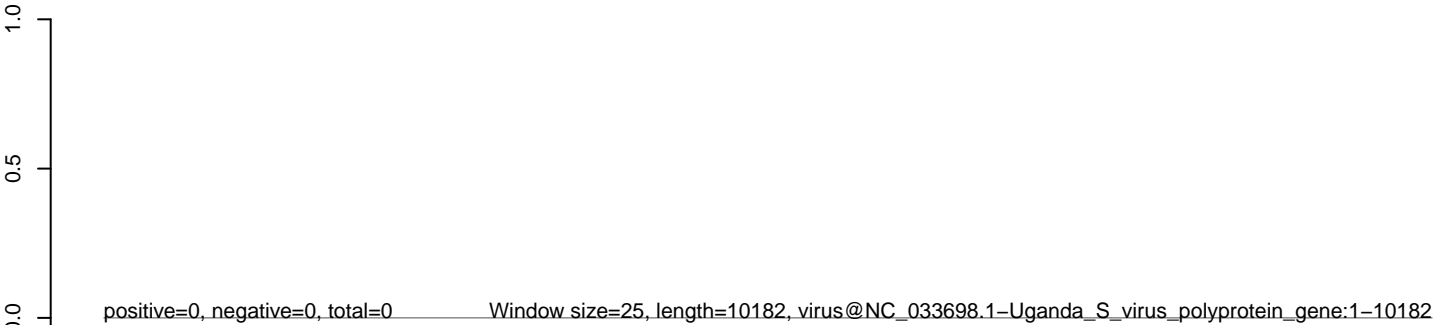
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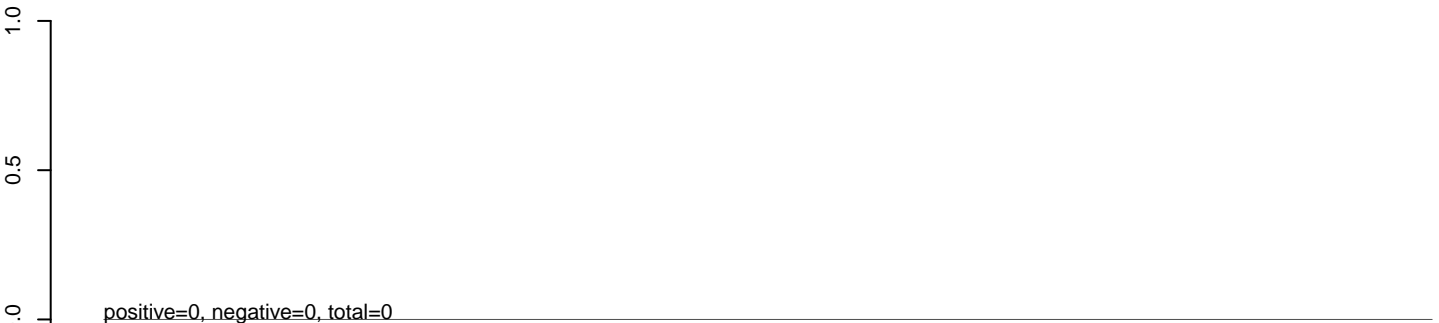


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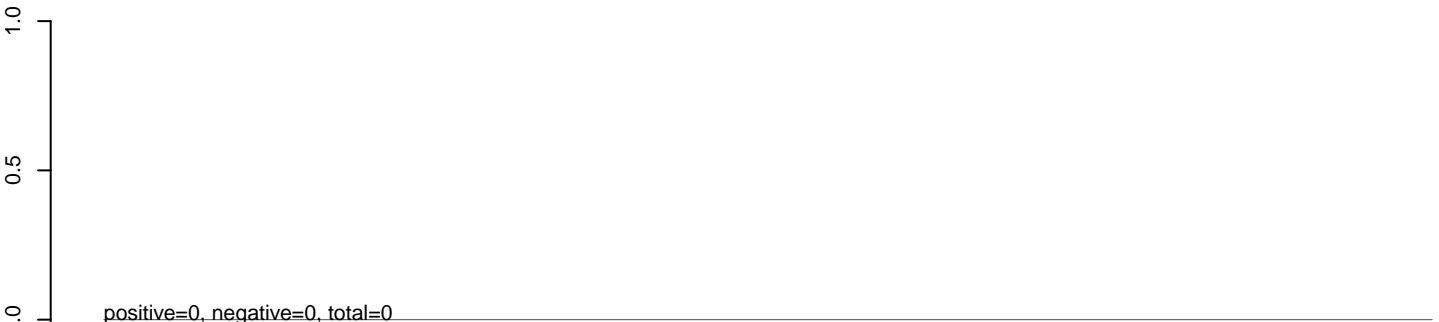


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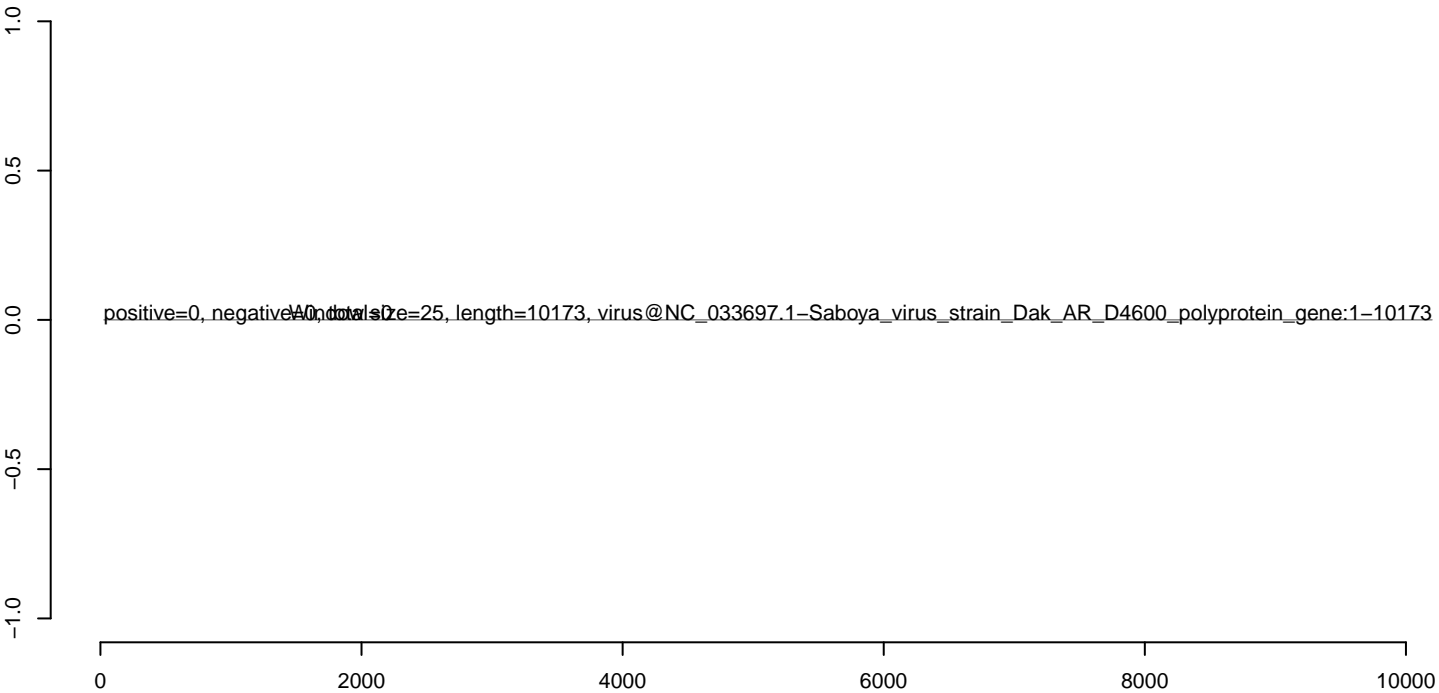
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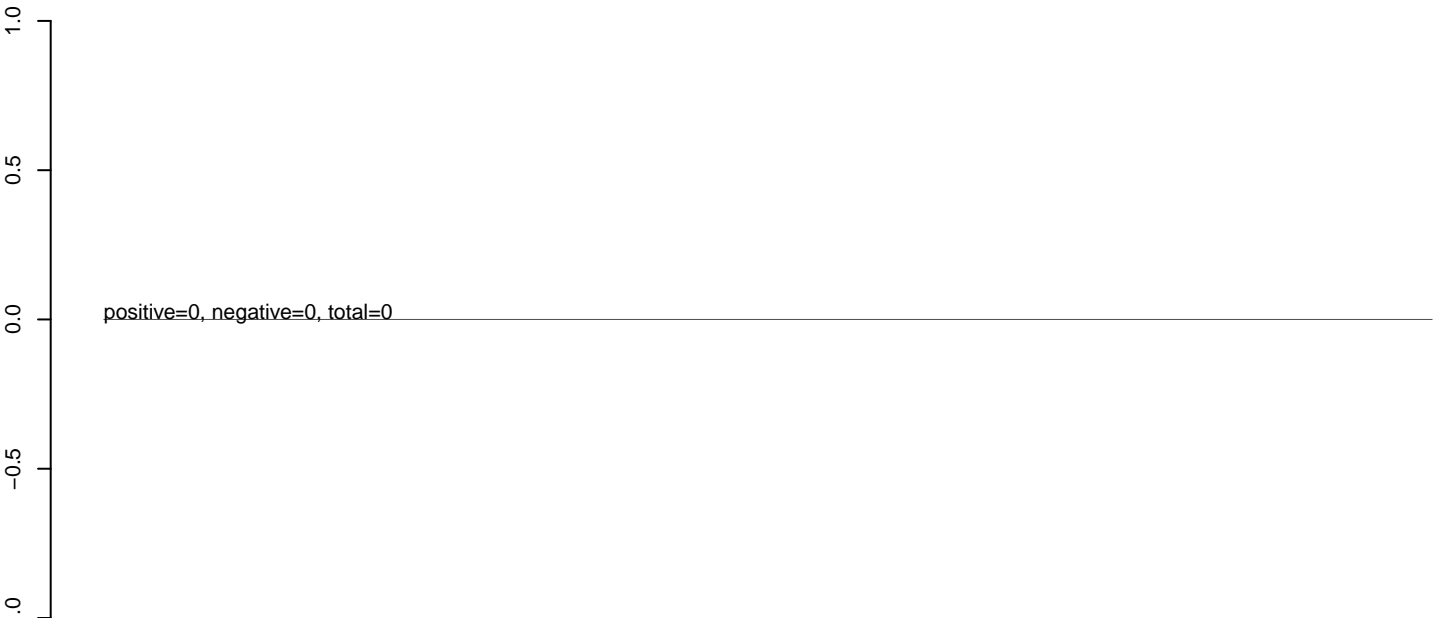
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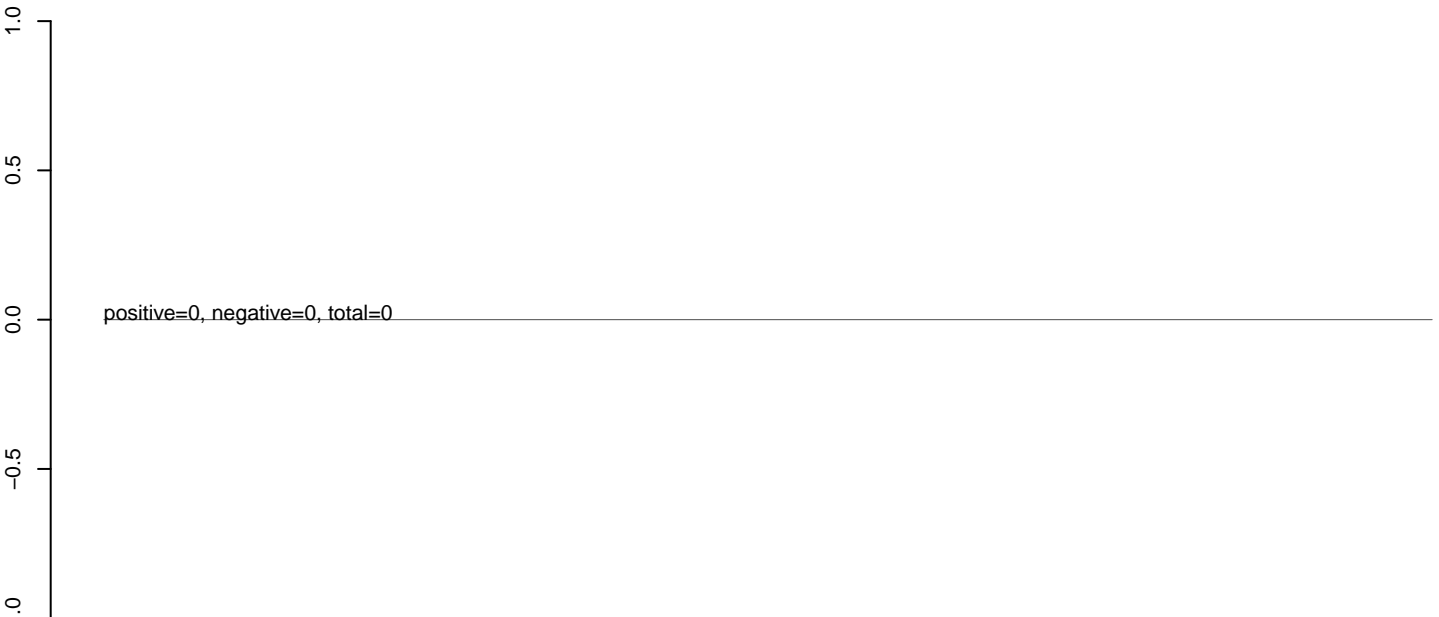
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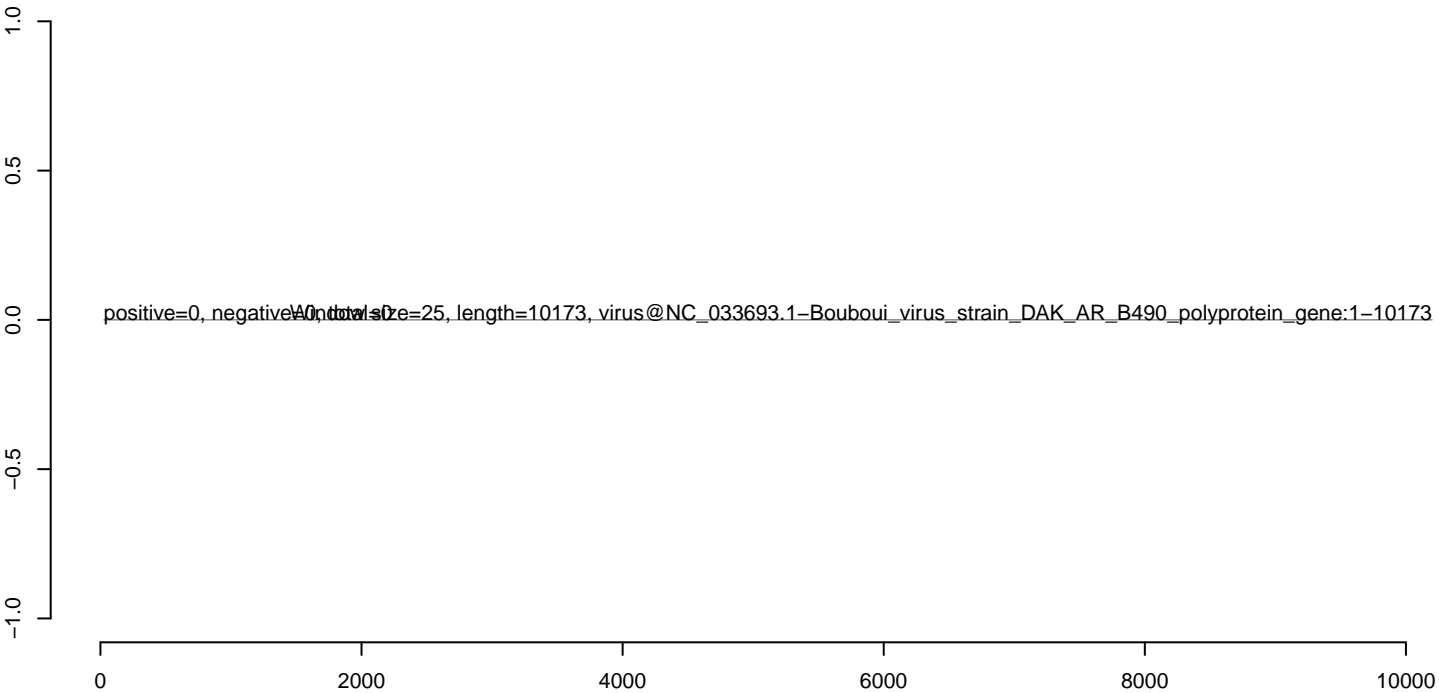
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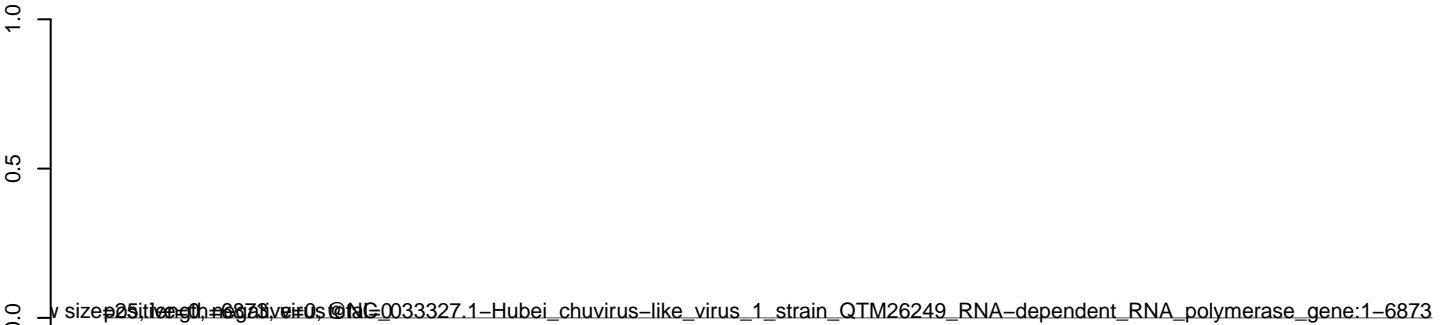
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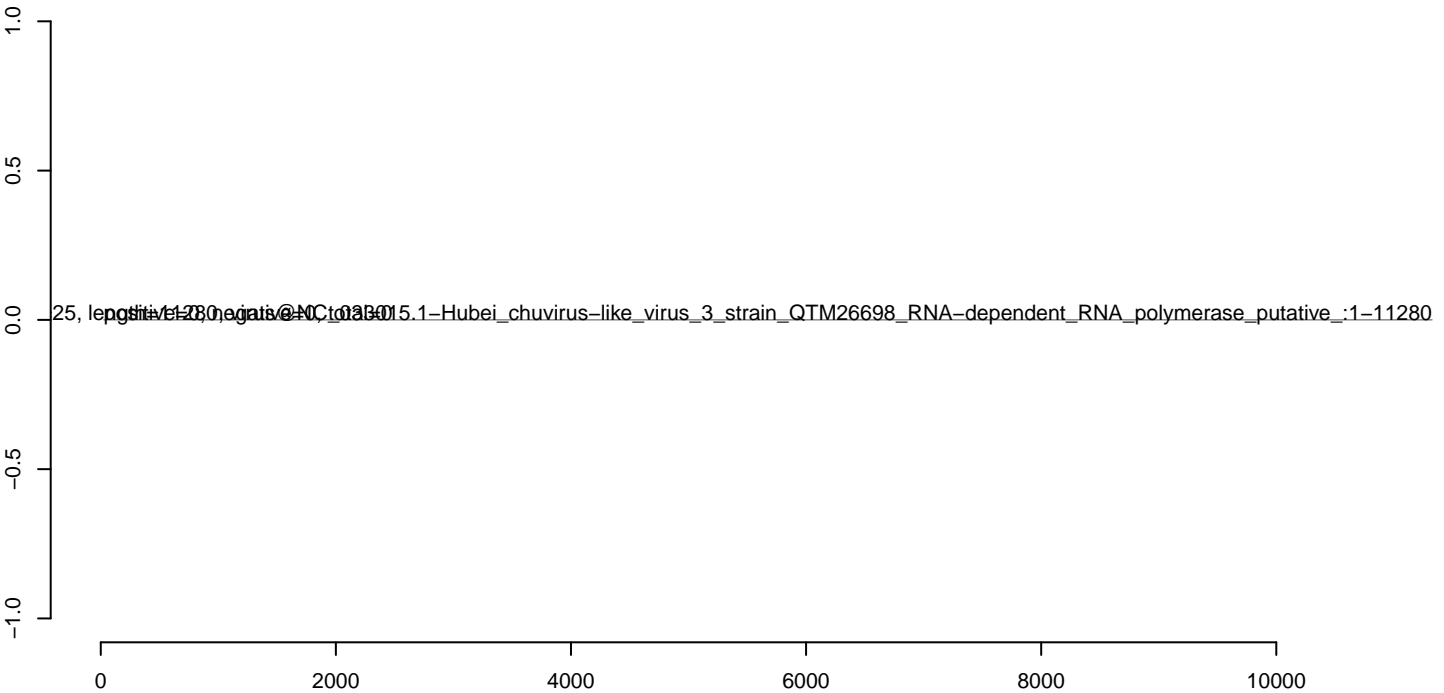
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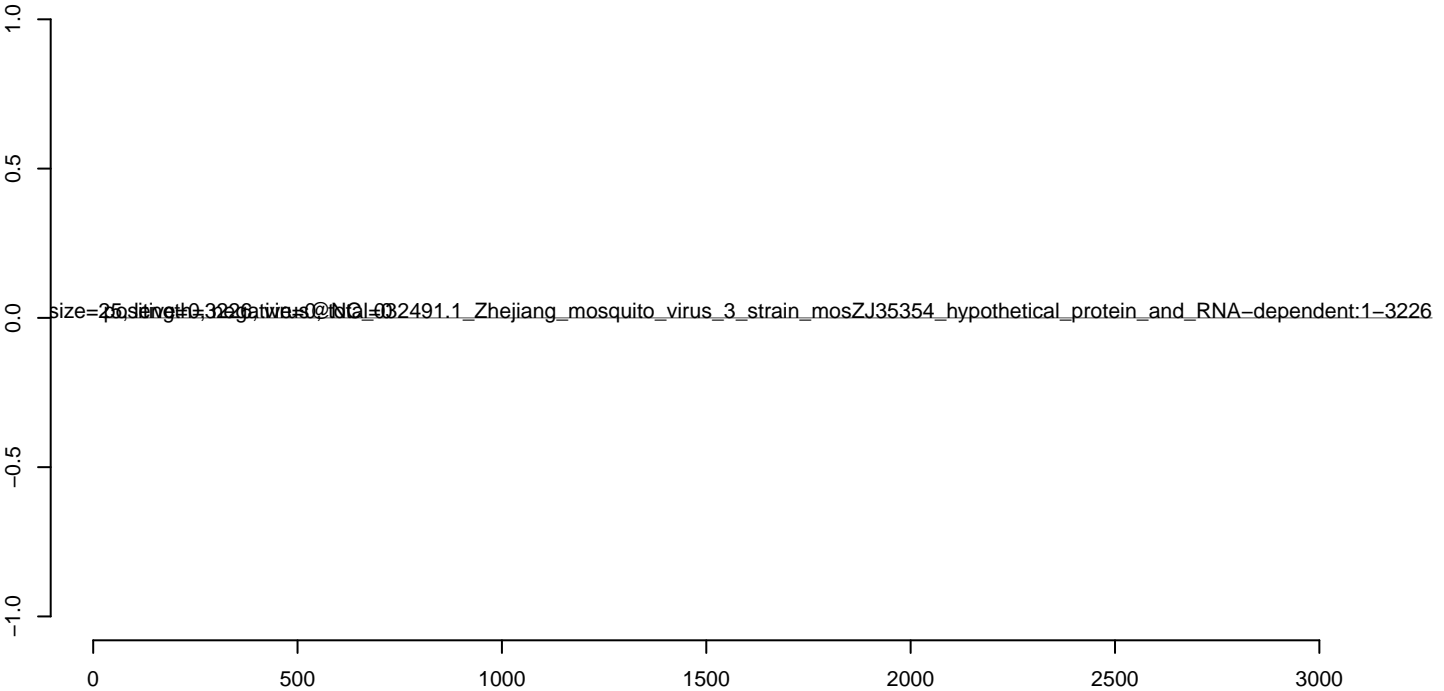
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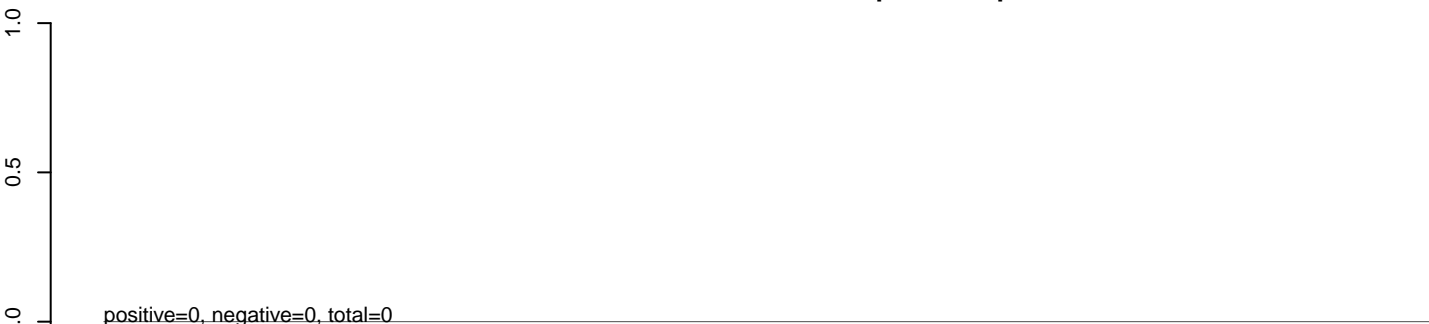
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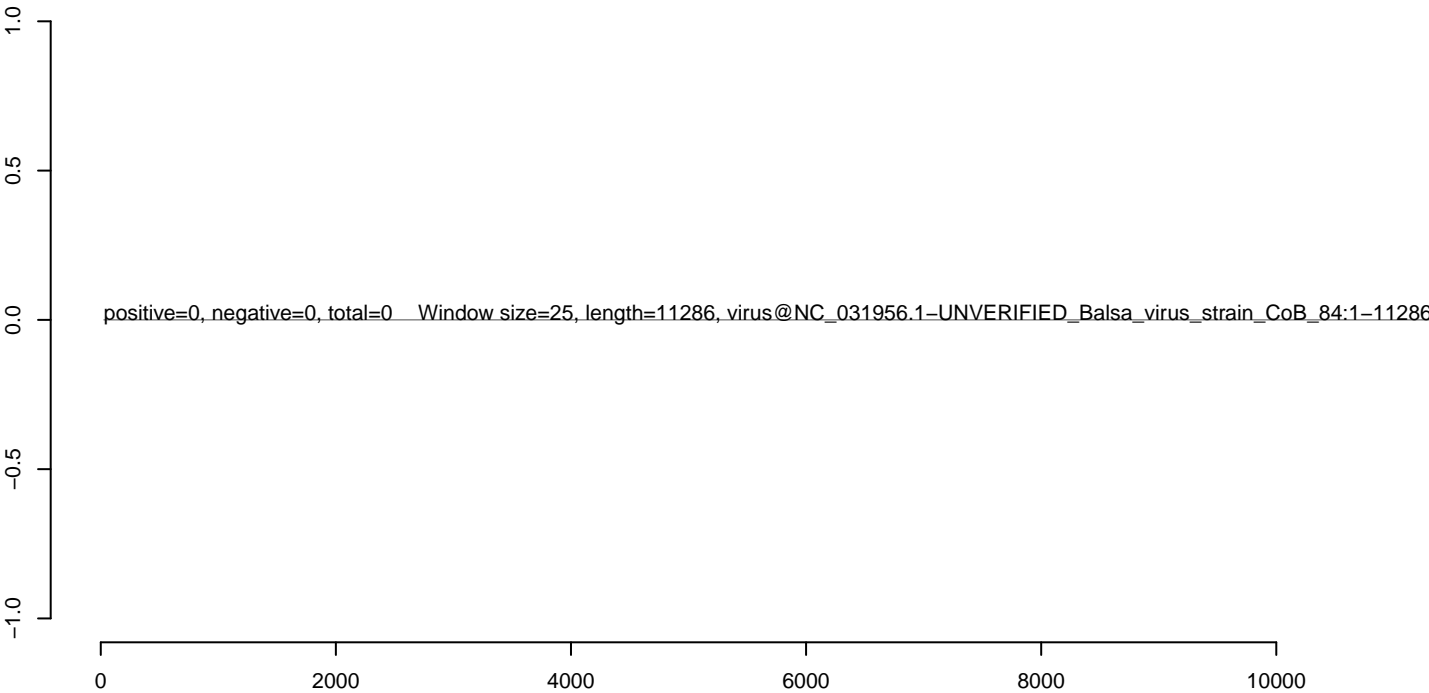
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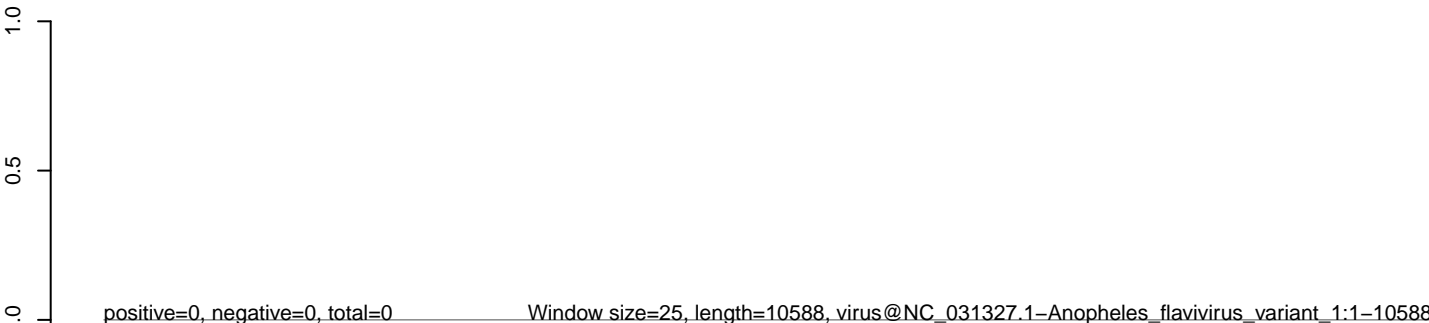
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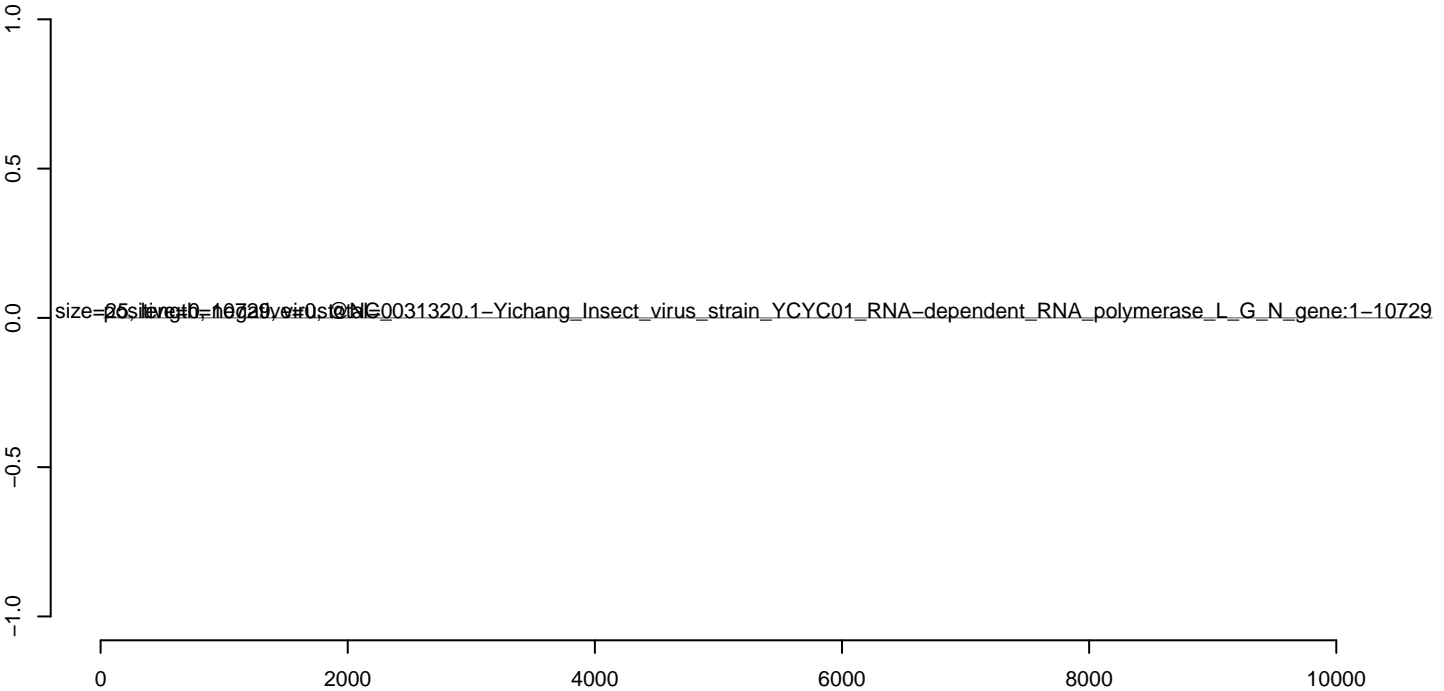
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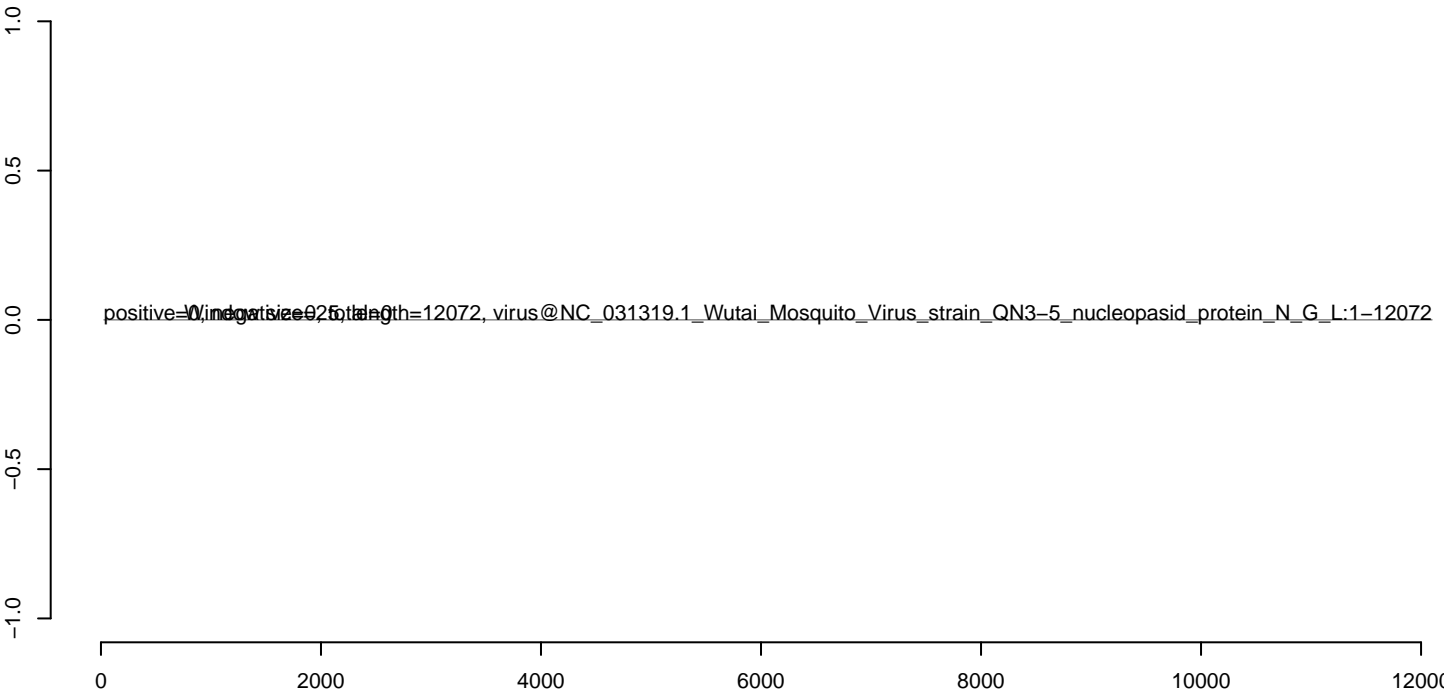
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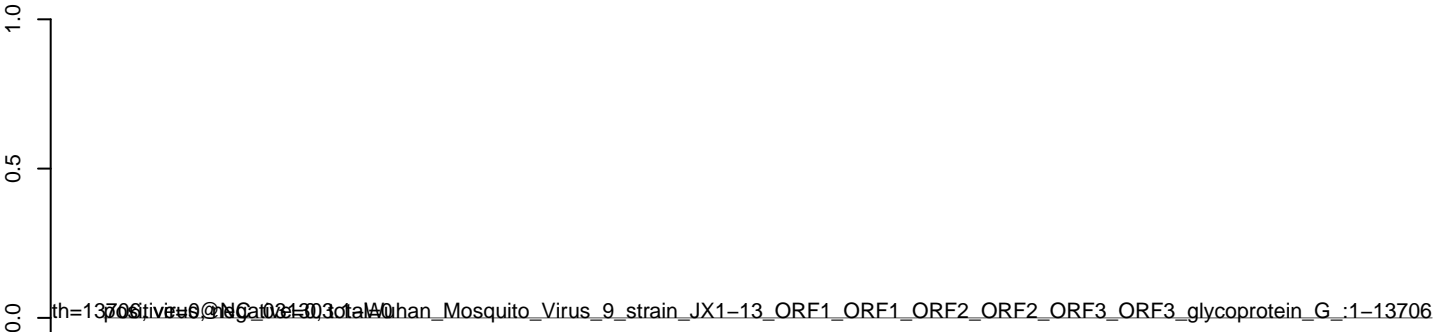
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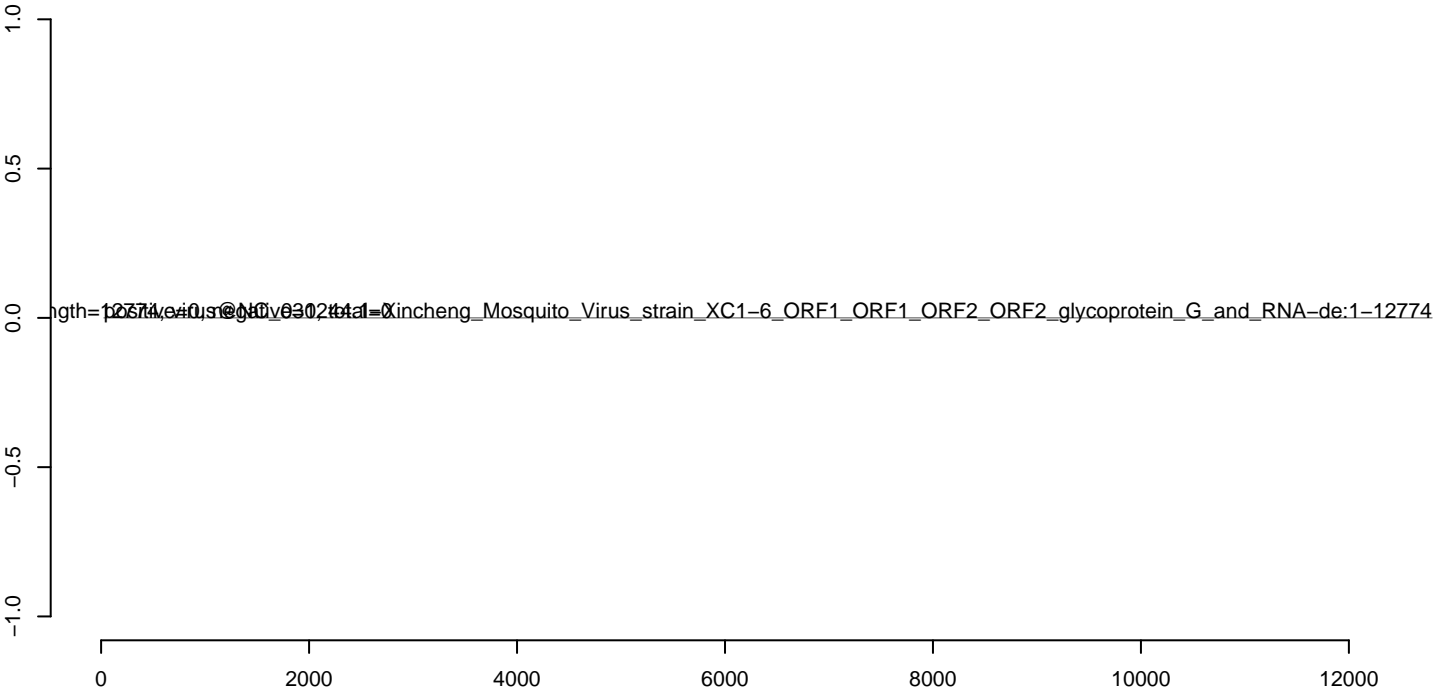
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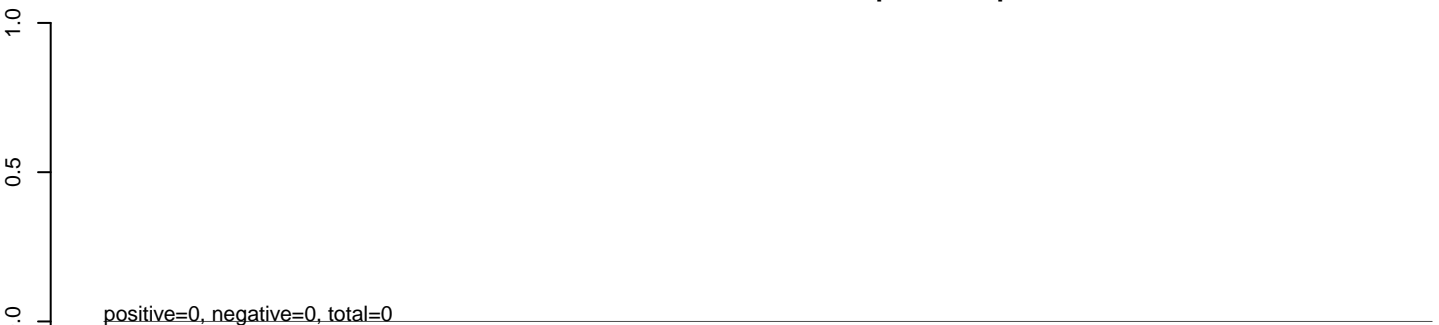
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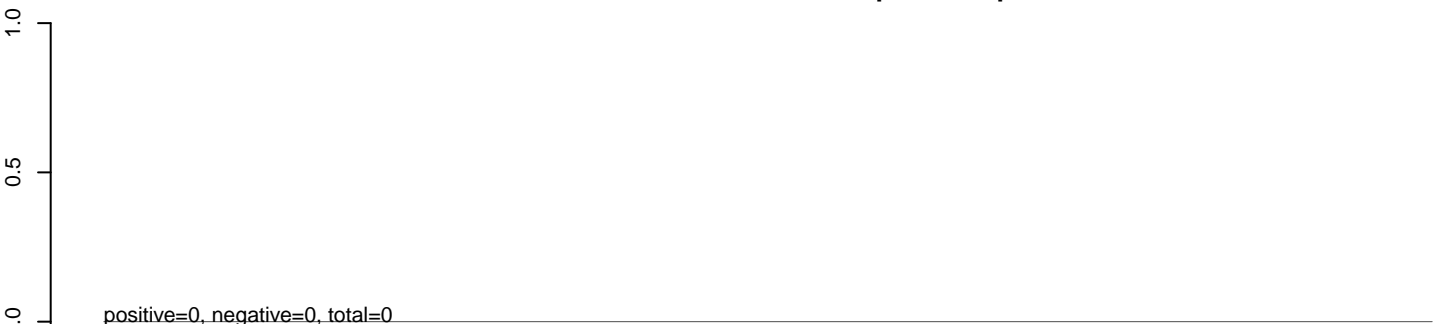
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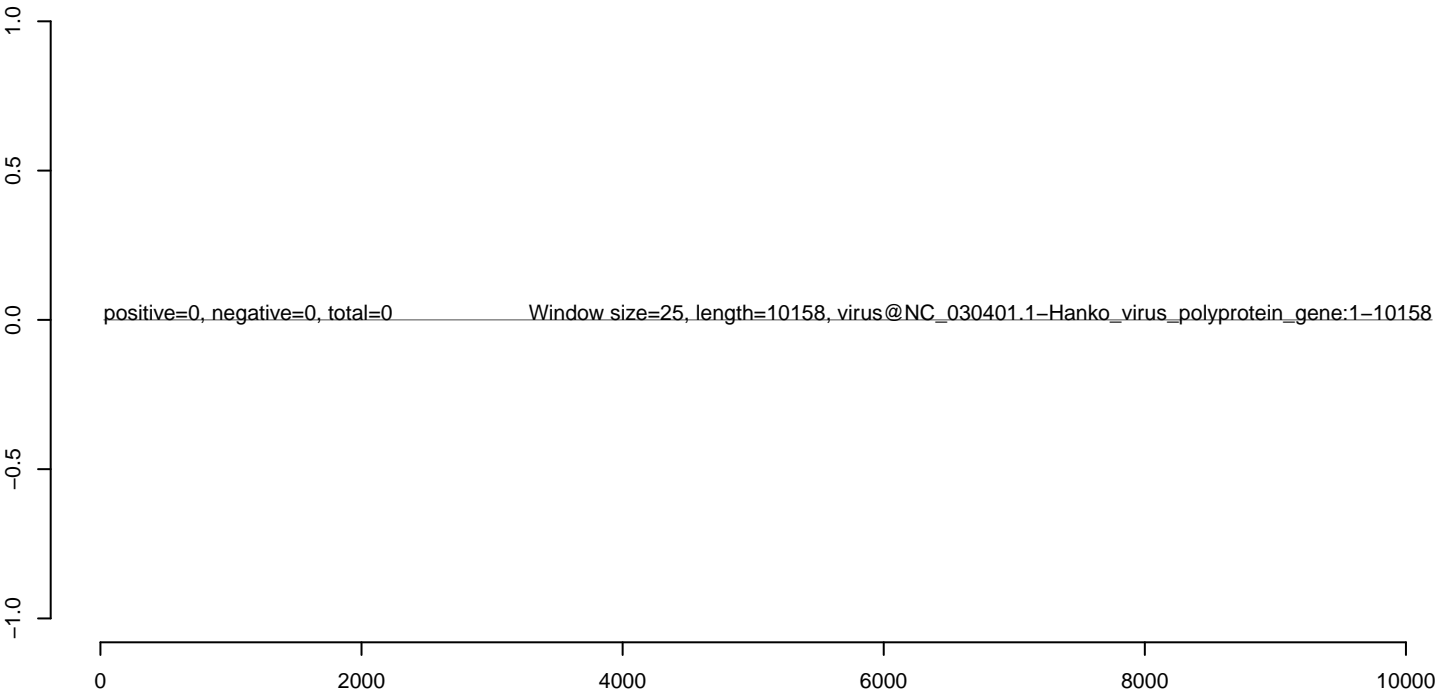
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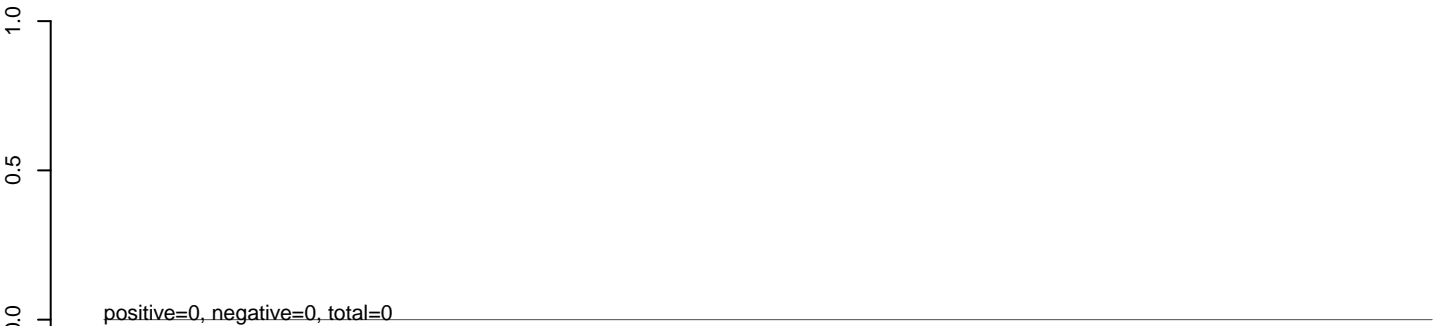
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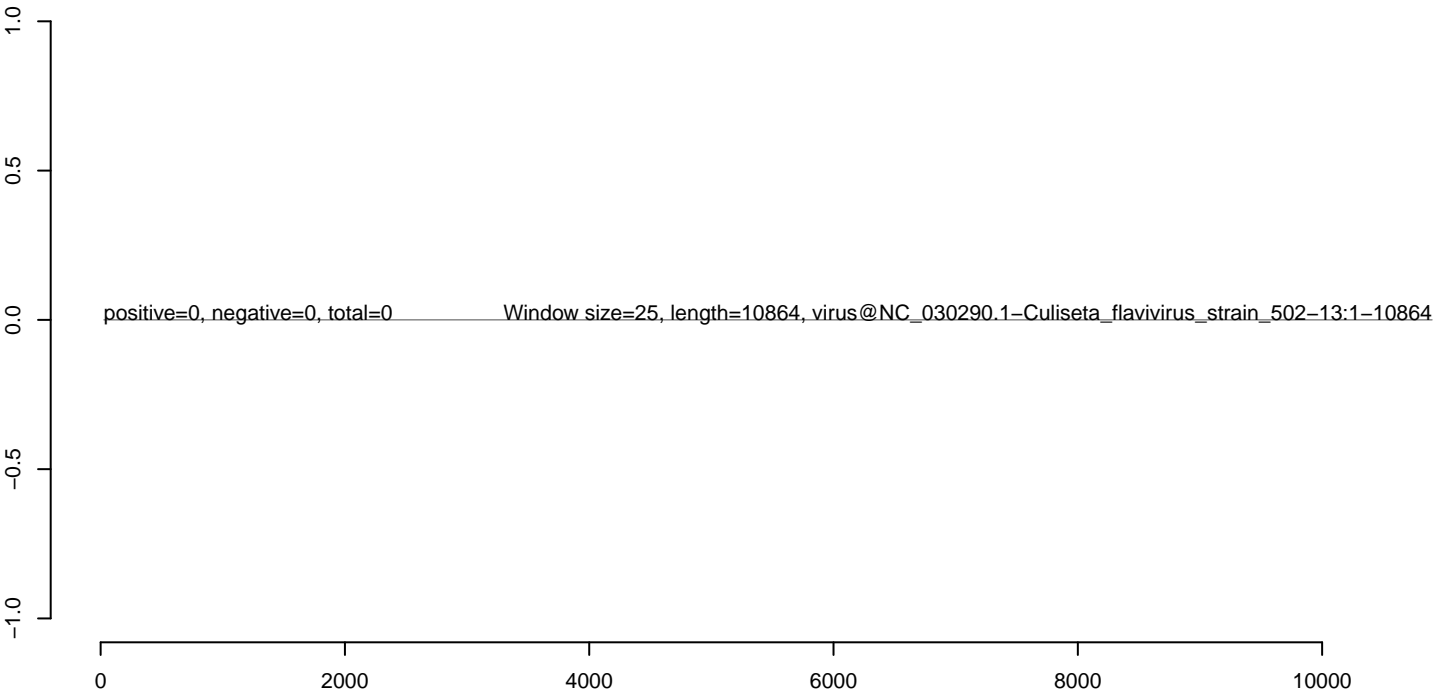
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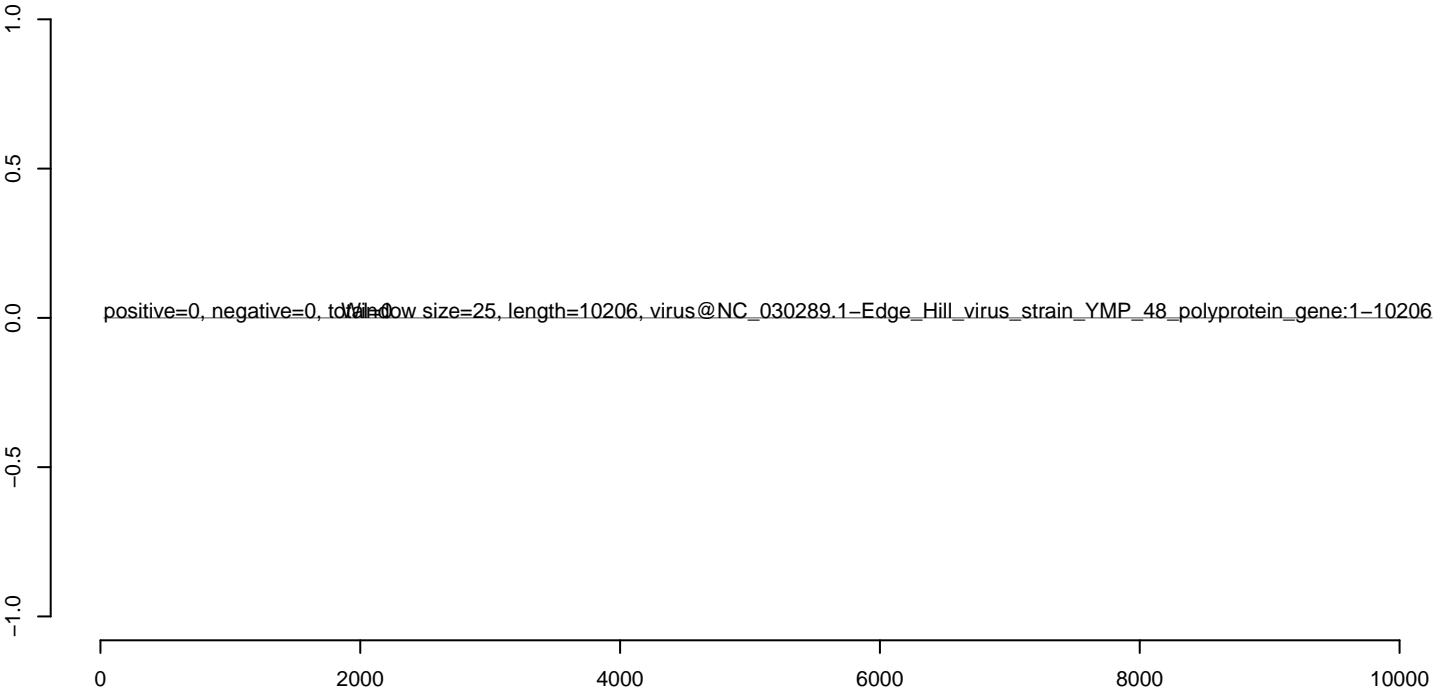
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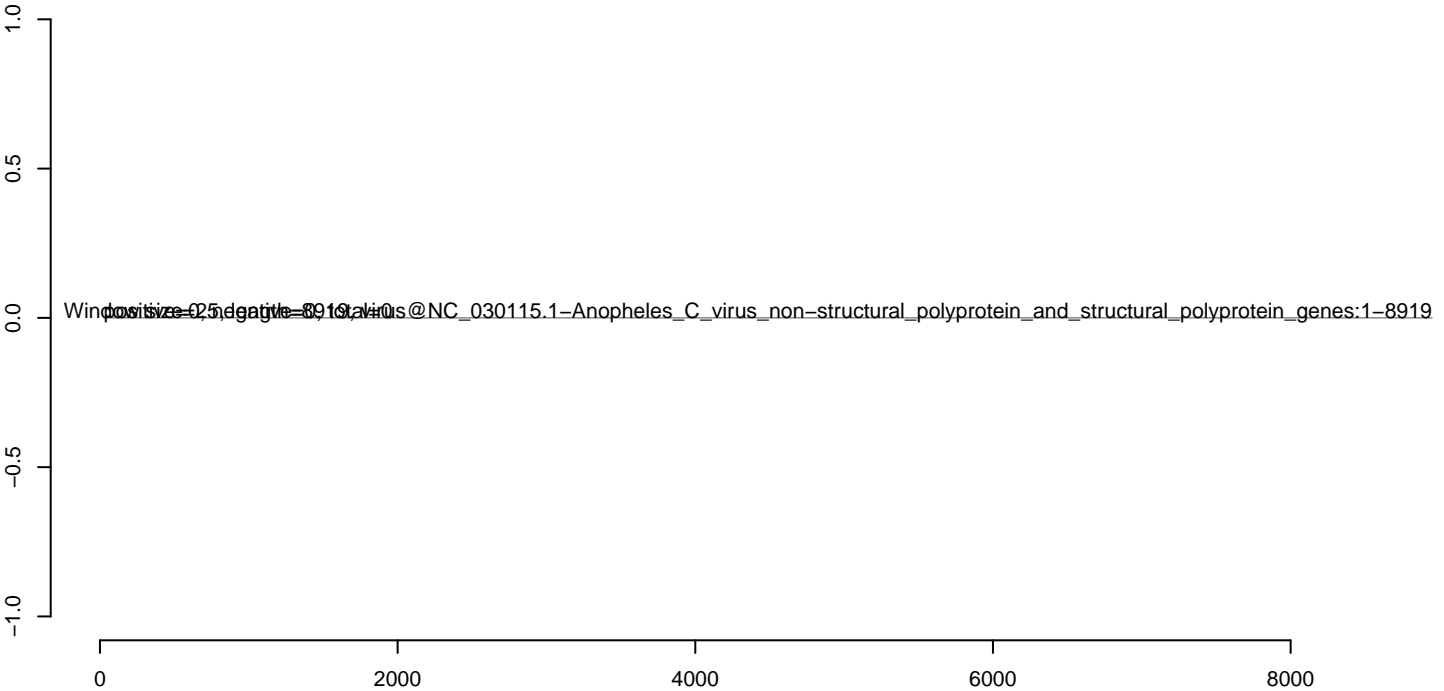
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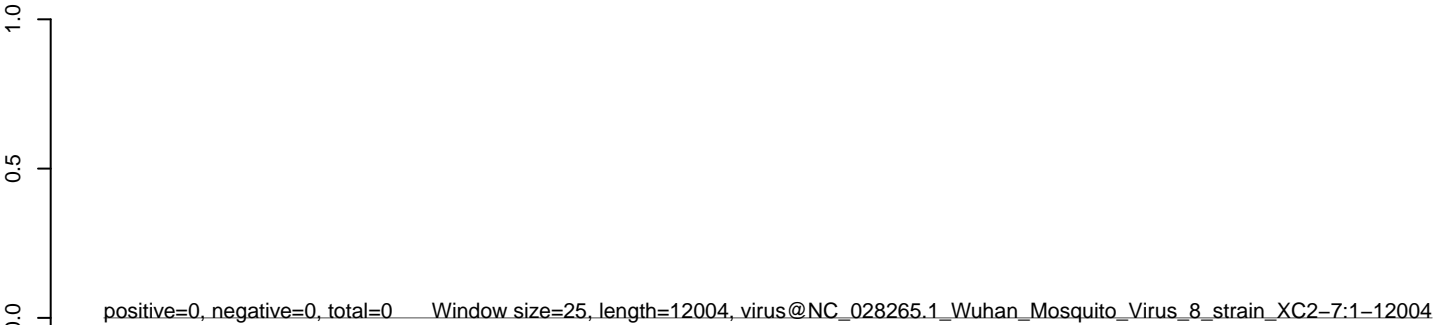
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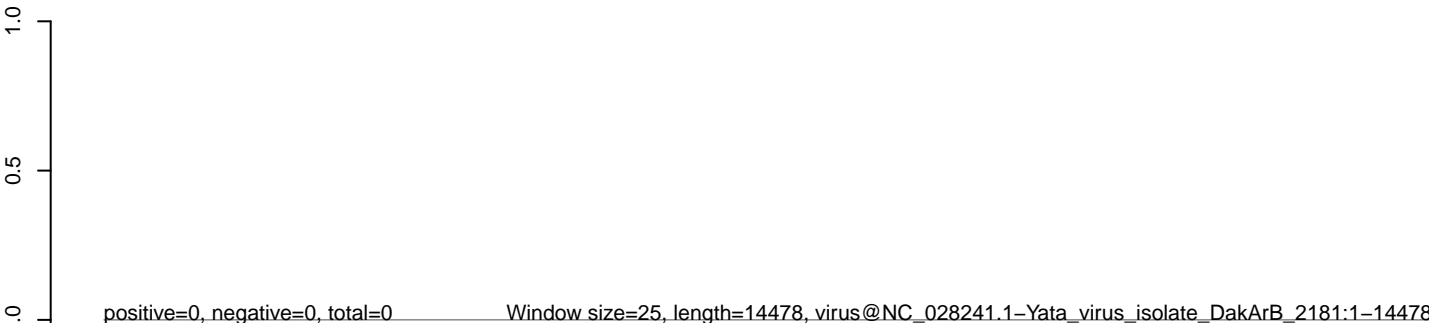
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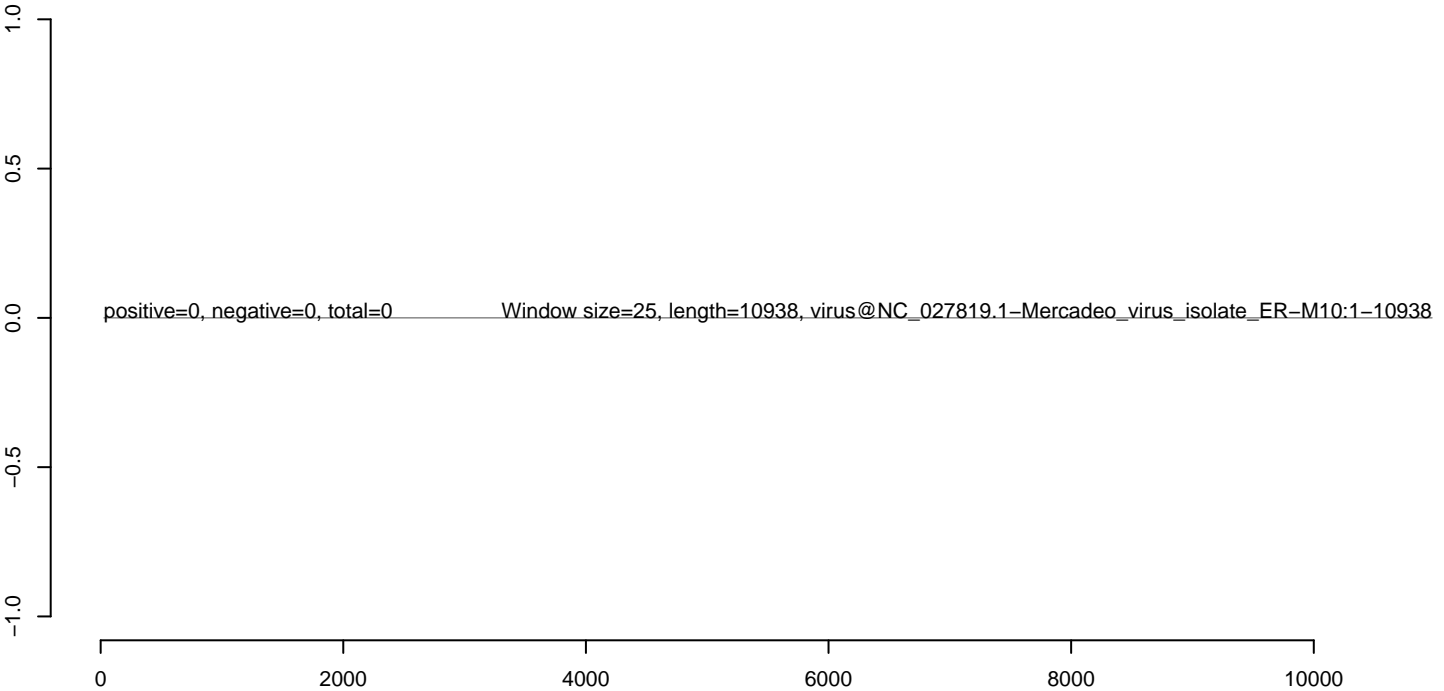
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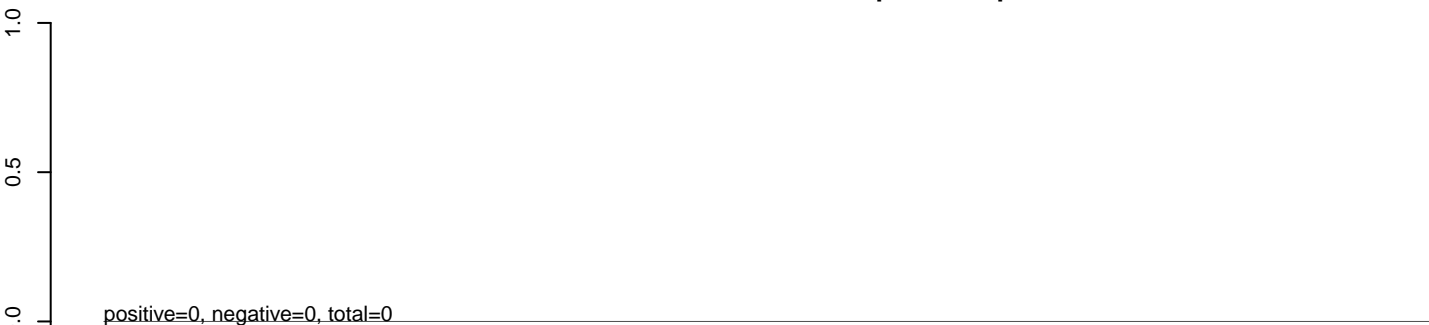
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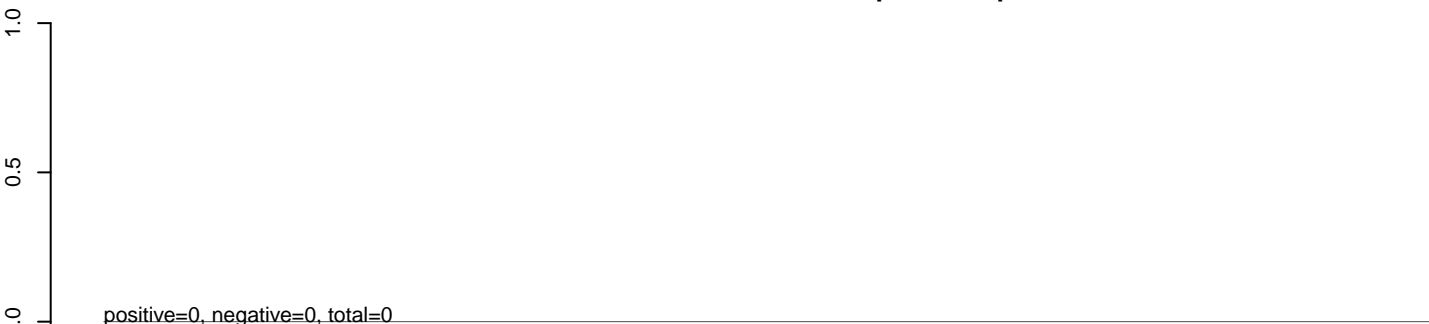
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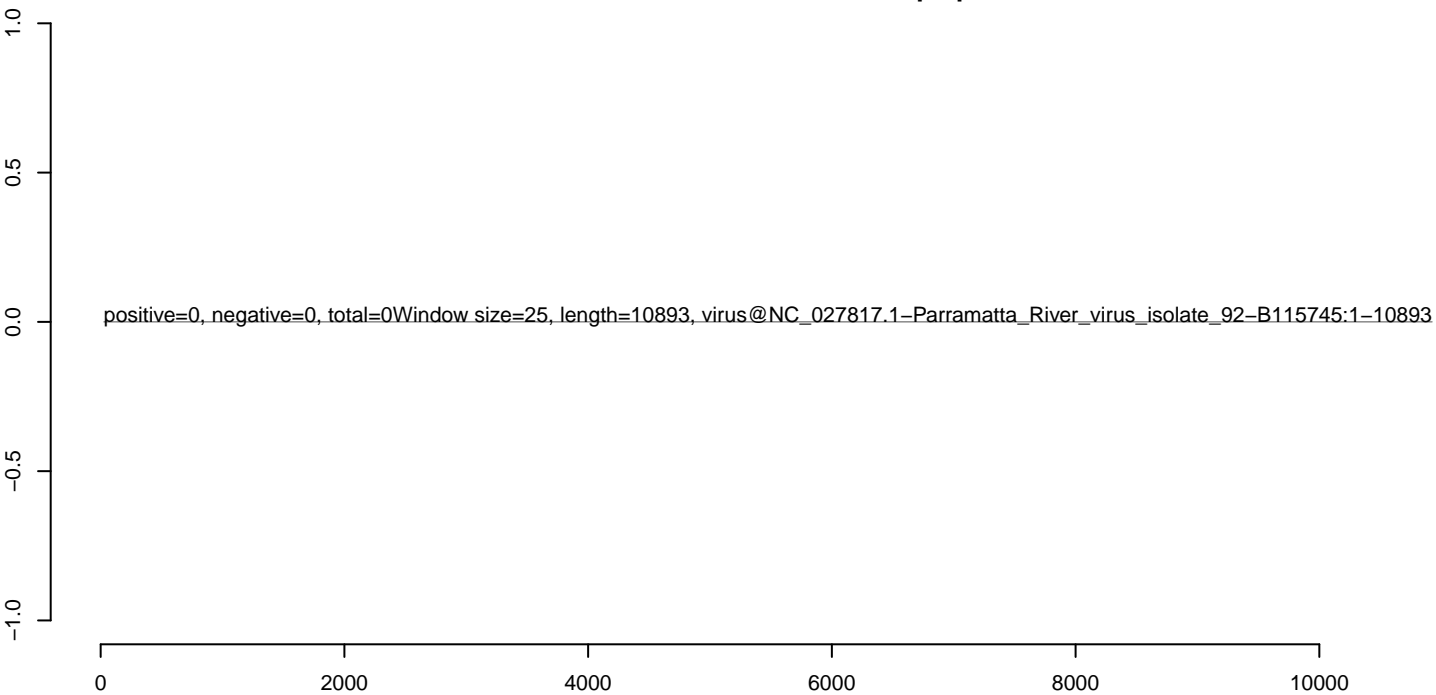
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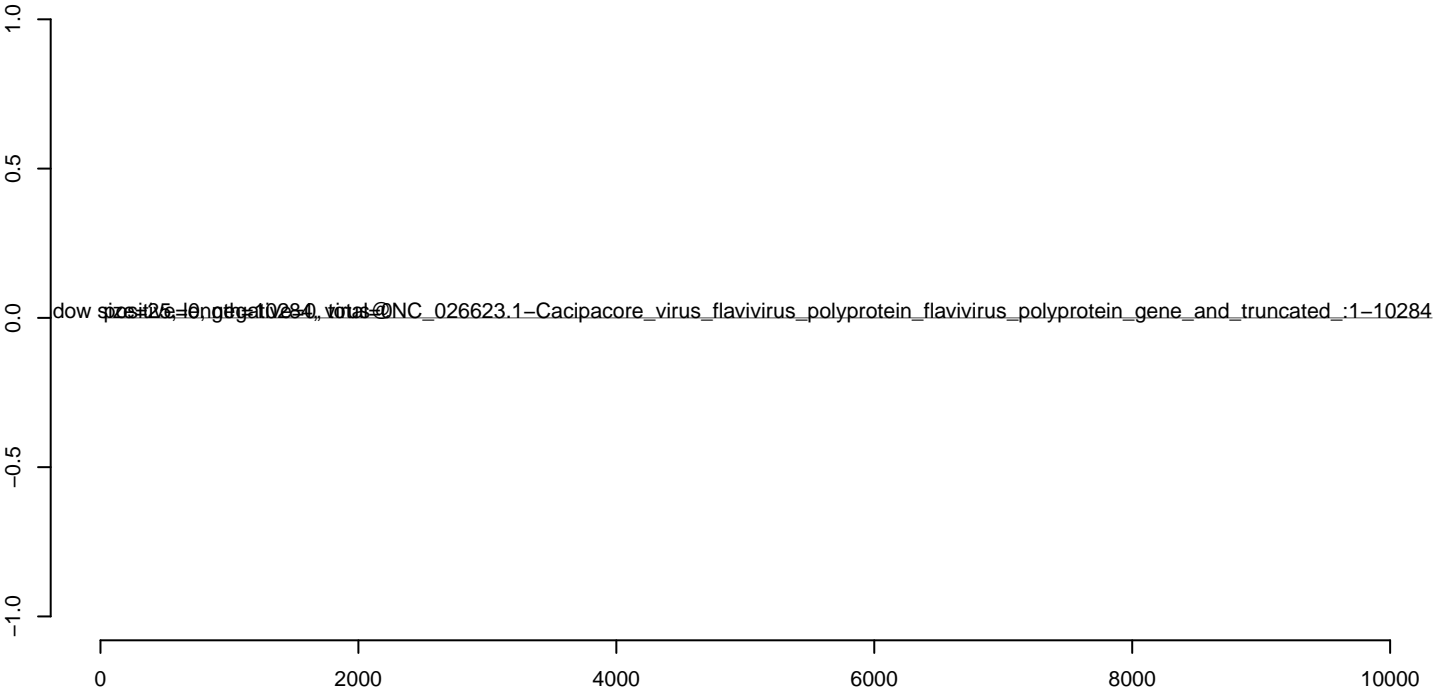
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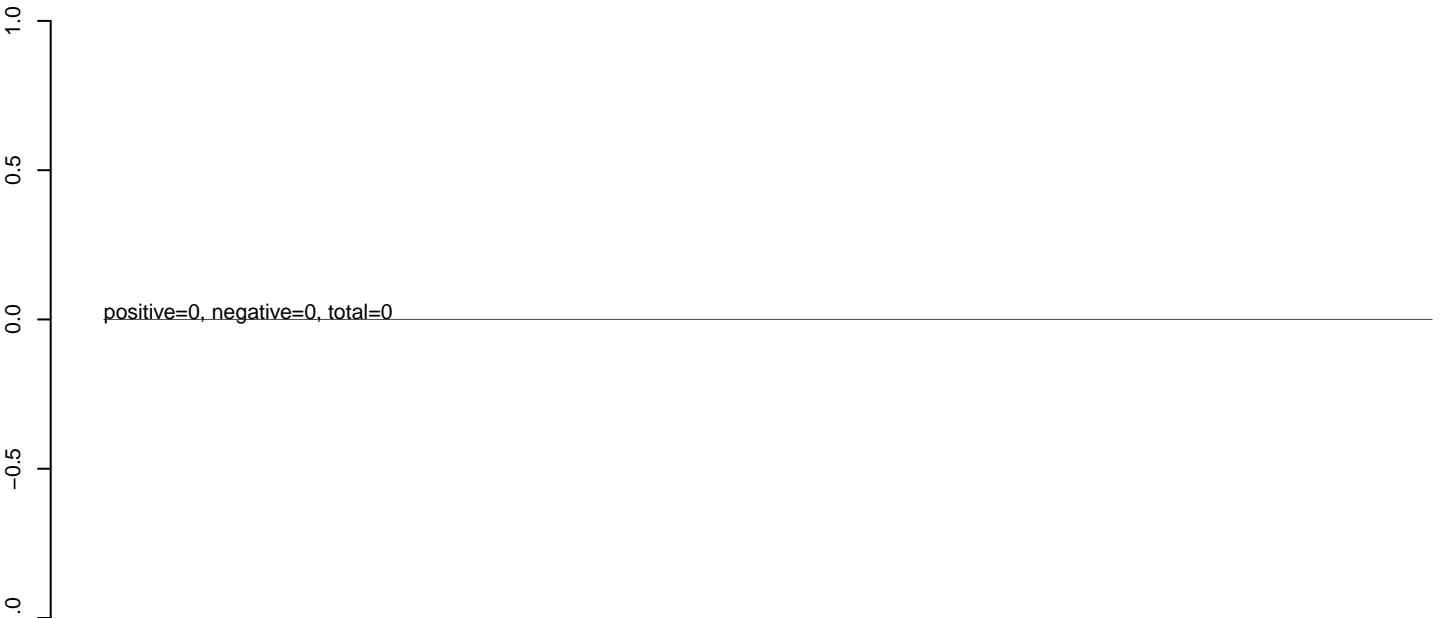
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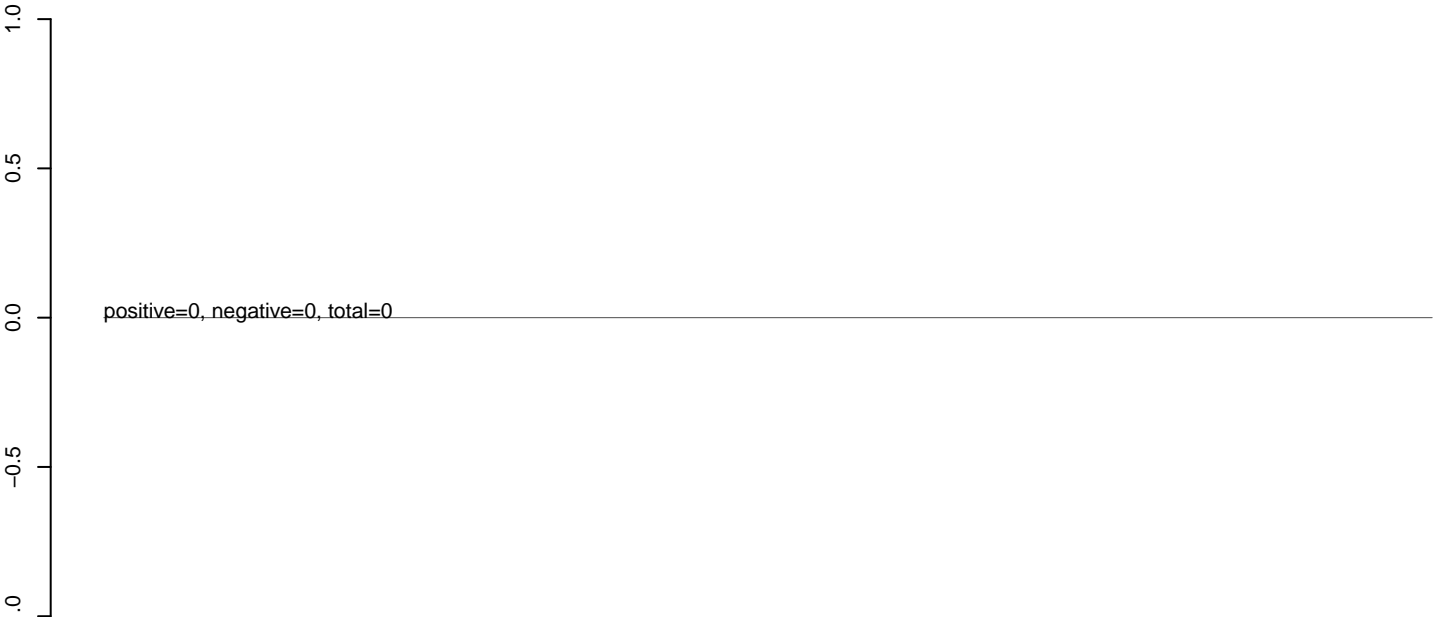
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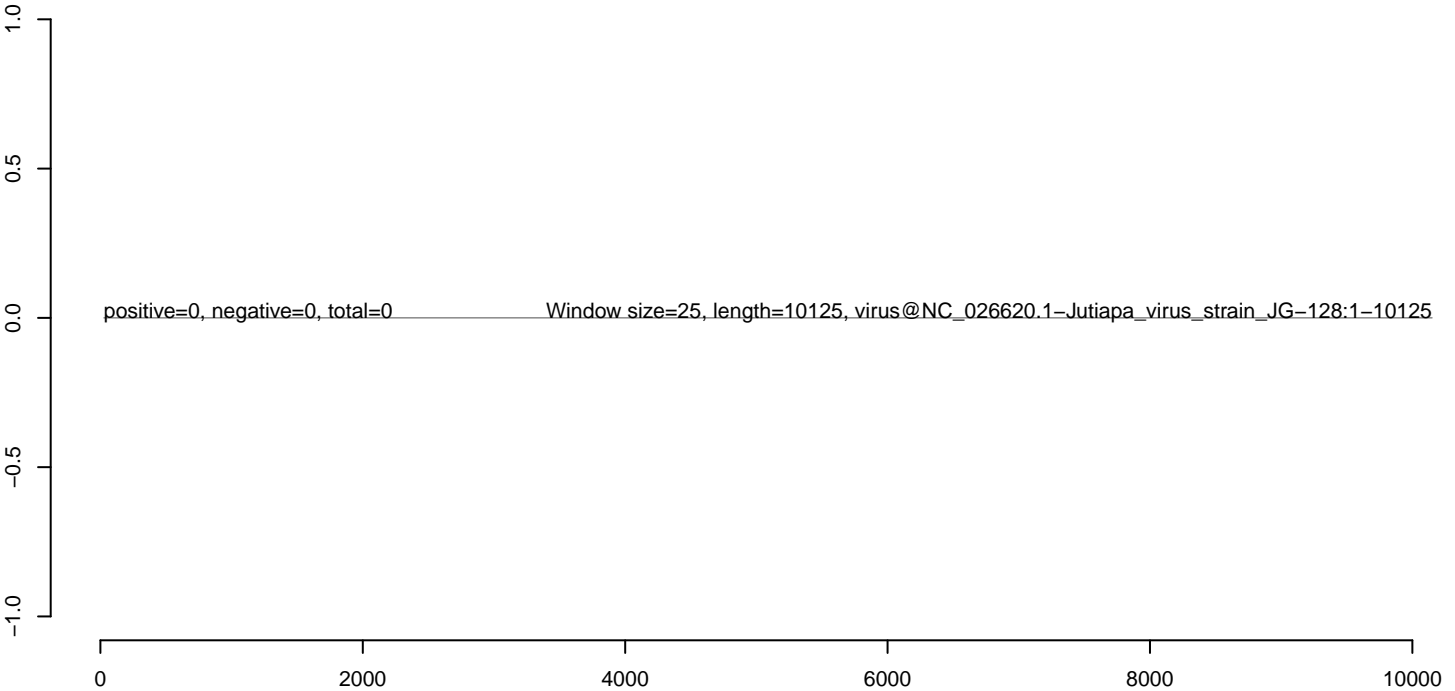
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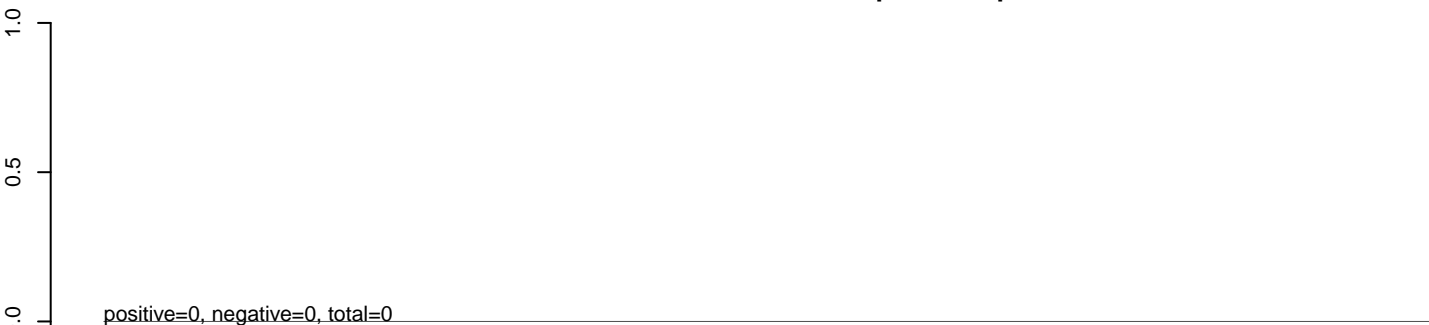
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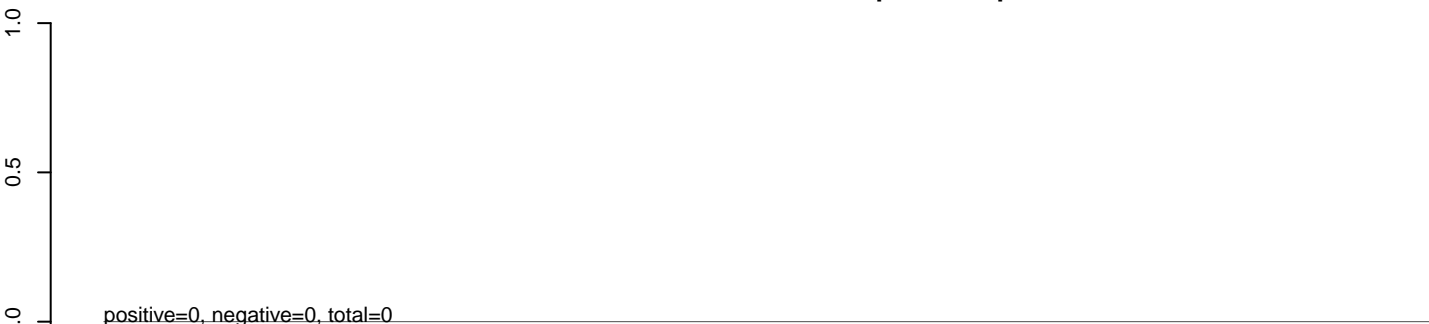
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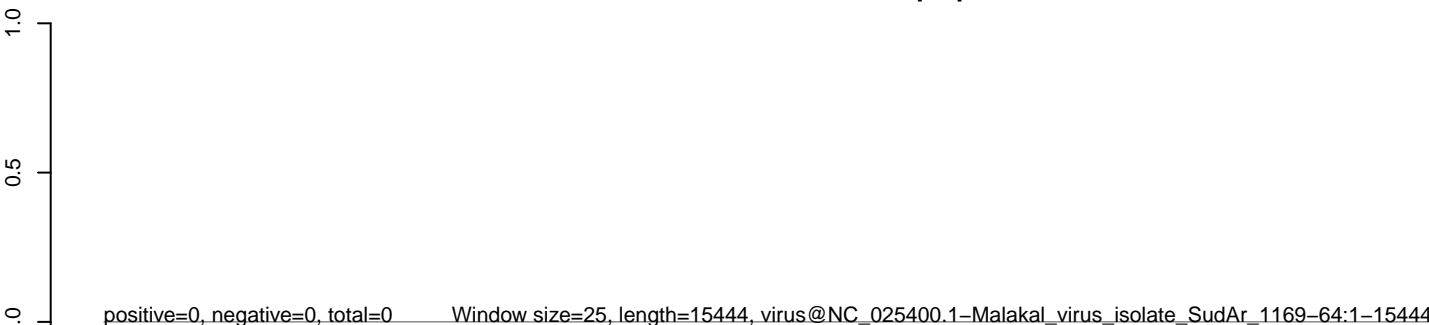
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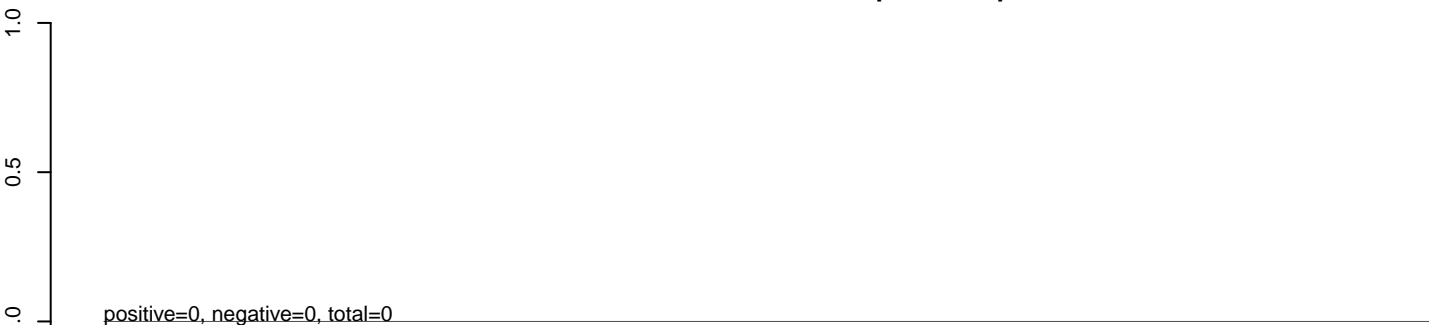
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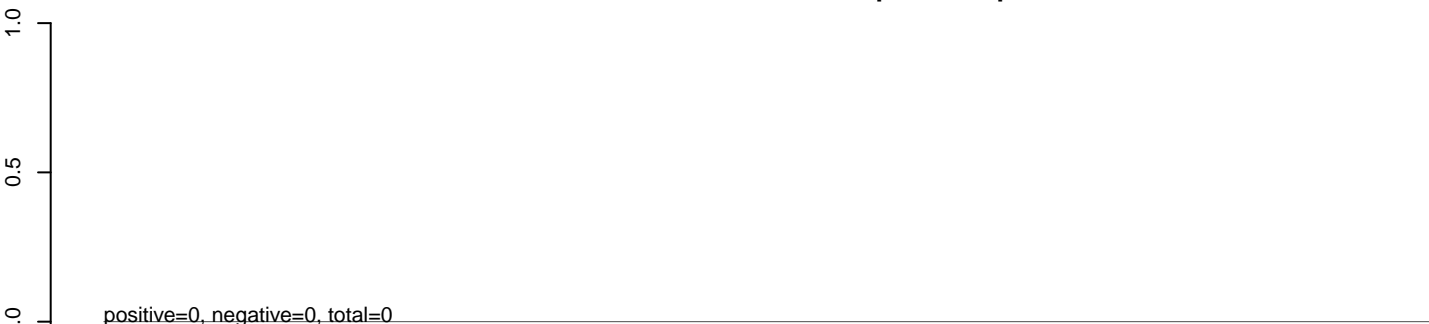
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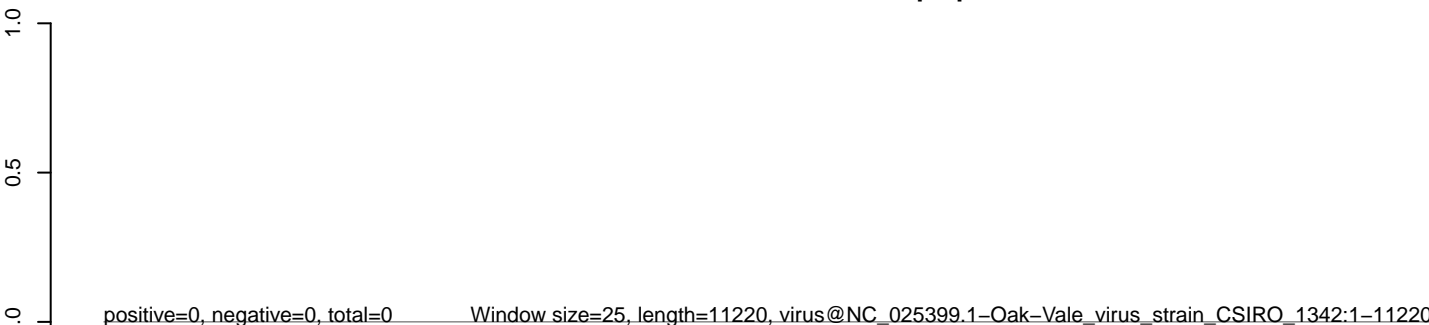
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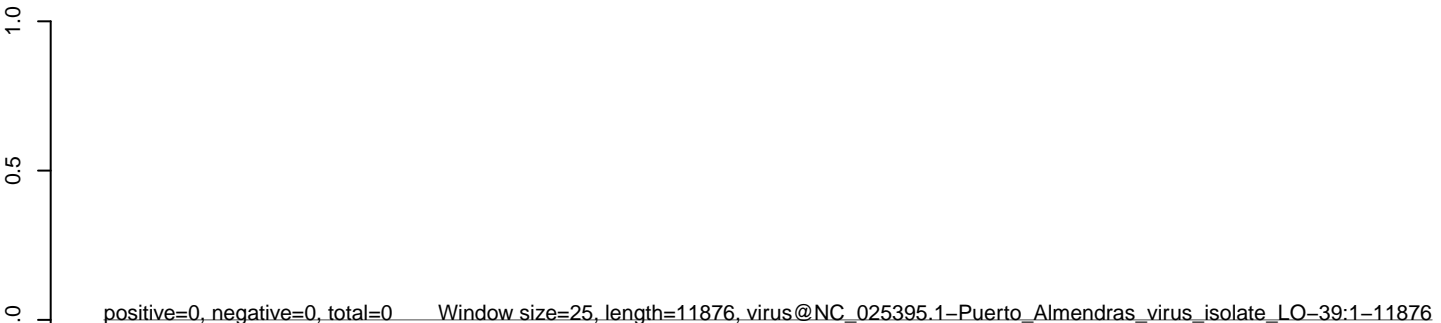
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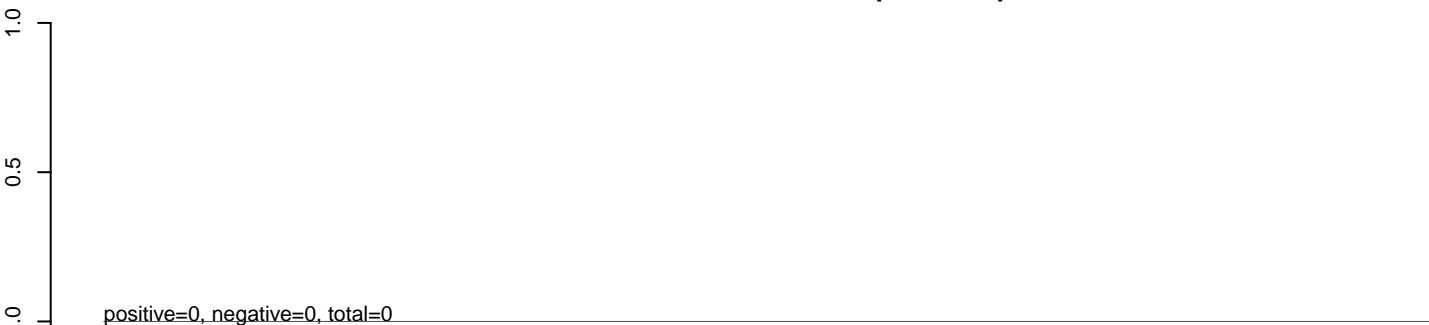
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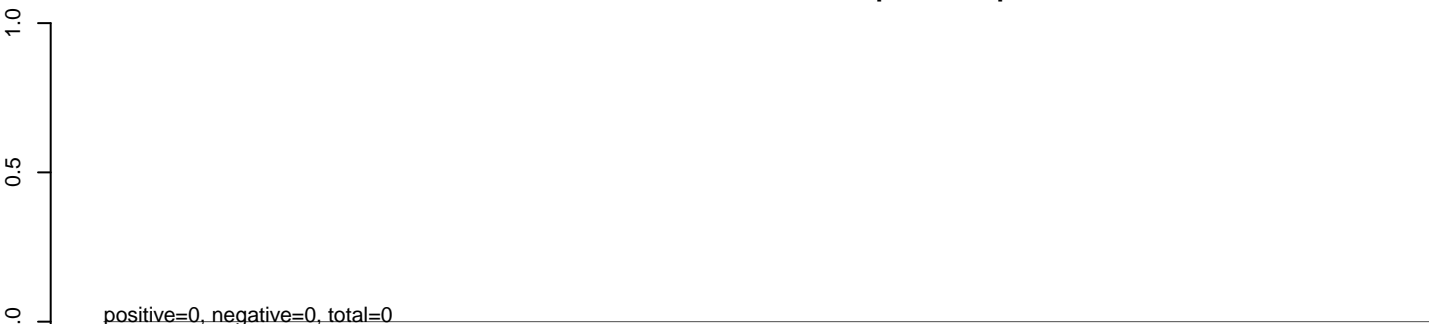
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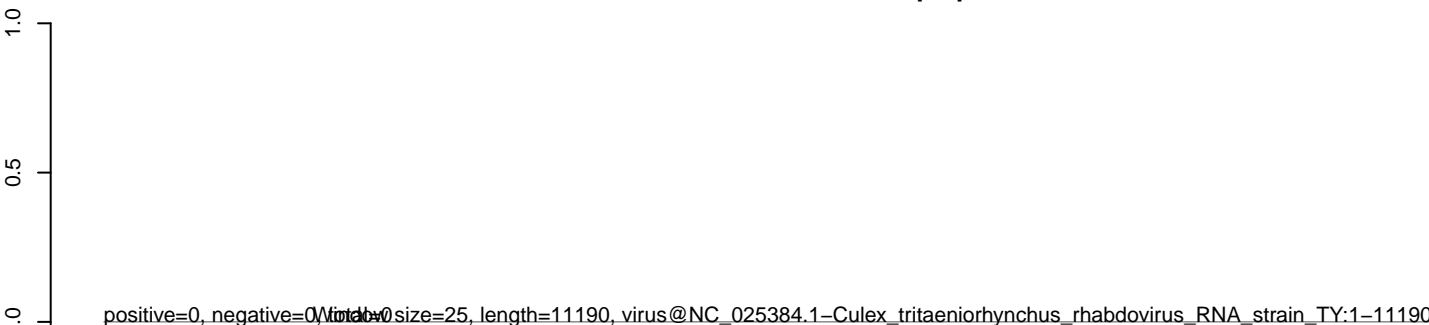
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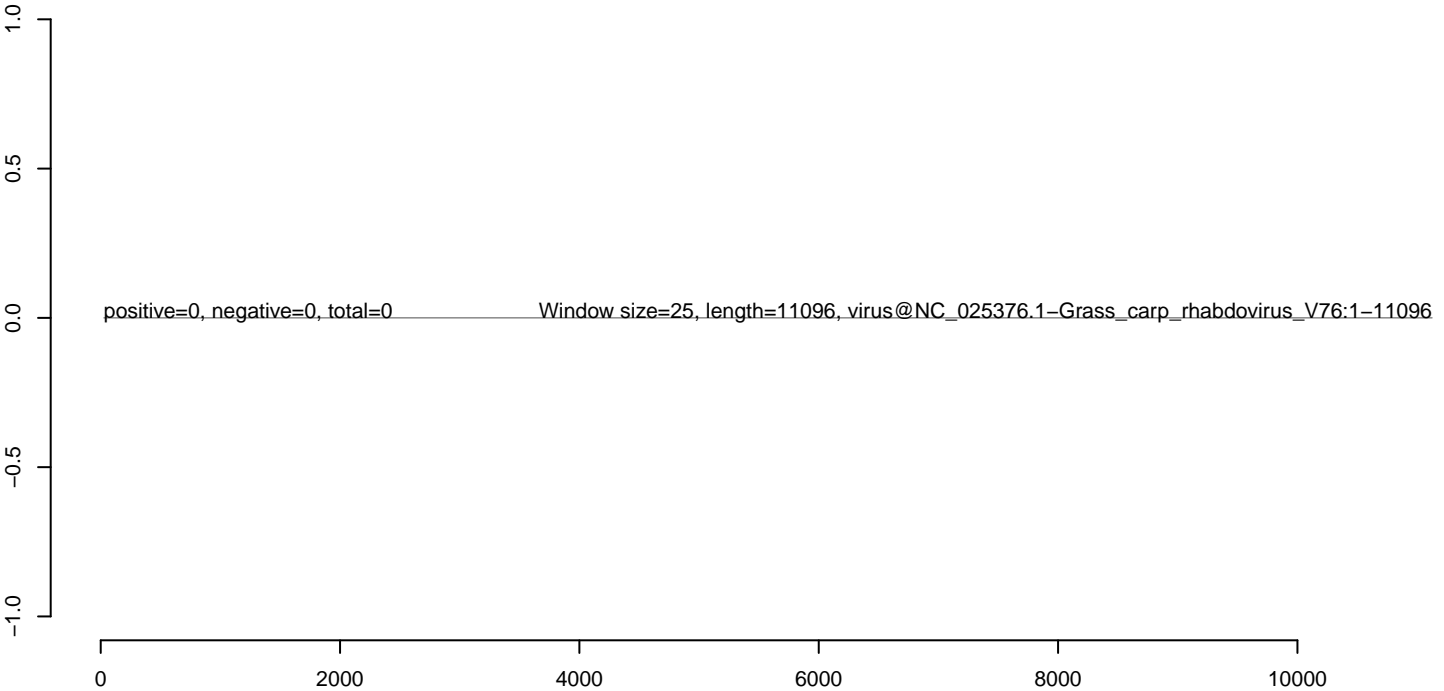
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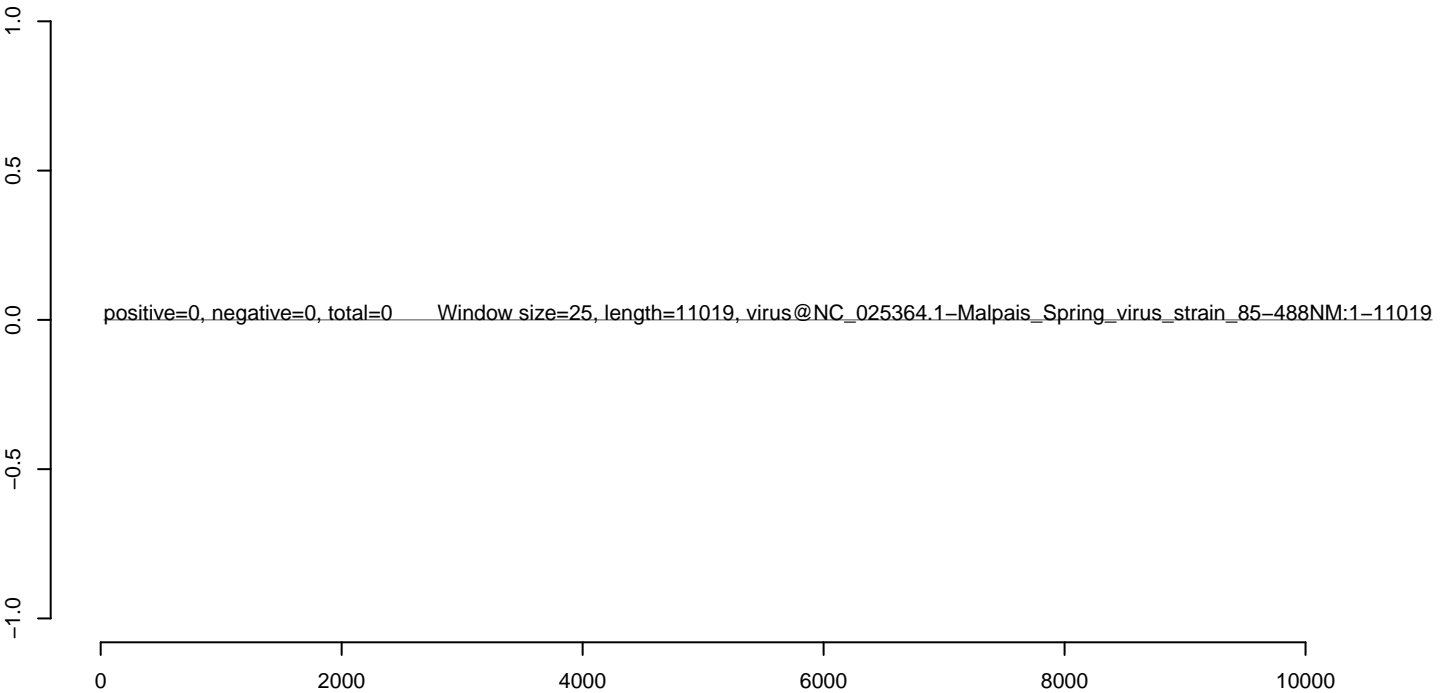
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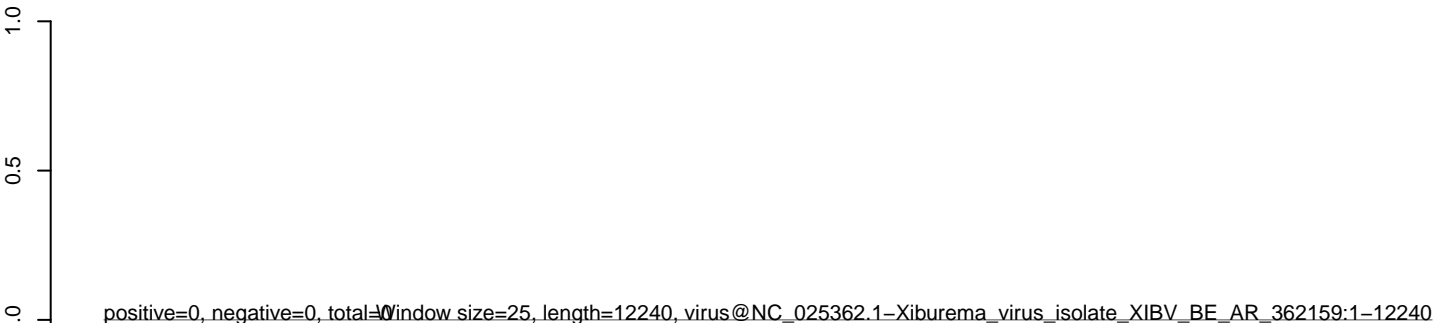
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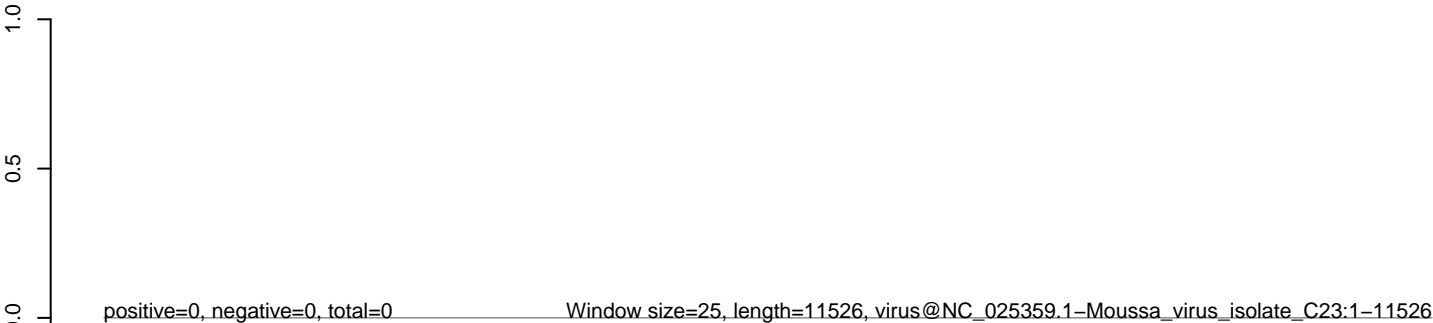
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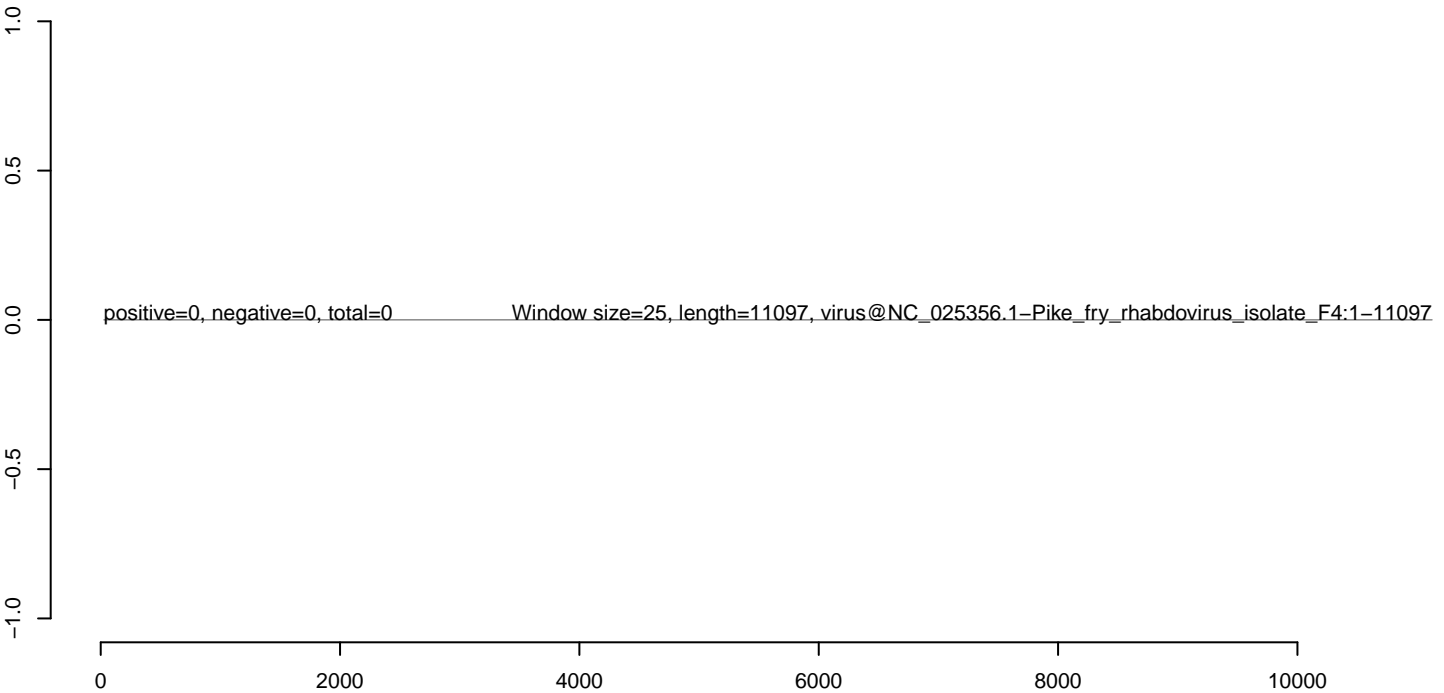
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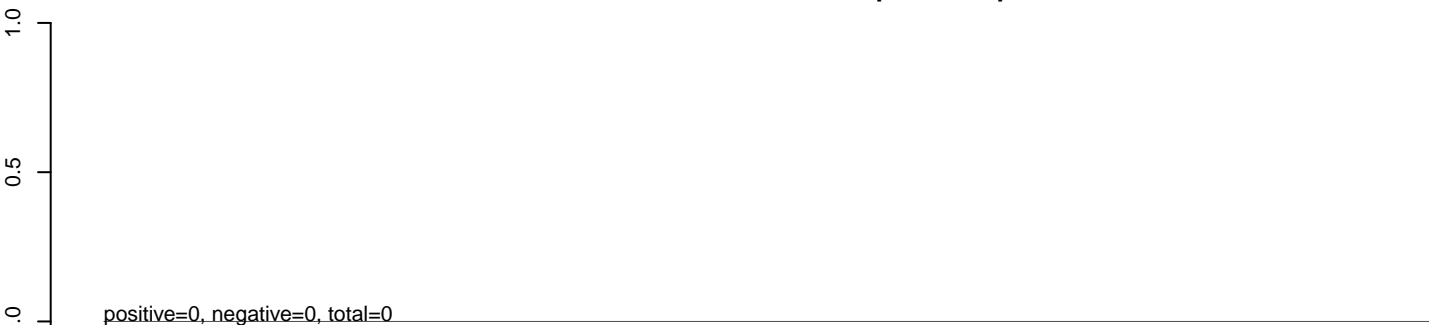
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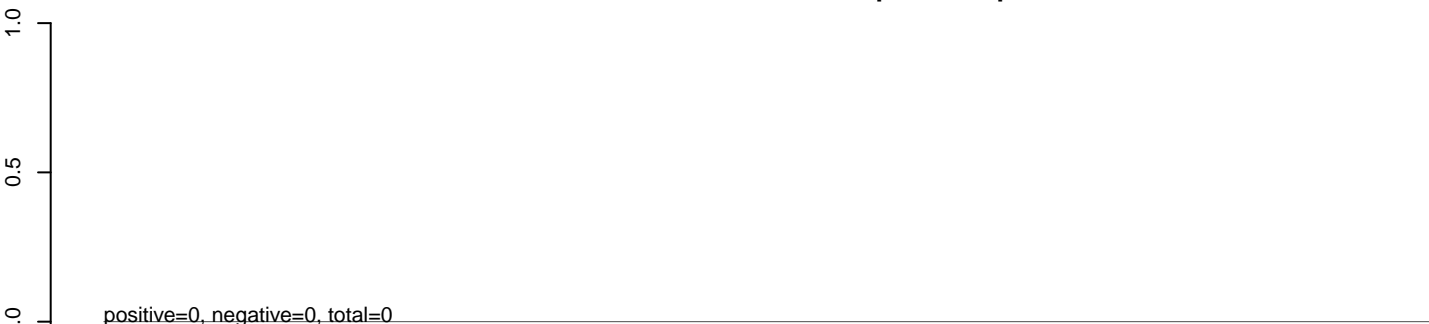
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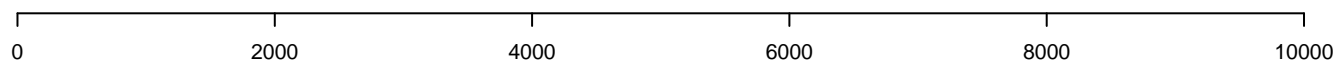
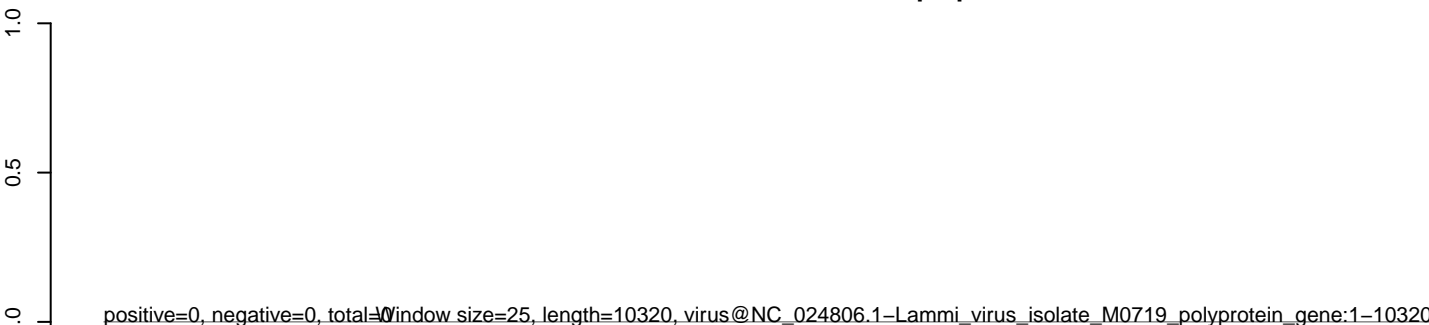
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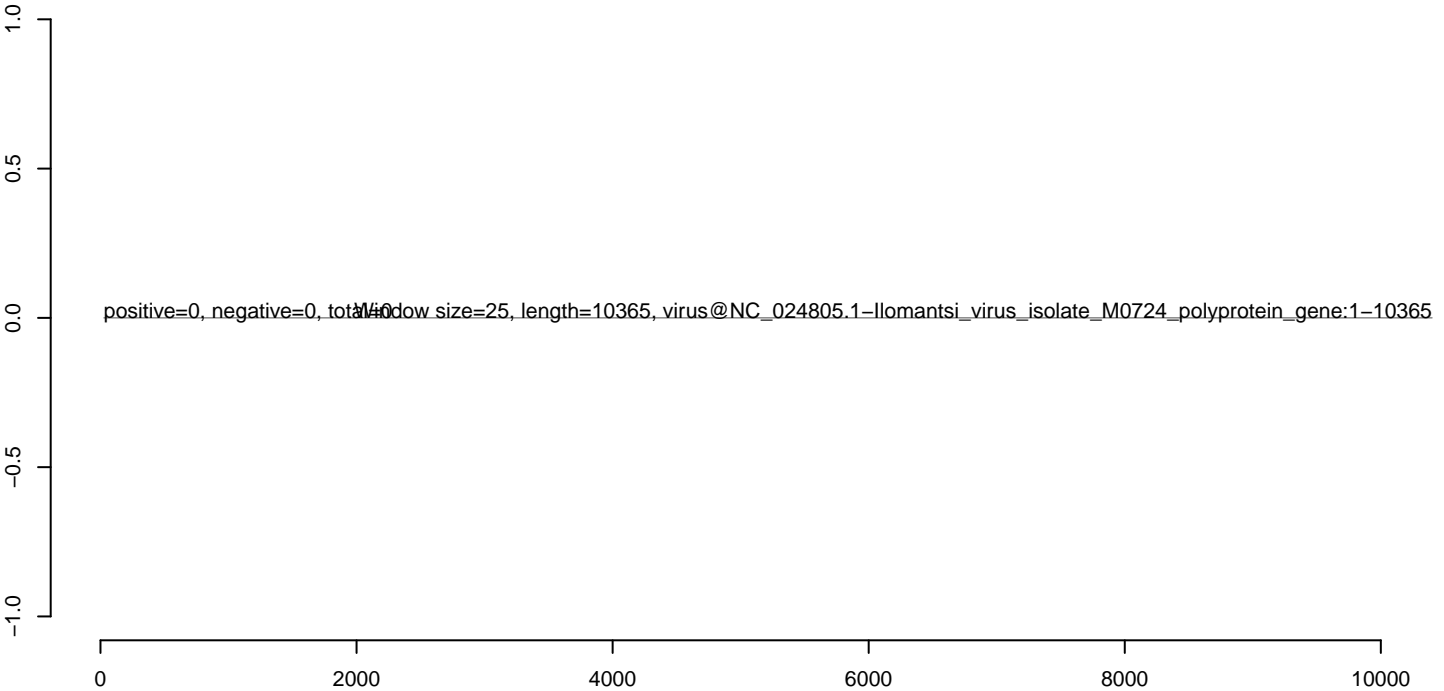
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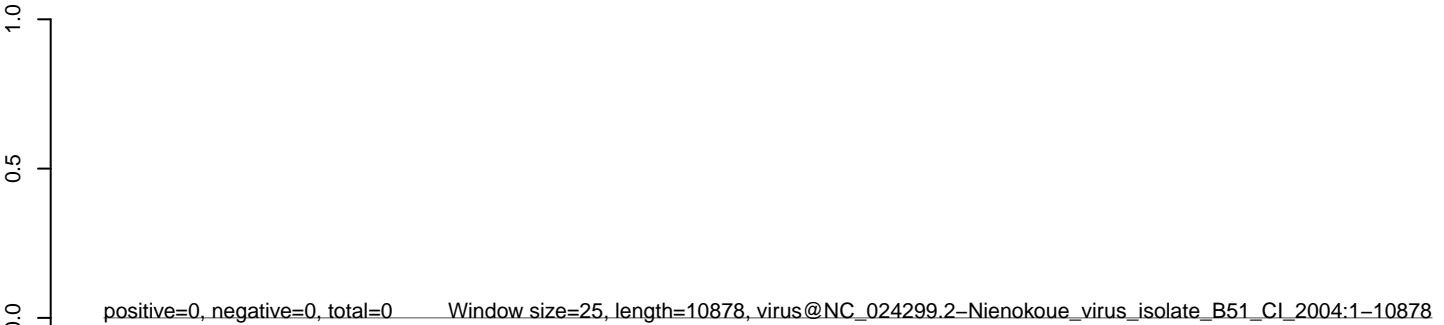
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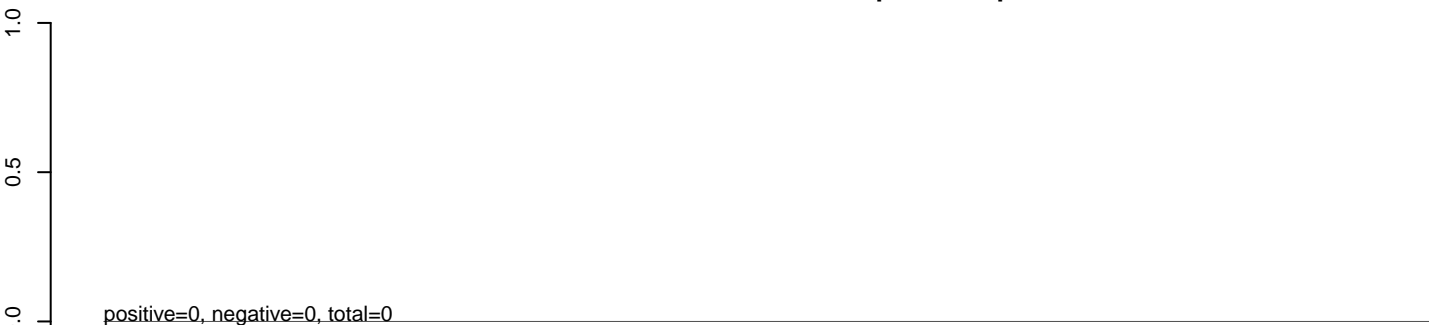
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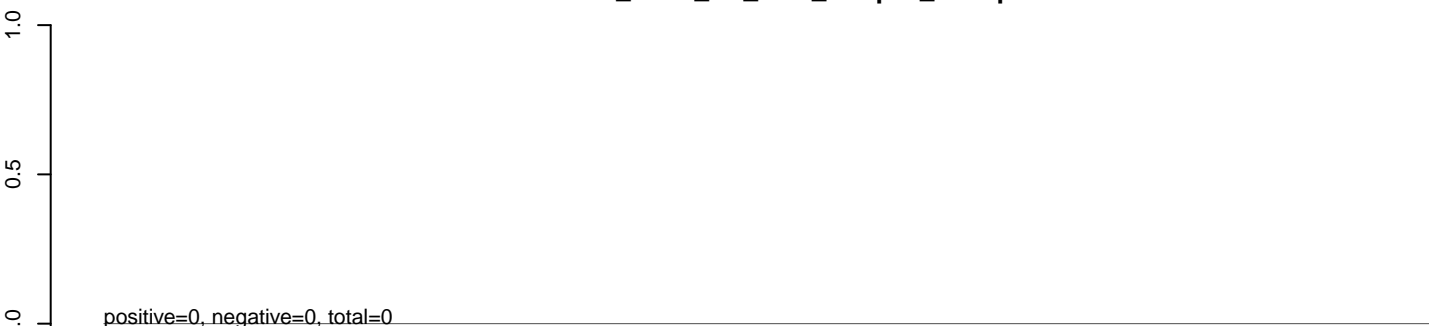
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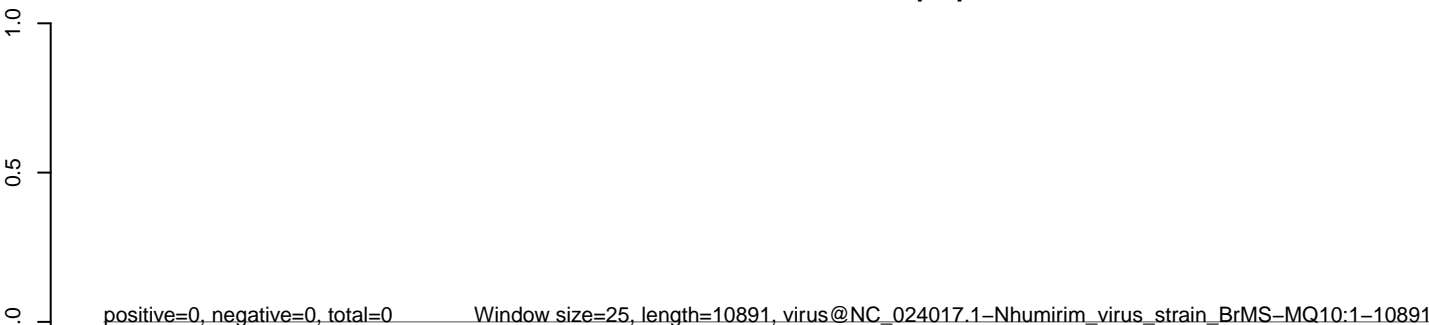
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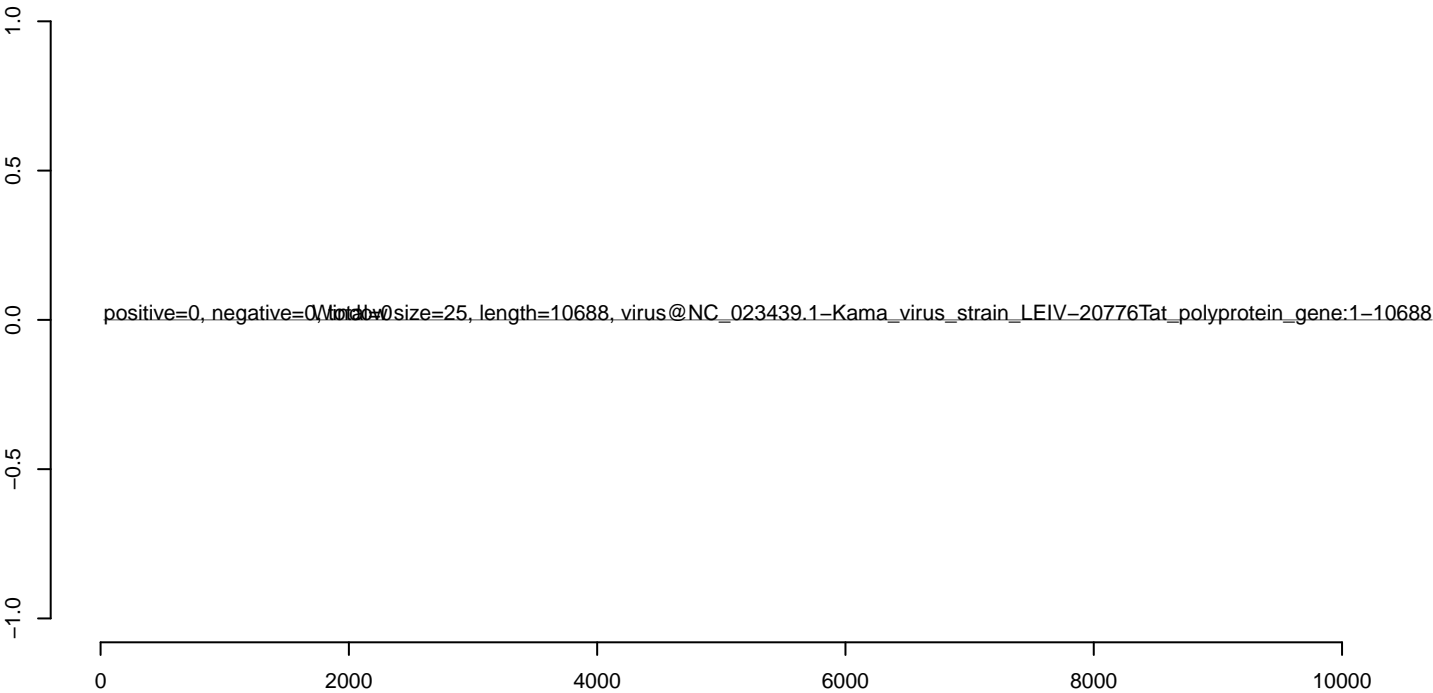
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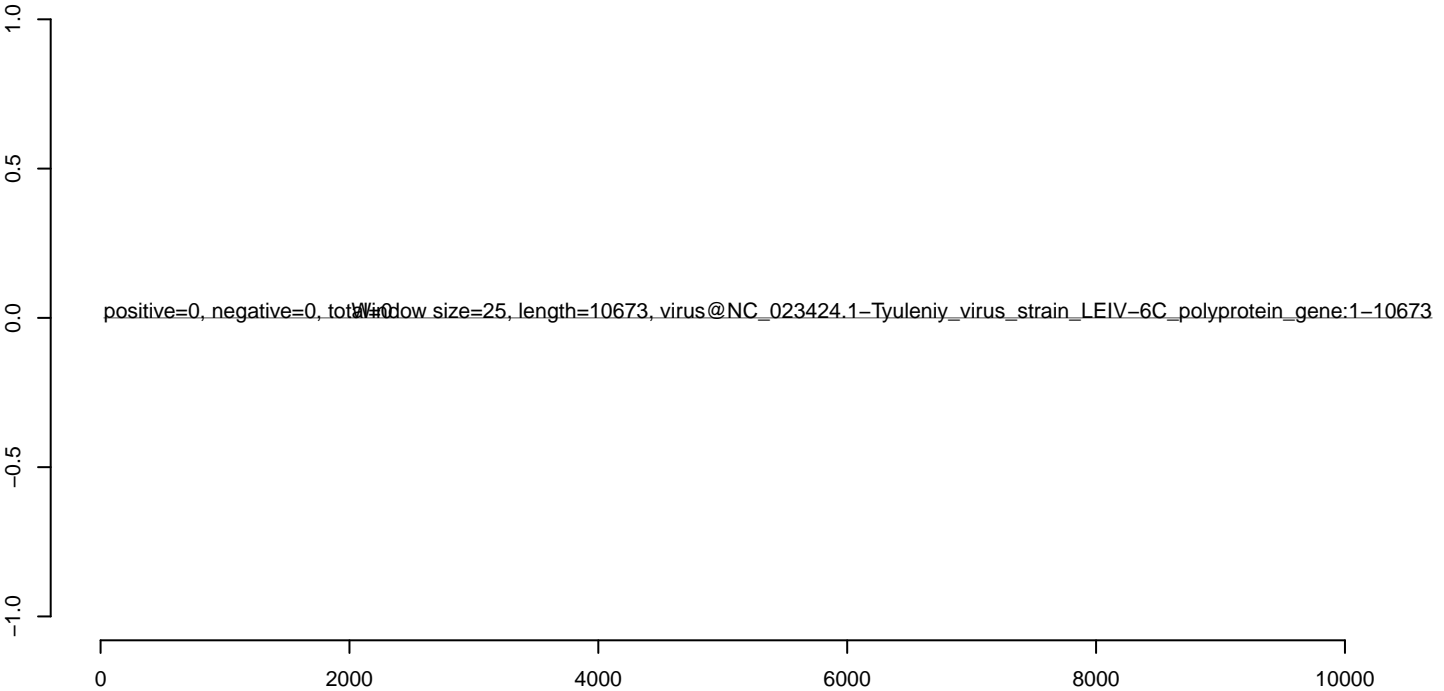
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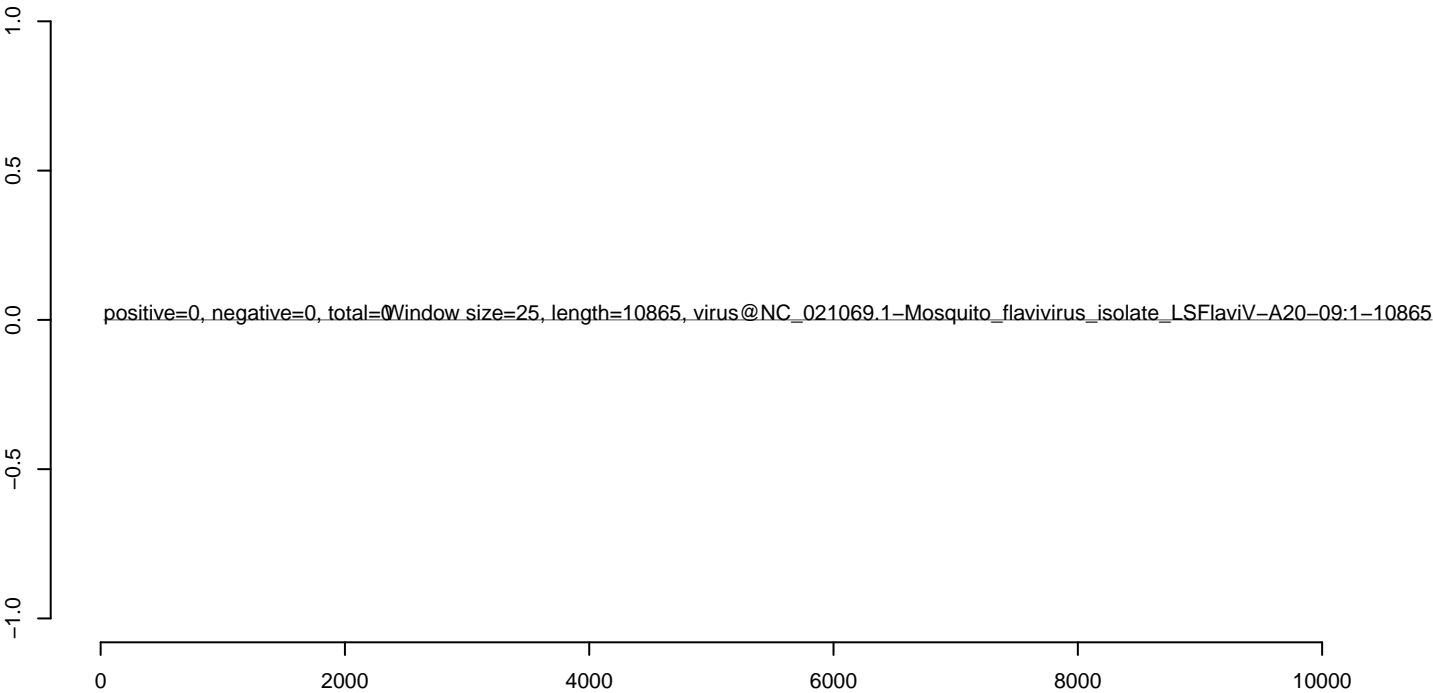
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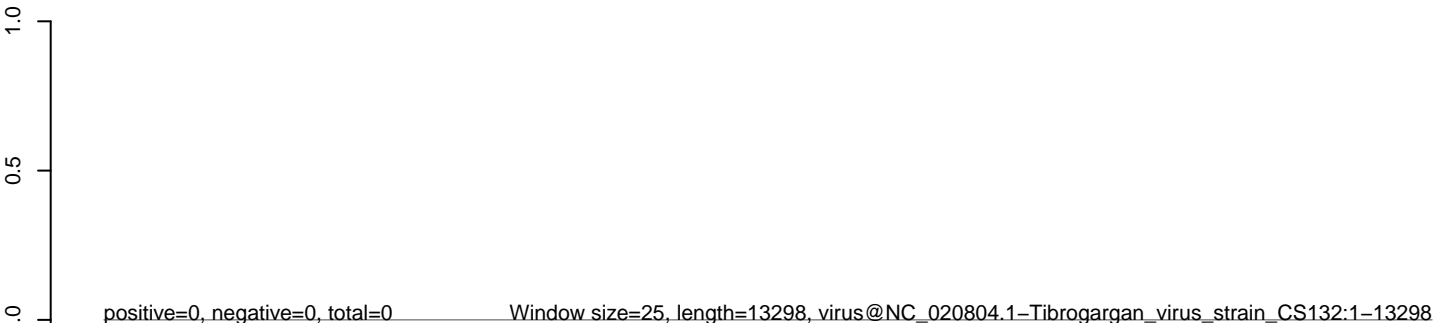
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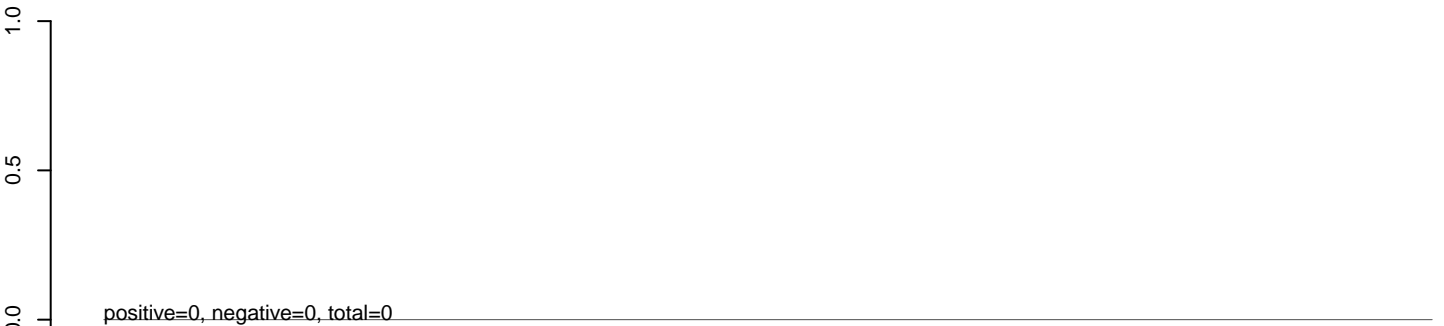
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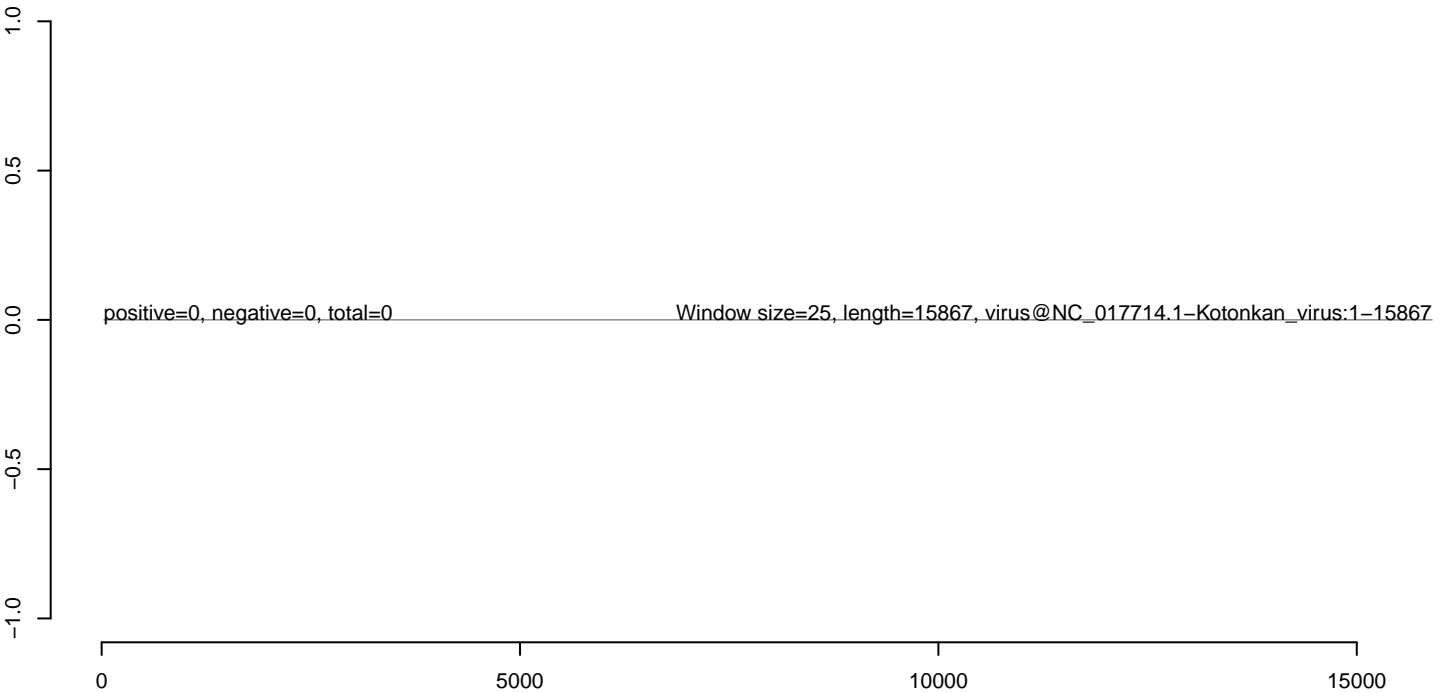
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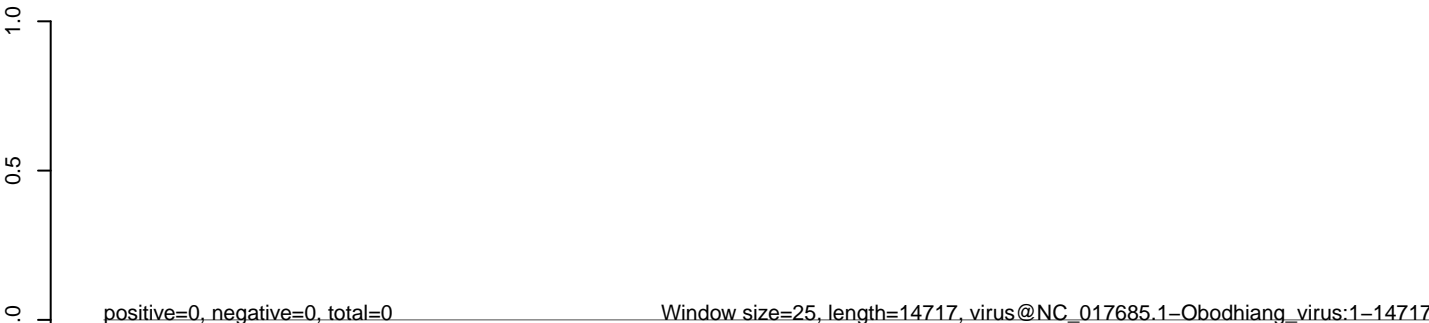
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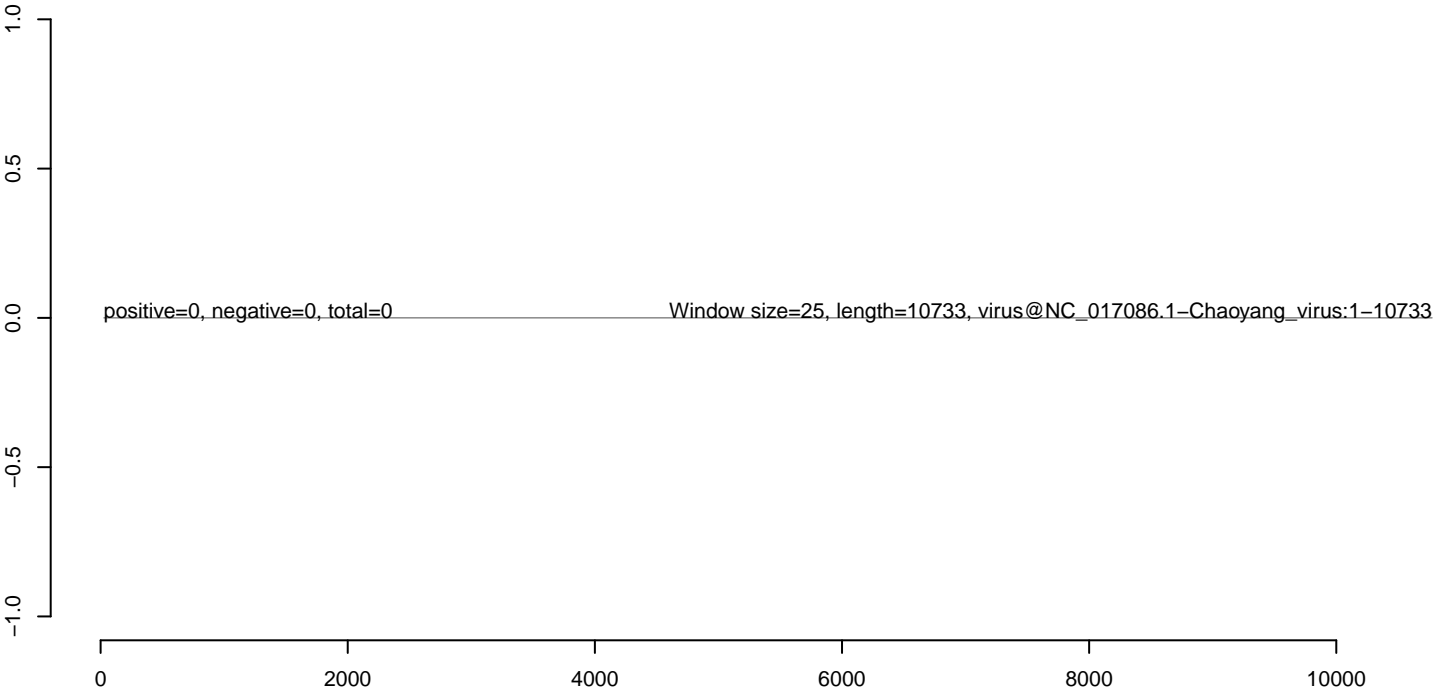
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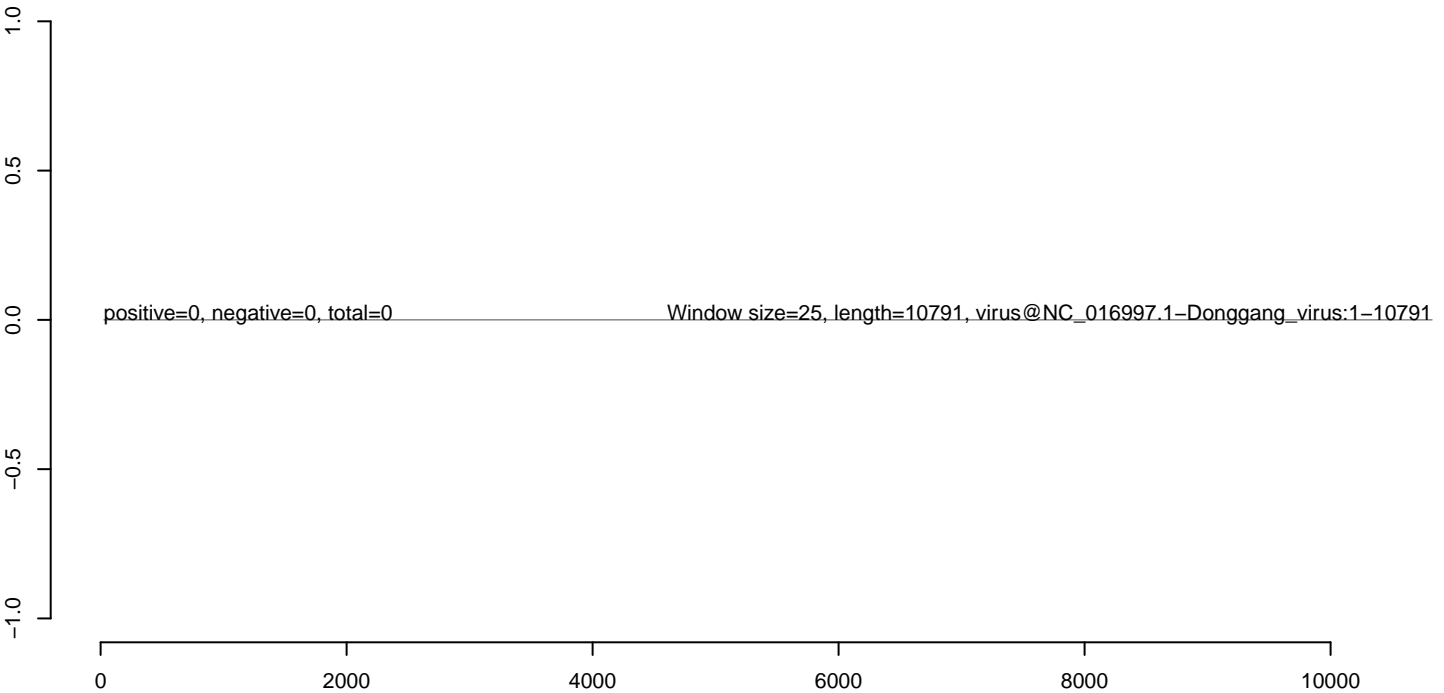
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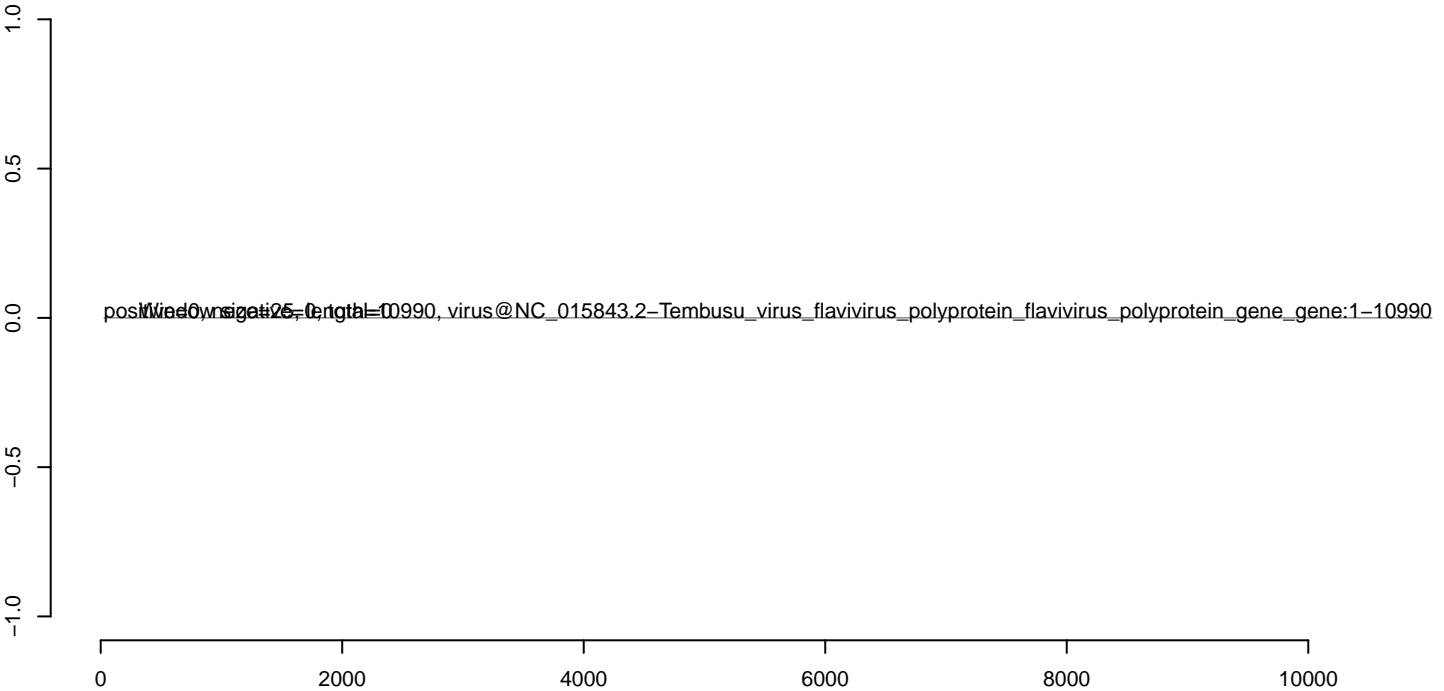
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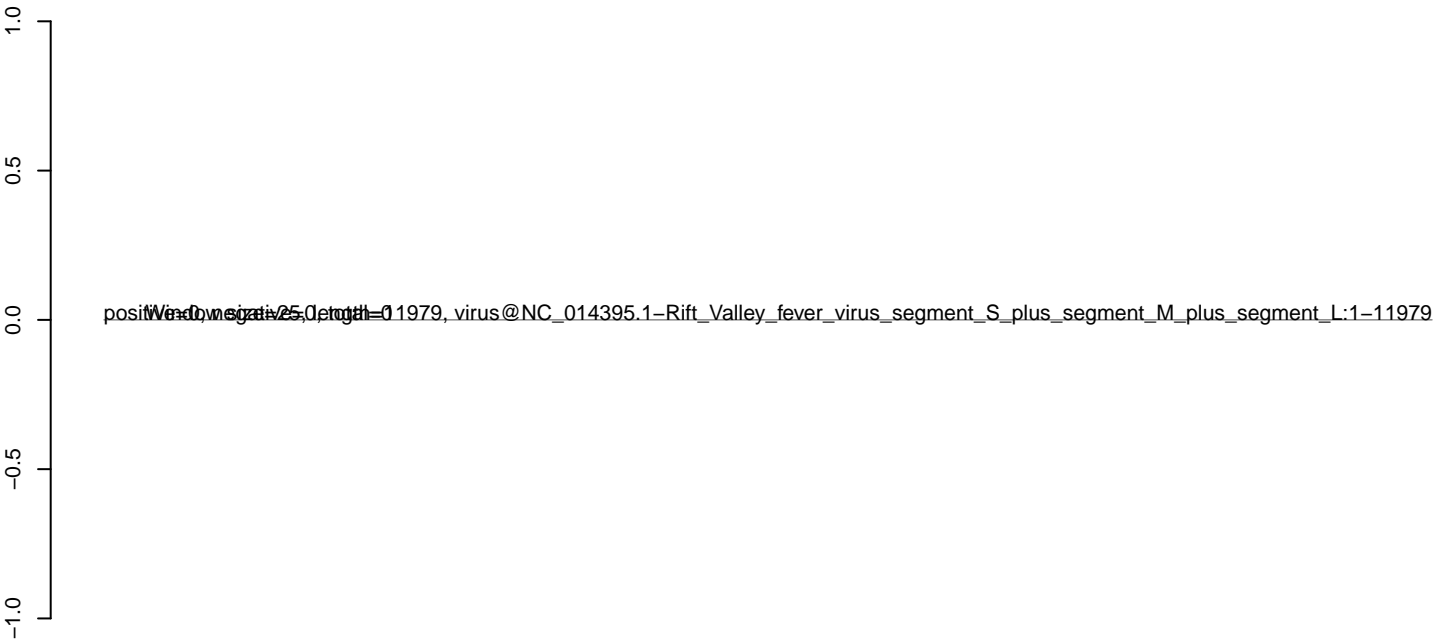
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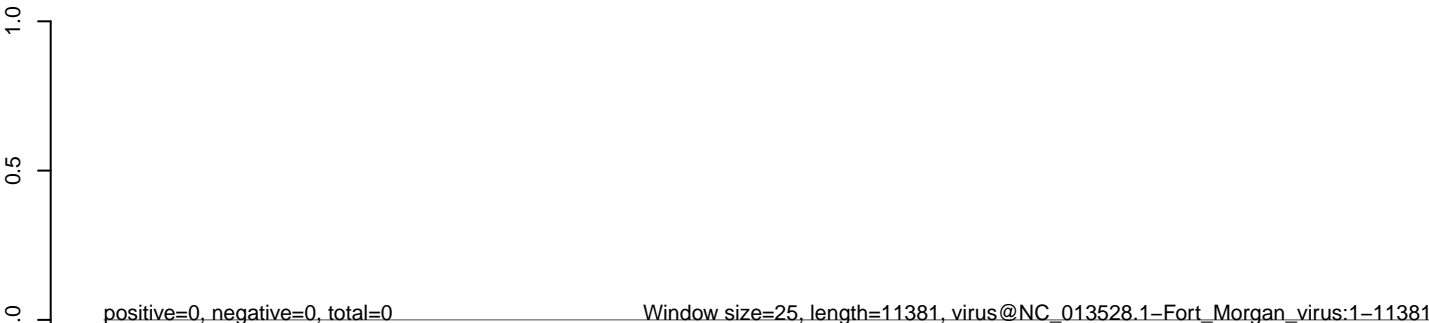
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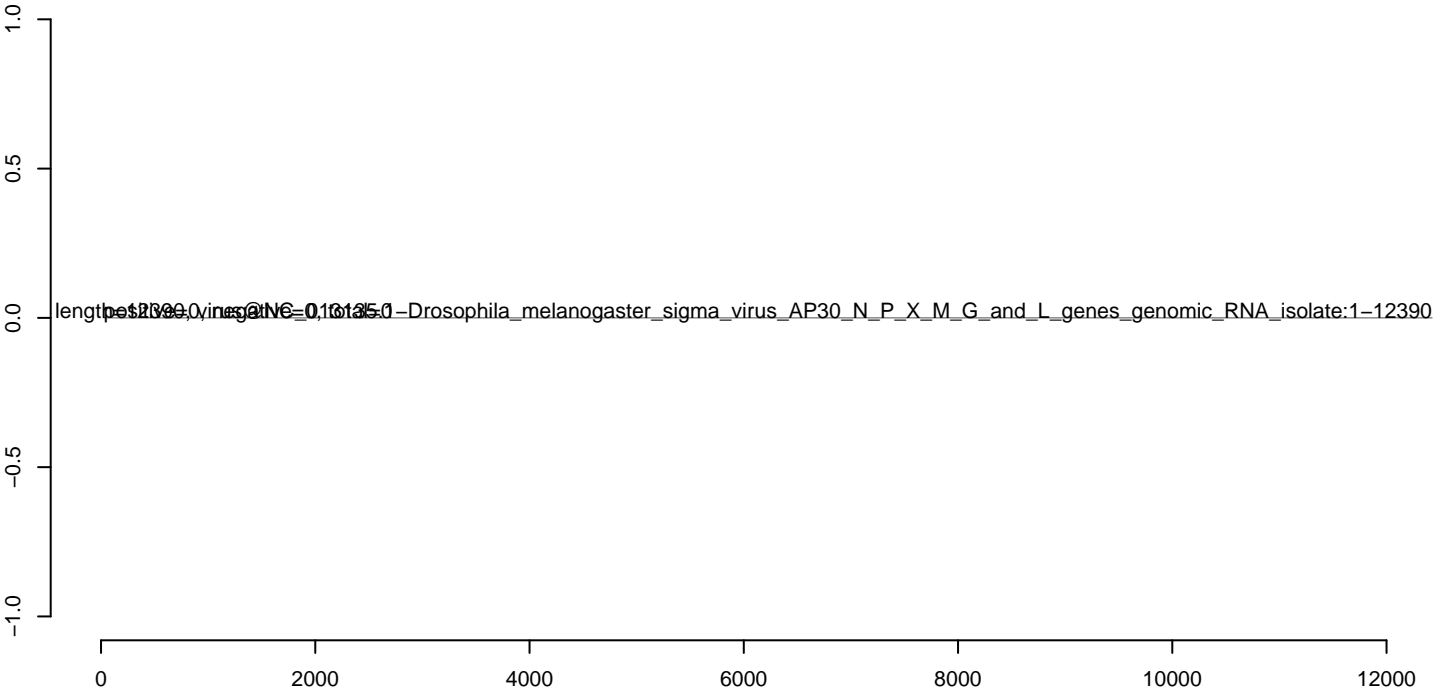
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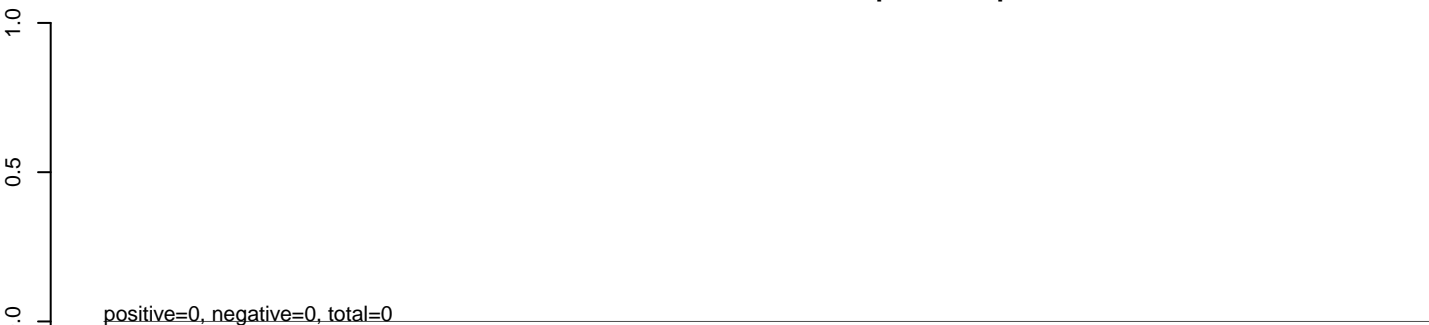
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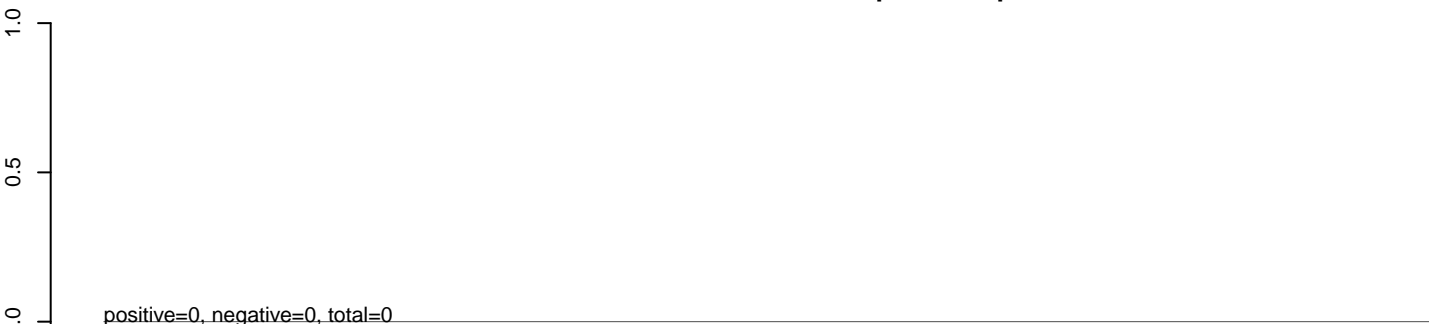
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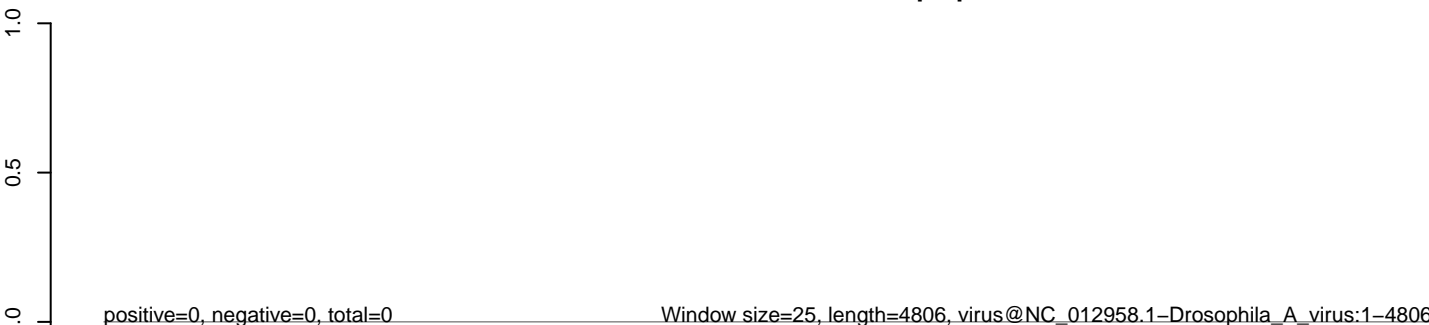
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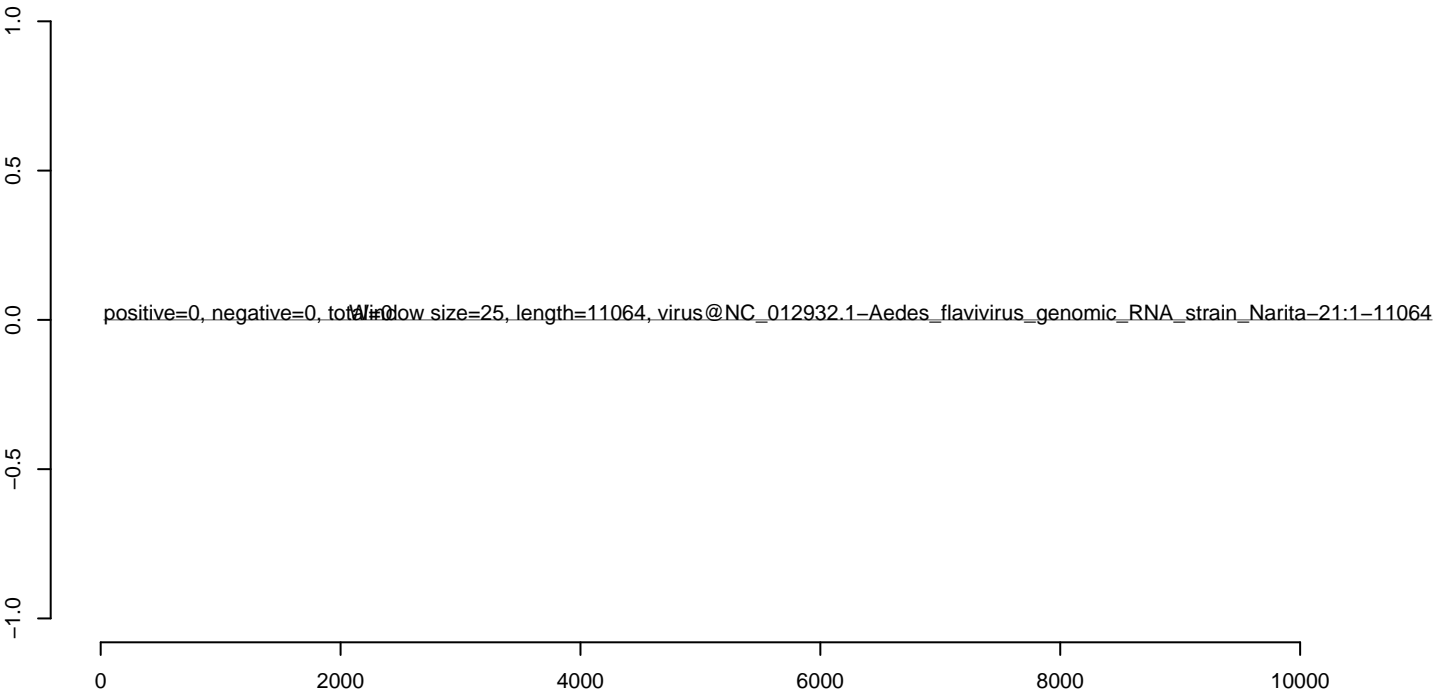
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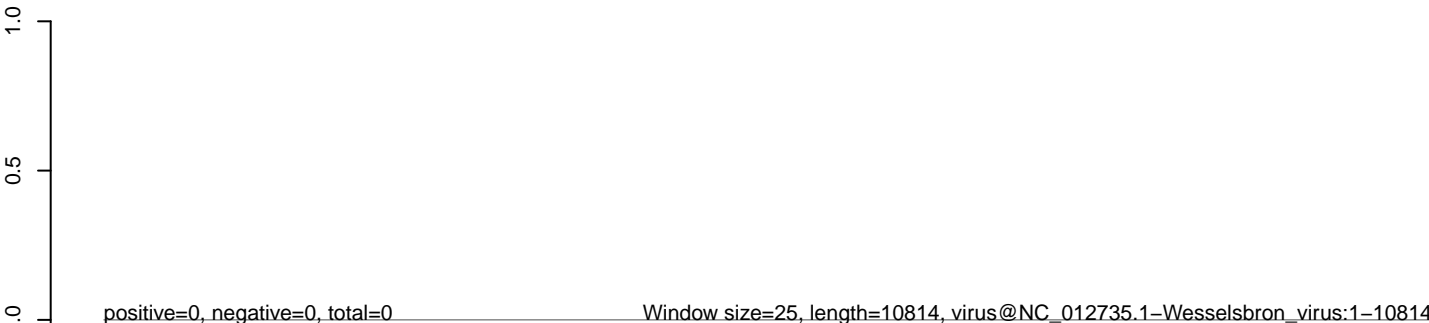
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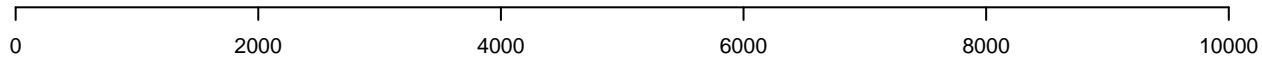
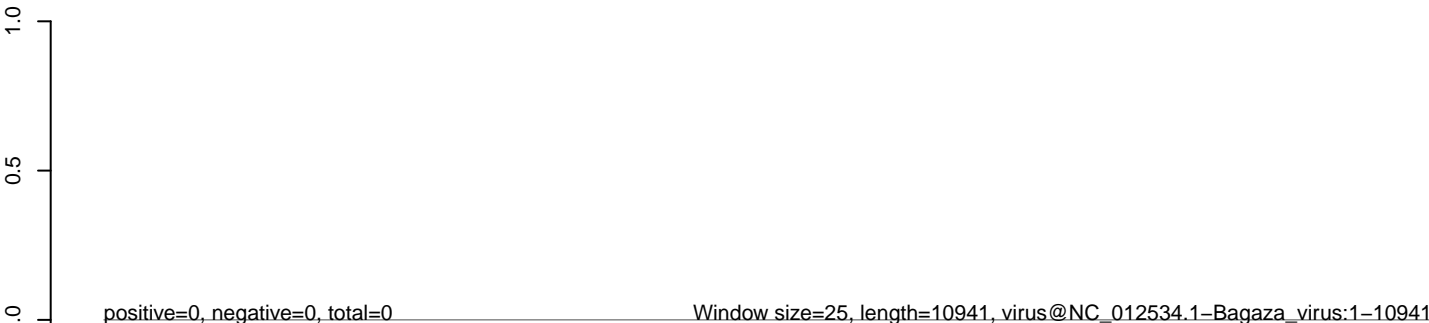
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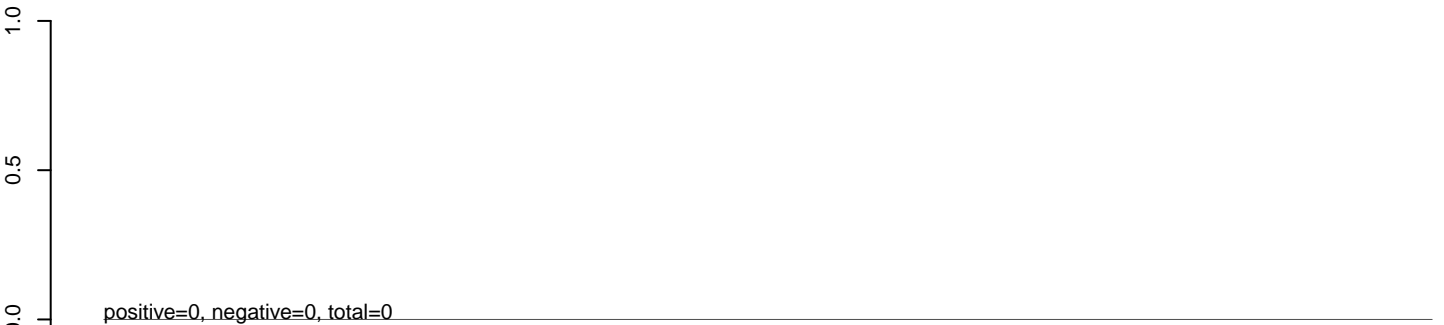
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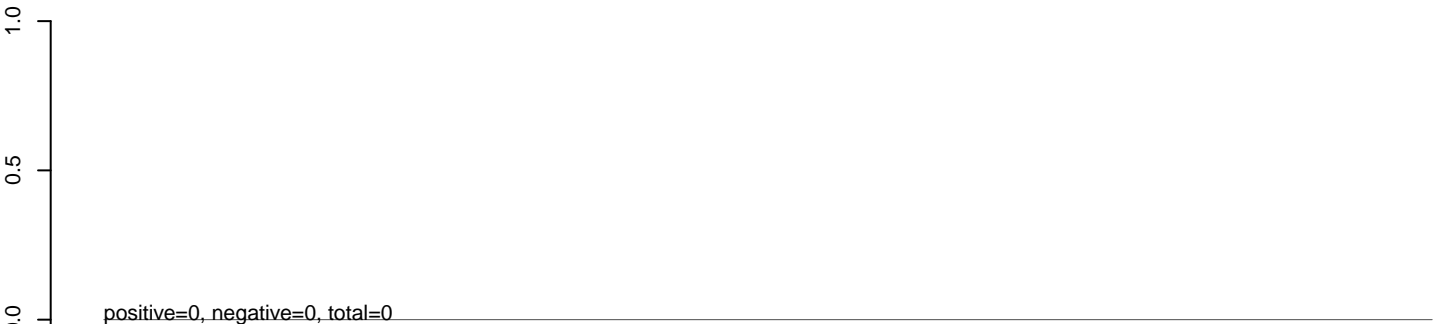
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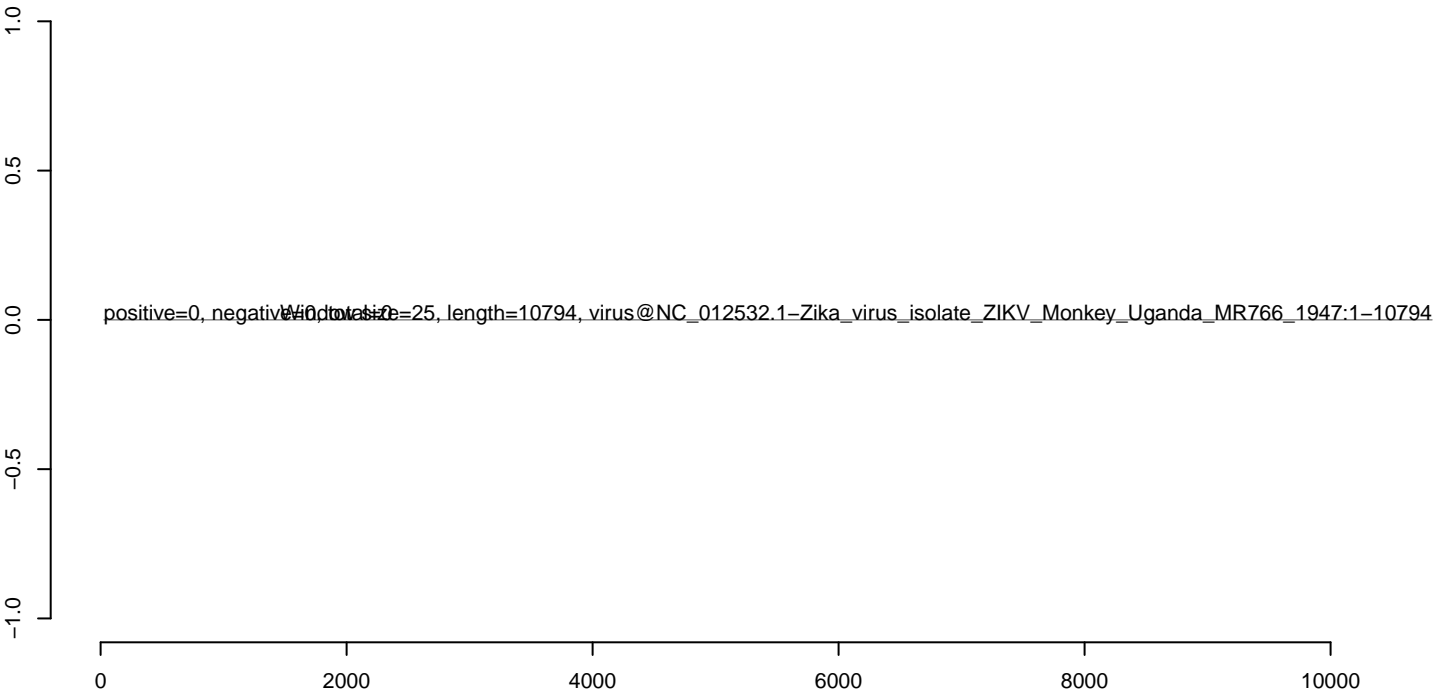
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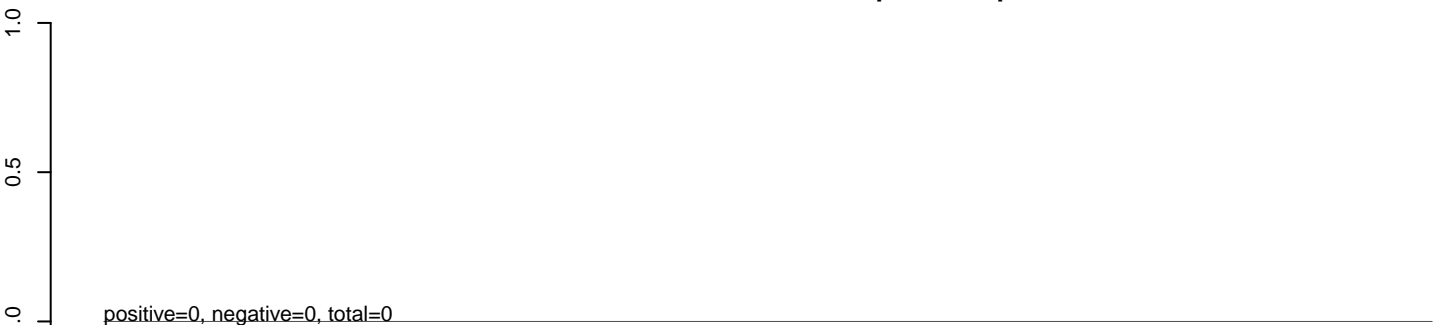
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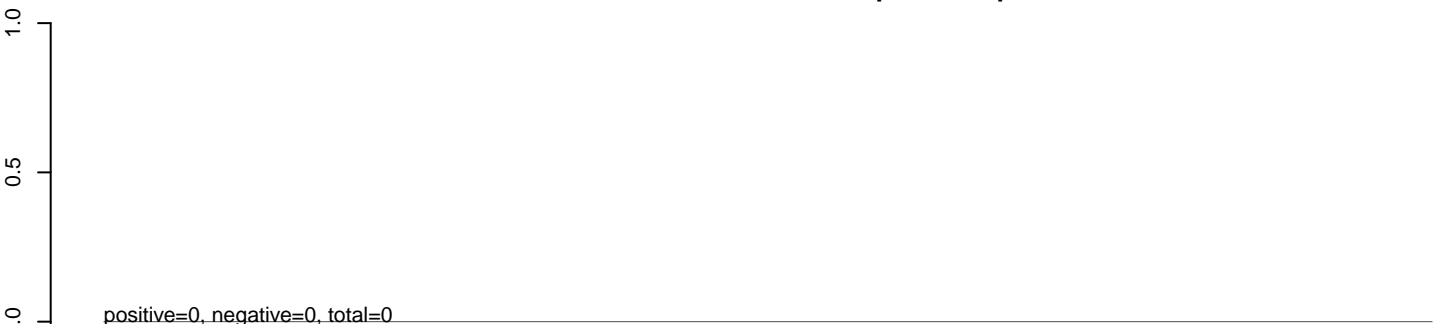
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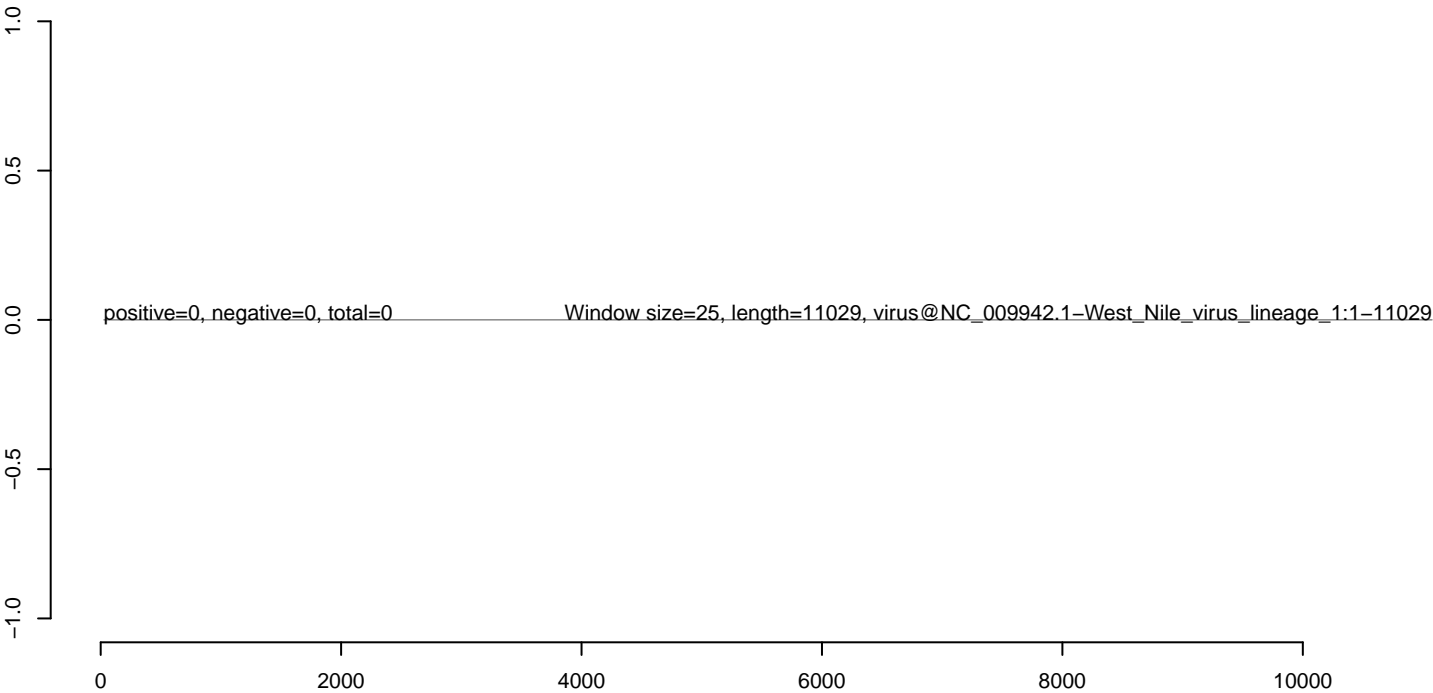
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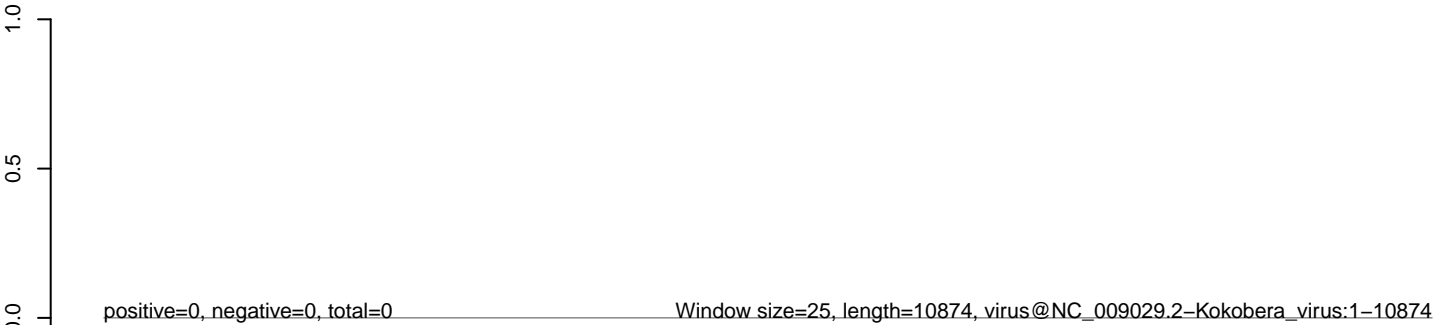
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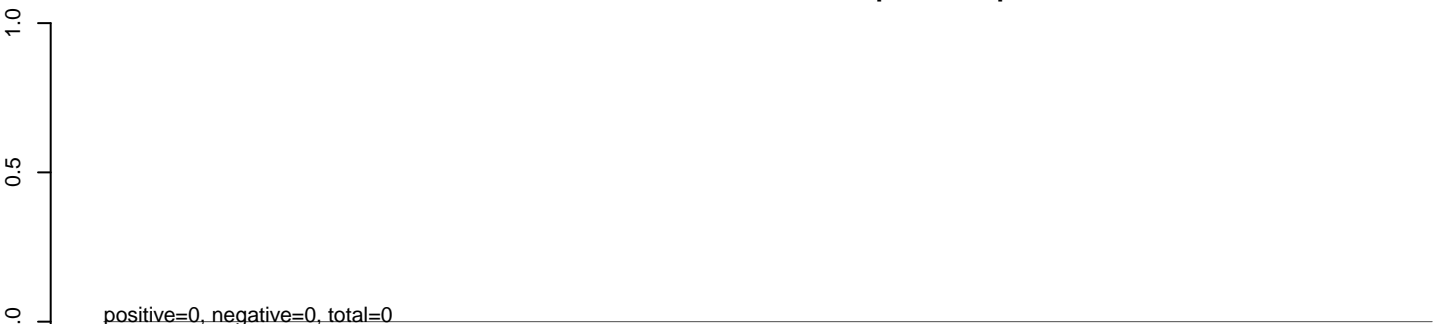


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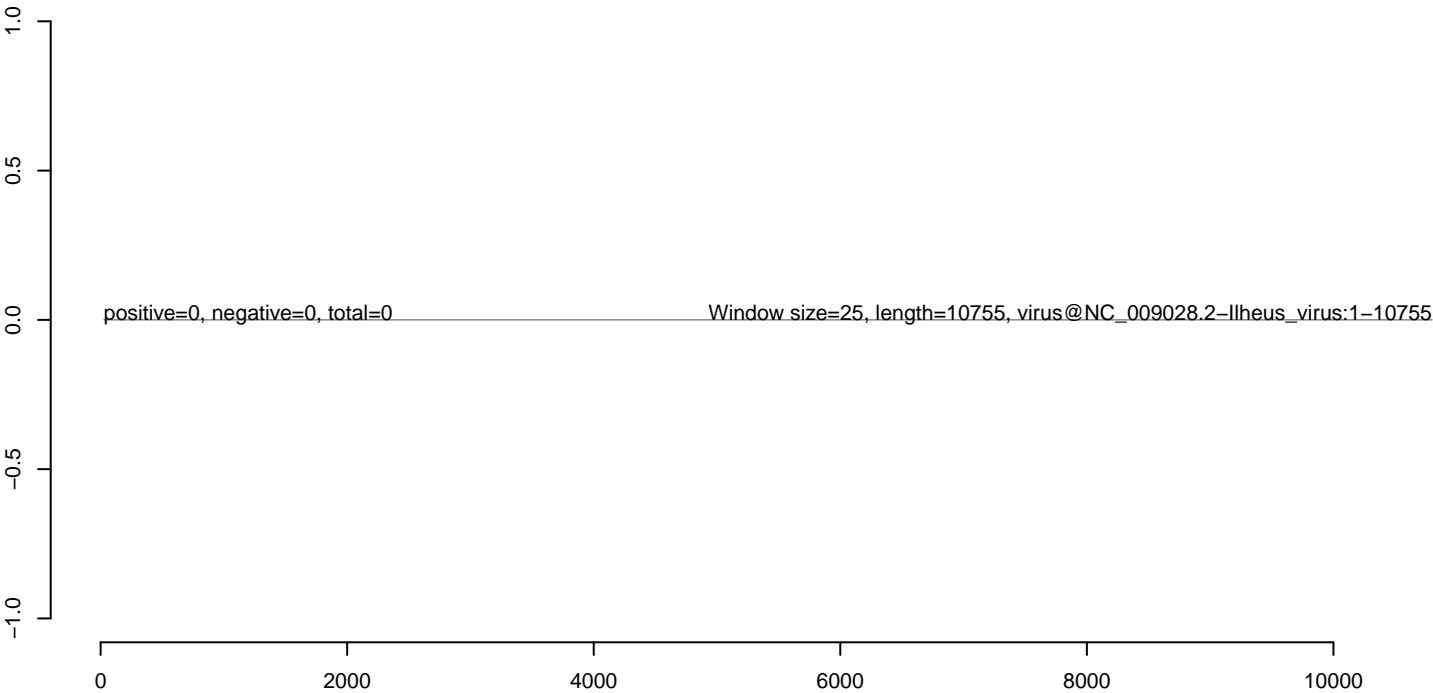
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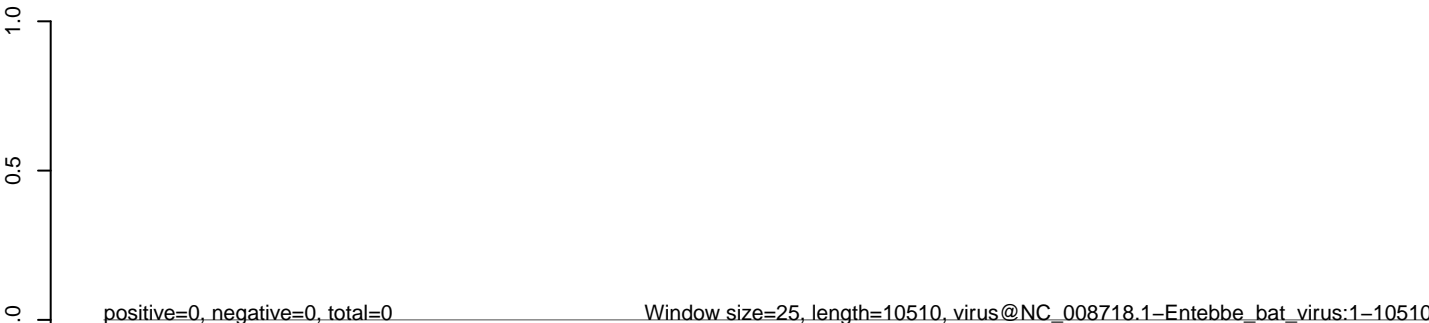
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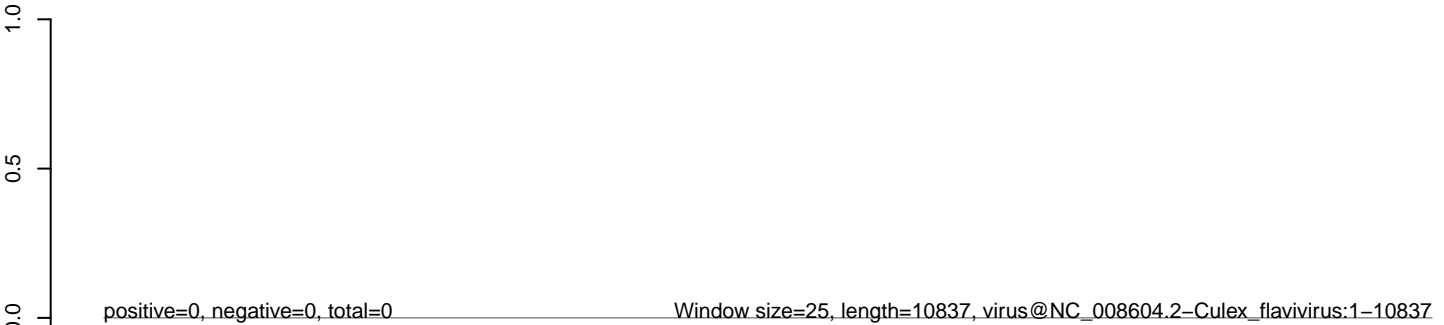
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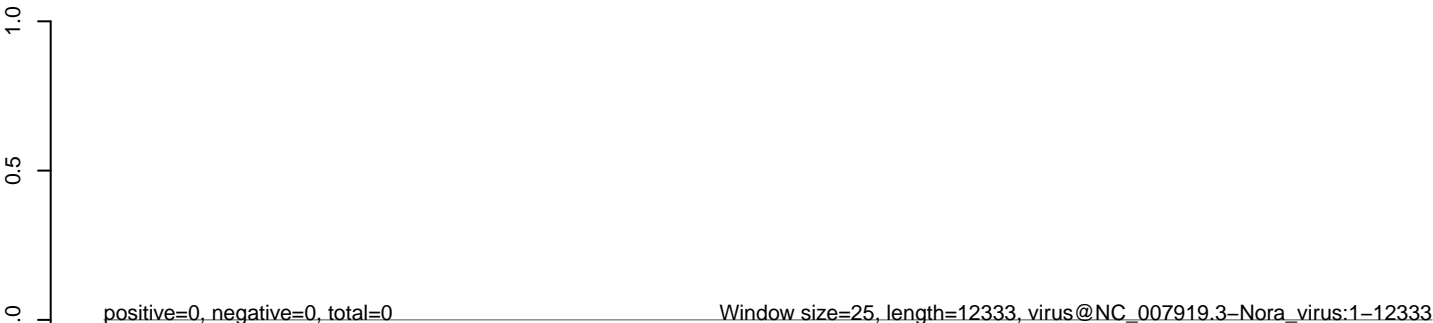
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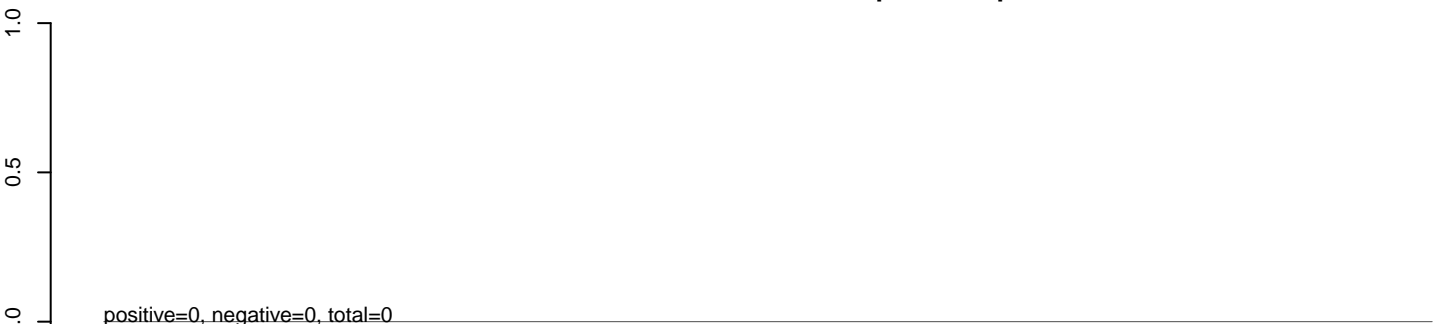
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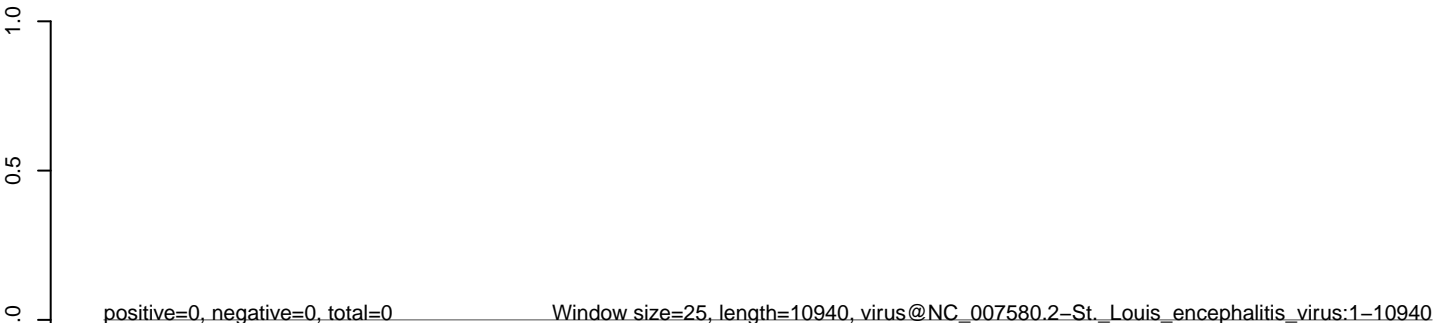
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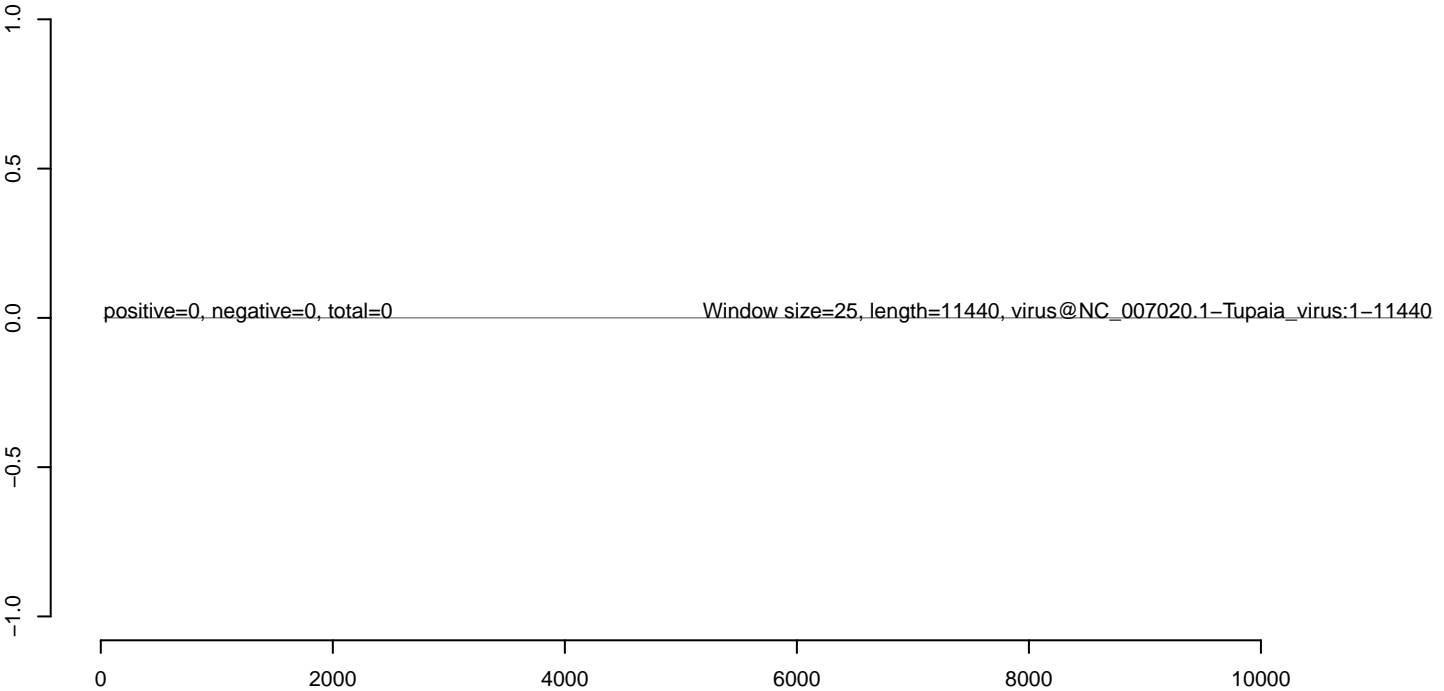
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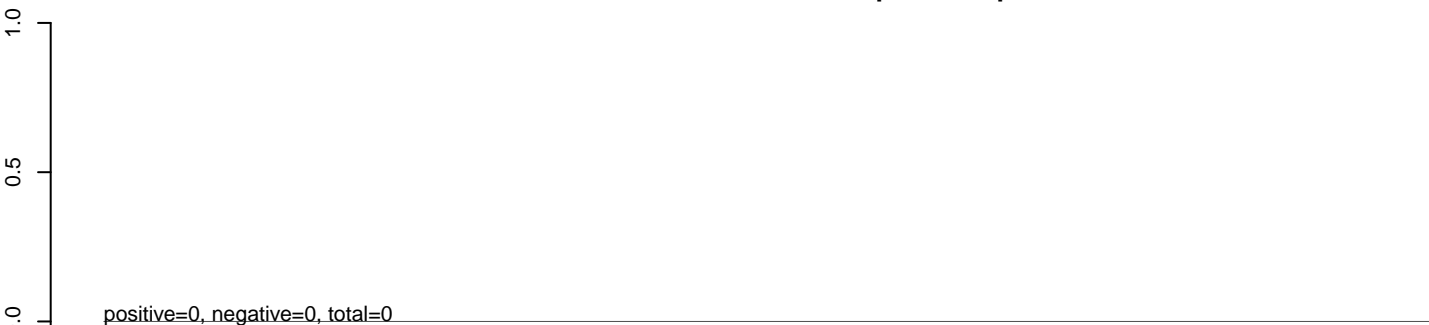
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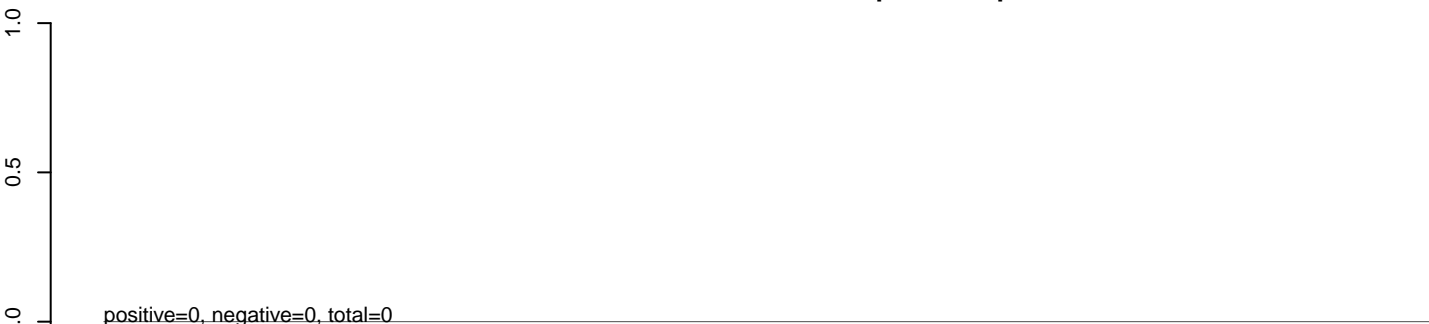
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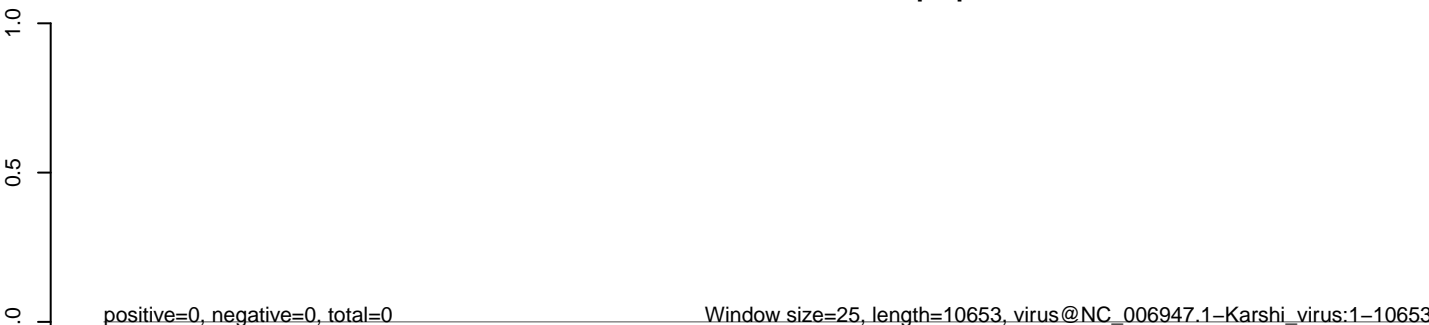
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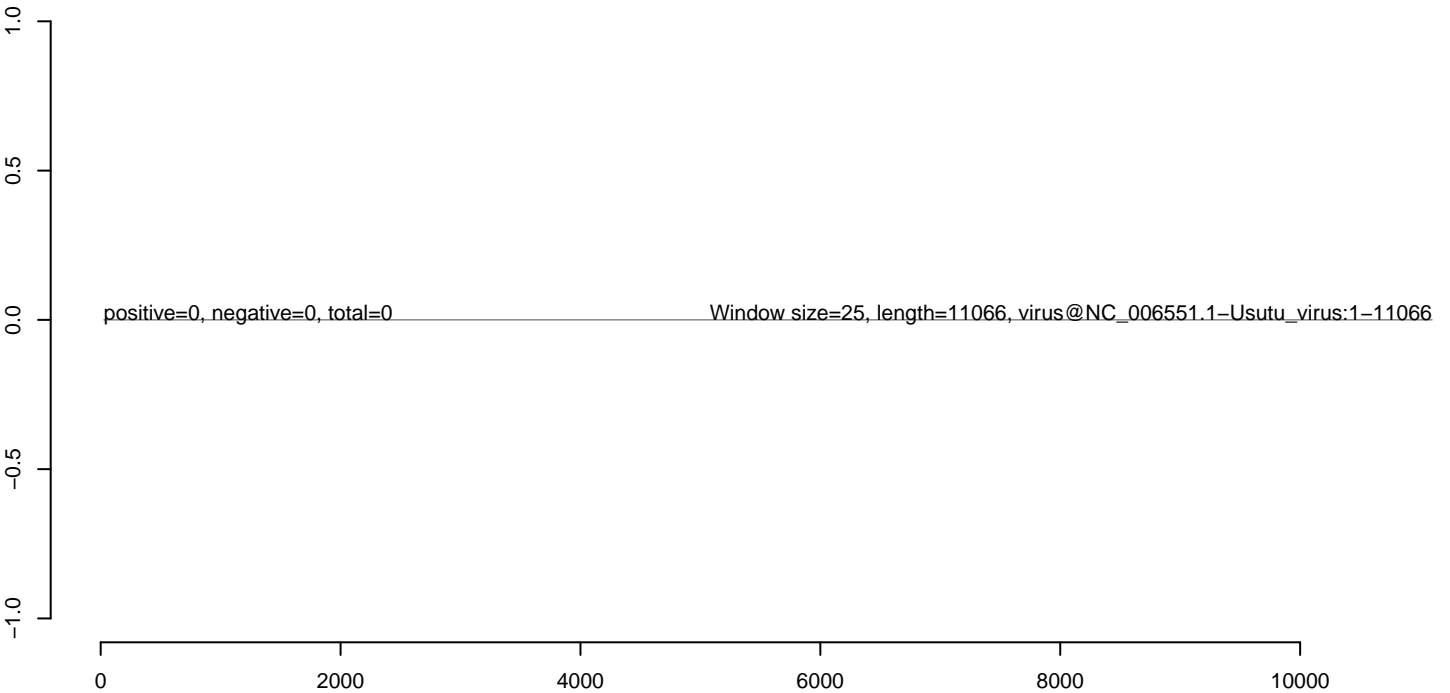
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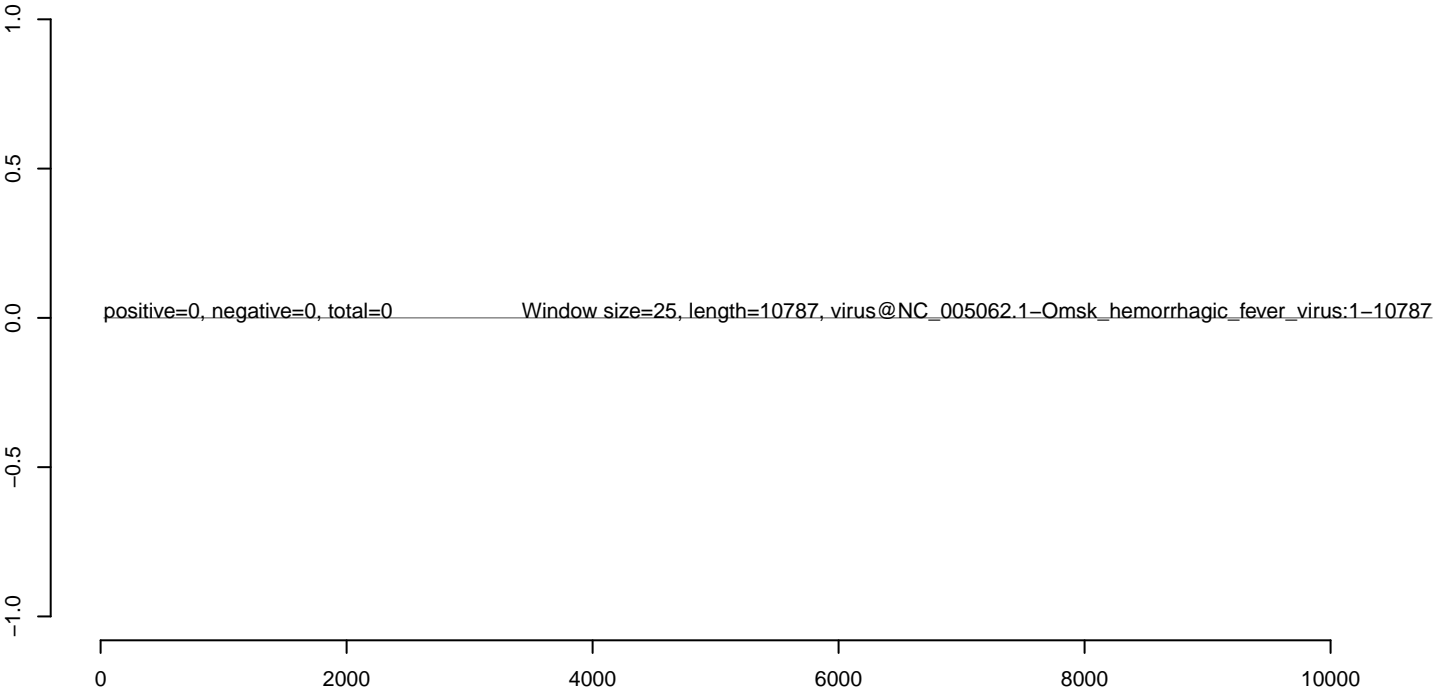
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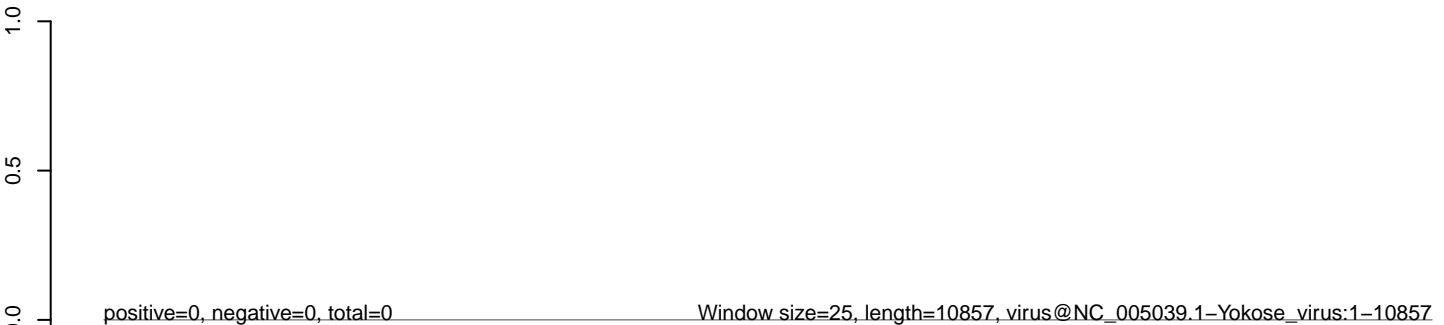
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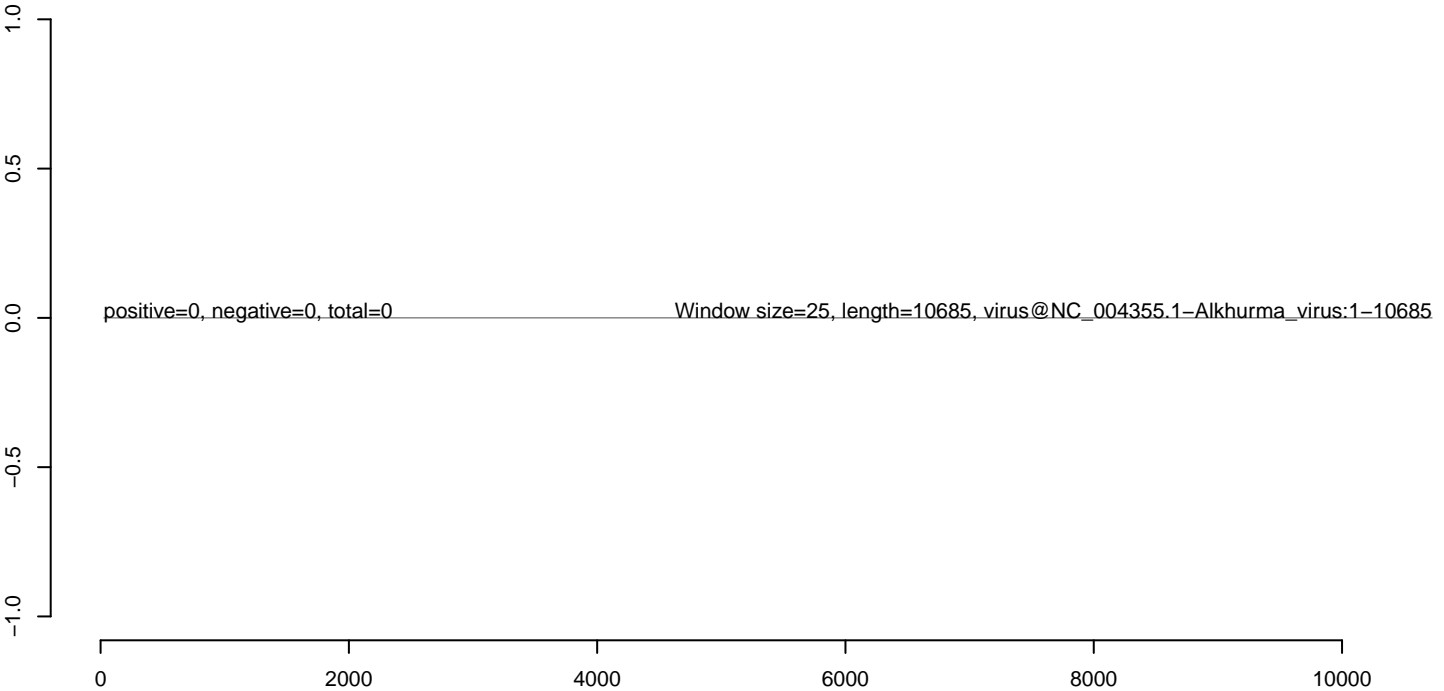
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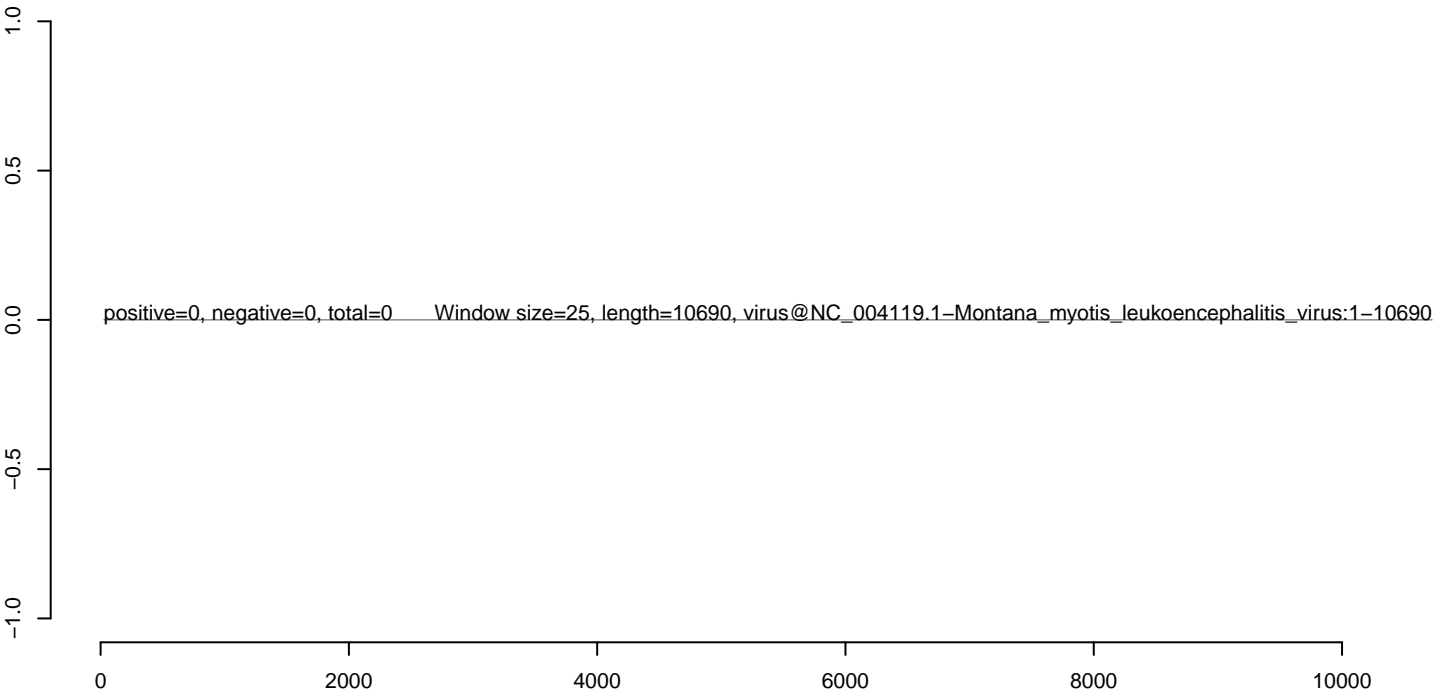
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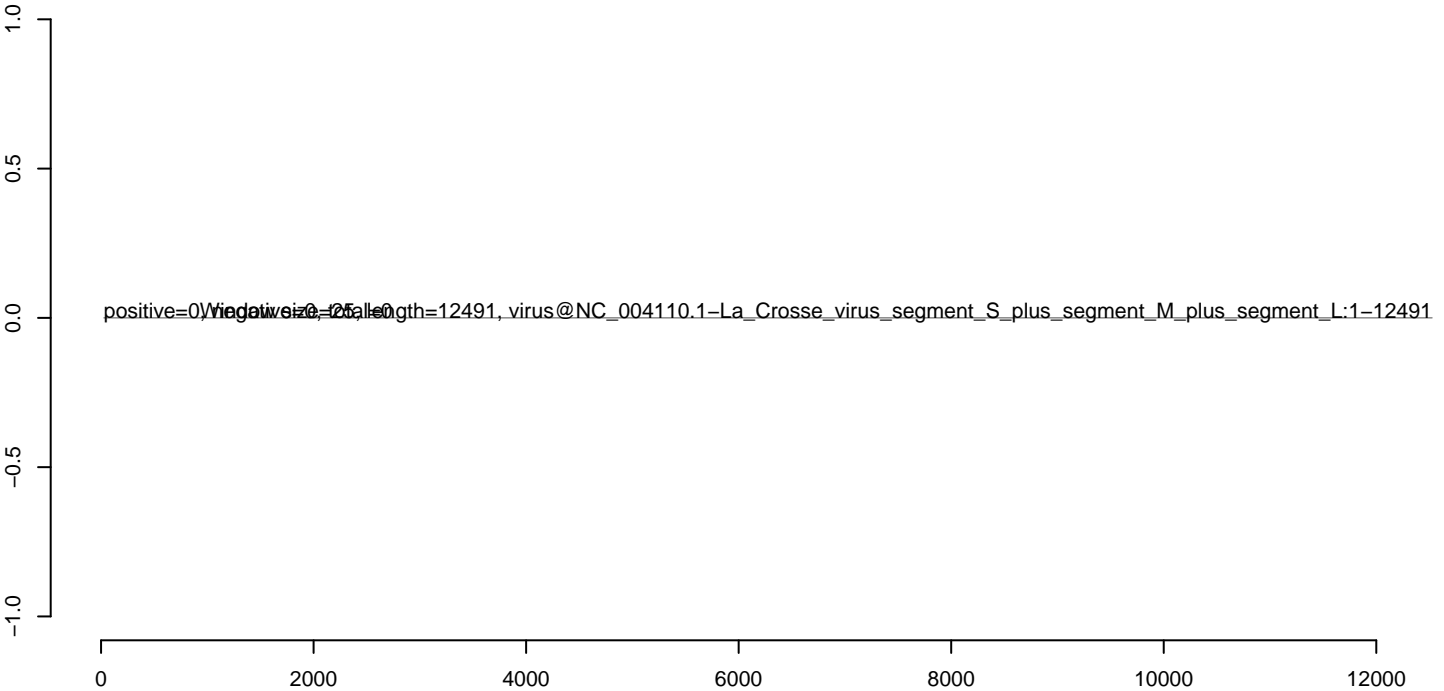
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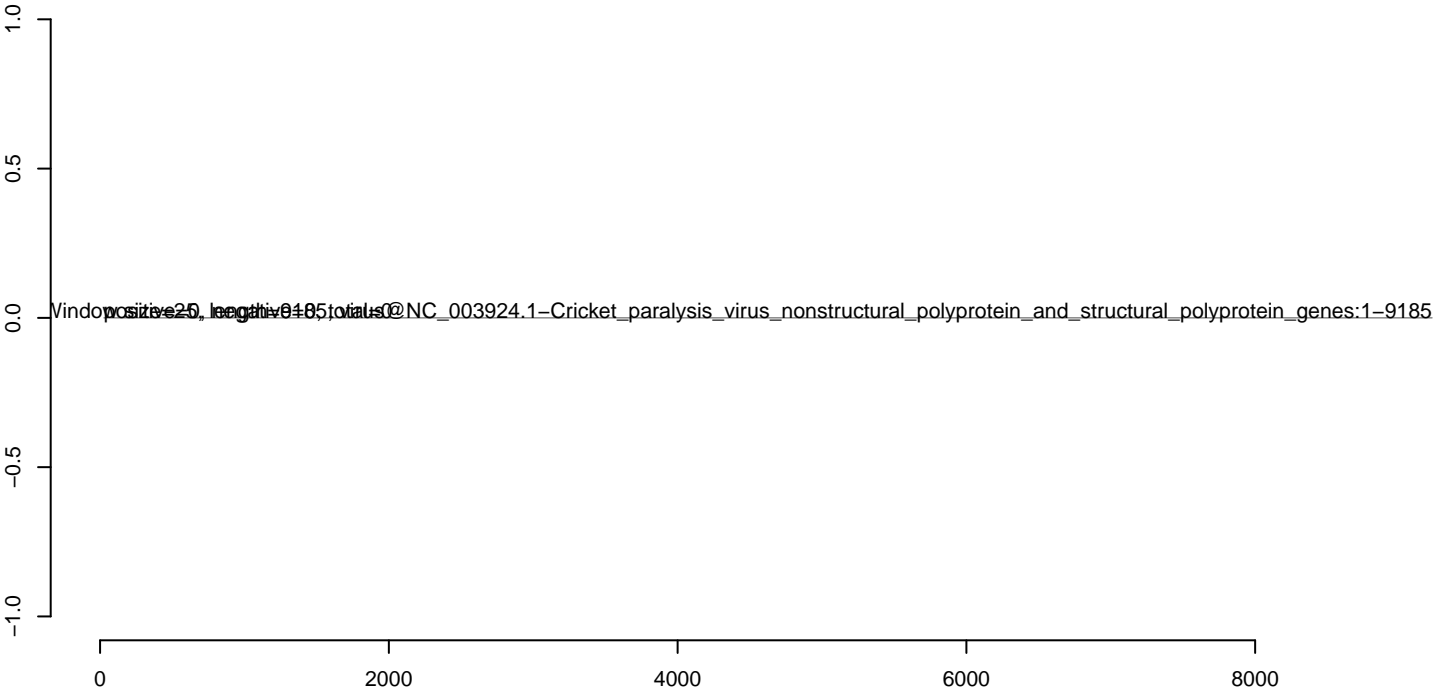
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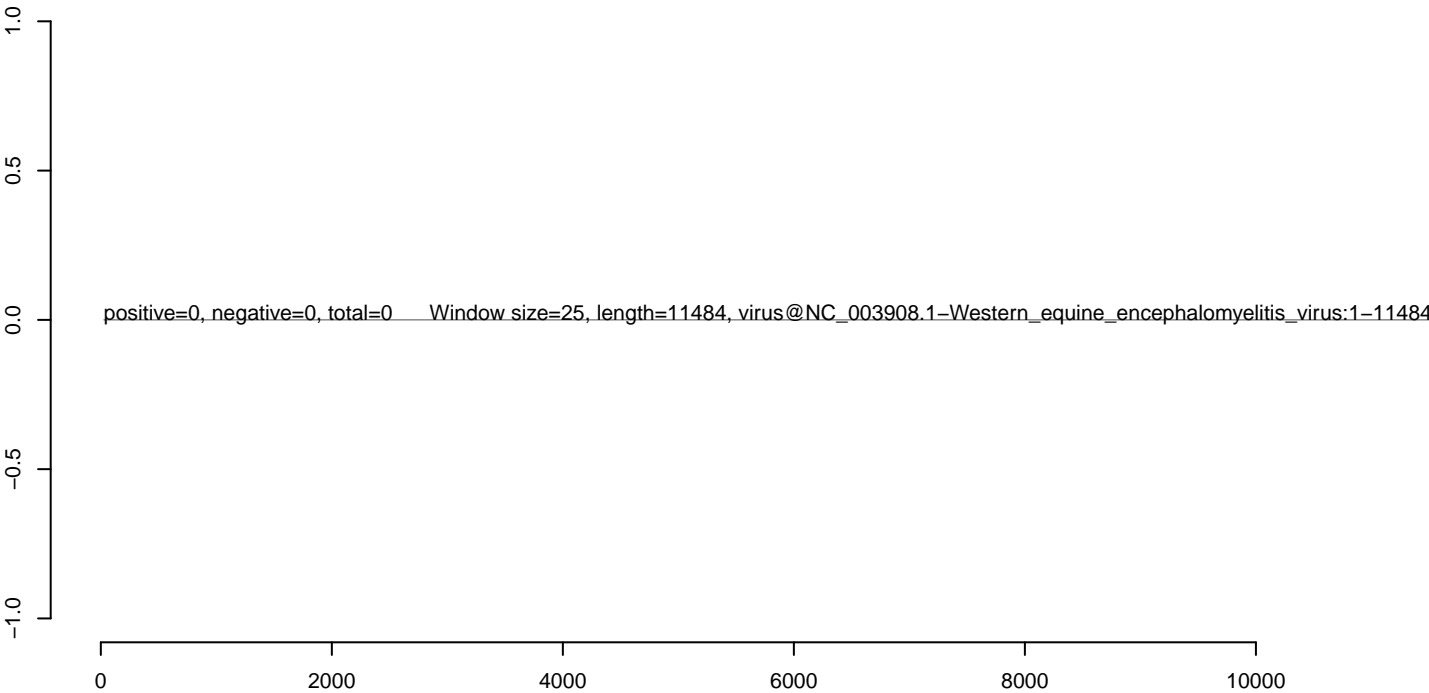
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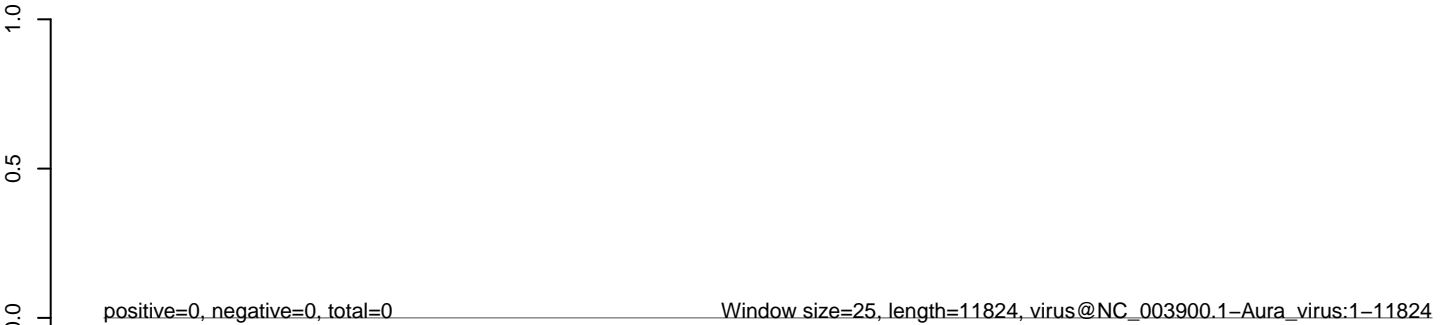
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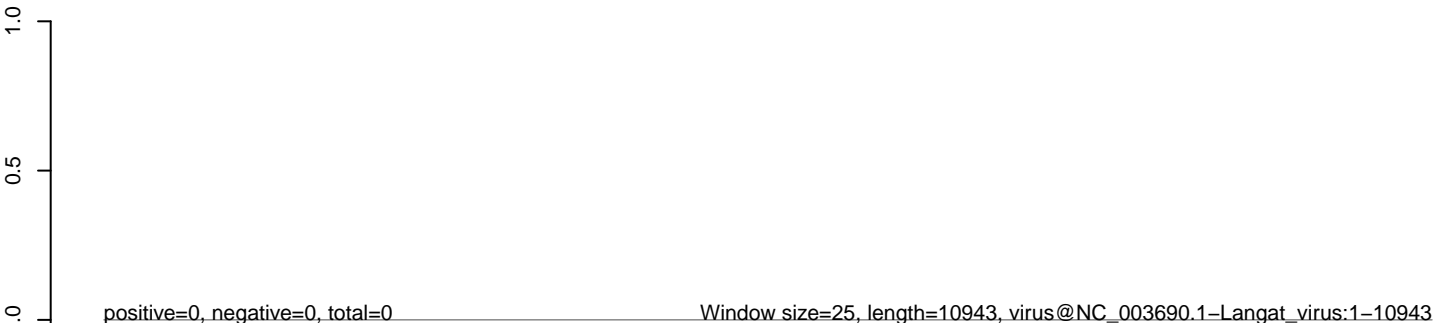
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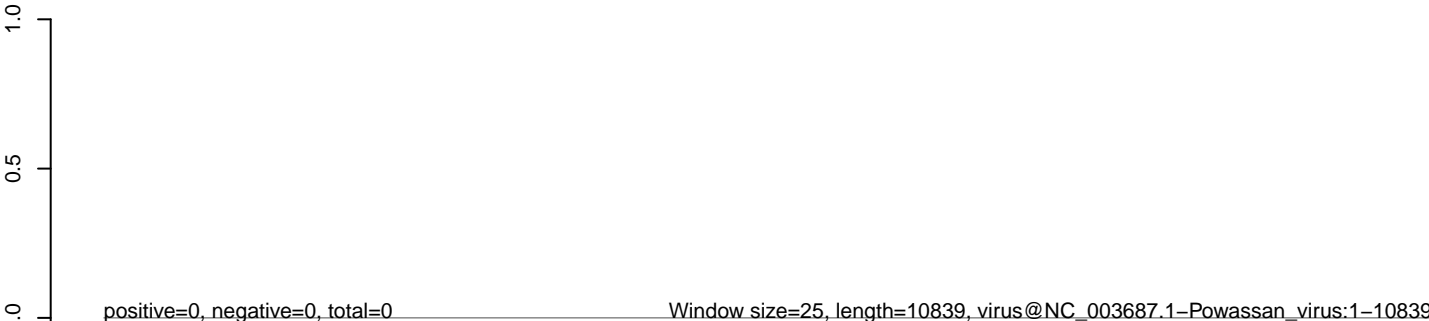
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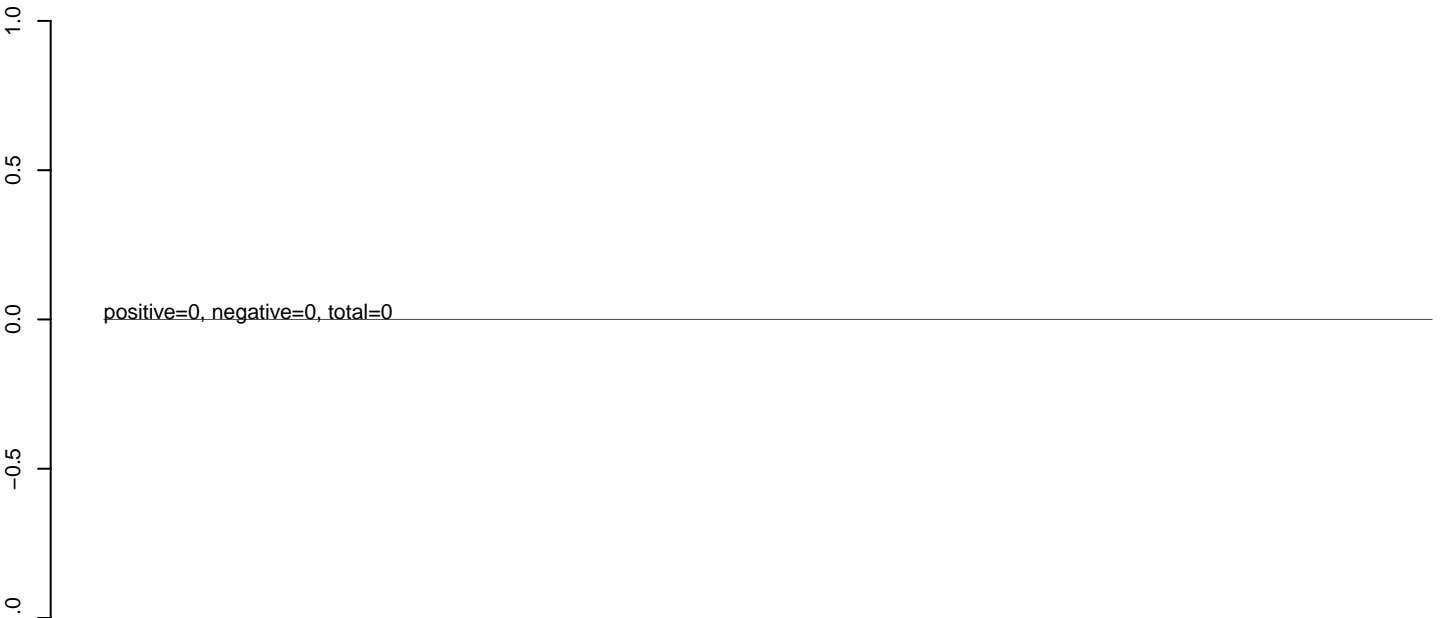


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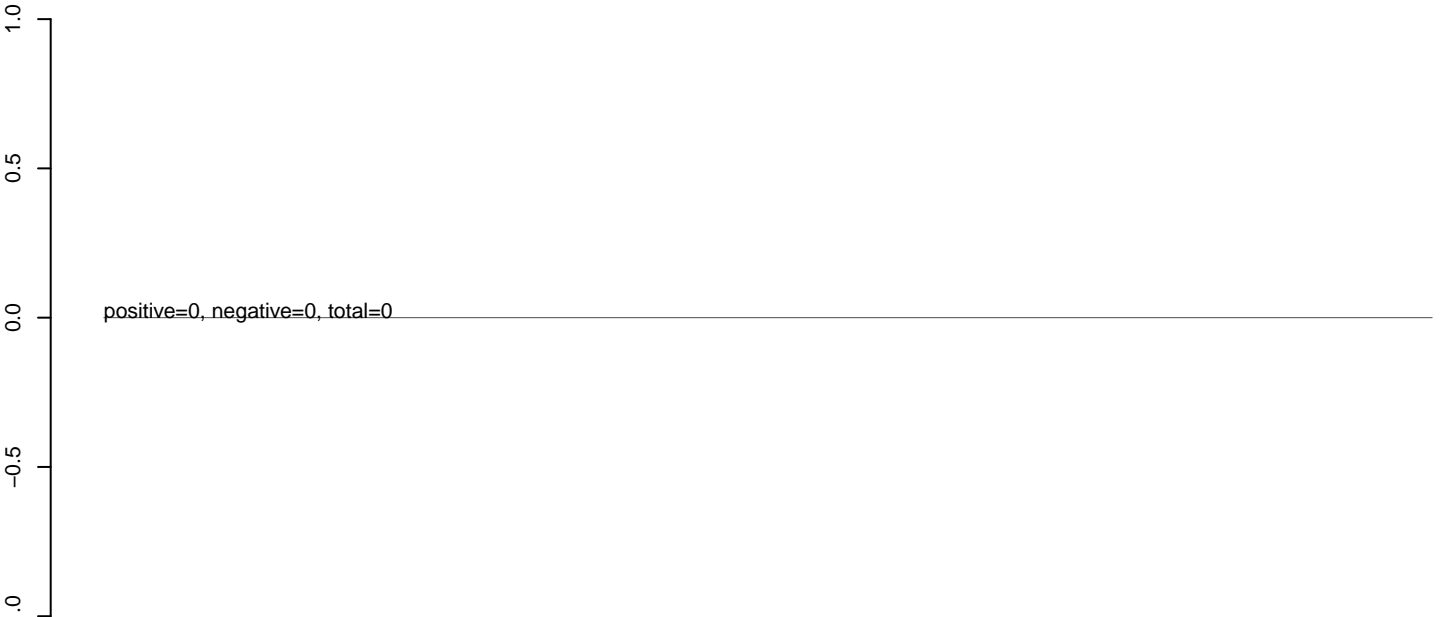


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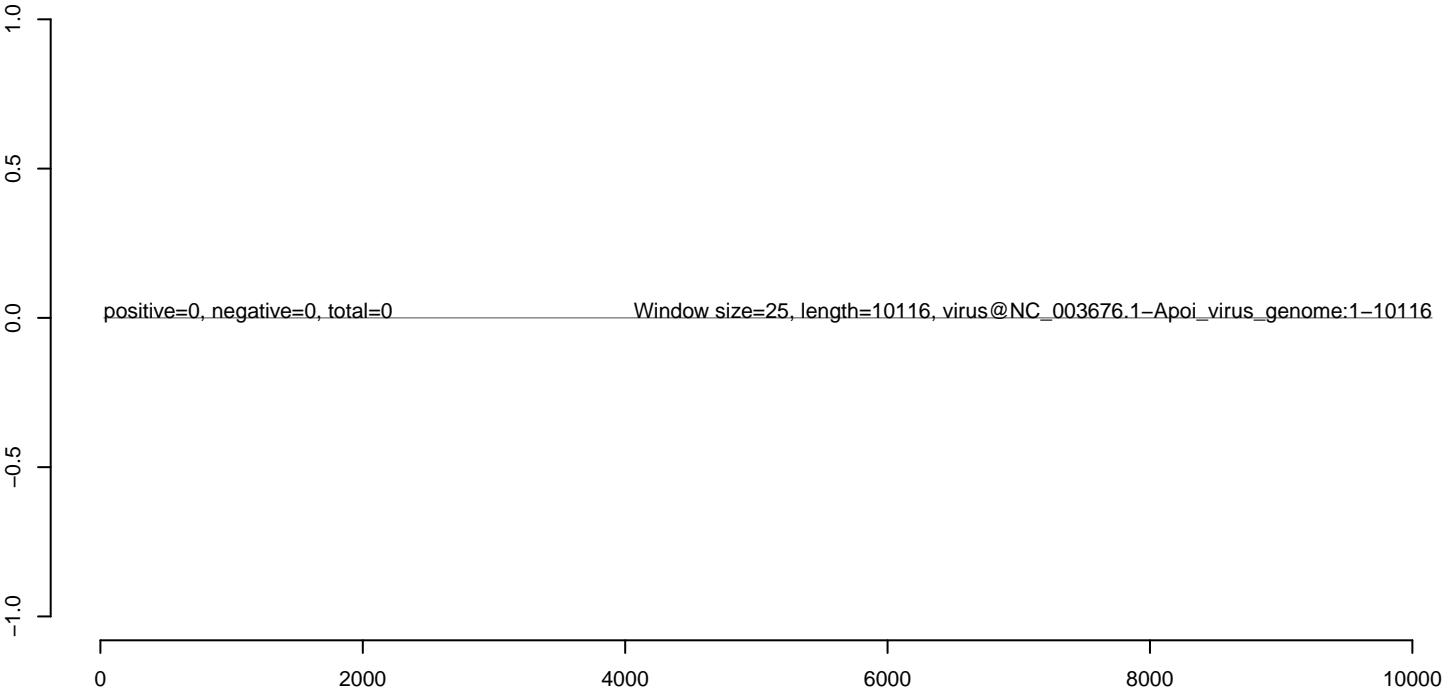
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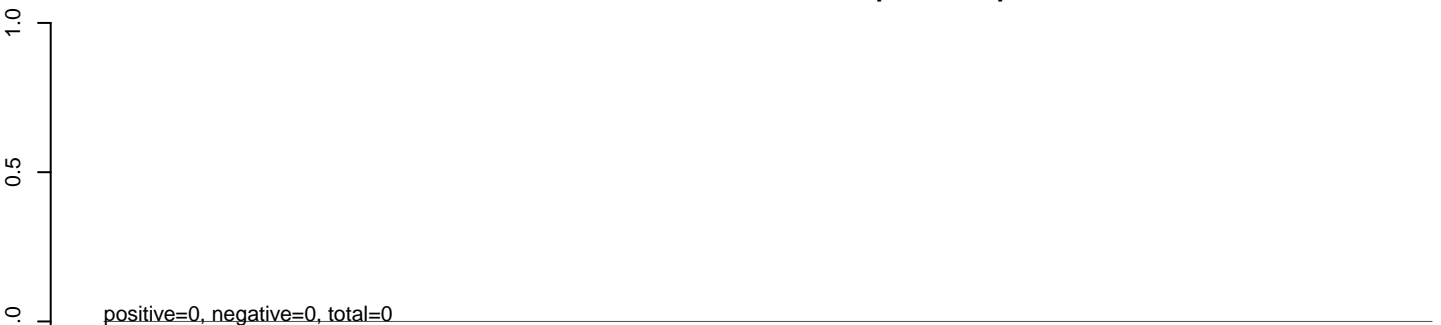
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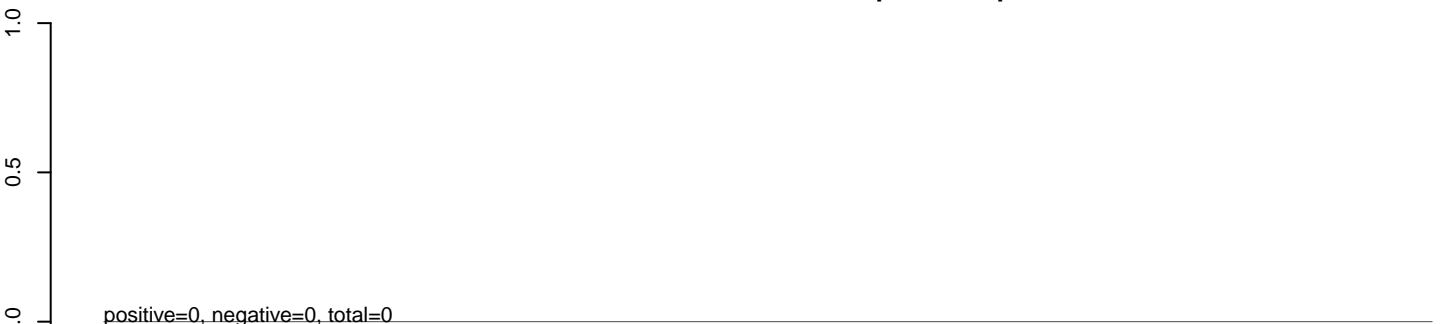
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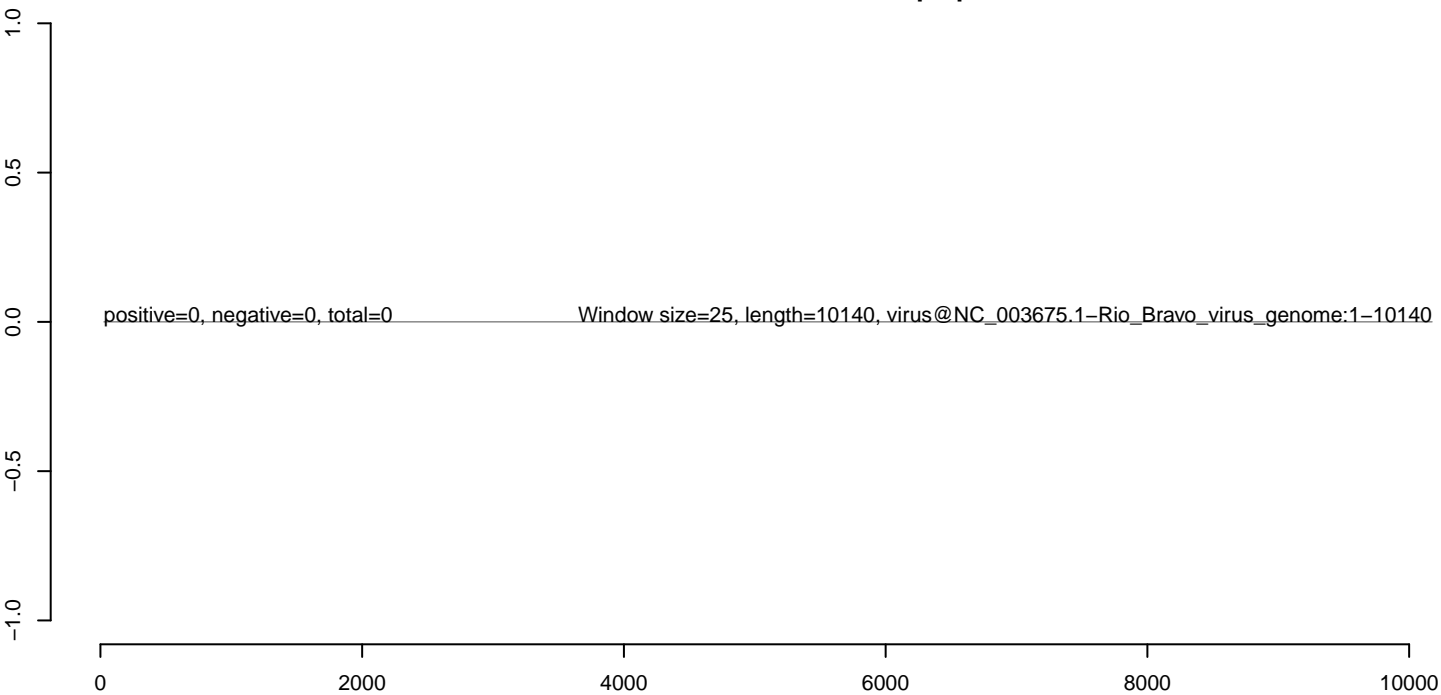
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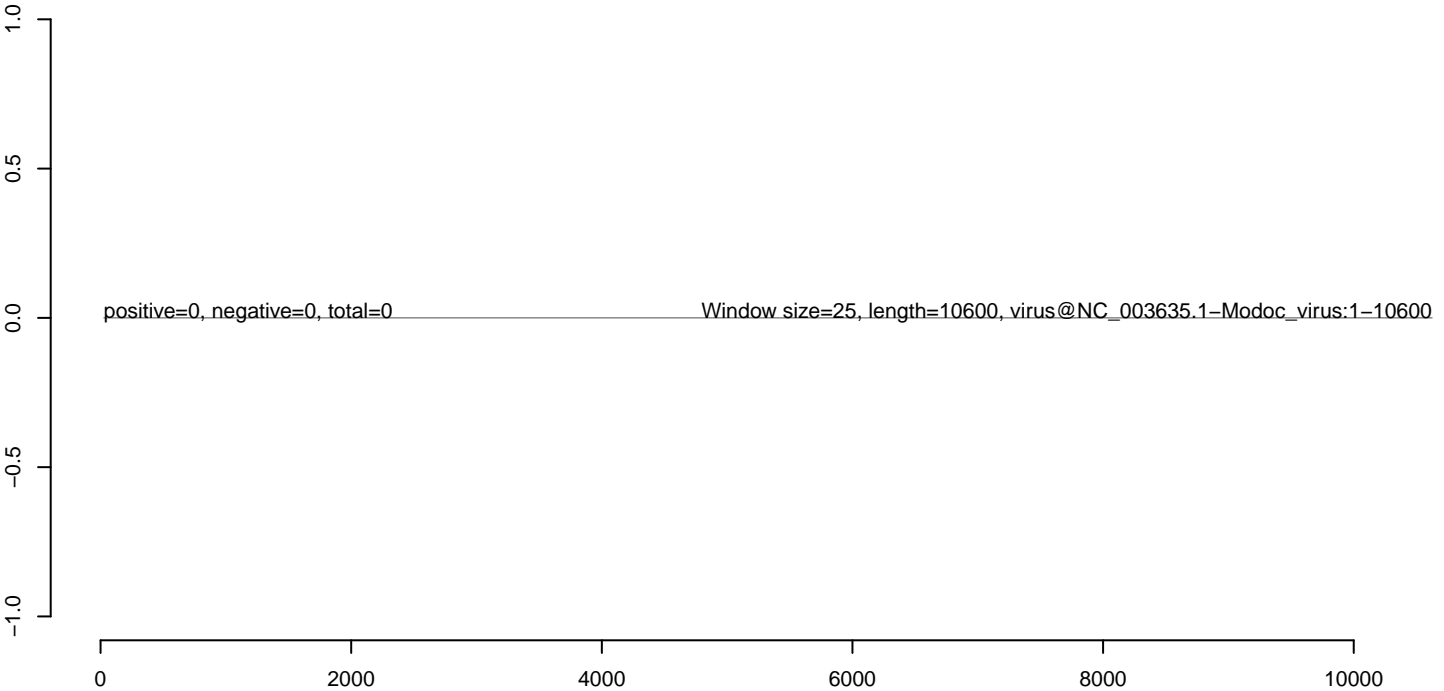
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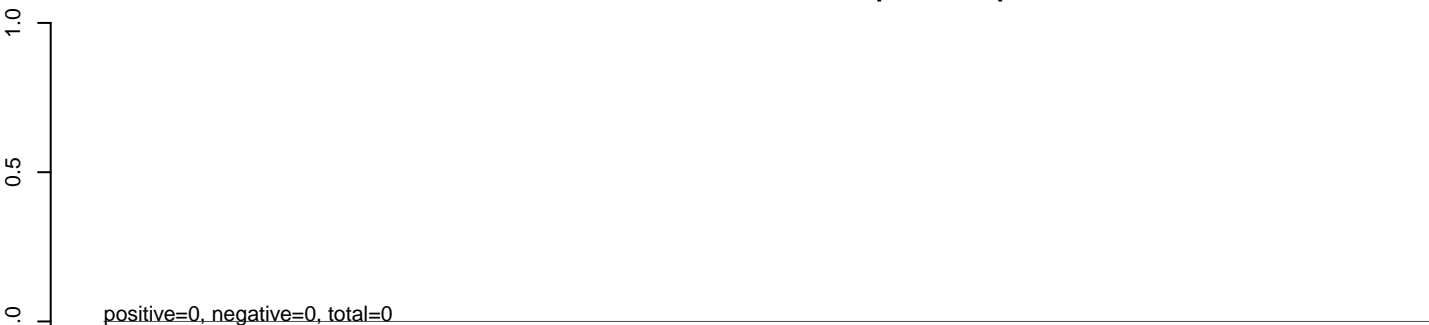
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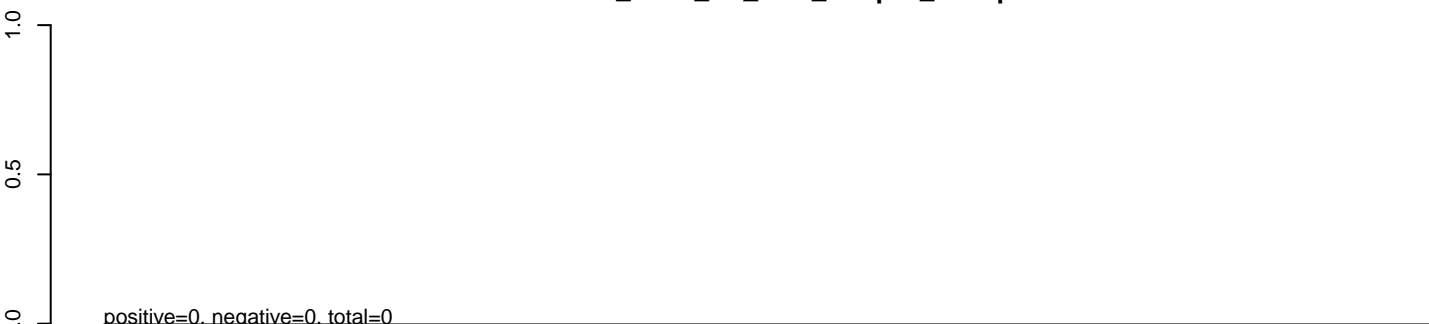
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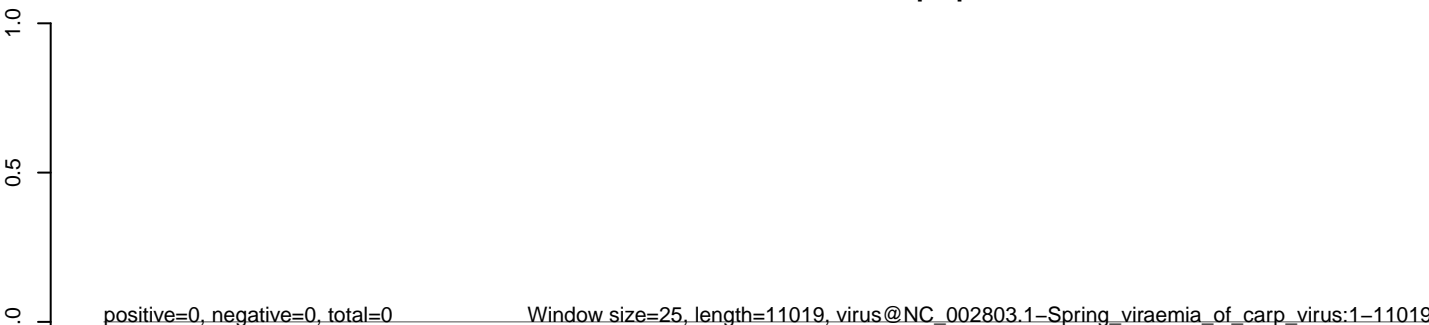
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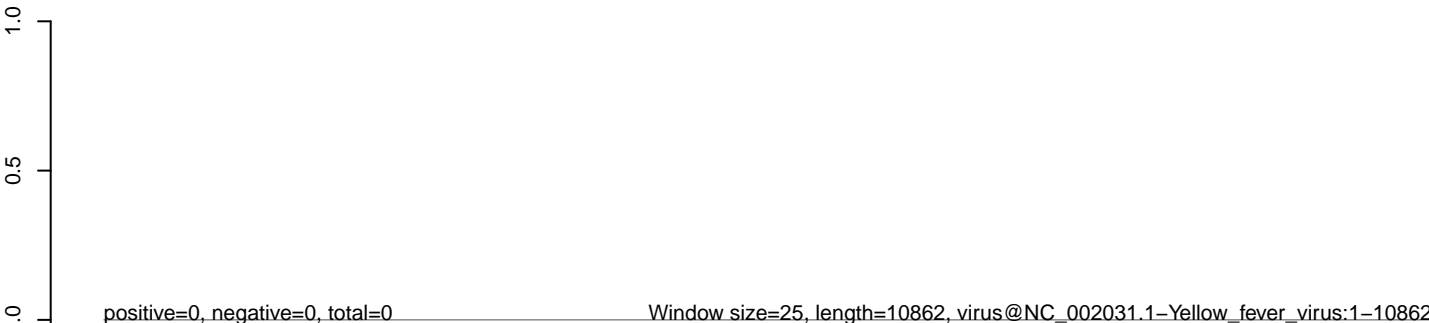
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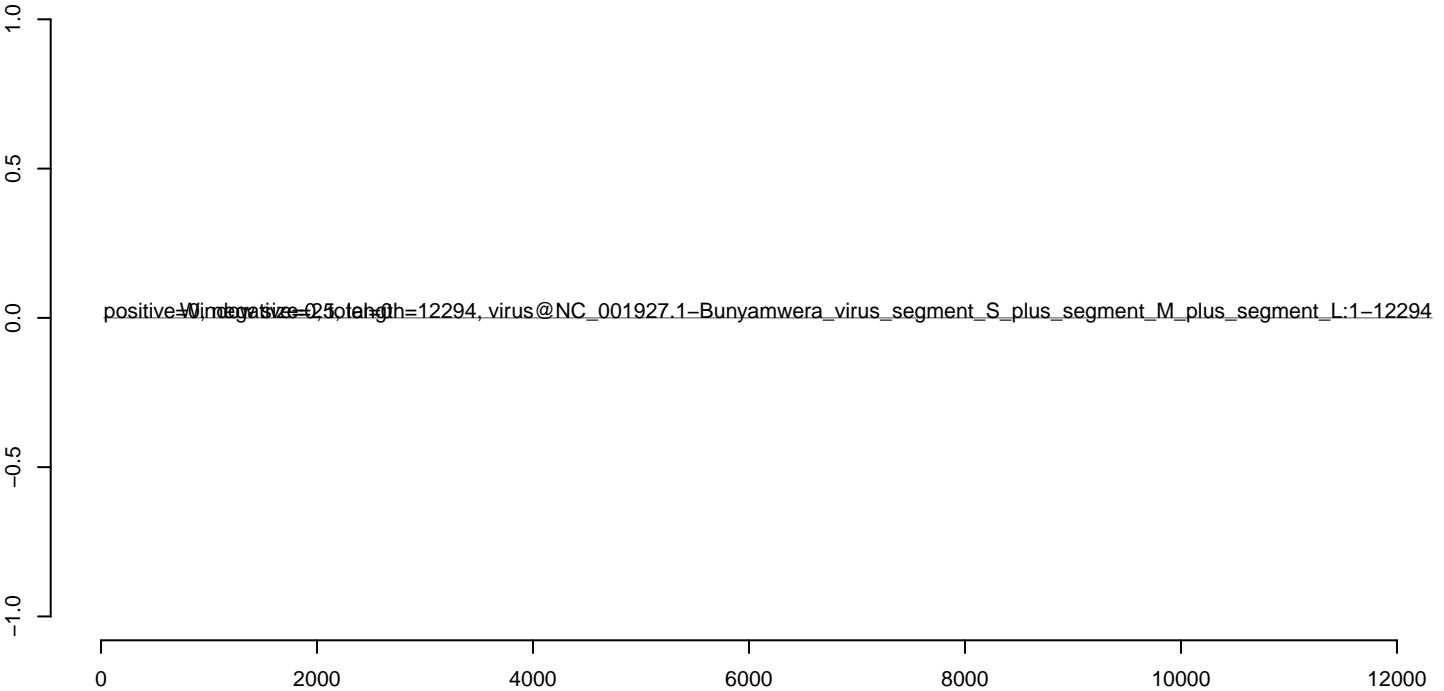
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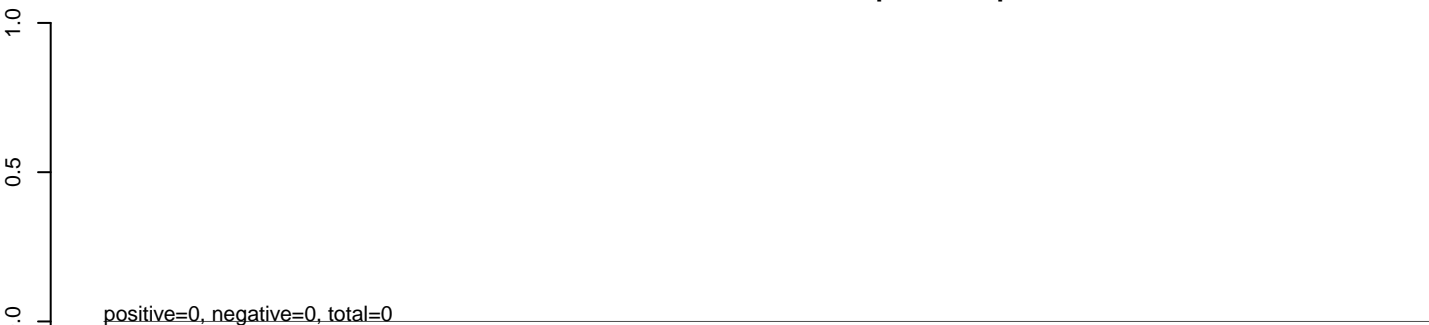
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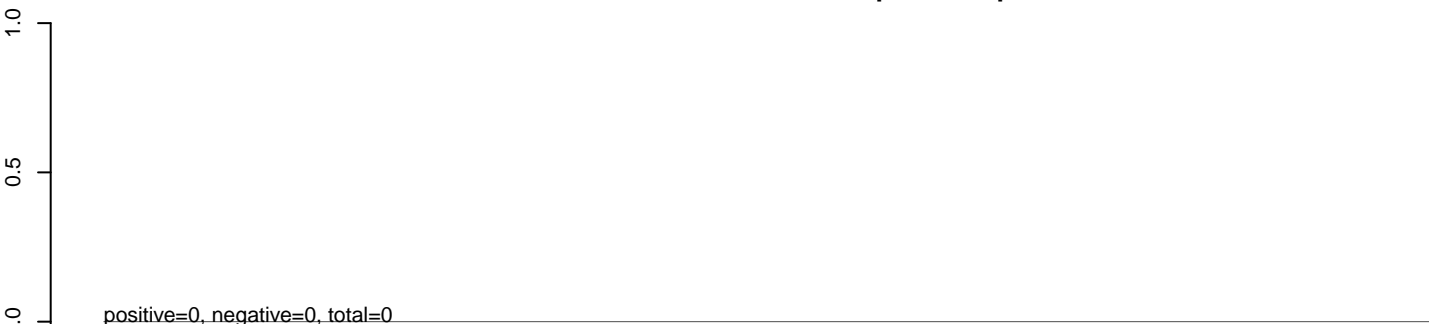
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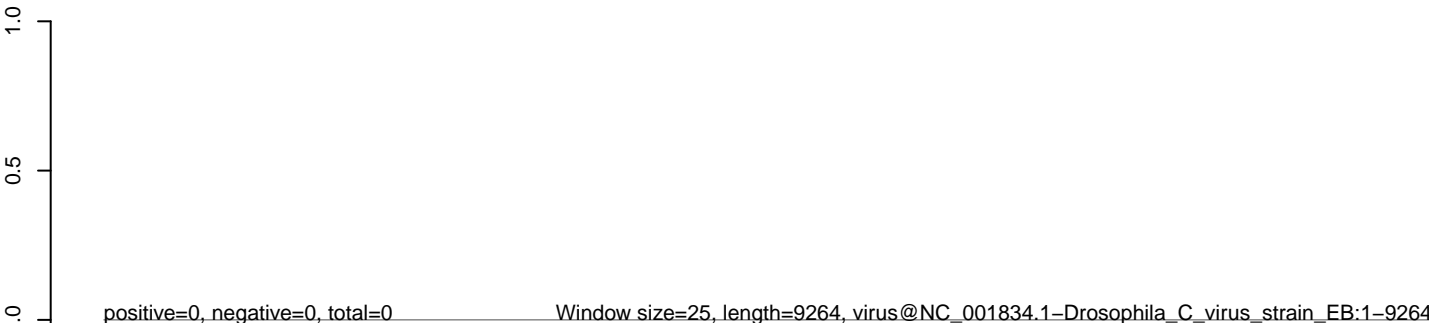
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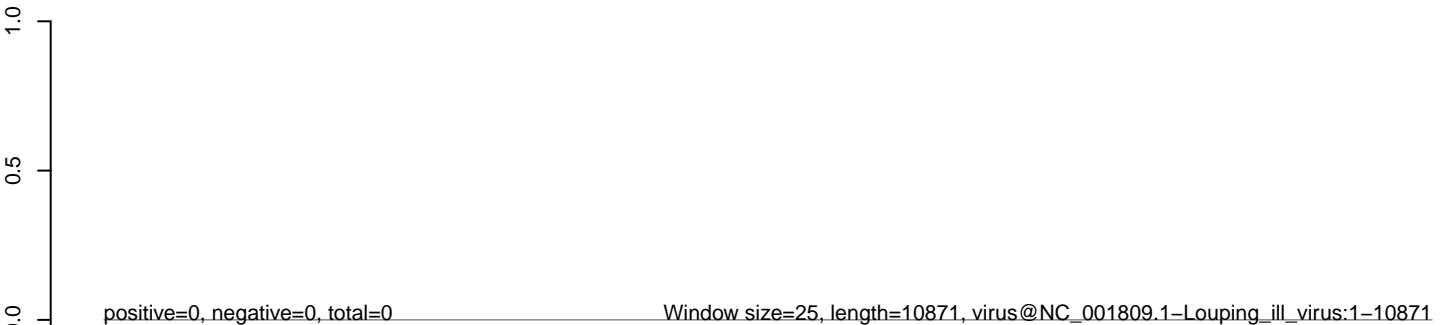
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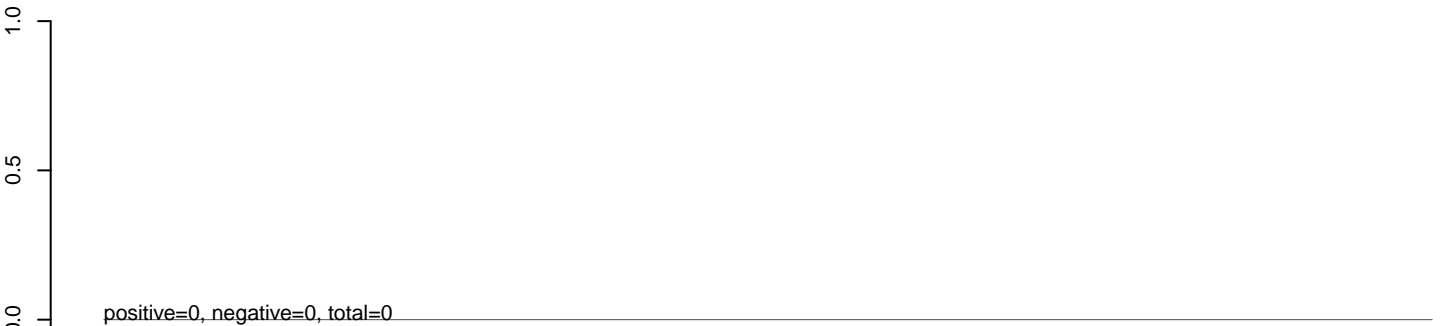


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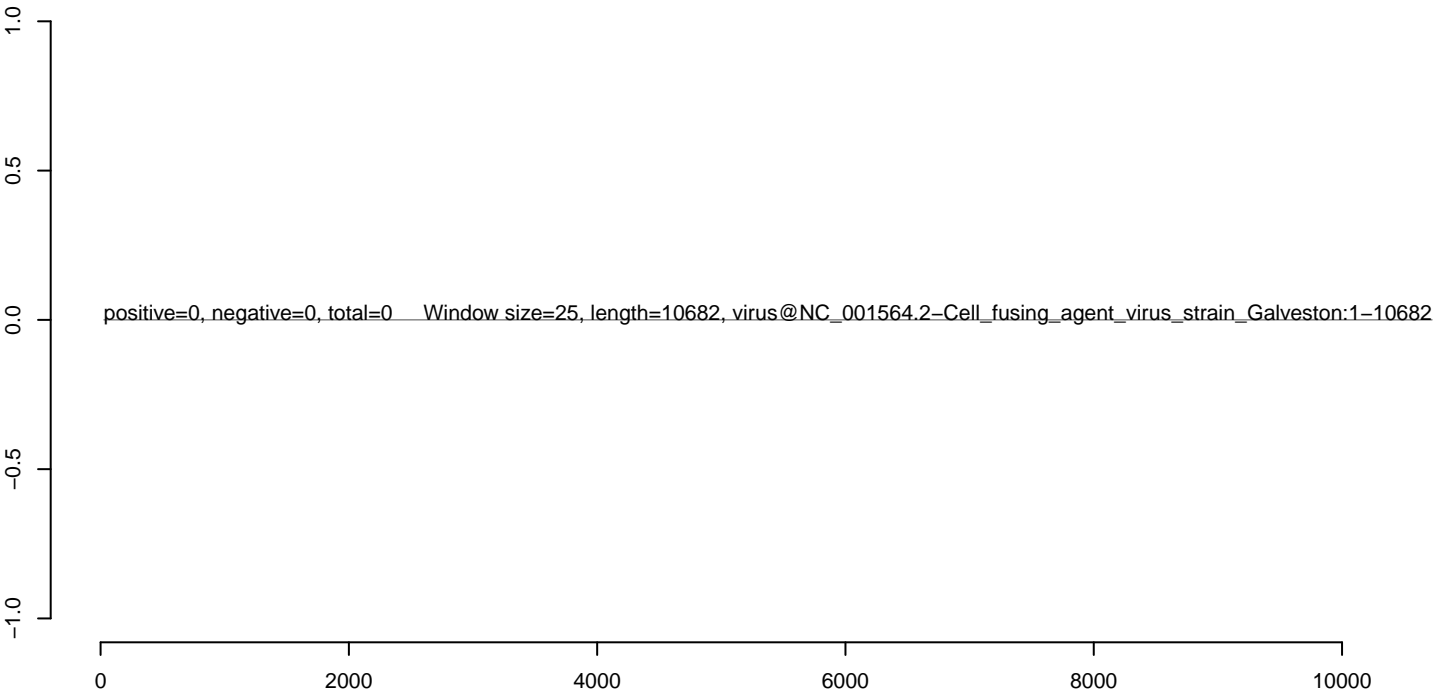
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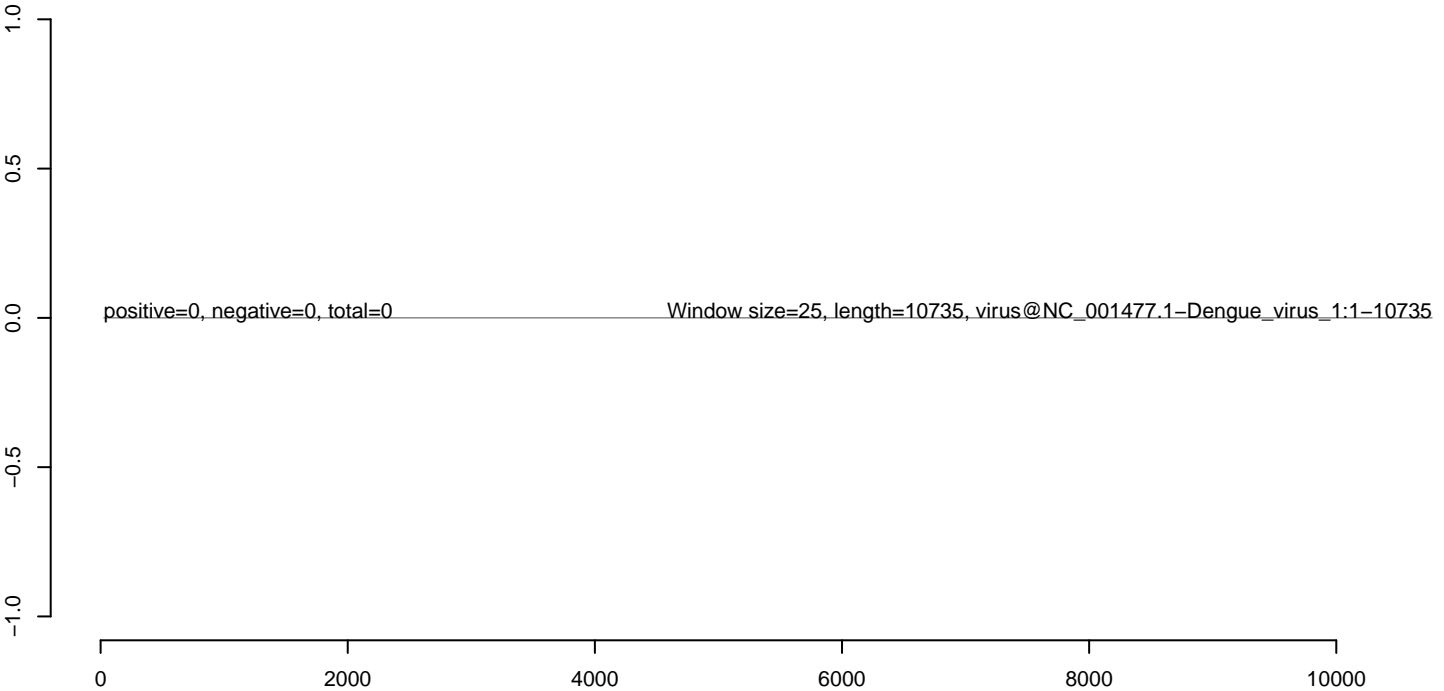
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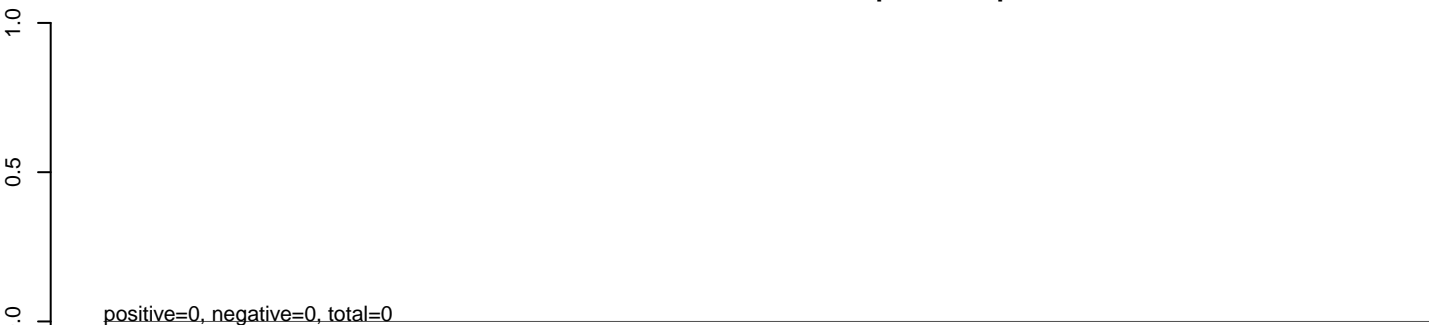
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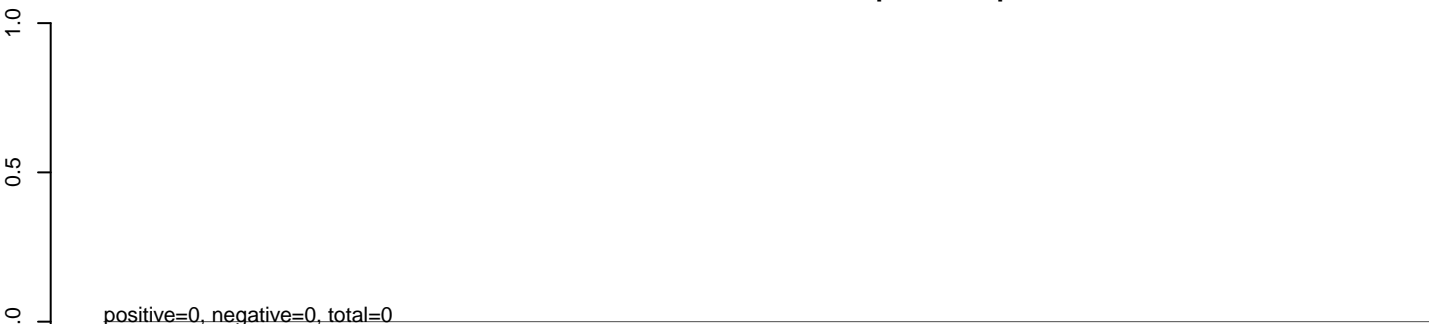
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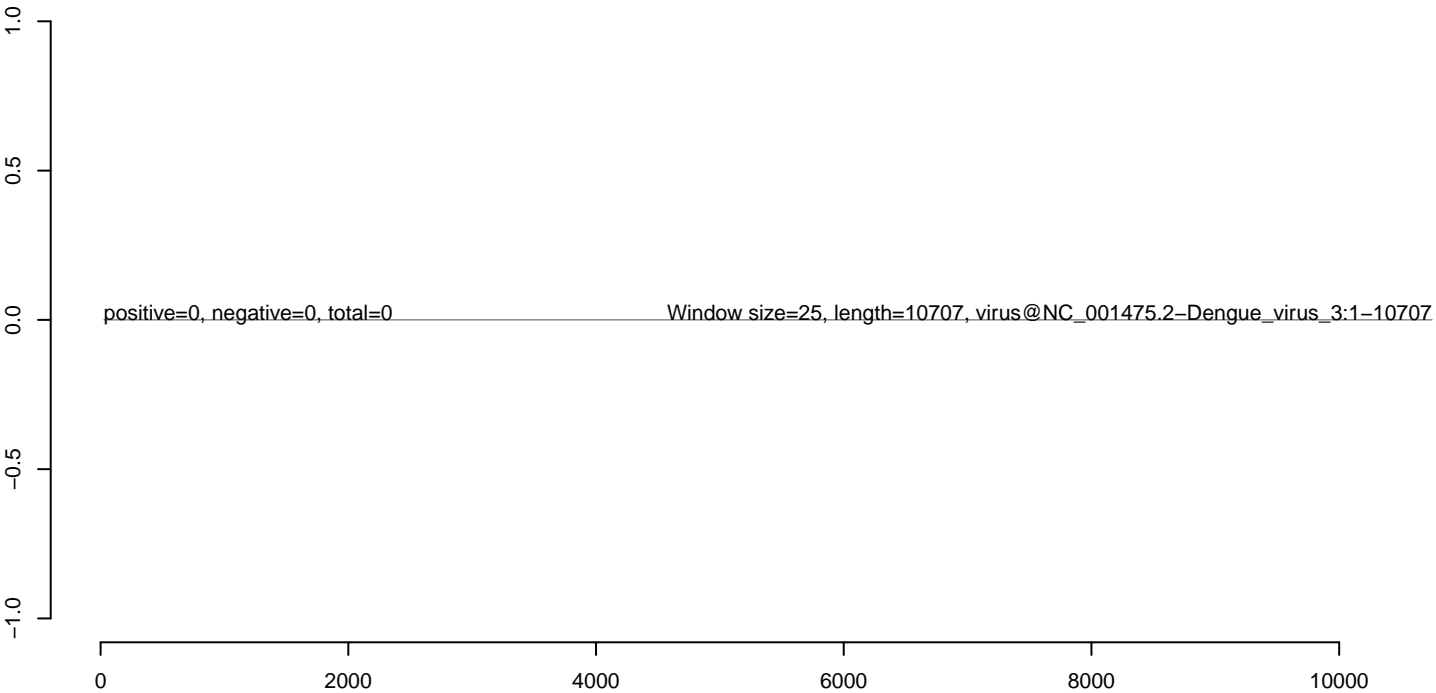
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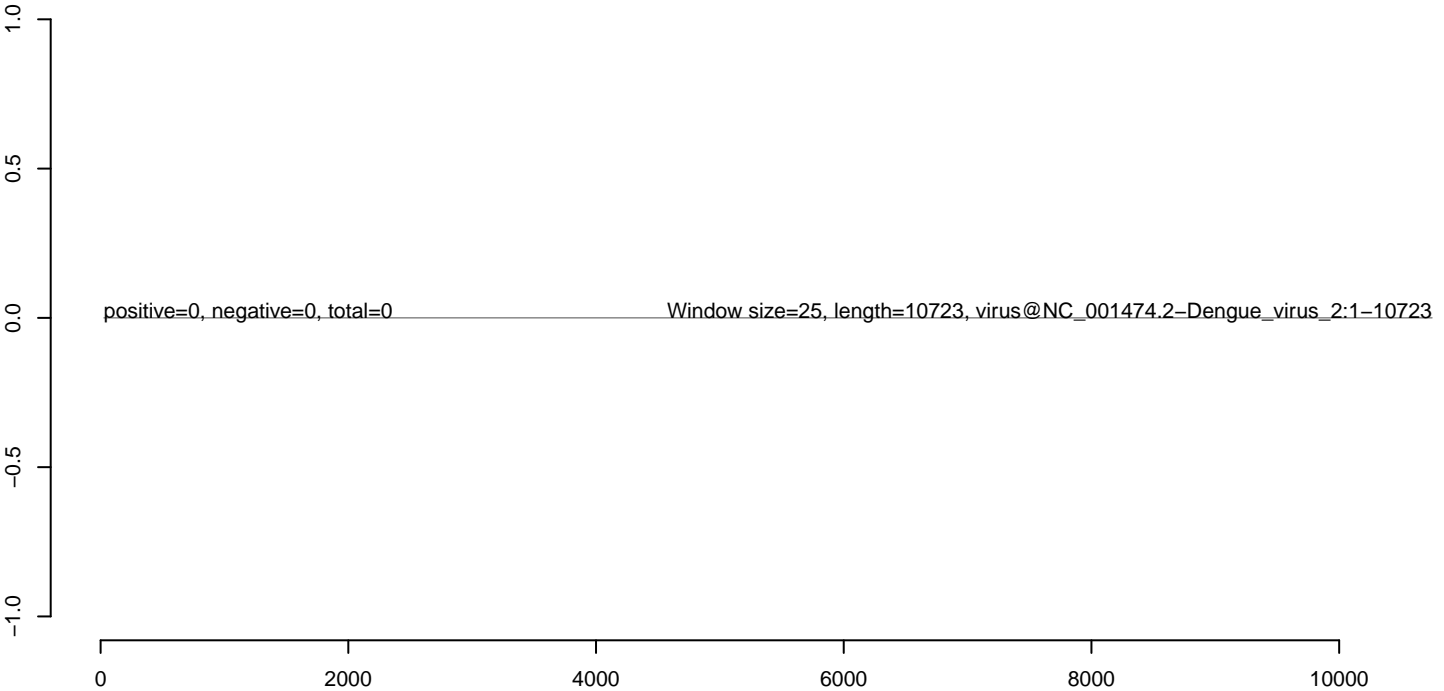
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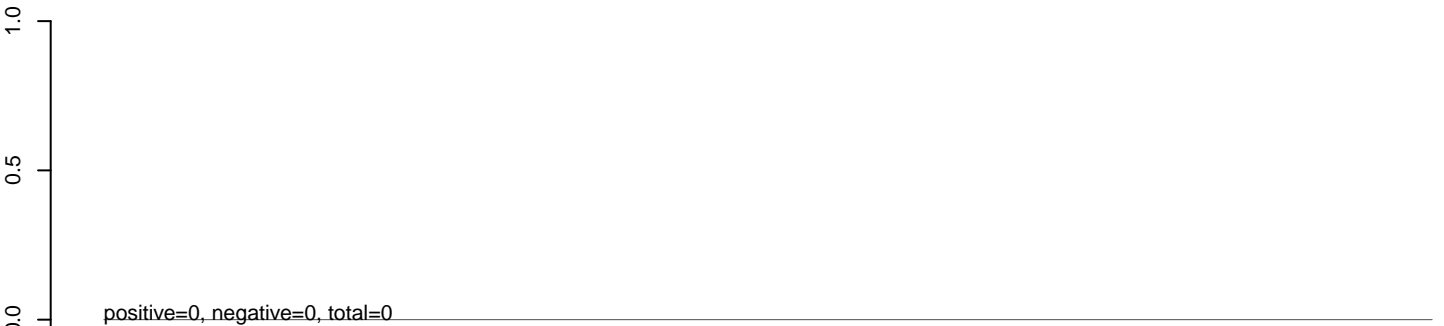
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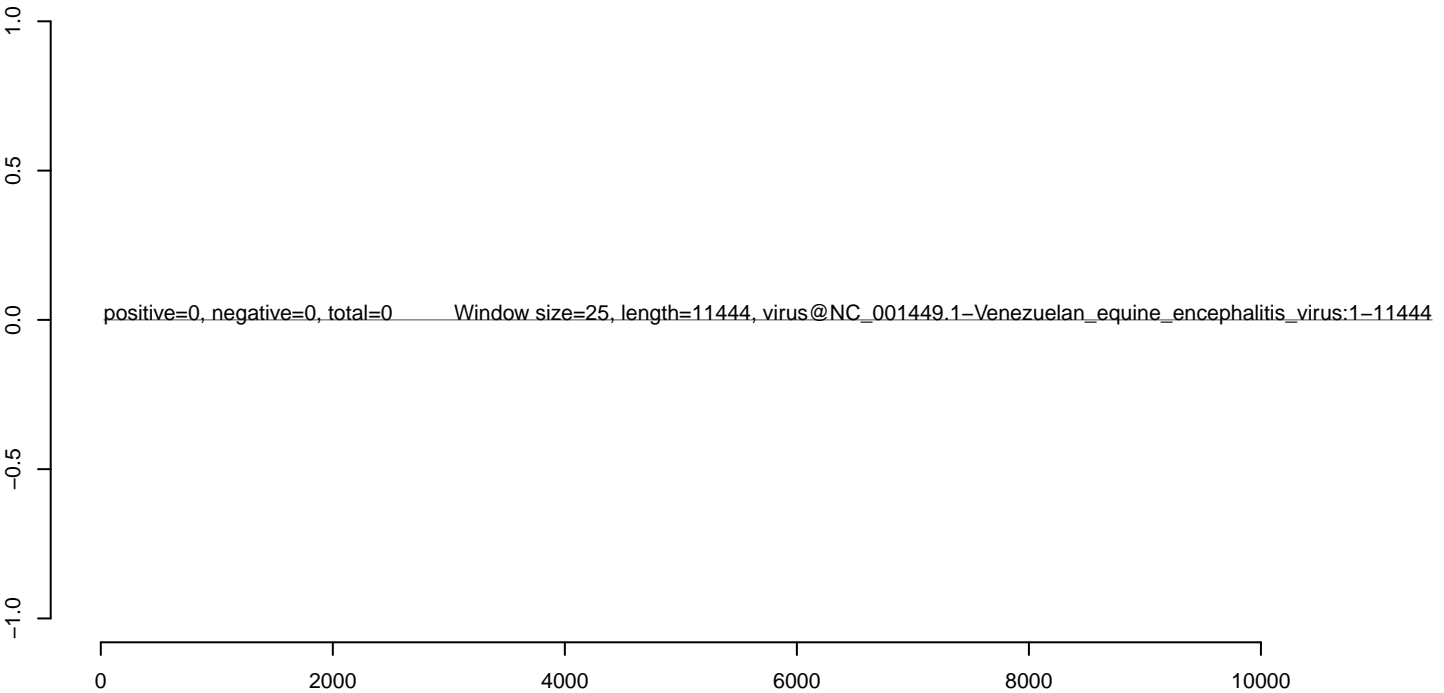
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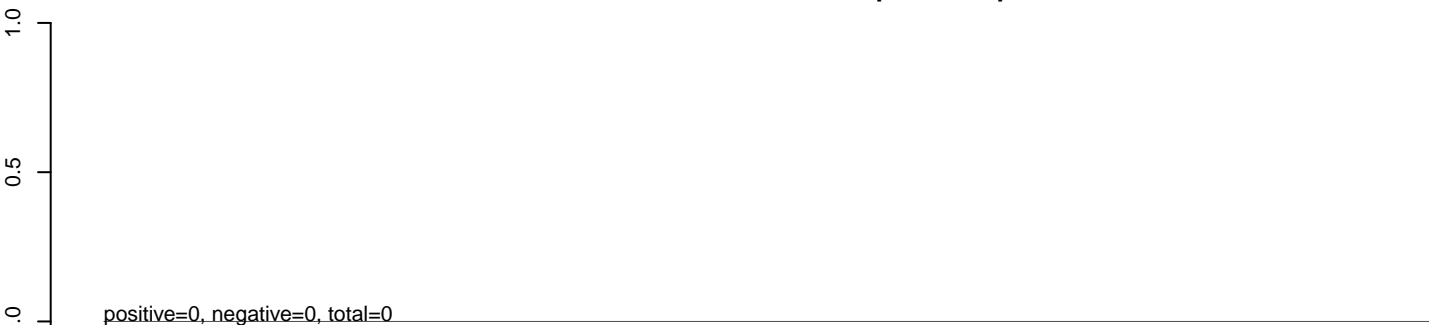
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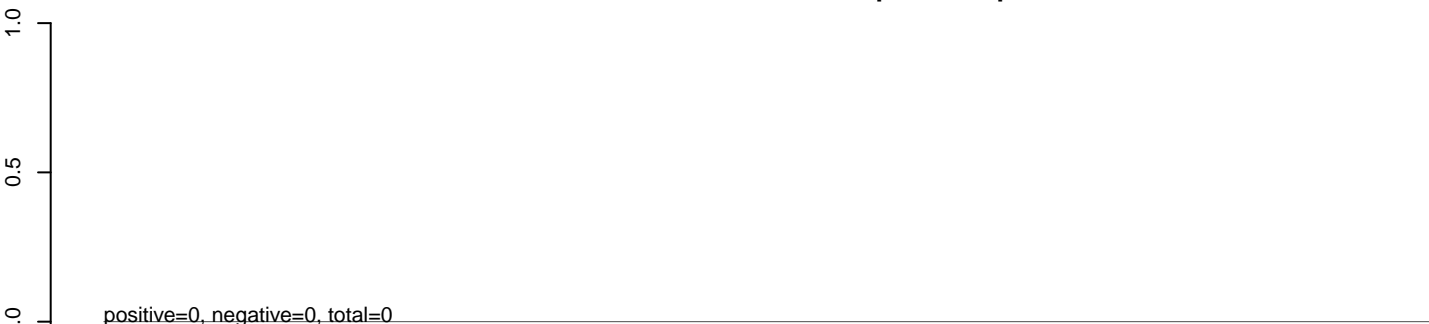
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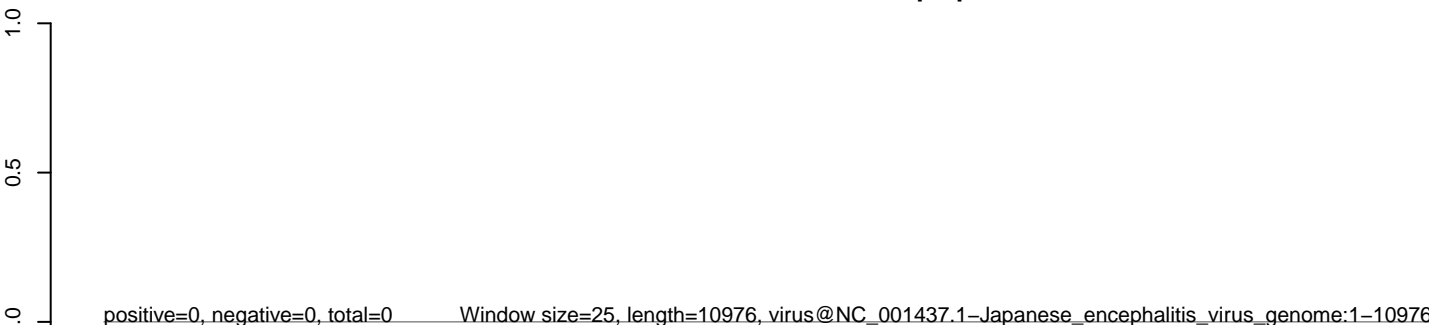
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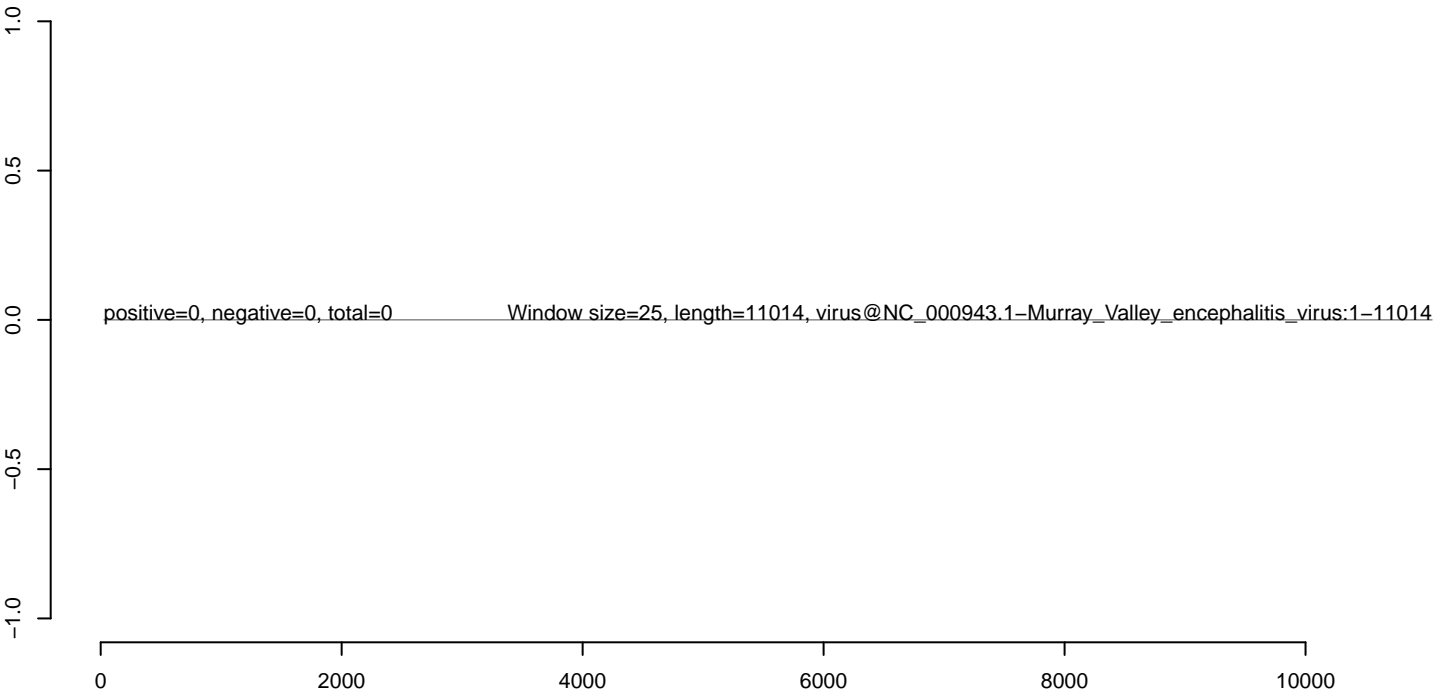
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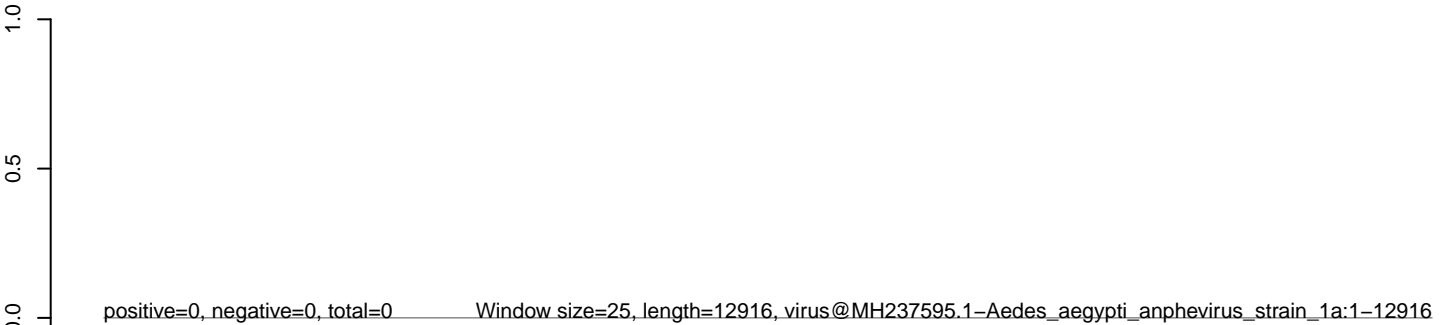
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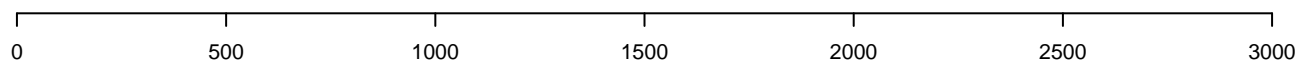
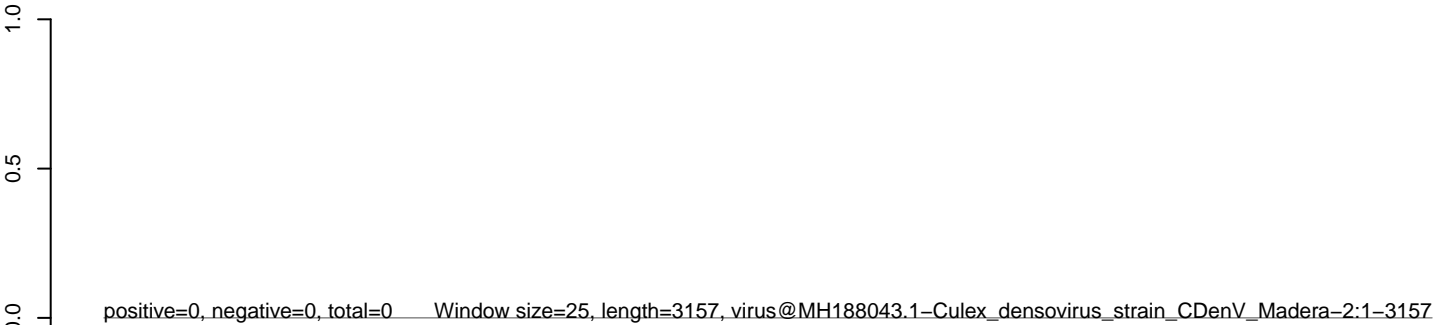
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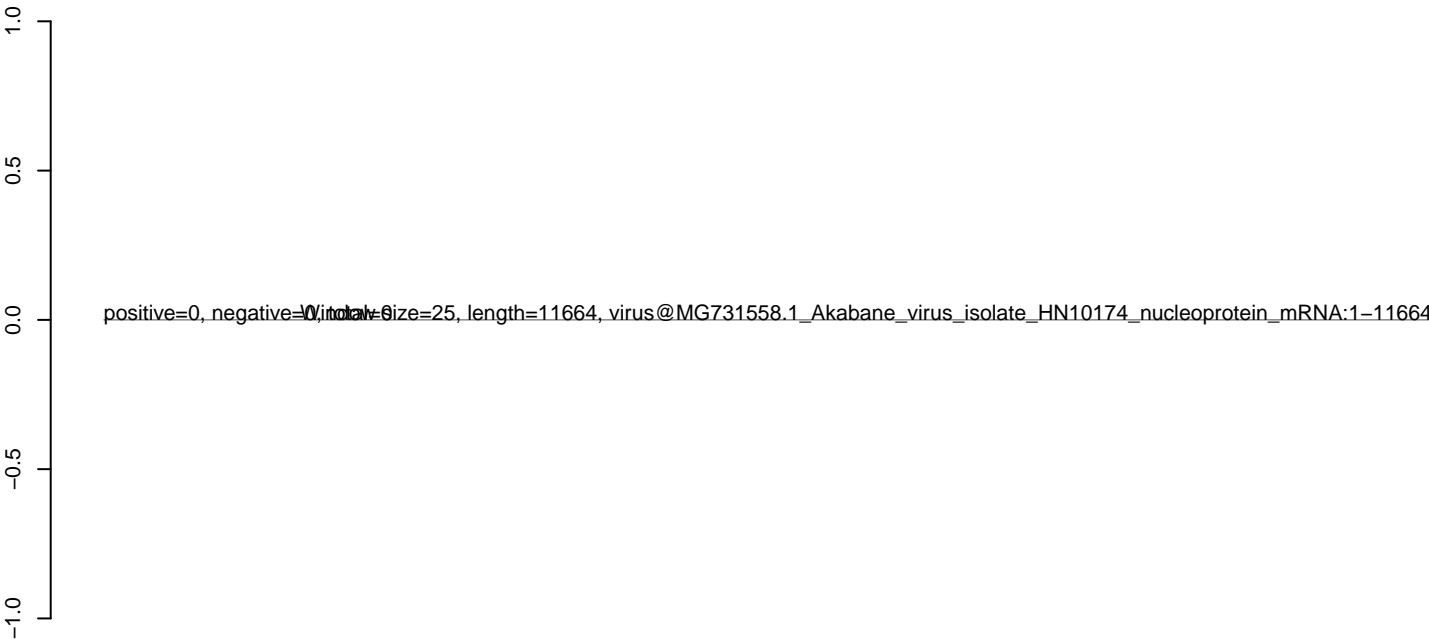
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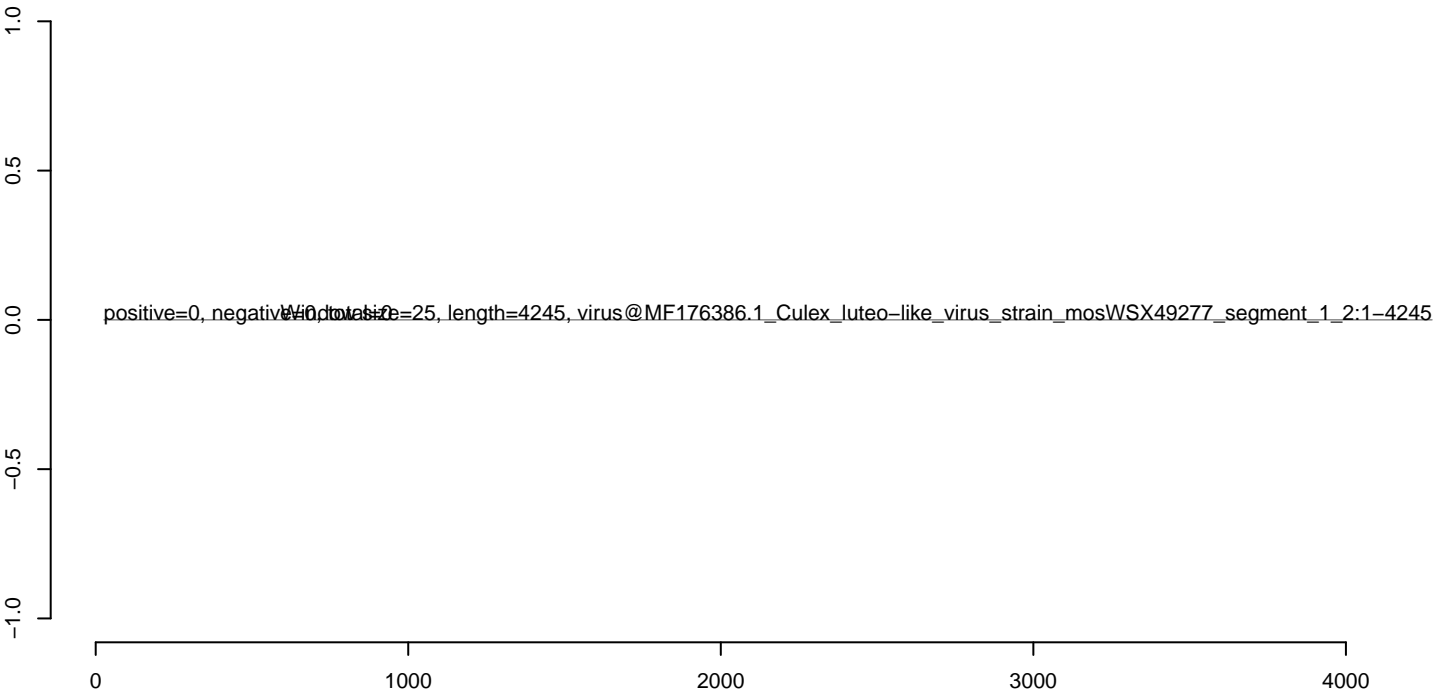
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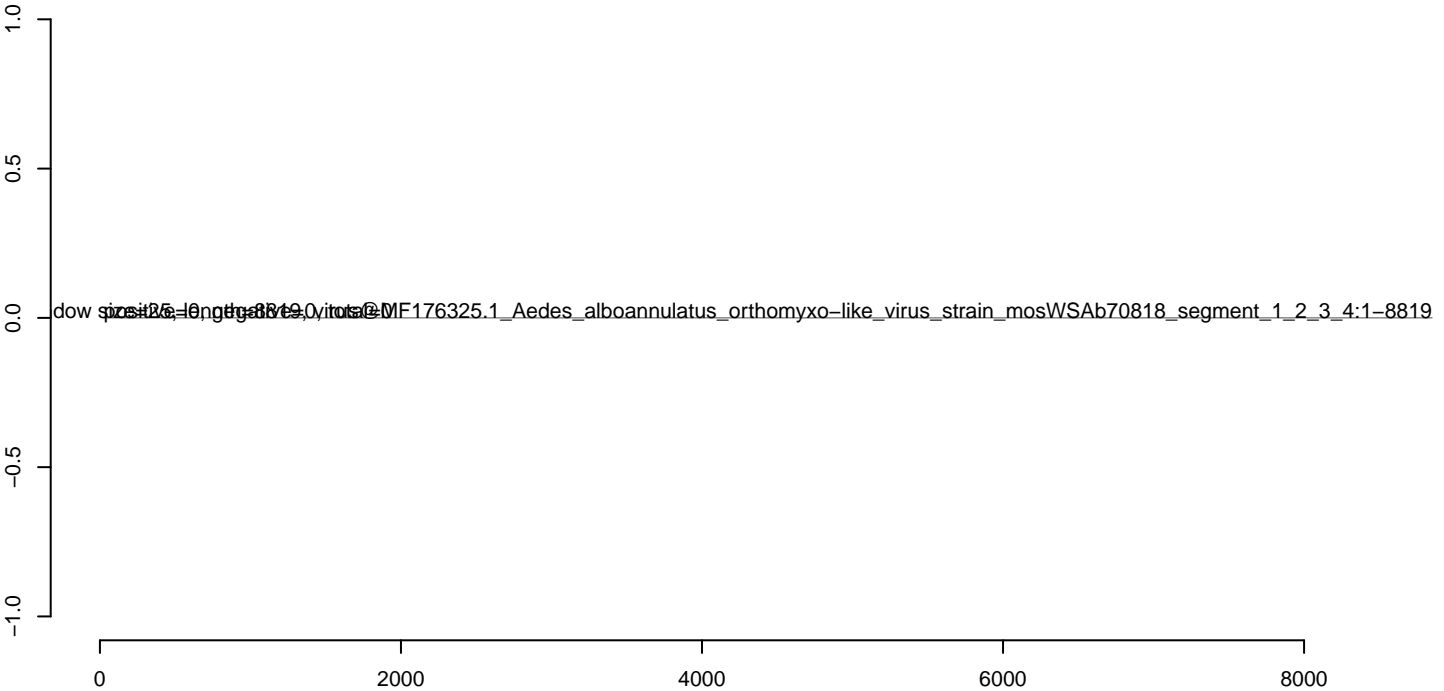
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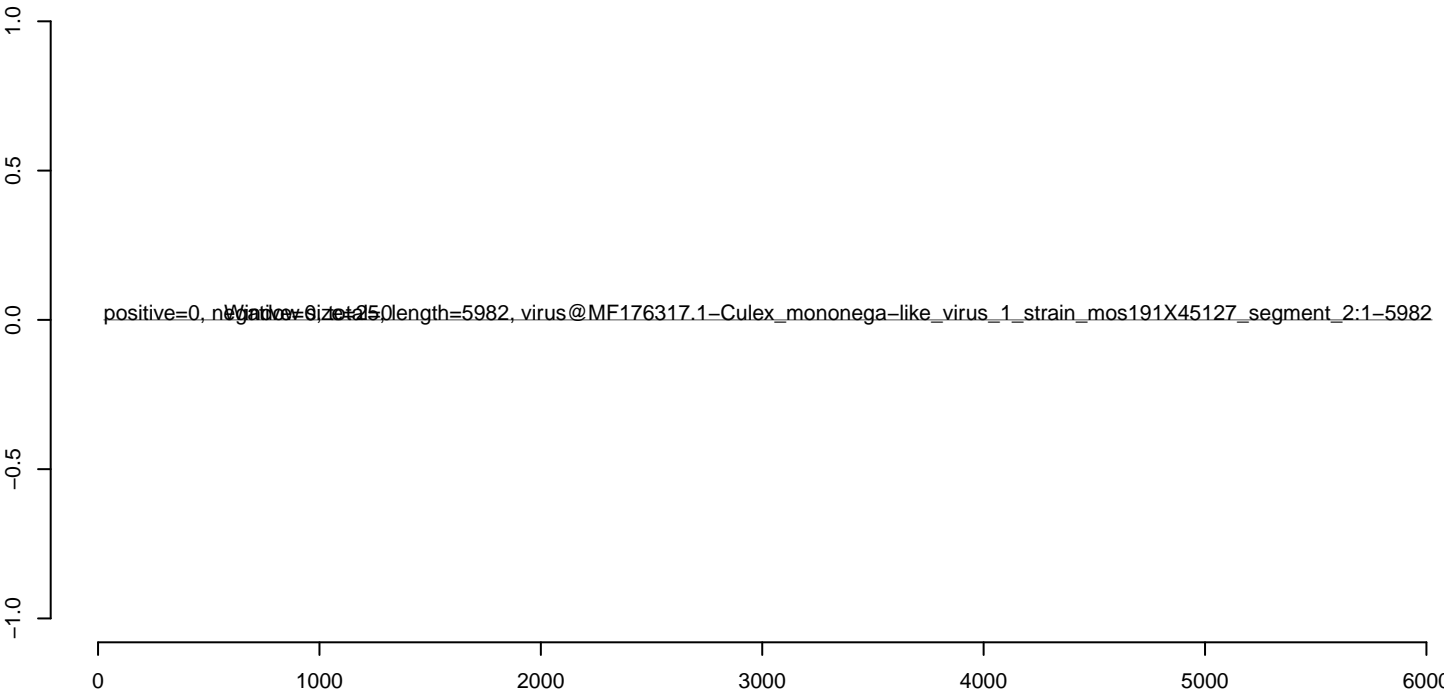
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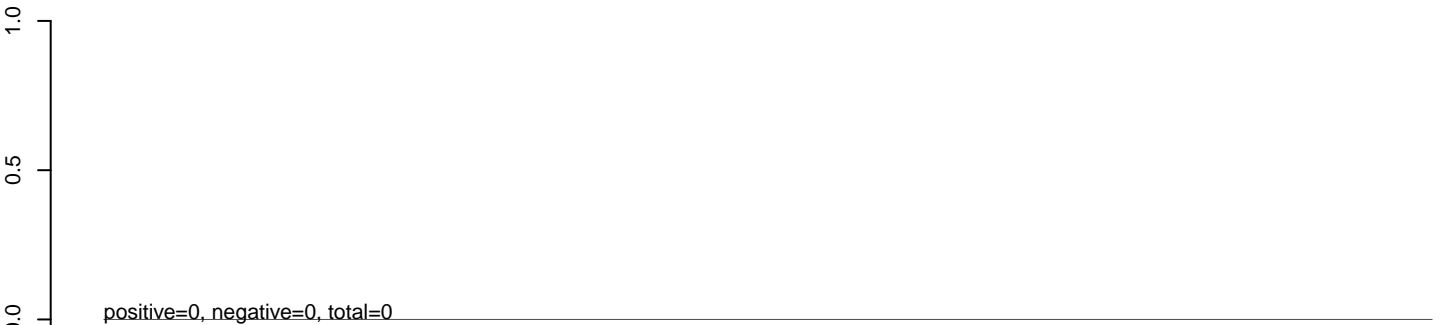
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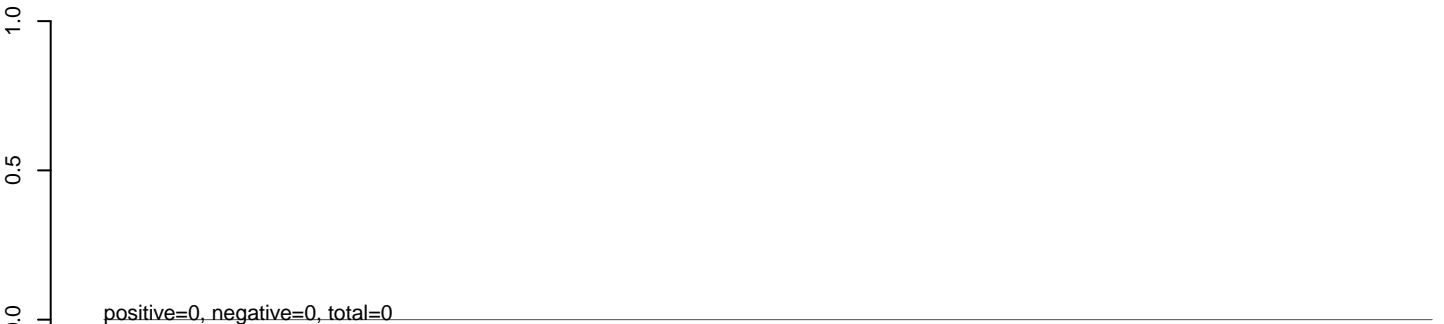
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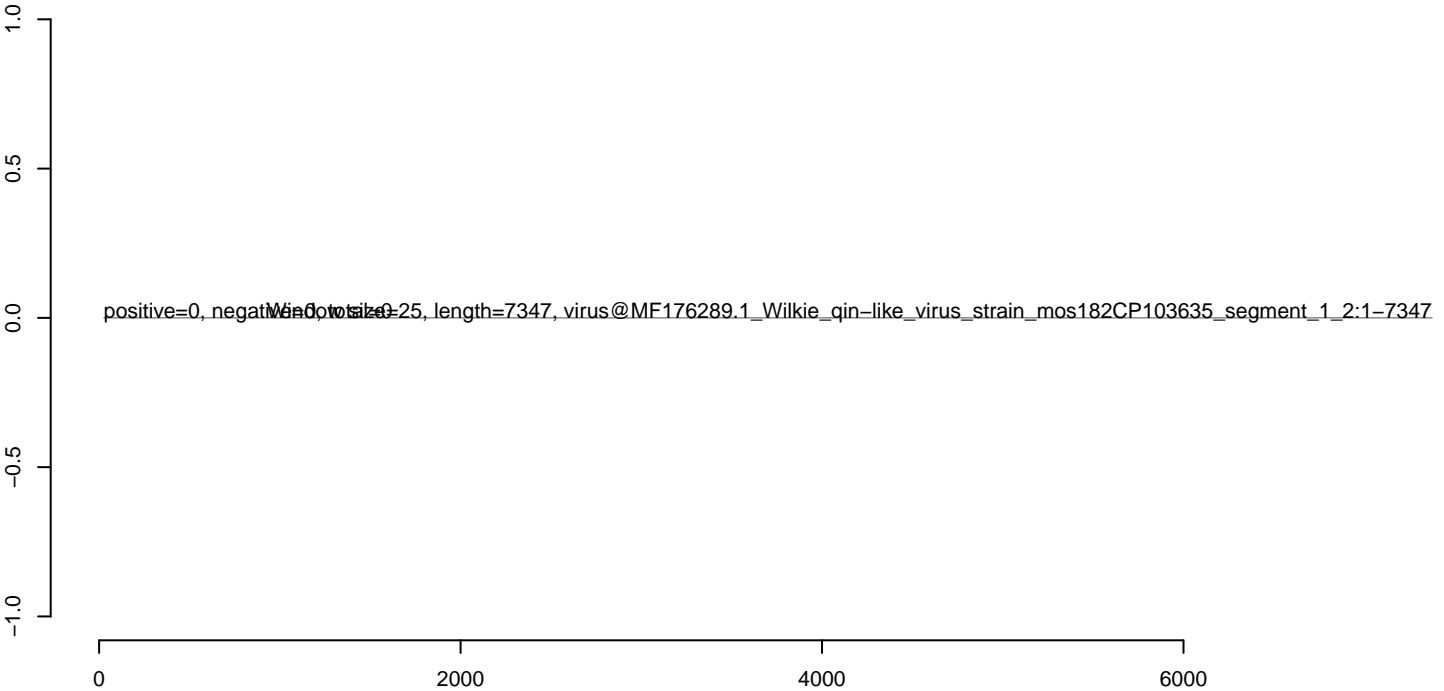
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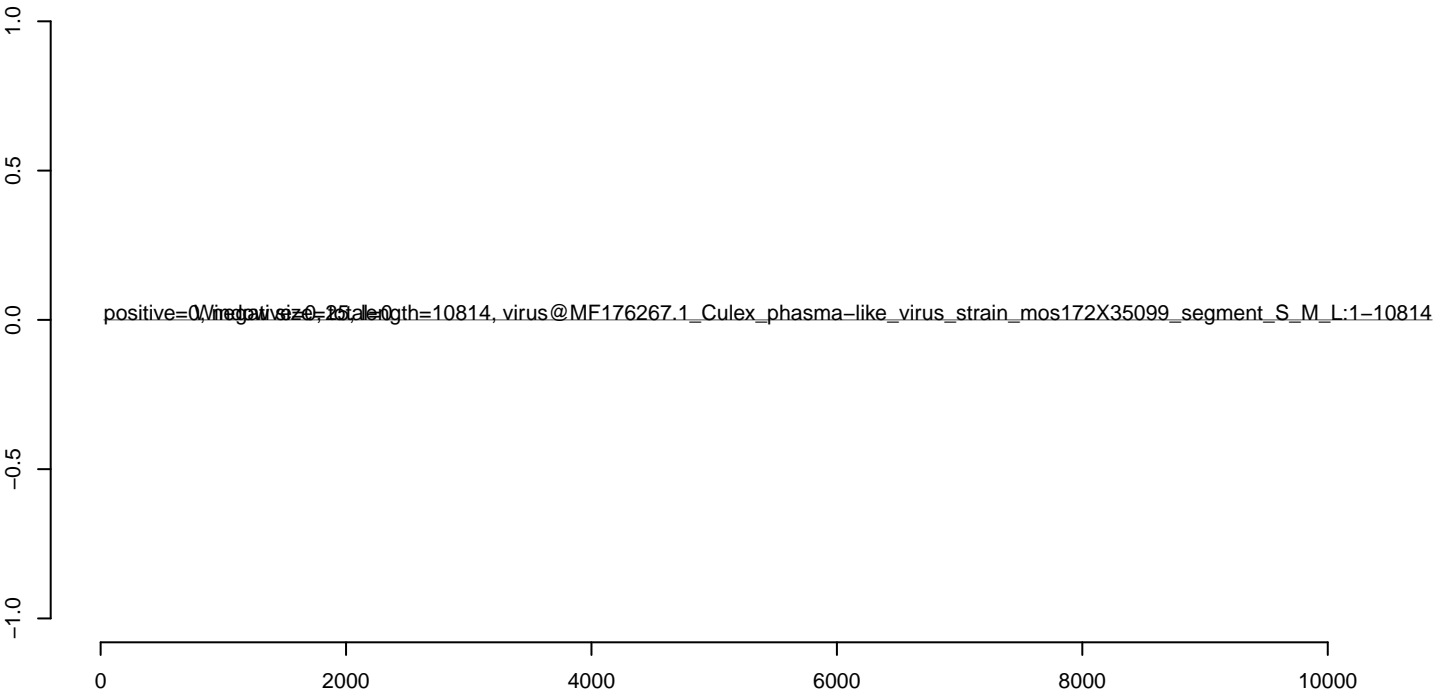
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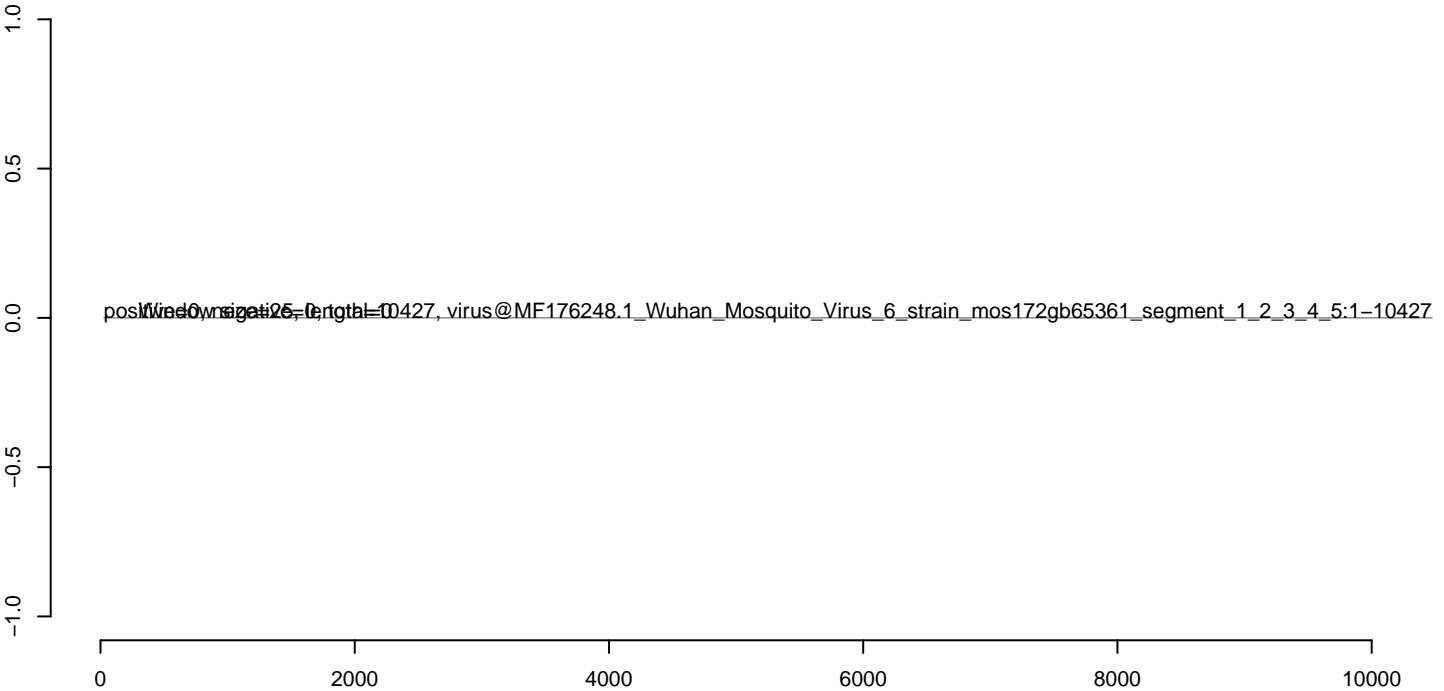
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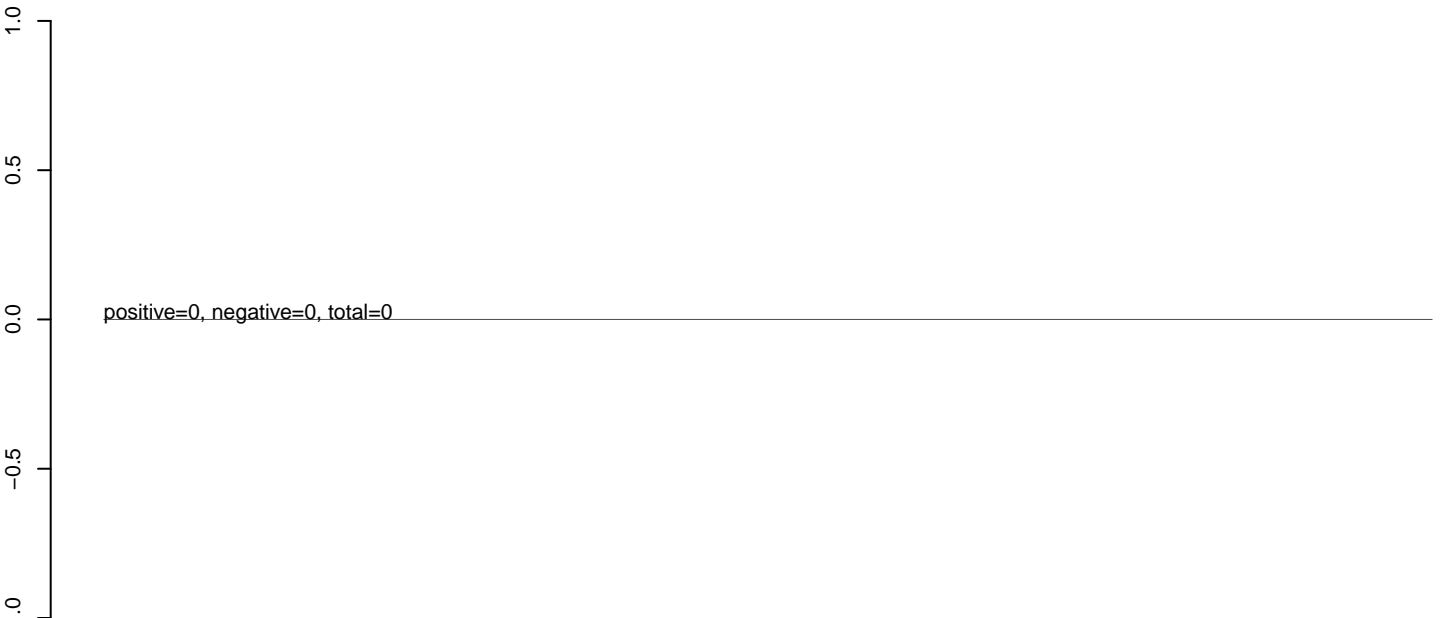
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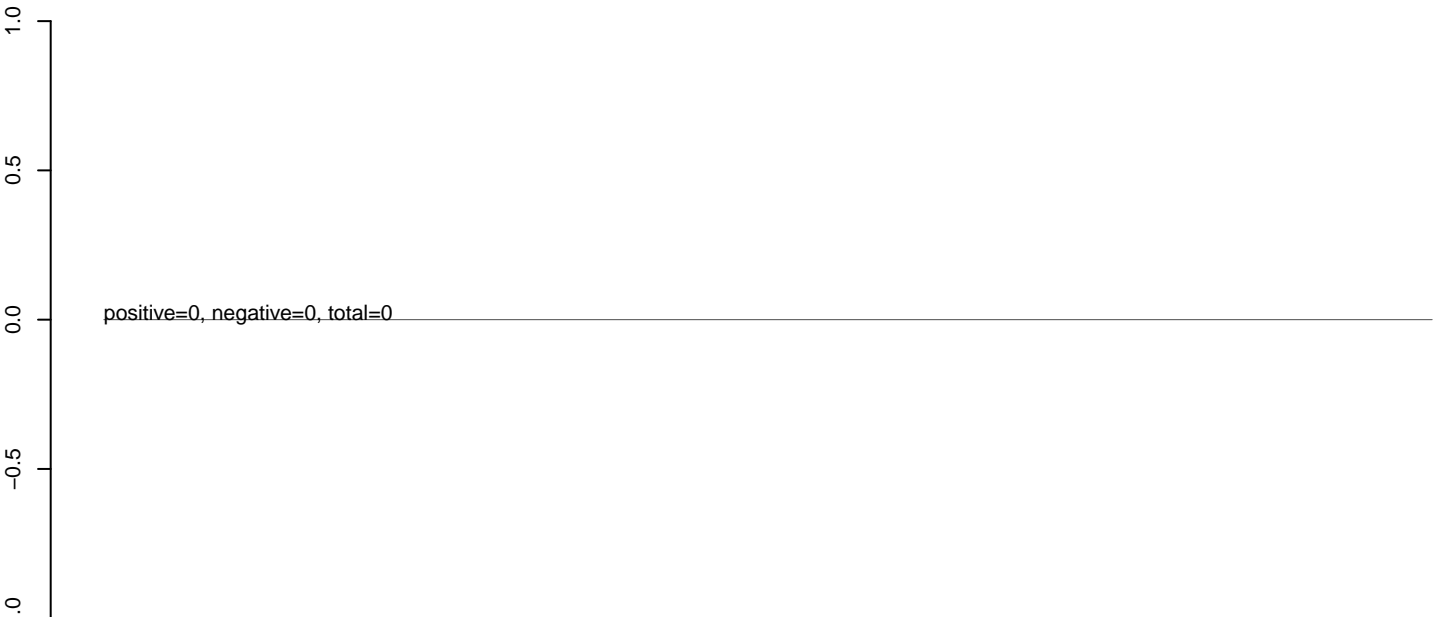
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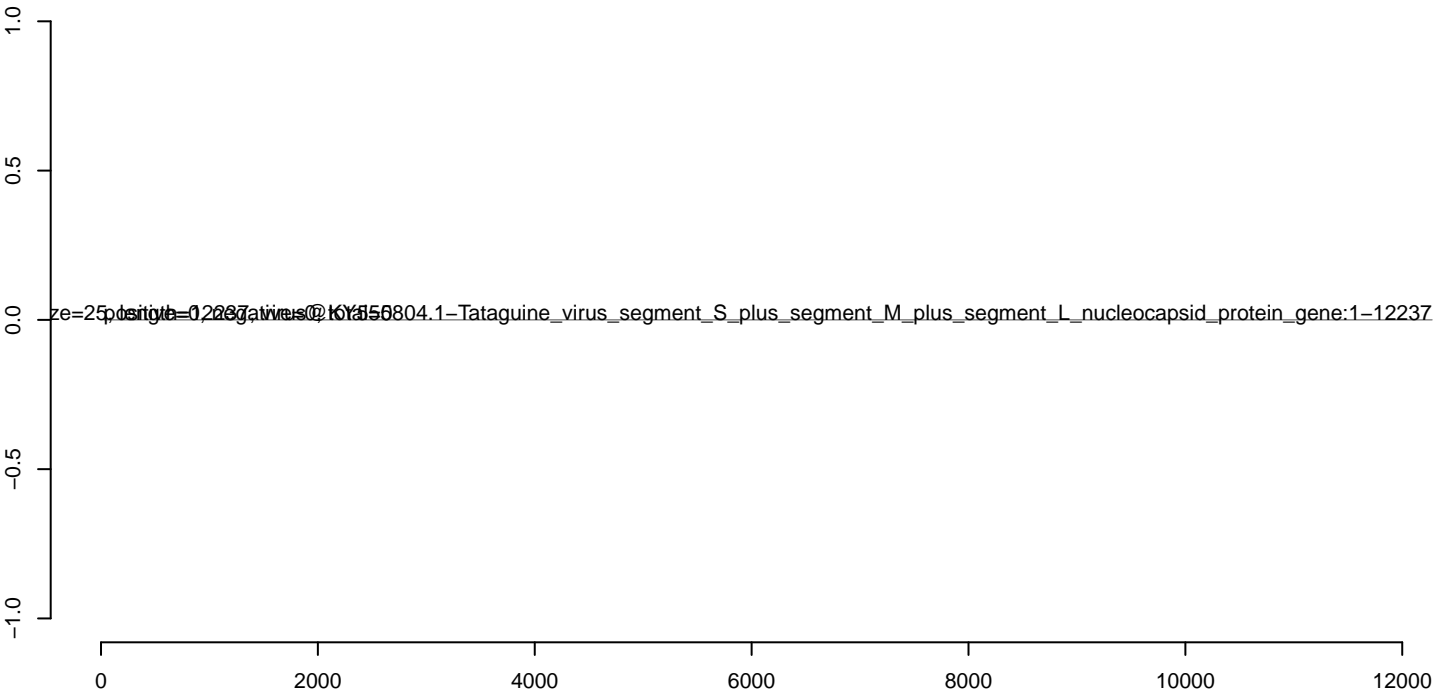
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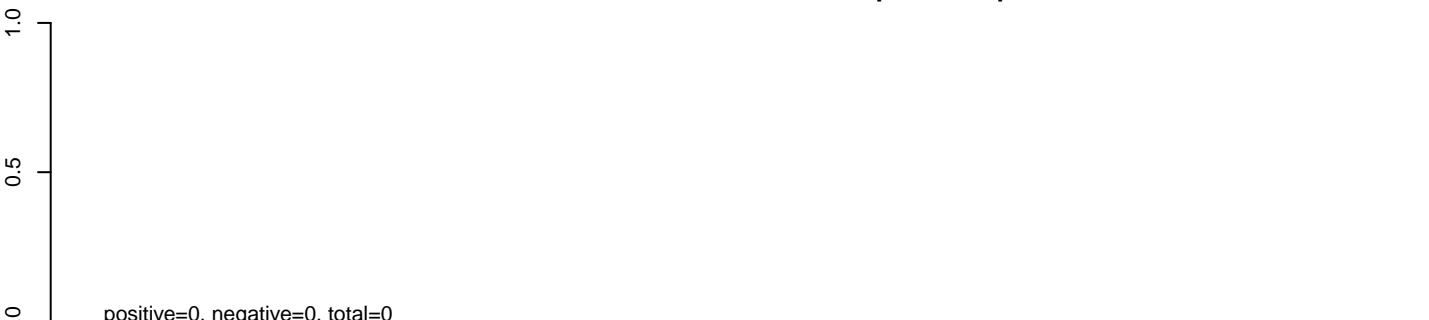
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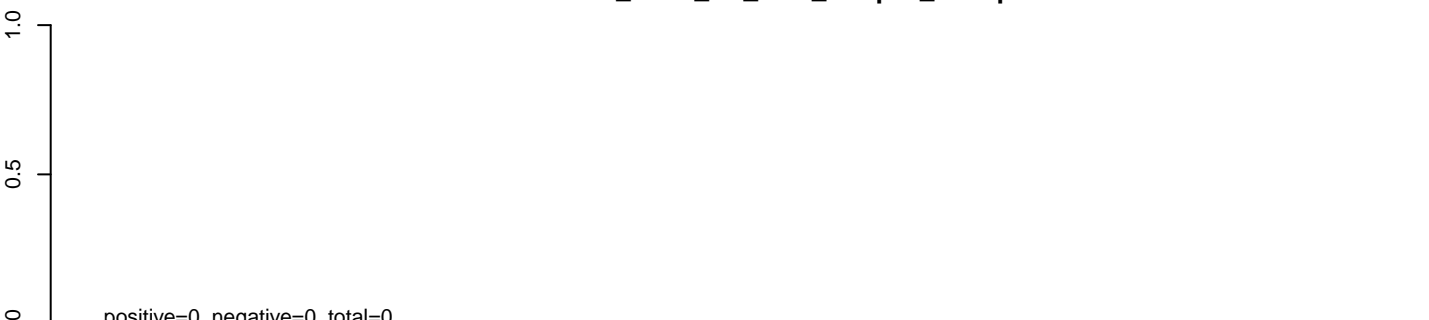
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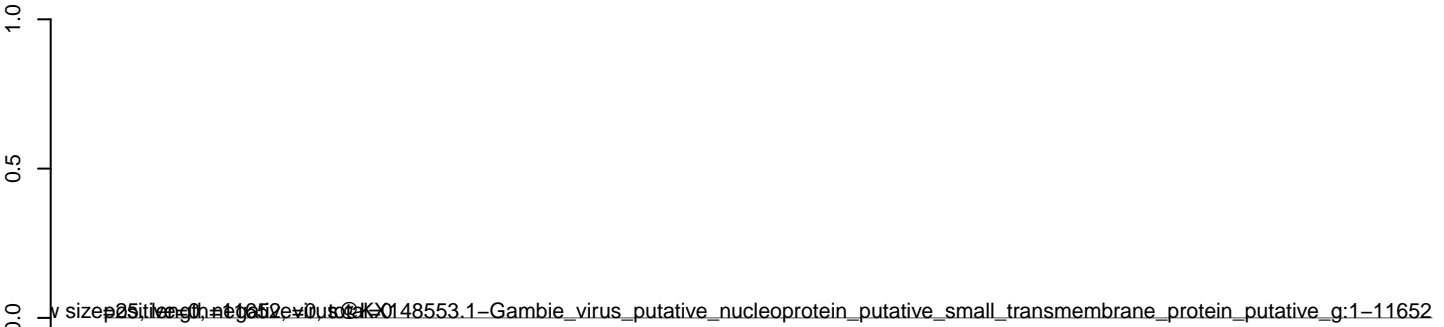
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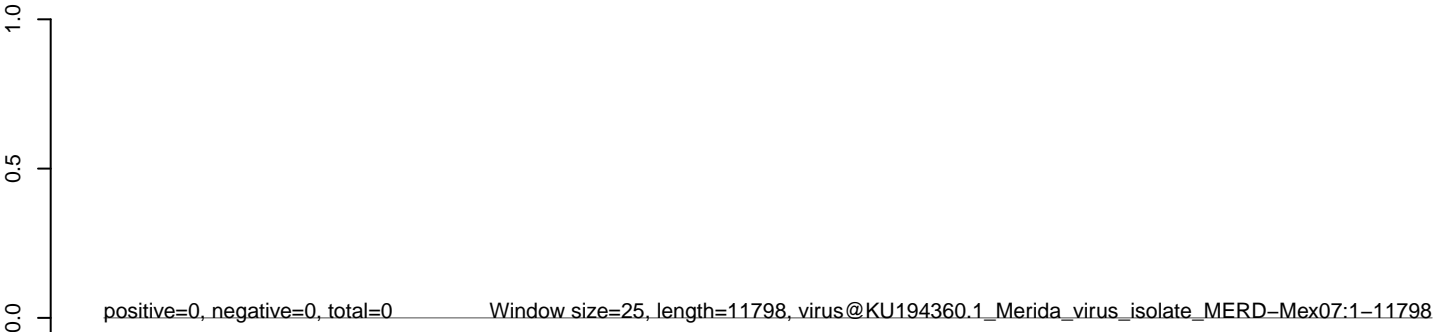
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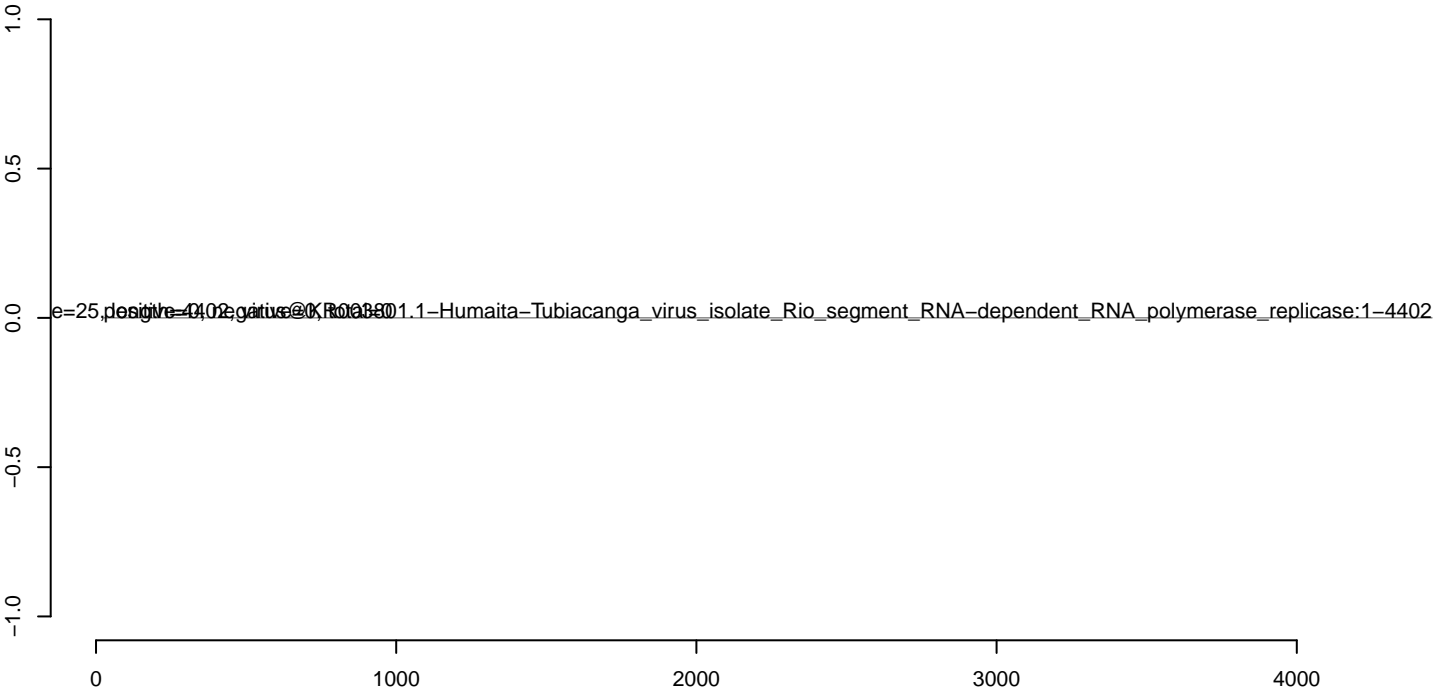
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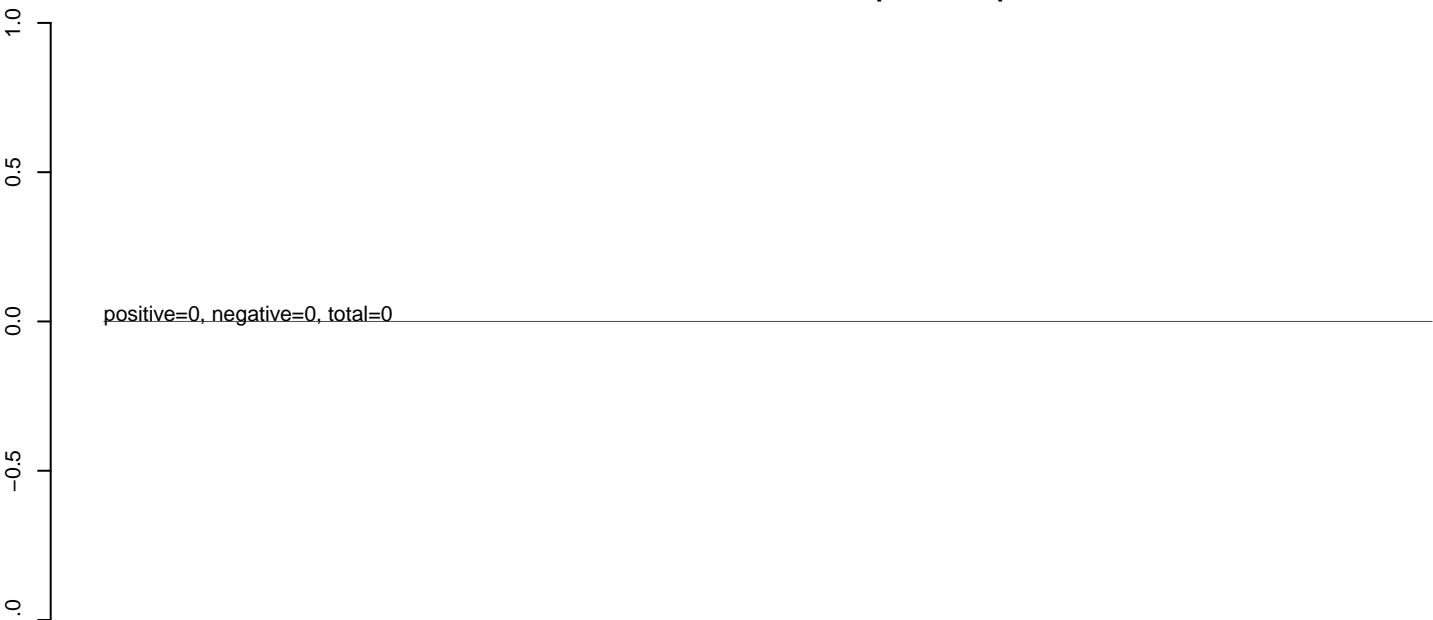
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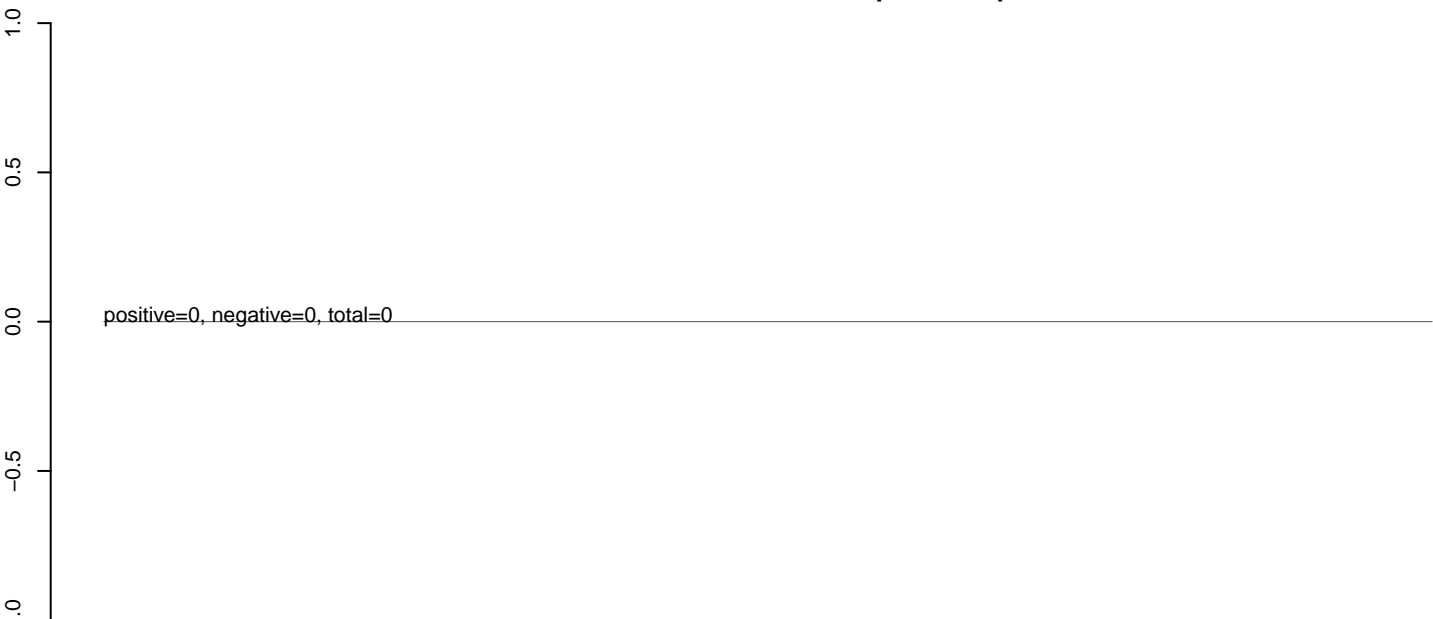
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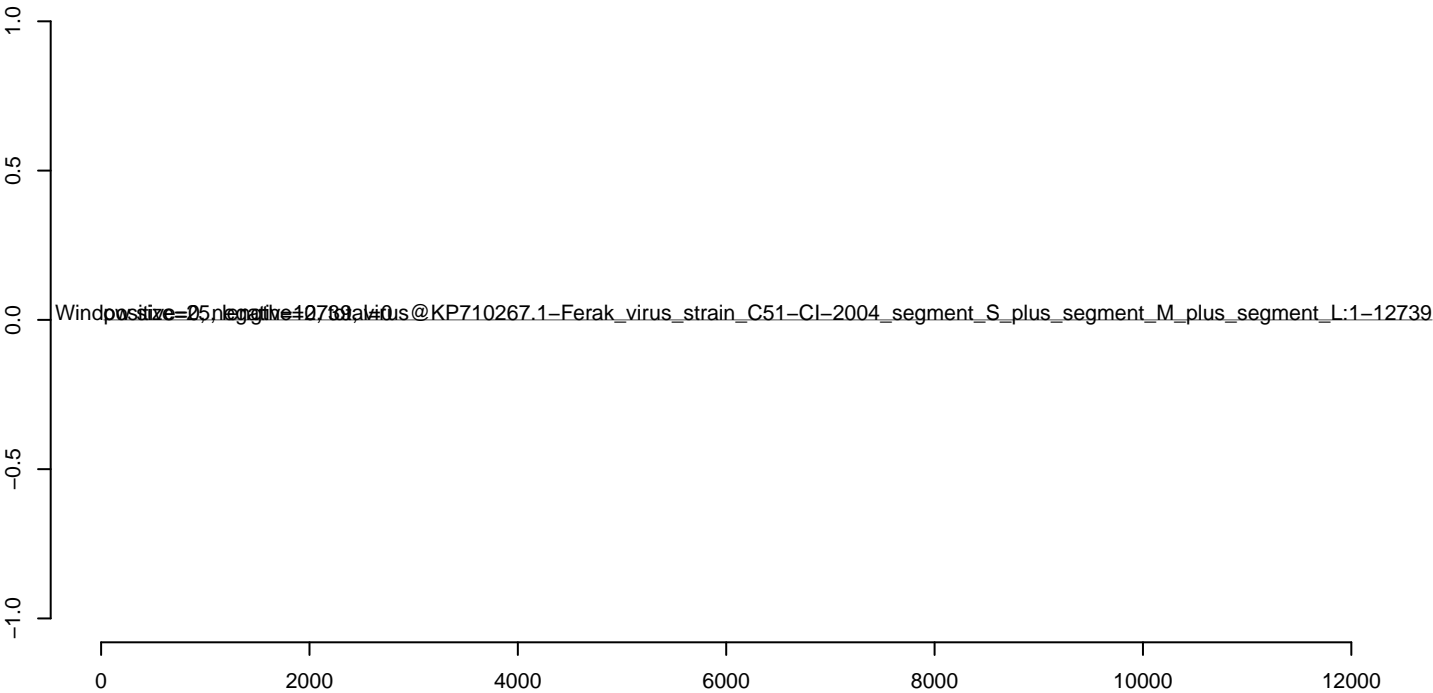
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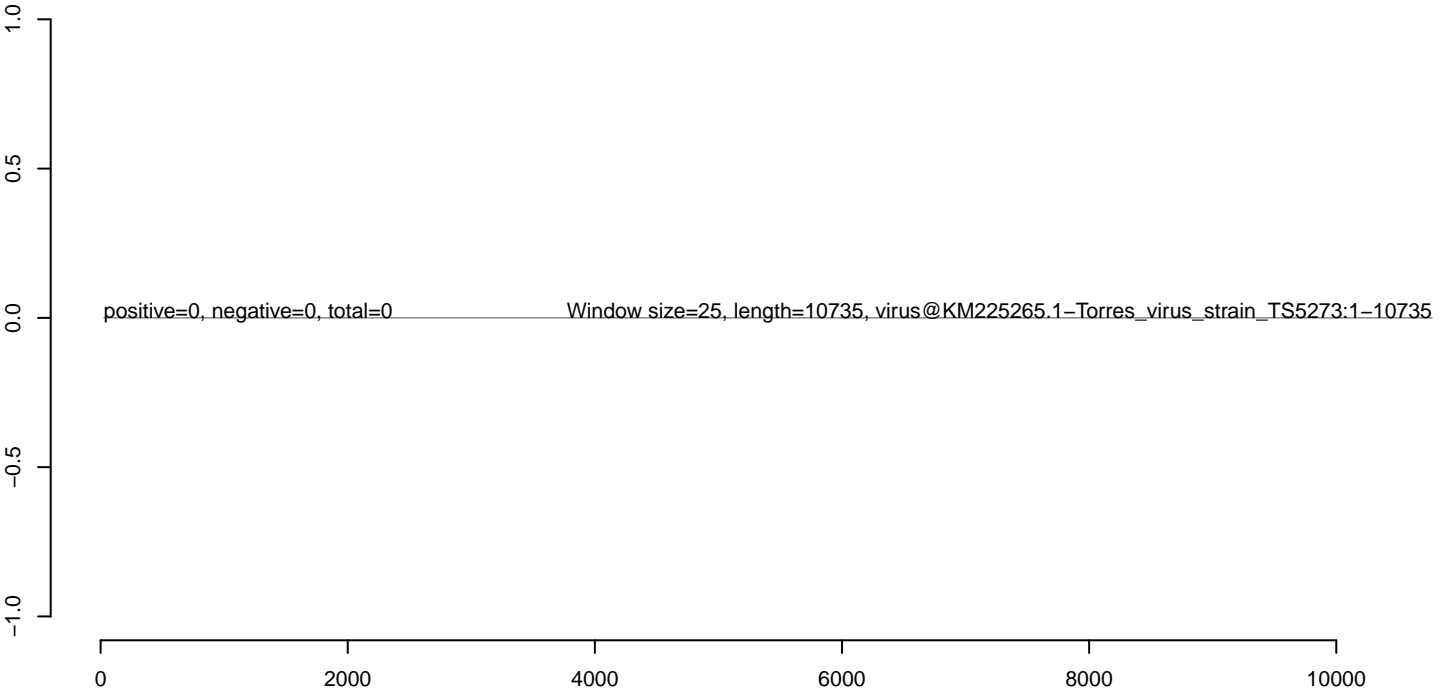
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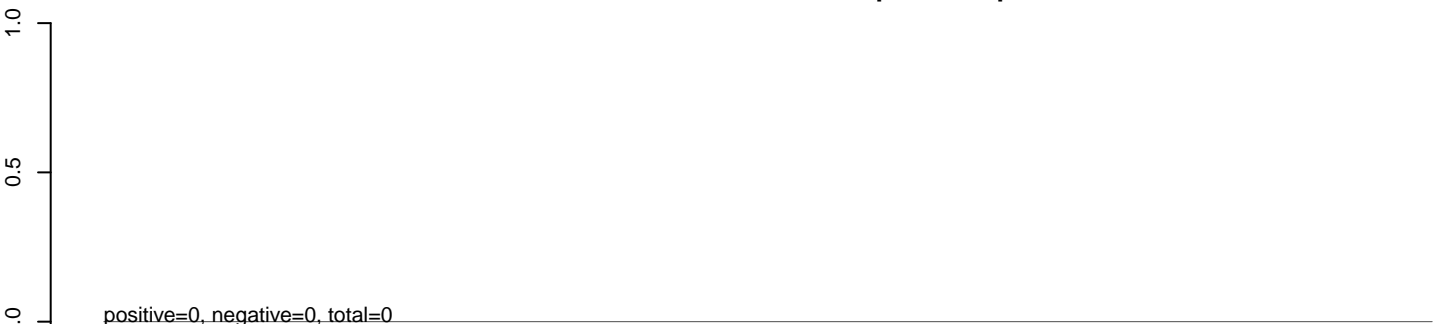
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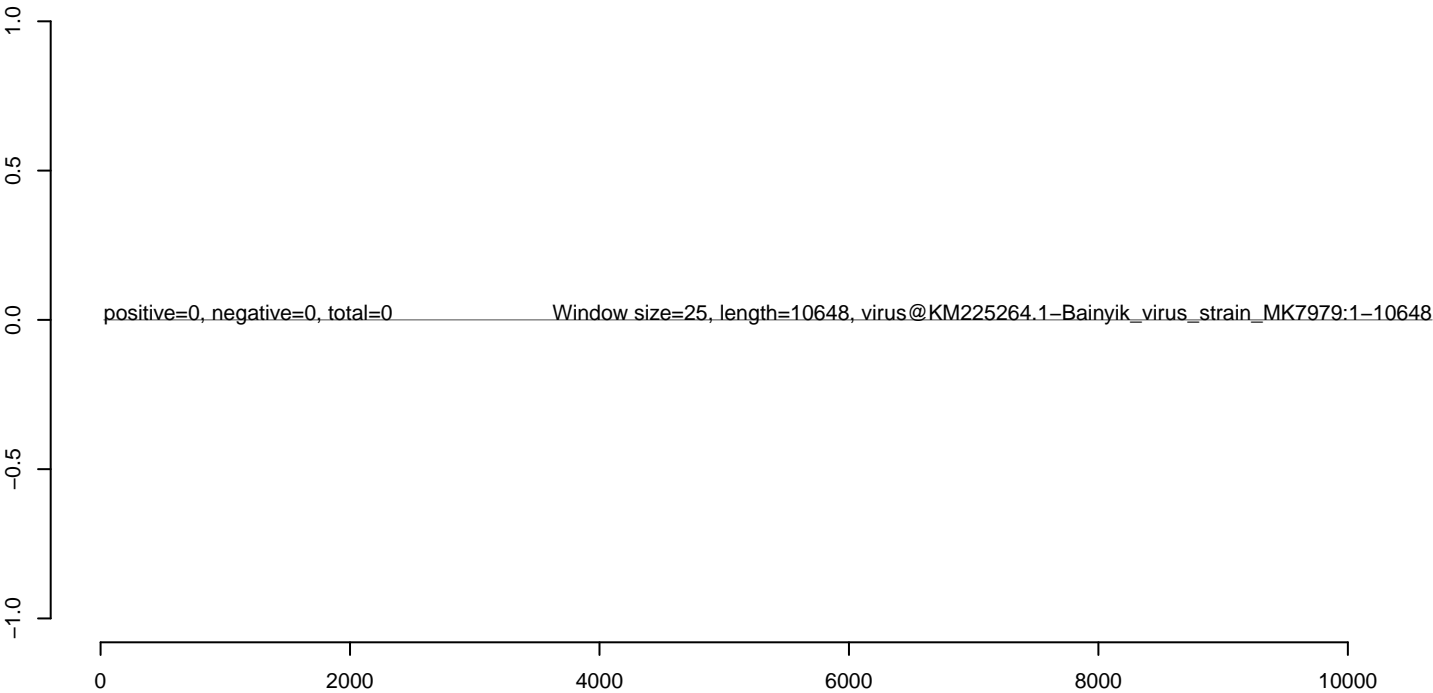
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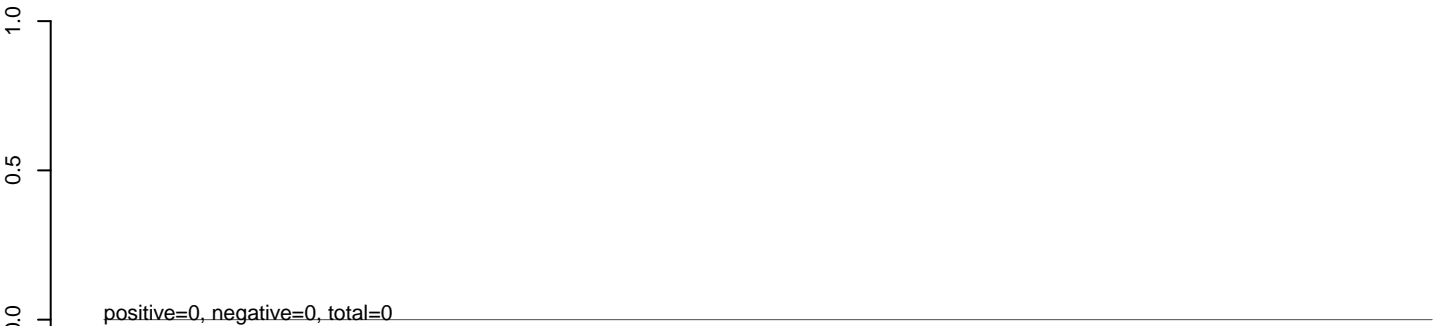
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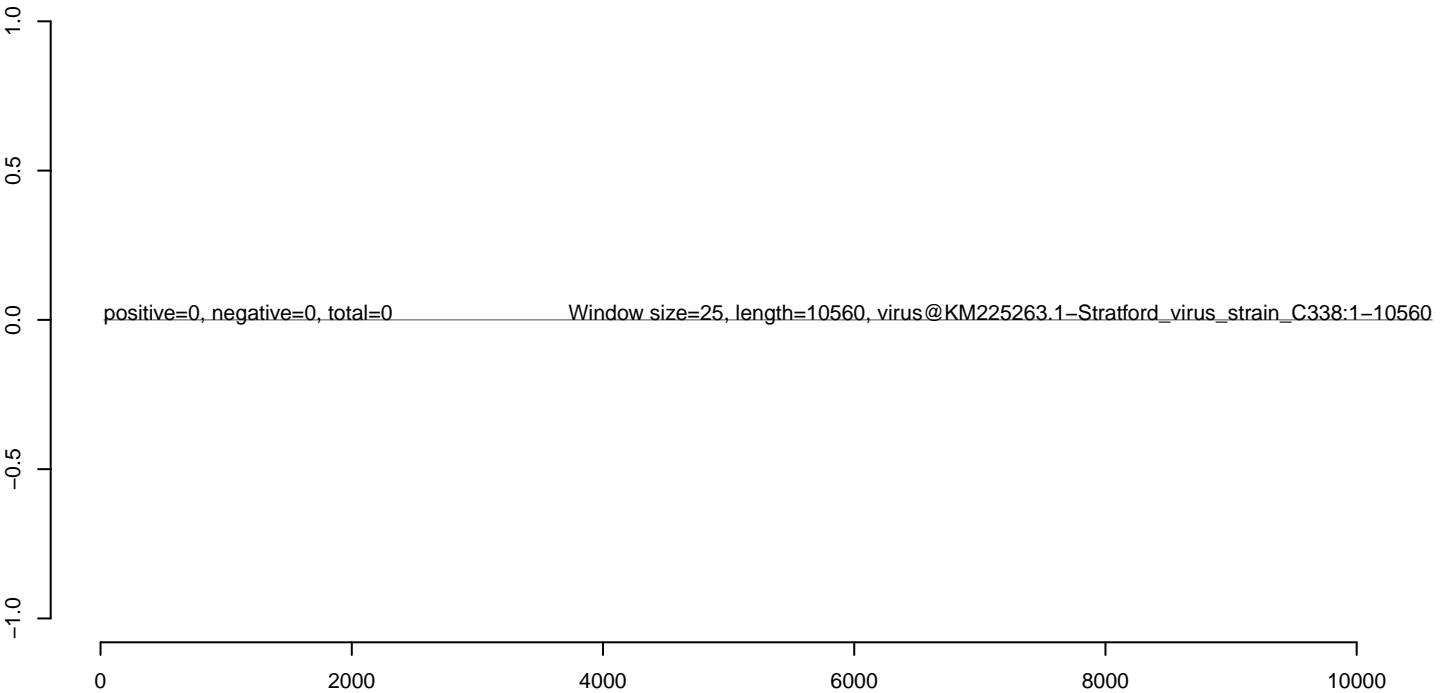
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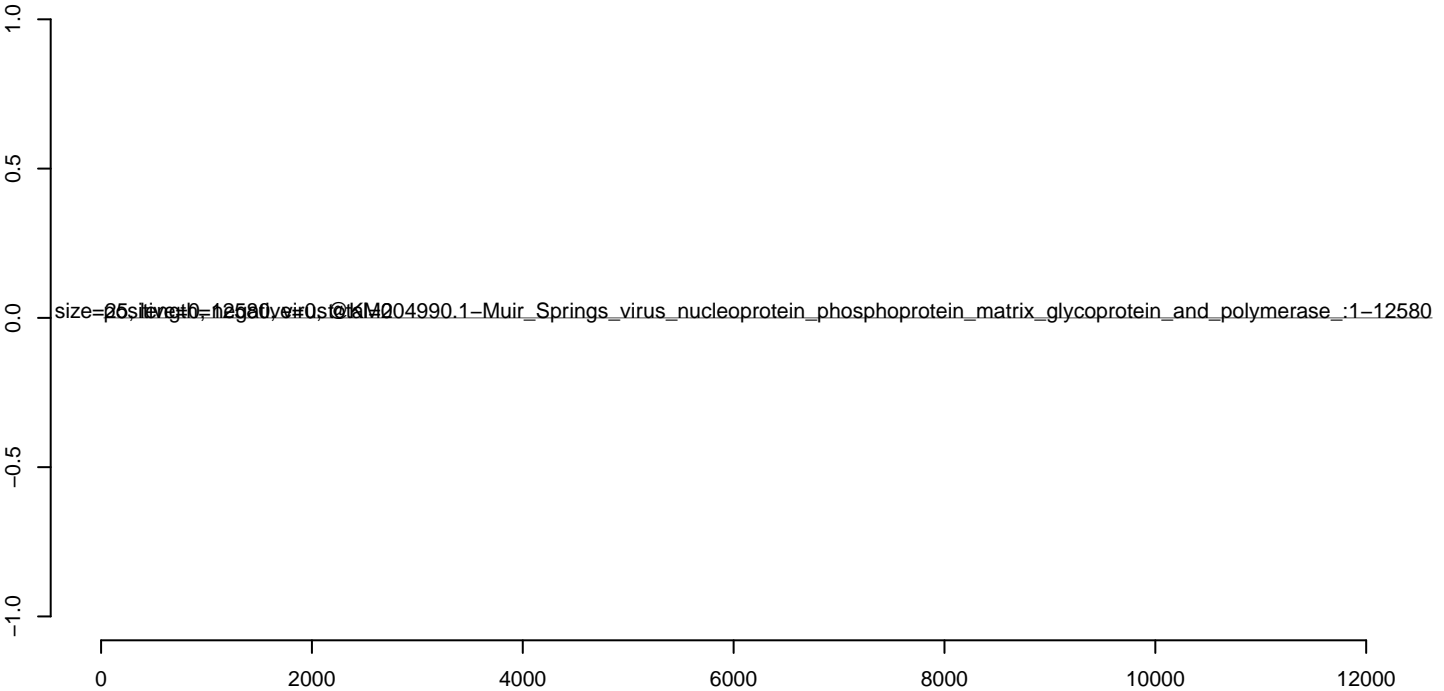
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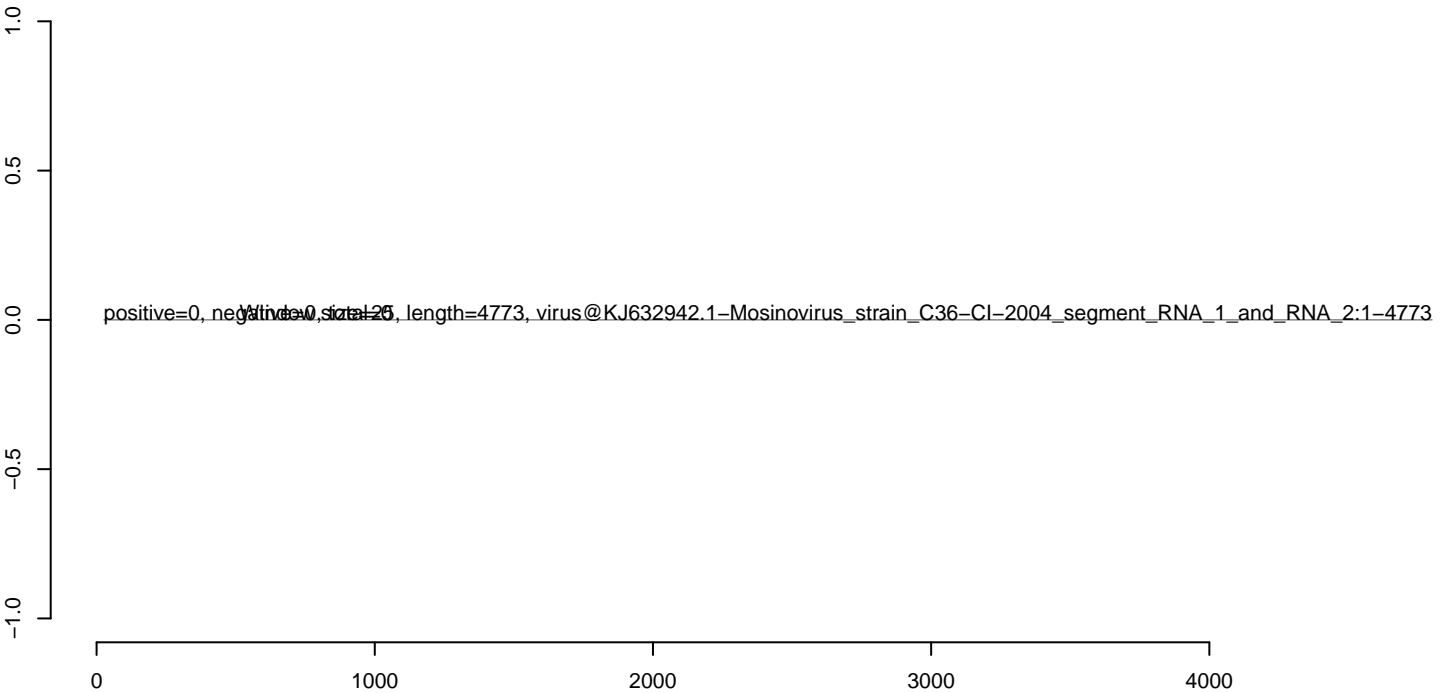
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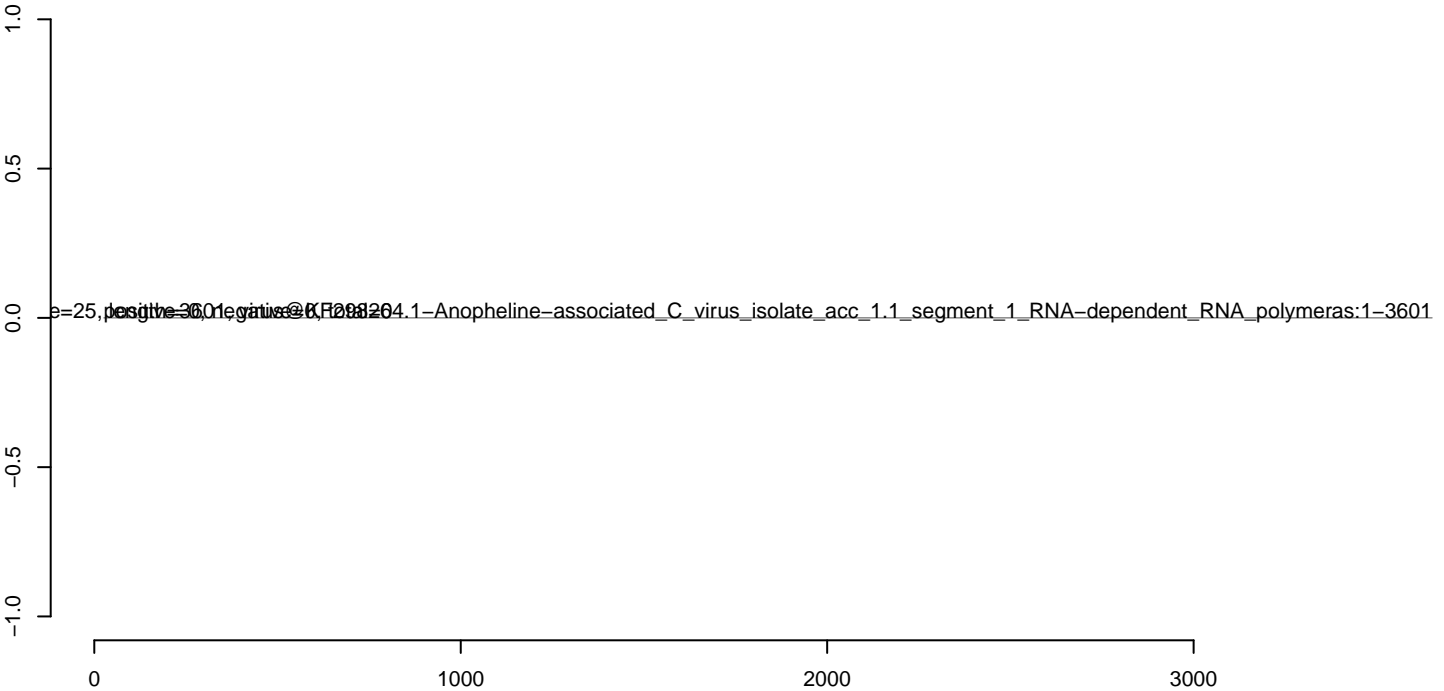
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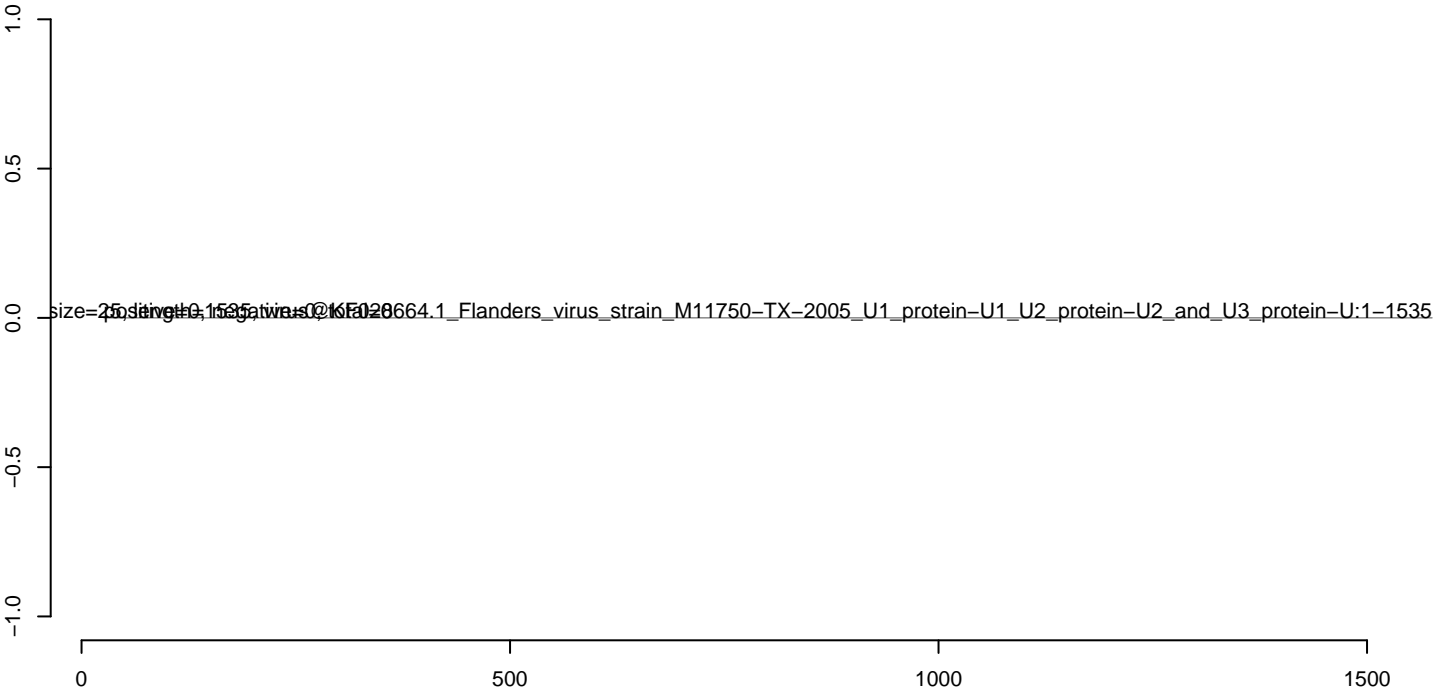
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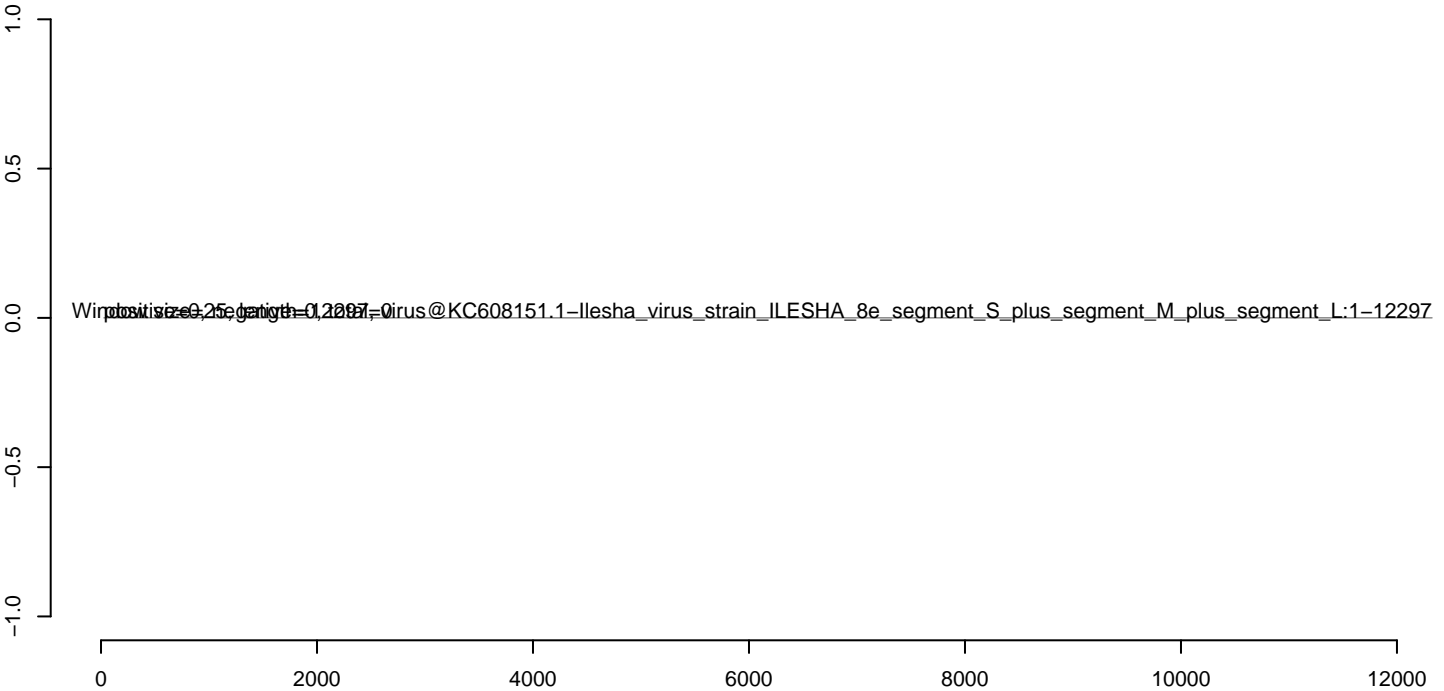
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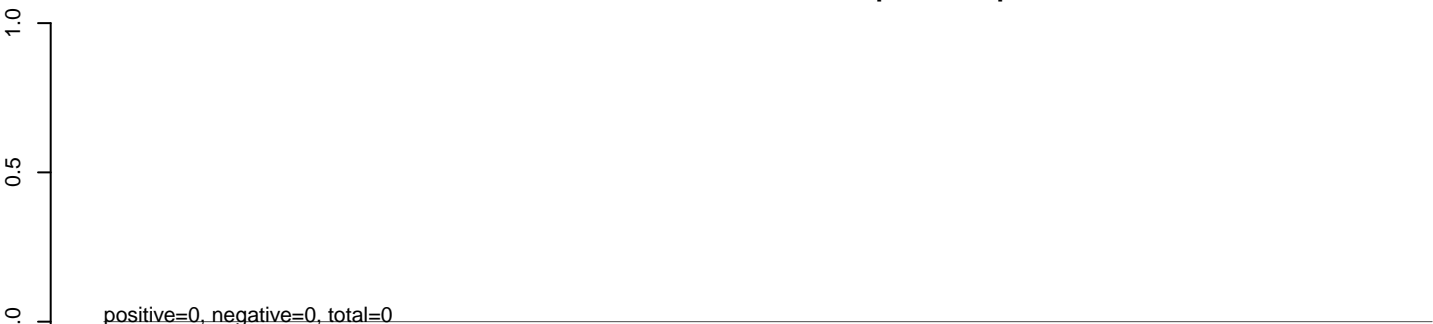
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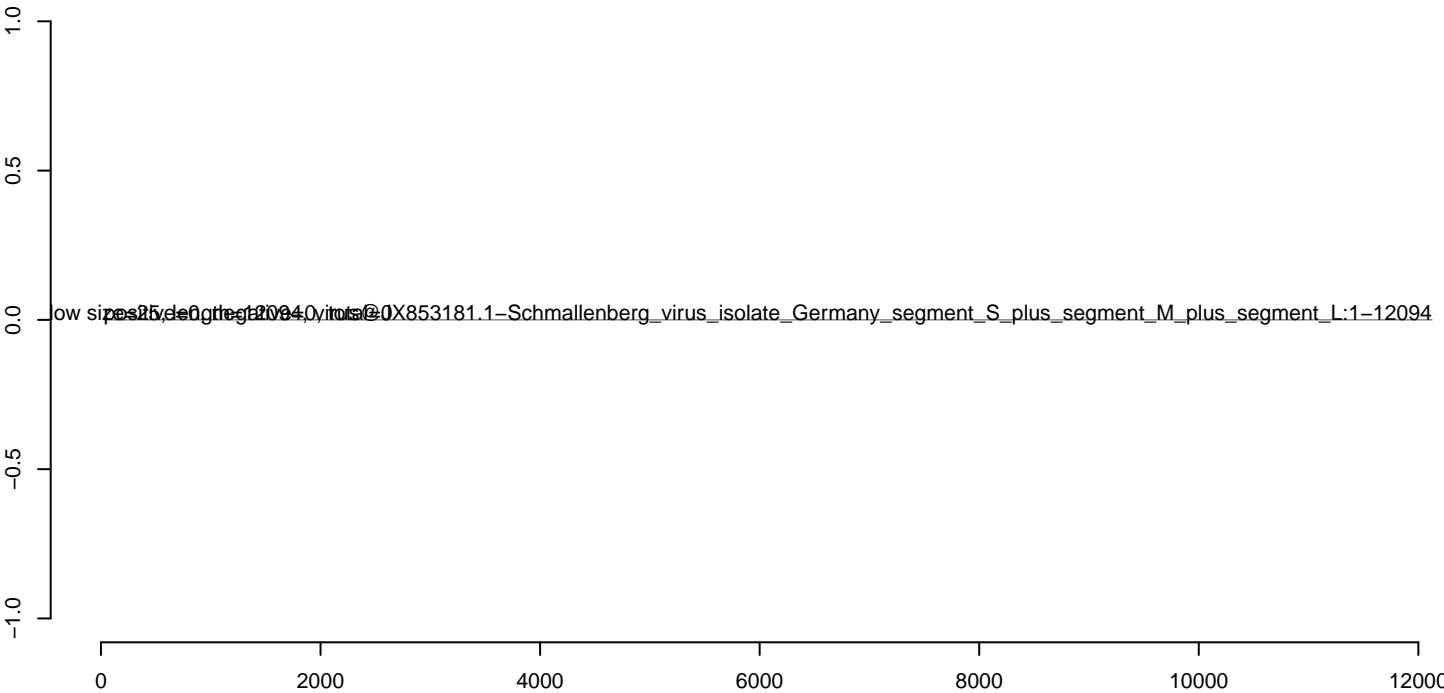
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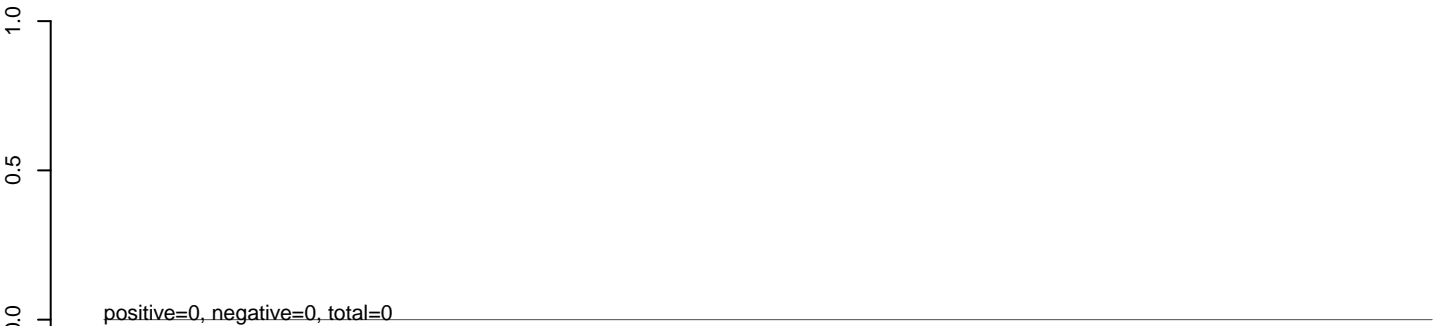
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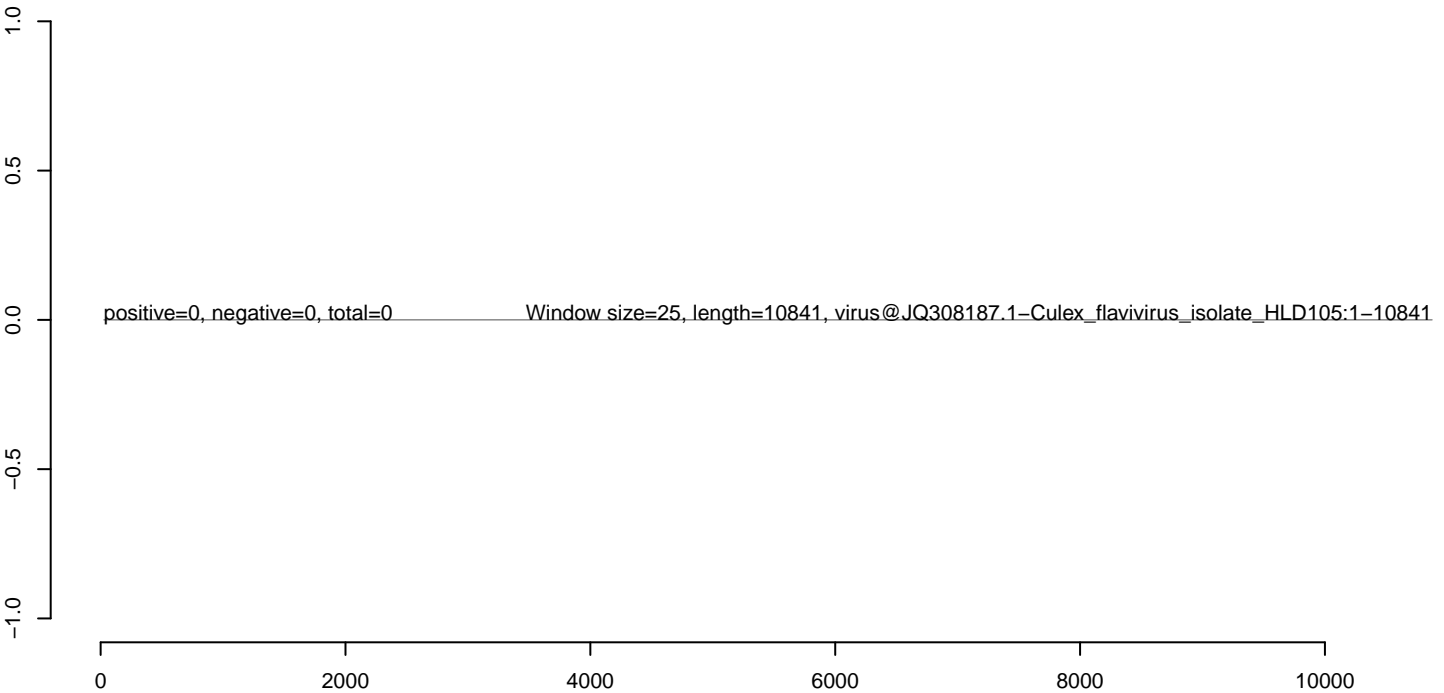
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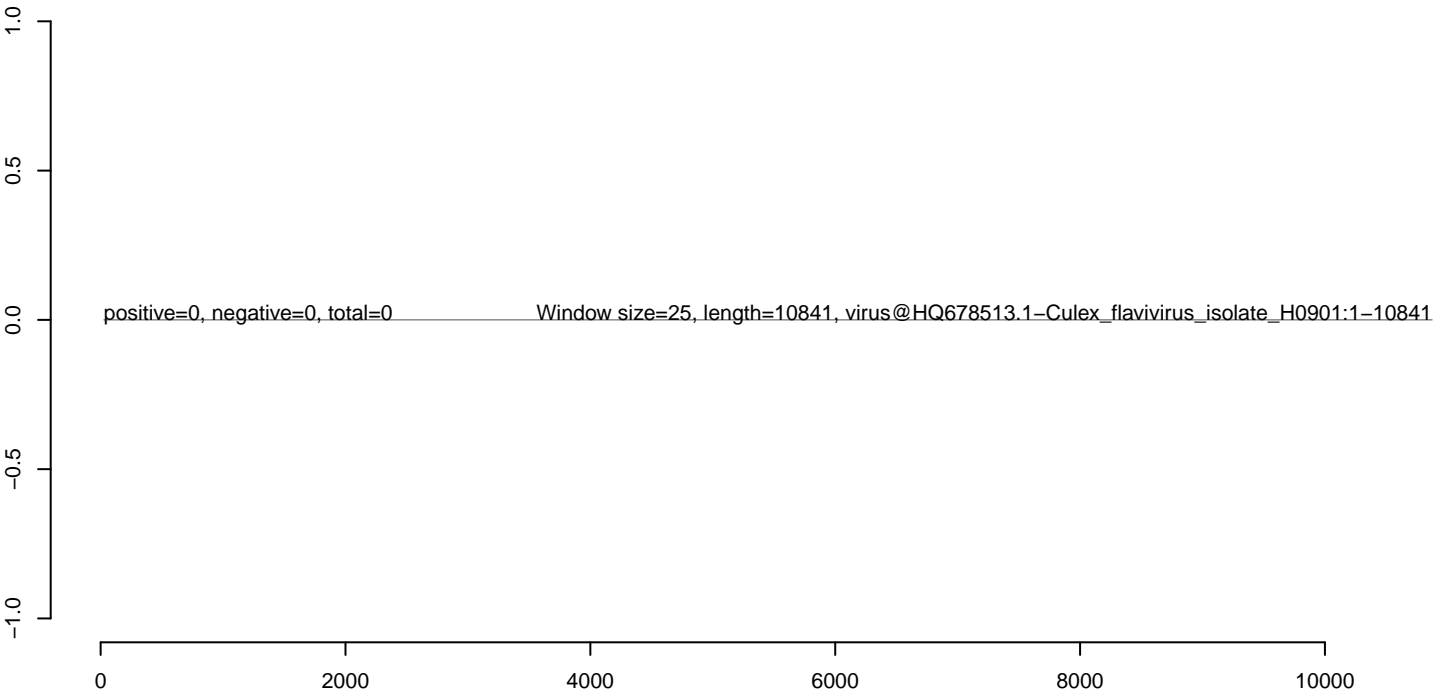
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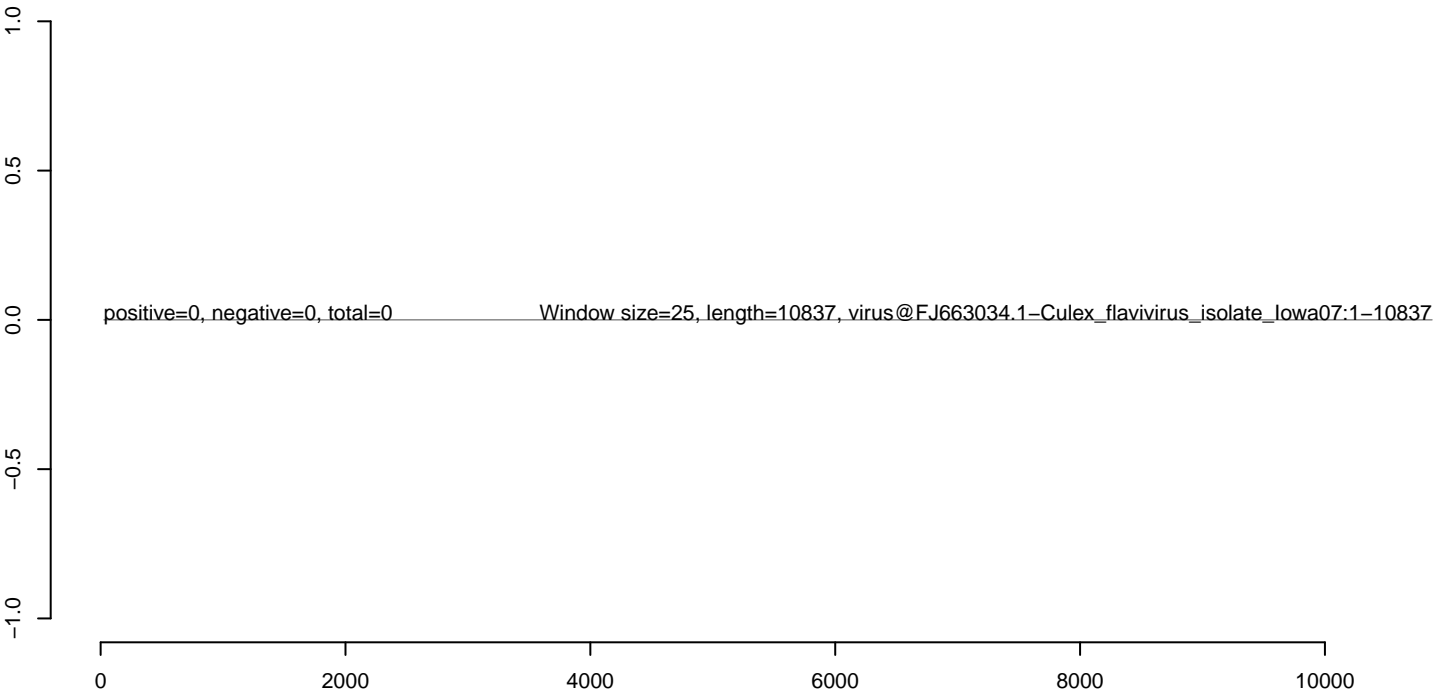
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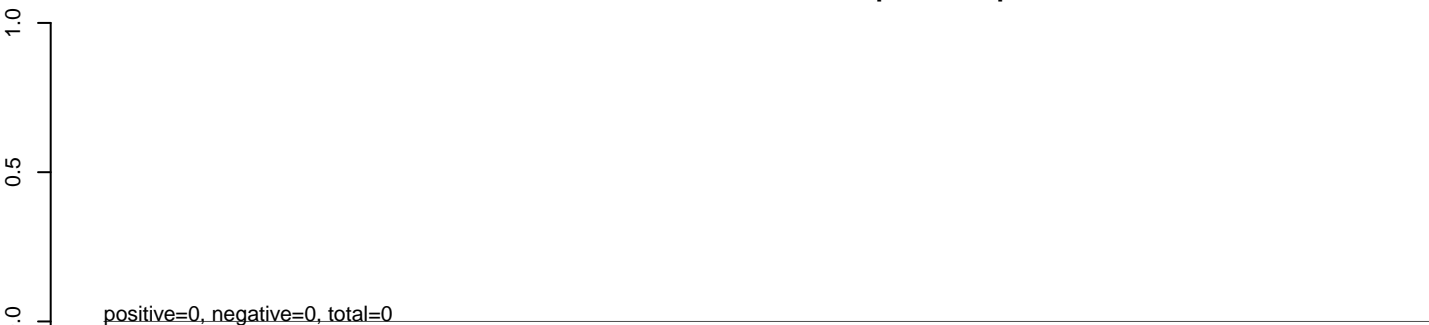
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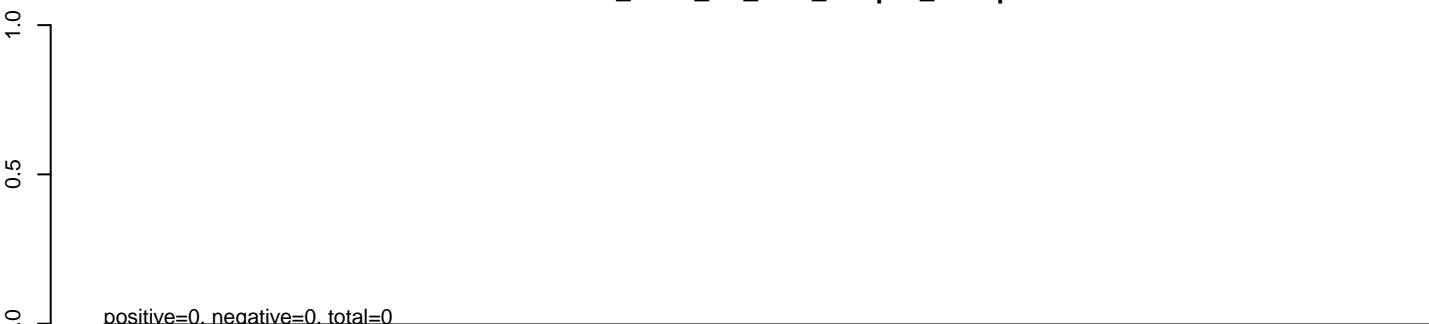
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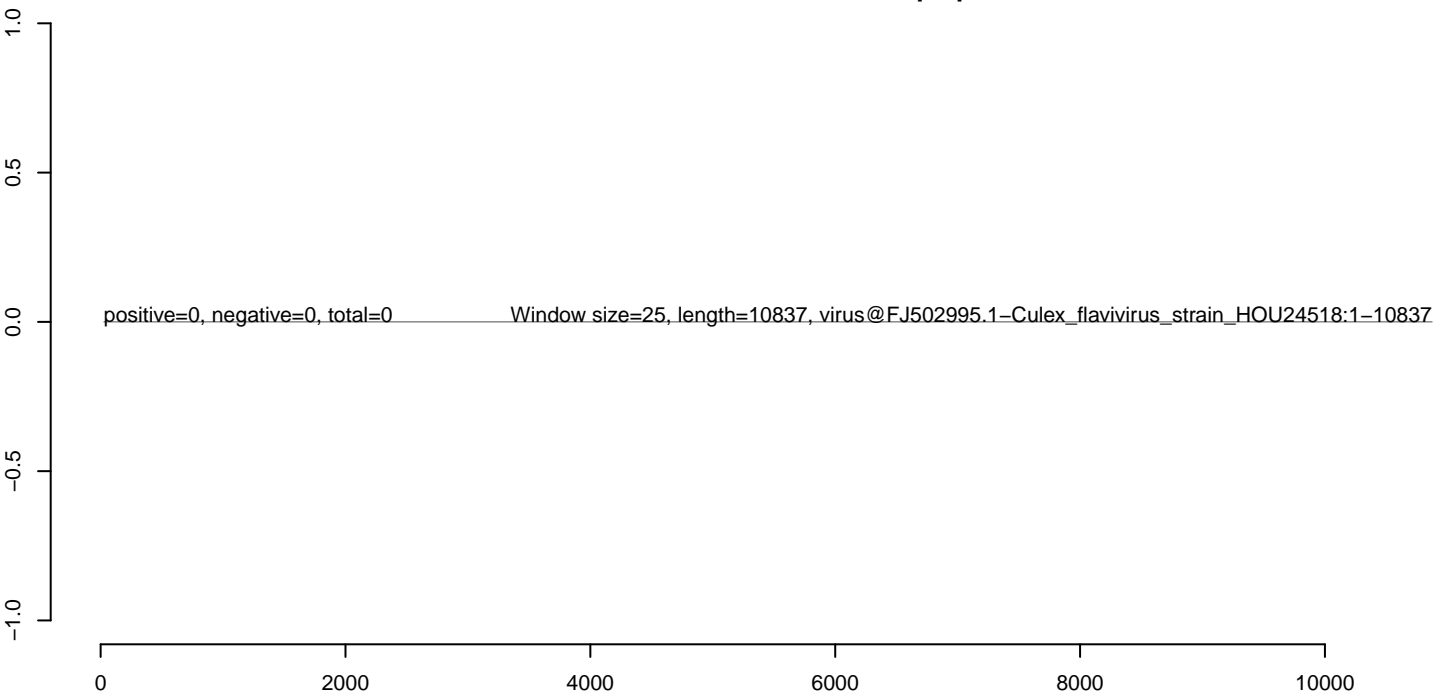
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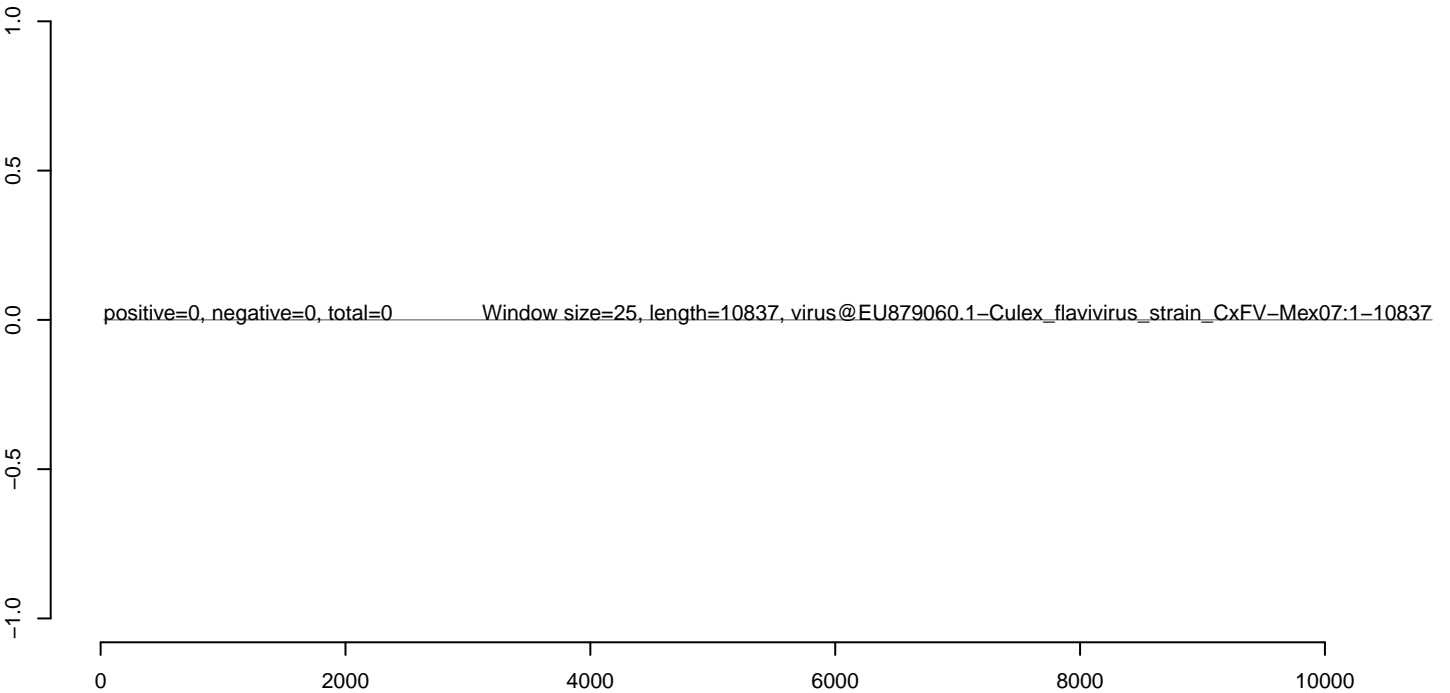
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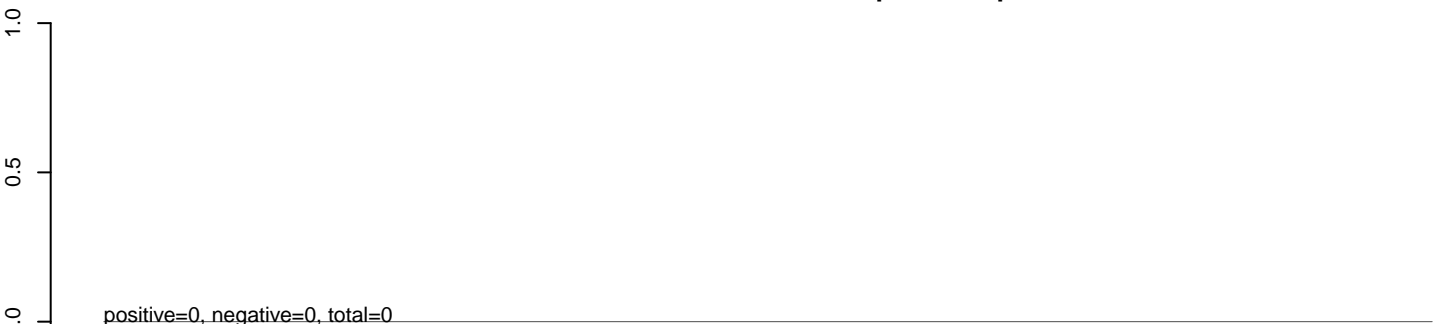
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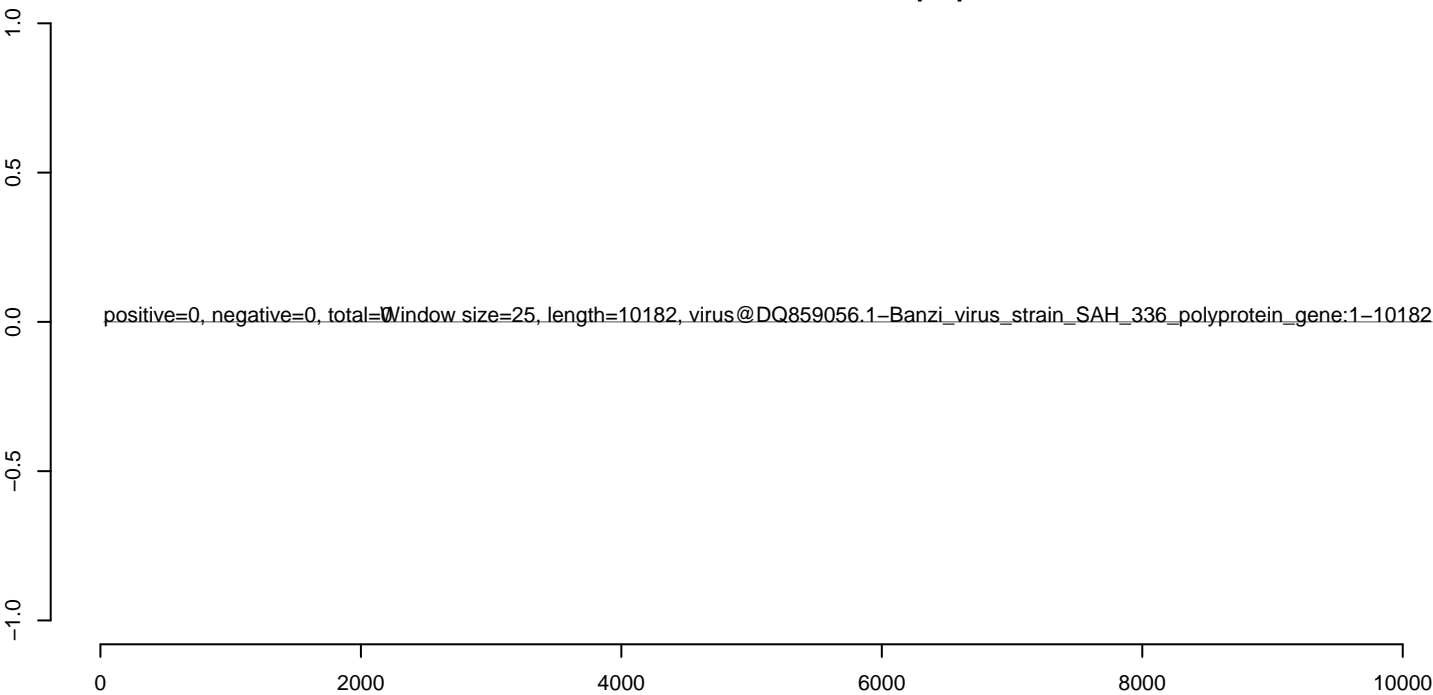
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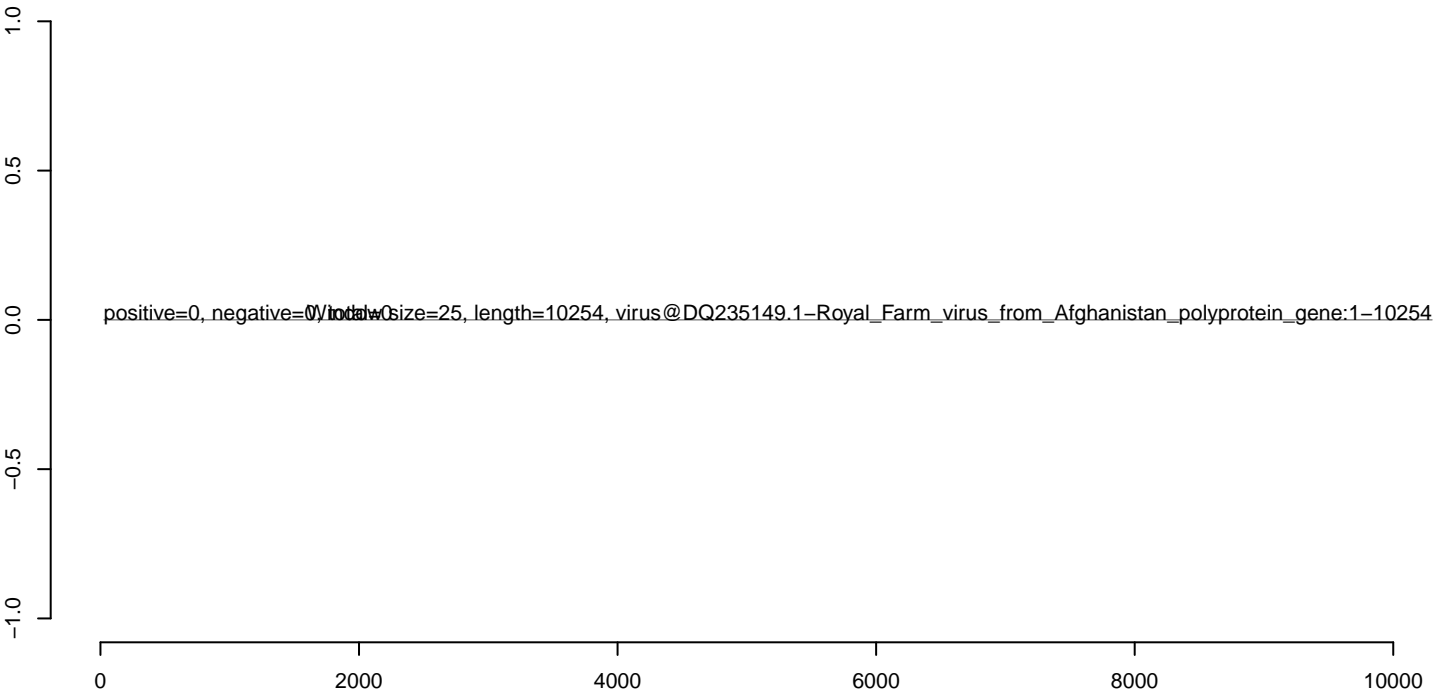
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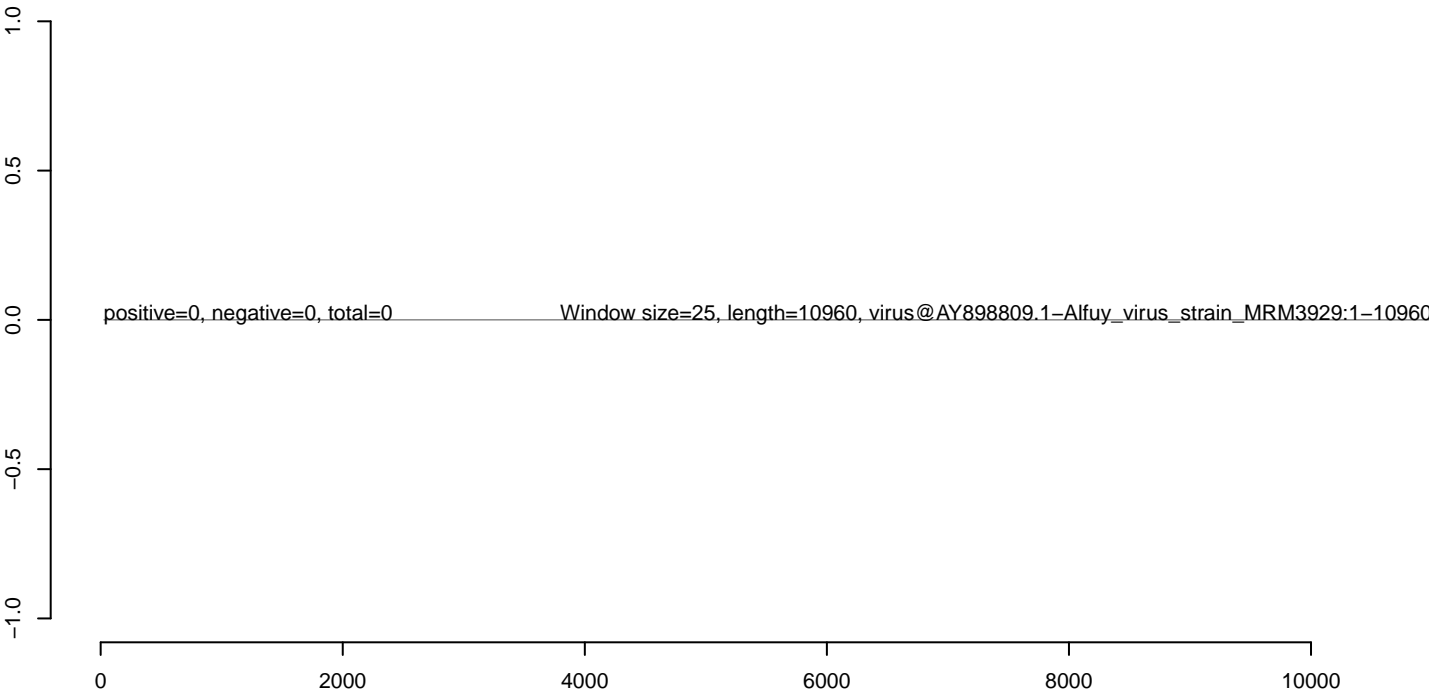
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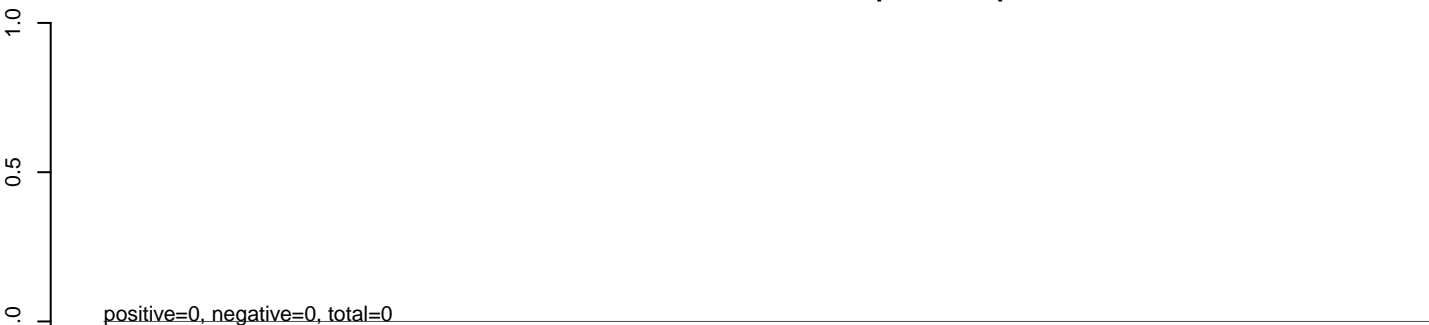
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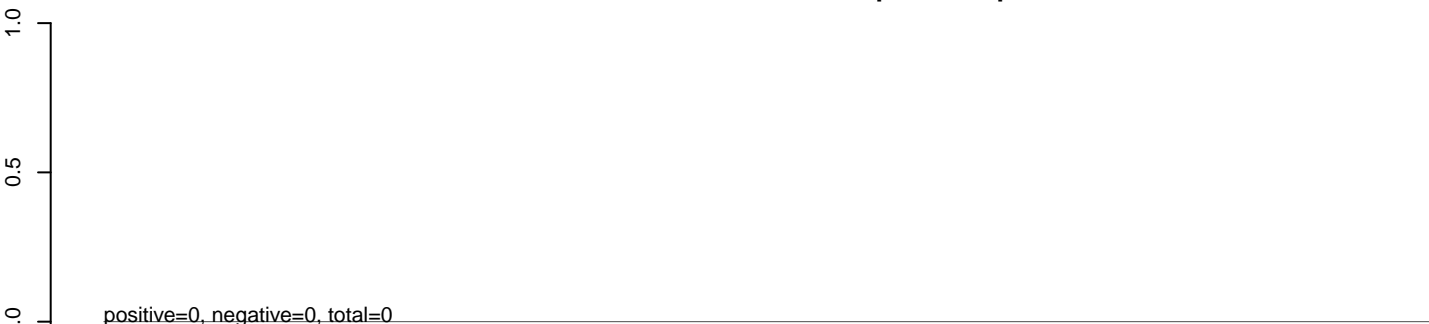
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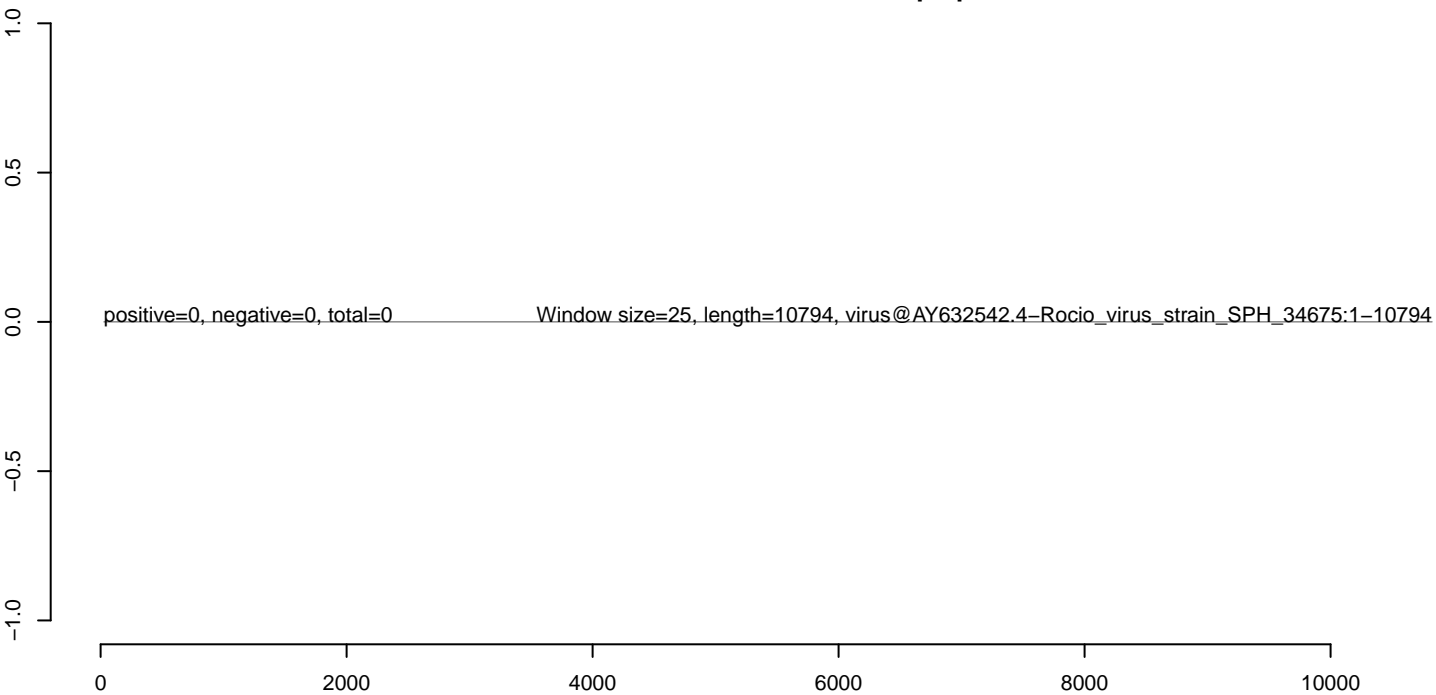
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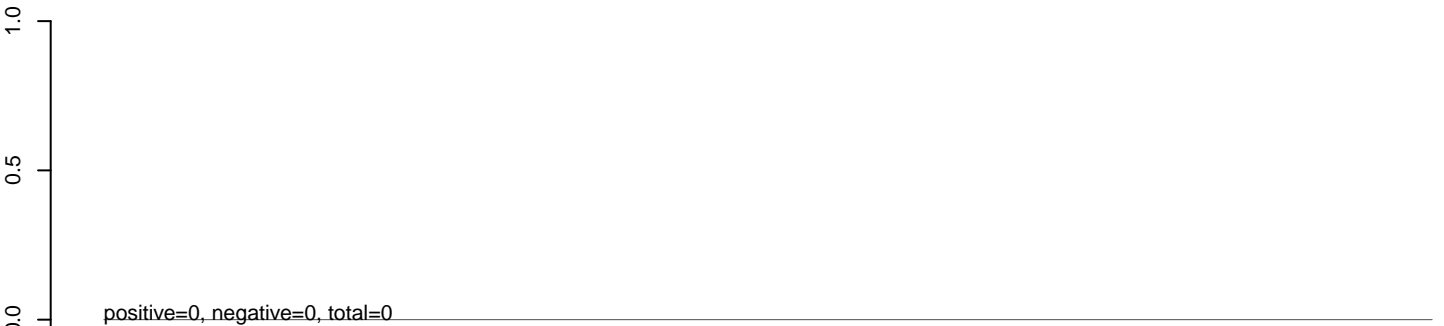
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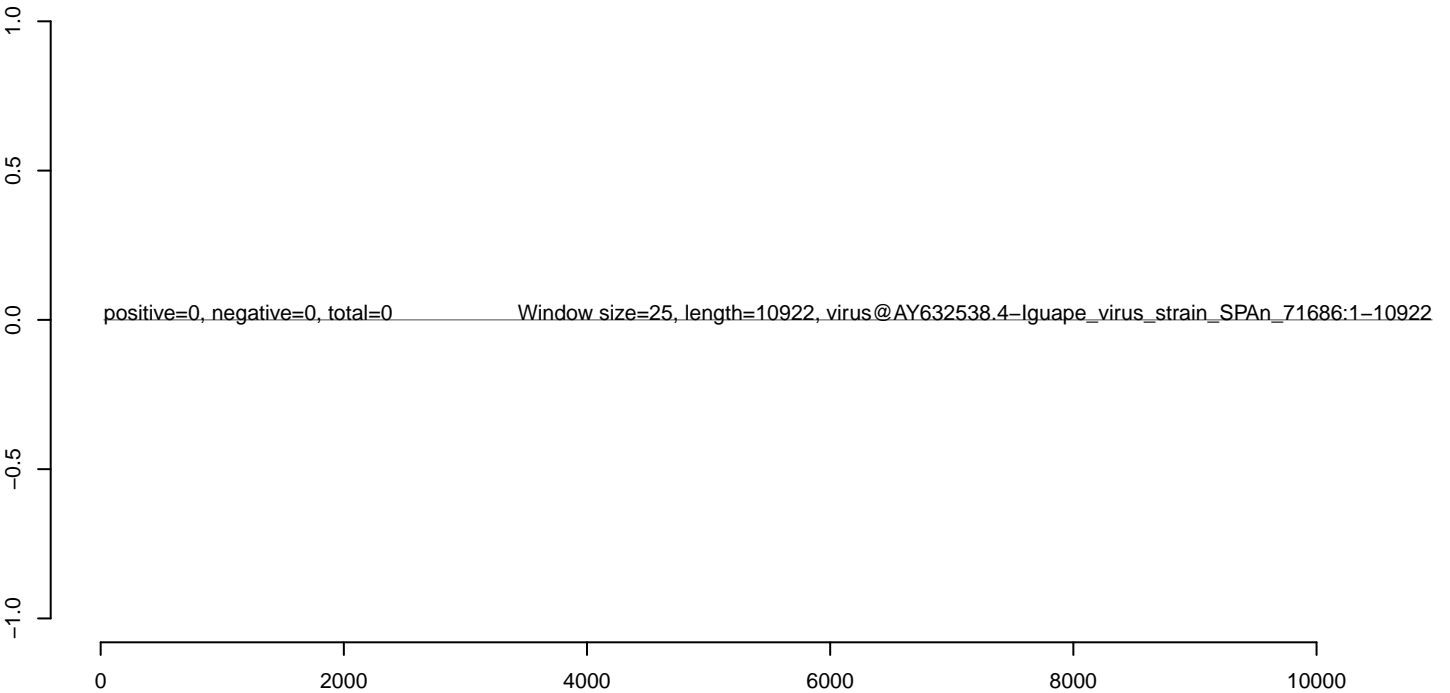
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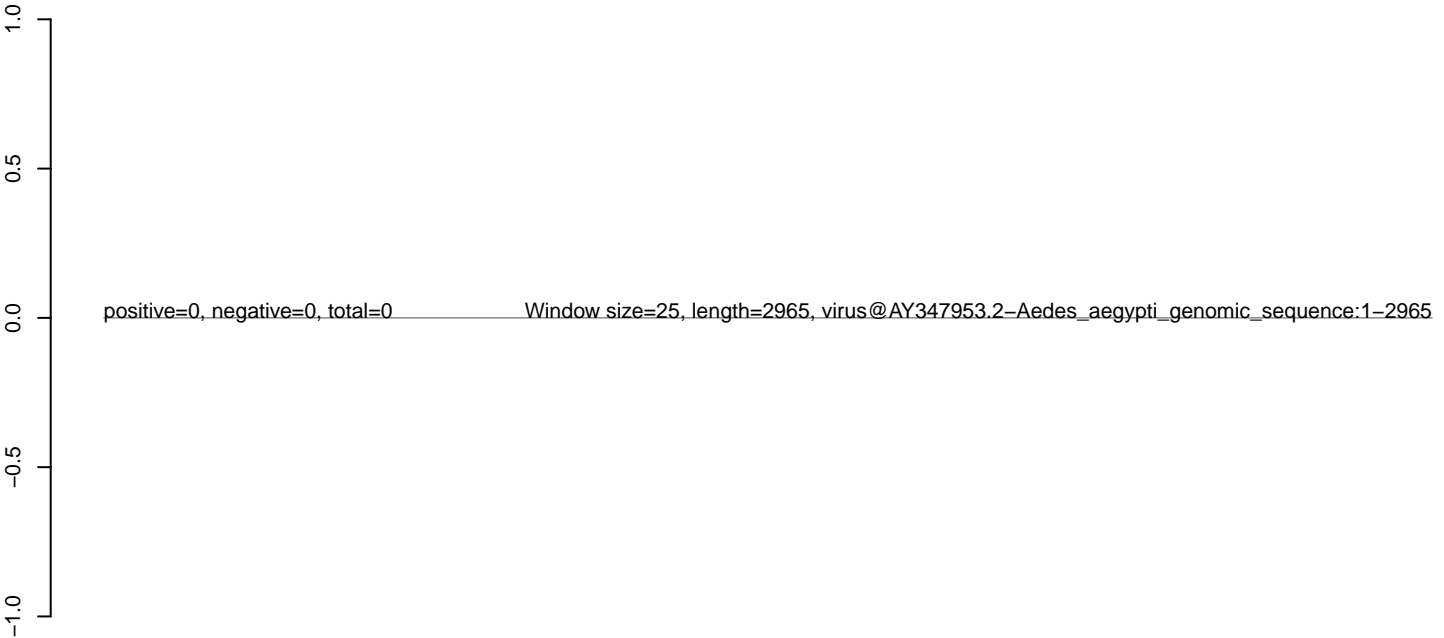
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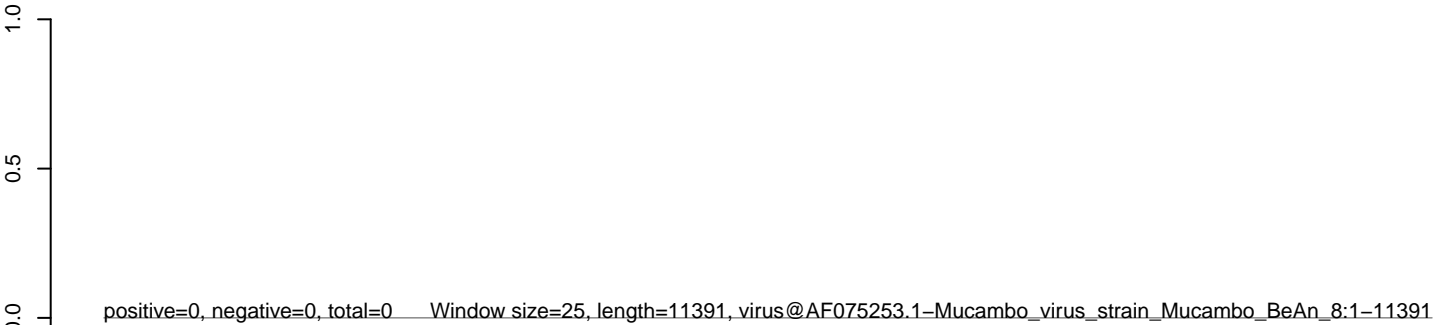
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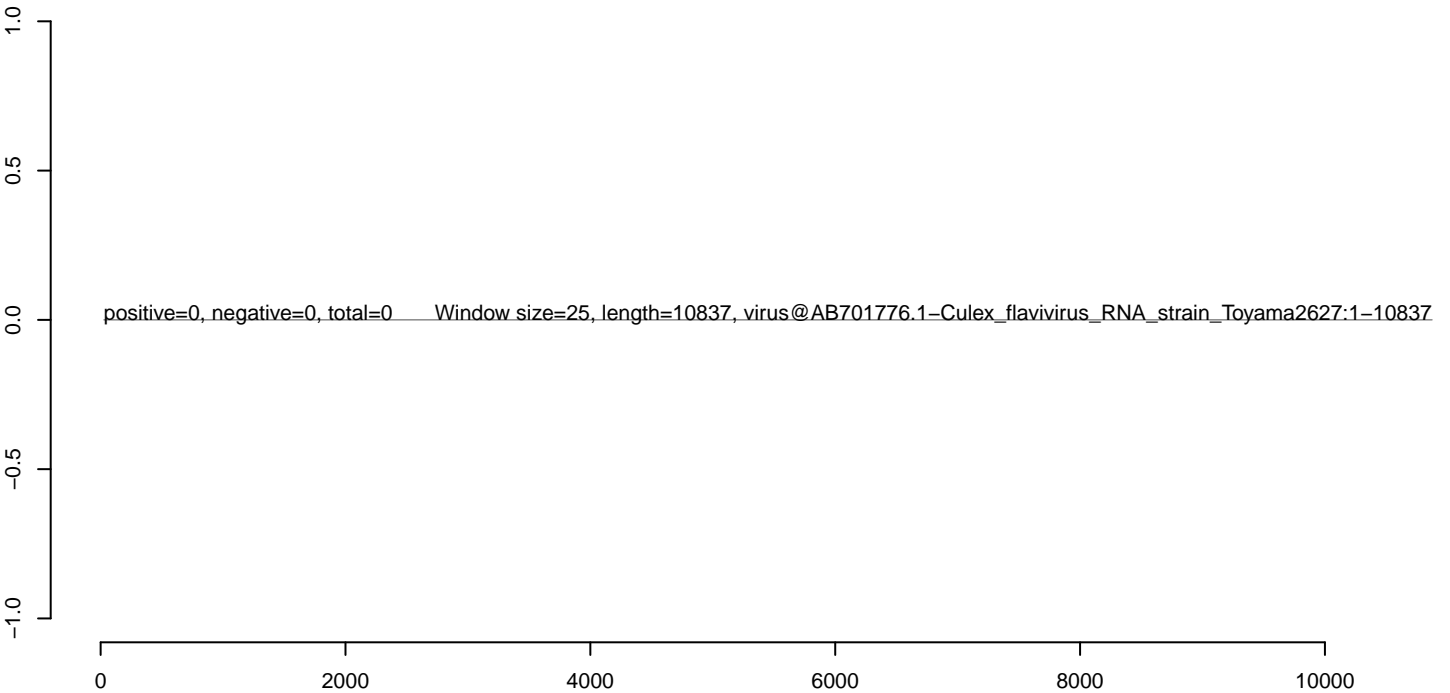
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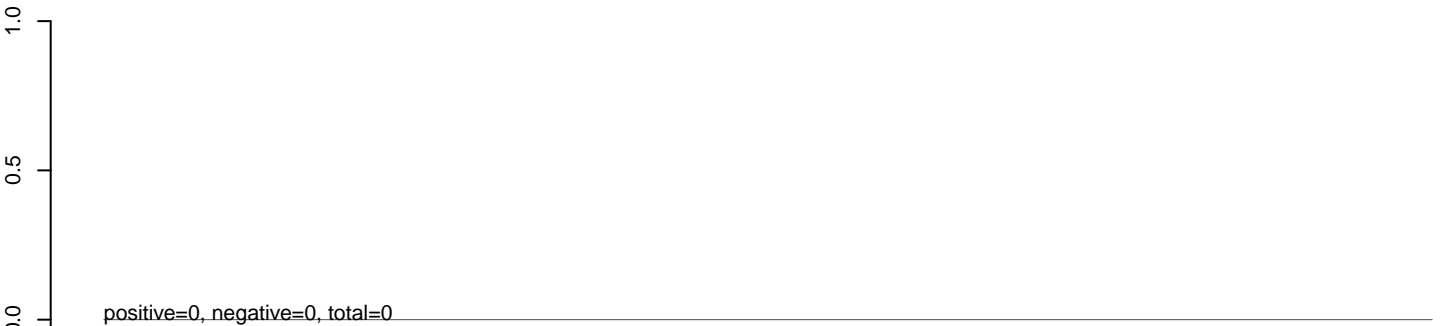
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