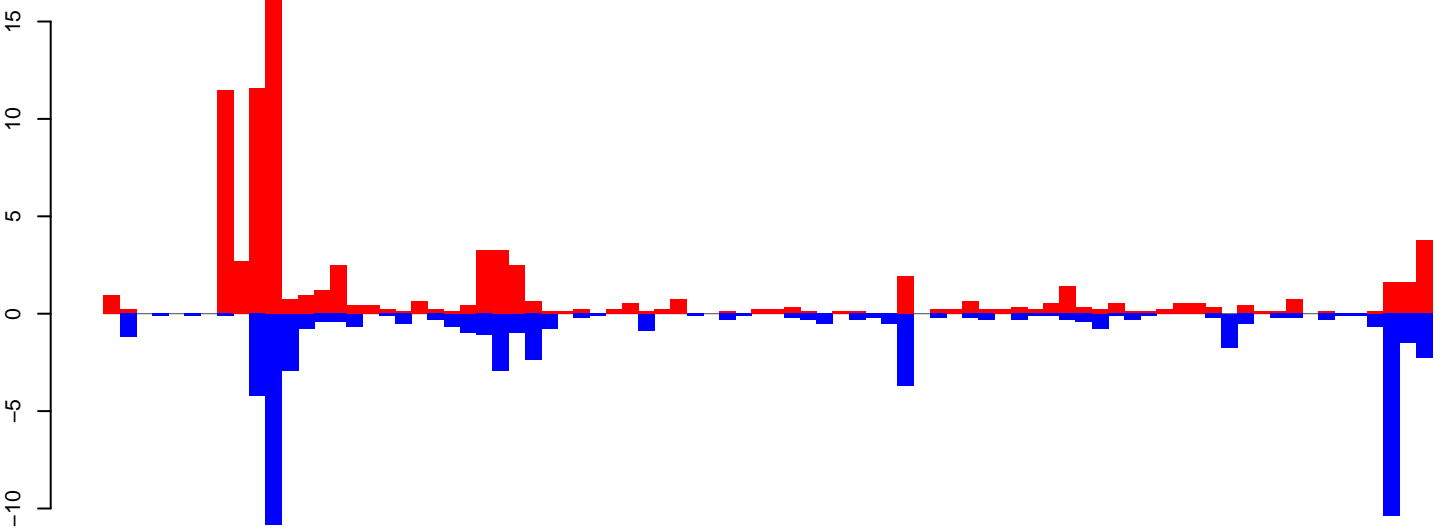
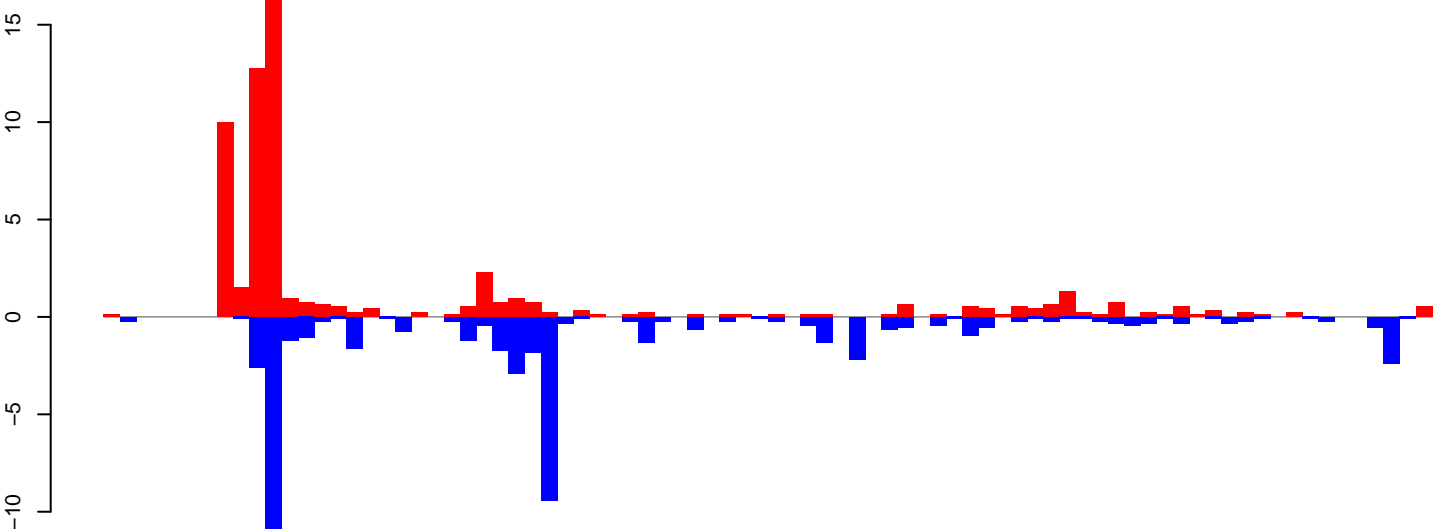


AnGam\_Sua5b\_cells.18\_23.rep



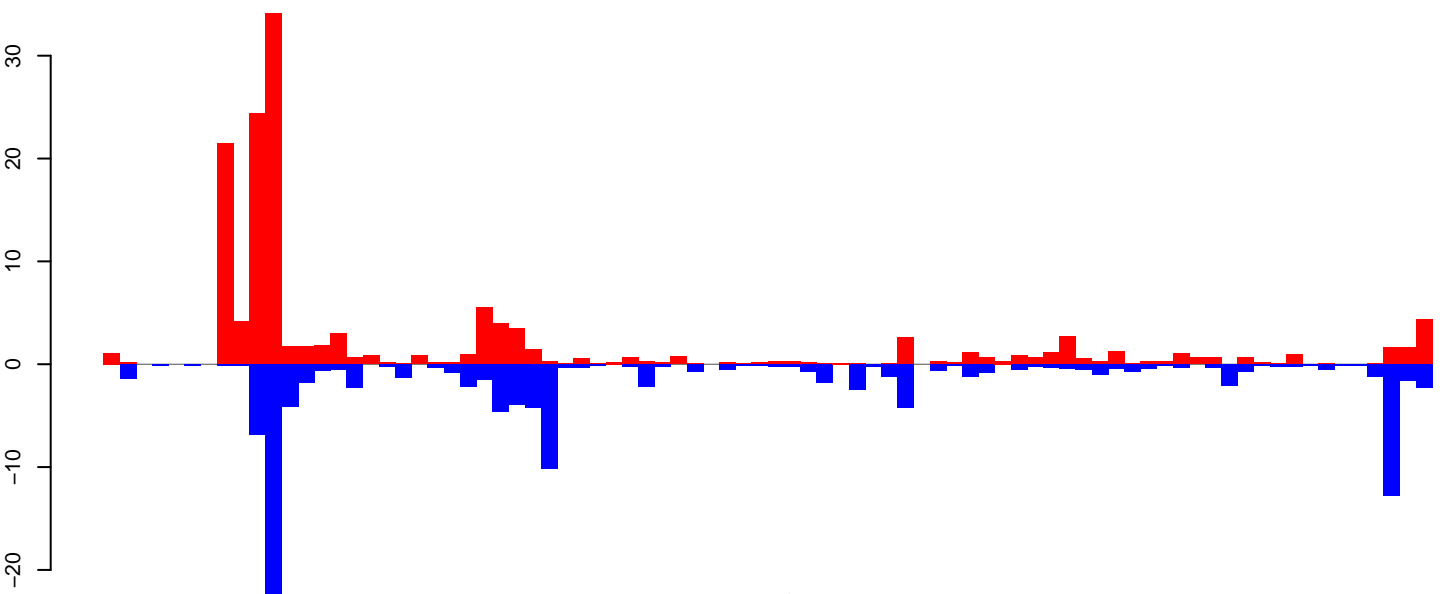
positive=87, negative=64, total=151

AnGam\_Sua5b\_cells.24\_35.rep



positive=61, negative=56, total=117

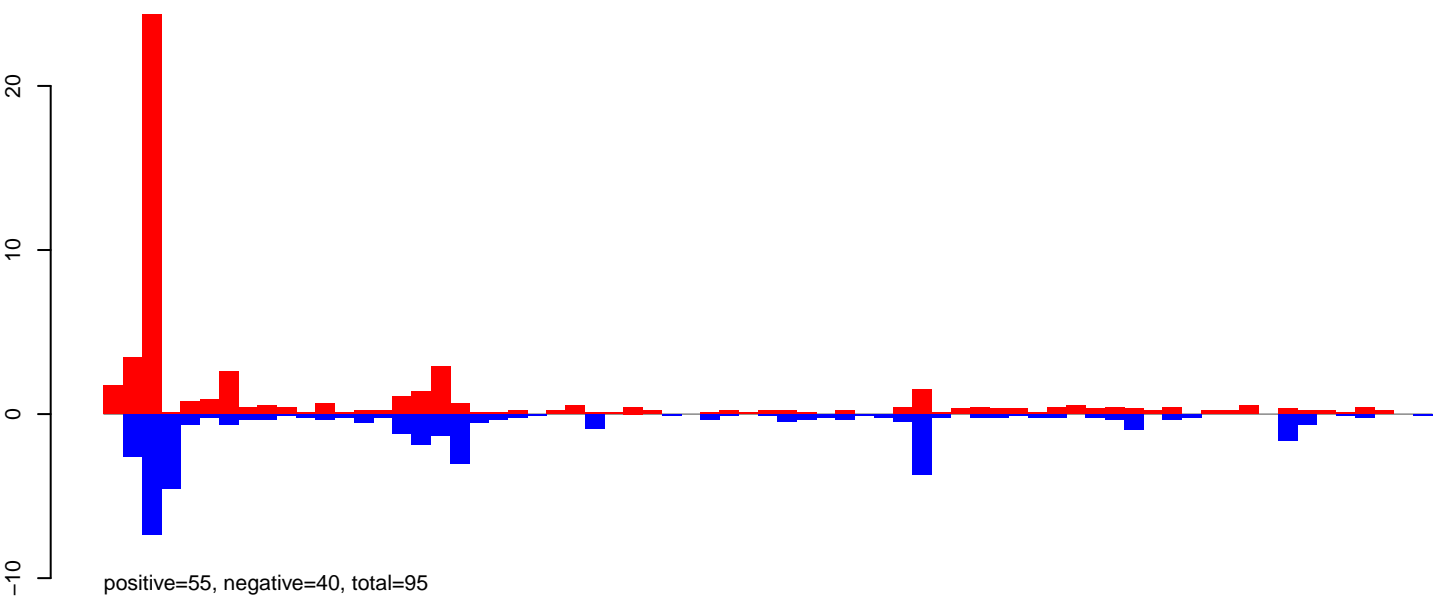
AnGam\_Sua5b\_cells.rep



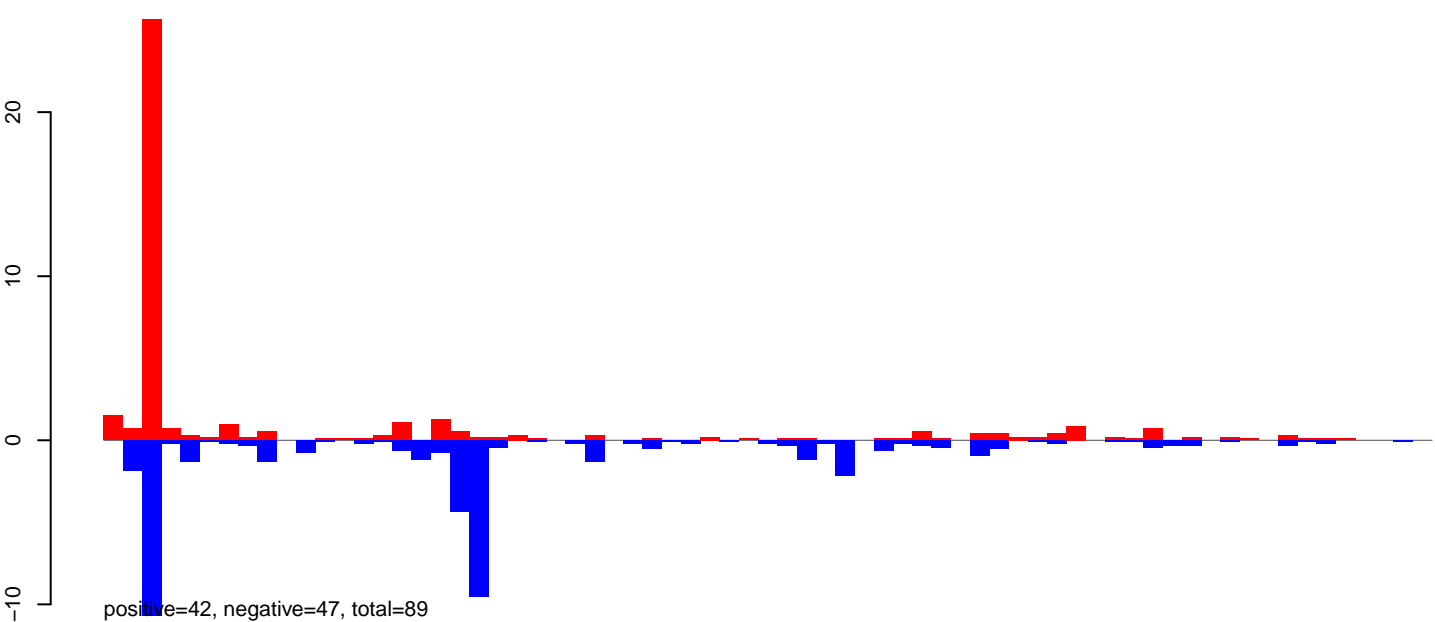
positive=148, negative=120, total=268 window size=50, length=4139, virus@NC\_011317.1-Anopheles\_gambiae\_densonucleosis\_virus:1-4139

0 1000 2000 3000 4000

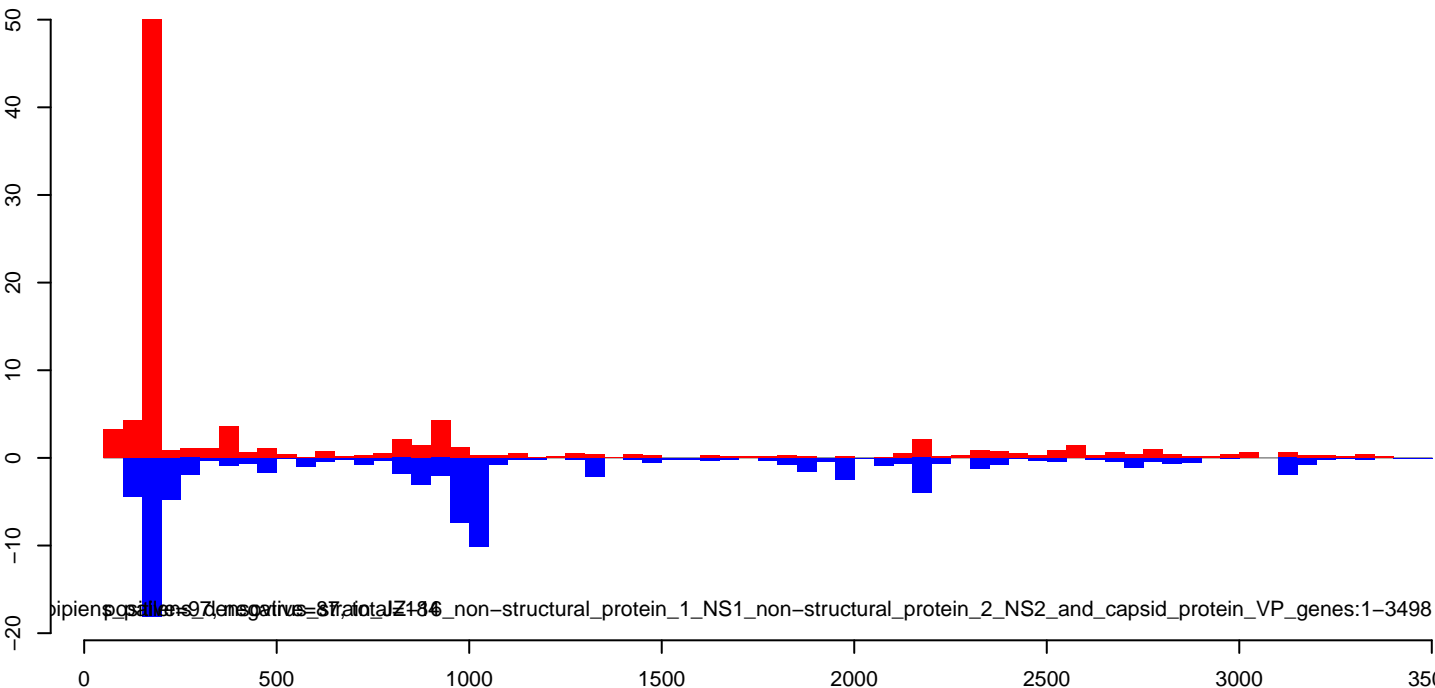
AnGam\_Sua5b\_cells.18\_23.rep



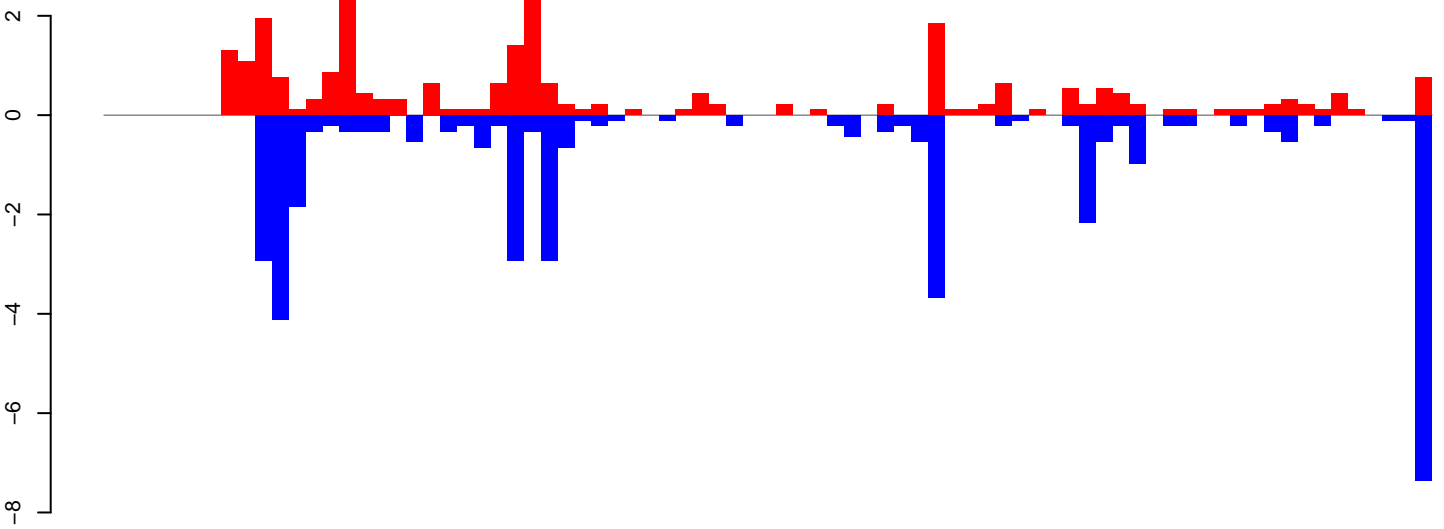
AnGam\_Sua5b\_cells.24\_35.rep



AnGam\_Sua5b\_cells.rep

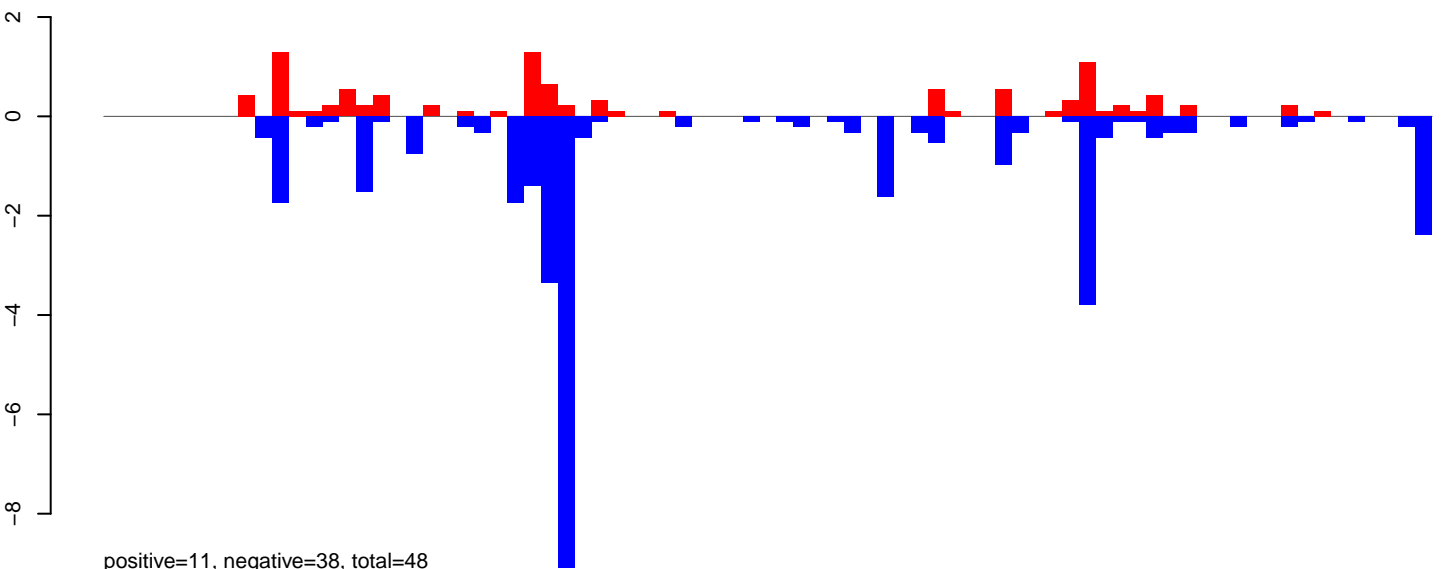


AnGam\_Sua5b\_cells.18\_23.rep



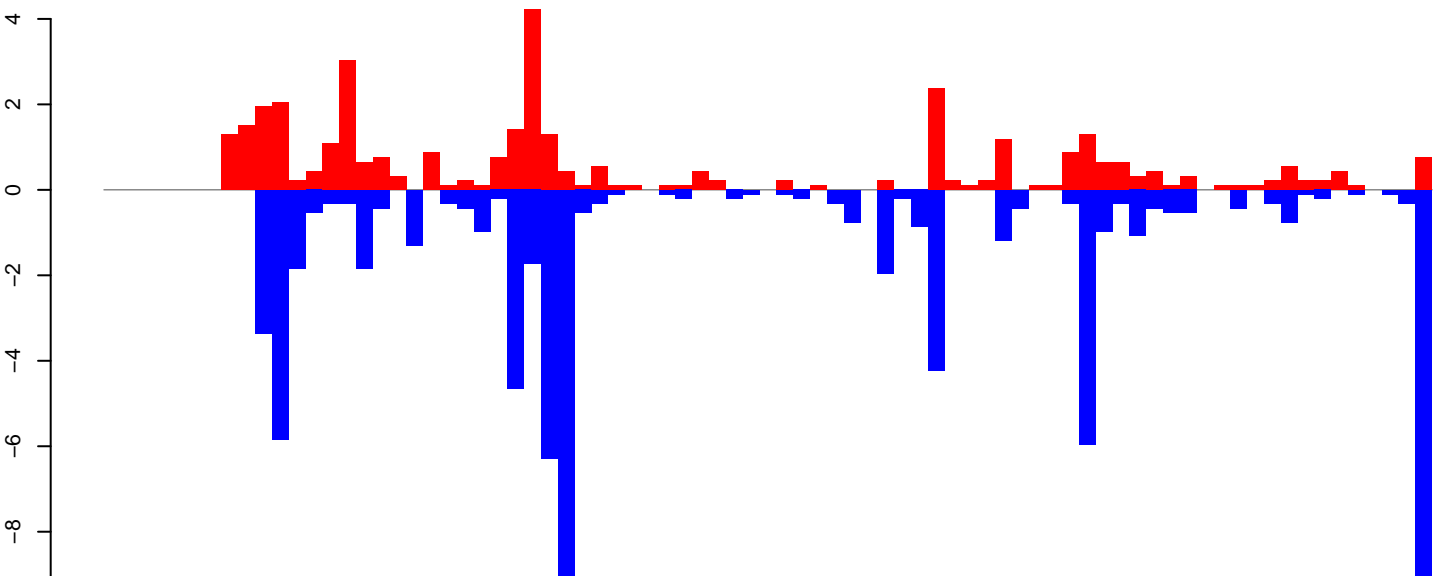
positive=27, negative=46, total=73

AnGam\_Sua5b\_cells.24\_35.rep



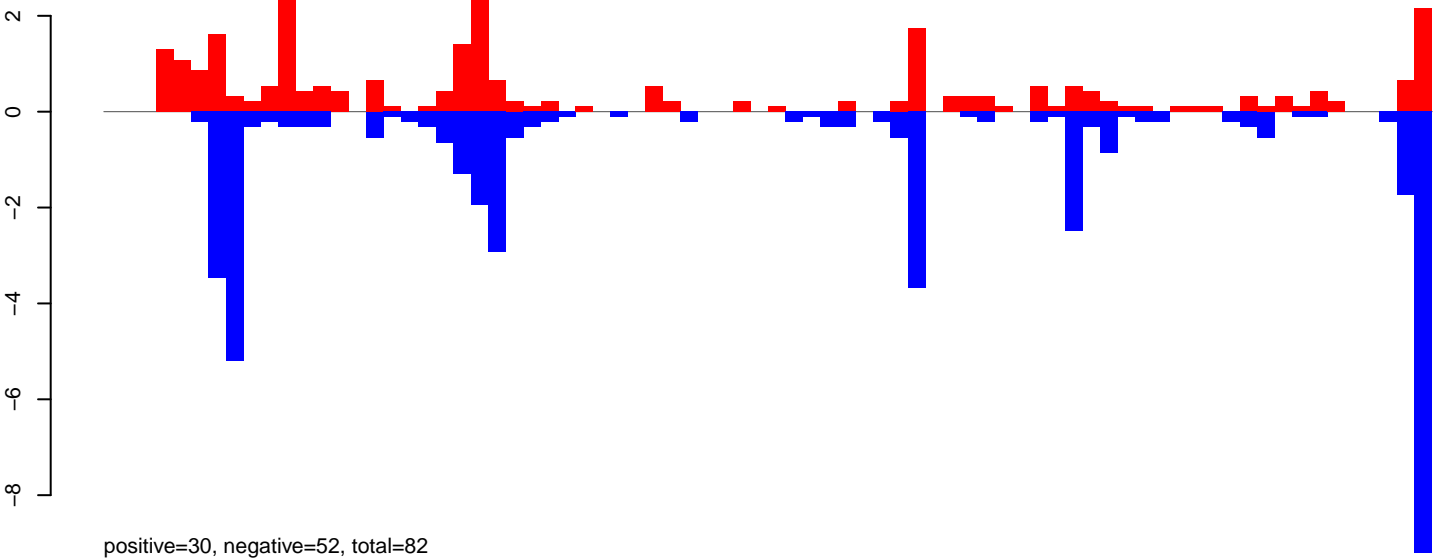
positive=11, negative=38, total=48

AnGam\_Sua5b\_cells.rep

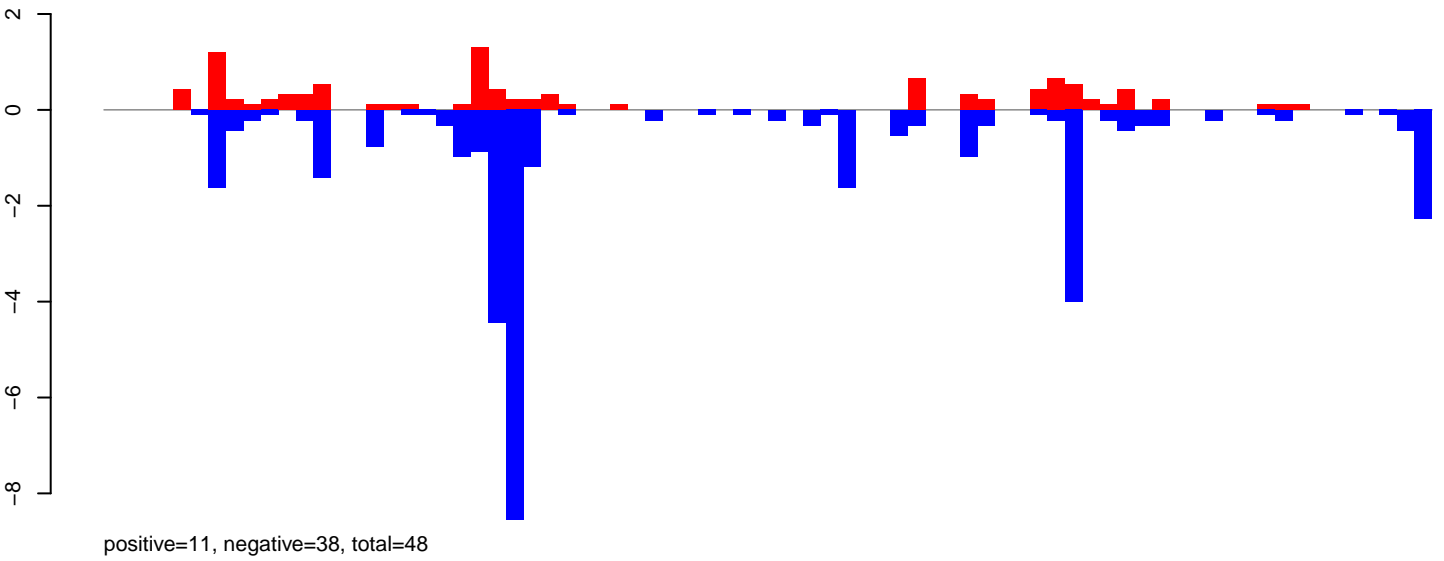


positive=38, negative=84, total=122, window size=50, length=3991, Virus@MH188044.1-Culex\_densovirus\_strain\_CDENV\_Madera-3:1-3991

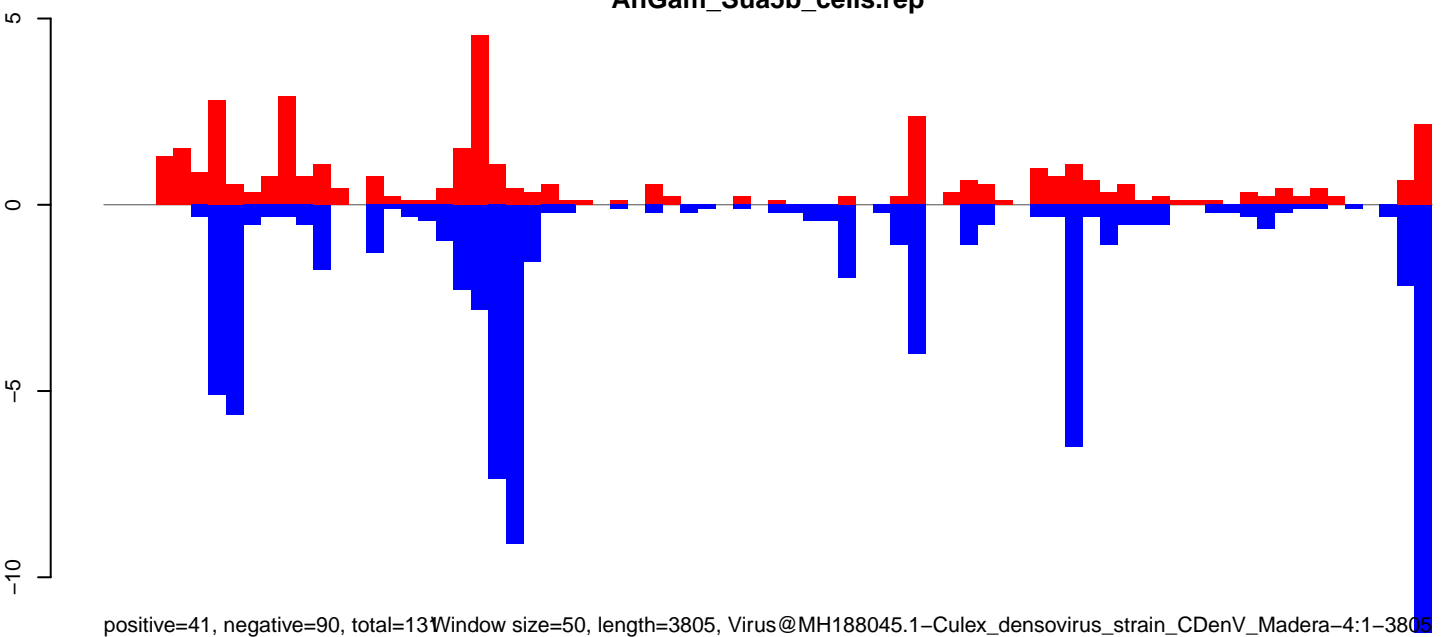
AnGam\_Sua5b\_cells.18\_23.rep



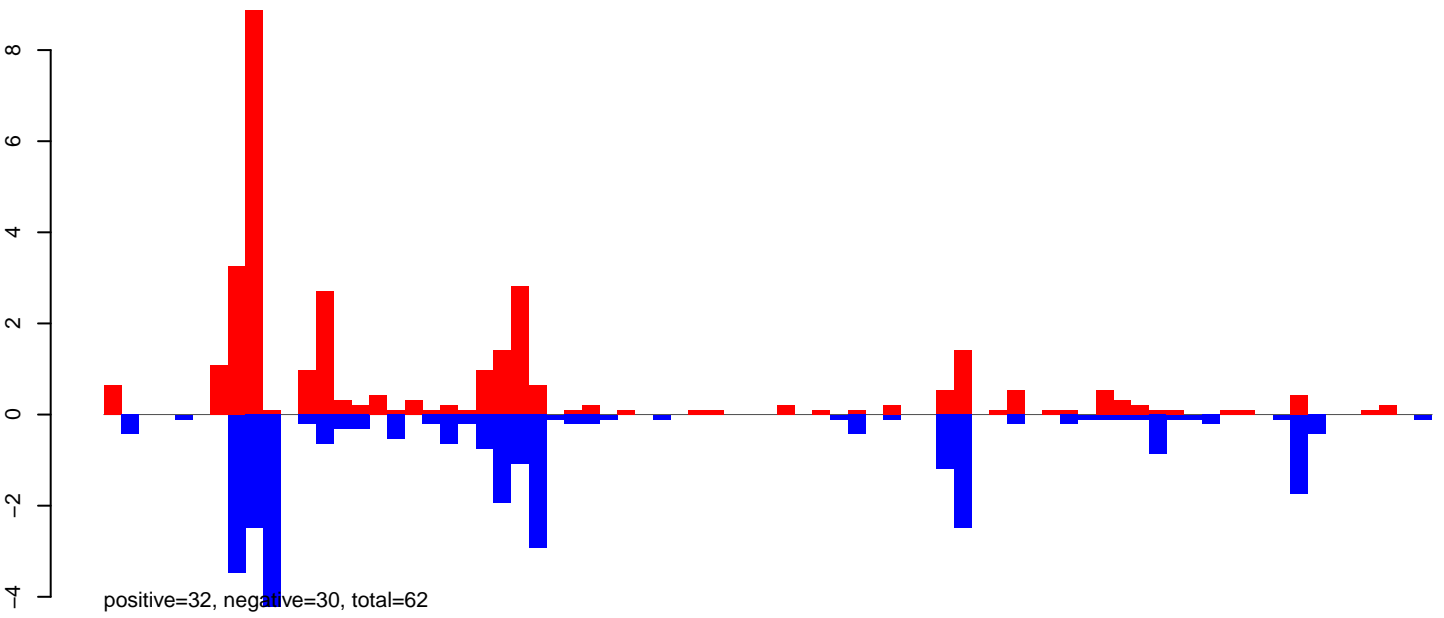
AnGam\_Sua5b\_cells.24\_35.rep



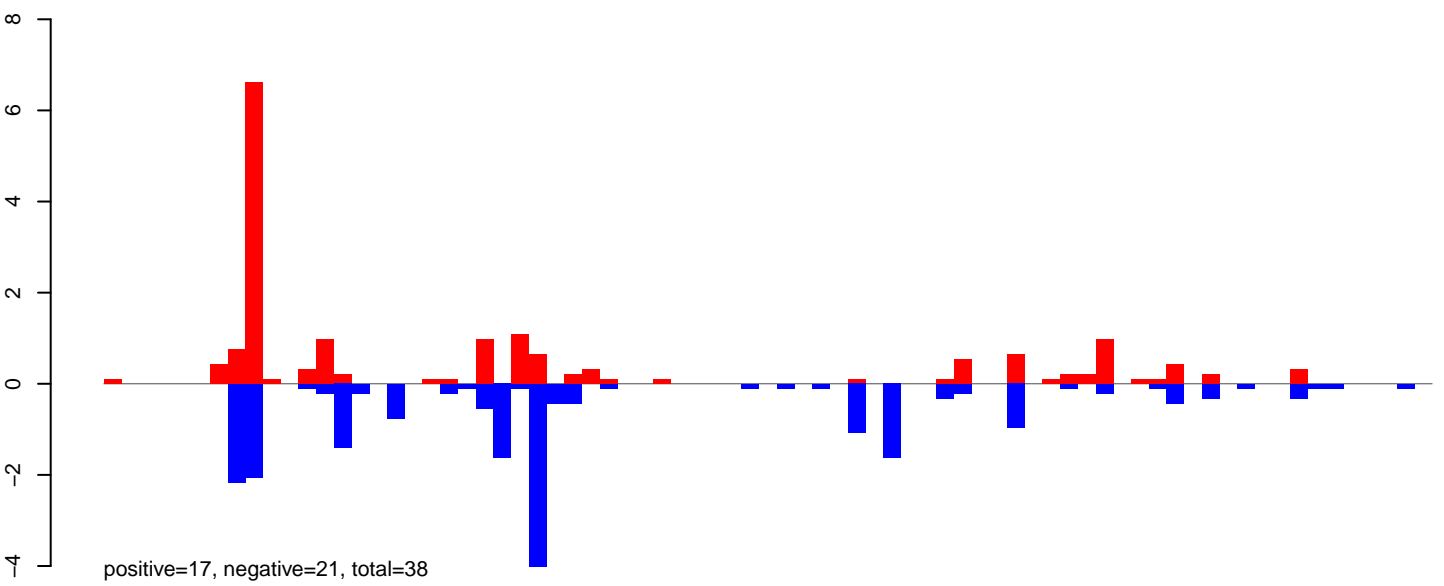
AnGam\_Sua5b\_cells.rep



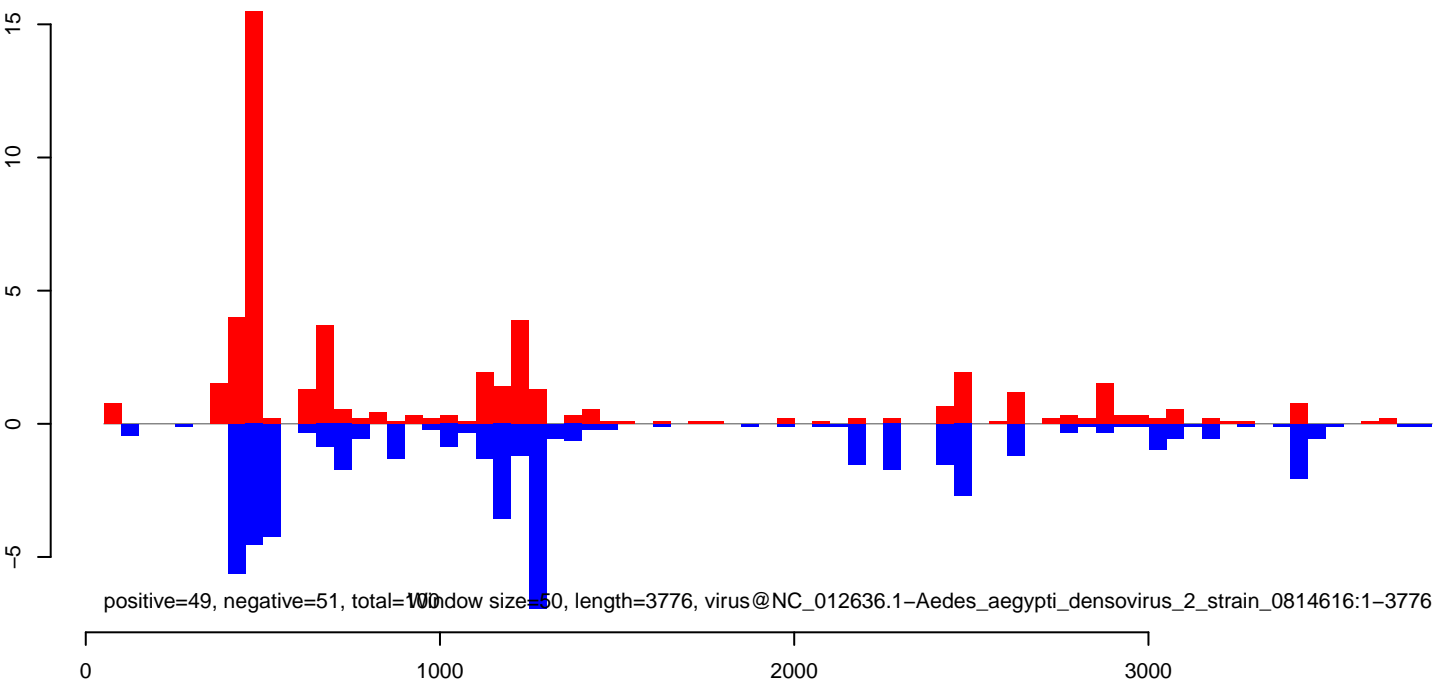
AnGam\_Sua5b\_cells.18\_23.rep



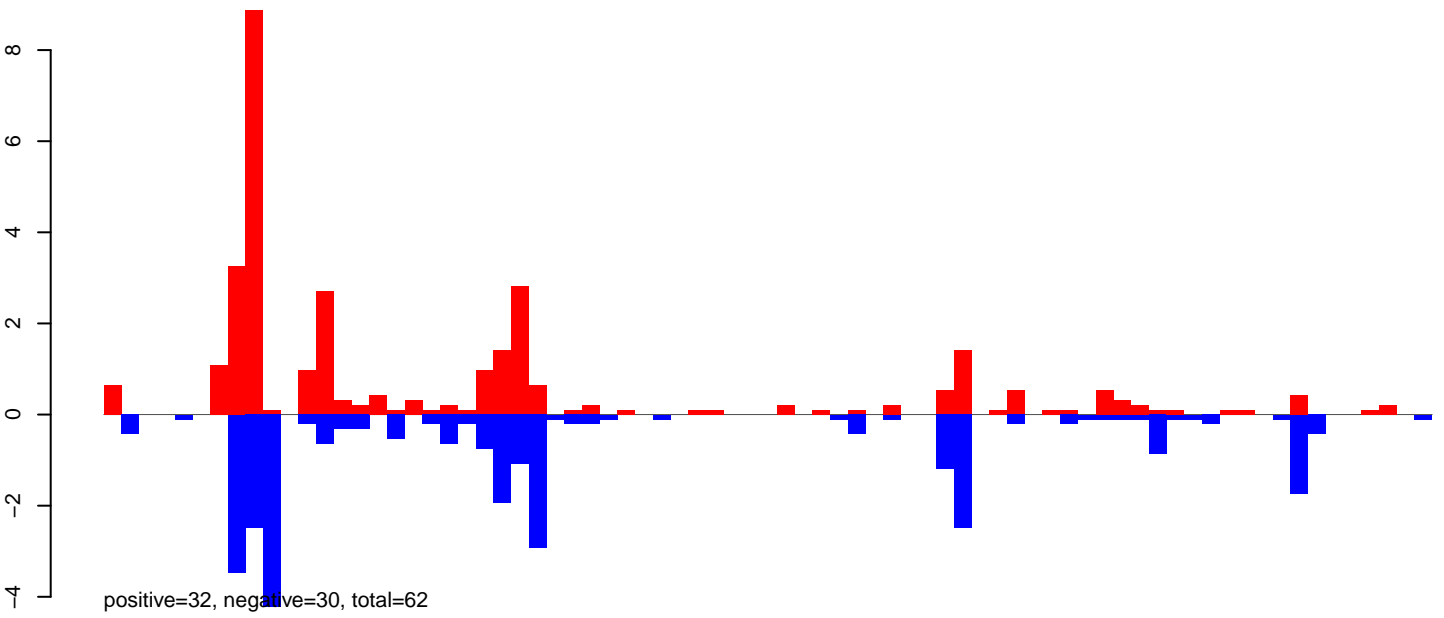
AnGam\_Sua5b\_cells.24\_35.rep



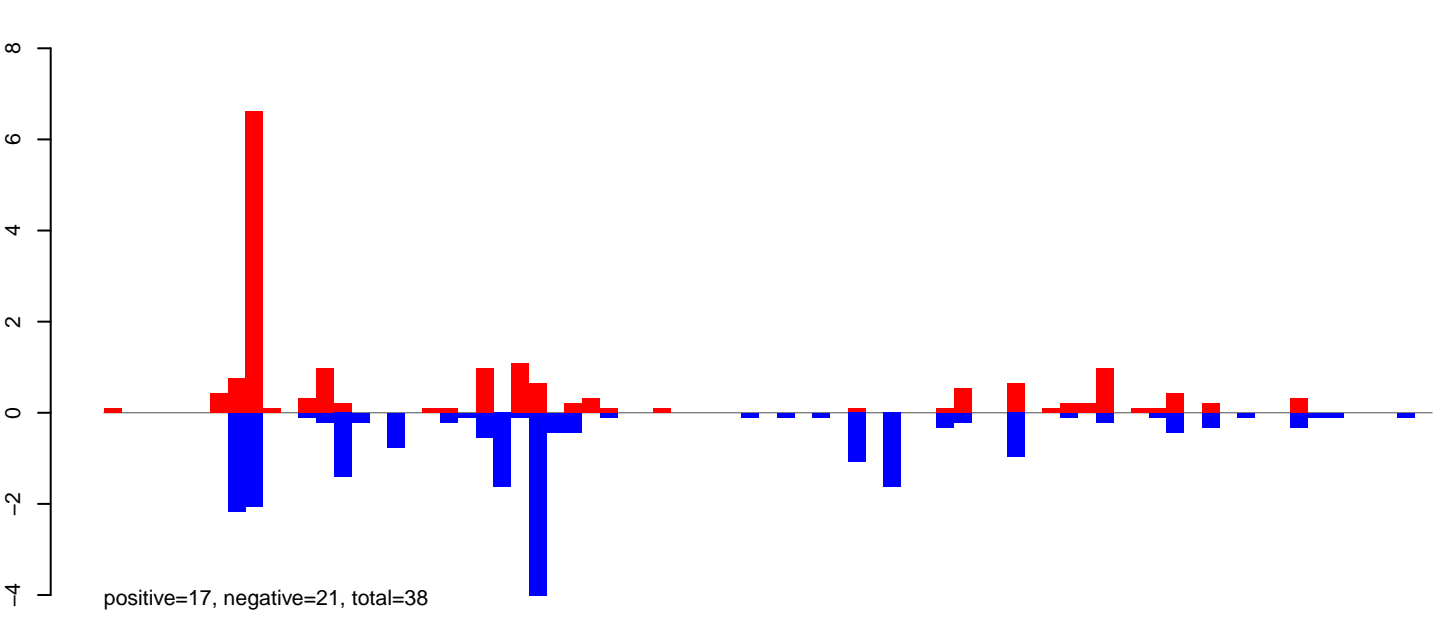
AnGam\_Sua5b\_cells.rep



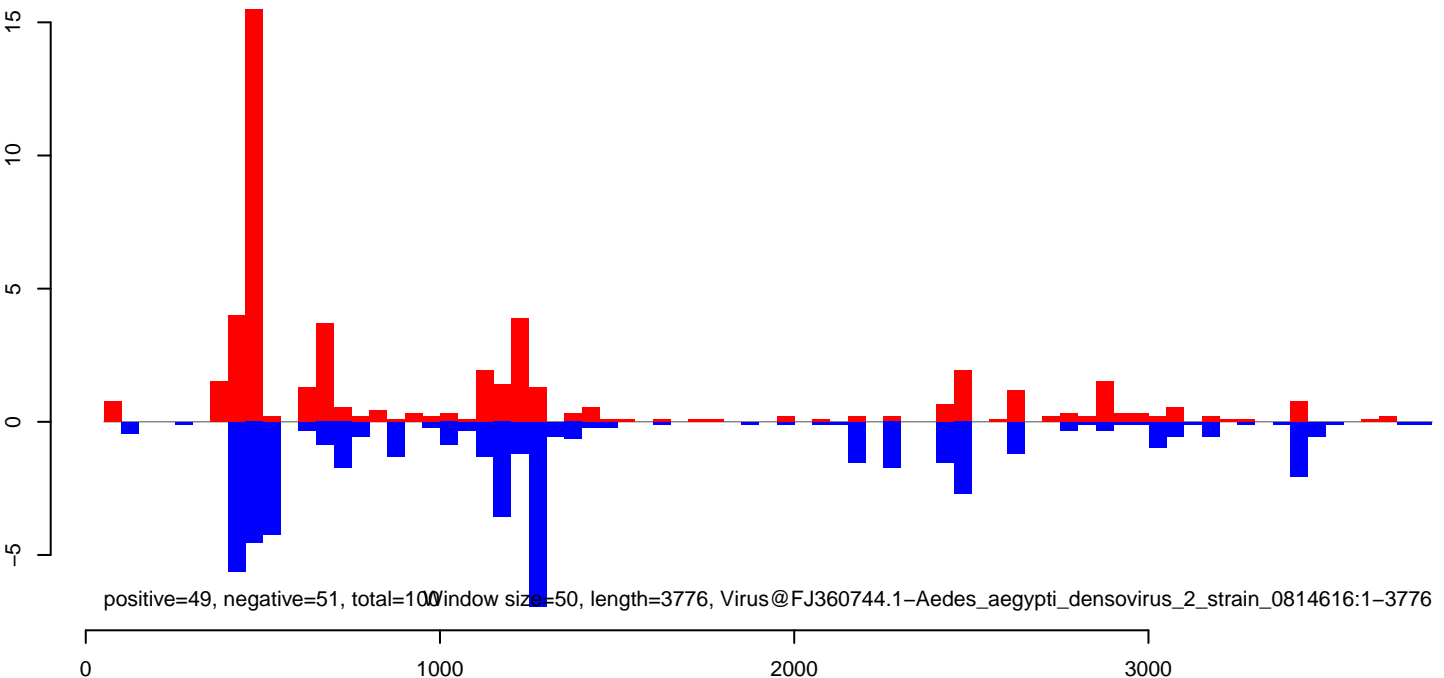
AnGam\_Sua5b\_cells.18\_23.rep



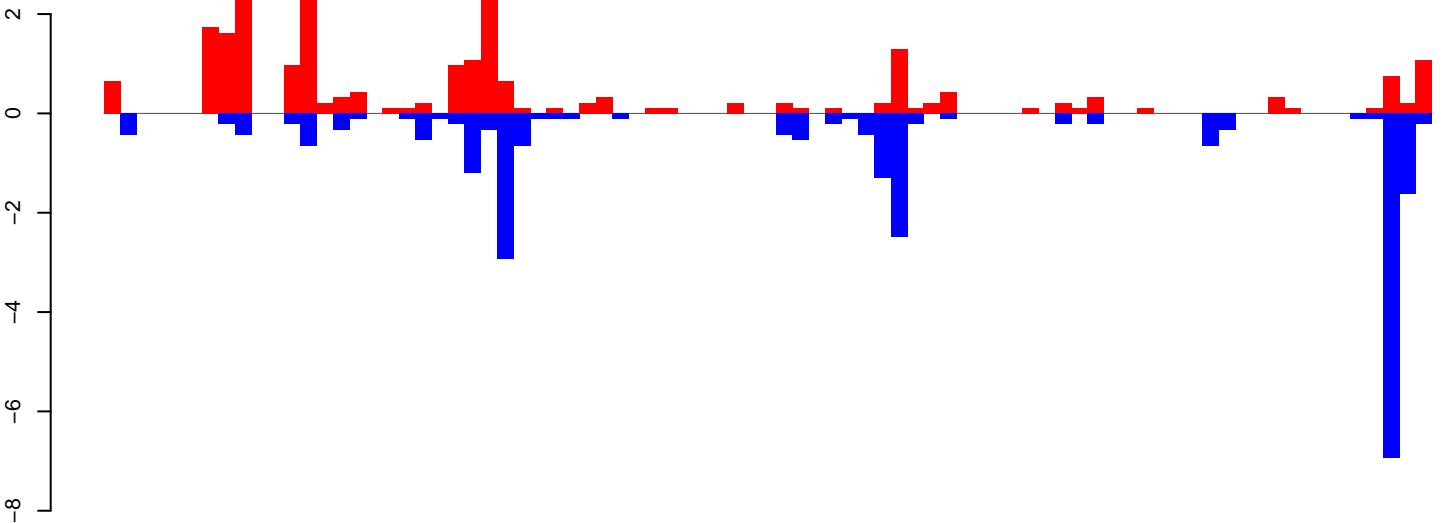
AnGam\_Sua5b\_cells.24\_35.rep



AnGam\_Sua5b\_cells.rep

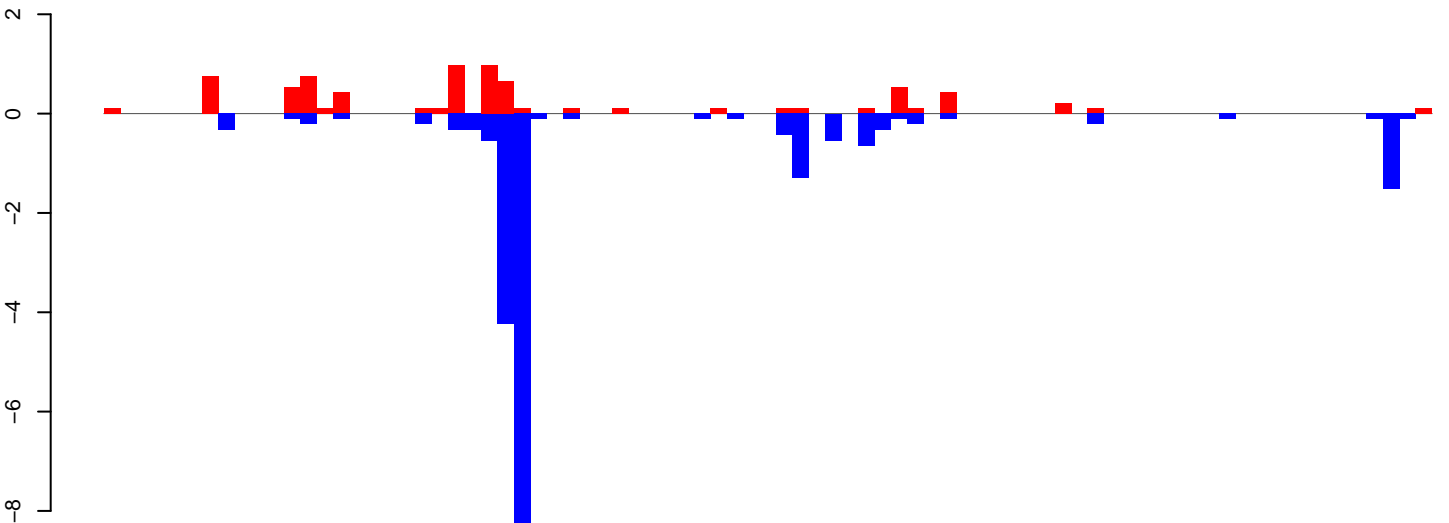


AnGam\_Sua5b\_cells.18\_23.rep



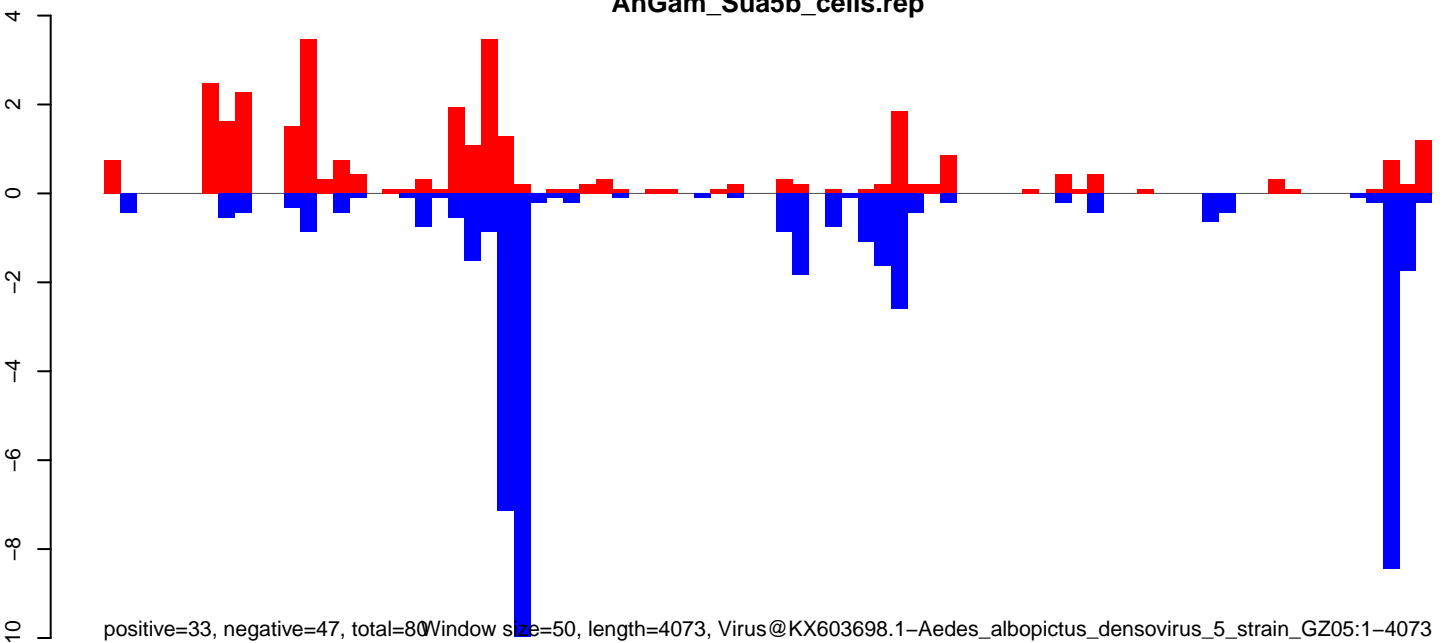
positive=25, negative=25, total=50

AnGam\_Sua5b\_cells.24\_35.rep



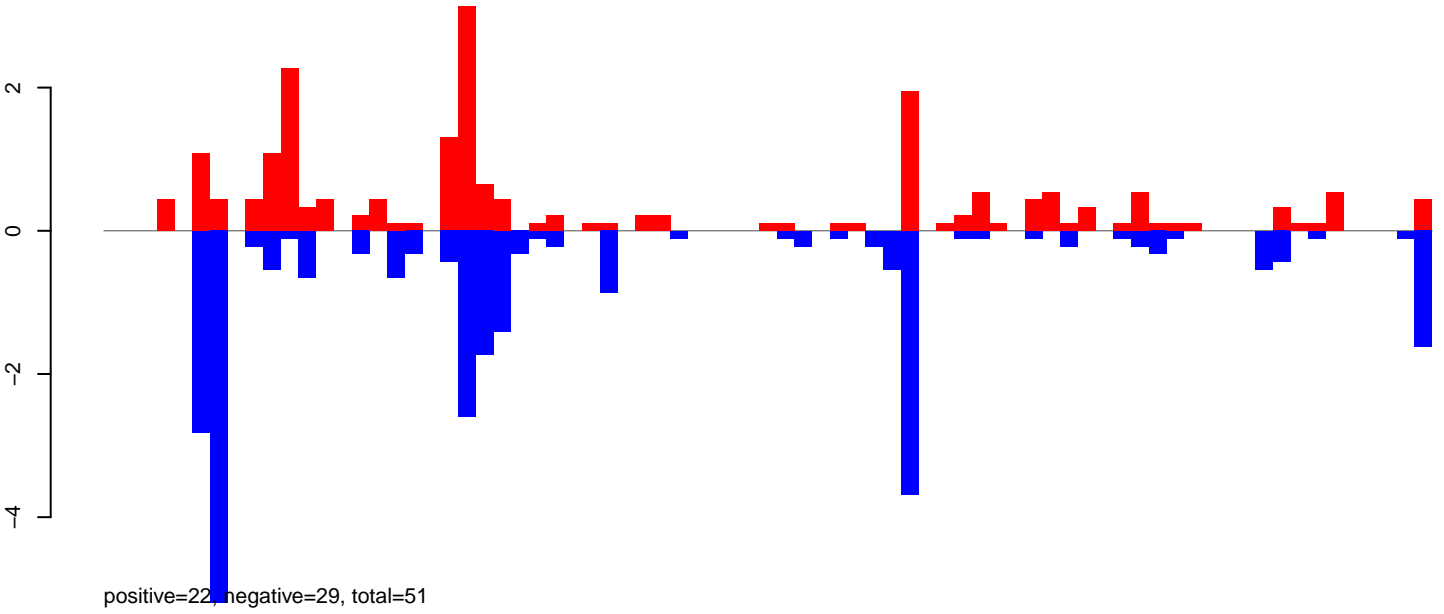
positive=8, negative=22, total=30

AnGam\_Sua5b\_cells.rep

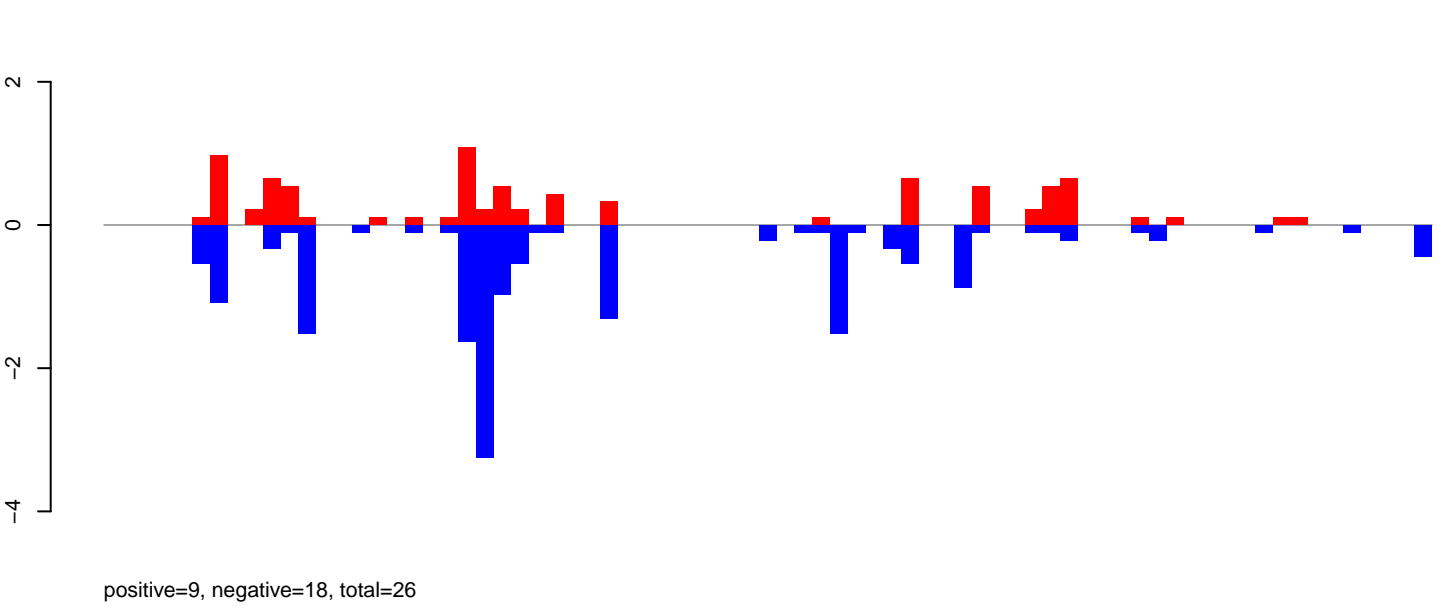


positive=33, negative=47, total=80, window size=50, length=4073, Virus@KX603698.1-Aedes\_albopictus\_densovirus\_5\_strain\_GZ05:1-4073

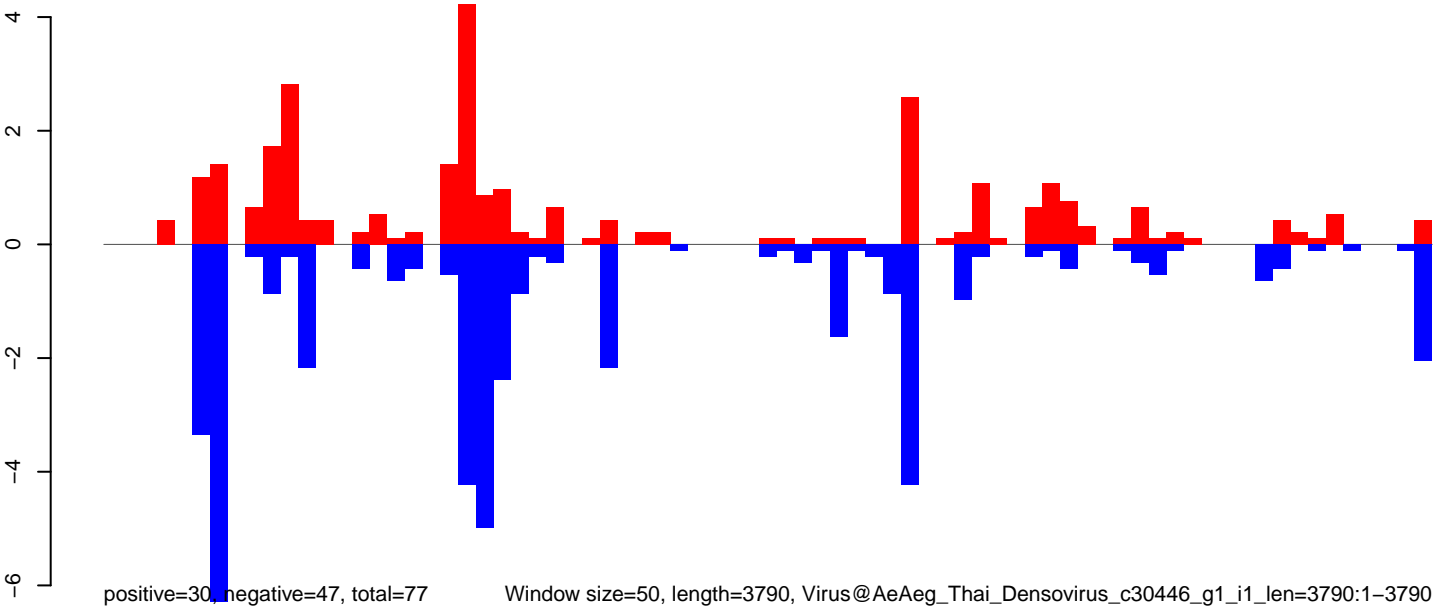
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep



AnGam\_Sua5b\_cells.rep

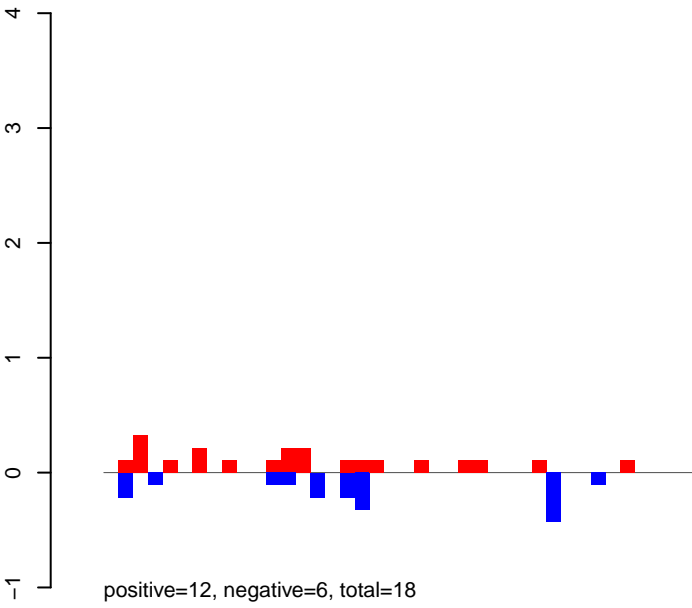


Window size=50, length=3790, Virus@AeAeg\_Thai\_Densovirus\_c30446\_g1\_i1\_len=3790:1-3790

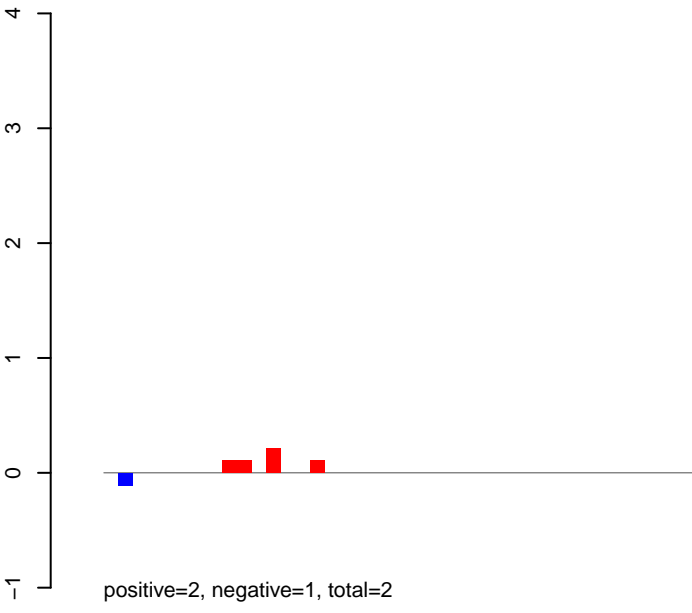
0 1000 2000 3000



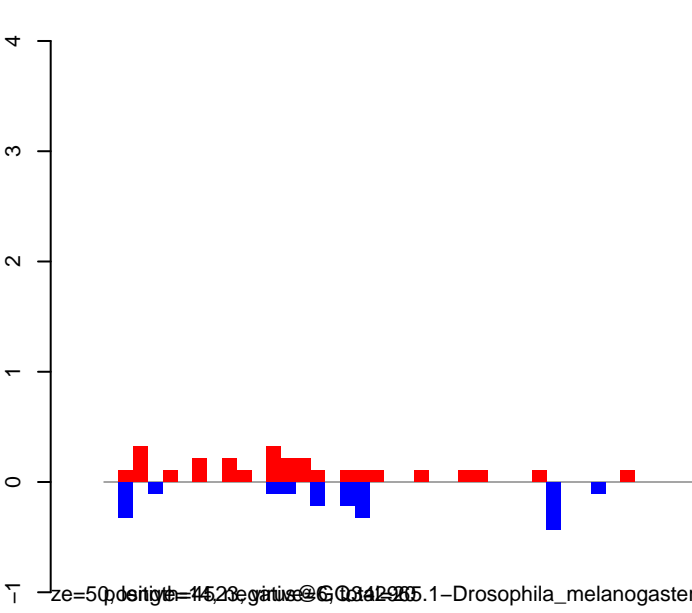
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep

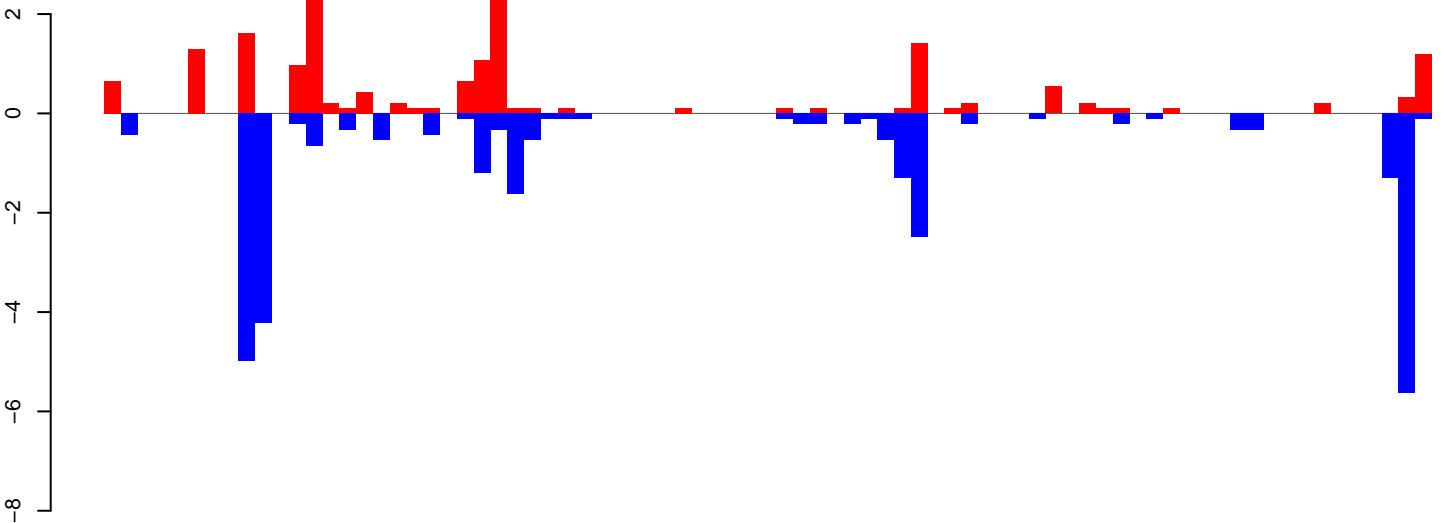


AnGam\_Sua5b\_cells.rep



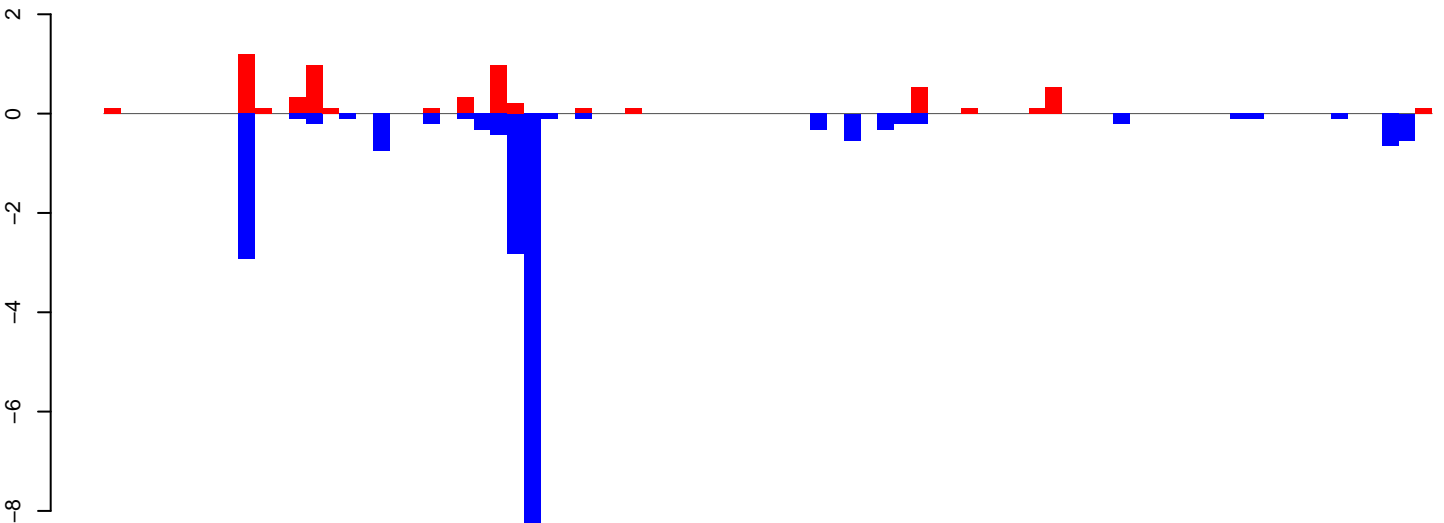
z=50 positive=15, negative=6, total=21 -Drosophila\_melanogaster\_American\_nodavirus\_ANV\_SW-2009a\_segment\_RNA1\_and\_RNA2:1-4523

AnGam\_Sua5b\_cells.18\_23.rep



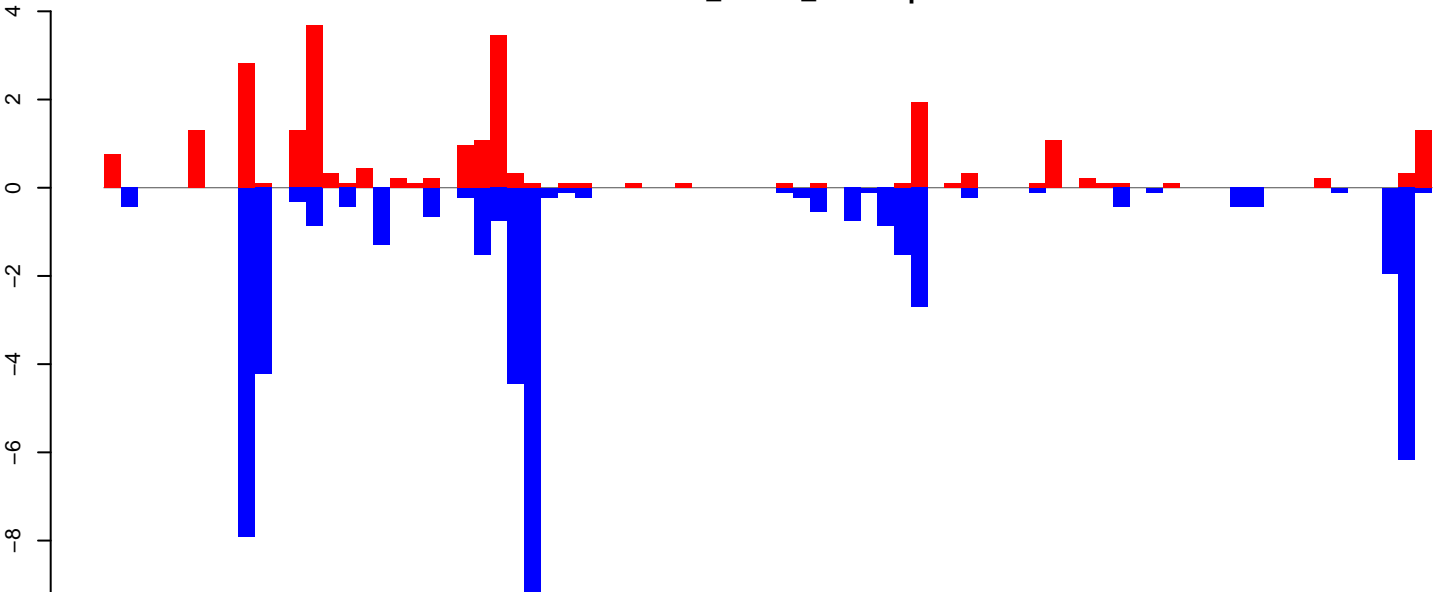
positive=19, negative=30, total=49

AnGam\_Sua5b\_cells.24\_35.rep



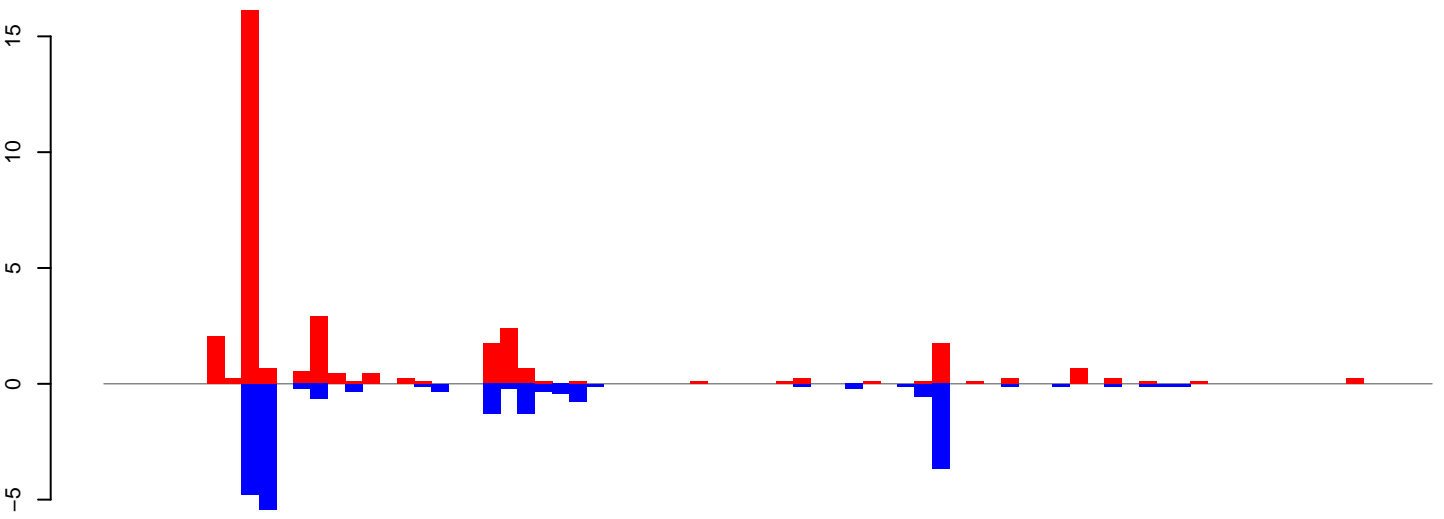
positive=6, negative=21, total=27

AnGam\_Sua5b\_cells.rep



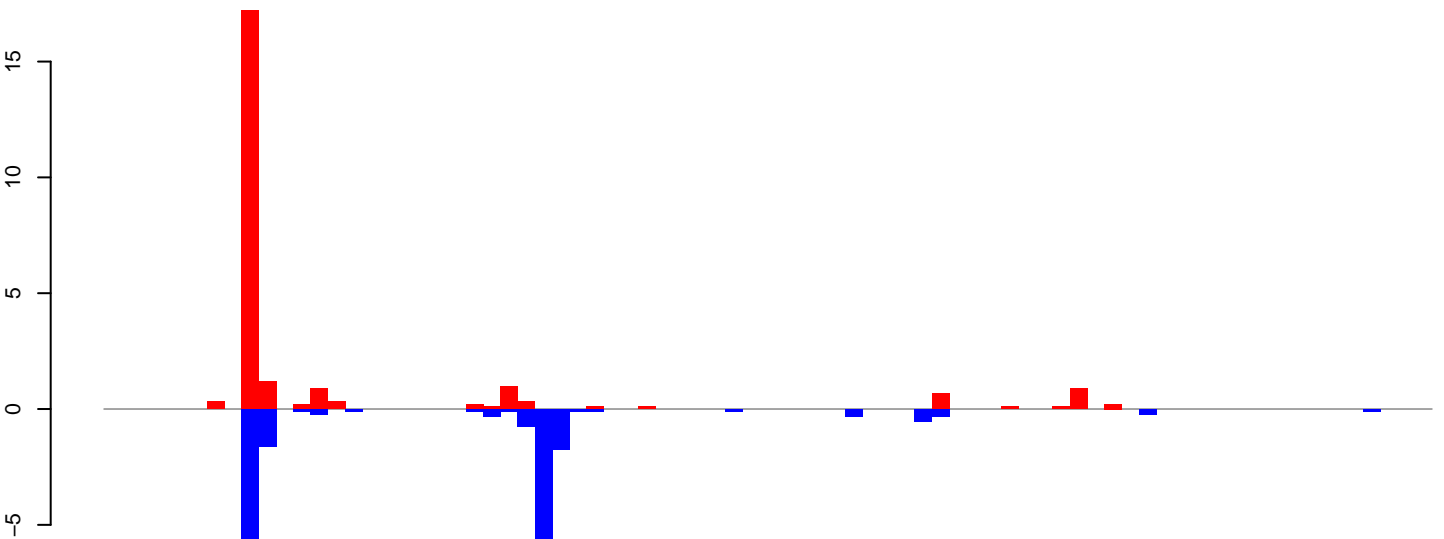
positive=25, negative=50, total=75, window size=50, length=3978, Virus@M37899.1-Aedes\_aegypti\_densovirus\_1\_genomic\_sequence:1-3978

AnGam\_Sua5b\_cells.18\_23.rep



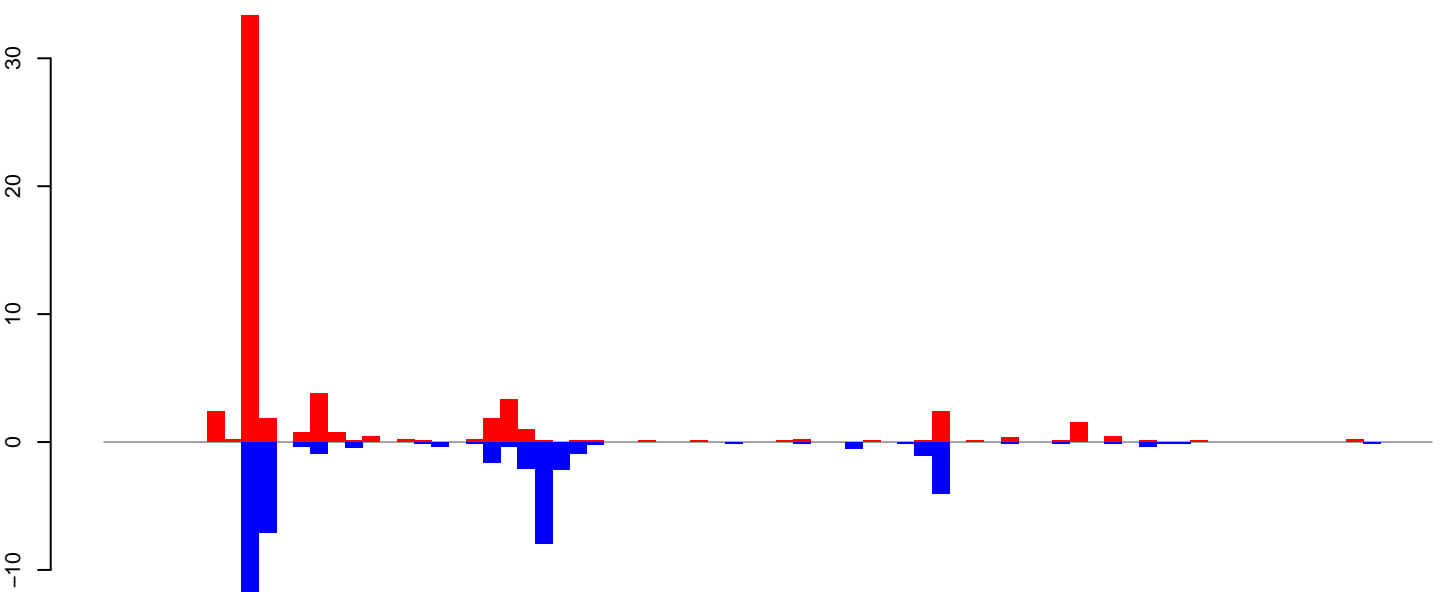
positive=33, negative=22, total=54

AnGam\_Sua5b\_cells.24\_35.rep



positive=24, negative=23, total=47

AnGam\_Sua5b\_cells.rep

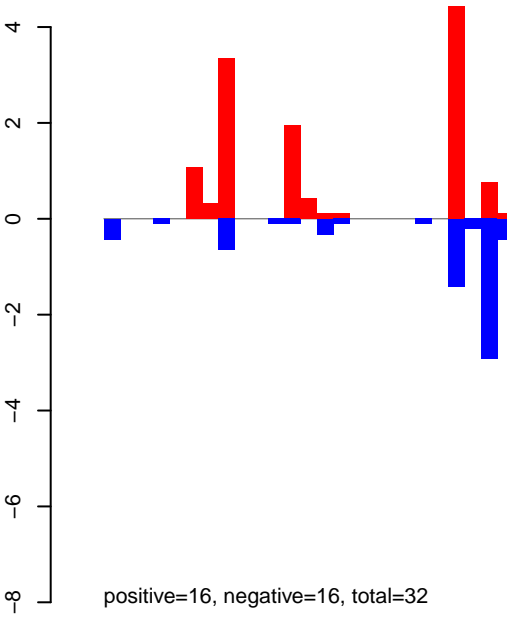


positive=57, negative=45, total=101

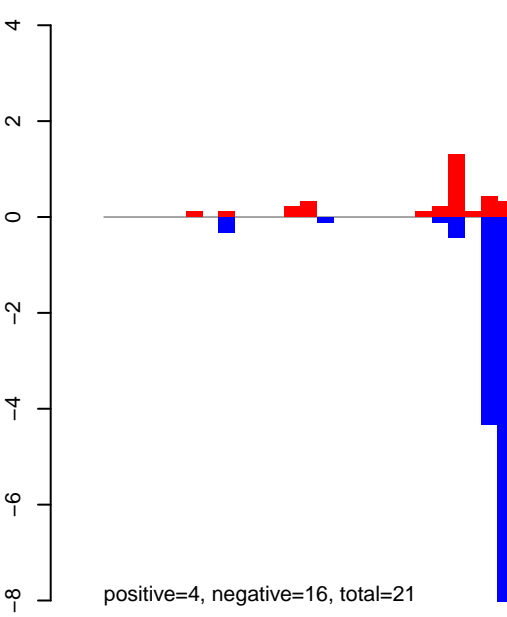
Window size=50, length=3893, Virus@GU452720.1-Mosquito\_densovirus\_BR\_07:1-3893

0 1000 2000 3000 4000

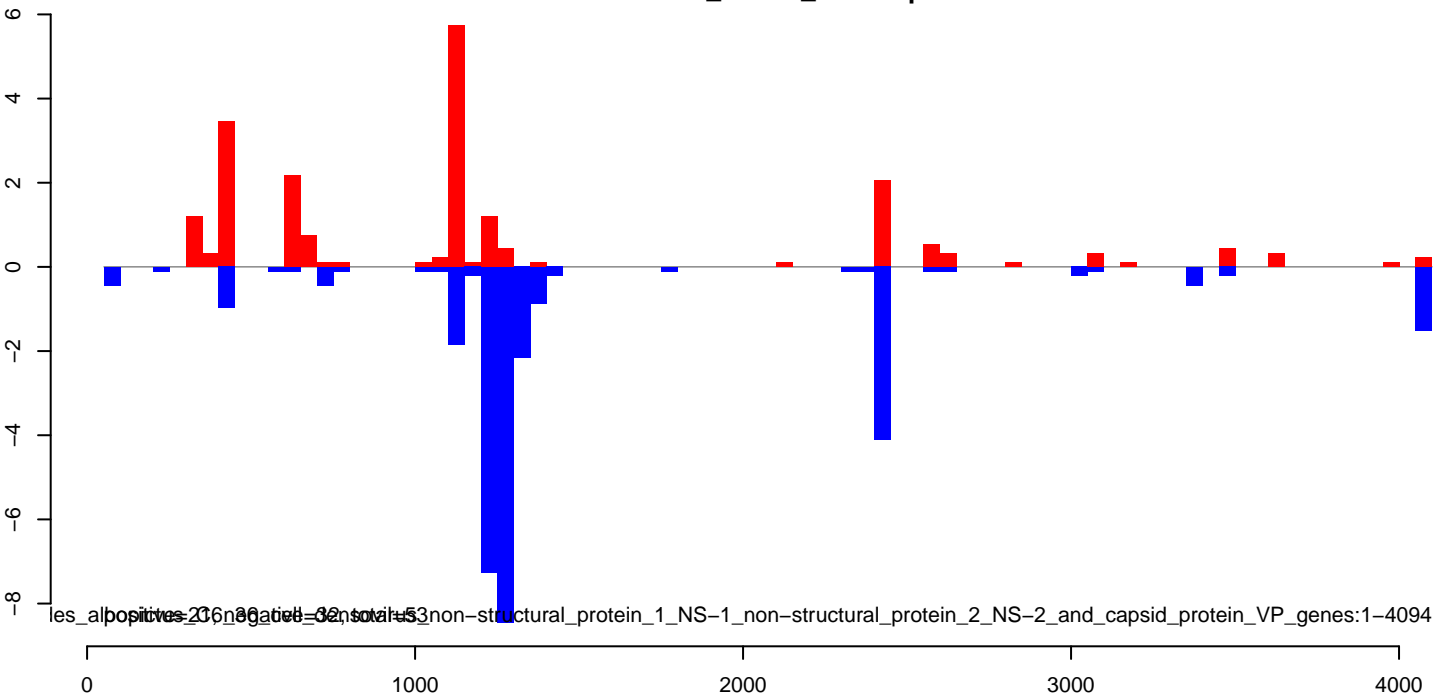
AnGam\_Sua5b\_cells.18\_23.rep



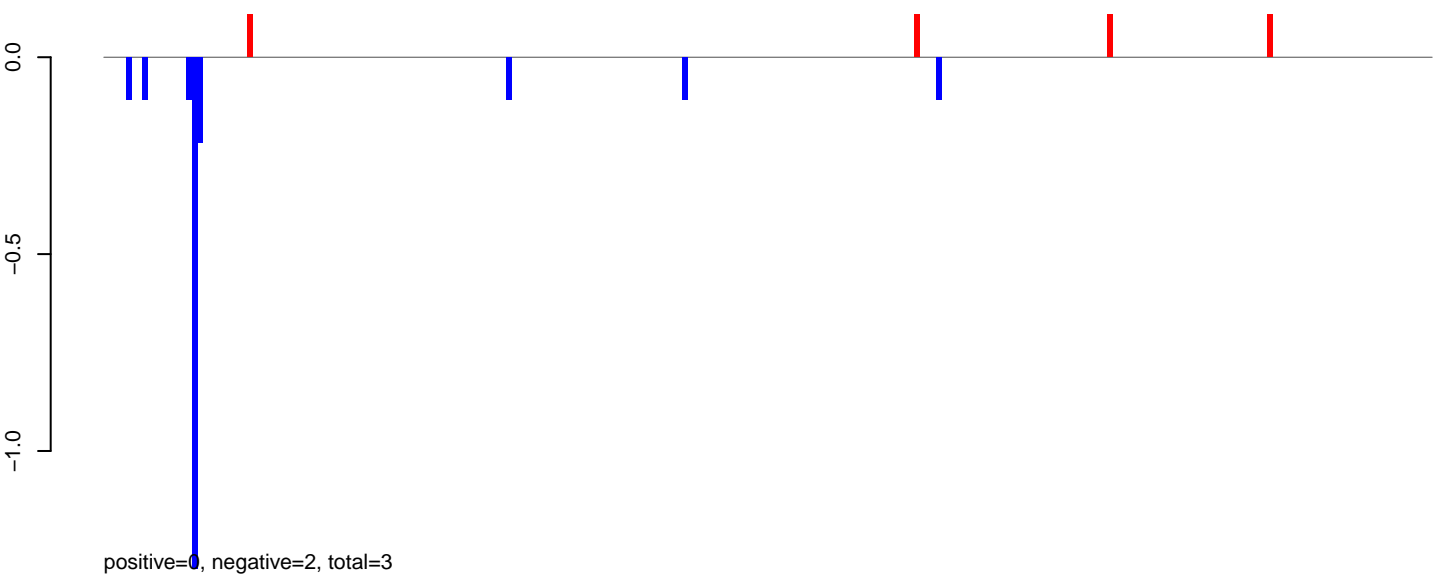
AnGam\_Sua5b\_cells.24\_35.rep



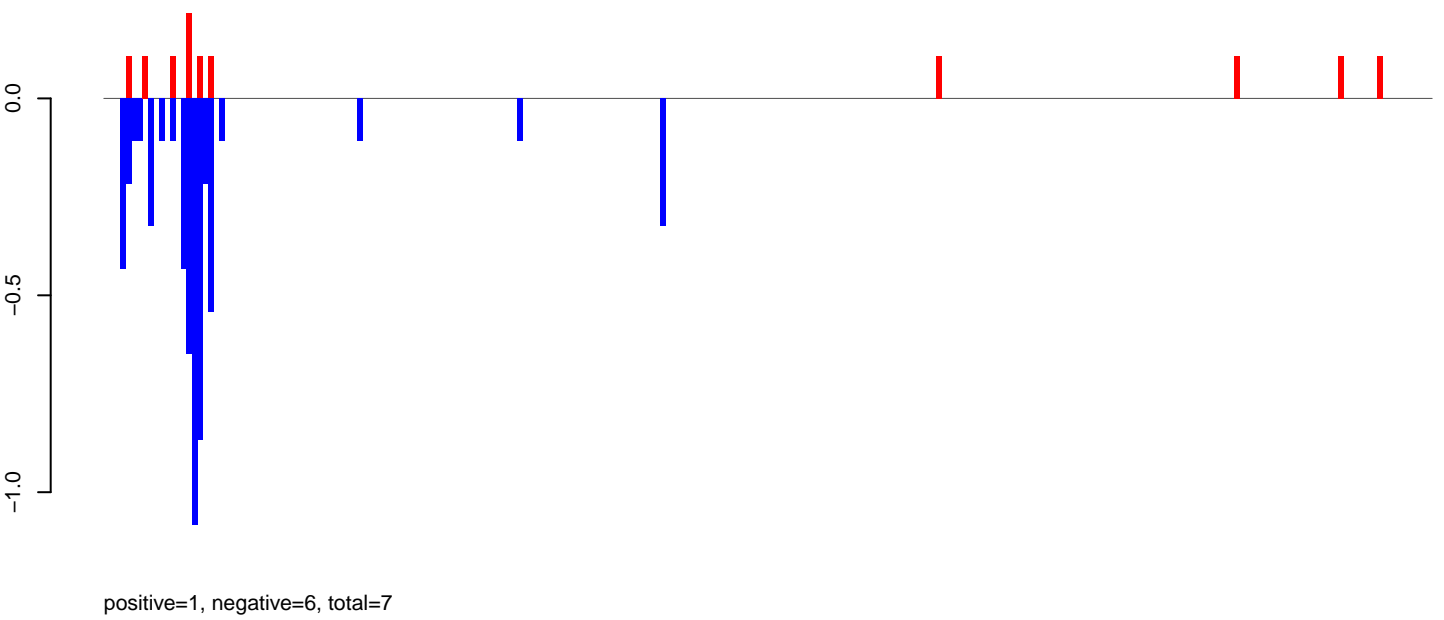
AnGam\_Sua5b\_cells.rep



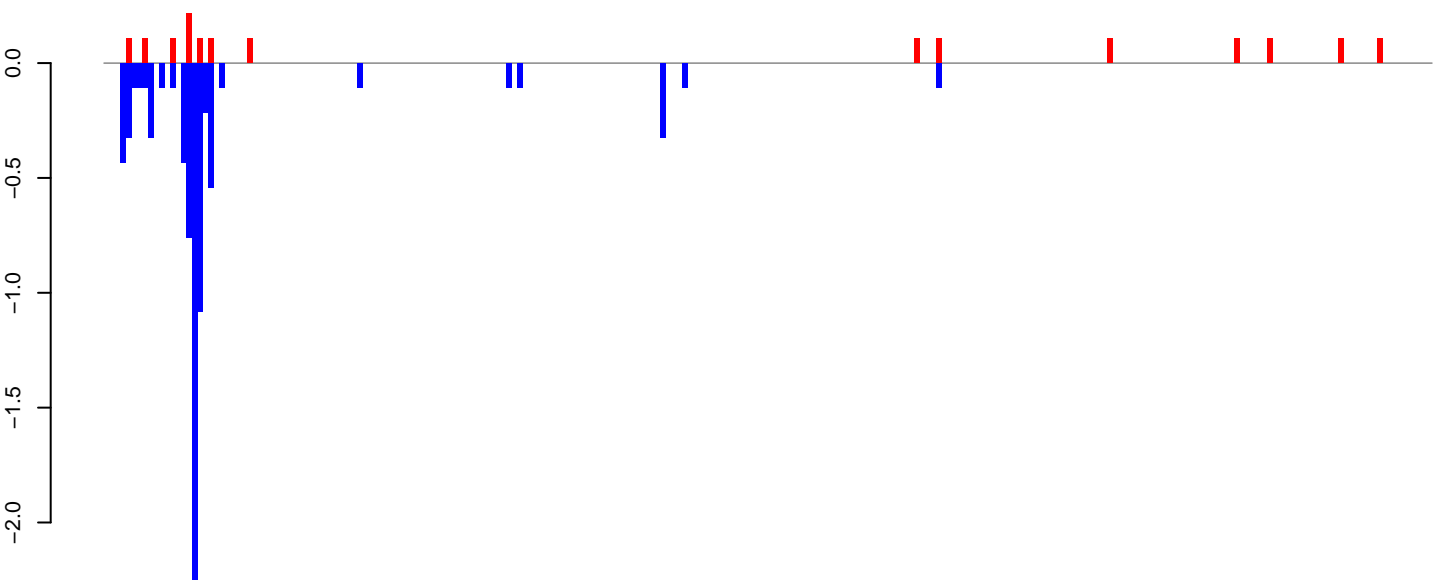
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep



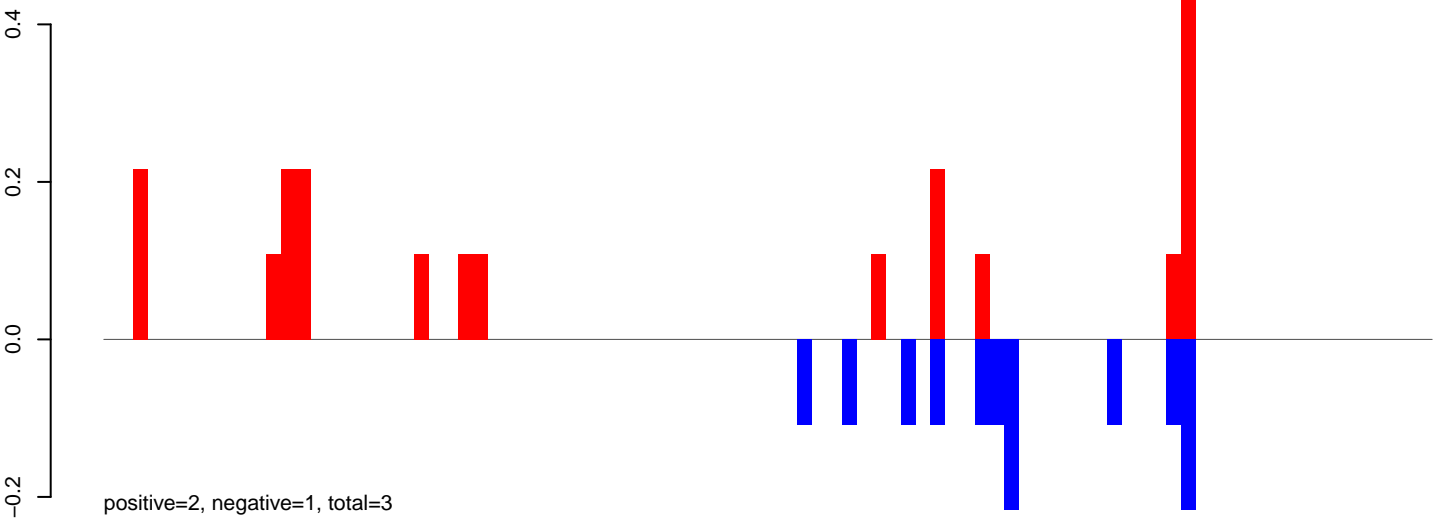
AnGam\_Sua5b\_cells.rep



2099, positive=42, negative=148, total=190. Charoen-like\_virus\_isolate\_Rio\_segment\_S\_plus\_segment\_M\_plus\_segment\_L\_Nucleocapsid\_gene:1-12099

0 2000 4000 6000 8000 10000 12000

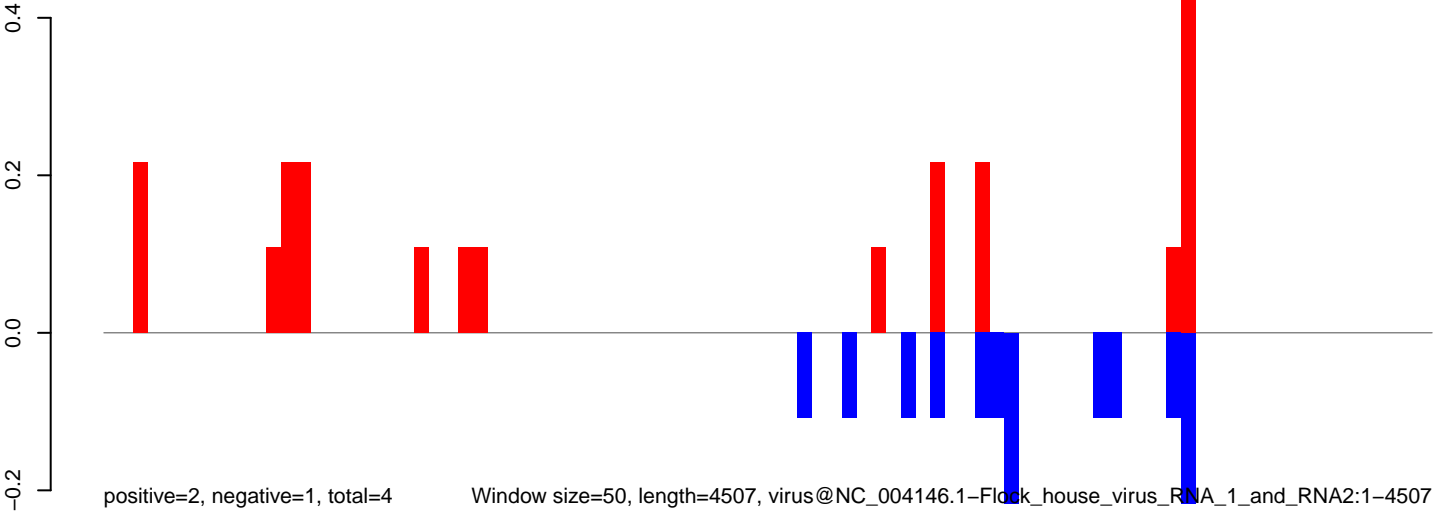
AnGam\_Sua5b\_cells.18\_23.rep



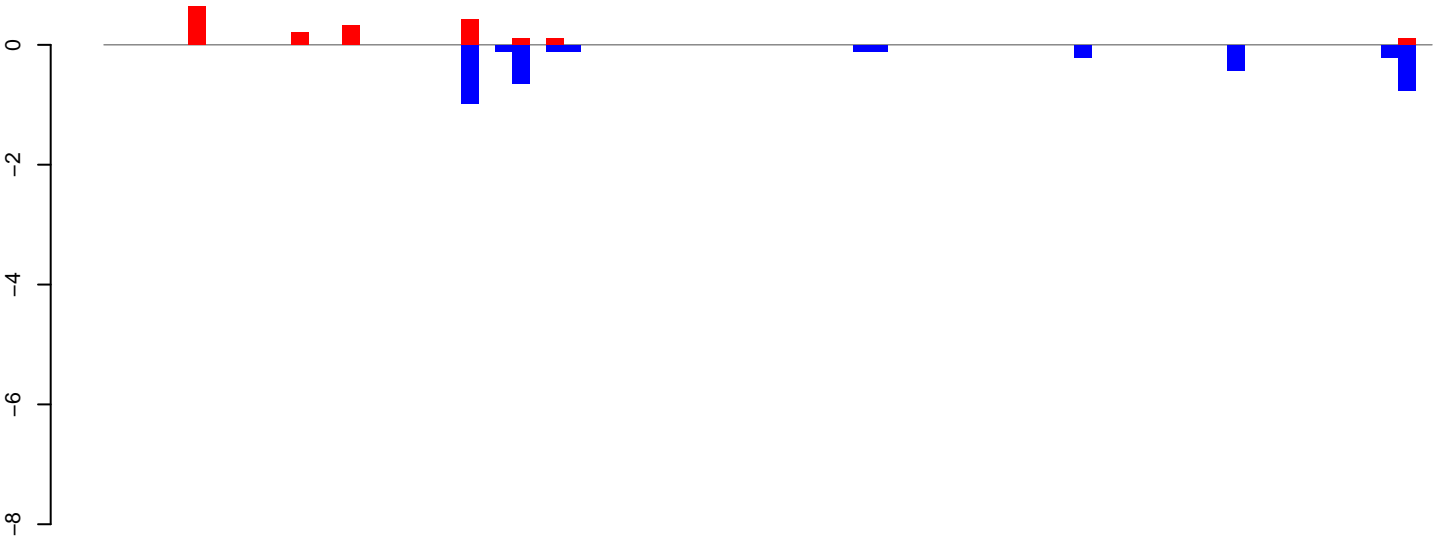
AnGam\_Sua5b\_cells.24\_35.rep



AnGam\_Sua5b\_cells.rep

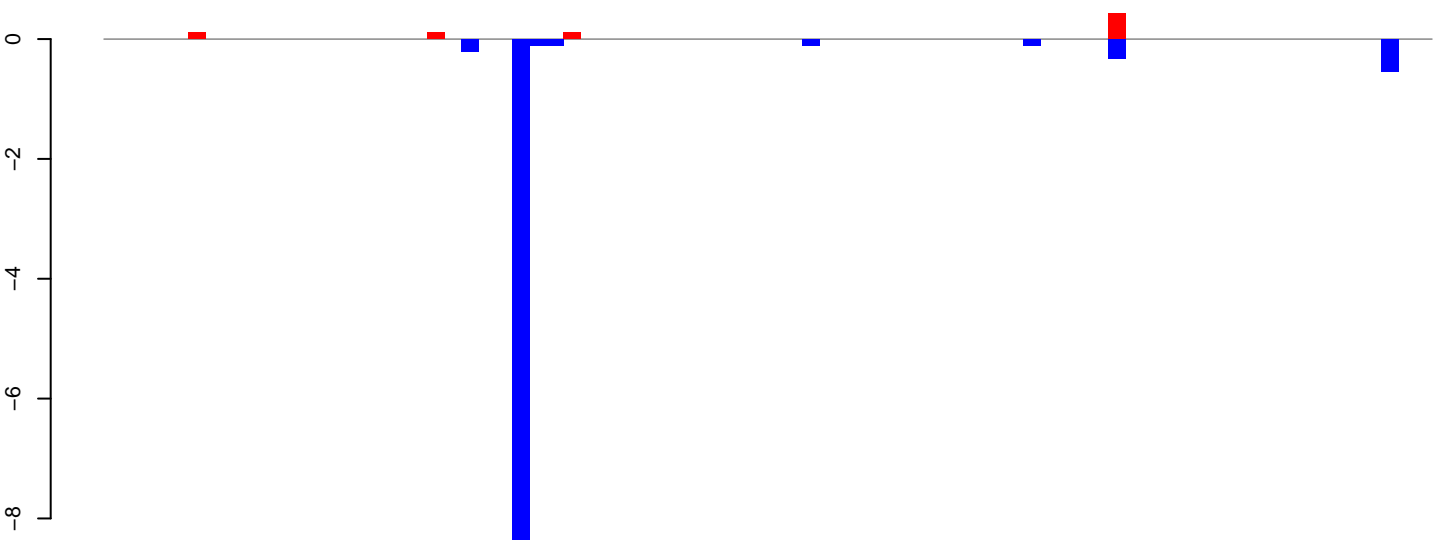


AnGam\_Sua5b\_cells.18\_23.rep



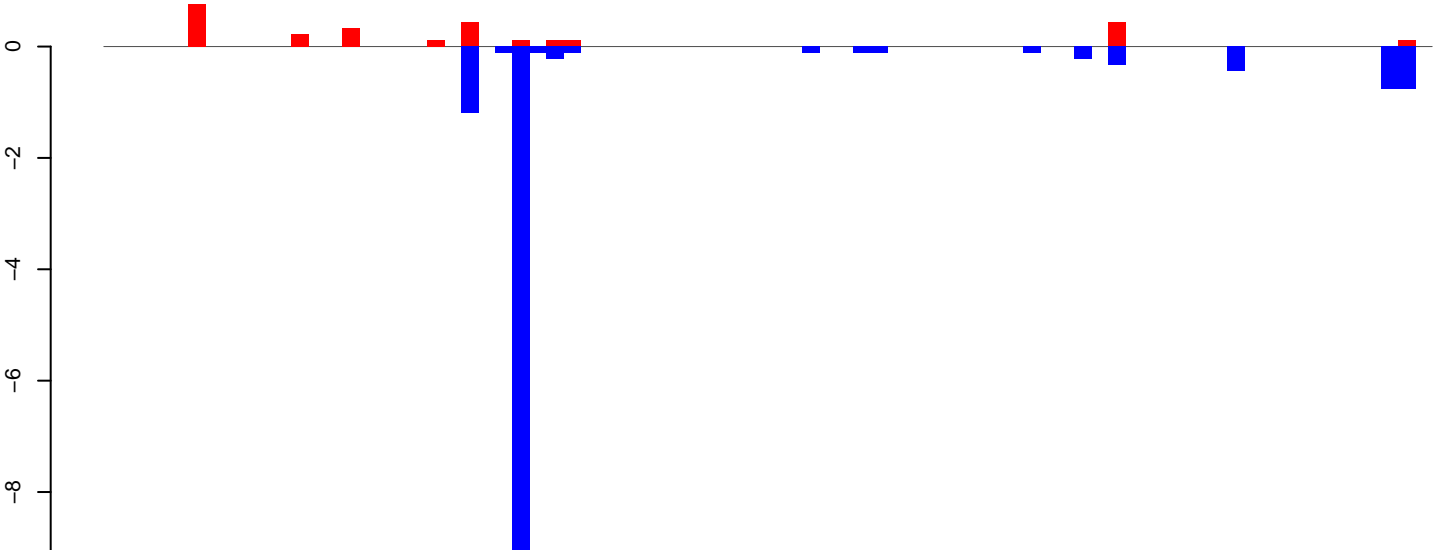
positive=2, negative=4, total=6

AnGam\_Sua5b\_cells.24\_35.rep



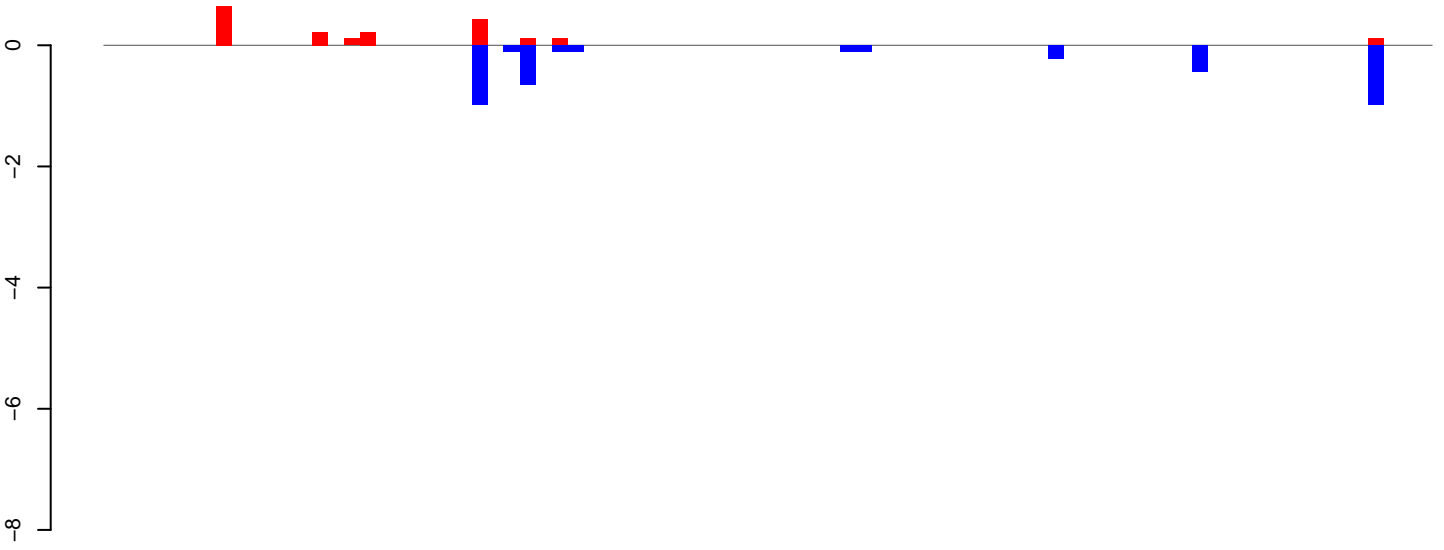
positive=1, negative=11, total=12

AnGam\_Sua5b\_cells.rep



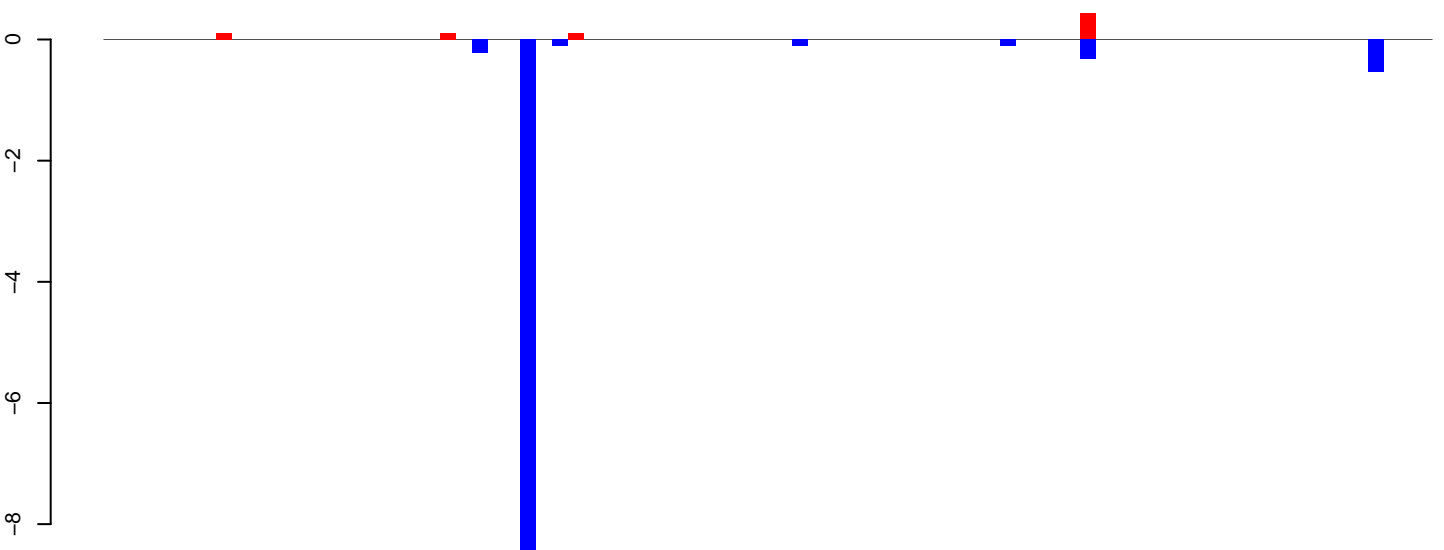
Window Size: 50 | Avg Length: 4687 | Virus@MH550148.1-Aedes\_albopictus\_densovirus\_isolate\_NC01\_L3\_1\_trimmed\_contig\_5889-RC:1-3937

AnGam\_Sua5b\_cells.18\_23.rep



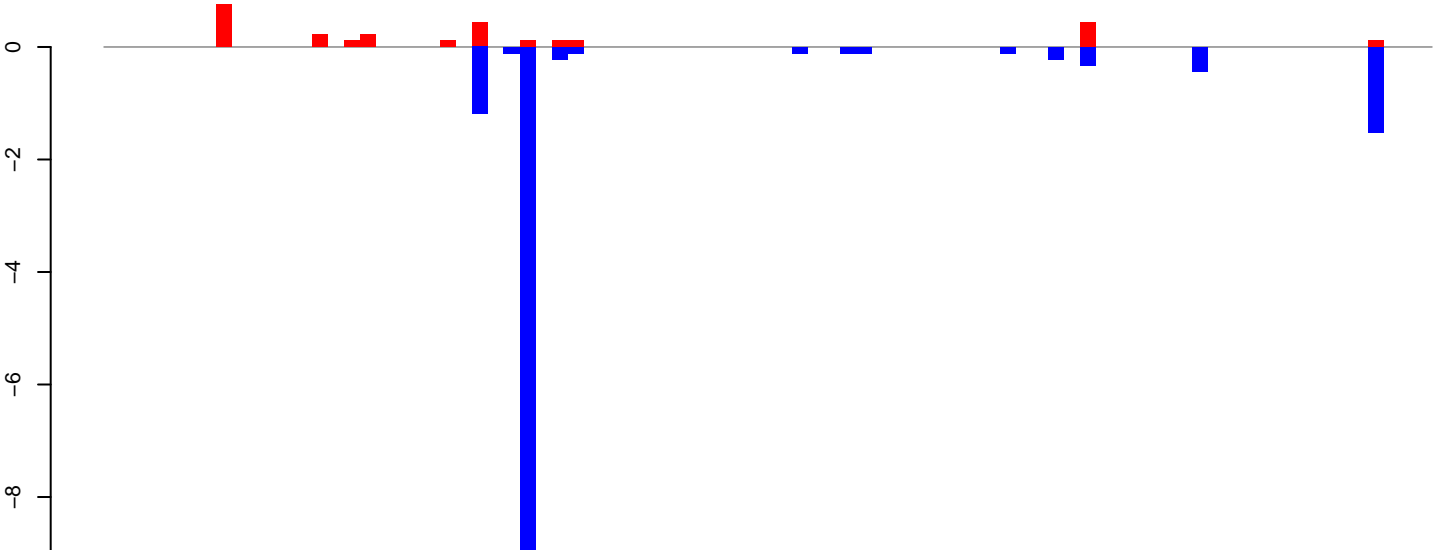
positive=2, negative=4, total=6

AnGam\_Sua5b\_cells.24\_35.rep



positive=1, negative=11, total=11

AnGam\_Sua5b\_cells.rep

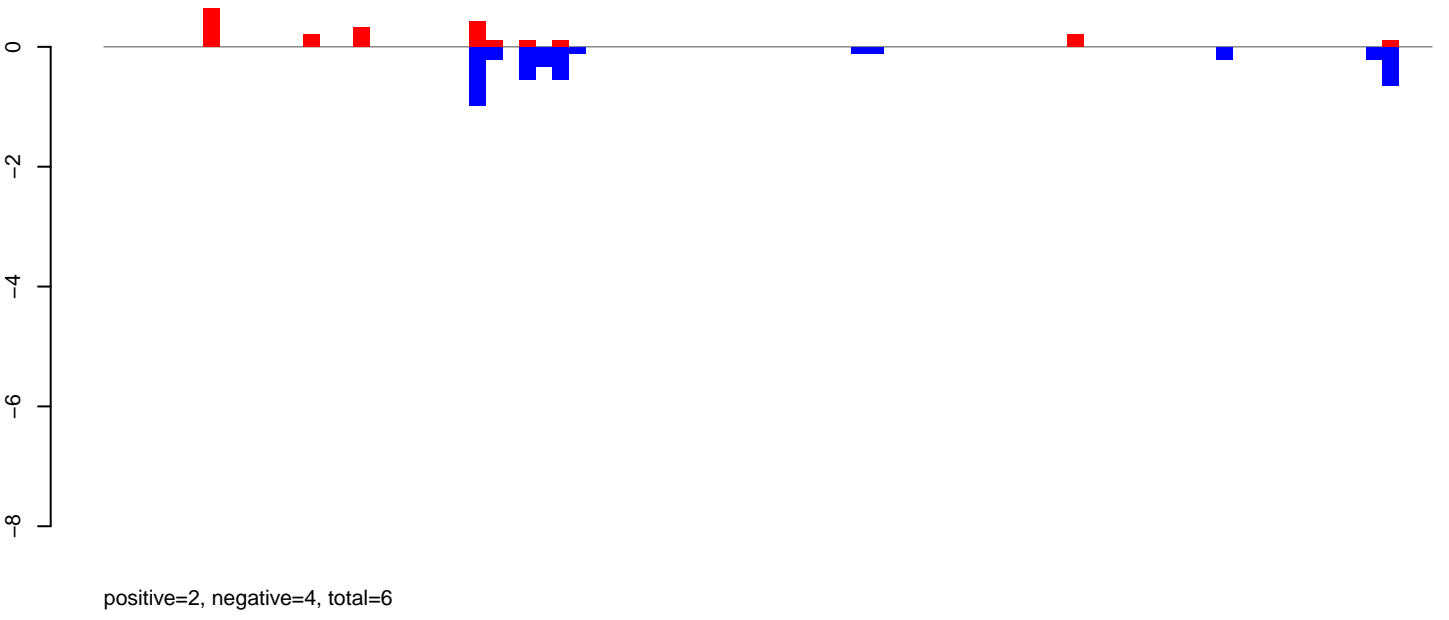


positive=3, negative=14, total=17

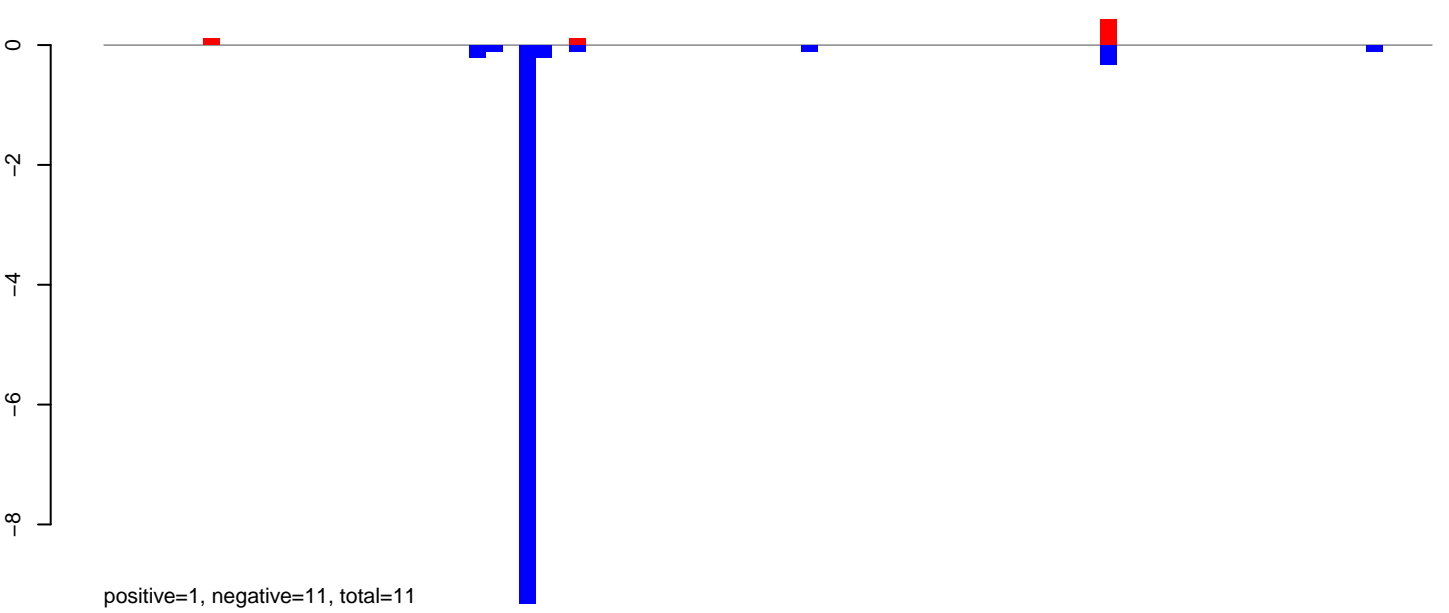
Window size=50, length=4176, Virus@X74945.1-Aedes\_albopictus\_densovirus\_2\_DNA:1-4176



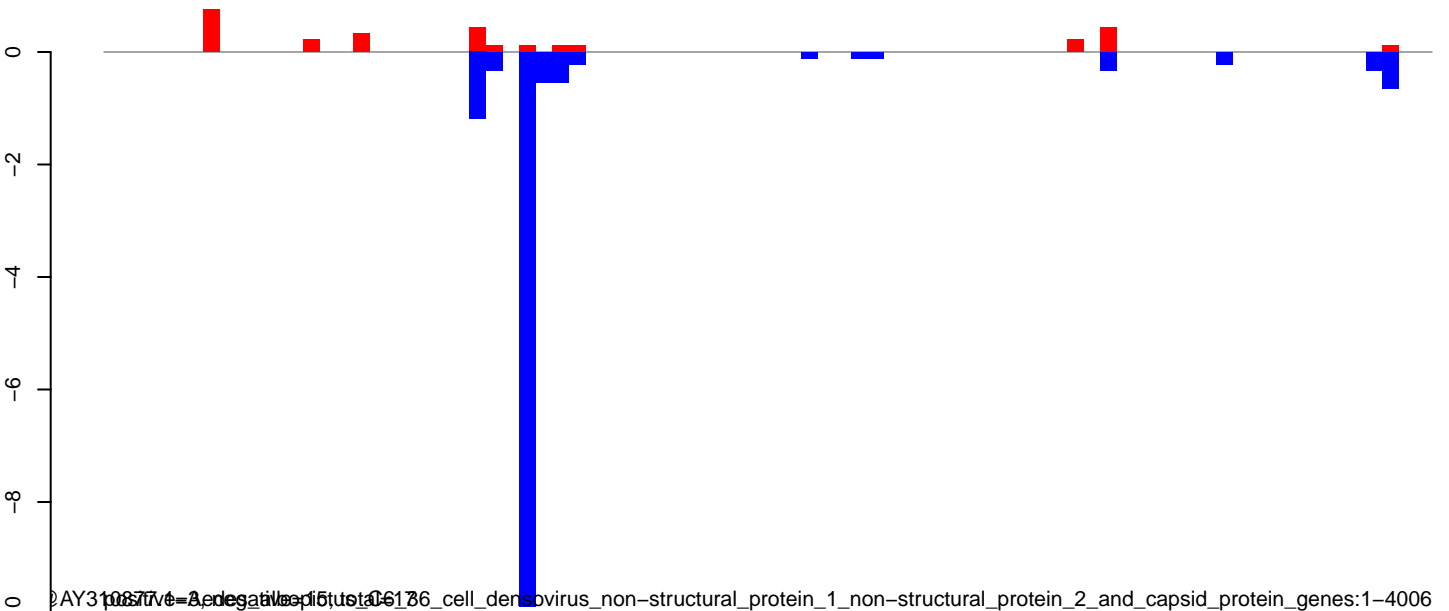
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep

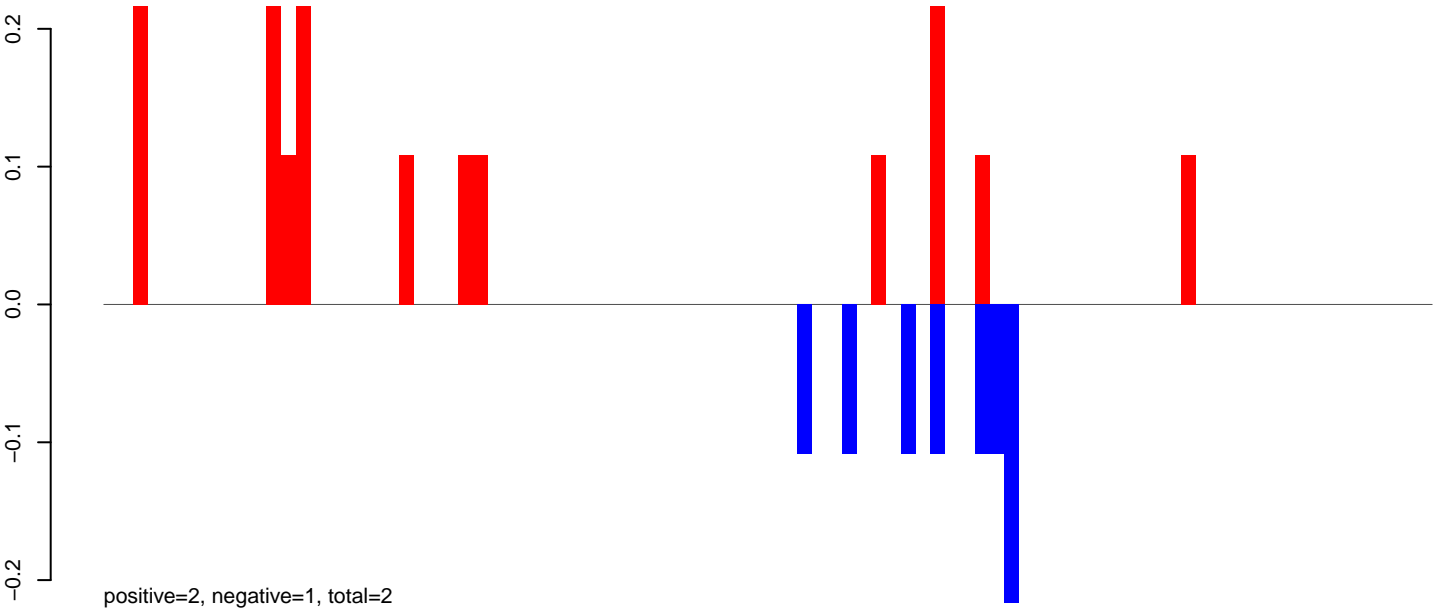


AnGam\_Sua5b\_cells.rep



AY310874 AY310875 AY310876 AY310877 AY310878 AY310879 AY310880 AY310881 AY310882 AY310883 AY310884 AY310885 AY310886 AY310887 AY310888 AY310889 AY310890 AY310891 AY310892 AY310893 AY310894 AY310895 AY310896 AY310897 AY310898 AY310899 AY310900 AY310901 AY310902 AY310903 AY310904 AY310905 AY310906 AY310907 AY310908 AY310909 AY310910 AY310911 AY310912 AY310913 AY310914 AY310915 AY310916 AY310917 AY310918 AY310919 AY310920 AY310921 AY310922 AY310923 AY310924 AY310925 AY310926 AY310927 AY310928 AY310929 AY310930 AY310931 AY310932 AY310933 AY310934 AY310935 AY310936 AY310937 AY310938 AY310939 AY310940 AY310941 AY310942 AY310943 AY310944 AY310945 AY310946 AY310947 AY310948 AY310949 AY310950 AY310951 AY310952 AY310953 AY310954 AY310955 AY310956 AY310957 AY310958 AY310959 AY310960 AY310961 AY310962 AY310963 AY310964 AY310965 AY310966 AY310967 AY310968 AY310969 AY310970 AY310971 AY310972 AY310973 AY310974 AY310975 AY310976 AY310977 AY310978 AY310979 AY310980 AY310981 AY310982 AY310983 AY310984 AY310985 AY310986 AY310987 AY310988 AY310989 AY310990 AY310991 AY310992 AY310993 AY310994 AY310995 AY310996 AY310997 AY310998 AY310999

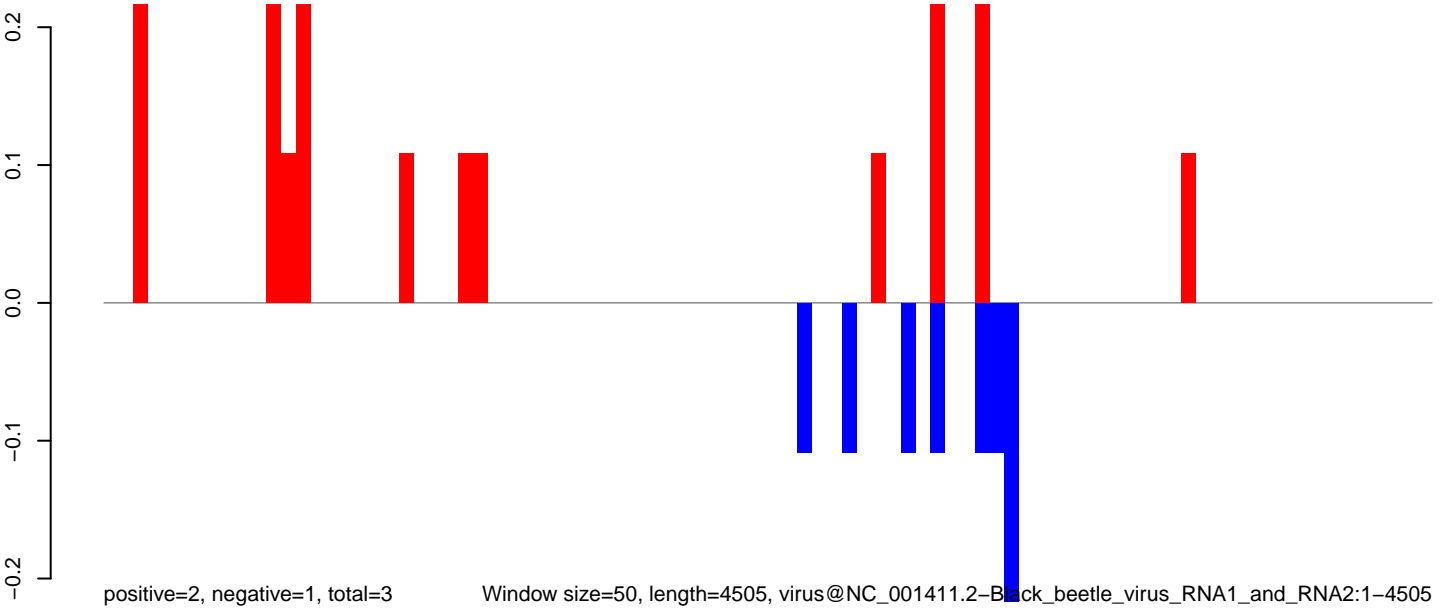
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep



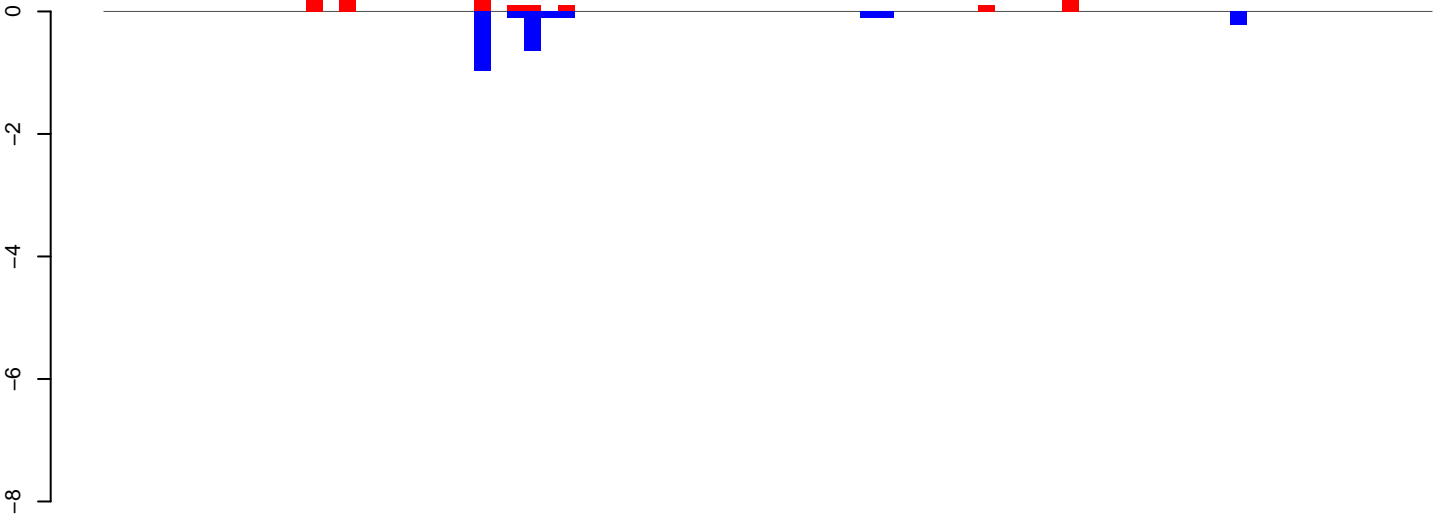
AnGam\_Sua5b\_cells.rep



Window size=50, length=4505, virus@NC\_001411.2-Black\_beetle\_virus\_RNA1\_and\_RNA2:1-4505

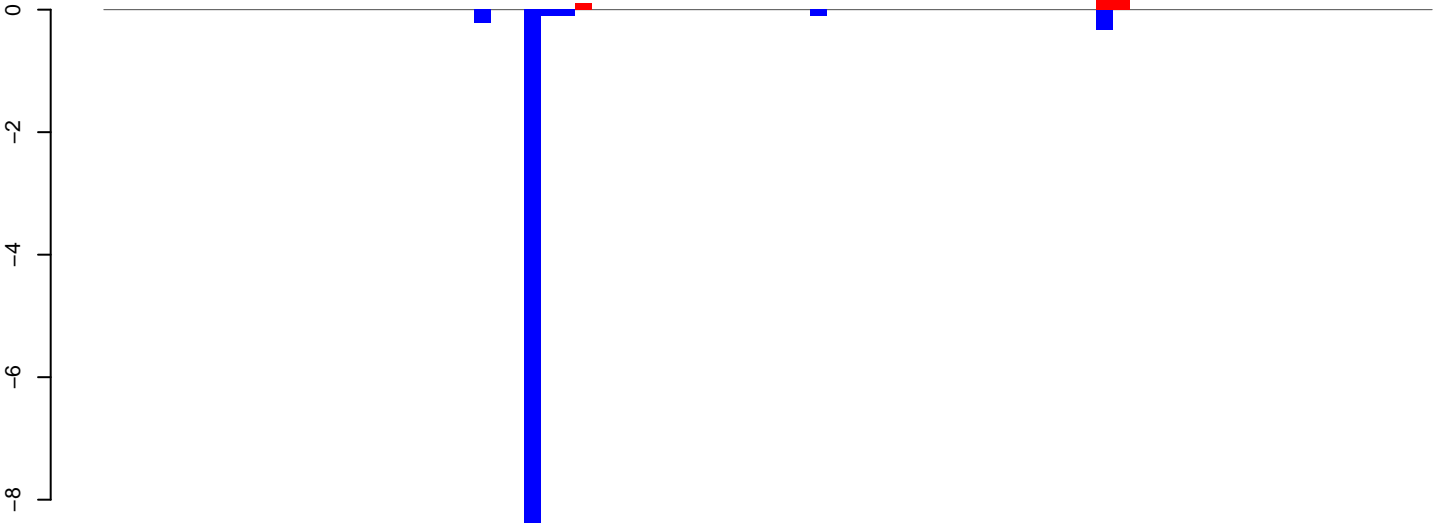
0 1000 2000 3000 4000

AnGam\_Sua5b\_cells.18\_23.rep



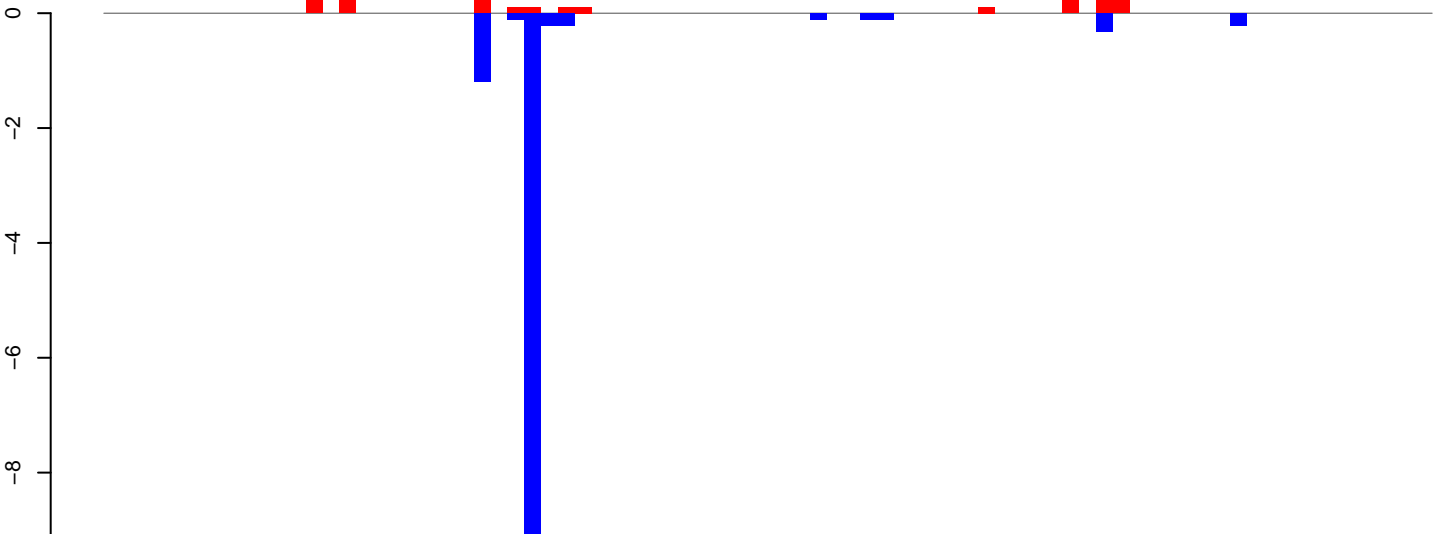
positive=2, negative=2, total=4

AnGam\_Sua5b\_cells.24\_35.rep



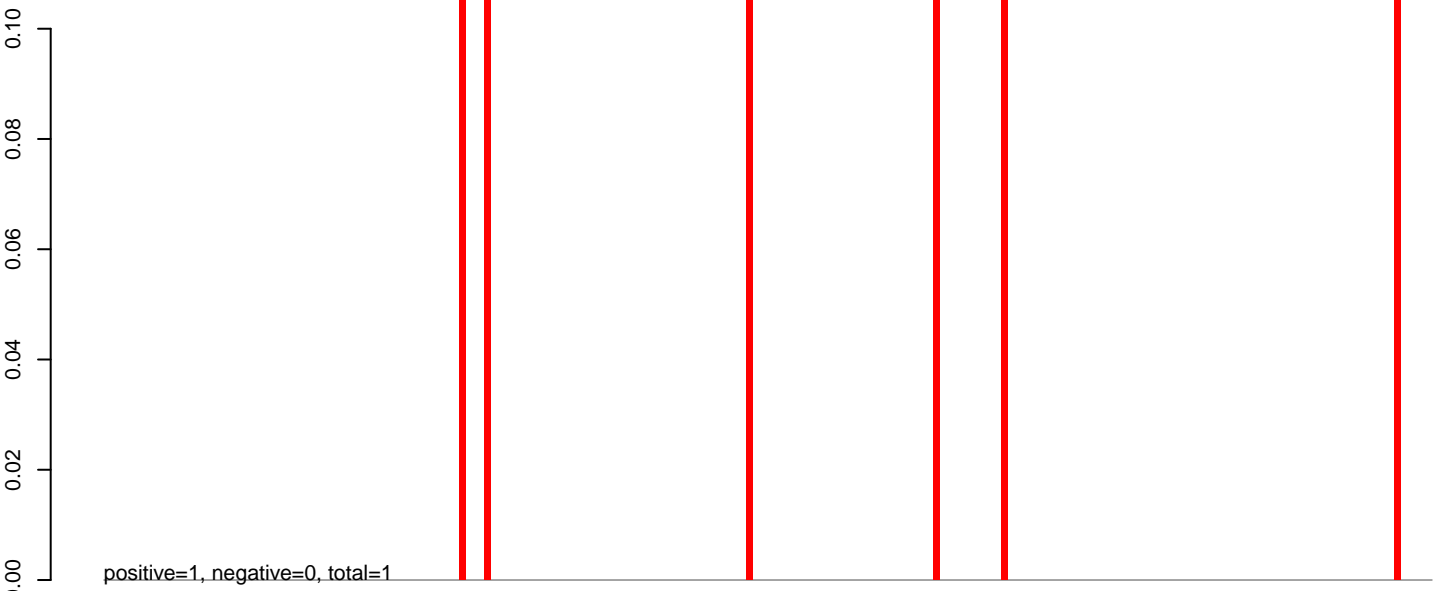
positive=1, negative=10, total=11

AnGam\_Sua5b\_cells.rep

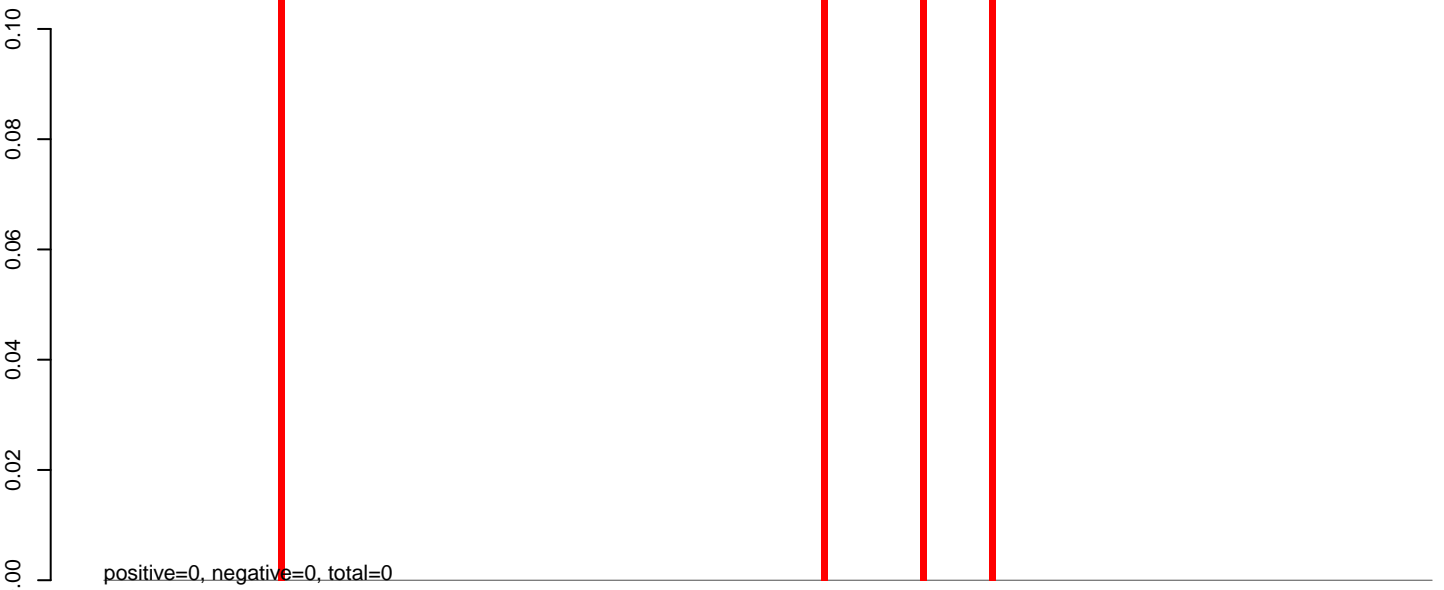


positive=4, negative=16, total=20

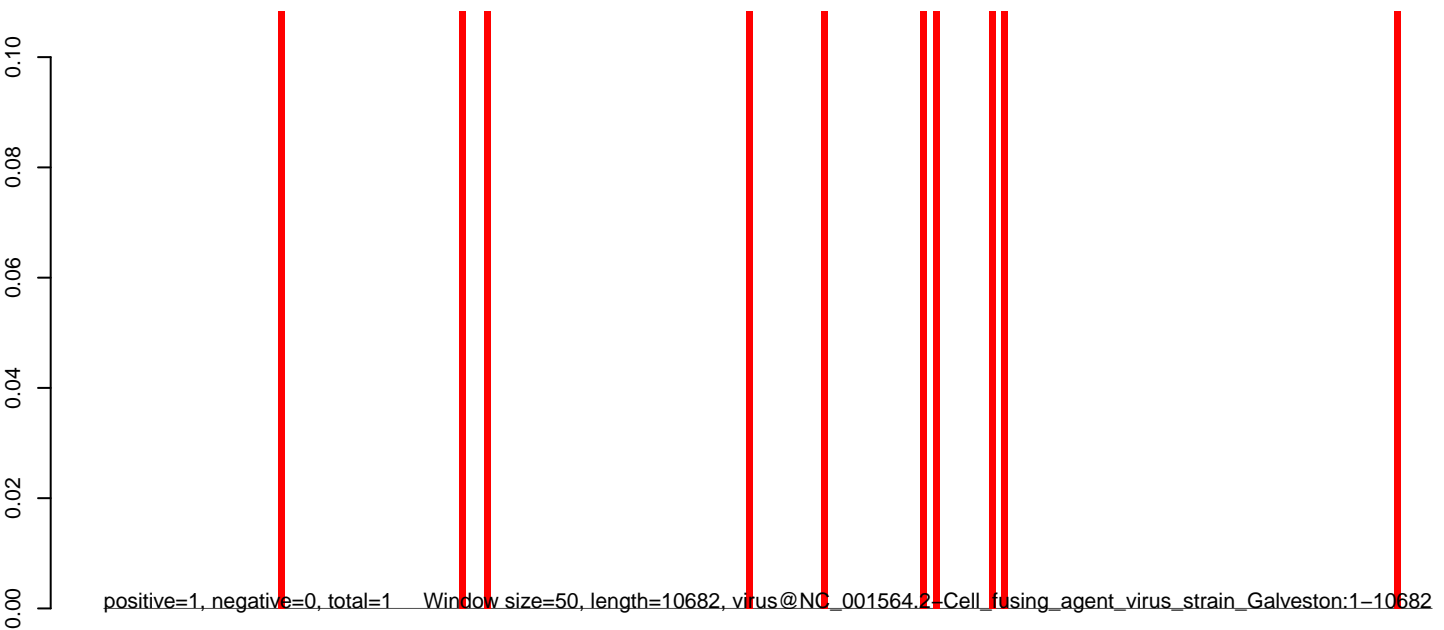
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep



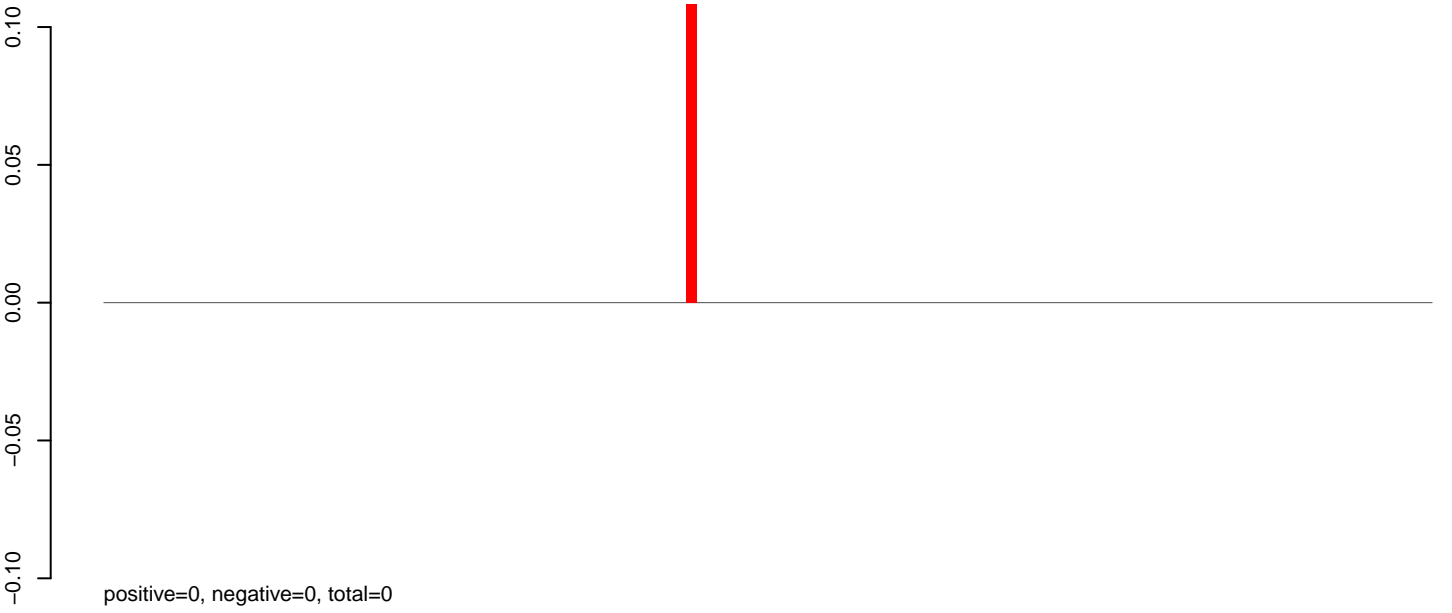
AnGam\_Sua5b\_cells.rep



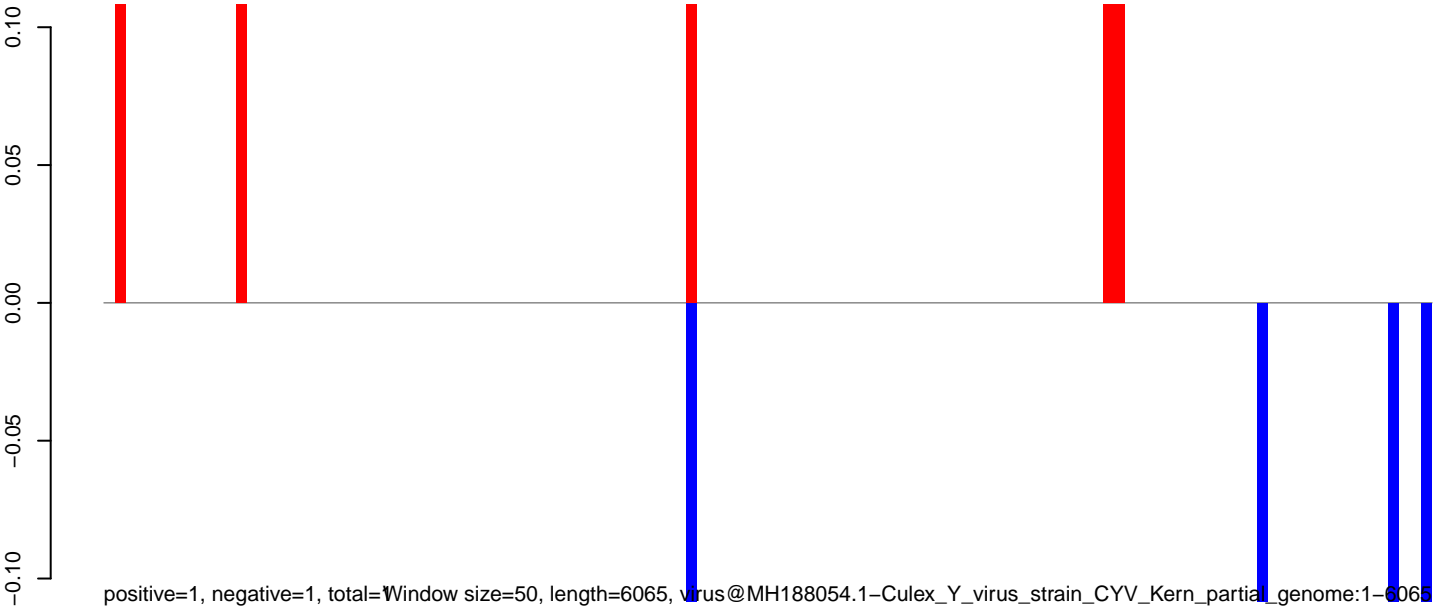
AnGam\_Sua5b\_cells.18\_23.rep



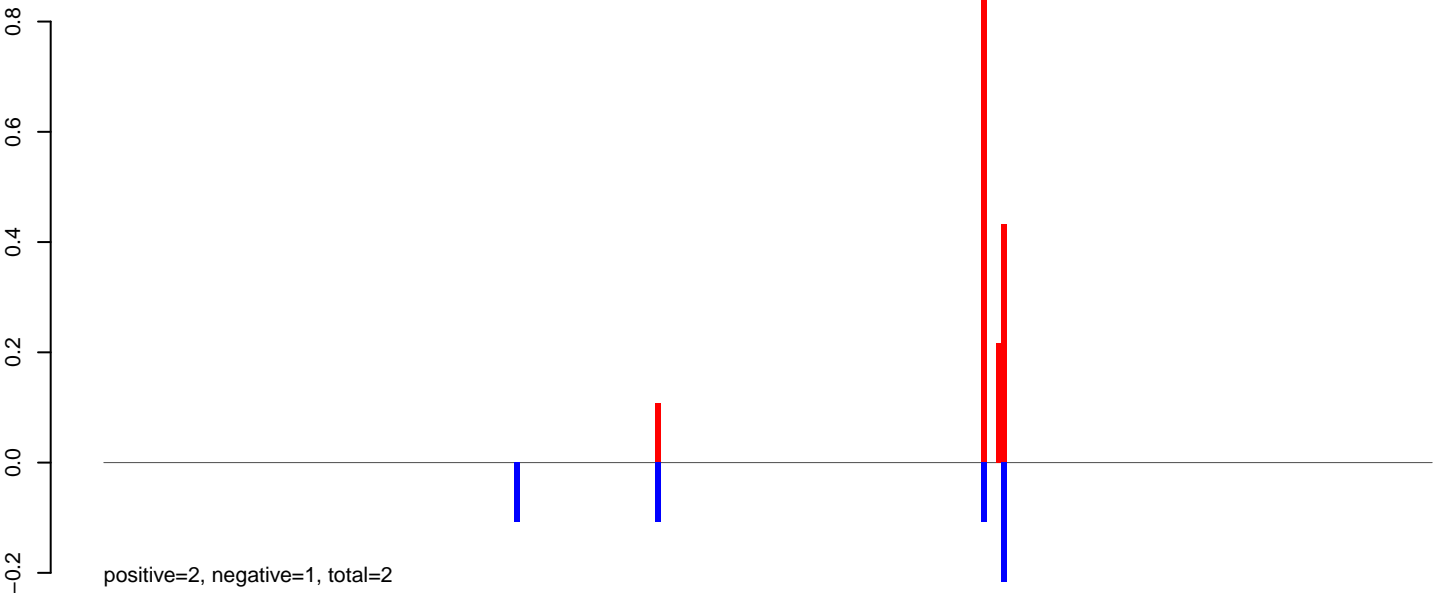
AnGam\_Sua5b\_cells.24\_35.rep



AnGam\_Sua5b\_cells.rep



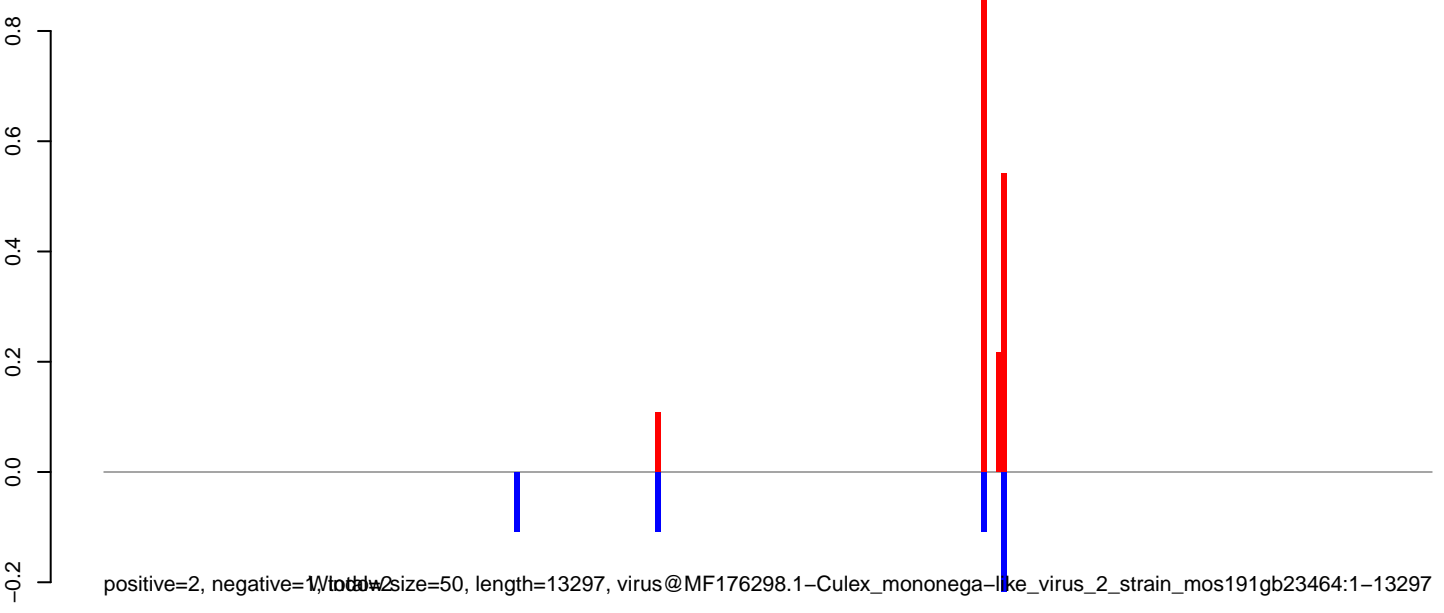
AnGam\_Sua5b\_cells.18\_23.rep



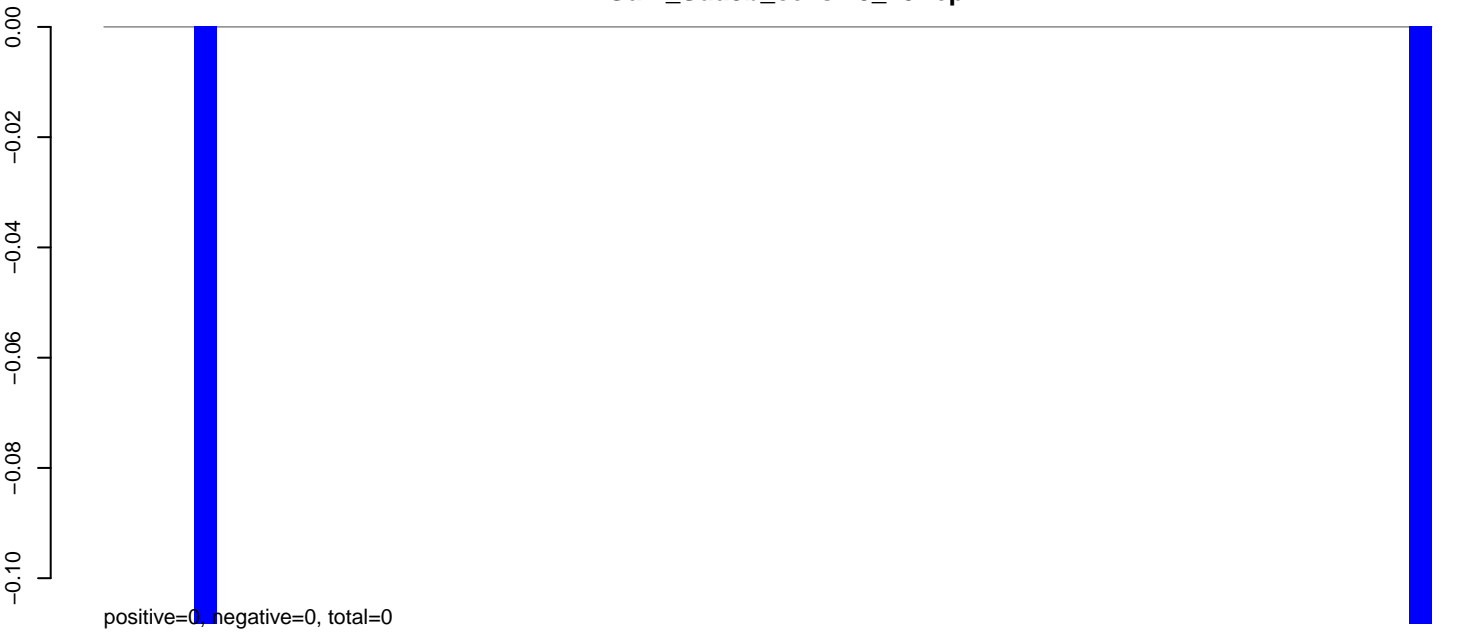
AnGam\_Sua5b\_cells.24\_35.rep



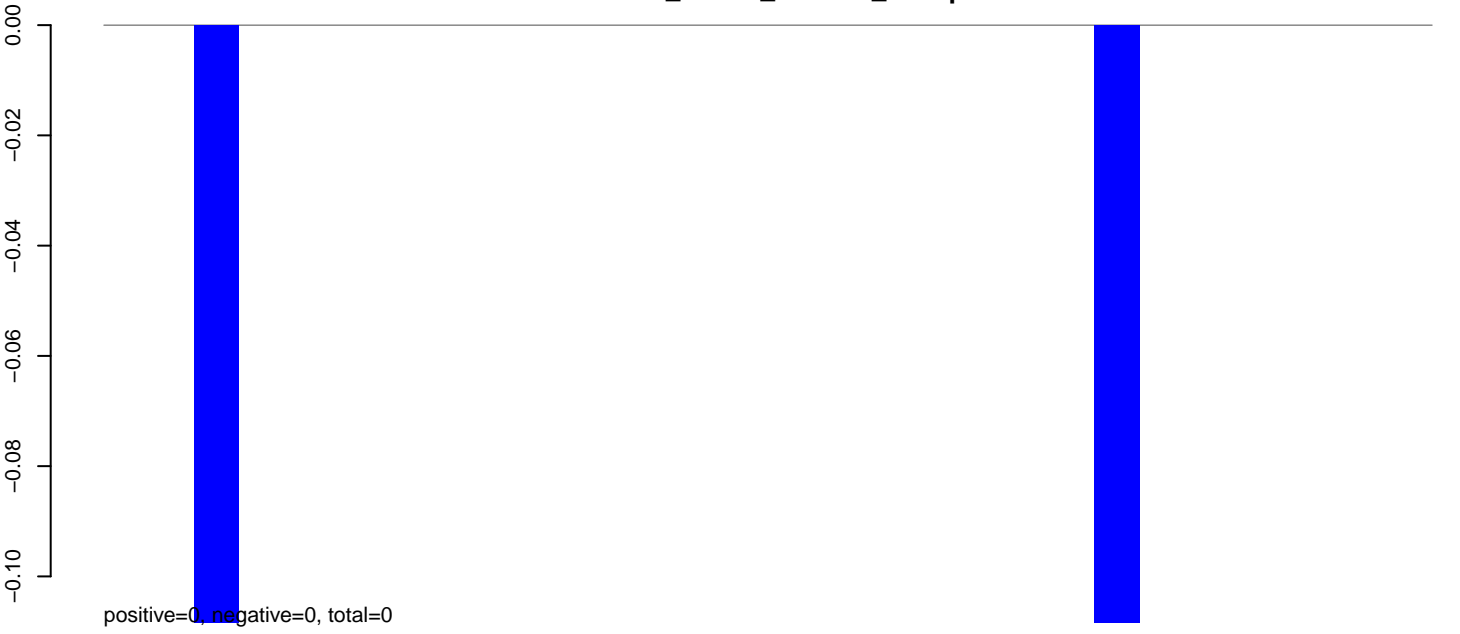
AnGam\_Sua5b\_cells.rep



AnGam\_Sua5b\_cells.18\_23.rep



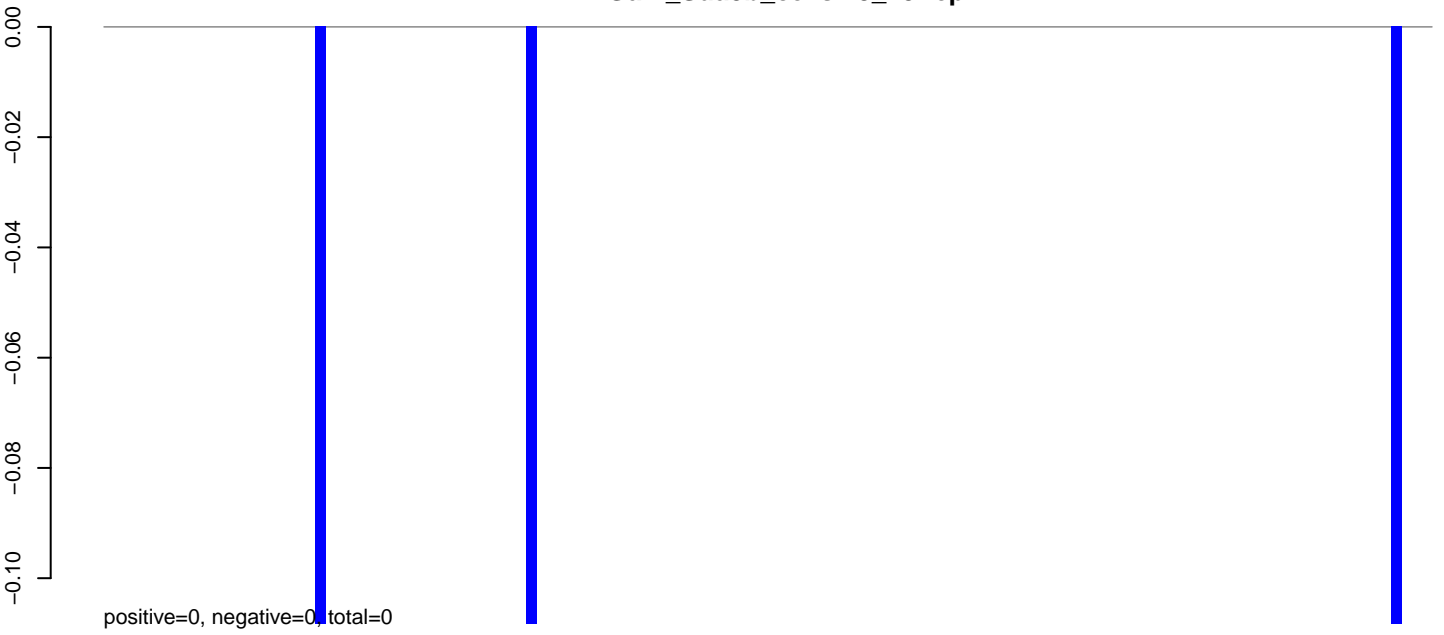
AnGam\_Sua5b\_cells.24\_35.rep



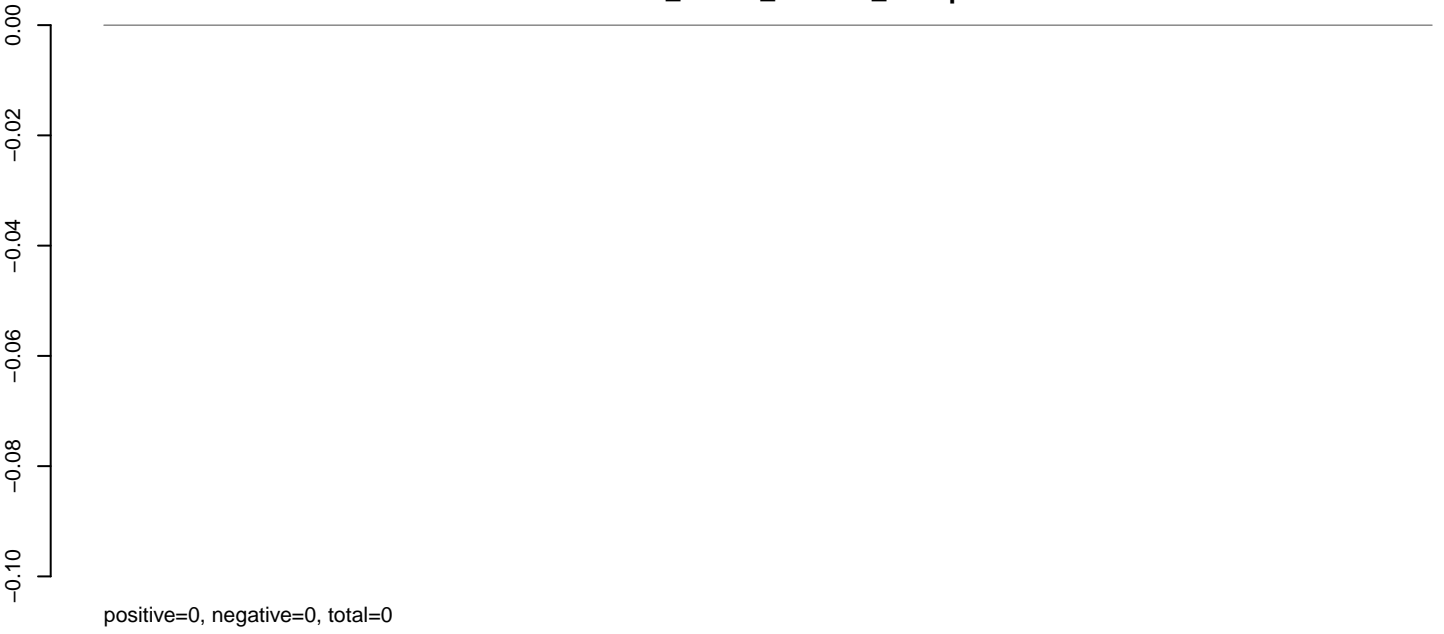
AnGam\_Sua5b\_cells.rep



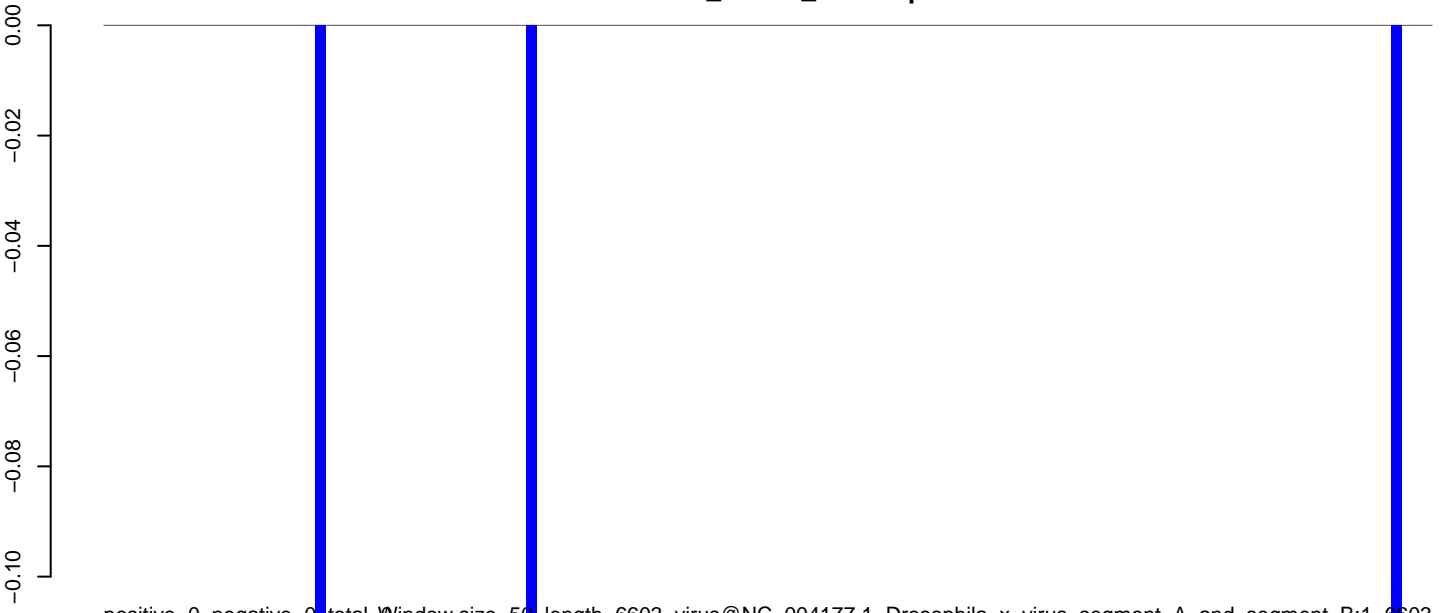
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep



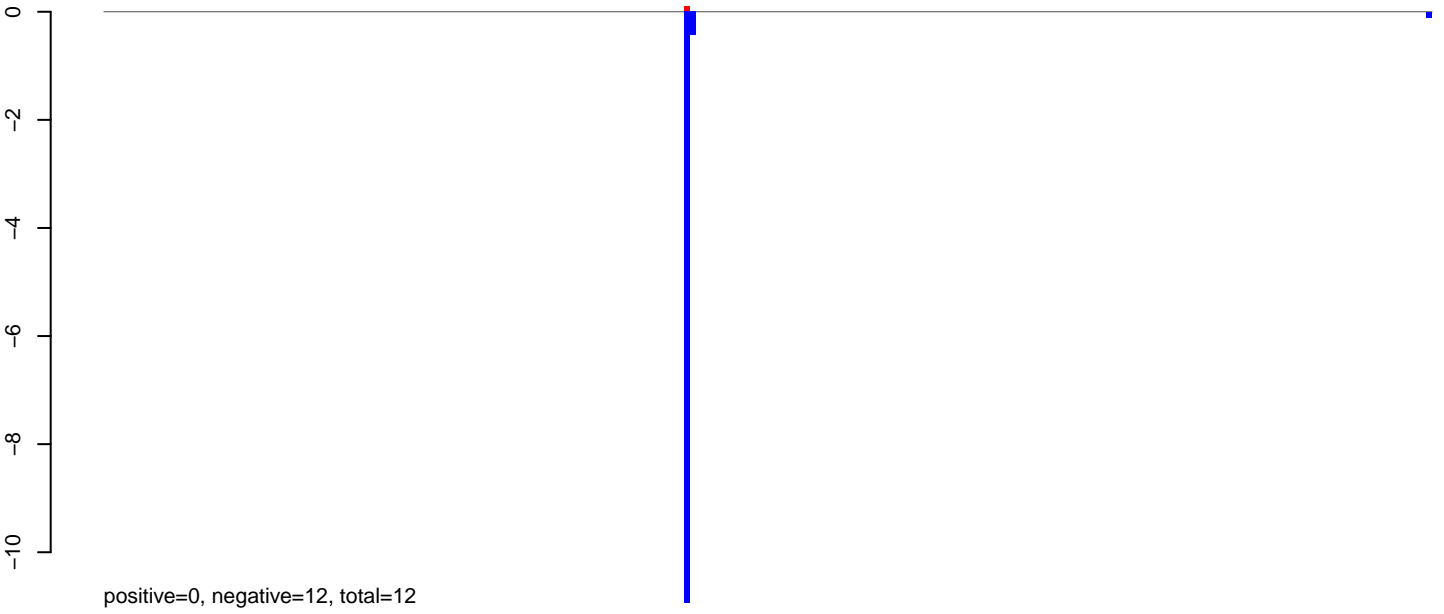
AnGam\_Sua5b\_cells.rep



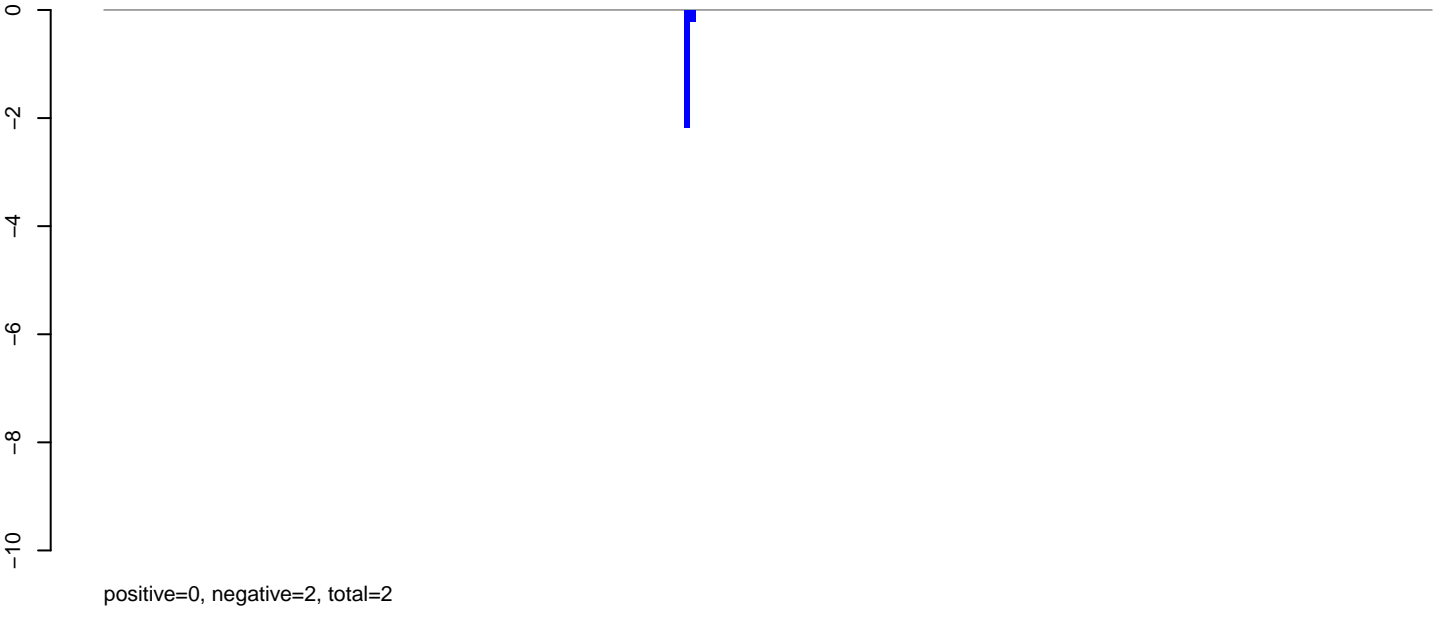
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window size=50, length=6603, virus@NC\_004177.1-Drosophila\_x\_virus\_segment\_A\_and\_segment\_B:1-6603



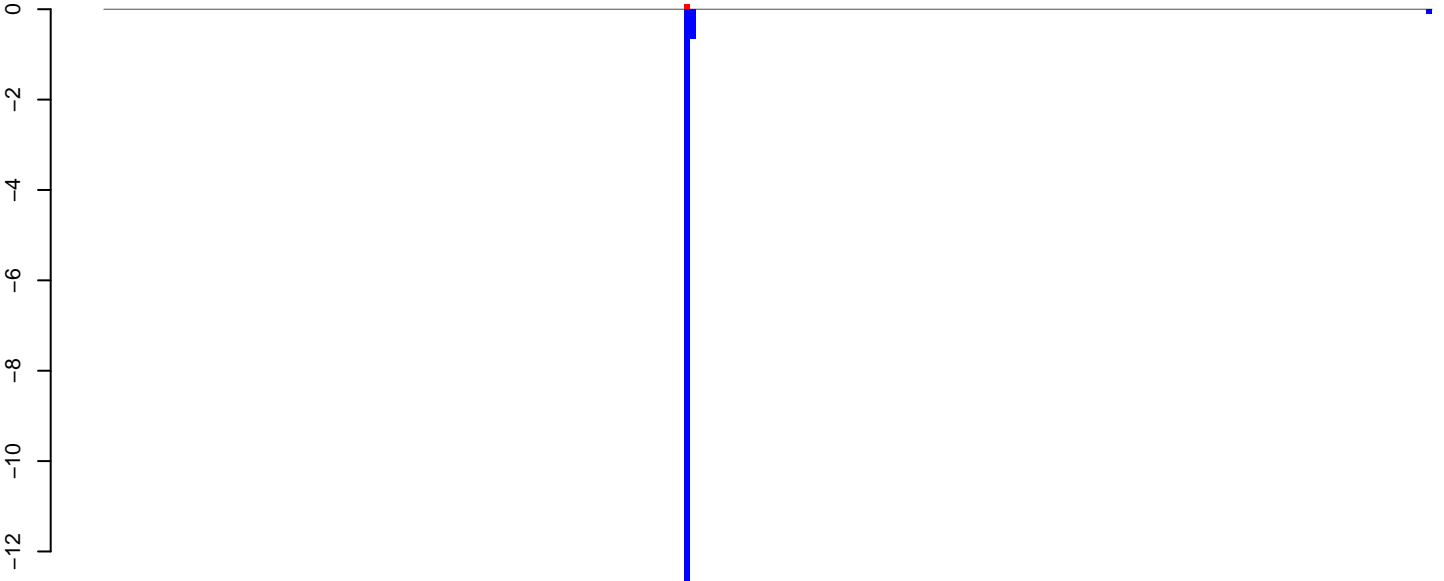
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep



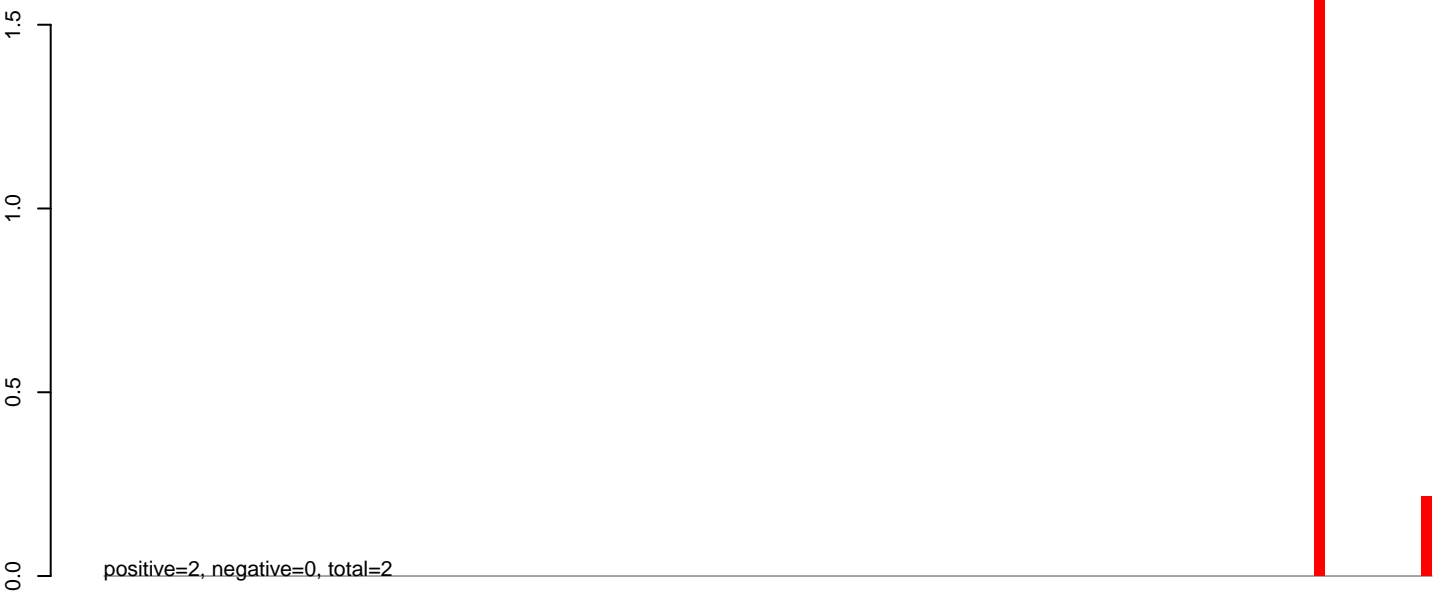
AnGam\_Sua5b\_cells.rep



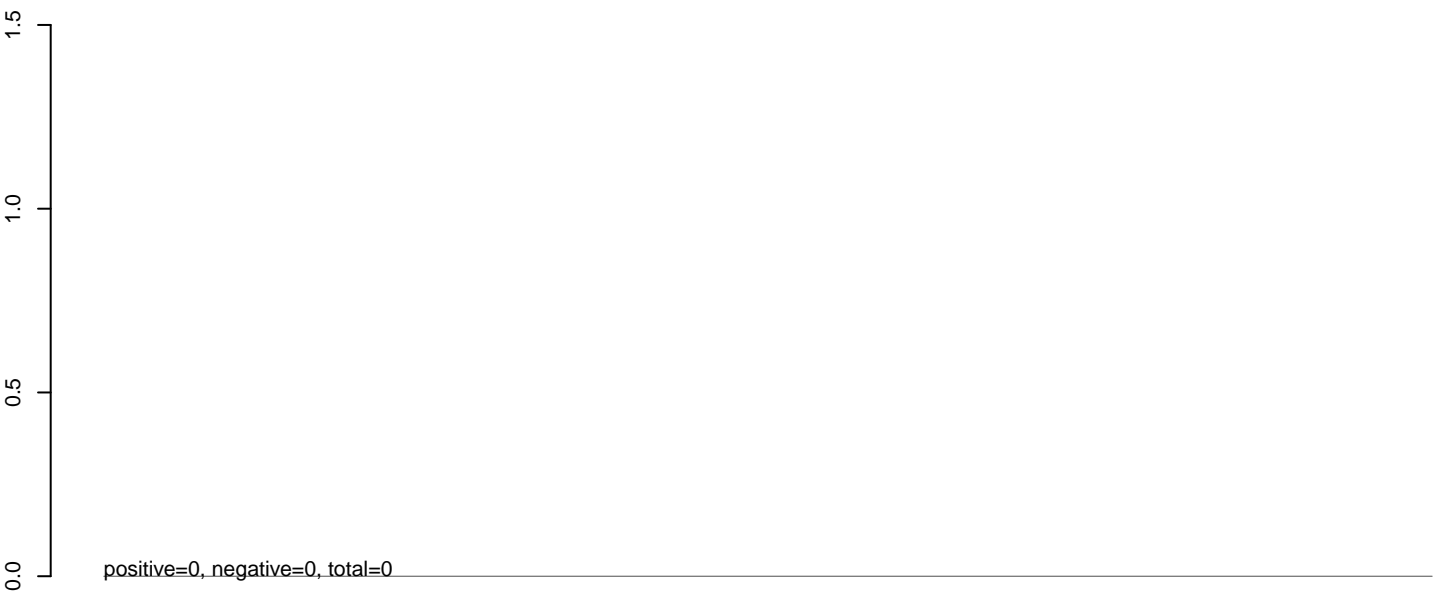
Window size=50, length=2369, virus@KJ716848.1-Ngali\_virus\_strain\_Adrar\_segment\_S\_plus\_segment\_M\_plus\_segment\_L:1-12369

0 2000 4000 6000 8000 10000 12000

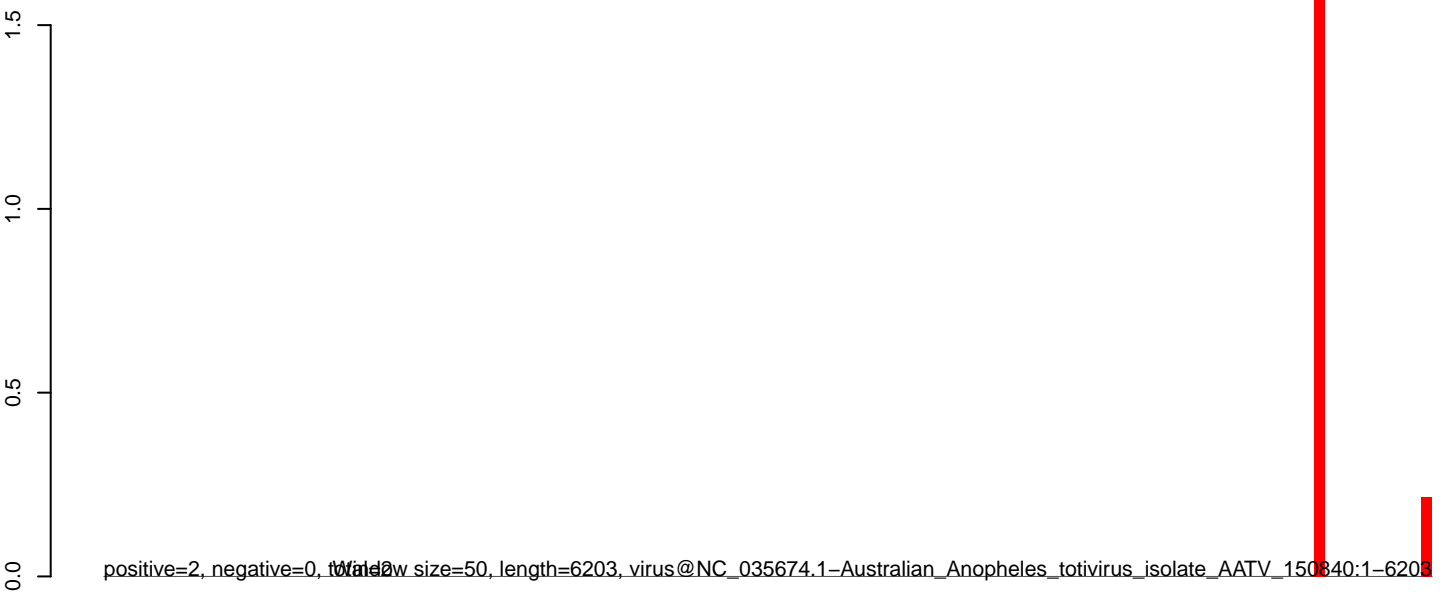
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep

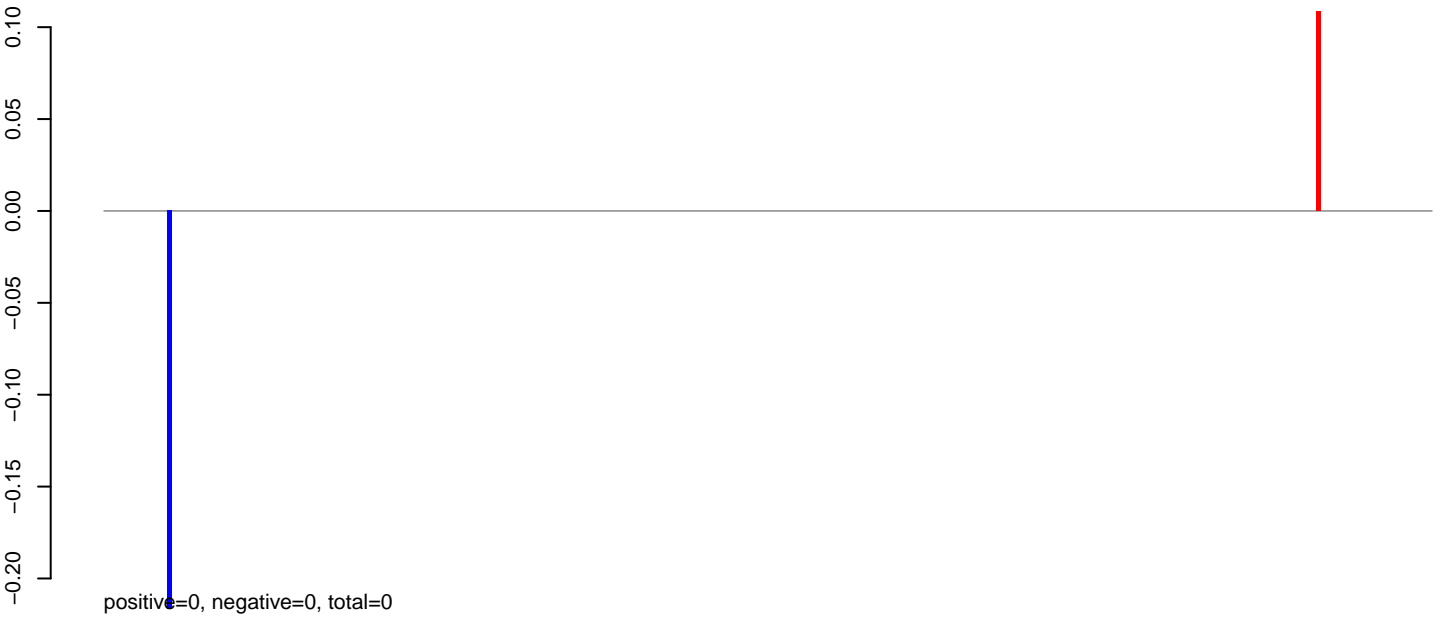


AnGam\_Sua5b\_cells.rep

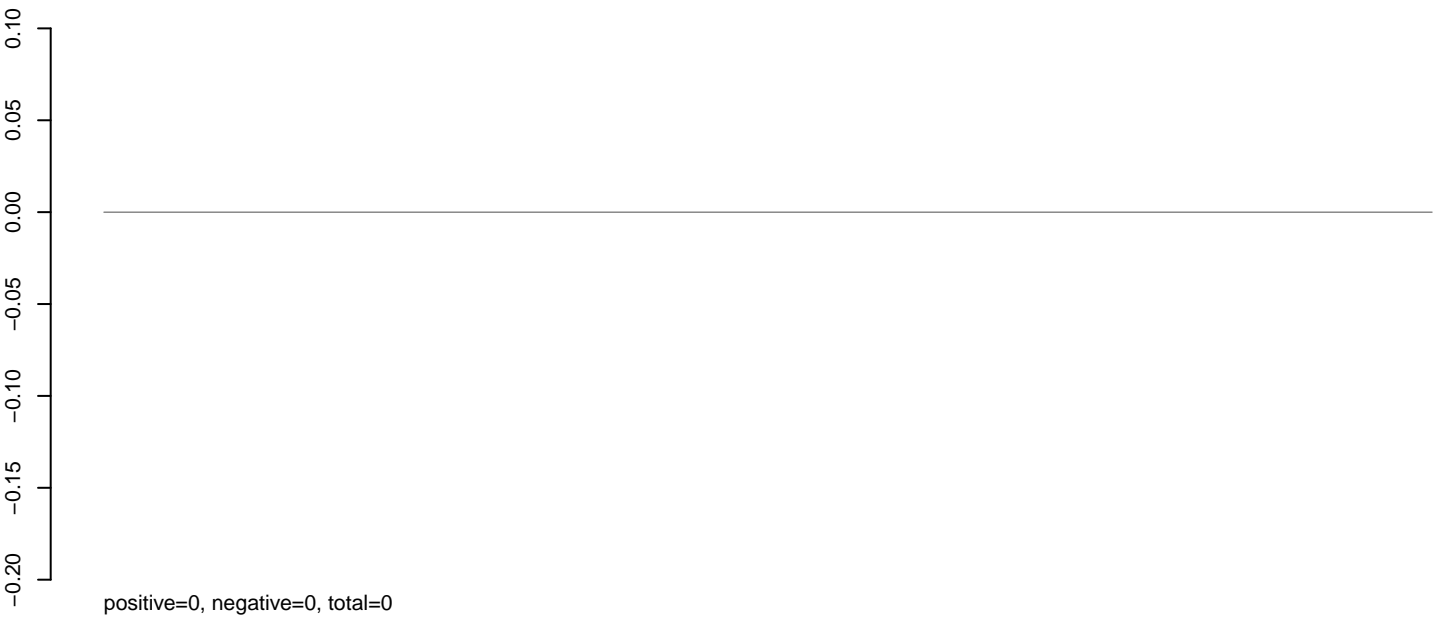


0 1000 2000 3000 4000 5000 6000

AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep



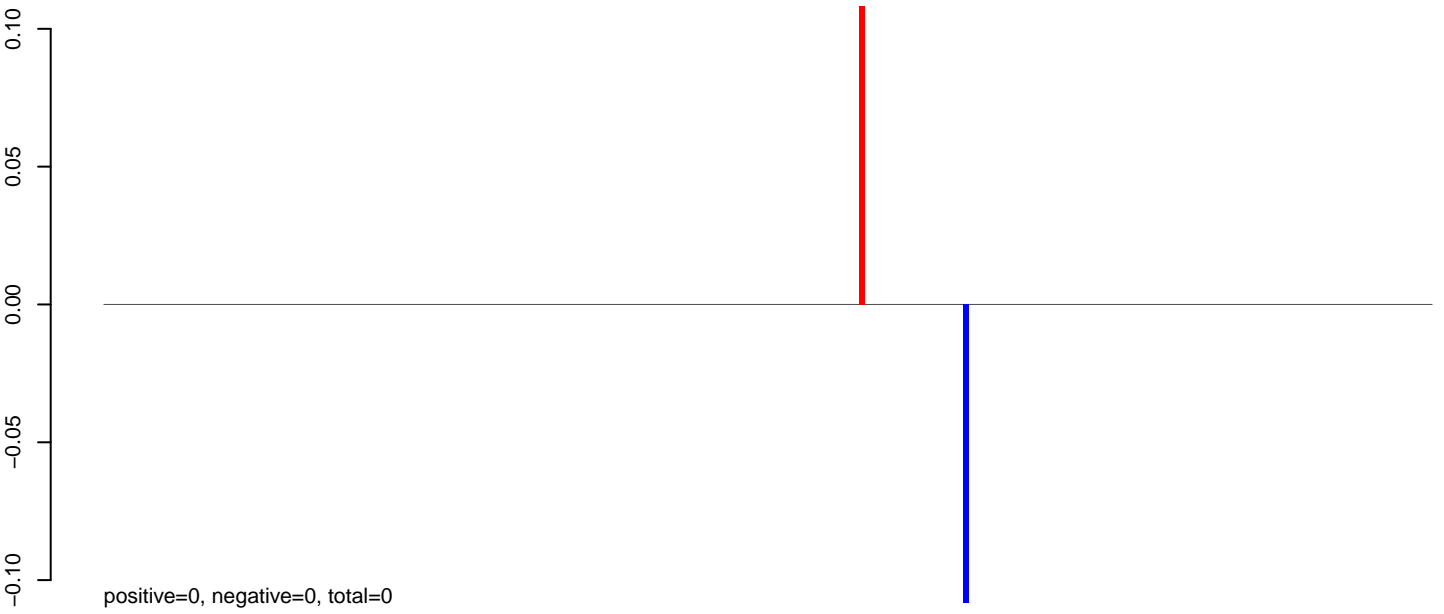
AnGam\_Sua5b\_cells.rep



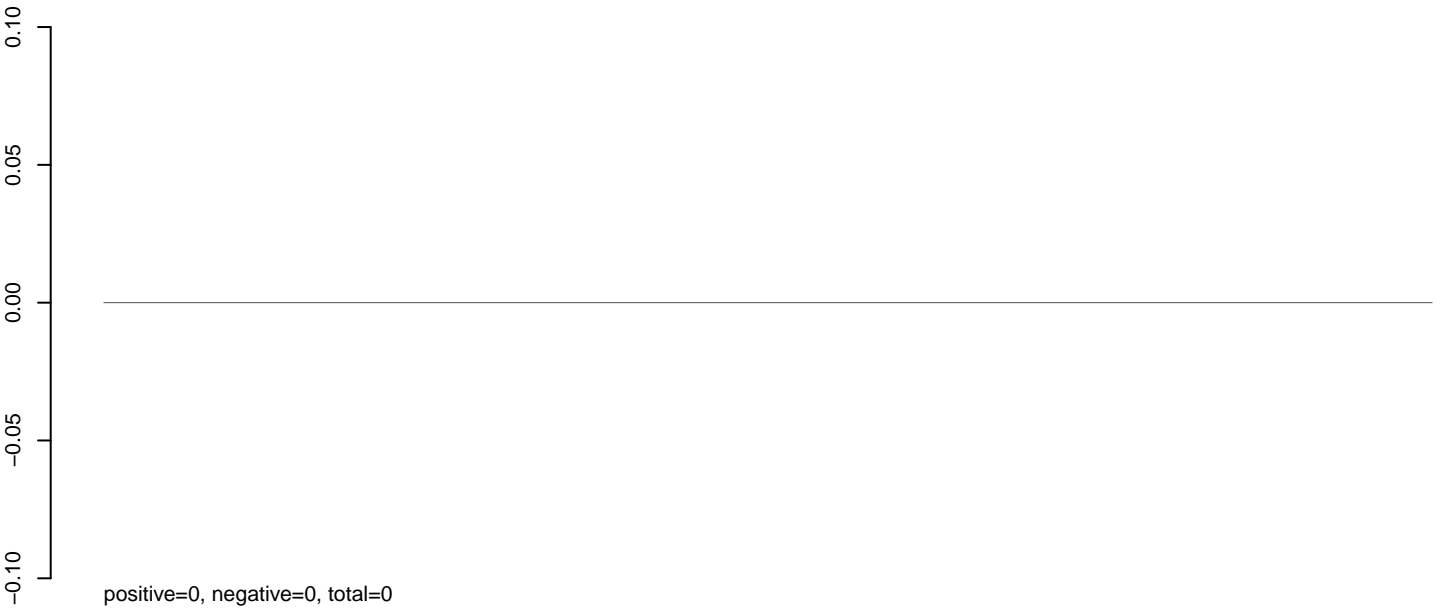
Manitoba positive, negative, total phosphoprotein\_hypothetical\_proteins\_matrix\_glycoprotein\_hypothetical\_protein\_and\_polymerase\_genes:1-13784

0 2000 4000 6000 8000 10000 12000 14000

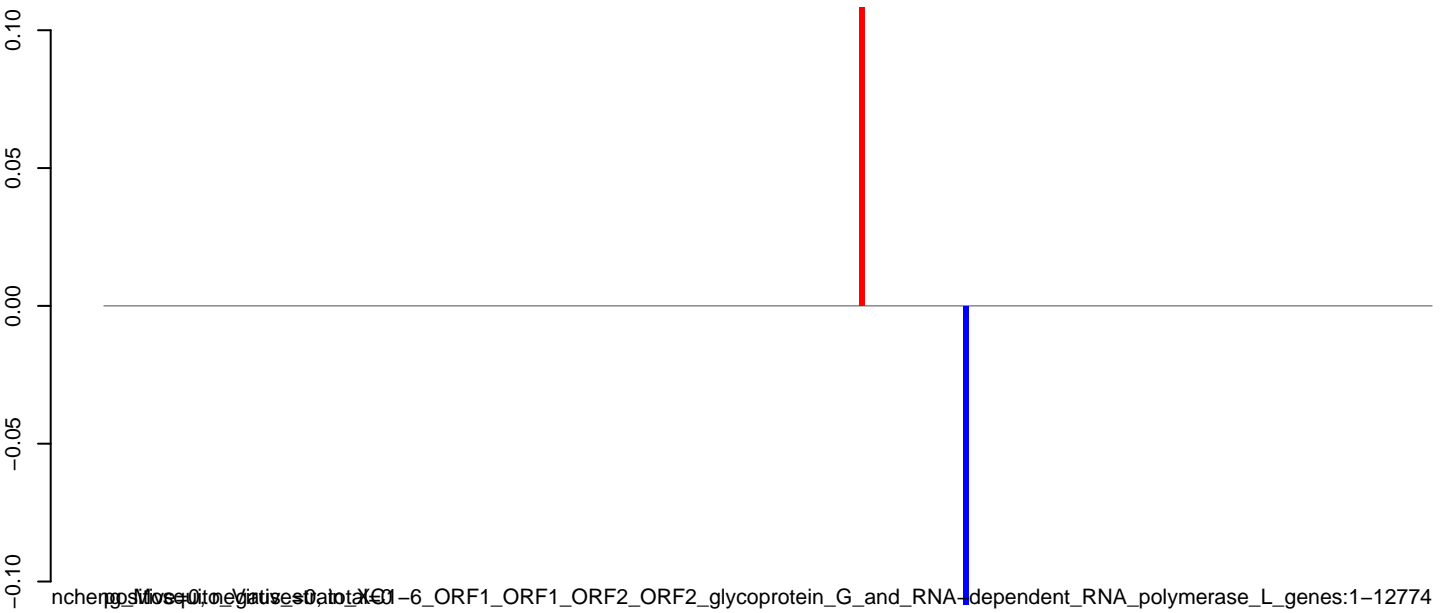
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep



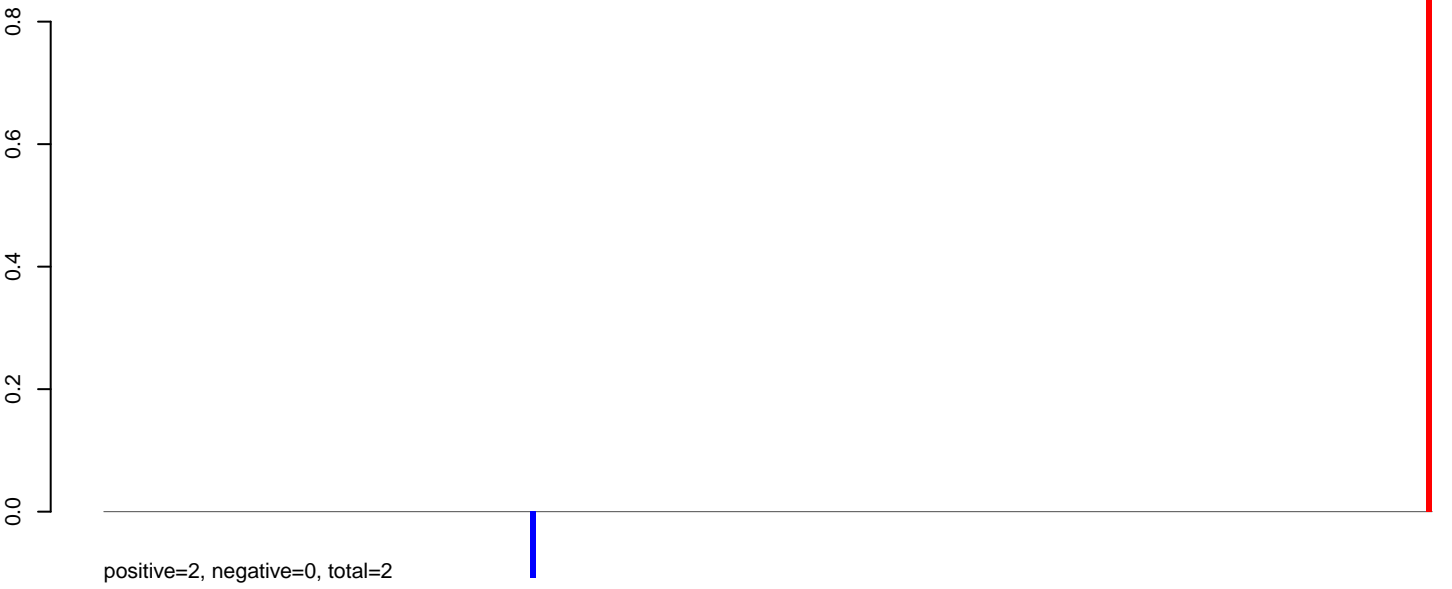
AnGam\_Sua5b\_cells.rep



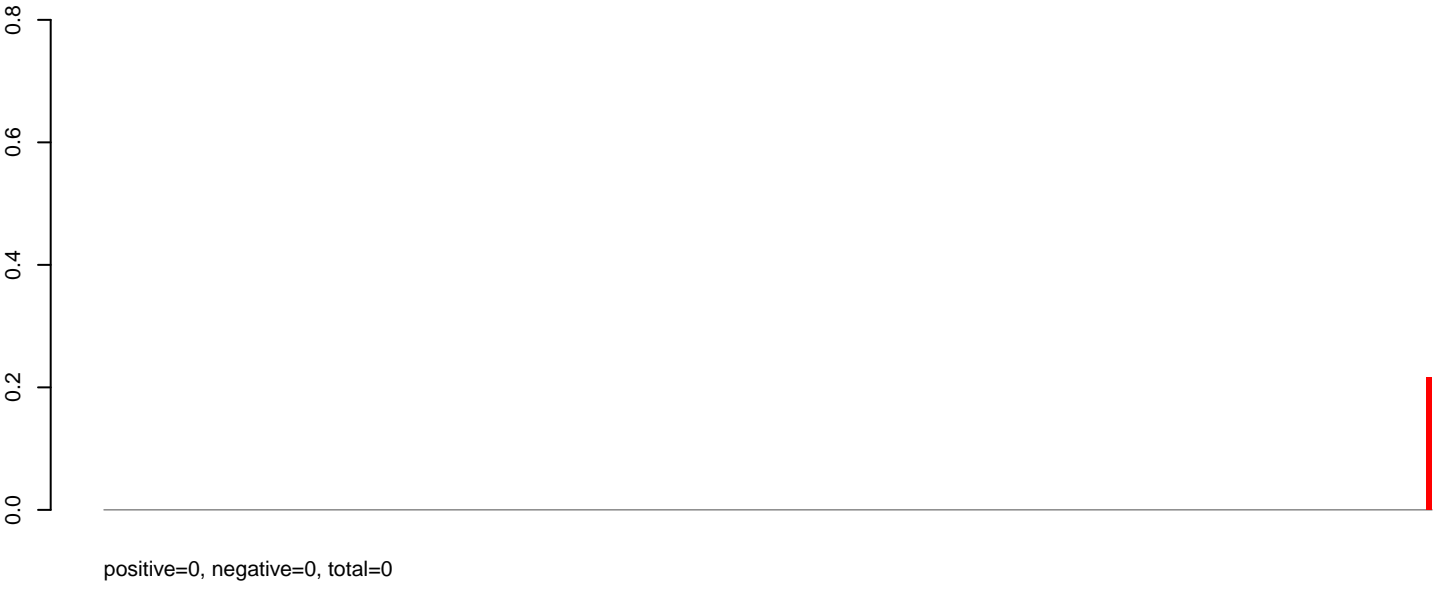
ncchemp110seq10megases0pan2401-6\_ORF1\_ORF1\_ORF2\_ORF2\_glycoprotein\_G\_and\_RNA-dependent\_RNA\_polymerase\_L\_genes:1-12774

0 2000 4000 6000 8000 10000 12000

AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep

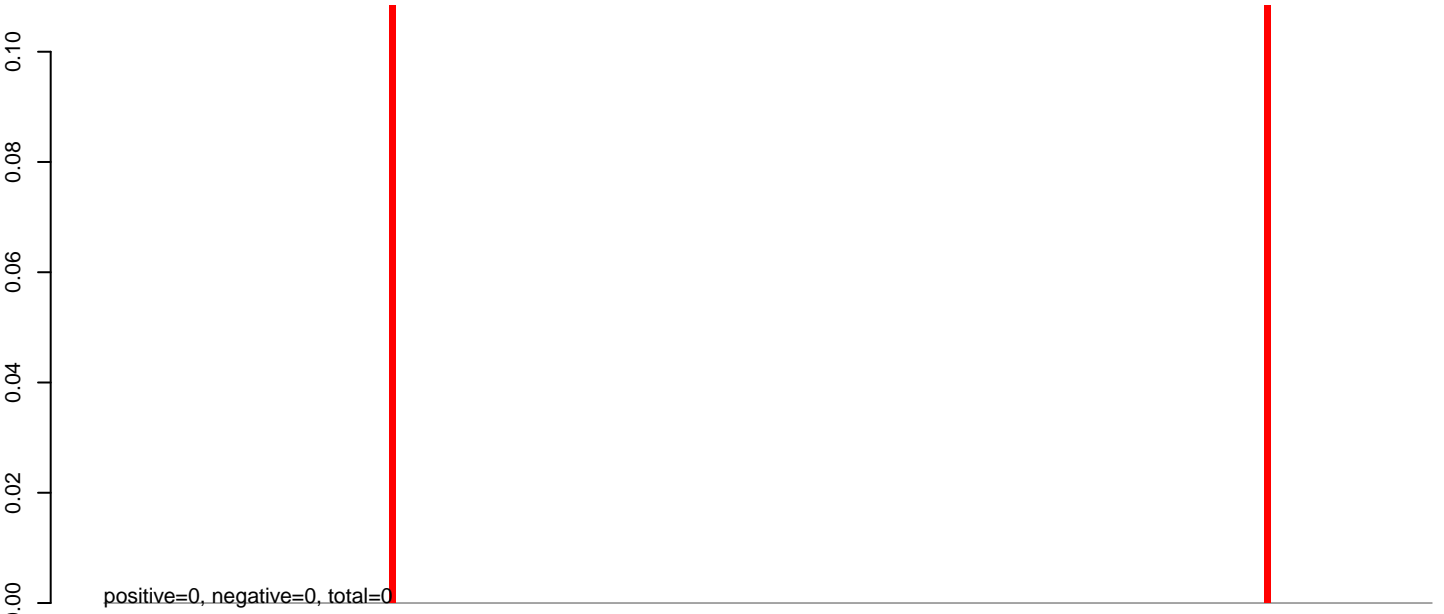


AnGam\_Sua5b\_cells.rep

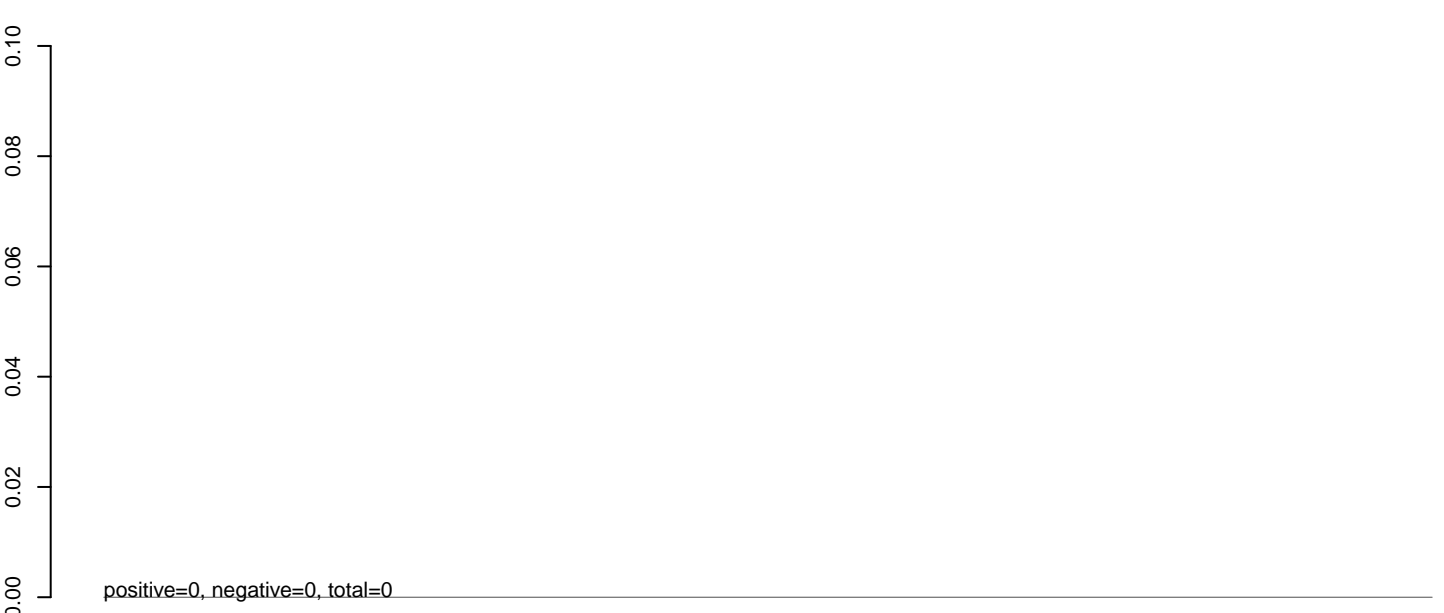


0 2000 4000 6000 8000 10000 12000

AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep

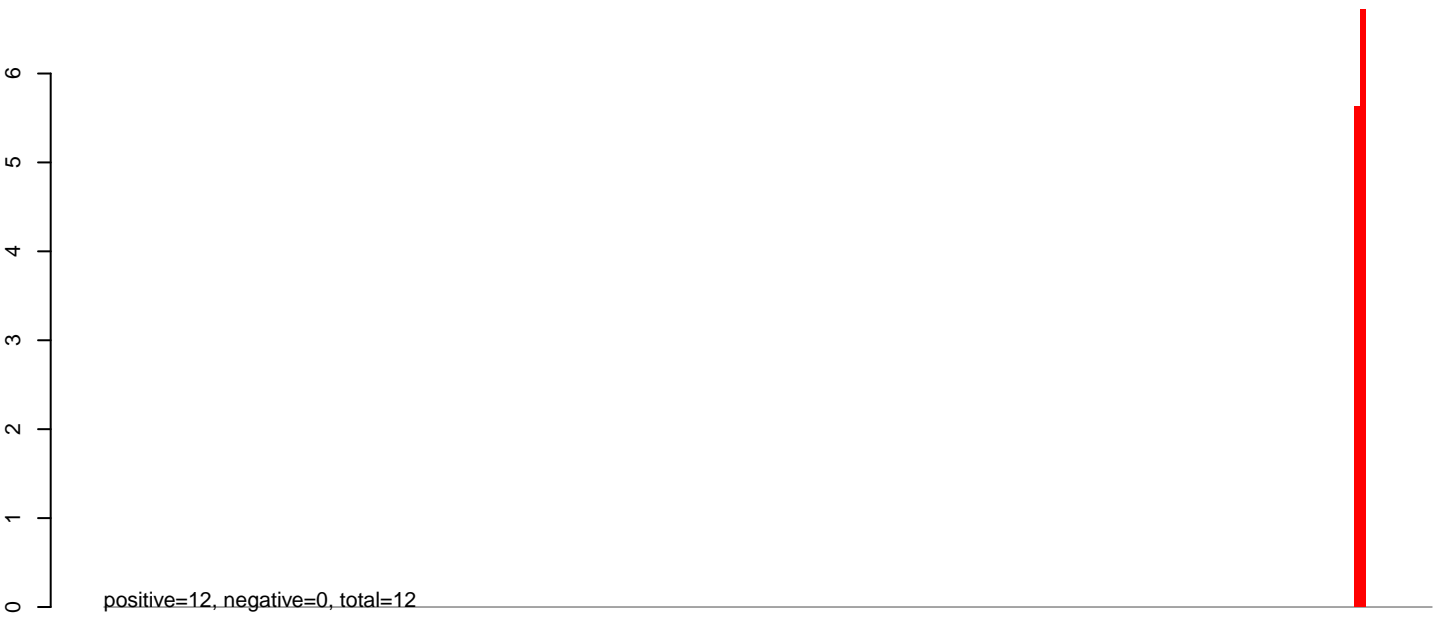


AnGam\_Sua5b\_cells.rep

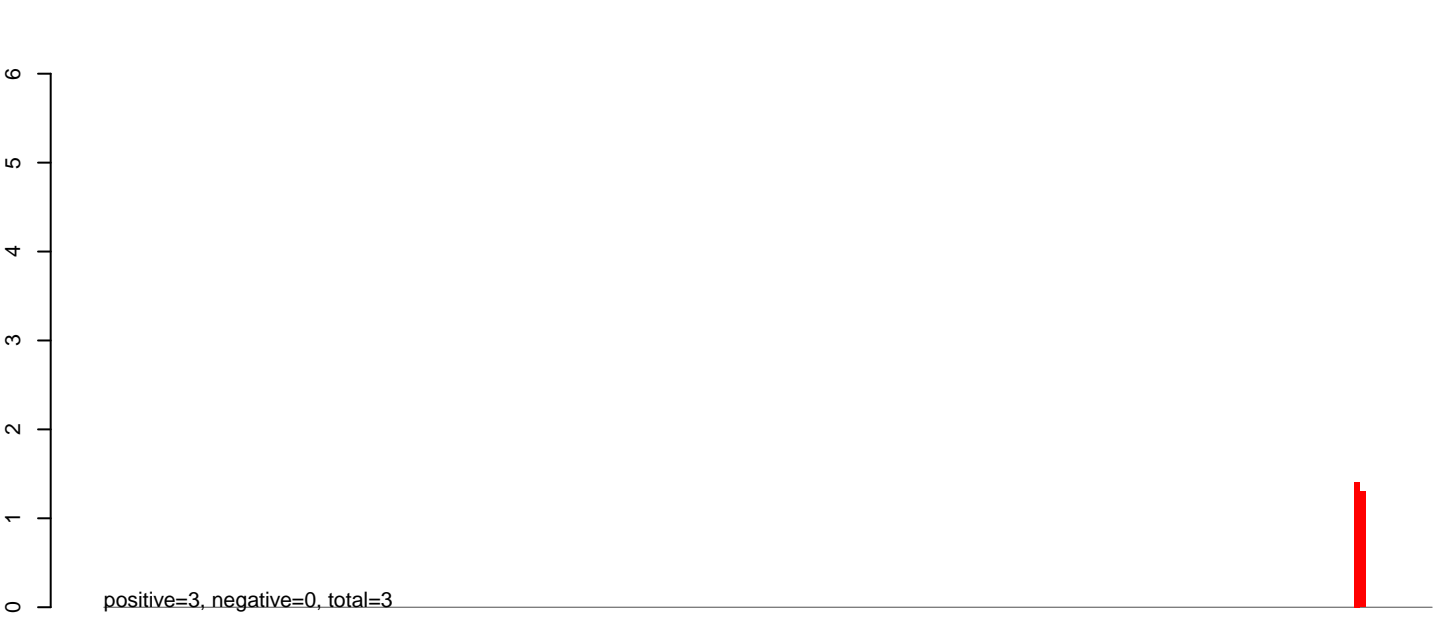


0 2000 4000 6000 8000 10000

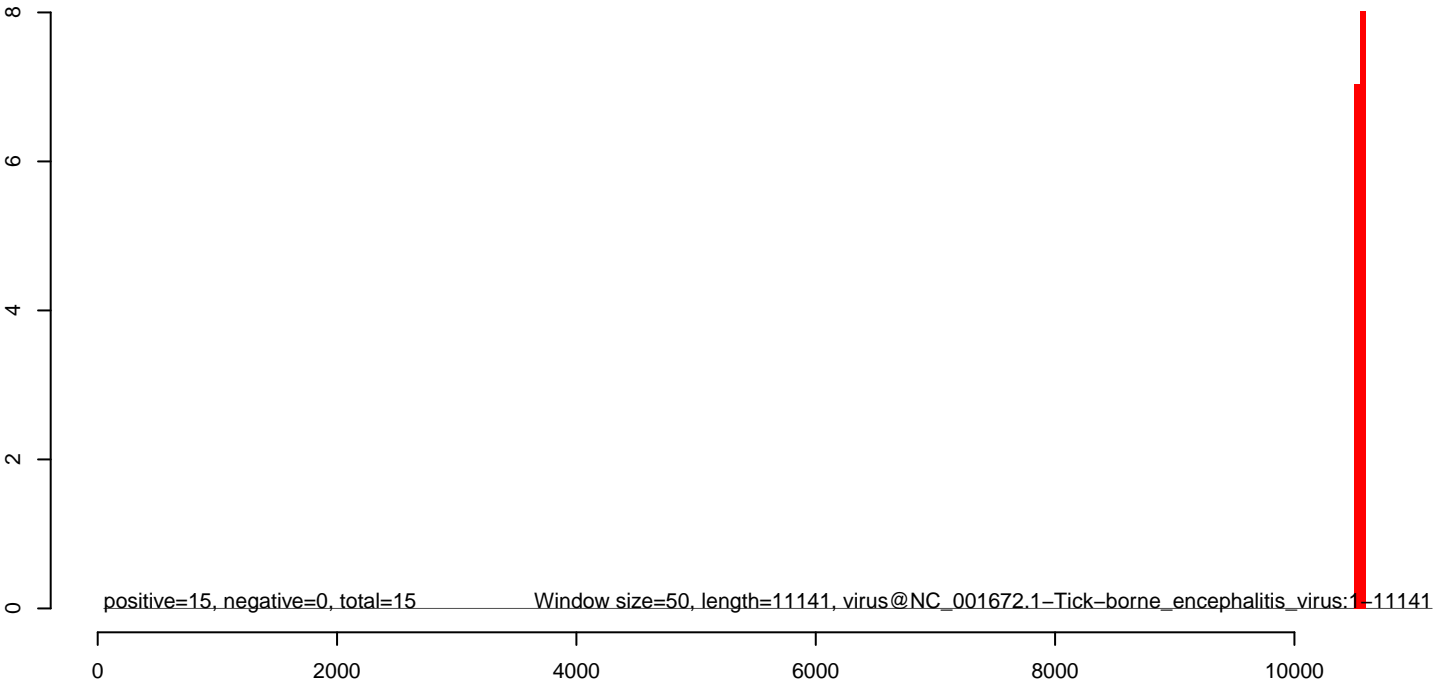
### AnGam\_Sua5b\_cells.18\_23.rep



### AnGam\_Sua5b\_cells.24\_35.rep



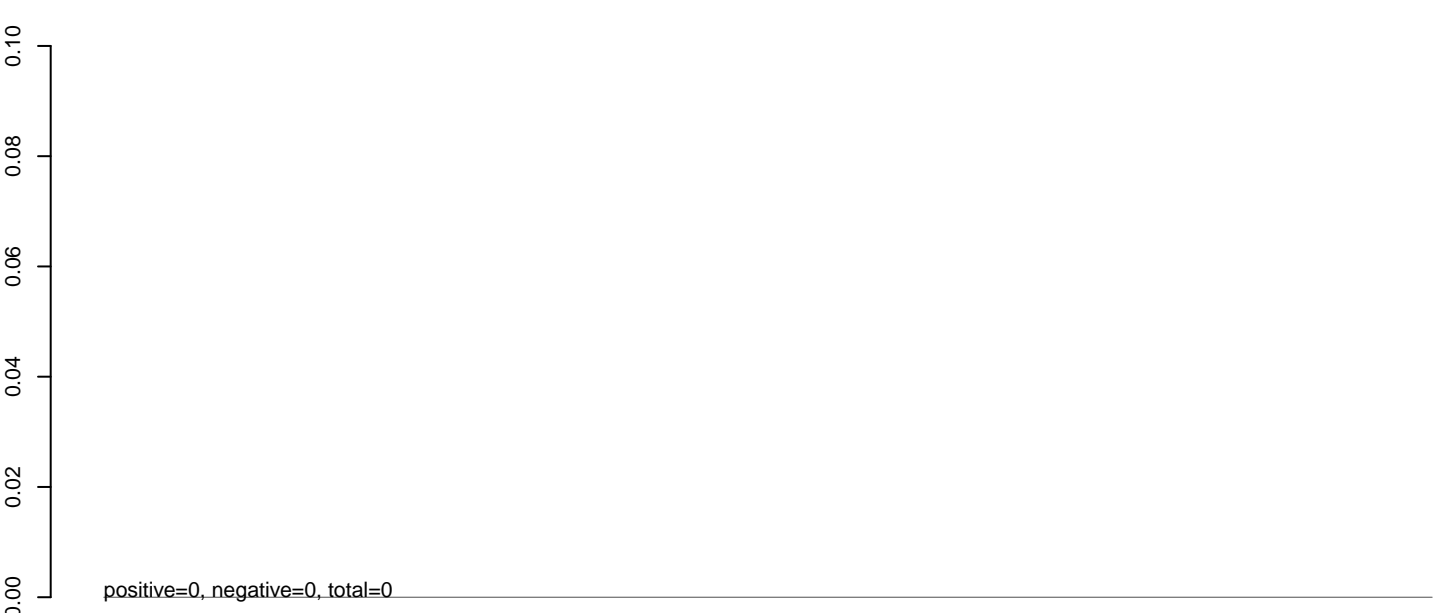
### AnGam\_Sua5b\_cells.rep



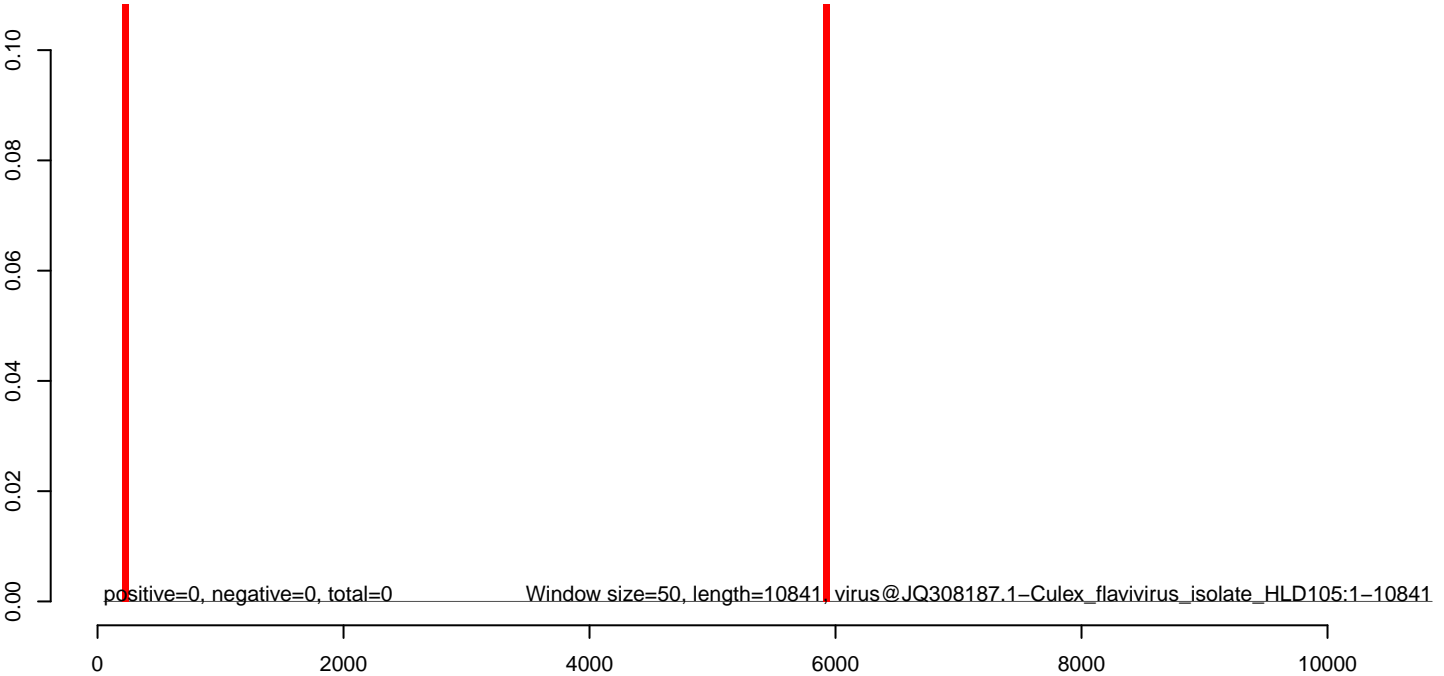
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep



AnGam\_Sua5b\_cells.rep

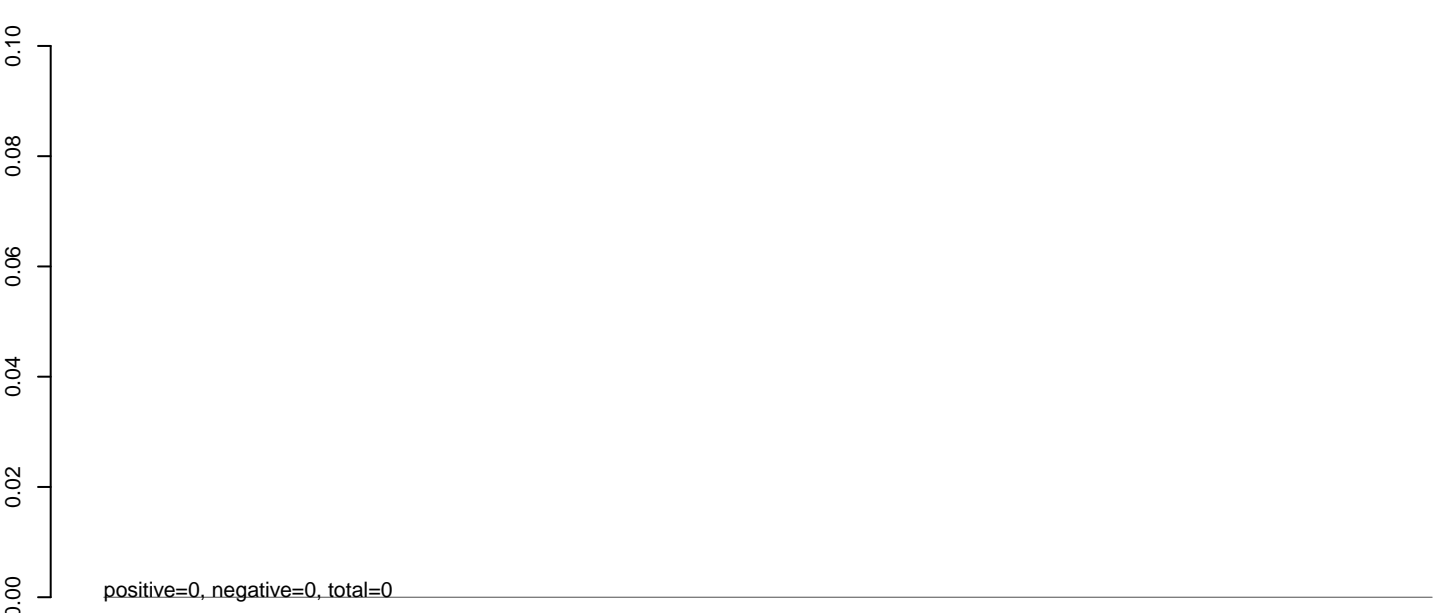




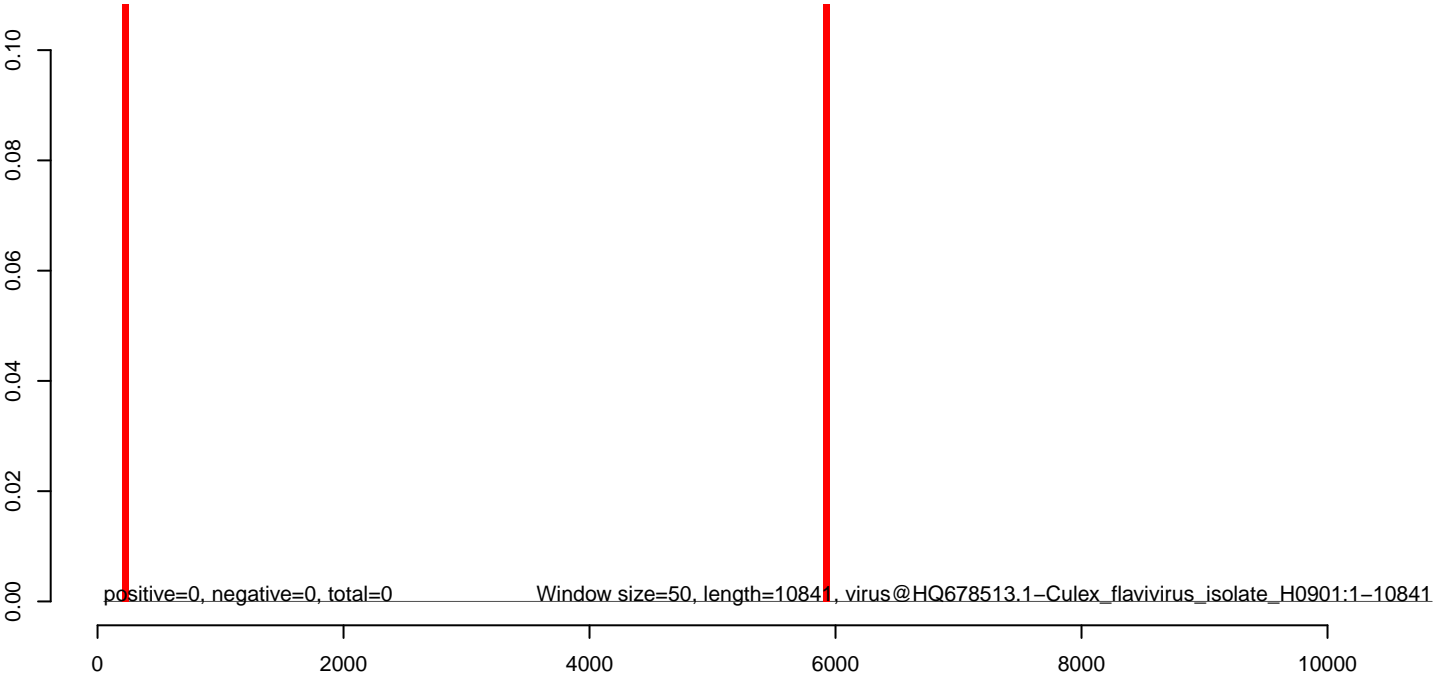
AnGam\_Sua5b\_cells.18\_23.rep



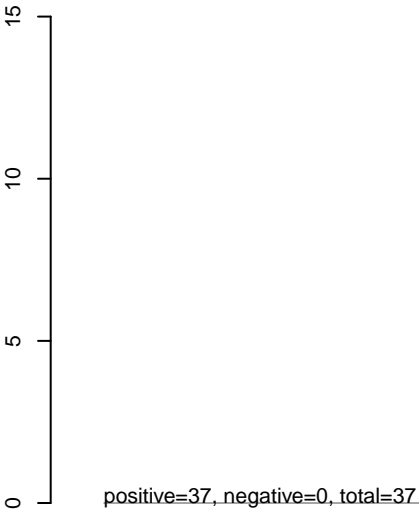
AnGam\_Sua5b\_cells.24\_35.rep



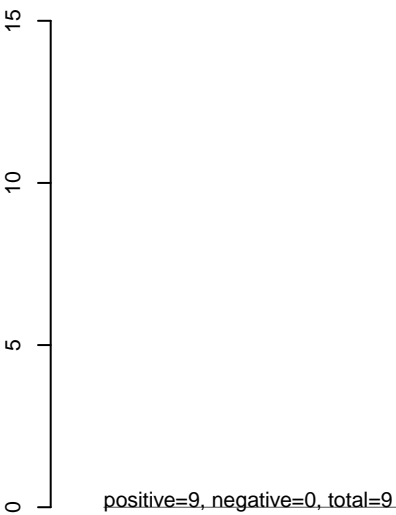
AnGam\_Sua5b\_cells.rep



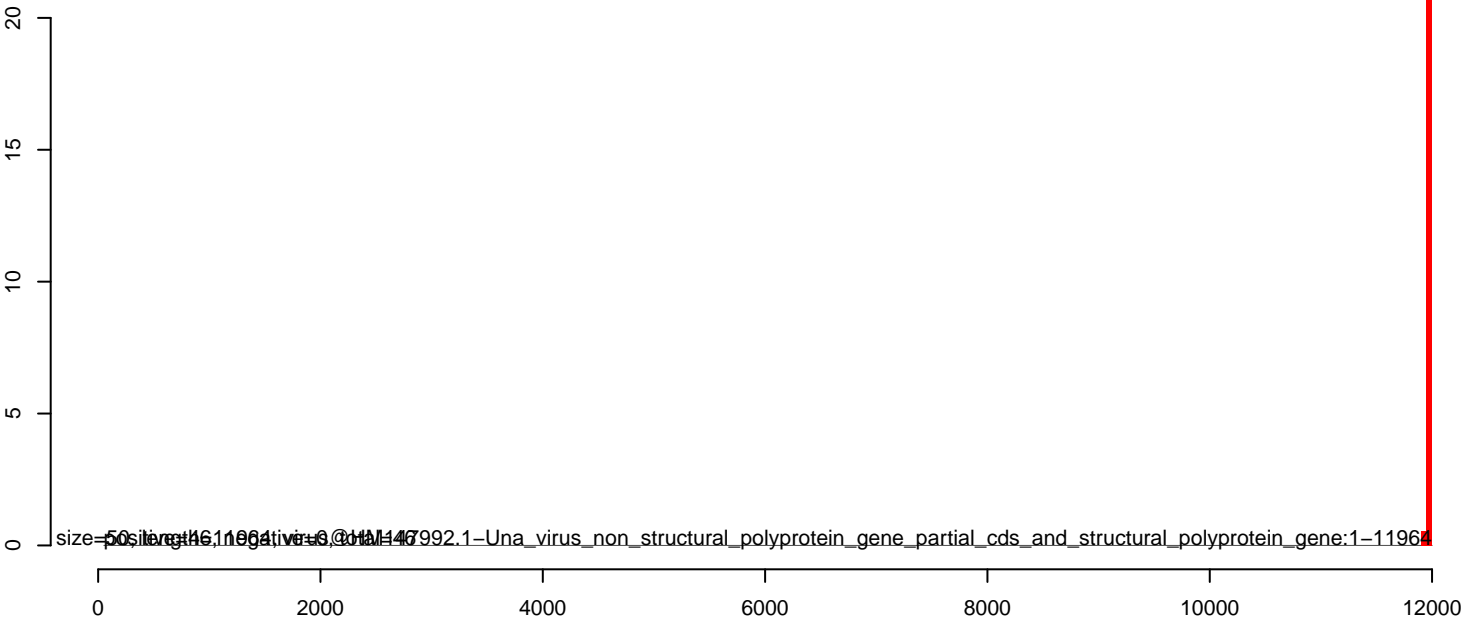
AnGam\_Sua5b\_cells.18\_23.rep



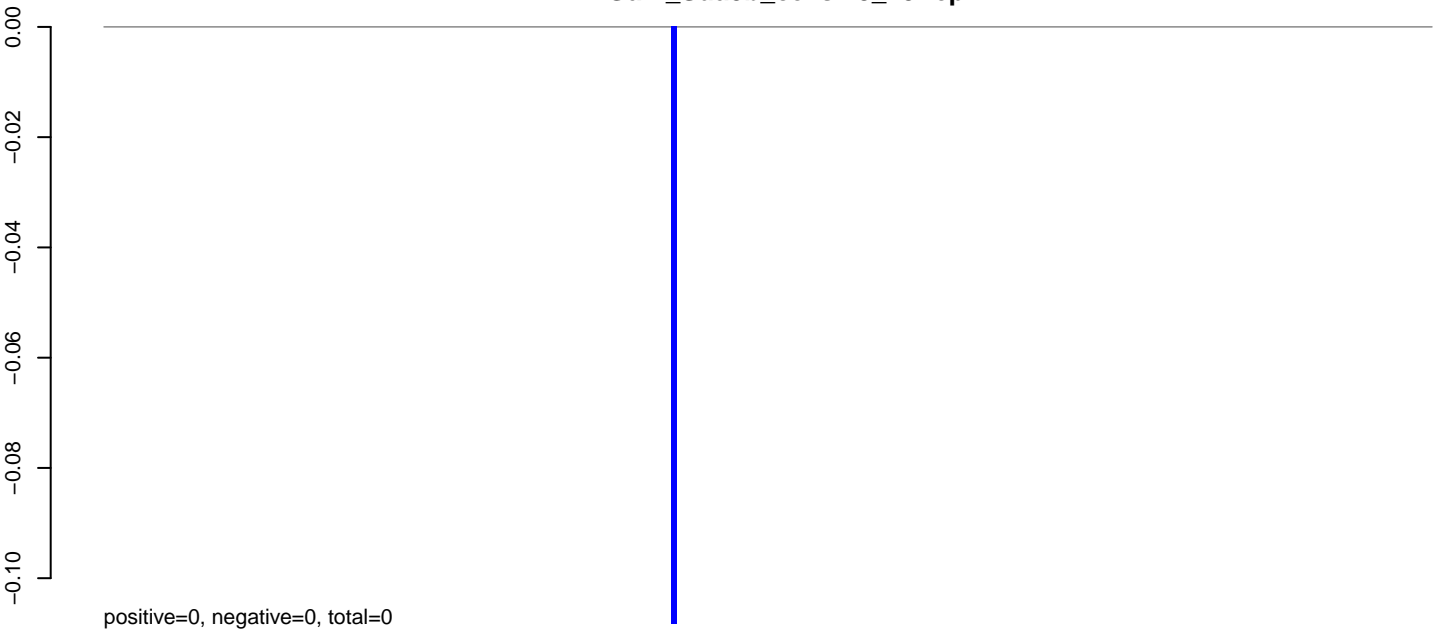
AnGam\_Sua5b\_cells.24\_35.rep



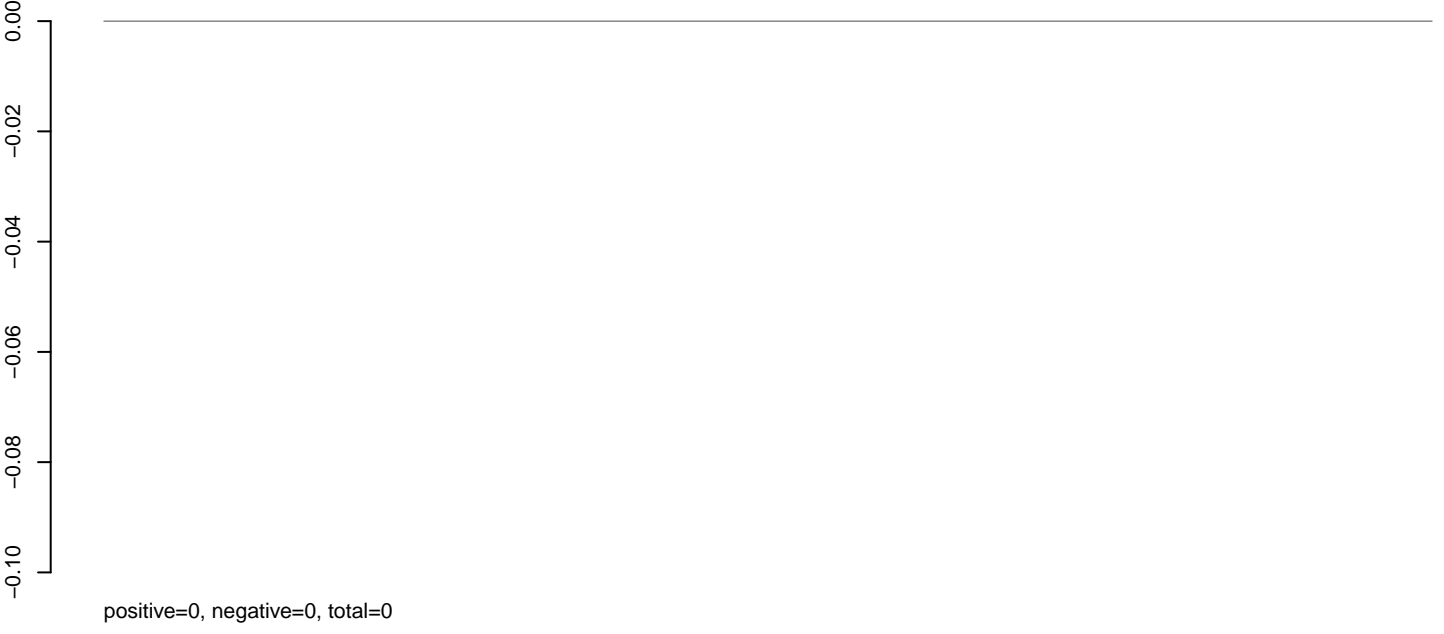
AnGam\_Sua5b\_cells.rep



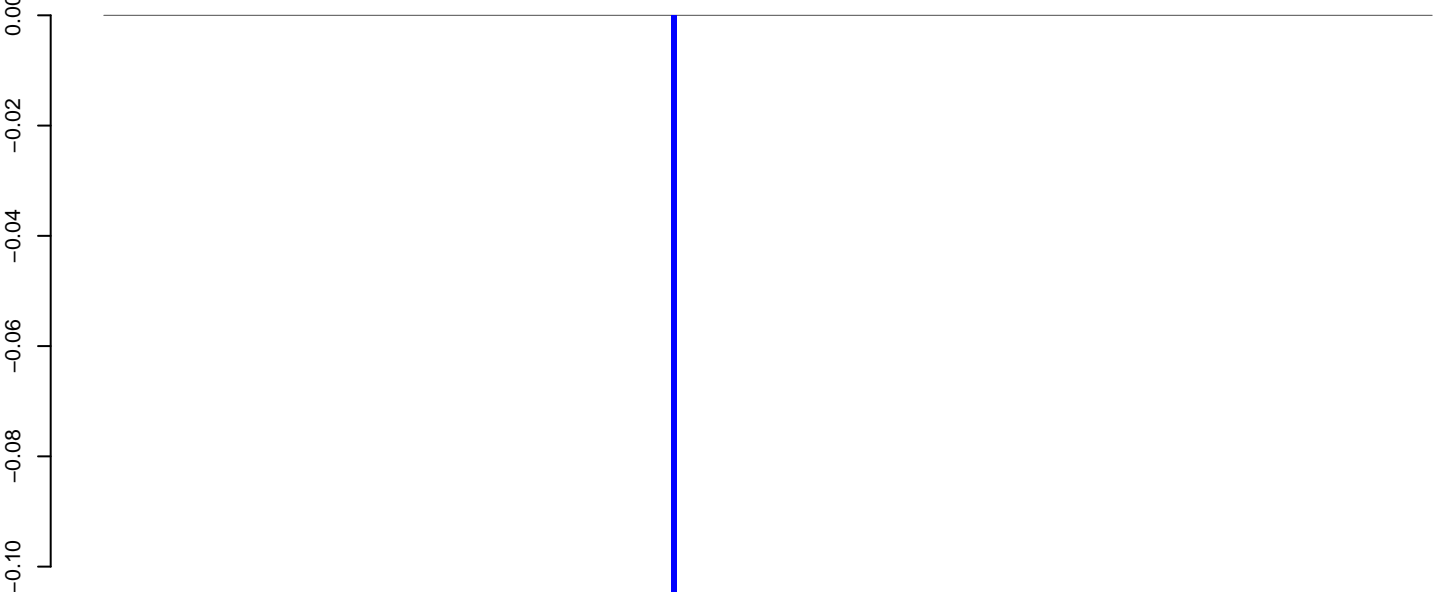
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep



AnGam\_Sua5b\_cells.rep



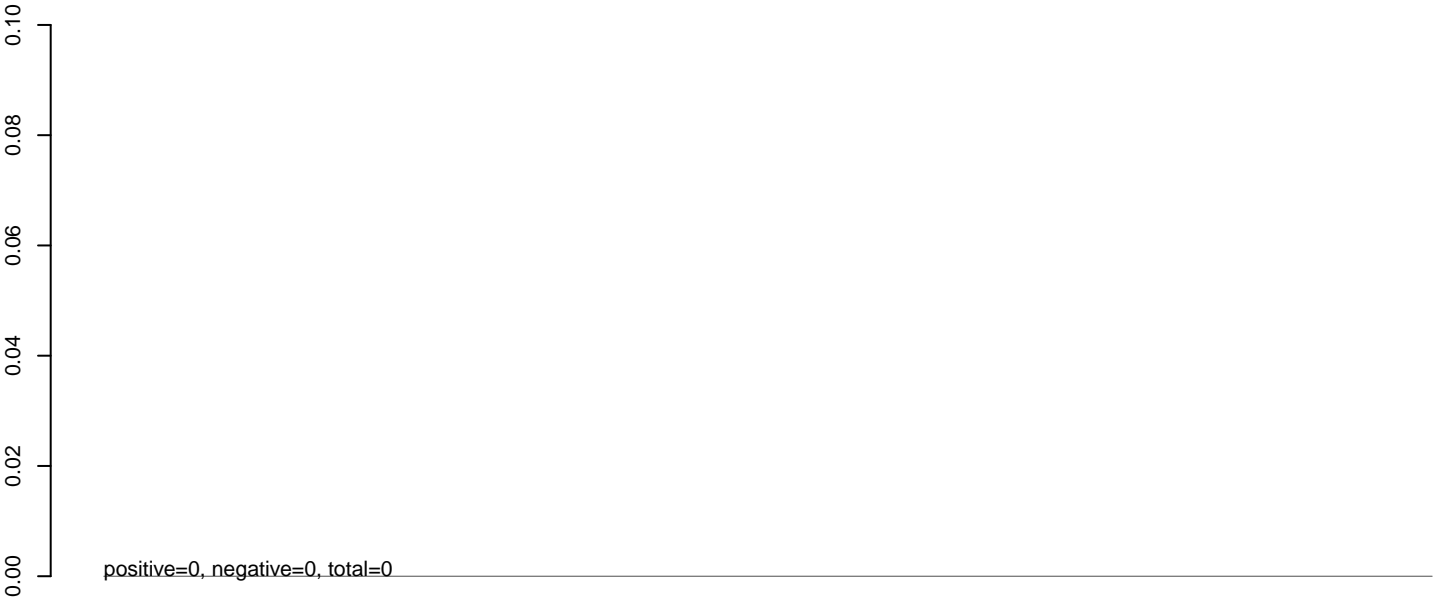
1728, virus\_010203405, total=0, virus\_strain\_F47\_CI\_2004\_segment\_S\_plus\_segment\_M\_plus\_segment\_L\_nucleocapsid\_N\_gene:1-11728

0 2000 4000 6000 8000 10000 12000

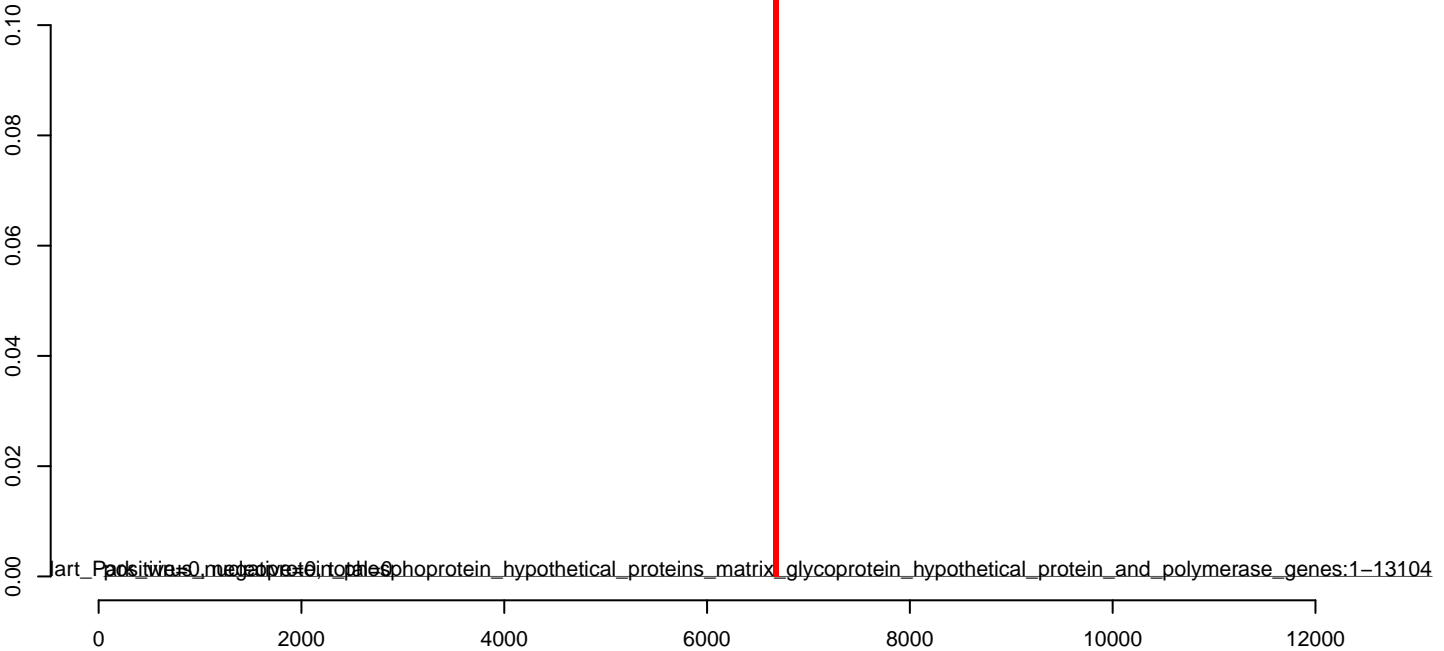
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep



AnGam\_Sua5b\_cells.rep



AnGam\_Sua5b\_cells.18\_23.rep

0.00  
-0.02  
-0.04  
-0.06  
-0.08  
-0.10

positive=0, negative=0, total=0

AnGam\_Sua5b\_cells.24\_35.rep

0.00  
-0.02  
-0.04  
-0.06  
-0.08  
-0.10

positive=0, negative=0, total=0

AnGam\_Sua5b\_cells.rep

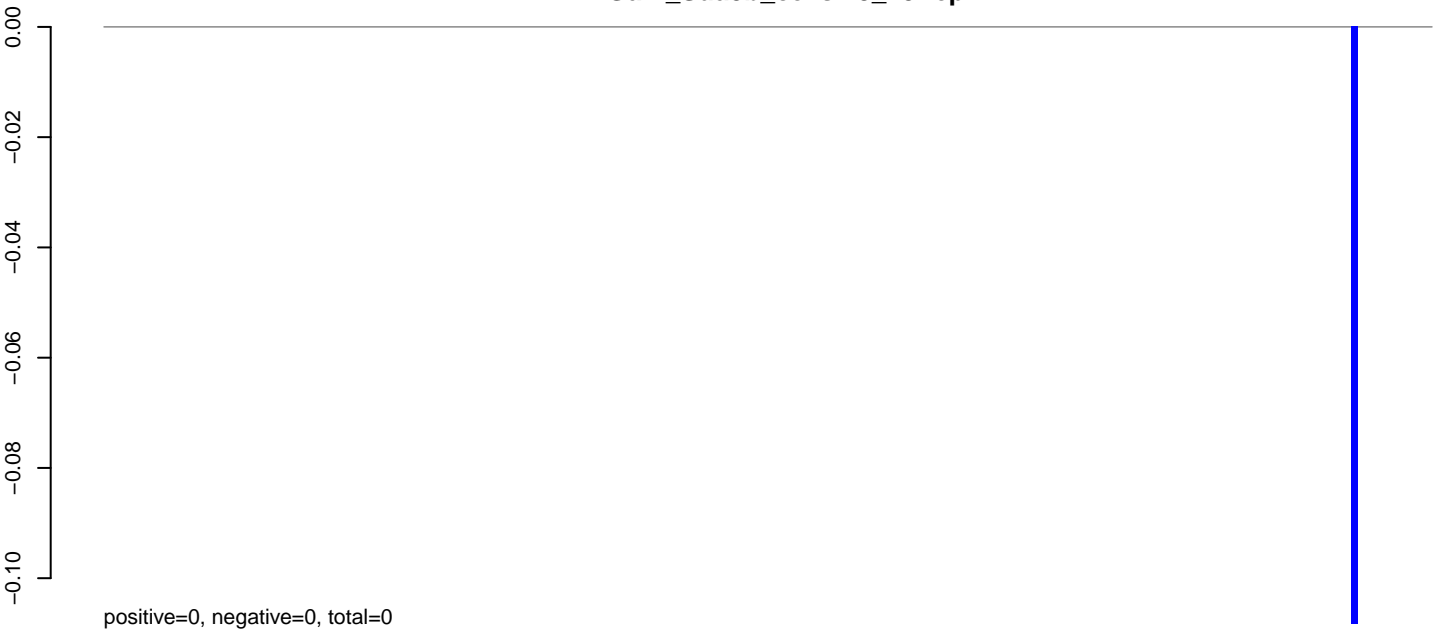
0.00  
-0.02  
-0.04  
-0.06  
-0.08  
-0.10

positive=0, negative=0, total=0

Window size=50, length=10588, virus@NC\_031327.1-Anopheles\_flavivirus\_variant\_1:1-10588

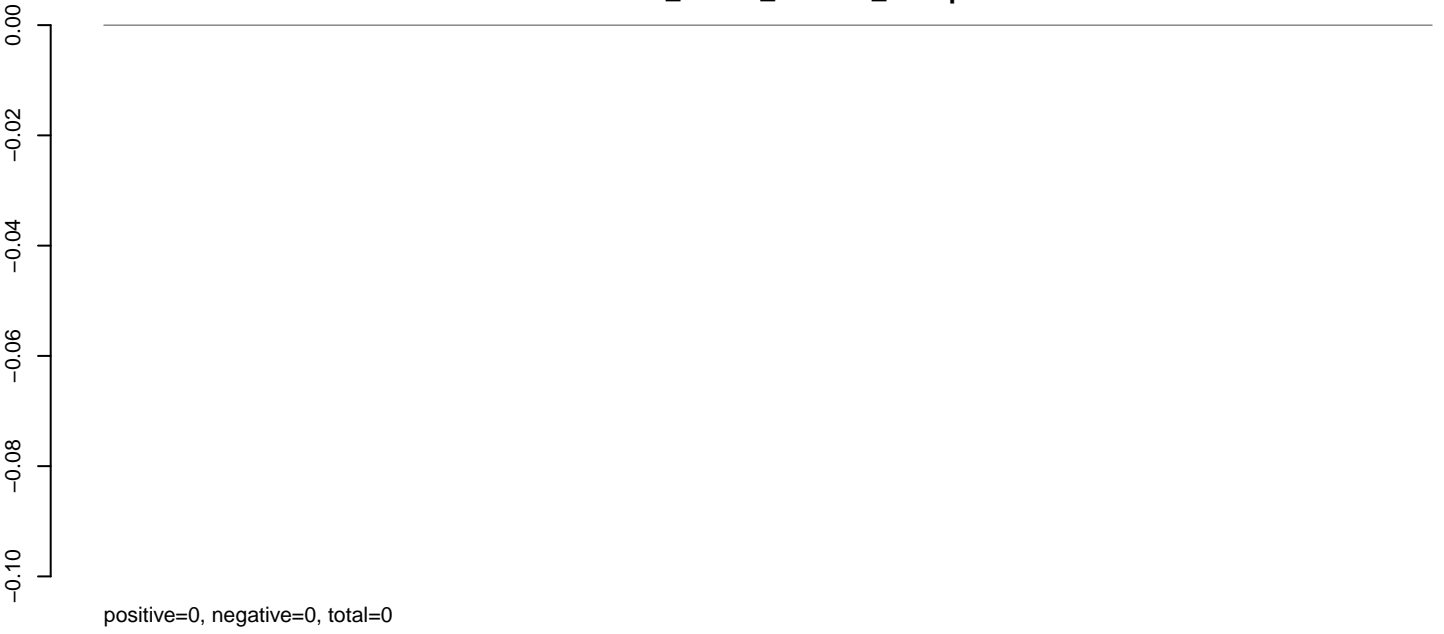
0 2000 4000 6000 8000 10000

AnGam\_Sua5b\_cells.18\_23.rep



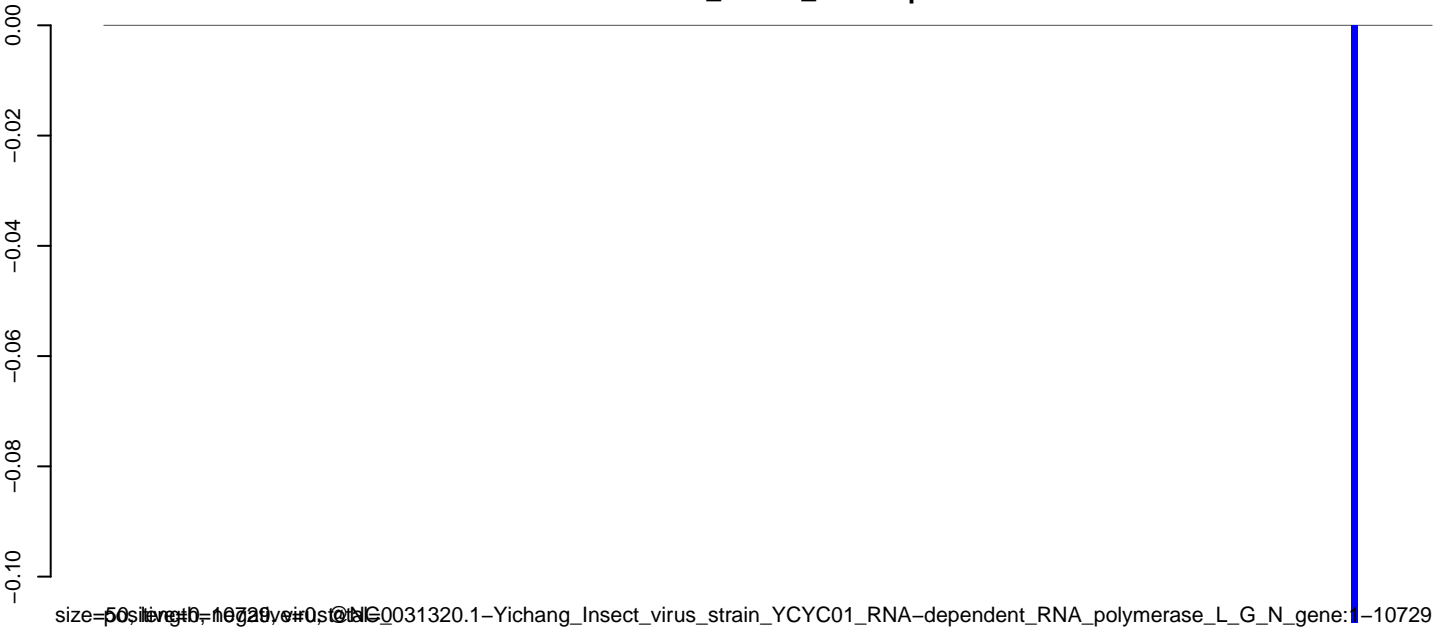
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AnGam\_Sua5b\_cells.24\_35.rep



positive=0, negative=0, total=0

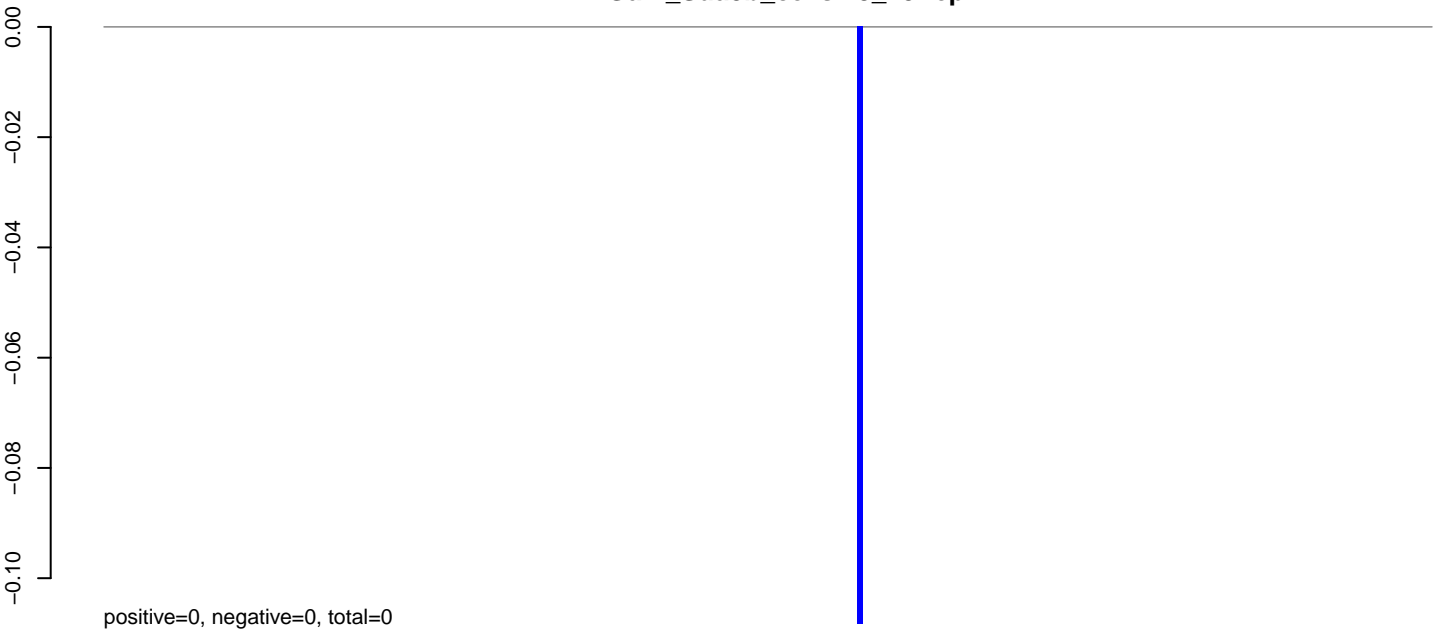
AnGam\_Sua5b\_cells.rep



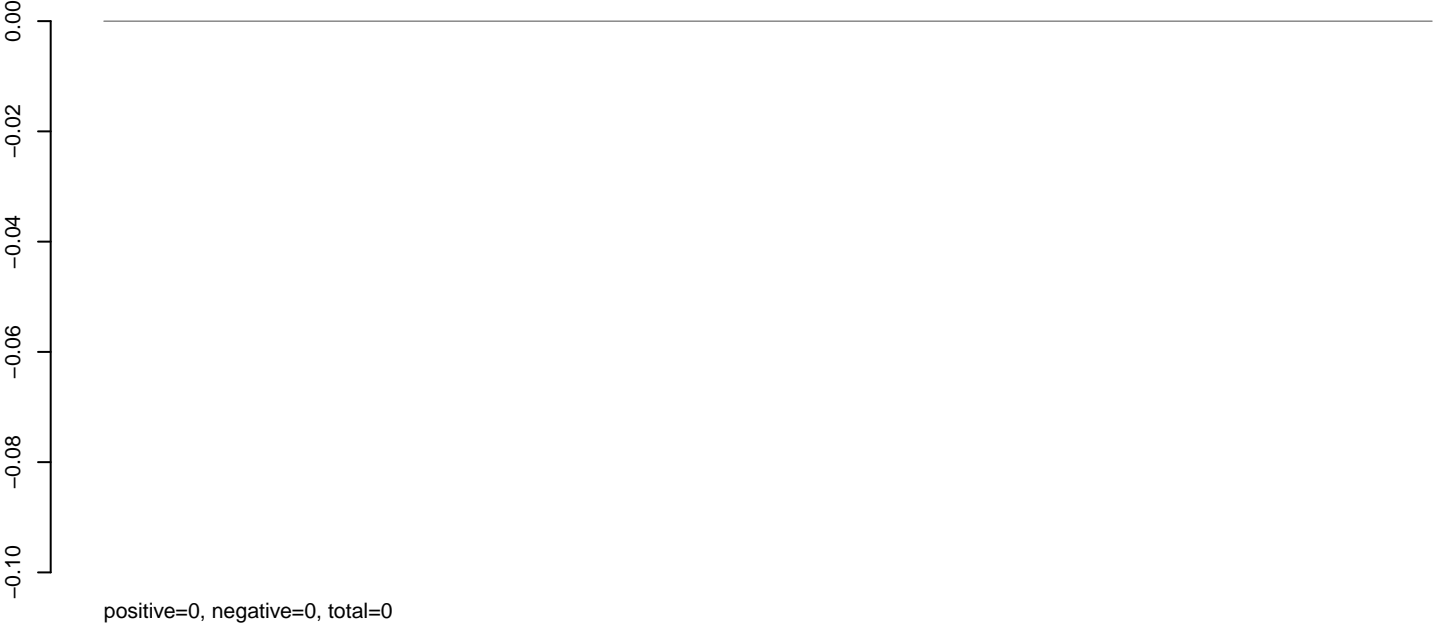
size=50, length=10729, version=0, chr=0, contig=0031320.1-Yichang\_Insect\_virus\_strain\_YCYC01\_RNA-dependent\_RNA\_polymerase\_L\_G\_N\_gene:1-10729

0 2000 4000 6000 8000 10000

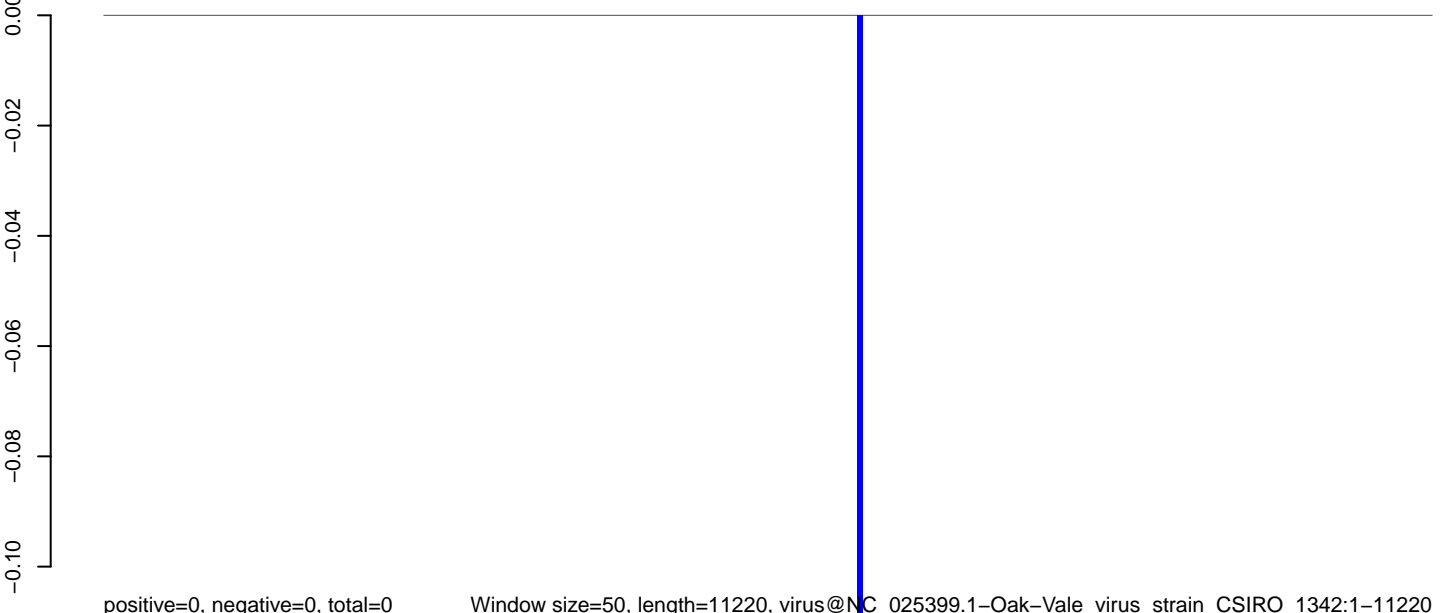
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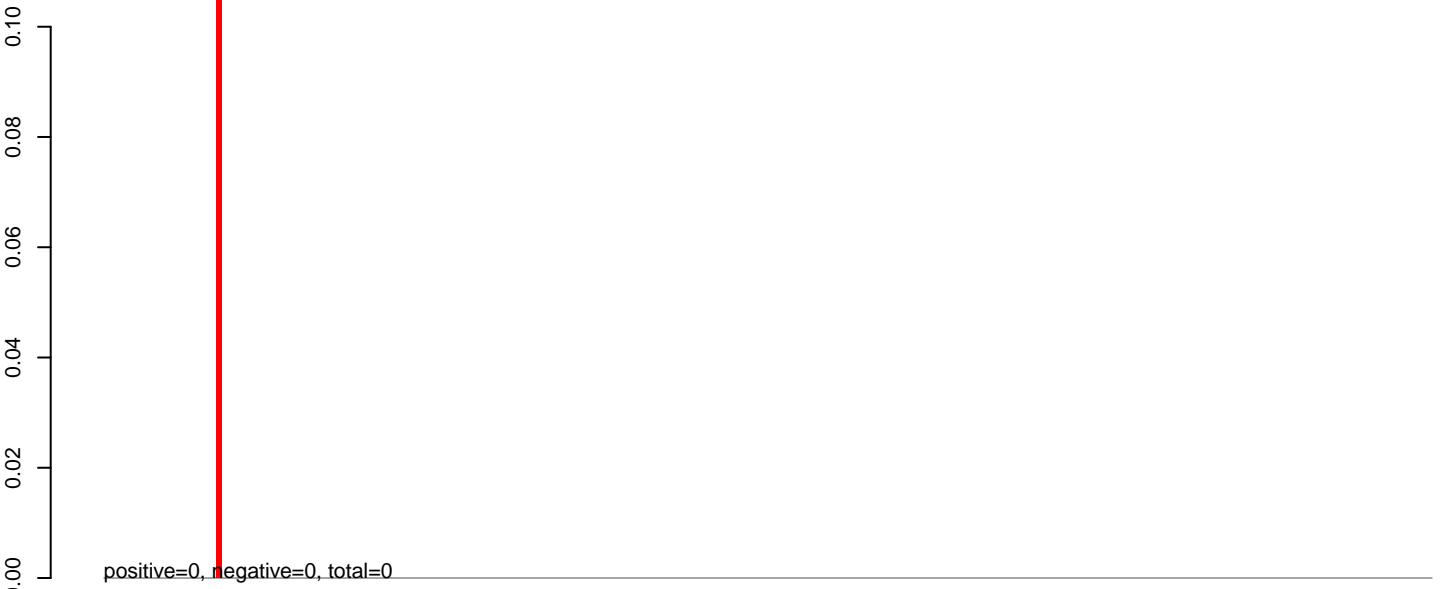
AnGam\_Sua5b\_cells.24\_35.rep



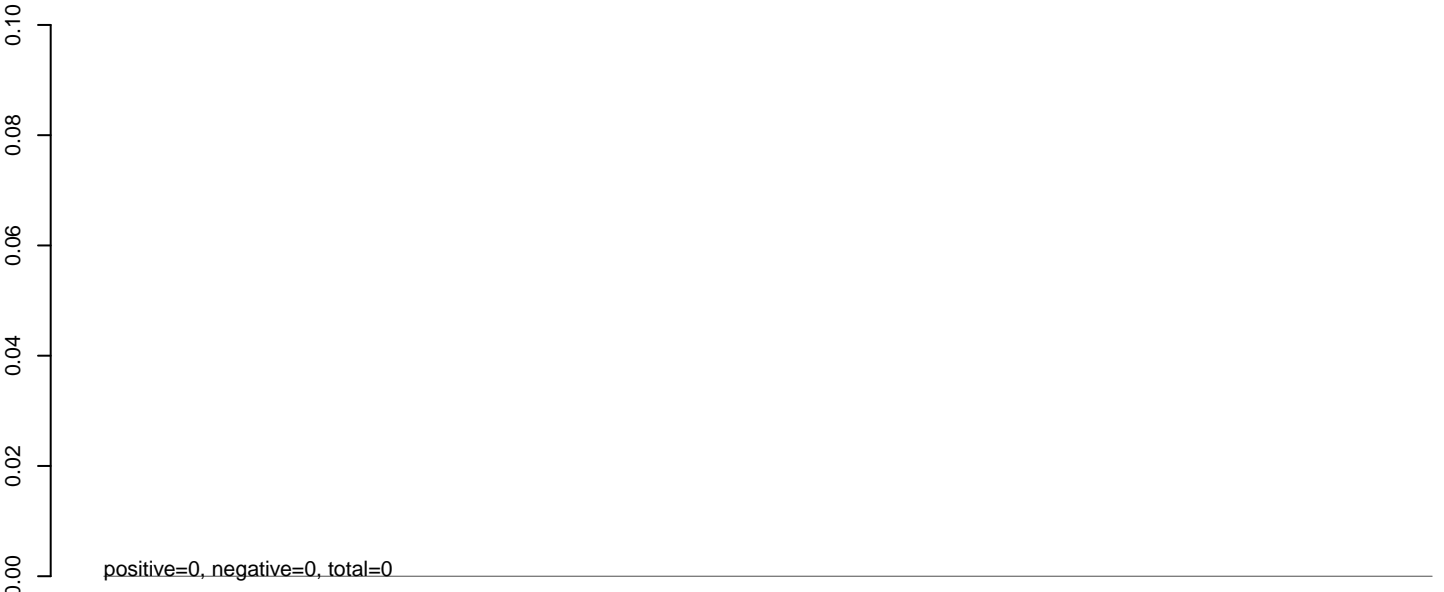
AnGam\_Sua5b\_cells.rep



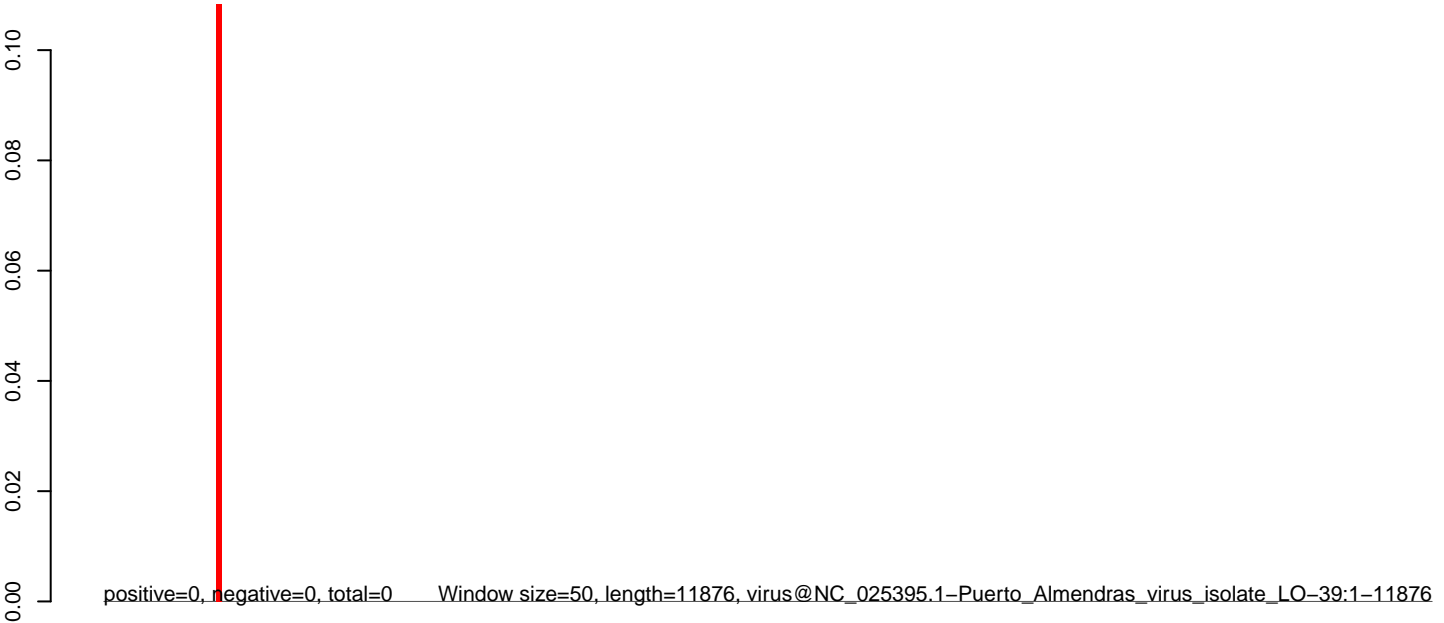
### AnGam\_Sua5b\_cells.18\_23.rep



### AnGam\_Sua5b\_cells.24\_35.rep



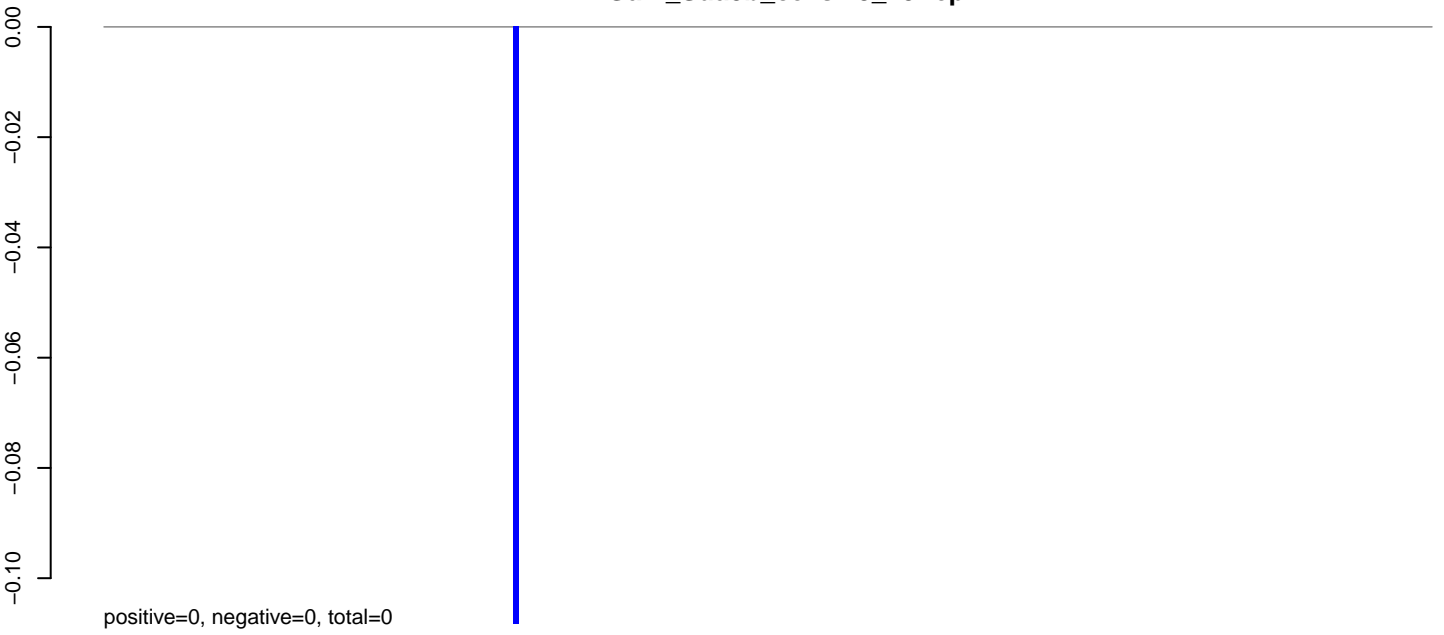
### AnGam\_Sua5b\_cells.rep



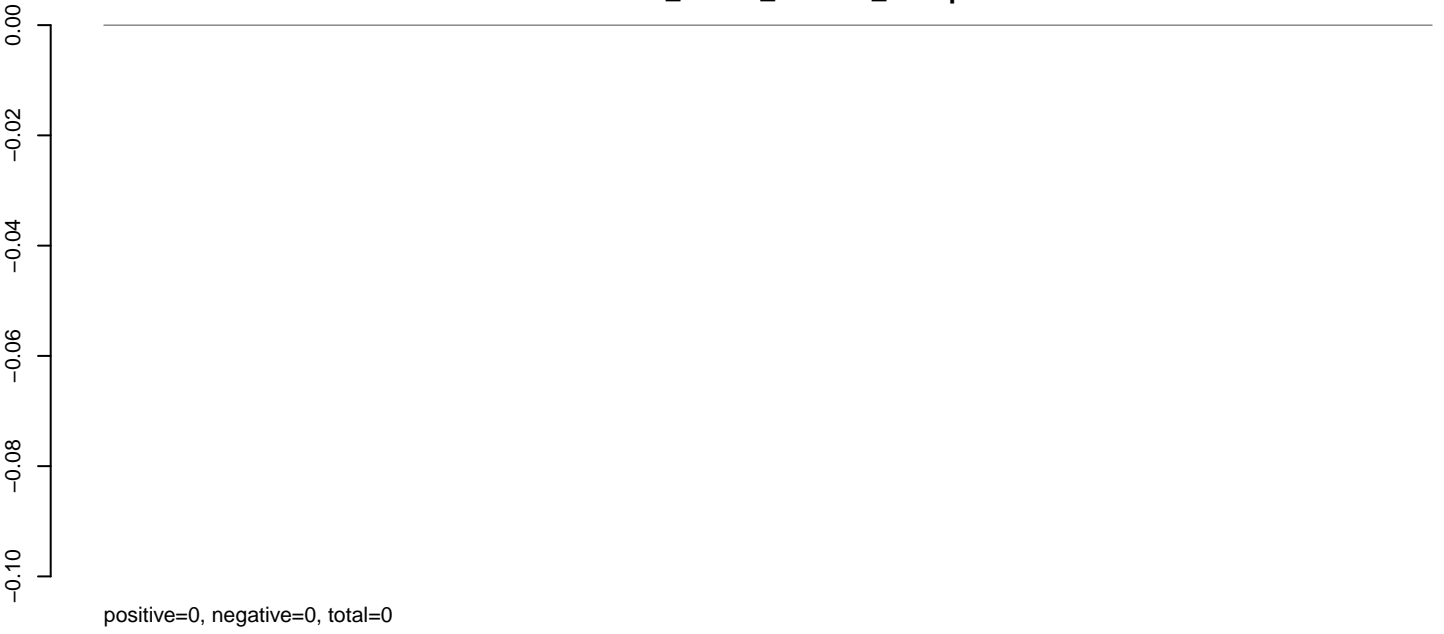
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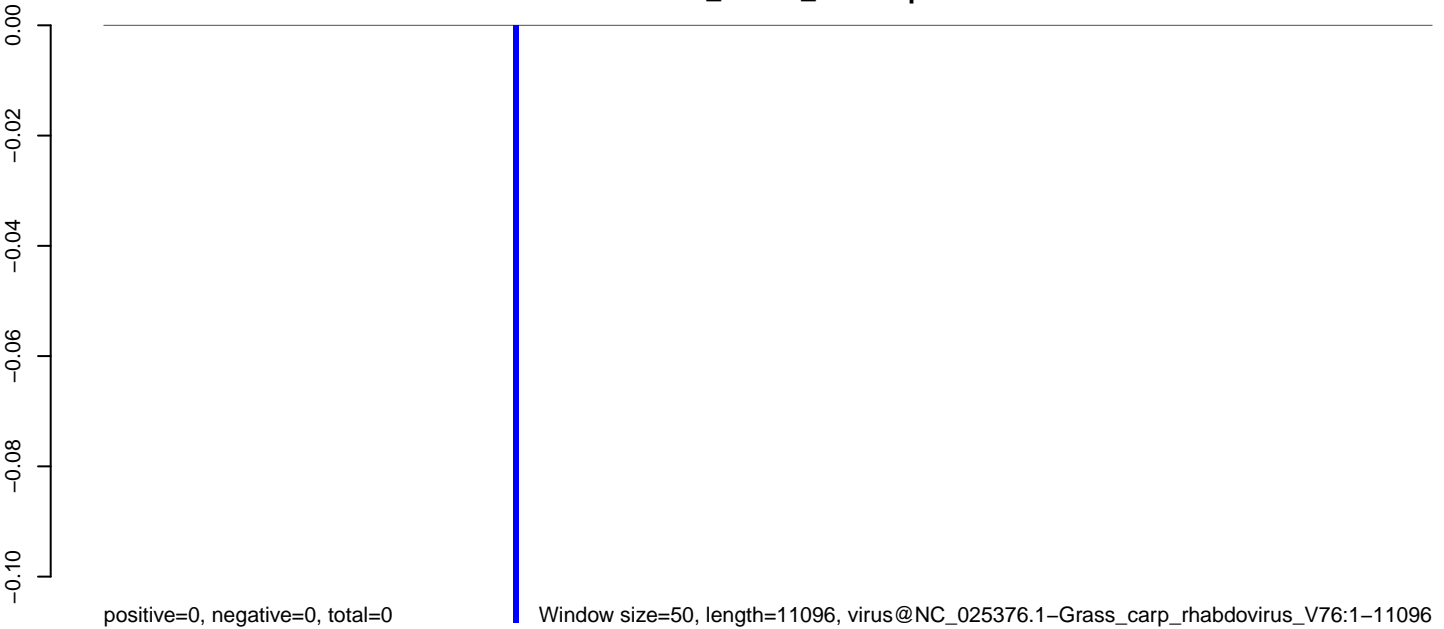
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep

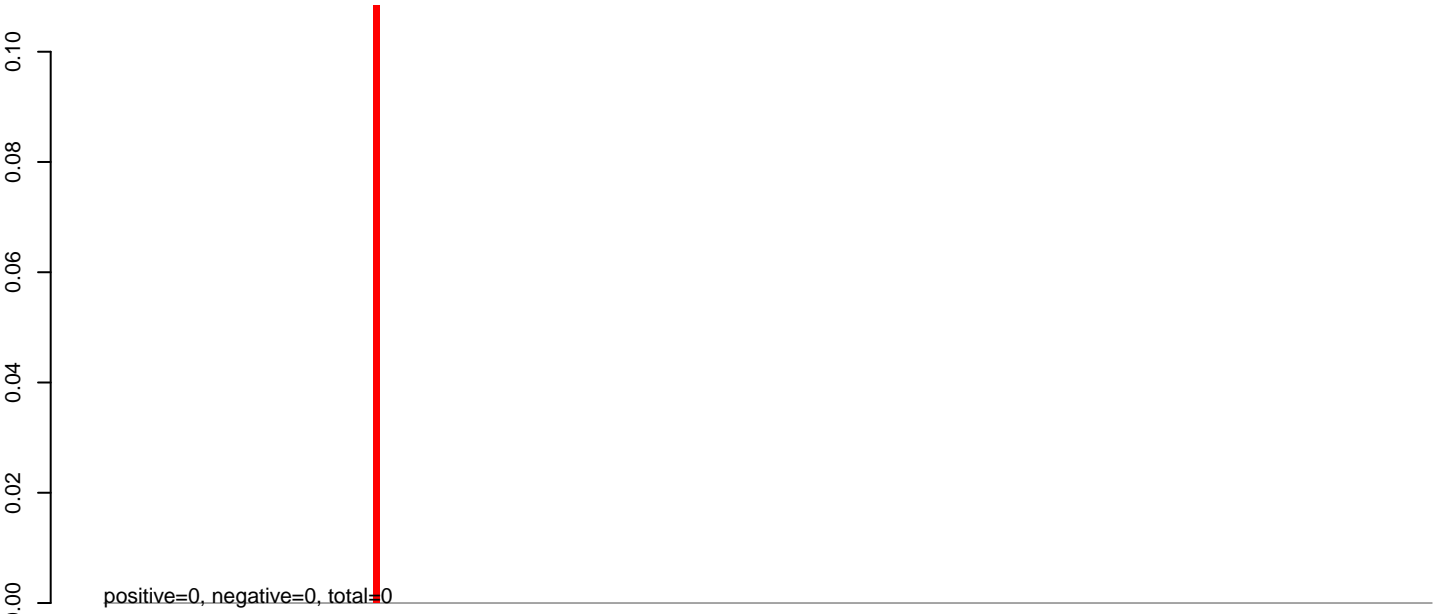


AnGam\_Sua5b\_cells.rep

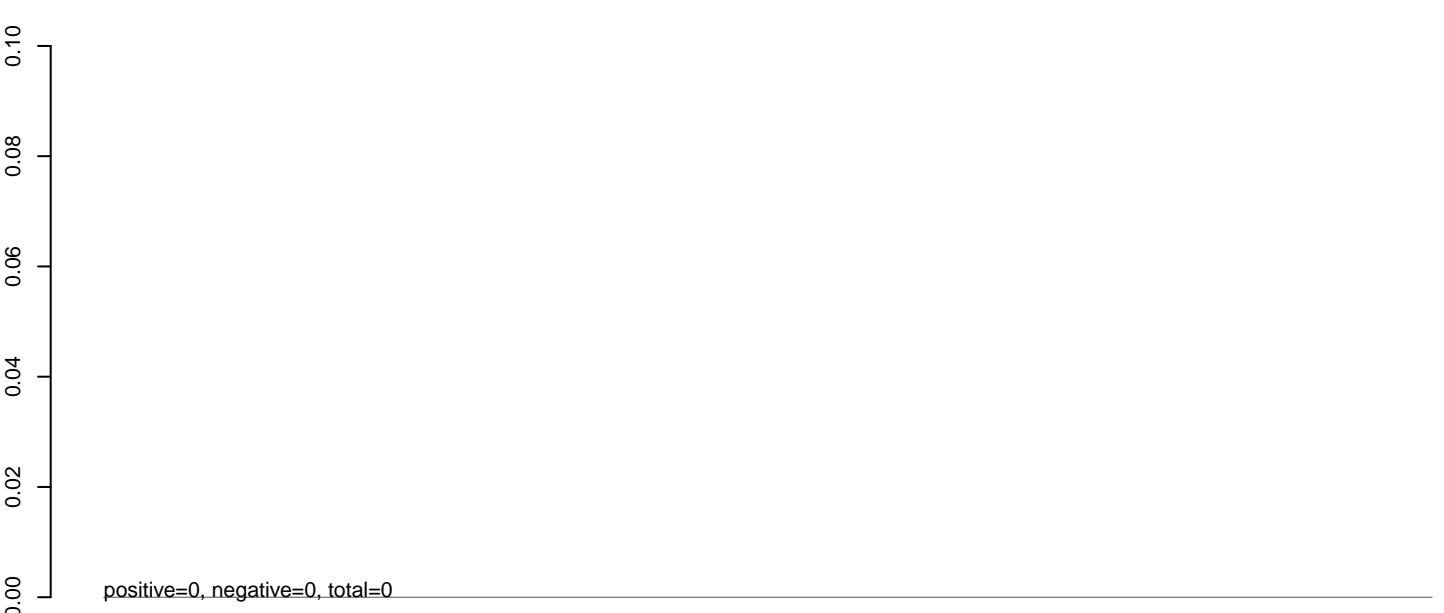


0 2000 4000 6000 8000 10000

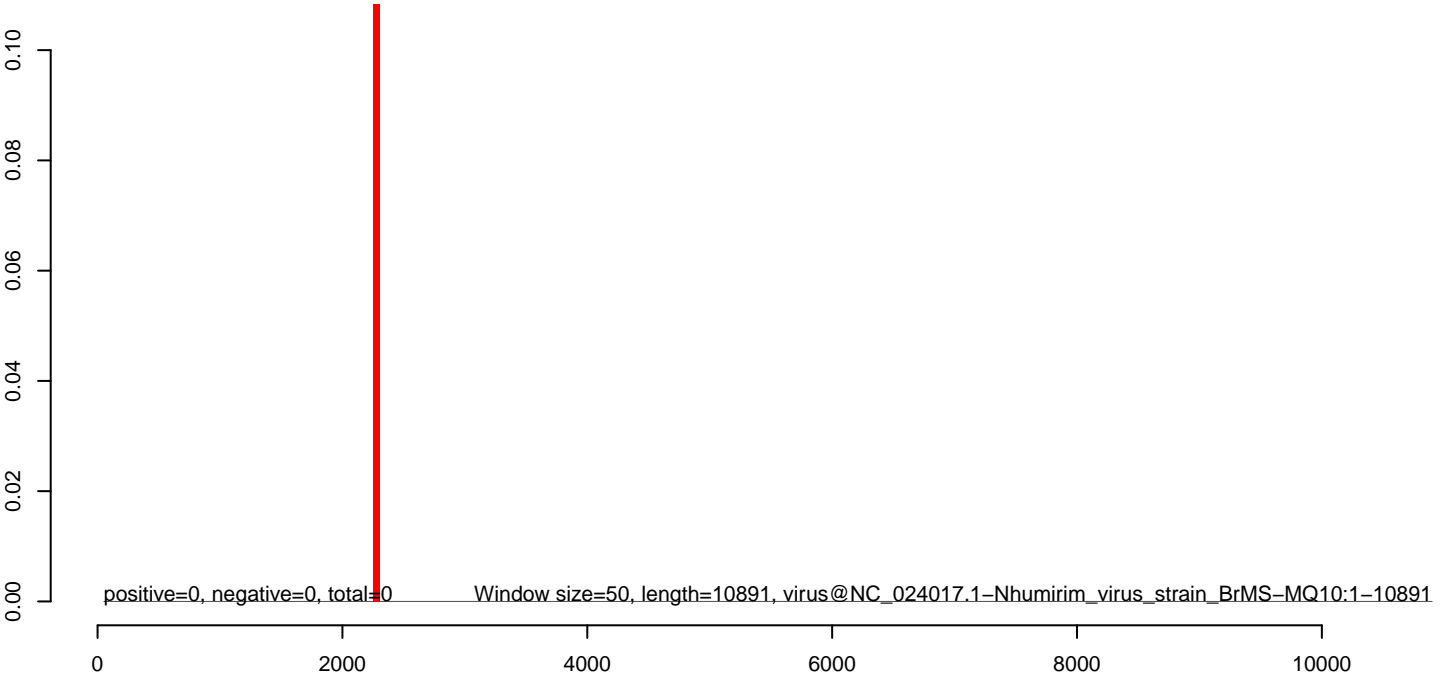
AnGam\_Sua5b\_cells.18\_23.rep



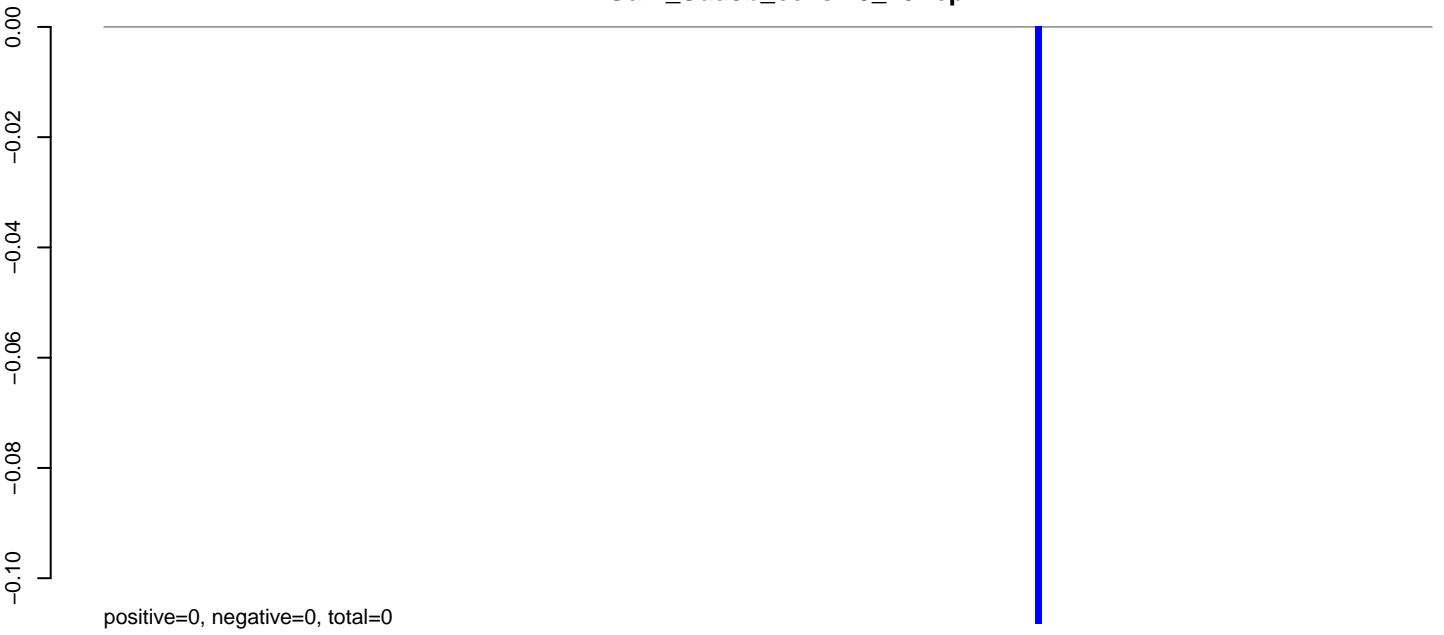
AnGam\_Sua5b\_cells.24\_35.rep



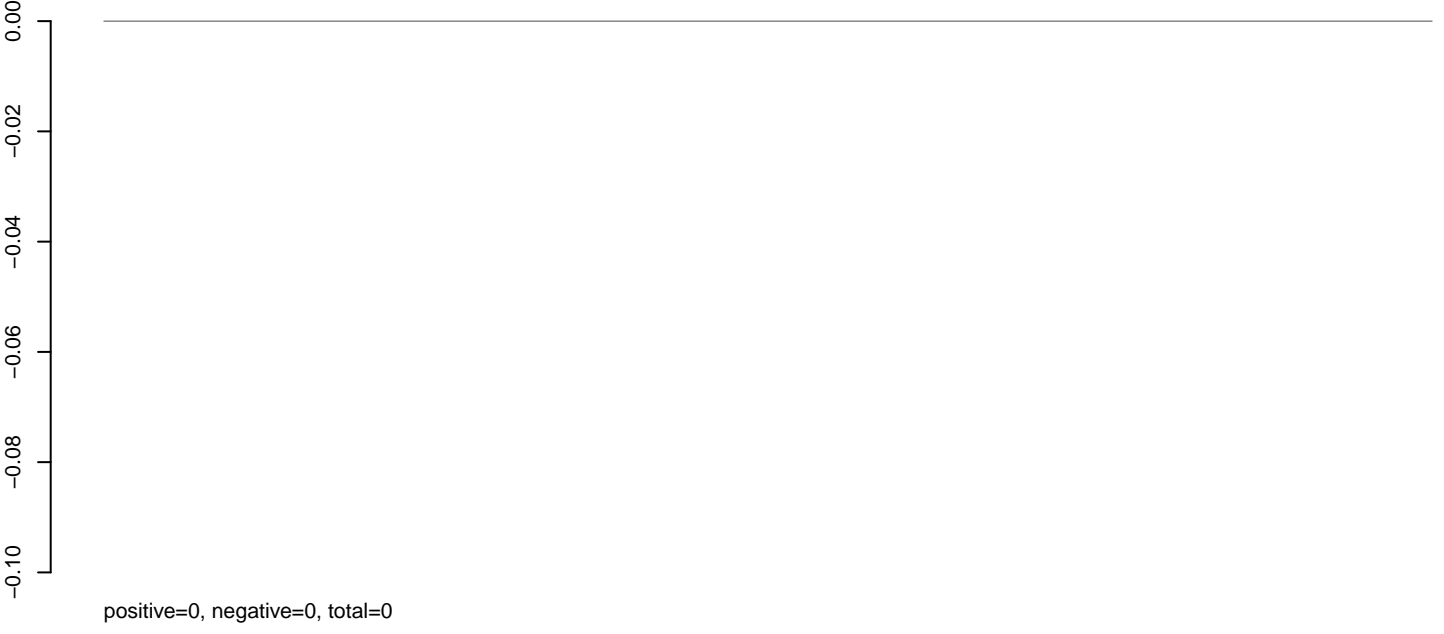
AnGam\_Sua5b\_cells.rep



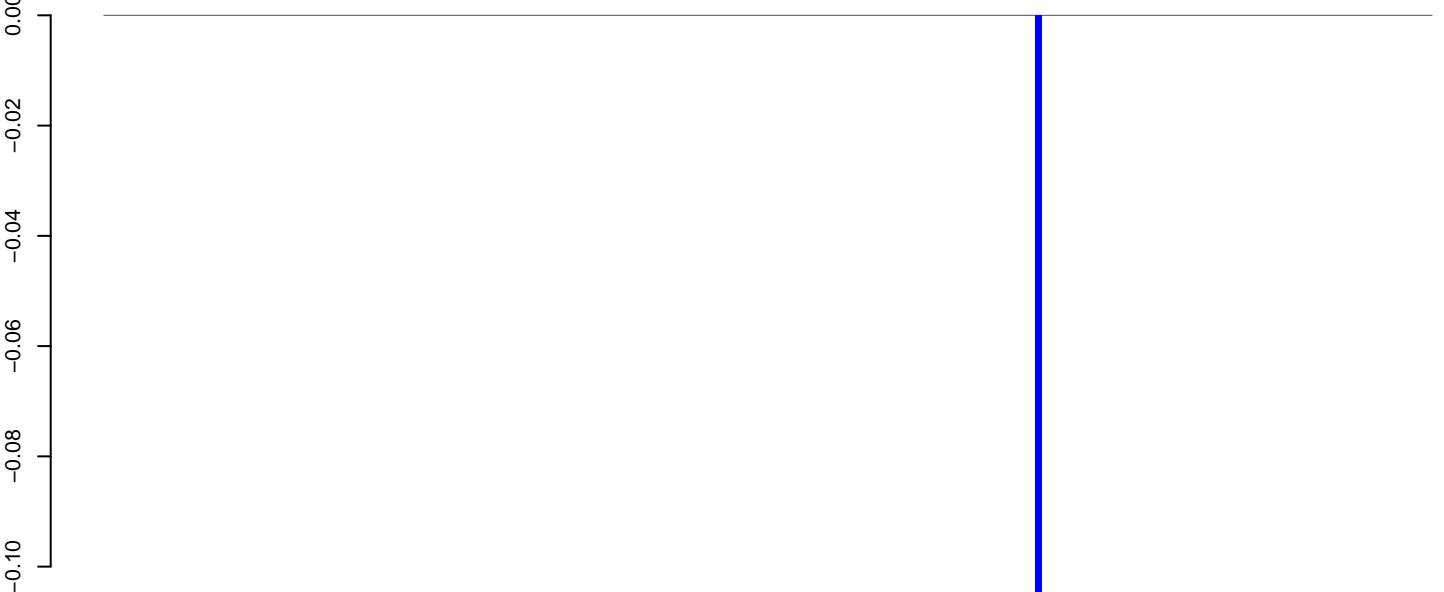
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AnGam\_Sua5b\_cells.24\_35.rep



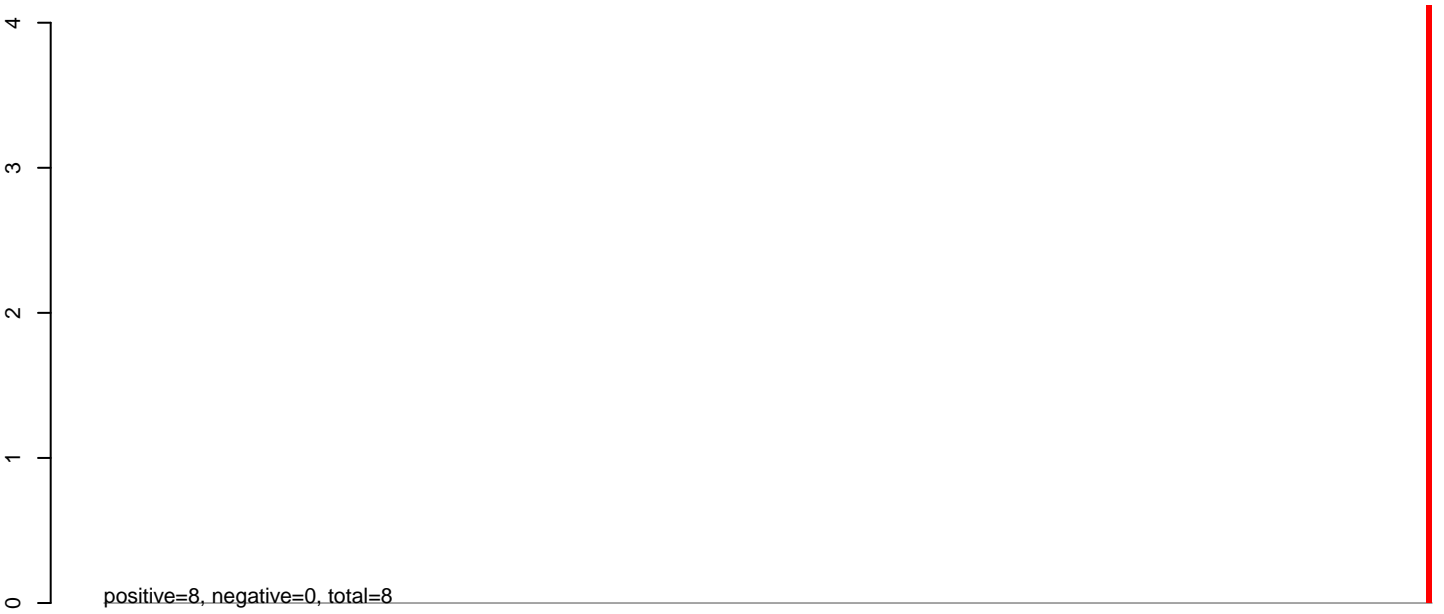
AnGam\_Sua5b\_cells.rep



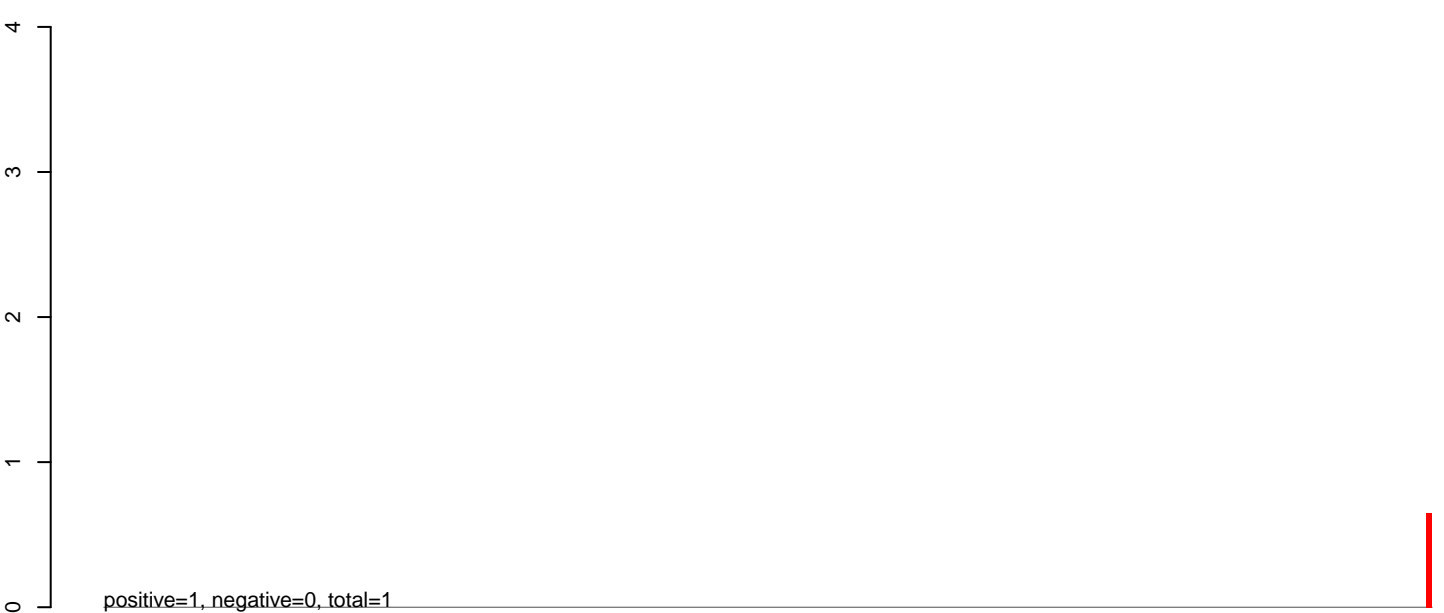
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0 2000 4000 6000 8000 10000

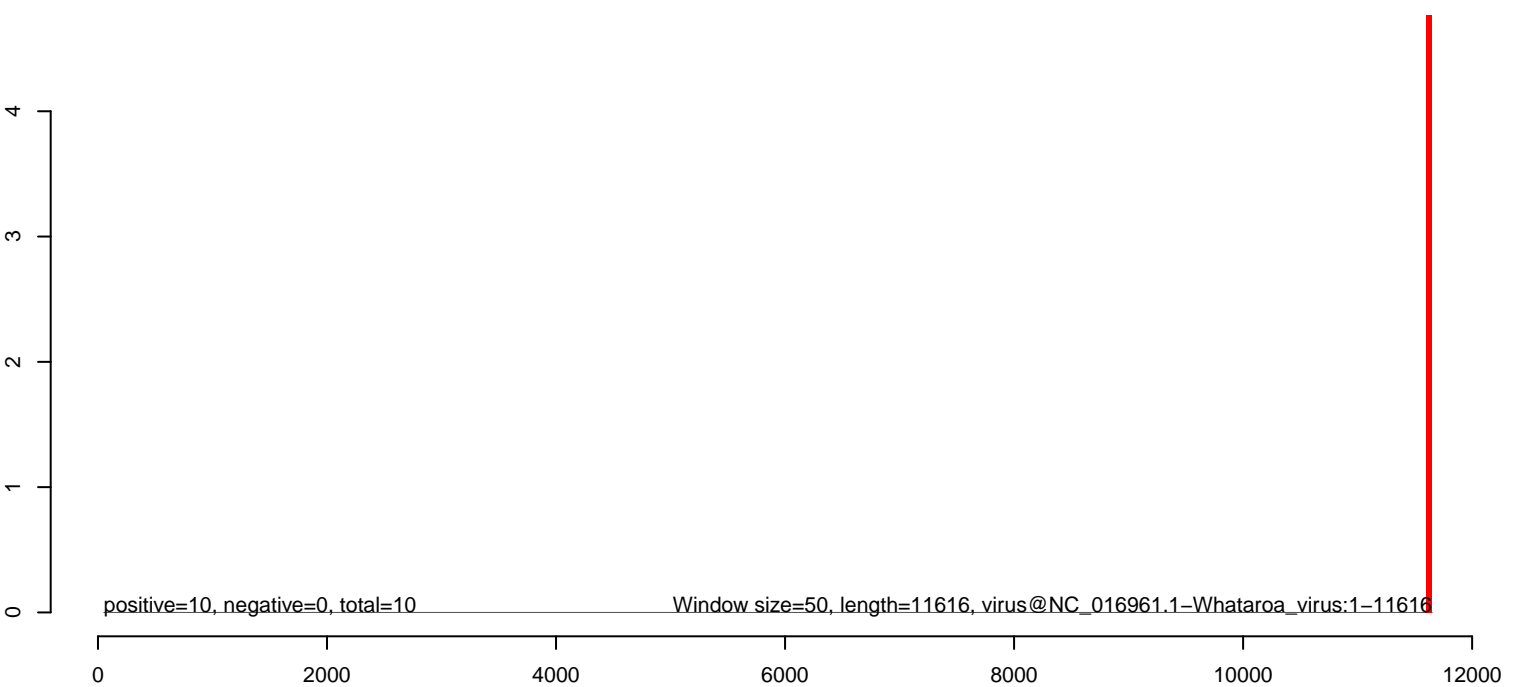
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AnGam\_Sua5b\_cells.24\_35.rep



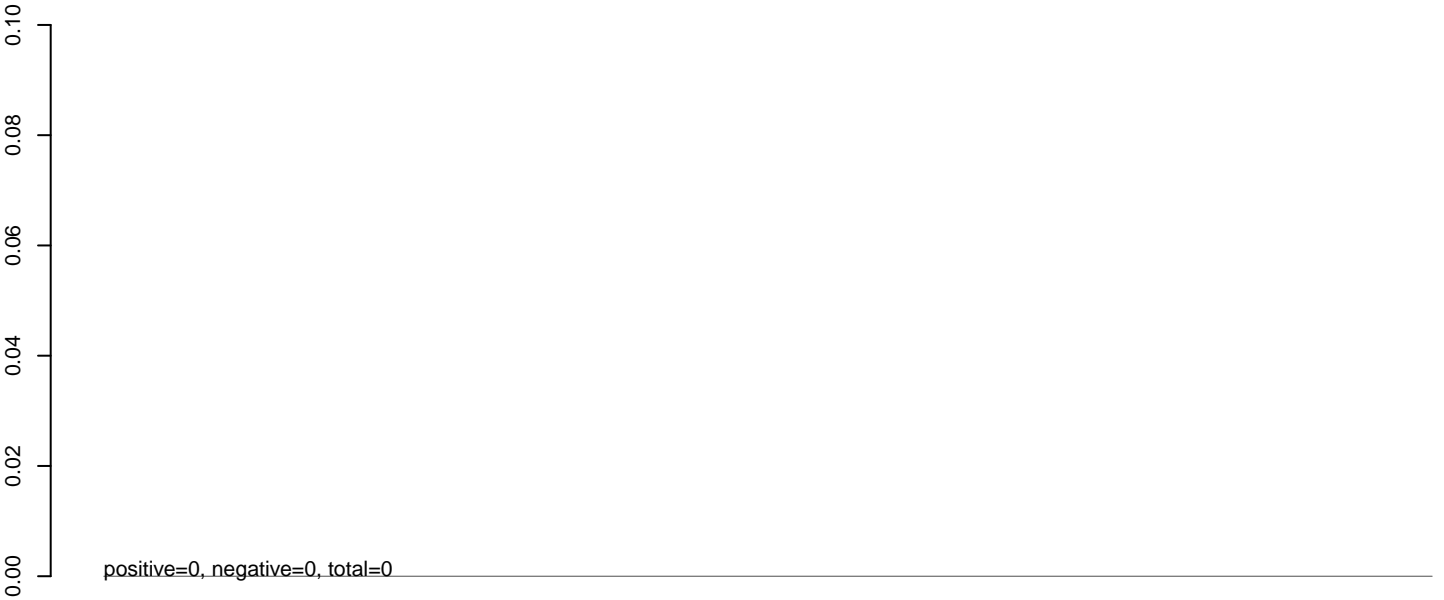
AnGam\_Sua5b\_cells.rep



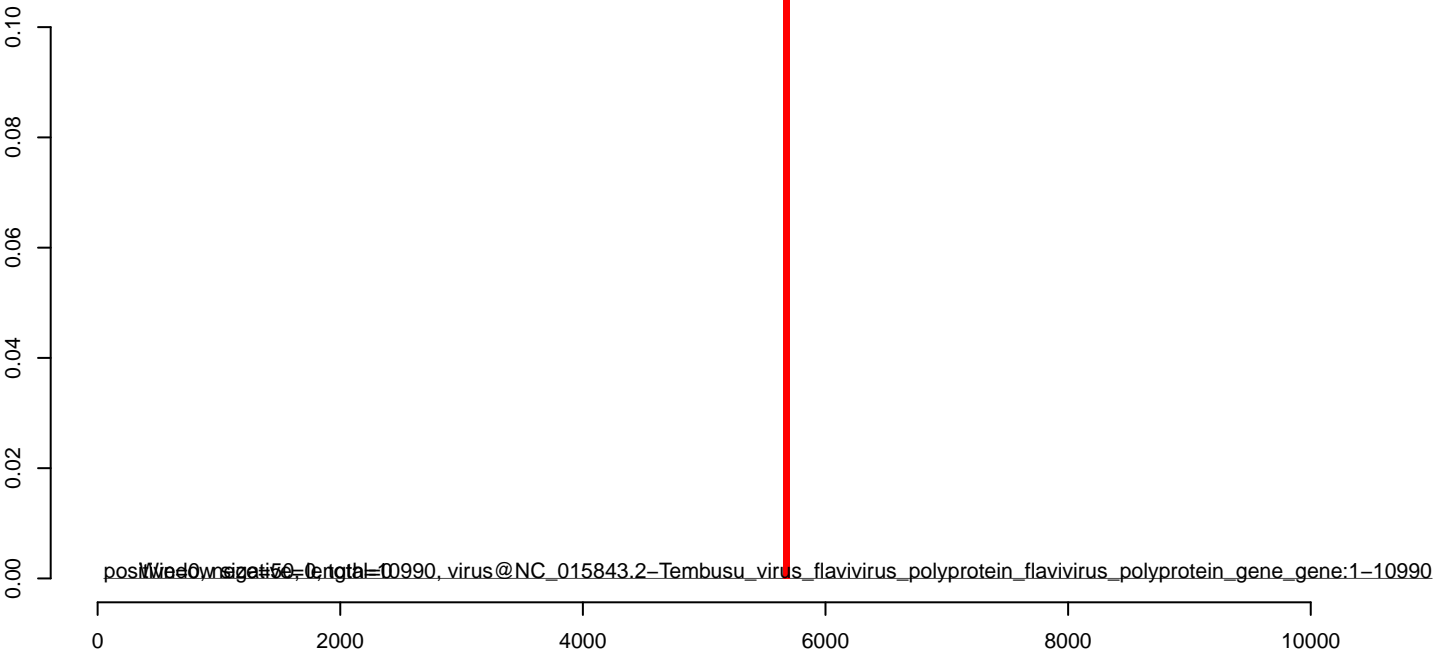
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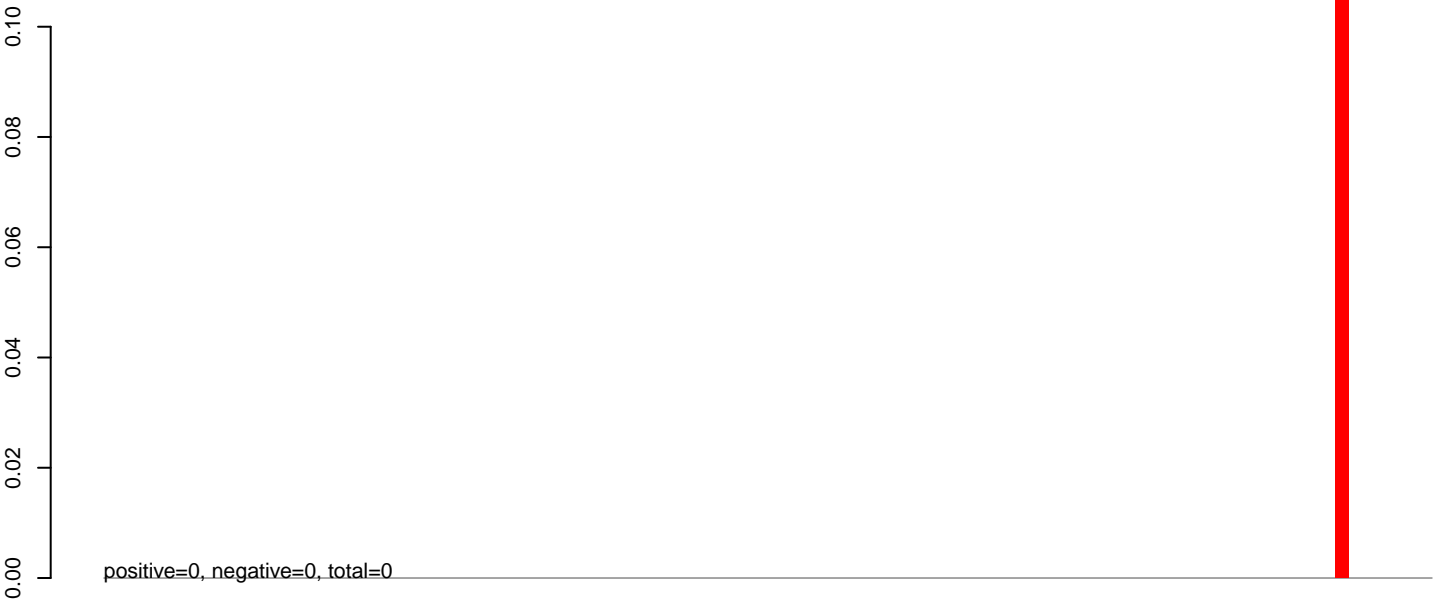
AnGam\_Sua5b\_cells.24\_35.rep



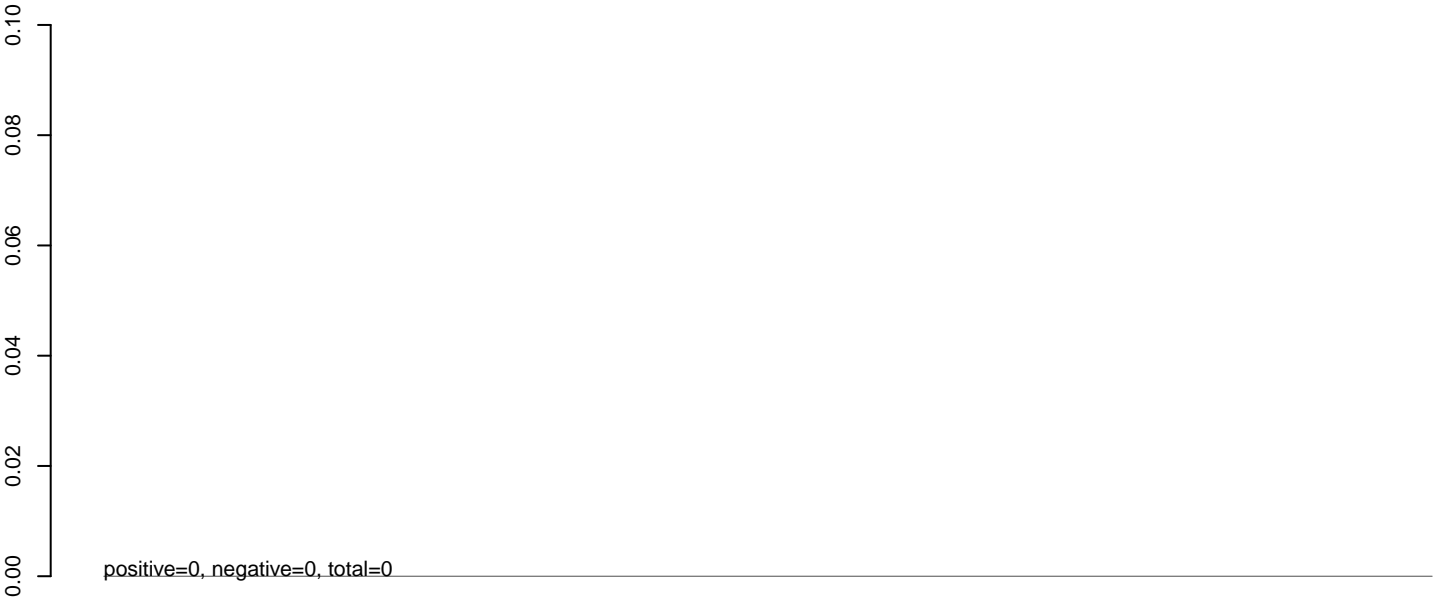
AnGam\_Sua5b\_cells.rep



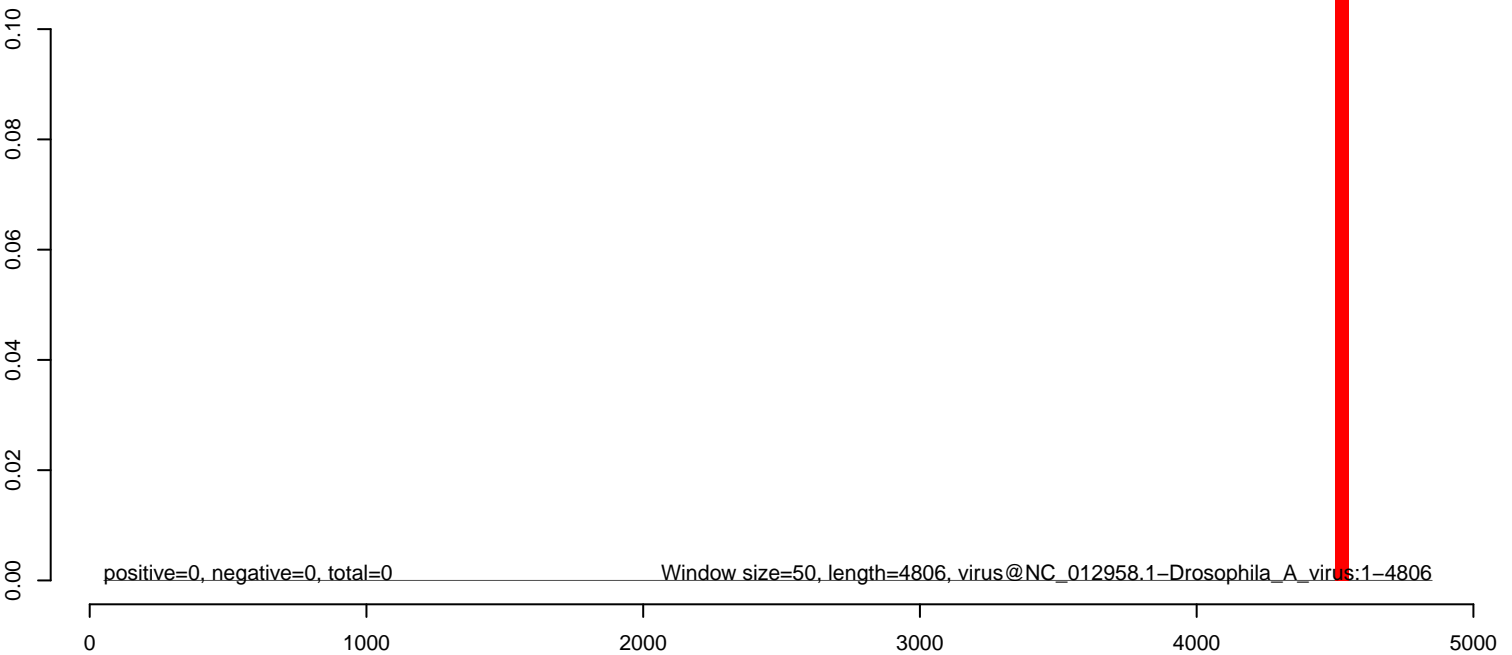
AnGam\_Sua5b\_cells.18\_23.rep



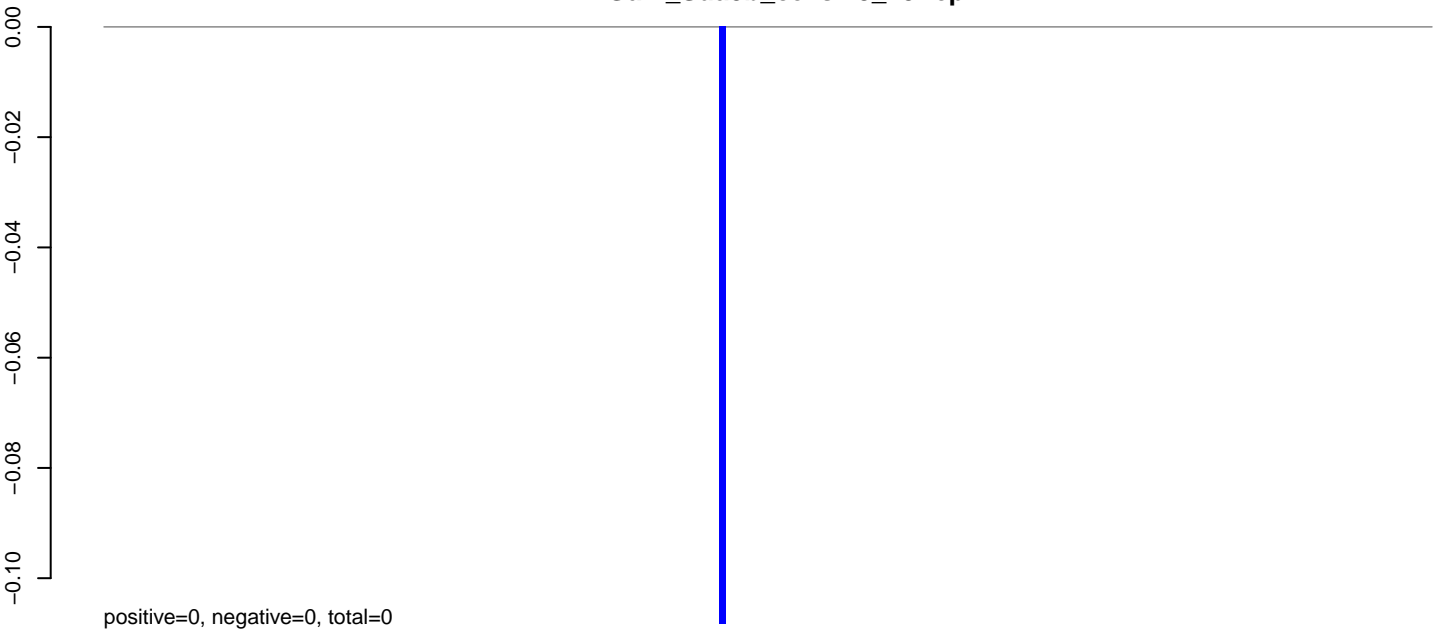
AnGam\_Sua5b\_cells.24\_35.rep



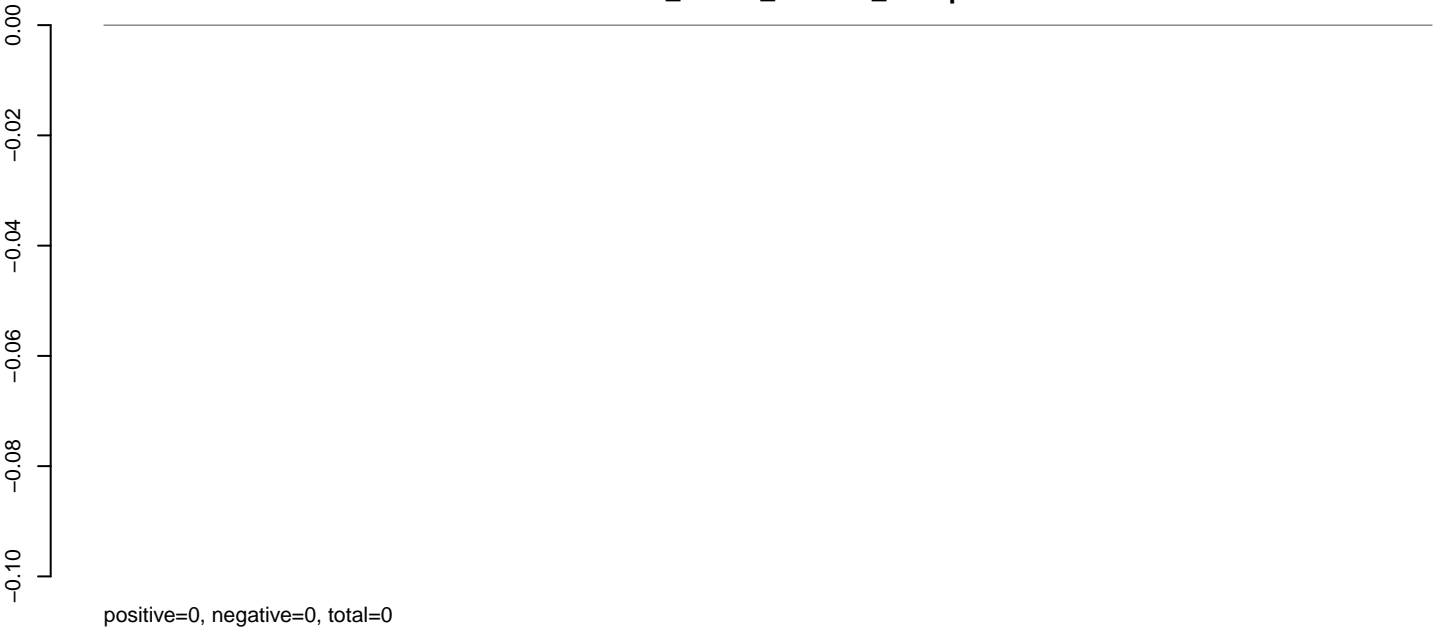
AnGam\_Sua5b\_cells.rep



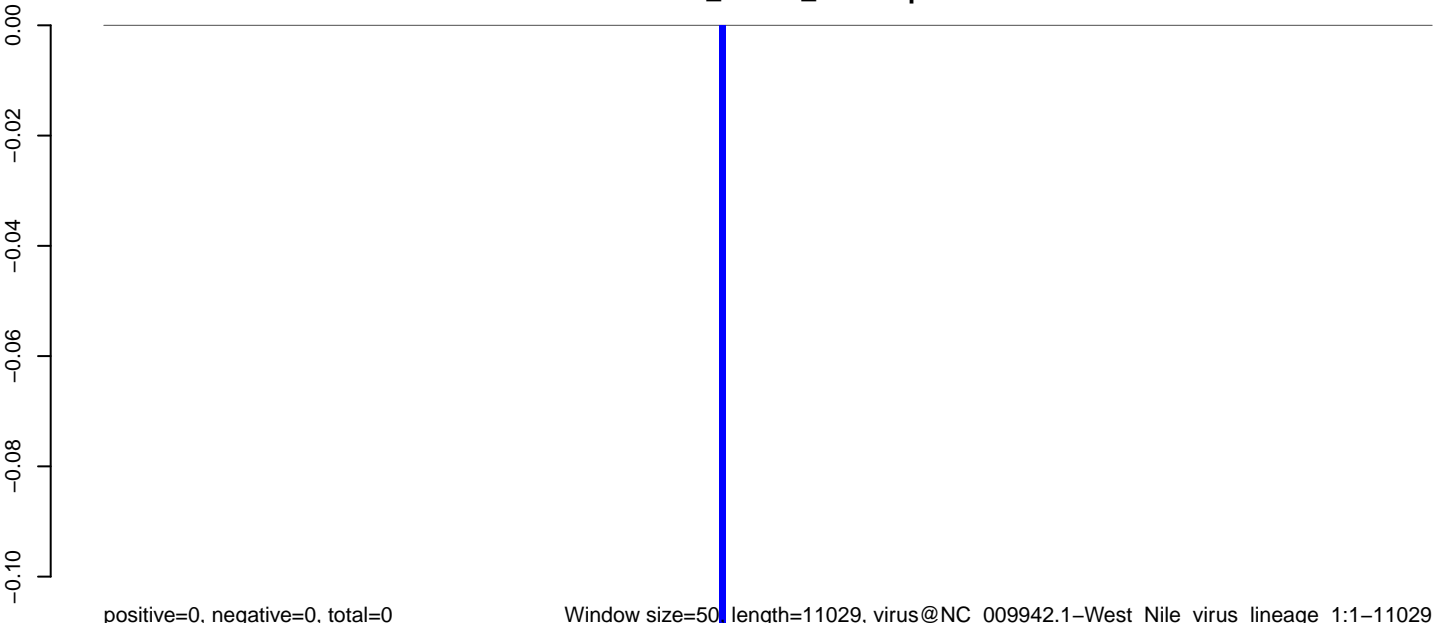
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep

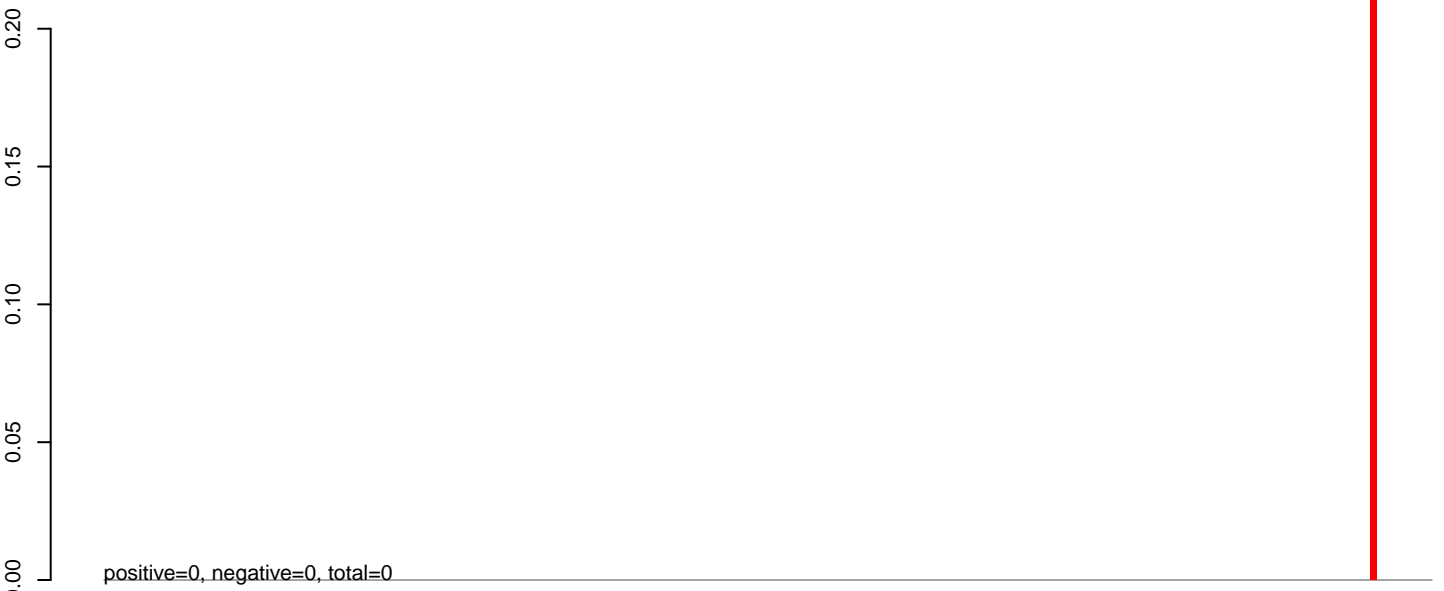


AnGam\_Sua5b\_cells.rep

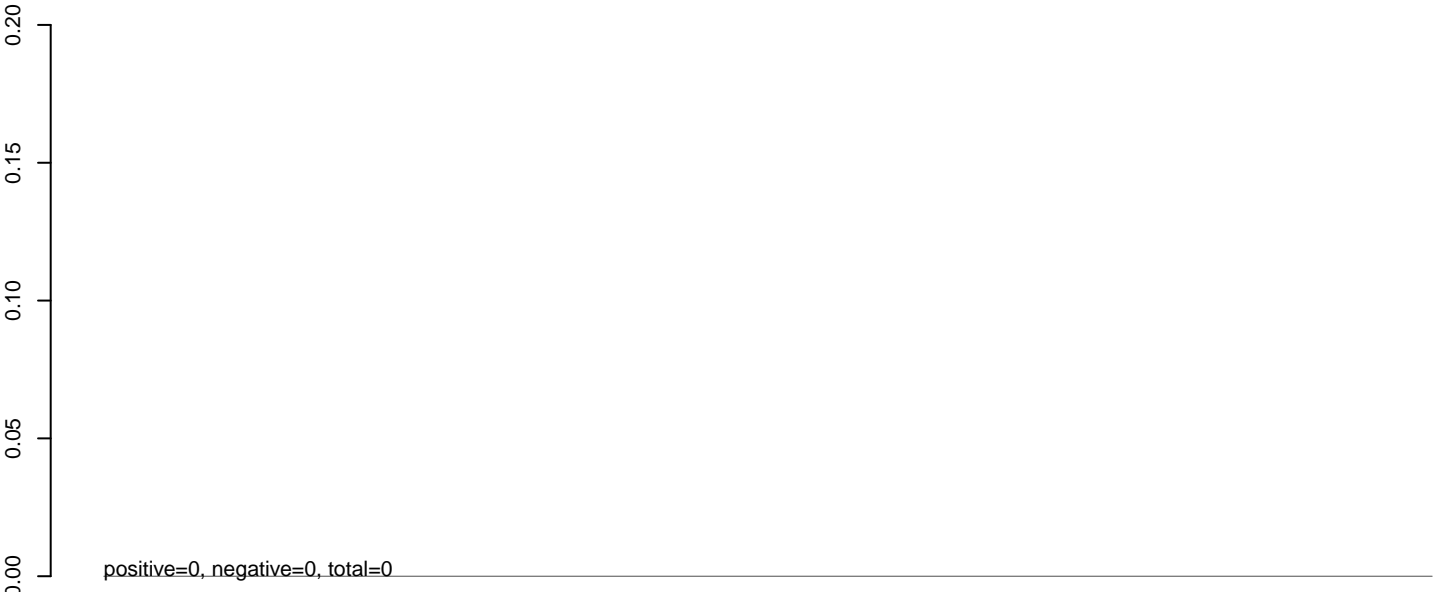


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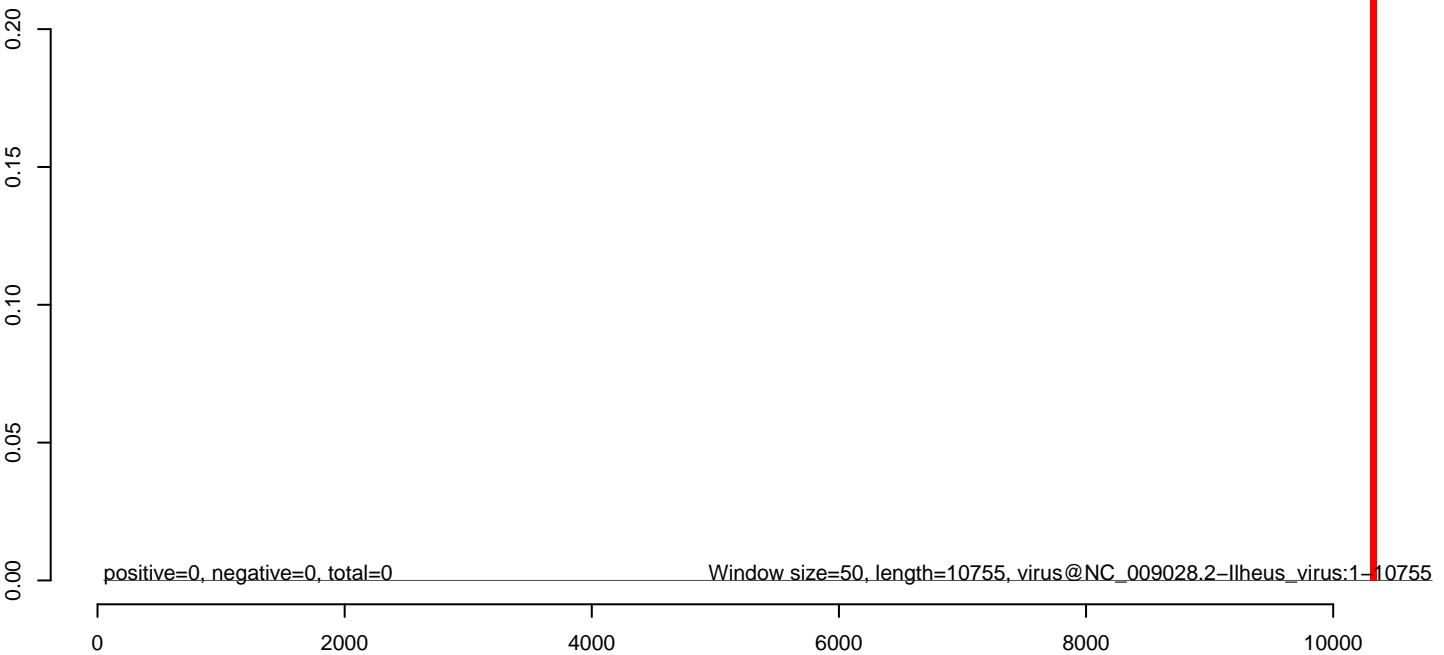
AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep

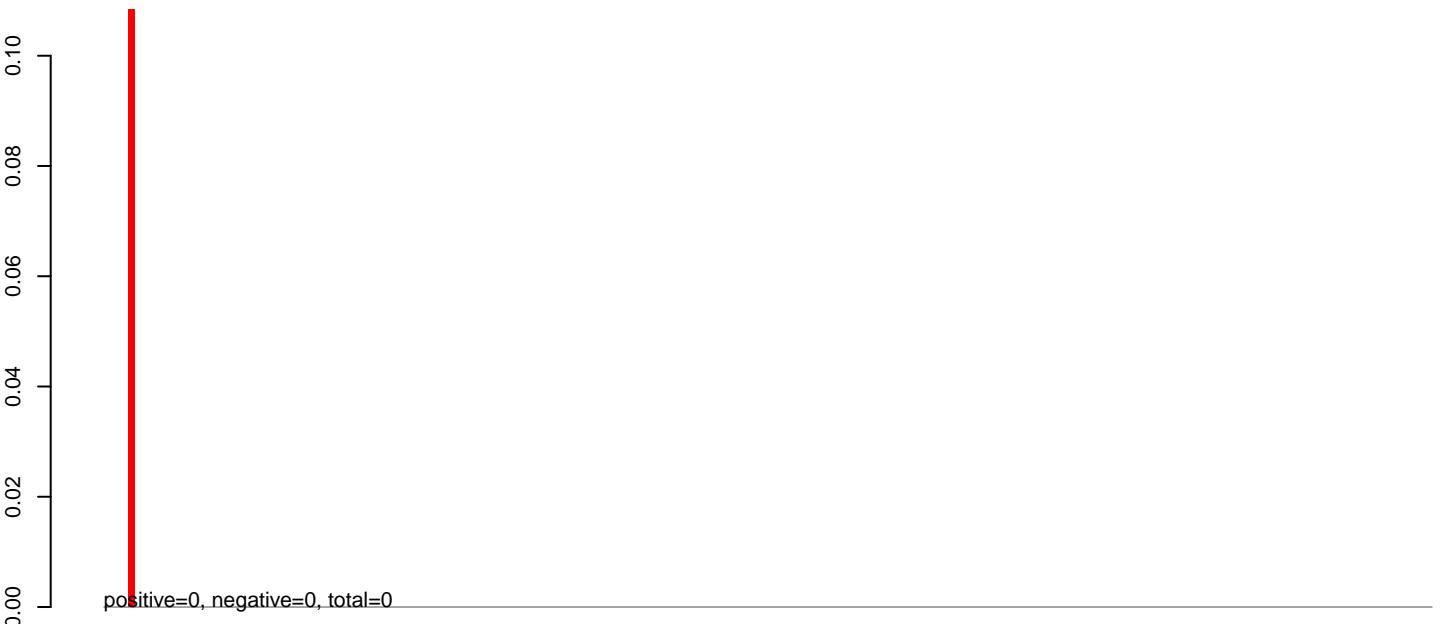


AnGam\_Sua5b\_cells.rep

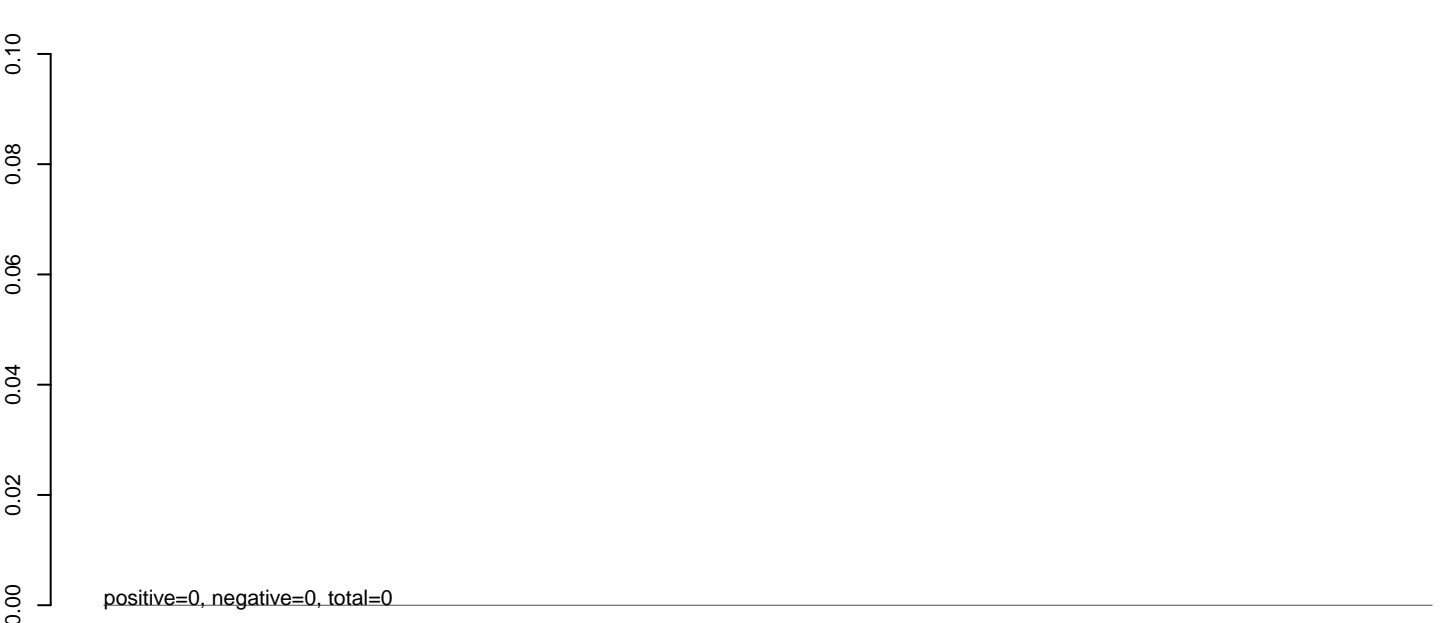




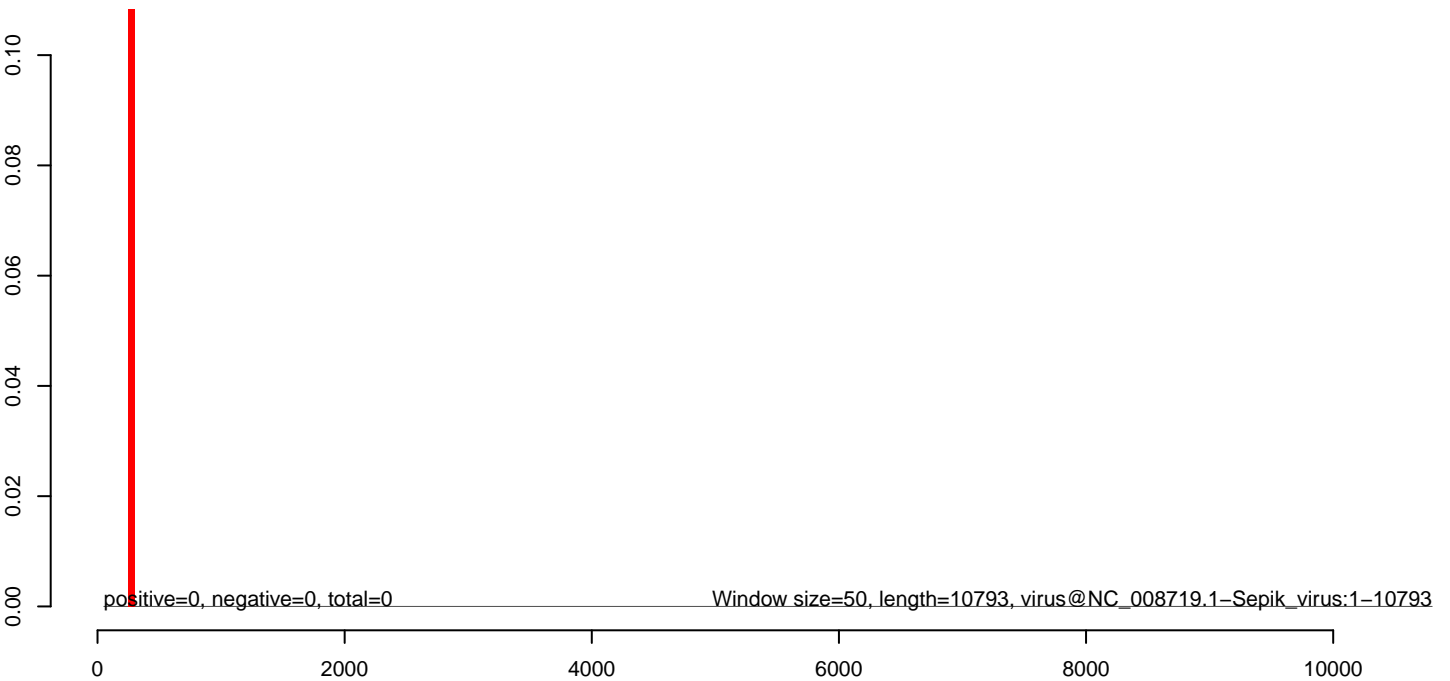
### AnGam\_Sua5b\_cells.18\_23.rep



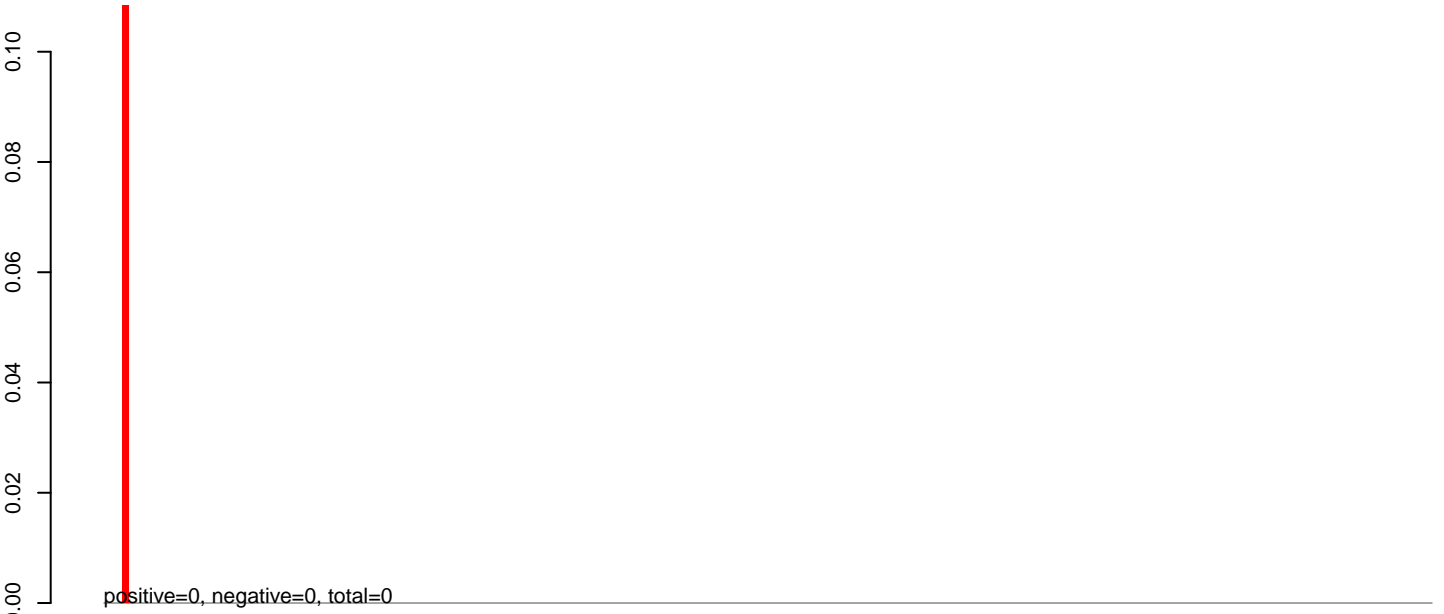
### AnGam\_Sua5b\_cells.24\_35.rep



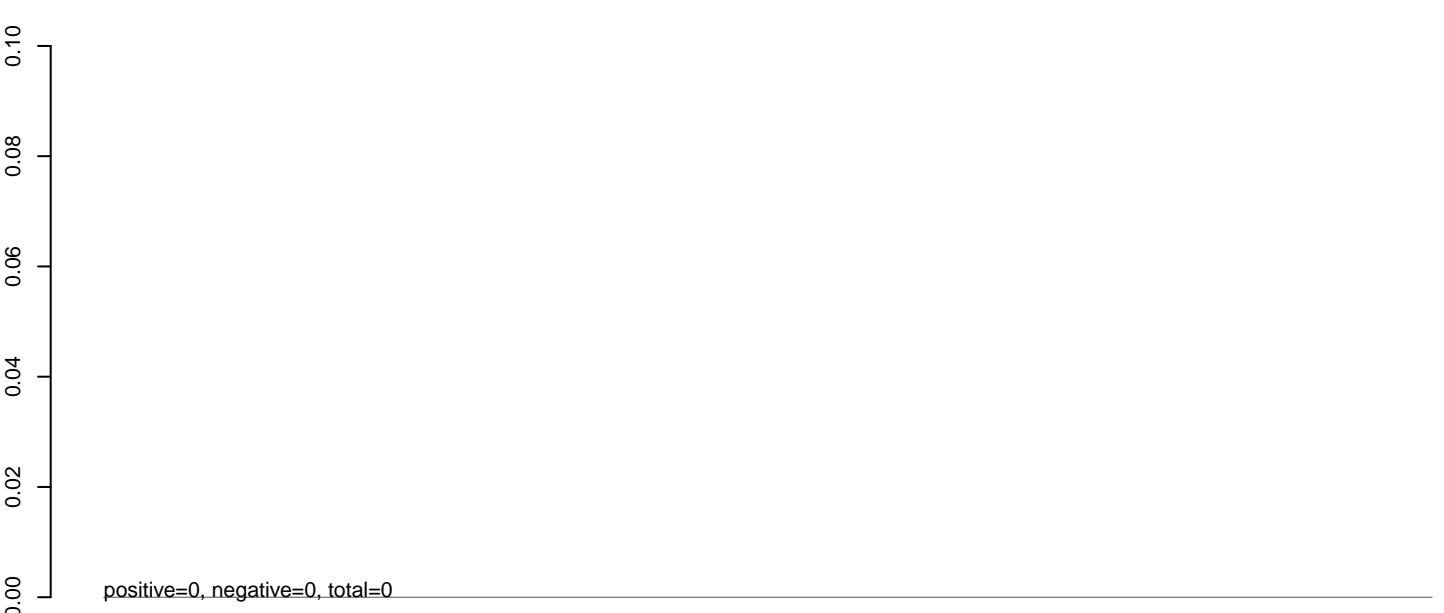
### AnGam\_Sua5b\_cells.rep



### AnGam\_Sua5b\_cells.18\_23.rep



### AnGam\_Sua5b\_cells.24\_35.rep

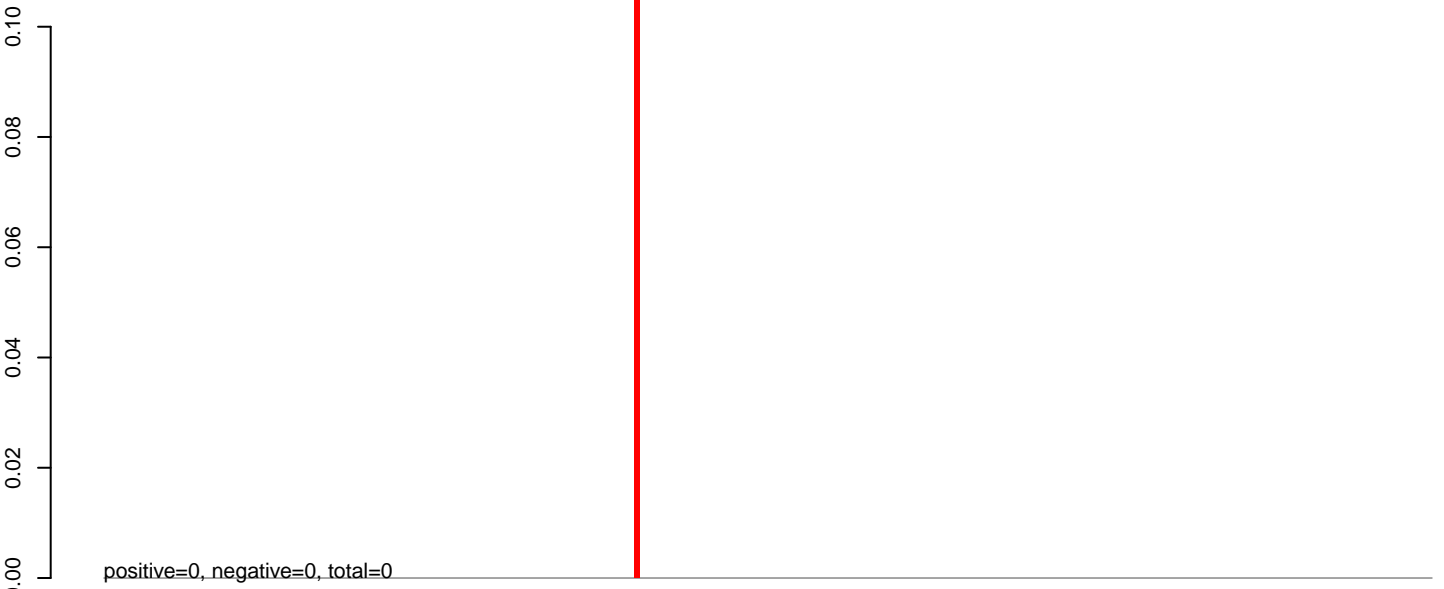


### AnGam\_Sua5b\_cells.rep

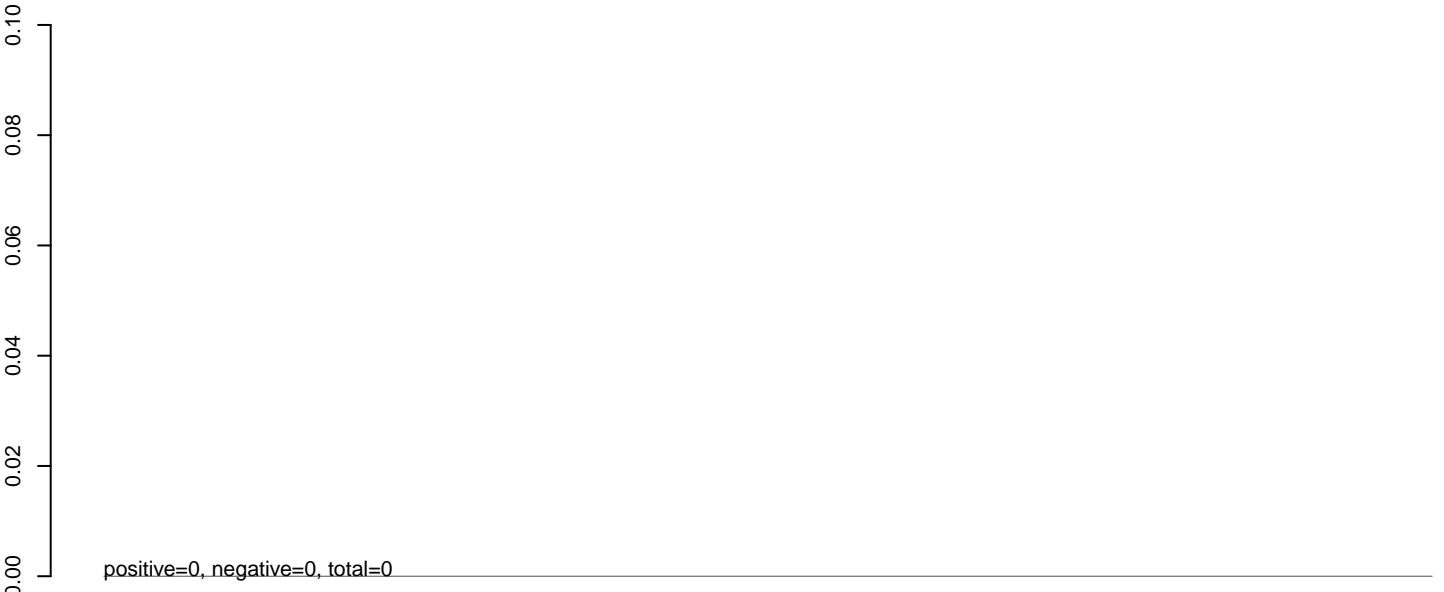


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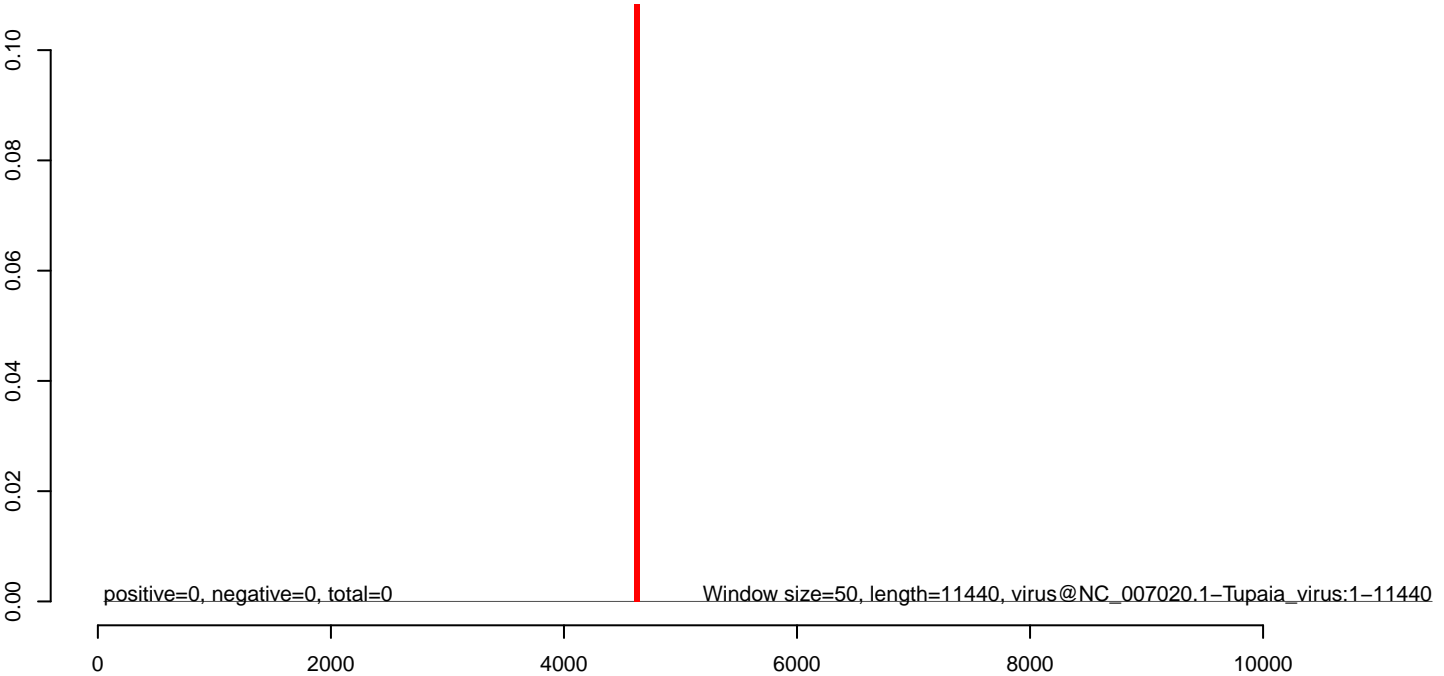
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AnGam\_Sua5b\_cells.24\_35.rep



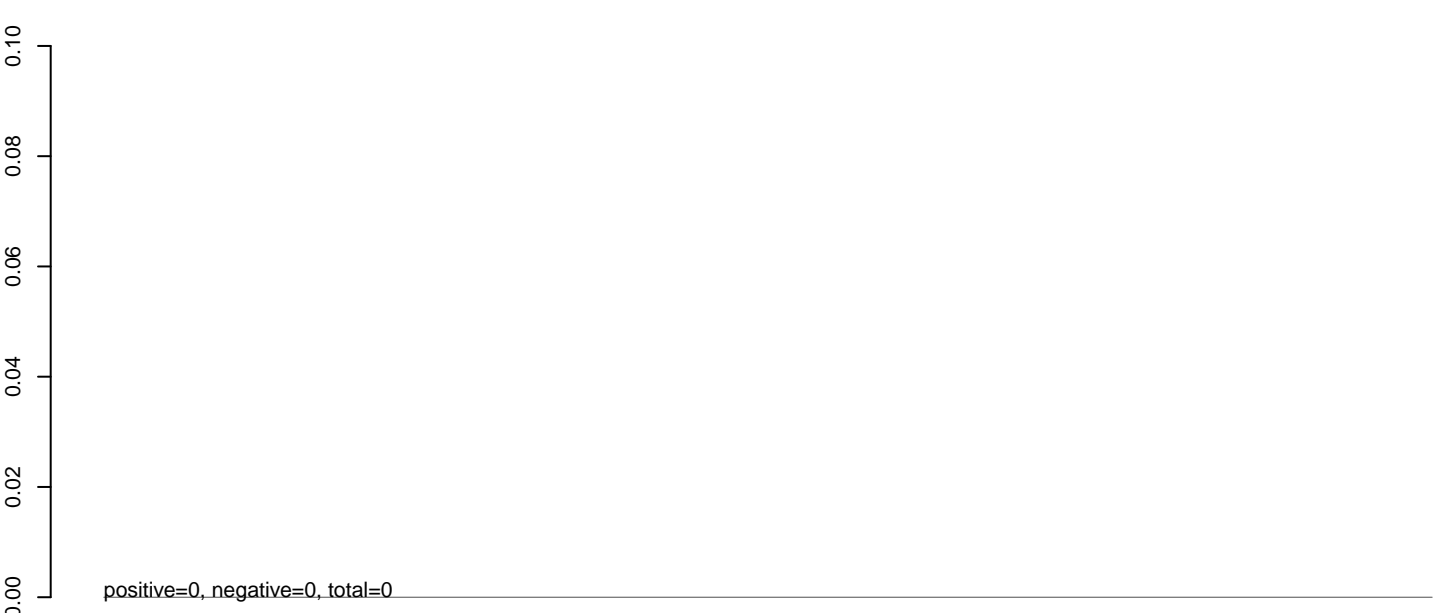
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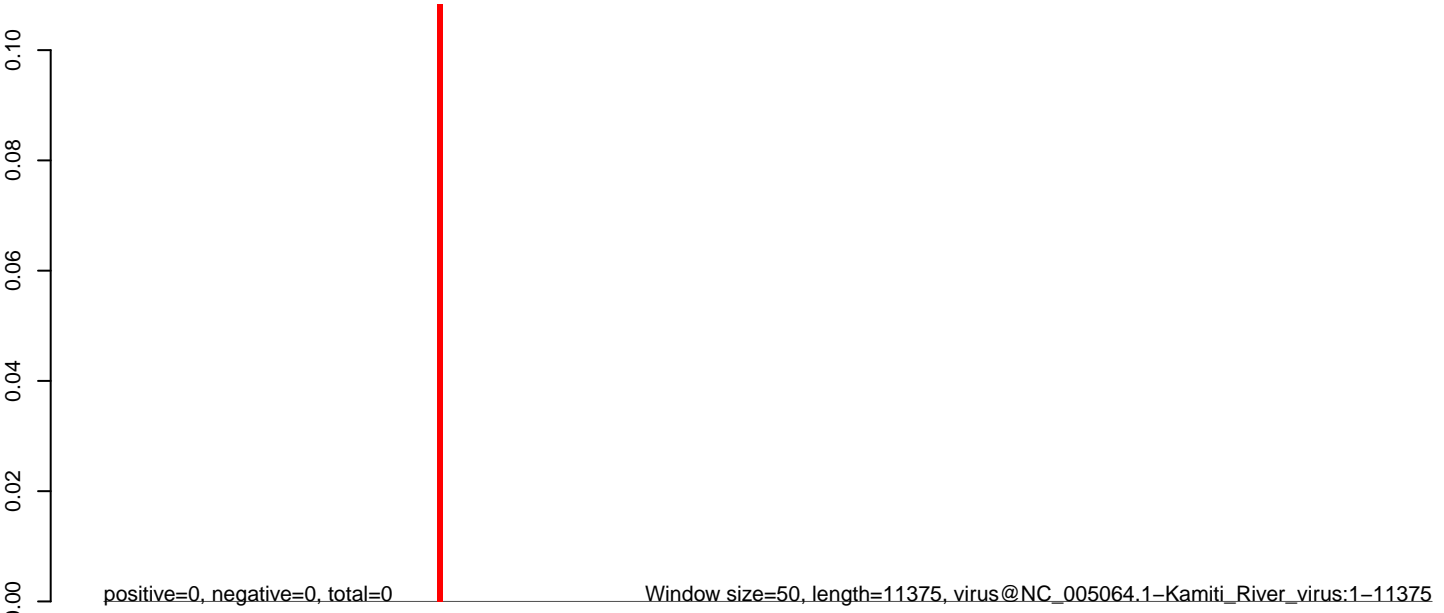
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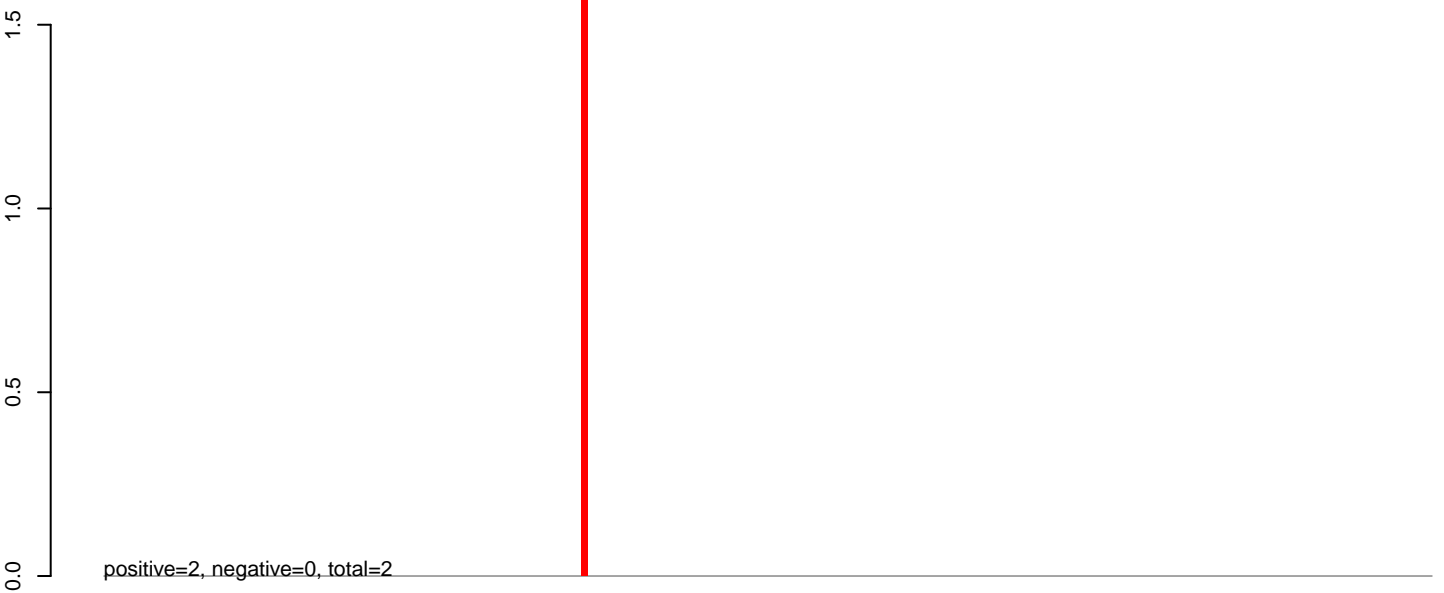
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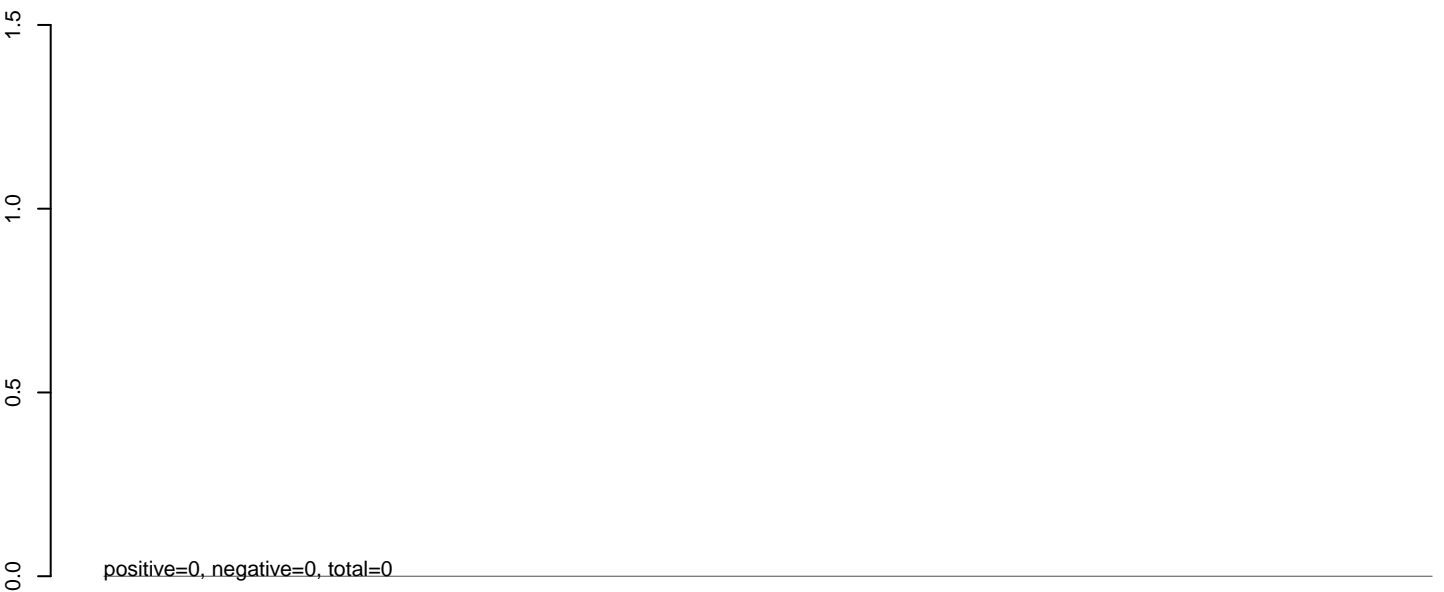
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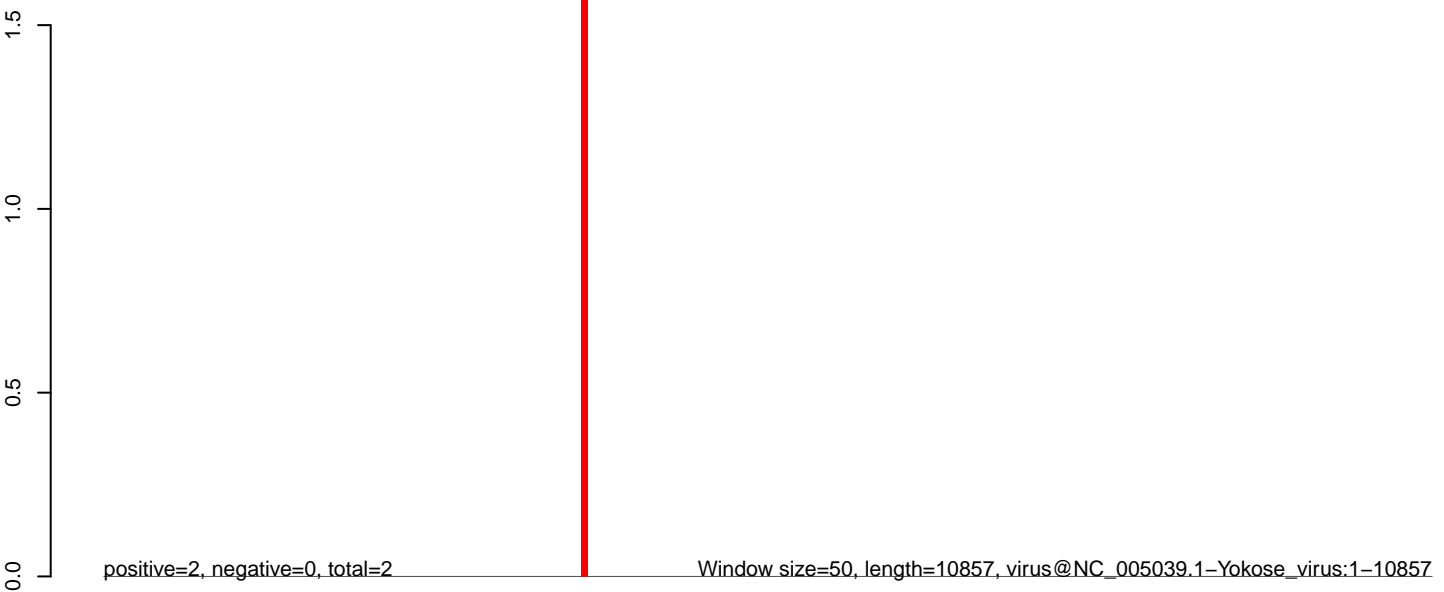
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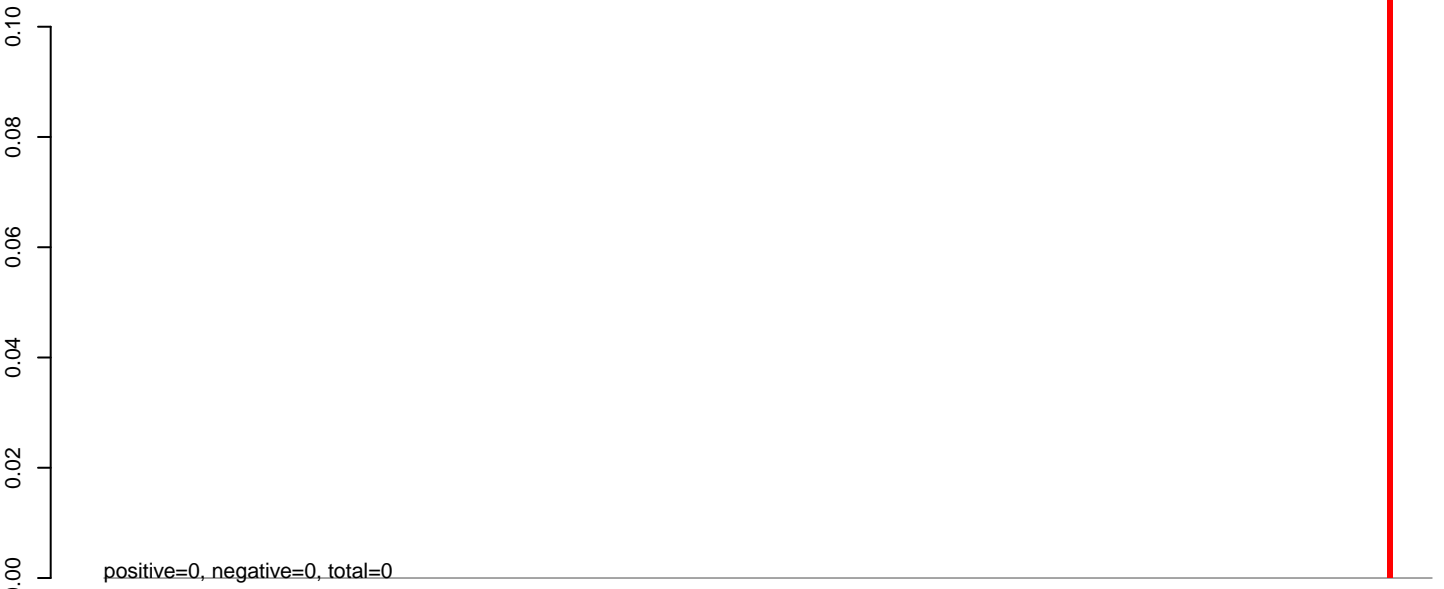
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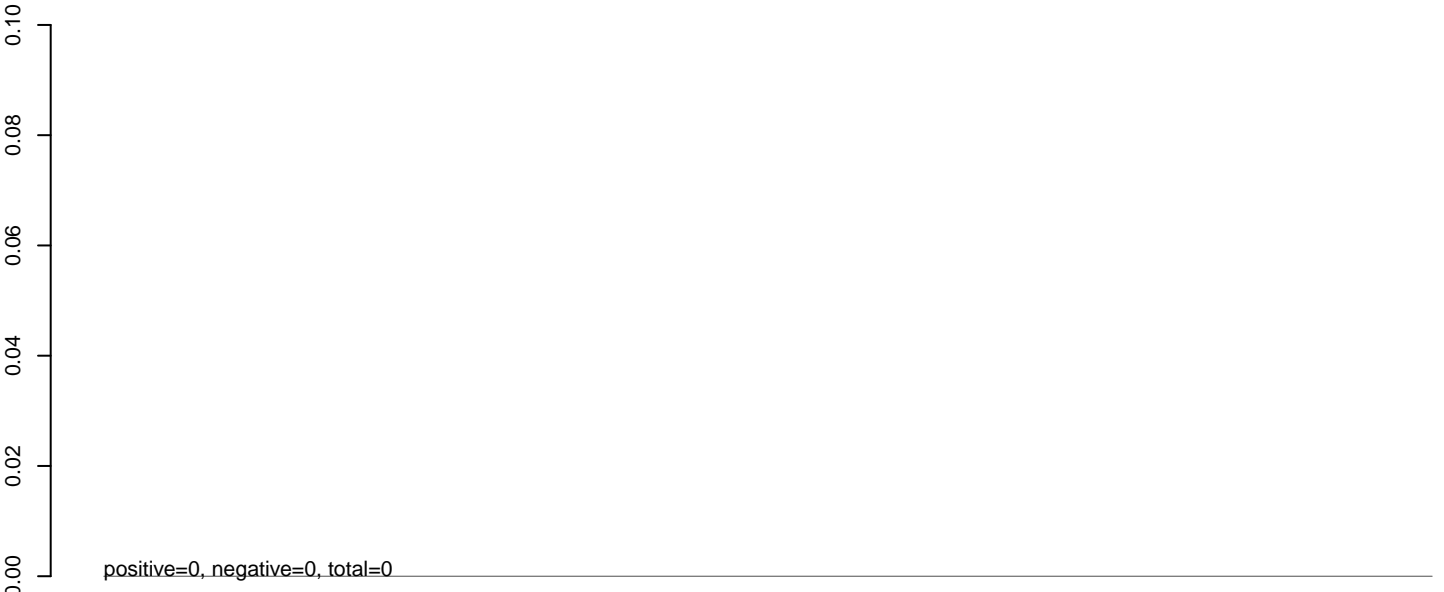
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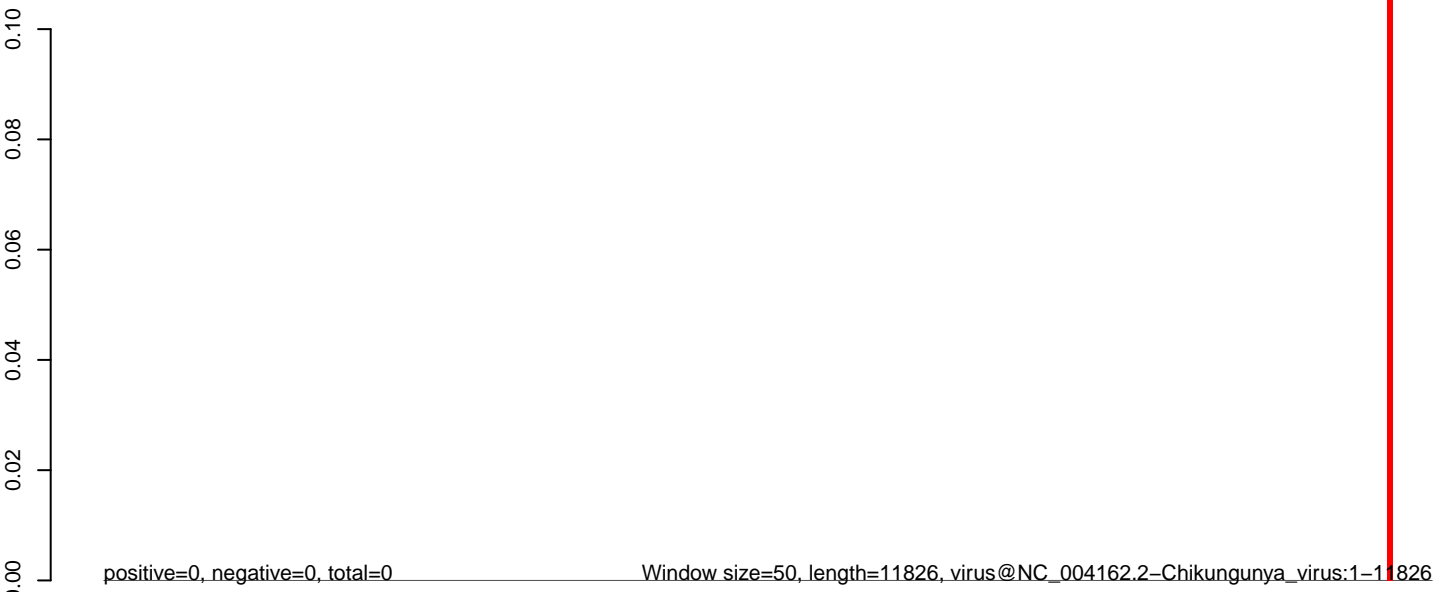
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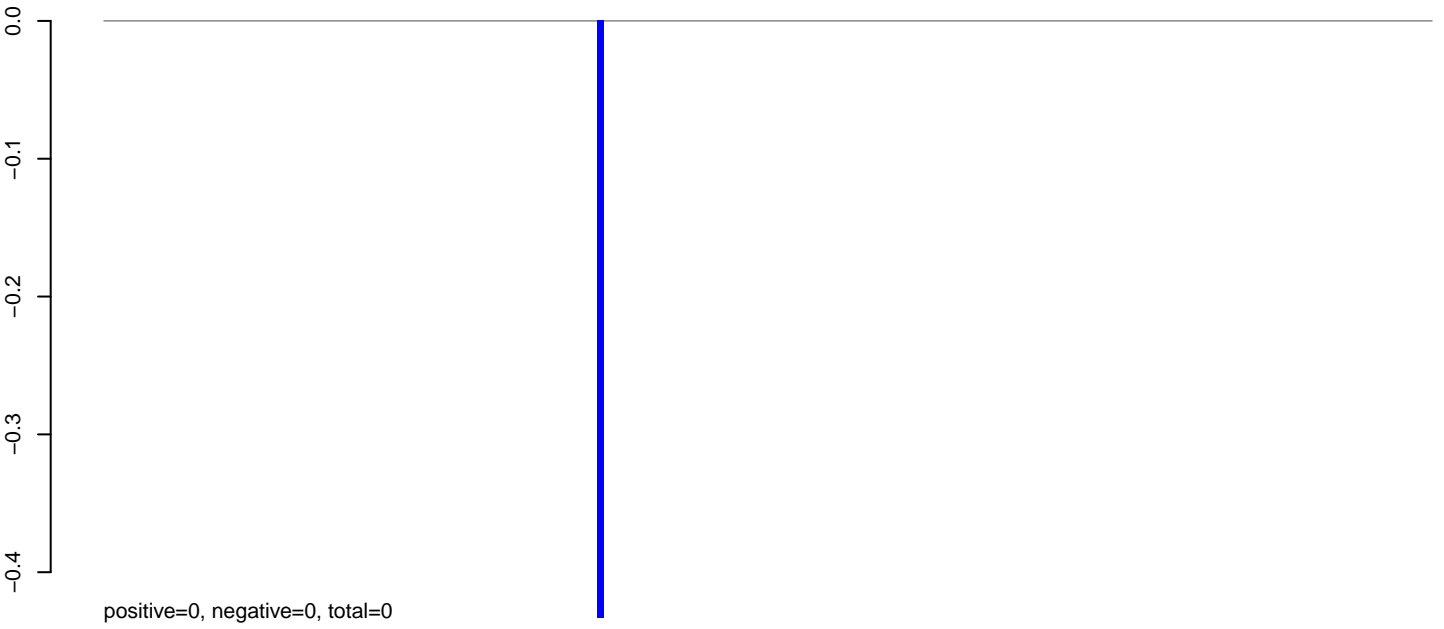


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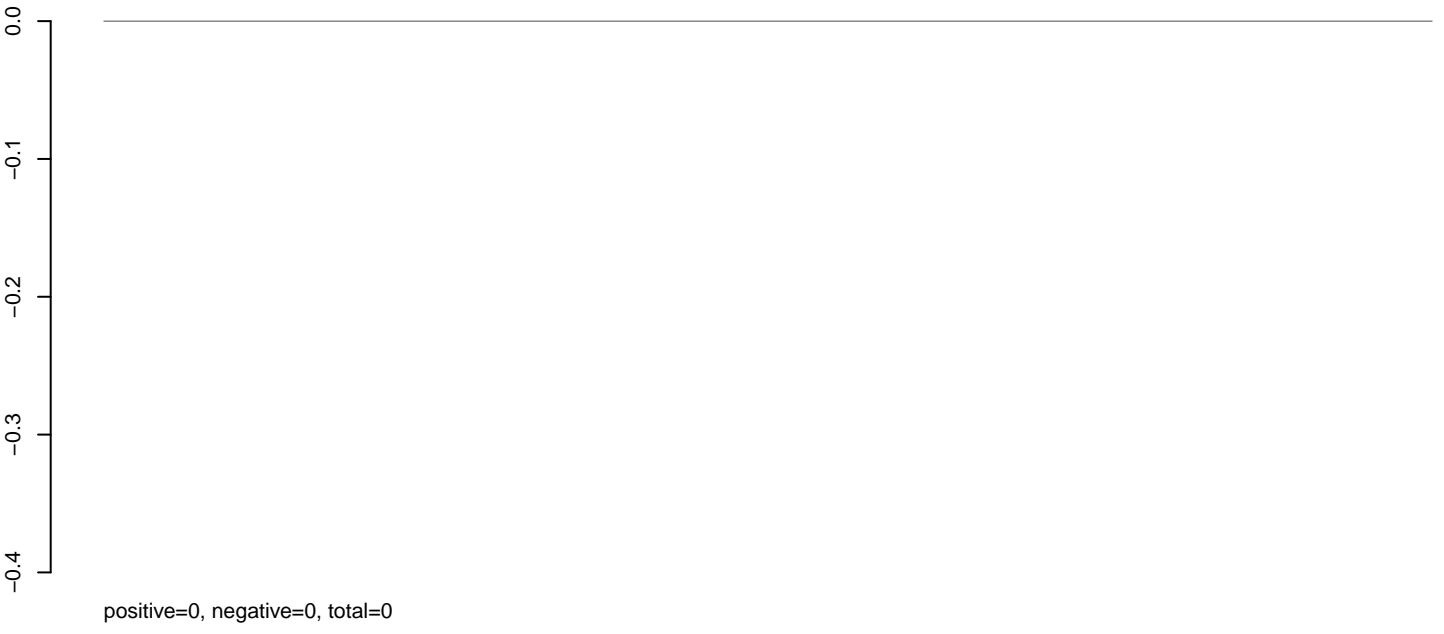


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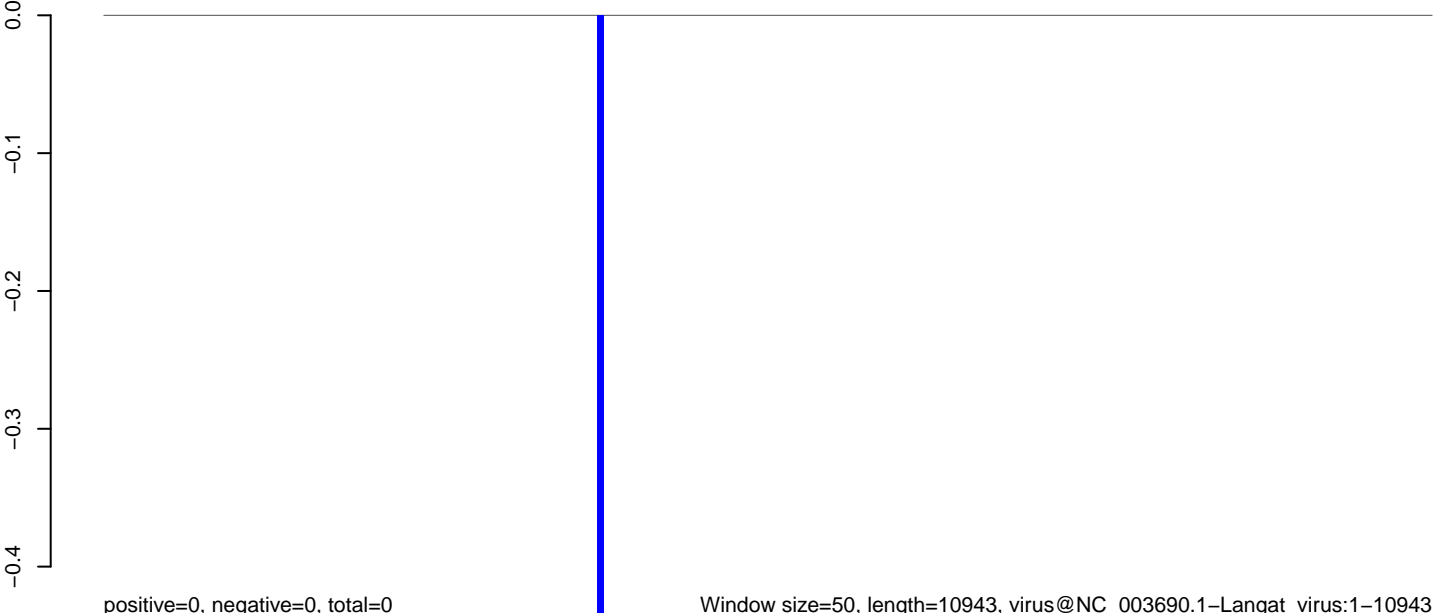
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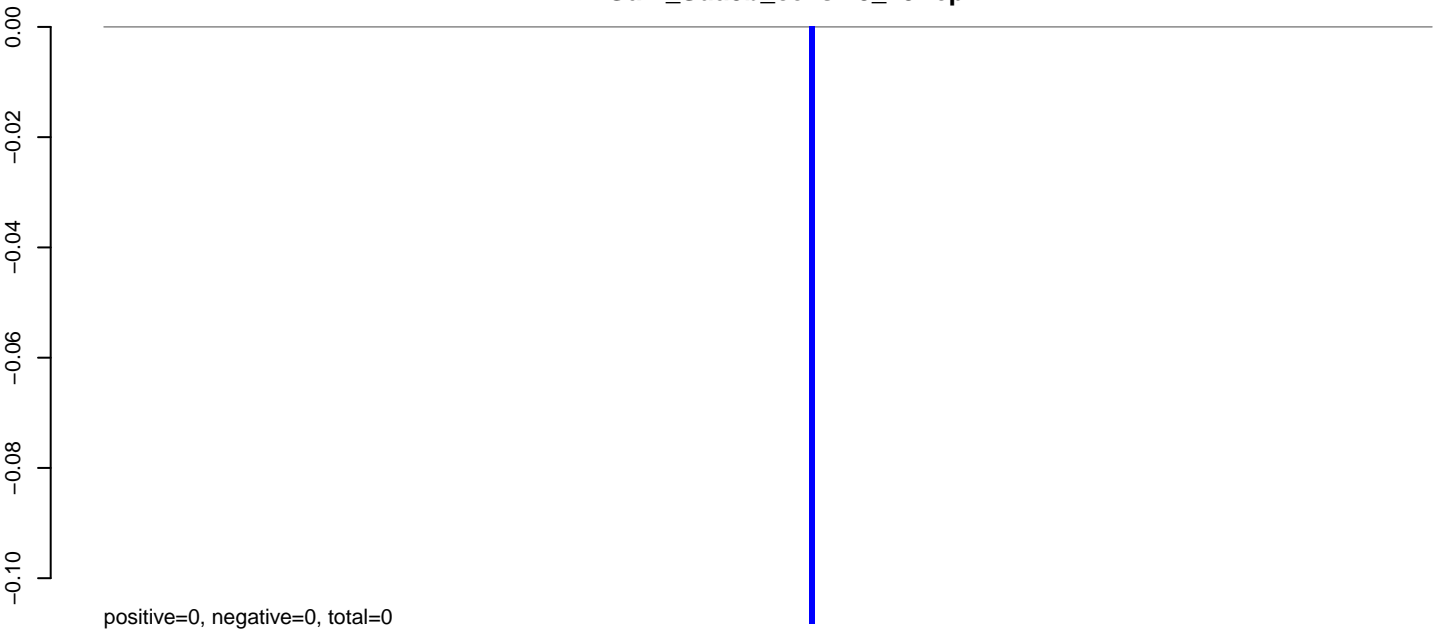
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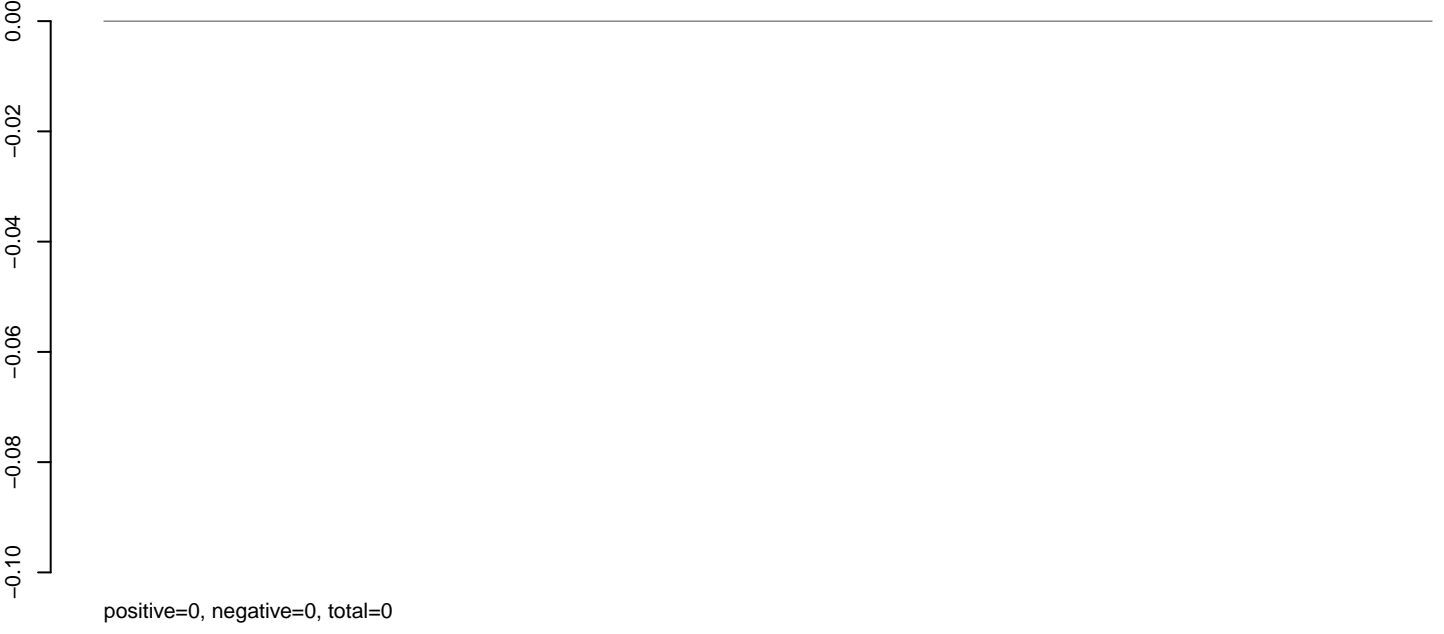
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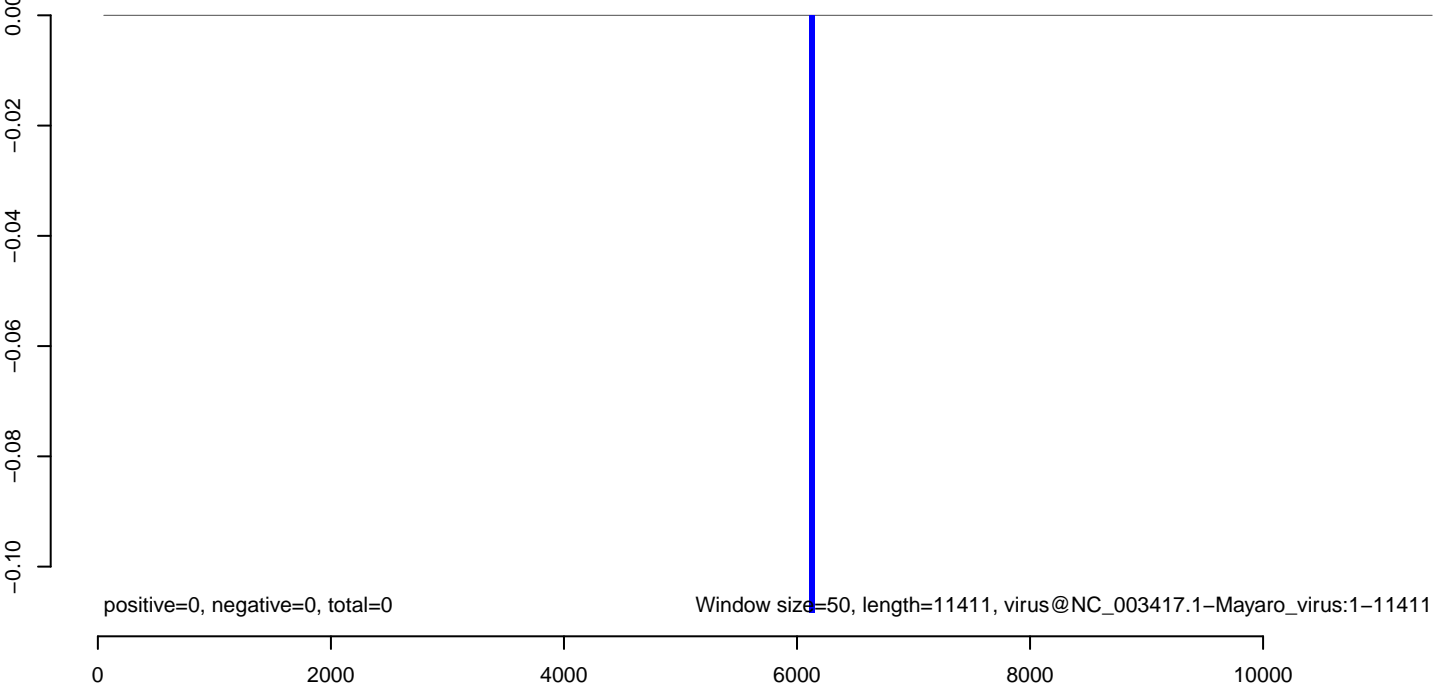
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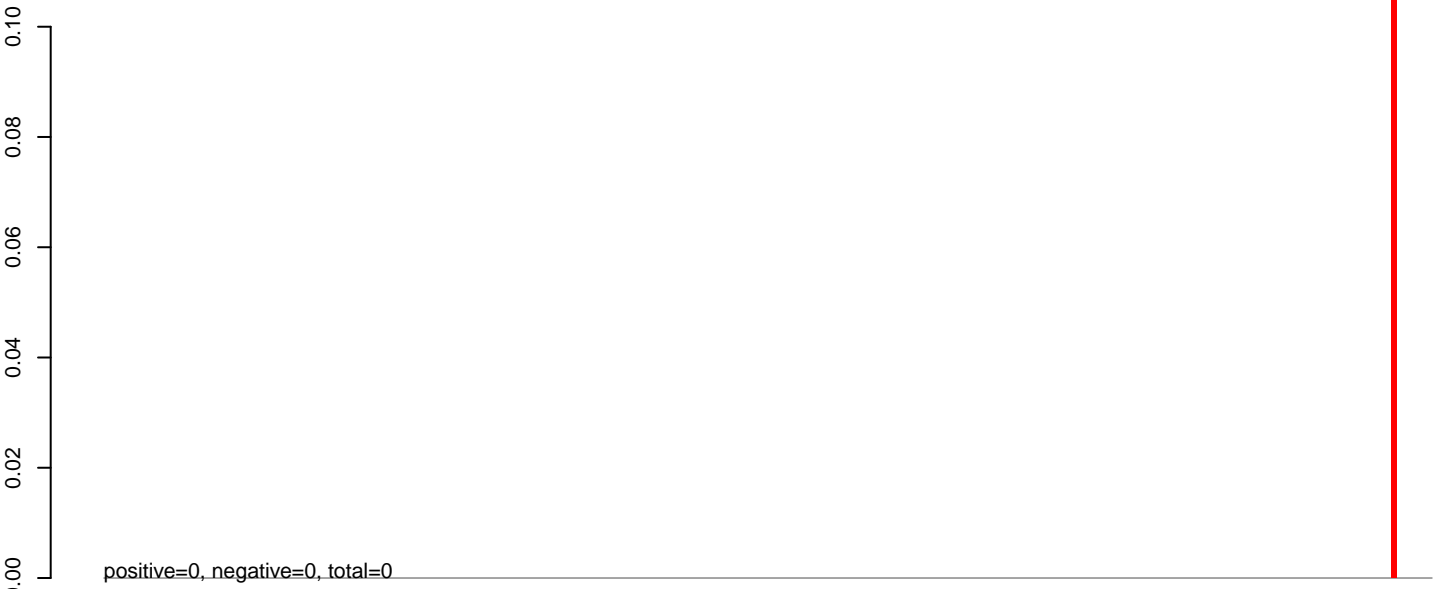


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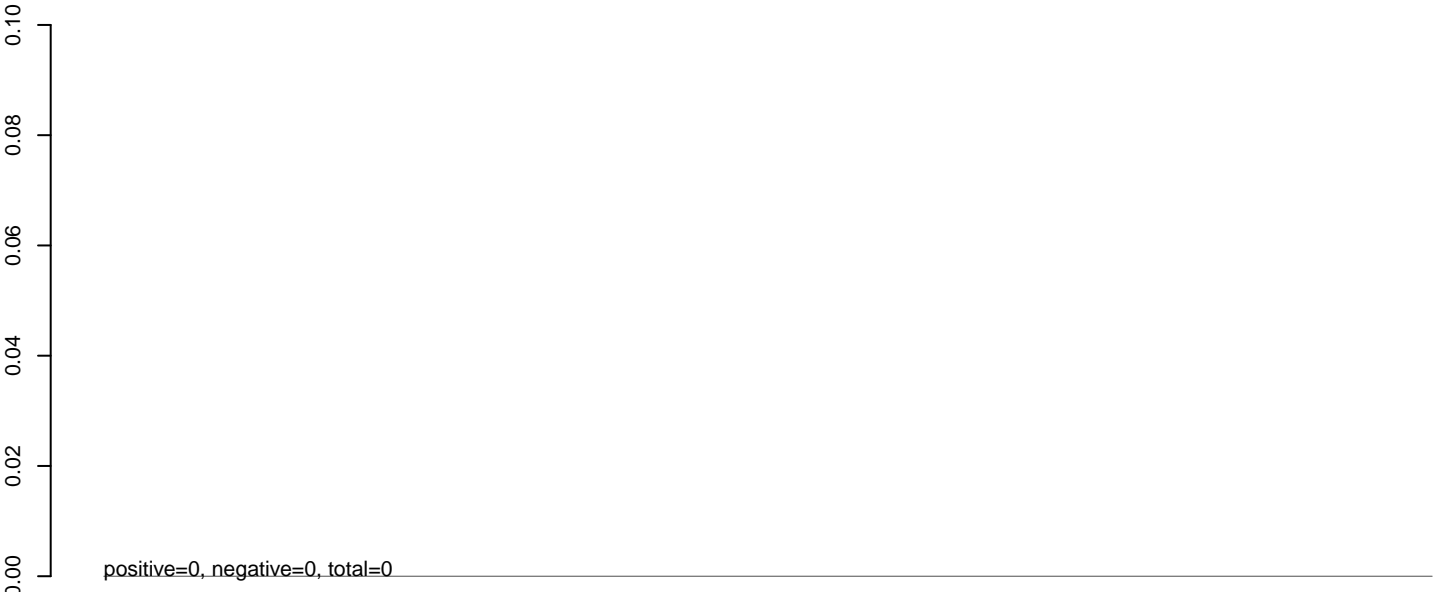




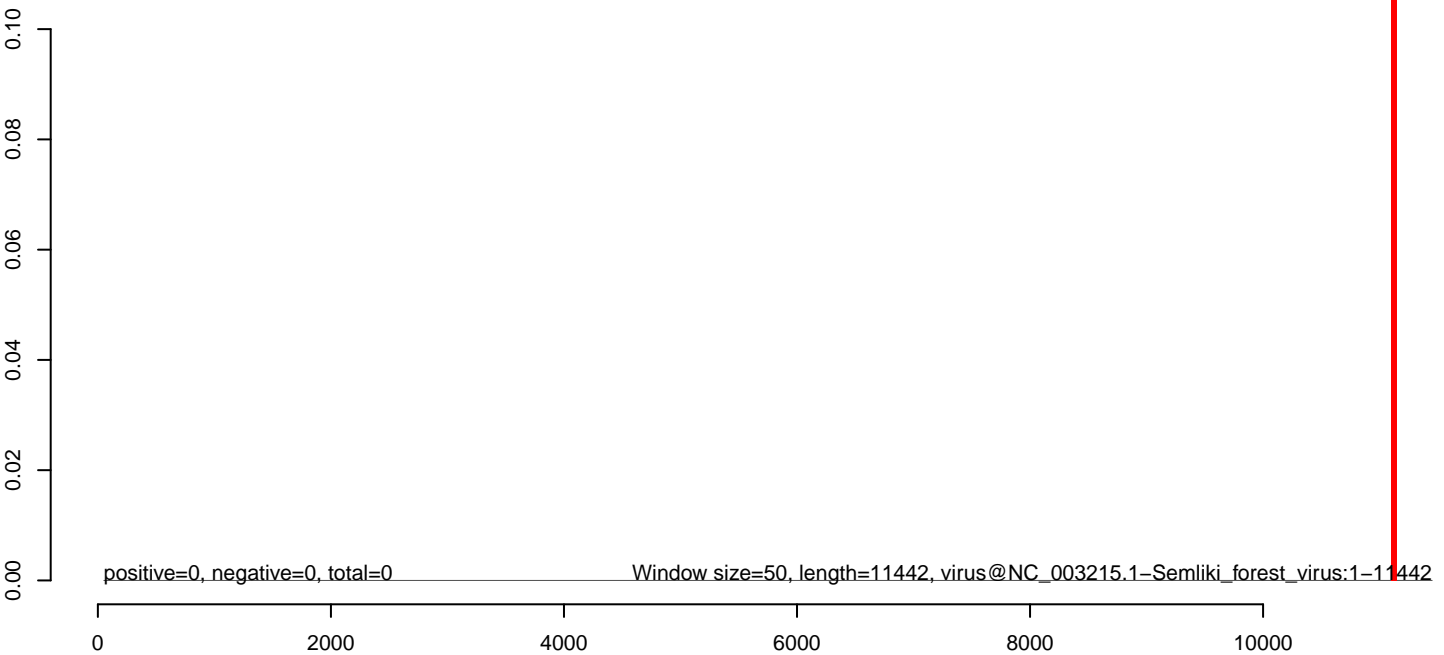
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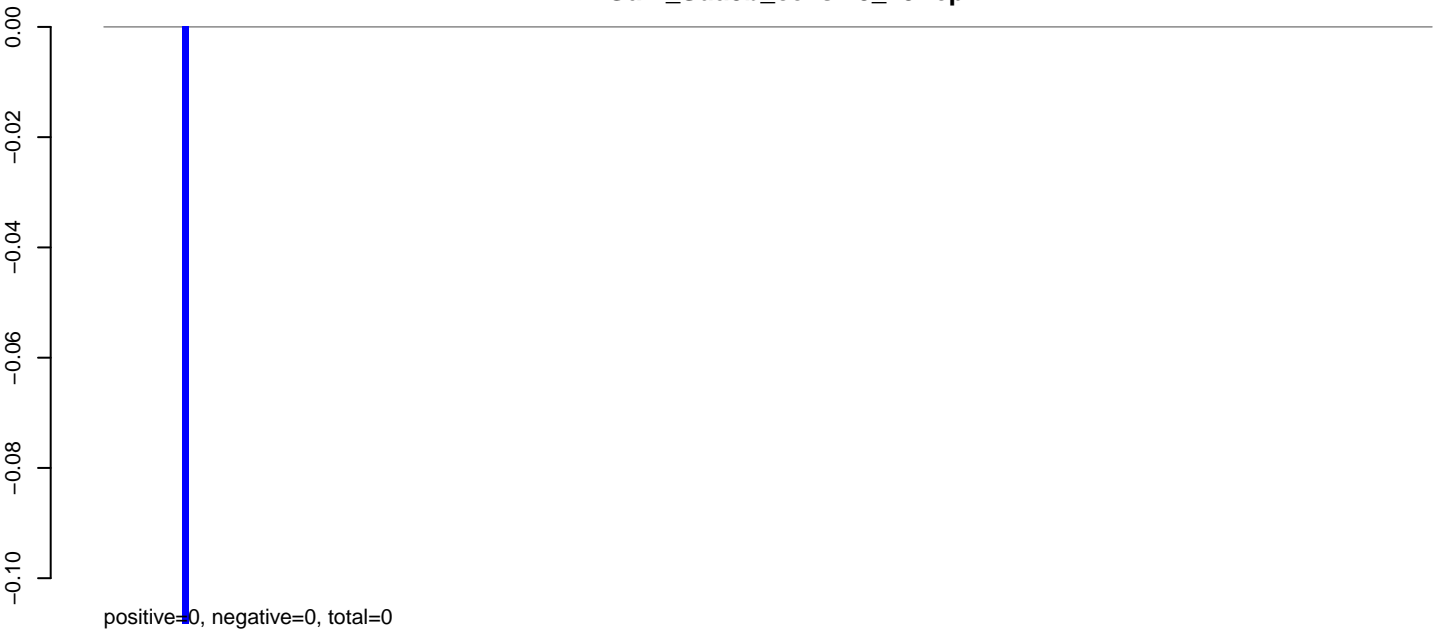
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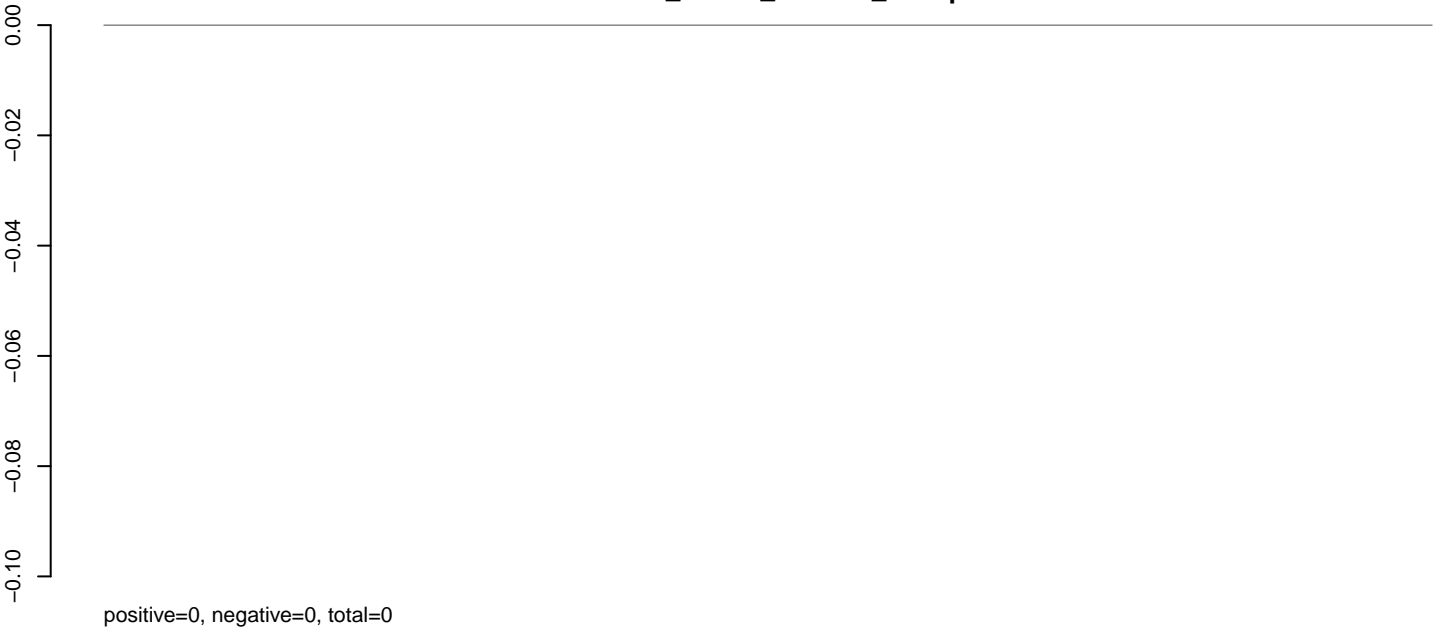
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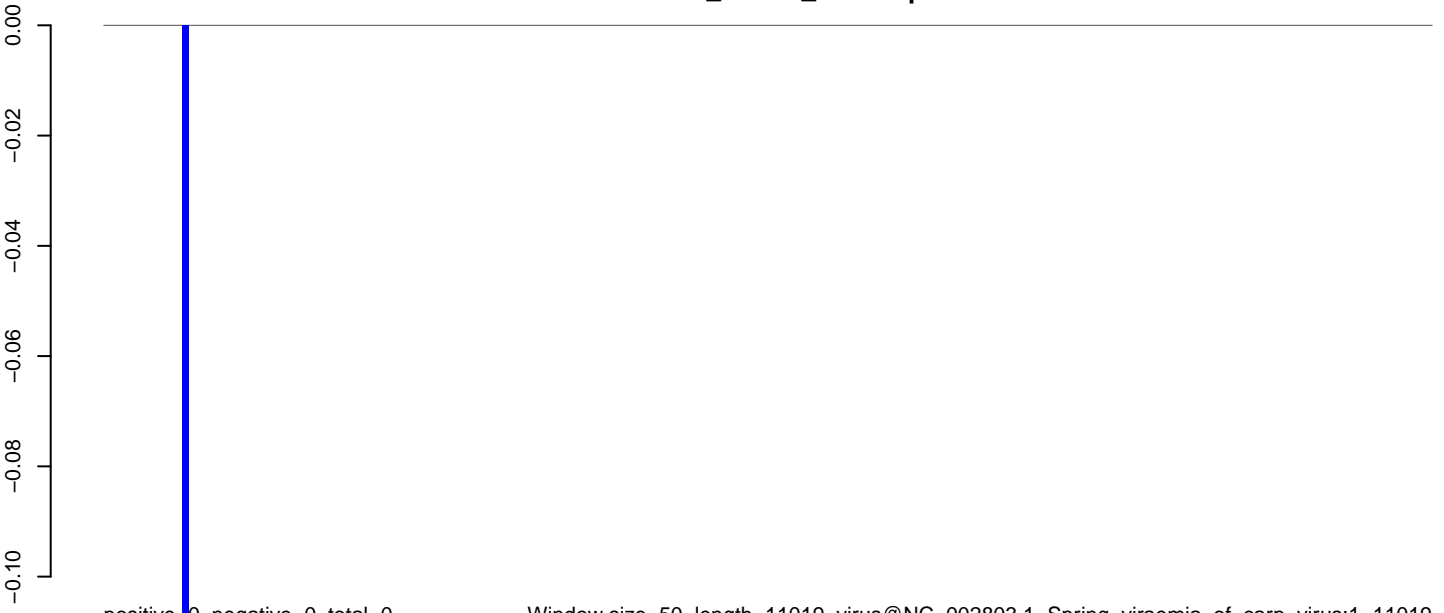
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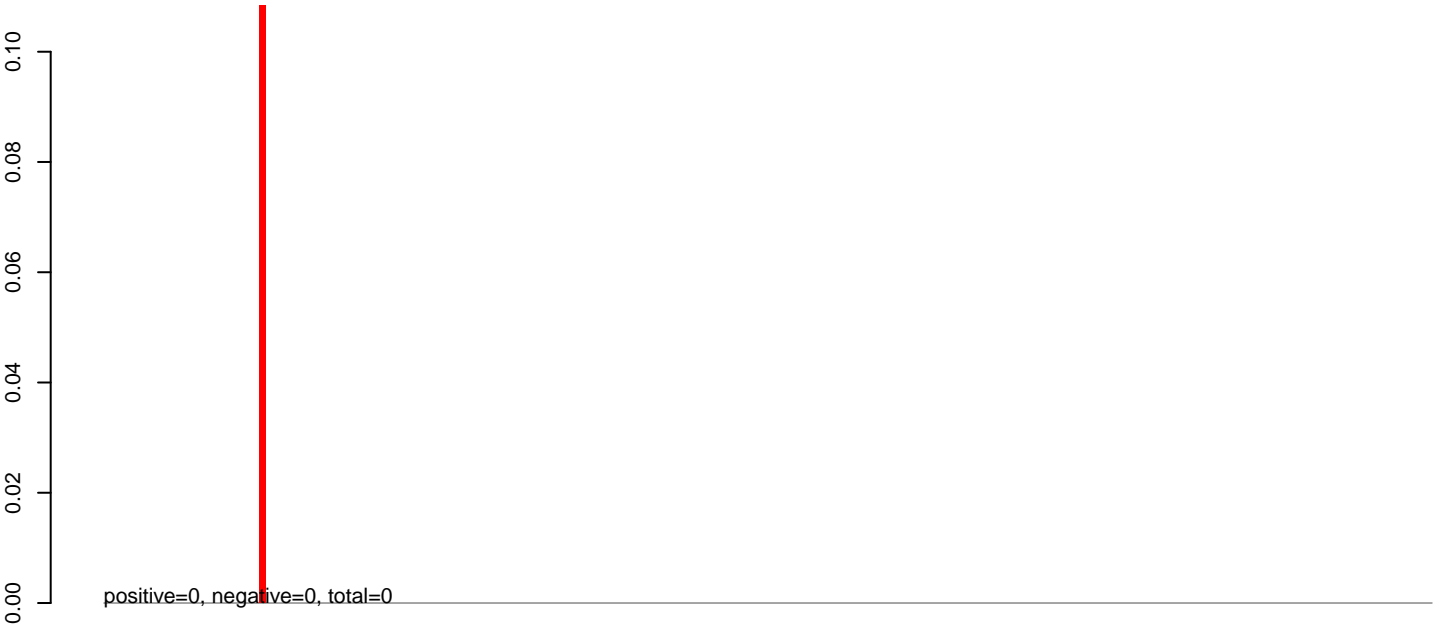
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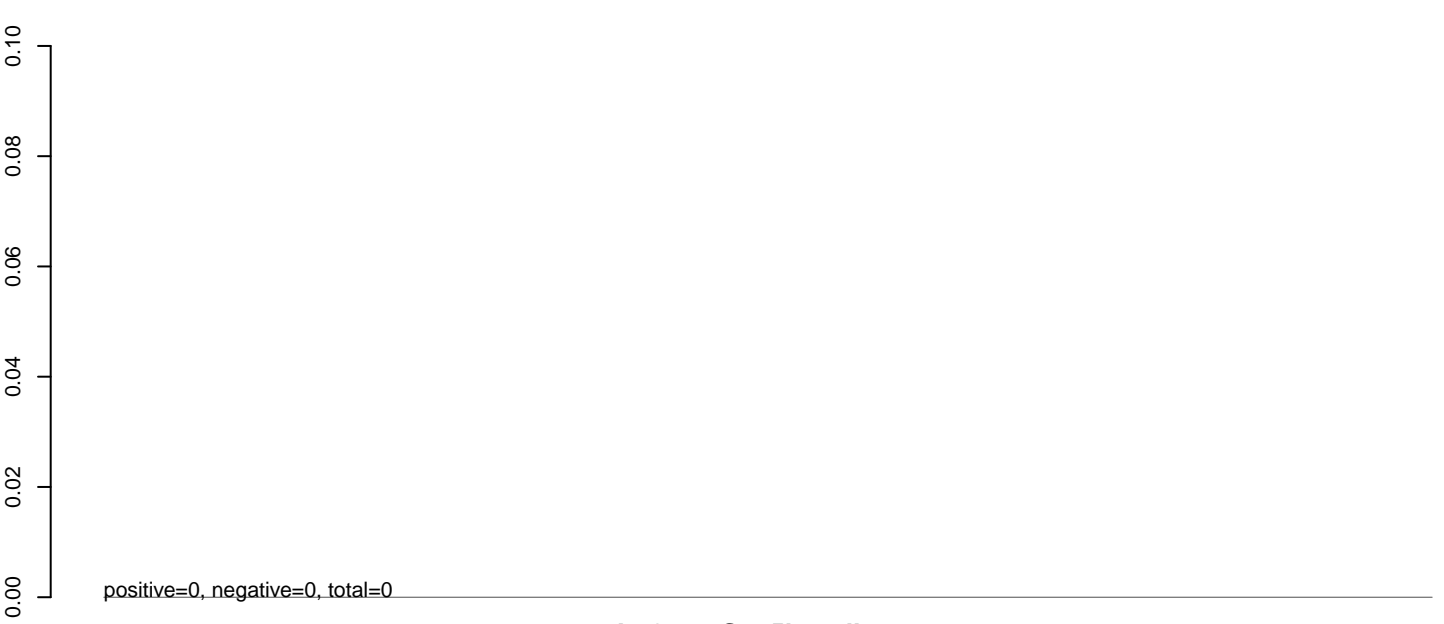
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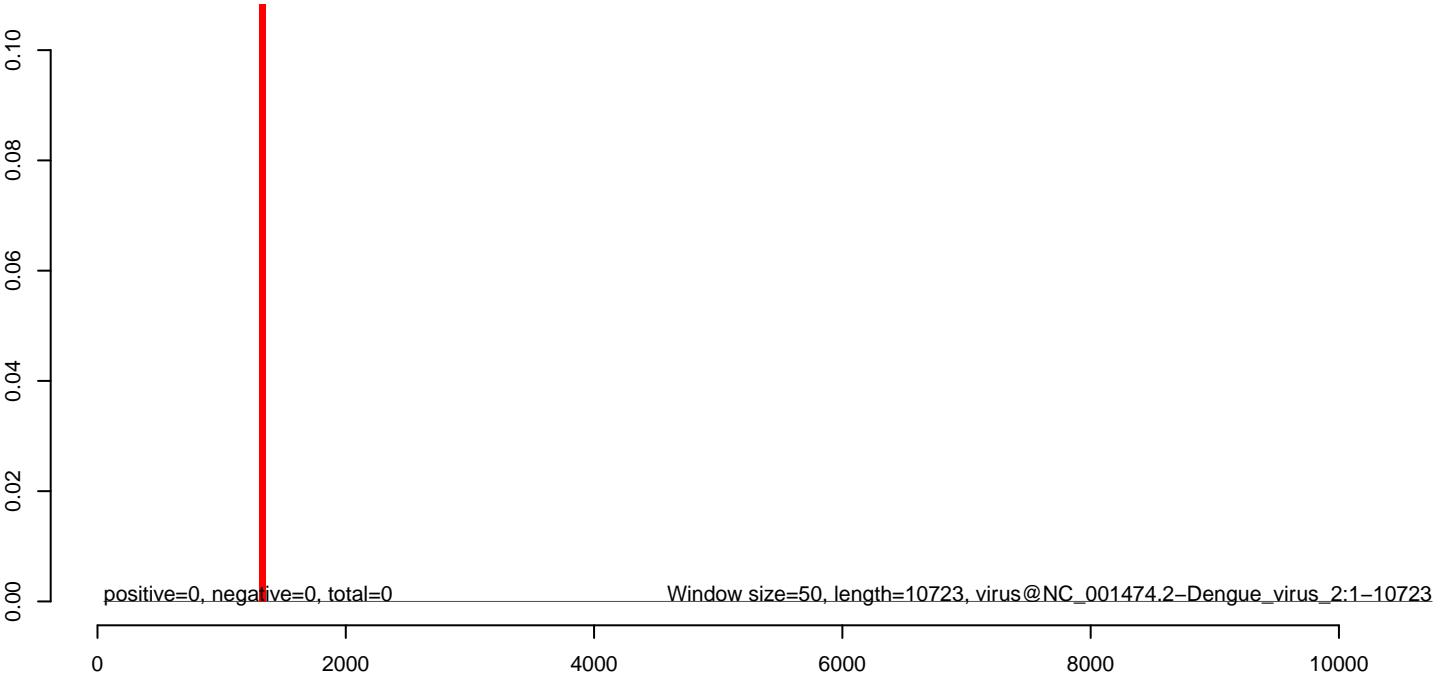
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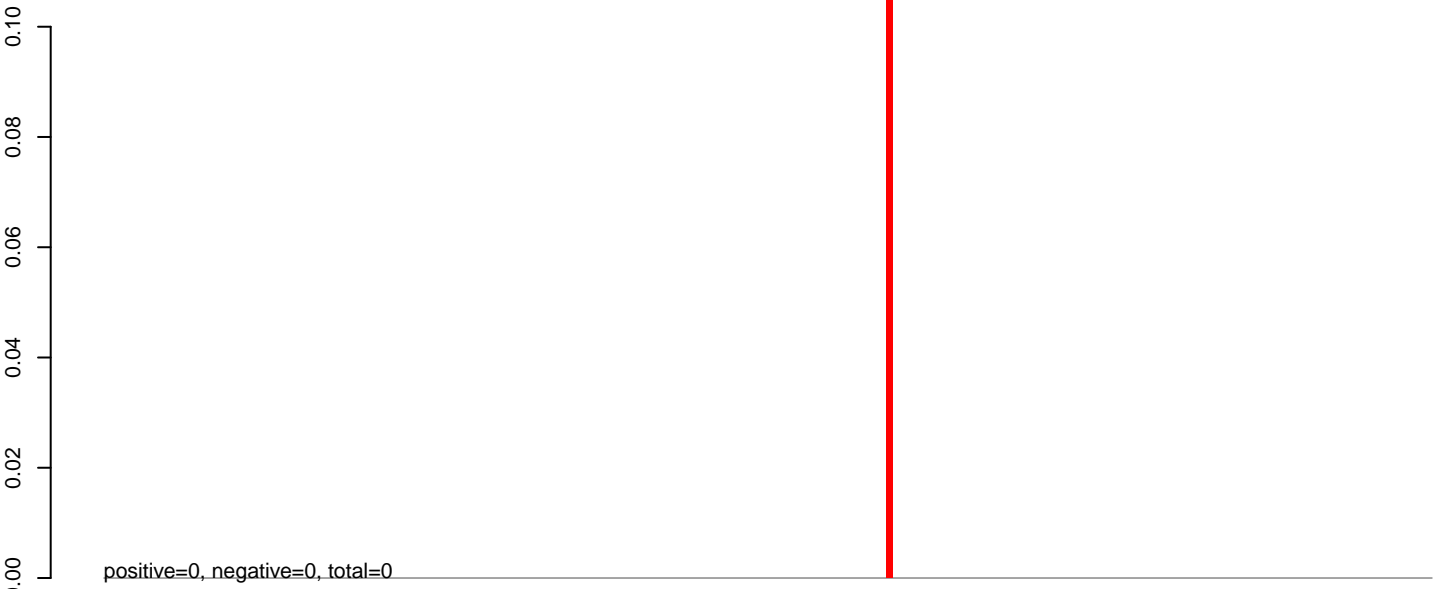
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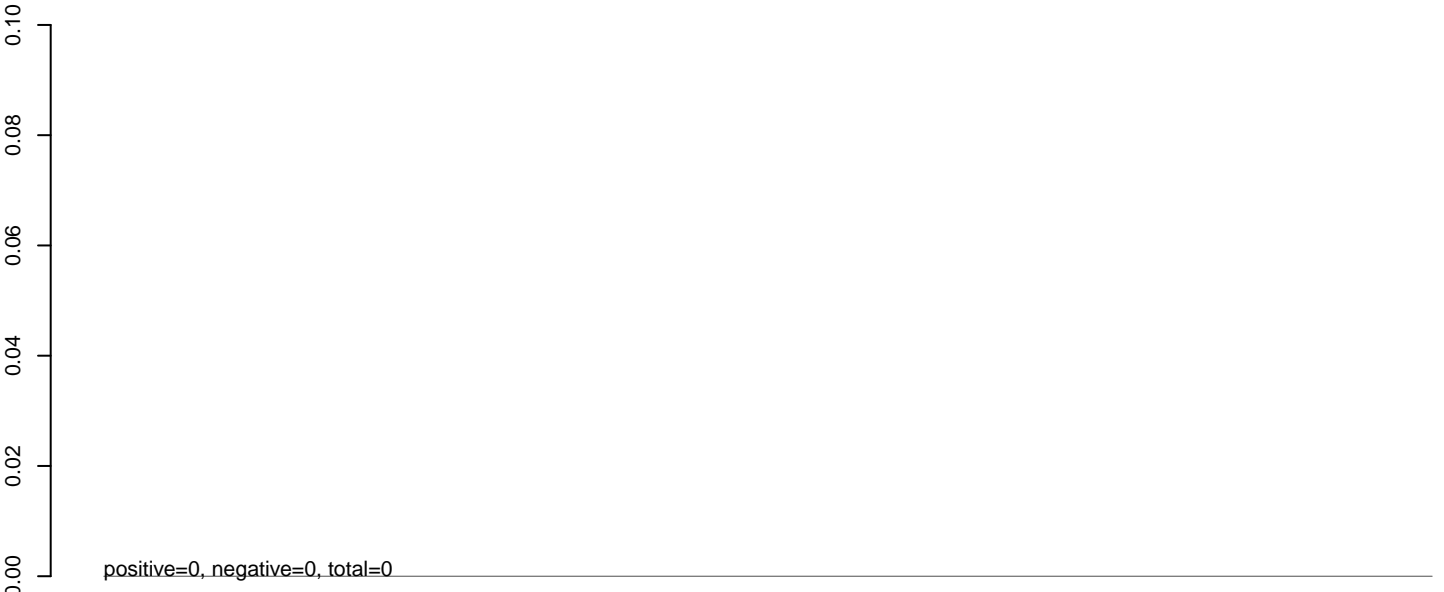
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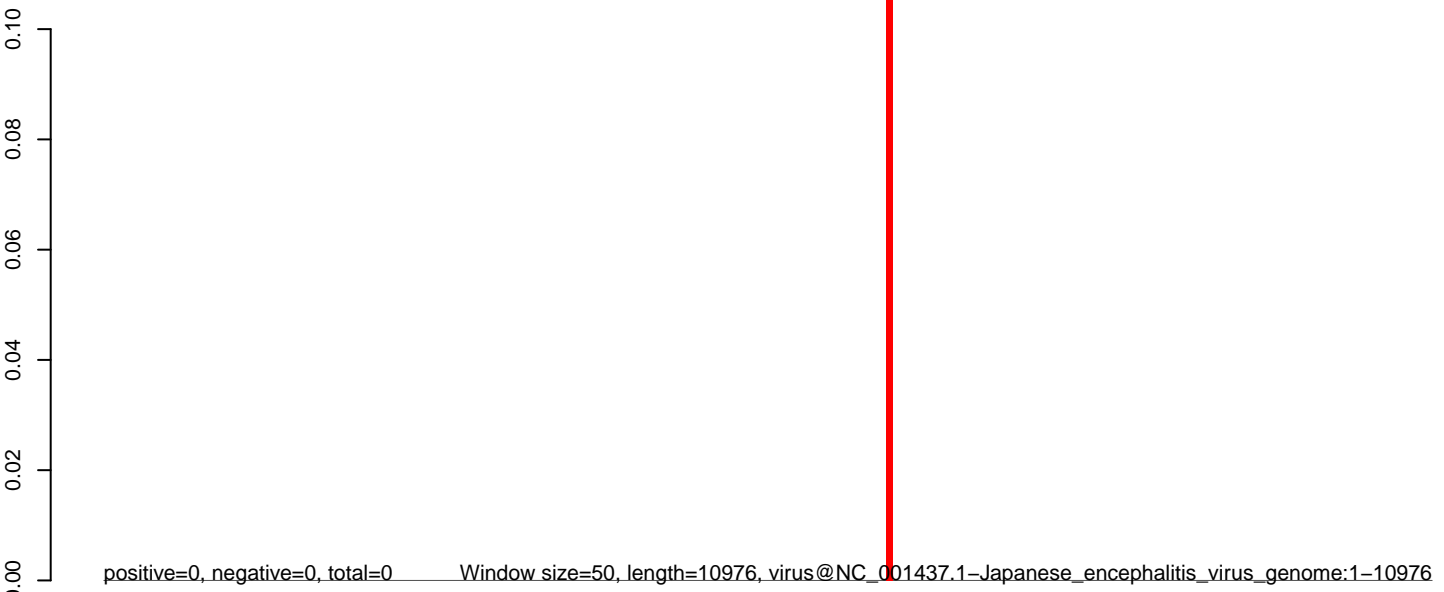
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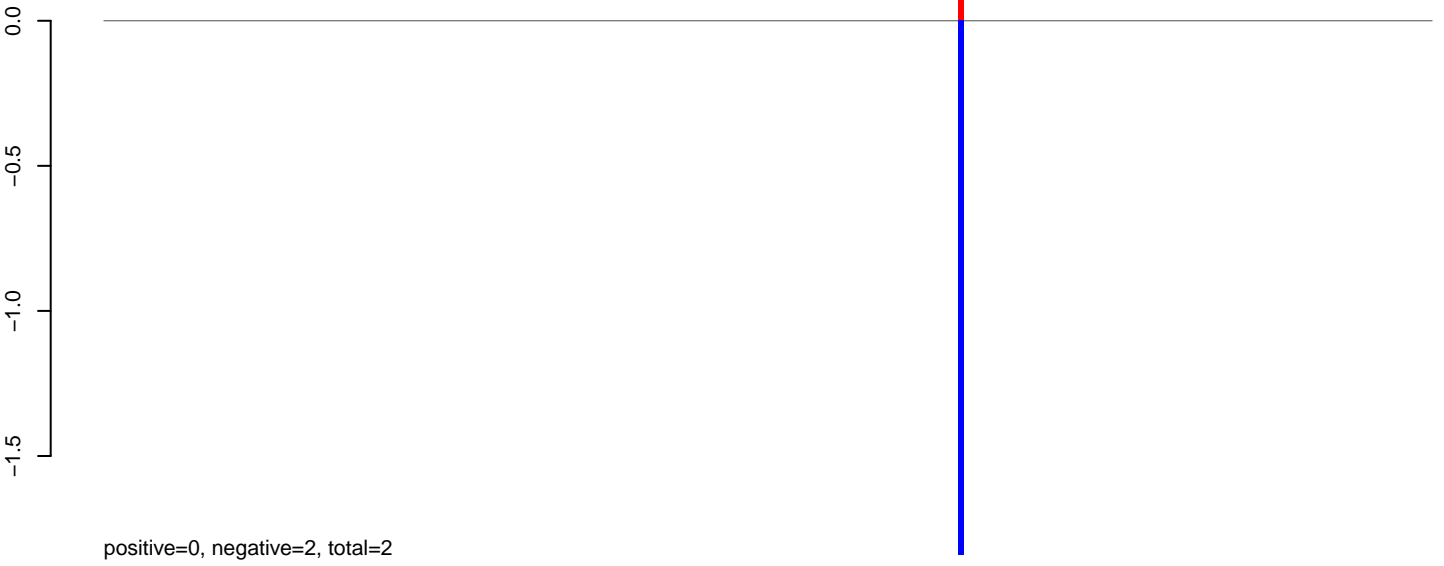


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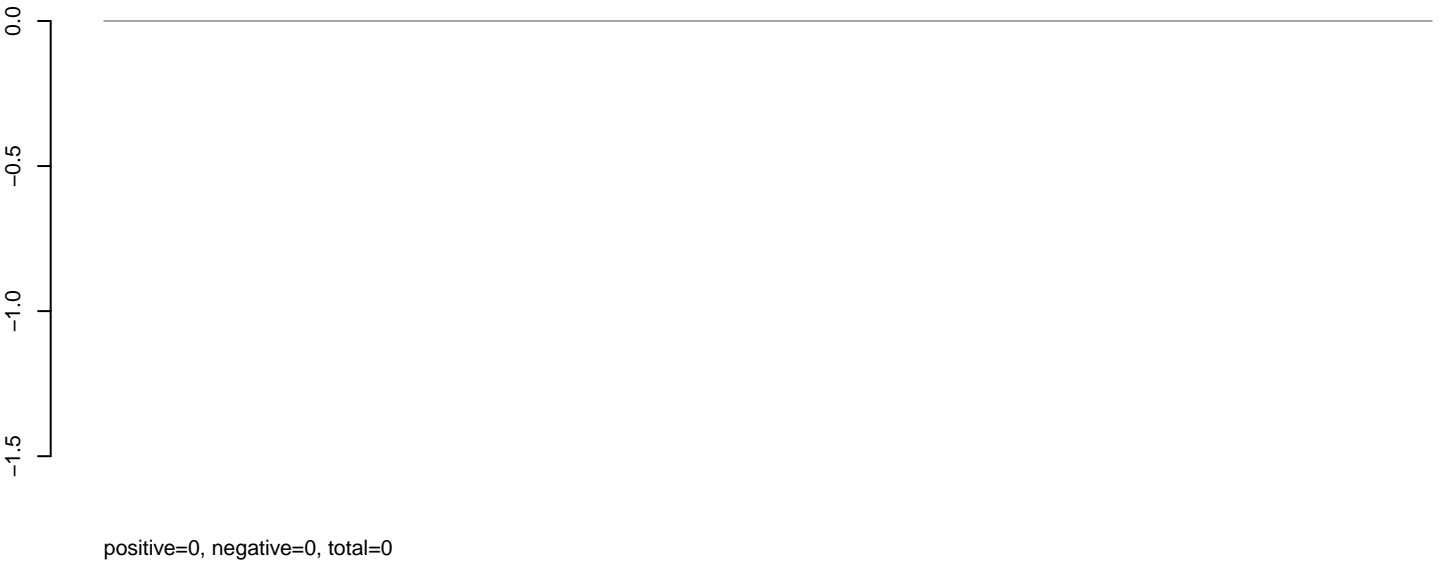


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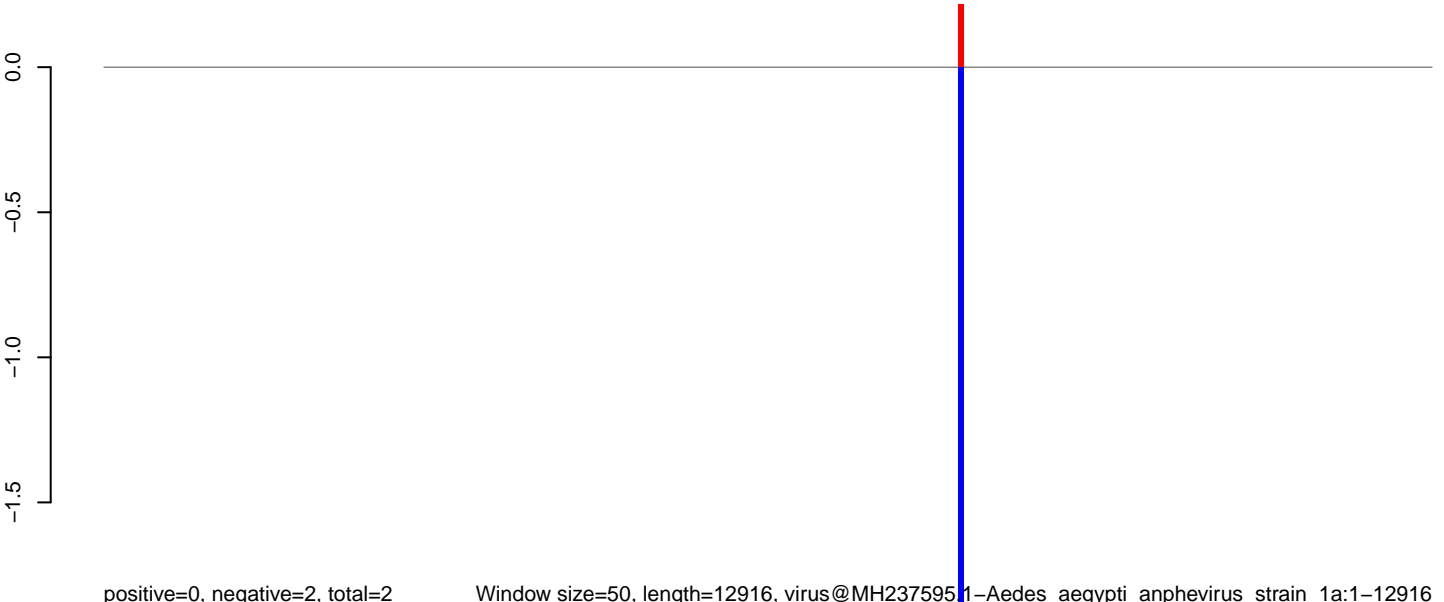
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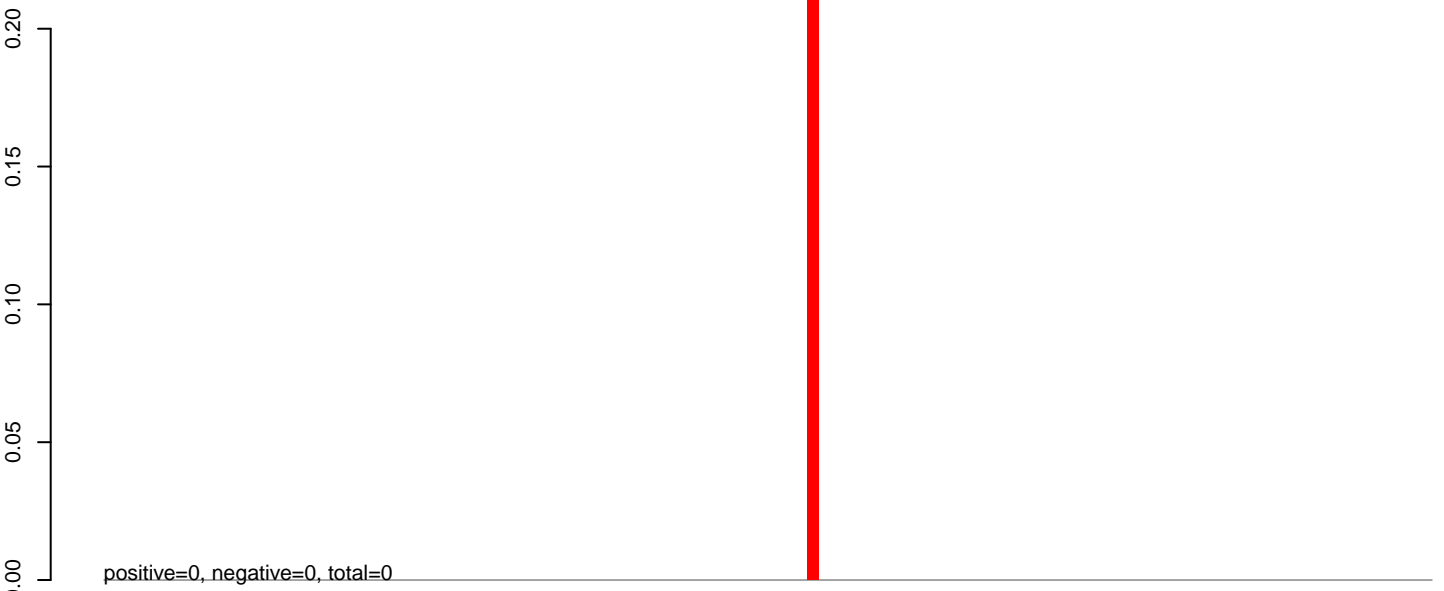
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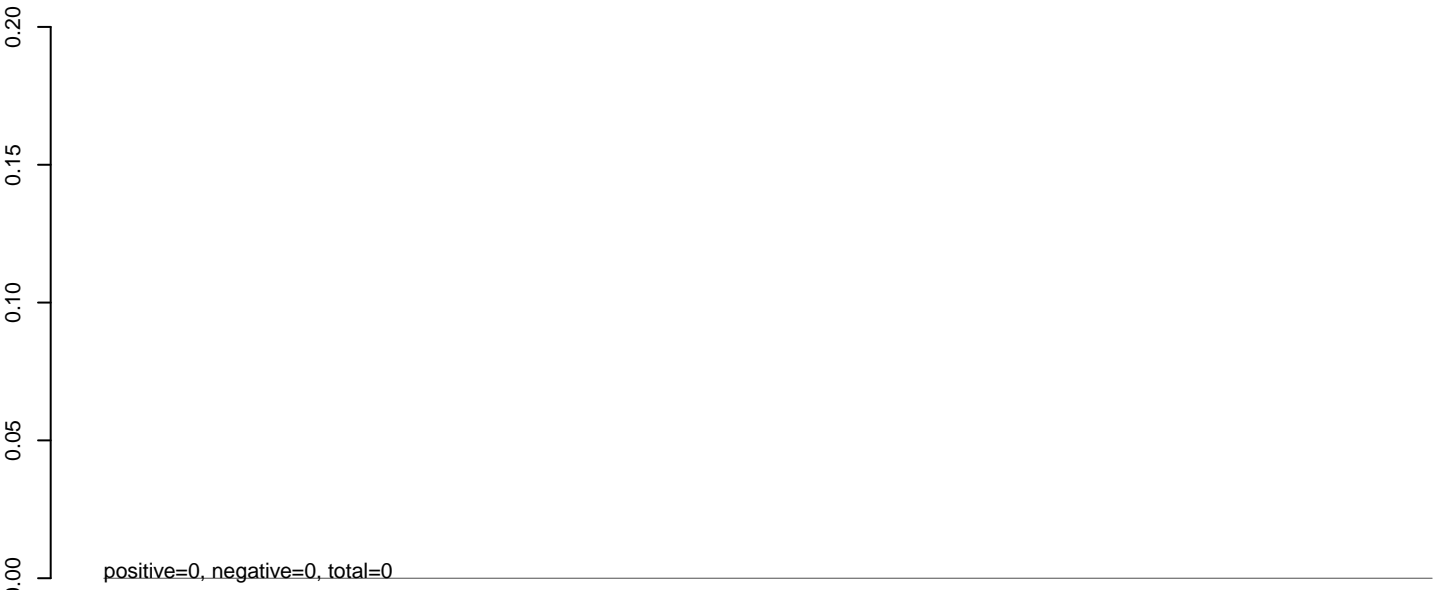
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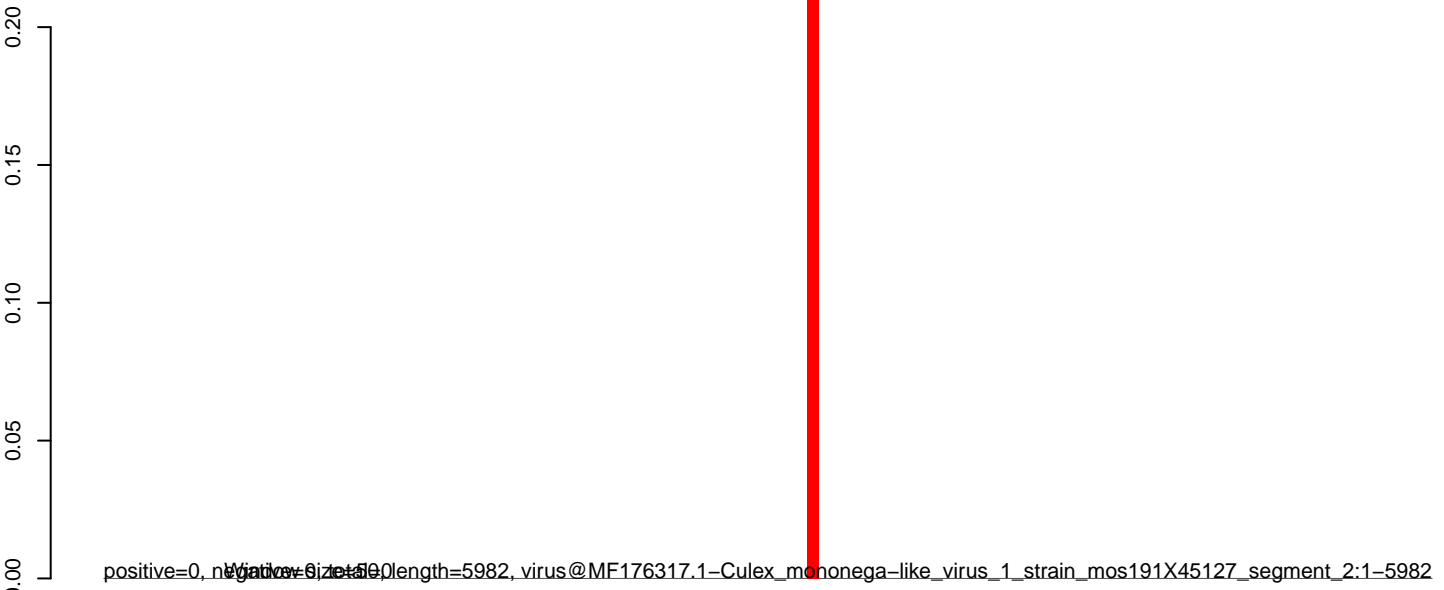
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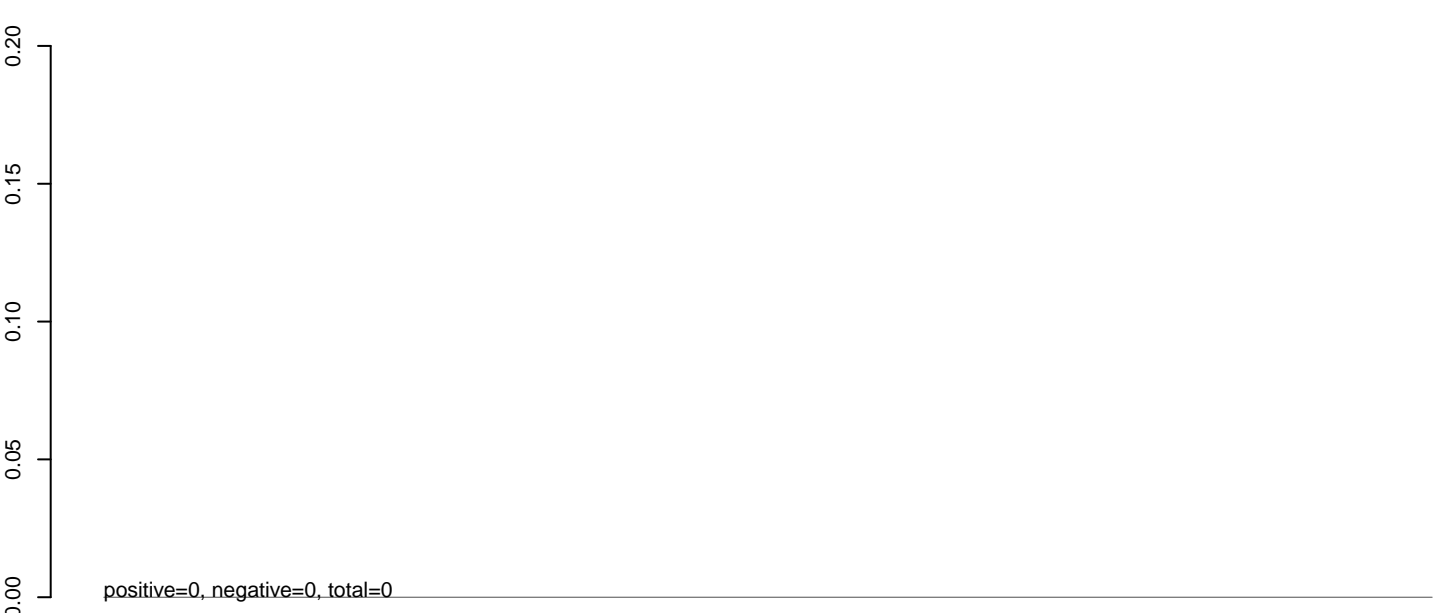


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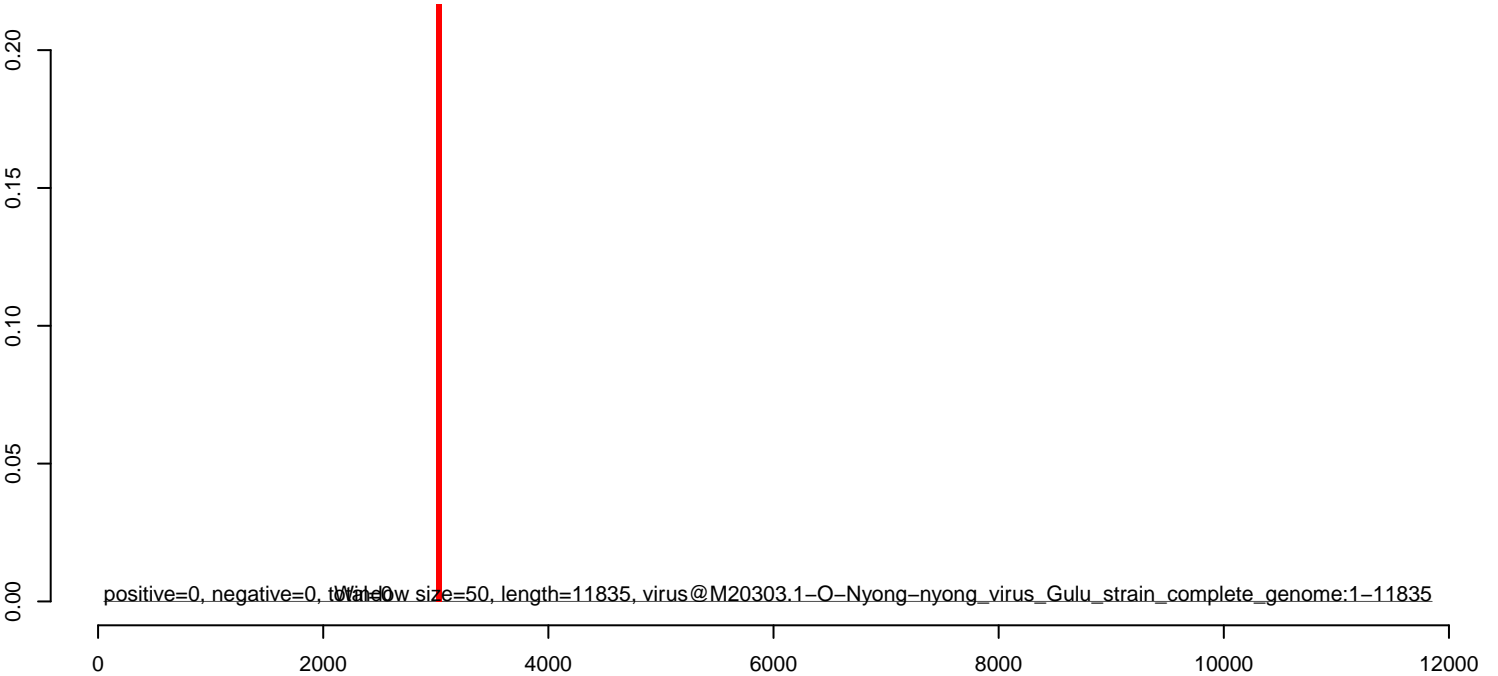
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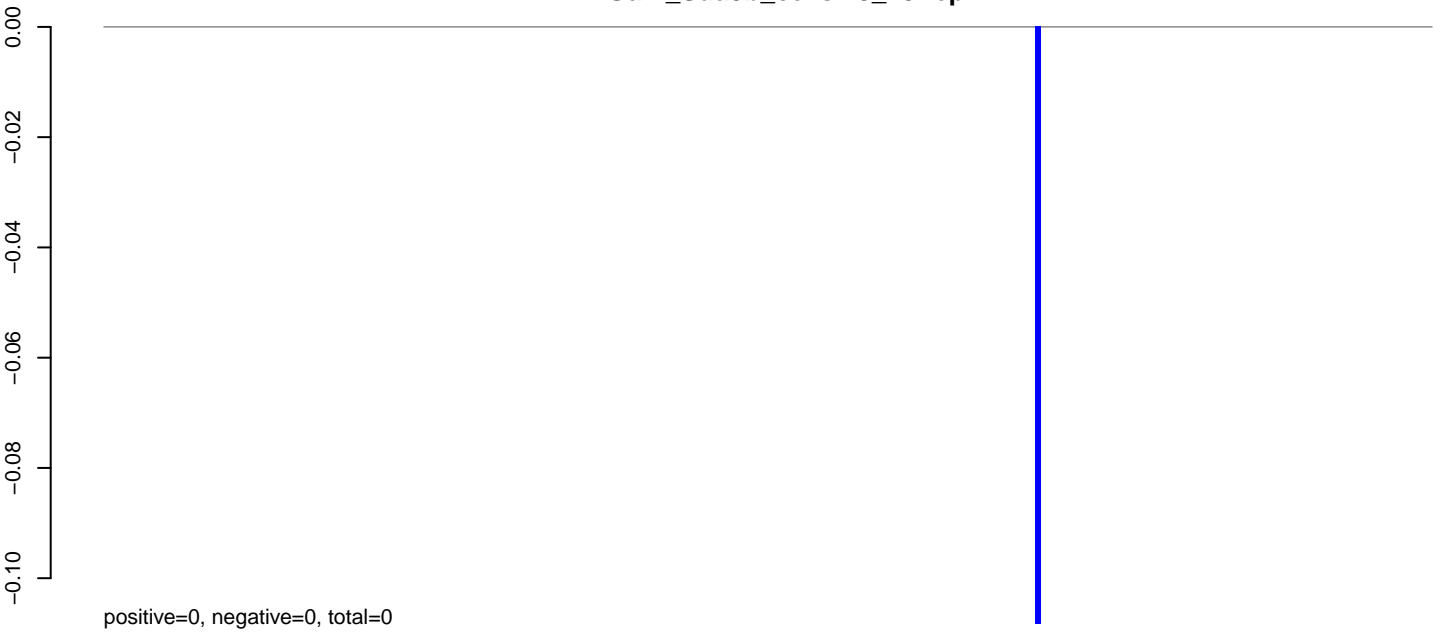
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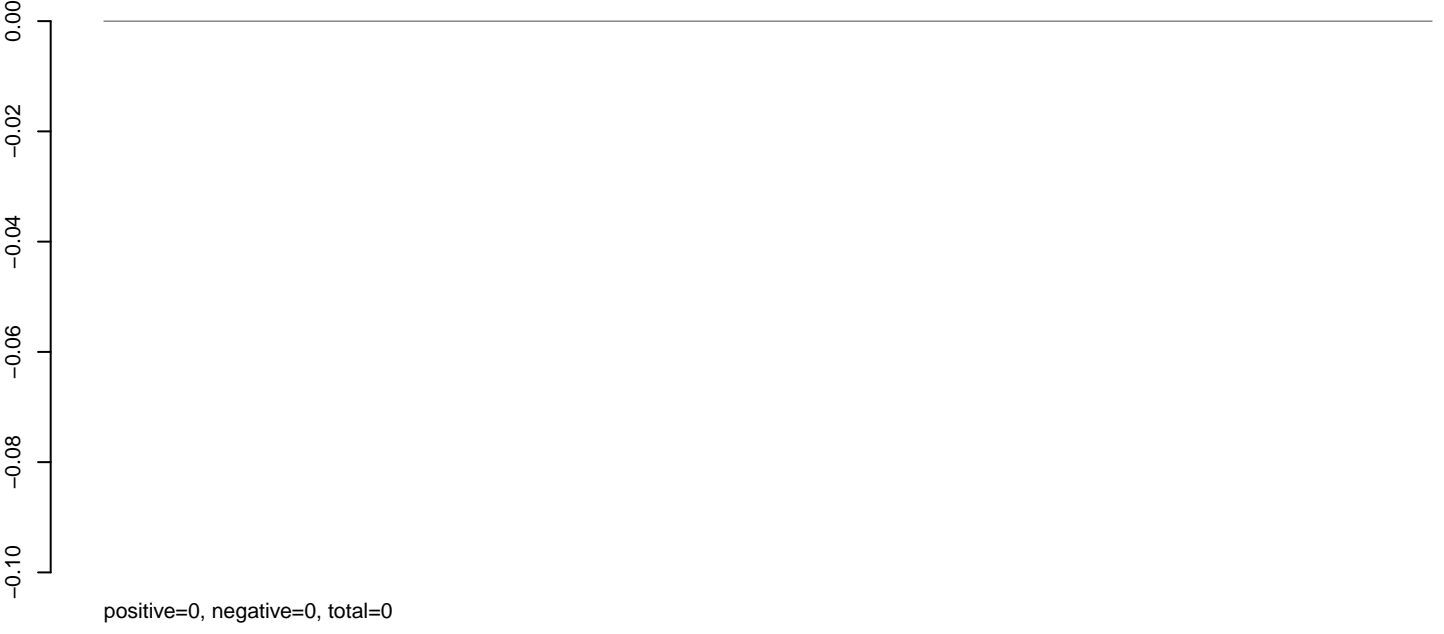
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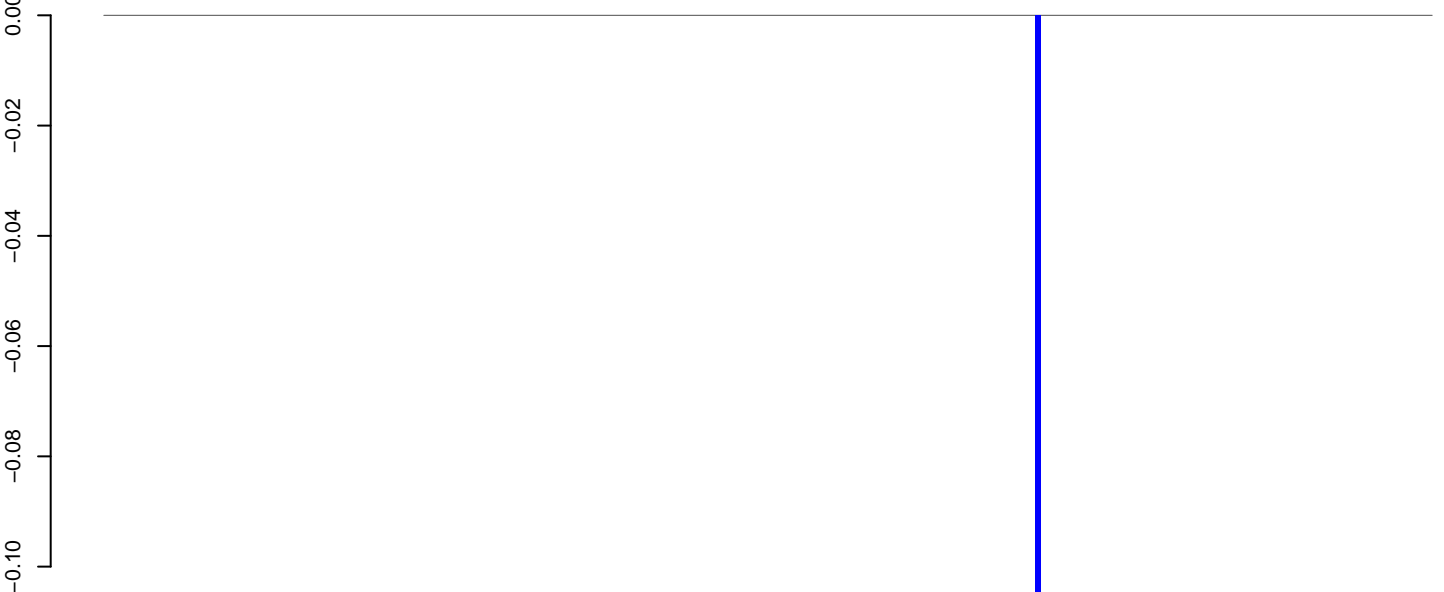
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AnGam\_Sua5b\_cells.rep

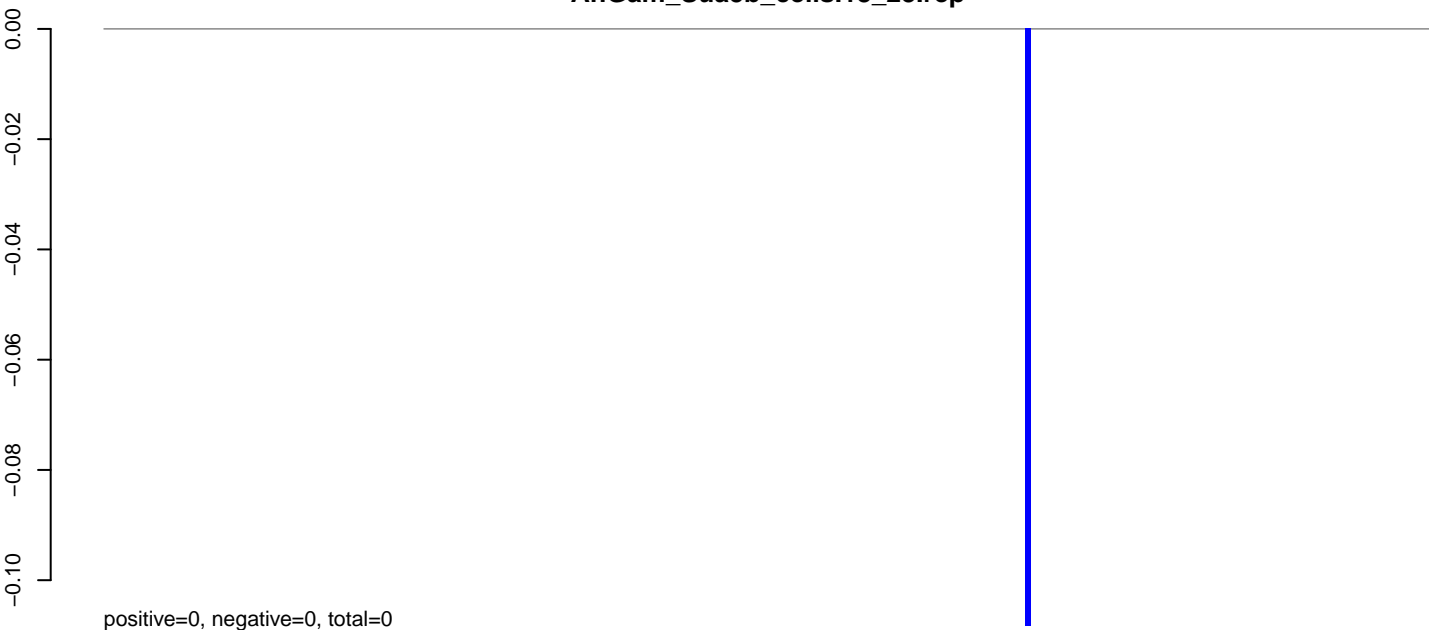


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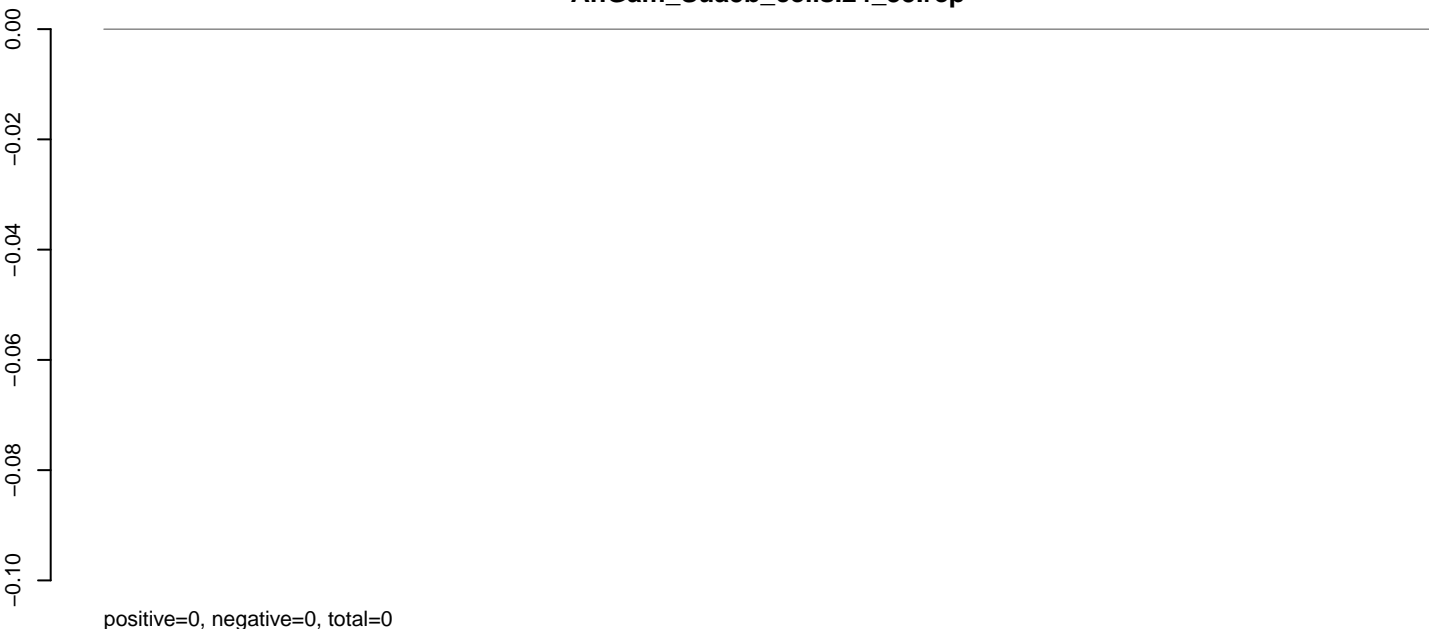
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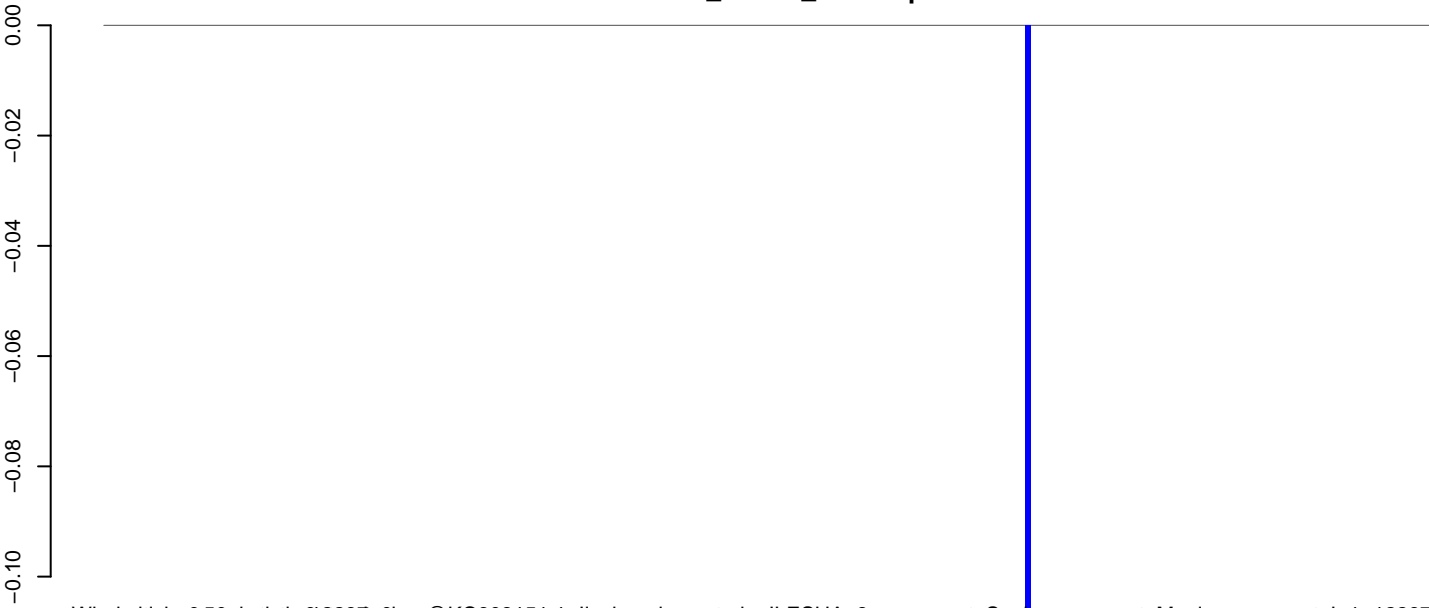
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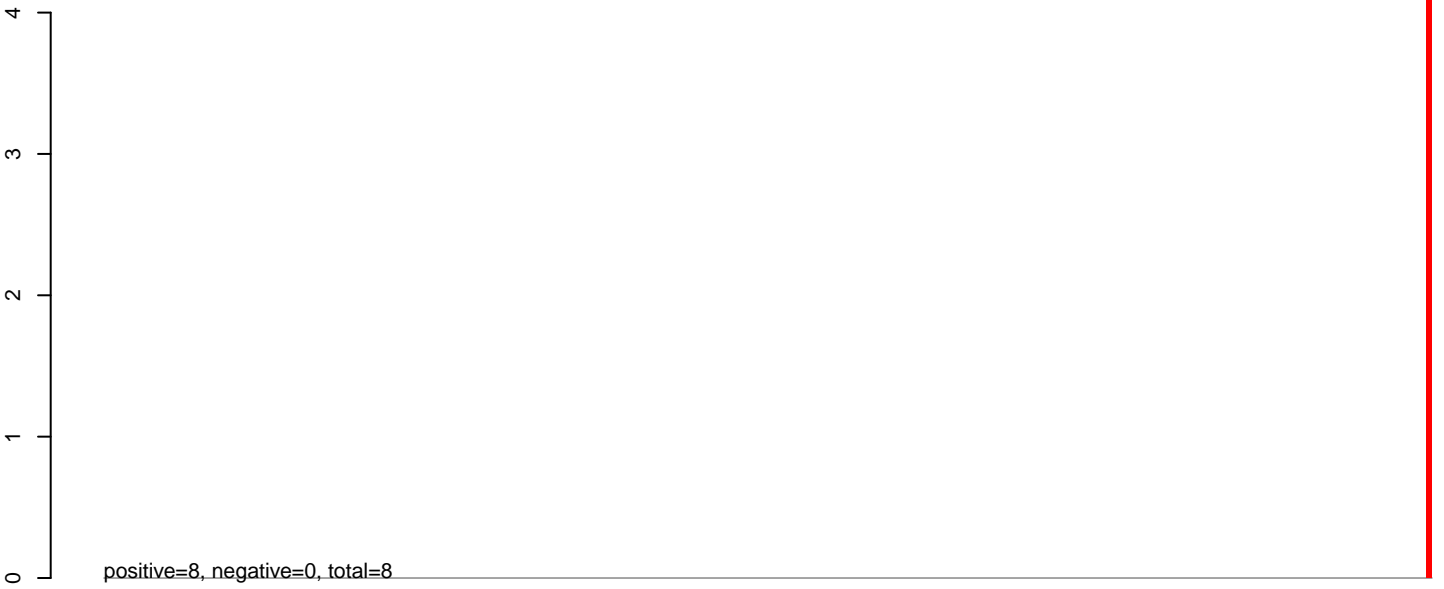
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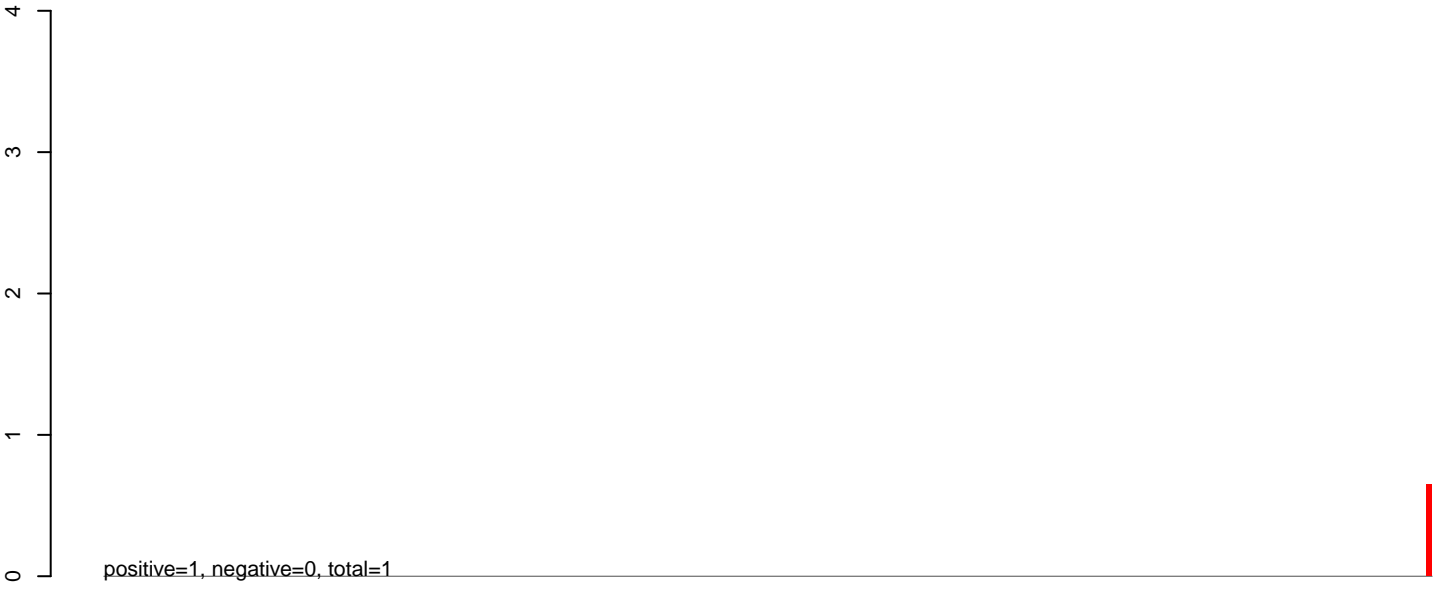
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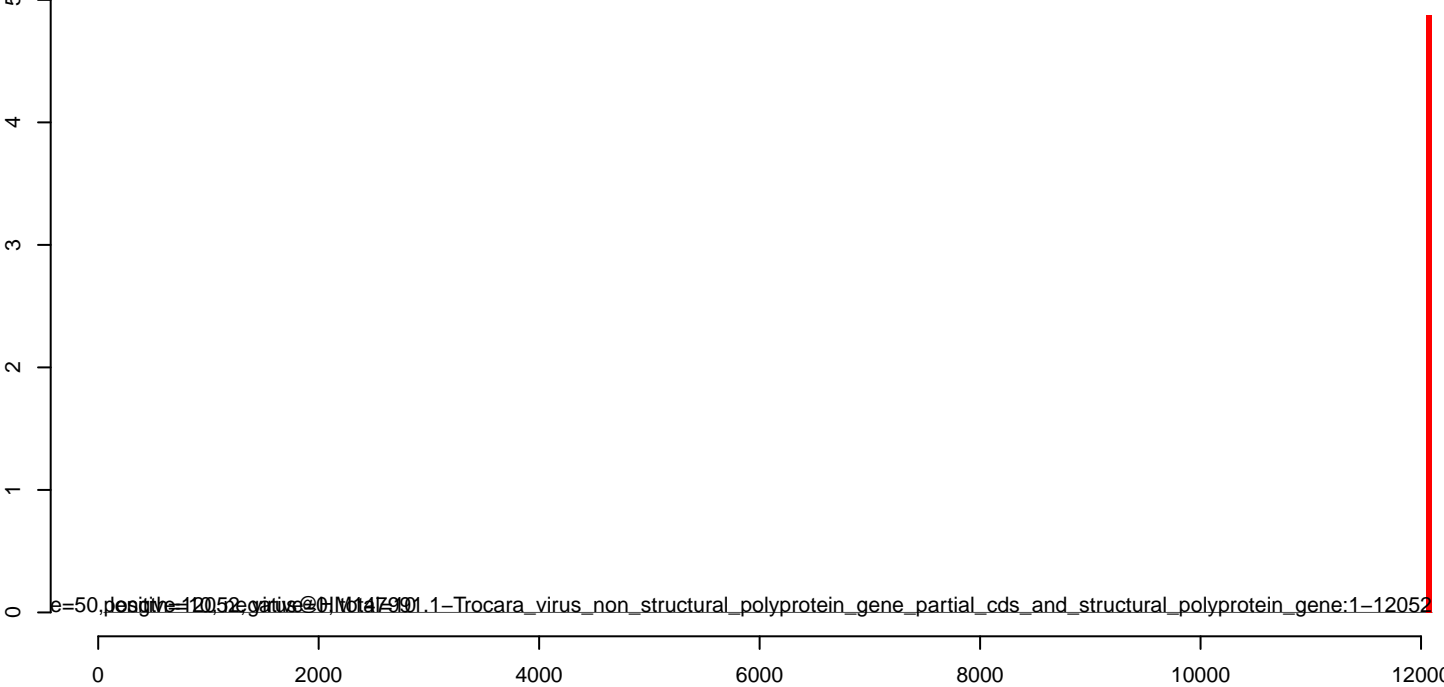
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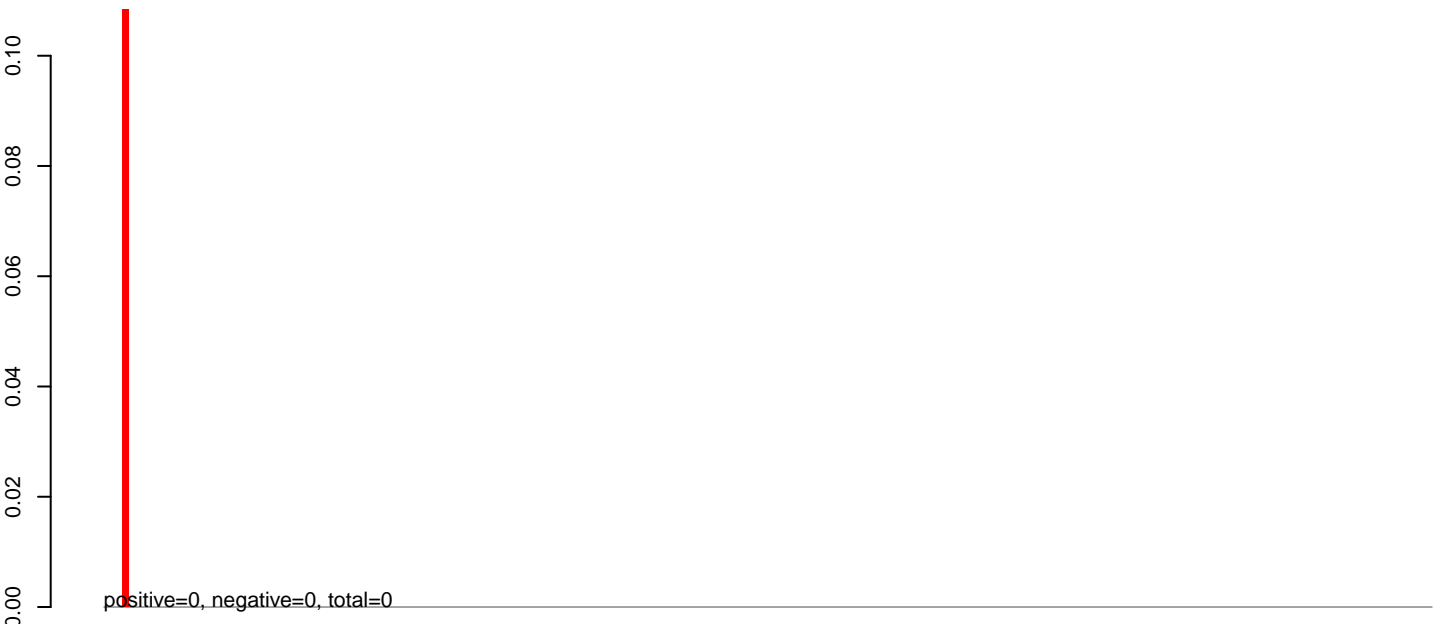
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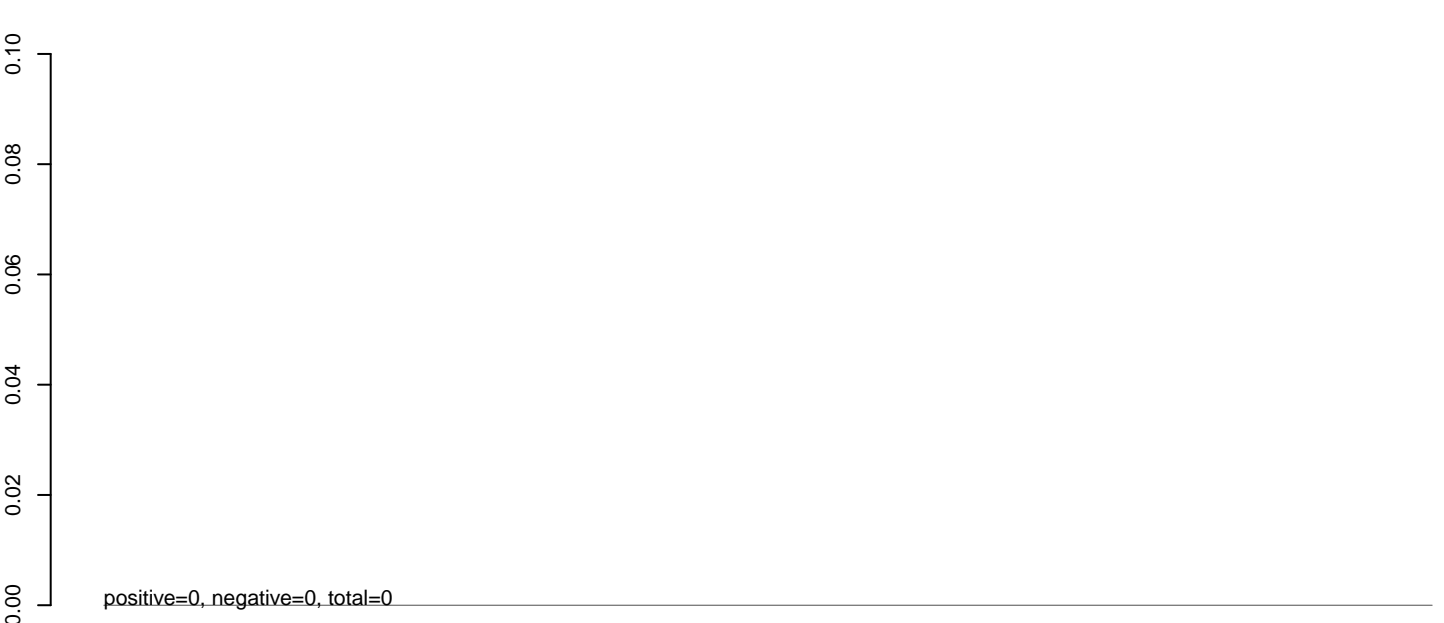
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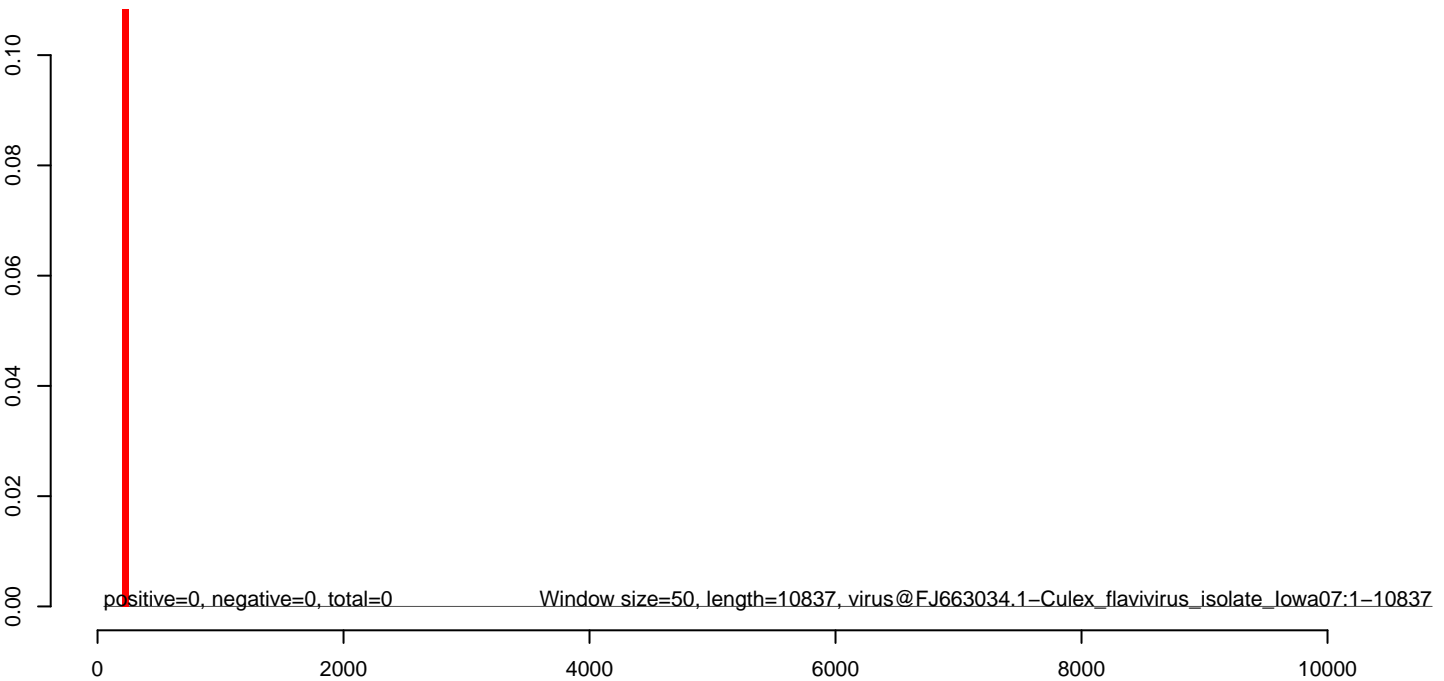
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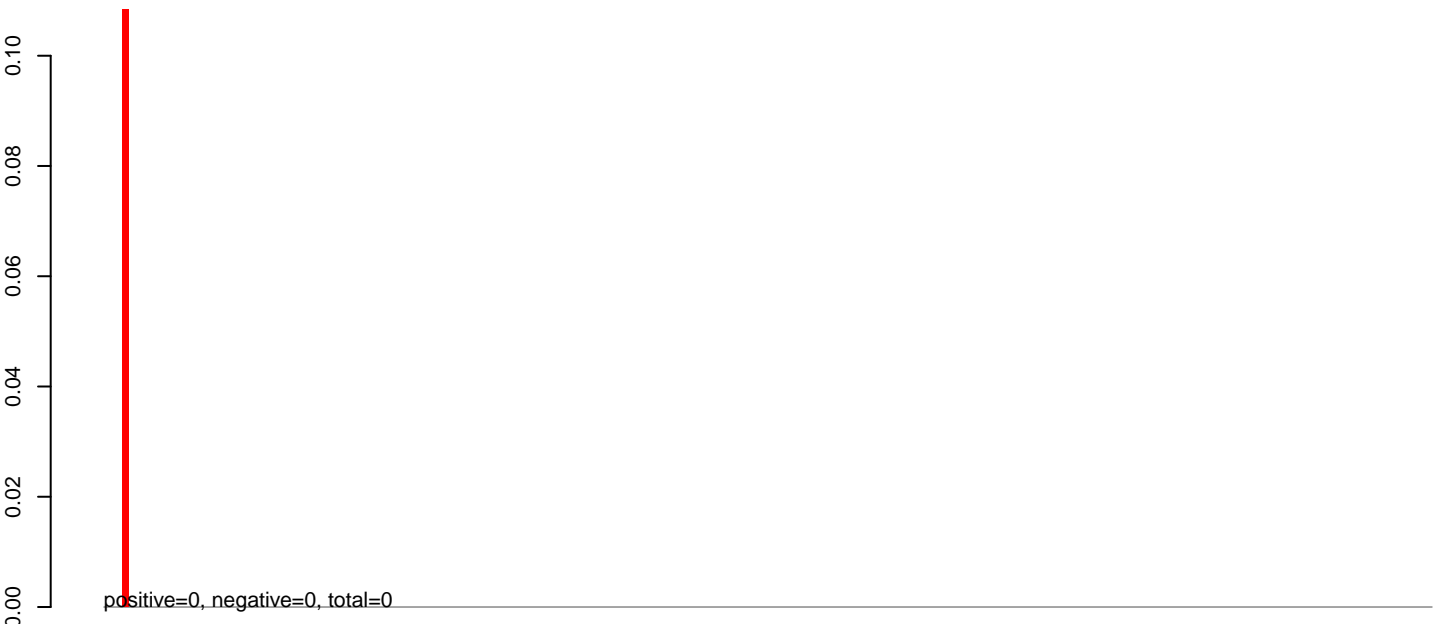
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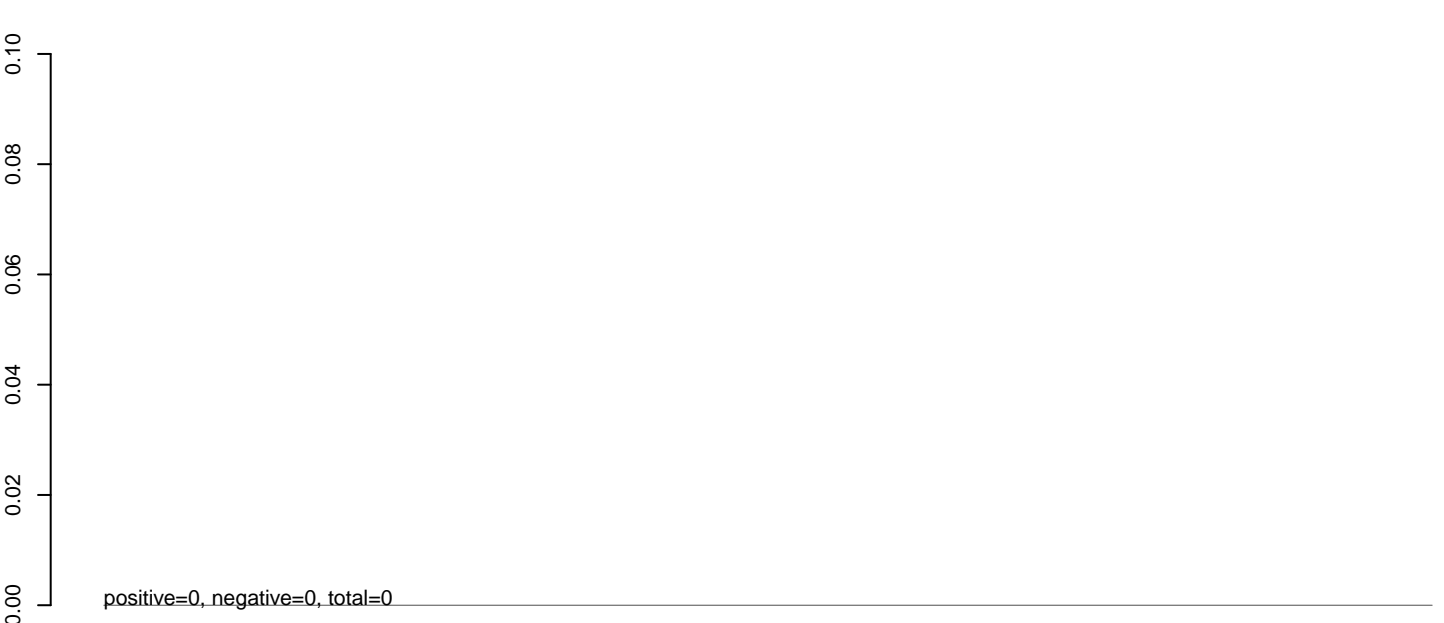
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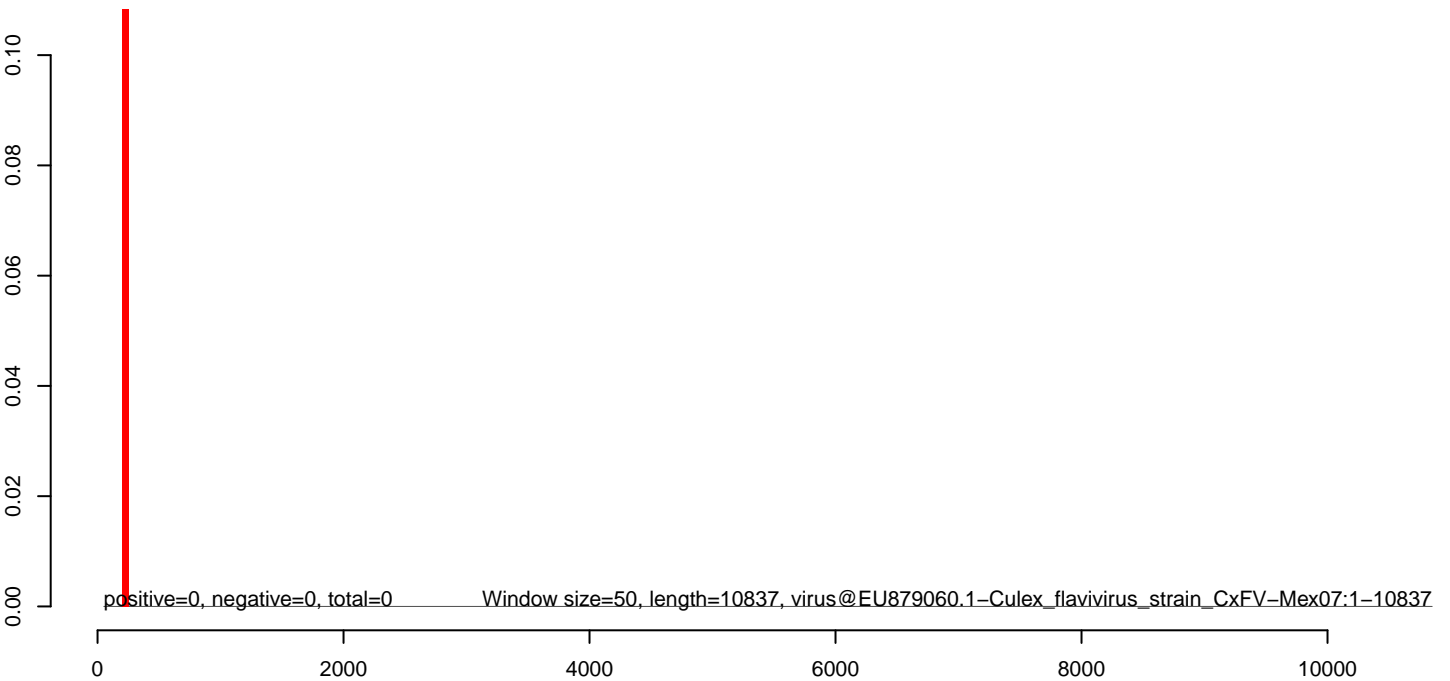
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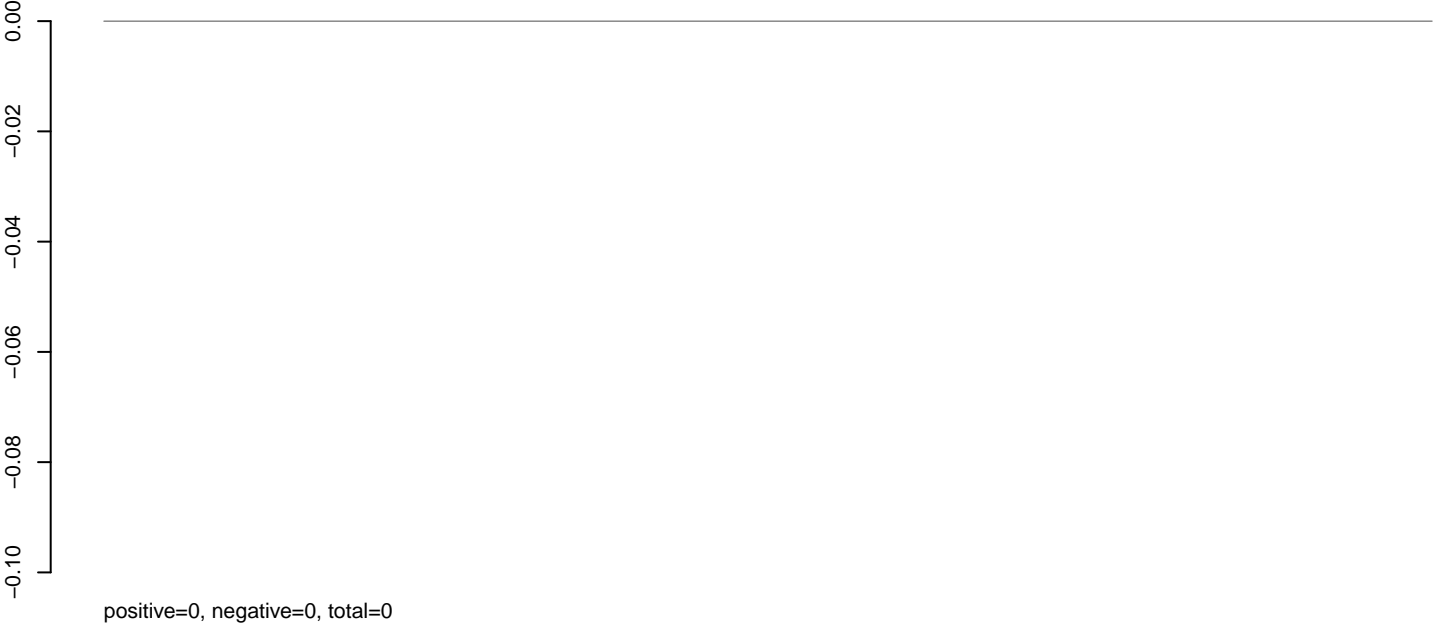
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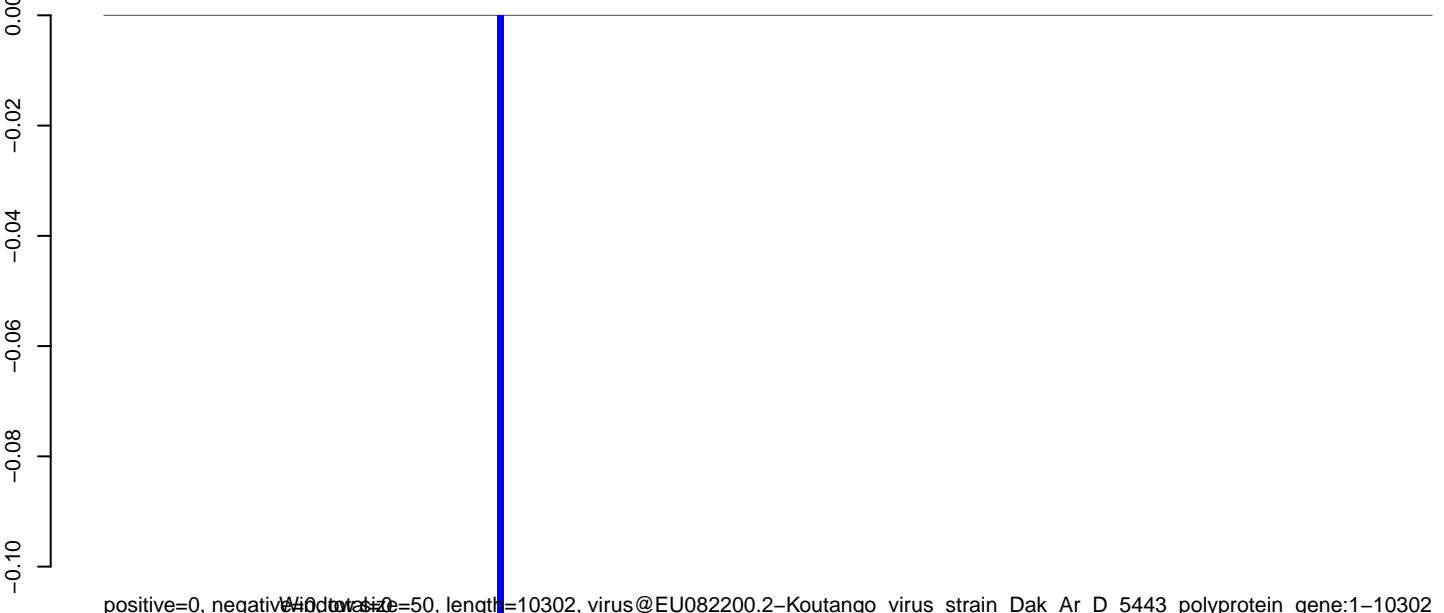
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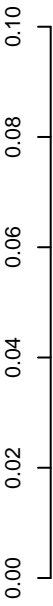
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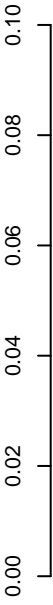
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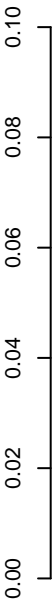
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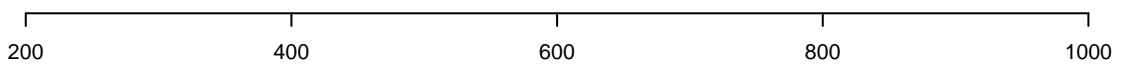


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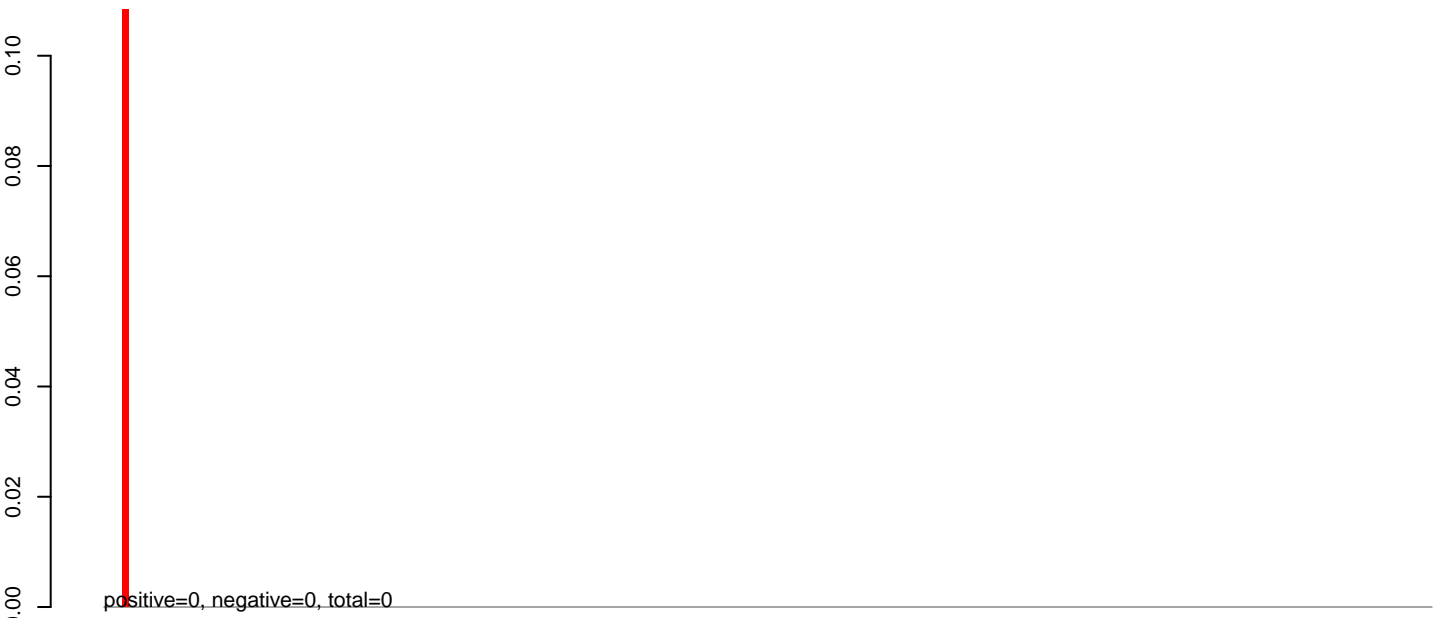
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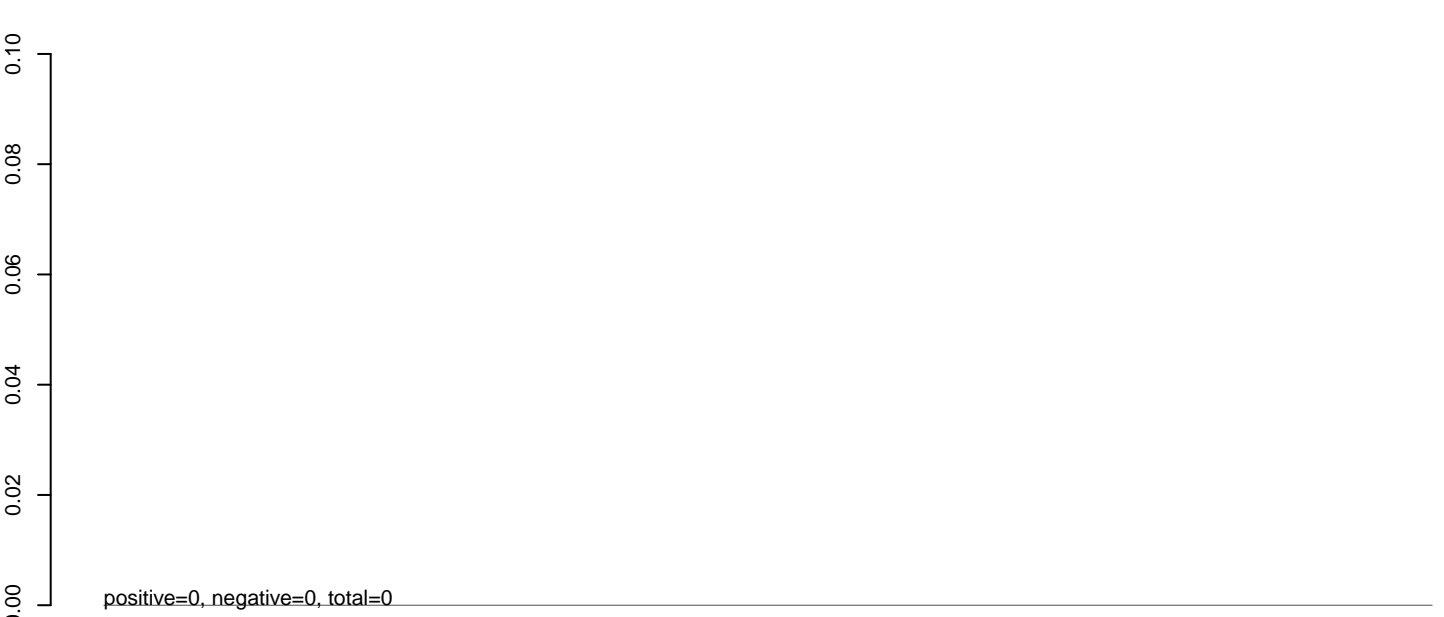
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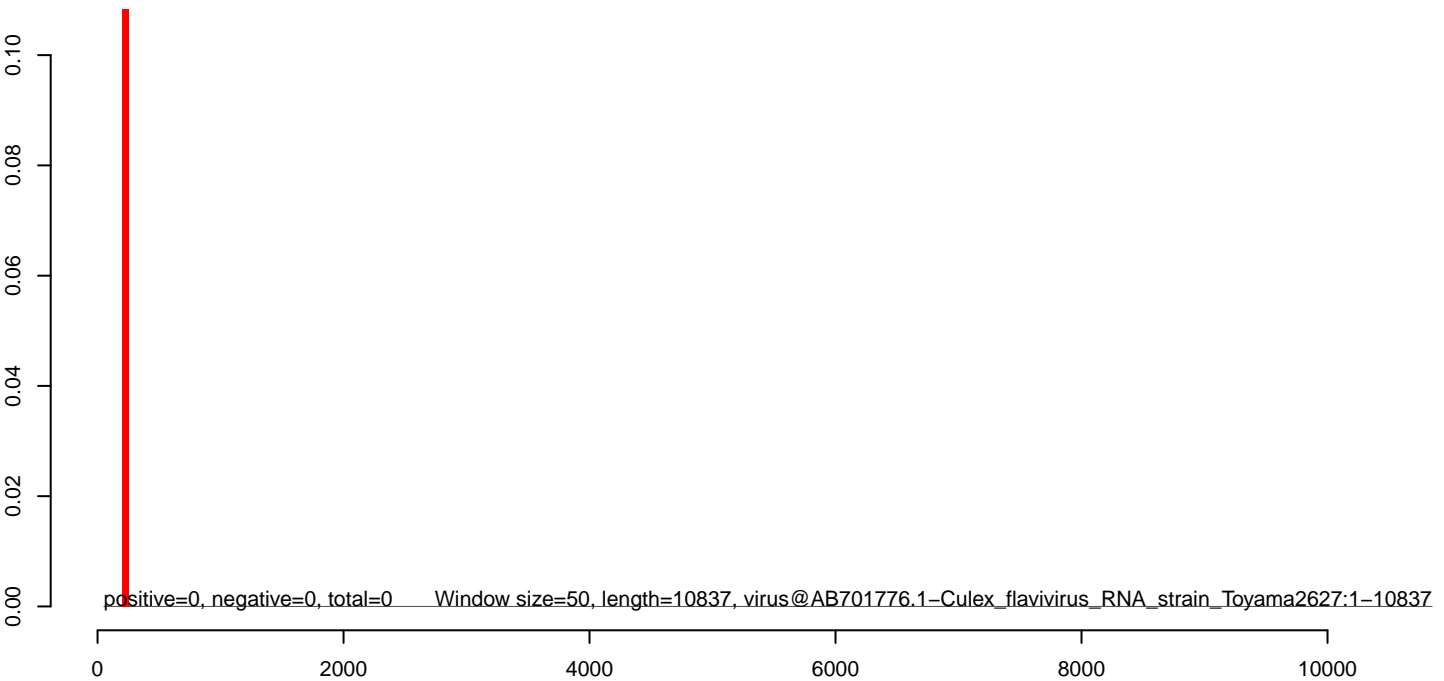
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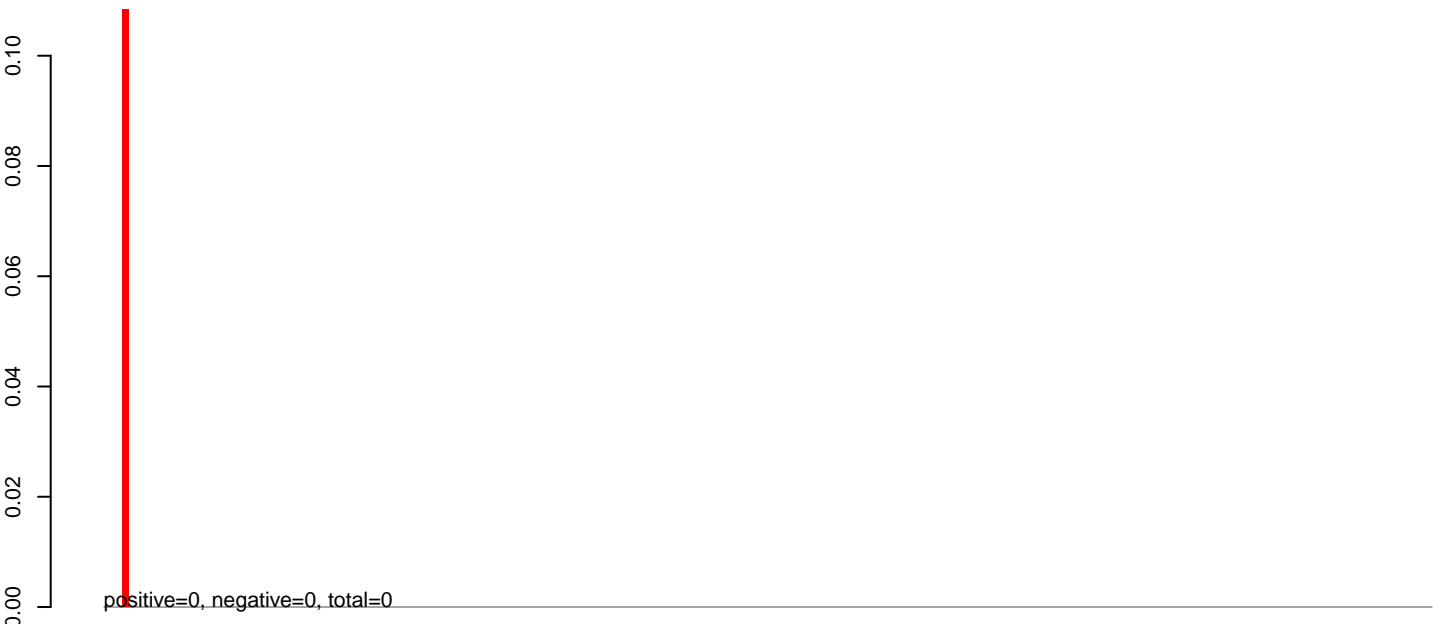
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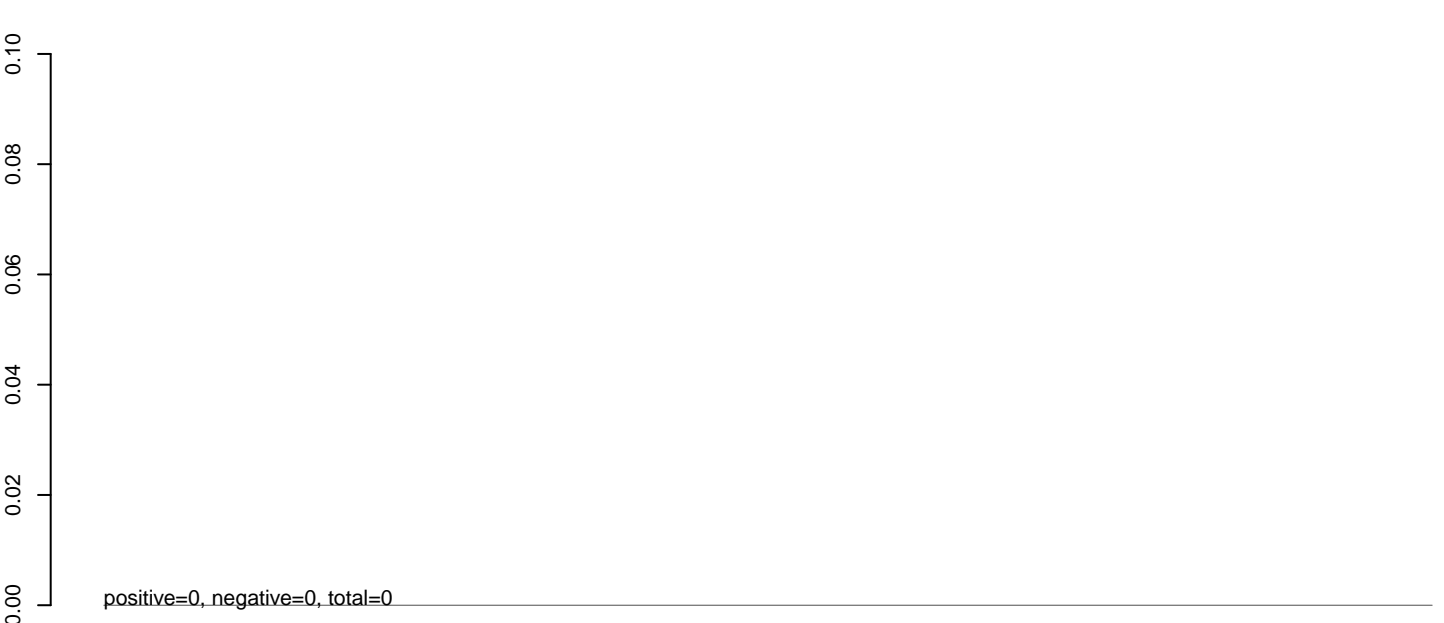
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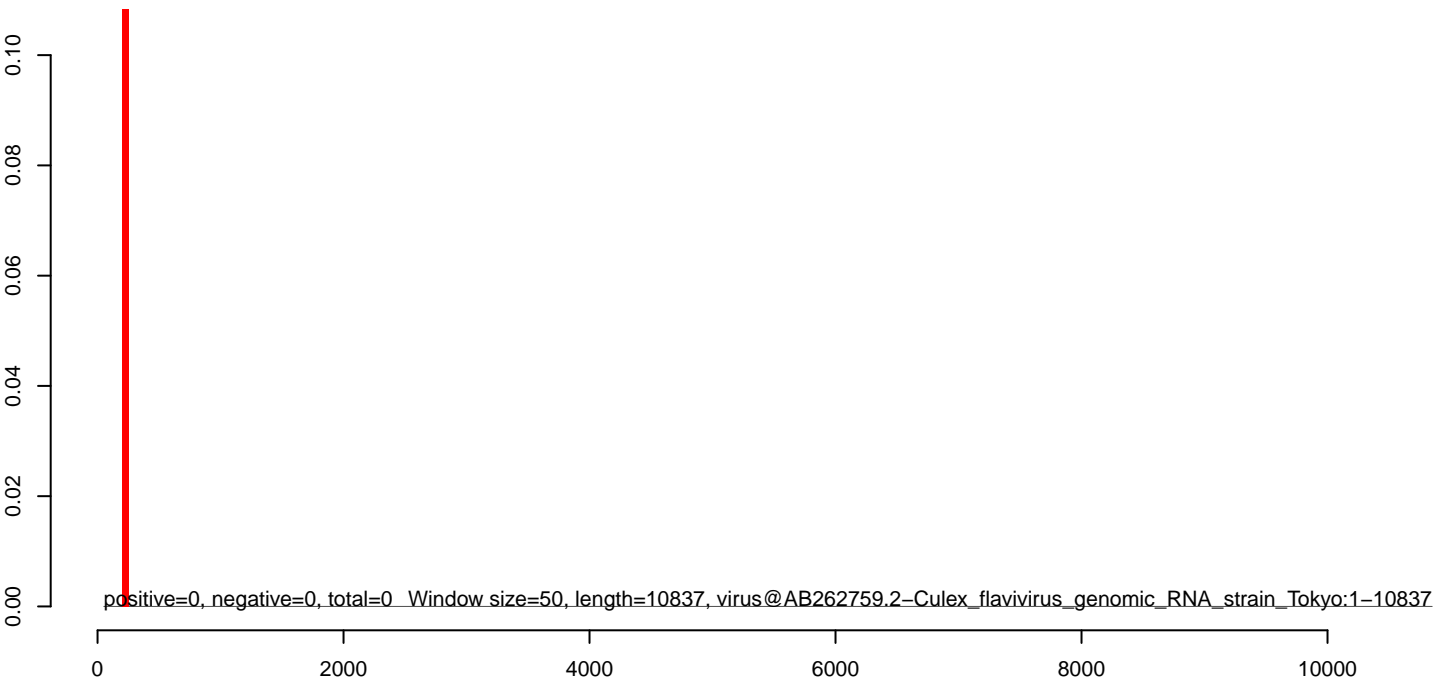
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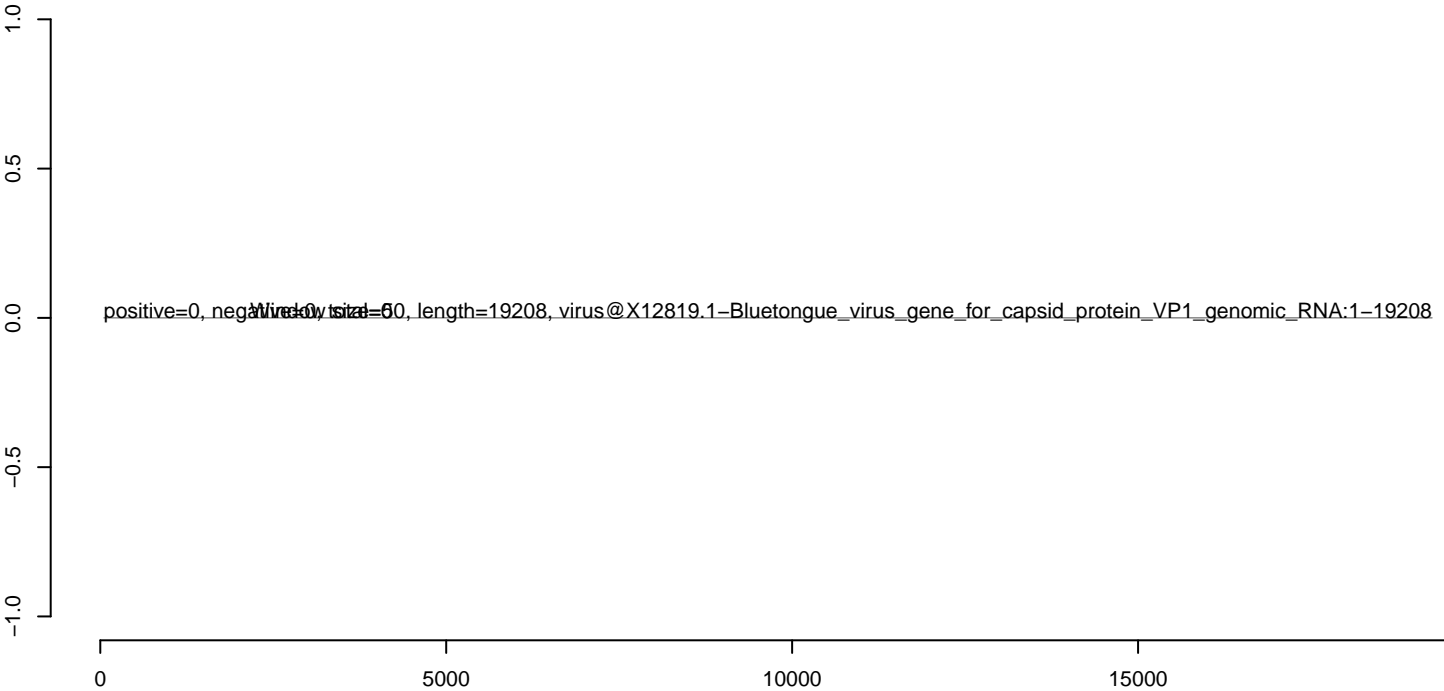
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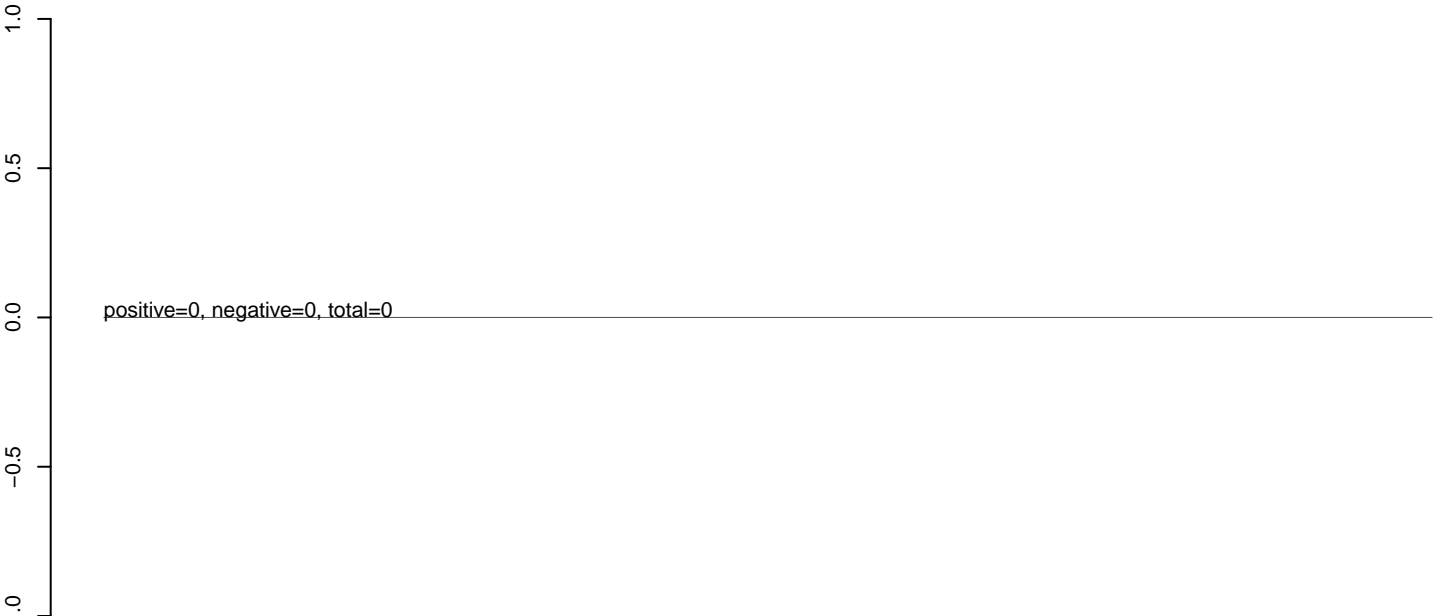
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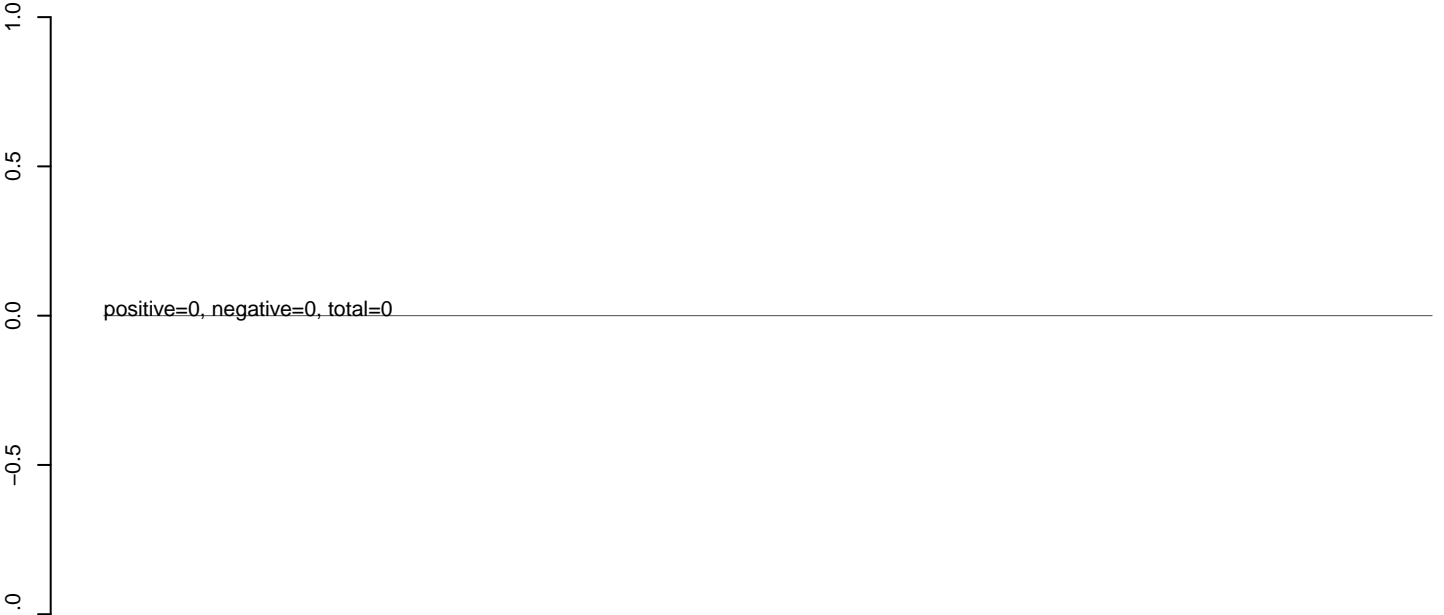
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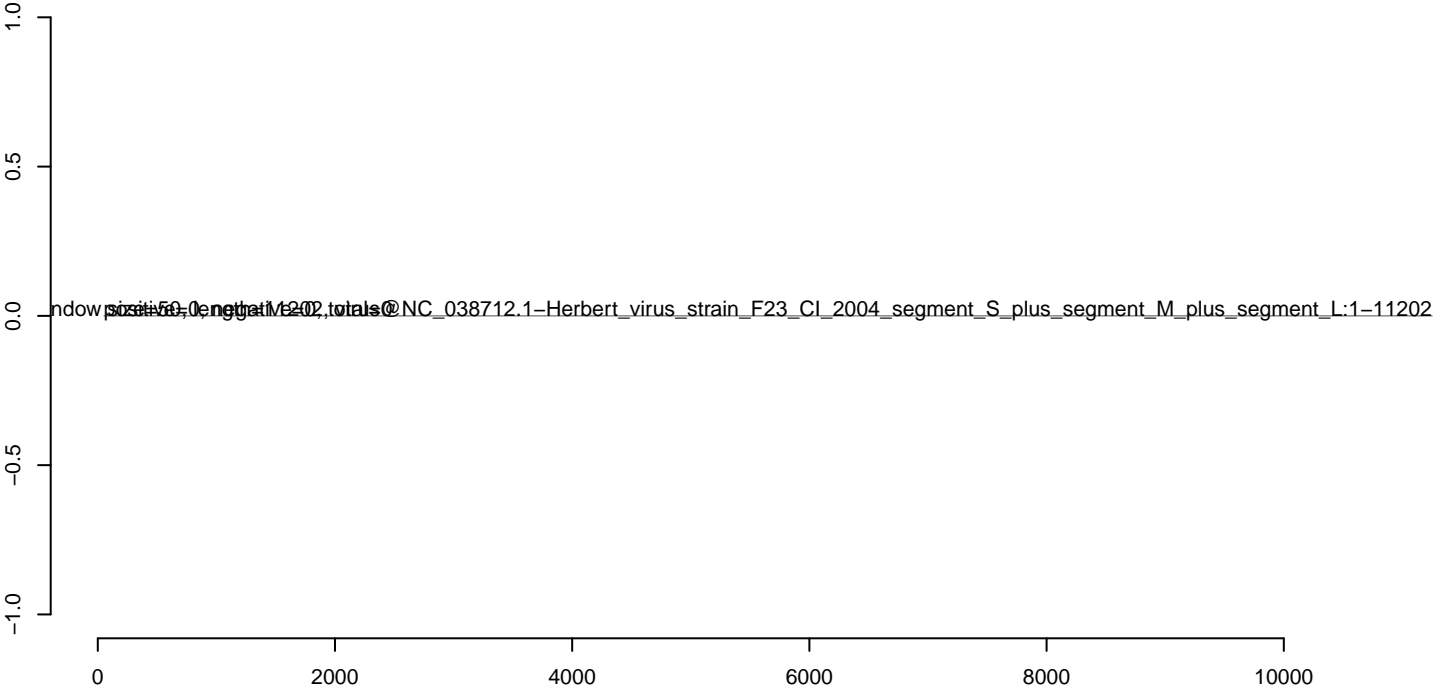
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AnGam\_Sua5b\_cells.rep



AnGam\_Sua5b\_cells.18\_23.rep



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AnGam\_Sua5b\_cells.rep



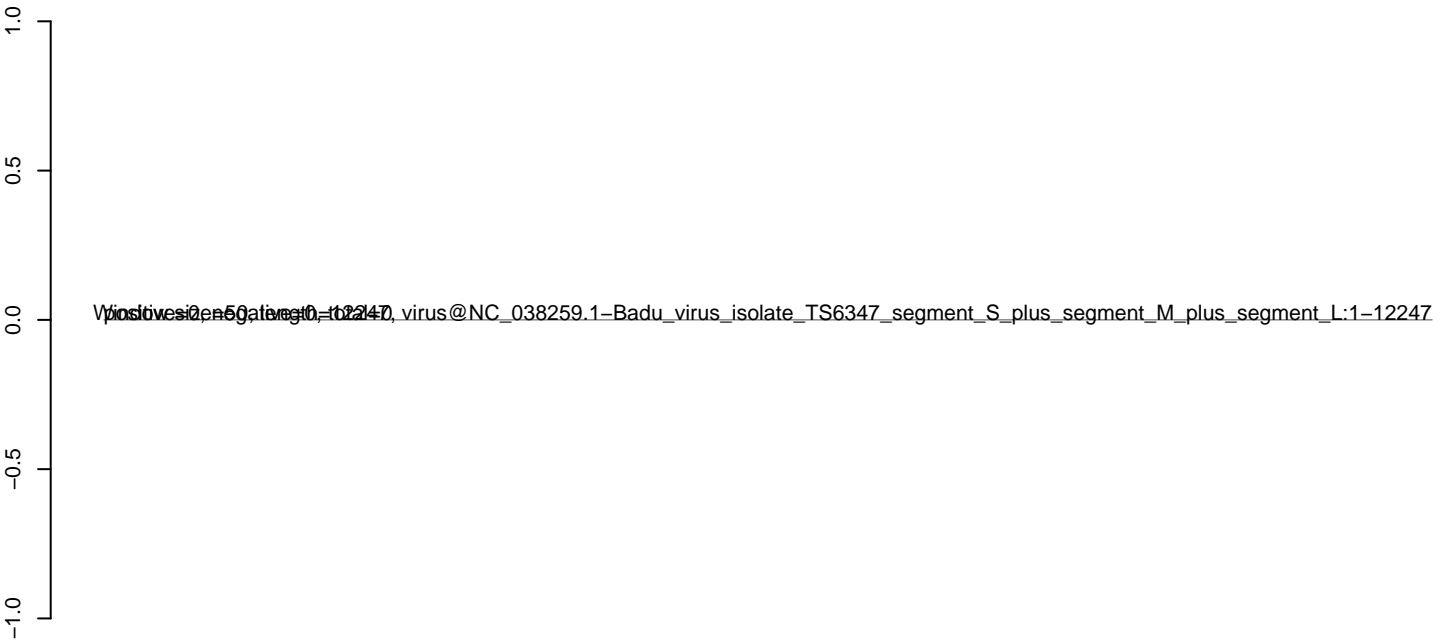
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AnGam\_Sua5b\_cells.rep



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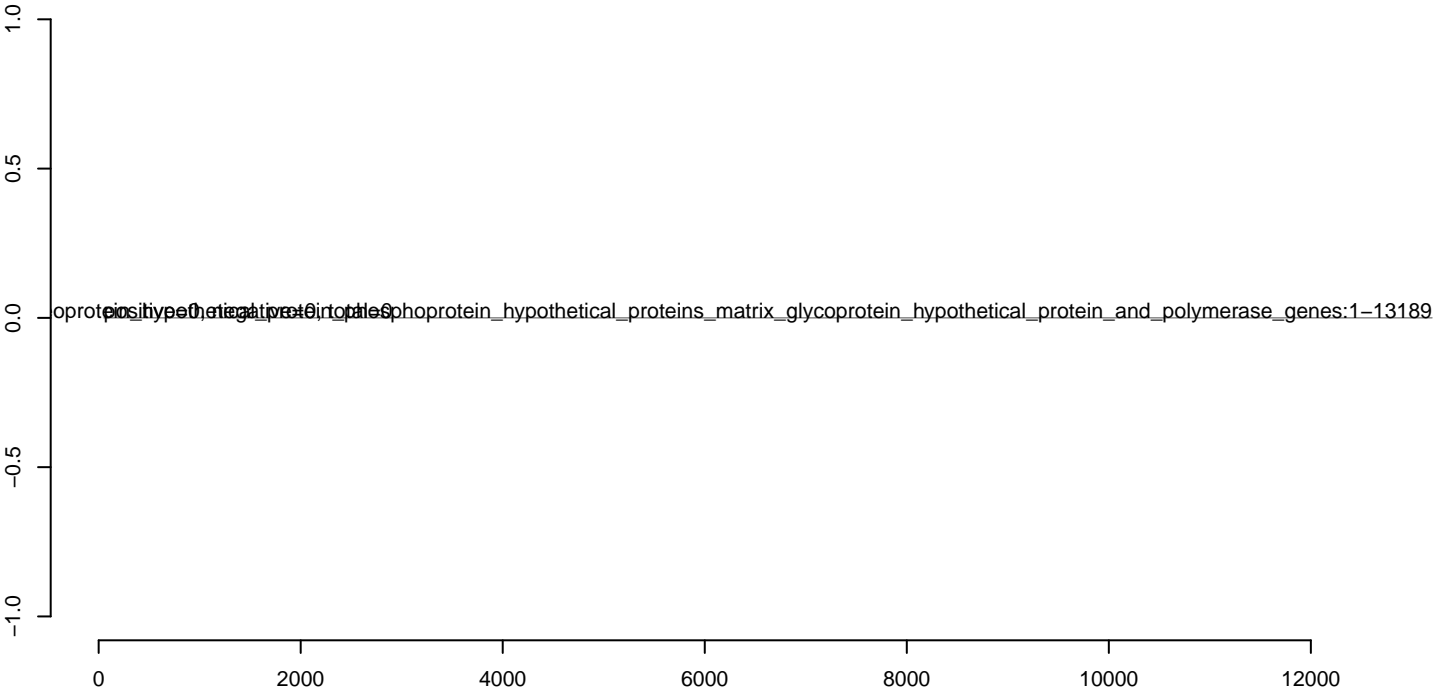
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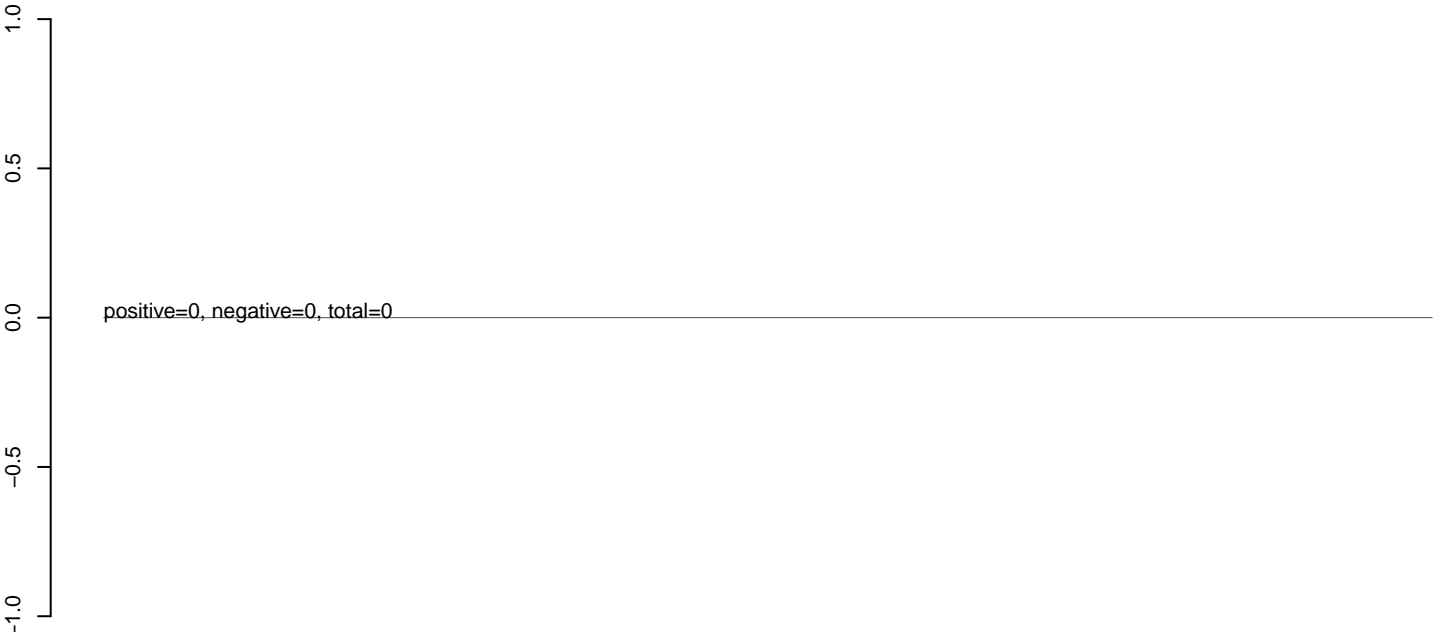
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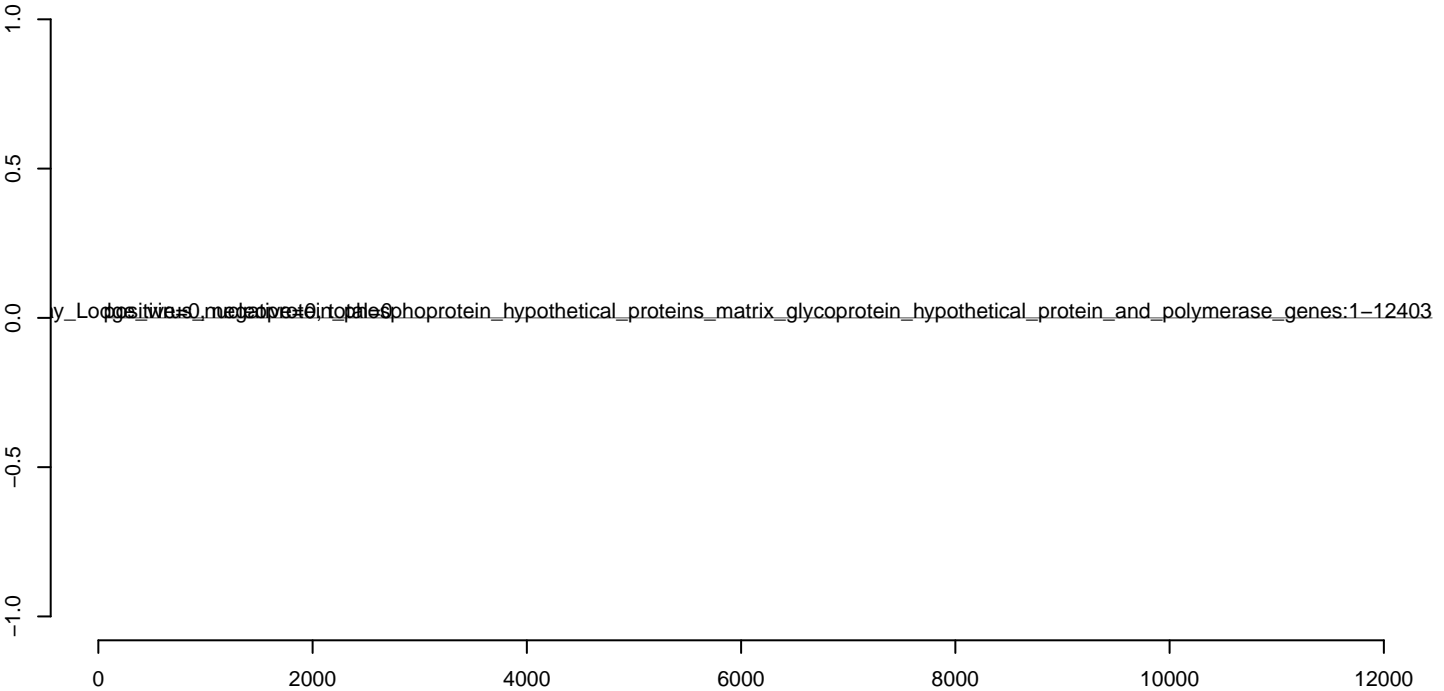
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**AnGam\_Sua5b\_cells.24\_35.rep**



**AnGam\_Sua5b\_cells.rep**



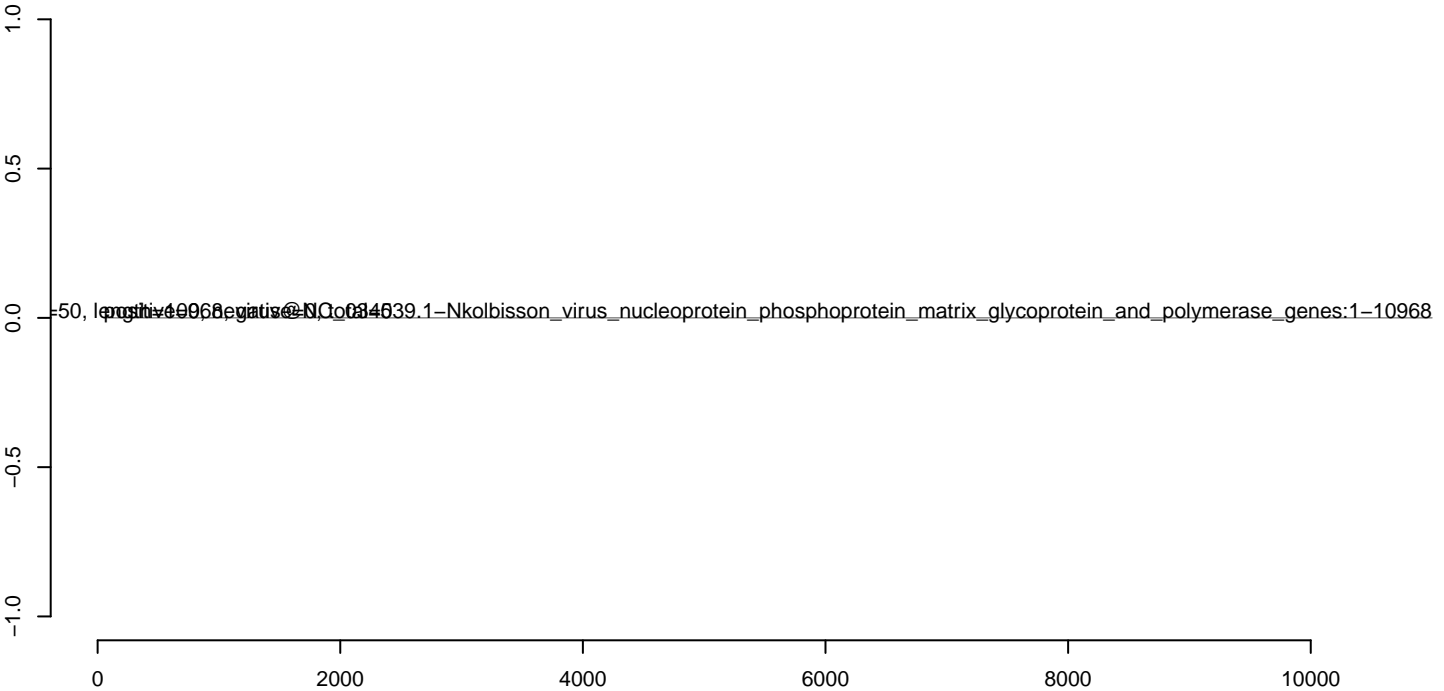
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AnGam\_Sua5b\_cells.rep



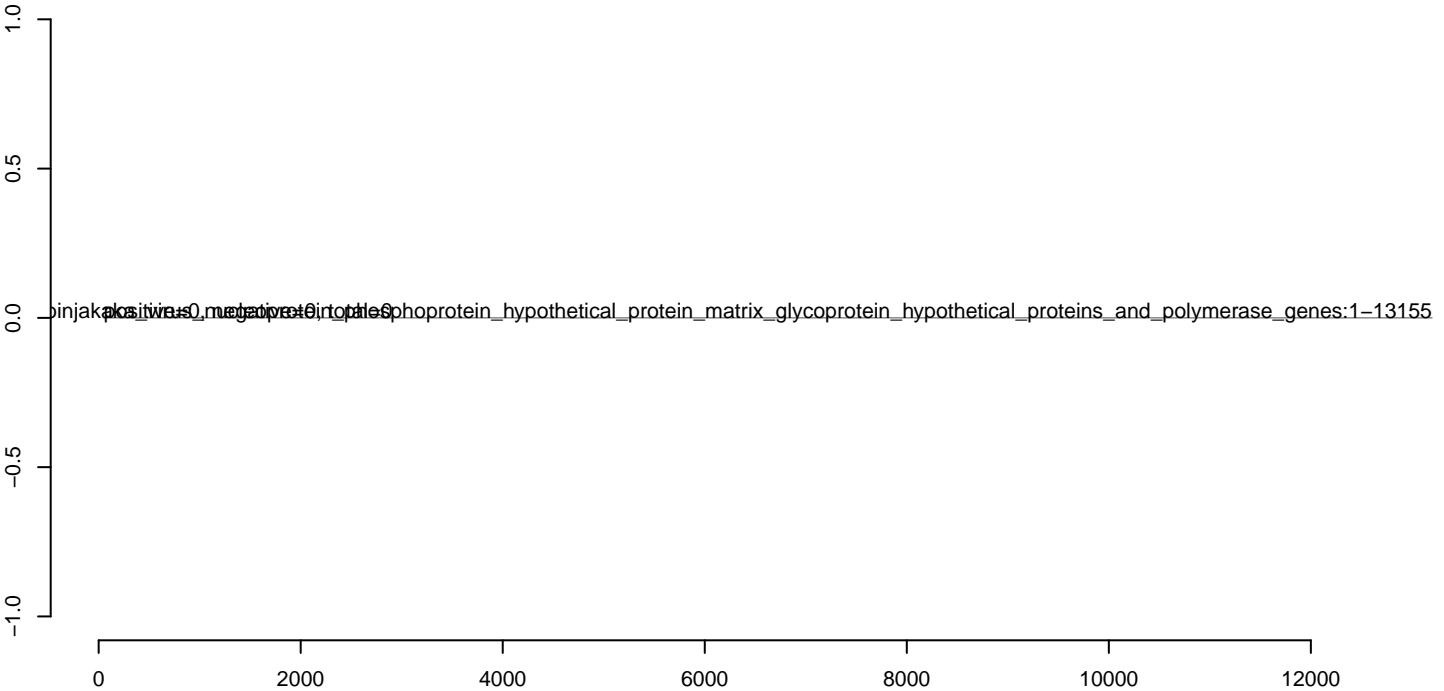
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AnGam\_Sua5b\_cells.rep





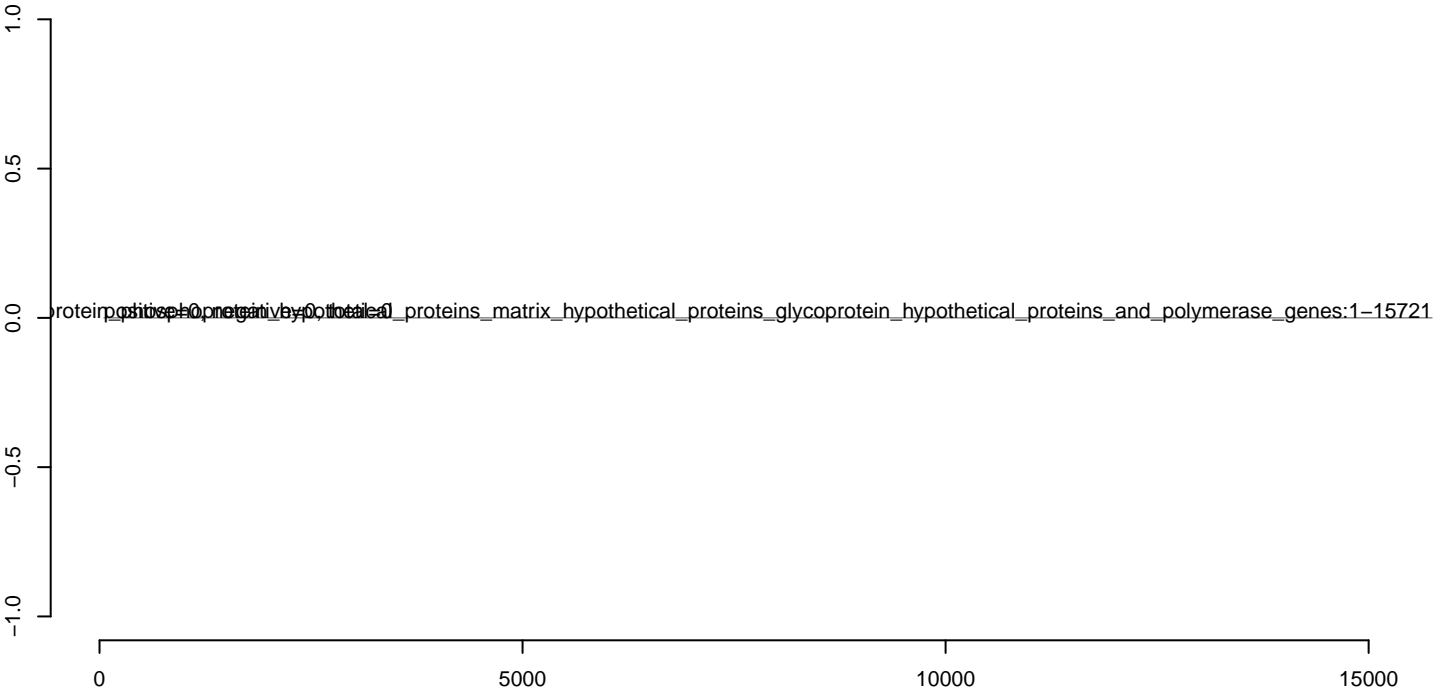
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AnGam\_Sua5b\_cells.rep



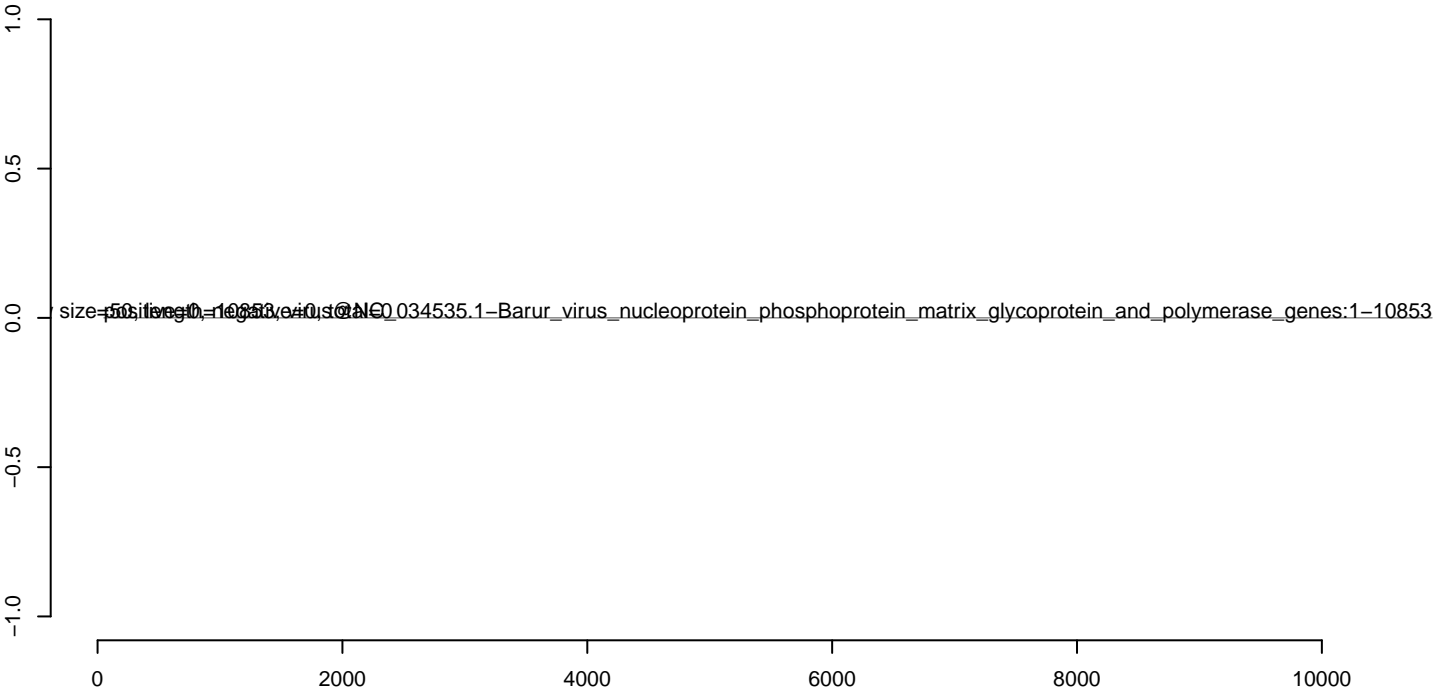
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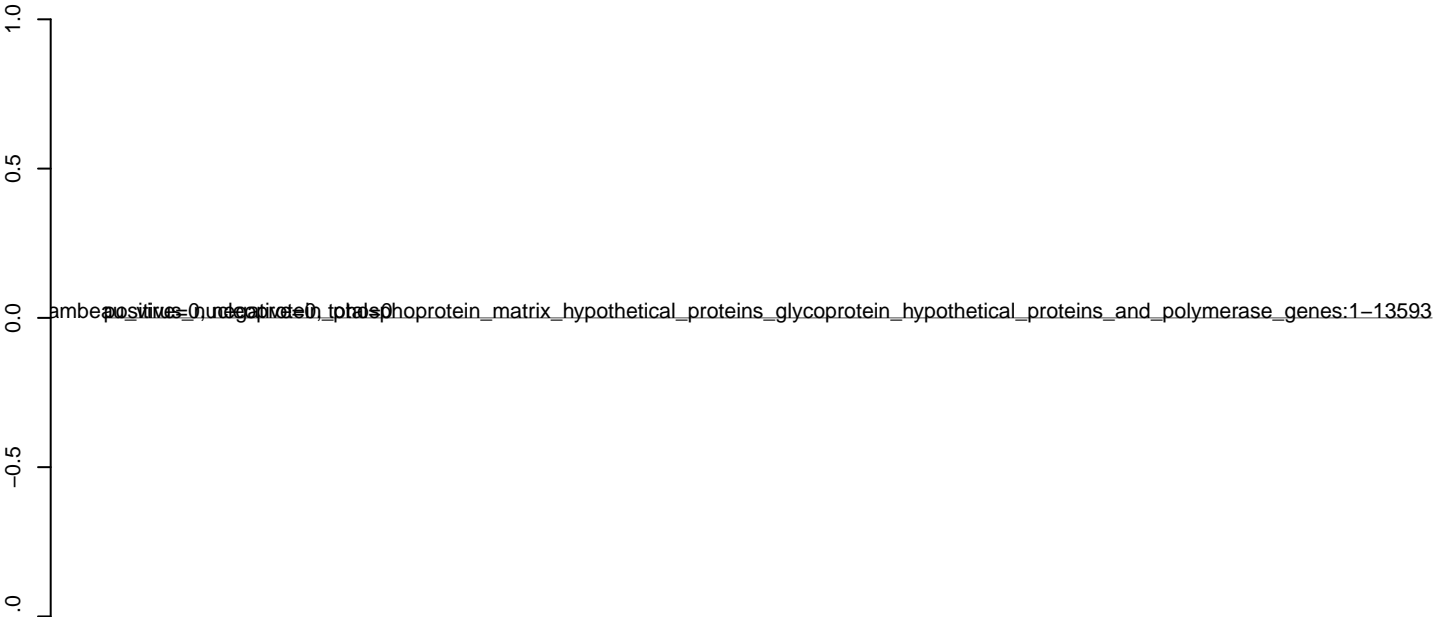
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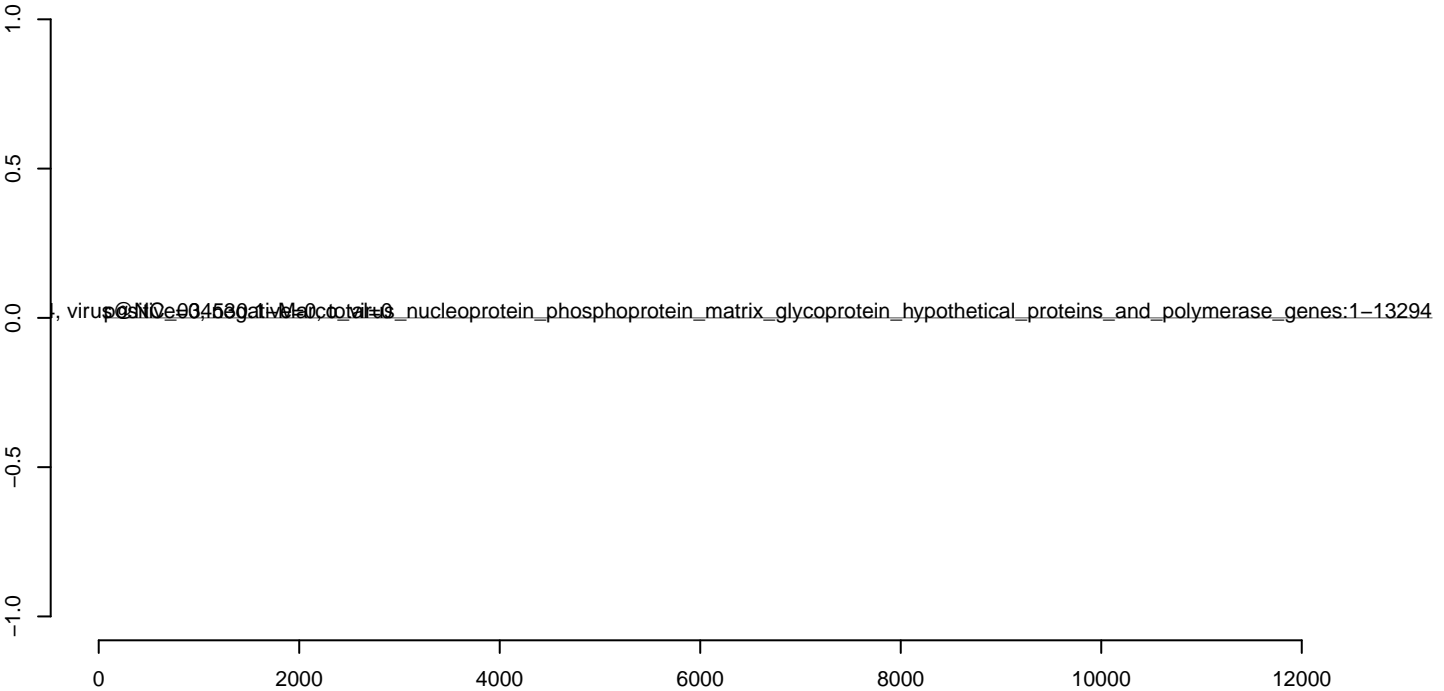
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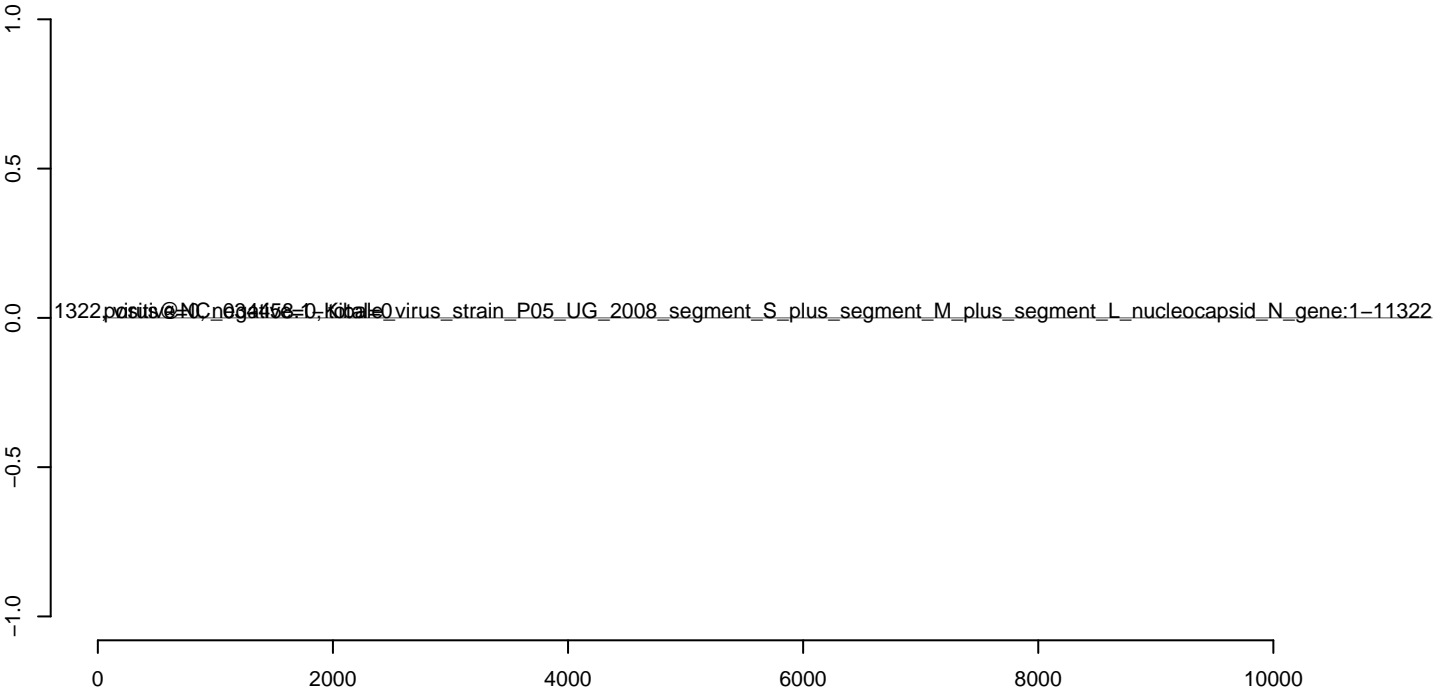
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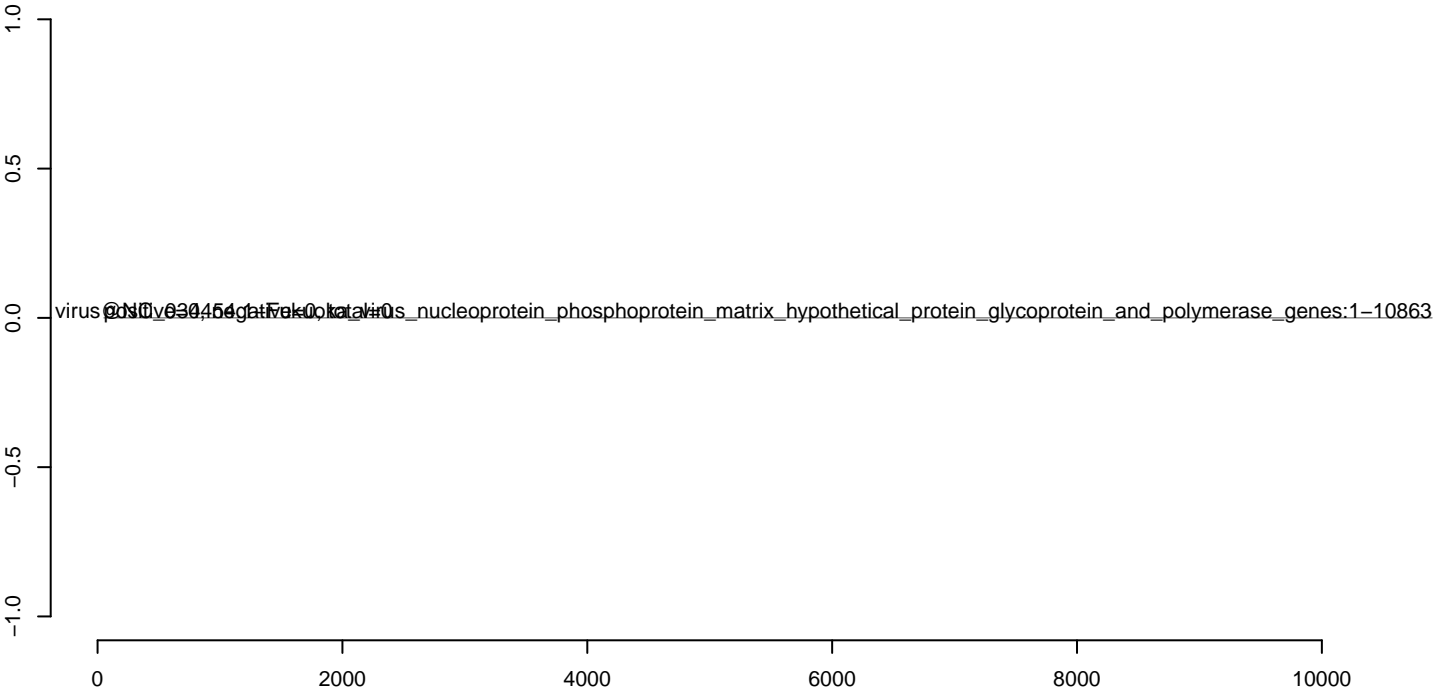
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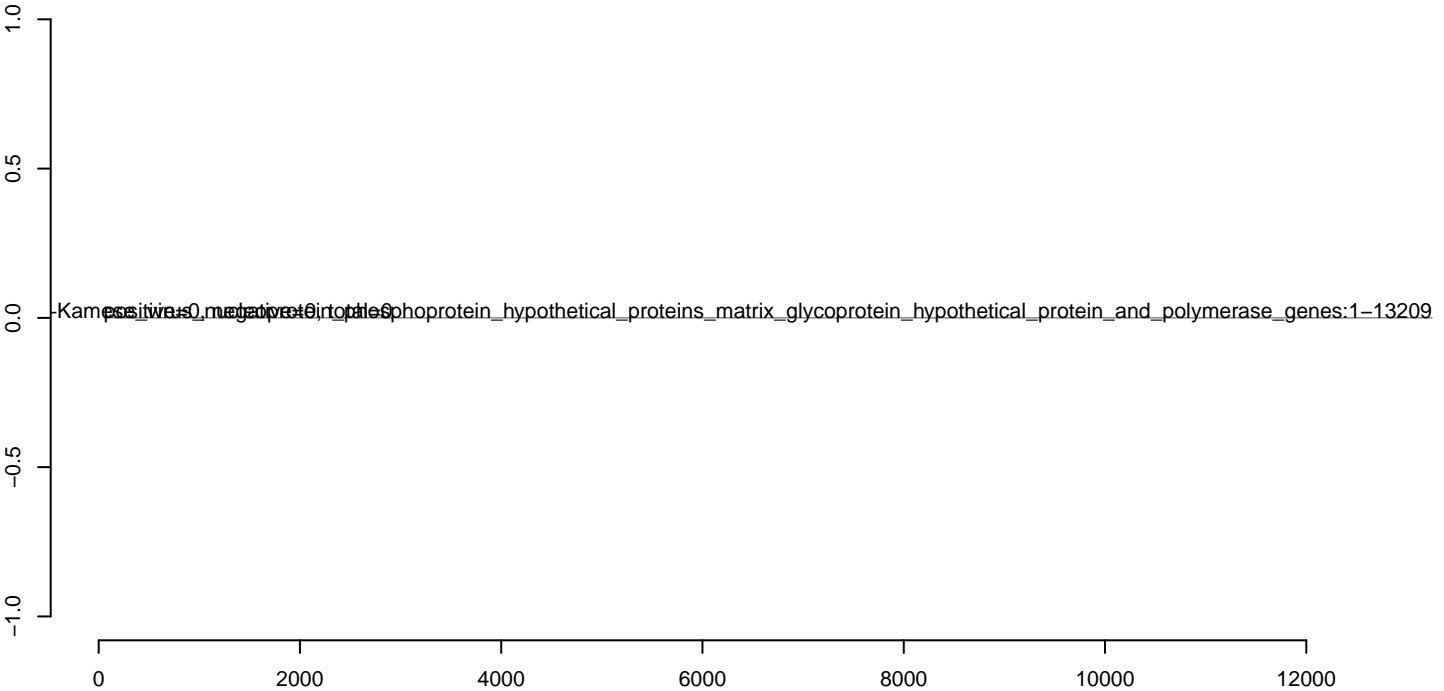
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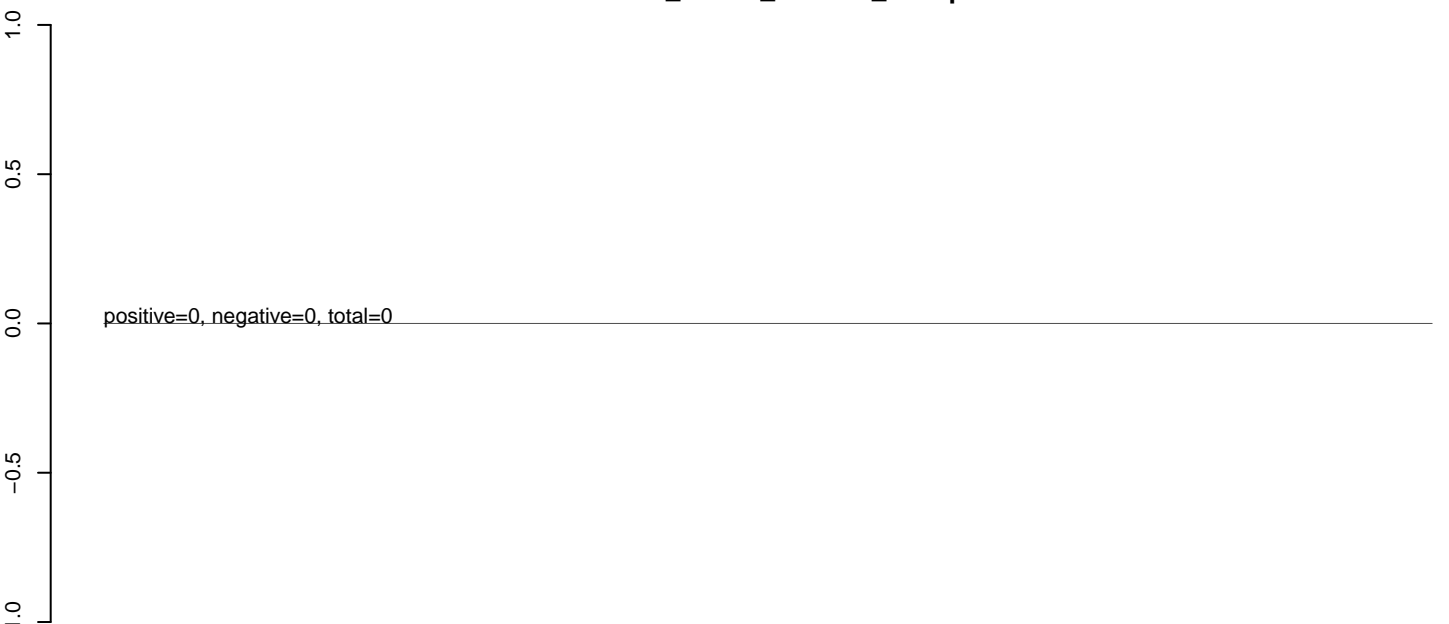
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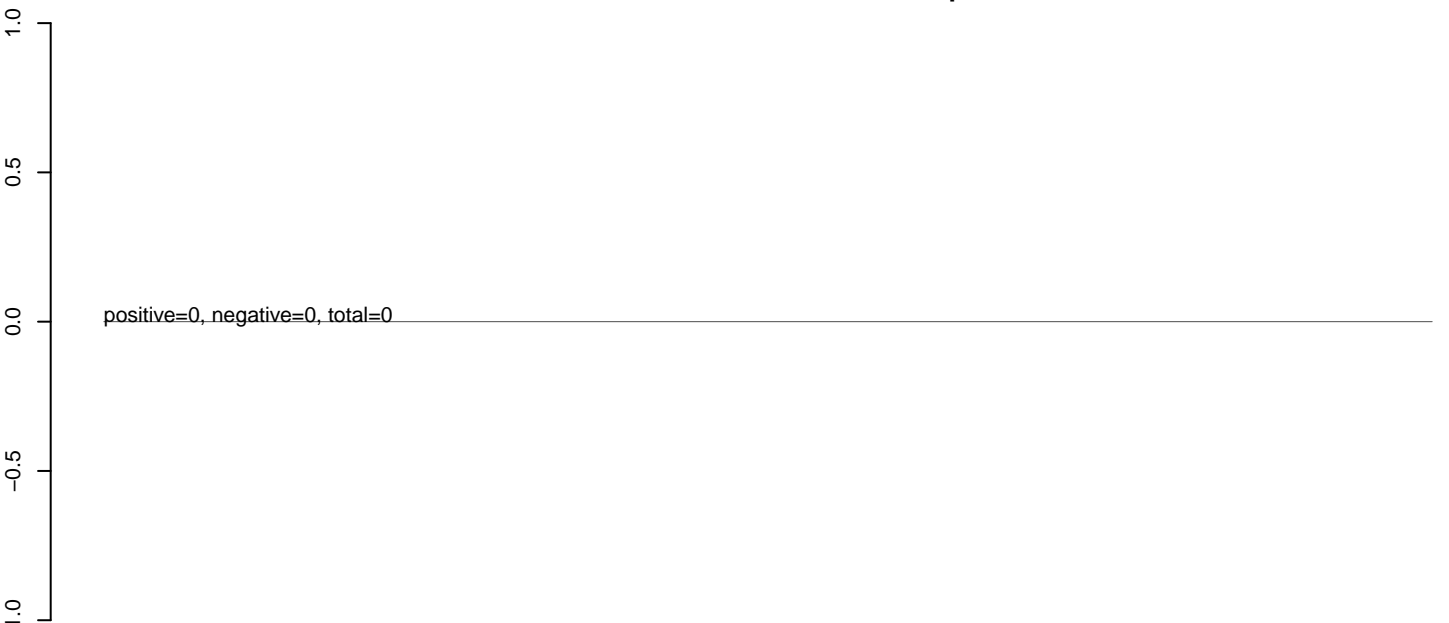
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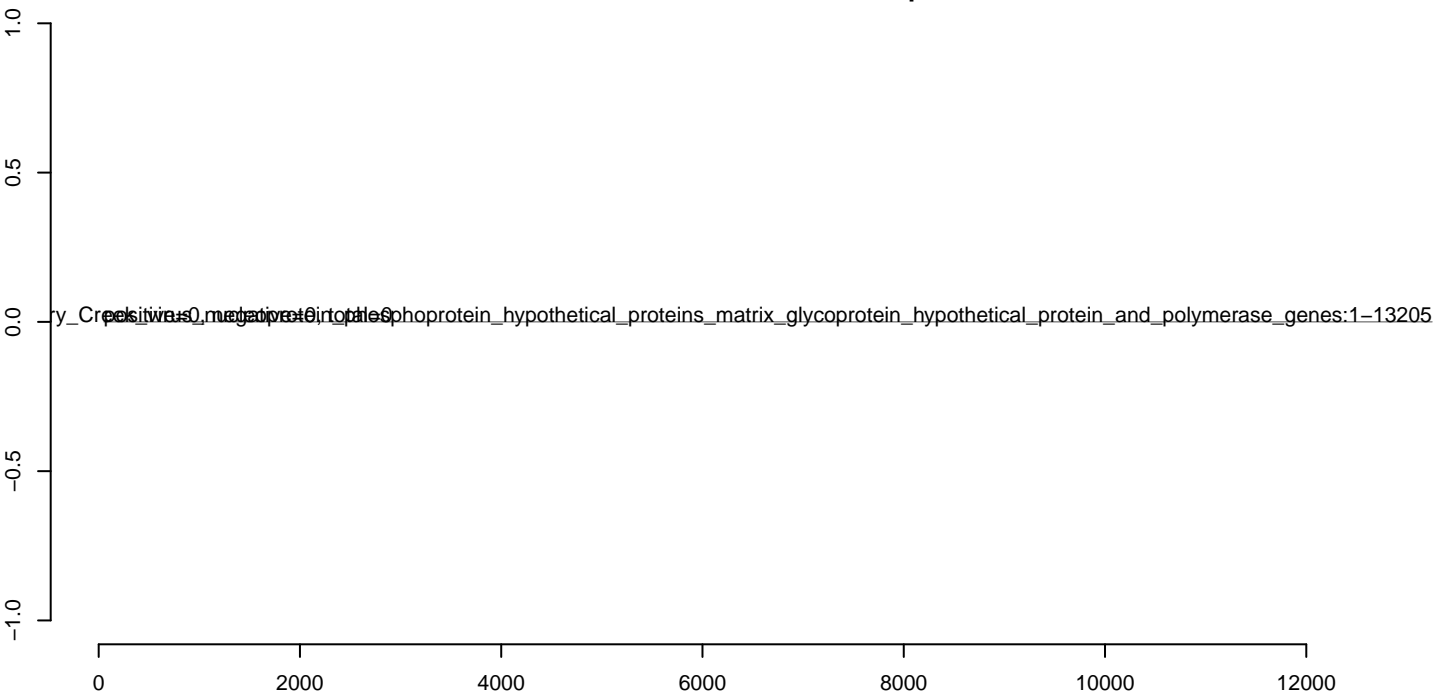
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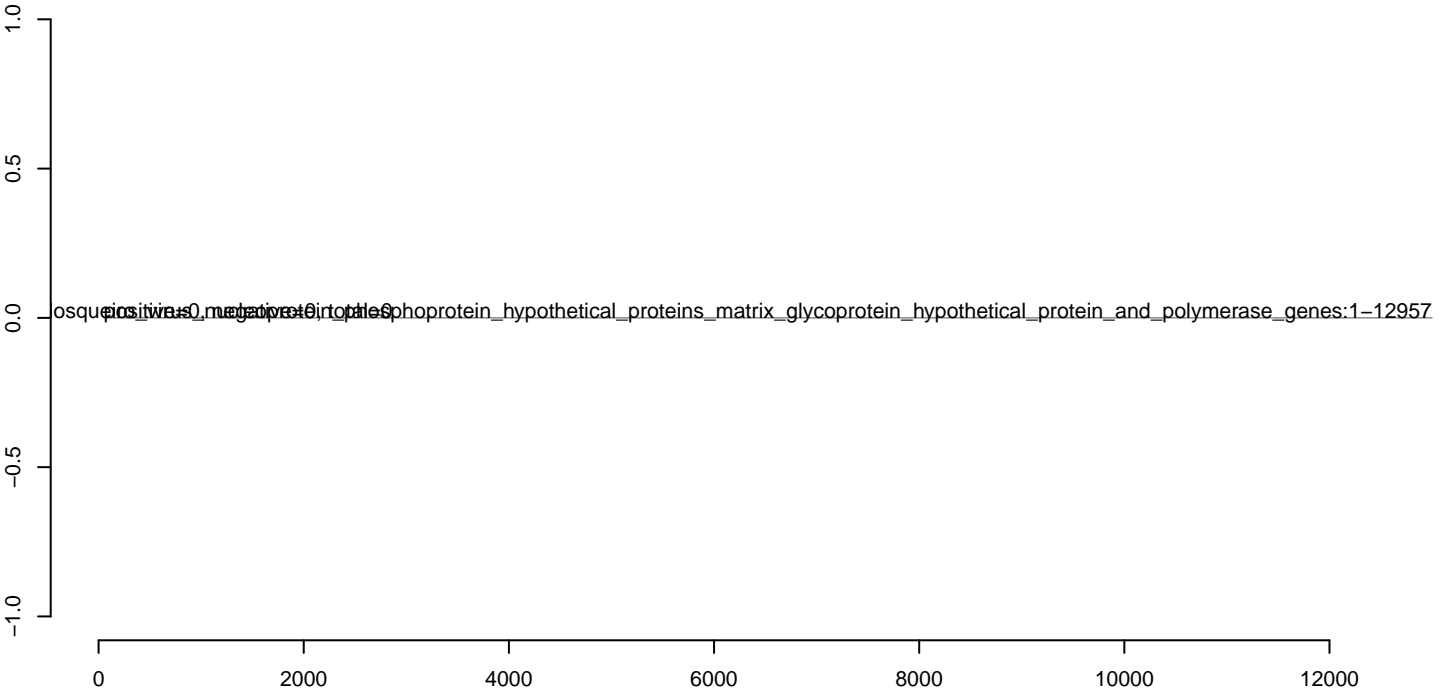
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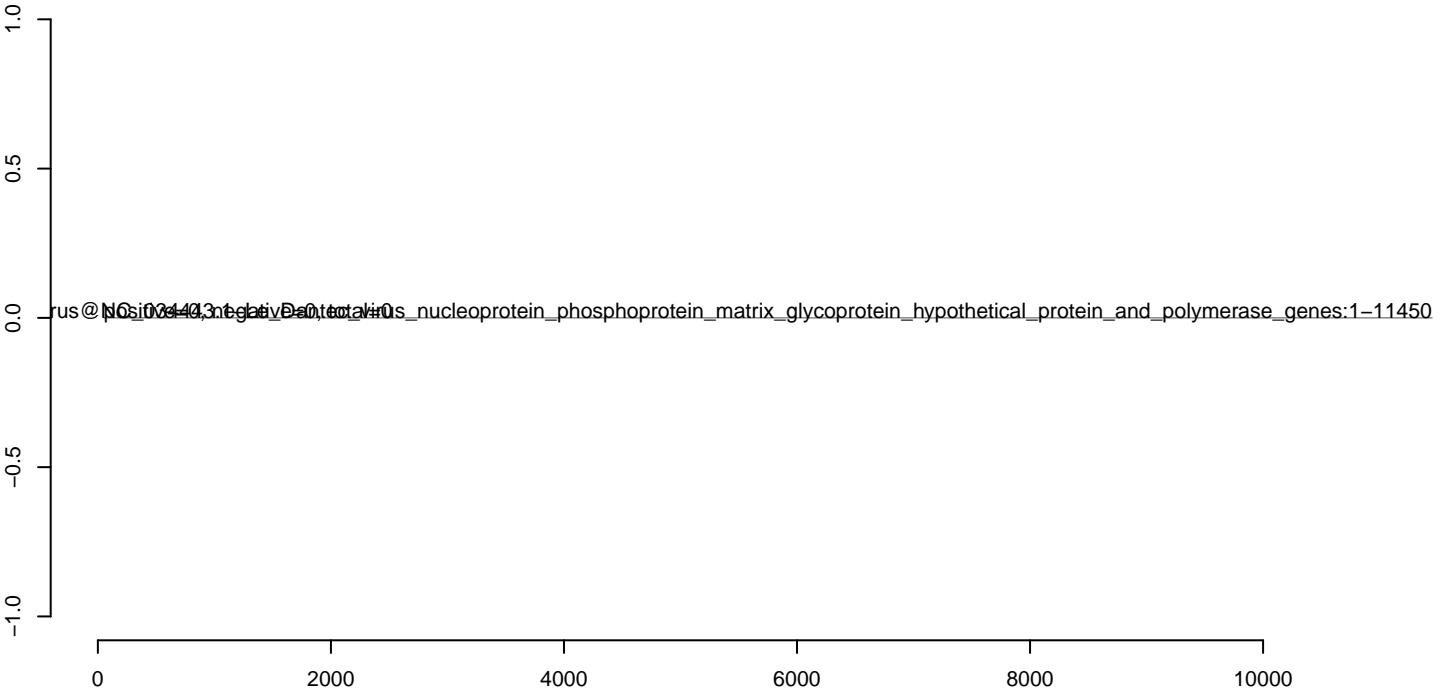
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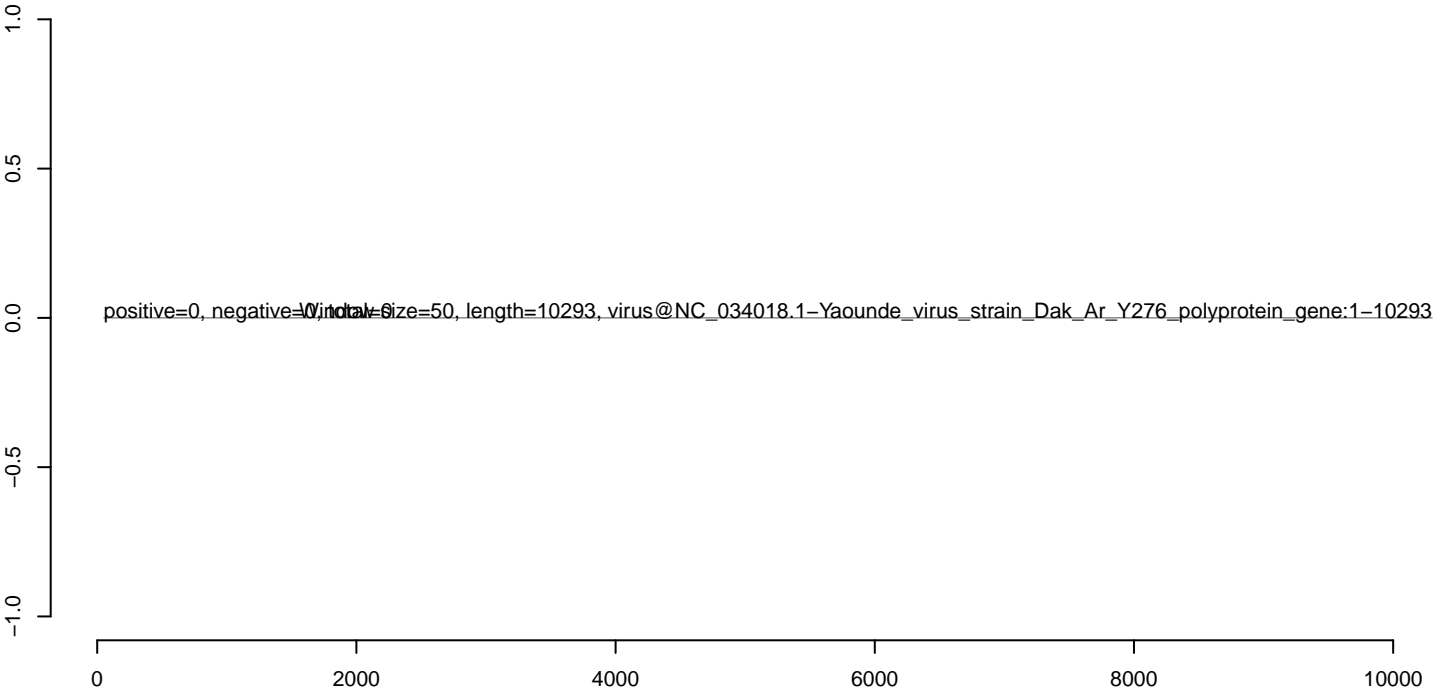
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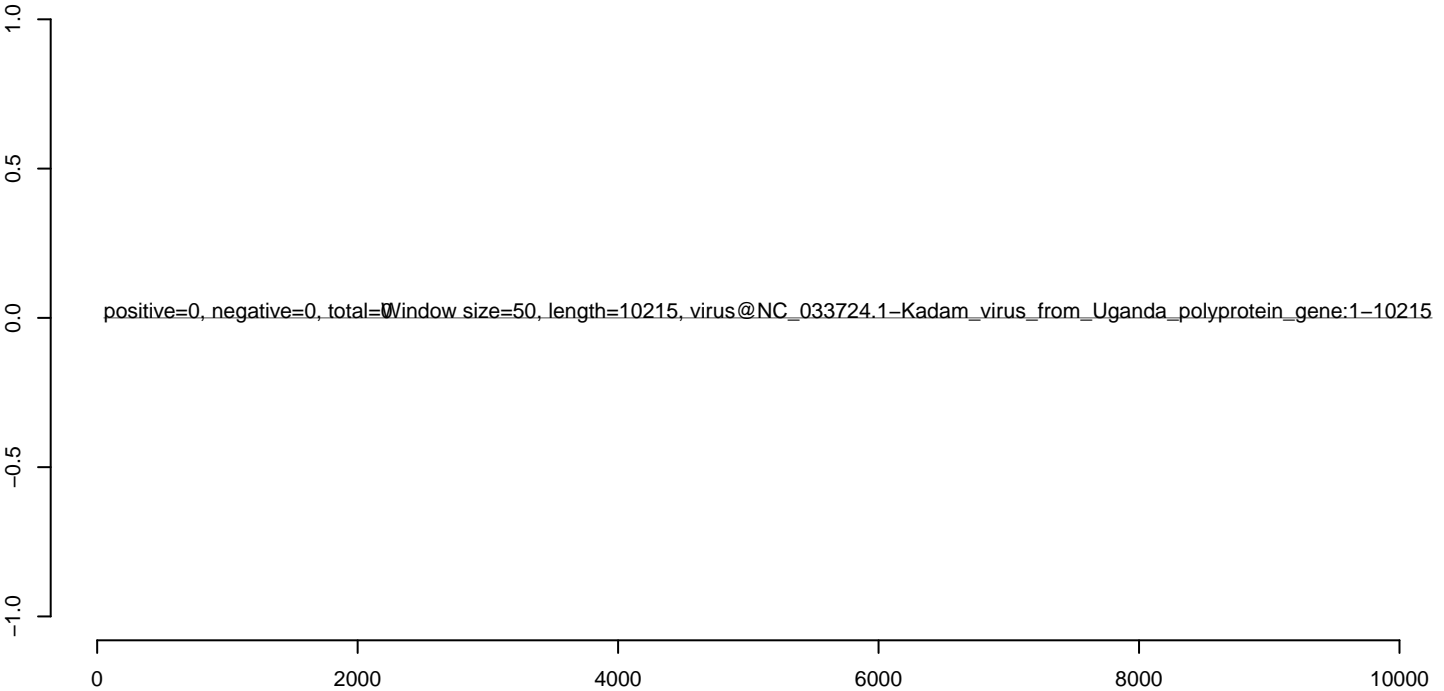
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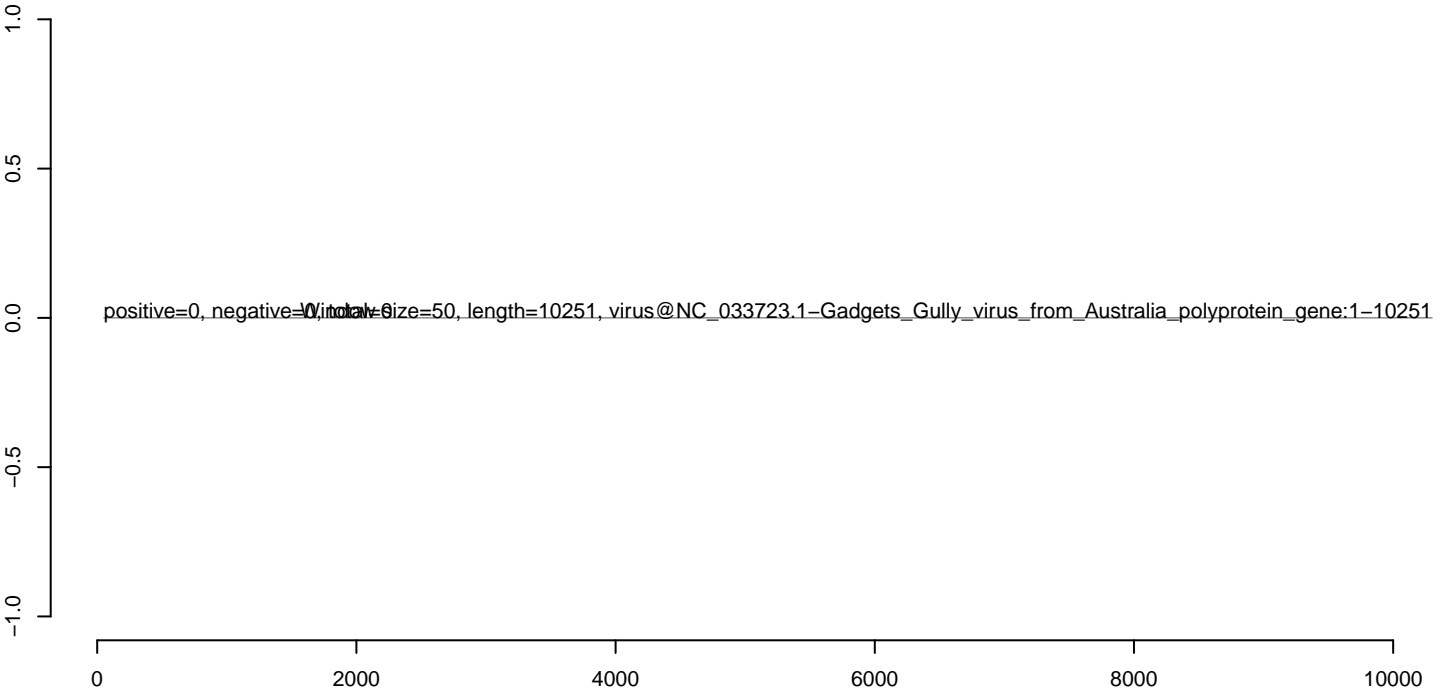
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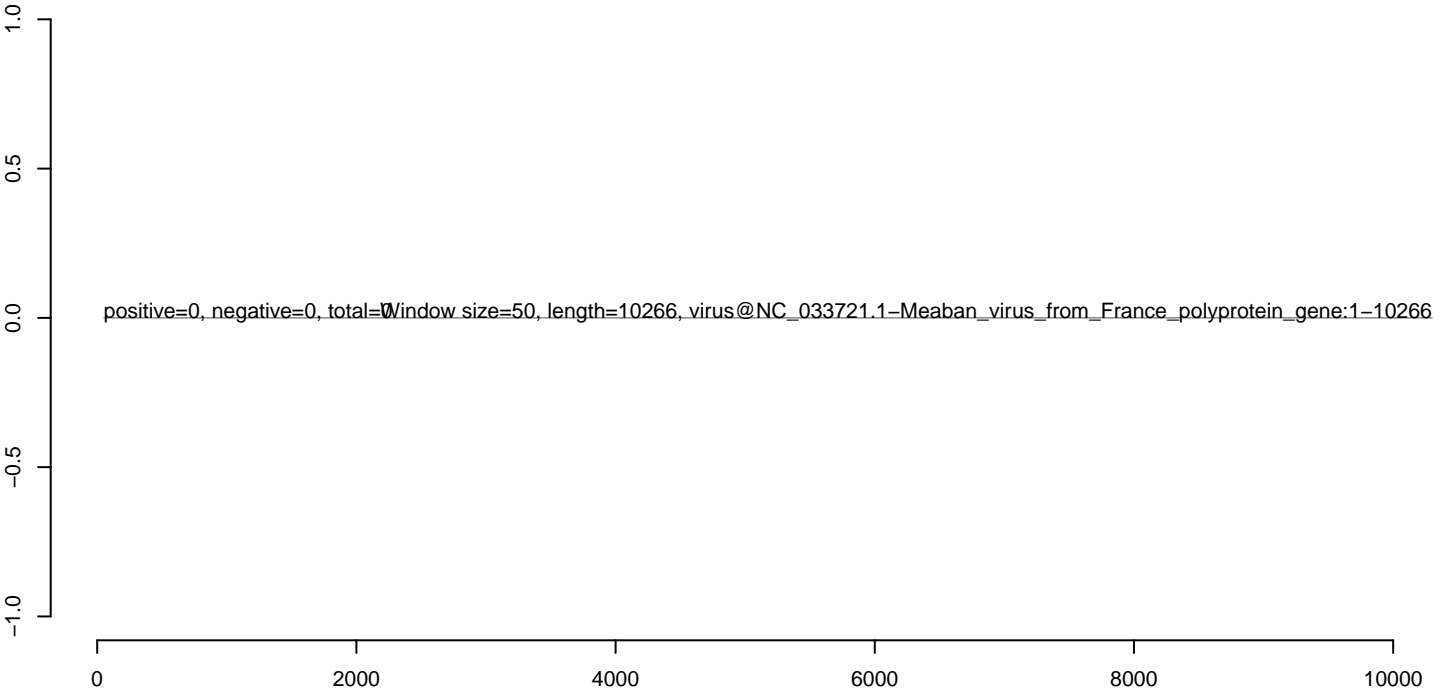
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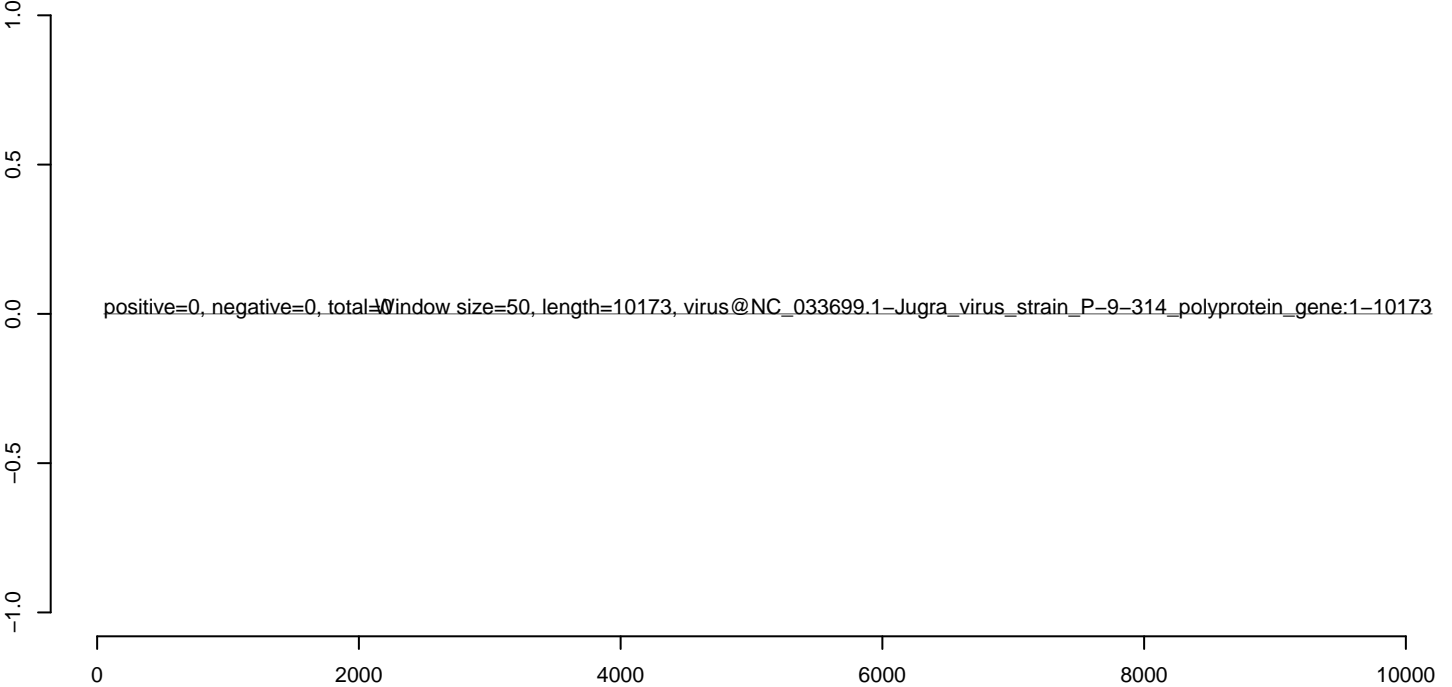
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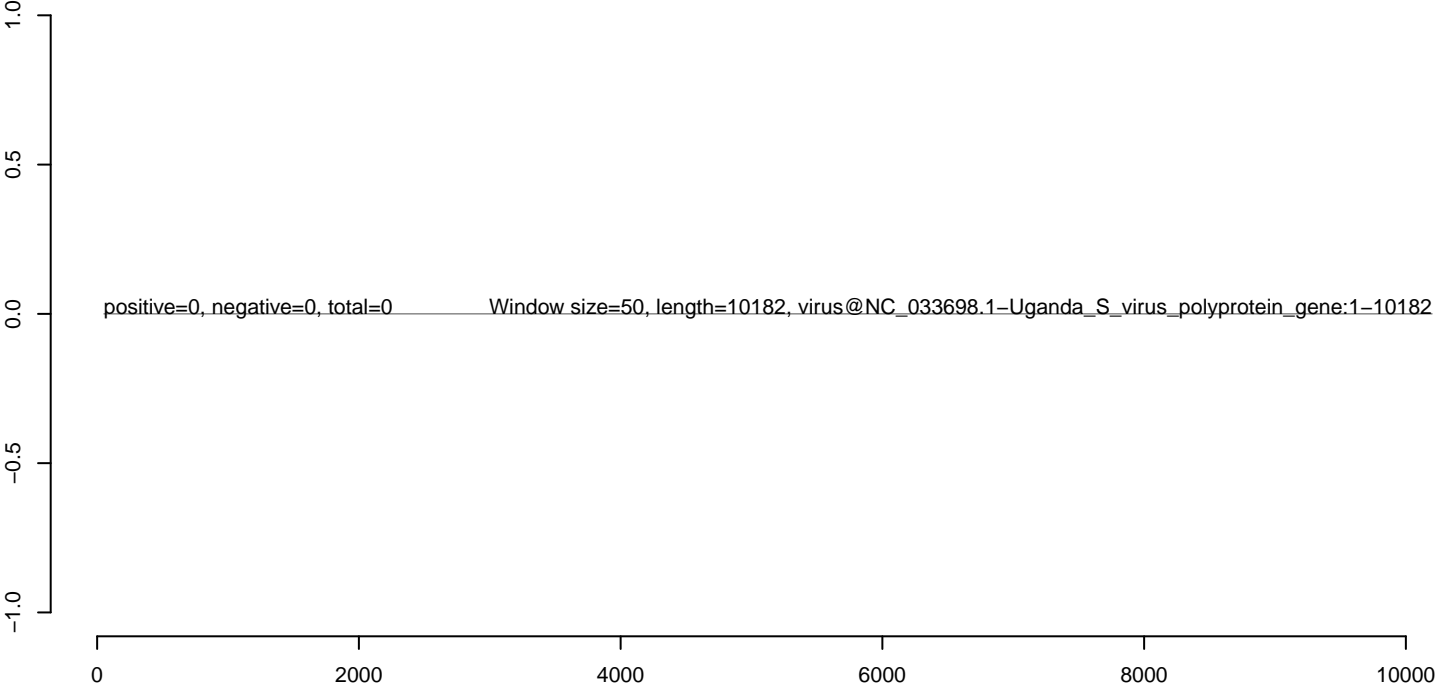
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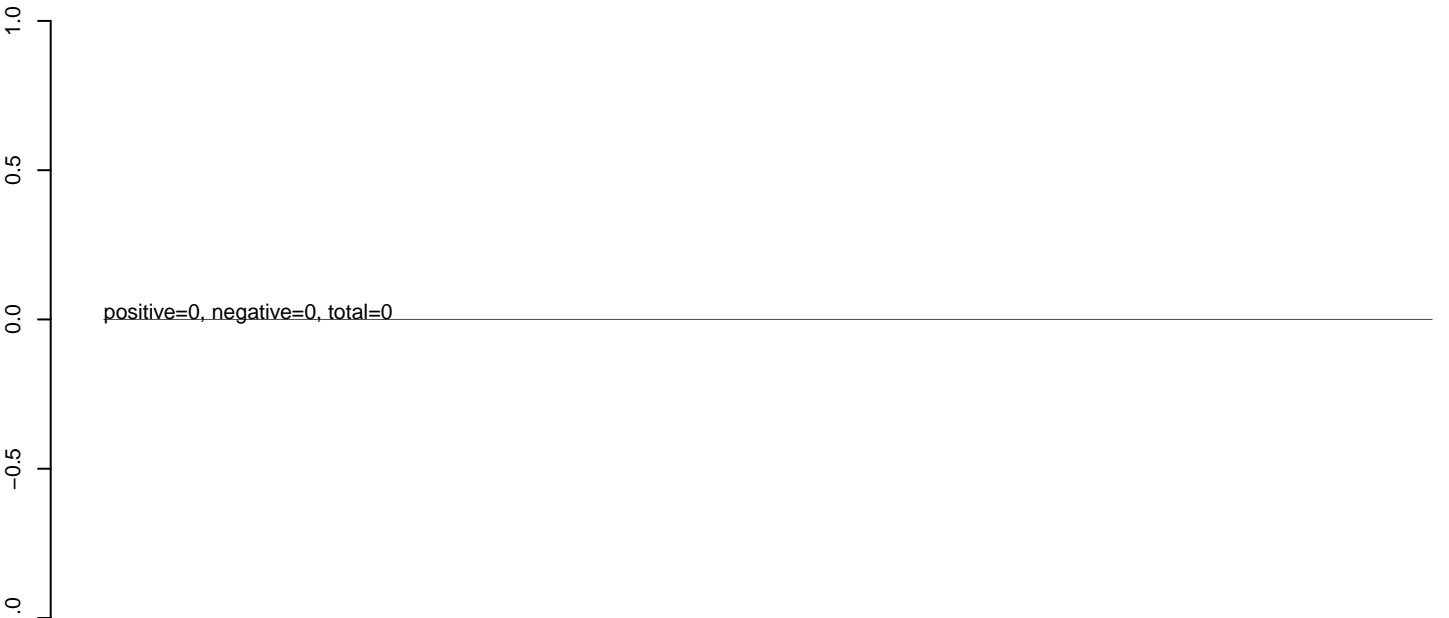


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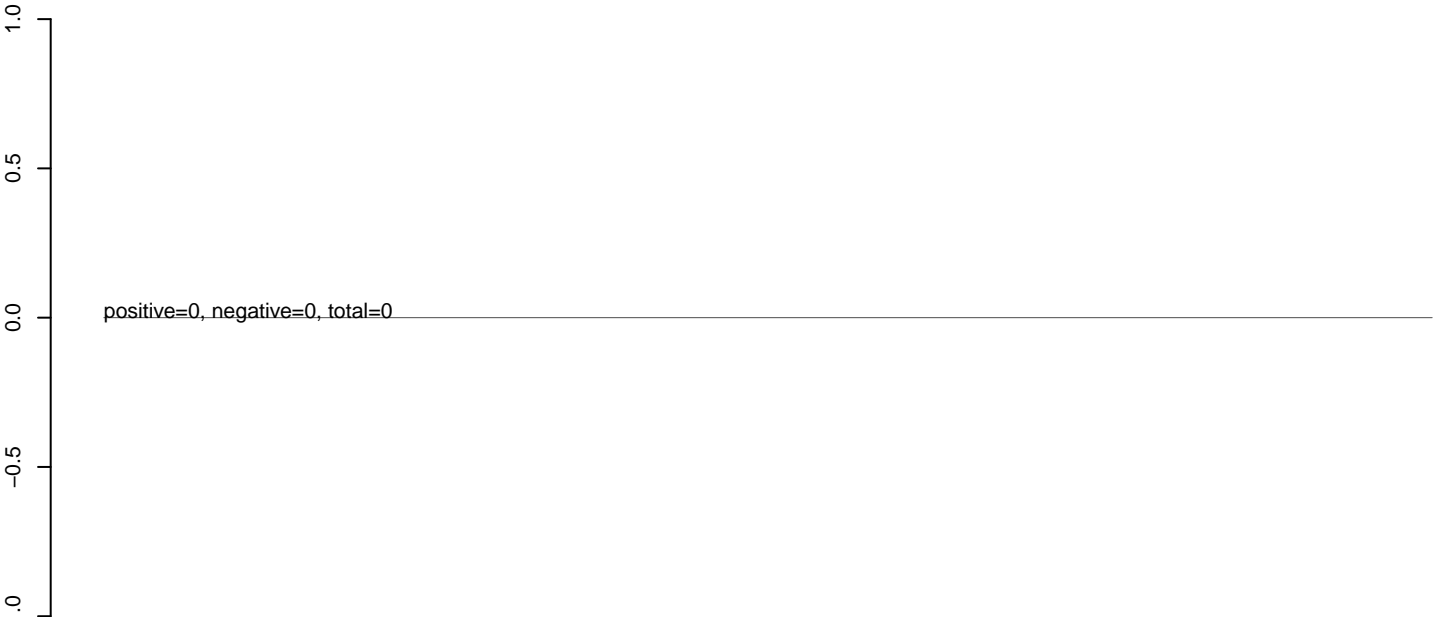




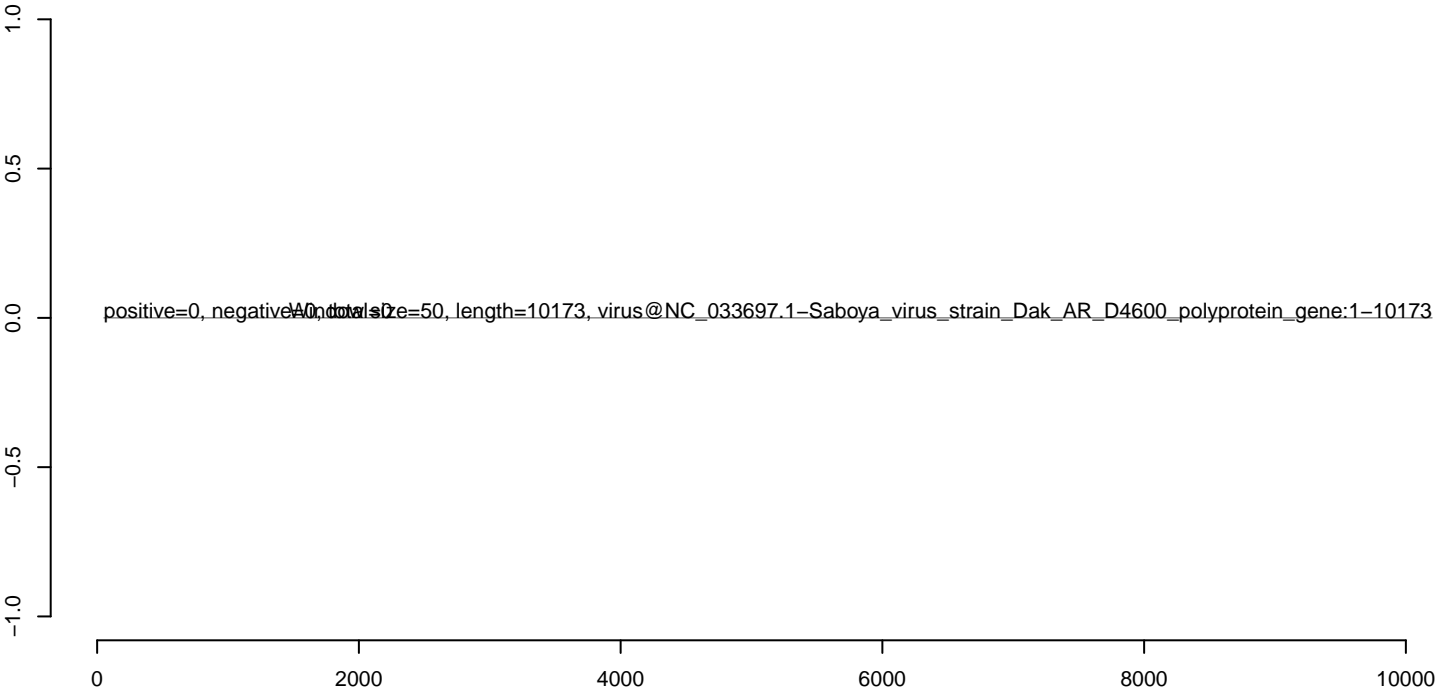
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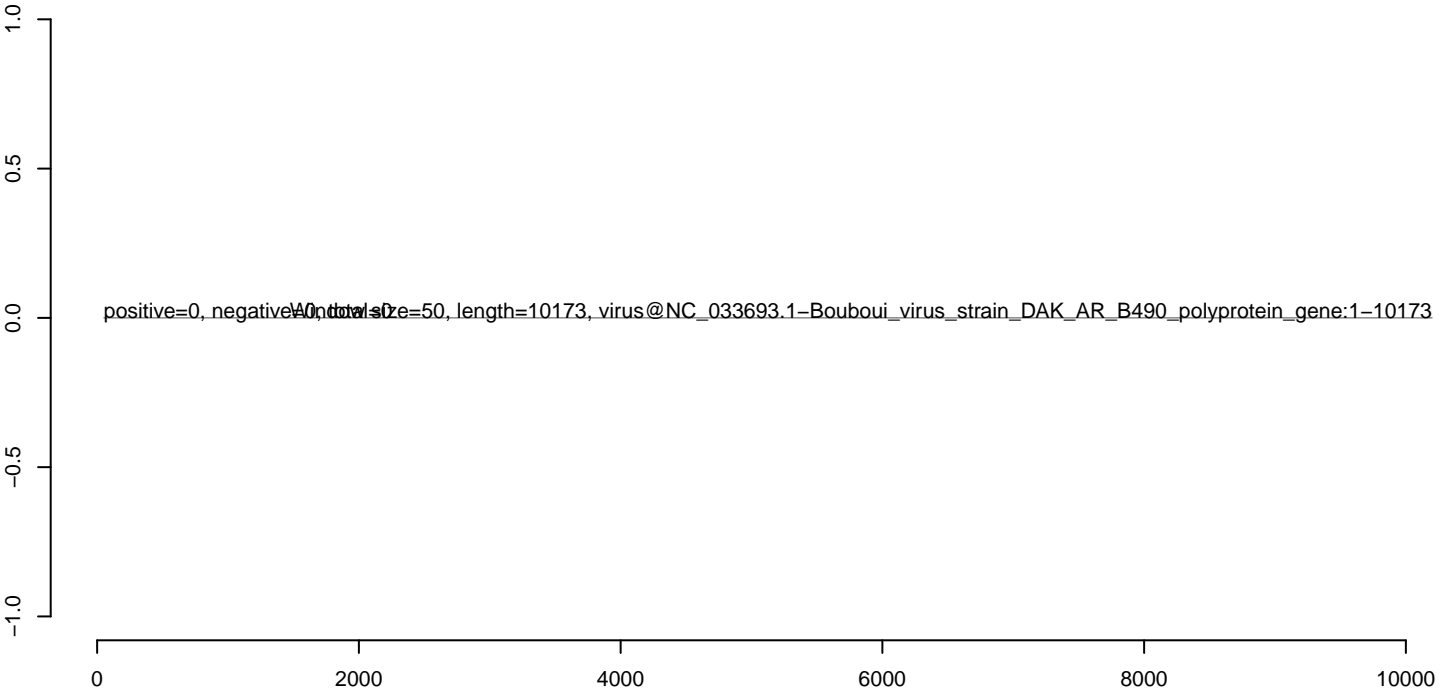
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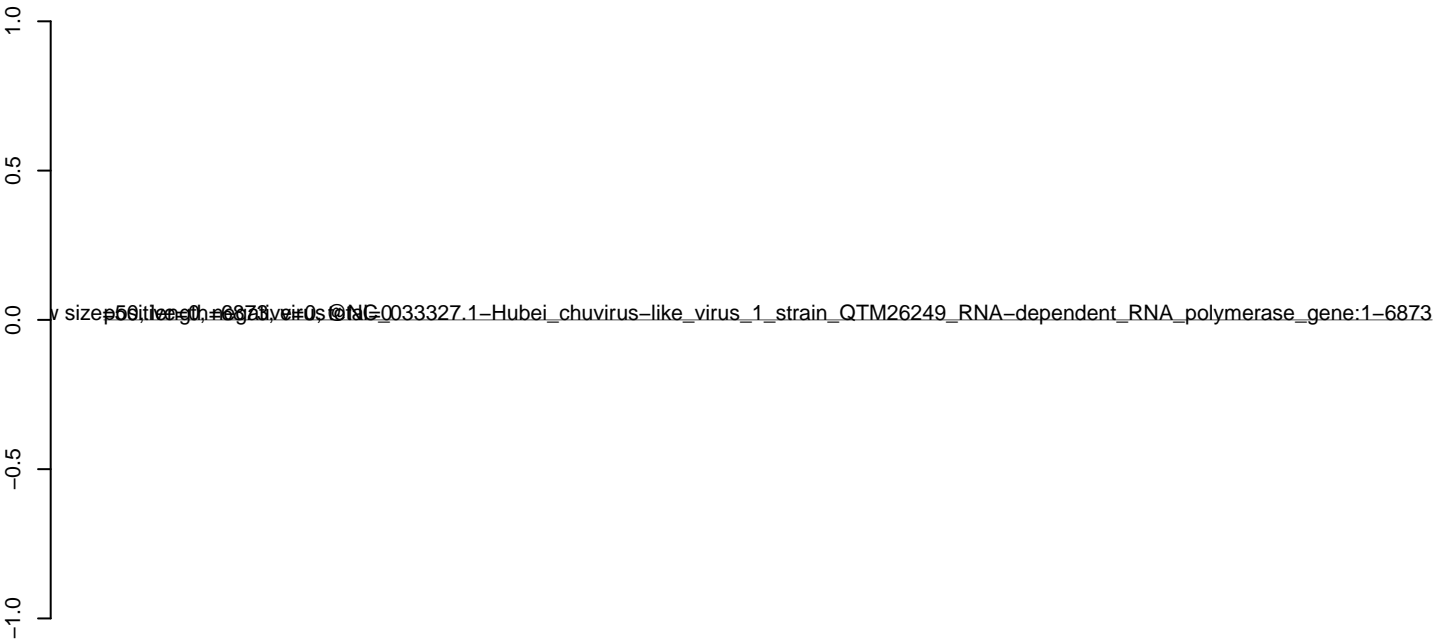
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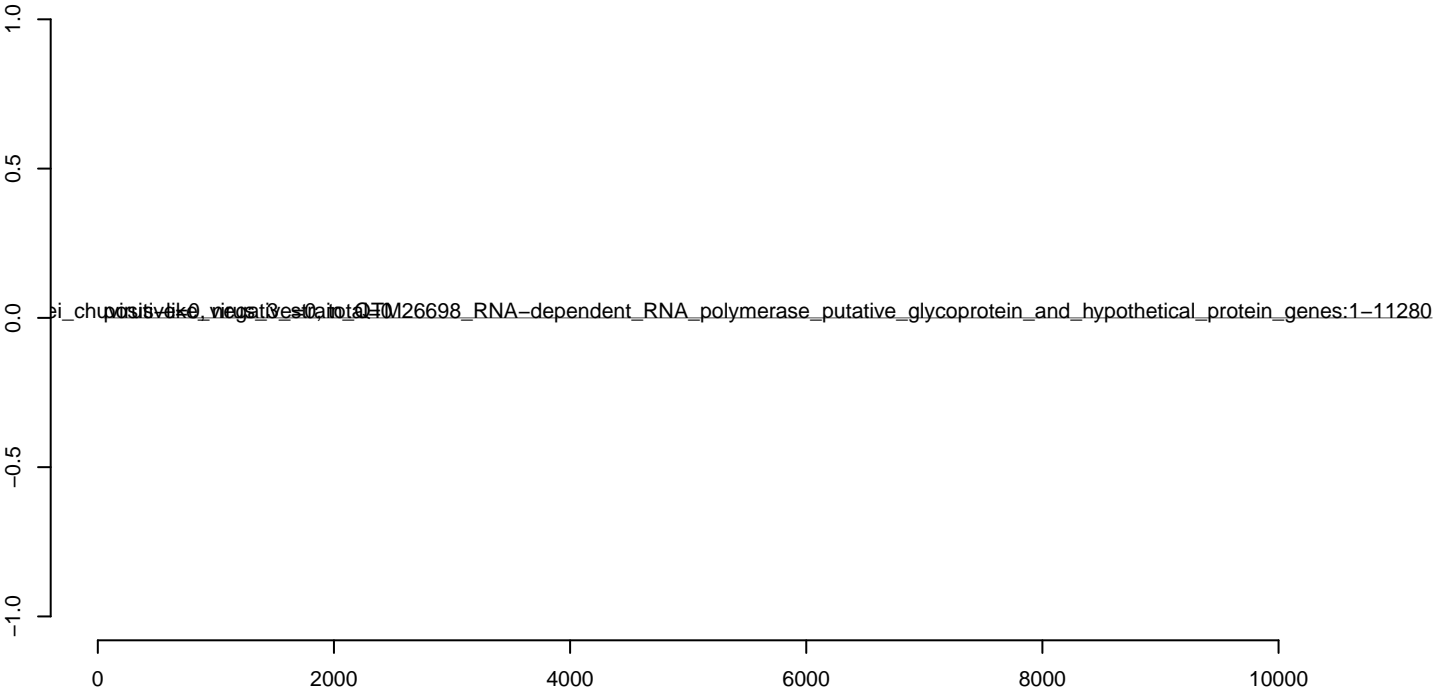
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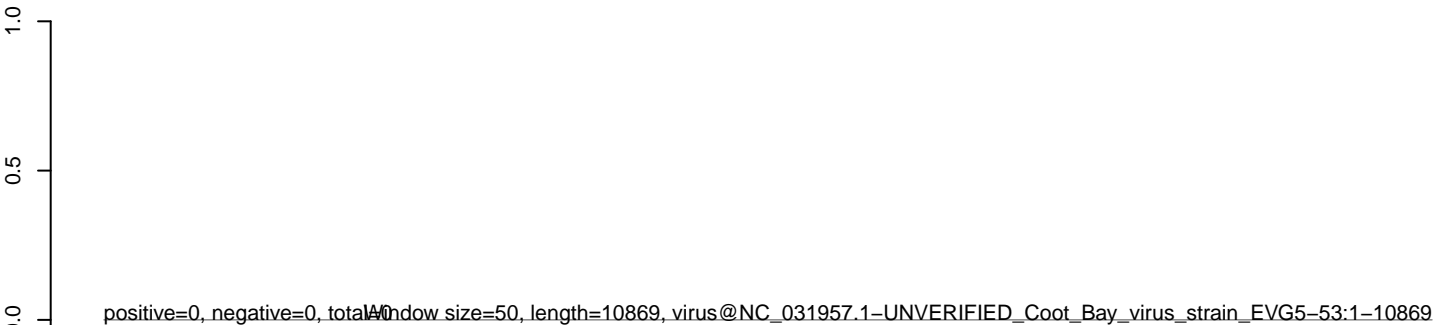
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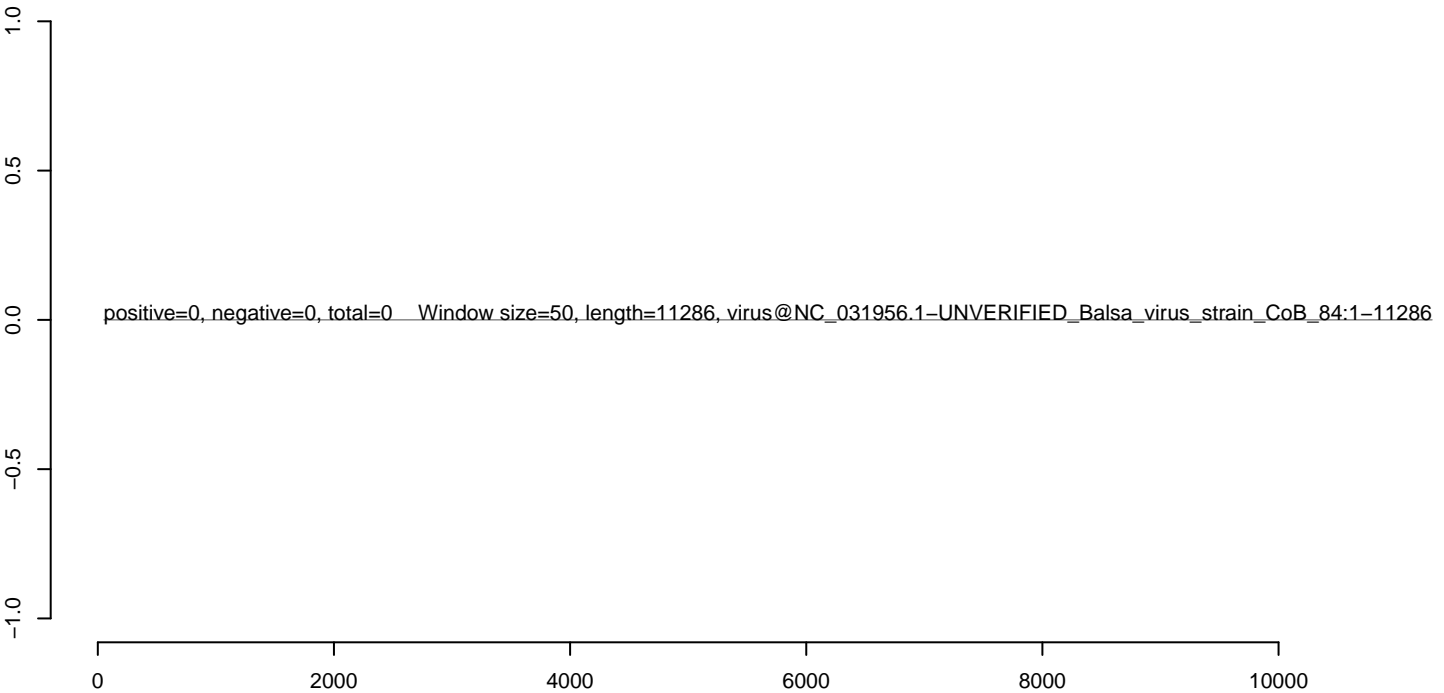
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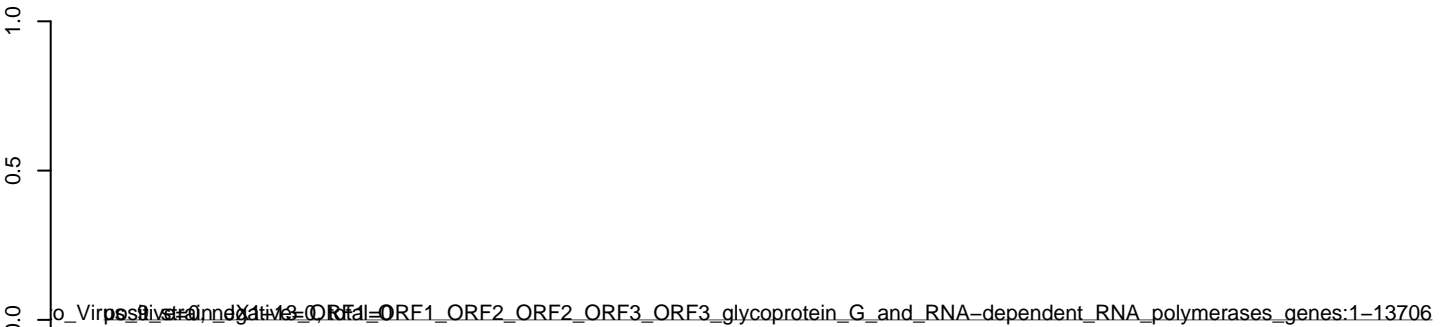
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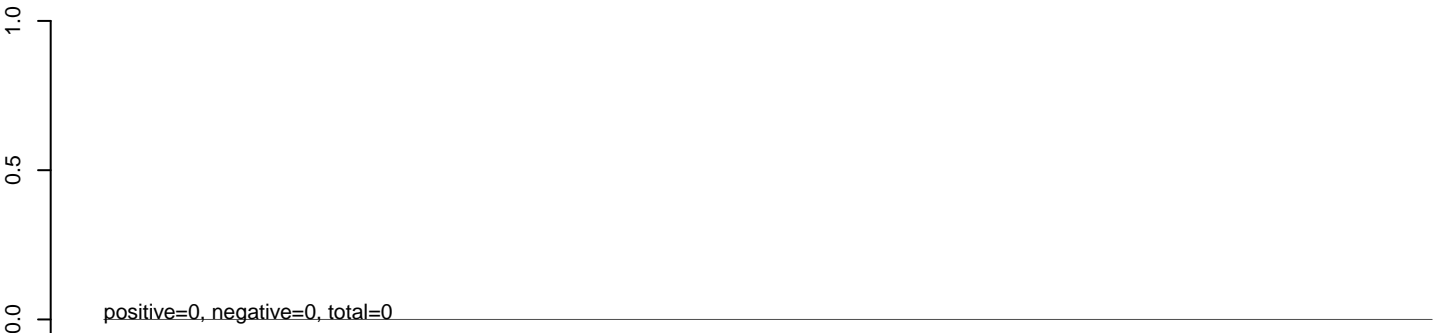


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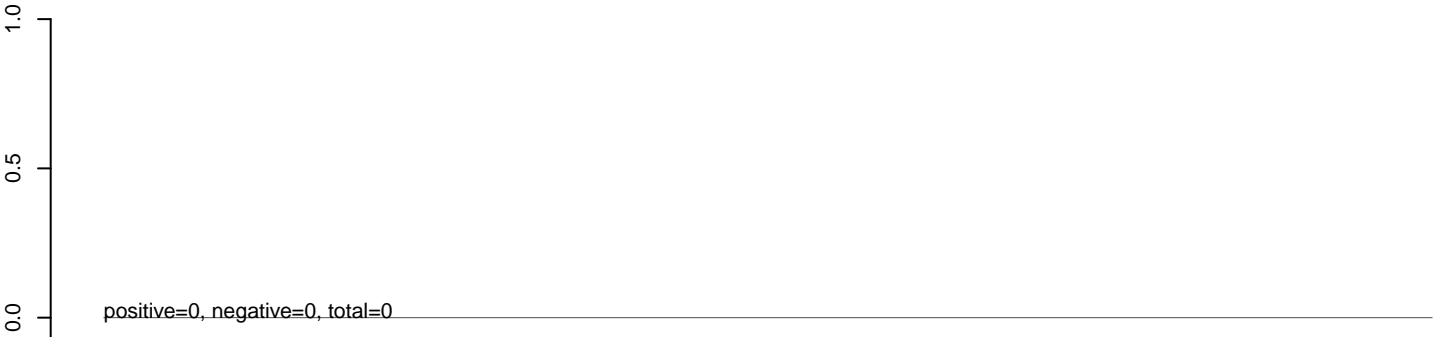


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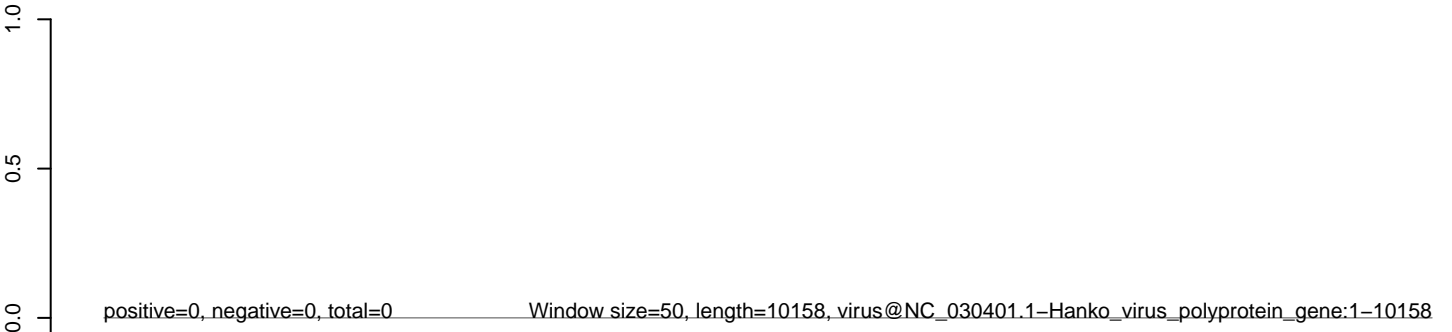
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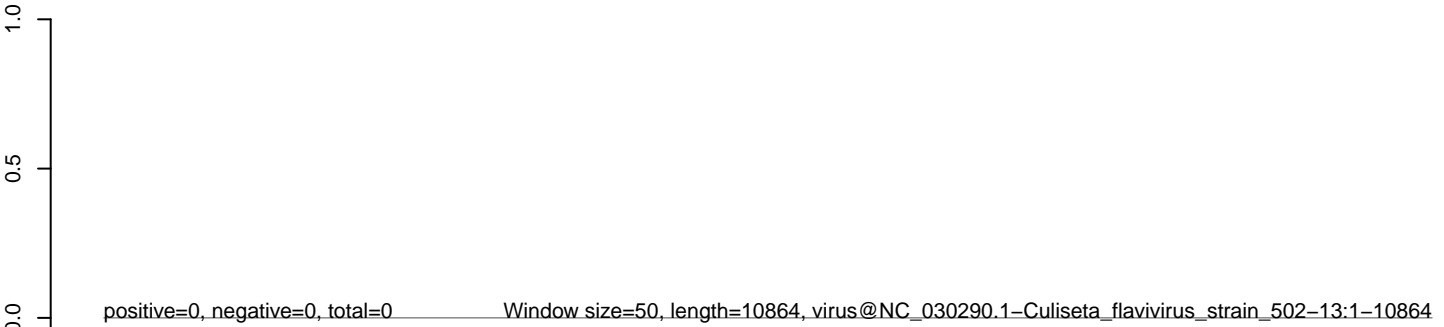
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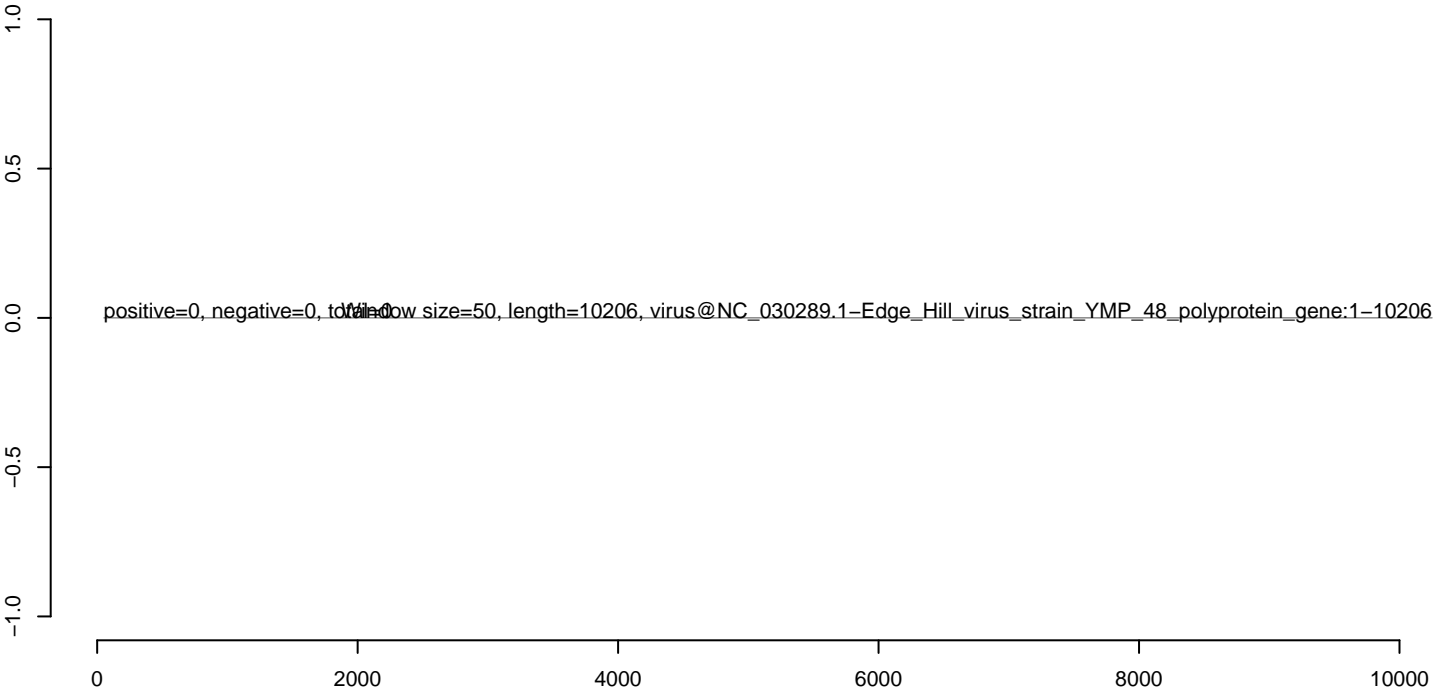
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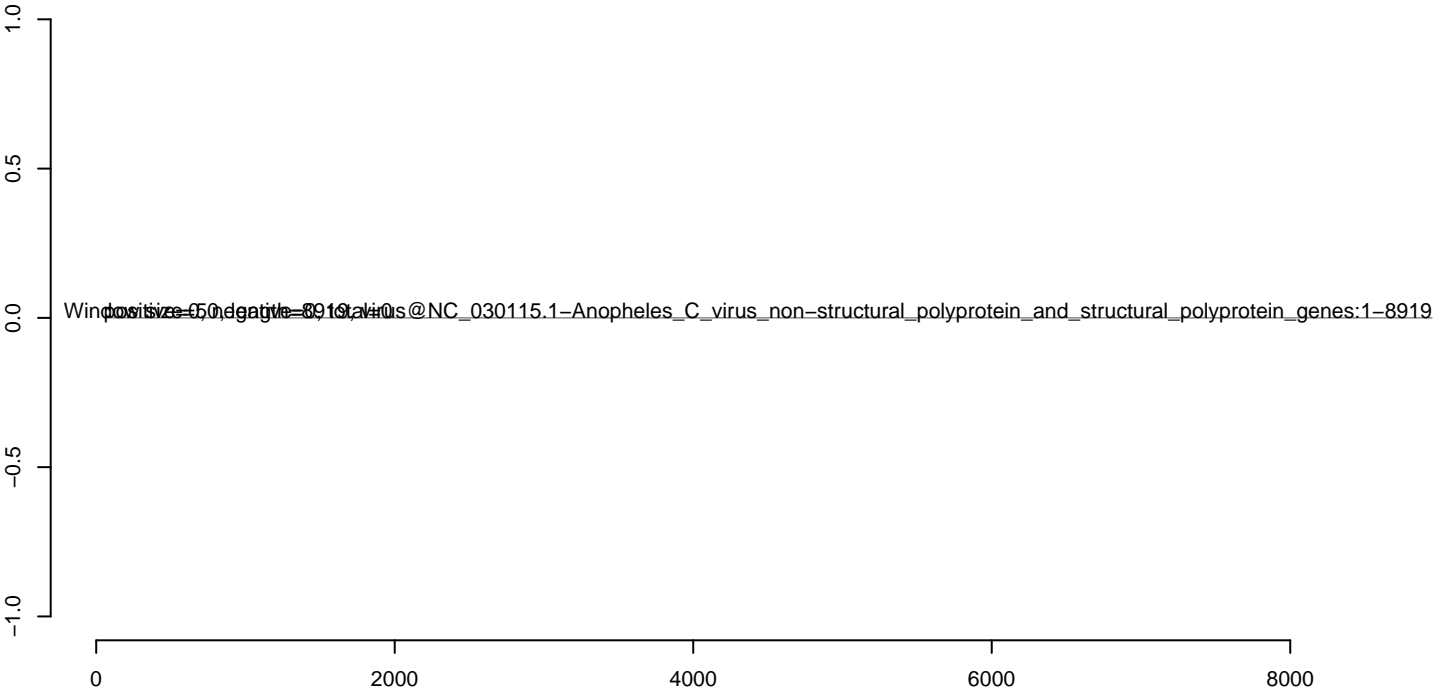
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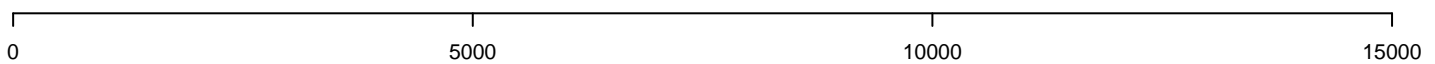
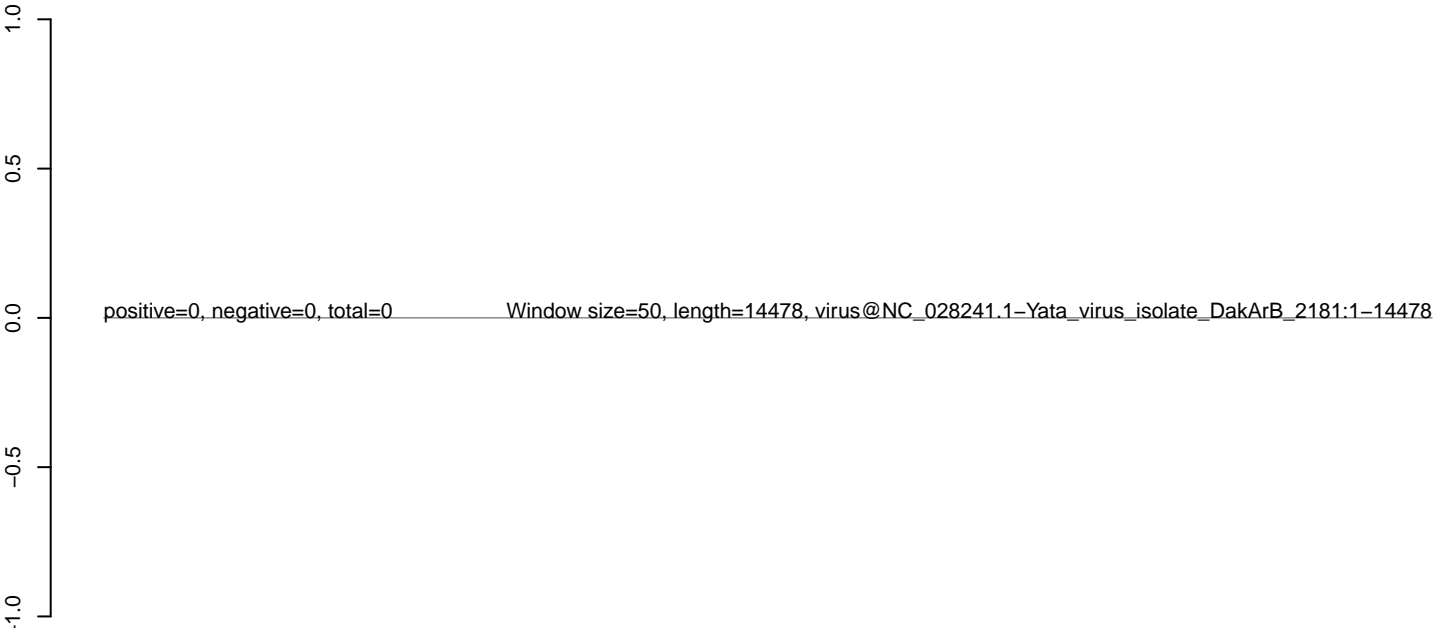
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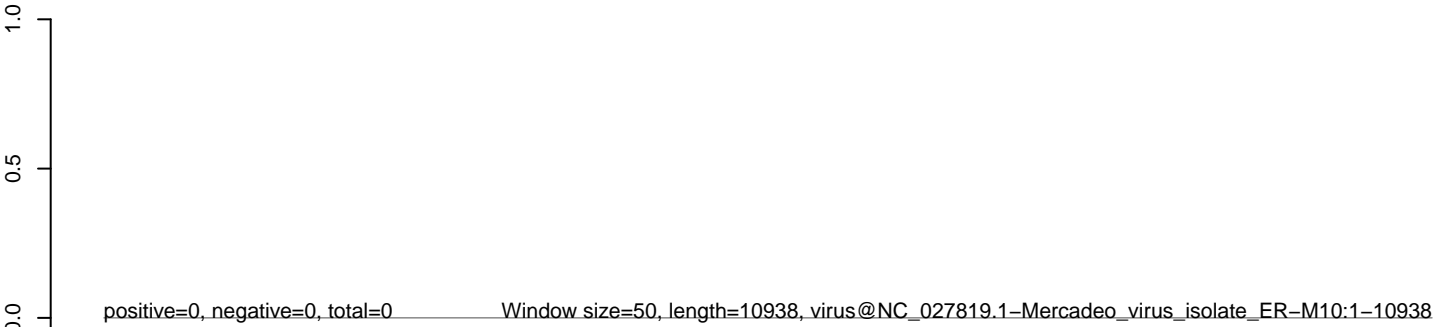
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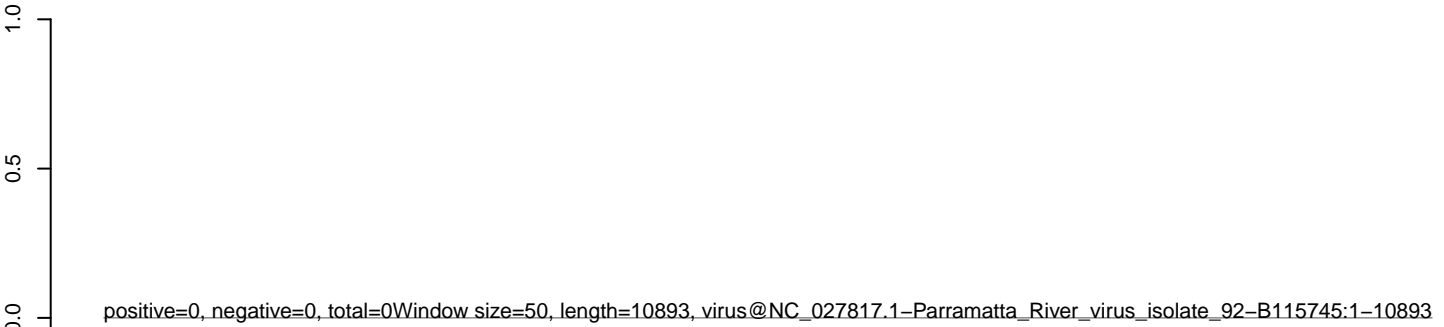
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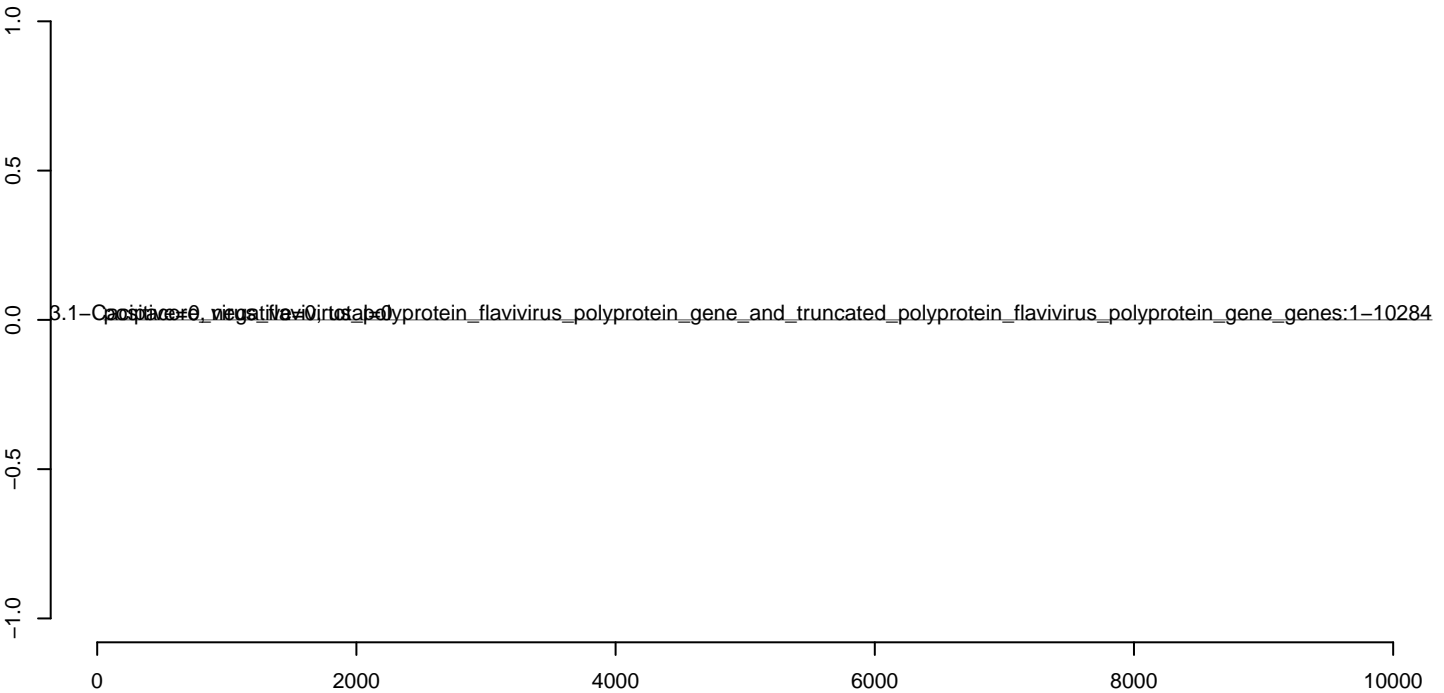
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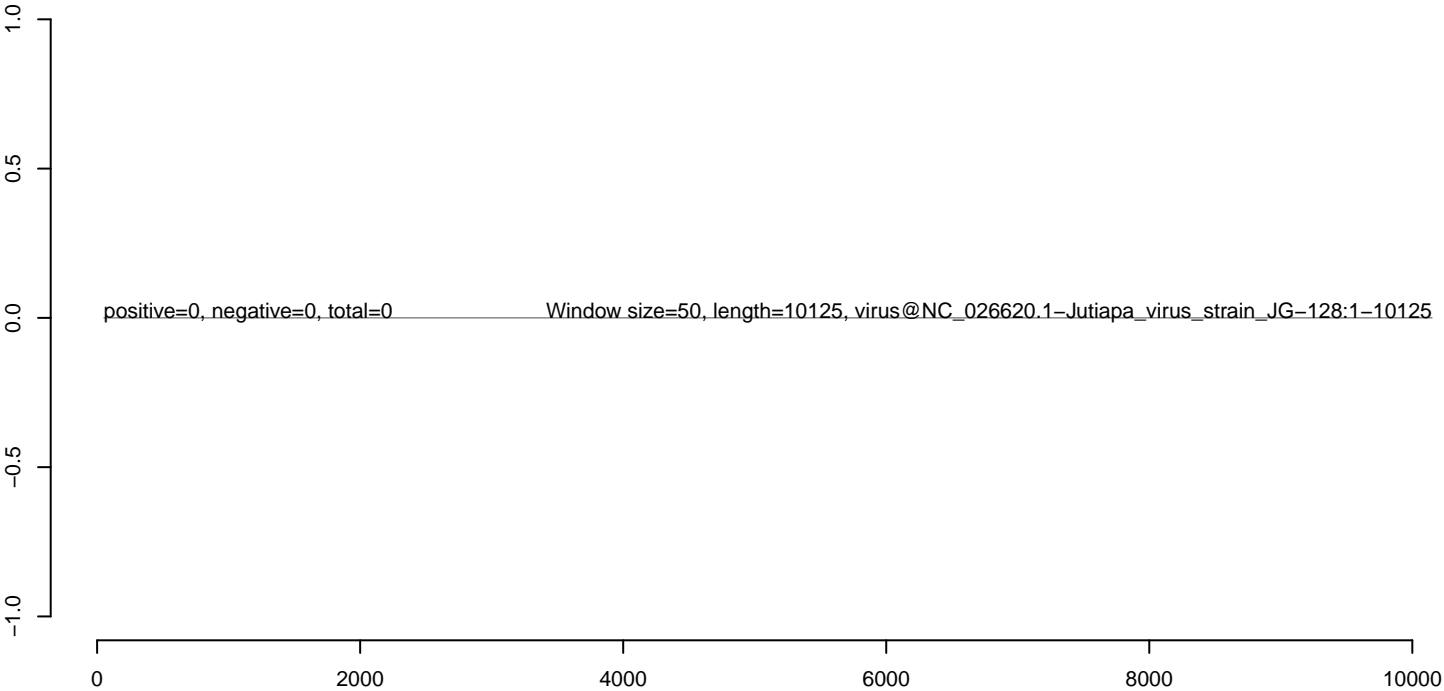
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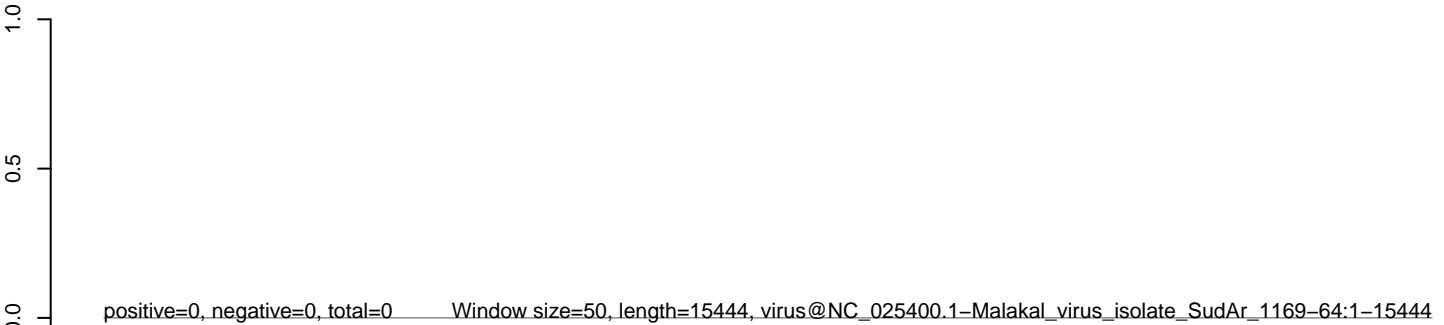
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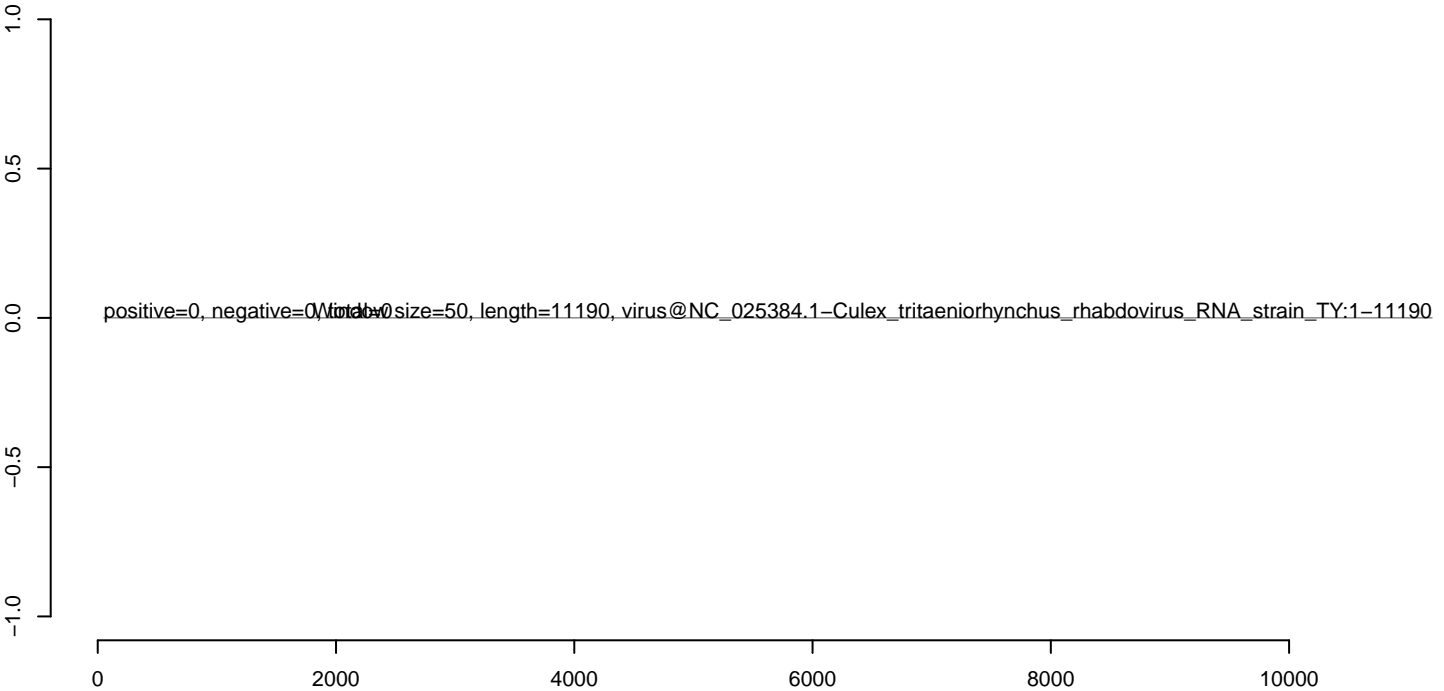
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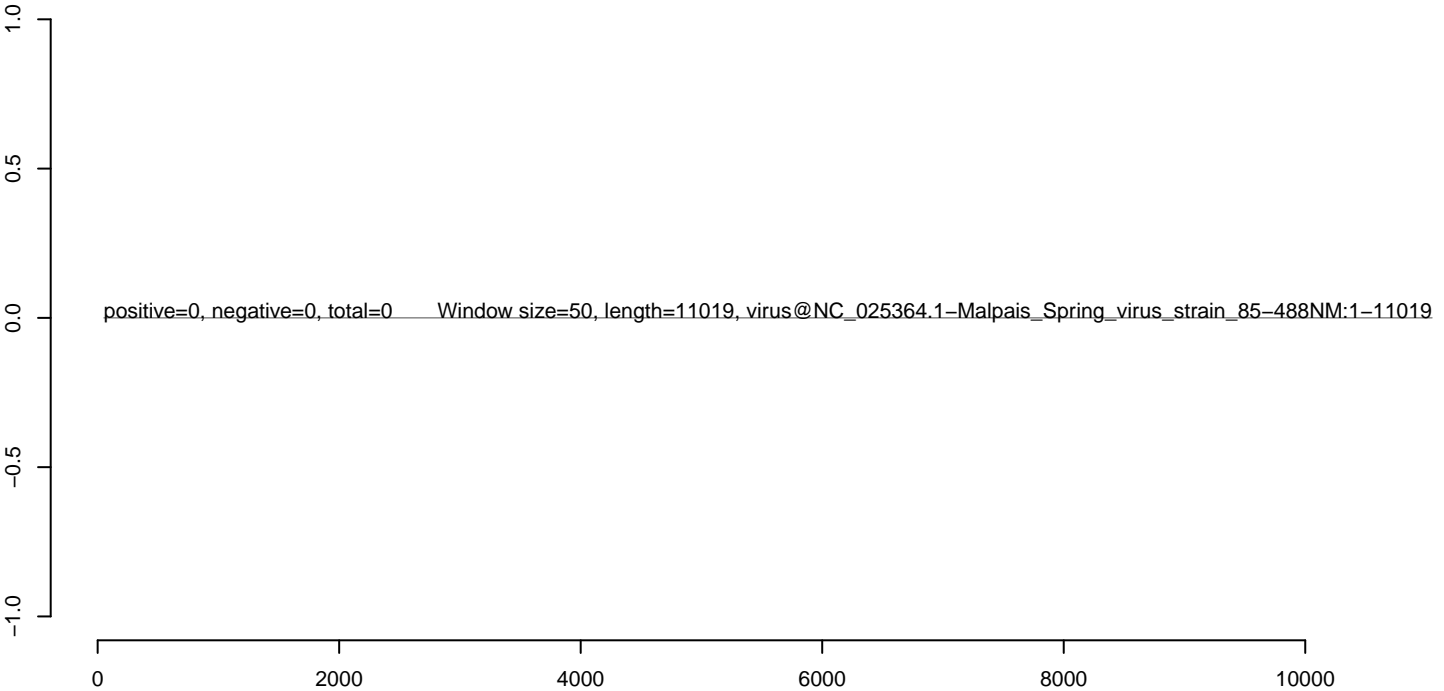
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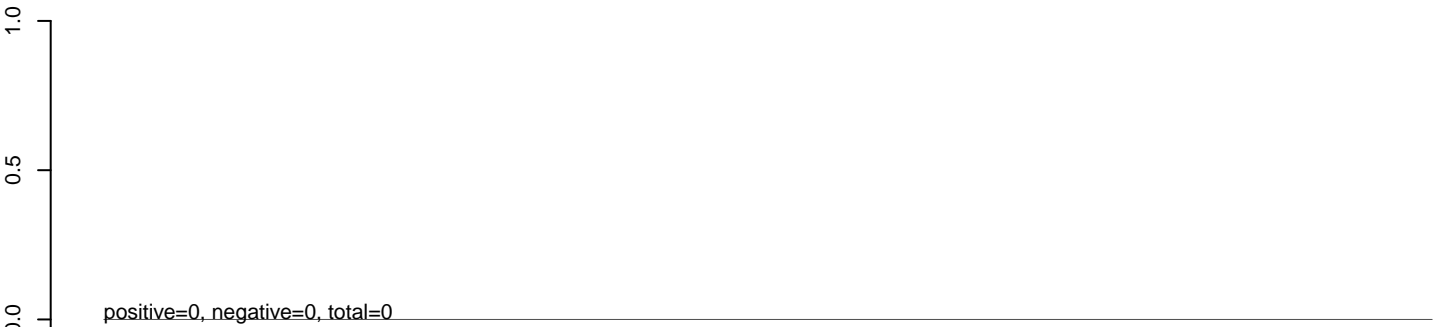
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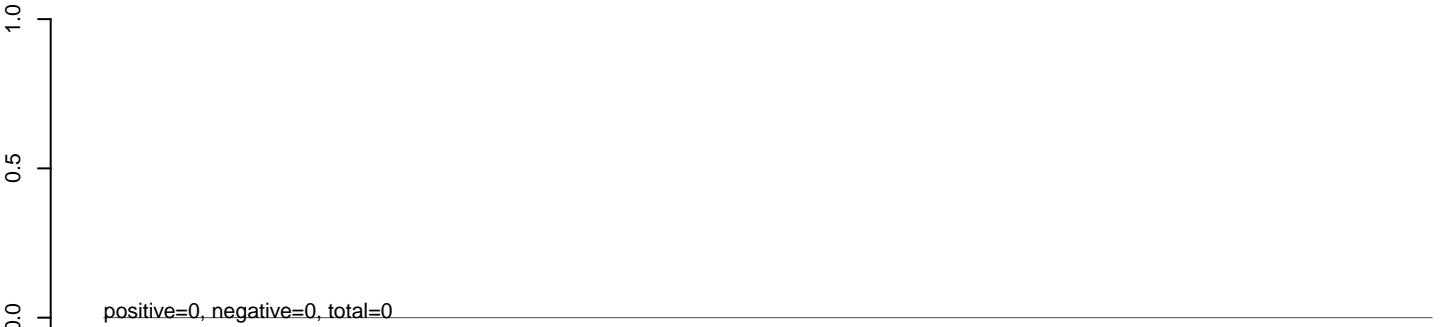
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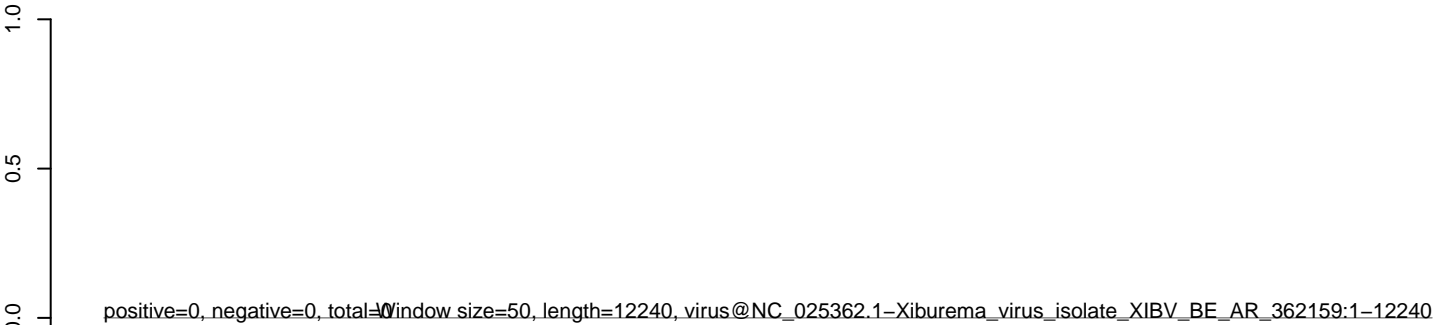
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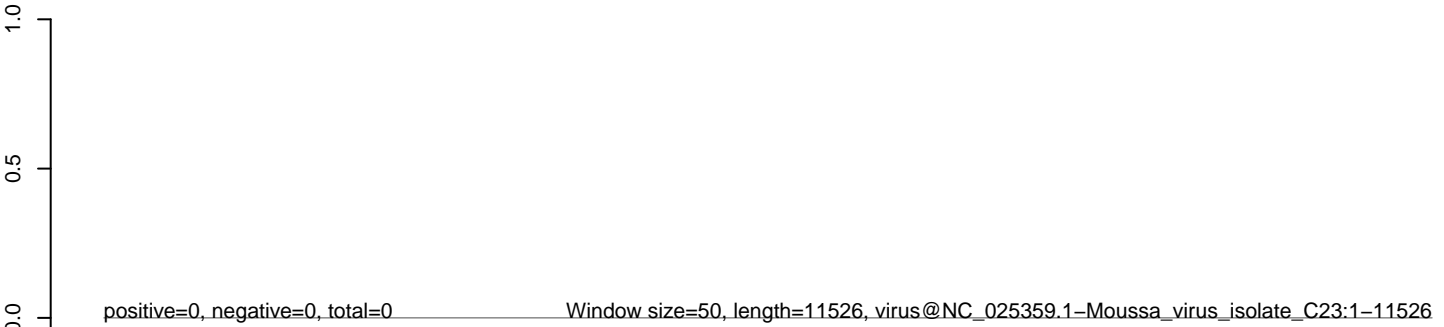
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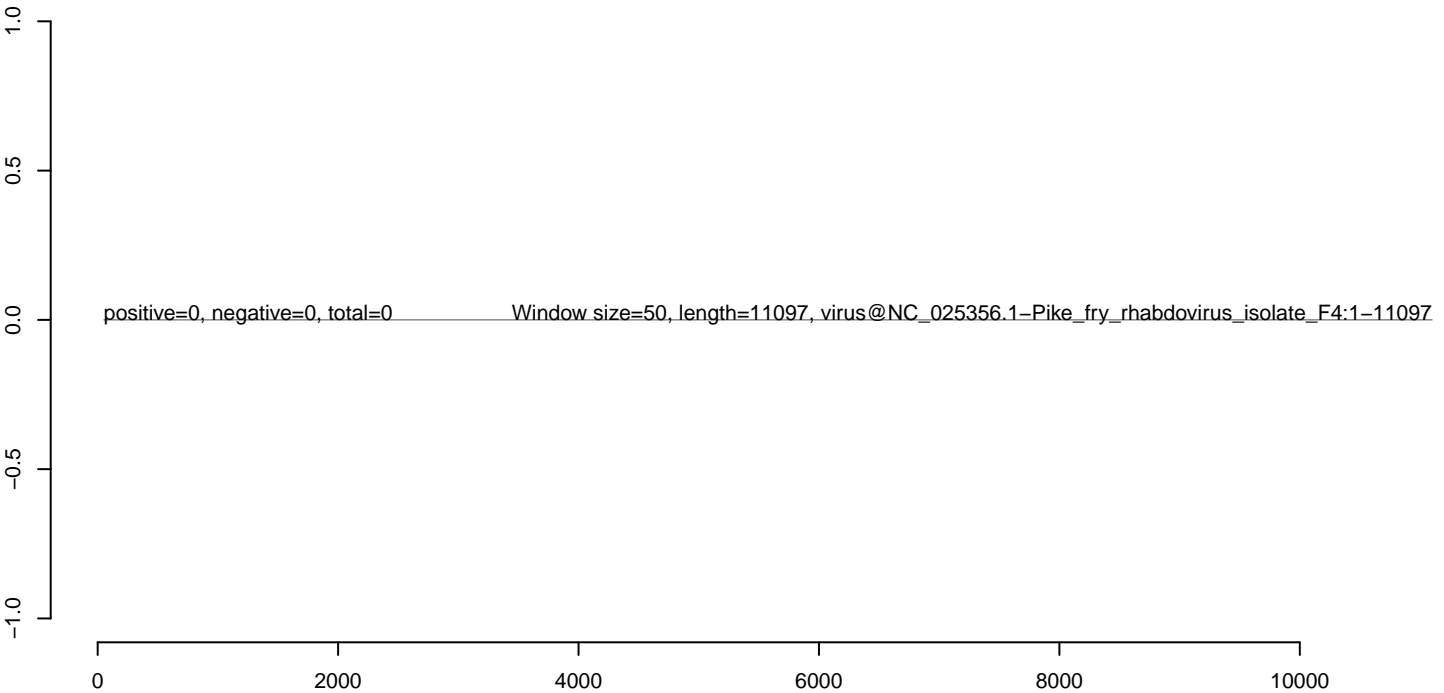
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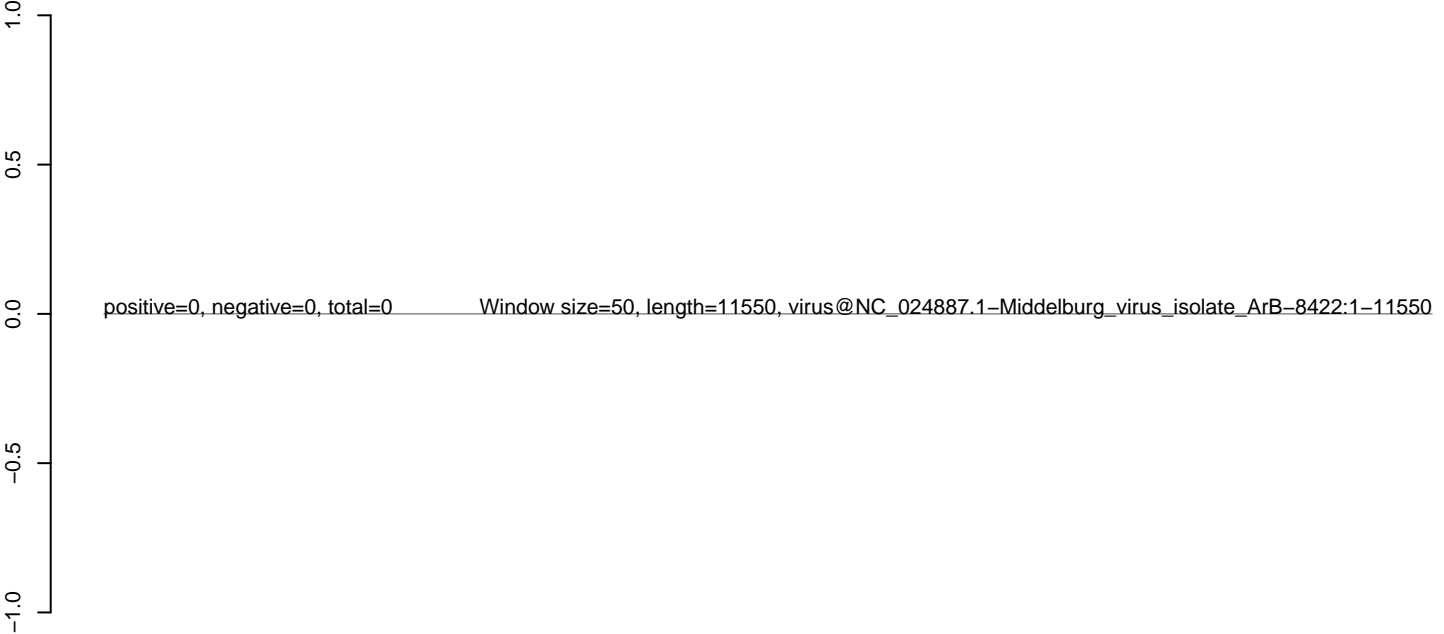
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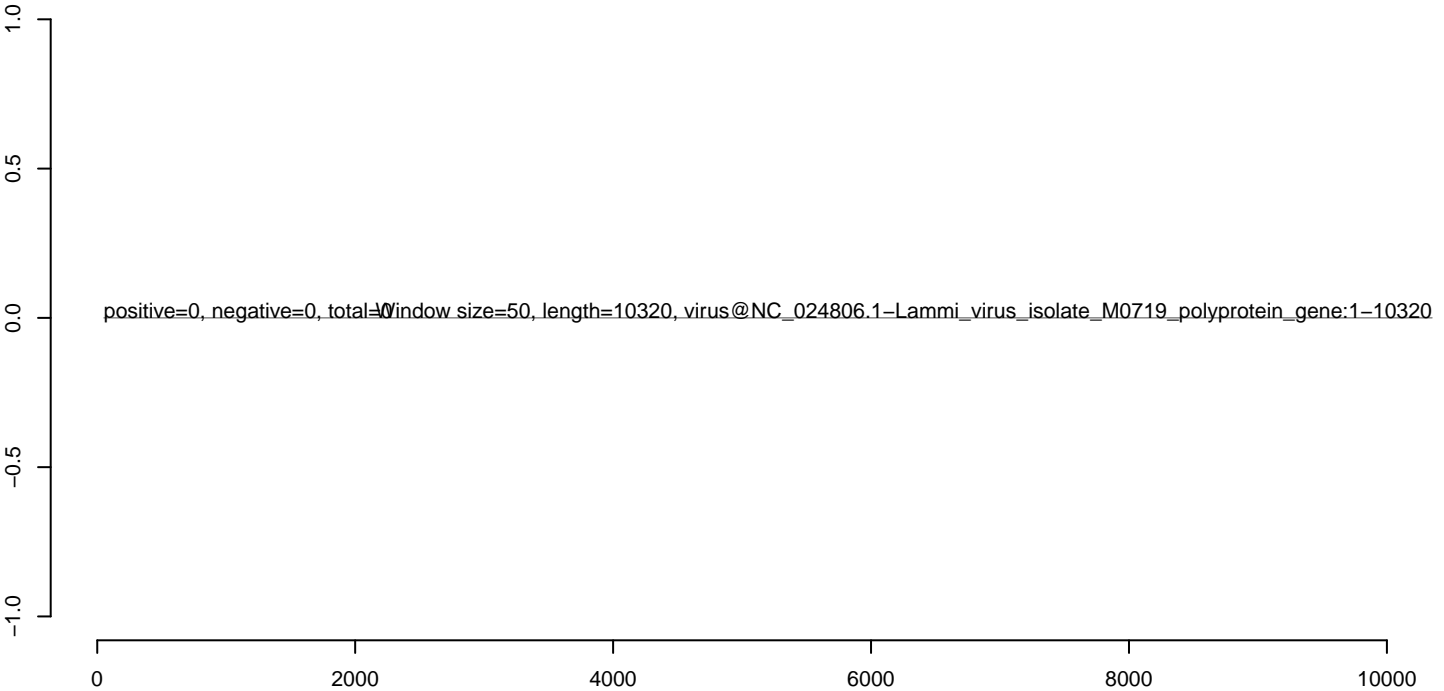
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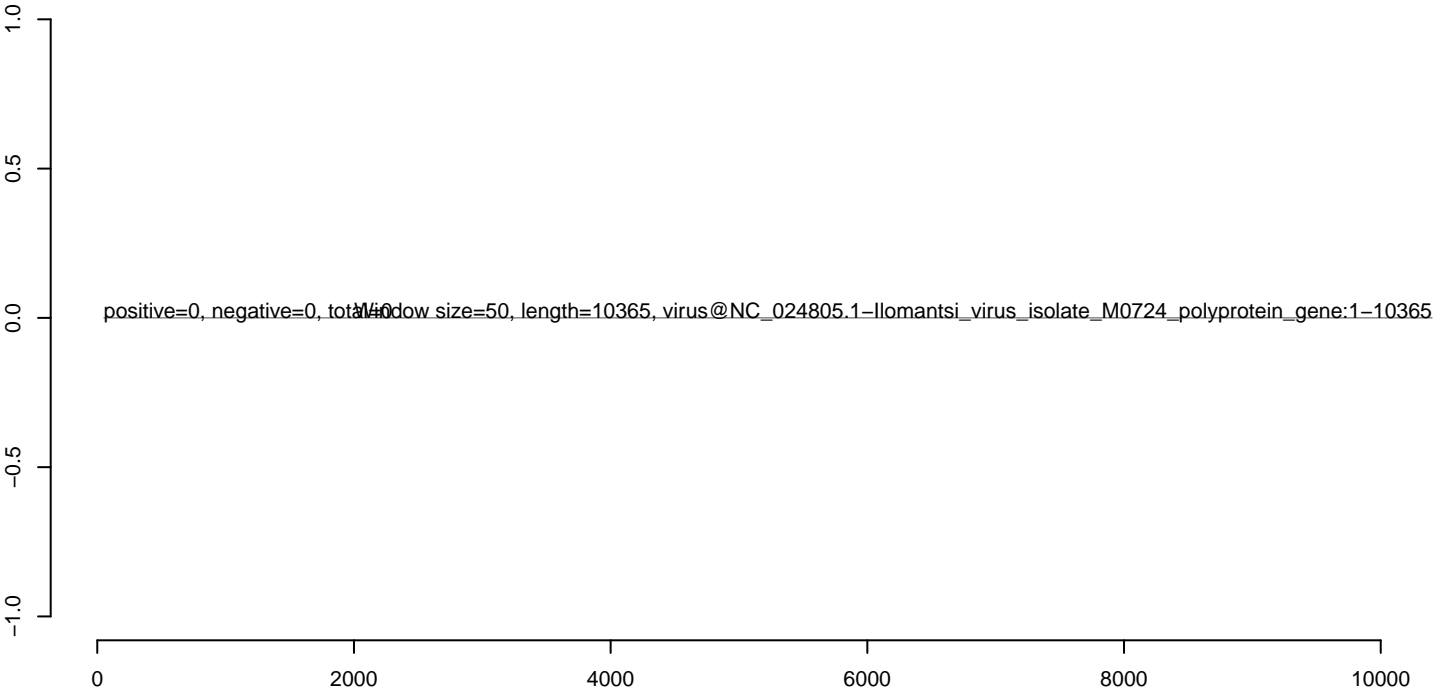
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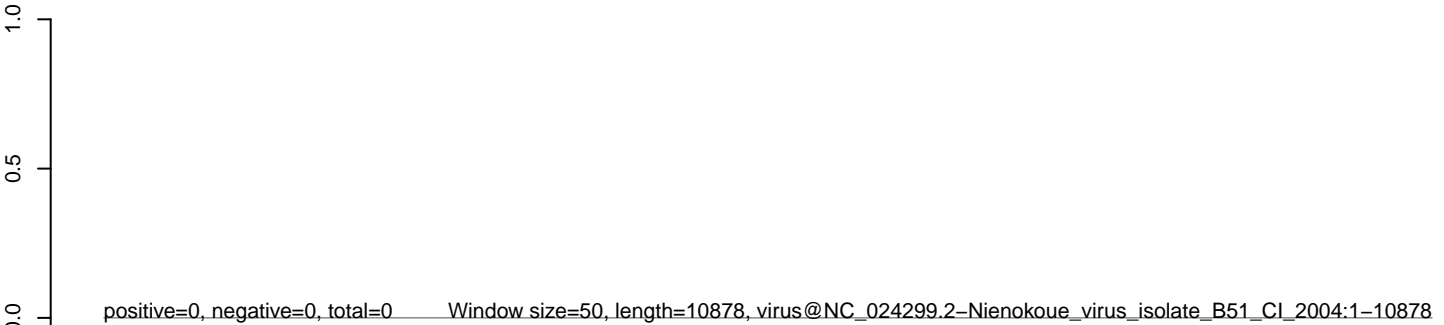
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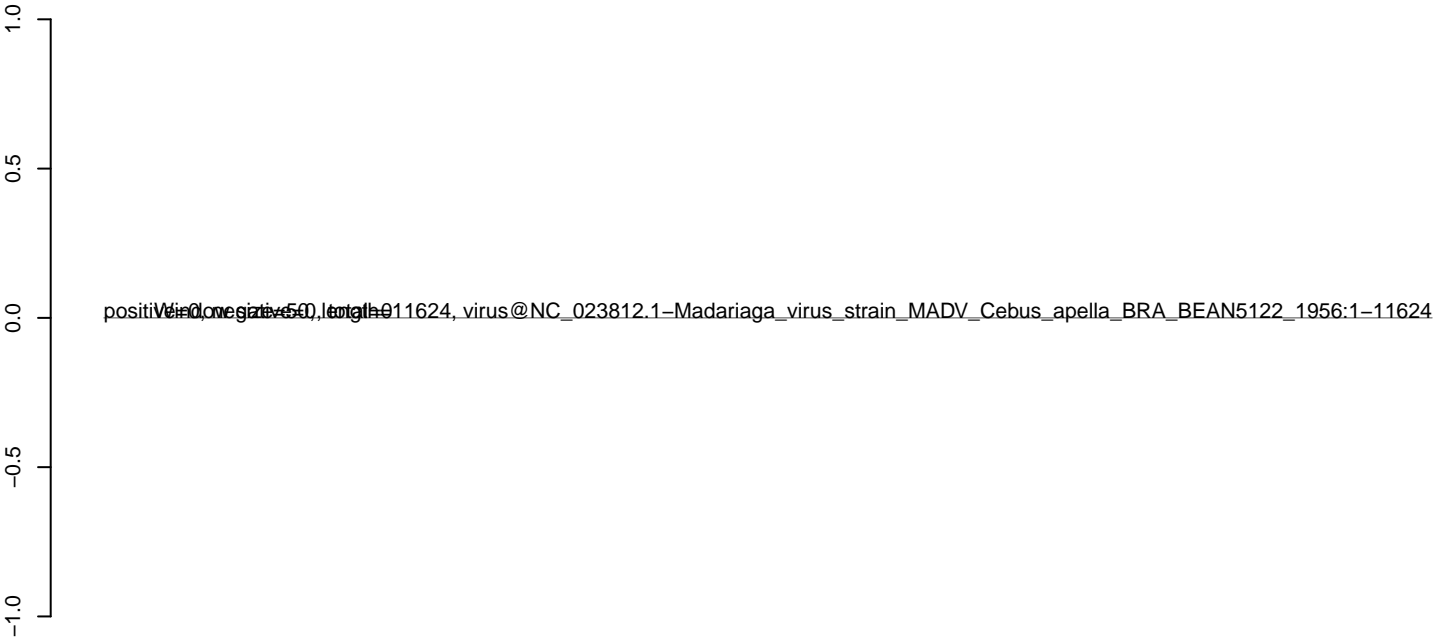
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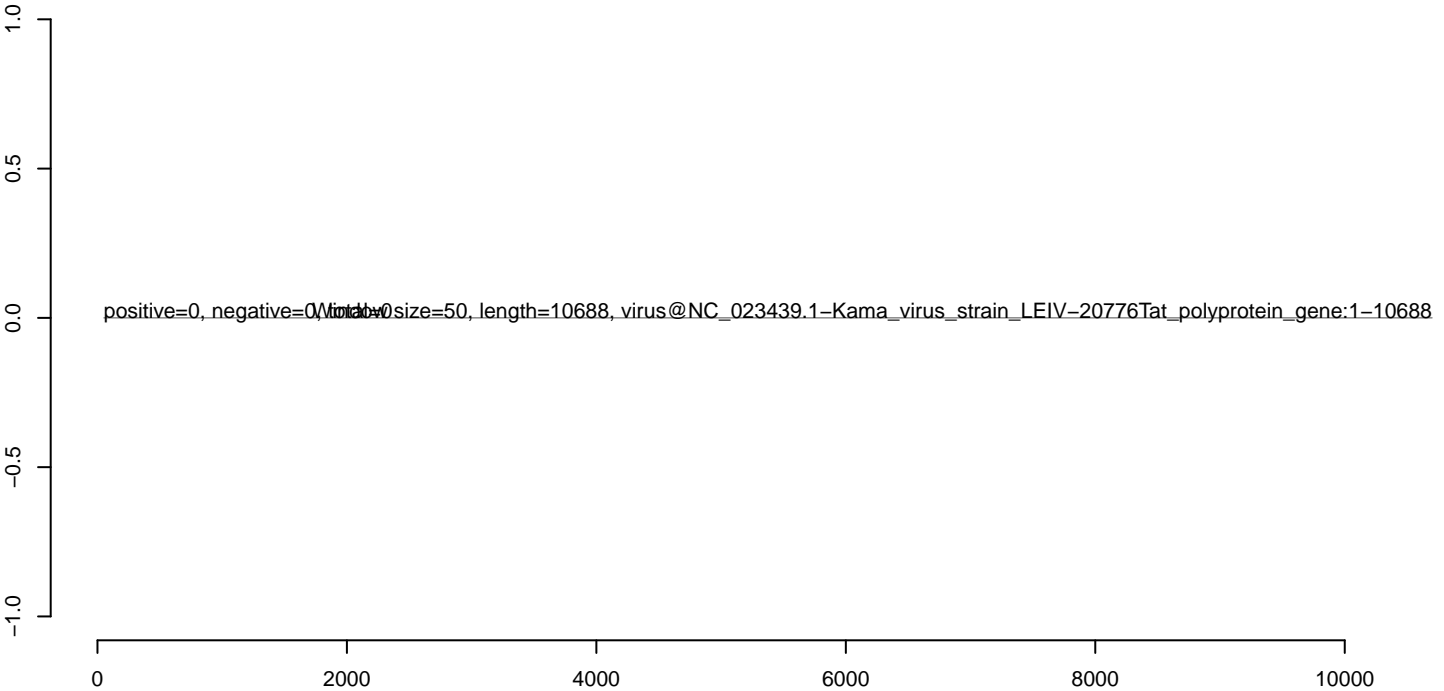
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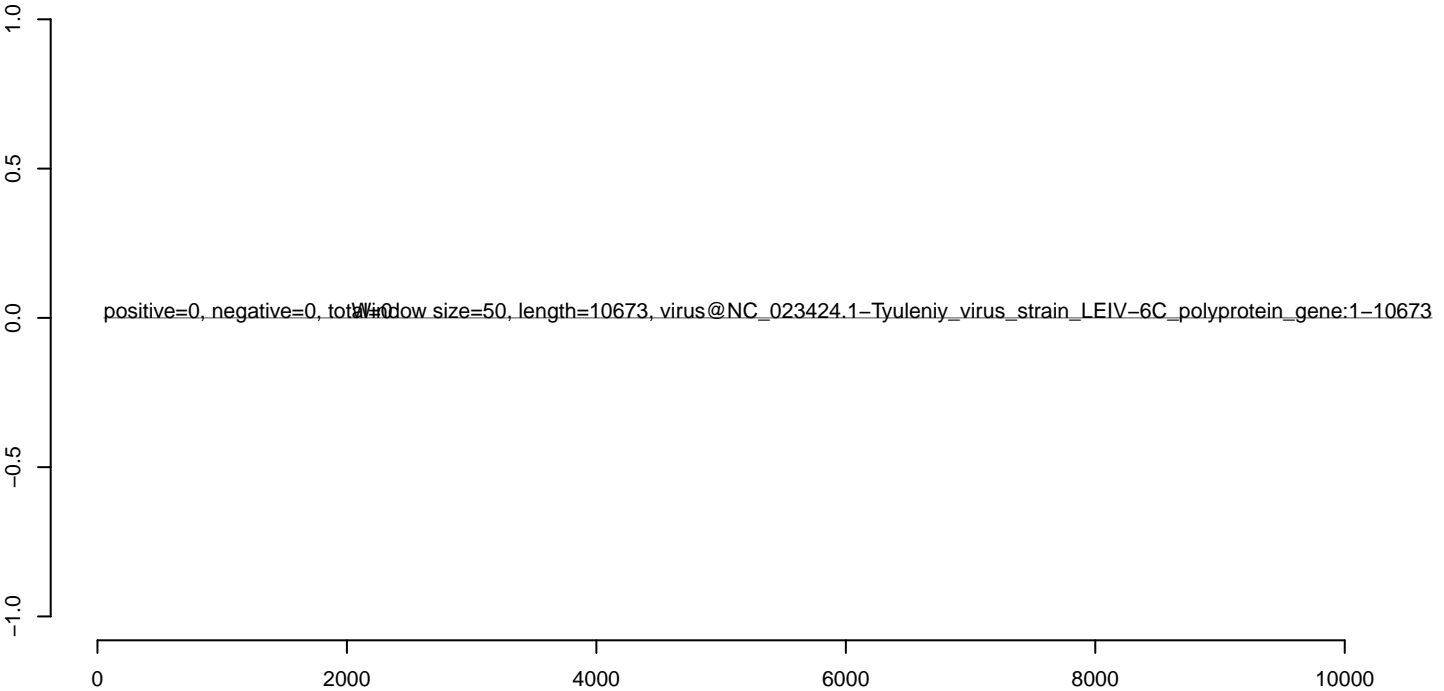
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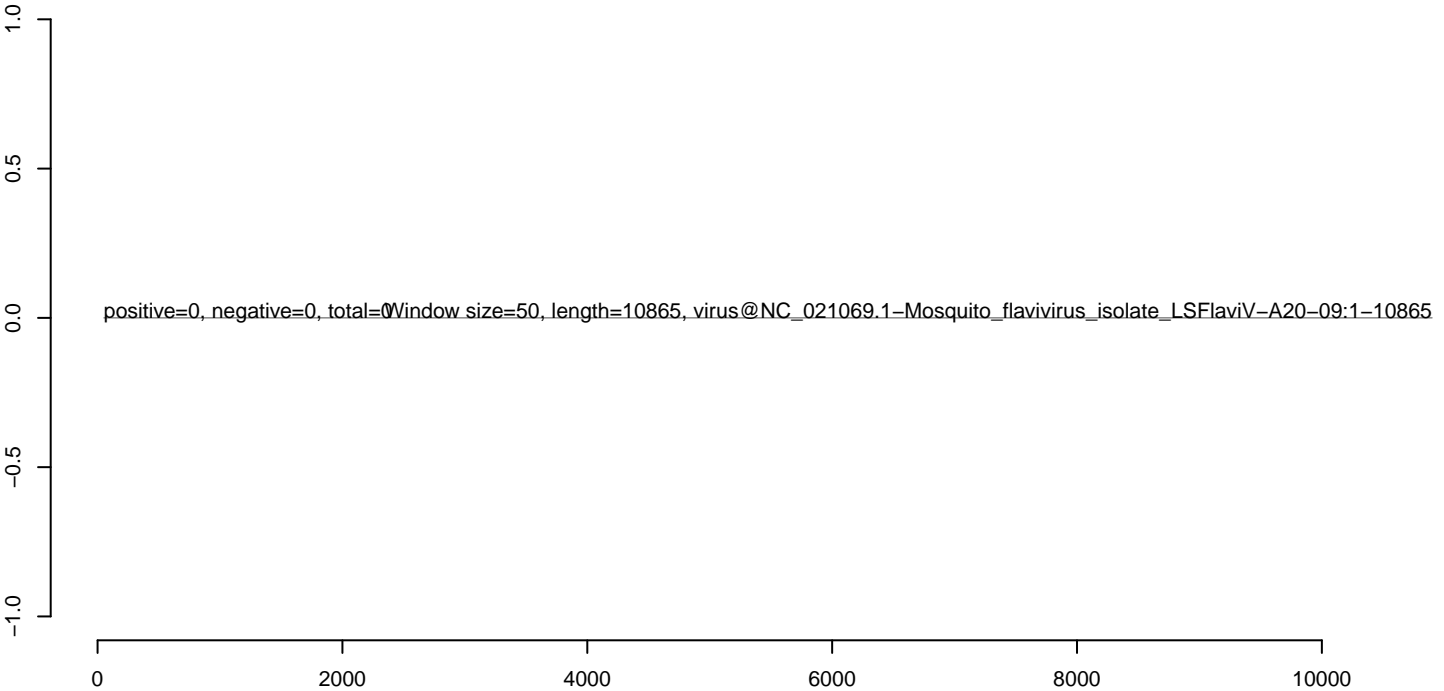
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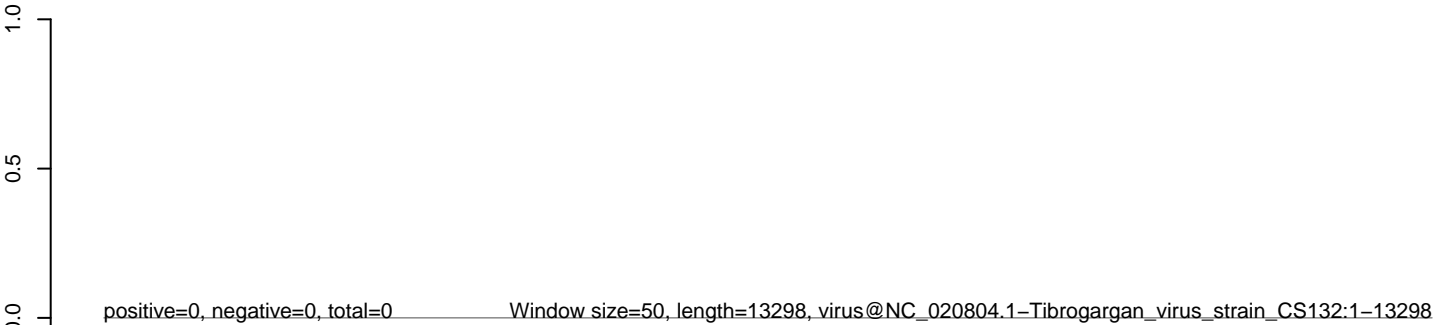
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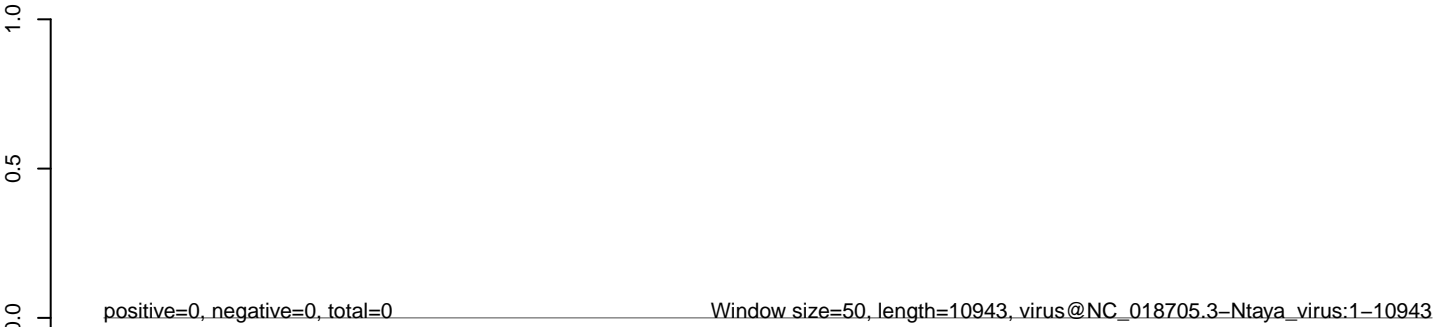
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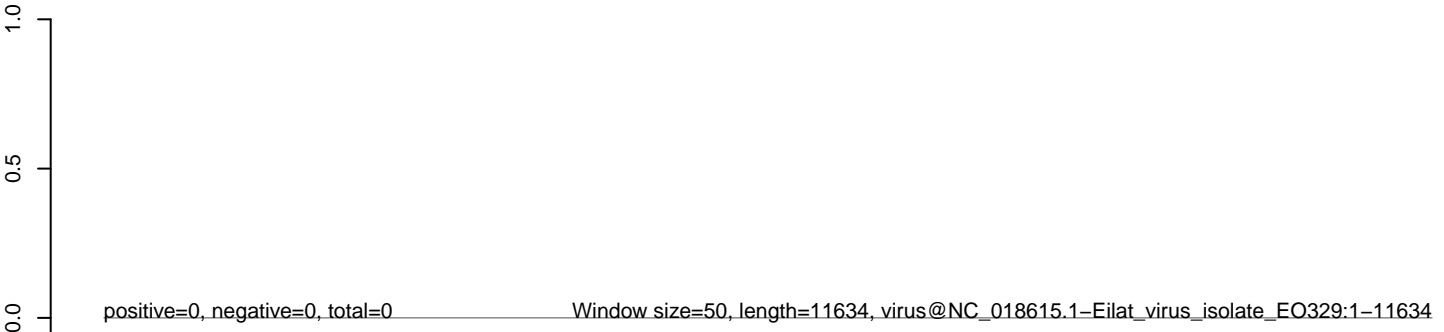
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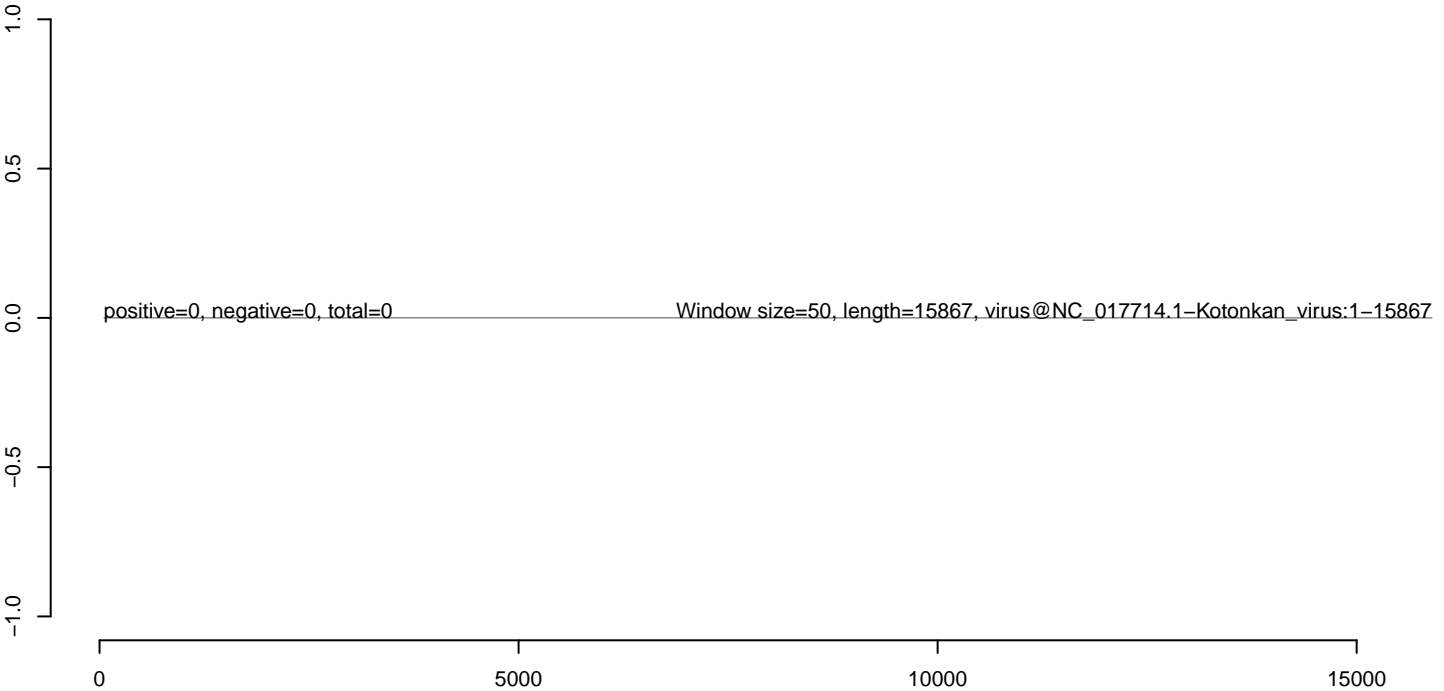
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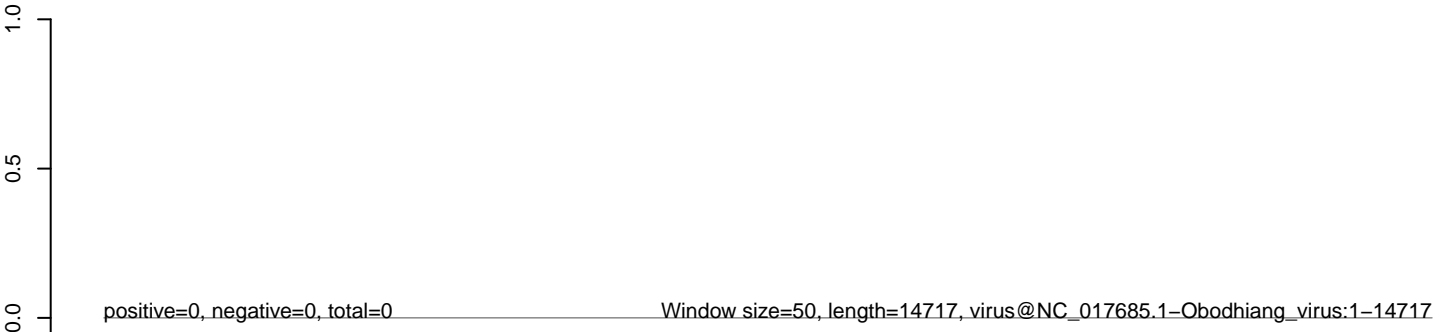
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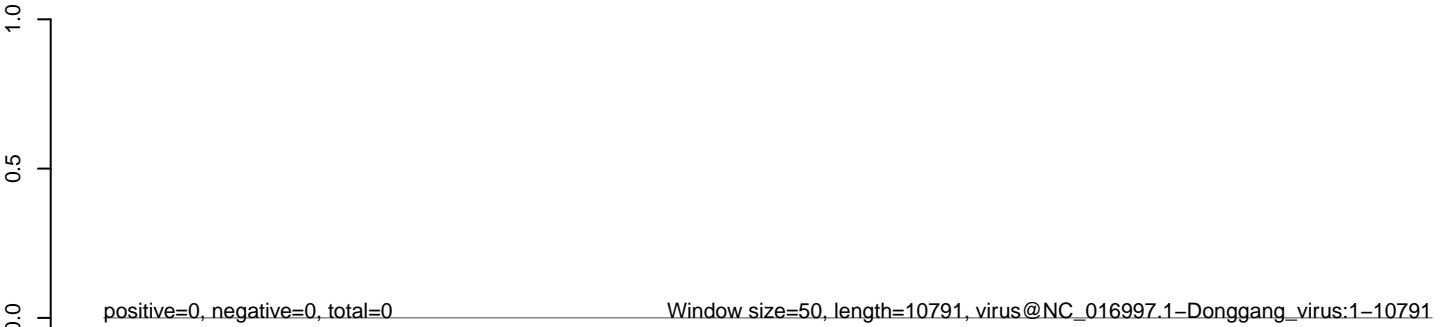
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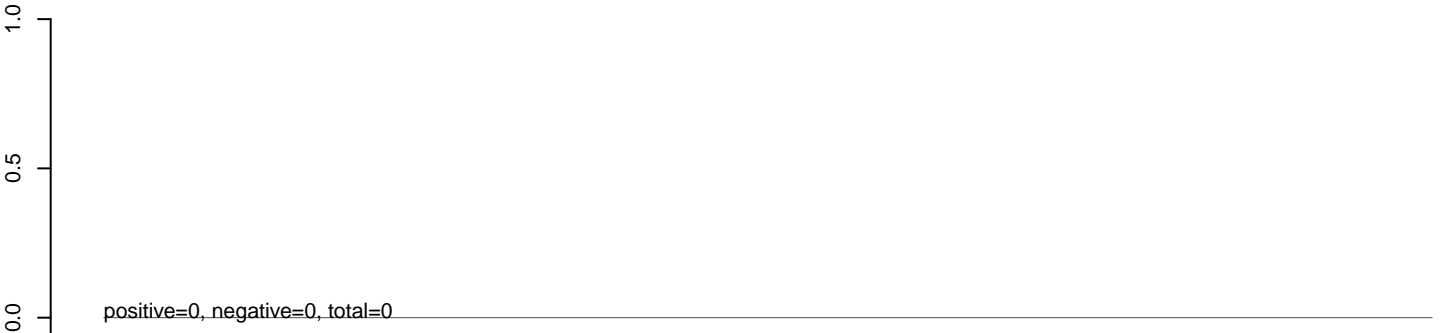


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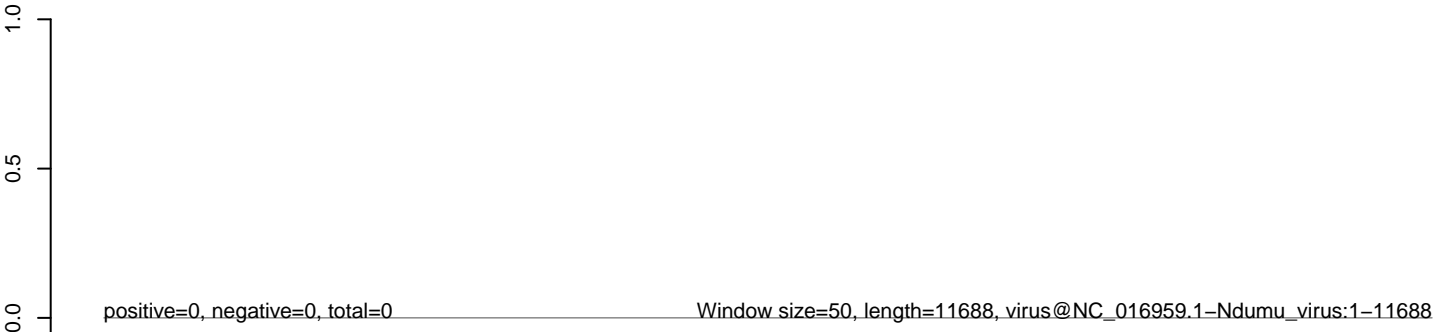
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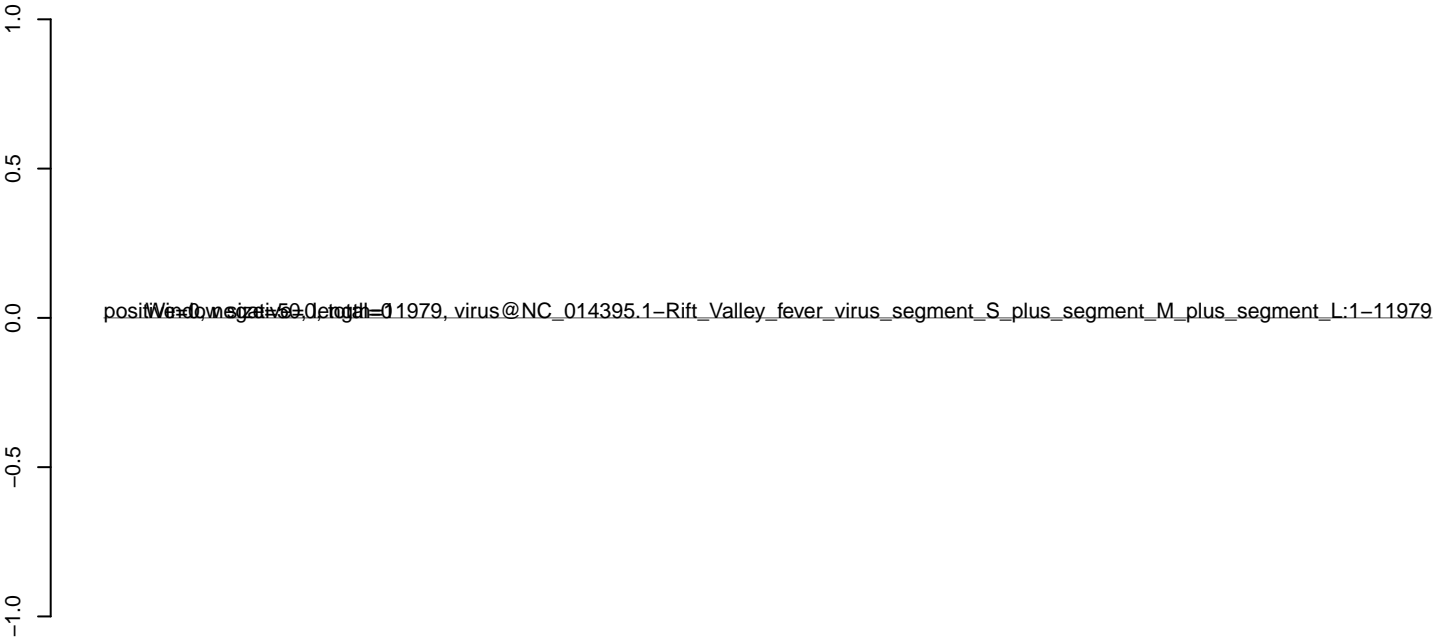
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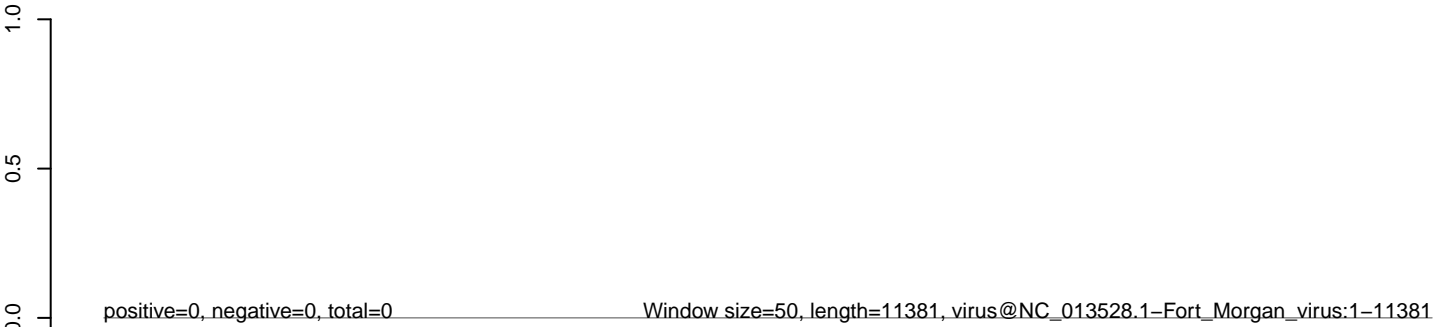
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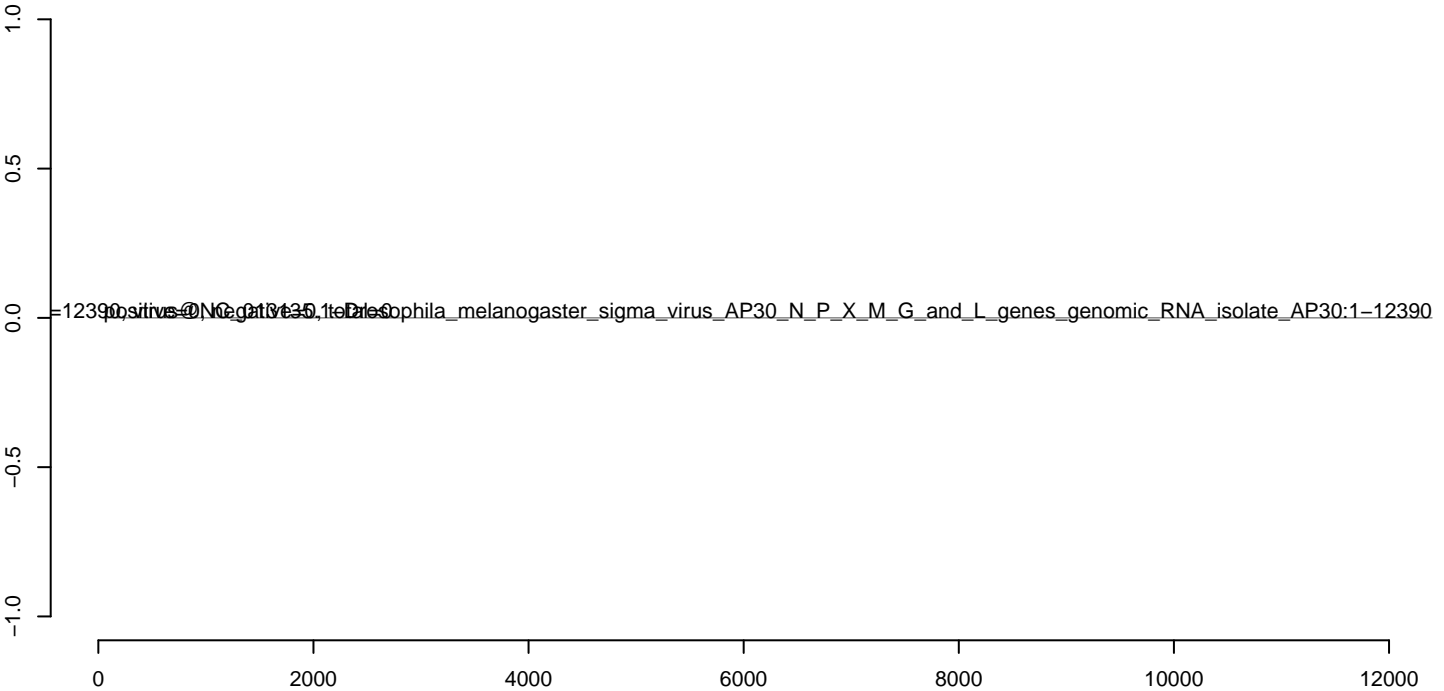
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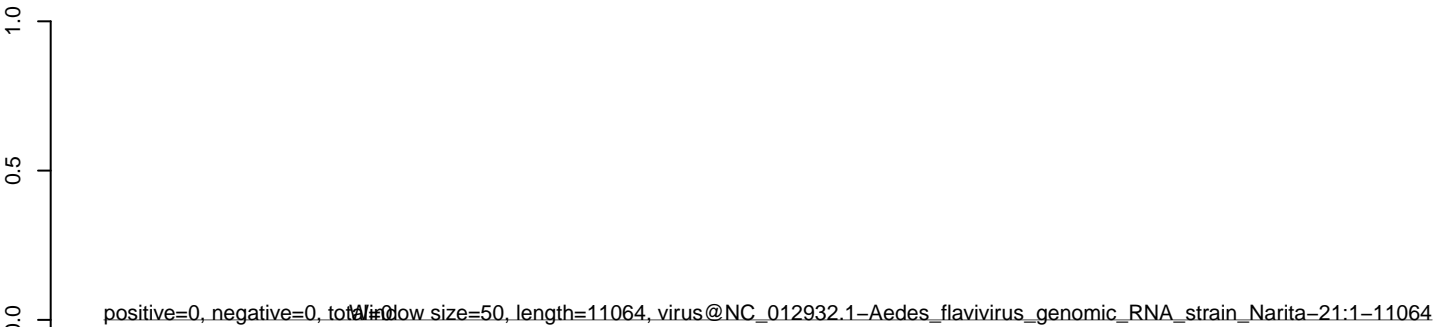
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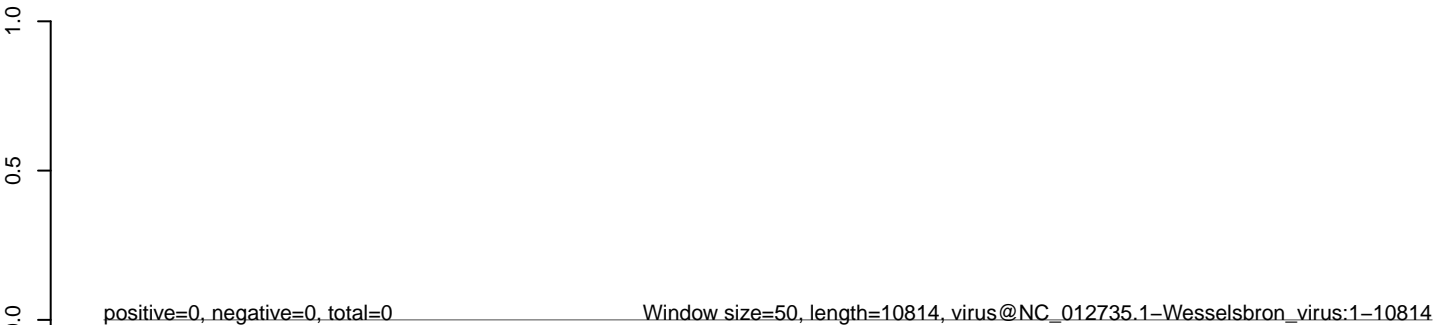
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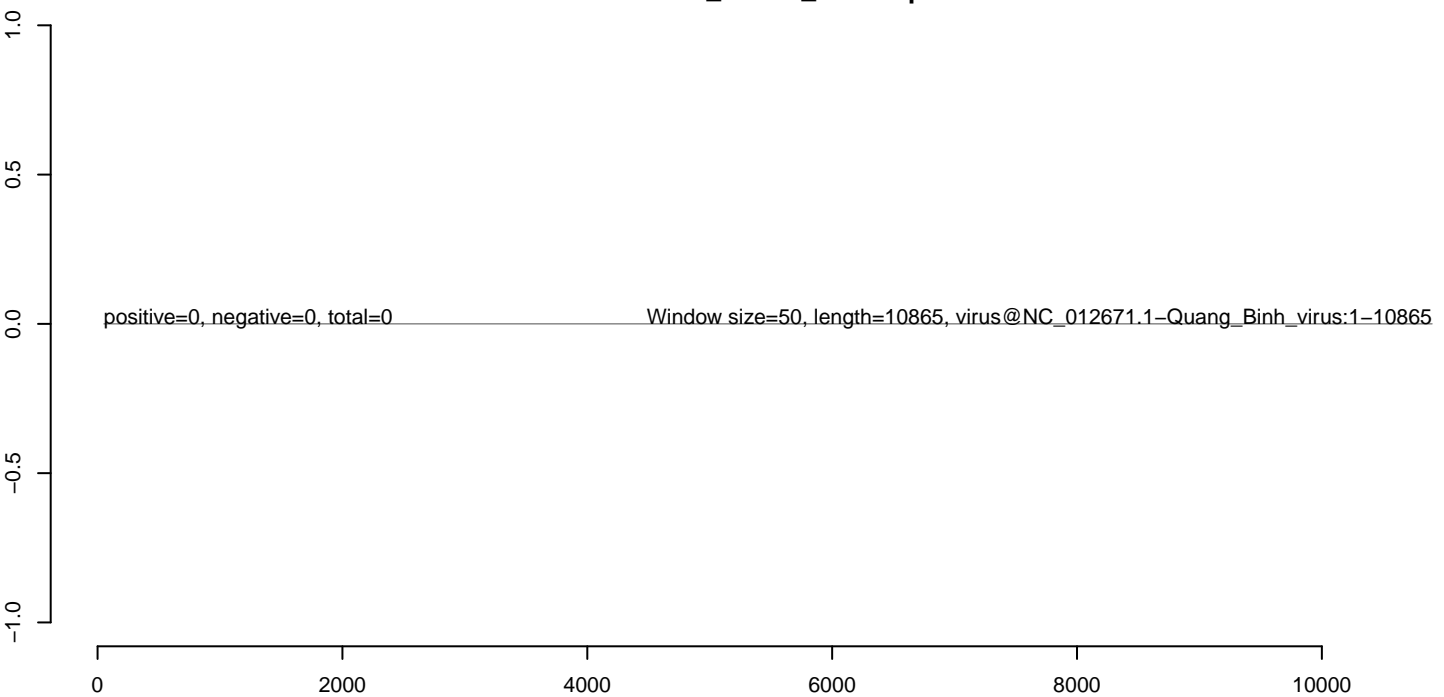
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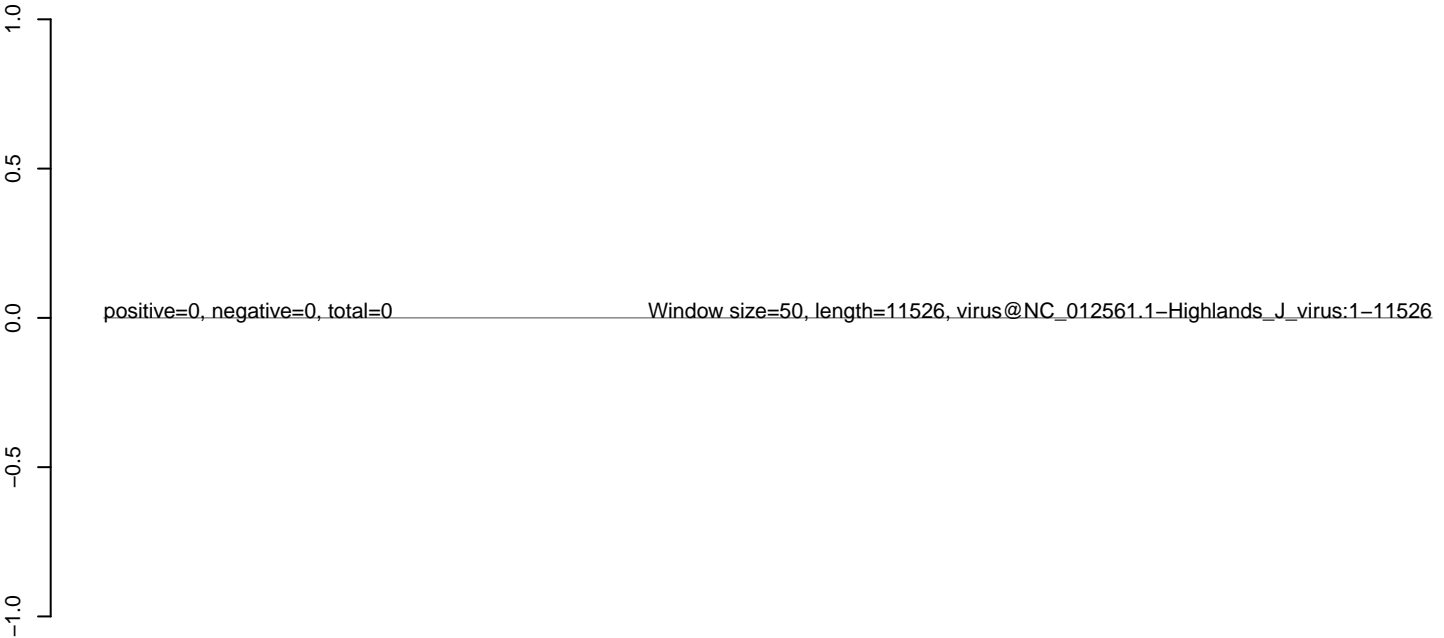
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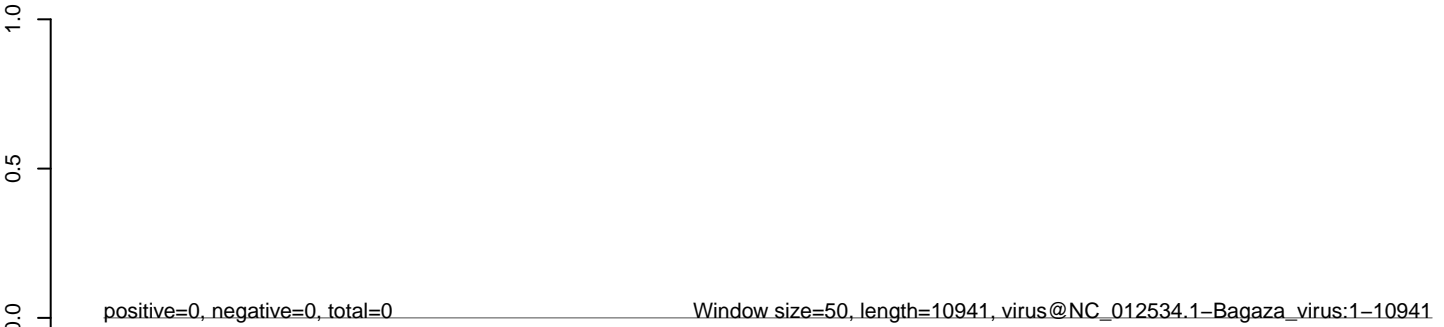
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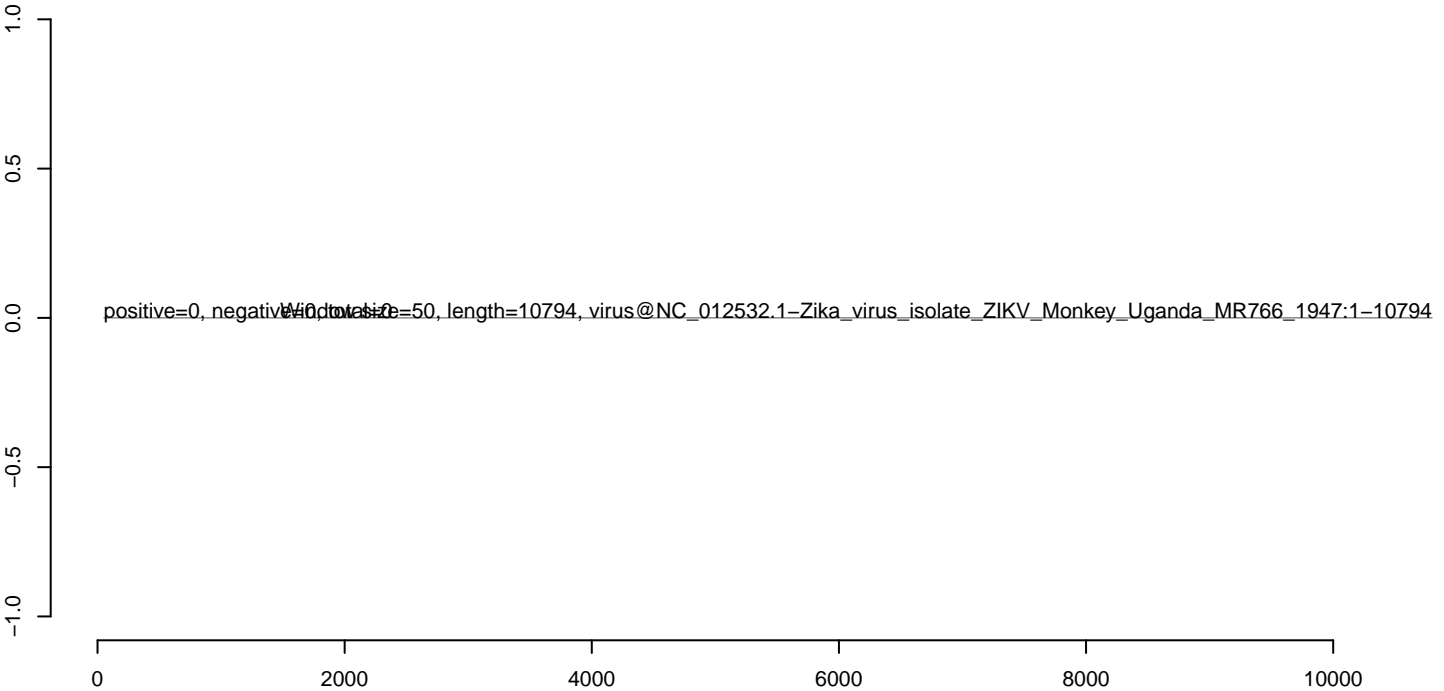
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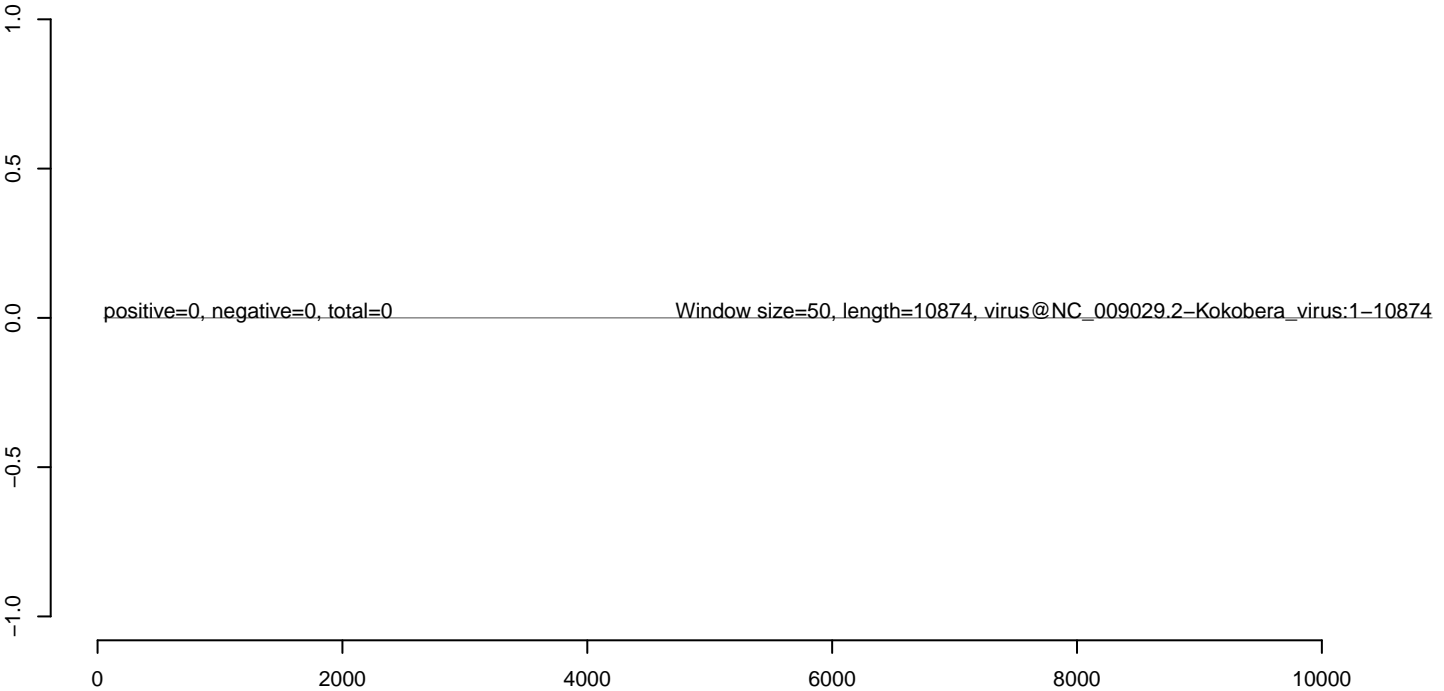
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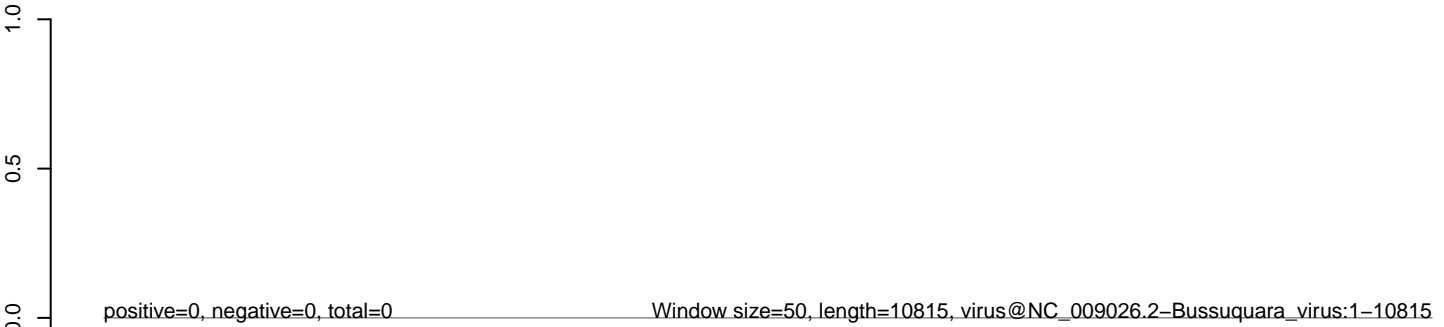
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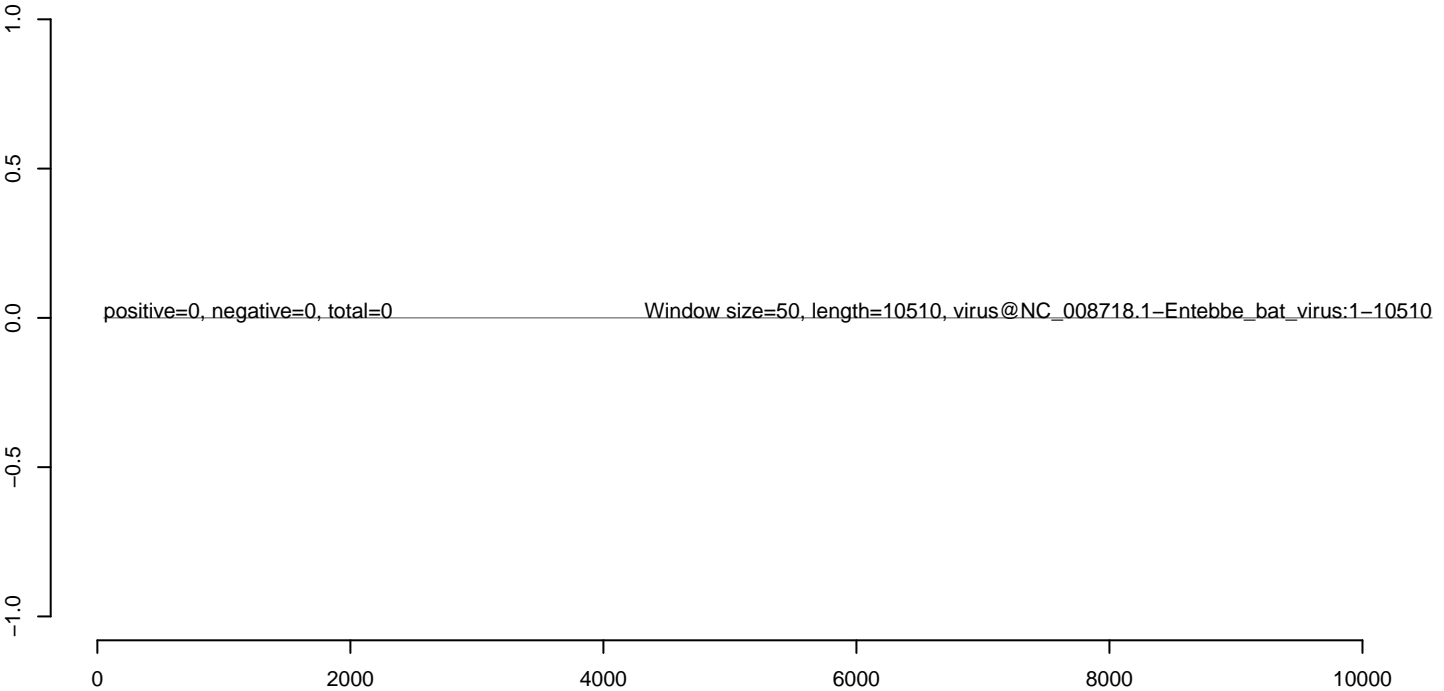
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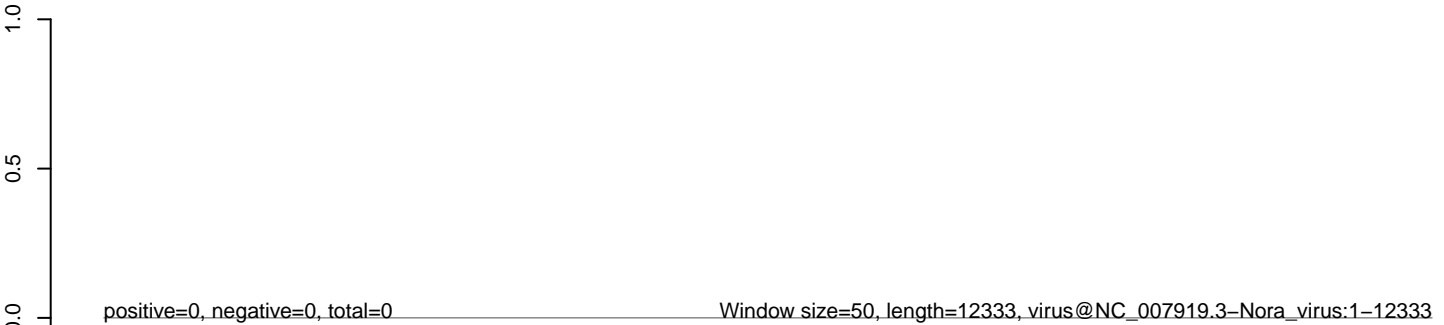
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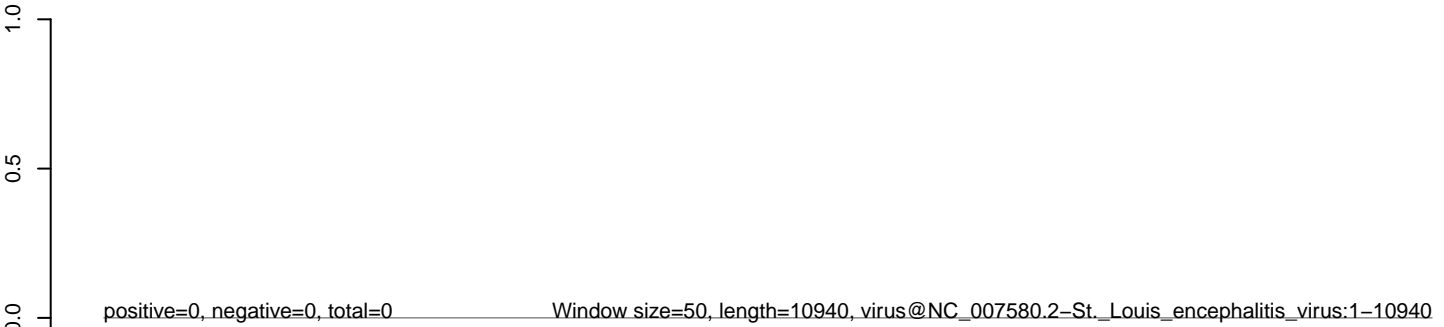
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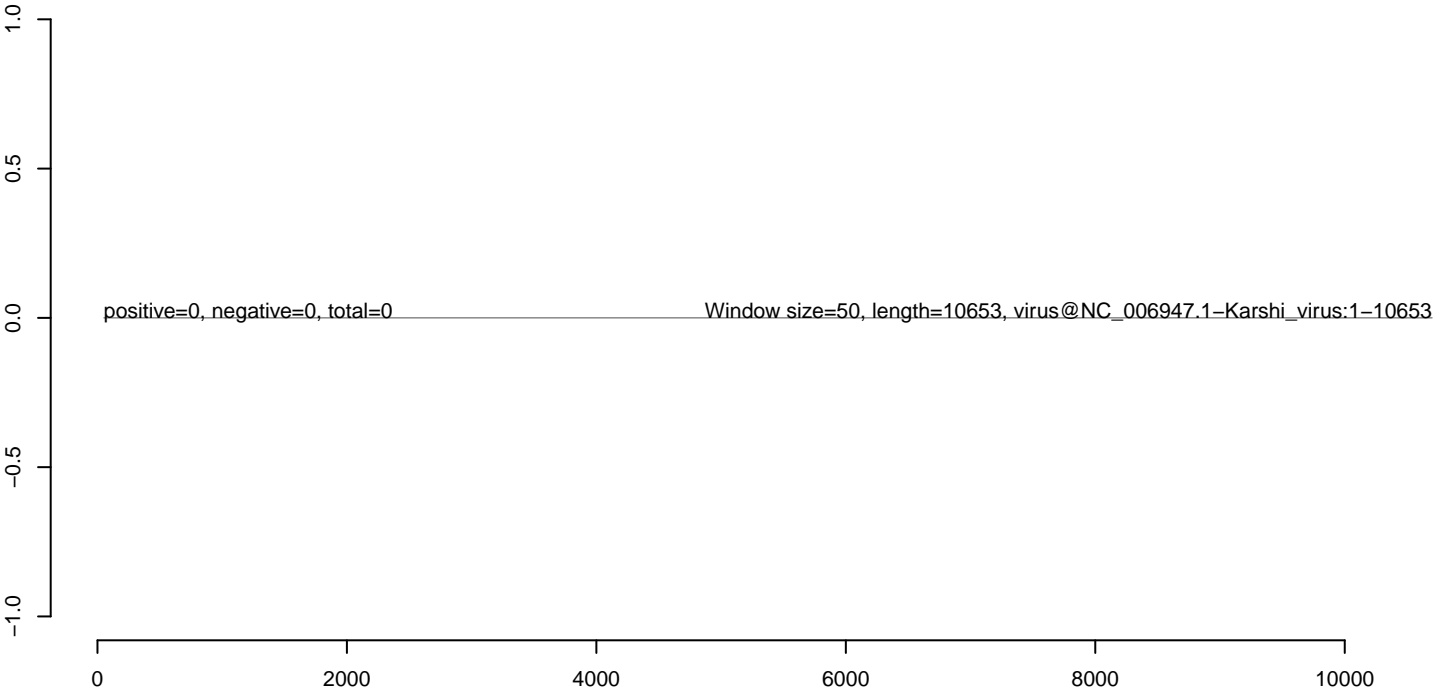
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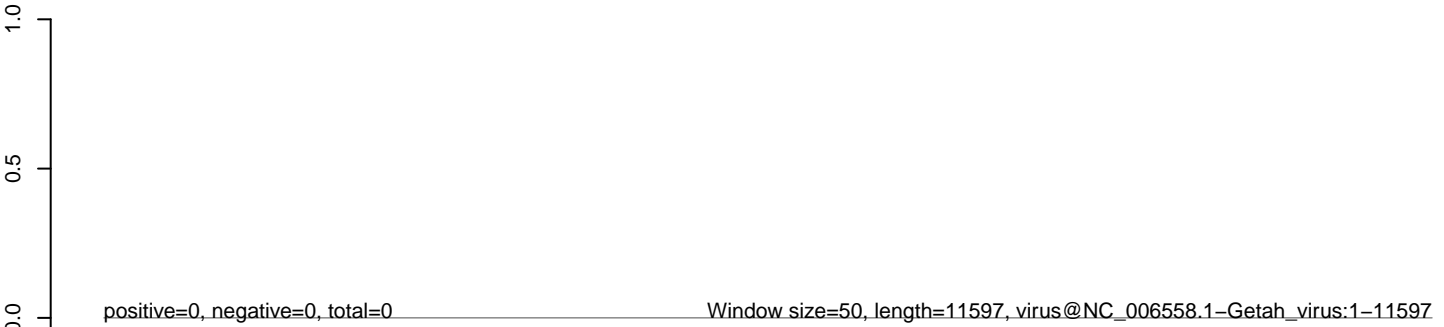
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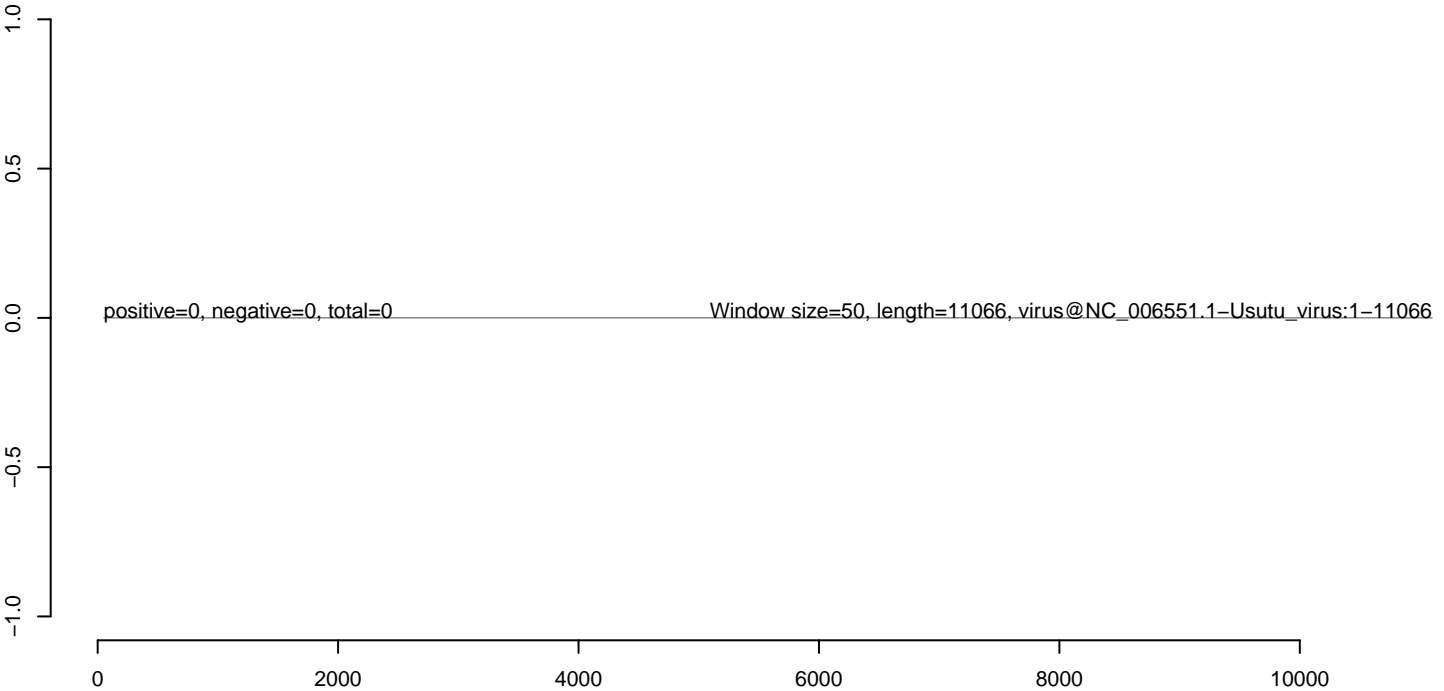
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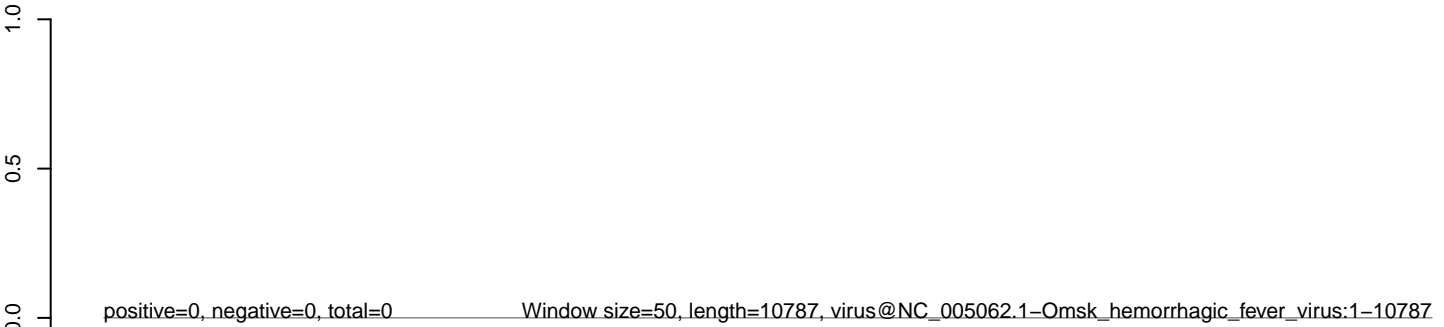
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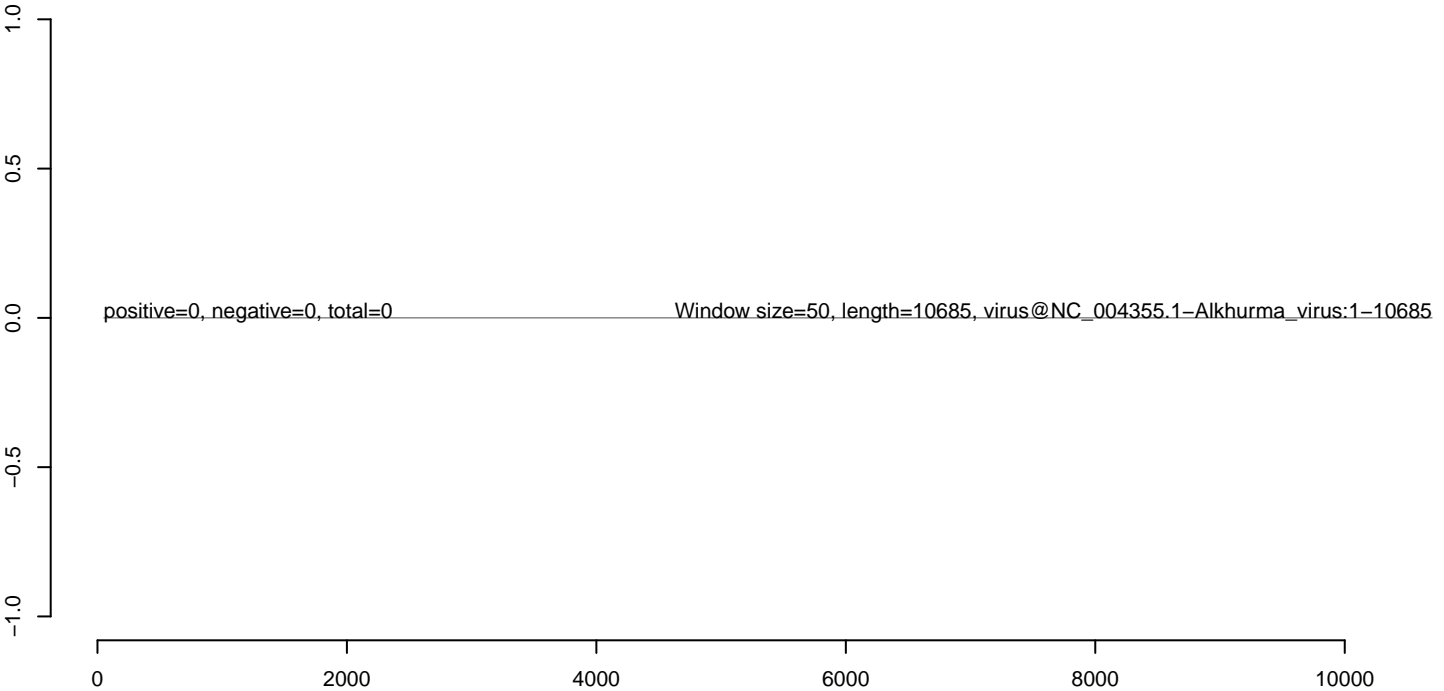
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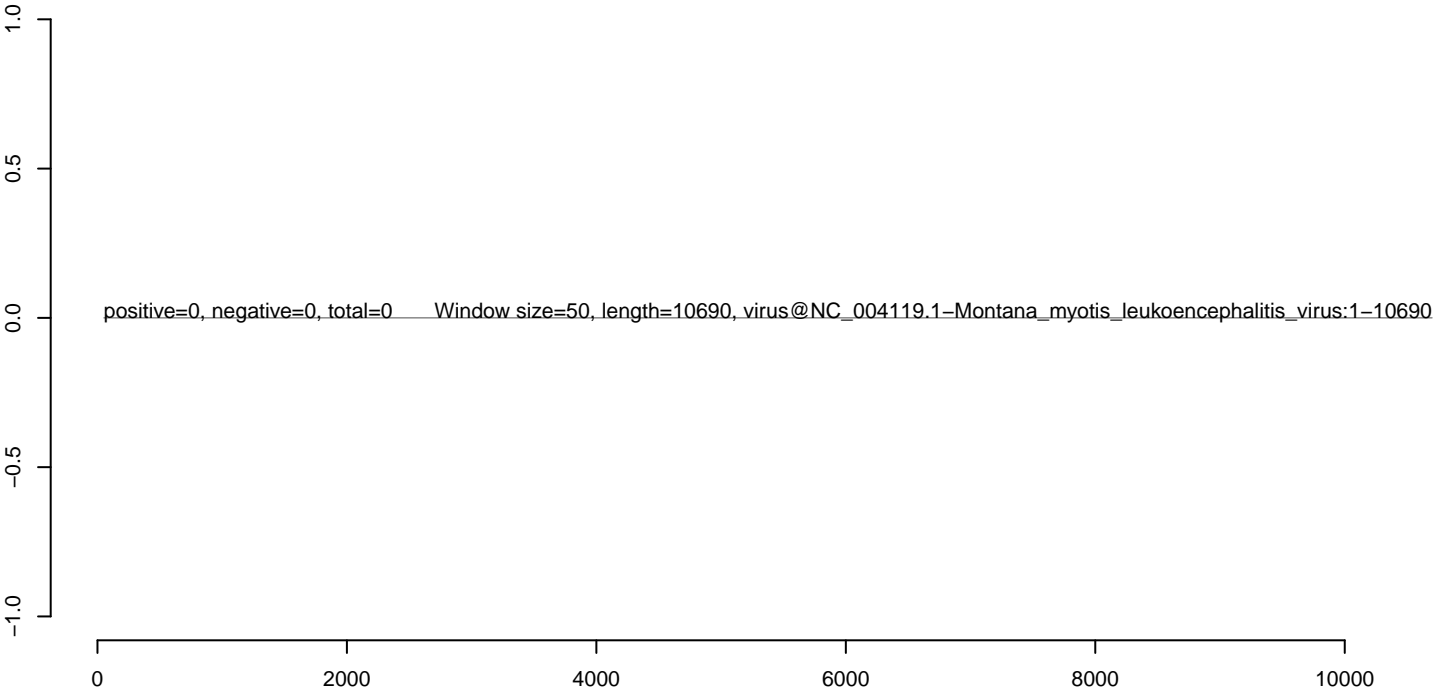
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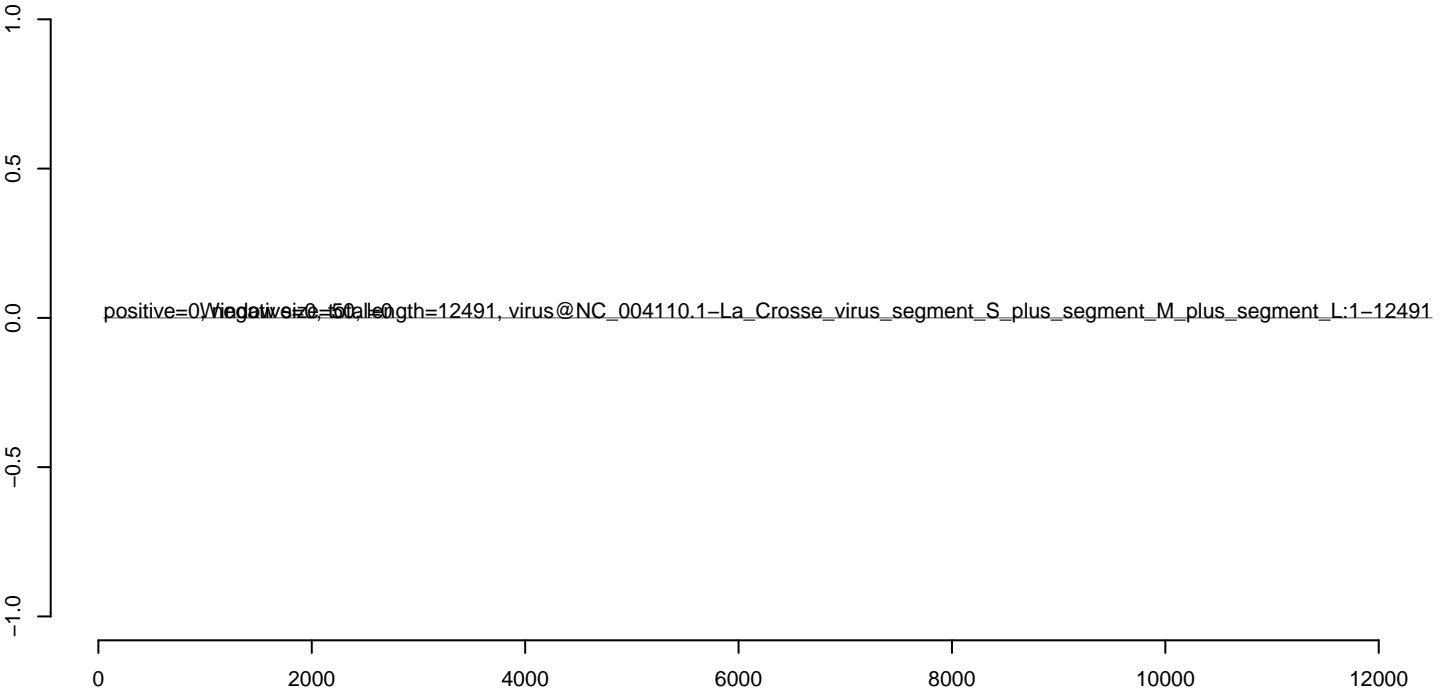
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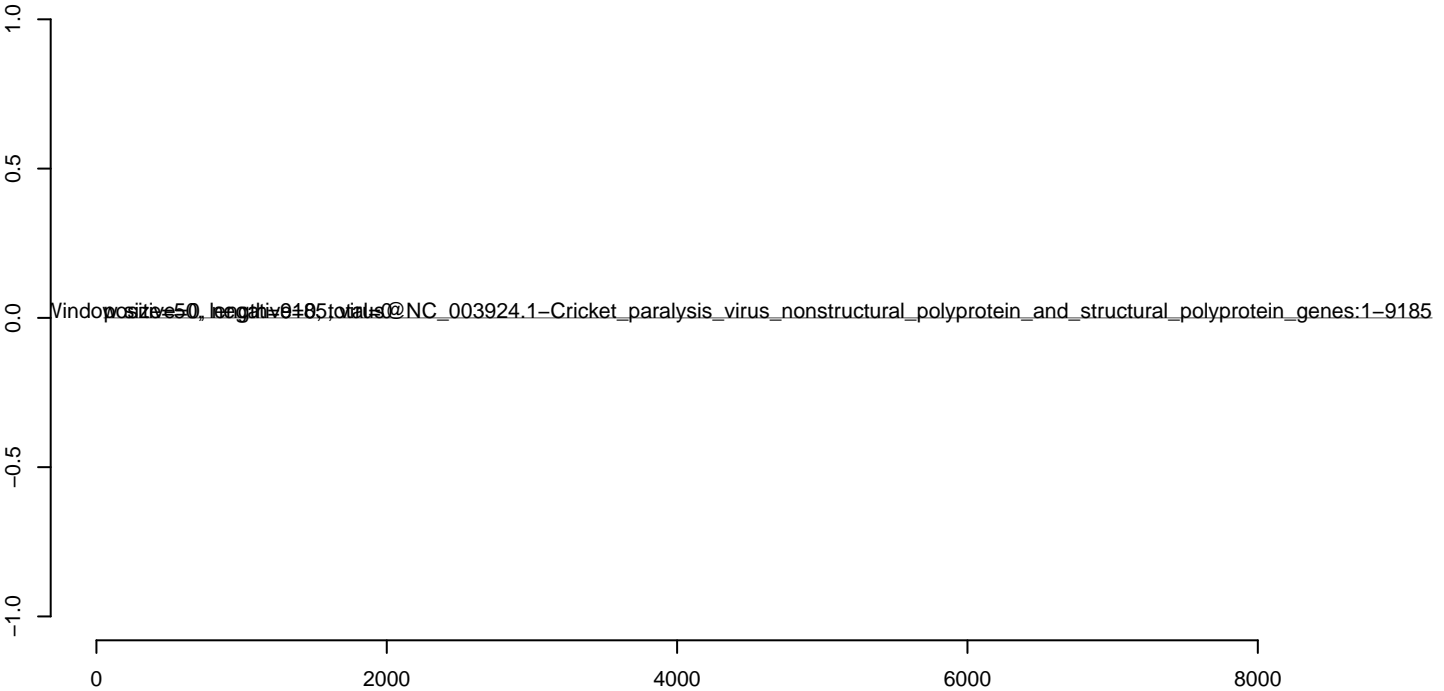
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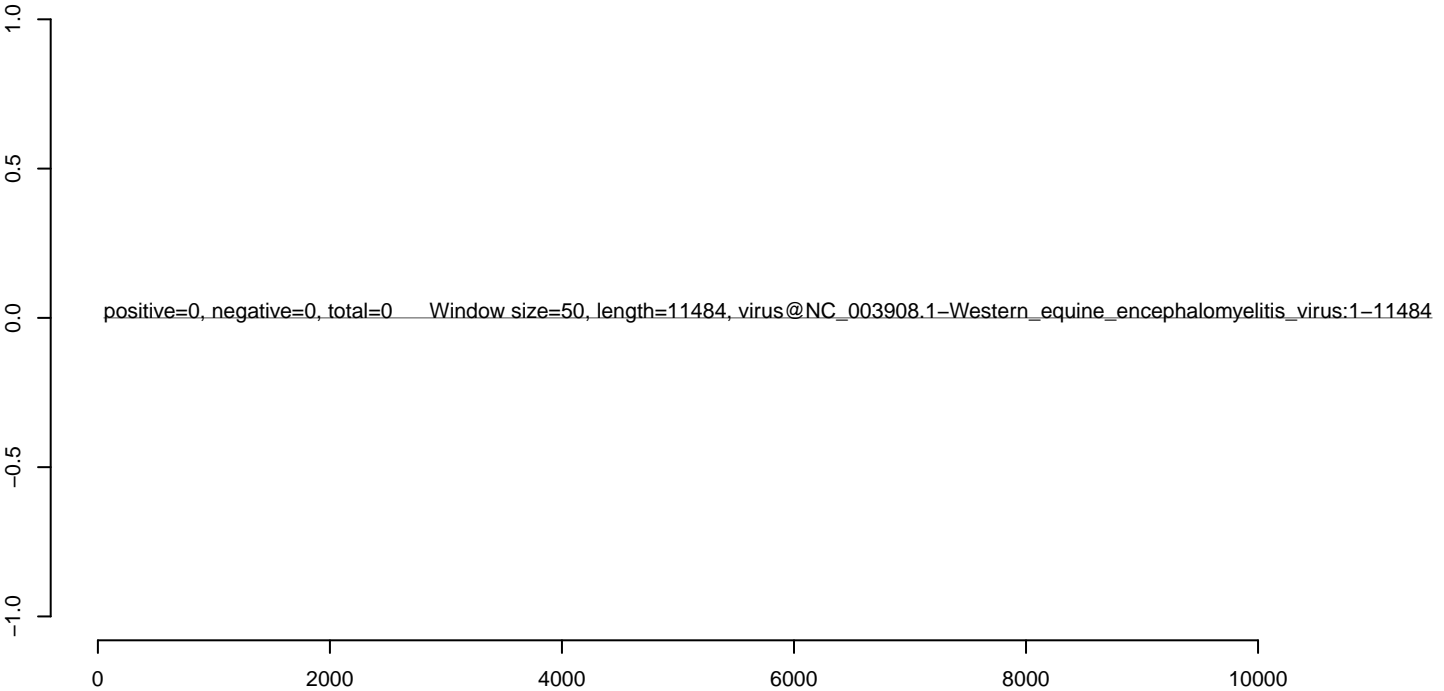
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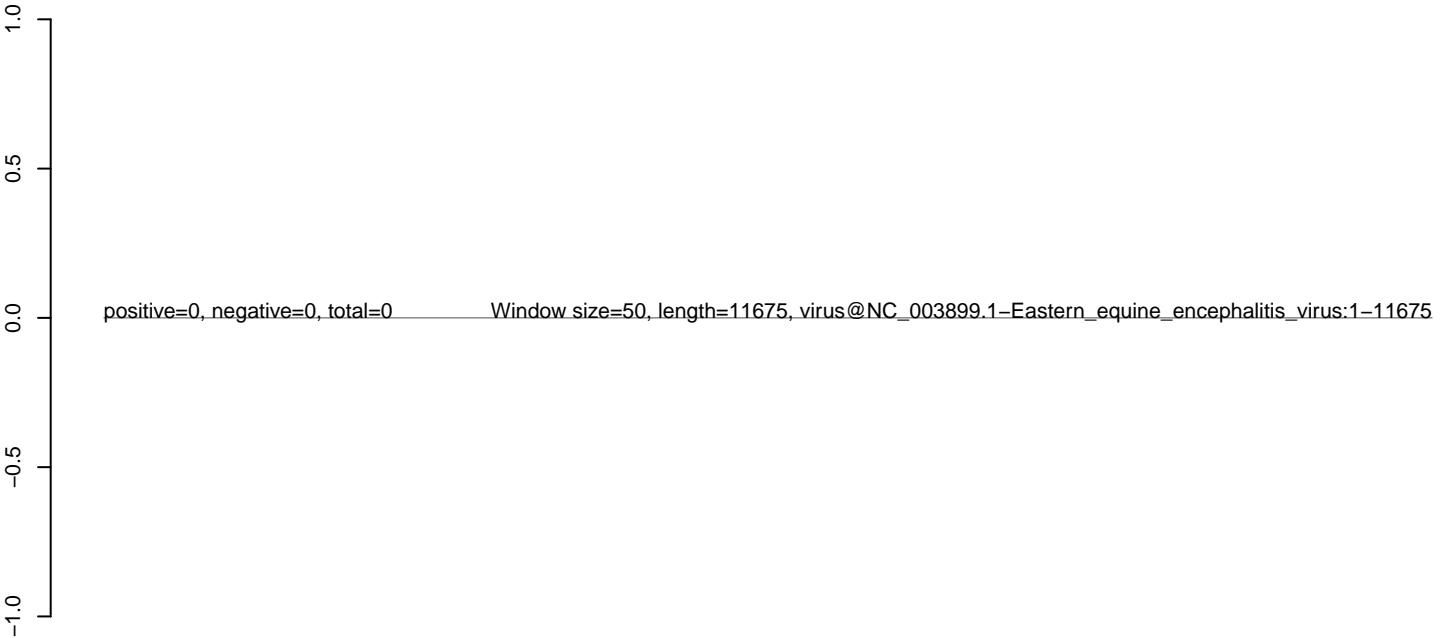
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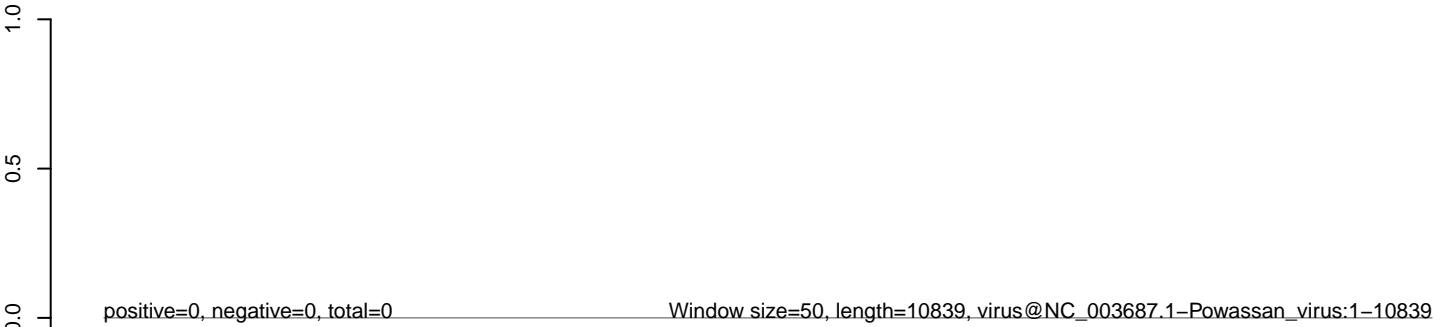
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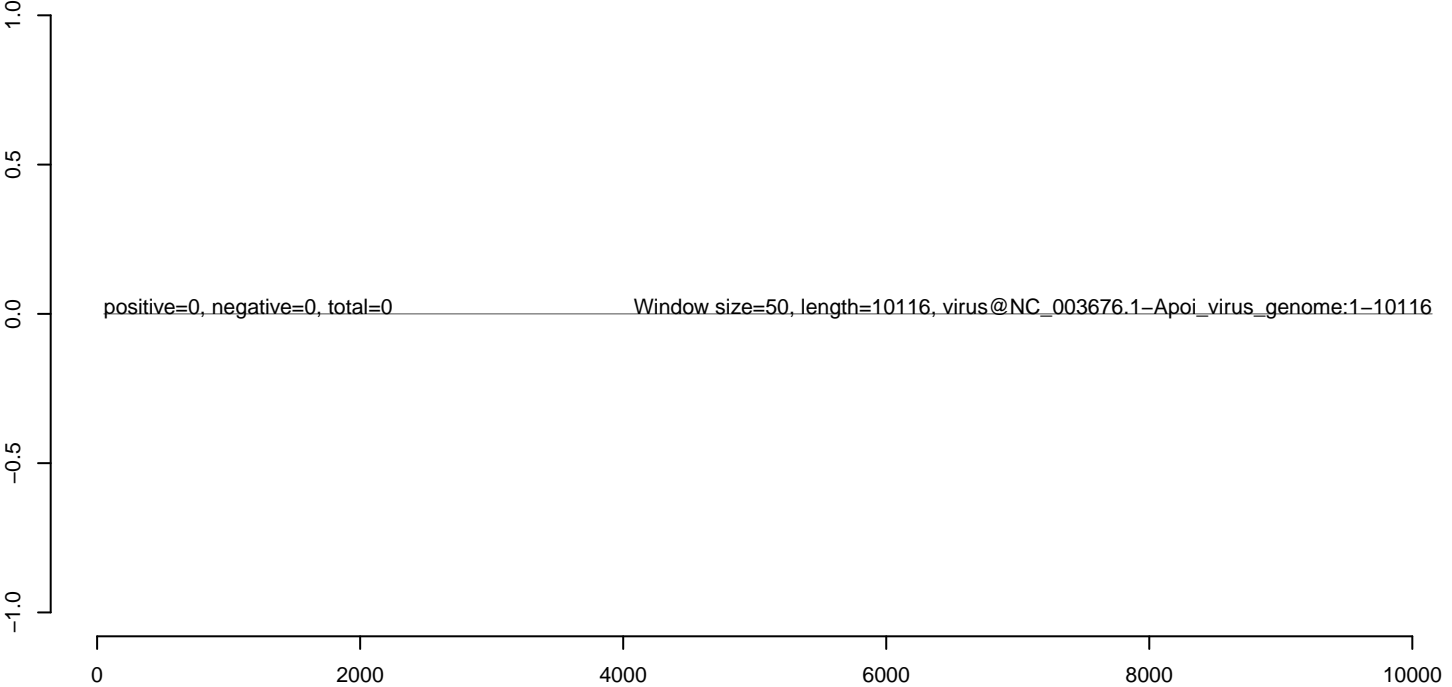
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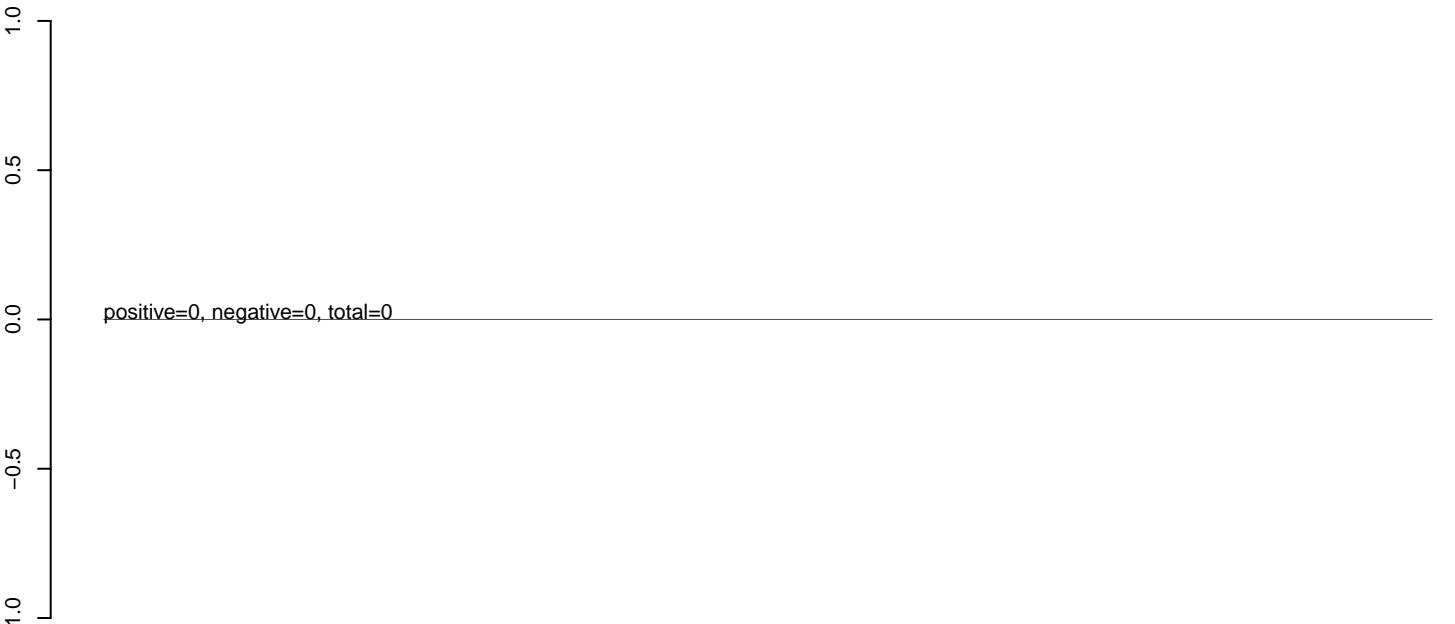


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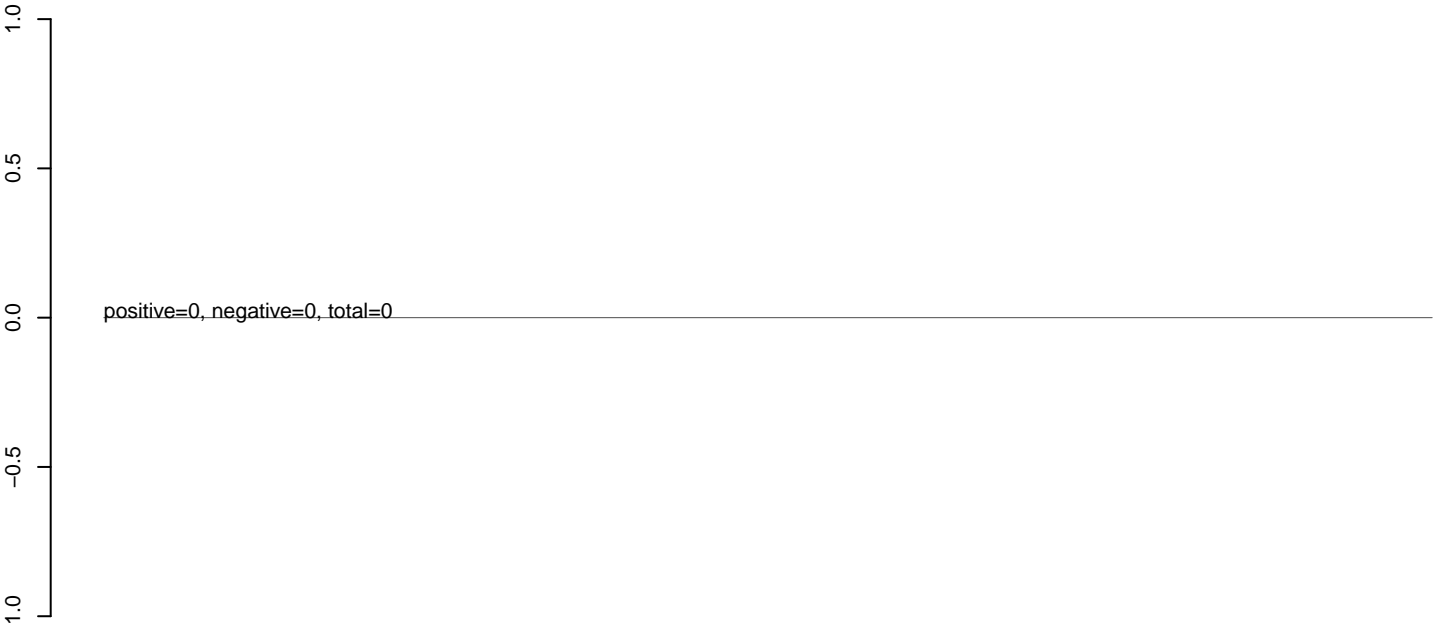




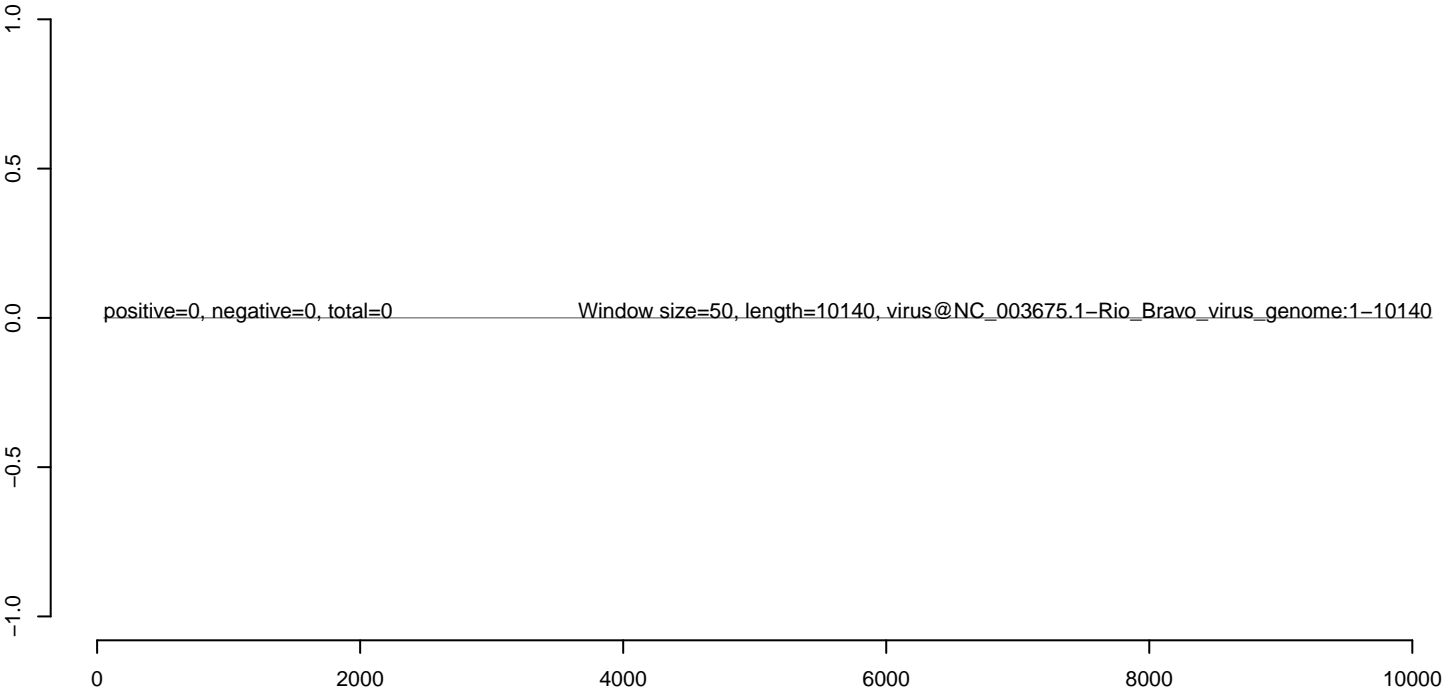
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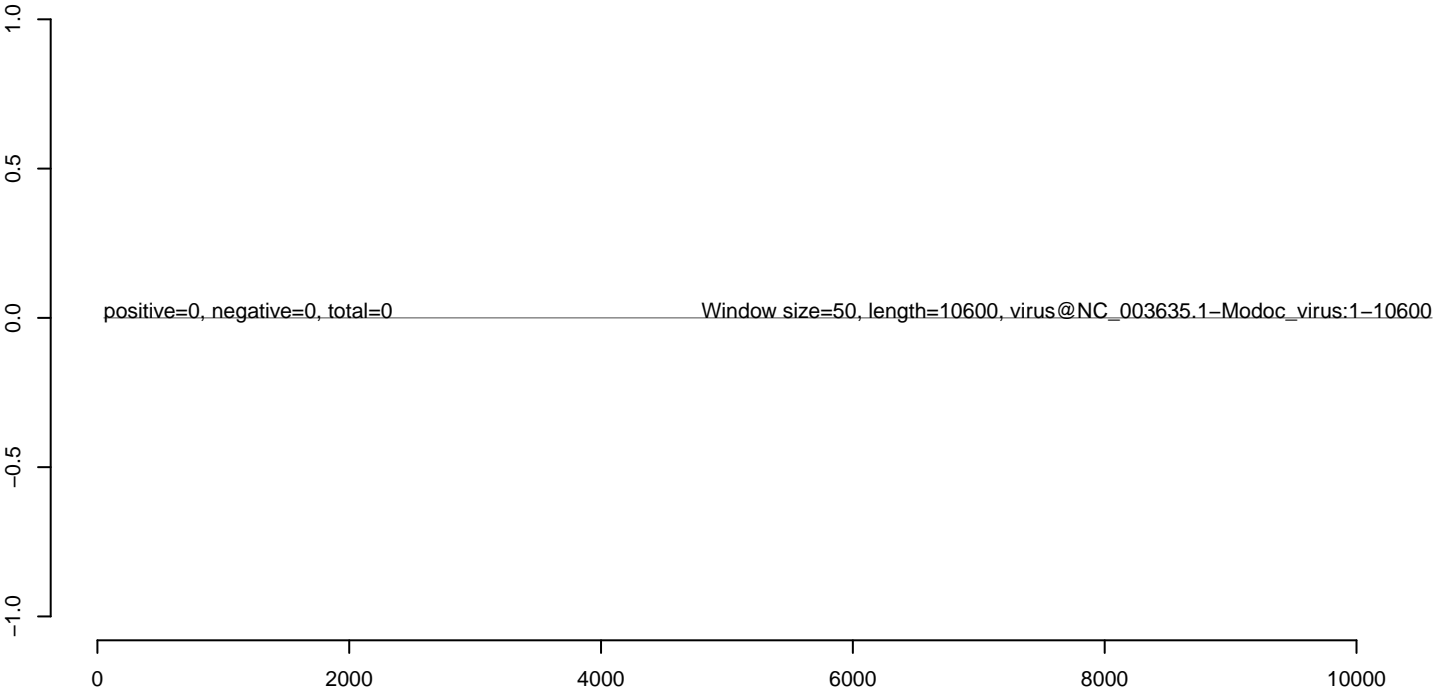
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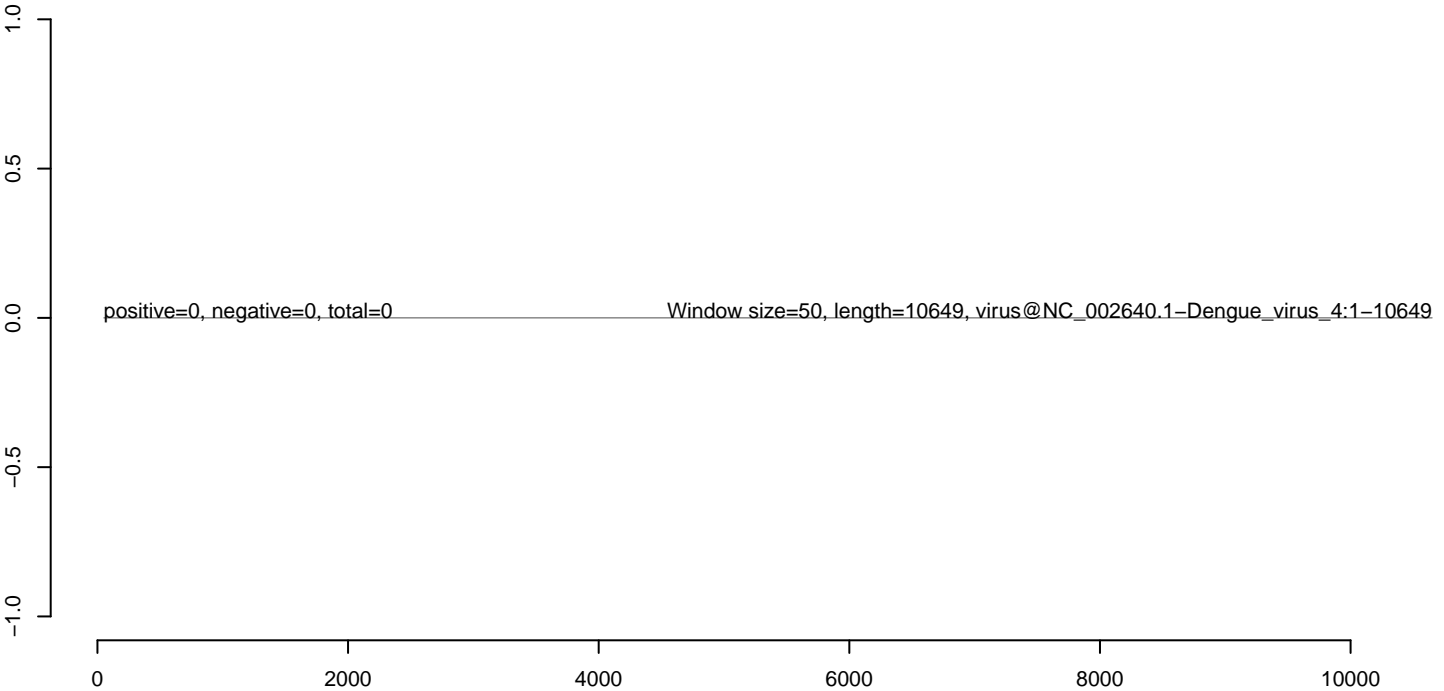
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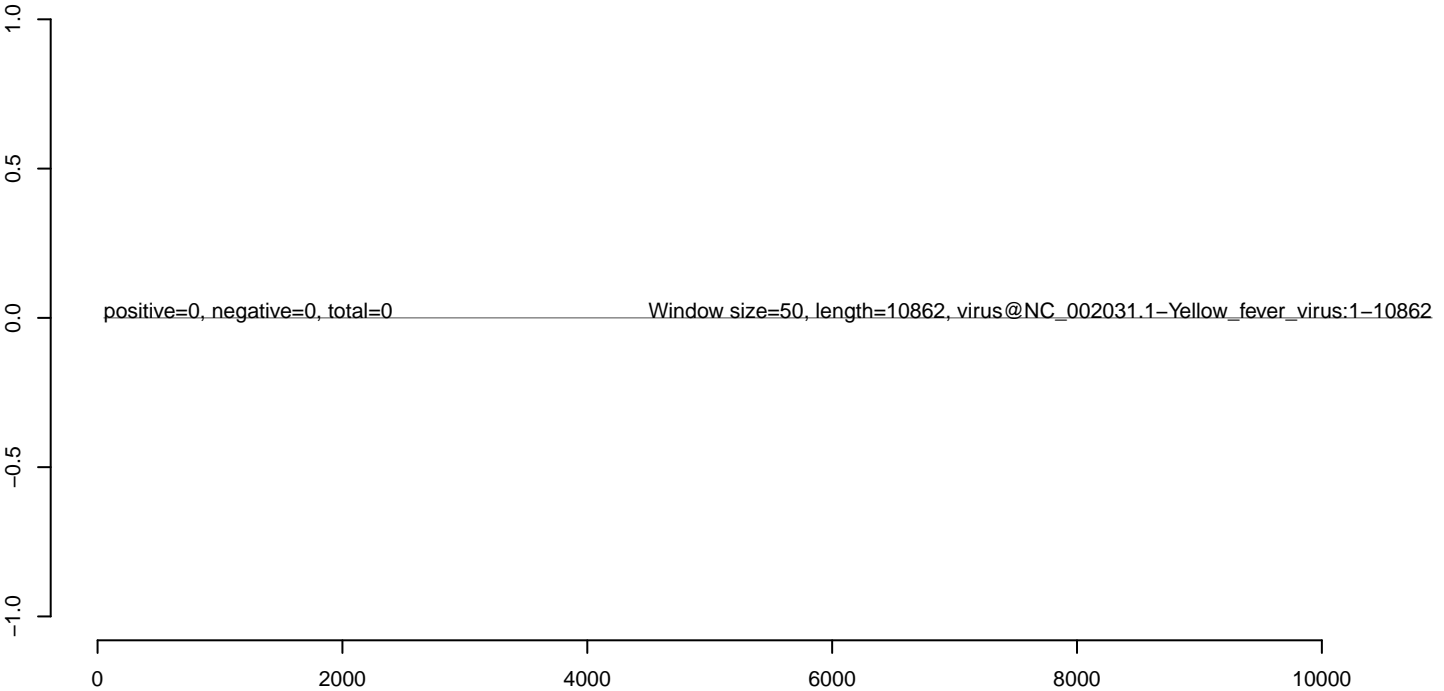
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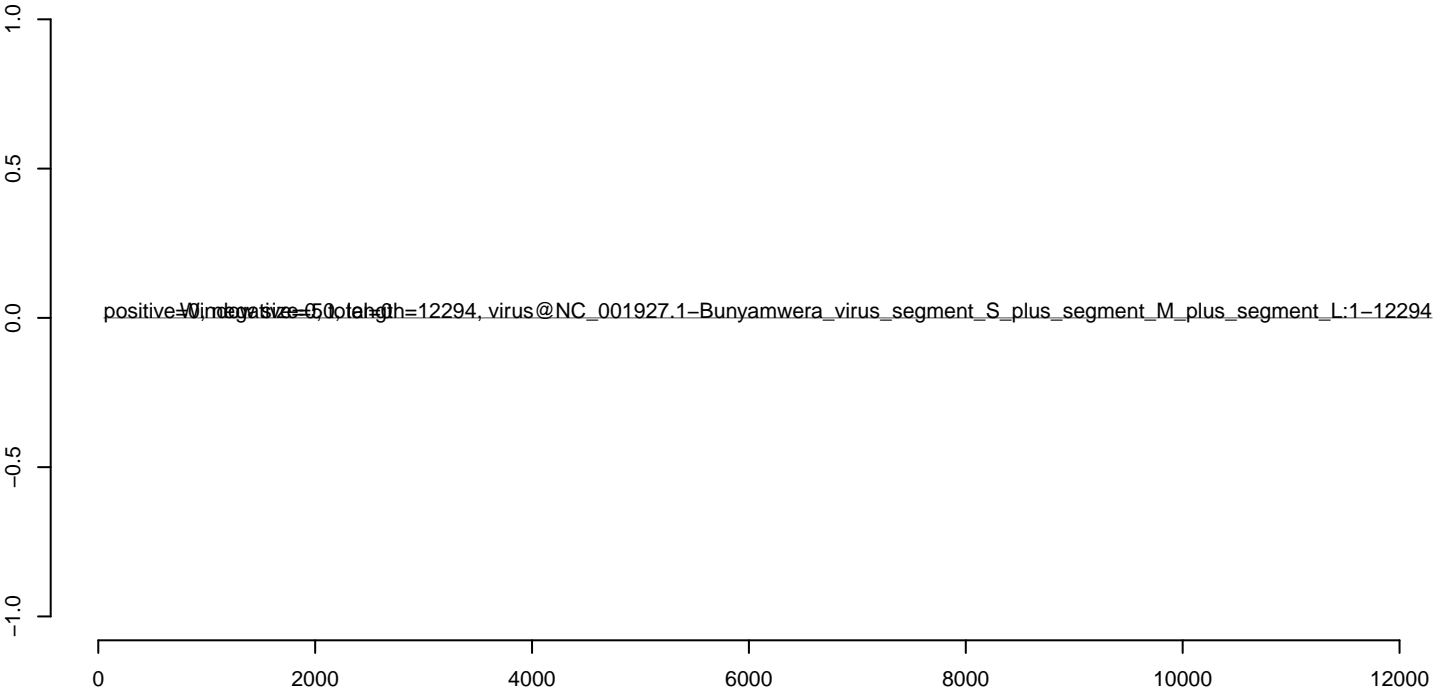
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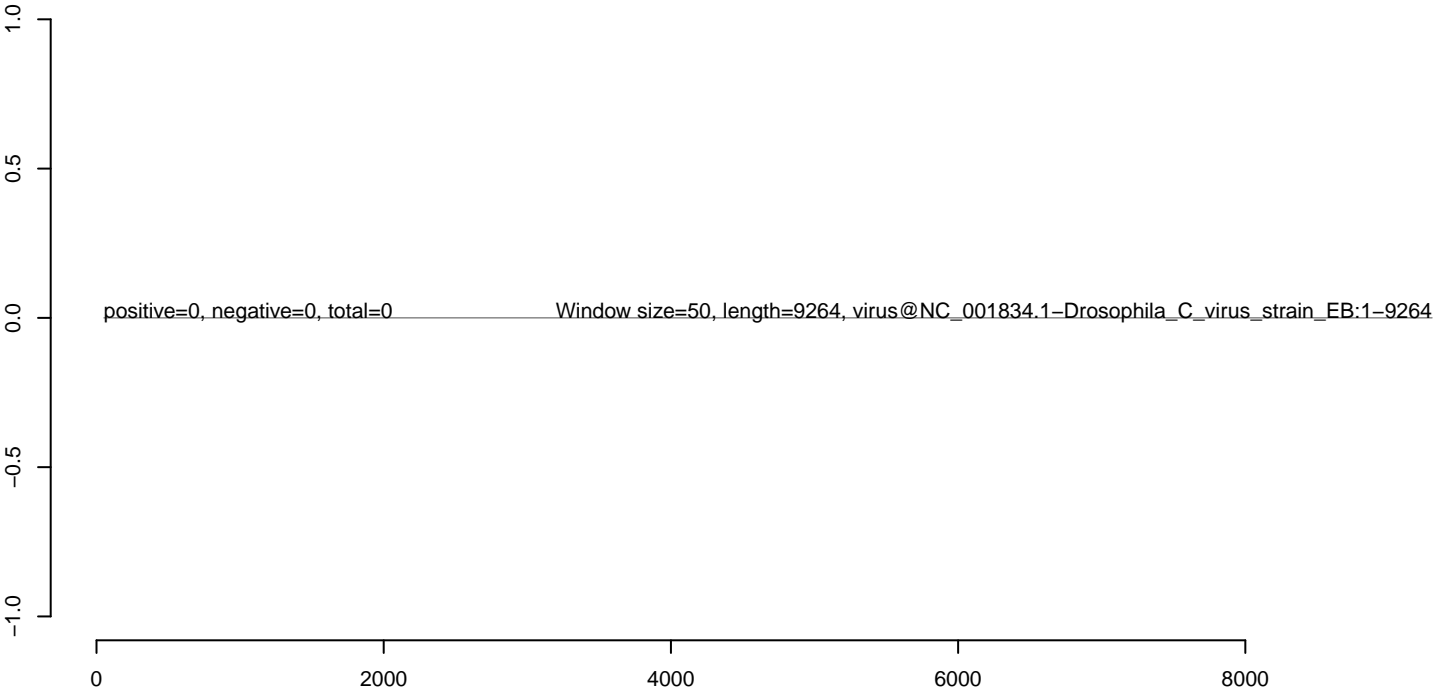
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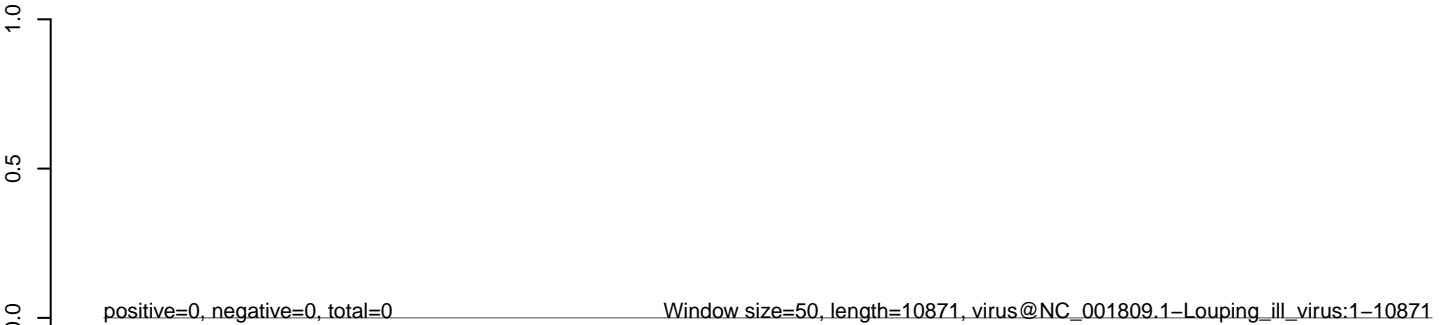
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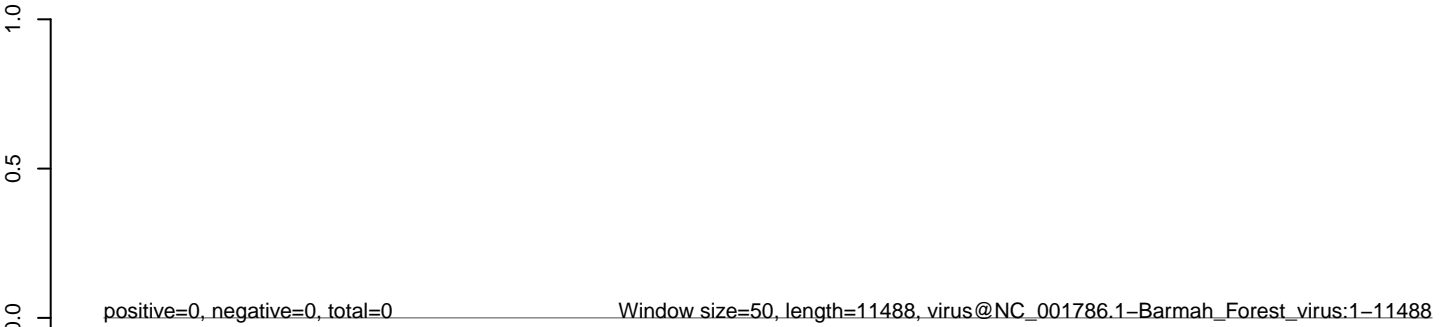
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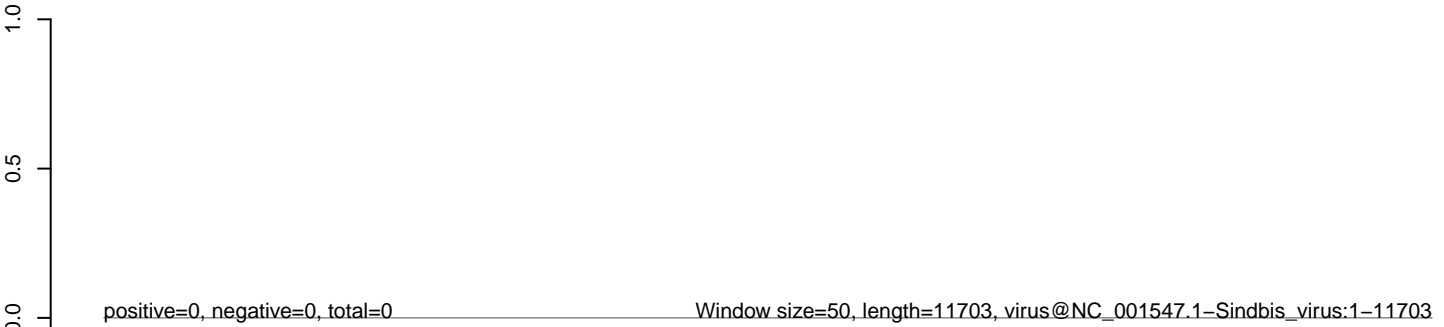
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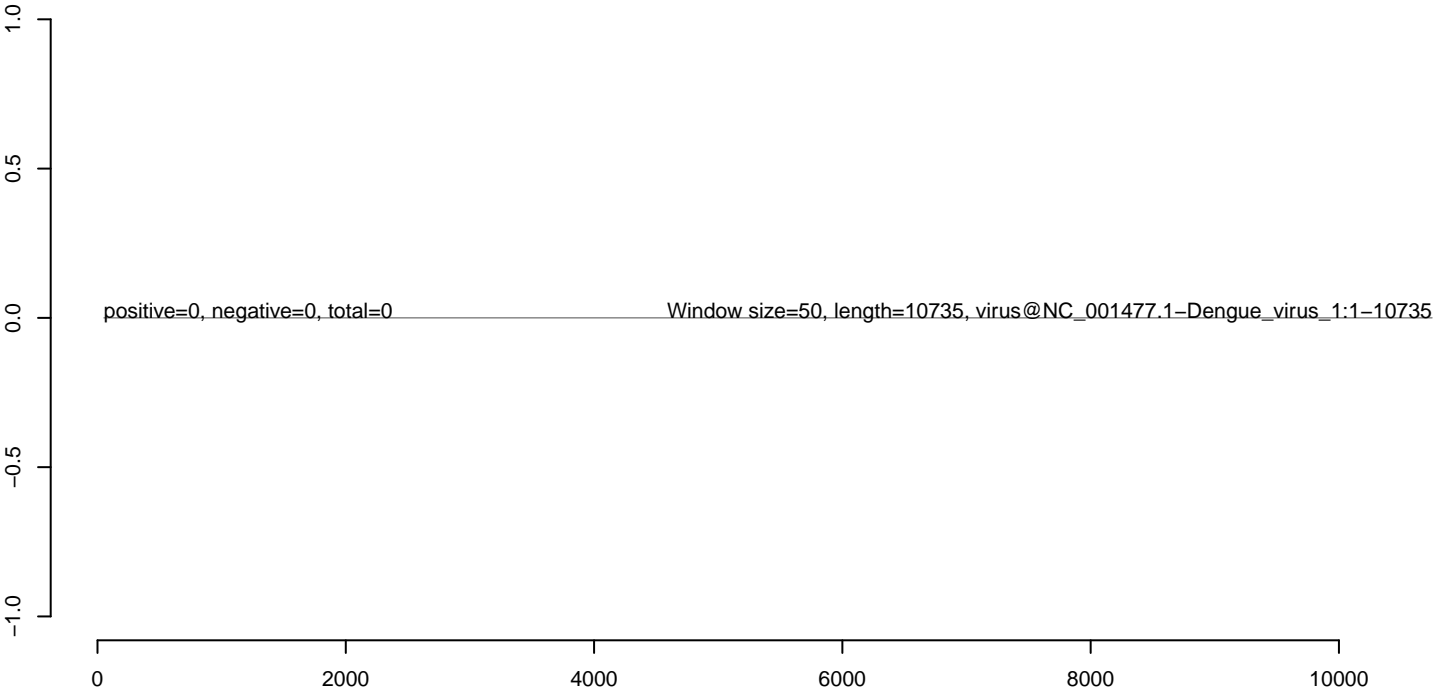
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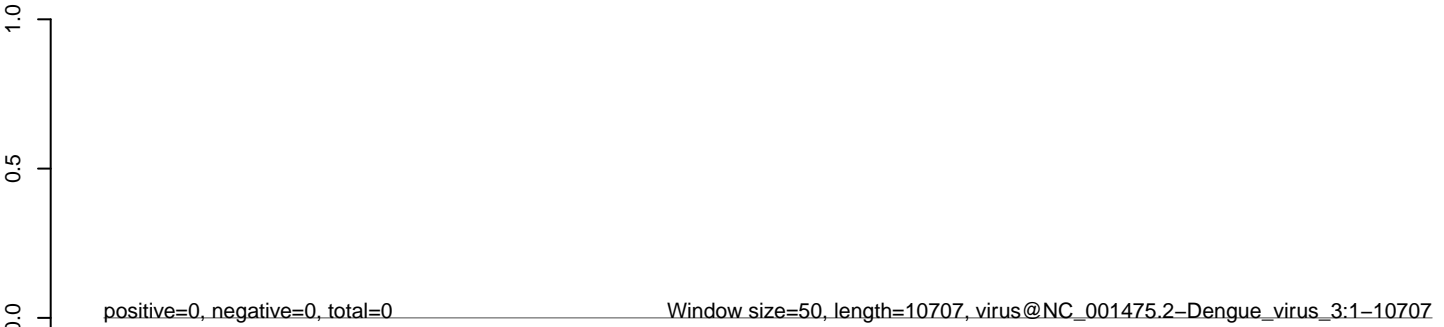
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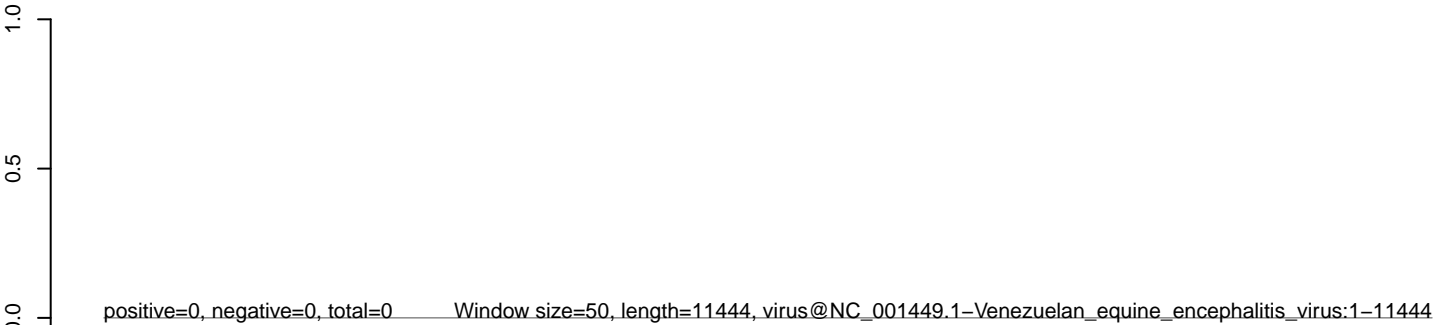
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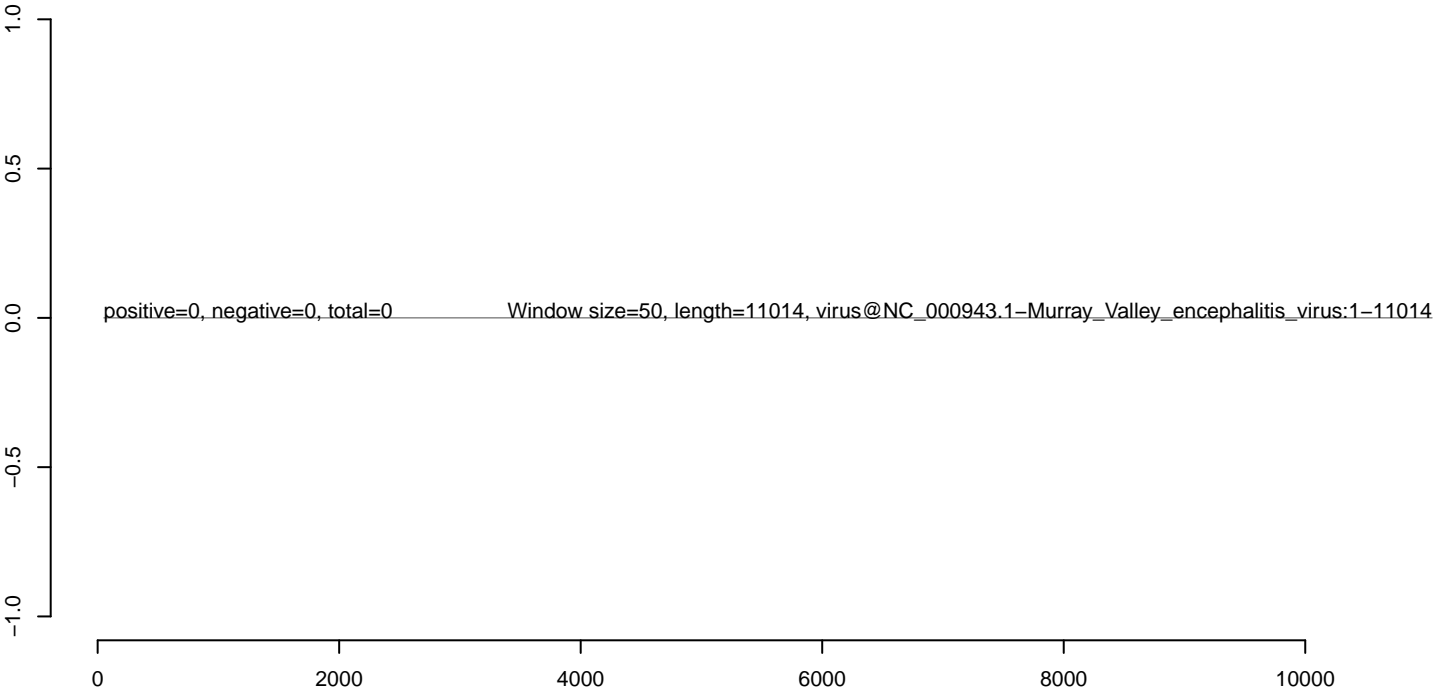
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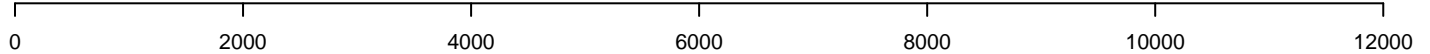
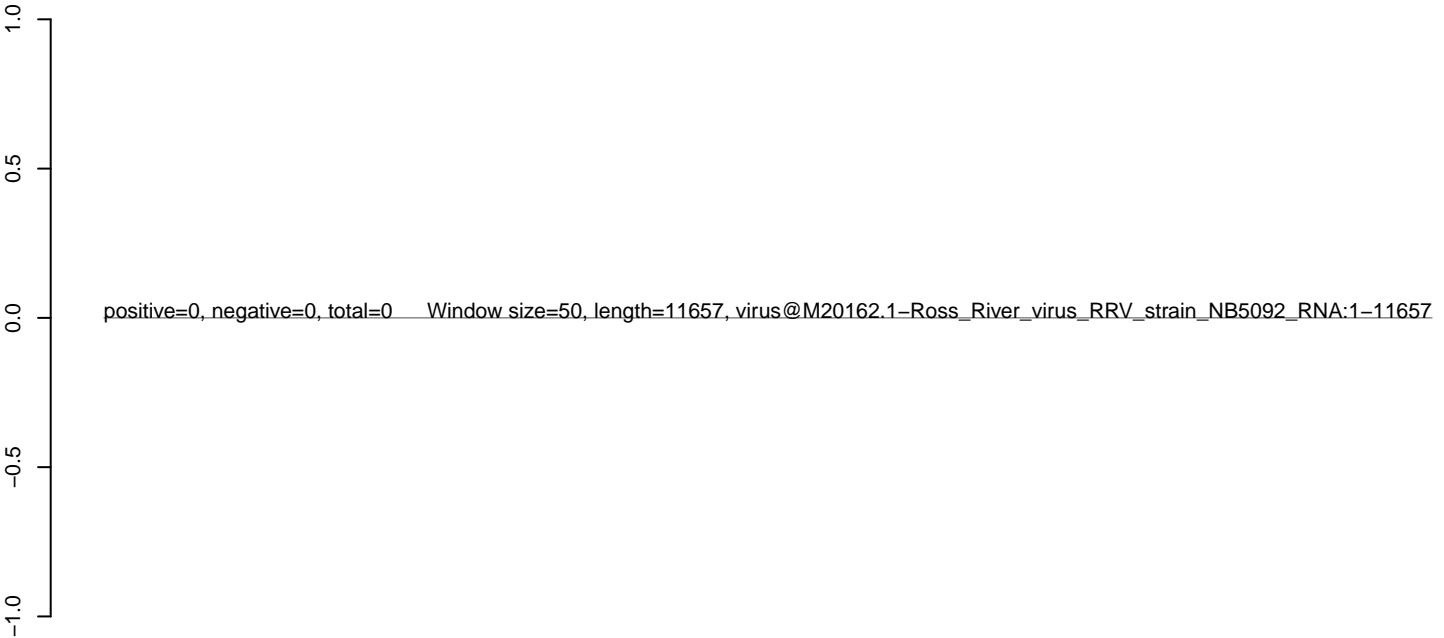
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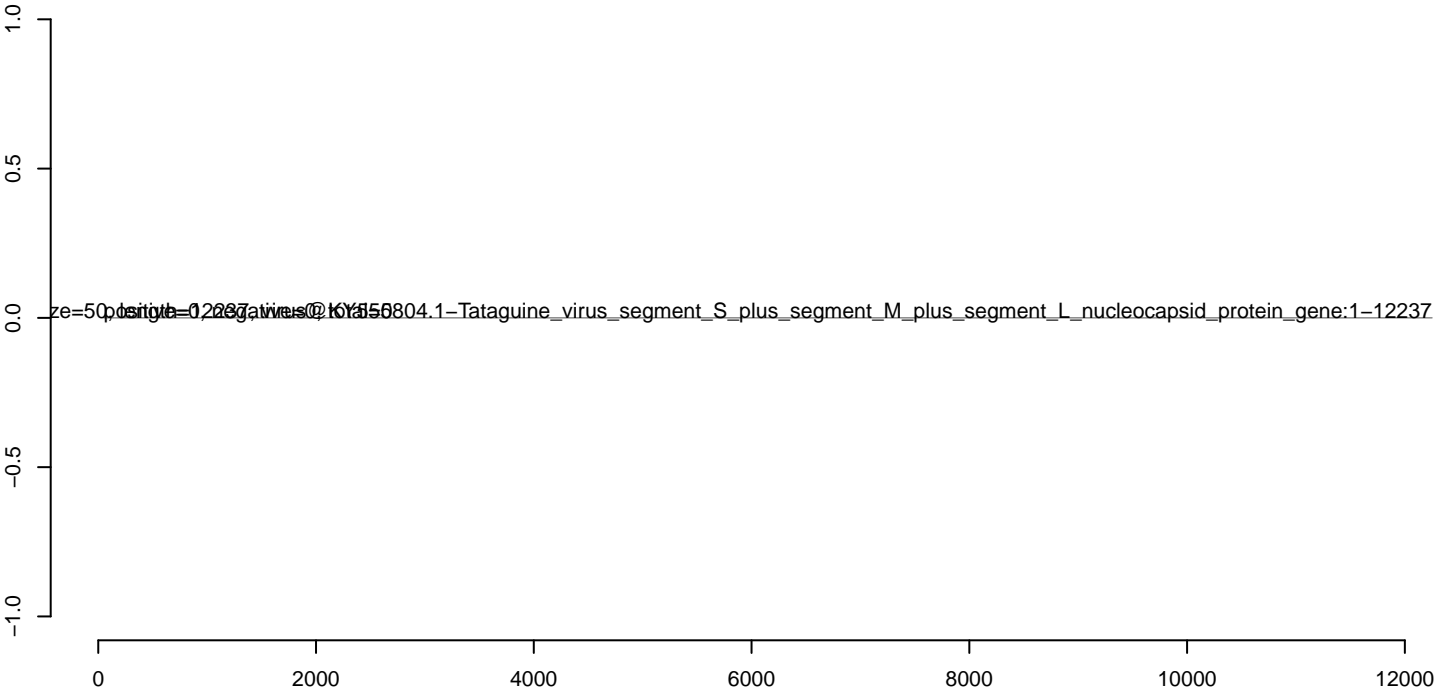
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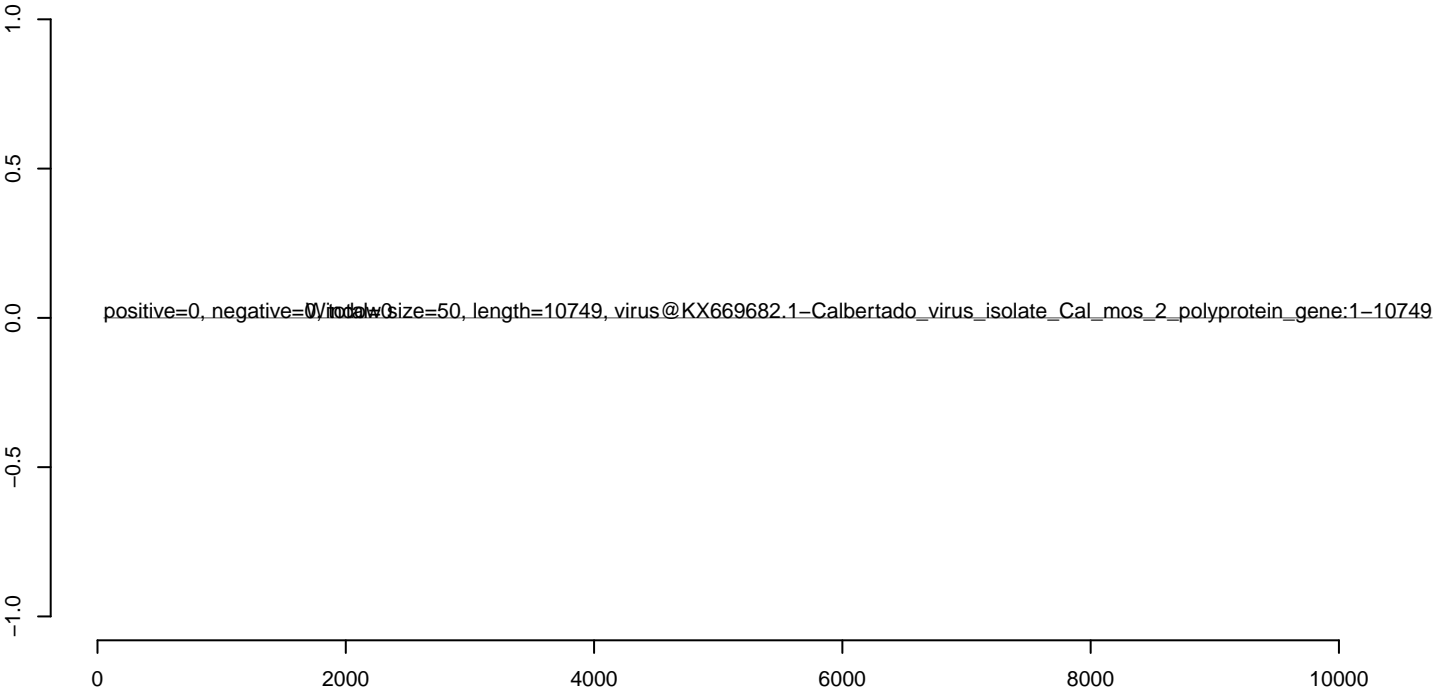
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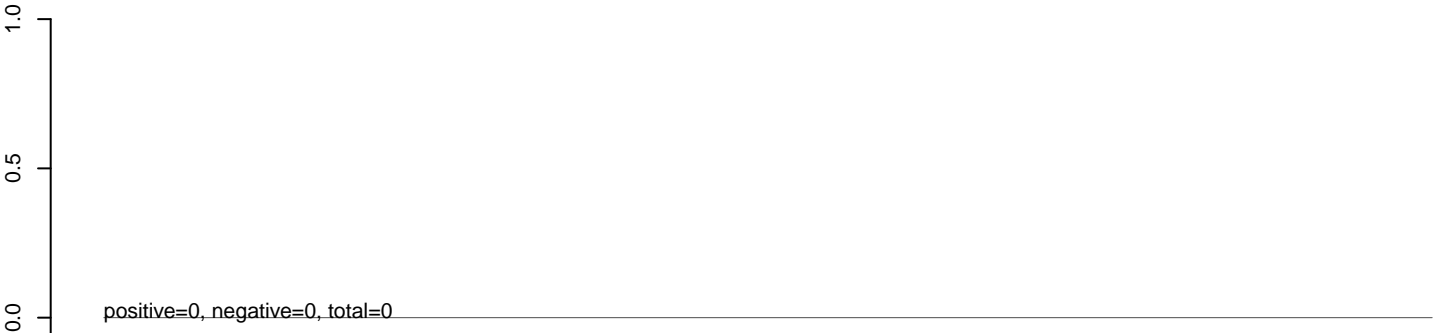




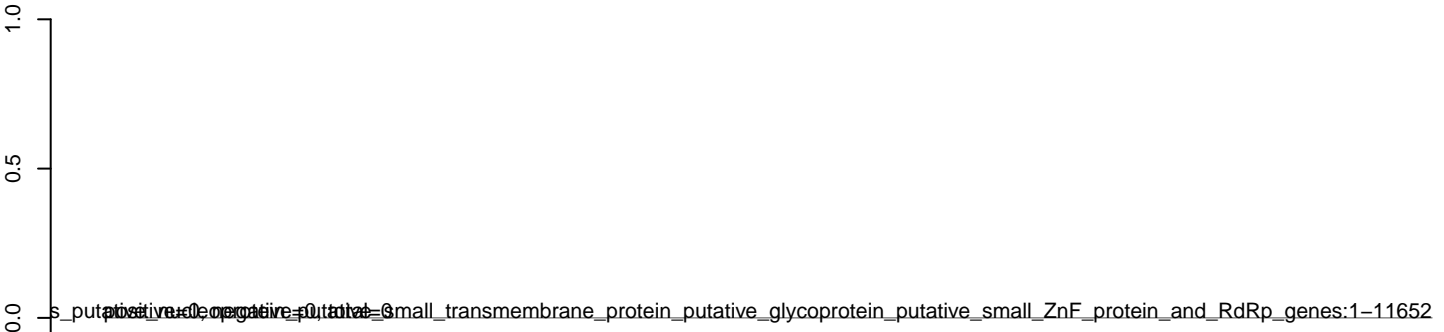
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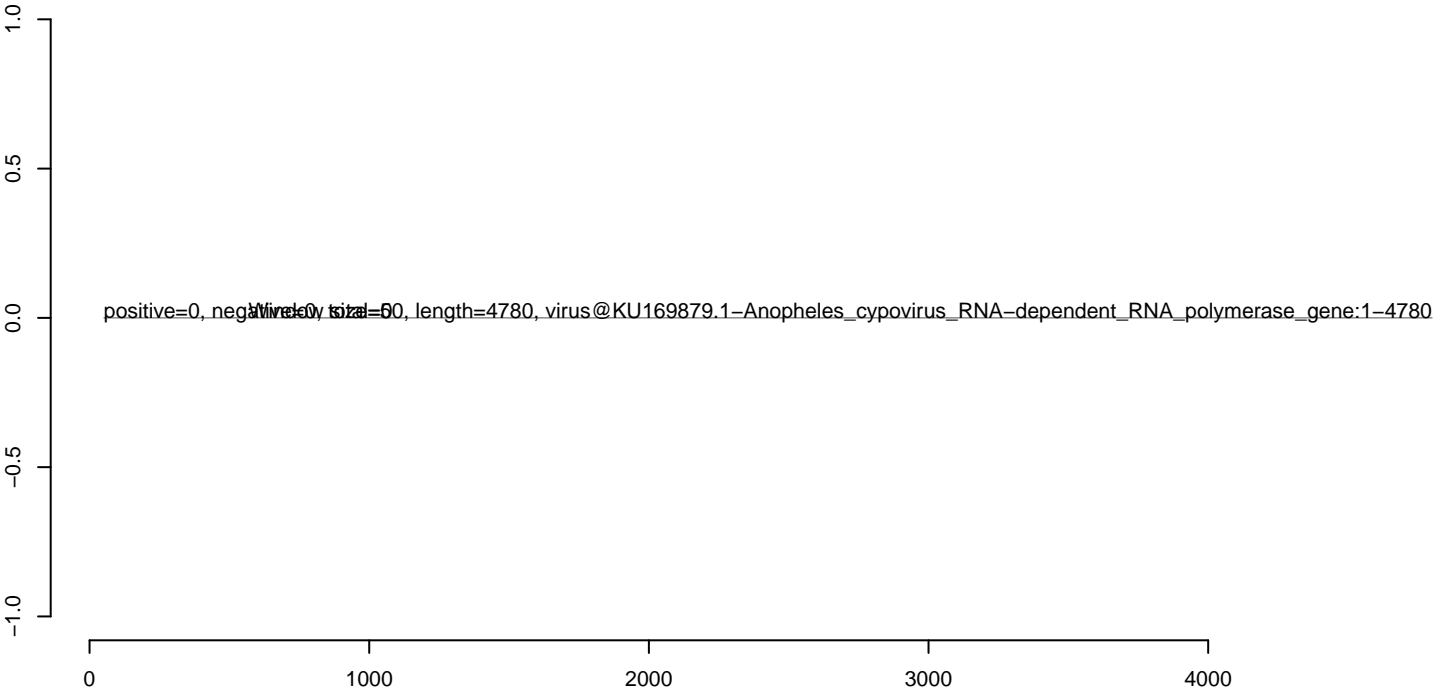
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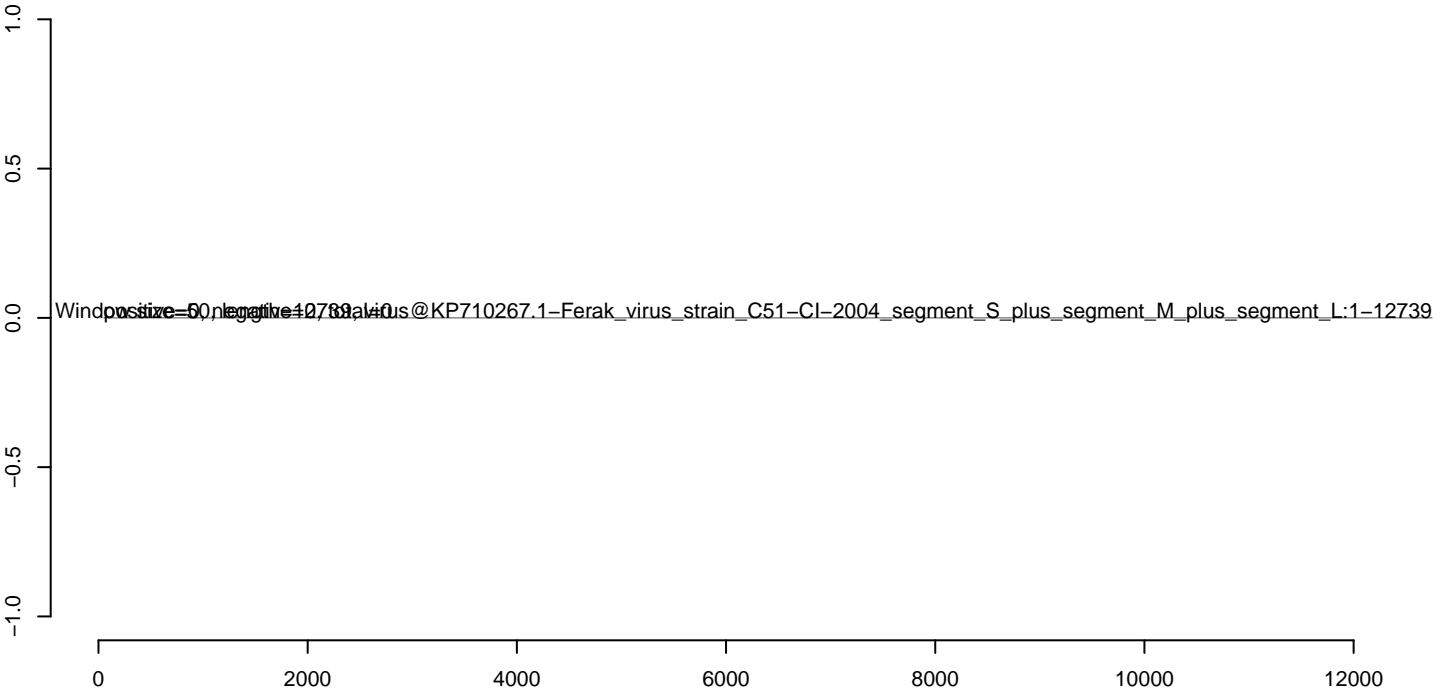
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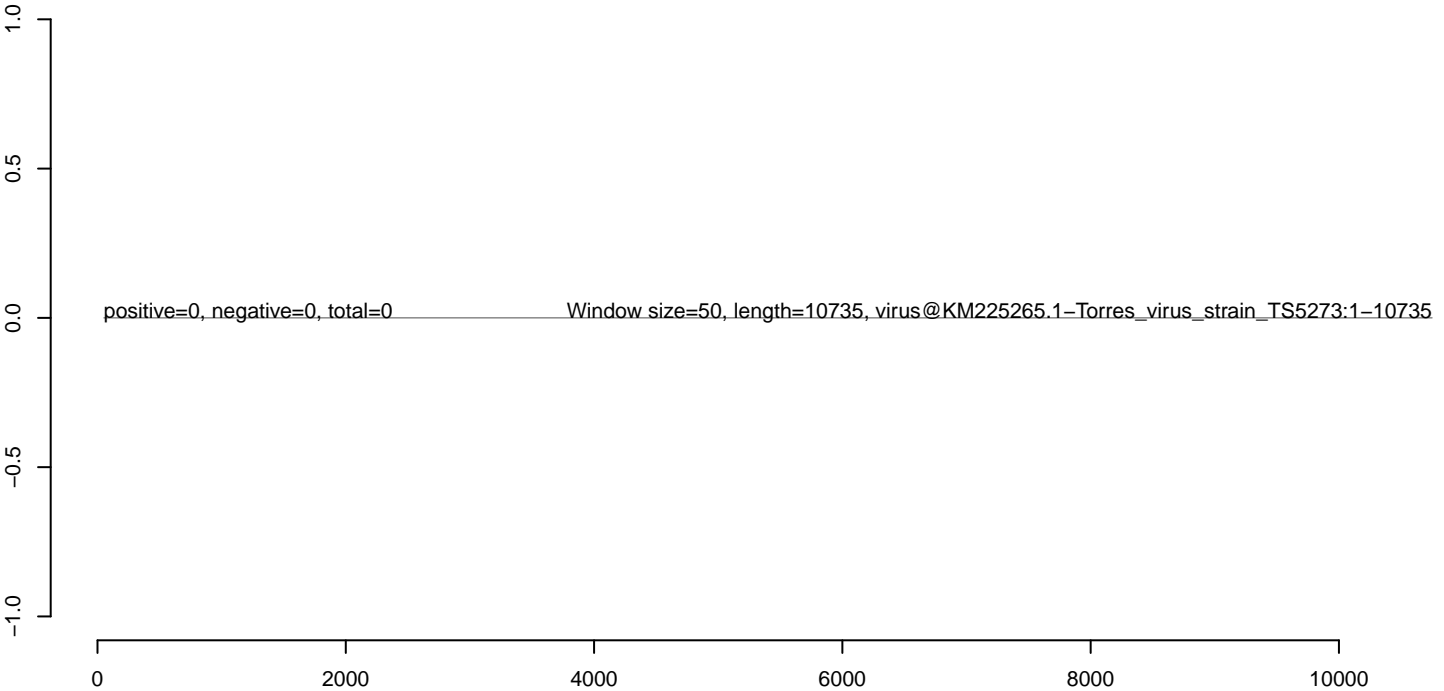
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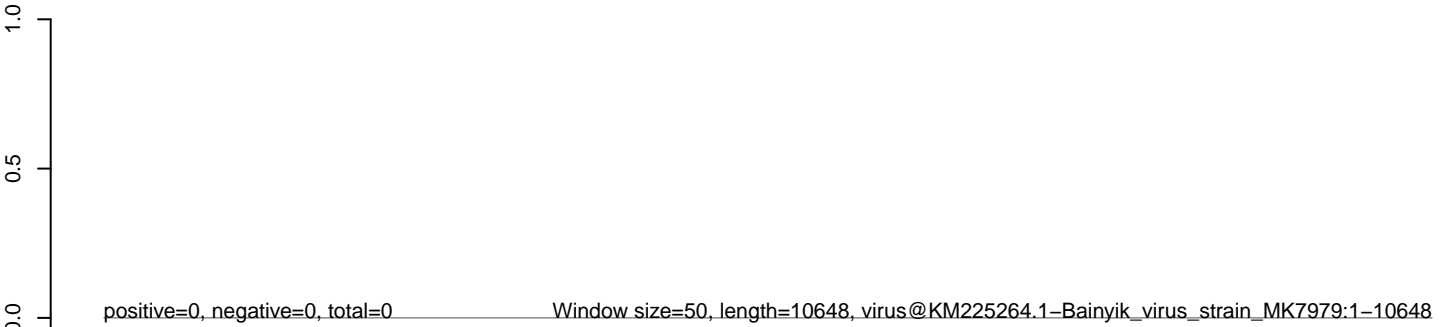
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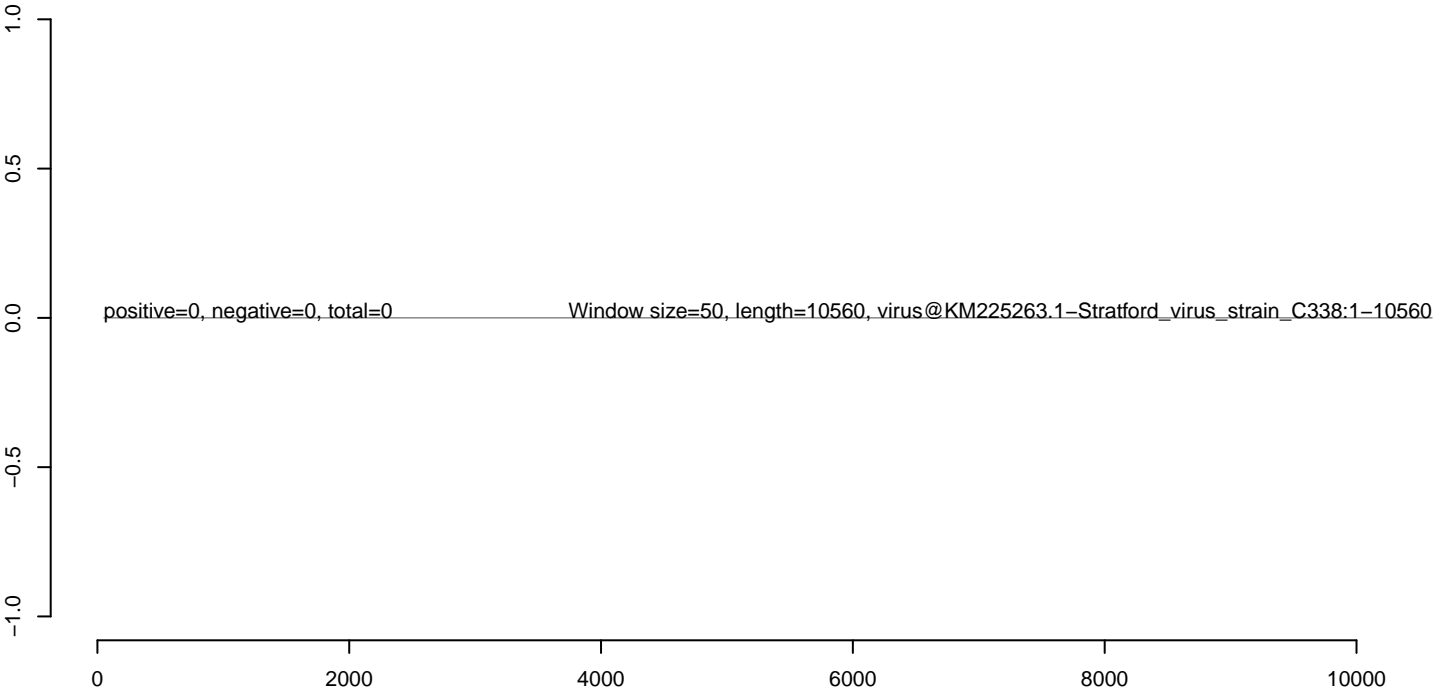
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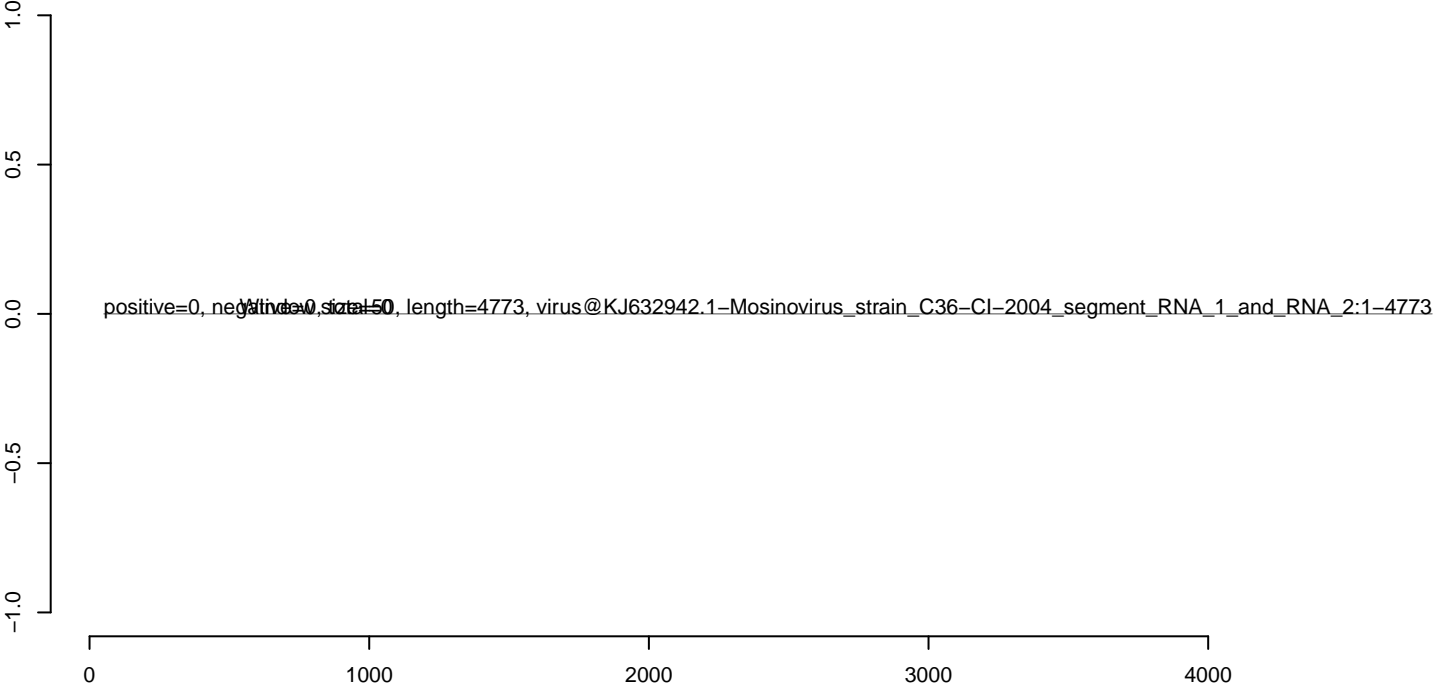
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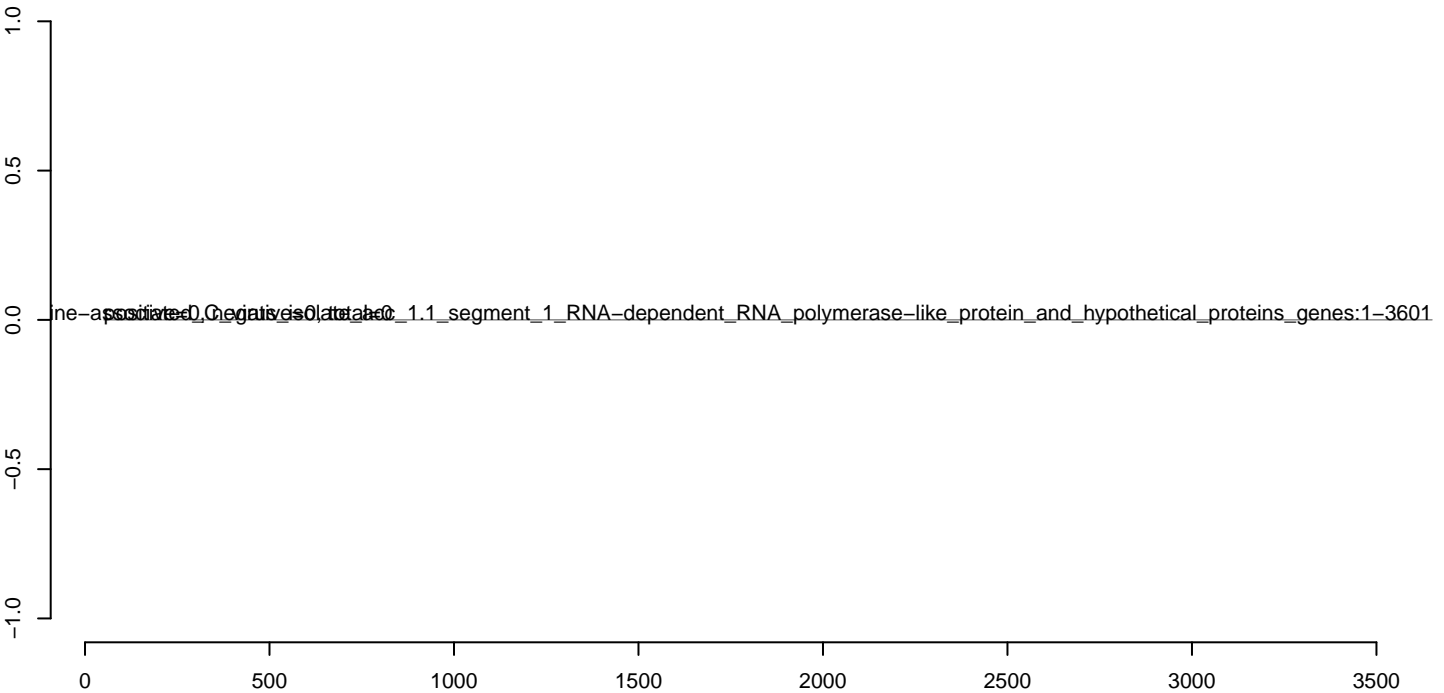
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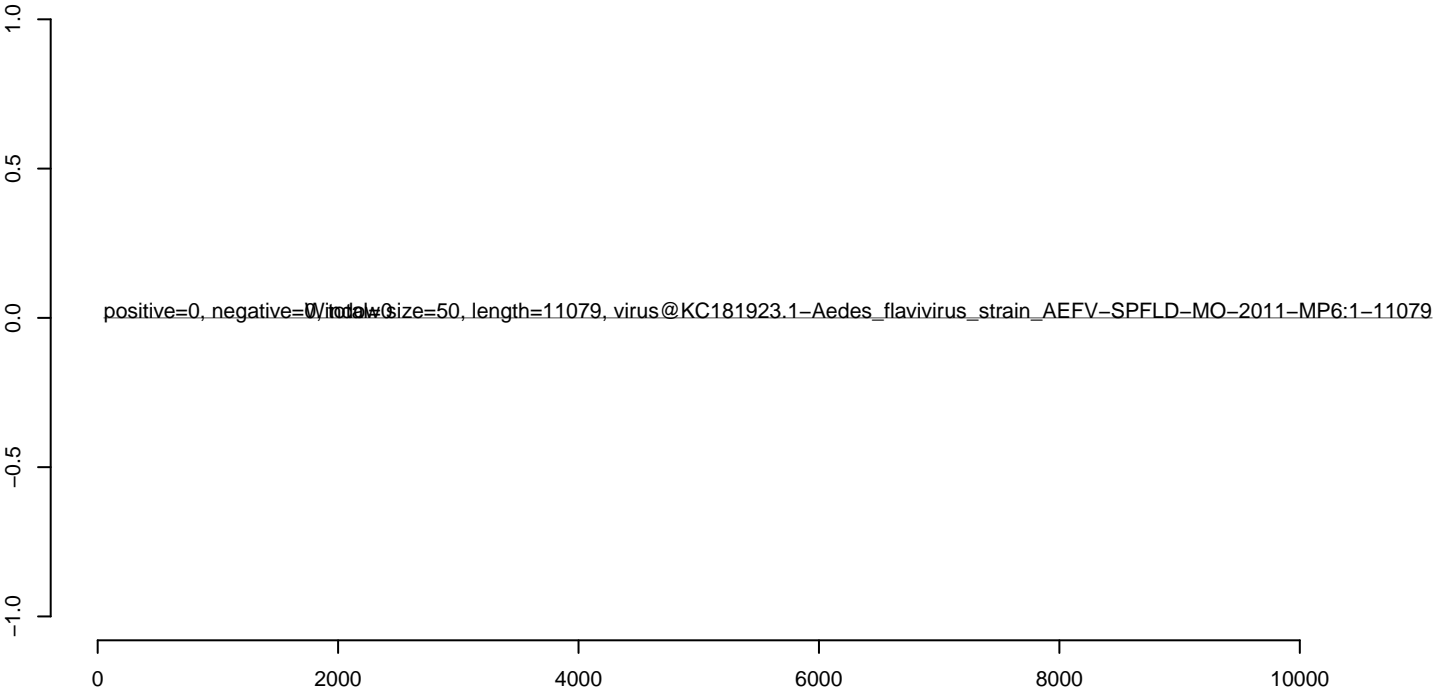
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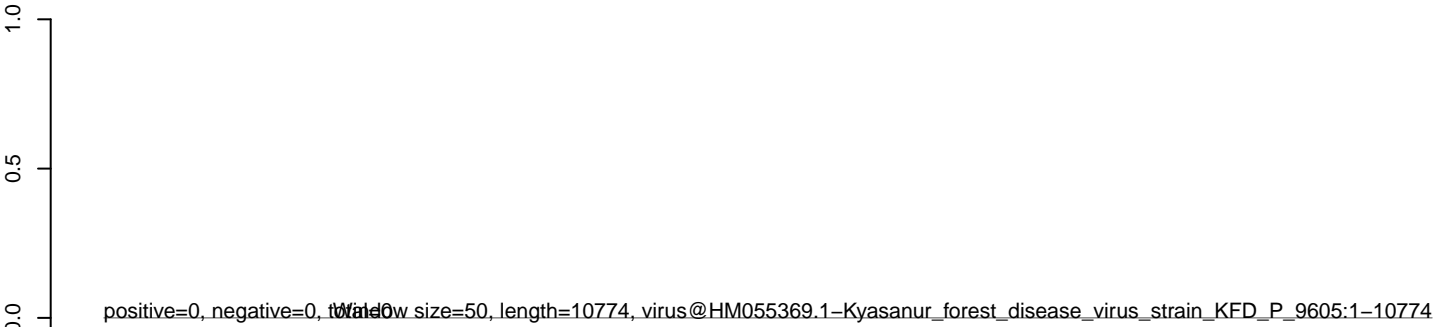
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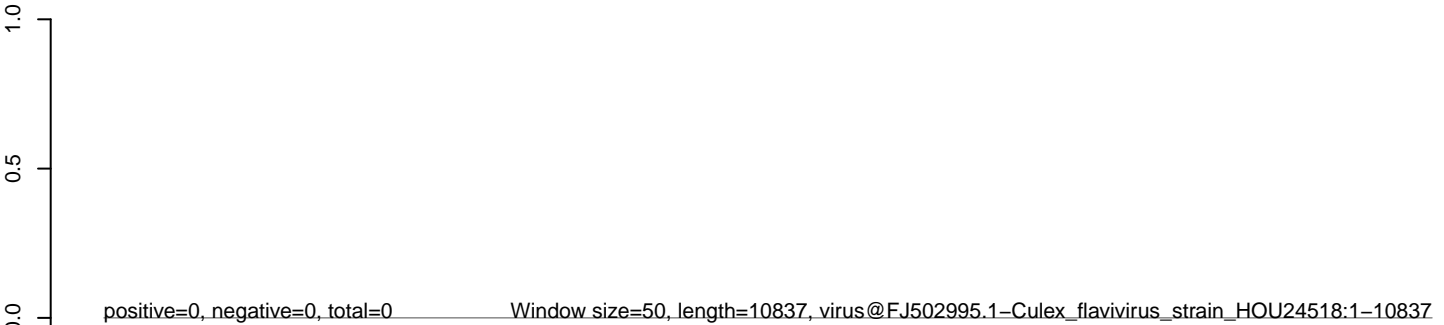
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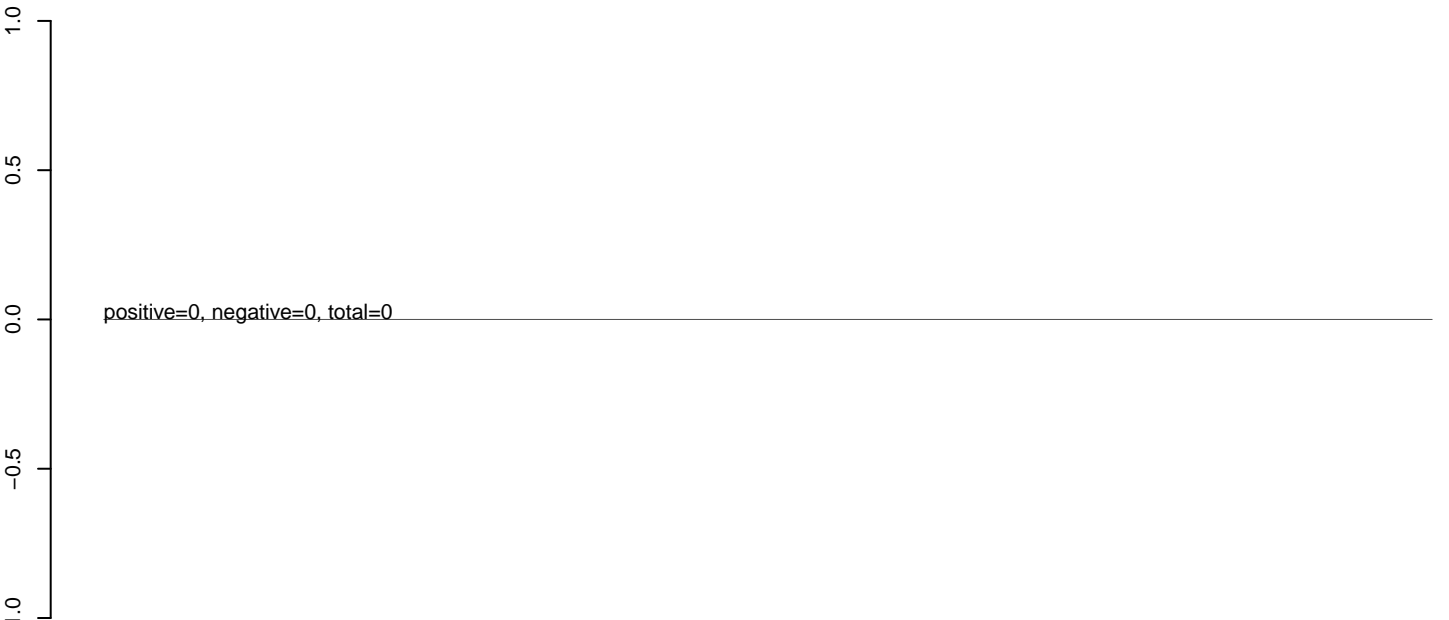


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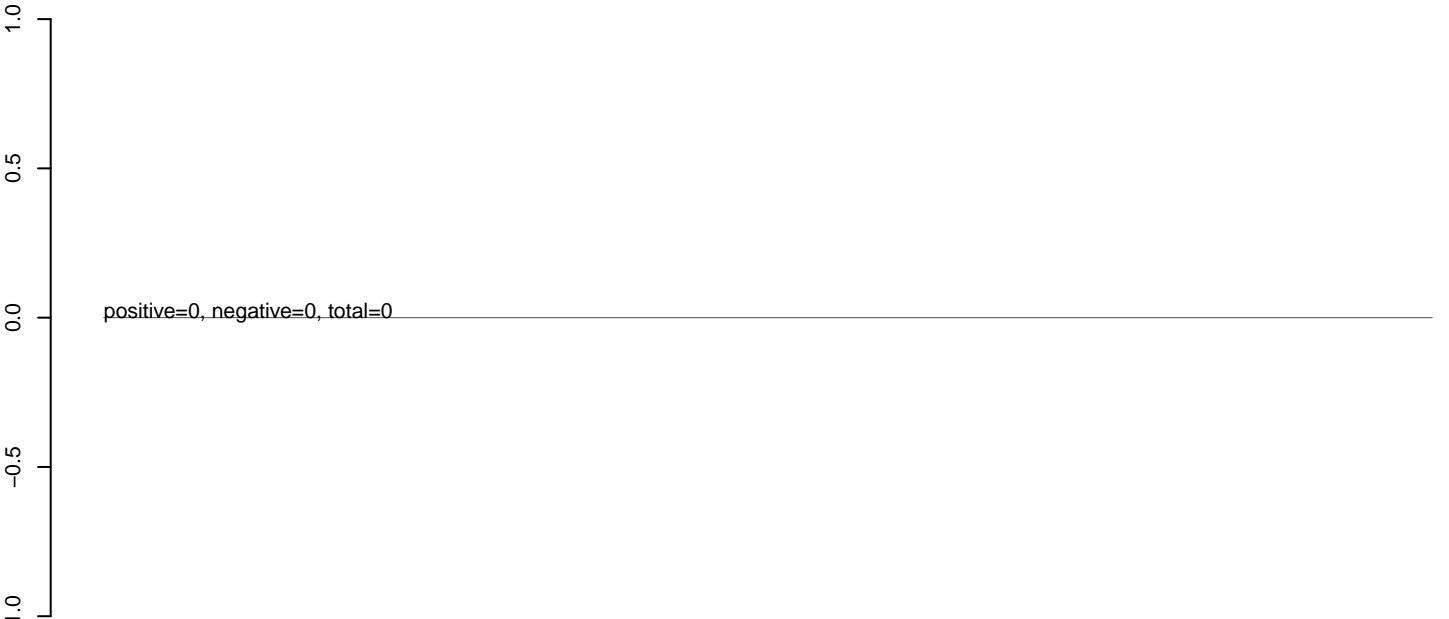


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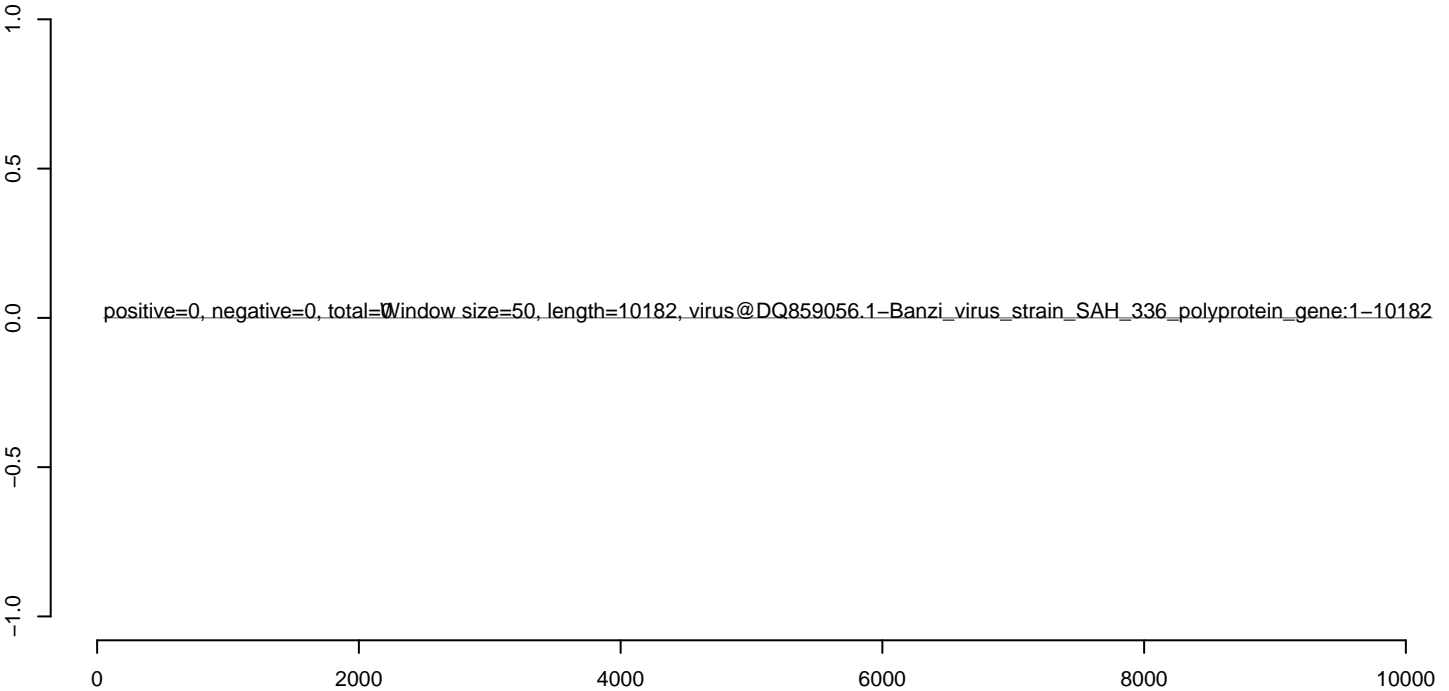
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**AnGam\_Sua5b\_cells.24\_35.rep**



**AnGam\_Sua5b\_cells.rep**



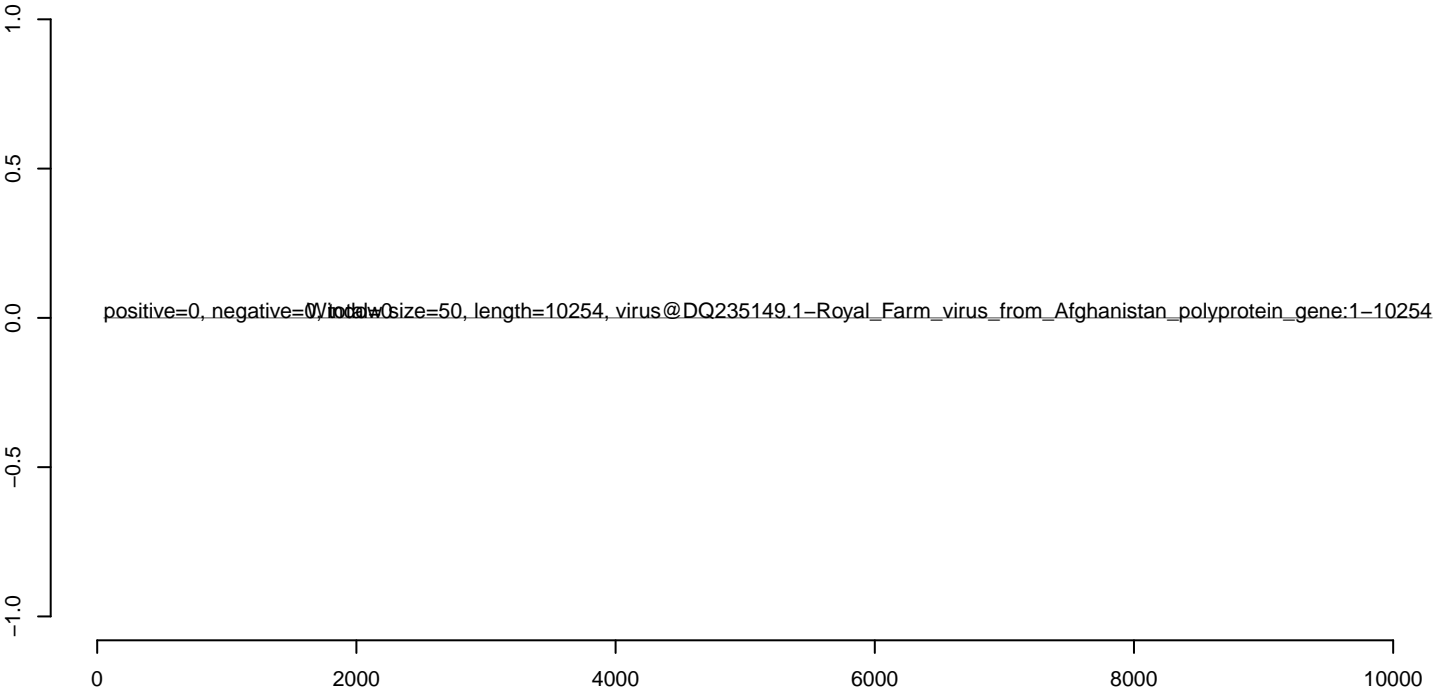
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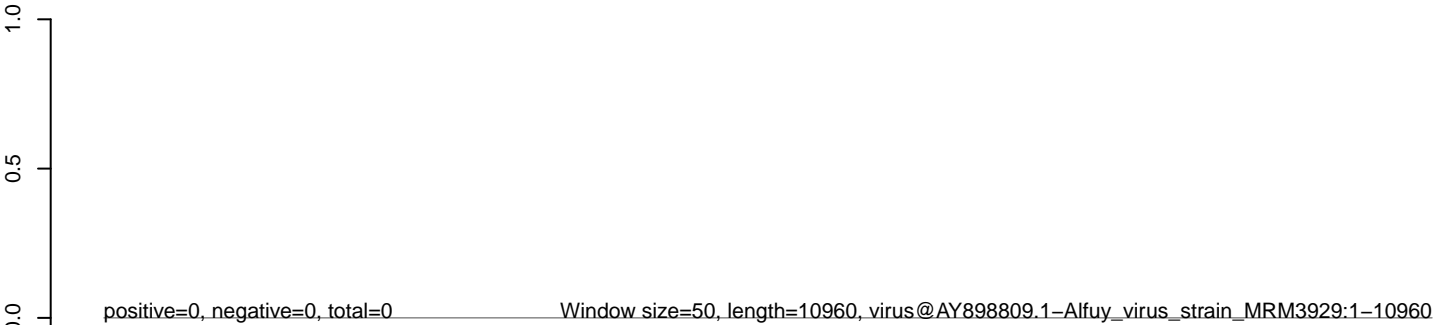
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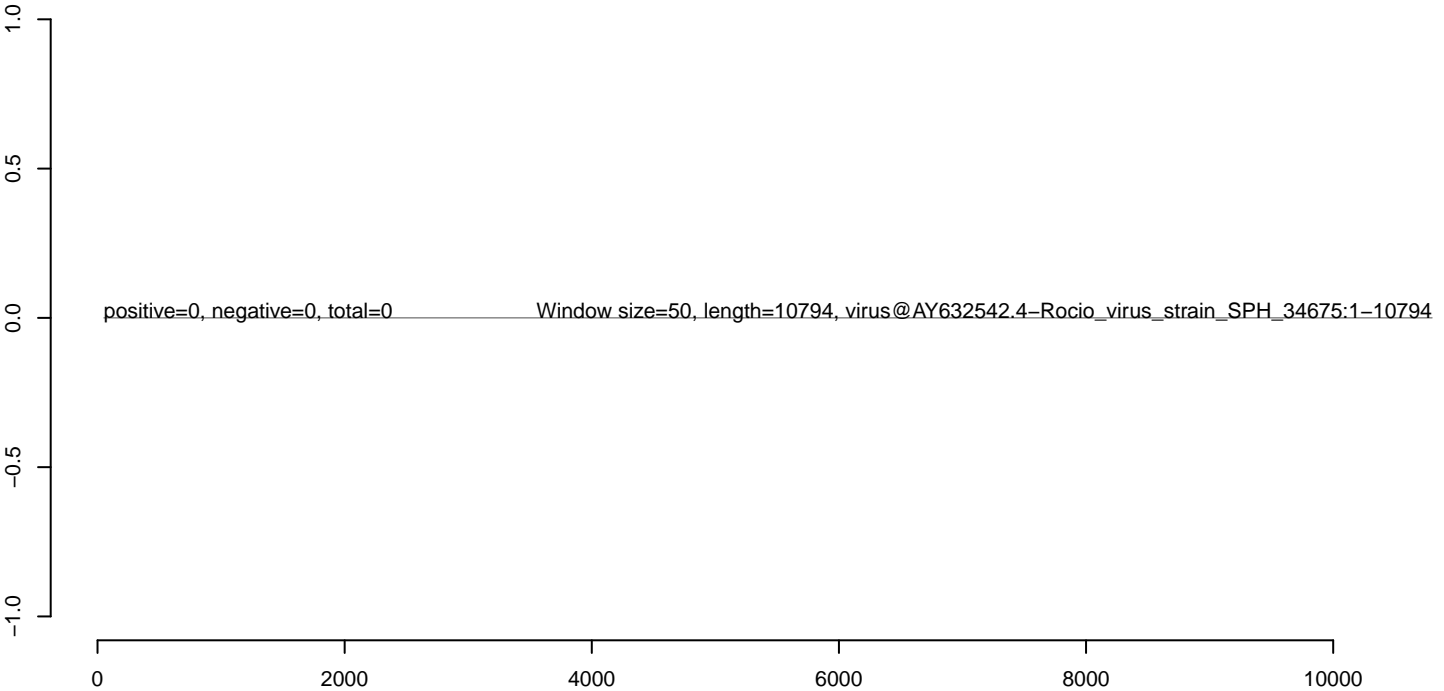
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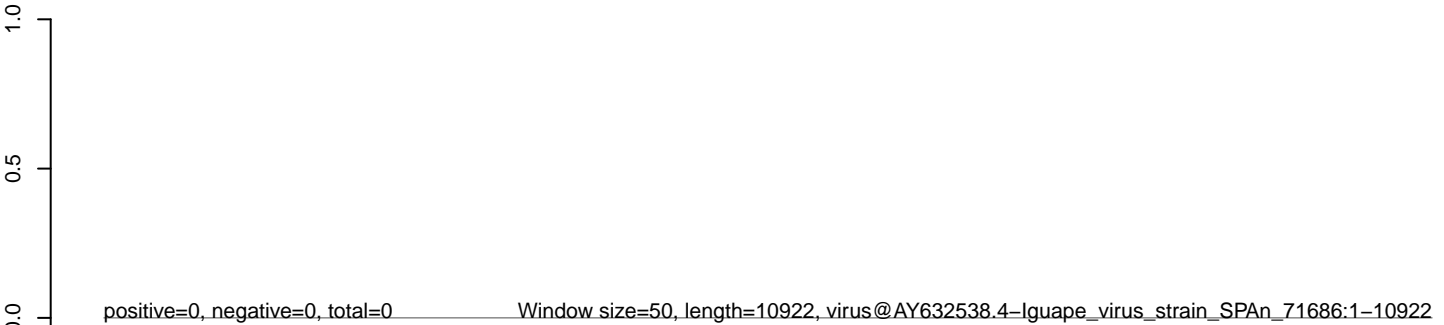
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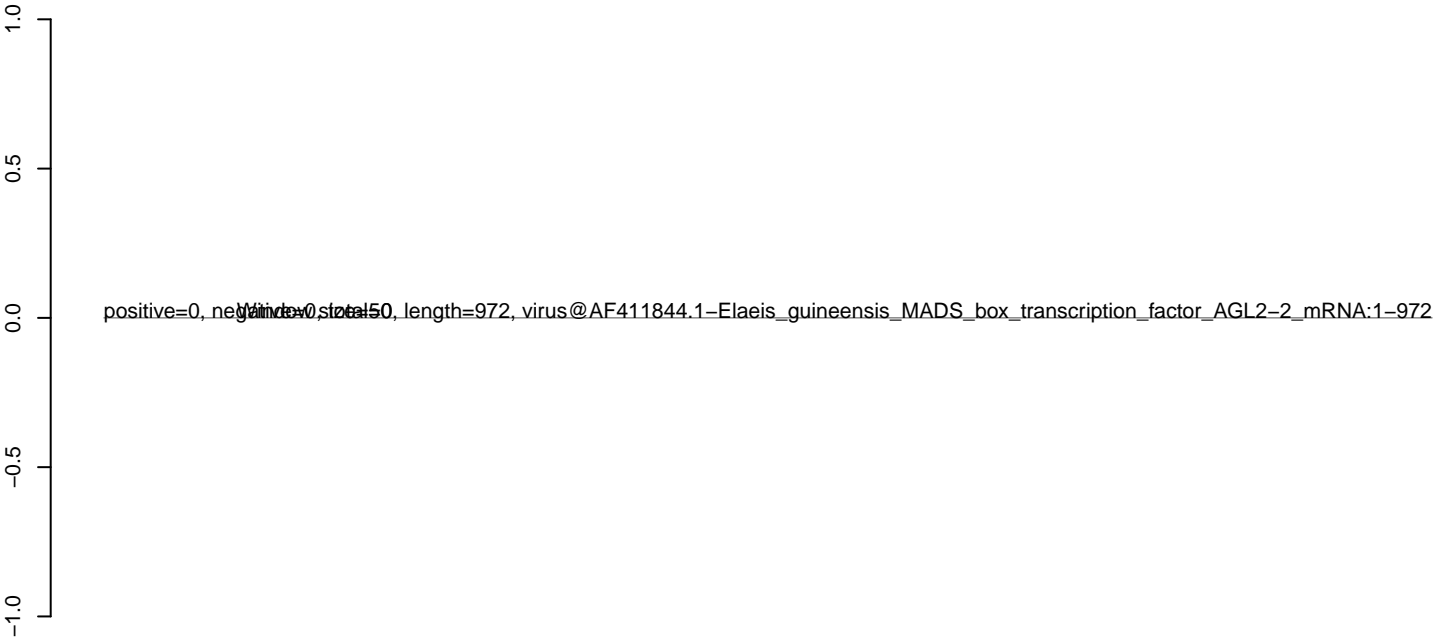
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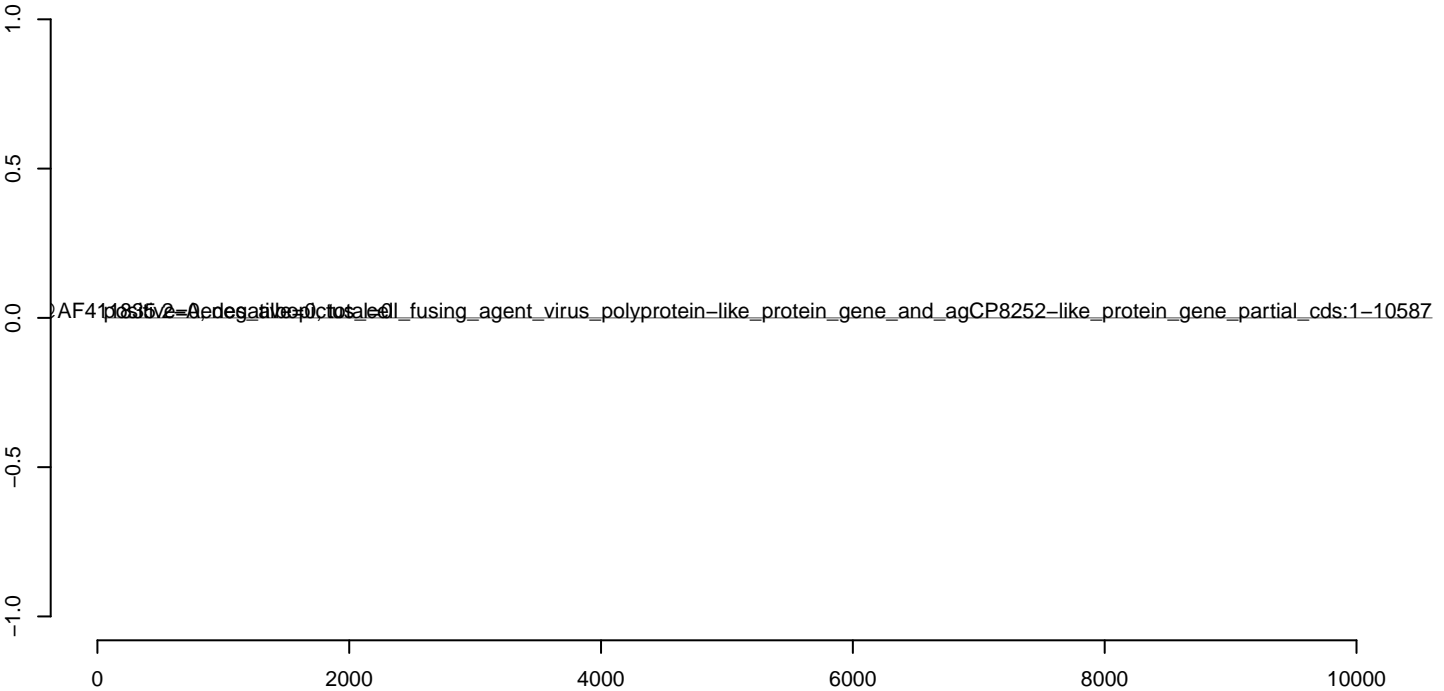
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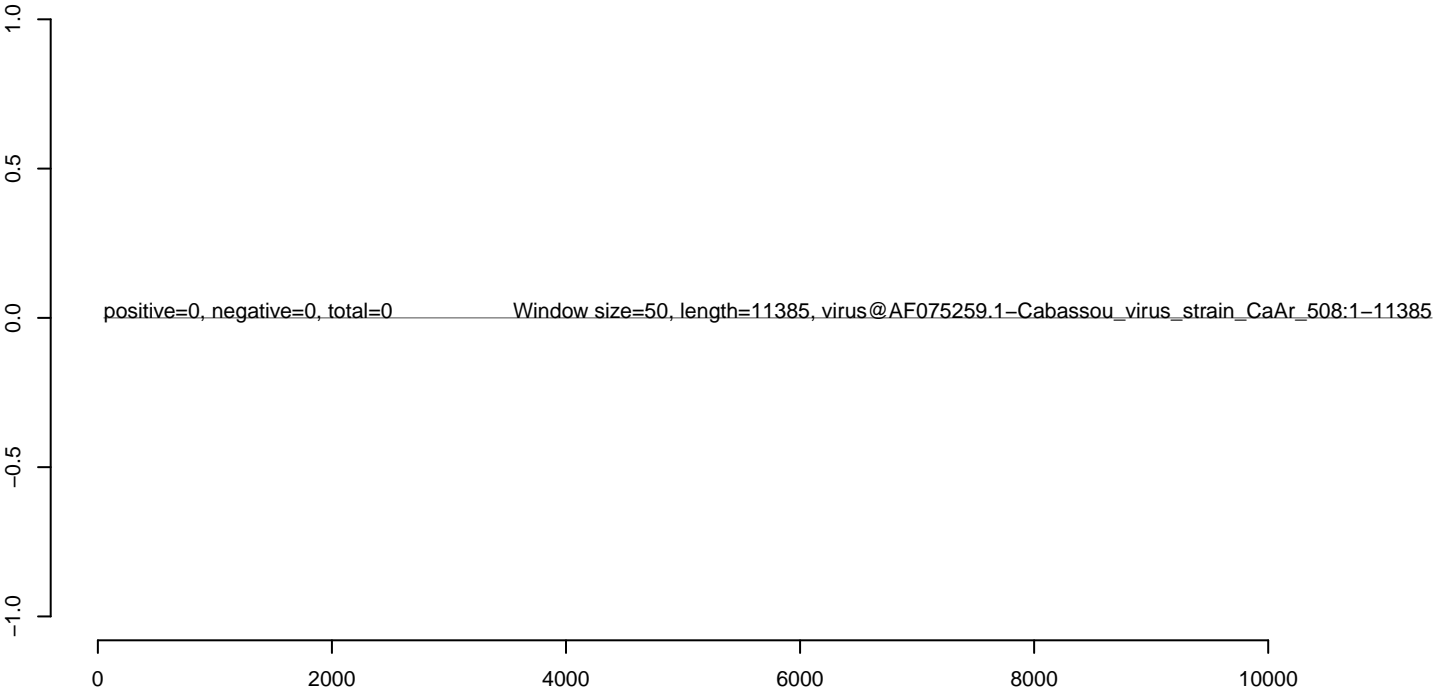
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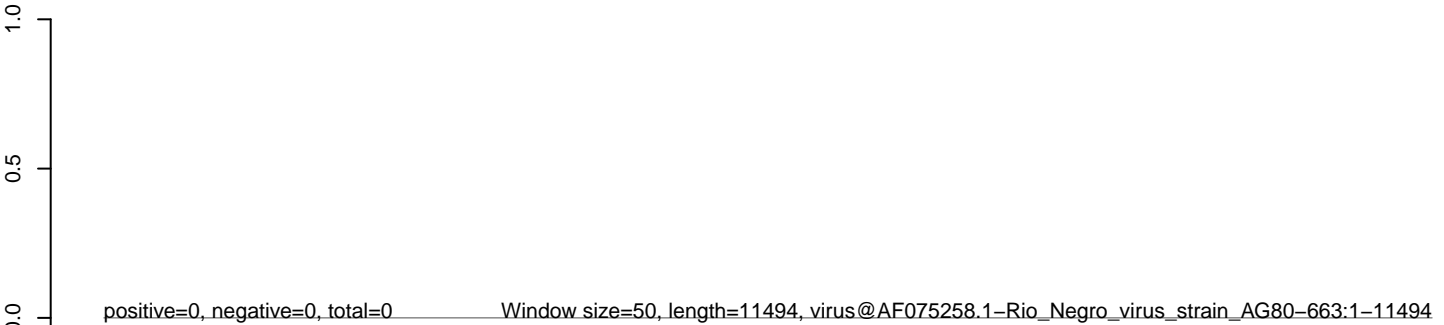
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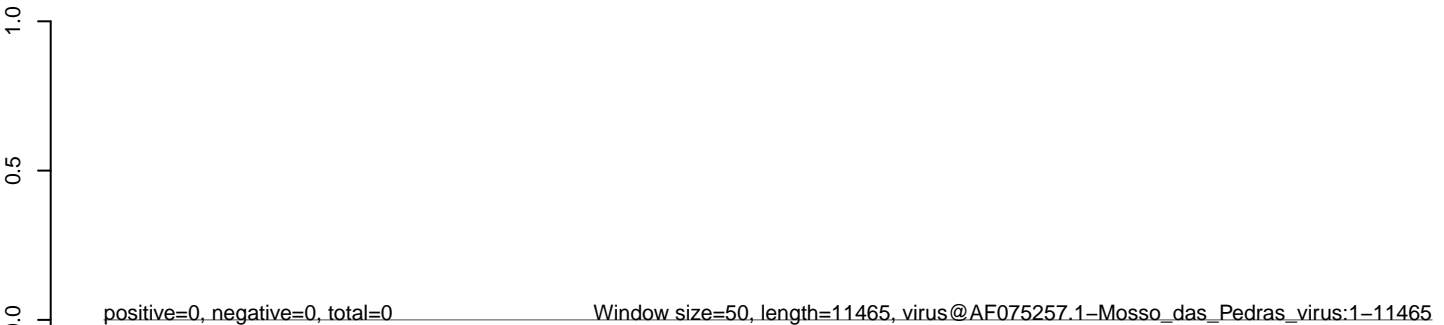
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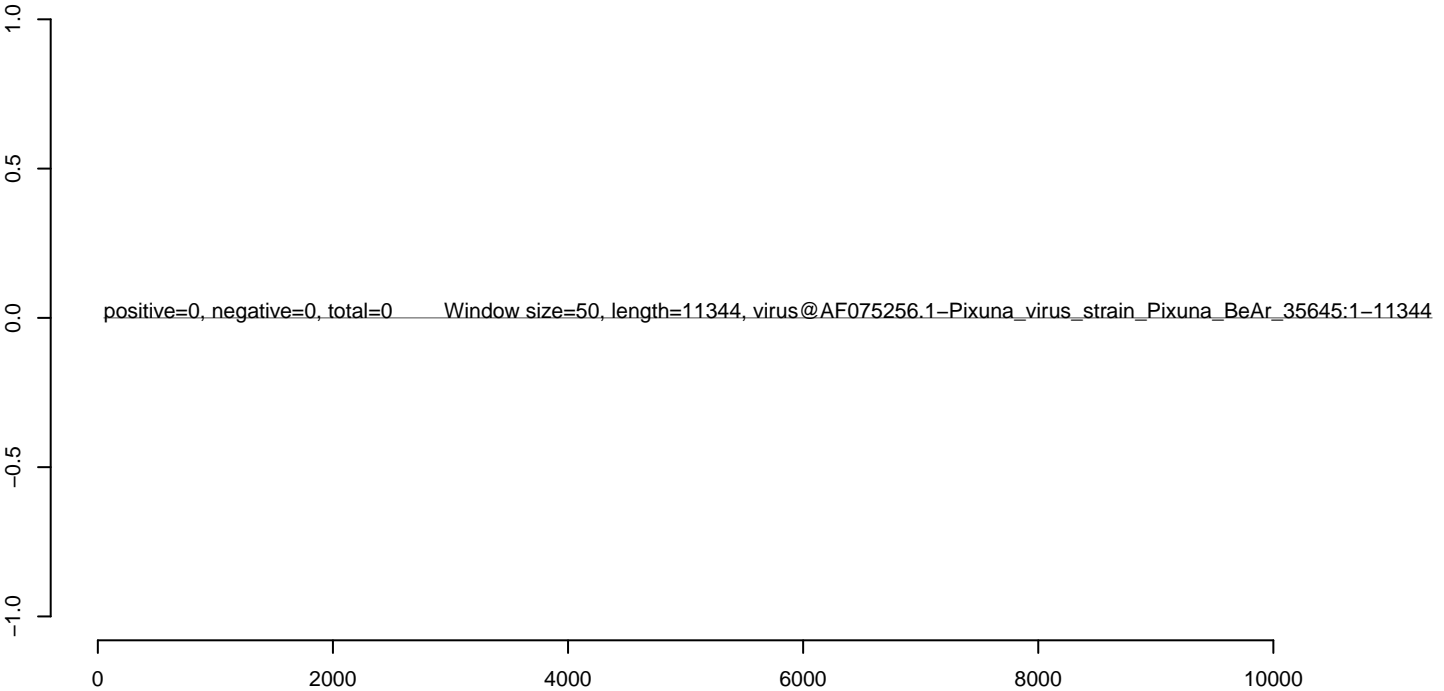
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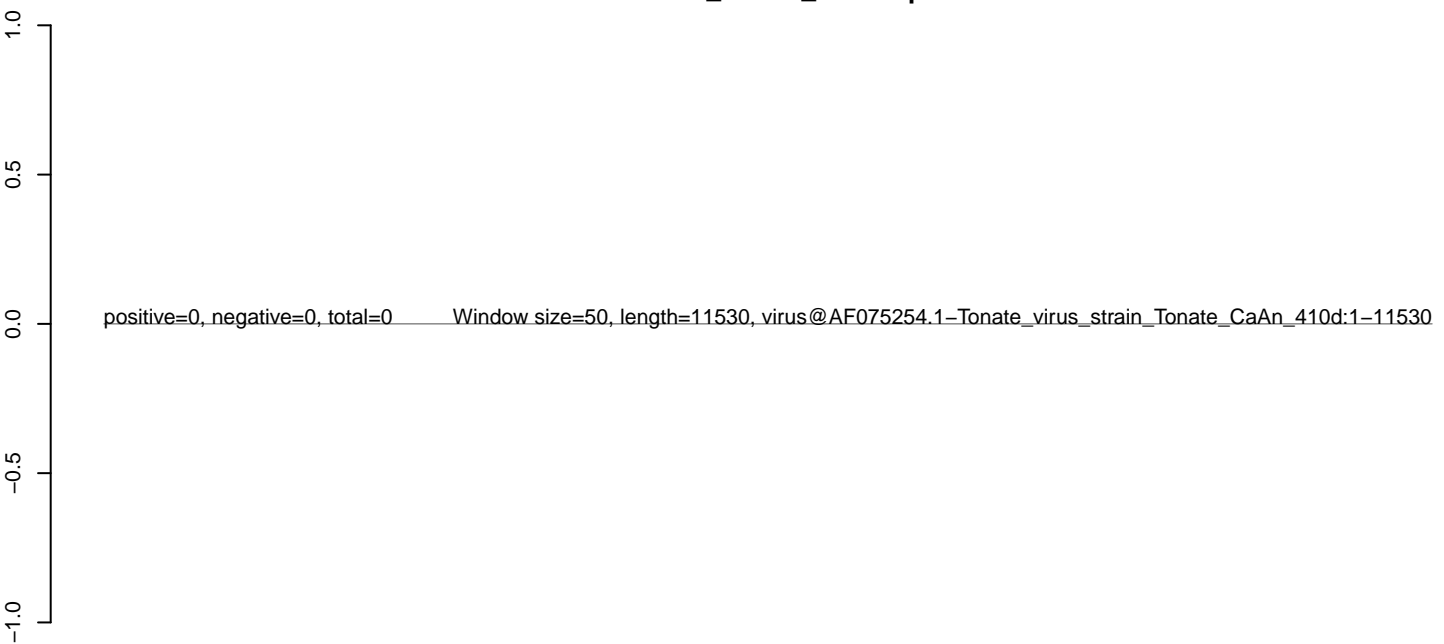
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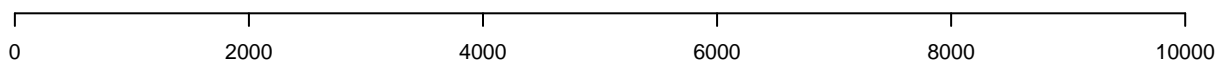
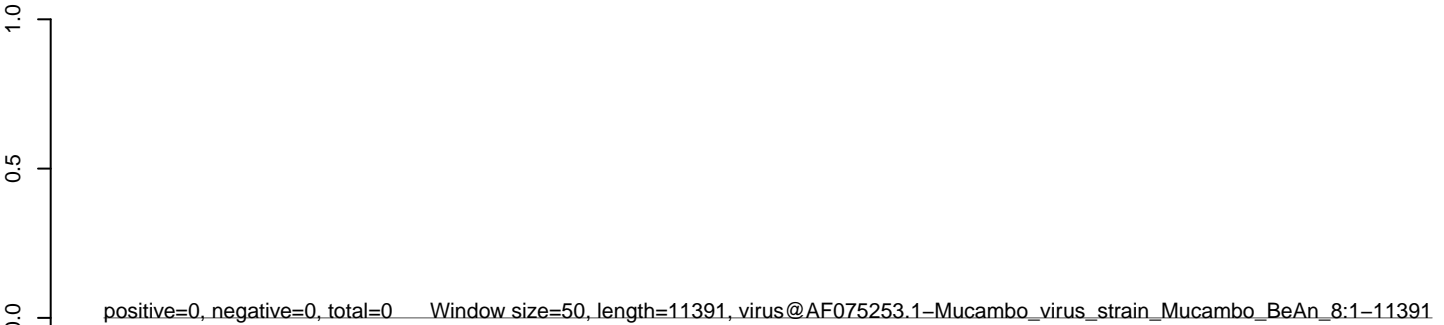
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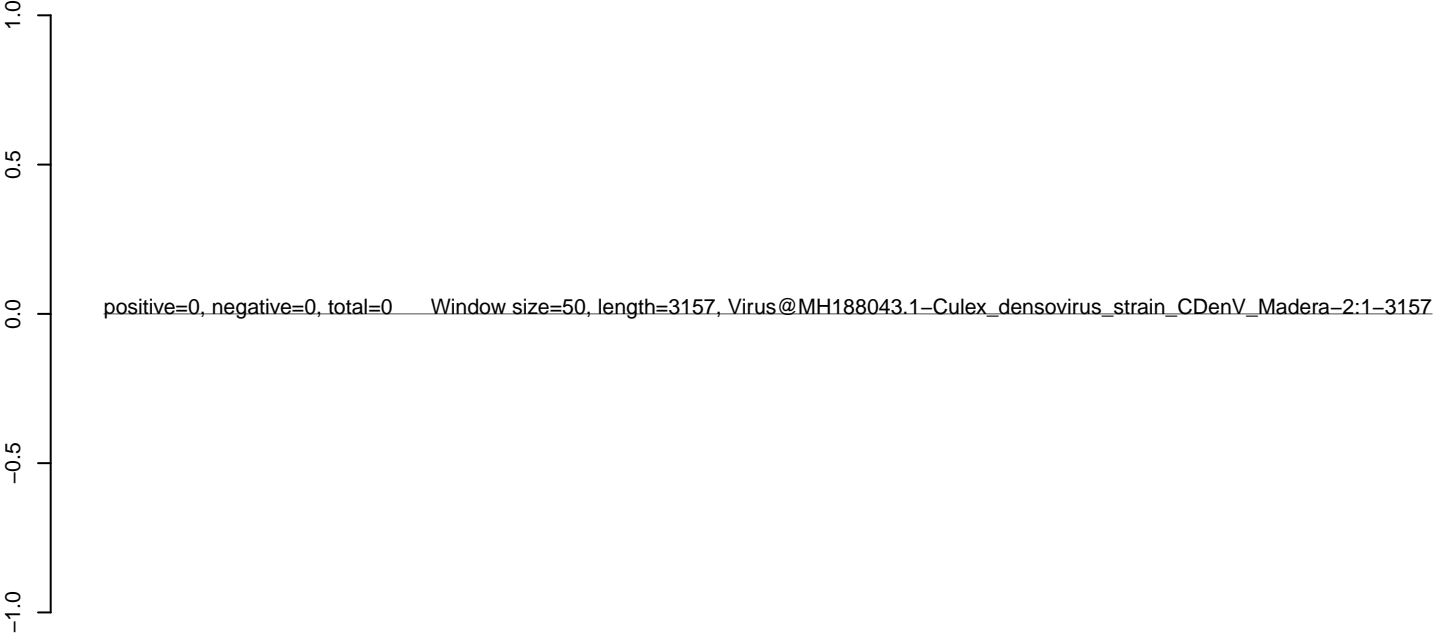
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**AnGam\_Sua5b\_cells.24\_35.rep**



**AnGam\_Sua5b\_cells.rep**



0 500 1000 1500 2000 2500 3000

AnGam\_Sua5b\_cells.18\_23.rep



AnGam\_Sua5b\_cells.24\_35.rep



AnGam\_Sua5b\_cells.rep

