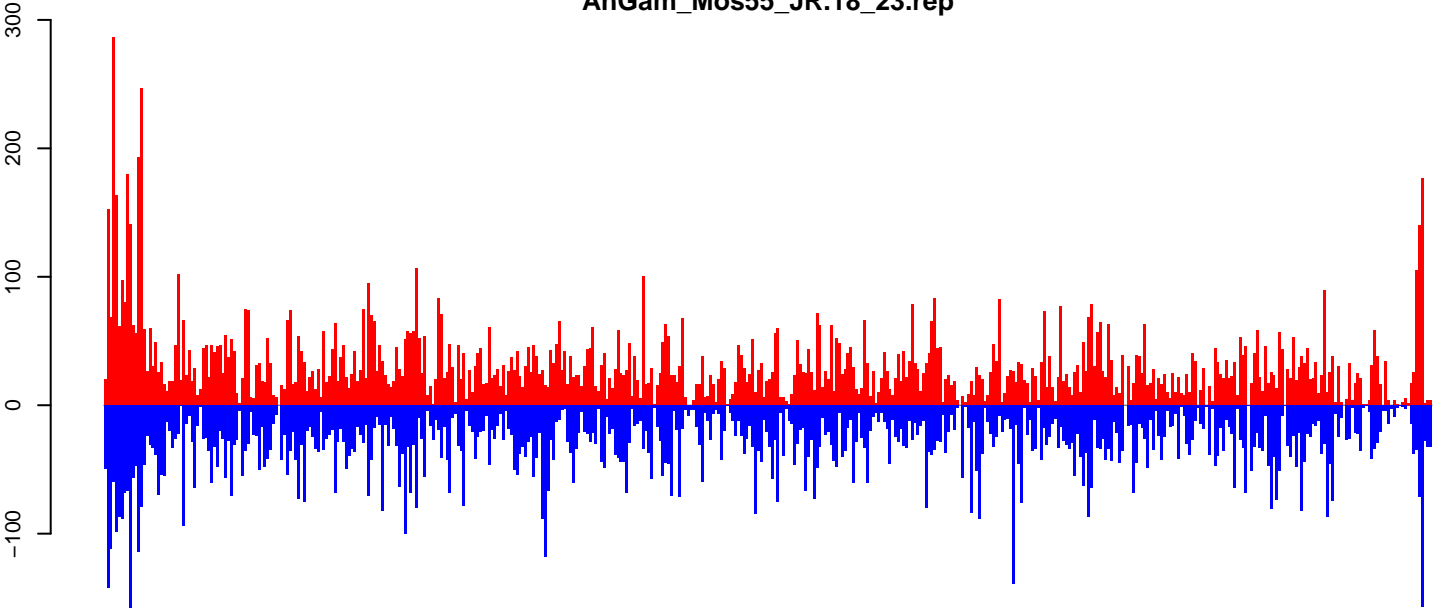


AnGam\_Mos55\_JR.18\_23.rep



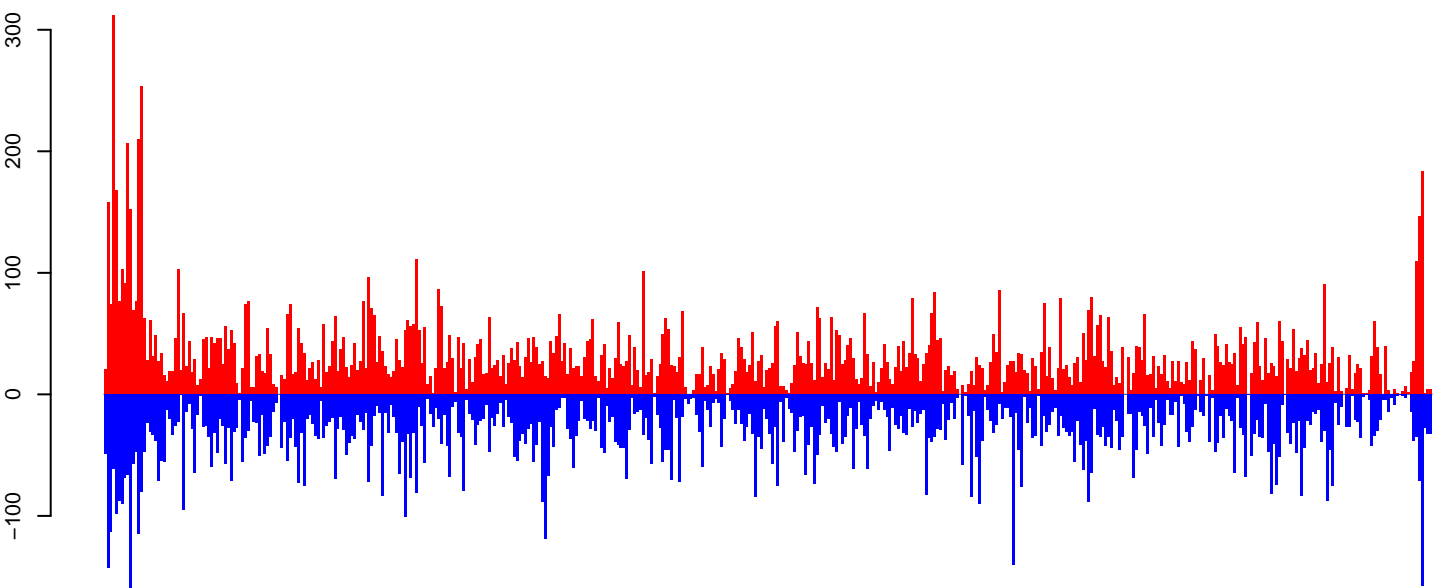
positive=14983, negative=14850, total=29833

AnGam\_Mos55\_JR.24\_35.rep



positive=459, negative=147, total=606

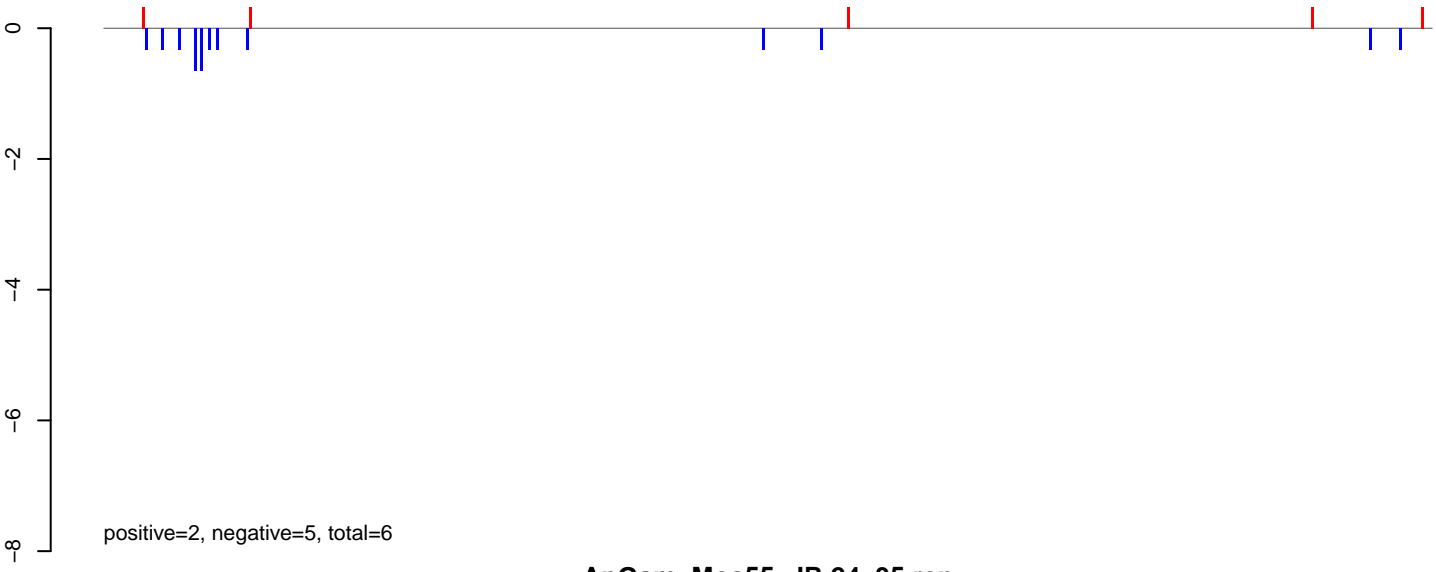
AnGam\_Mos55\_JR.rep



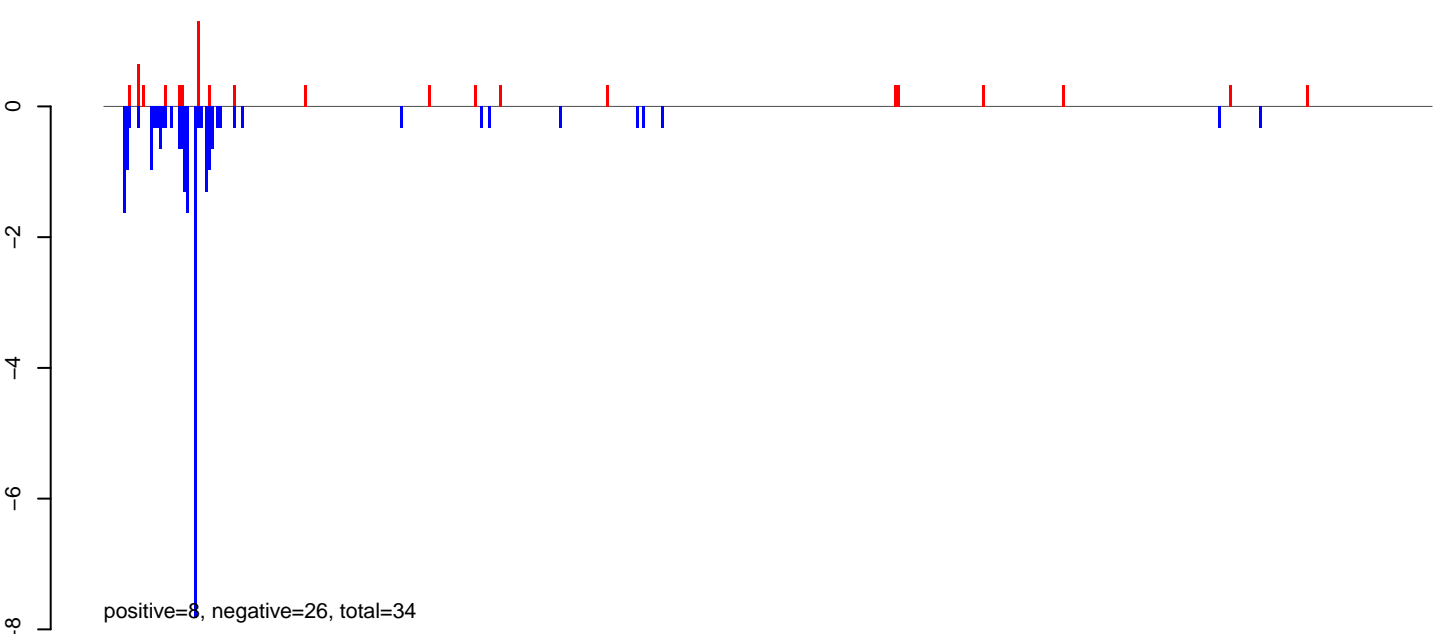
positive=15442, negative=14997, total=30439, virus@M20303.1-O-Nyong-nyong\_virus\_Gulu\_strain\_complete\_genome:1-11835

0 2000 4000 6000 8000 10000 12000

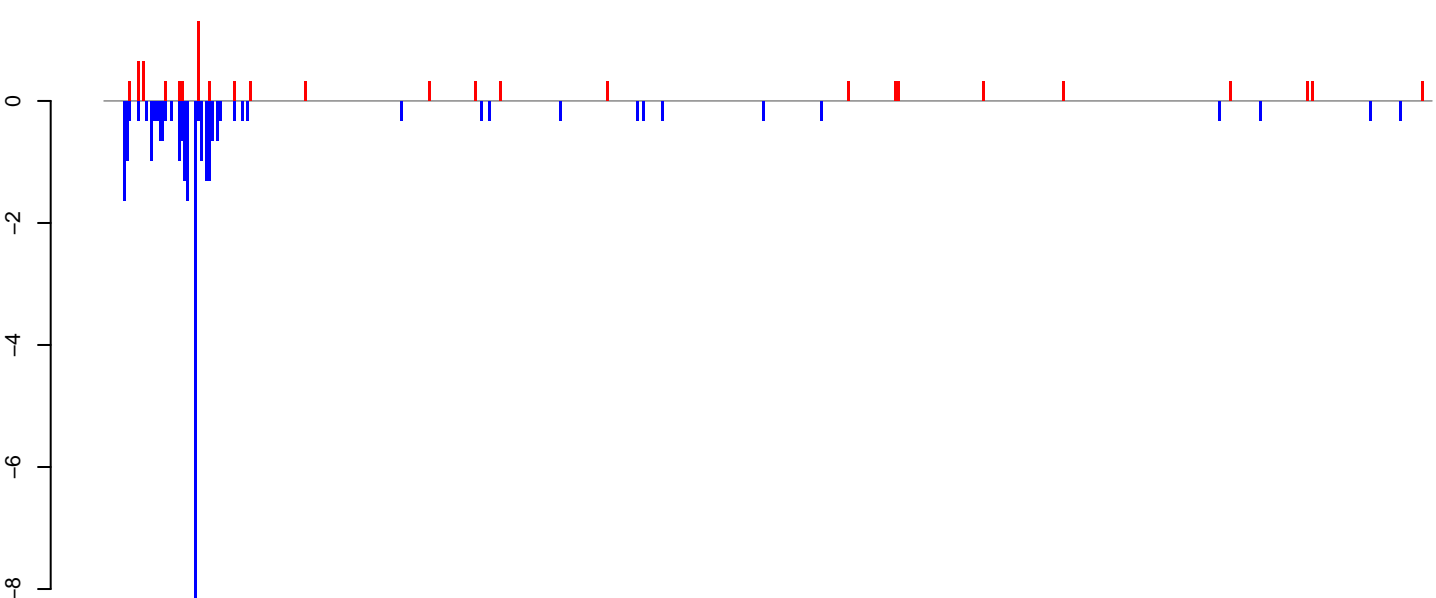
AnGam\_Mos55\_JR.18\_23.rep



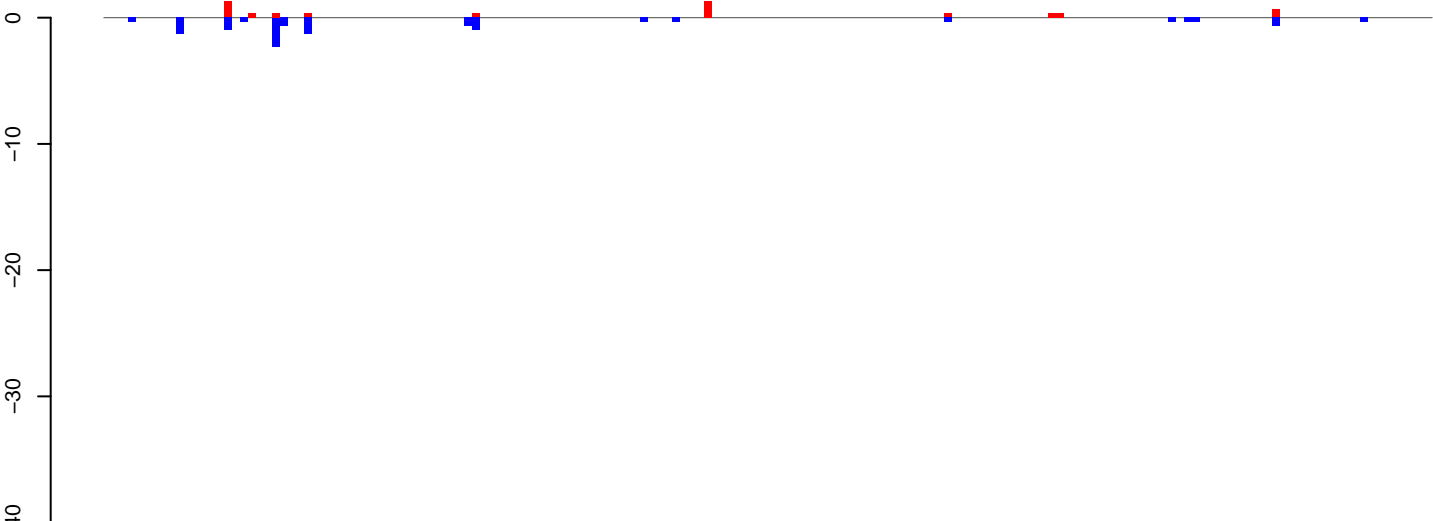
AnGam\_Mos55\_JR.24\_35.rep



AnGam\_Mos55\_JR.rep

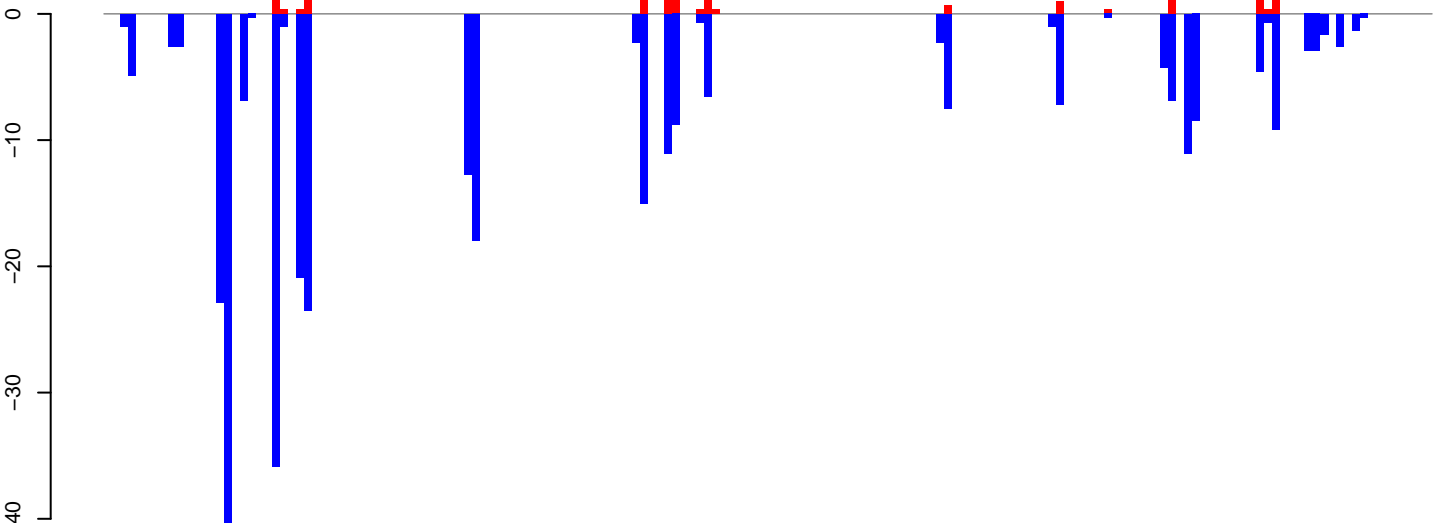


AnGam\_Mos55\_JR.18\_23.rep



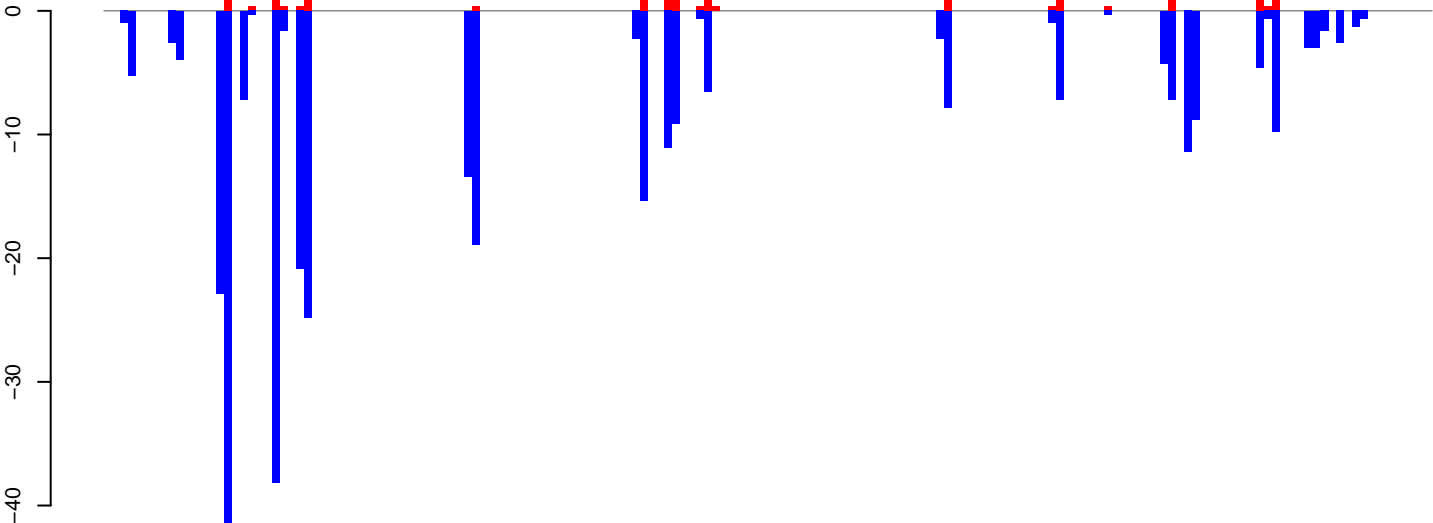
positive=6, negative=12, total=17

AnGam\_Mos55\_JR.24\_35.rep



positive=25, negative=315, total=340

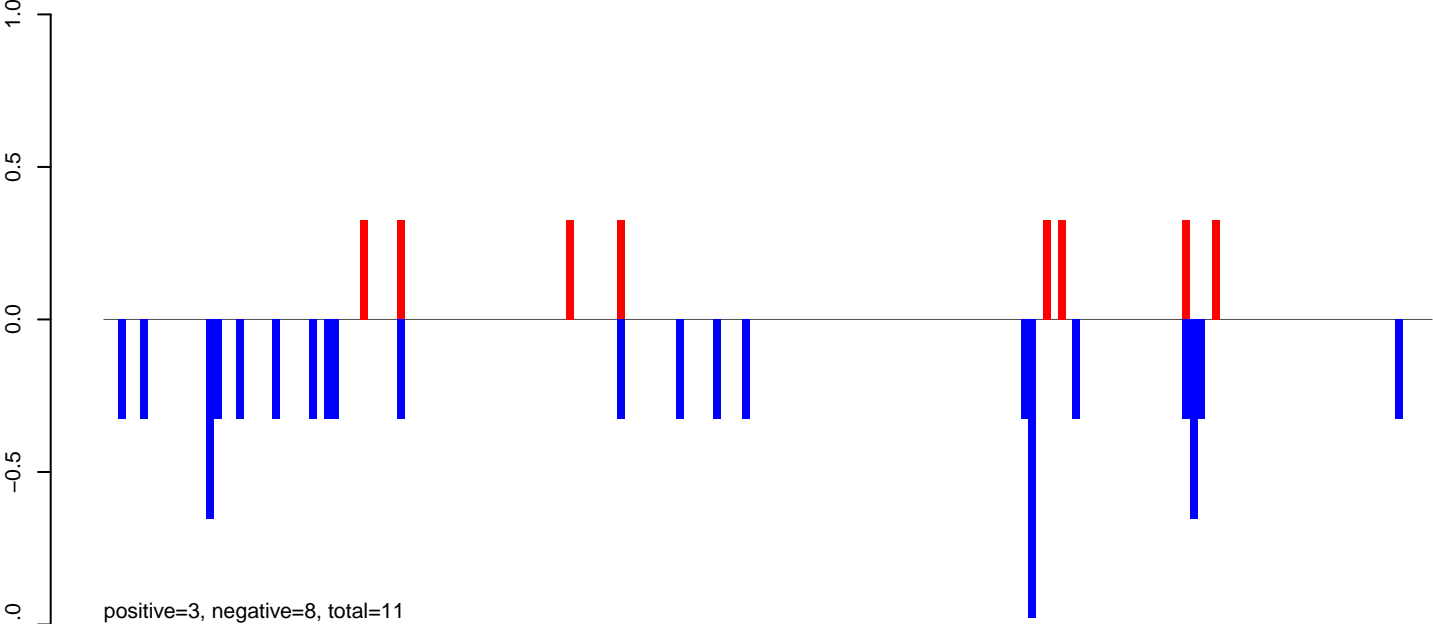
AnGam\_Mos55\_JR.rep



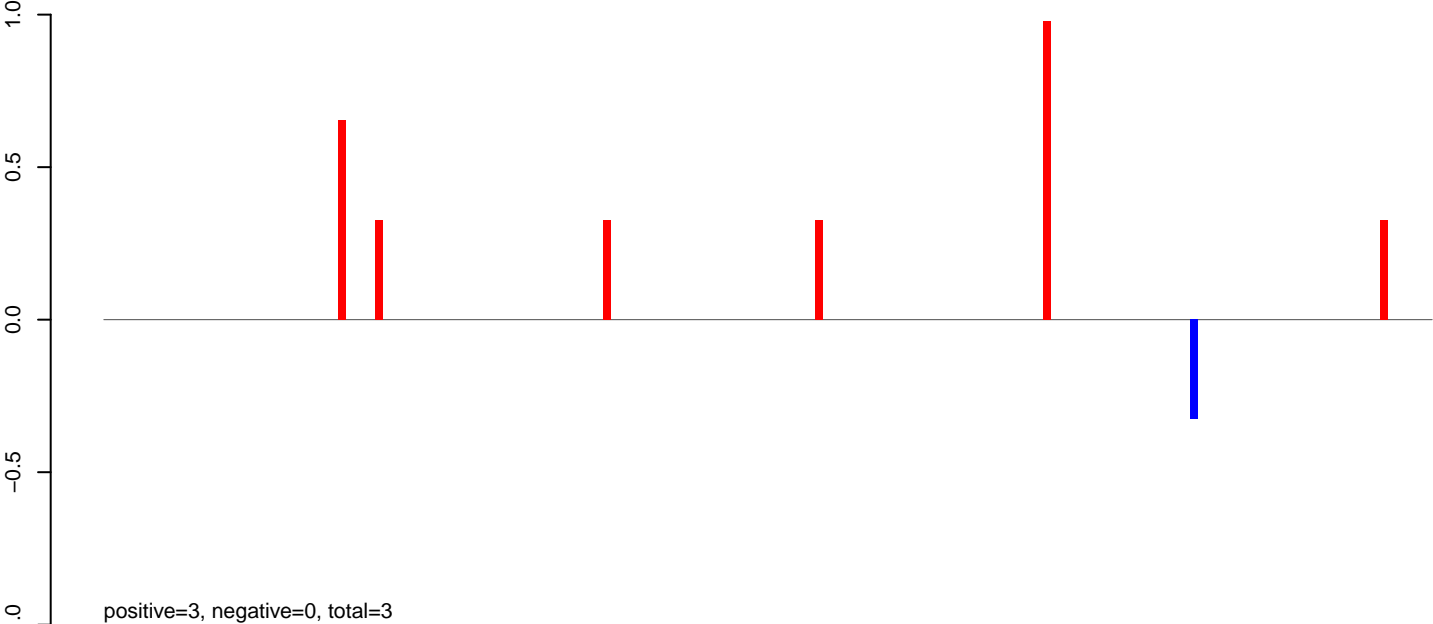
positive=30, negative=327, total=357, window size=25, length=4139, virus@NC\_011317.1-Anopheles\_gambiae\_densovirus:1-4139

0 1000 2000 3000 4000

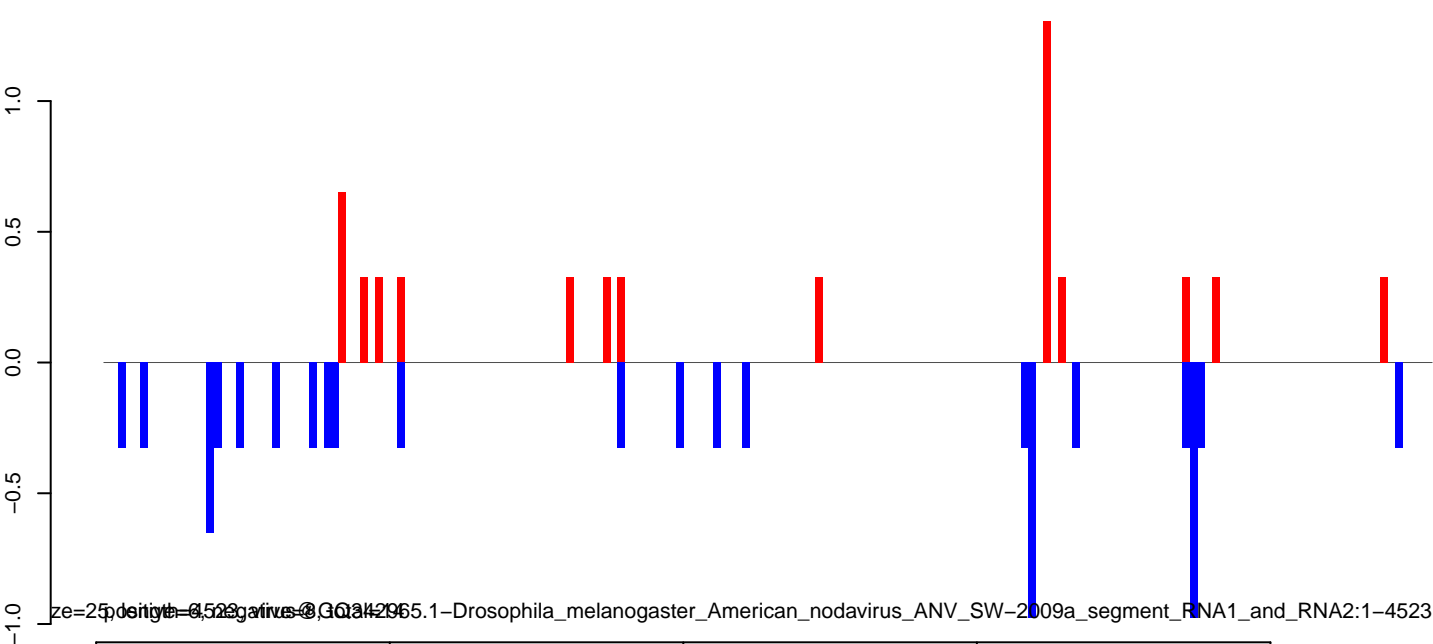
AnGam\_Mos55\_JR.18\_23.rep



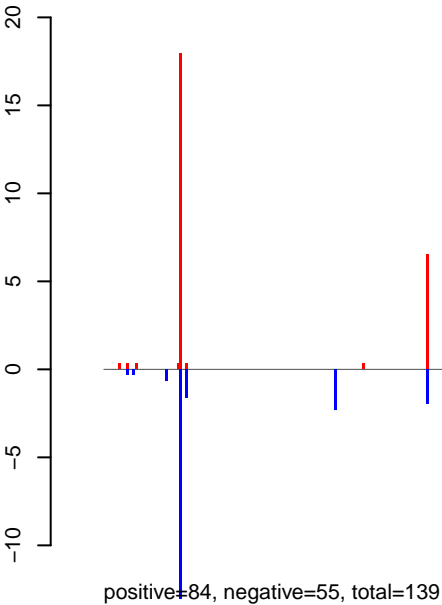
AnGam\_Mos55\_JR.24\_35.rep



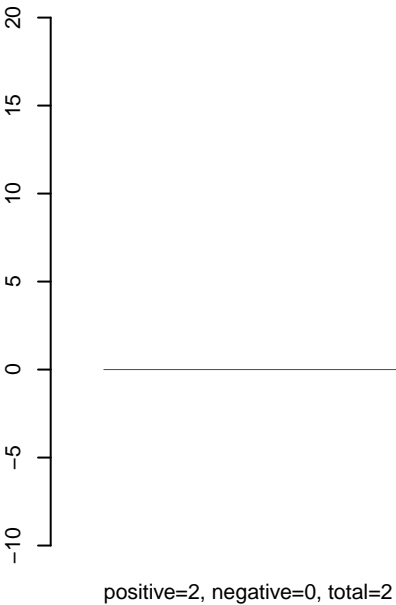
AnGam\_Mos55\_JR.rep



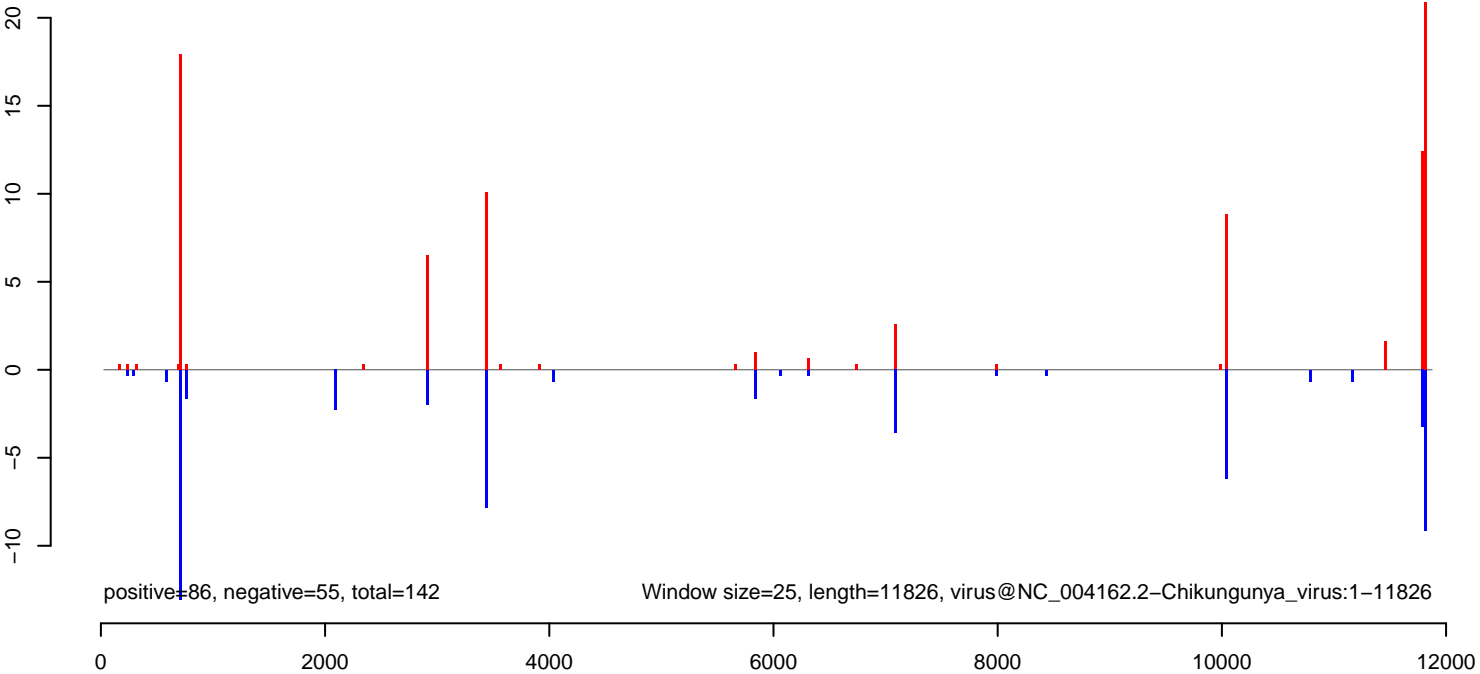
AnGam\_Mos55\_JR.18\_23.rep



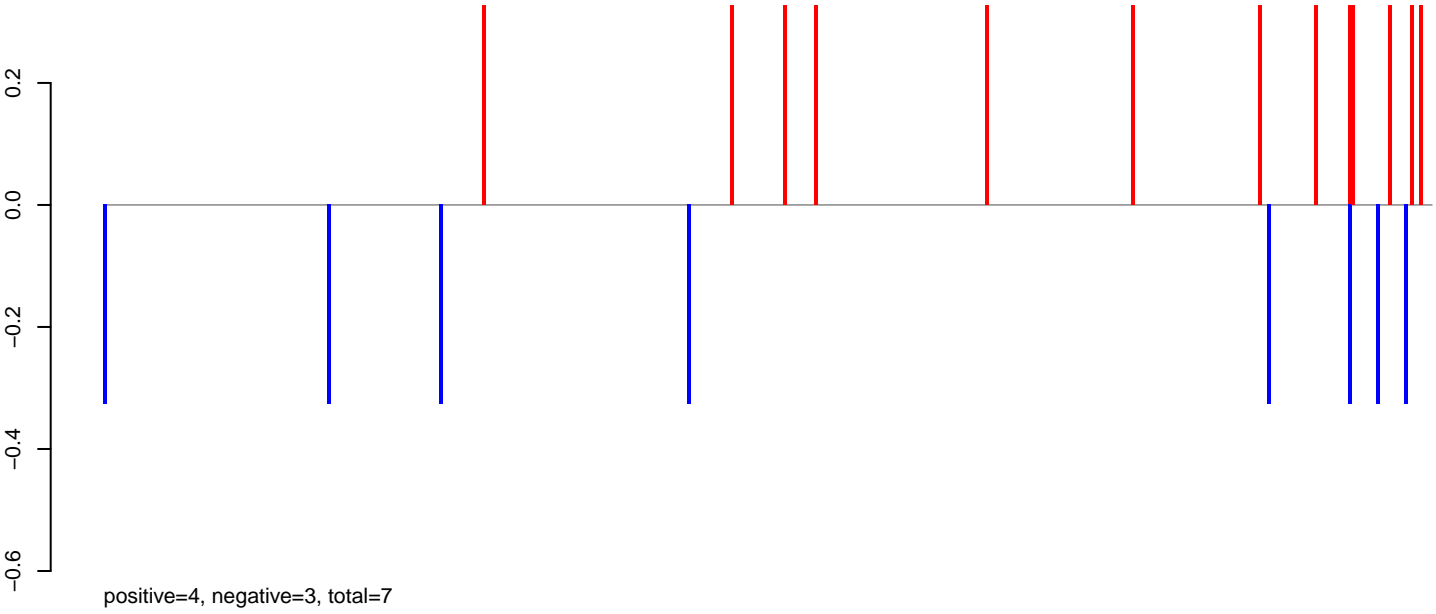
AnGam\_Mos55\_JR.24\_35.rep



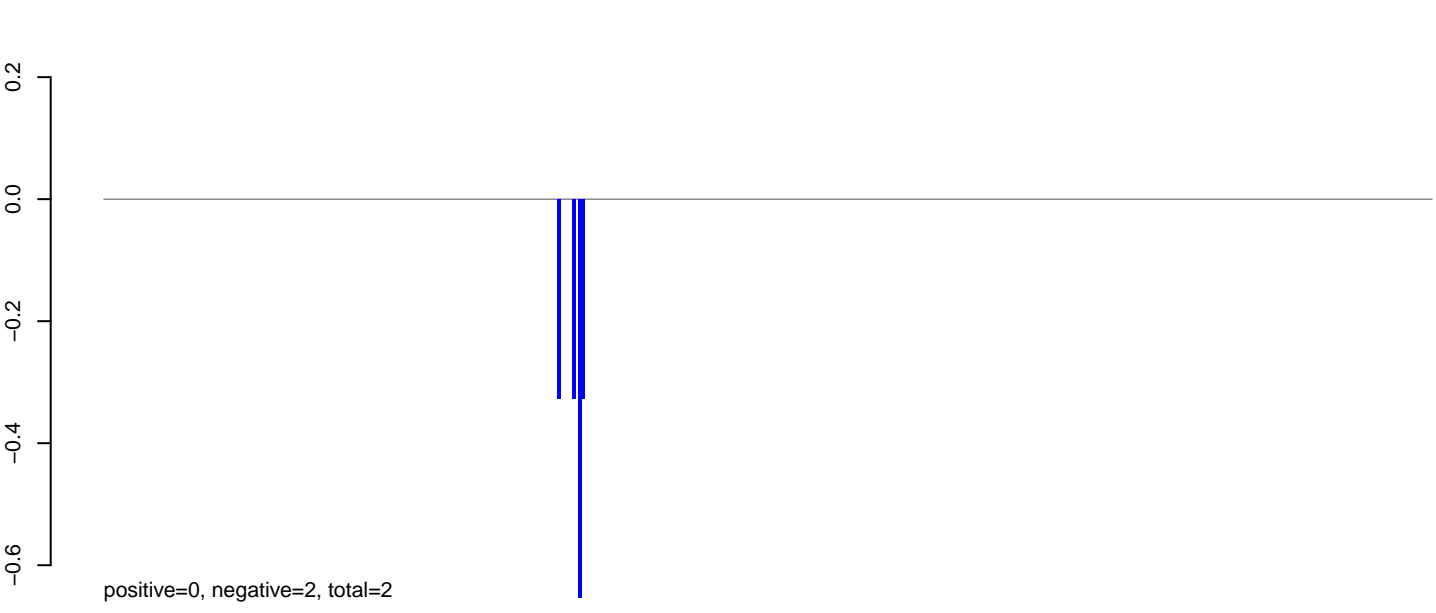
AnGam\_Mos55\_JR.rep



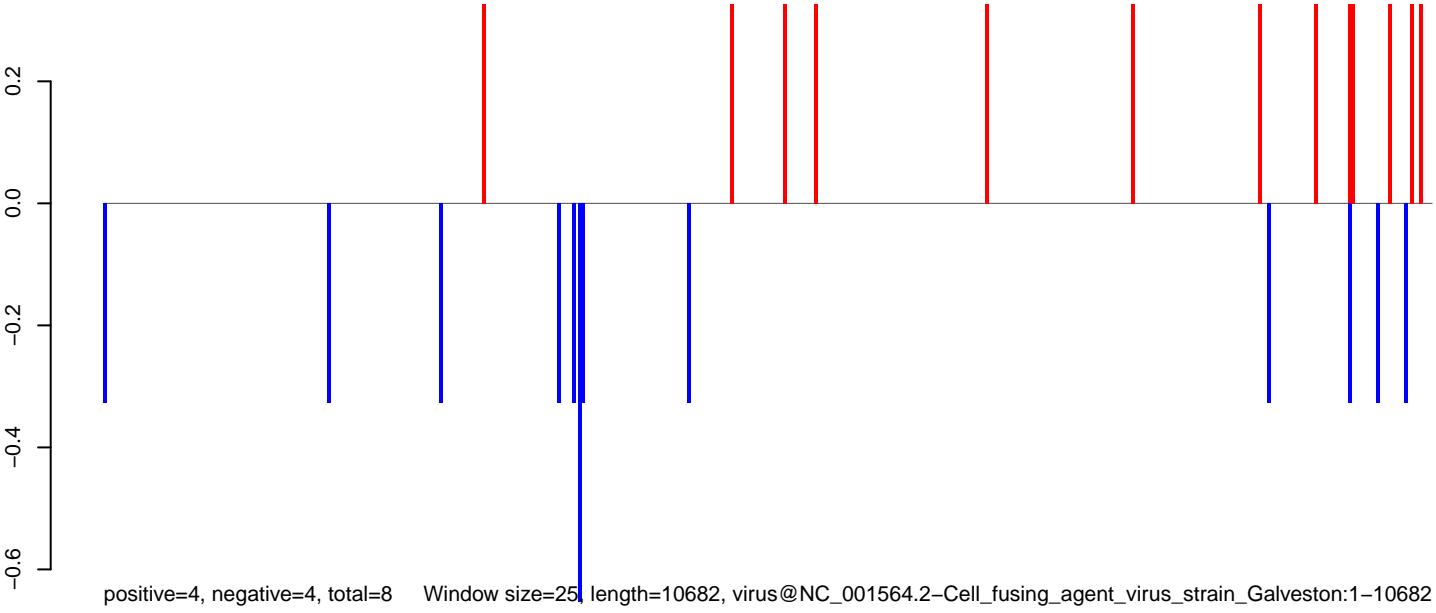
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



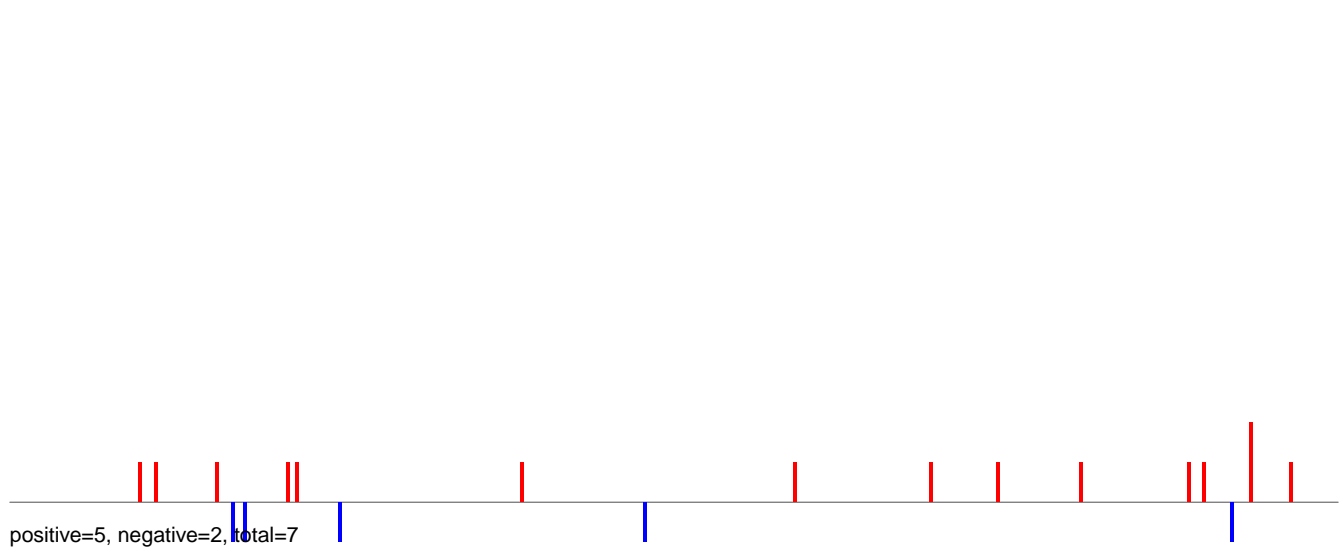
AnGam\_Mos55\_JR.rep



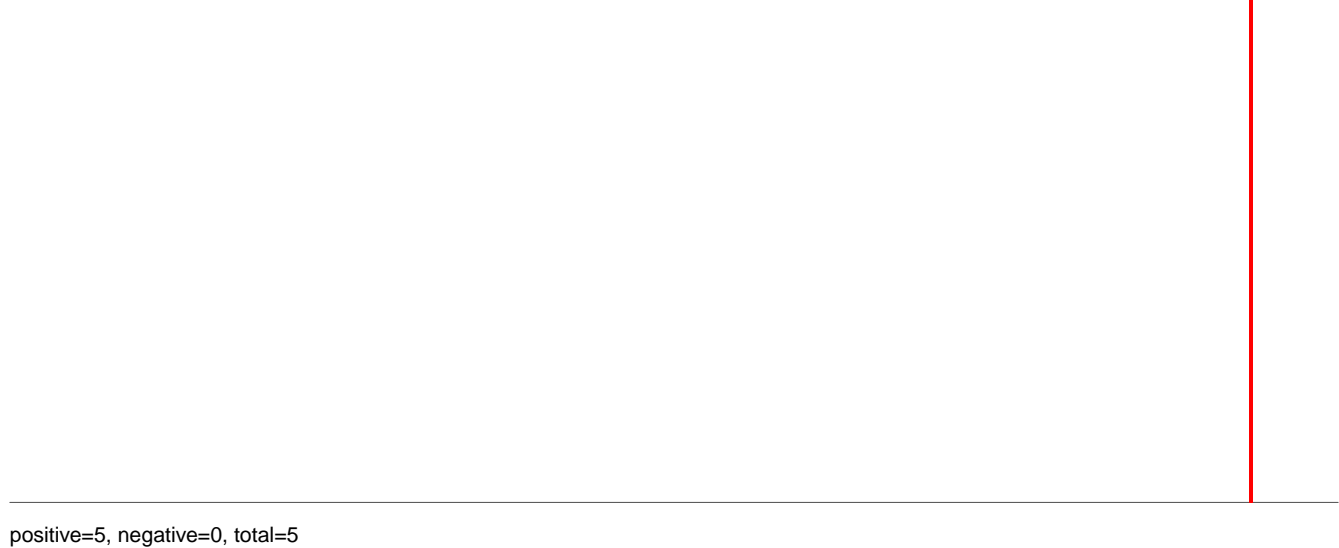
Window size=25, length=10682, virus@NC\_001564.2-Cell\_fusing\_agent\_virus\_strain\_Galveston:1-10682

0 2000 4000 6000 8000 10000

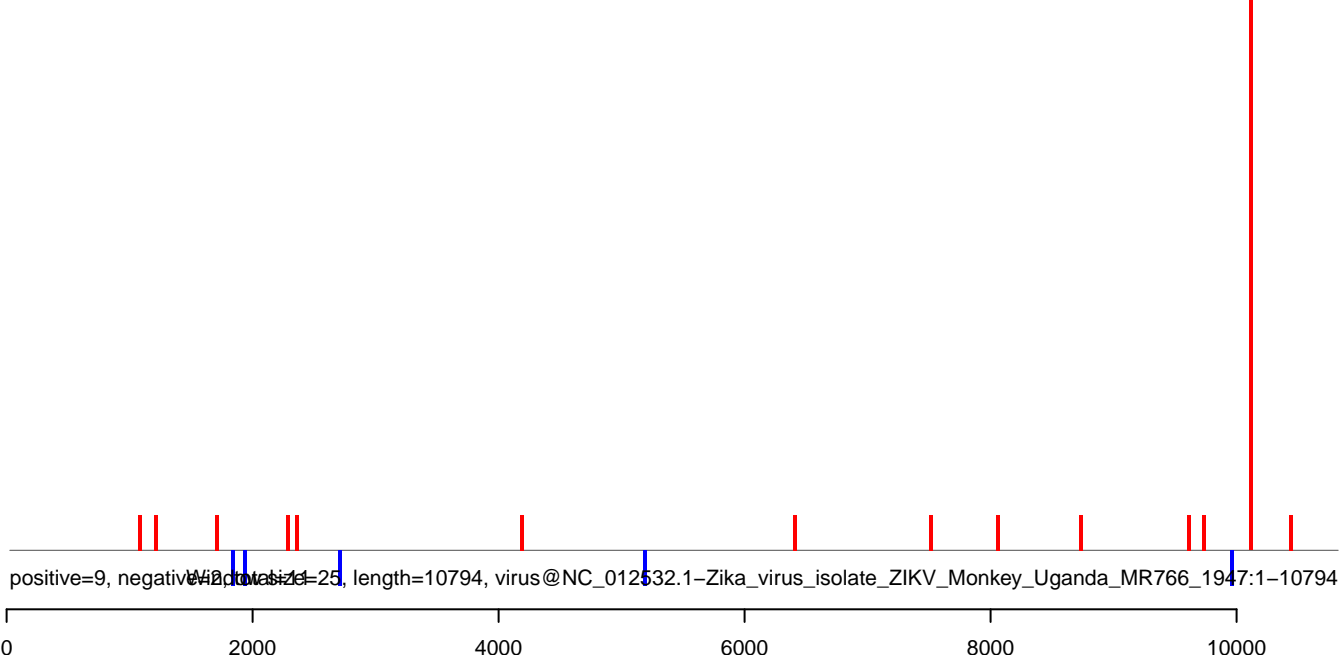
AnGam\_Mos55\_JR.18\_23.rep



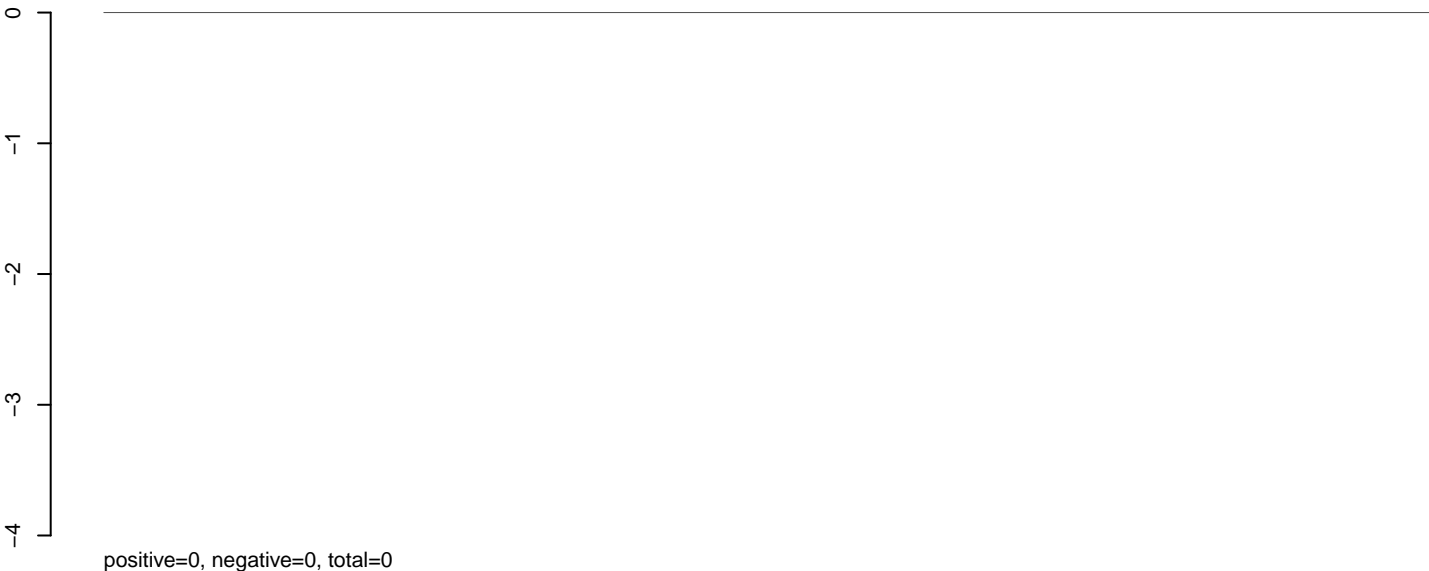
AnGam\_Mos55\_JR.24\_35.rep



AnGam\_Mos55\_JR.rep



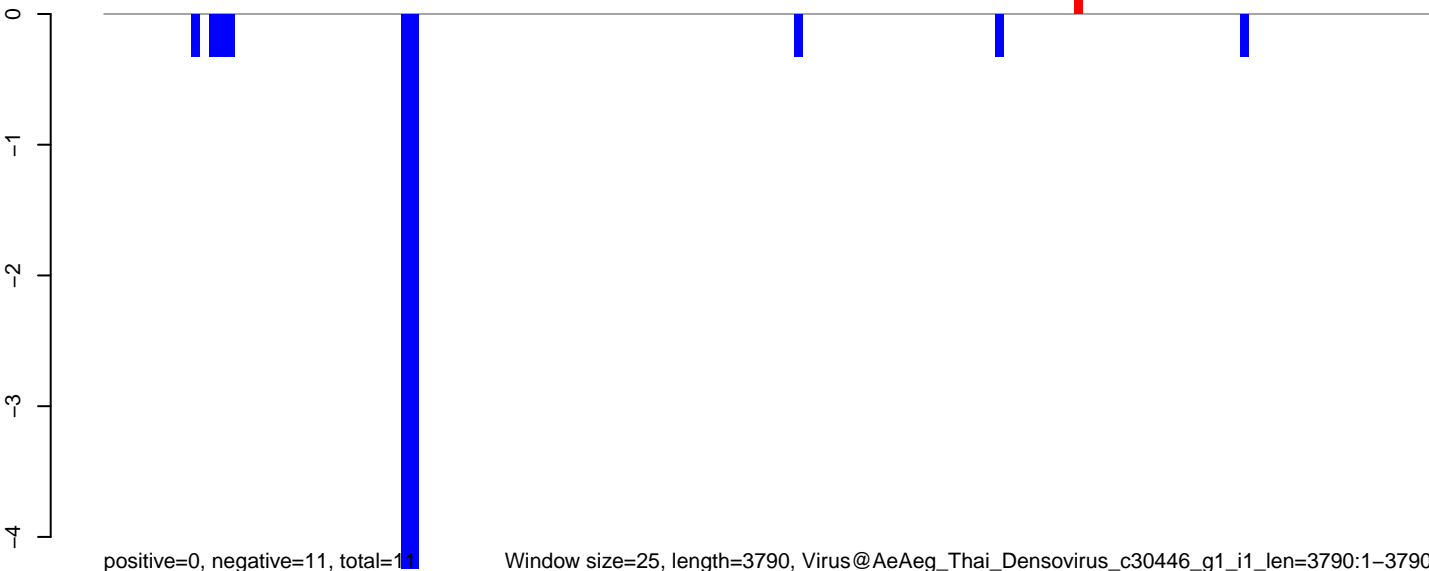
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



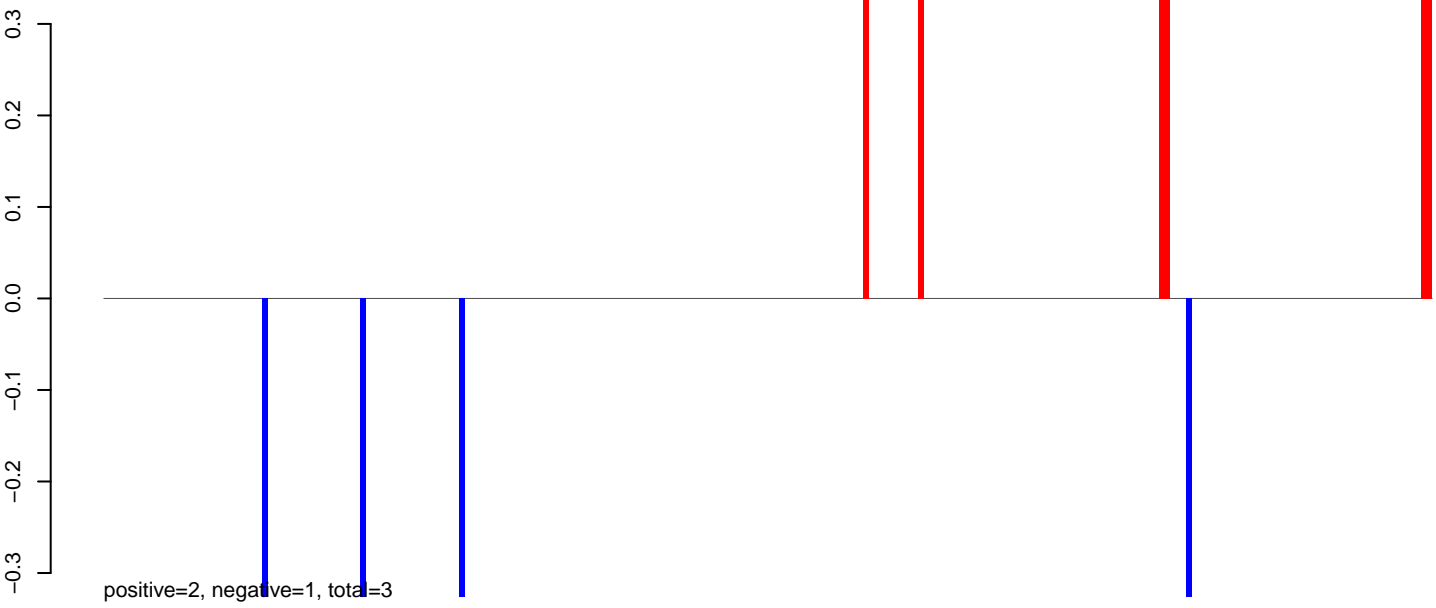
AnGam\_Mos55\_JR.rep



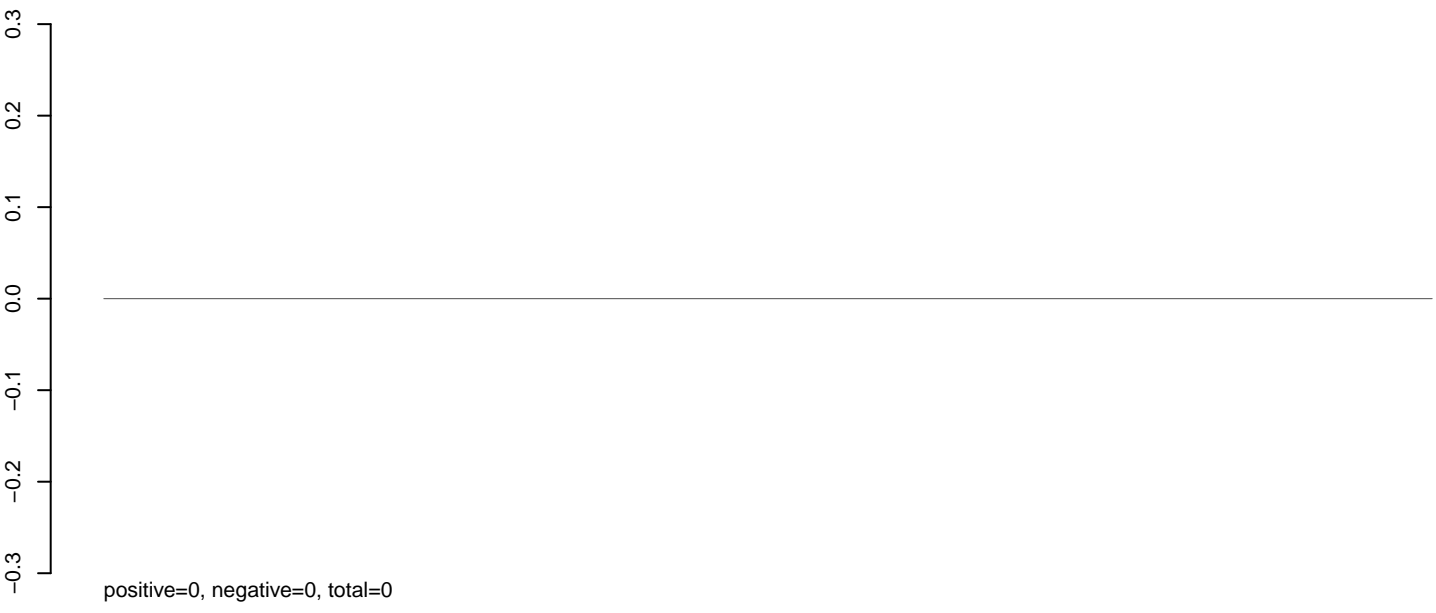
0 1000 2000 3000



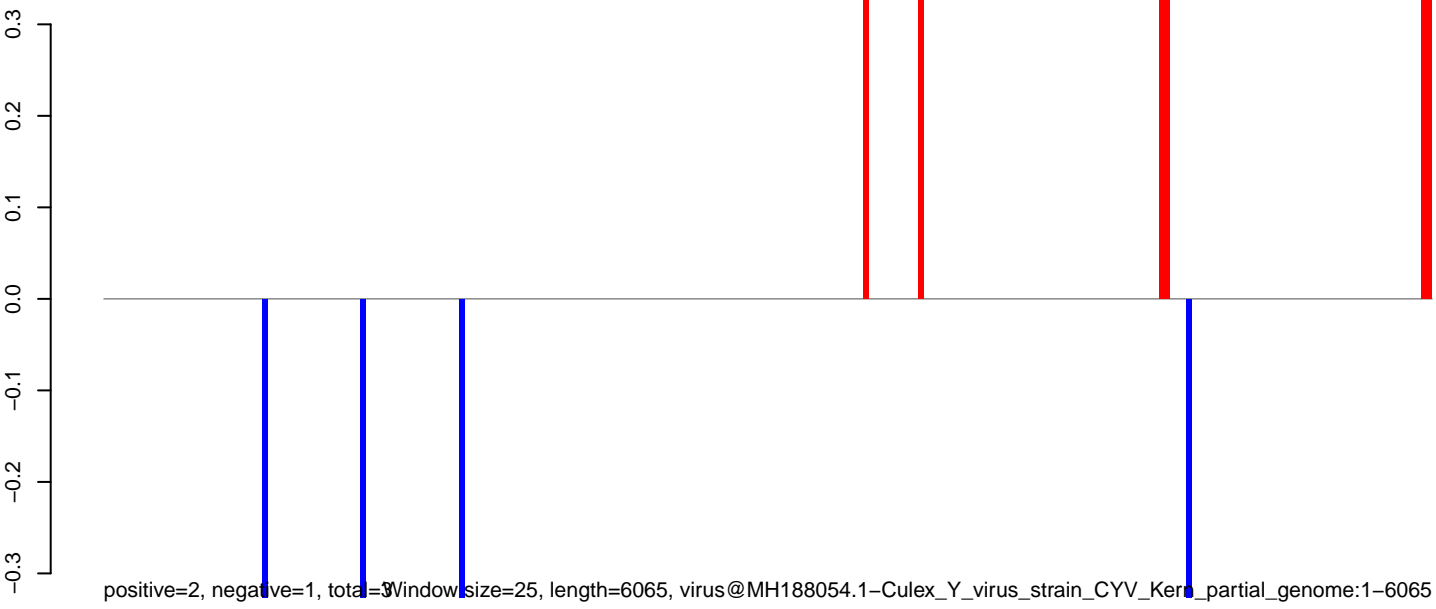
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep

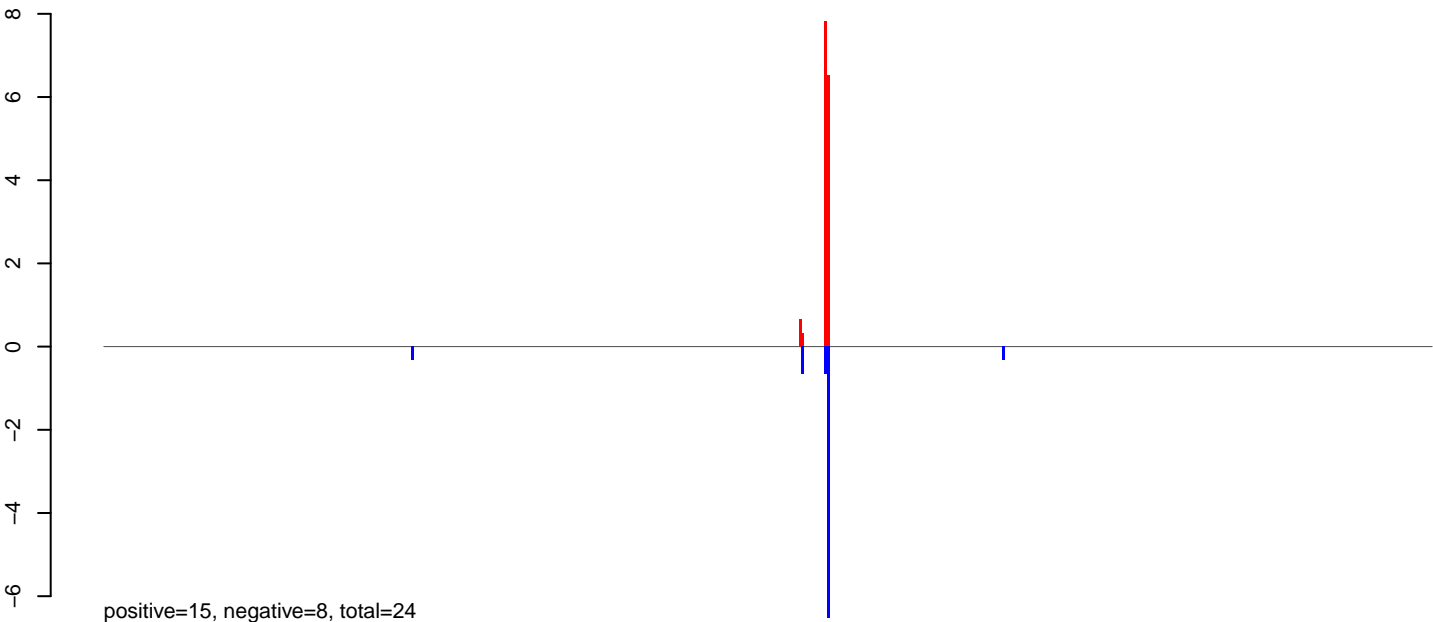


AnGam\_Mos55\_JR.rep

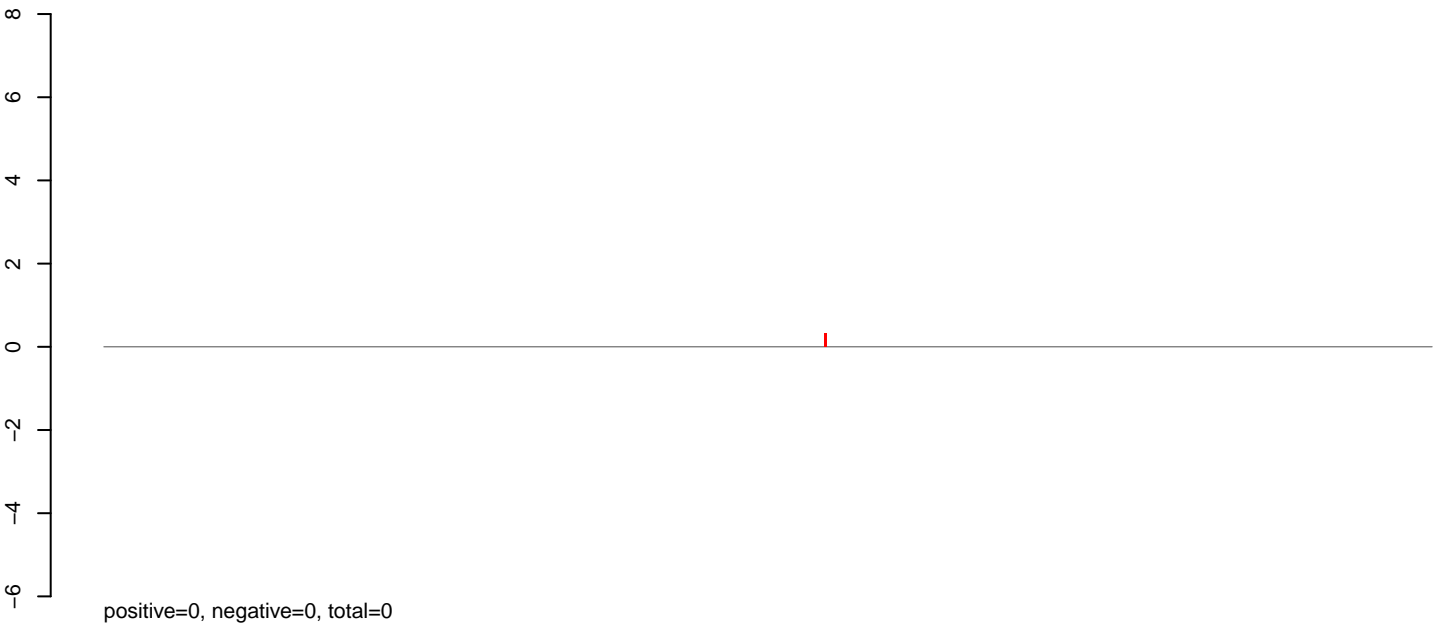


Window size=25, length=6065, virus@MH188054.1-Culex\_Y\_virus\_strain\_CYV\_Kero\_partial\_genome:1-6065

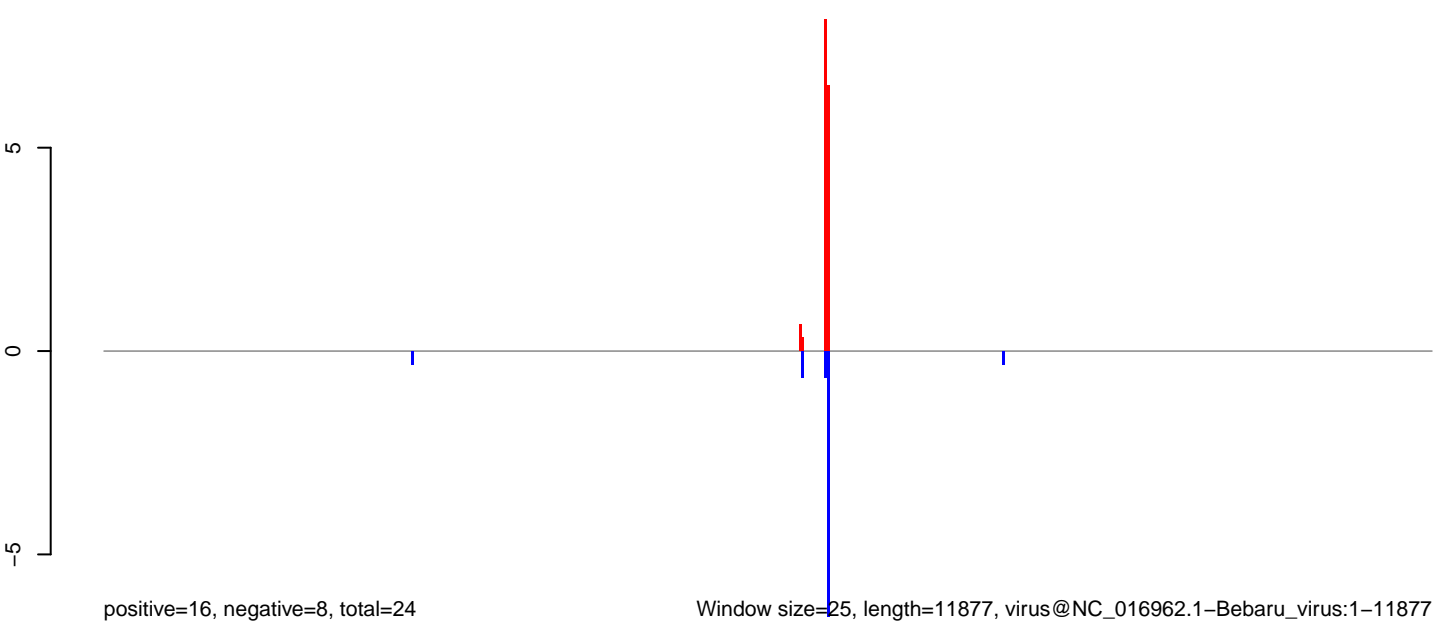
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



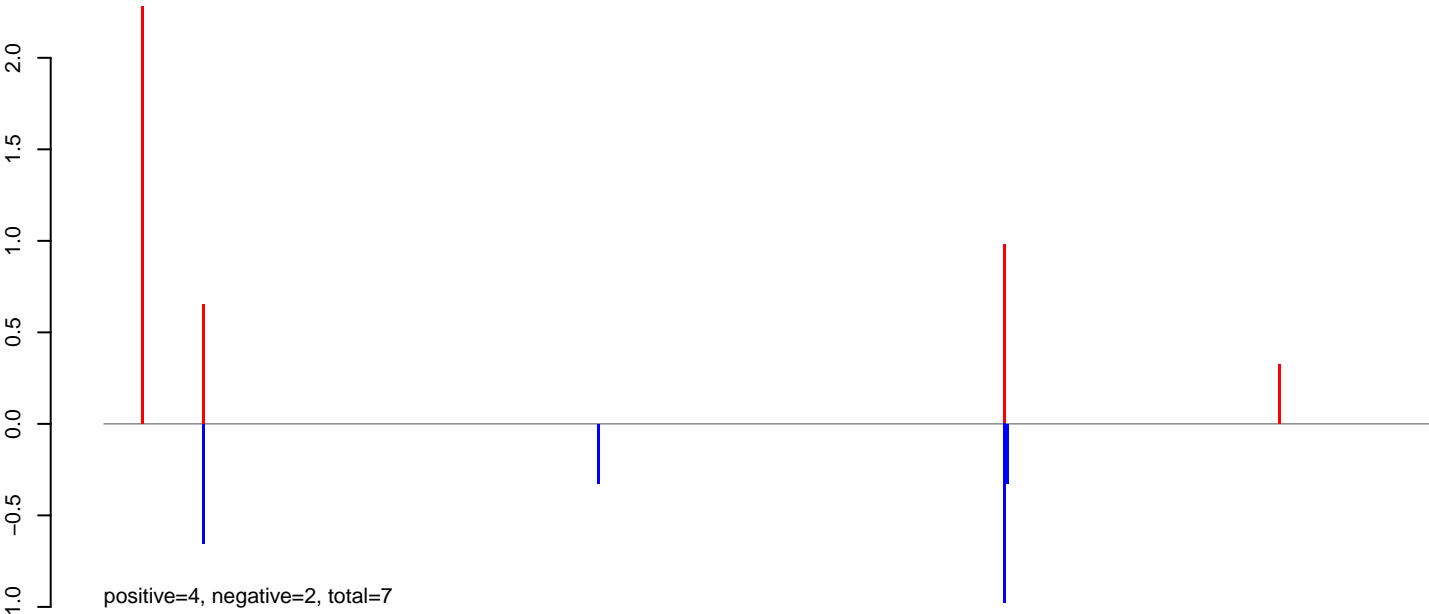
AnGam\_Mos55\_JR.rep



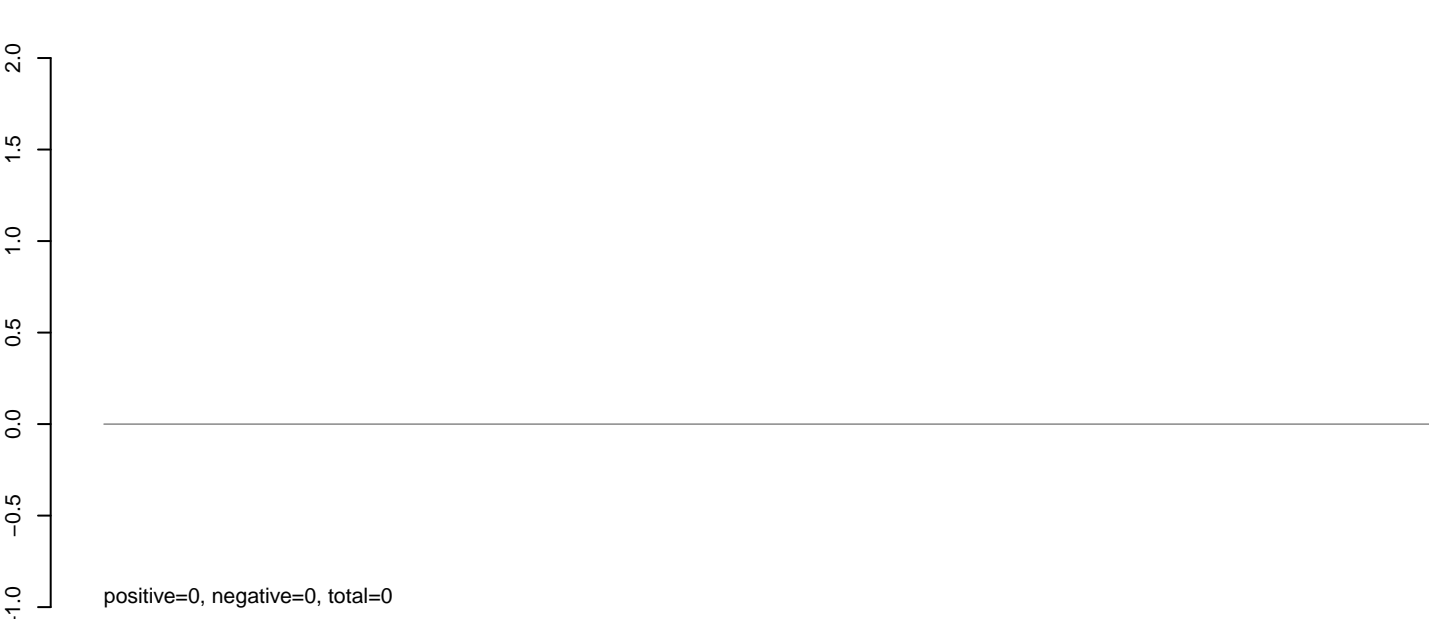
Window size=25, length=11877, virus@NC\_016962.1-Bebaru\_virus:1-11877

0 2000 4000 6000 8000 10000 12000

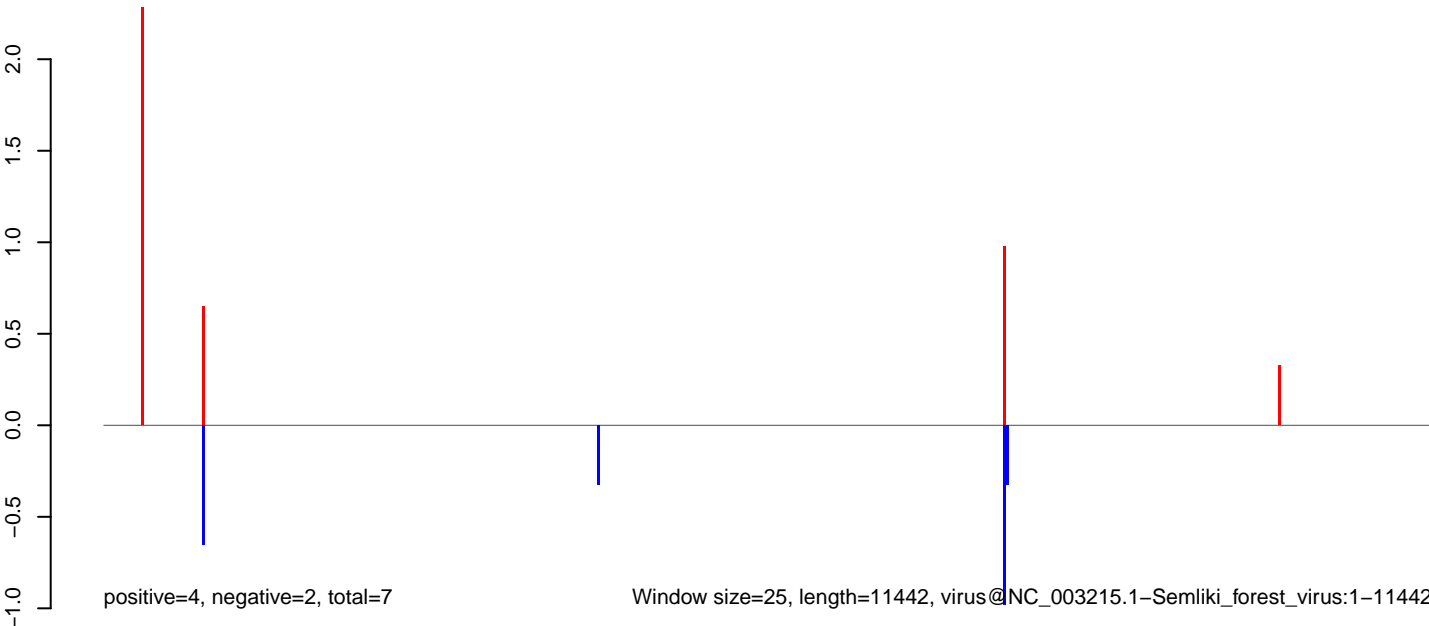
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep

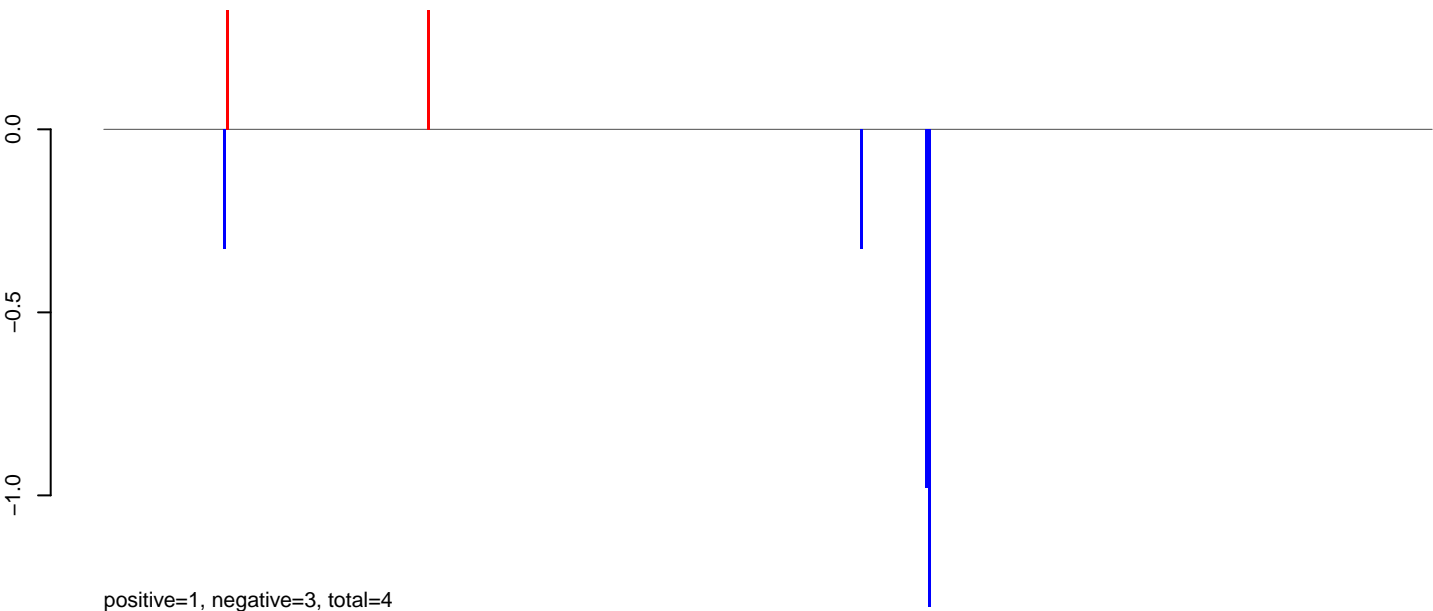


AnGam\_Mos55\_JR.rep

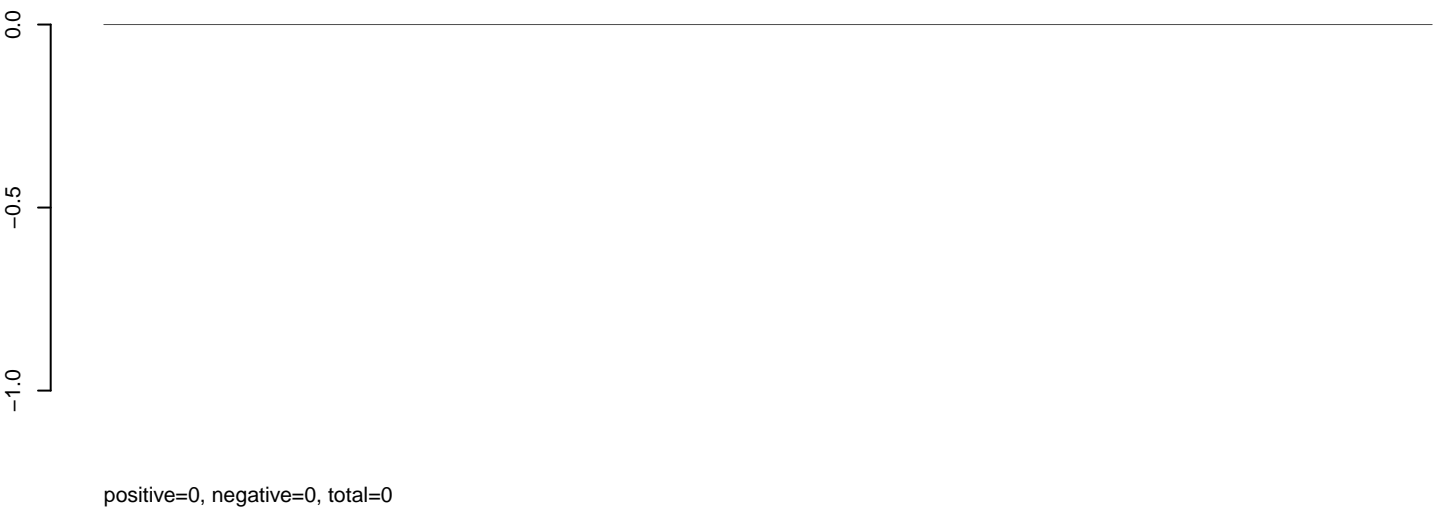


Window size=25, length=11442, virus@NC\_003215.1-Semliki\_forest\_virus:1-11442

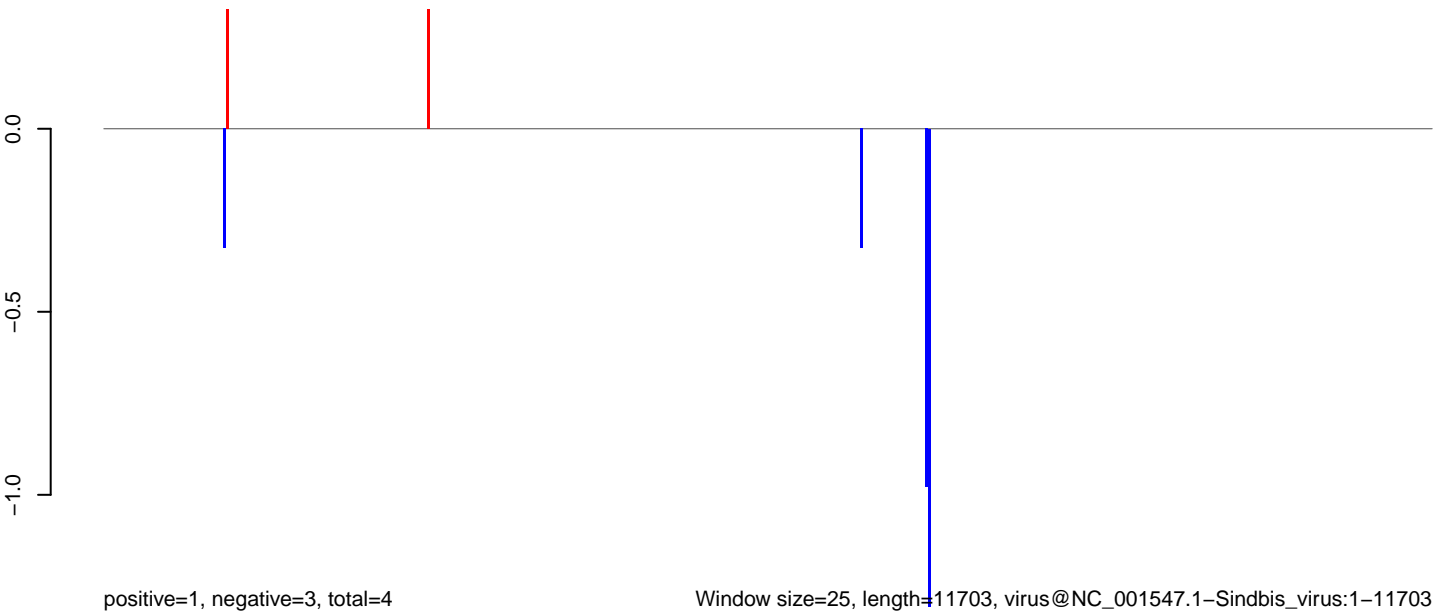
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



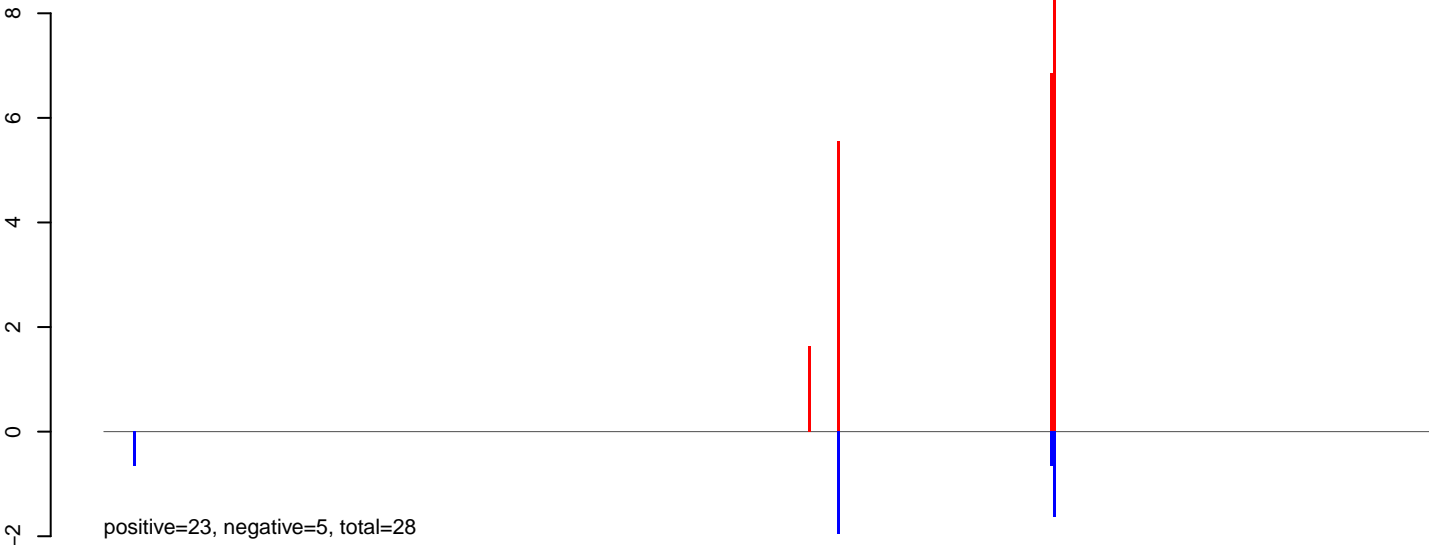
AnGam\_Mos55\_JR.rep



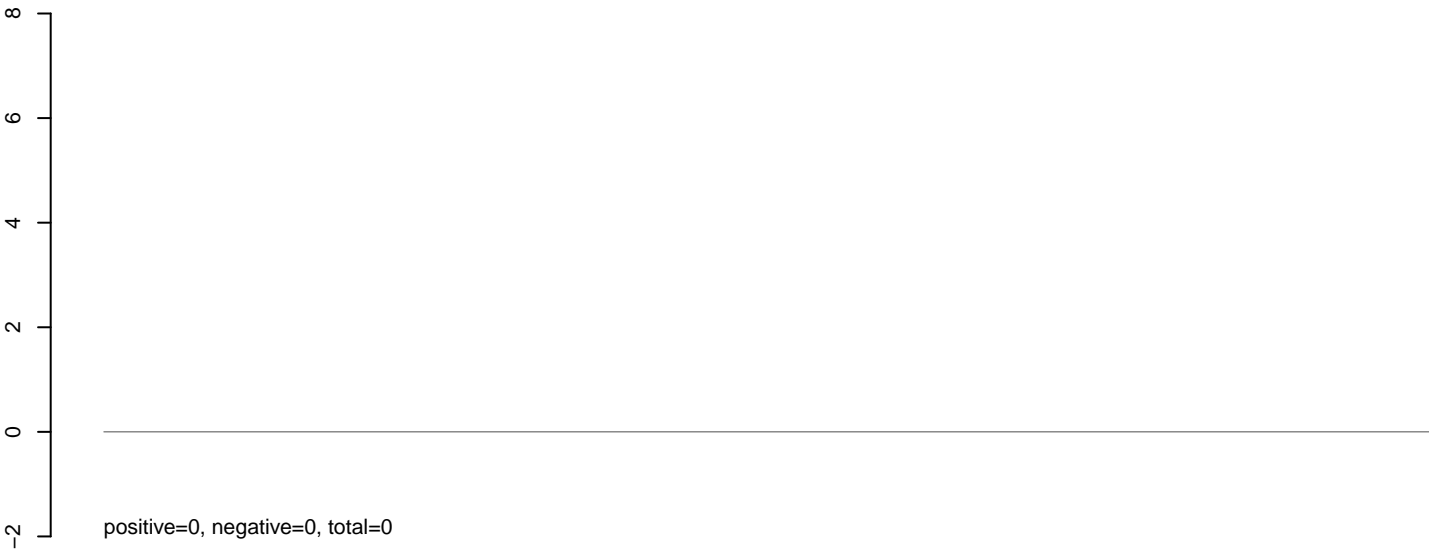
Window size=25, length=11703, virus@NC\_001547.1-Sindbis\_virus:1-11703

0 2000 4000 6000 8000 10000 12000

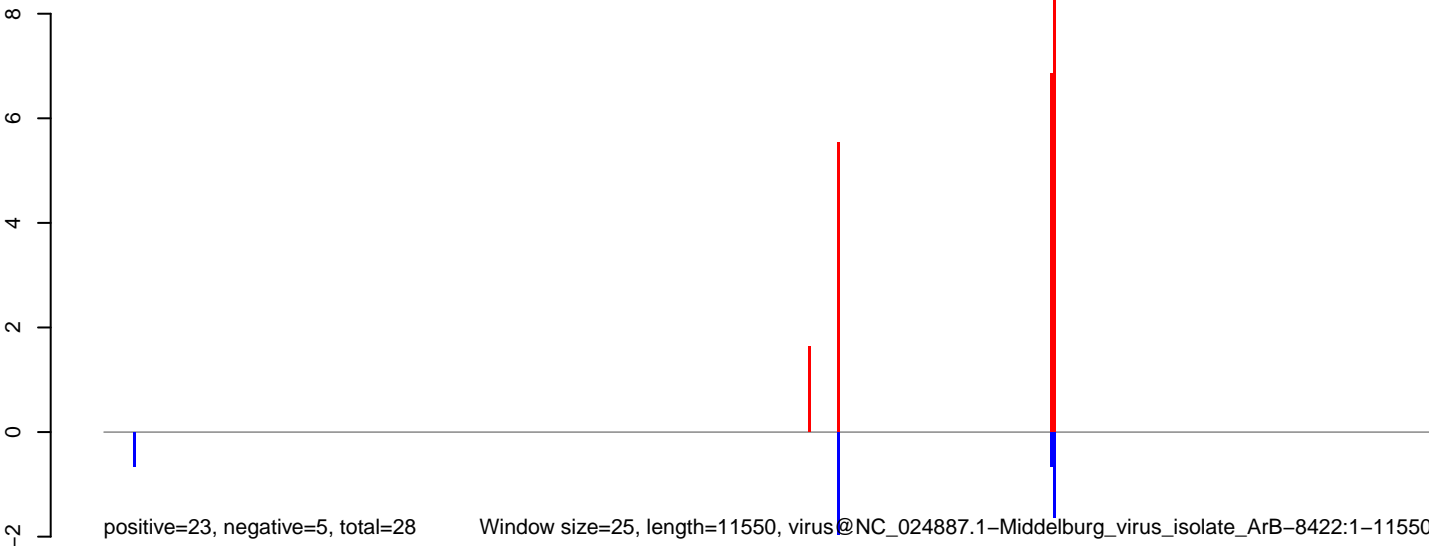
AnGam\_Mos55\_JR.18\_23.rep



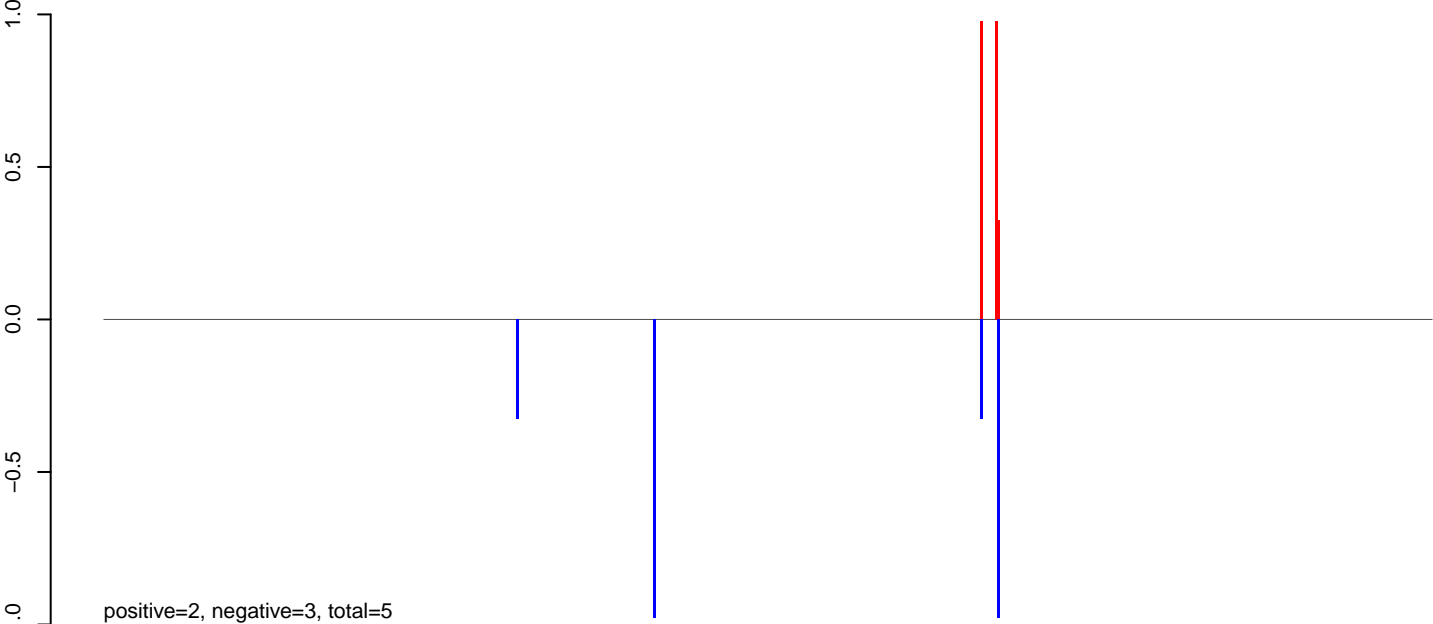
AnGam\_Mos55\_JR.24\_35.rep



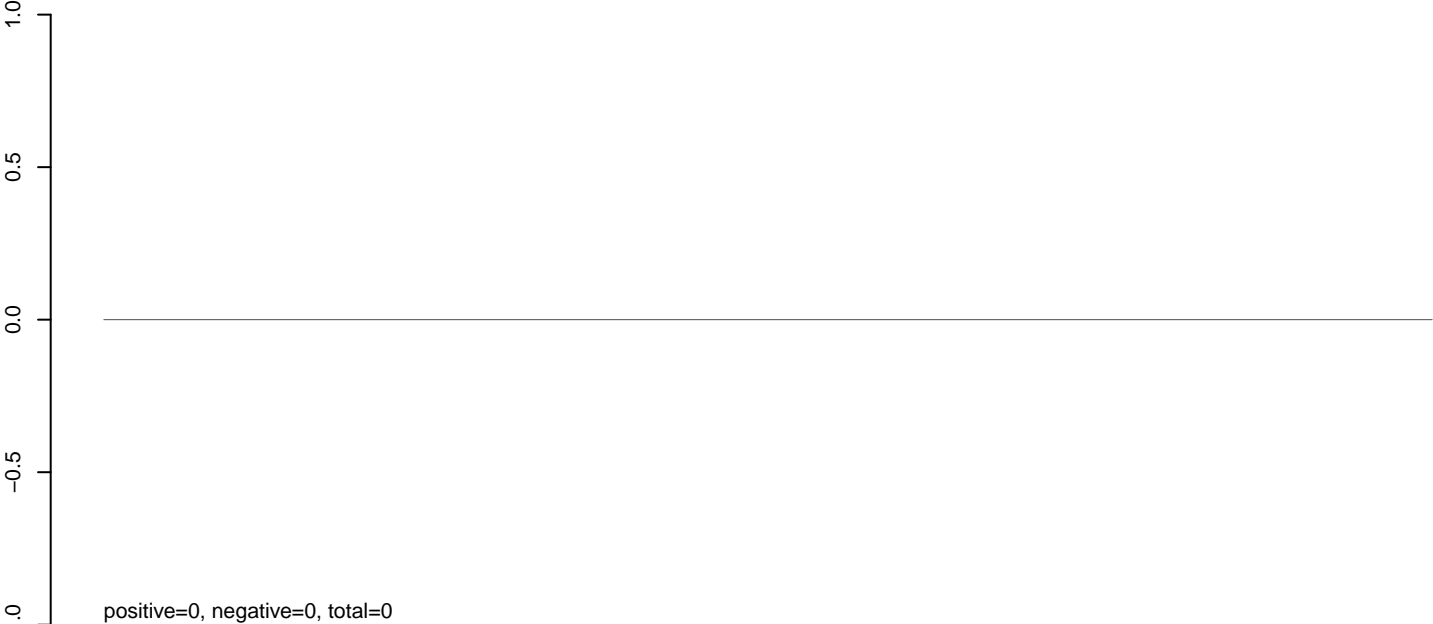
AnGam\_Mos55\_JR.rep



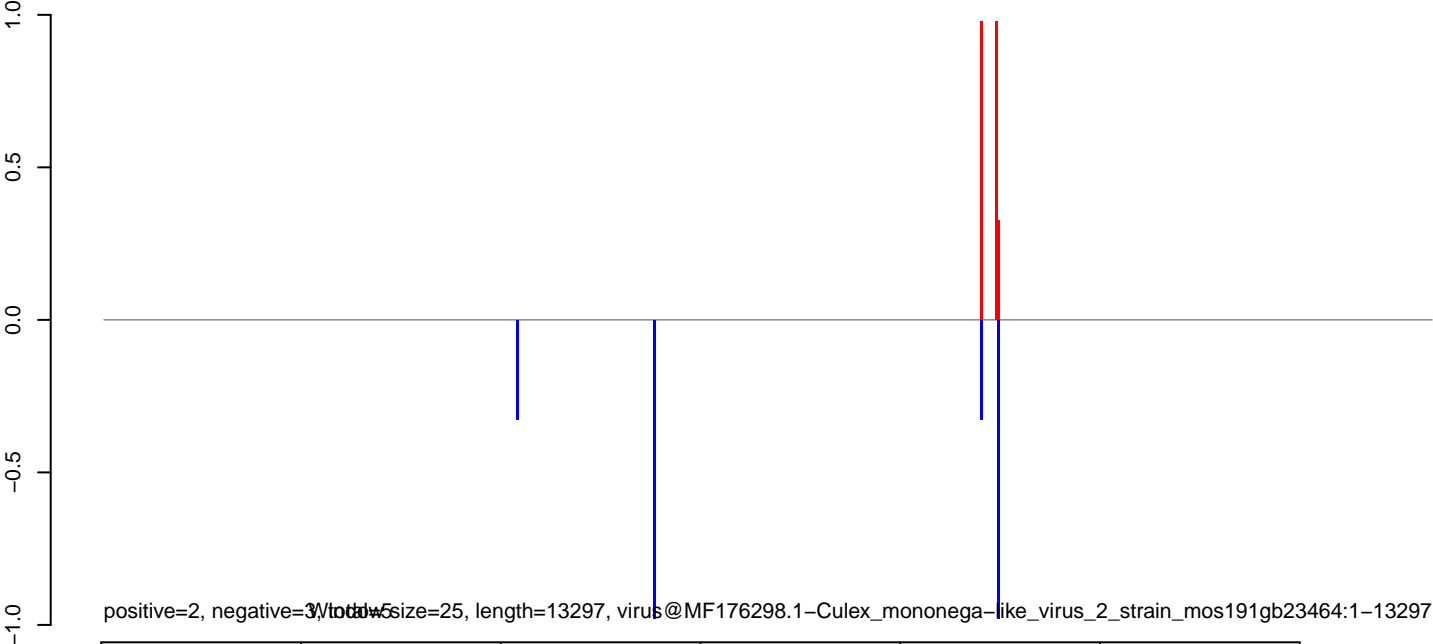
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



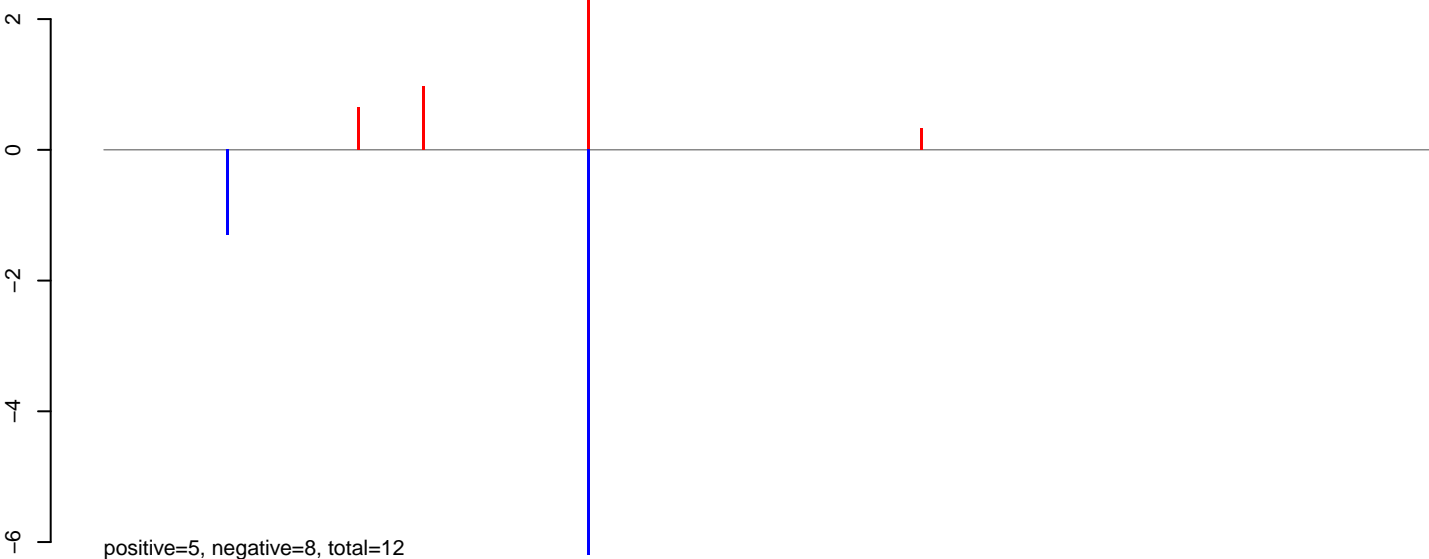
AnGam\_Mos55\_JR.rep



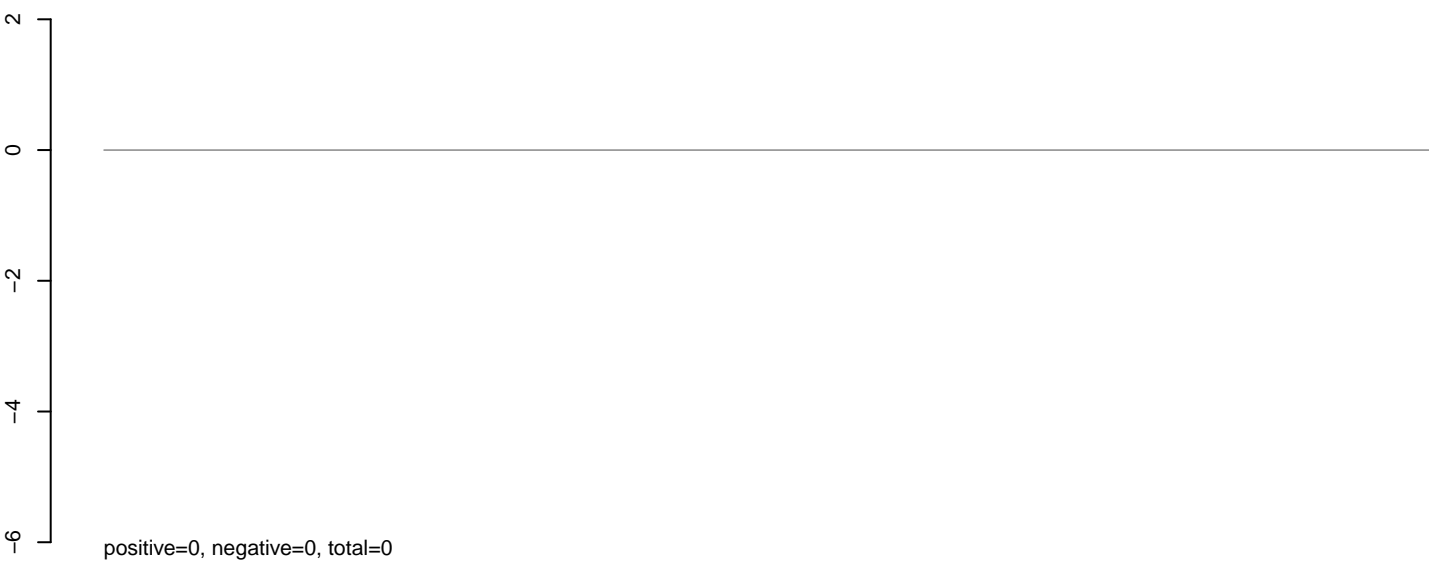
positive=2, negative=3, total=5, window size=25, length=13297, virus@MF176298.1-Culex\_mononega-like\_virus\_2\_strain\_mos191gb23464:1-13297

0 2000 4000 6000 8000 10000 12000

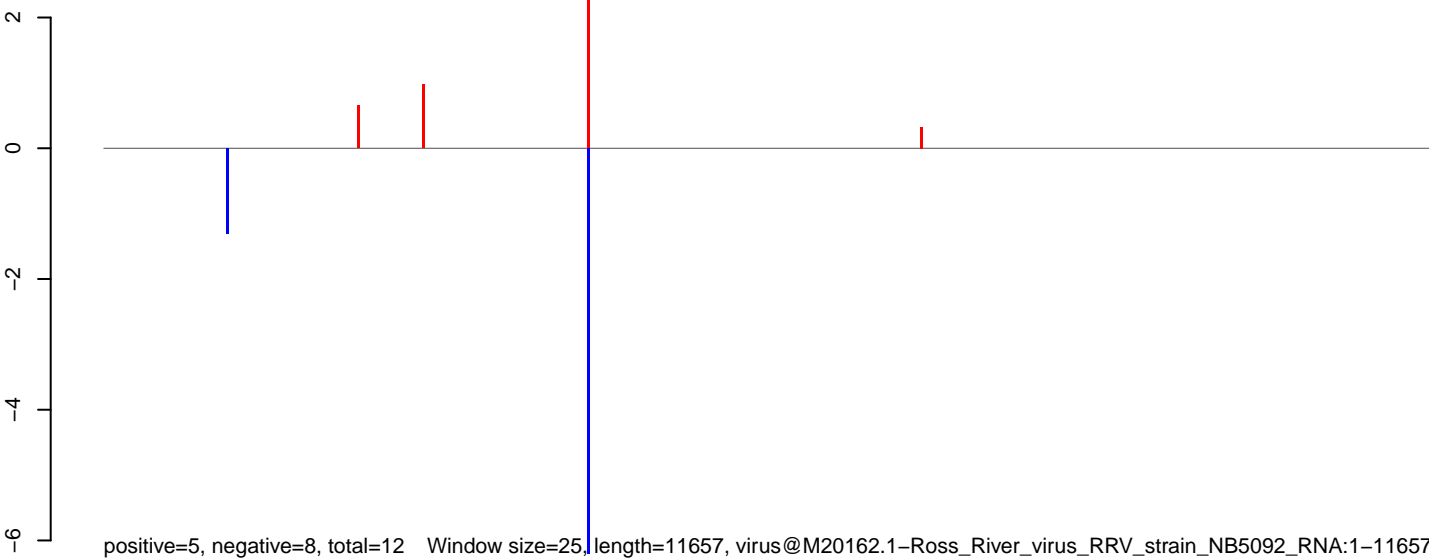
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep

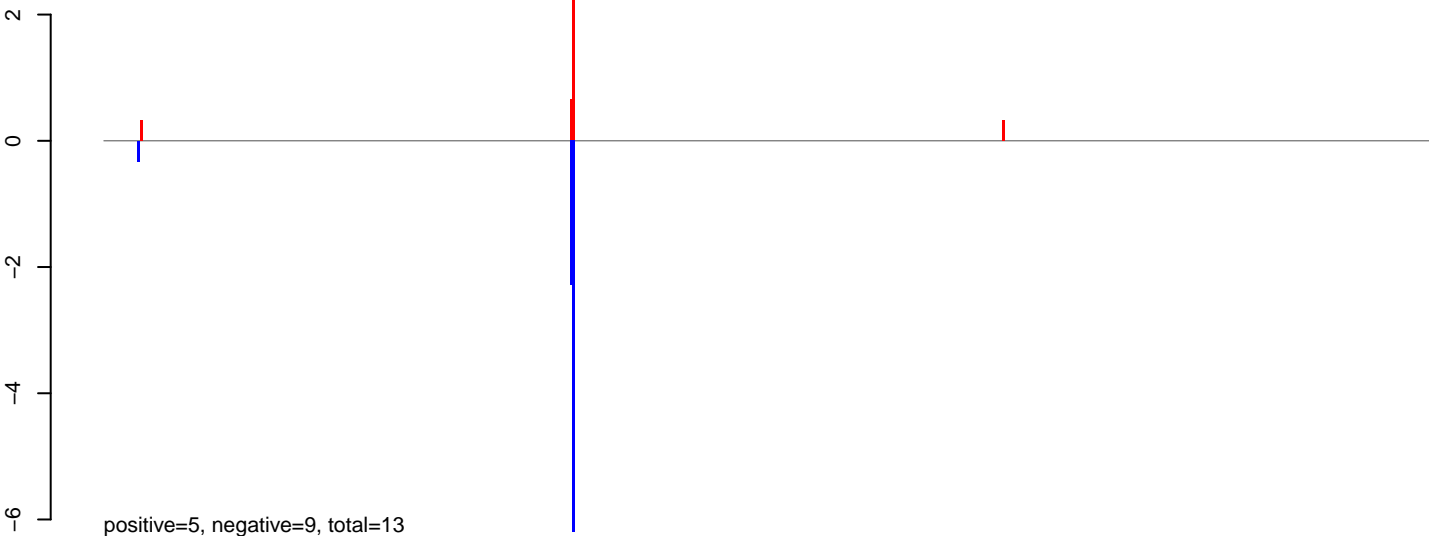


AnGam\_Mos55\_JR.rep

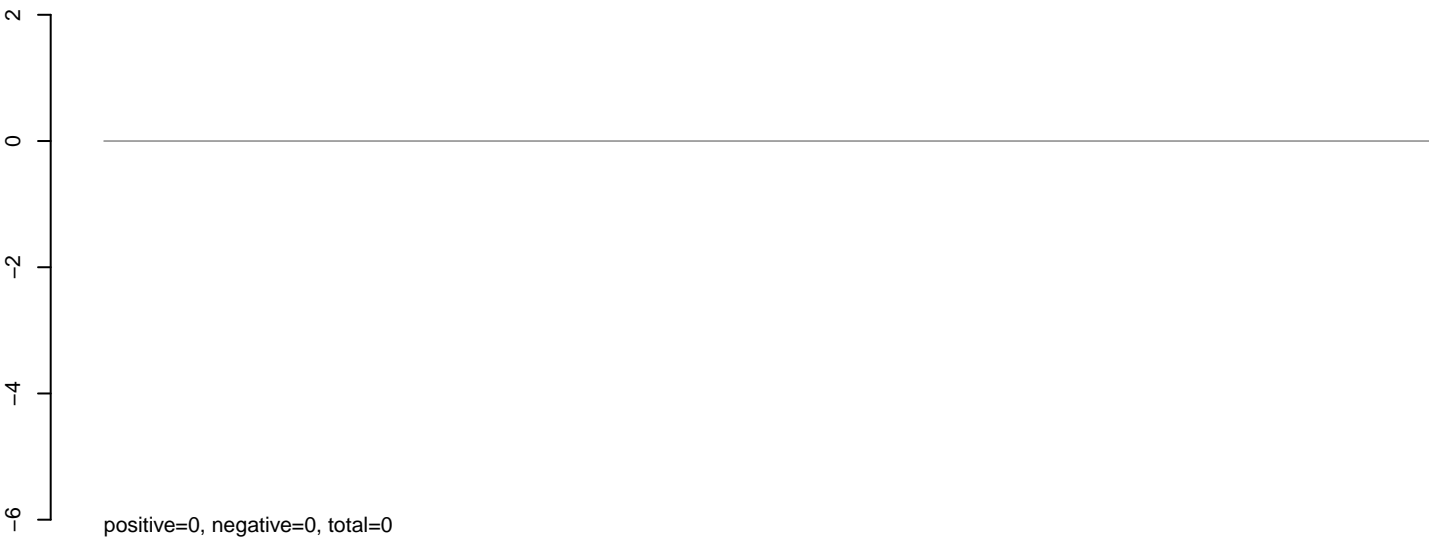


Window size=25, length=11657, virus@M20162.1-Ross\_River\_virus\_RRV\_strain\_NB5092\_RNA:1-11657

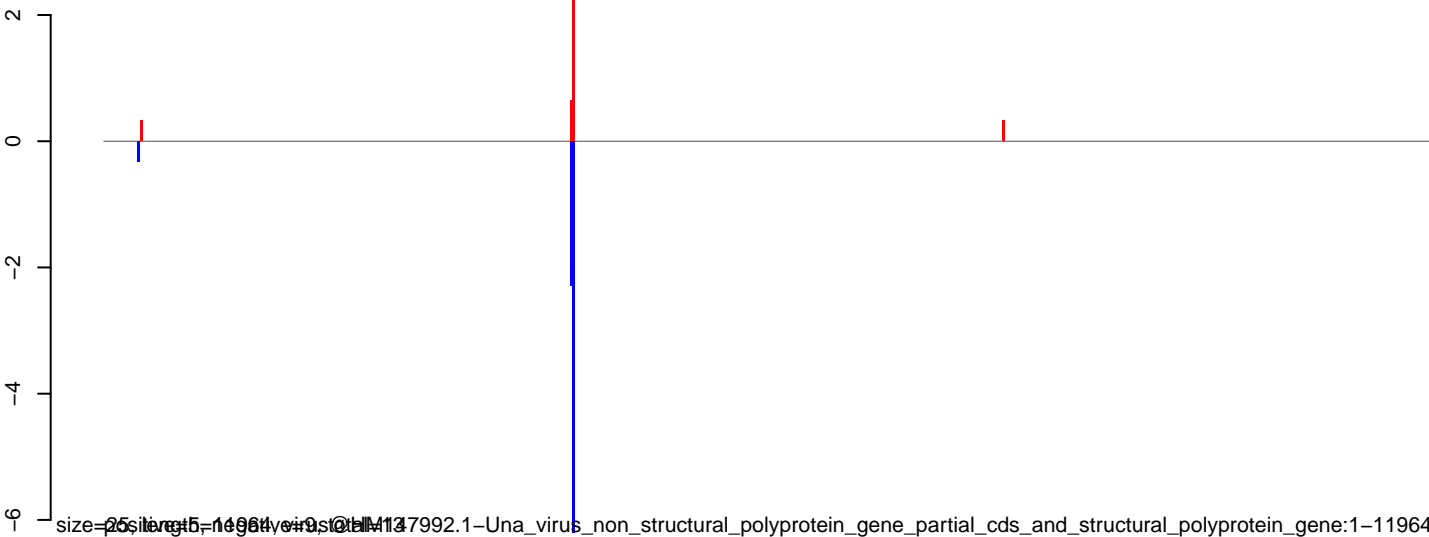
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



AnGam\_Mos55\_JR.rep

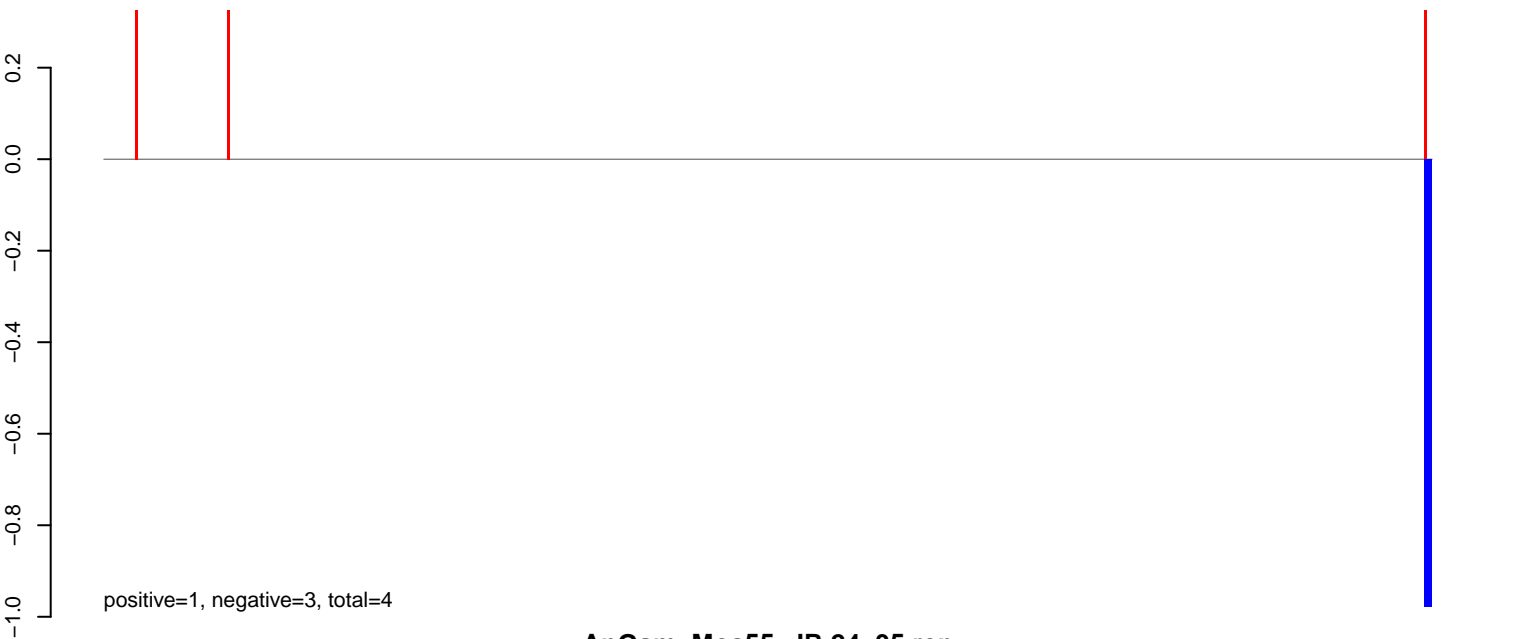


size=25, length=14961, view=9, obs=13  
1-Una\_virus\_non\_structural\_polyprotein\_gene\_partial\_cds\_and\_structural\_polyprotein\_gene:1-11964

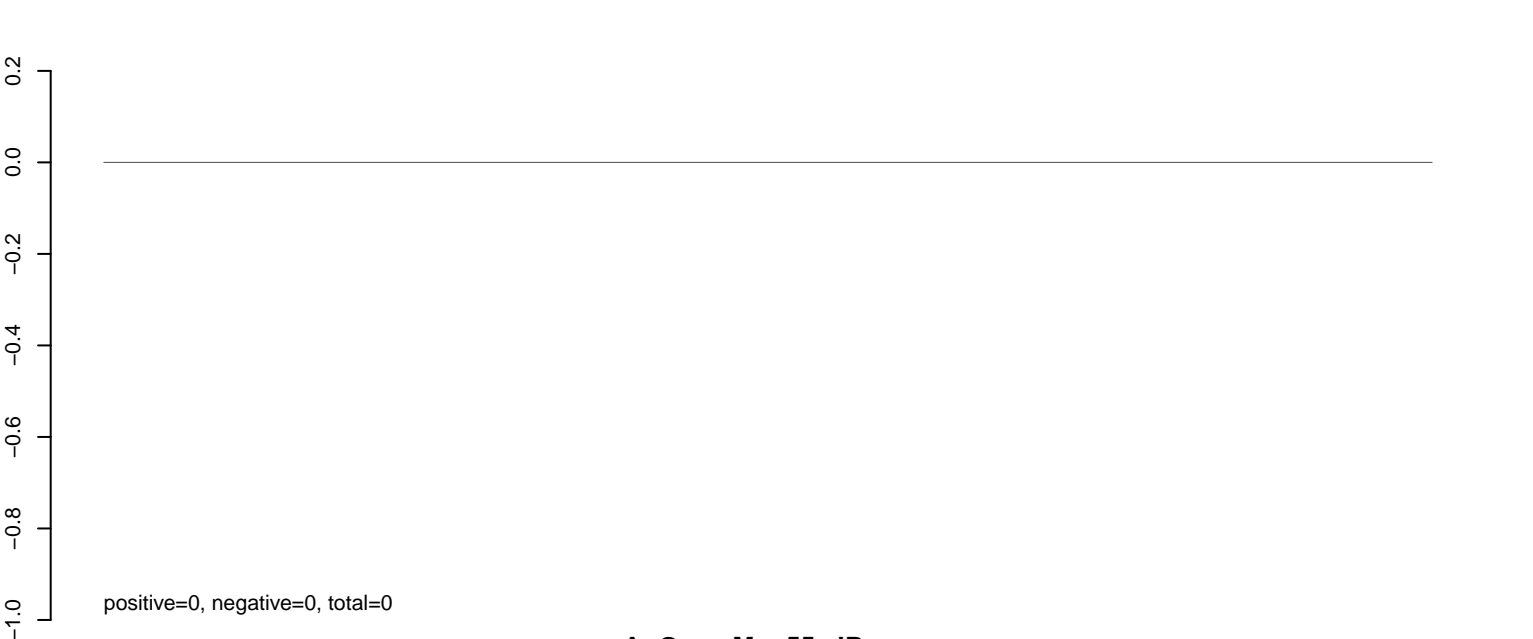
0 2000 4000 6000 8000 10000 12000



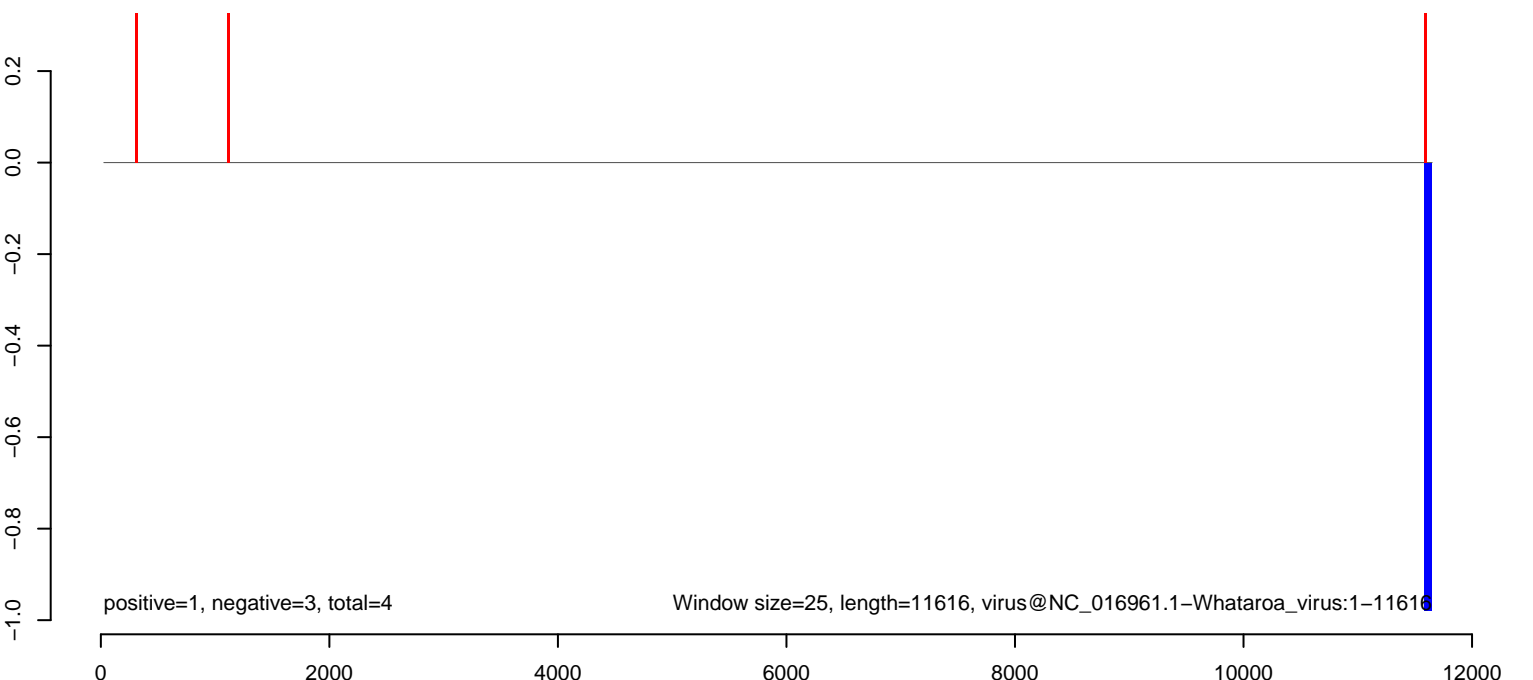
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



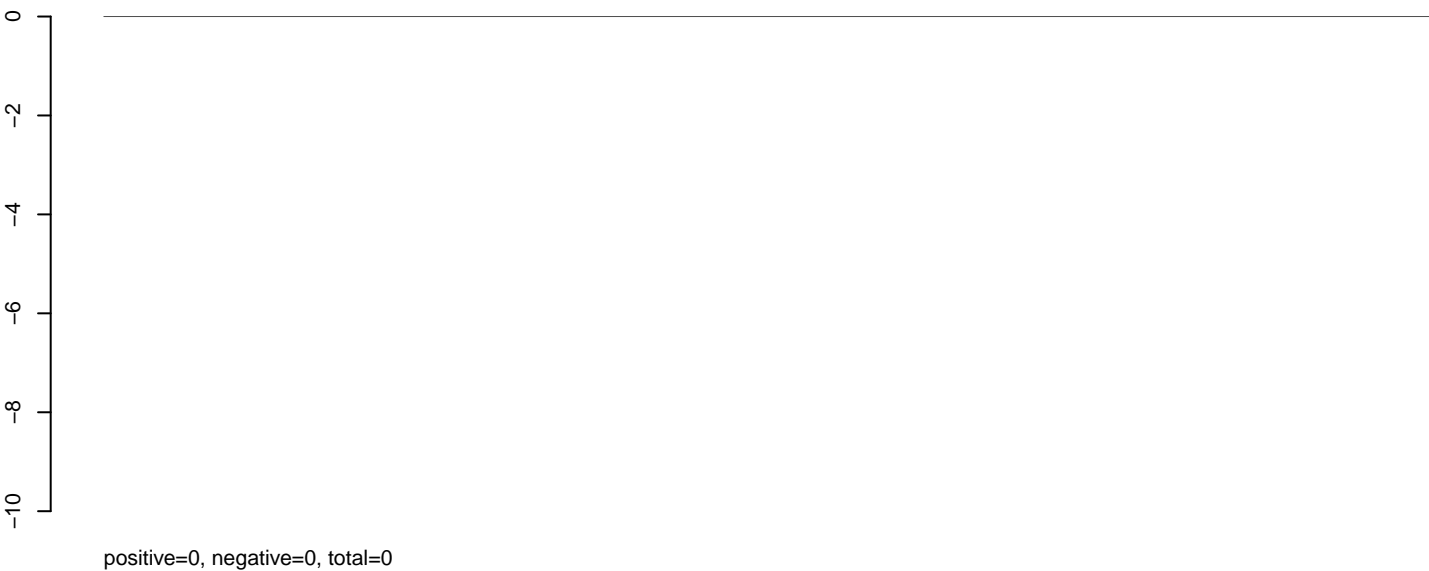
AnGam\_Mos55\_JR.rep



AnGam\_Mos55\_JR.18\_23.rep



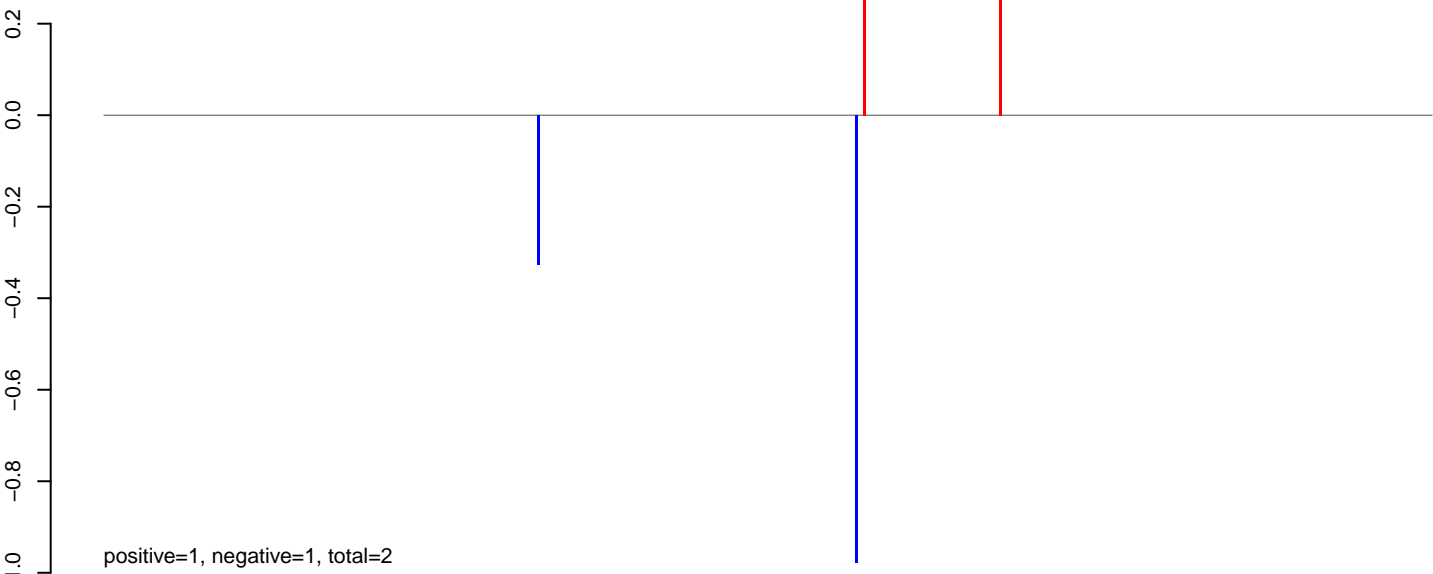
AnGam\_Mos55\_JR.24\_35.rep



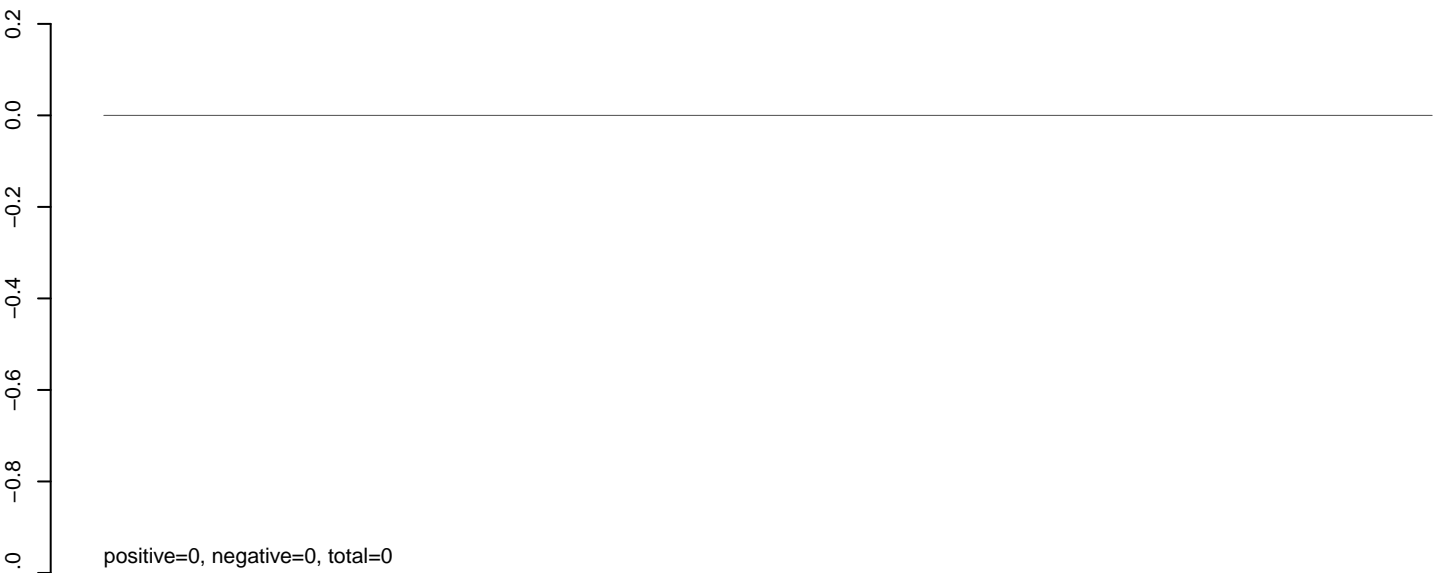
AnGam\_Mos55\_JR.rep



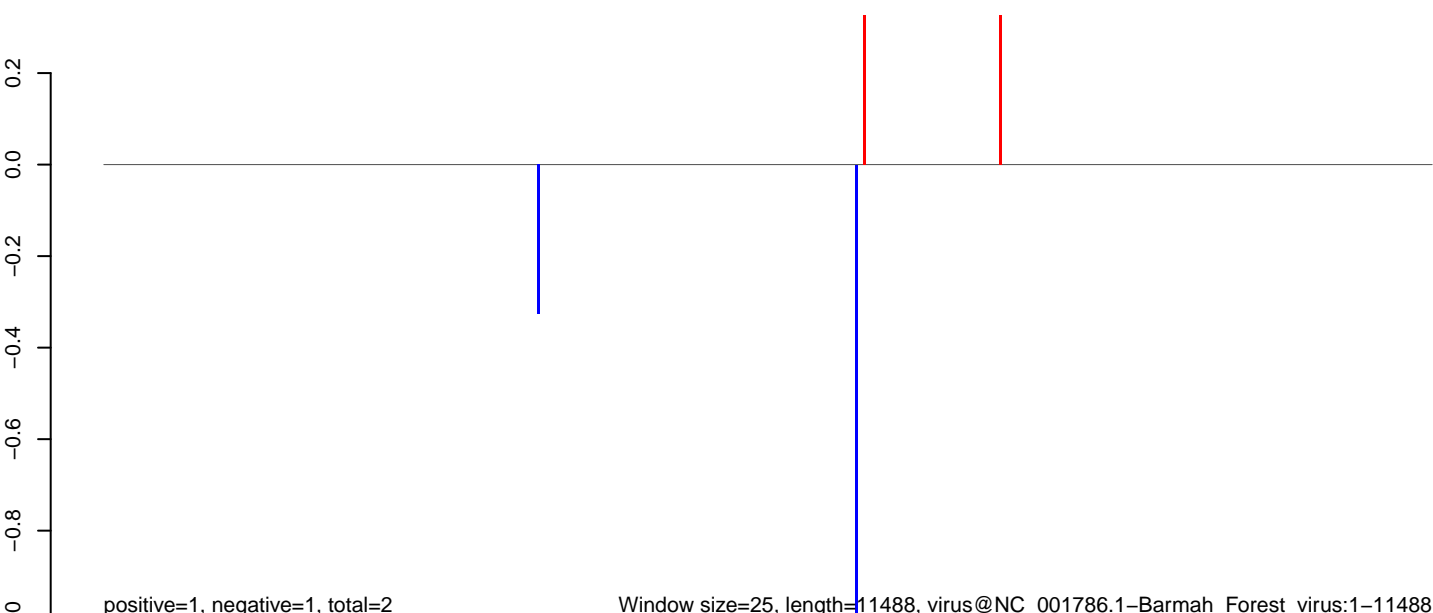
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



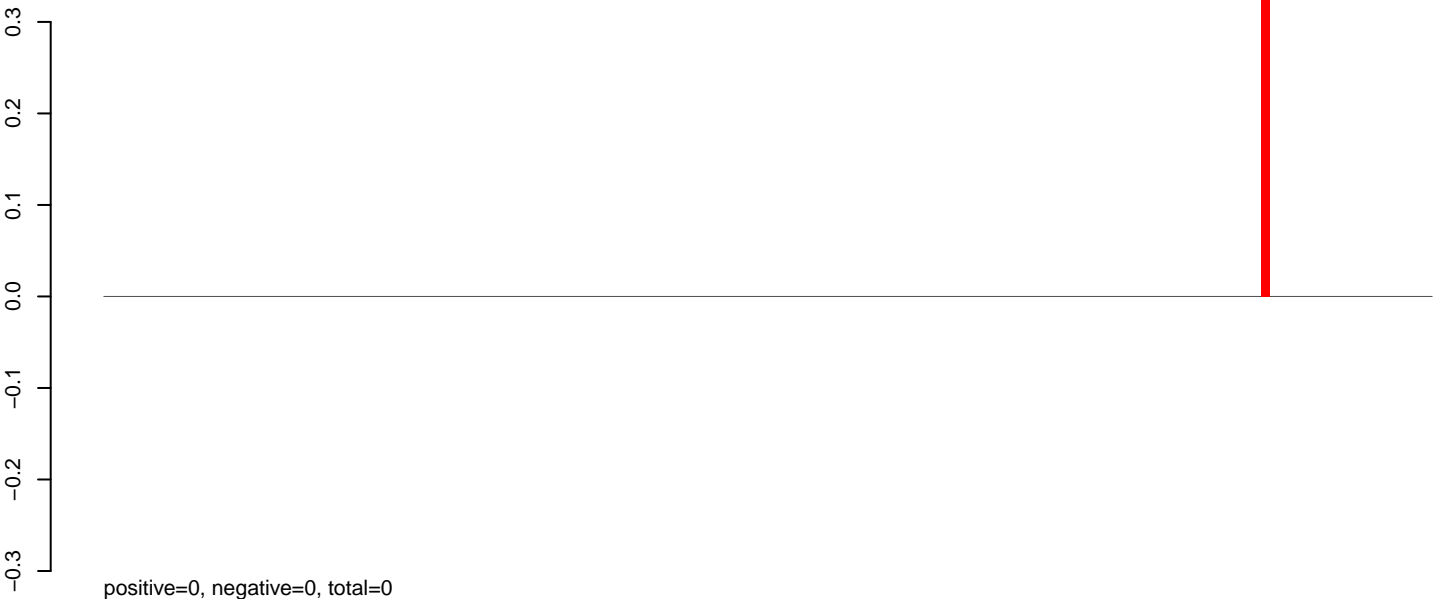
AnGam\_Mos55\_JR.rep



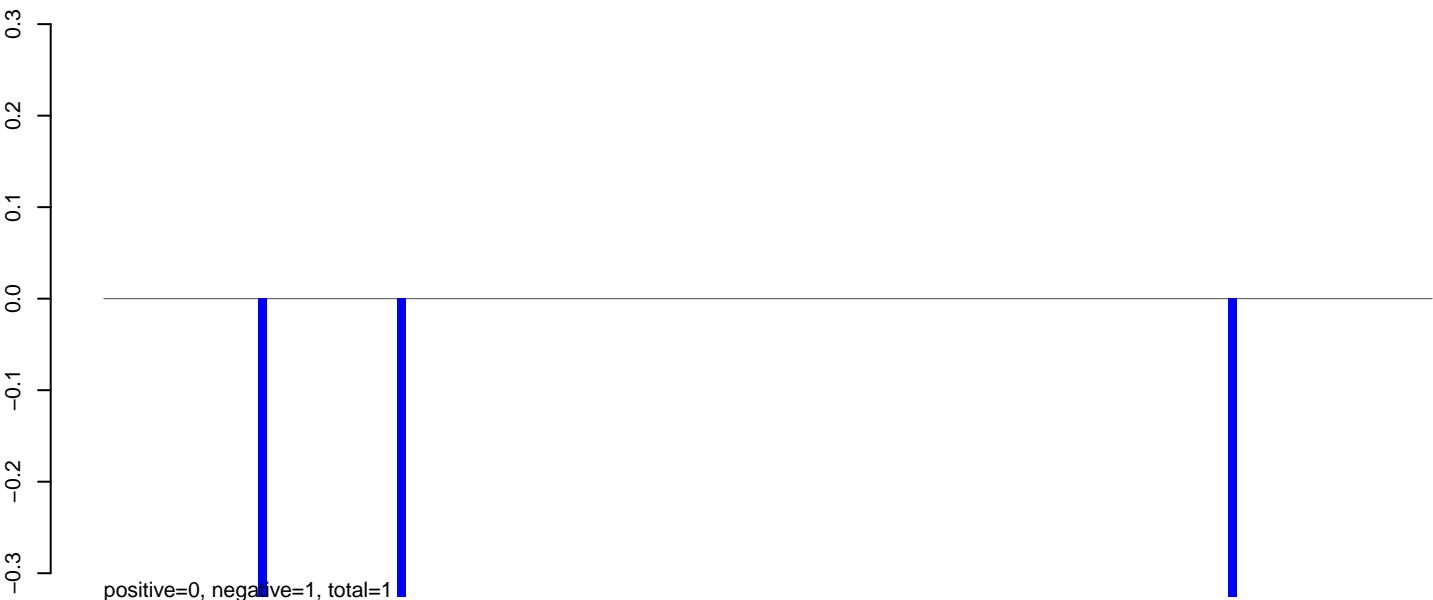
Window size=25, length=11488, virus@NC\_001786.1-Barmah\_Forest\_virus:1-11488

0 2000 4000 6000 8000 10000

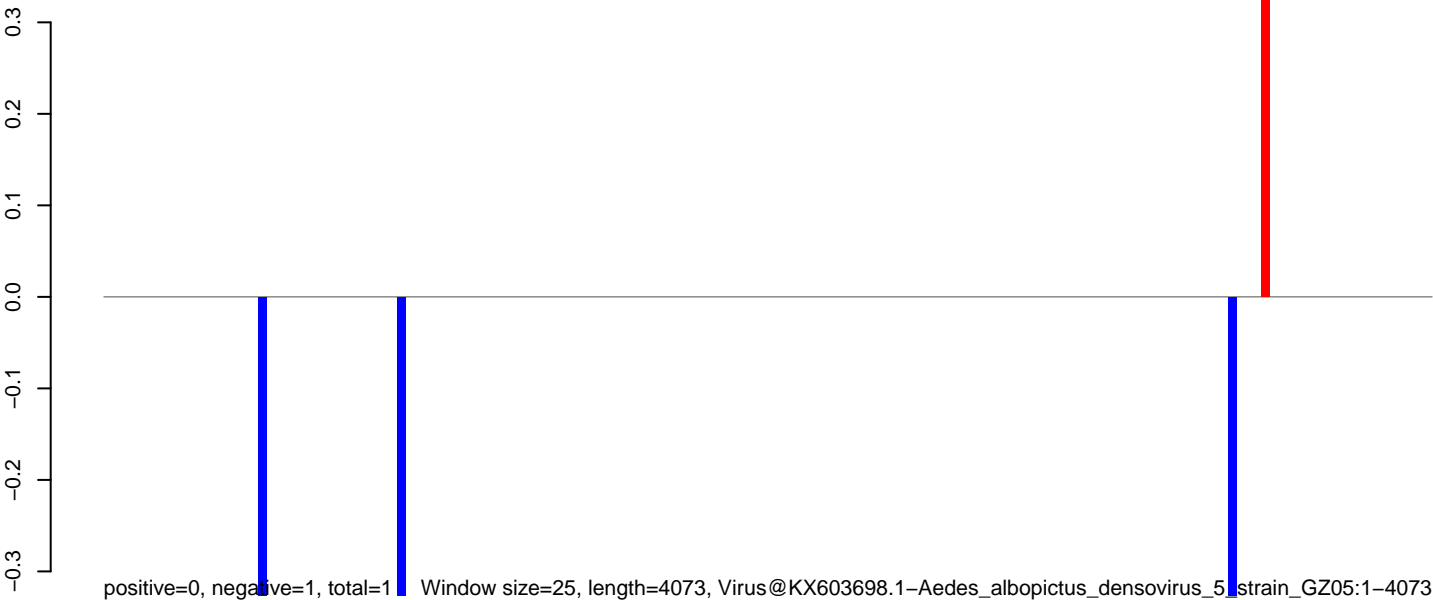
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



AnGam\_Mos55\_JR.rep

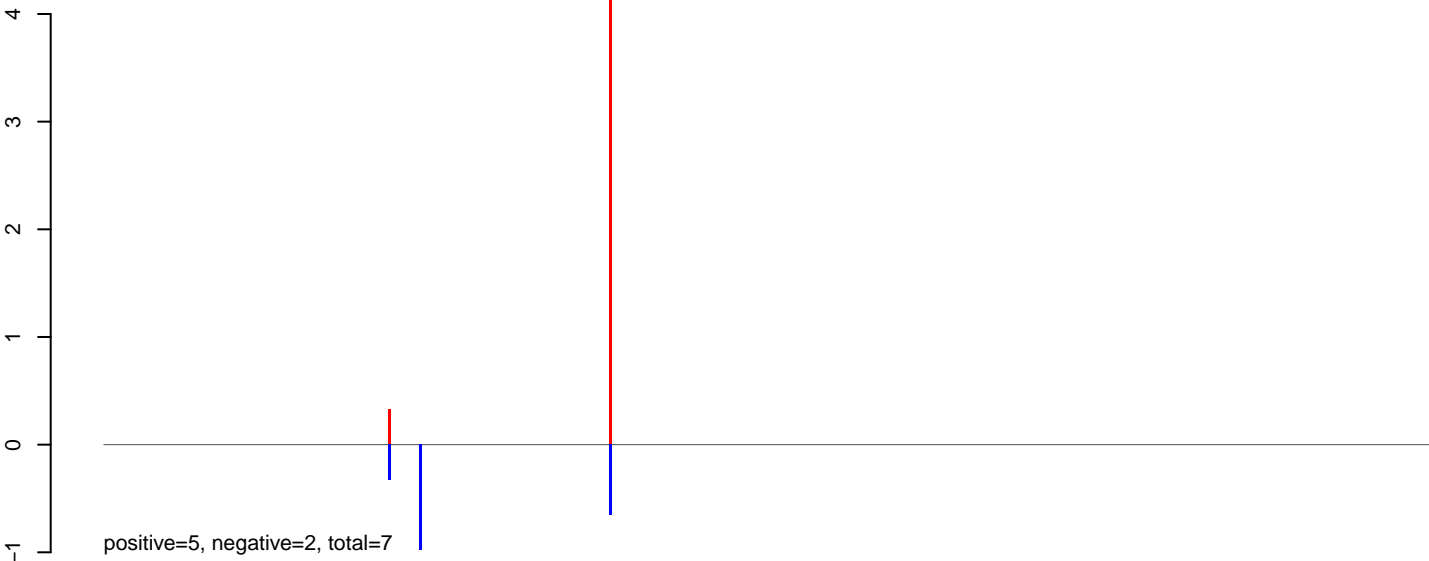


Window size=25, length=4073, Virus@KX603698.1-Aedes albopictus\_densovirus\_5 strain\_GZ05:1-4073

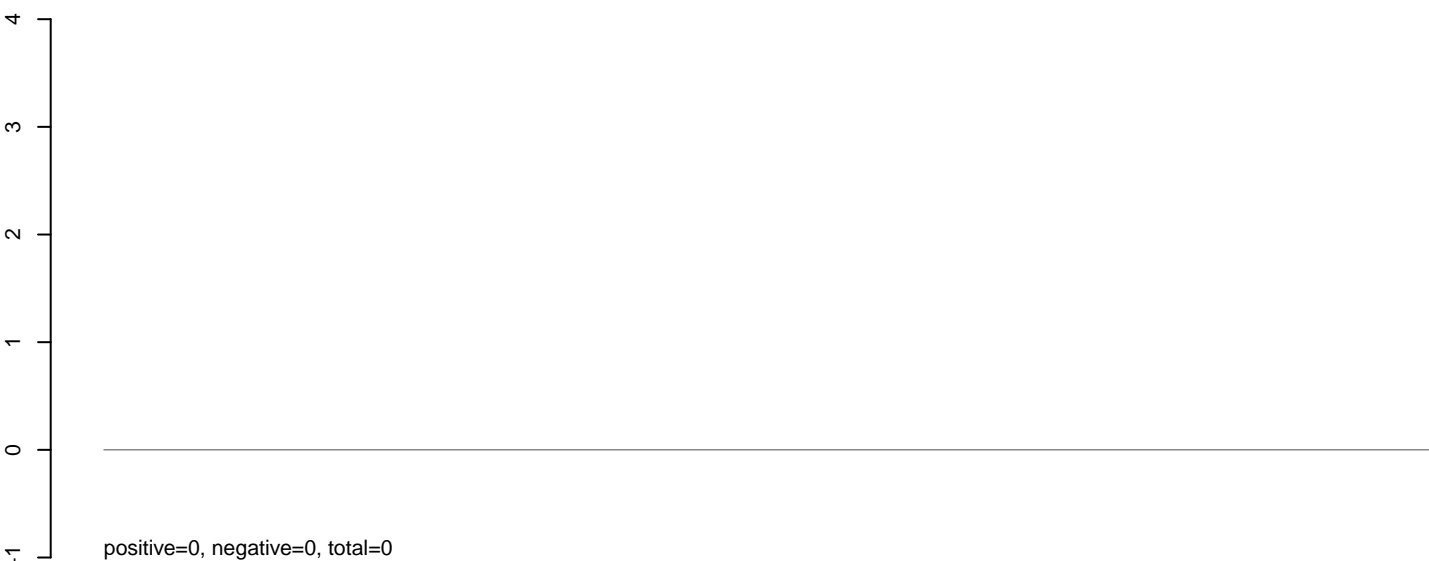
0 1000 2000 3000 4000



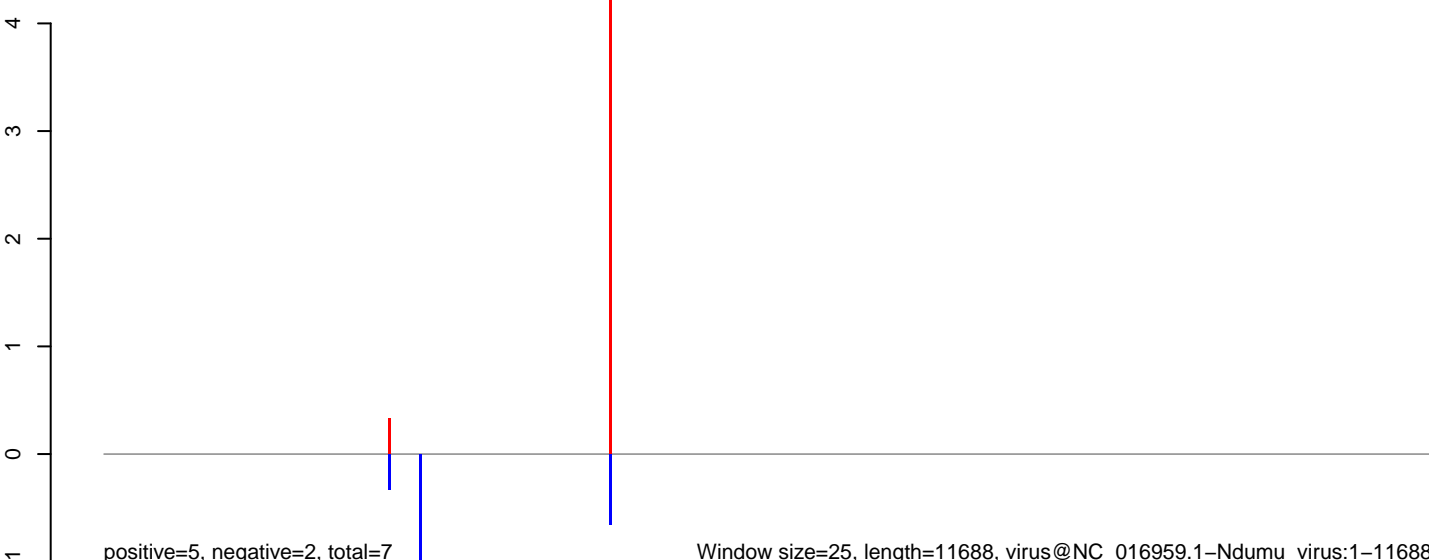
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



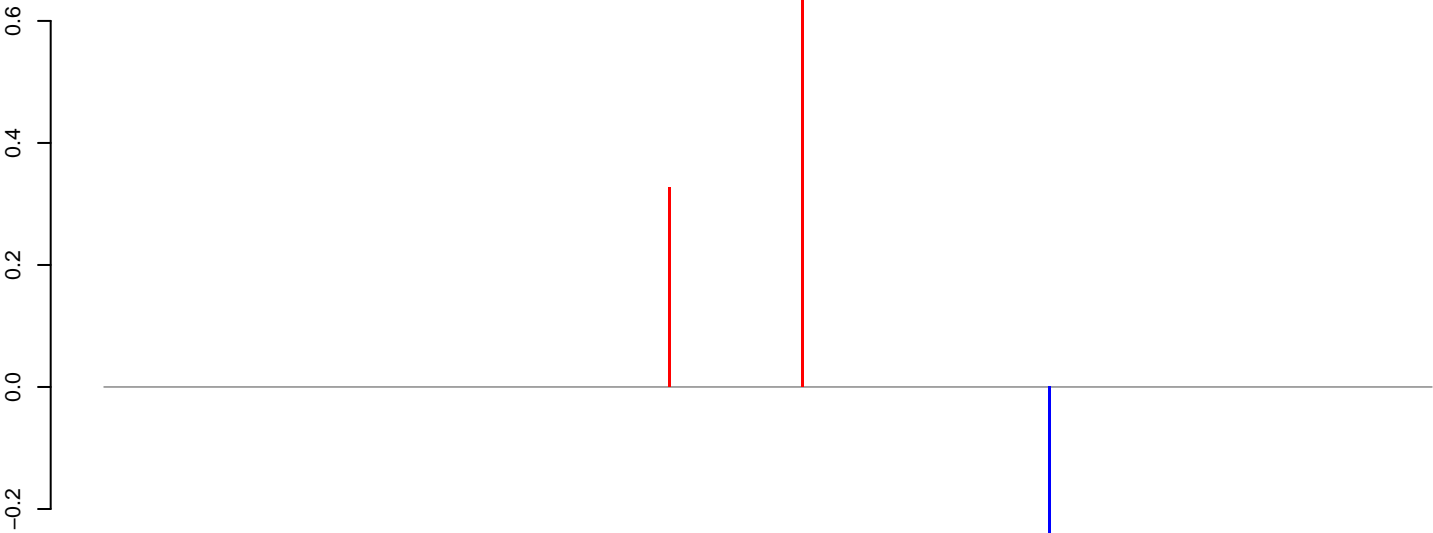
AnGam\_Mos55\_JR.rep



Window size=25, length=11688, virus@NC\_016959.1-Ndumu\_virus:1-11688

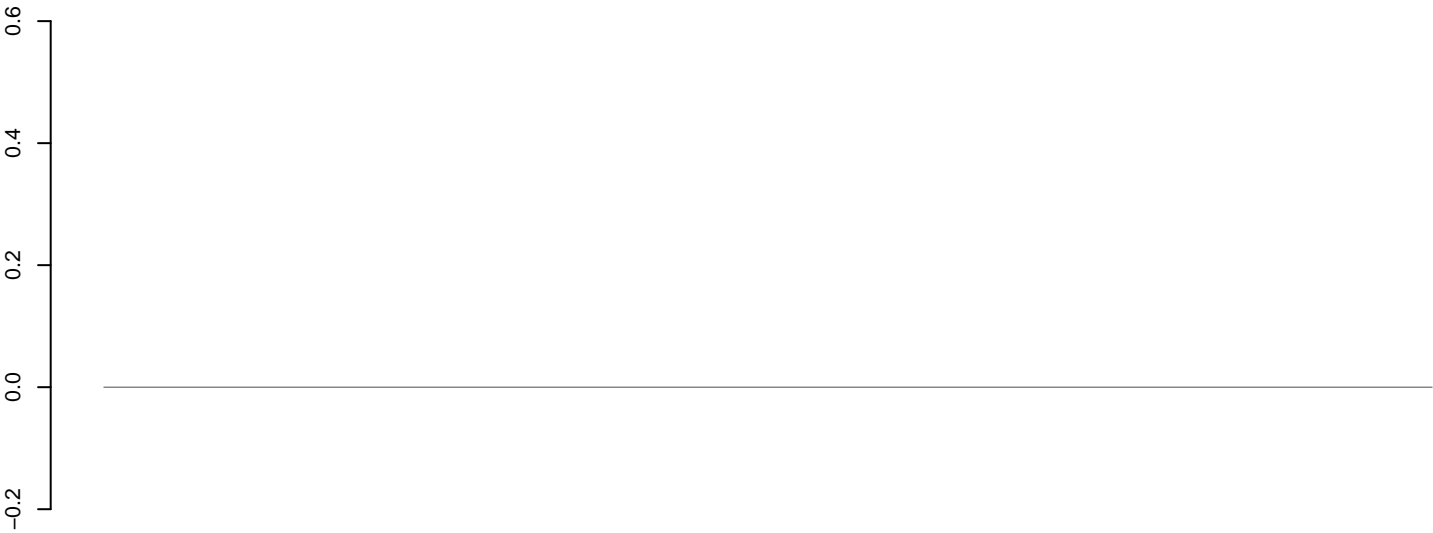
0 2000 4000 6000 8000 10000 12000

AnGam\_Mos55\_JR.18\_23.rep



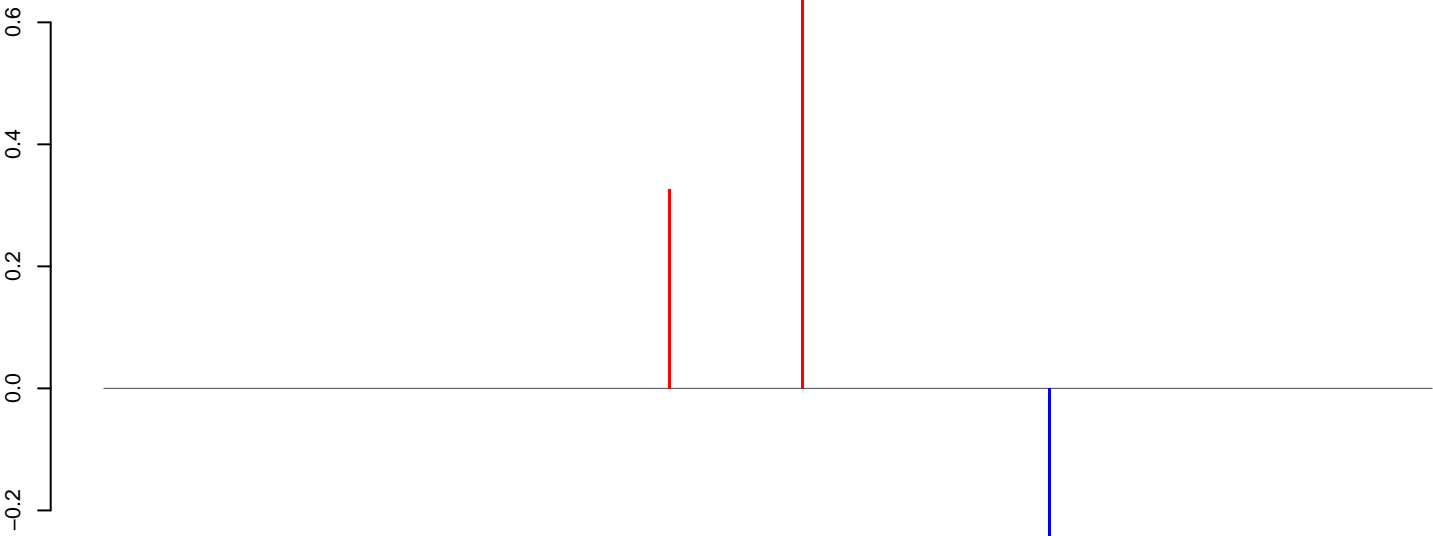
positive=1, negative=0, total=1

AnGam\_Mos55\_JR.24\_35.rep



positive=0, negative=0, total=0

AnGam\_Mos55\_JR.rep

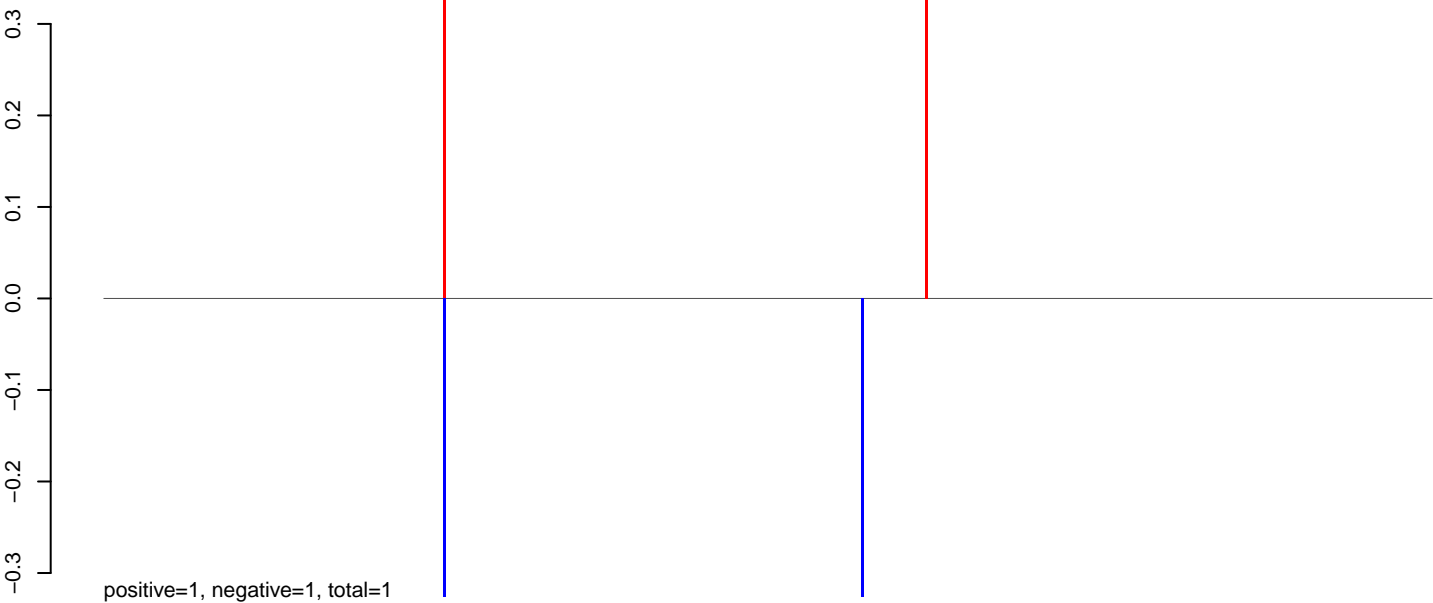


positive=1, negative=0, total=1

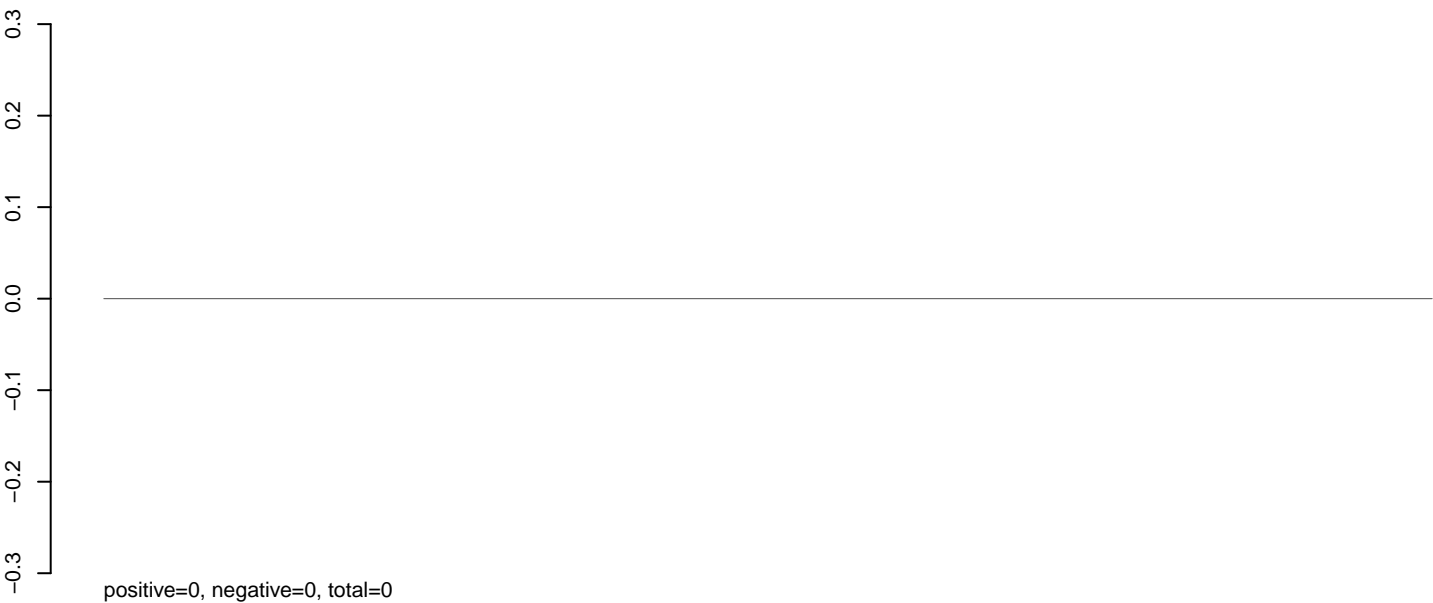
Window size=25, length=11411, virus@NC\_003417.1-Mayaro\_virus:1-11411



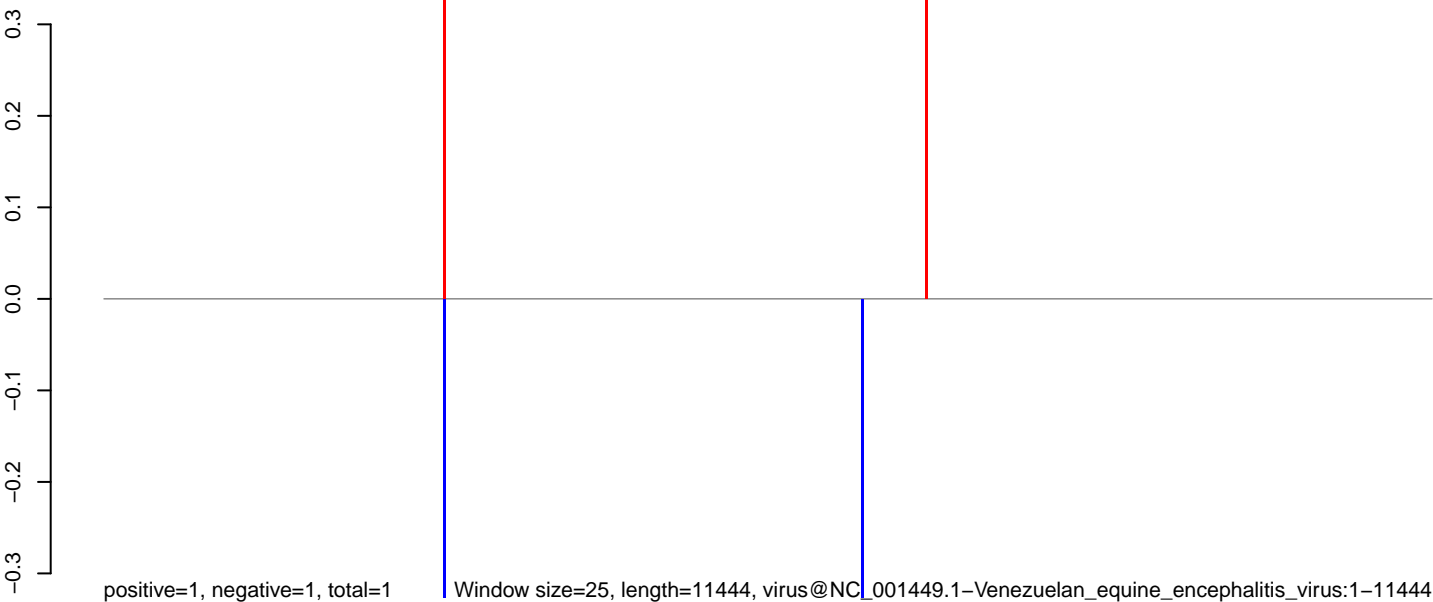
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



AnGam\_Mos55\_JR.rep

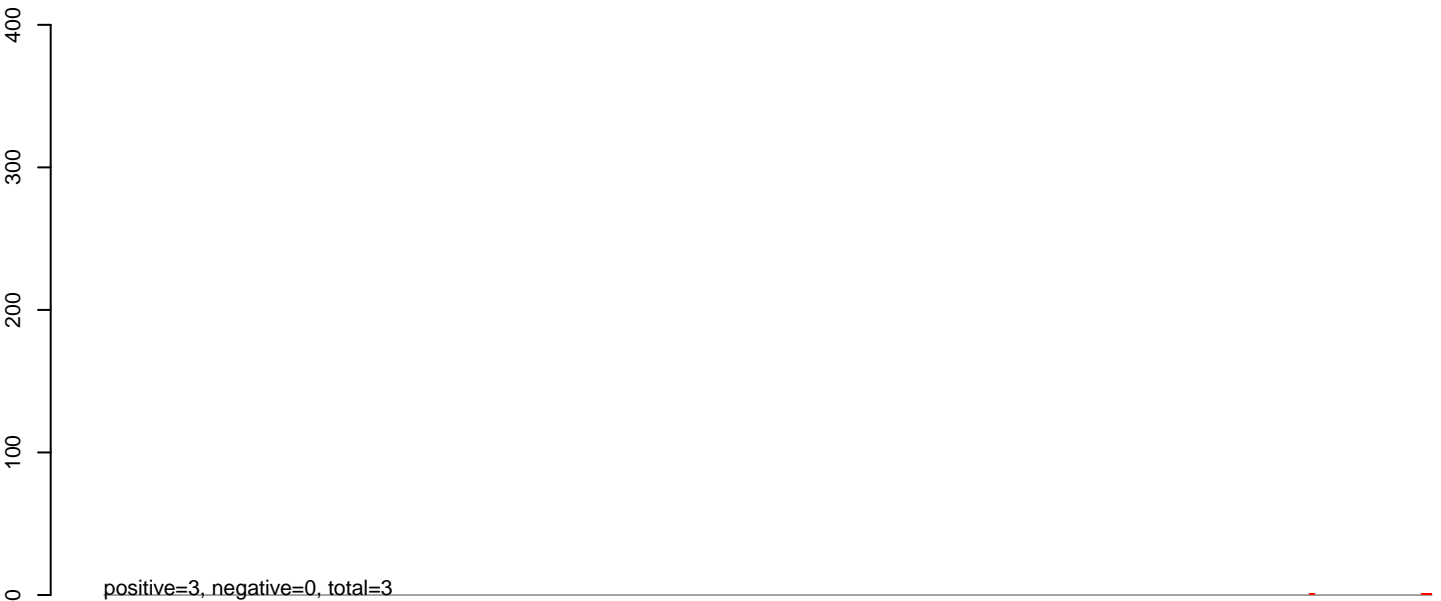


Window size=25, length=11444, virus@NC\_001449.1-Venezuelan\_equine\_encephalitis\_virus:1-11444

0 2000 4000 6000 8000 10000



AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep

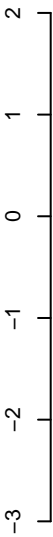


AnGam\_Mos55\_JR.rep



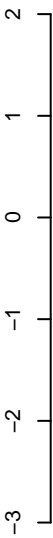
Window size=25, length=6203, virus@NC\_035674.1-Australian\_Anopheles\_totivirus\_isolate\_AATV\_150840:1-6203

AnGam\_Mos55\_JR.18\_23.rep



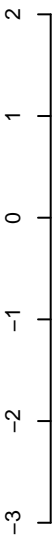
positive=5, negative=7, total=12

AnGam\_Mos55\_JR.24\_35.rep

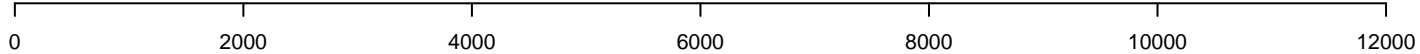


positive=0, negative=0, total=0

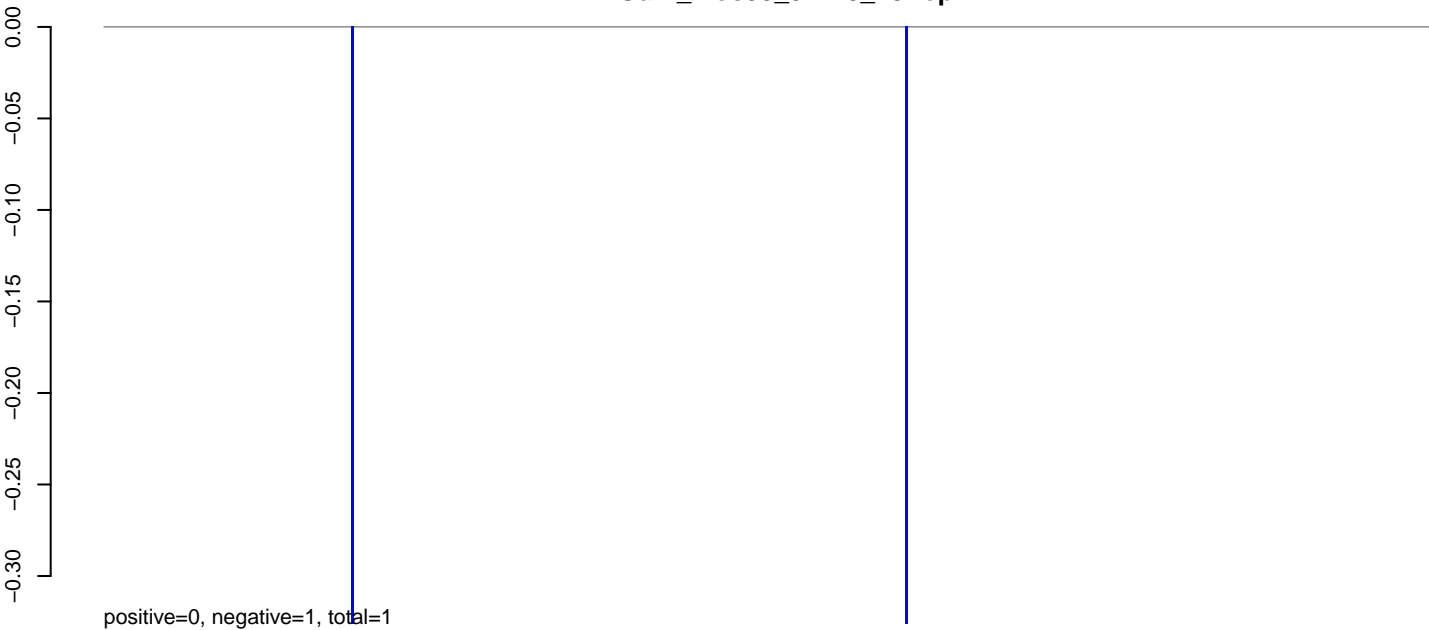
AnGam\_Mos55\_JR.rep



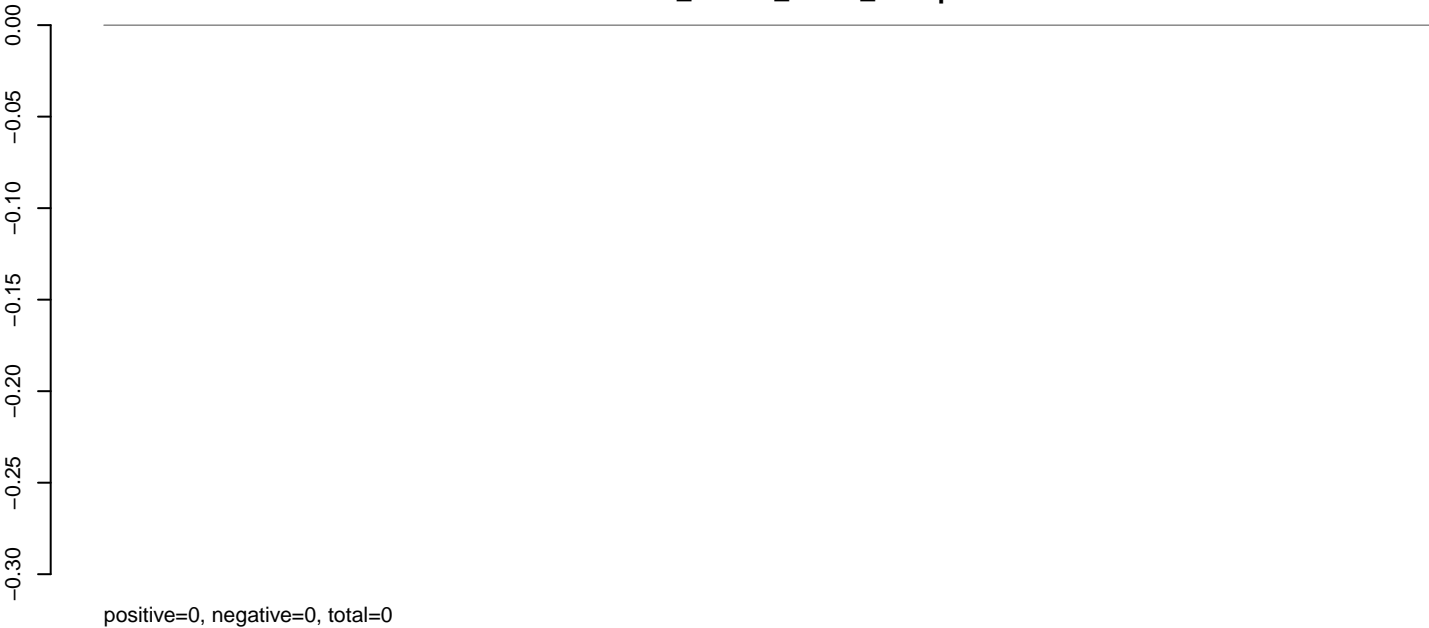
positive=5, negative=25, total=30, length=11624, virus@NC\_023812.1-Madariaga\_virus\_strain\_MADV\_Cebus\_apella\_BRA\_BEAN5122\_1956:1-11624



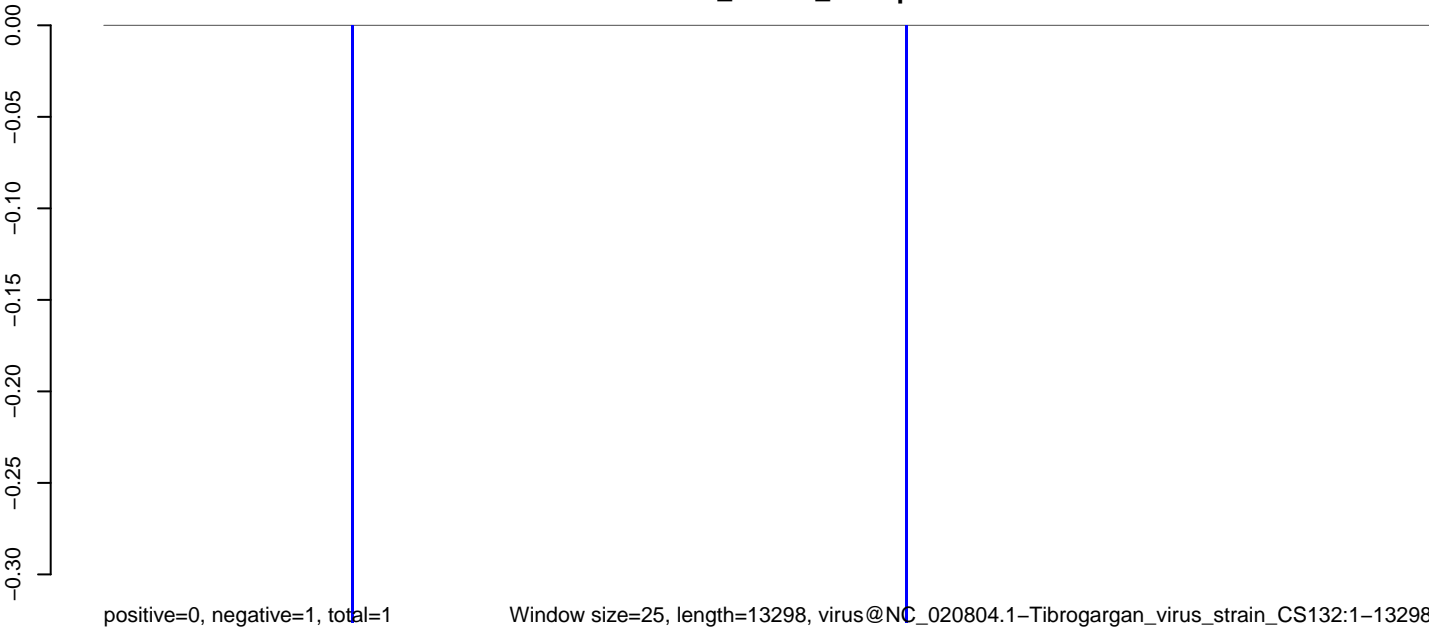
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



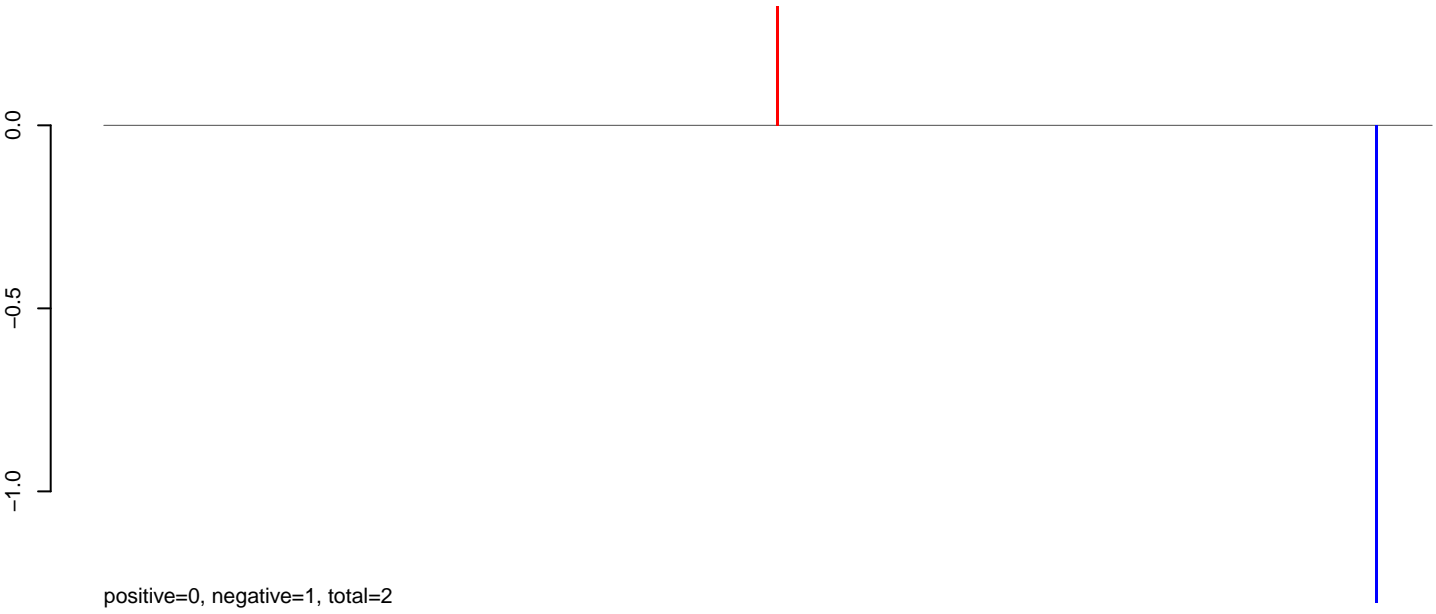
AnGam\_Mos55\_JR.rep



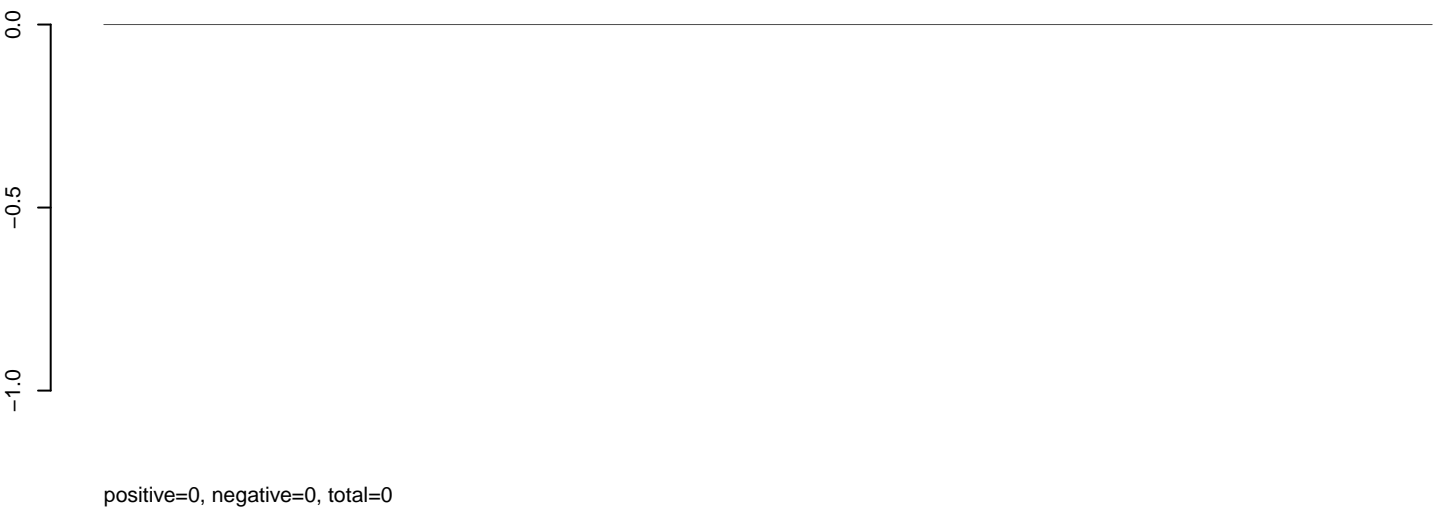
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0 2000 4000 6000 8000 10000 12000

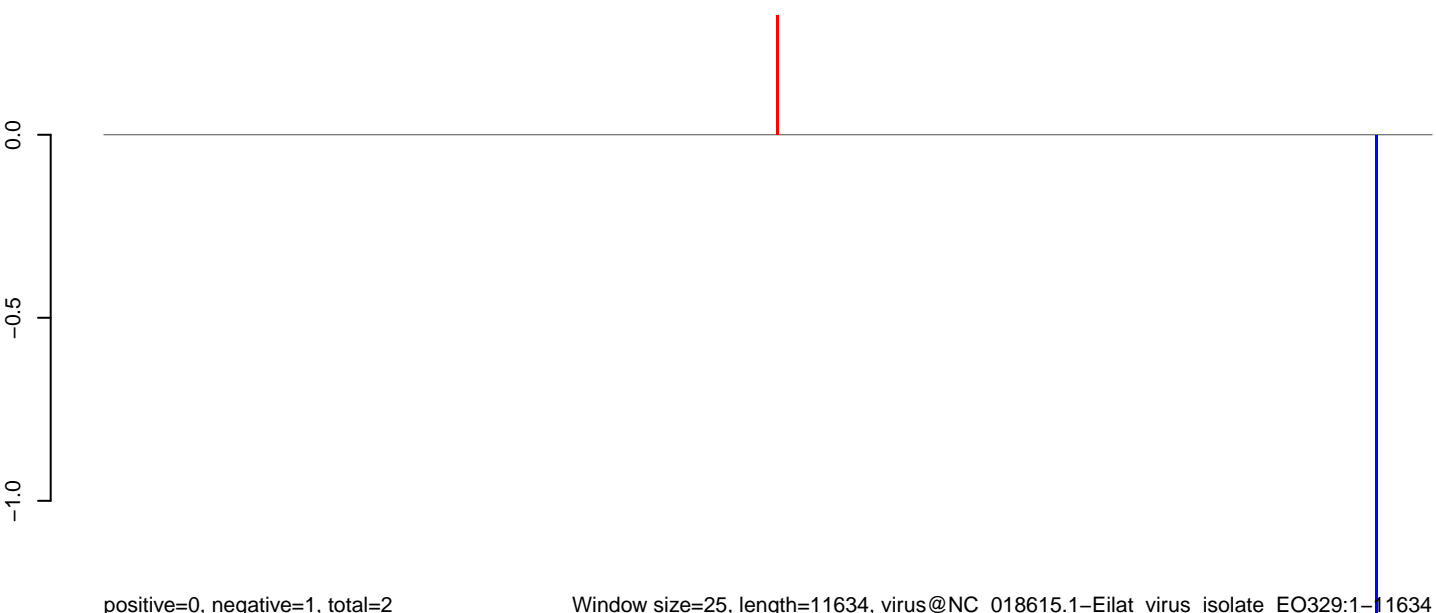
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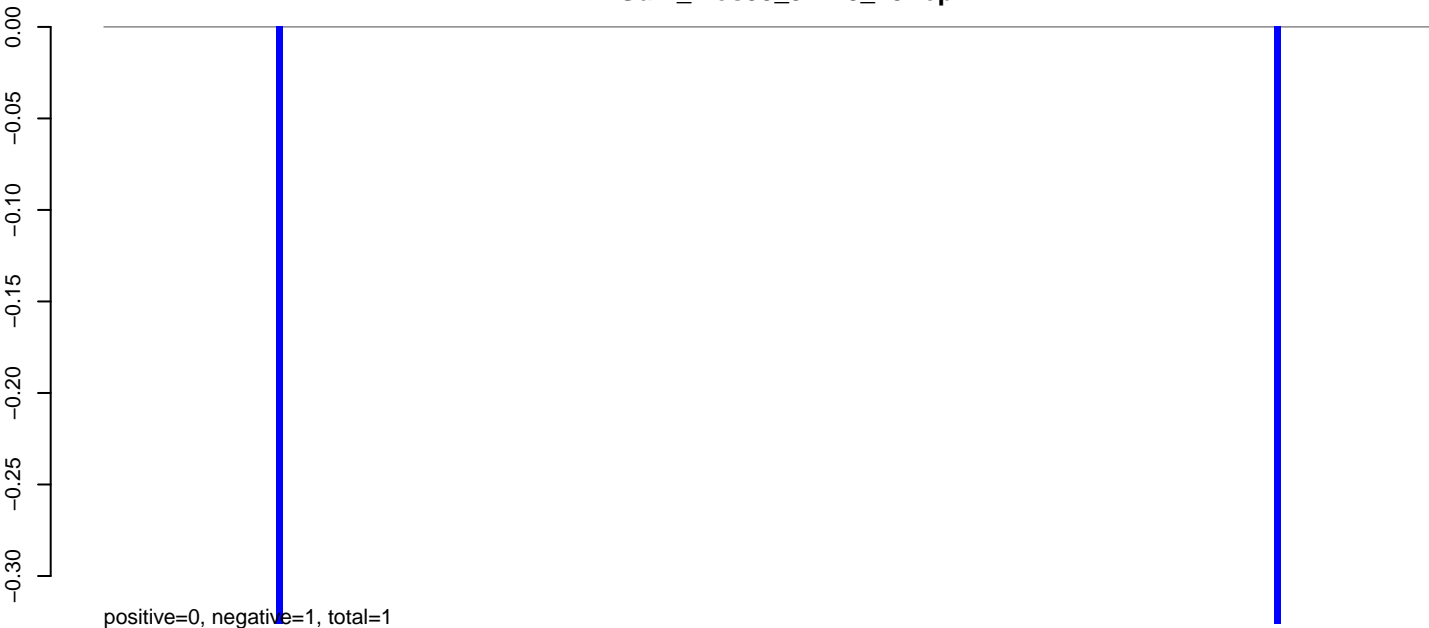
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AnGam\_Mos55\_JR.rep

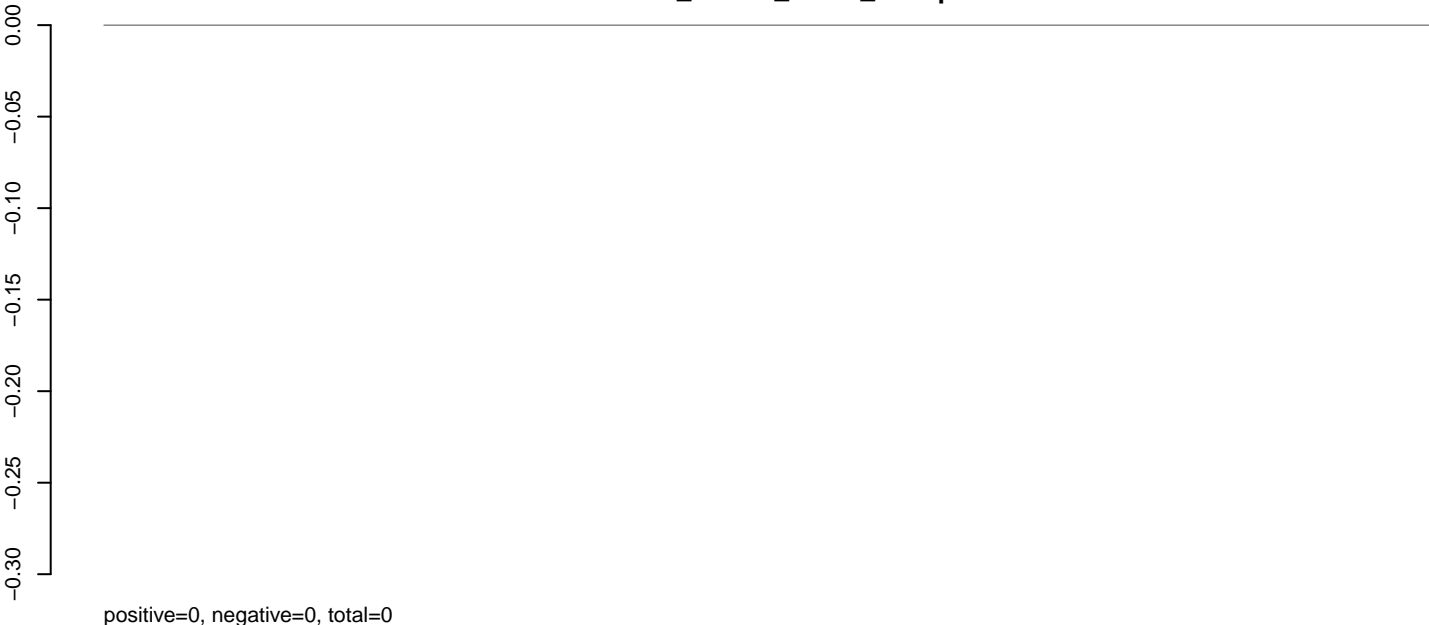


AnGam\_Mos55\_JR.18\_23.rep



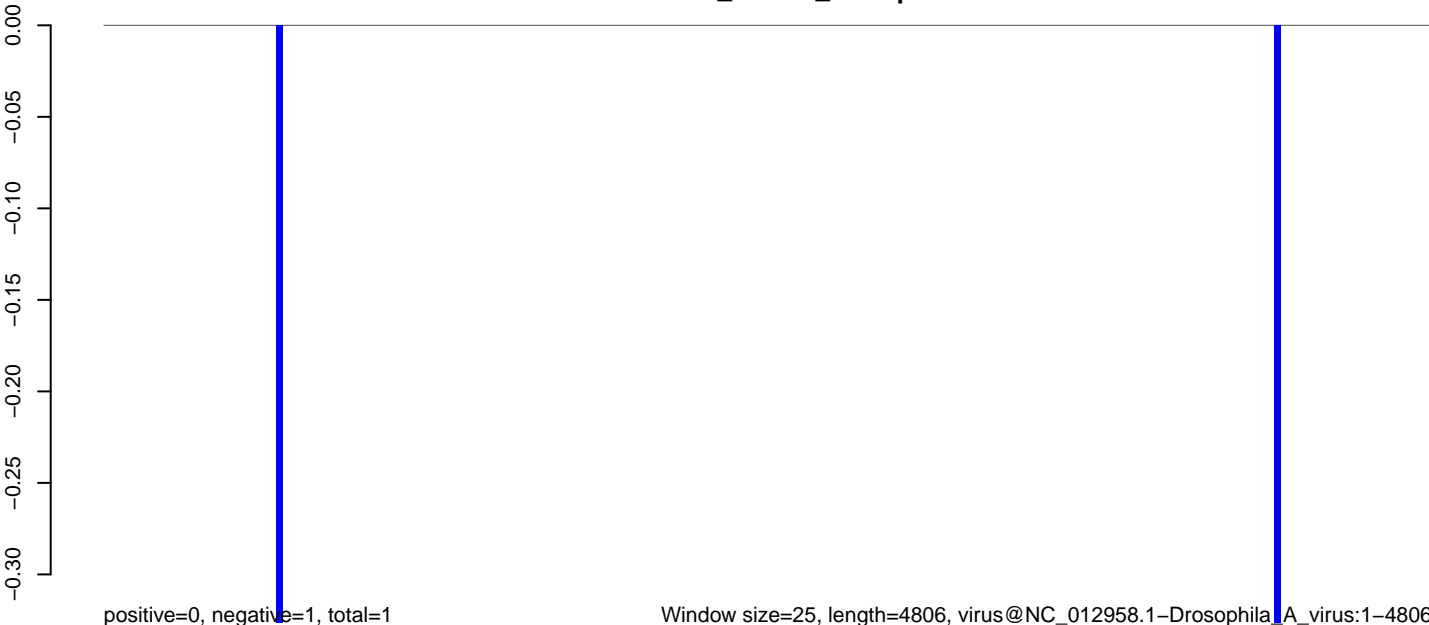
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AnGam\_Mos55\_JR.24\_35.rep



positive=0, negative=0, total=0

AnGam\_Mos55\_JR.rep

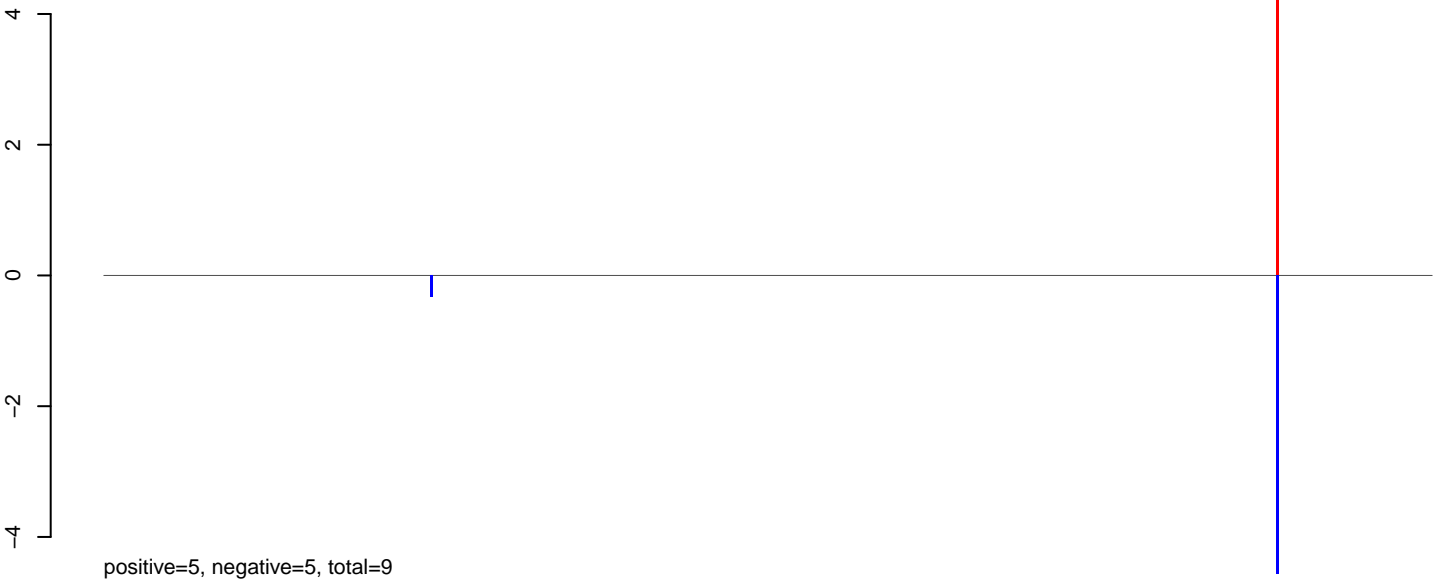


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Window size=25, length=4806, virus@NC\_012958.1-Drosophila\_A\_virus:1-4806

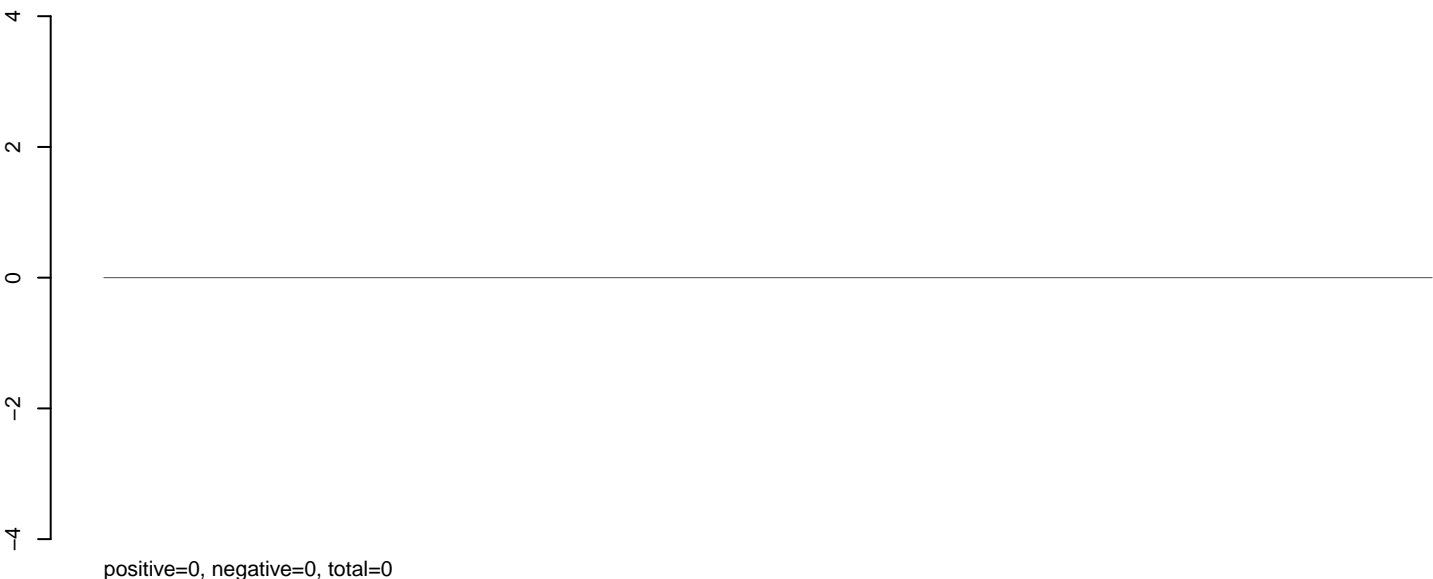
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AnGam\_Mos55\_JR.18\_23.rep



positive=5, negative=5, total=9

AnGam\_Mos55\_JR.24\_35.rep



positive=0, negative=0, total=0

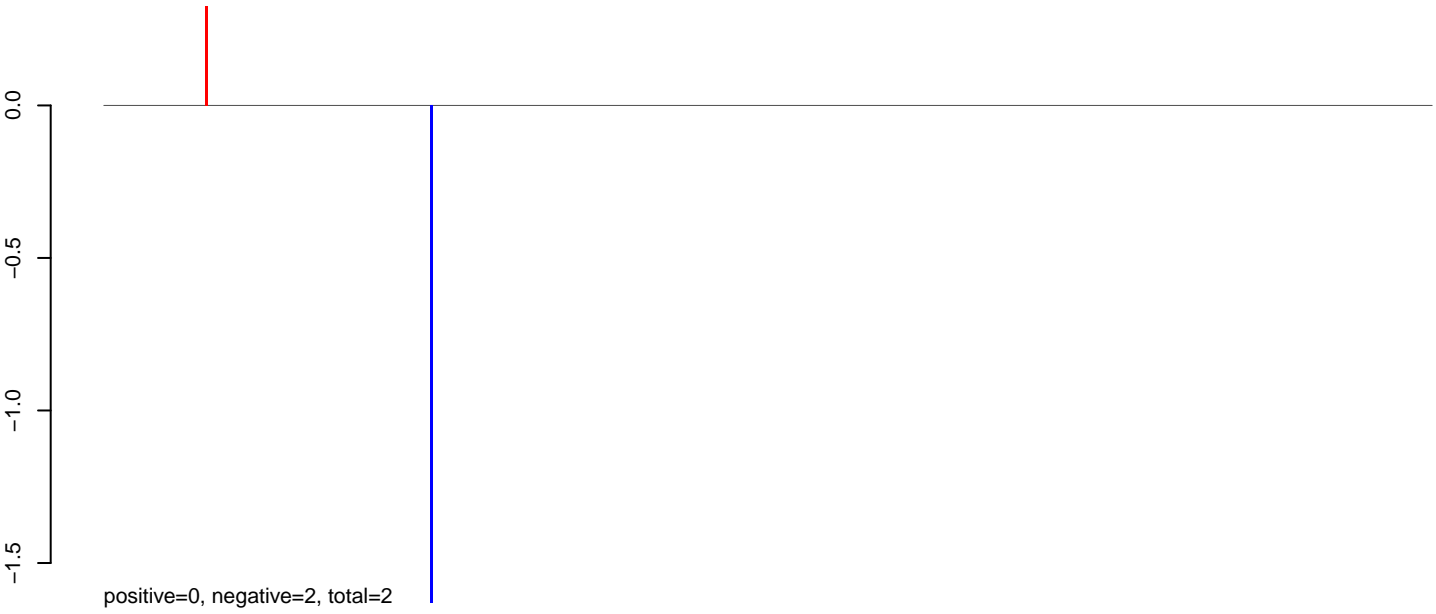
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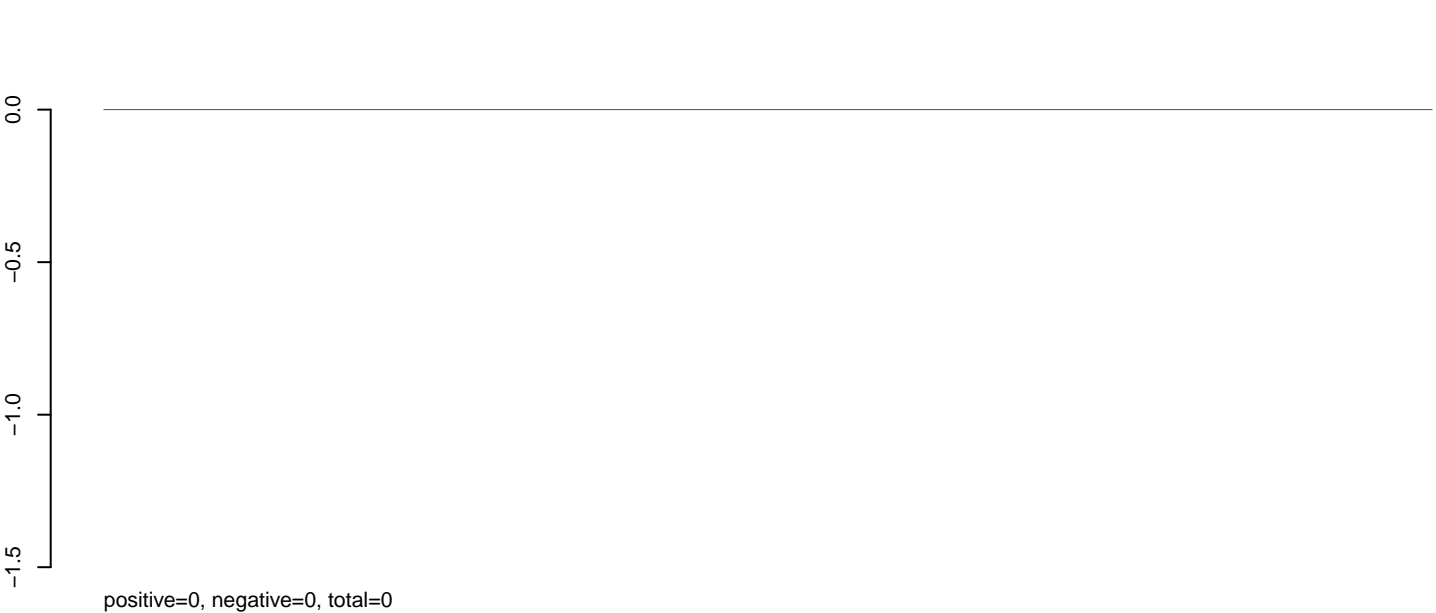
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0 2000 4000 6000 8000 10000

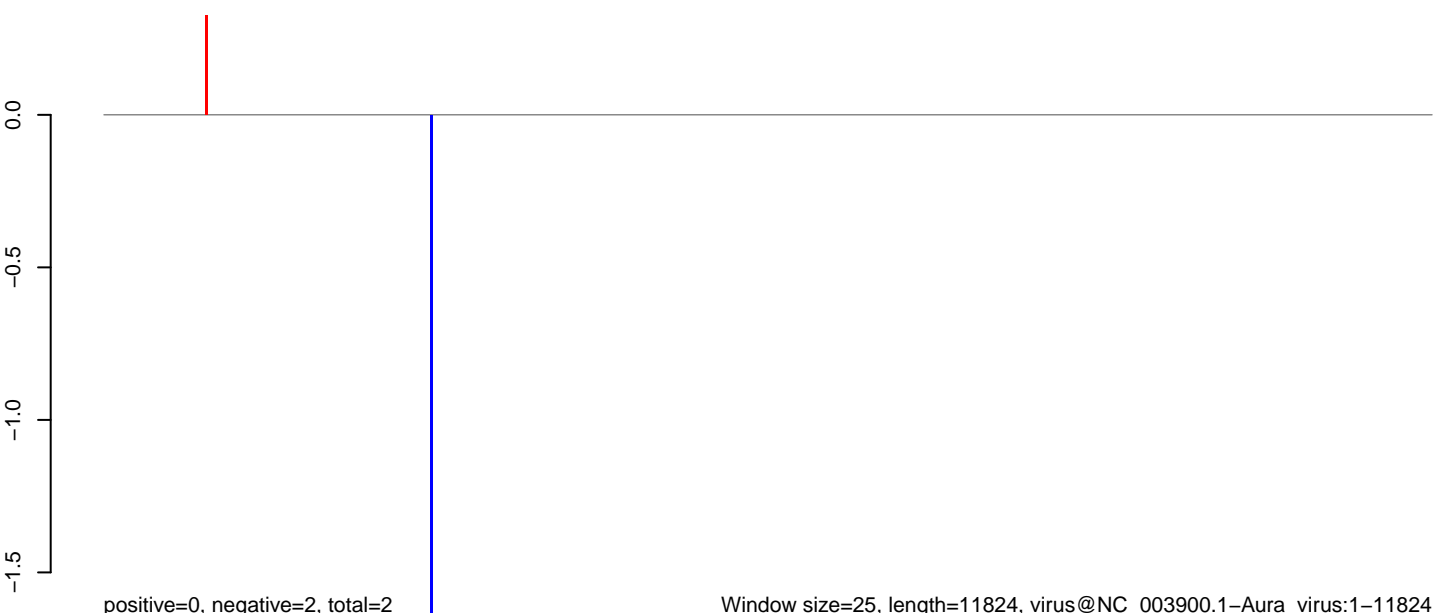
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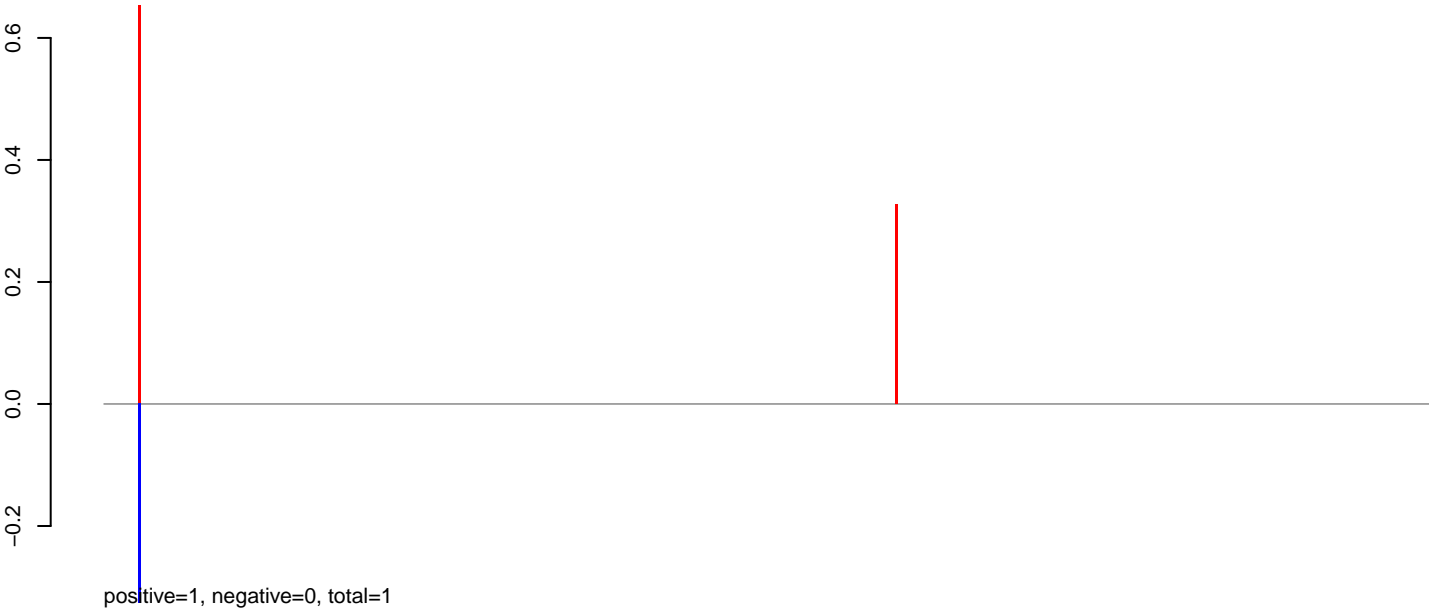
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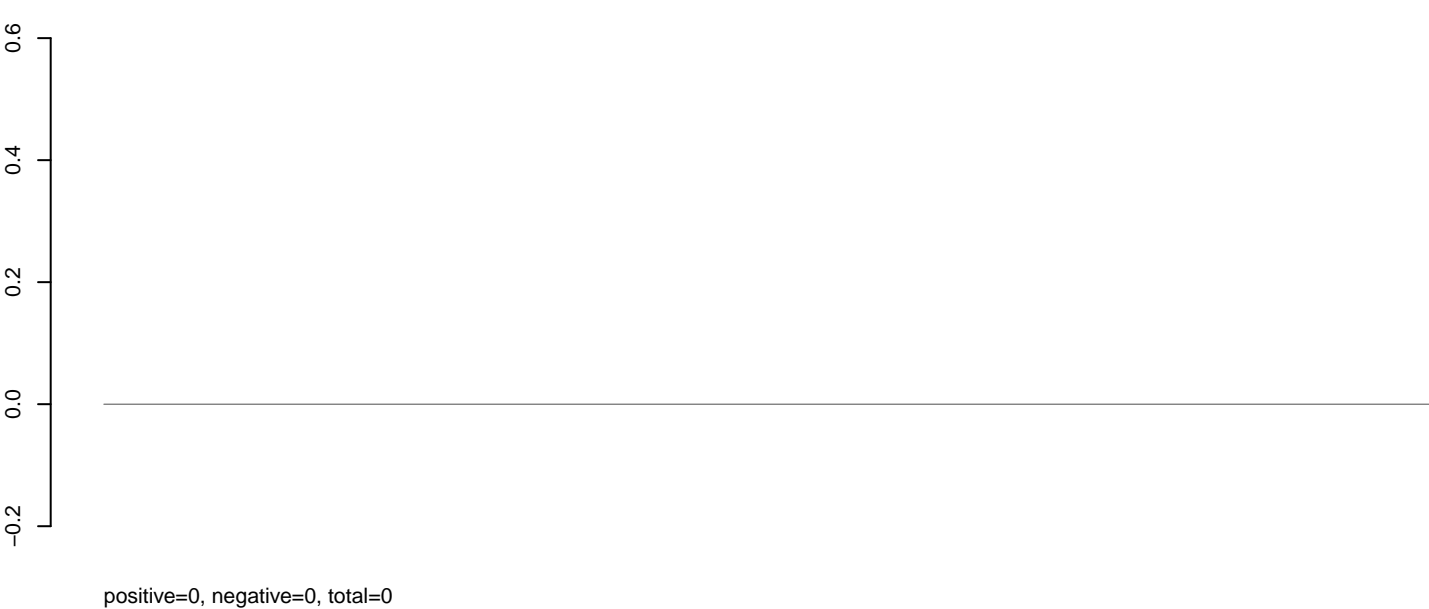
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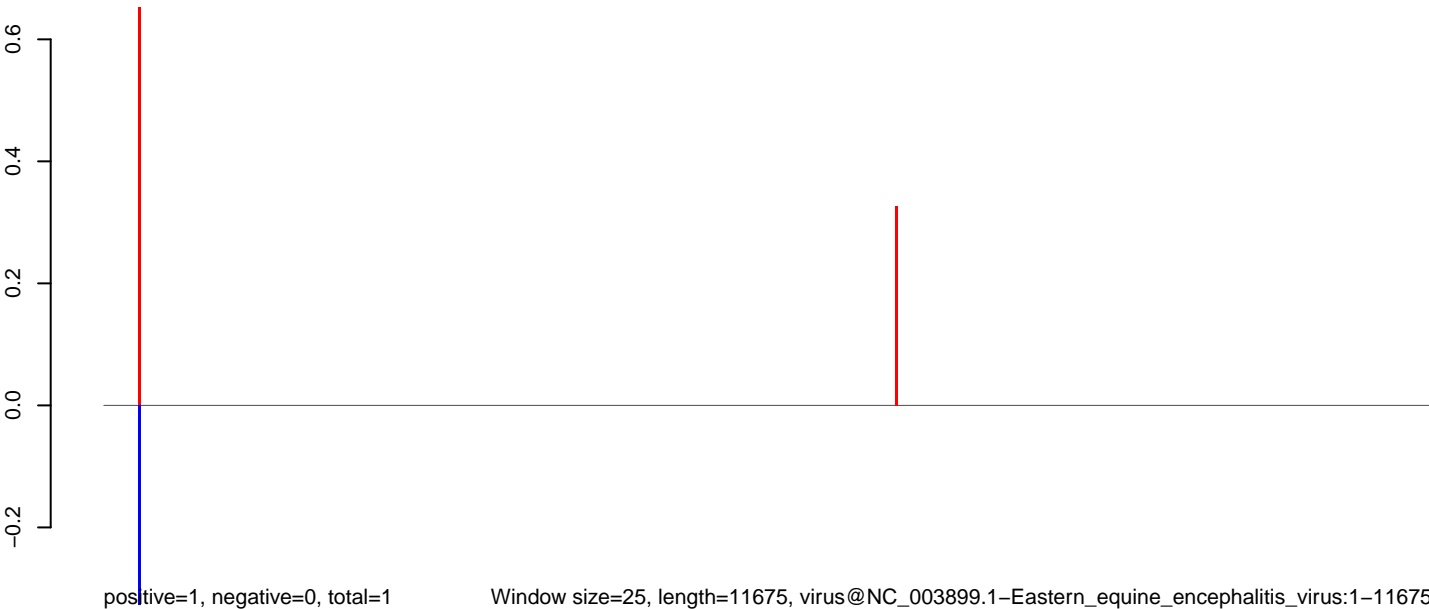
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AnGam\_Mos55\_JR.24\_35.rep



AnGam\_Mos55\_JR.rep

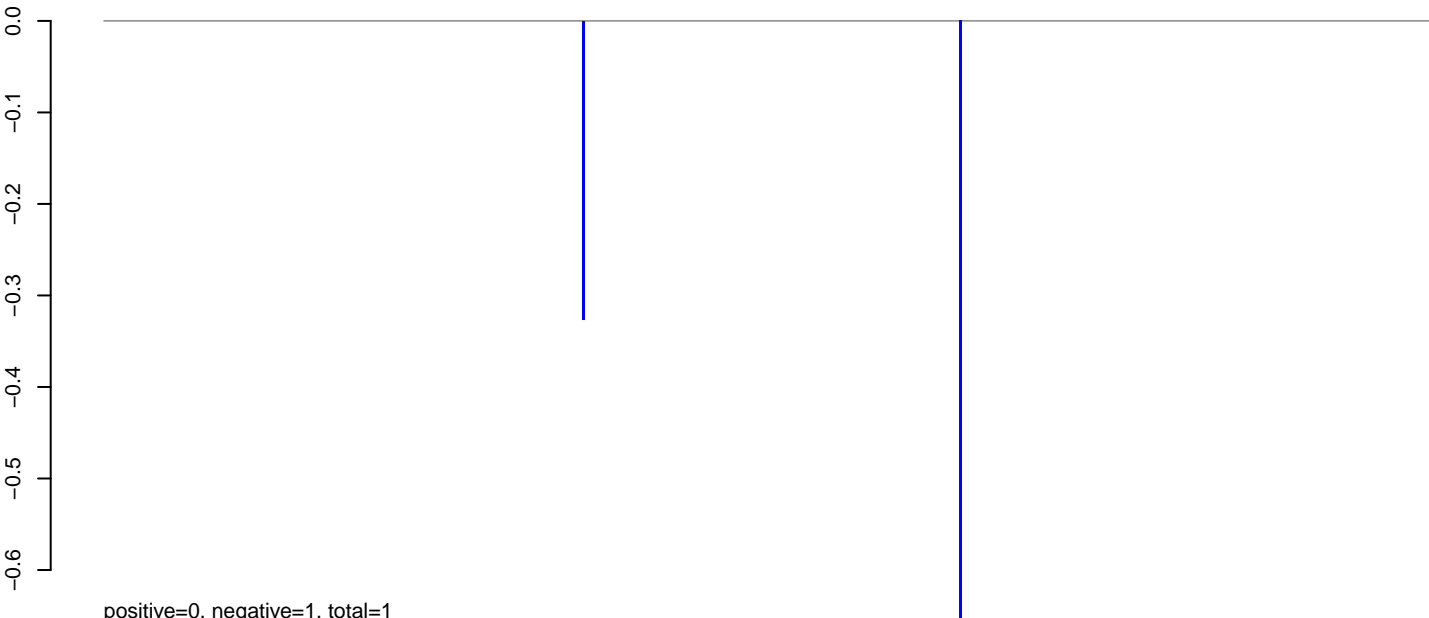


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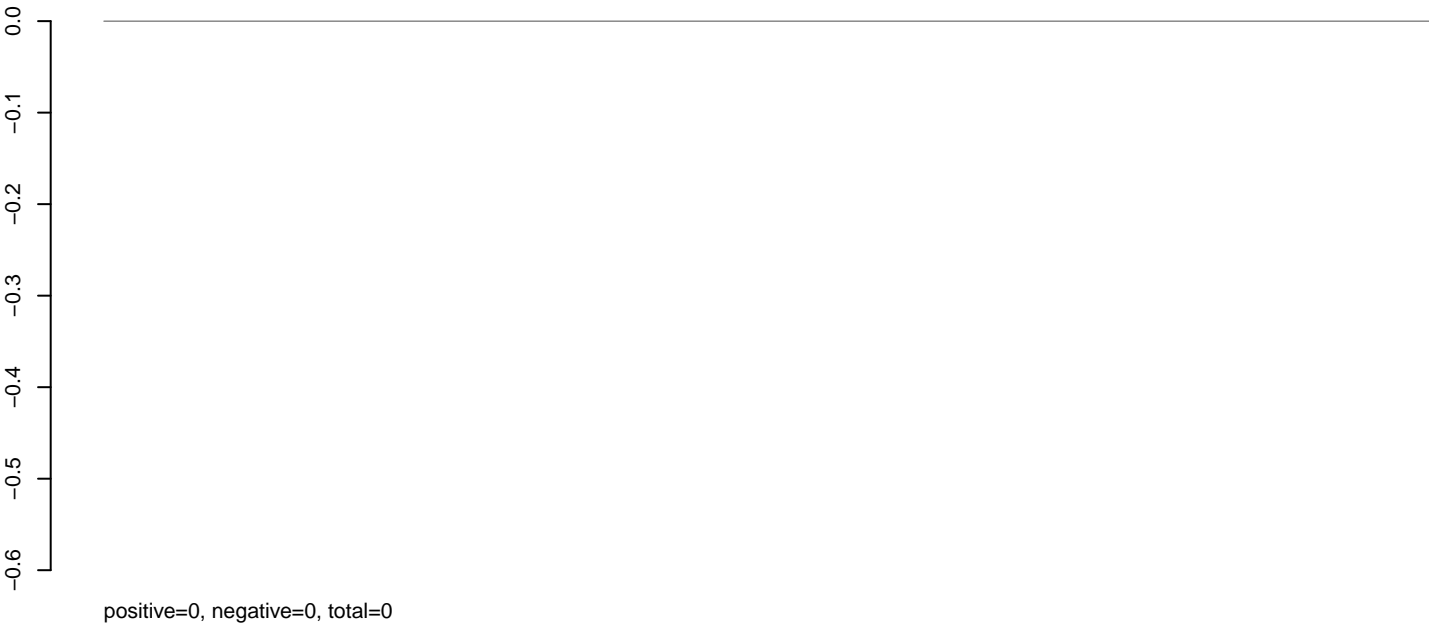
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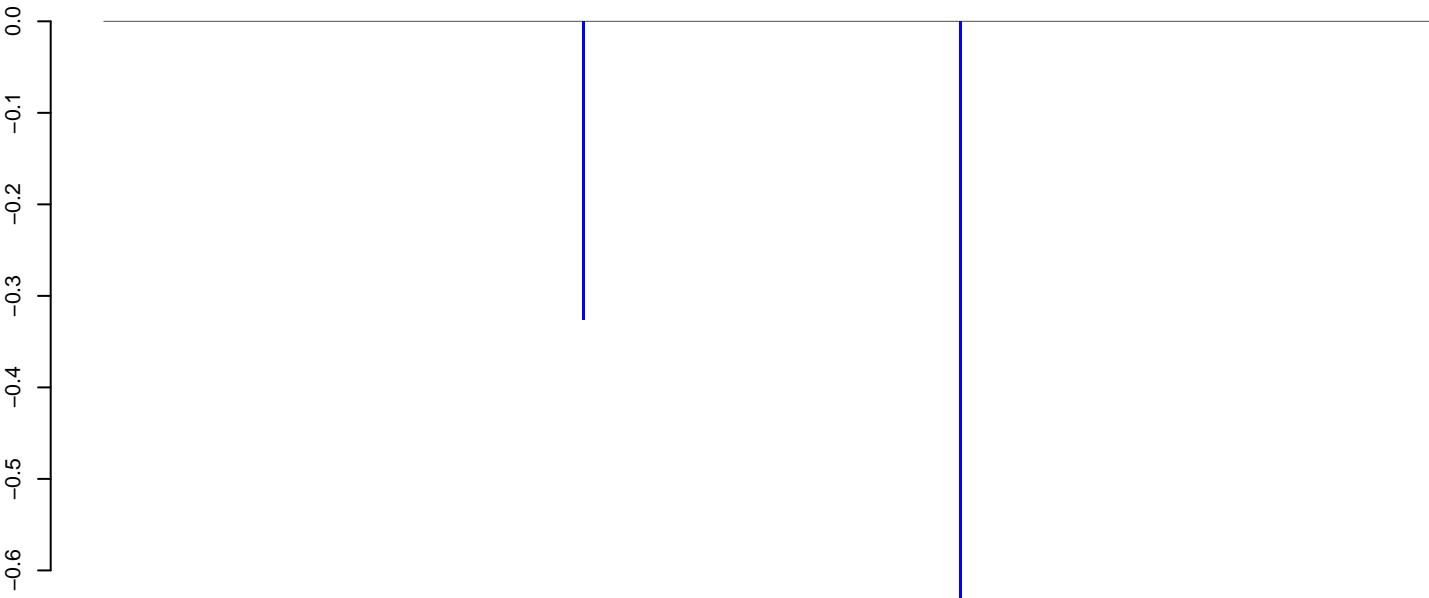
AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



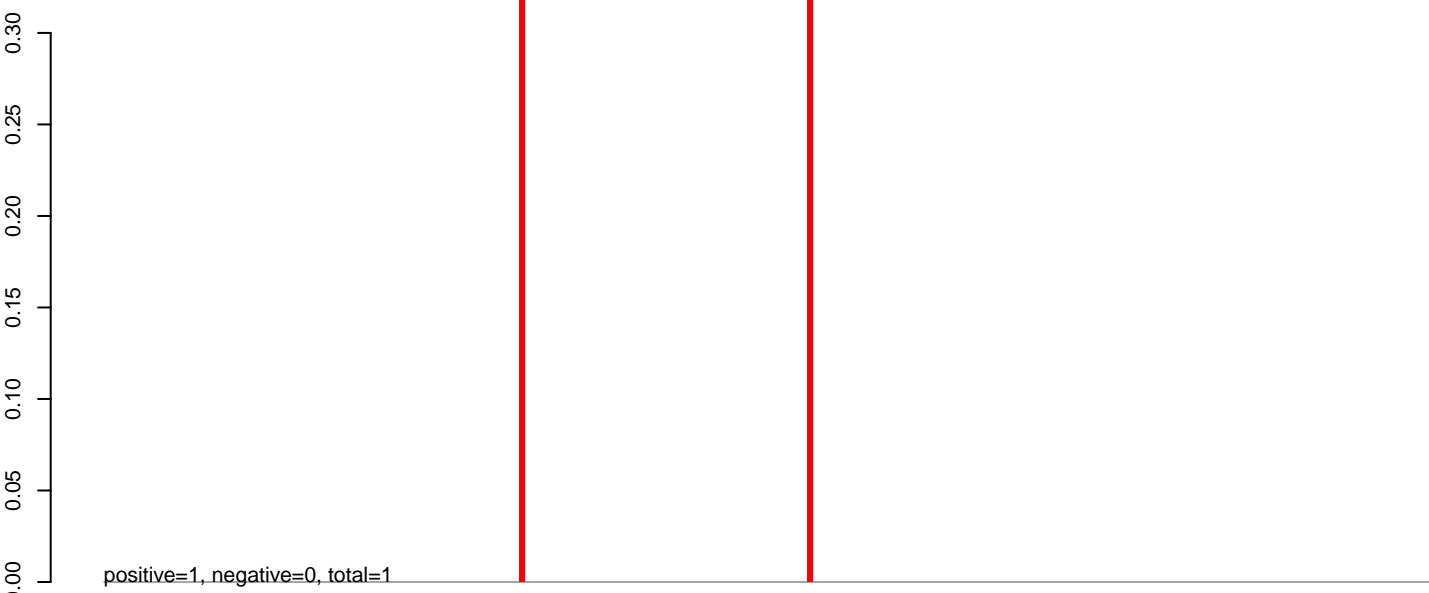
AnGam\_Mos55\_JR.rep



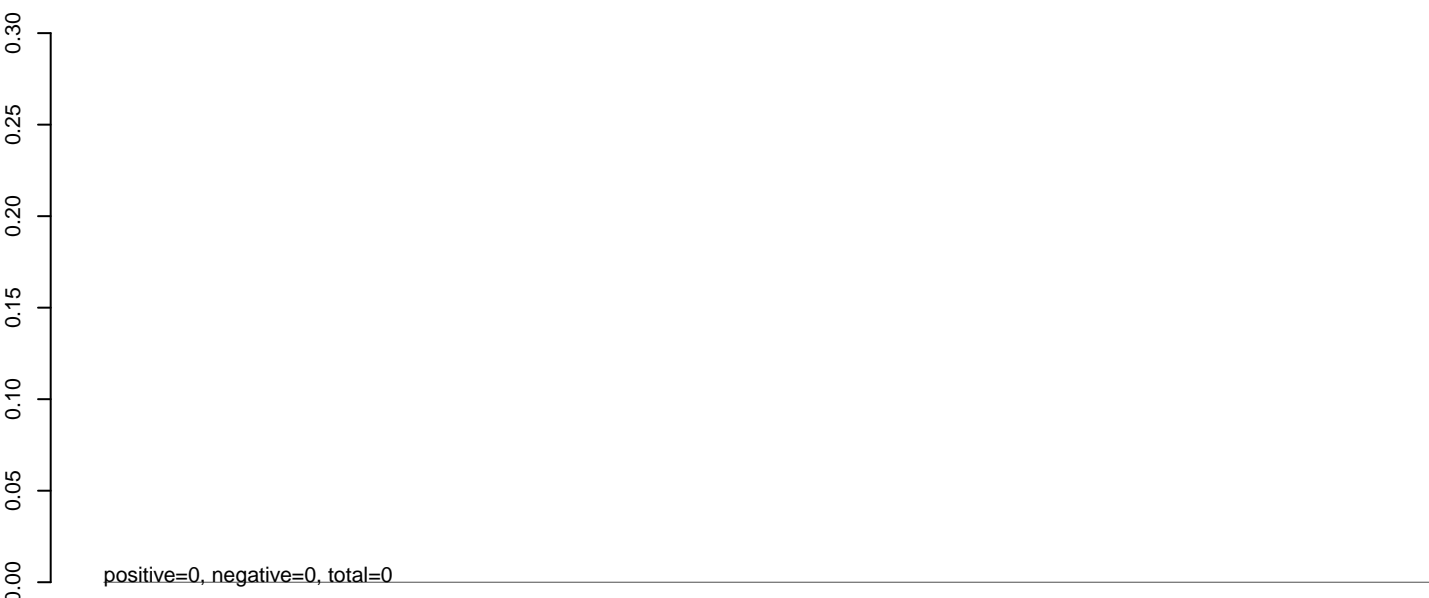
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0 2000 4000 6000 8000 10000 12000

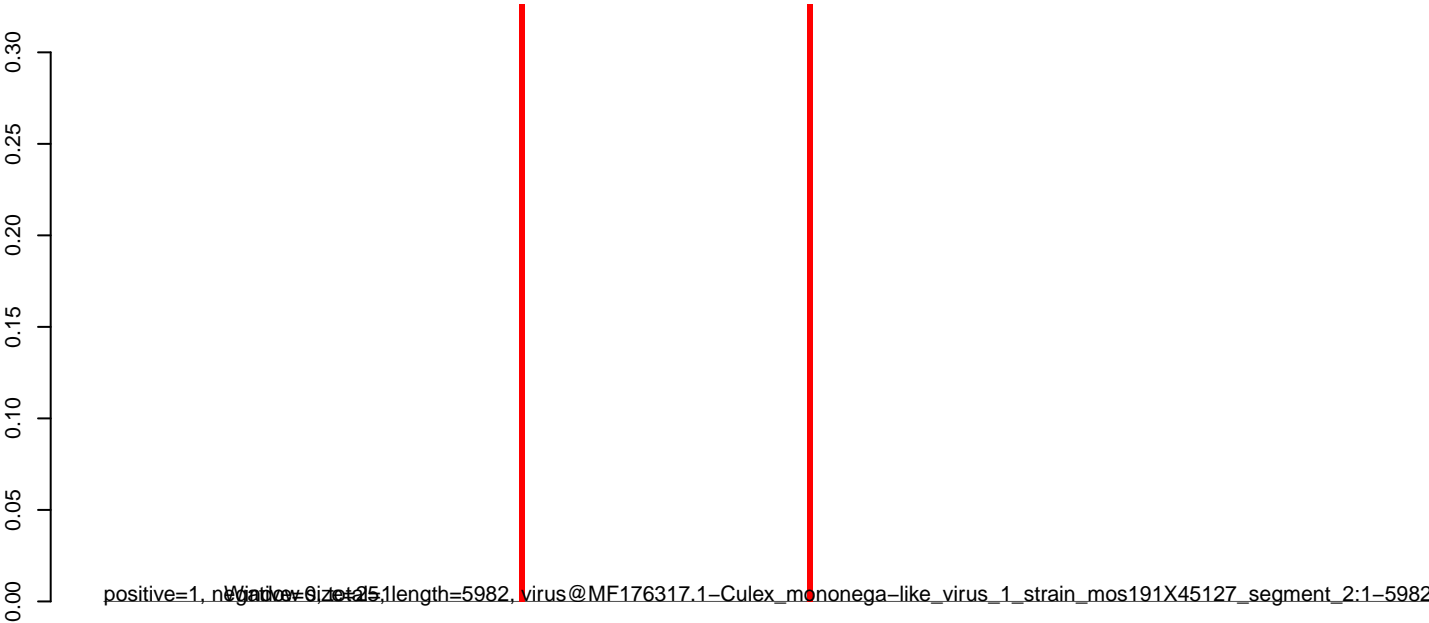
### AnGam\_Mos55\_JR.18\_23.rep



### AnGam\_Mos55\_JR.24\_35.rep



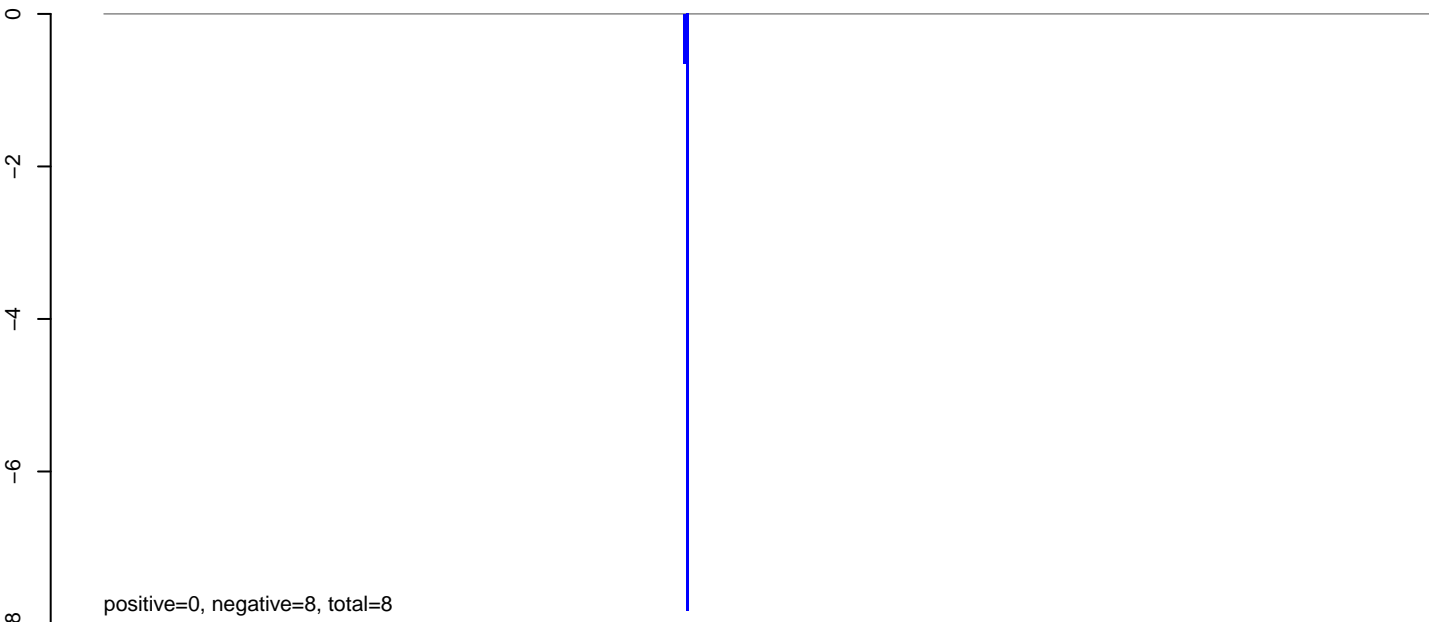
### AnGam\_Mos55\_JR.rep



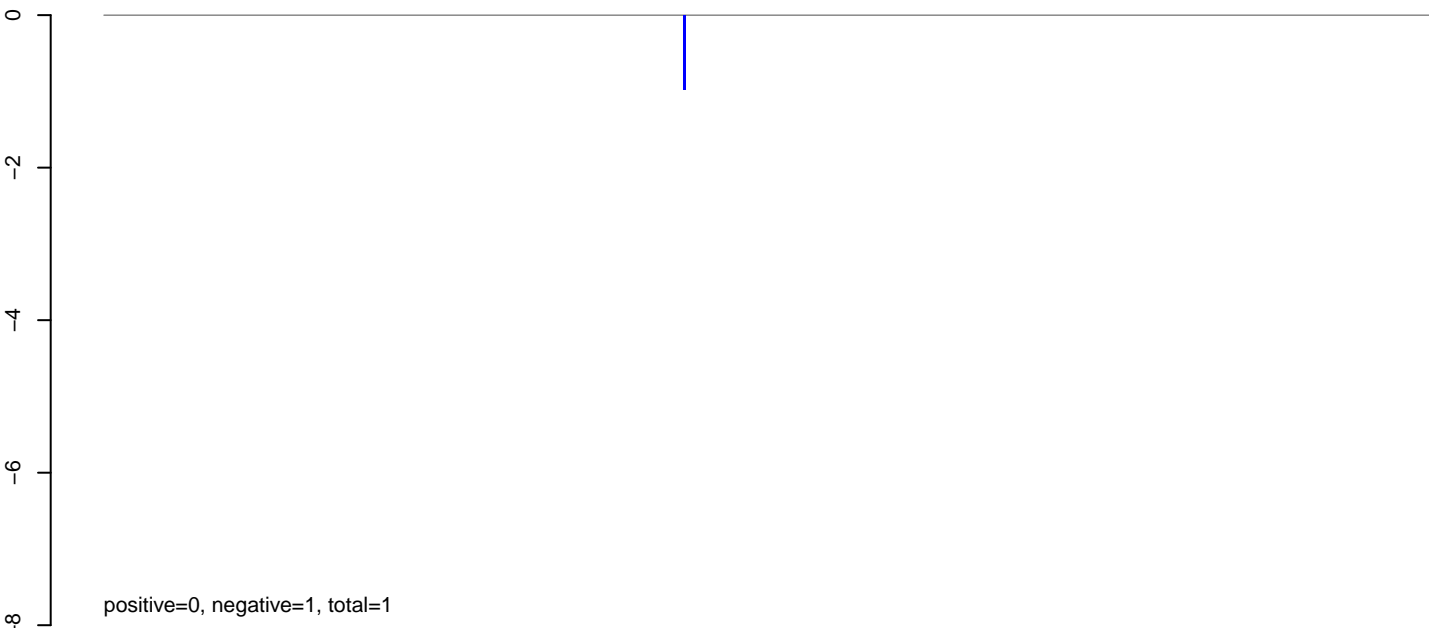
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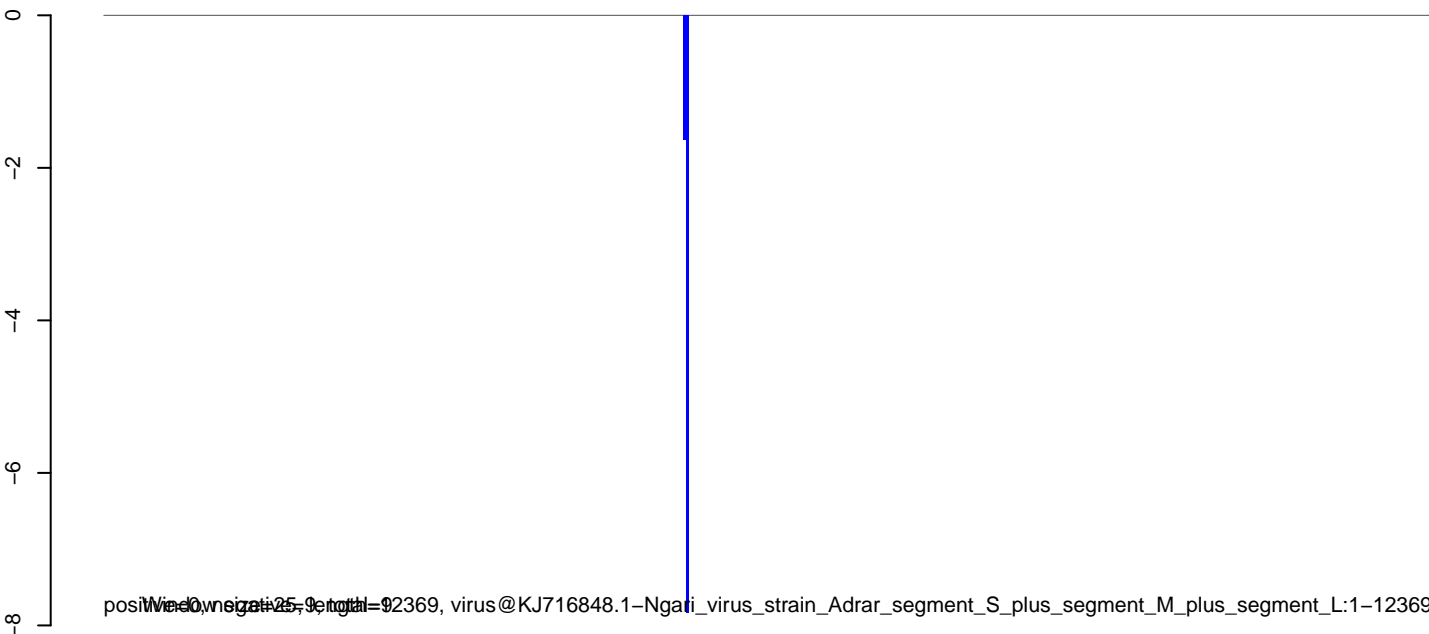
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AnGam\_Mos55\_JR.24\_35.rep

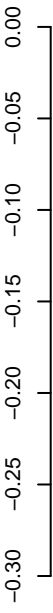


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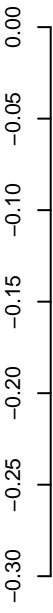
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AnGam\_Mos55\_JR.18\_23.rep



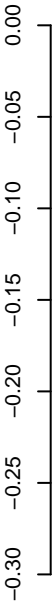
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AnGam\_Mos55\_JR.24\_35.rep



positive=0, negative=0, total=0

AnGam\_Mos55\_JR.rep



positive=0, negative=1, total=1

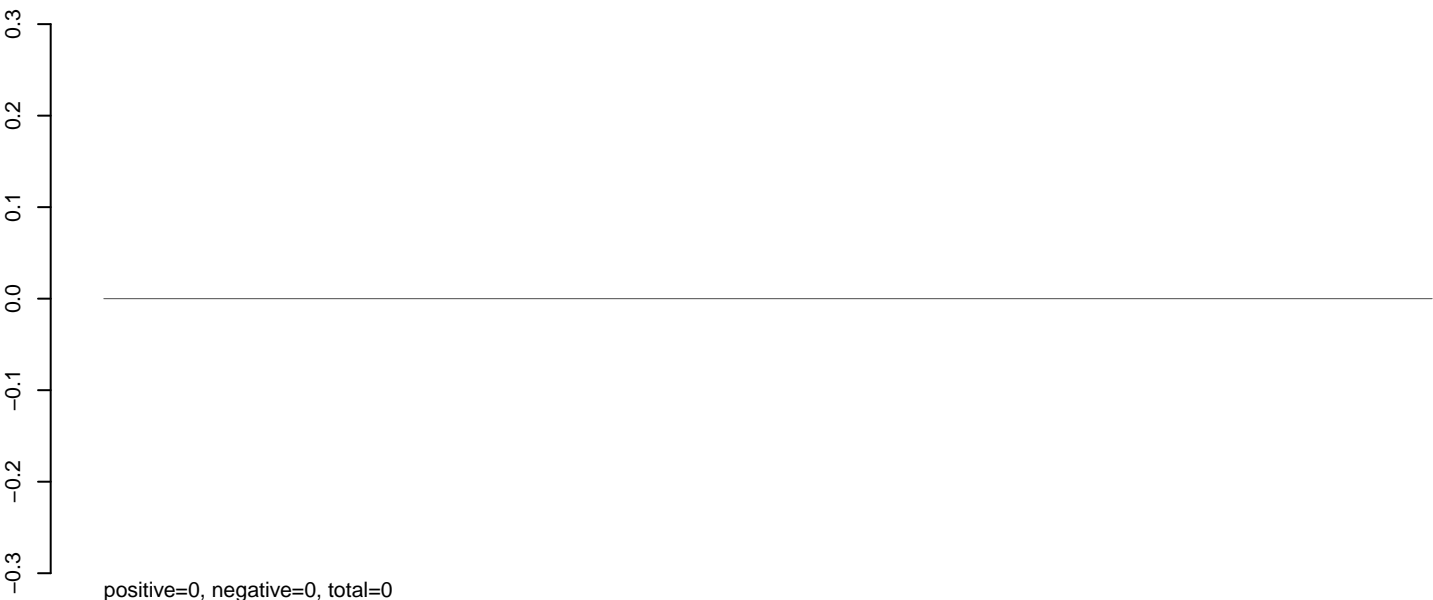
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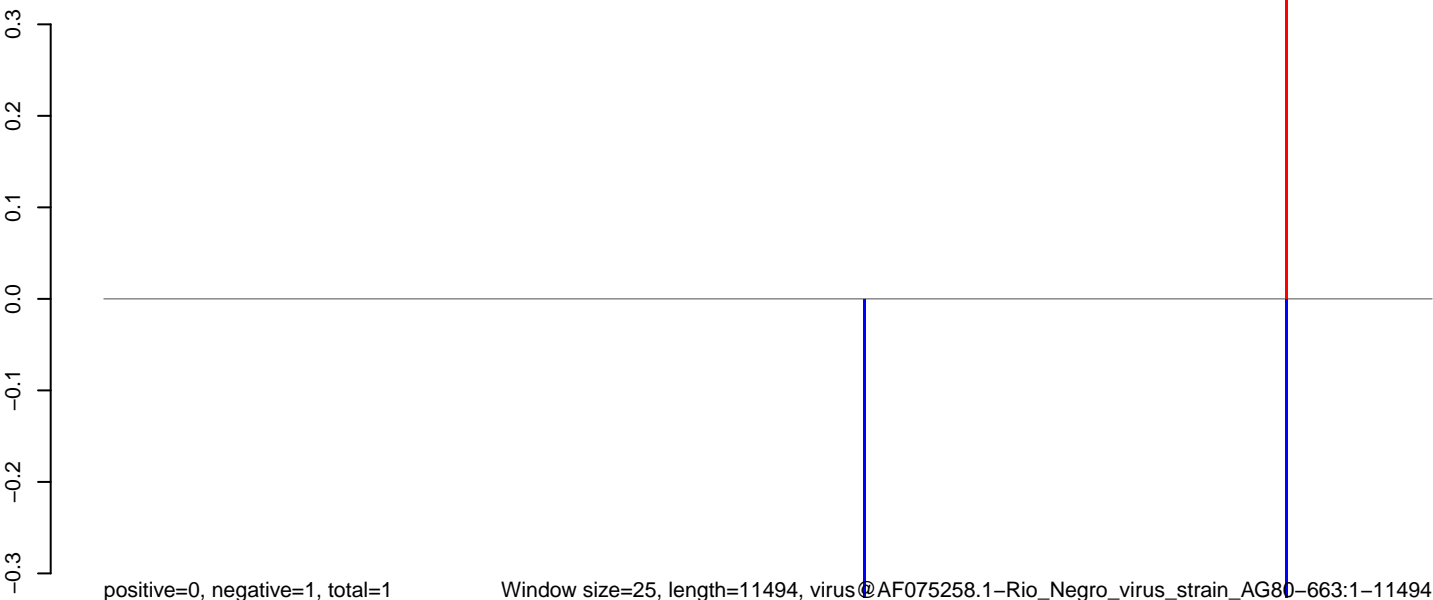
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AnGam\_Mos55\_JR.24\_35.rep



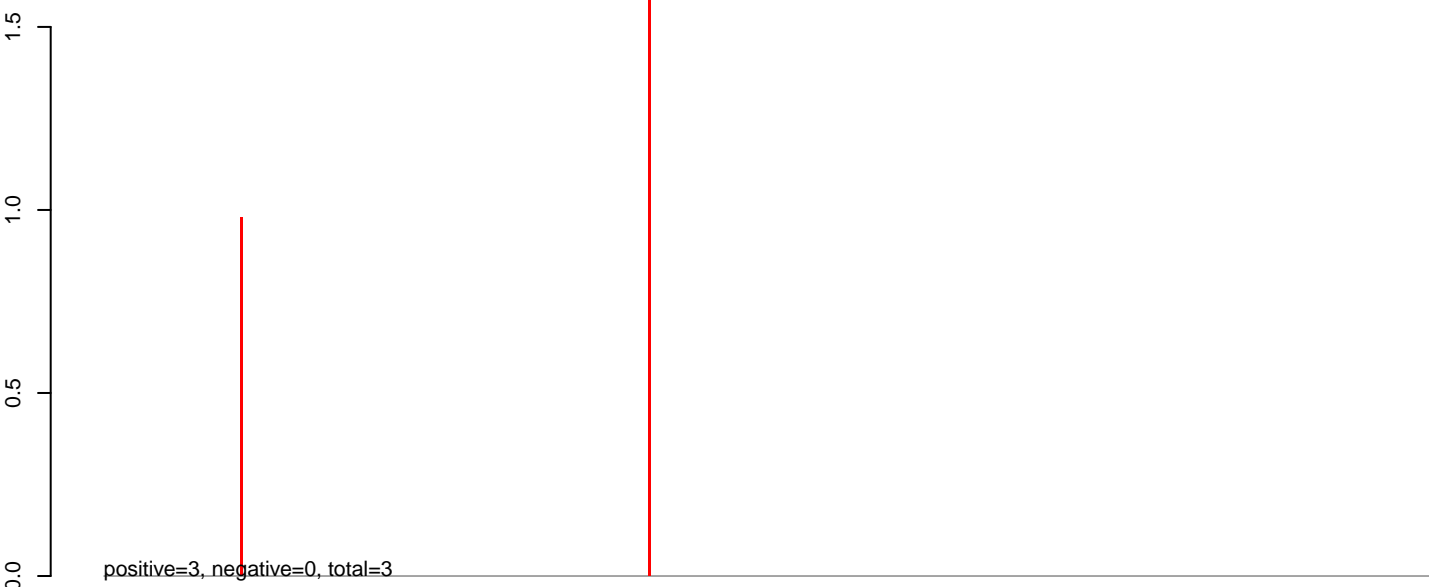
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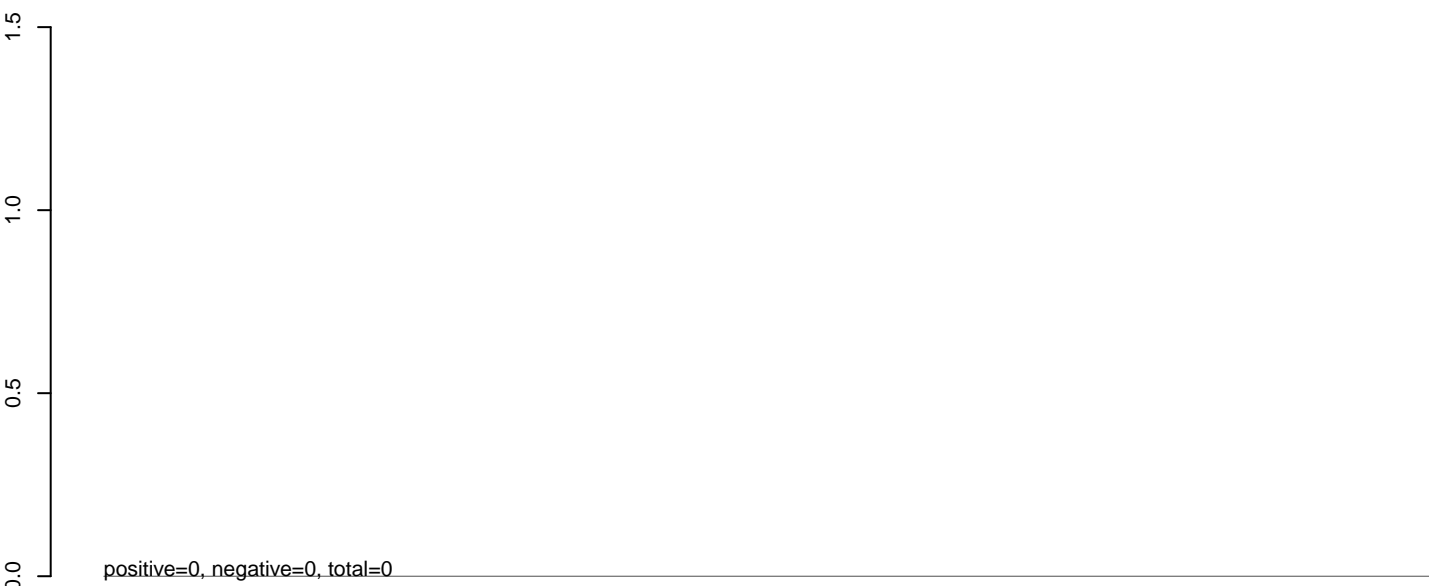
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AnGam\_Mos55\_JR.18\_23.rep



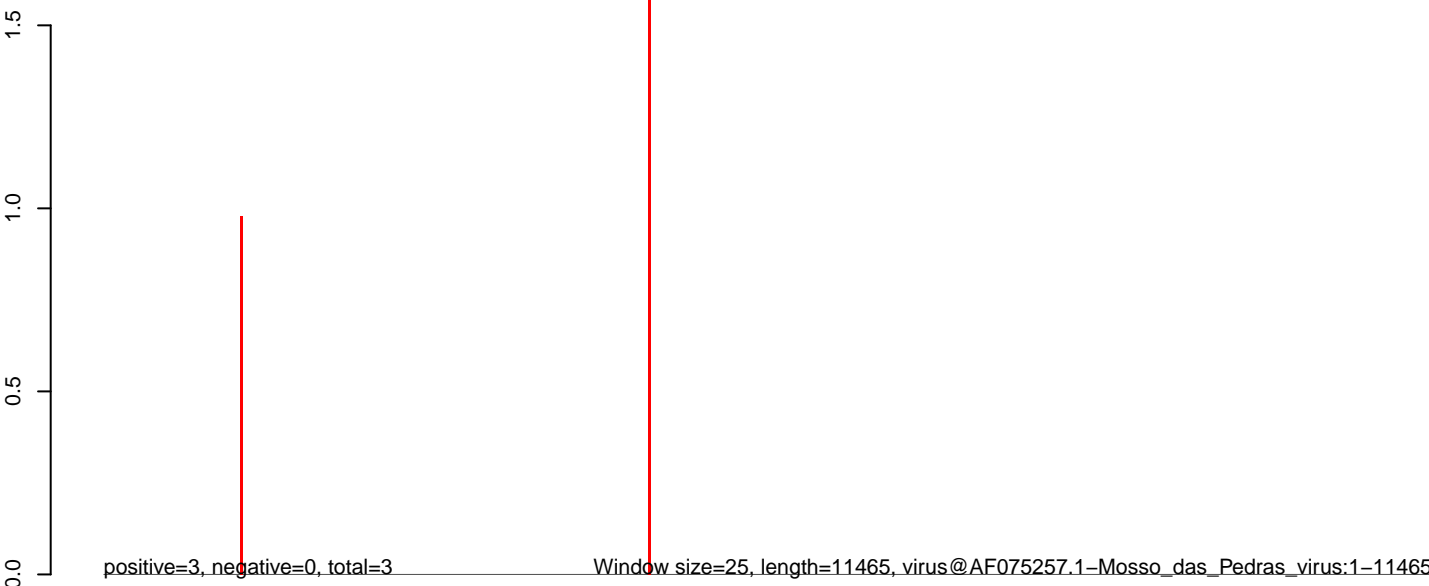
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AnGam\_Mos55\_JR.24\_35.rep



positive=0, negative=0, total=0

AnGam\_Mos55\_JR.rep

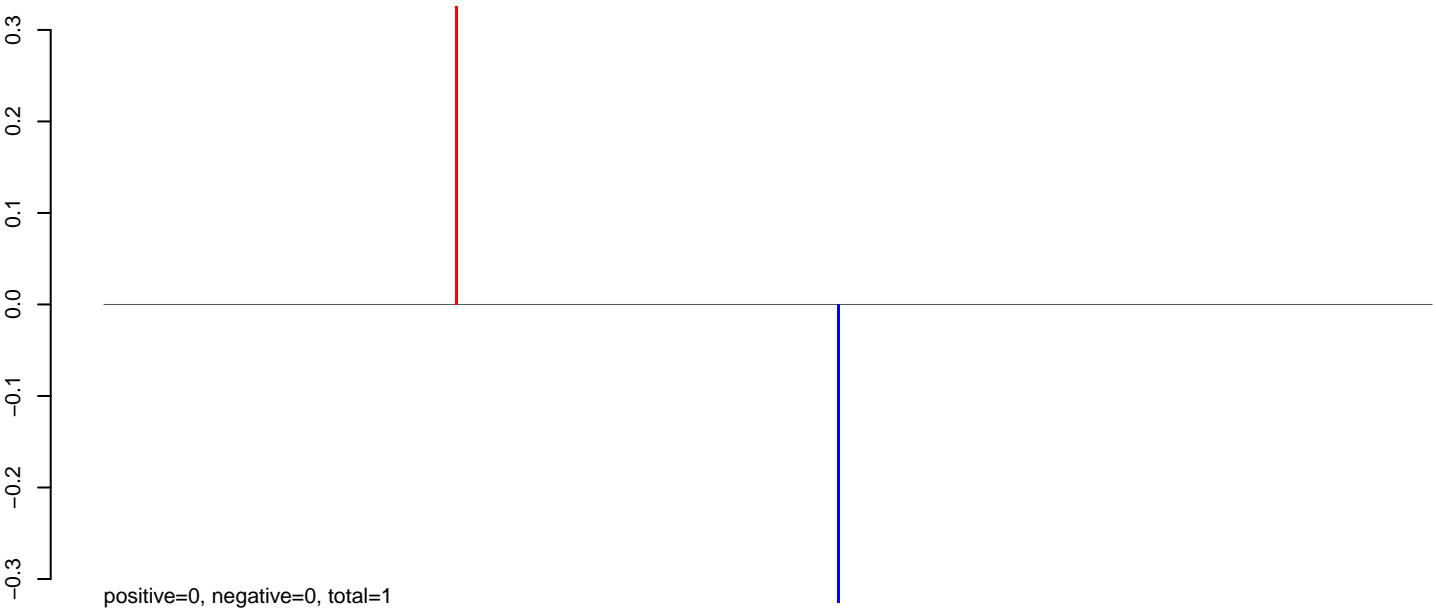


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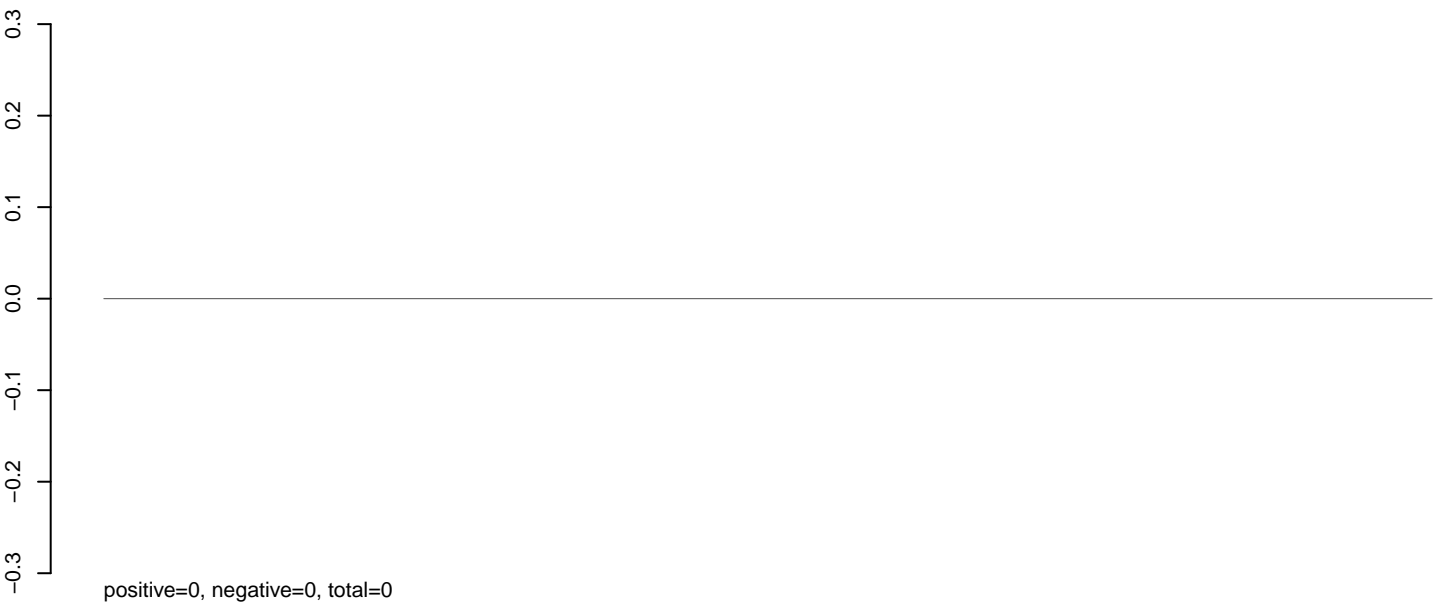
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0 2000 4000 6000 8000 10000

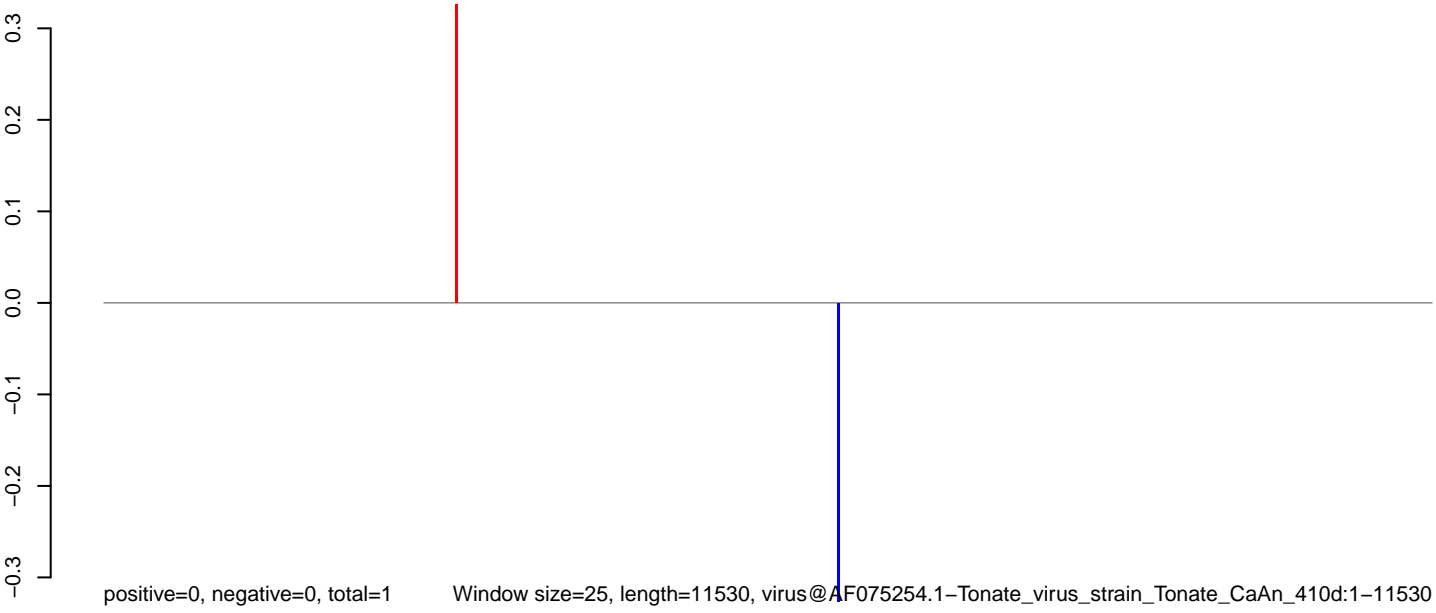
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AnGam\_Mos55\_JR.24\_35.rep

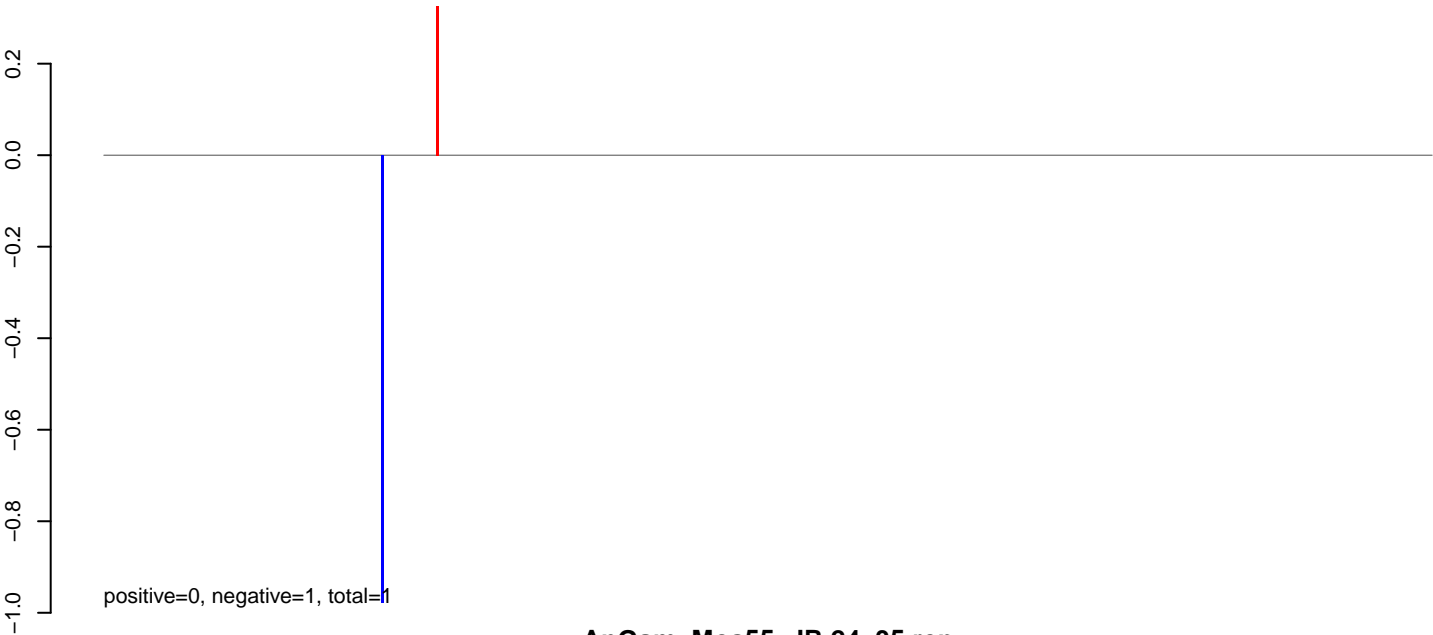


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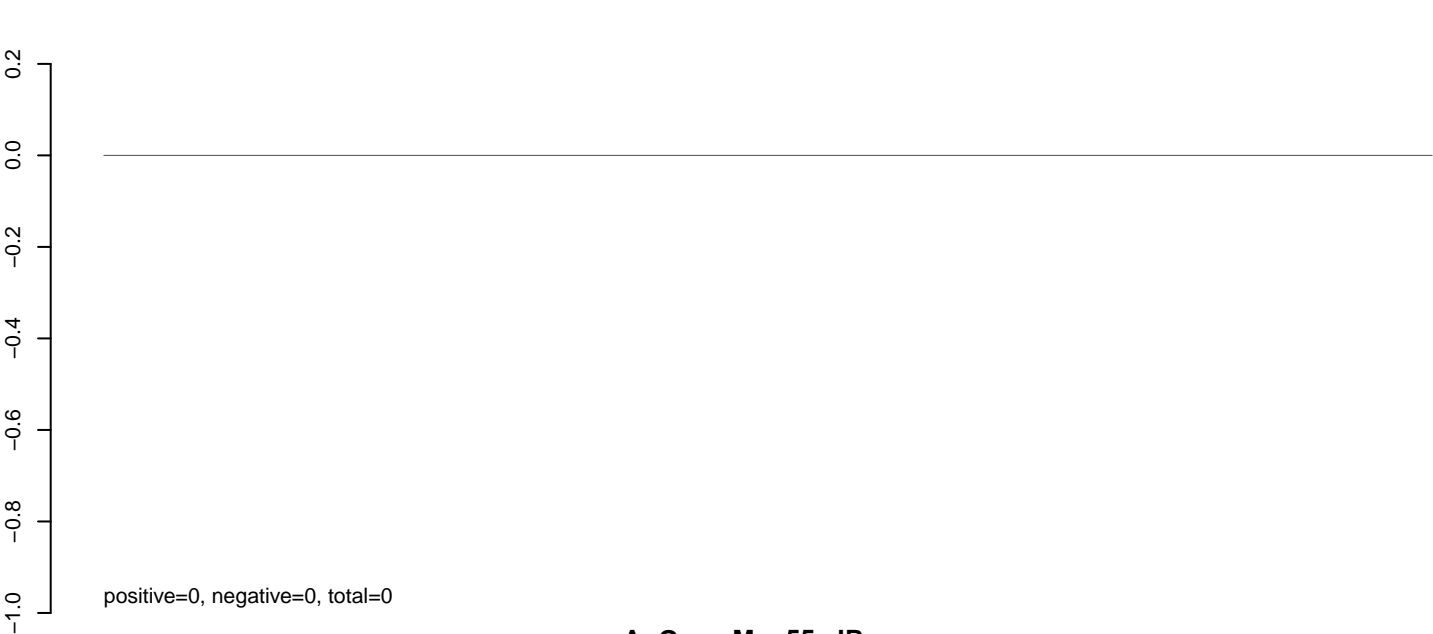


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AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



AnGam\_Mos55\_JR.rep

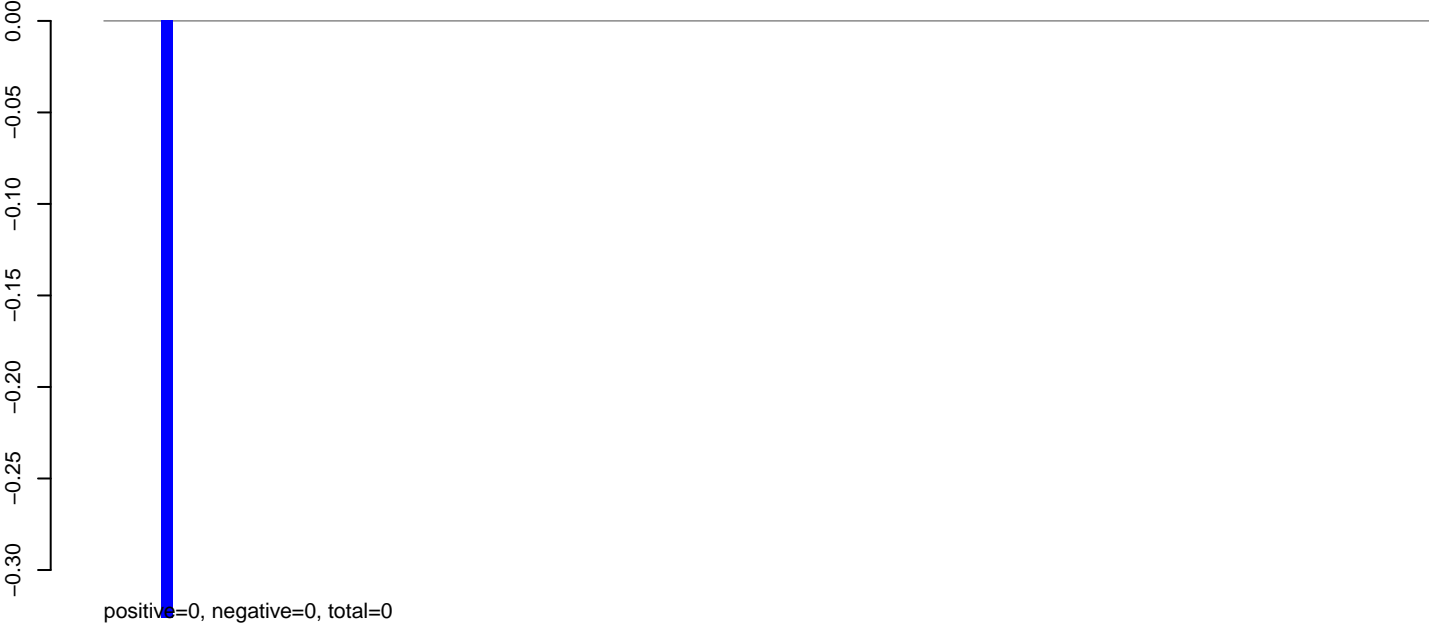


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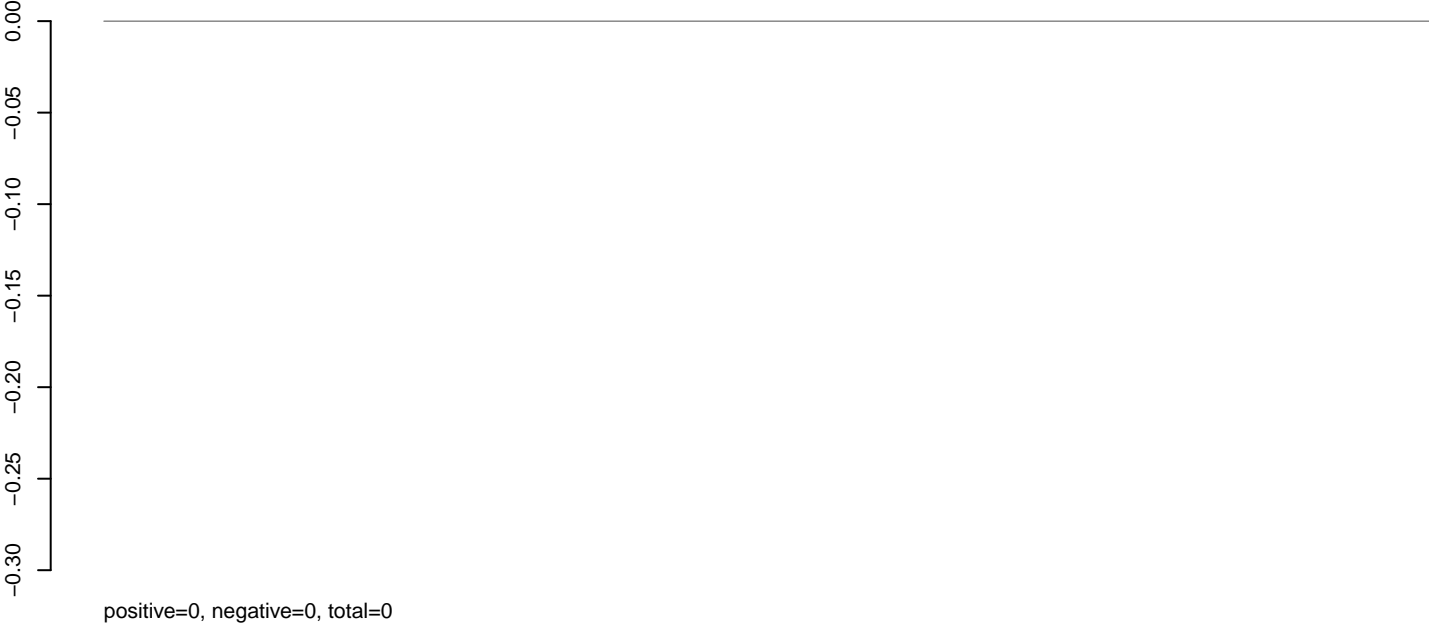
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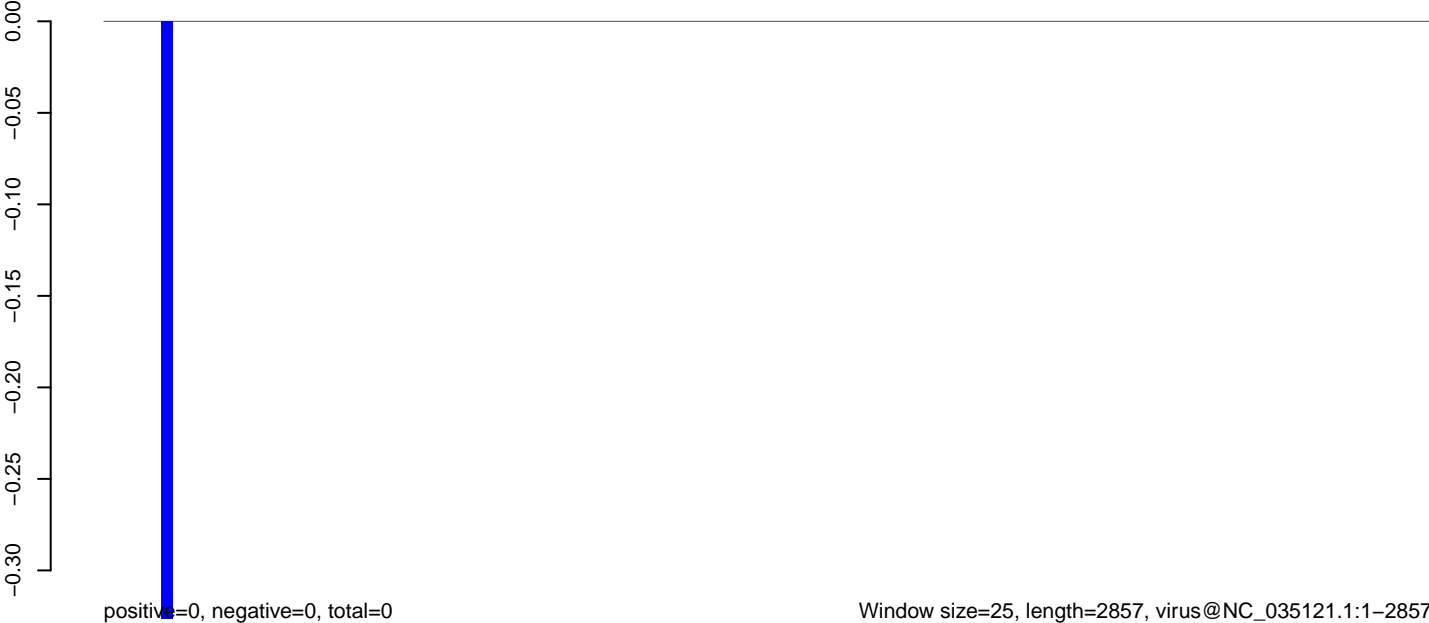
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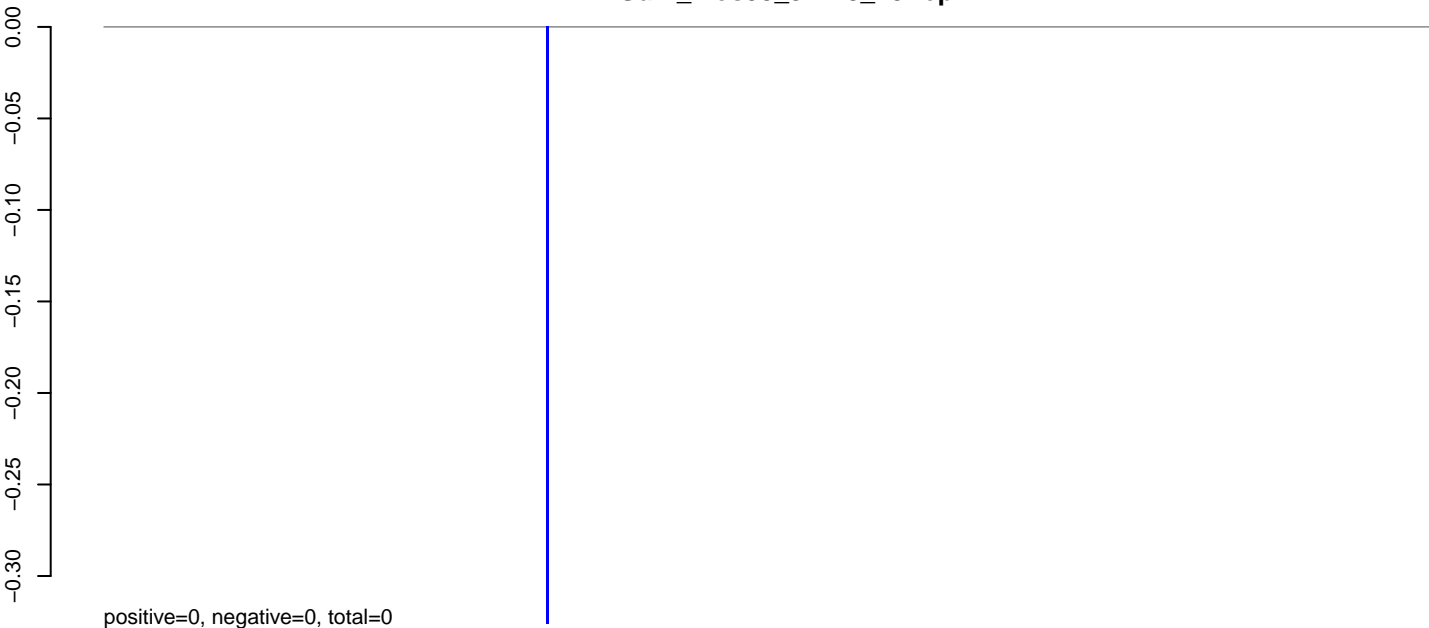
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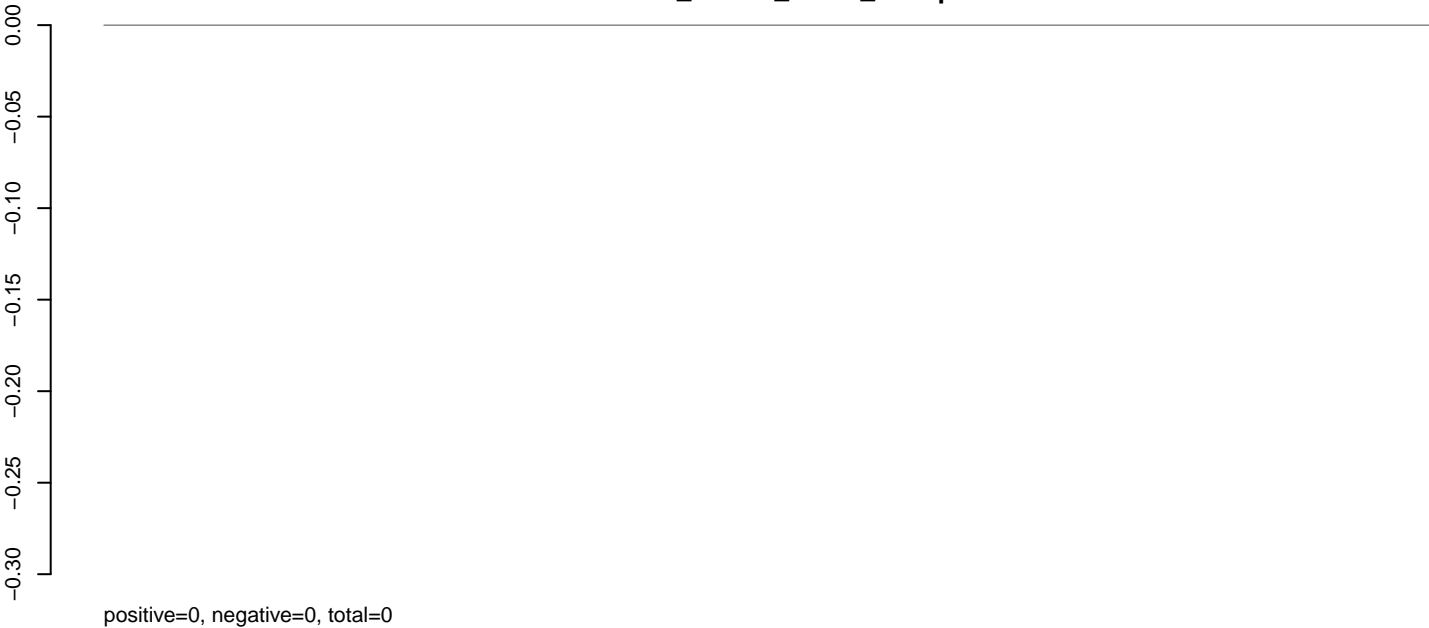
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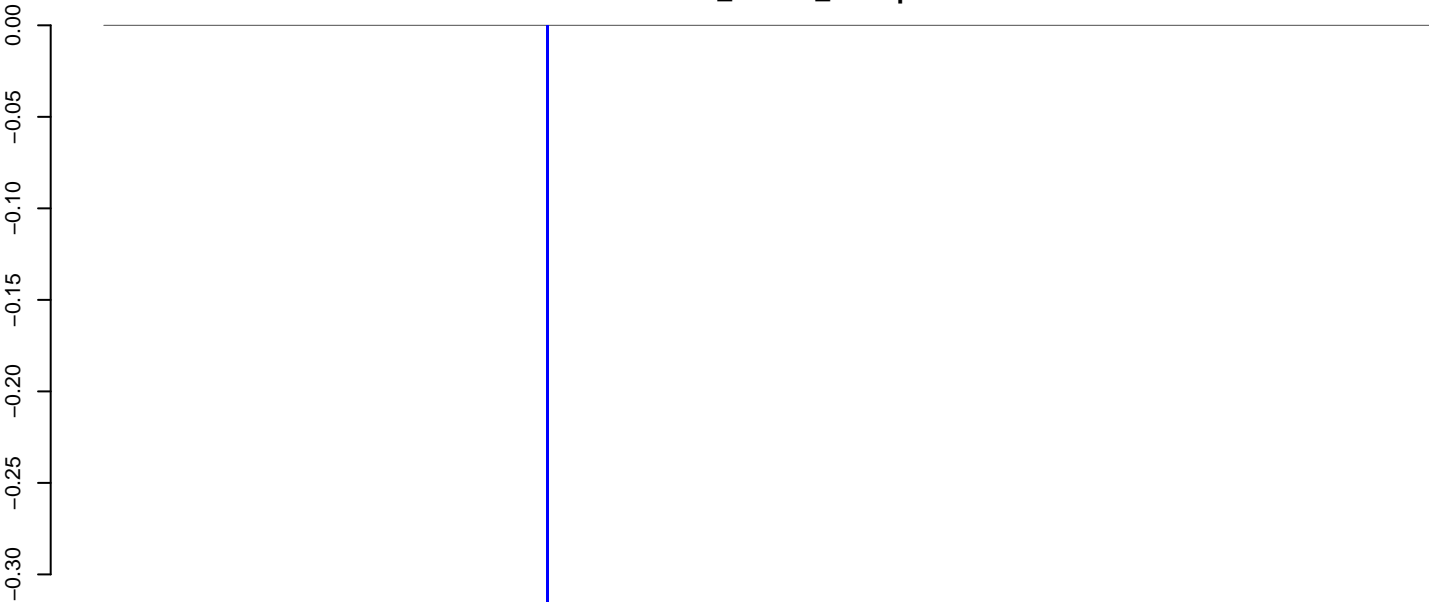
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AnGam\_Mos55\_JR.rep

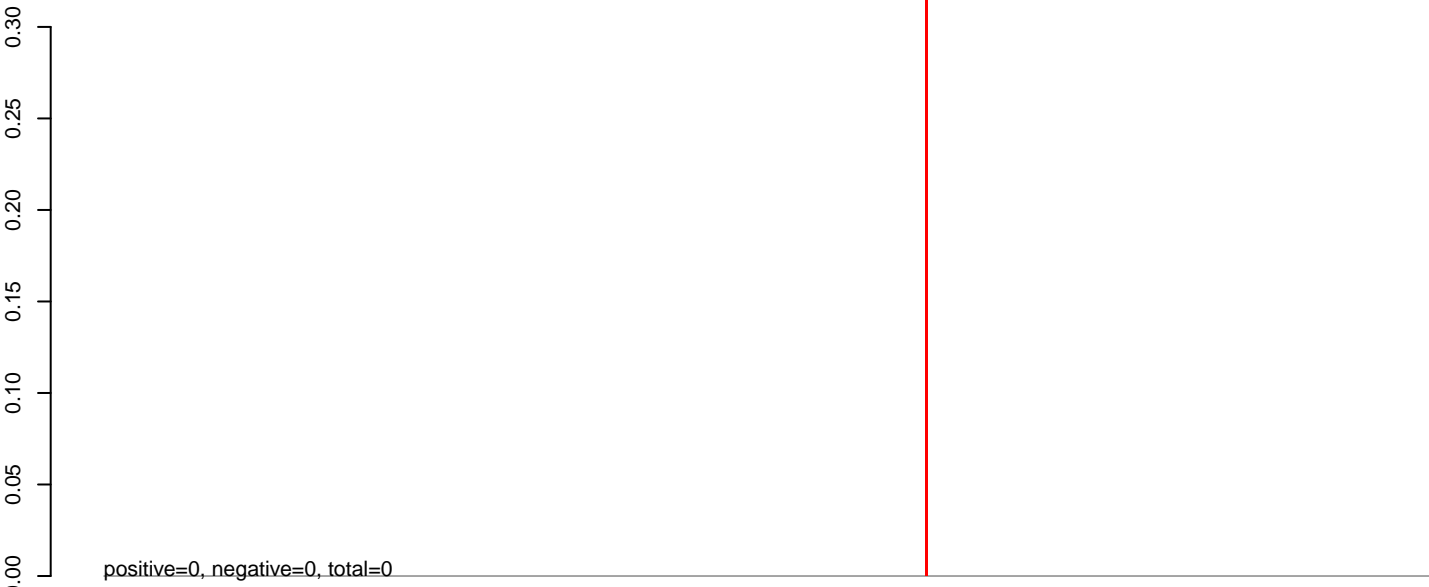


-Kamessitis, M. et al. (2015) Phosphoprotein\_hypothetical\_proteins\_matrix\_glycoprotein\_hypothetical\_protein\_and\_polymerase\_genes:1-13209

0 2000 4000 6000 8000 10000 12000

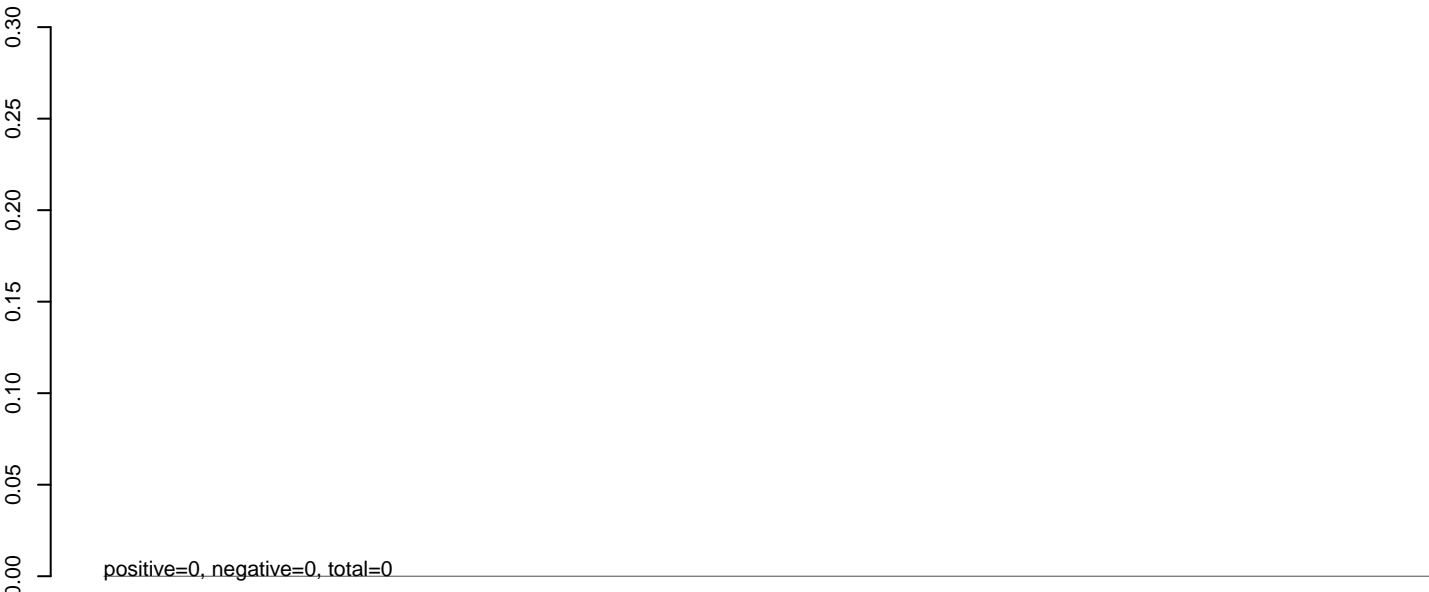


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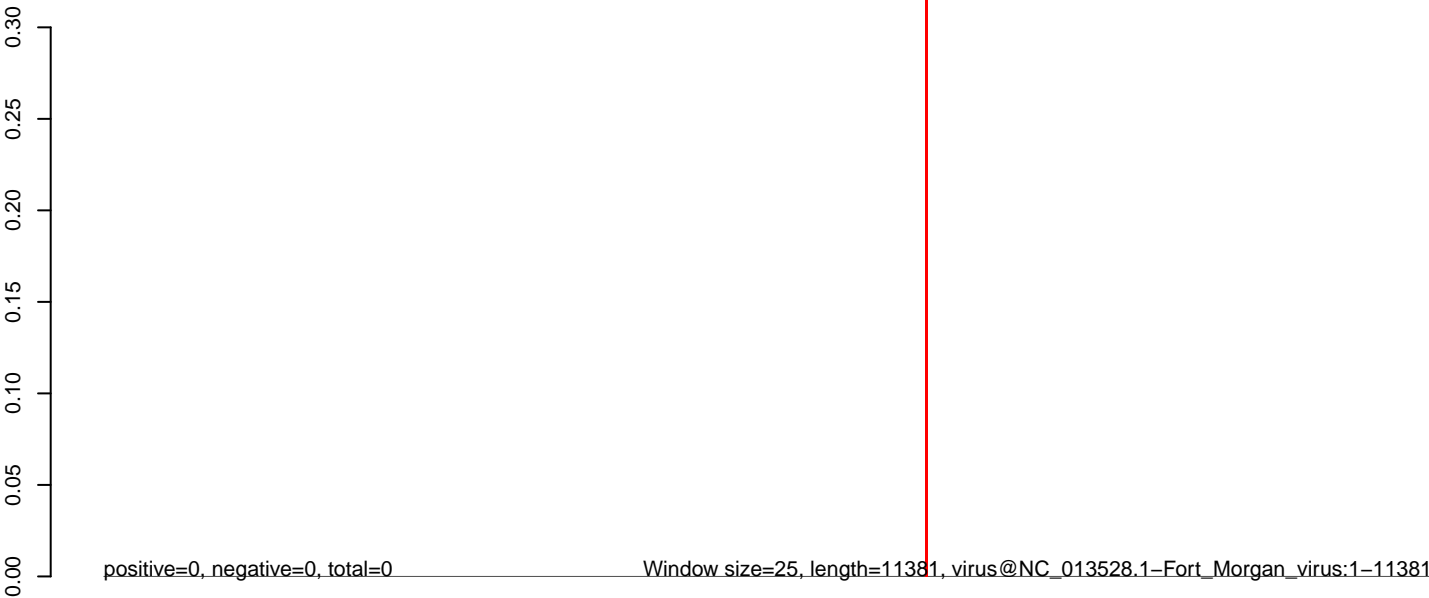
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AnGam\_Mos55\_JR.24\_35.rep



positive=0, negative=0, total=0

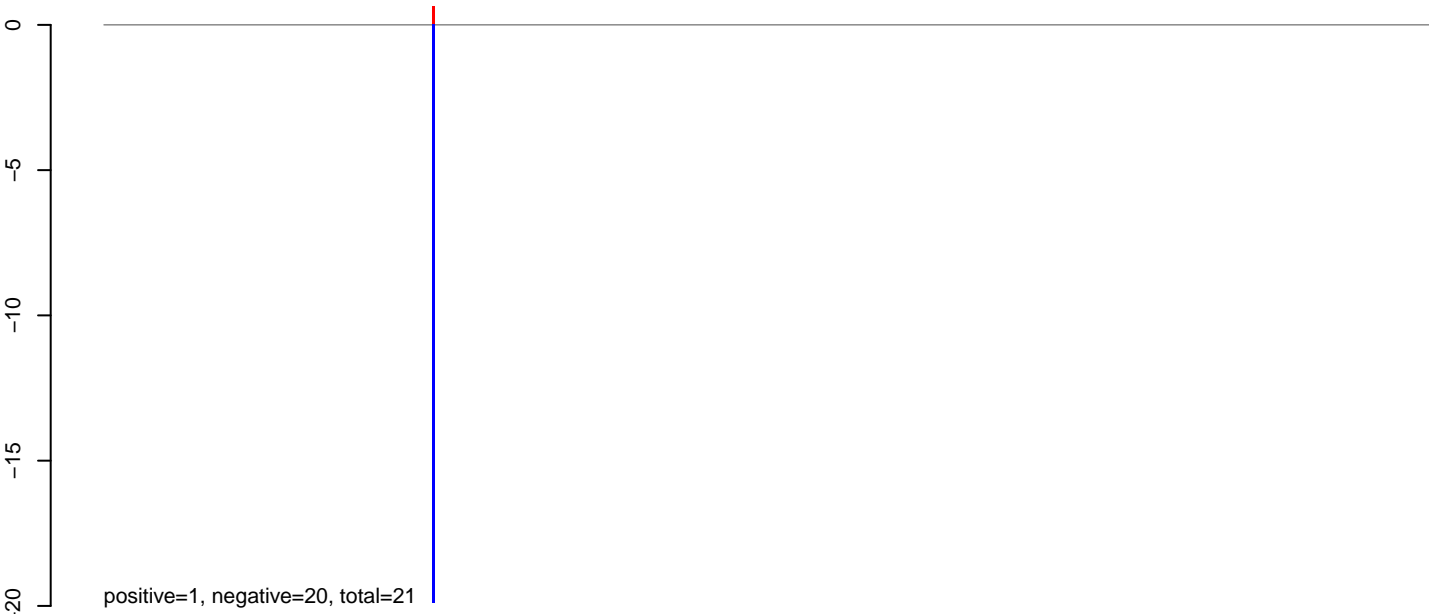
AnGam\_Mos55\_JR.rep



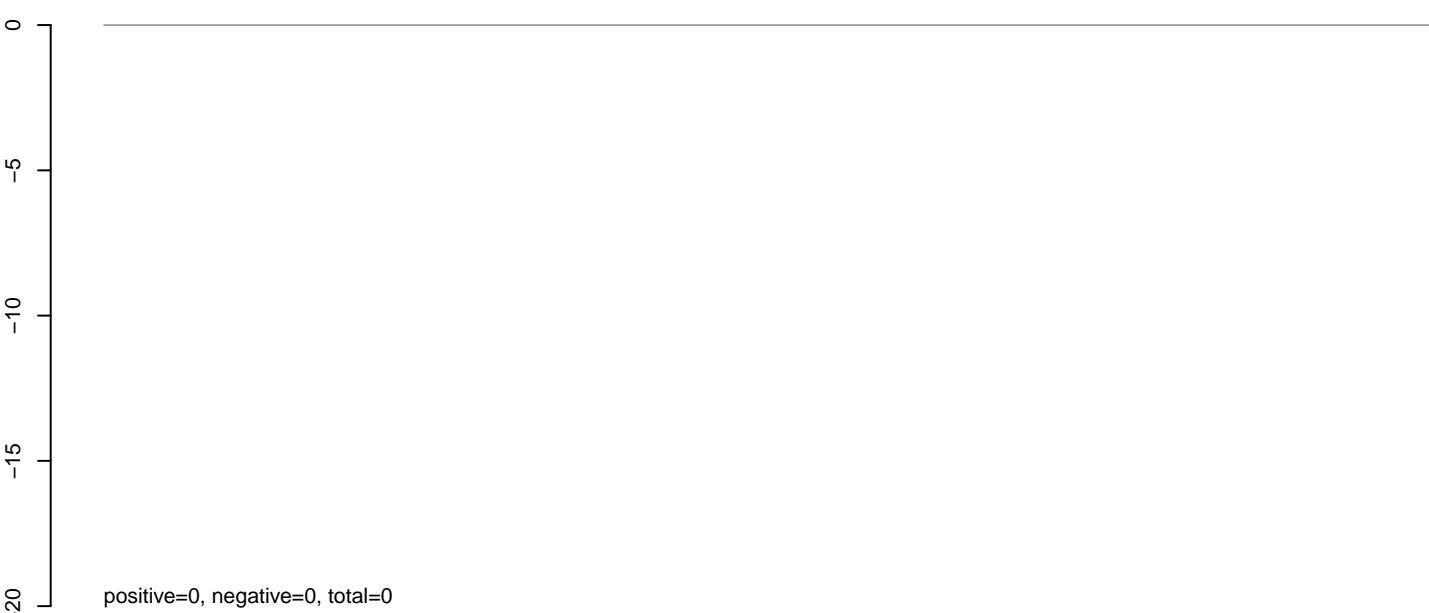
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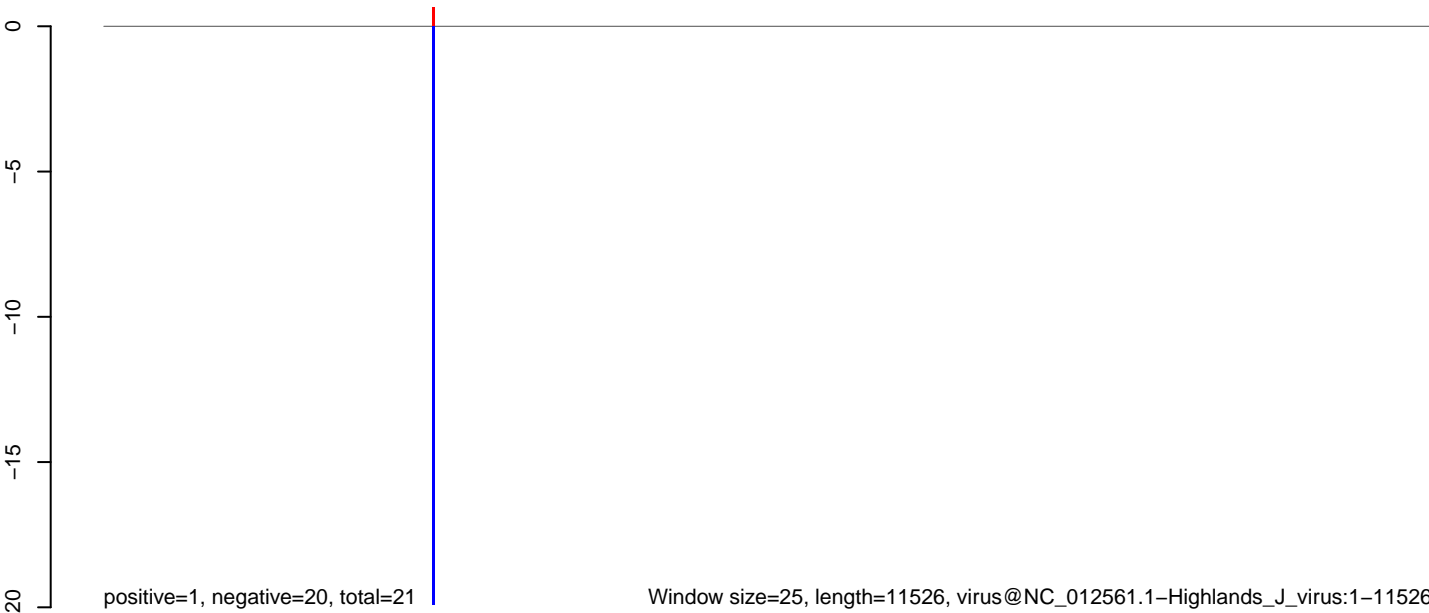
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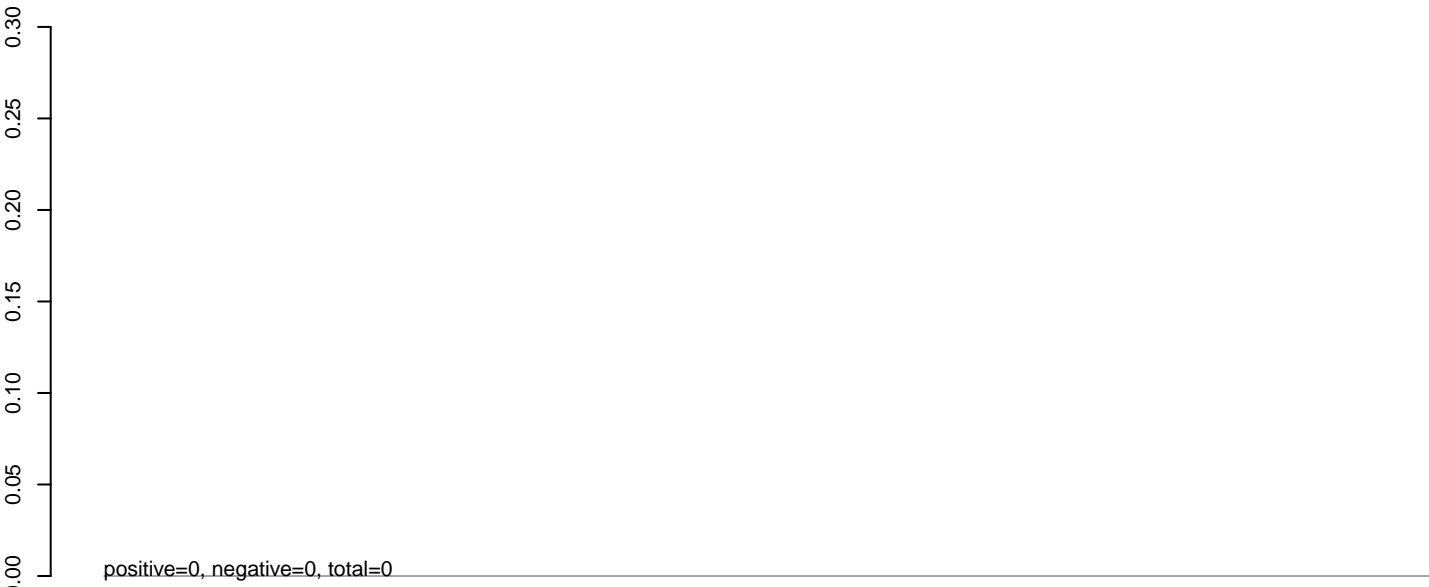
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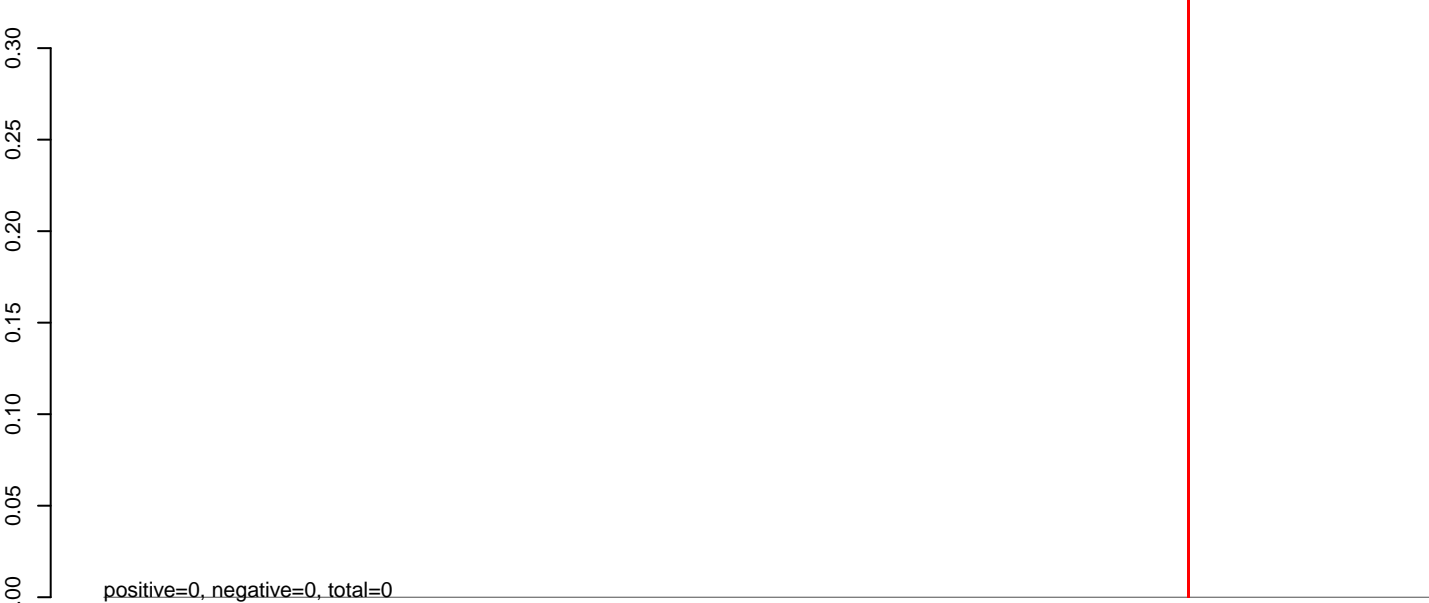
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AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep

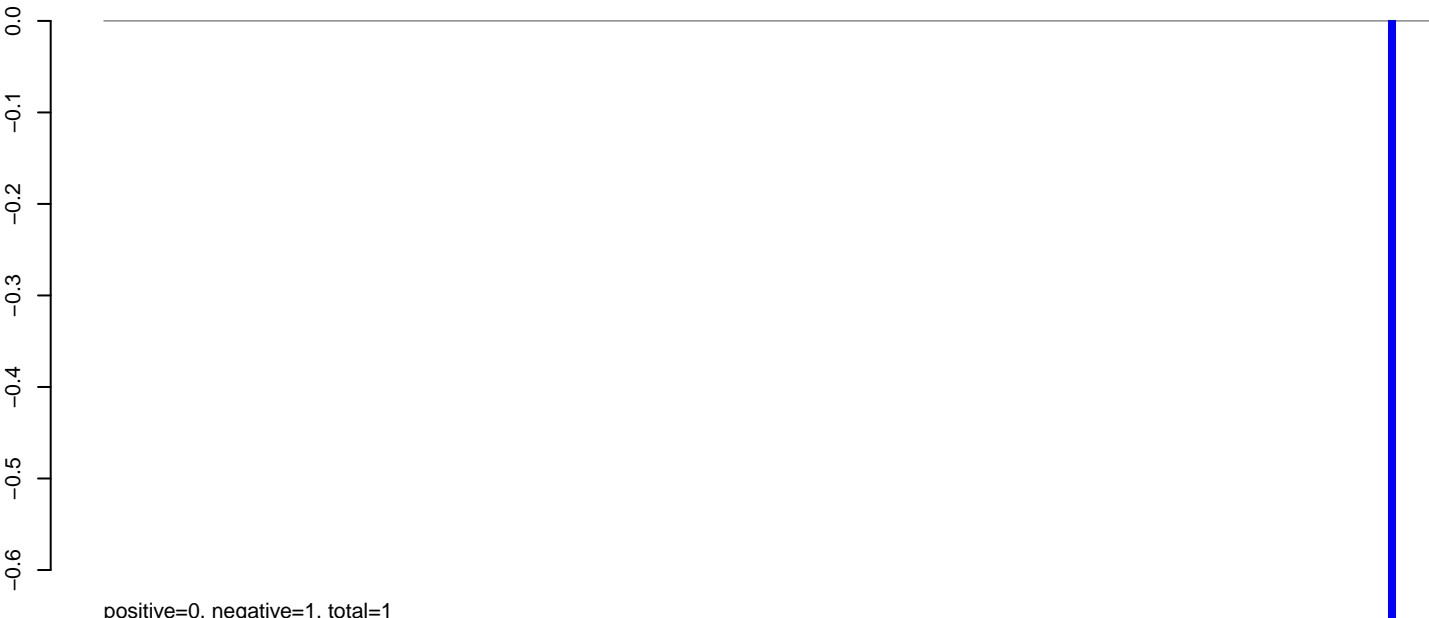


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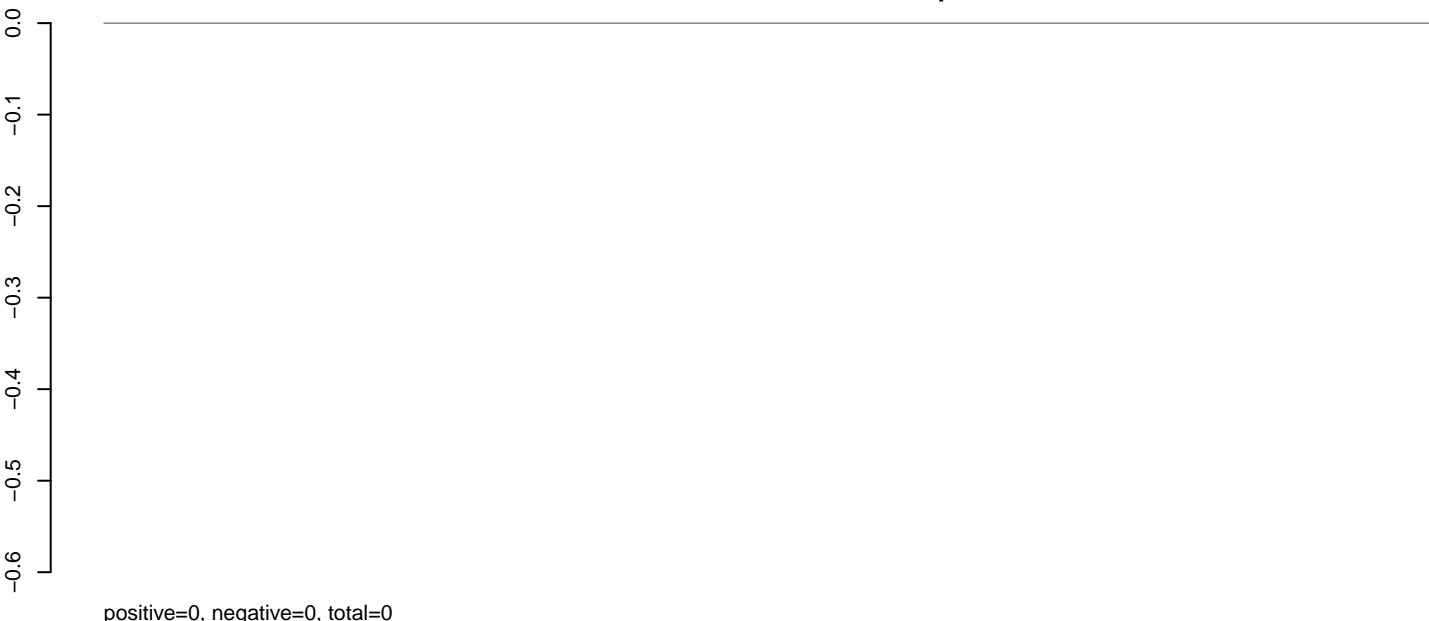
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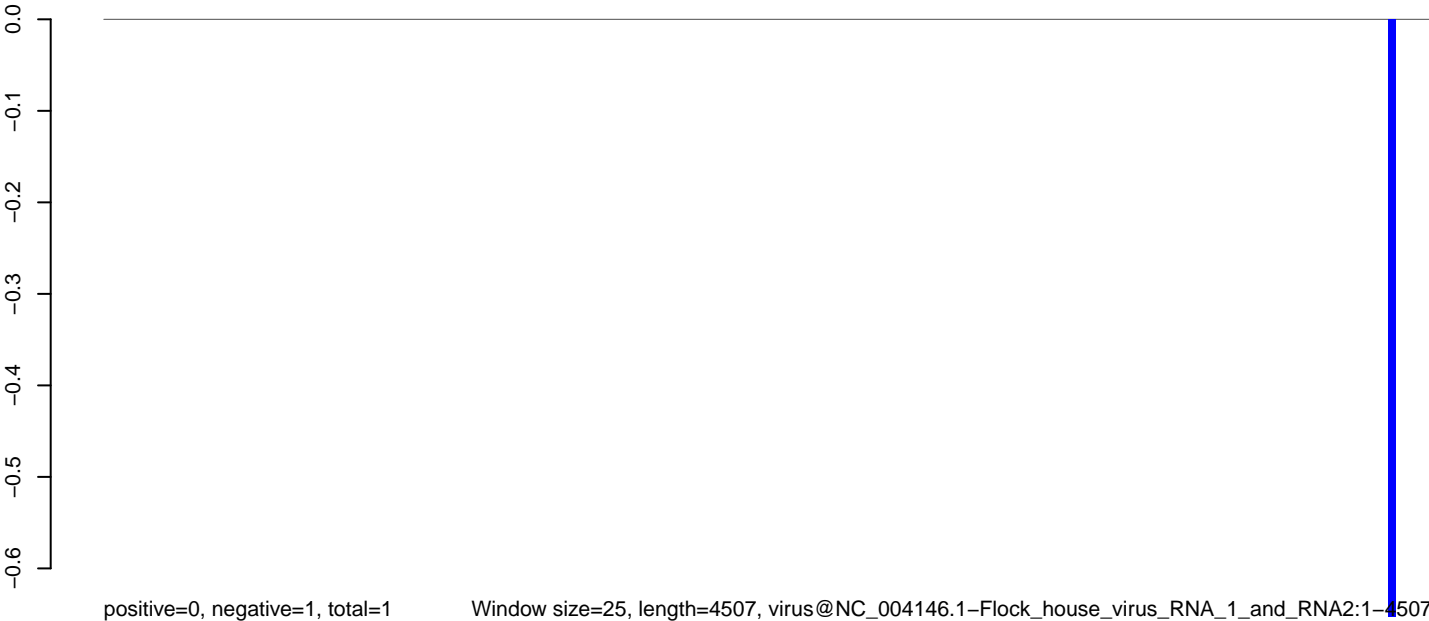
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AnGam\_Mos55\_JR.24\_35.rep



positive=0, negative=0, total=0

AnGam\_Mos55\_JR.rep

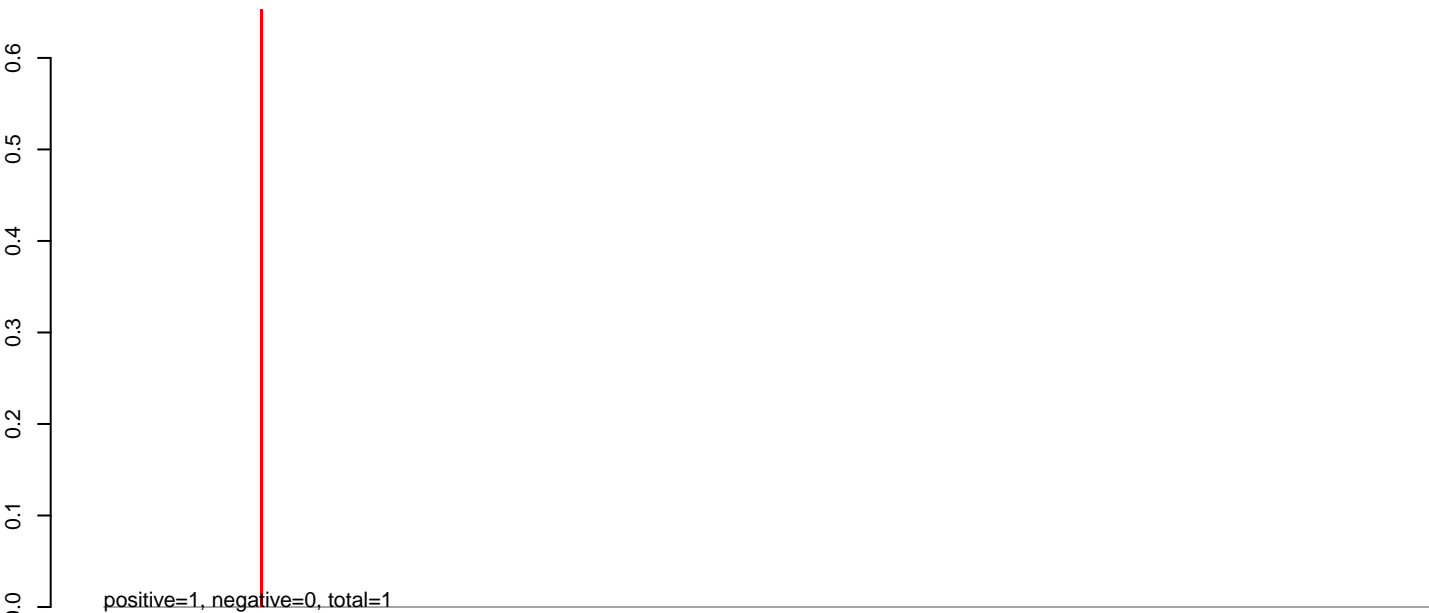


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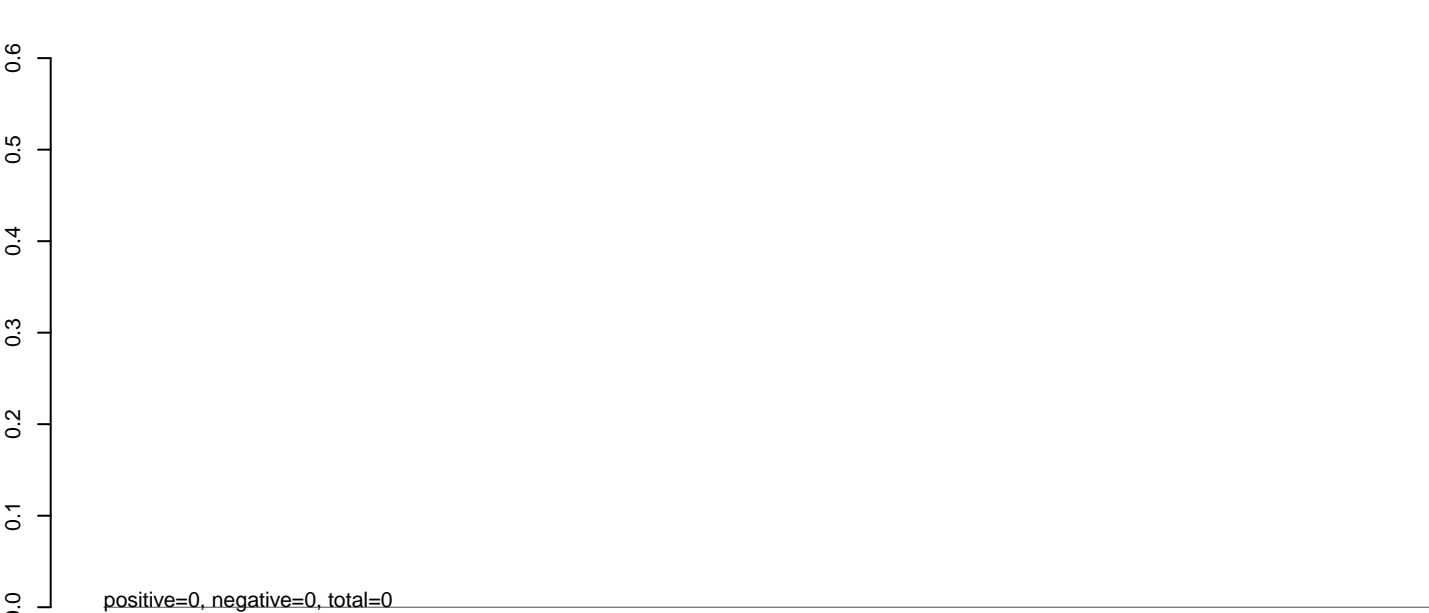
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0 1000 2000 3000 4000

AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep

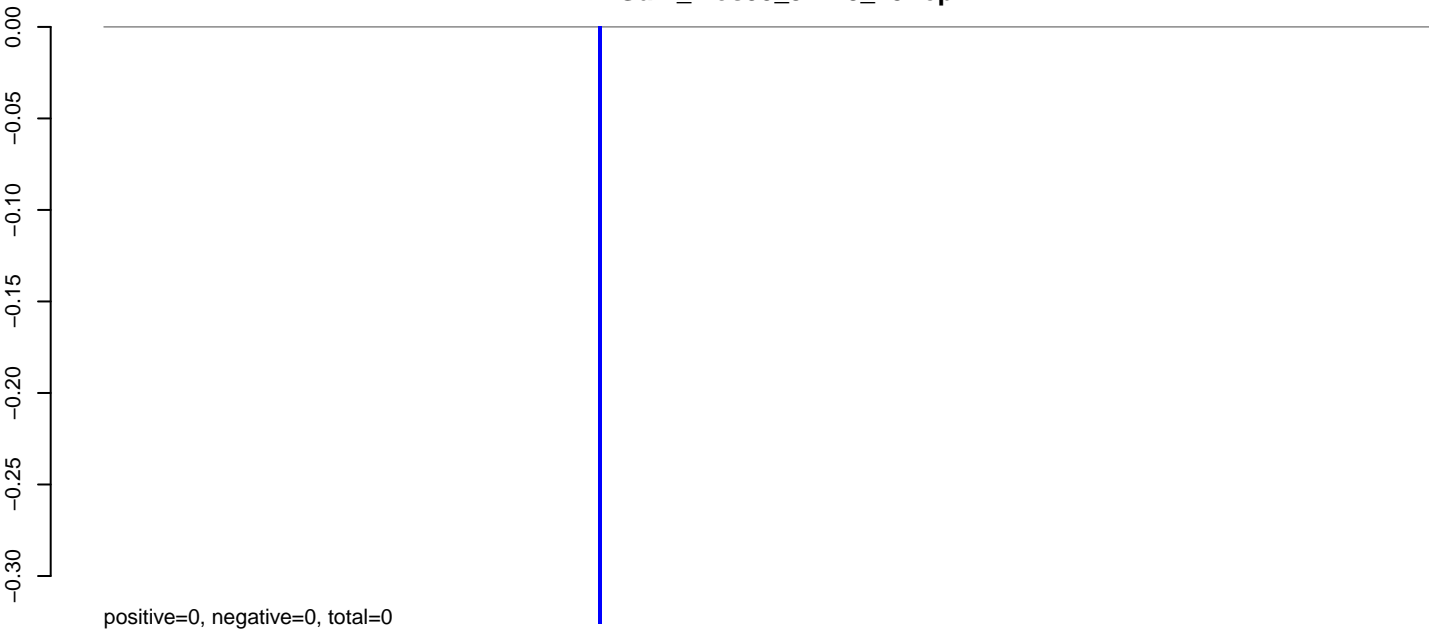


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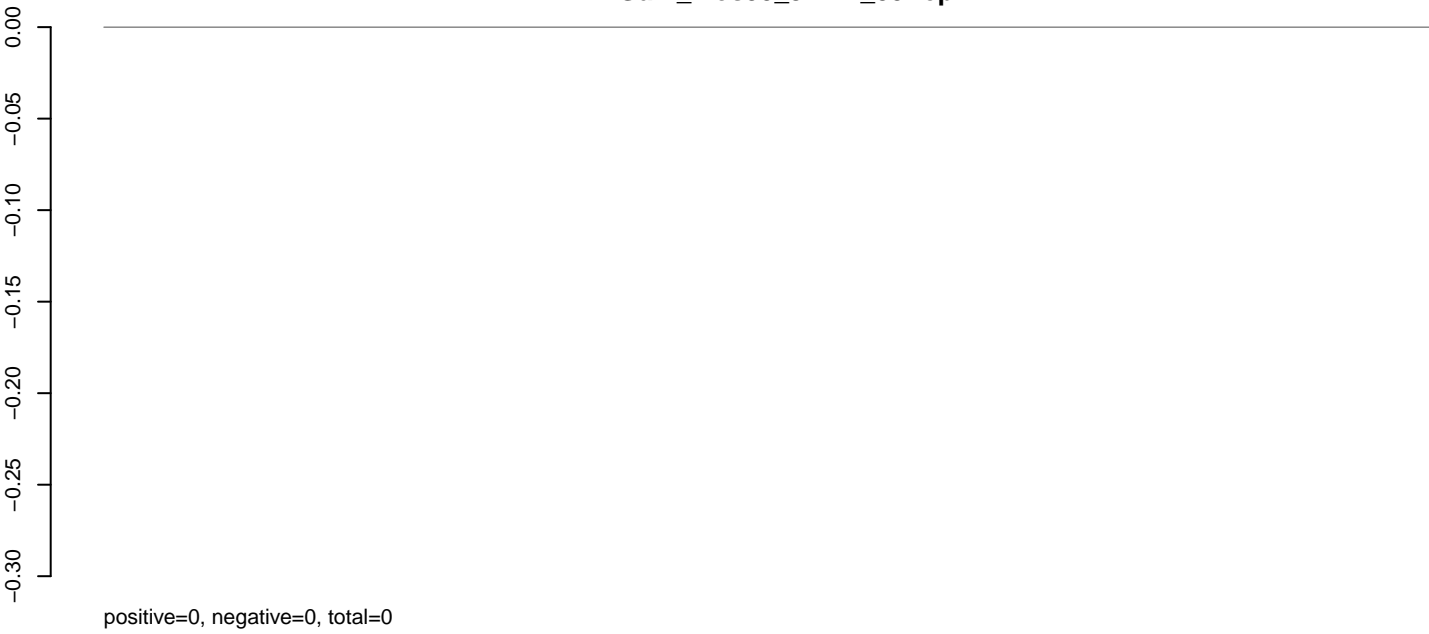




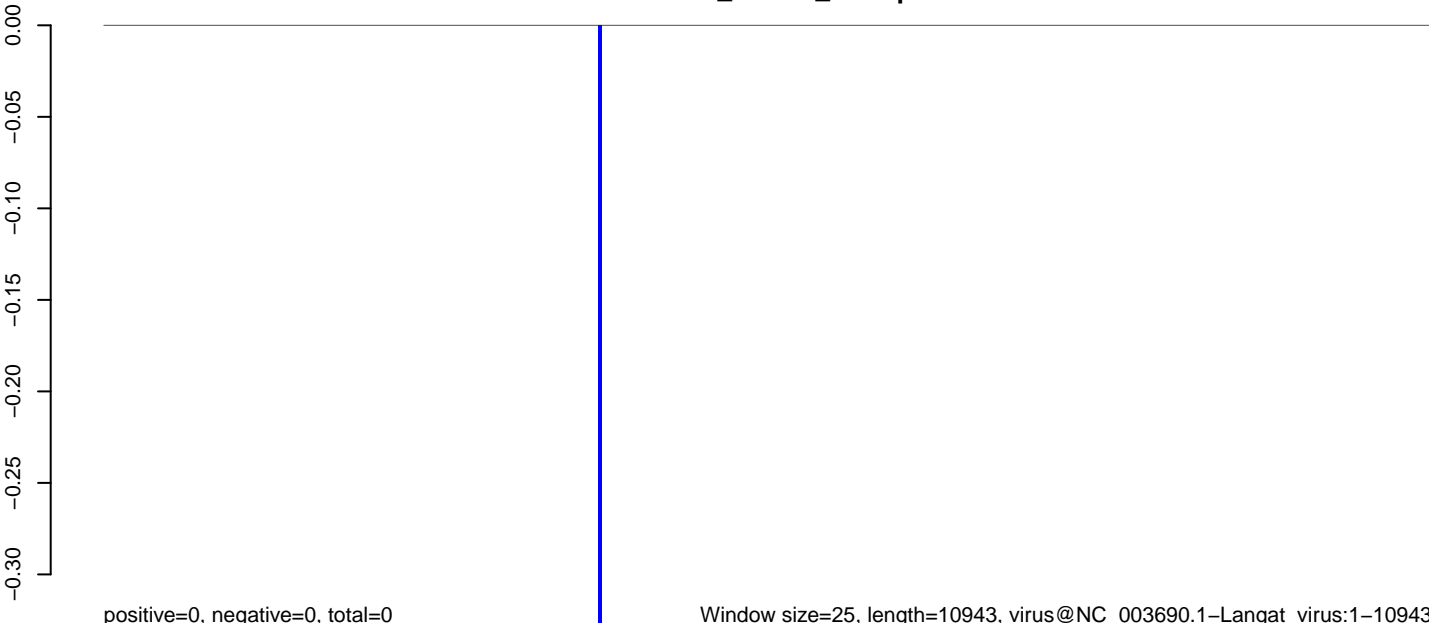
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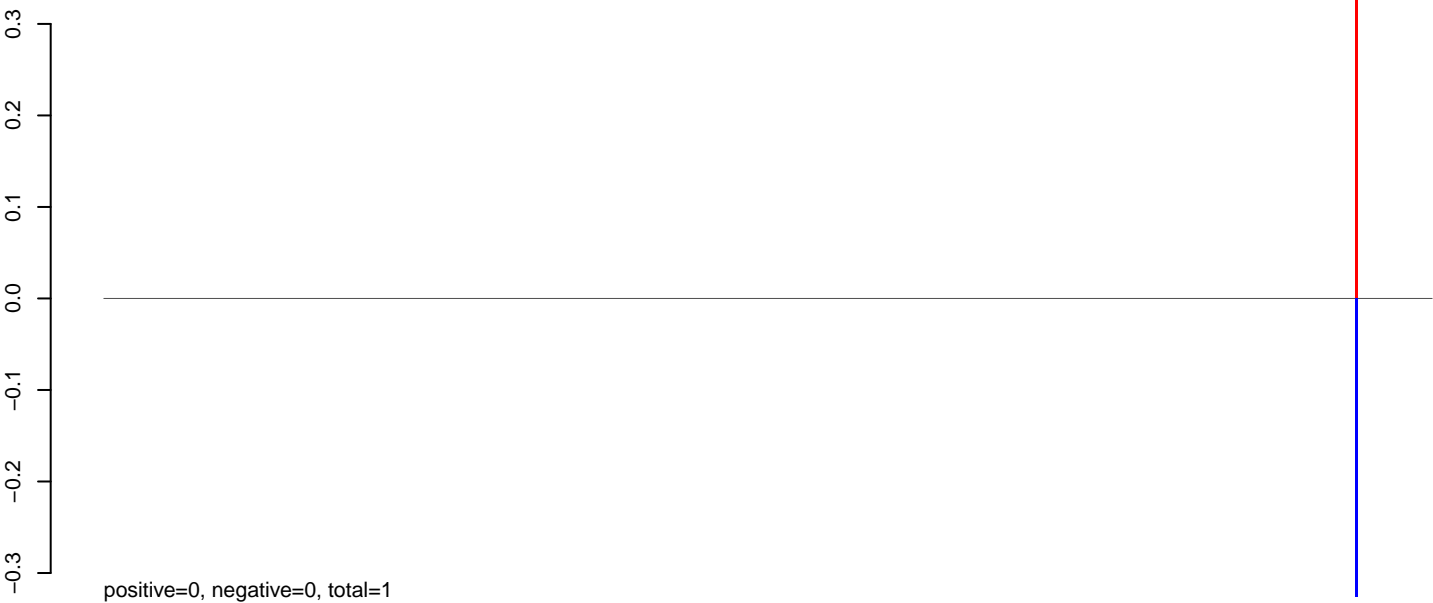
AnGam\_Mos55\_JR.24\_35.rep



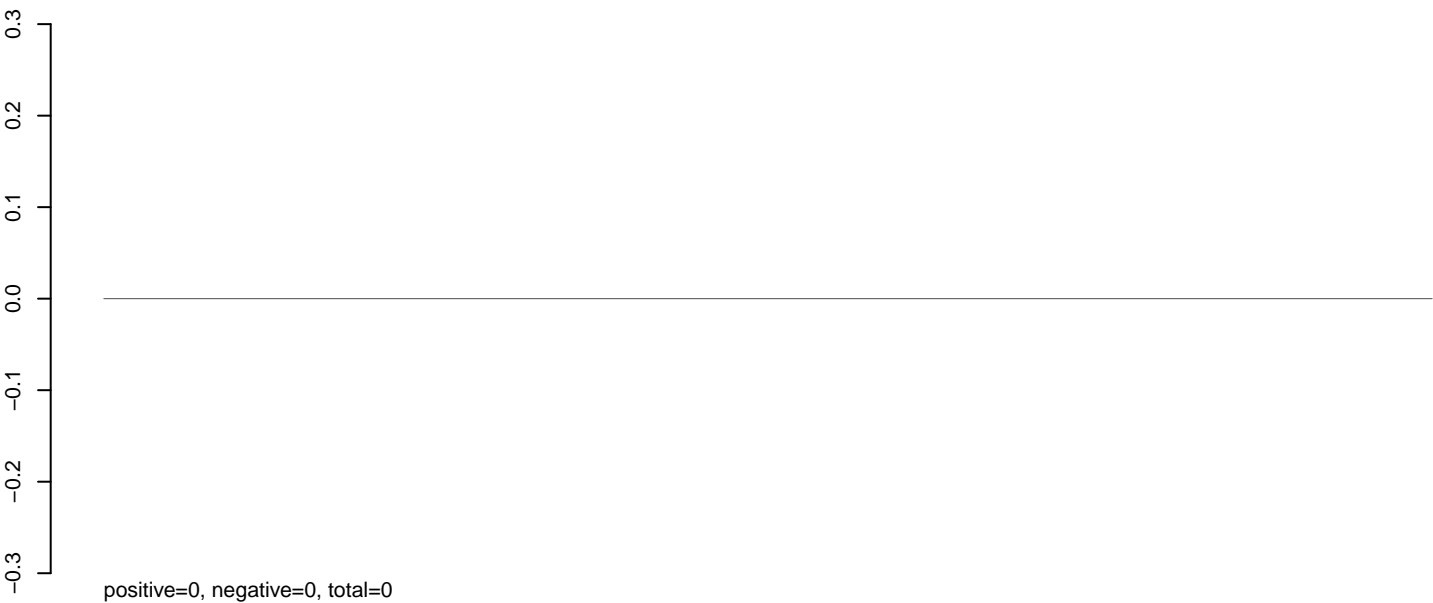
AnGam\_Mos55\_JR.rep



AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



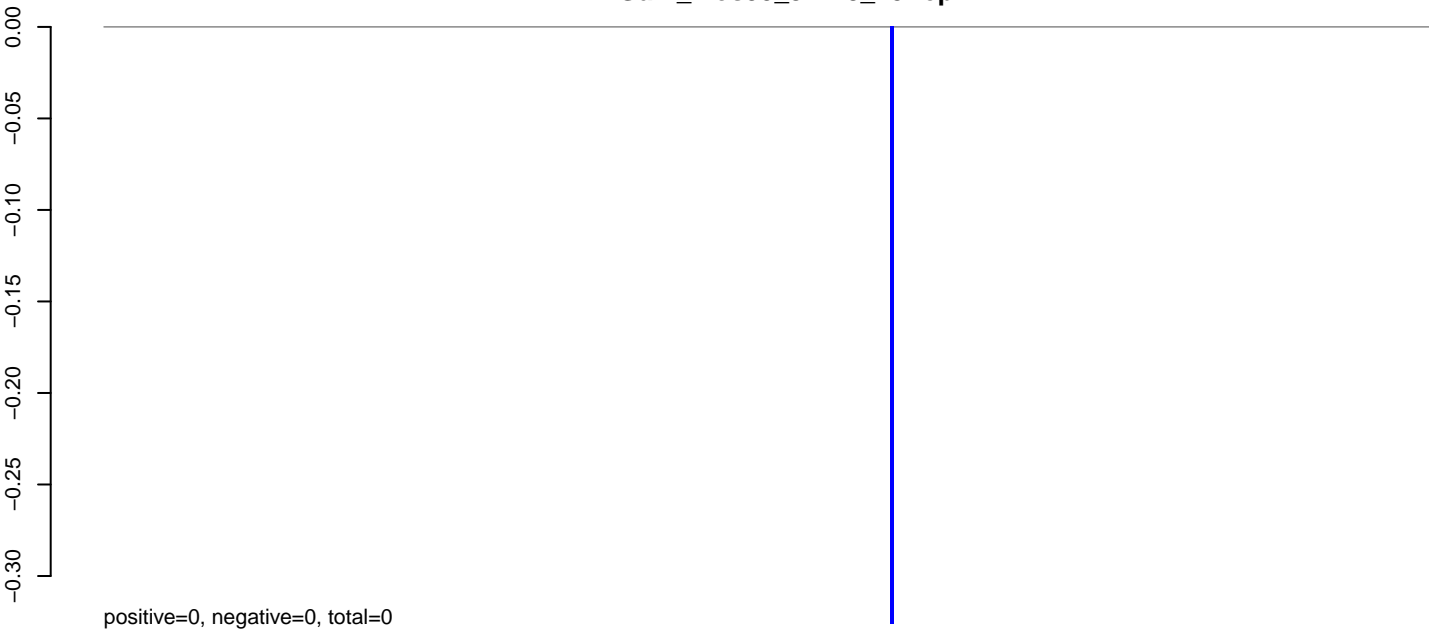
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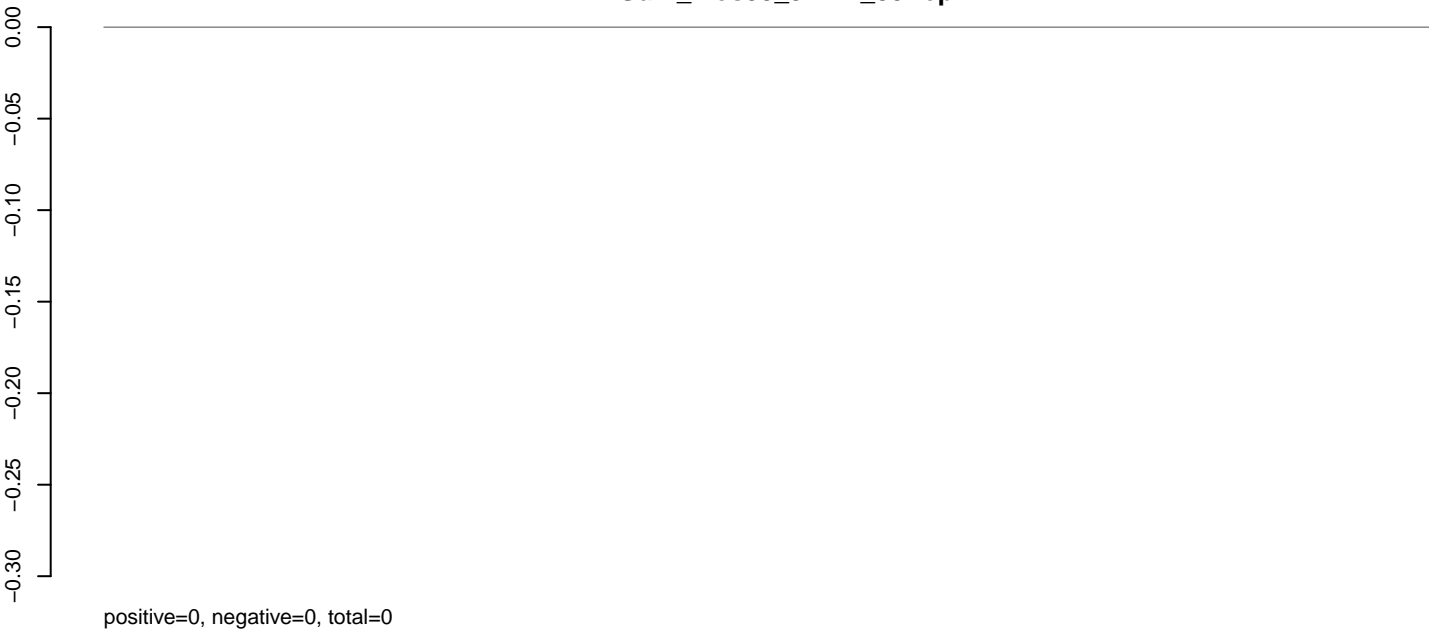
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0 2000 4000 6000 8000 10000

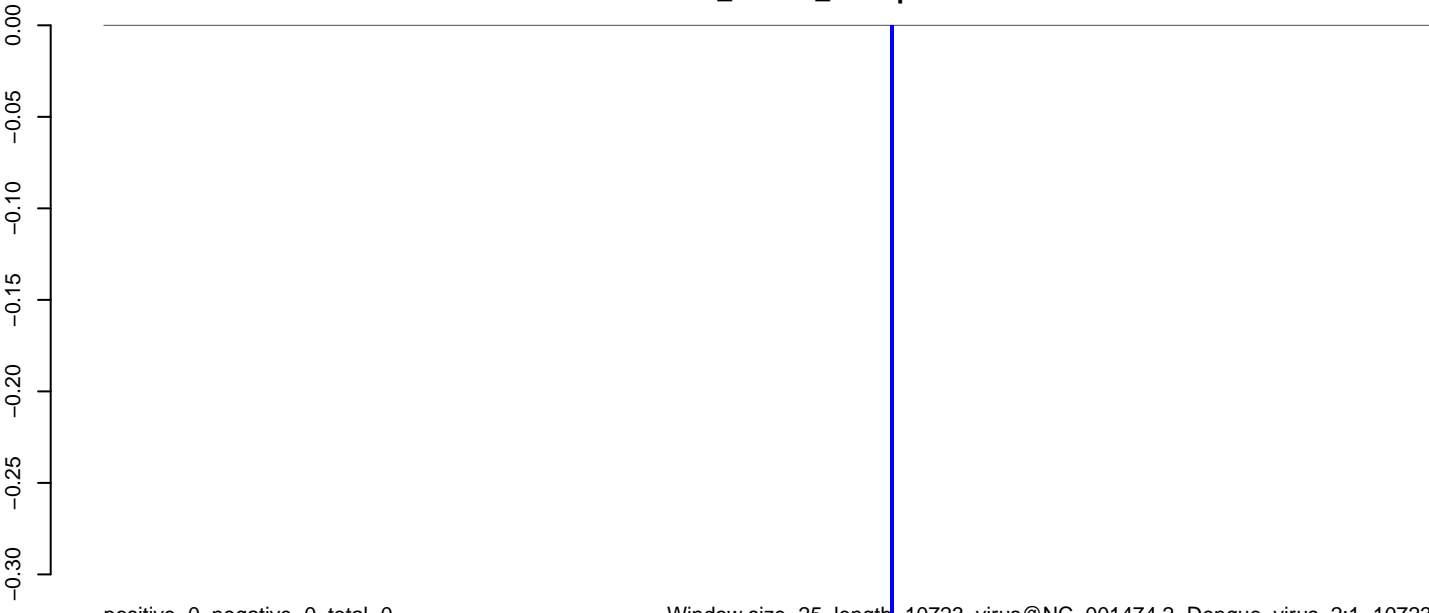
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AnGam\_Mos55\_JR.24\_35.rep



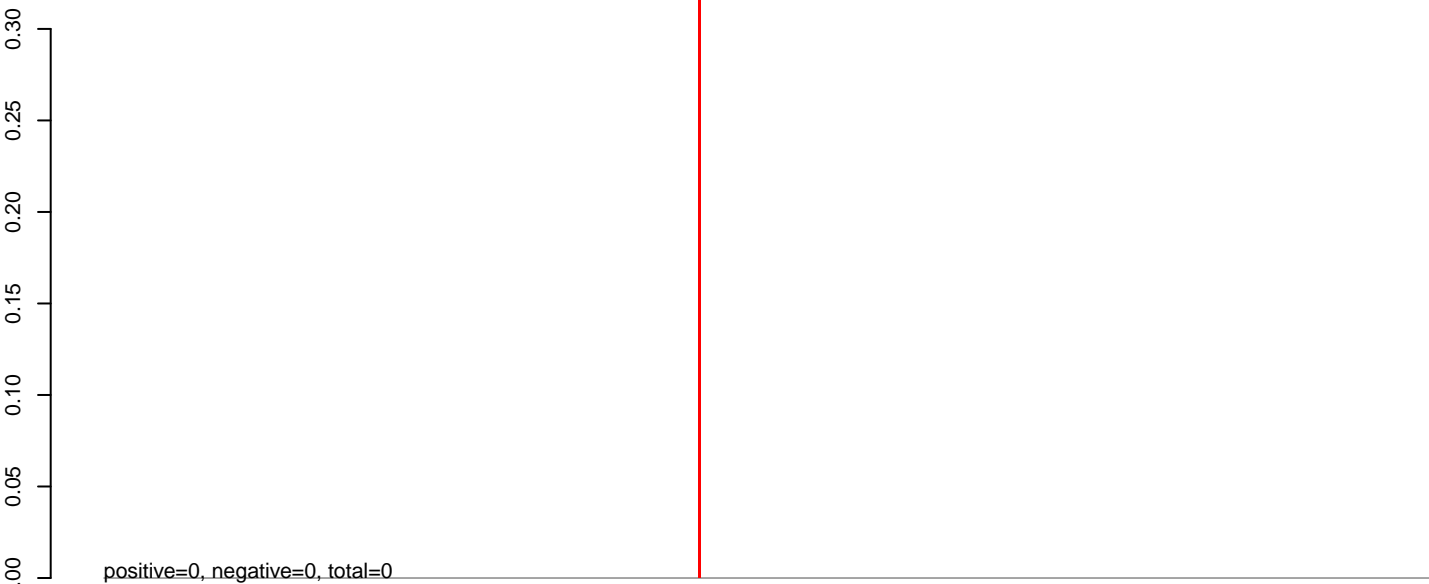
AnGam\_Mos55\_JR.rep



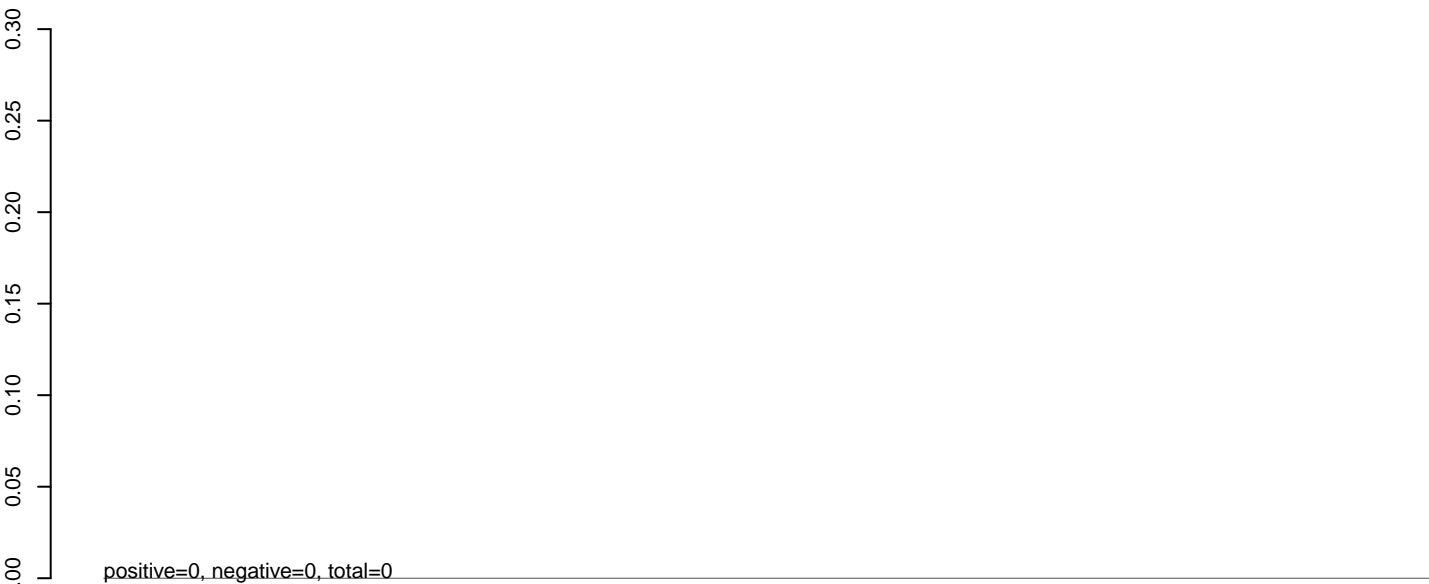
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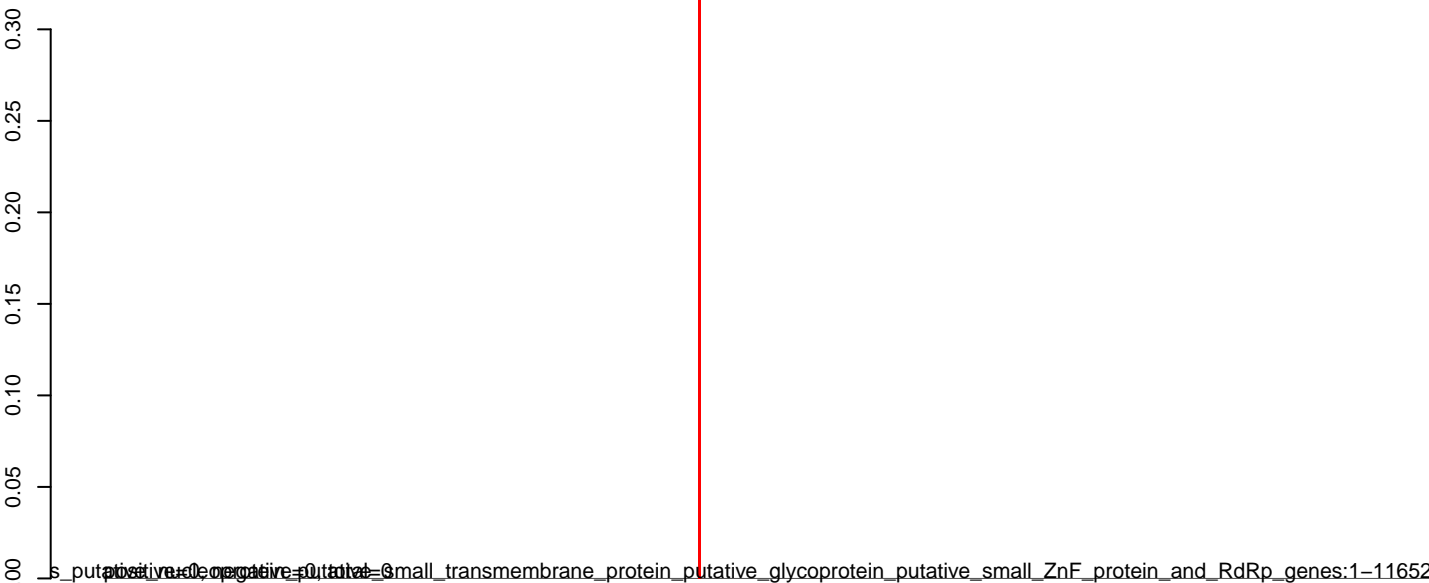
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AnGam\_Mos55\_JR.24\_35.rep

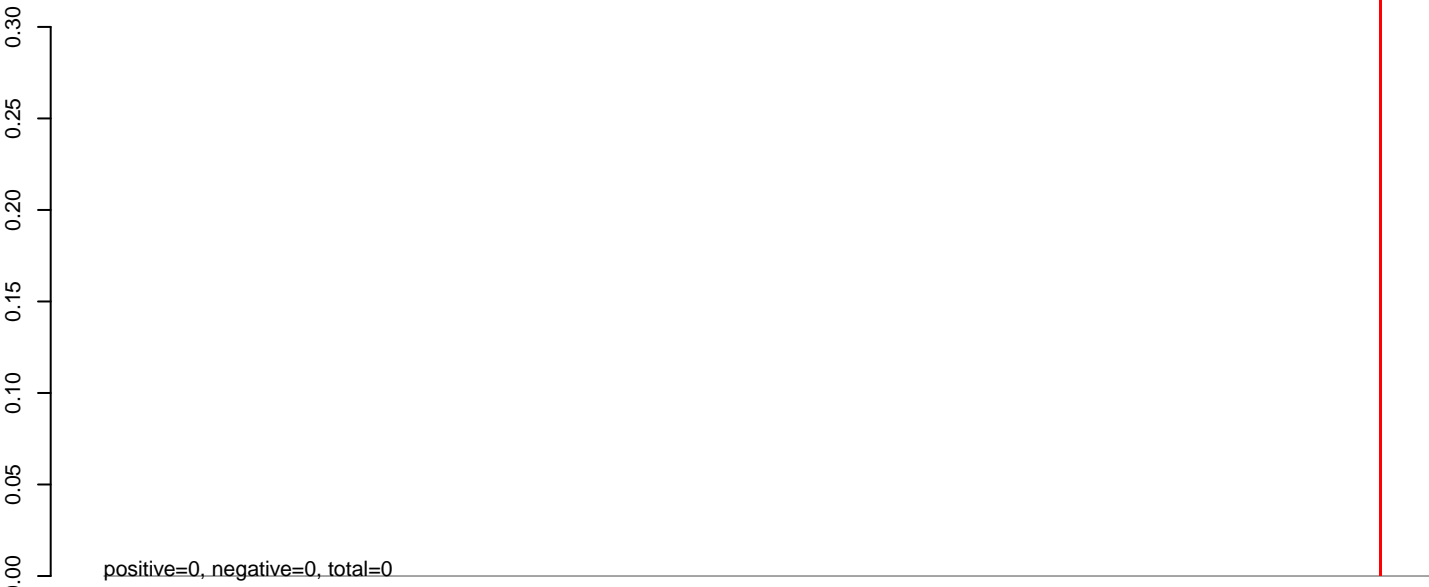


AnGam\_Mos55\_JR.rep

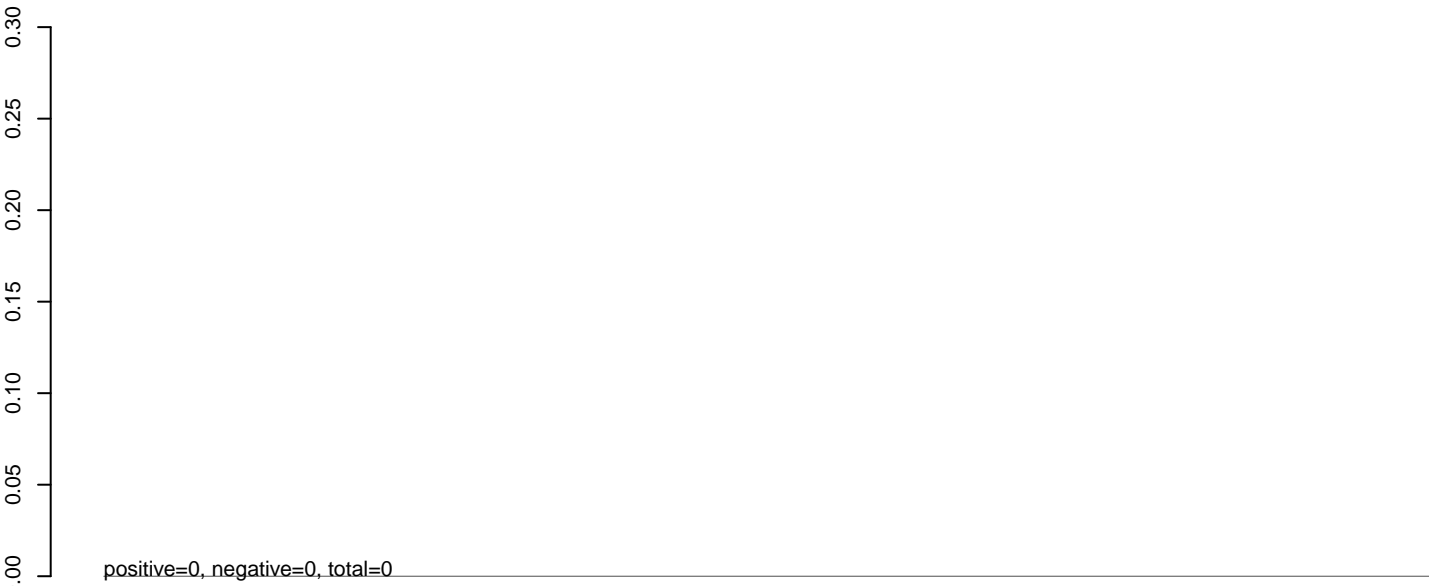


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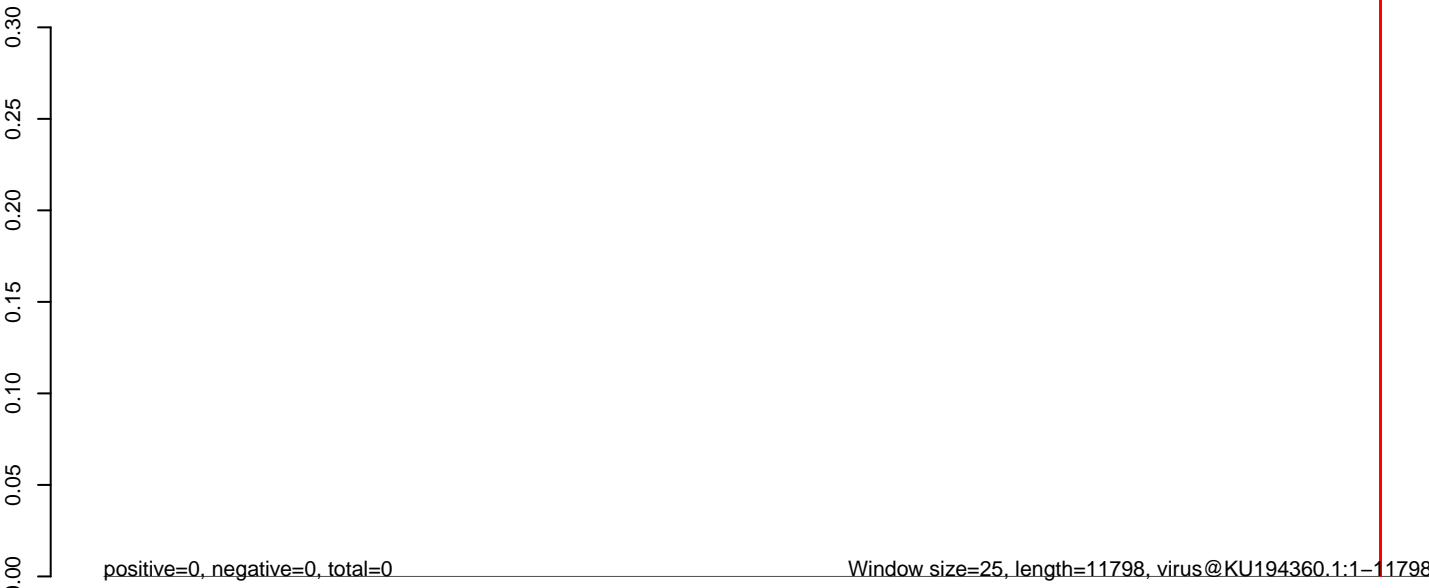
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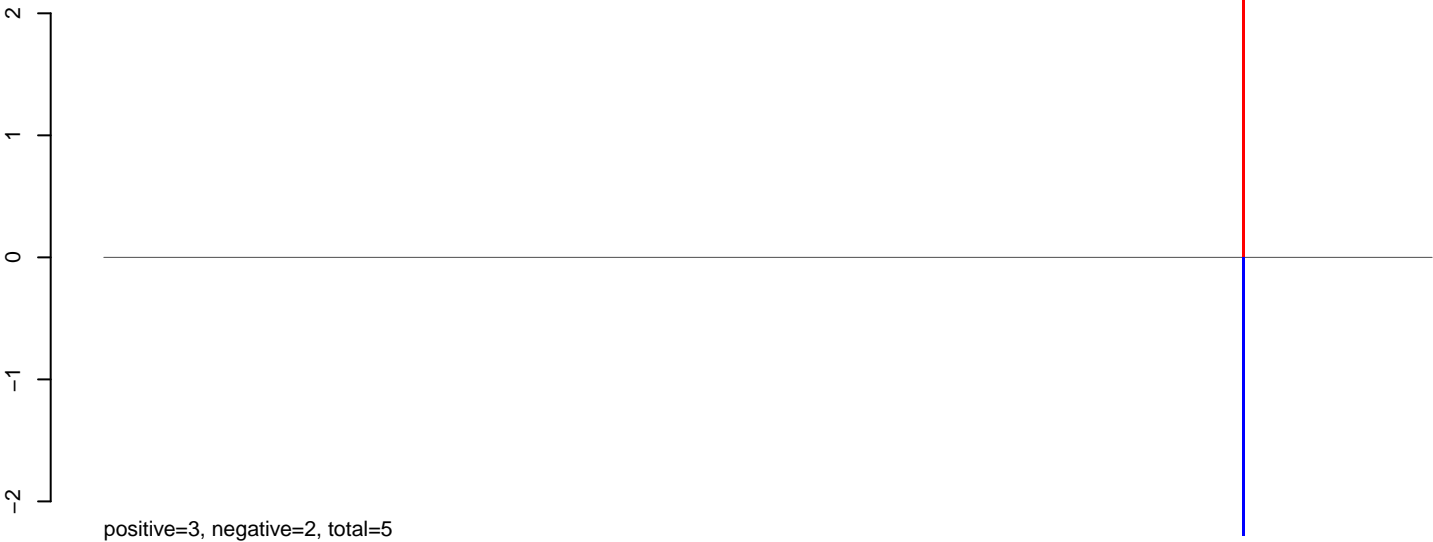


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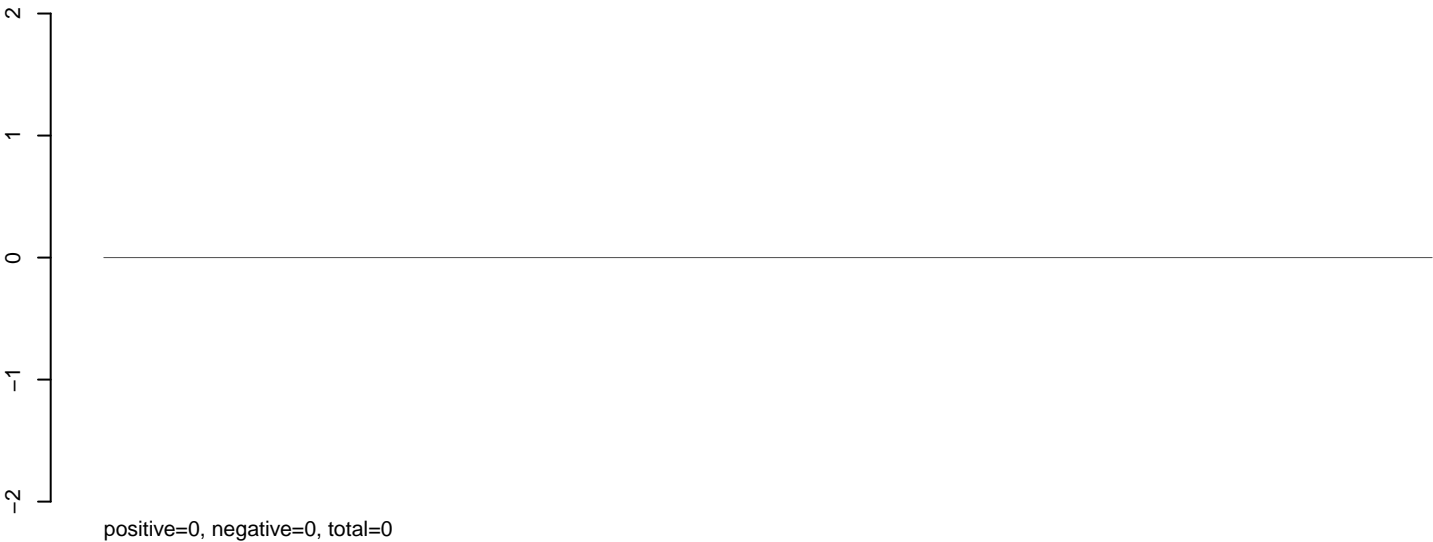


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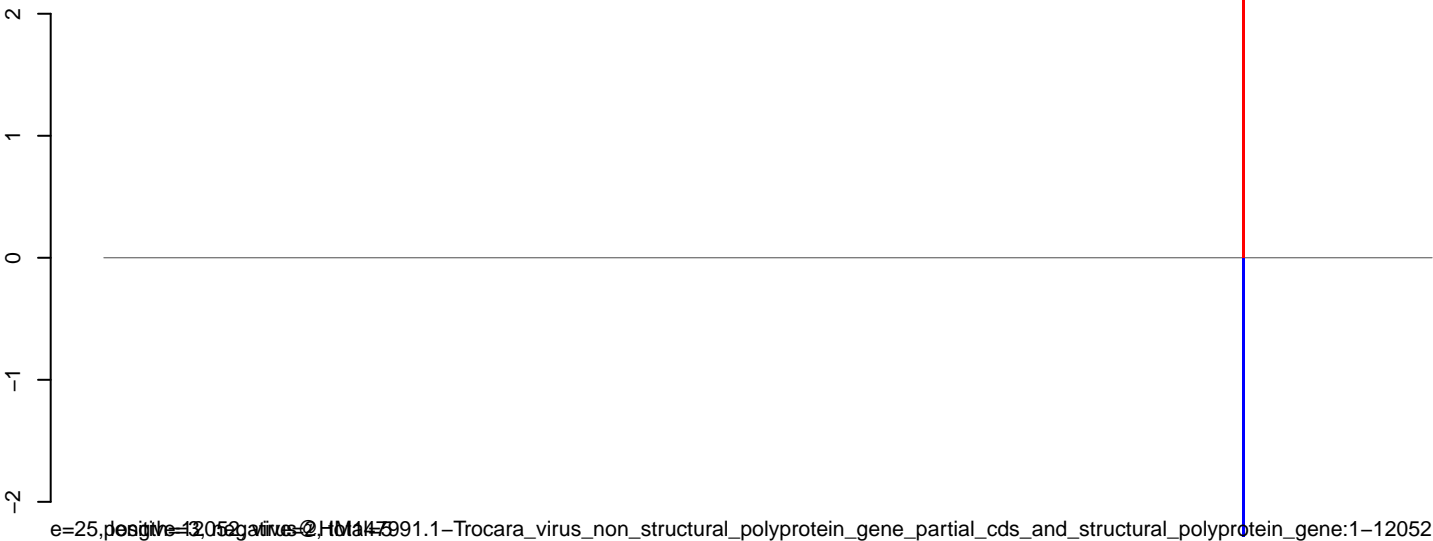
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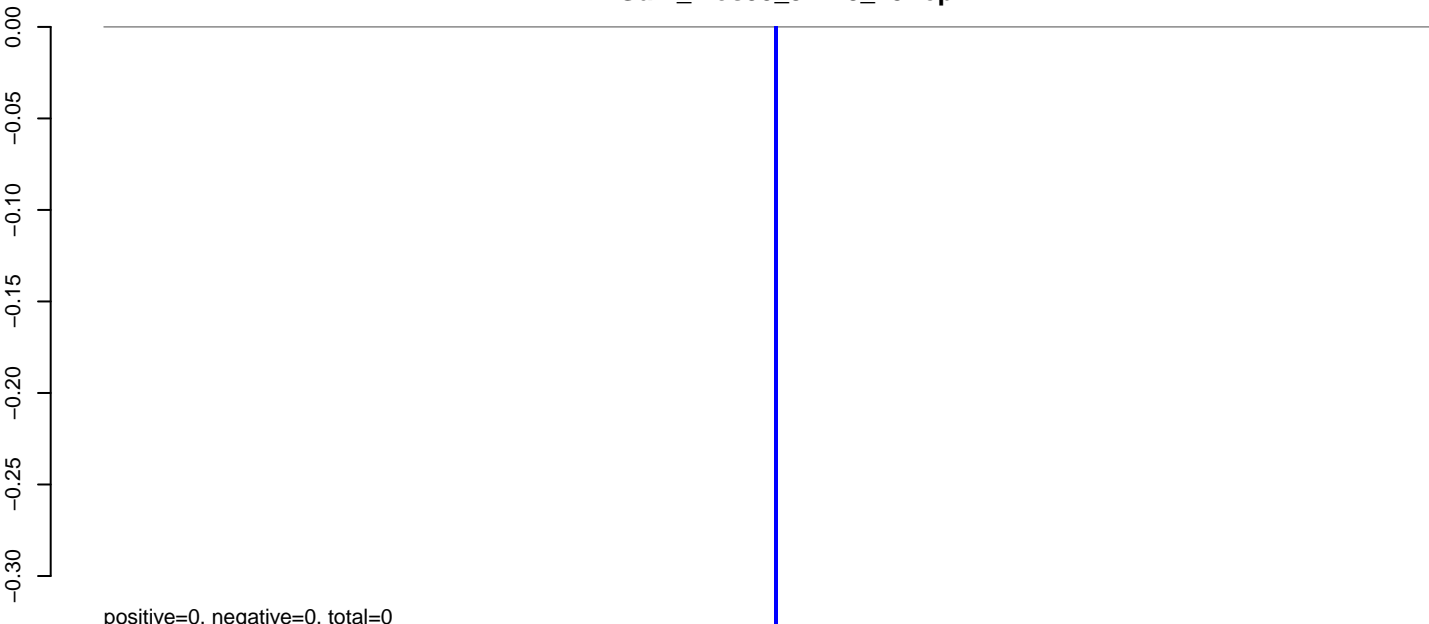
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12052-gene-2-HM147991.1-Trocaravirus\_non\_structural\_polyprotein\_gene\_partial\_cds\_and\_structural\_polyprotein\_gene:1-12052

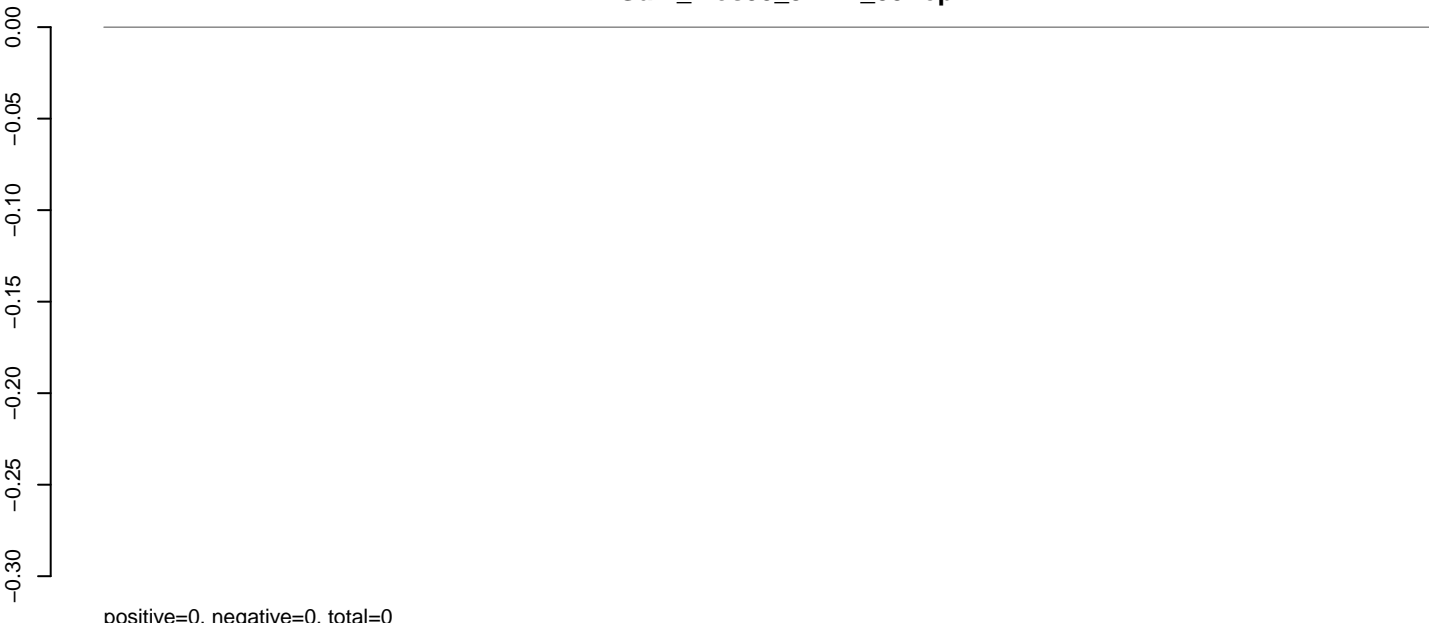
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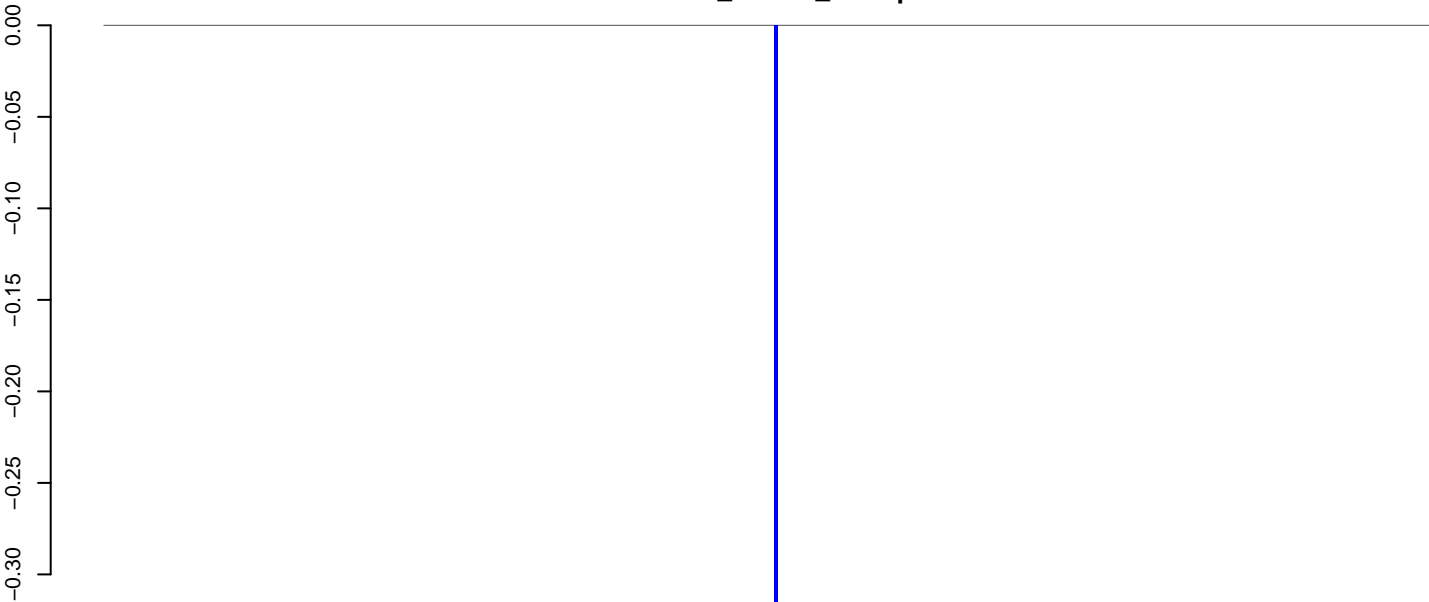
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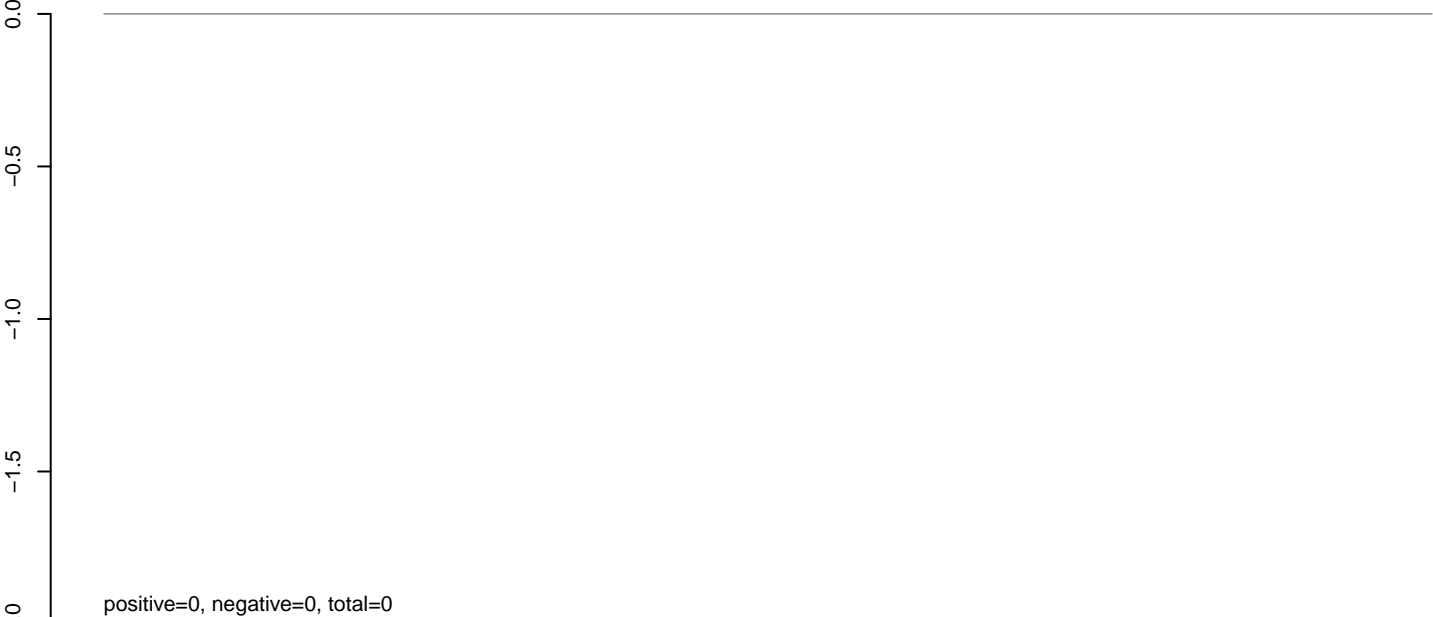
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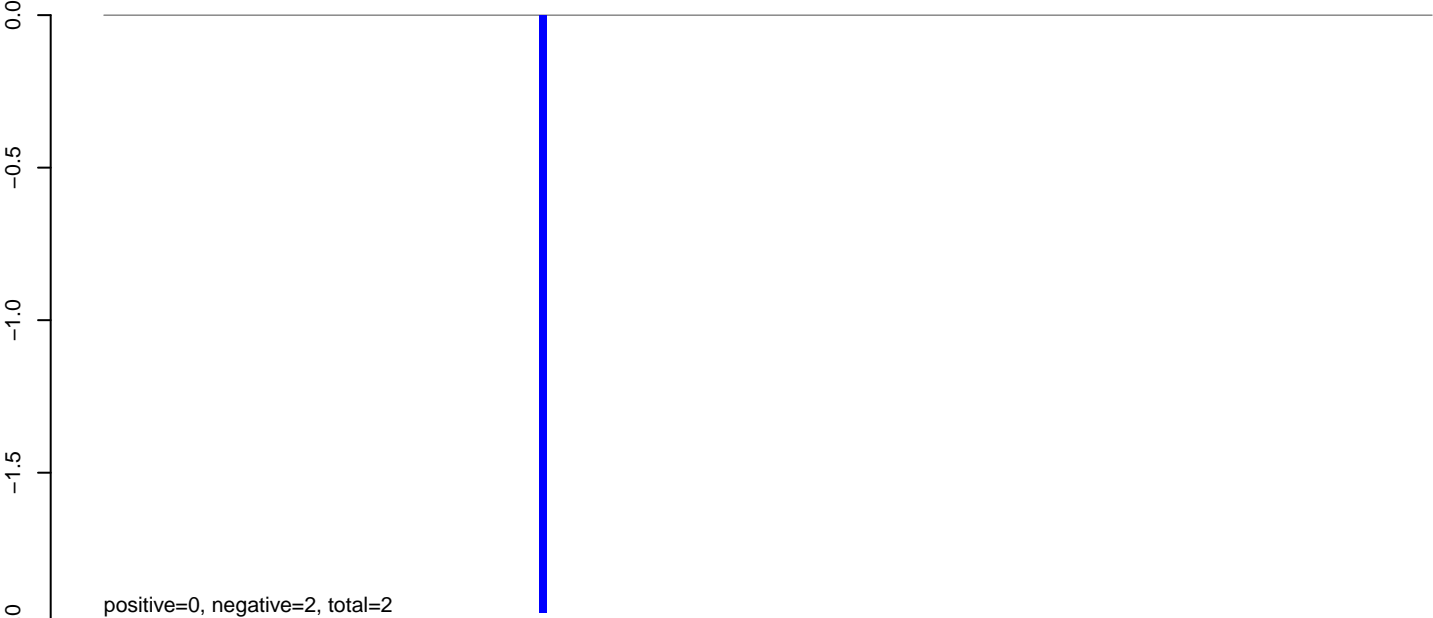
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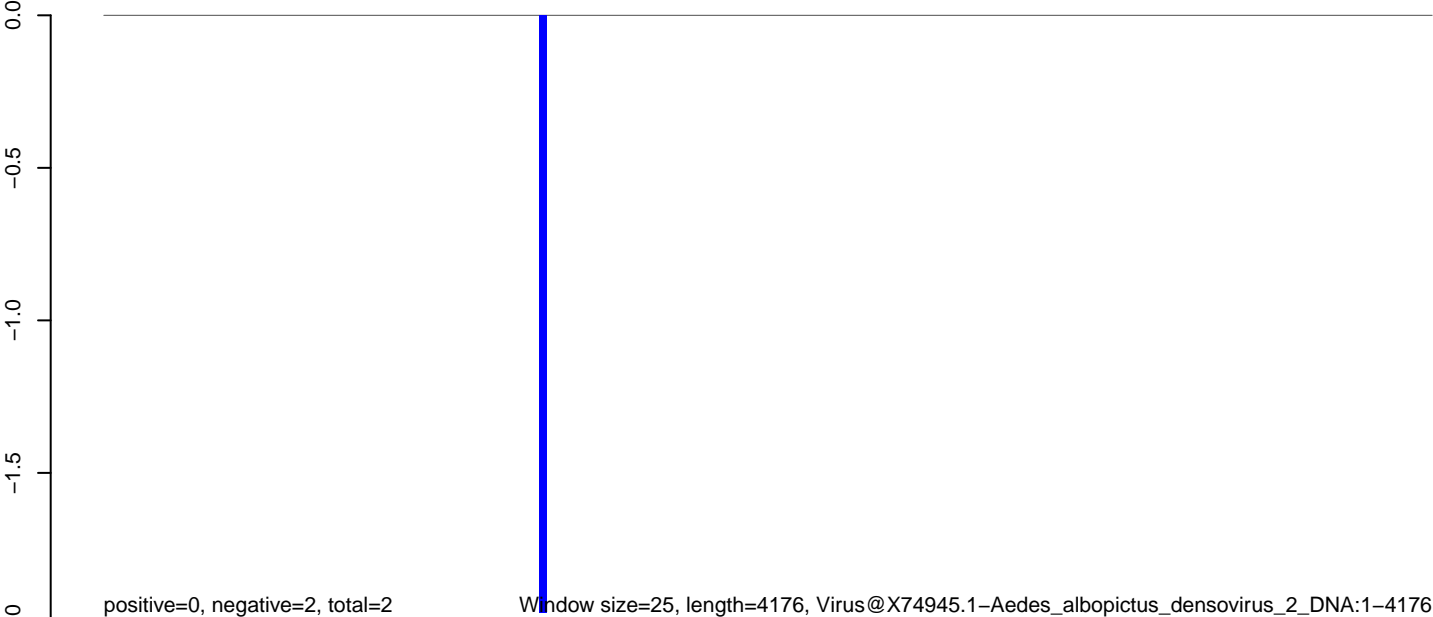
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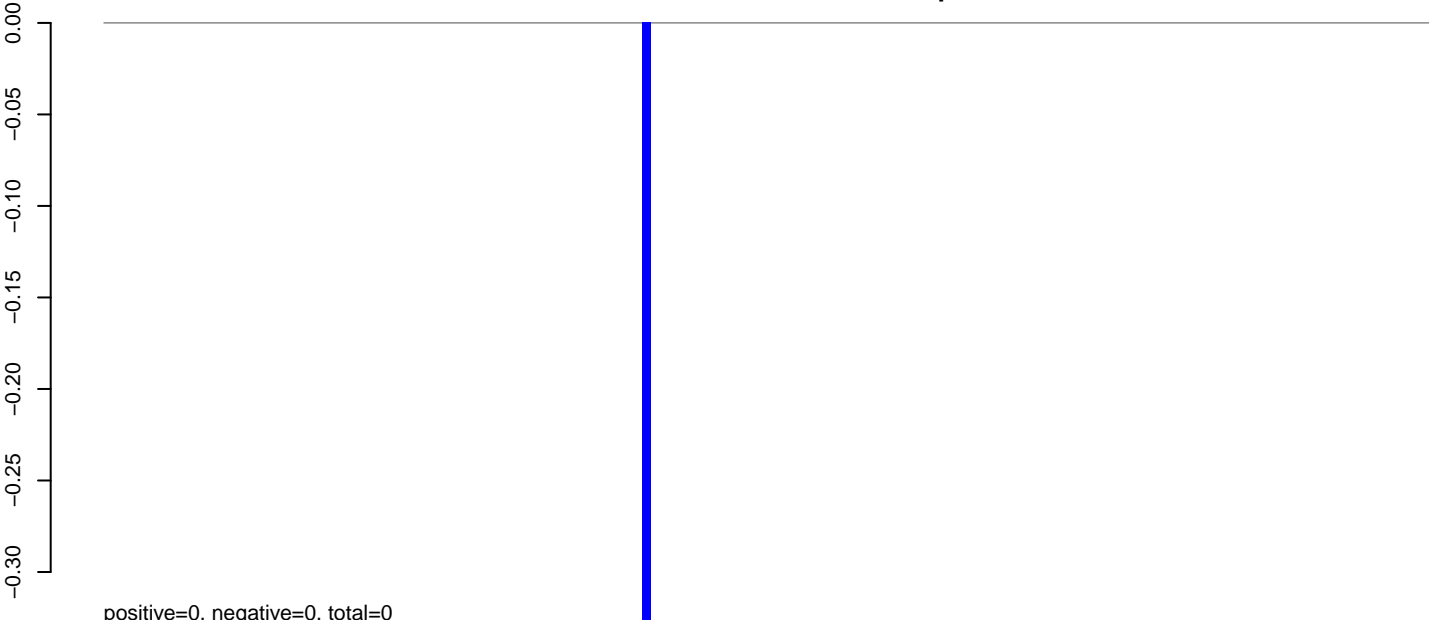


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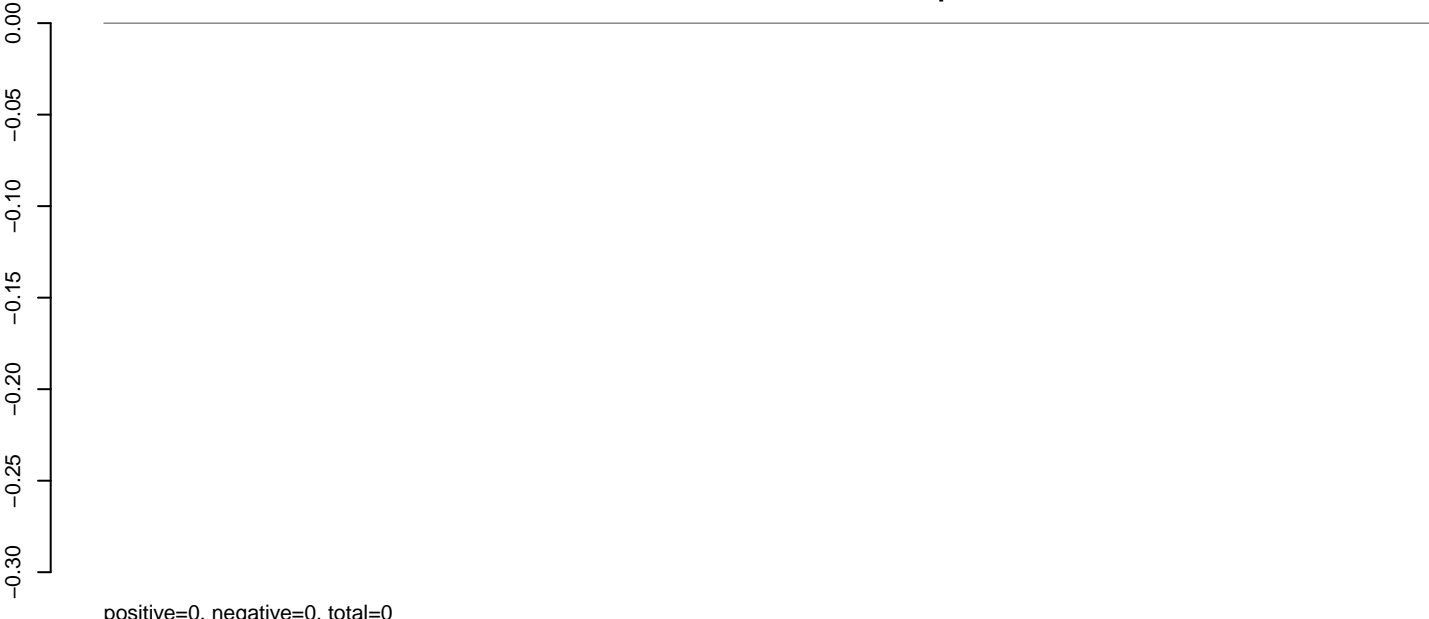




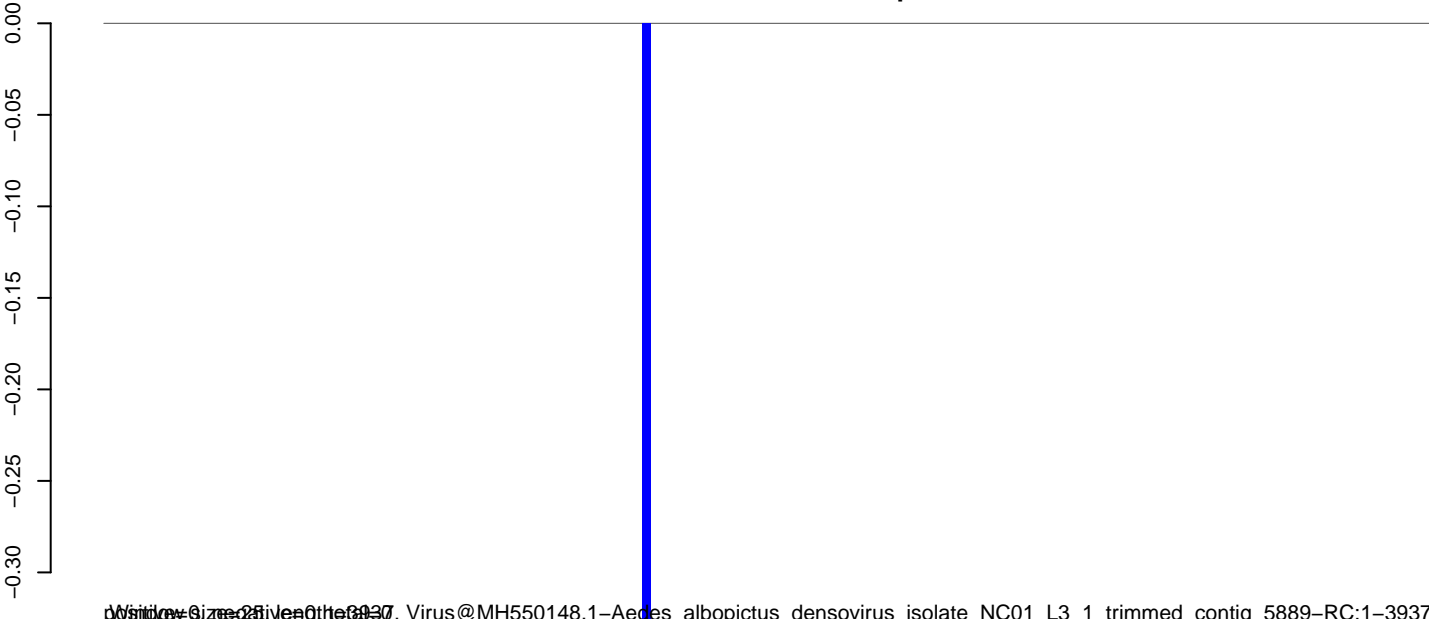
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AnGam\_Mos55\_JR.rep



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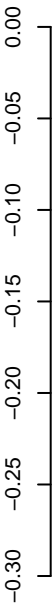
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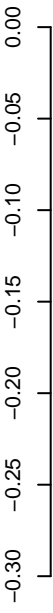
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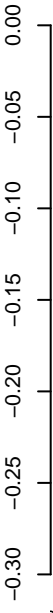
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AnGam\_Mos55\_JR.rep



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AnGam\_Mos55\_JR.24\_35.rep



AnGam\_Mos55\_JR.rep



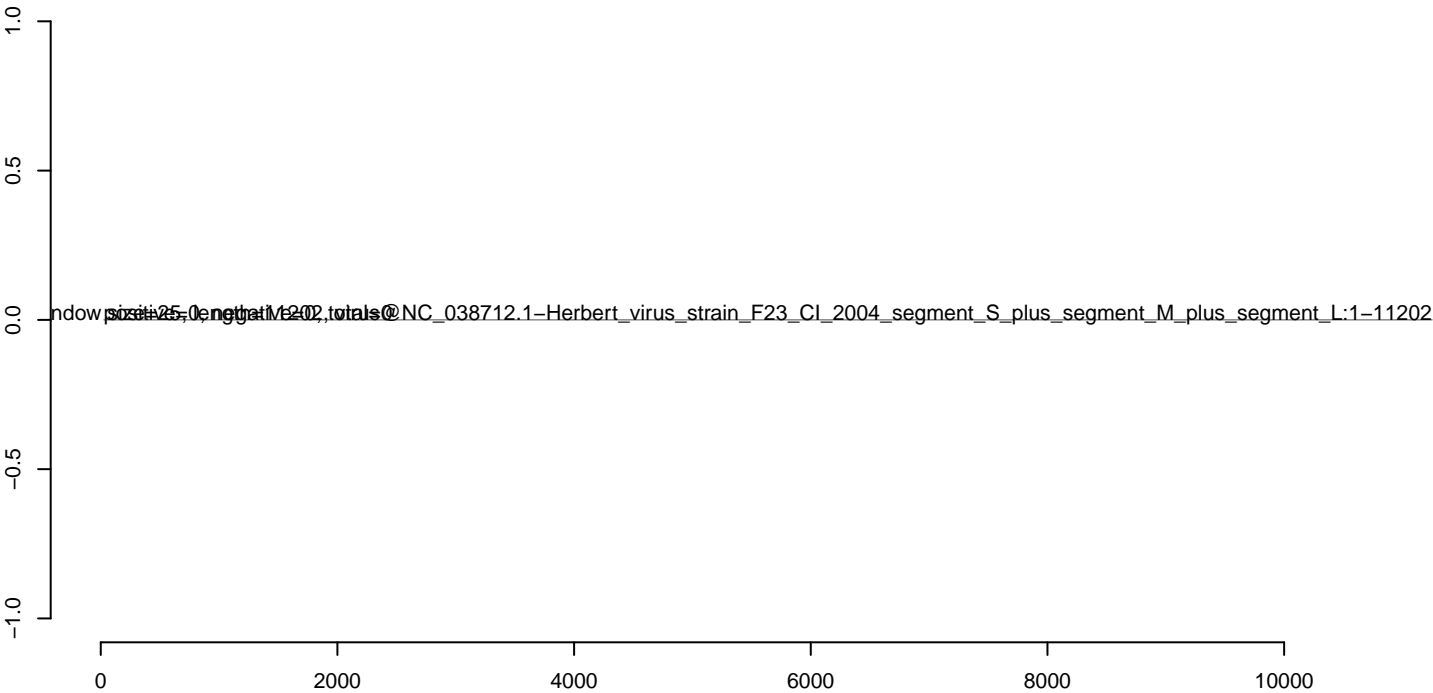
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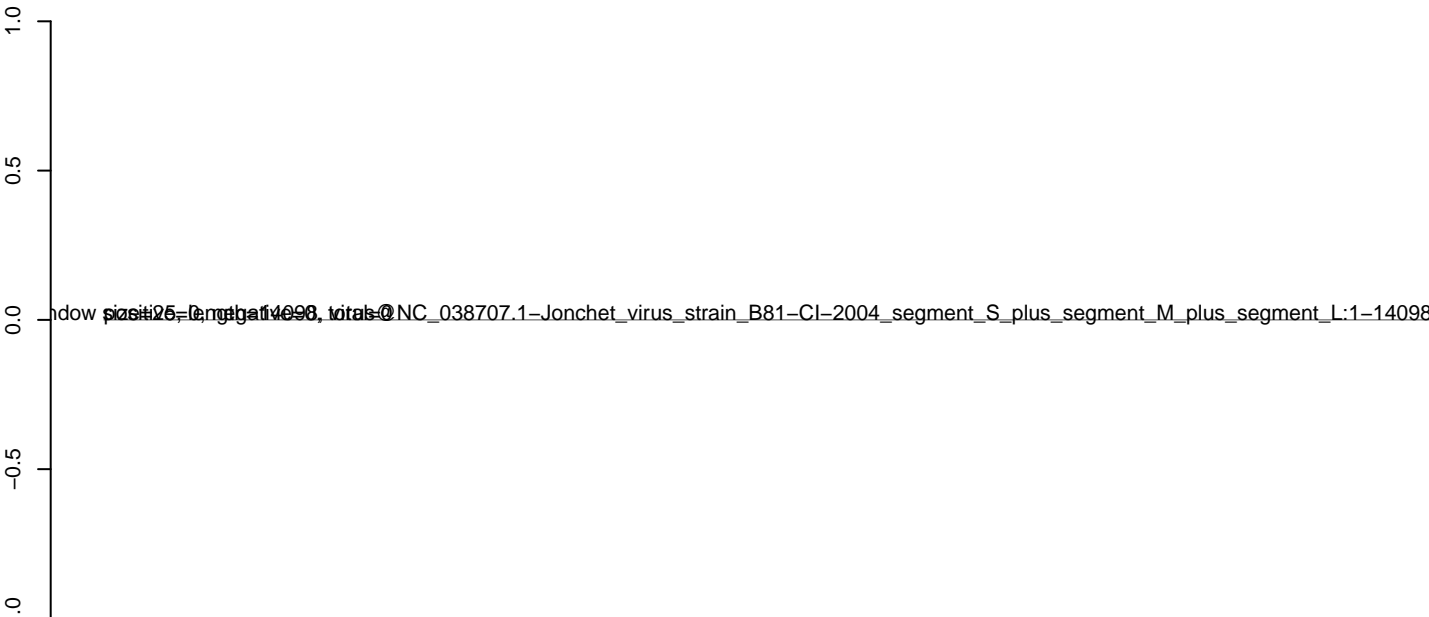
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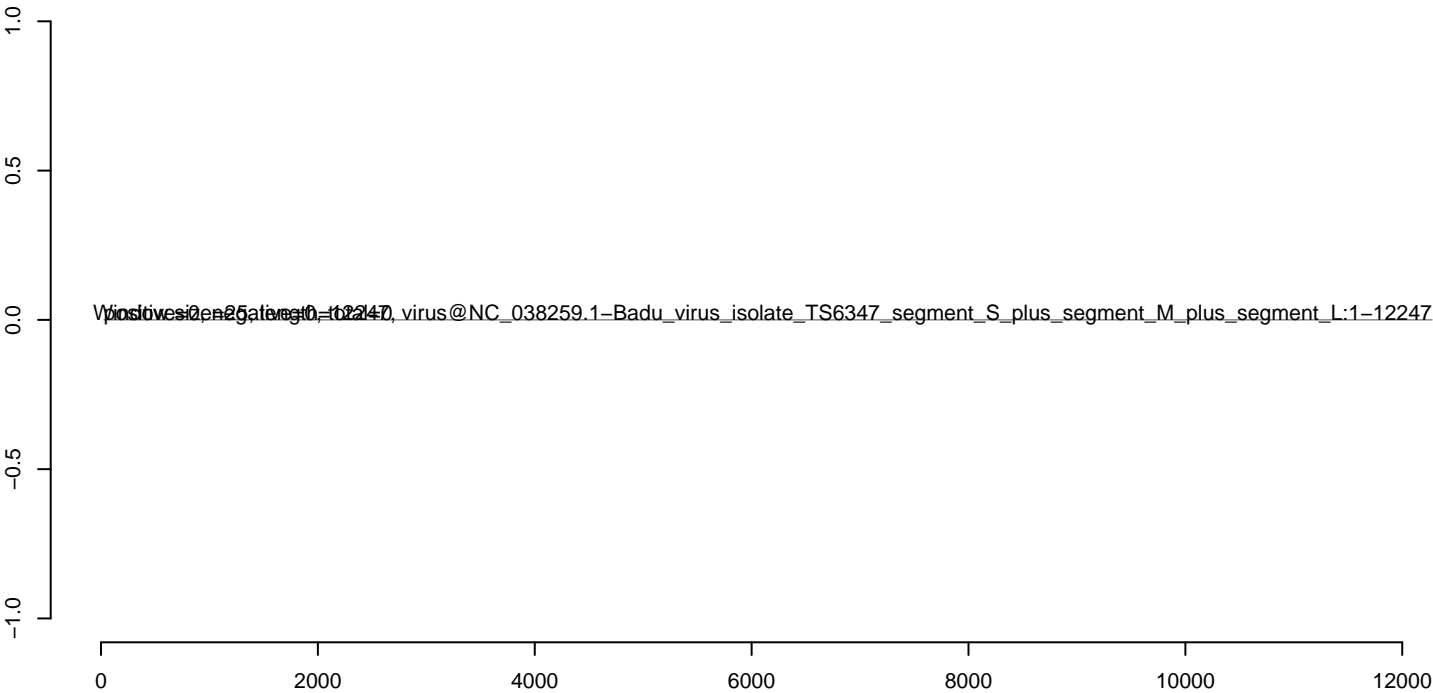
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AnGam\_Mos55\_JR.24\_35.rep



AnGam\_Mos55\_JR.rep





**AnGam\_Mos55\_JR.18\_23.rep**



**AnGam\_Mos55\_JR.24\_35.rep**



**AnGam\_Mos55\_JR.rep**



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AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



AnGam\_Mos55\_JR.rep



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AnGam\_Mos55\_JR.rep



AnGam\_Mos55\_JR.18\_23.rep



AnGam\_Mos55\_JR.24\_35.rep



AnGam\_Mos55\_JR.rep



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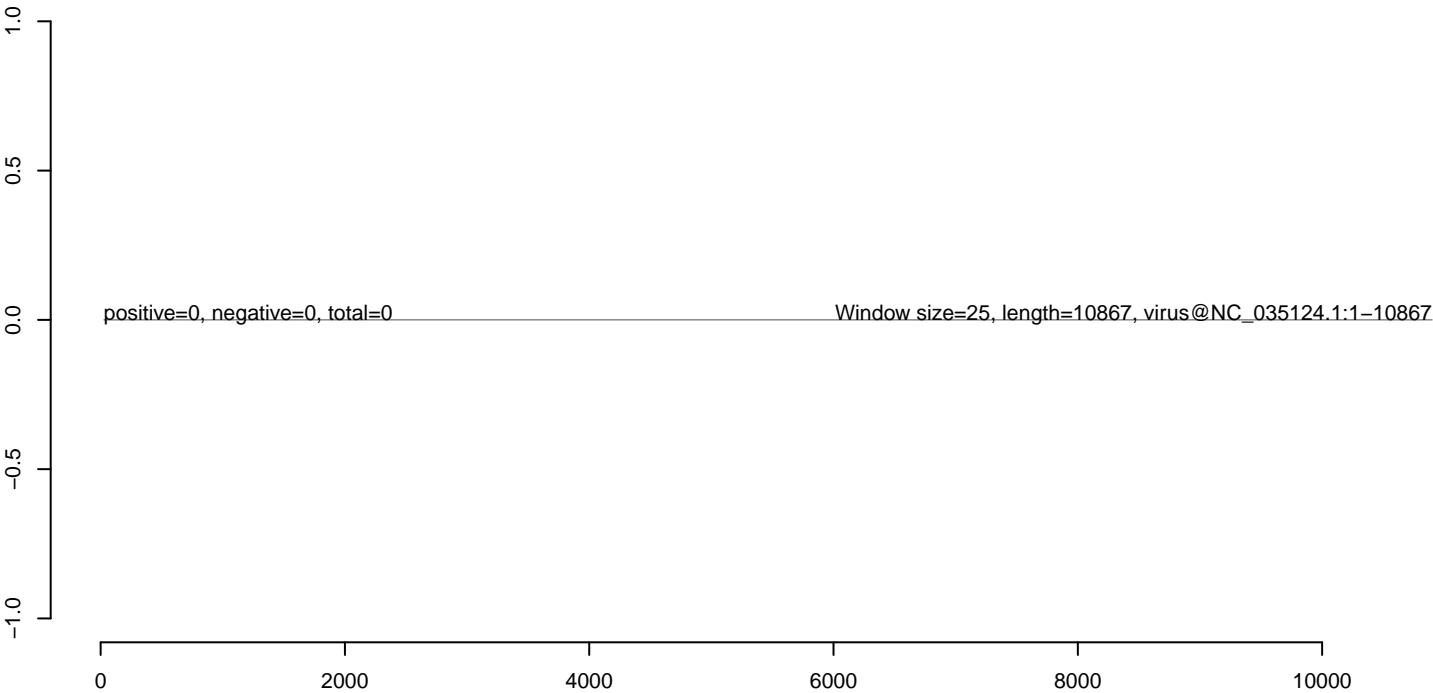
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AnGam\_Mos55\_JR.rep

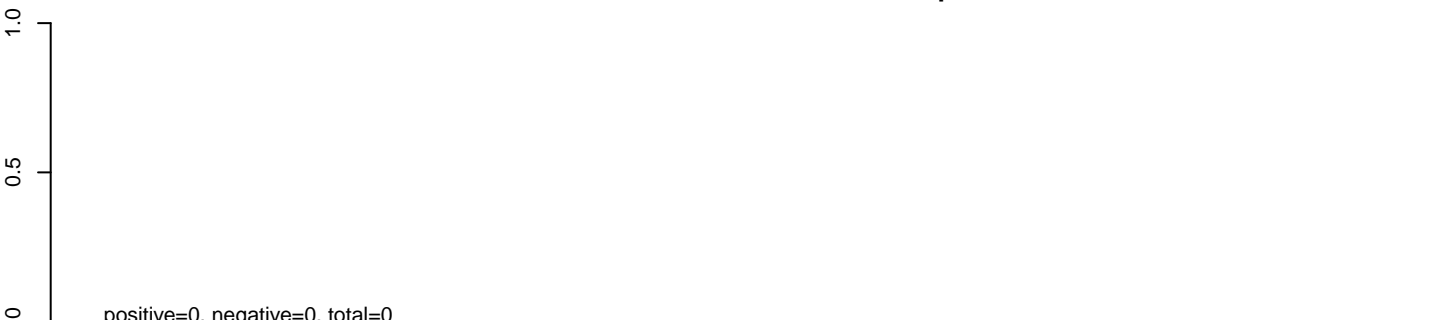


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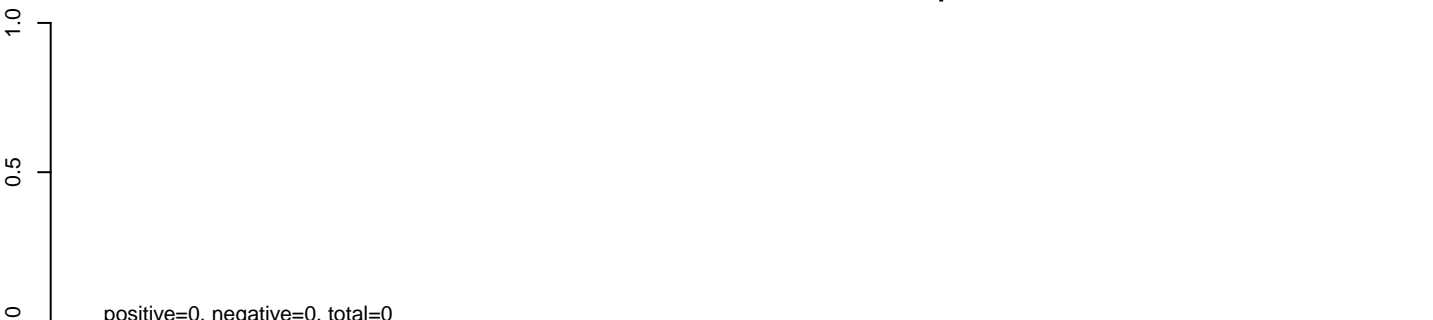


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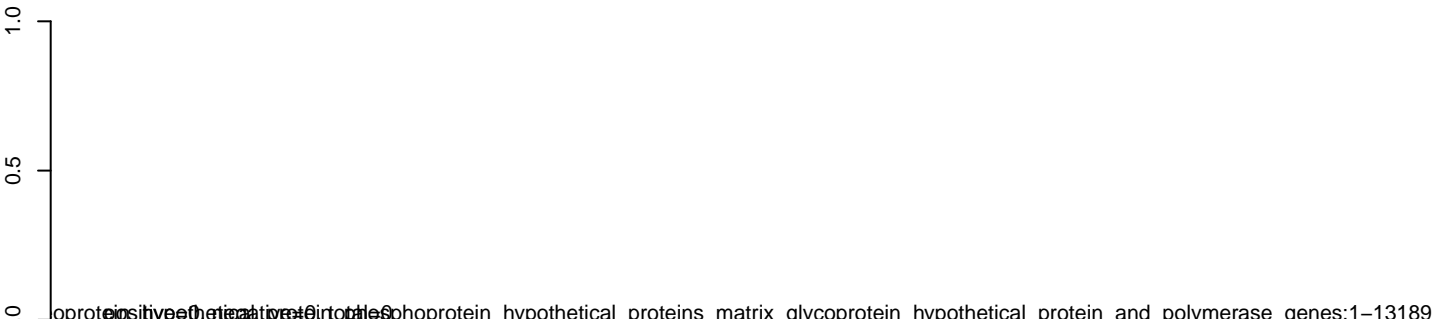
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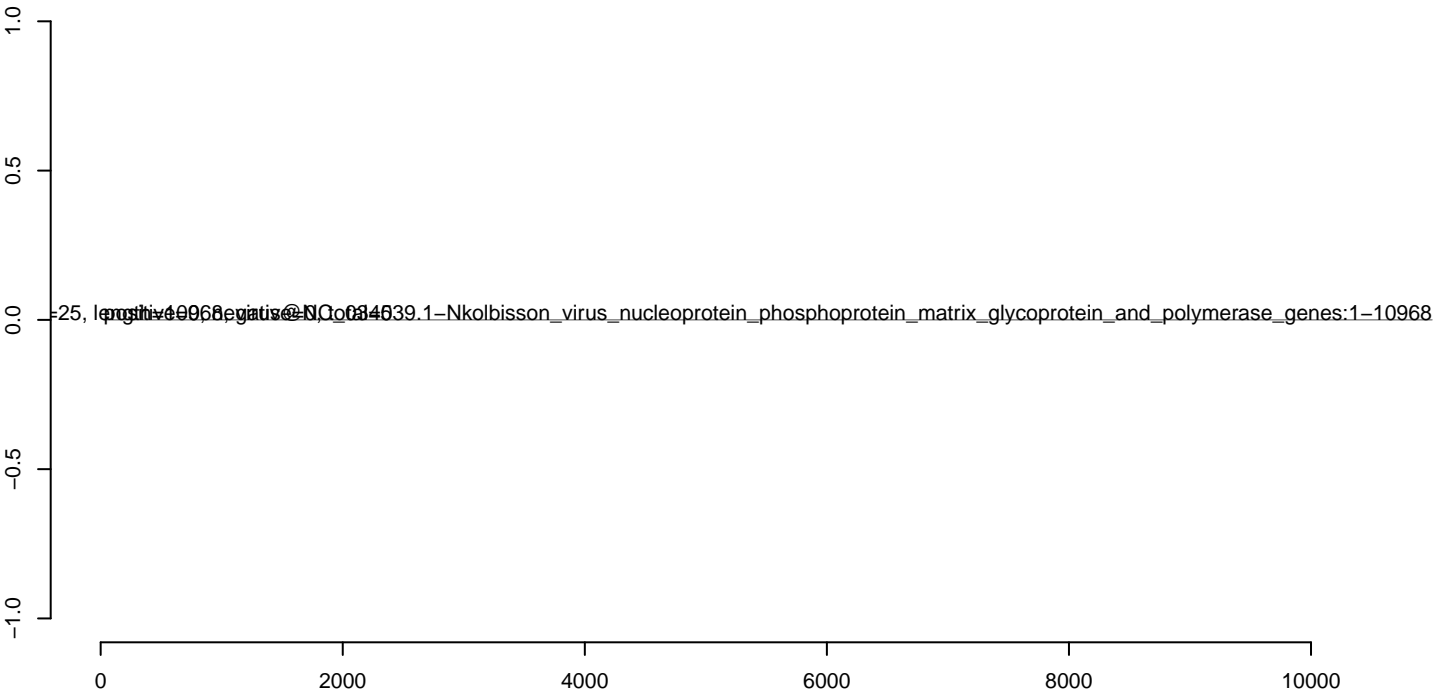
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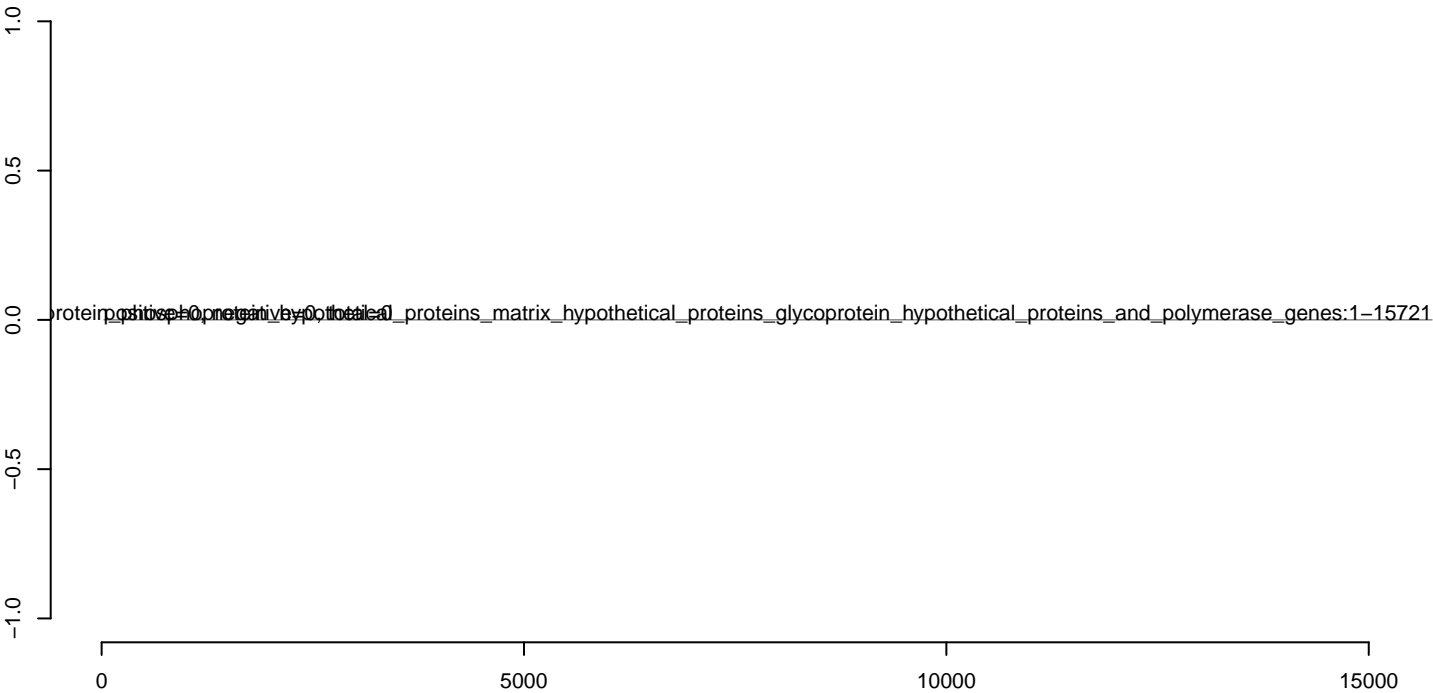
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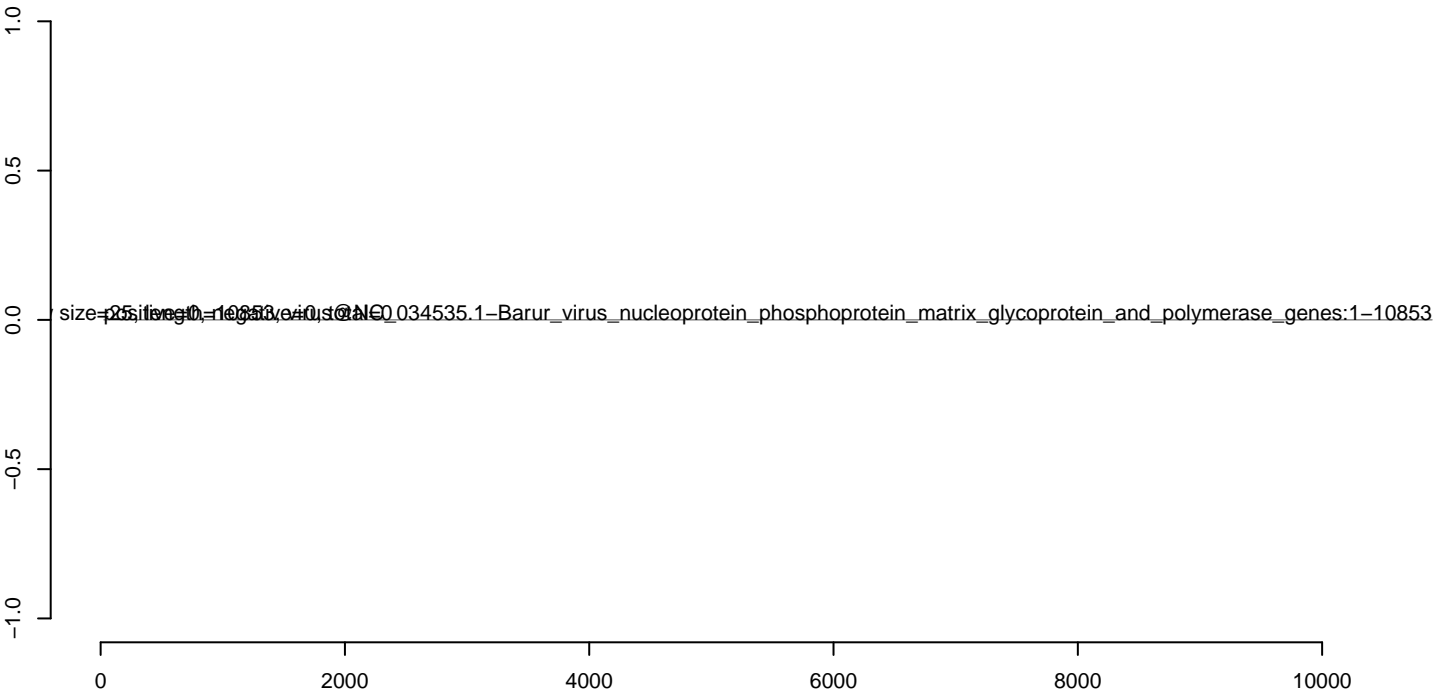
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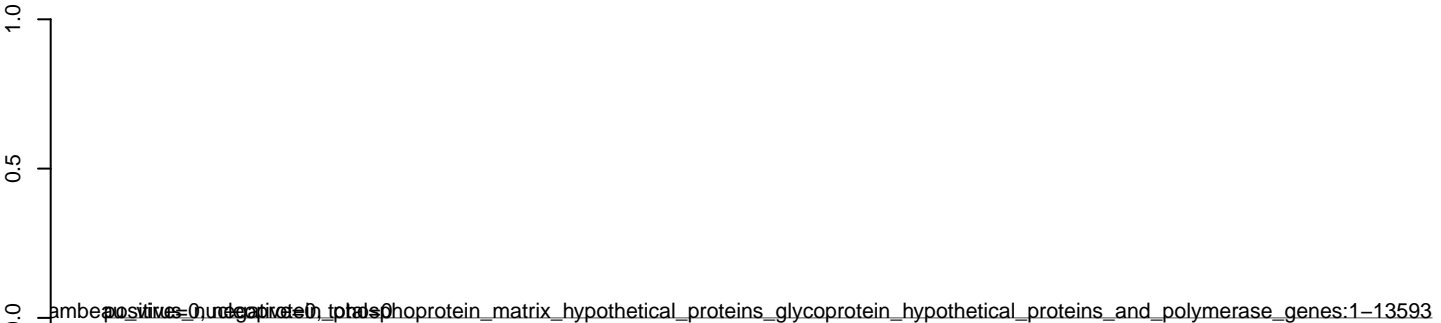
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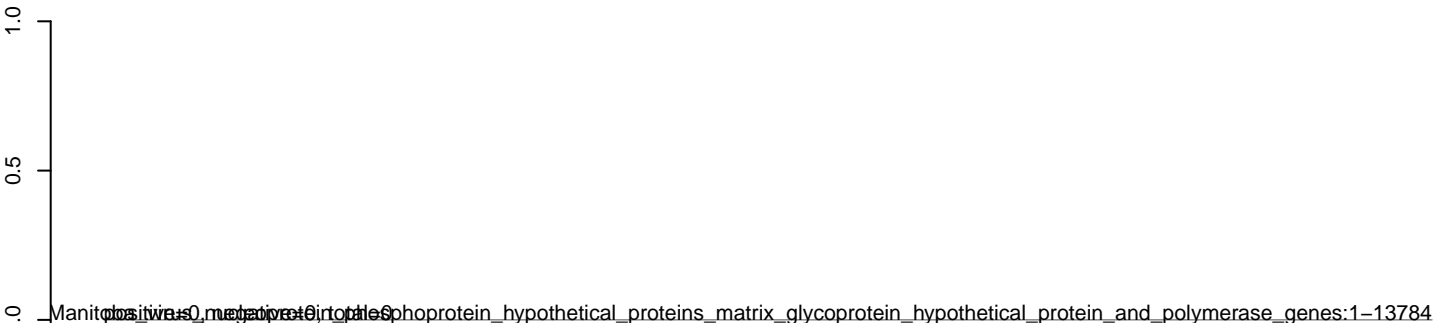
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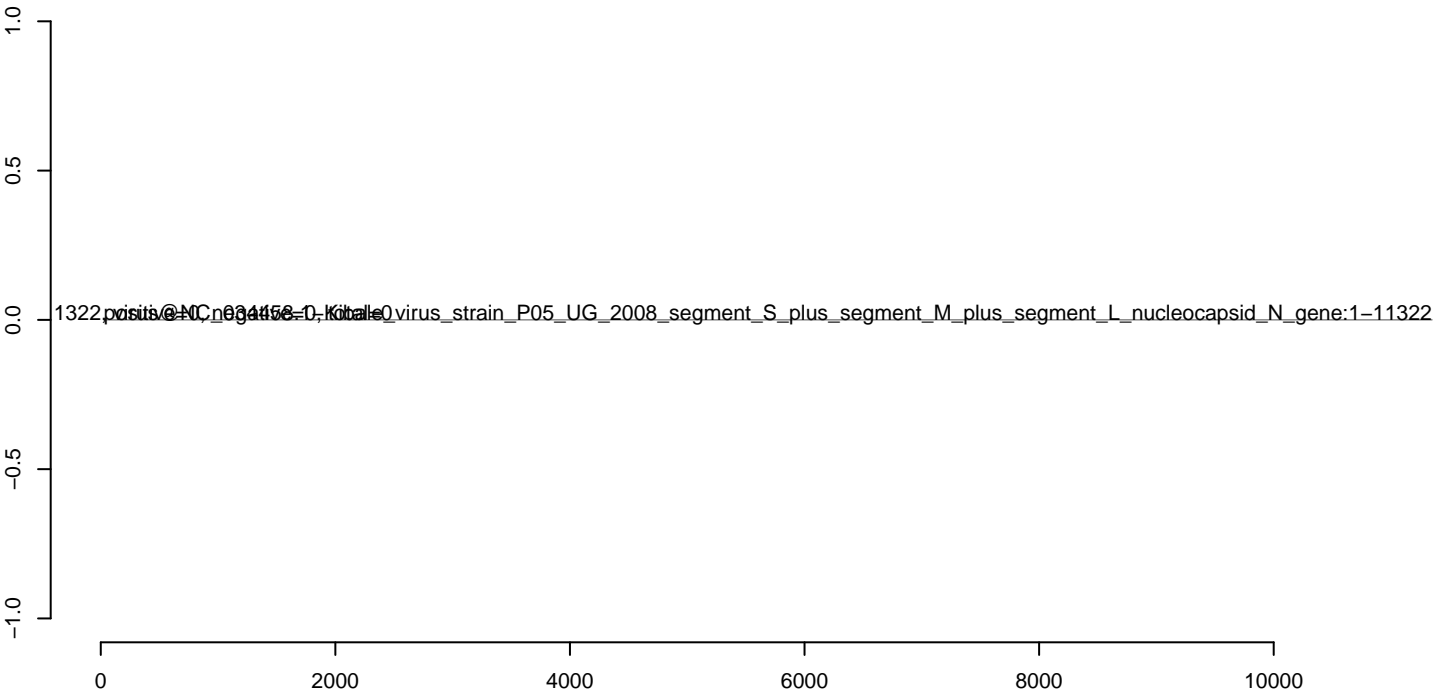
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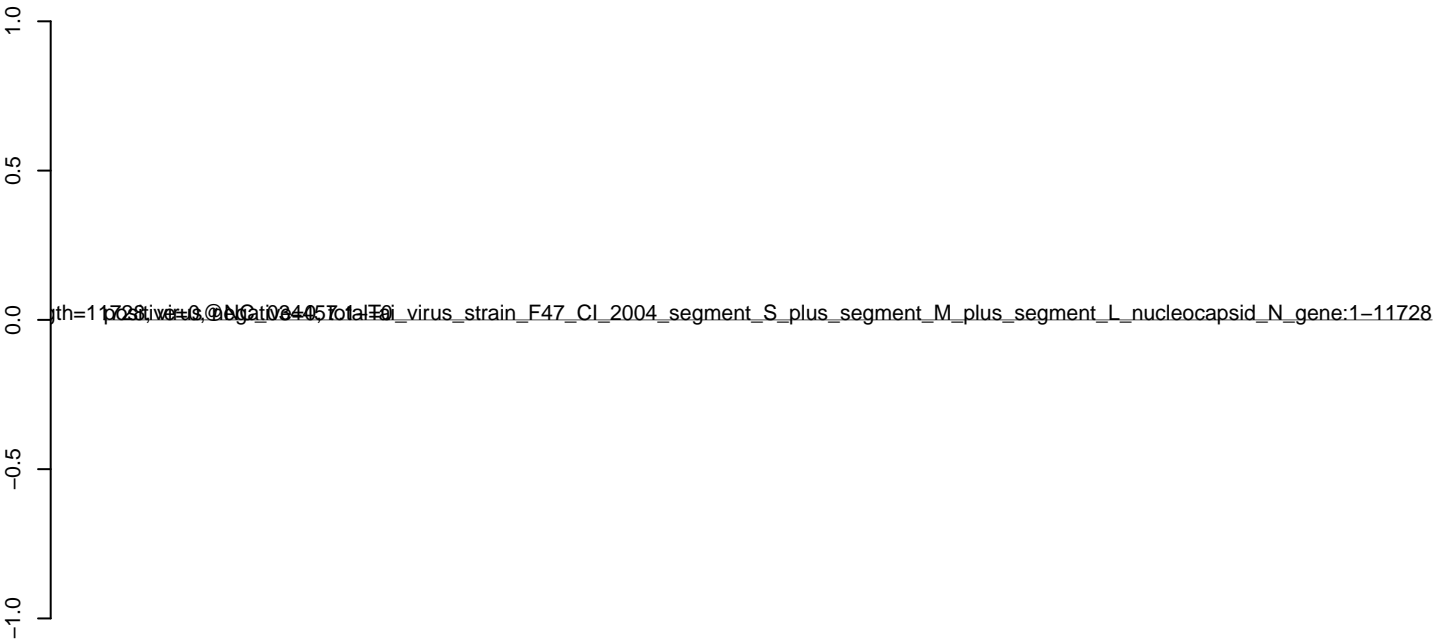
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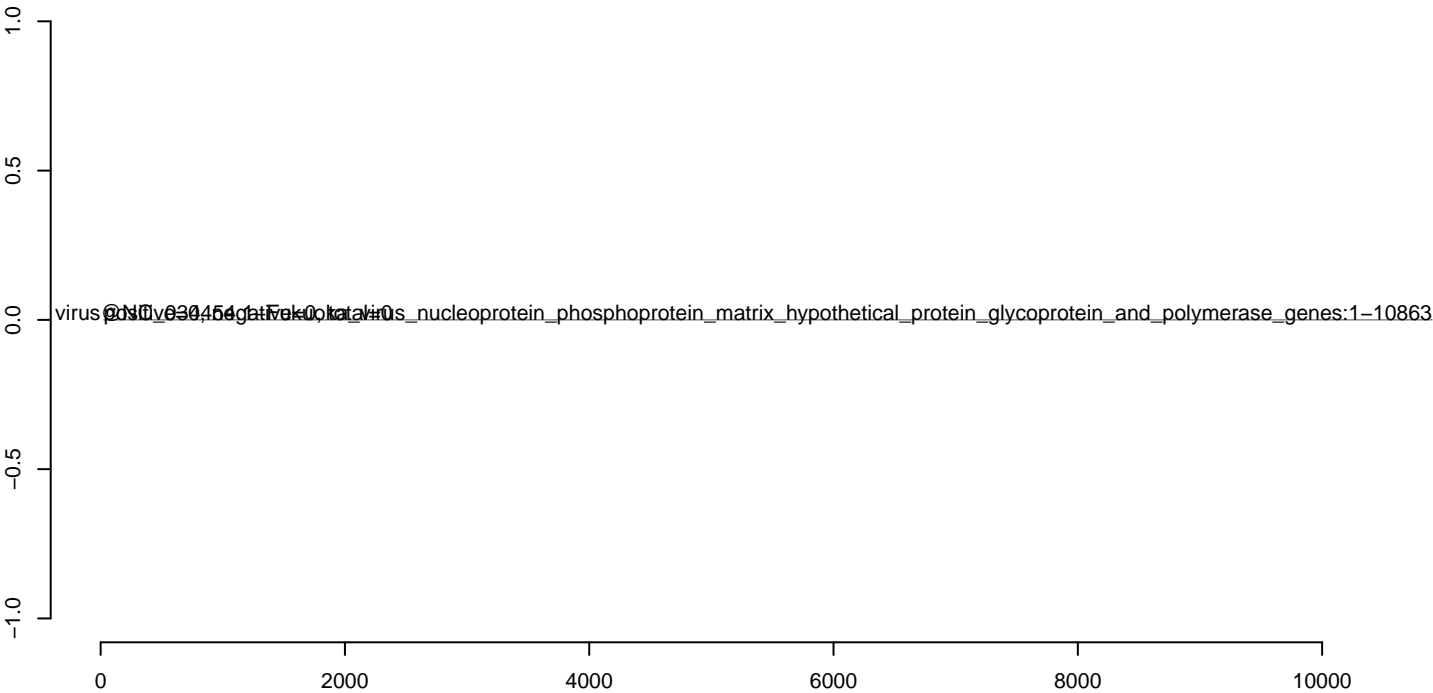
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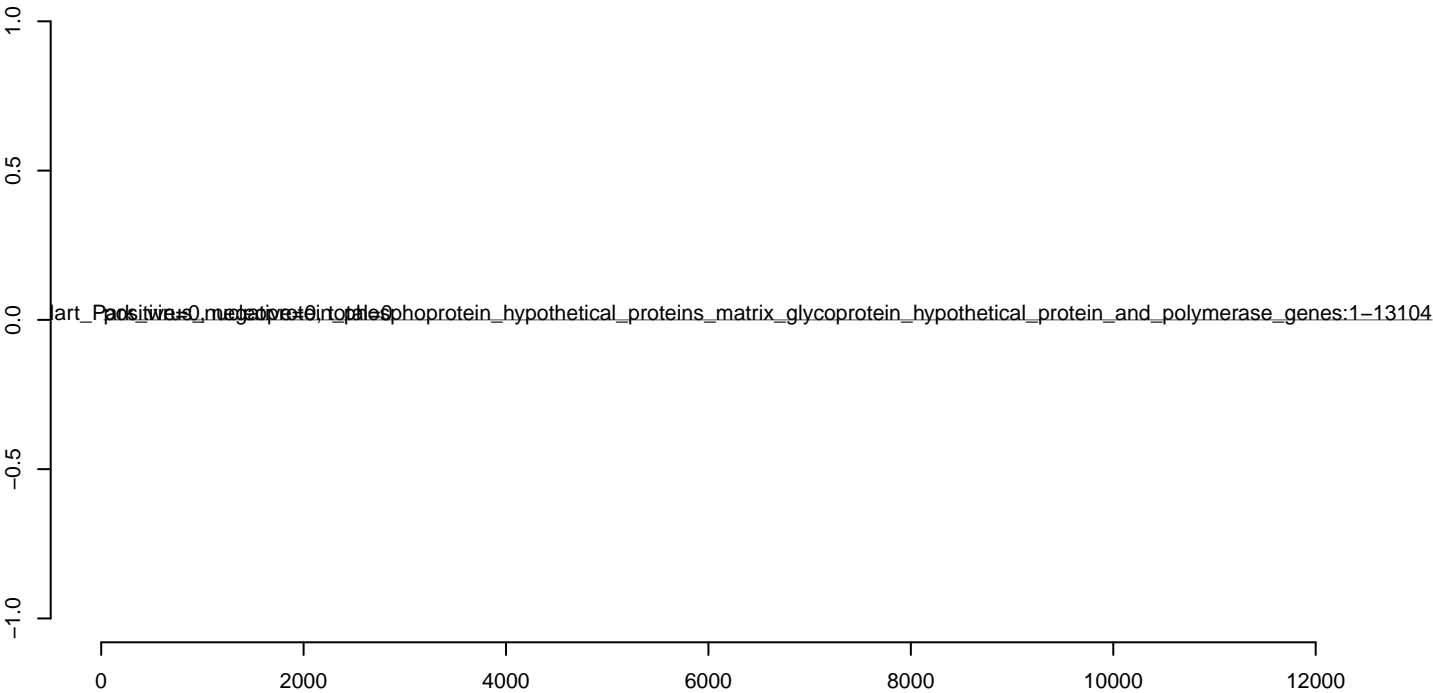
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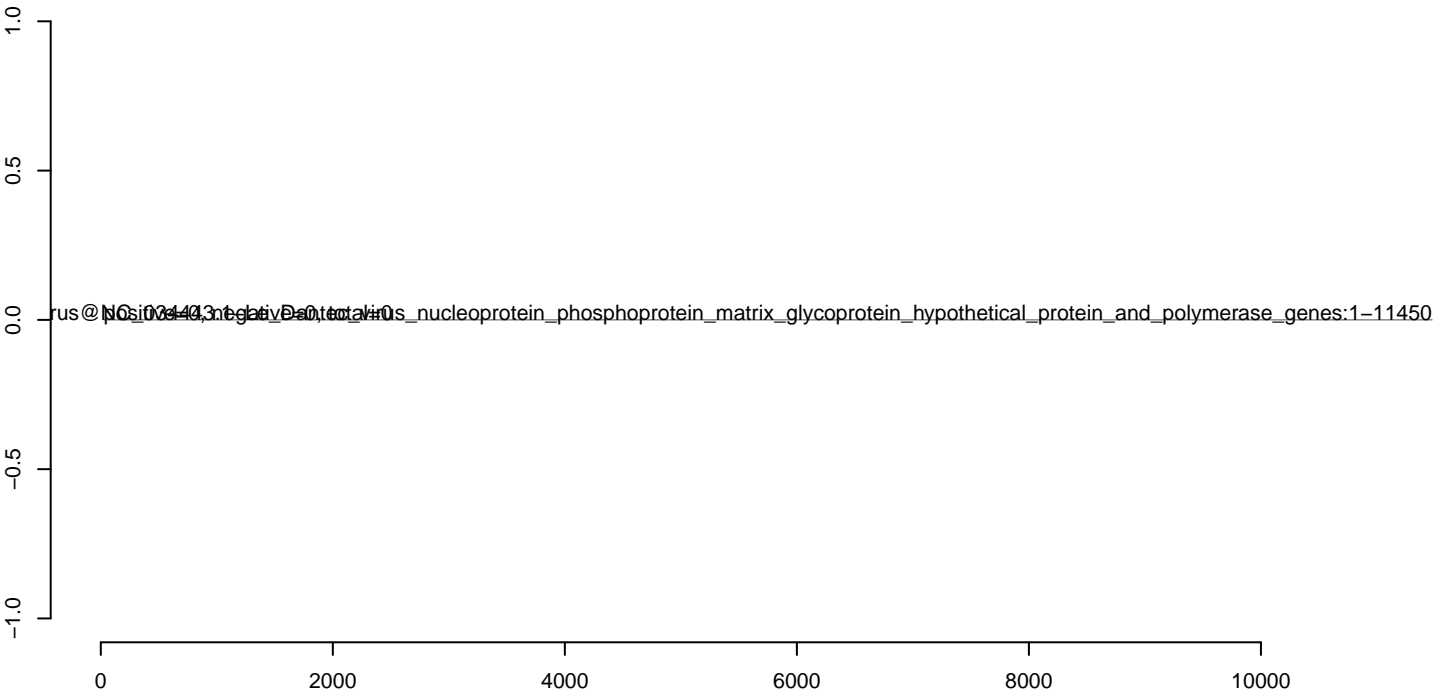
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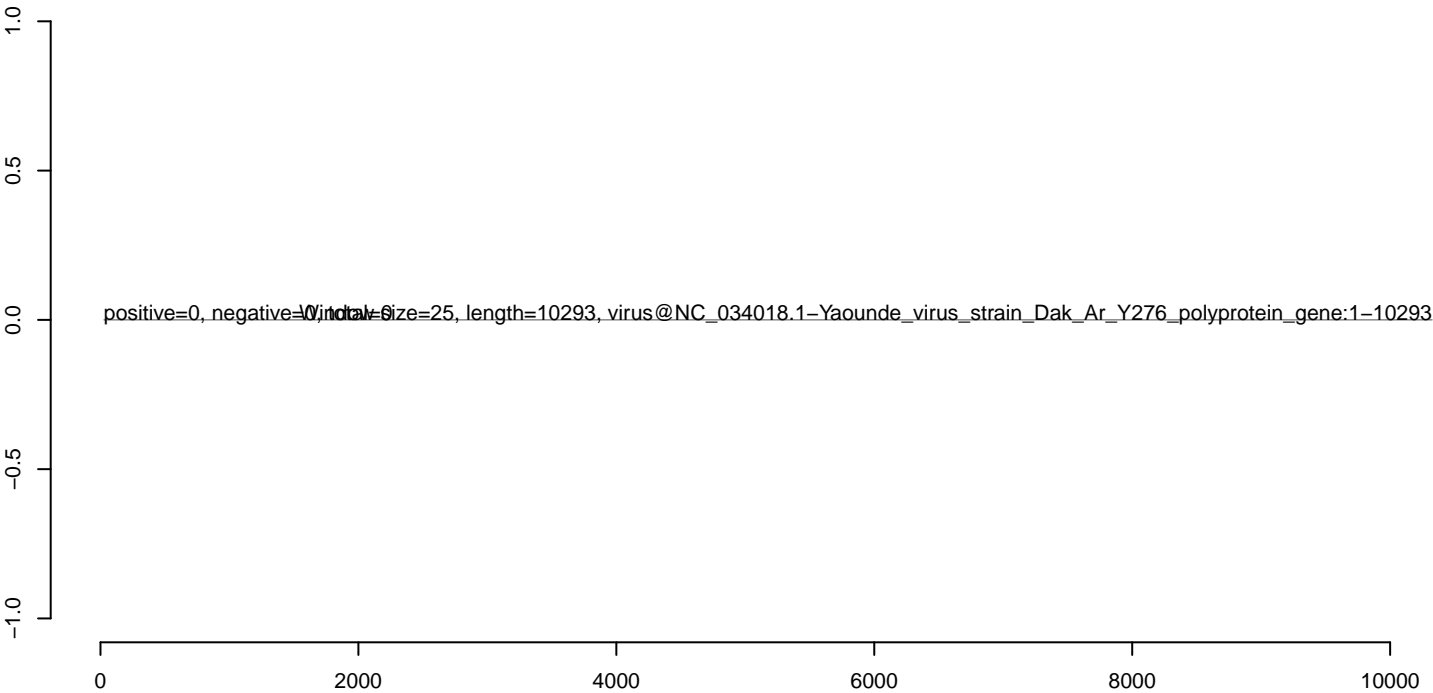
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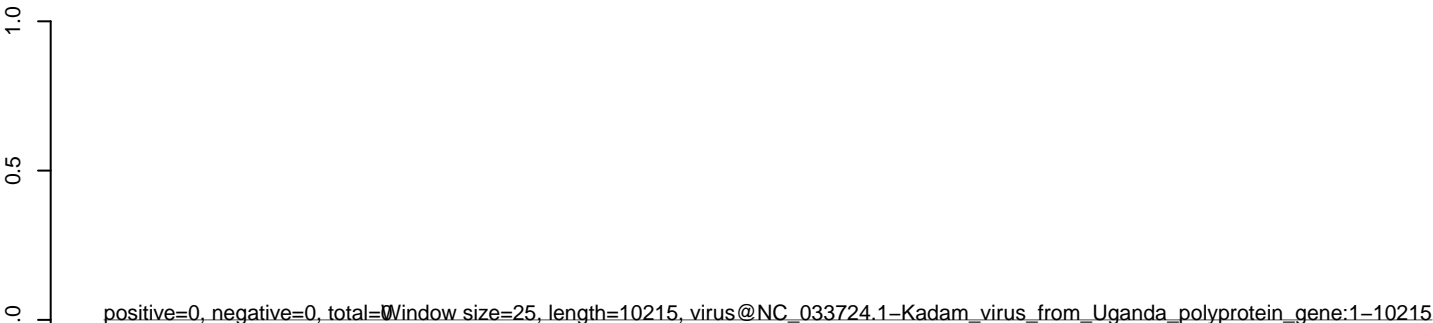
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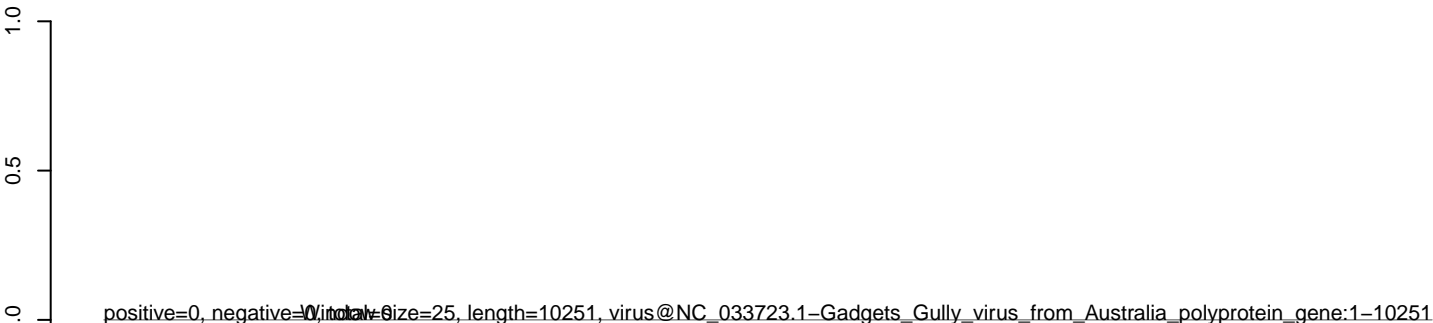
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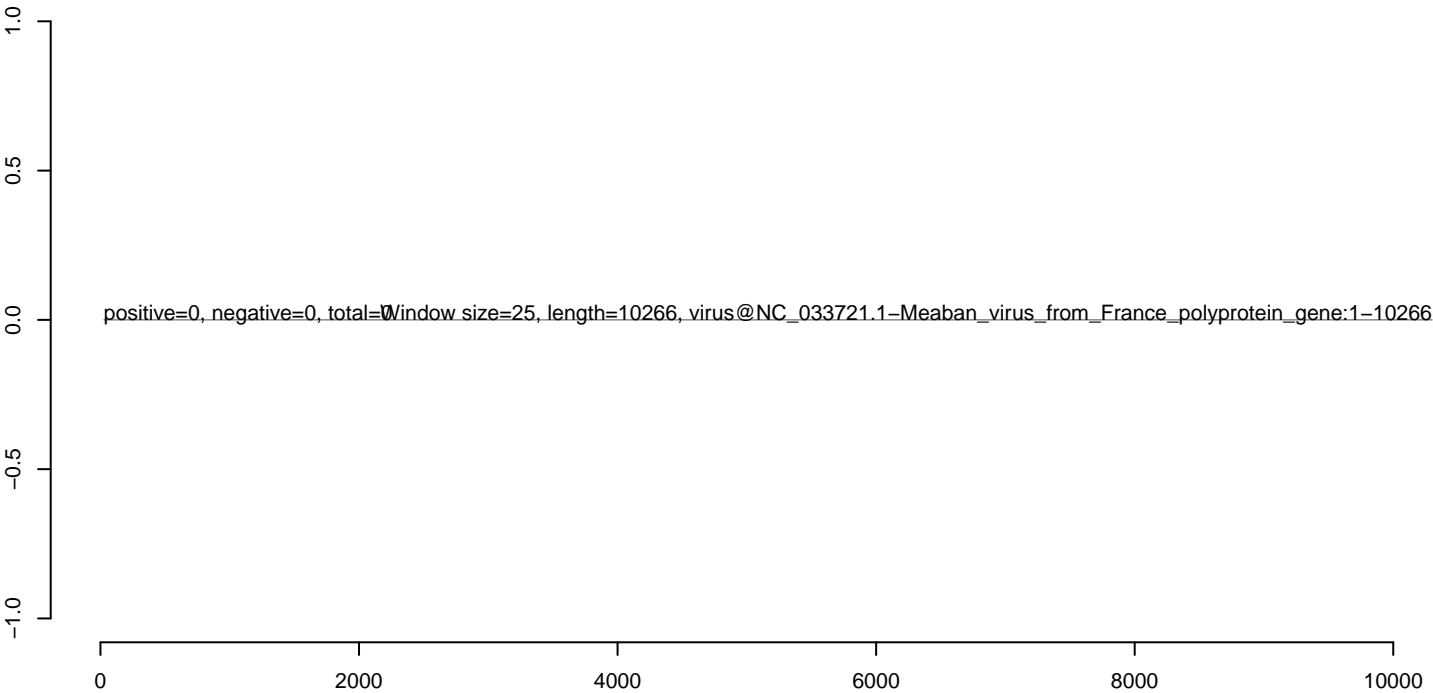
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AnGam\_Mos55\_JR.24\_35.rep



AnGam\_Mos55\_JR.rep



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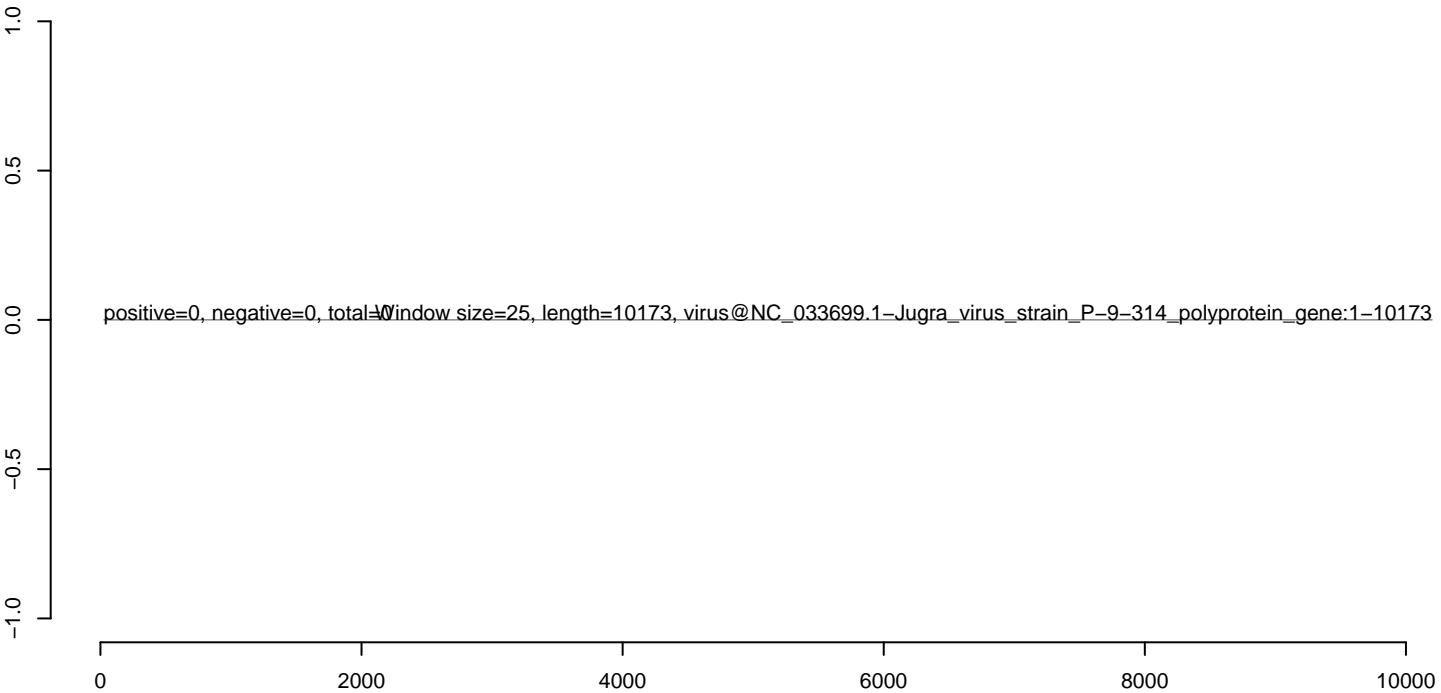
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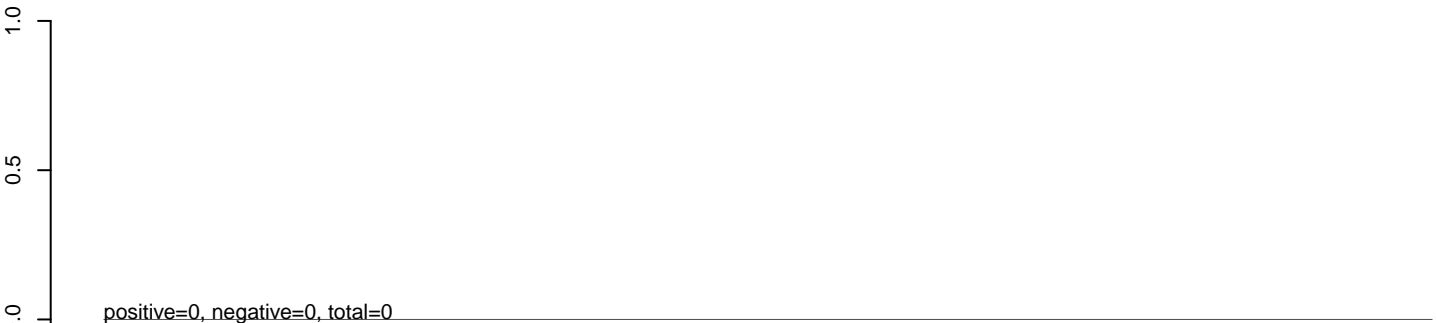
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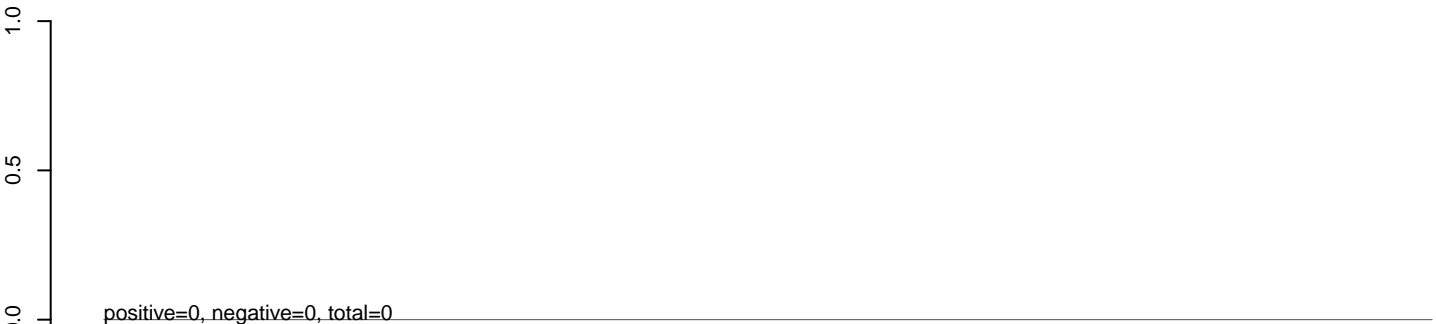
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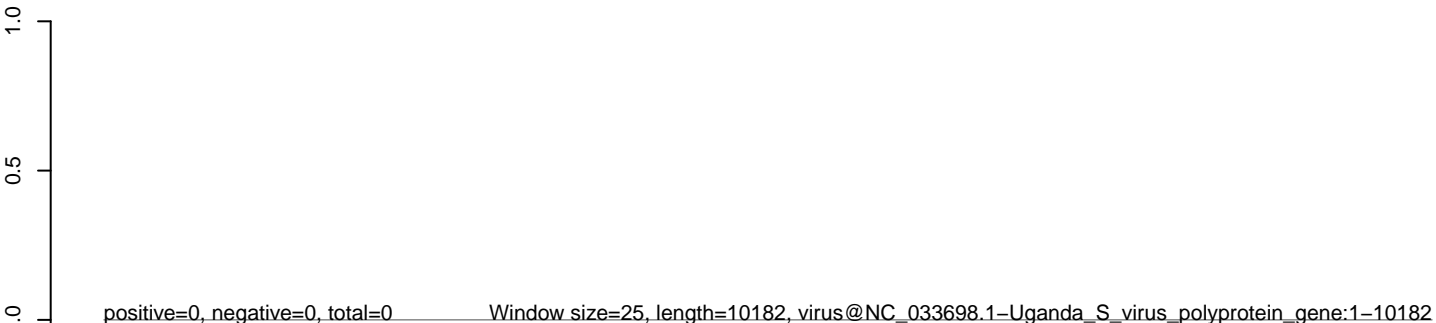
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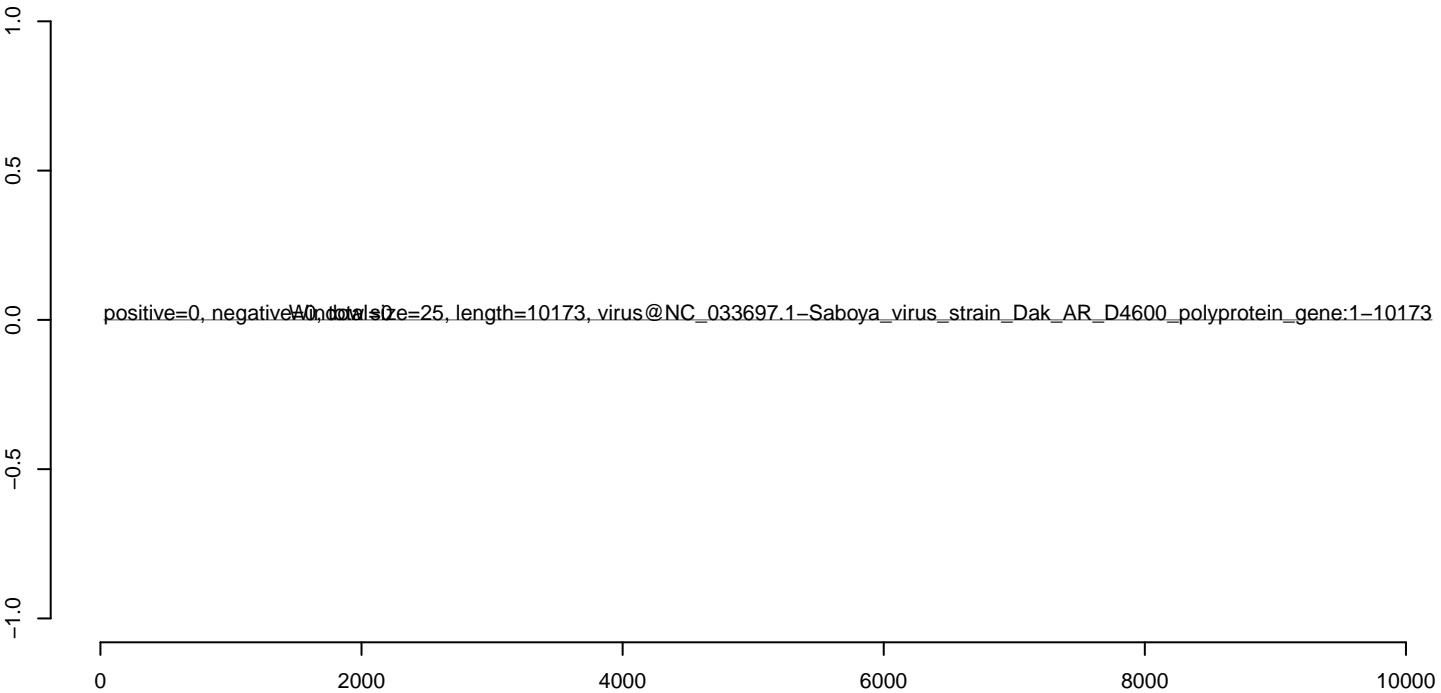
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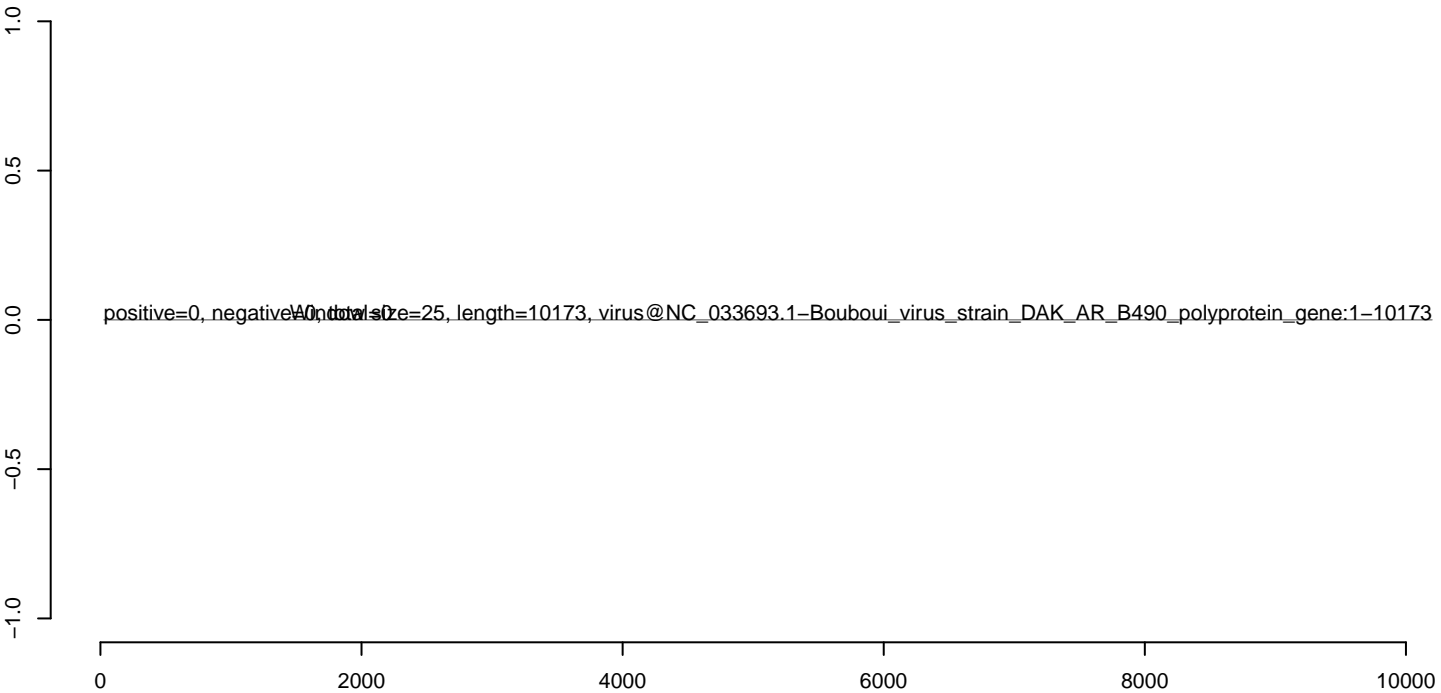
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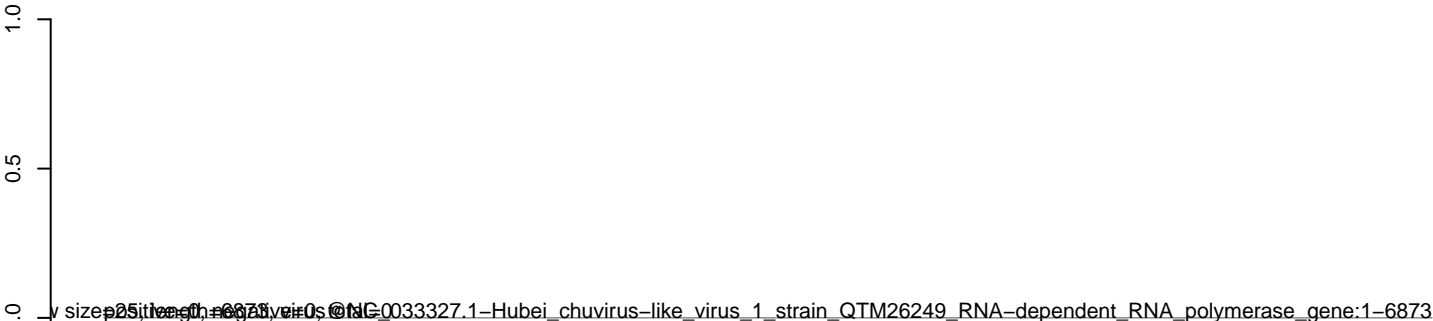
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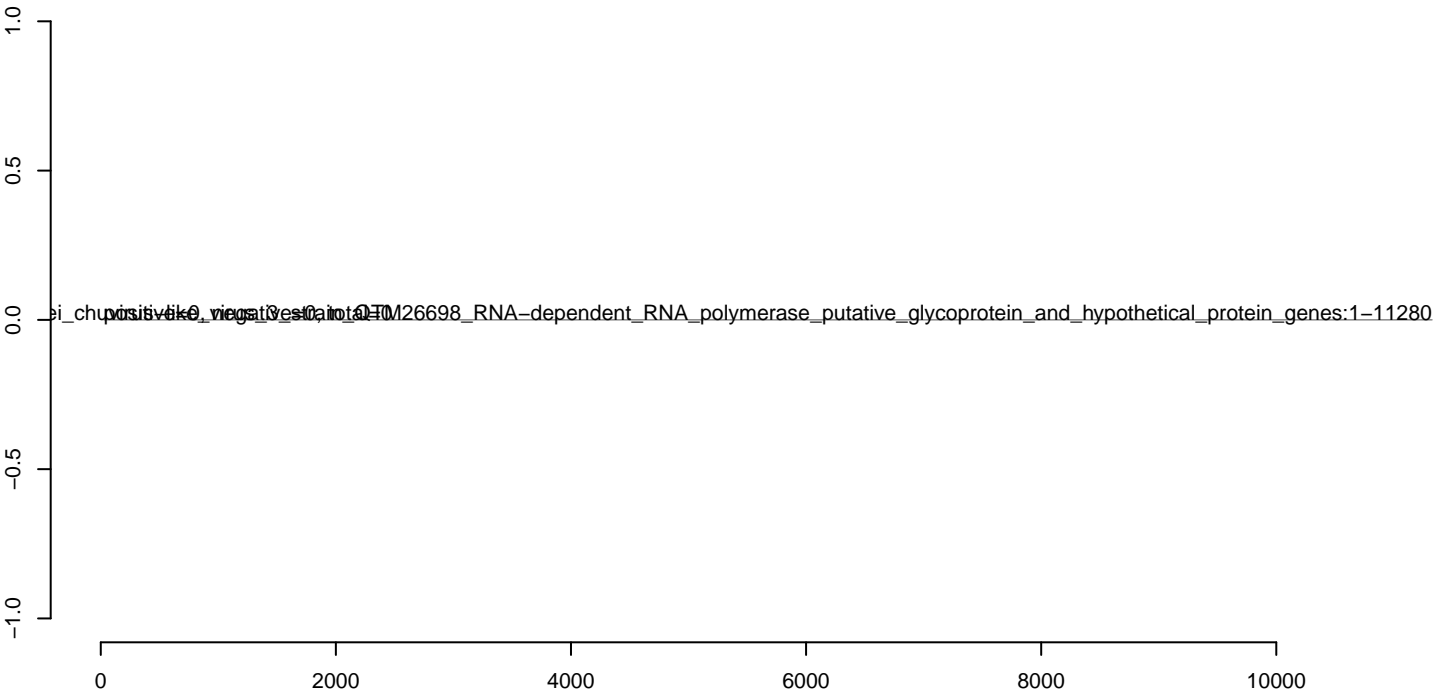
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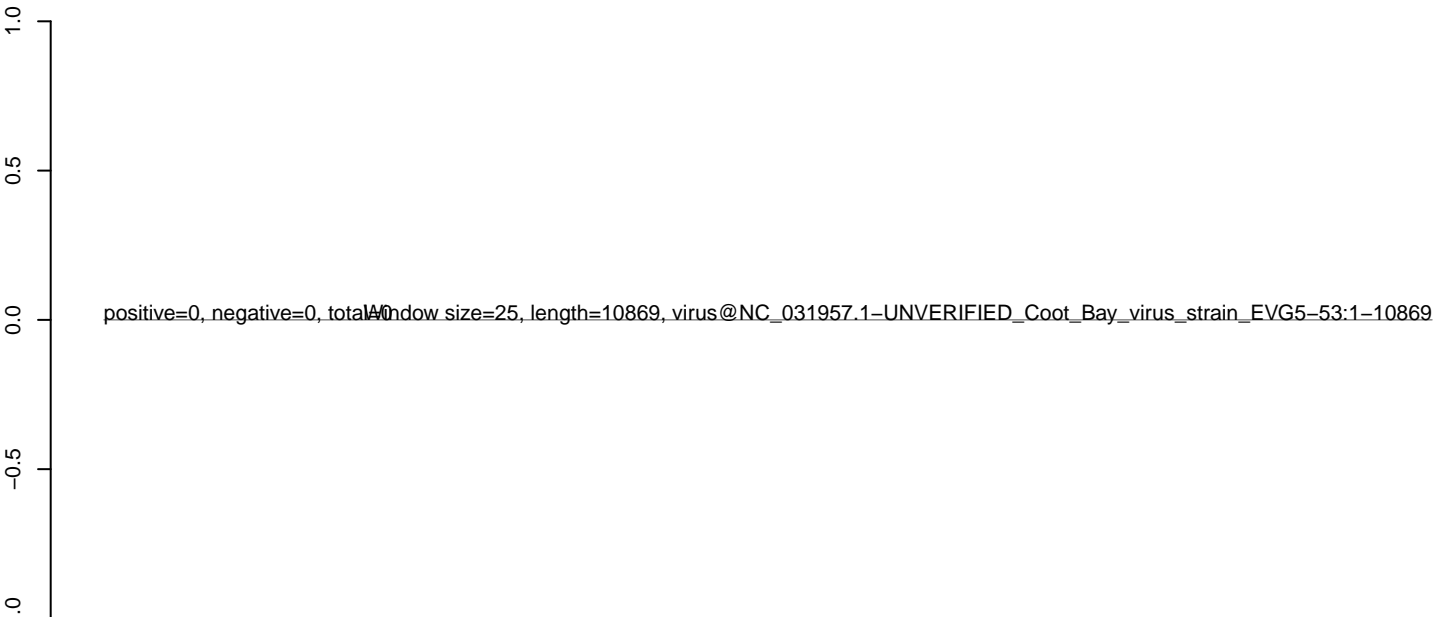
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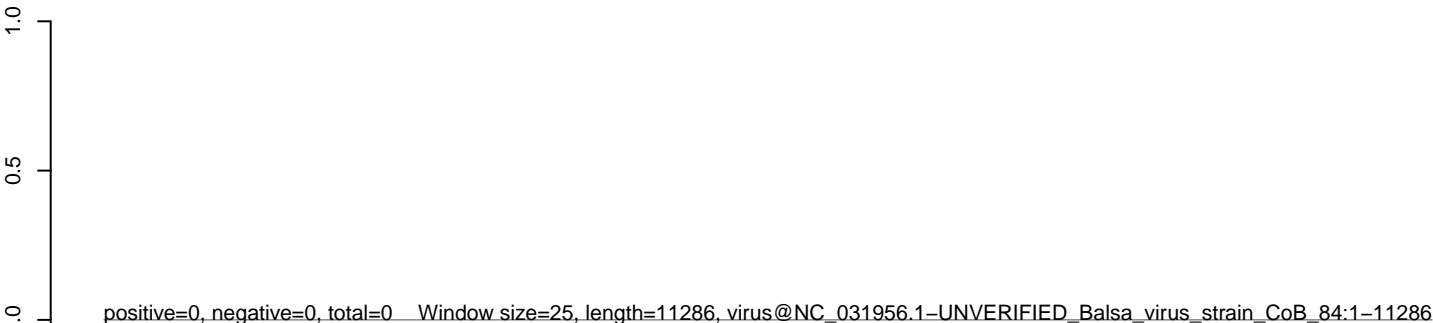
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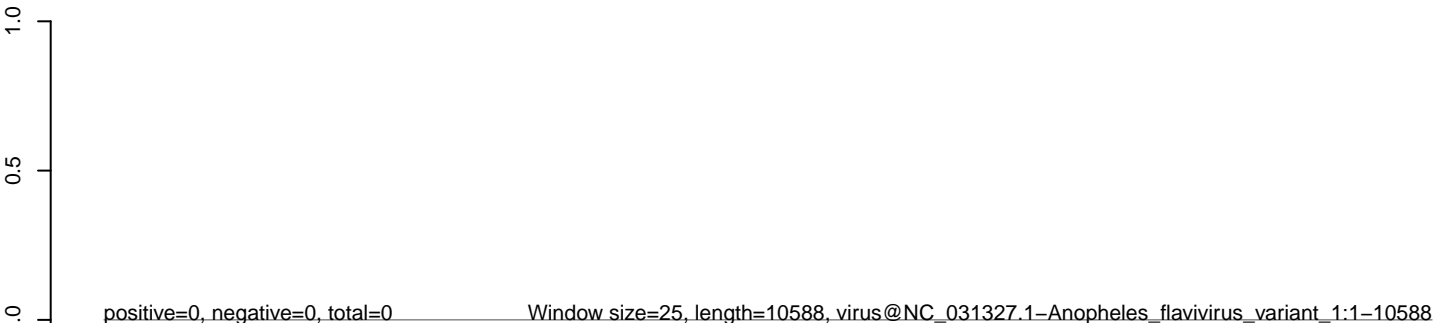
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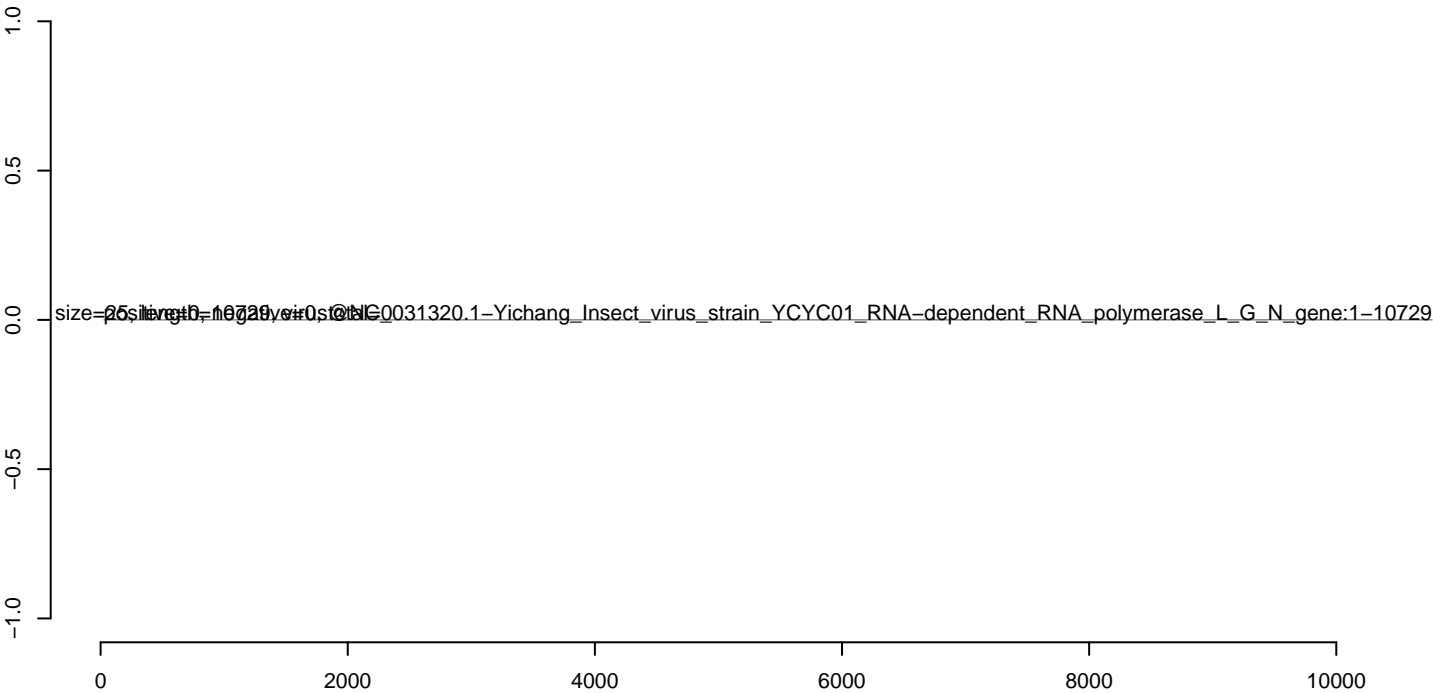
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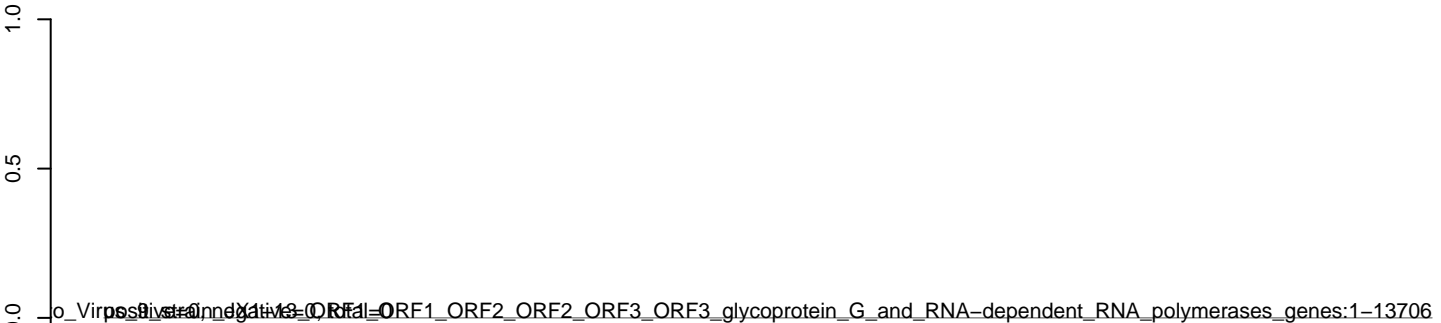
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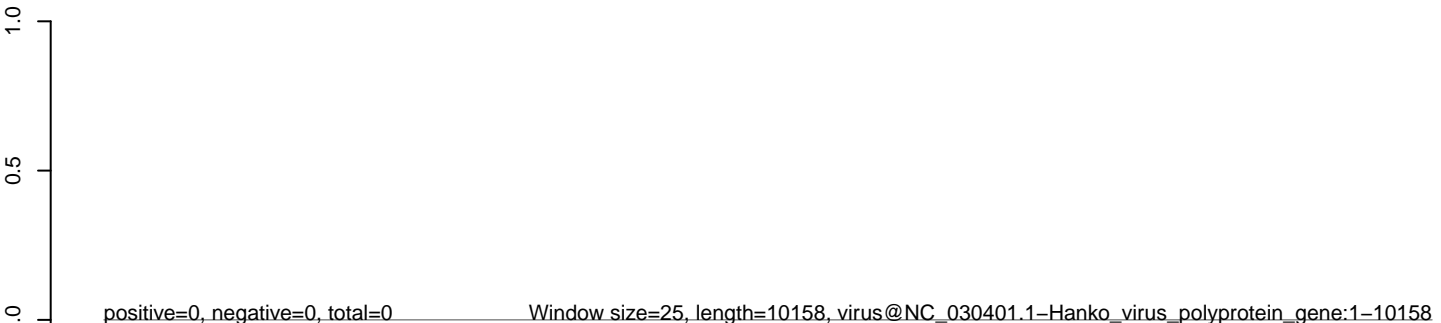
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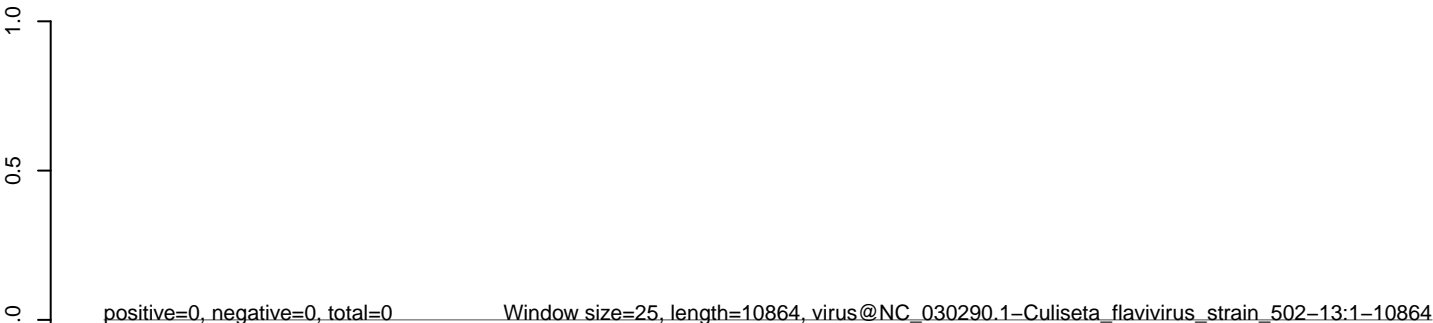
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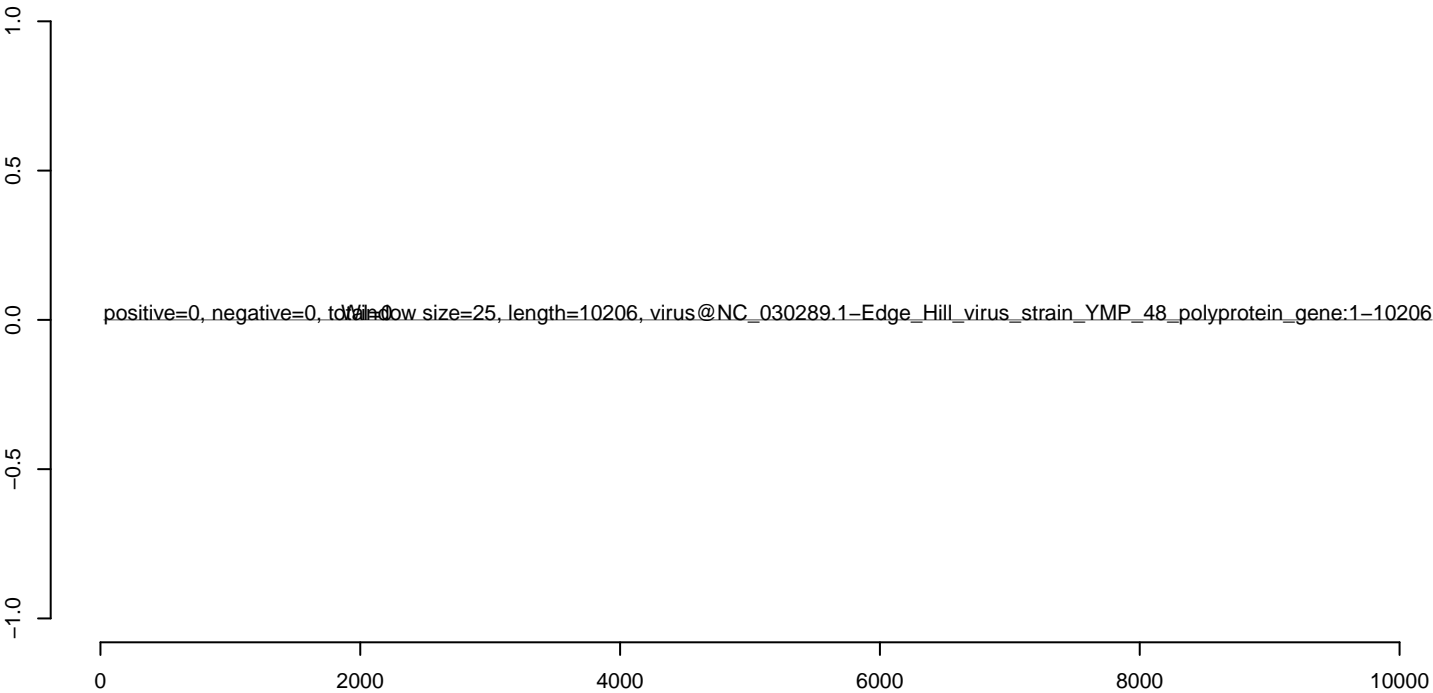
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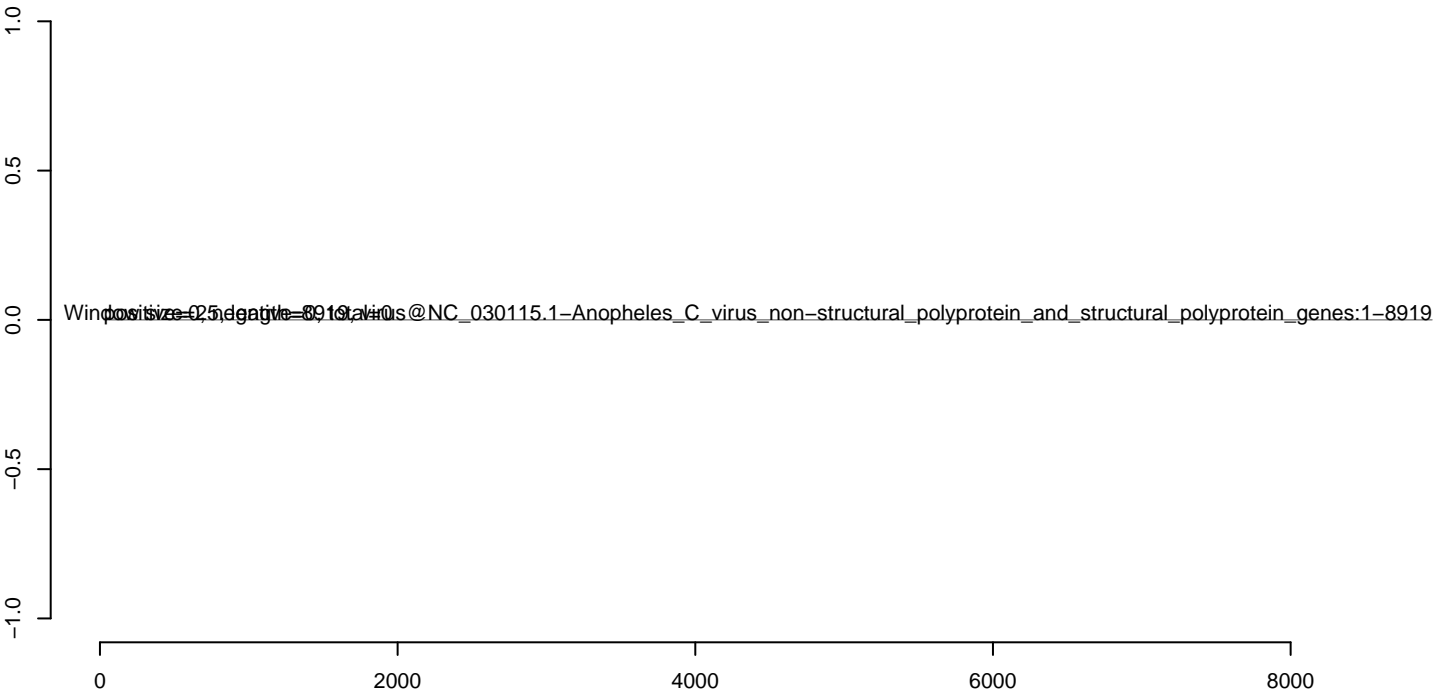
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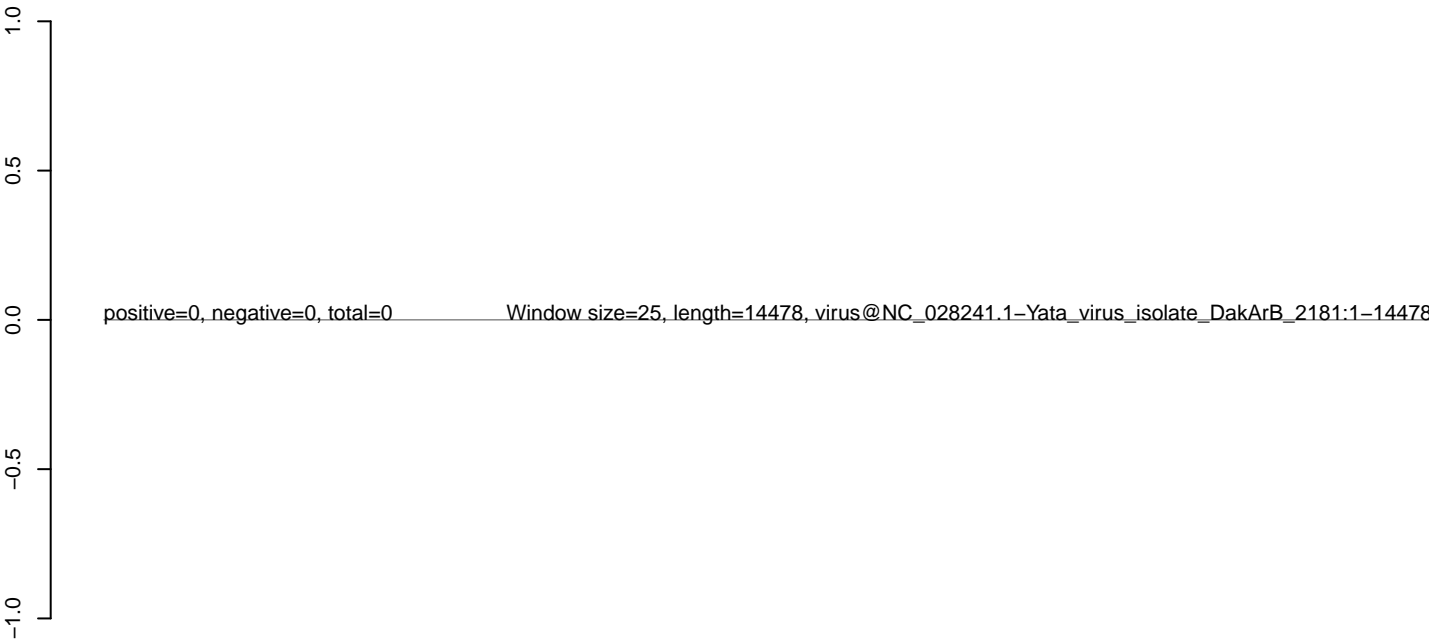
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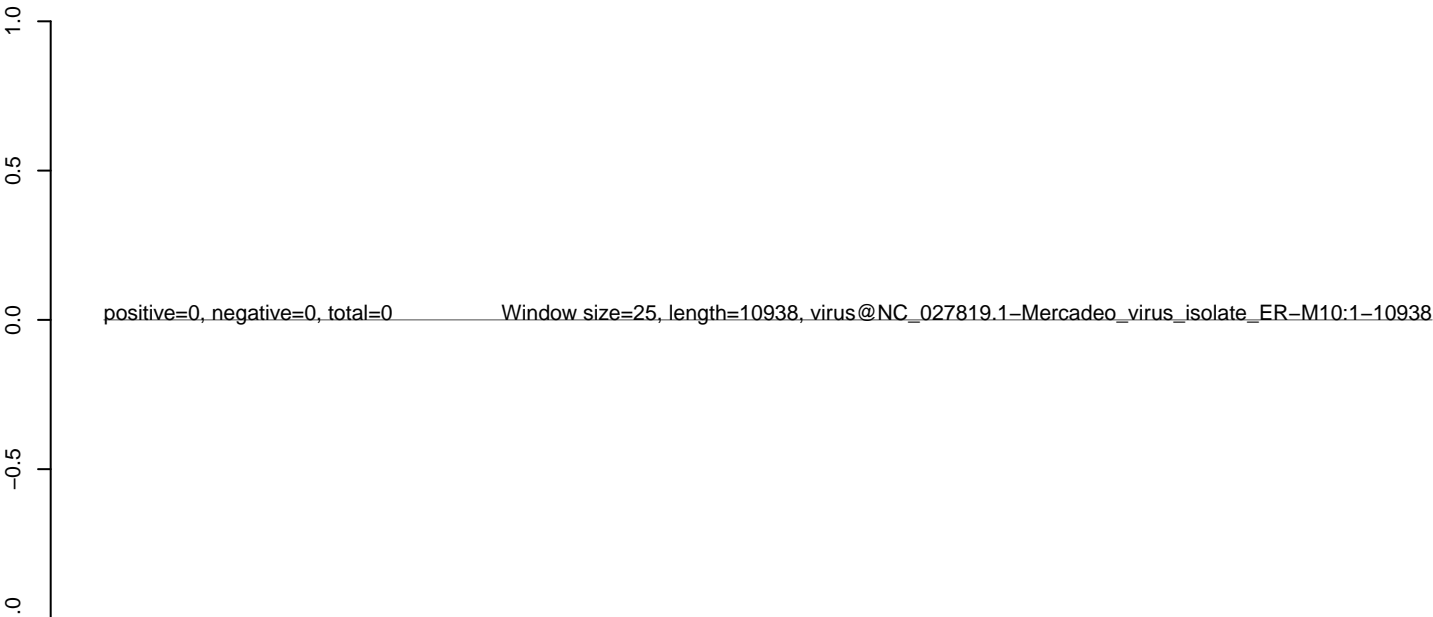
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**AnGam\_Mos55\_JR.18\_23.rep**



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**AnGam\_Mos55\_JR.rep**



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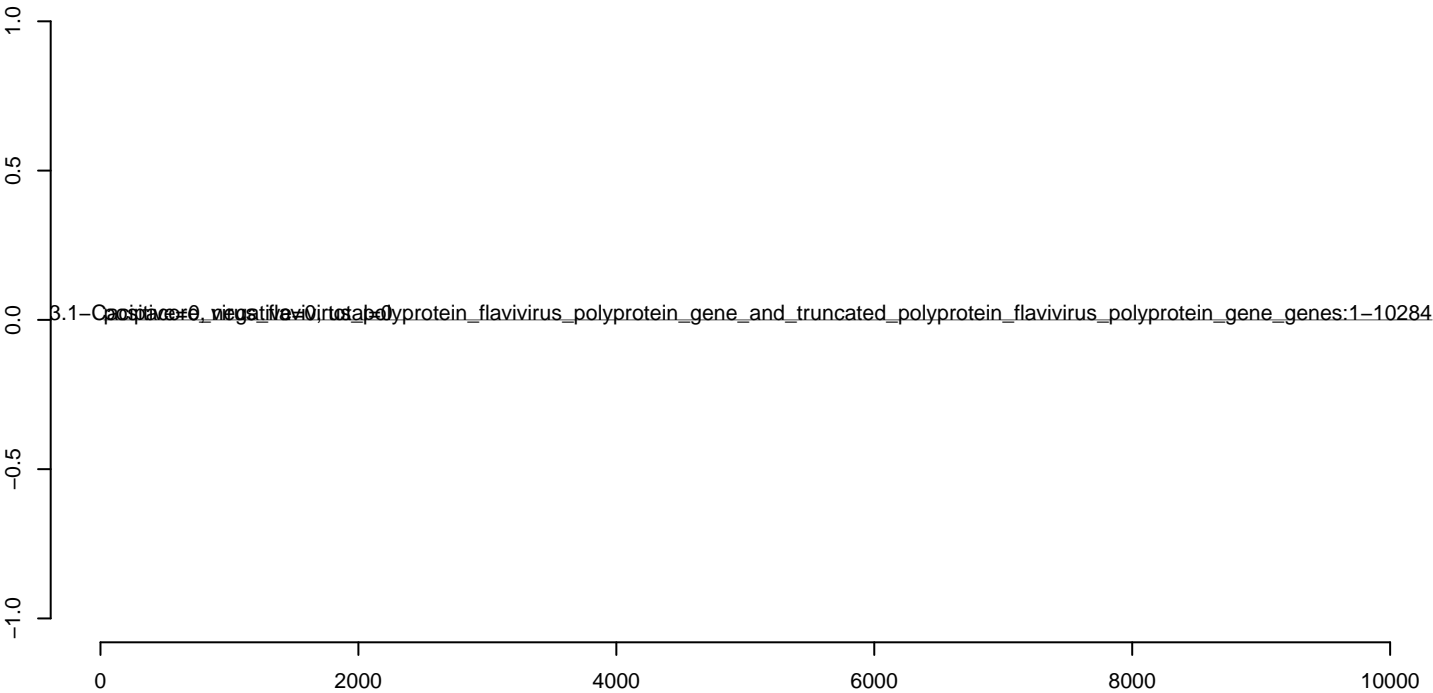
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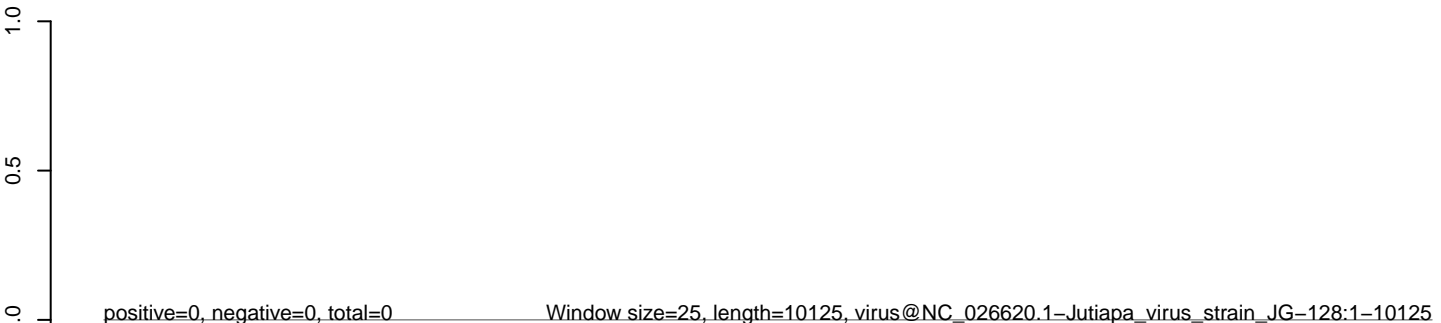
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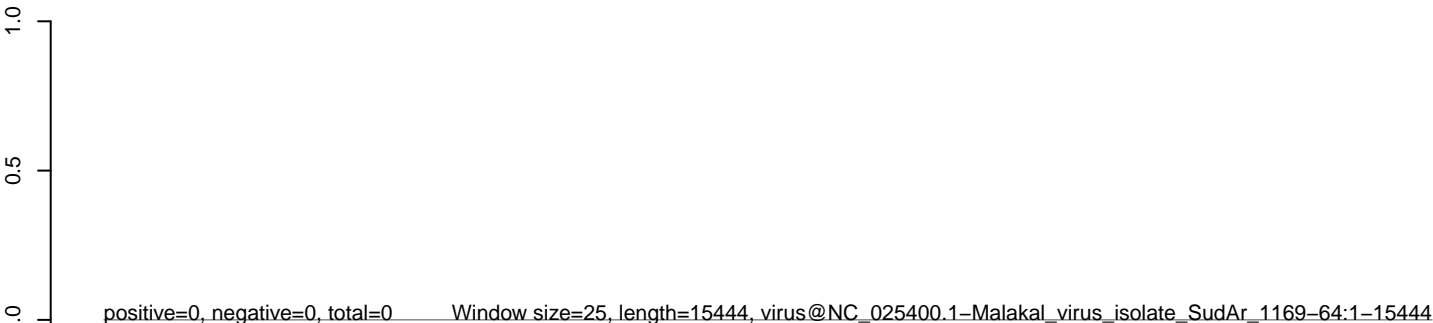
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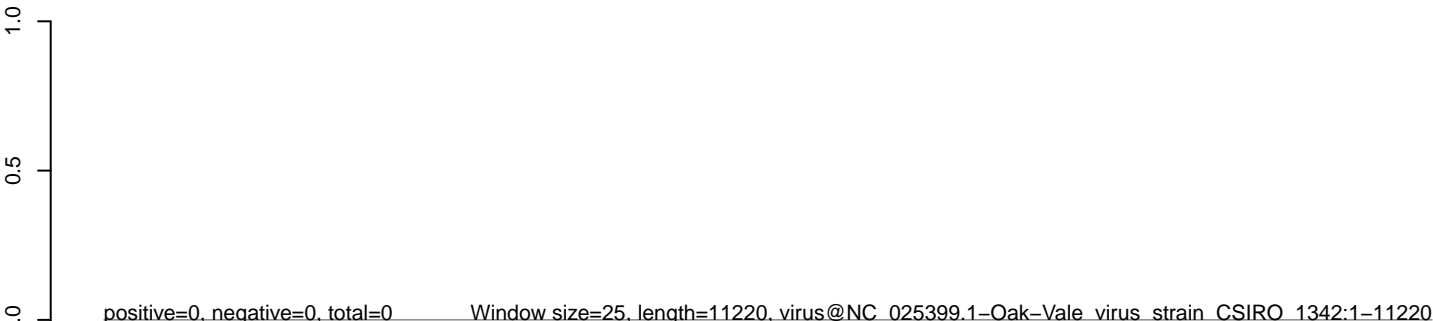
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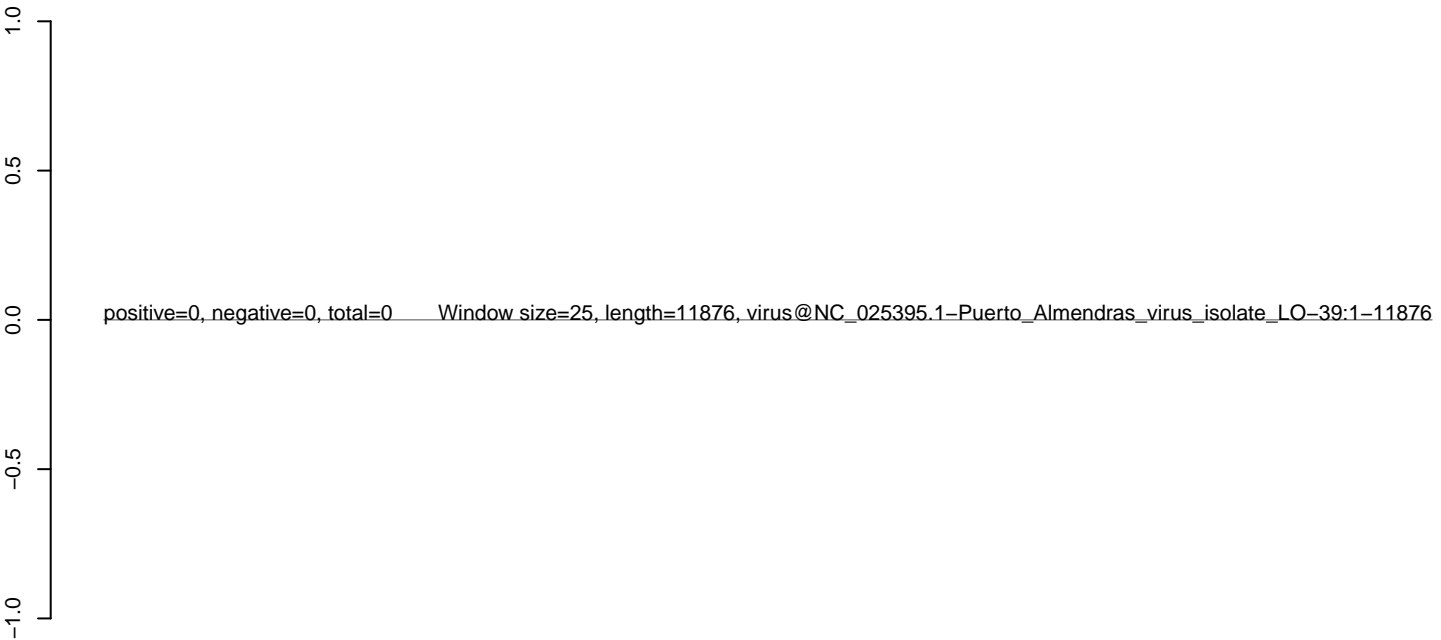
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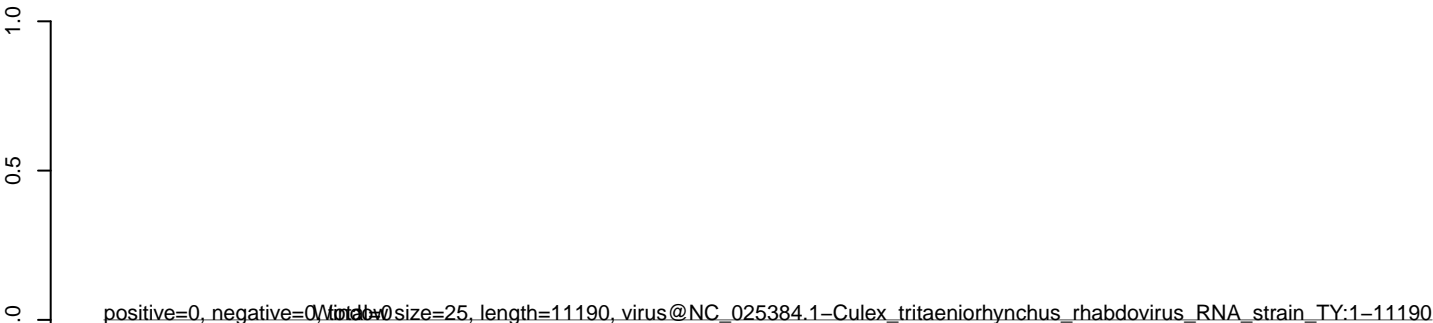
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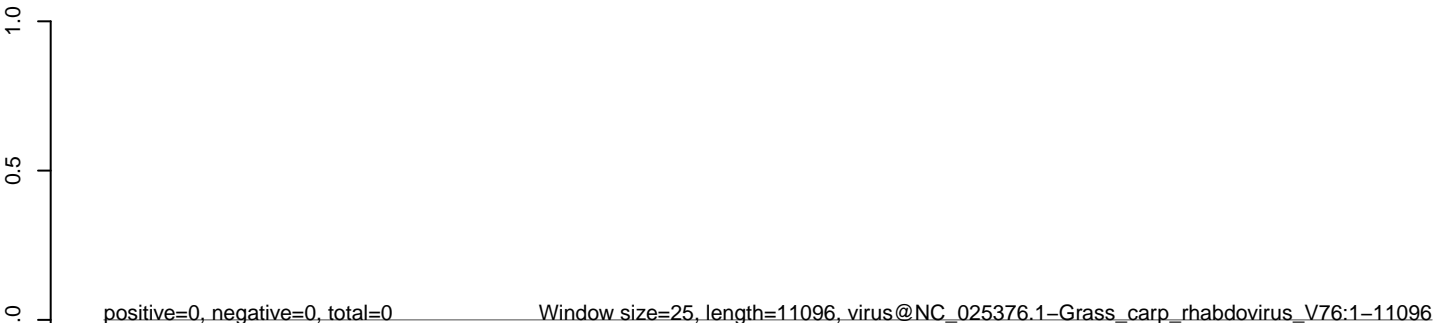
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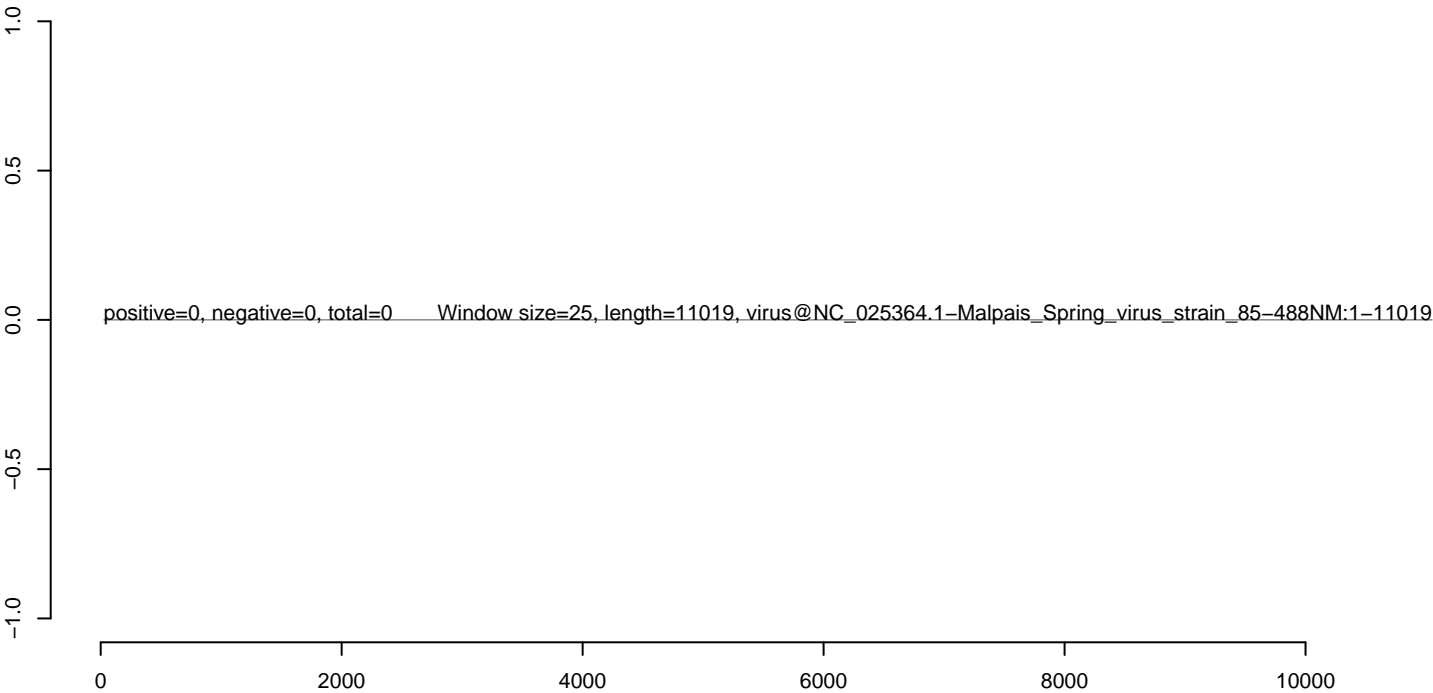
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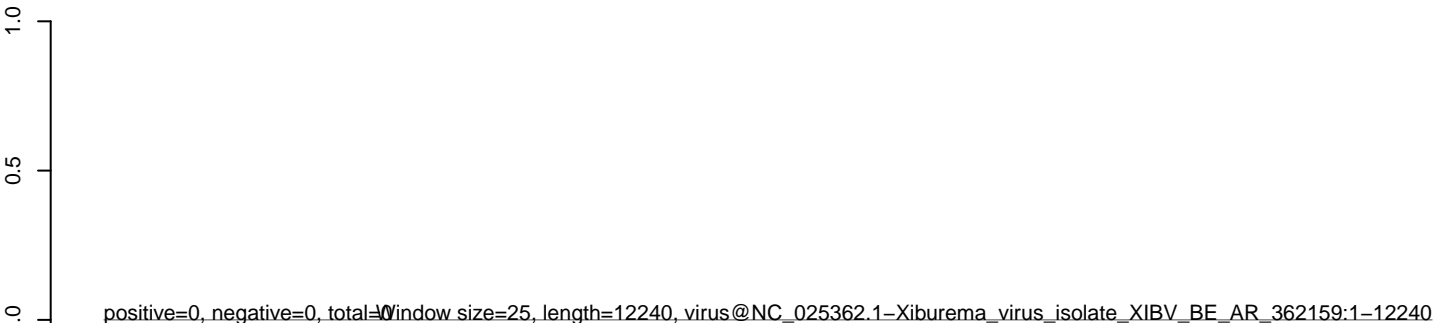
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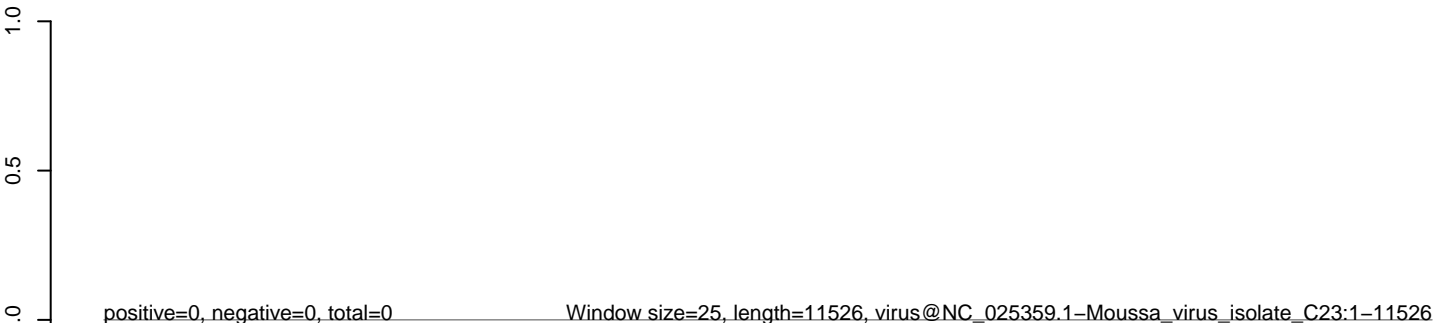
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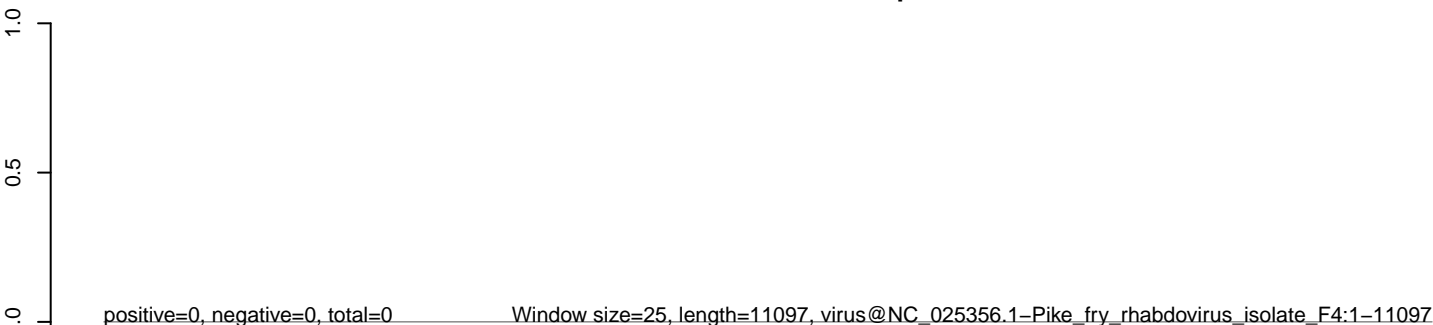
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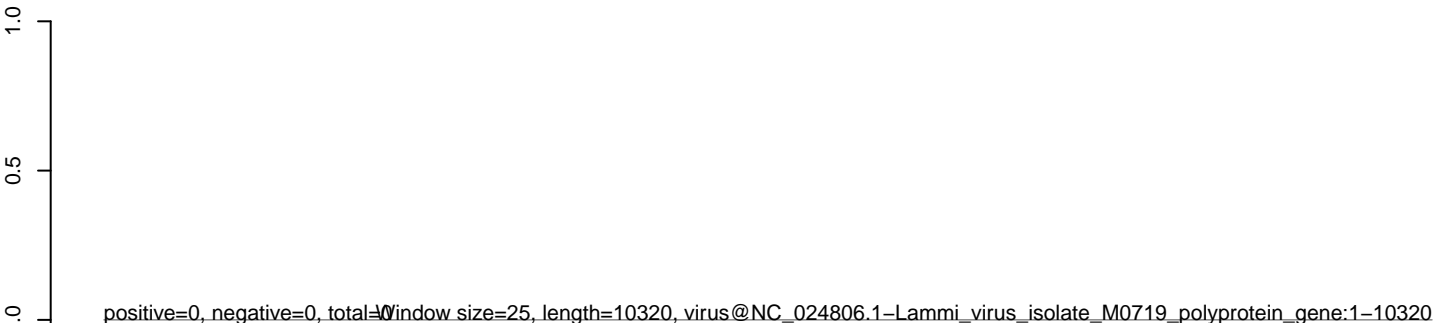
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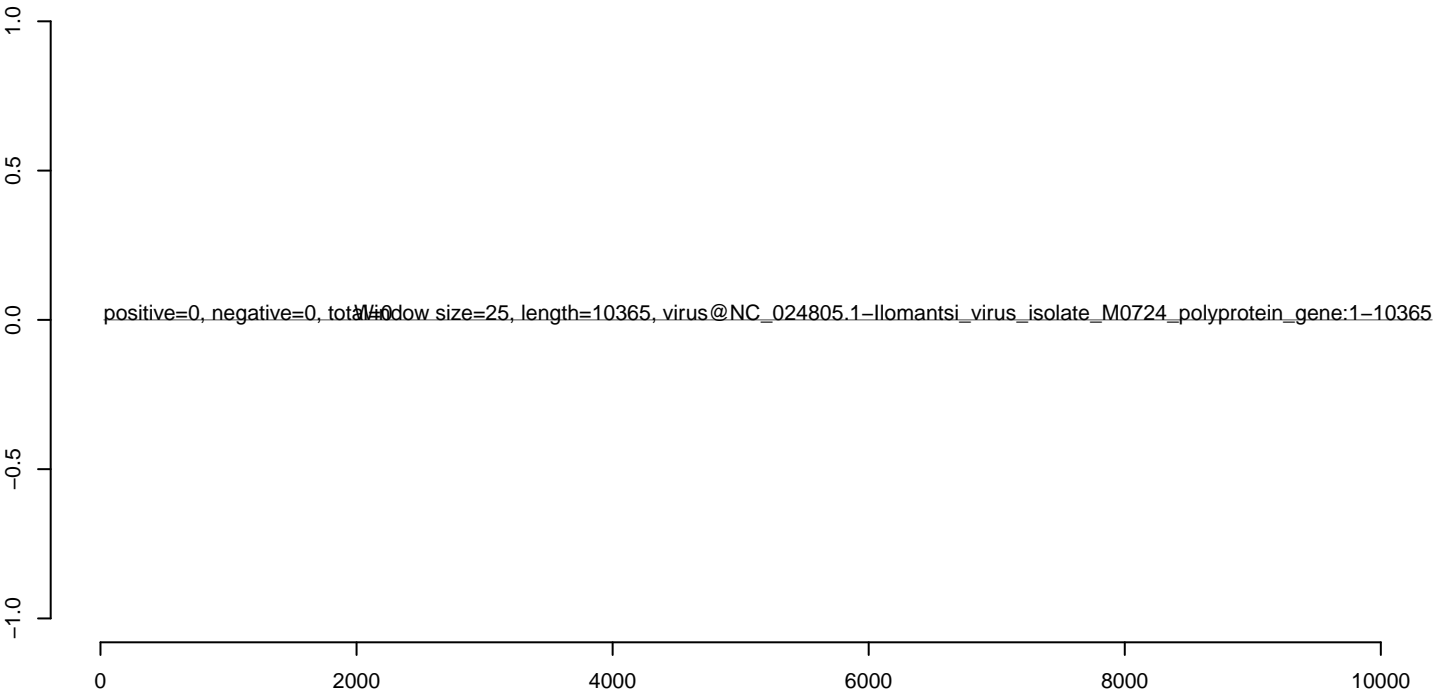
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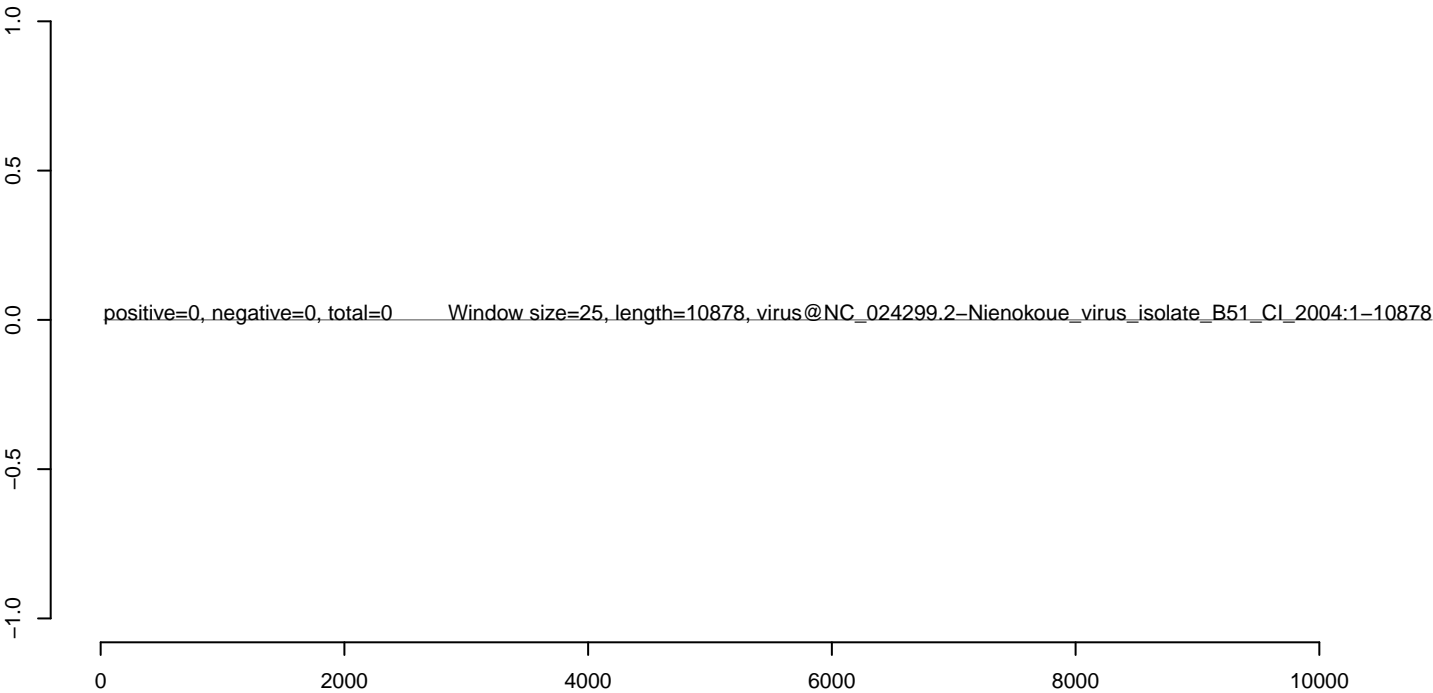
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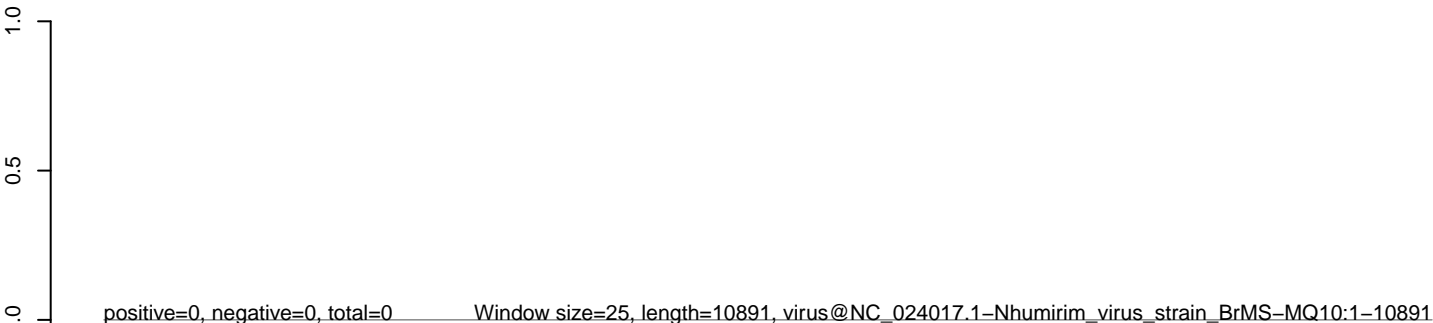
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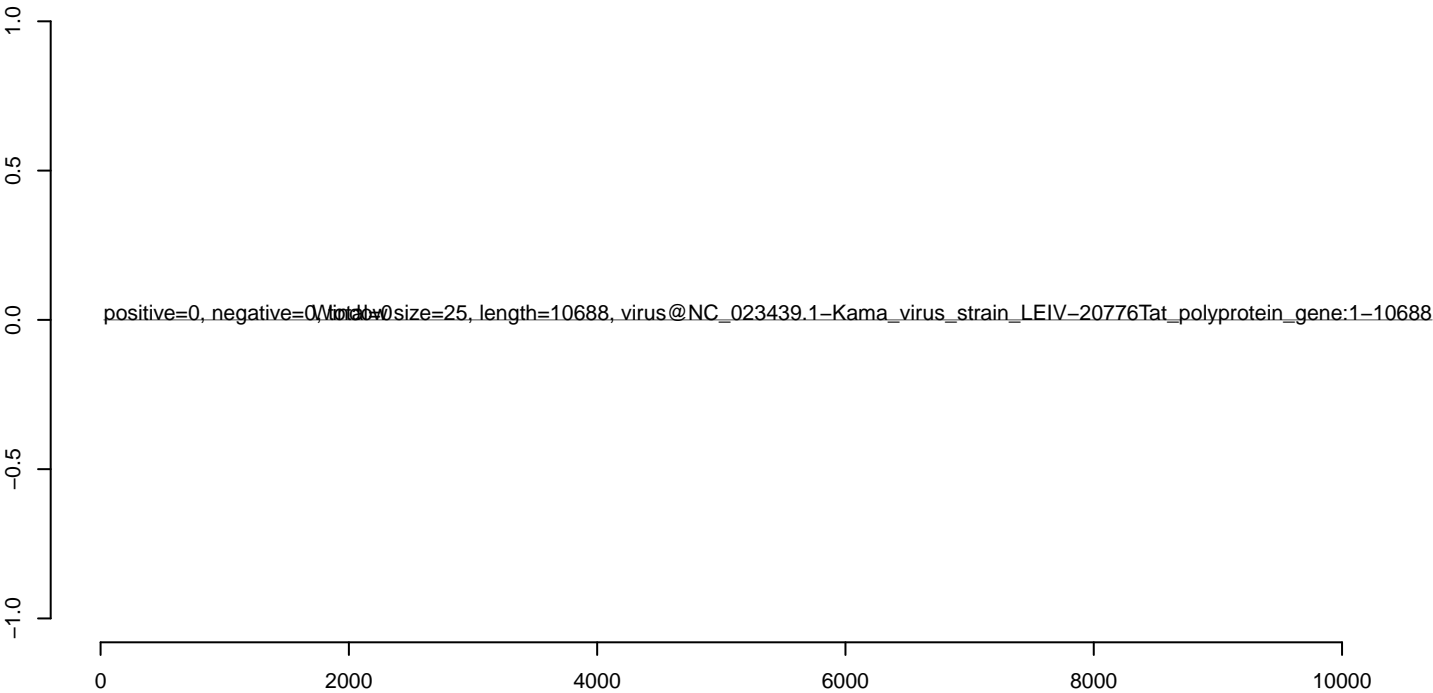
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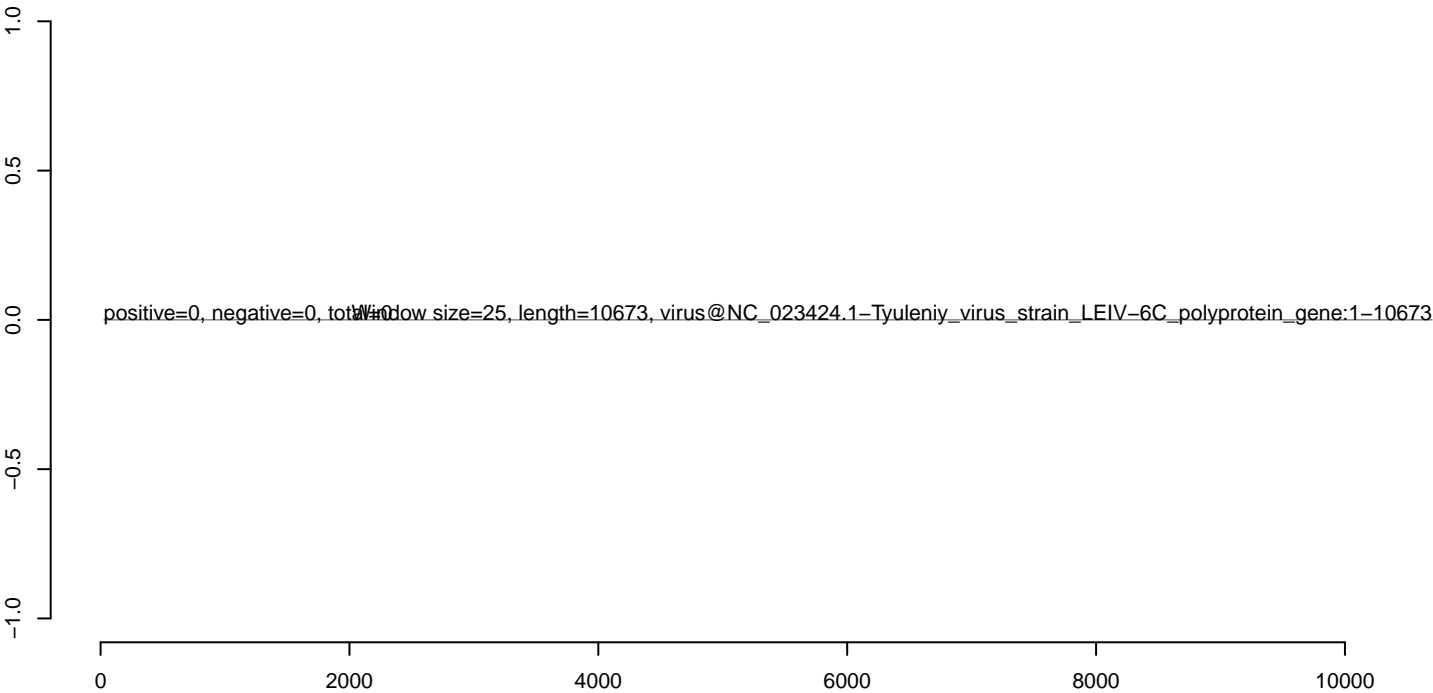
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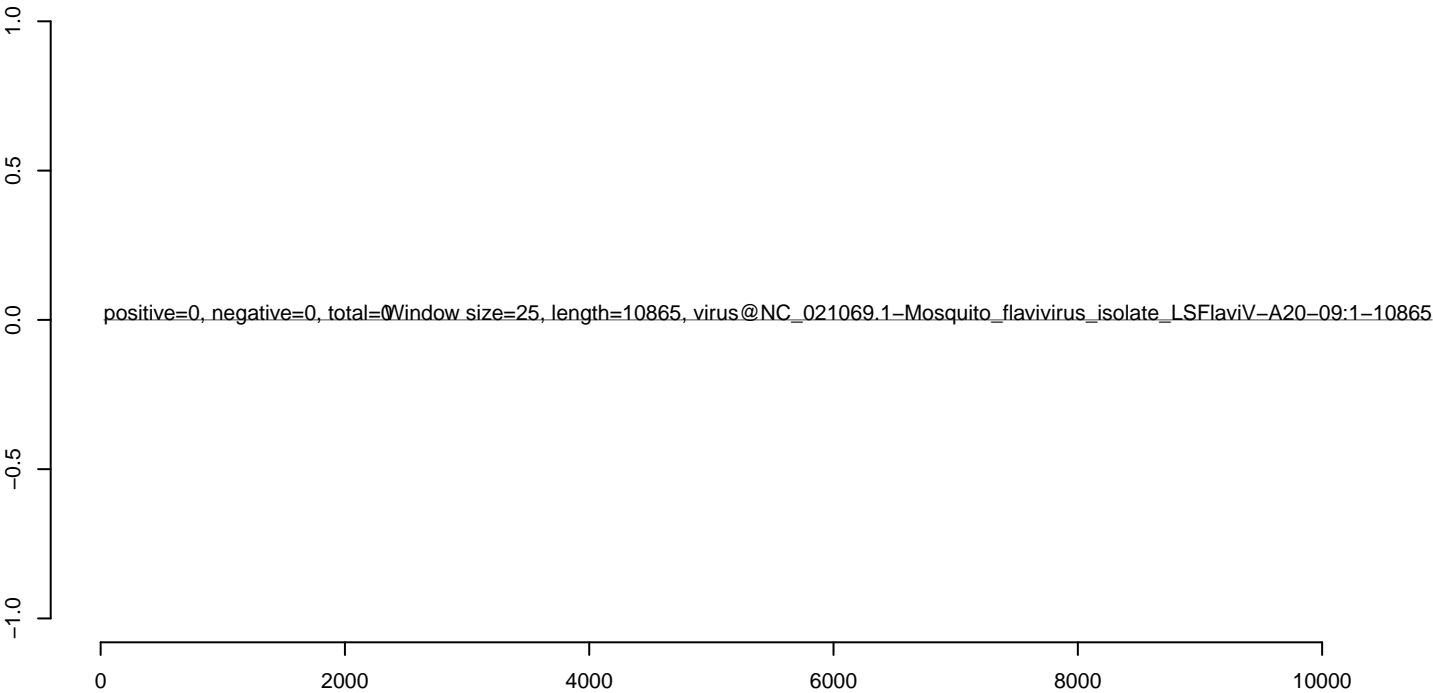
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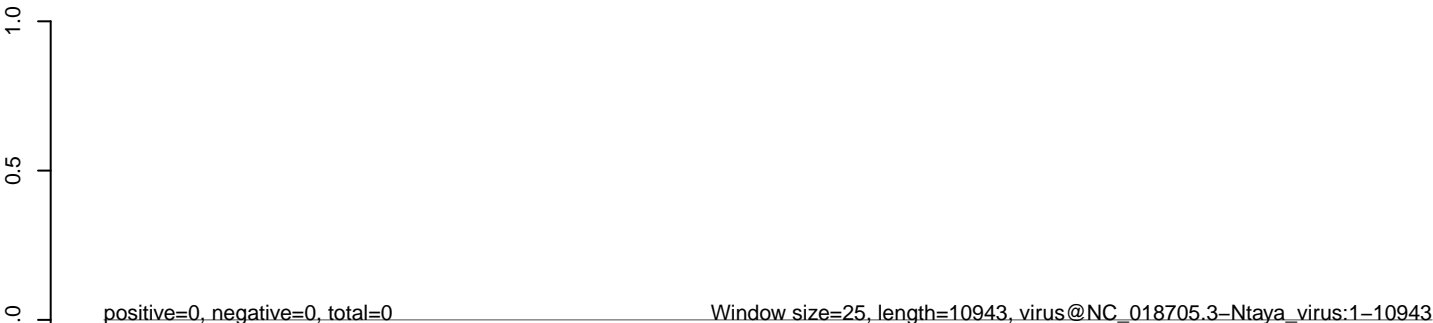
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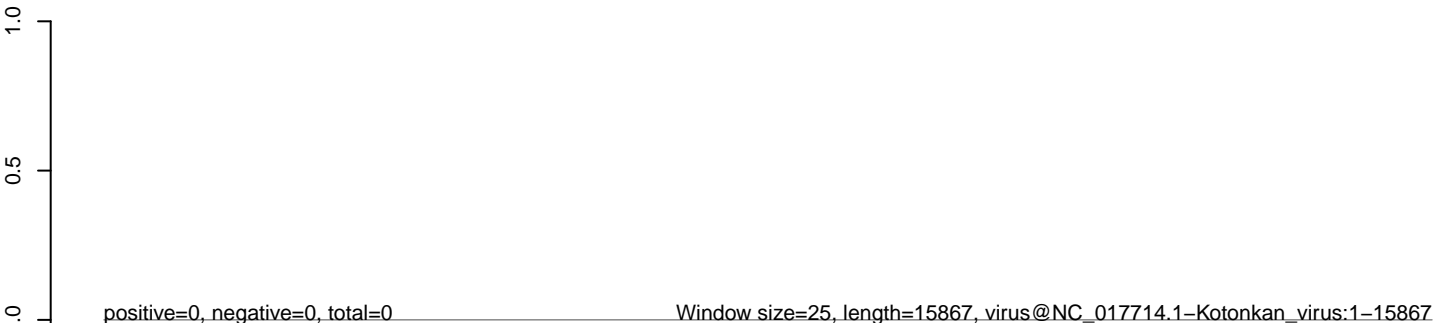
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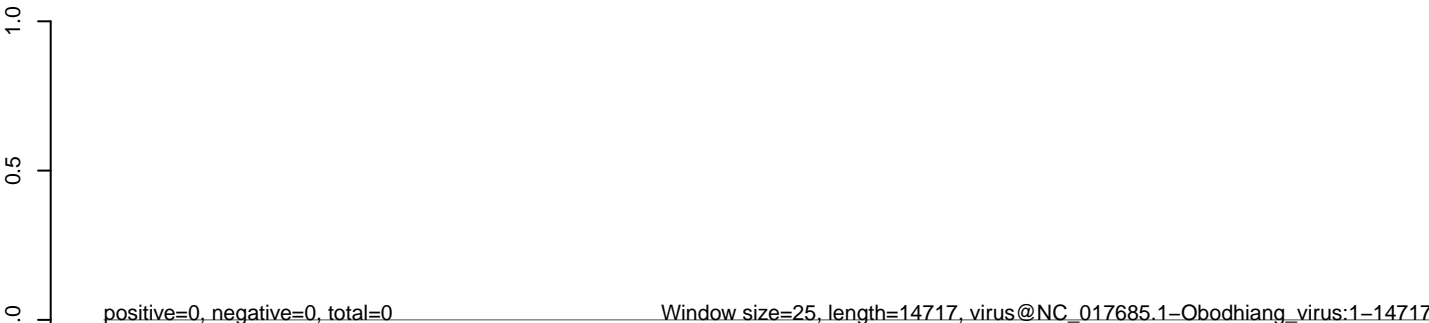
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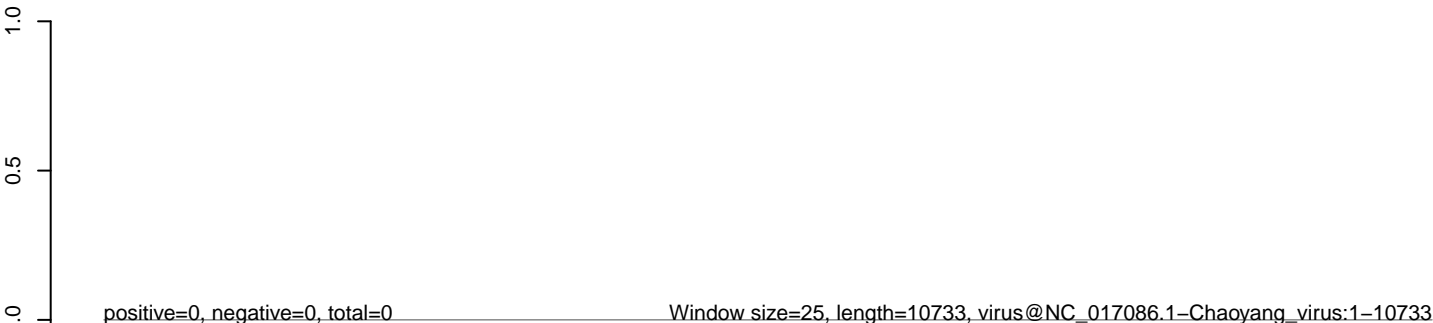
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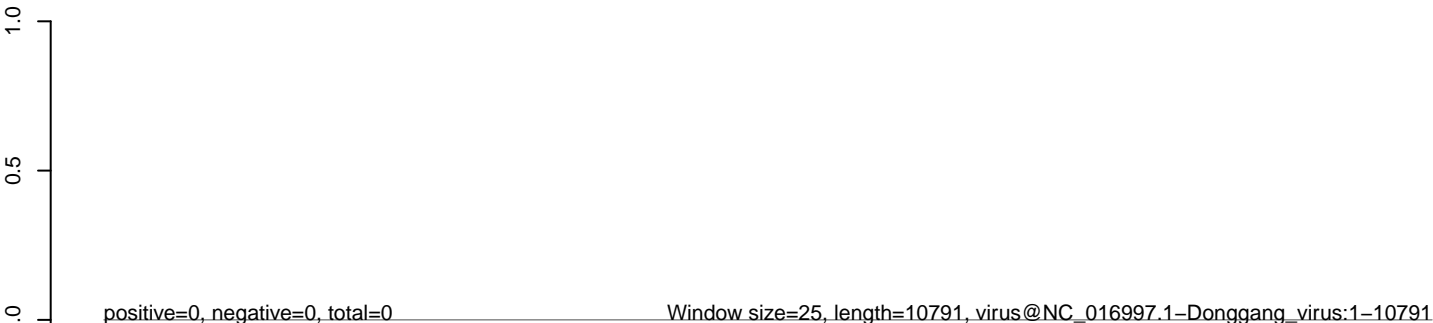
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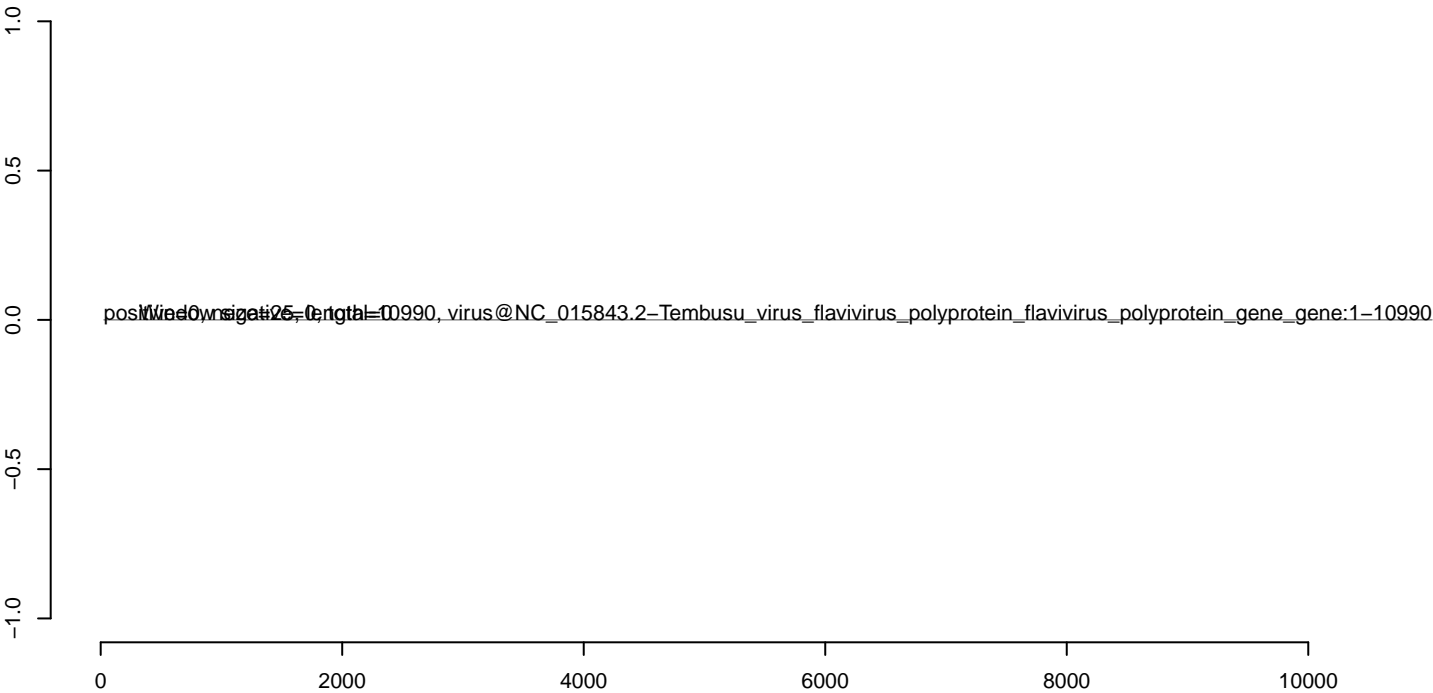
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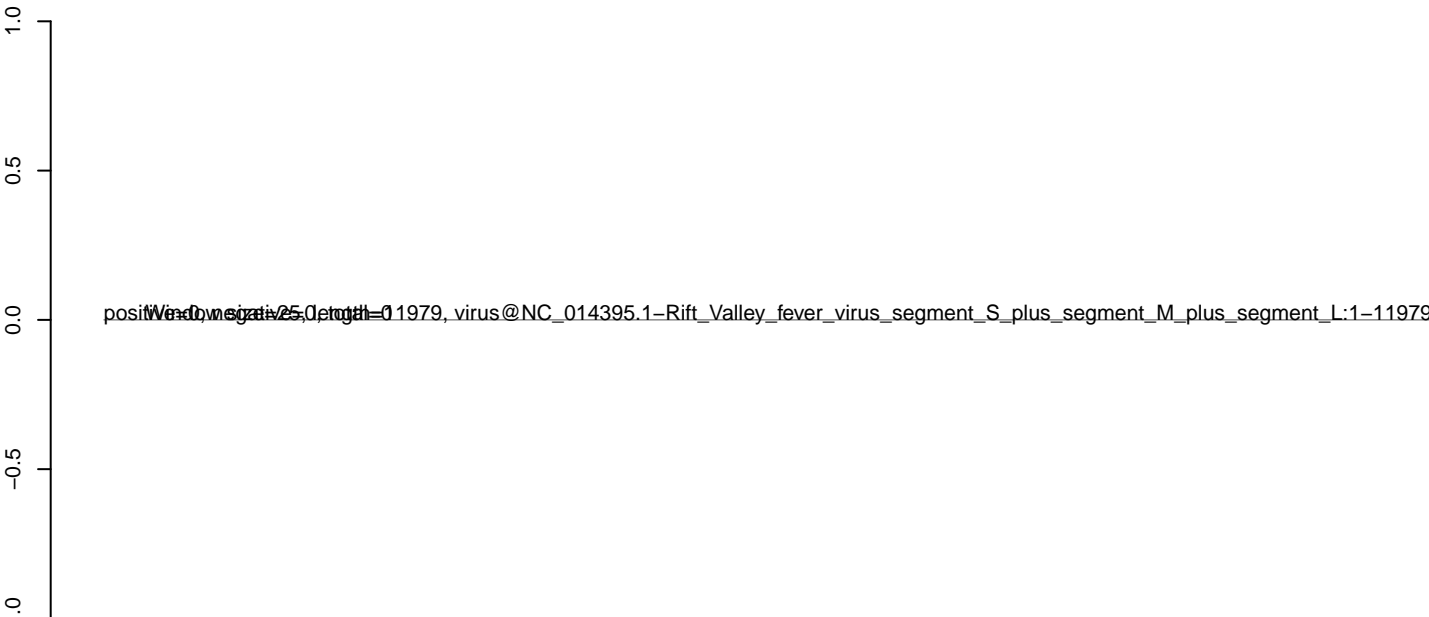
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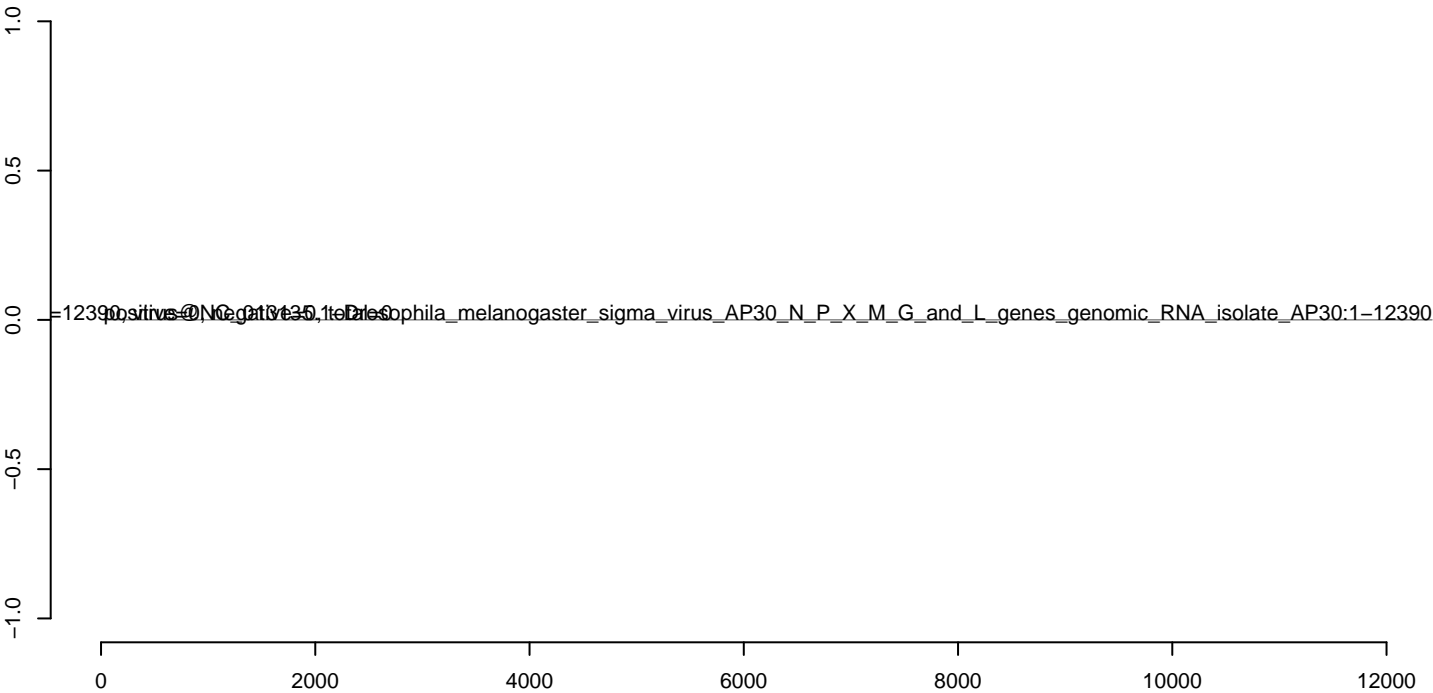
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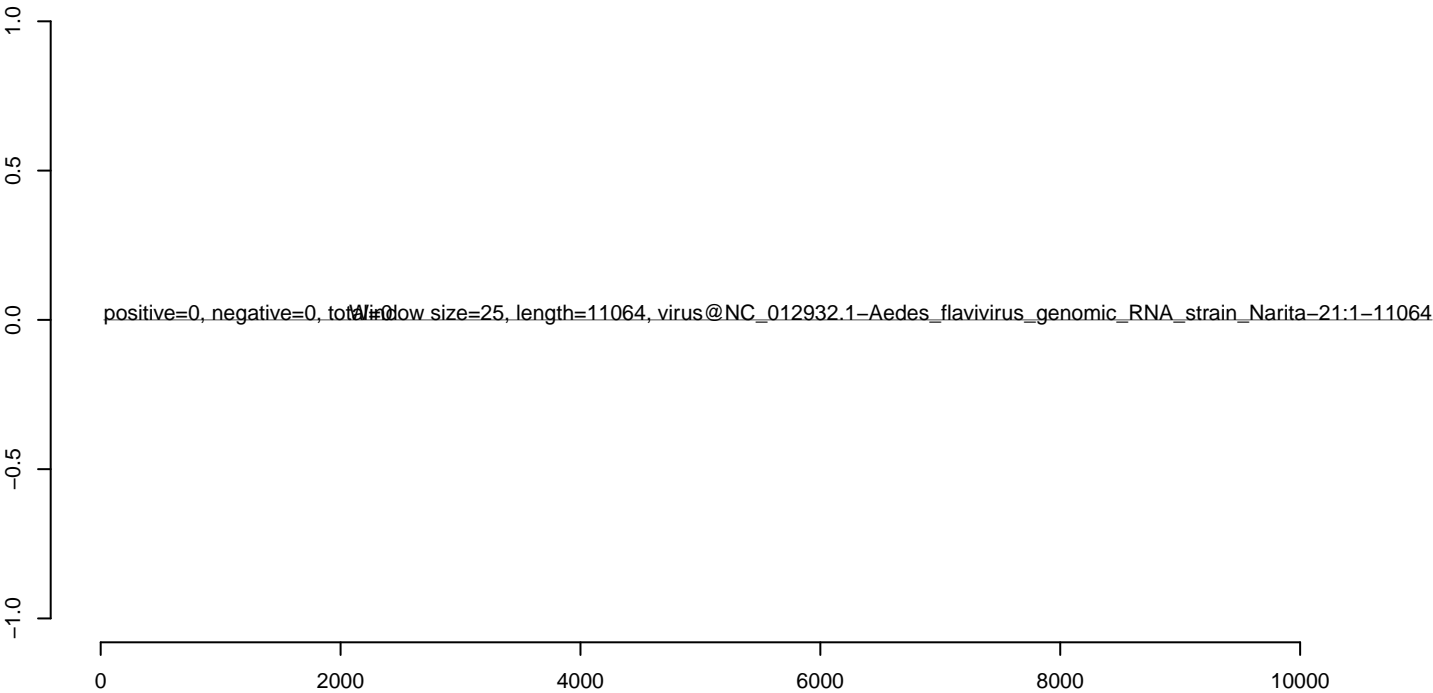
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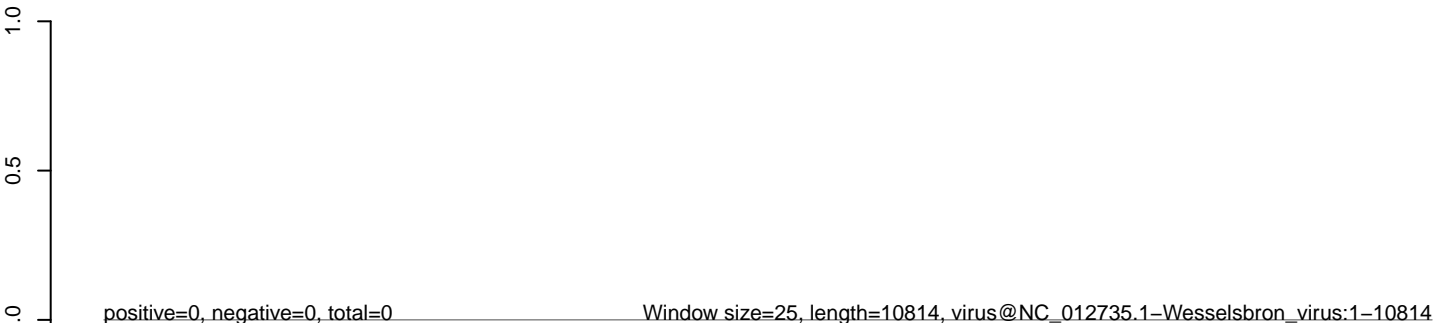
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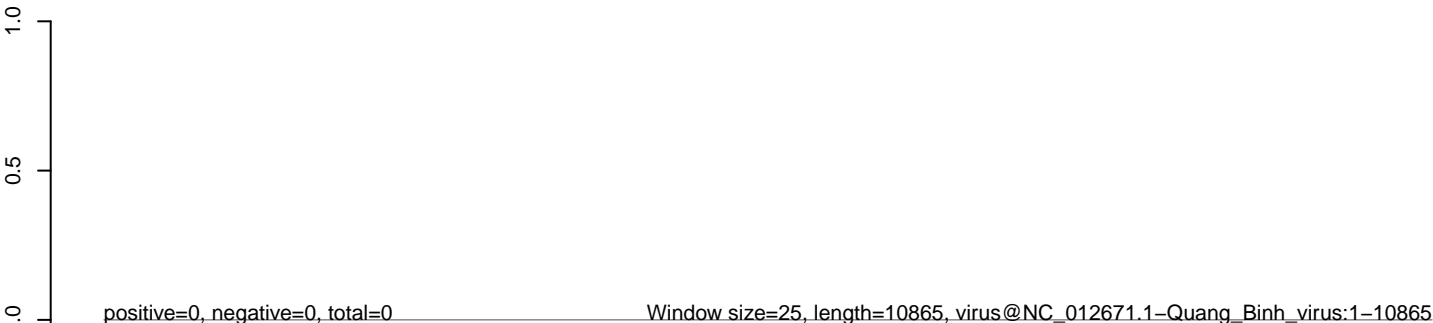
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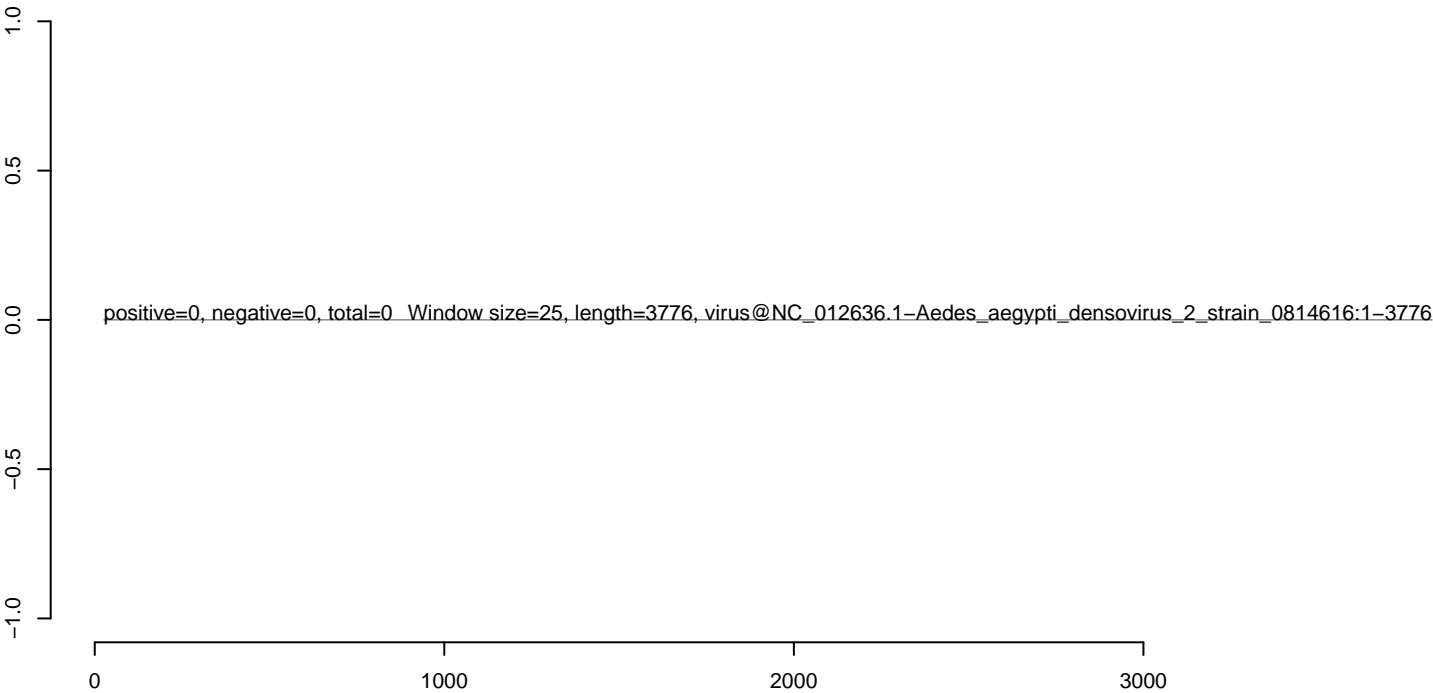
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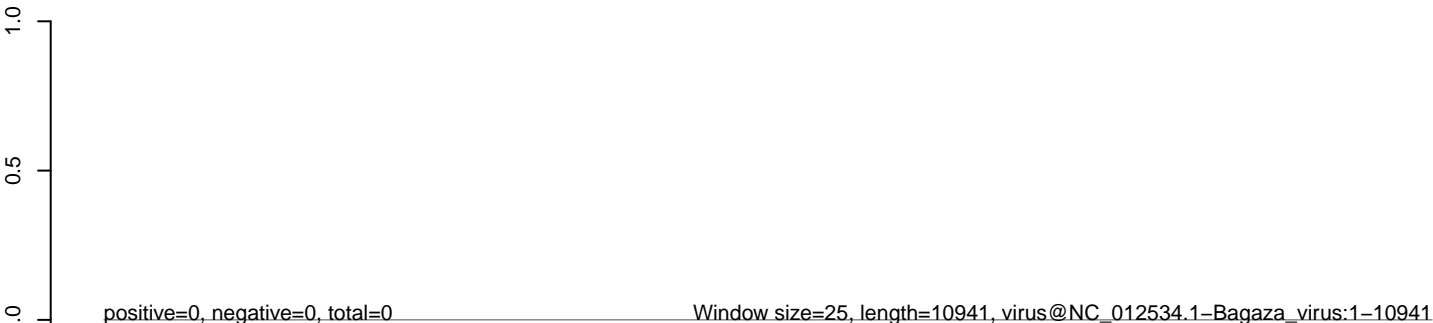
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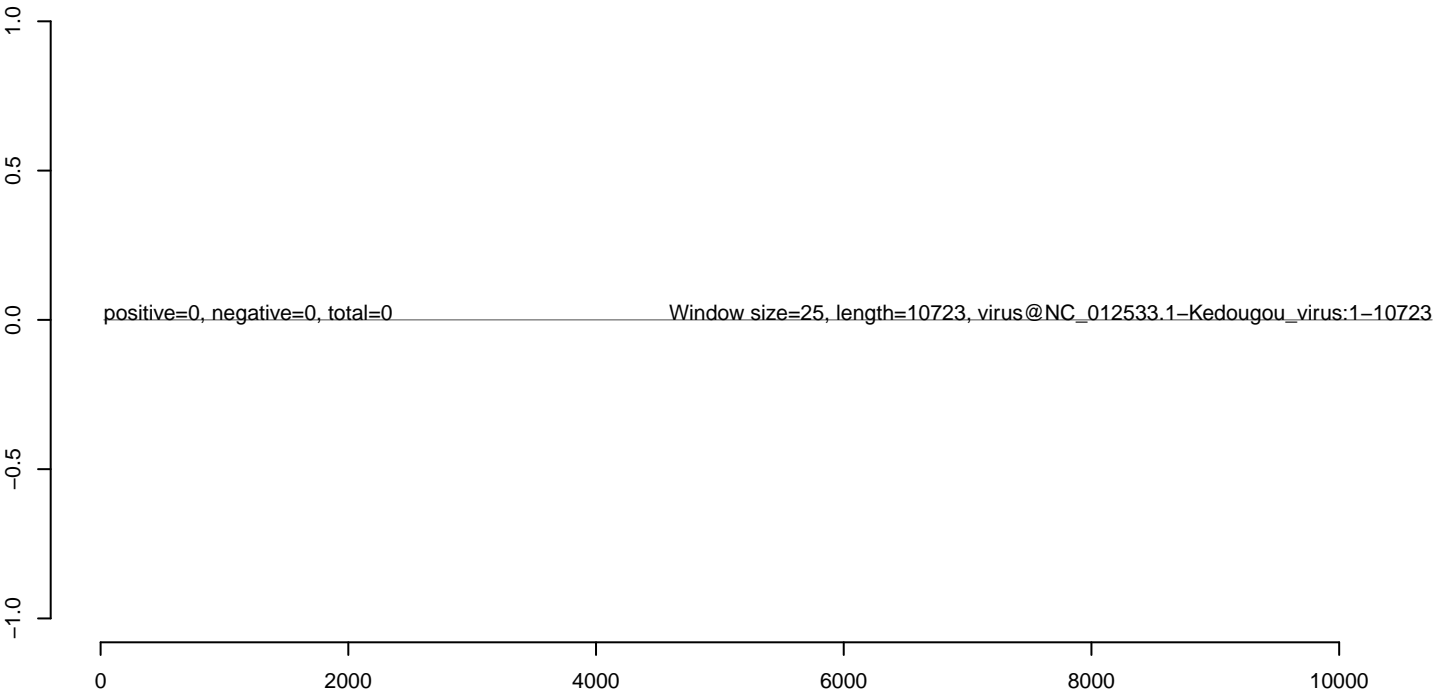
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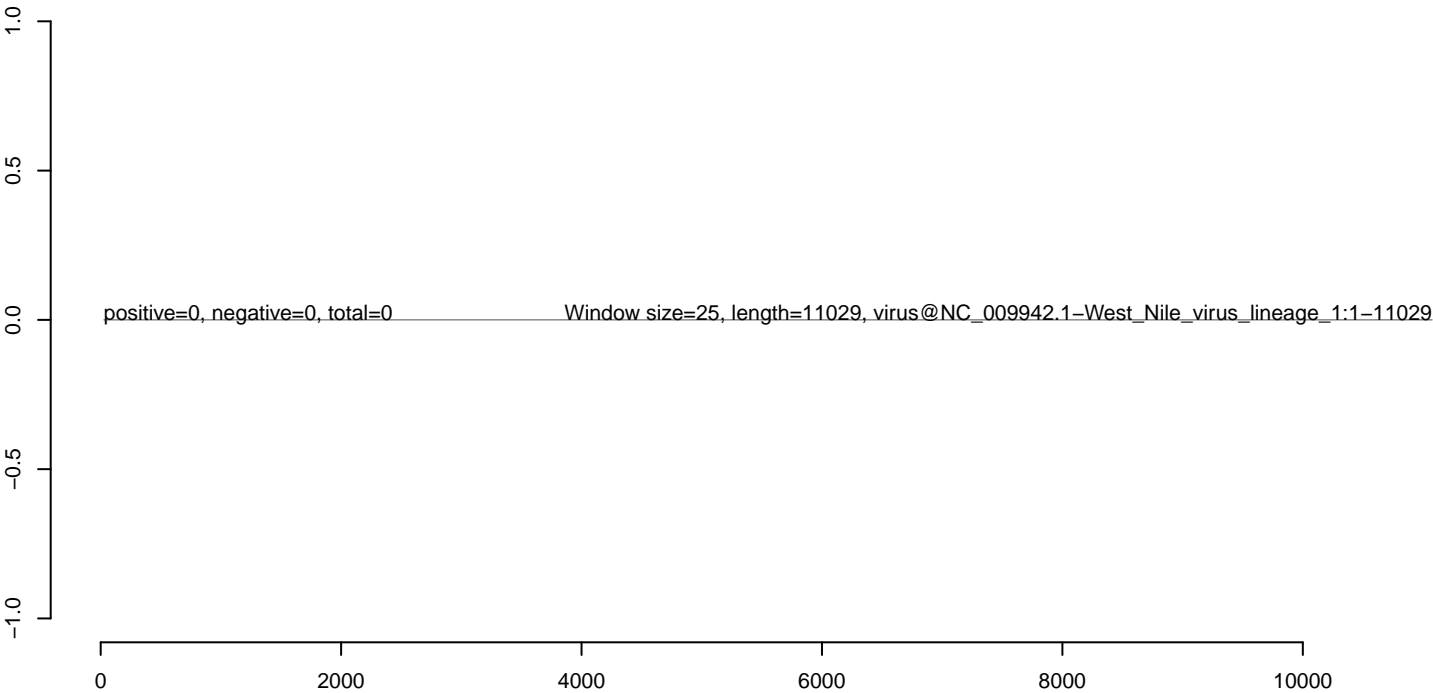
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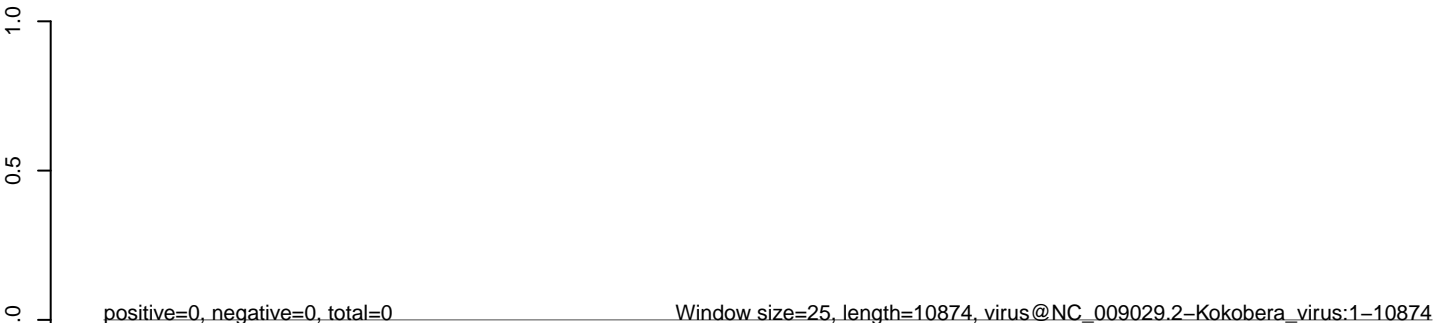
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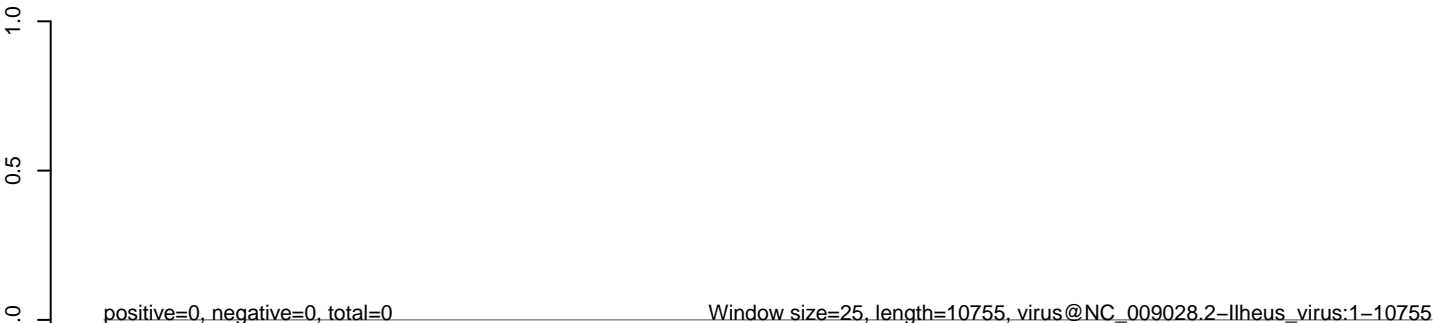
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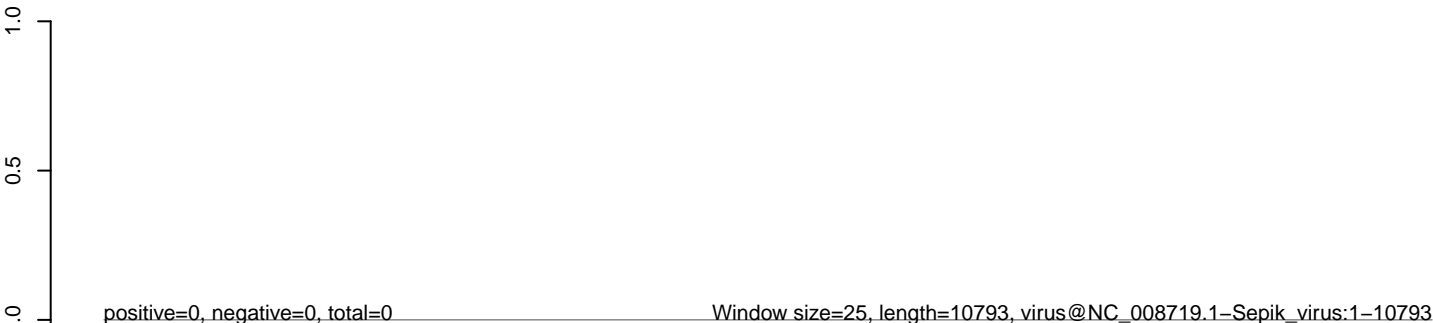
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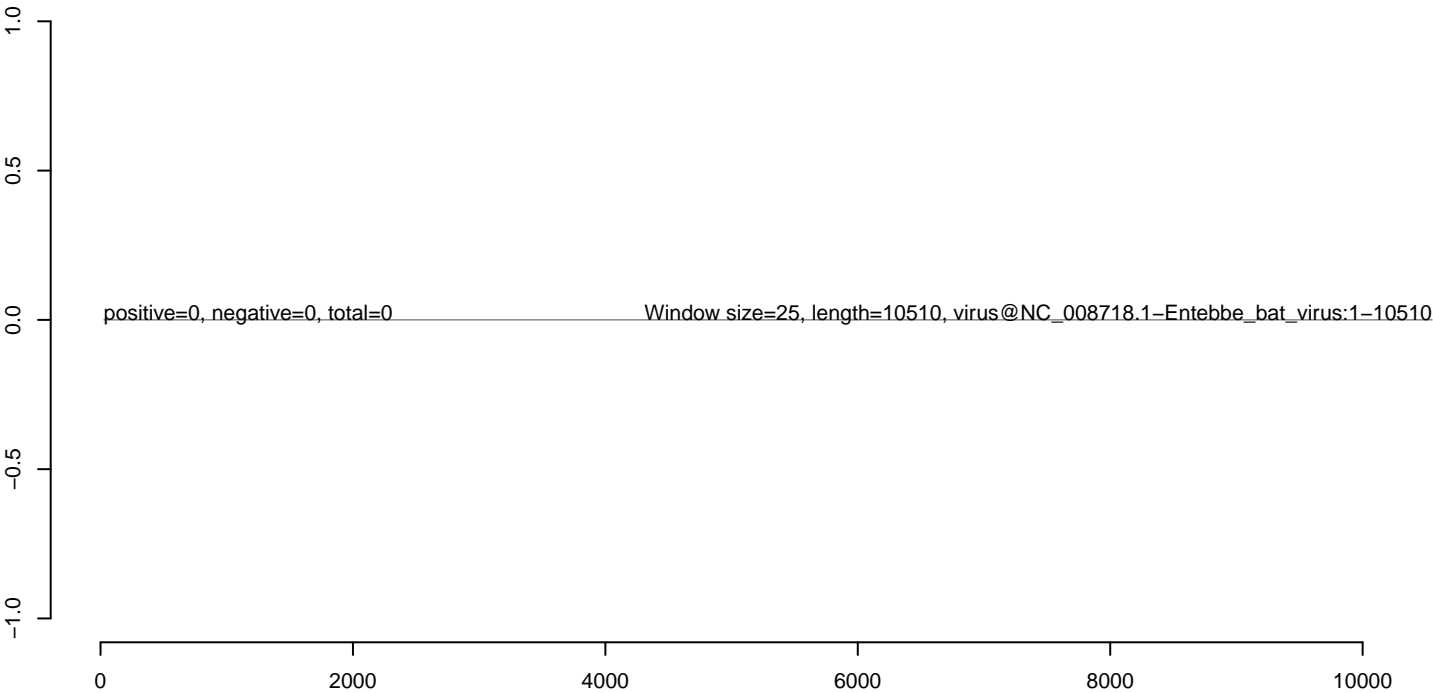
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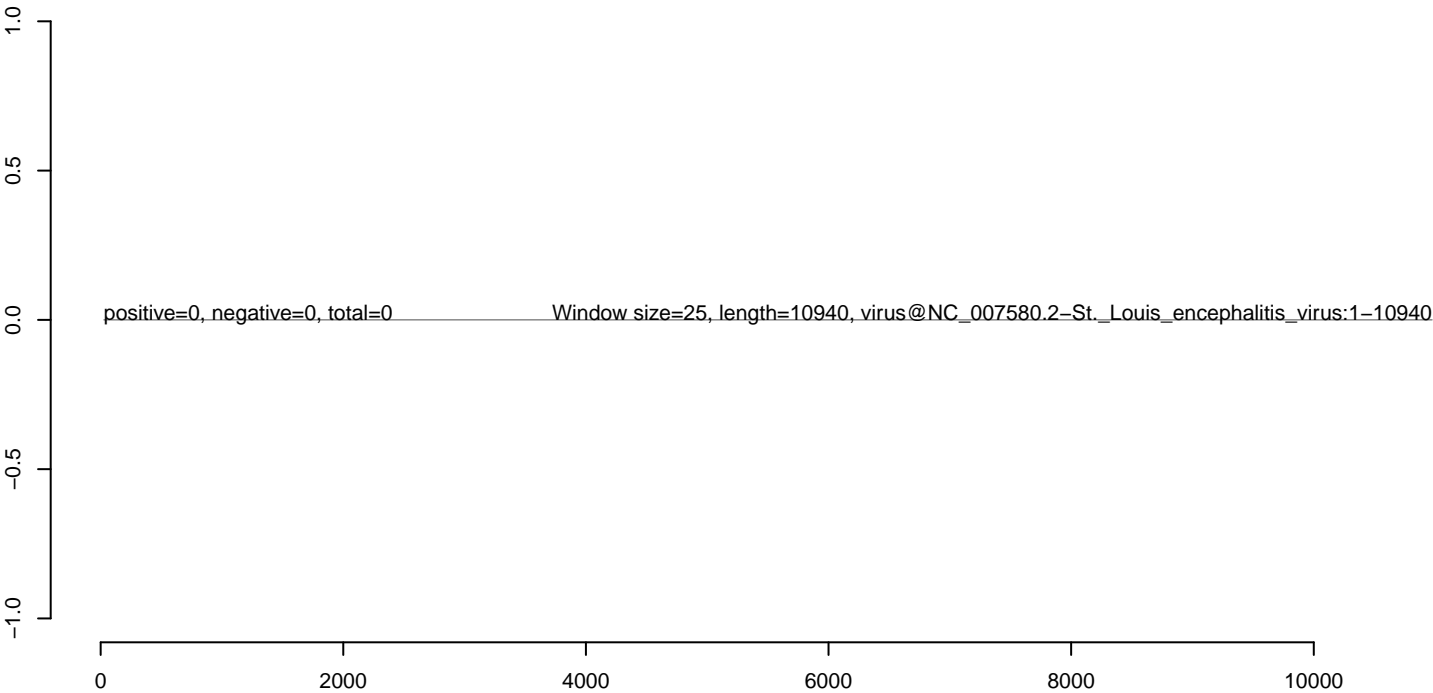
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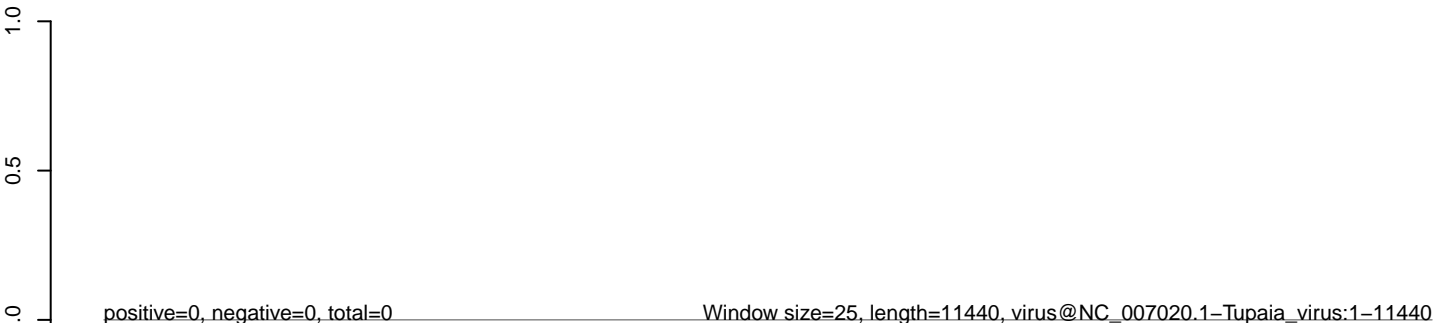
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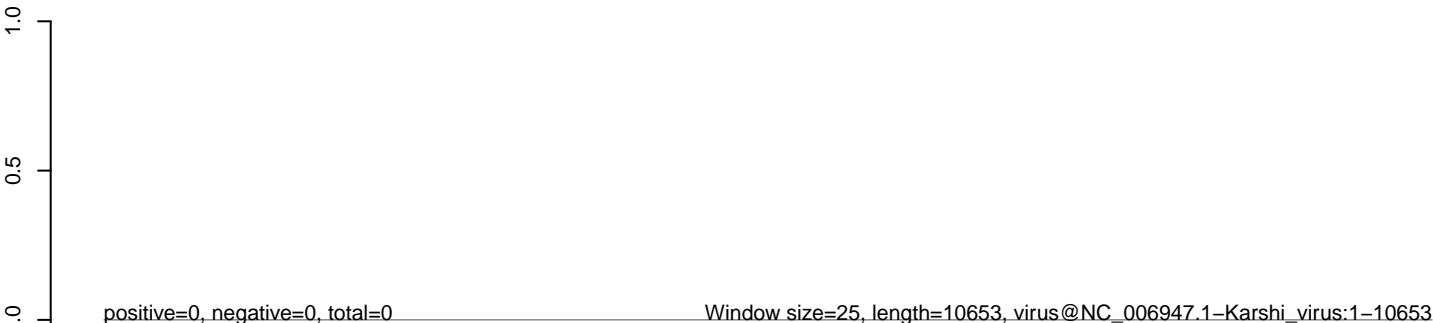
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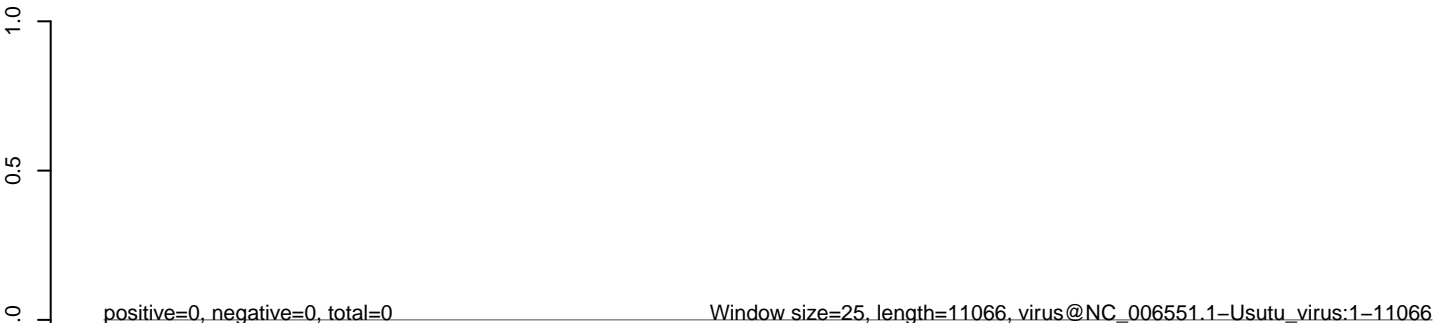
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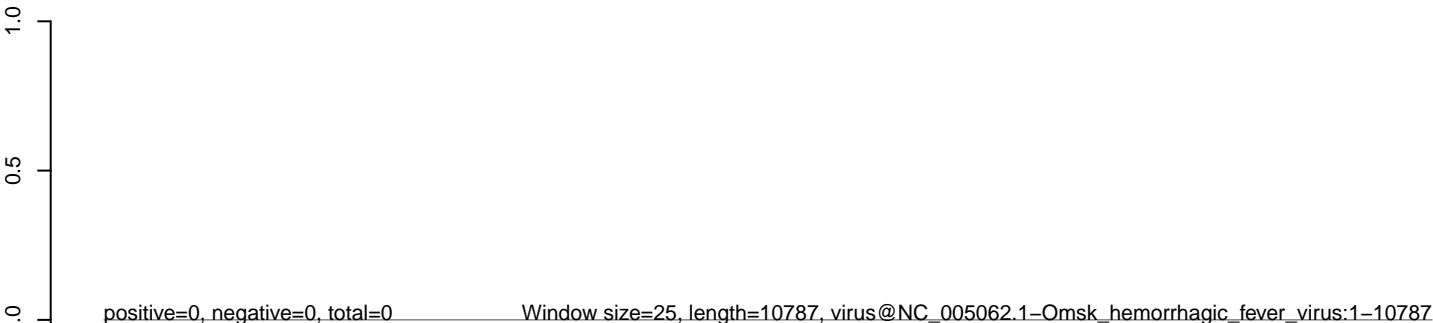
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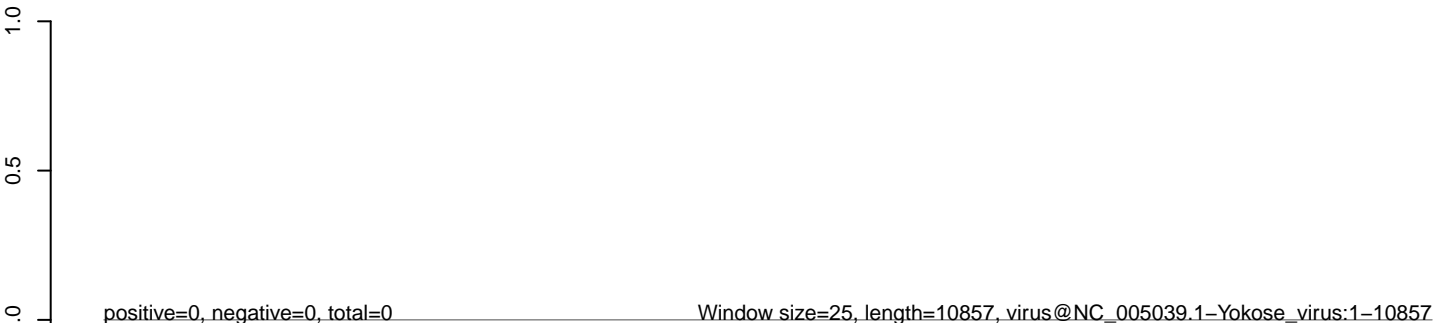
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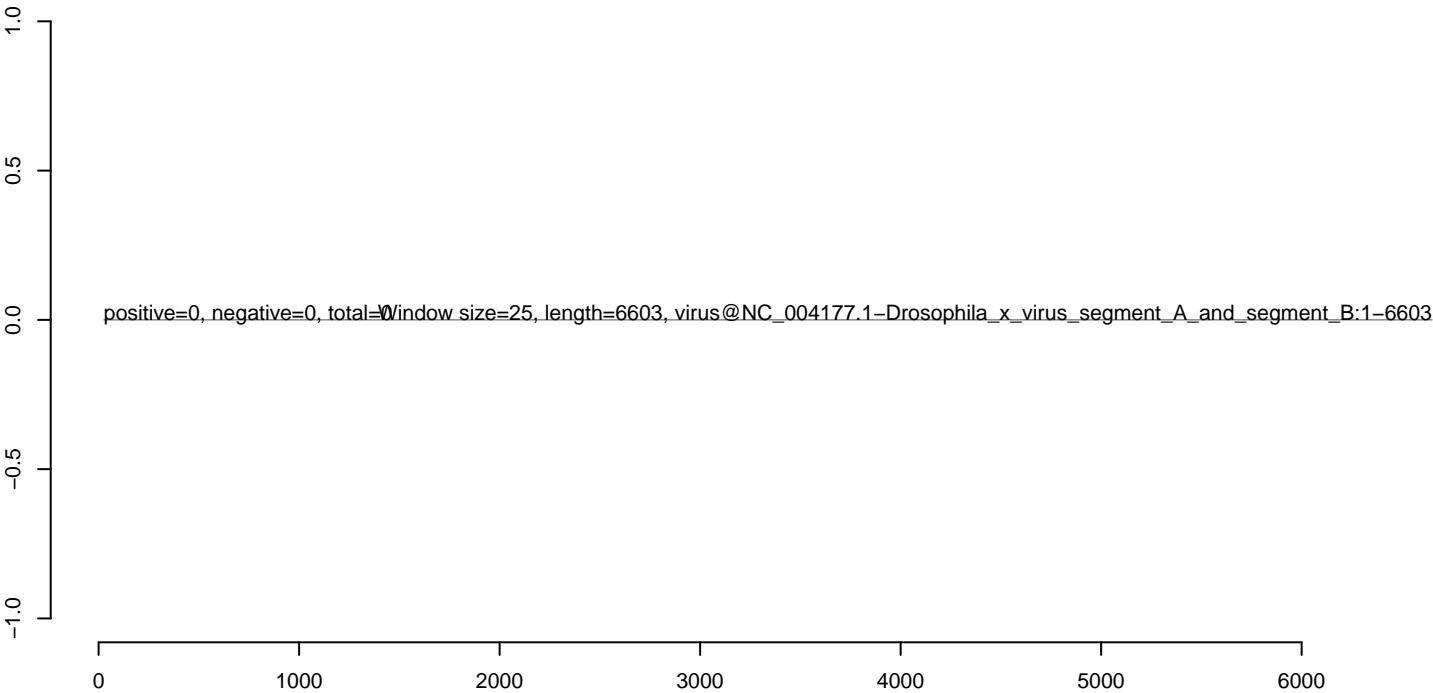
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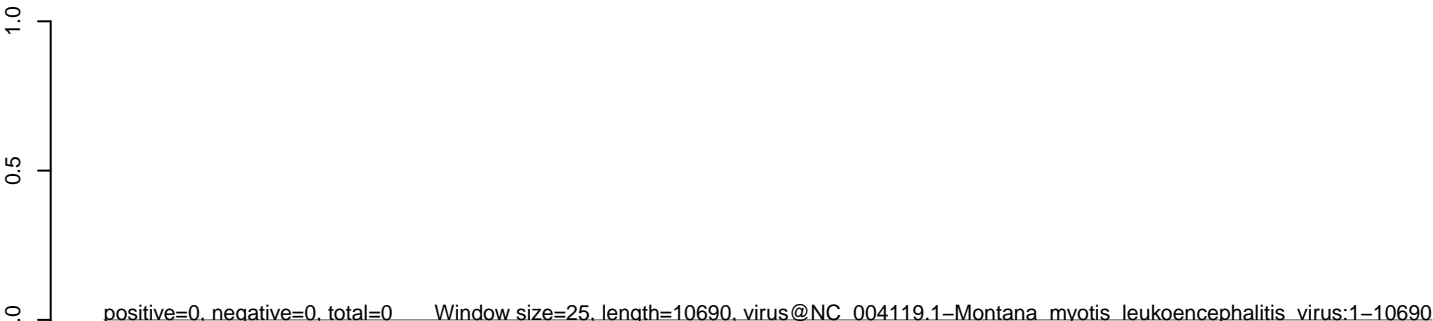
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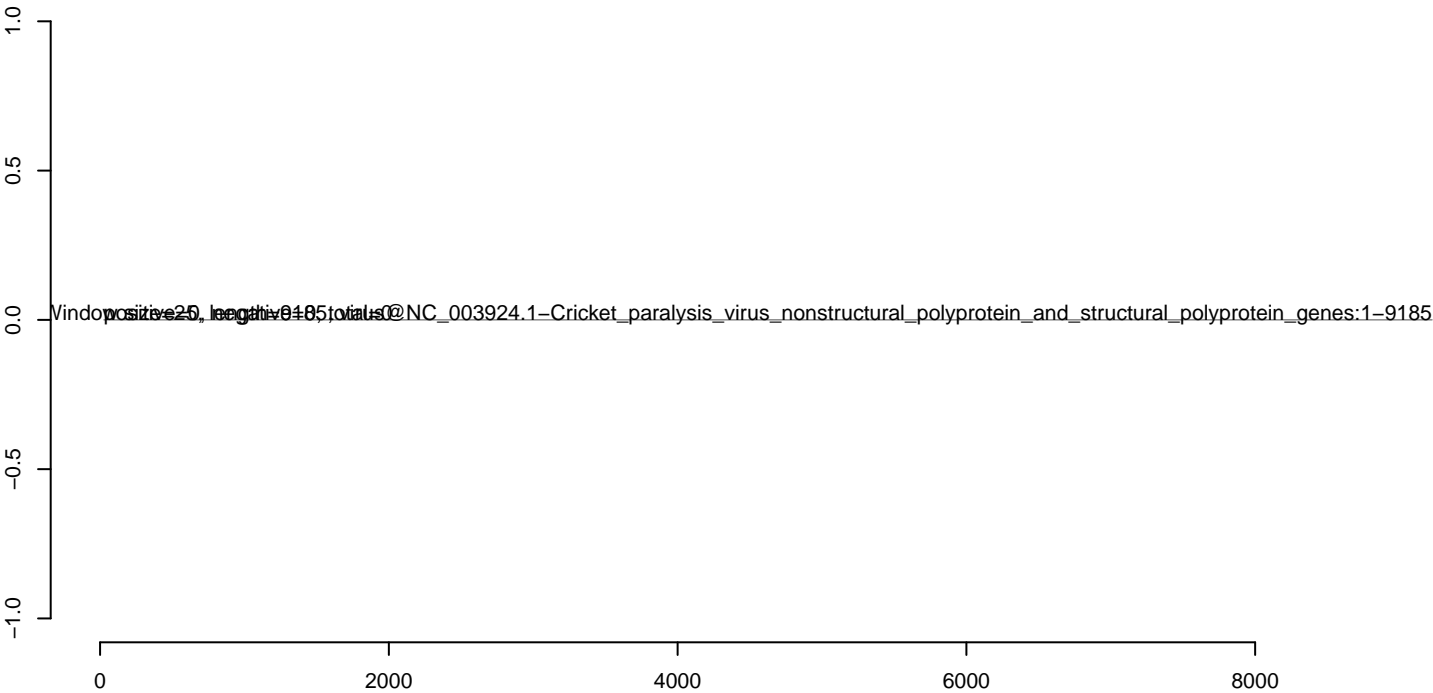
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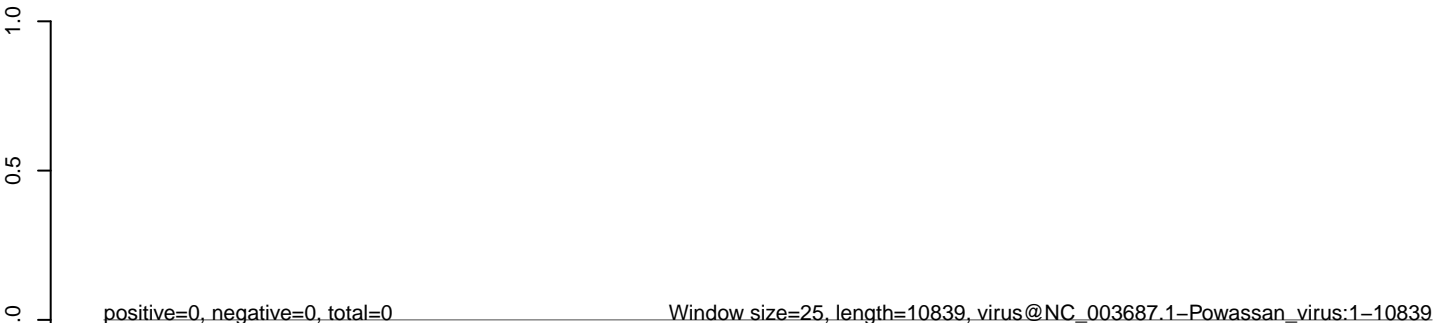
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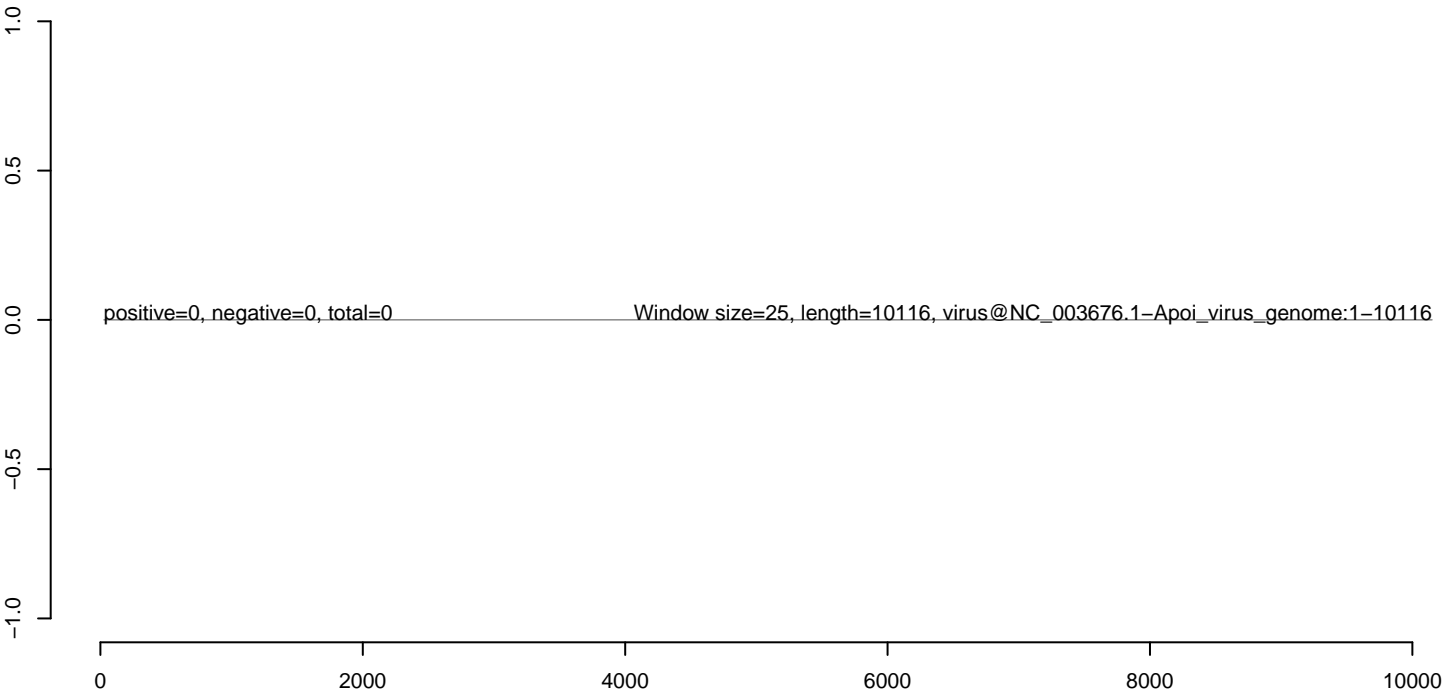
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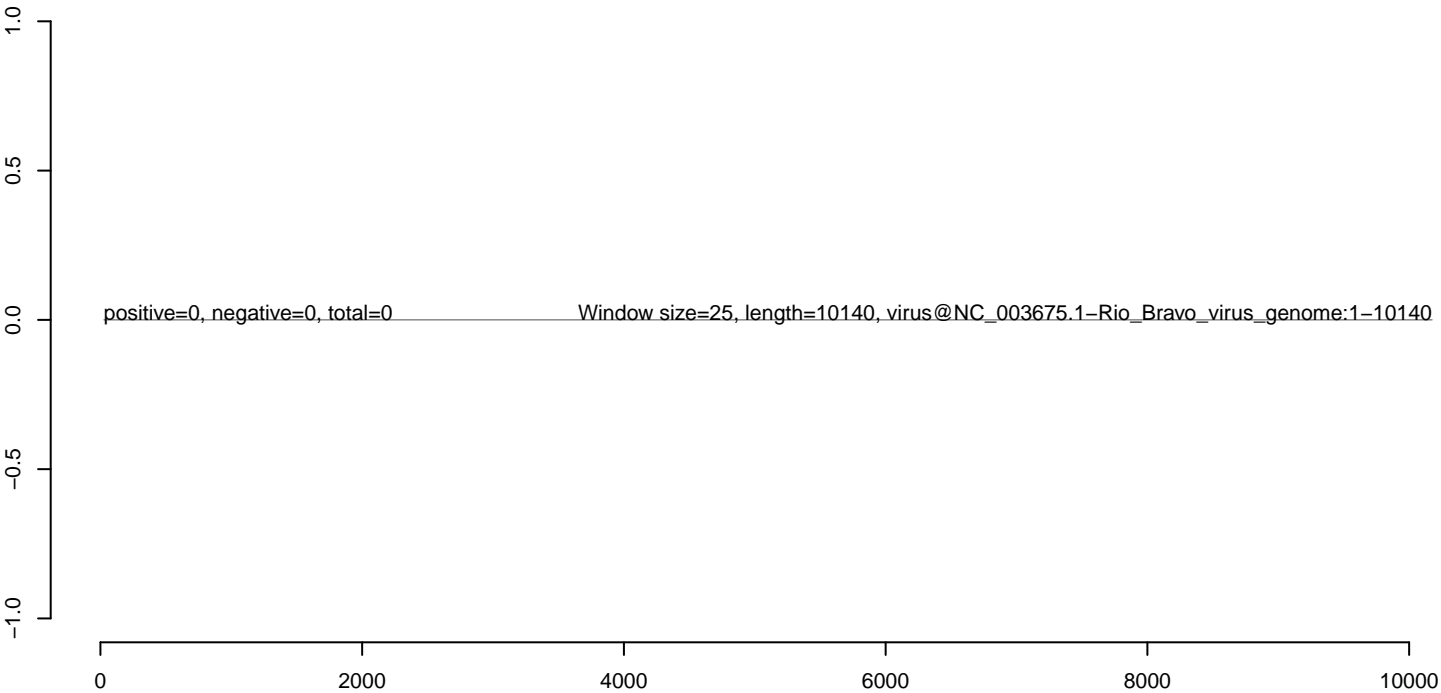
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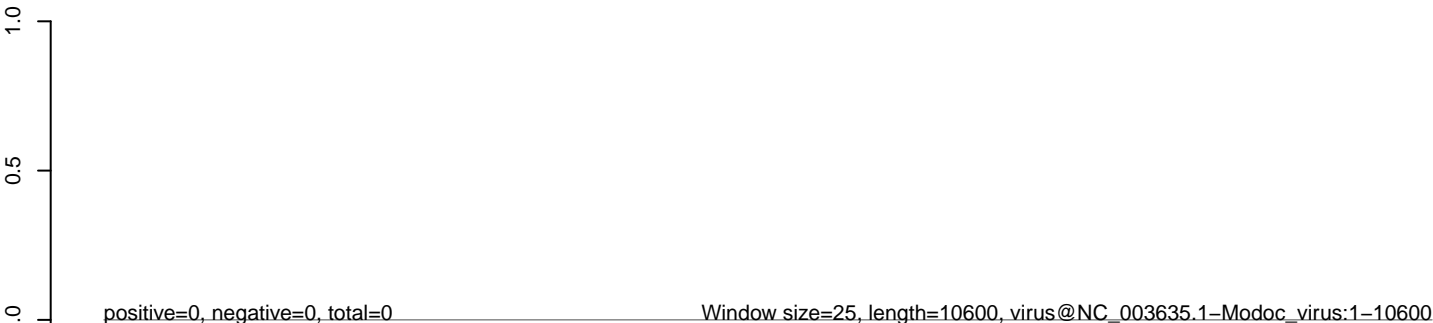
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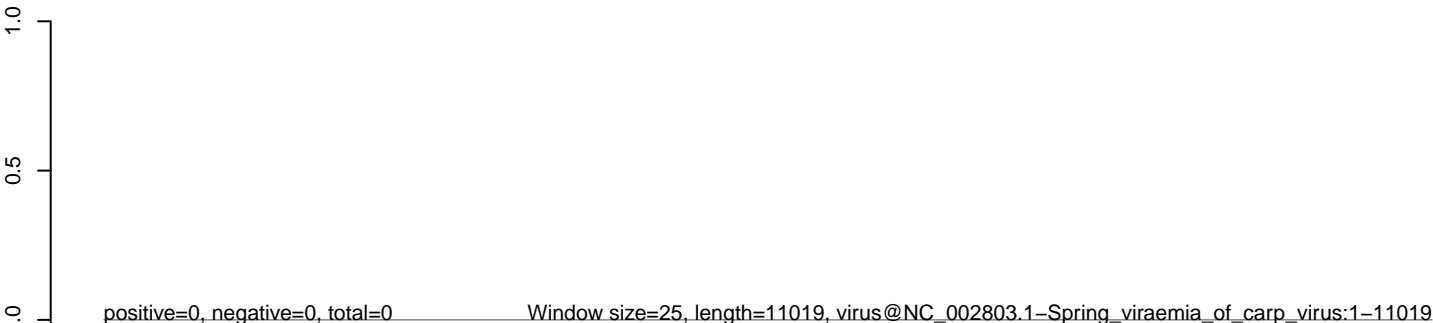
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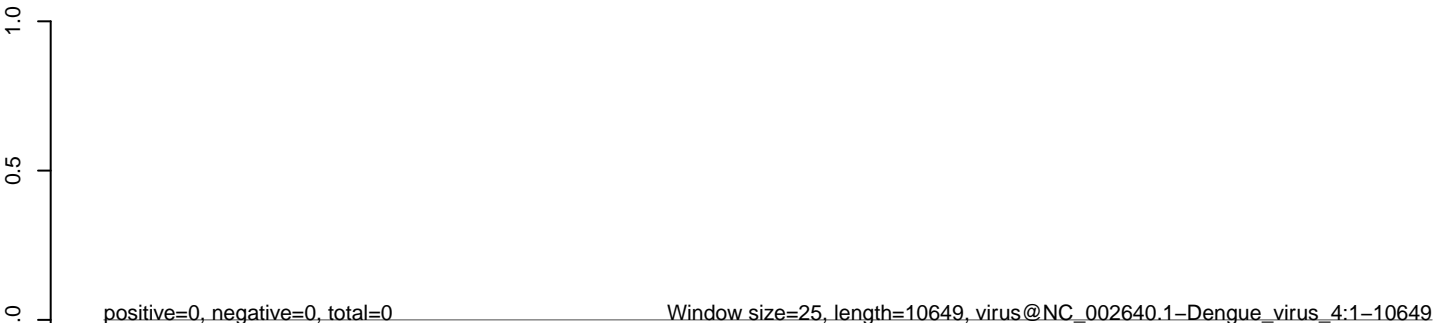
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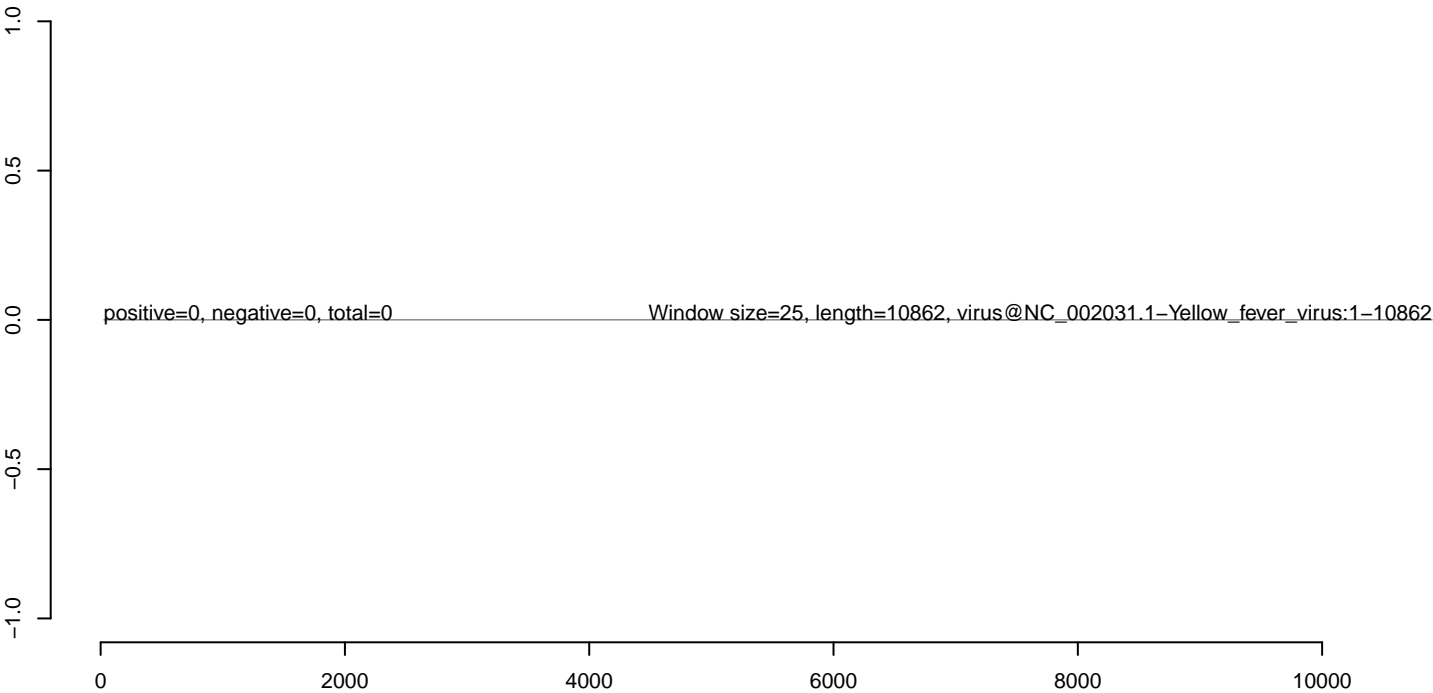
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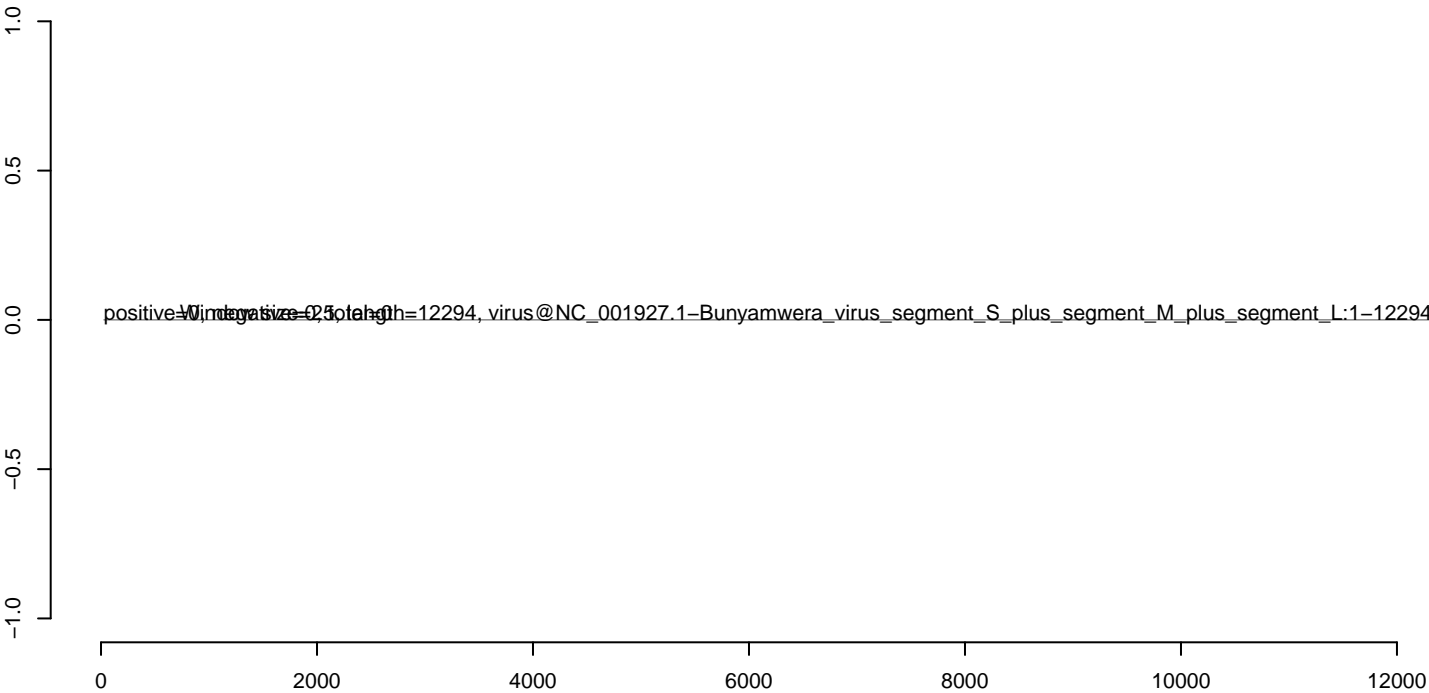
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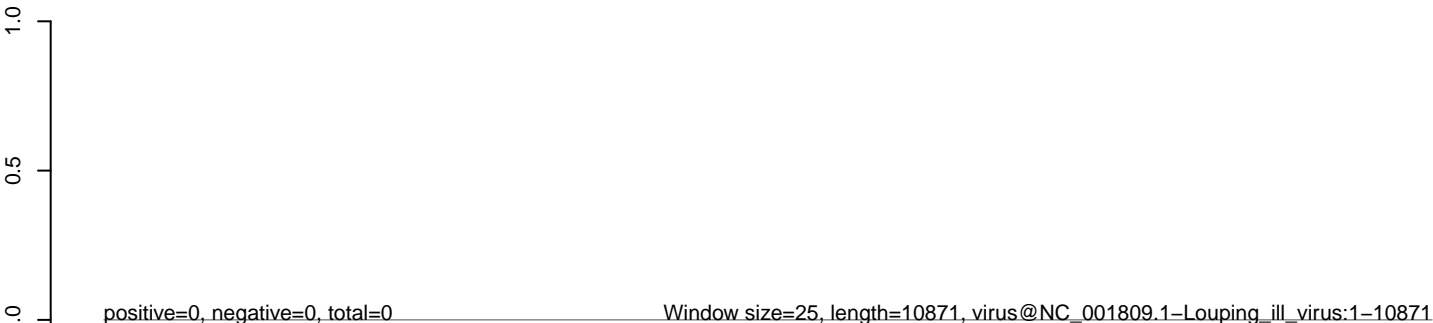
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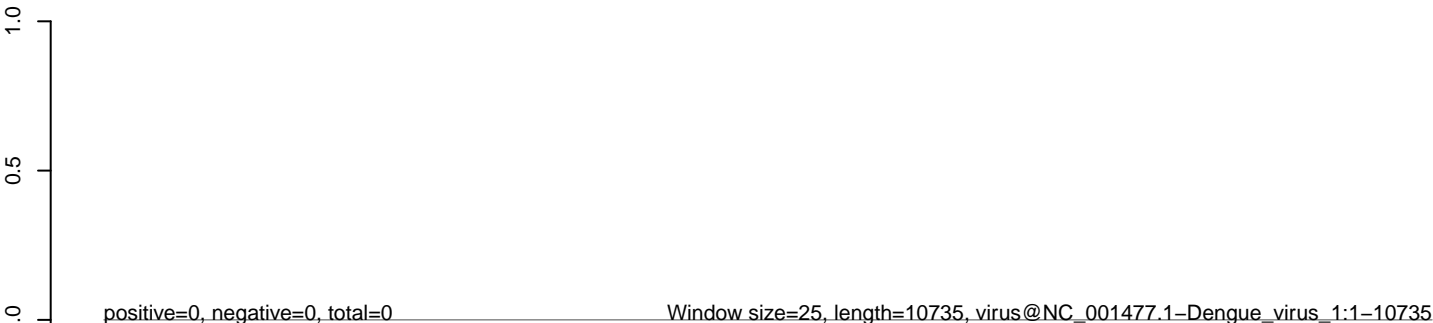
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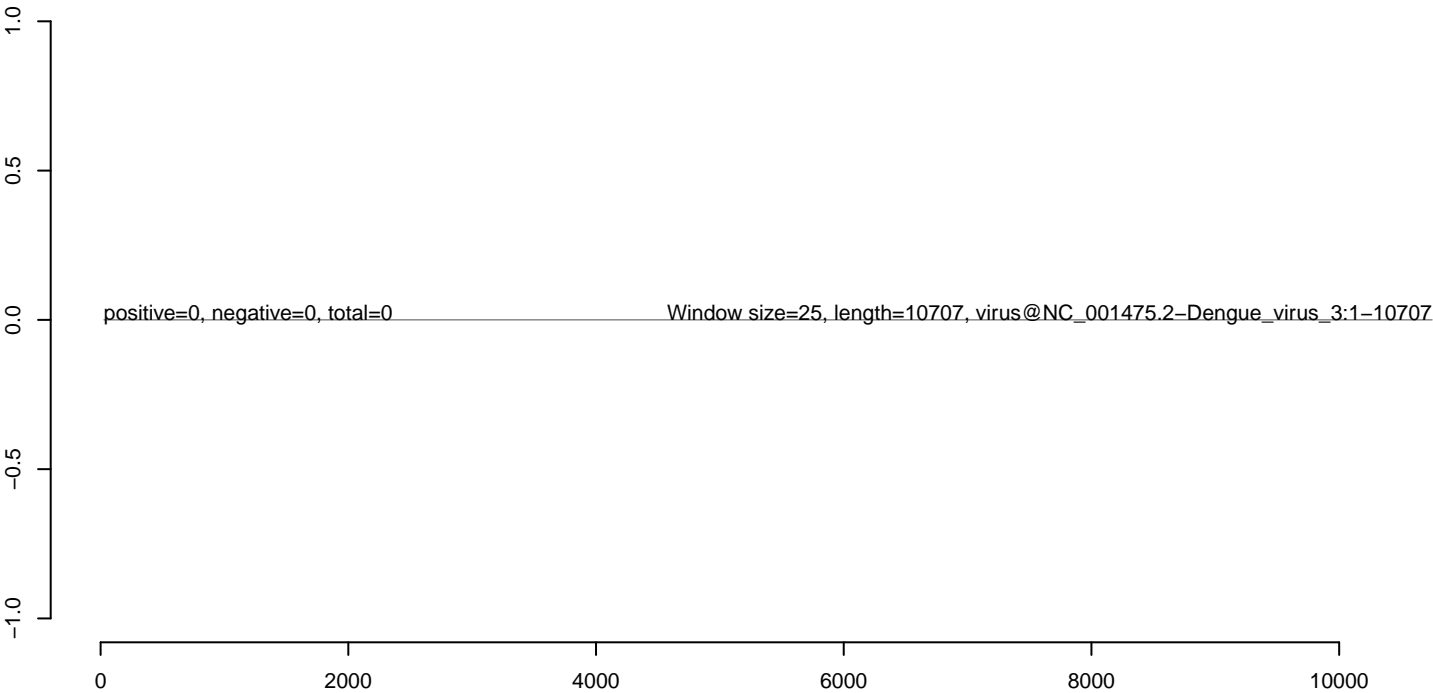
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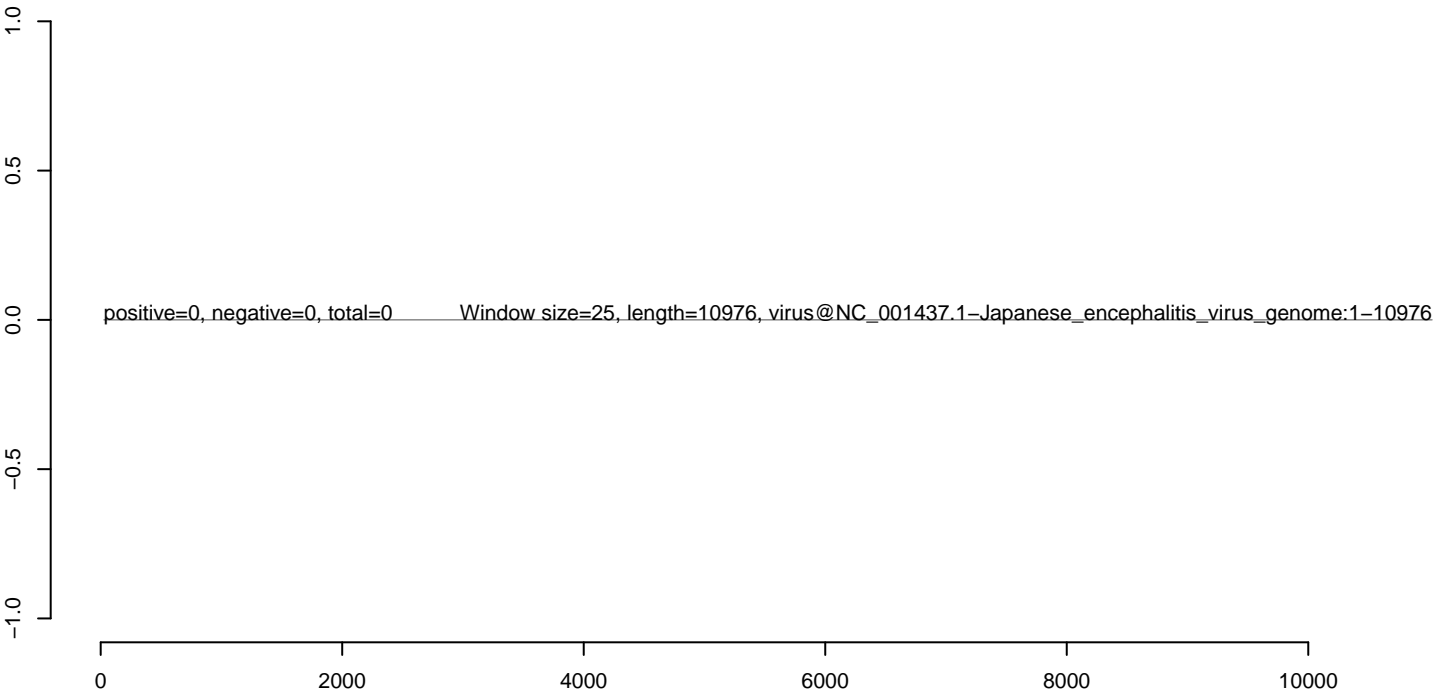
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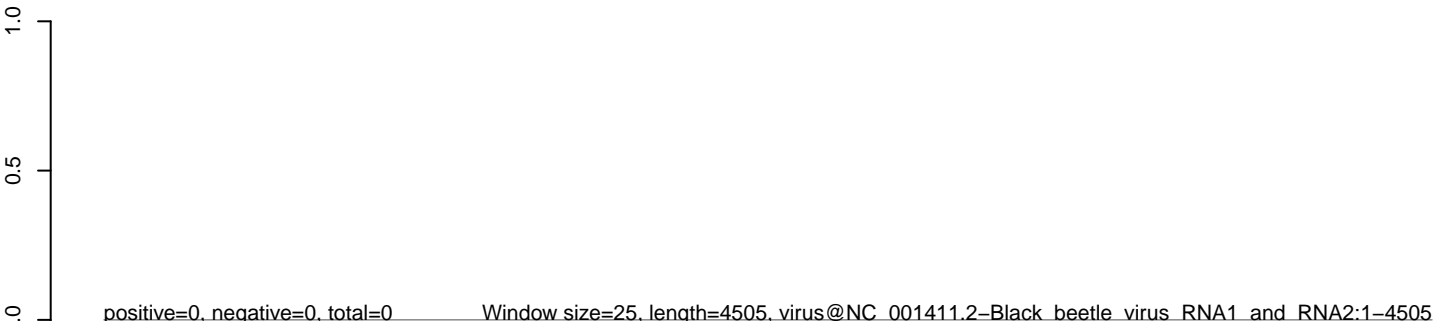
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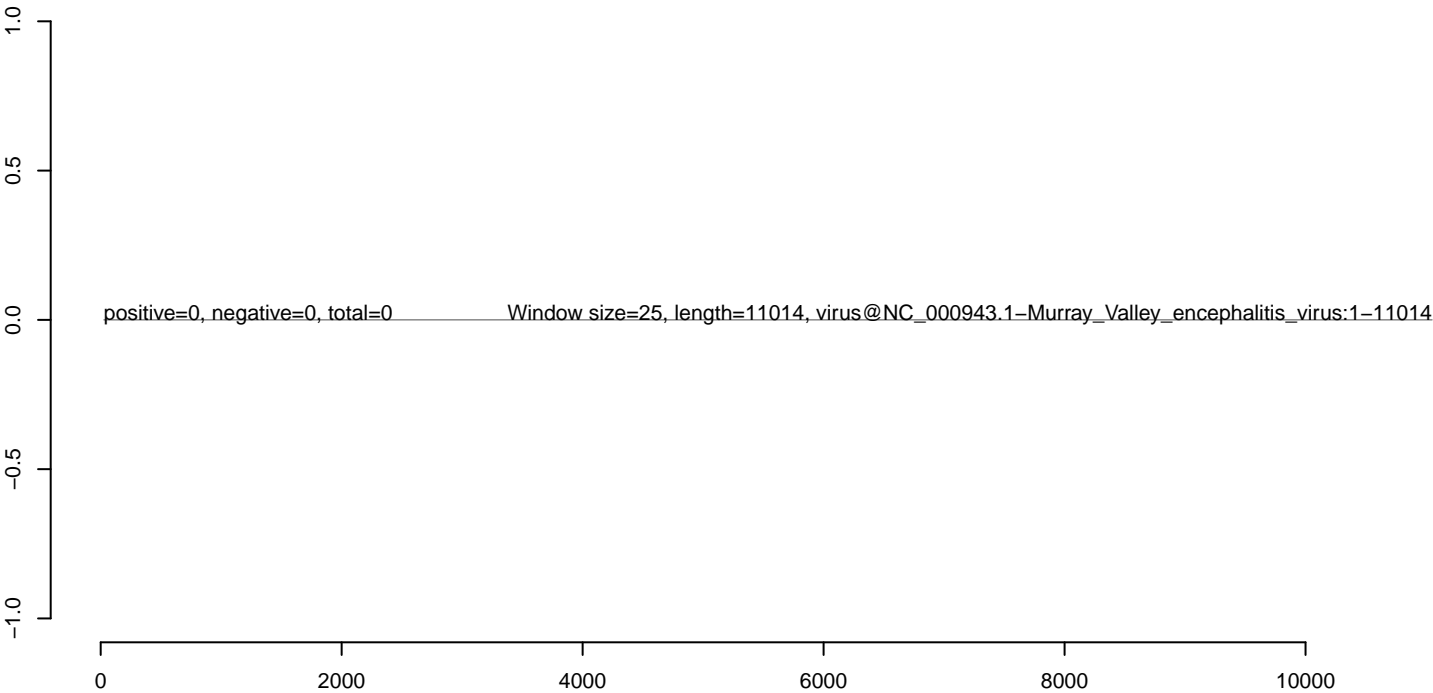
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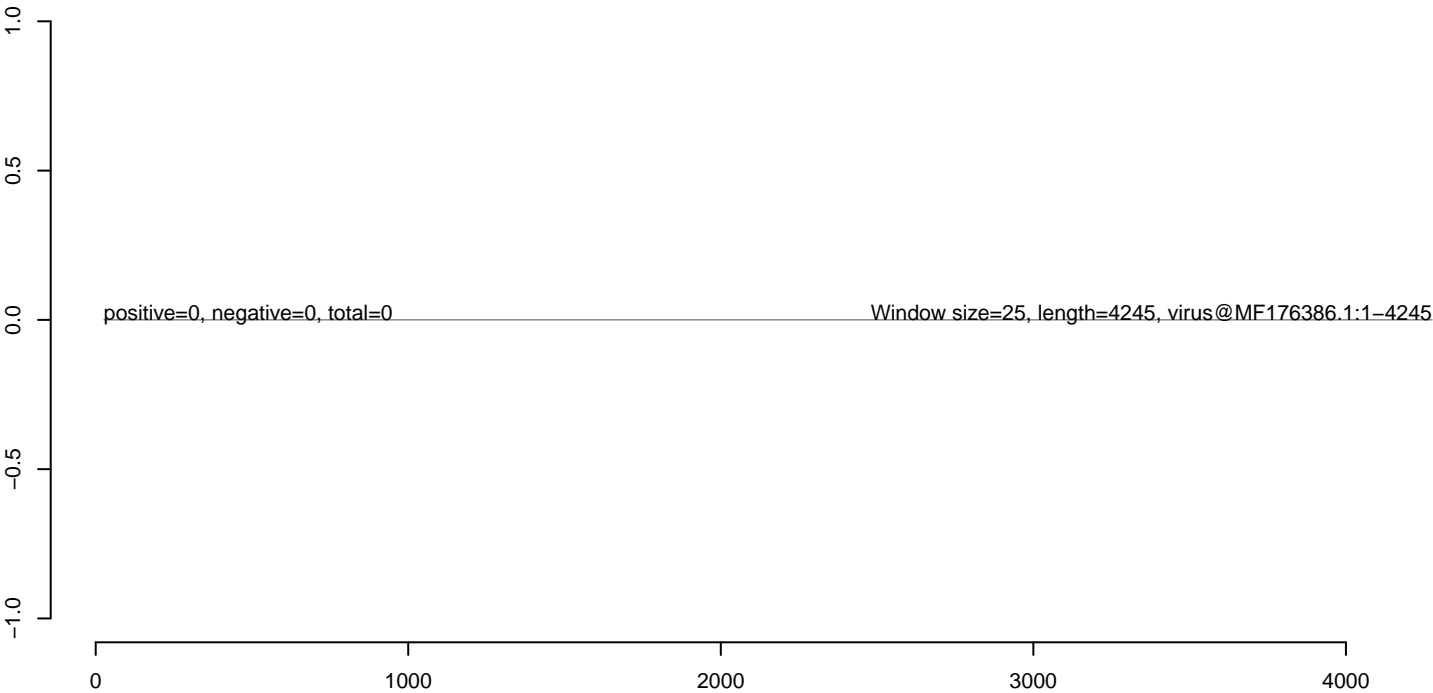
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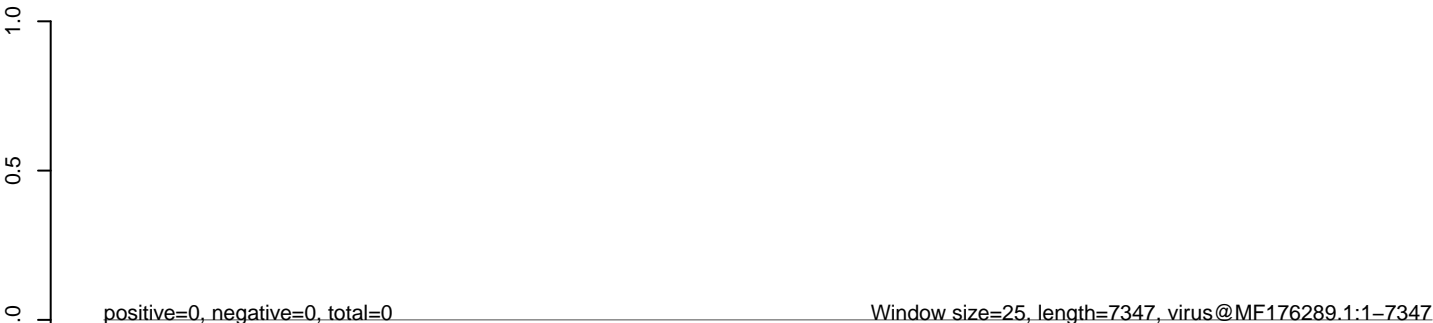
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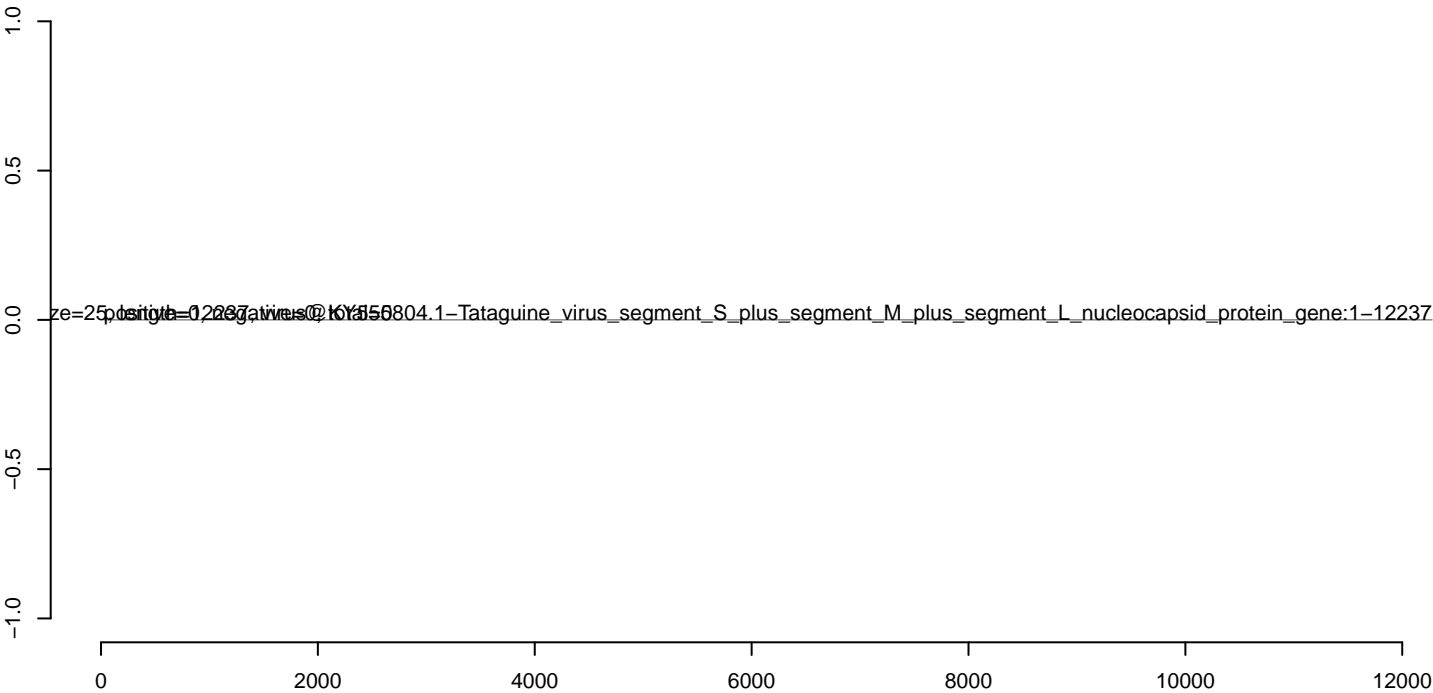
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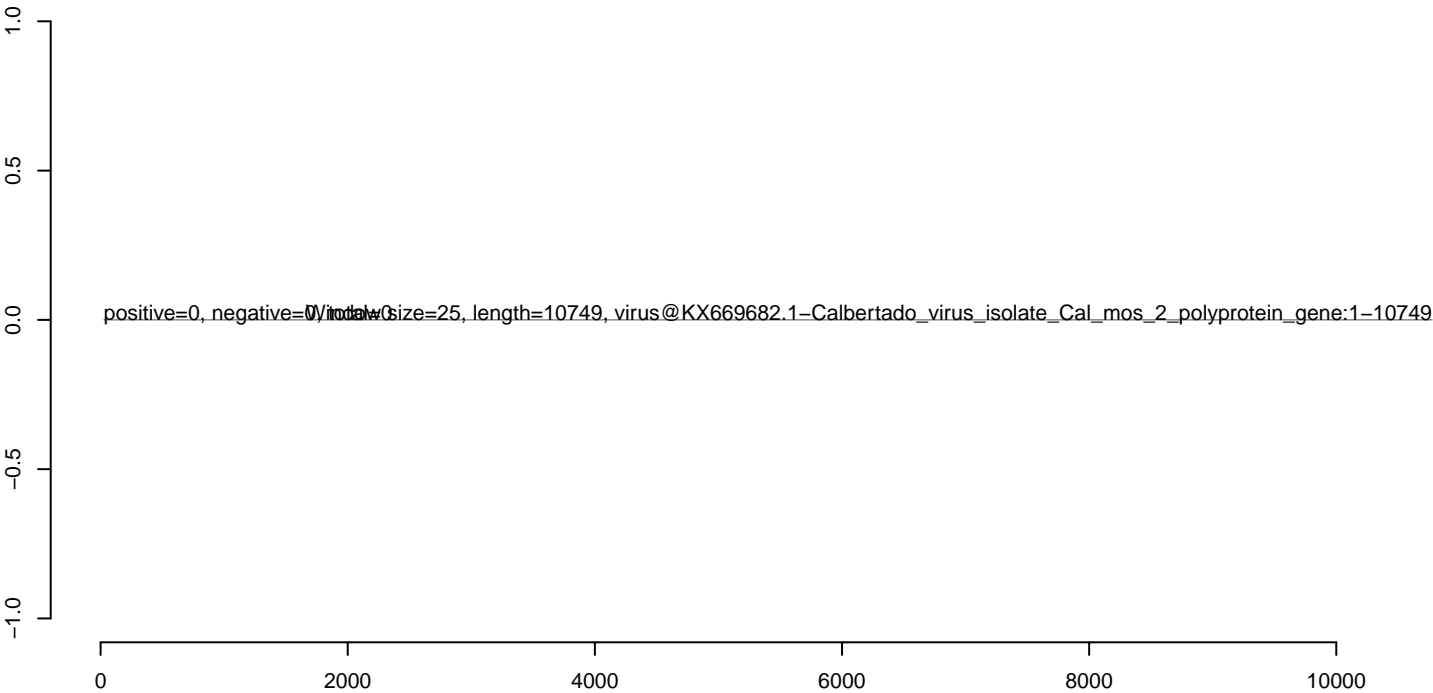
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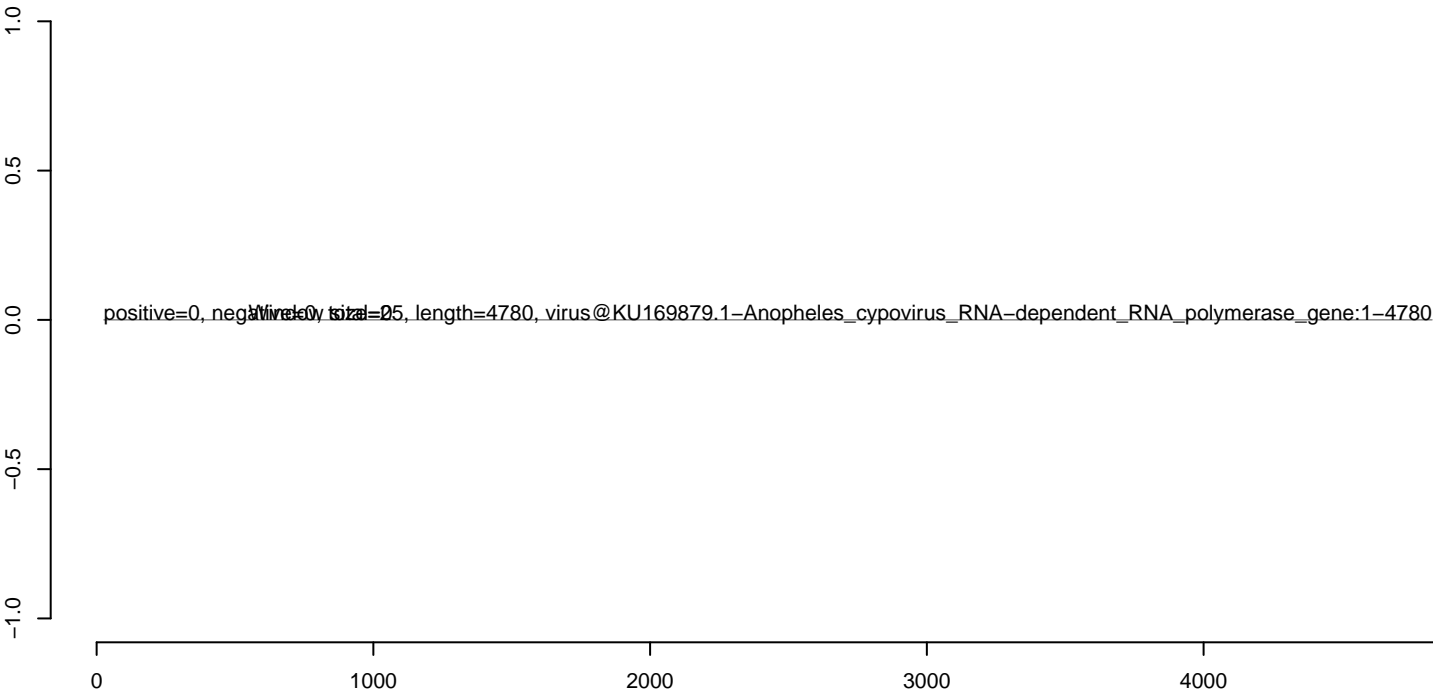
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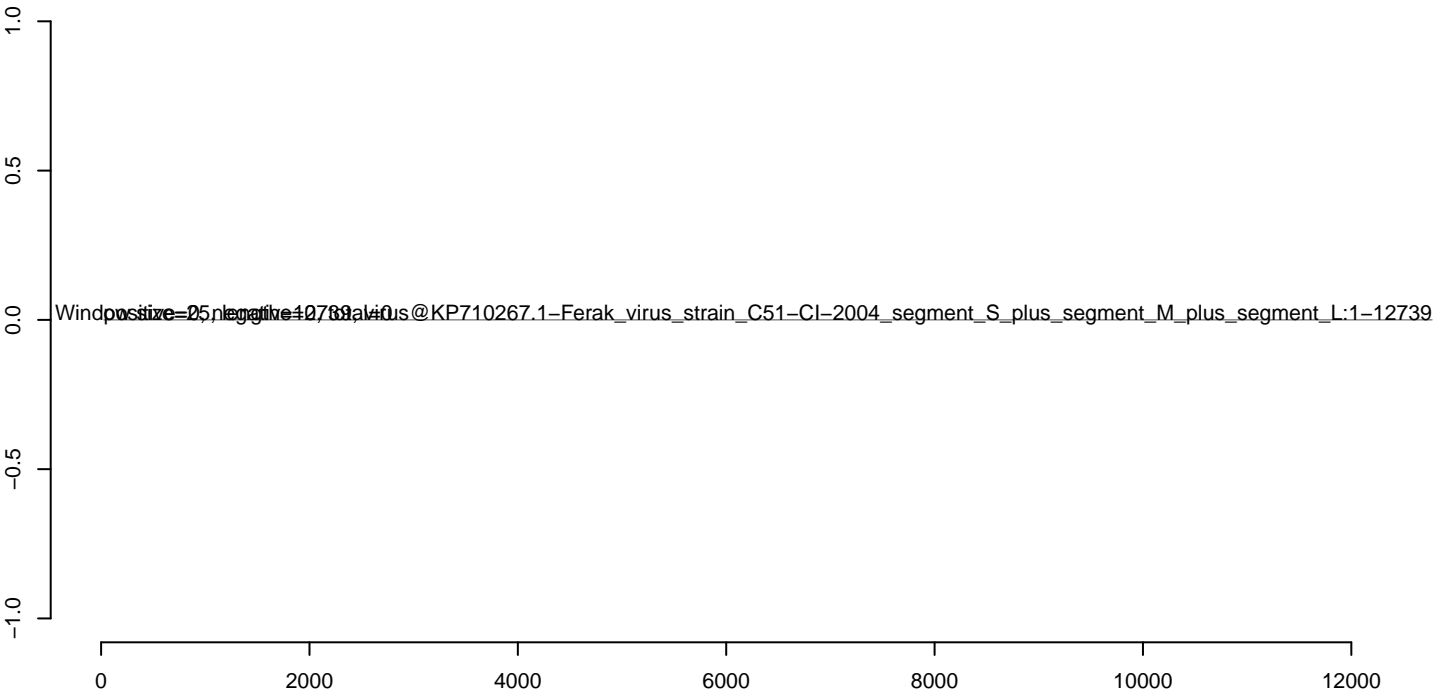
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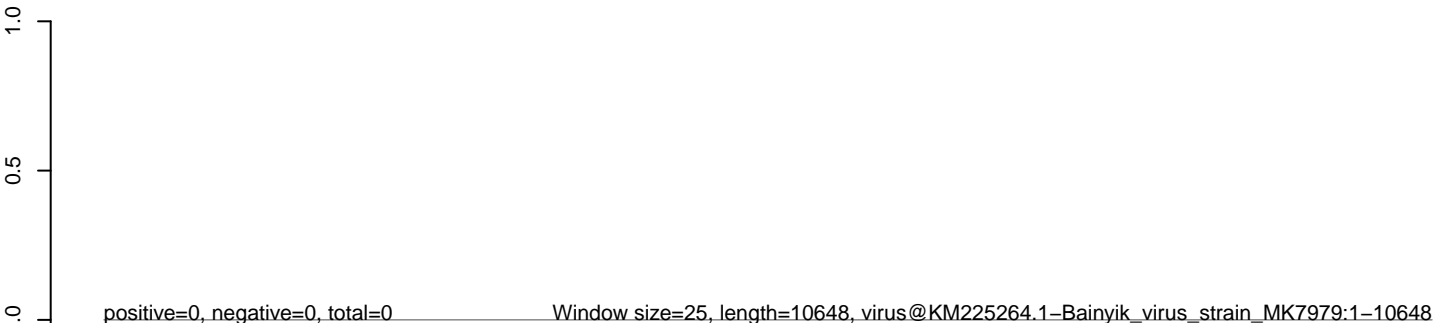
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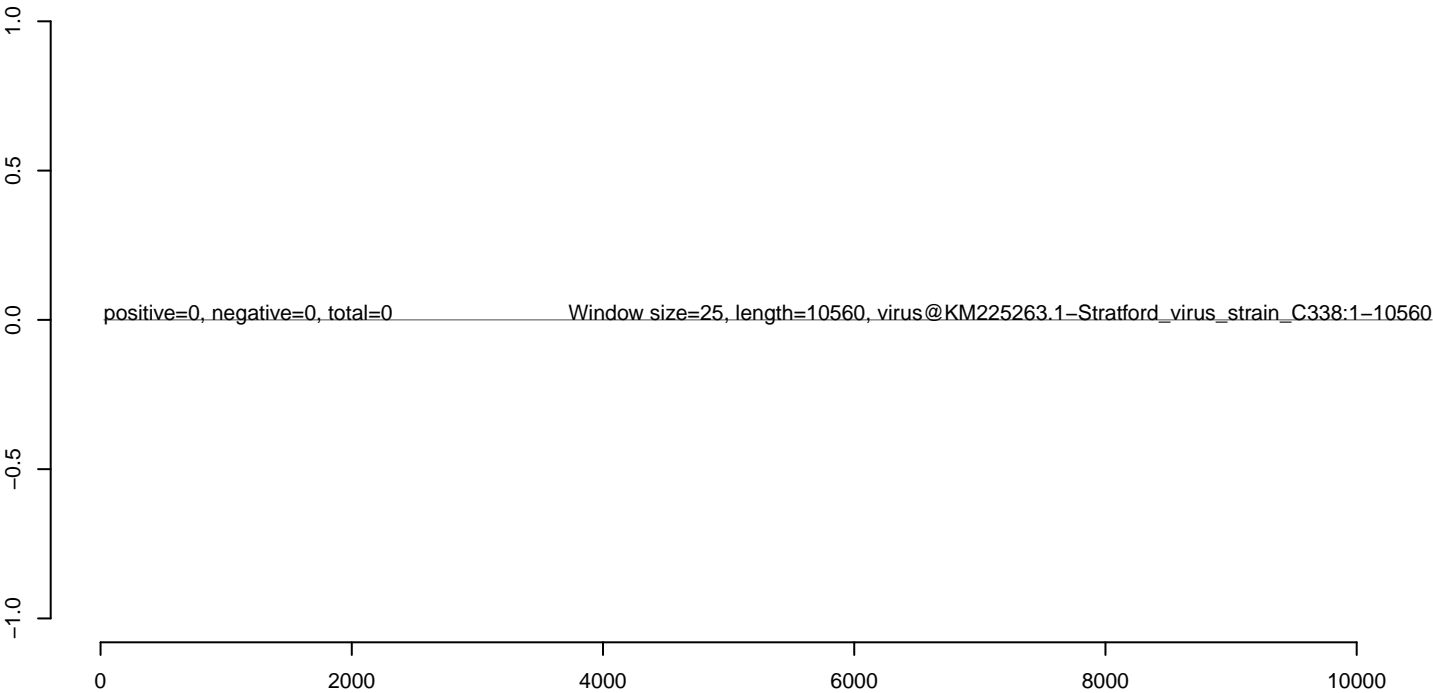
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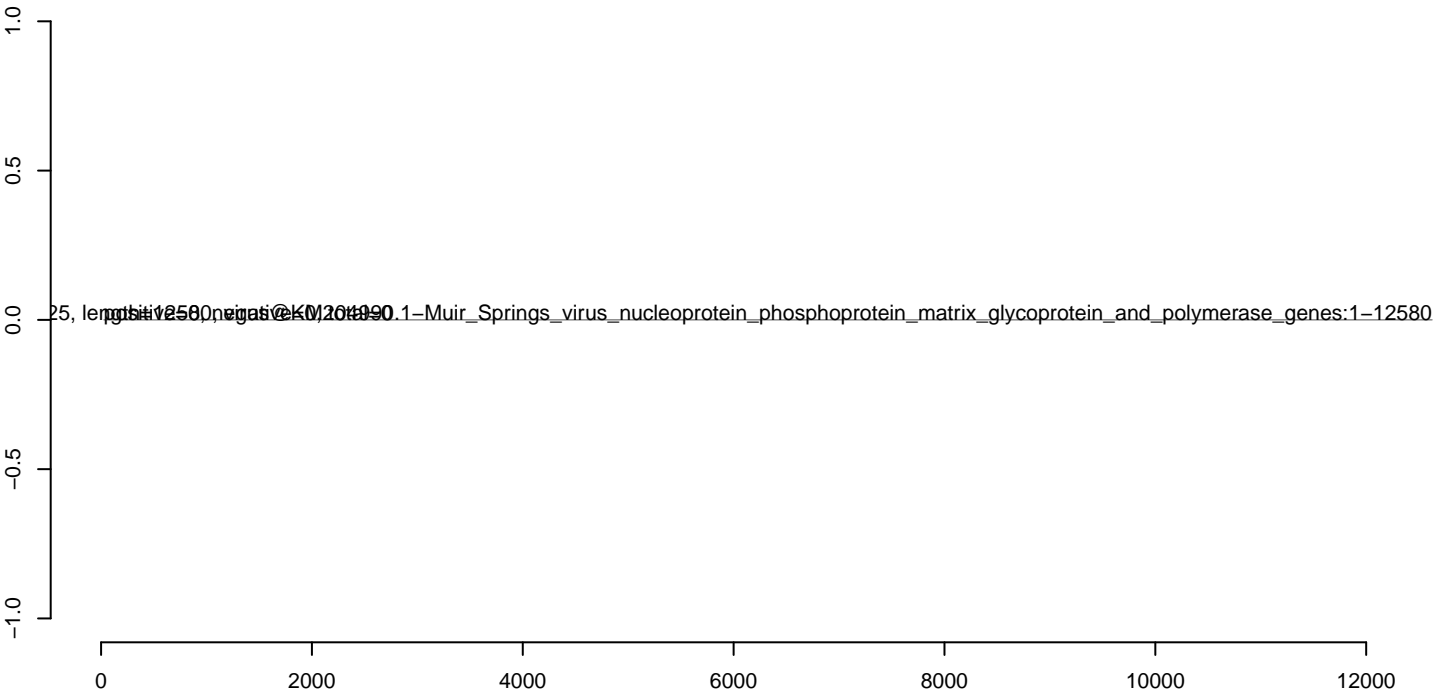
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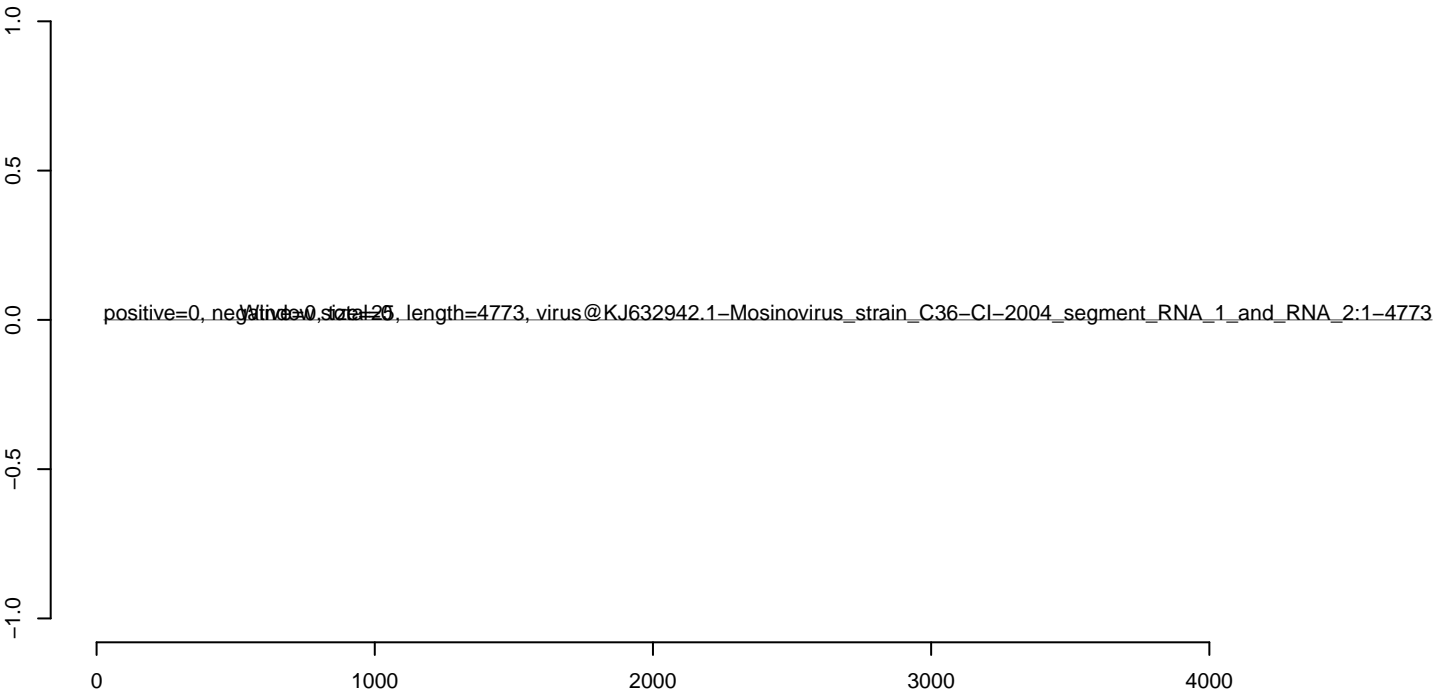
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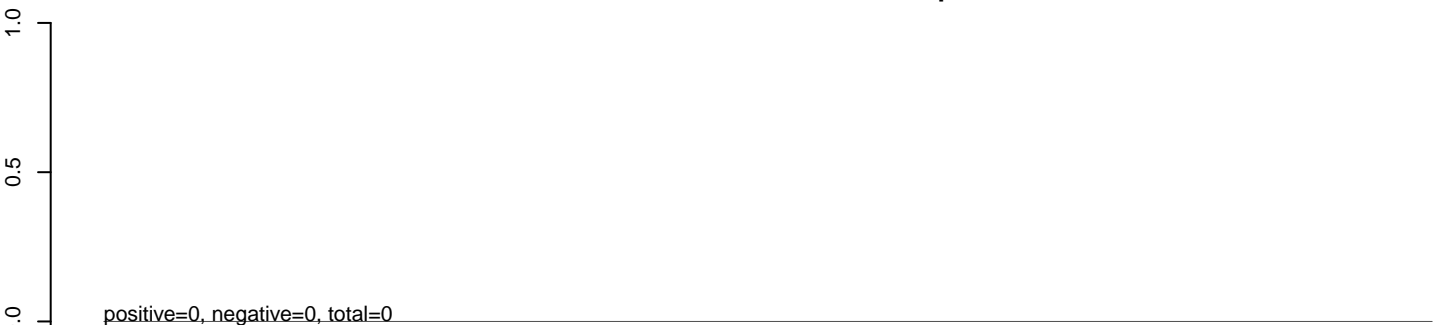
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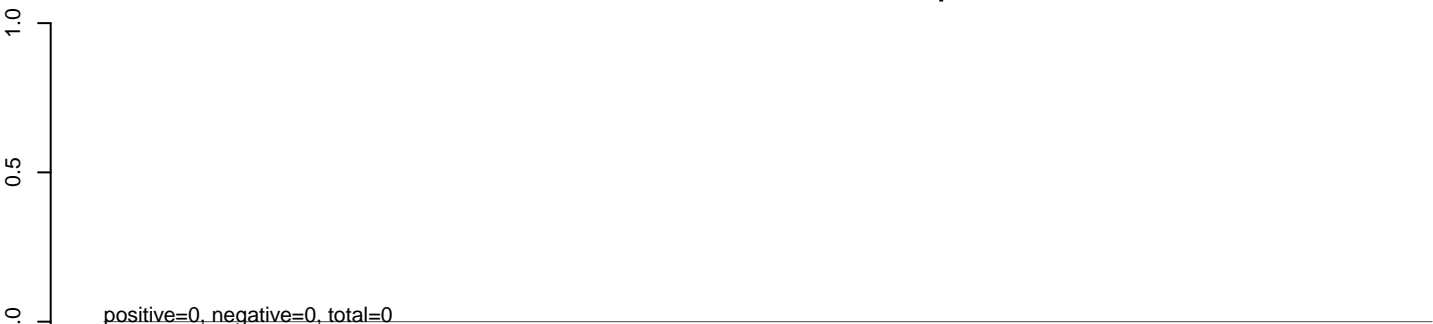
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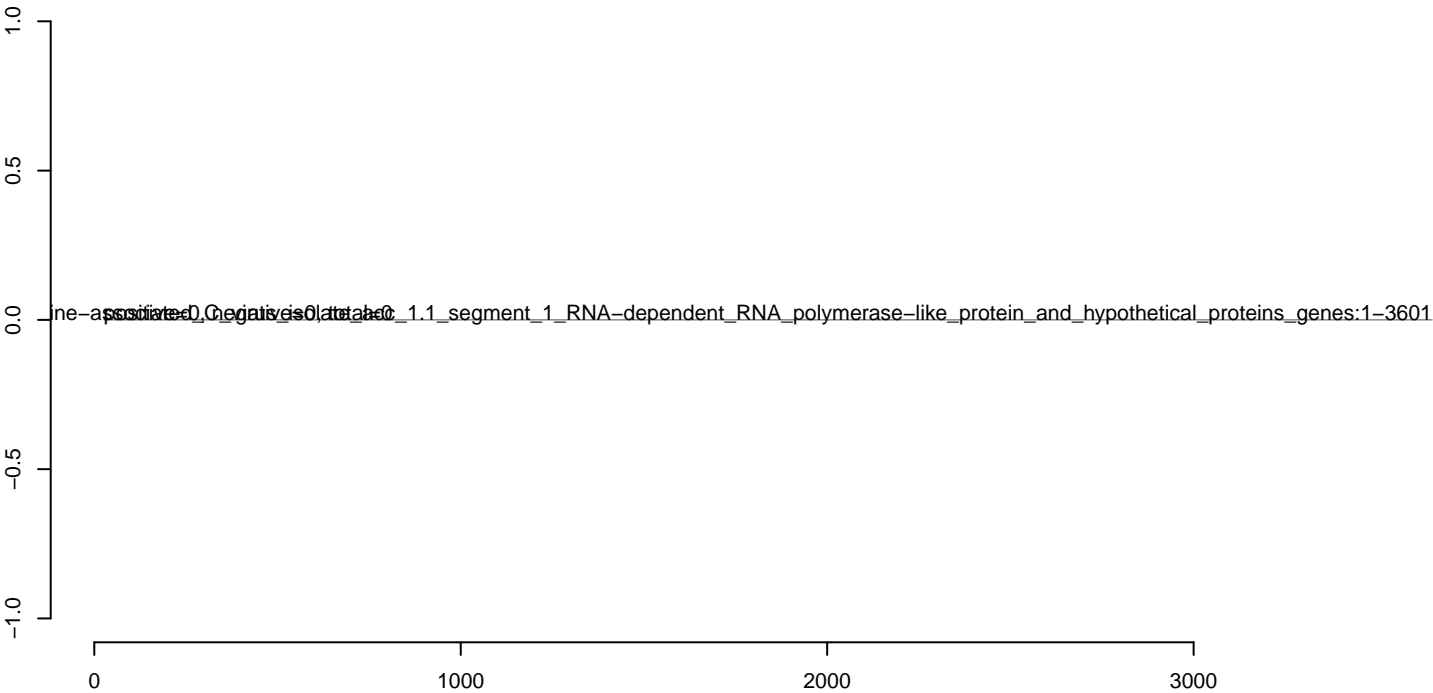
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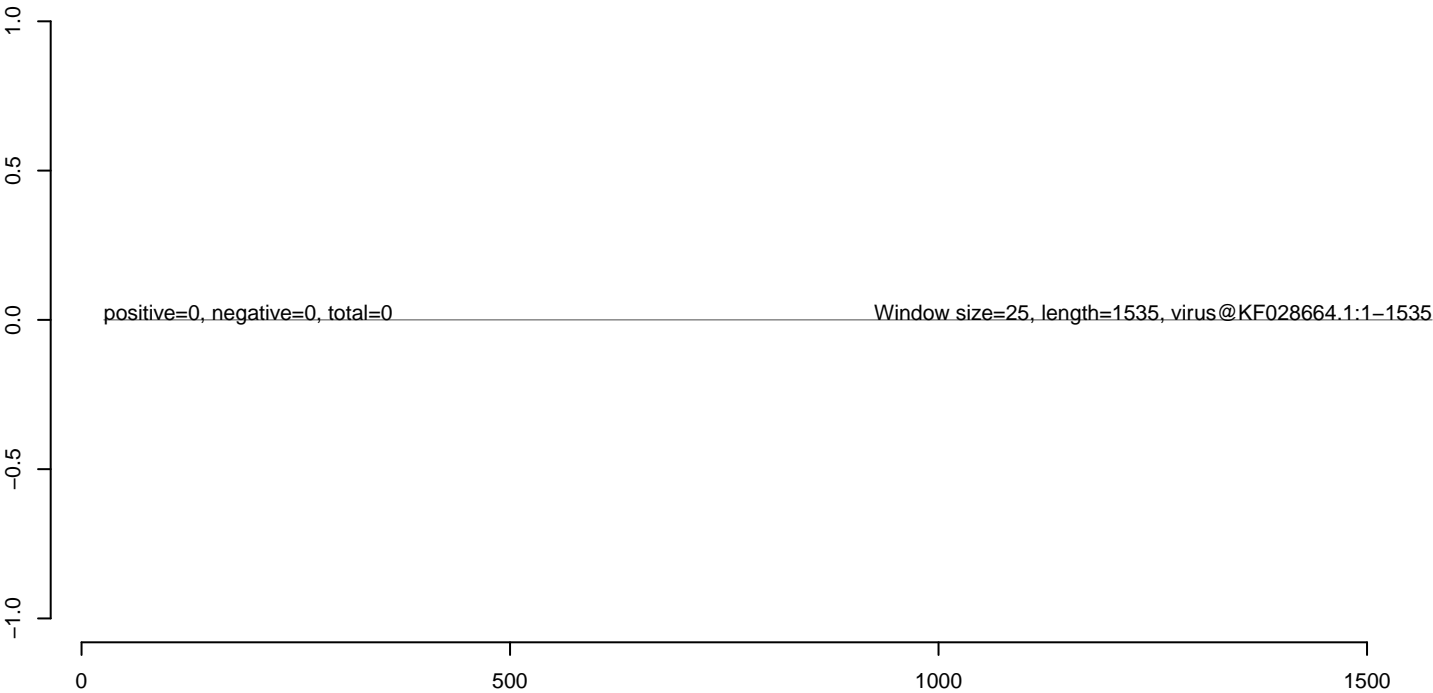
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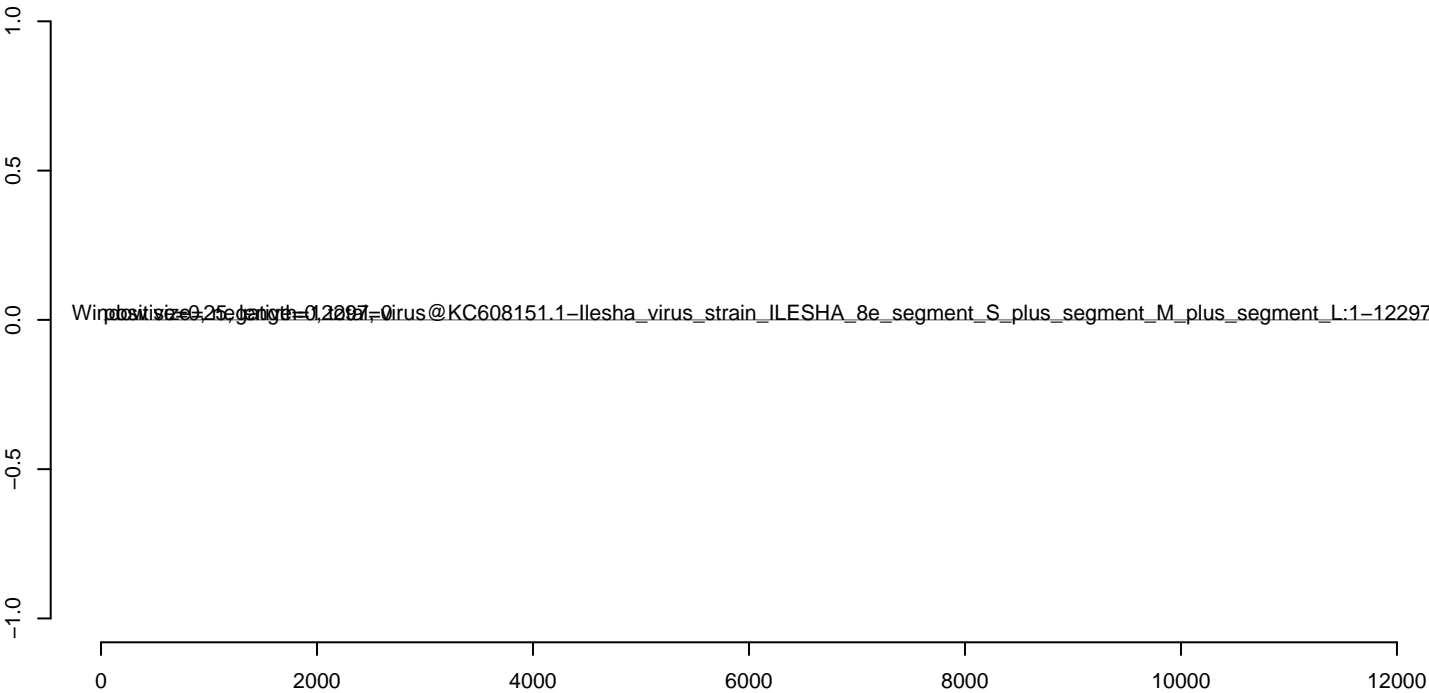
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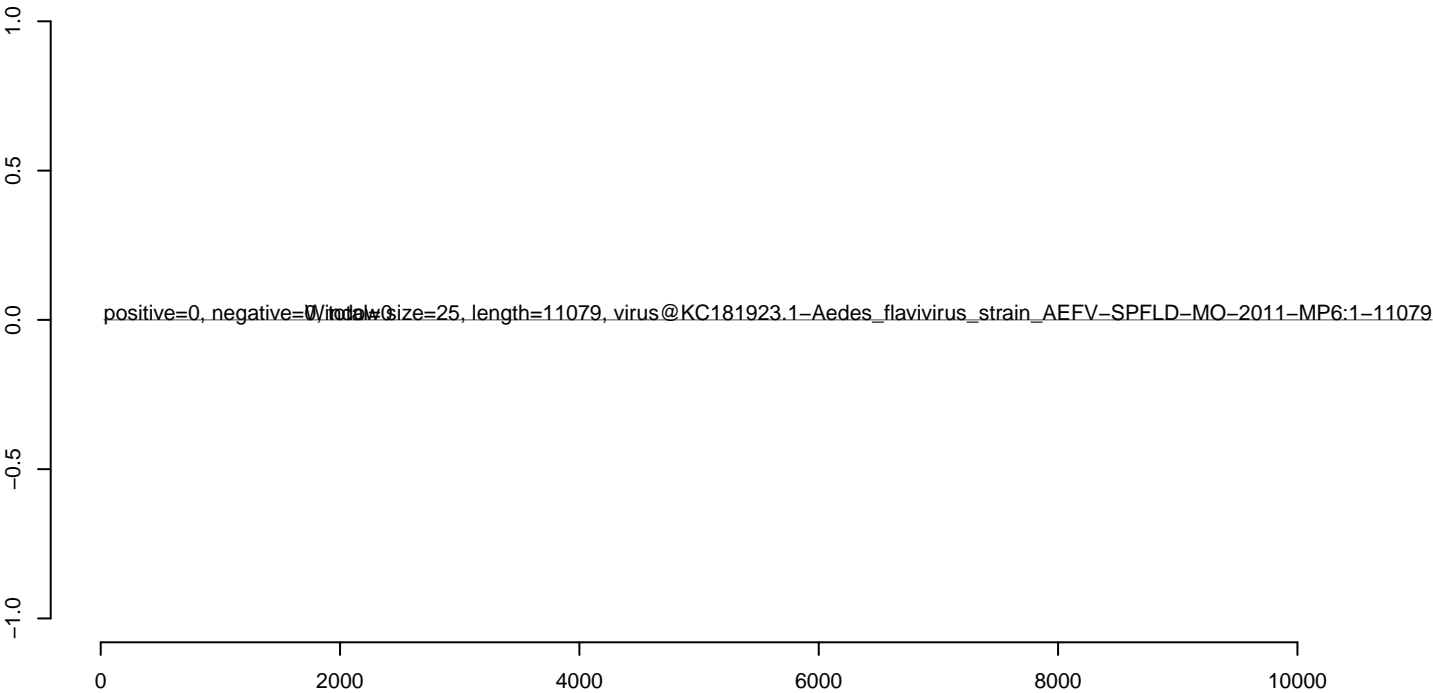
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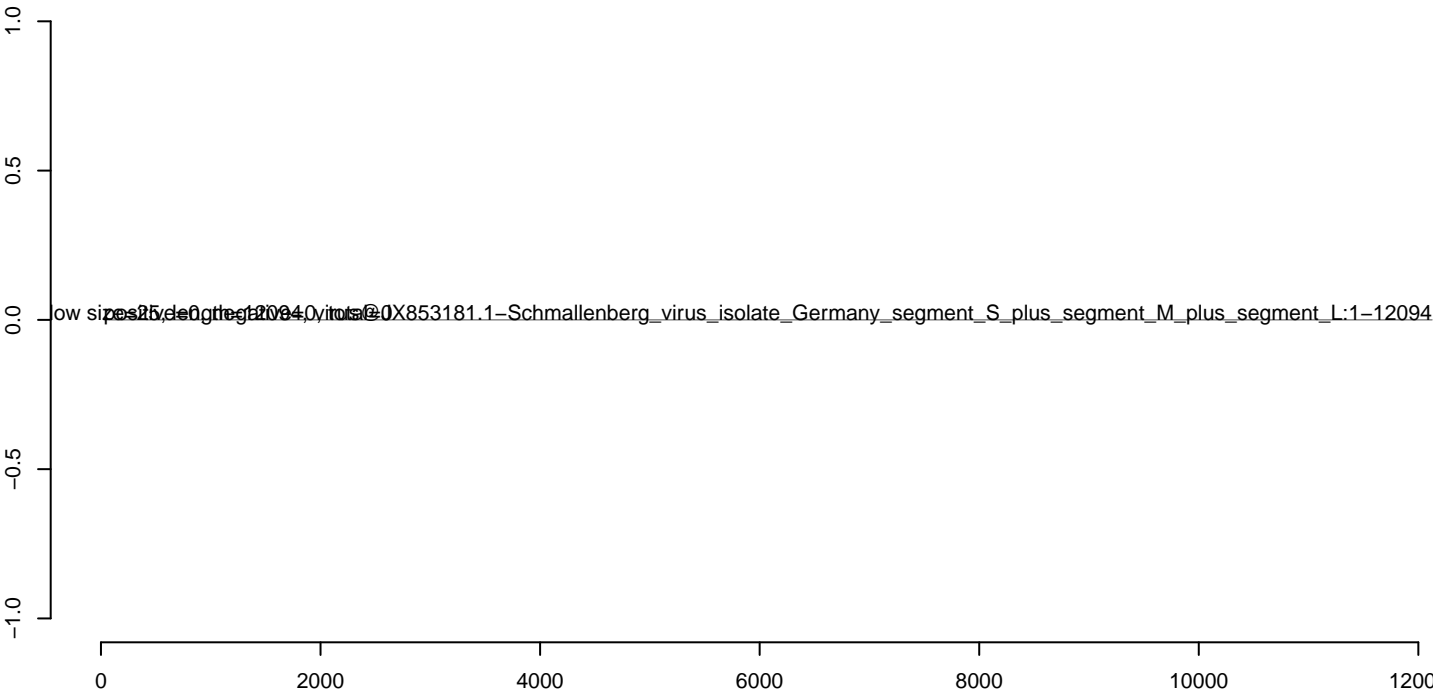
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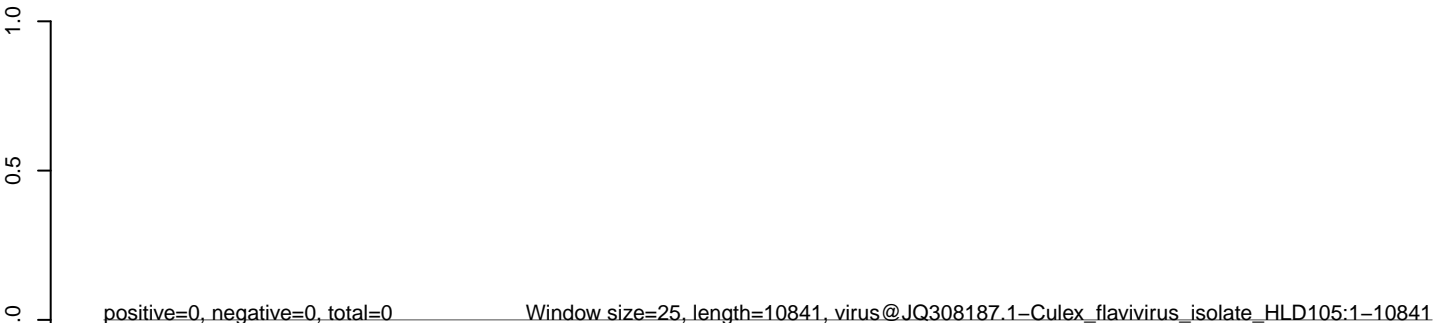
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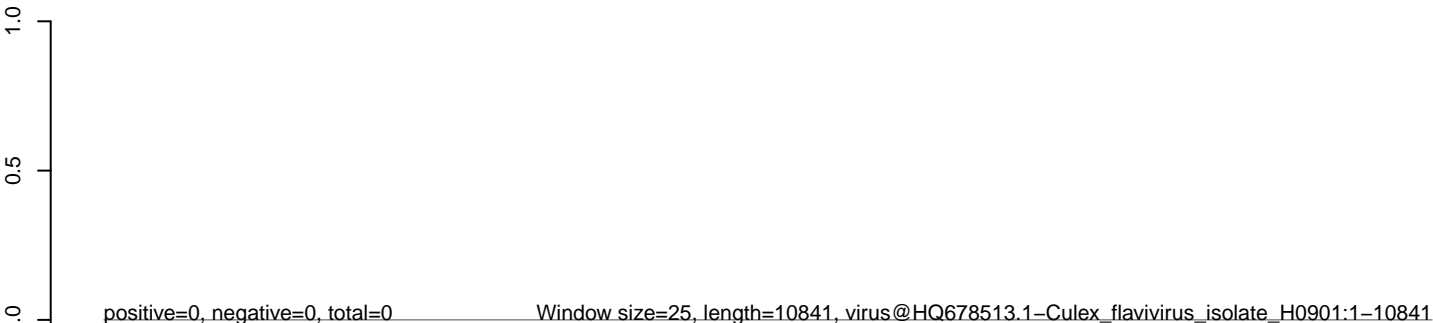
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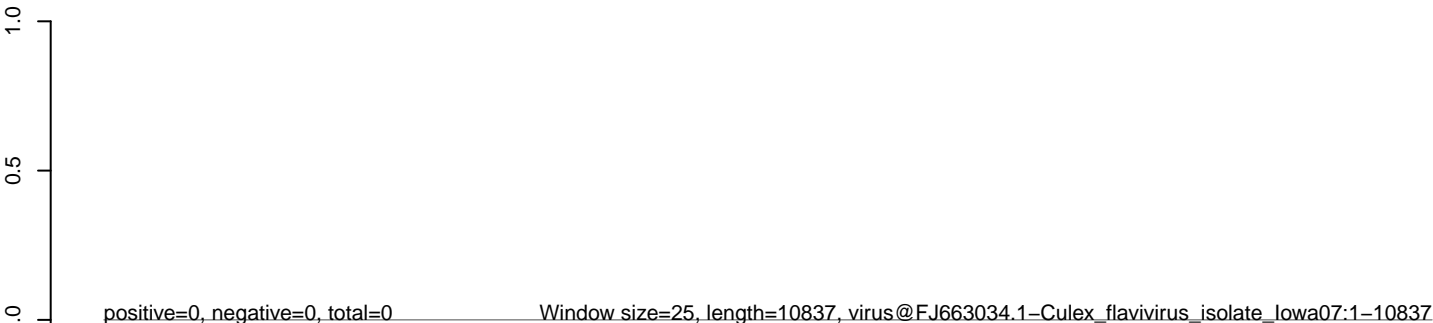
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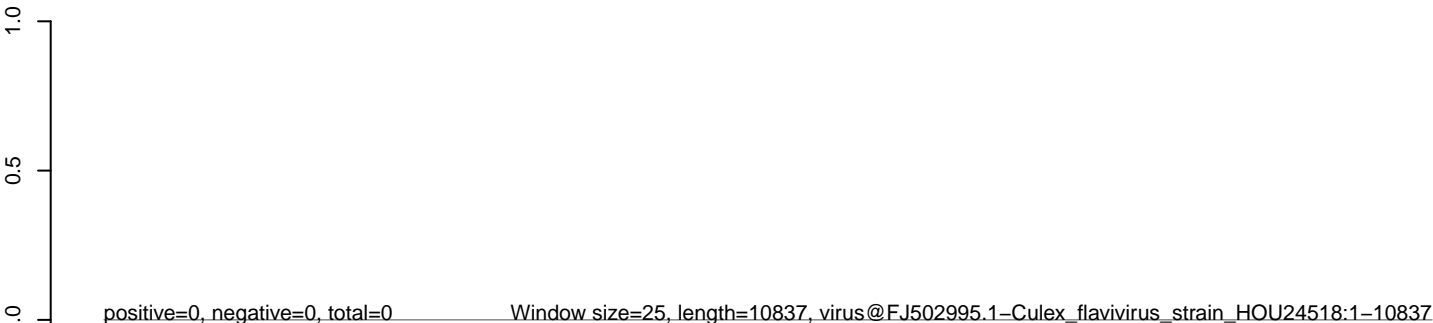
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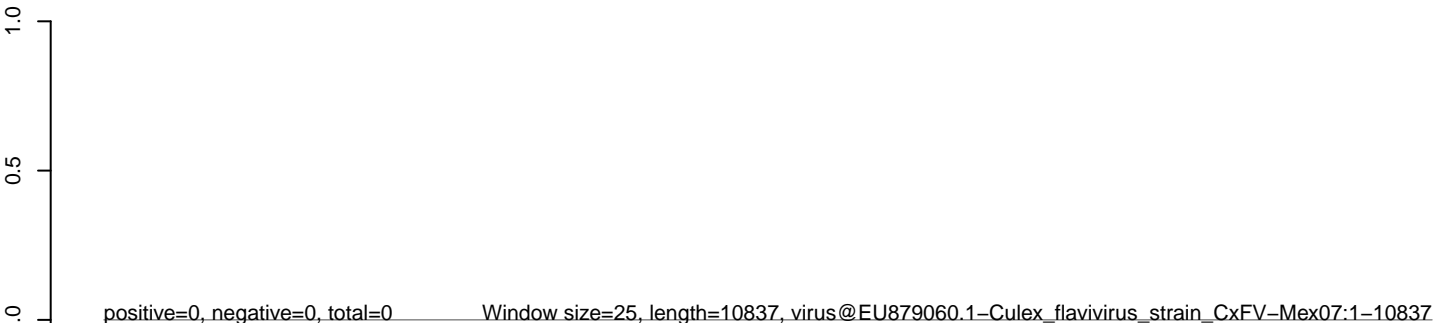
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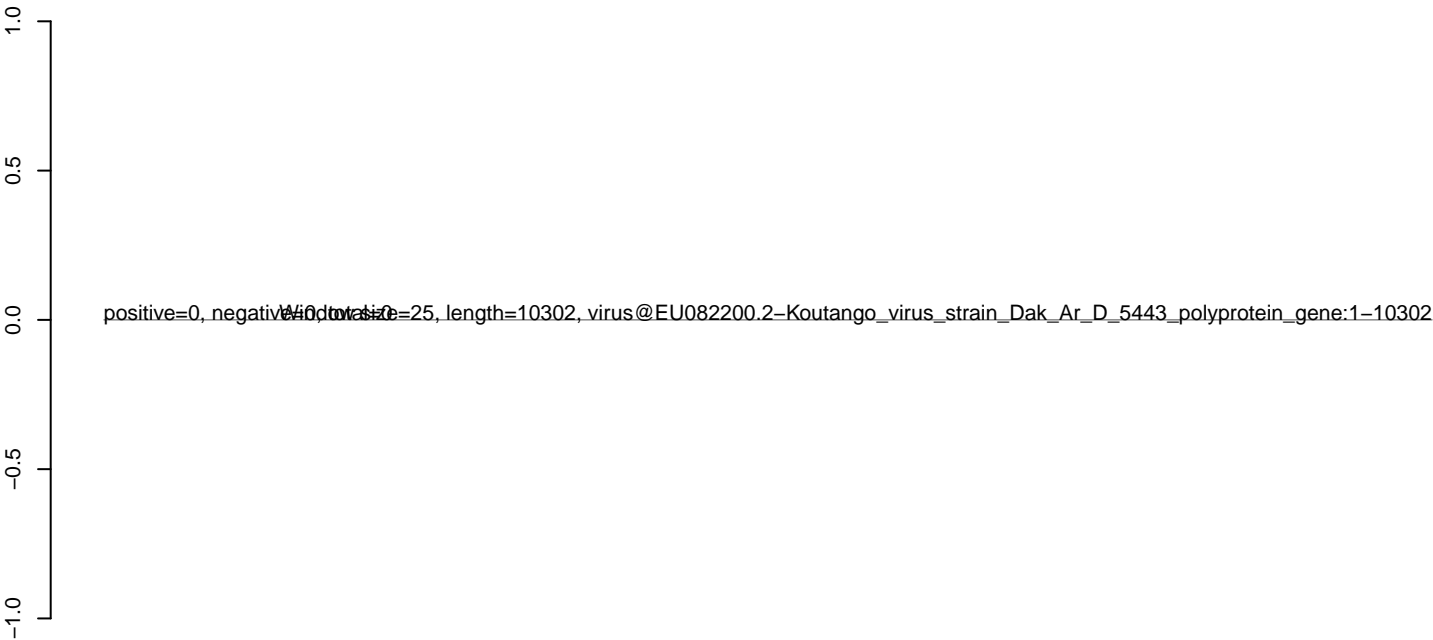
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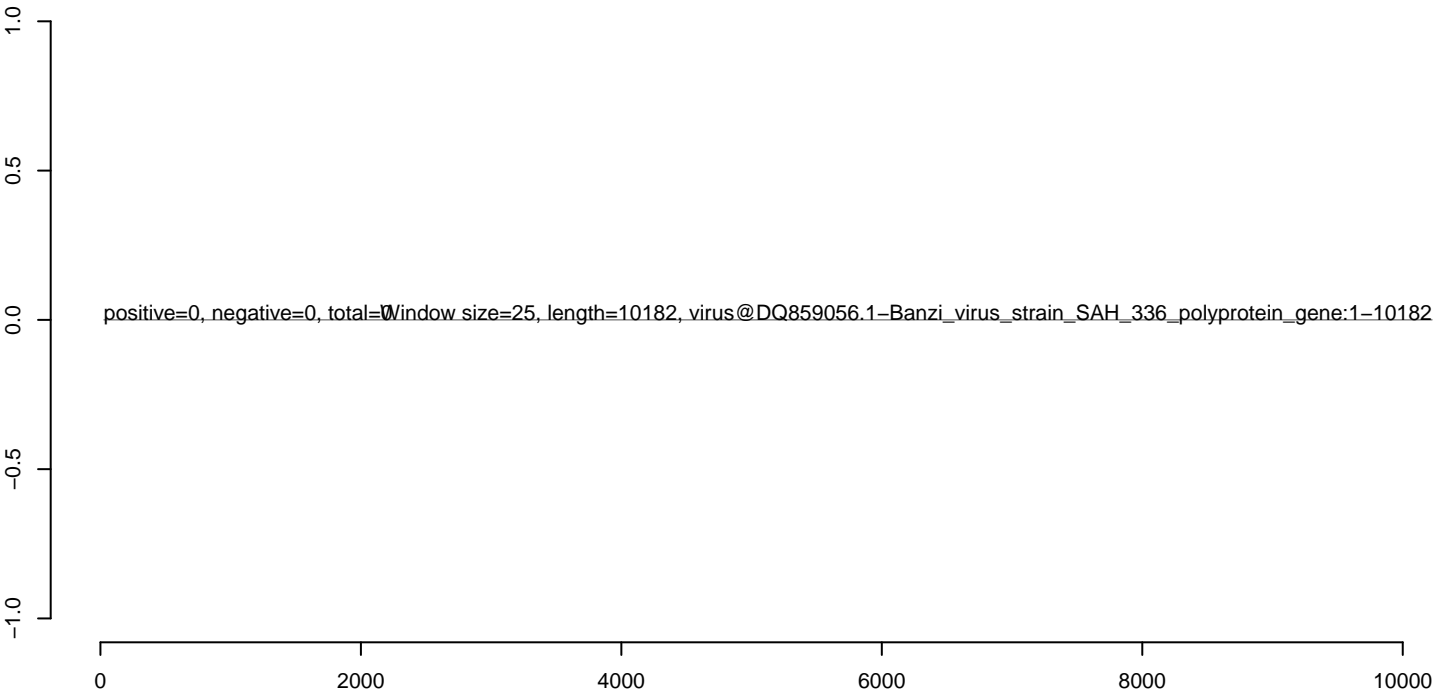
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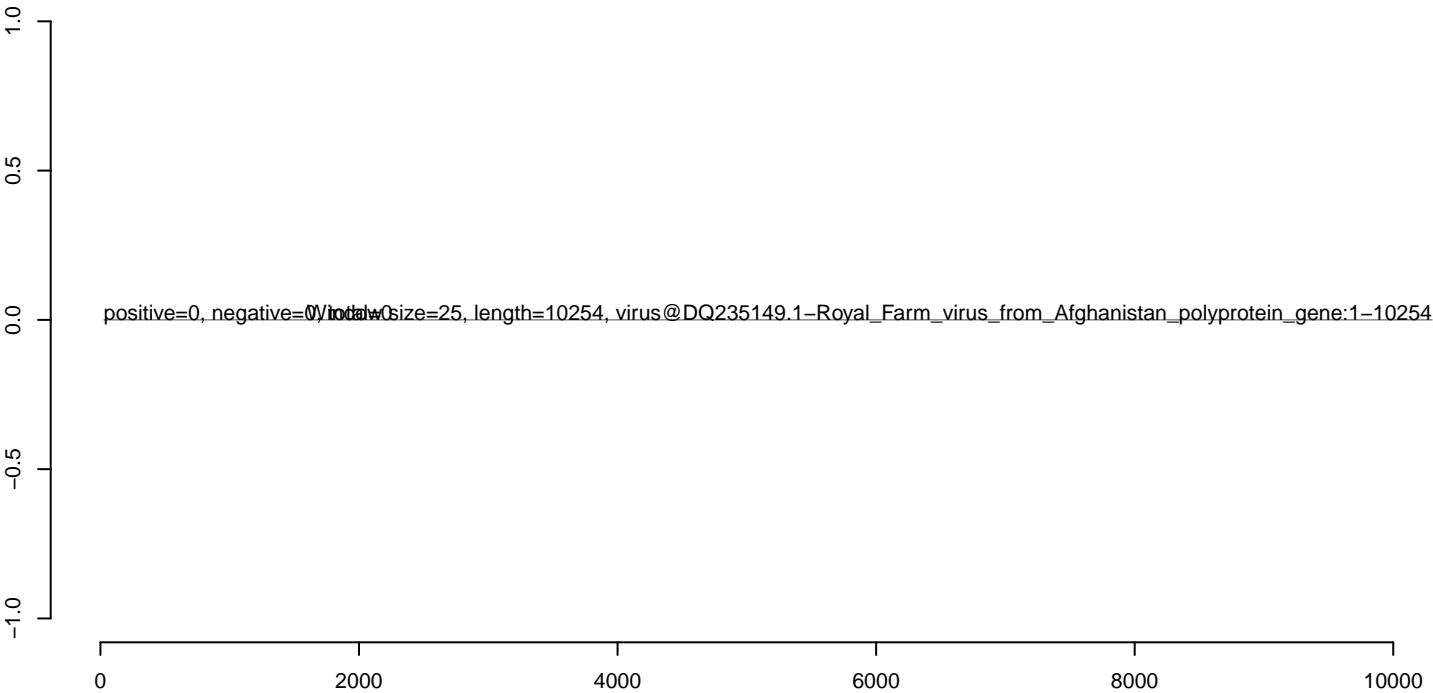
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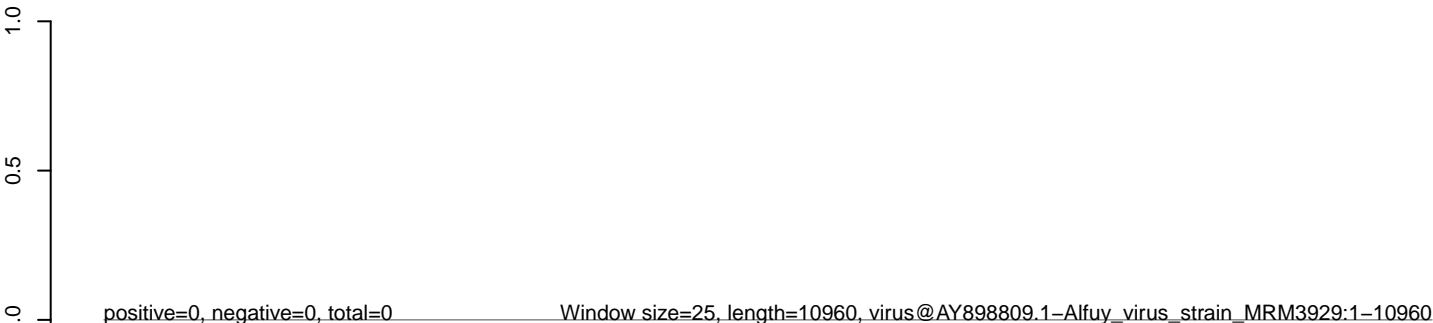
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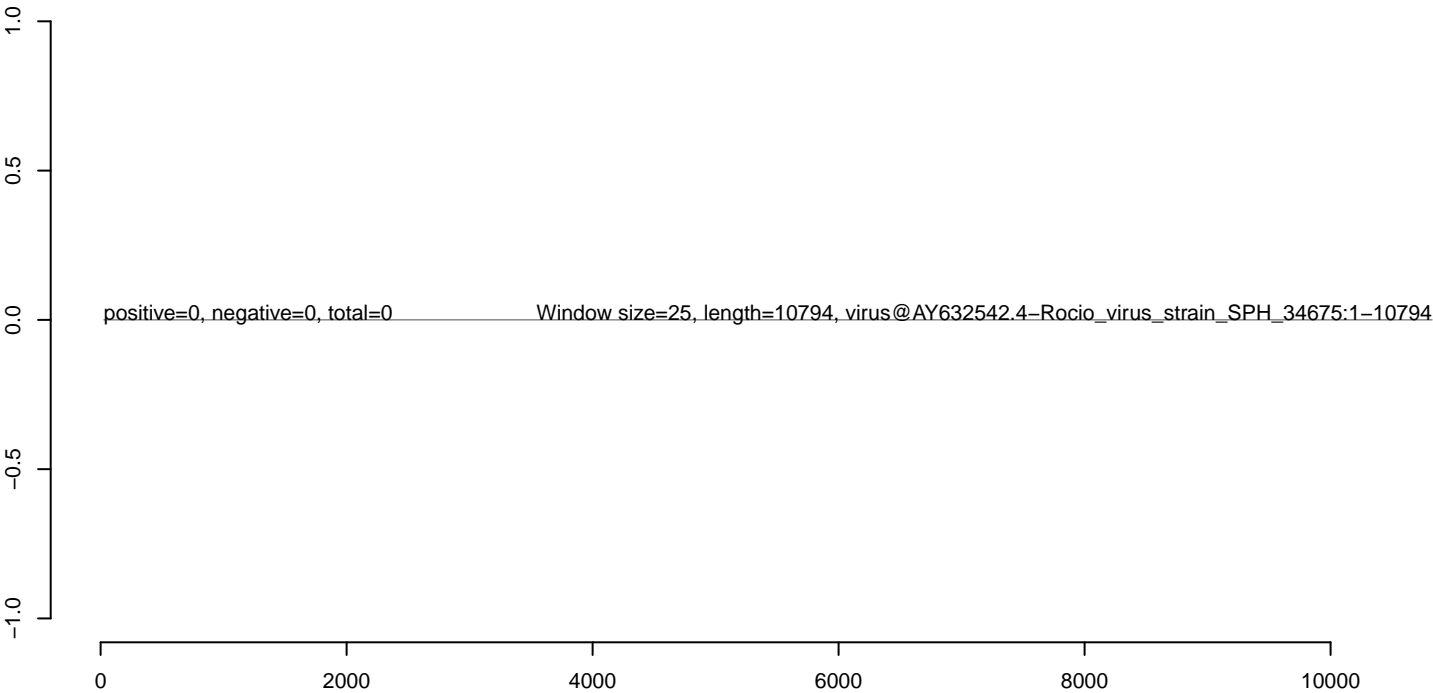
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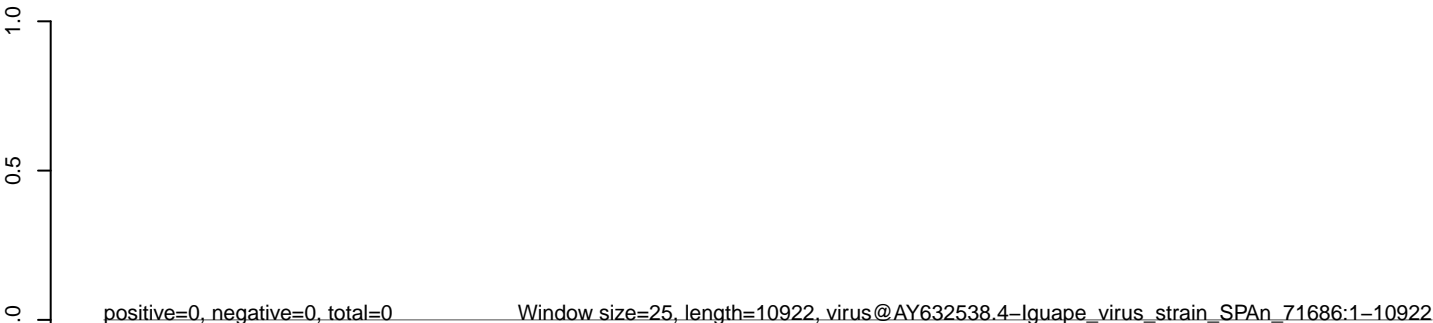
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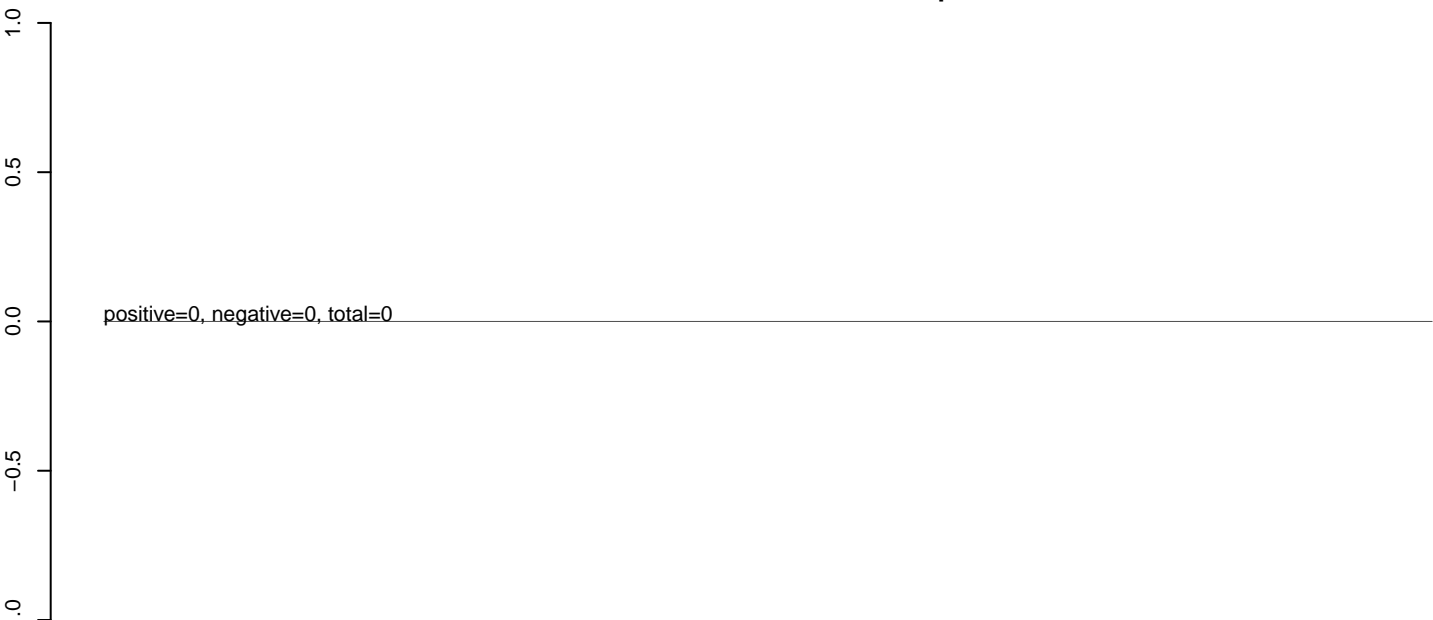
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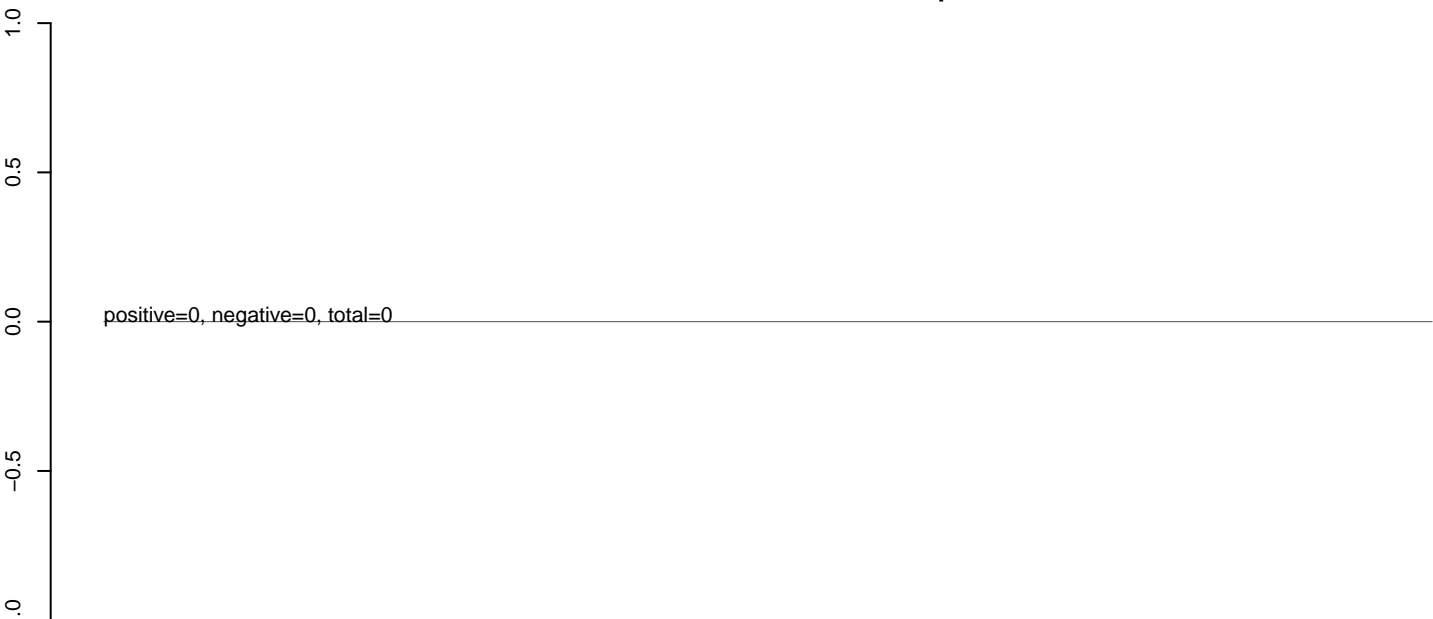
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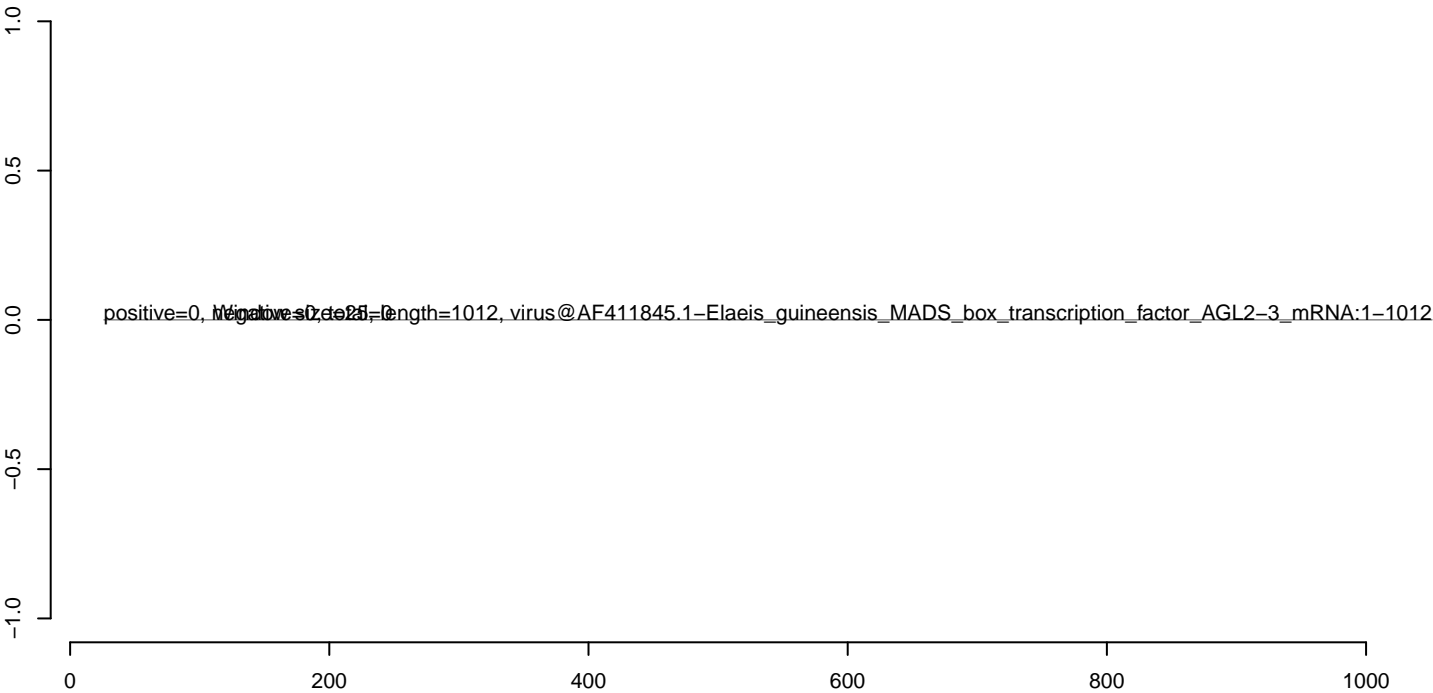
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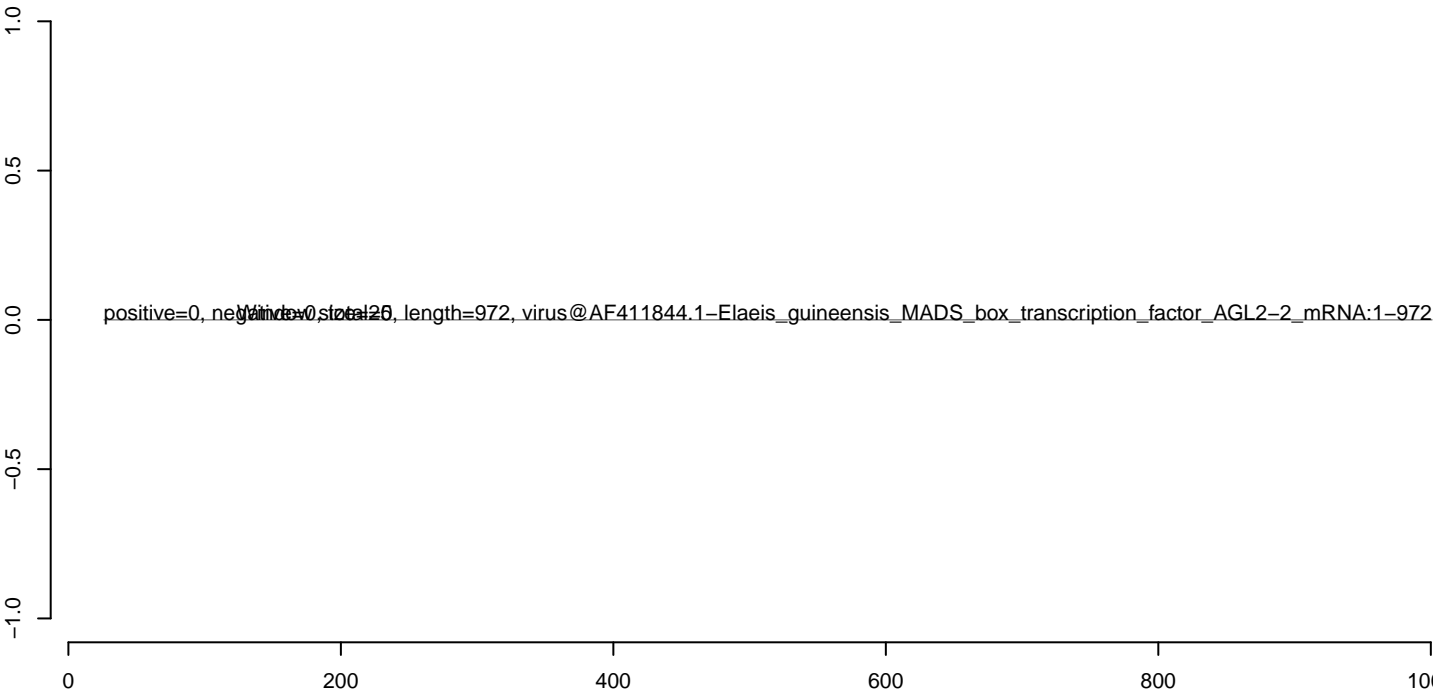
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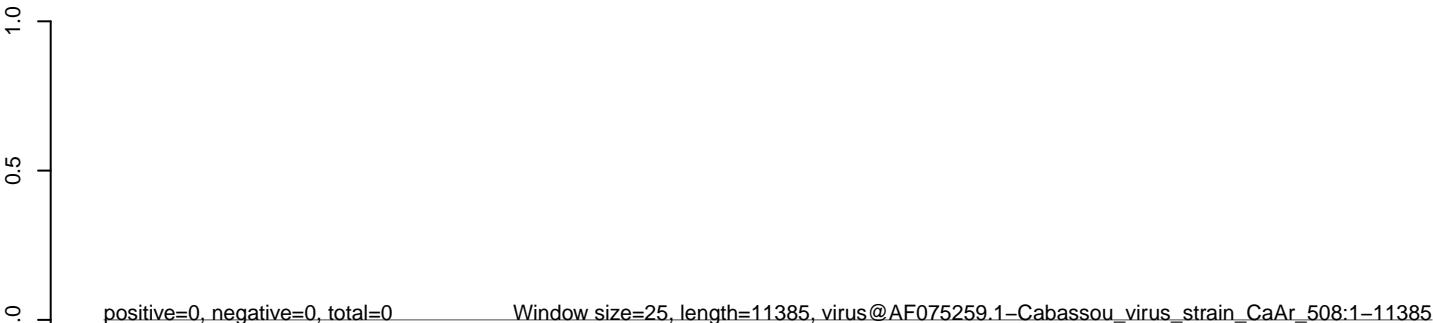
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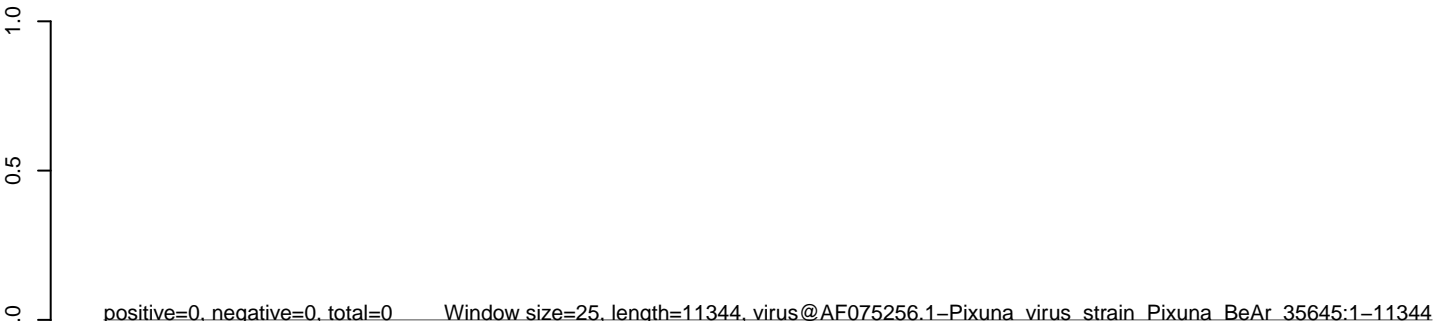
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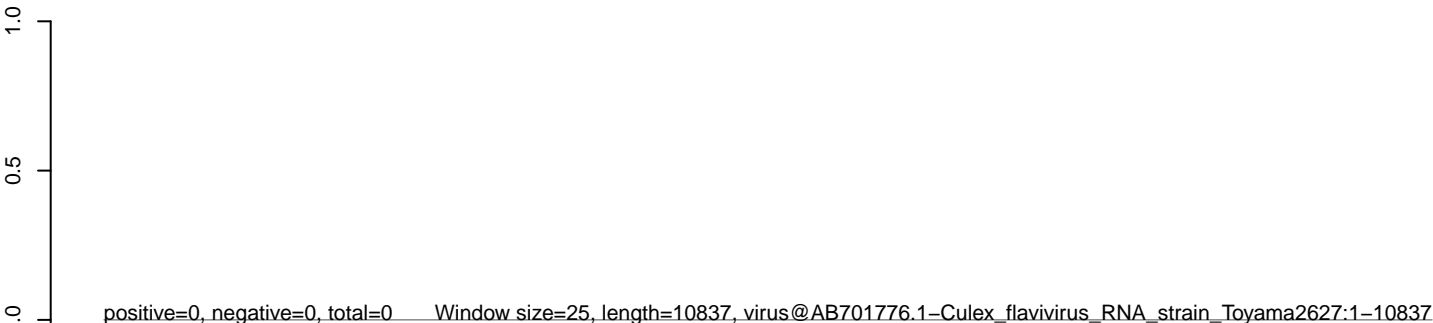
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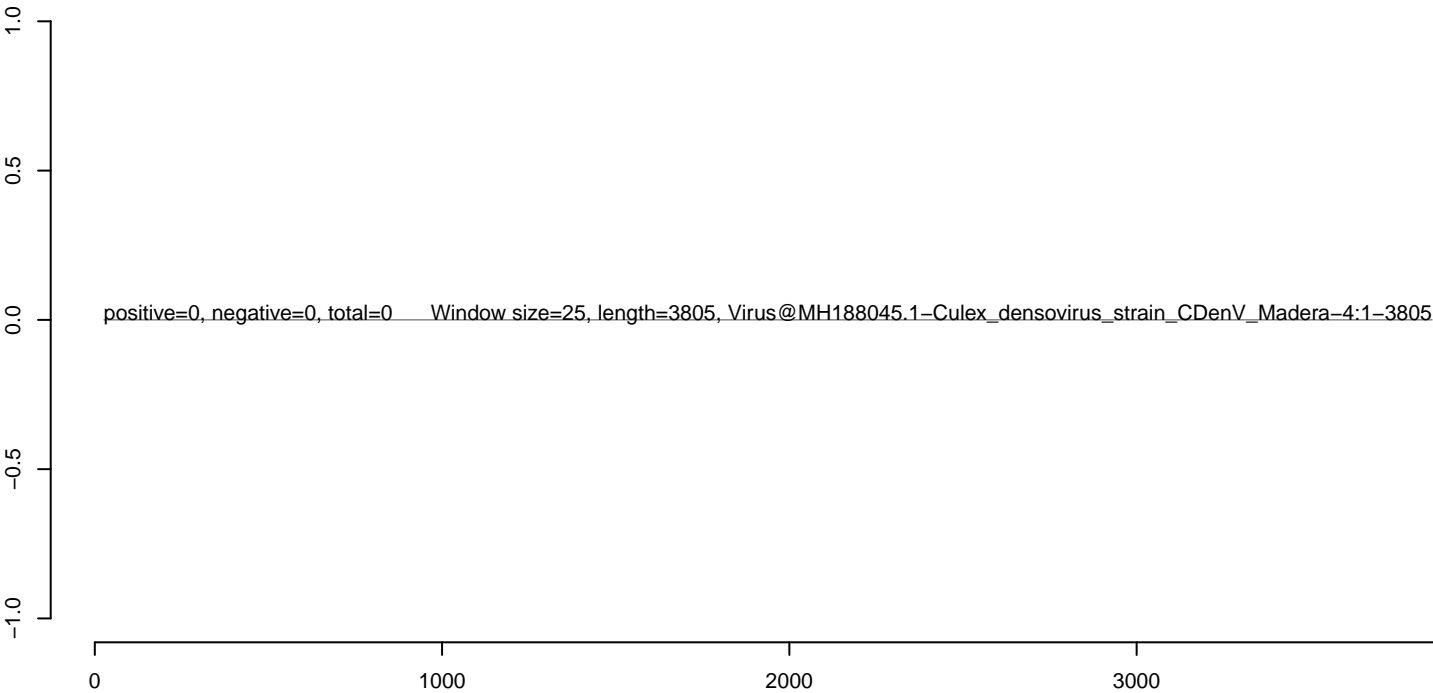
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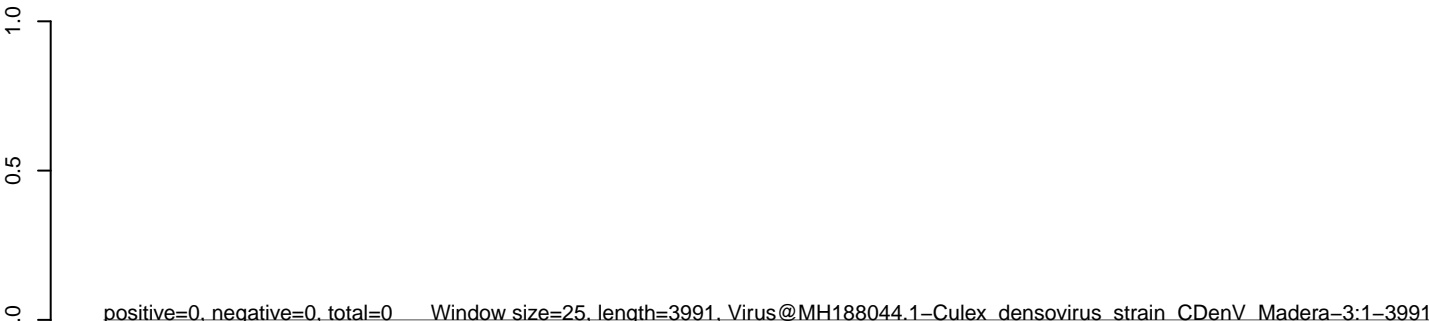
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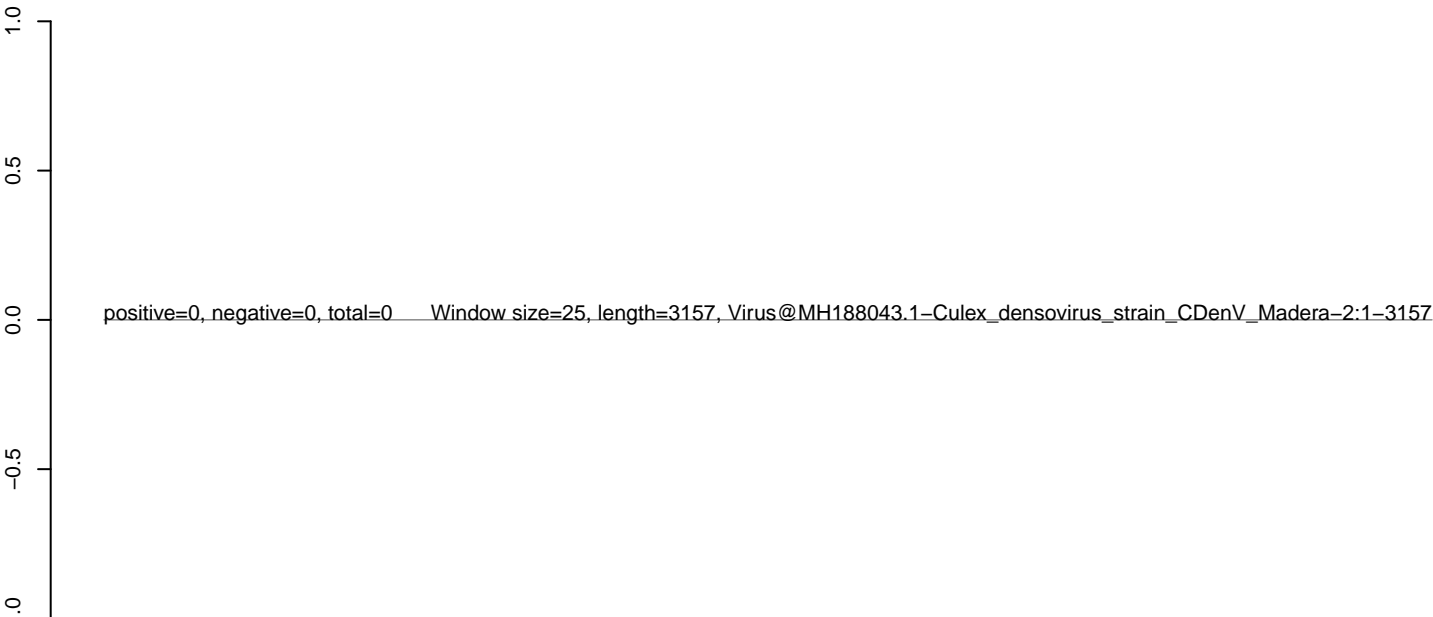
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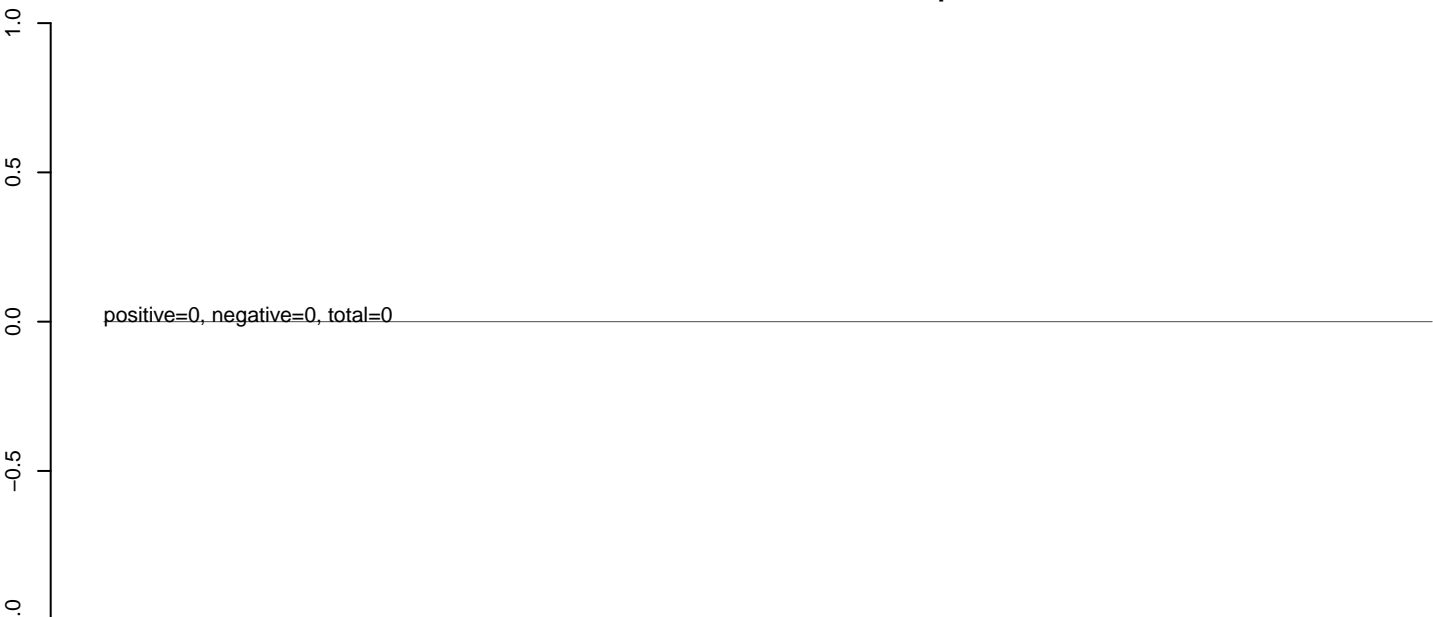
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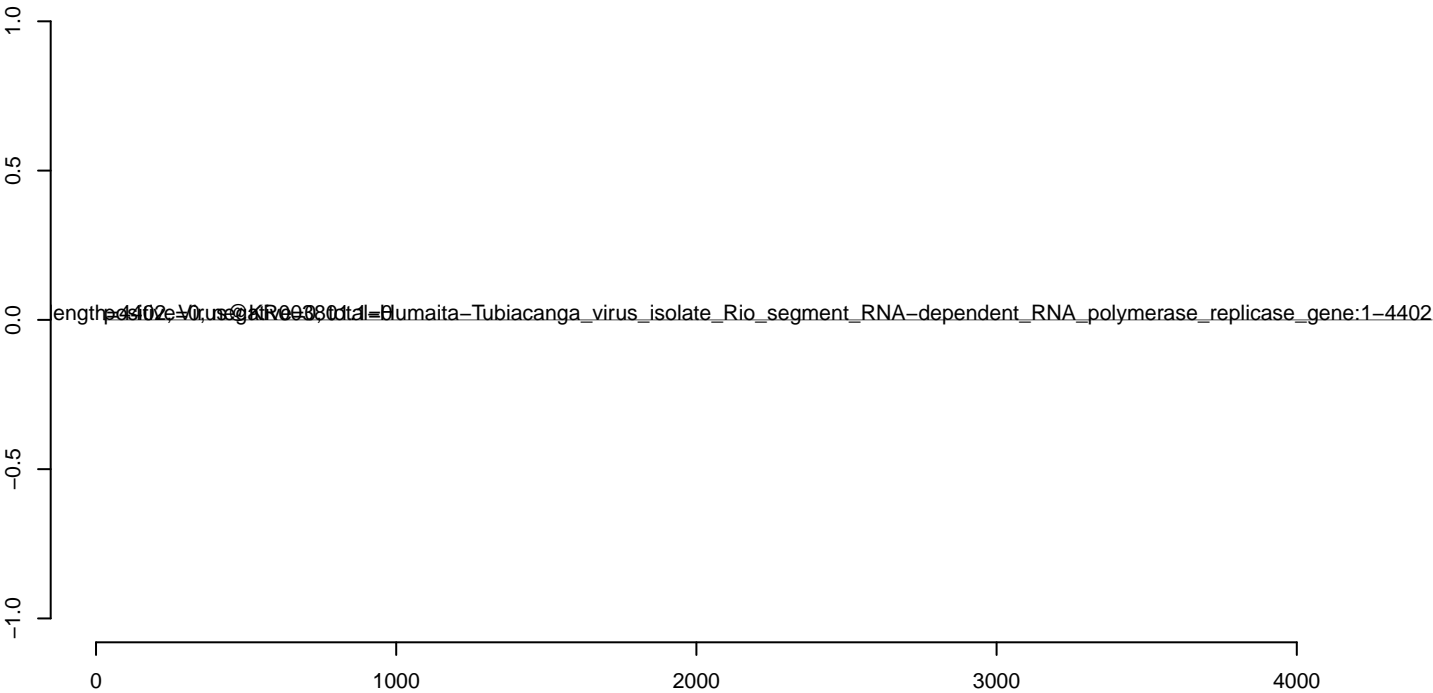
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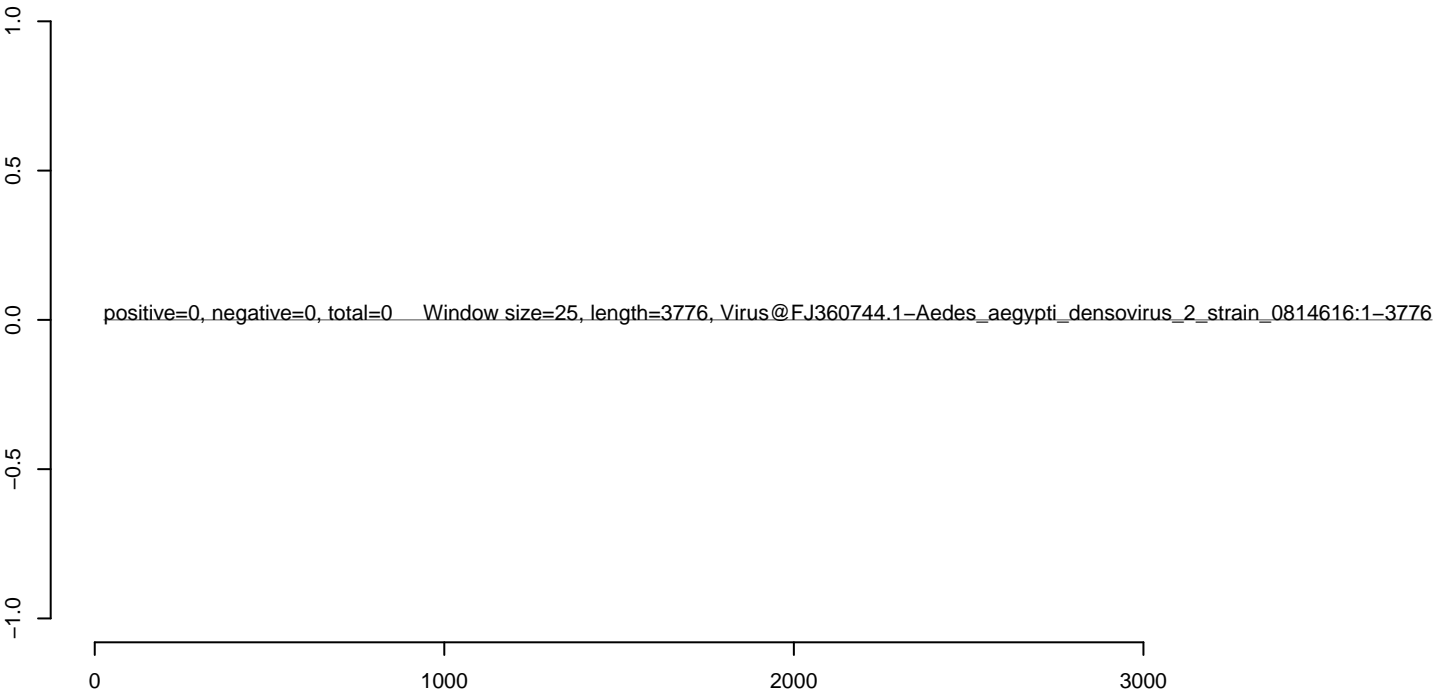
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