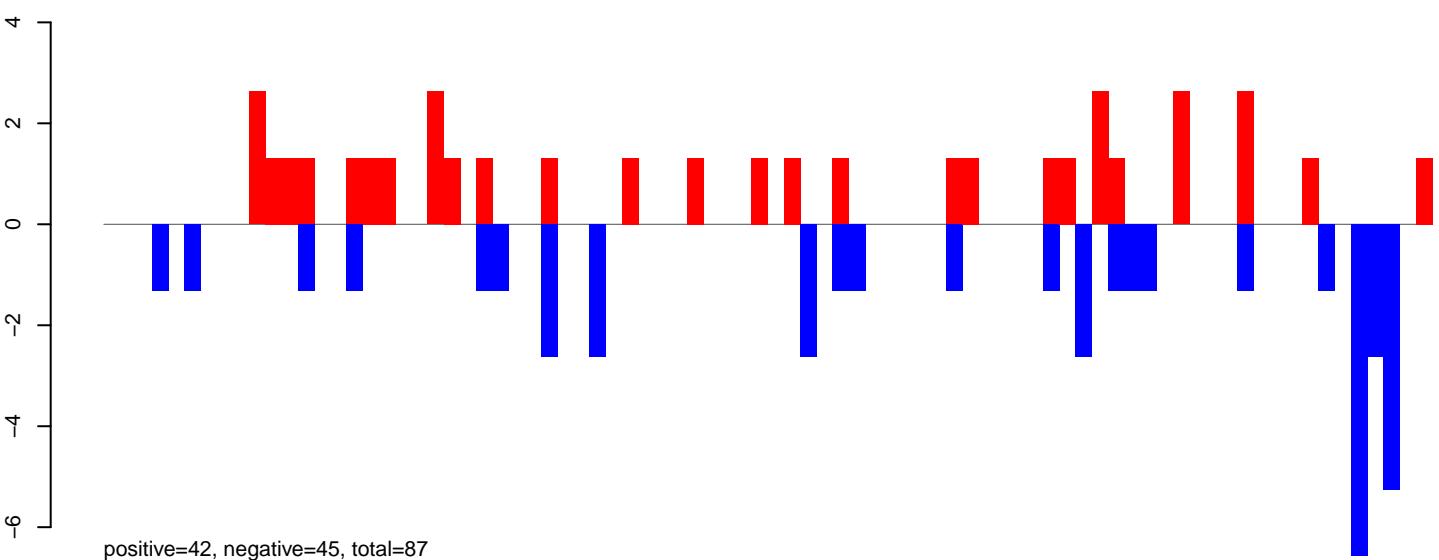
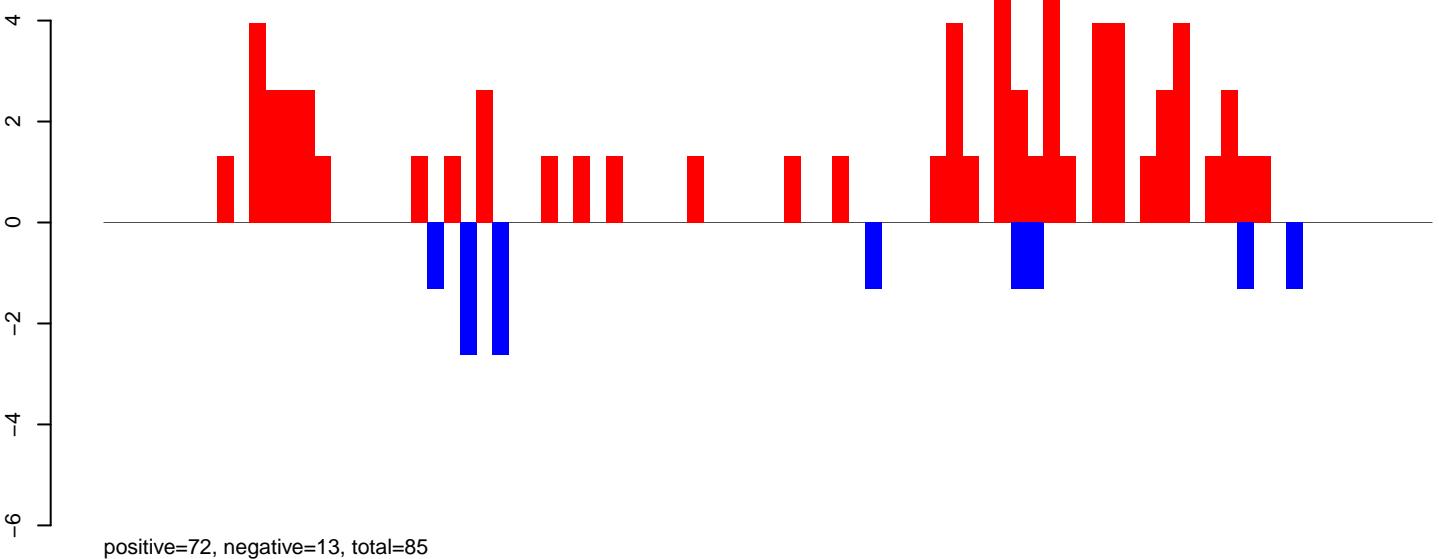


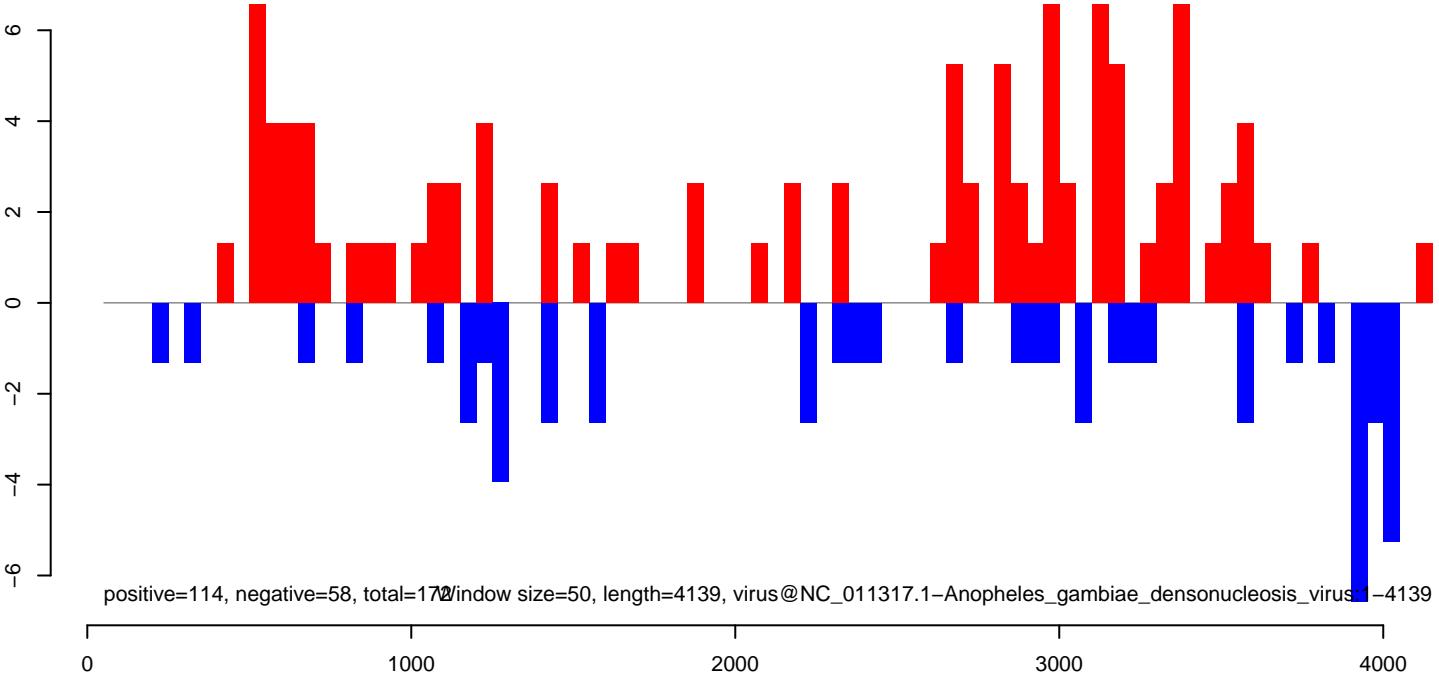
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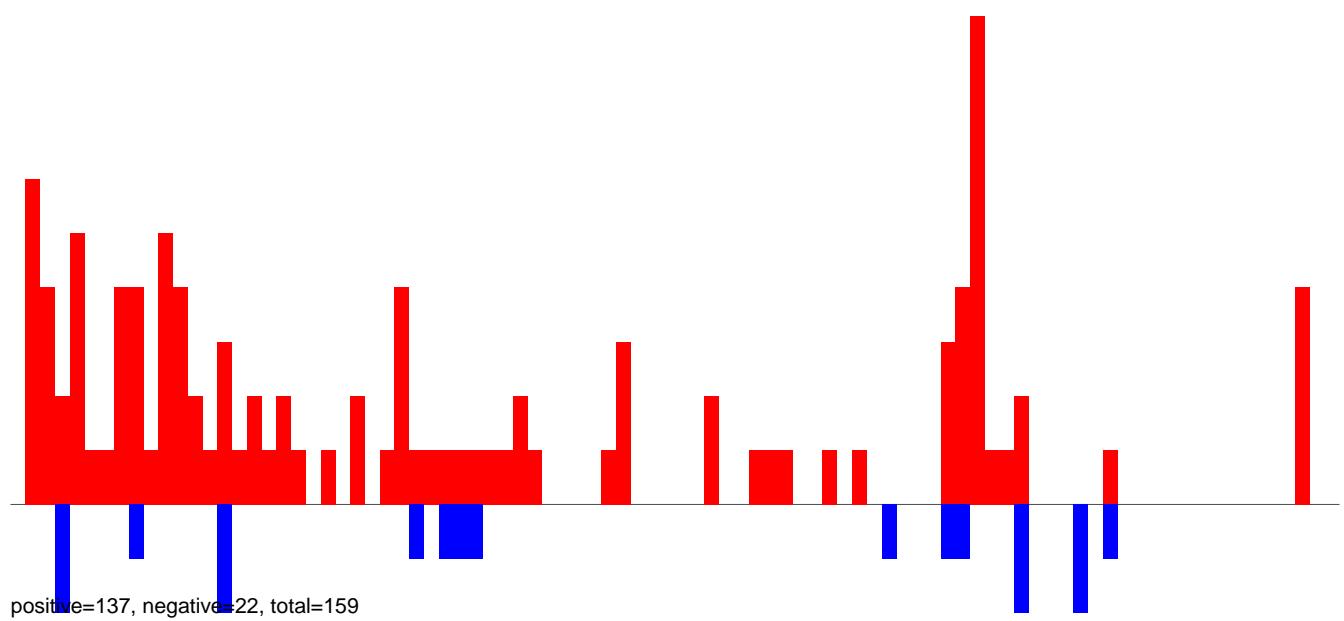
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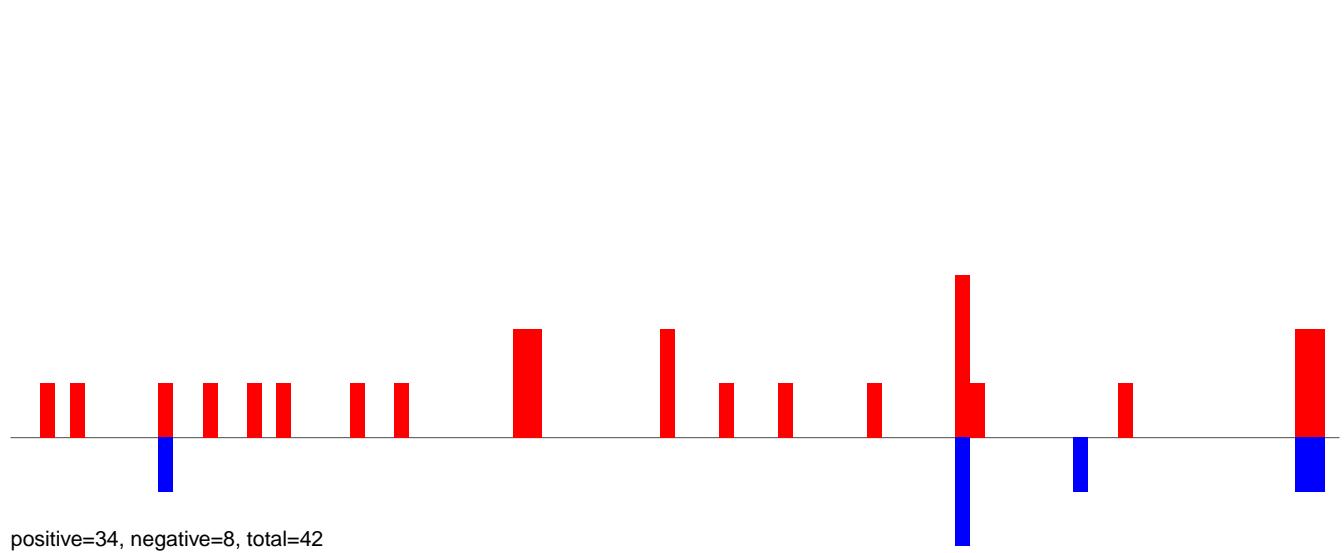
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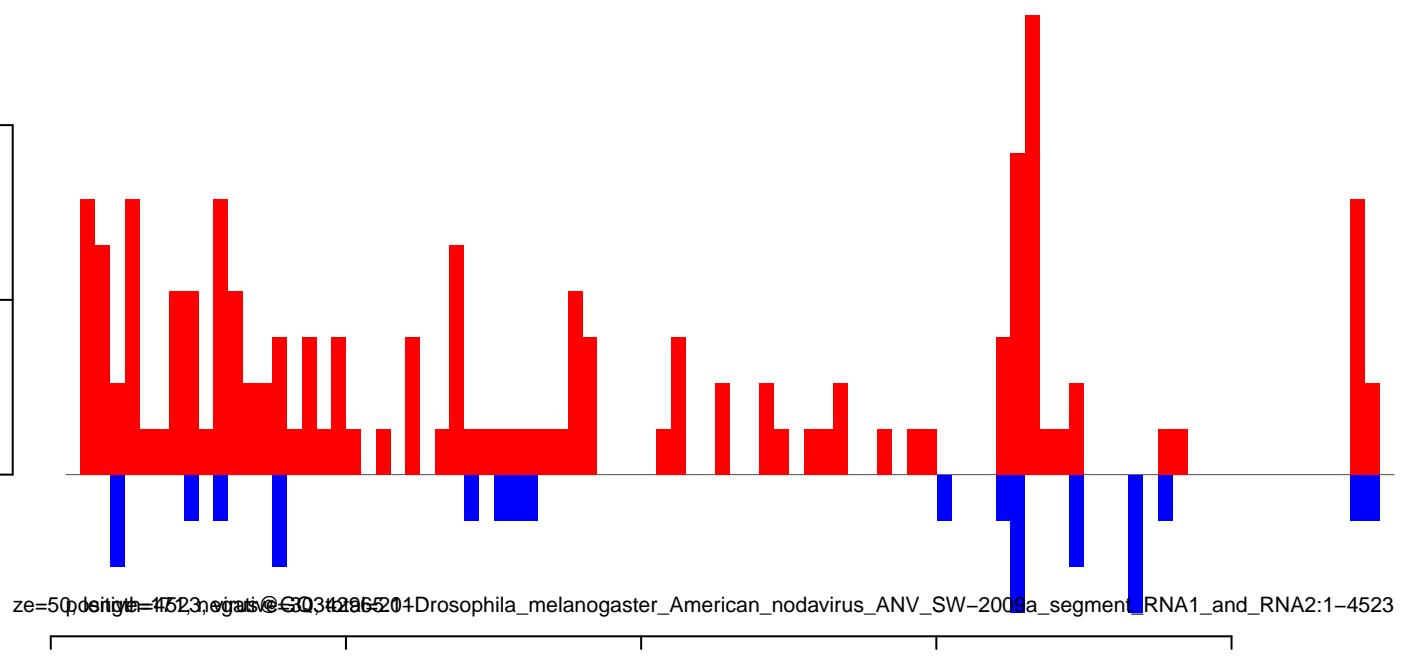
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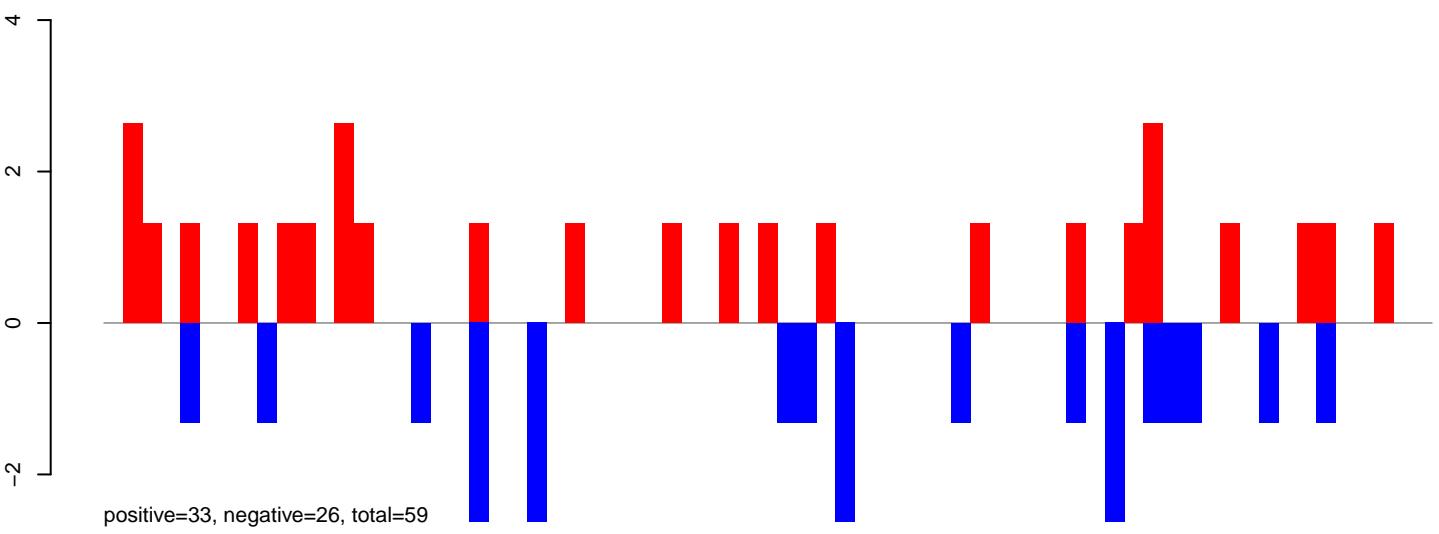
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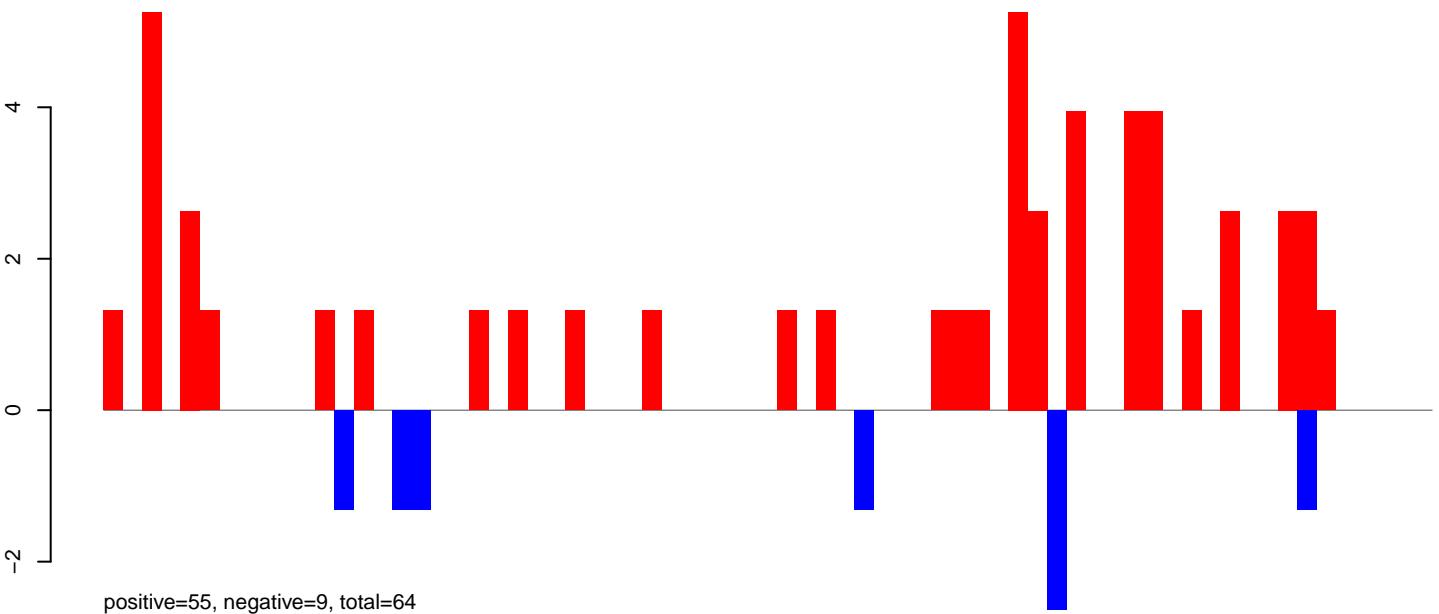
AnGam\_Fcanc\_TN.rep



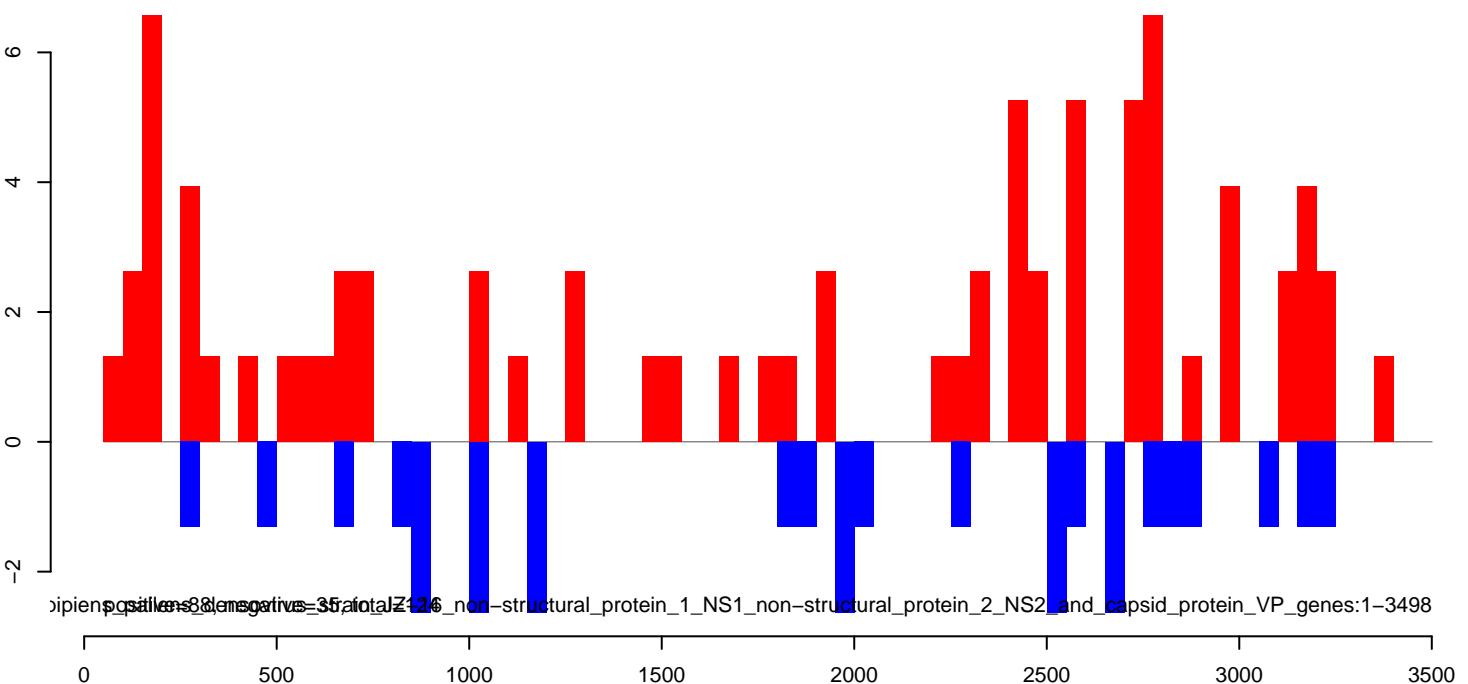
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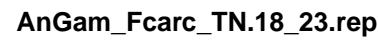


### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep





positive=11, negative=20, total=30

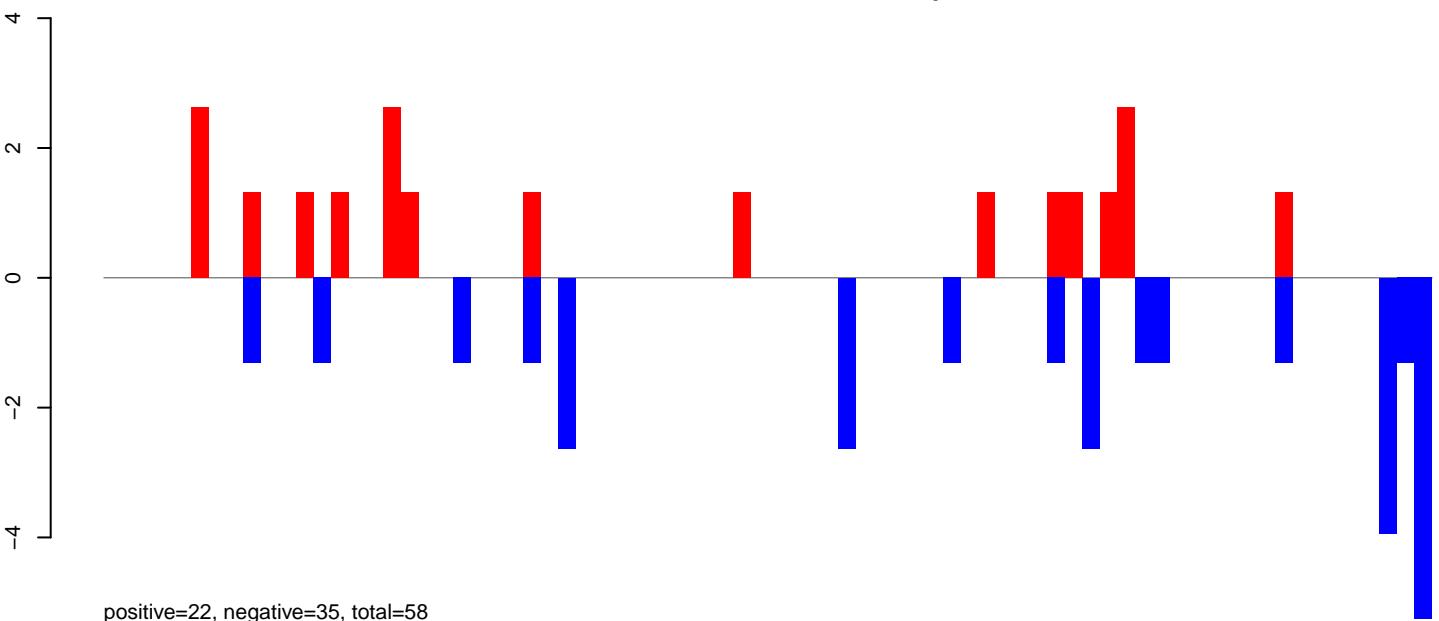


positive=21, negative=88, total=109

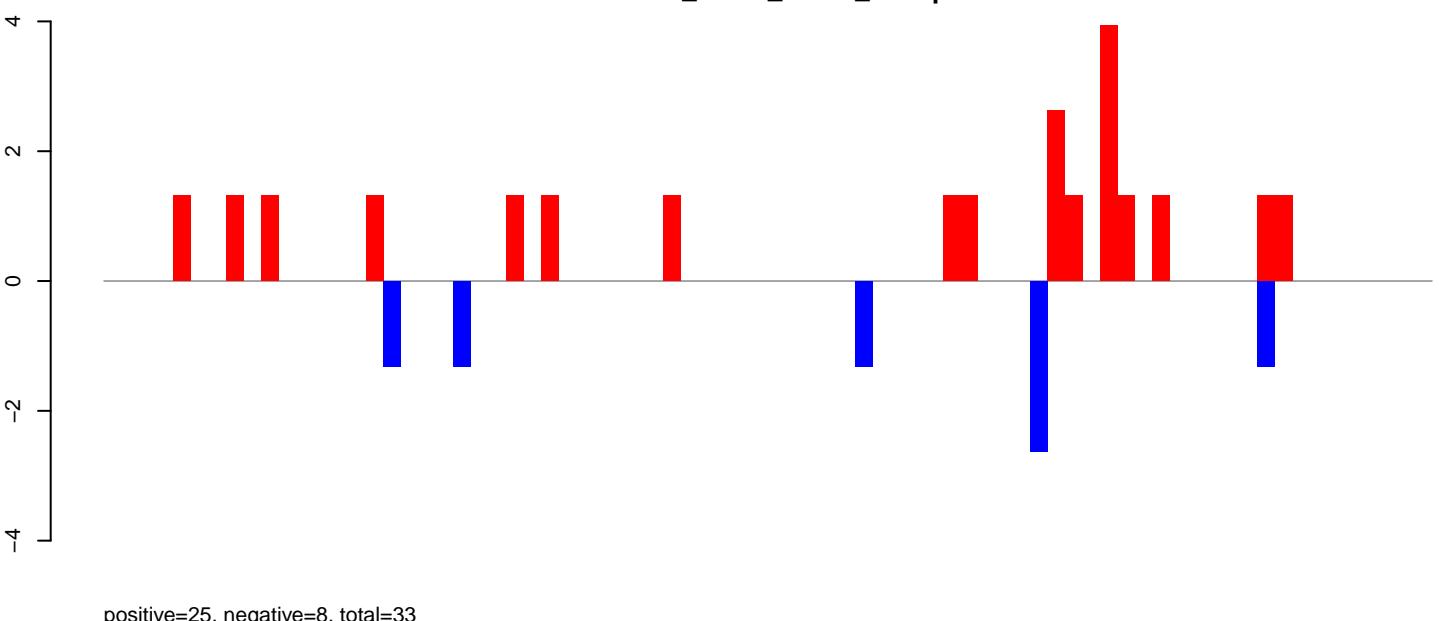


2099, piso12nucleo2036260v=P08stoG1at30n-like\_virus\_isolate\_Rio\_segment\_S\_plus\_segment\_M\_plus\_segment\_L\_Nucleocapsid\_gene:1-12099

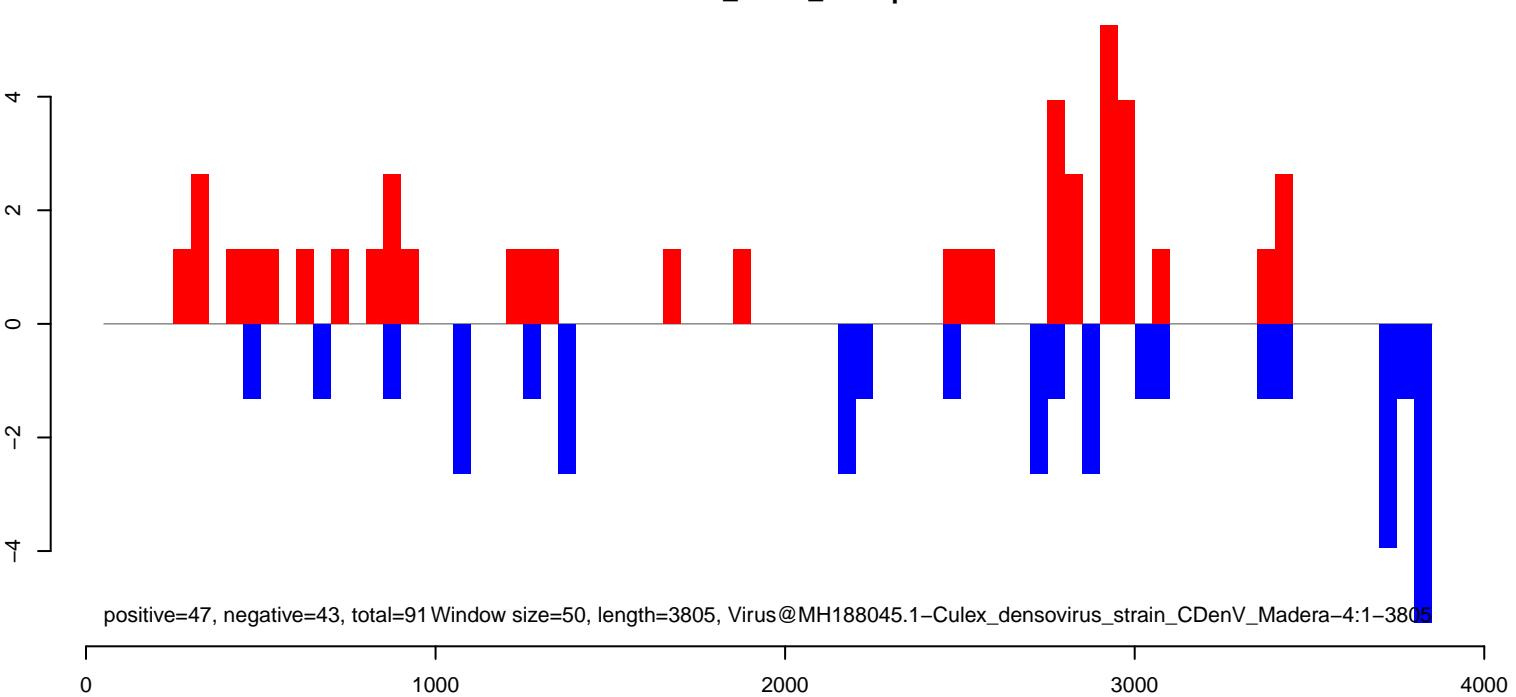
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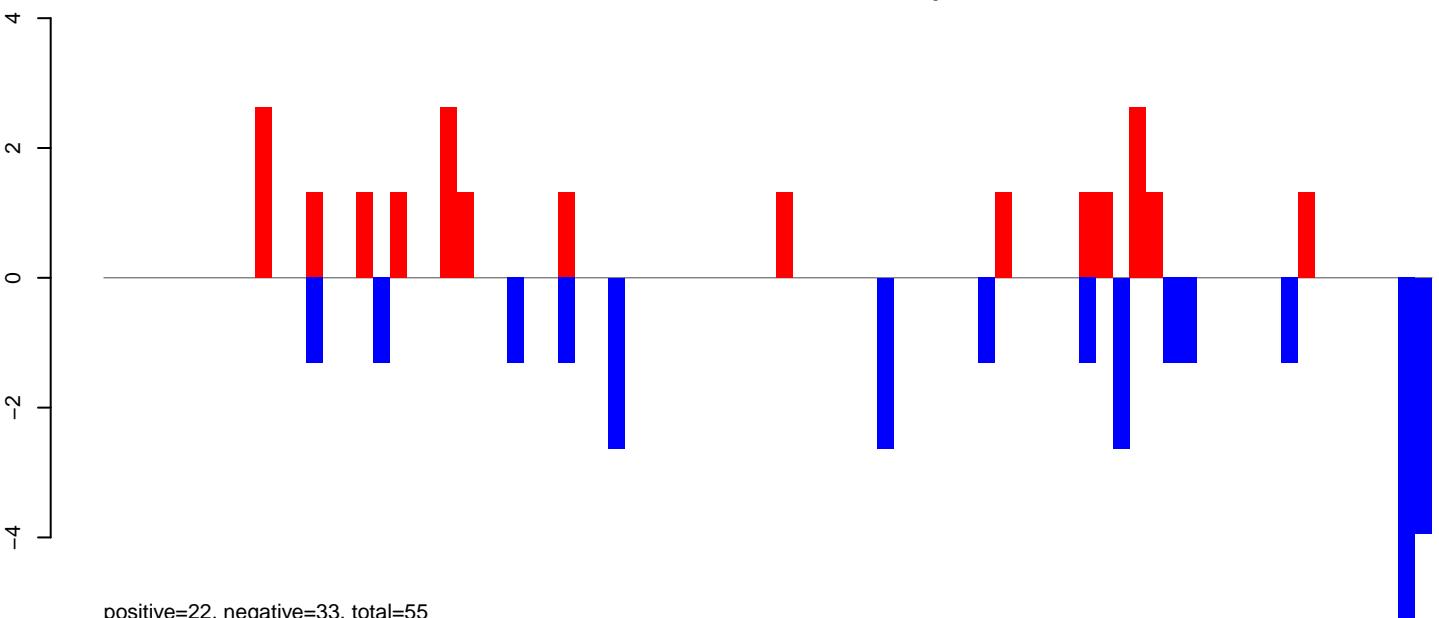
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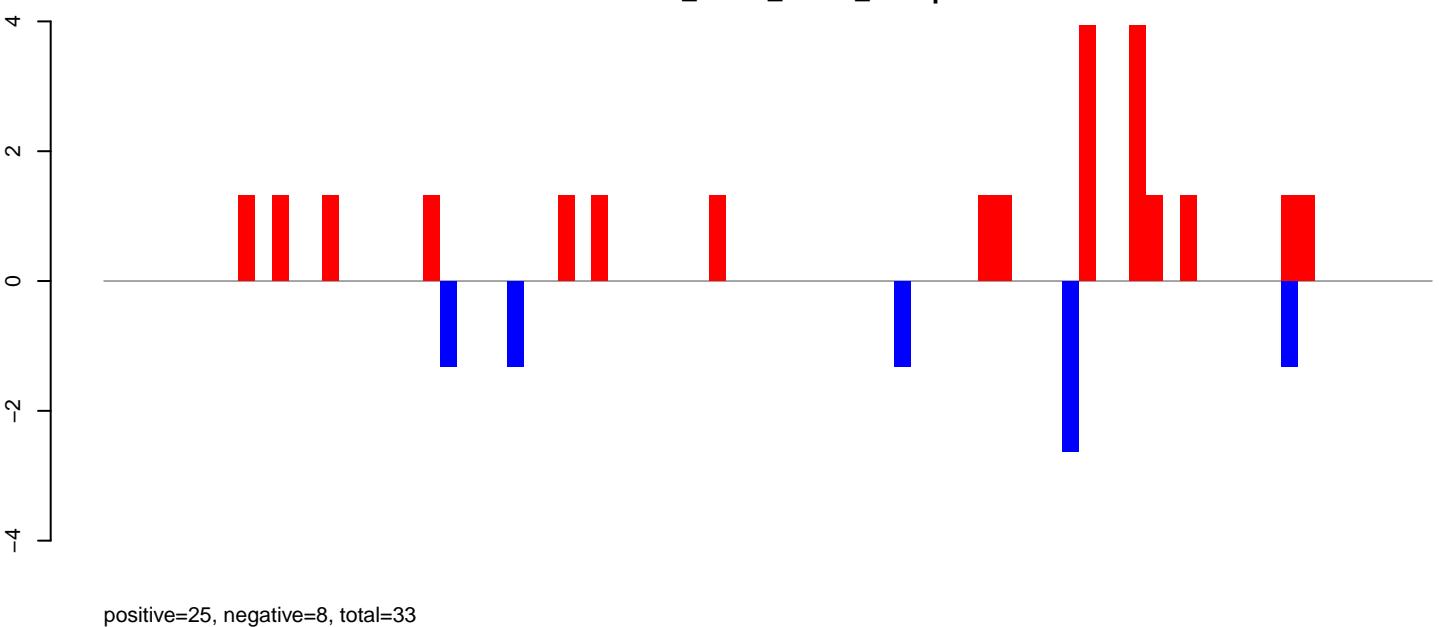
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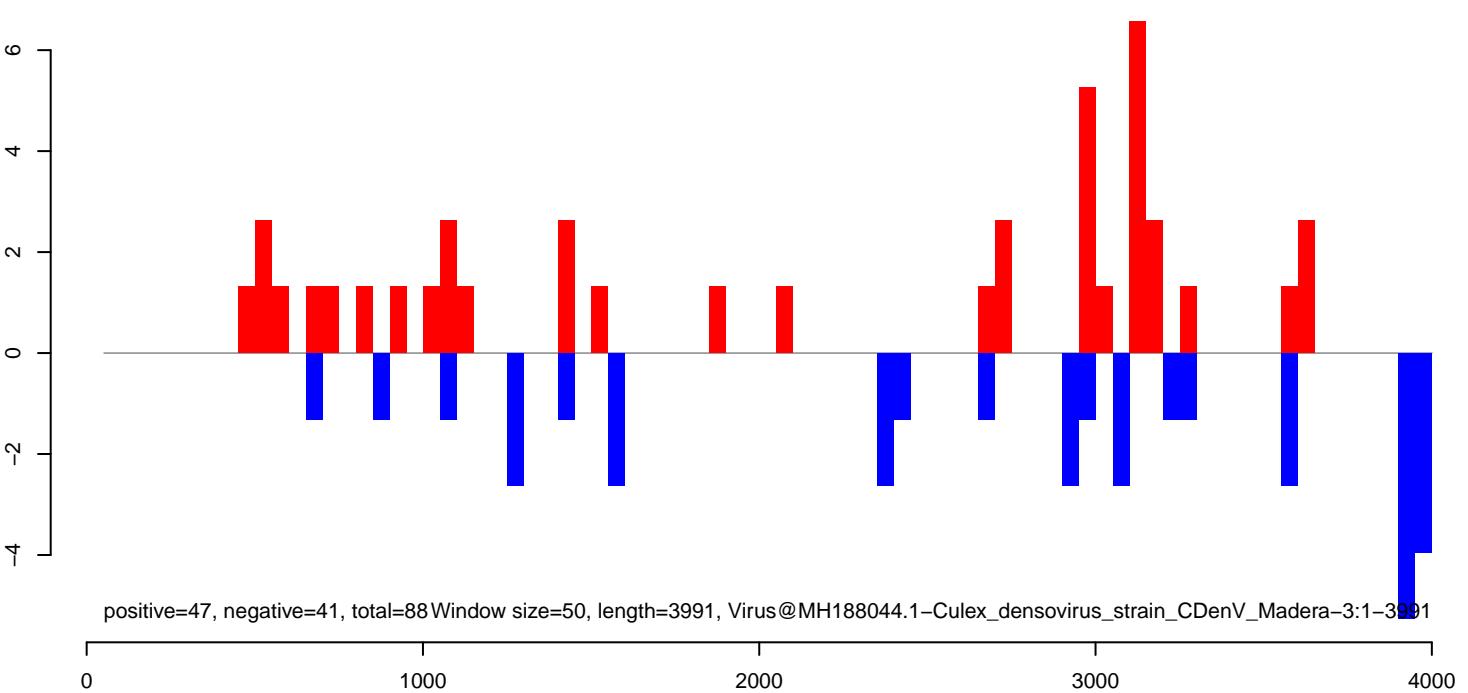
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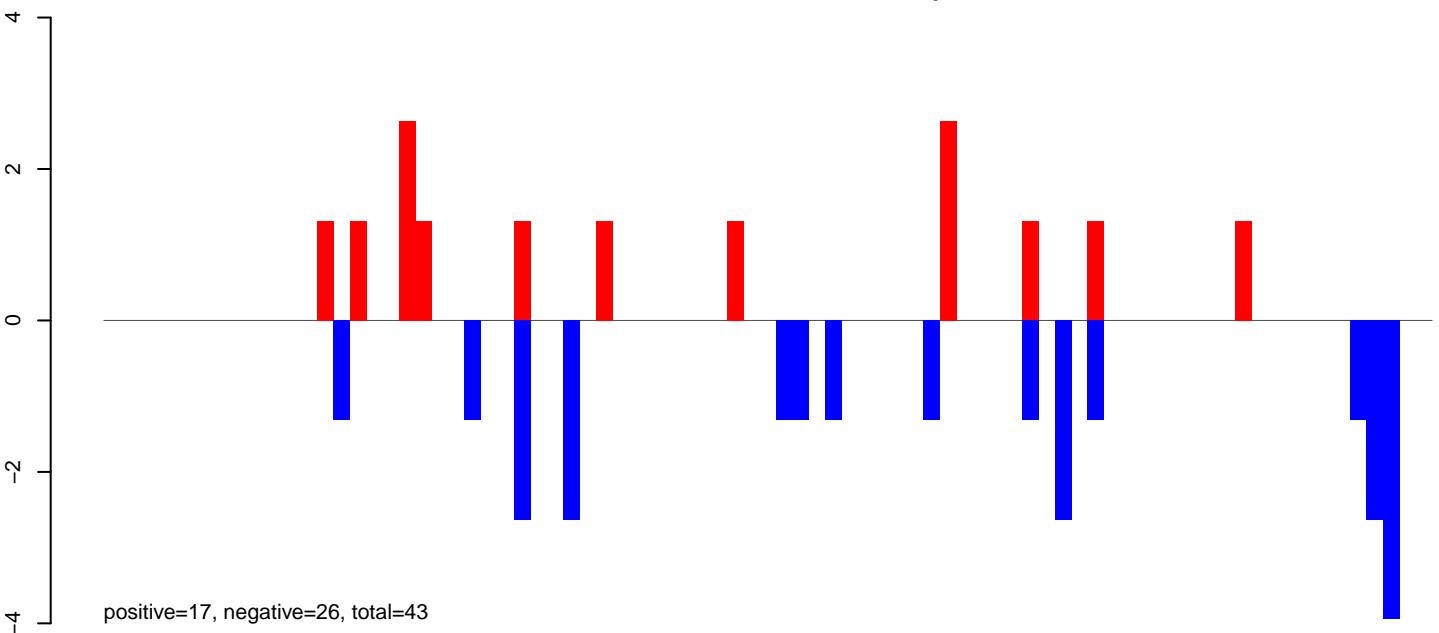
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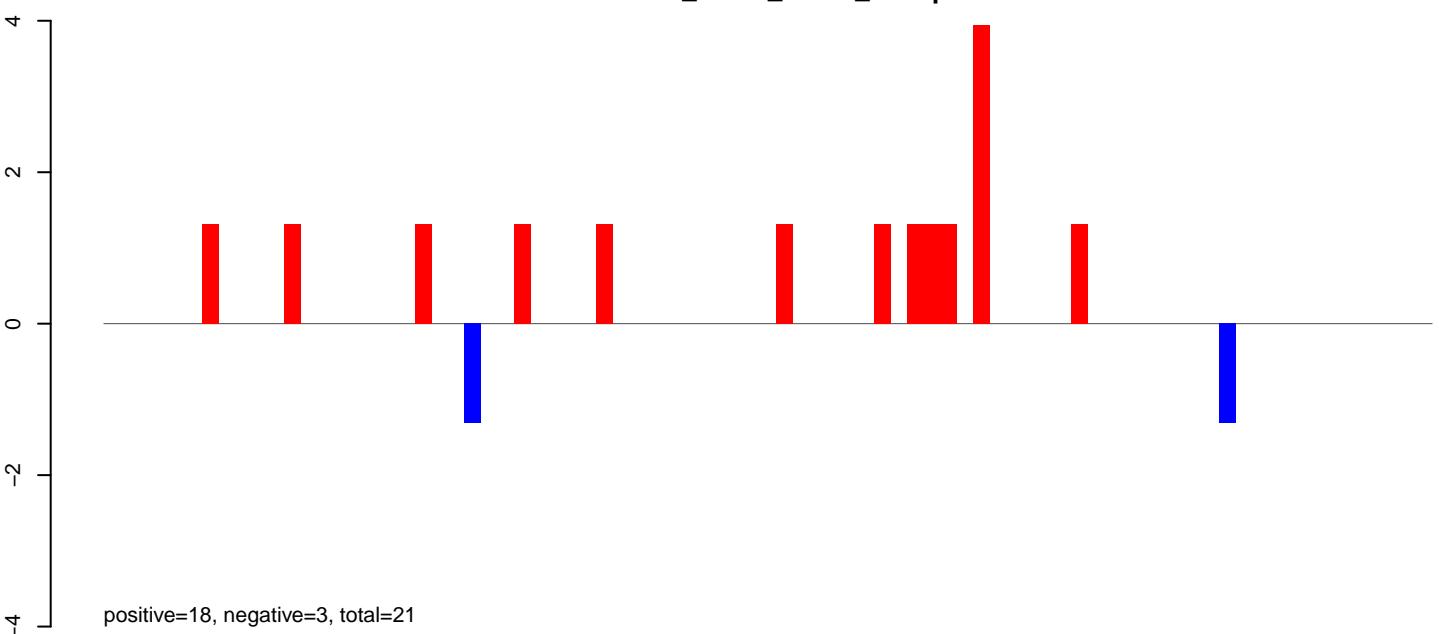
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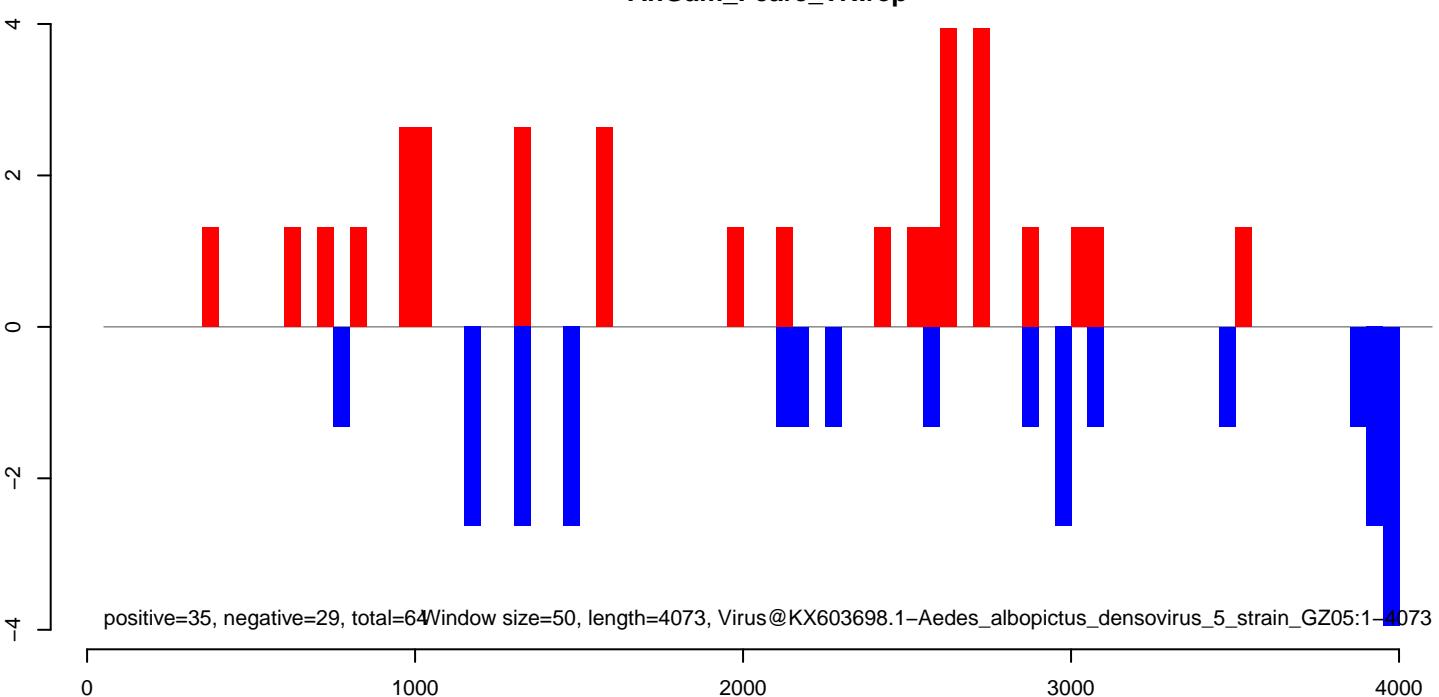
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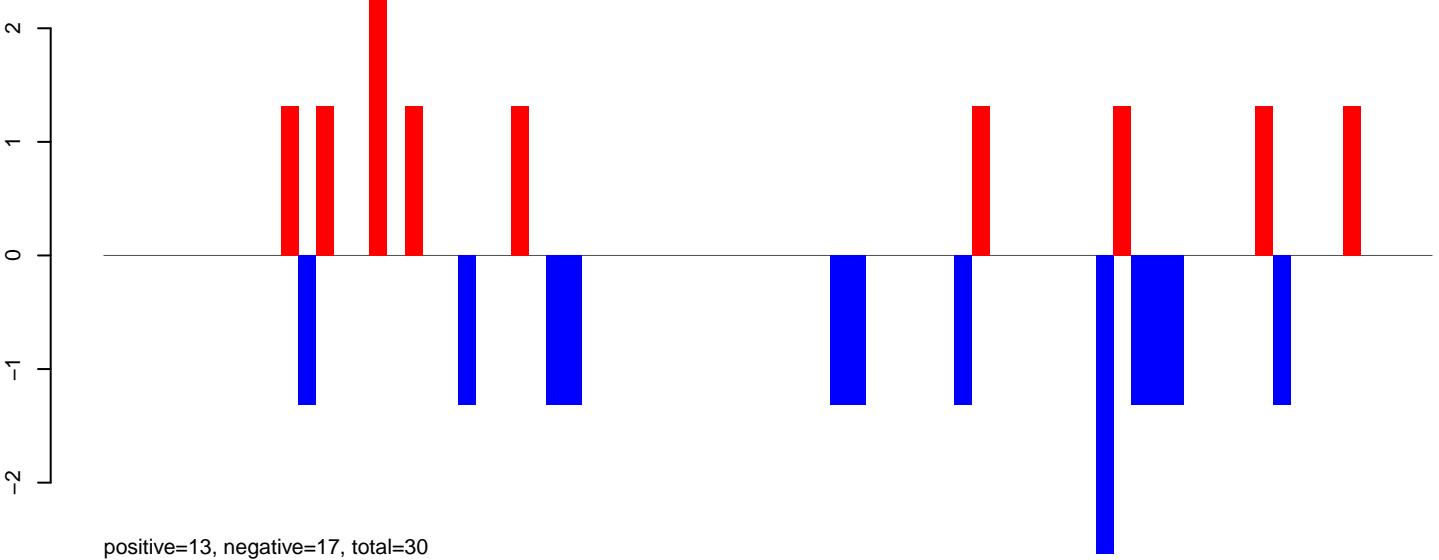
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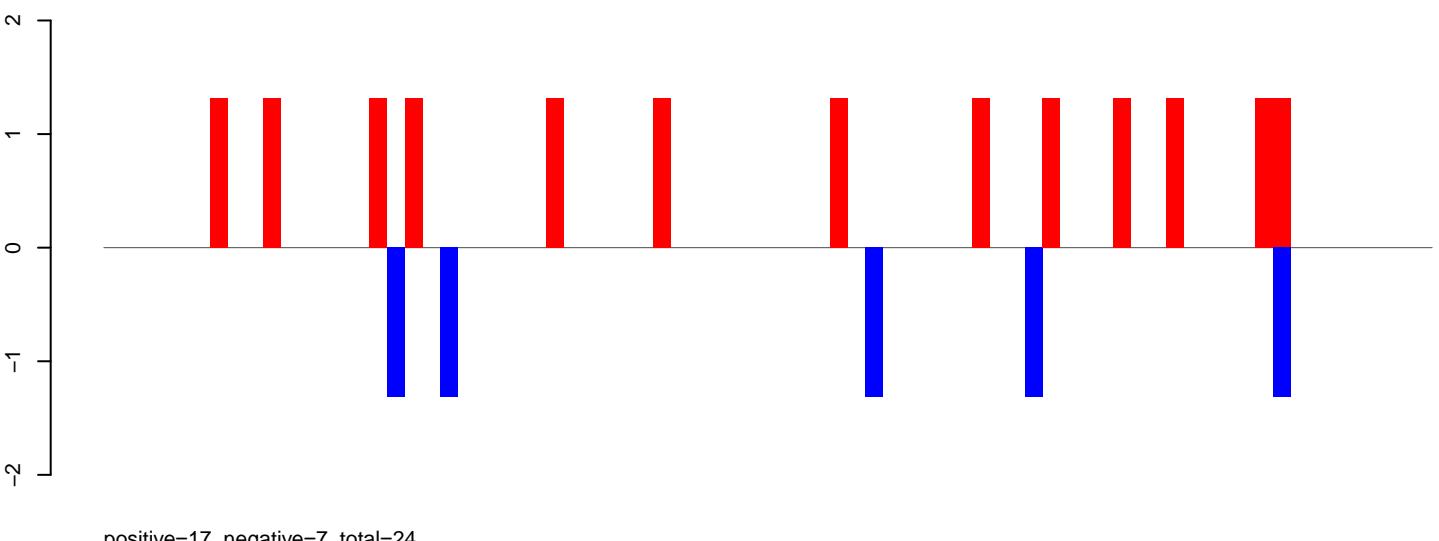
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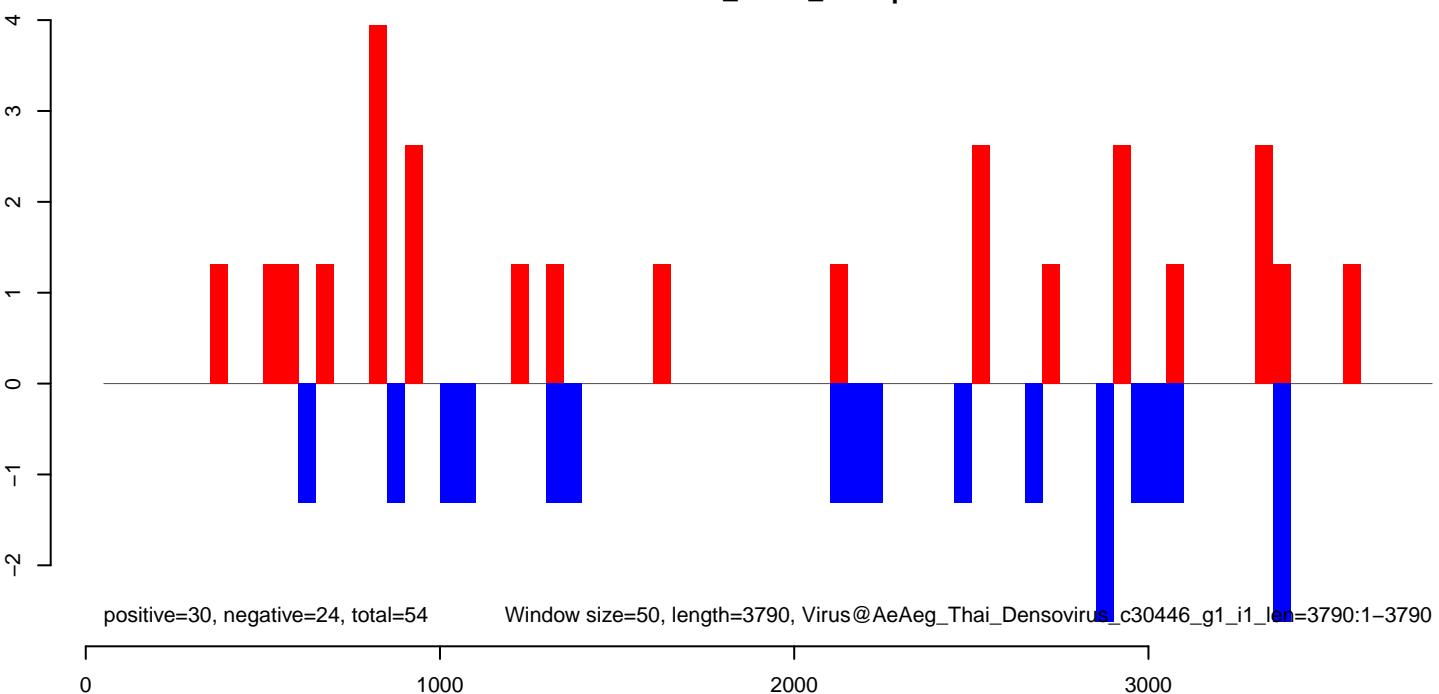
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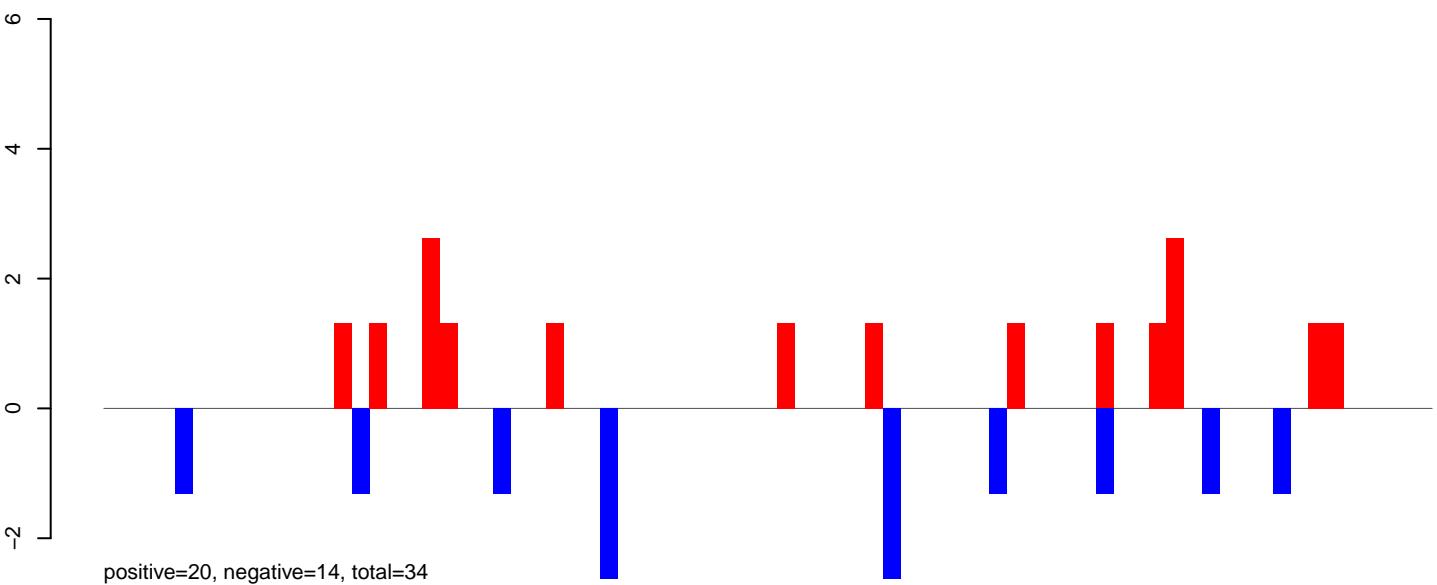
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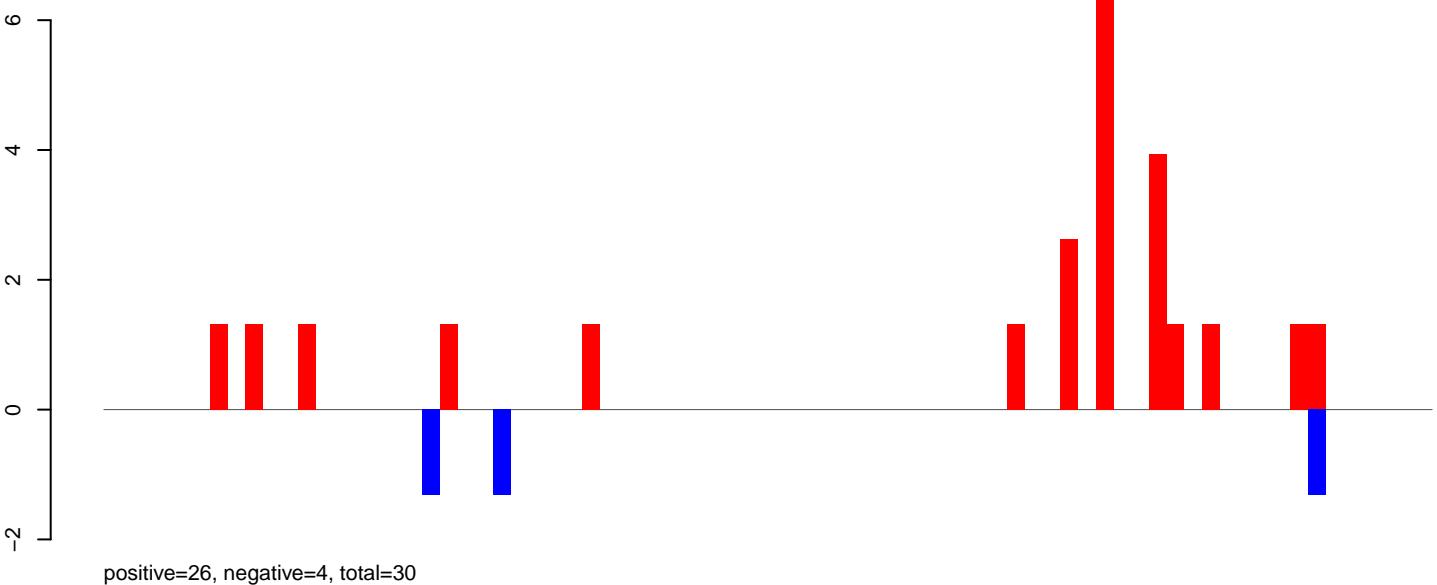
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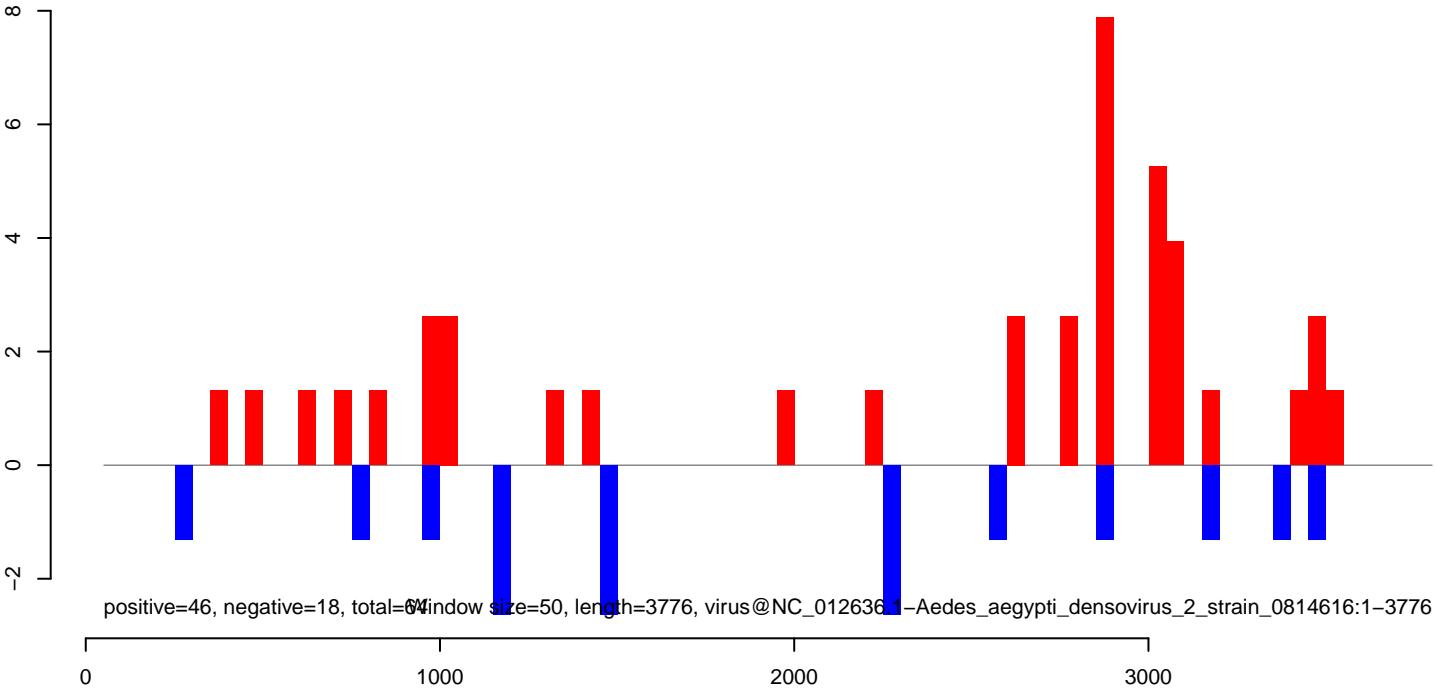
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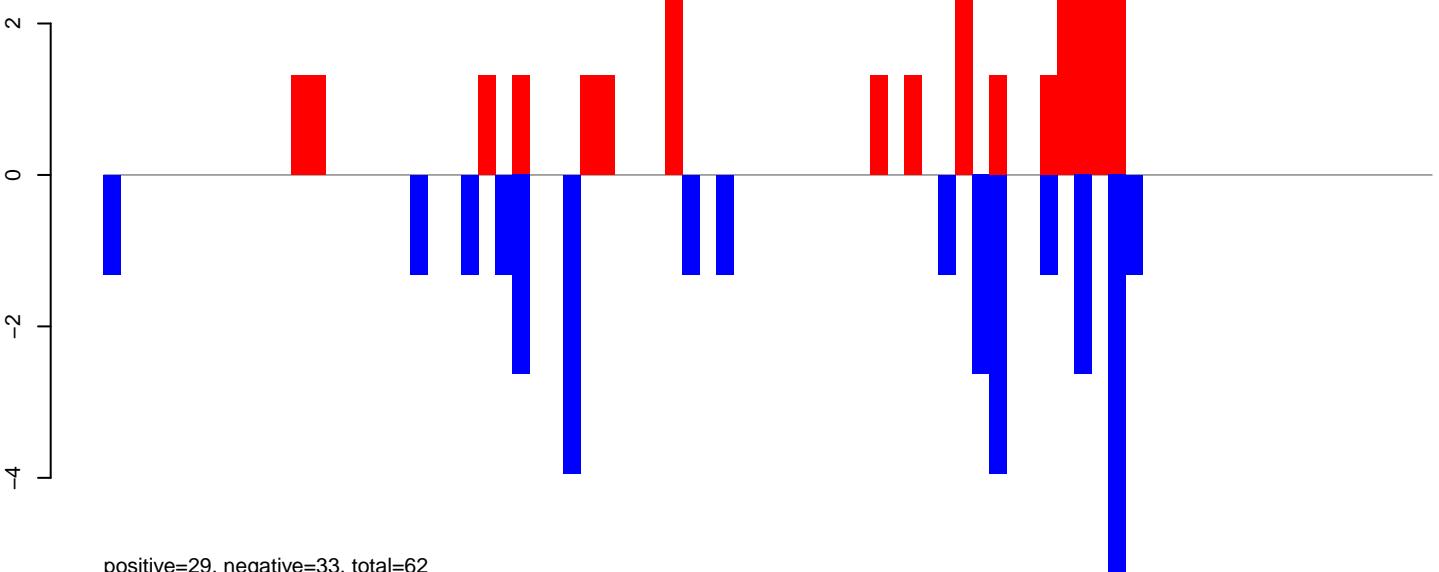


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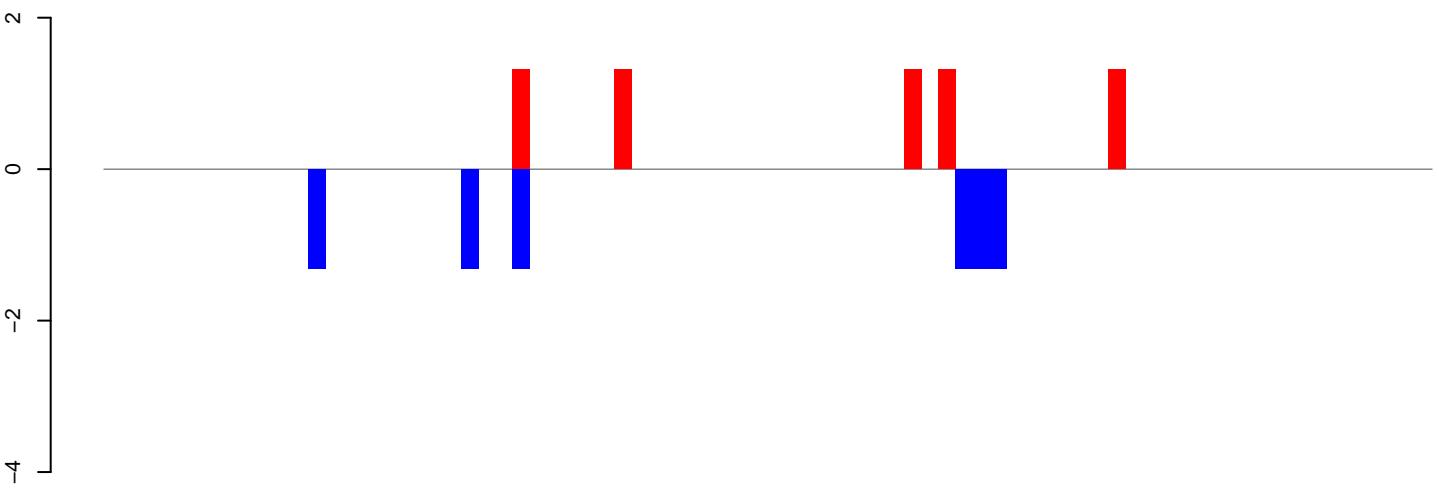


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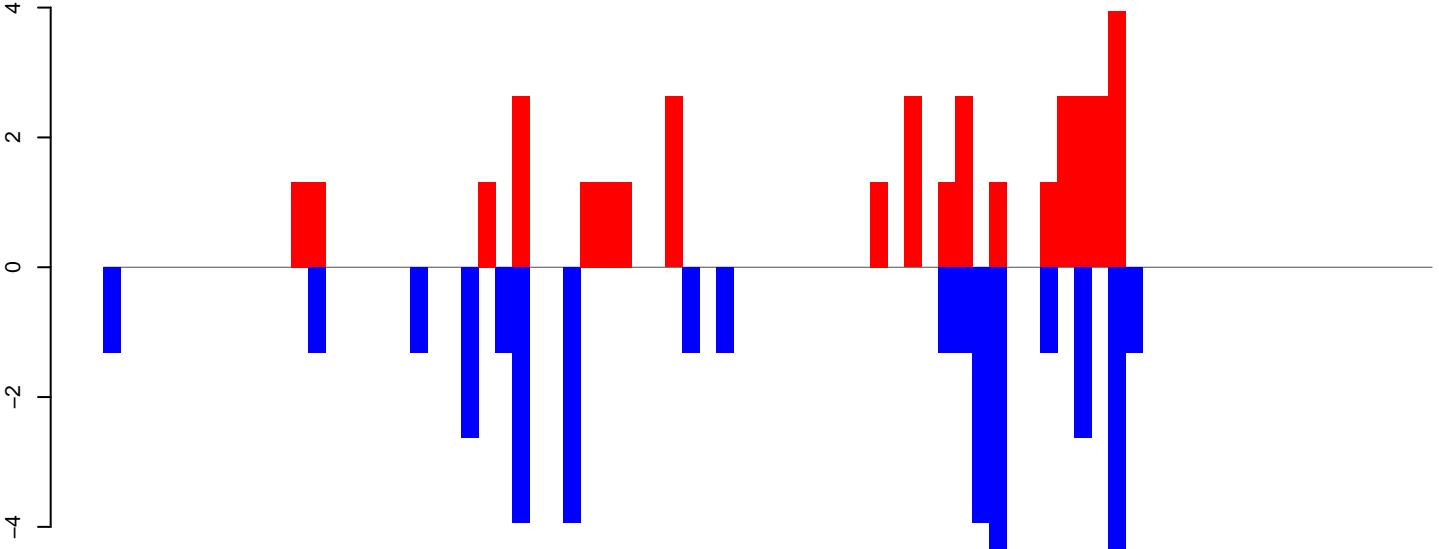


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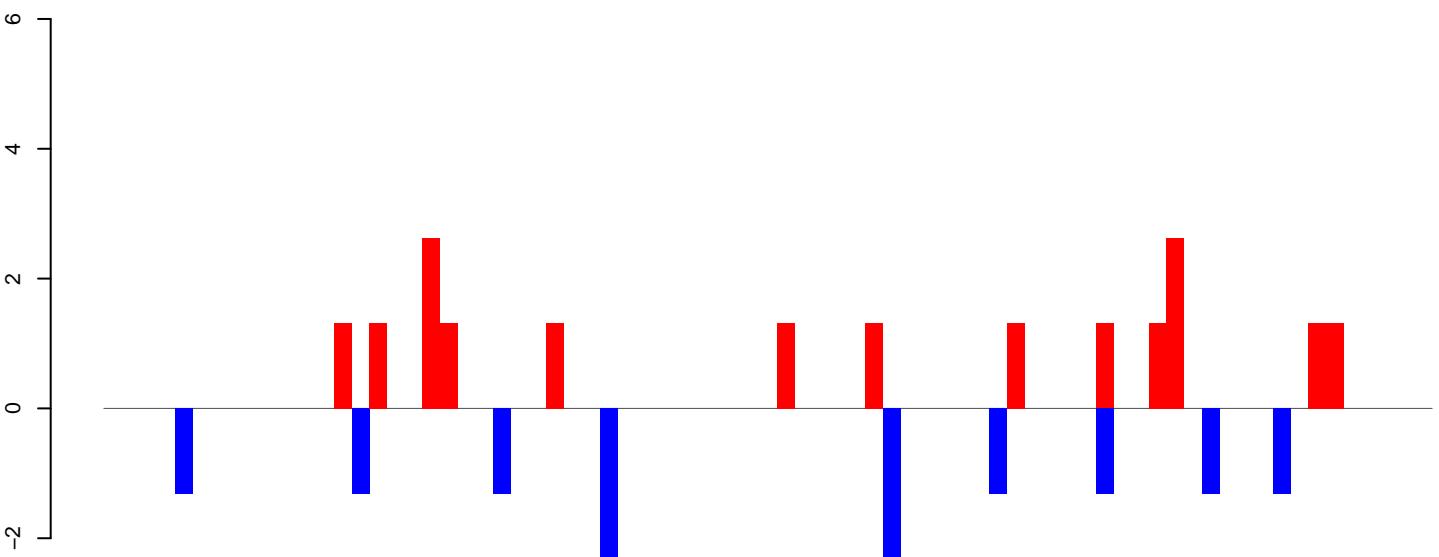
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AnGam\_Fcanc\_TN.rep

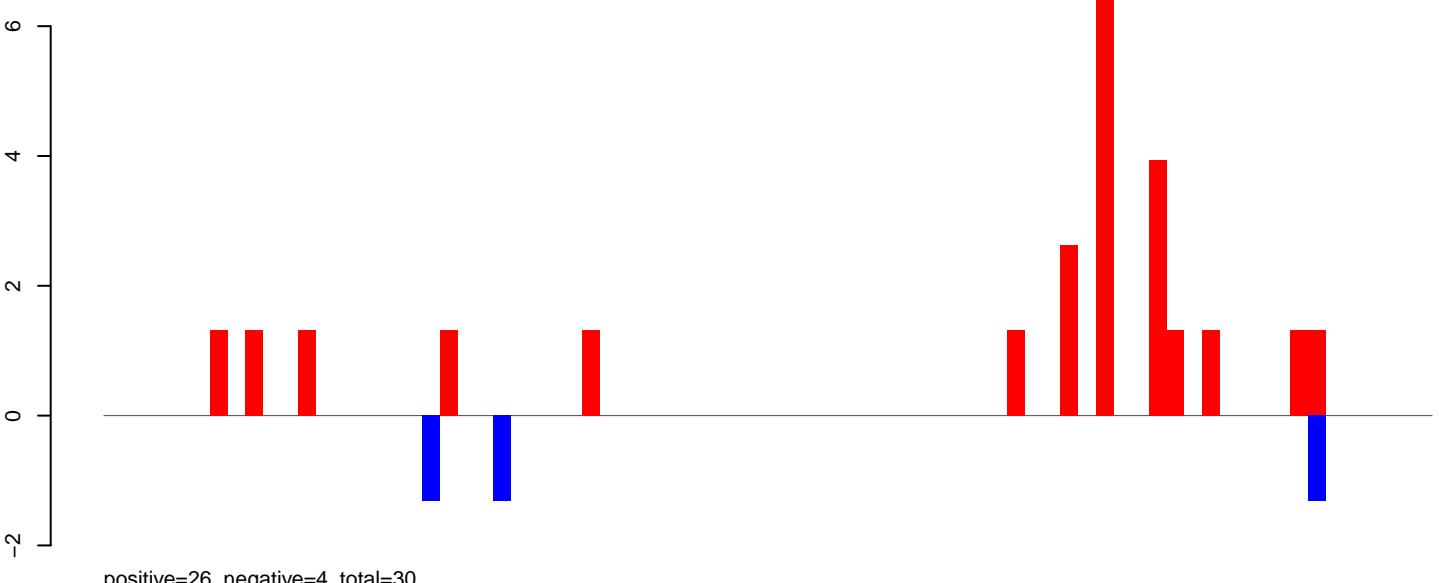


division 32, section 41, 3937-76@us@MH550148.1-Aedes\_albopictus\_densovirus\_isolate\_NC01\_L3\_1\_hmmmed\_contig\_5889-RC:1-3937

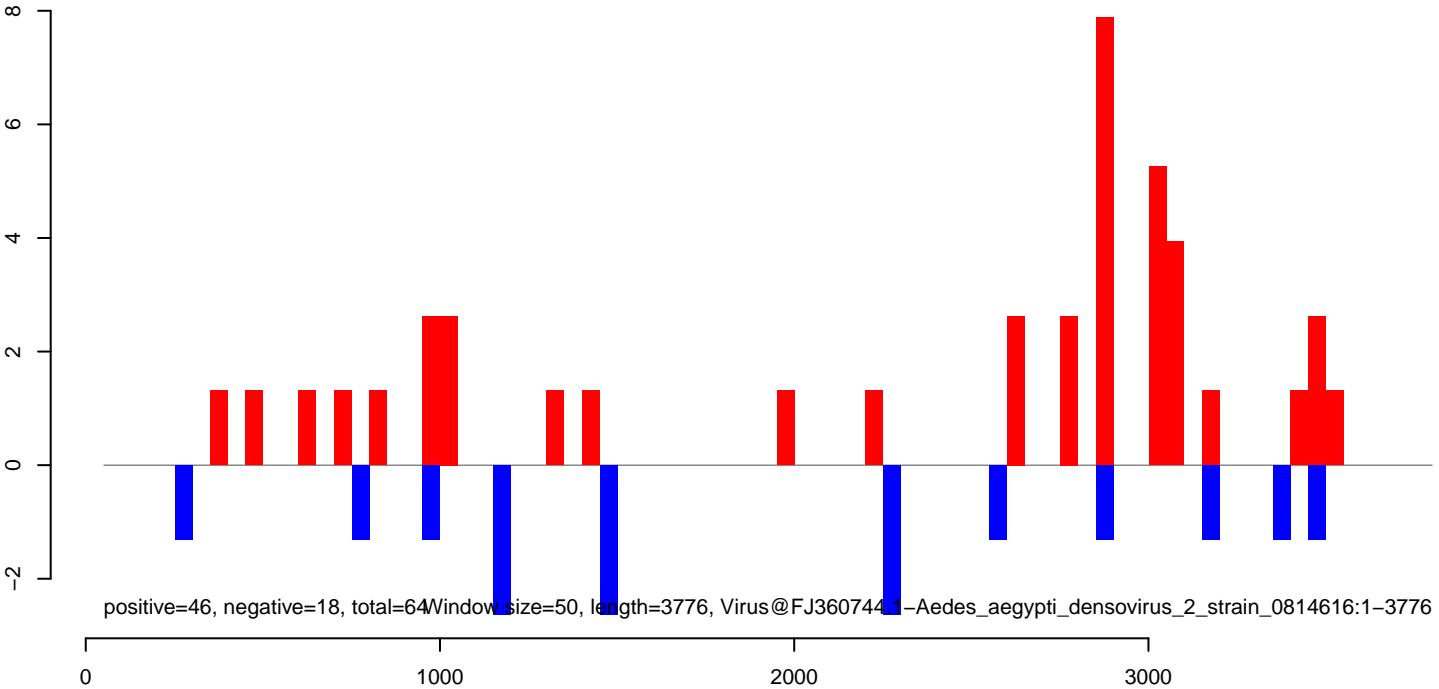
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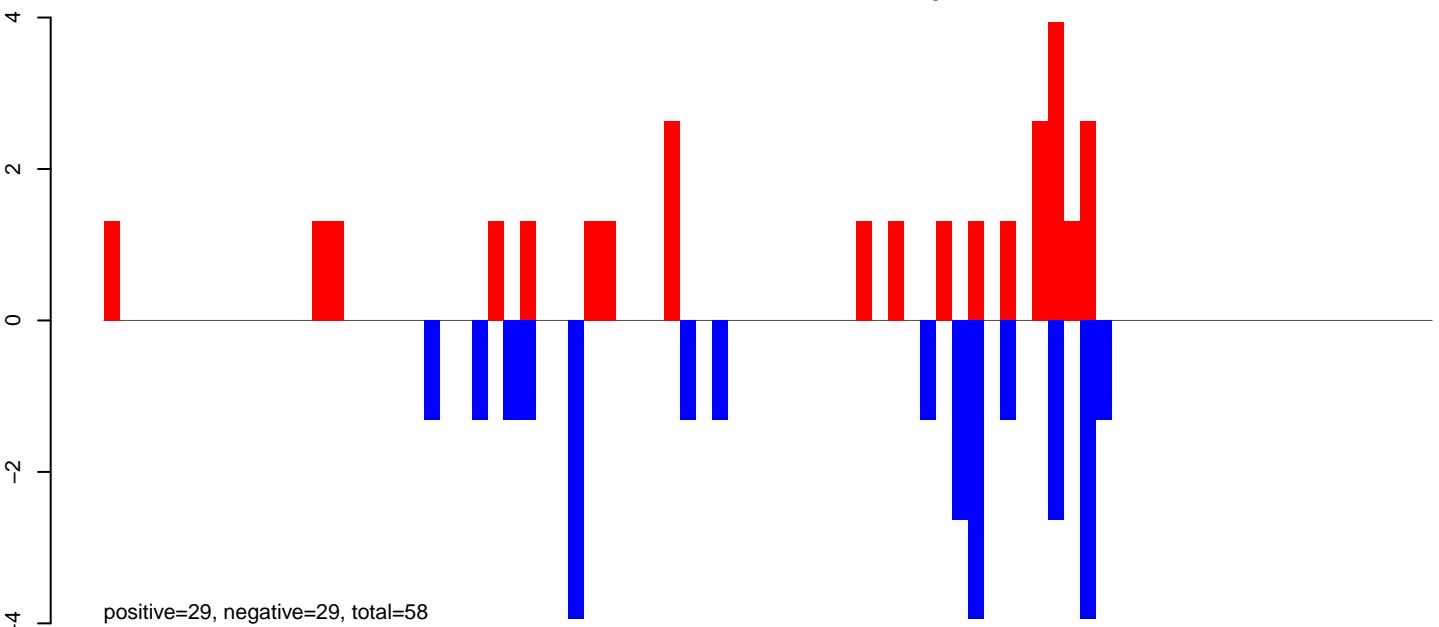
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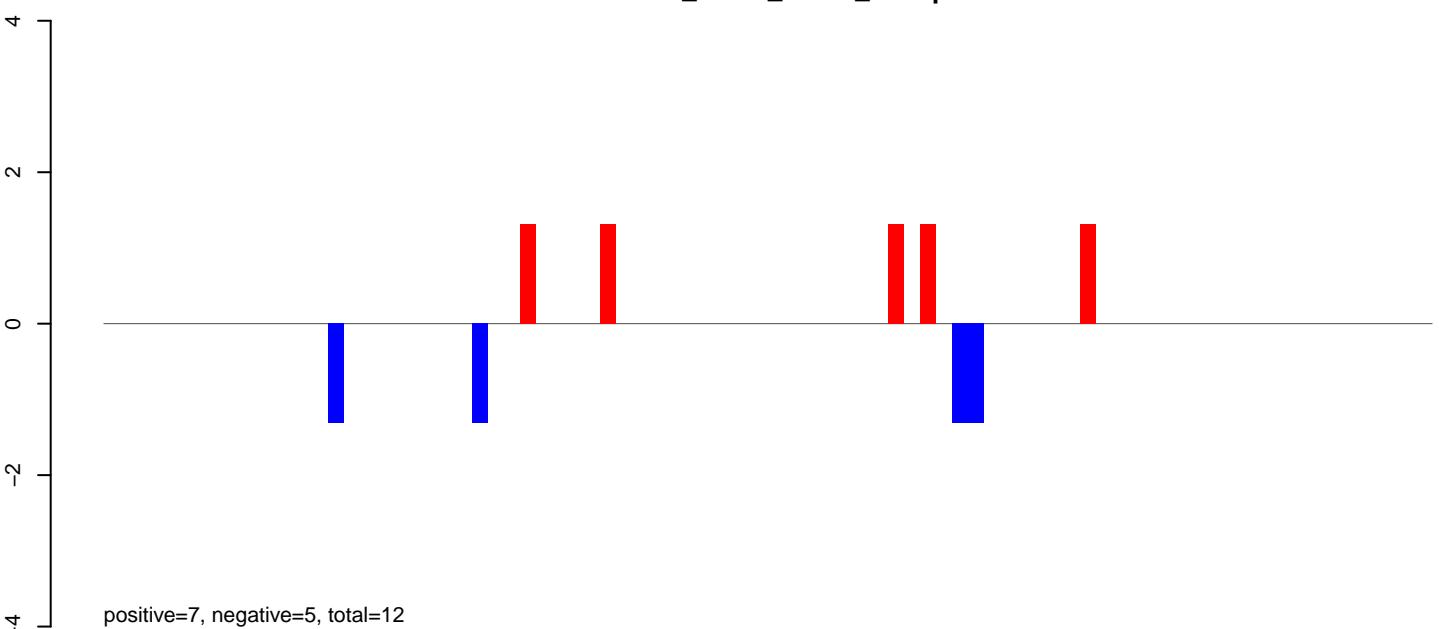
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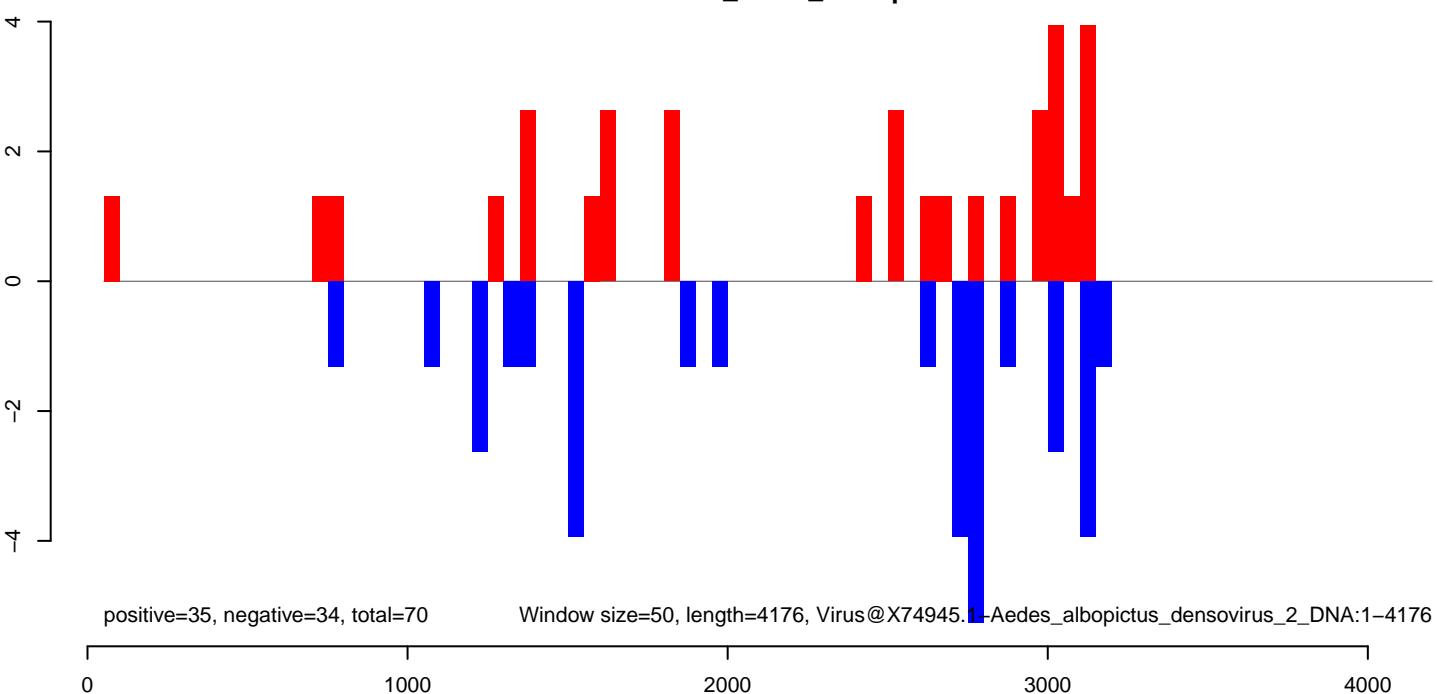
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AnGam\_Fcanc\_TN.24\_35.rep

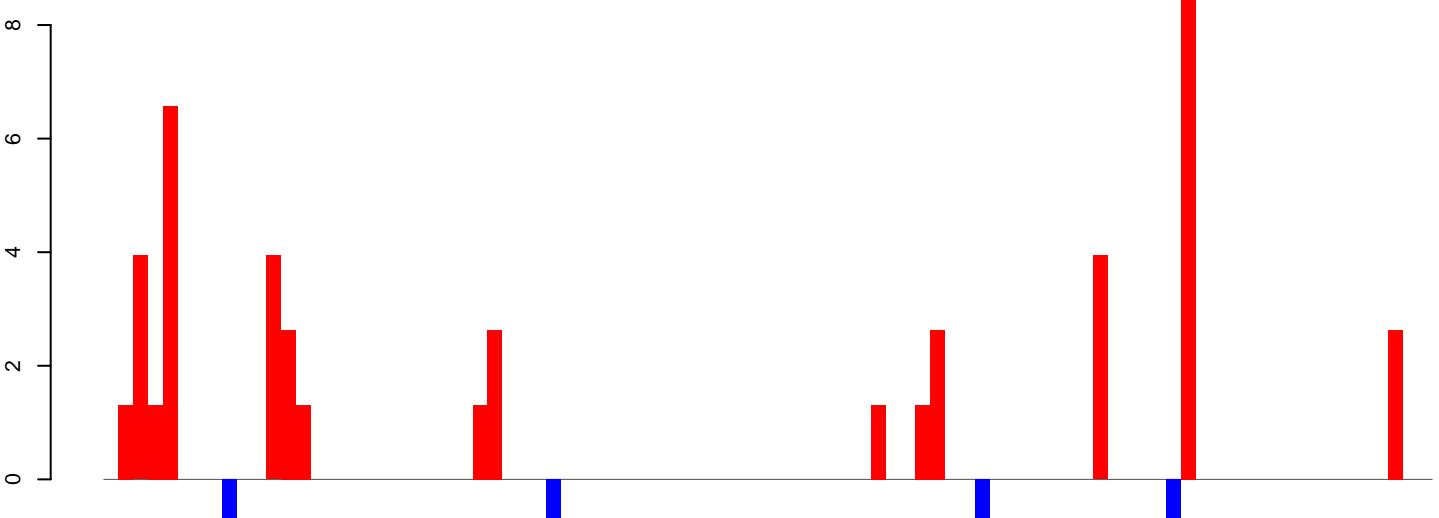


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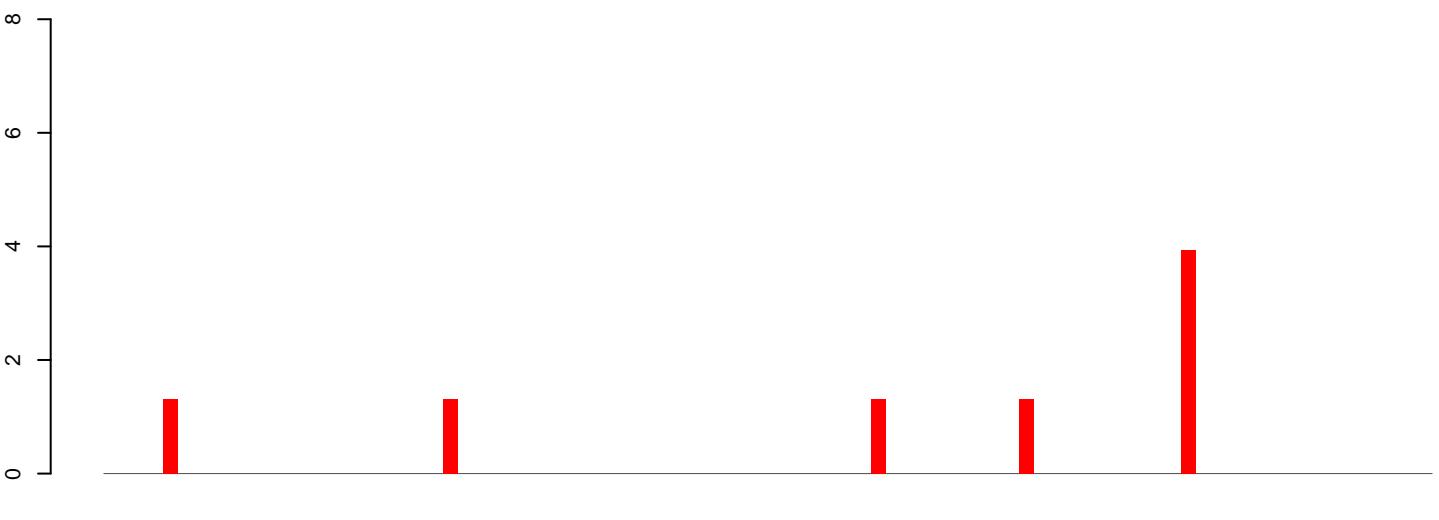
### AnGam\_Fcanc\_TN.18\_23.rep

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### AnGam\_Fcanc\_TN.24\_35.rep

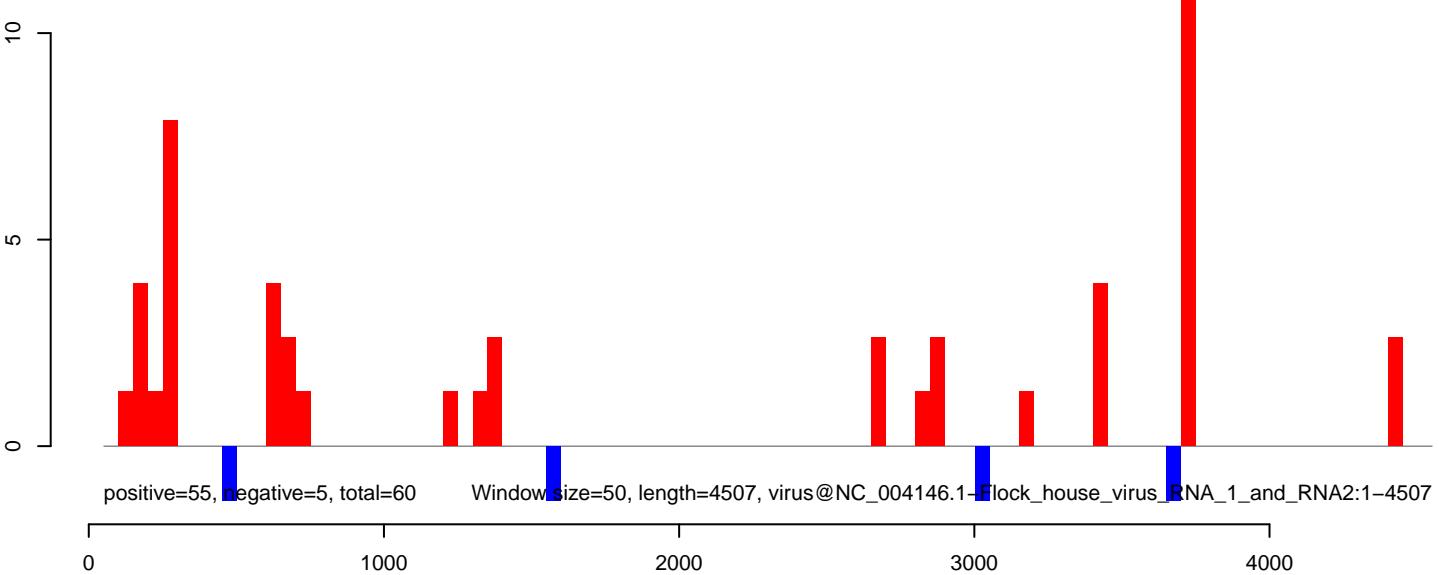
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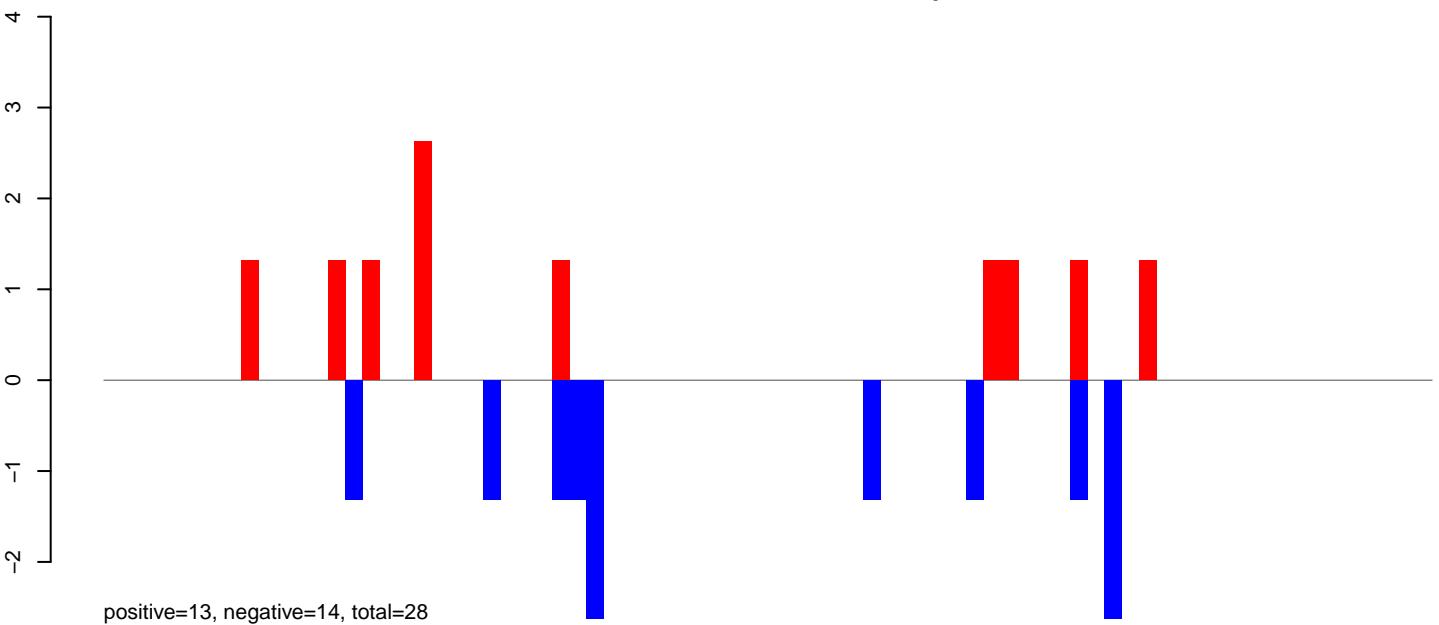
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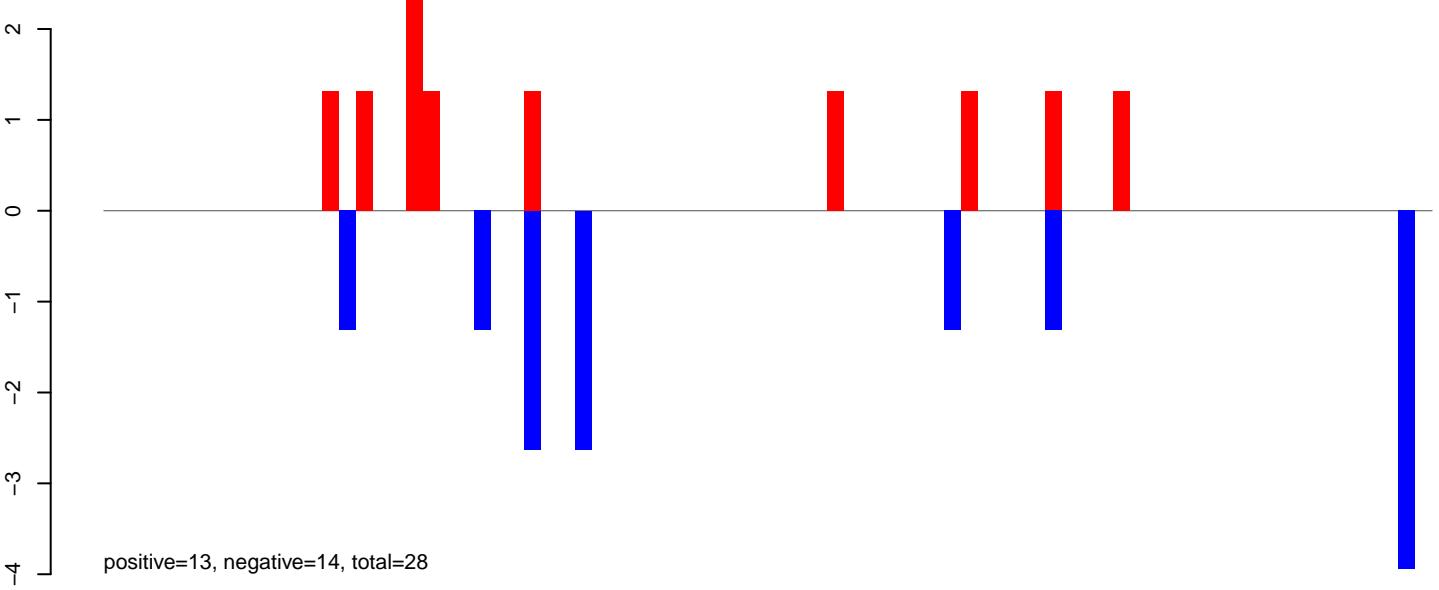
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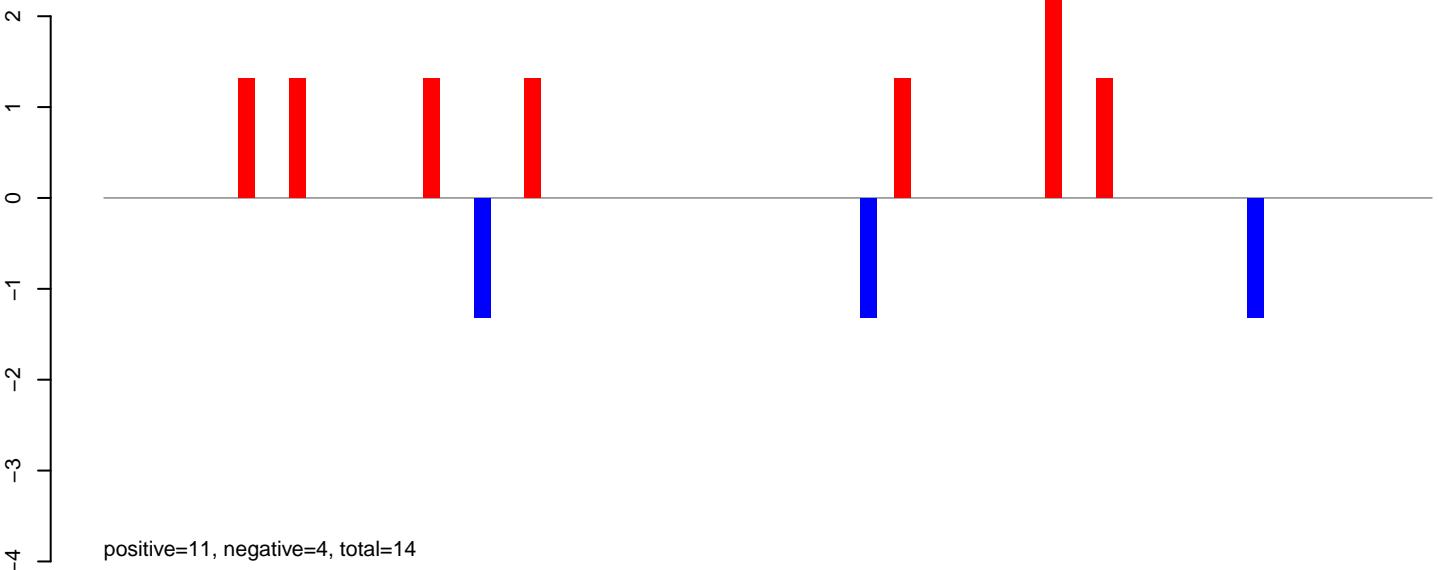
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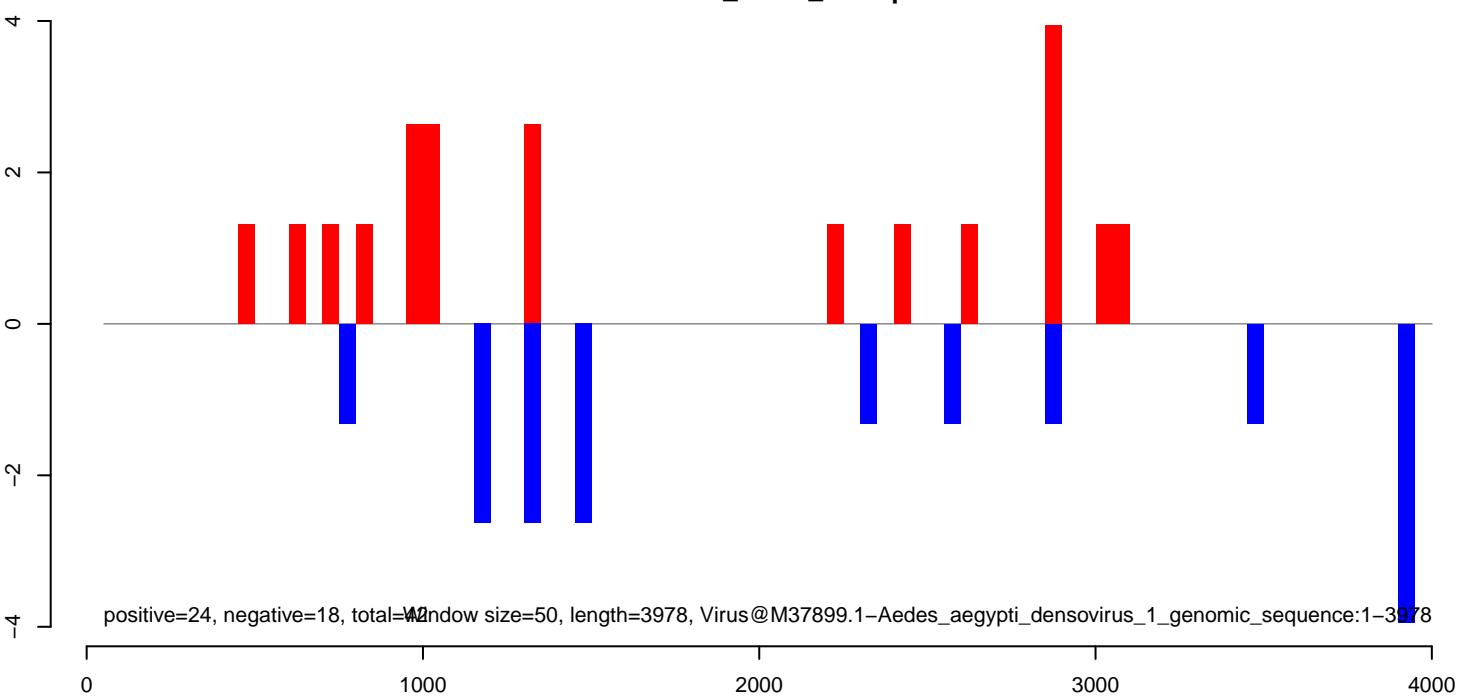
### AnGam\_Fcanc\_TN.18\_23.rep



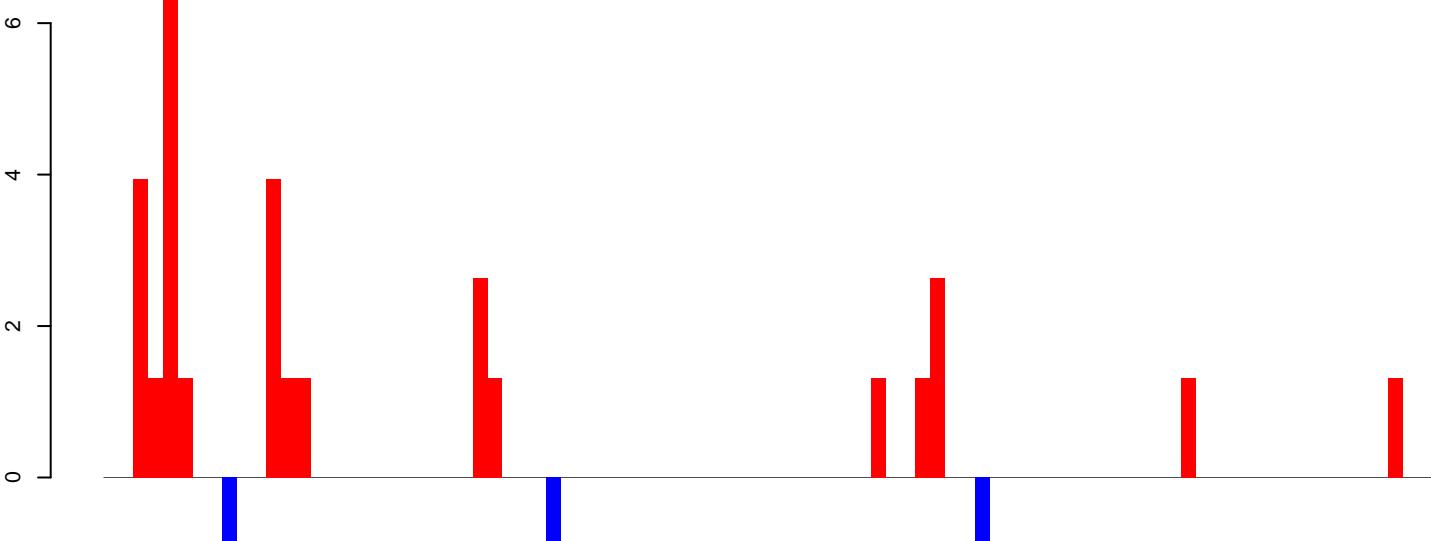
### AnGam\_Fcanc\_TN.24\_35.rep



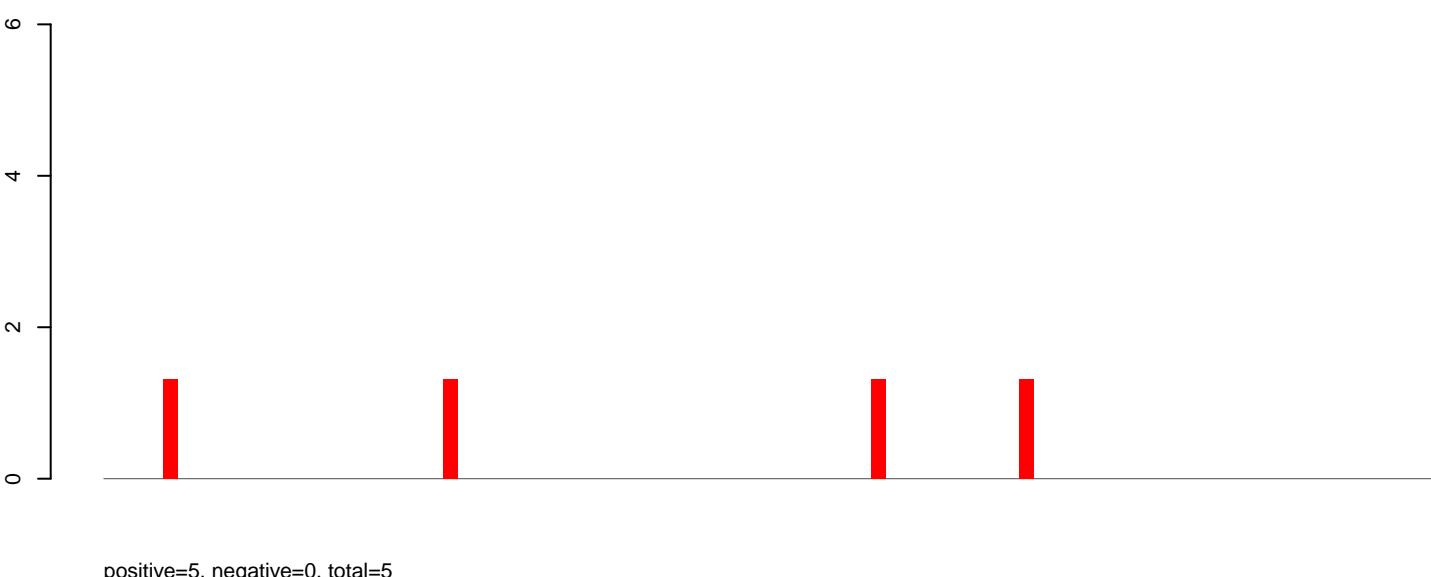
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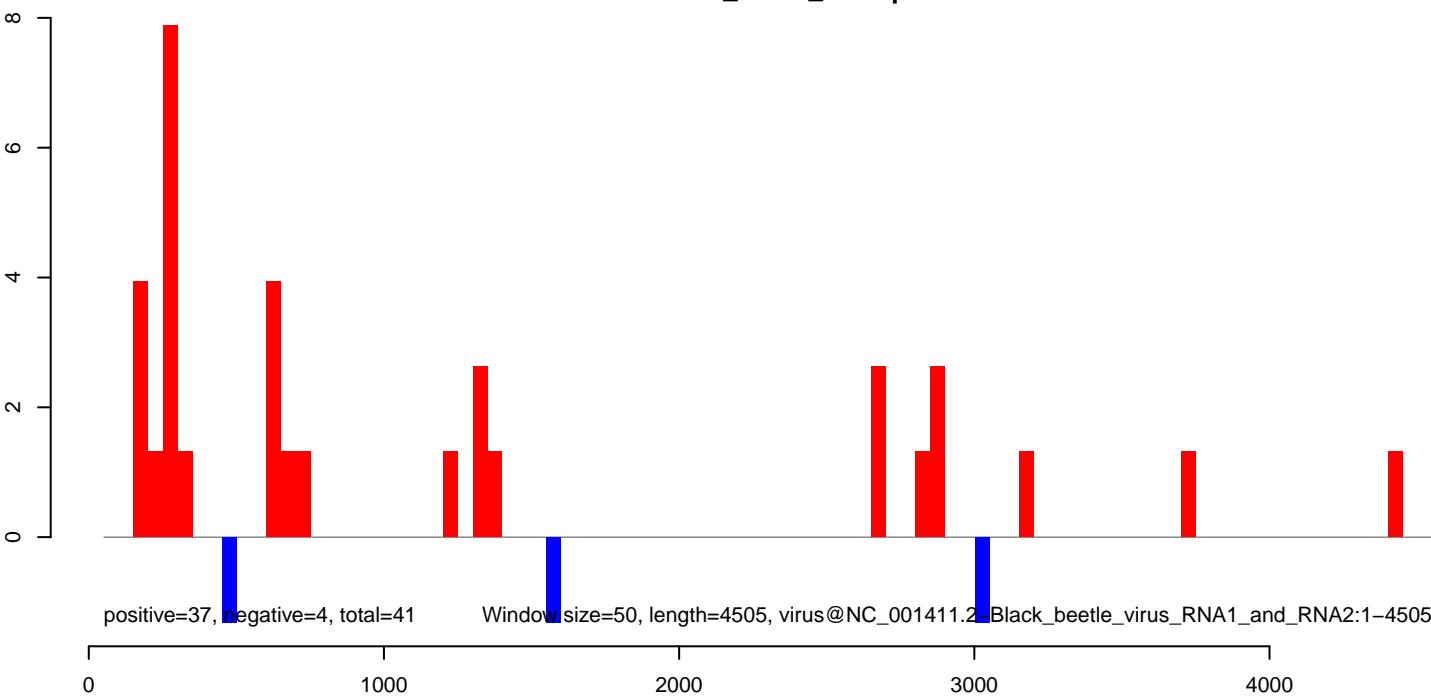
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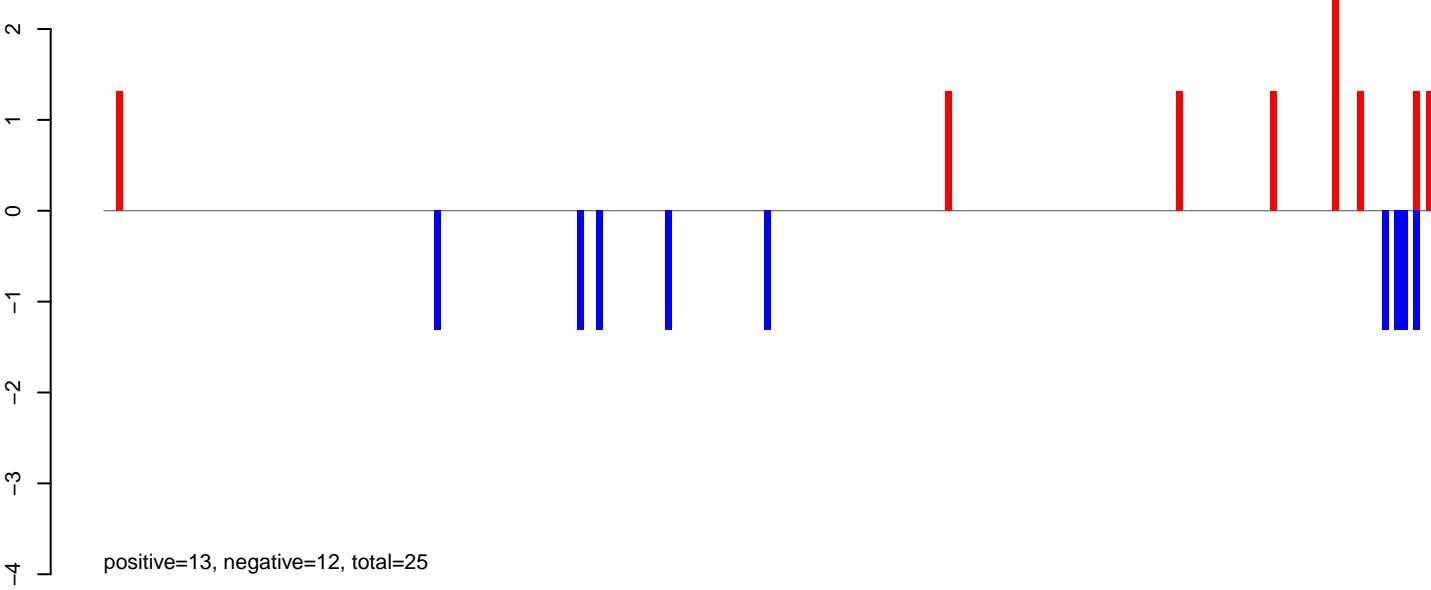
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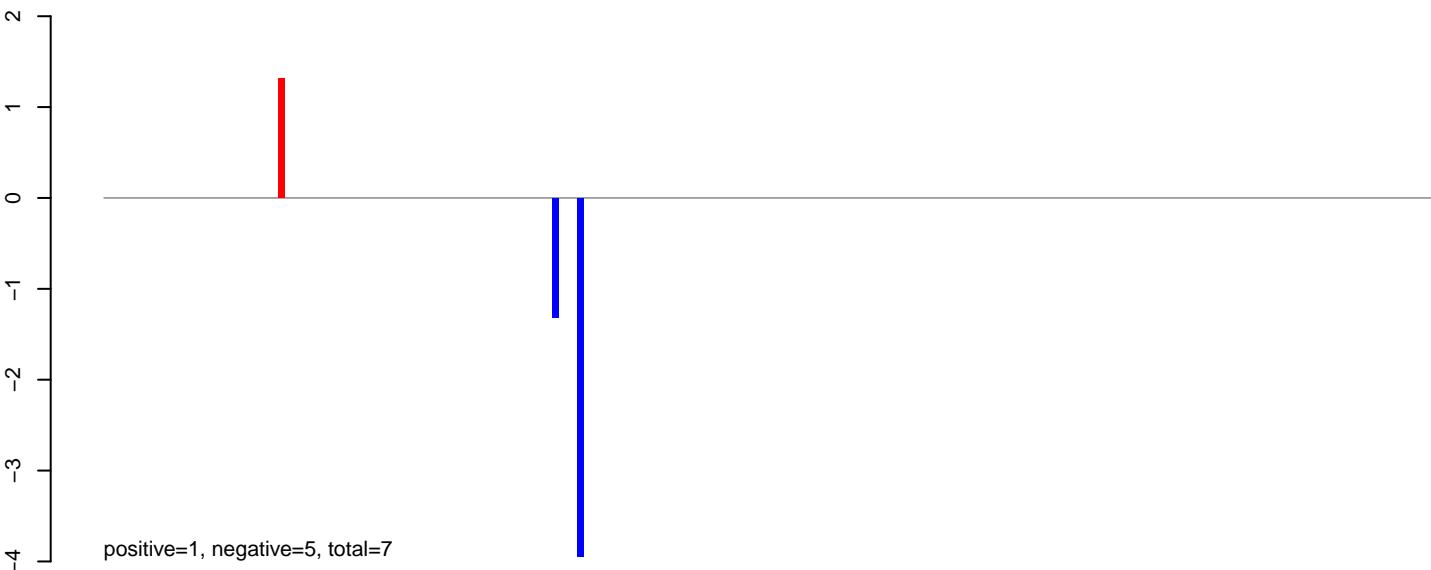
### AnGam\_Fcanc\_TN.rep



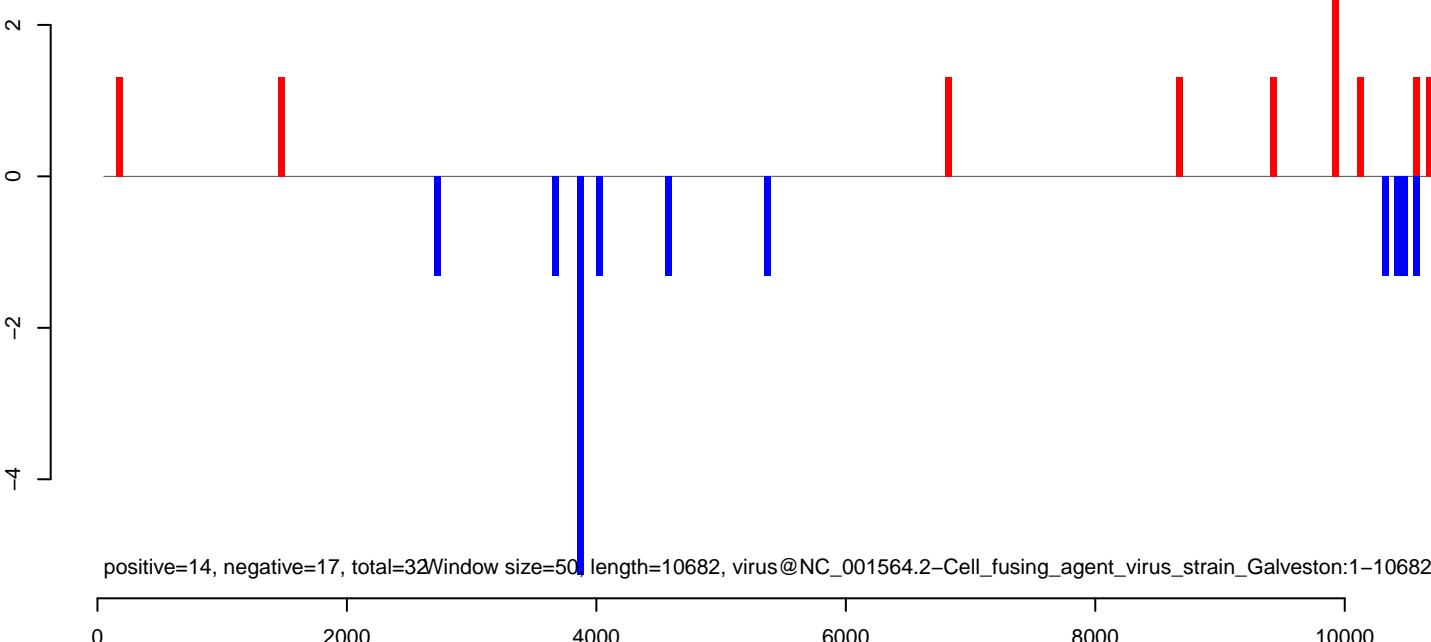
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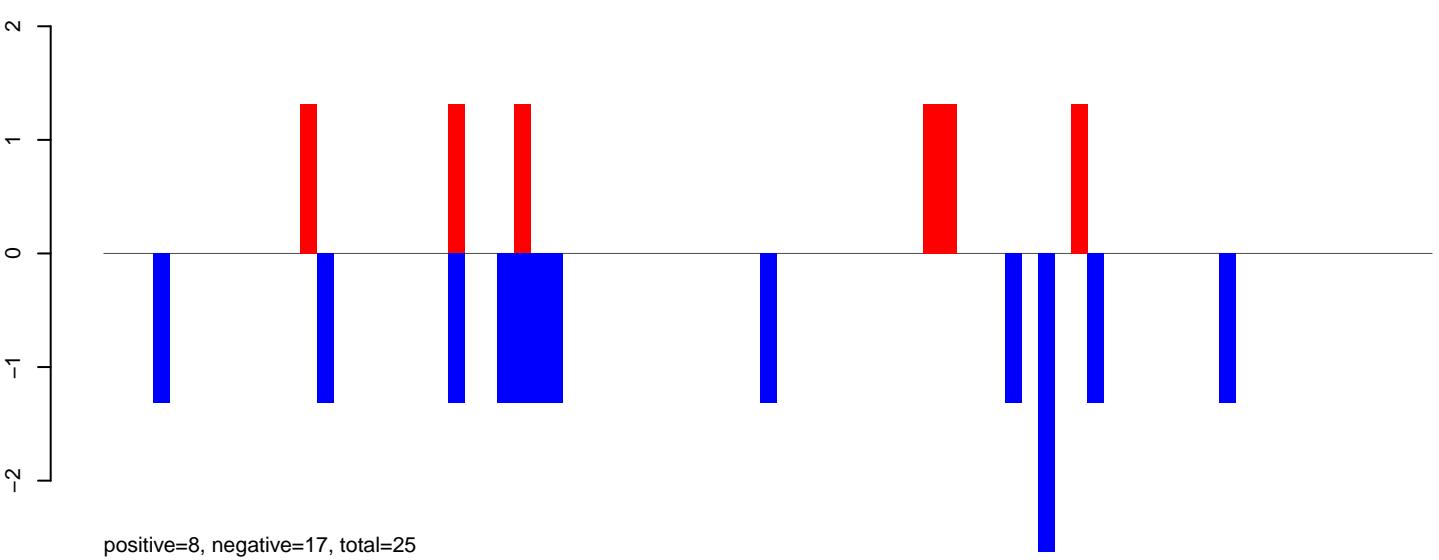
AnGam\_Fcanc\_TN.24\_35.rep



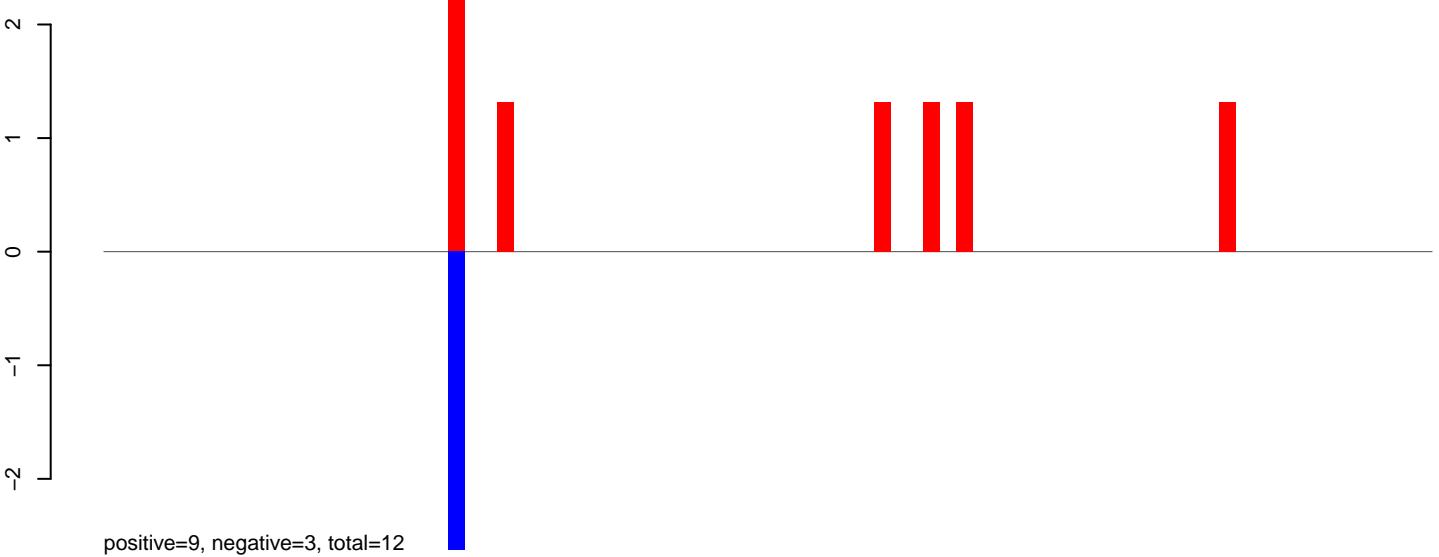
AnGam\_Fcanc\_TN.rep



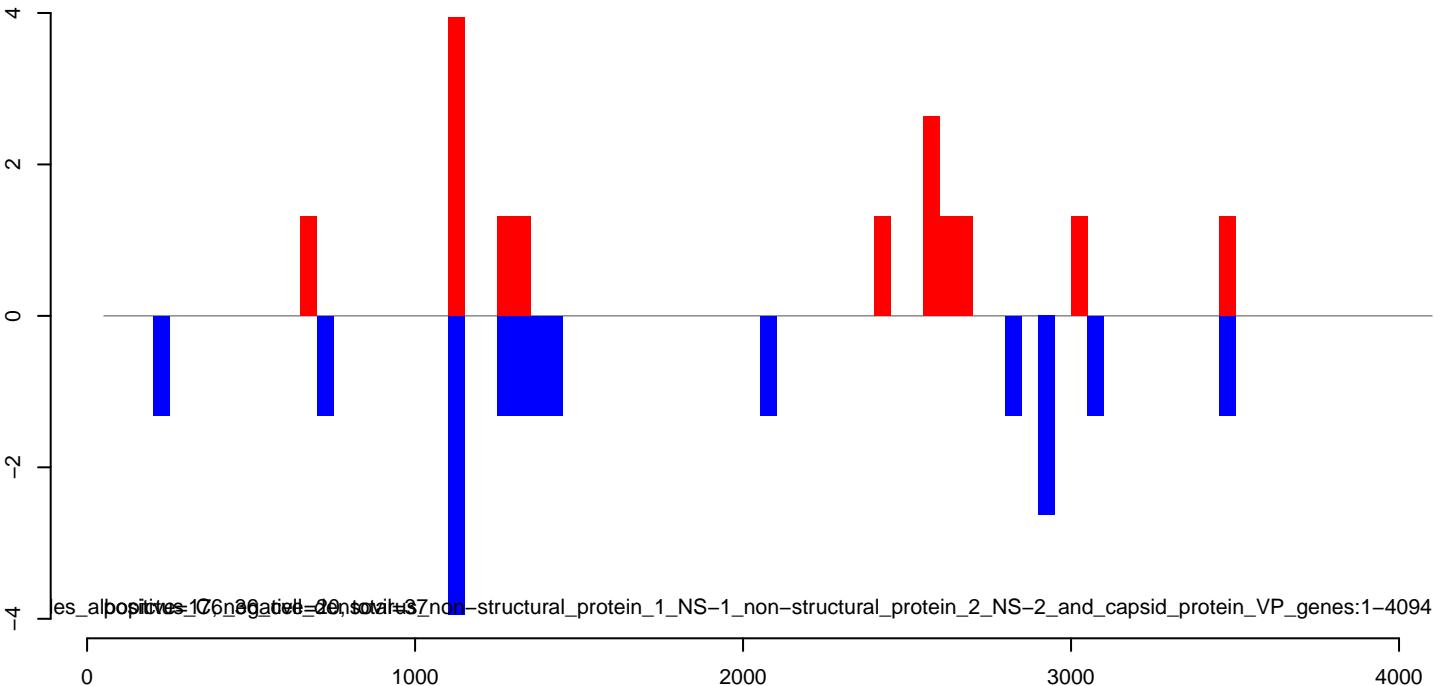
### AnGam\_Fcanc\_TN.18\_23.rep



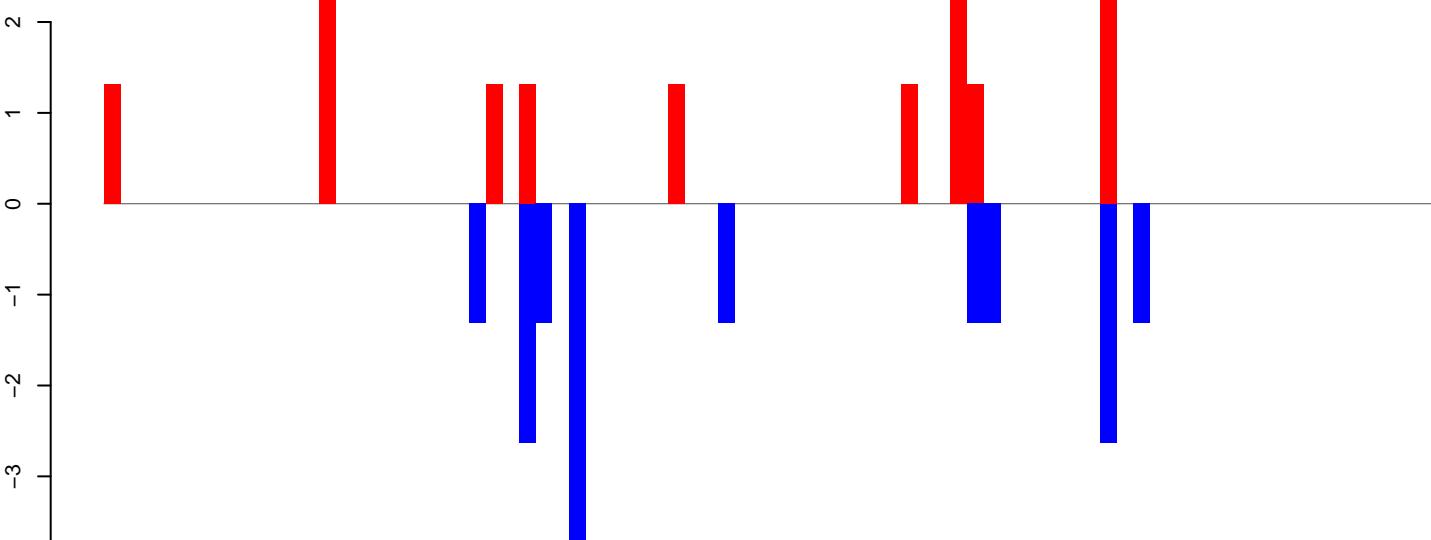
### AnGam\_Fcanc\_TN.24\_35.rep



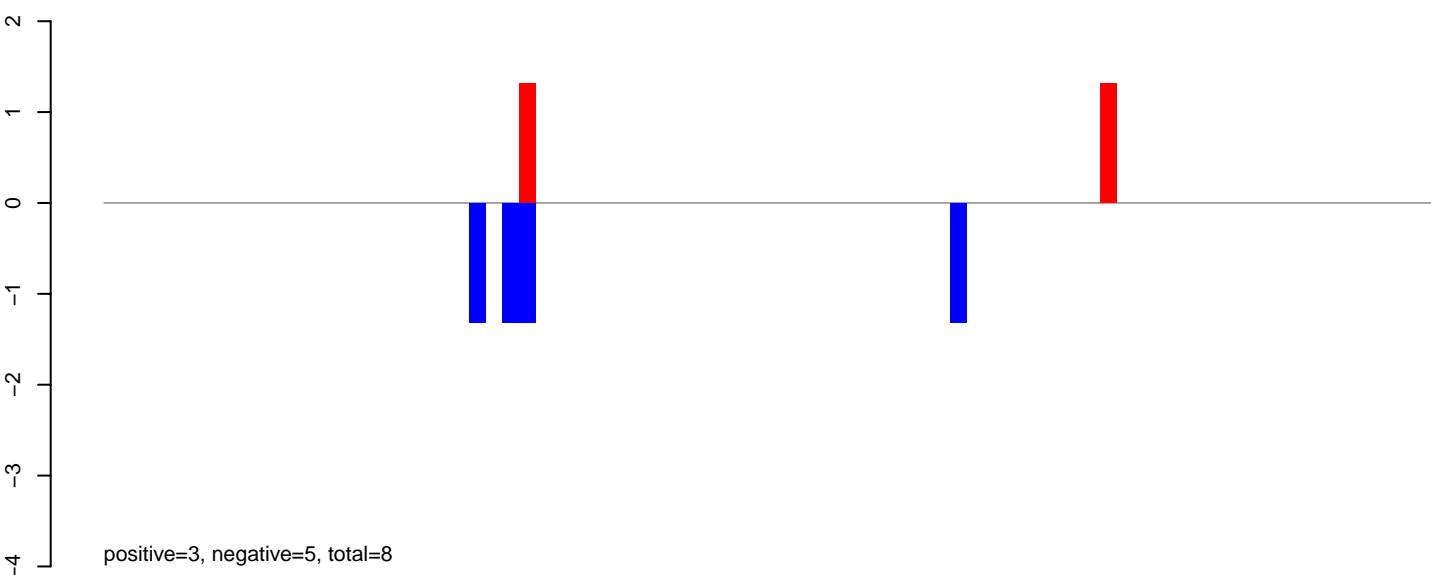
### AnGam\_Fcanc\_TN.rep



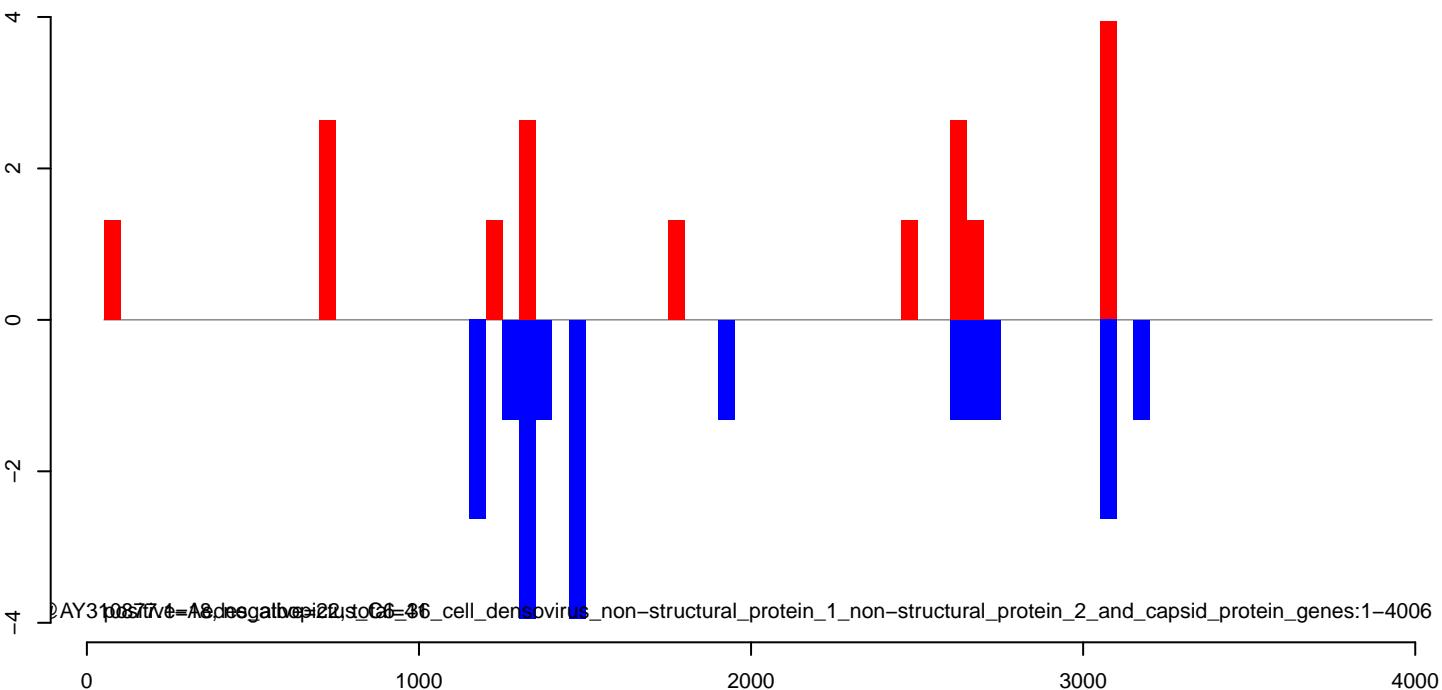
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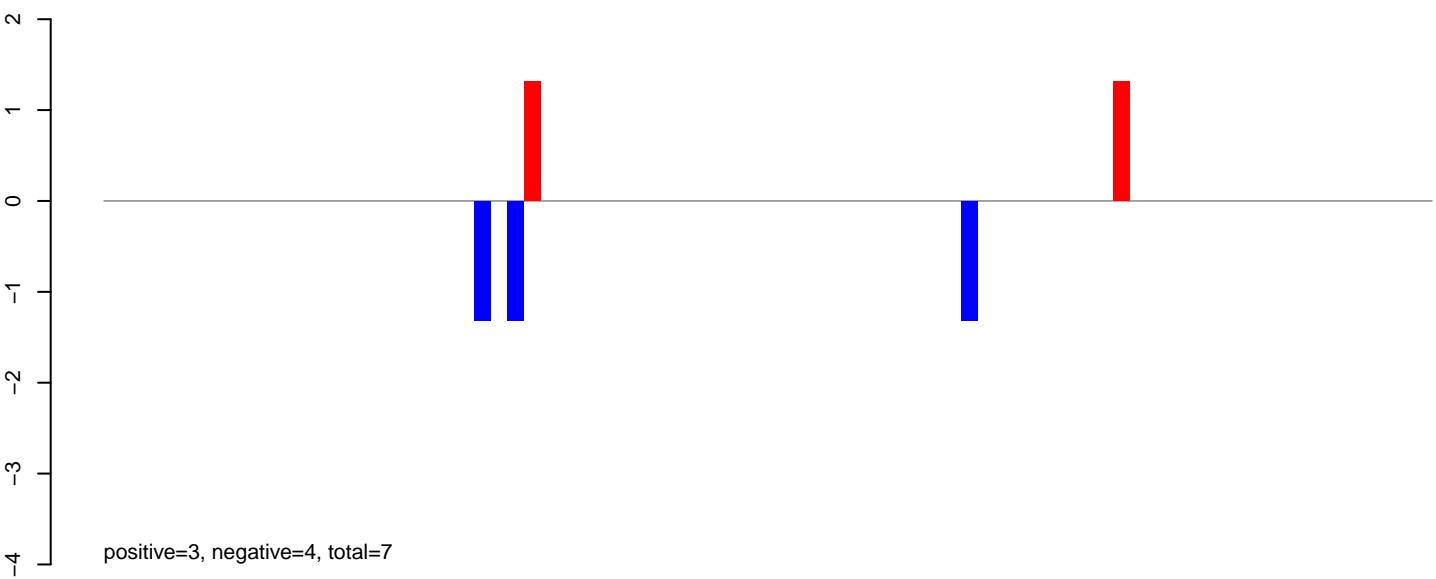
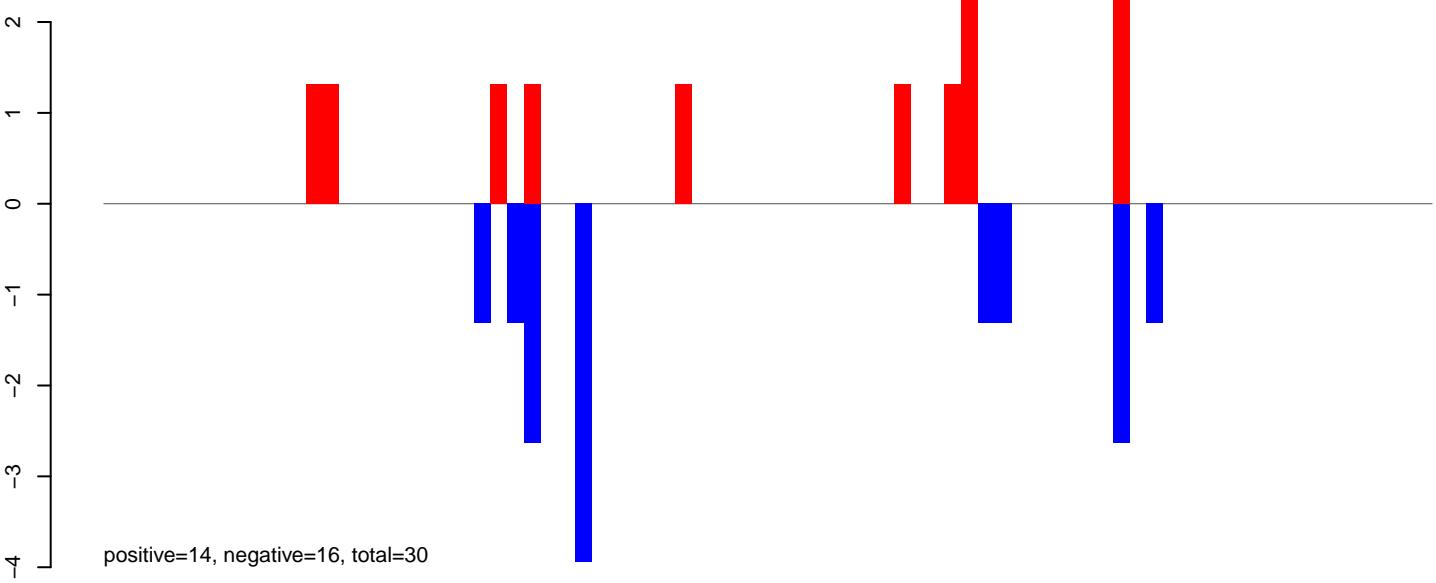
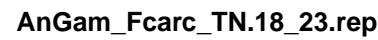


AnGam\_Fcanc\_TN.24\_35.rep

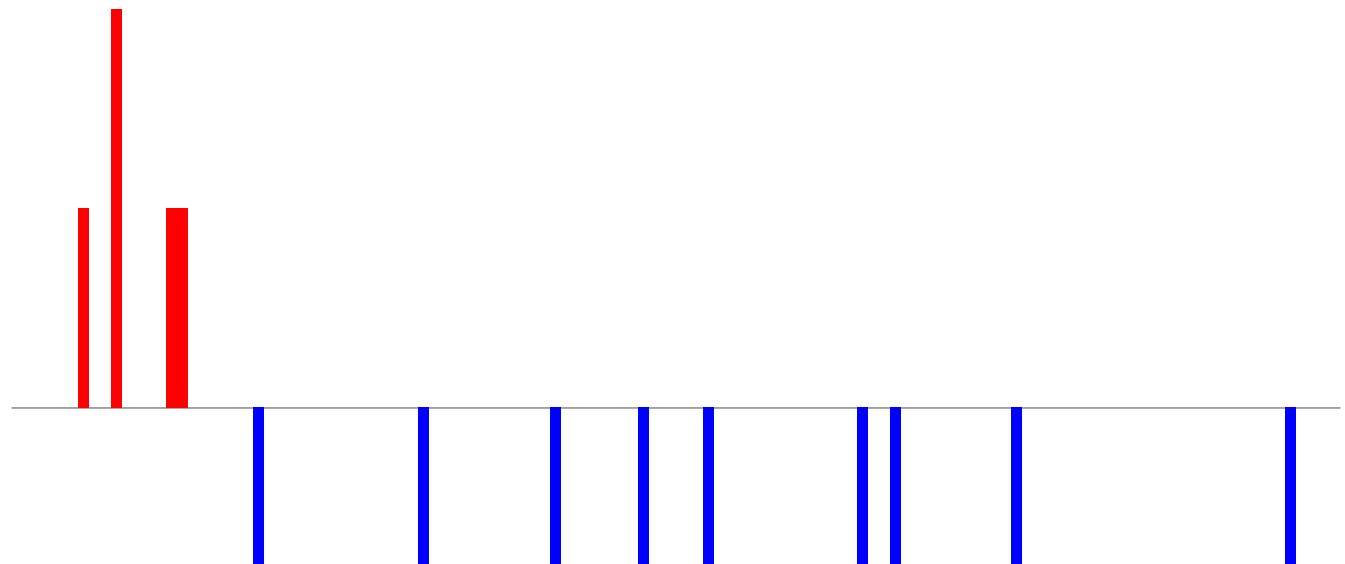


AnGam\_Fcanc\_TN.rep

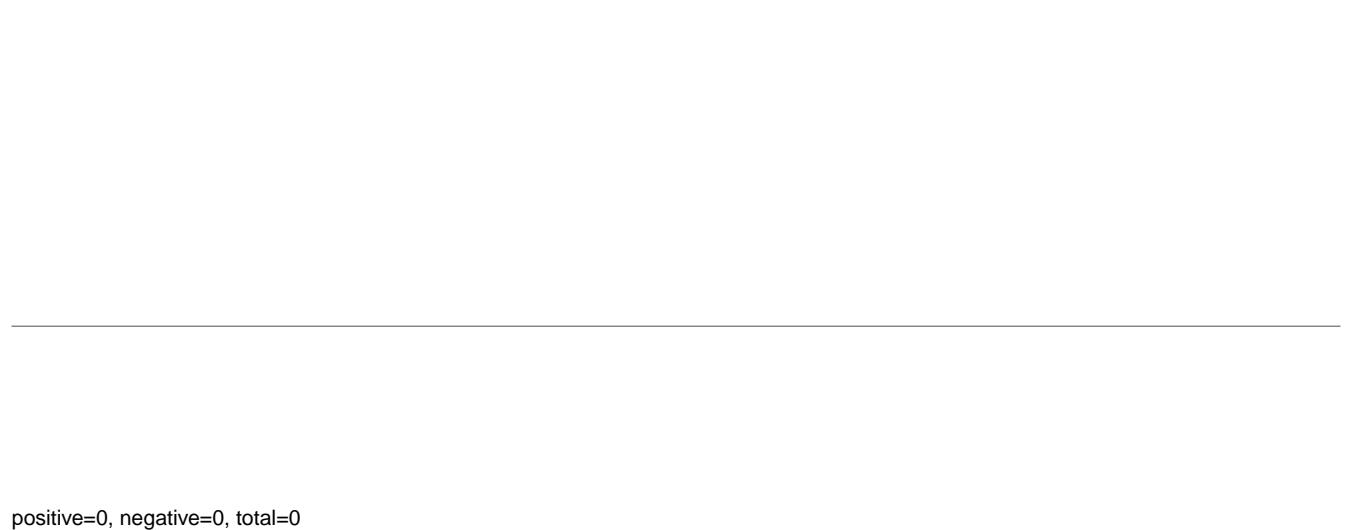




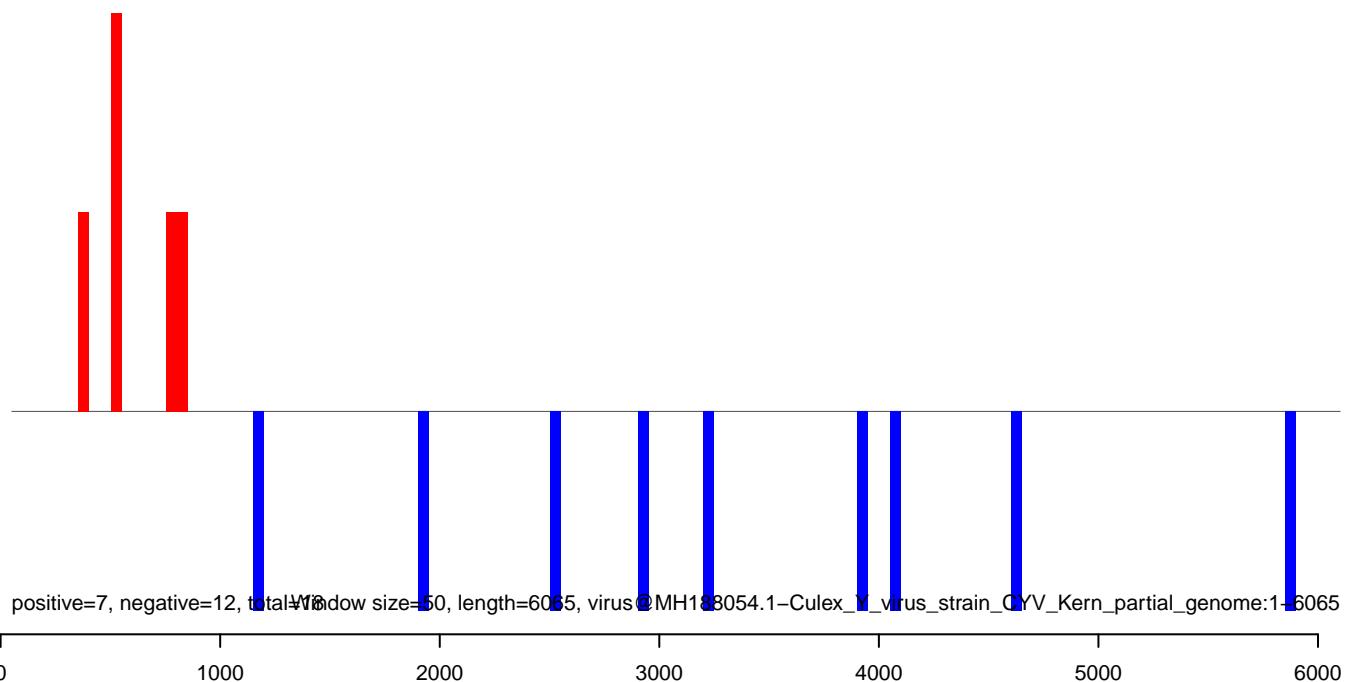
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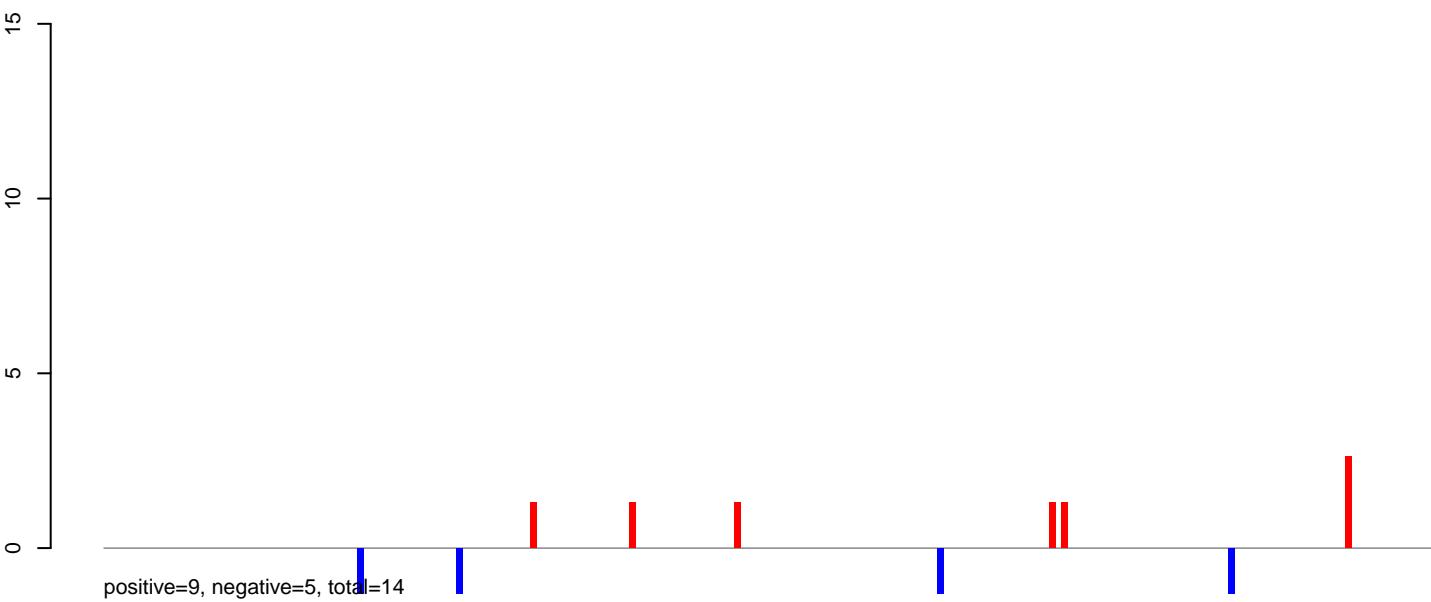
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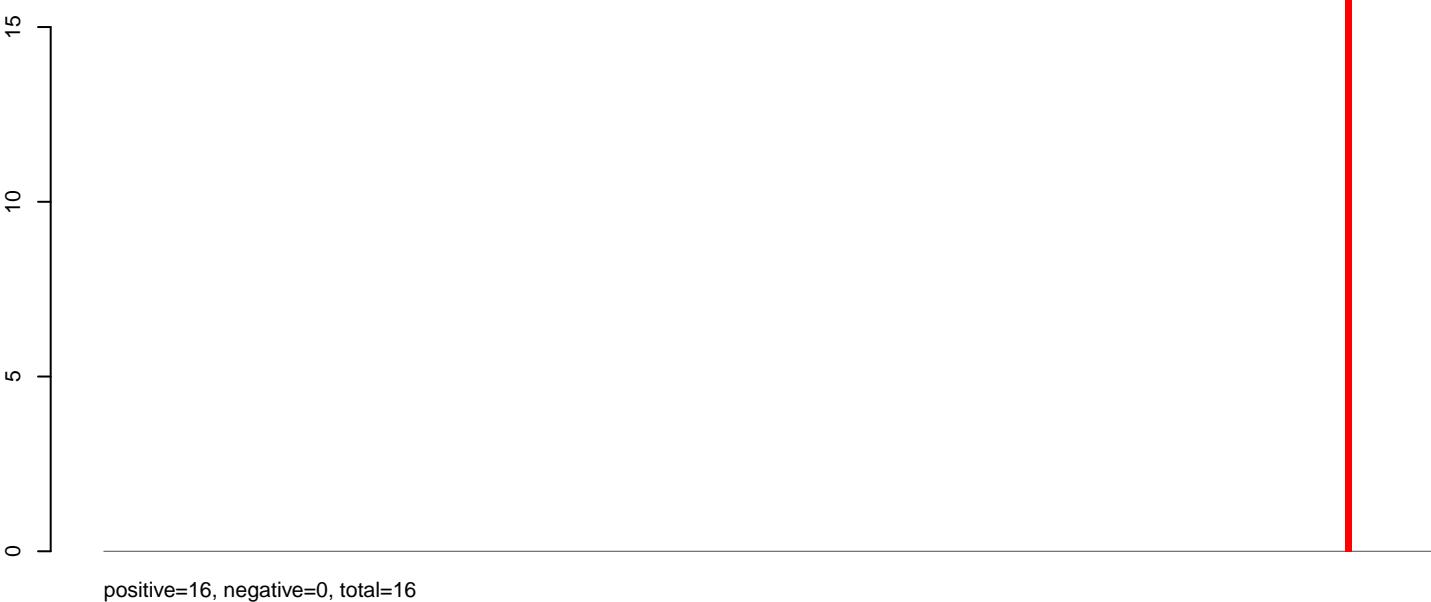
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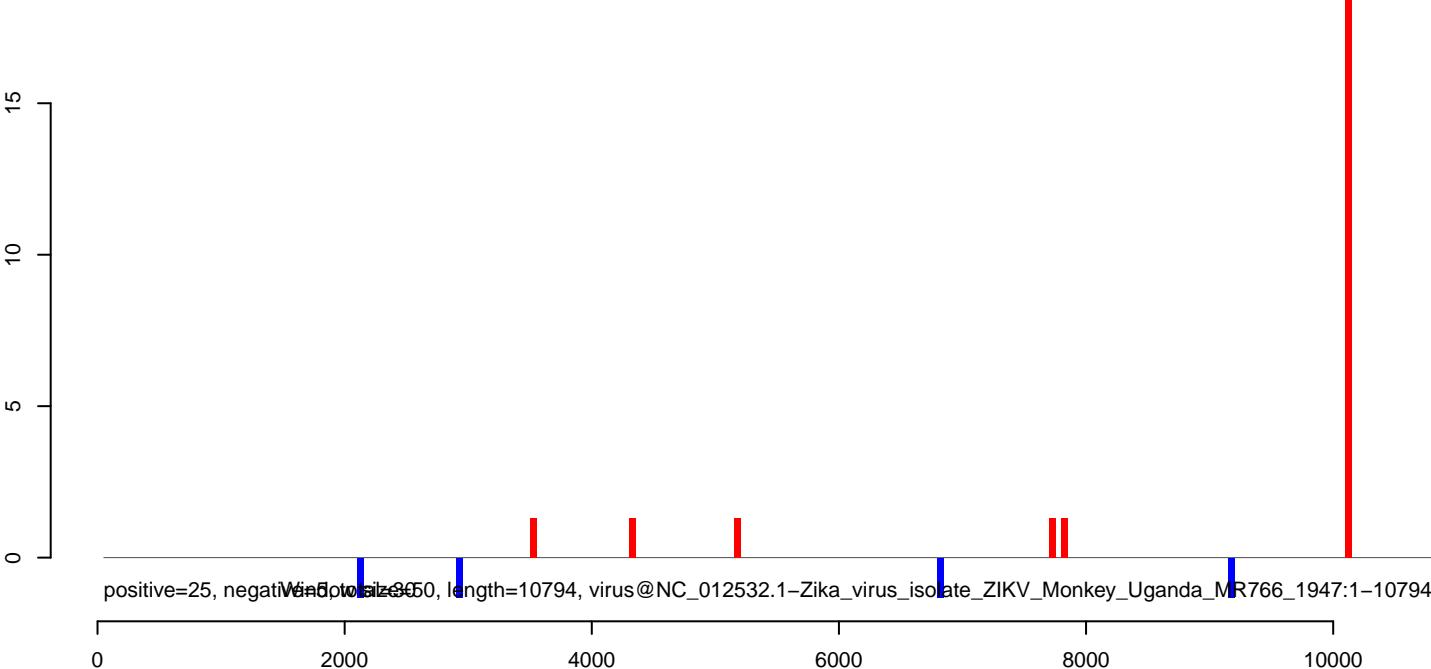
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### AnGam\_Fcanc\_TN.24\_35.rep

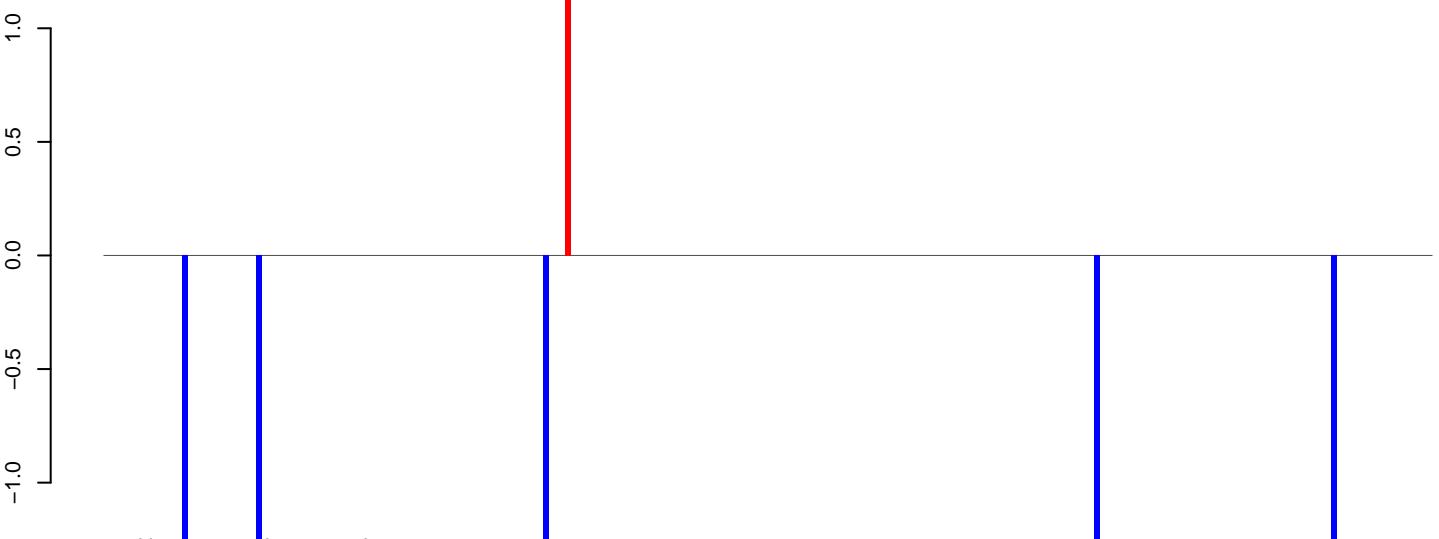


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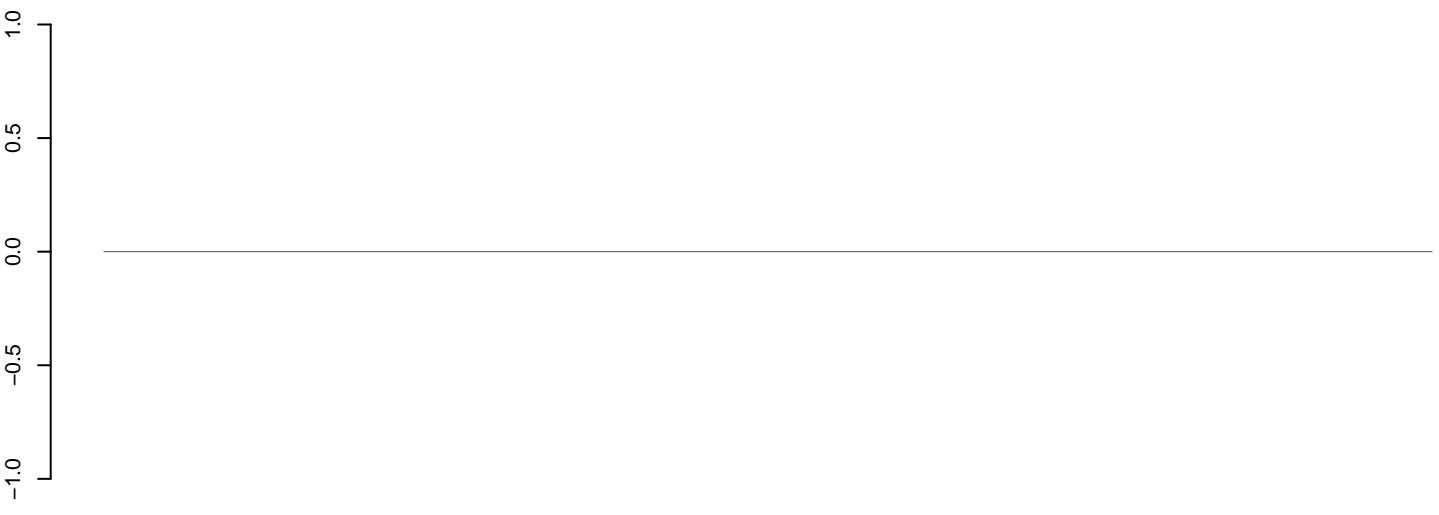
### AnGam\_Fcanc\_TN.18\_23.rep

positive=1, negative=7, total=8



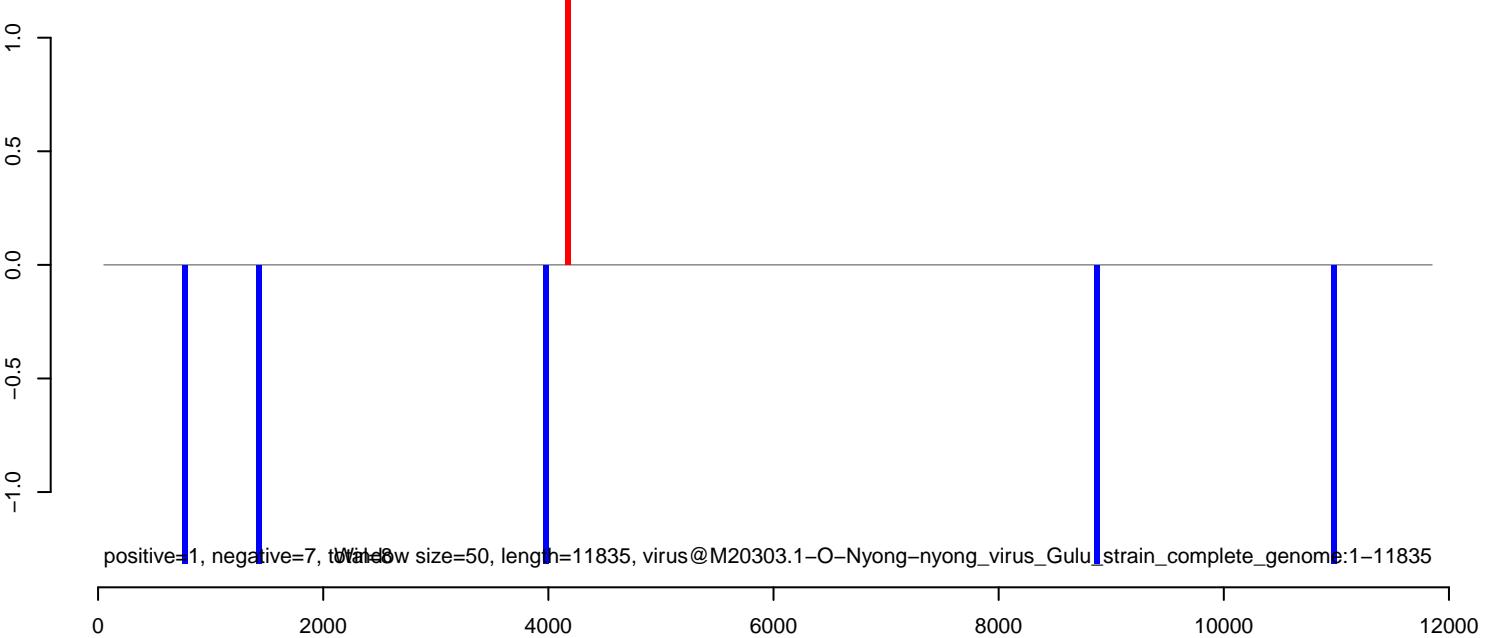
### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

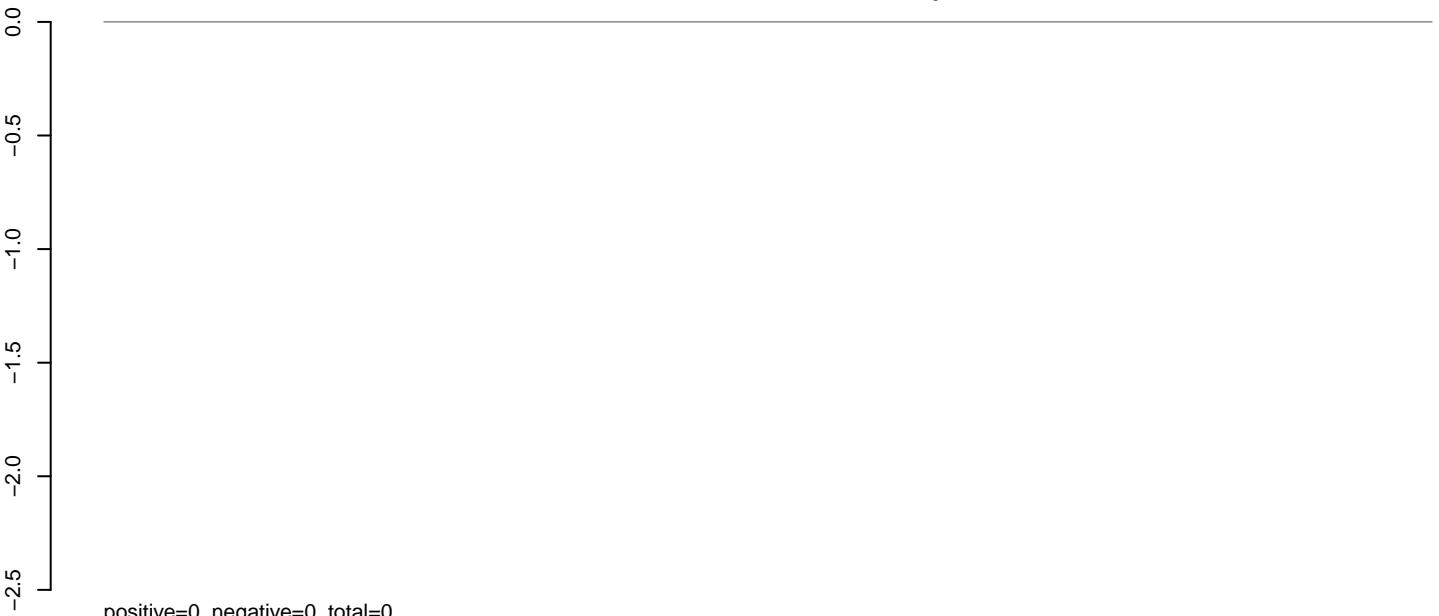


### AnGam\_Fcanc\_TN.rep

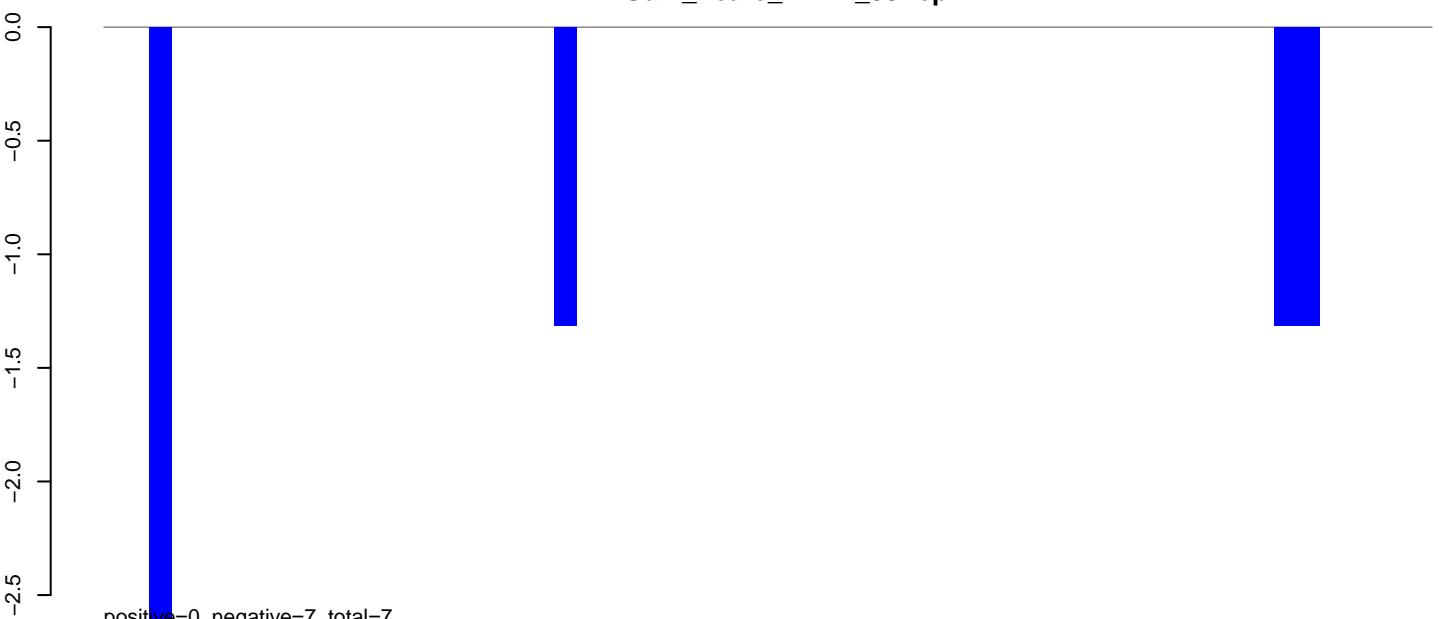
positive=1, negative=7, total=8, window size=50, length=11835, virus@M20303.1-O-Nyong-nyong\_virus\_Gulu strain\_complete\_genome:1-11835



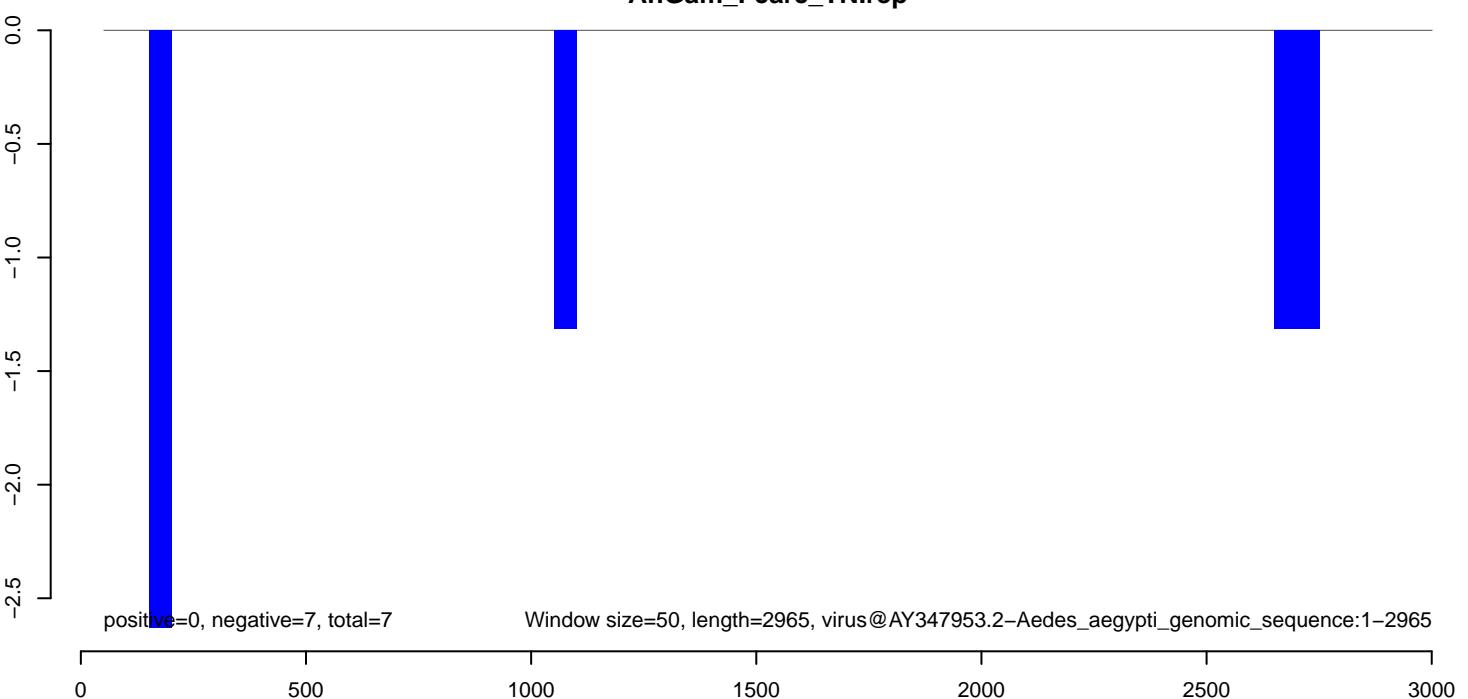
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



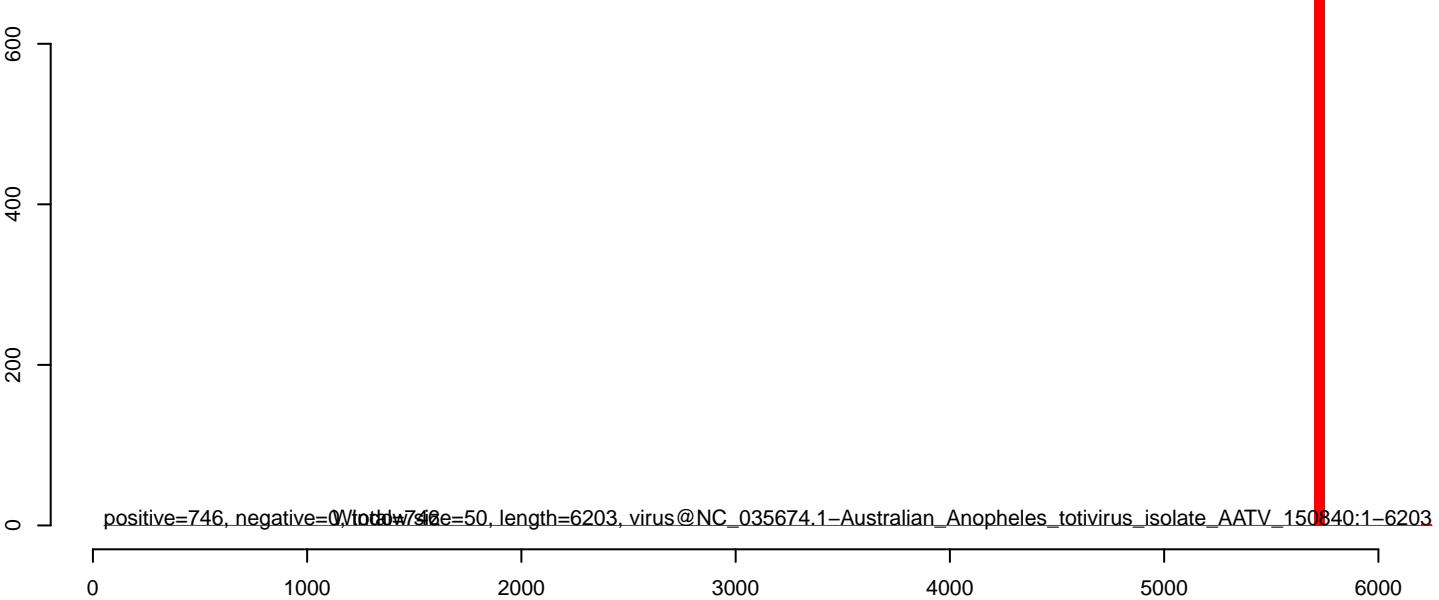
### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep



### AnGam\_Fcarc\_TN.rep



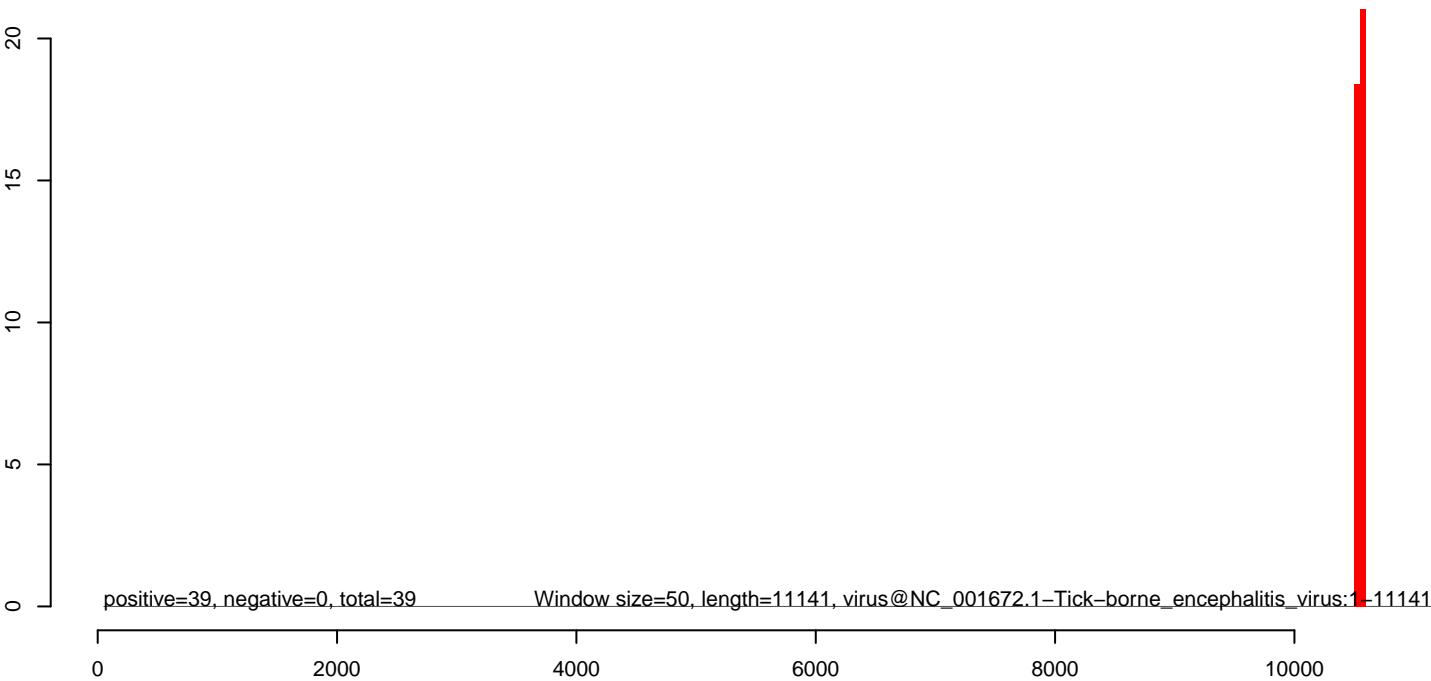
### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep



### AnGam\_Fcarc\_TN.rep



### AnGam\_Fcarc\_TN.18\_23.rep

positive=112, negative=0, total=112

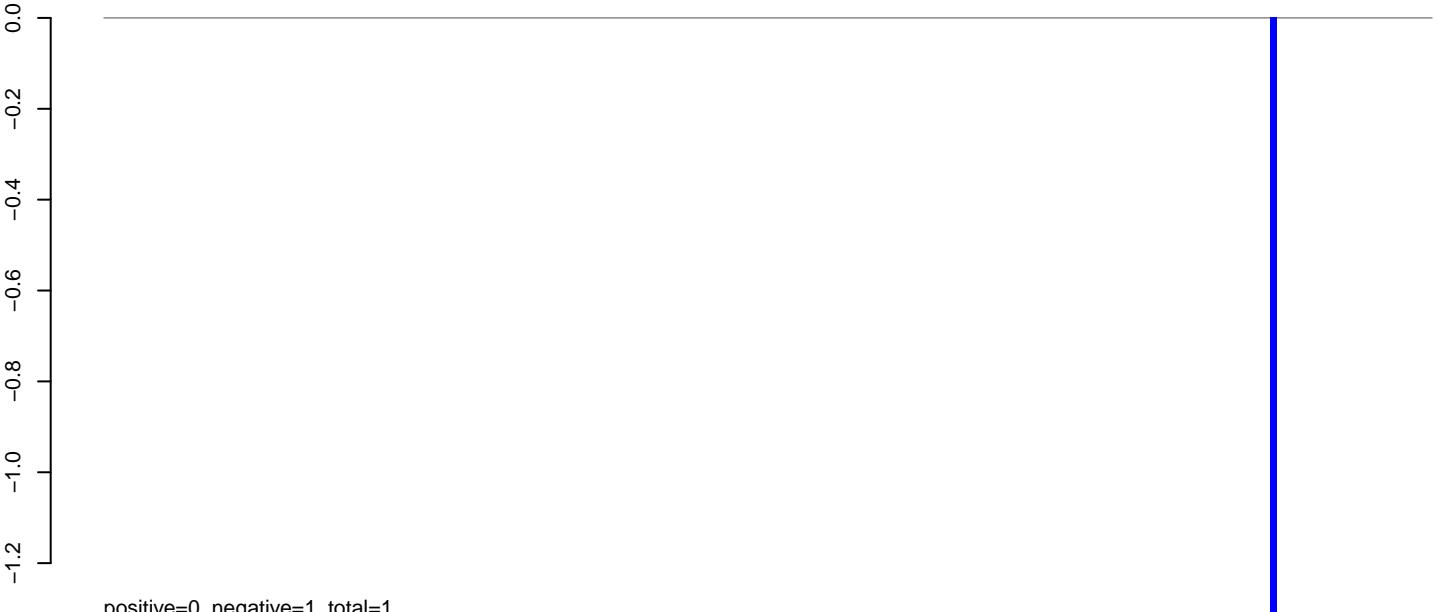
### AnGam\_Fcarc\_TN.24\_35.rep

positive=0, negative=0, total=0

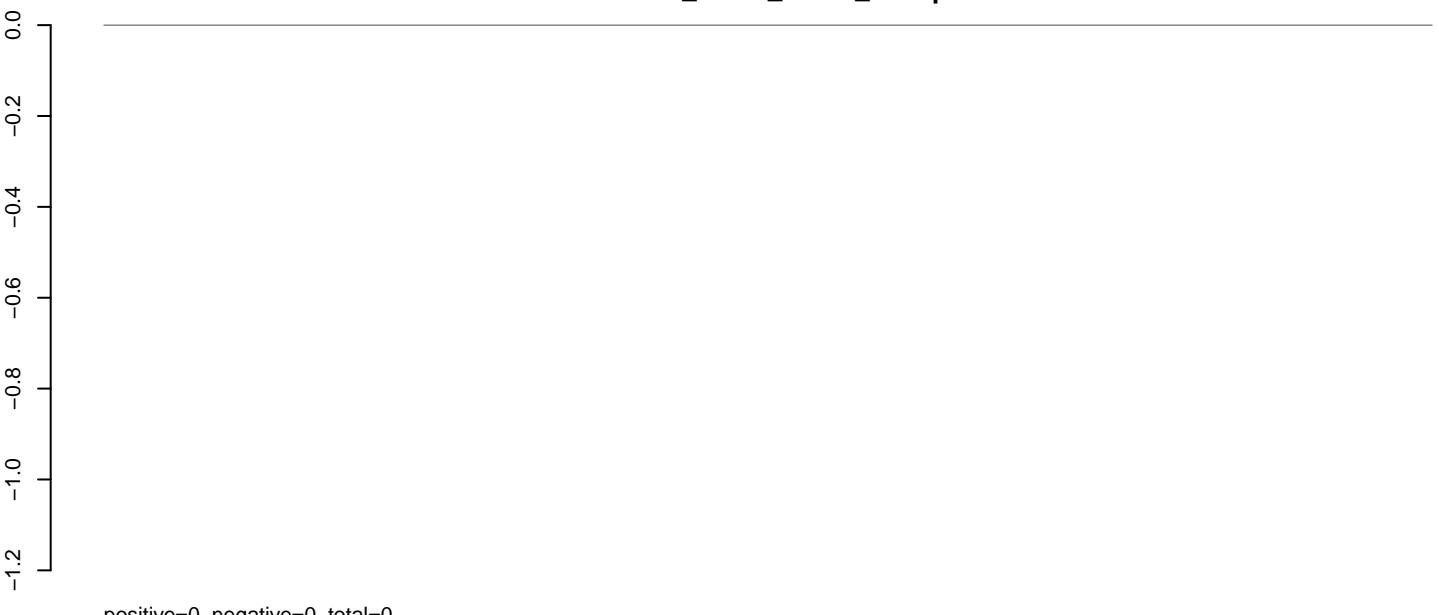
### AnGam\_Fcarc\_TN.rep

size=50 length=12196 positives@ 11147992.1 Una\_virus\_non\_structural\_polyprotein\_gene\_partial\_cds\_and\_structural\_polyprotein\_gene:1-1196

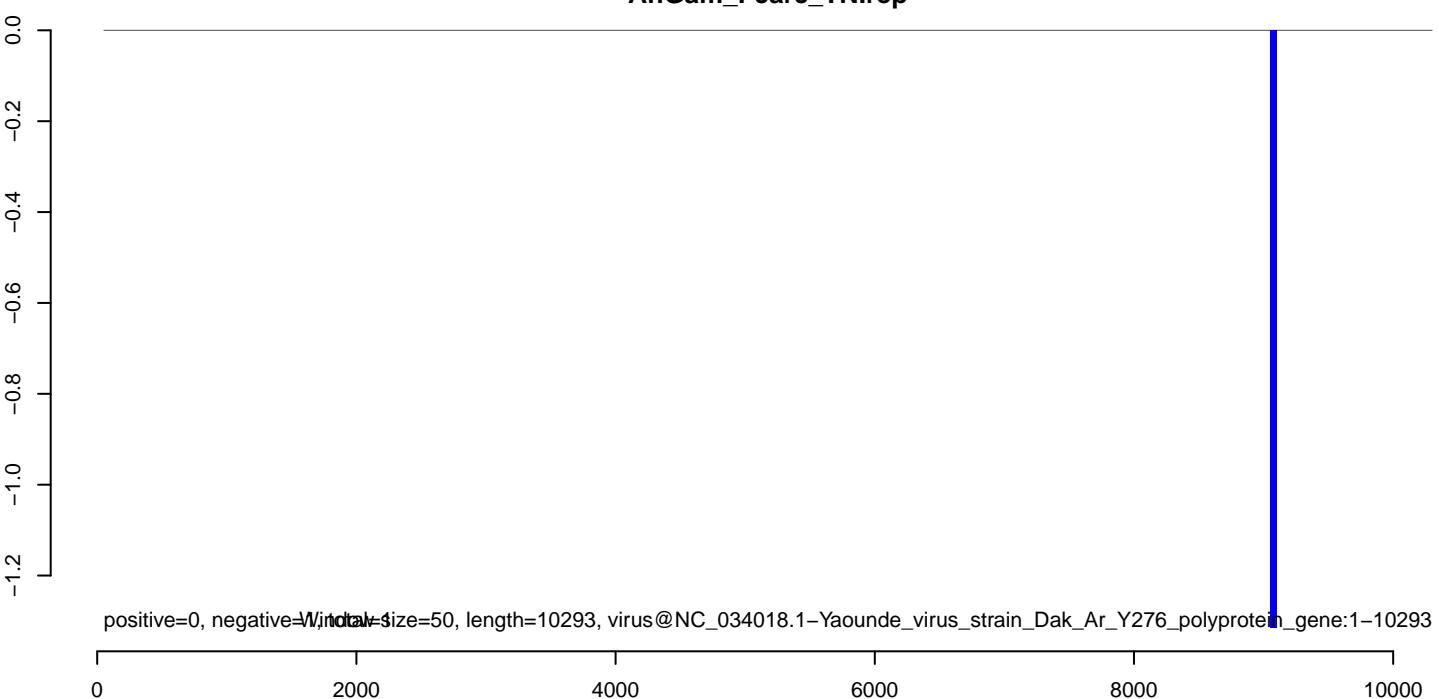
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



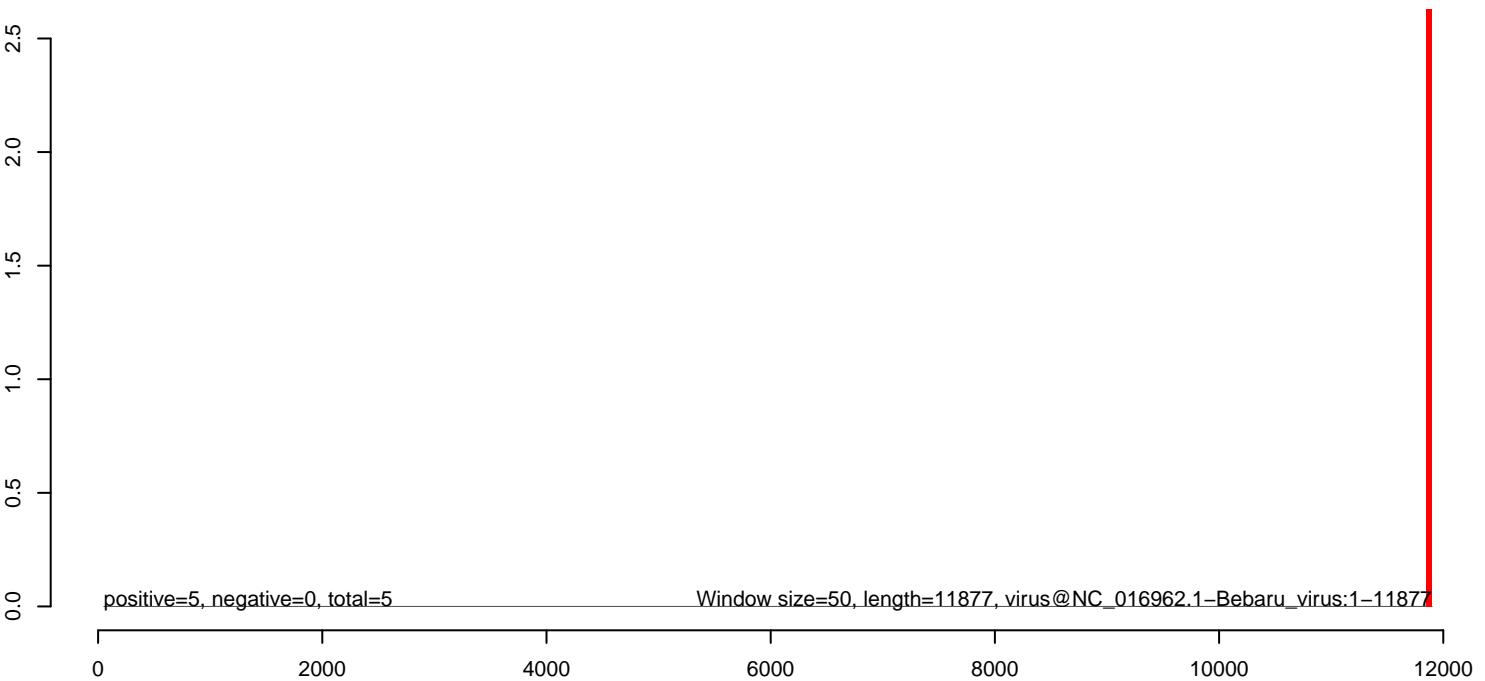
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



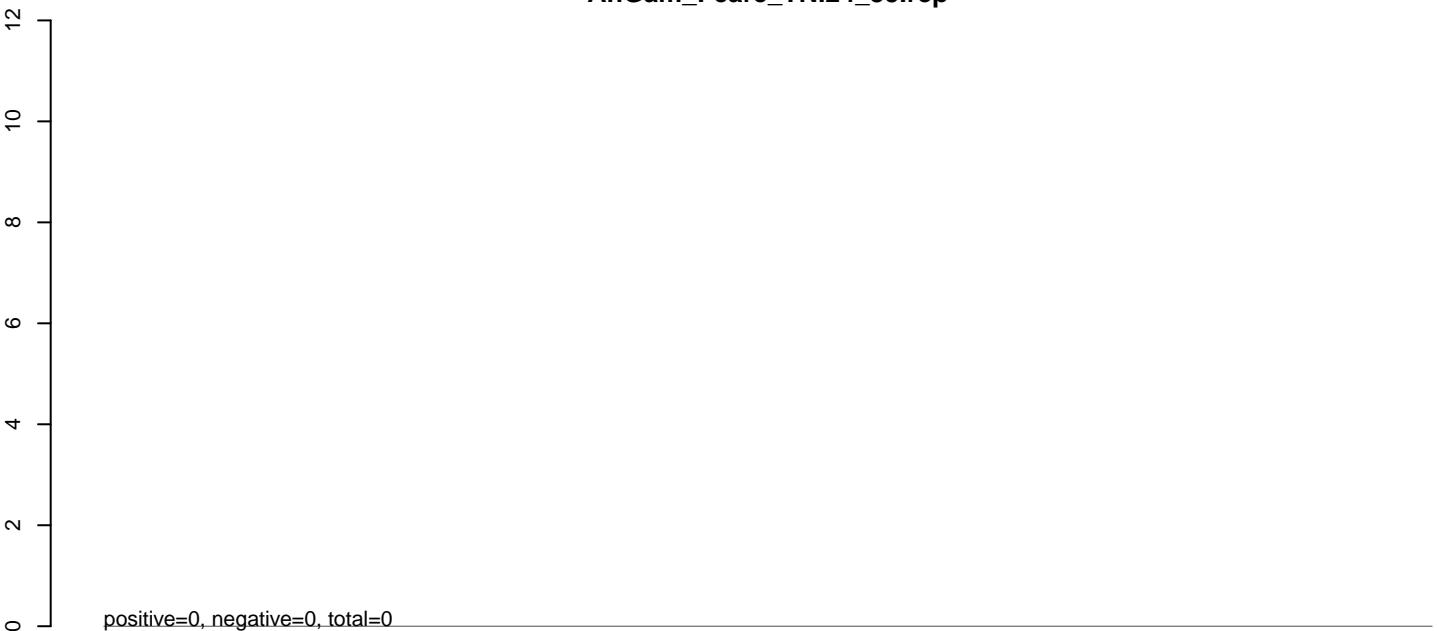
### AnGam\_Fcanc\_TN.rep



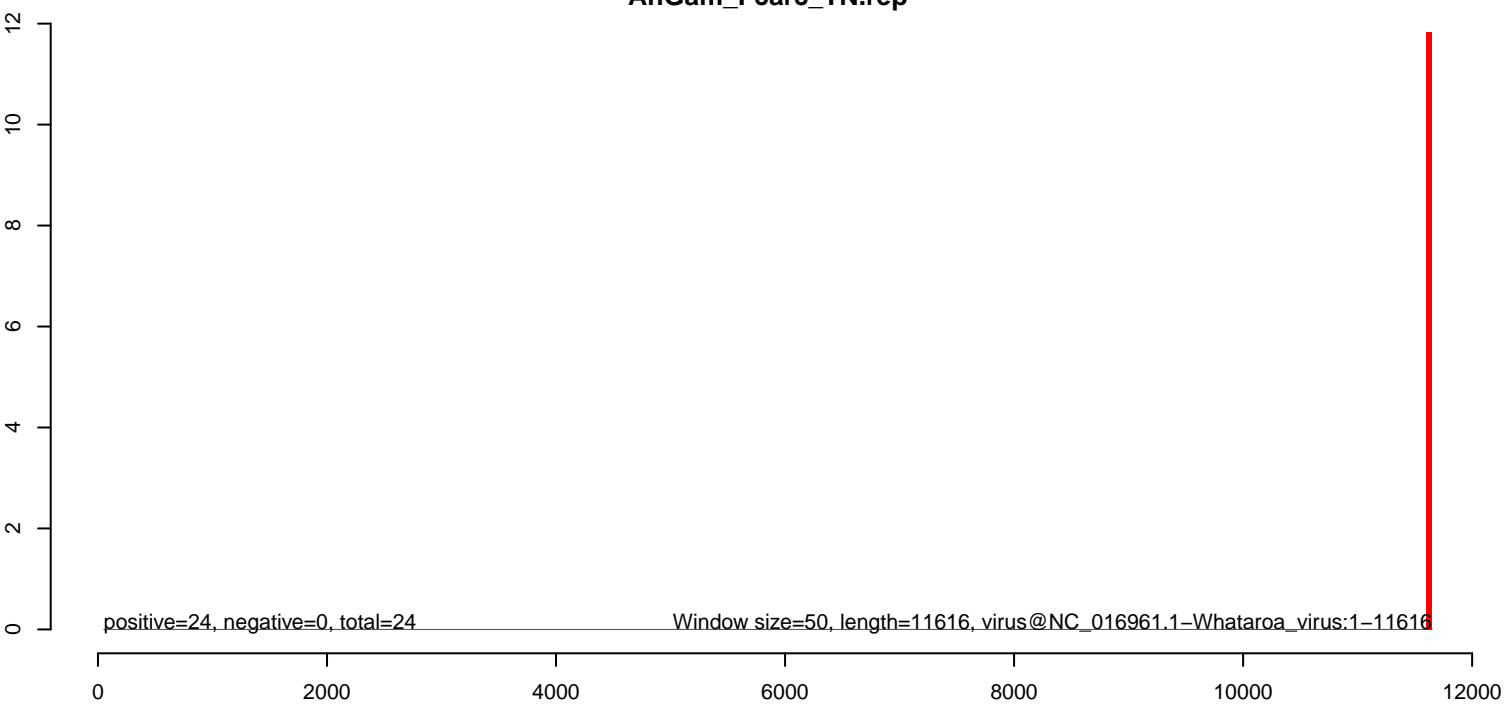
### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep



### AnGam\_Fcarc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=1, negative=0, total=1

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

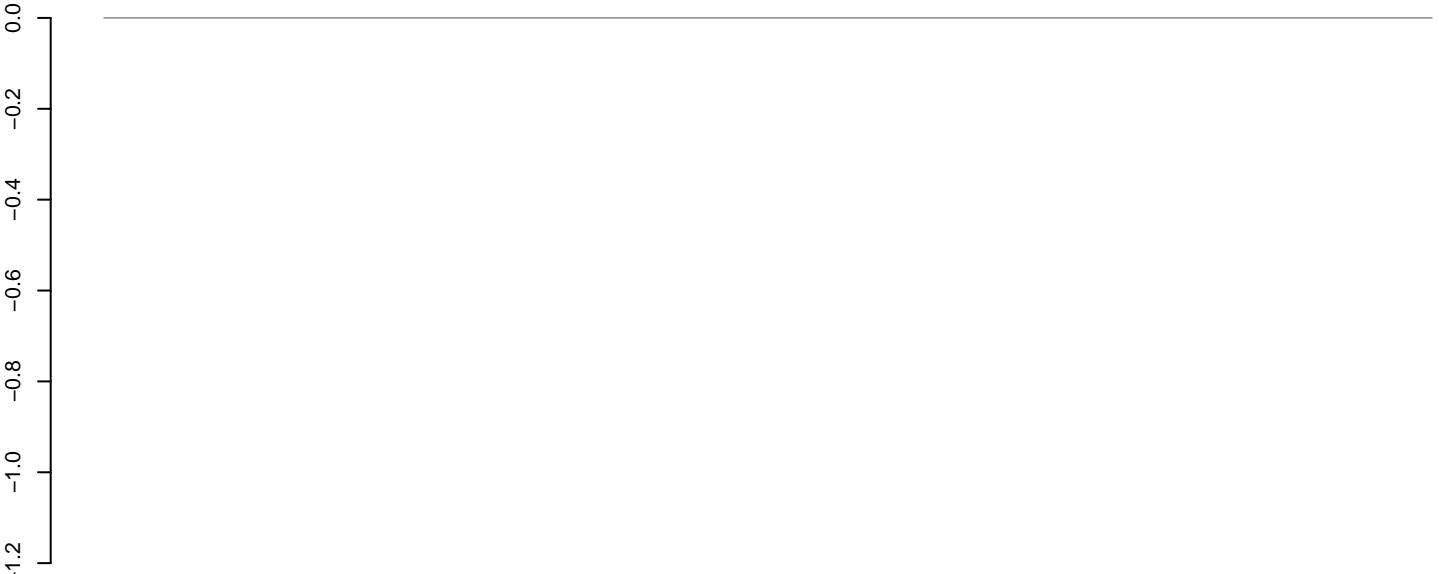
### AnGam\_Fcanc\_TN.rep

positive=1, negative=0, total=1

Window size=50, length=4806, virus@NC\_012958.1-Drosophila\_A\_virus:1-4806

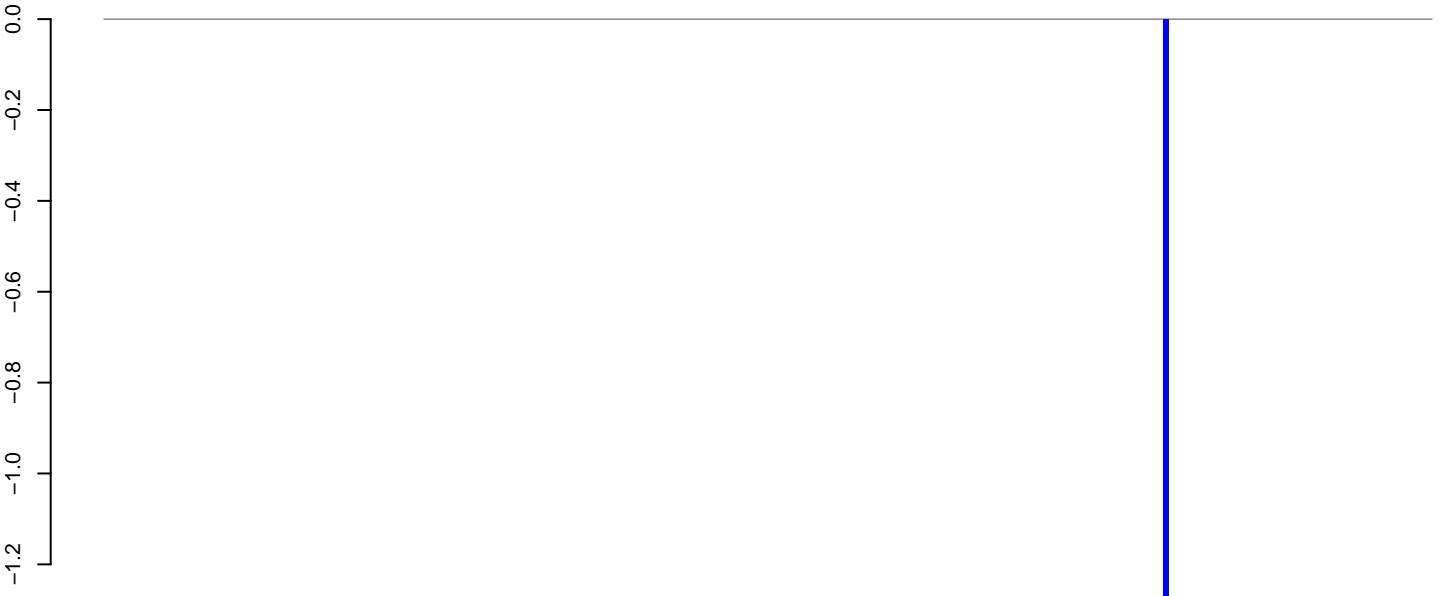
0 1000 2000 3000 4000 5000

### AnGam\_Fcanc\_TN.18\_23.rep



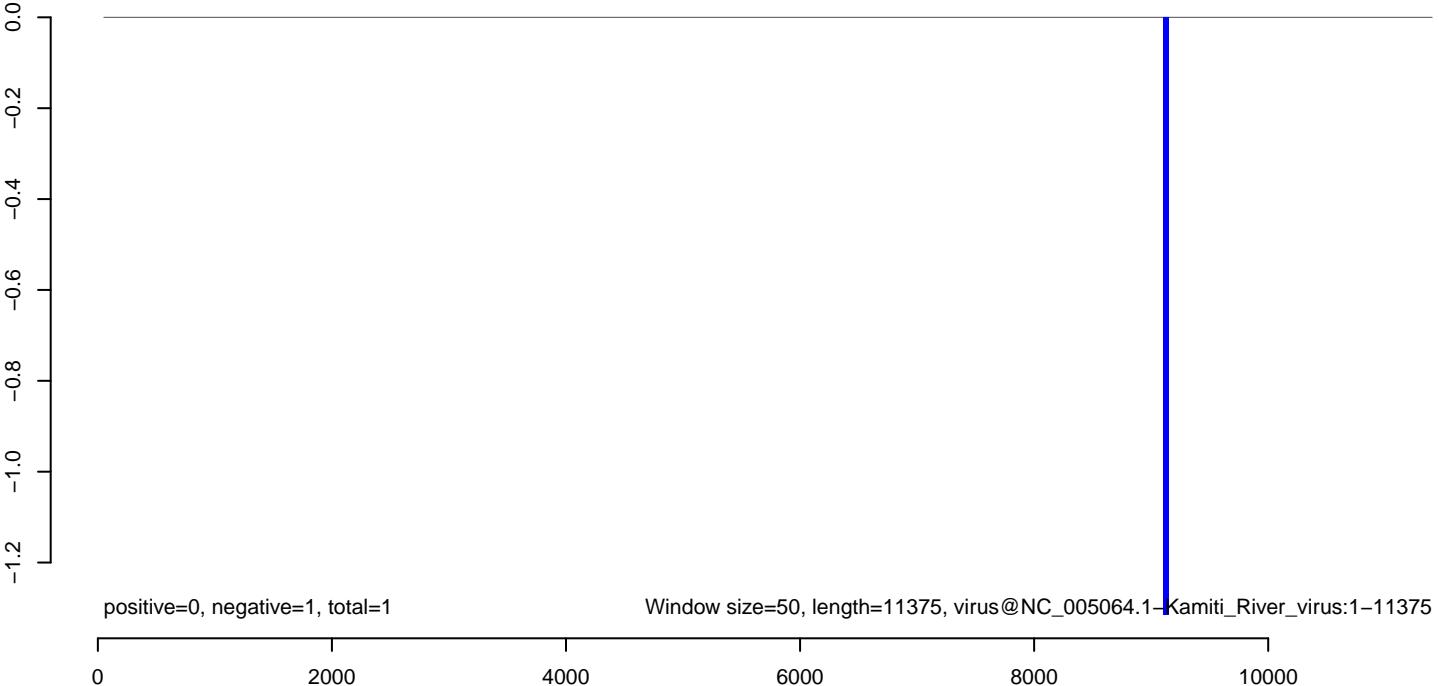
positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep



positive=0, negative=1, total=1

### AnGam\_Fcanc\_TN.rep



positive=0, negative=1, total=1

Window size=50, length=11375, virus@NC\_005064.1-Kamiti\_River\_virus:1-11375

0 2000 4000 6000 8000 10000

### AnGam\_Fcarc\_TN.18\_23.rep



positive=0, negative=1, total=1

### AnGam\_Fcarc\_TN.24\_35.rep



positive=0, negative=0, total=0

### AnGam\_Fcarc\_TN.rep



positive=0, negative=1, total=1, window size=50, length=6603, virus@NC\_004177.1-Drosophila\_x\_virus\_segment\_A\_and\_segment\_B:1-6603

0 1000 2000 3000 4000 5000 6000

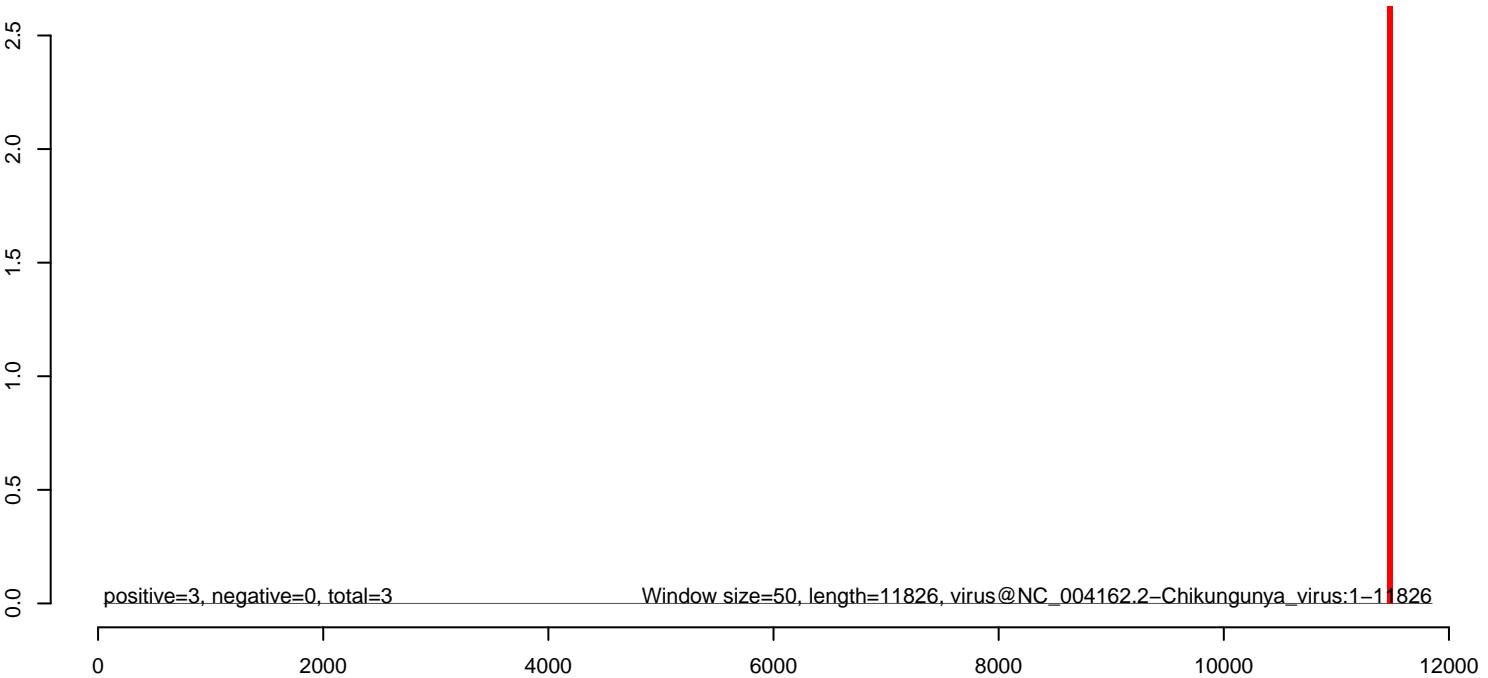
### AnGam\_Fcarc\_TN.18\_23.rep



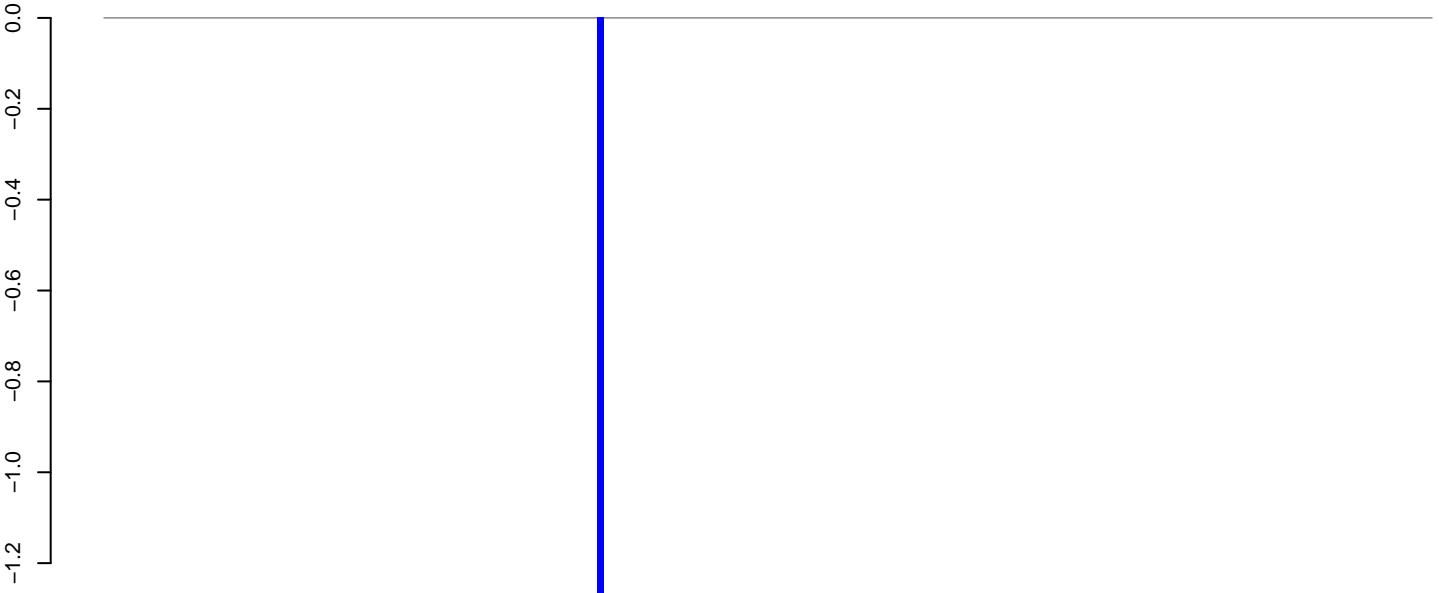
### AnGam\_Fcarc\_TN.24\_35.rep



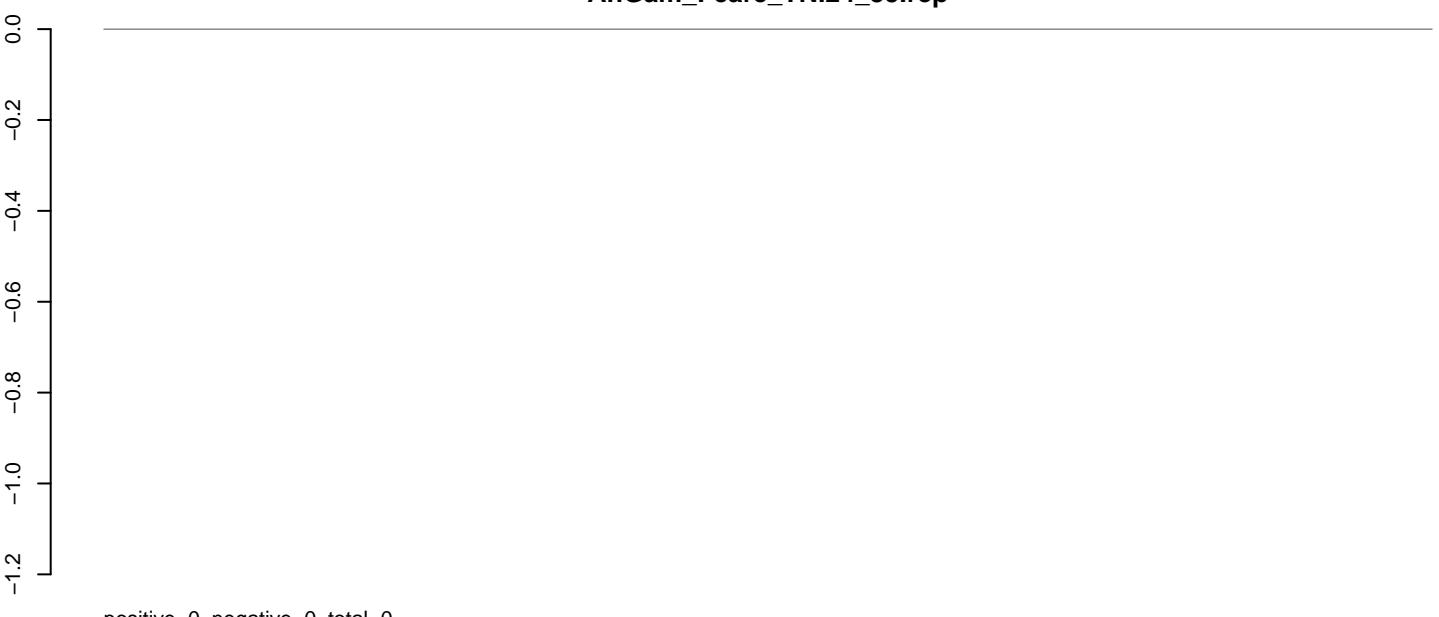
### AnGam\_Fcarc\_TN.rep



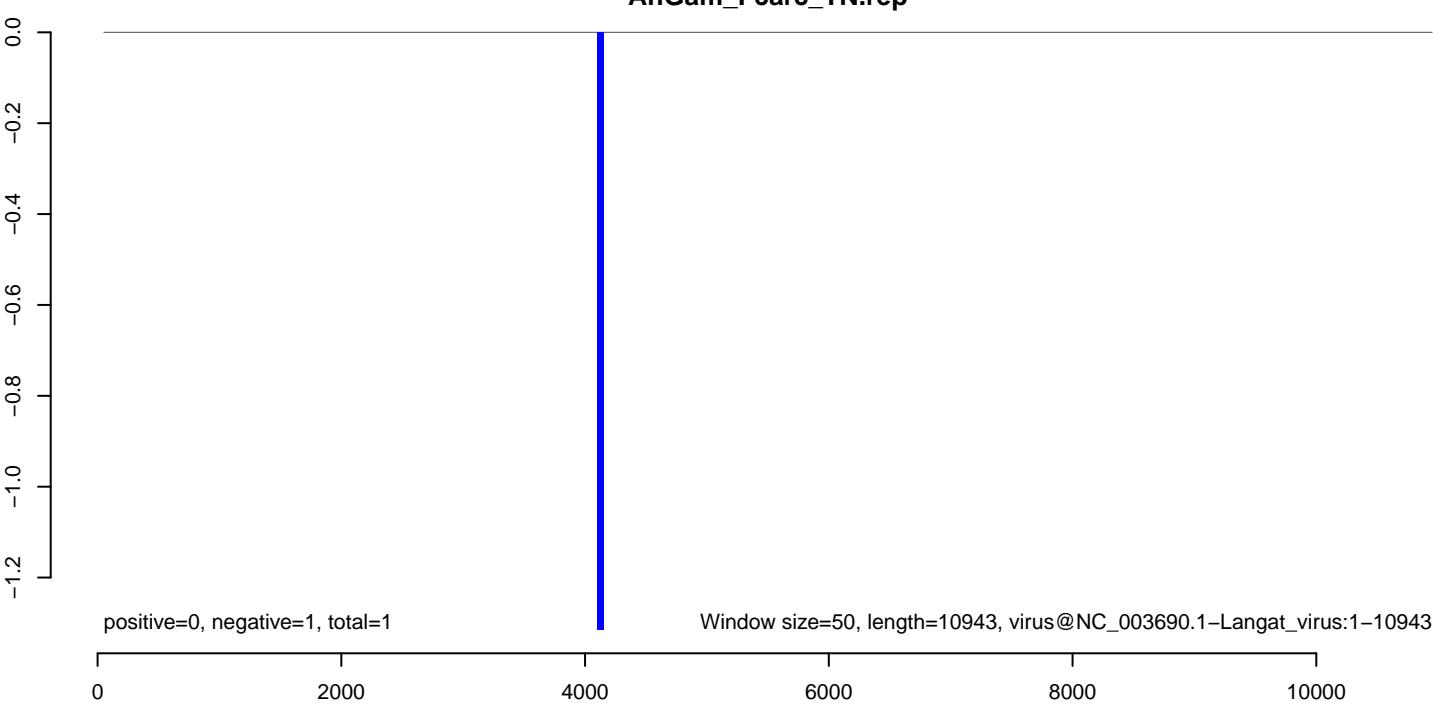
### AnGam\_Fcanc\_TN.18\_23.rep



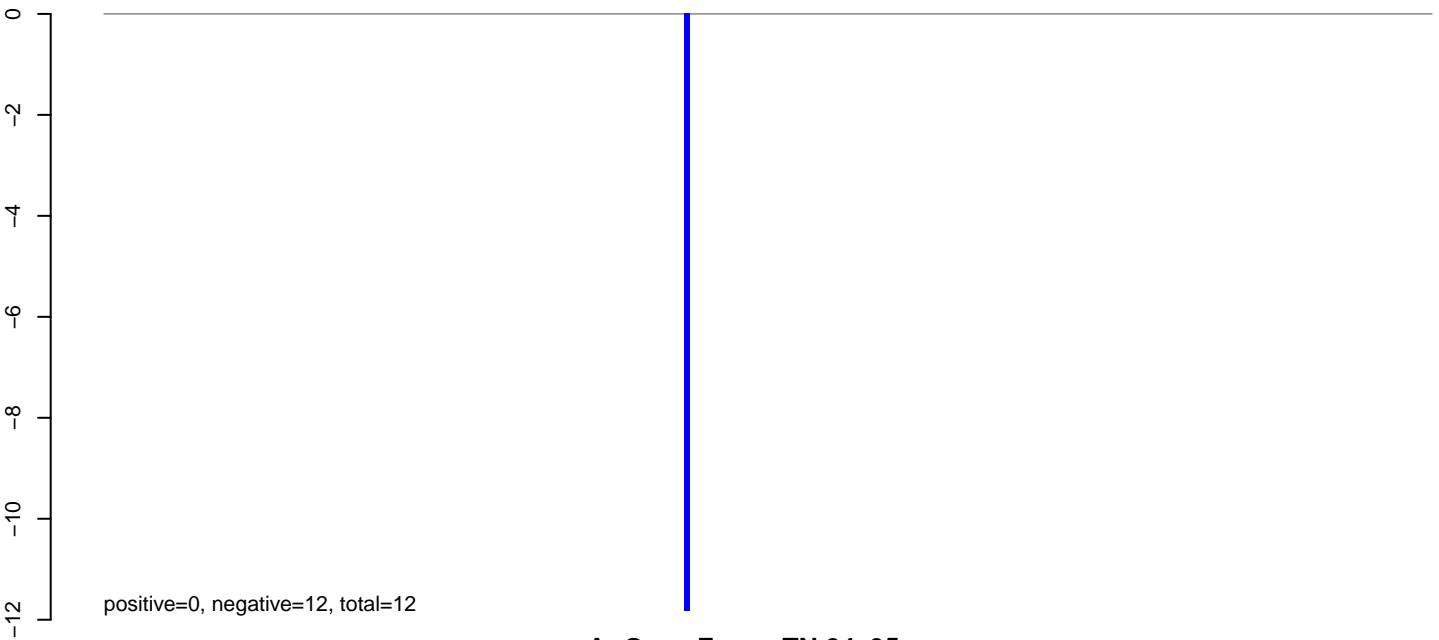
### AnGam\_Fcanc\_TN.24\_35.rep



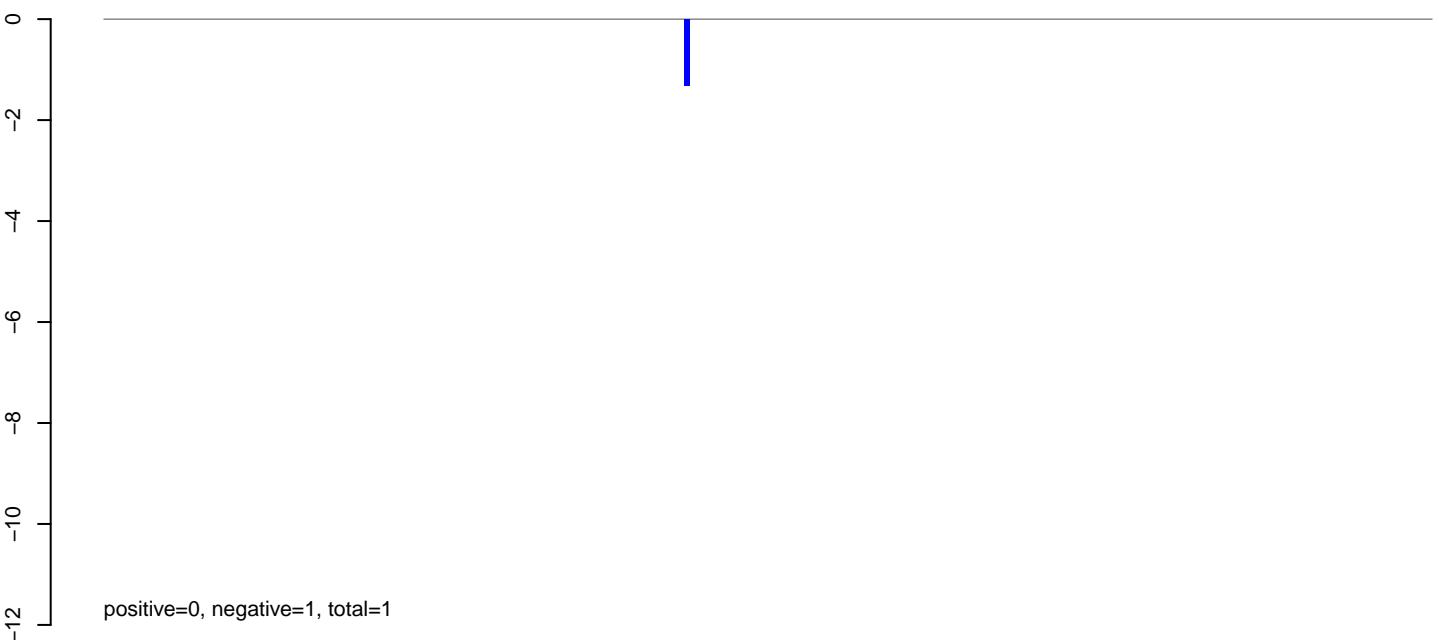
### AnGam\_Fcanc\_TN.rep



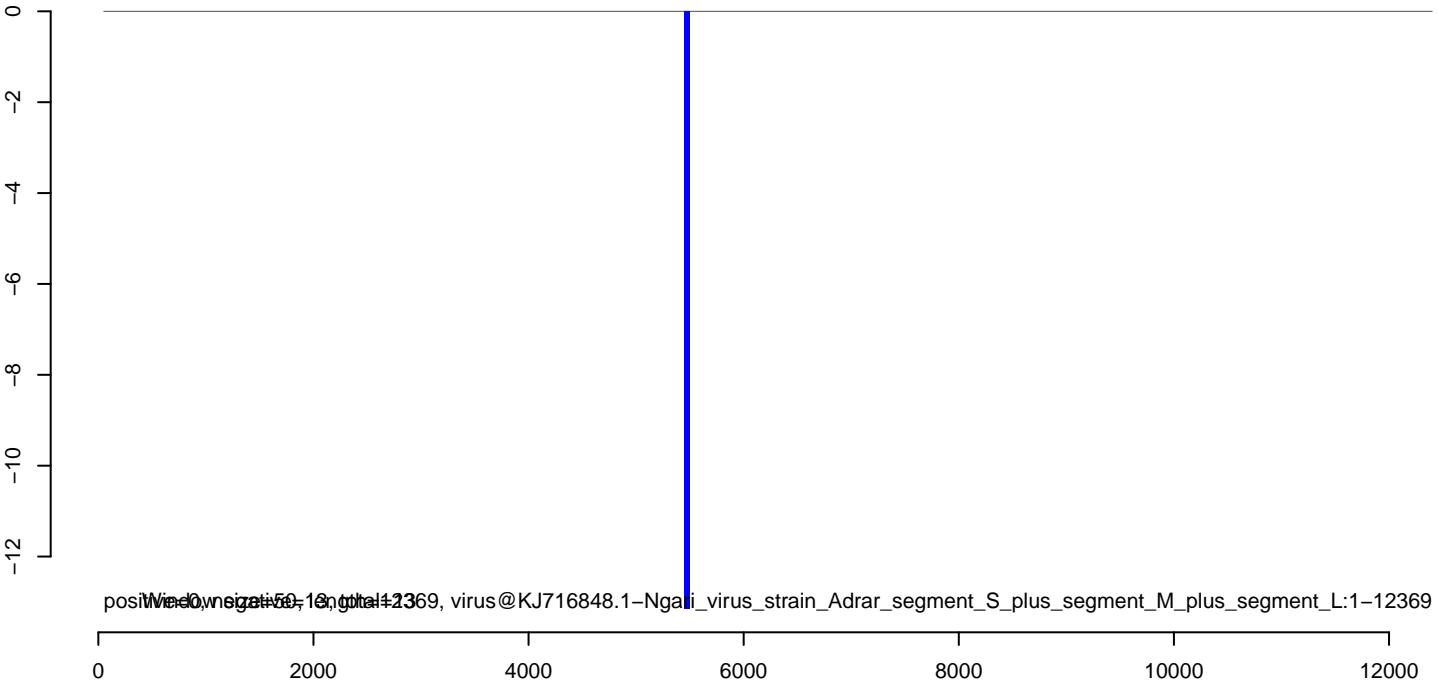
### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep



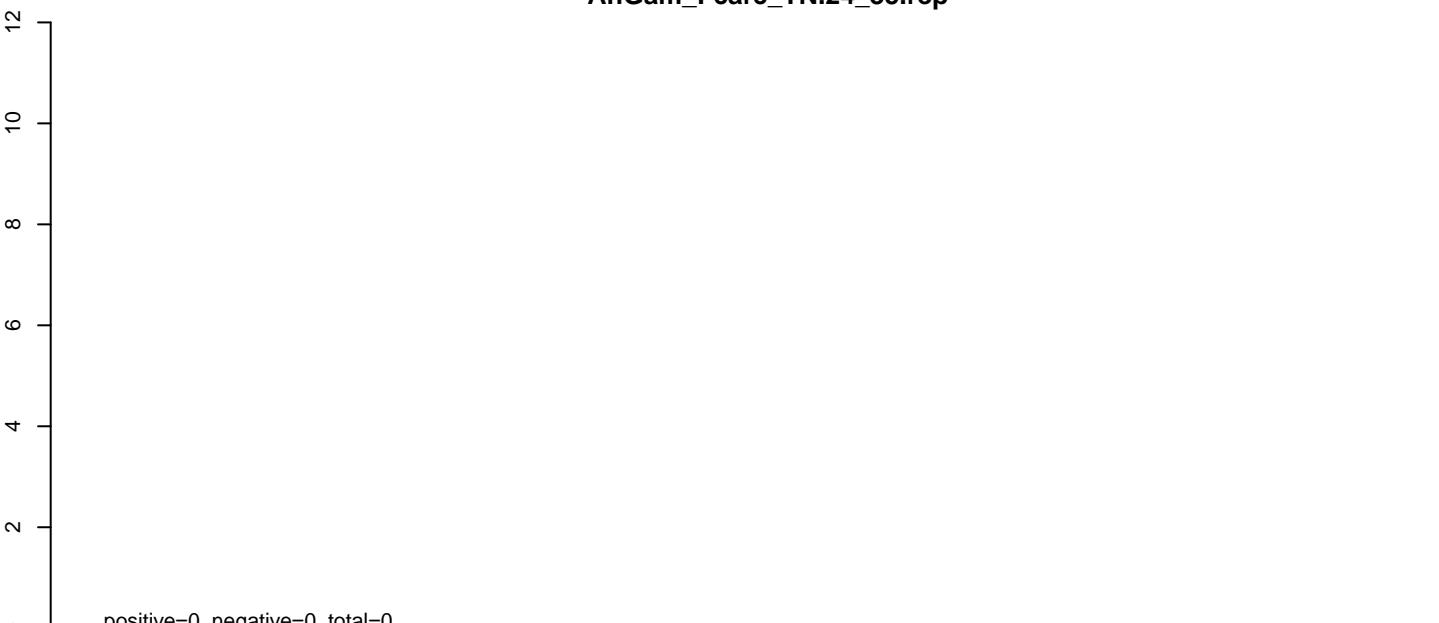
### AnGam\_Fcarc\_TN.rep



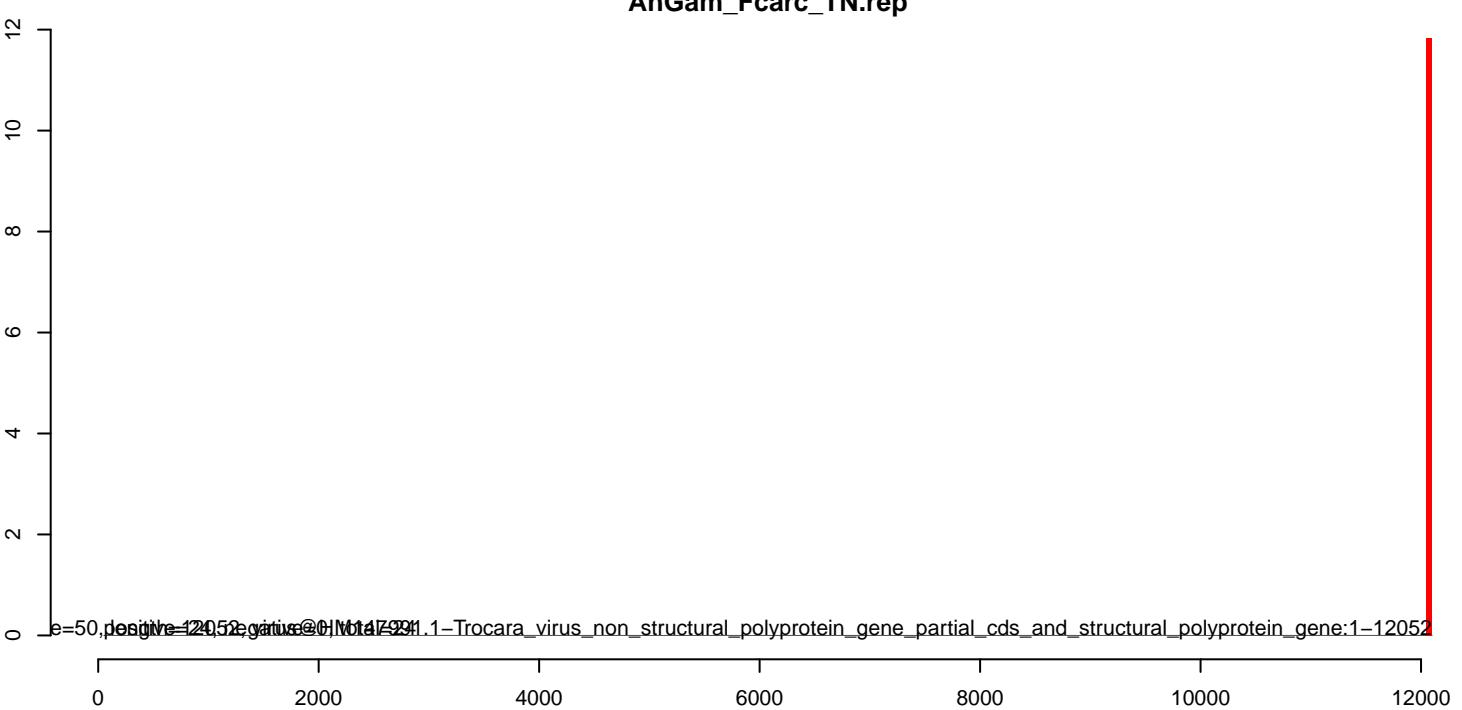
### AnGam\_Fcanc\_TN.18\_23.rep



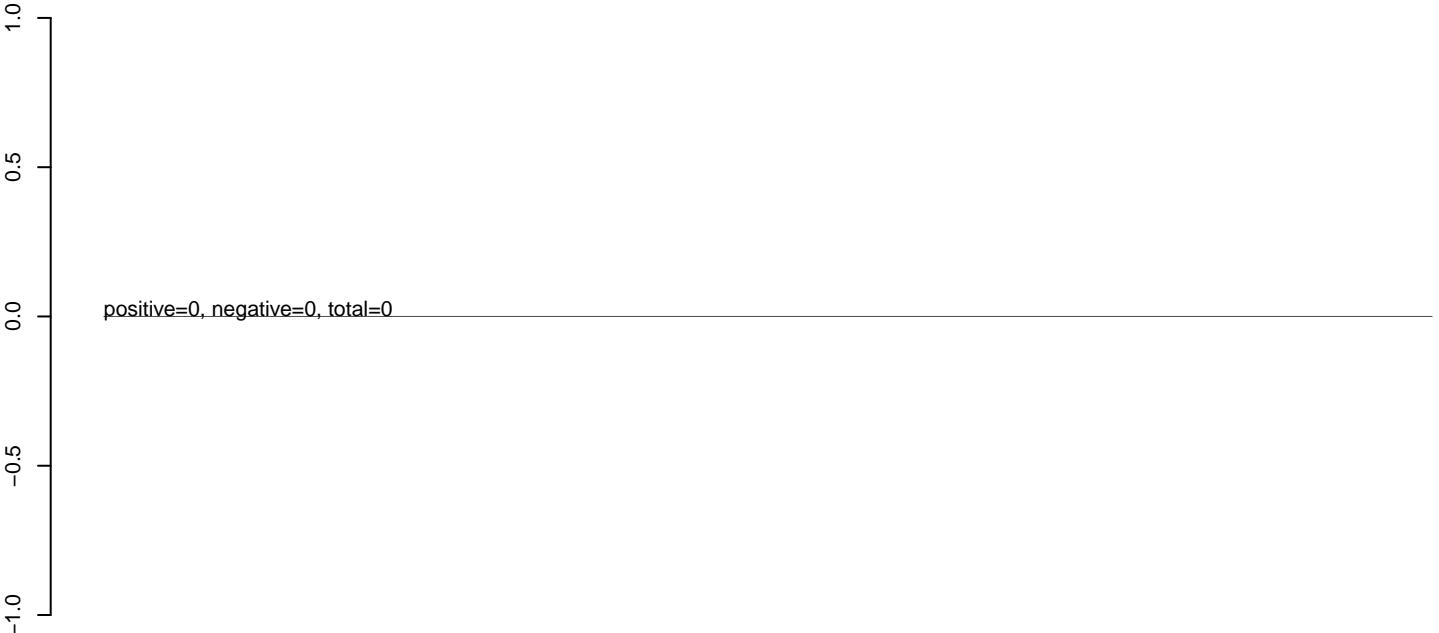
### AnGam\_Fcanc\_TN.24\_35.rep



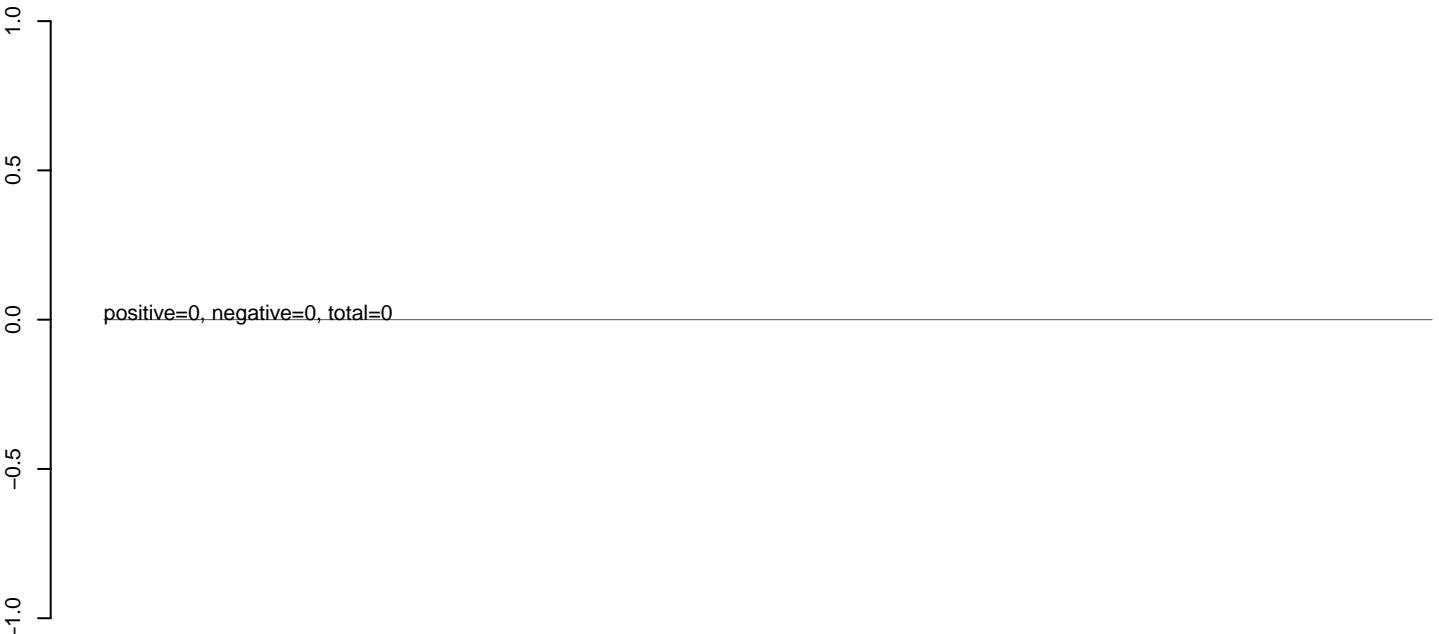
### AnGam\_Fcanc\_TN.rep



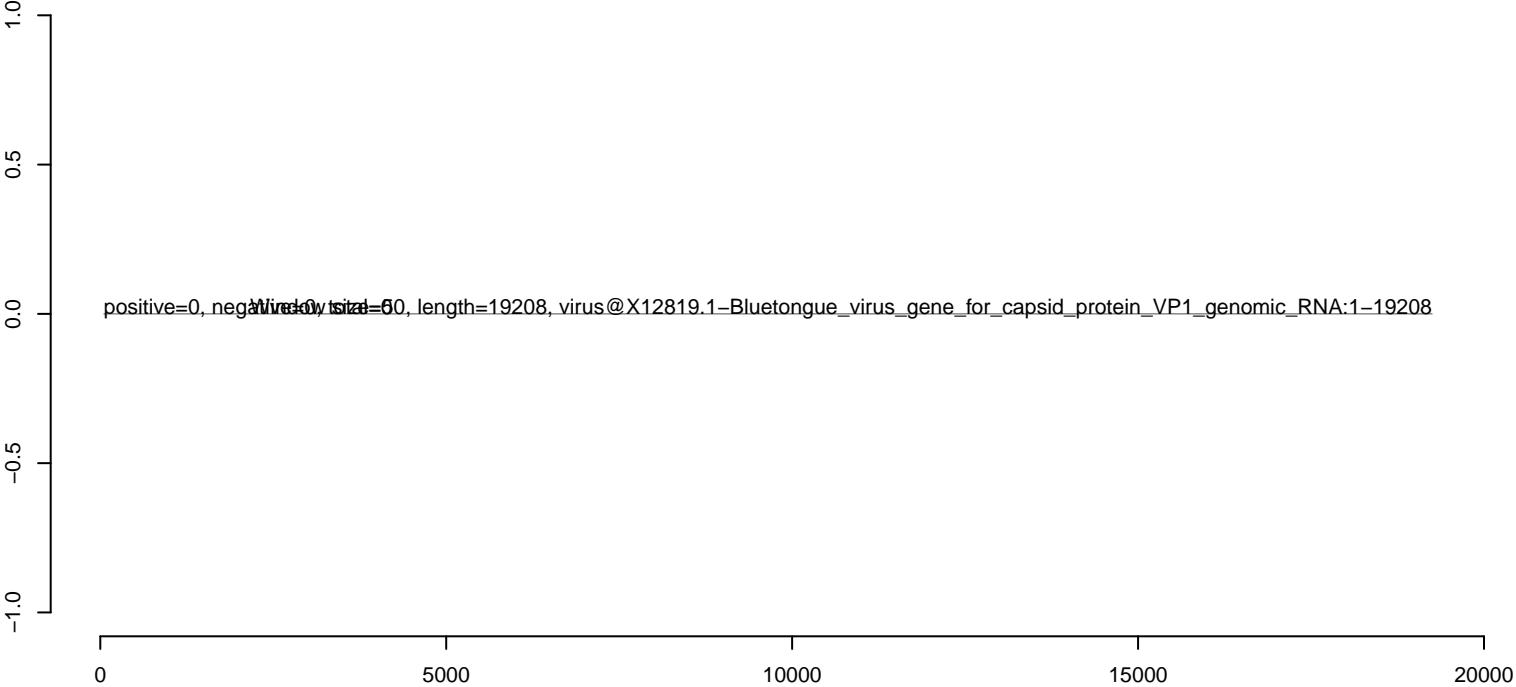
### AnGam\_Fcanc\_TN.18\_23.rep



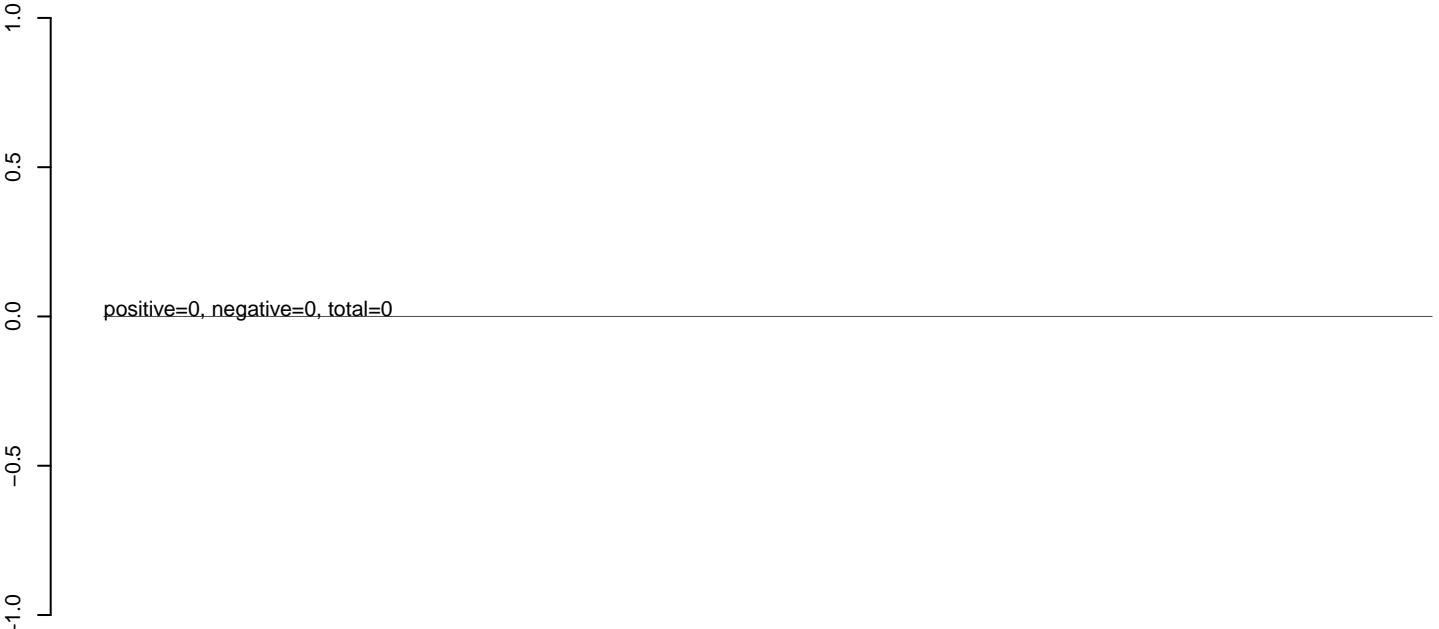
### AnGam\_Fcanc\_TN.24\_35.rep



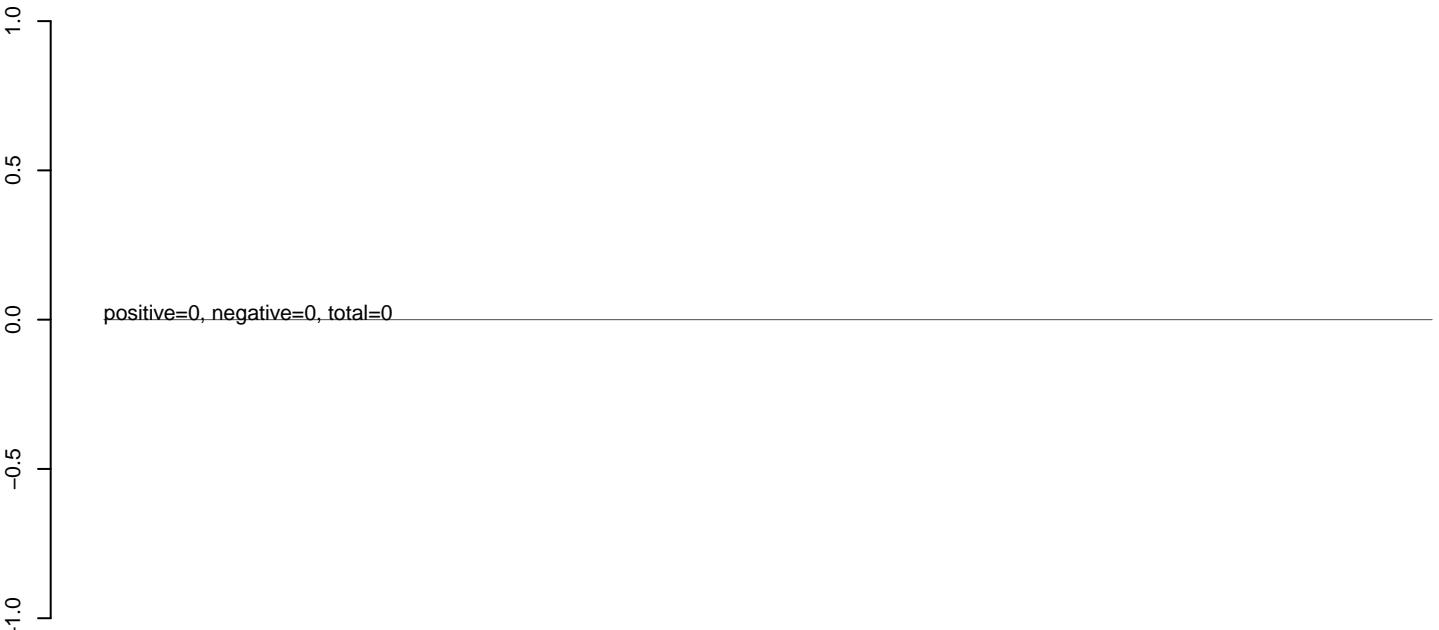
### AnGam\_Fcanc\_TN.rep



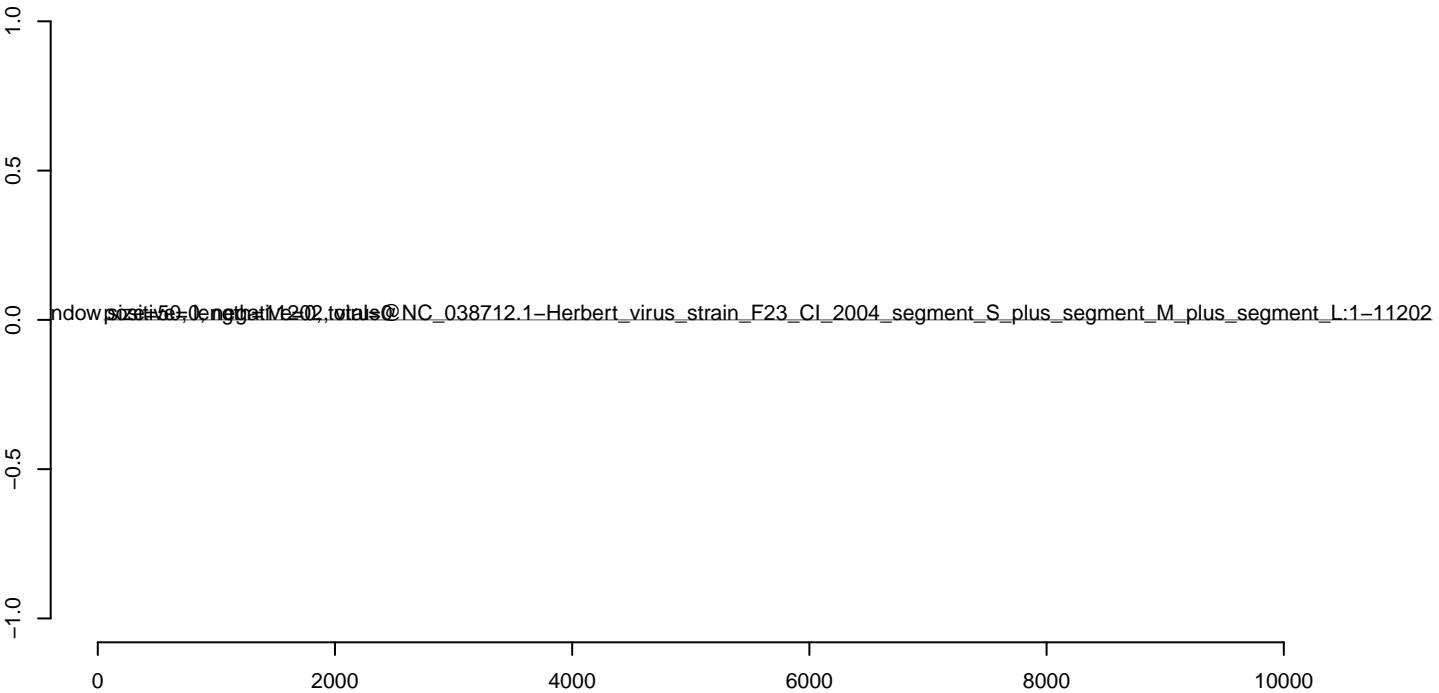
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.rep

window positive=0, negative=0, total=0 NC\_038707.1-Jonchet\_virus\_strain\_B81-CI-2004\_segment\_S\_plus\_segment\_M\_plus\_segment\_L:1-14098



positive=0, negative=0, total=0



positive=0, negative=0, total=0





positive=0, negative=0, total=0



positive=0, negative=0, total=0



**C** - oprotein positive hypothetical proteins chaperone protein hypothetical proteins matrix glycoprotein hypothetical protein and polymerase genes:1-13189

0



positive=0, negative=0, total=0

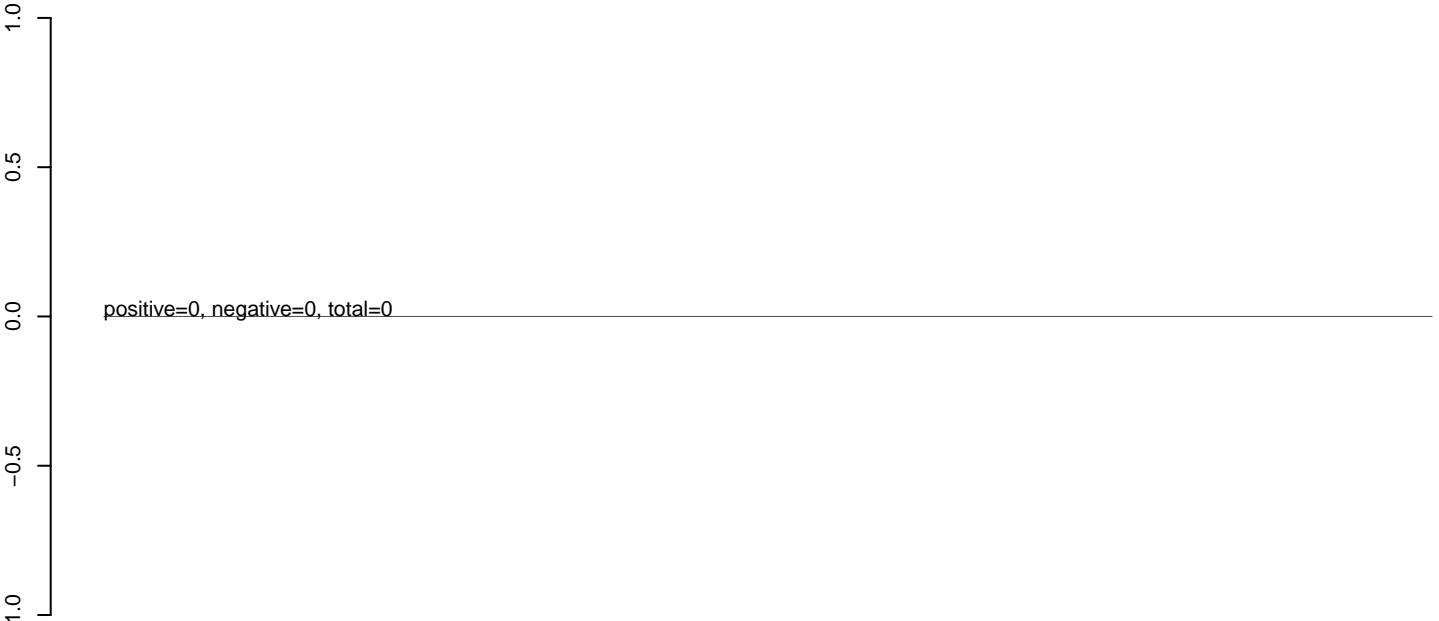


positive=0, negative=0, total=0

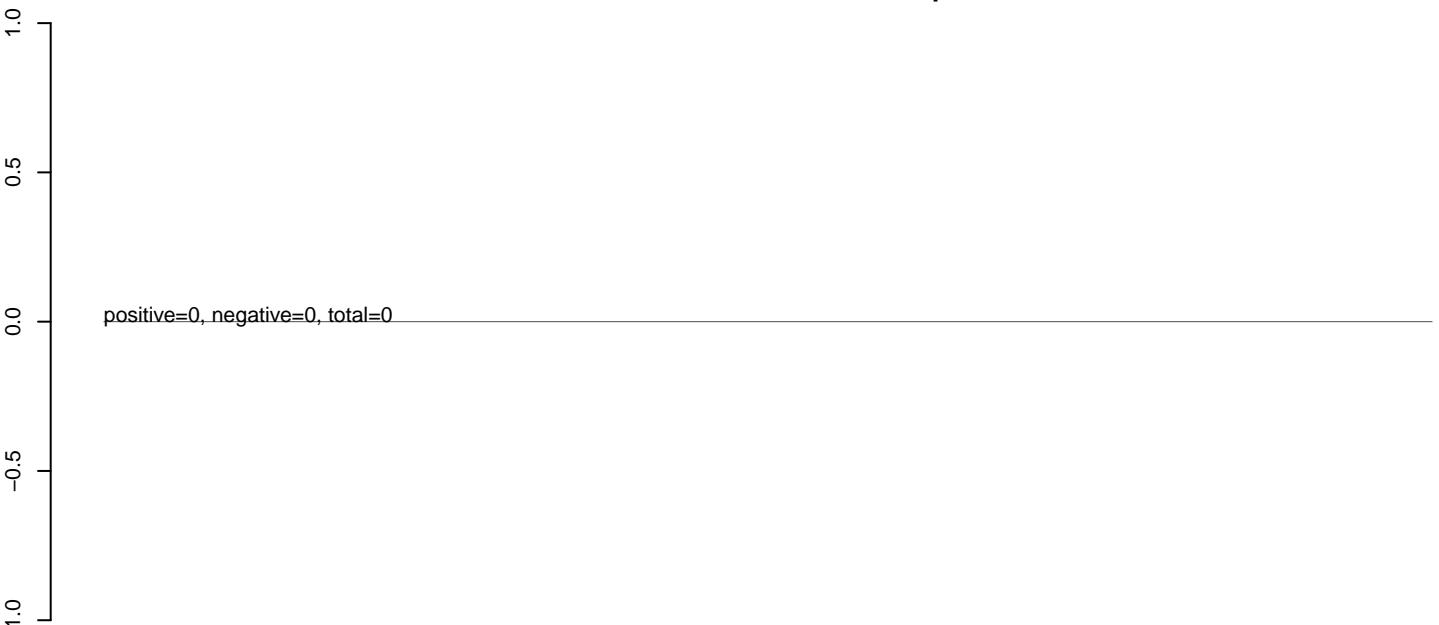


positive=0 mutations=distribution of phosphoprotein hypothetical proteins matrix glycoprotein hypothetical protein and polymerase genes:1-12403

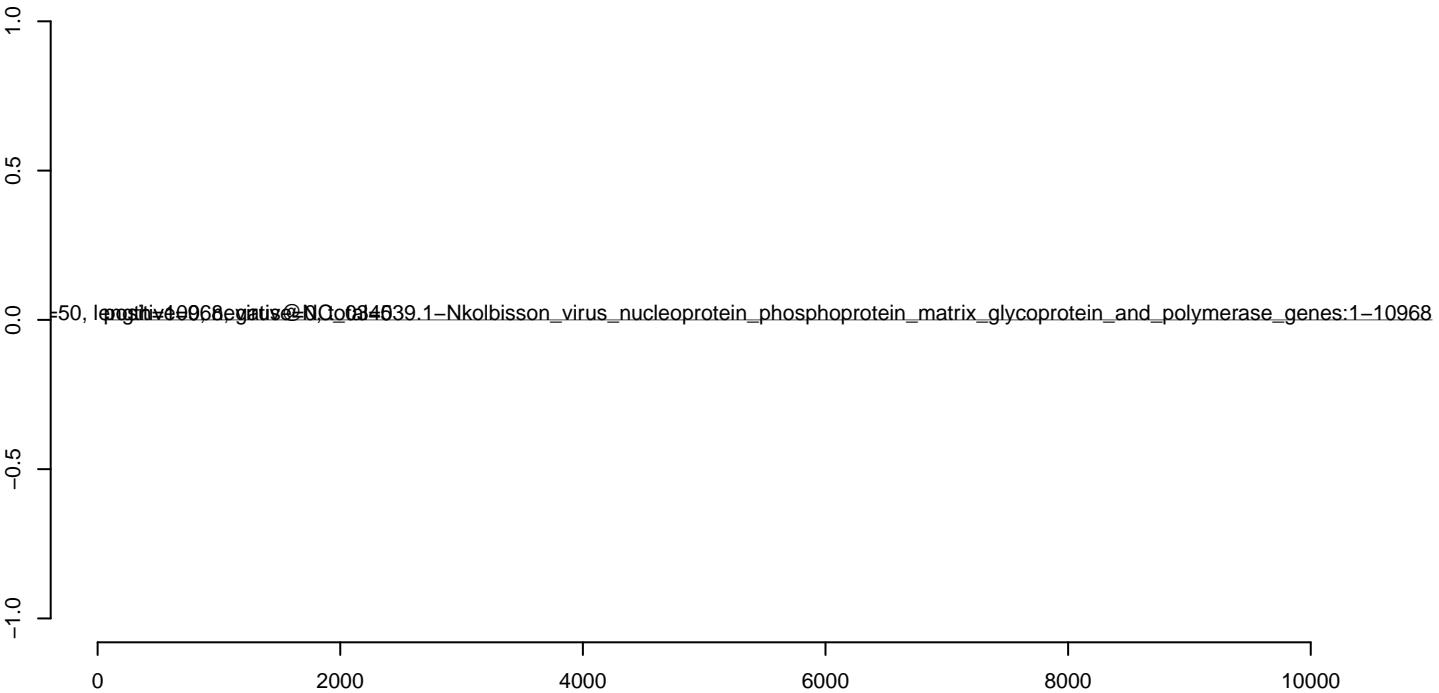
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep





positive=0, negative=0, total=0



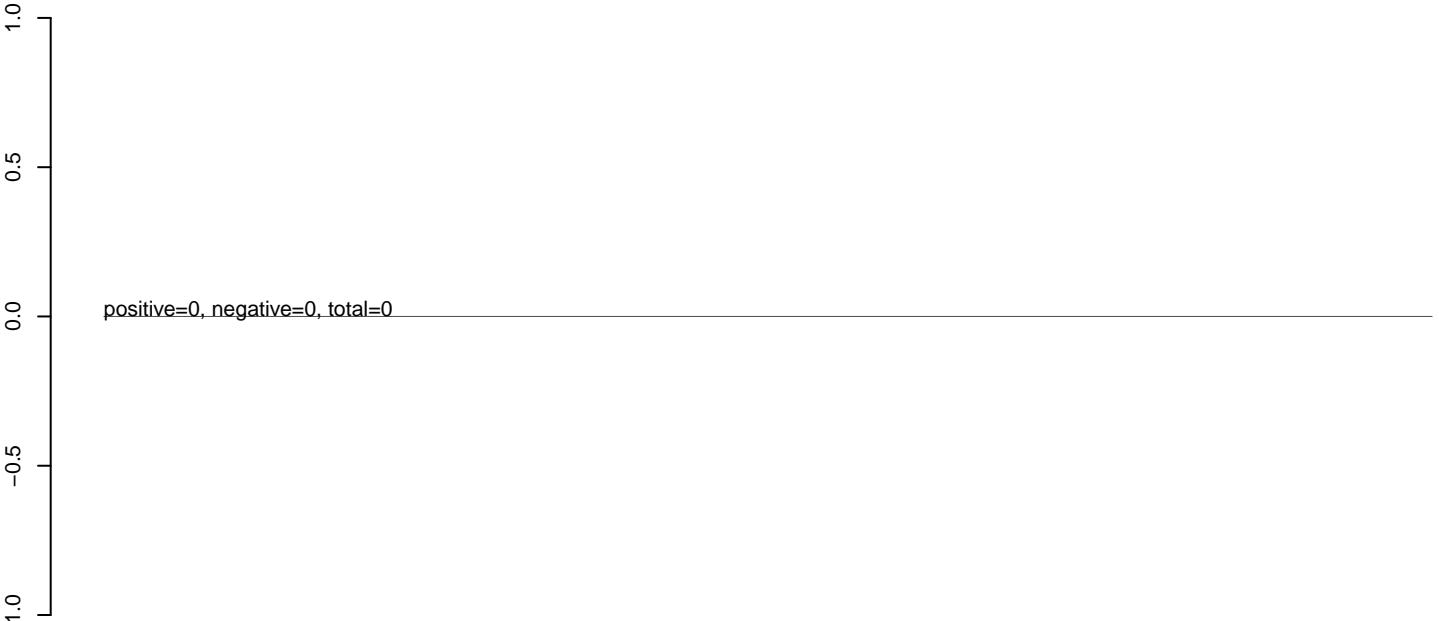
positive=0, negative=0, total=0



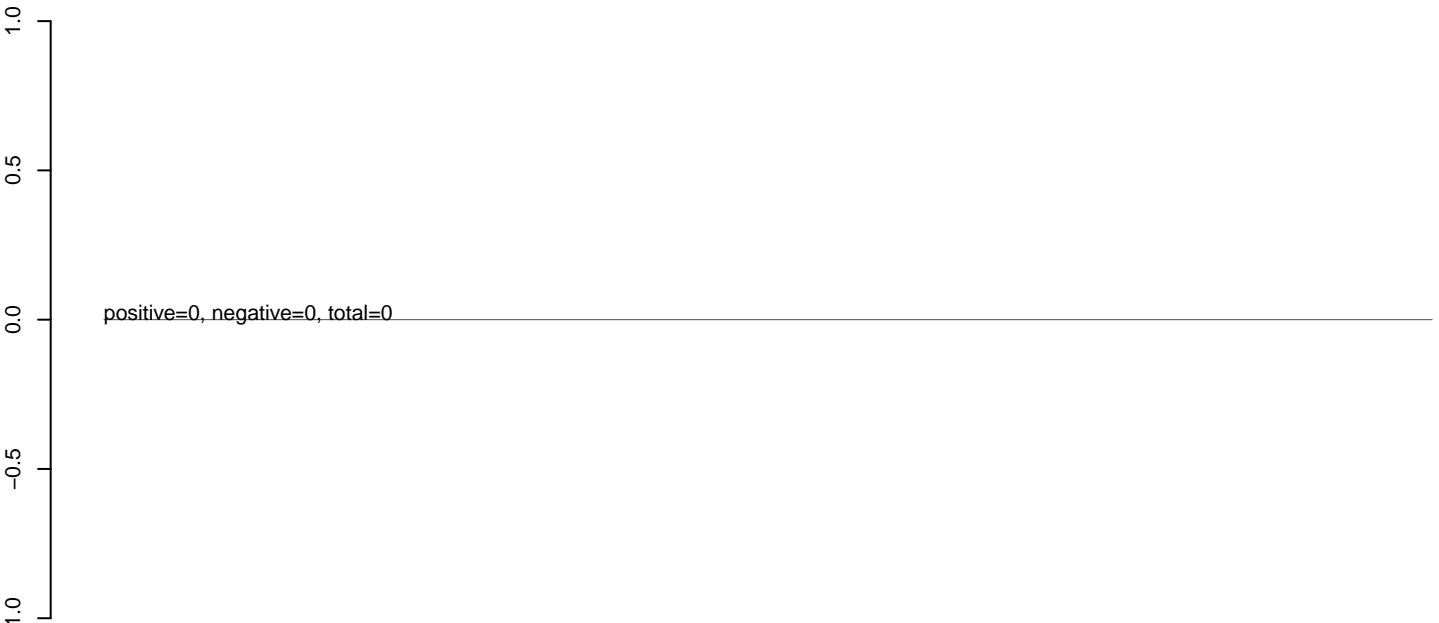
kalosivirus+0 mutations in the phosphoprotein, hypothetical protein, matrix, glycoprotein, hypothetical proteins and polymerase genes: 1-13155

பாகு\_பாகு, பாகுபாகு, பாகு

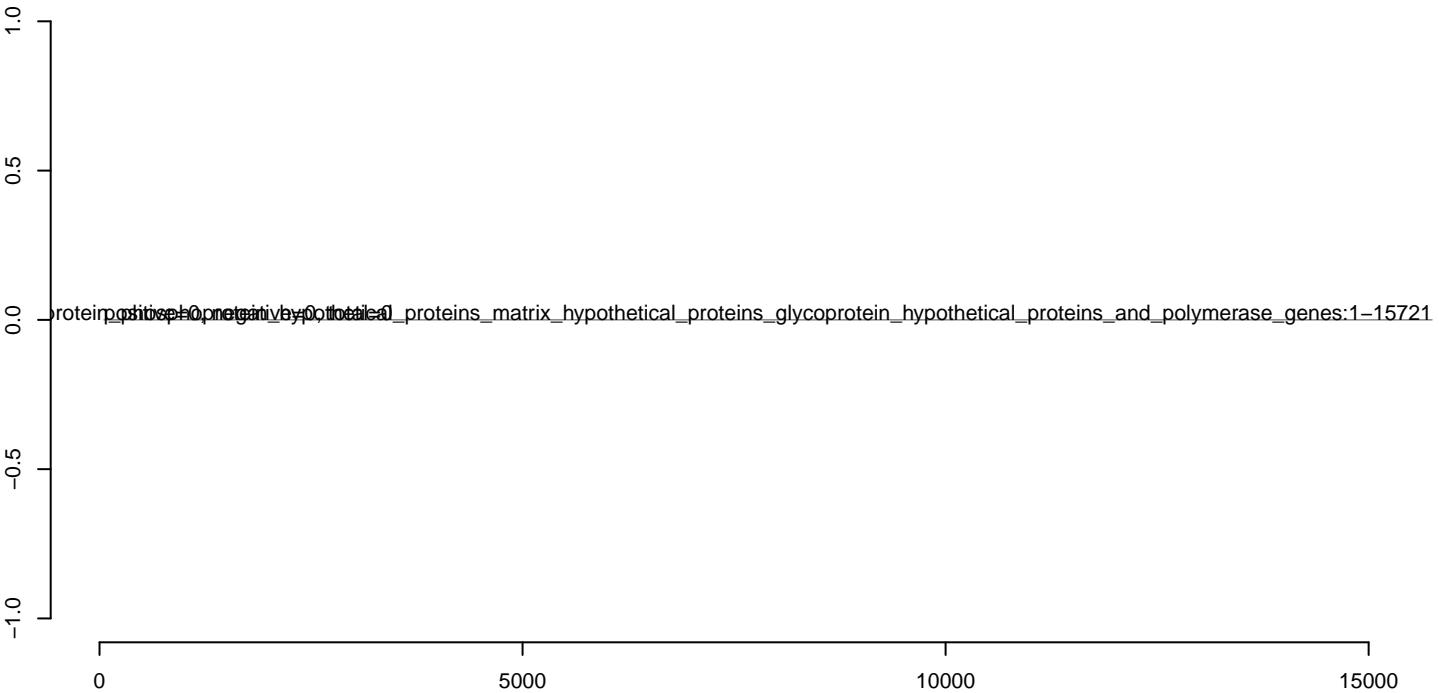
### AnGam\_Fcanc\_TN.18\_23.rep



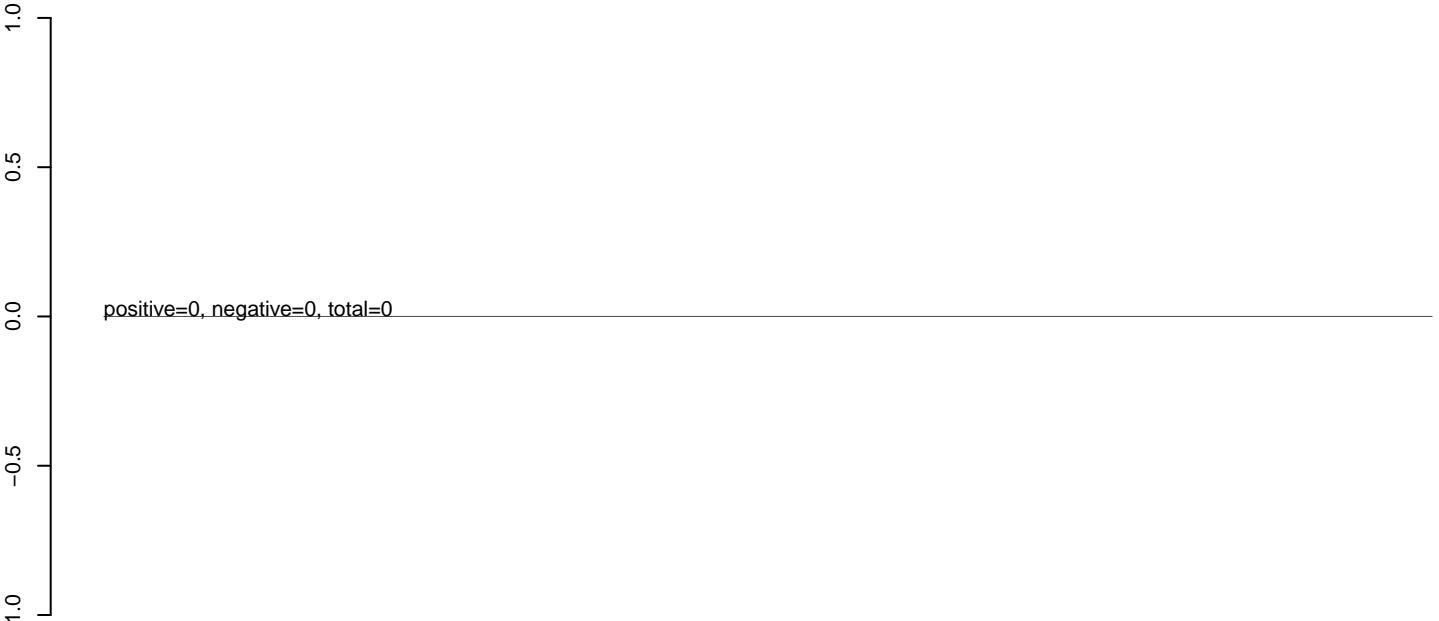
### AnGam\_Fcanc\_TN.24\_35.rep



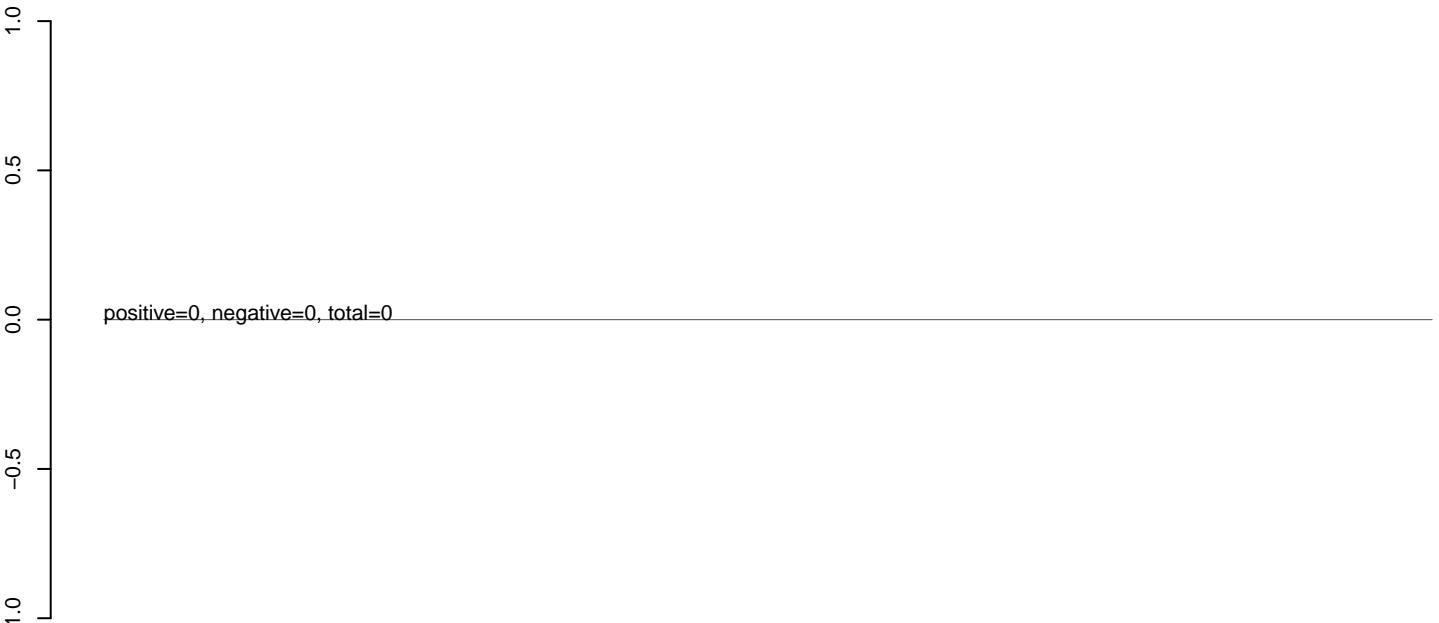
### AnGam\_Fcanc\_TN.rep



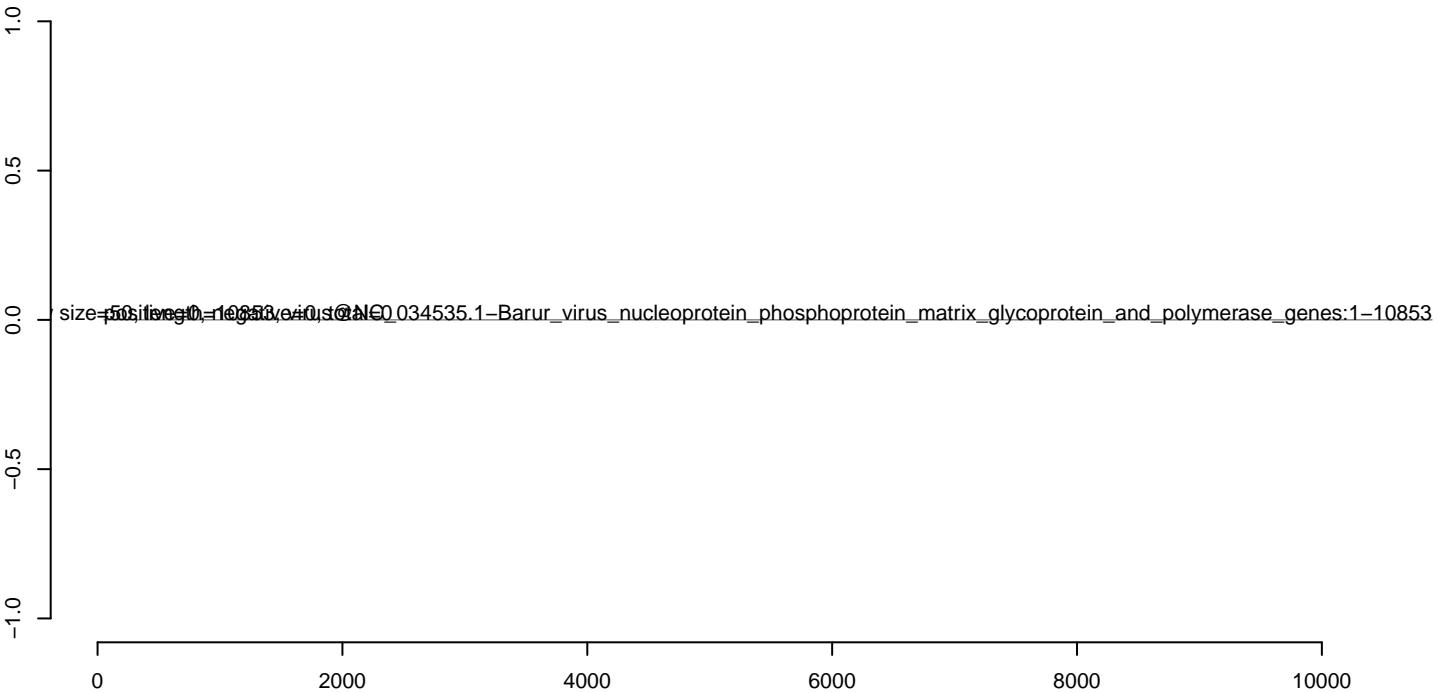
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep





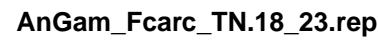
positive=0, negative=0, total=0



positive=0, negative=0, total=0



**positive- $\beta$ -undercarboxylic tripeptide-rich protein matrix hypothetical proteins glycoprotein hypothetical proteins and polymerase genes:1-13593**



positive=0, negative=0, total=0



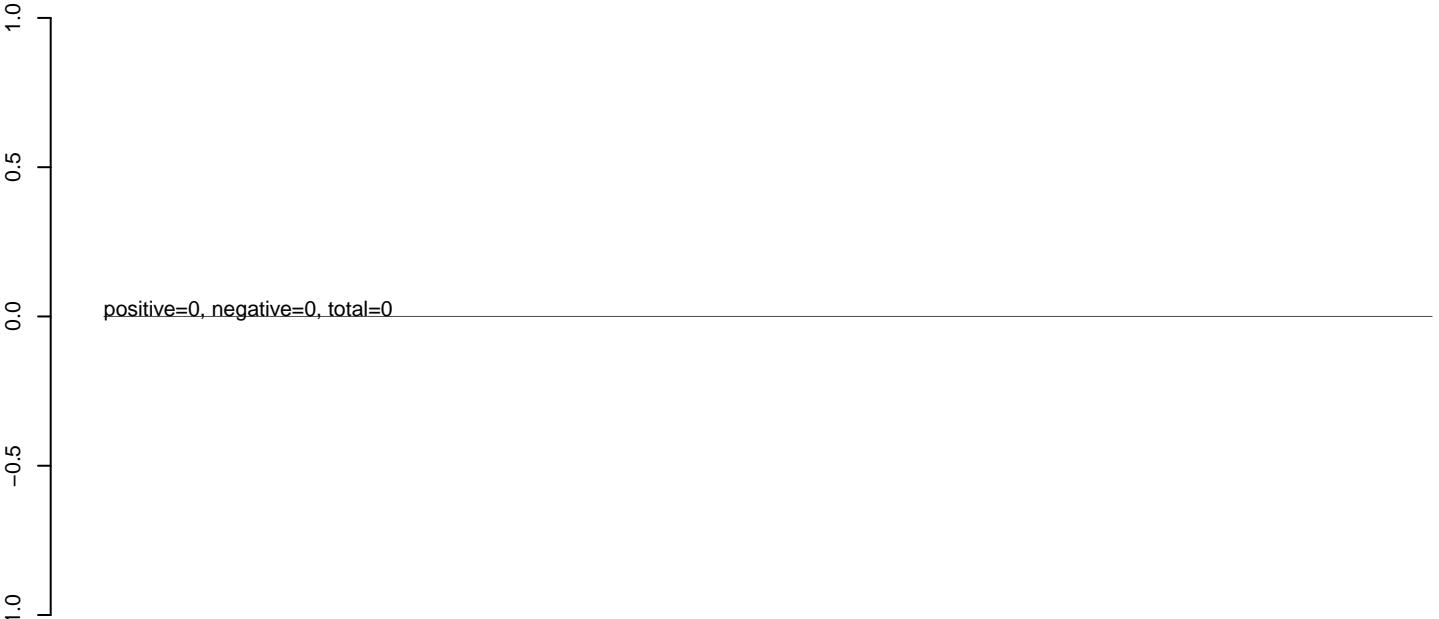
positive=0, negative=0, total=0



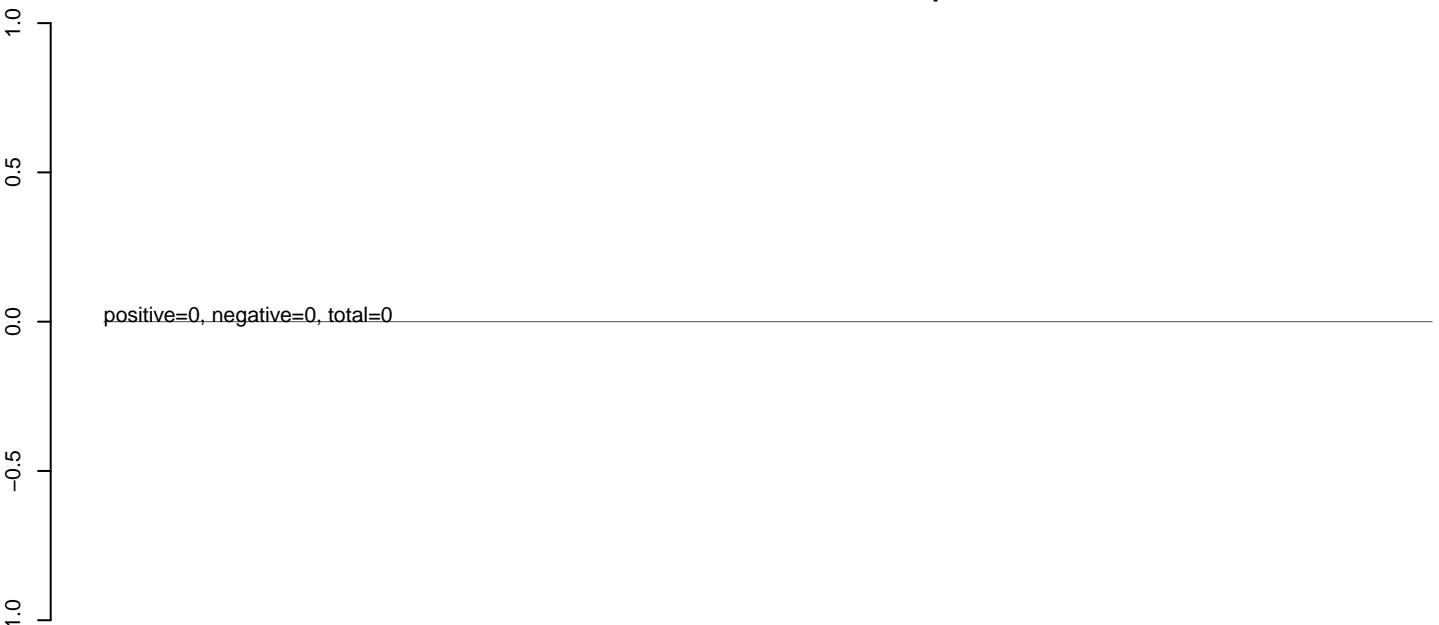
Manitobaproteins0 metaproteins0 ribophoprotein hypothetical proteins matrix glycoprotein hypothetical protein and polymerase genes:1-13784

Warrington, Cheshire, England

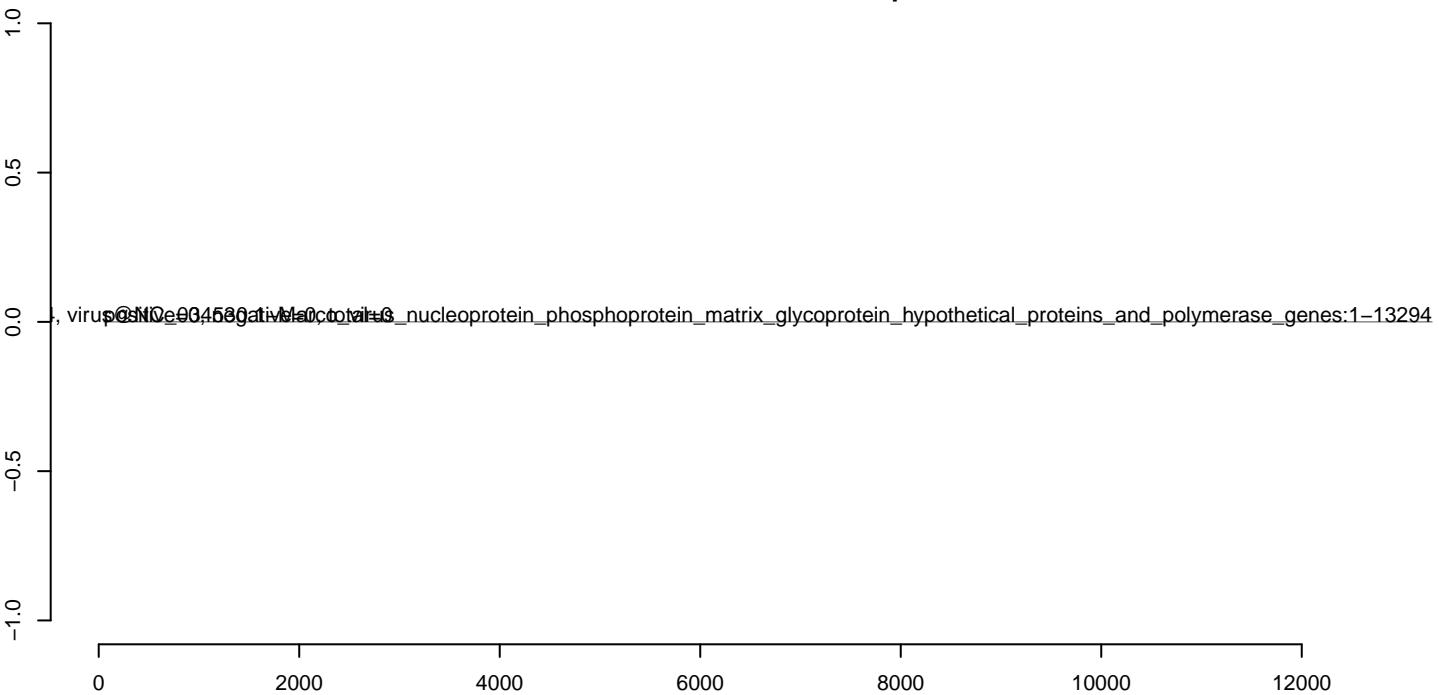
### AnGam\_Fcanc\_TN.18\_23.rep



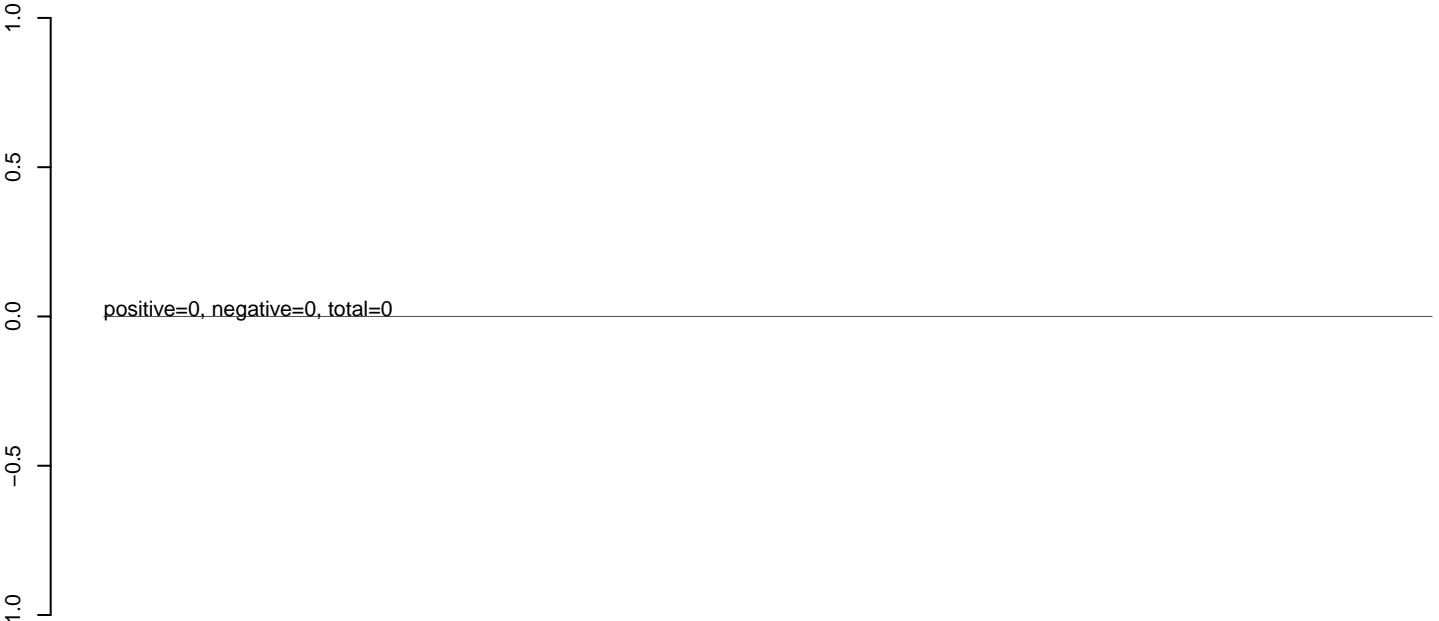
### AnGam\_Fcanc\_TN.24\_35.rep



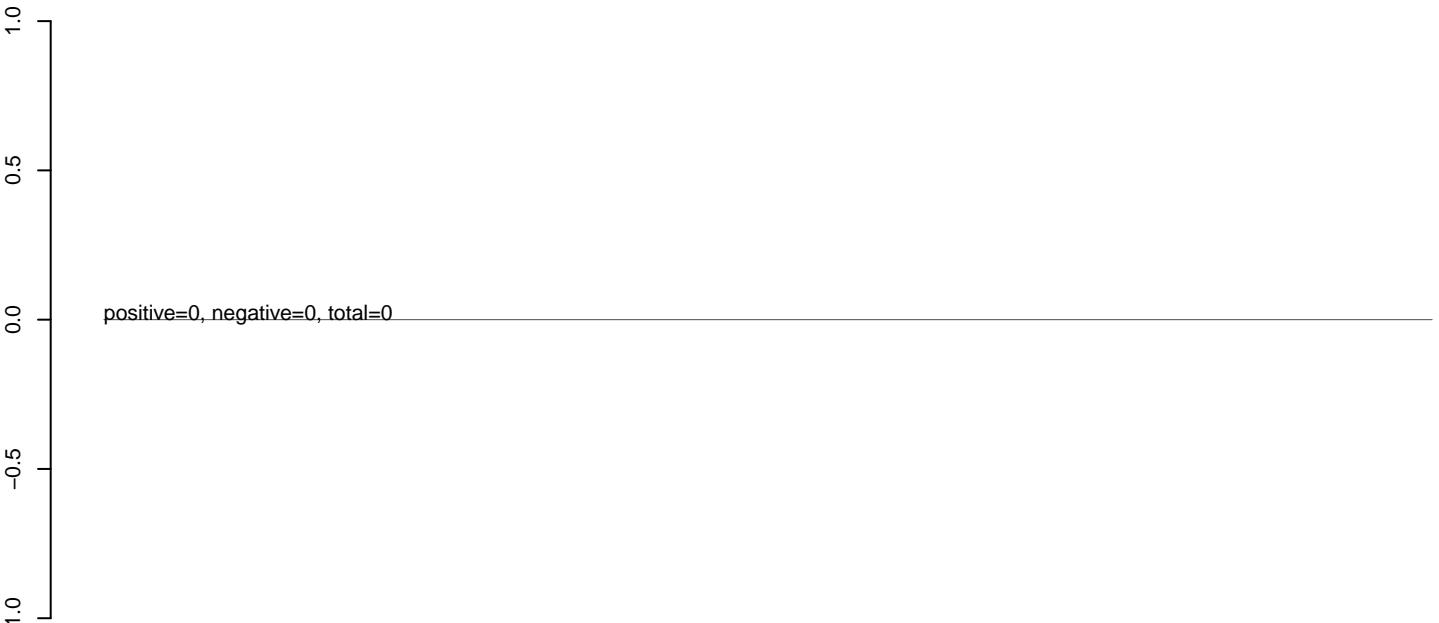
### AnGam\_Fcanc\_TN.rep



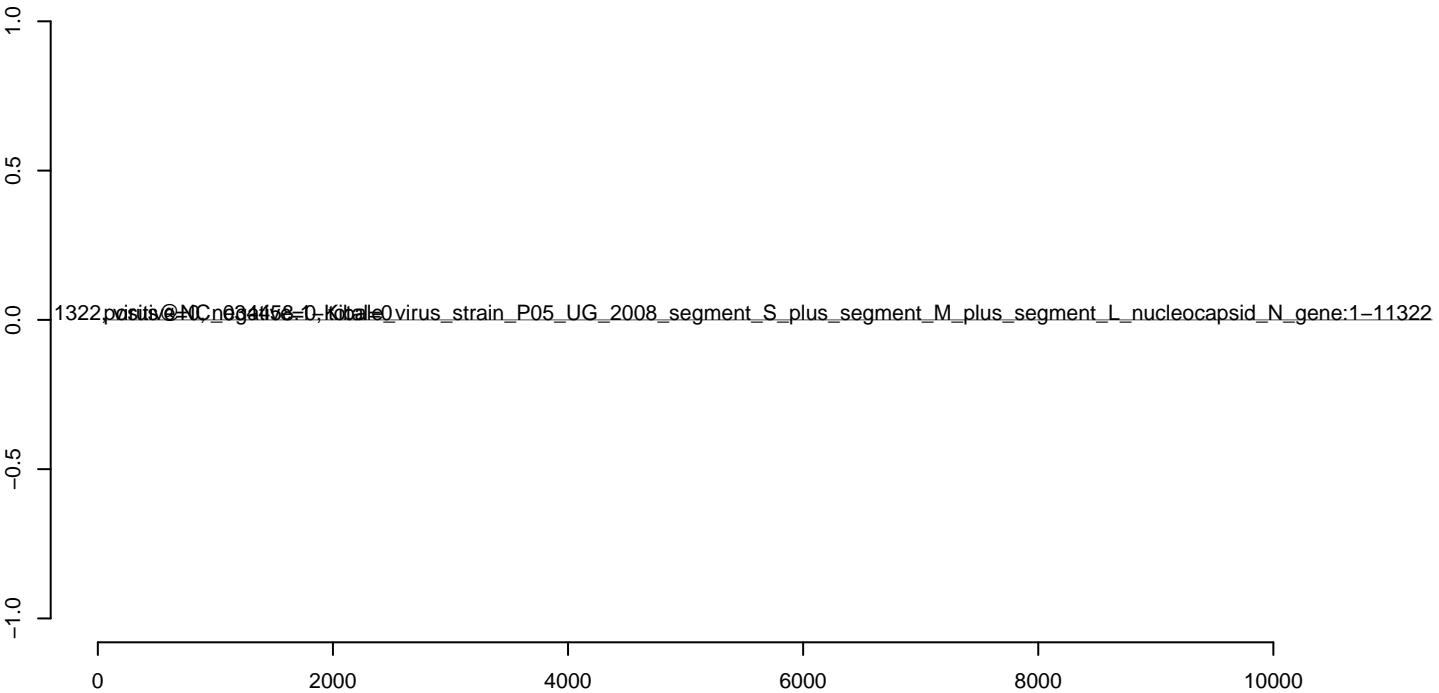
### AnGam\_Fcanc\_TN.18\_23.rep



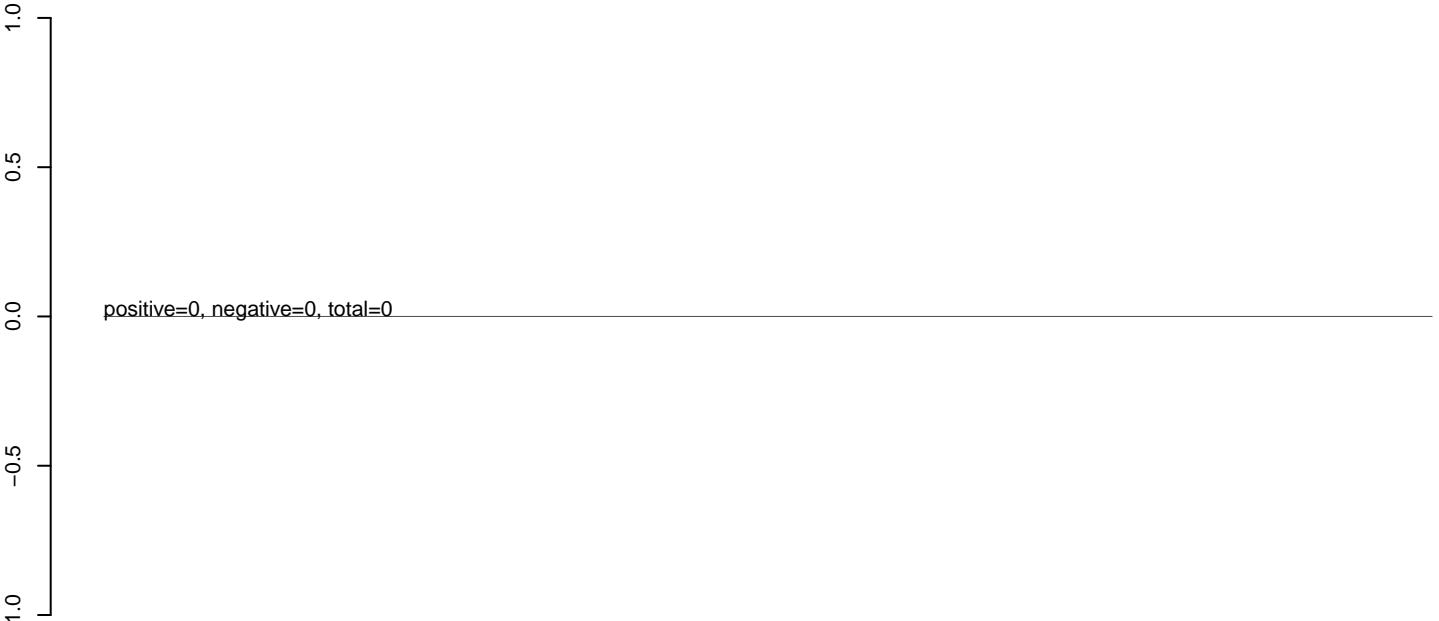
### AnGam\_Fcanc\_TN.24\_35.rep



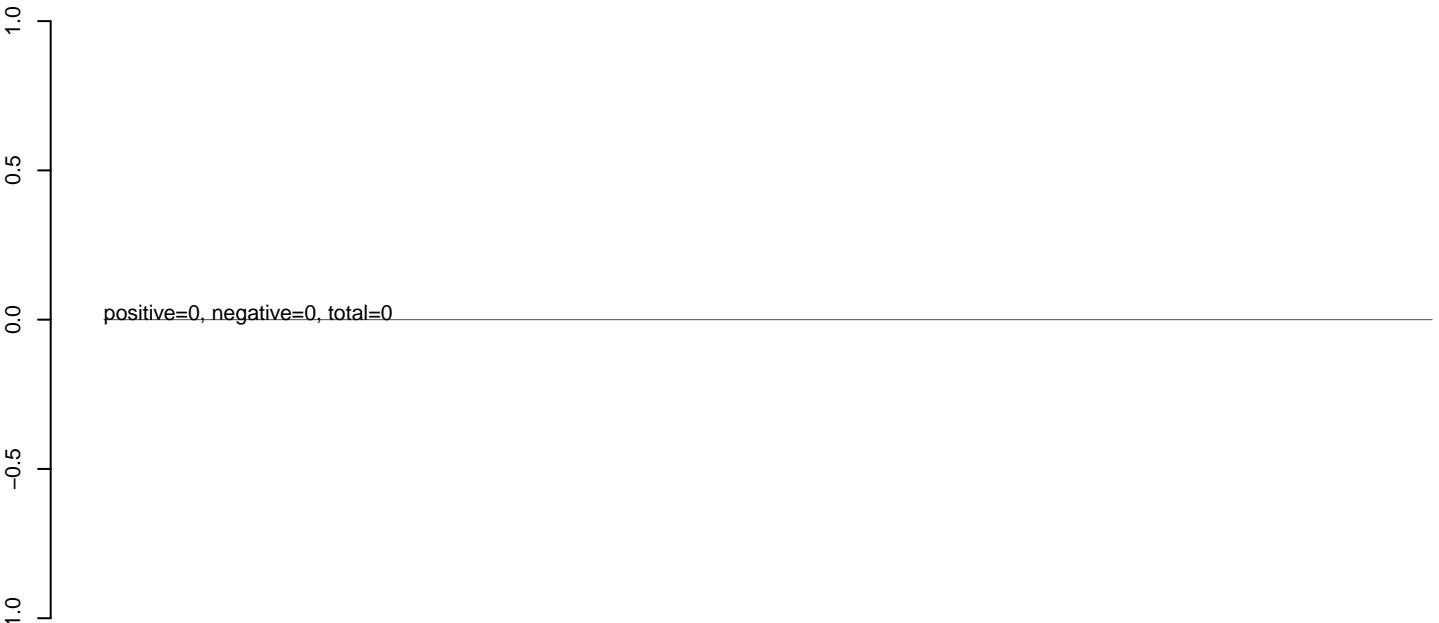
### AnGam\_Fcanc\_TN.rep



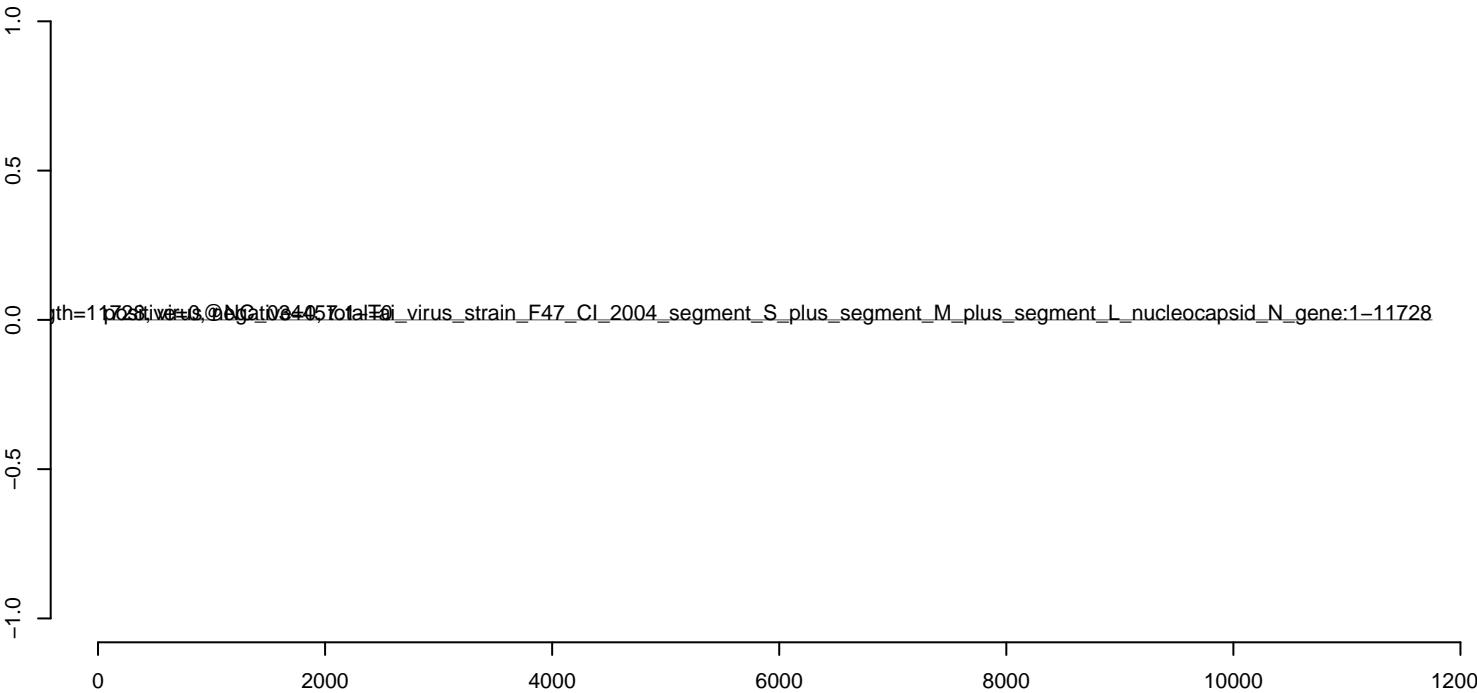
### AnGam\_Fcanc\_TN.18\_23.rep



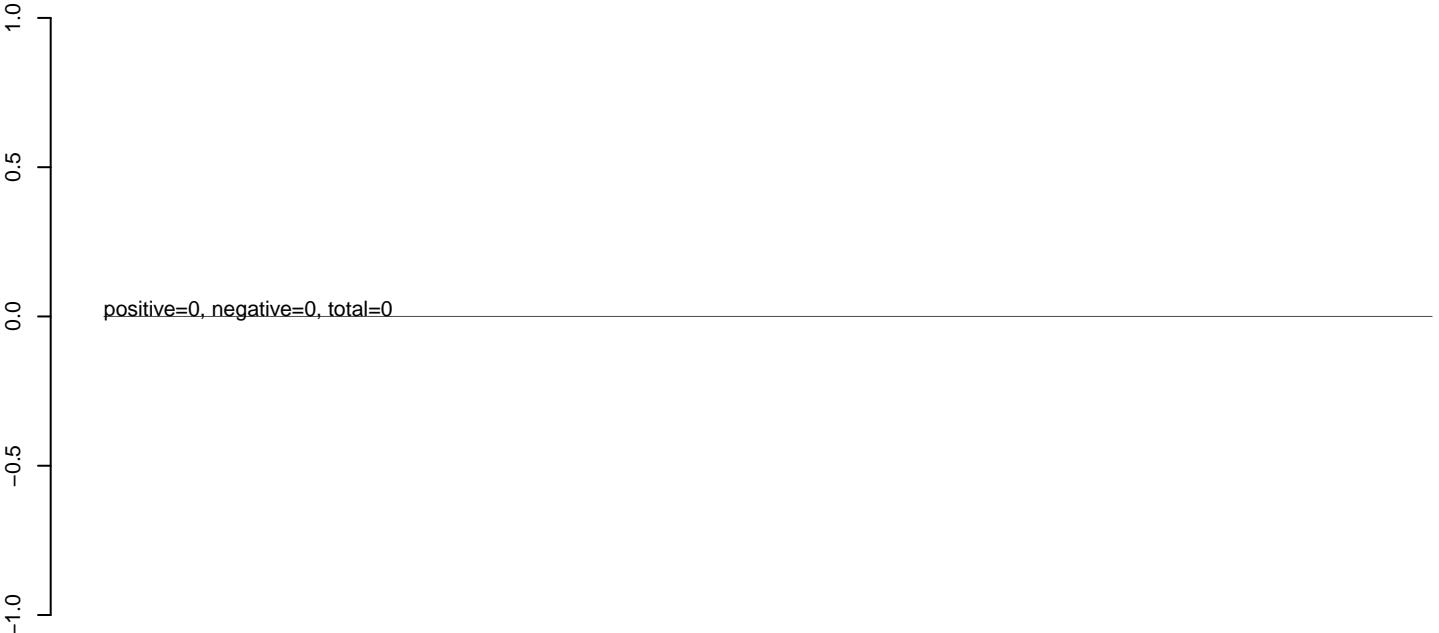
### AnGam\_Fcanc\_TN.24\_35.rep



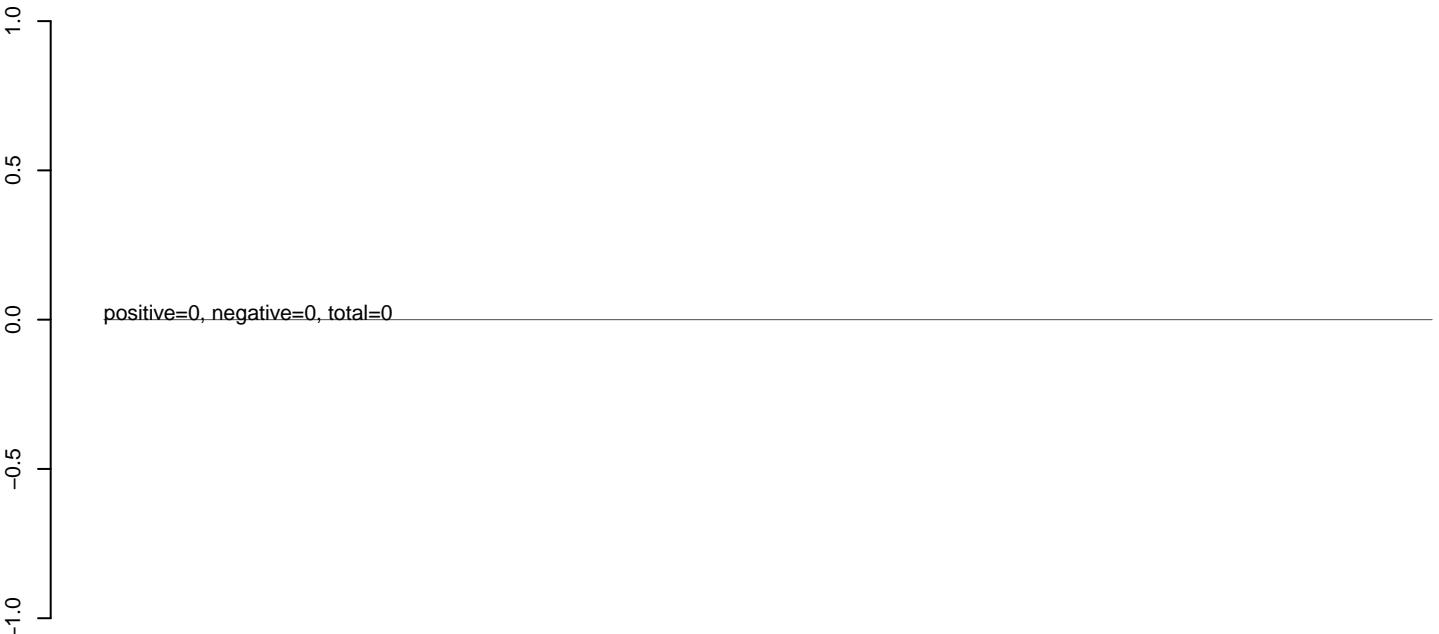
### AnGam\_Fcanc\_TN.rep



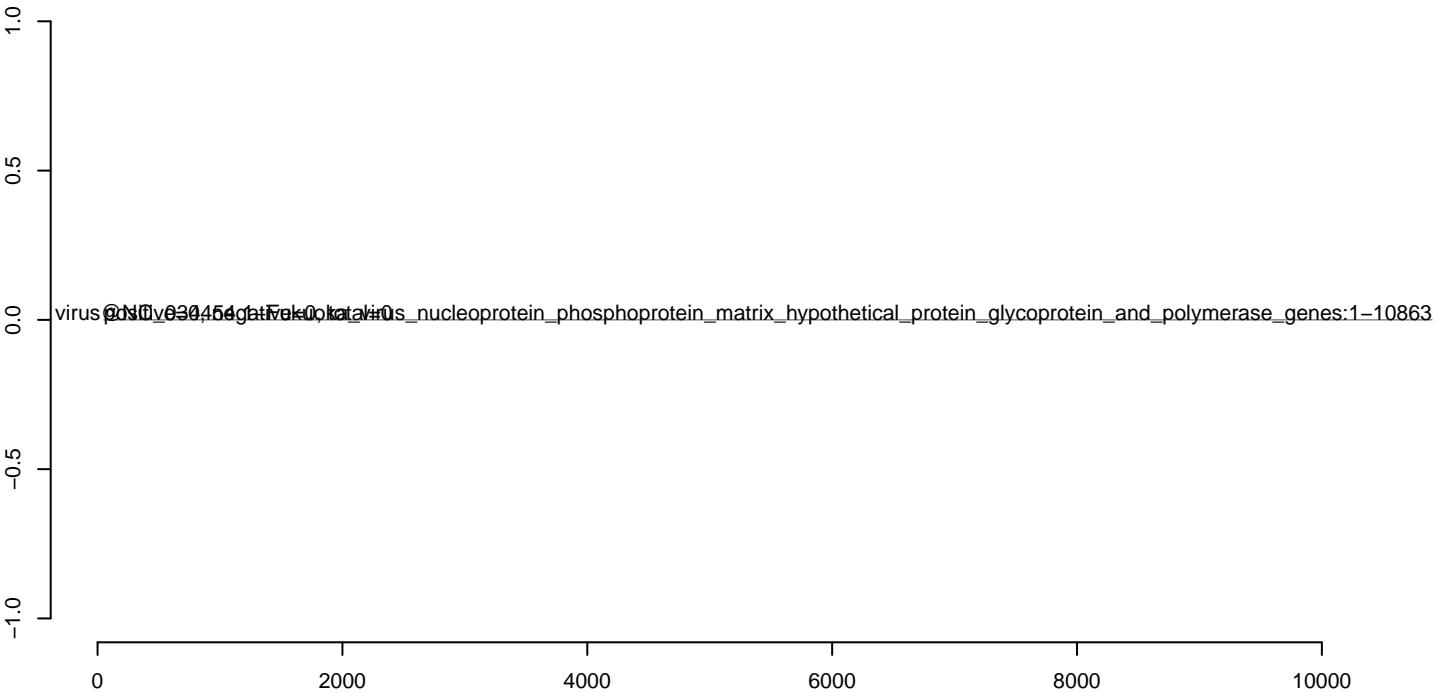
### AnGam\_Fcanc\_TN.18\_23.rep



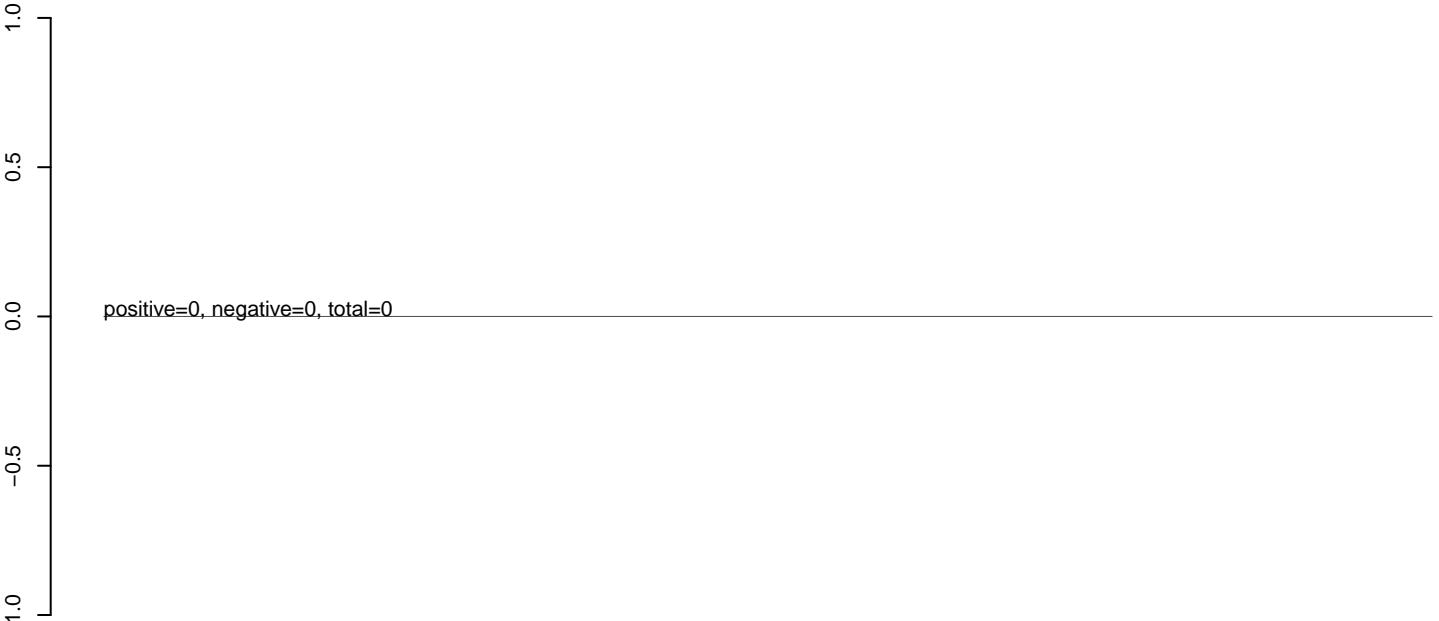
### AnGam\_Fcanc\_TN.24\_35.rep



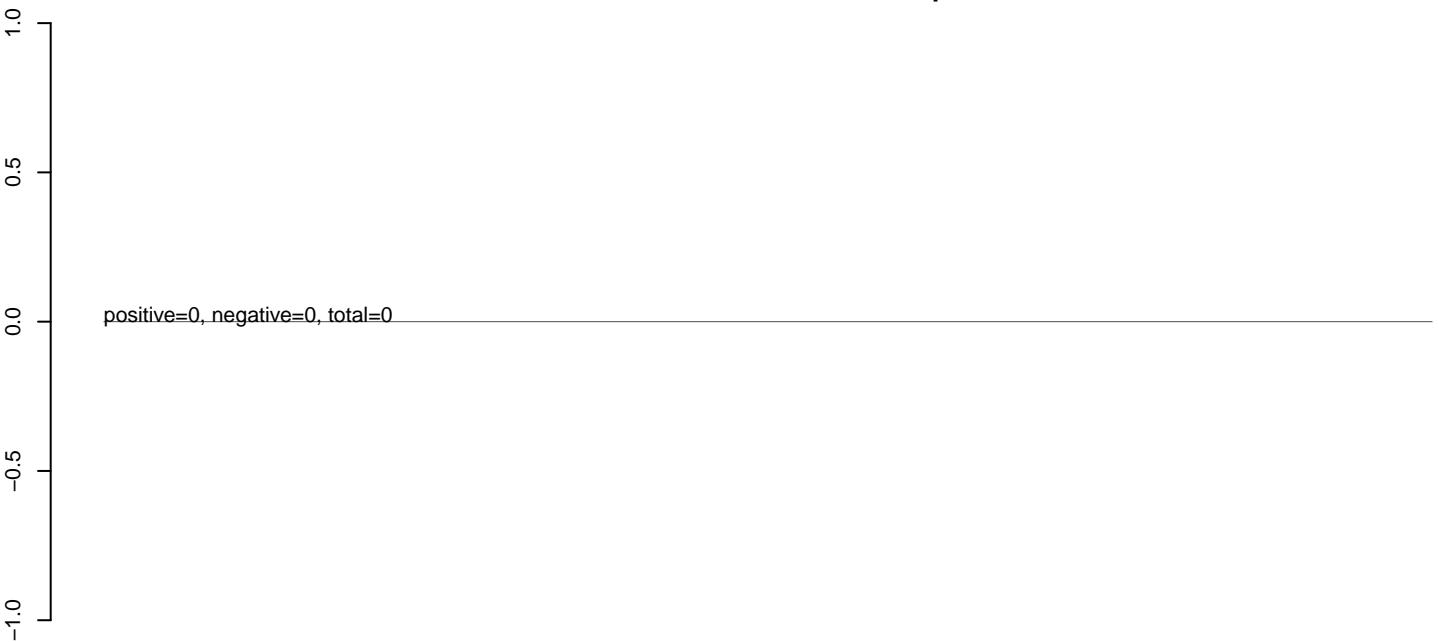
### AnGam\_Fcanc\_TN.rep



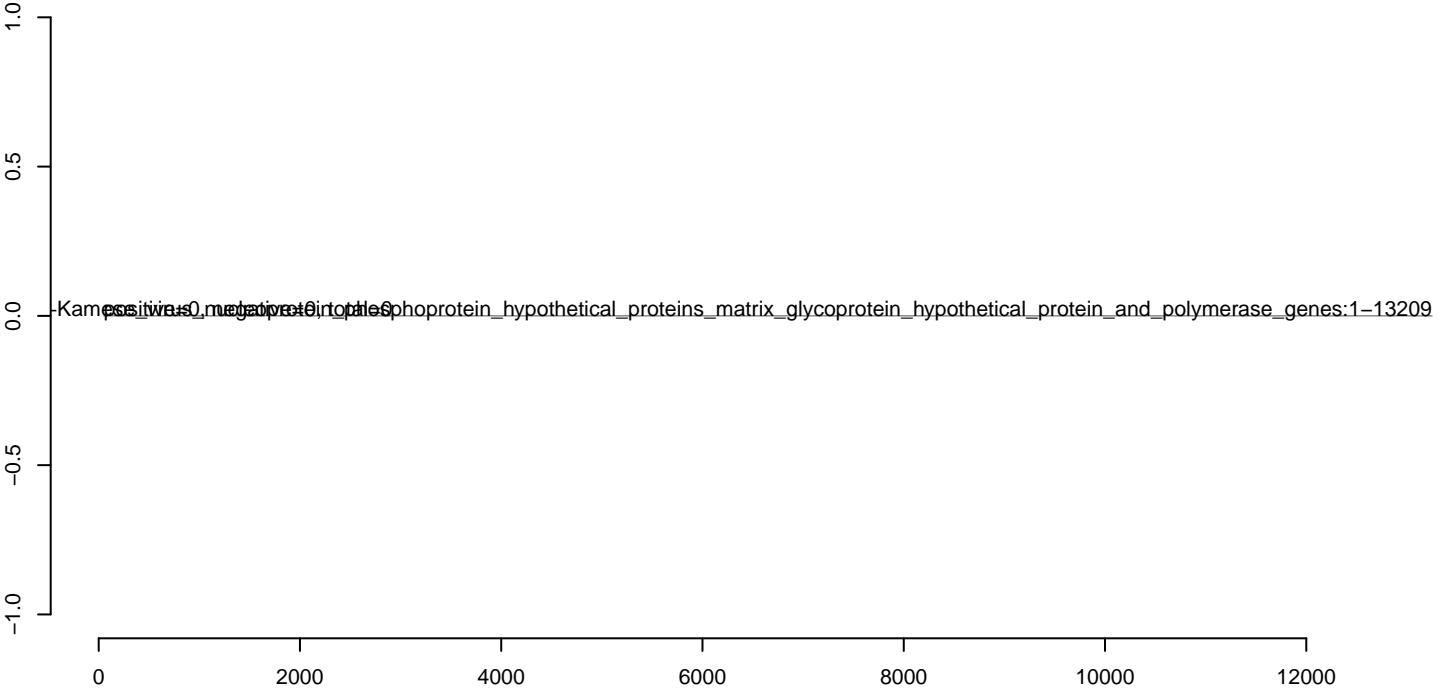
### AnGam\_Fcanc\_TN.18\_23.rep



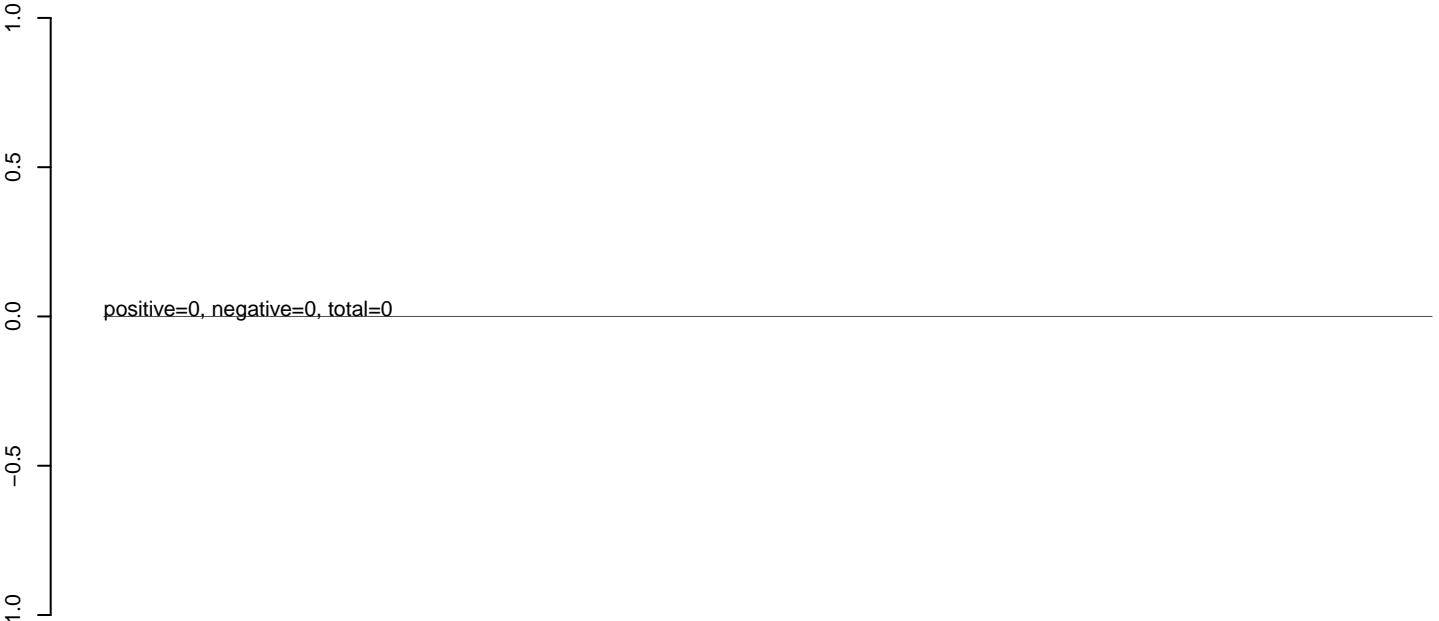
### AnGam\_Fcanc\_TN.24\_35.rep



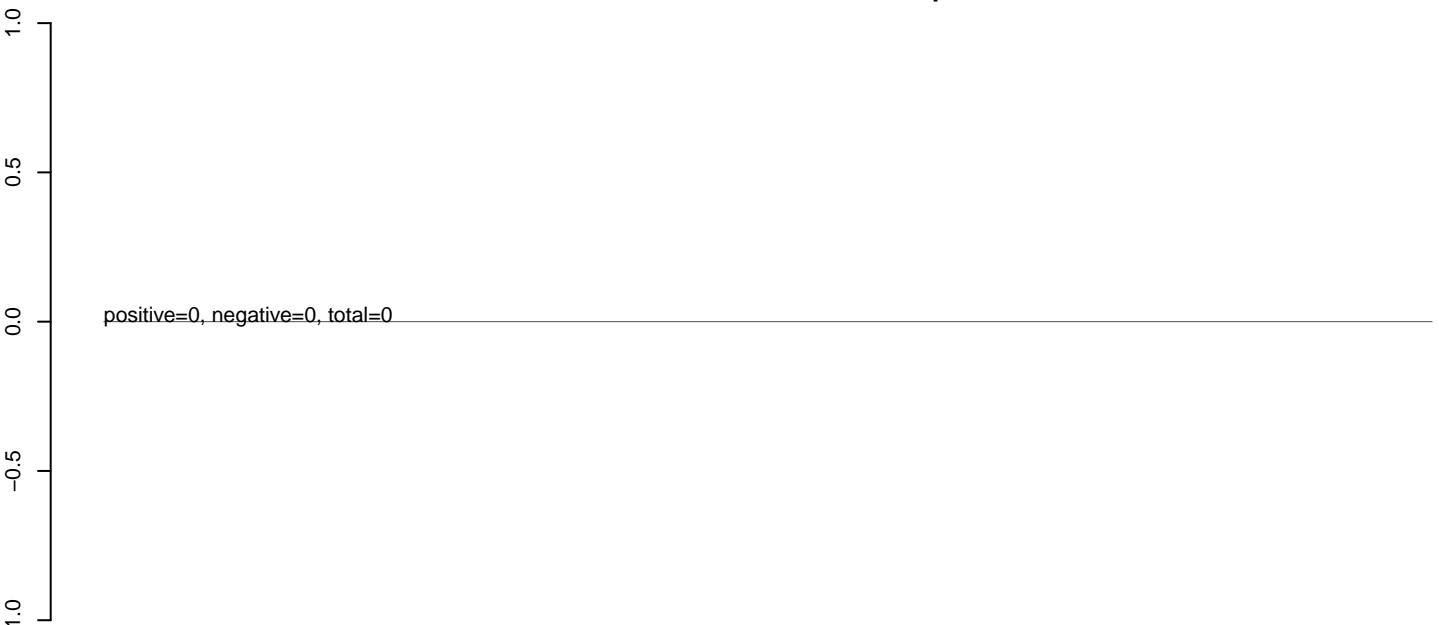
### AnGam\_Fcanc\_TN.rep



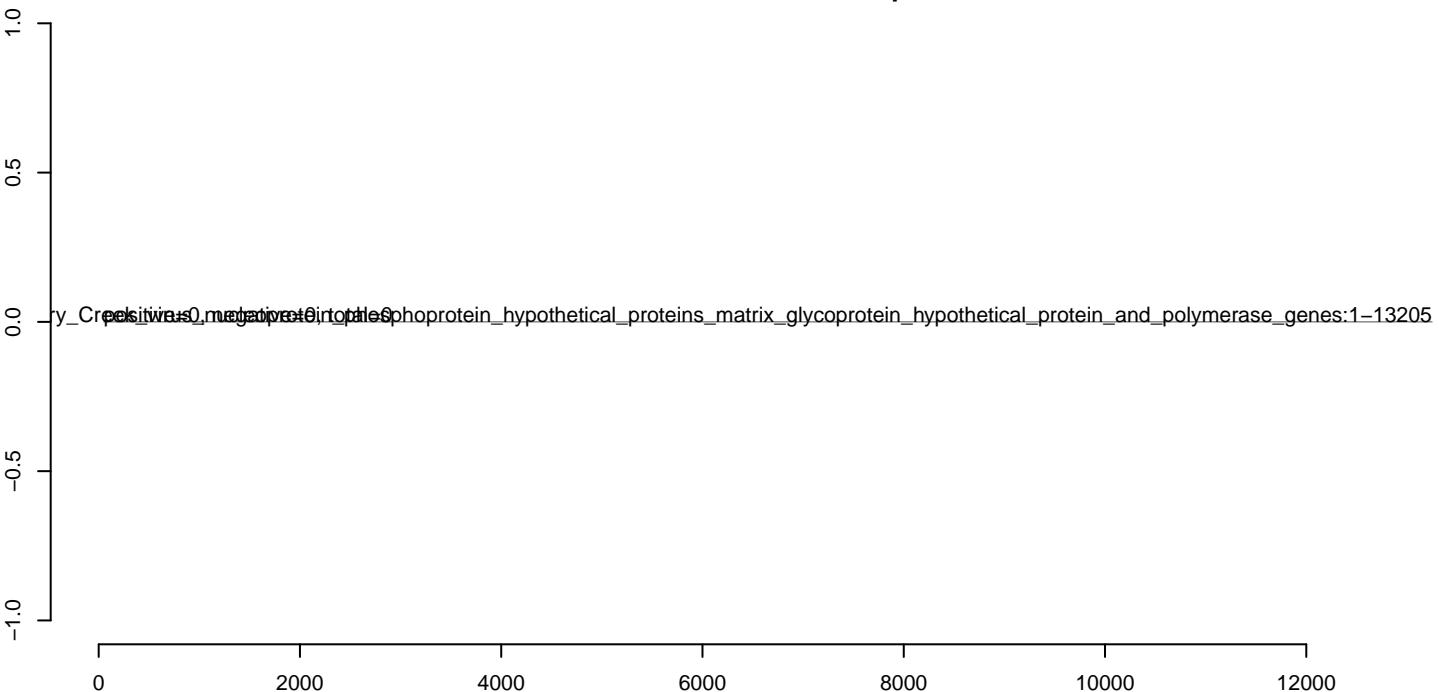
### AnGam\_Fcanc\_TN.18\_23.rep



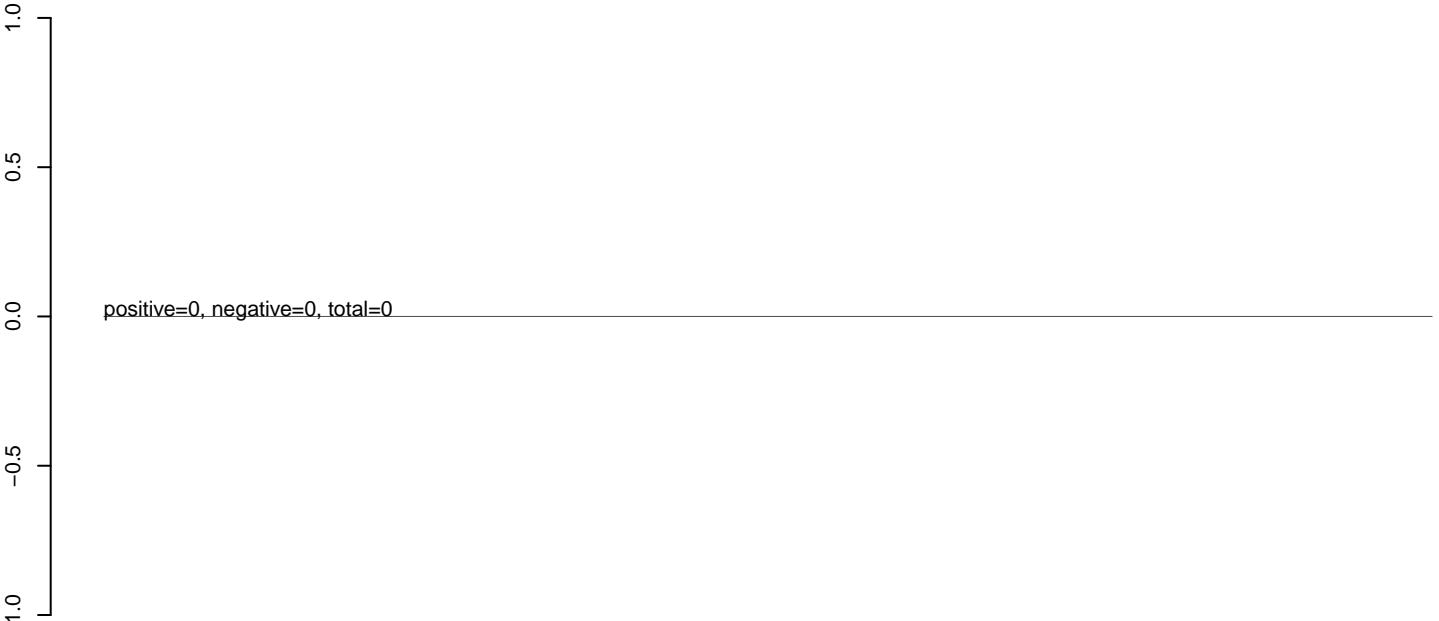
### AnGam\_Fcanc\_TN.24\_35.rep



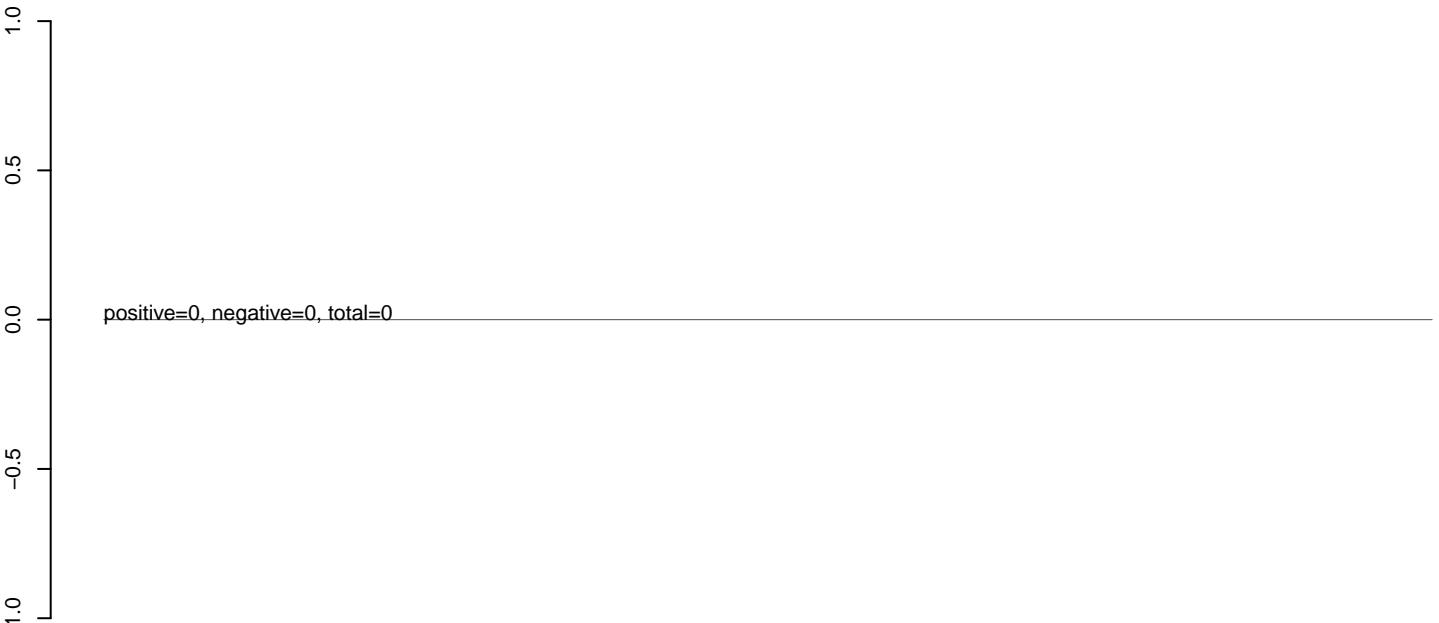
### AnGam\_Fcanc\_TN.rep



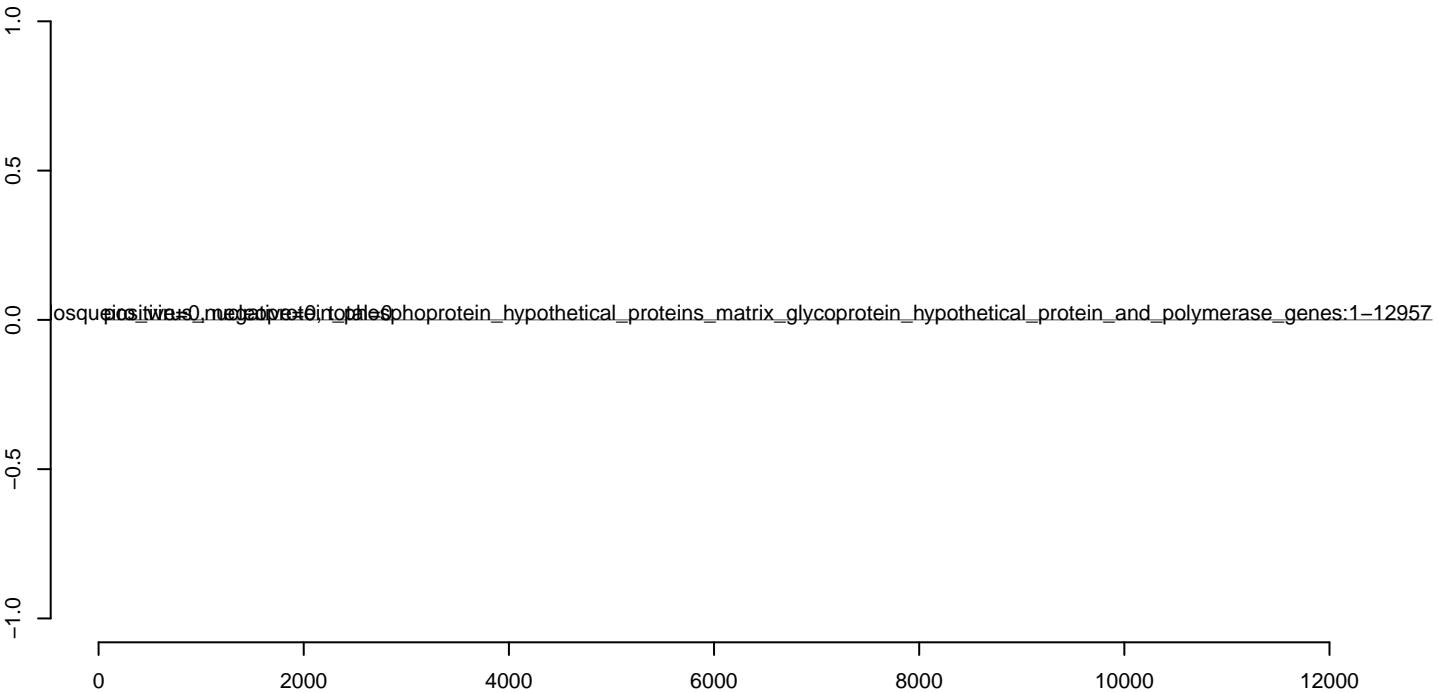
### AnGam\_Fcanc\_TN.18\_23.rep



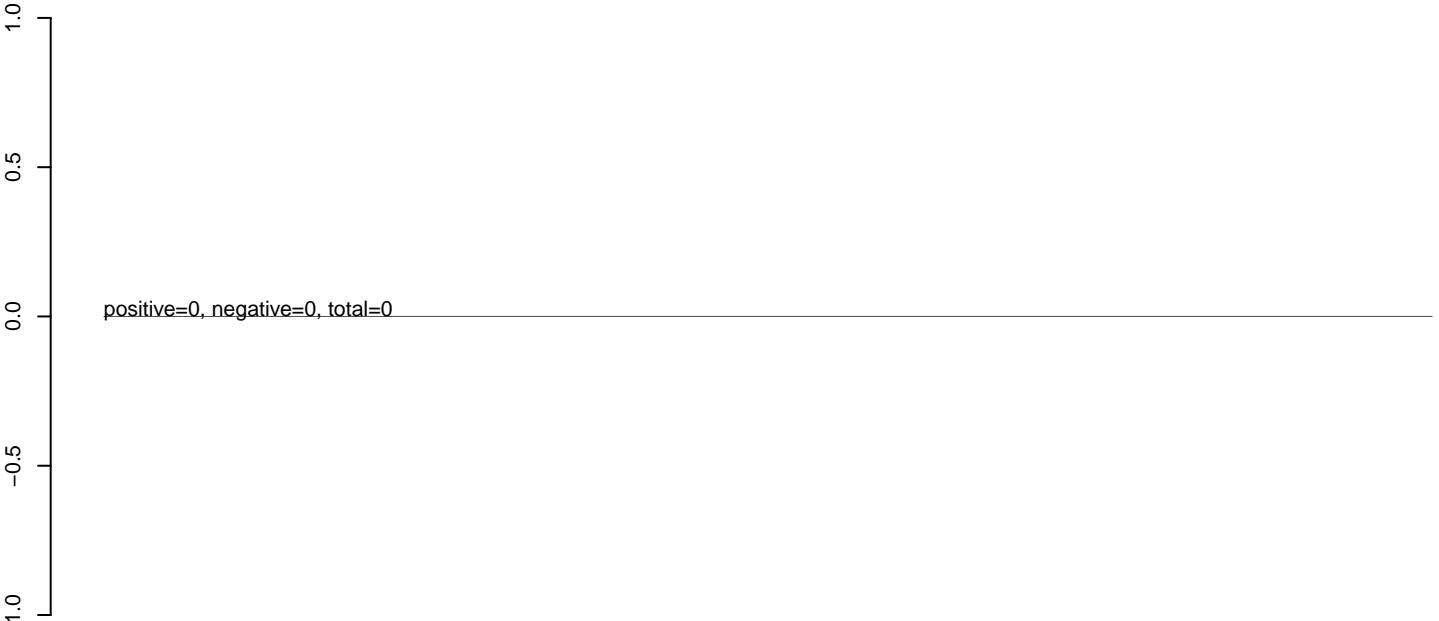
### AnGam\_Fcanc\_TN.24\_35.rep



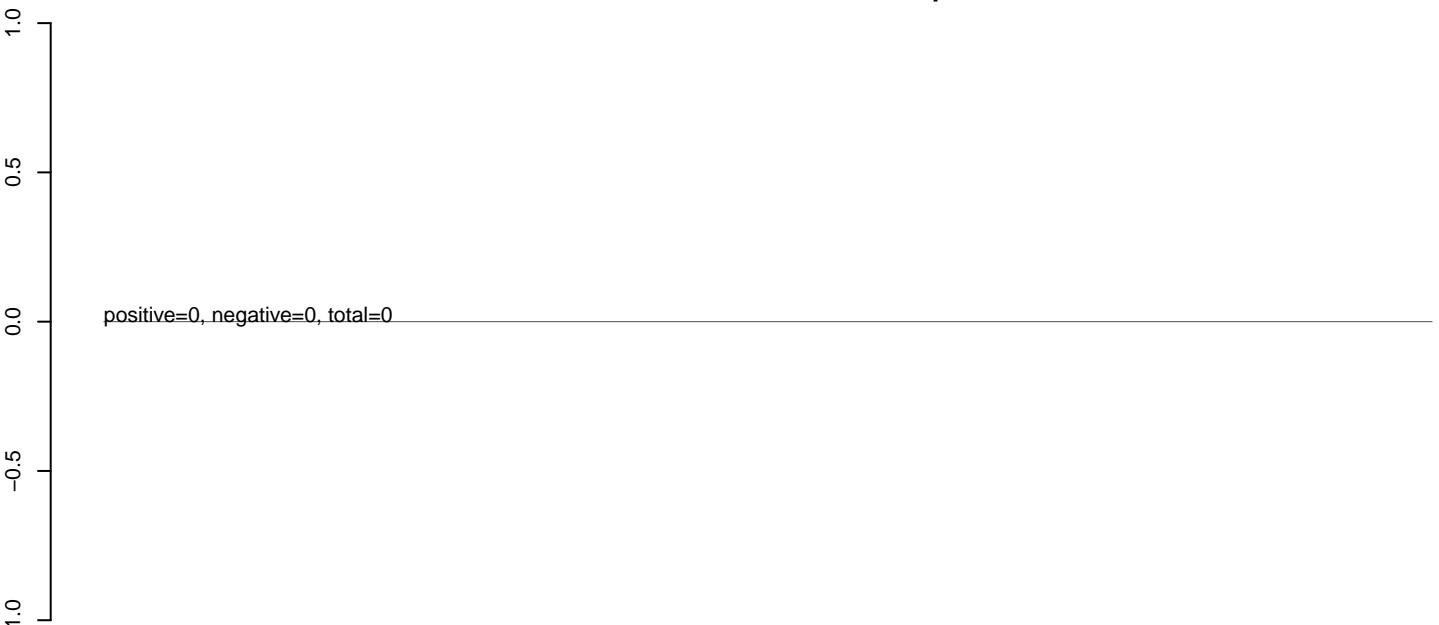
### AnGam\_Fcanc\_TN.rep



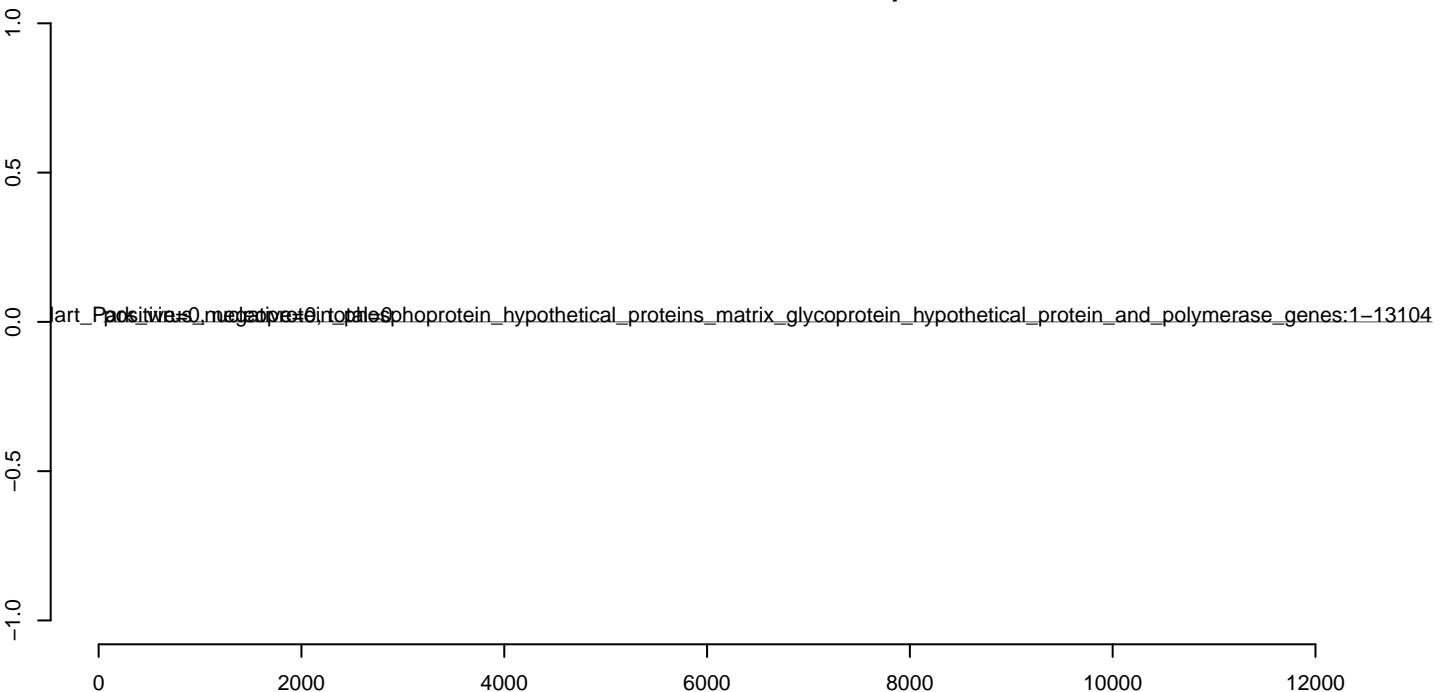
### AnGam\_Fcanc\_TN.18\_23.rep



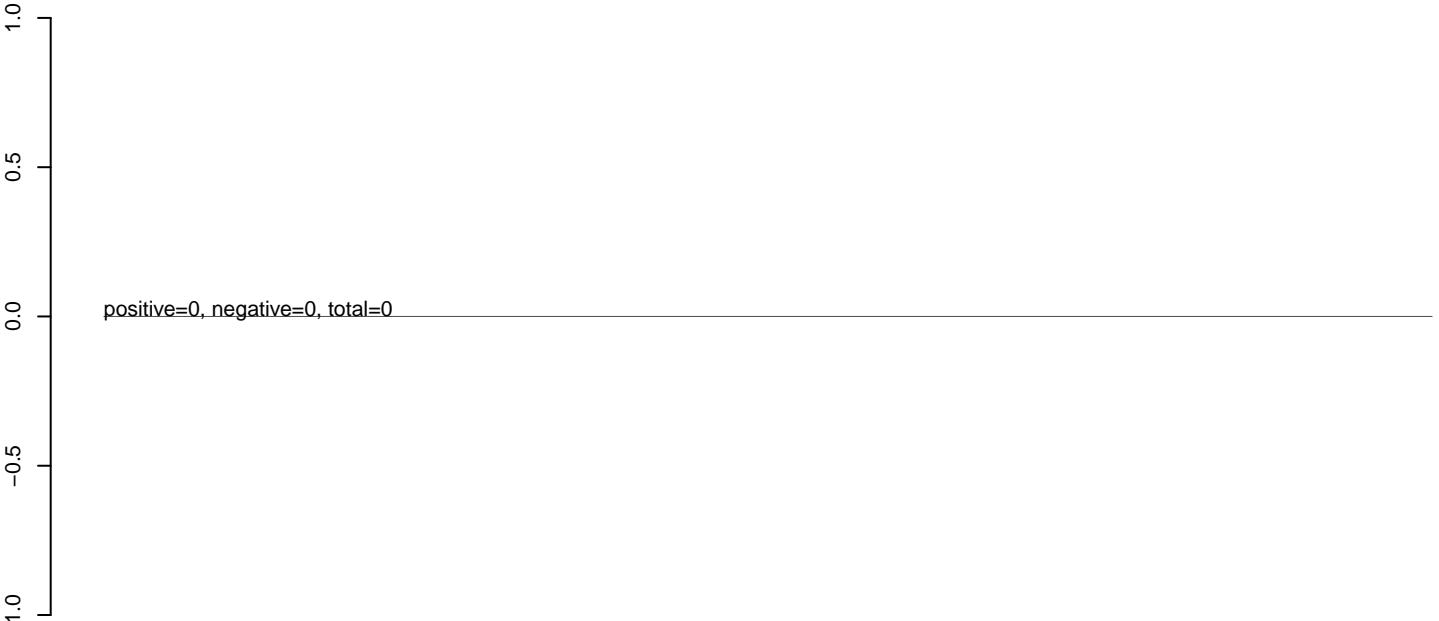
### AnGam\_Fcanc\_TN.24\_35.rep



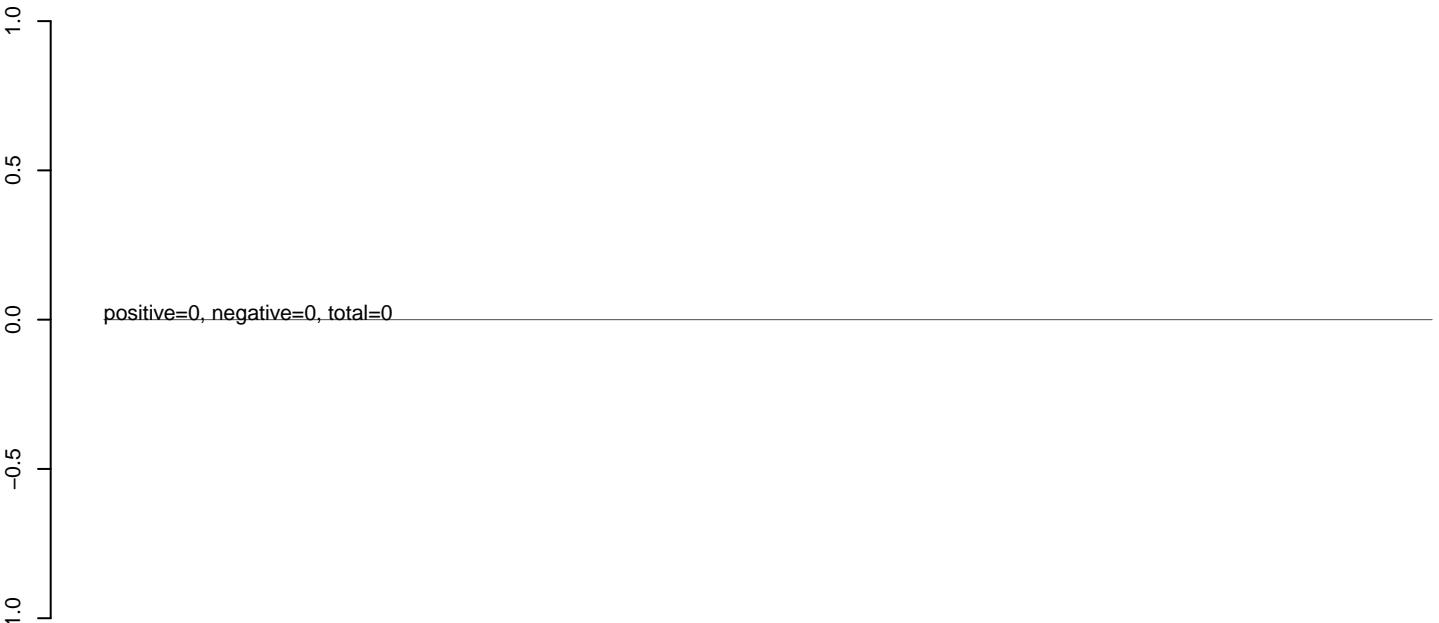
### AnGam\_Fcanc\_TN.rep



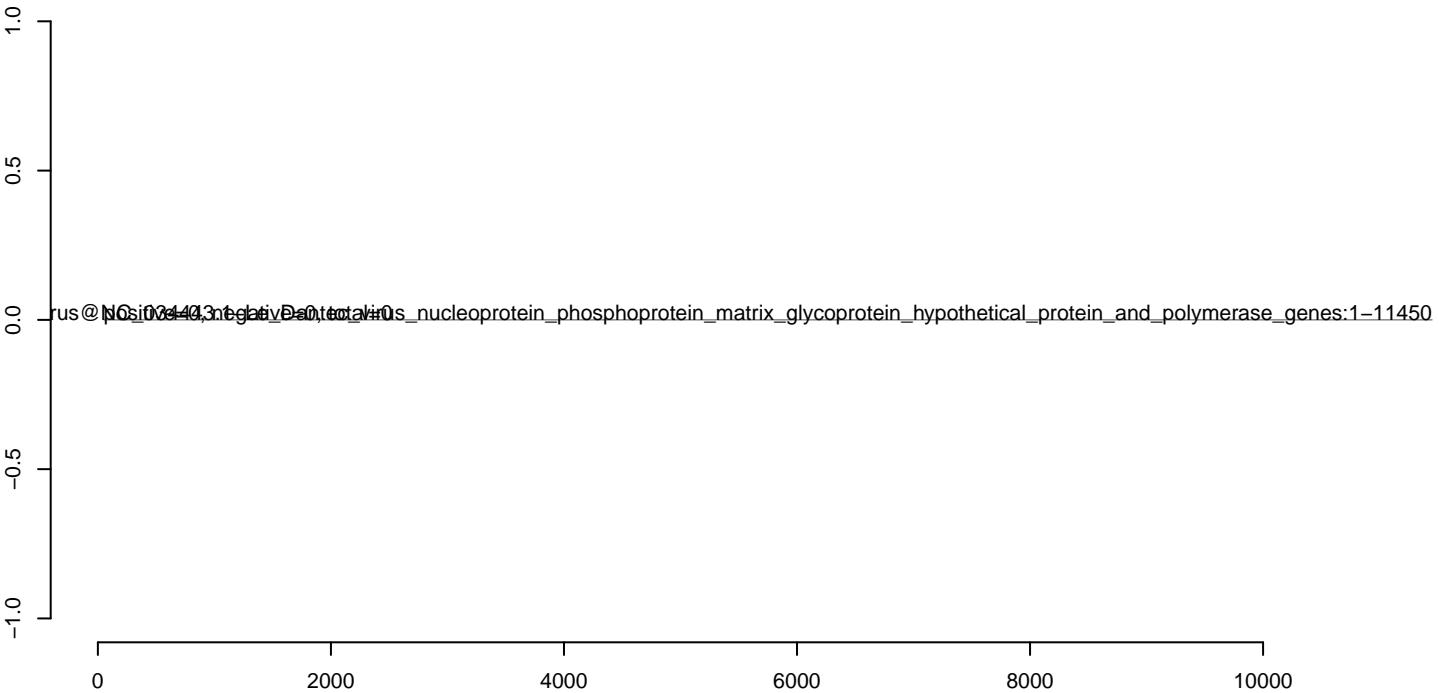
### AnGam\_Fcanc\_TN.18\_23.rep



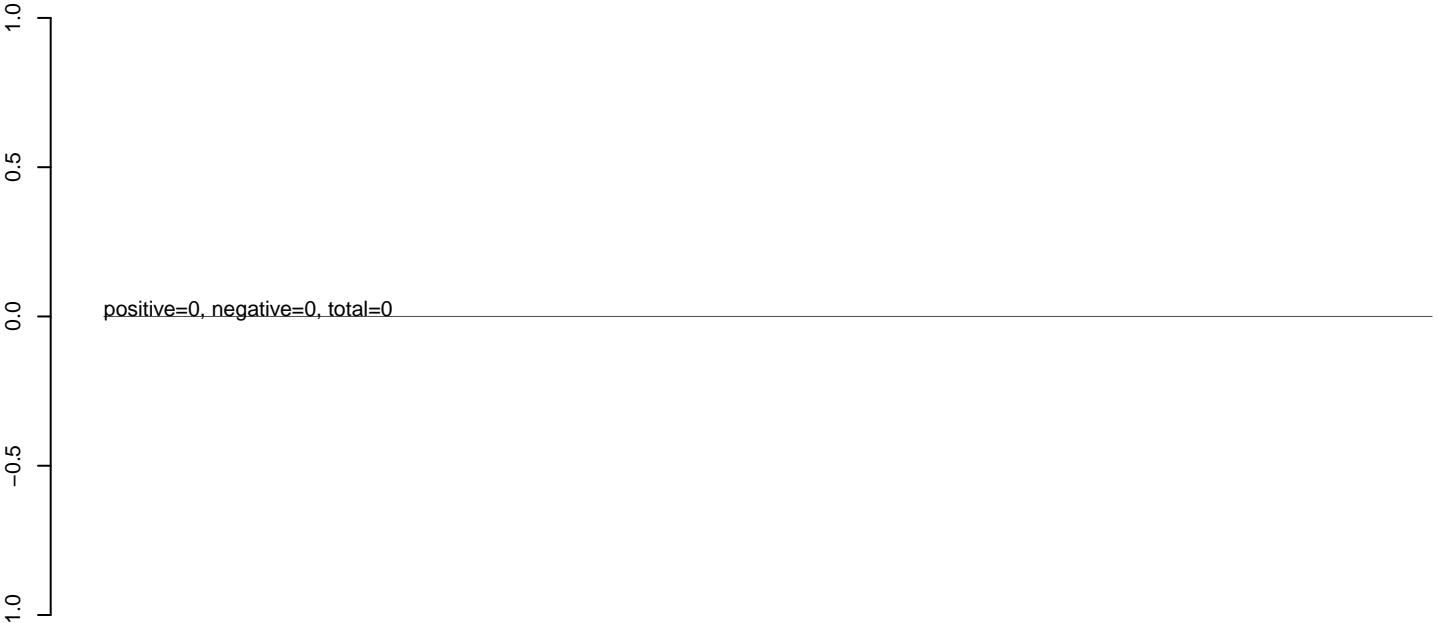
### AnGam\_Fcanc\_TN.24\_35.rep



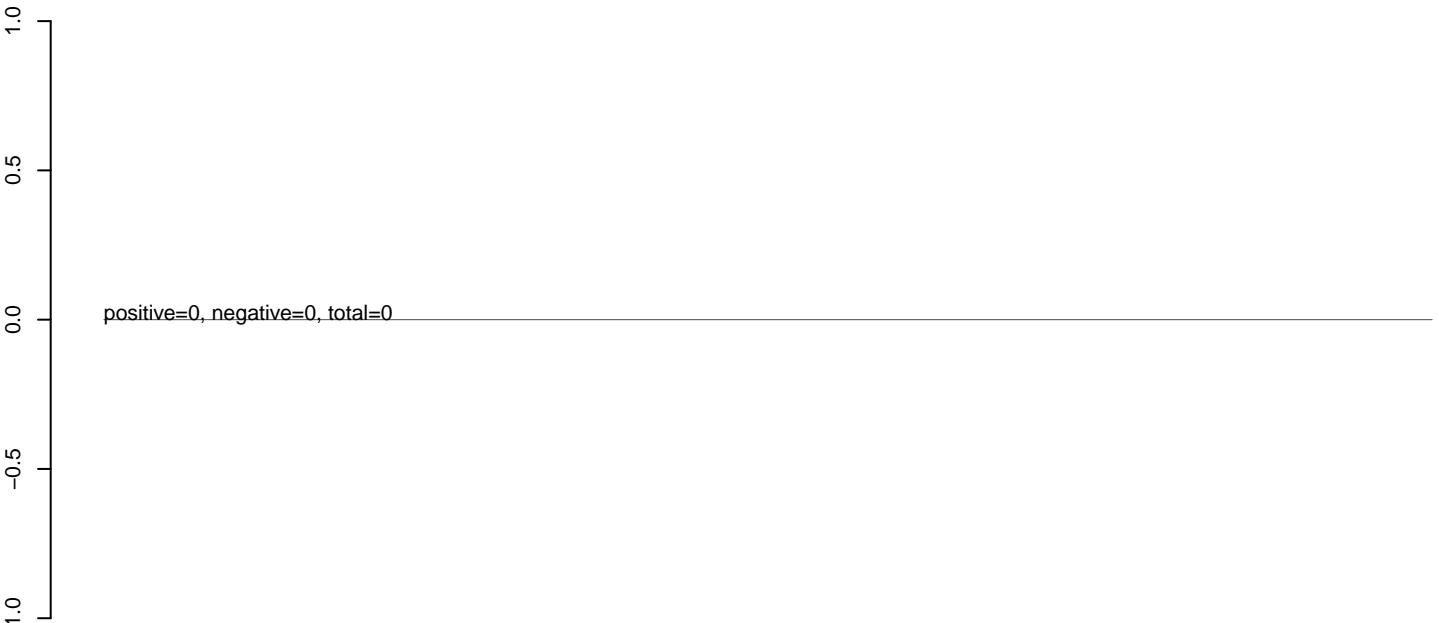
### AnGam\_Fcanc\_TN.rep



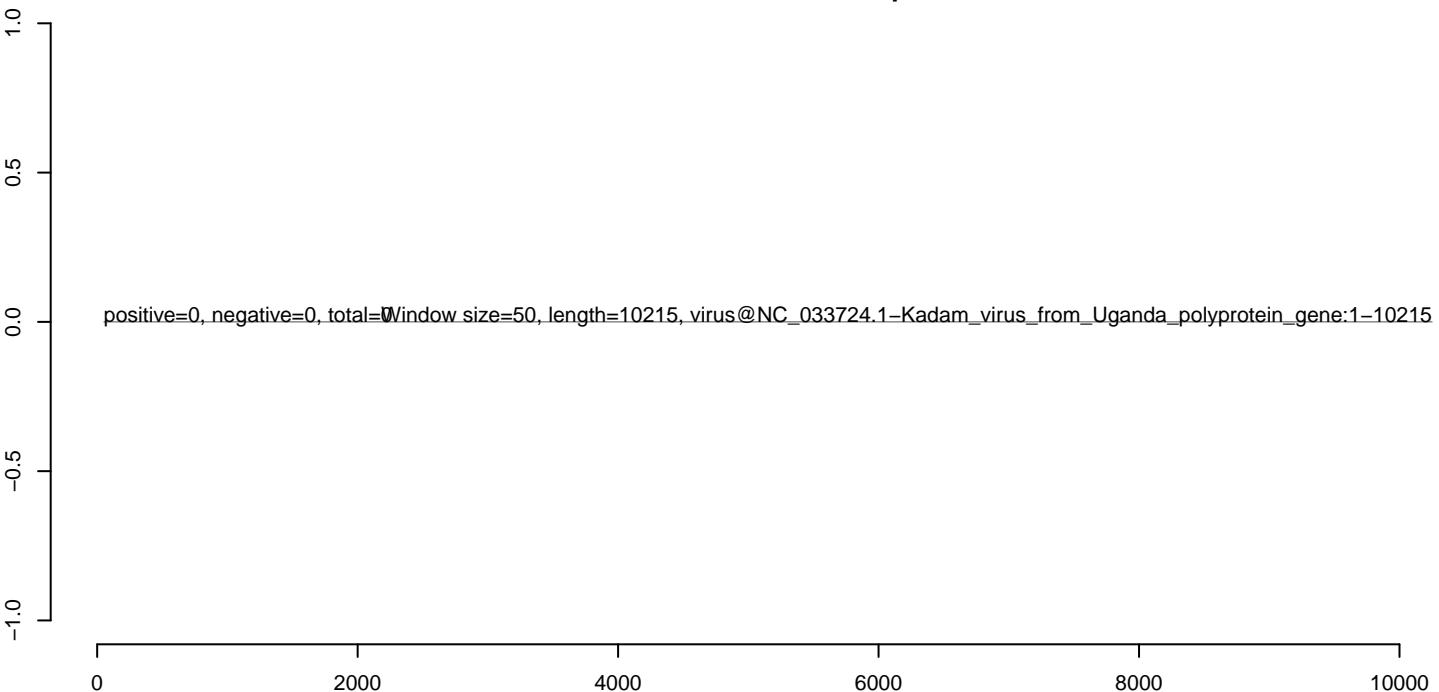
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep





positive=0, negative=0, total=0



positive=0, negative=0, total=0



positive=0, negative=Window, size=50, length=10251, virus@NC\_033723.1-Gadgets\_Gully\_virus\_from\_Australia\_polyprotein\_gene:1-10251

### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

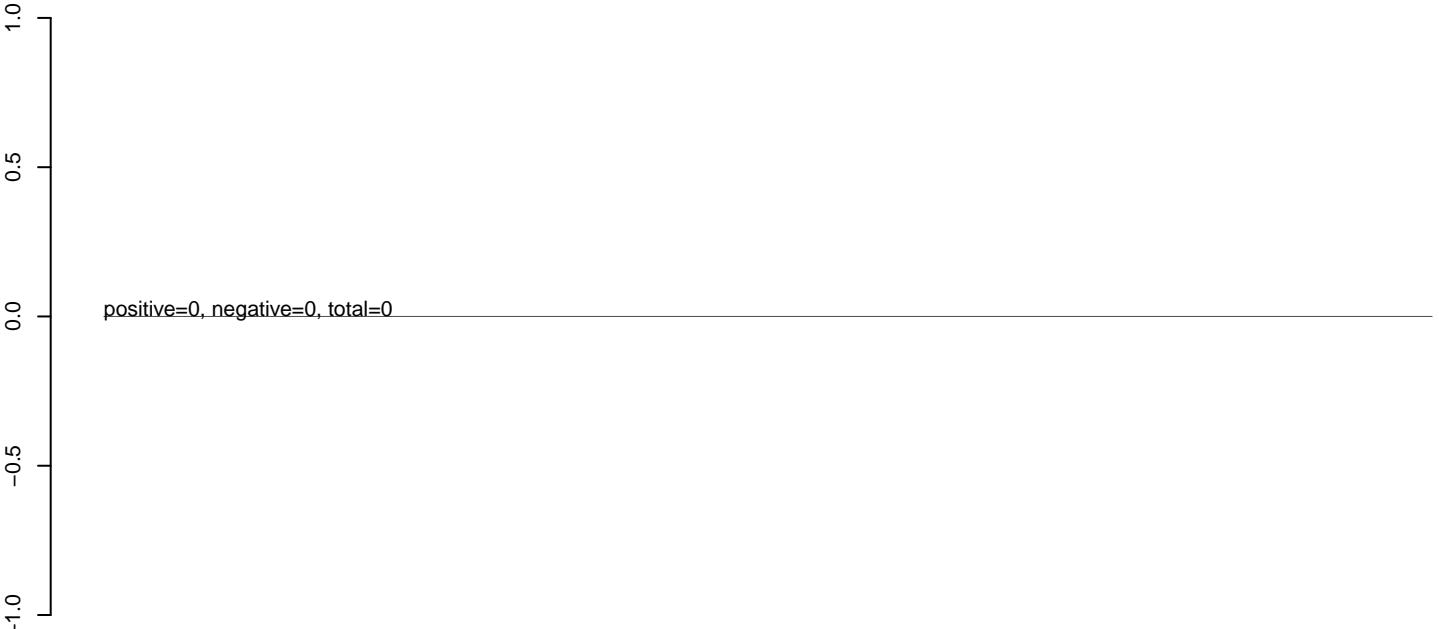
positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.rep

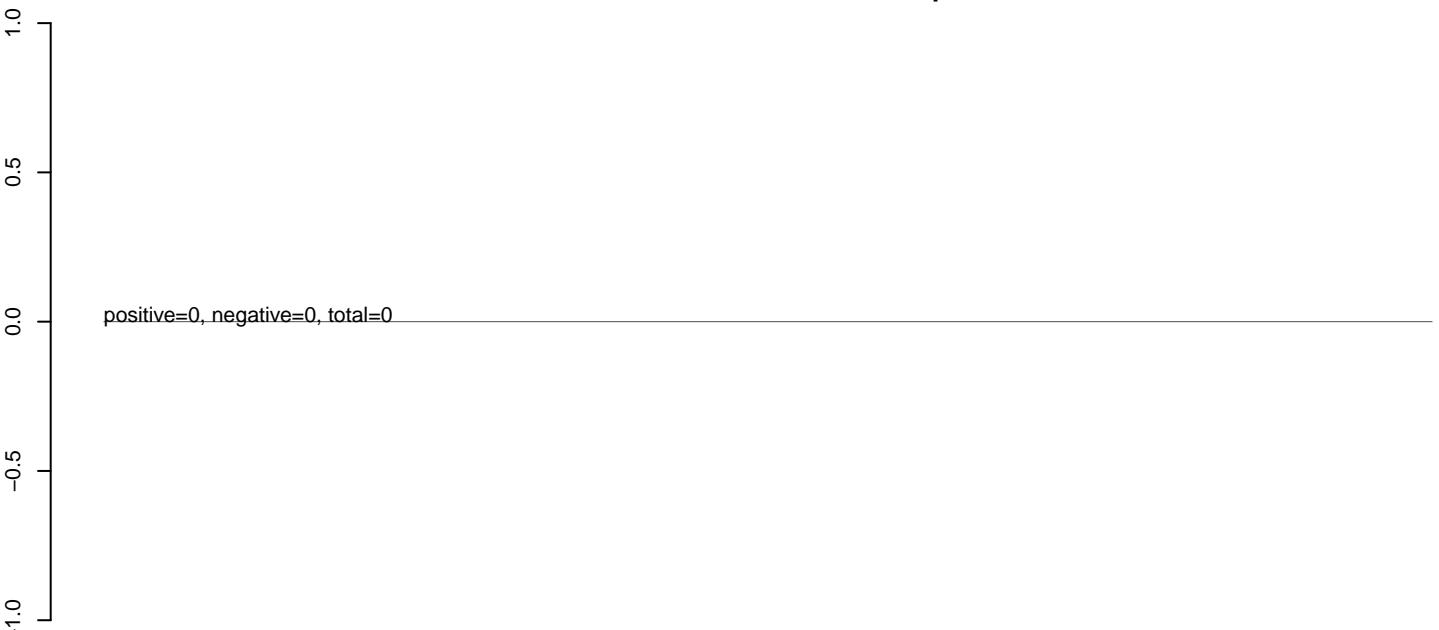
positive=0, negative=0, total=0 window size=50, length=10266, virus@NC\_033721.1-Meaban\_virus\_from\_France\_polyprotein\_gene:1-10266

0 2000 4000 6000 8000 10000

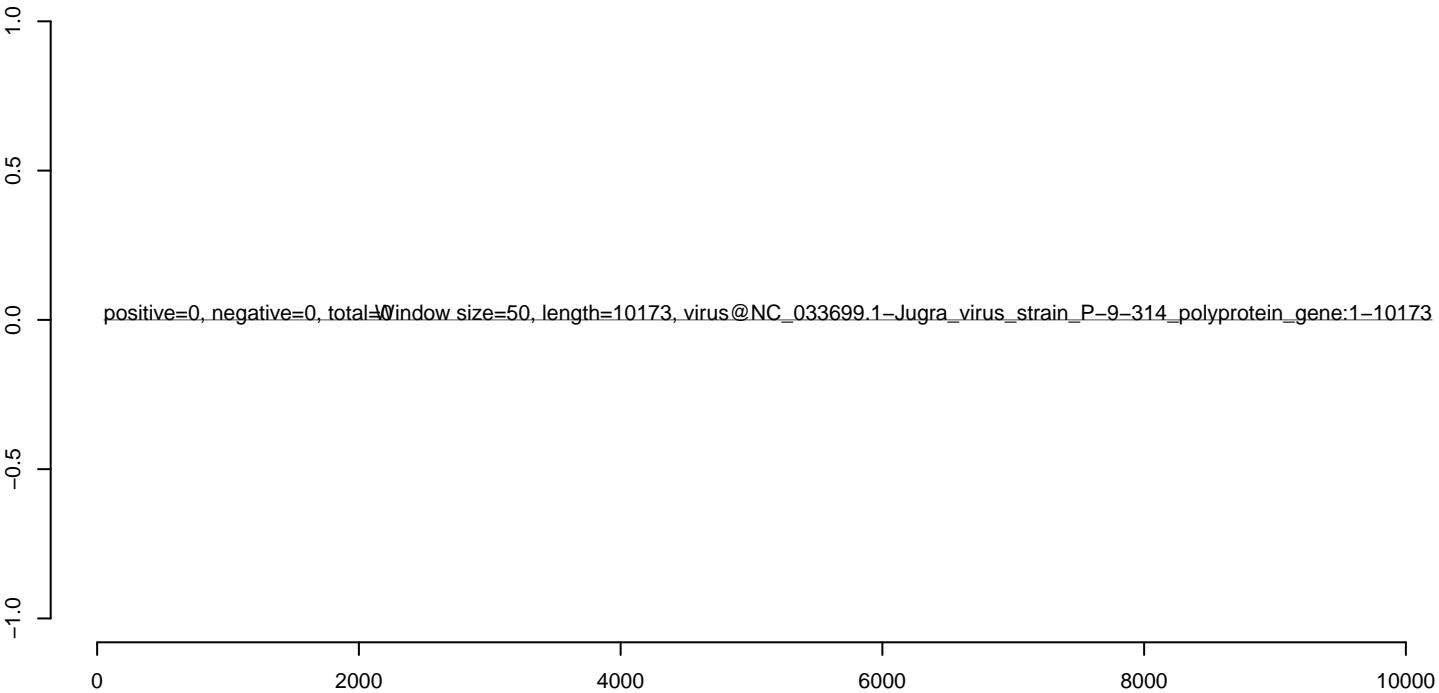
### AnGam\_Fcanc\_TN.18\_23.rep



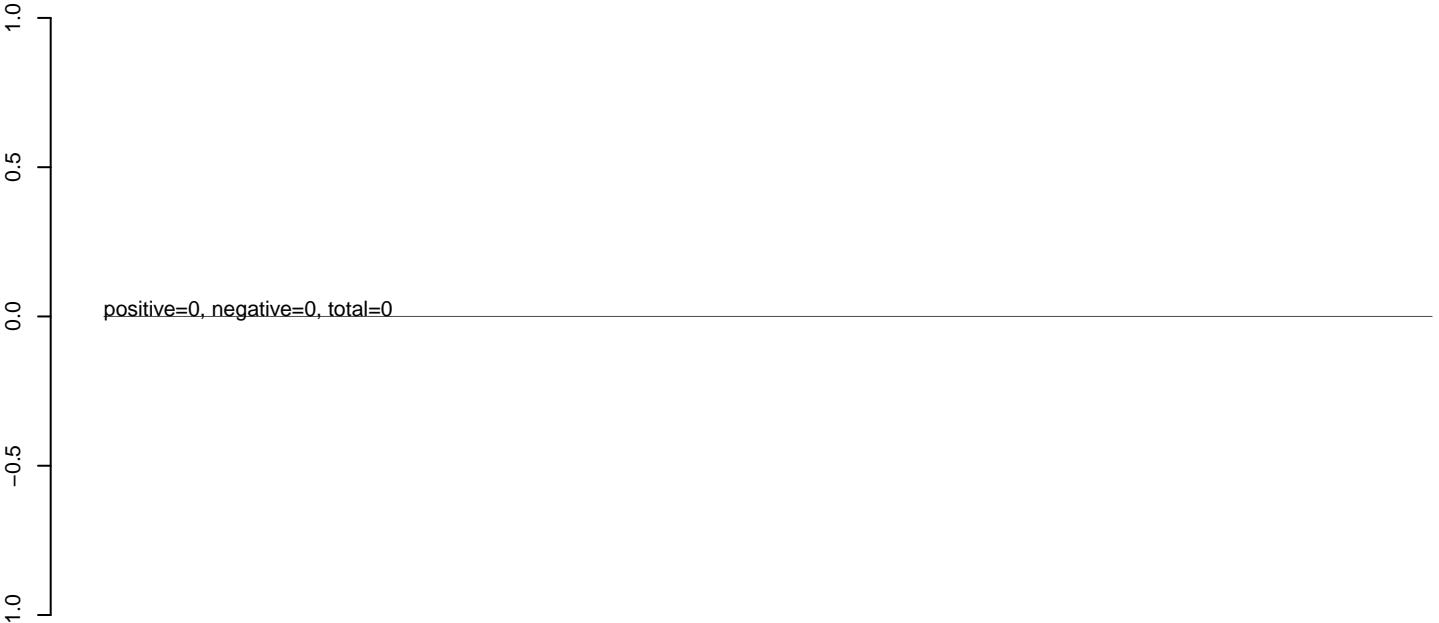
### AnGam\_Fcanc\_TN.24\_35.rep



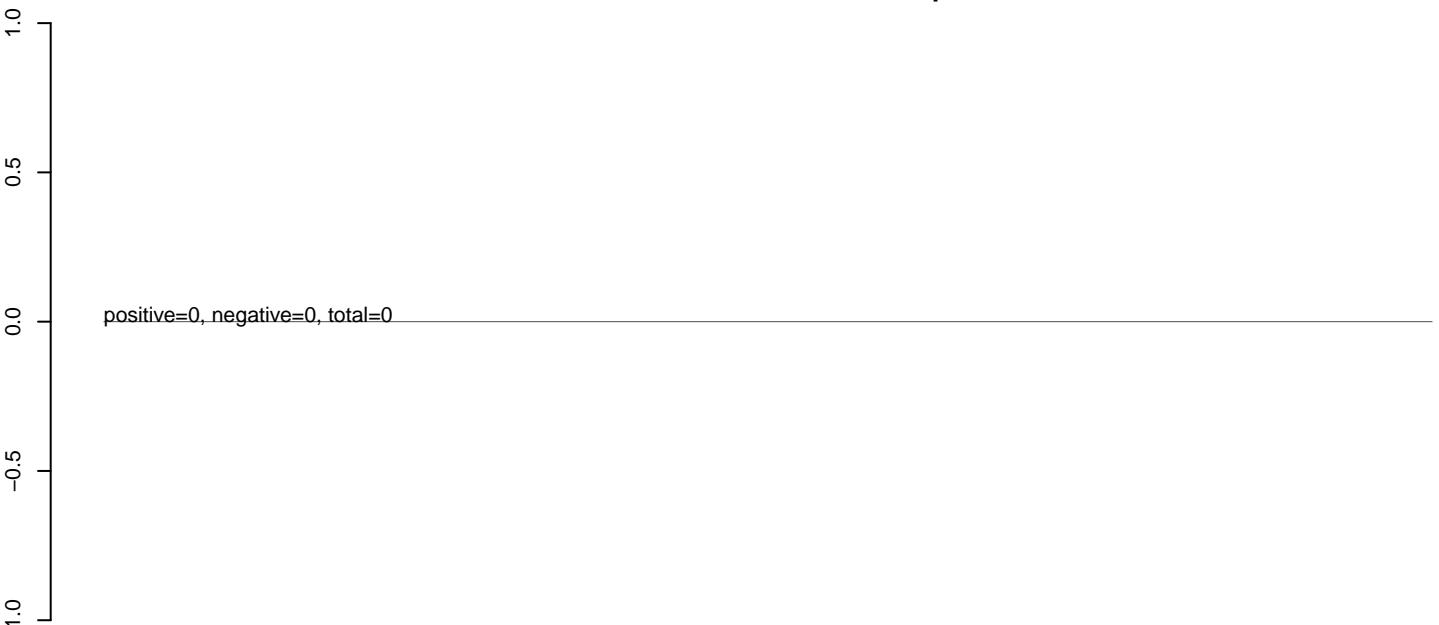
### AnGam\_Fcanc\_TN.rep



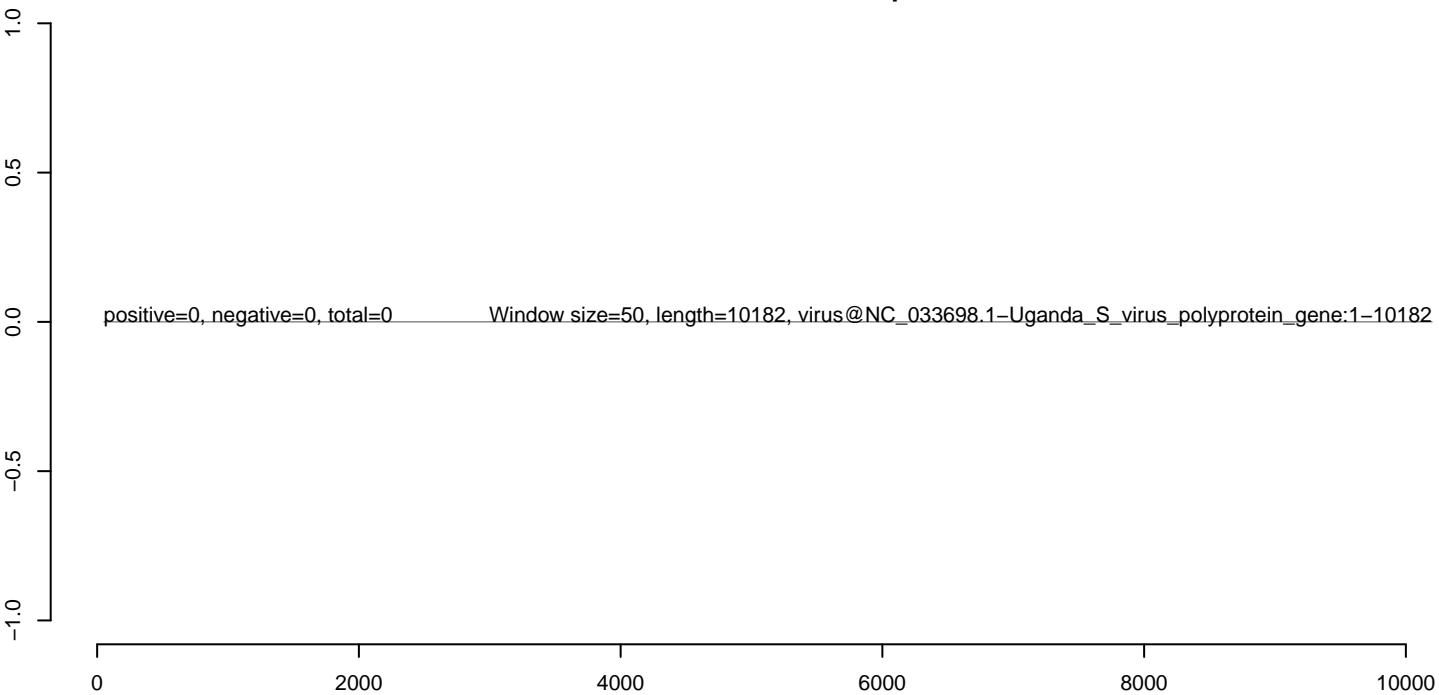
### AnGam\_Fcanc\_TN.18\_23.rep

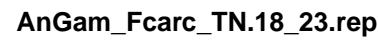


### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep





positive=0, negative=0, total=0

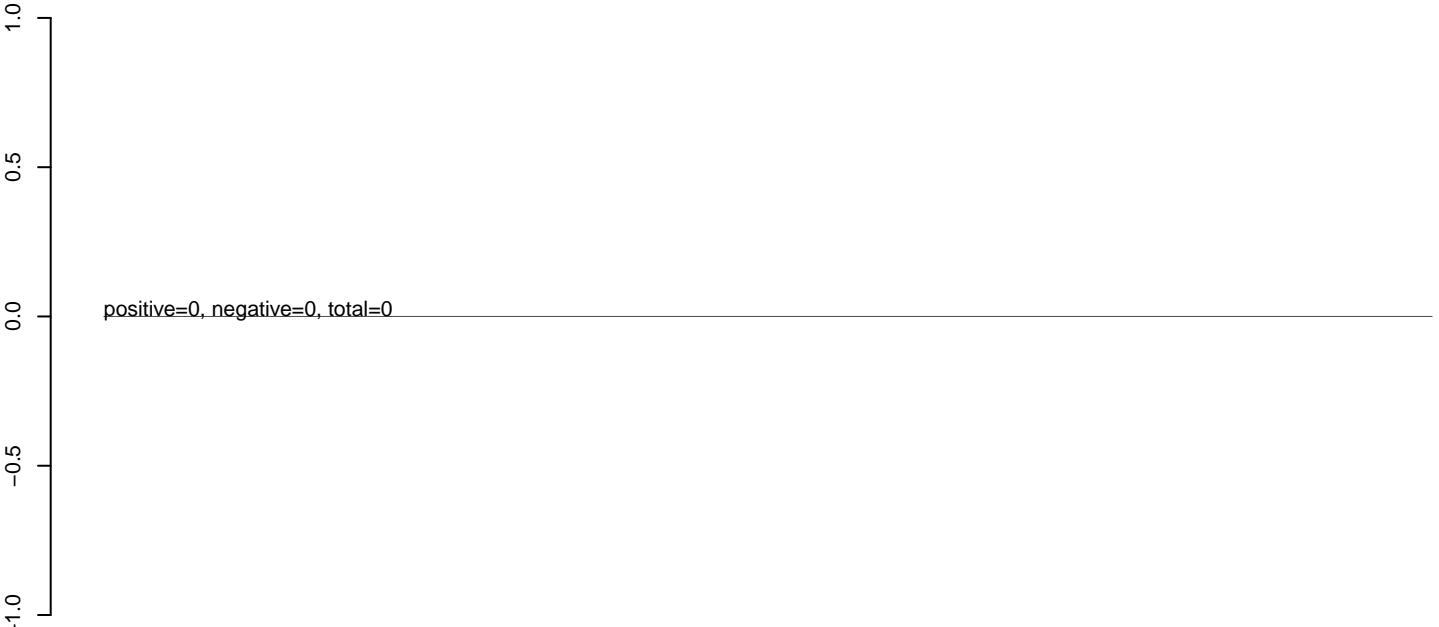


positive=0, negative=0, total=0

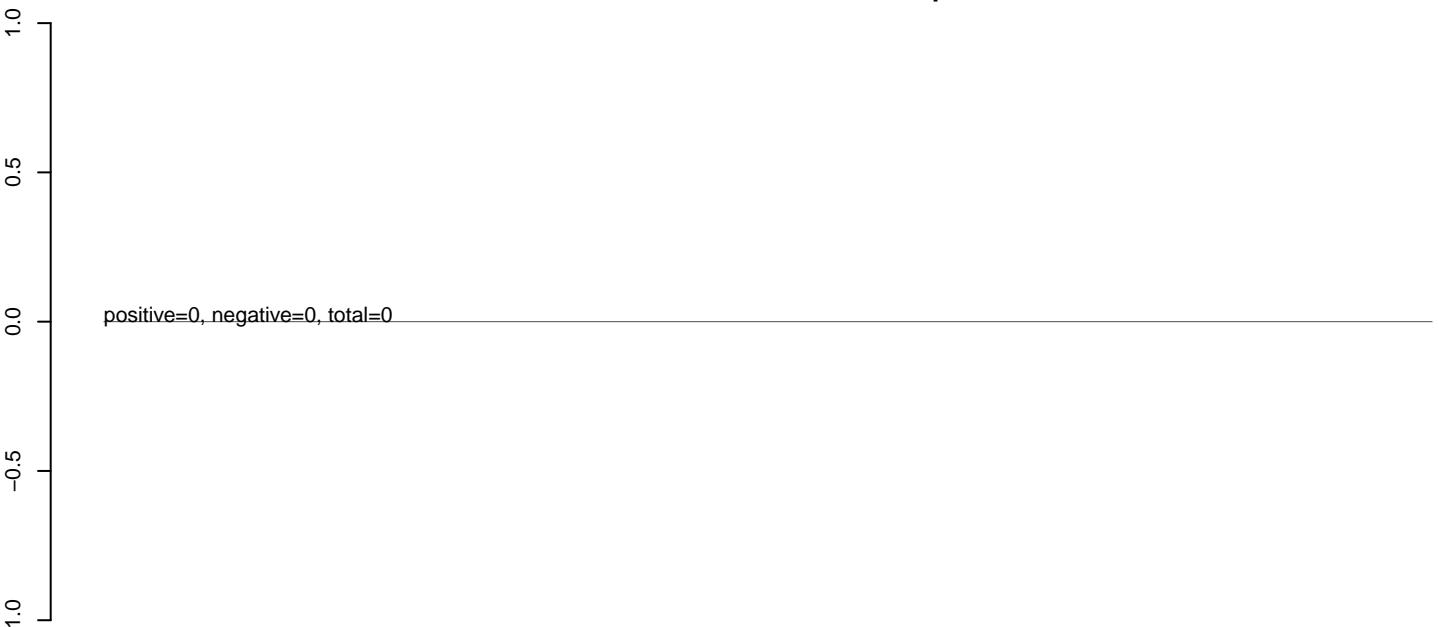


positive=0, negative=0, nontypeable=50, length=10173, virus@NC\_033697.1-Saboya\_virus strain Dak AR D4600 polyprotein gene:1-10173

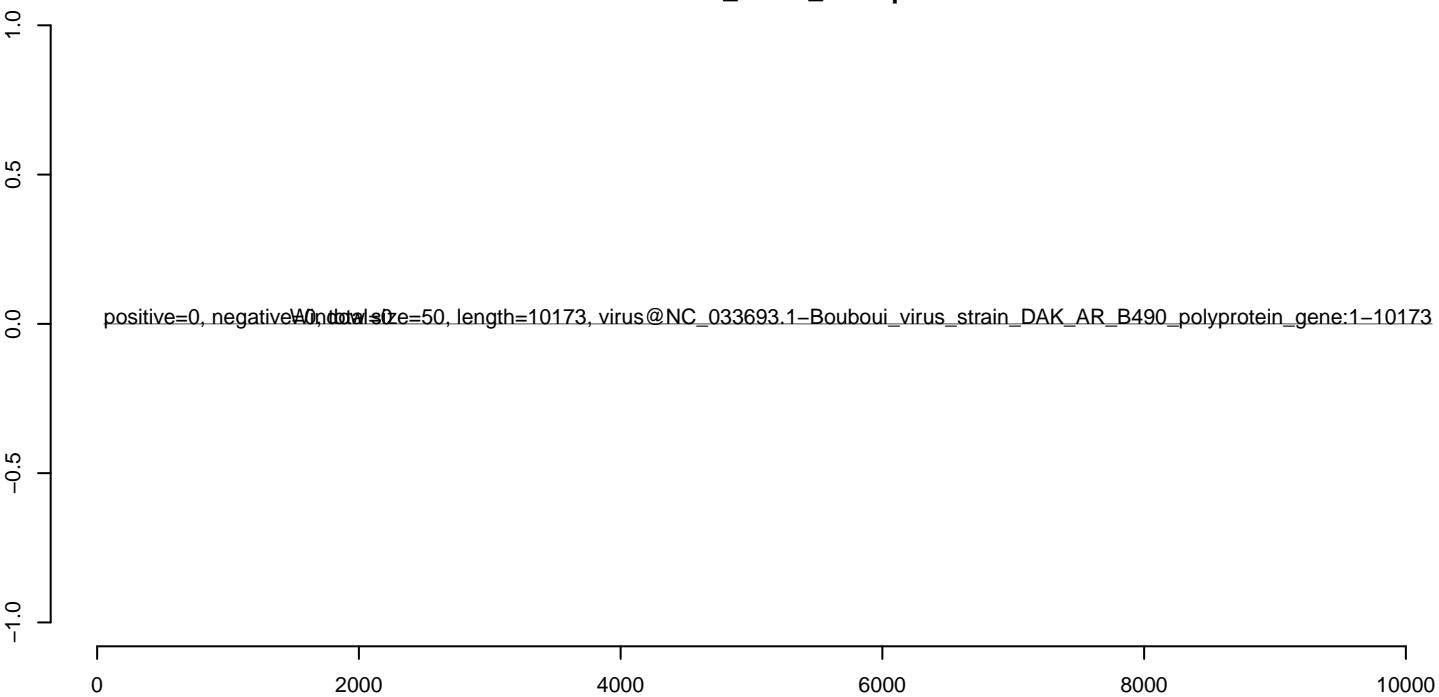
### AnGam\_Fcanc\_TN.18\_23.rep



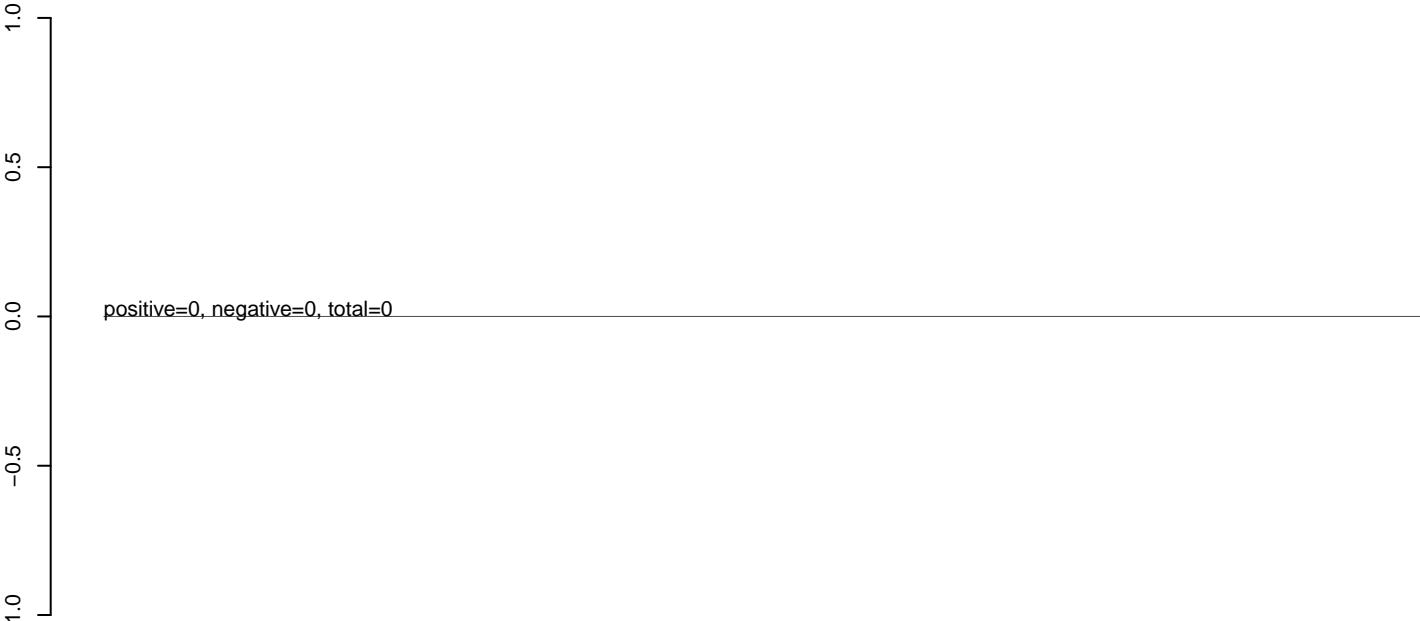
### AnGam\_Fcanc\_TN.24\_35.rep



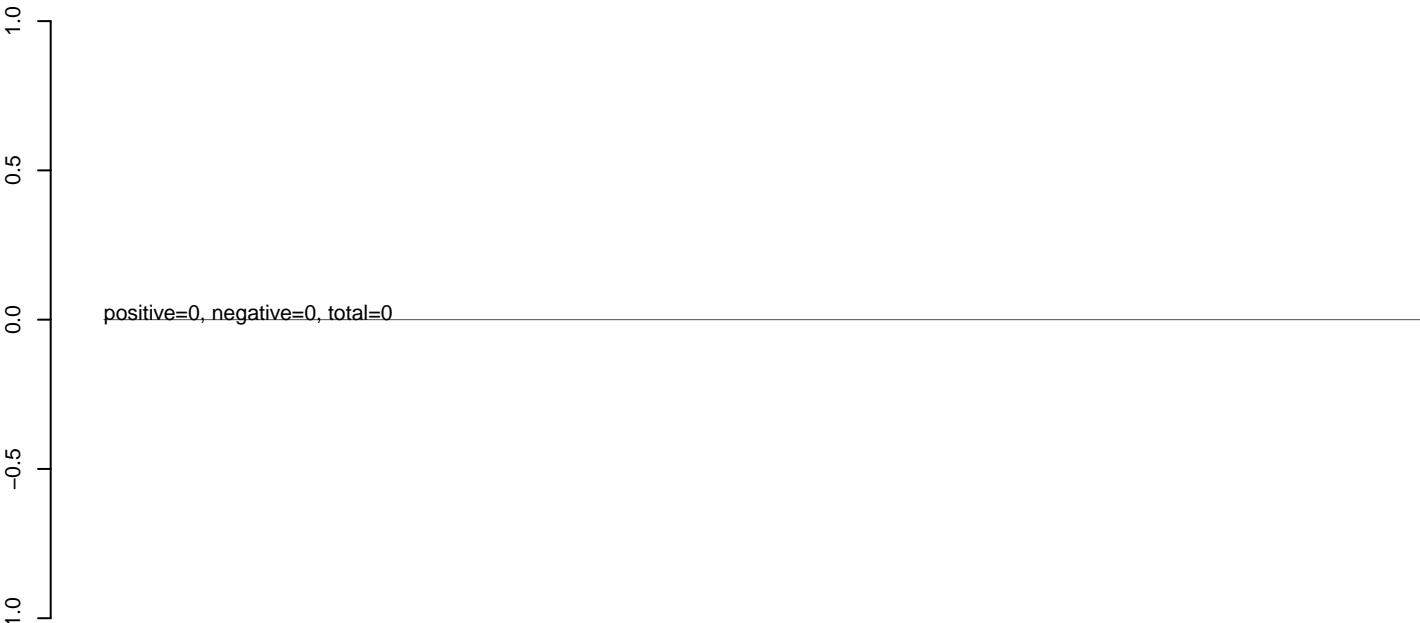
### AnGam\_Fcanc\_TN.rep



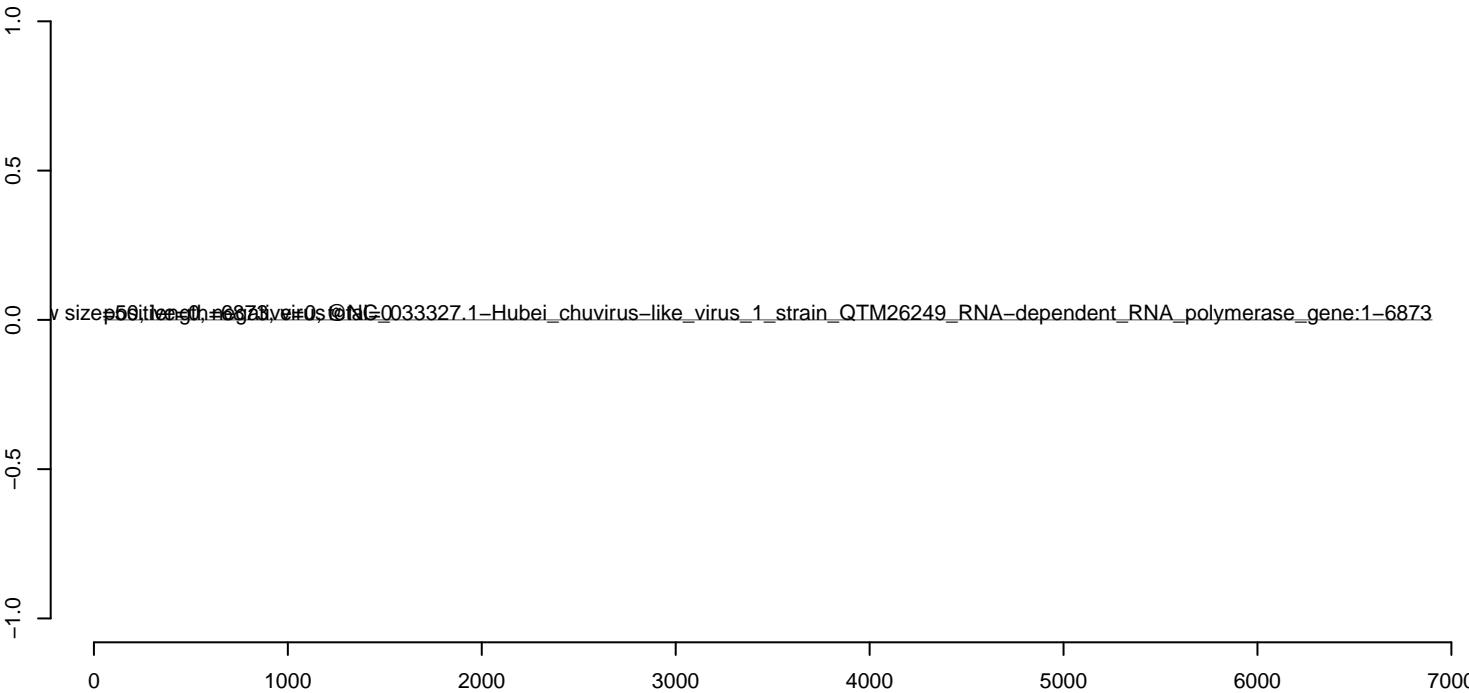
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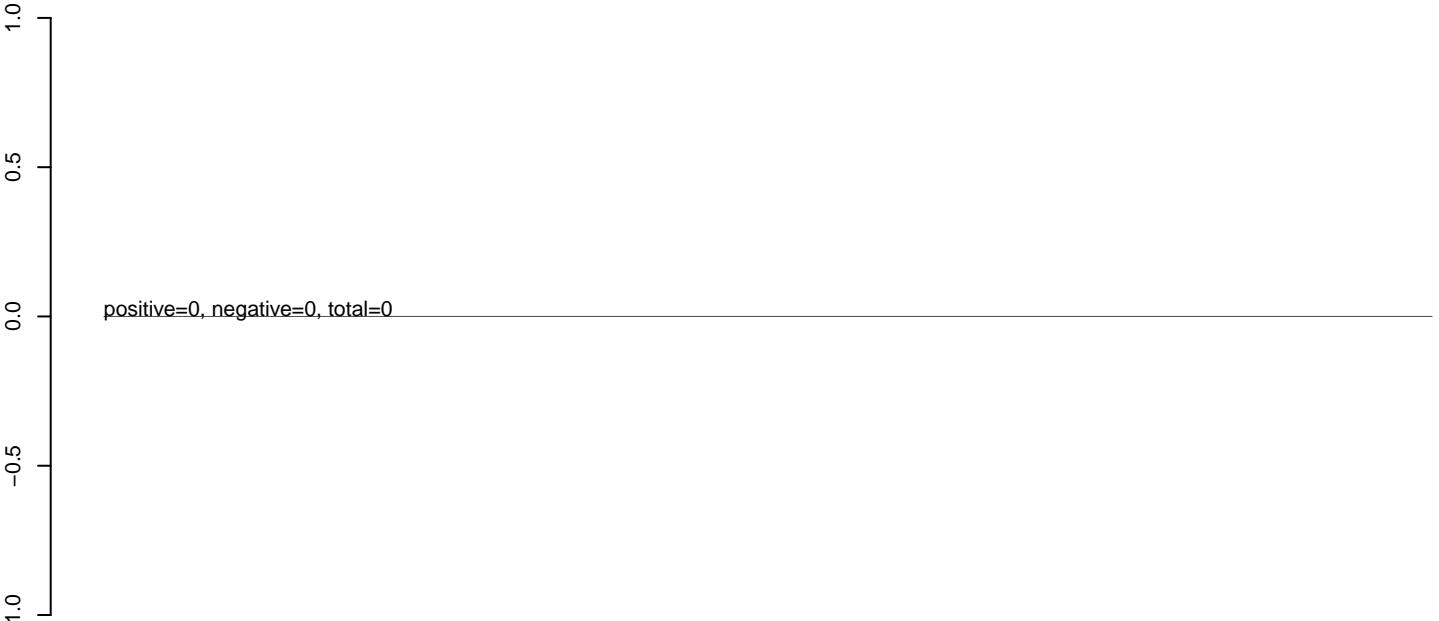
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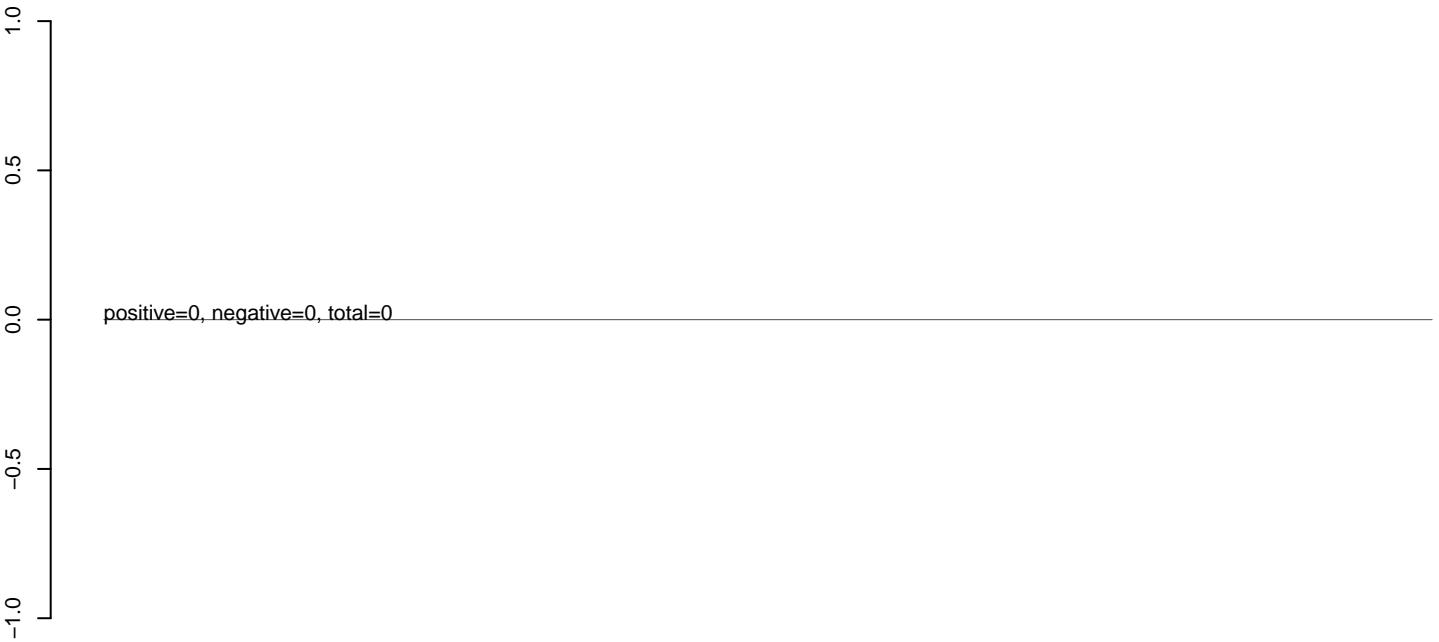
### AnGam\_Fcanc\_TN.rep



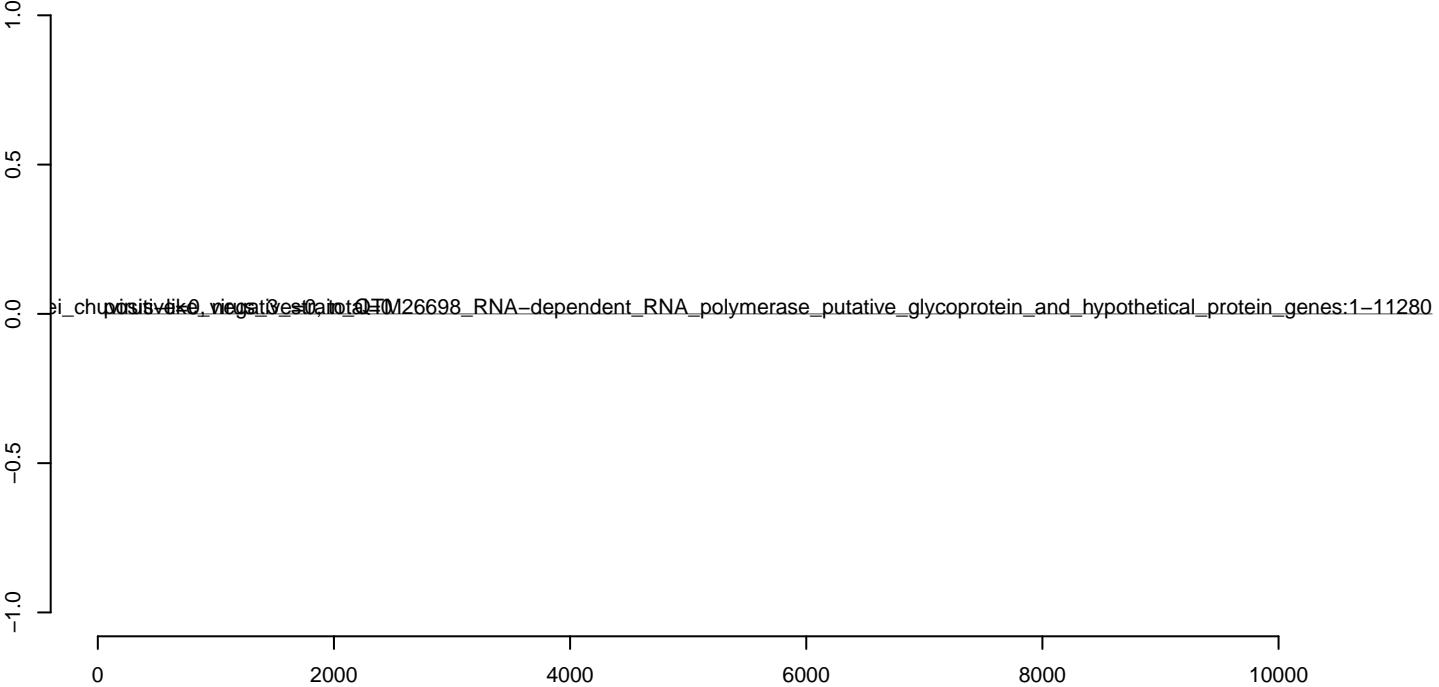
### AnGam\_Fcanc\_TN.18\_23.rep



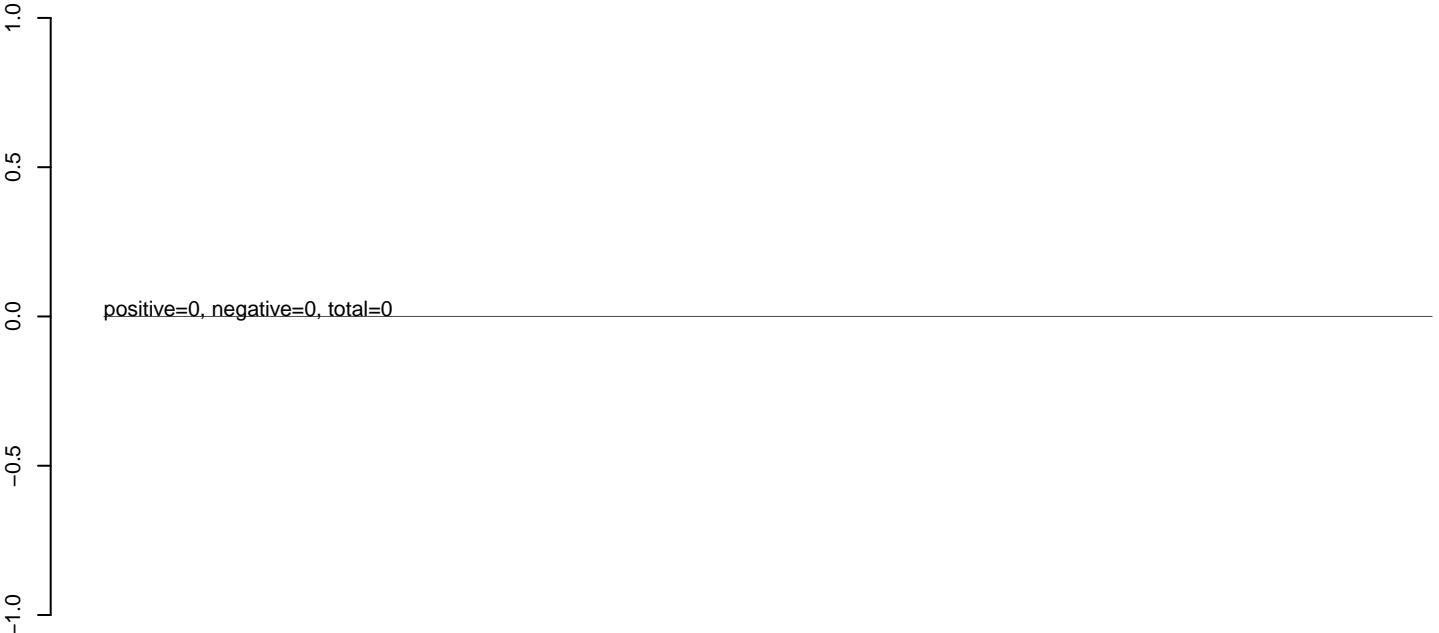
### AnGam\_Fcanc\_TN.24\_35.rep



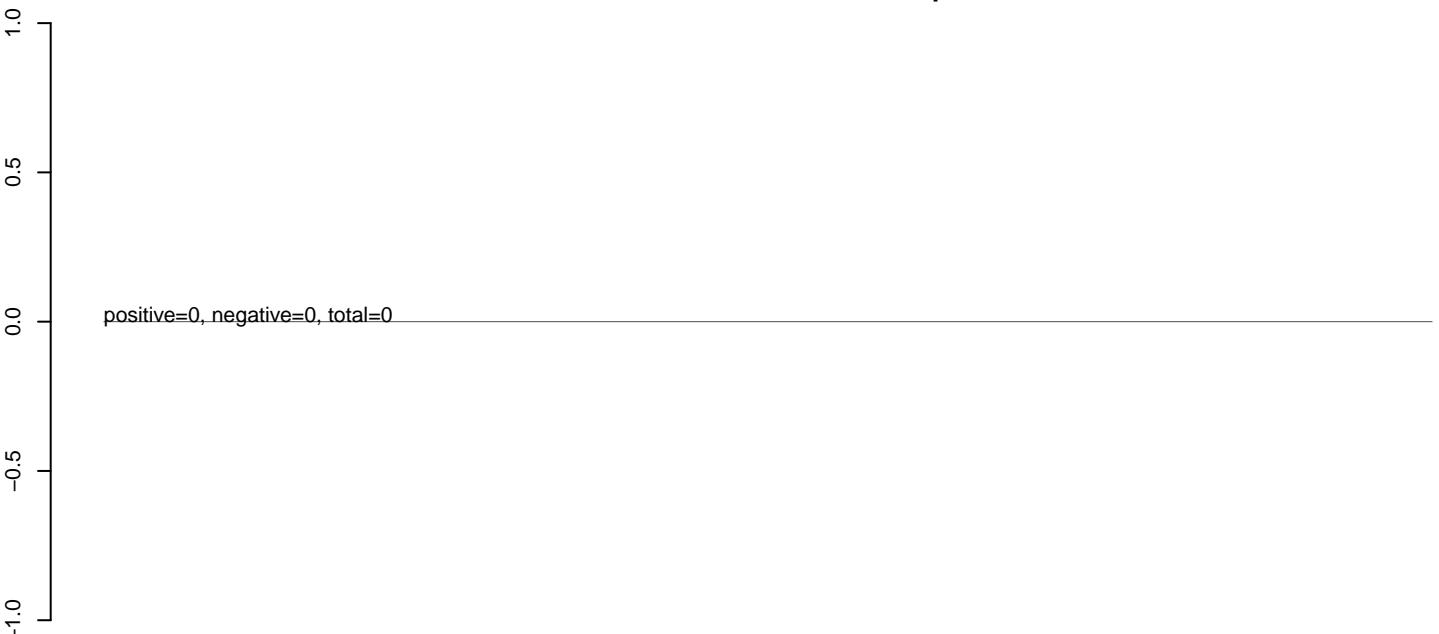
### AnGam\_Fcanc\_TN.rep



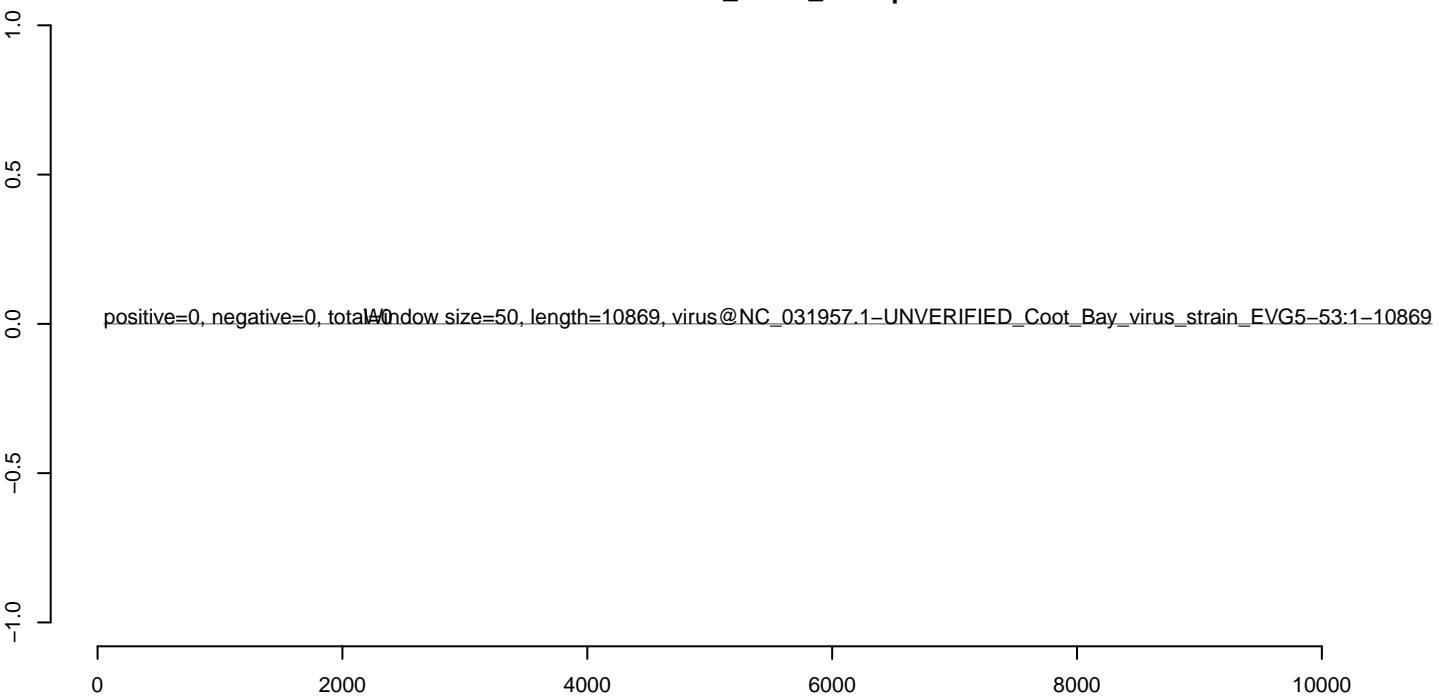
### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep



### AnGam\_Fcarc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0 Window.size=50, length=11286, virus@NC\_031956.1-UNVERIFIED\_Balsa\_virus\_strain\_CoB\_84:1-11286

0 2000 4000 6000 8000 10000

### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

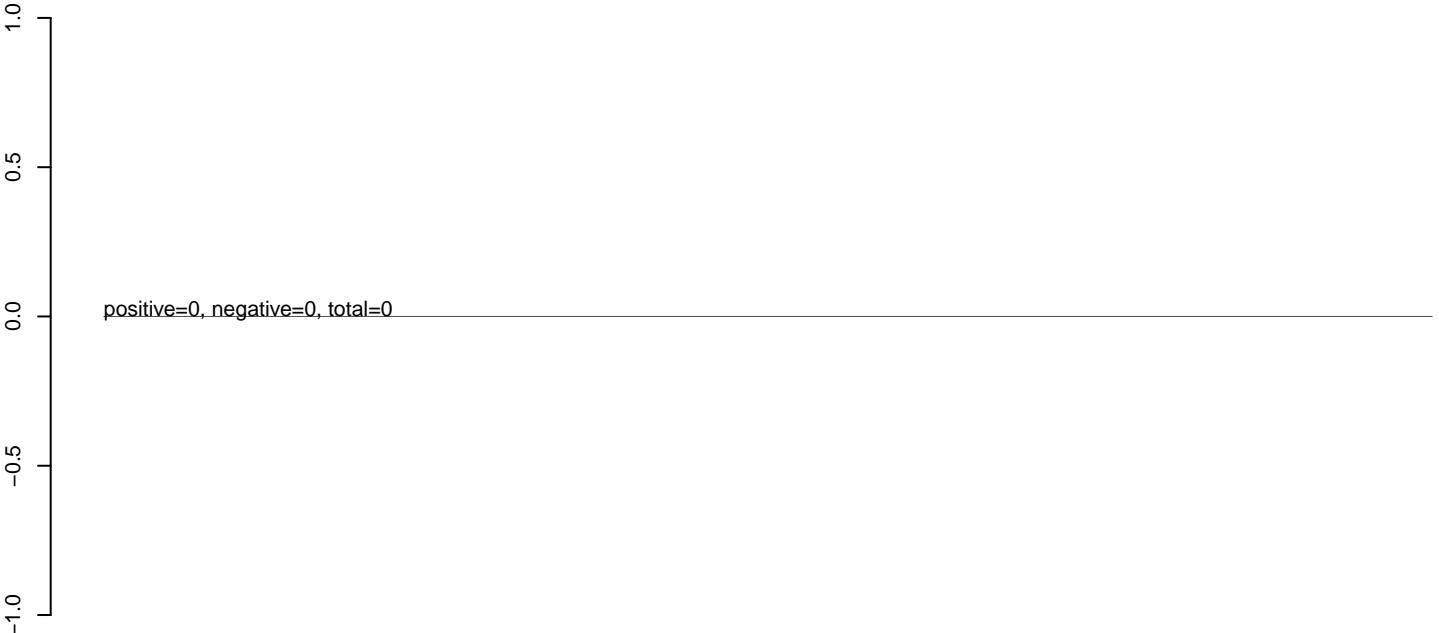
positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.rep

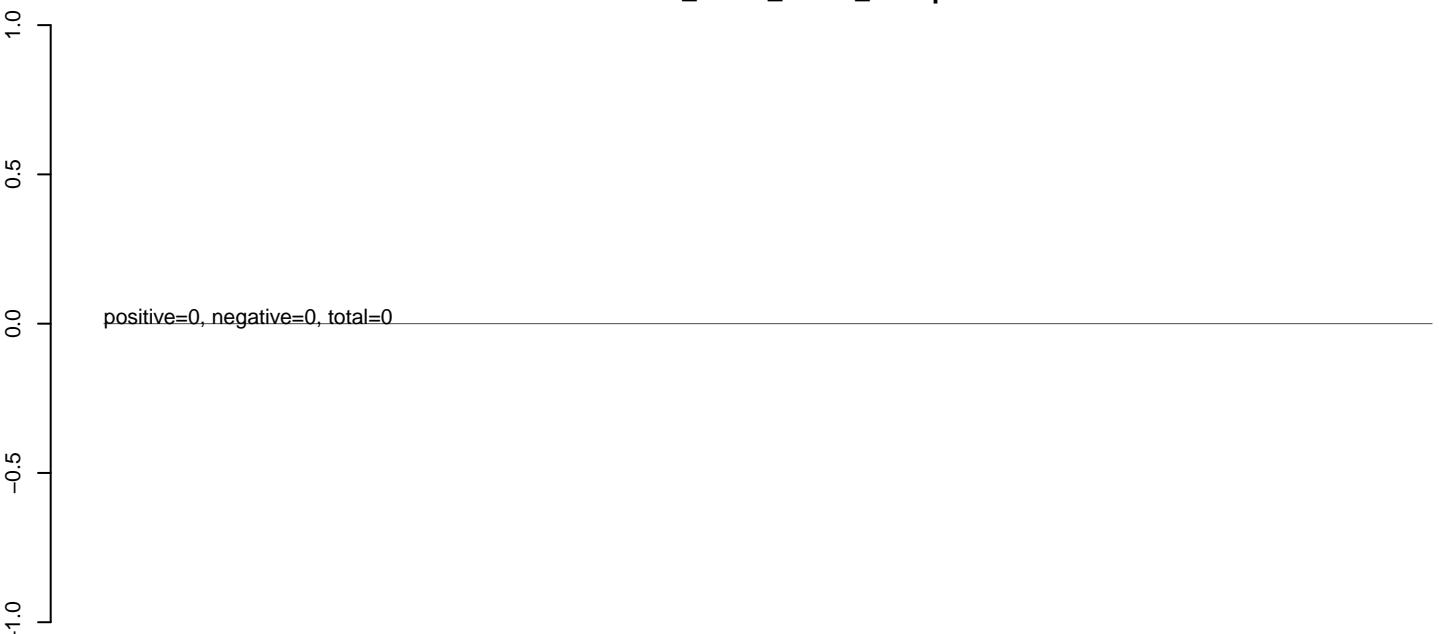
positive=0, negative=0, total=0

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

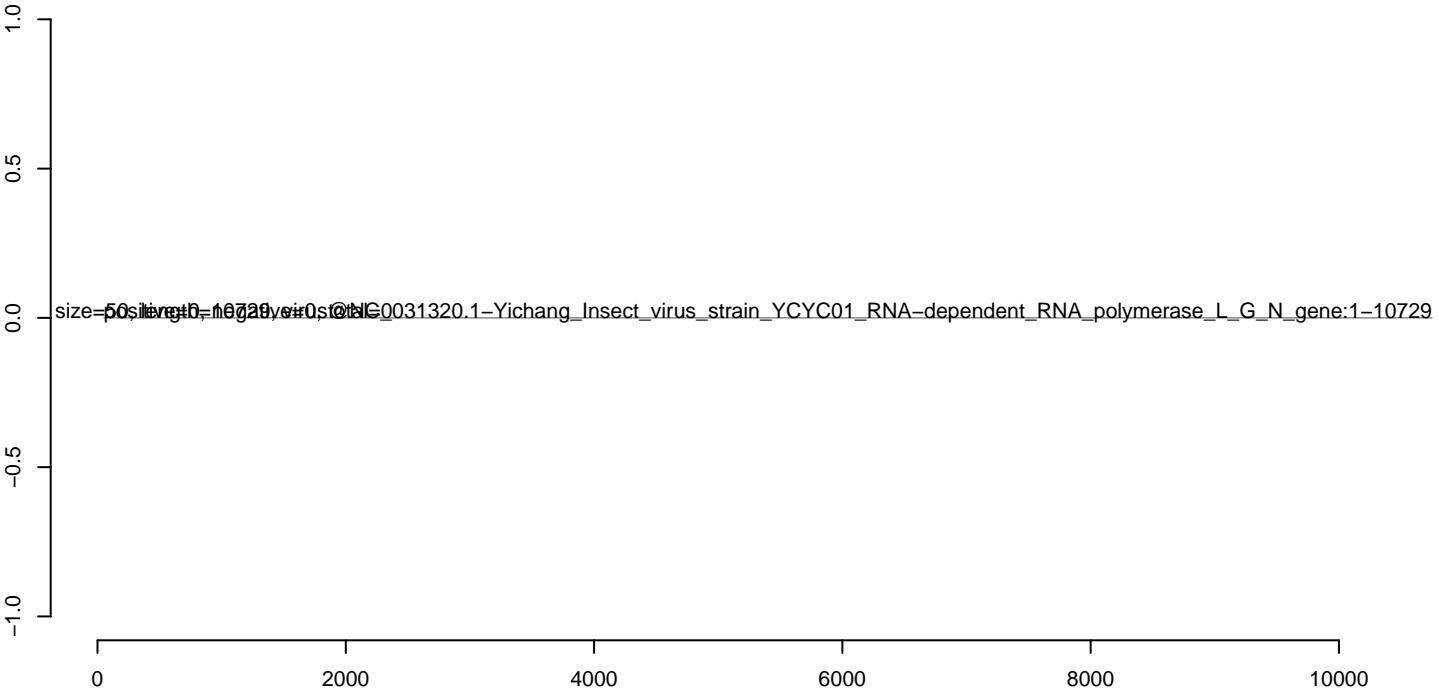
### AnGam\_Fcanc\_TN.18\_23.rep



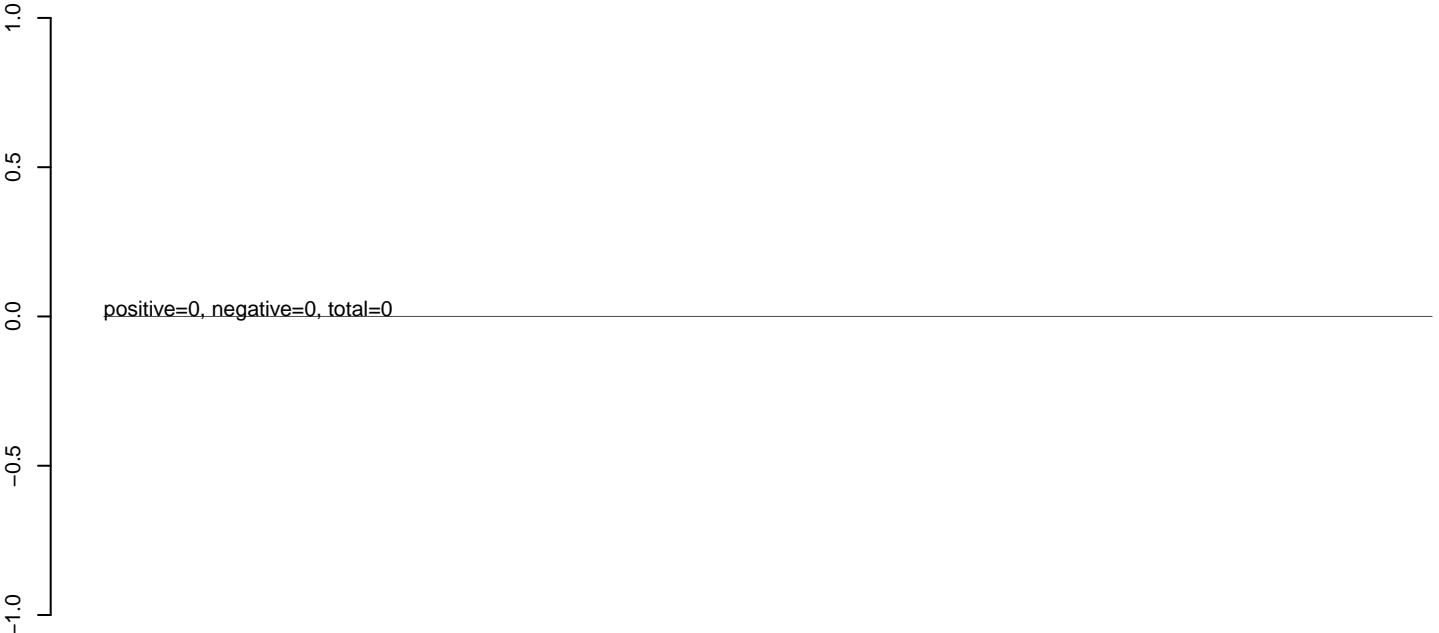
### AnGam\_Fcanc\_TN.24\_35.rep



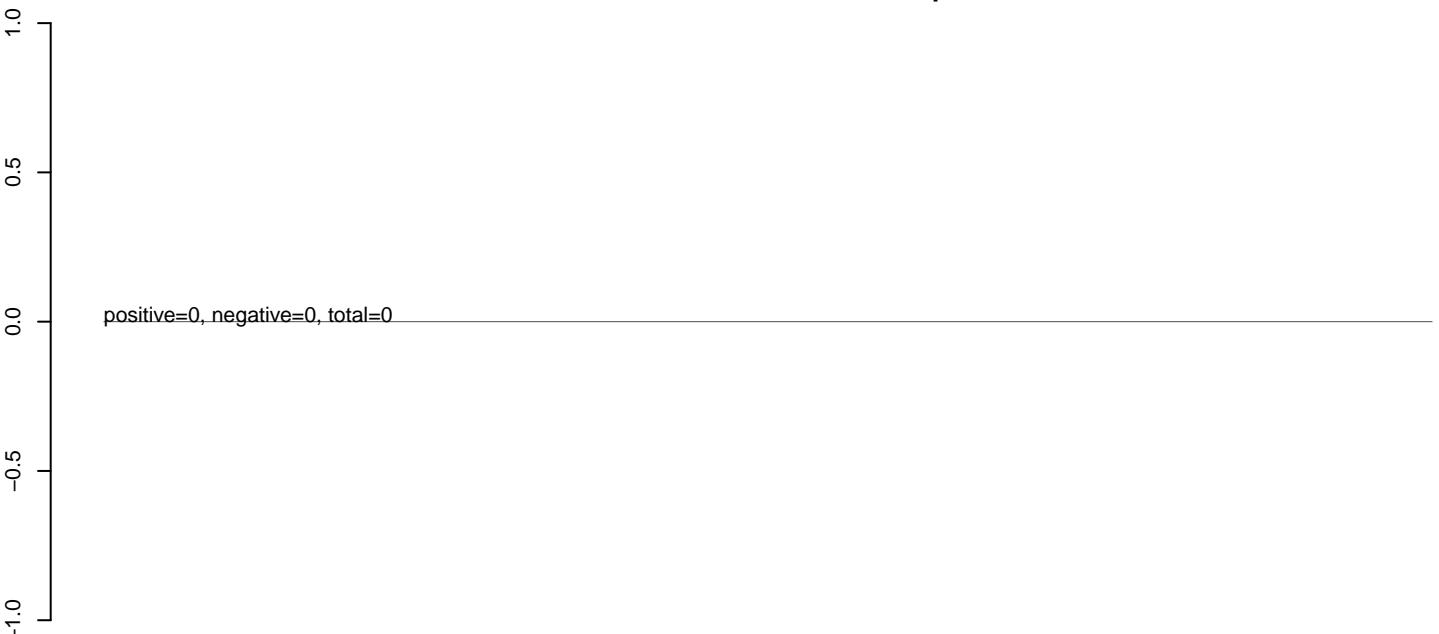
### AnGam\_Fcanc\_TN.rep



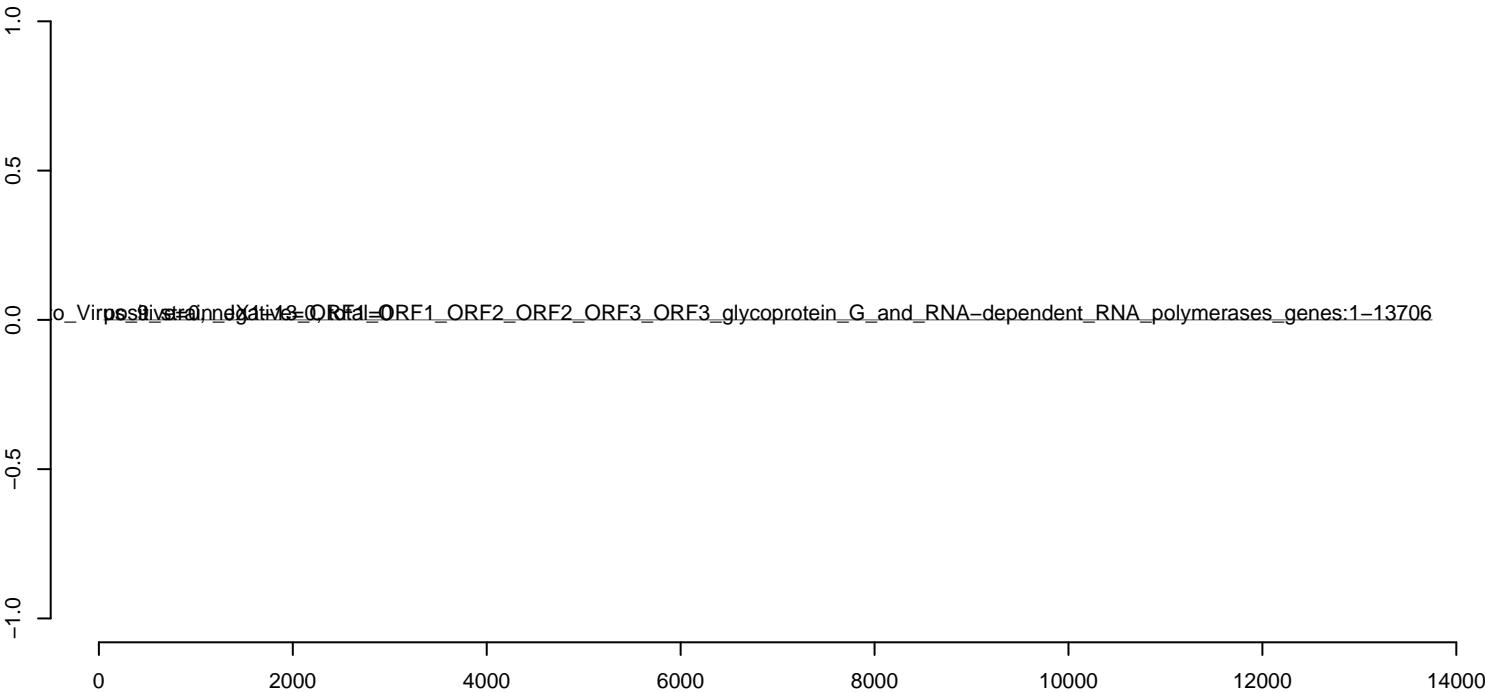
### AnGam\_Fcanc\_TN.18\_23.rep



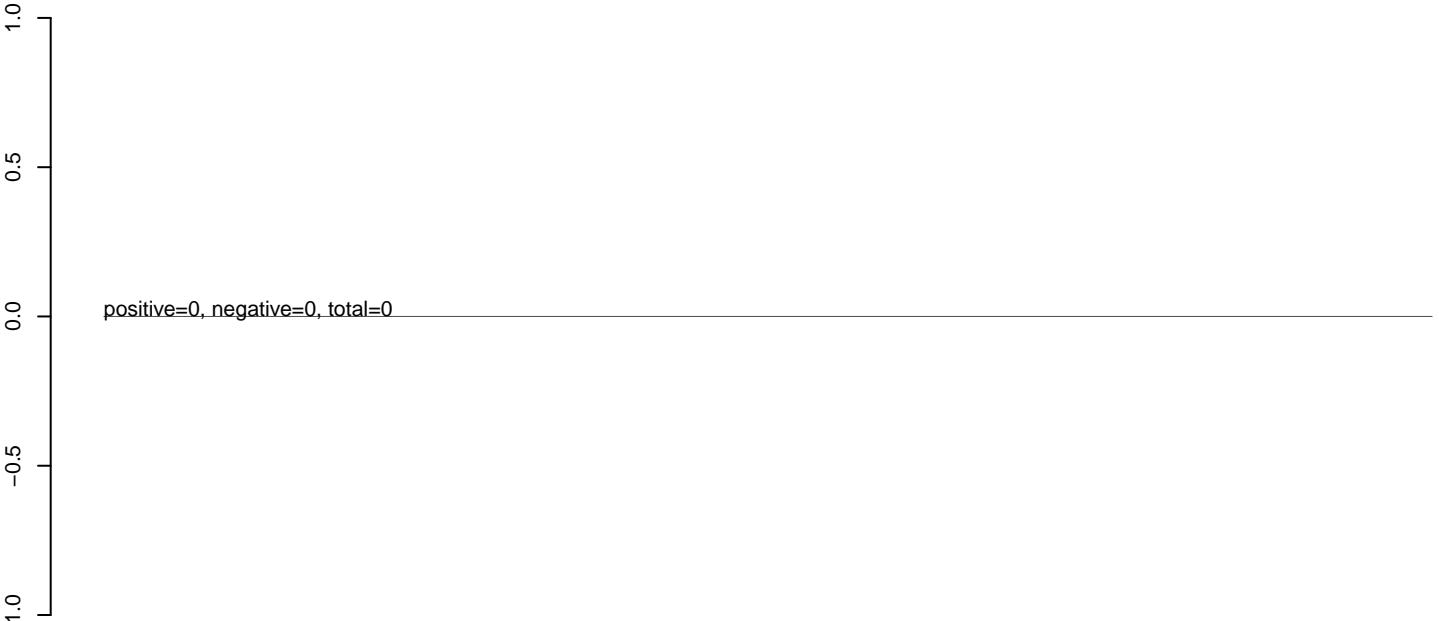
### AnGam\_Fcanc\_TN.24\_35.rep



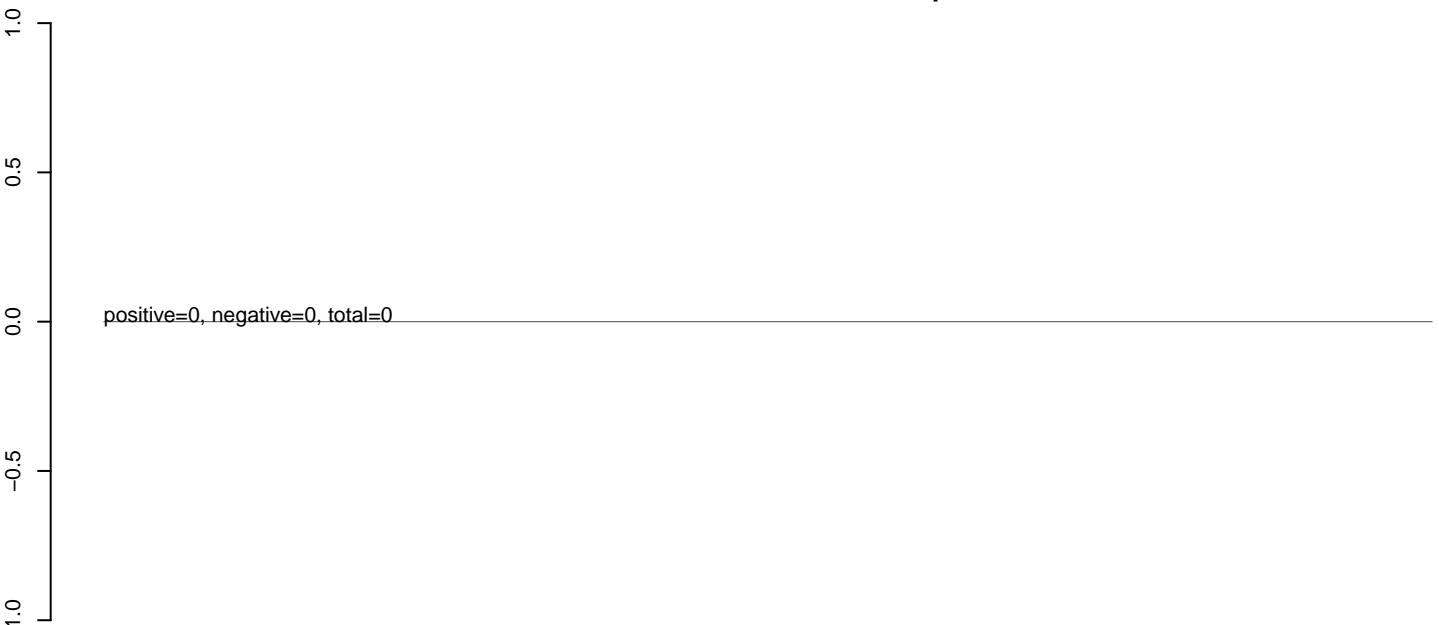
### AnGam\_Fcanc\_TN.rep



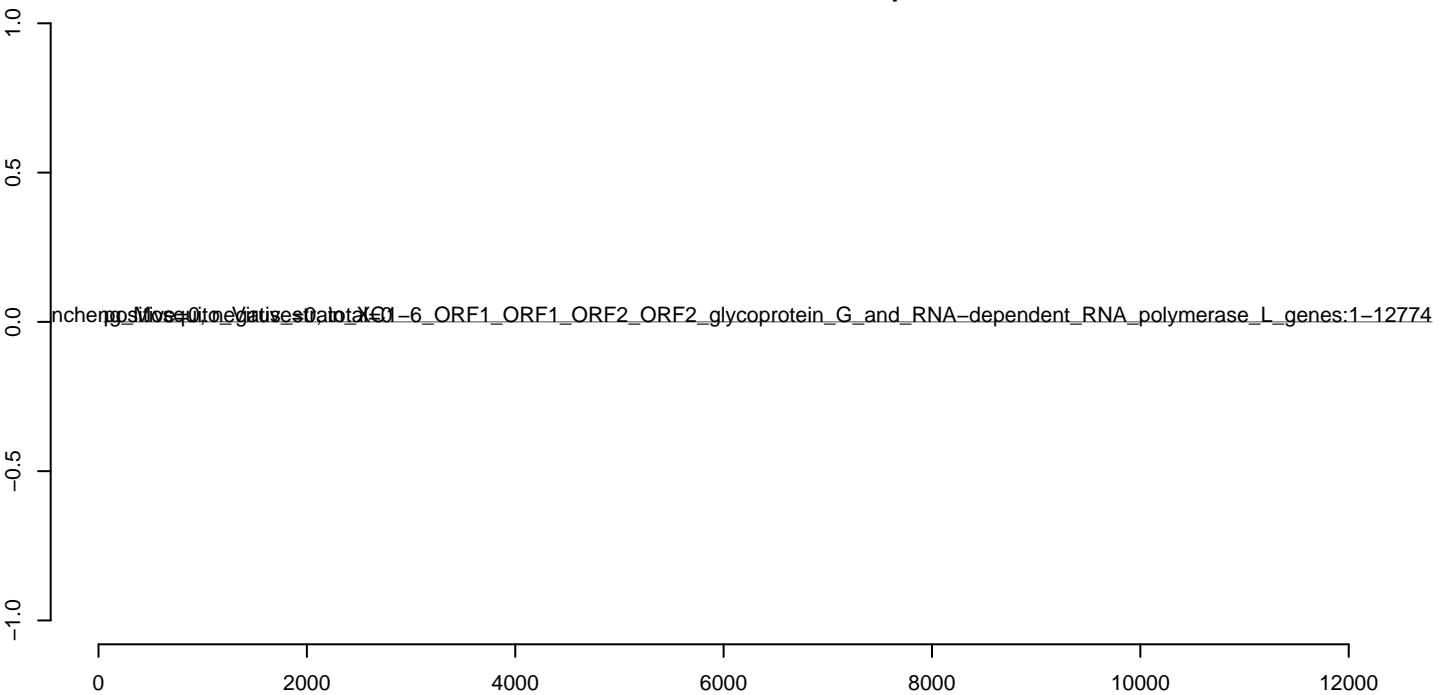
### AnGam\_Fcanc\_TN.18\_23.rep



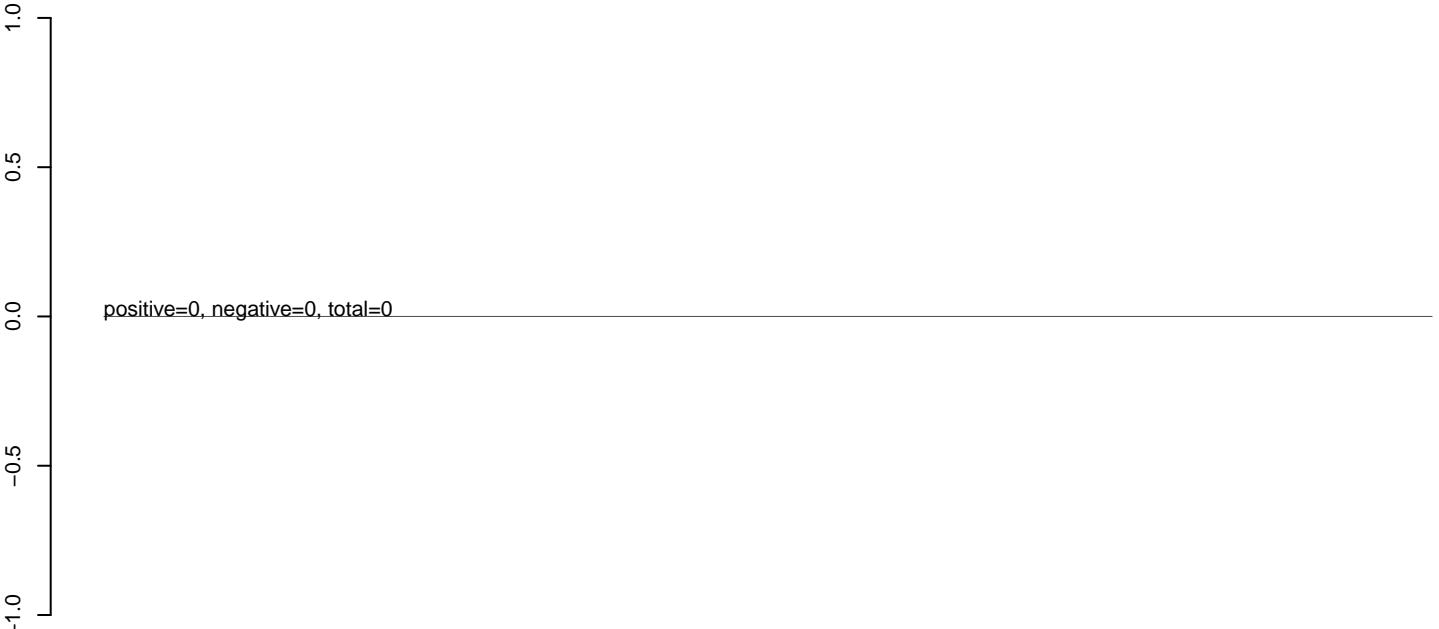
### AnGam\_Fcanc\_TN.24\_35.rep



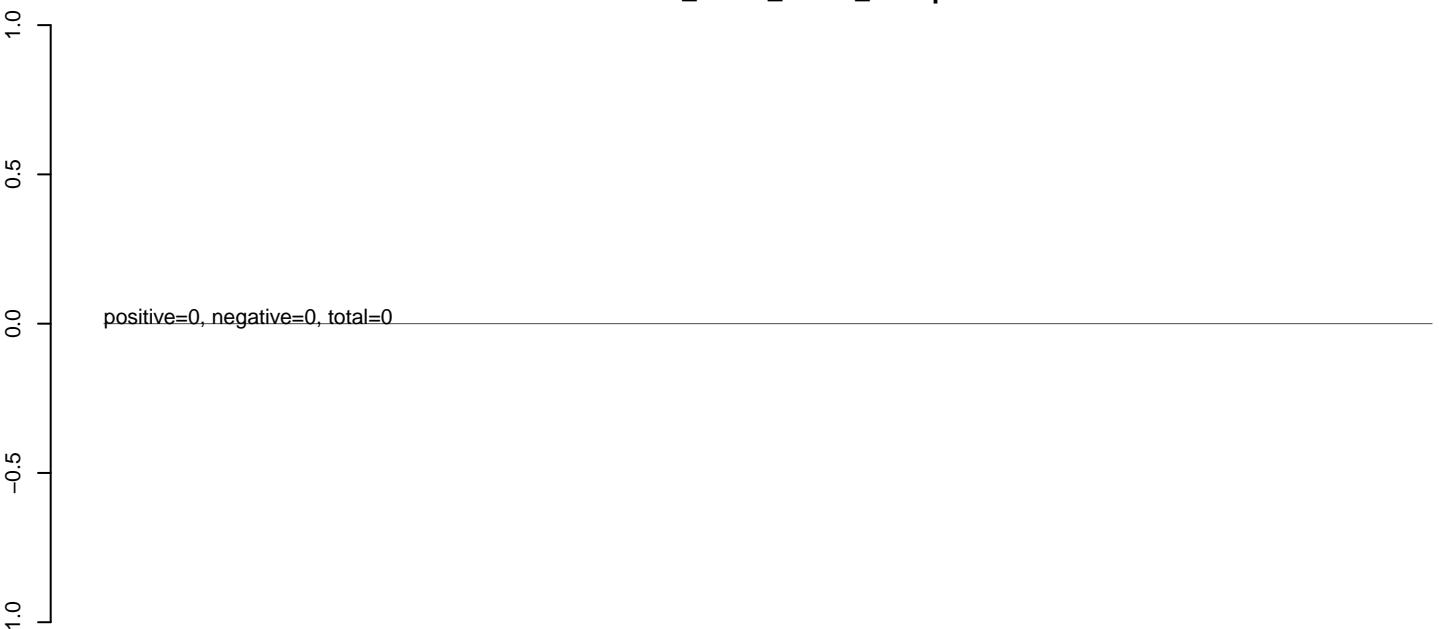
### AnGam\_Fcanc\_TN.rep



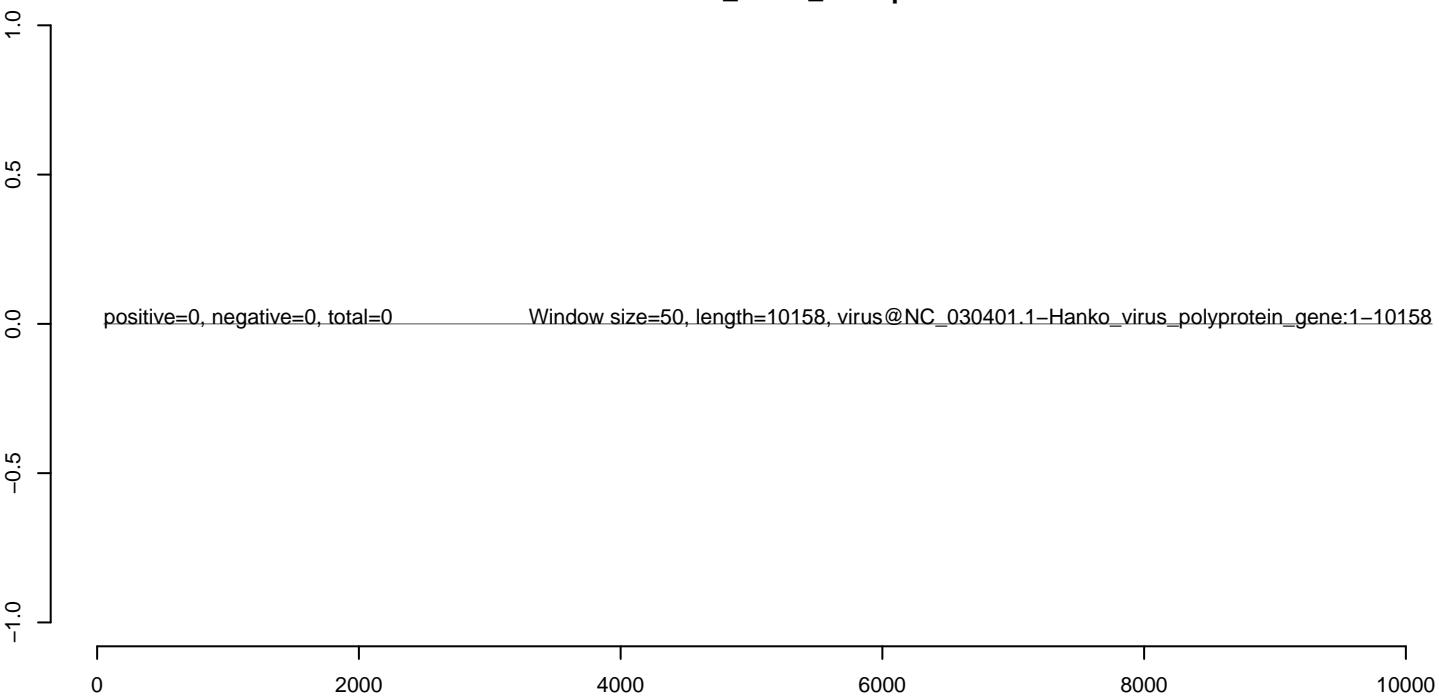
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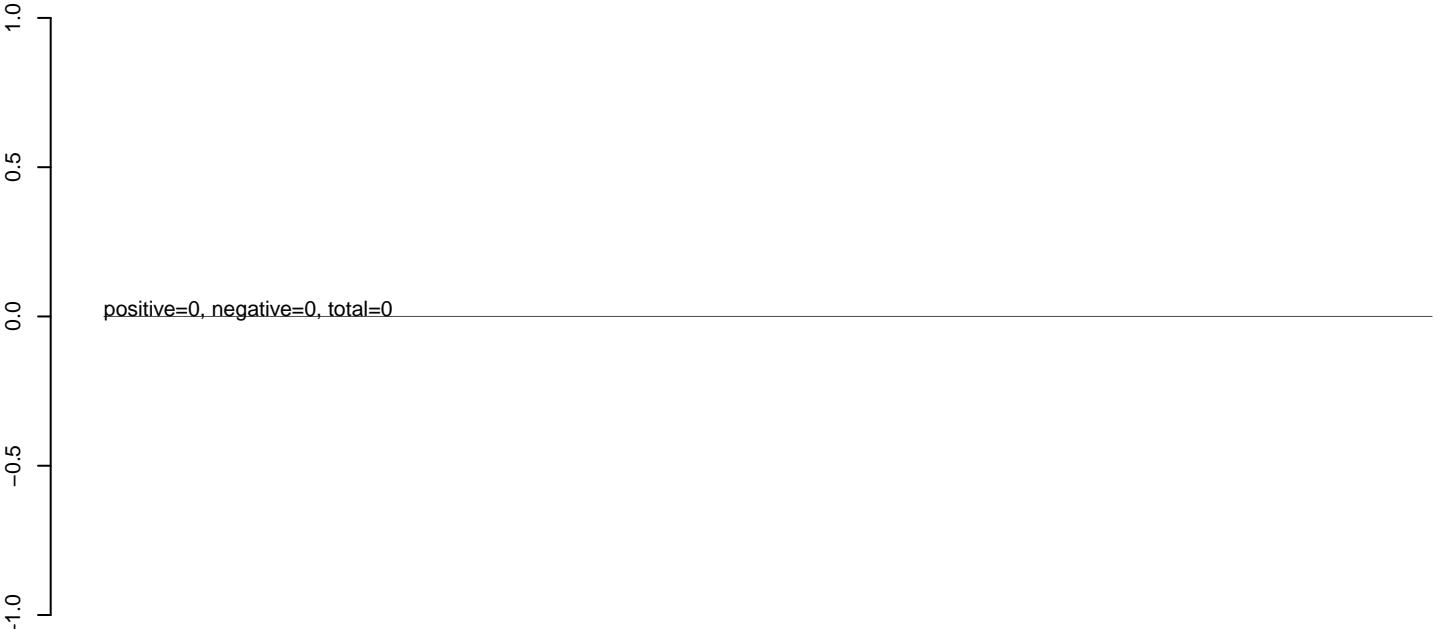
### AnGam\_Fcanc\_TN.24\_35.rep



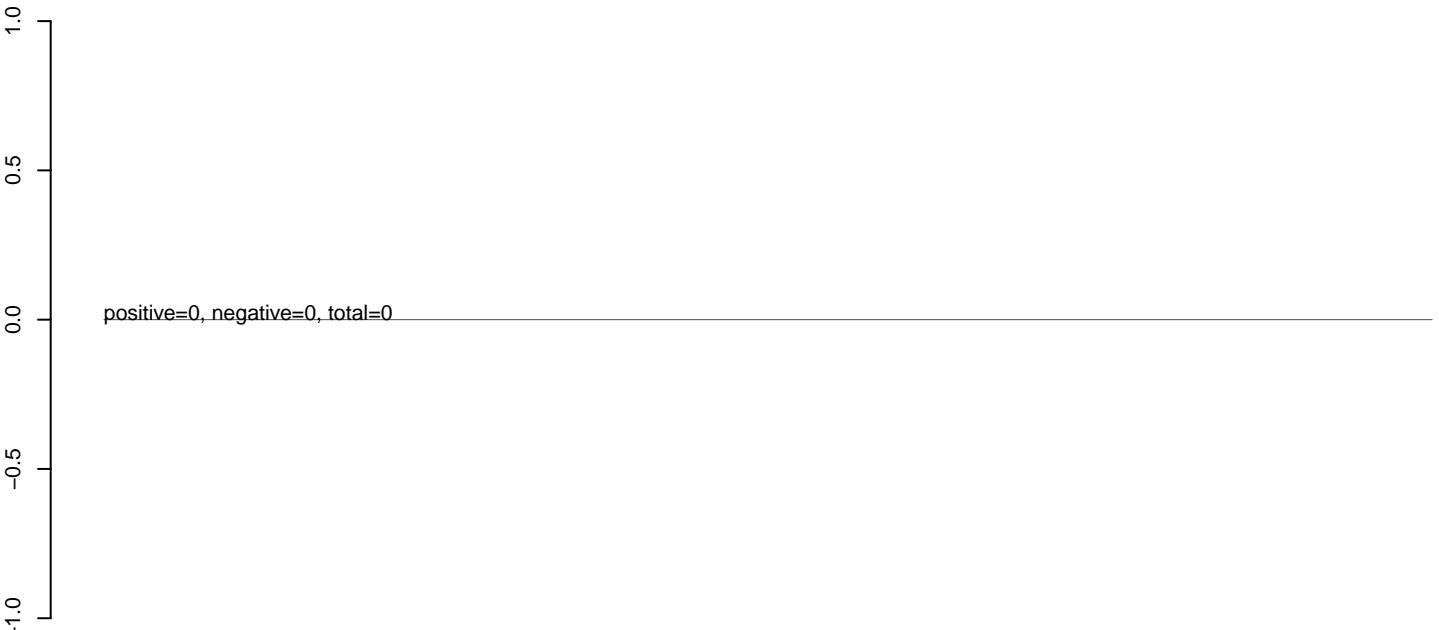
### AnGam\_Fcanc\_TN.rep



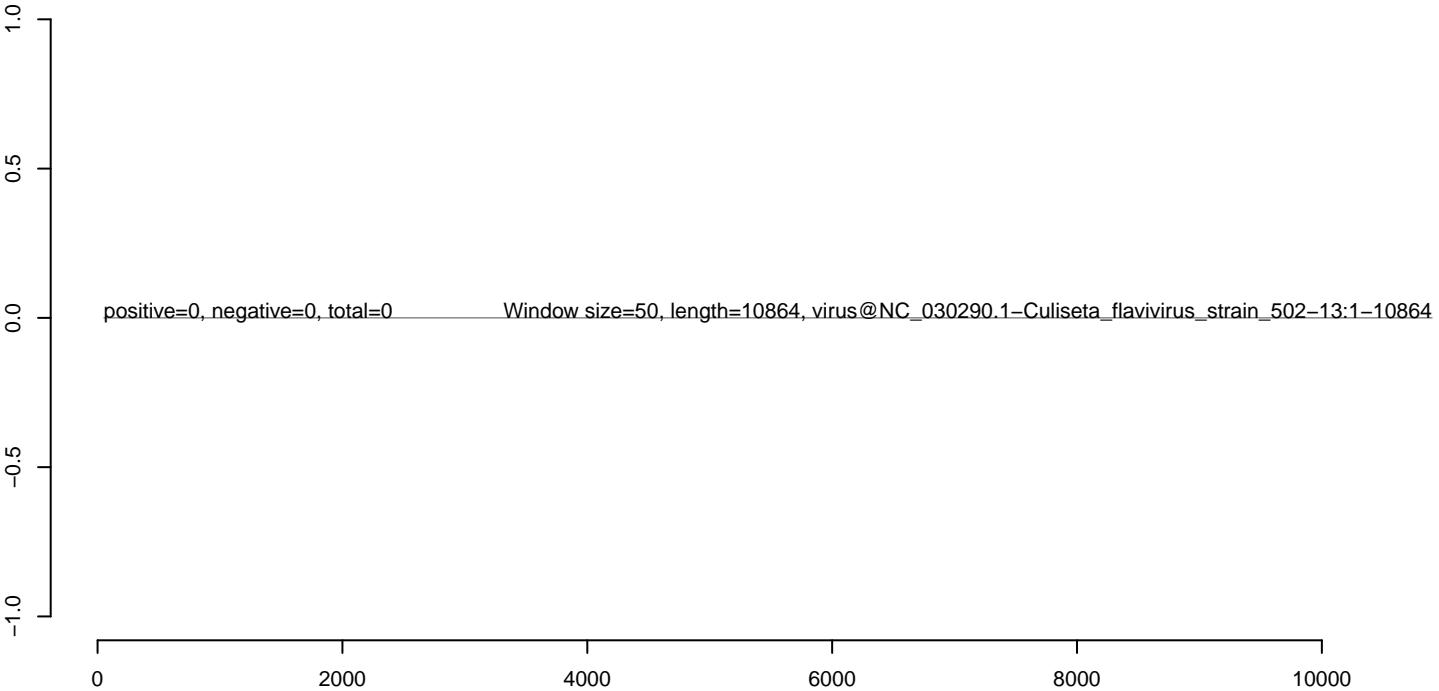
### AnGam\_Fcanc\_TN.18\_23.rep



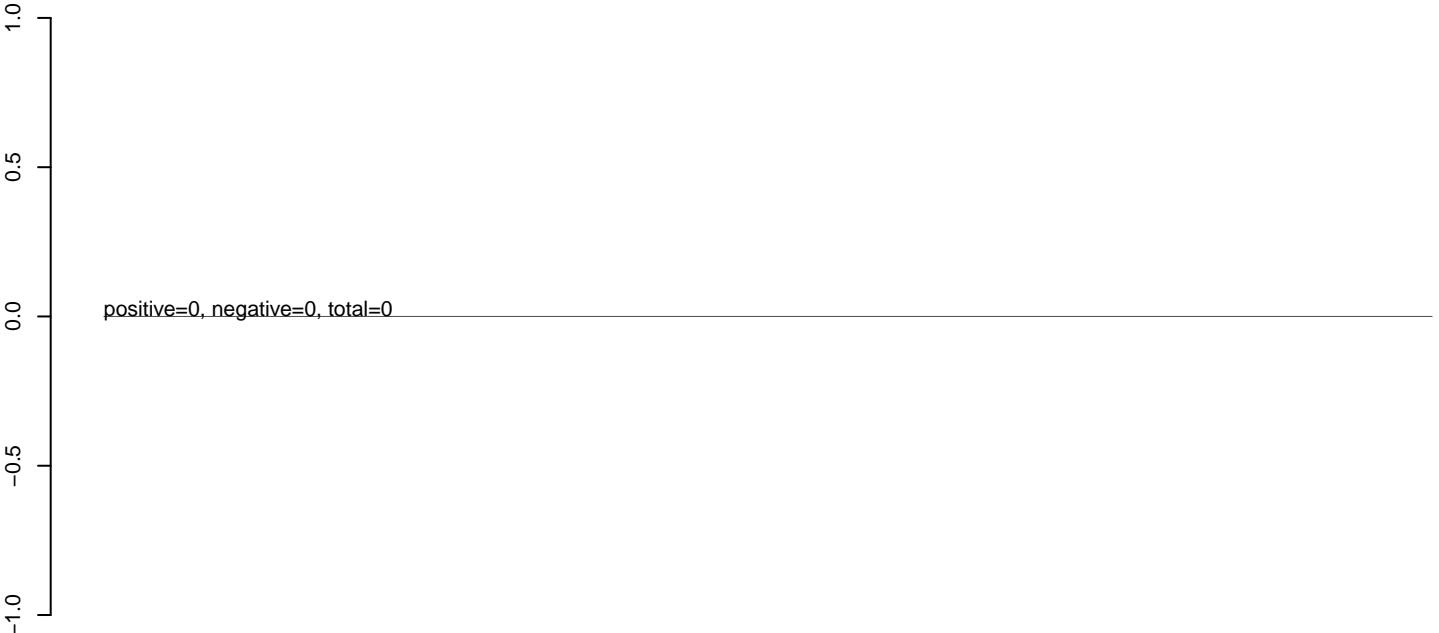
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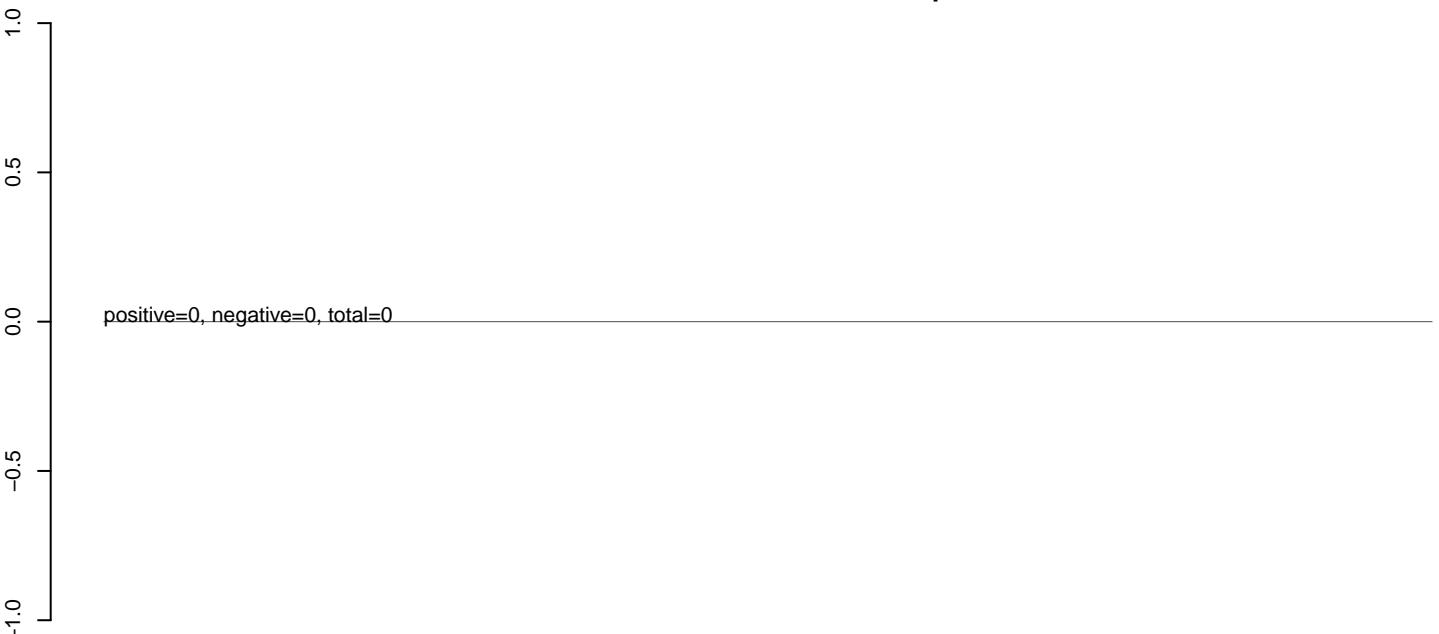
### AnGam\_Fcanc\_TN.rep



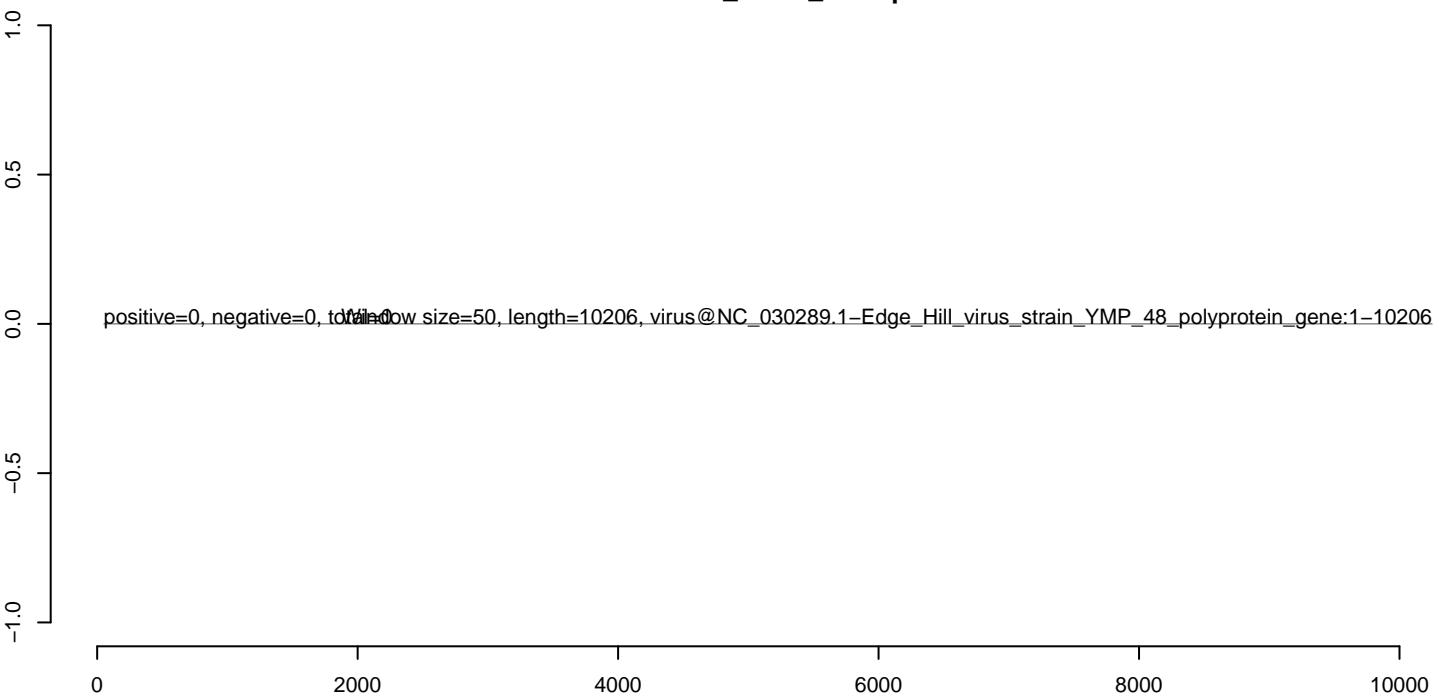
### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep



### AnGam\_Fcarc\_TN.rep





positive=0, negative=0, total=0

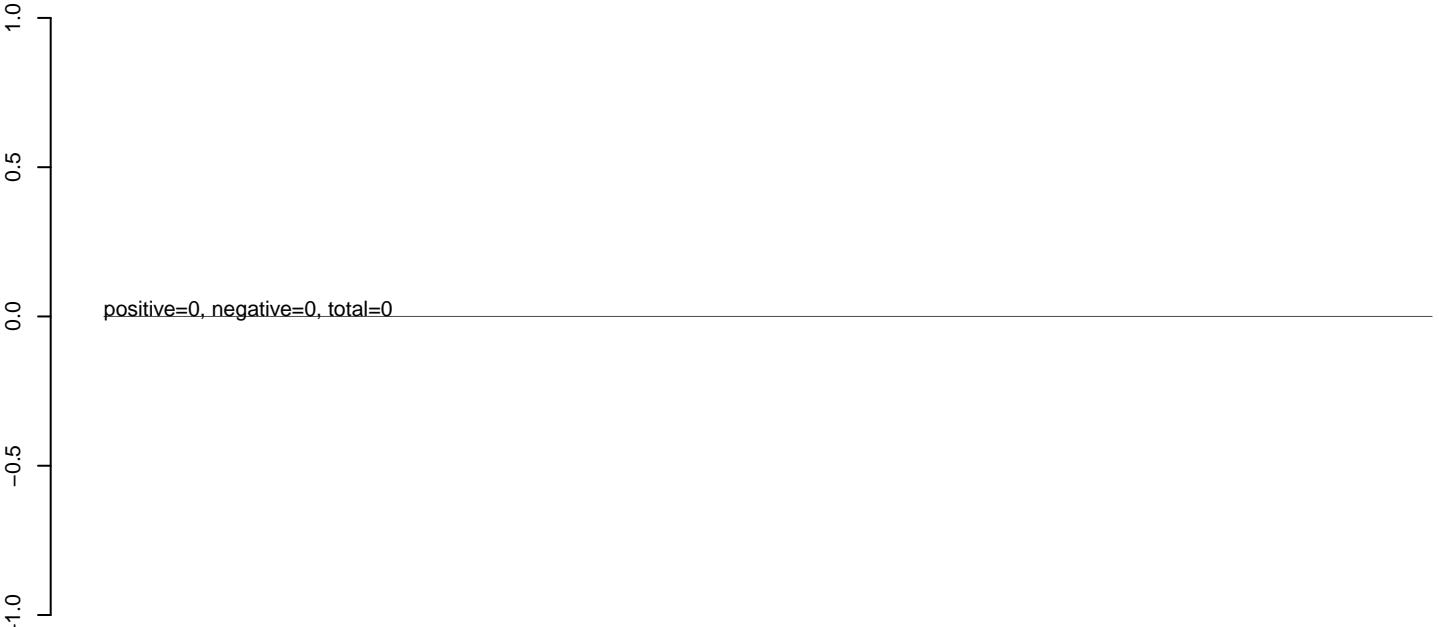


positive=0, negative=0, total=0

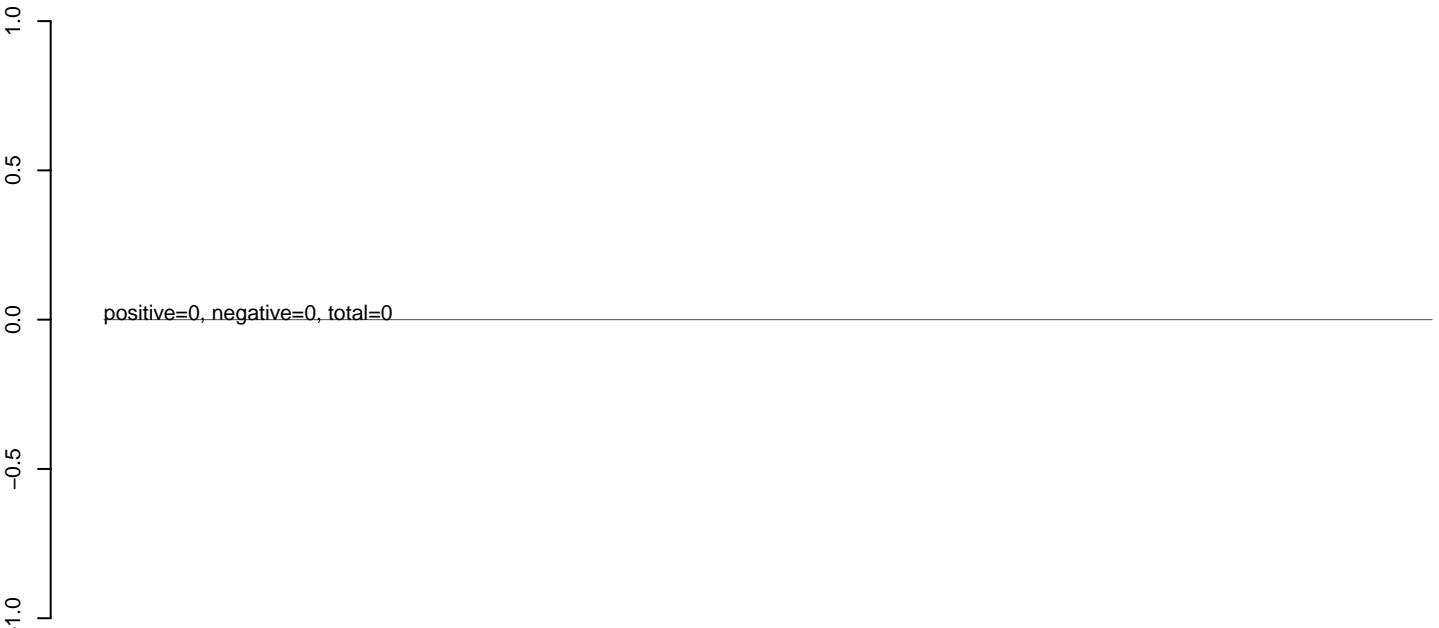


0                    2000                    4000                    6000                    8000

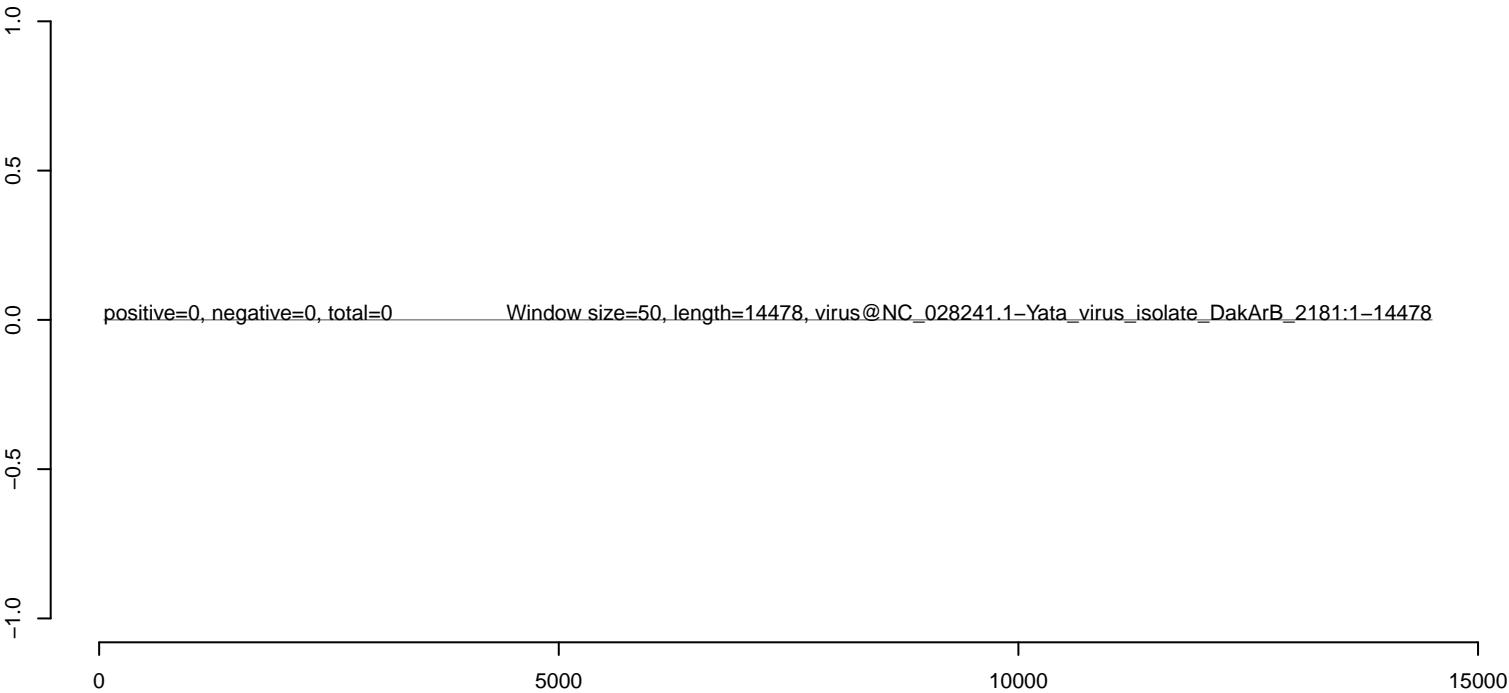
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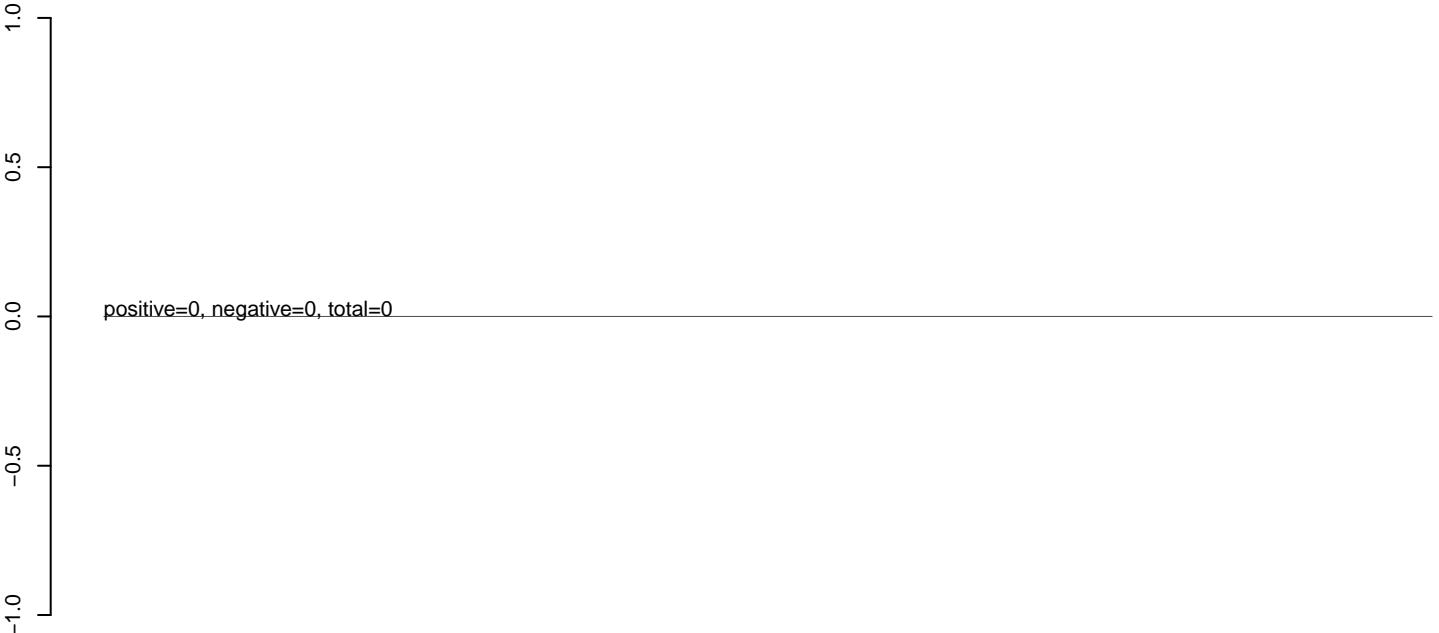
### AnGam\_Fcanc\_TN.24\_35.rep



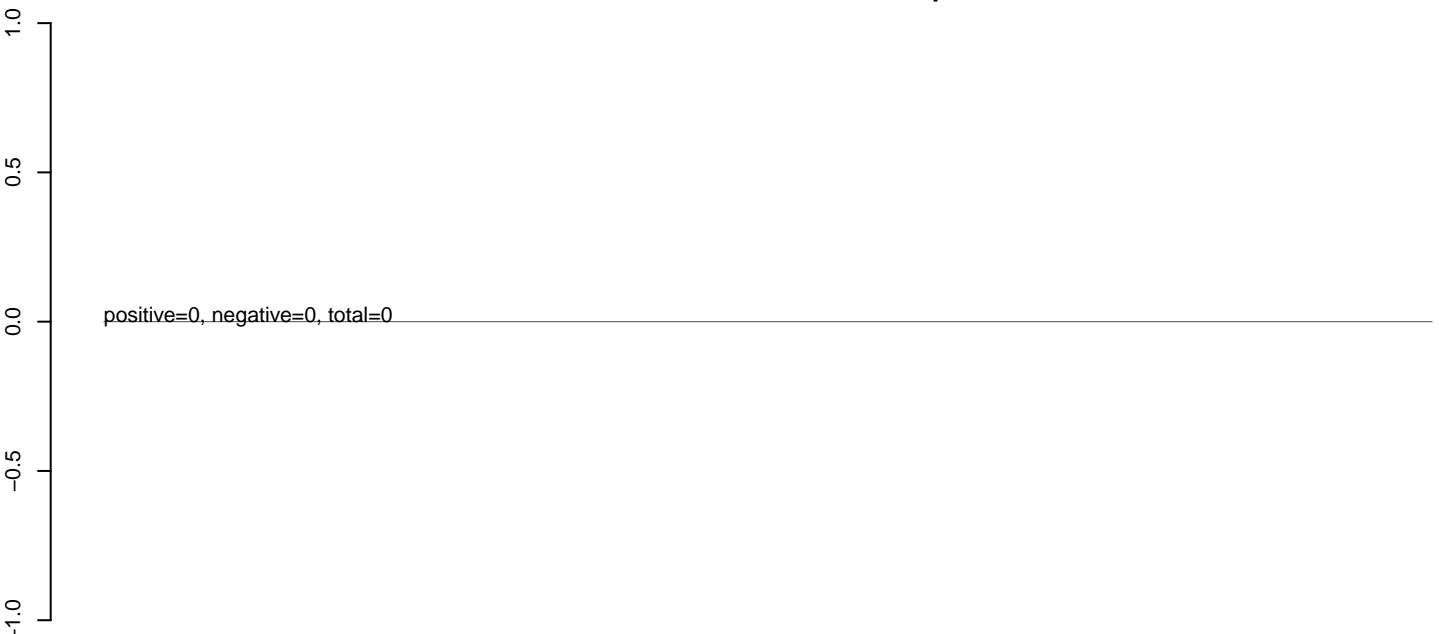
### AnGam\_Fcanc\_TN.rep



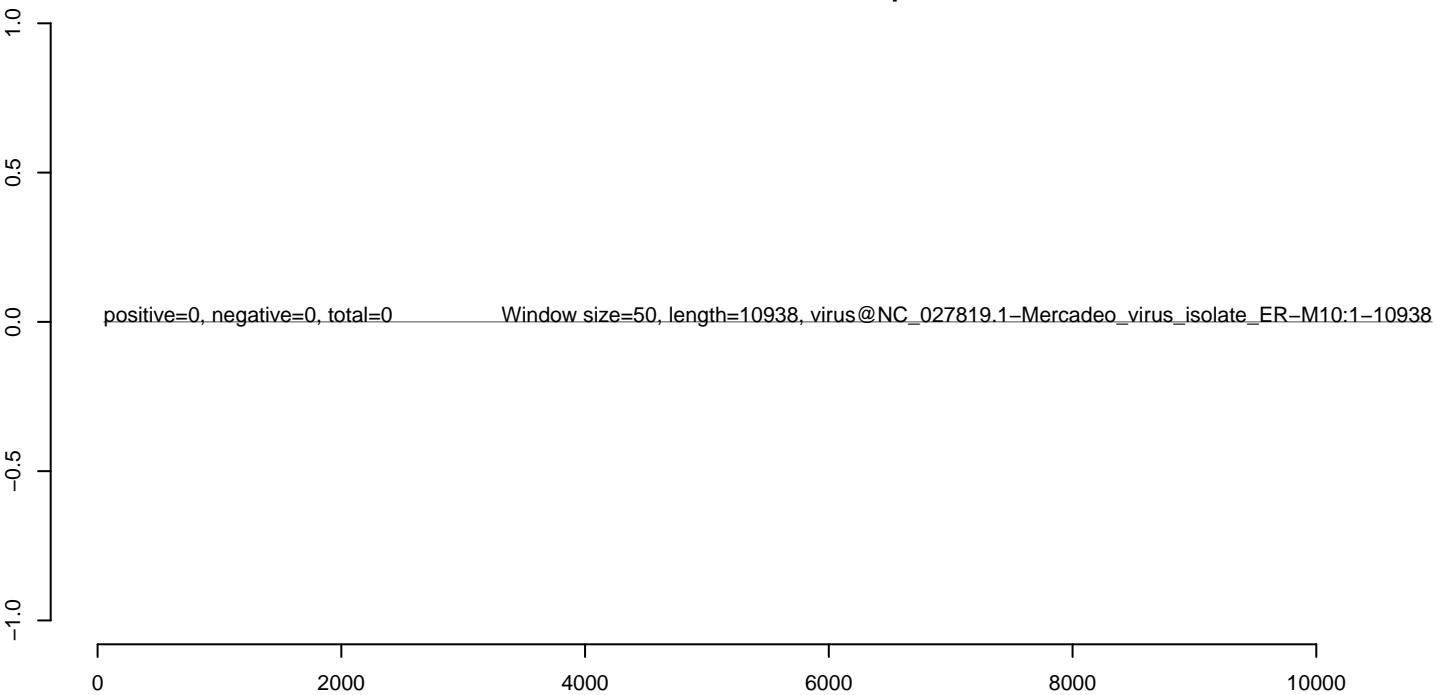
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0|Window size=50, length=10893, virus@NC\_027817.1-Parramatta\_River\_virus\_isolate\_92-B115745:1-10893

0 2000 4000 6000 8000 10000



positive=0, negative=0, total=0



positive=0, negative=0, total=0



S-3.1-Capitellidae-pipistrellivirustab01protein\_flavivirus\_polyprotein\_gene\_and\_truncated\_polyprotein\_flavivirus\_polyprotein\_gene\_genes1-10284

### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

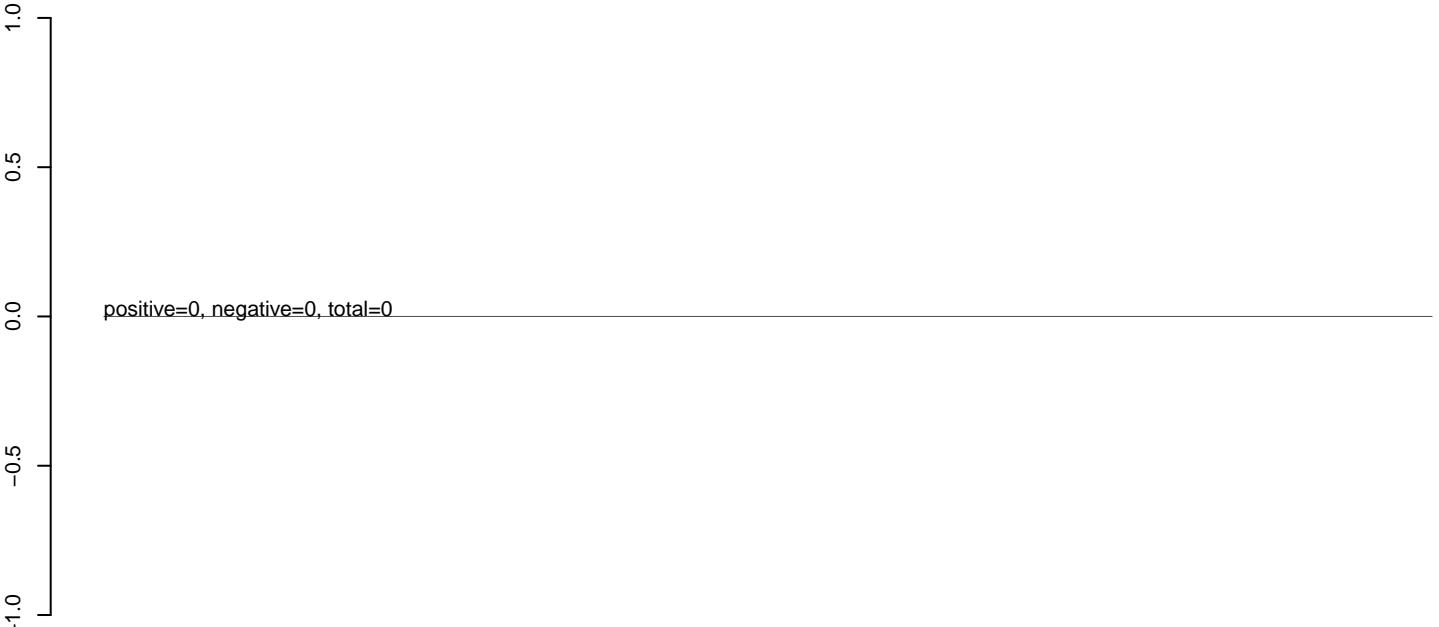
### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0

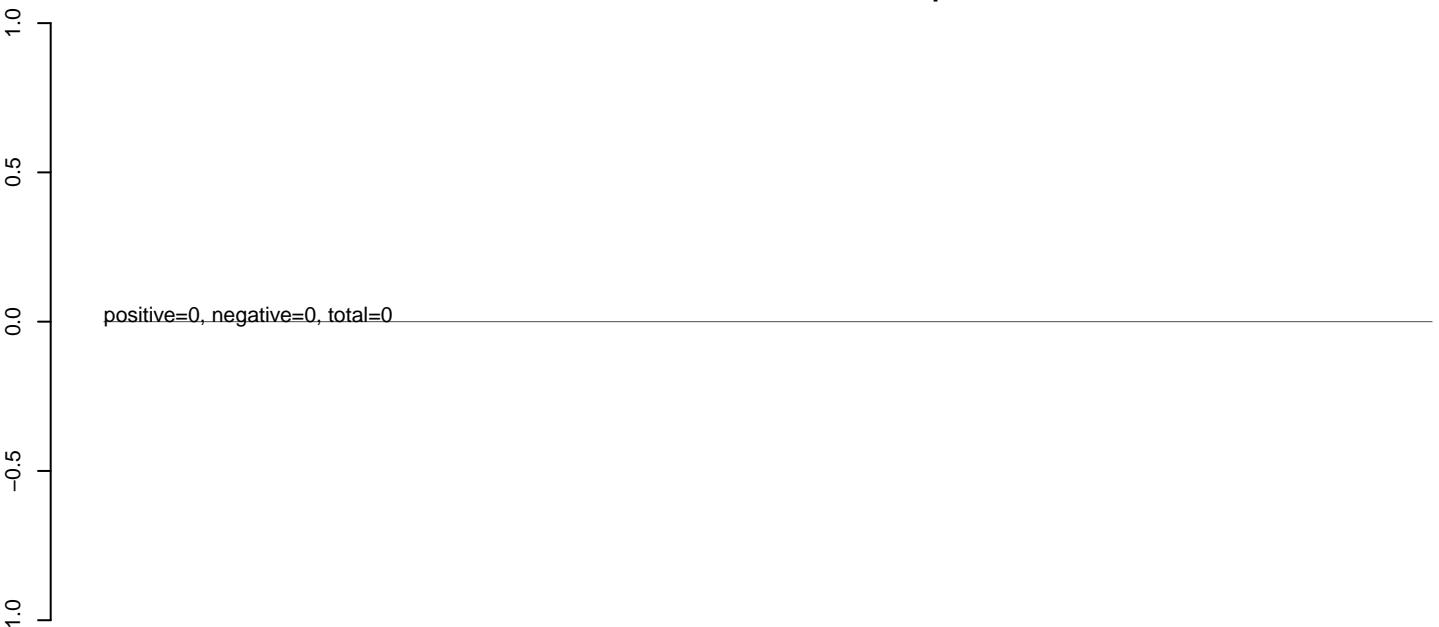
Window size=50, length=10125, virus@NC\_026620.1-Jutiapa\_virus\_strain\_JG-128:1-10125

0 2000 4000 6000 8000 10000

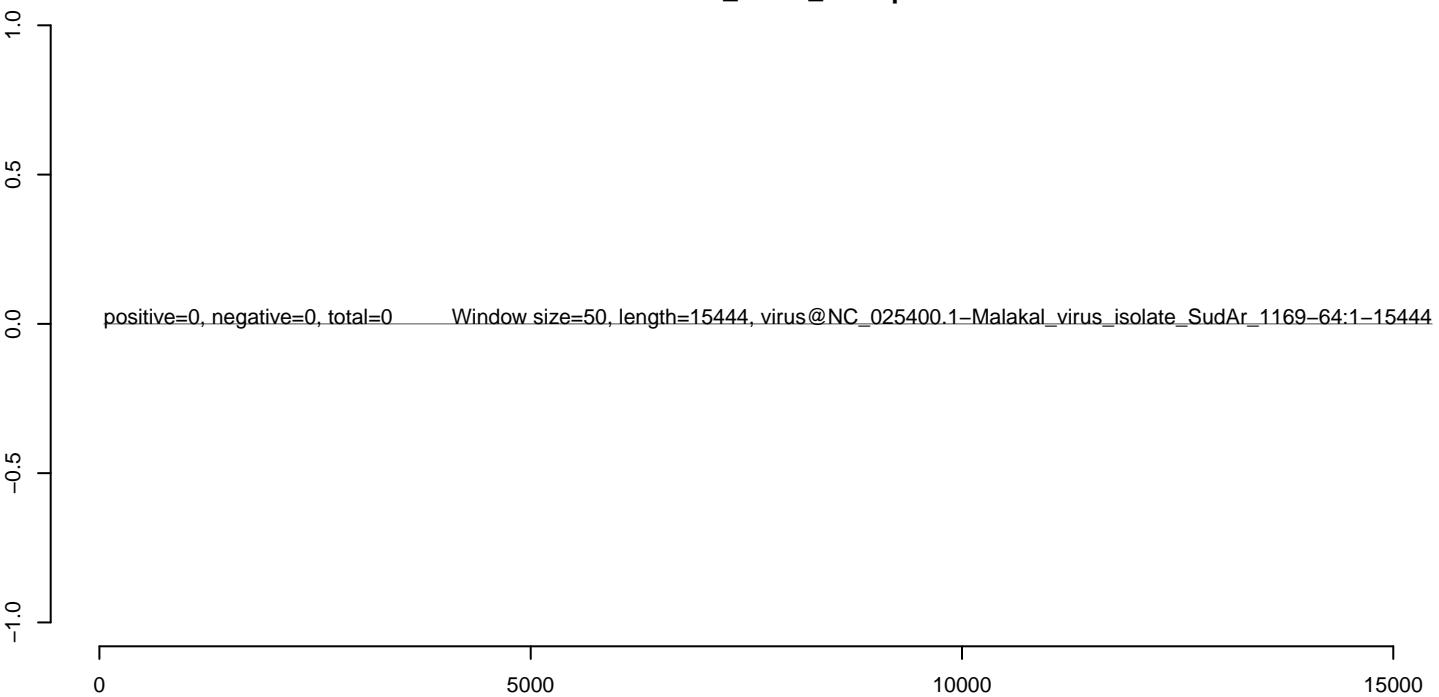
### AnGam\_Fcarc\_TN.18\_23.rep



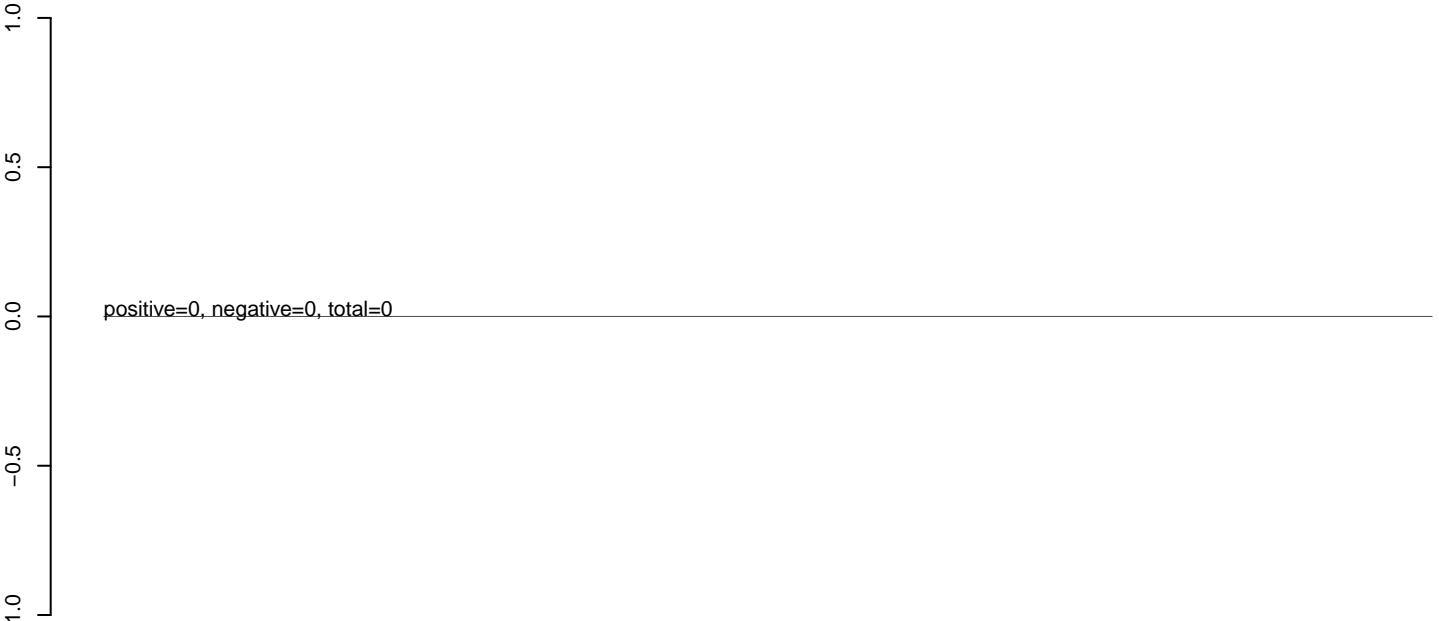
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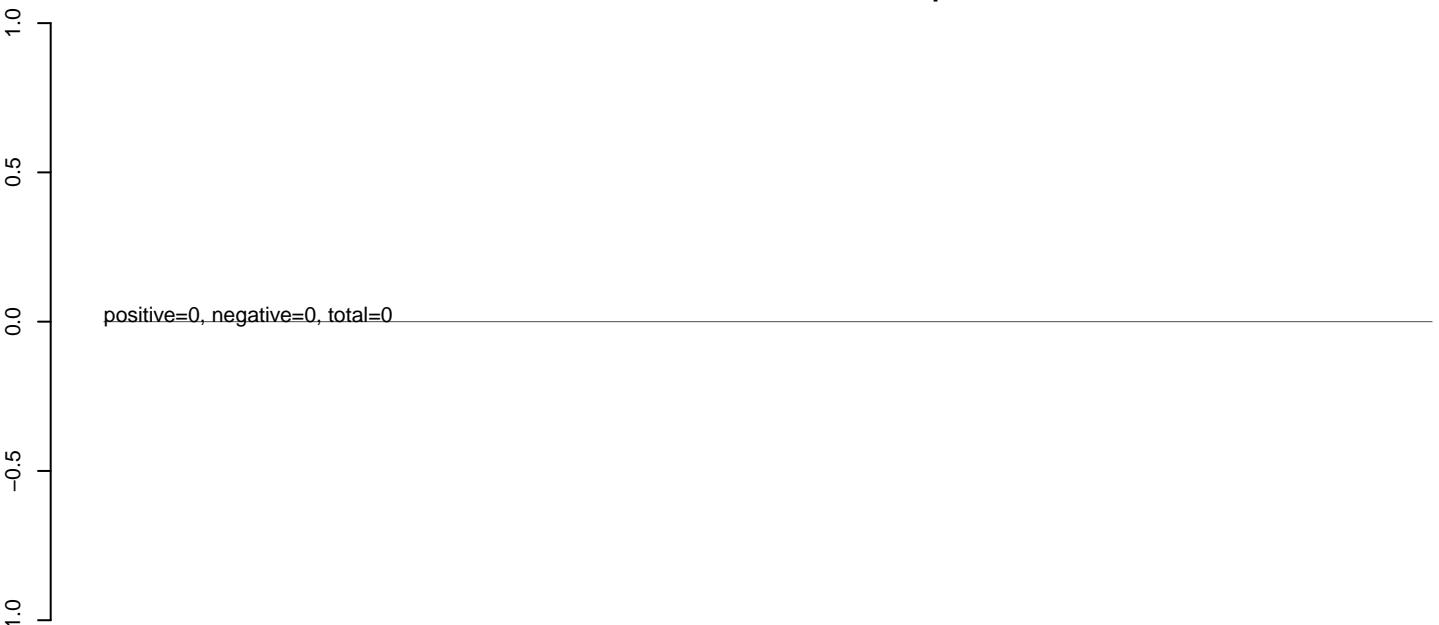
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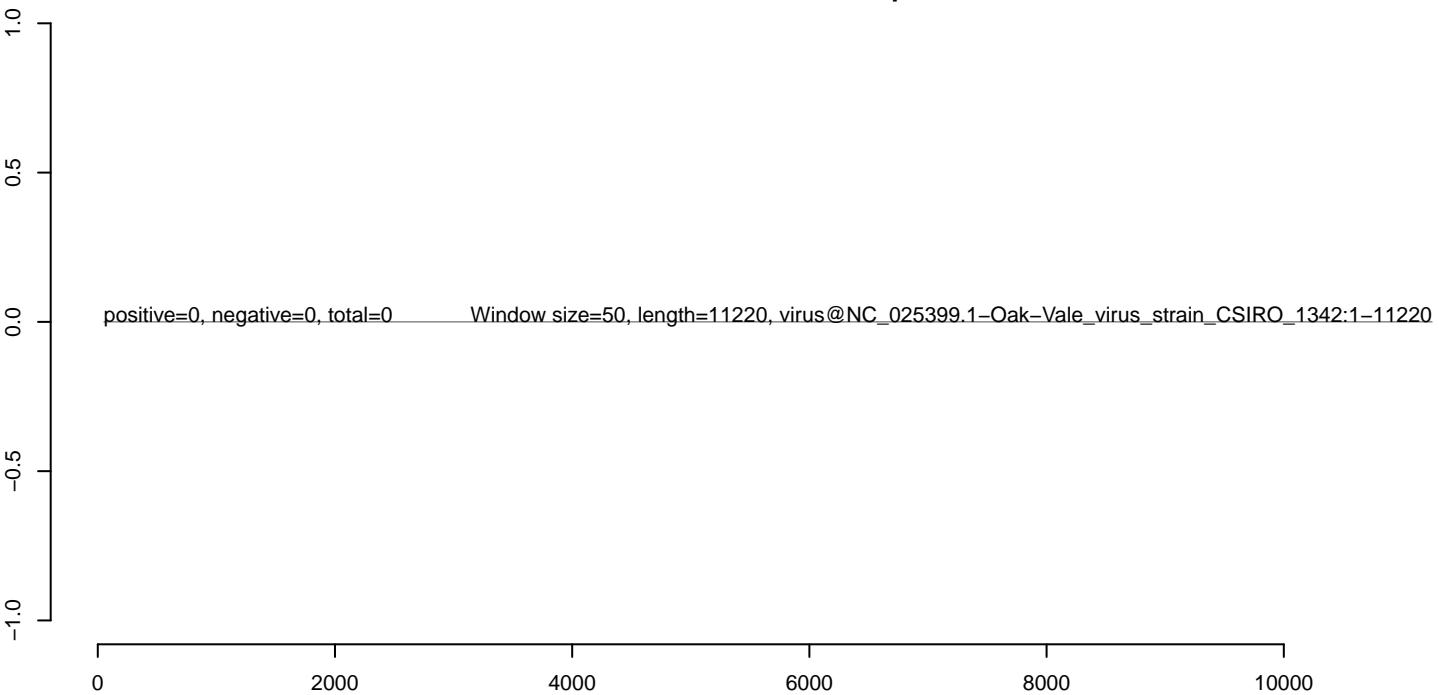
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

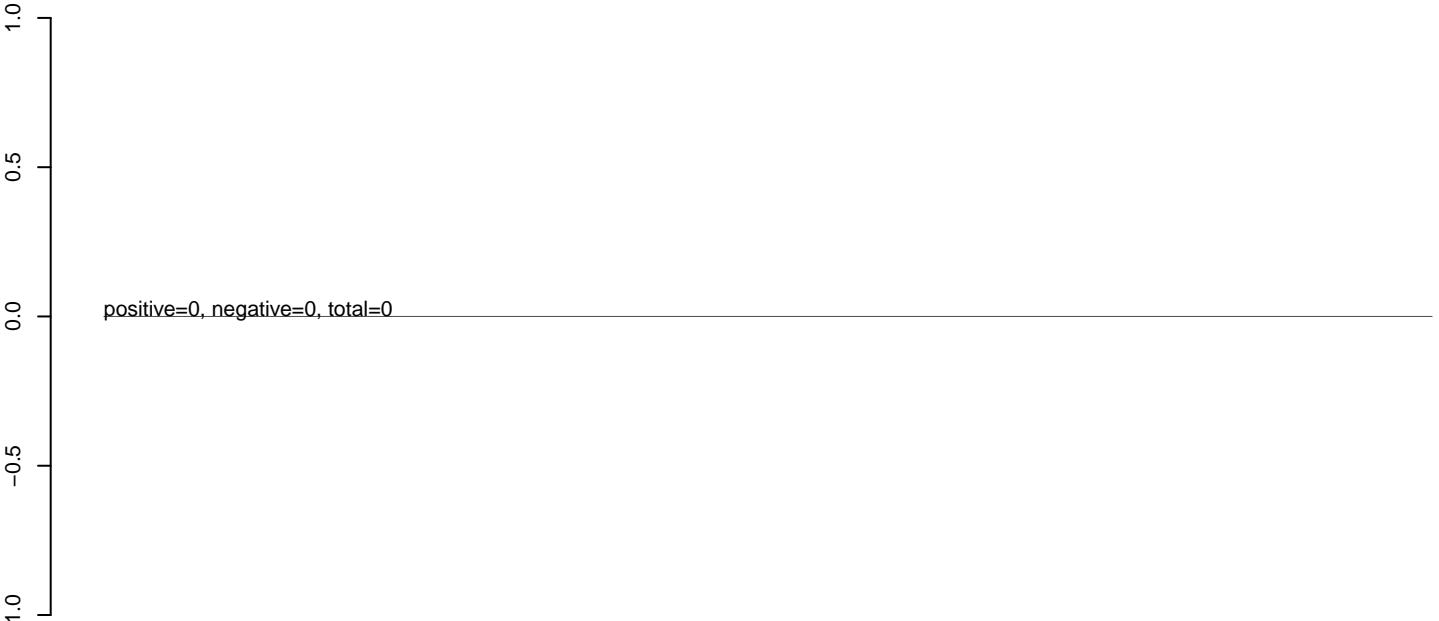
### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

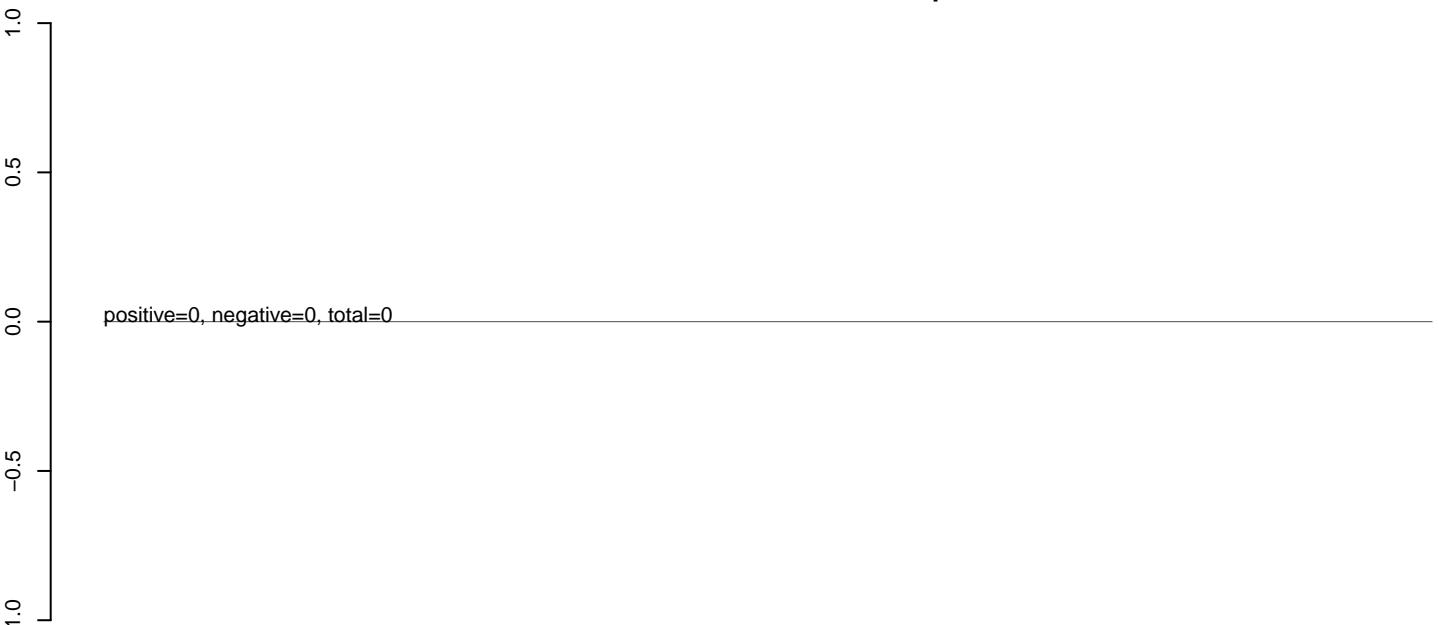
### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0    Window size=50, length=11876, virus@NC\_025395.1–Puerto\_Almendras\_virus\_isolate\_LO-39:1–11876

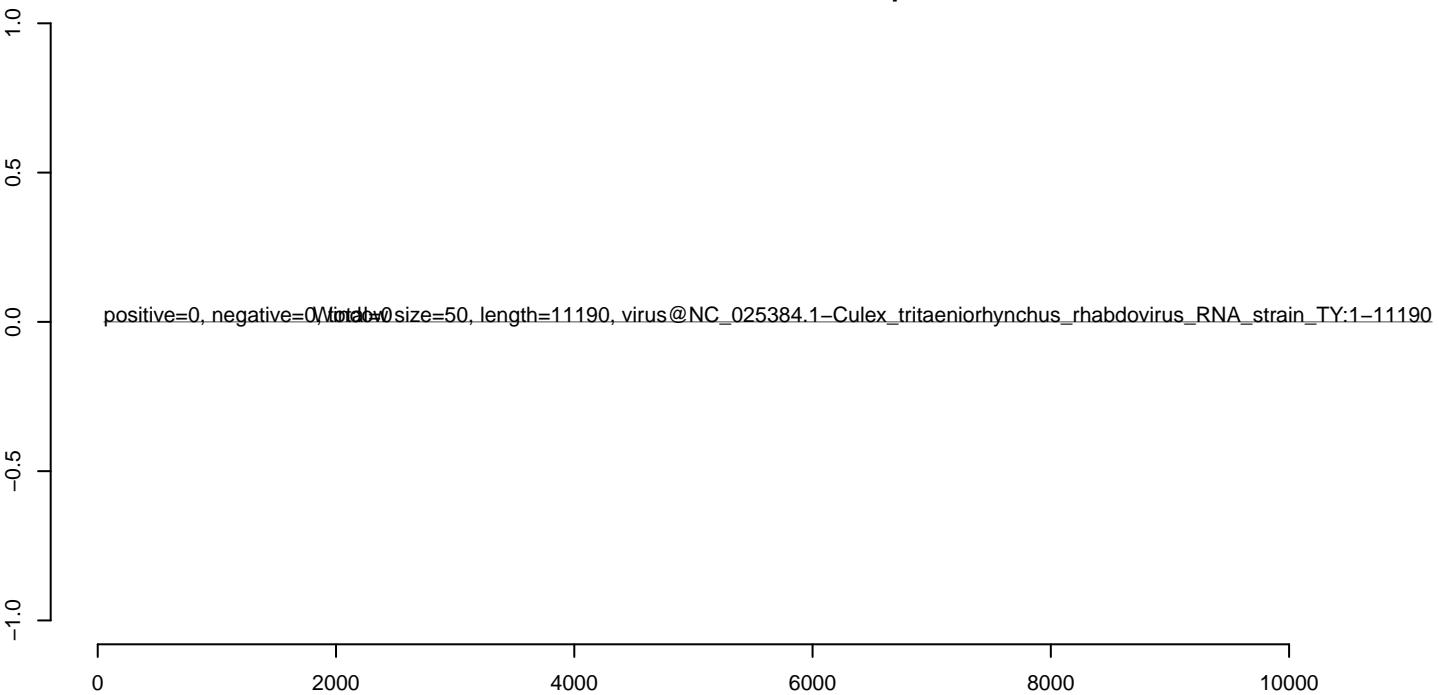
### AnGam\_Fcanc\_TN.18\_23.rep



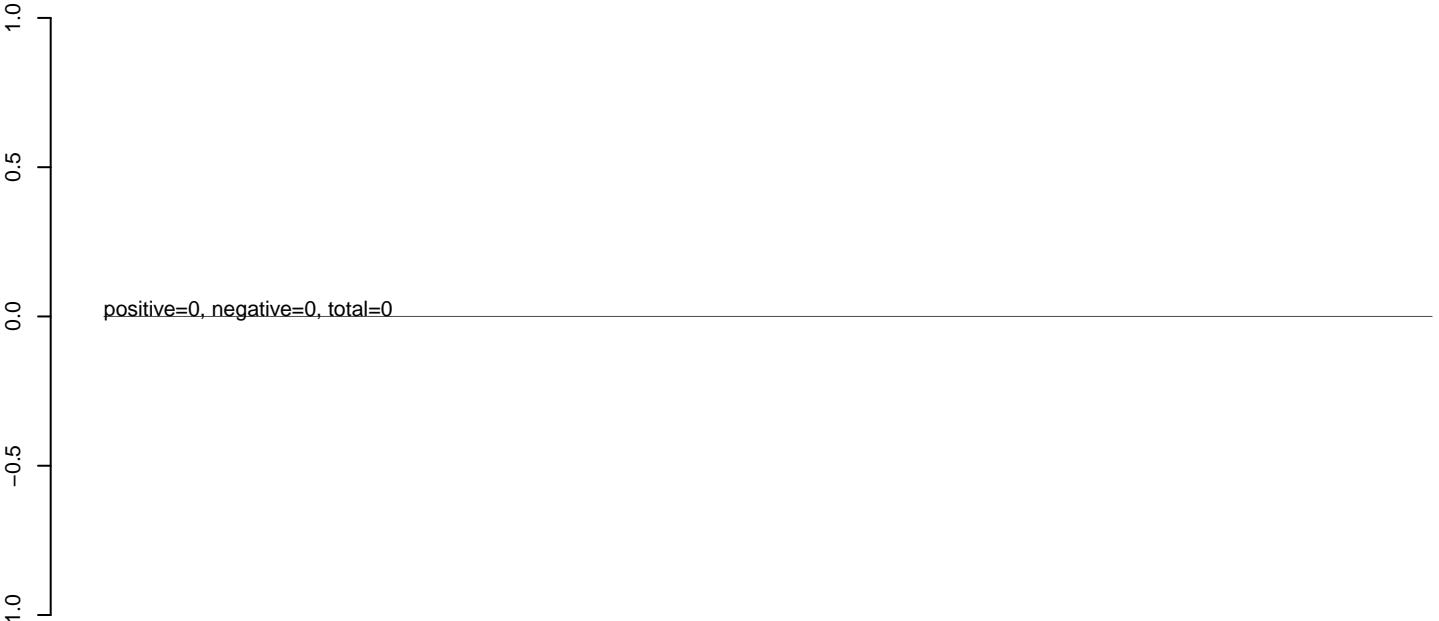
### AnGam\_Fcanc\_TN.24\_35.rep



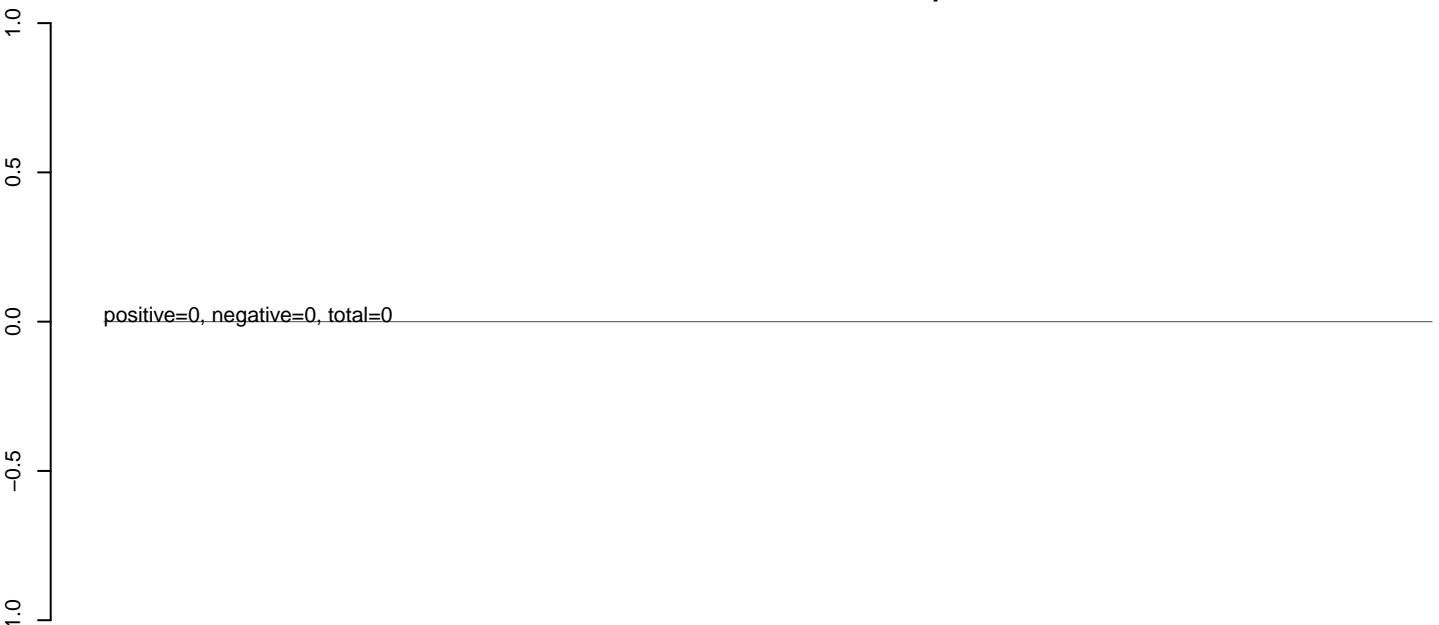
### AnGam\_Fcanc\_TN.rep



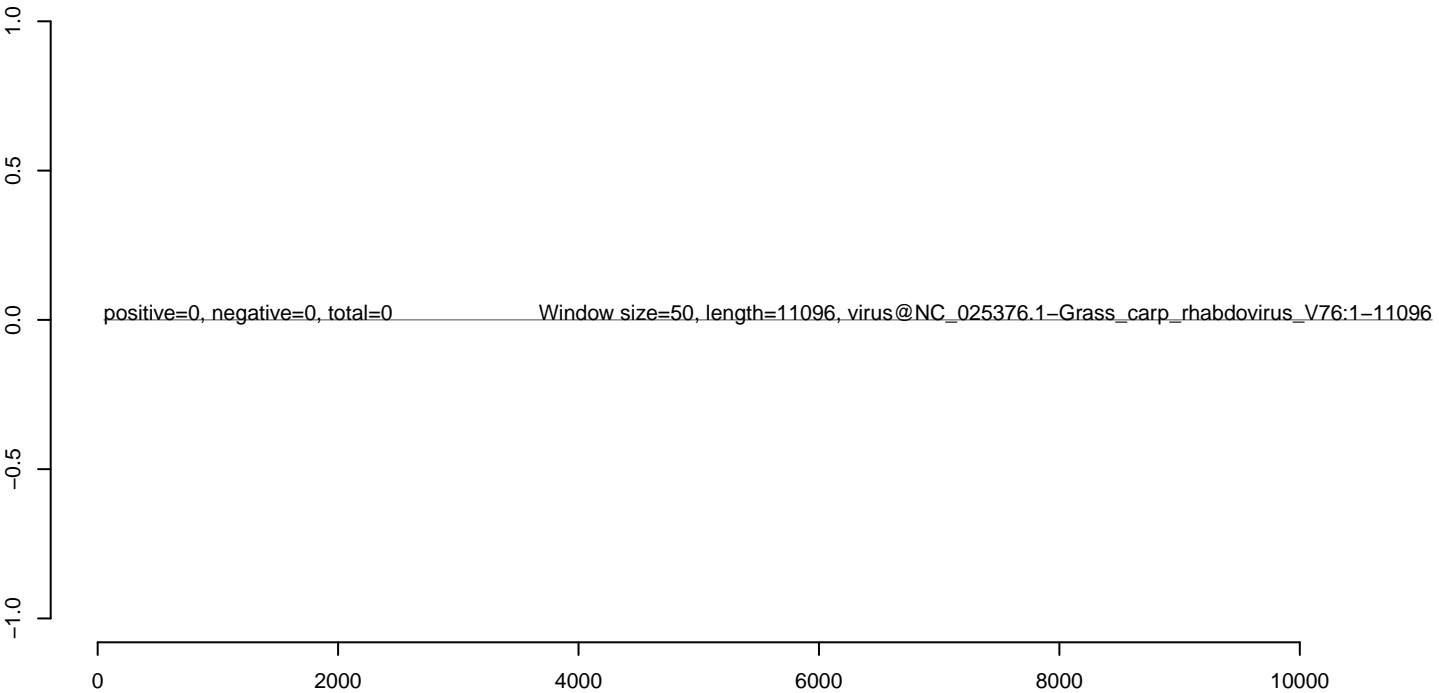
### AnGam\_Fcanc\_TN.18\_23.rep



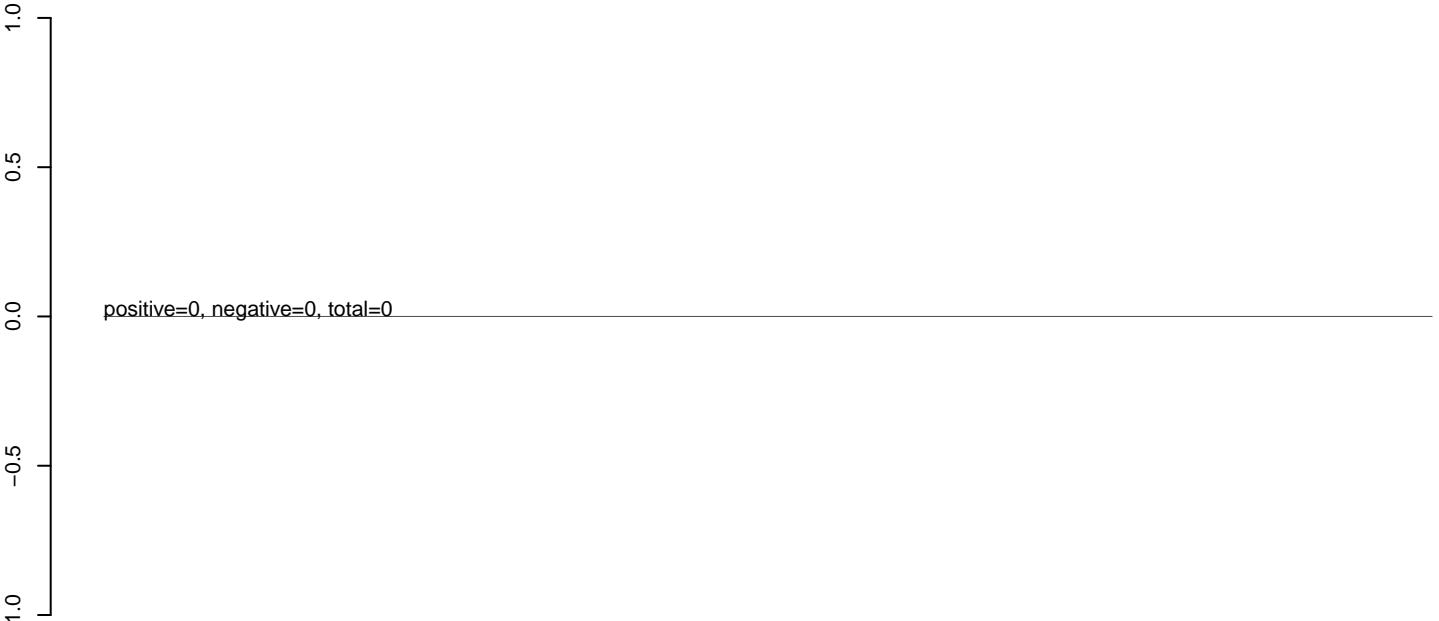
### AnGam\_Fcanc\_TN.24\_35.rep



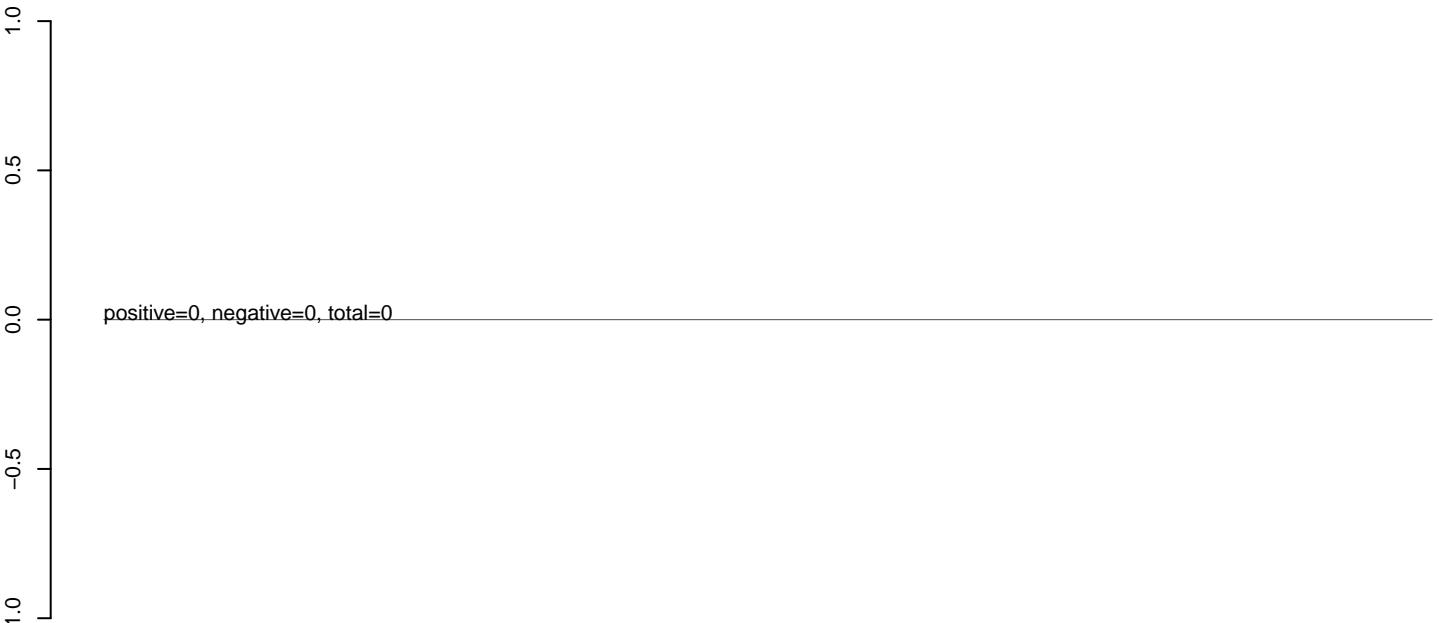
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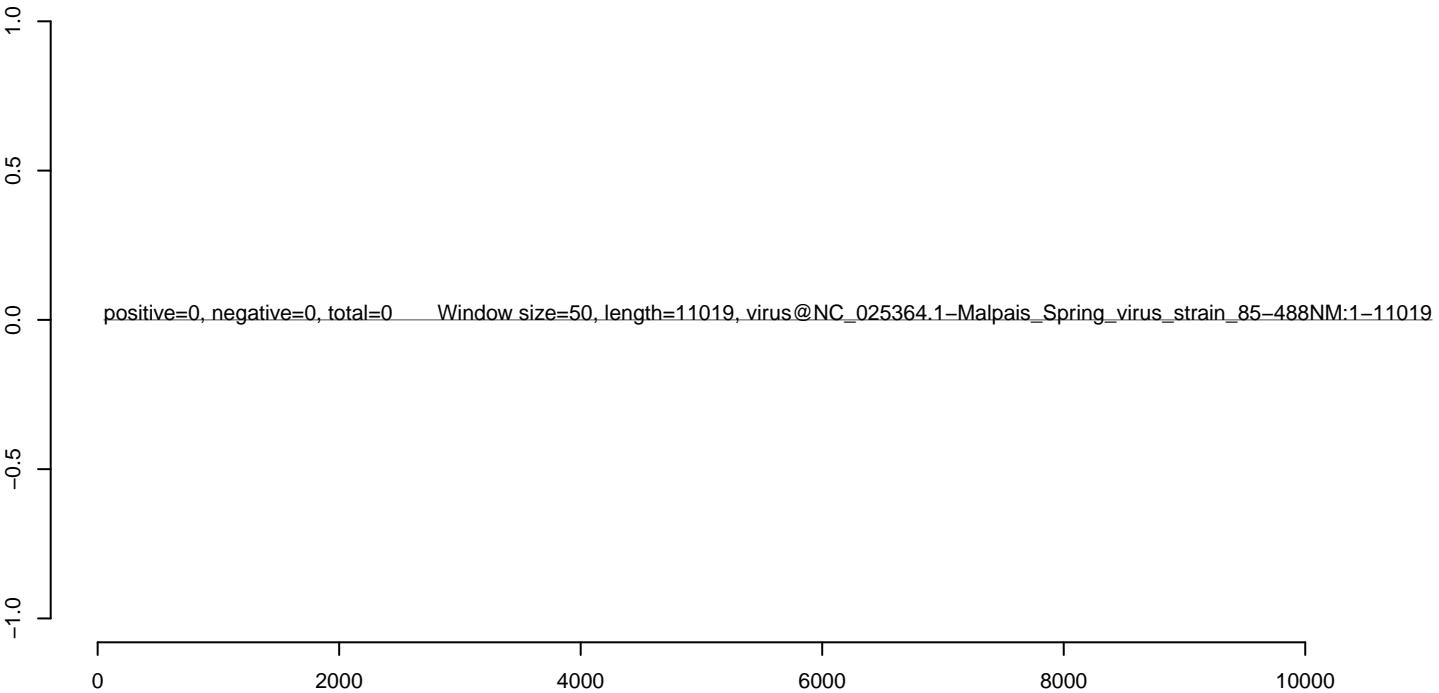
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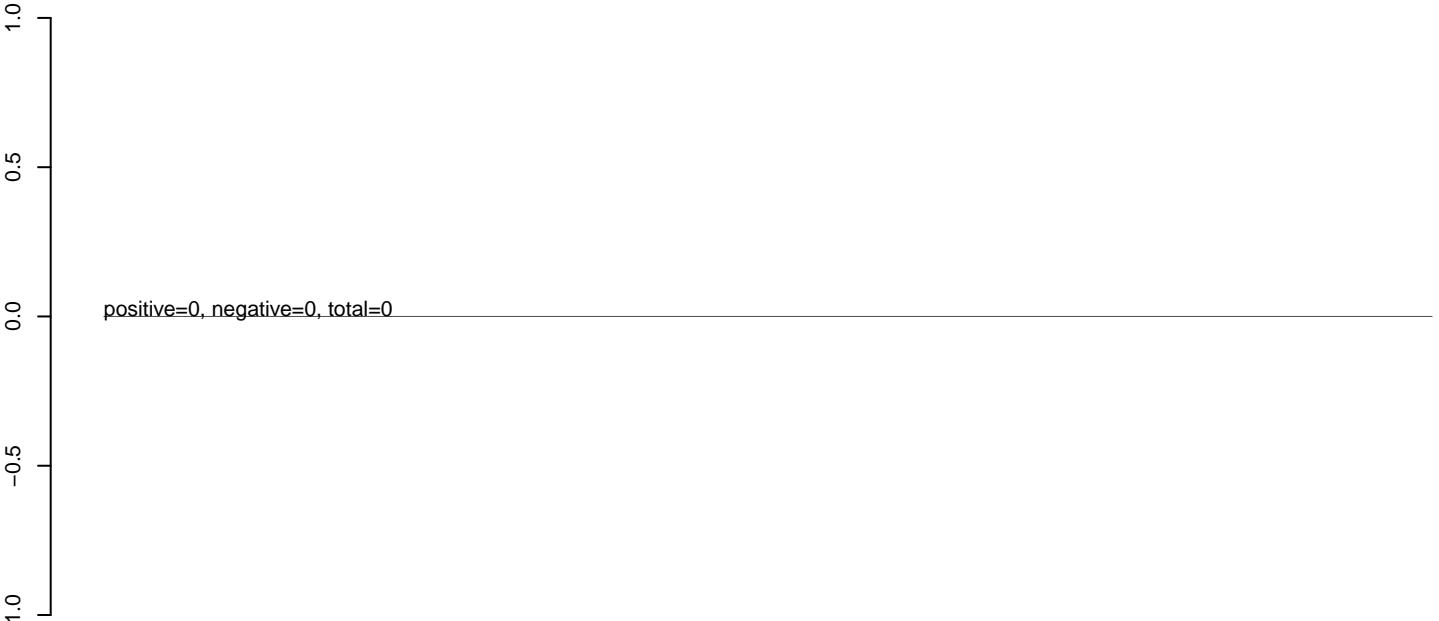
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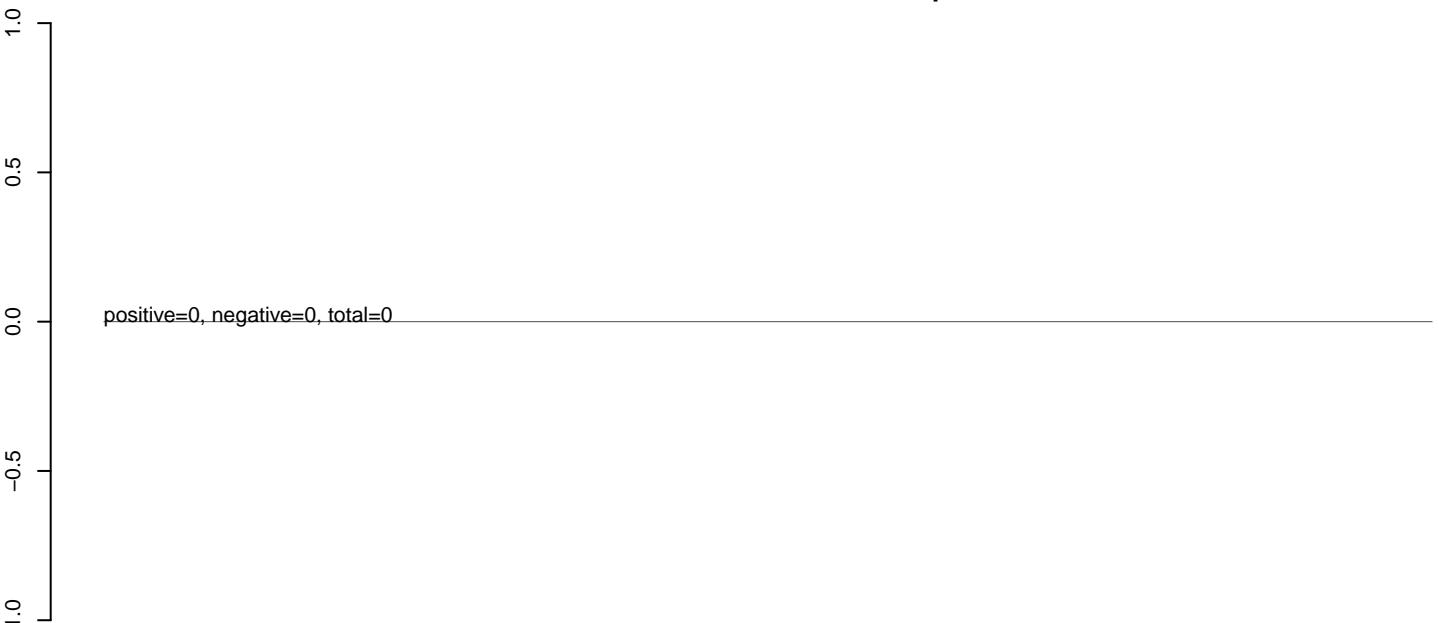
### AnGam\_Fcanc\_TN.rep



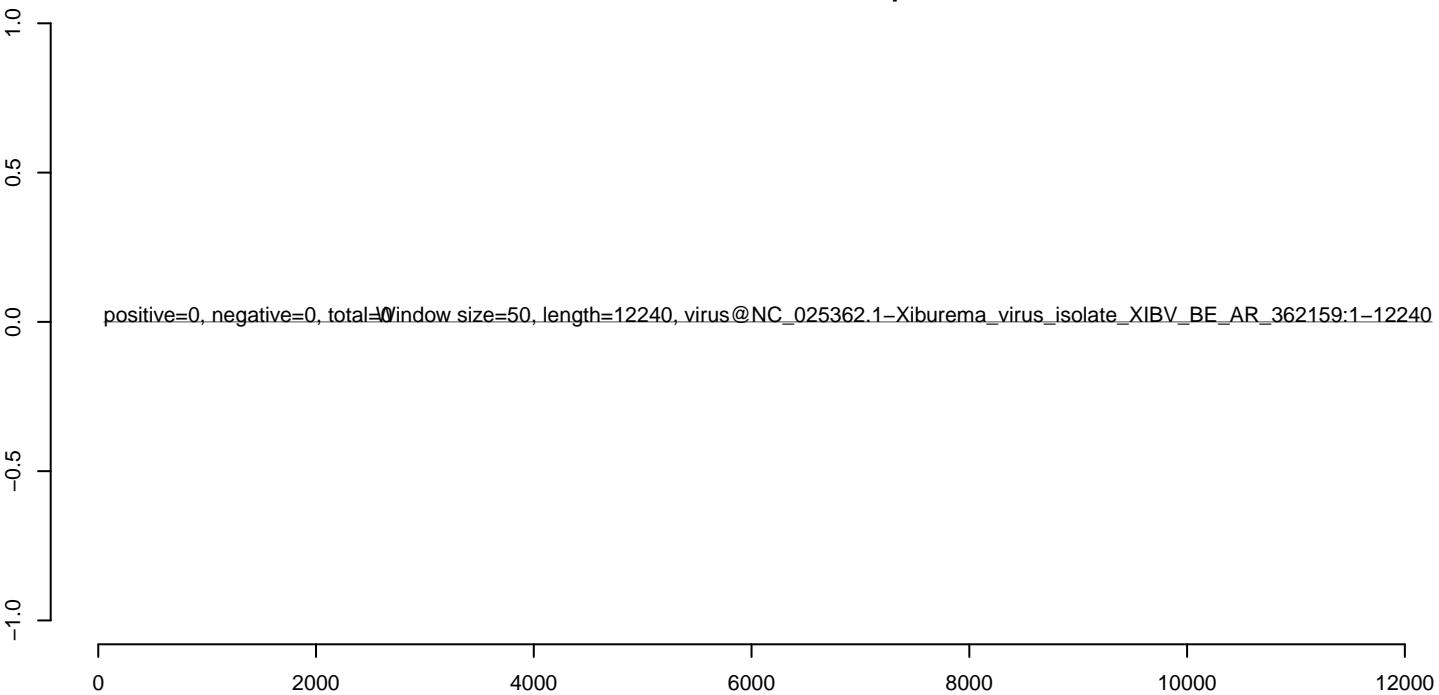
### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep



### AnGam\_Fcarc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

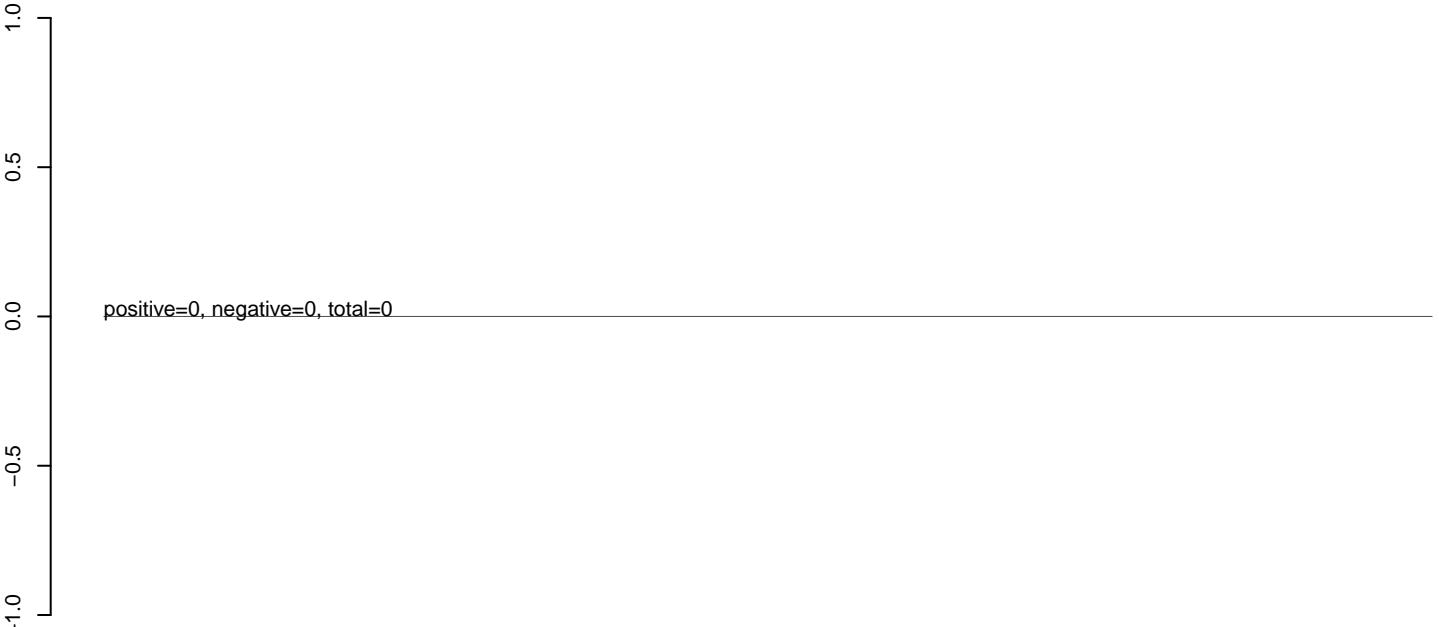
### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0

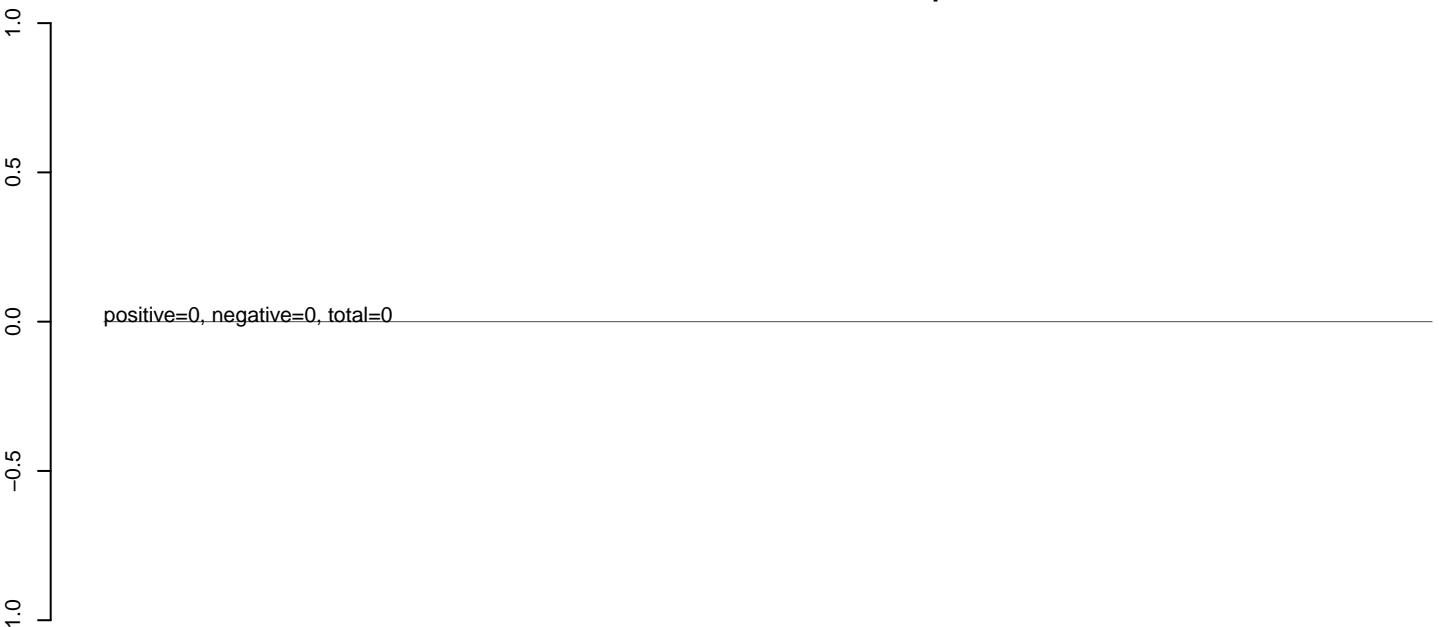
Window size=50, length=11526, virus@NC\_025359.1-Moussa\_virus\_isolate\_C23:1-11526

0 2000 4000 6000 8000 10000 12000

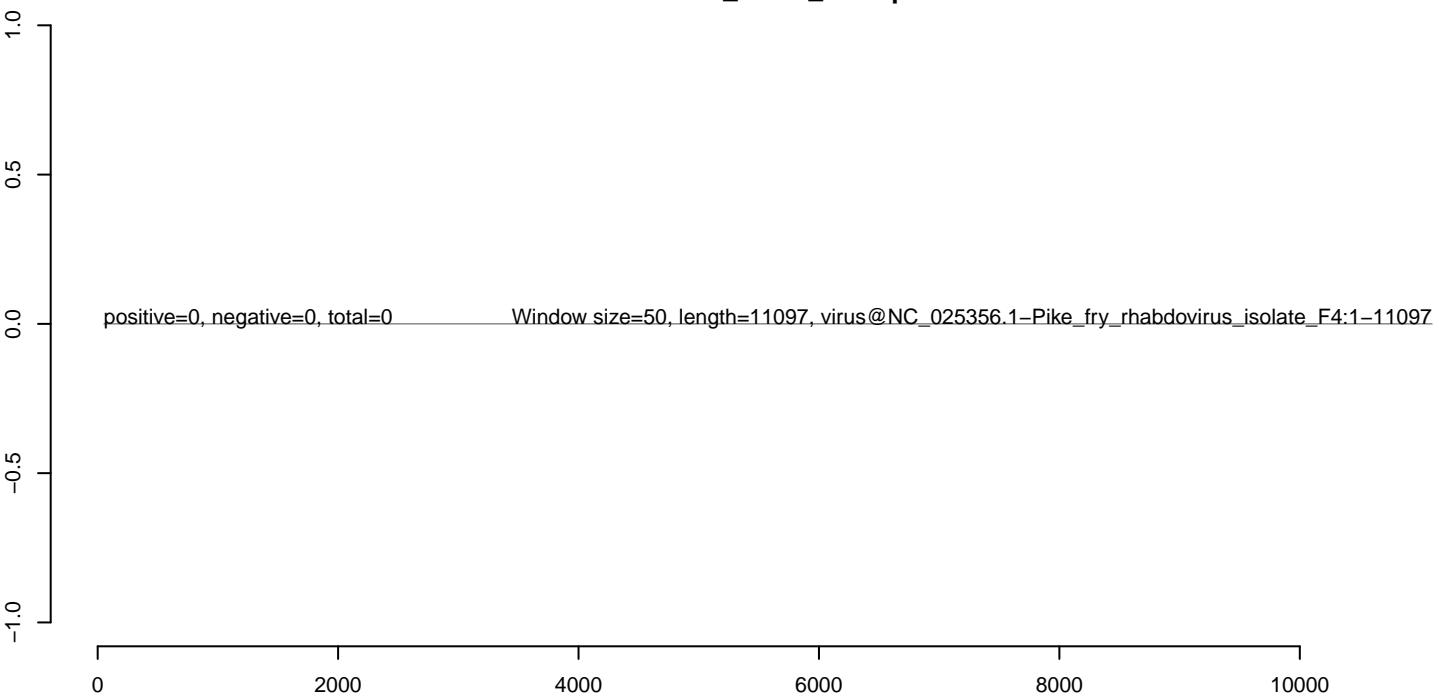
### AnGam\_Fcanc\_TN.18\_23.rep



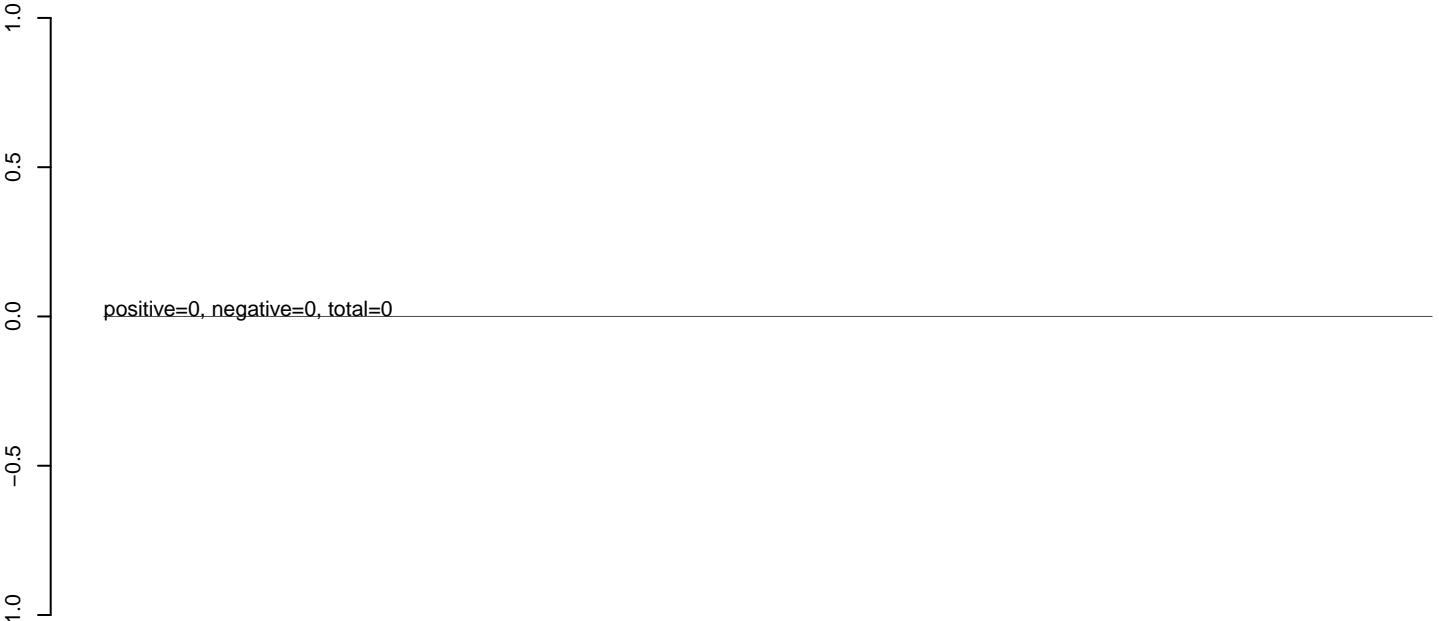
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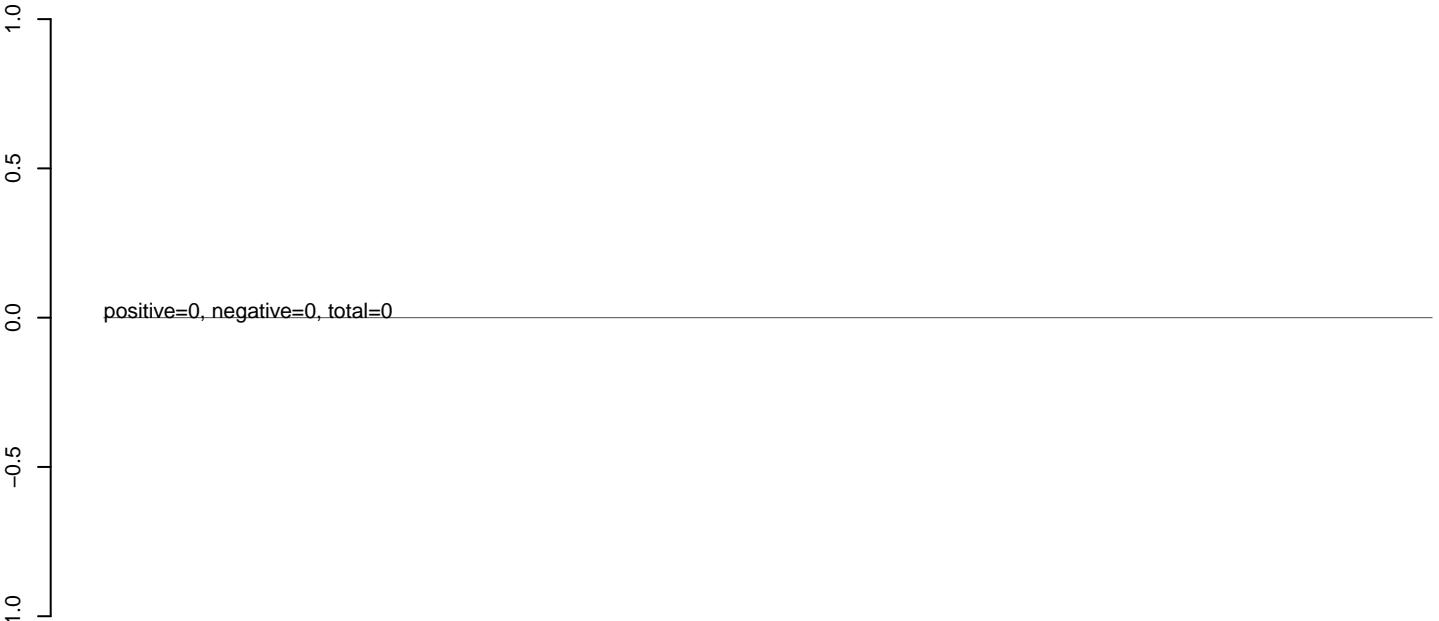
### AnGam\_Fcanc\_TN.rep



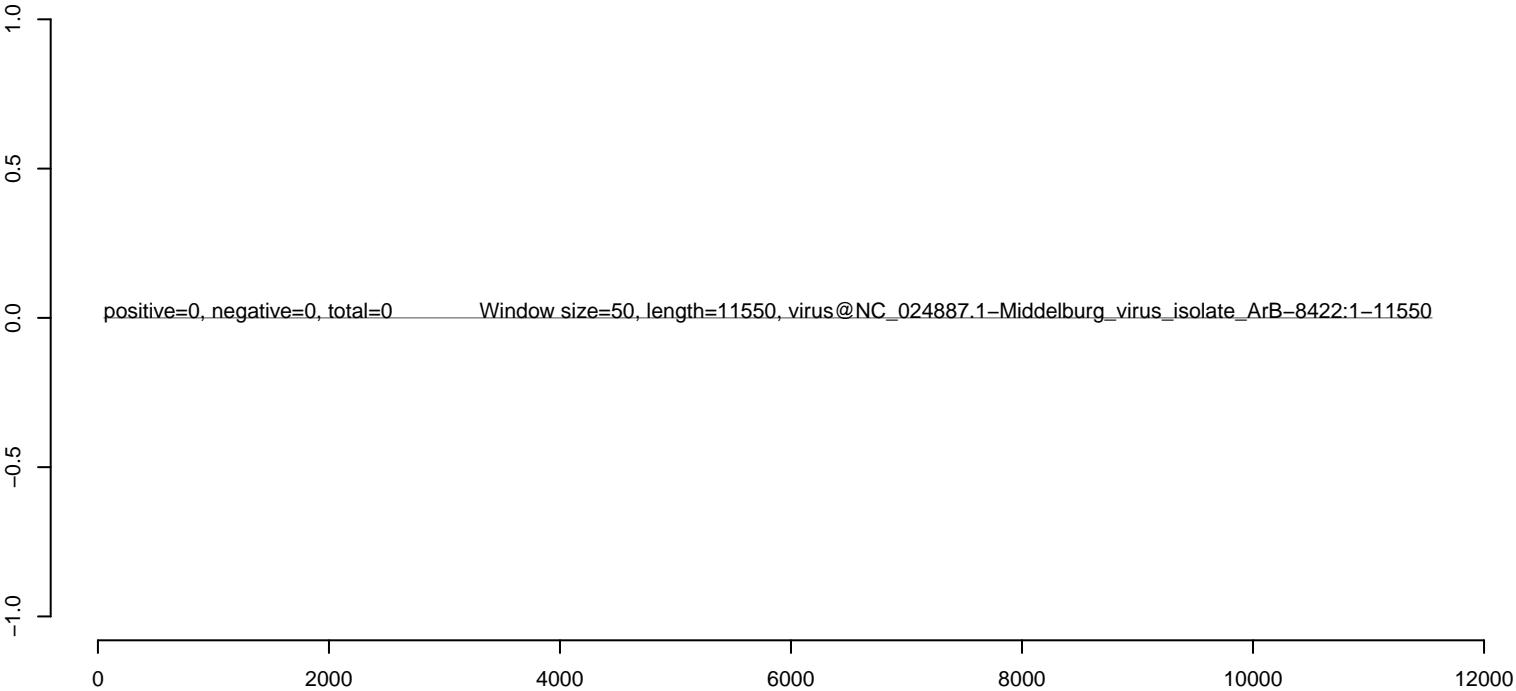
### AnGam\_Fcanc\_TN.18\_23.rep



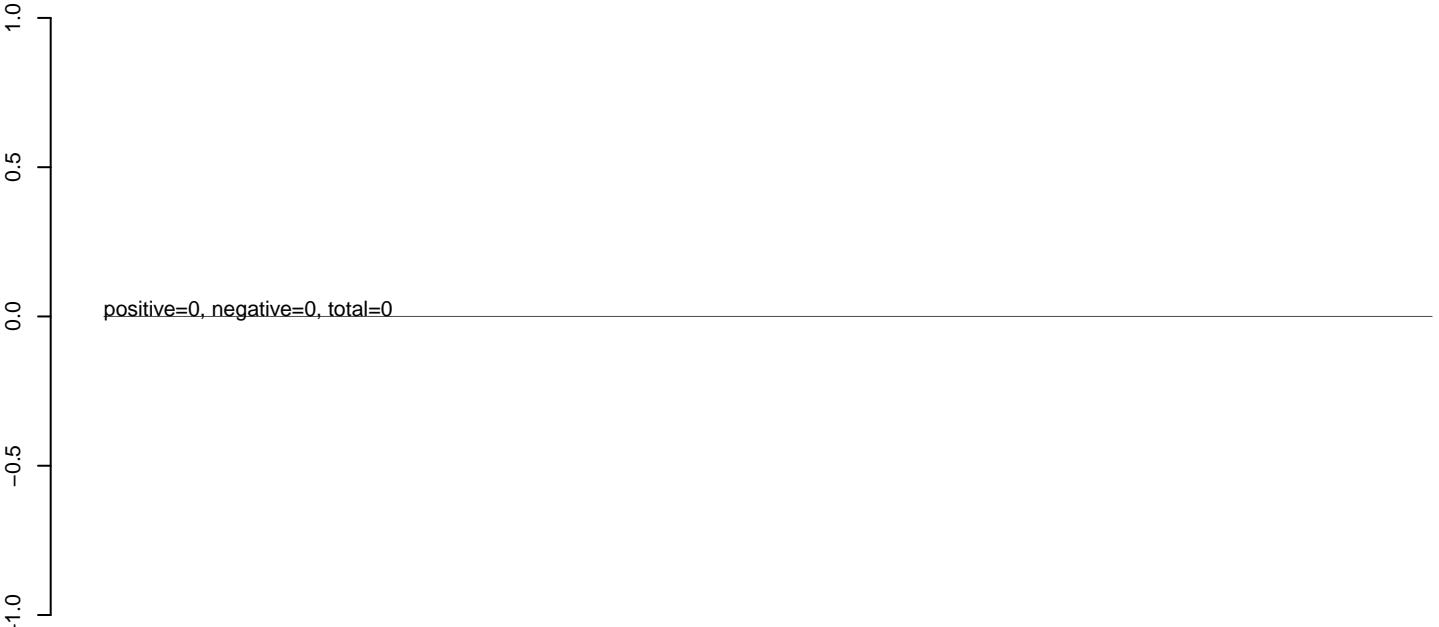
### AnGam\_Fcanc\_TN.24\_35.rep



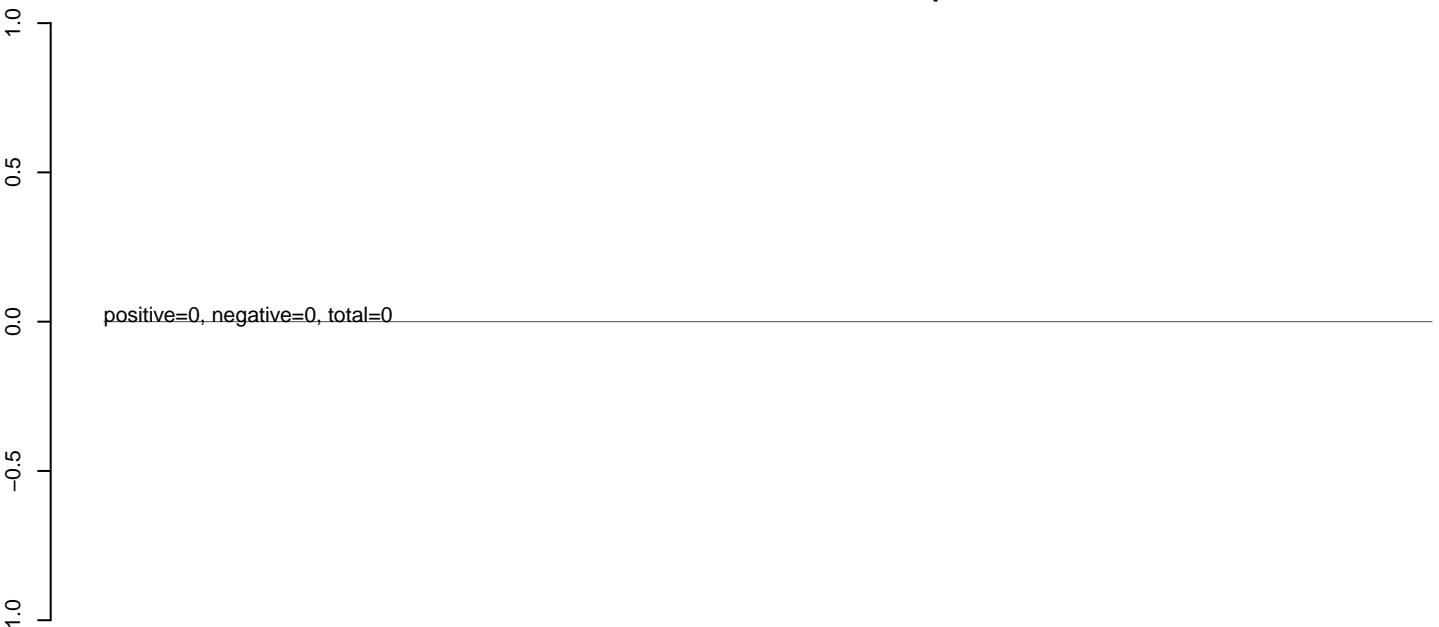
### AnGam\_Fcanc\_TN.rep



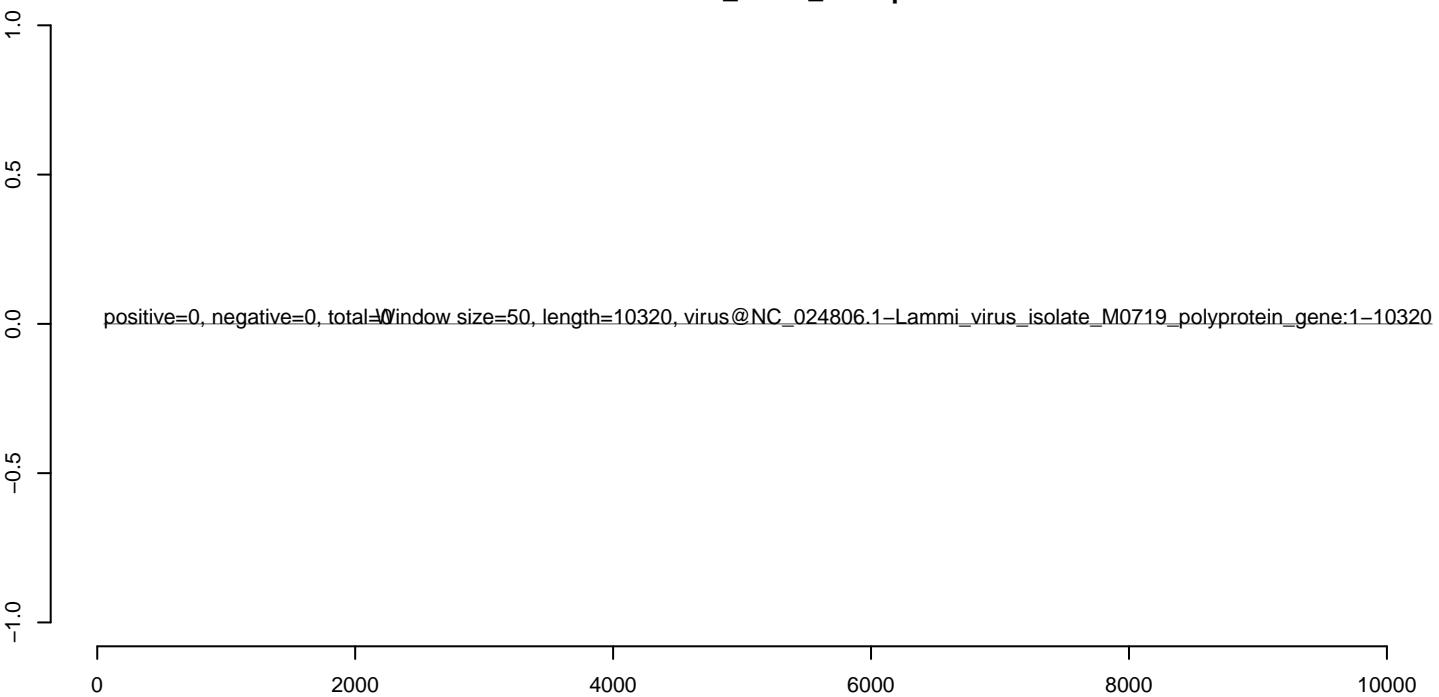
### AnGam\_Fcanc\_TN.18\_23.rep



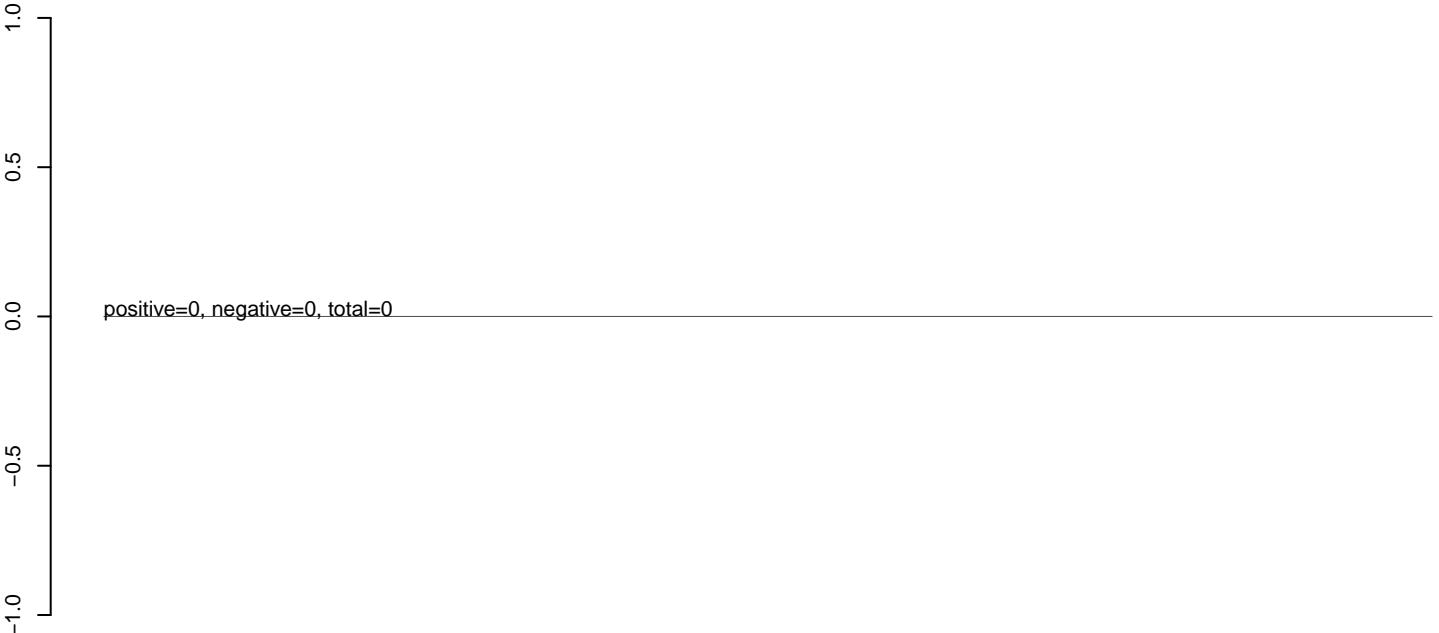
### AnGam\_Fcanc\_TN.24\_35.rep



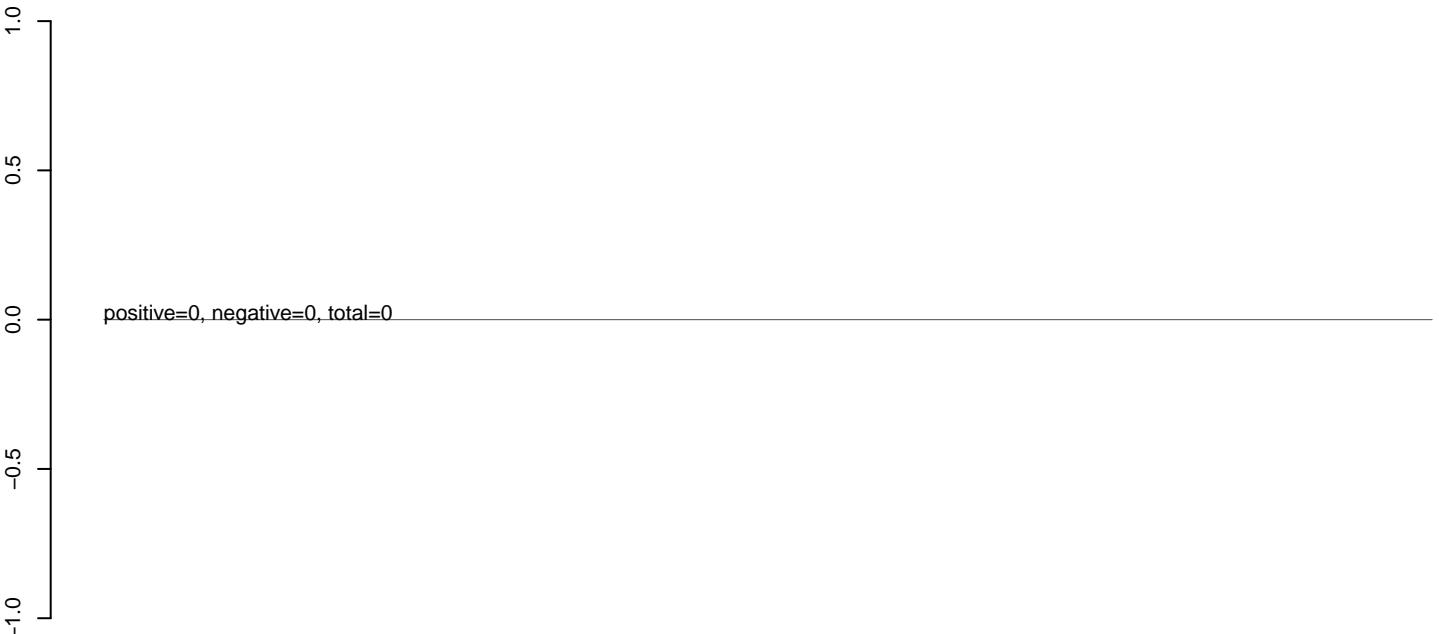
### AnGam\_Fcanc\_TN.rep



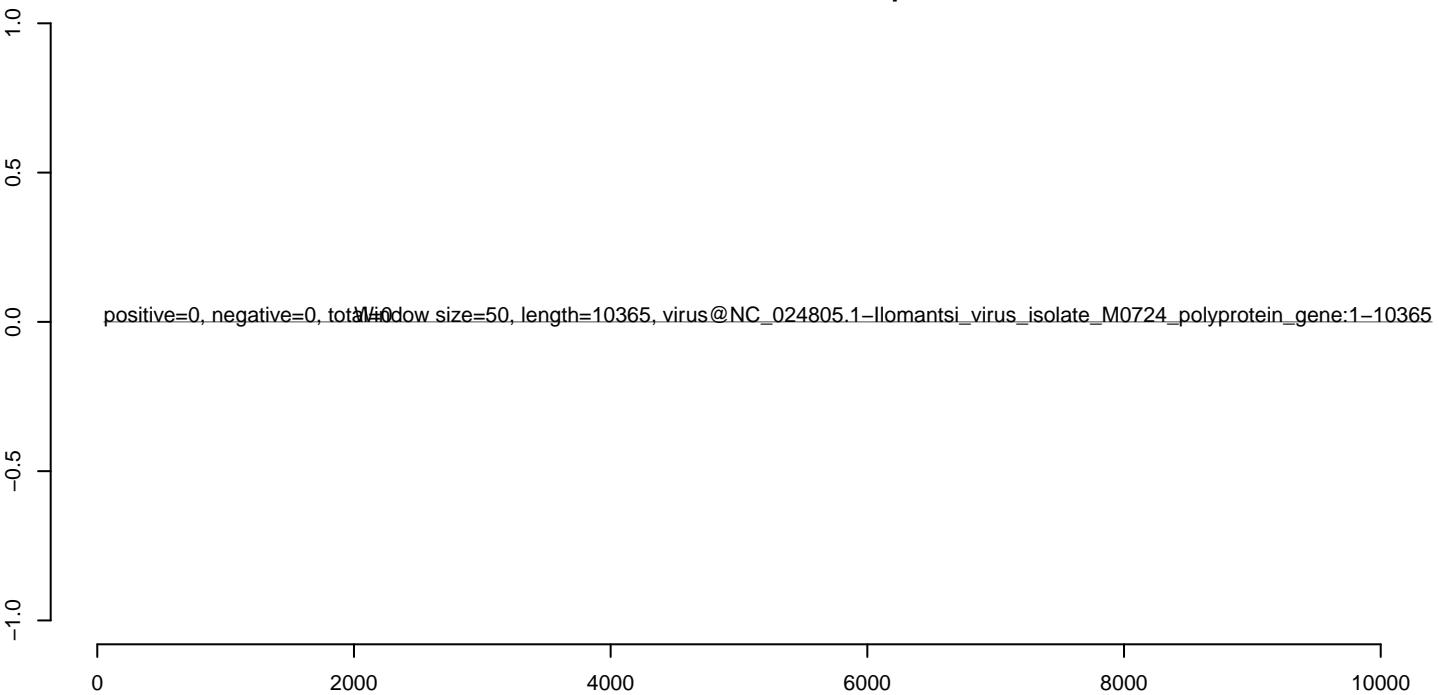
### AnGam\_Fcarc\_TN.18\_23.rep



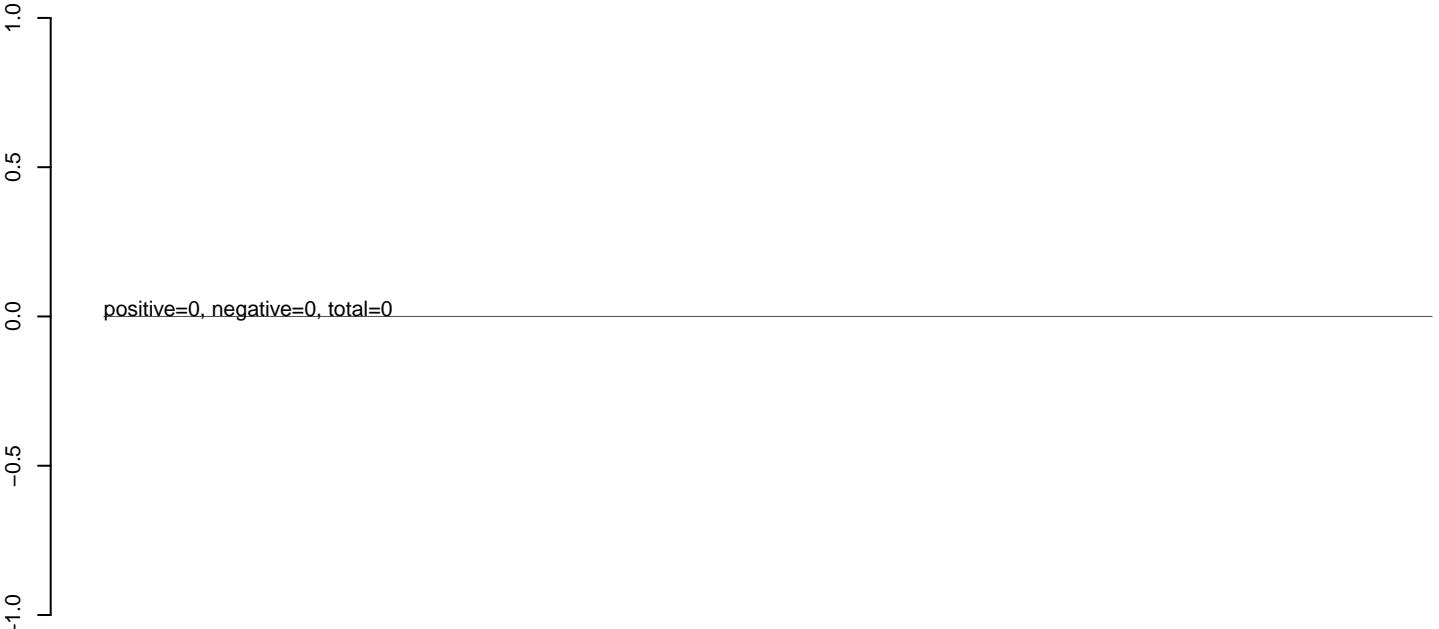
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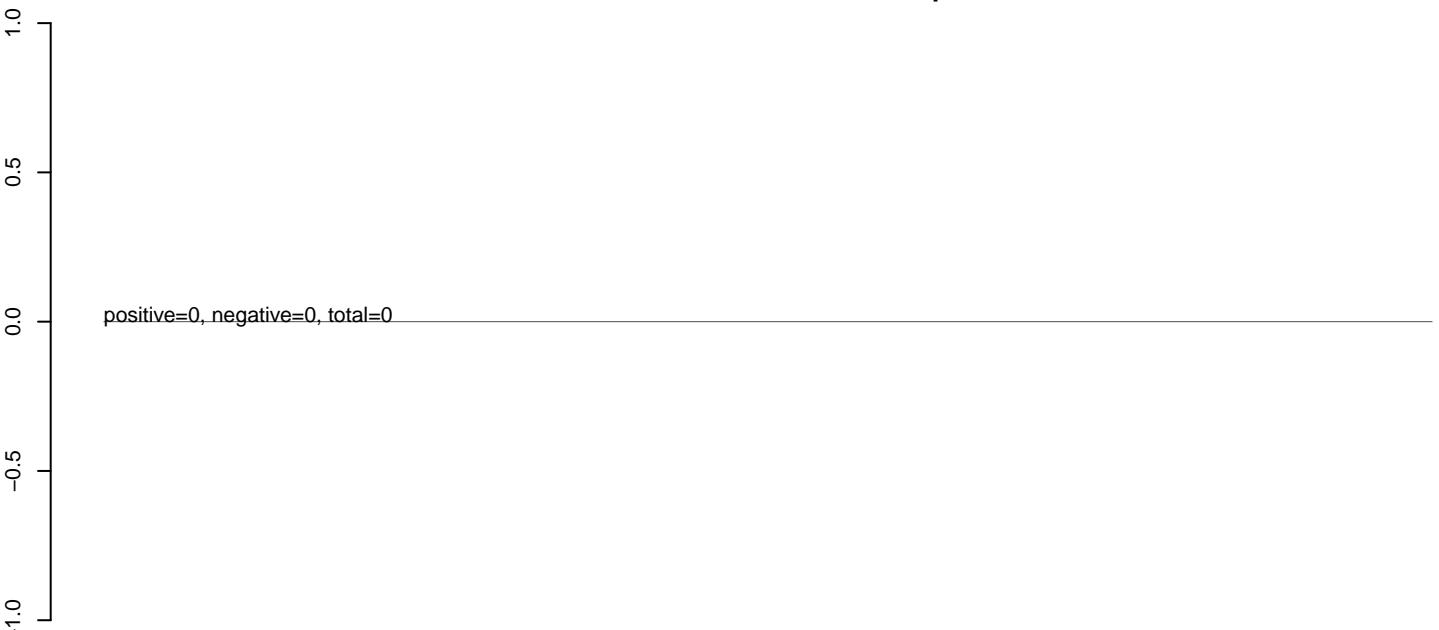
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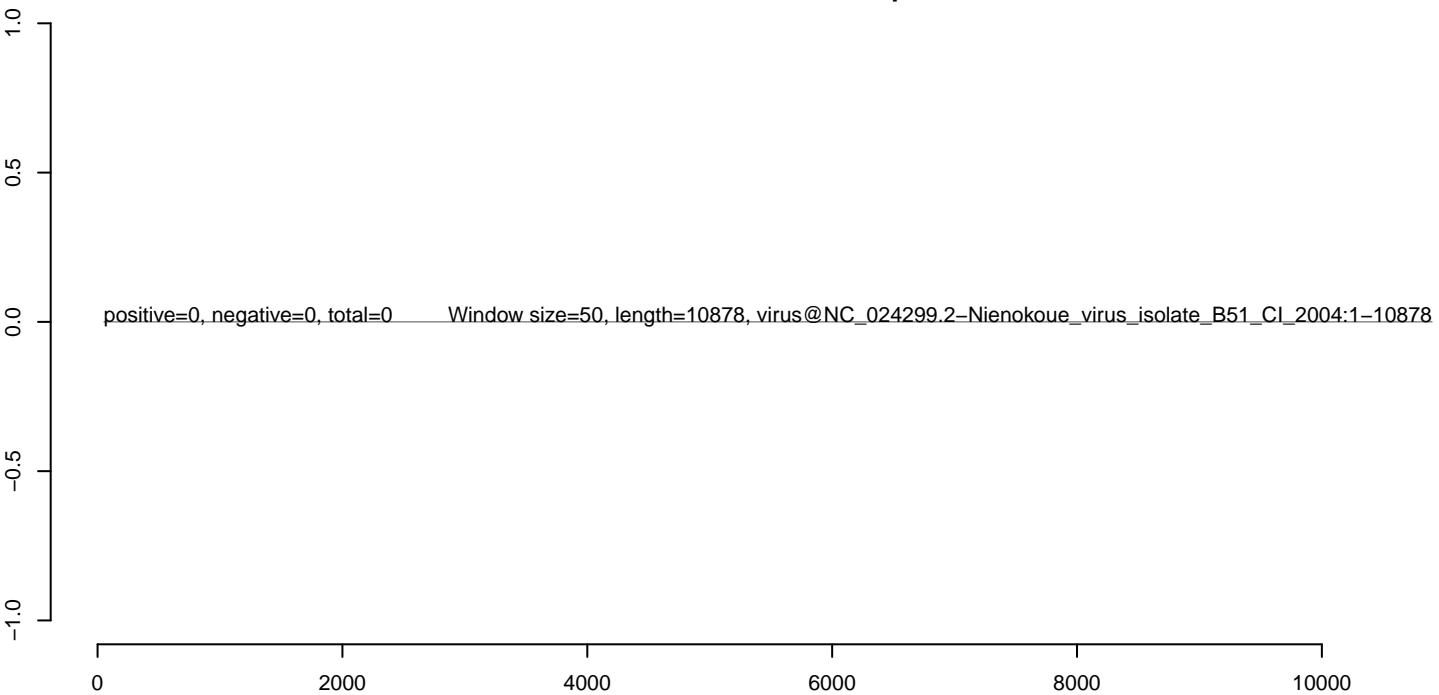
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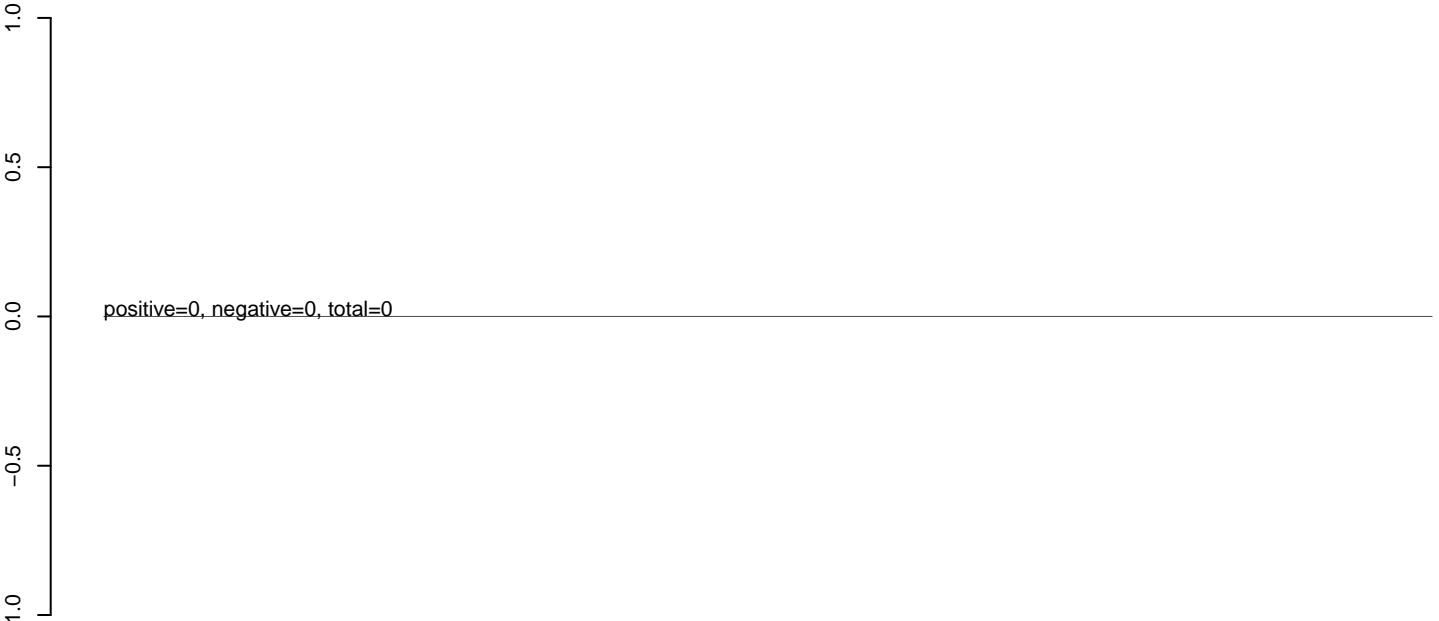
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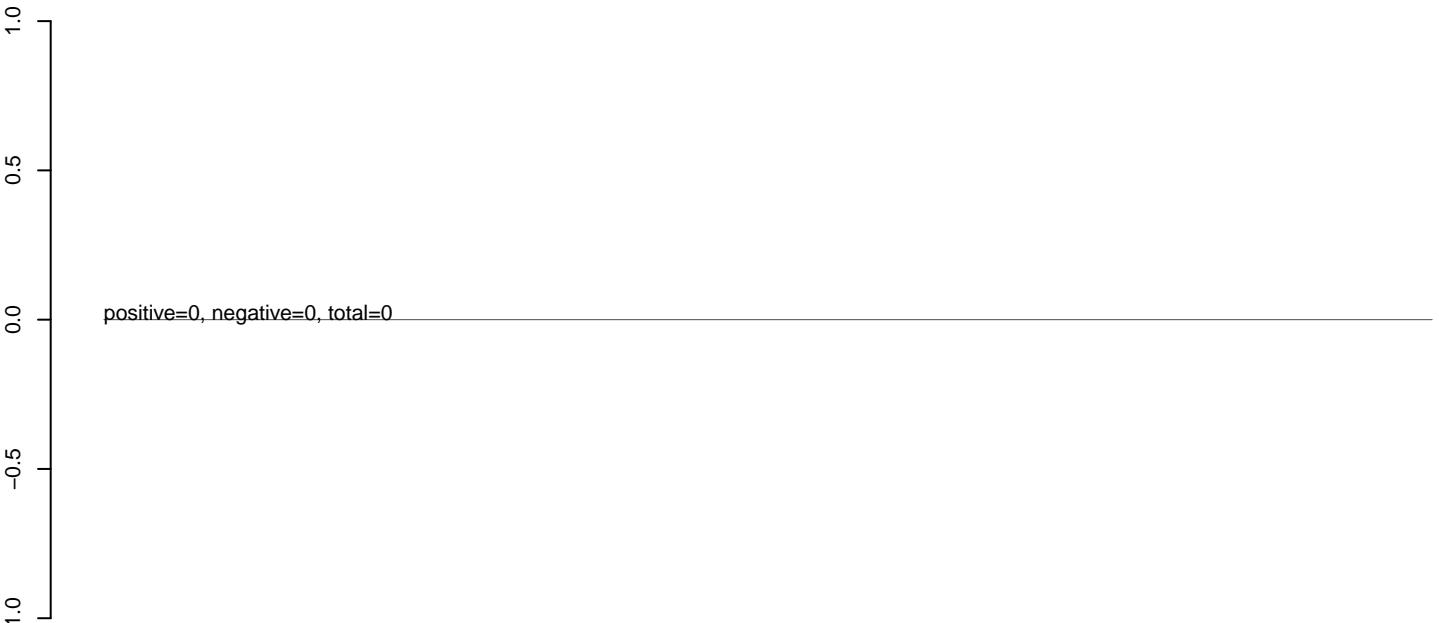
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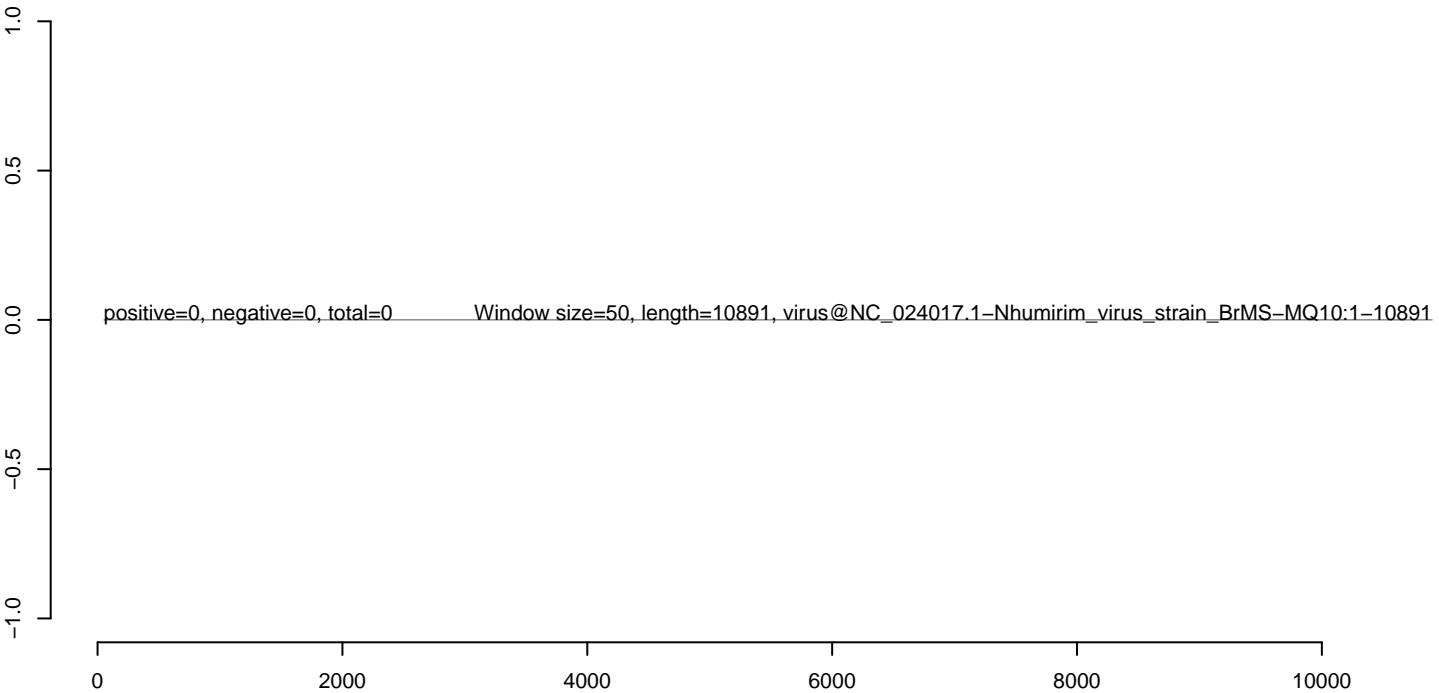
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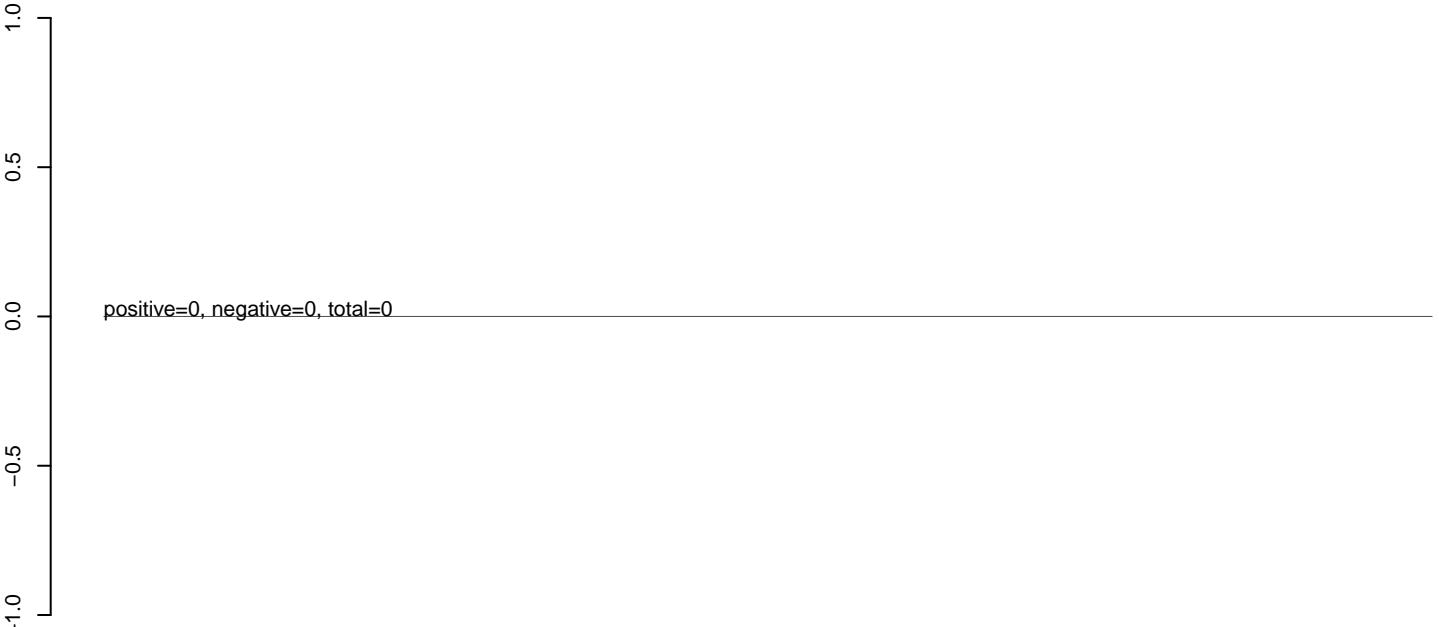
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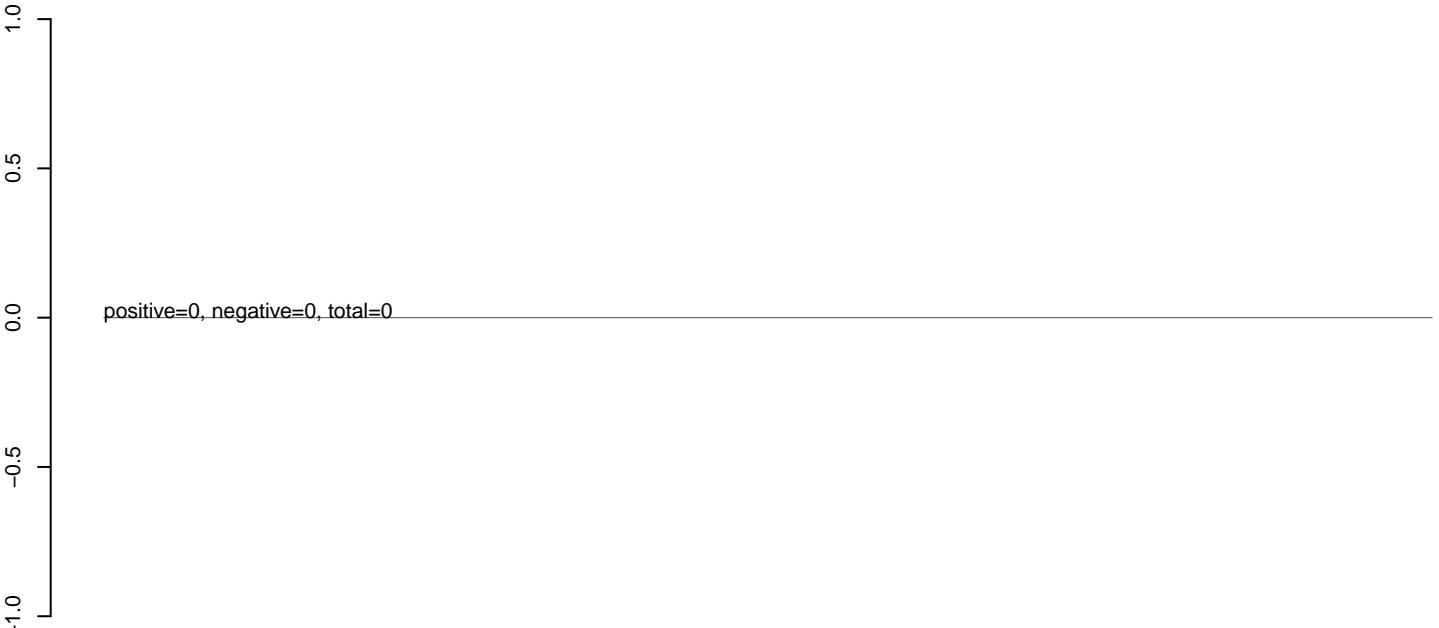
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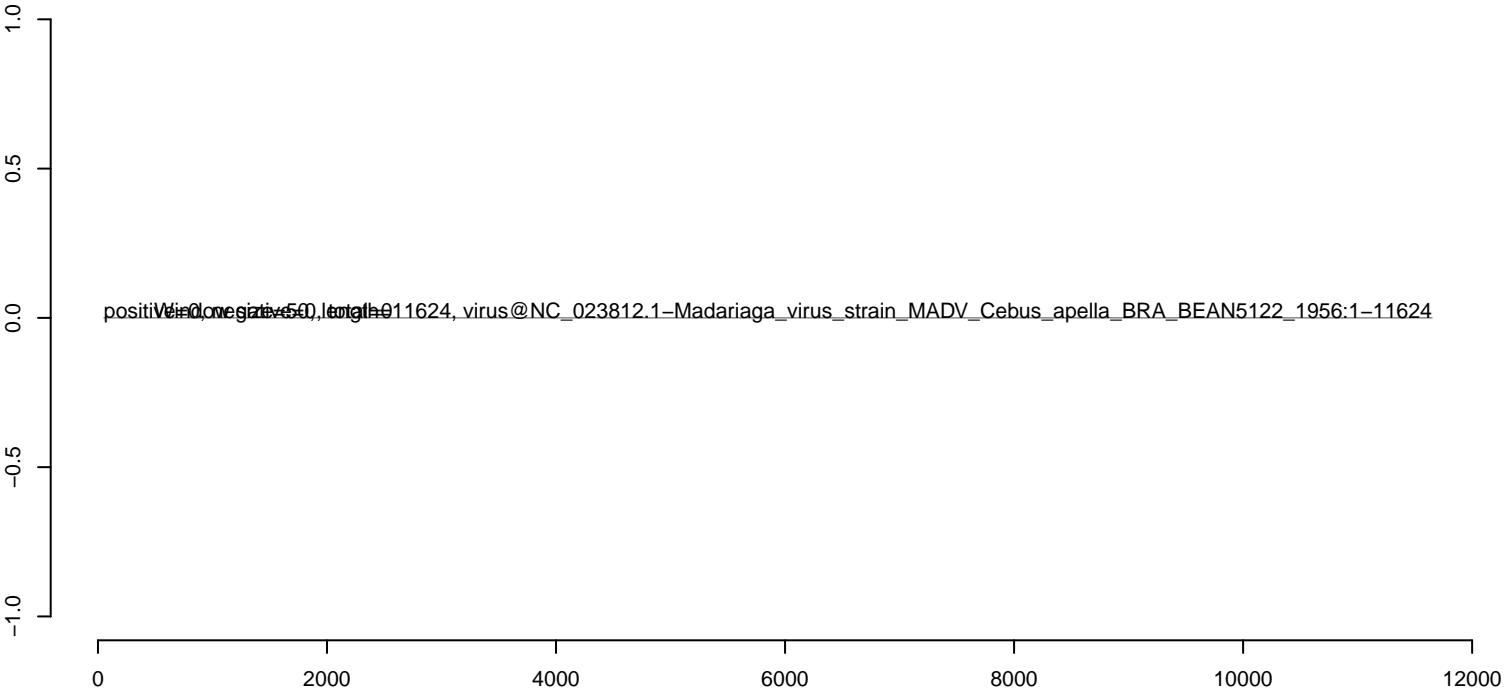
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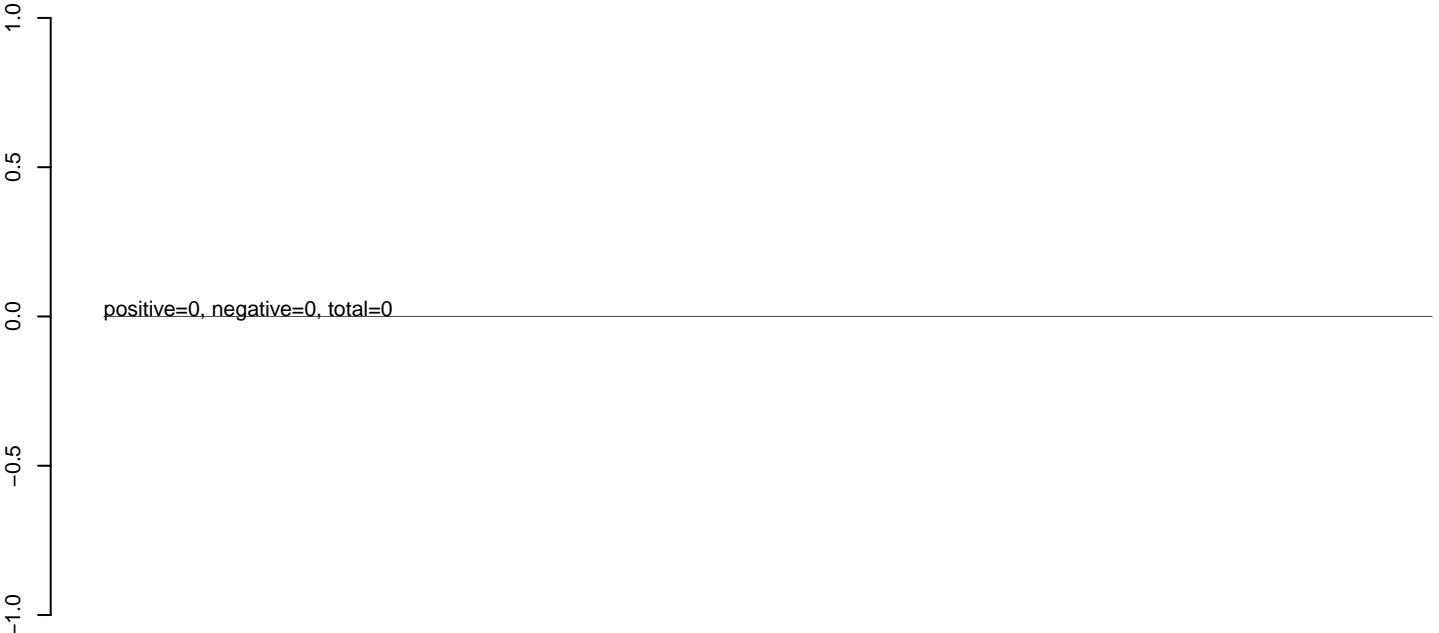
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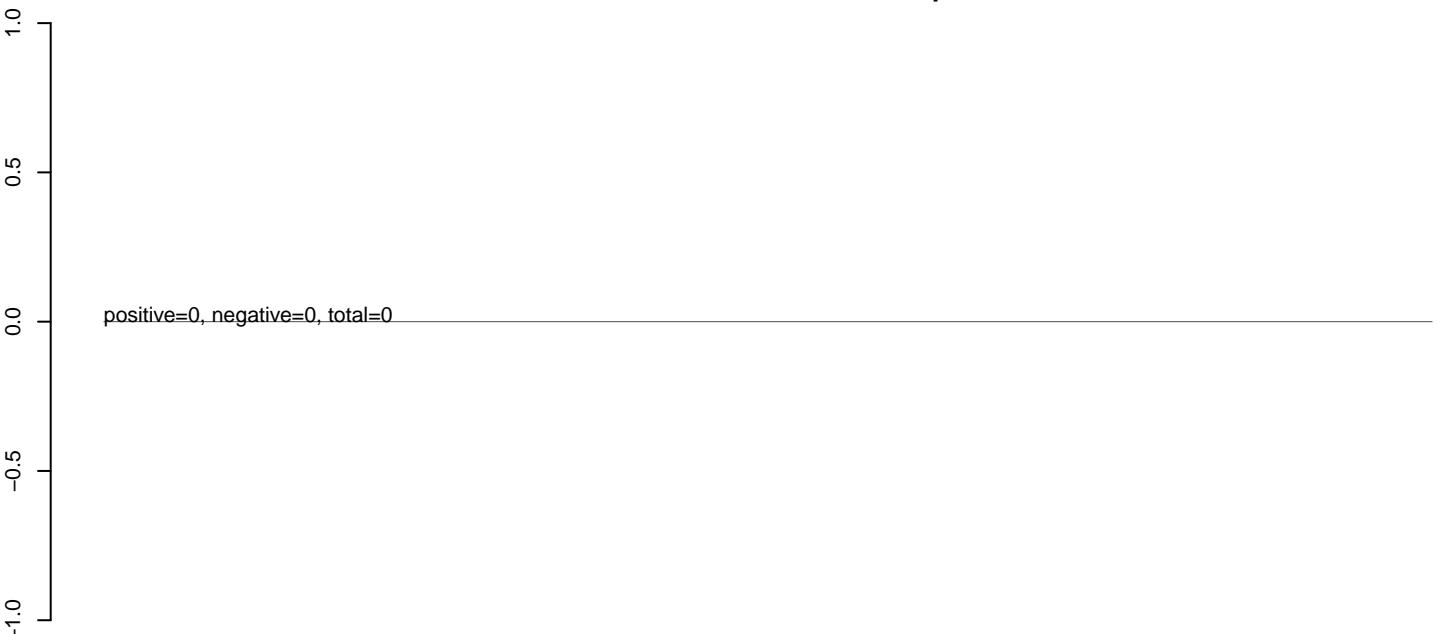
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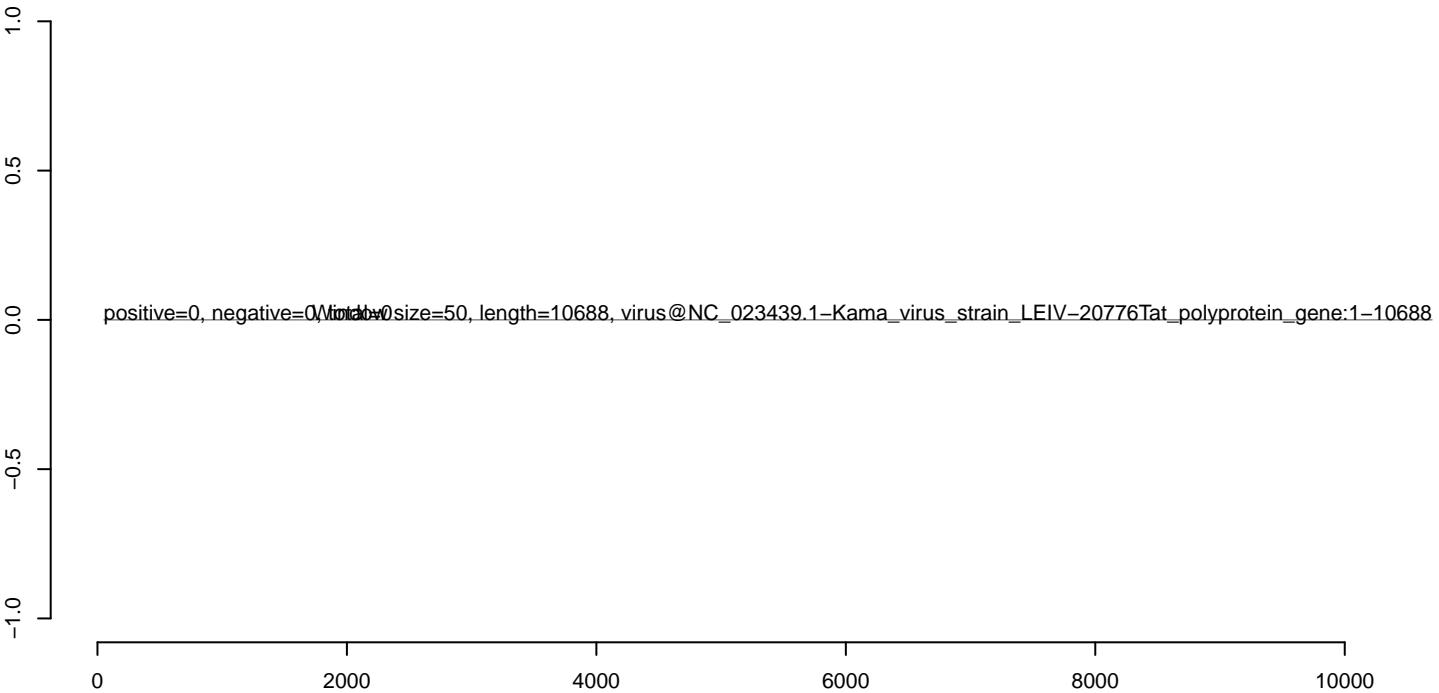
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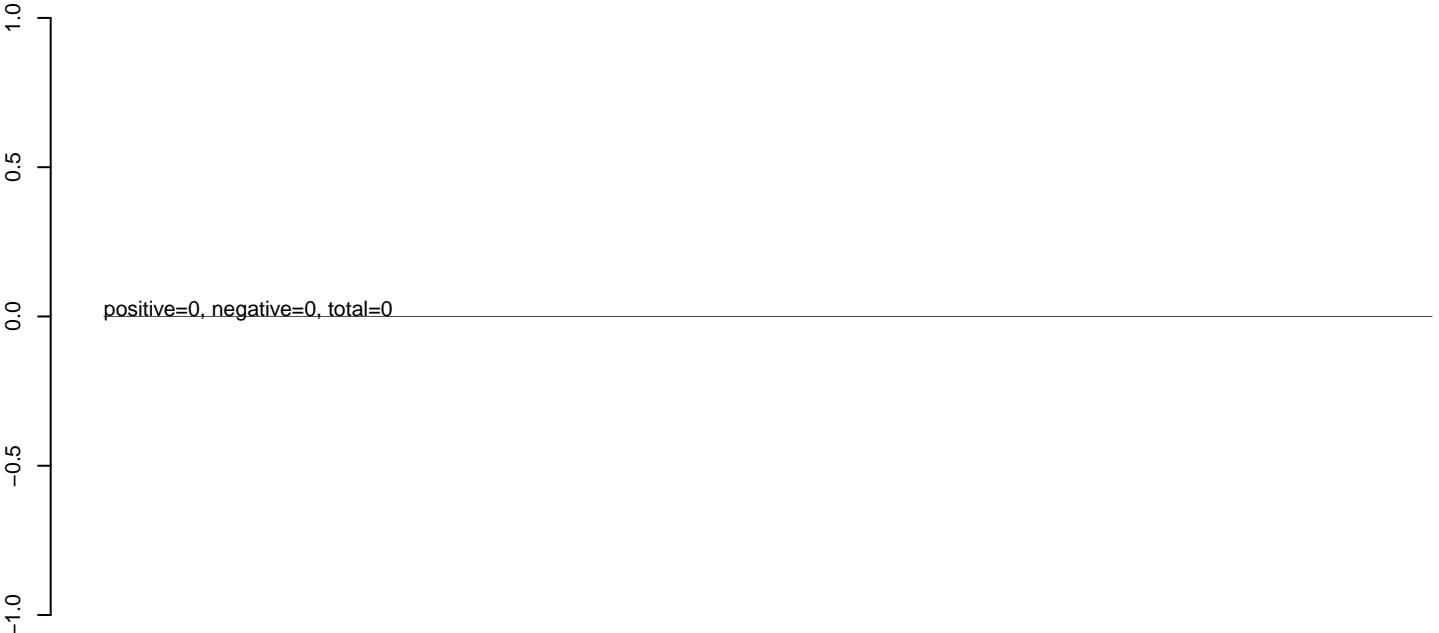
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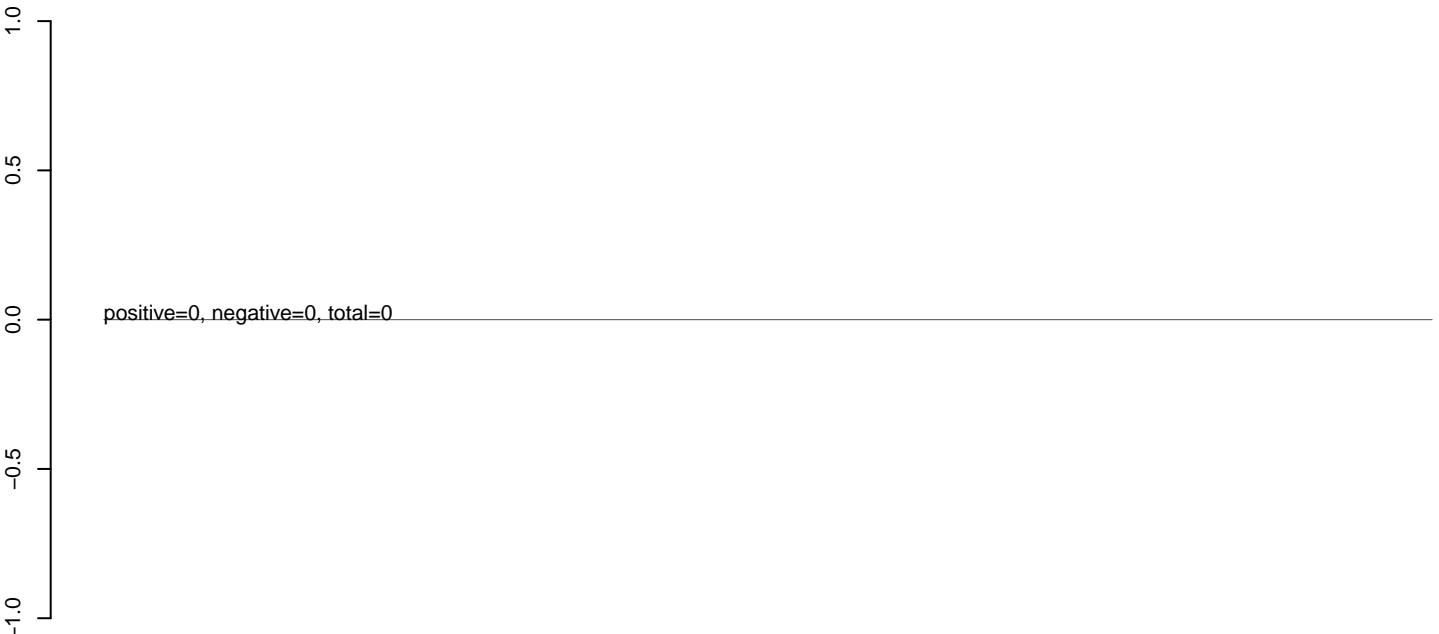
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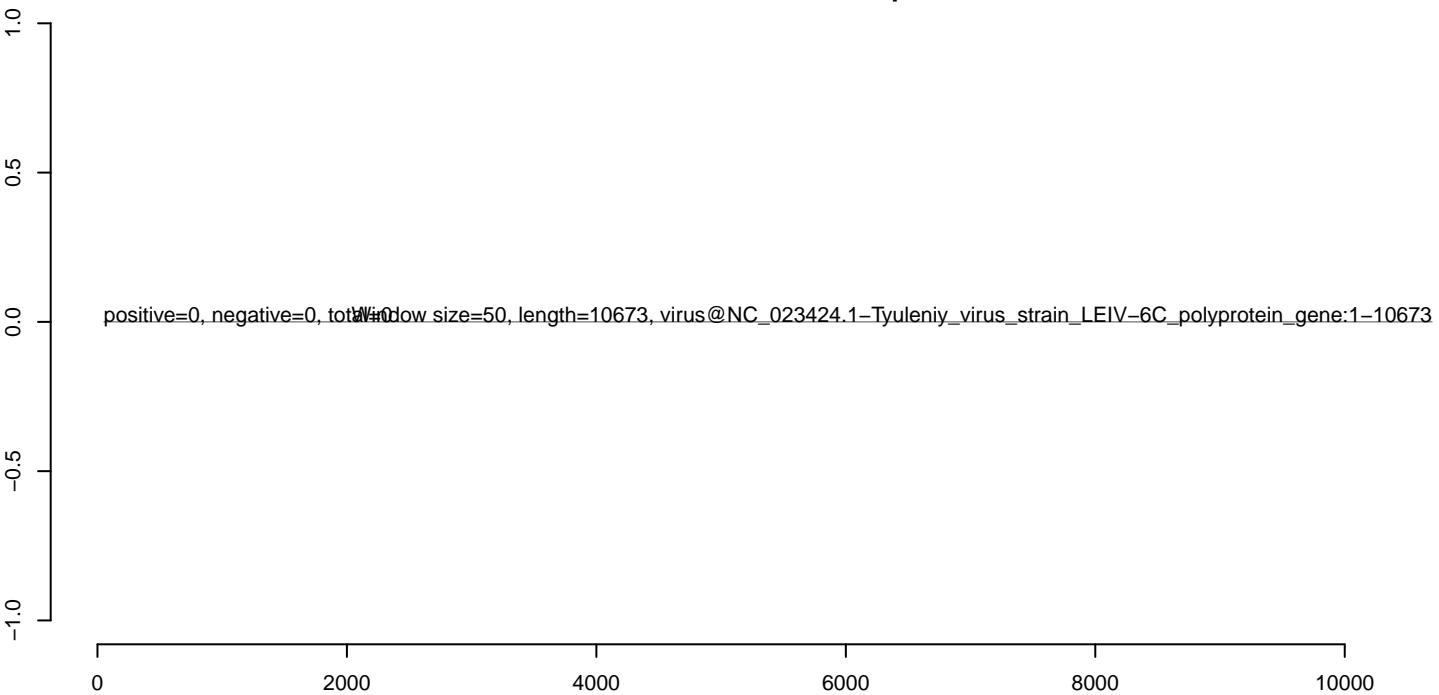
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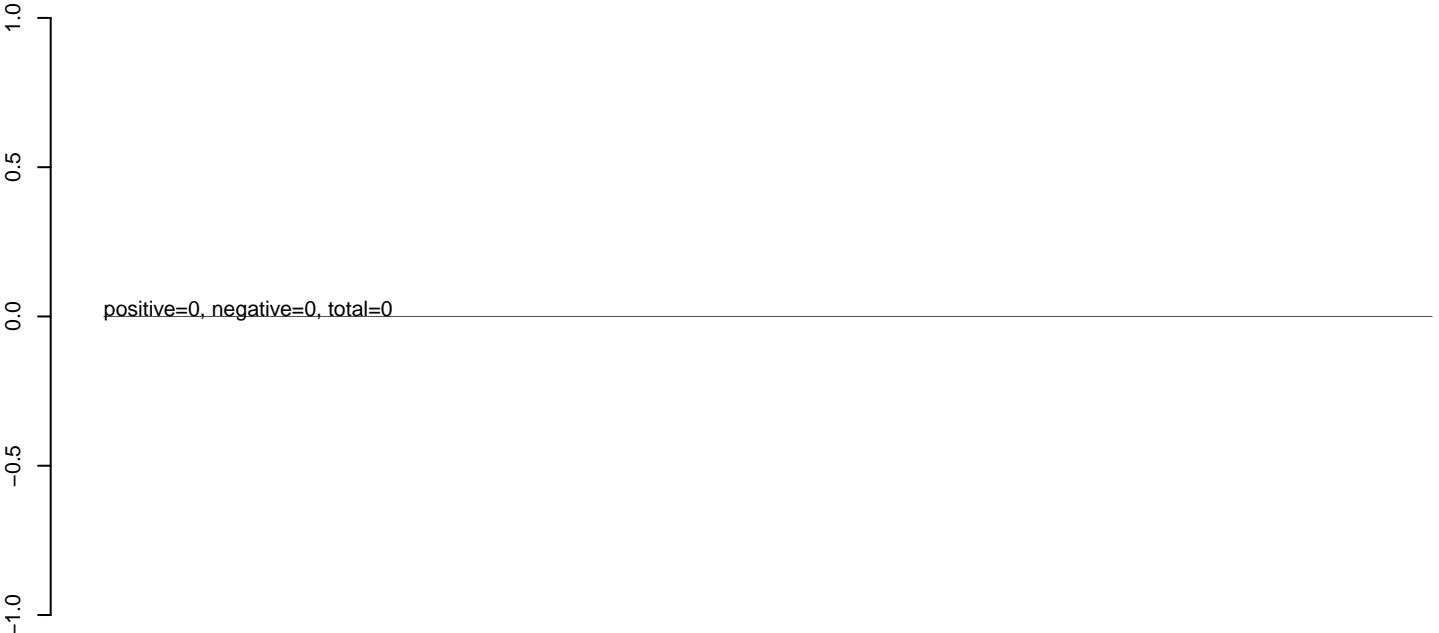
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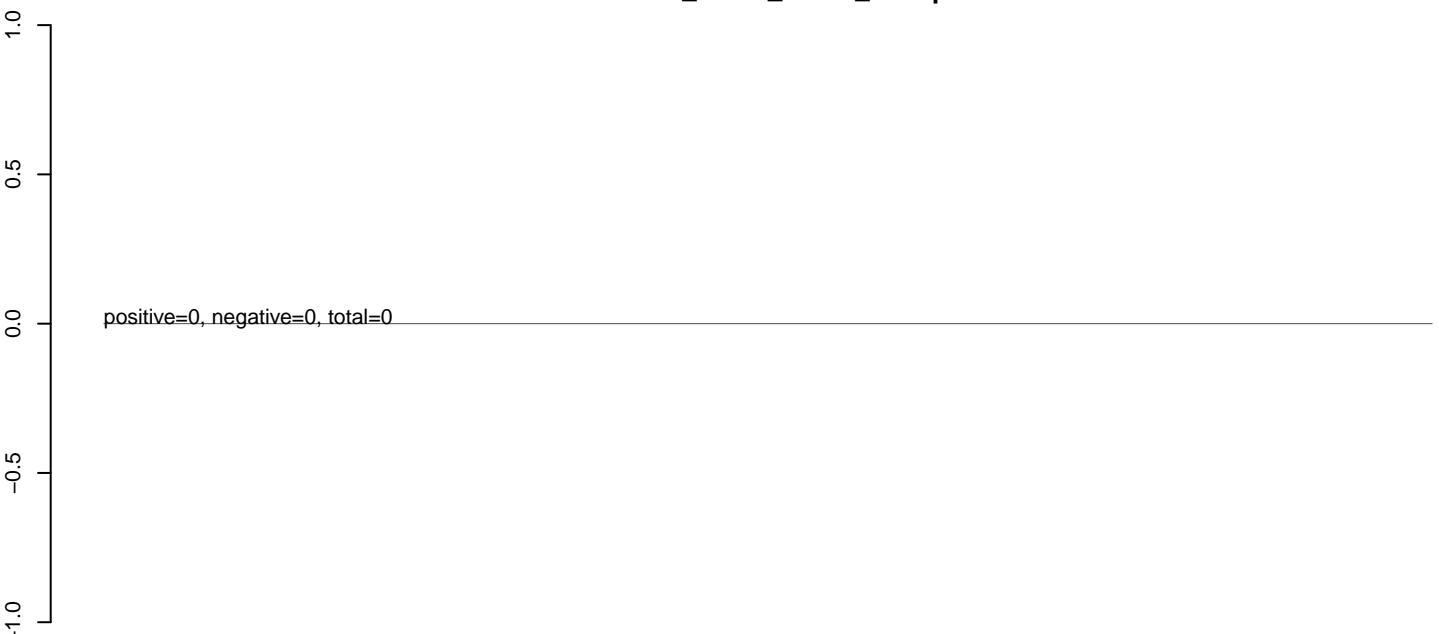
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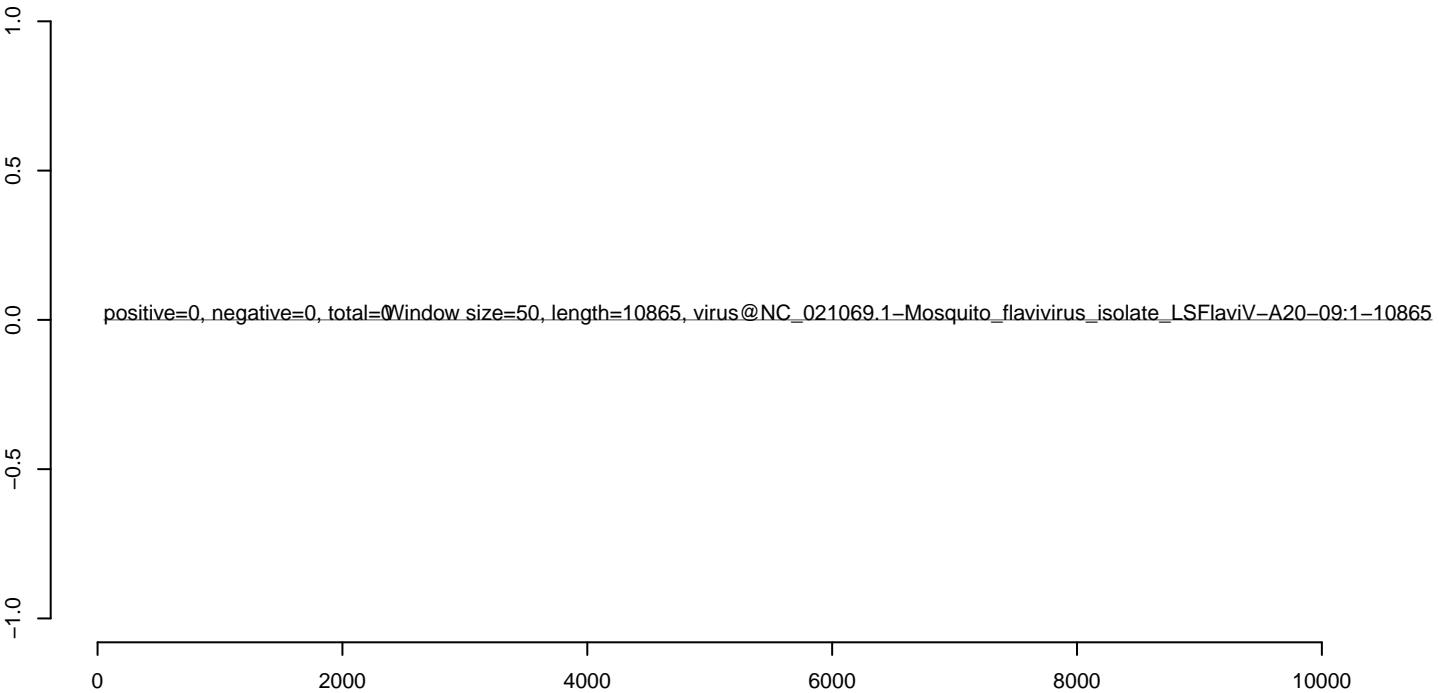
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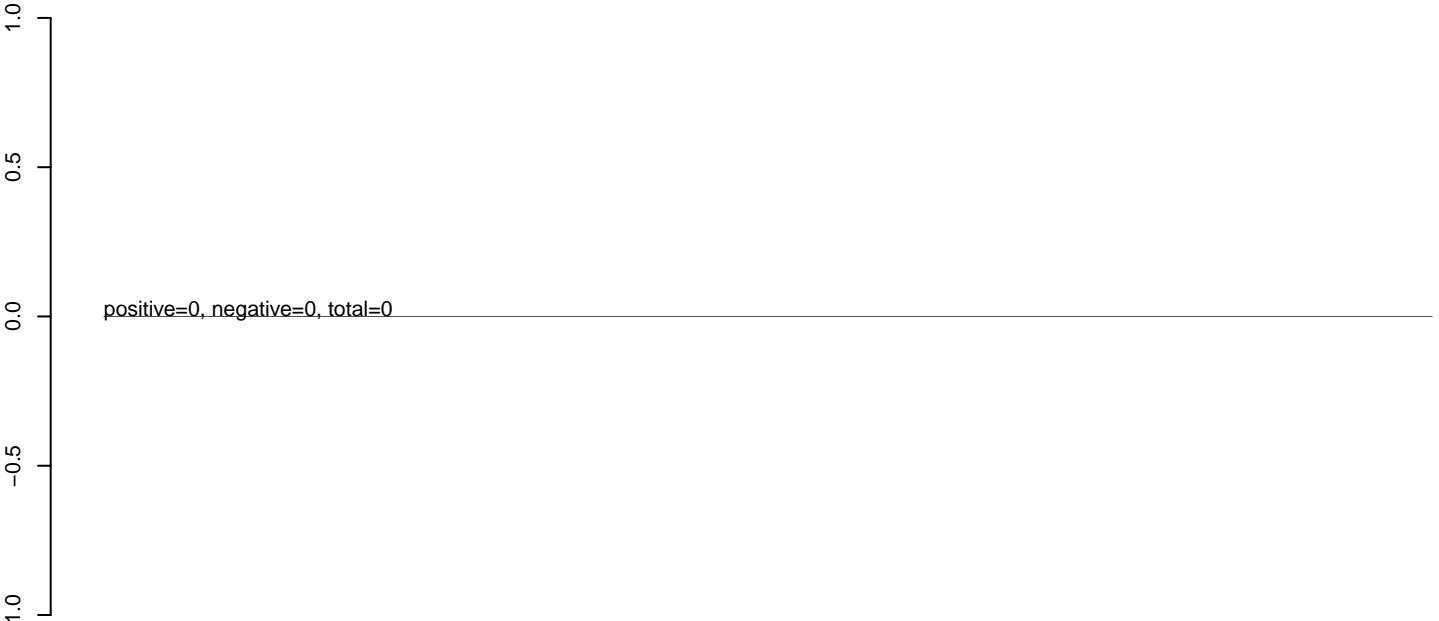
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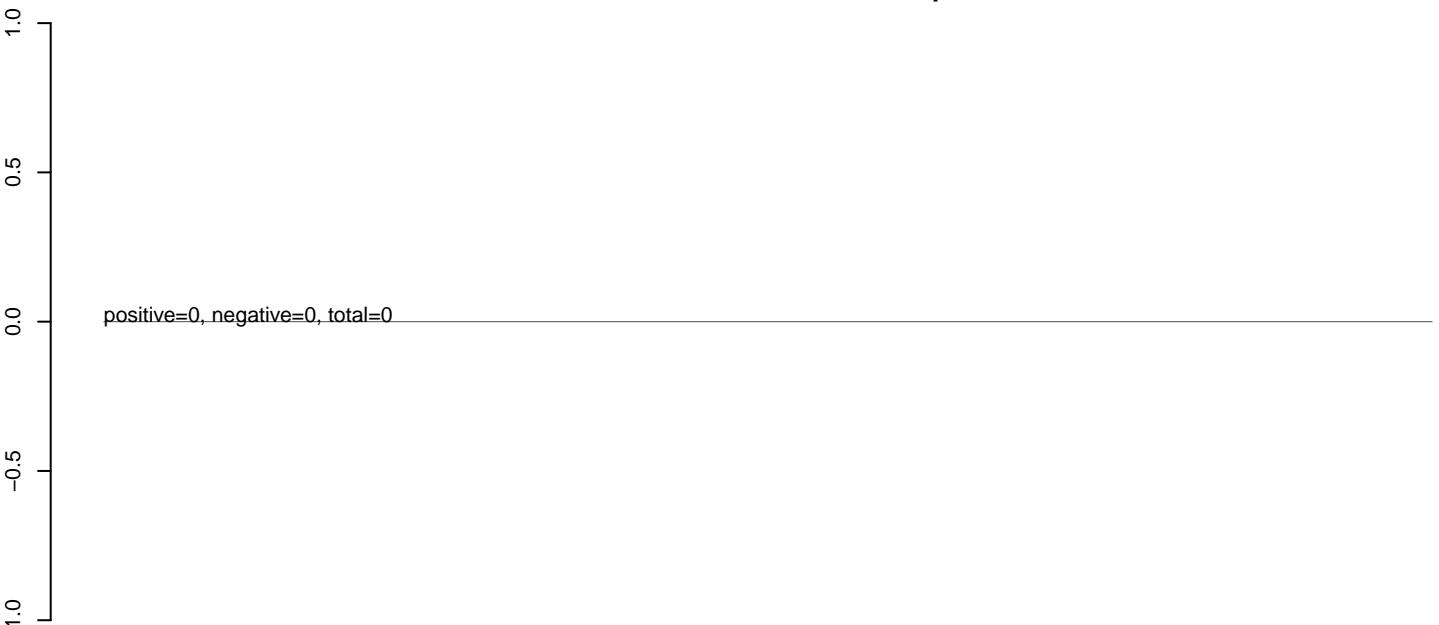
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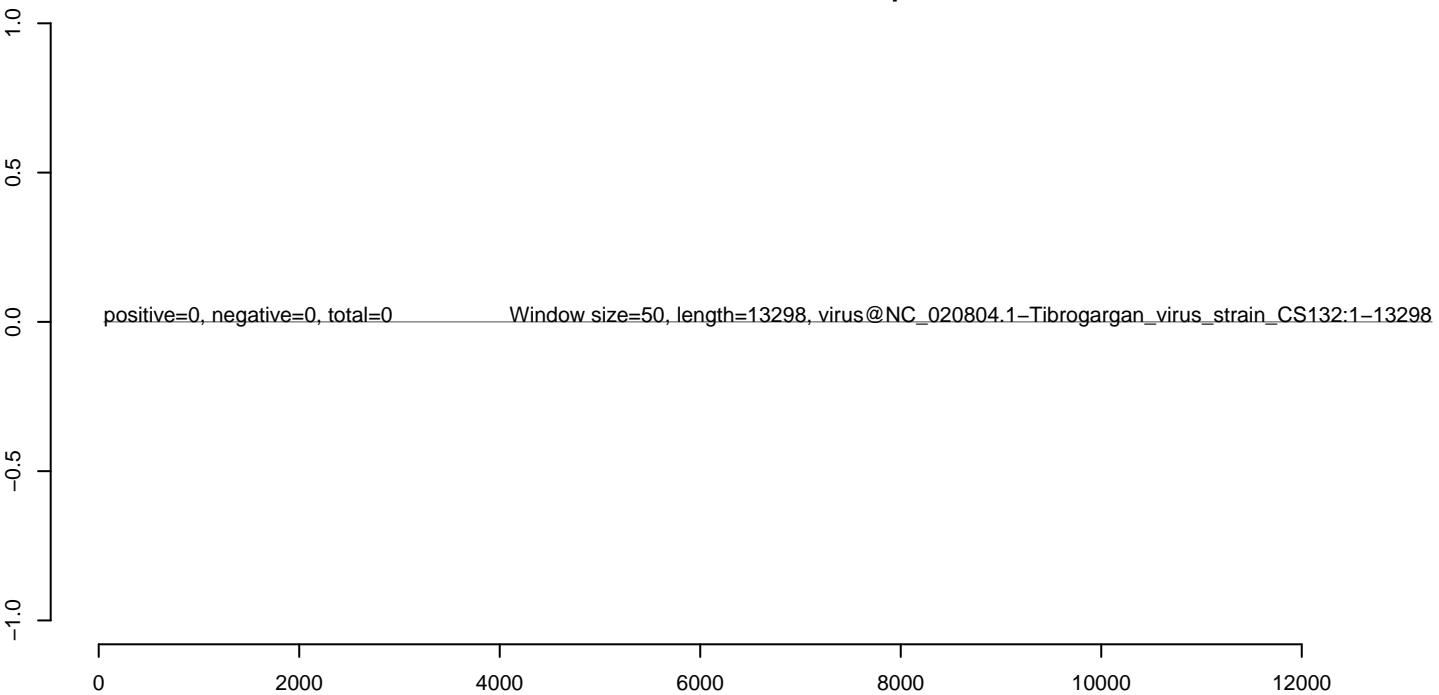
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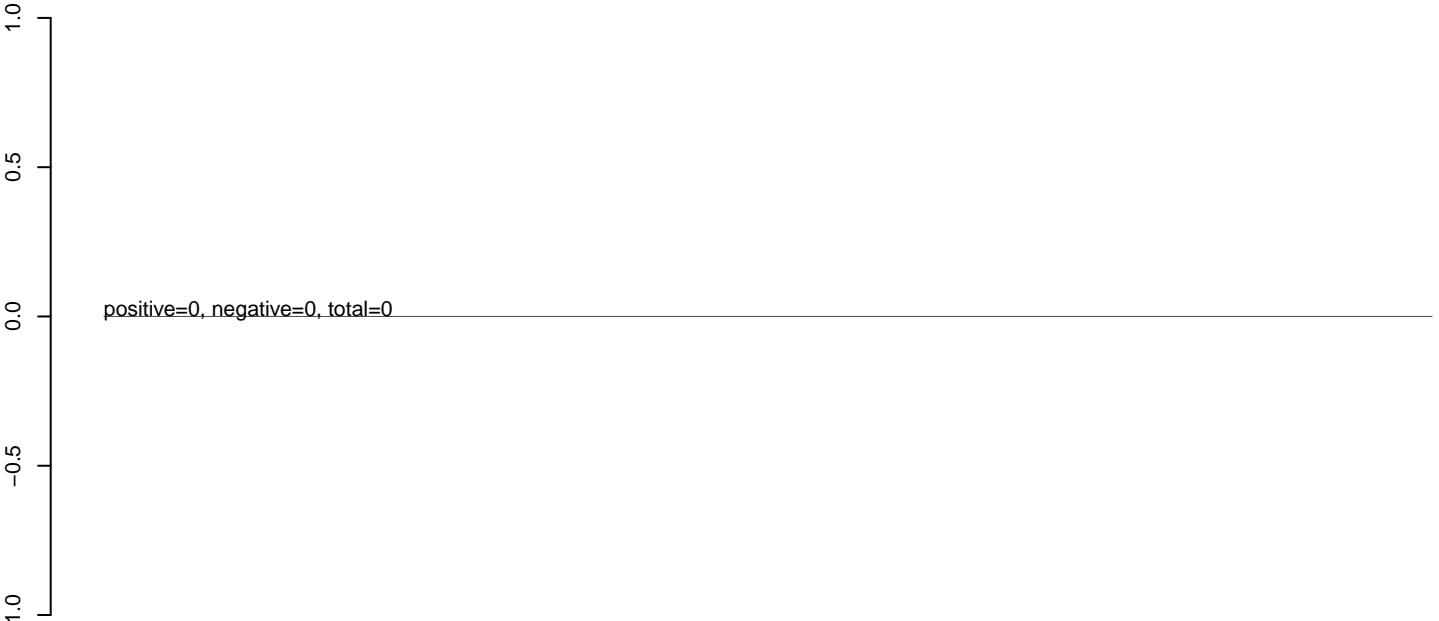
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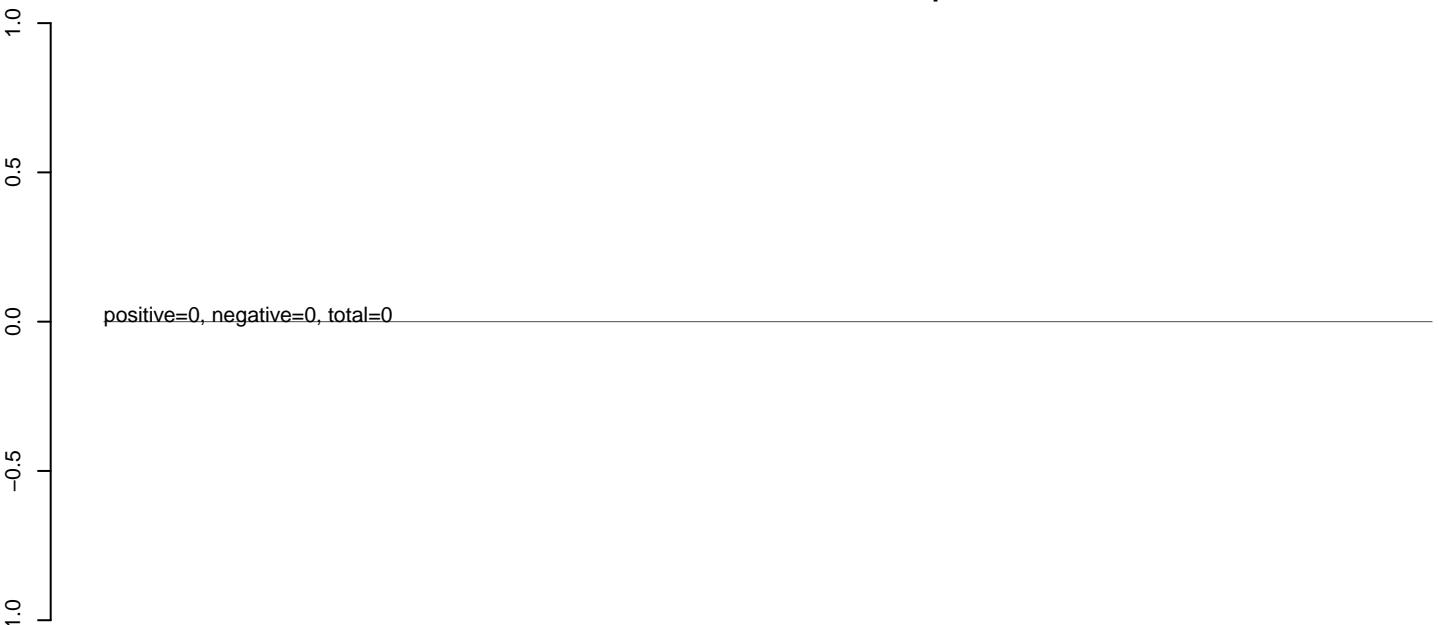
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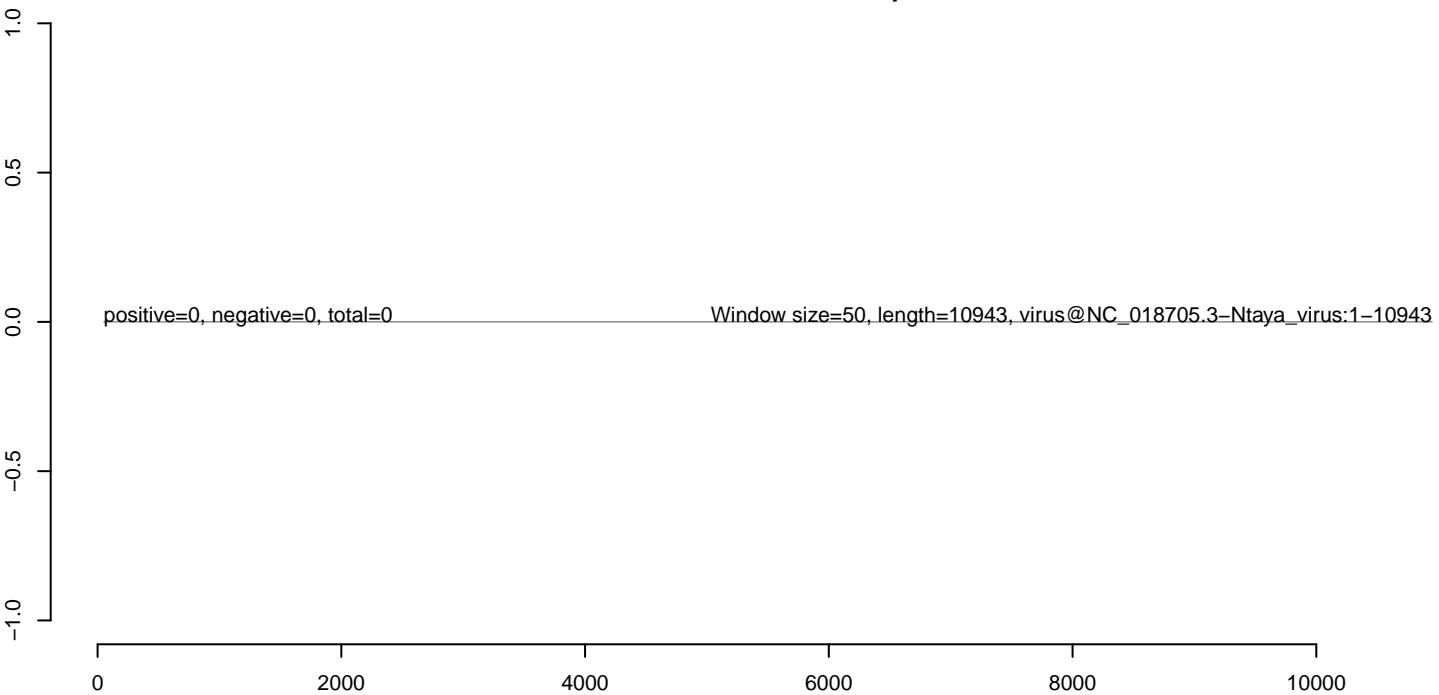
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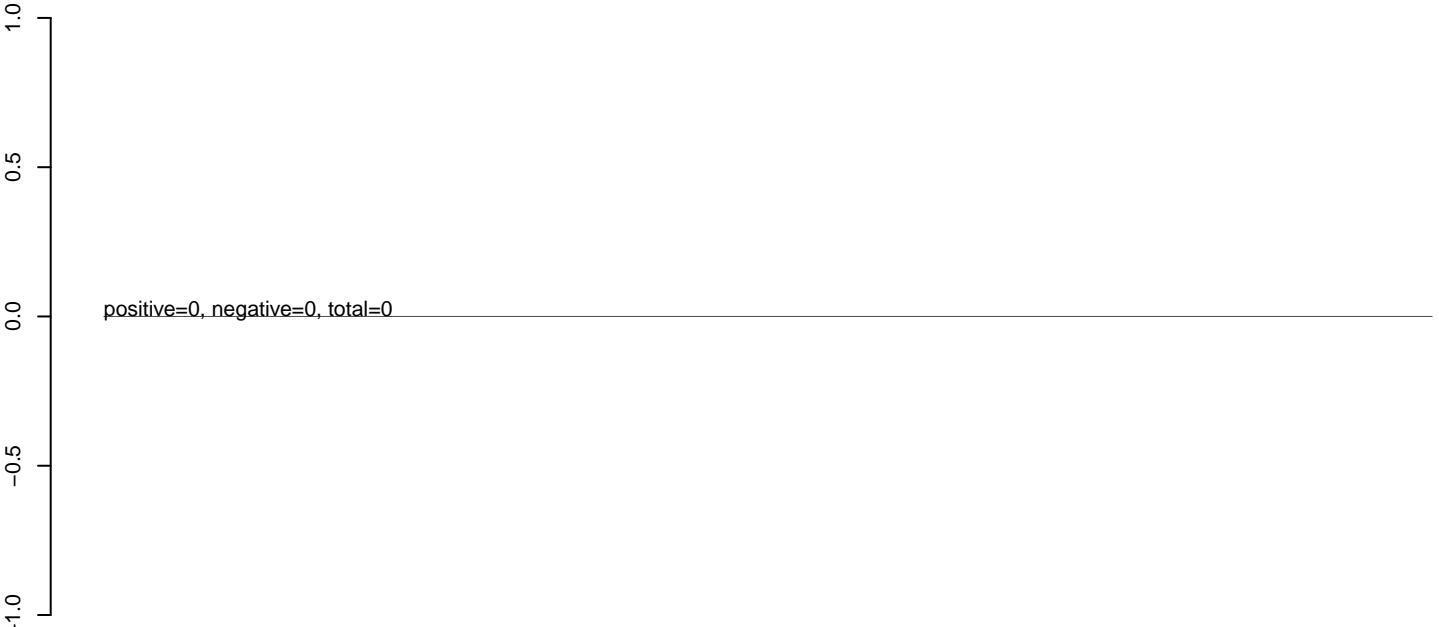
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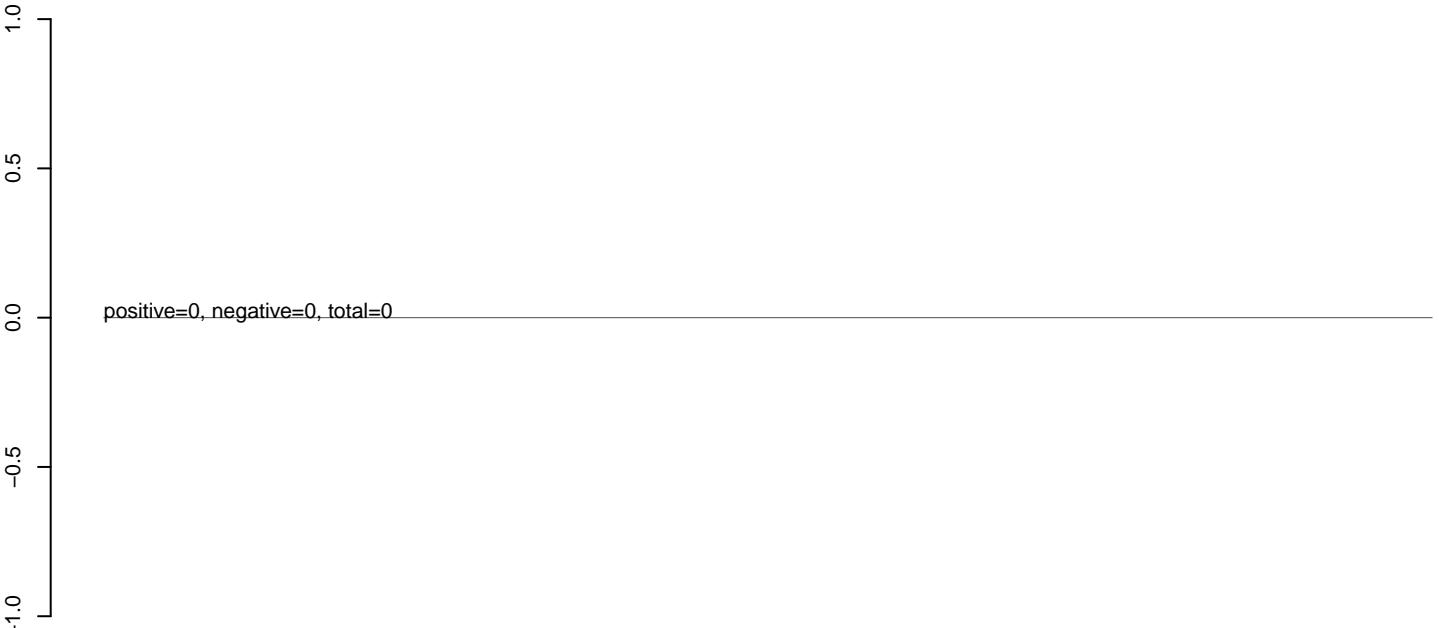
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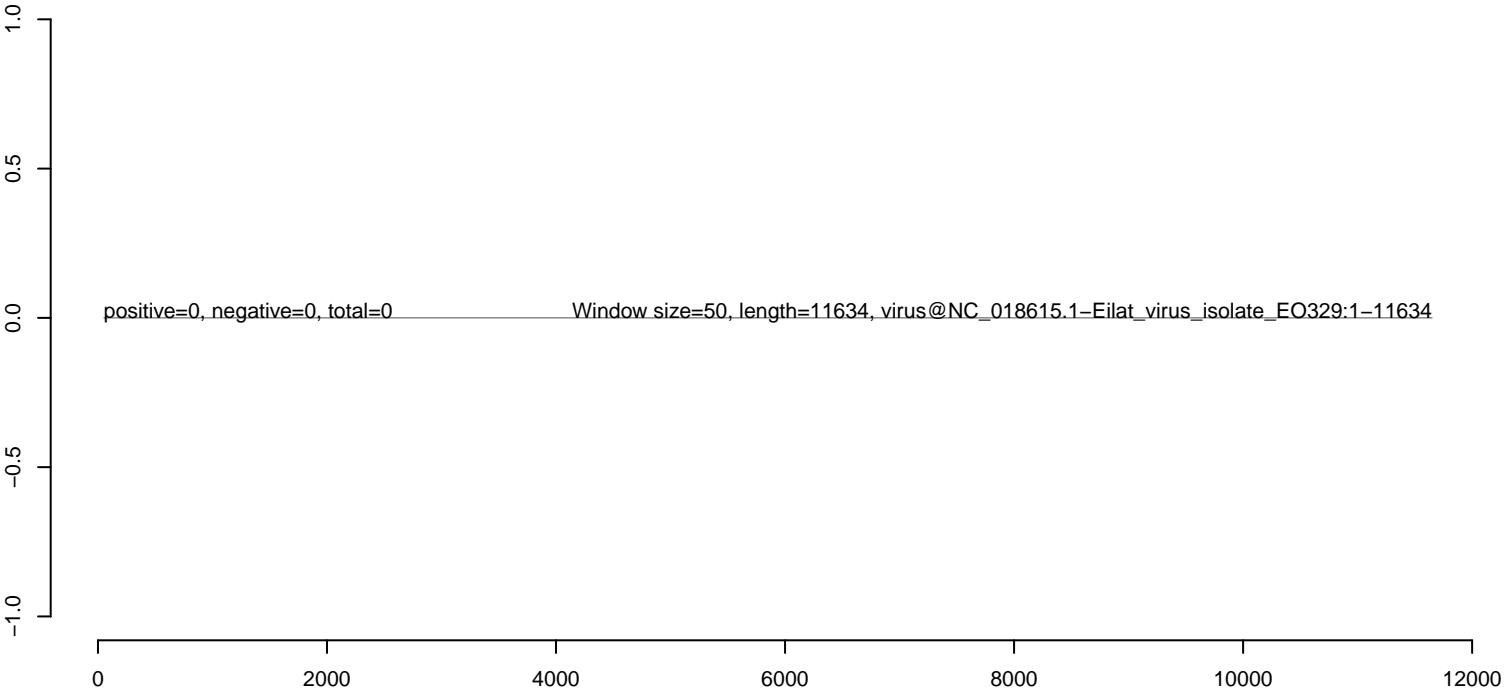
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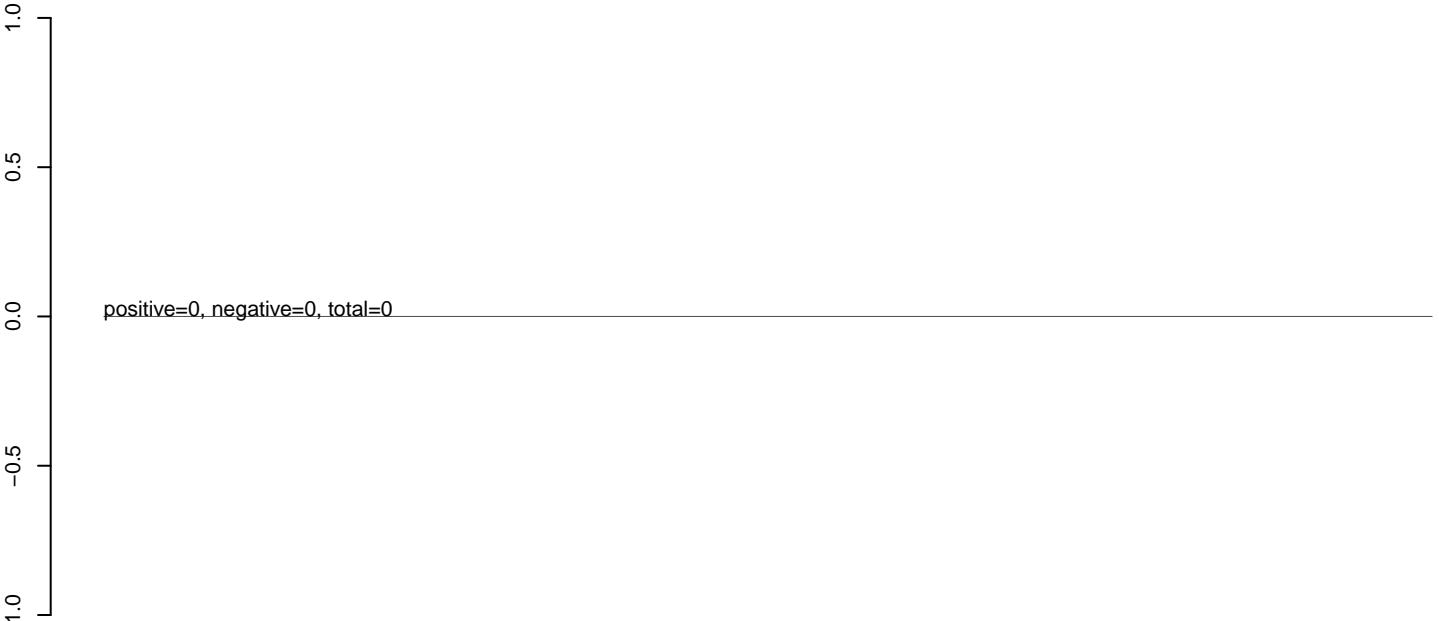
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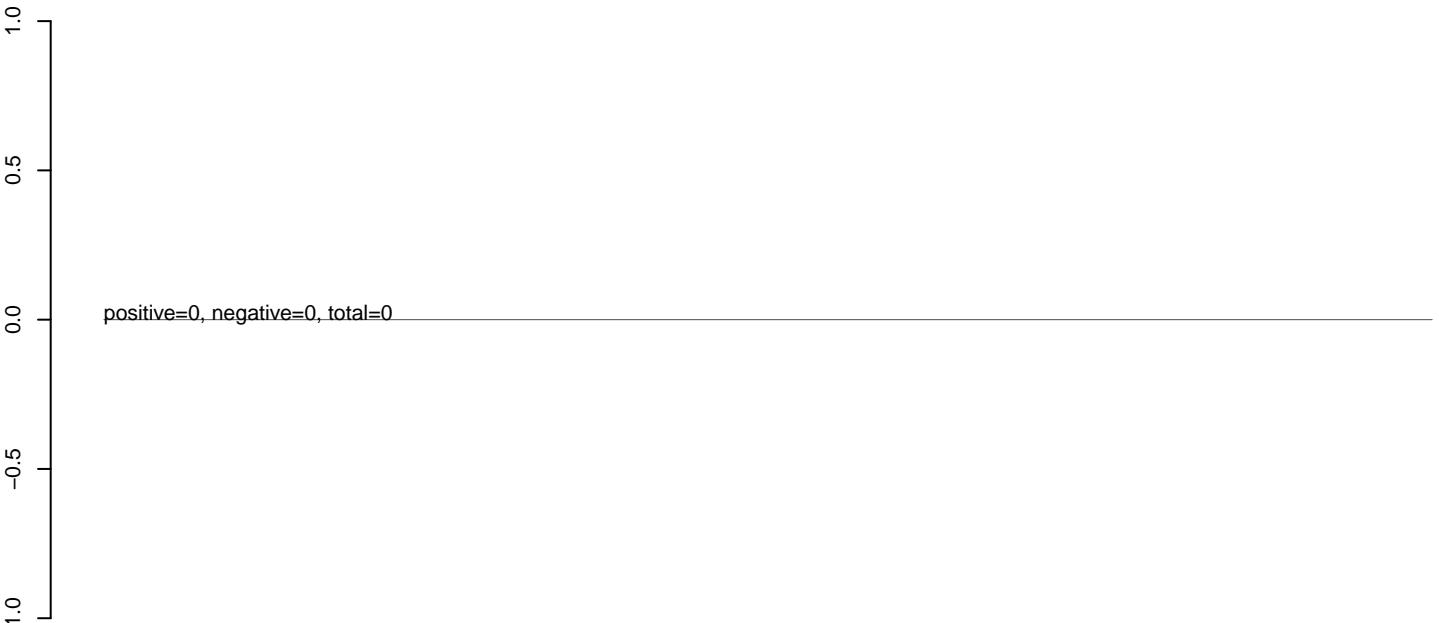
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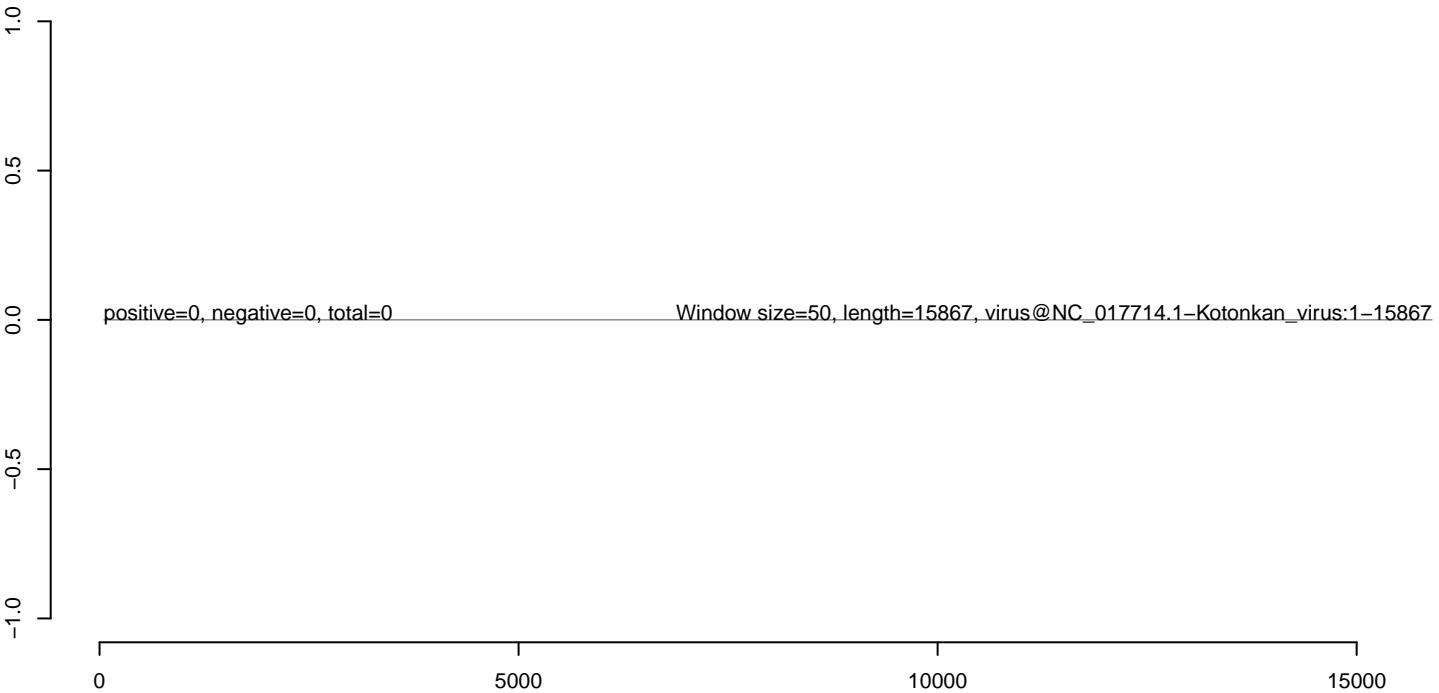
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### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

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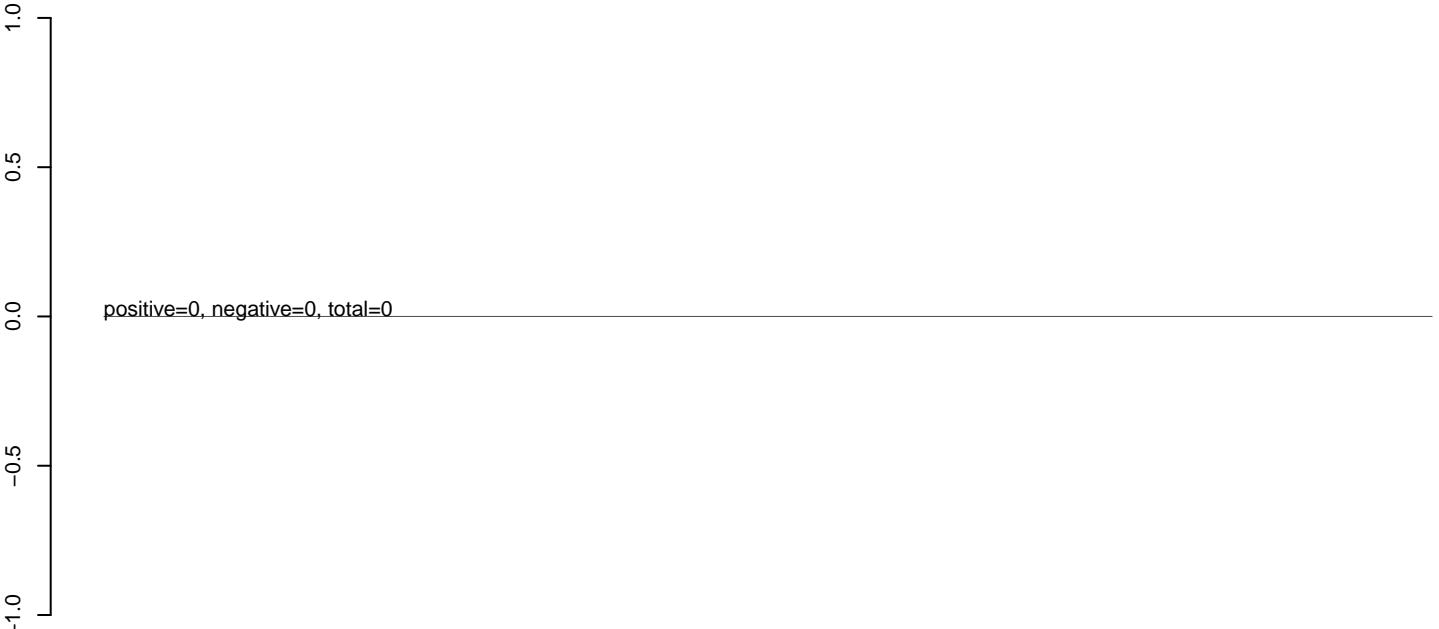
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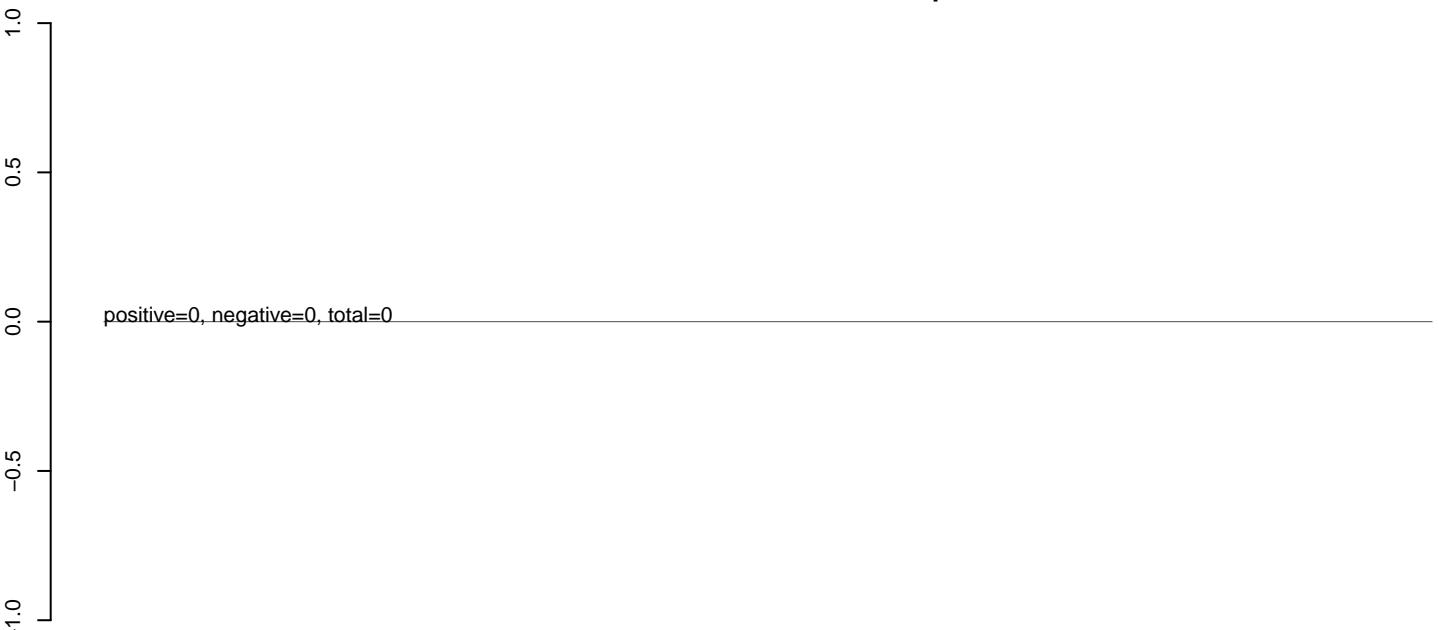
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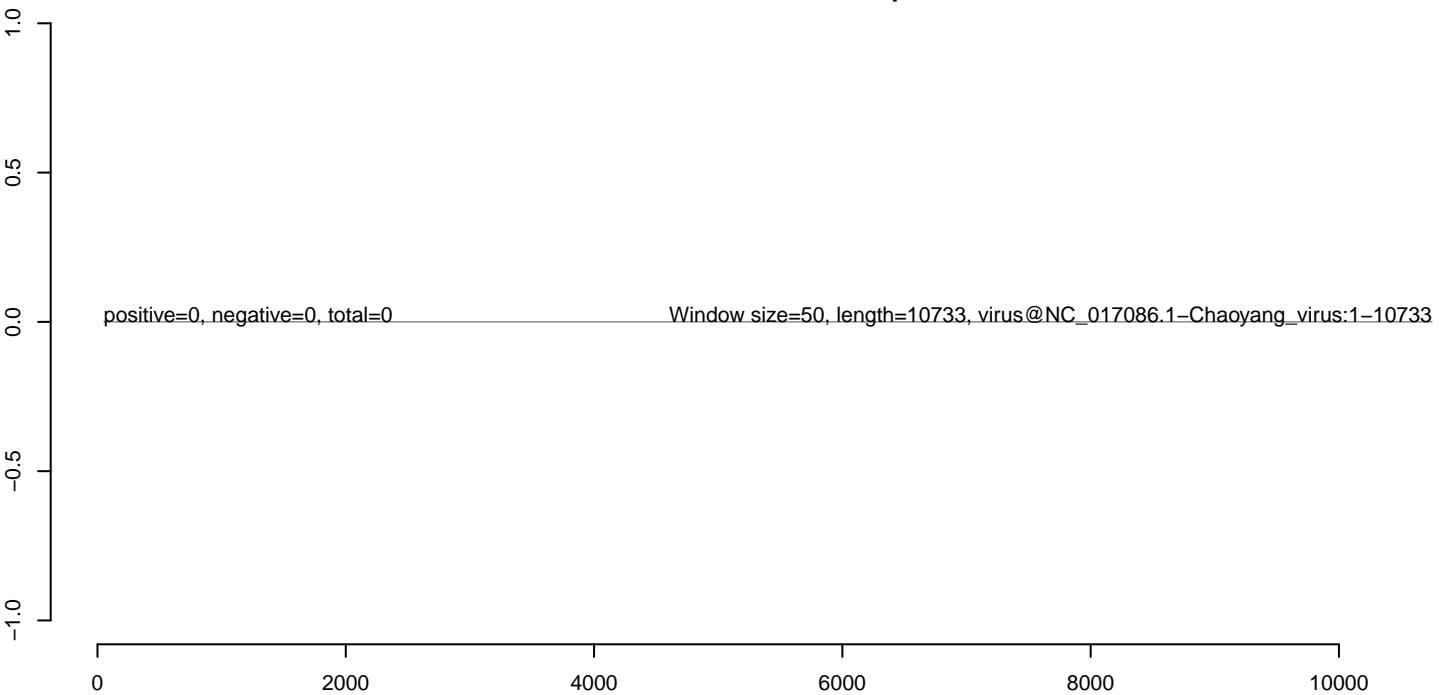
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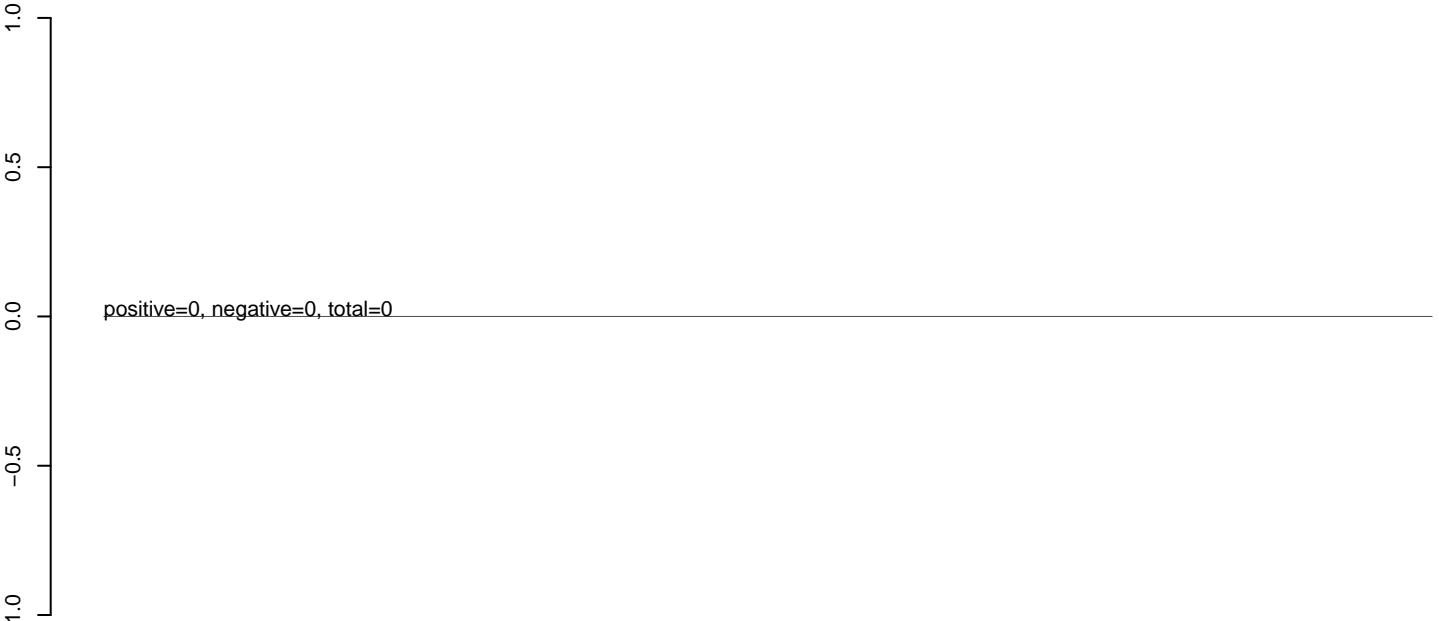
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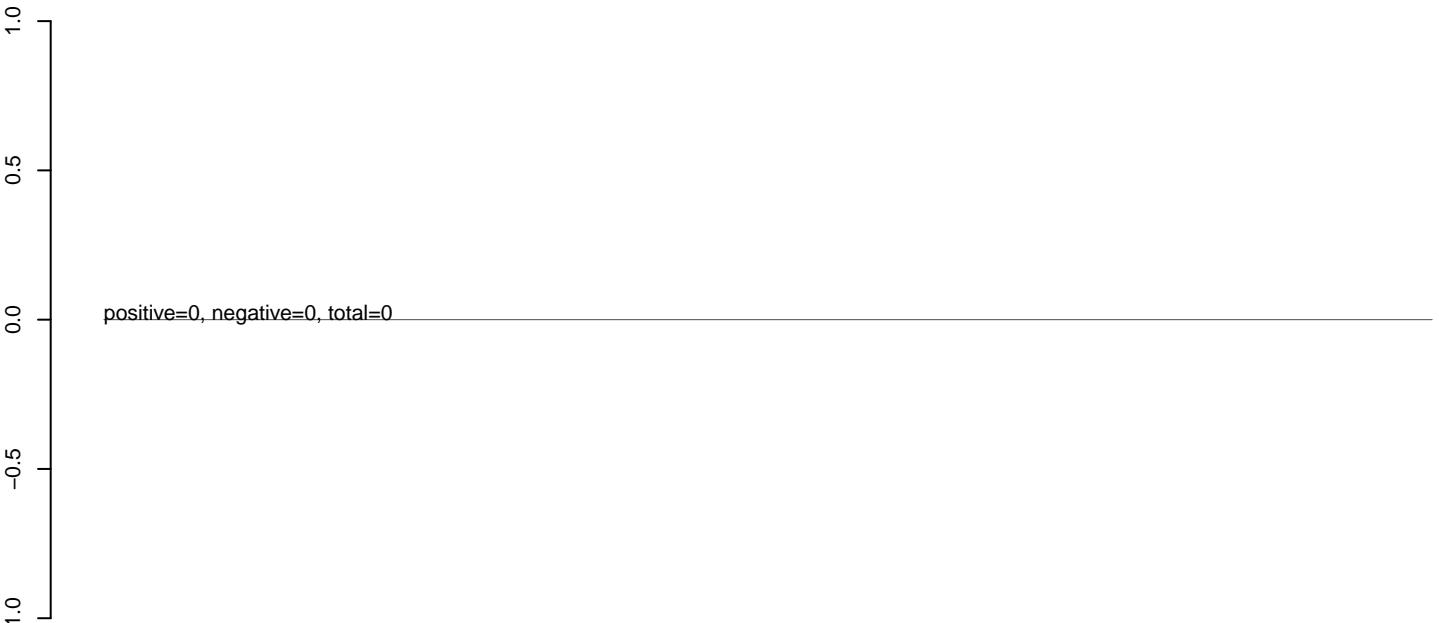
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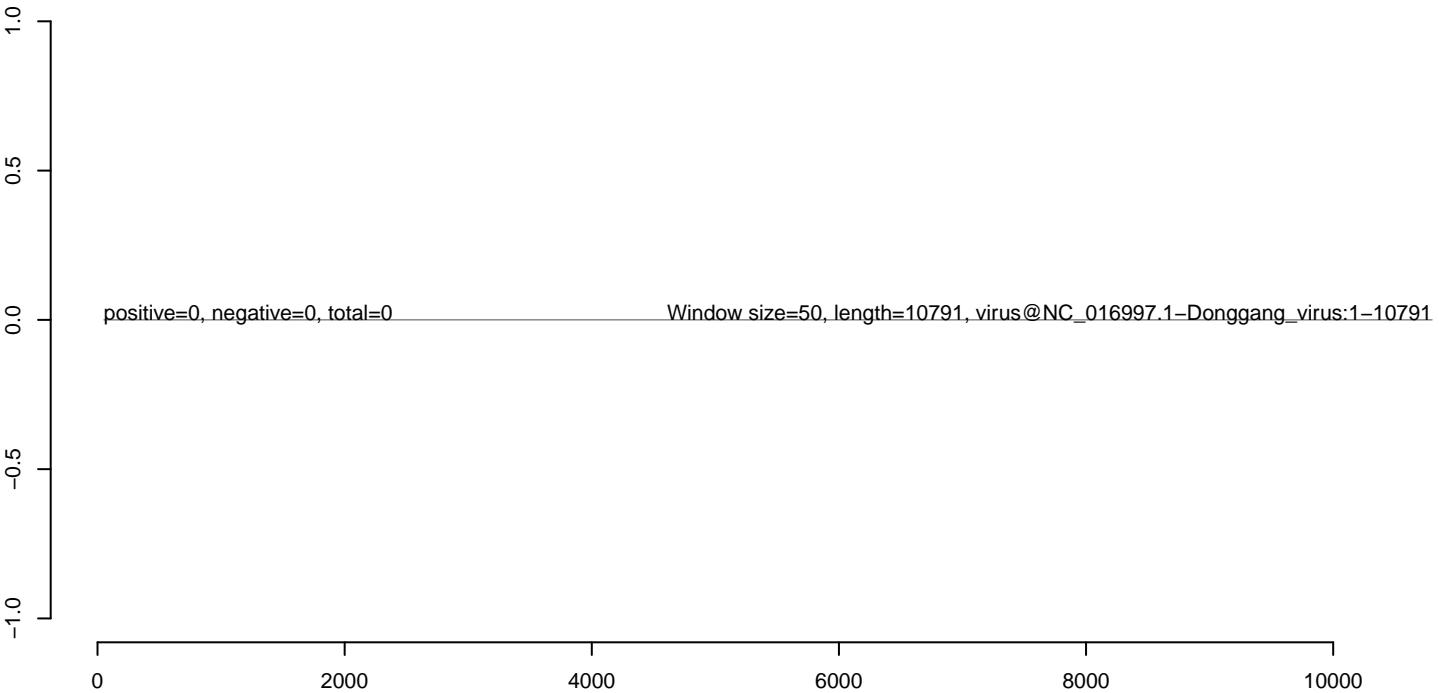
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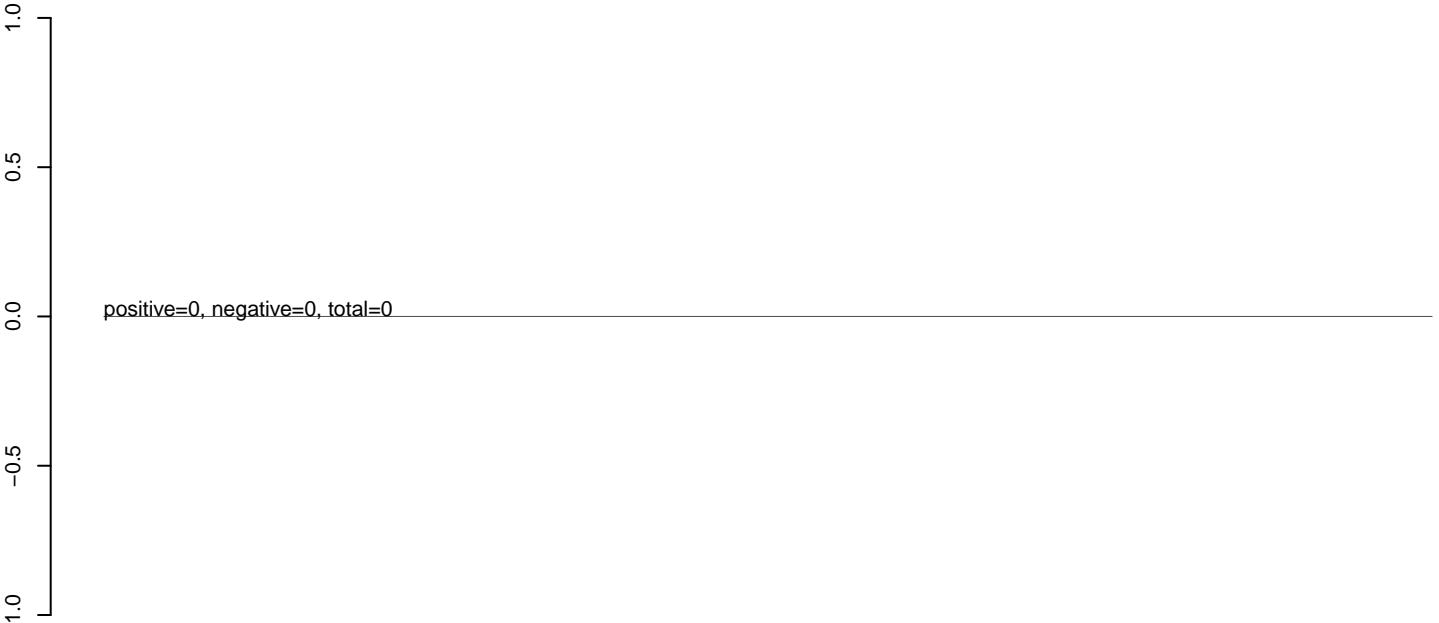
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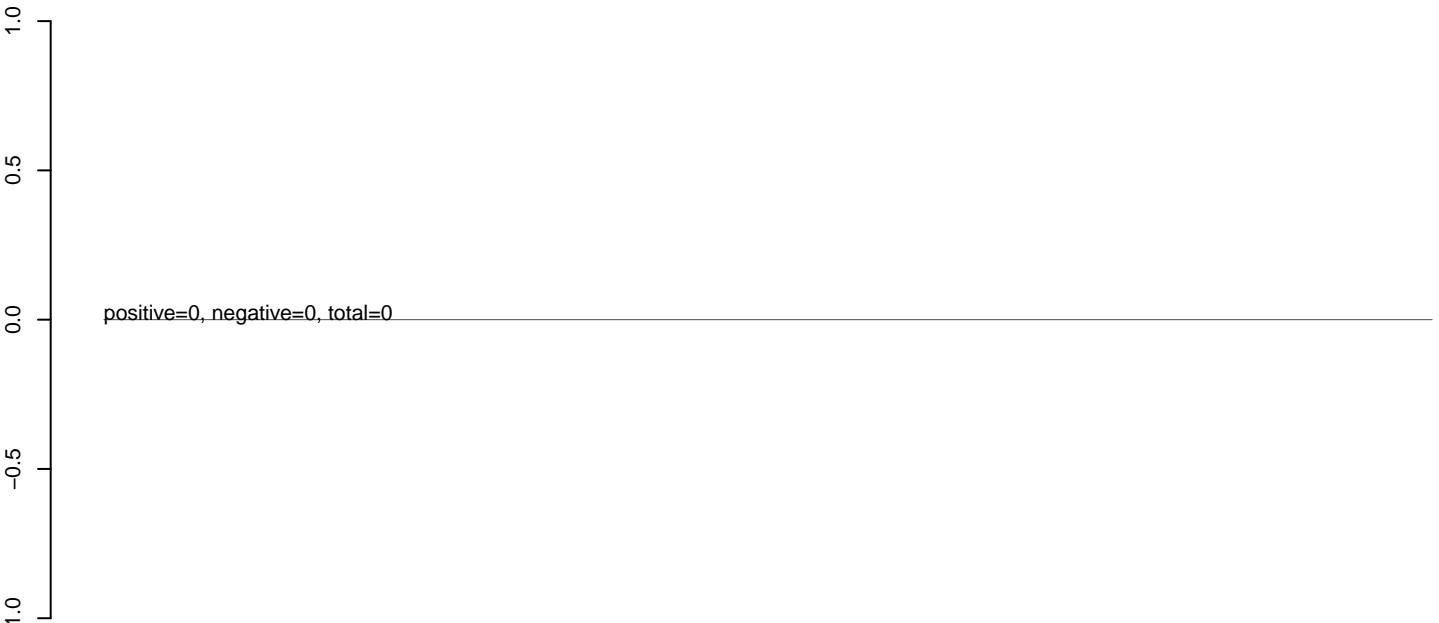
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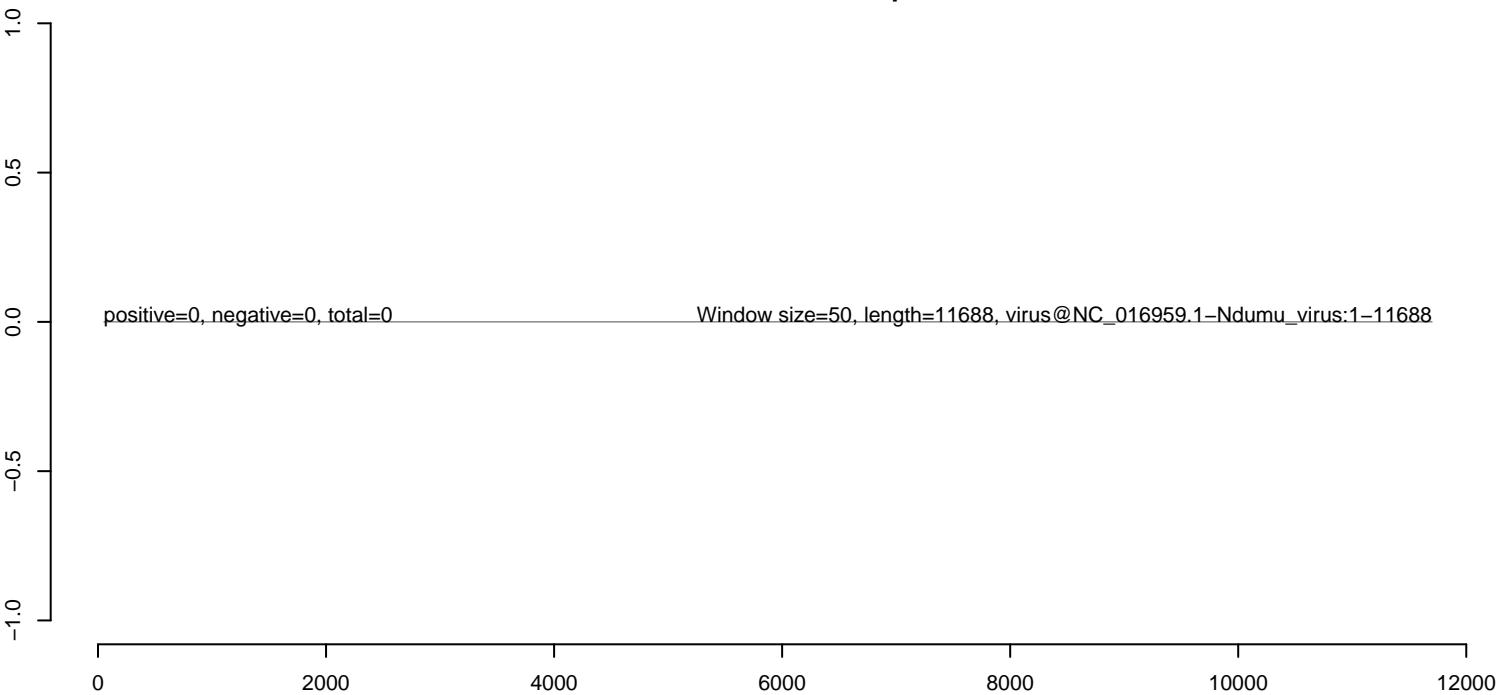
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### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep





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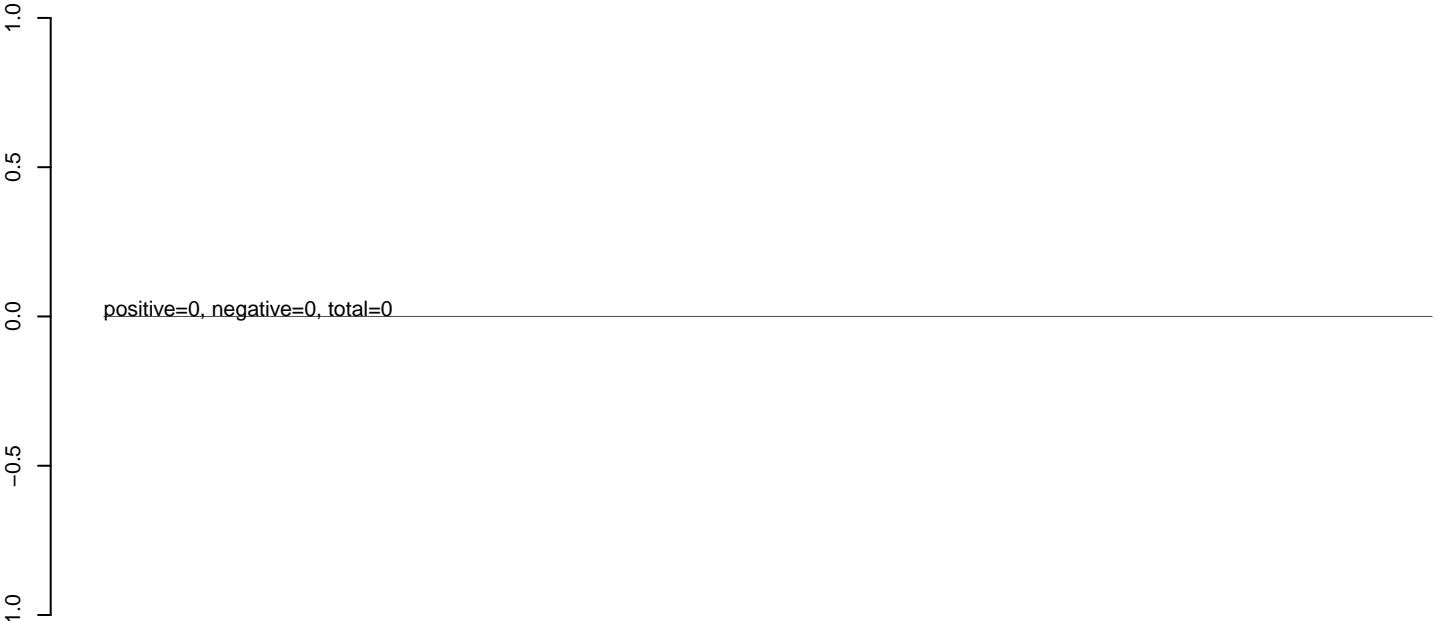


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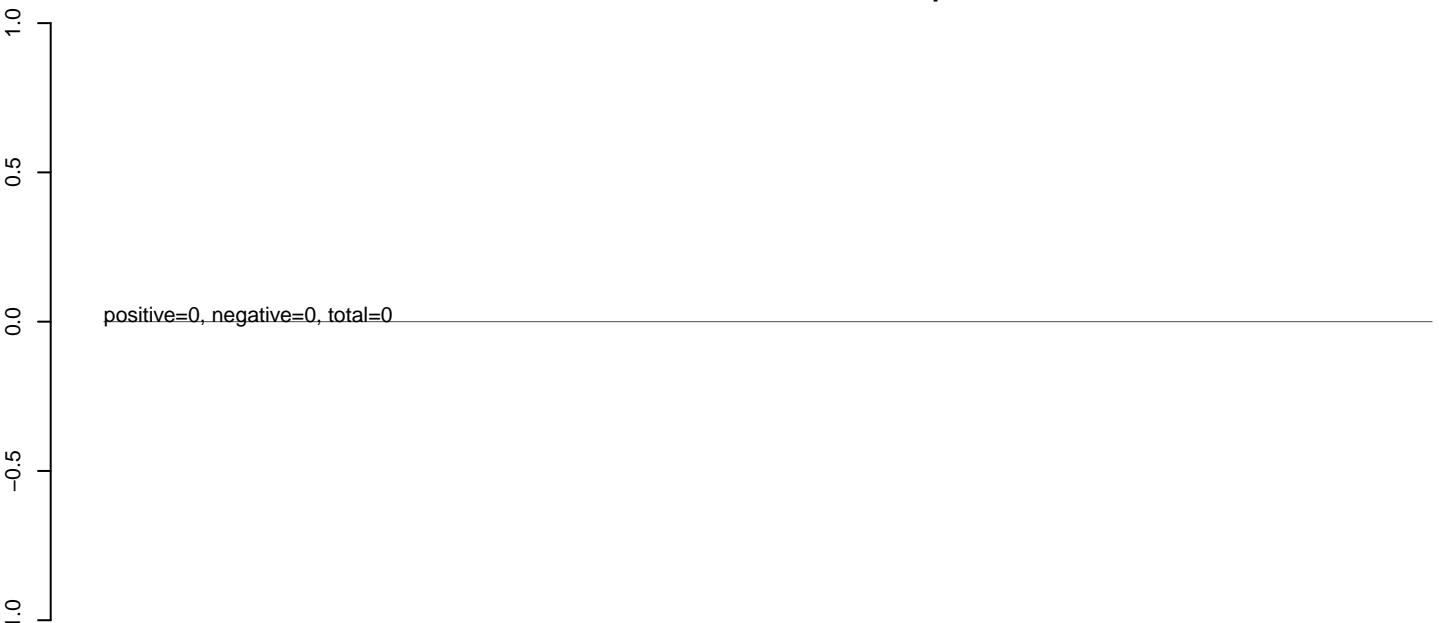


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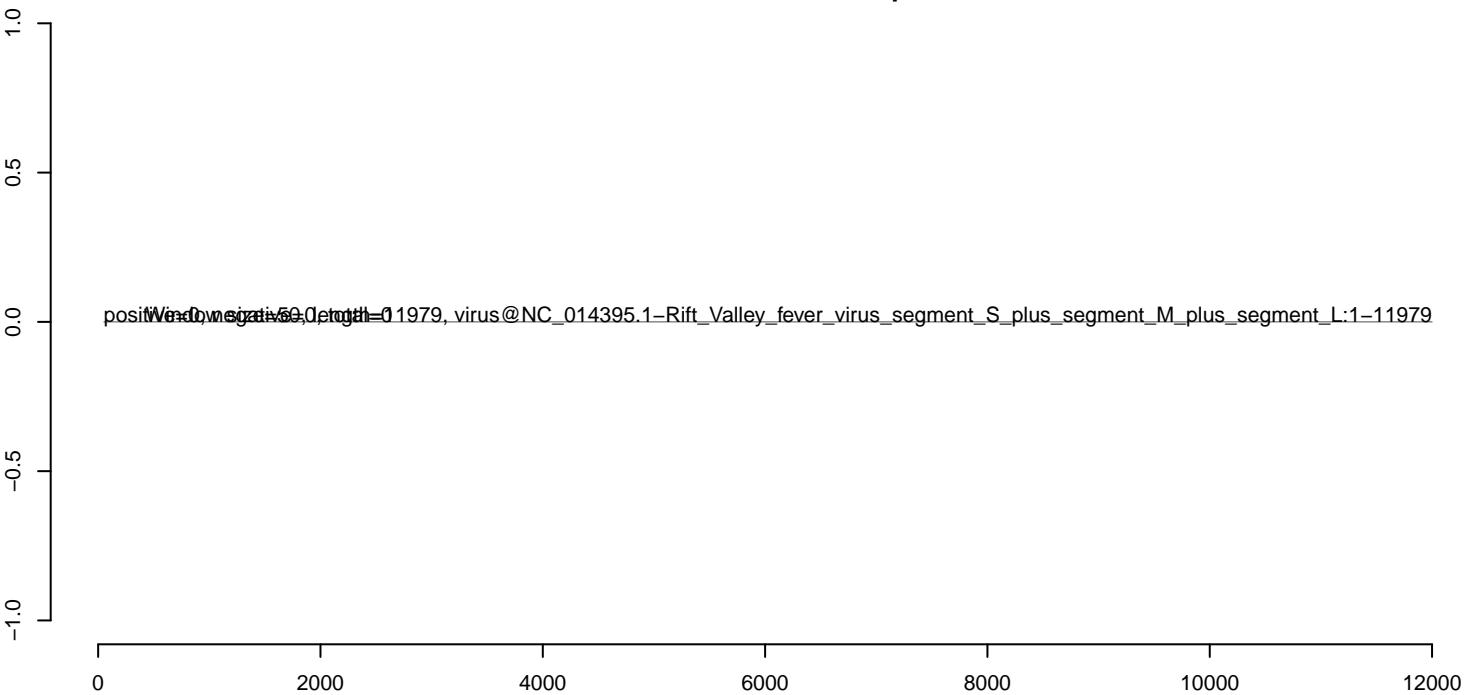
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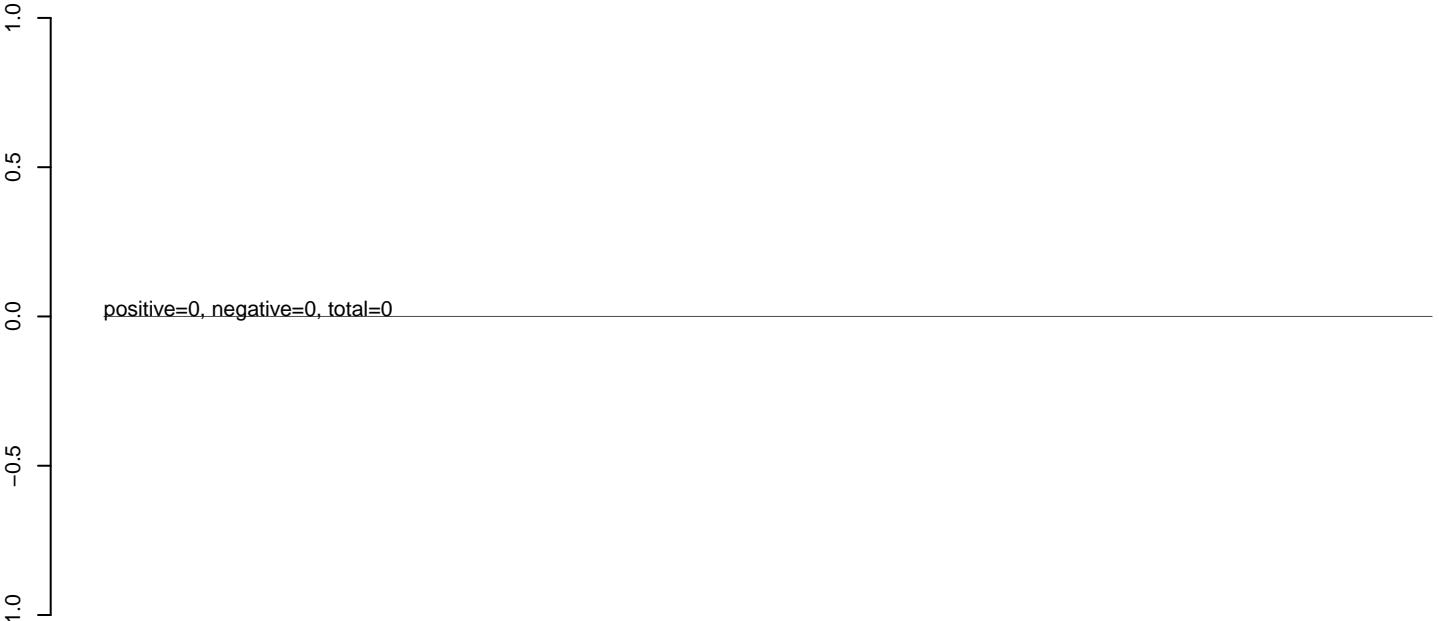
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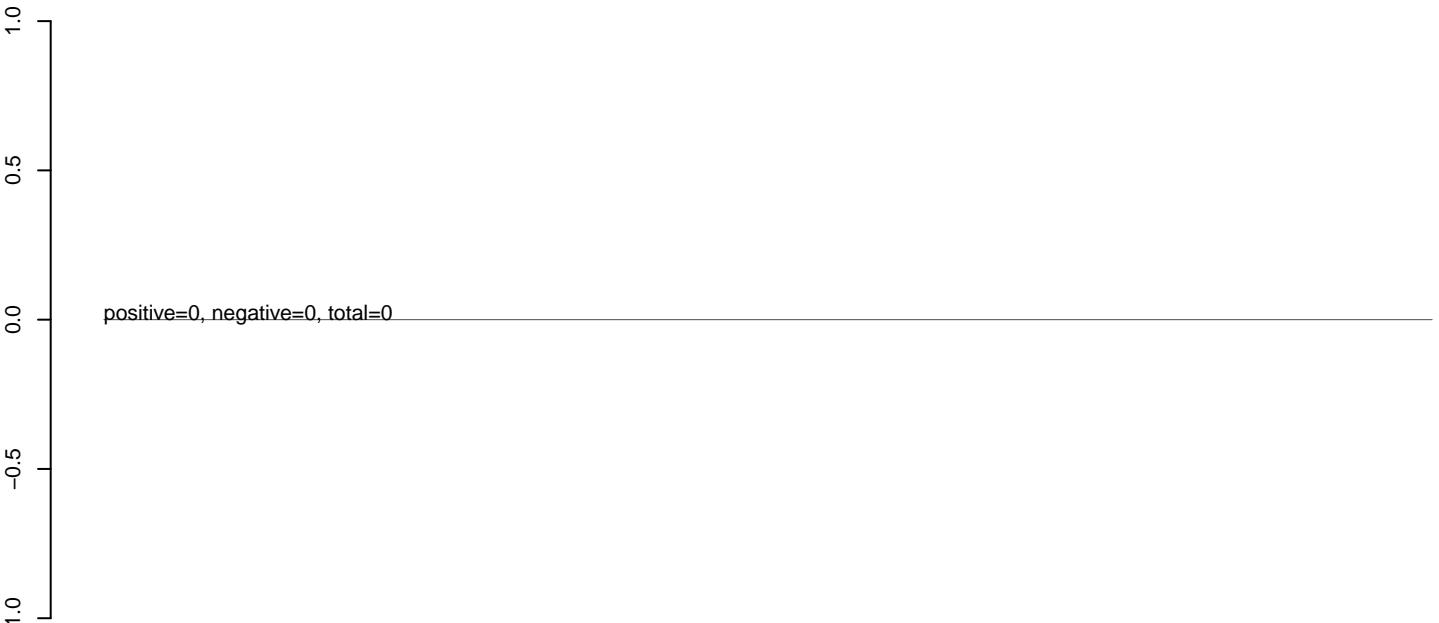
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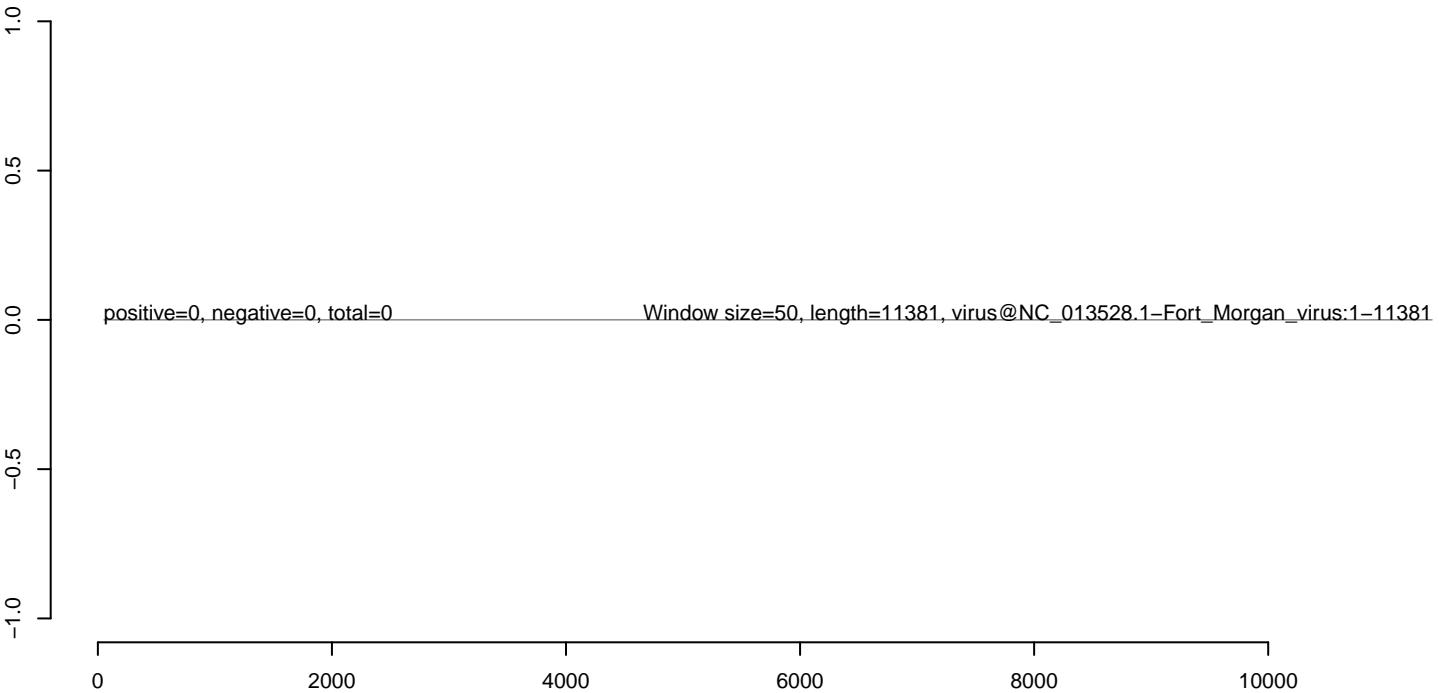
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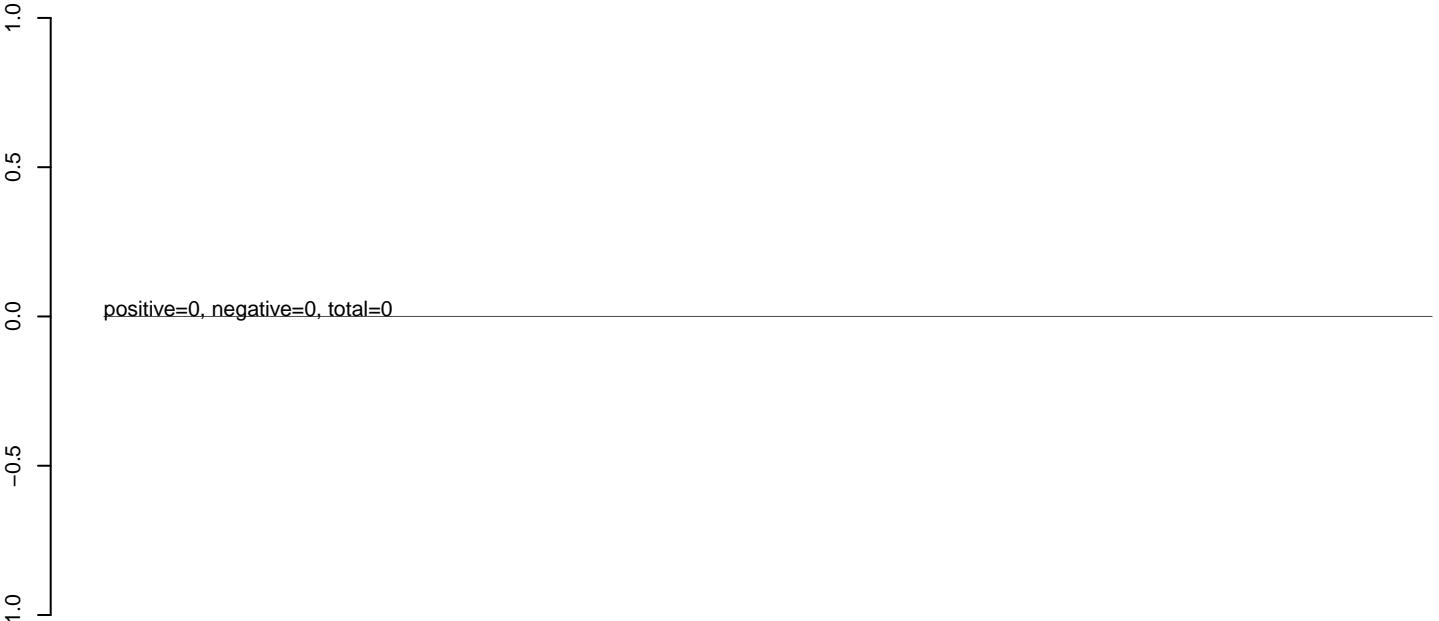
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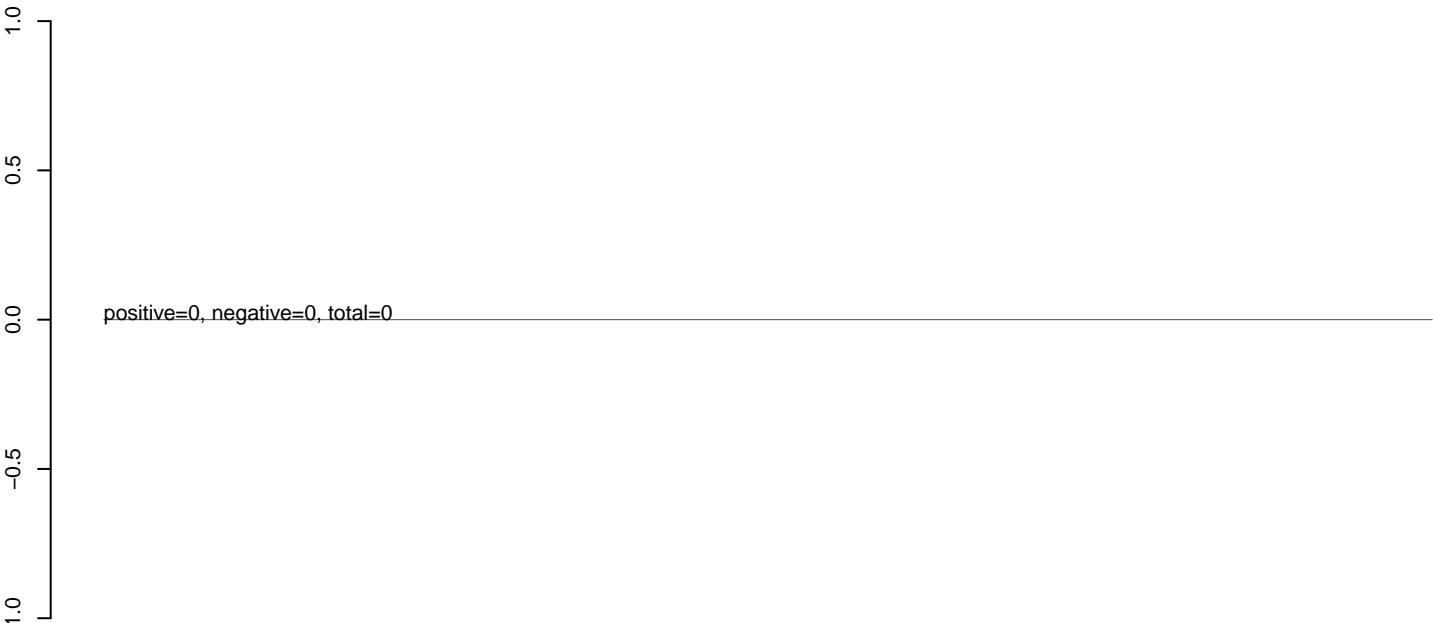
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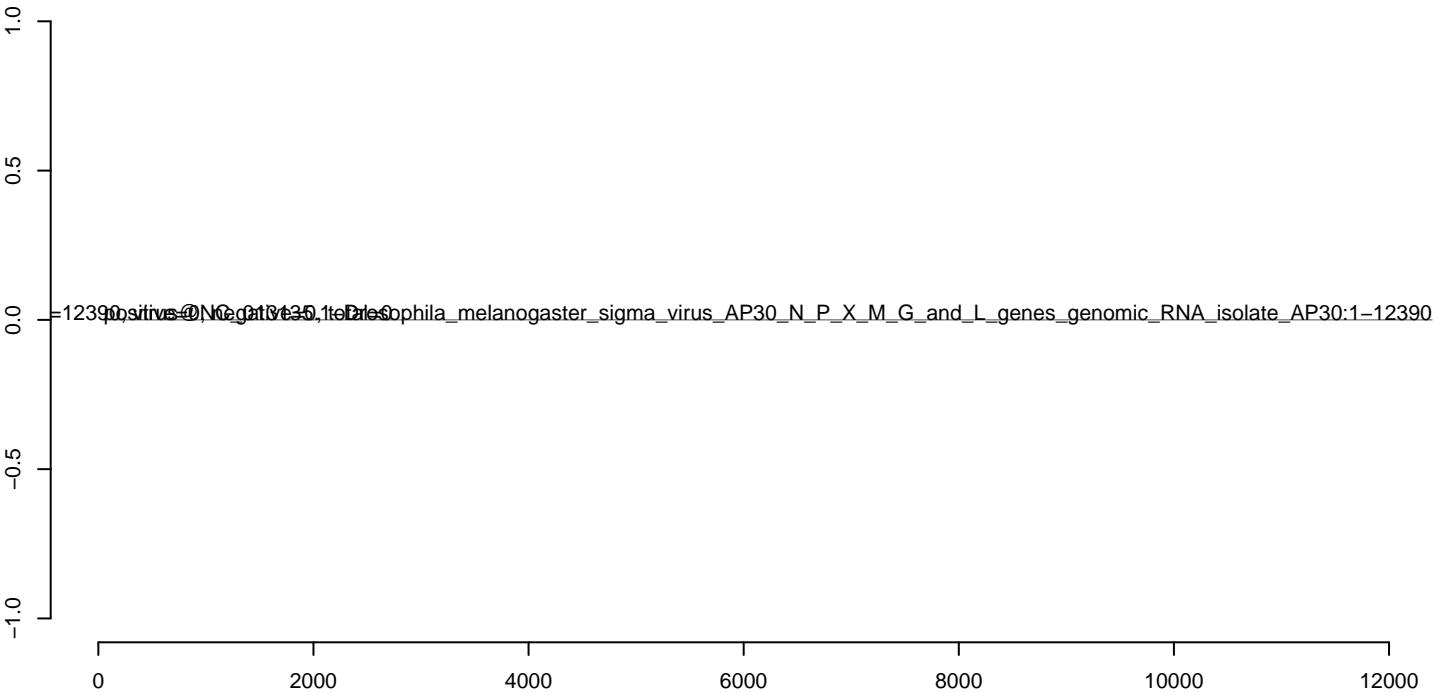
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### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

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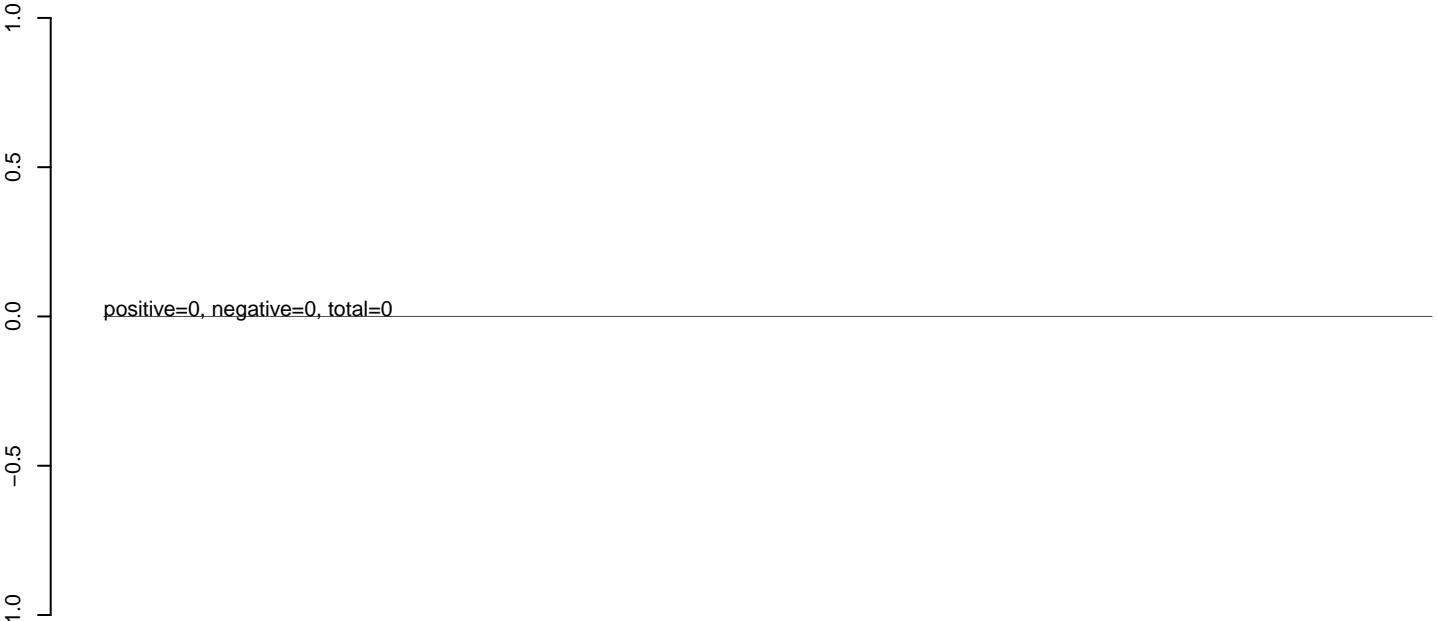
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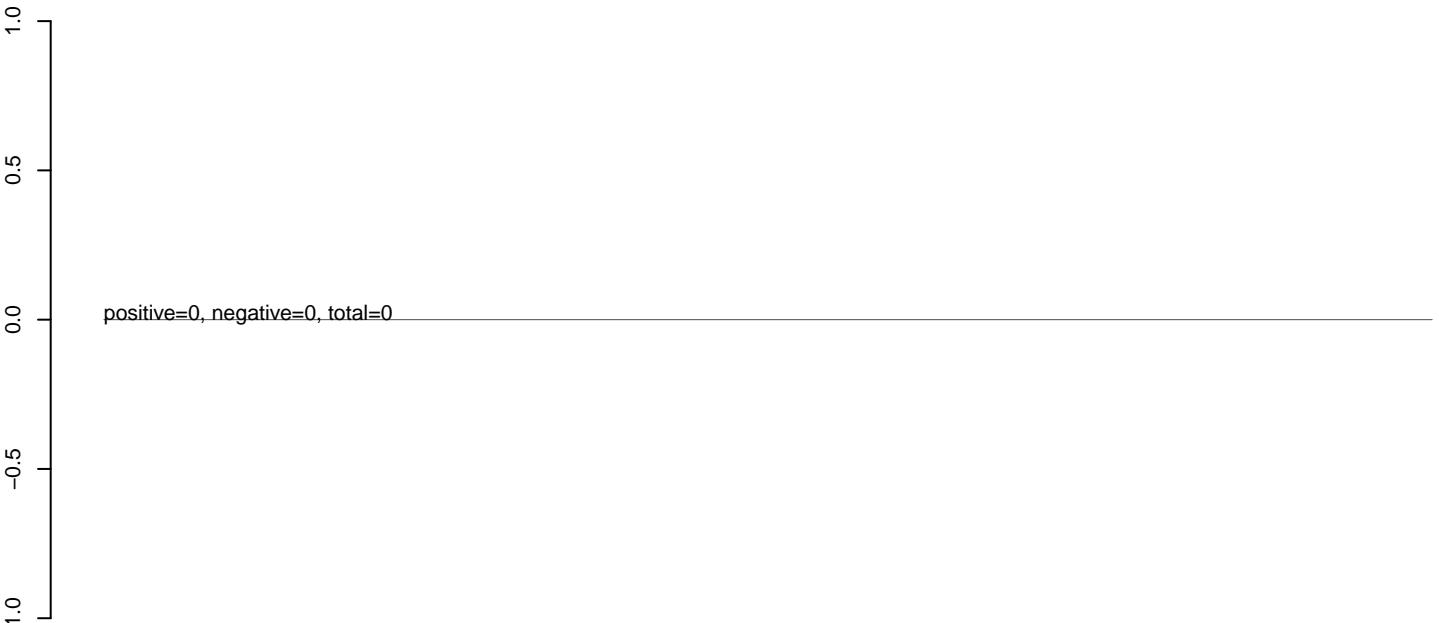
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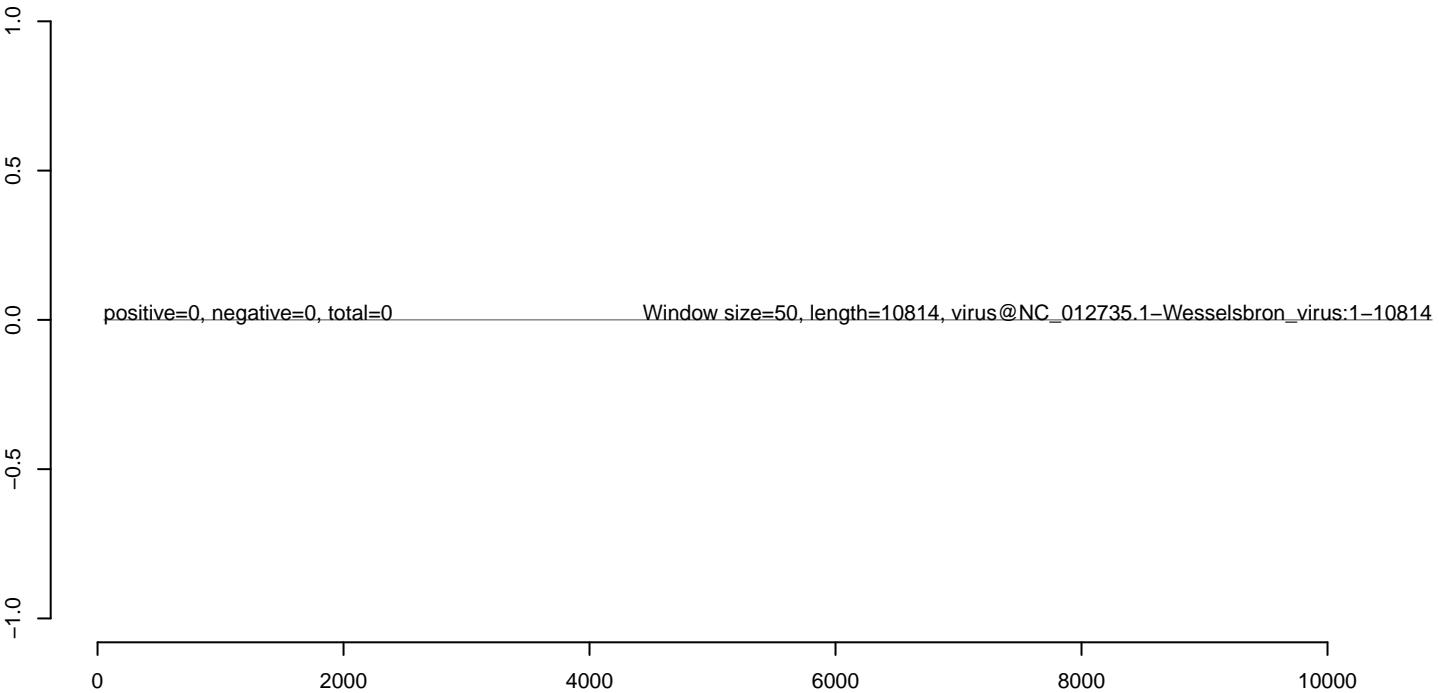
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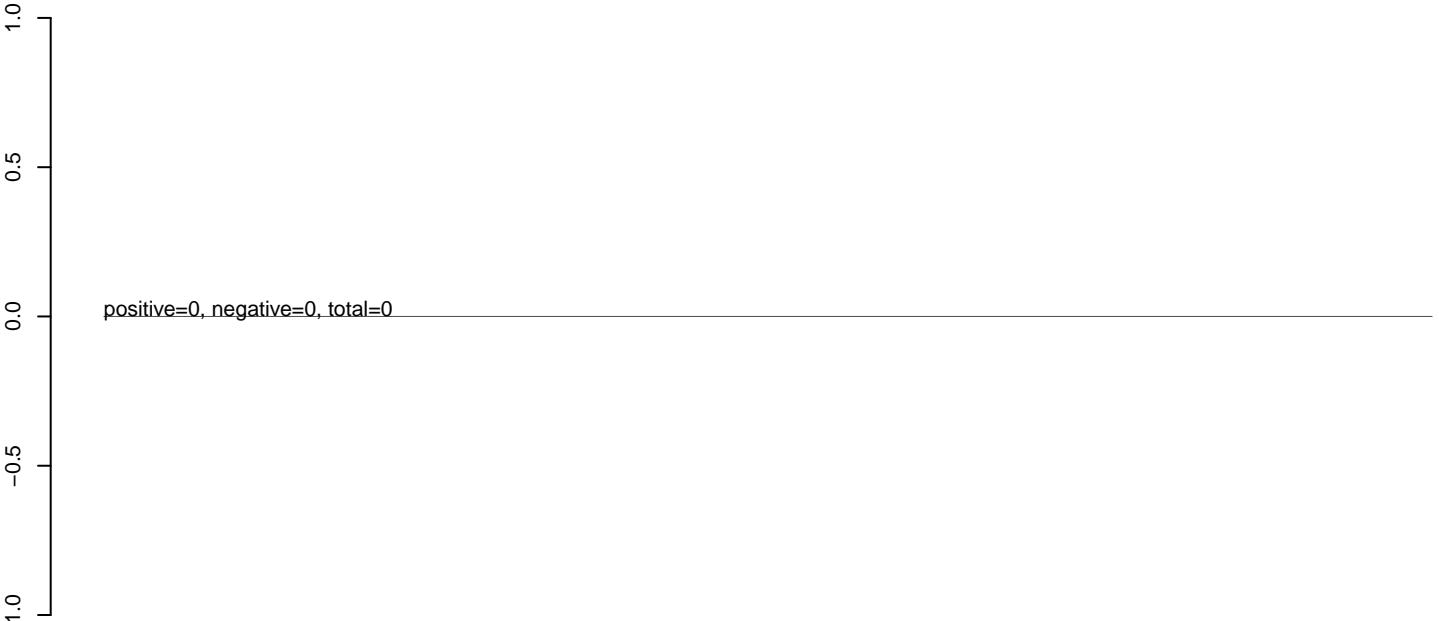
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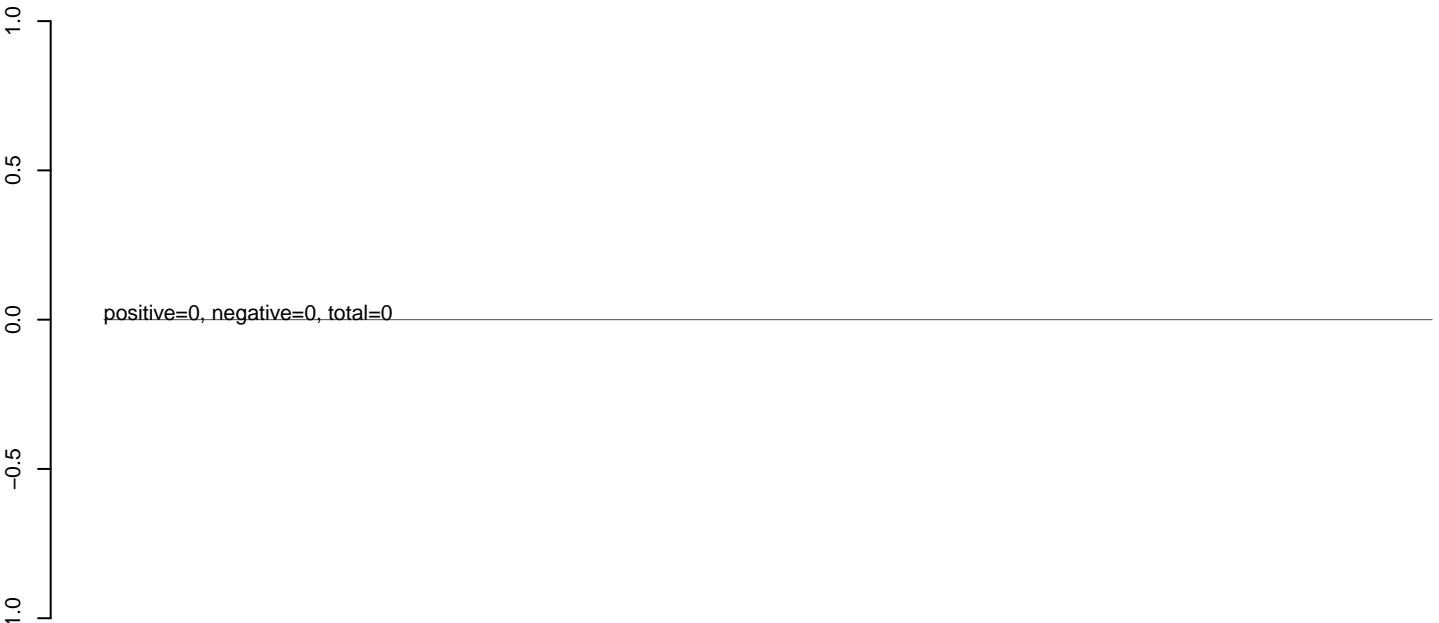
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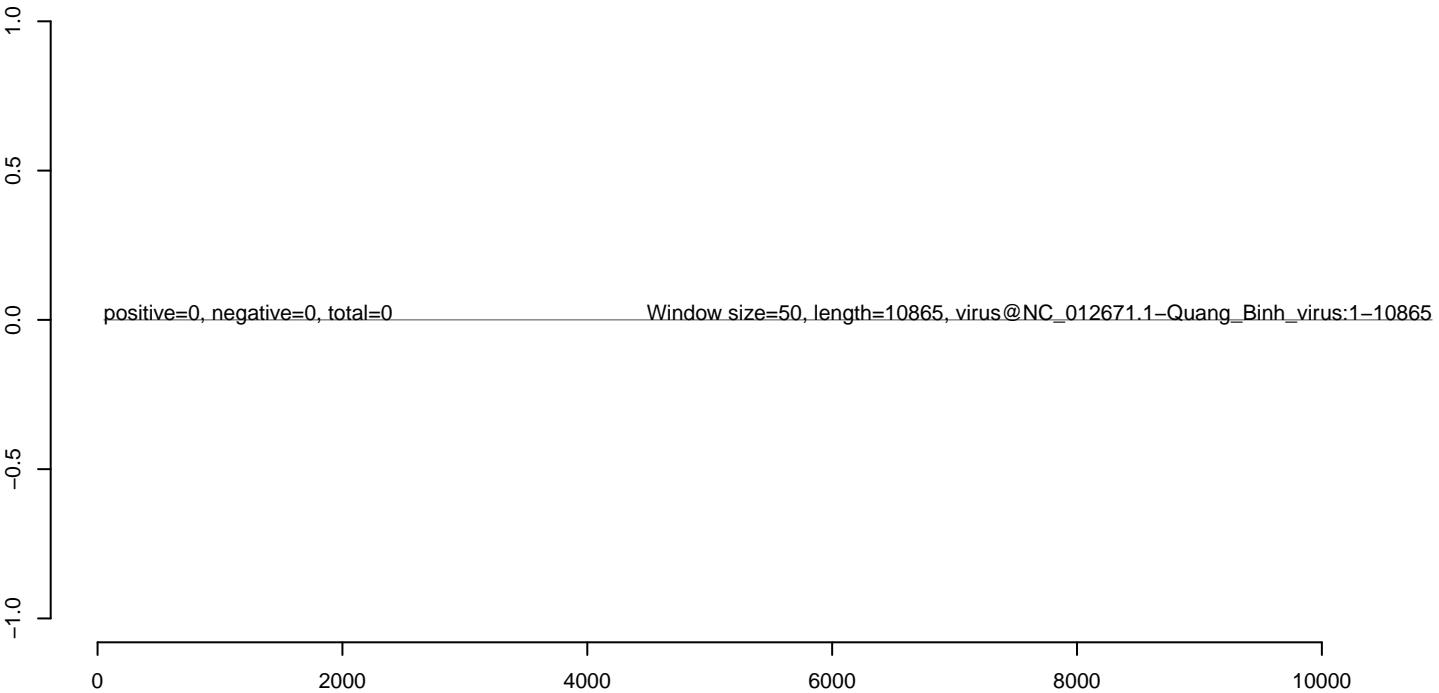
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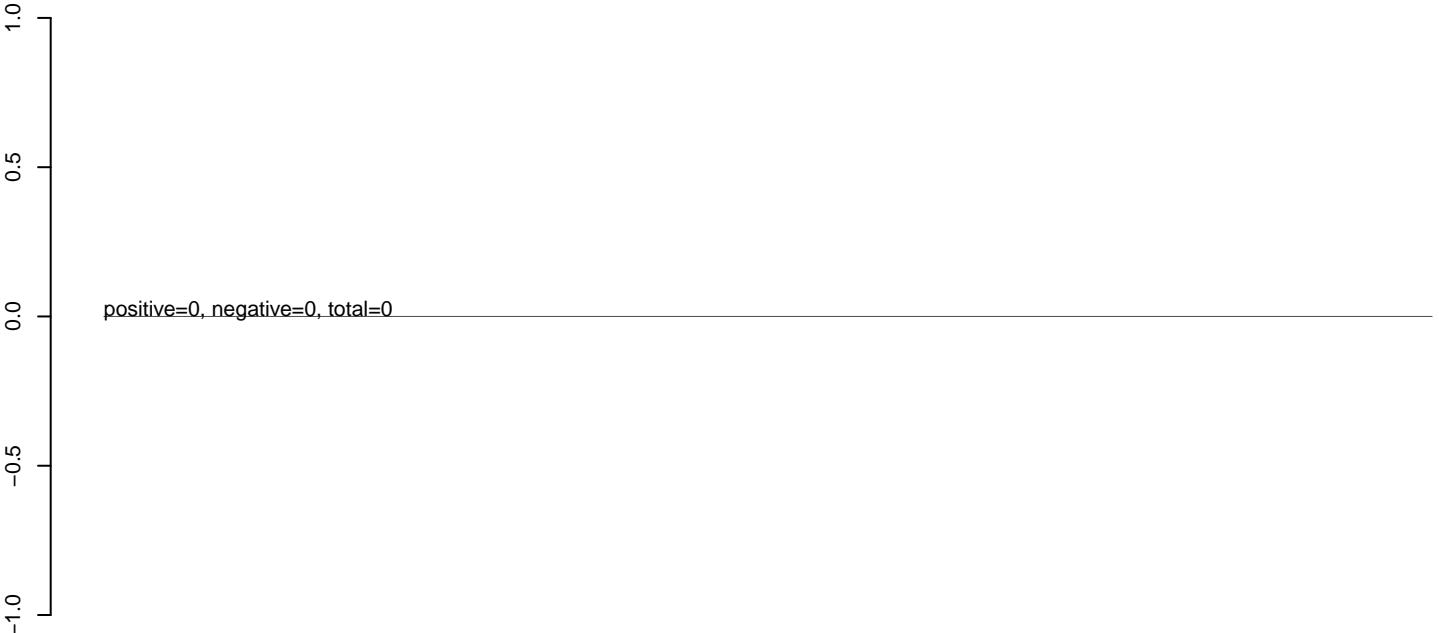
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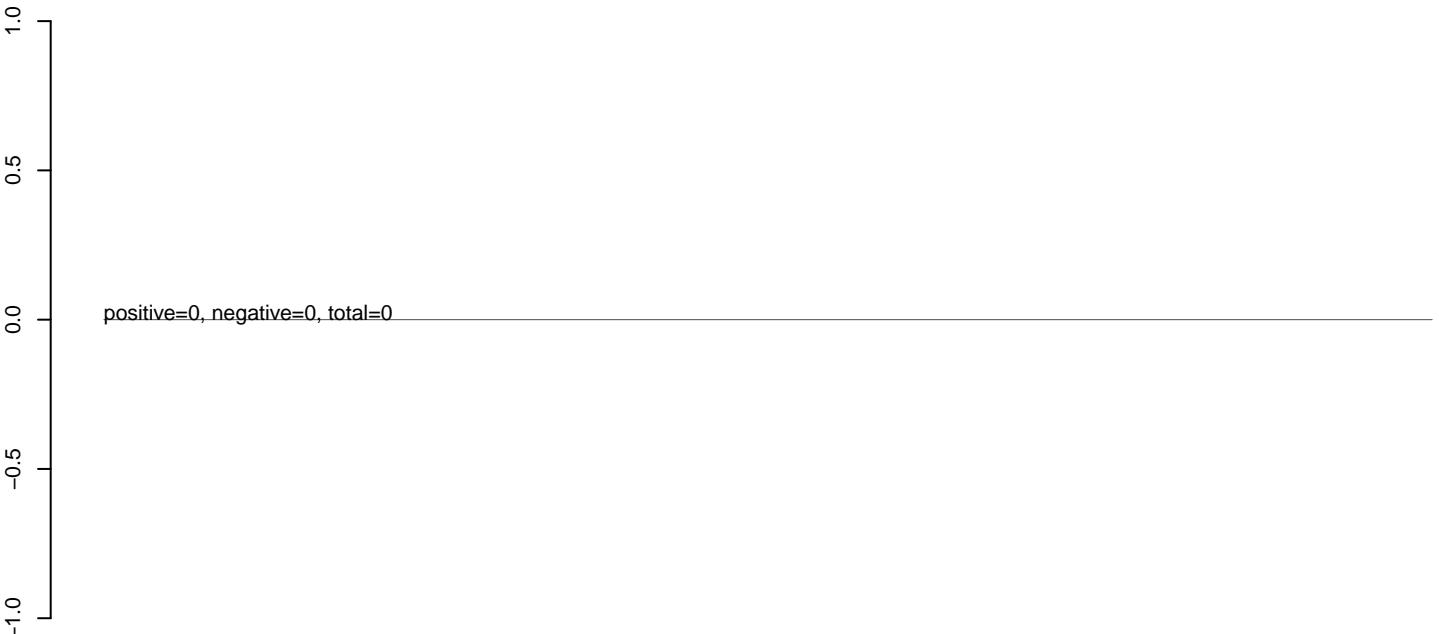
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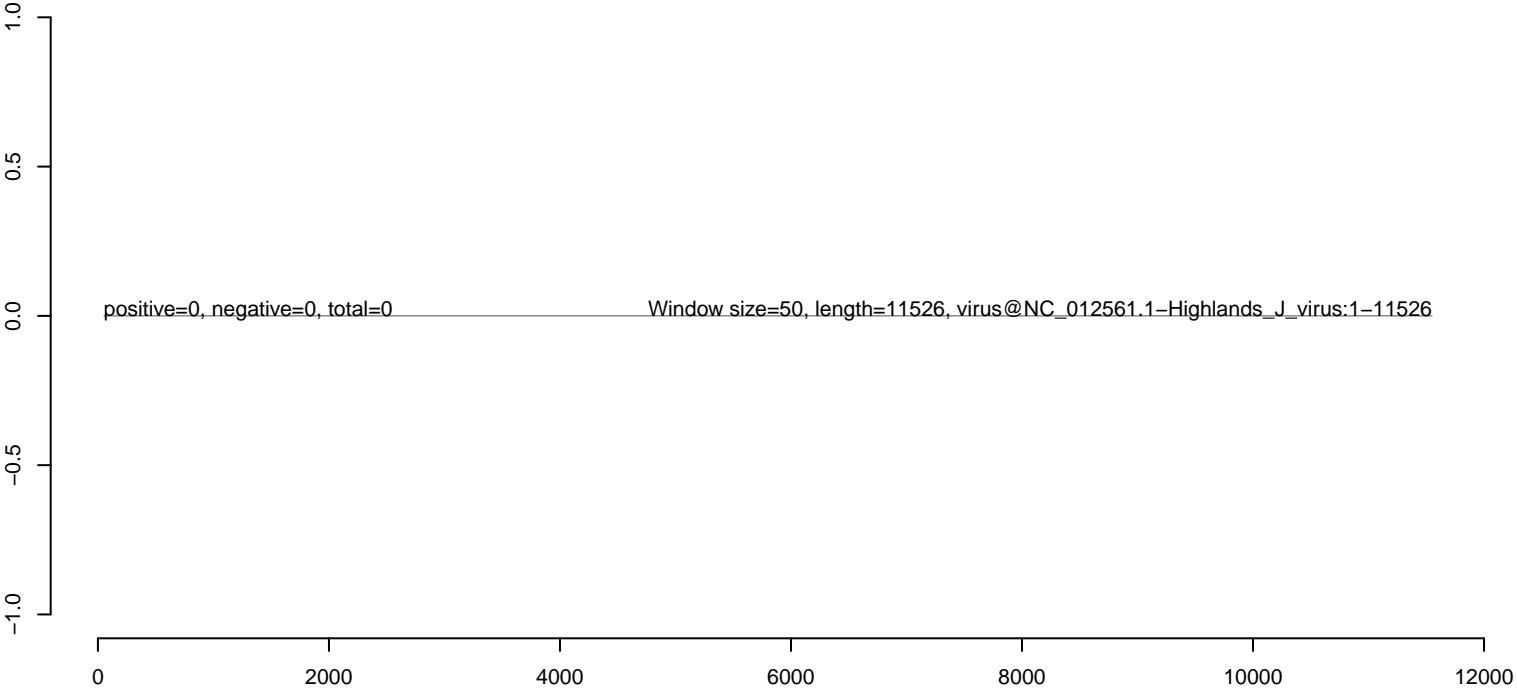
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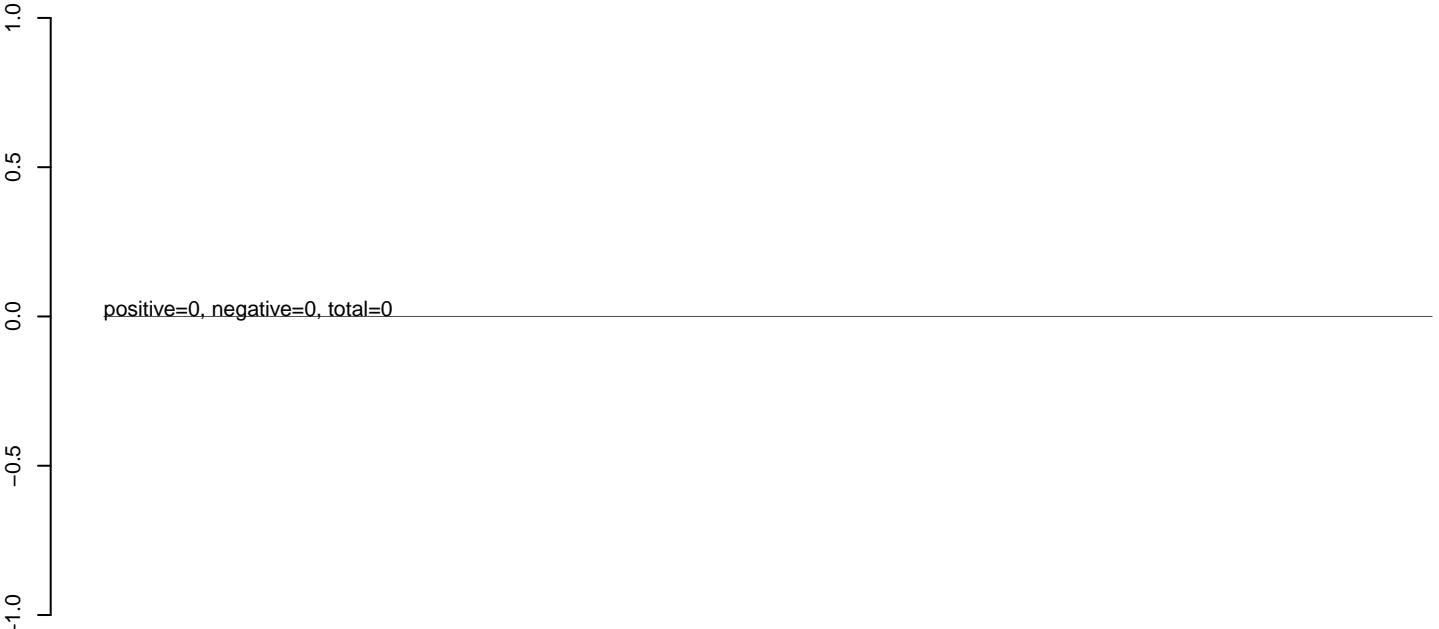
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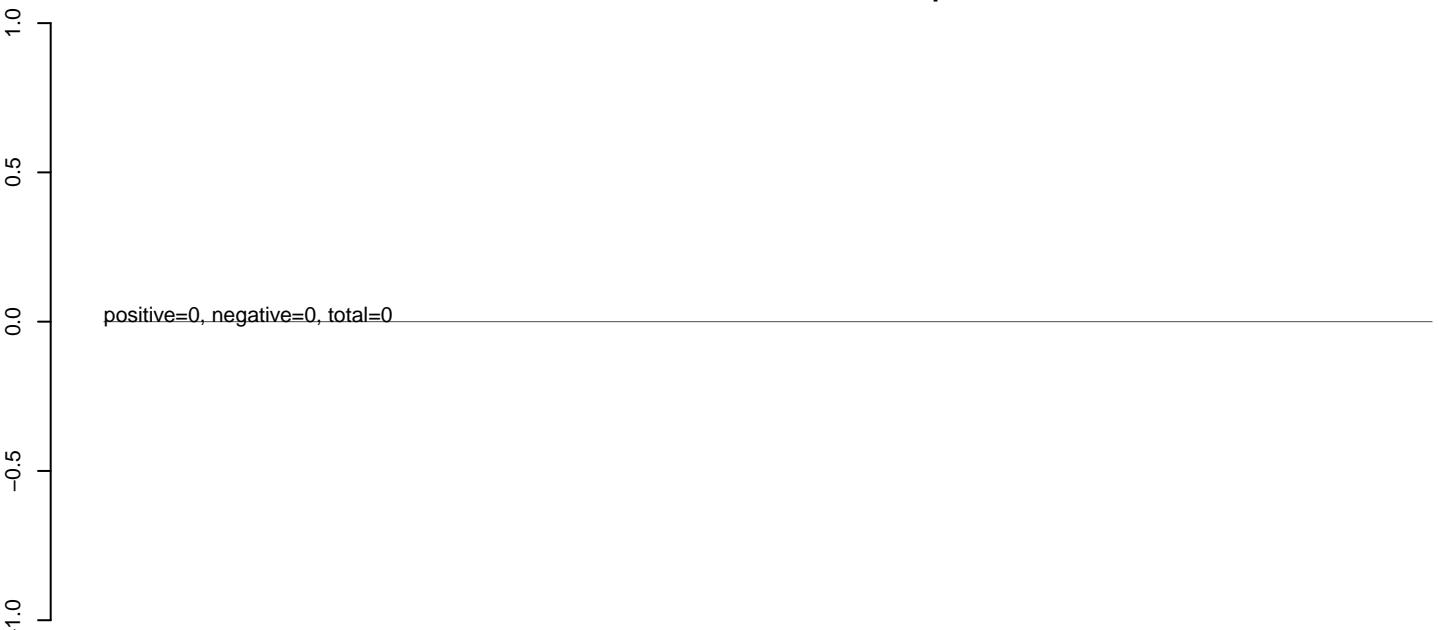
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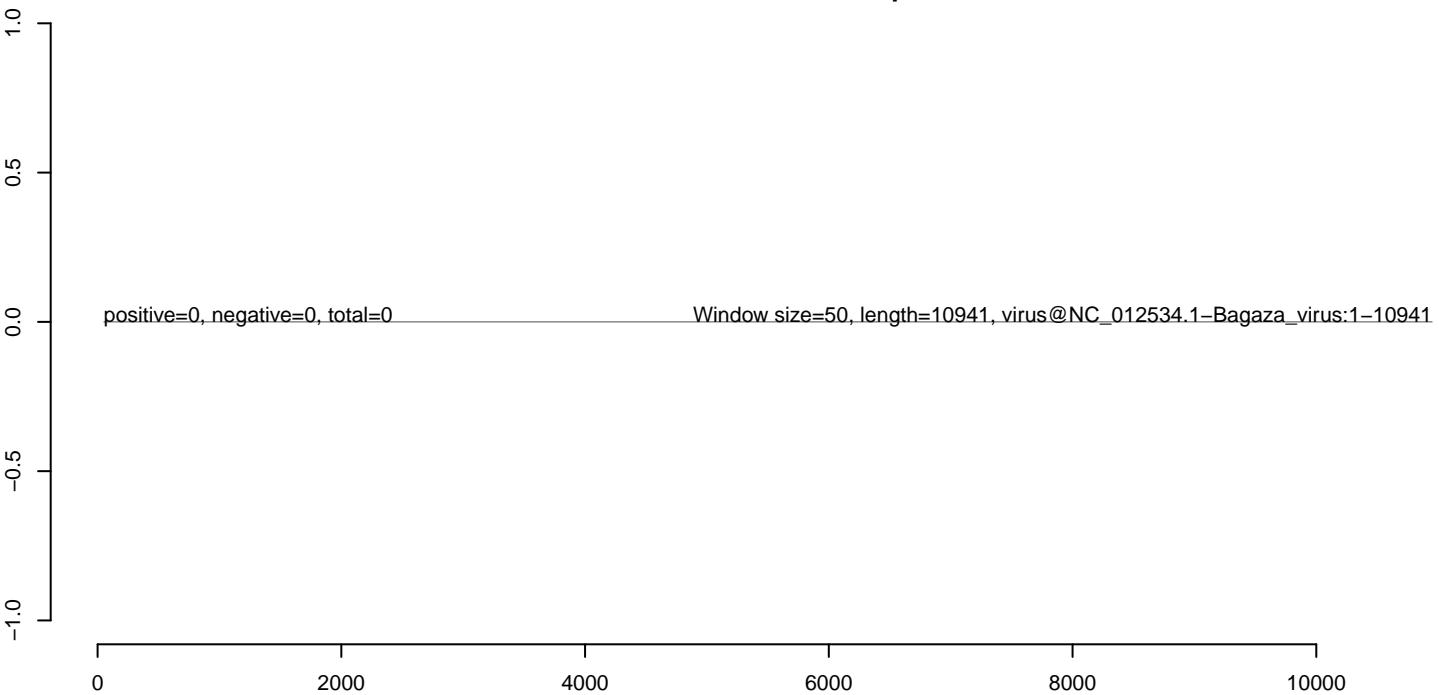
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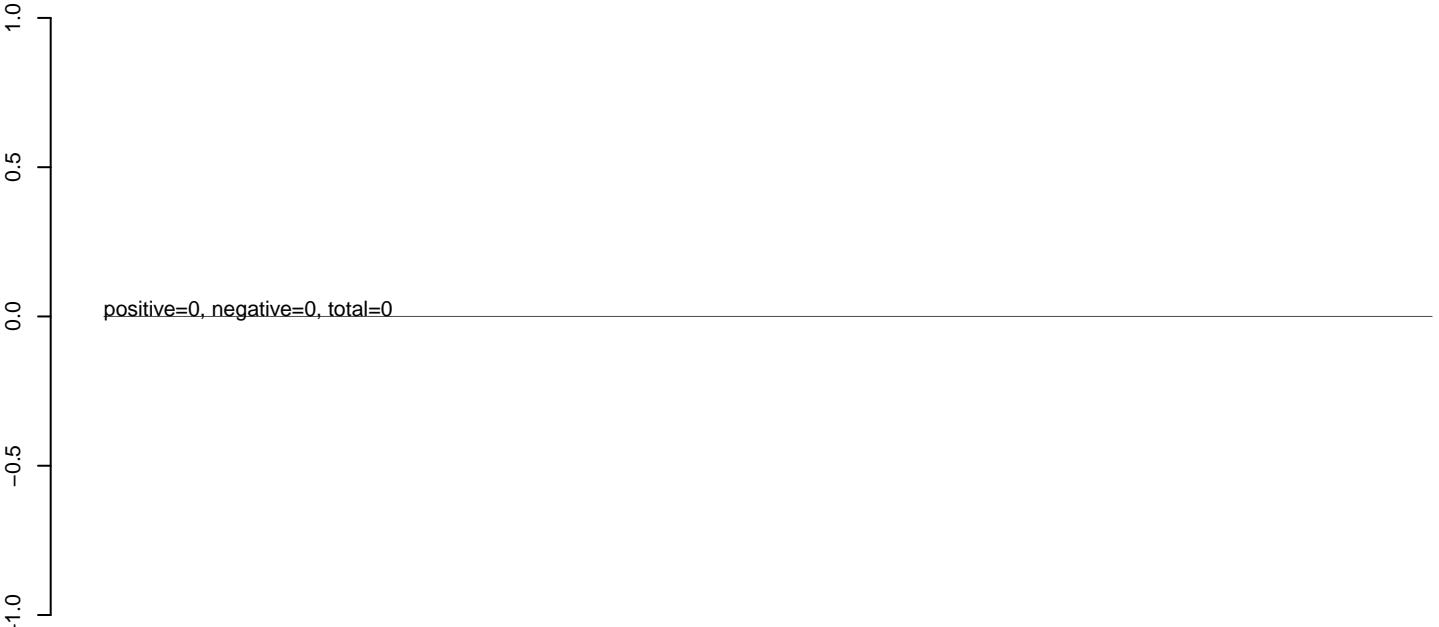
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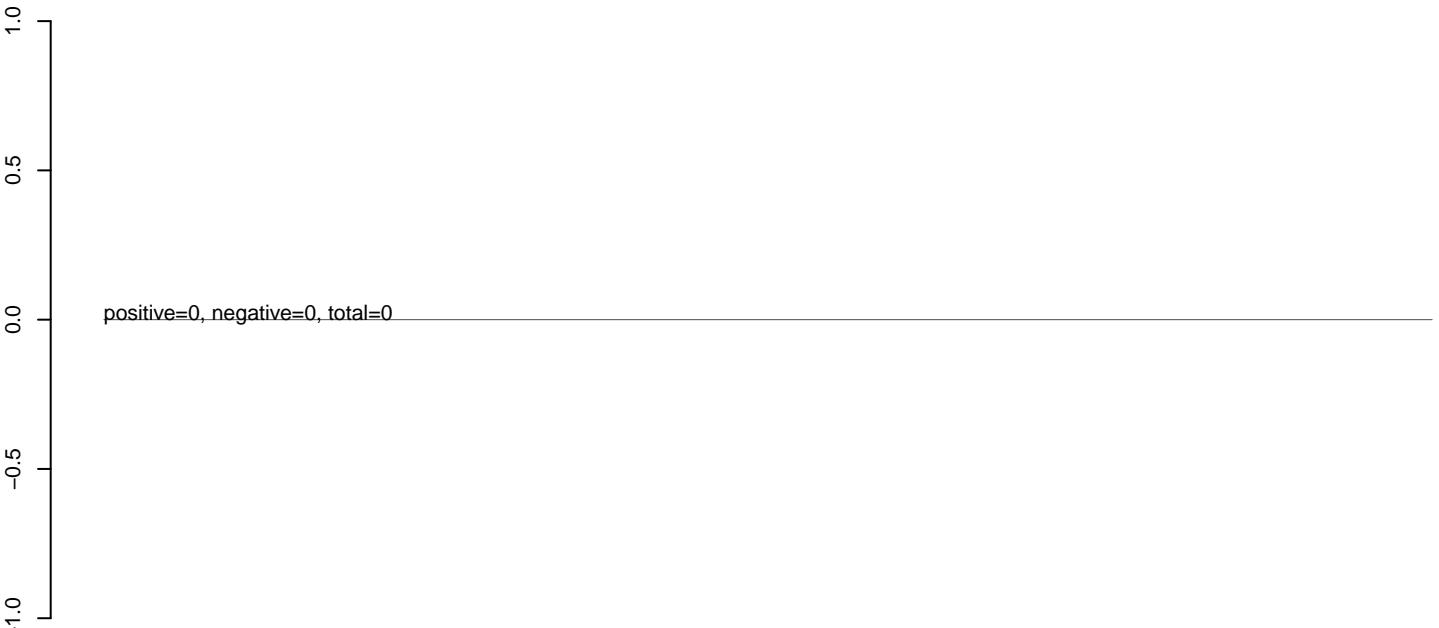
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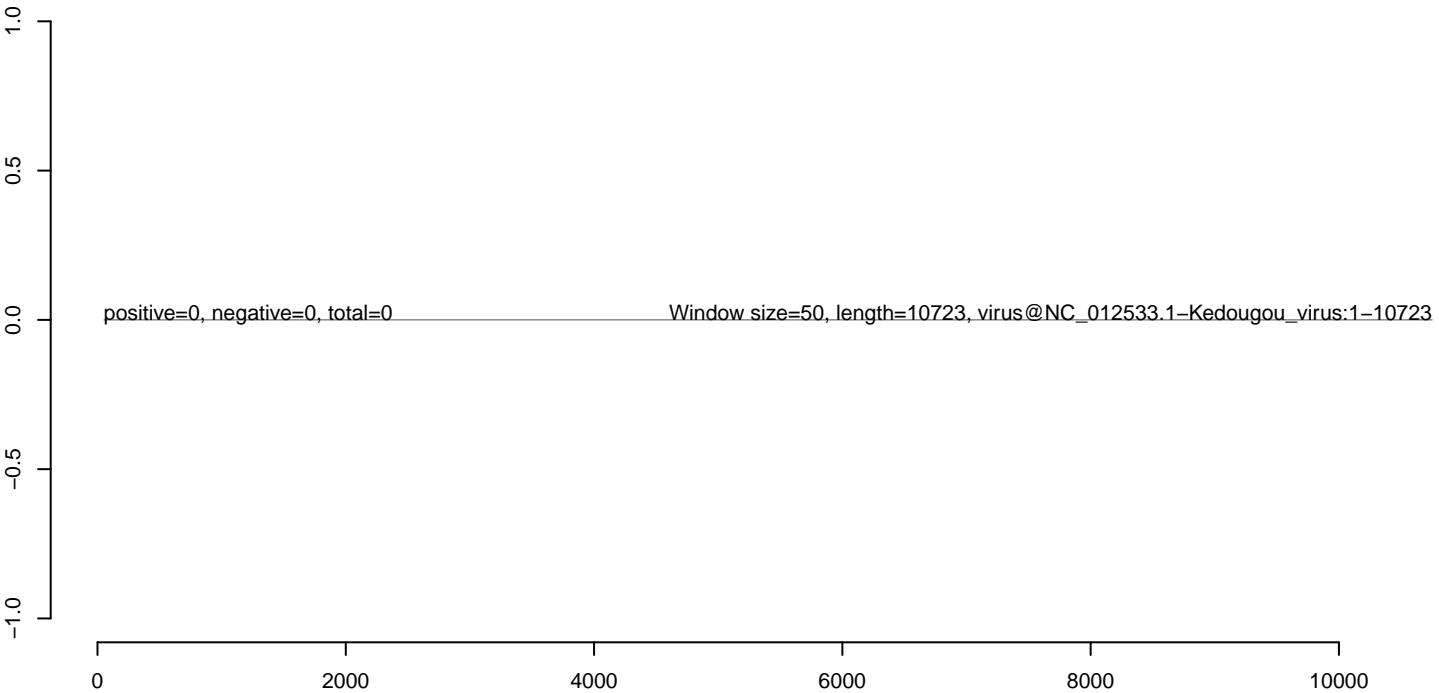
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### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

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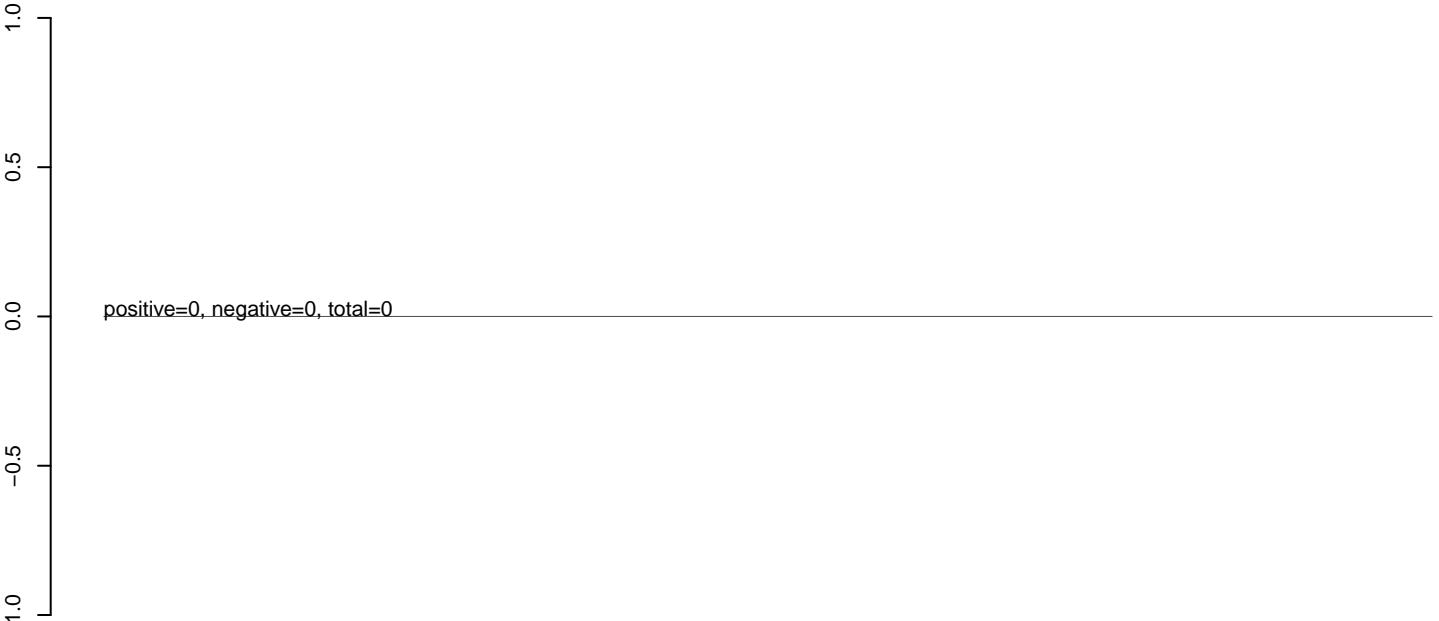
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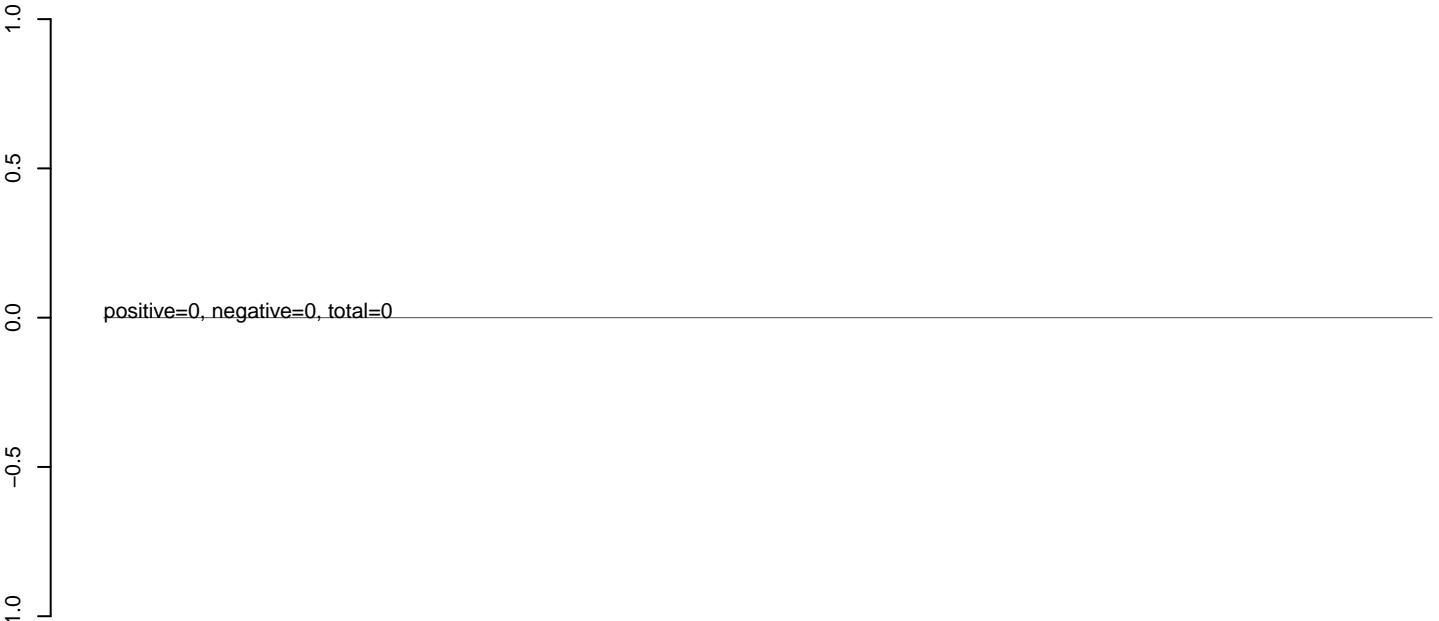
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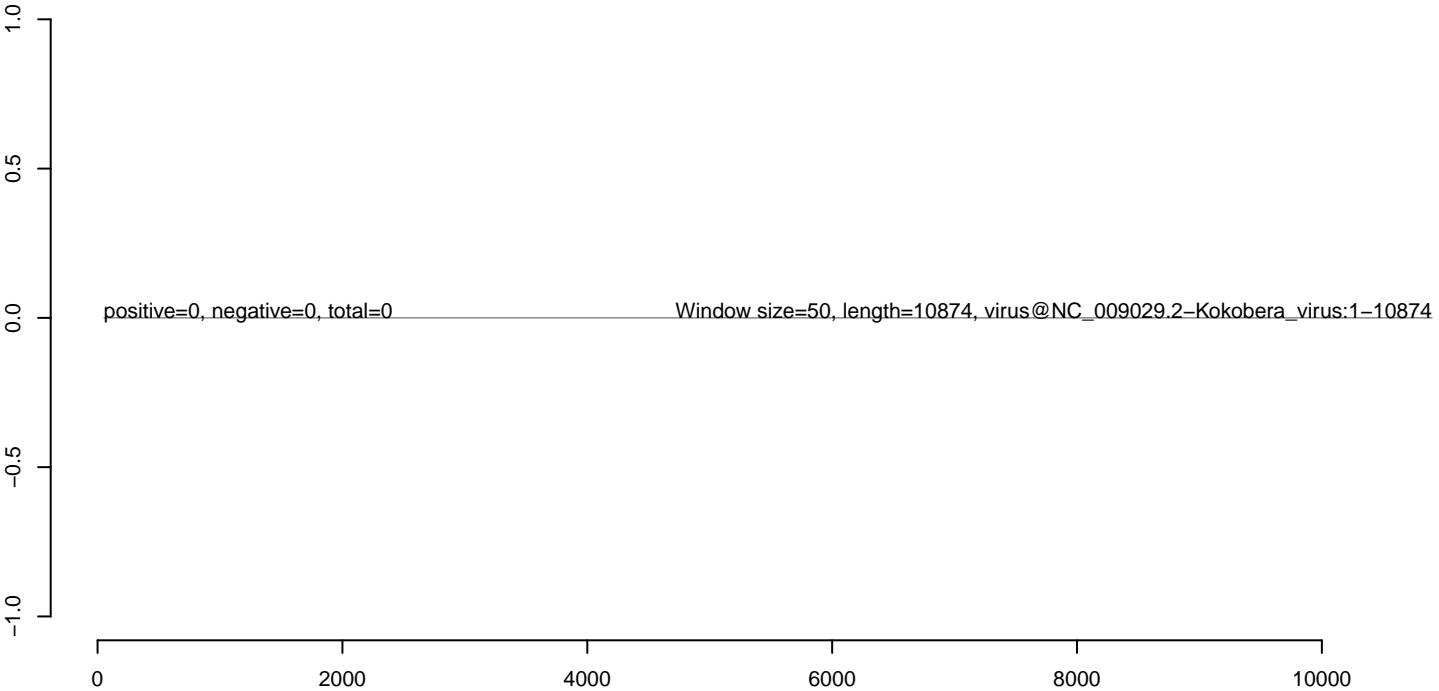
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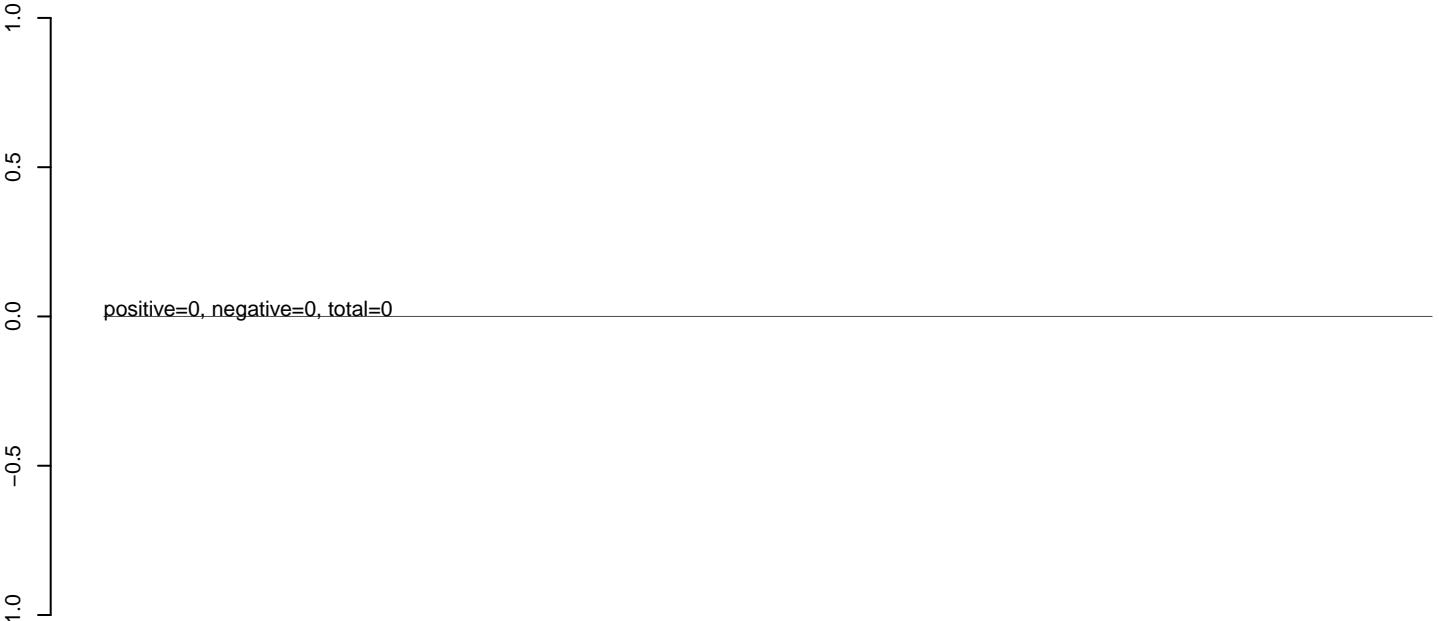
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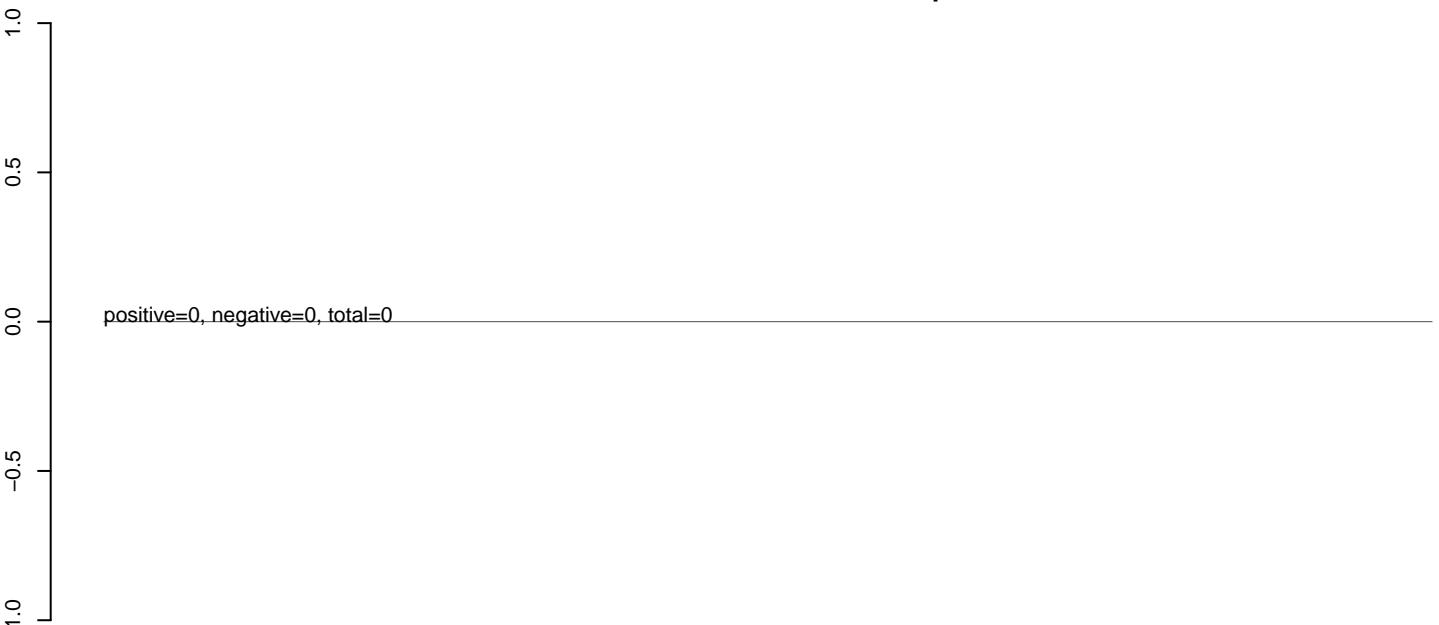
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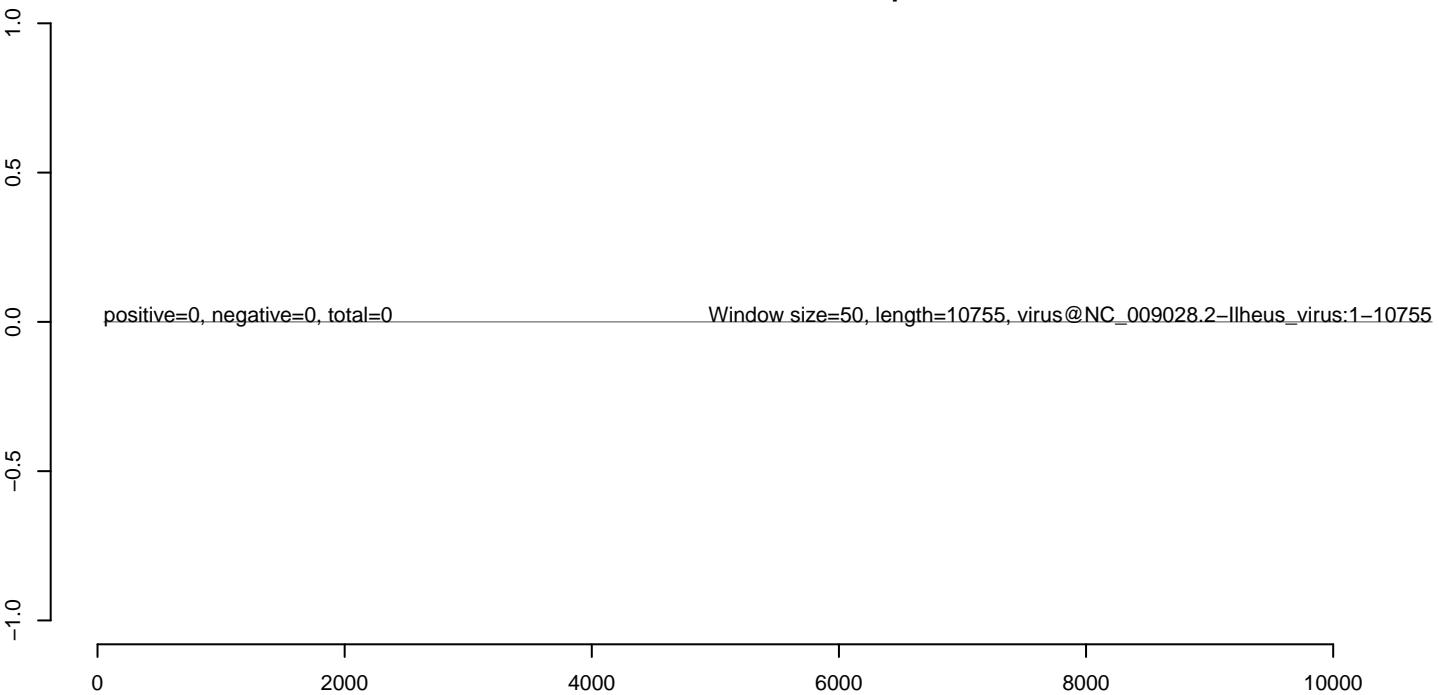
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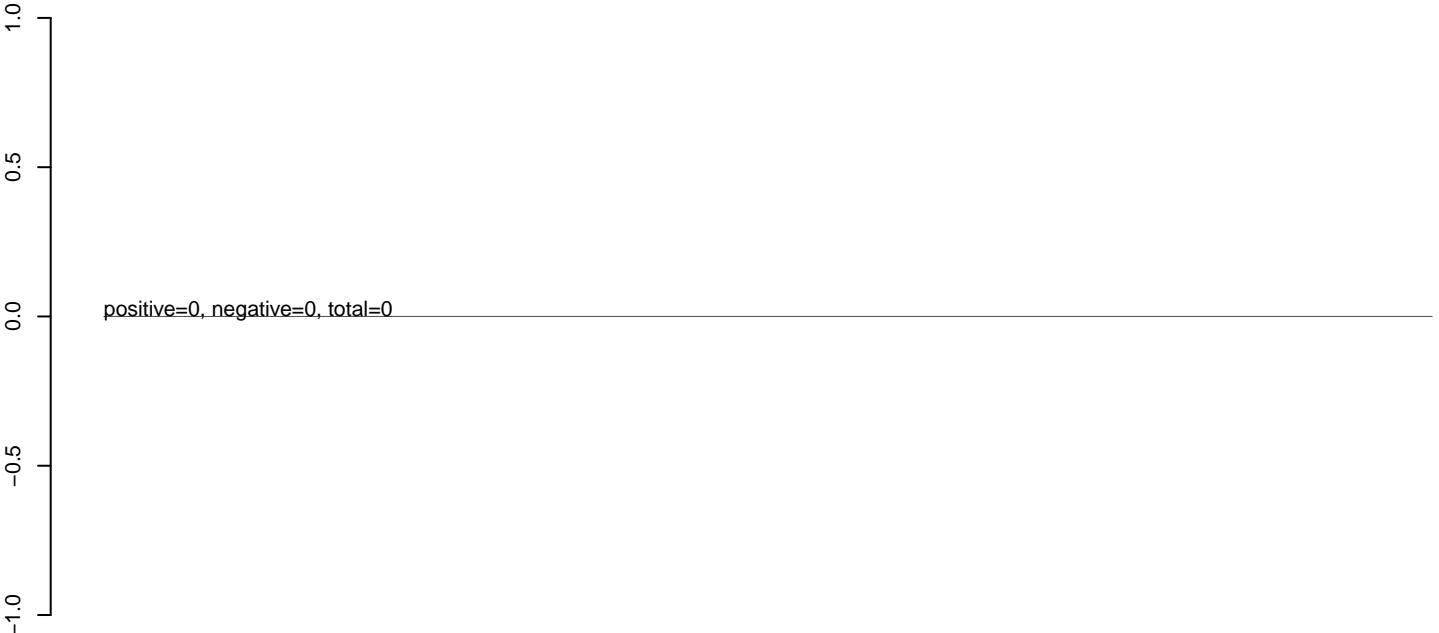
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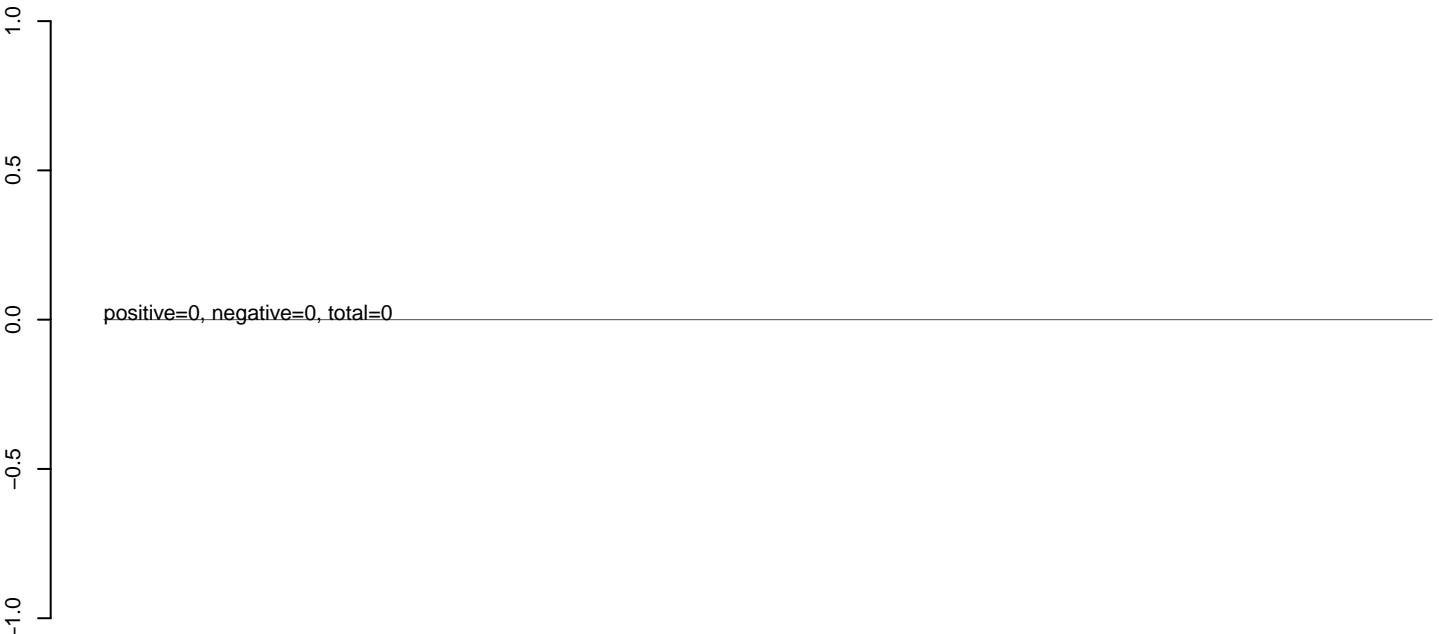
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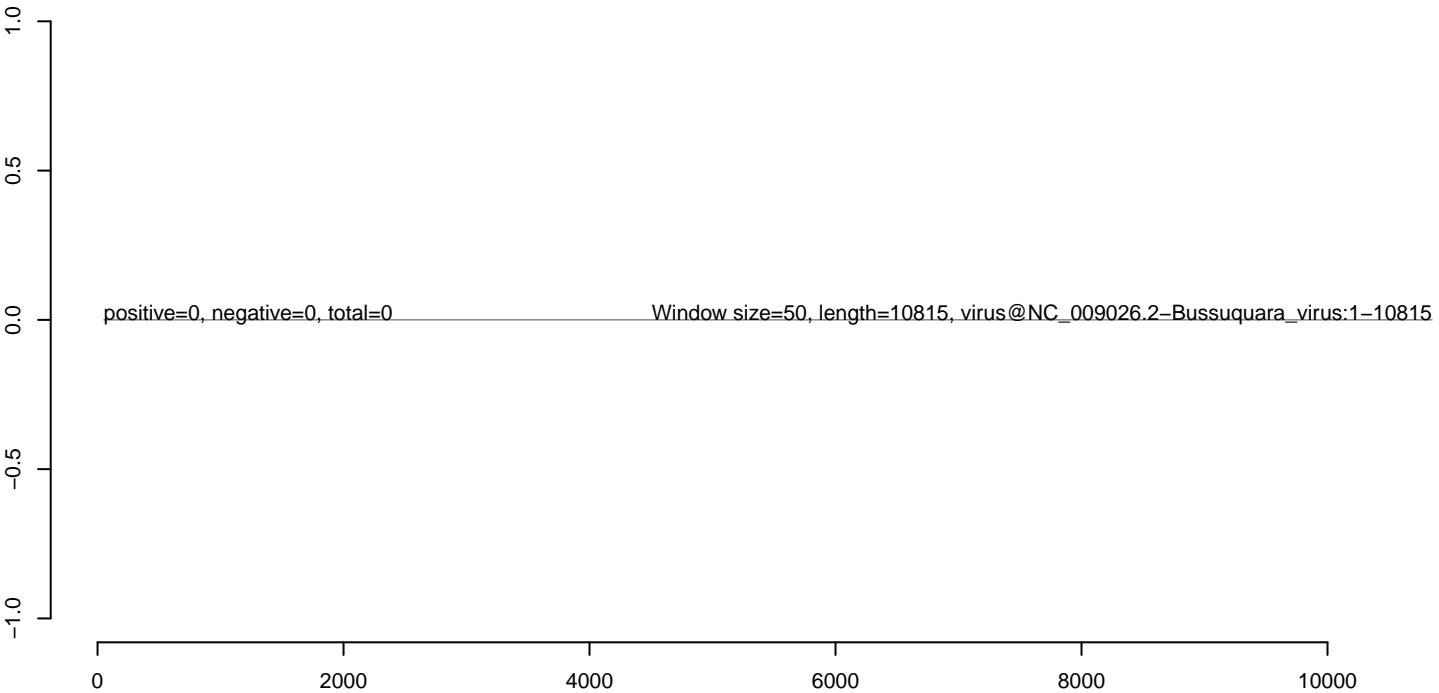
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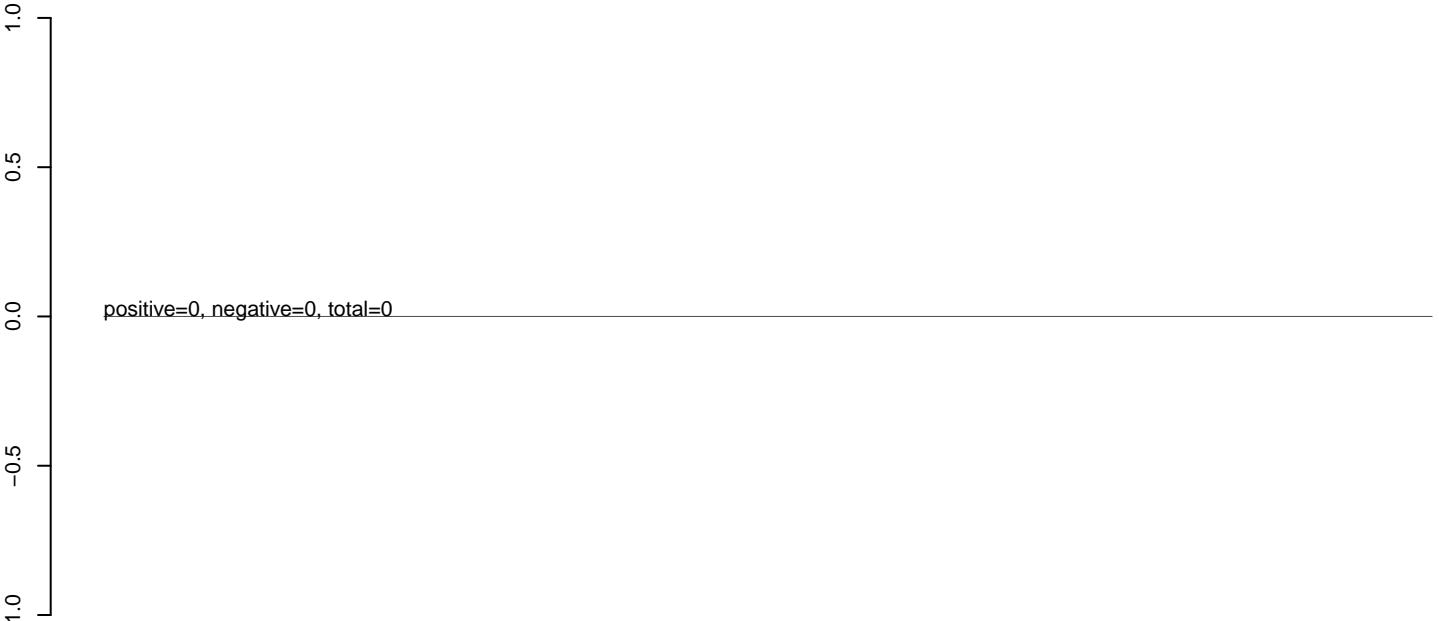
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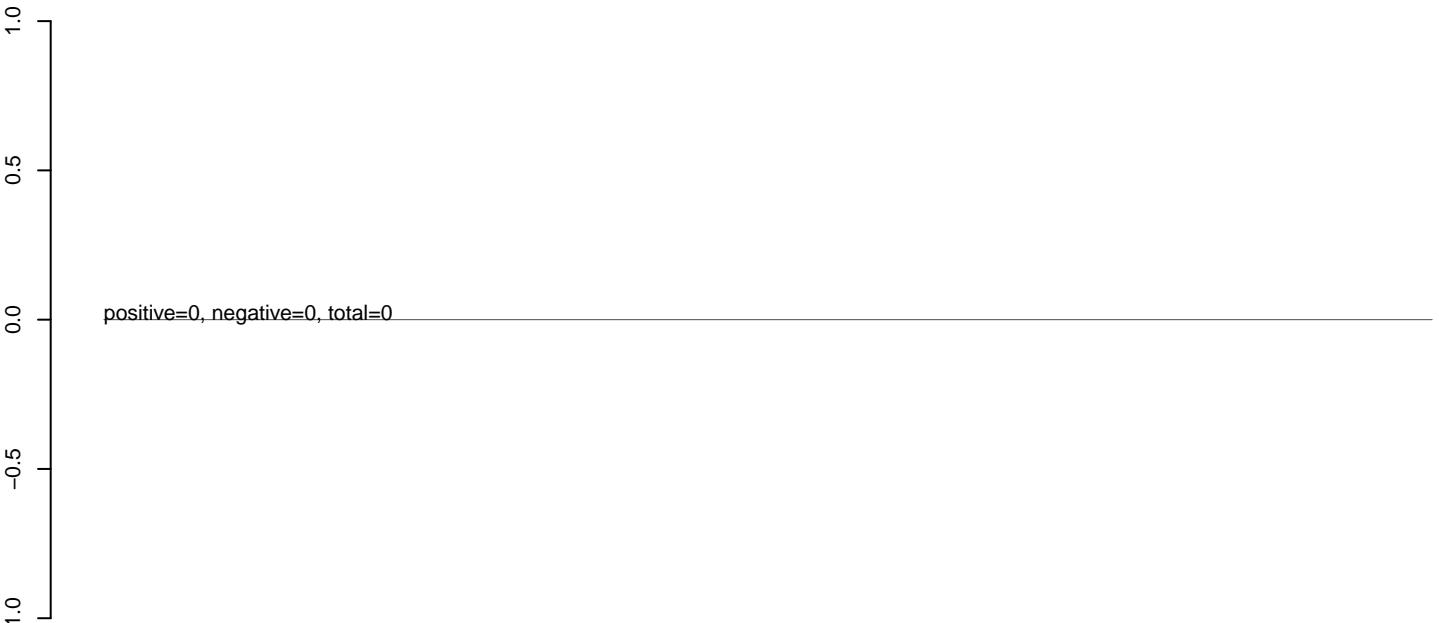
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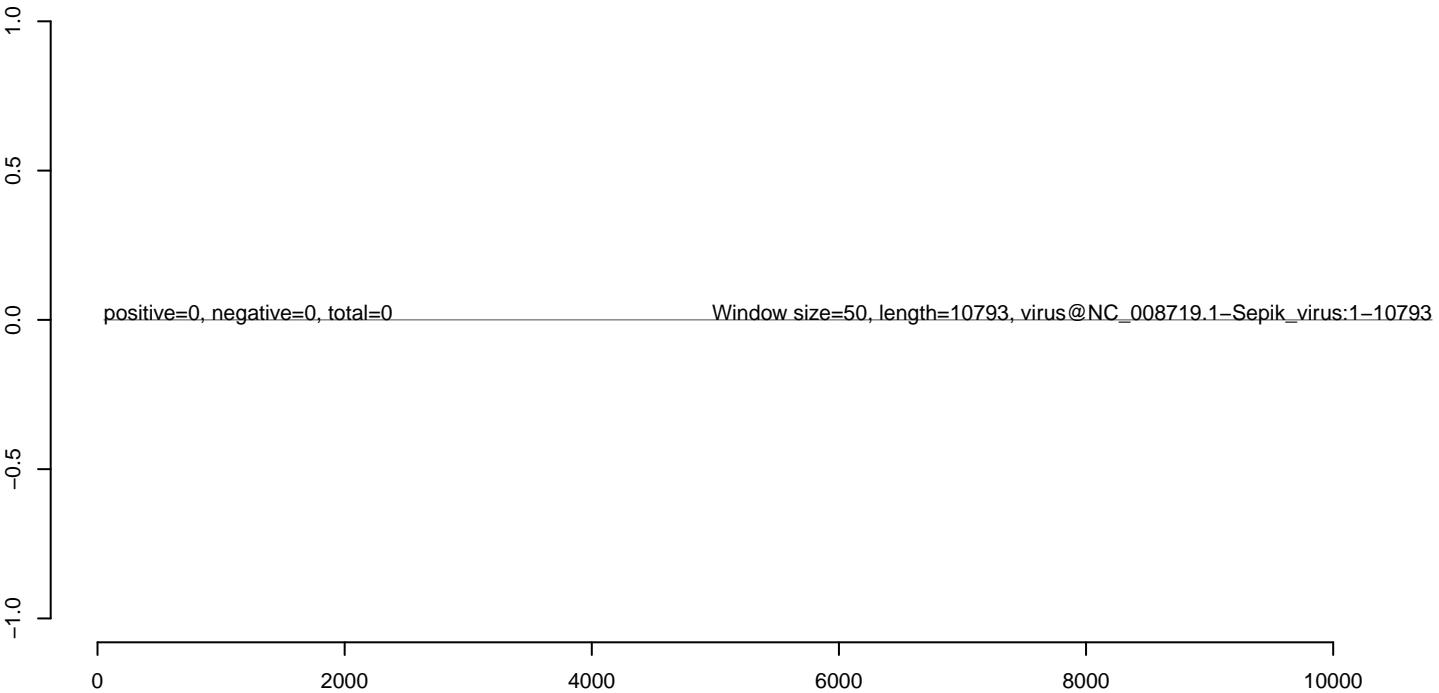
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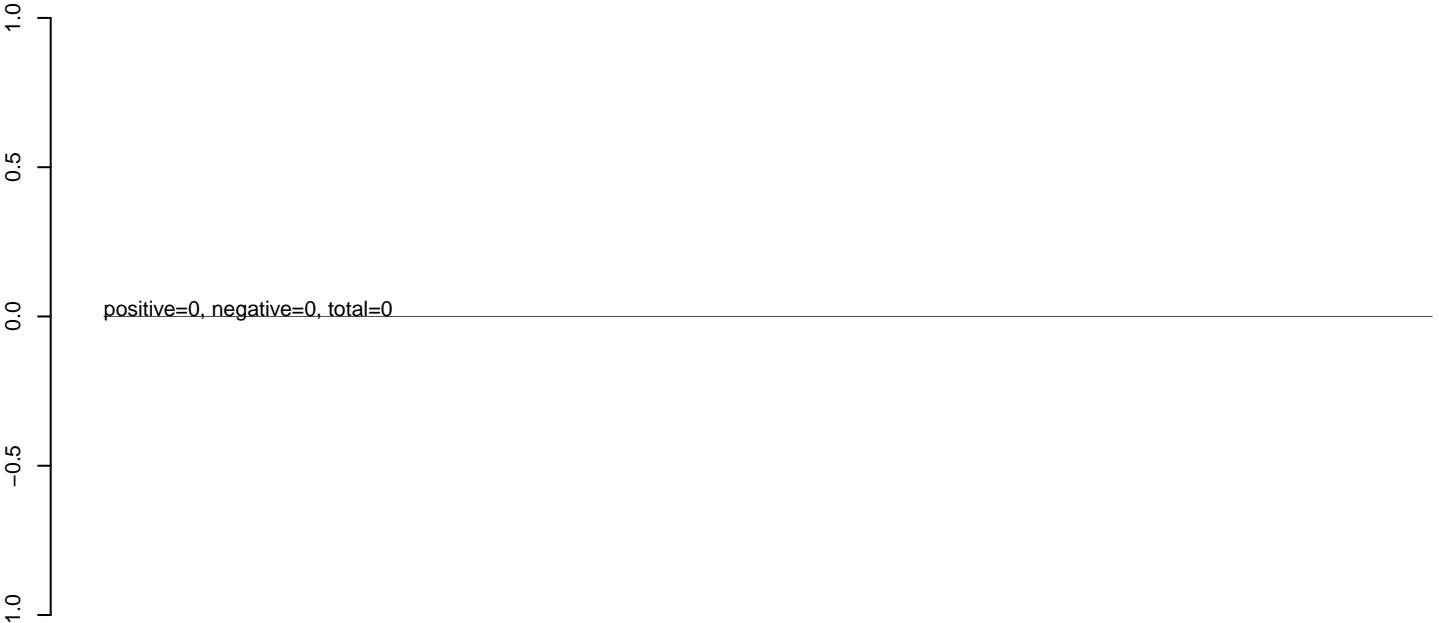
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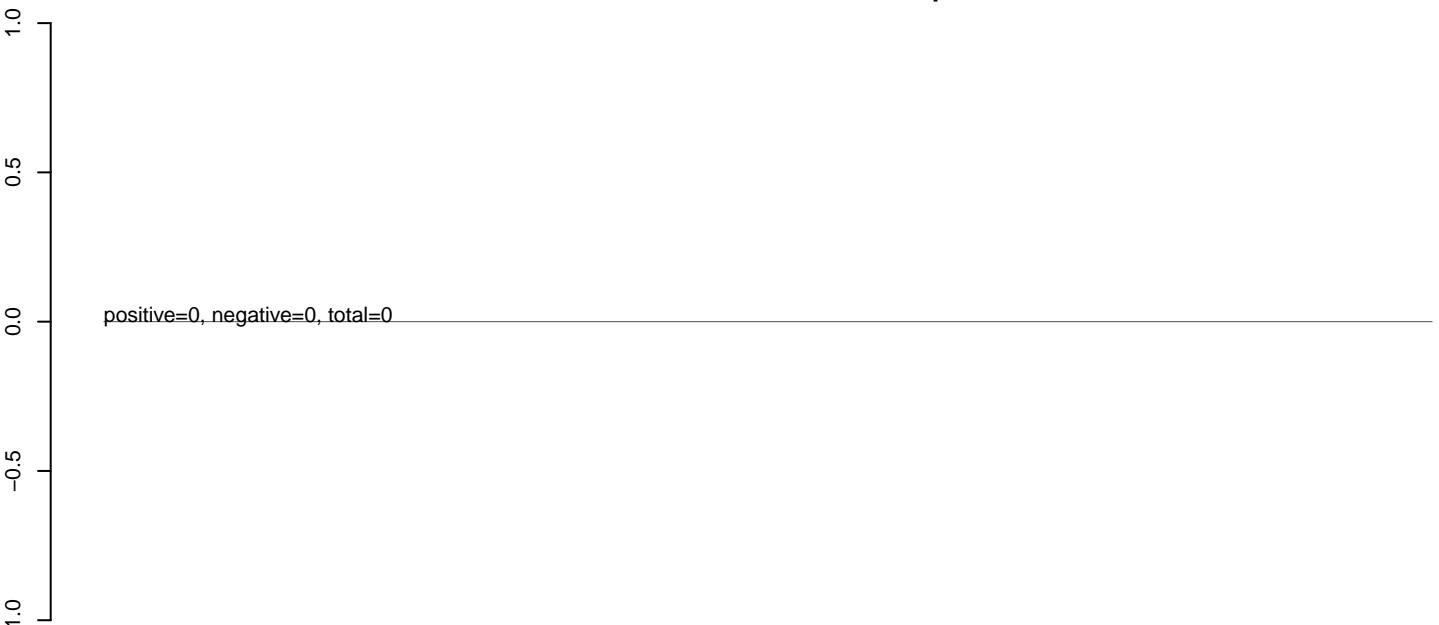
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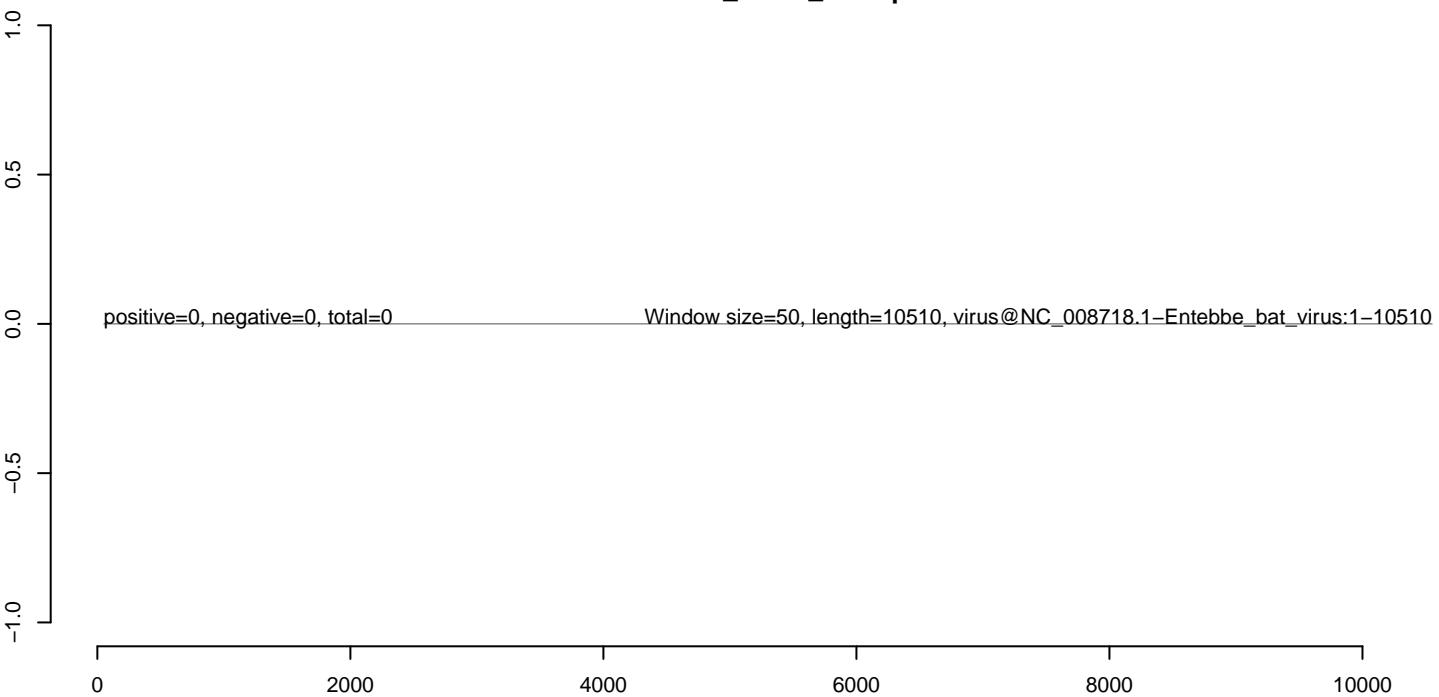
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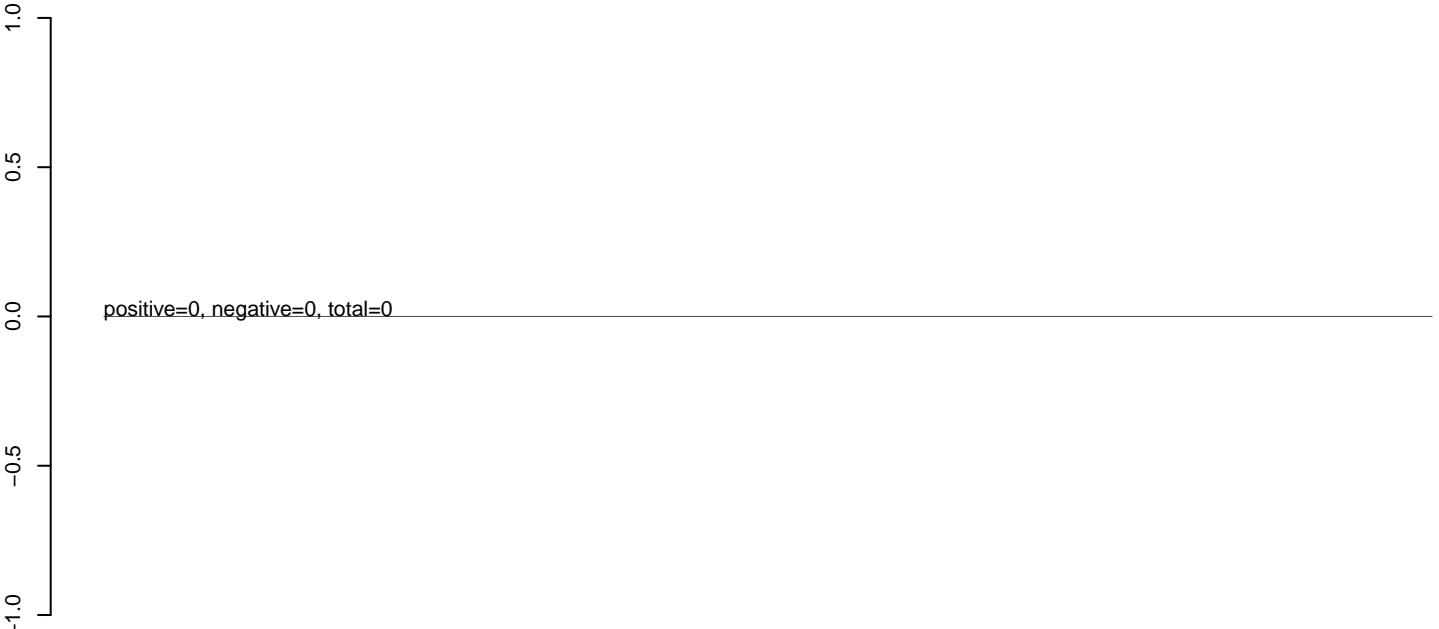
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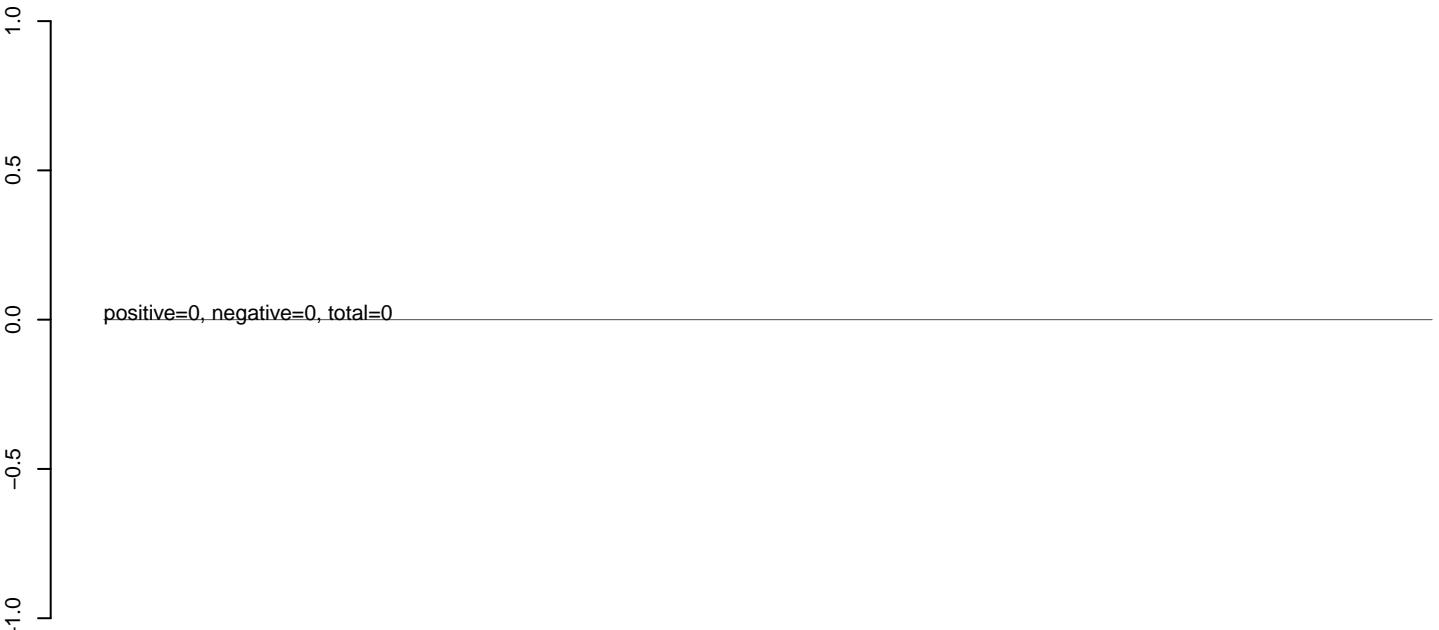
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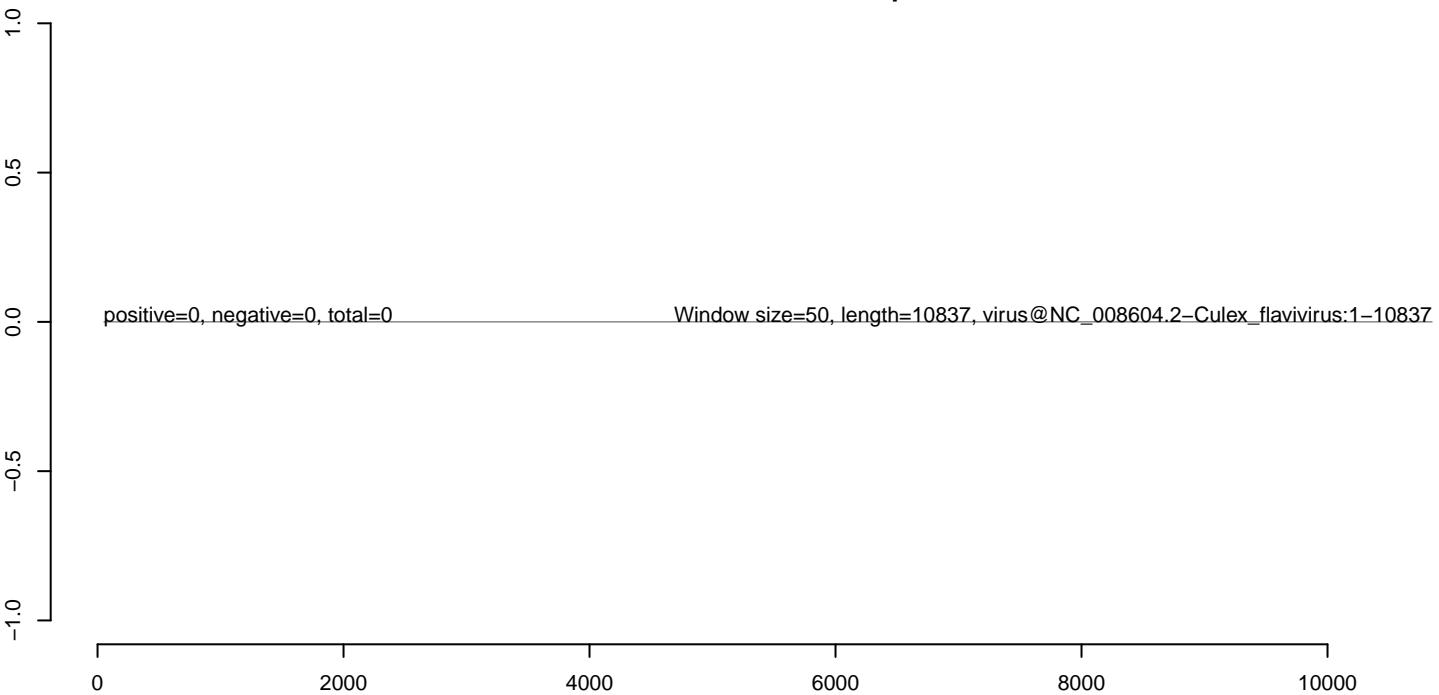
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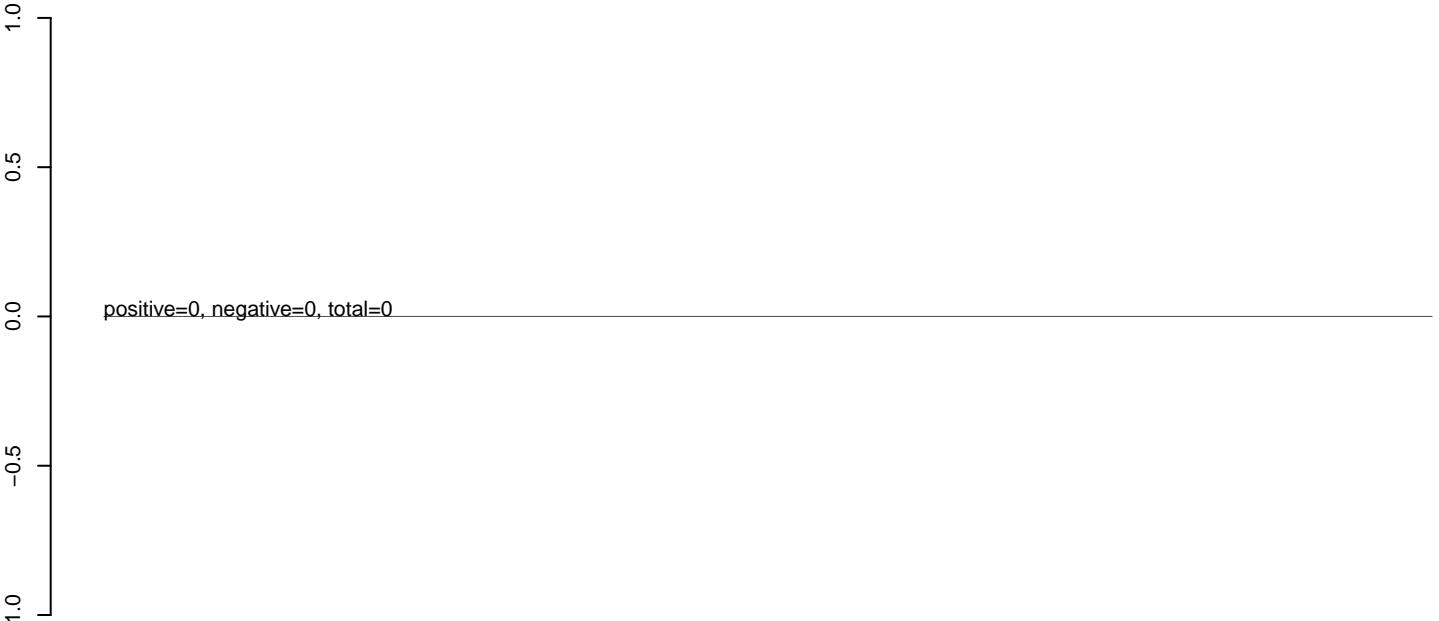
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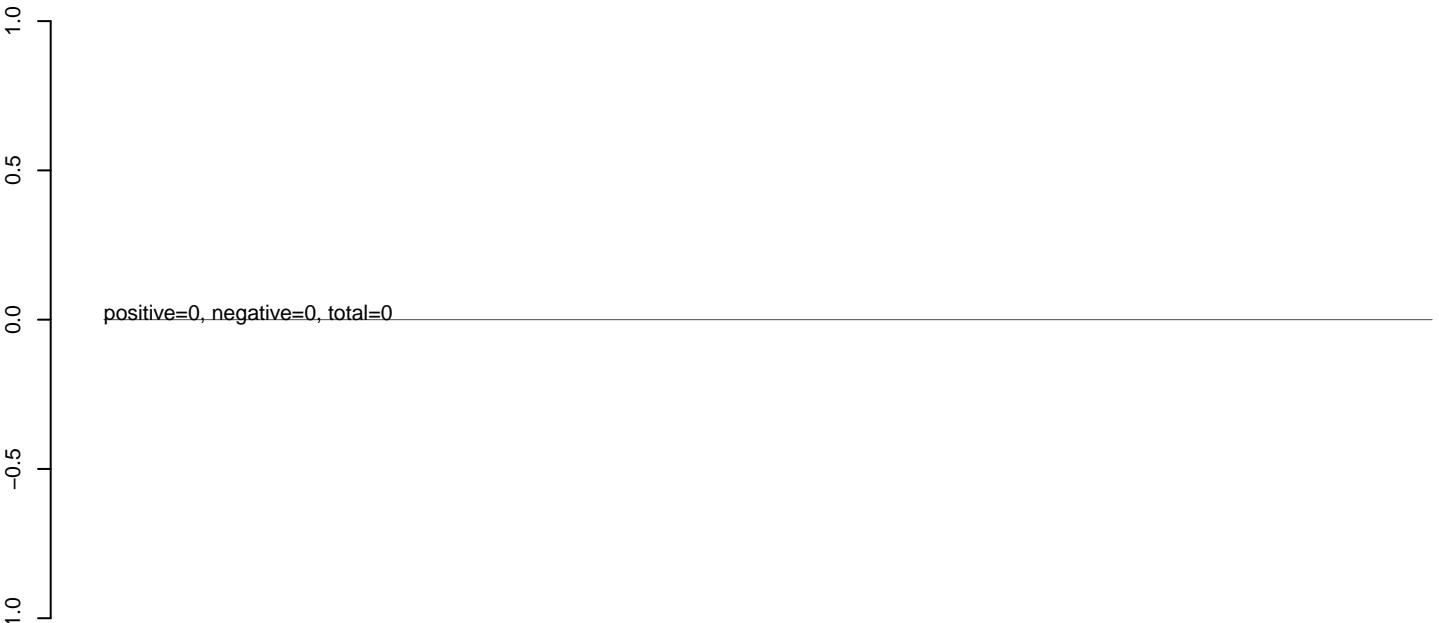
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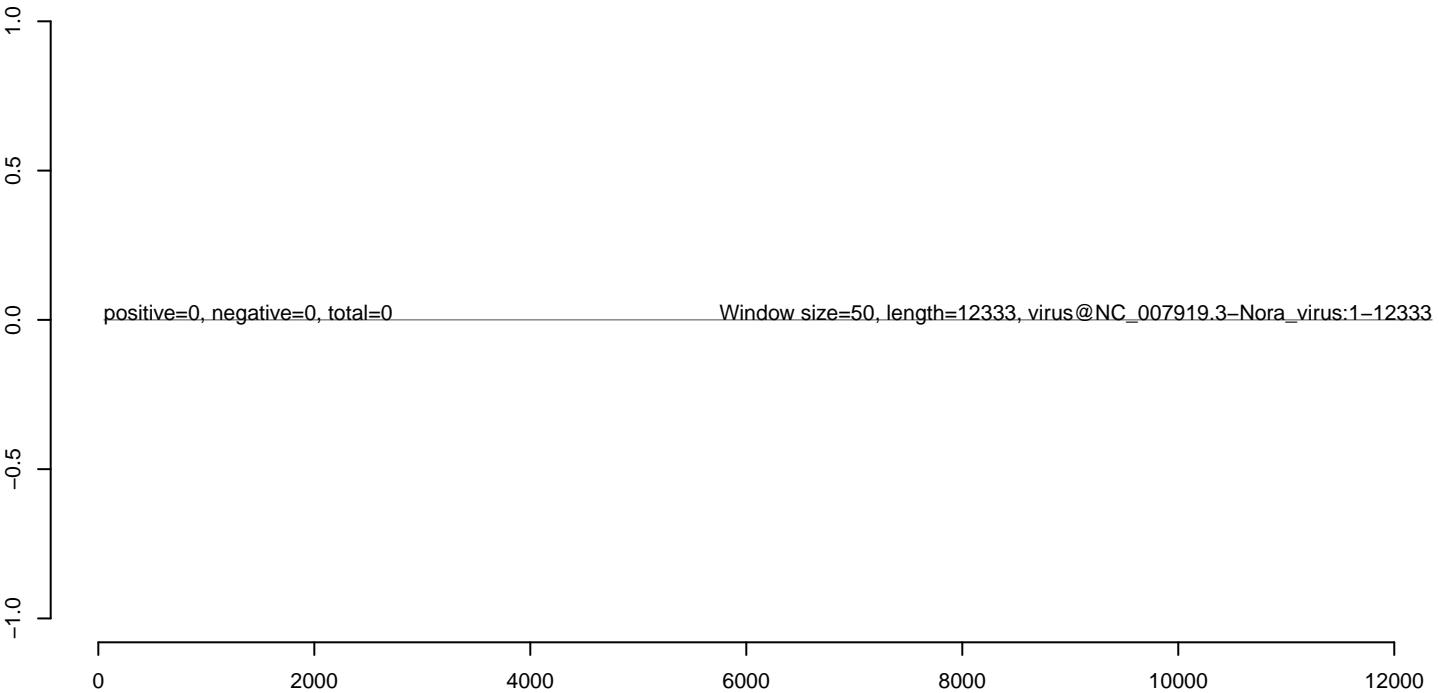
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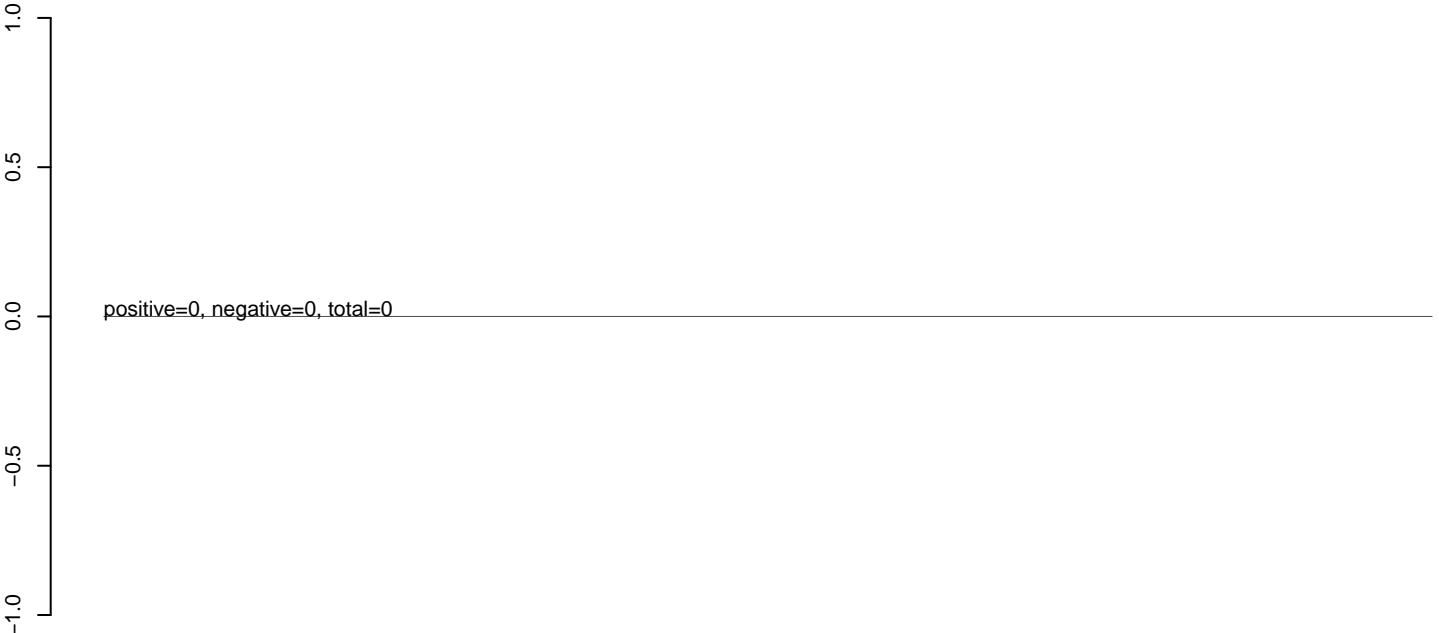
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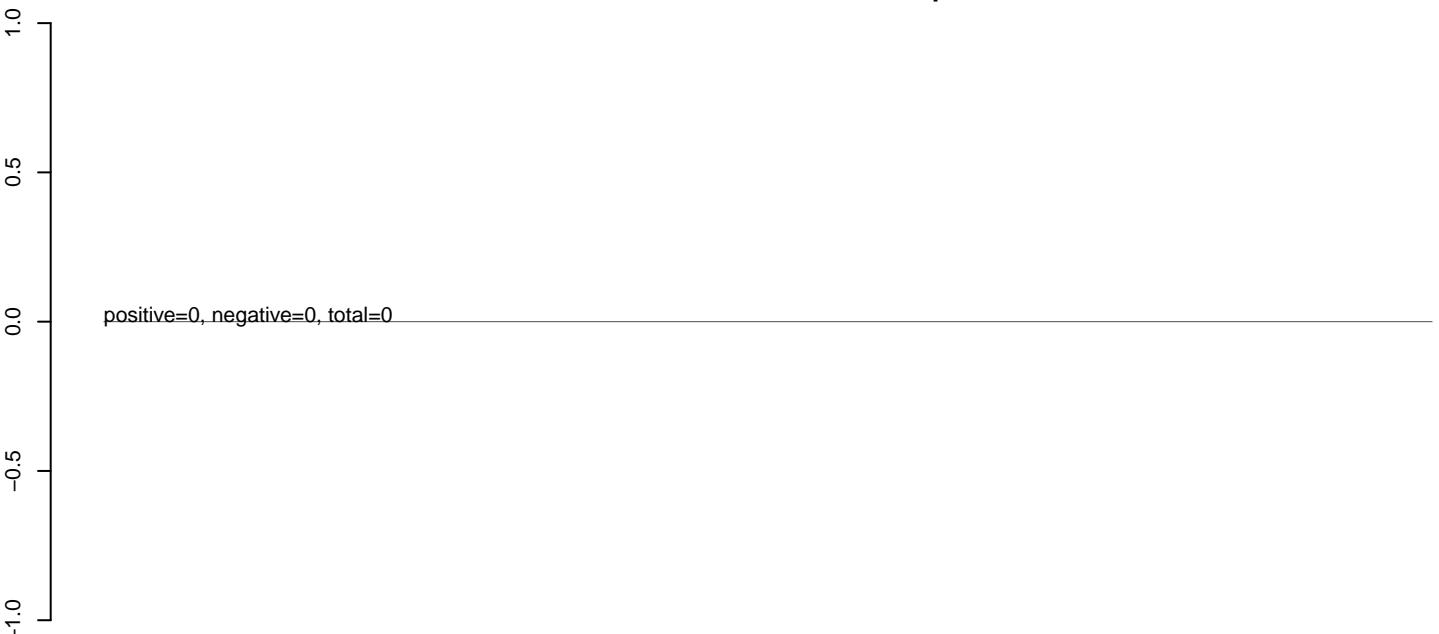
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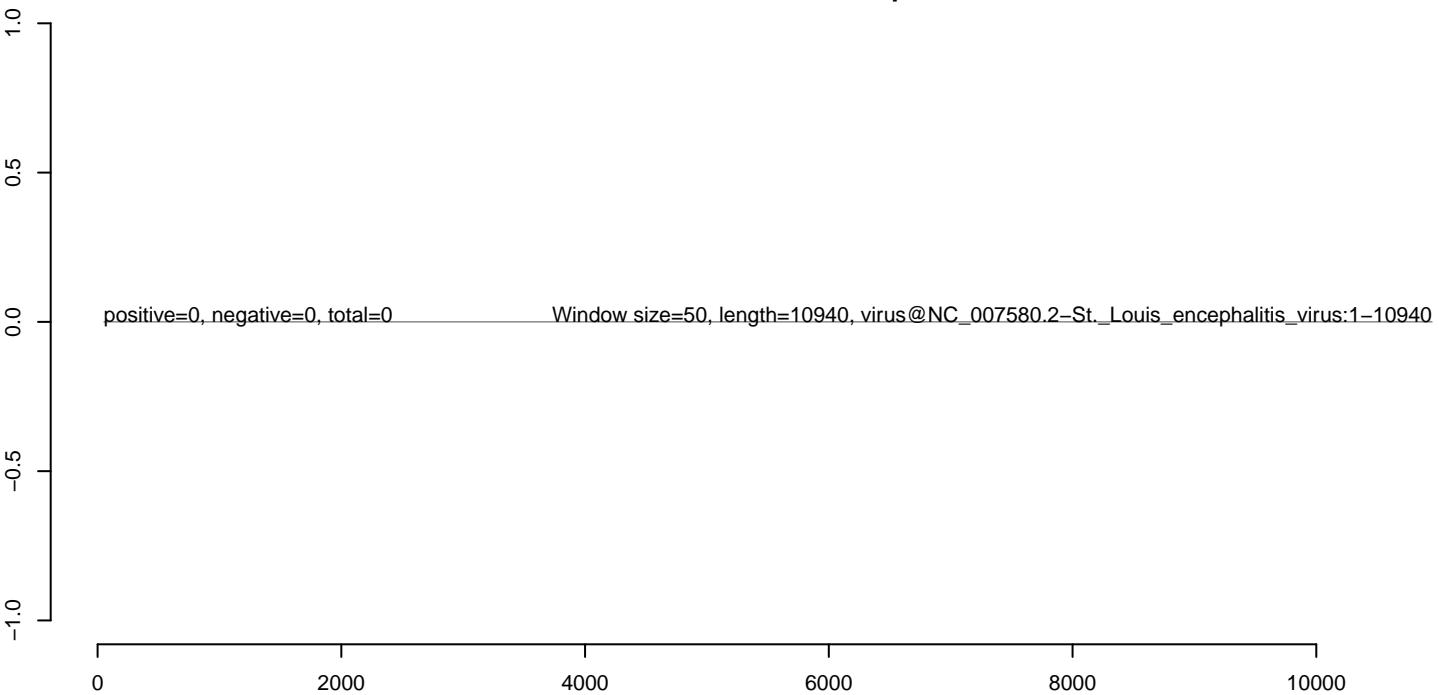
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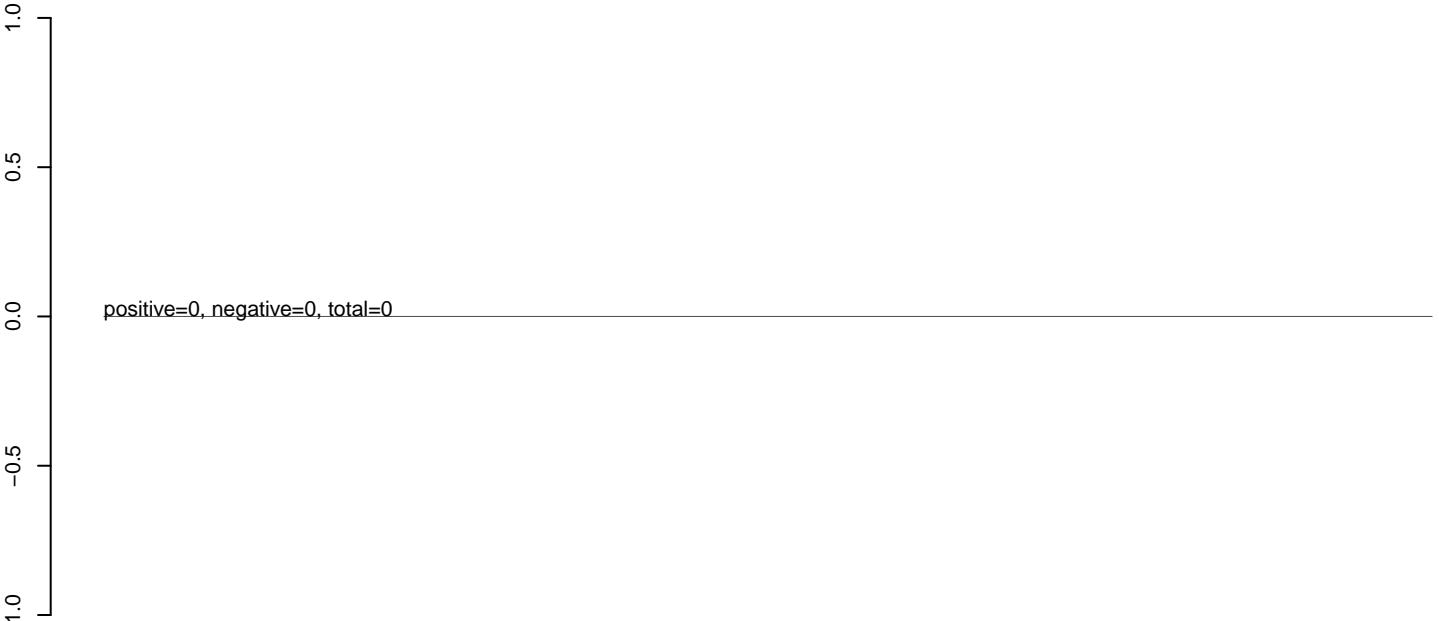
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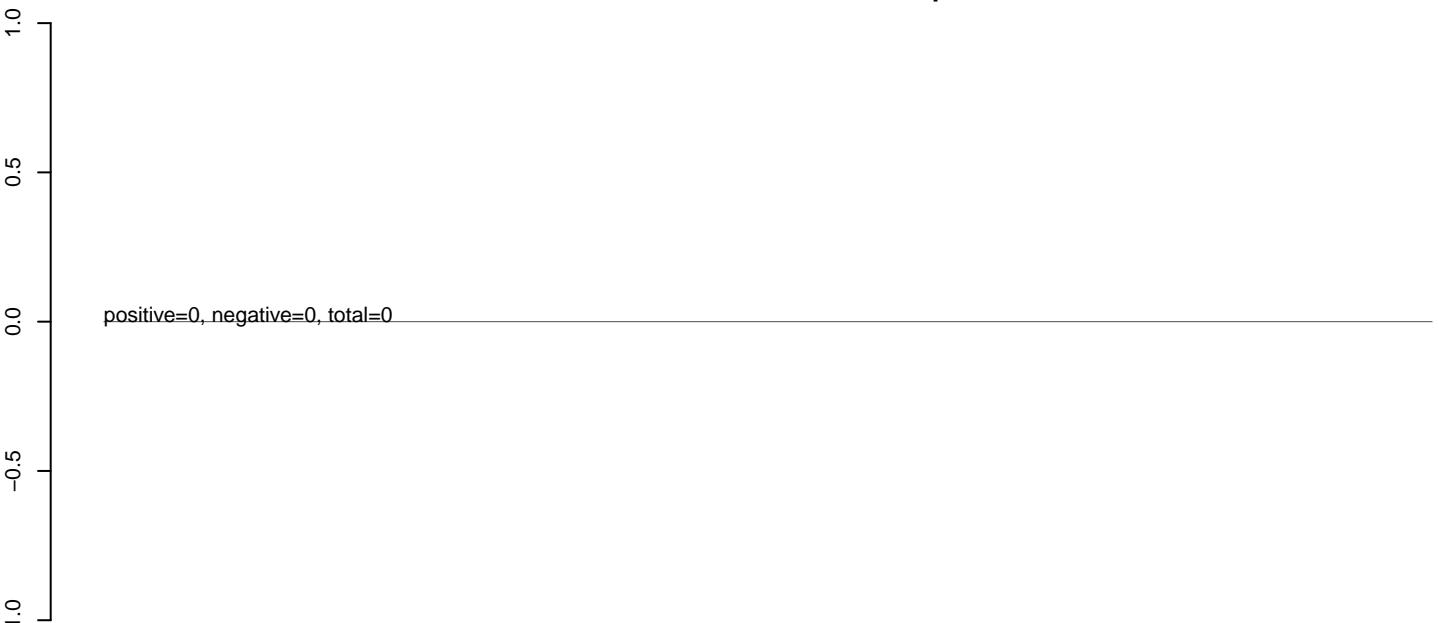
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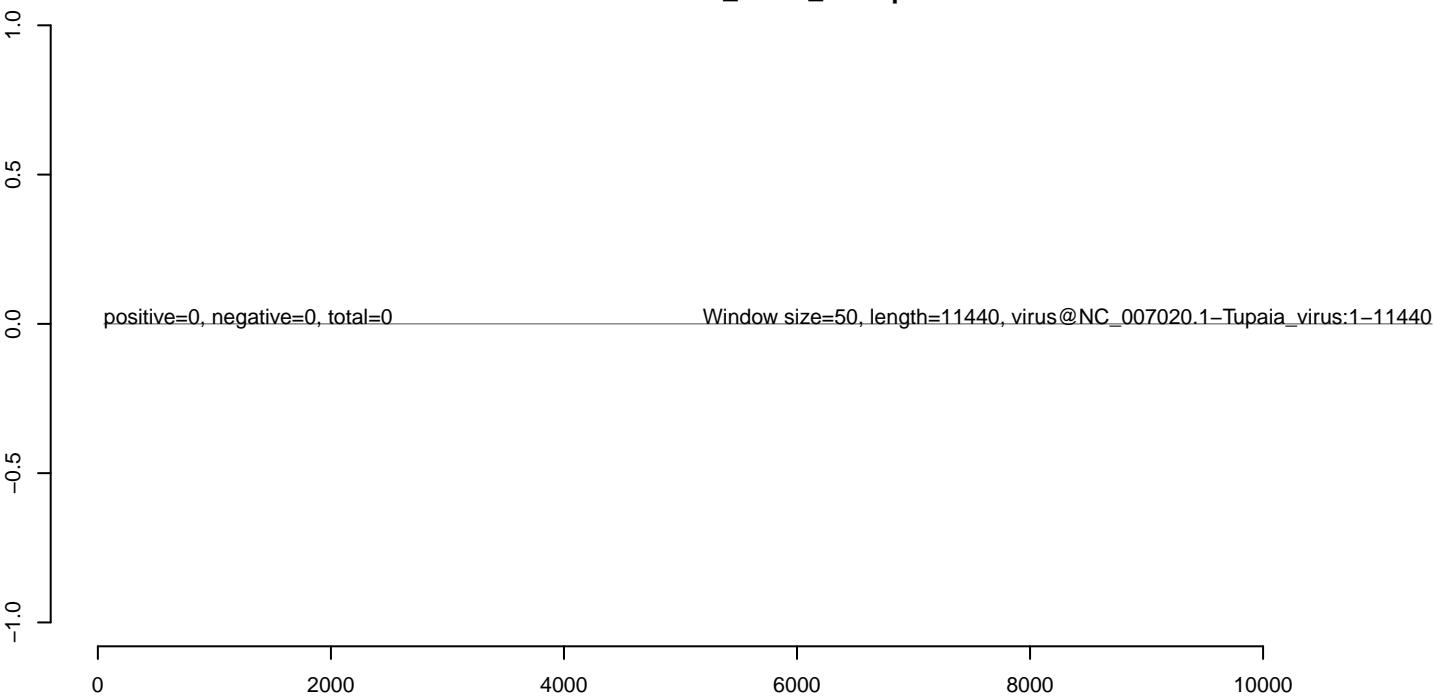
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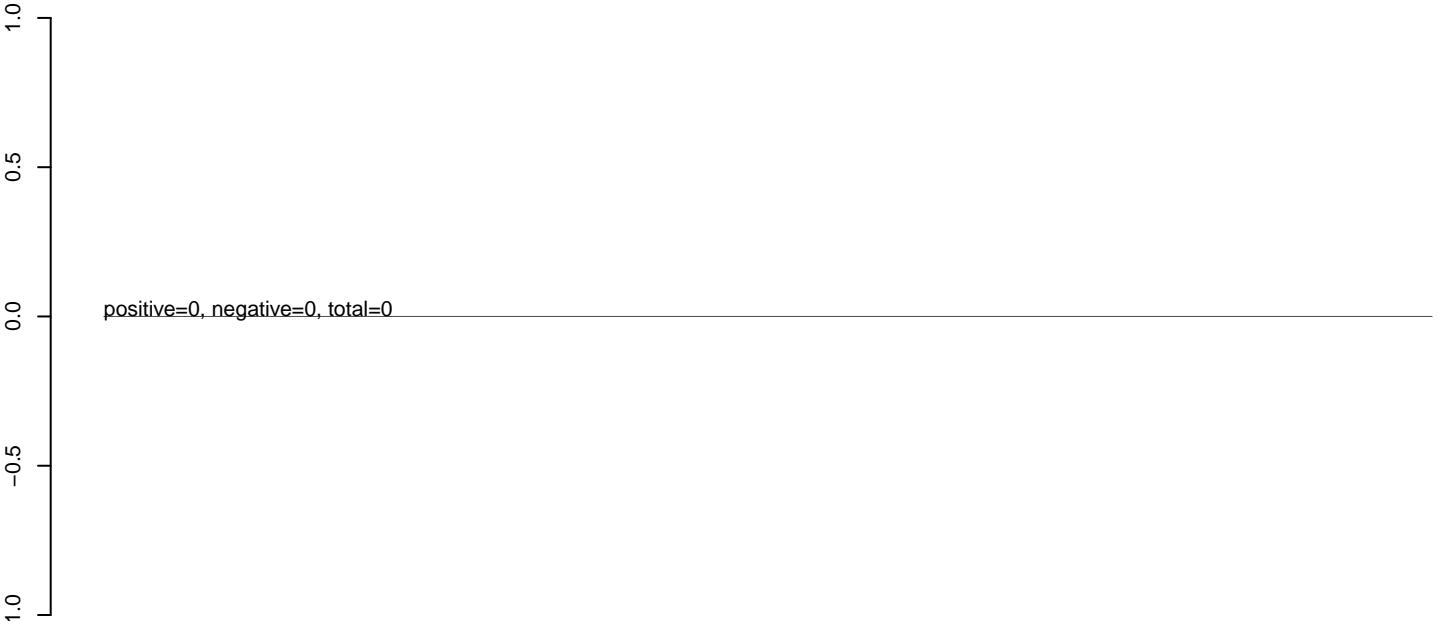
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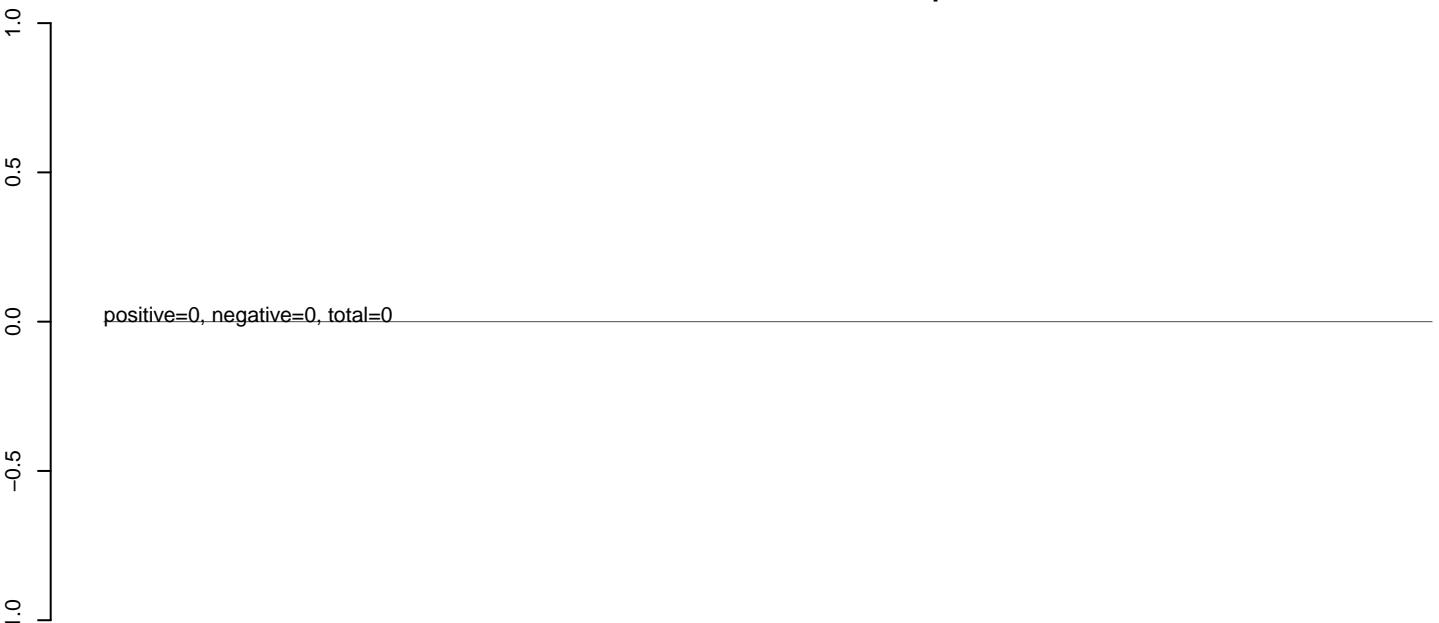
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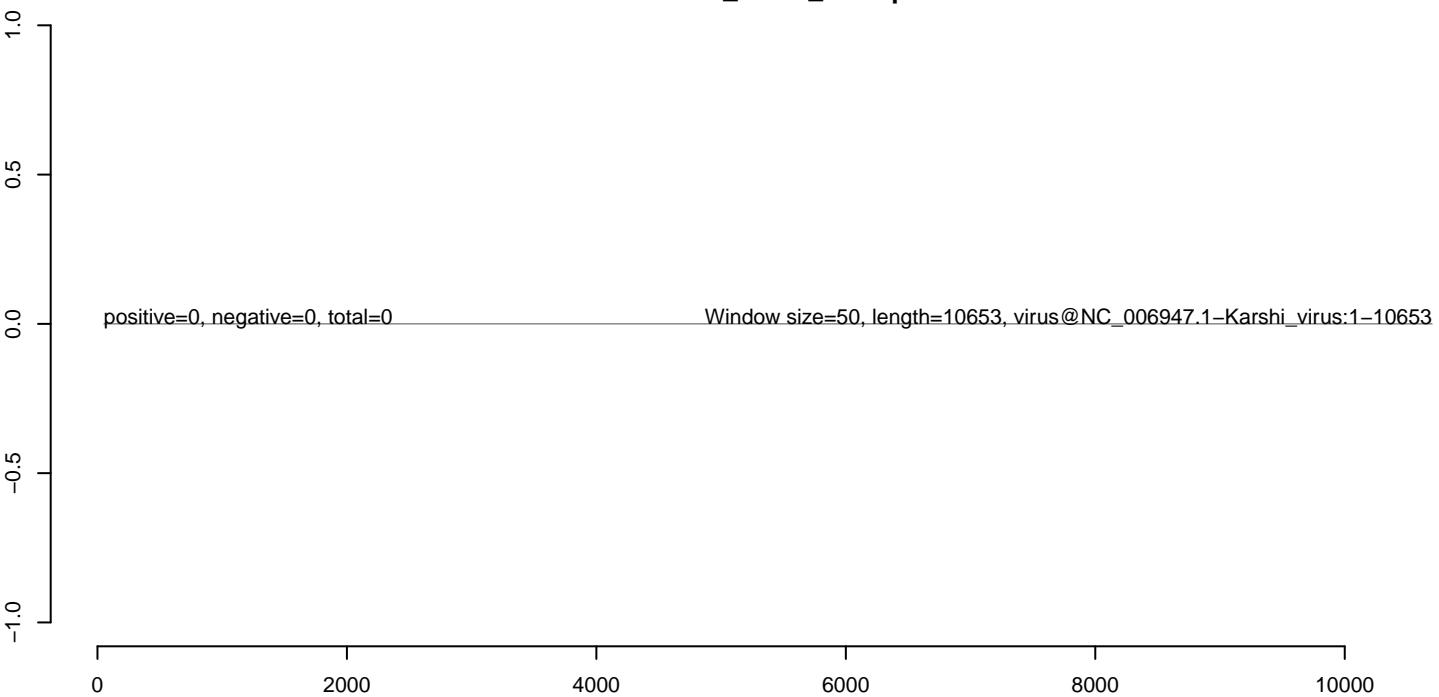
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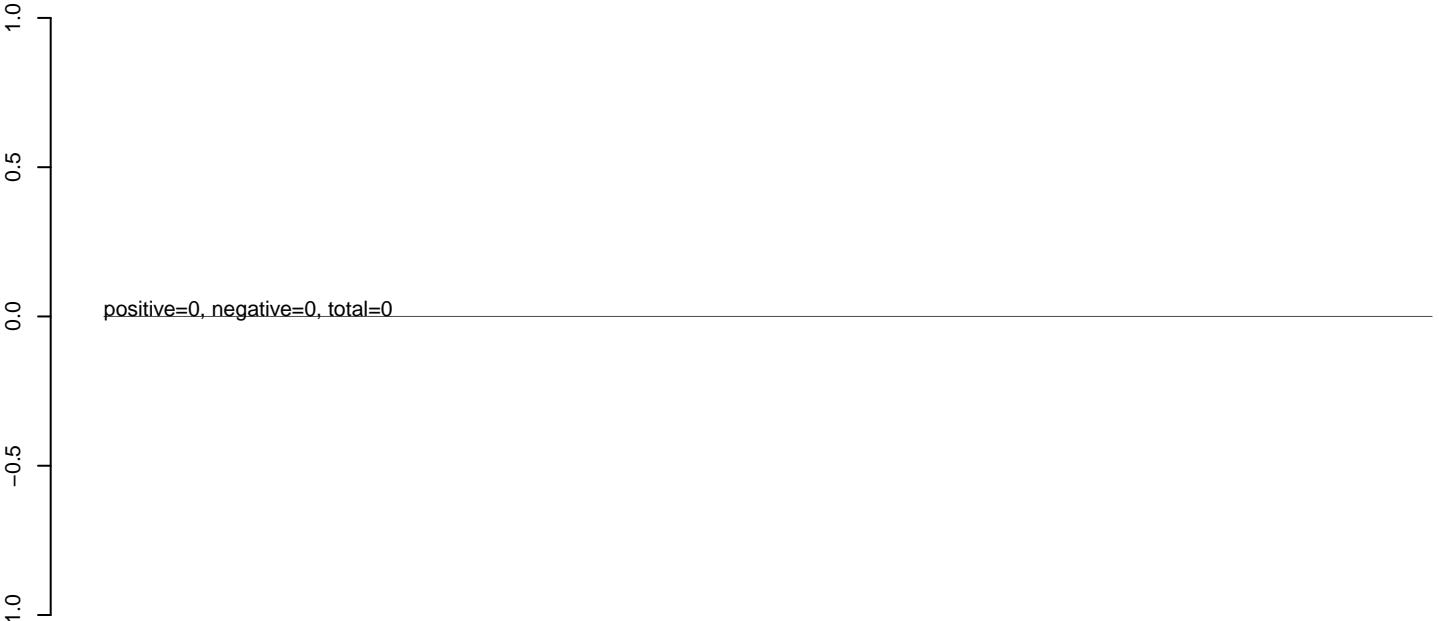
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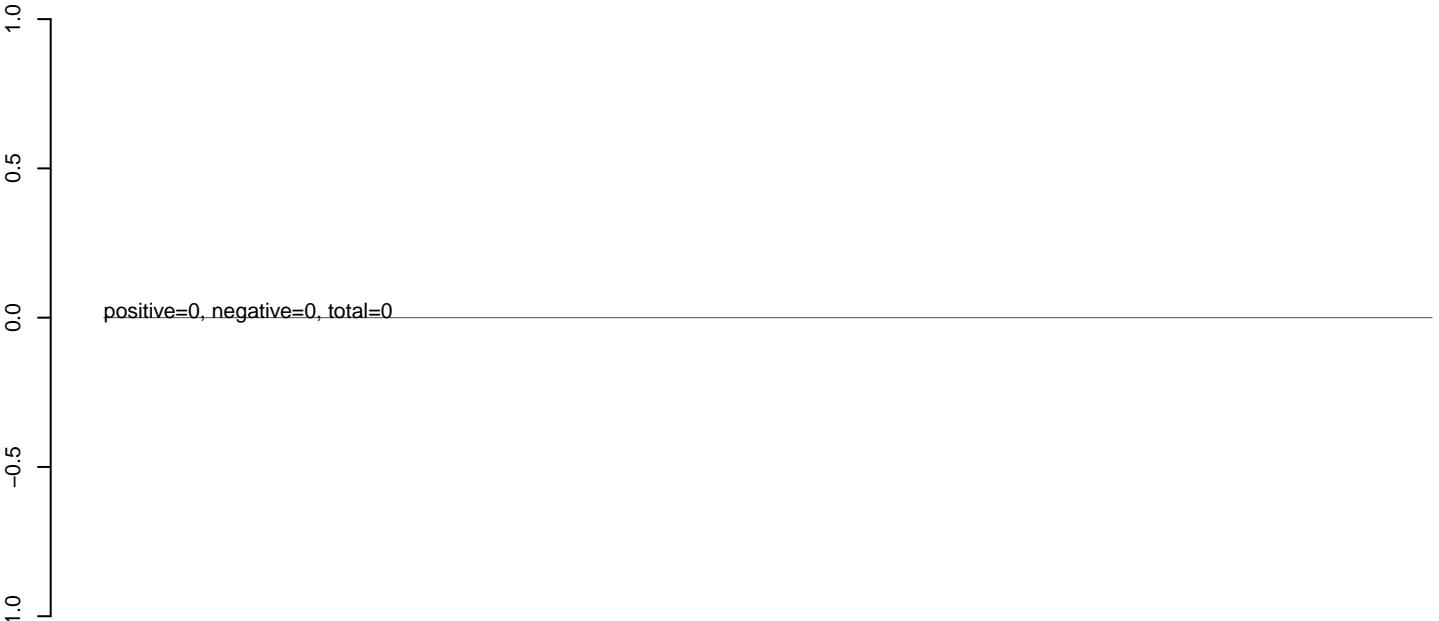
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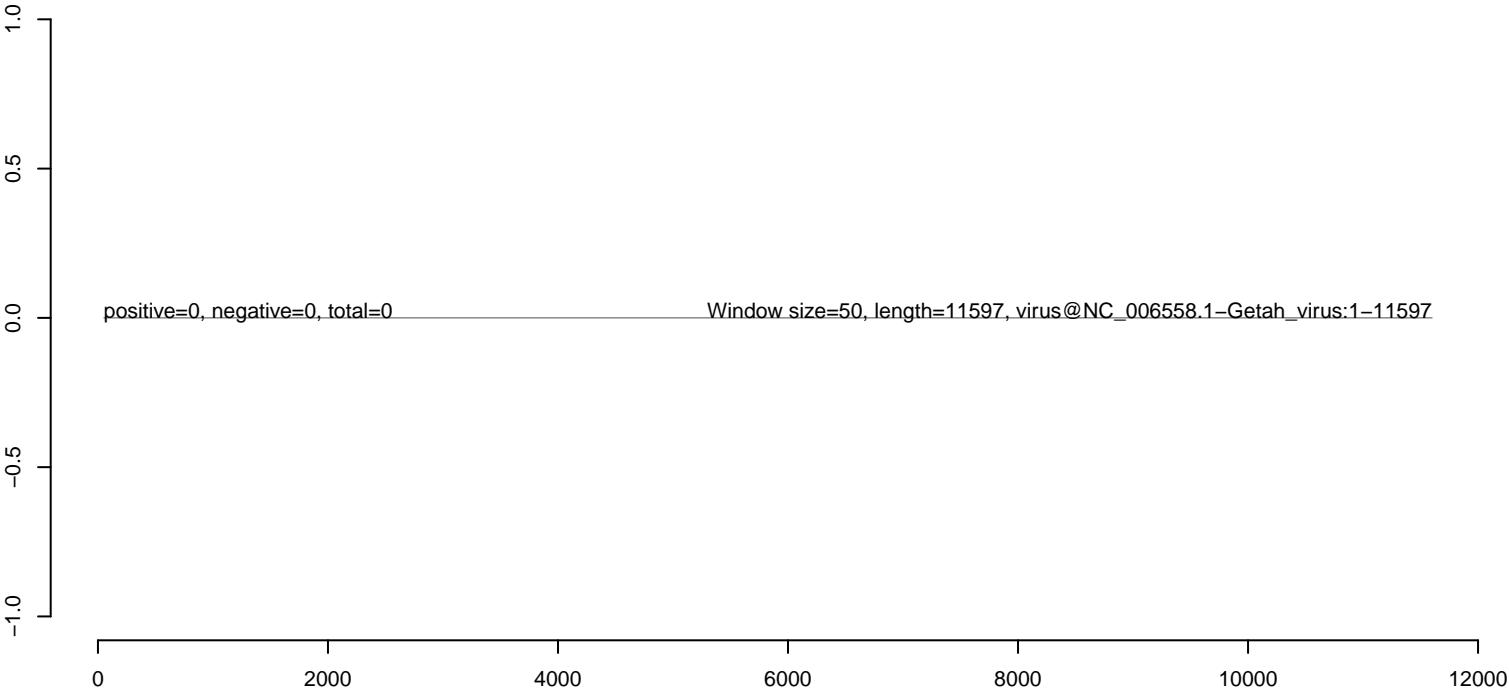
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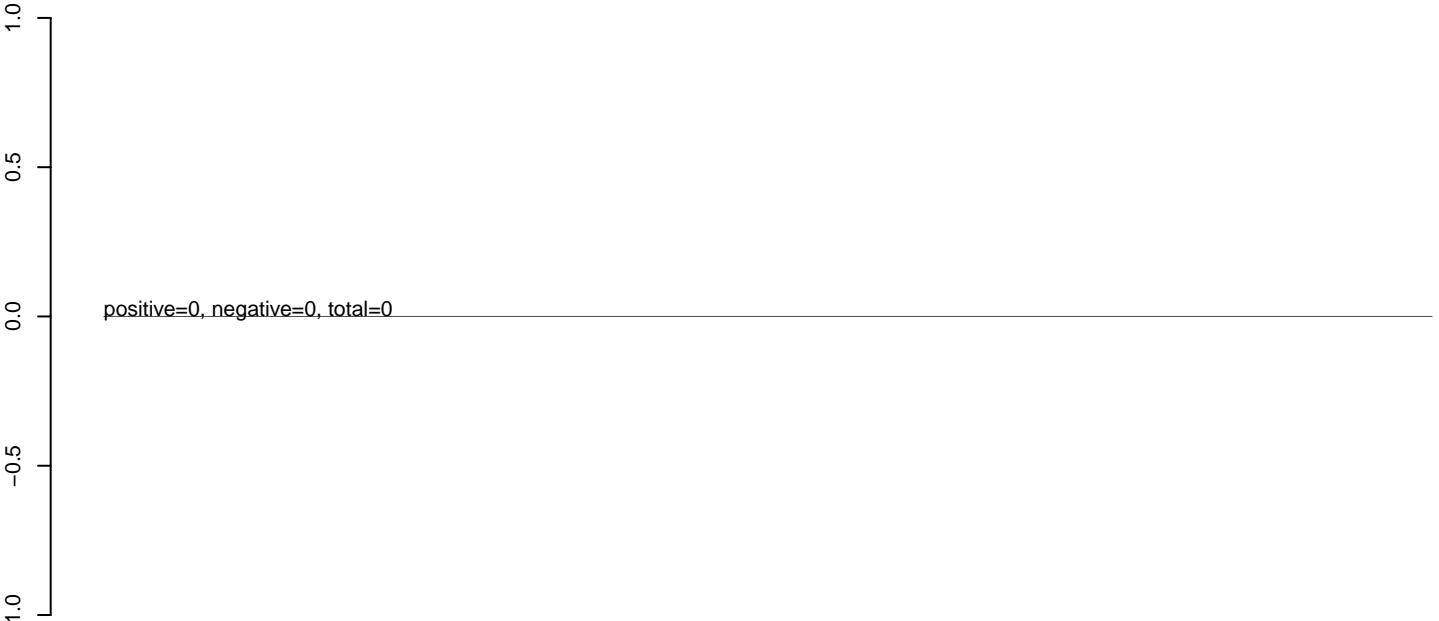
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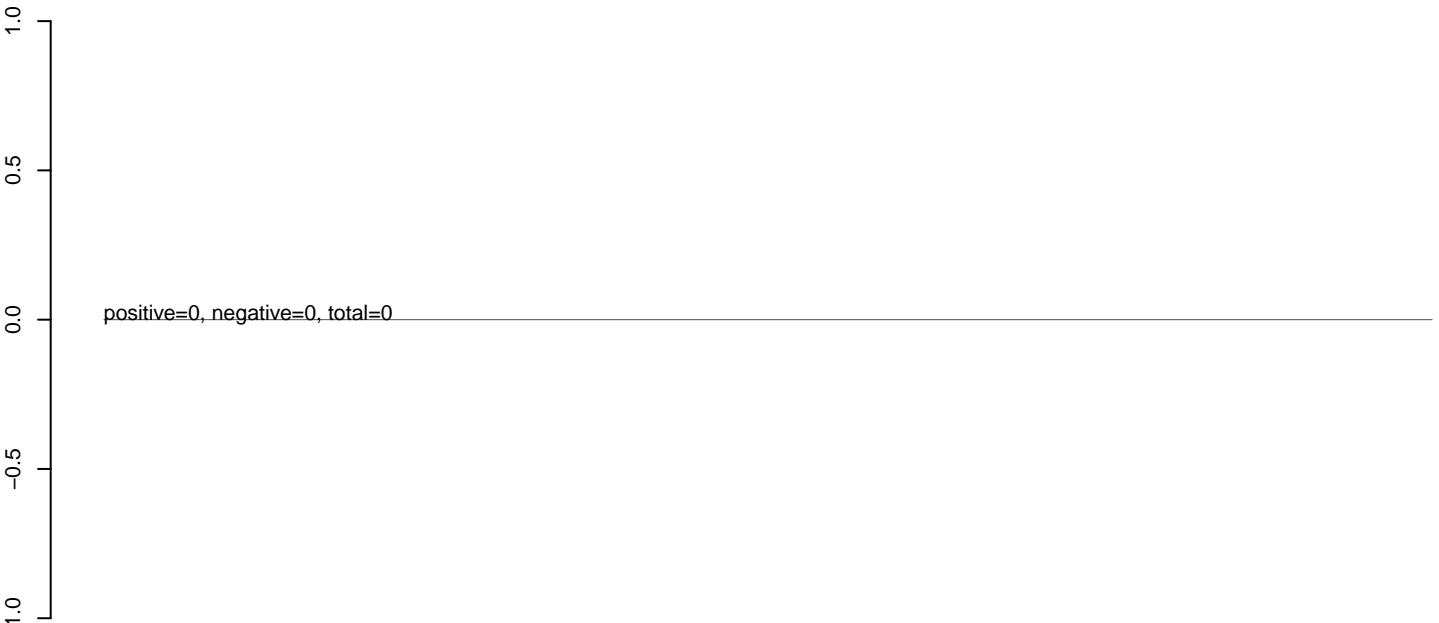
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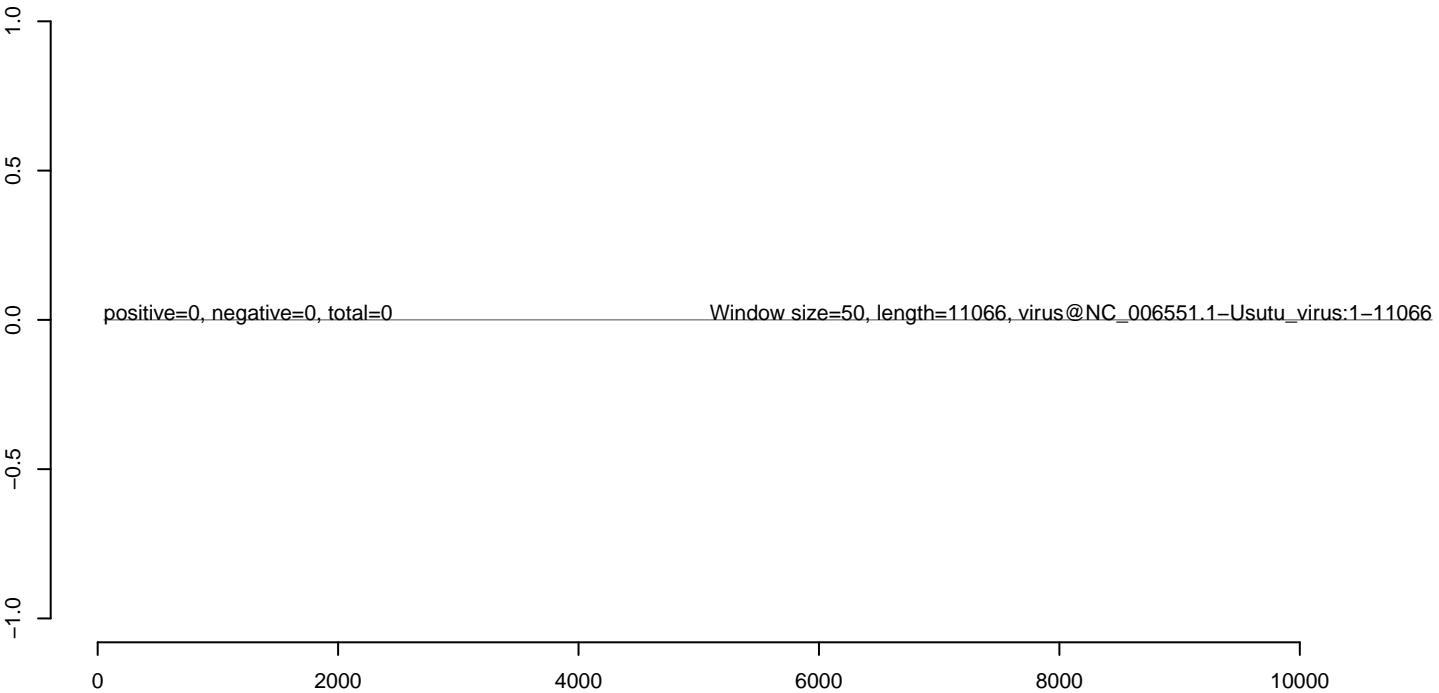
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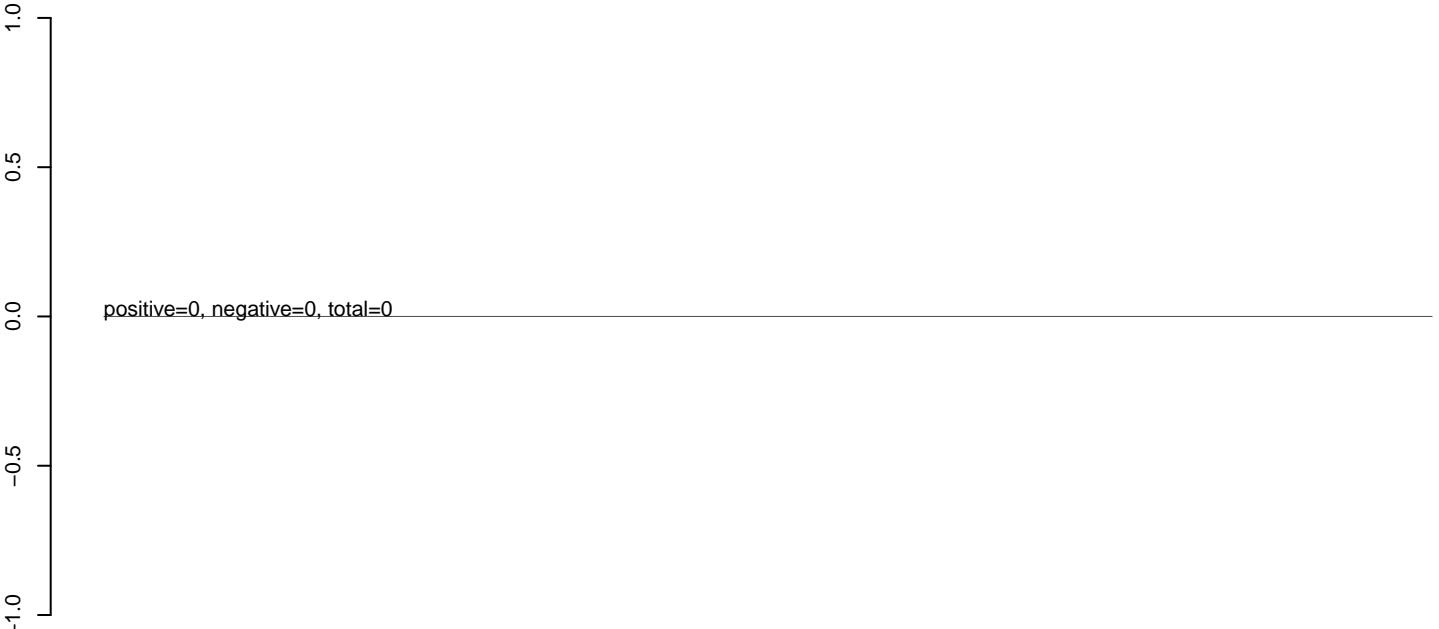
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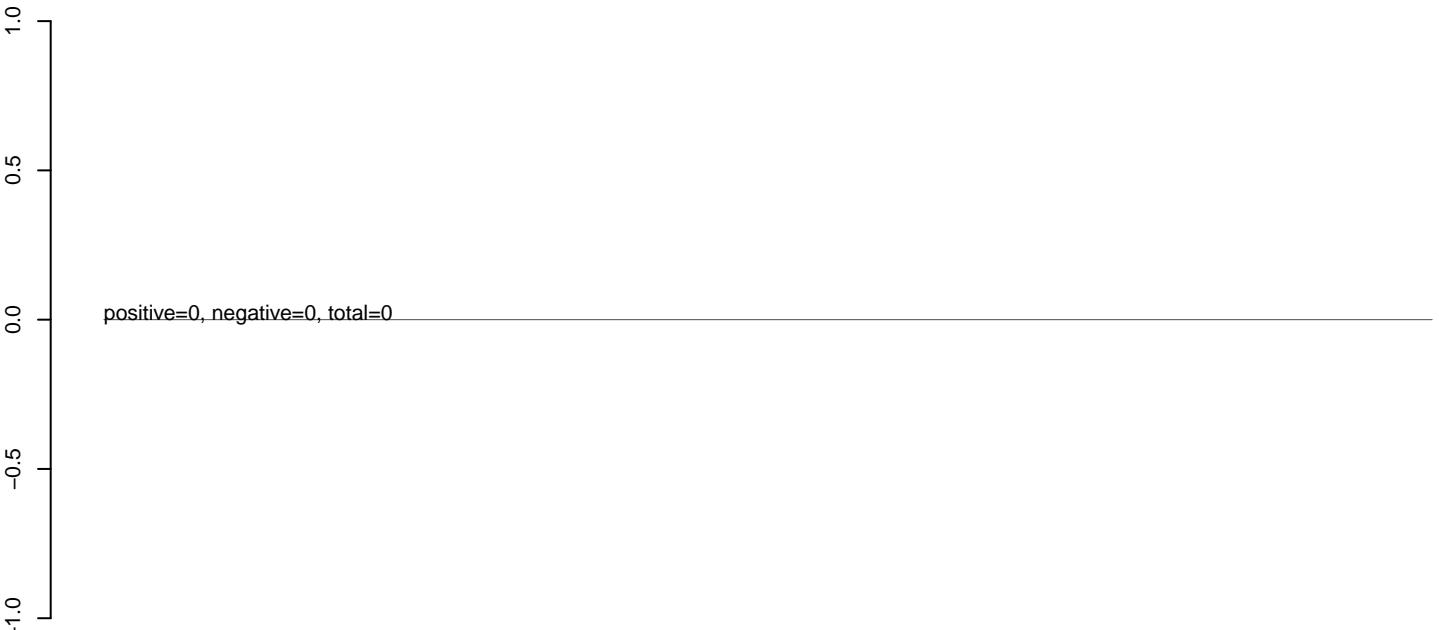
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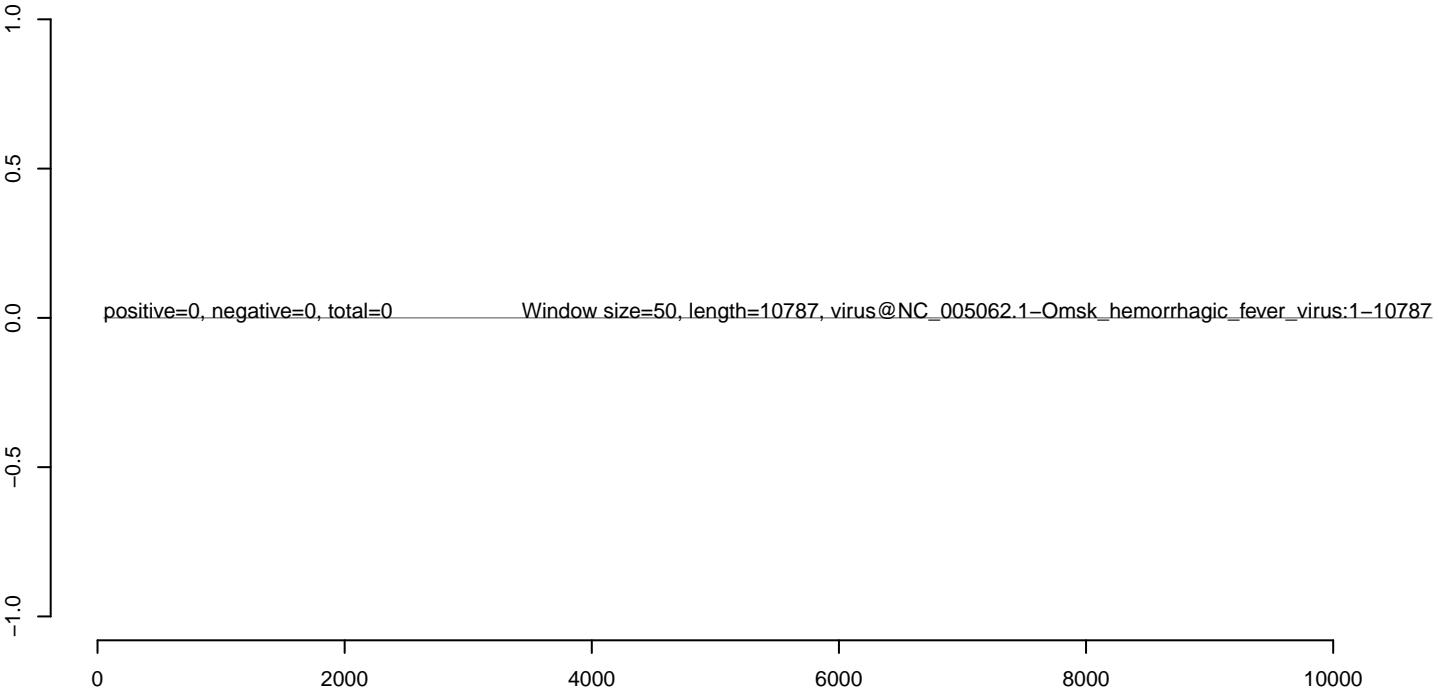
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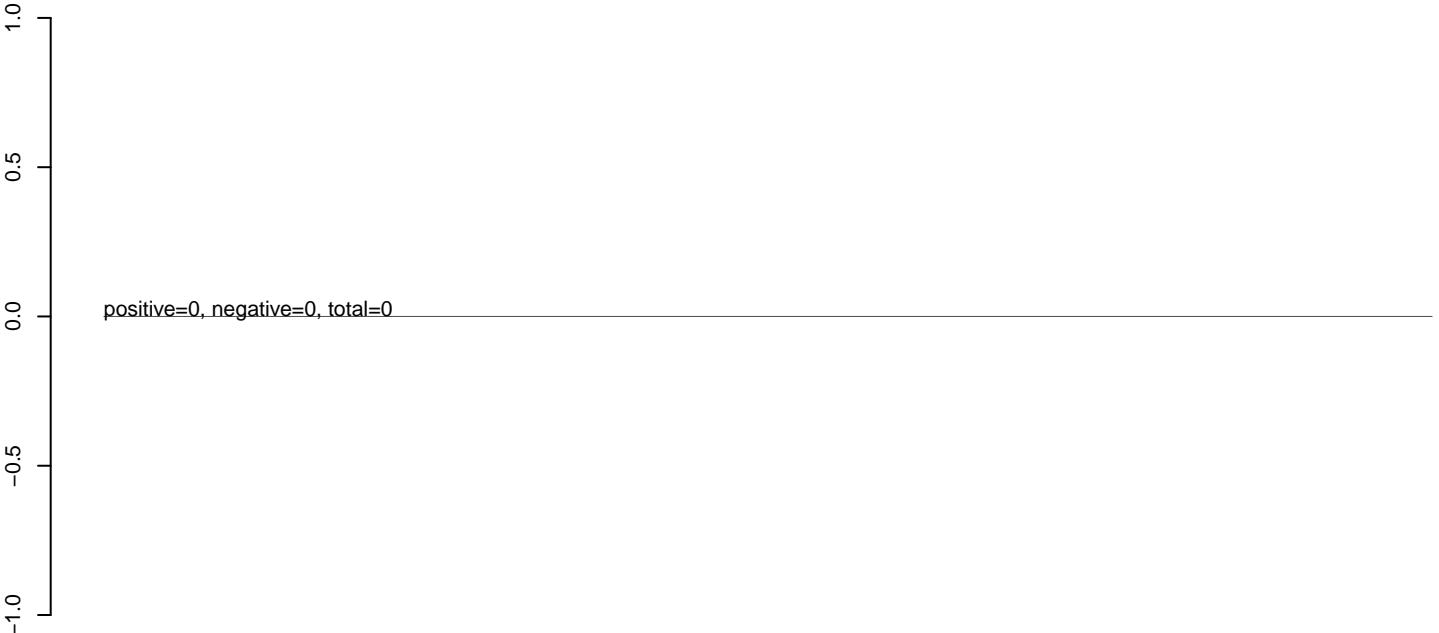
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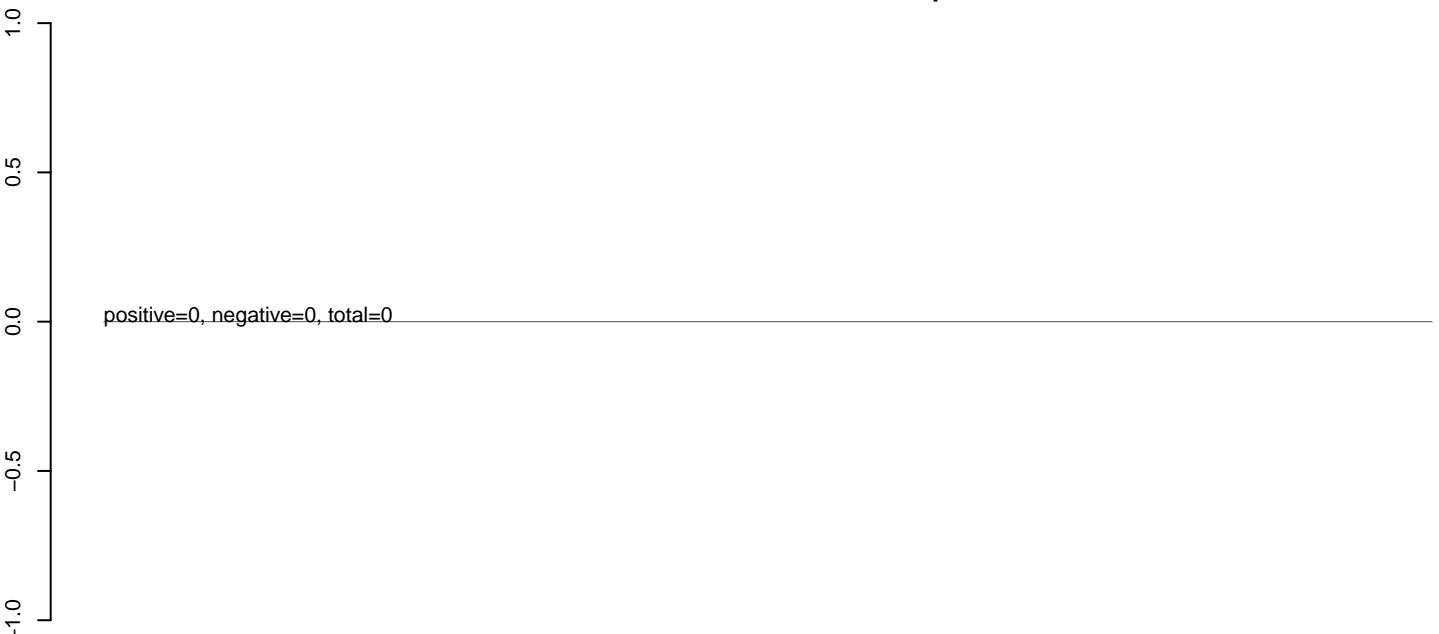
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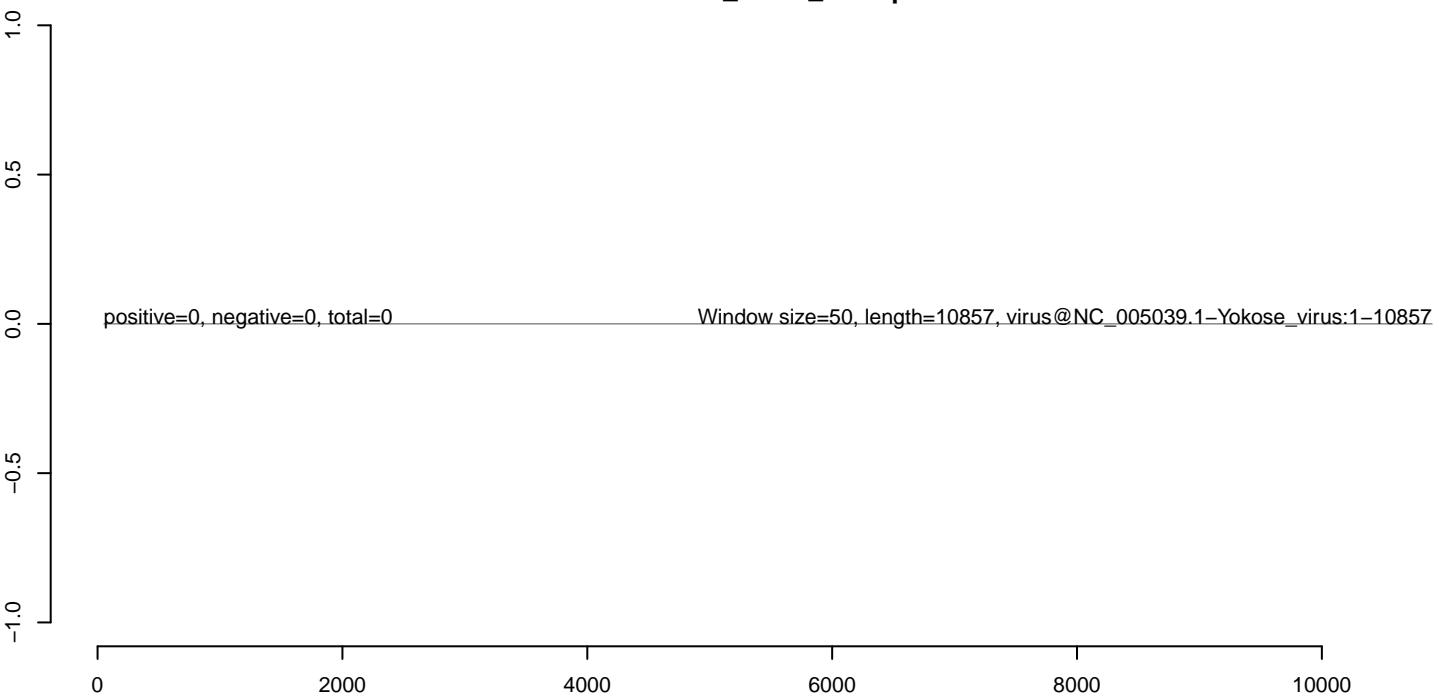
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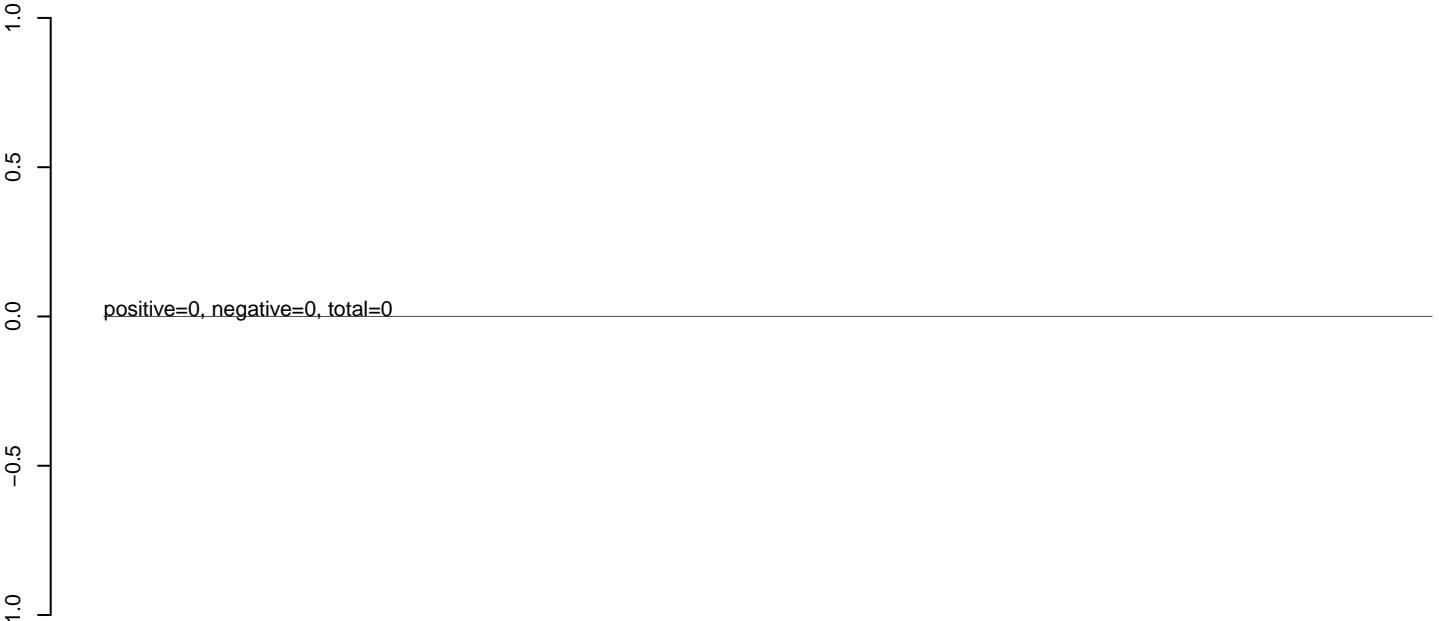
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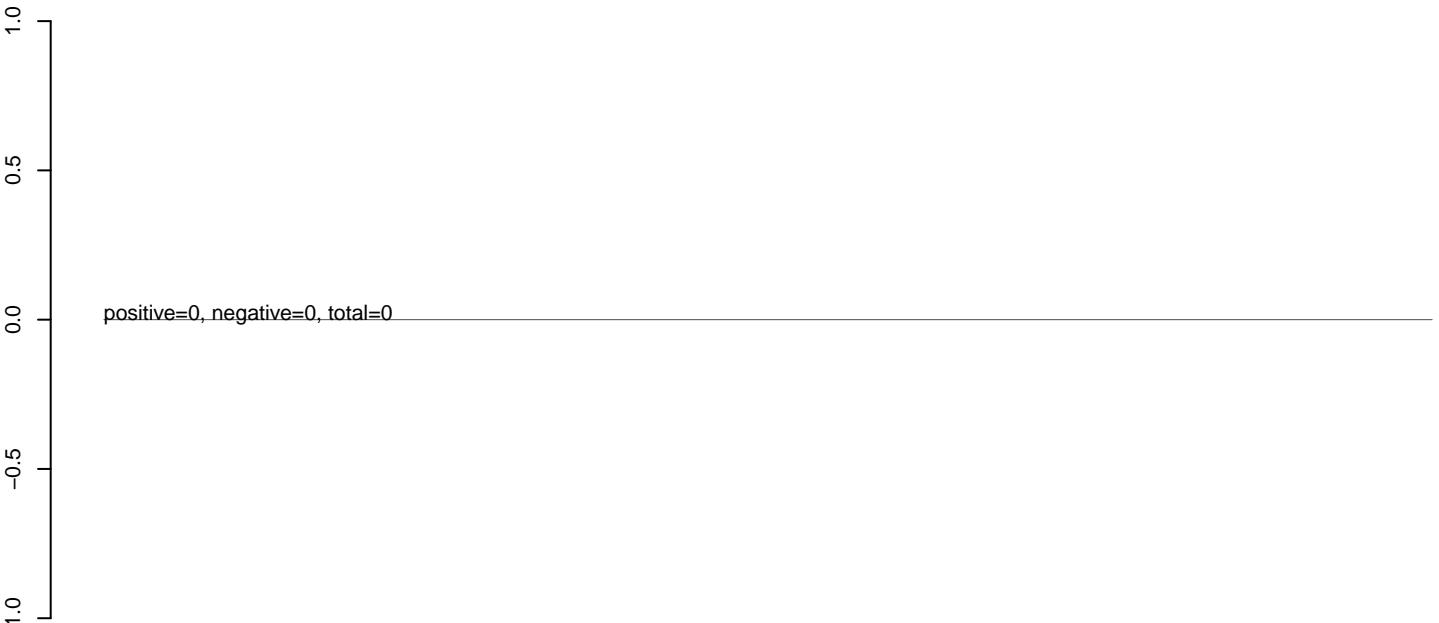
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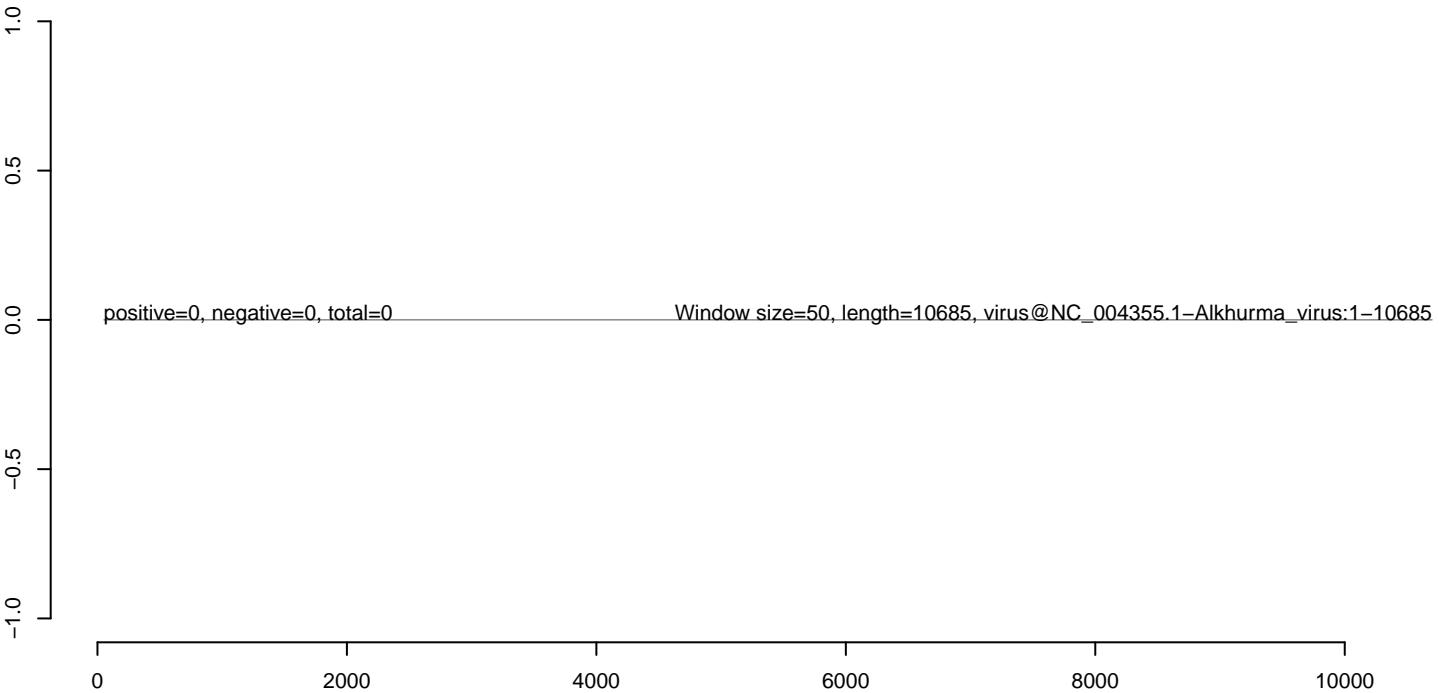
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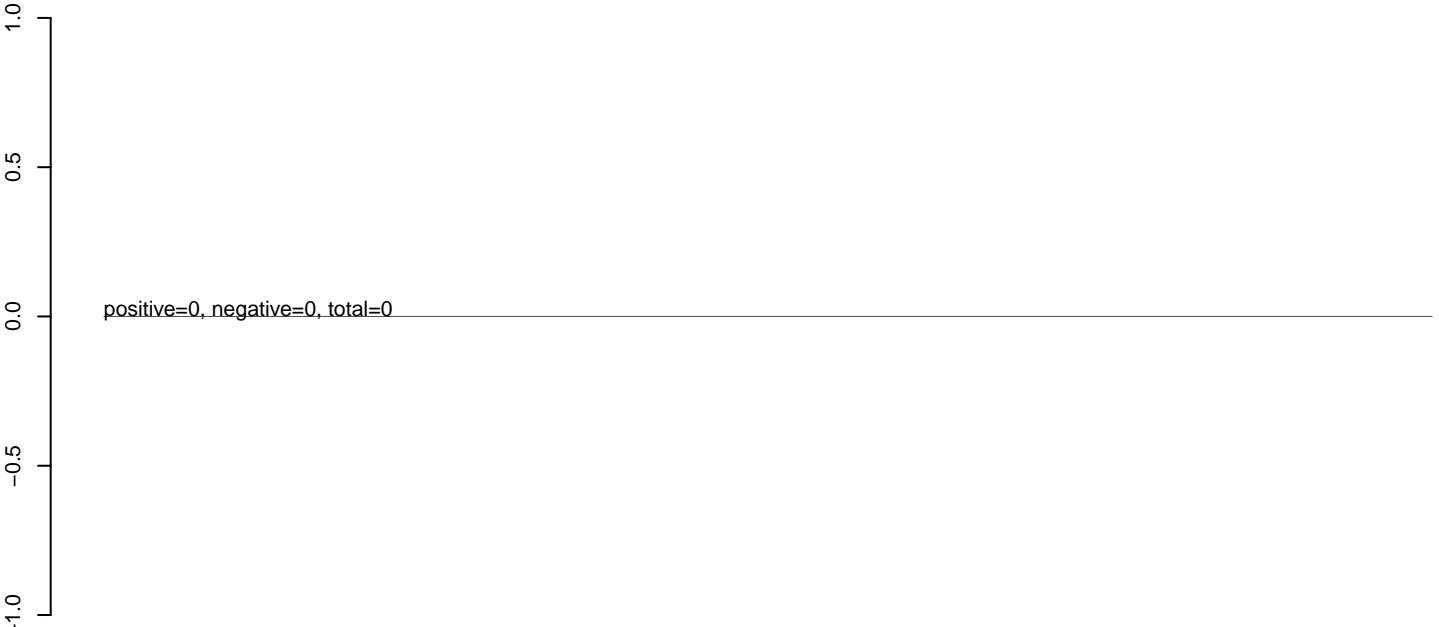
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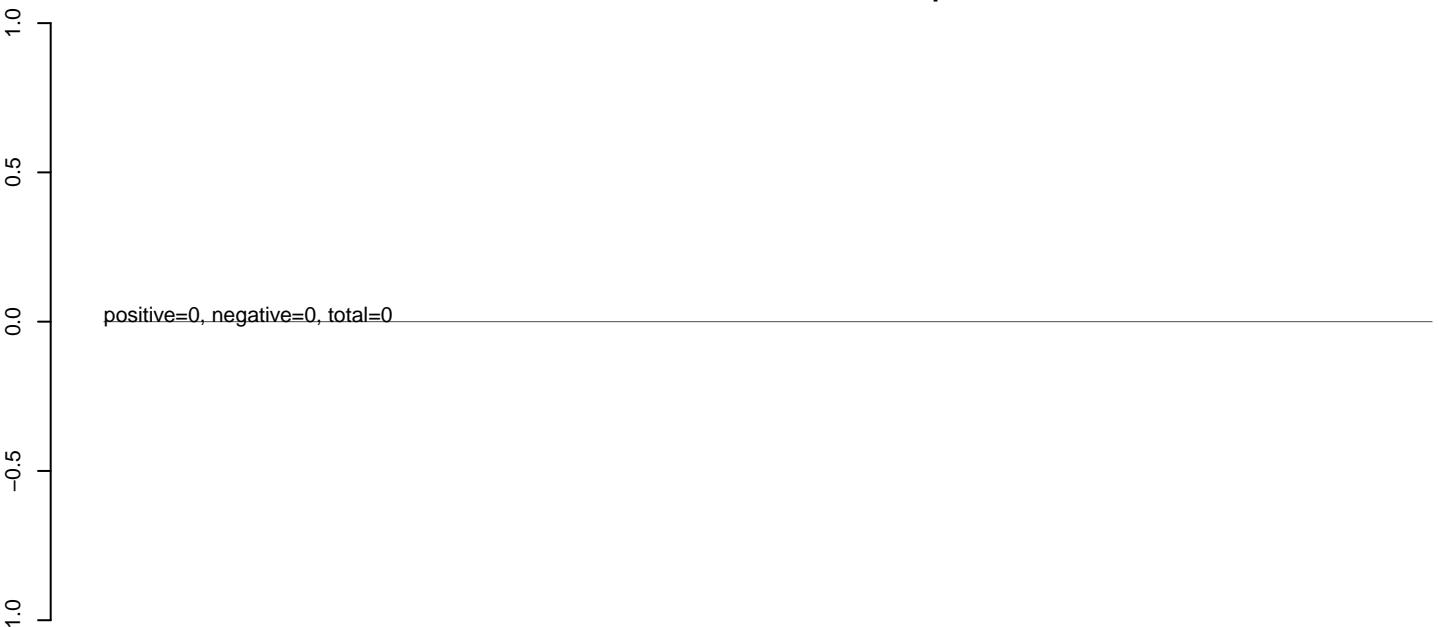
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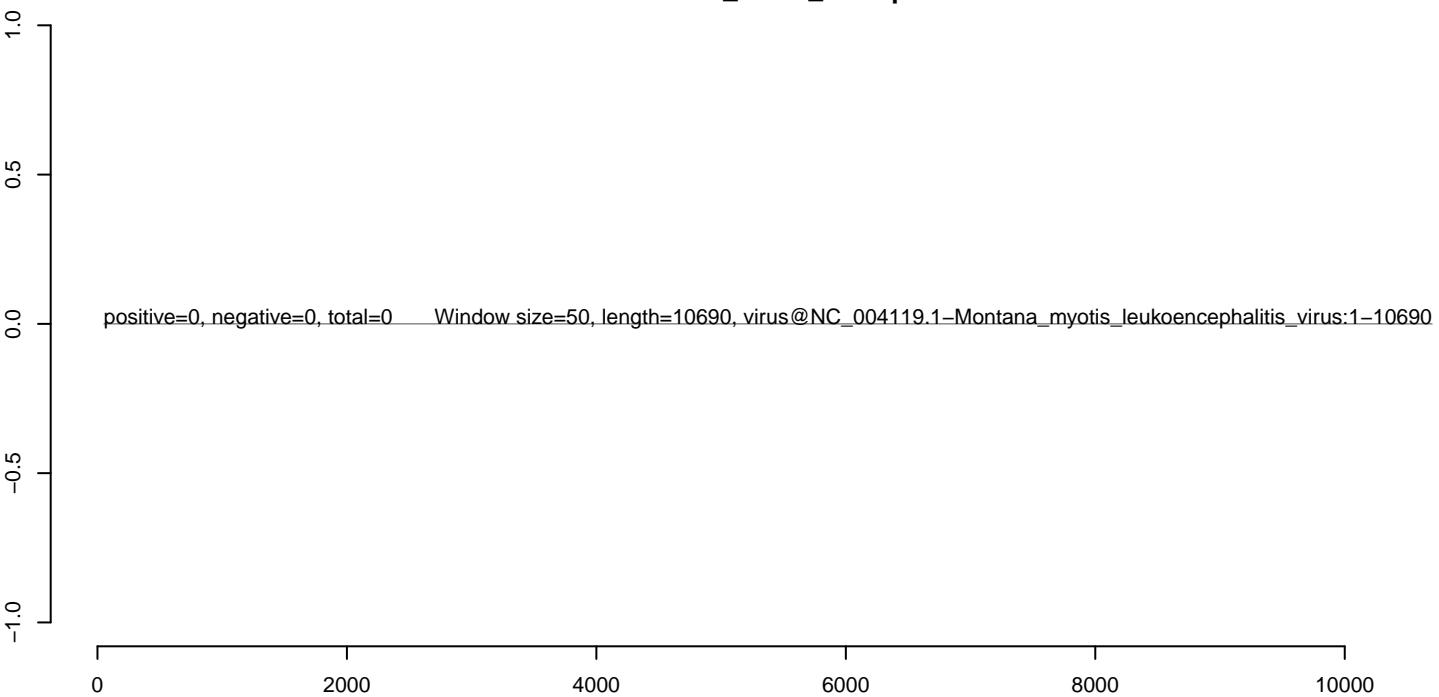
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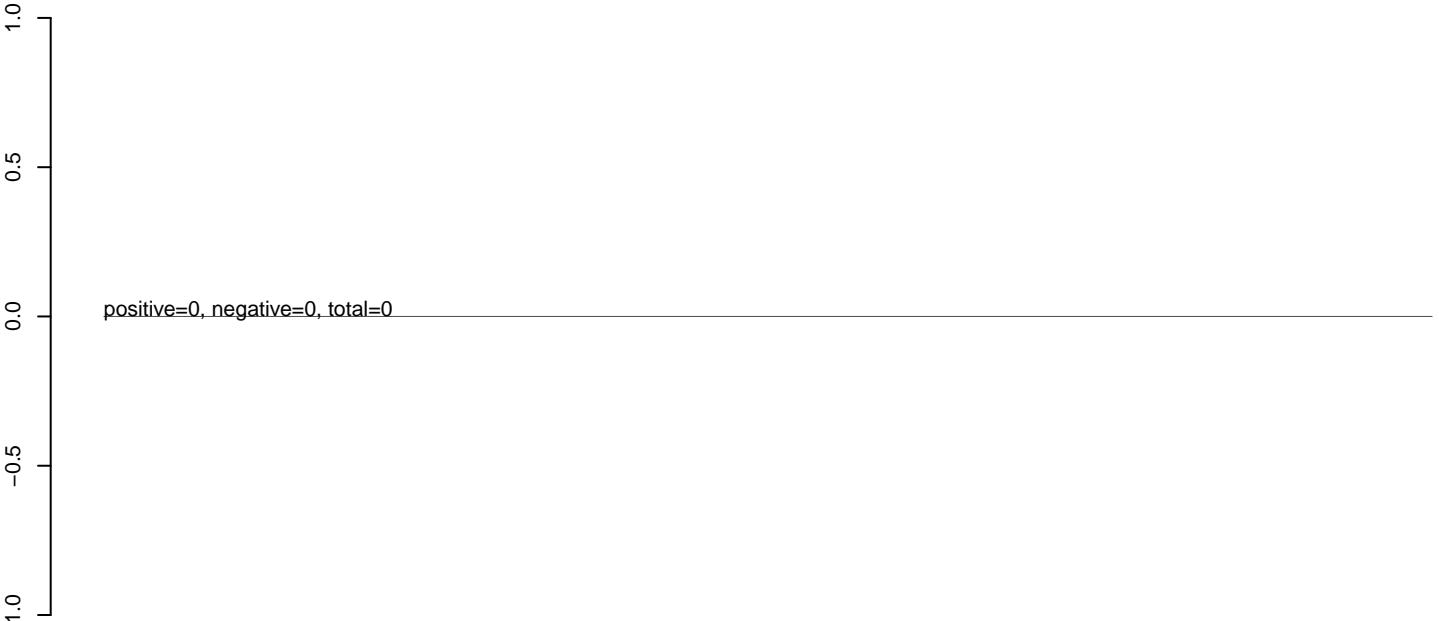
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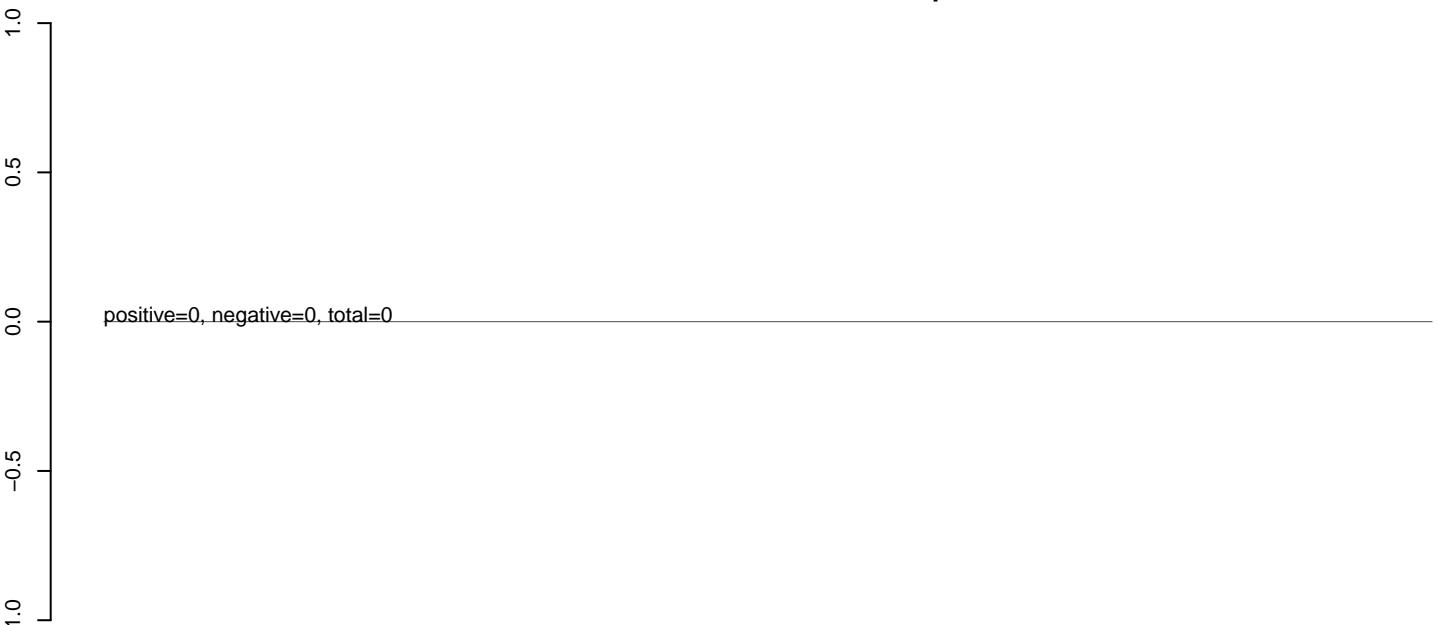
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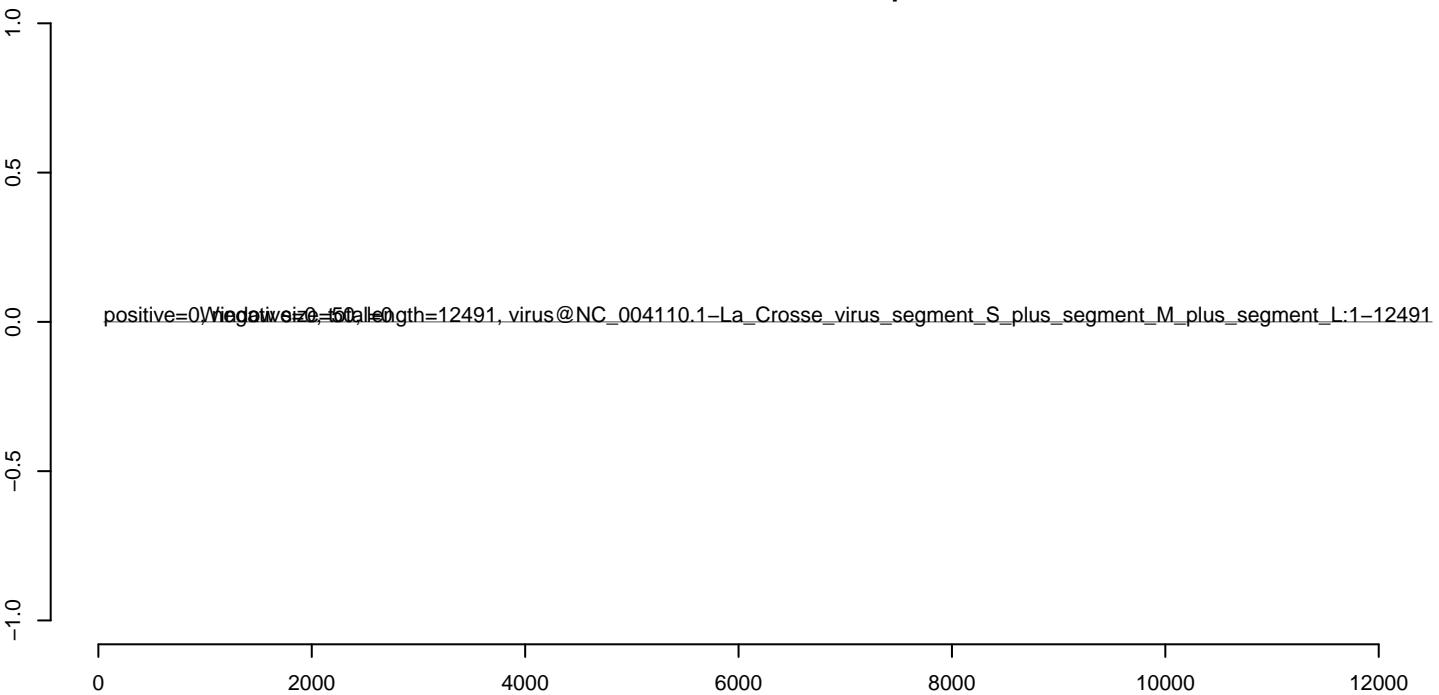
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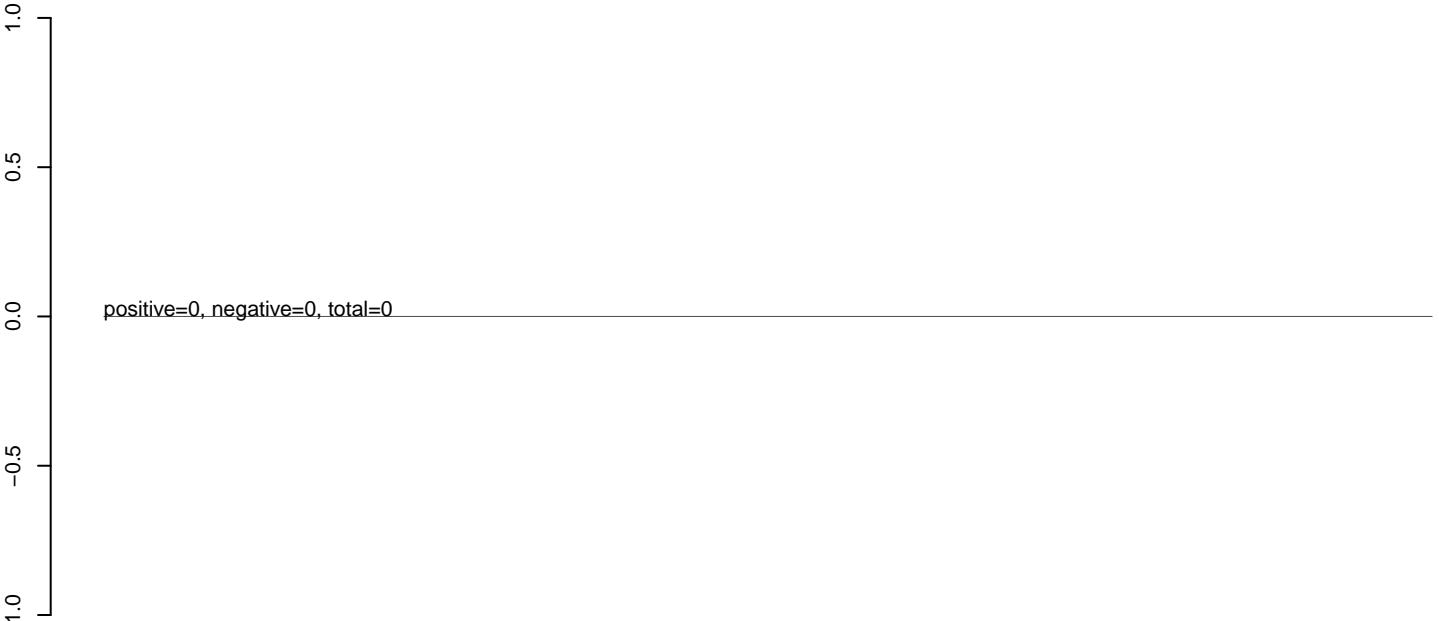
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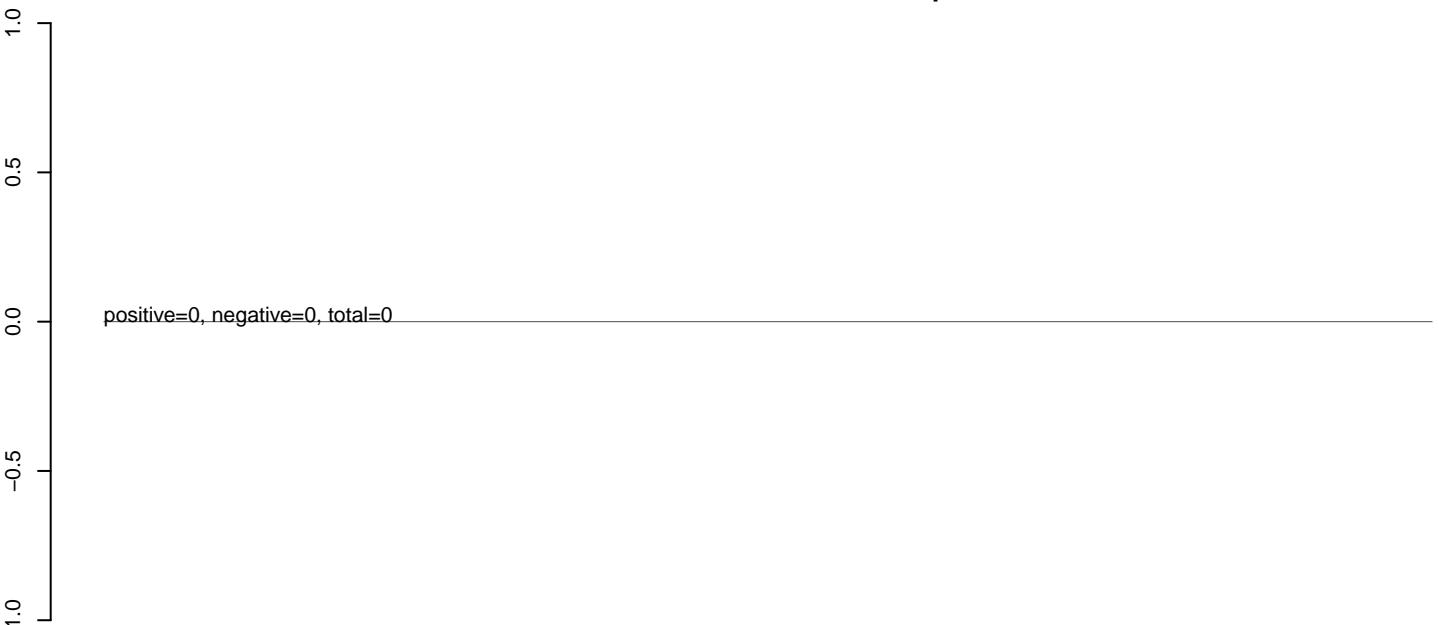
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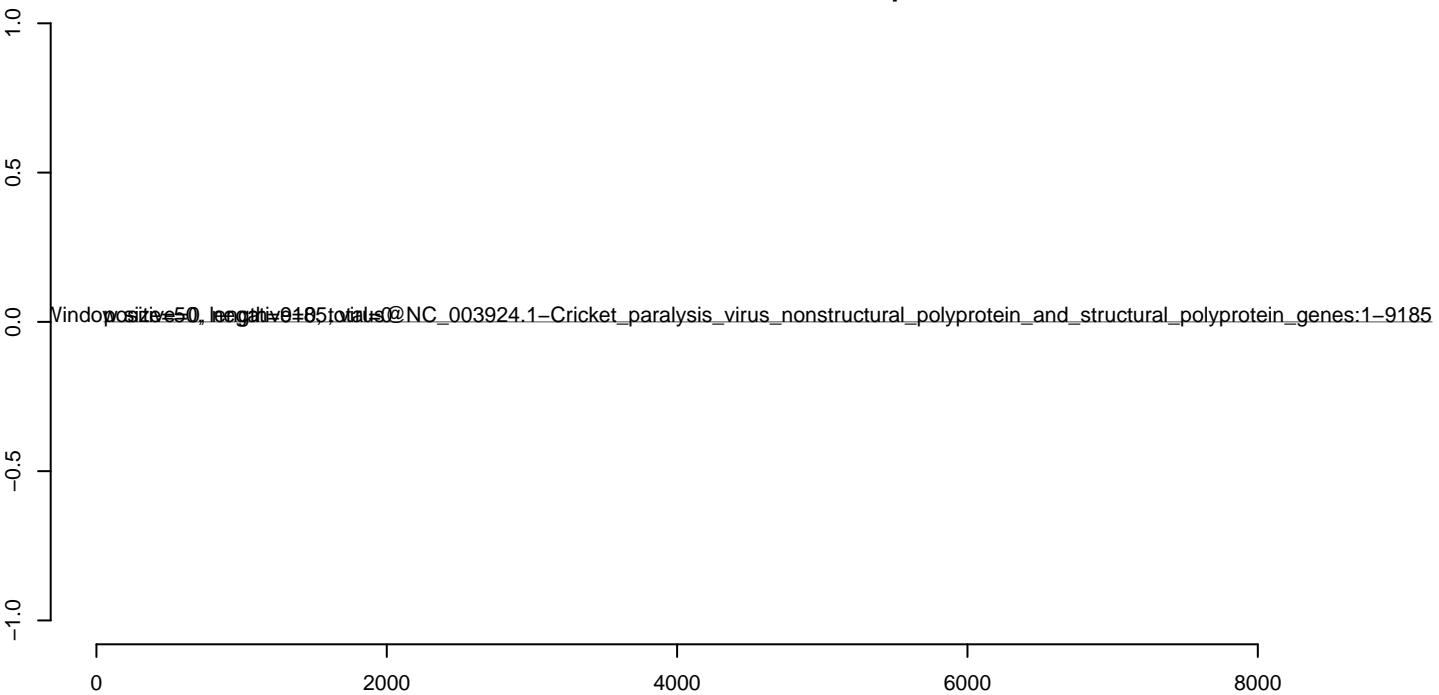
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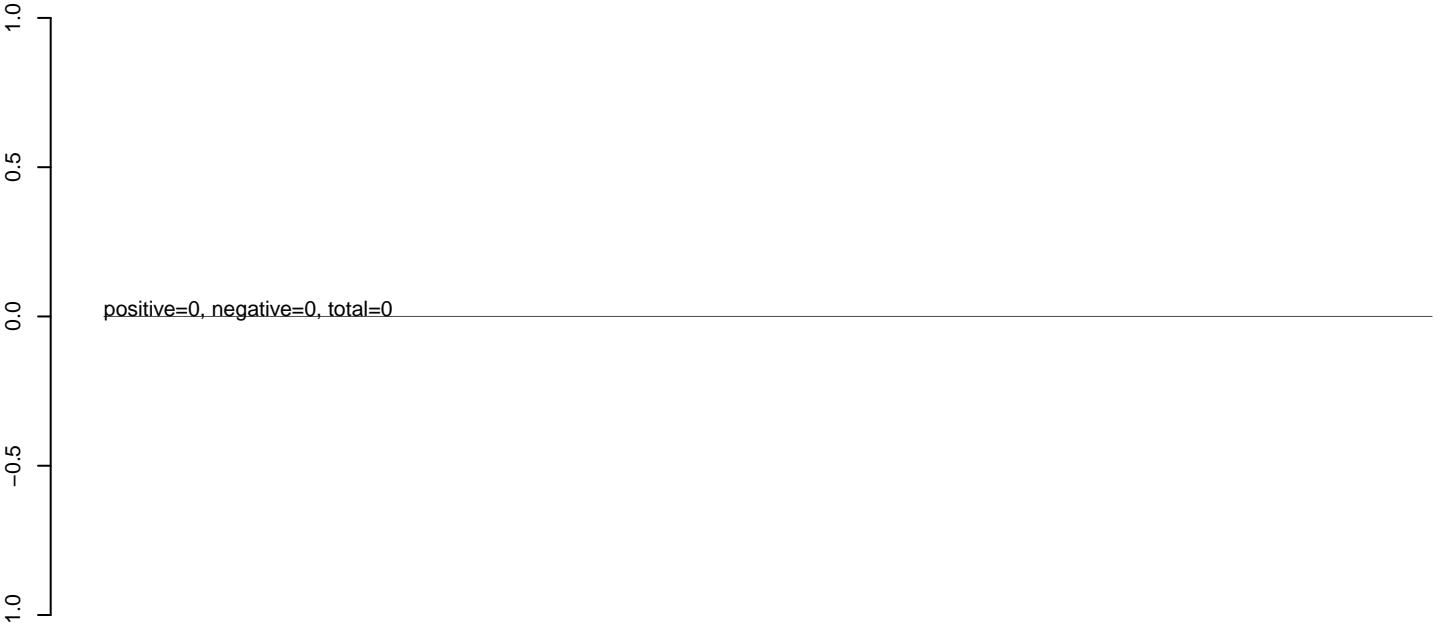
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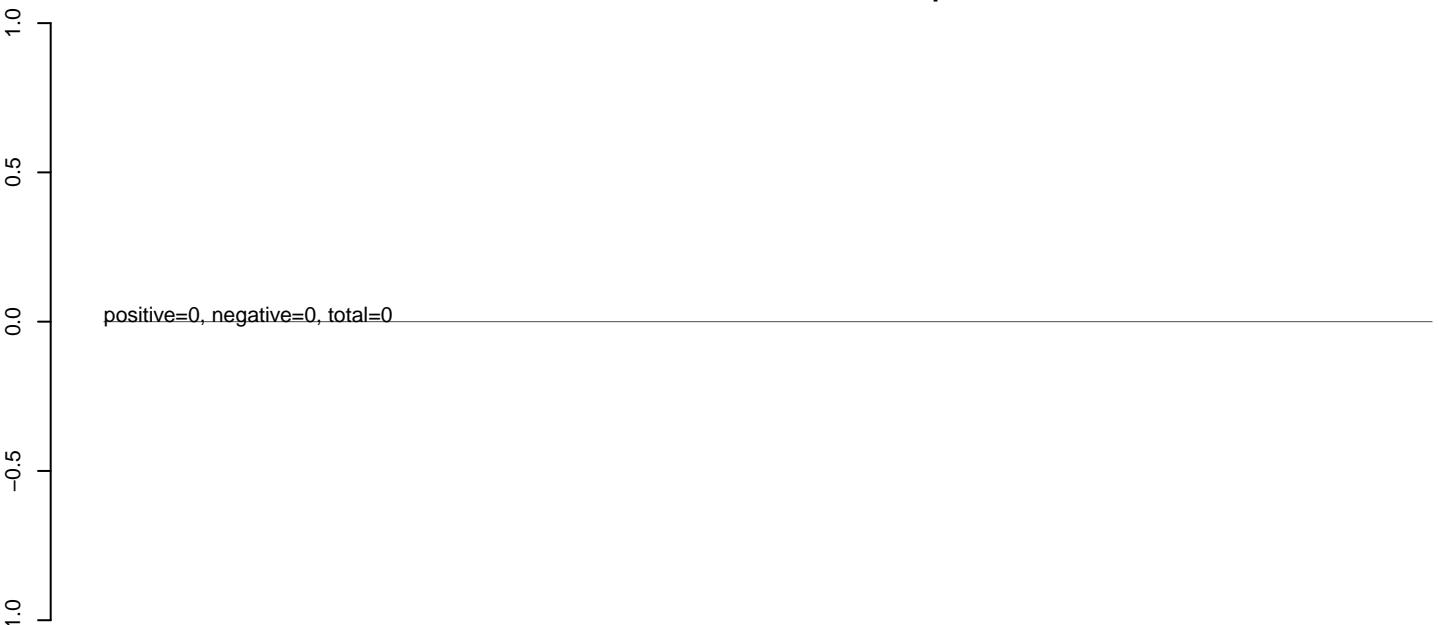
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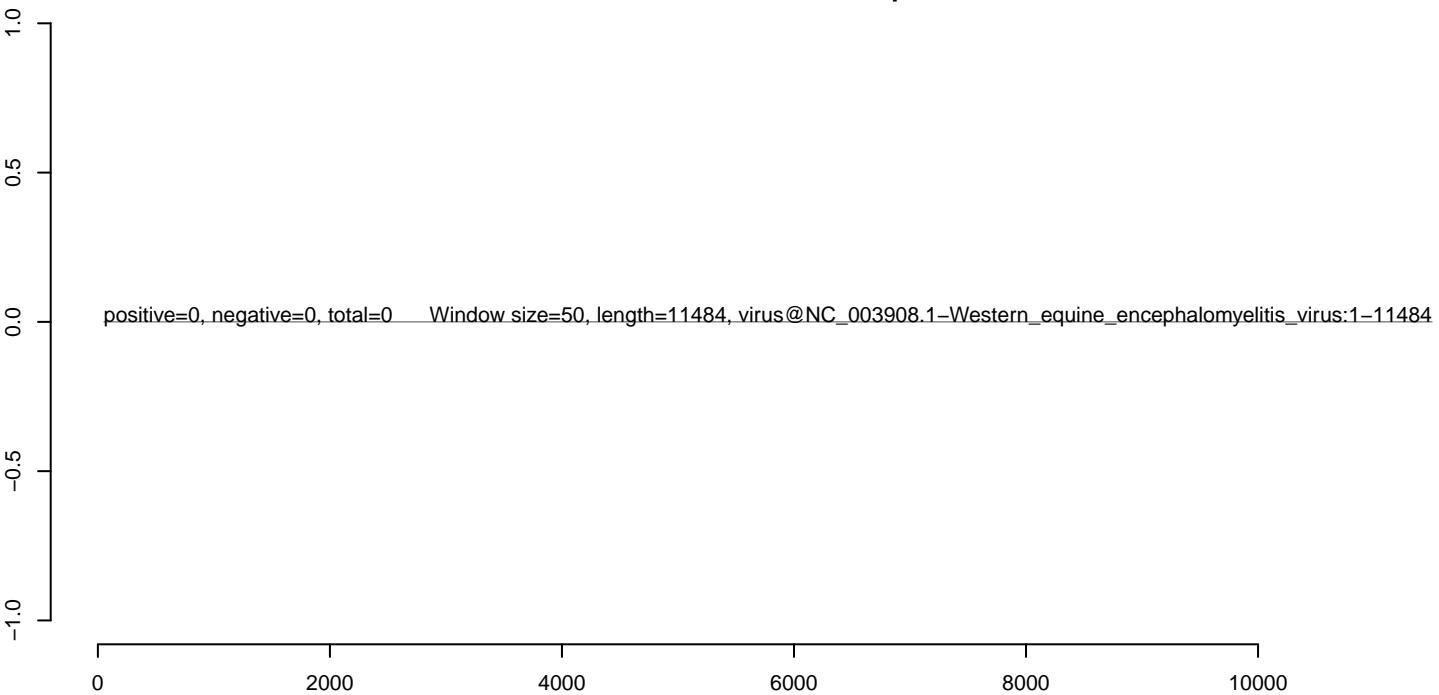
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### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0

Window size=50, length=11824, virus@NC\_003900.1-Aura\_virus:1-11824

0 2000 4000 6000 8000 10000 12000

### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

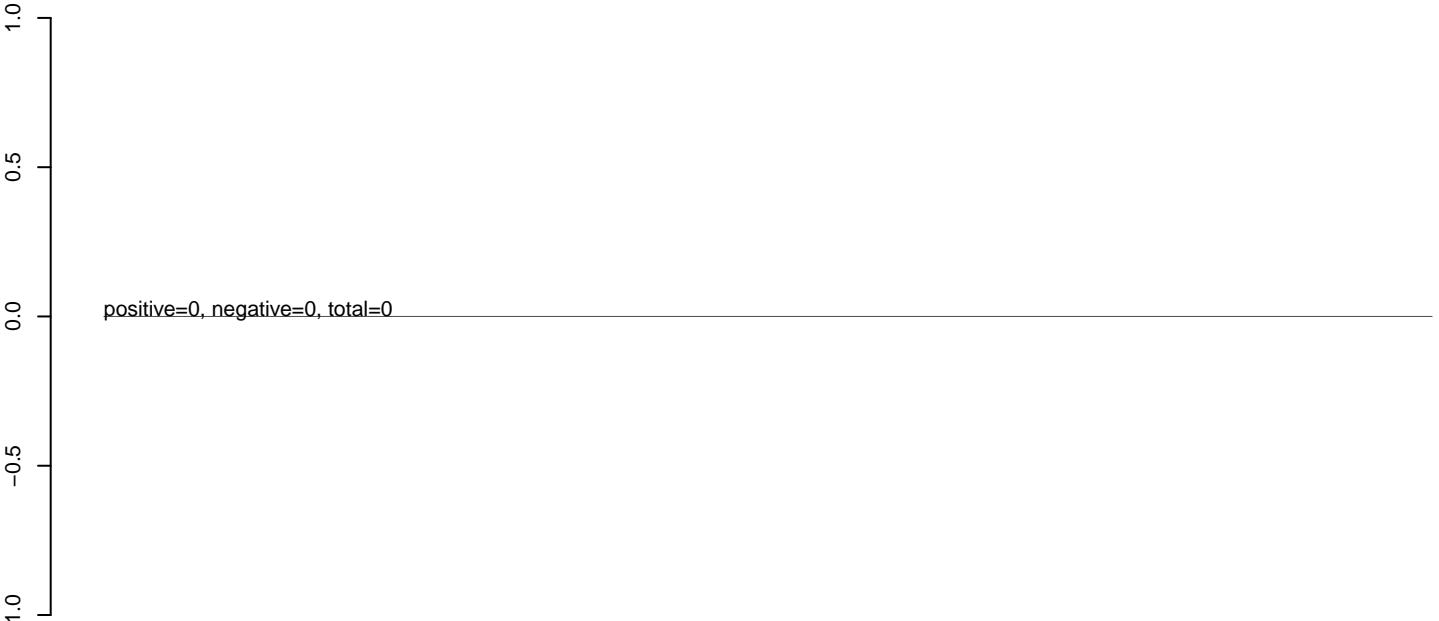
### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0

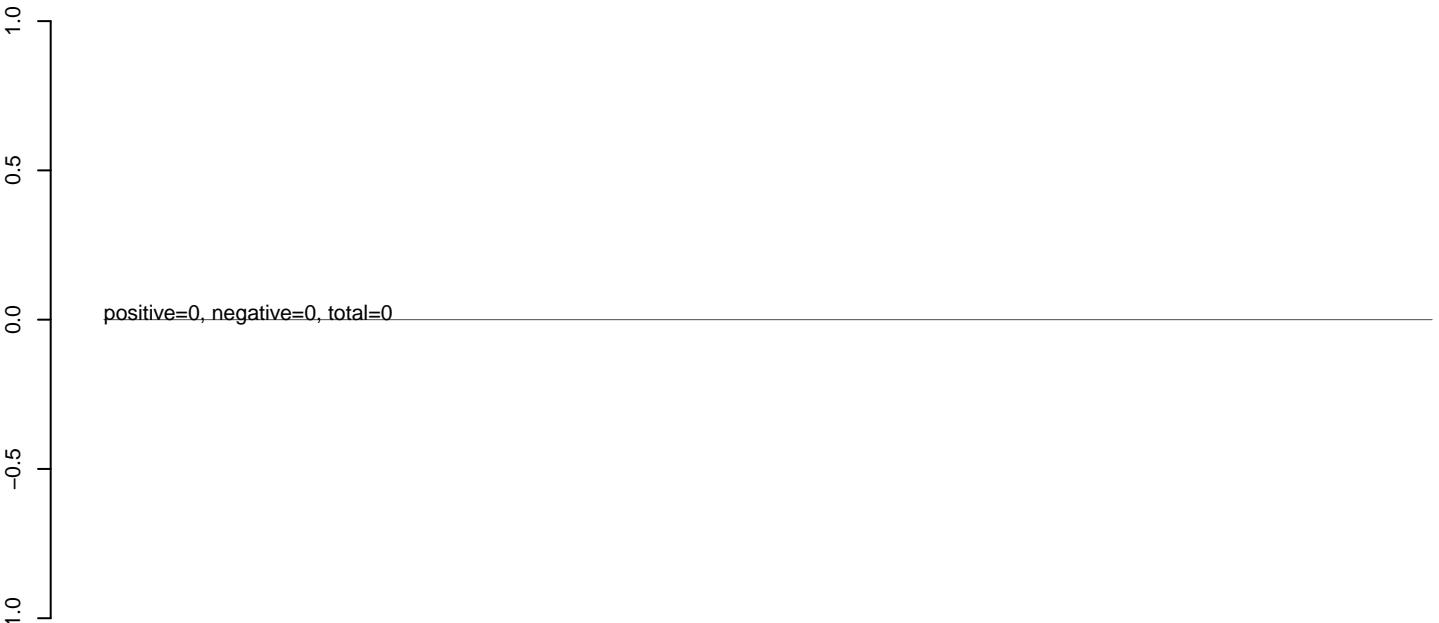
Window size=50, length=11675, virus@NC\_003899.1-Eastern\_equine\_encephalitis\_virus:1-11675

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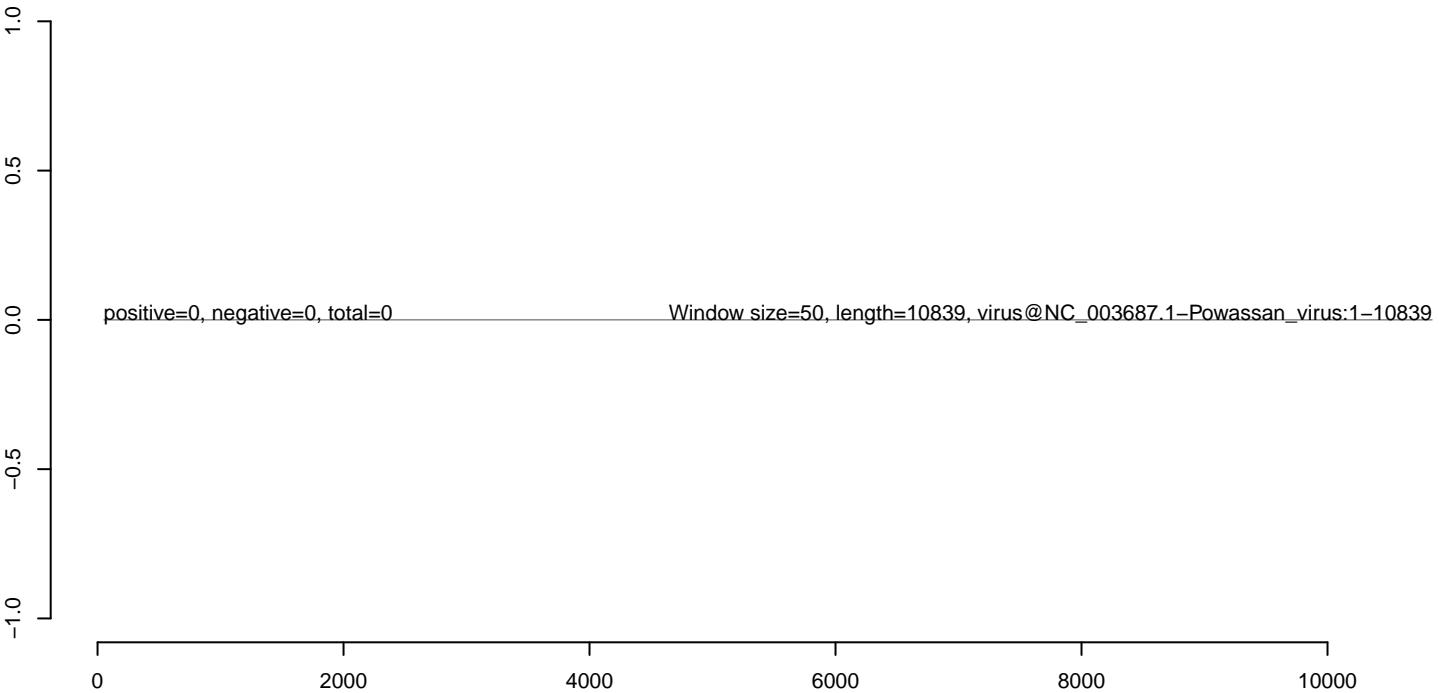
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0

Window size=50, length=10116, virus@NC\_003676.1-Apoi\_virus\_genome:1-10116

0 2000 4000 6000 8000 10000

### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

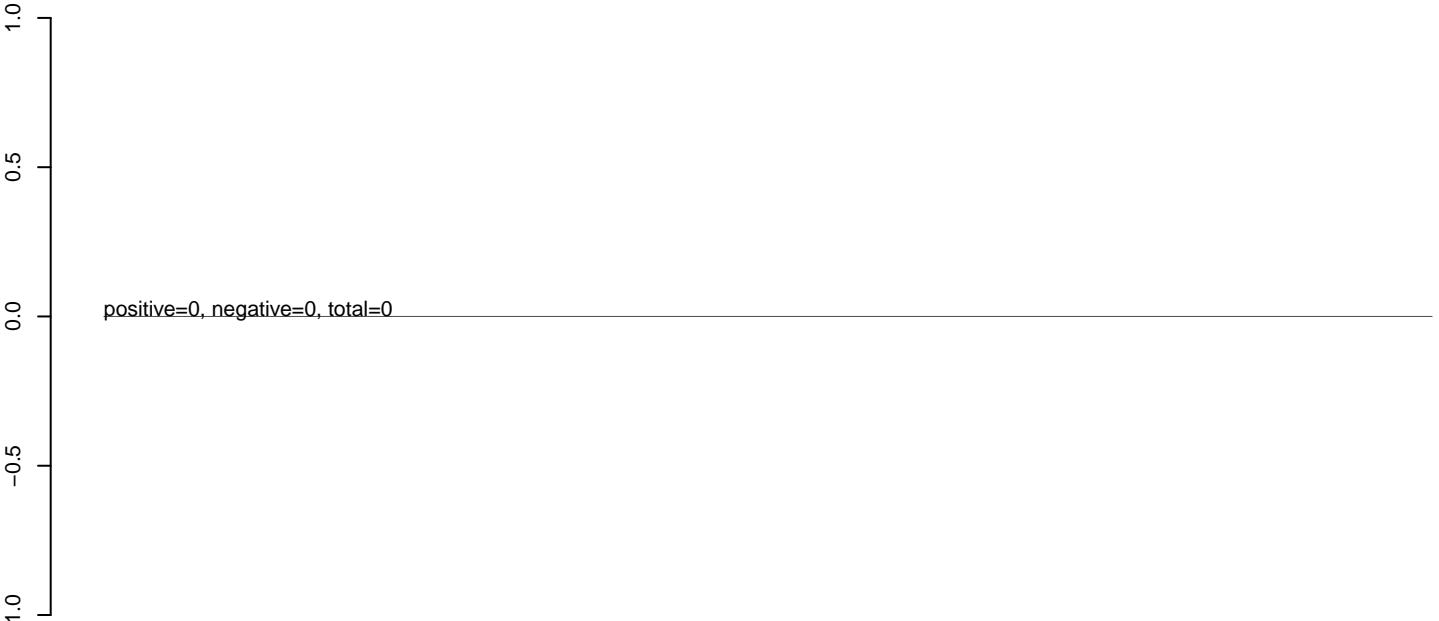
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### AnGam\_Fcanc\_TN.rep

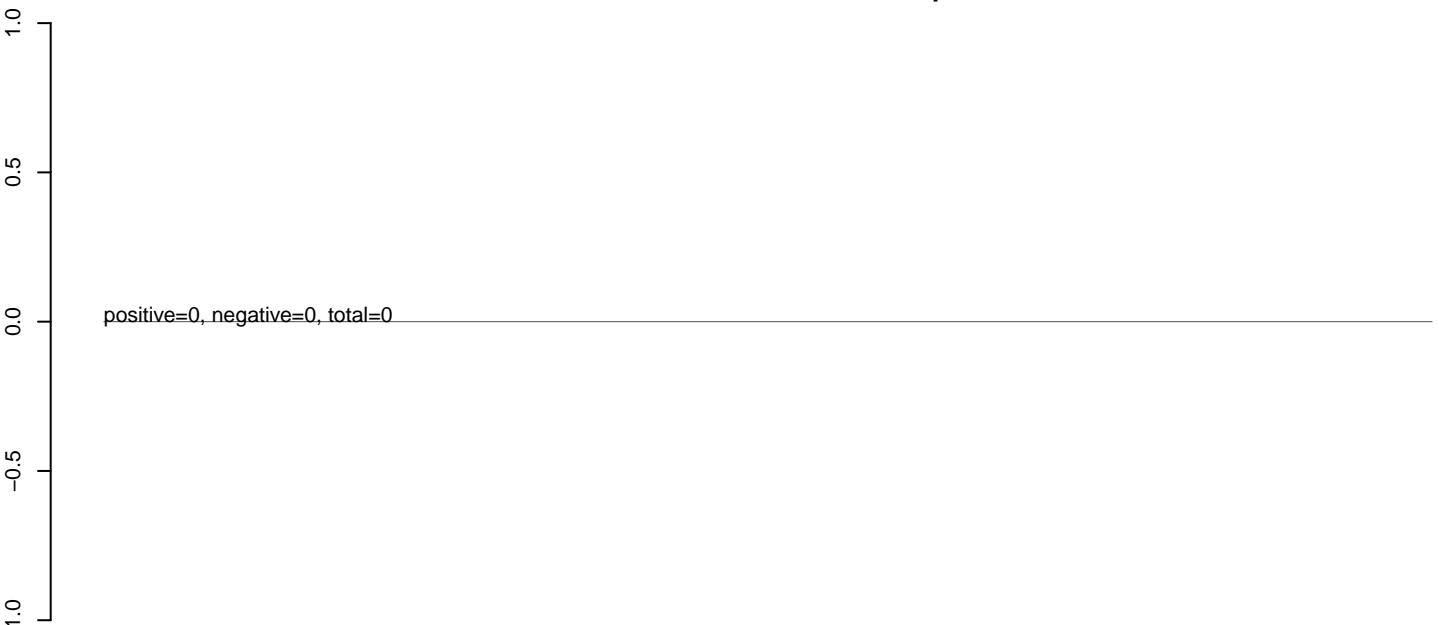
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Window size=50, length=10140, virus@NC\_003675.1–Rio\_Bravo\_virus\_genome:1–10140

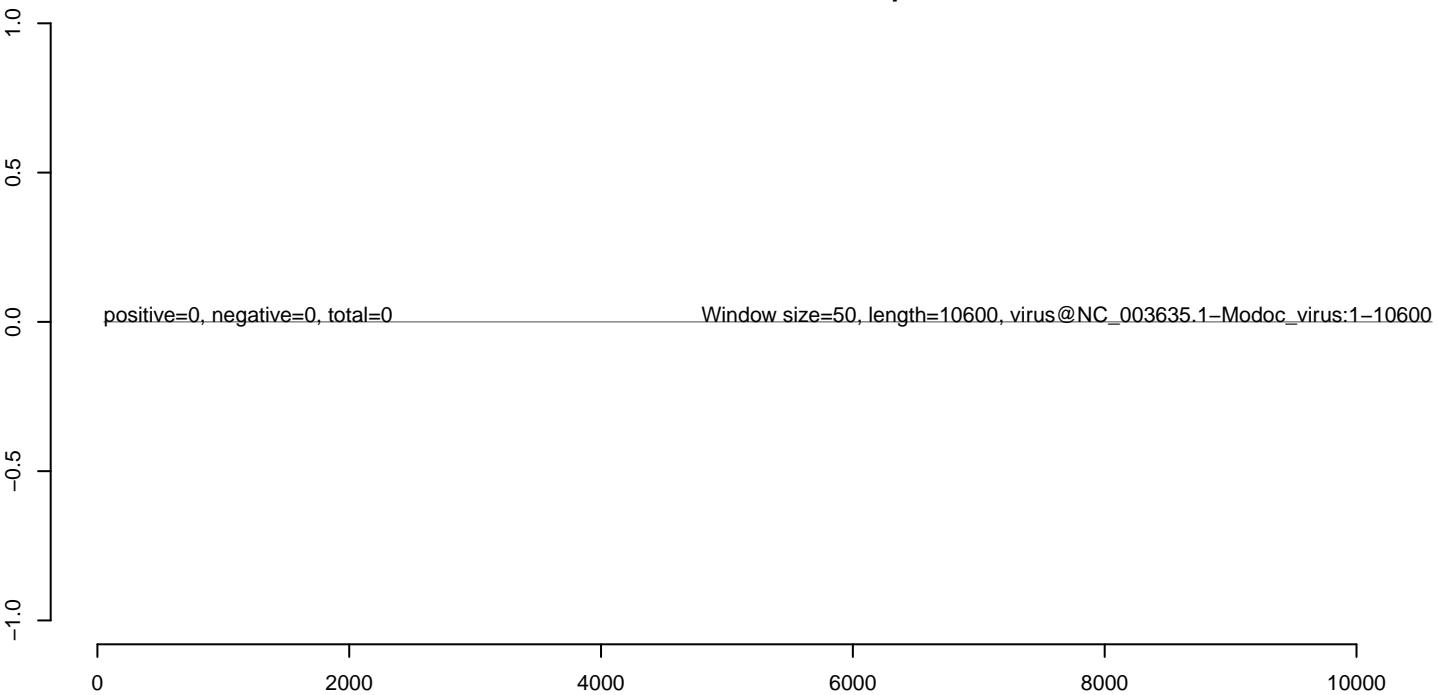
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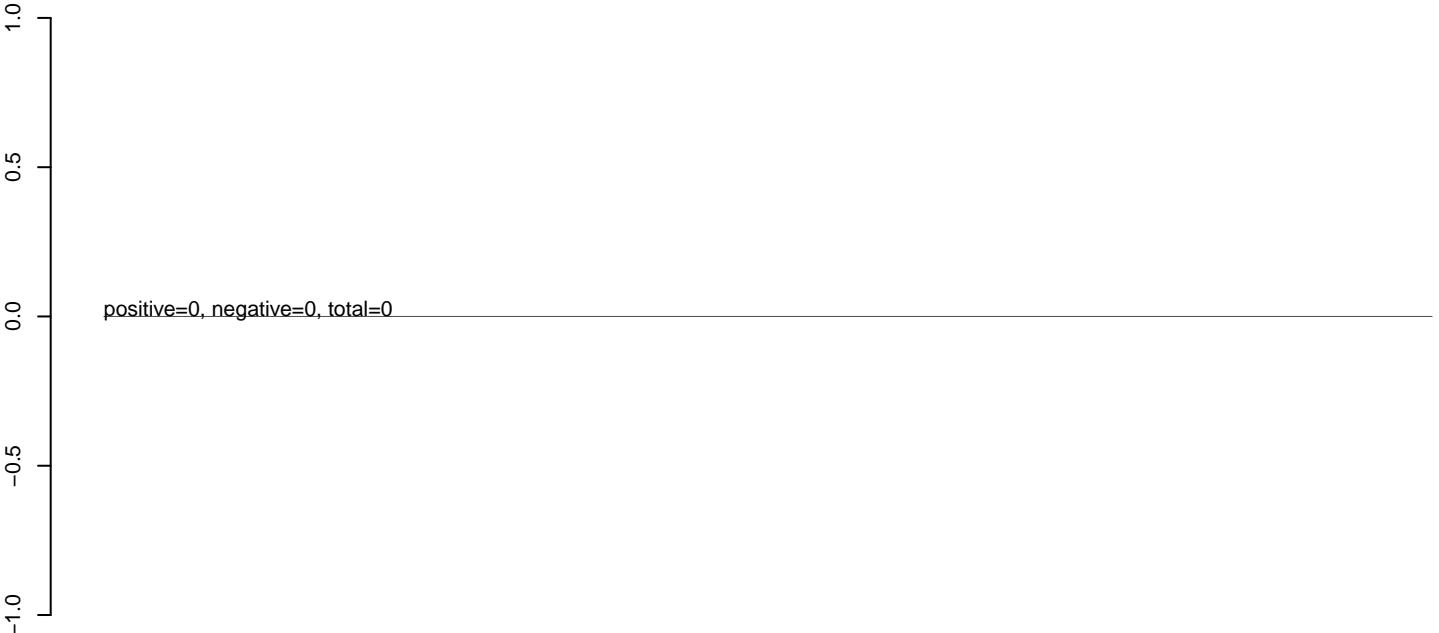
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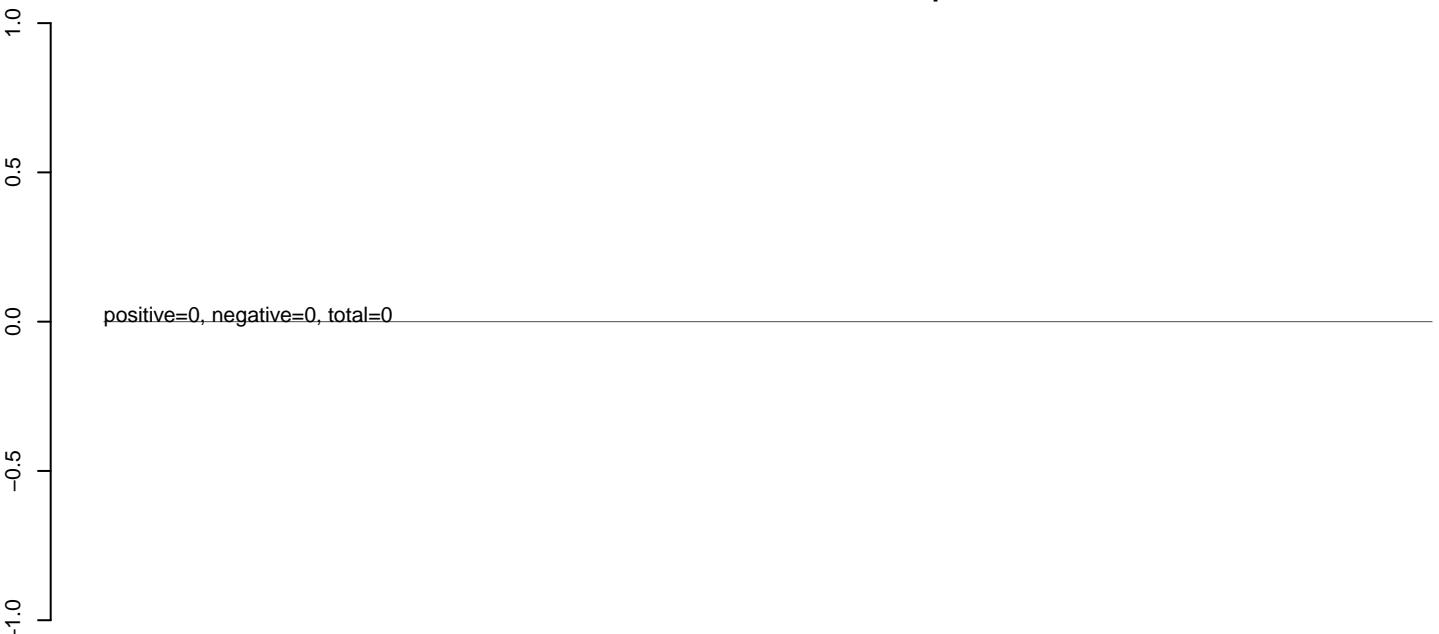
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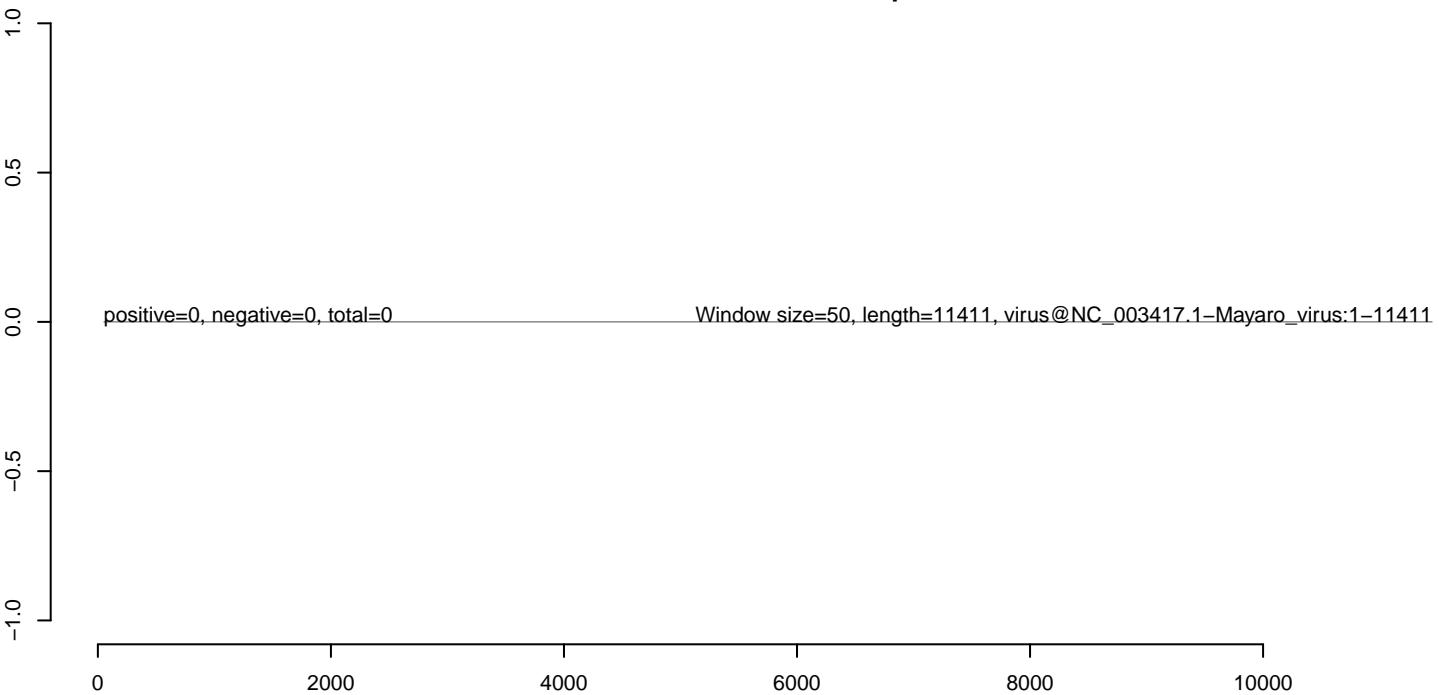
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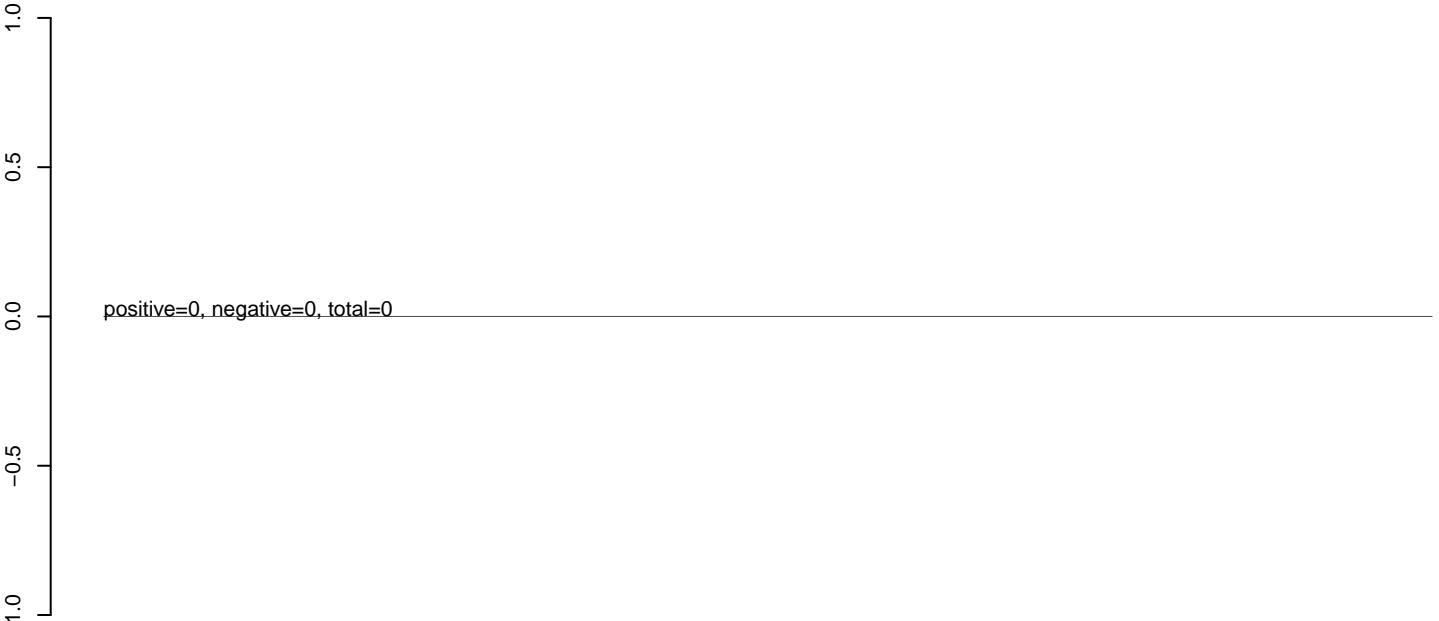
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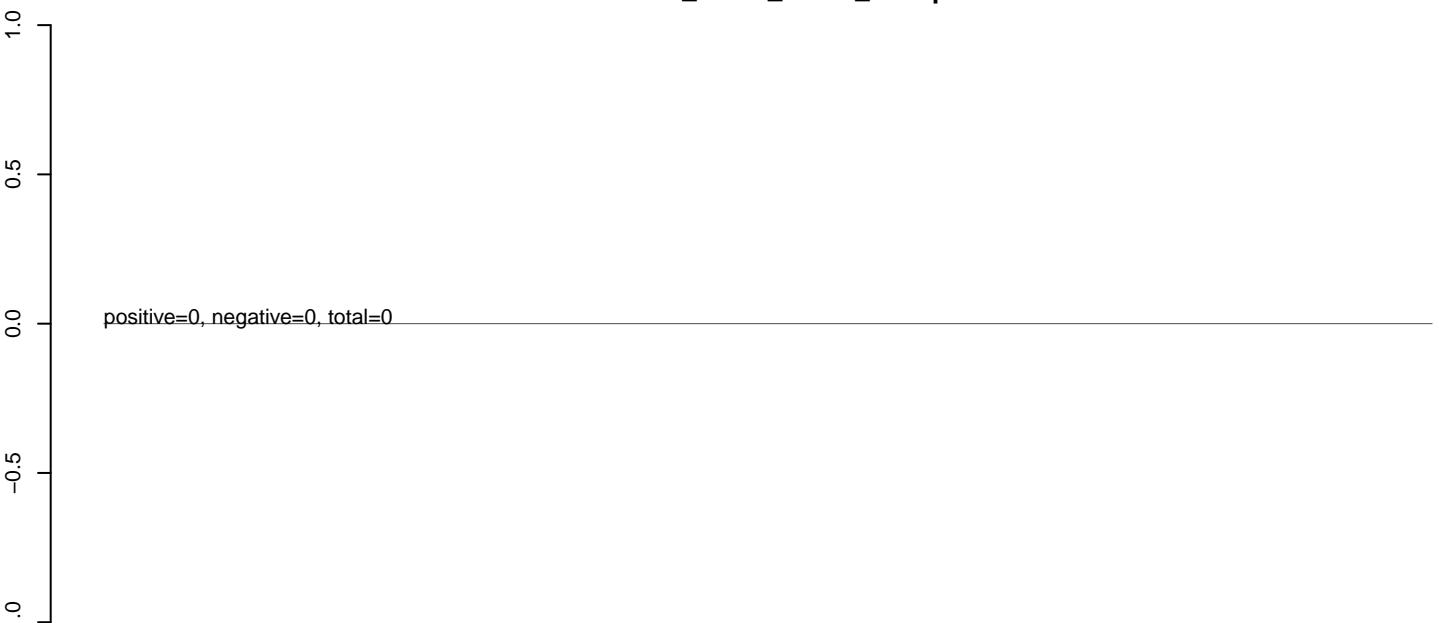
### AnGam\_Fcanc\_TN.rep



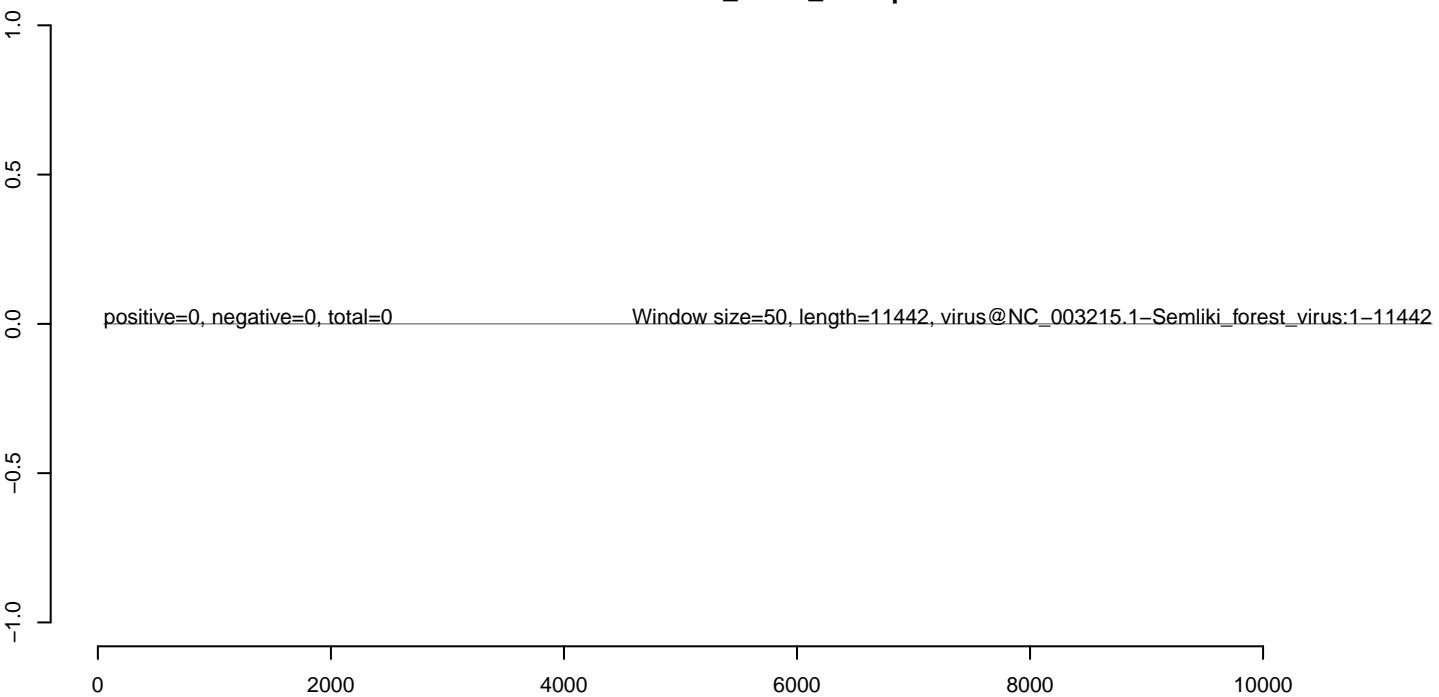
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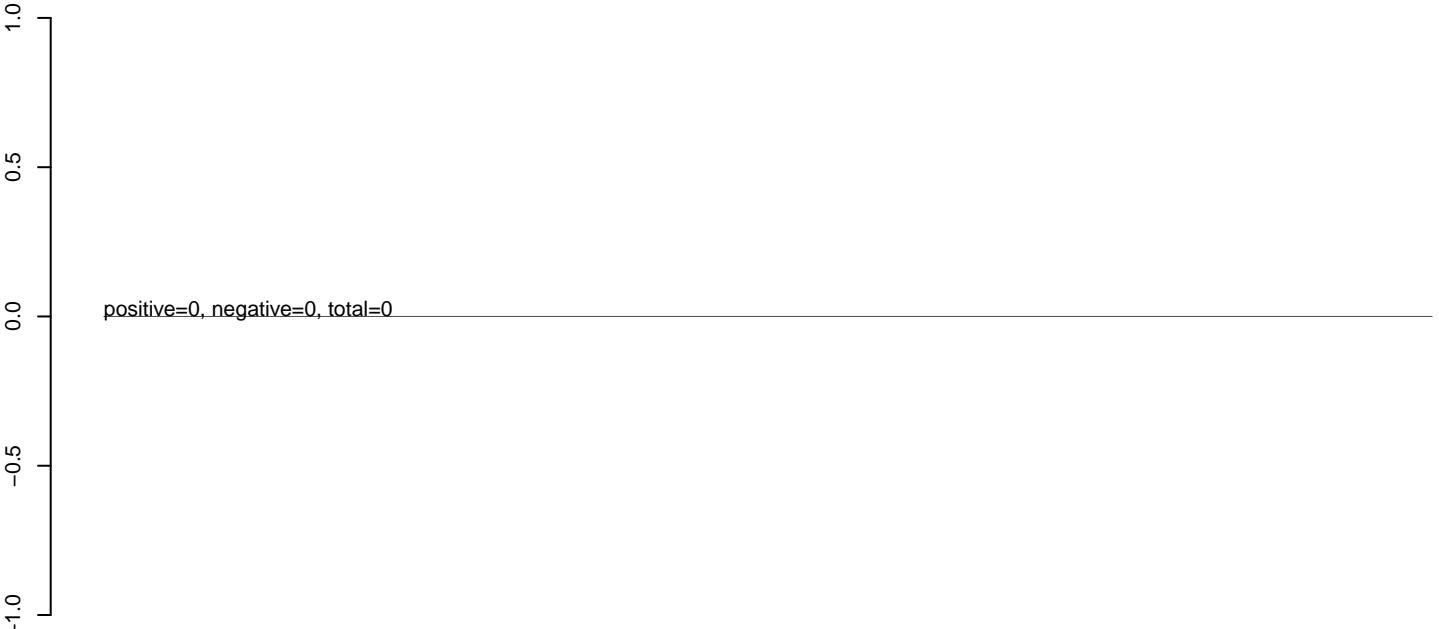
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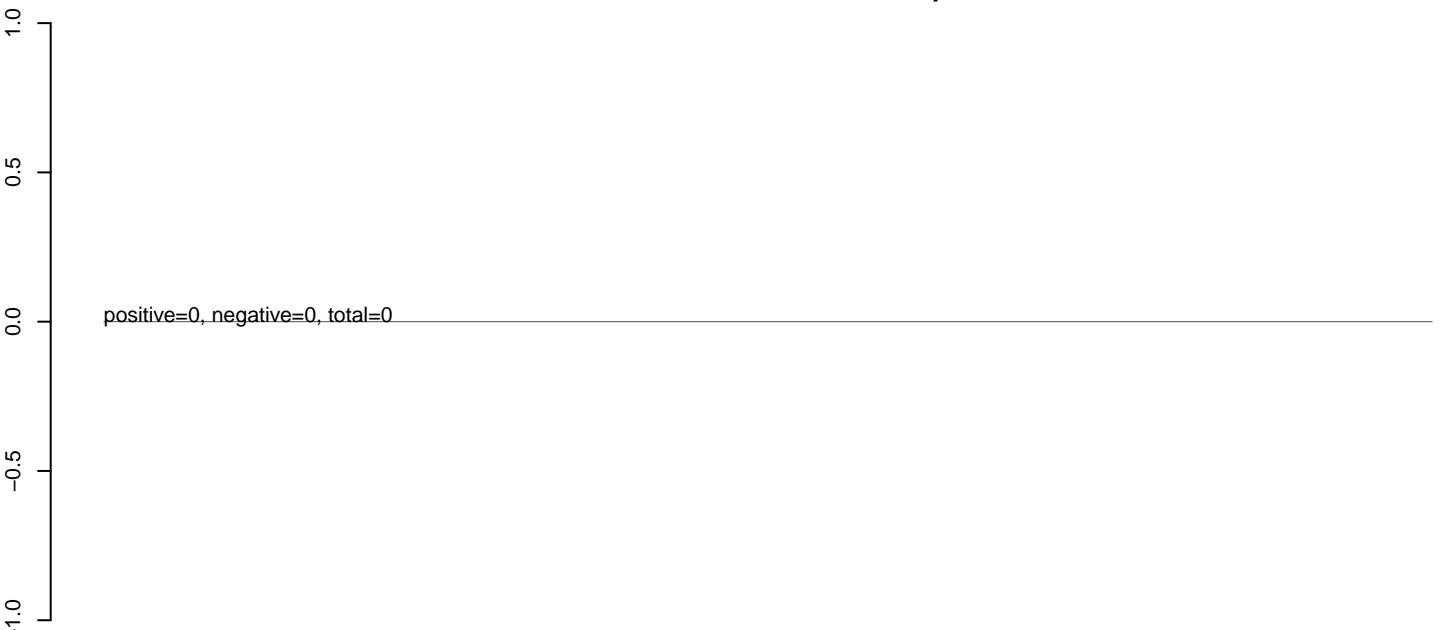
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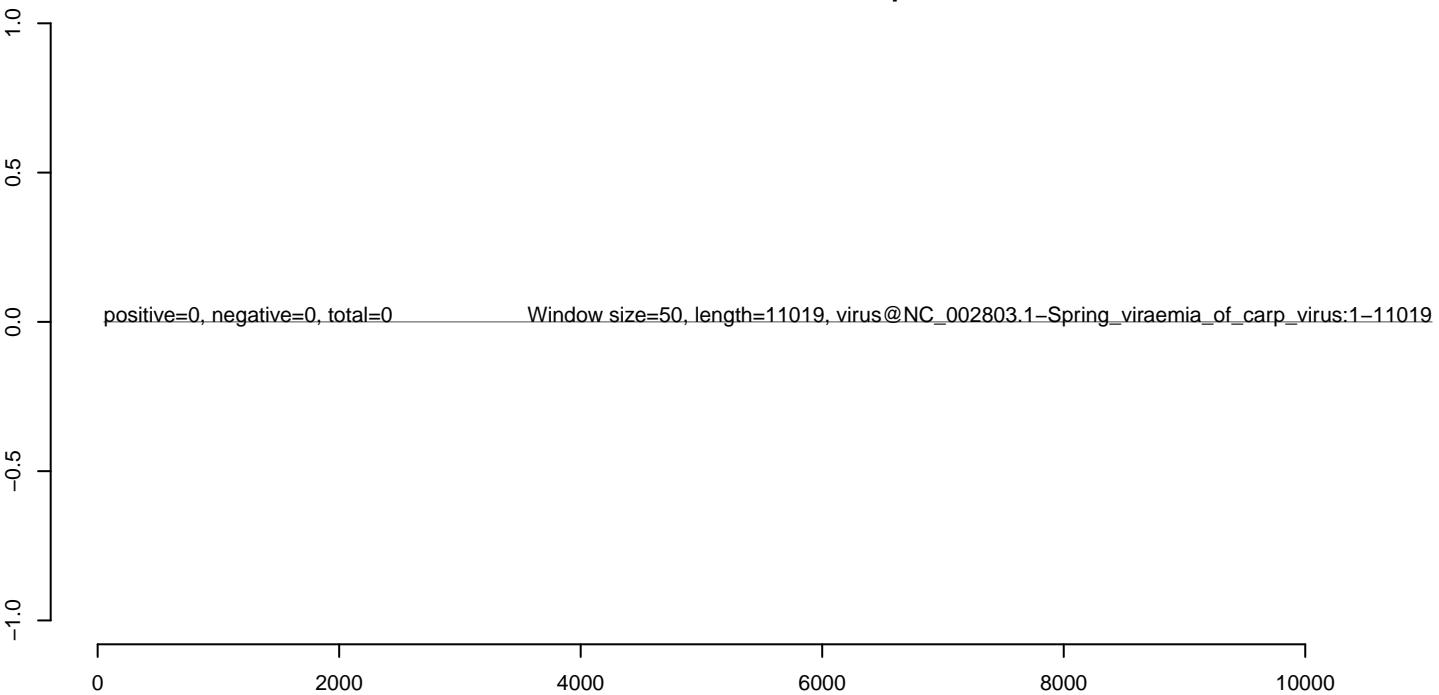
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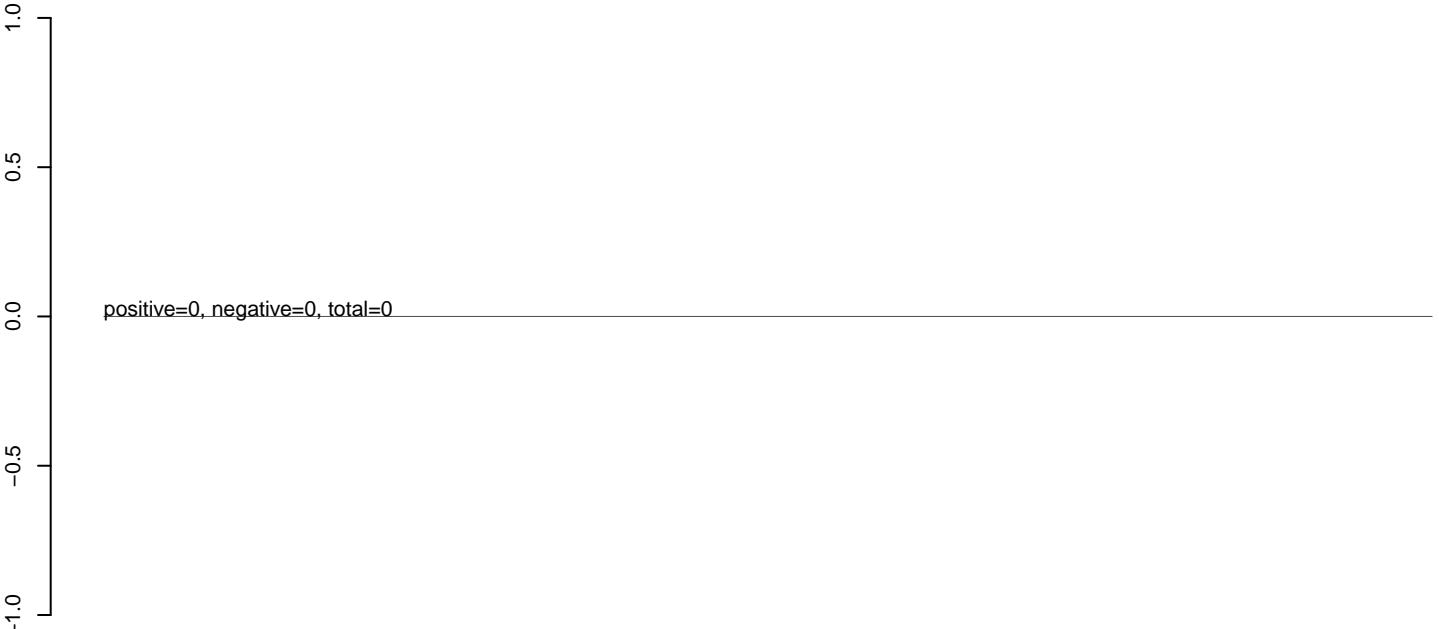
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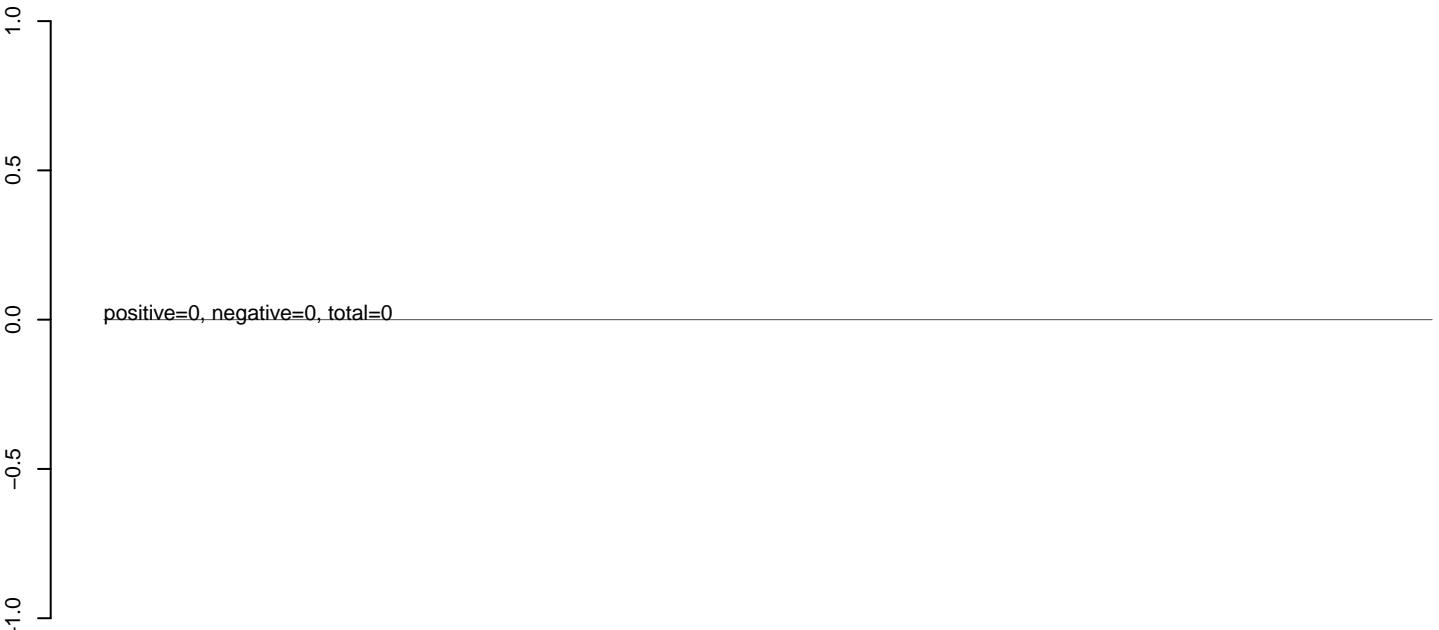
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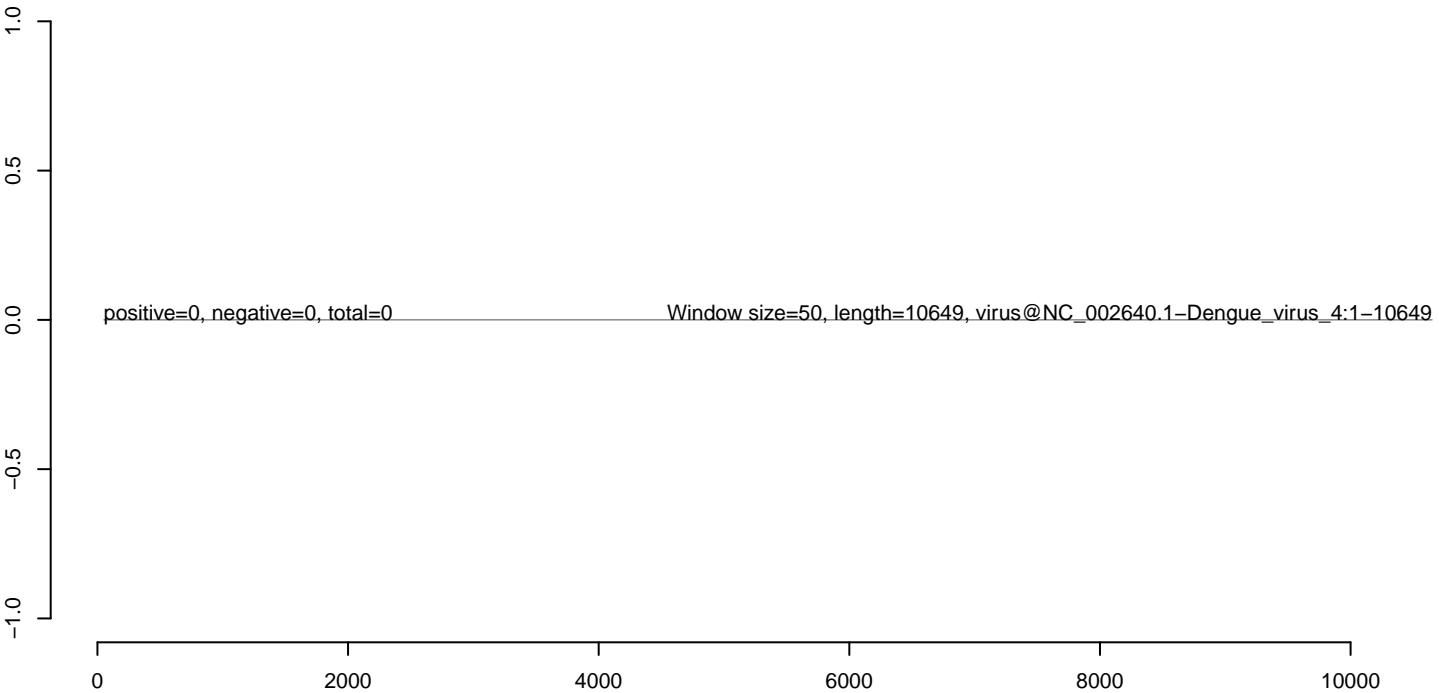
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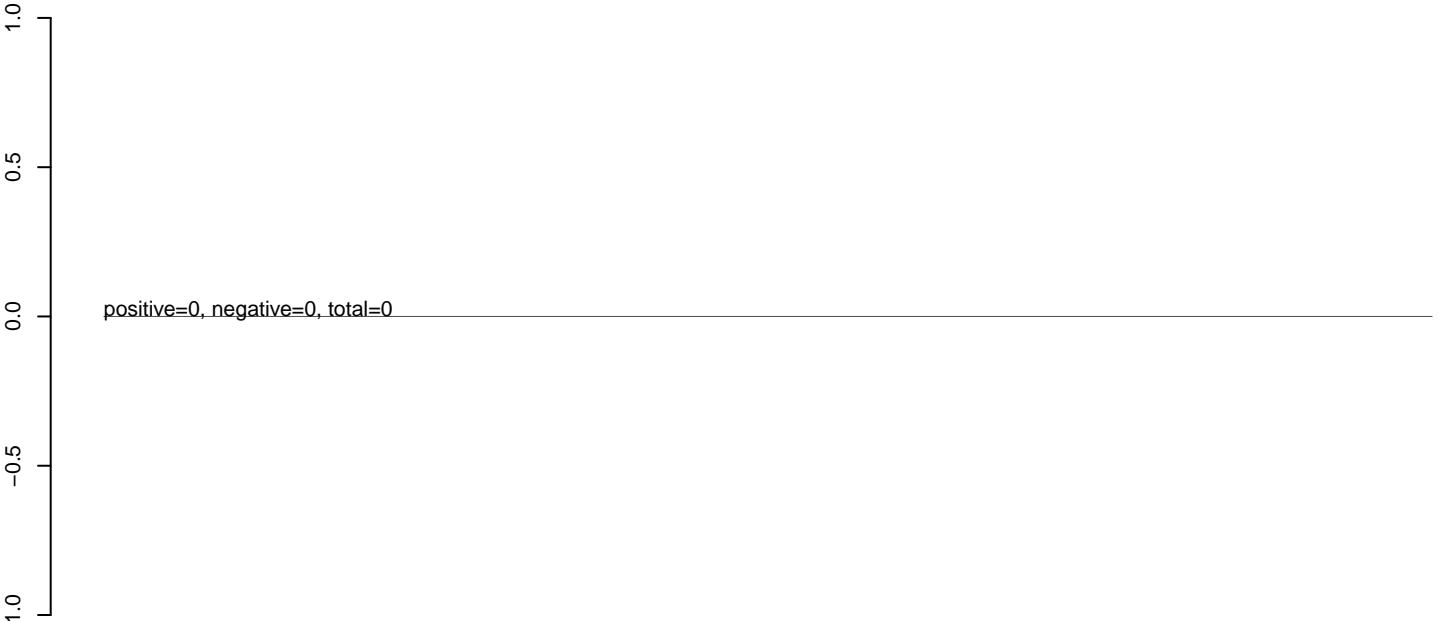
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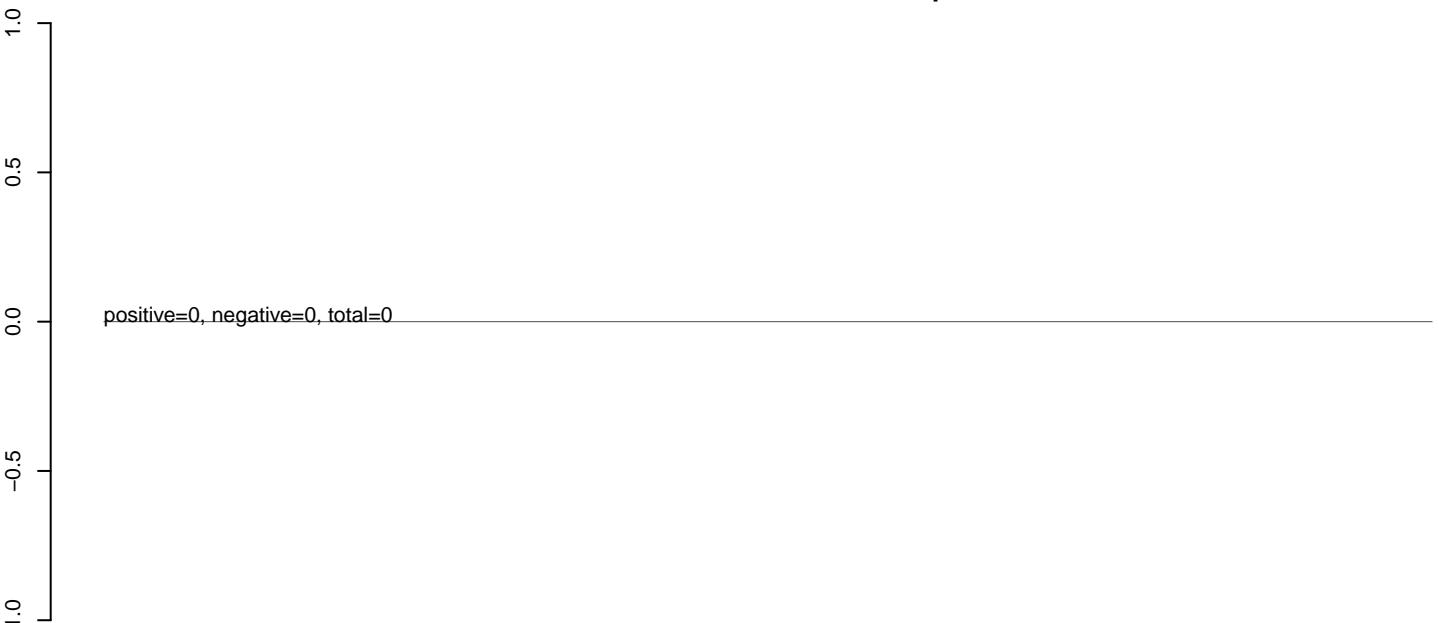
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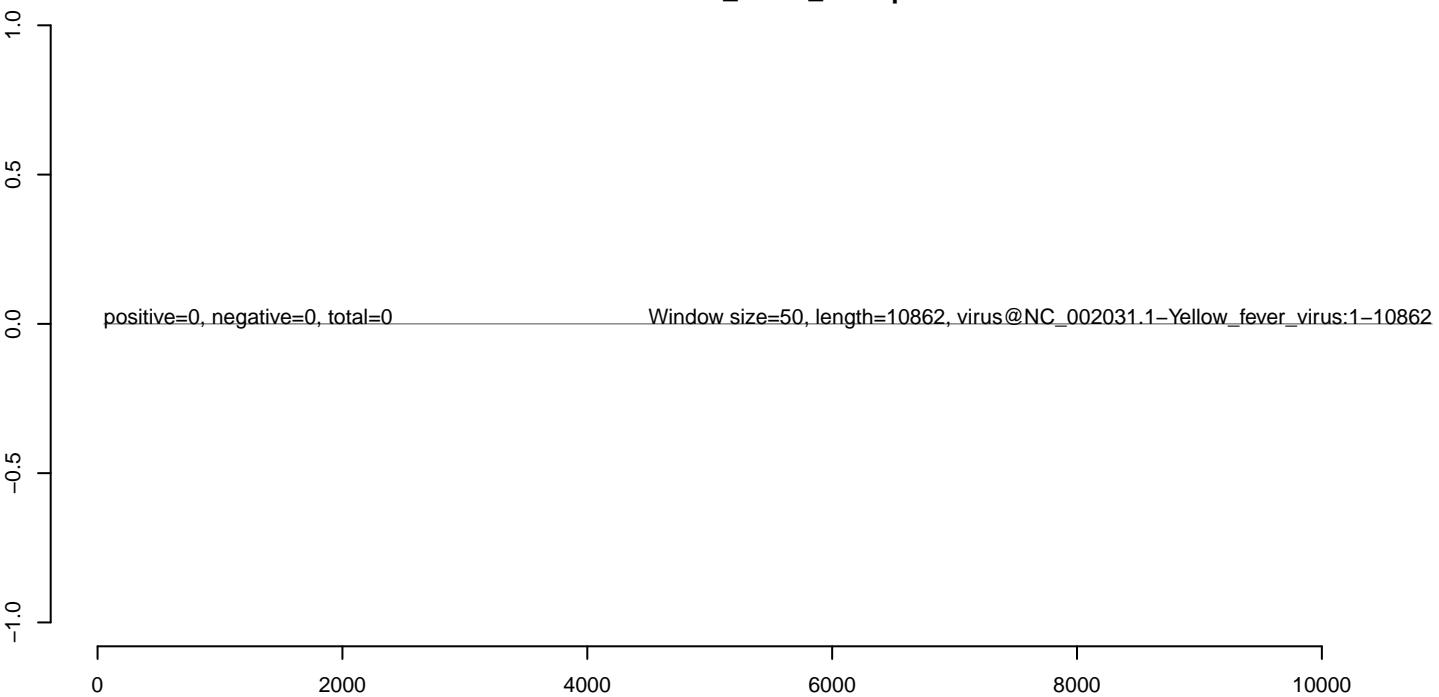
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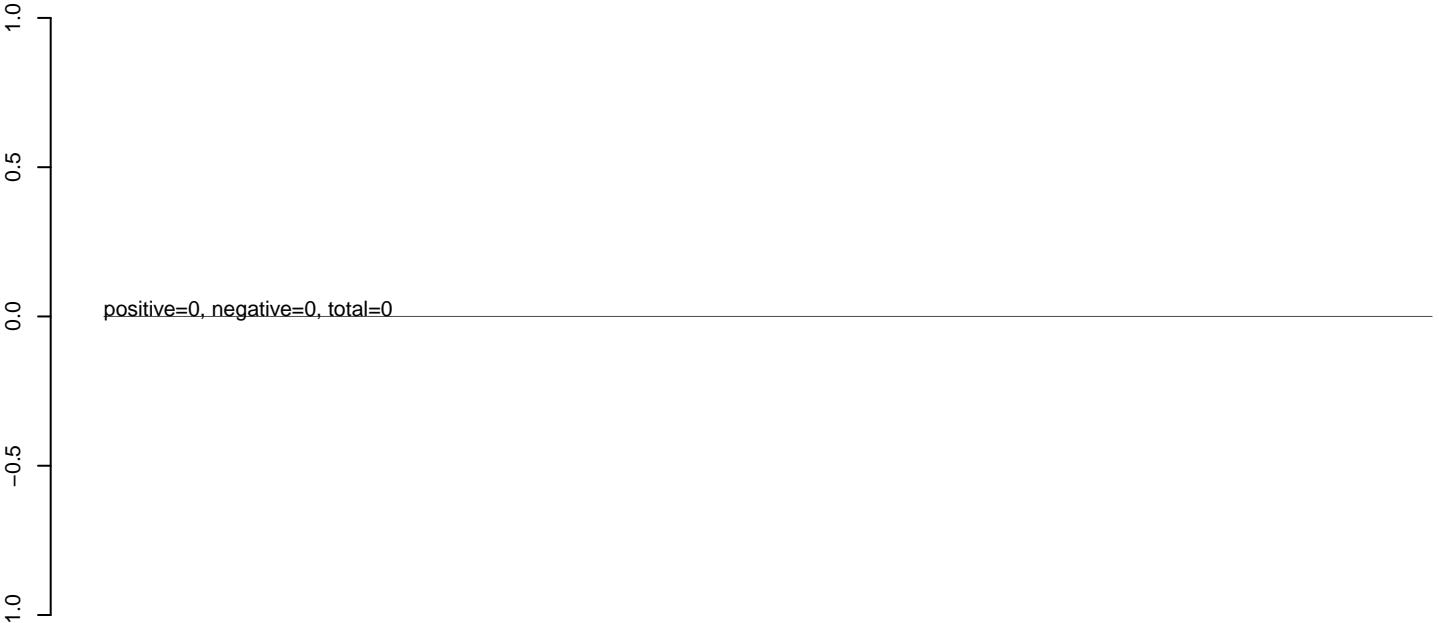
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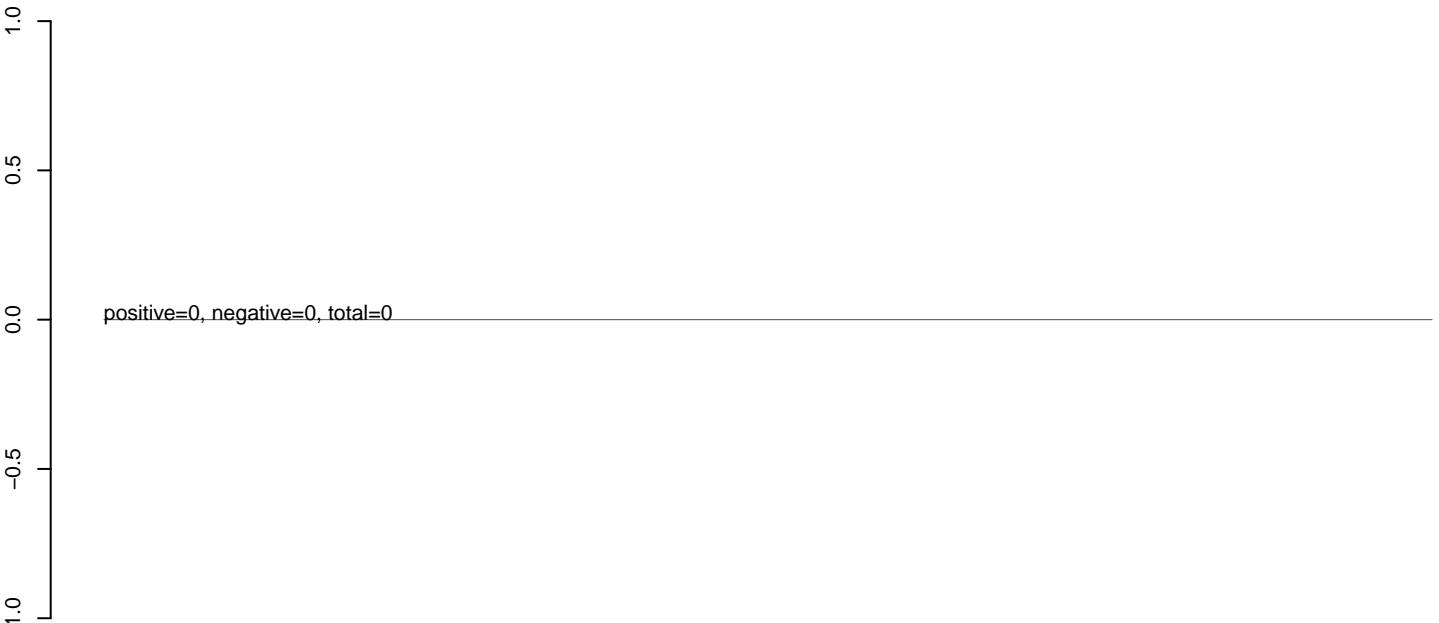
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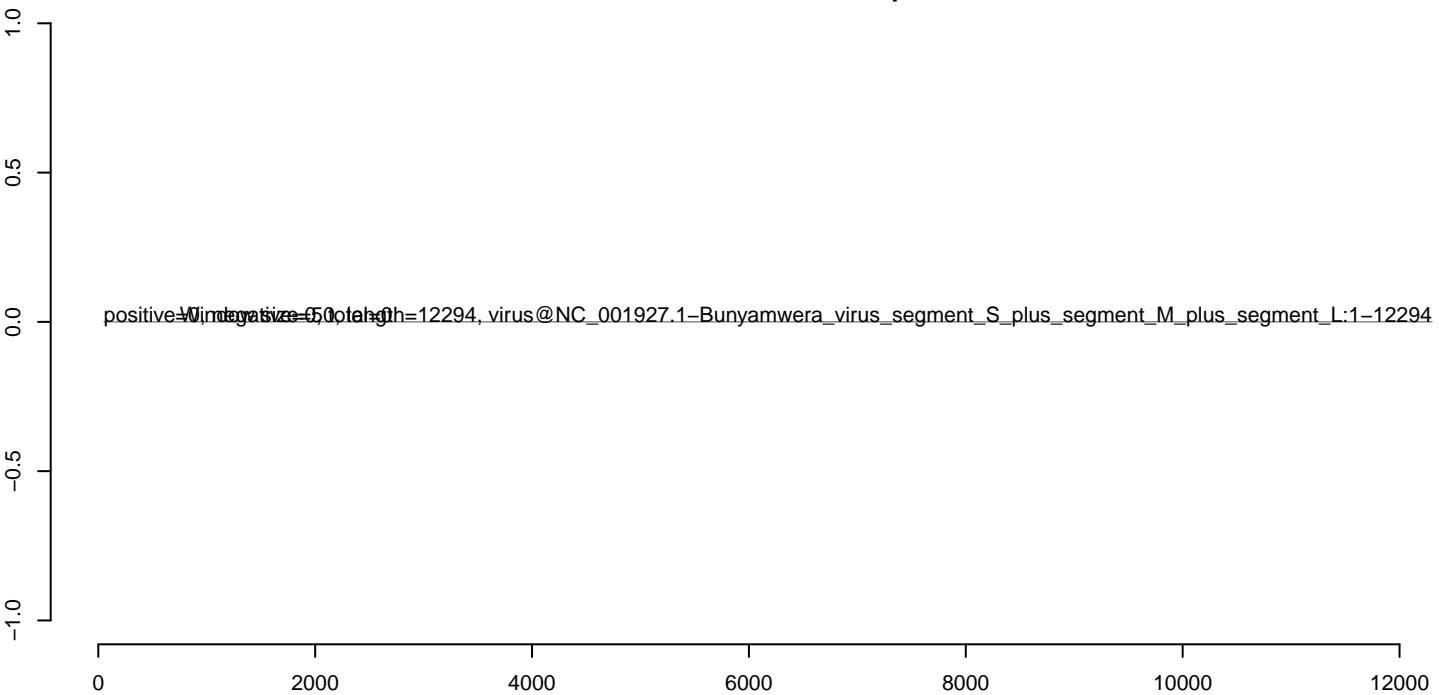
### AnGam\_Fcarc\_TN.18\_23.rep



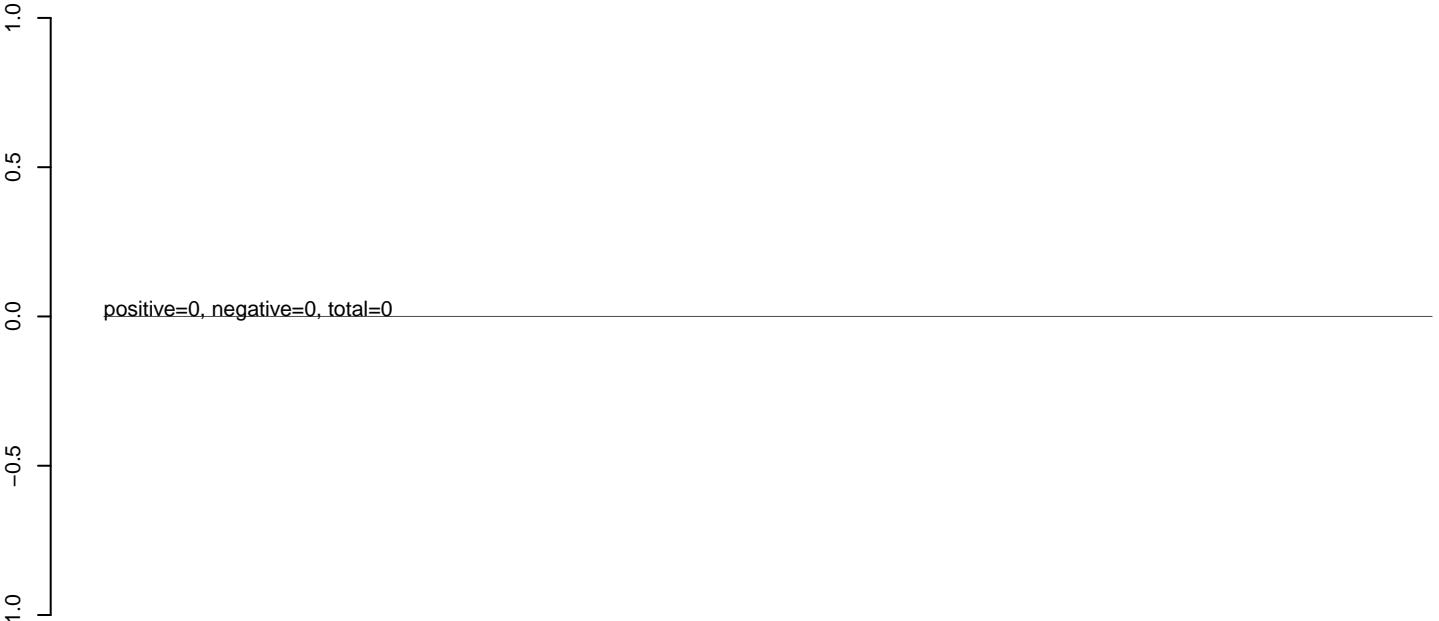
### AnGam\_Fcarc\_TN.24\_35.rep



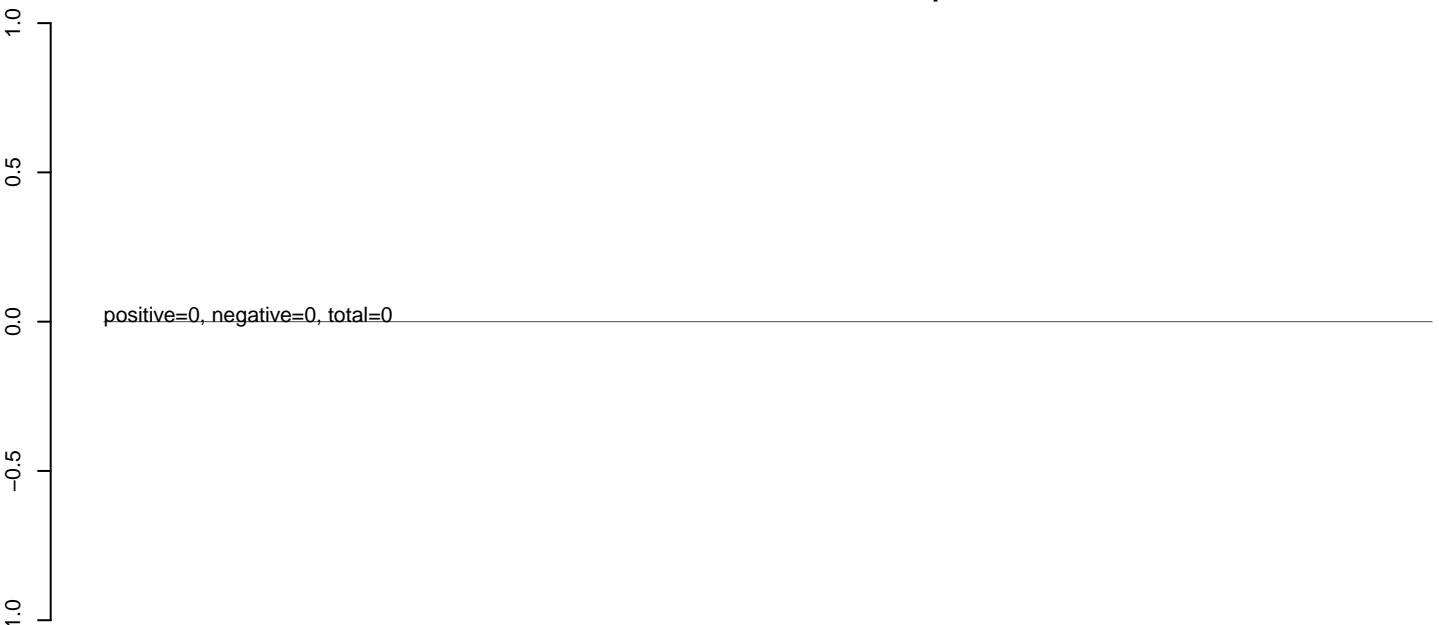
### AnGam\_Fcarc\_TN.rep



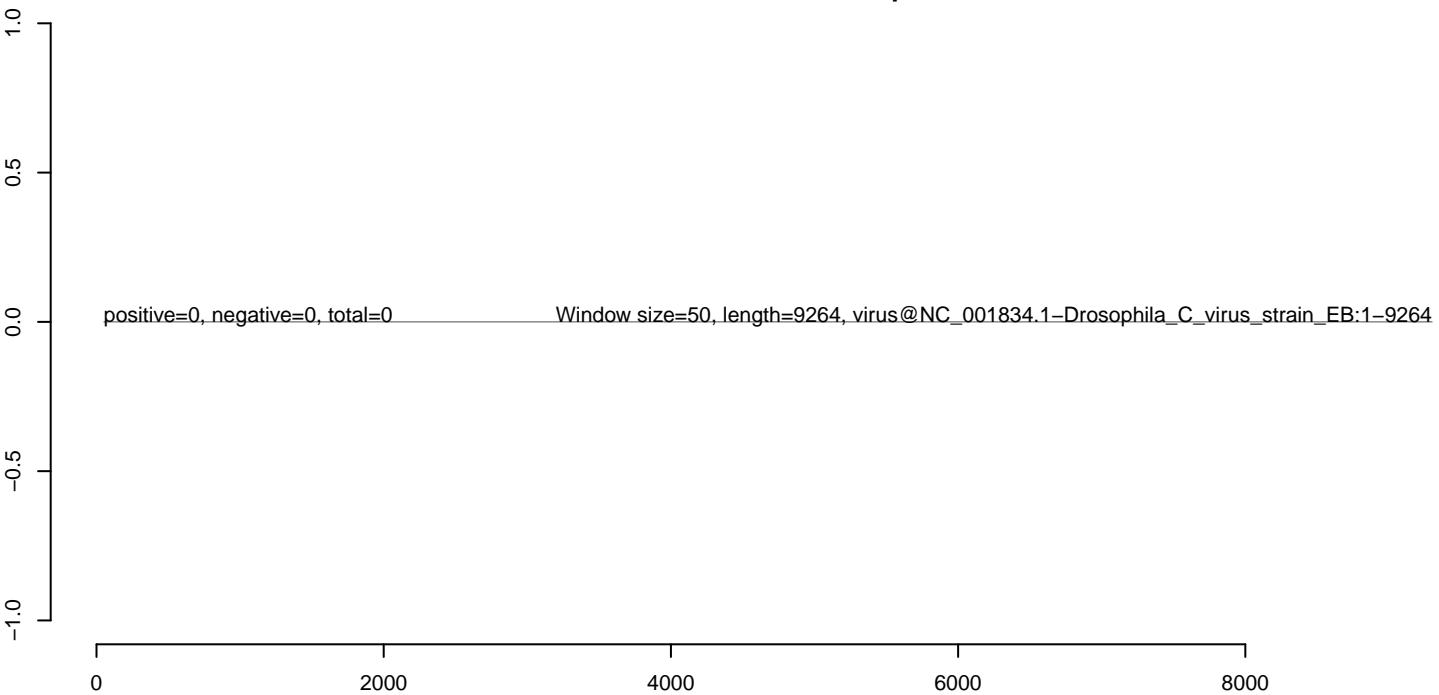
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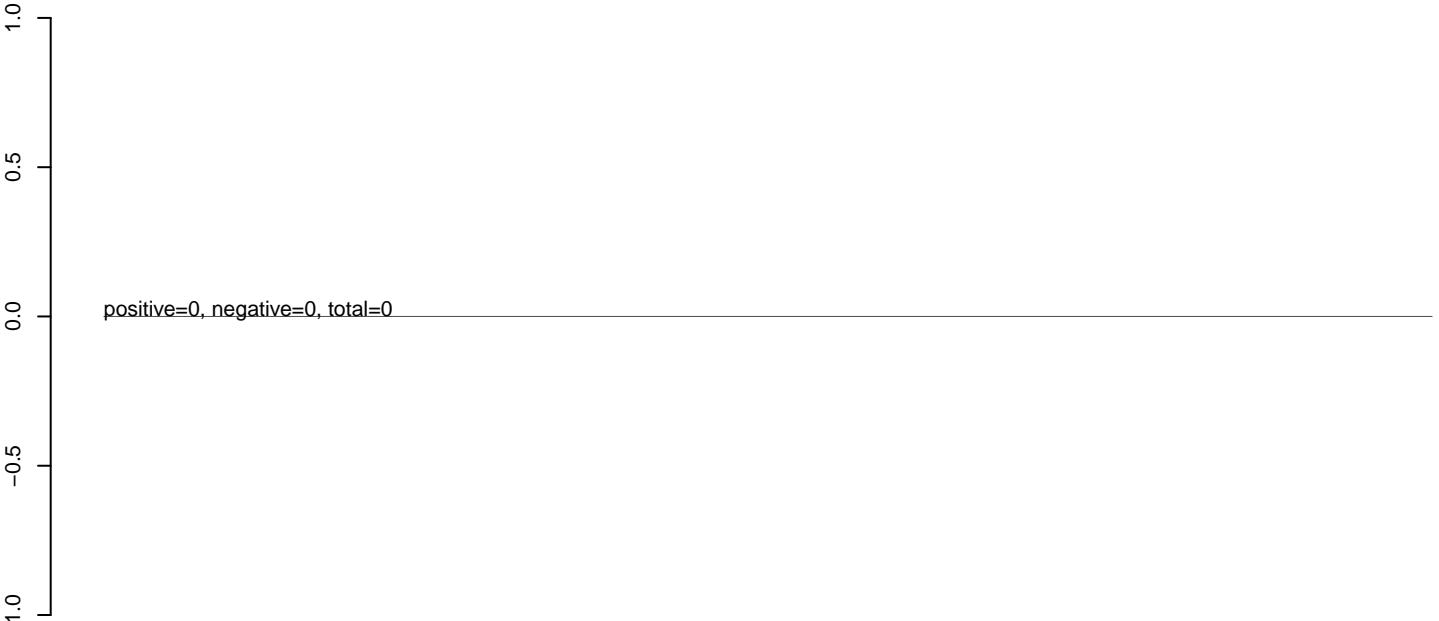
### AnGam\_Fcanc\_TN.24\_35.rep



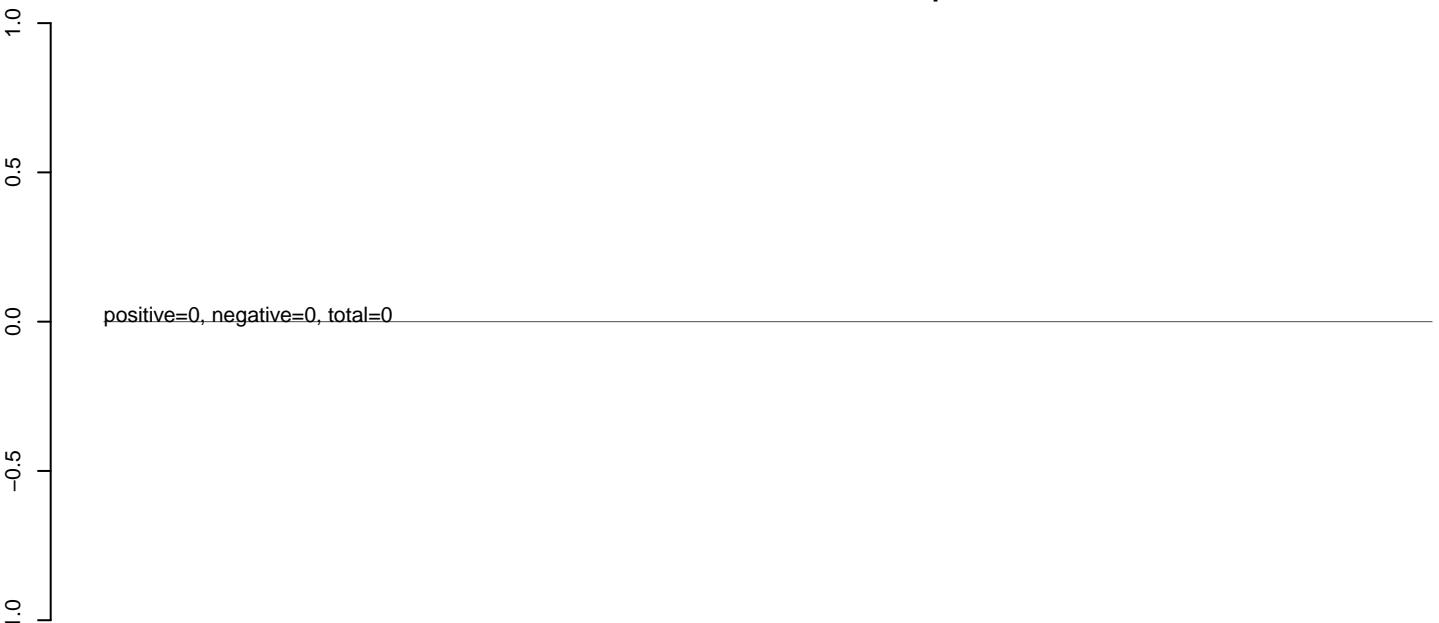
### AnGam\_Fcanc\_TN.rep



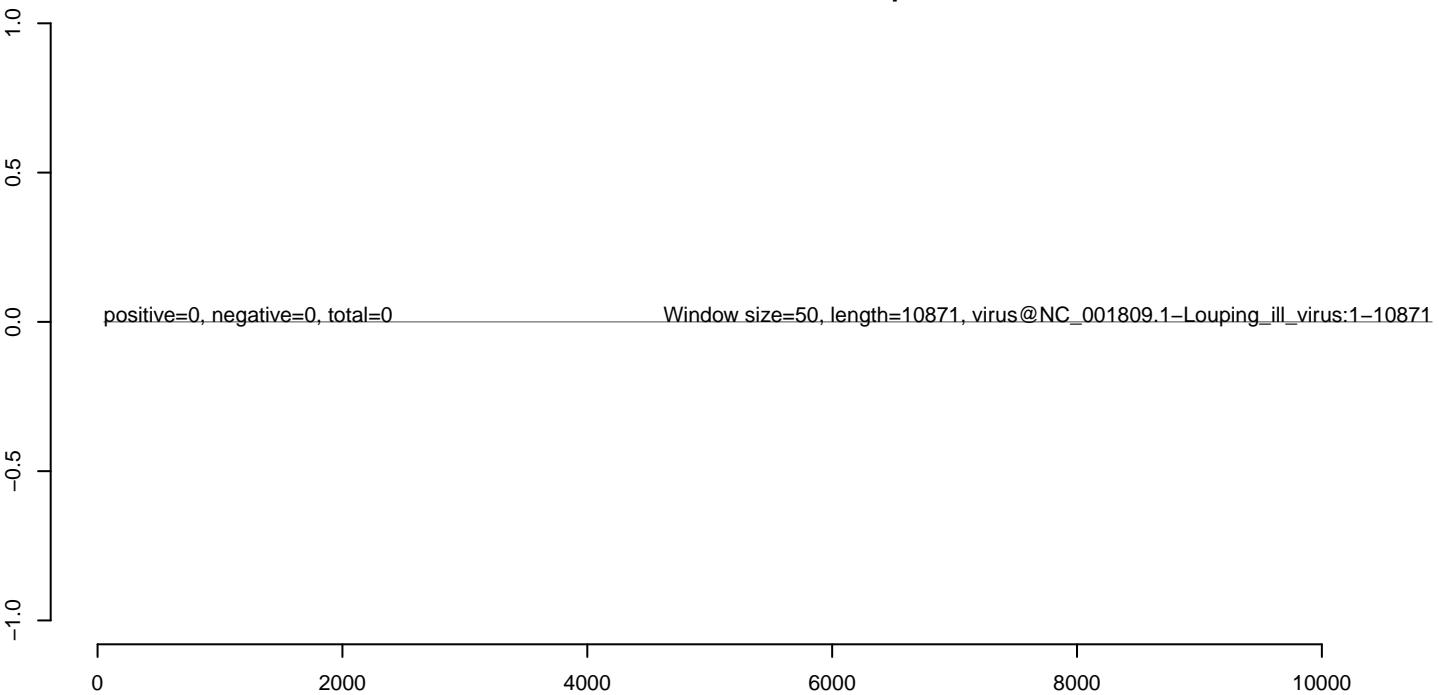
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0

Window size=50, length=11488, virus@NC\_001786.1–Barmah\_Forest\_virus:1–11488

0 2000 4000 6000 8000 10000

### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

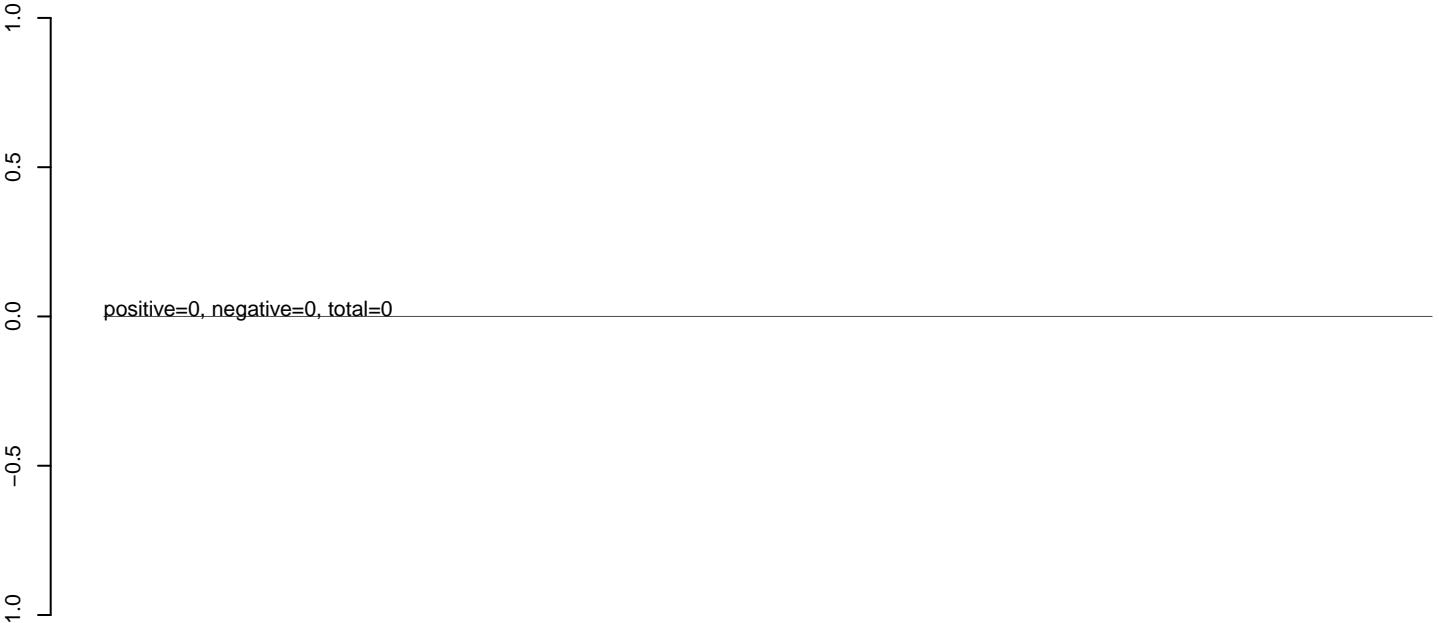
### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0

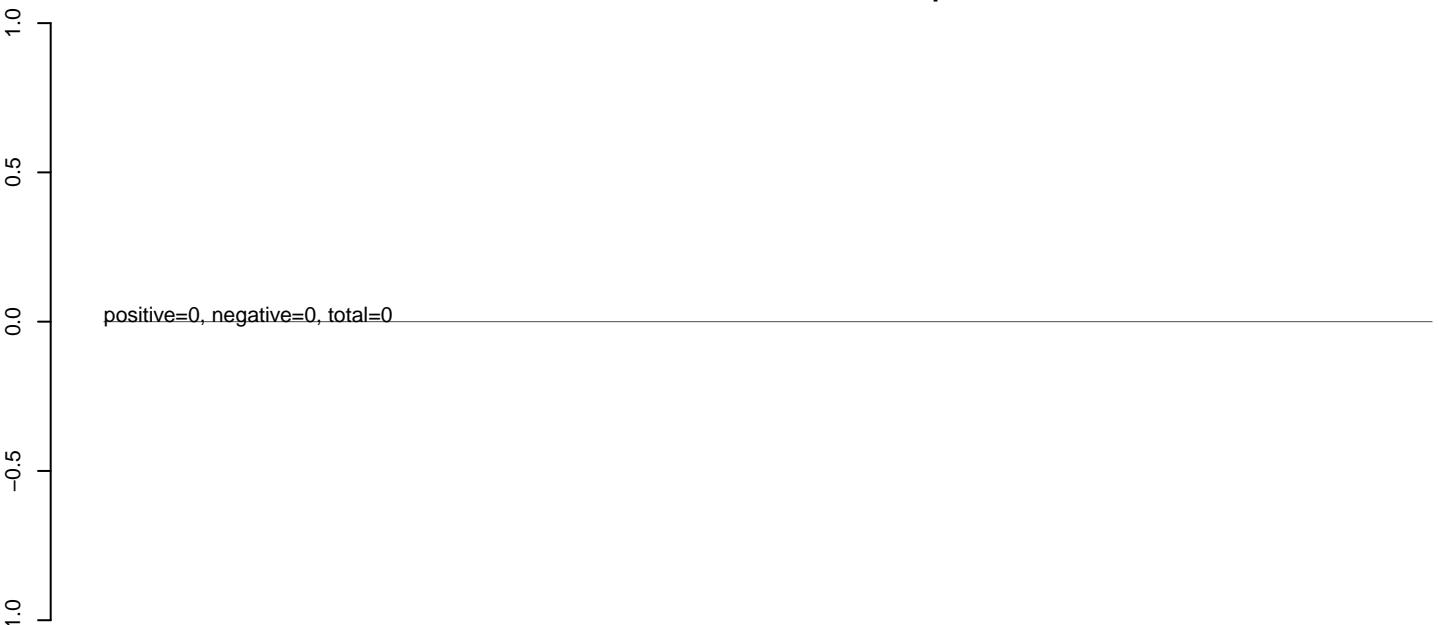
Window size=50, length=11703, virus@NC\_001547.1-Sindbis\_virus:1-11703

0 2000 4000 6000 8000 10000 12000

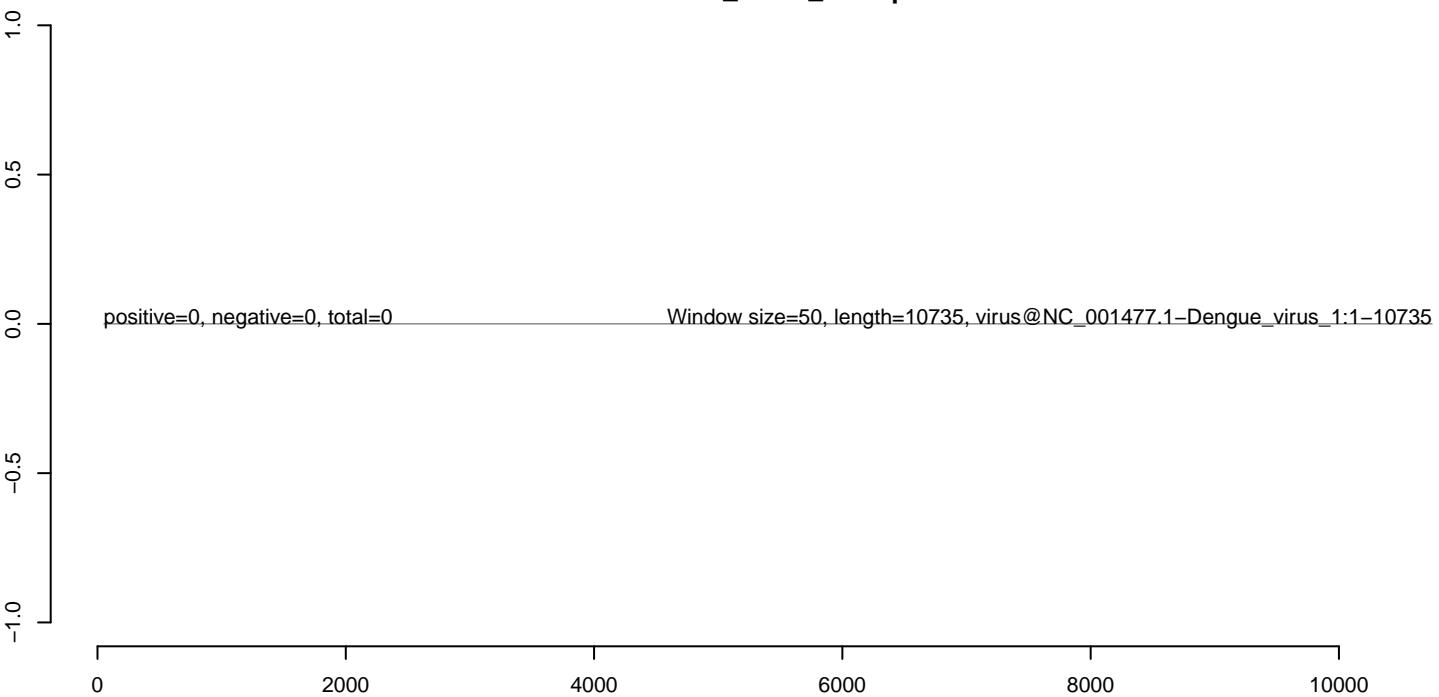
### AnGam\_Fcanc\_TN.18\_23.rep



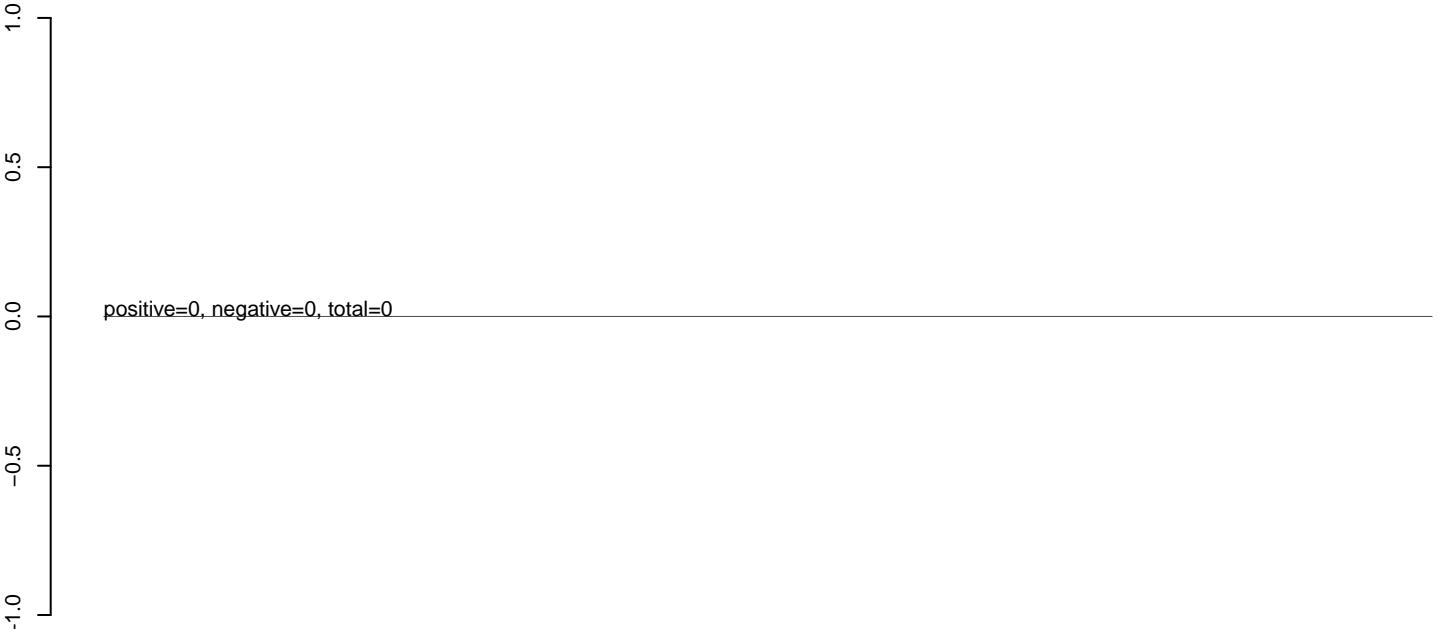
### AnGam\_Fcanc\_TN.24\_35.rep



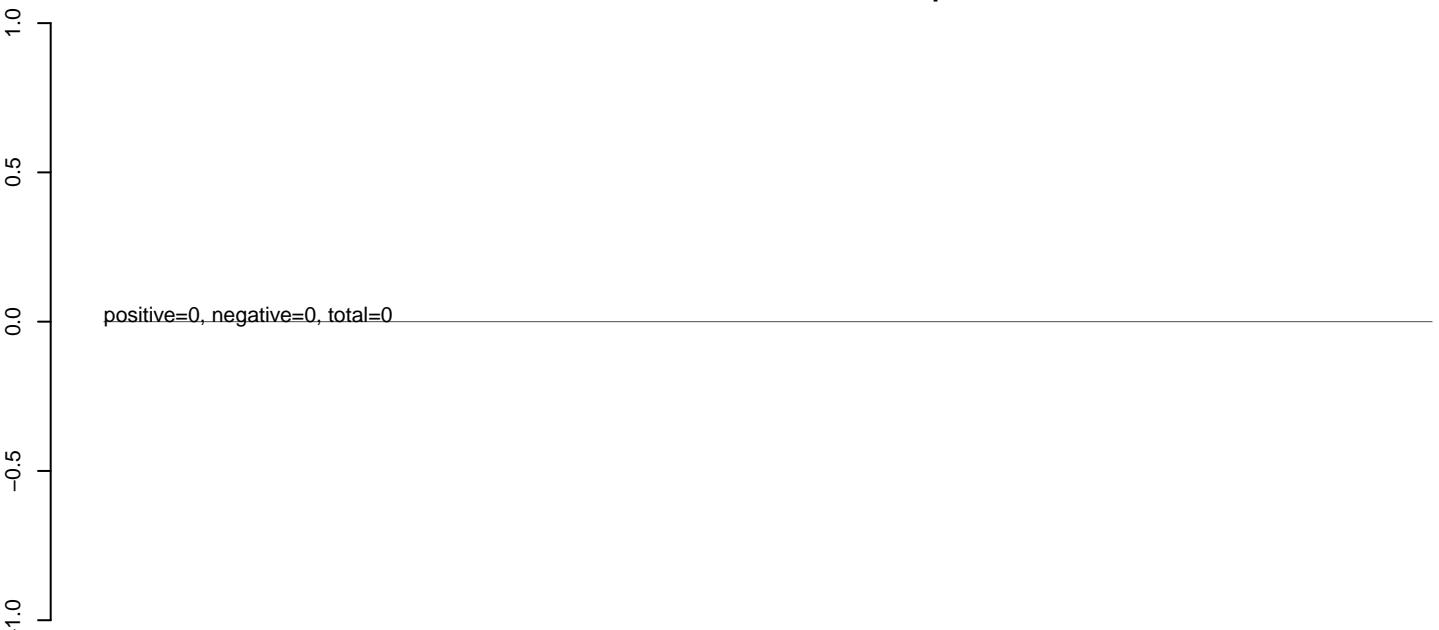
### AnGam\_Fcanc\_TN.rep



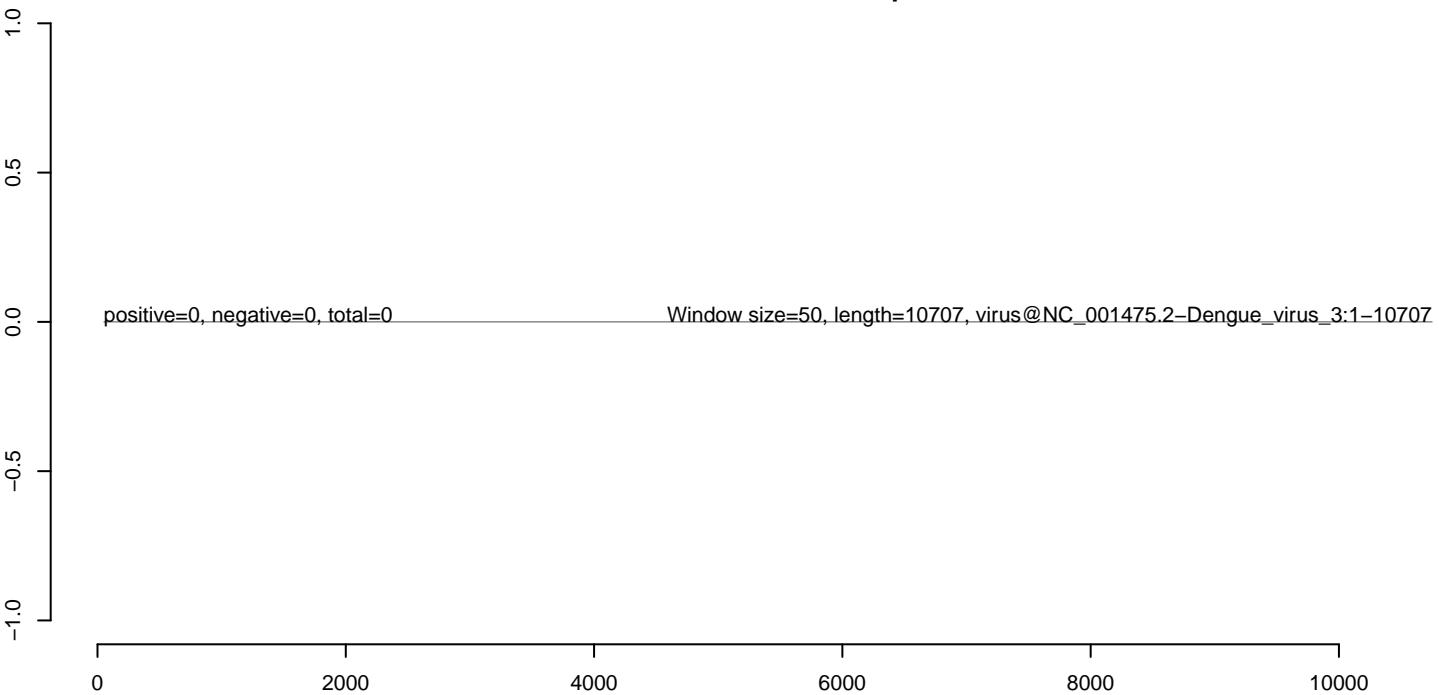
### AnGam\_Fcanc\_TN.18\_23.rep



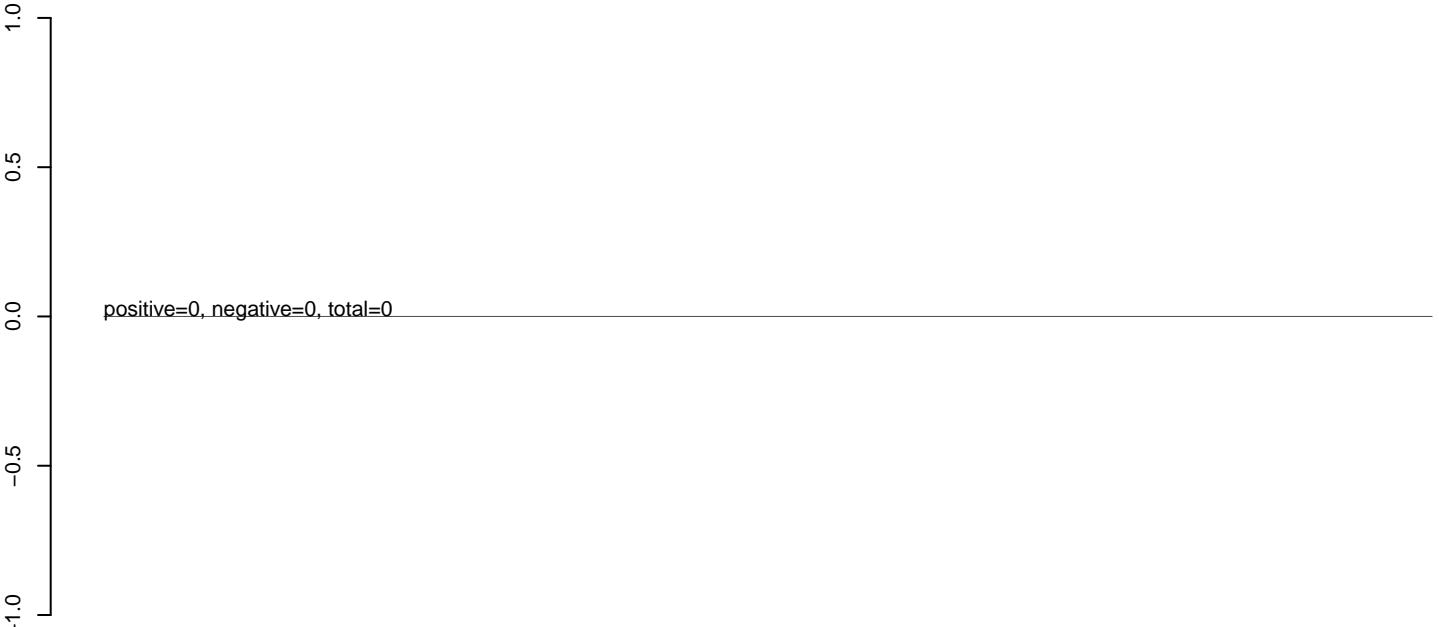
### AnGam\_Fcanc\_TN.24\_35.rep



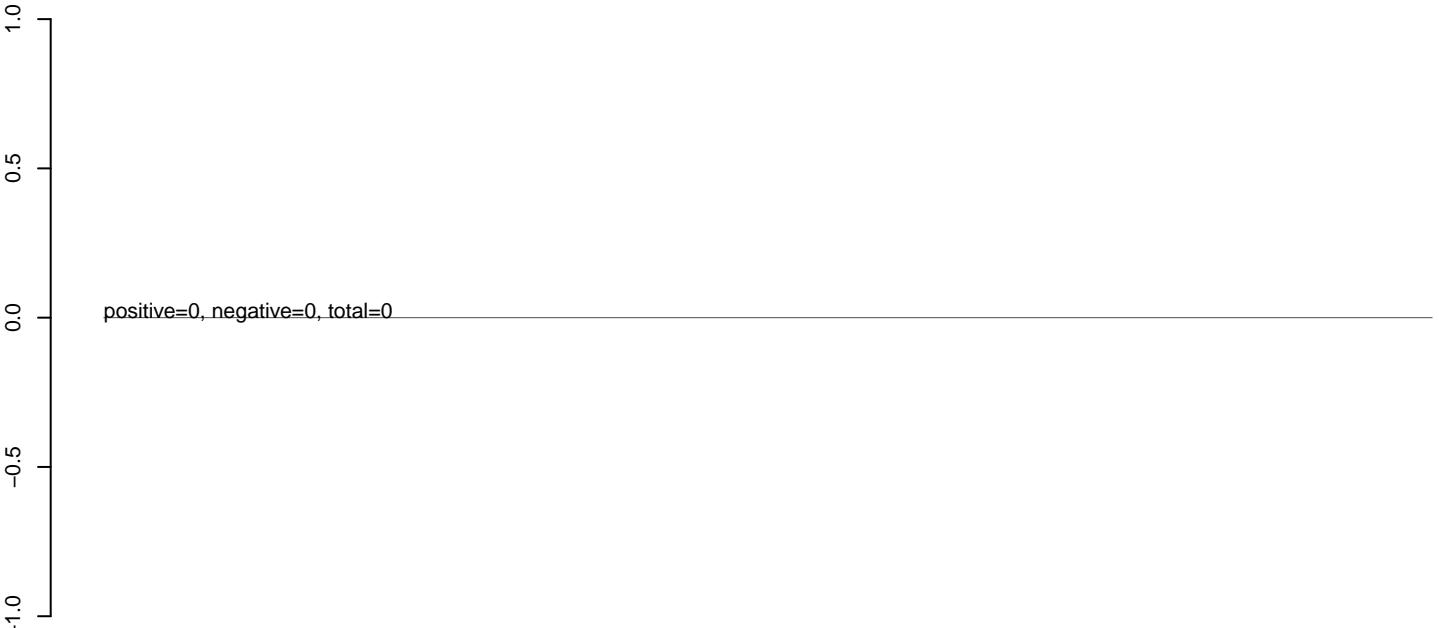
### AnGam\_Fcanc\_TN.rep



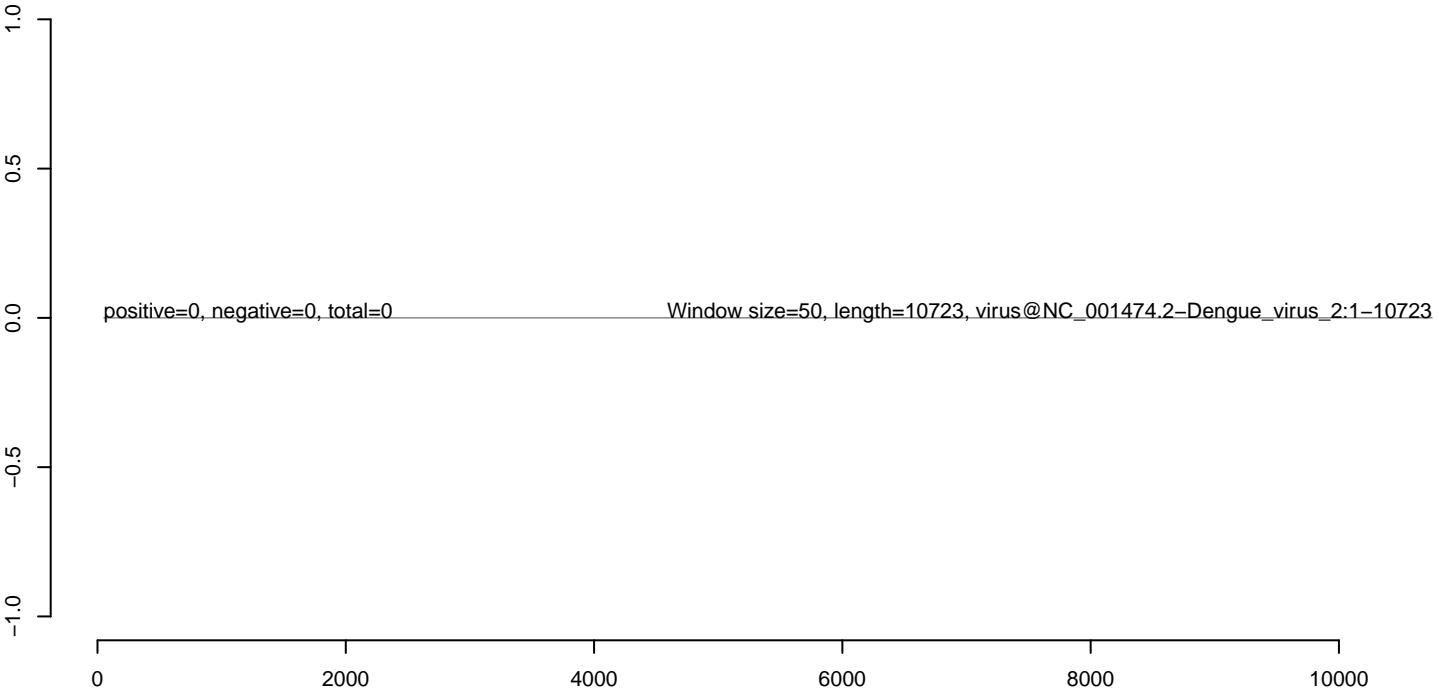
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

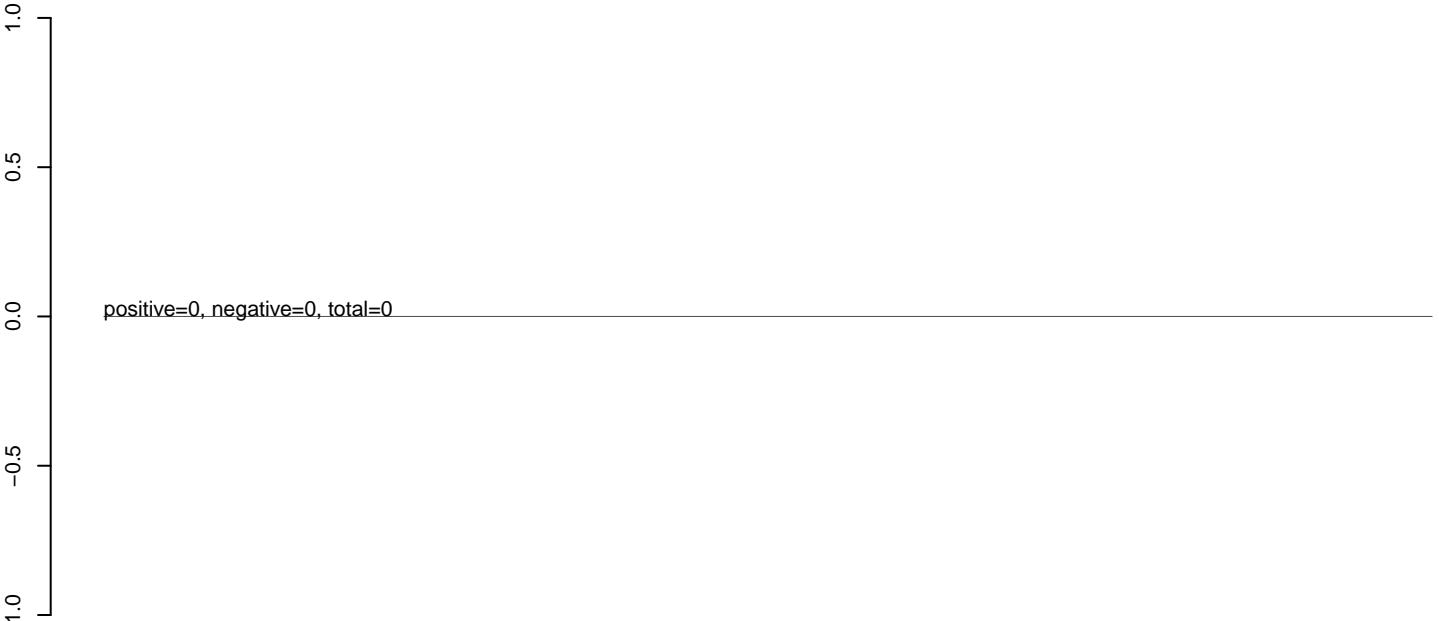
positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.rep

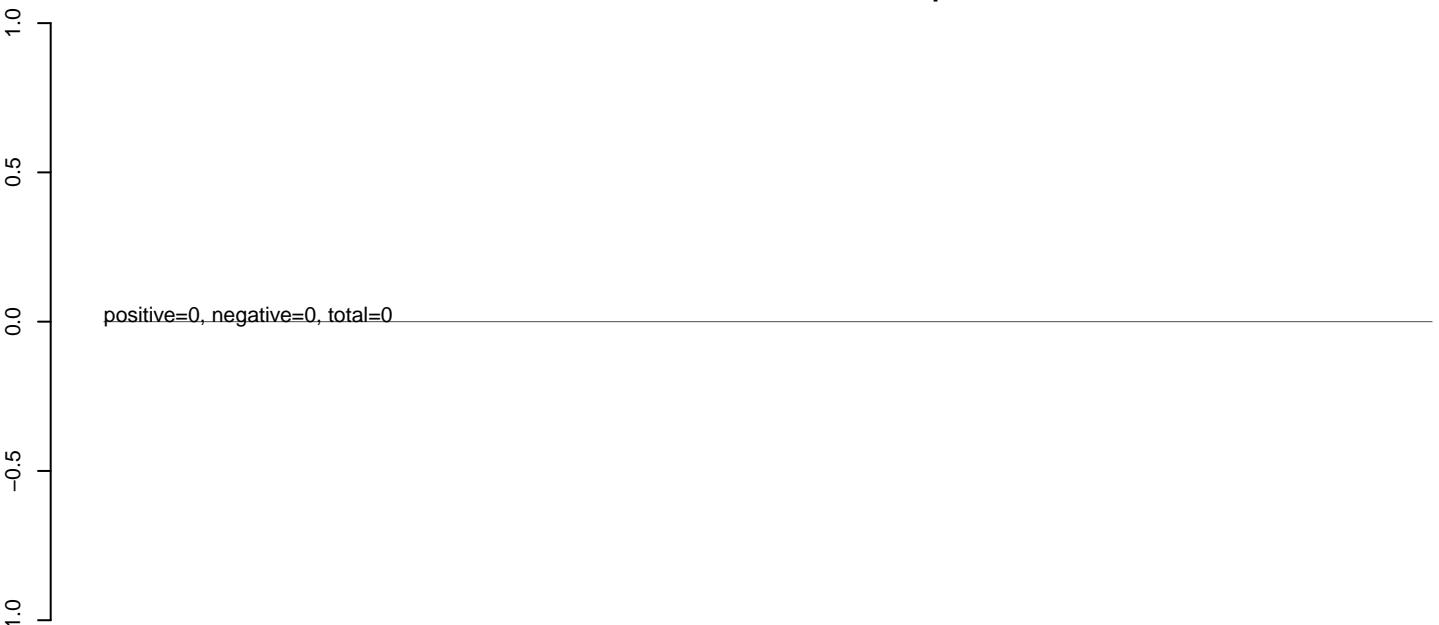
positive=0, negative=0, total=0

Window size=50, length=11444, virus@NC\_001449.1–Venezuelan equine encephalitis virus:1–11444

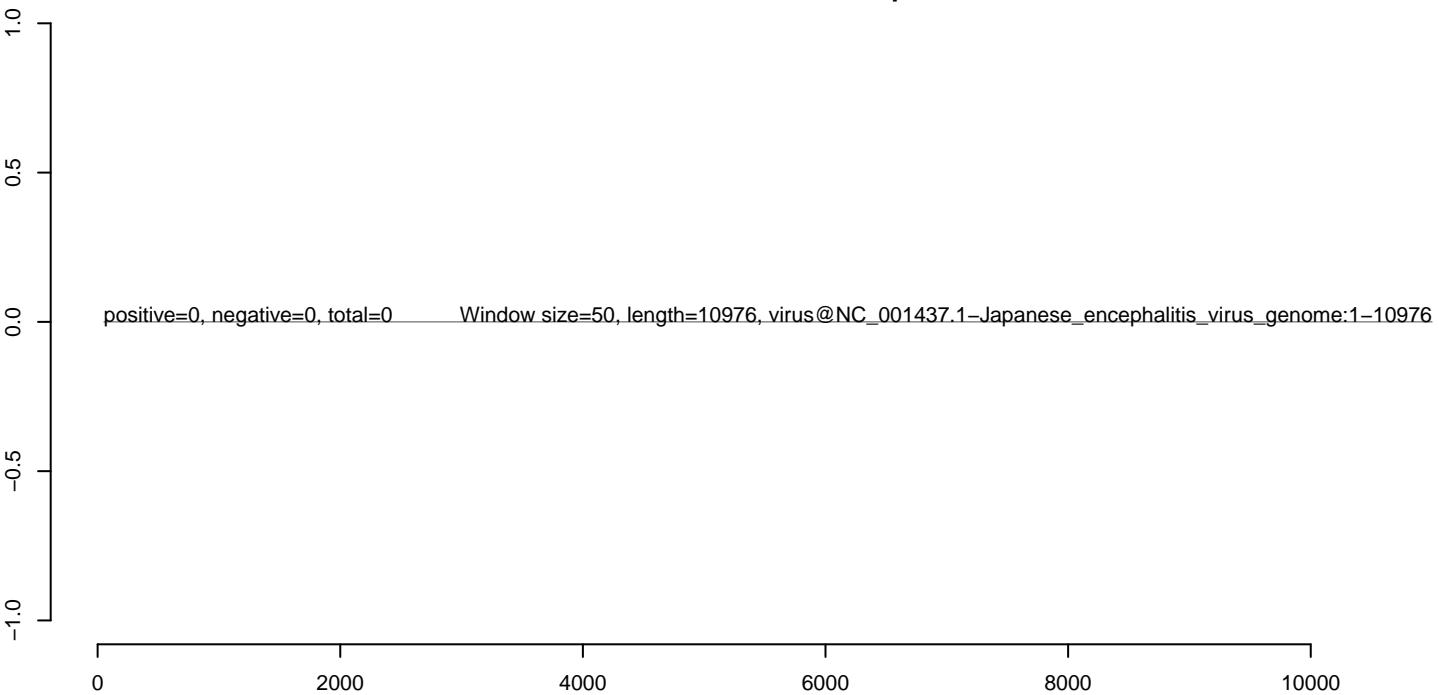
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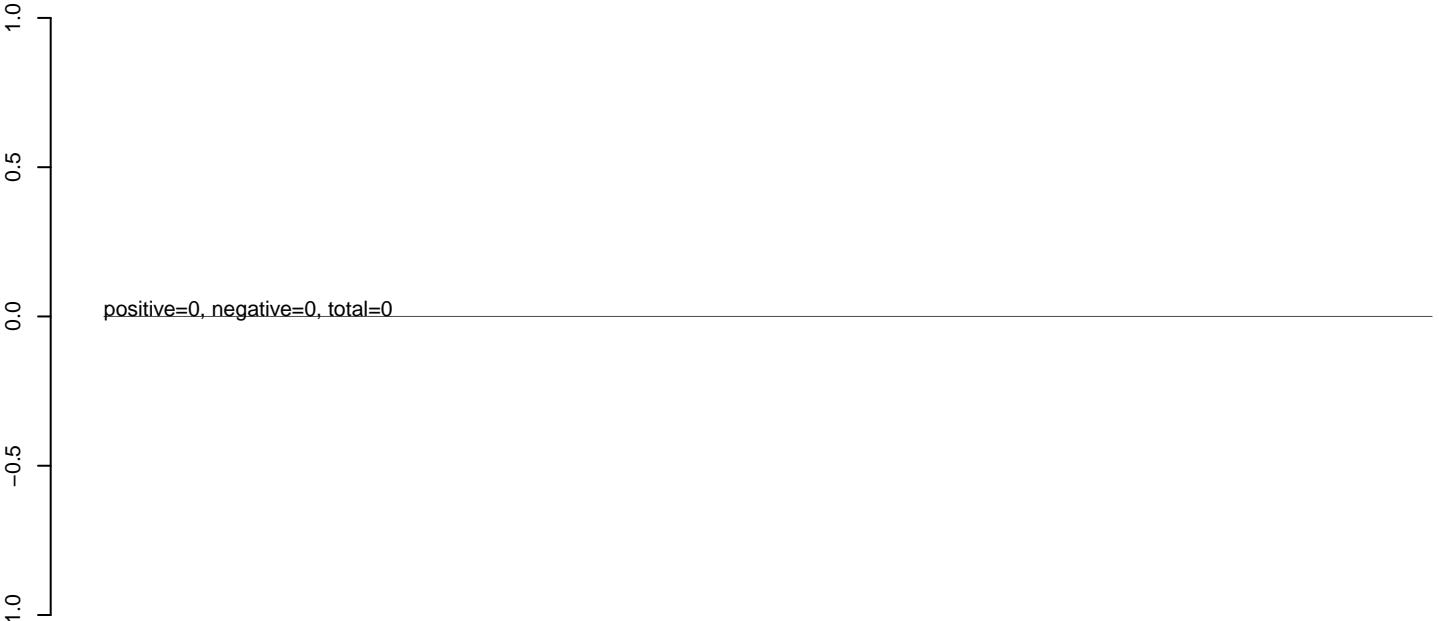
### AnGam\_Fcanc\_TN.24\_35.rep



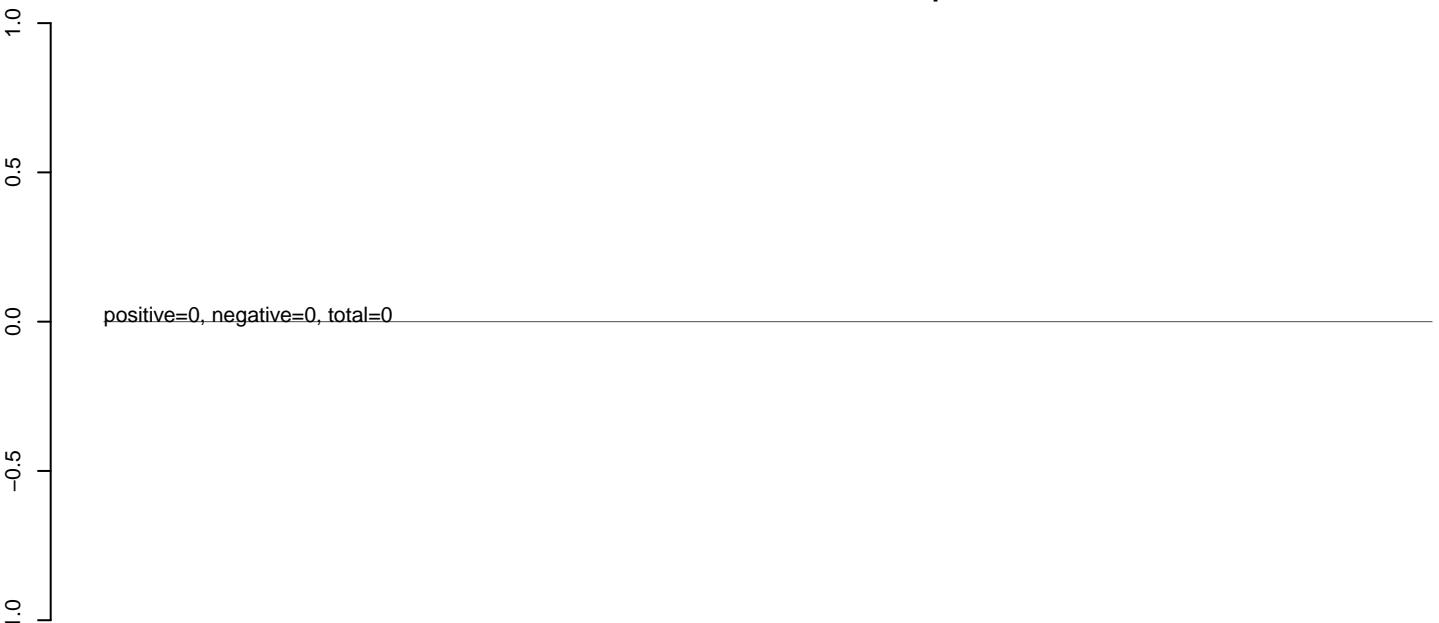
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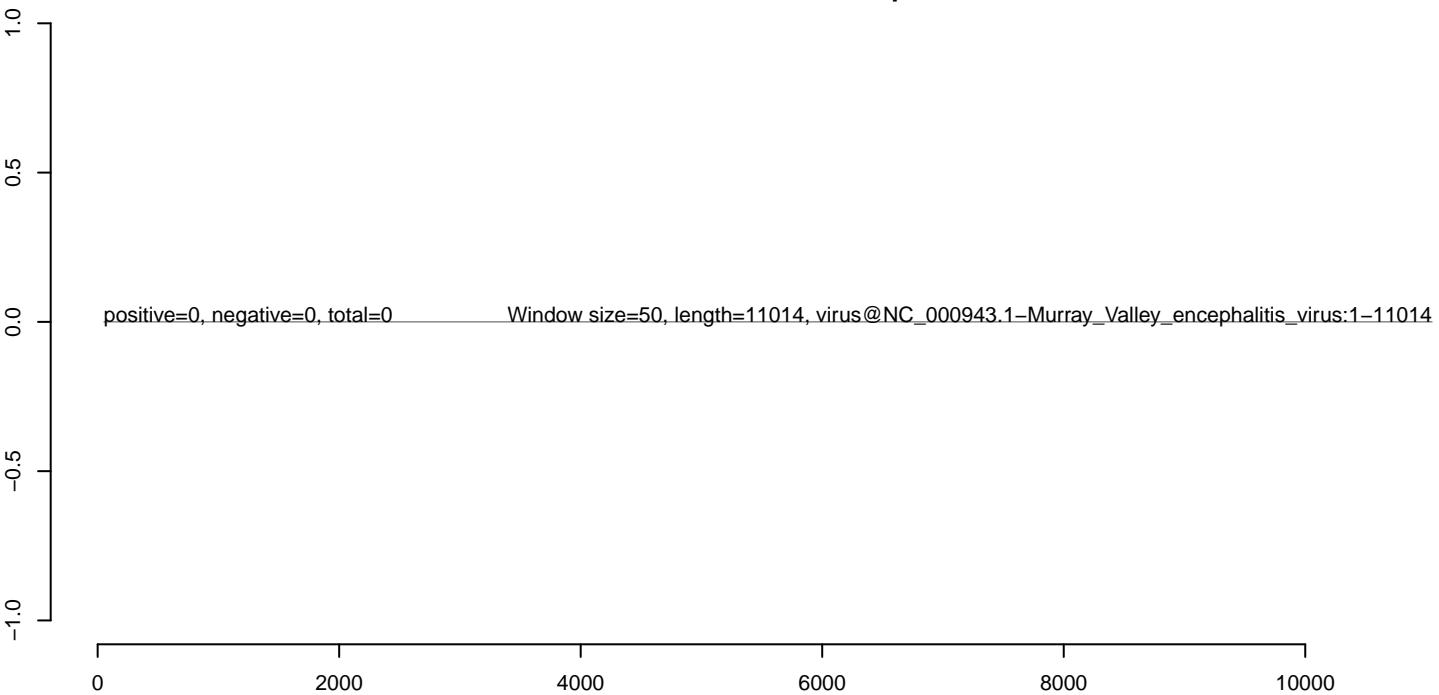
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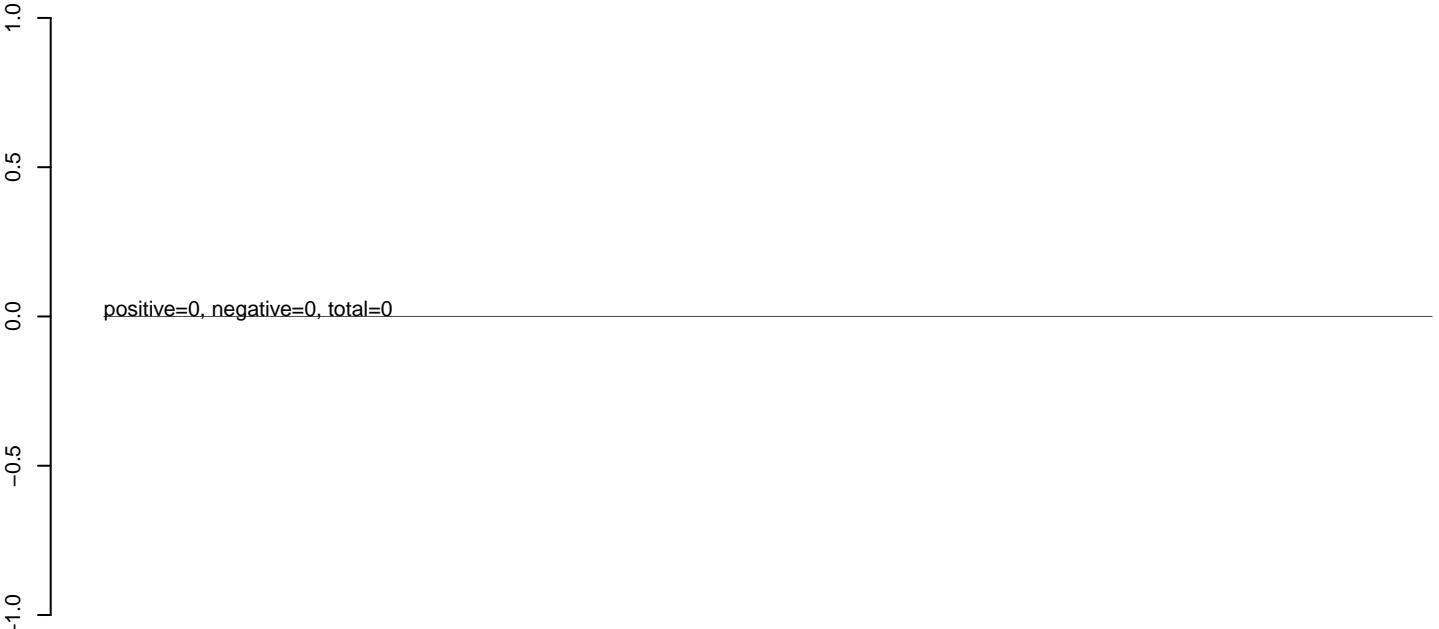
### AnGam\_Fcanc\_TN.24\_35.rep



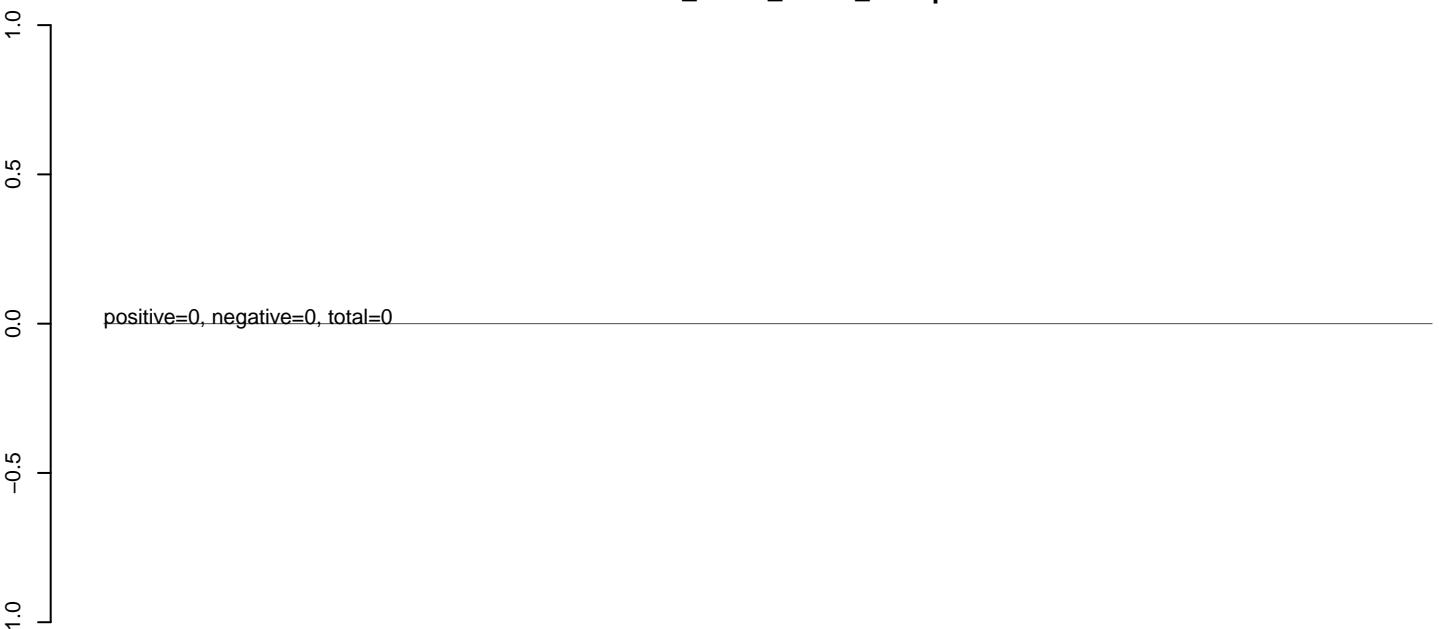
### AnGam\_Fcanc\_TN.rep



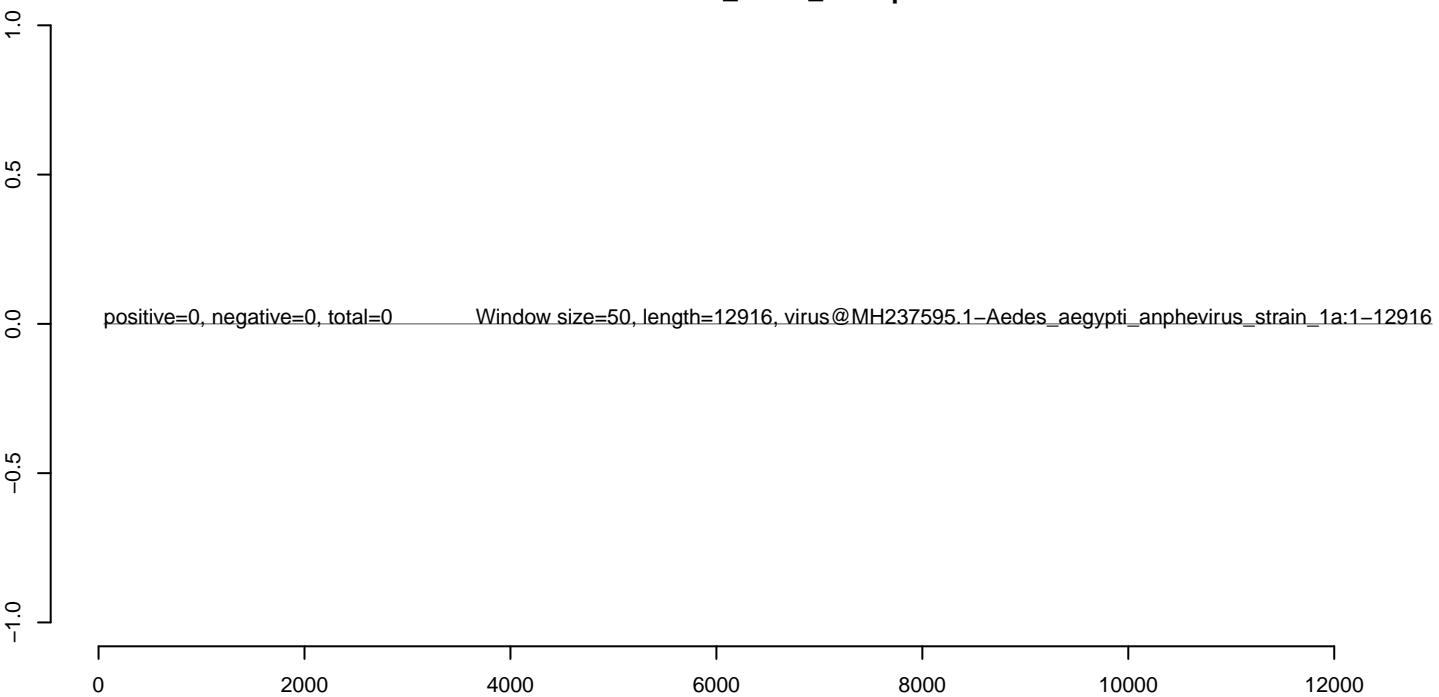
### AnGam\_Fcanc\_TN.18\_23.rep



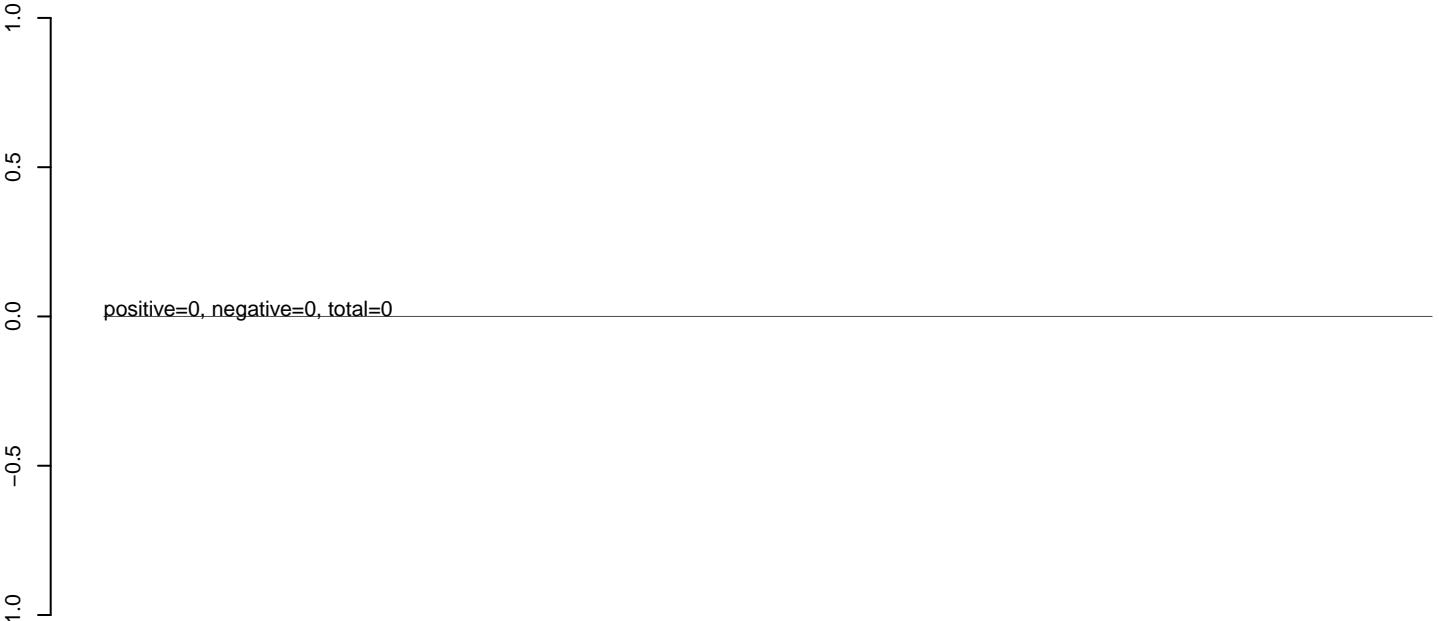
### AnGam\_Fcanc\_TN.24\_35.rep



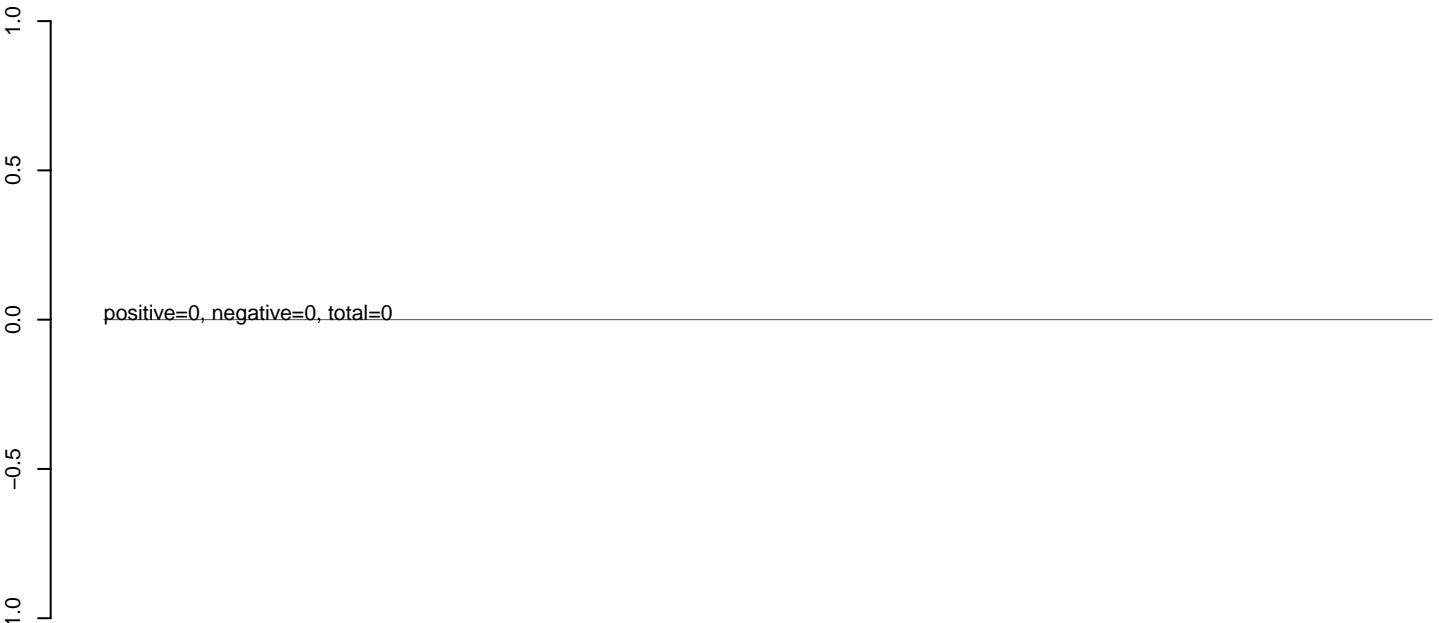
### AnGam\_Fcanc\_TN.rep



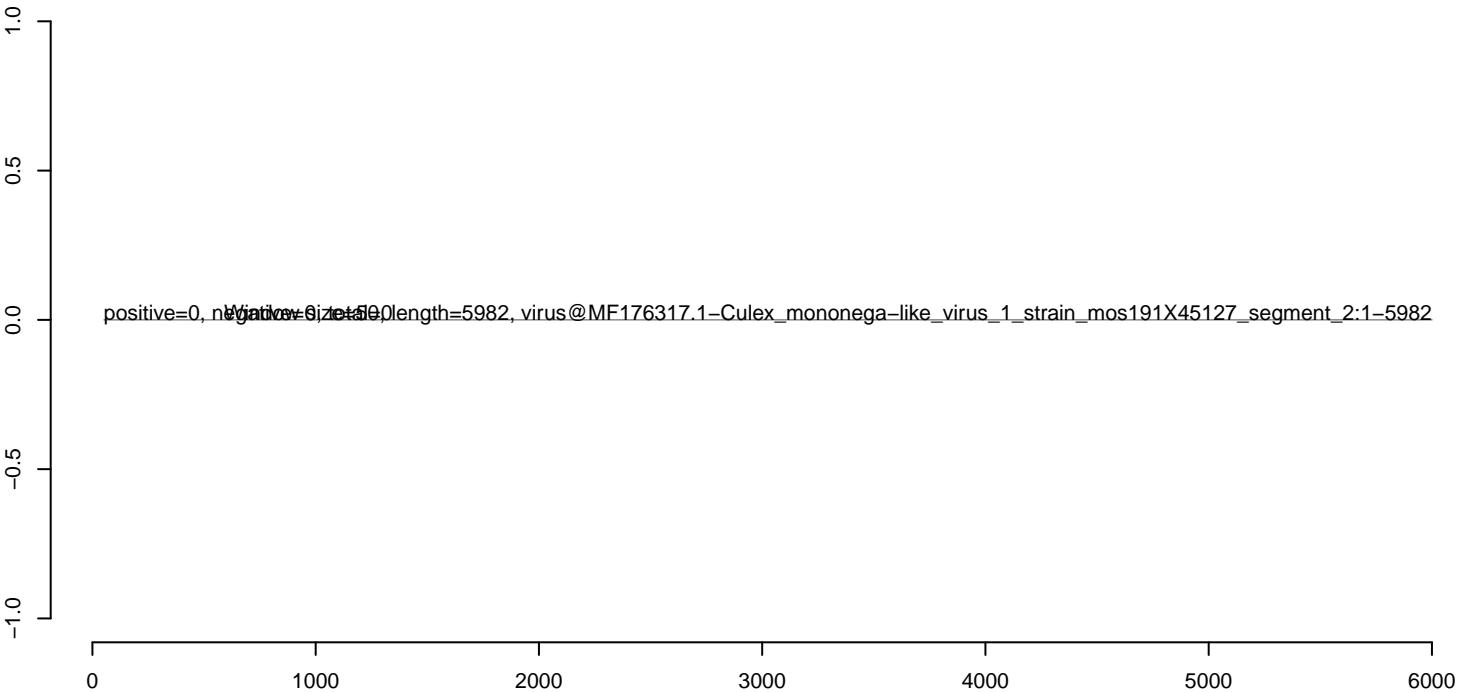
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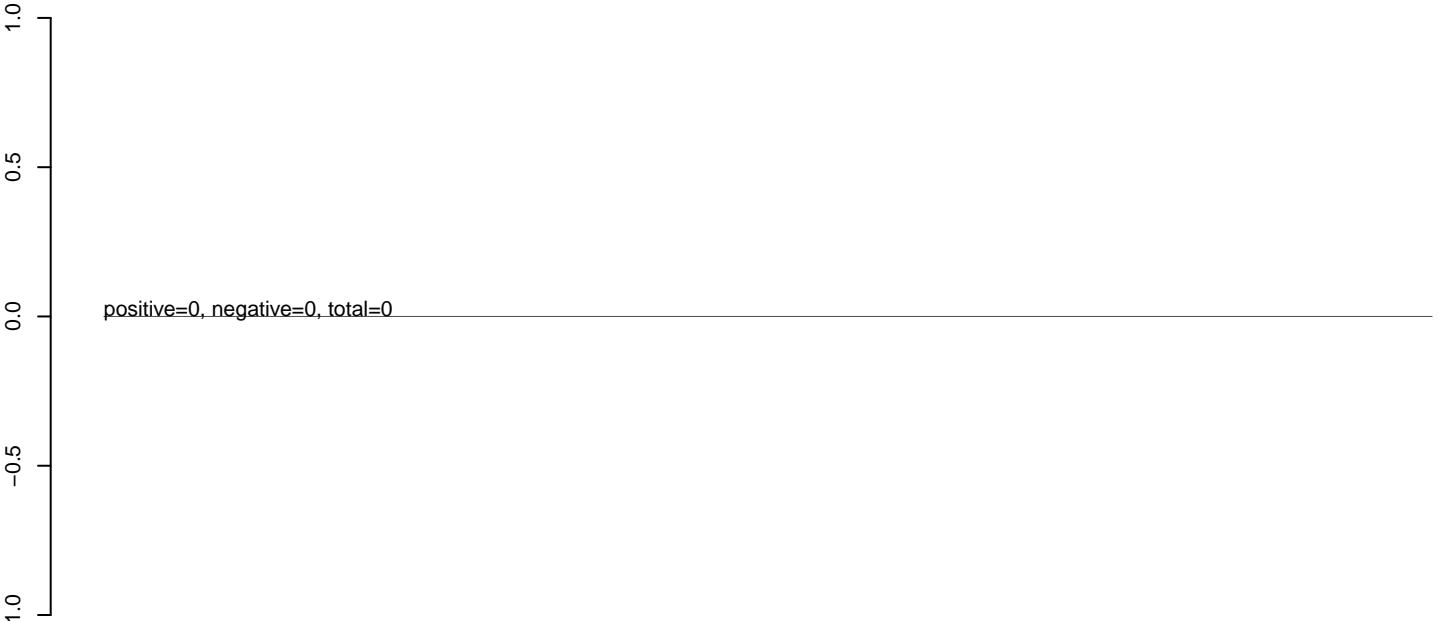
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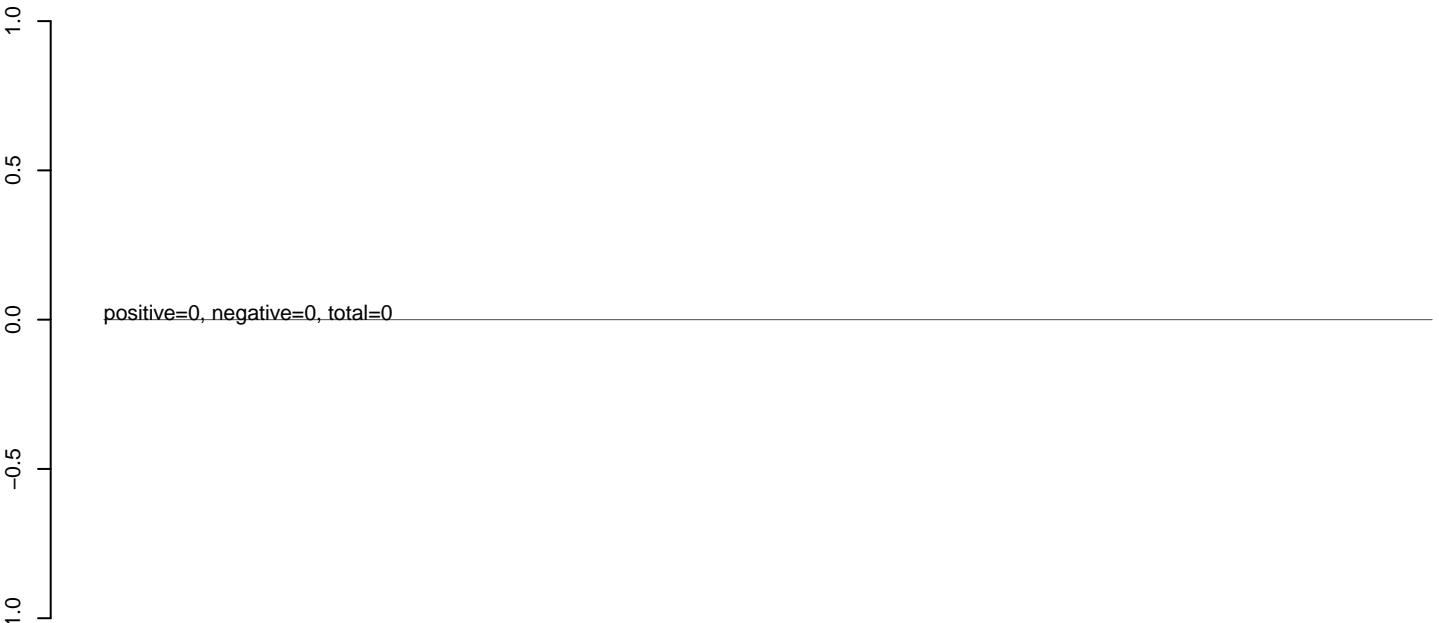
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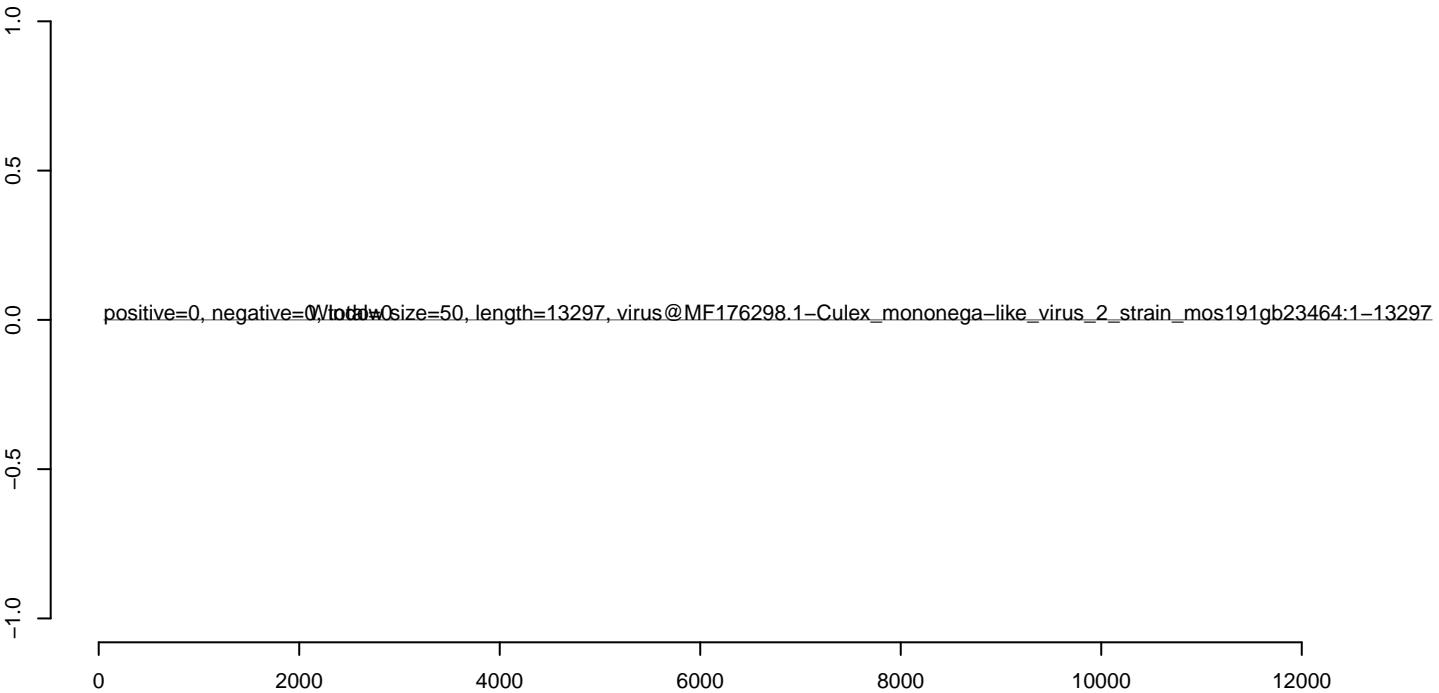
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

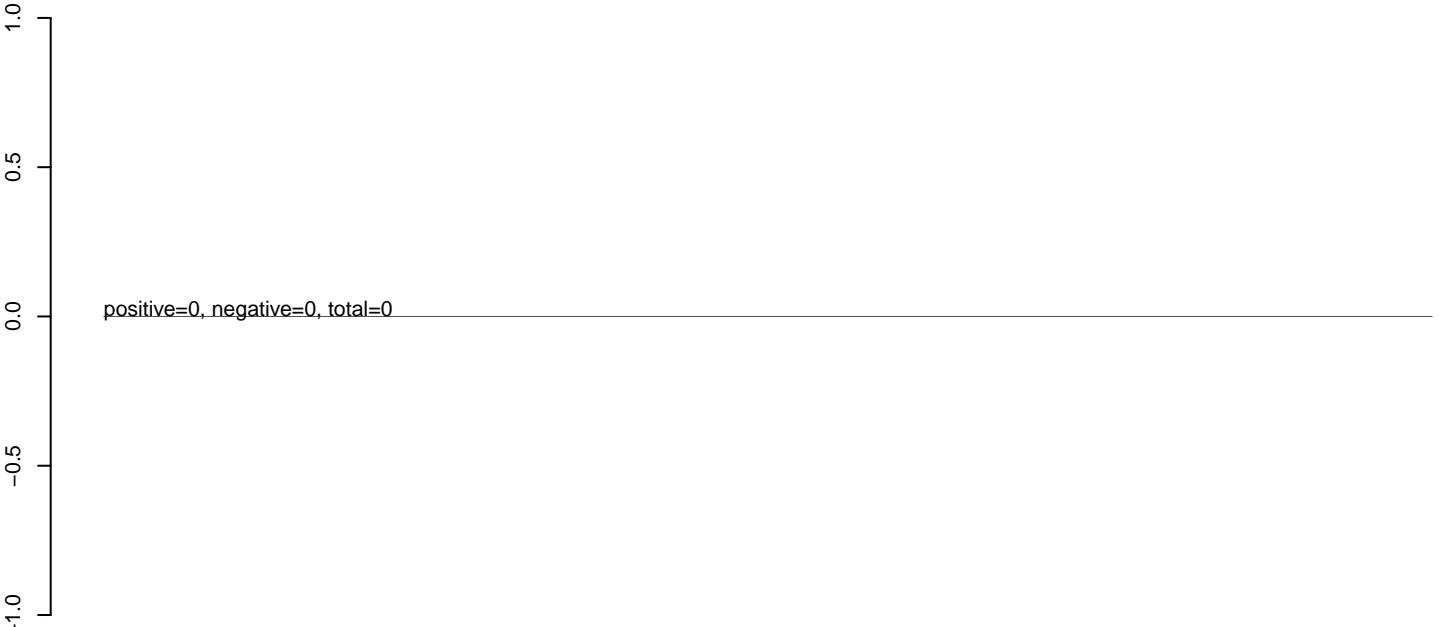
### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

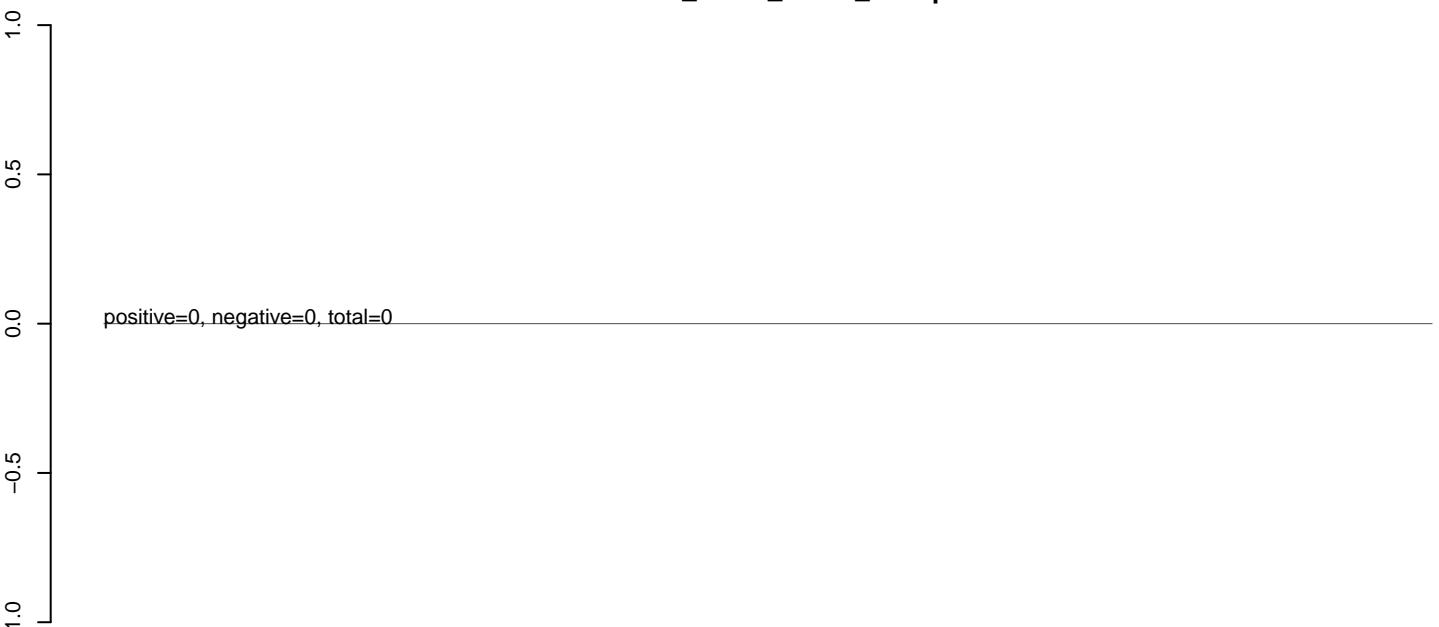
### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0    Window size=50, length=11657, virus@M20162.1–Ross\_River\_virus\_RRV\_strain\_NB5092\_RNA:1–11657

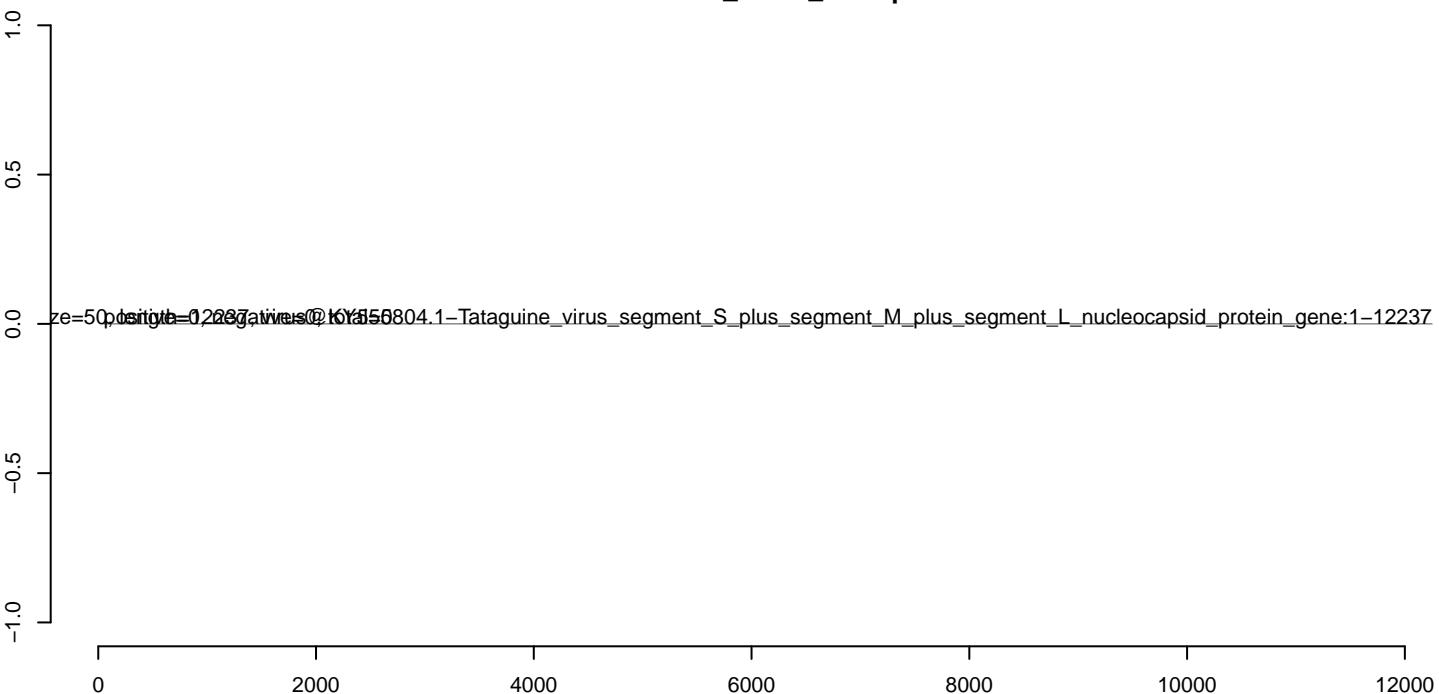
### AnGam\_Fcanc\_TN.18\_23.rep



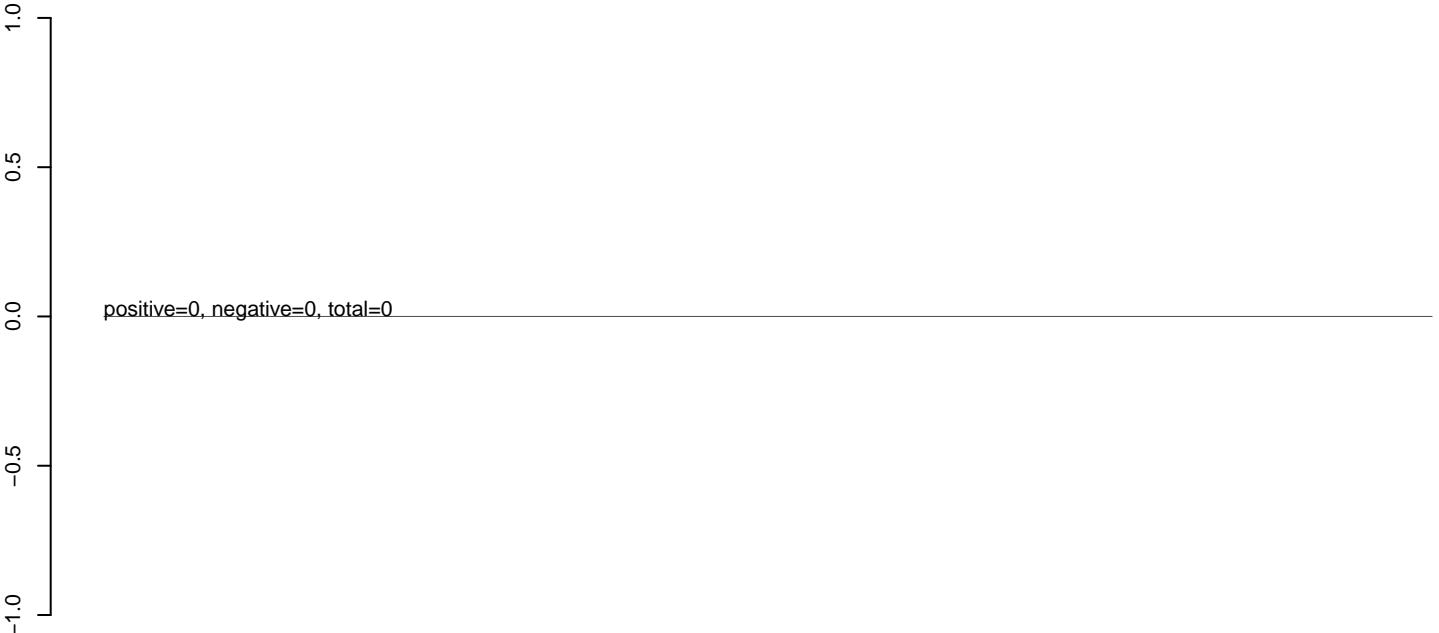
### AnGam\_Fcanc\_TN.24\_35.rep



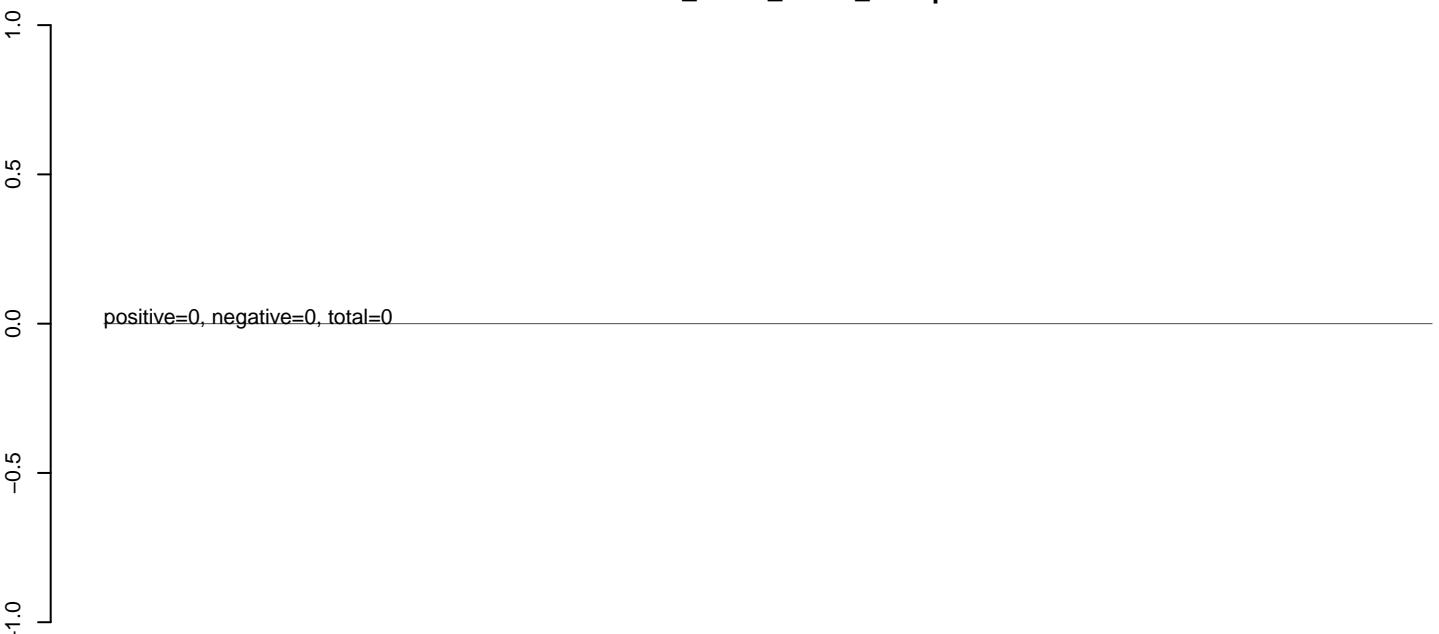
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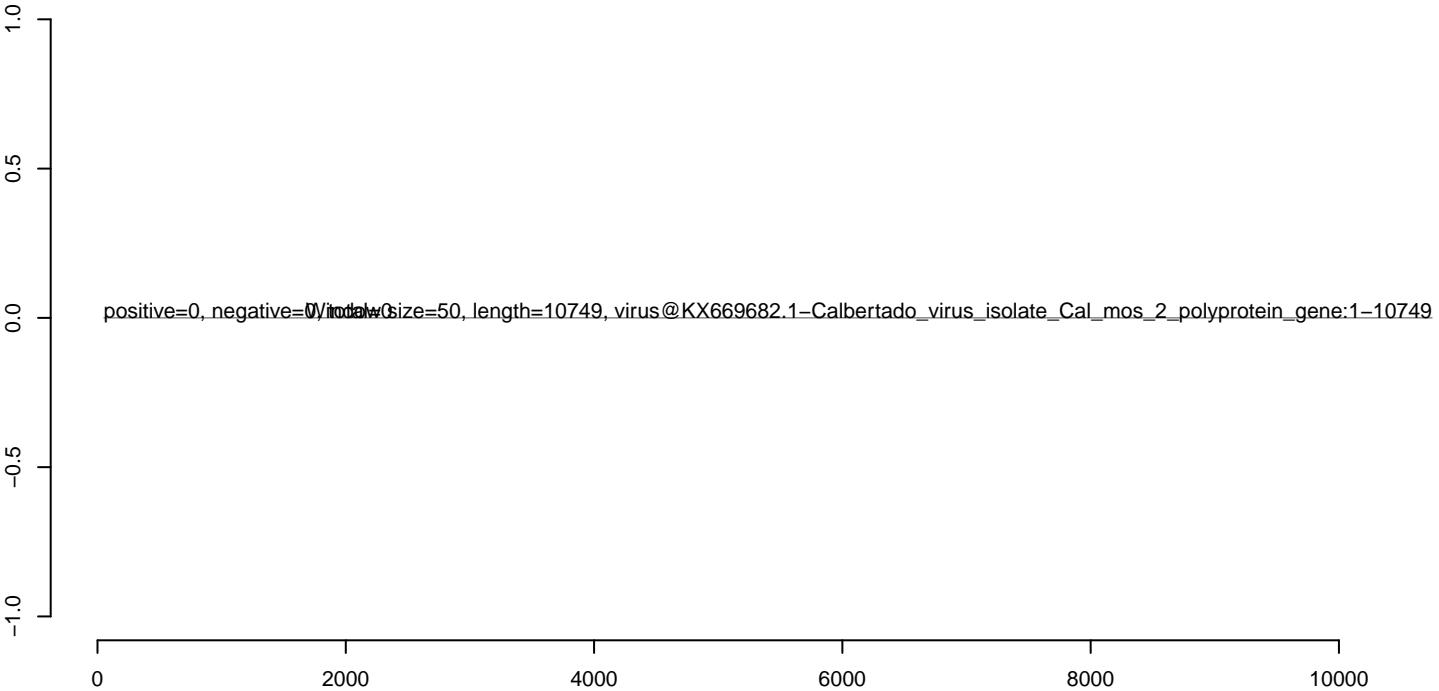
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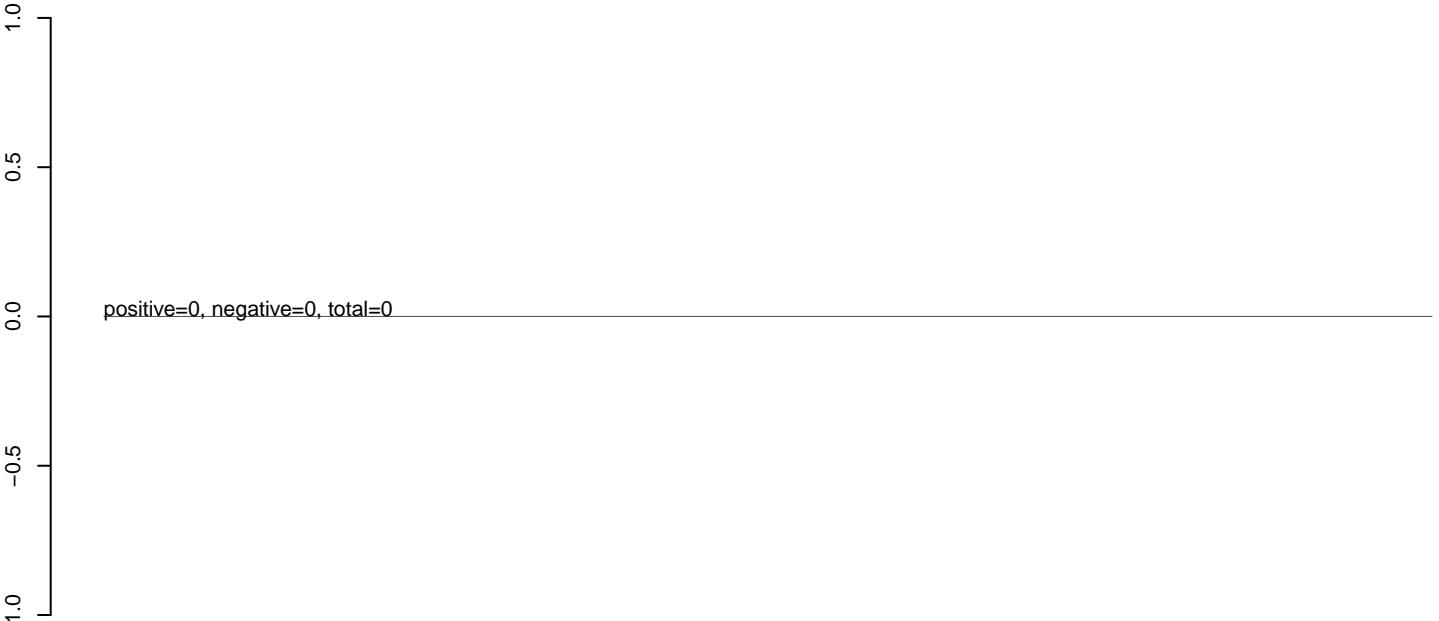
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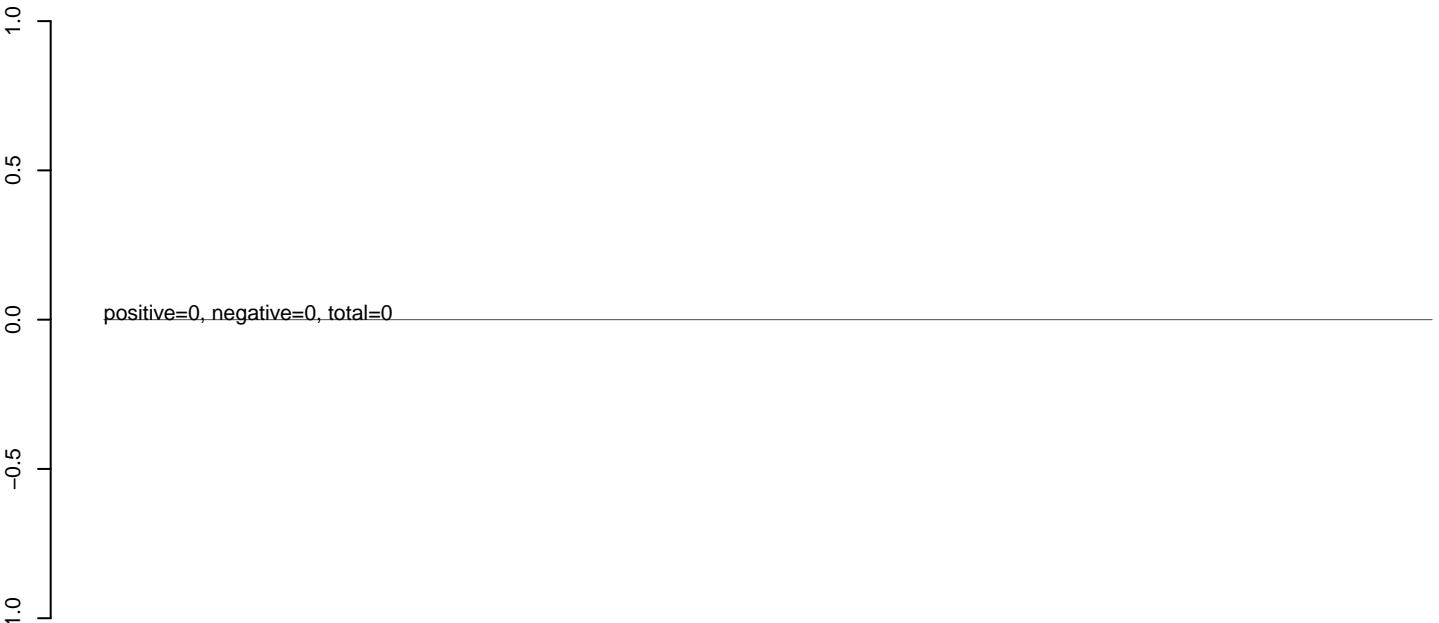
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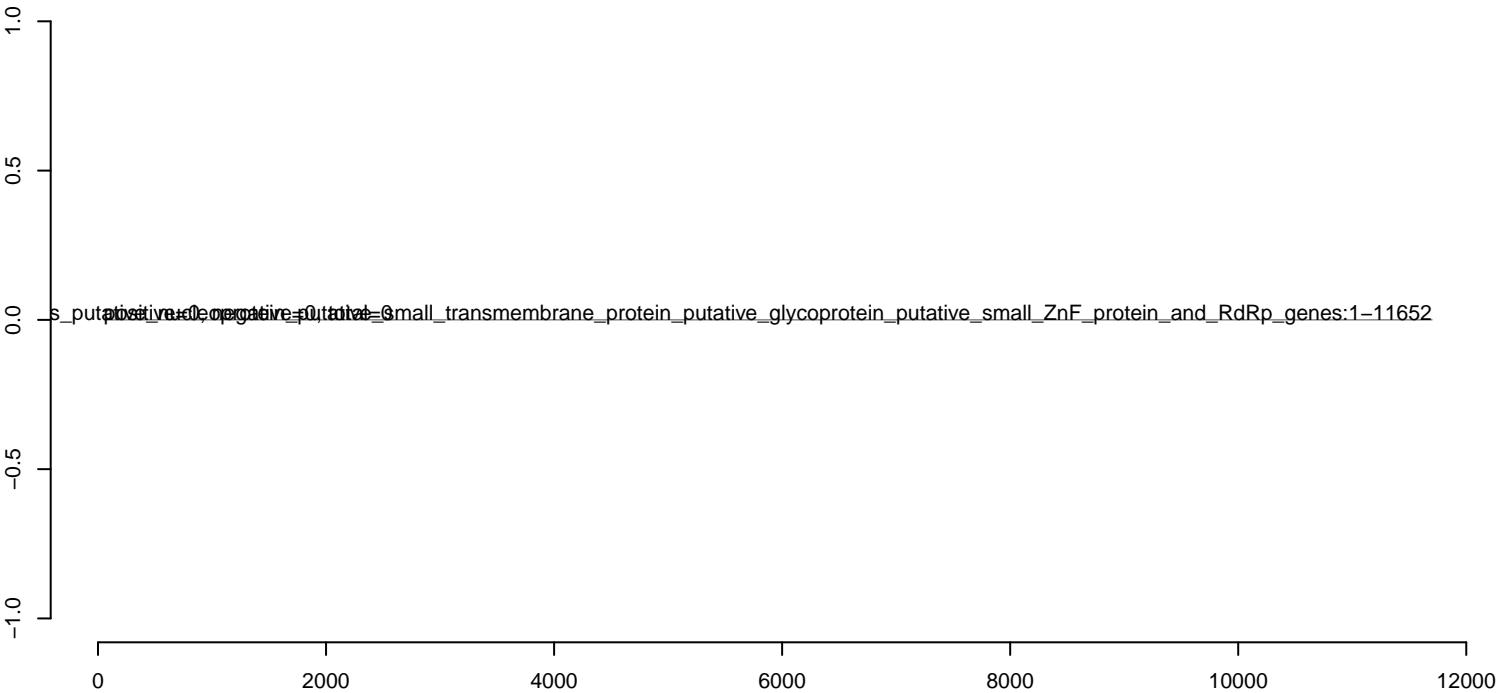
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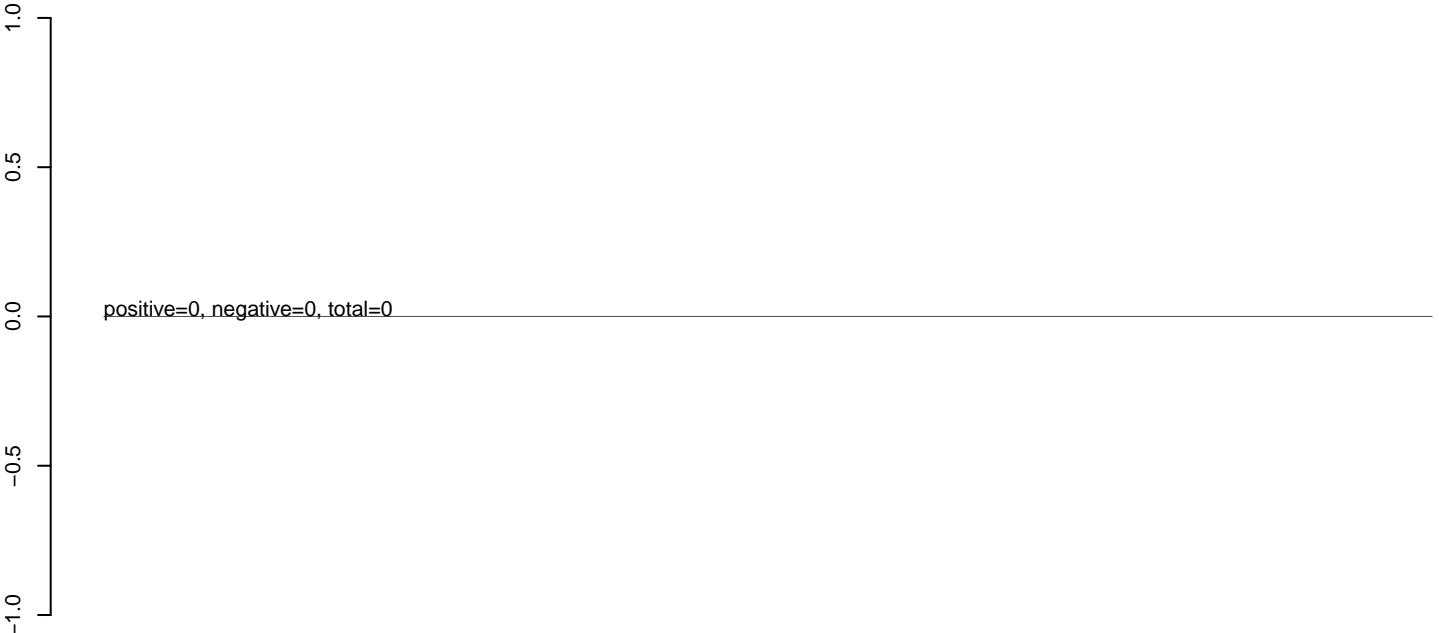
### AnGam\_Fcanc\_TN.24\_35.rep



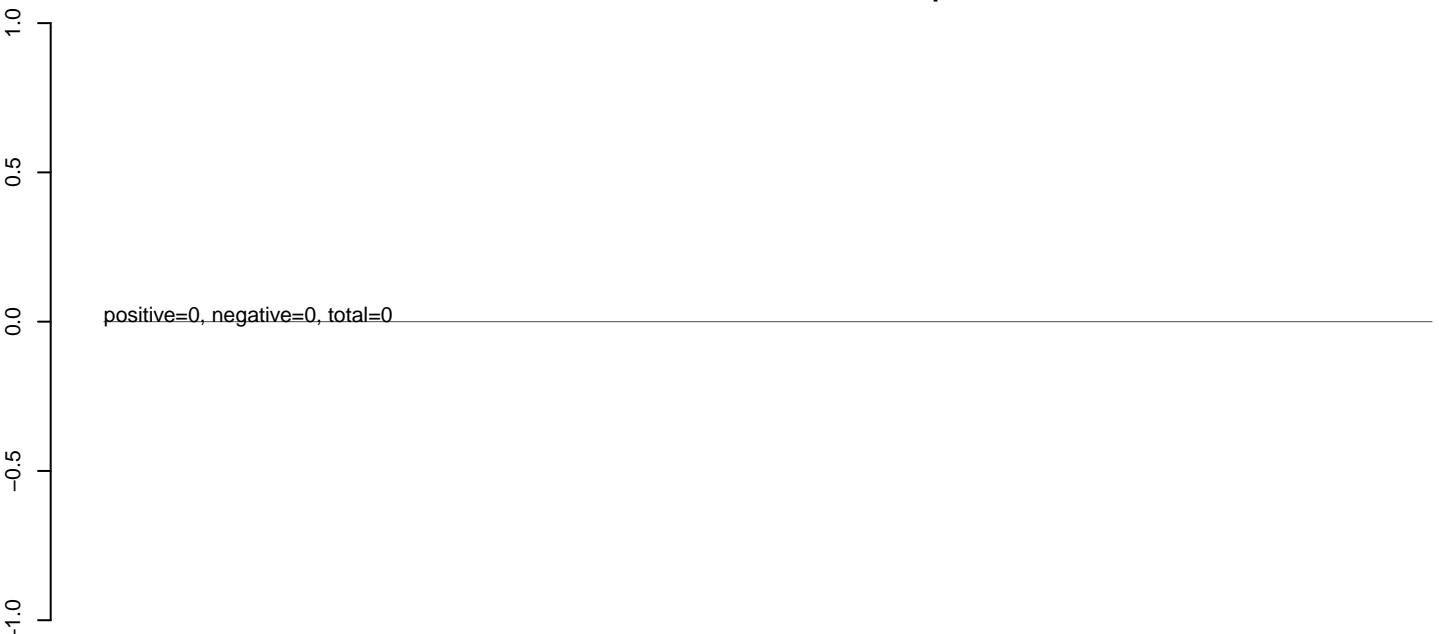
### AnGam\_Fcanc\_TN.rep



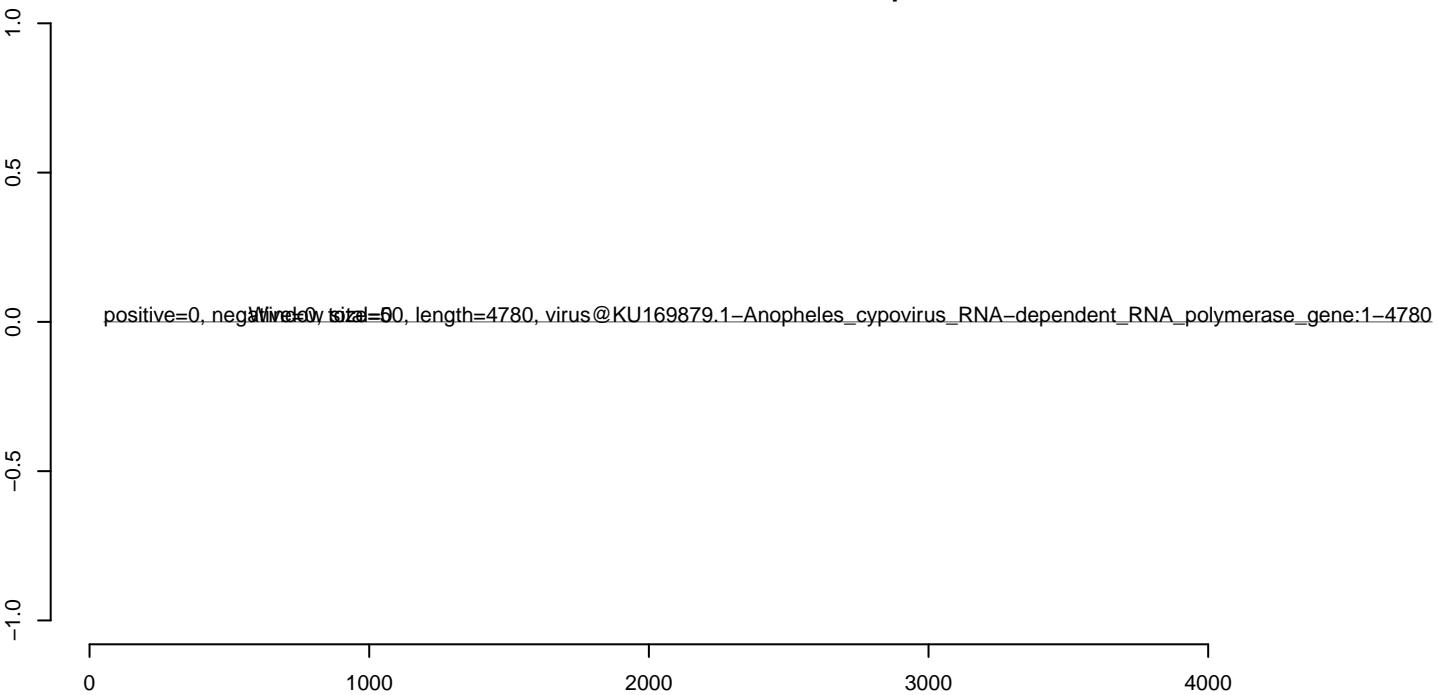
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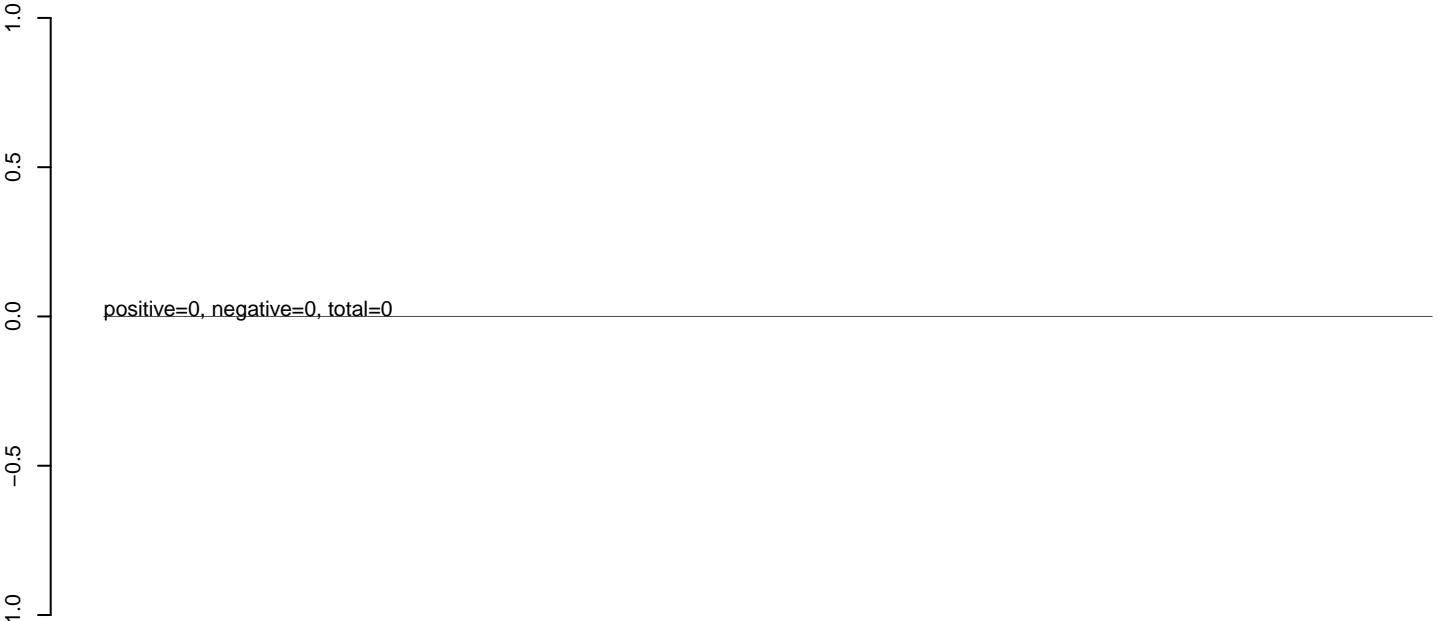
### AnGam\_Fcanc\_TN.24\_35.rep



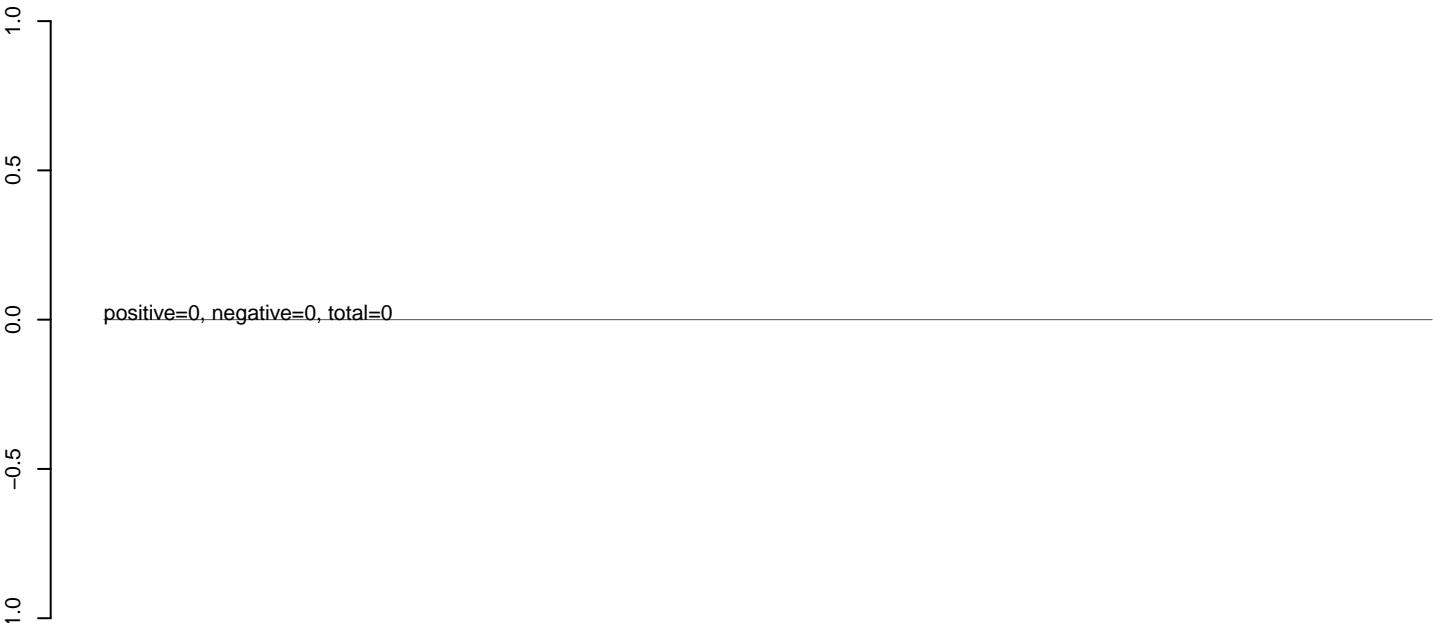
### AnGam\_Fcanc\_TN.rep



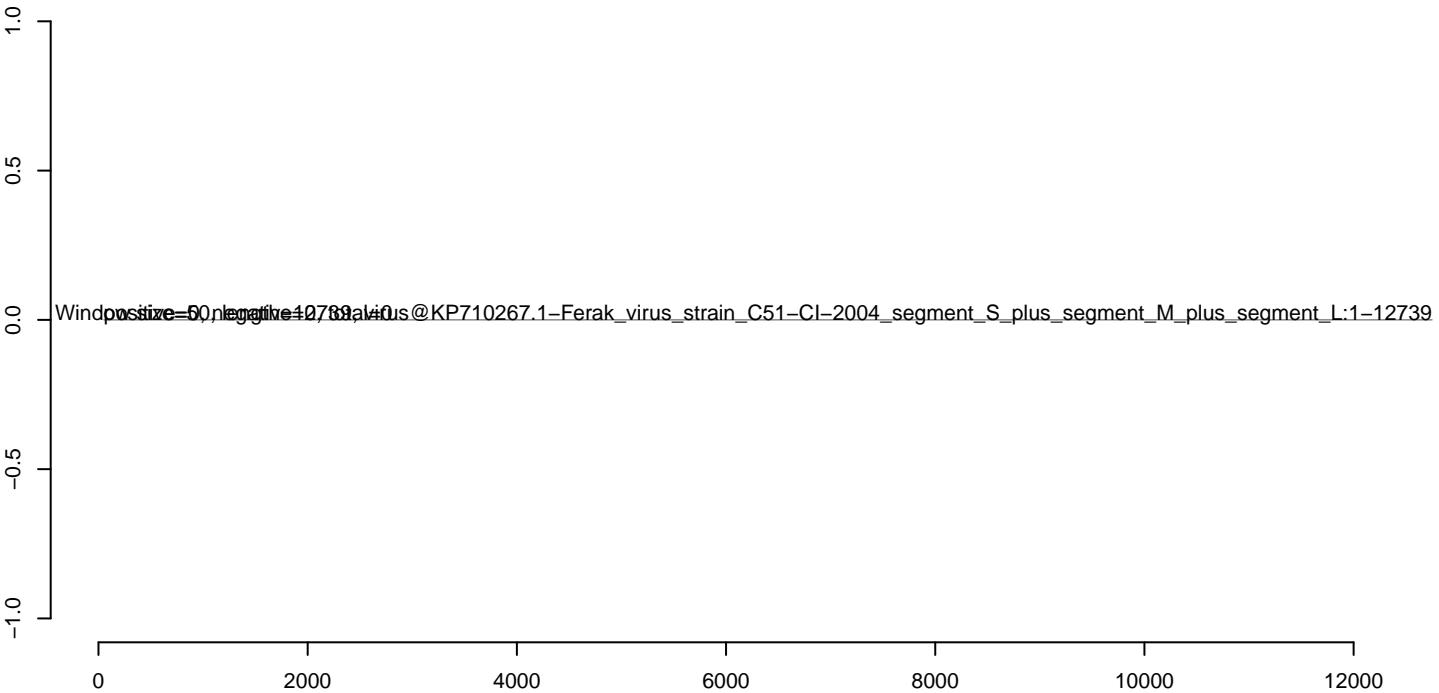
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

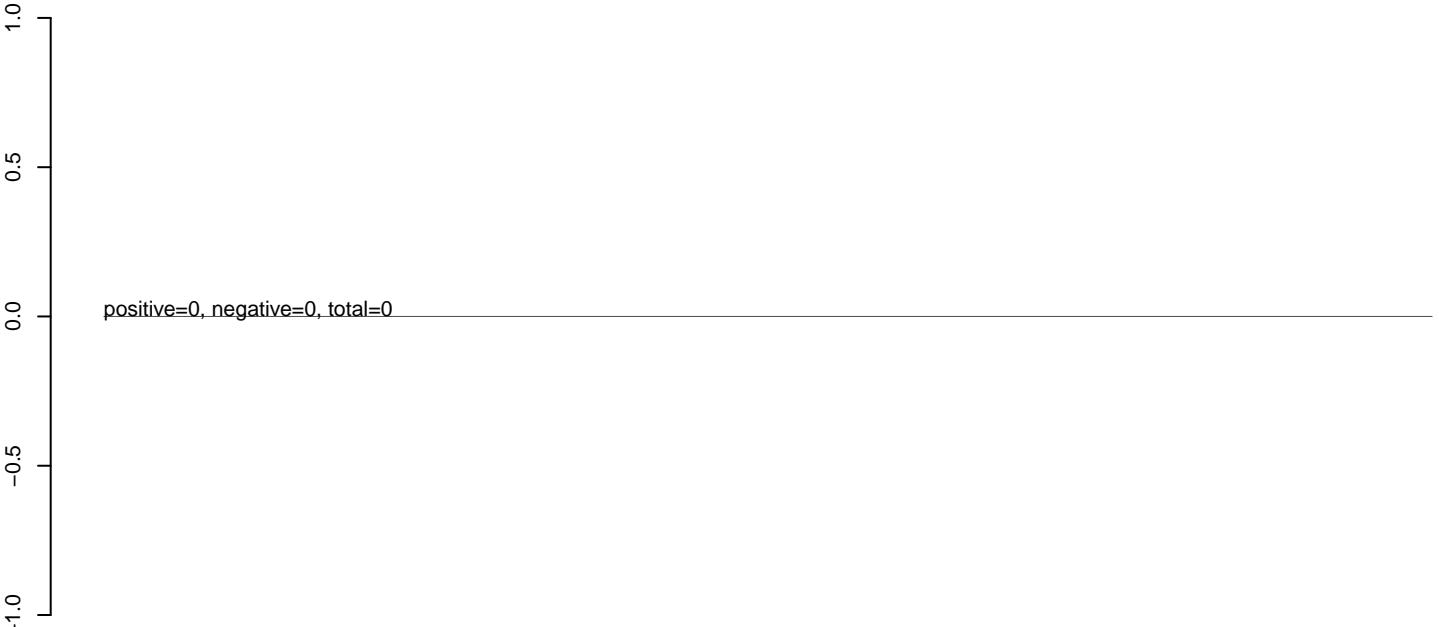
### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0

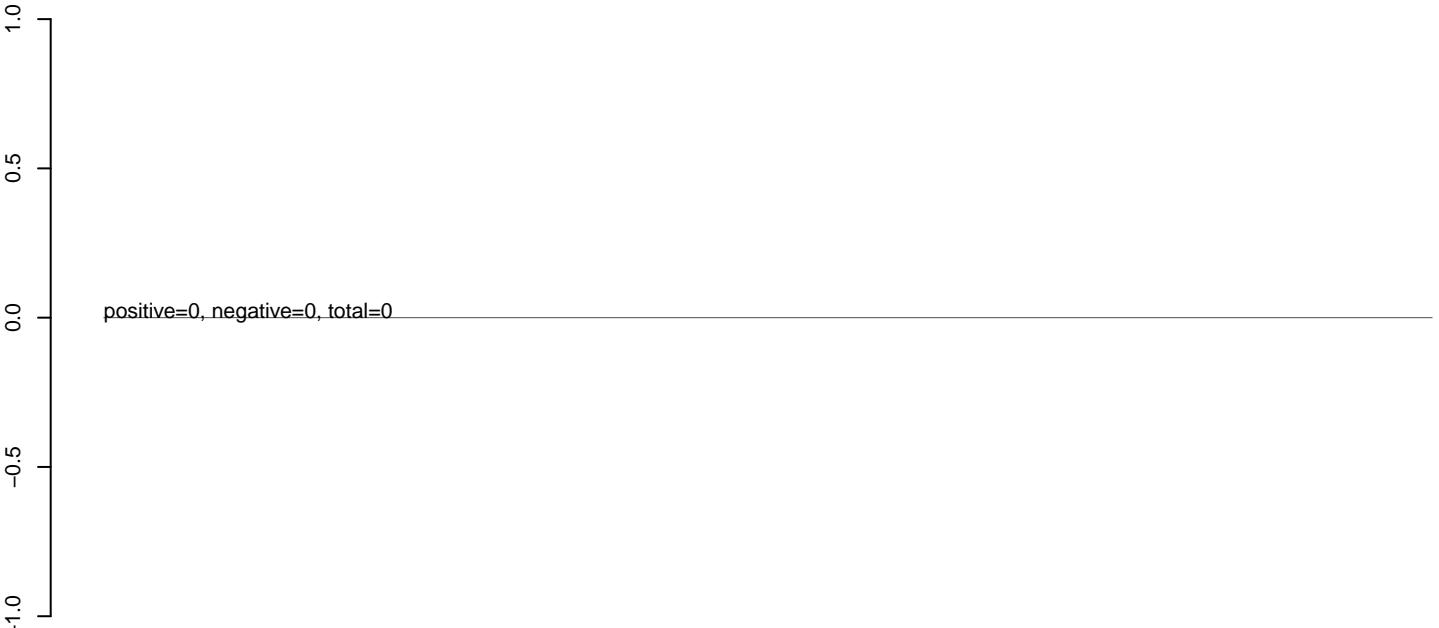
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0 2000 4000 6000 8000 10000

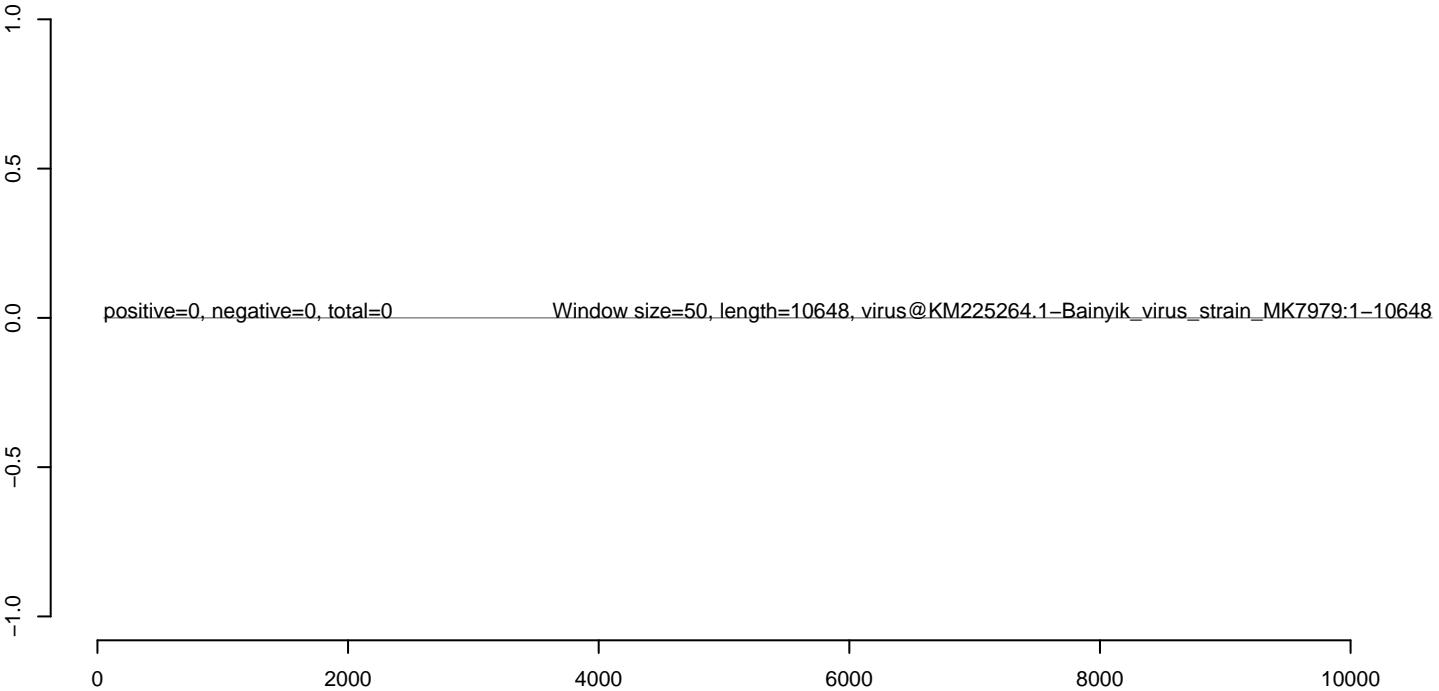
### AnGam\_Fcanc\_TN.18\_23.rep



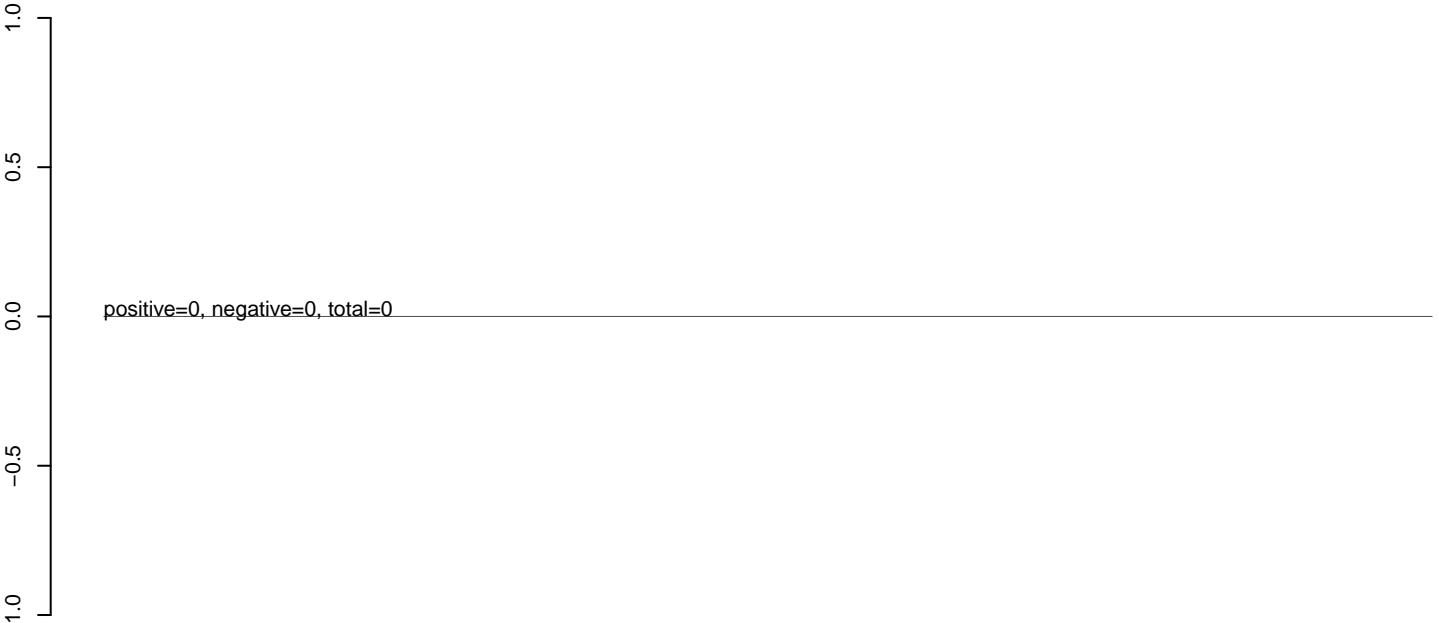
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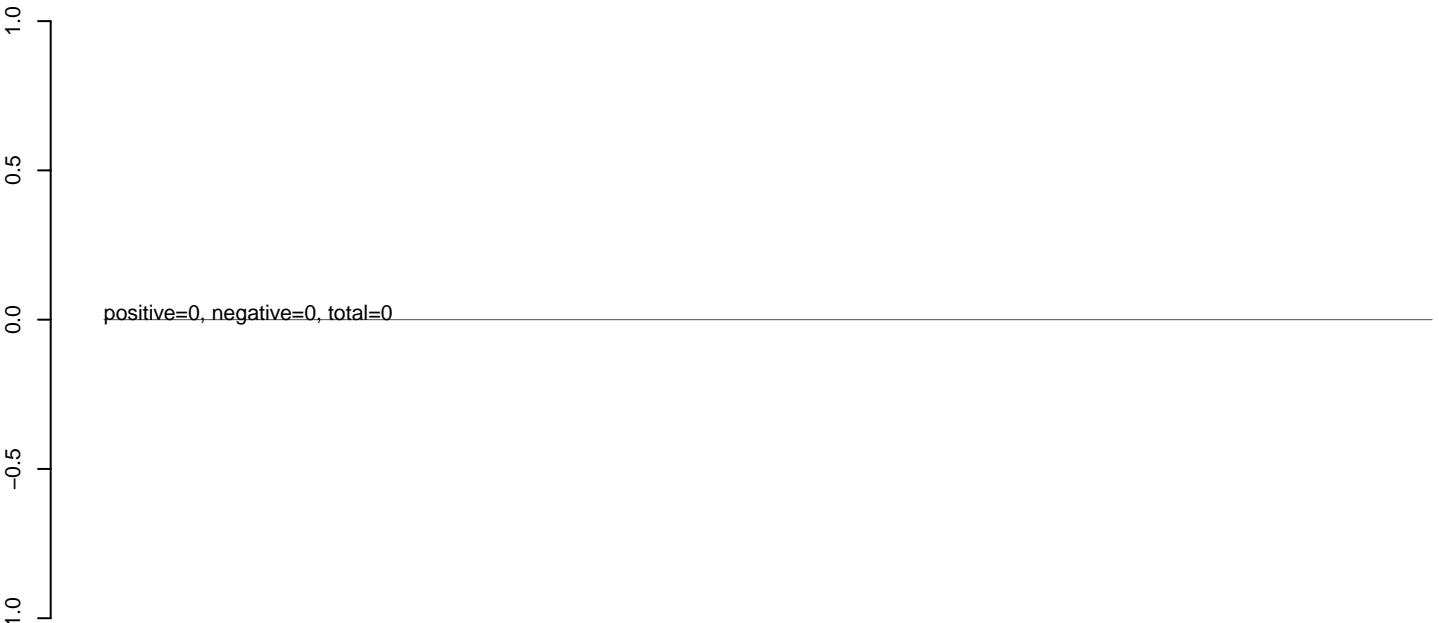
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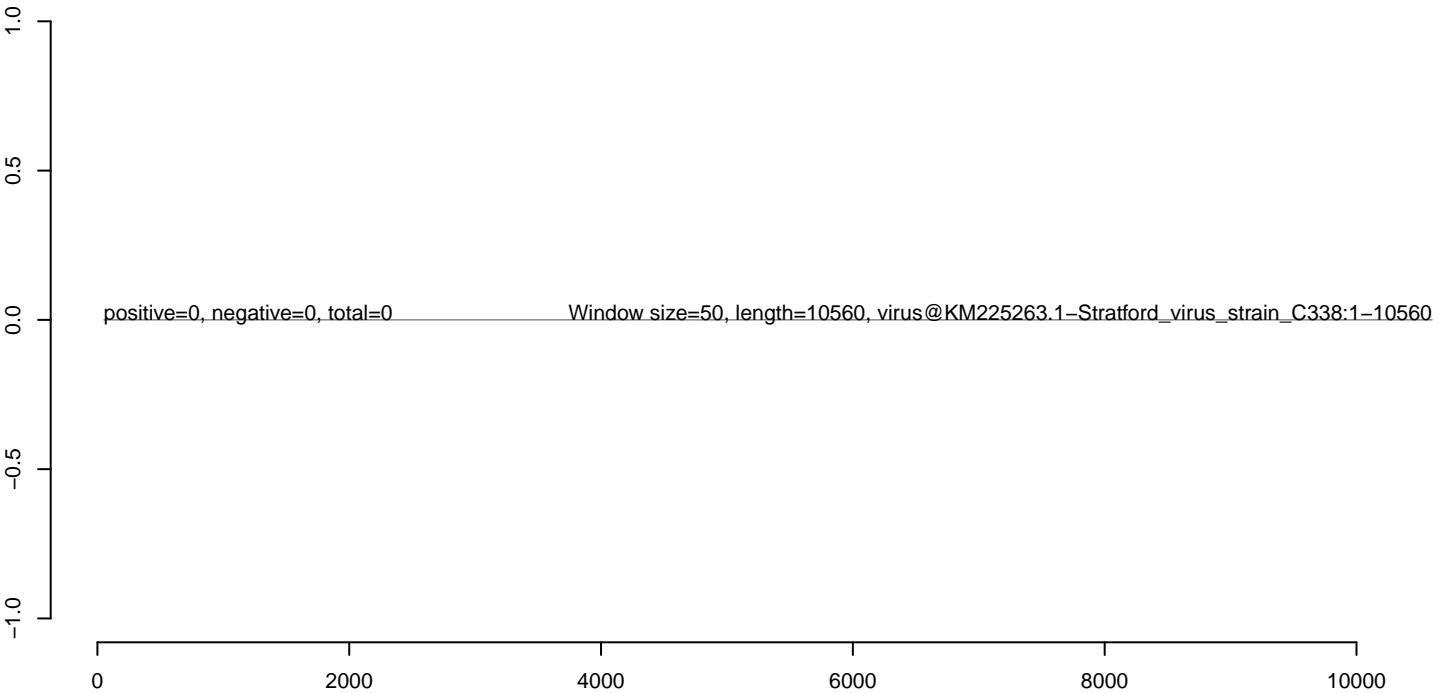
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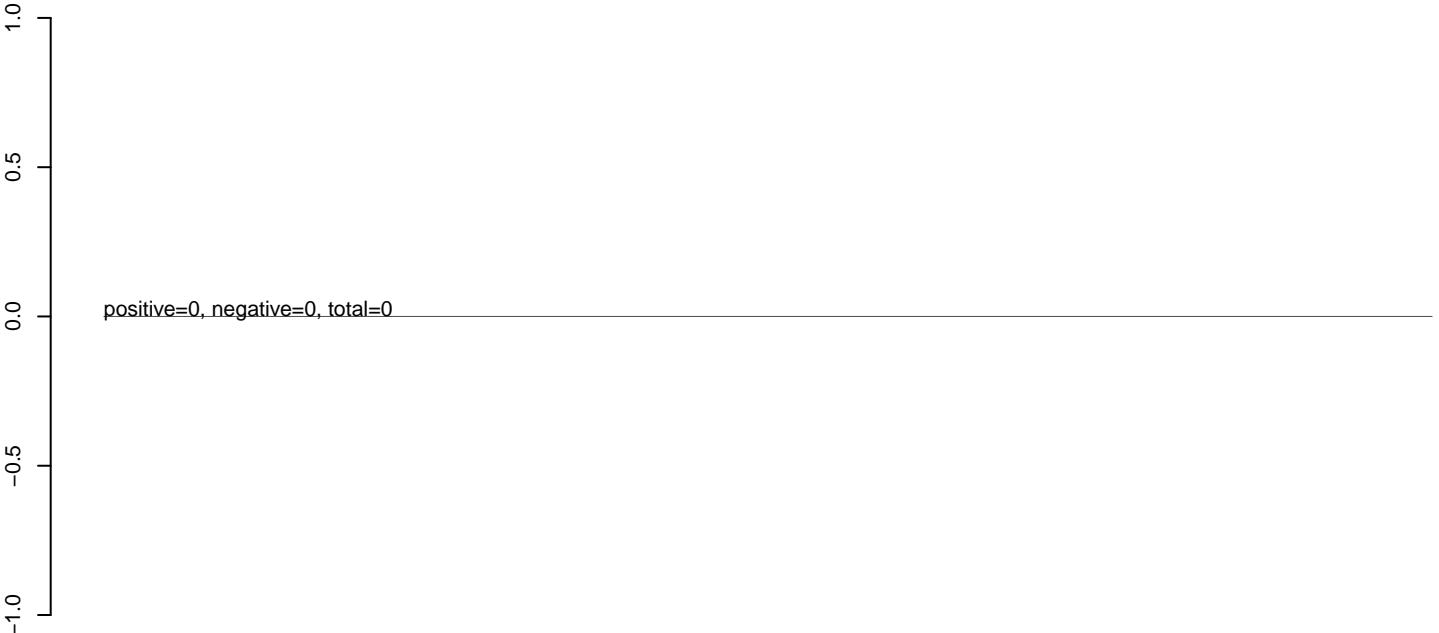
### AnGam\_Fcanc\_TN.24\_35.rep



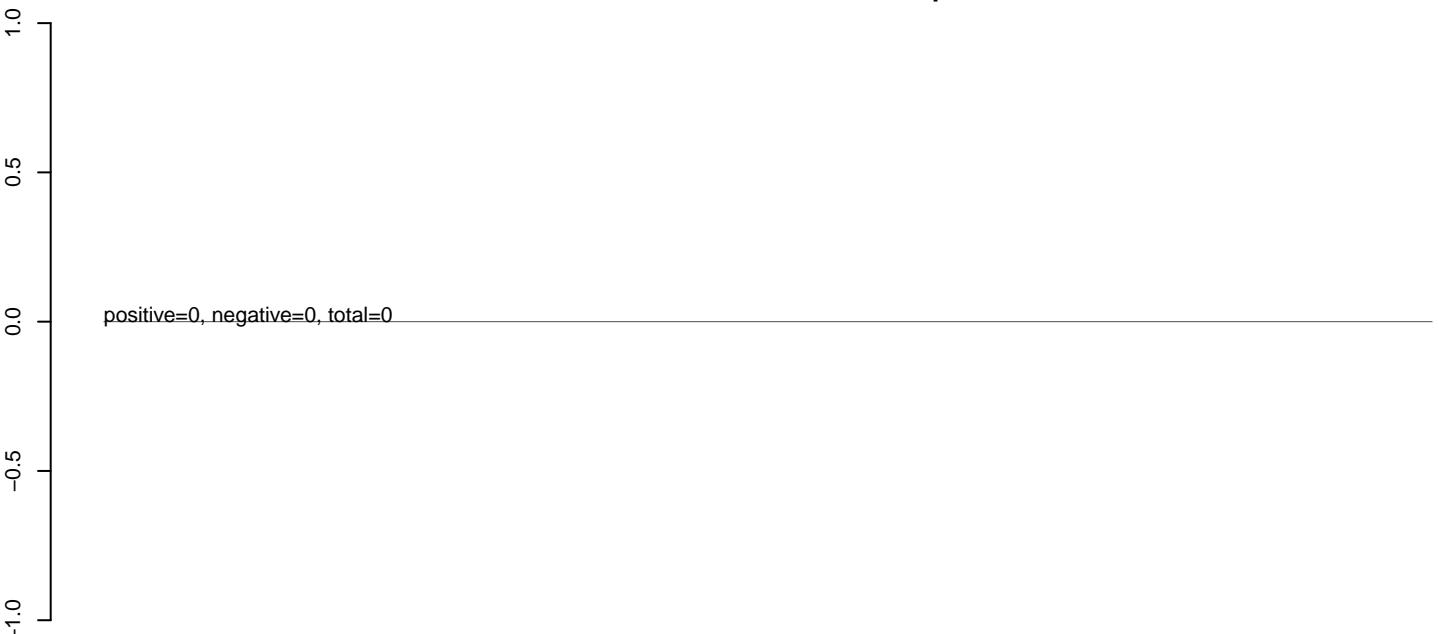
### AnGam\_Fcanc\_TN.rep



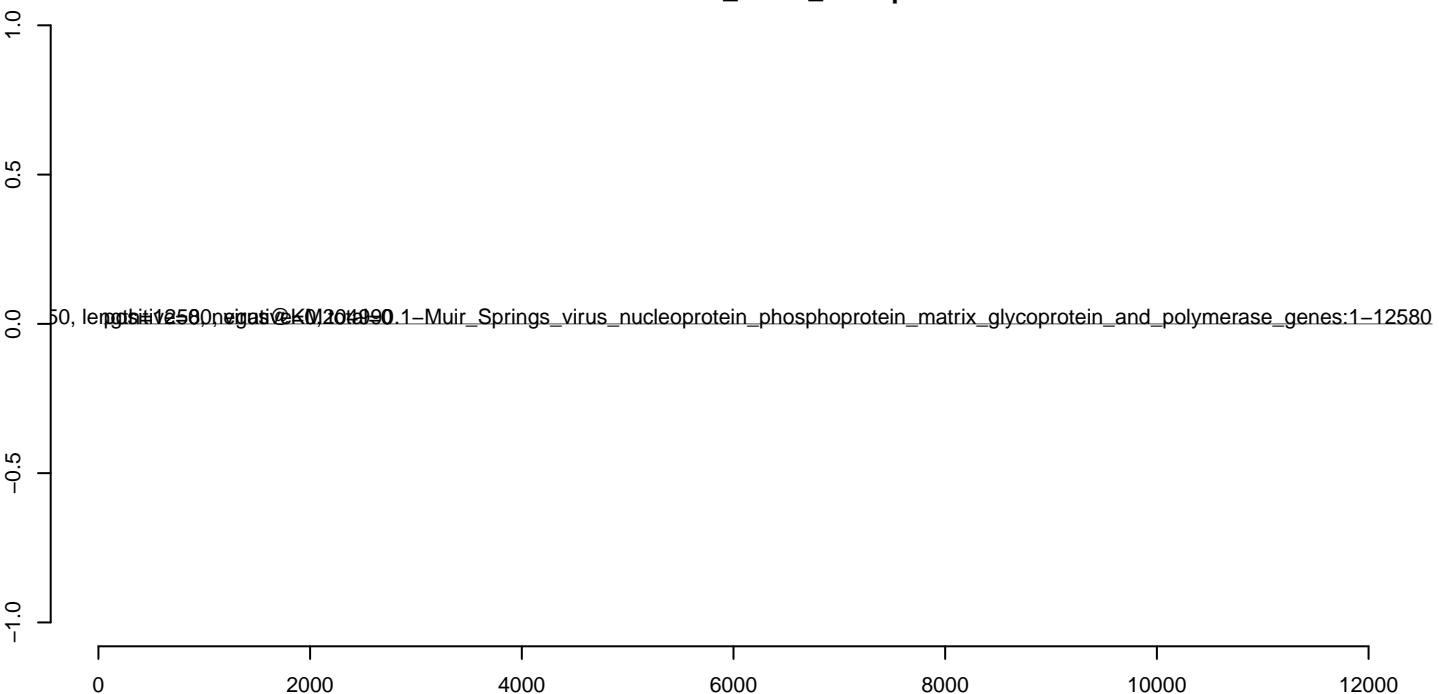
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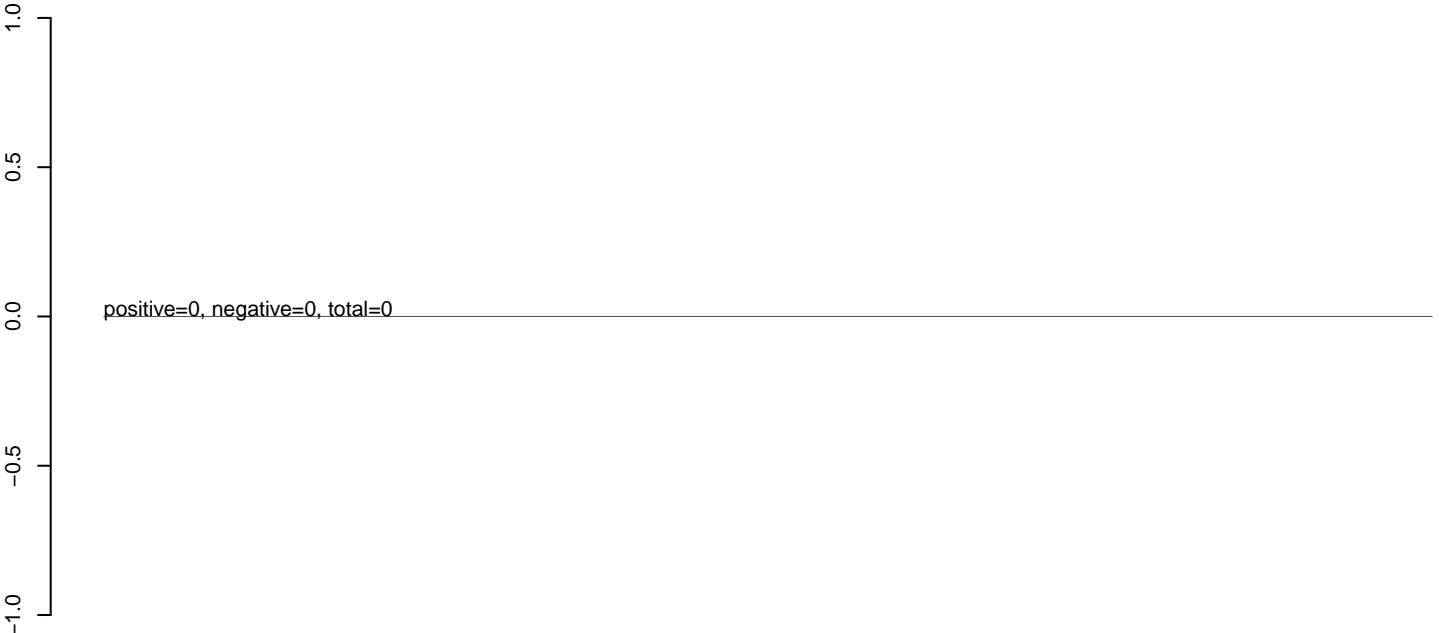
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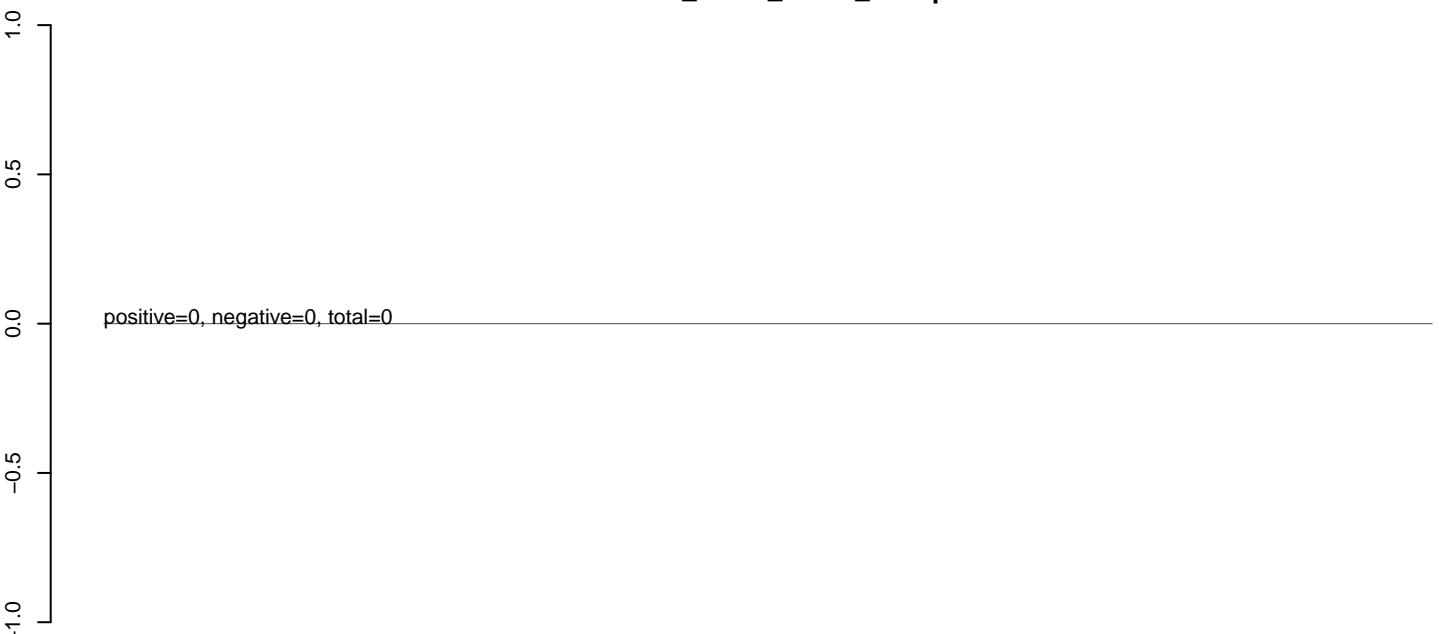
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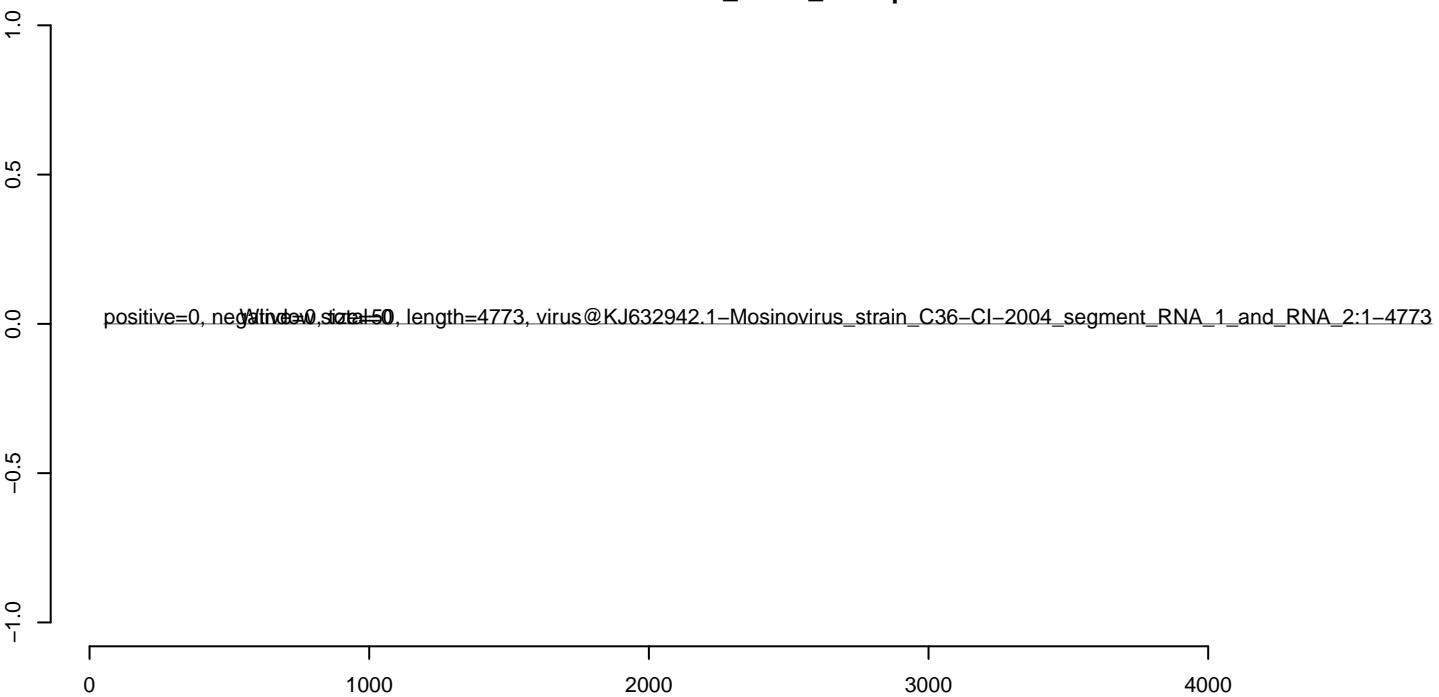
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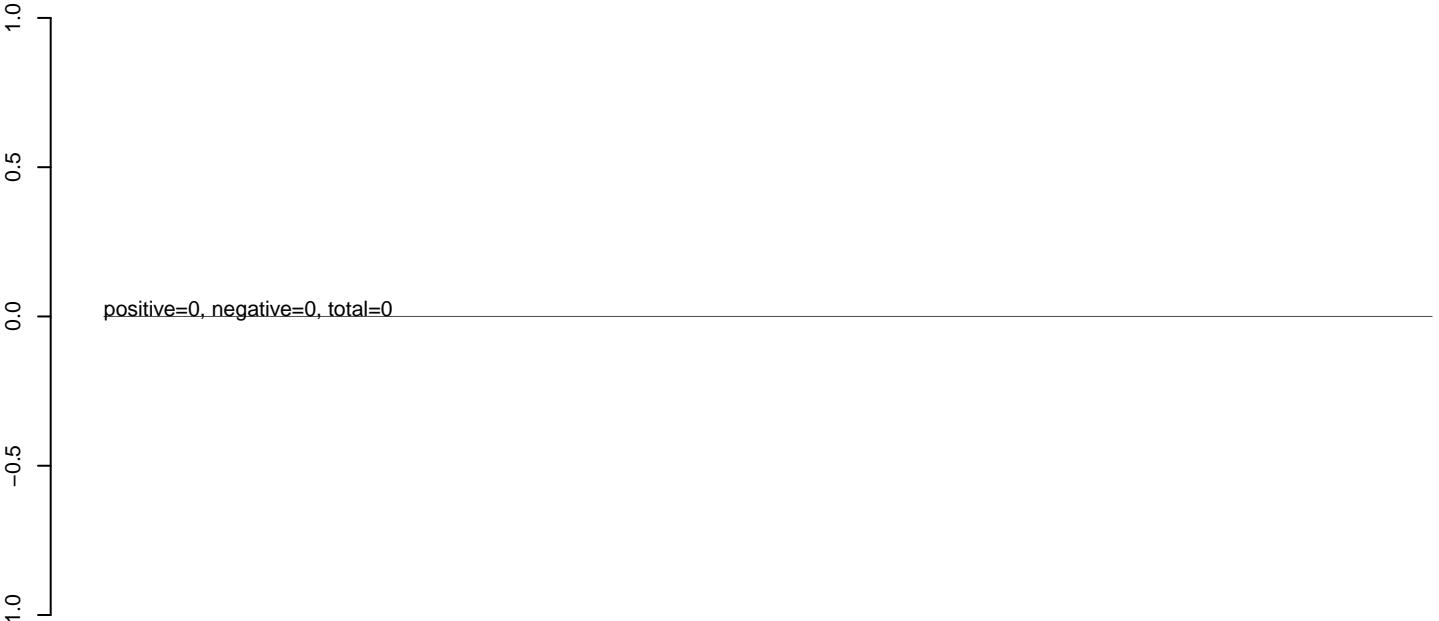
### AnGam\_Fcanc\_TN.24\_35.rep



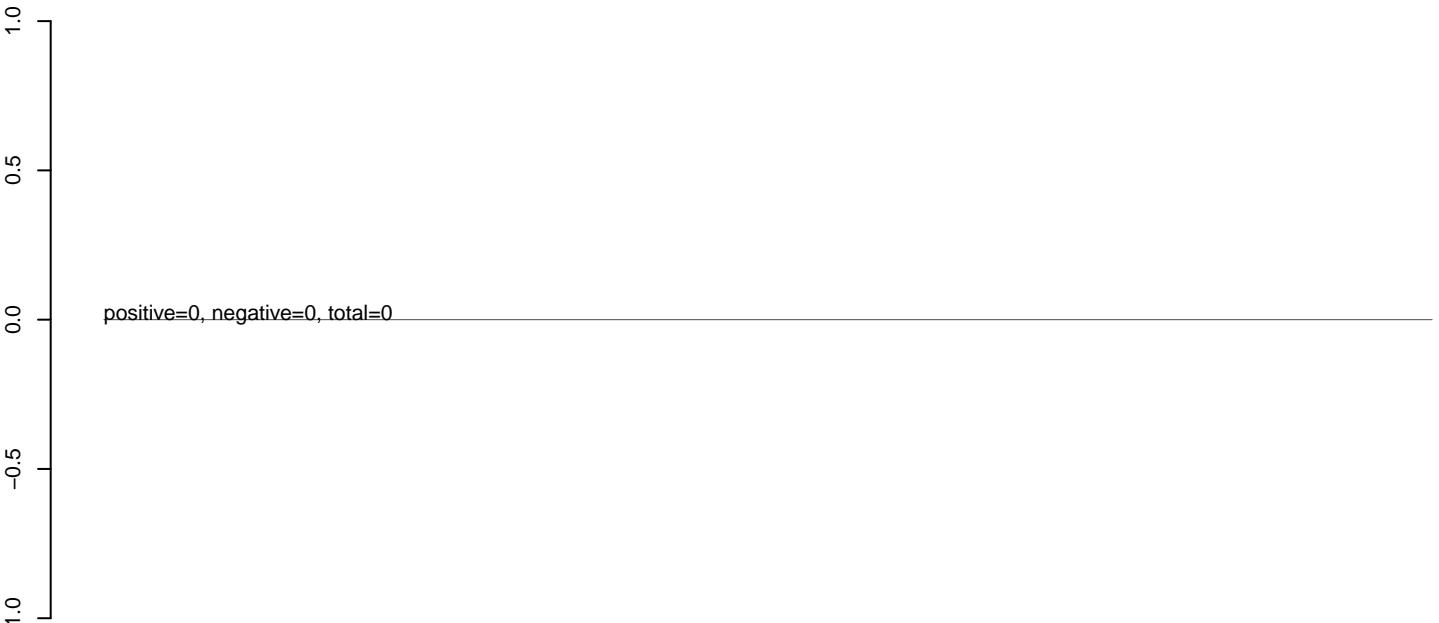
### AnGam\_Fcanc\_TN.rep



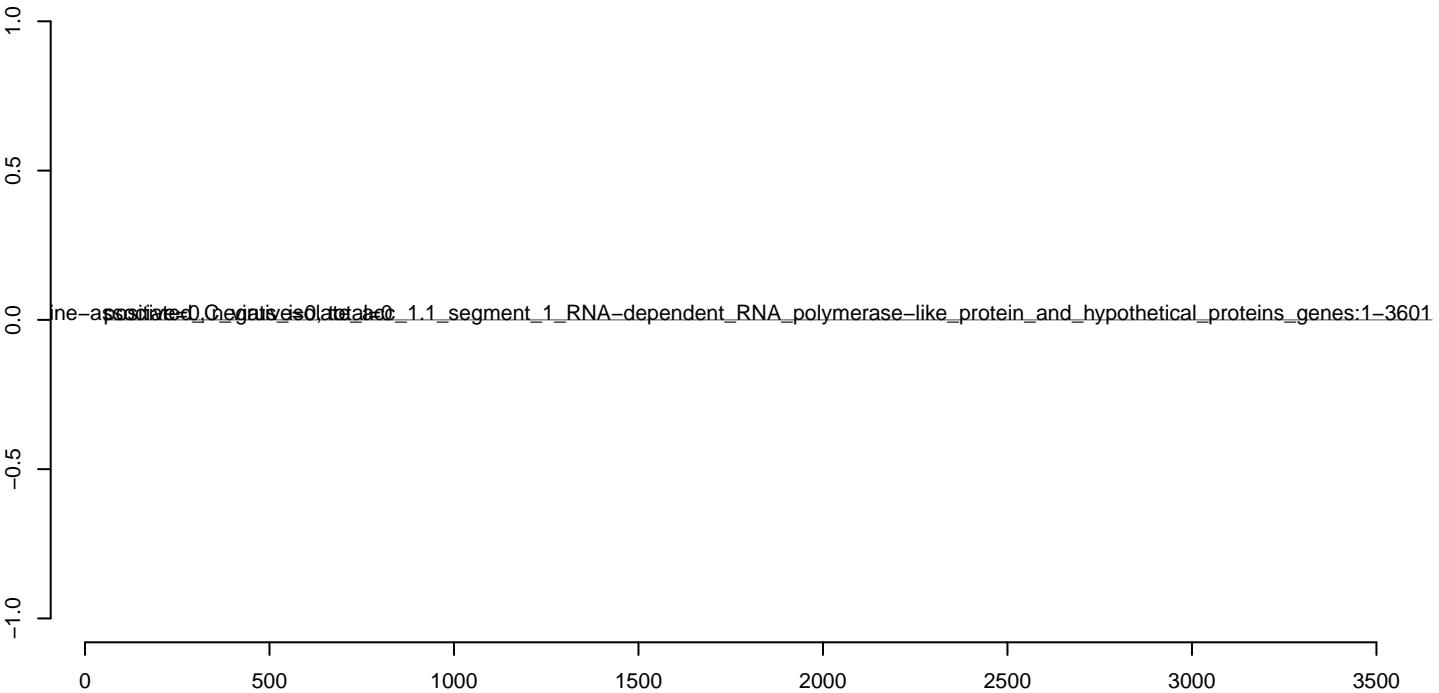
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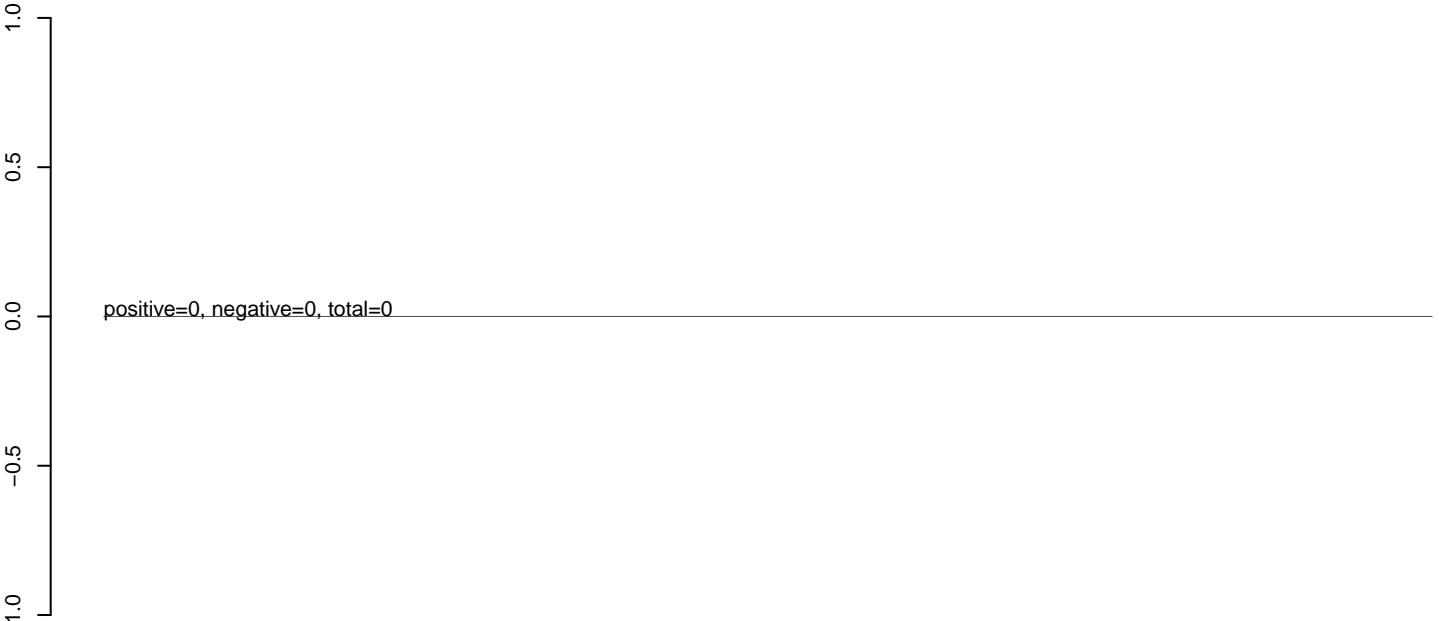
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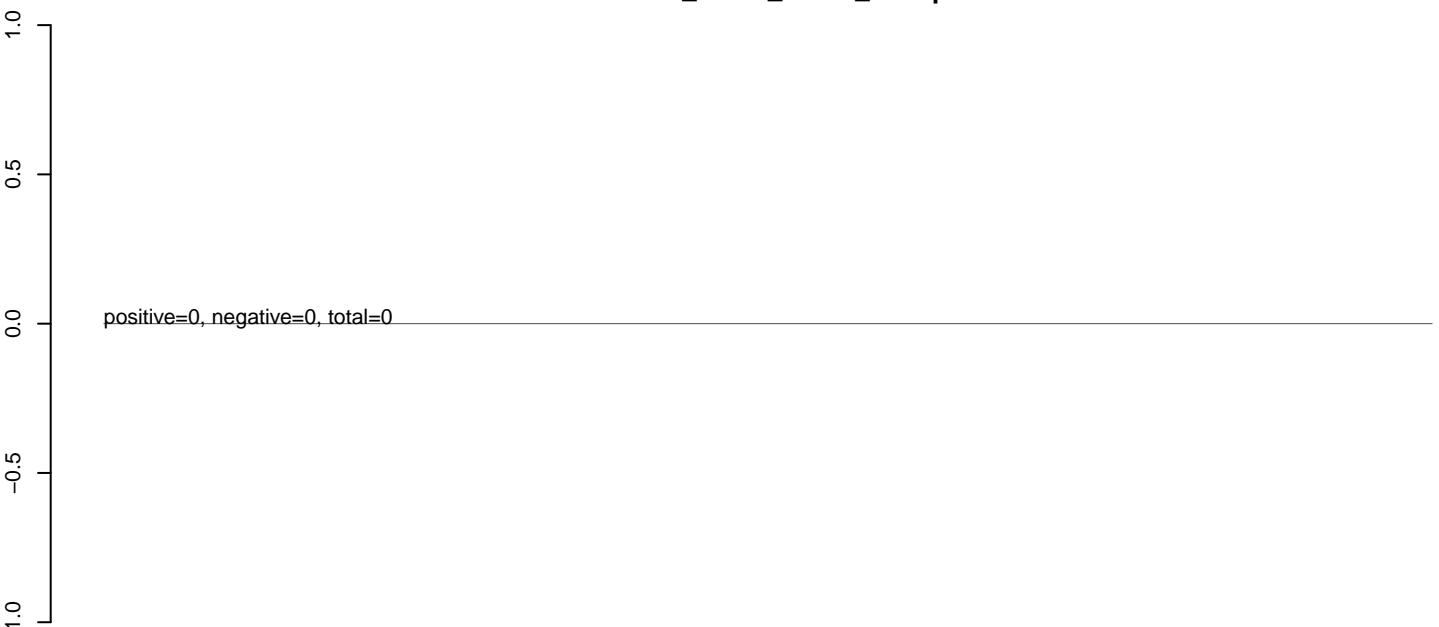
### AnGam\_Fcanc\_TN.rep



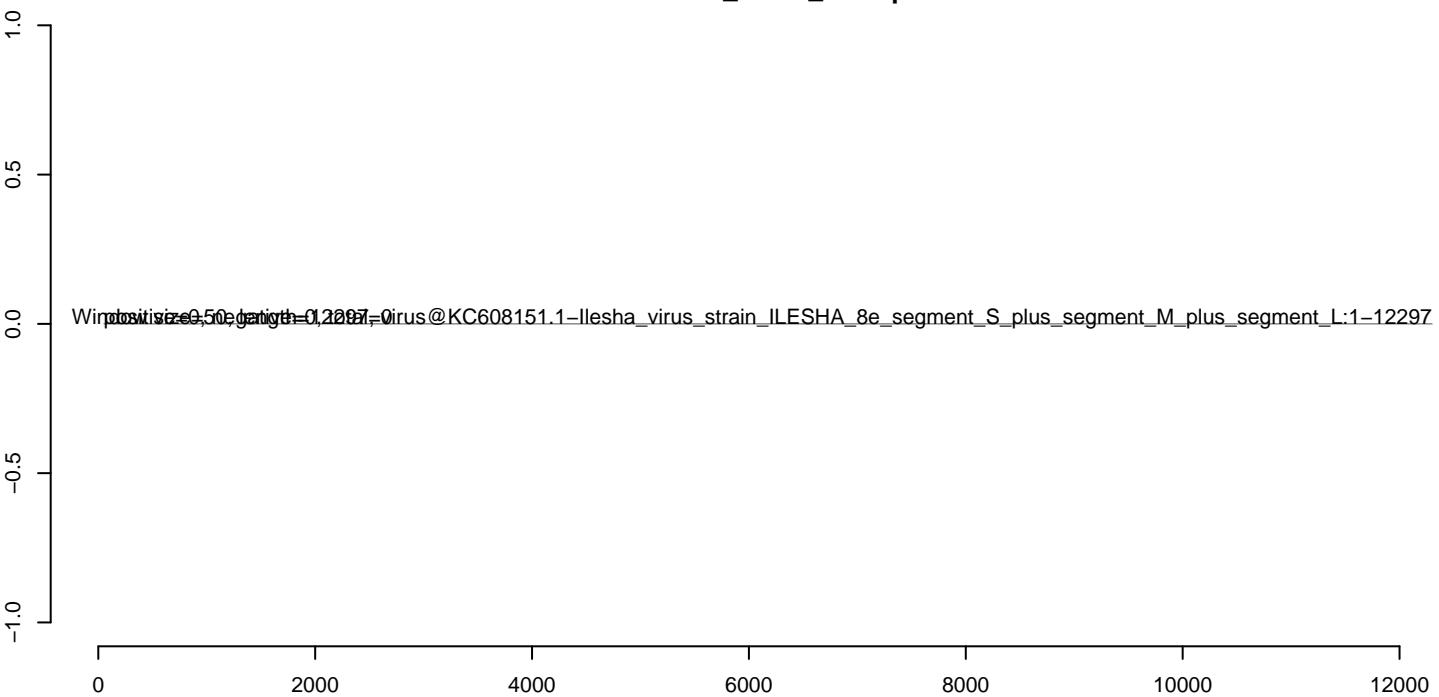
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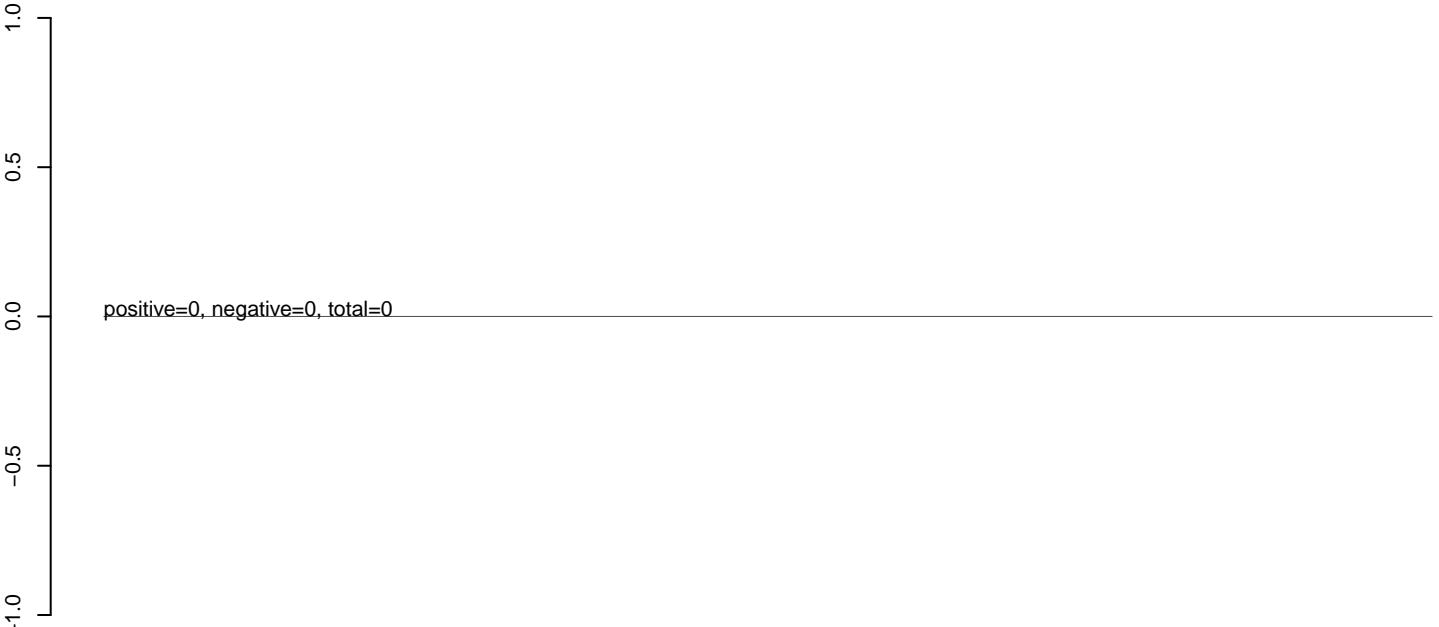
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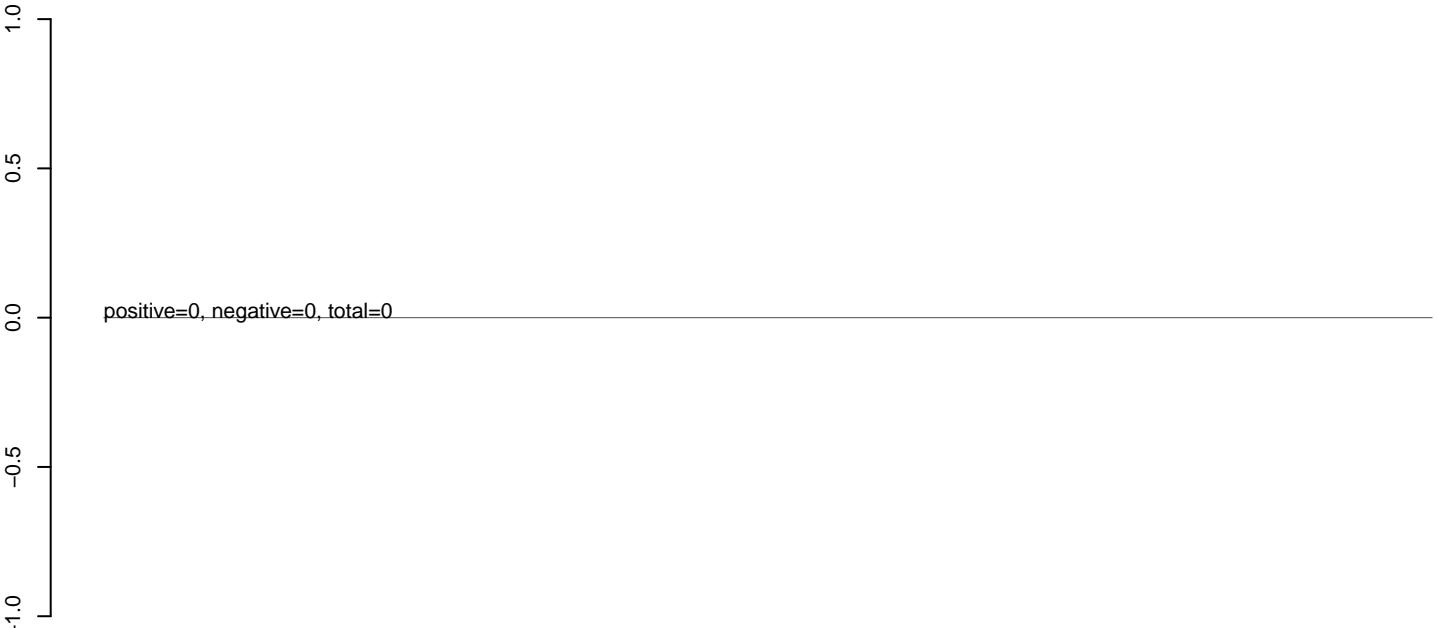
### AnGam\_Fcanc\_TN.rep



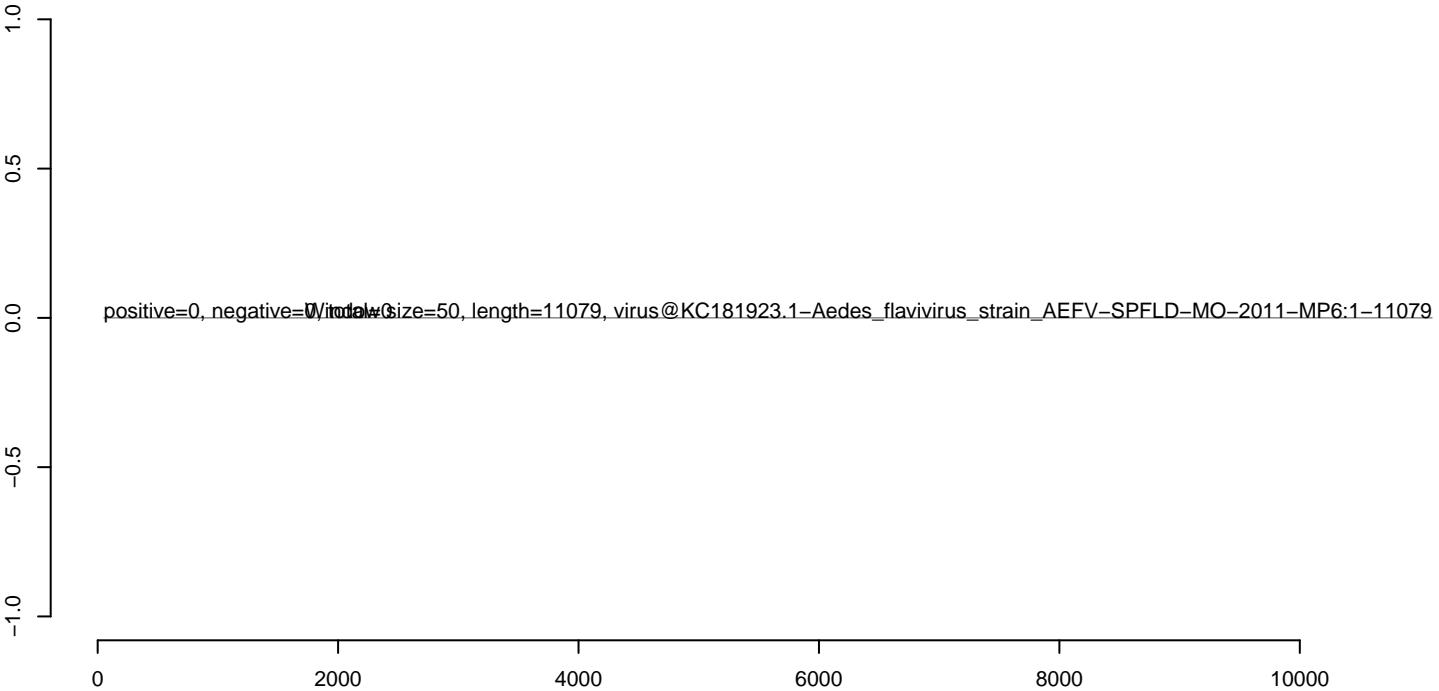
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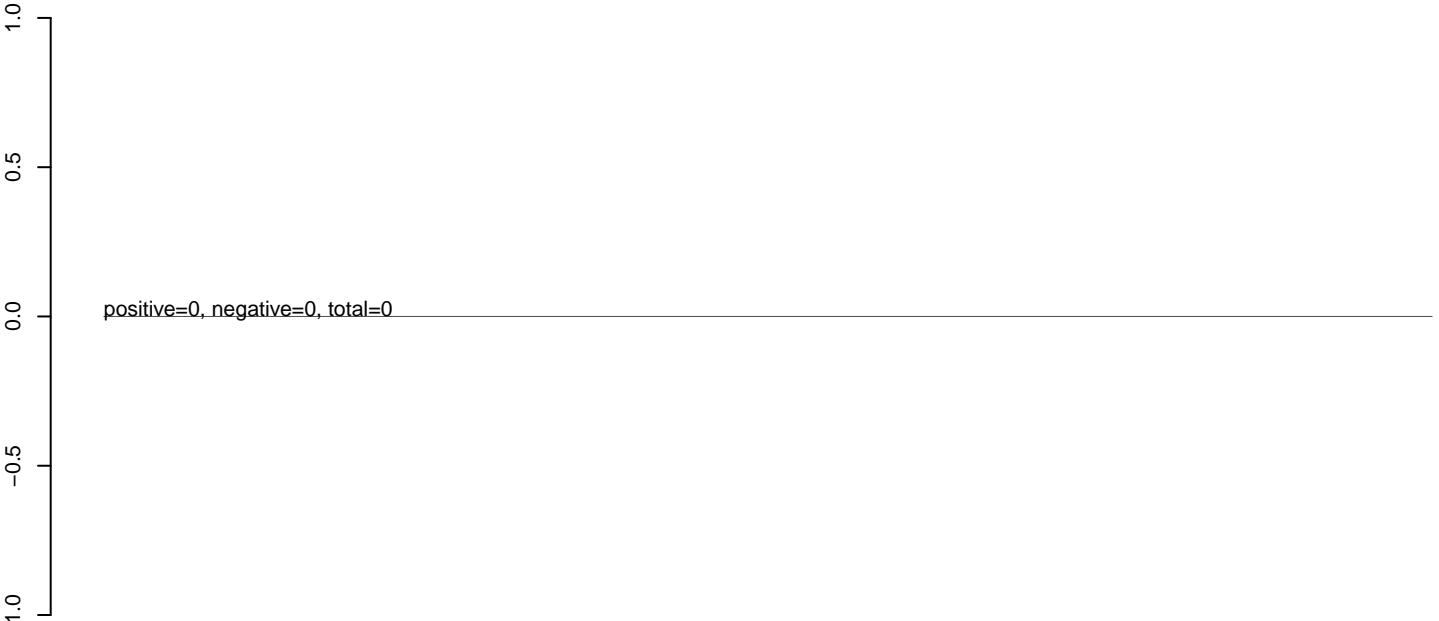
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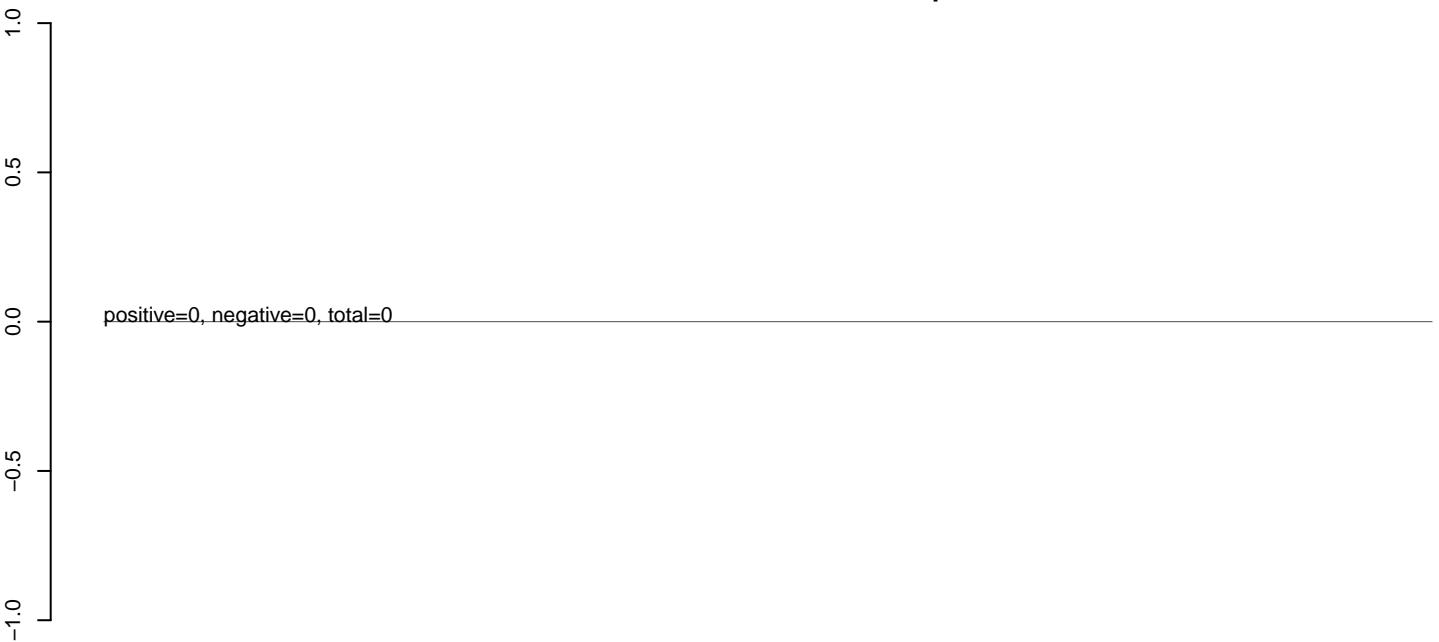
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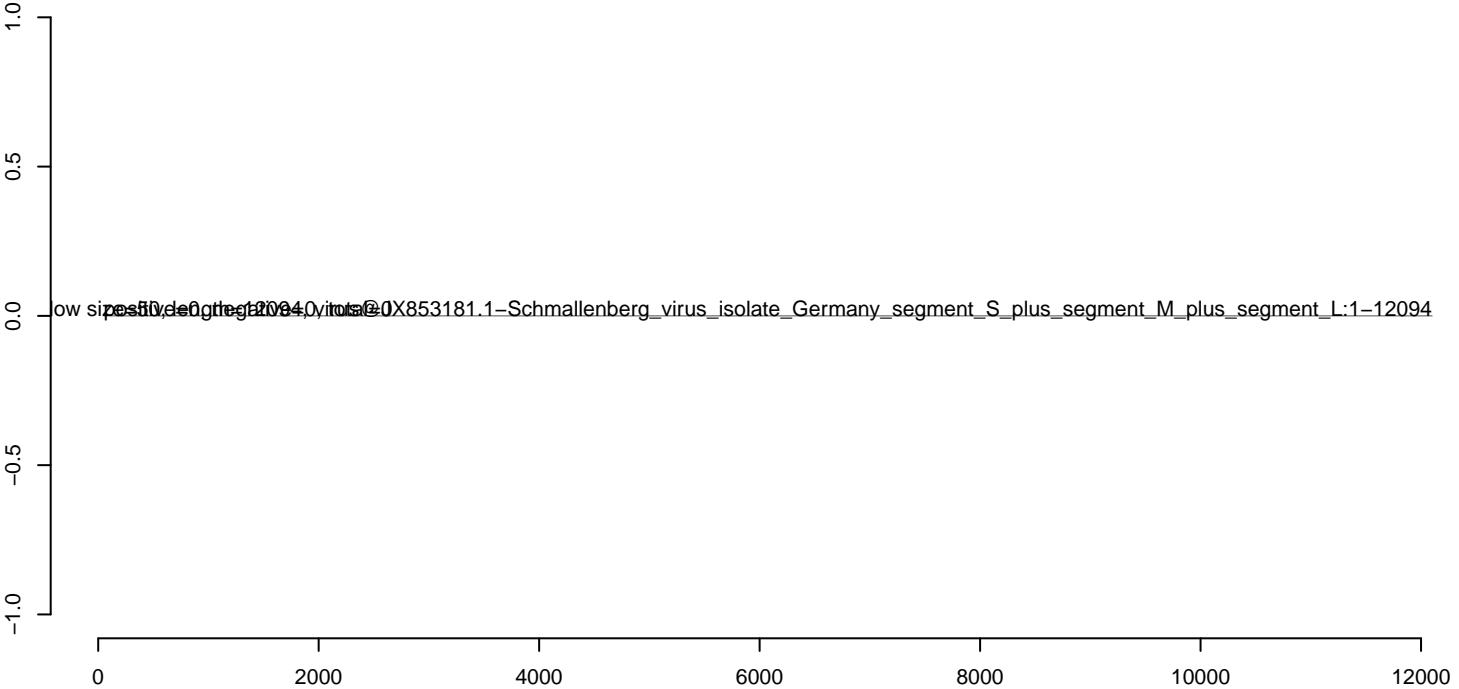
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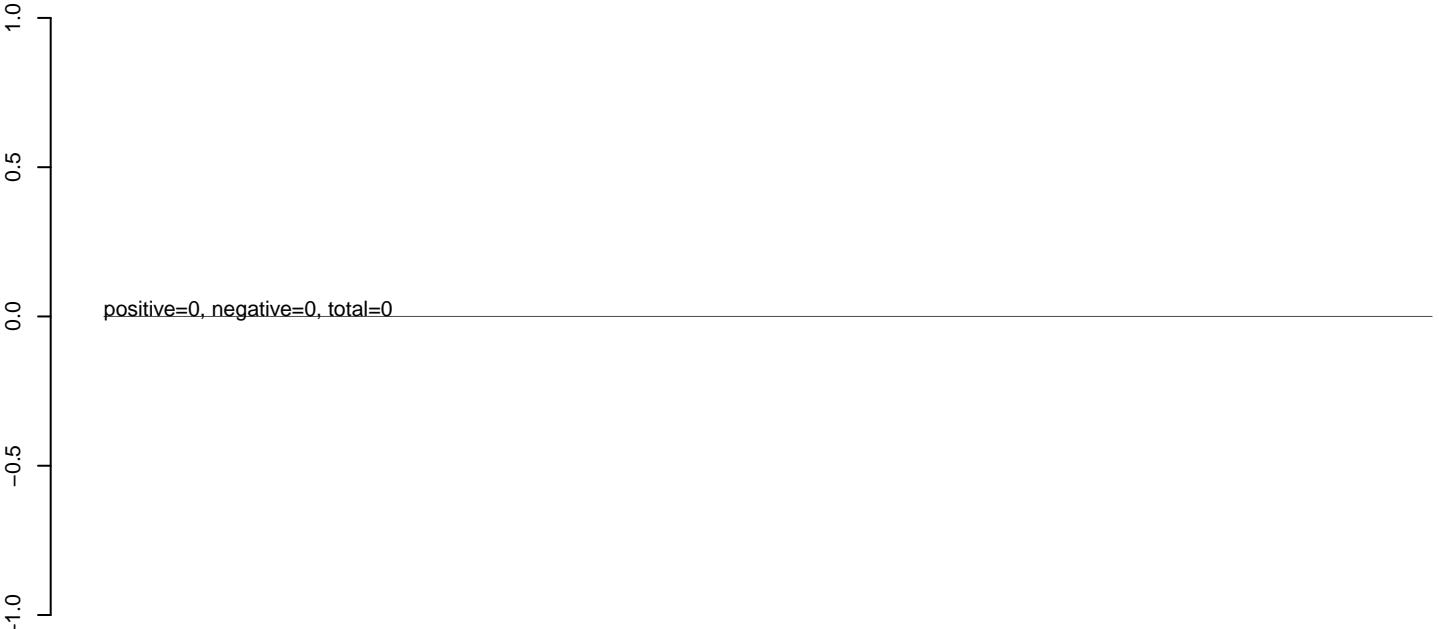
### AnGam\_Fcanc\_TN.24\_35.rep



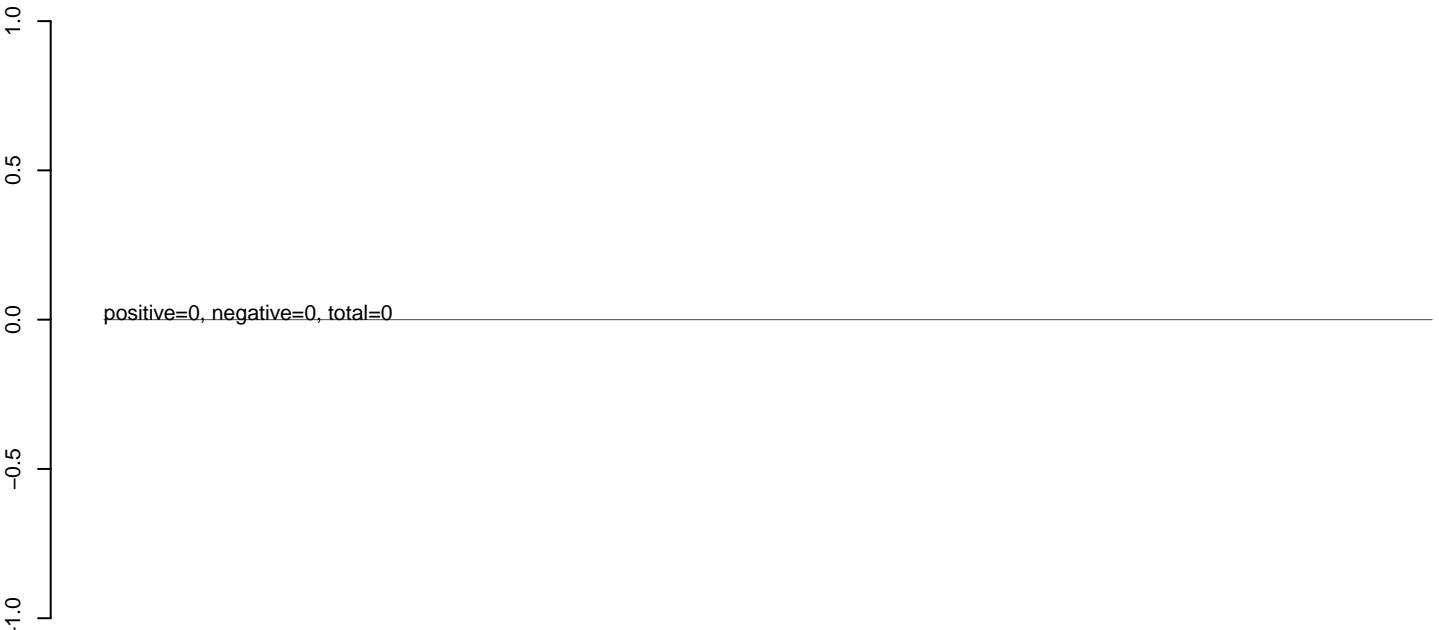
### AnGam\_Fcanc\_TN.rep



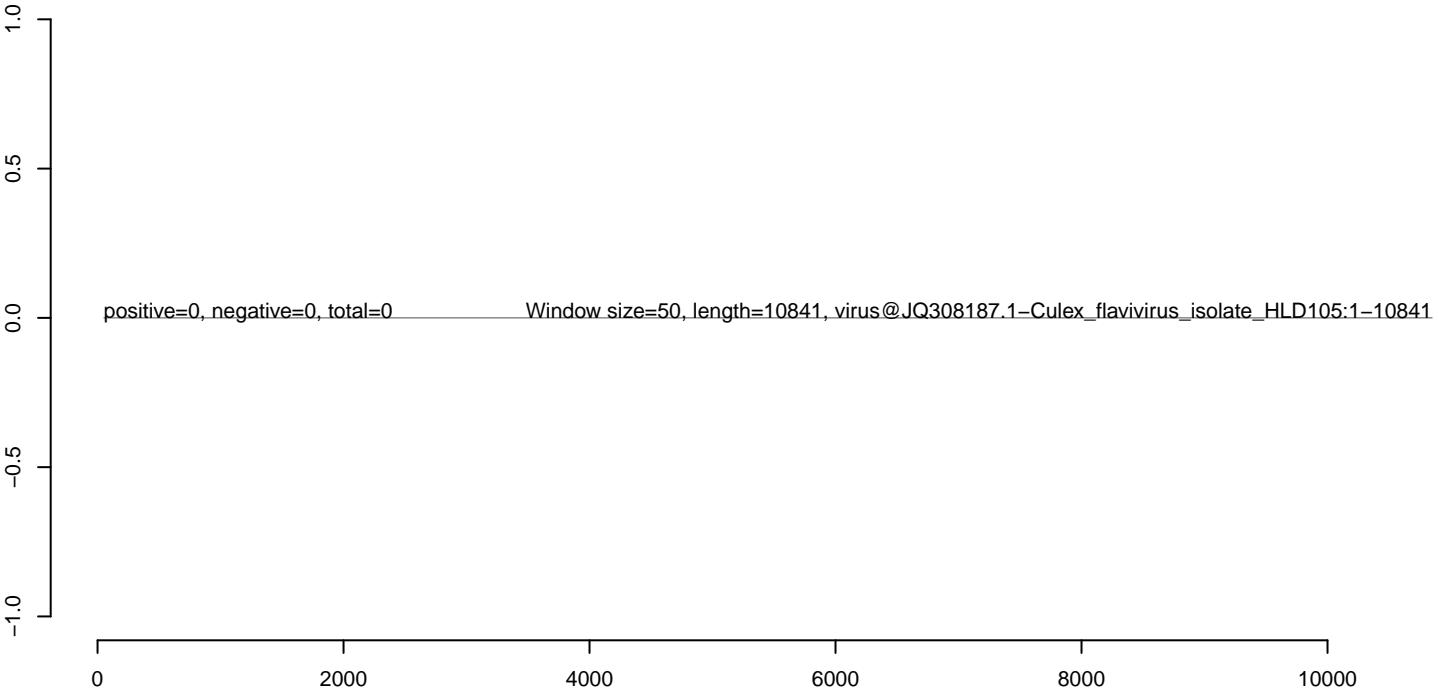
### AnGam\_Fcanc\_TN.18\_23.rep



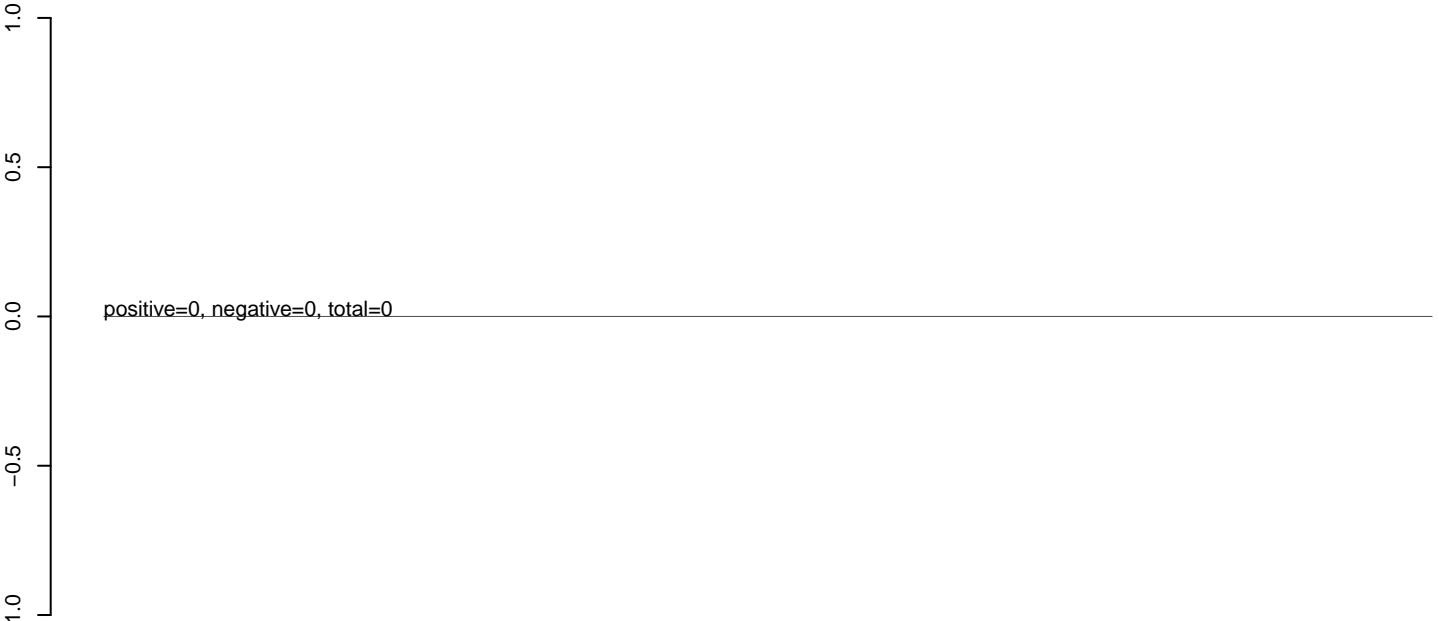
### AnGam\_Fcanc\_TN.24\_35.rep



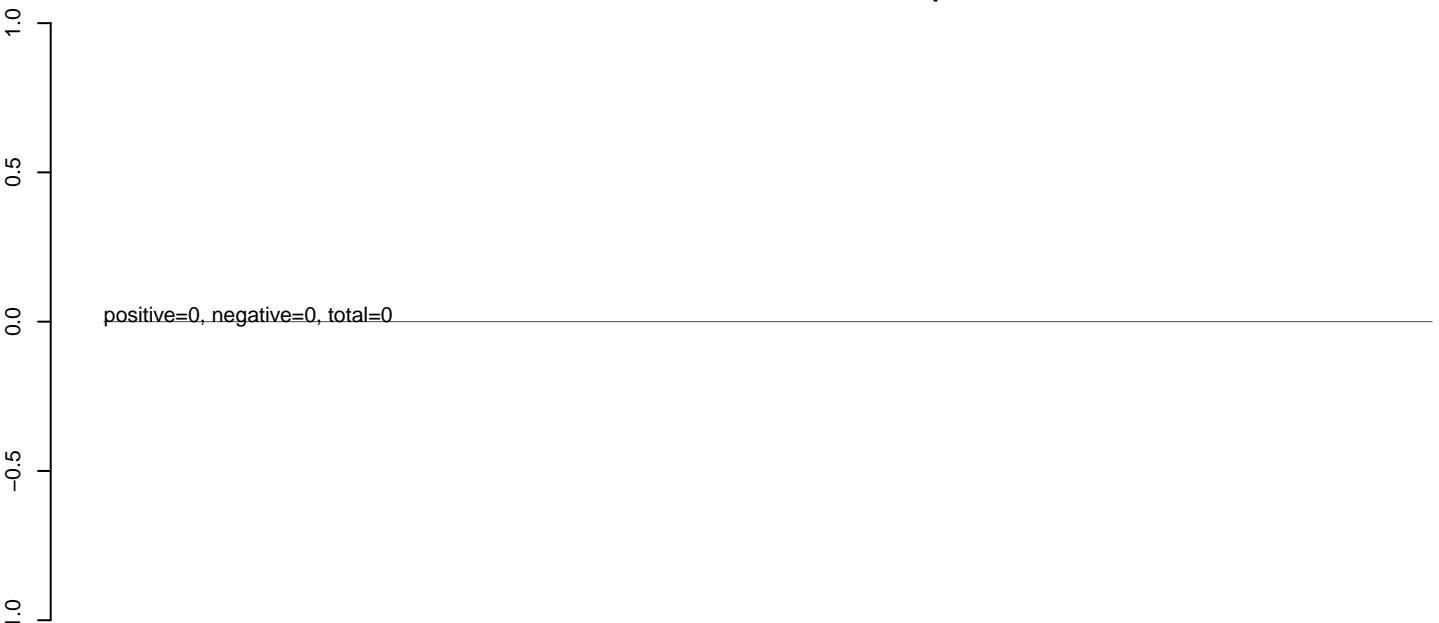
### AnGam\_Fcanc\_TN.rep



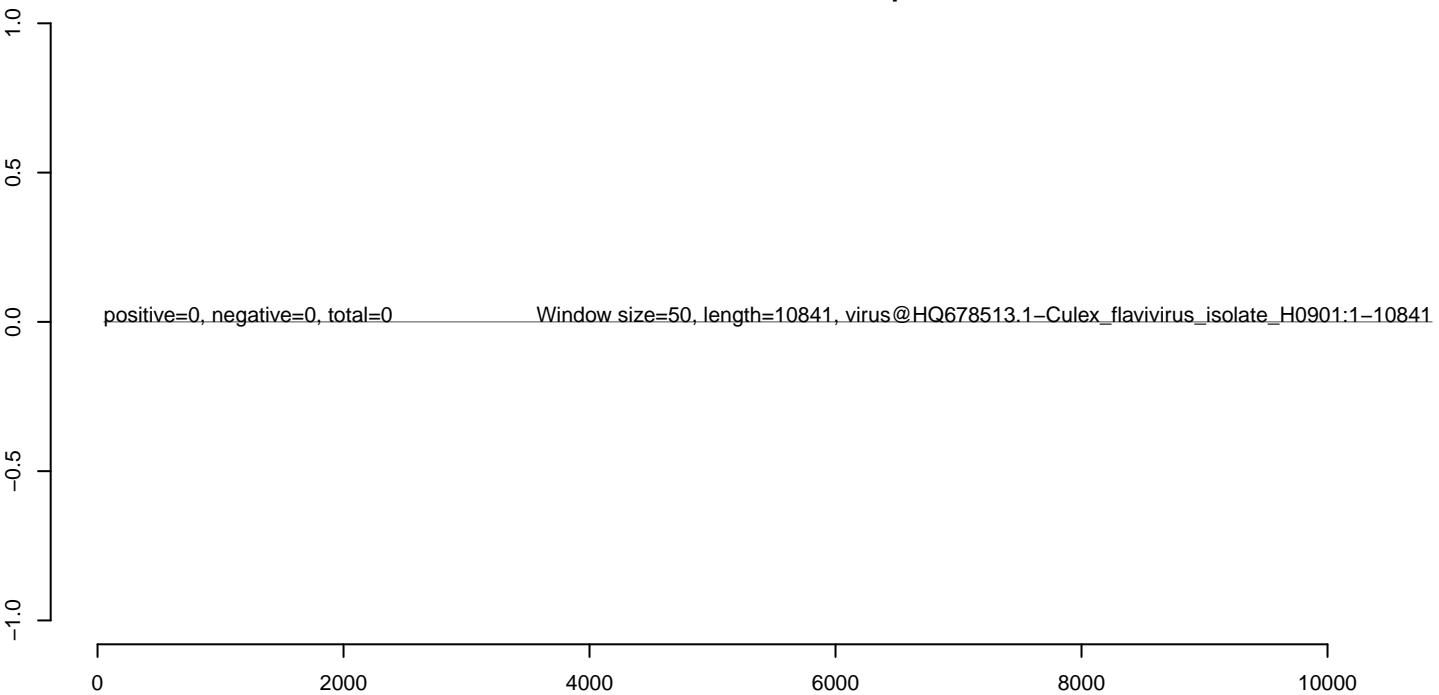
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcarc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcarc\_TN.24\_35.rep

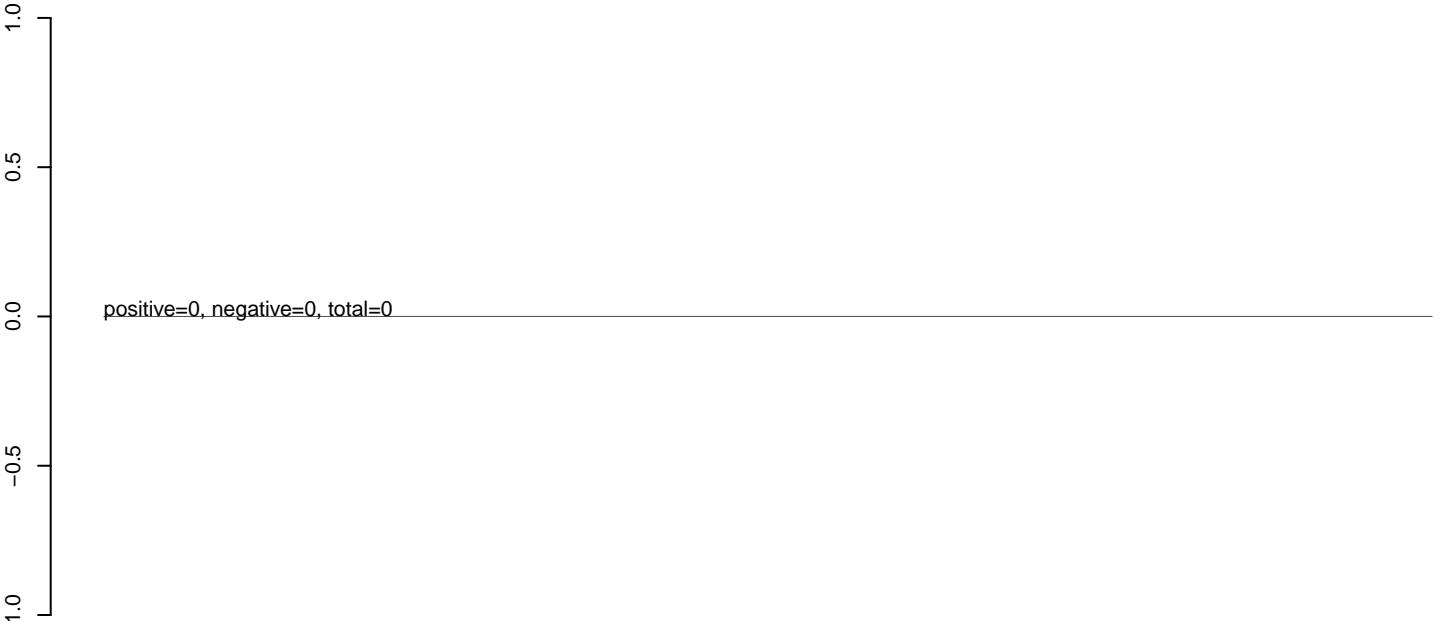
positive=0, negative=0, total=0

### AnGam\_Fcarc\_TN.rep

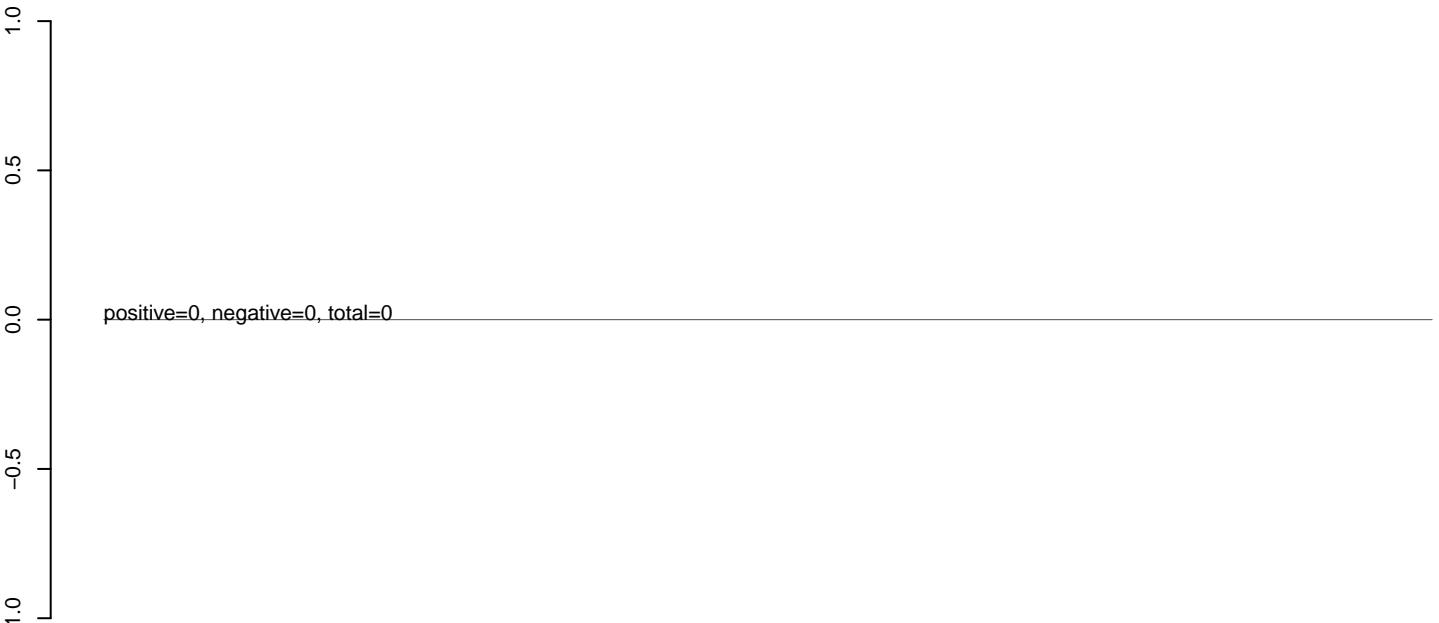
positive=0, negative=0, total=0, window size=50, length=10774, virus@HM055369.1-Kyasanur\_forest\_disease\_virus\_strain\_KFD\_P\_9605:1-10774

0 2000 4000 6000 8000 10000

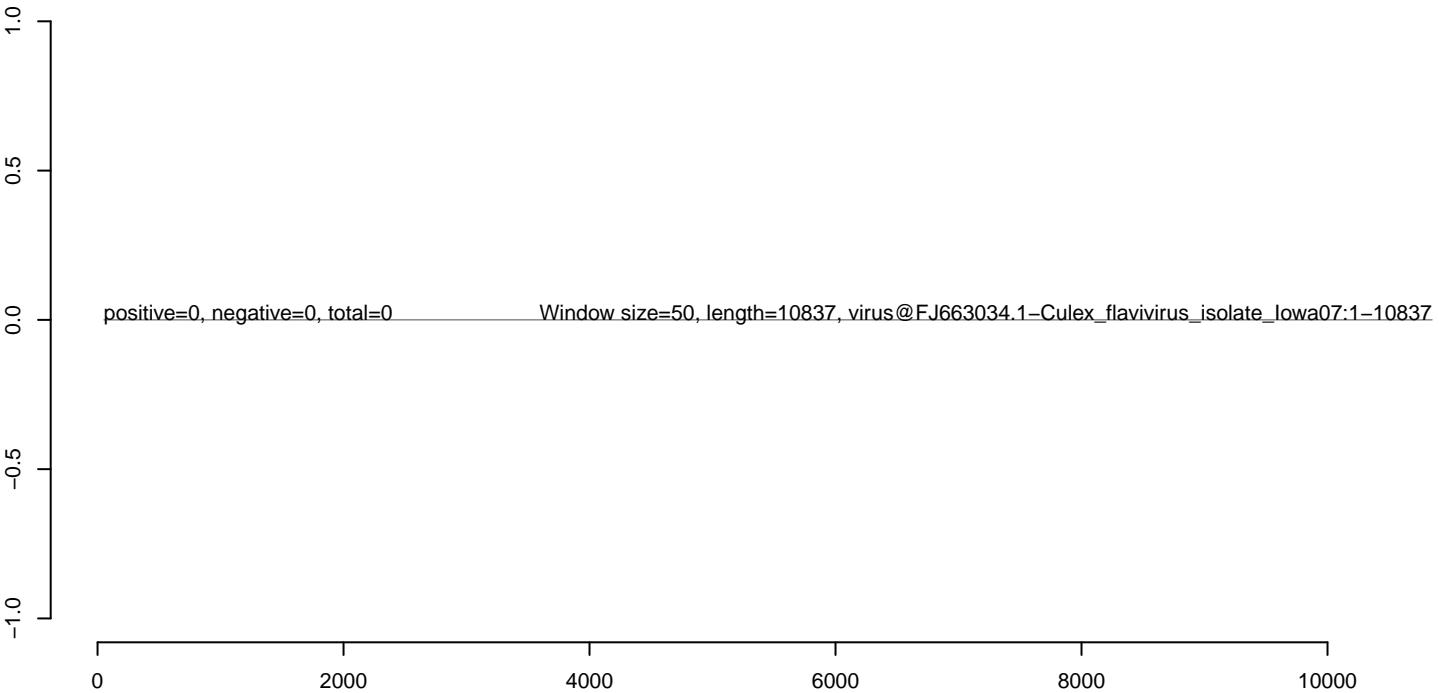
### AnGam\_Fcanc\_TN.18\_23.rep



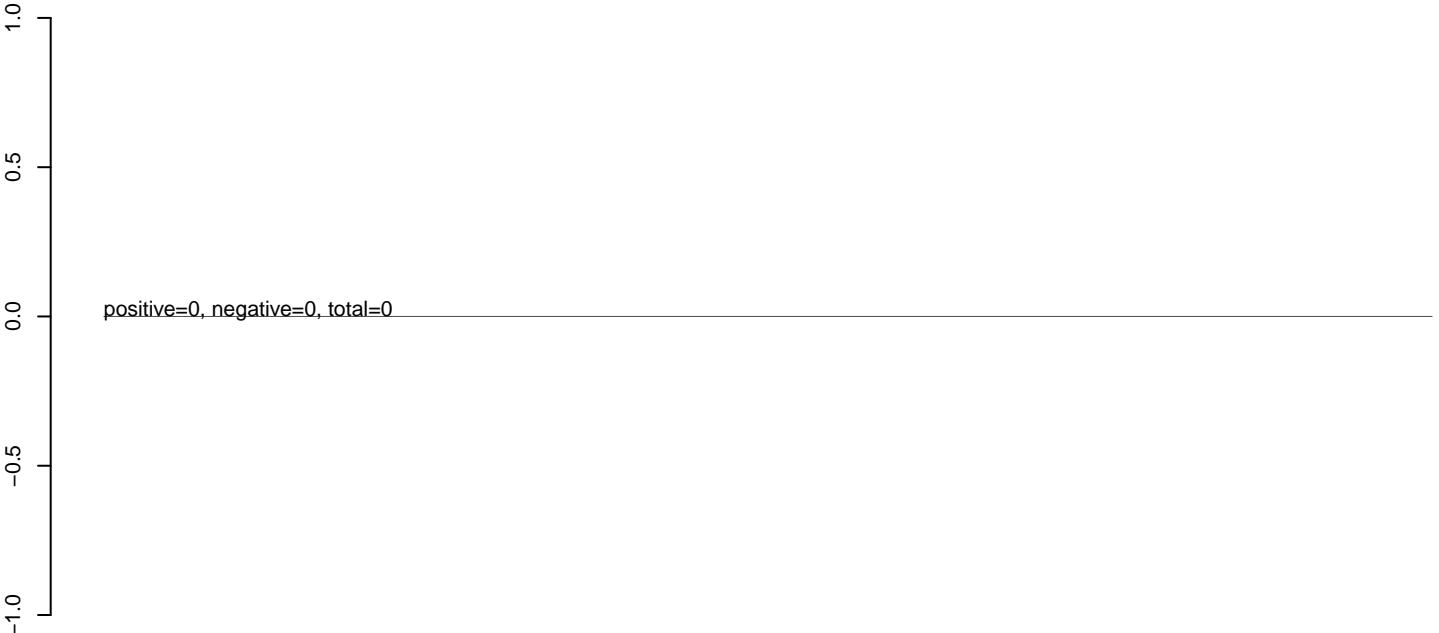
### AnGam\_Fcanc\_TN.24\_35.rep



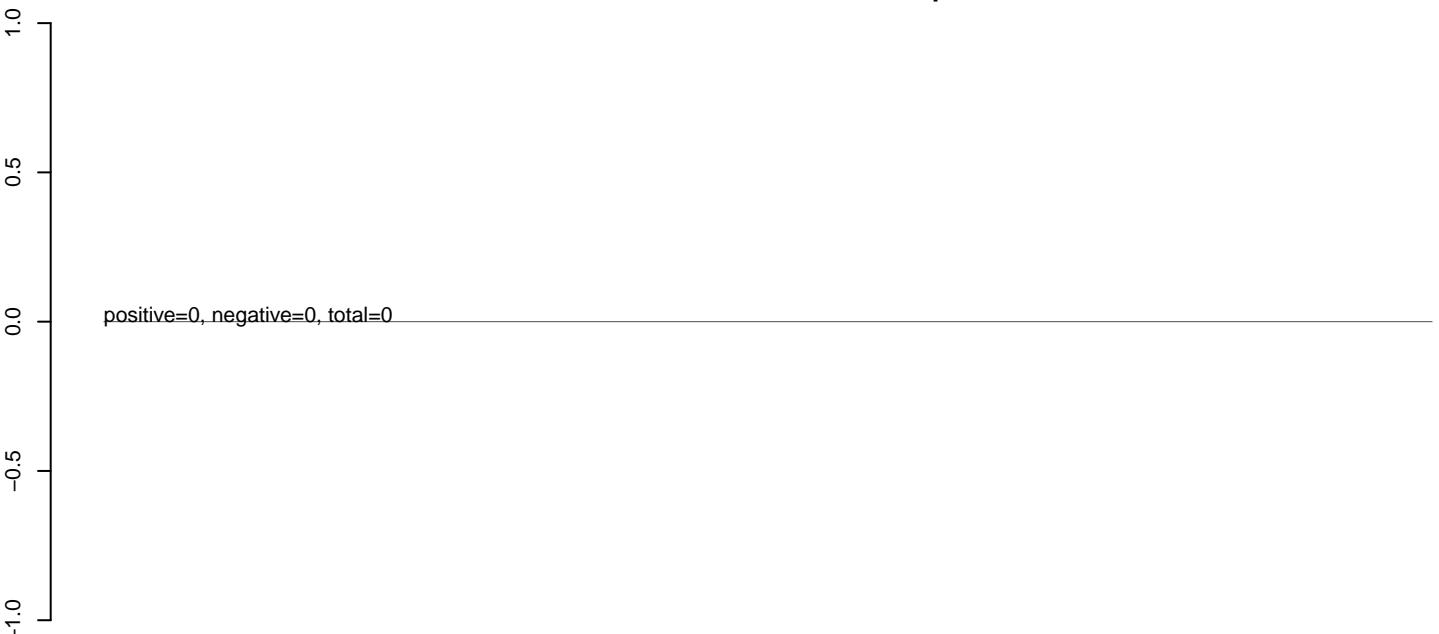
### AnGam\_Fcanc\_TN.rep



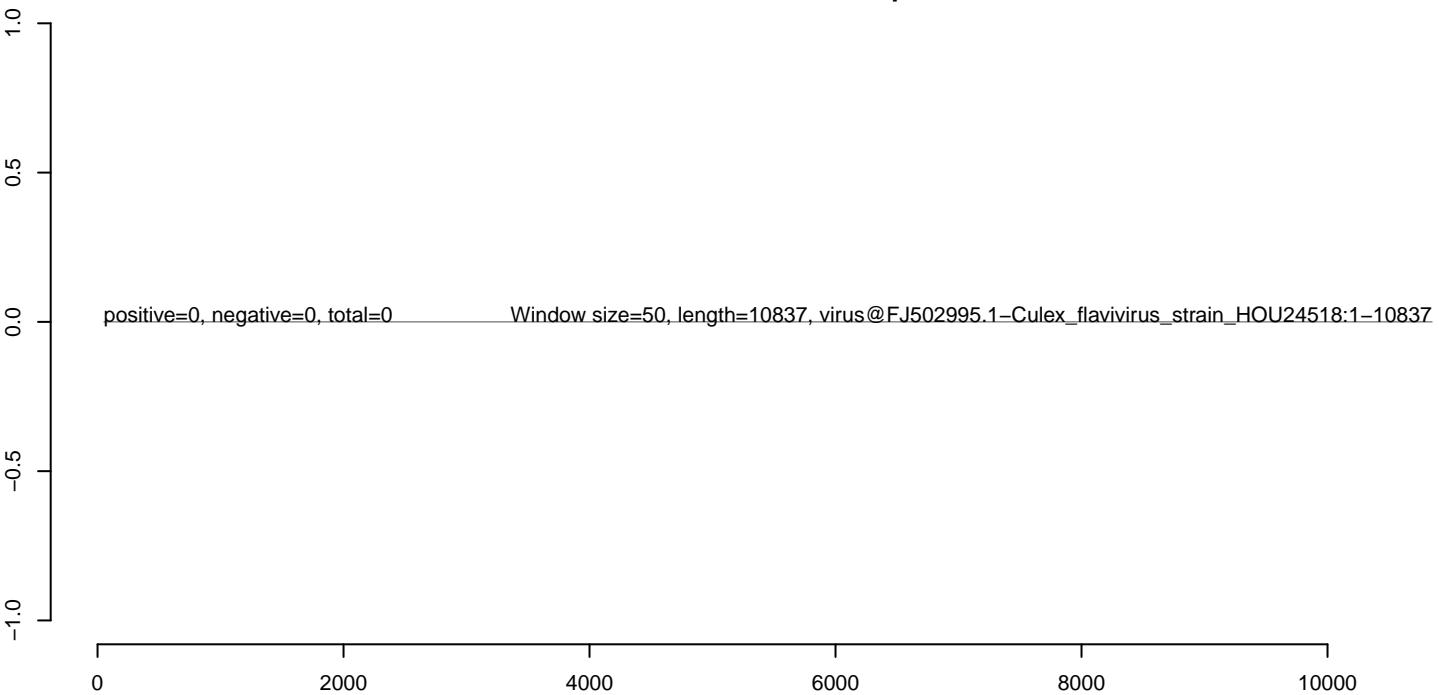
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0

Window size=50, length=10837, virus@EU879060.1-Culex\_flavivirus\_strain\_CxFV-Mex07:1-10837

0 2000 4000 6000 8000 10000

### AnGam\_Fcarc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcarc\_TN.24\_35.rep

positive=0, negative=0, total=0

### AnGam\_Fcarc\_TN.rep

positive=0, negative=0, total=0, length=10302, virus@EU082200.2-Koutango\_virus\_strain\_Dak\_Ar\_D\_5443\_polyprotein\_gene:1-10302

0 2000 4000 6000 8000 10000

### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

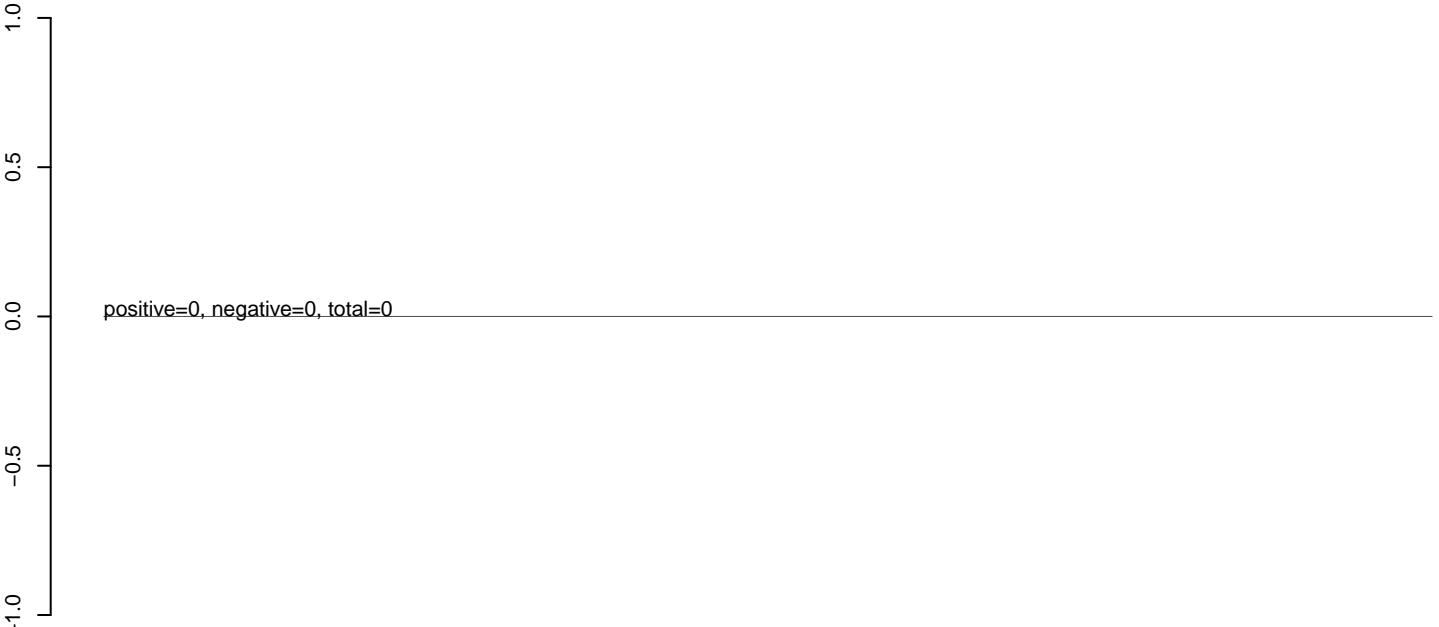
### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

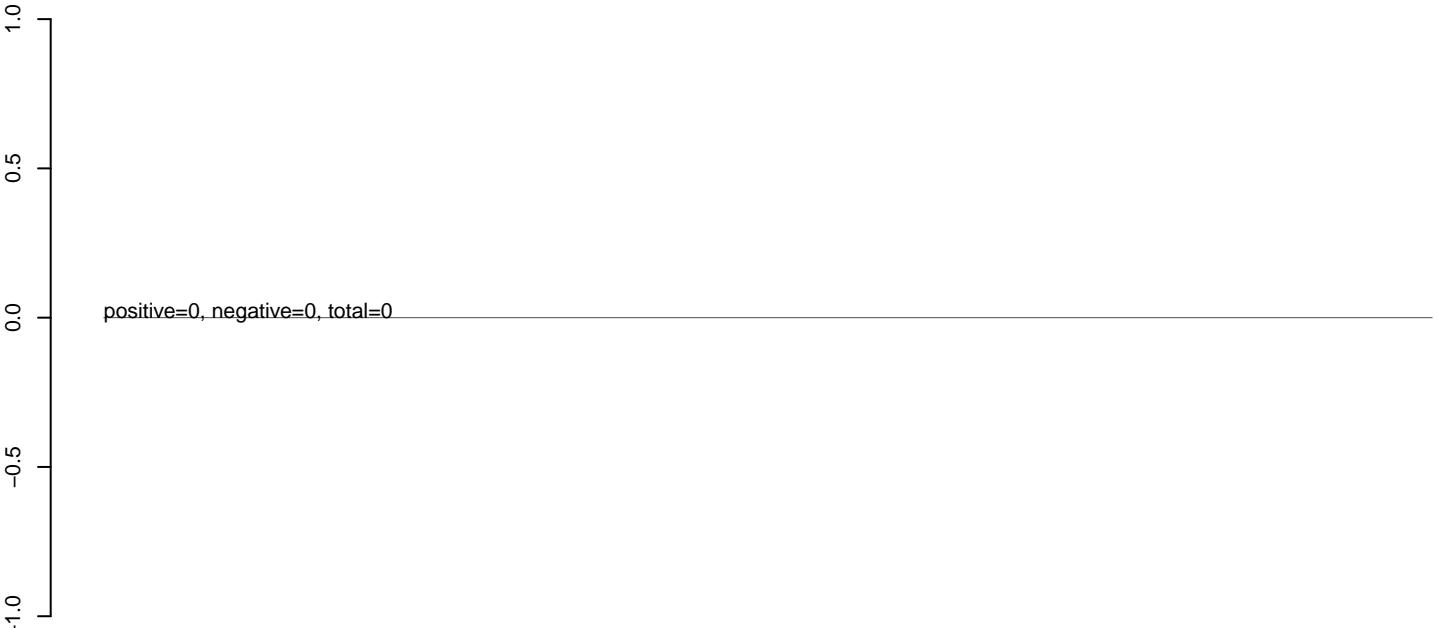
### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0 window size=50, length=10182, virus@DQ859056.1-Banzi\_virus\_strain\_SAH\_336\_polyprotein\_gene:1-10182

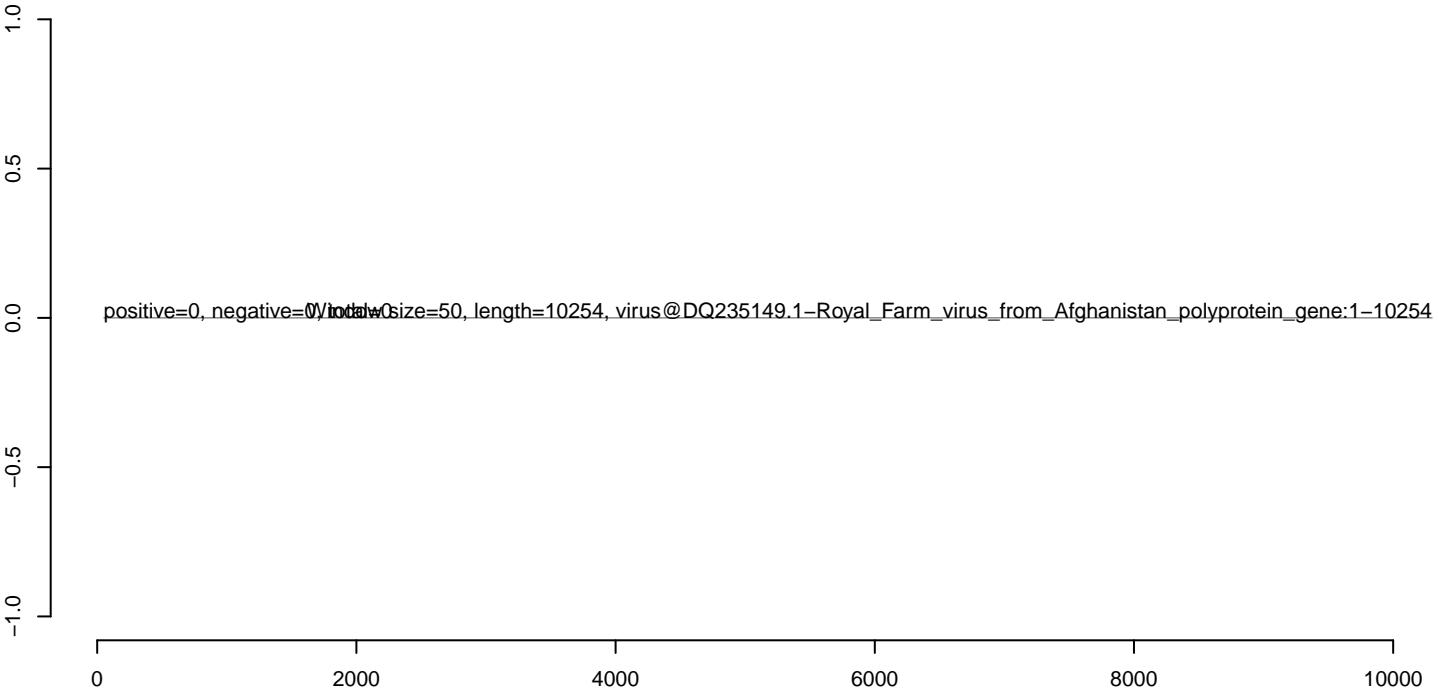
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

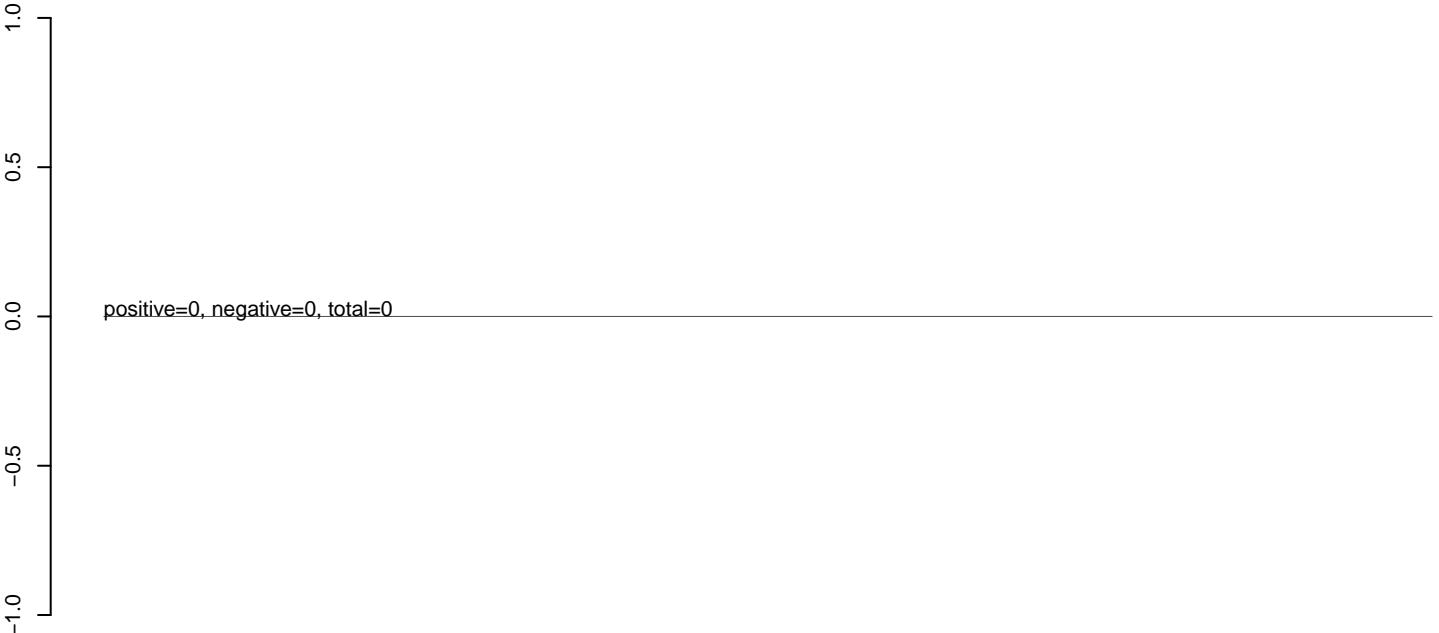
positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.rep

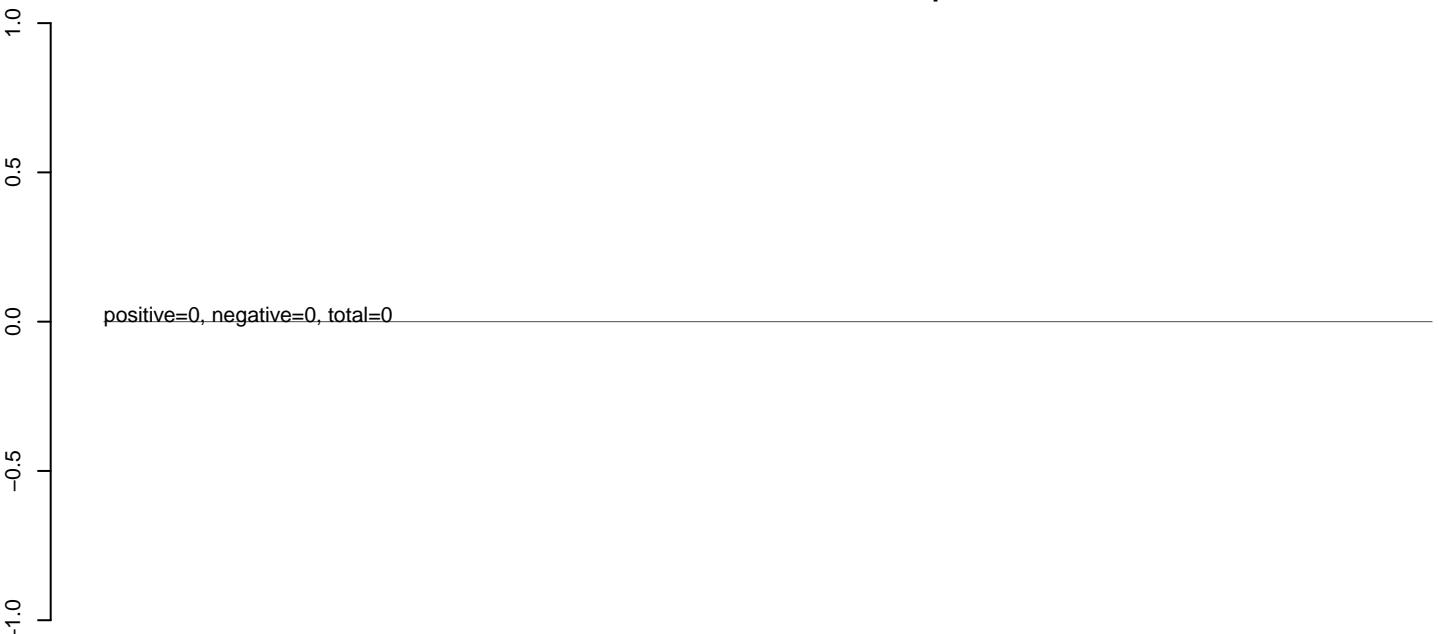
positive=0, negative=0, total=0

Window size=50, length=10960, virus@AY898809.1-Alfuy\_virus\_strain\_MRM3929:1-10960

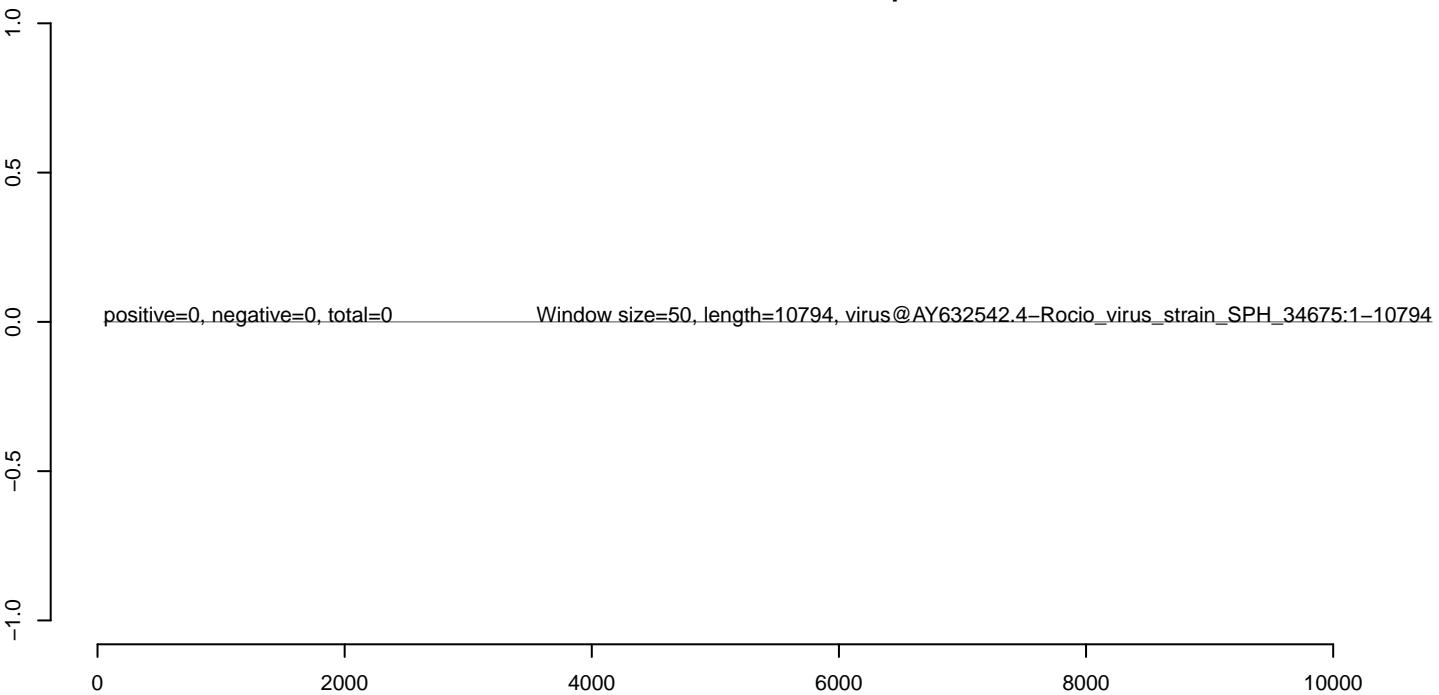
### AnGam\_Fcanc\_TN.18\_23.rep



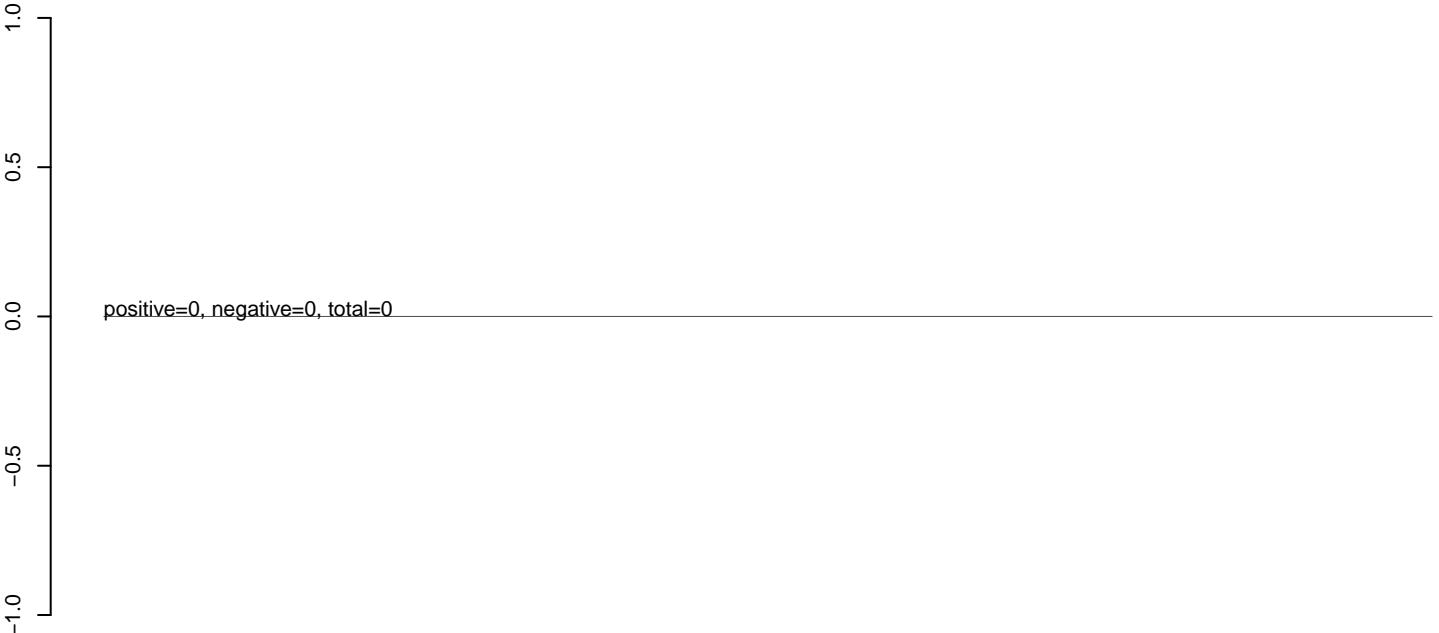
### AnGam\_Fcanc\_TN.24\_35.rep



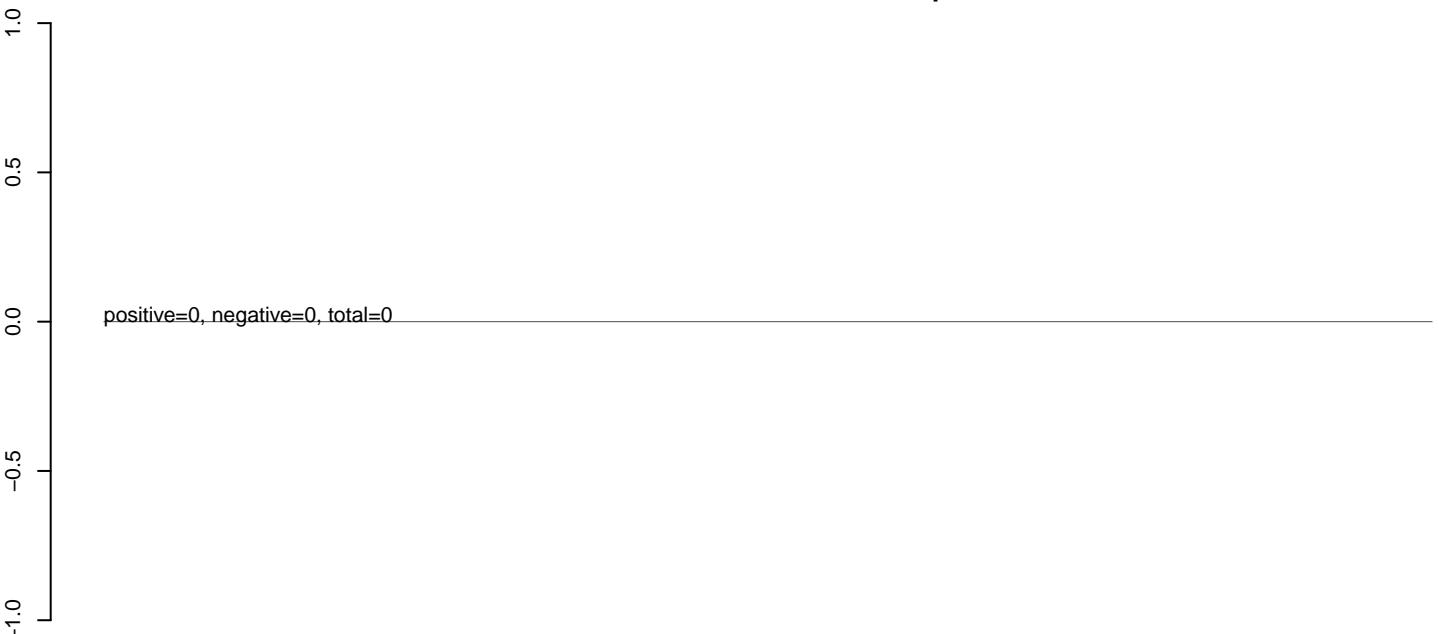
### AnGam\_Fcanc\_TN.rep



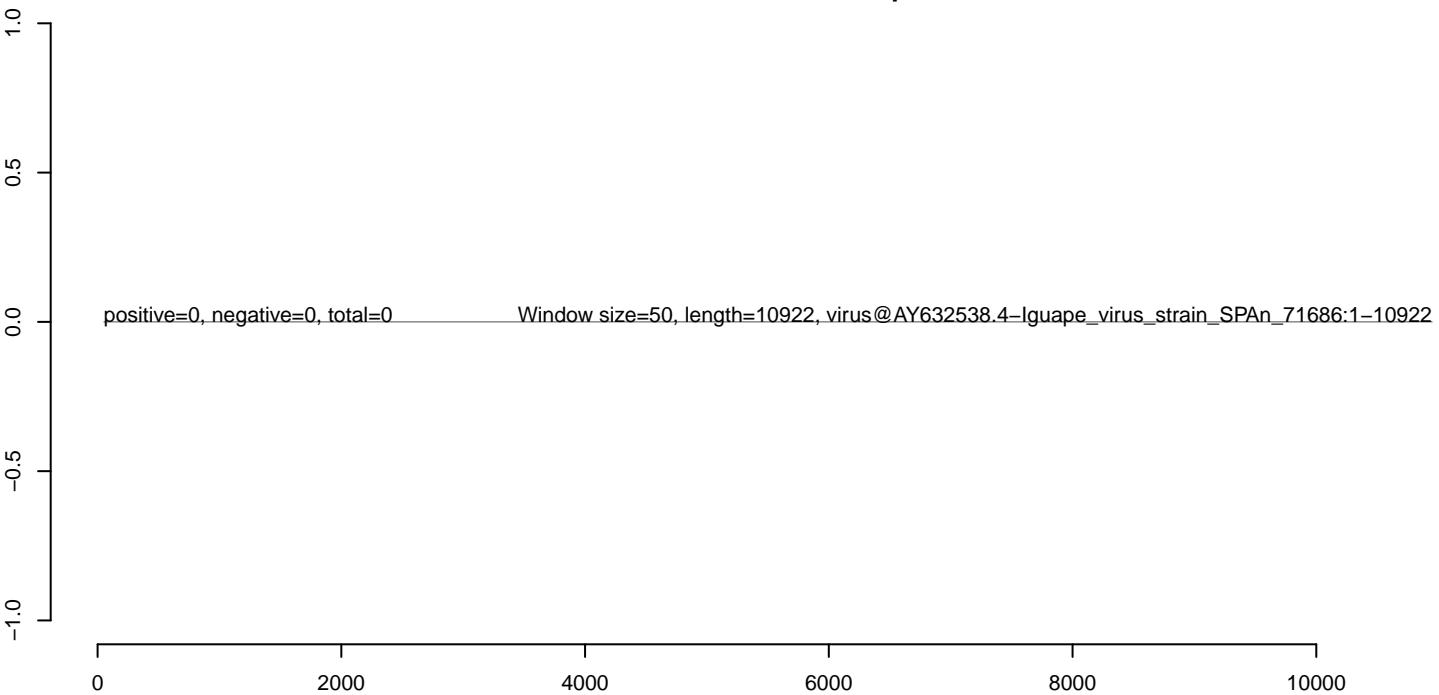
### AnGam\_Fcanc\_TN.18\_23.rep



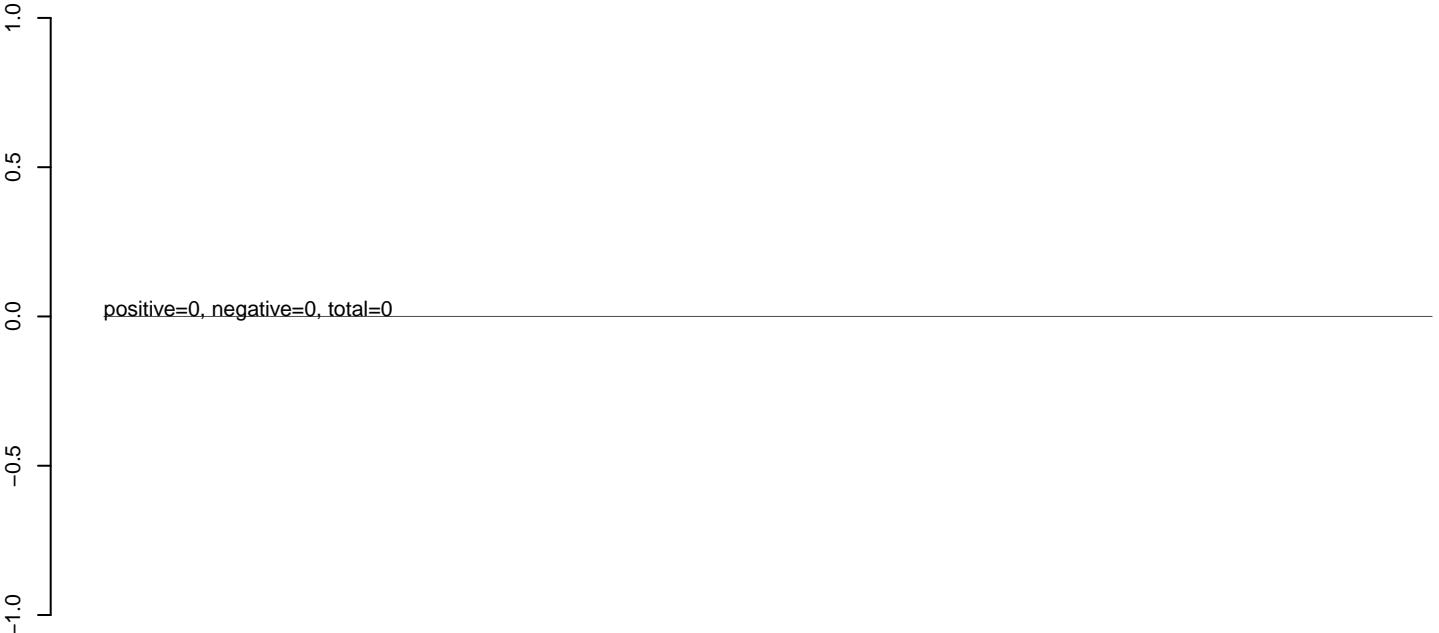
### AnGam\_Fcanc\_TN.24\_35.rep



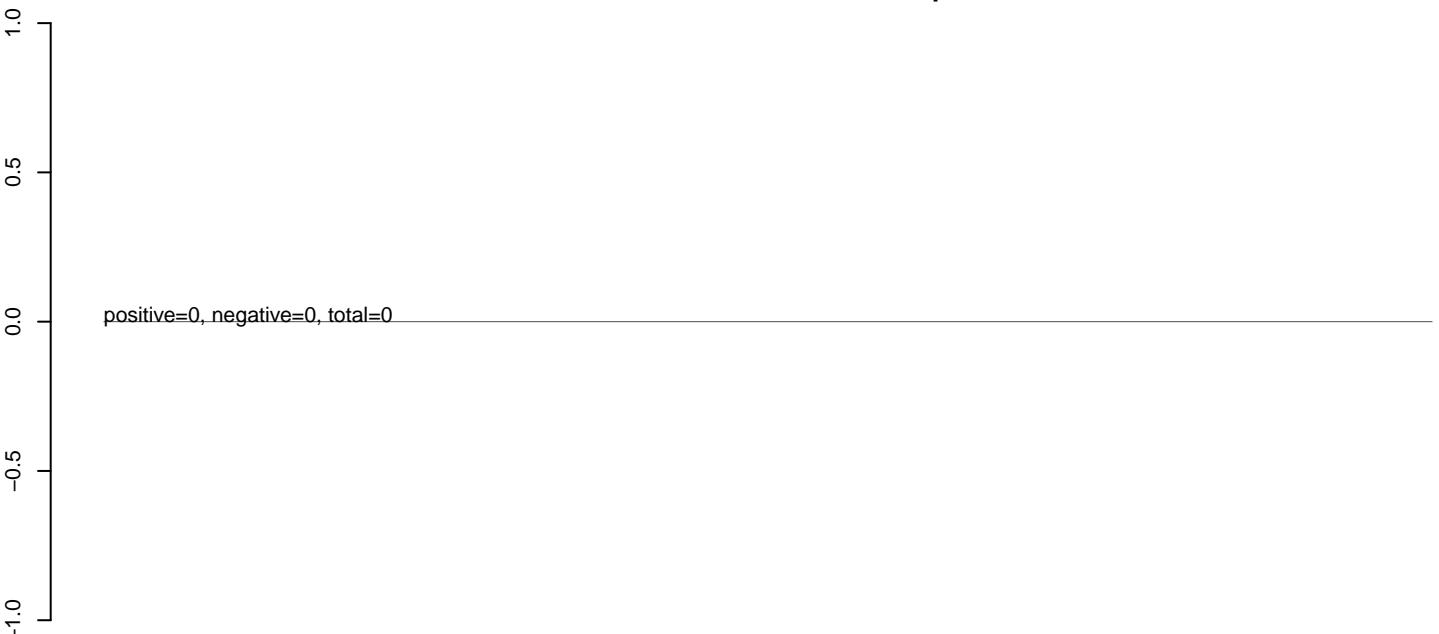
### AnGam\_Fcanc\_TN.rep



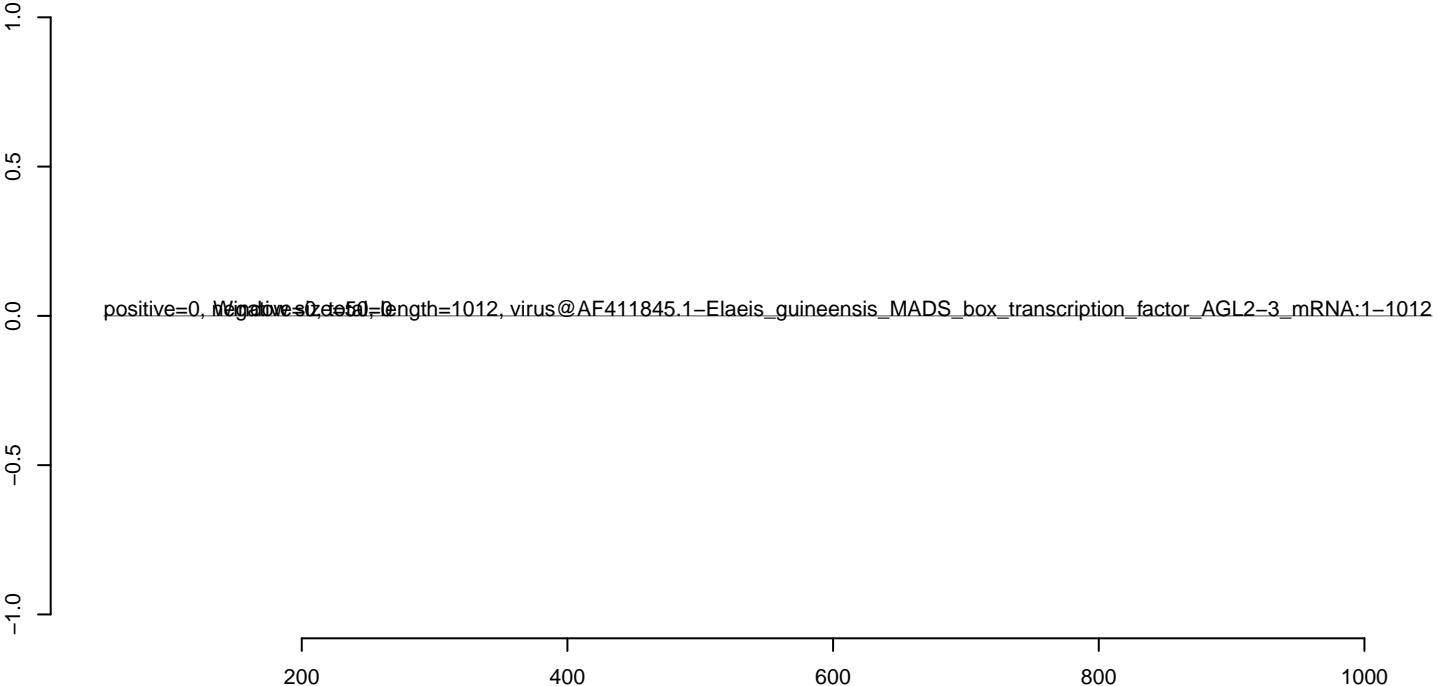
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

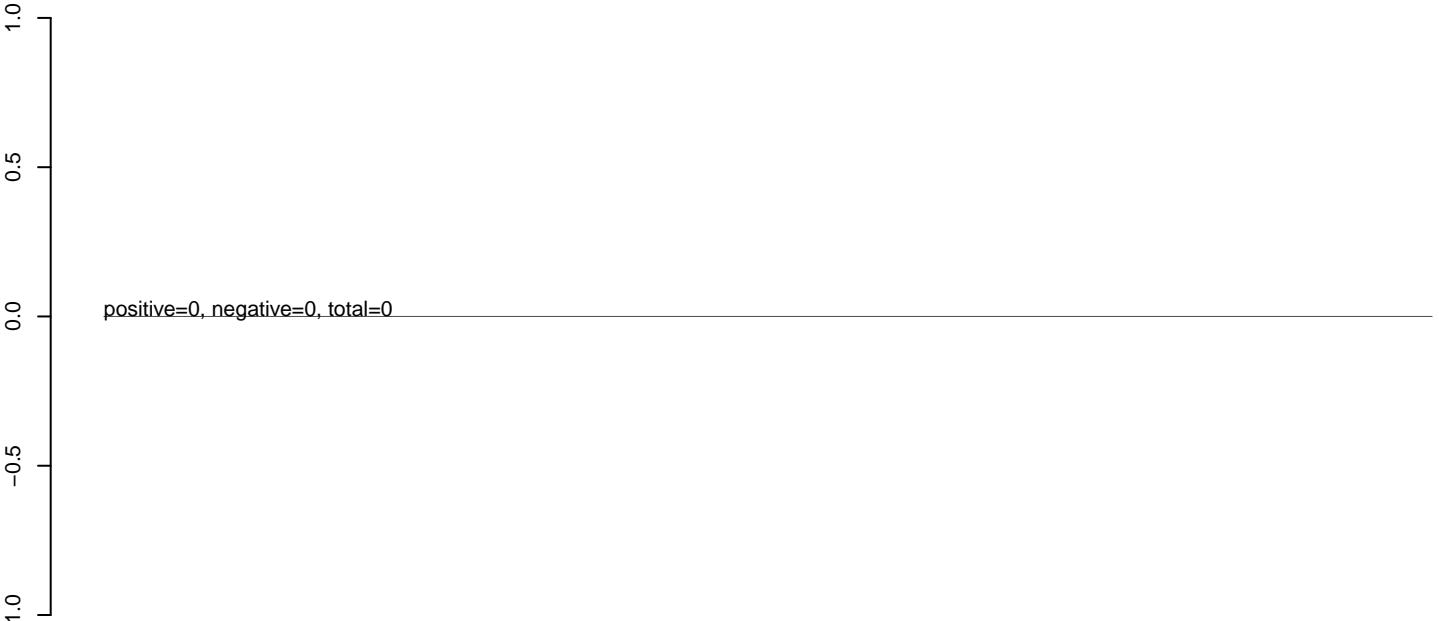
### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

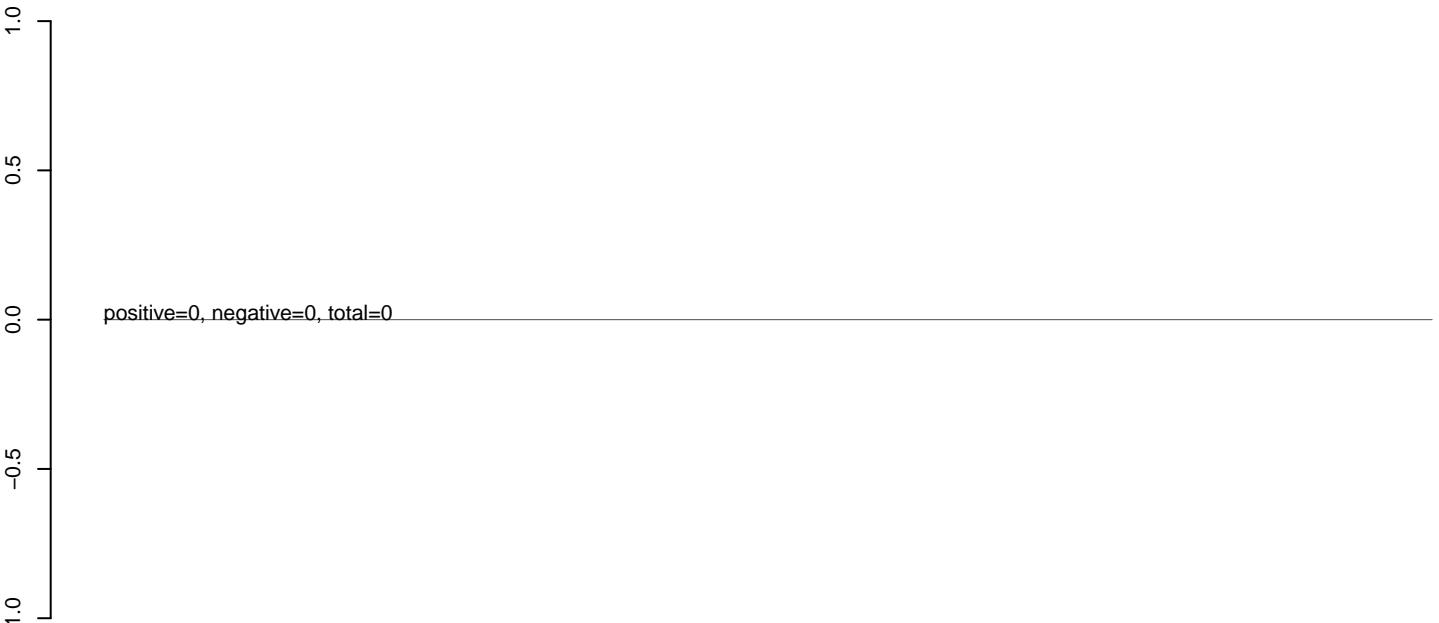
### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=150, length=972, virus@AF411844.1-Elaeis\_guineensis\_MADS\_box\_transcription\_factor\_AGL2-2\_mRNA:1-972

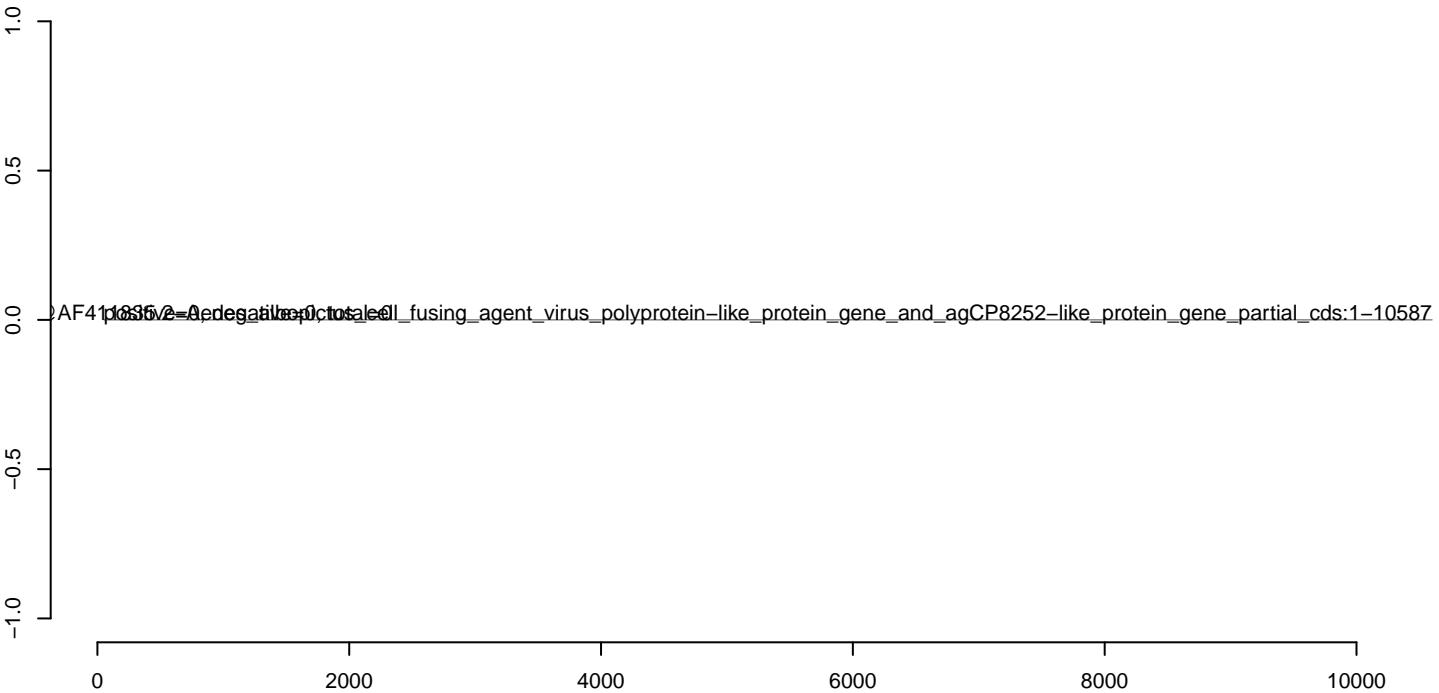
### AnGam\_Fcanc\_TN.18\_23.rep



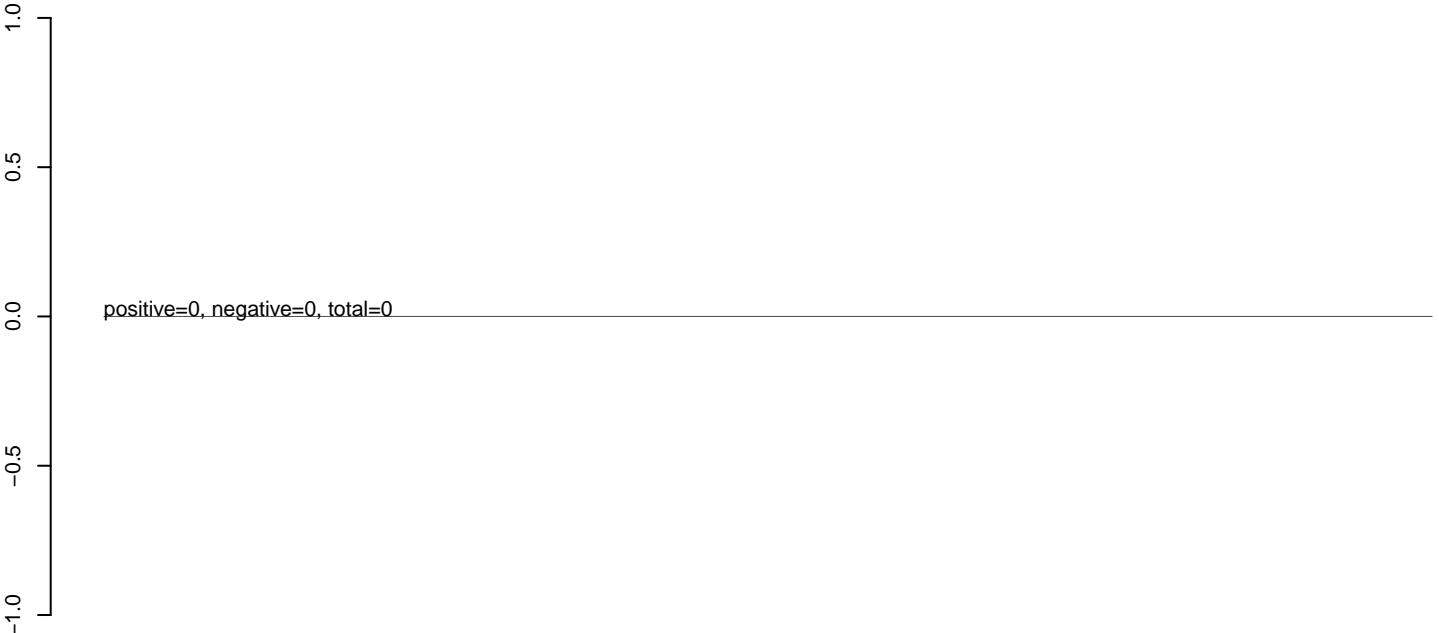
### AnGam\_Fcanc\_TN.24\_35.rep



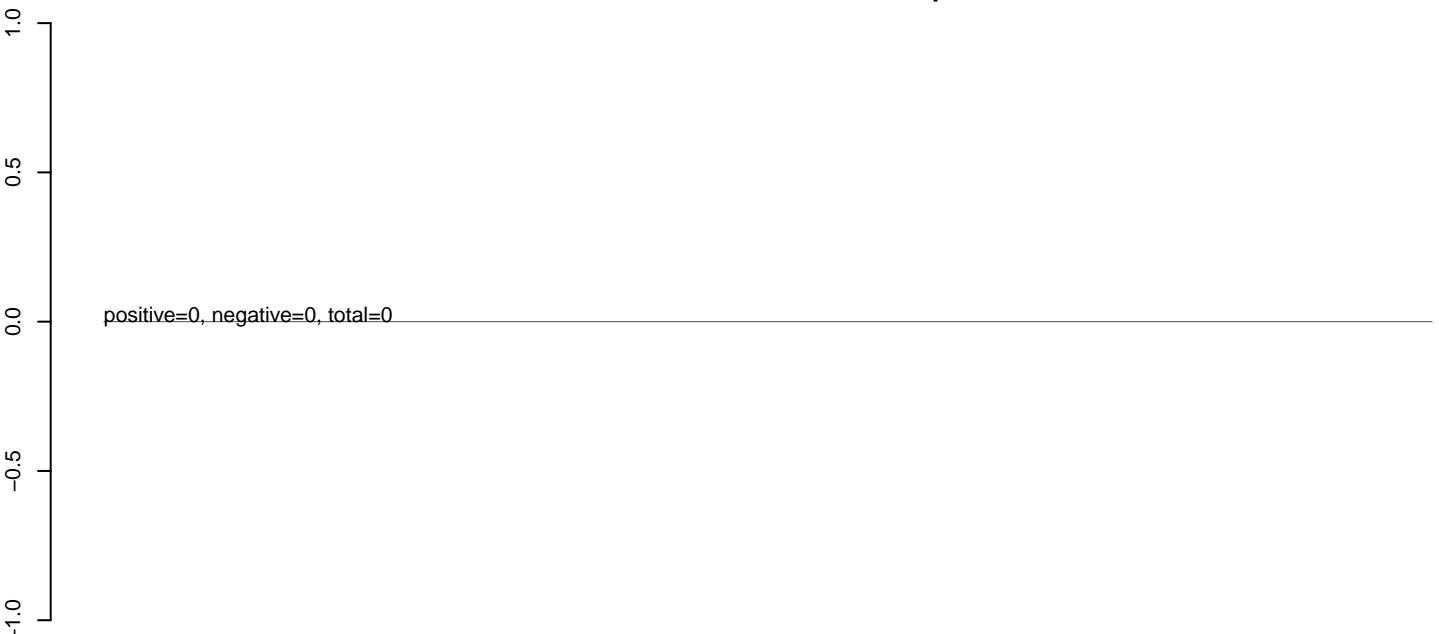
### AnGam\_Fcanc\_TN.rep



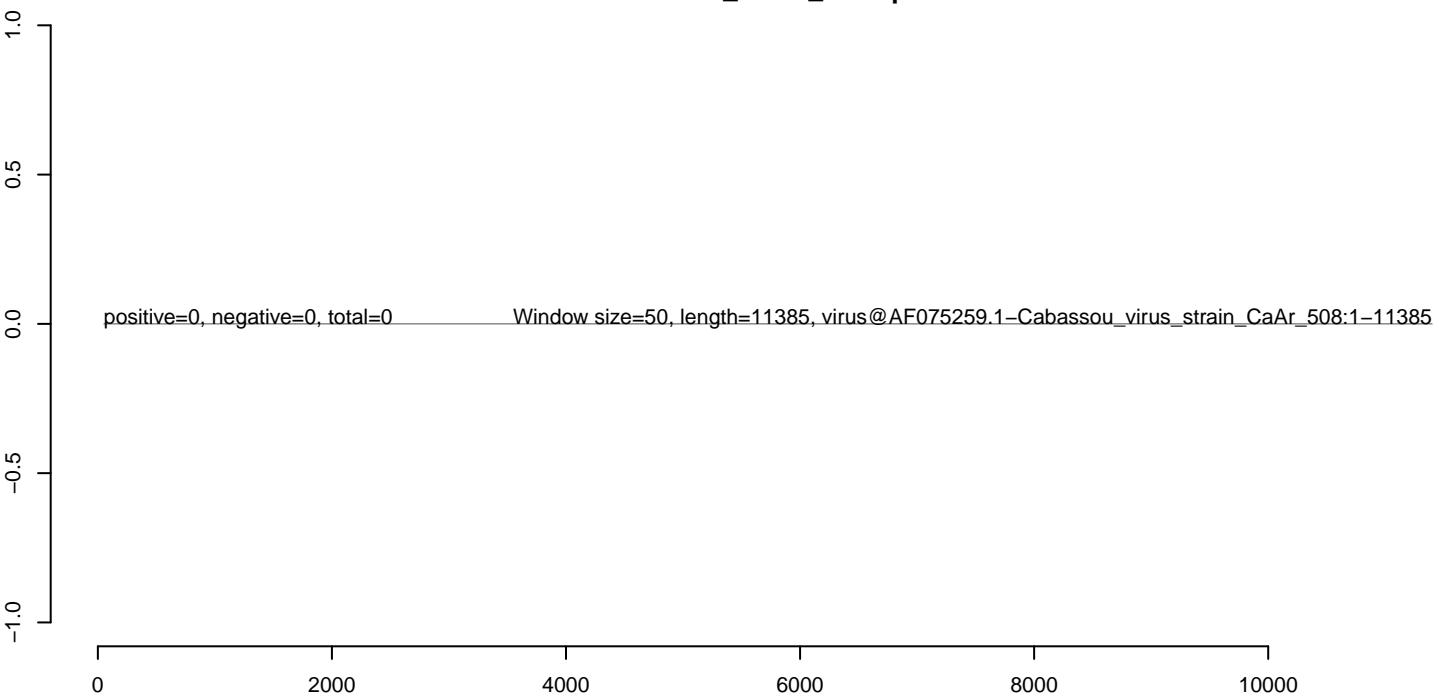
### AnGam\_Fcanc\_TN.18\_23.rep



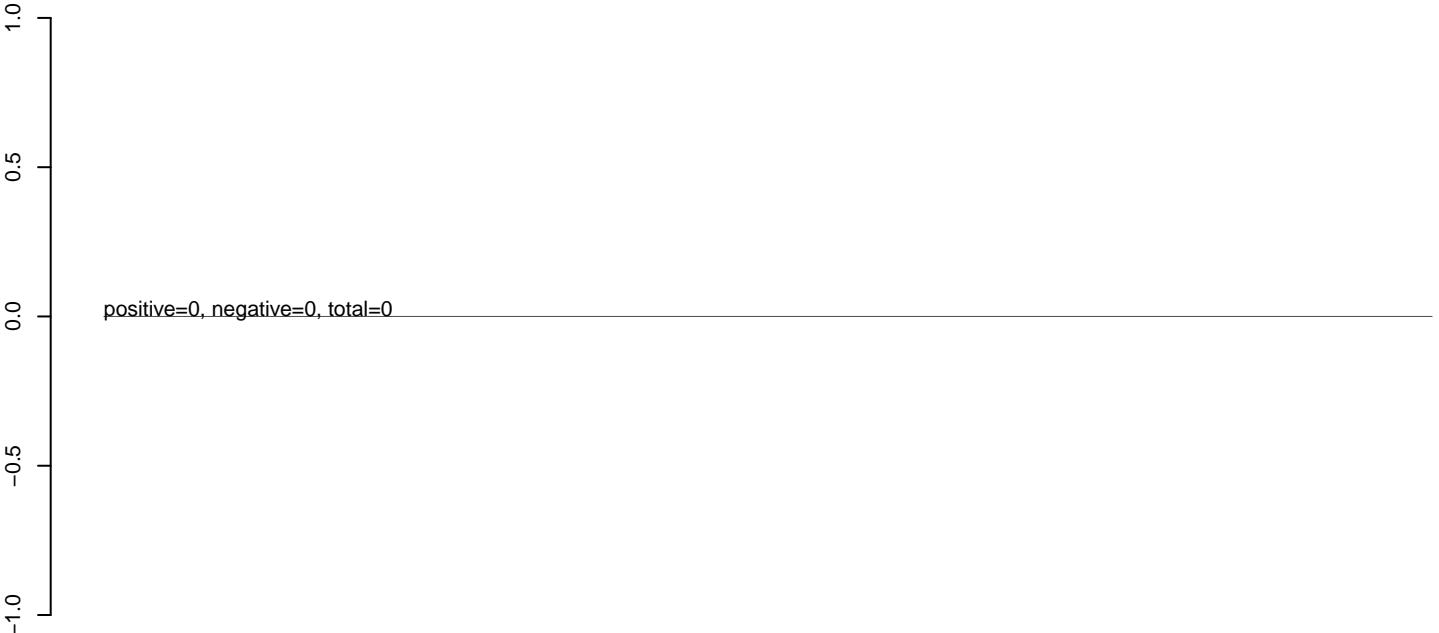
### AnGam\_Fcanc\_TN.24\_35.rep



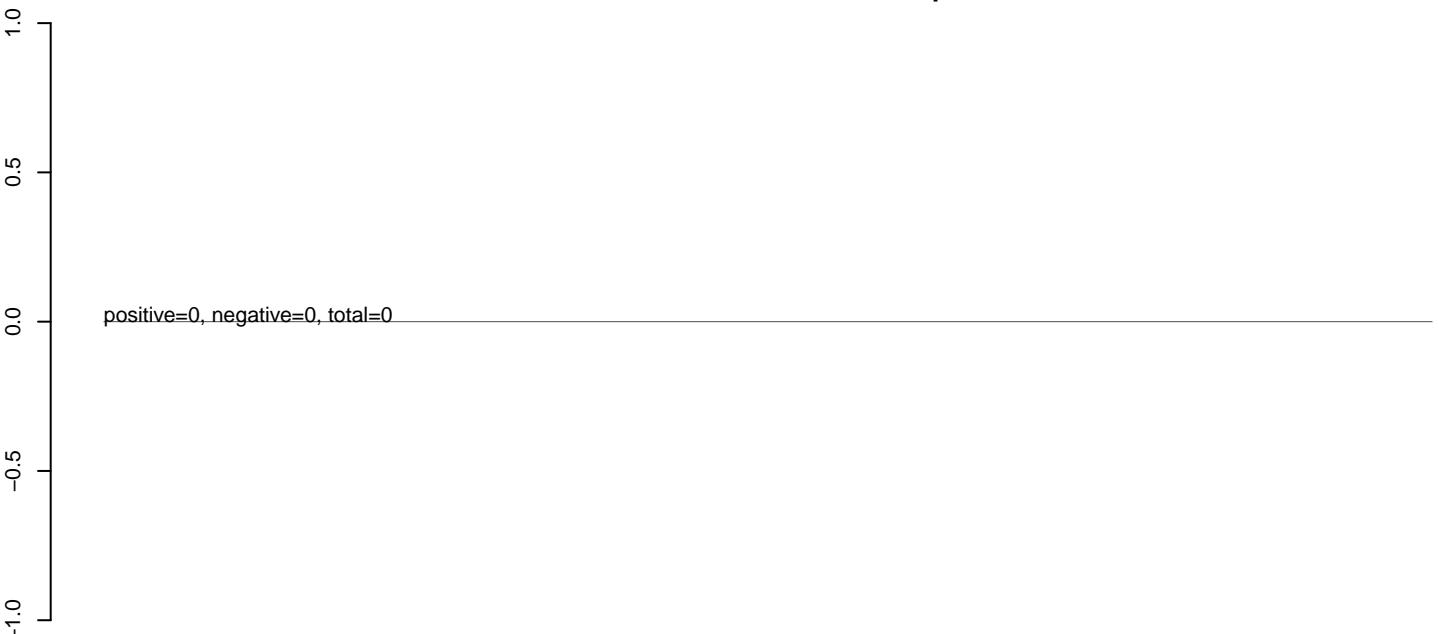
### AnGam\_Fcanc\_TN.rep



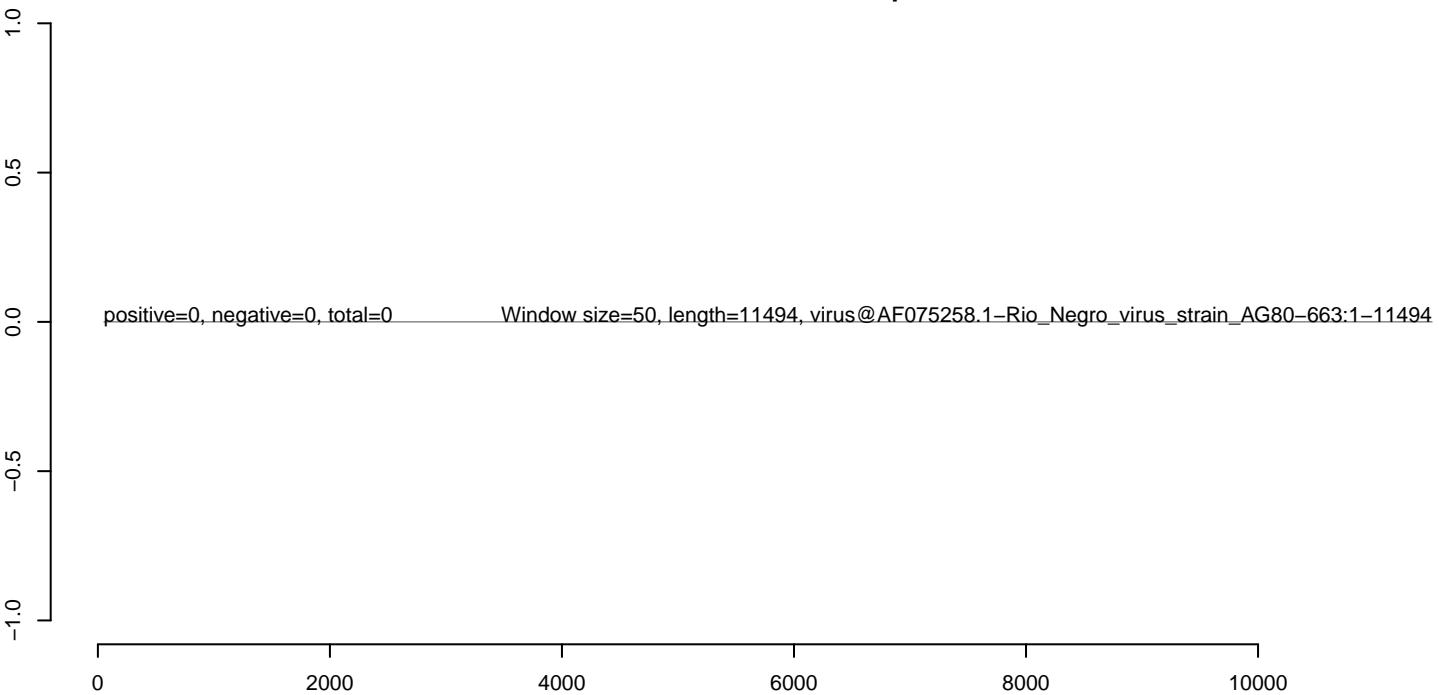
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

positive=0, negative=0, total=0

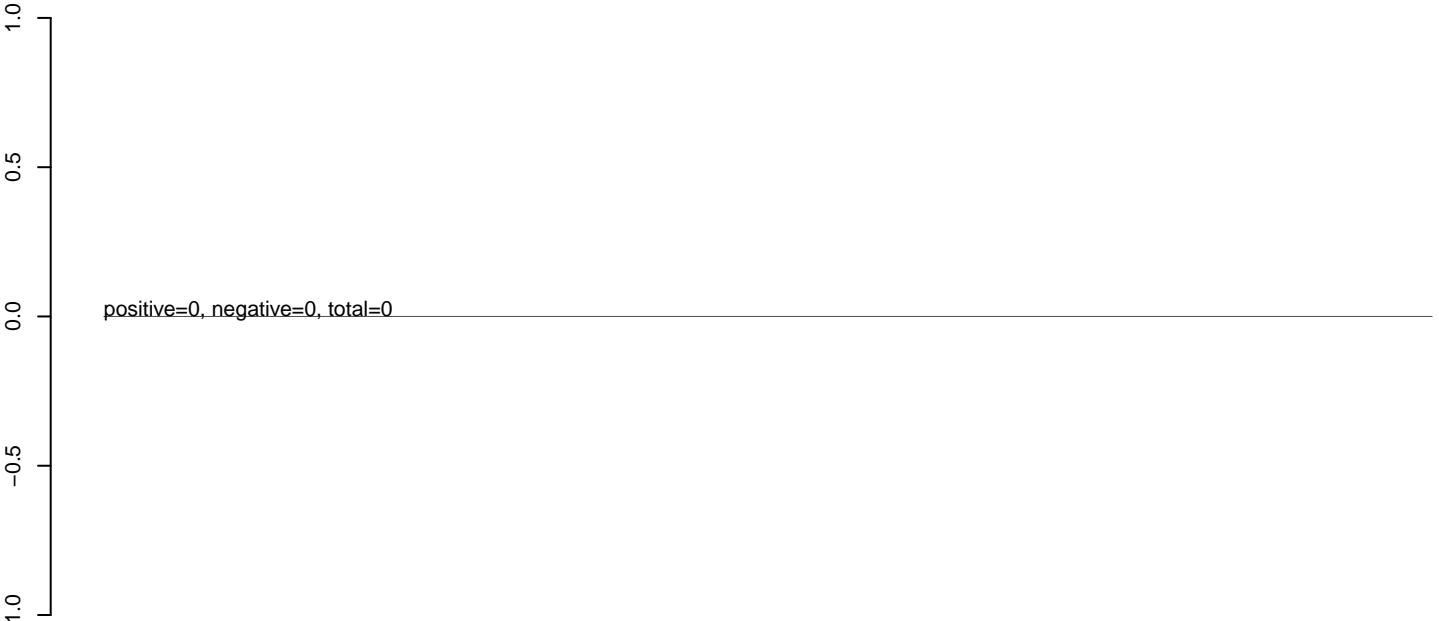
### AnGam\_Fcanc\_TN.rep

positive=0, negative=0, total=0

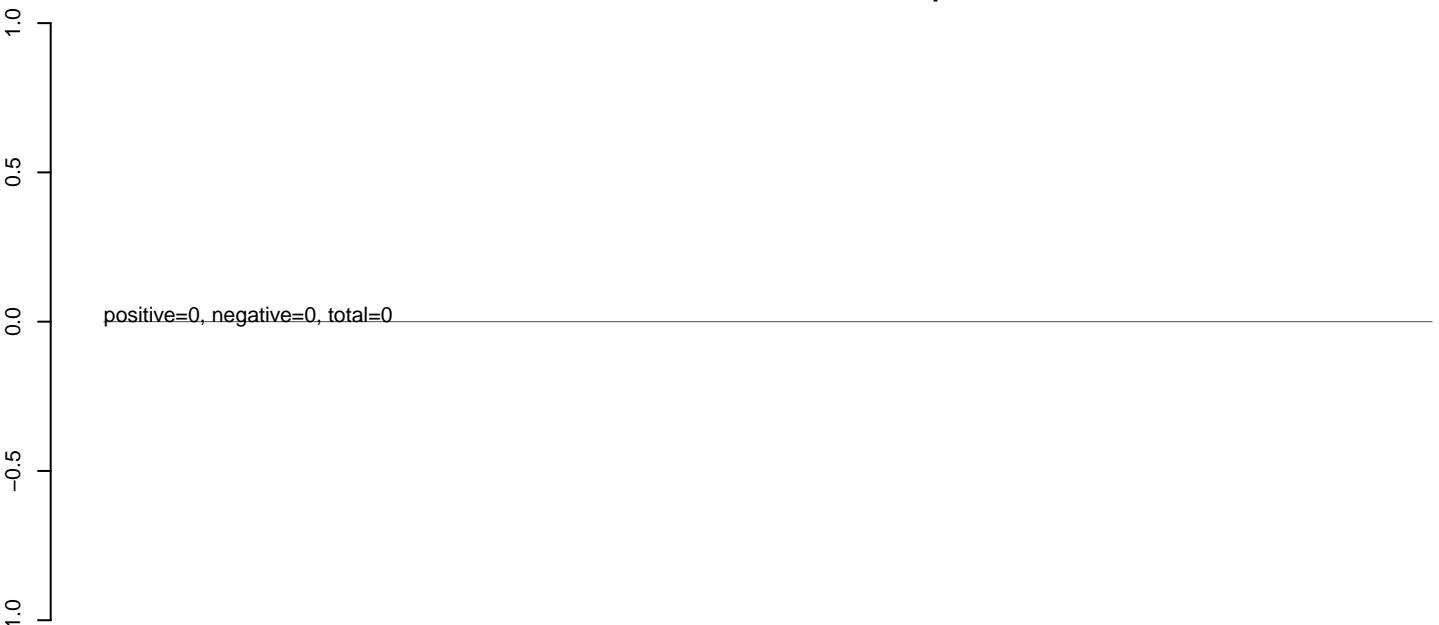
Window size=50, length=11465, virus@AF075257.1-Mosso\_das\_Pedras\_virus:1-11465

0 2000 4000 6000 8000 10000

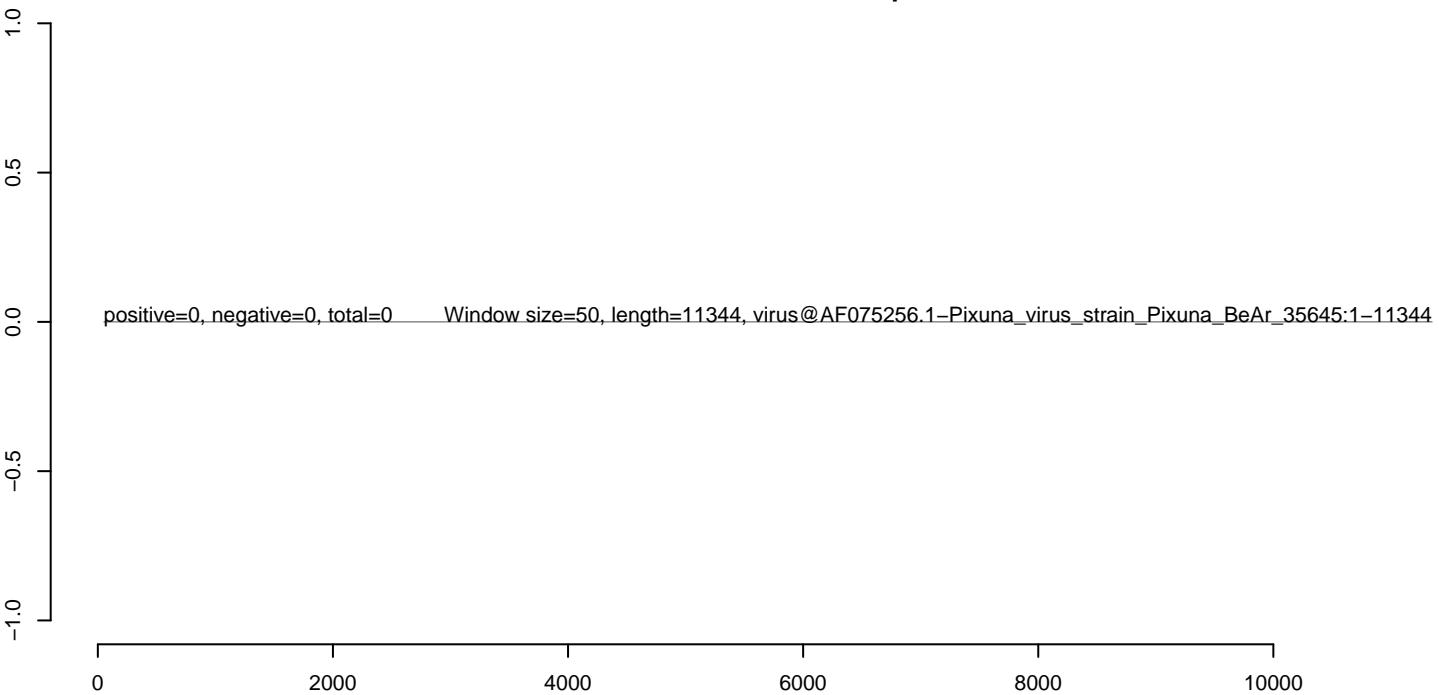
### AnGam\_Fcanc\_TN.18\_23.rep



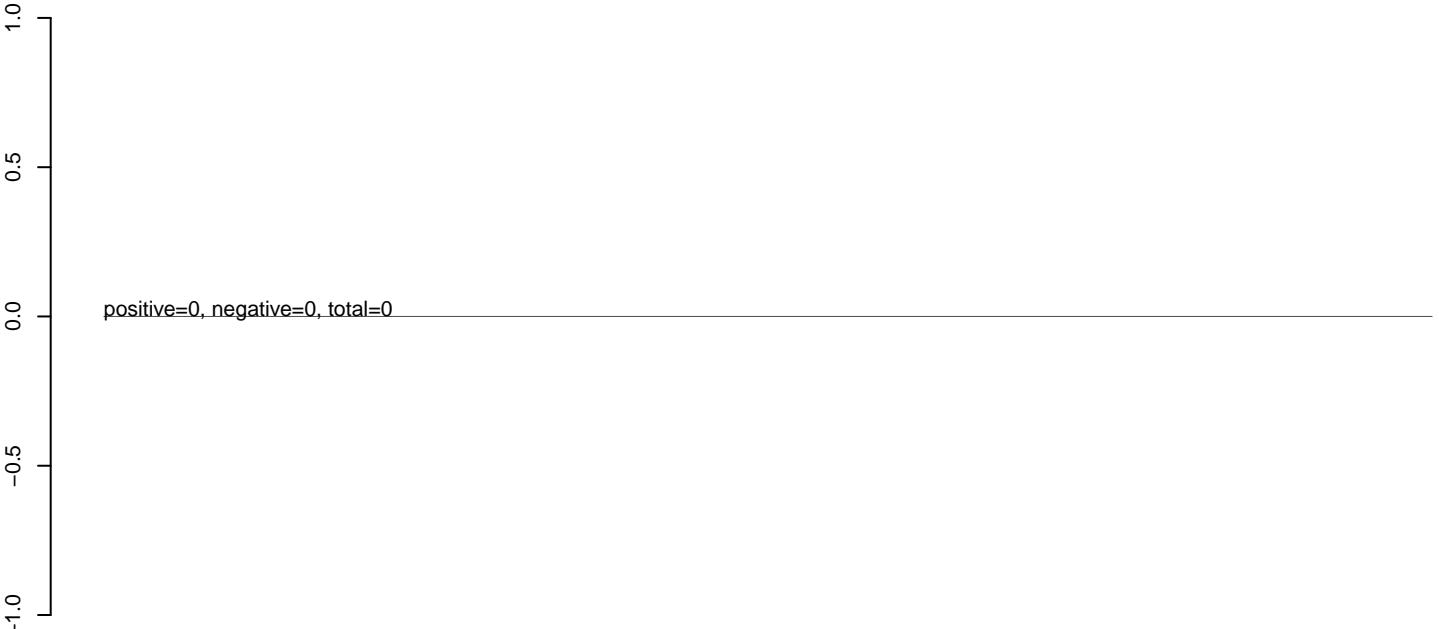
### AnGam\_Fcanc\_TN.24\_35.rep



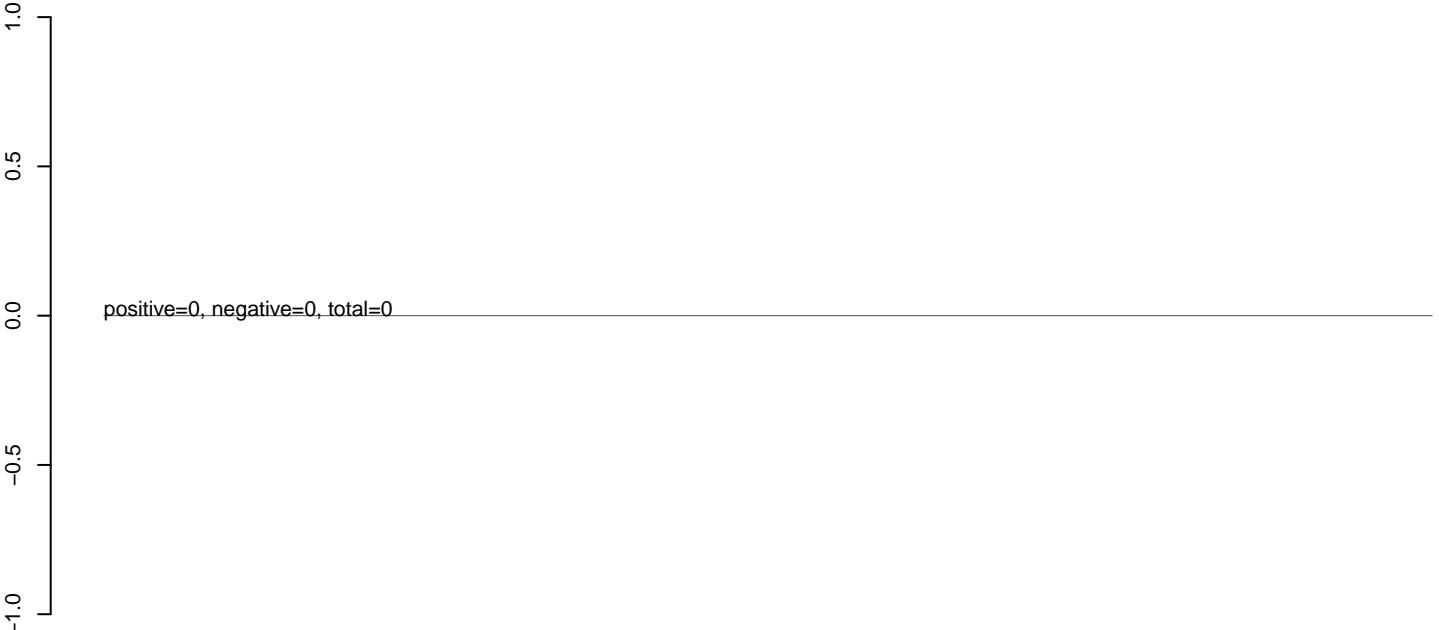
### AnGam\_Fcanc\_TN.rep



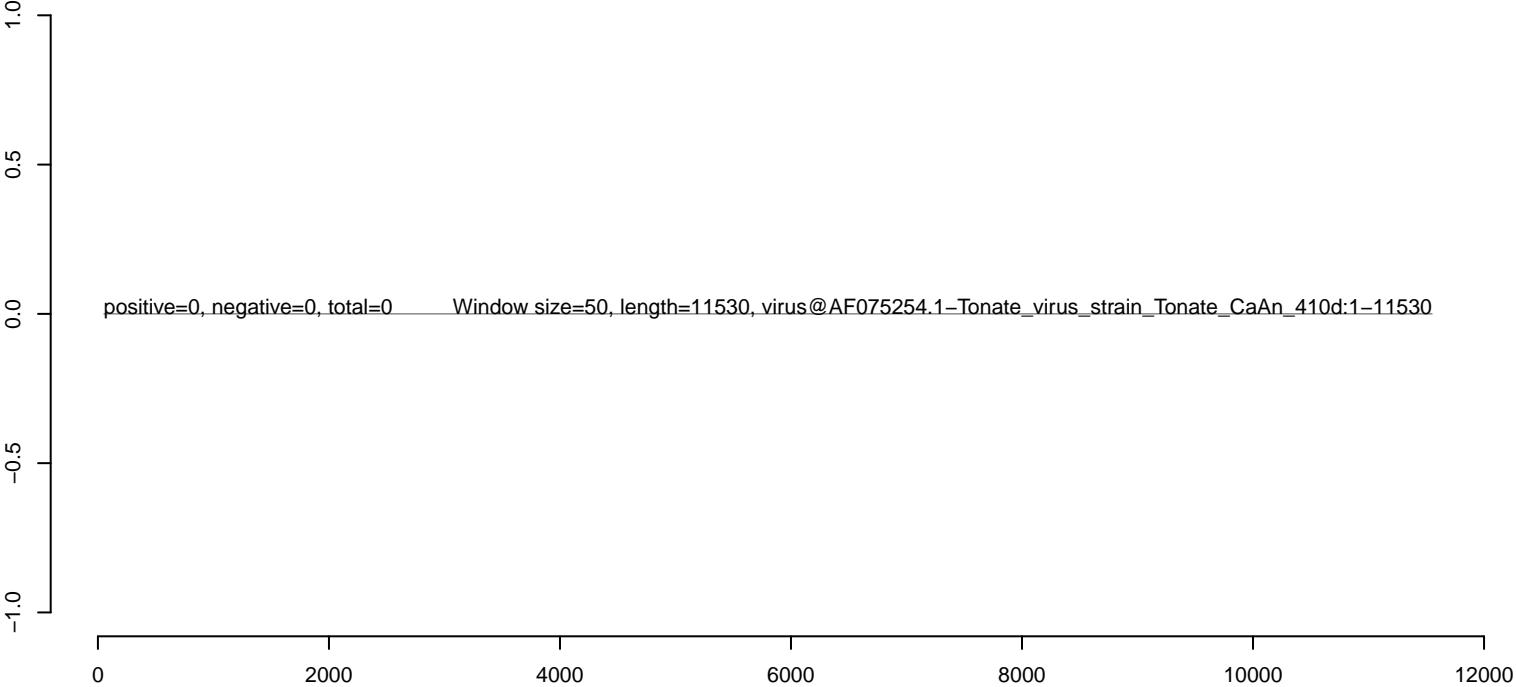
### AnGam\_Fcanc\_TN.18\_23.rep



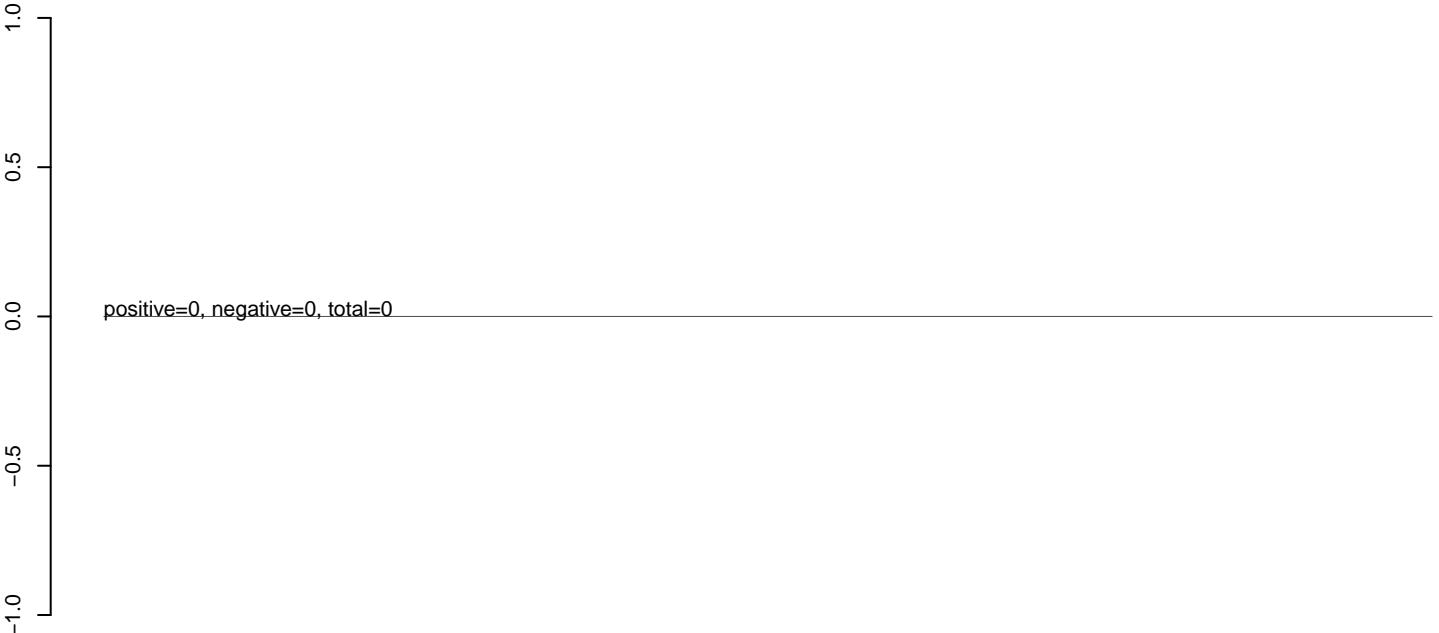
### AnGam\_Fcanc\_TN.24\_35.rep



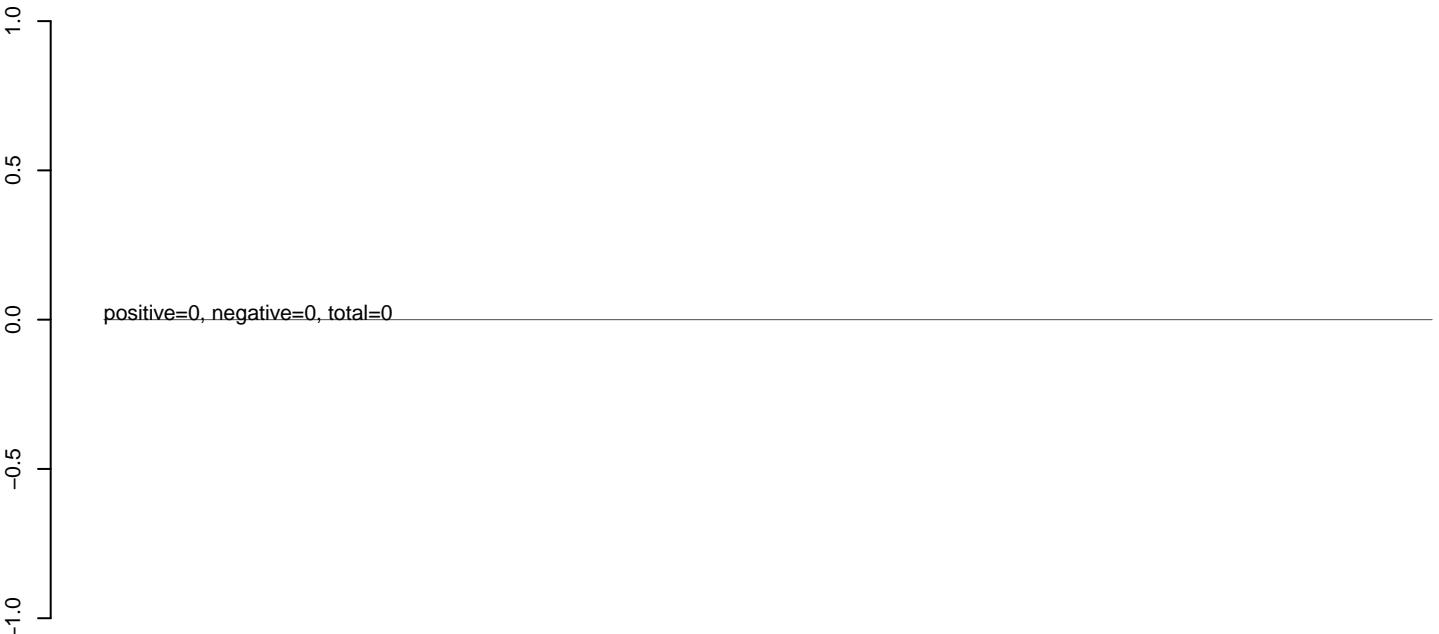
### AnGam\_Fcanc\_TN.rep



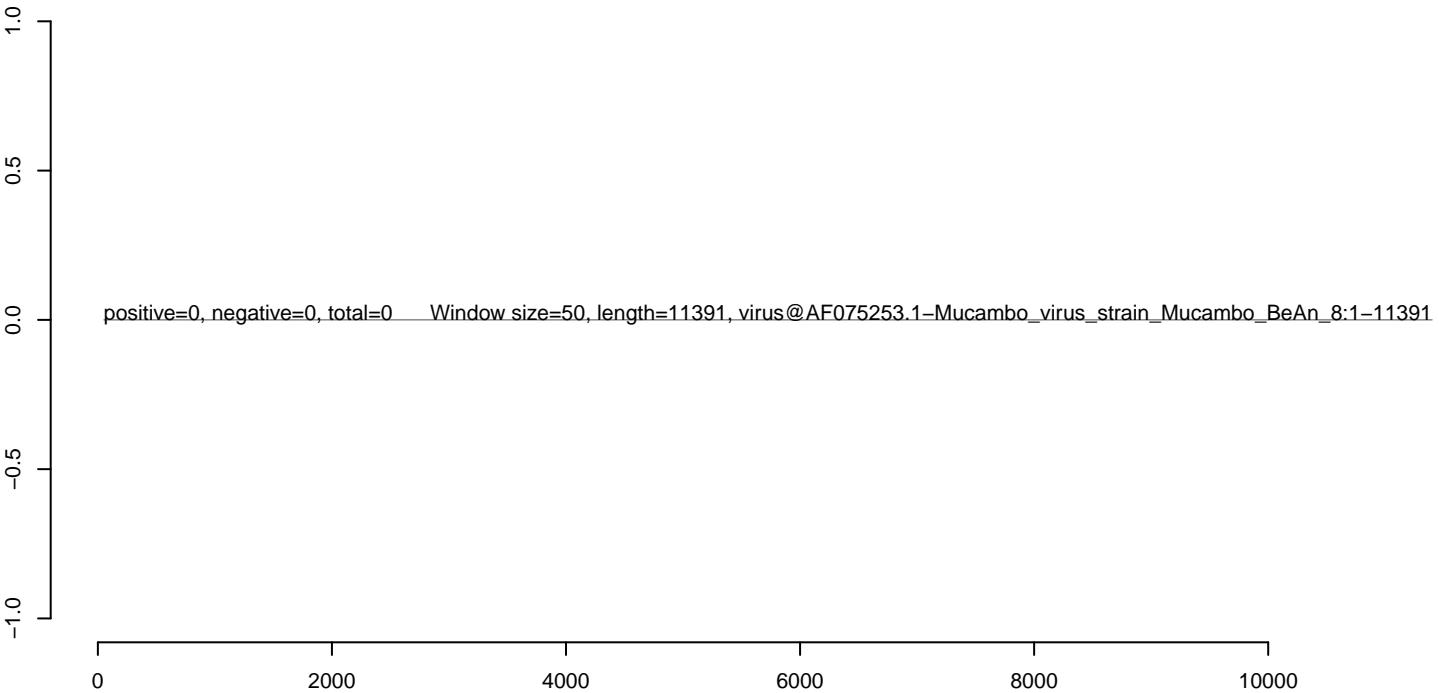
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

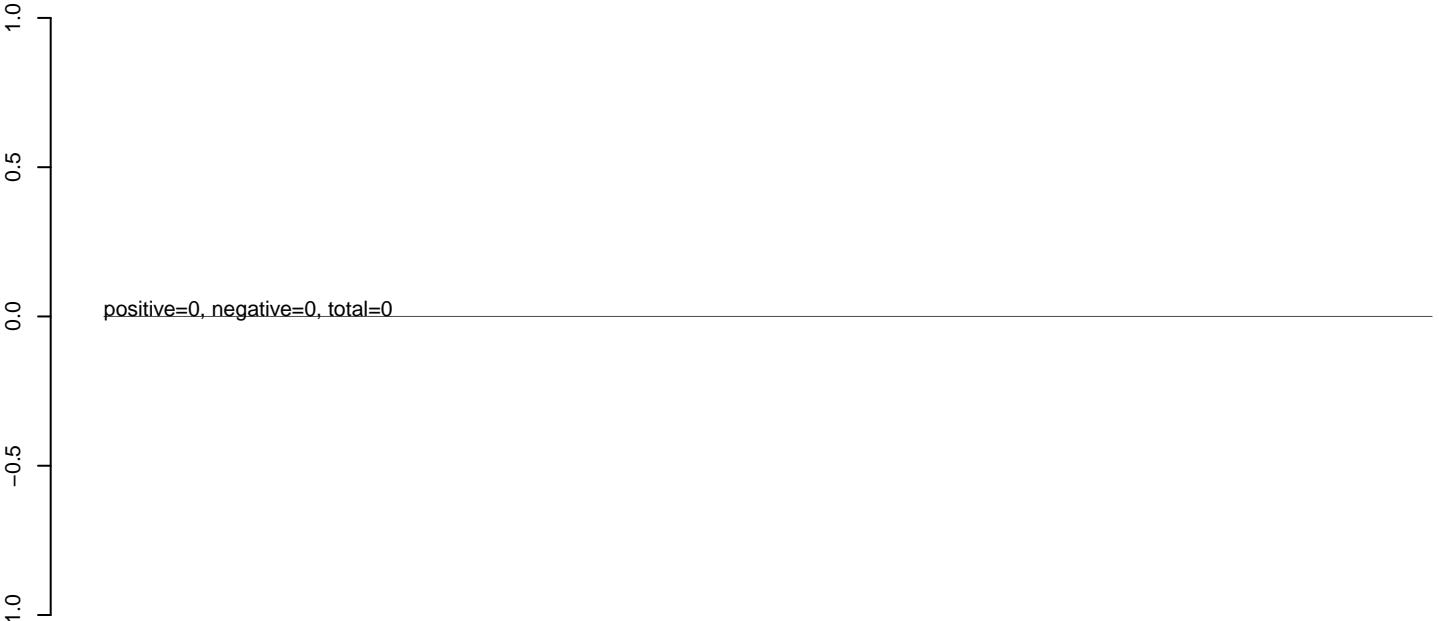
positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.rep

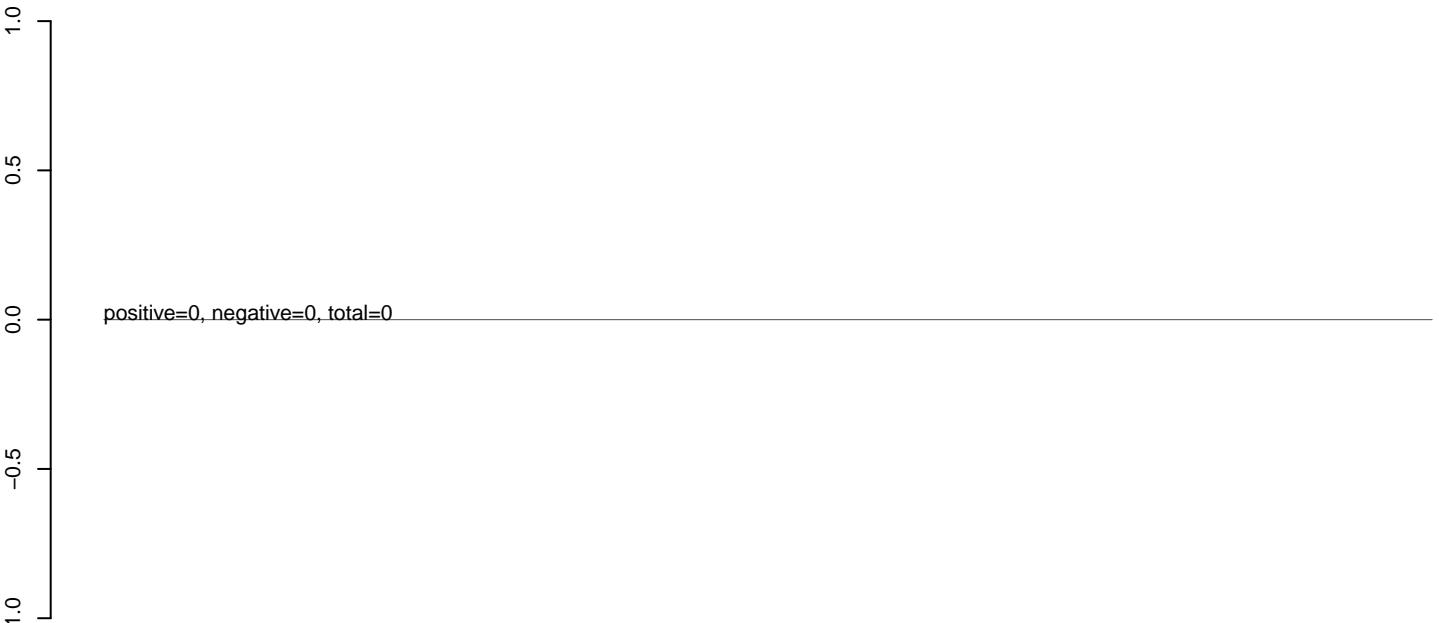
positive=0, negative=0, total=0    Window size=50, length=10837, virus@AB701776.1-Culex\_flavivirus\_RNA\_strain\_Toyama2627:1-10837

0 2000 4000 6000 8000 10000

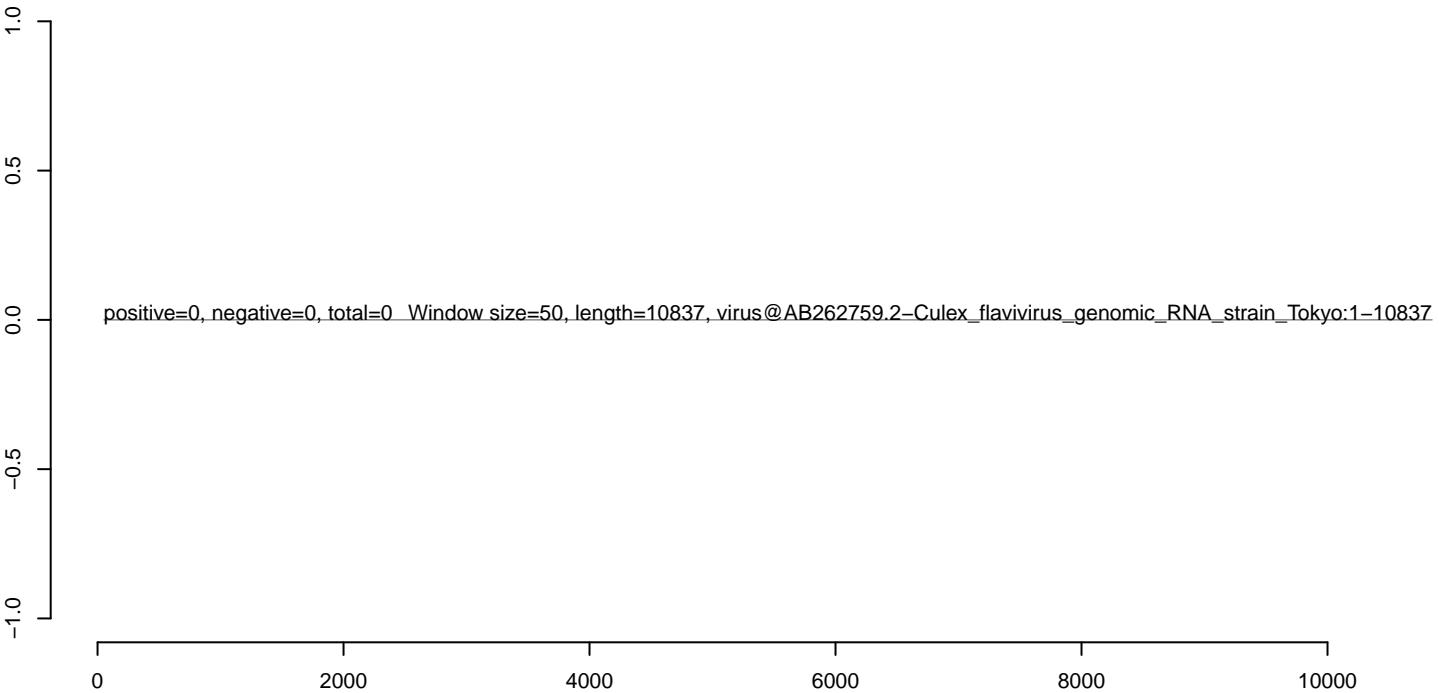
### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep



### AnGam\_Fcanc\_TN.18\_23.rep

positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.24\_35.rep

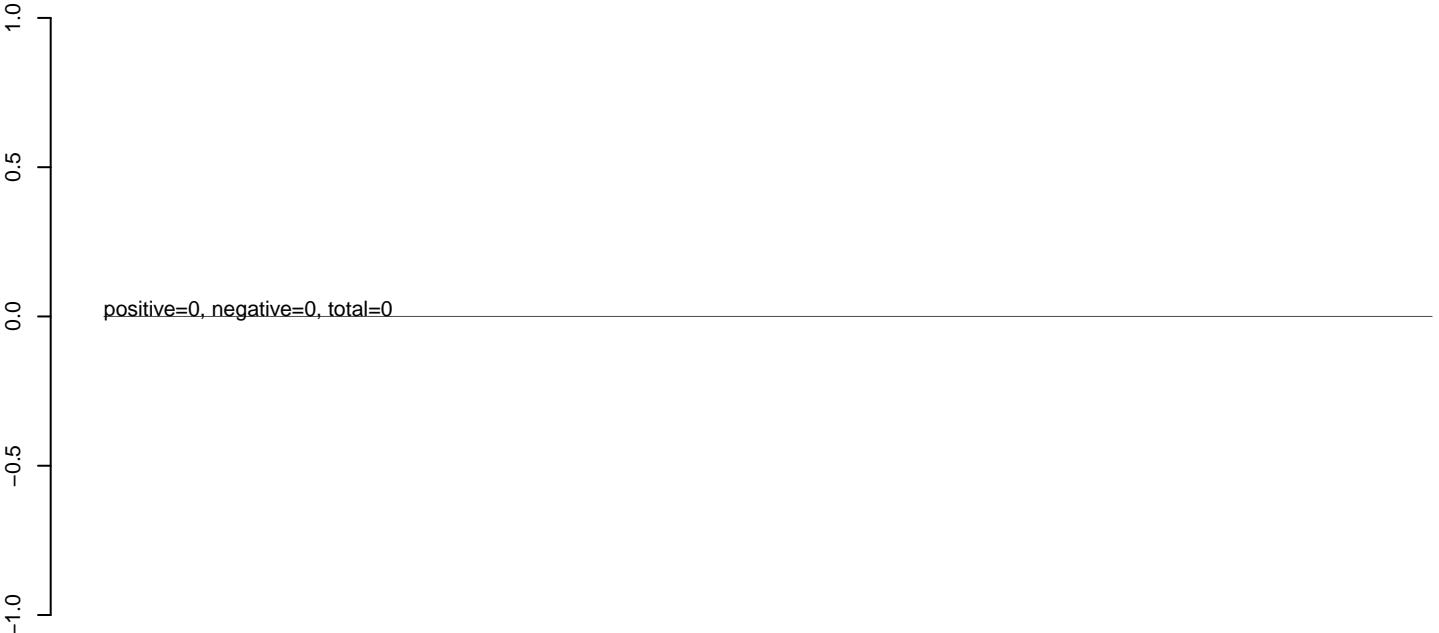
positive=0, negative=0, total=0

### AnGam\_Fcanc\_TN.rep

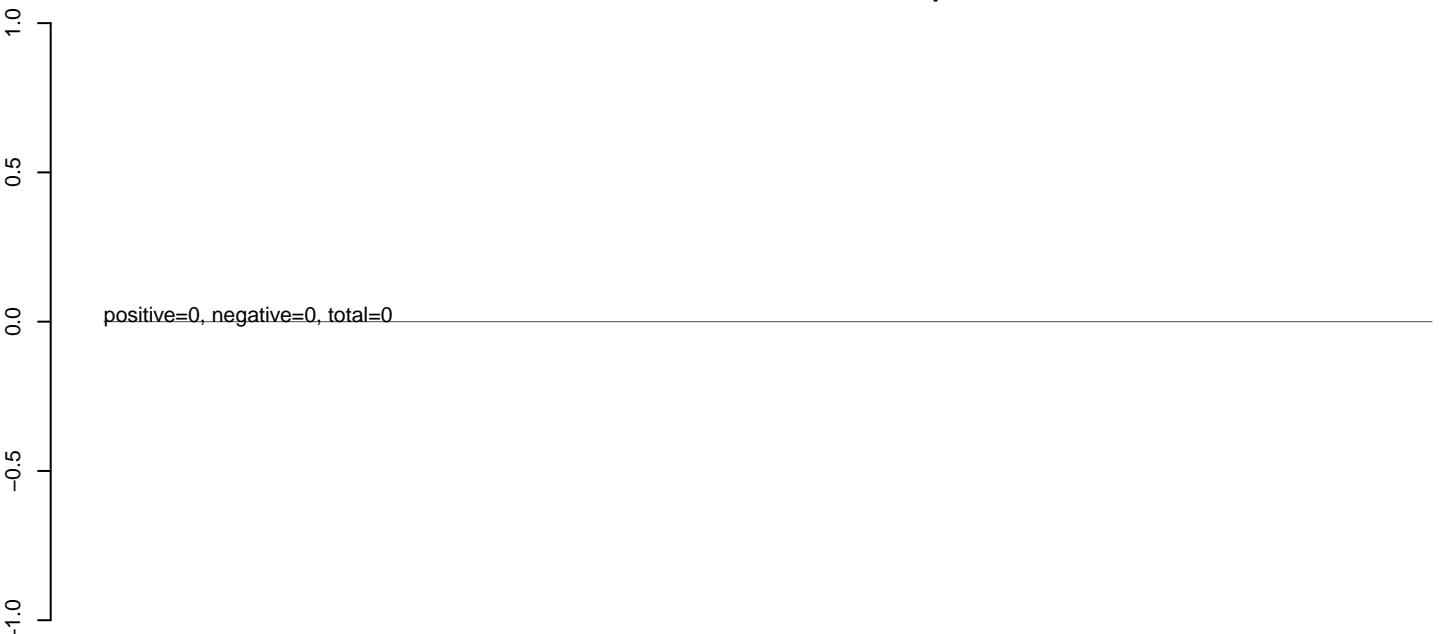
positive=0, negative=0, total=0    Window.size=50, length=3157, Virus@MH188043.1-Culex\_densovirus\_strain\_CDenv\_Madera-2:1-3157

0 500 1000 1500 2000 2500 3000

### AnGam\_Fcanc\_TN.18\_23.rep



### AnGam\_Fcanc\_TN.24\_35.rep



### AnGam\_Fcanc\_TN.rep

