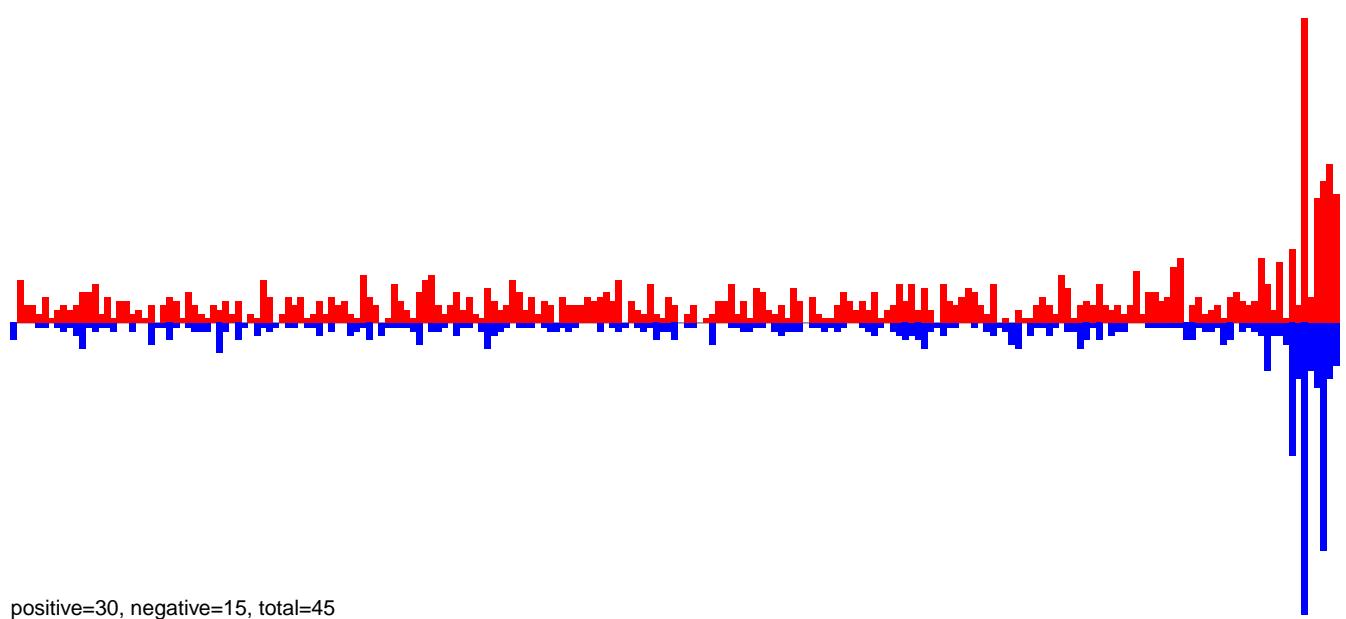
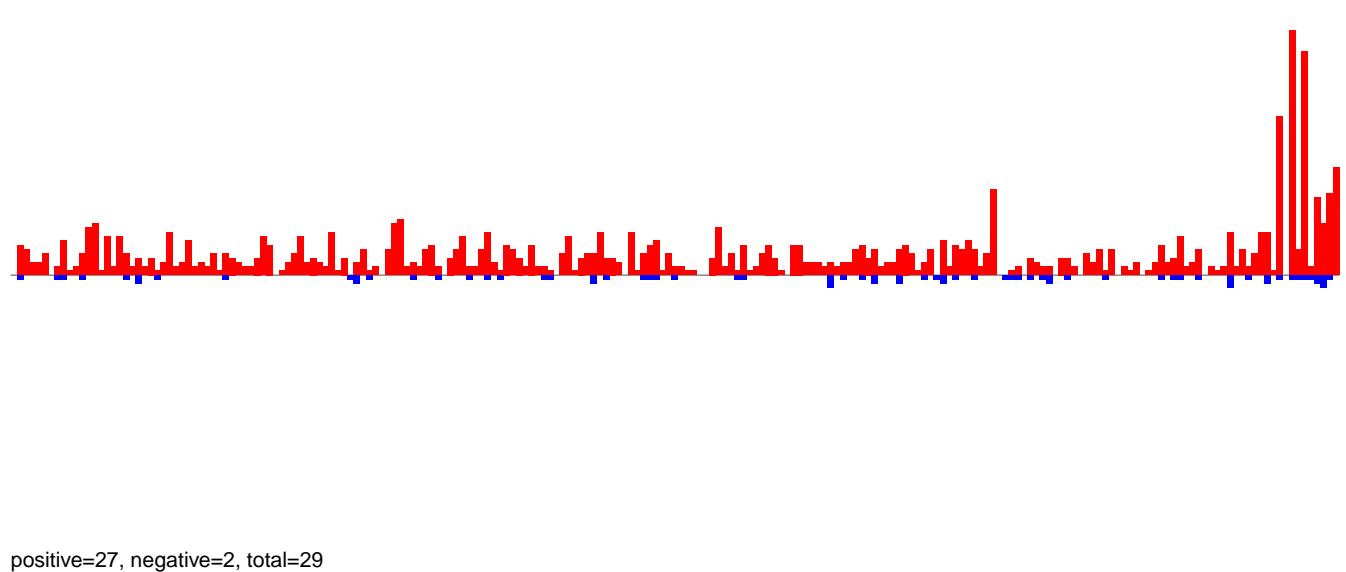


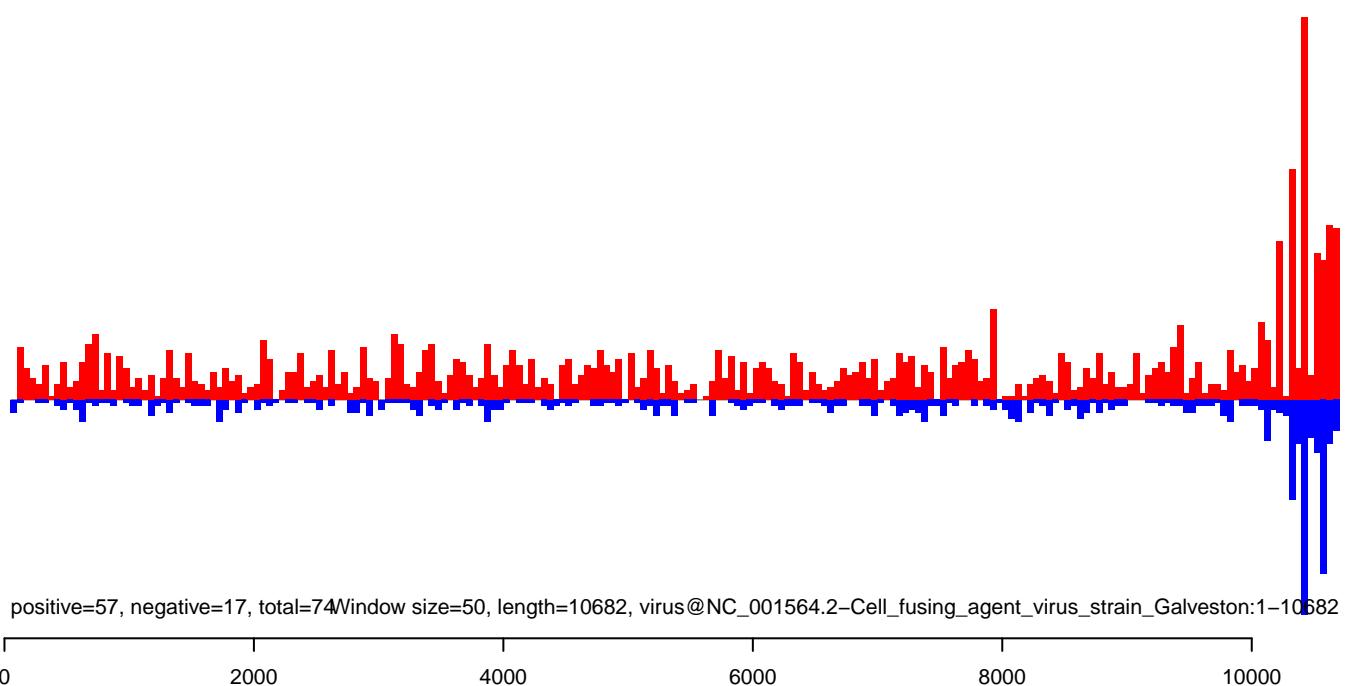
### AeAeg\_Fcanc\_GH.18\_23.rep



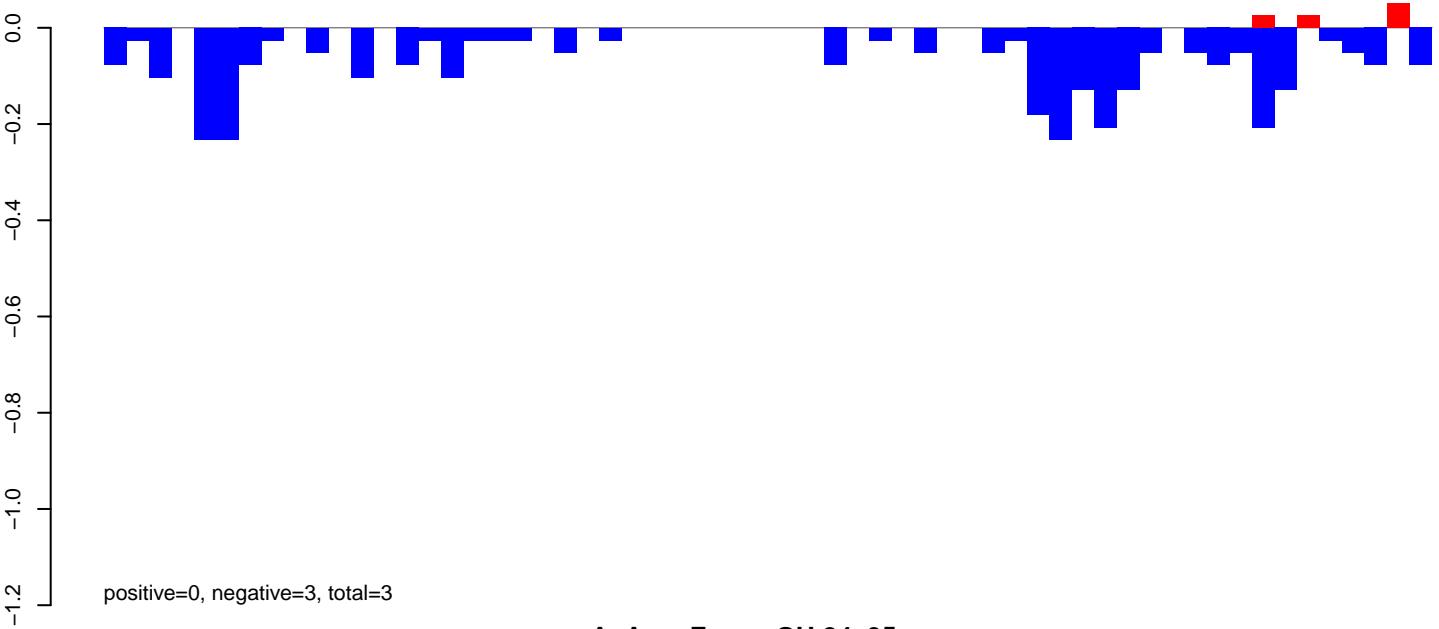
### AeAeg\_Fcanc\_GH.24\_35.rep



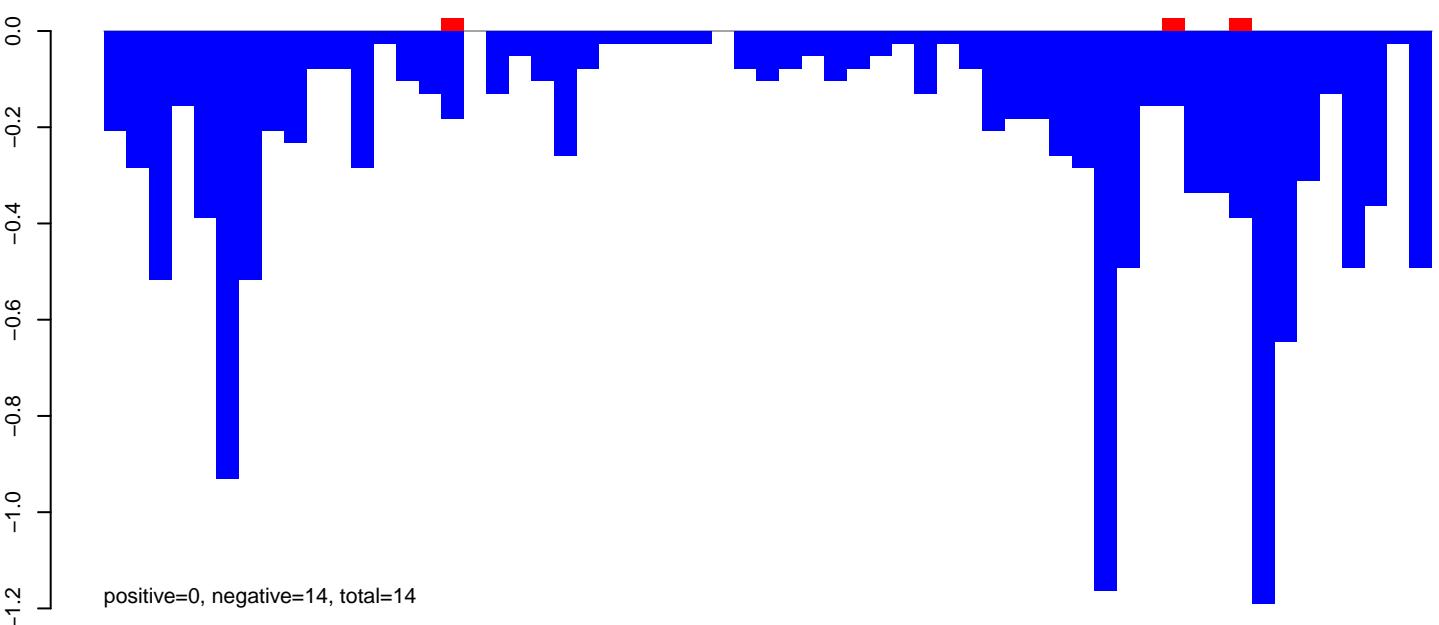
### AeAeg\_Fcanc\_GH.rep



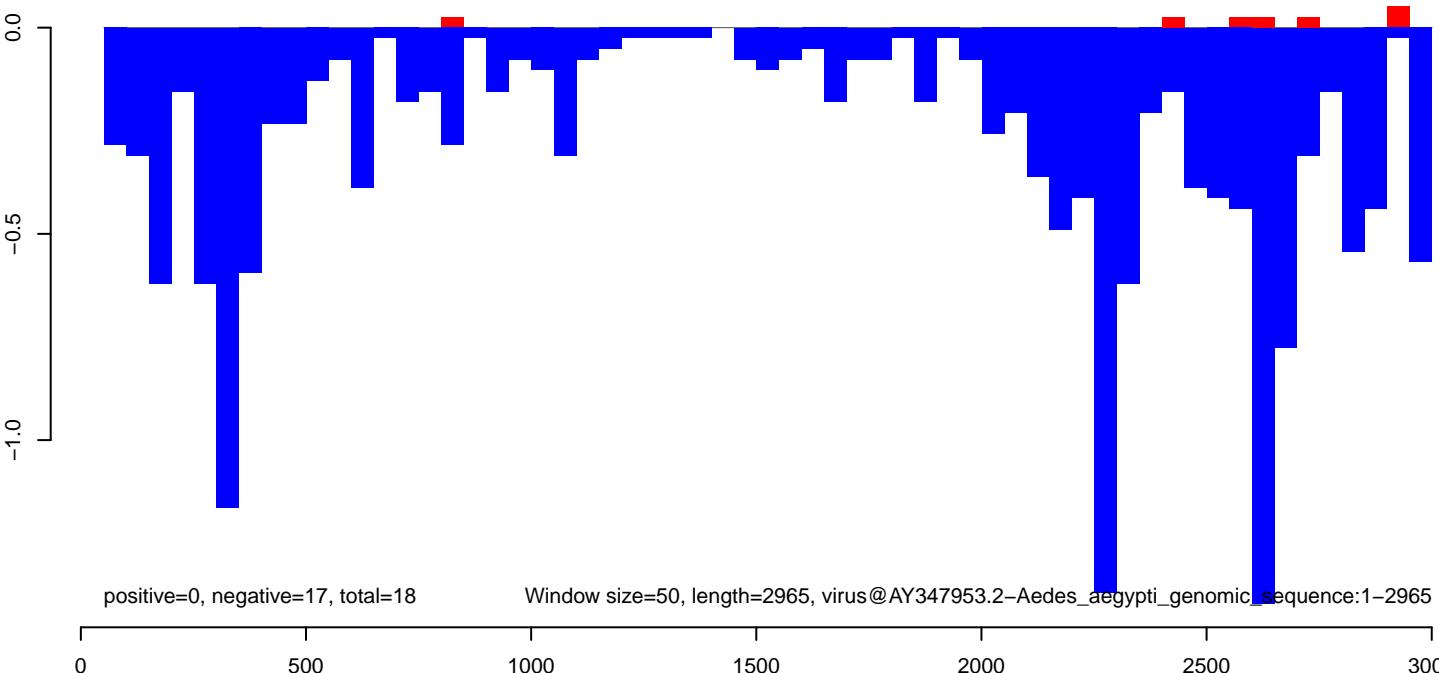
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep

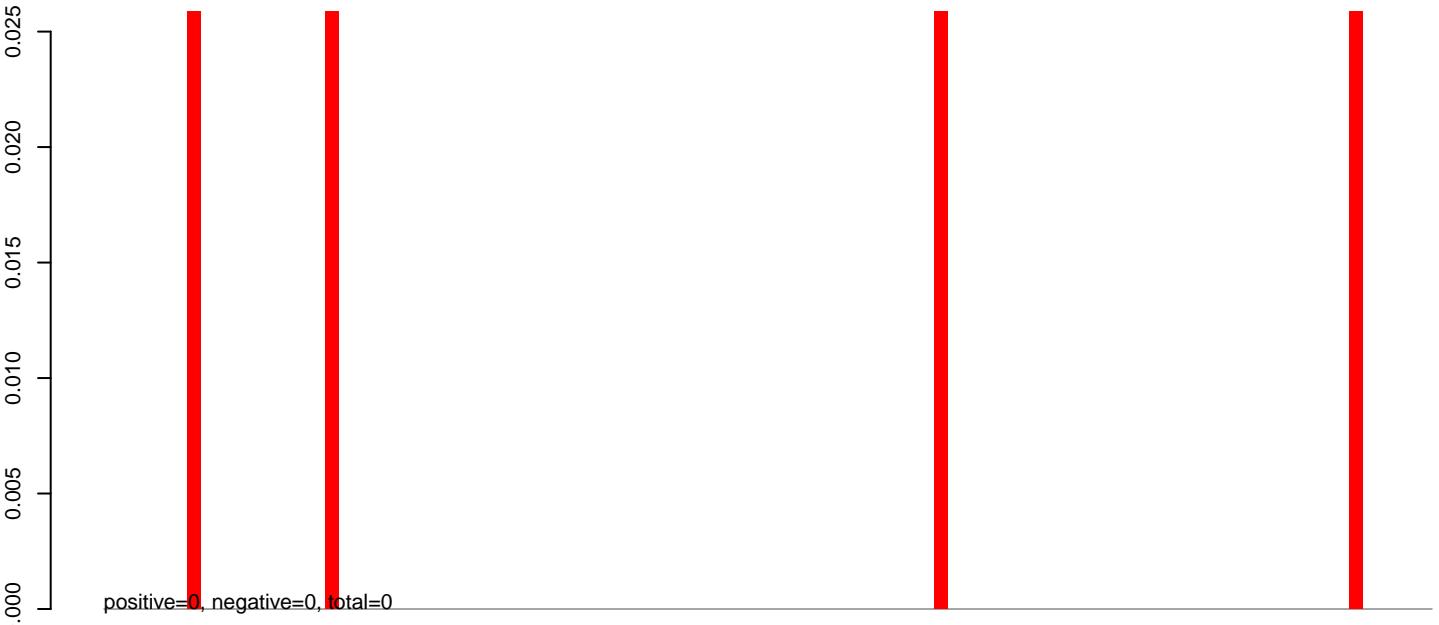


### AeAeg\_Fcanc\_GH.rep



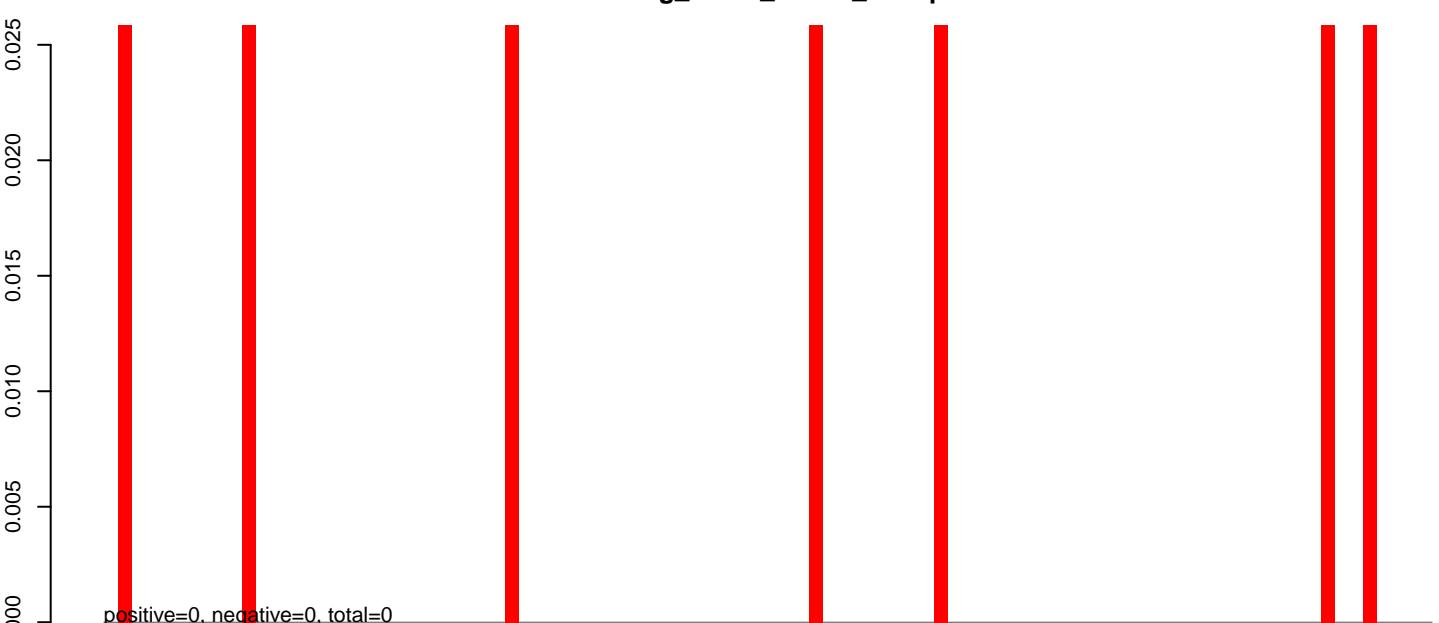
AeAeg\_Fcarc\_GH.18\_23.rep

positive=0, negative=0, total=0



AeAeg\_Fcarc\_GH.24\_35.rep

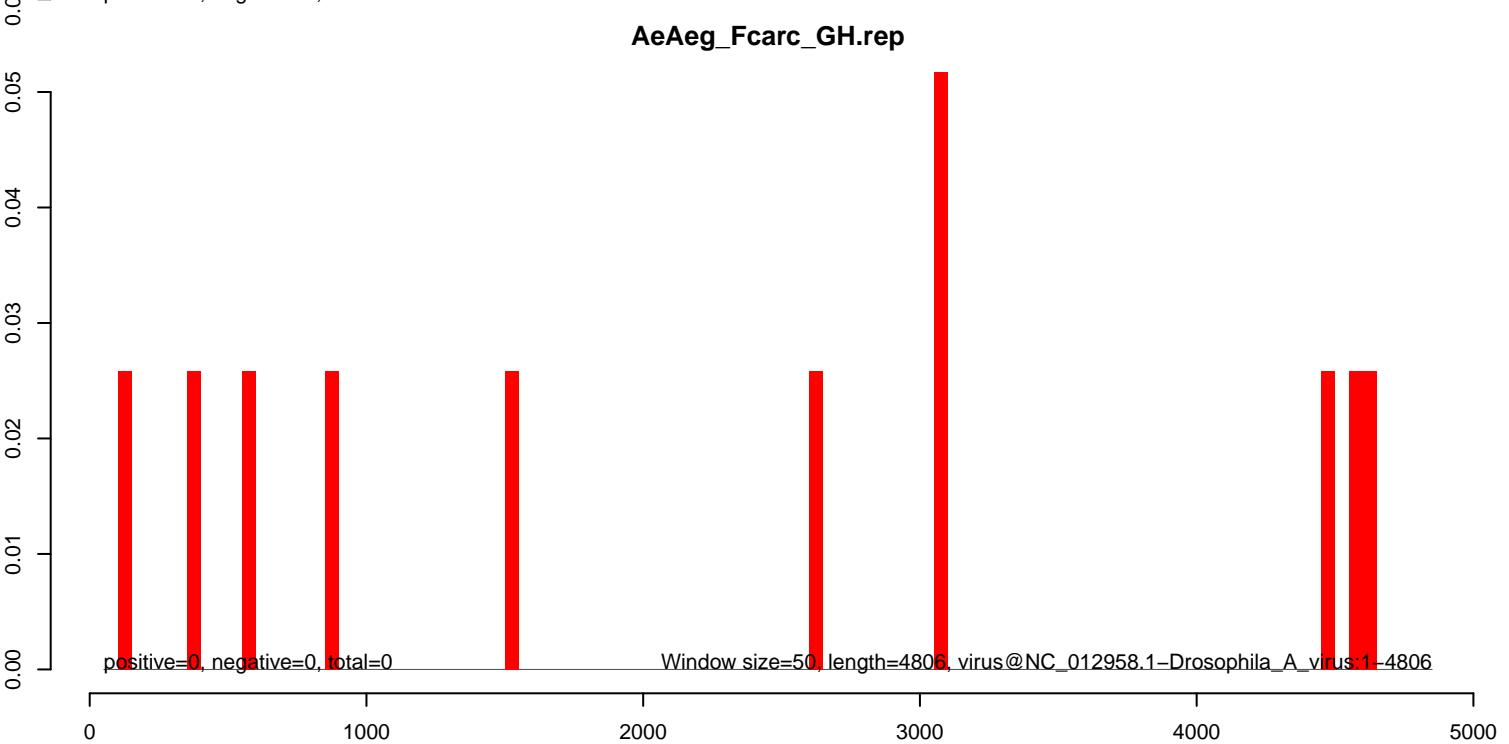
positive=0, negative=0, total=0



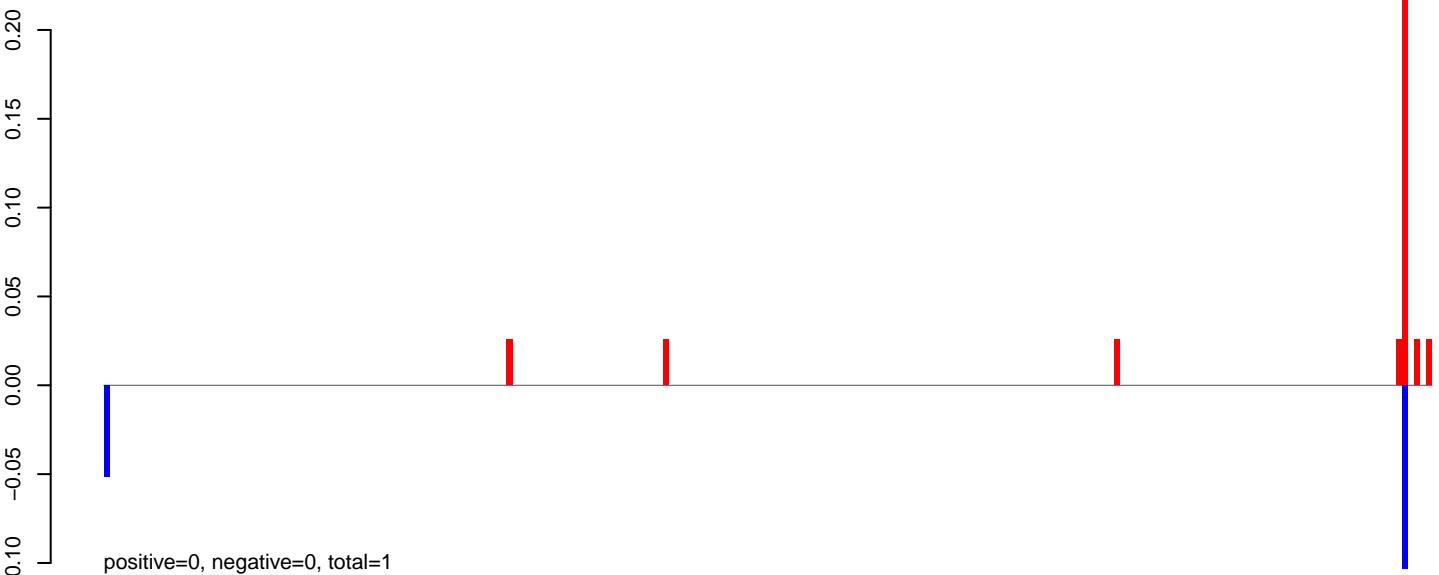
AeAeg\_Fcarc\_GH.rep

positive=0, negative=0, total=0

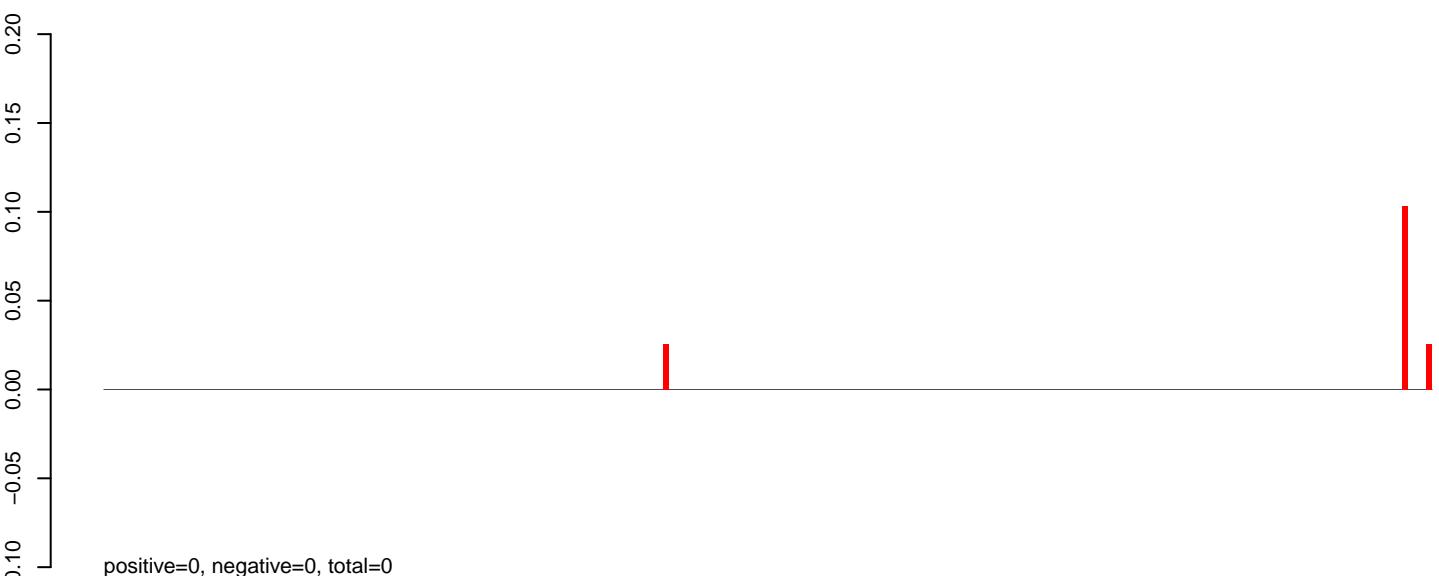
Window size=50, length=4806, virus@NC\_012958.1-Drosophila\_A\_virus\_1-4806



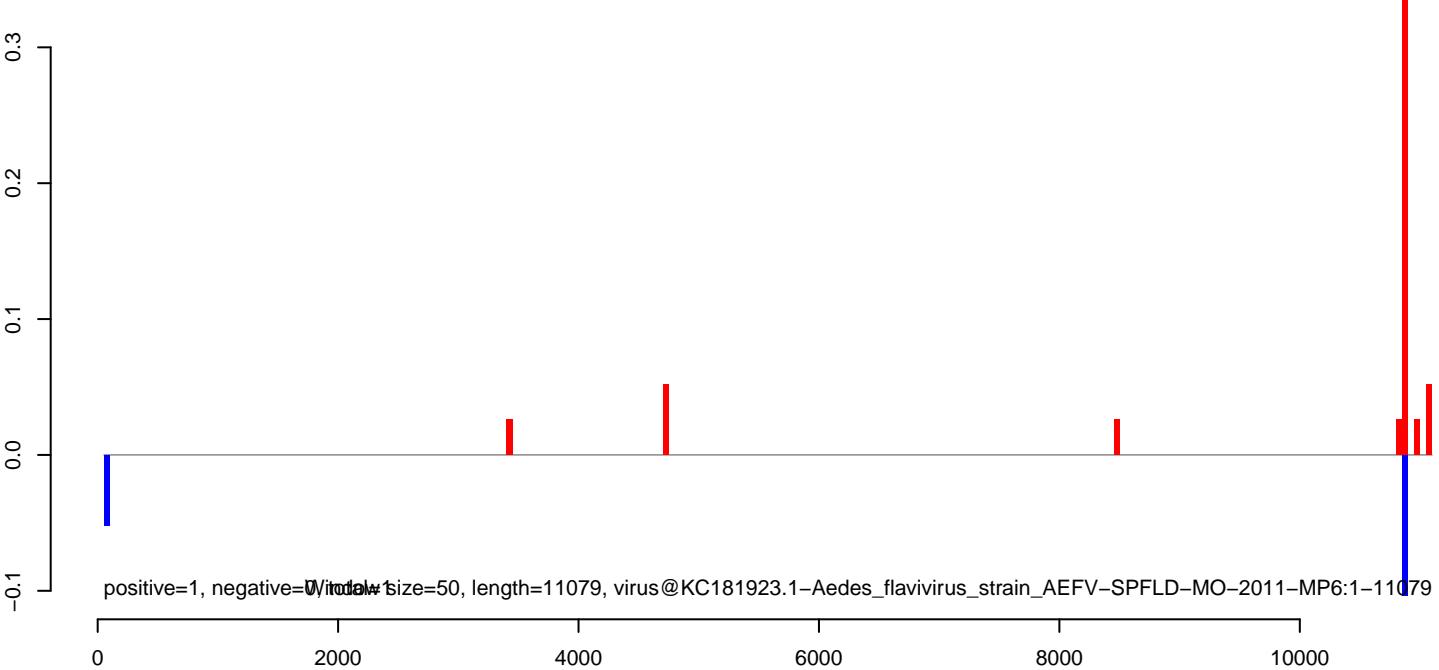
### AeAeg\_Fcanc\_GH.18\_23.rep



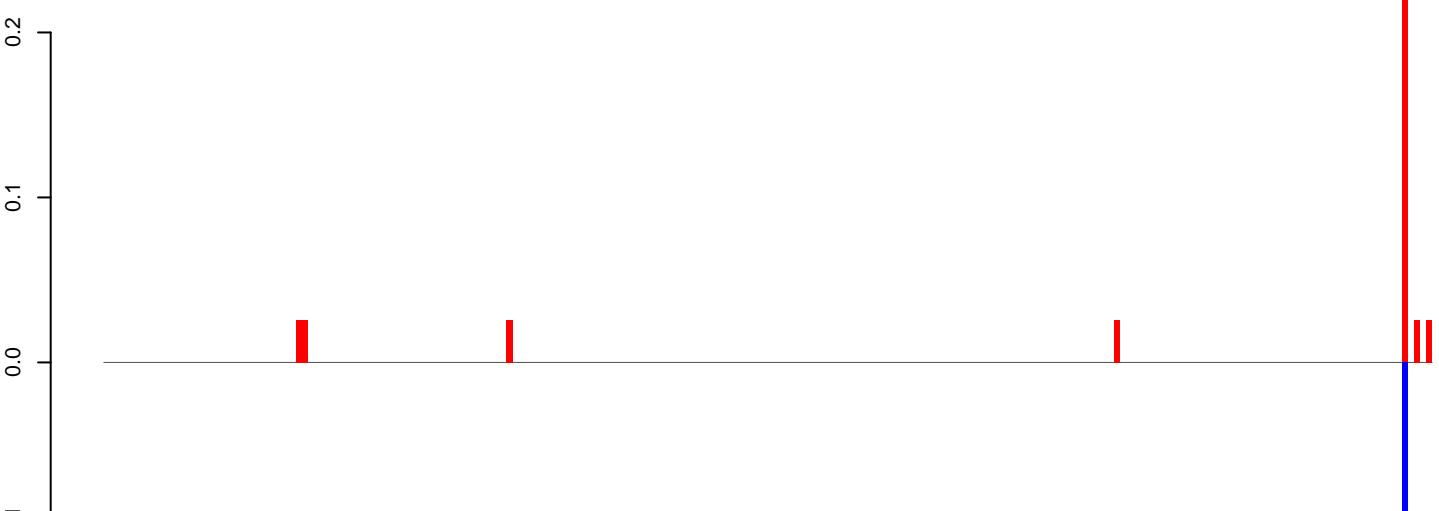
### AeAeg\_Fcanc\_GH.24\_35.rep



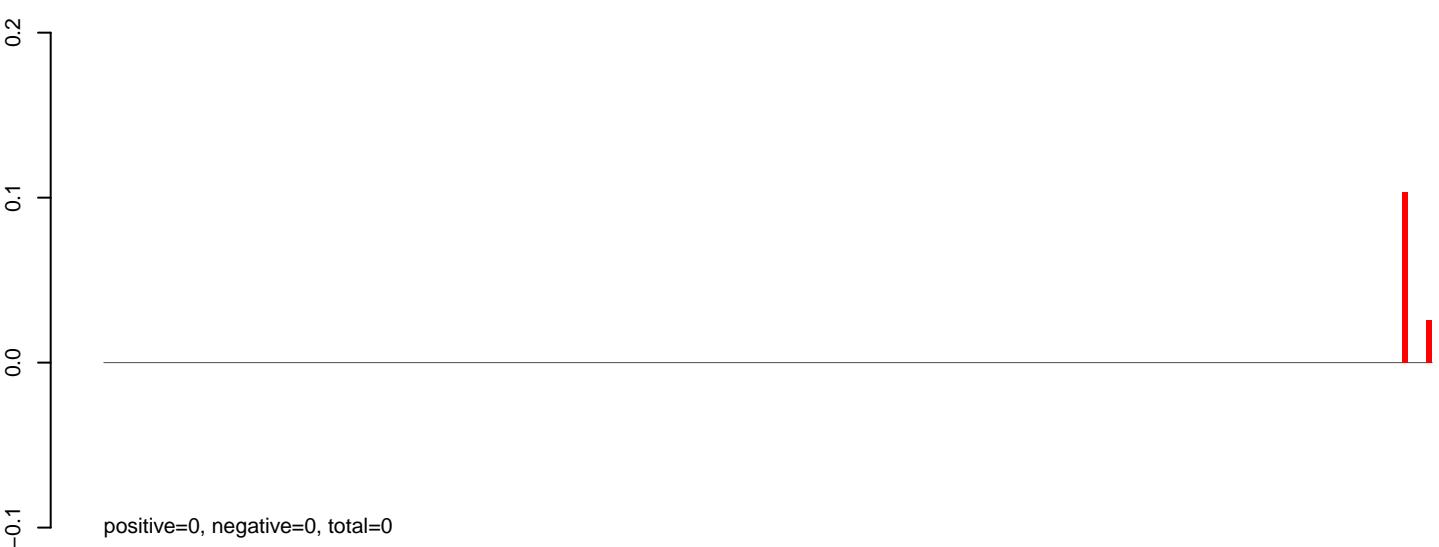
### AeAeg\_Fcanc\_GH.rep



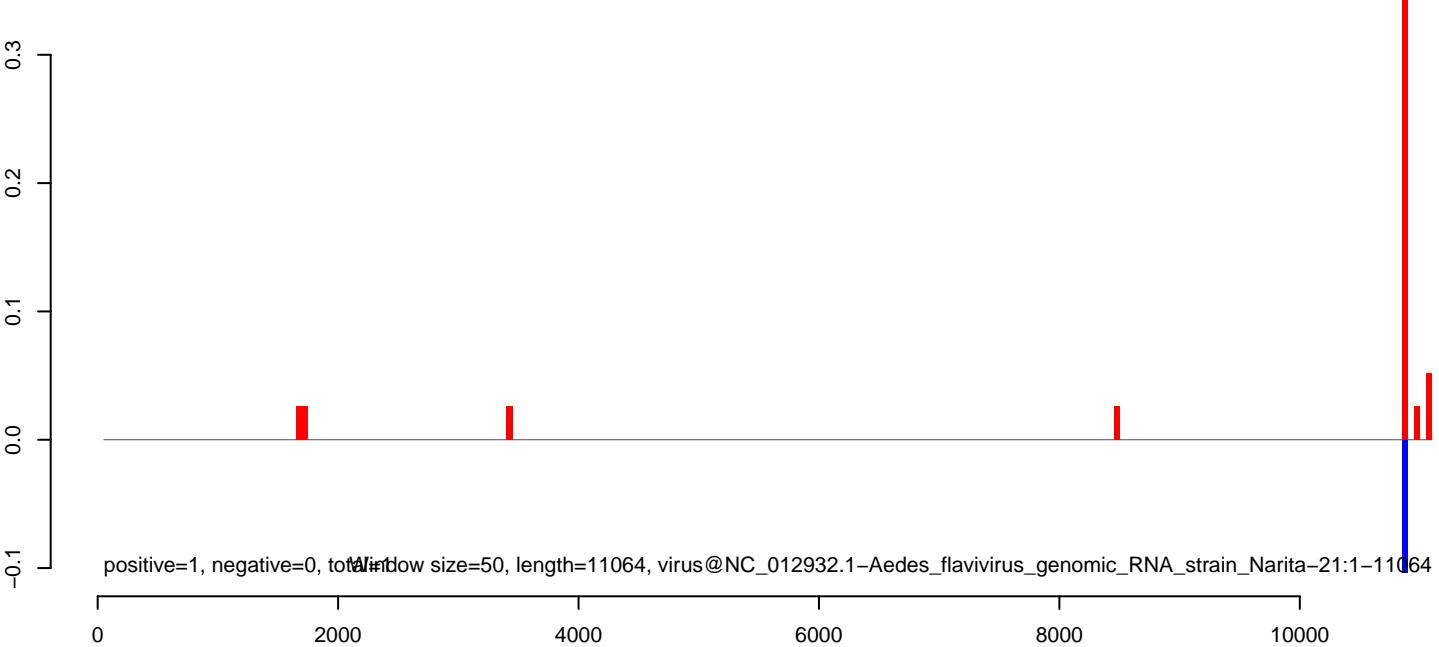
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep

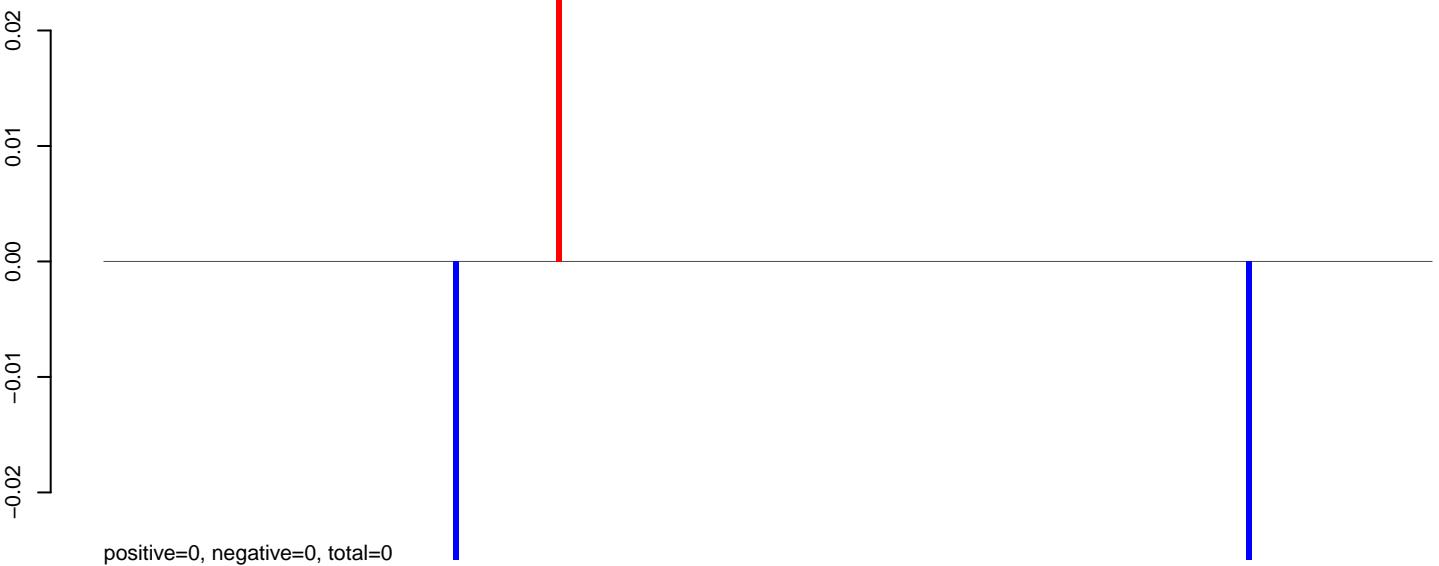


### AeAeg\_Fcanc\_GH.rep



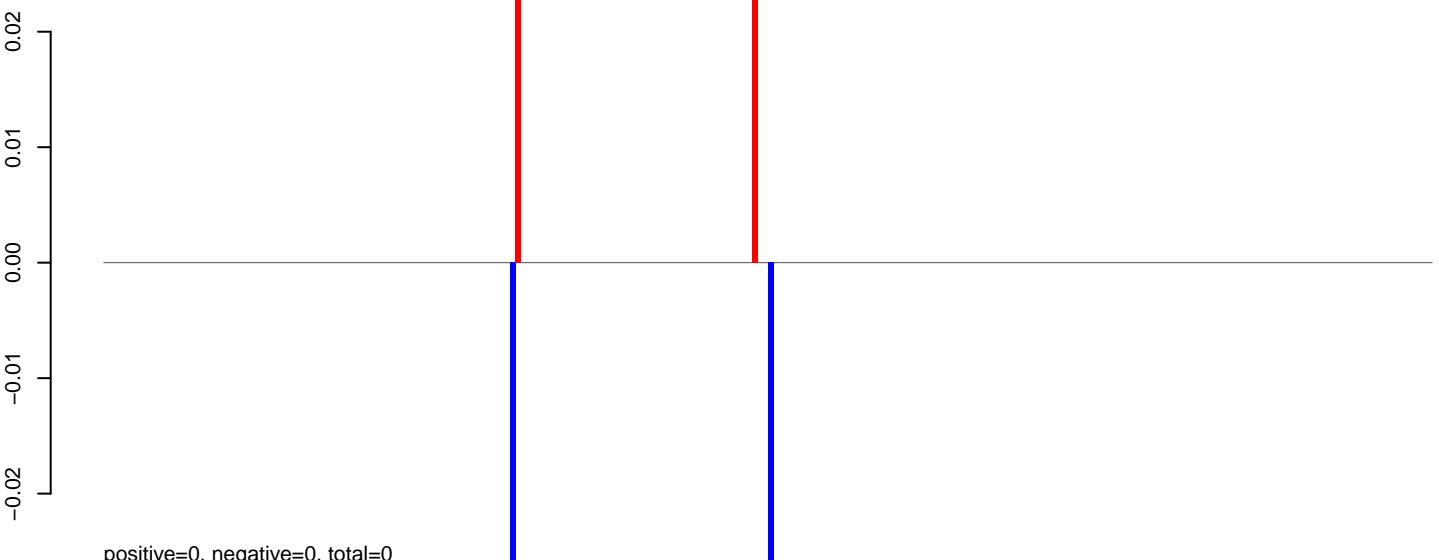
### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

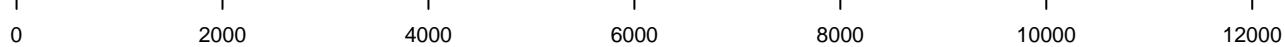
positive=0, negative=0, total=0



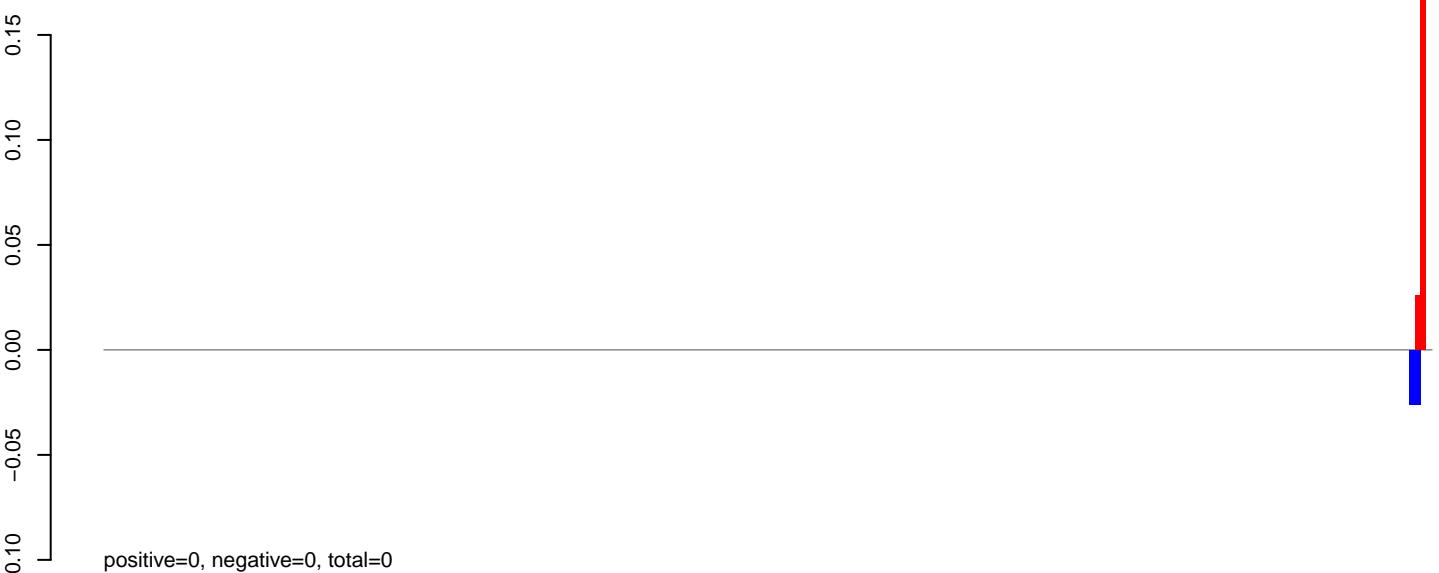
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

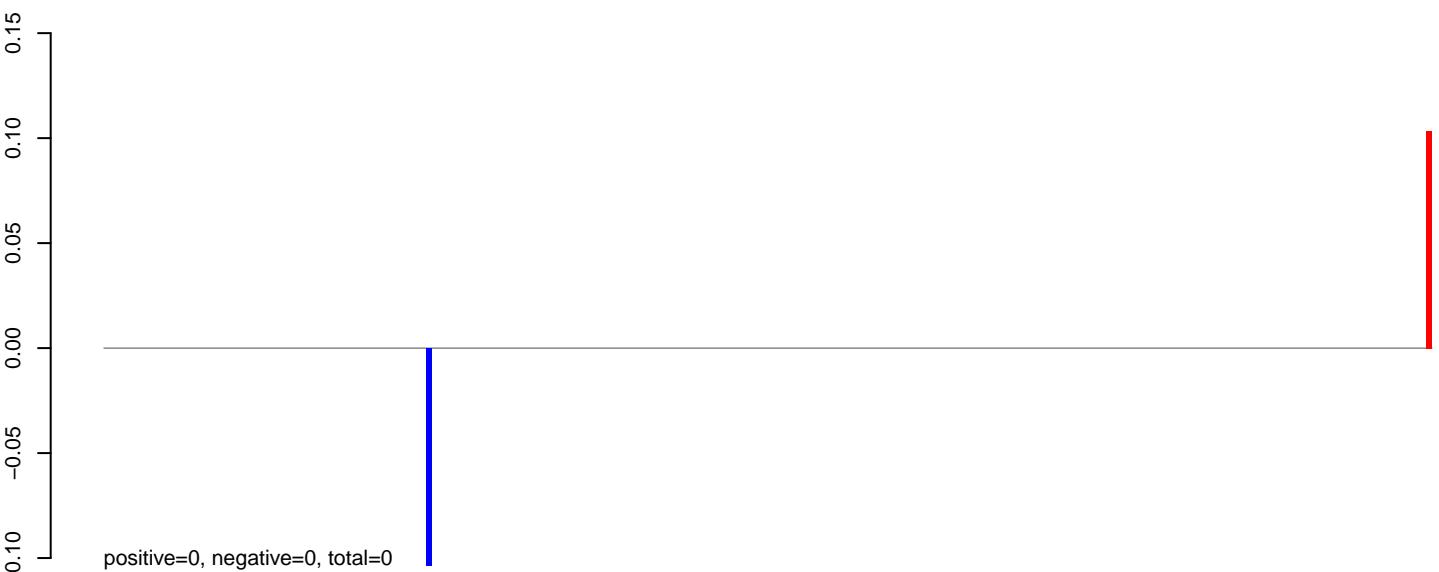
Window size=50, length=12916, virus@MH237595.1-Aedes\_aegypti\_anphevirus, strain\_1a:1-12916



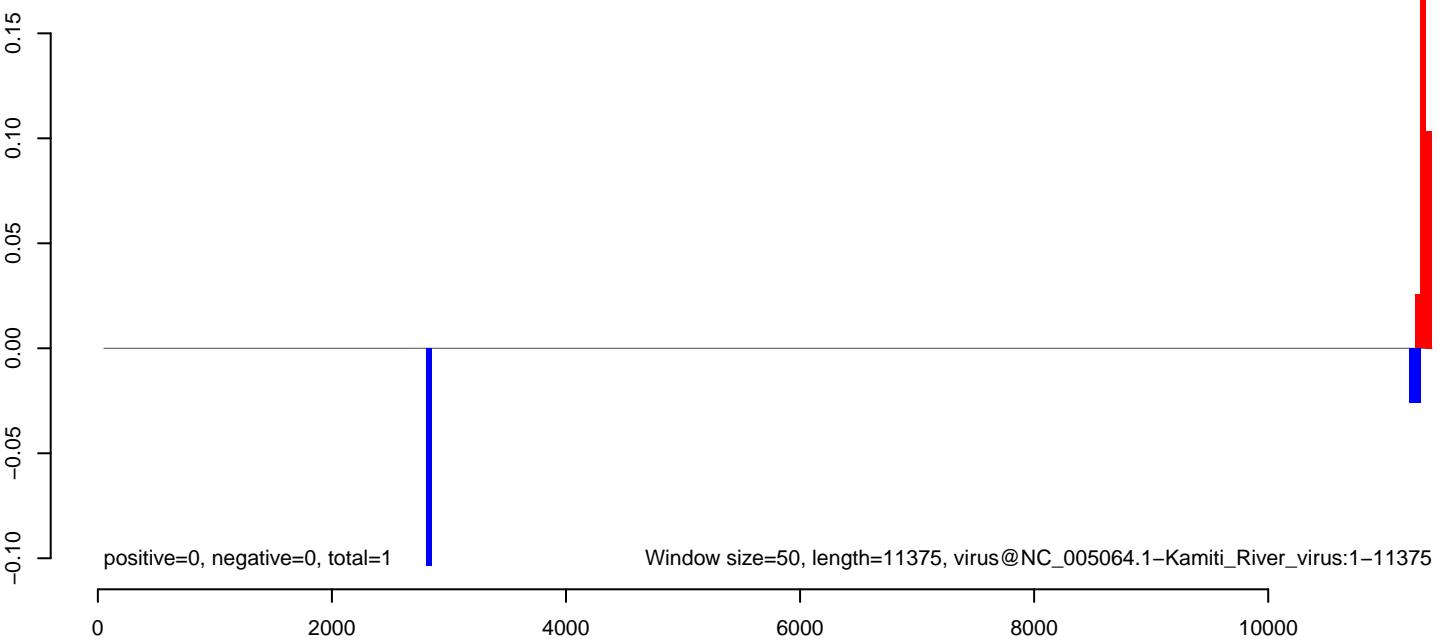
### AeAeg\_Fcanc\_GH.18\_23.rep



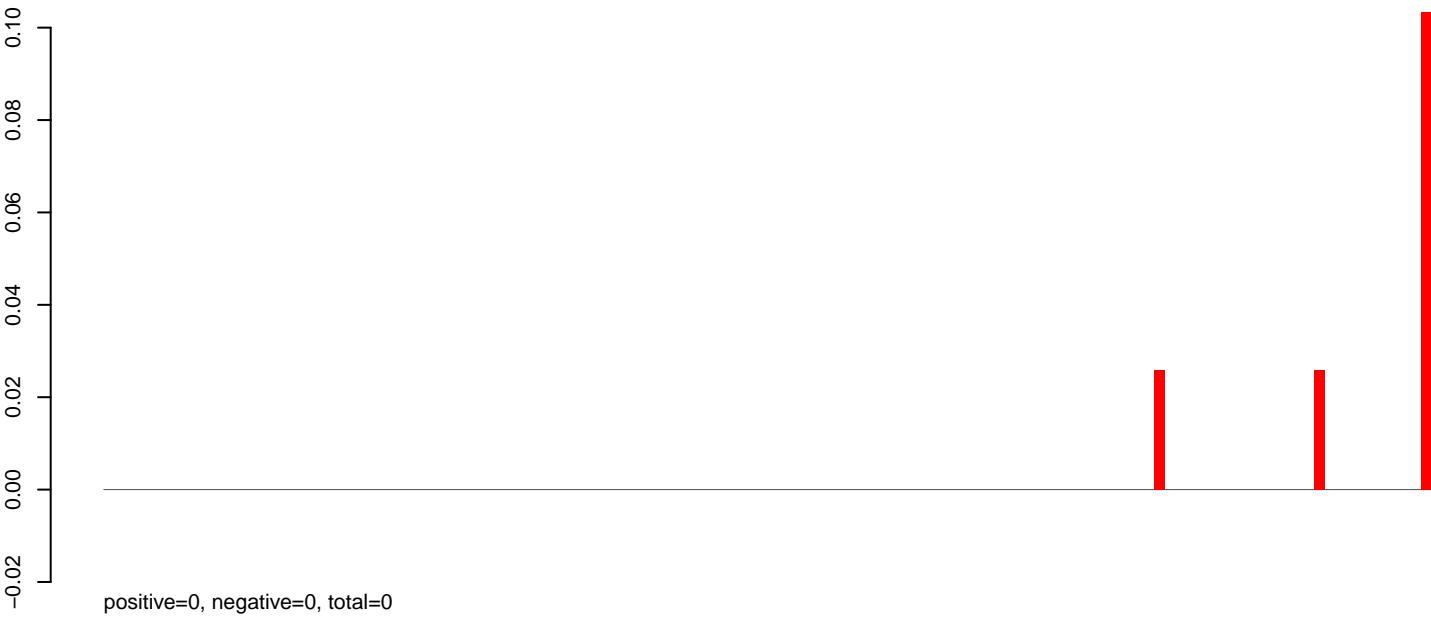
### AeAeg\_Fcanc\_GH.24\_35.rep



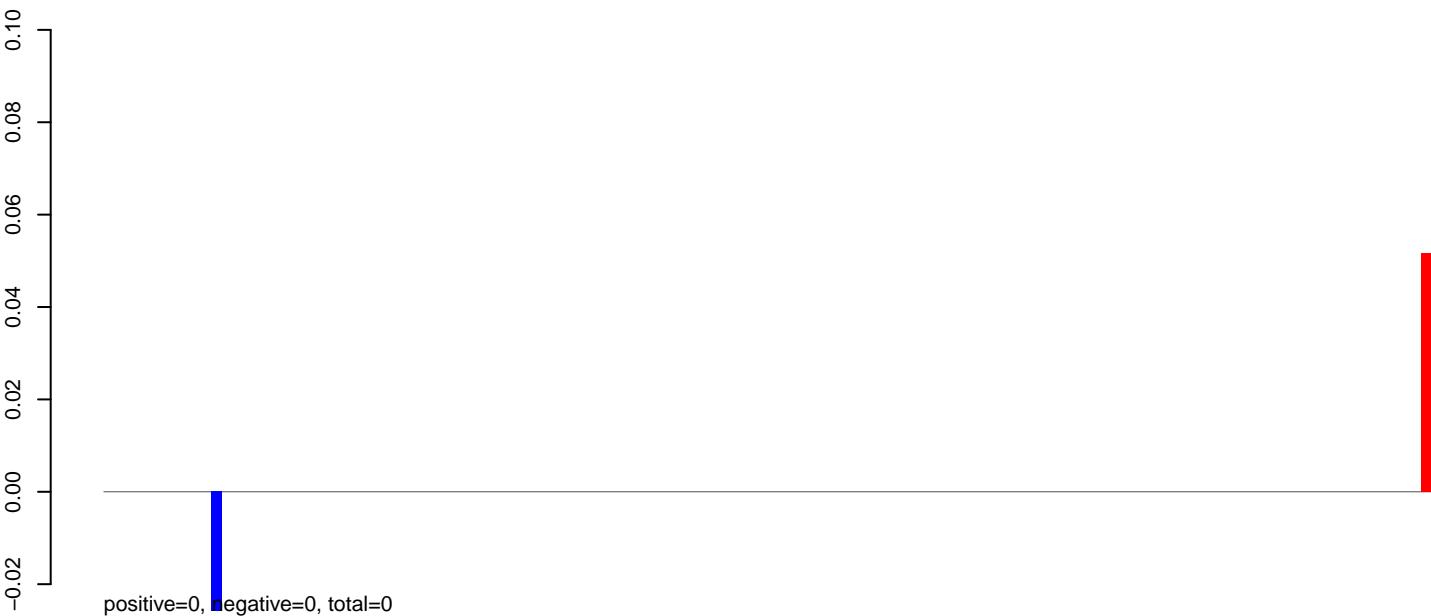
### AeAeg\_Fcanc\_GH.rep



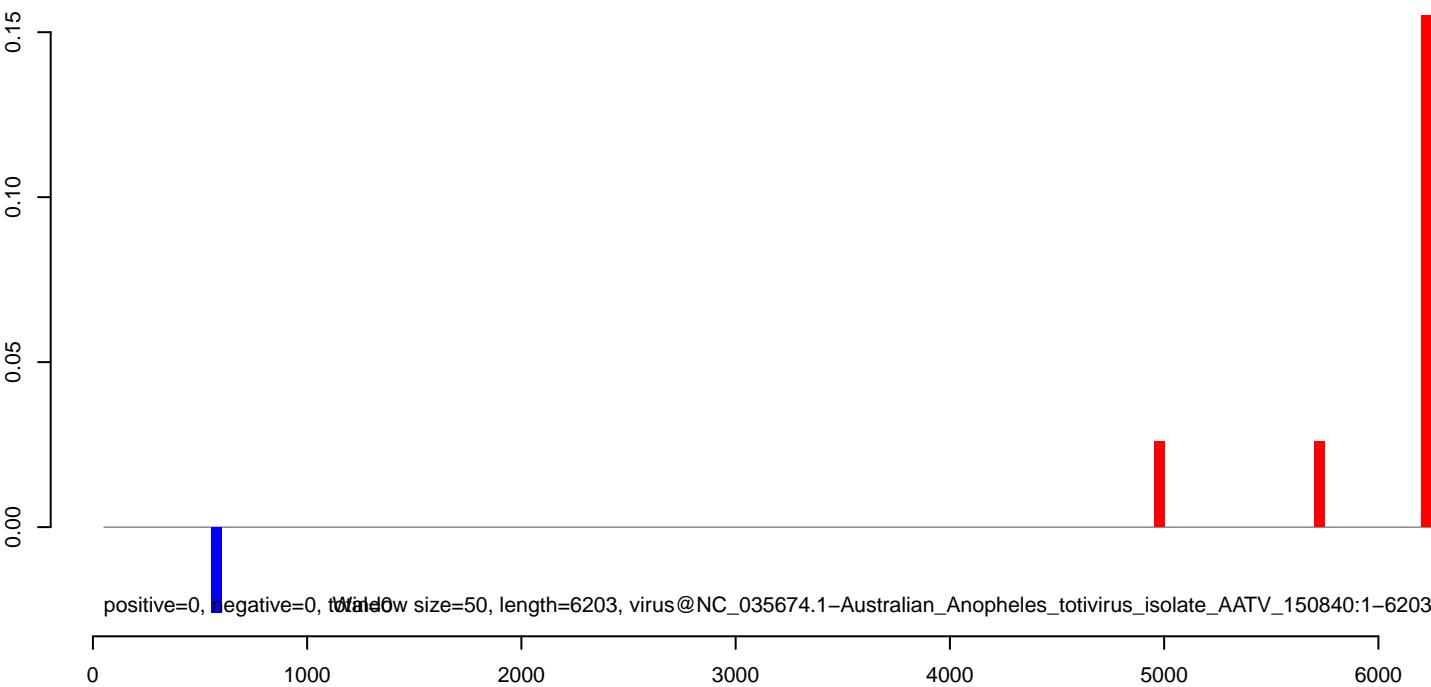
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



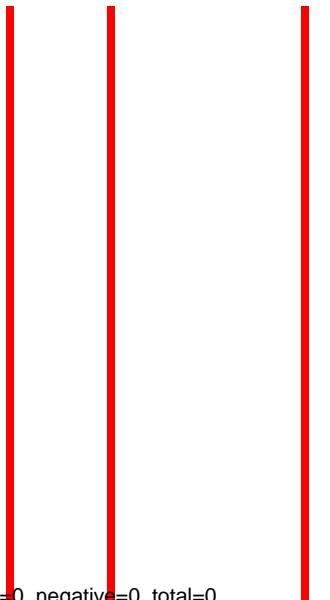
### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

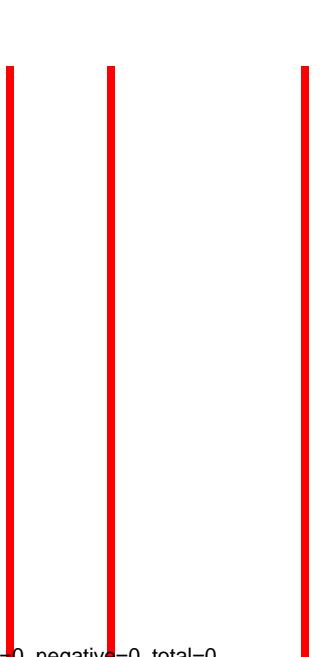
positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

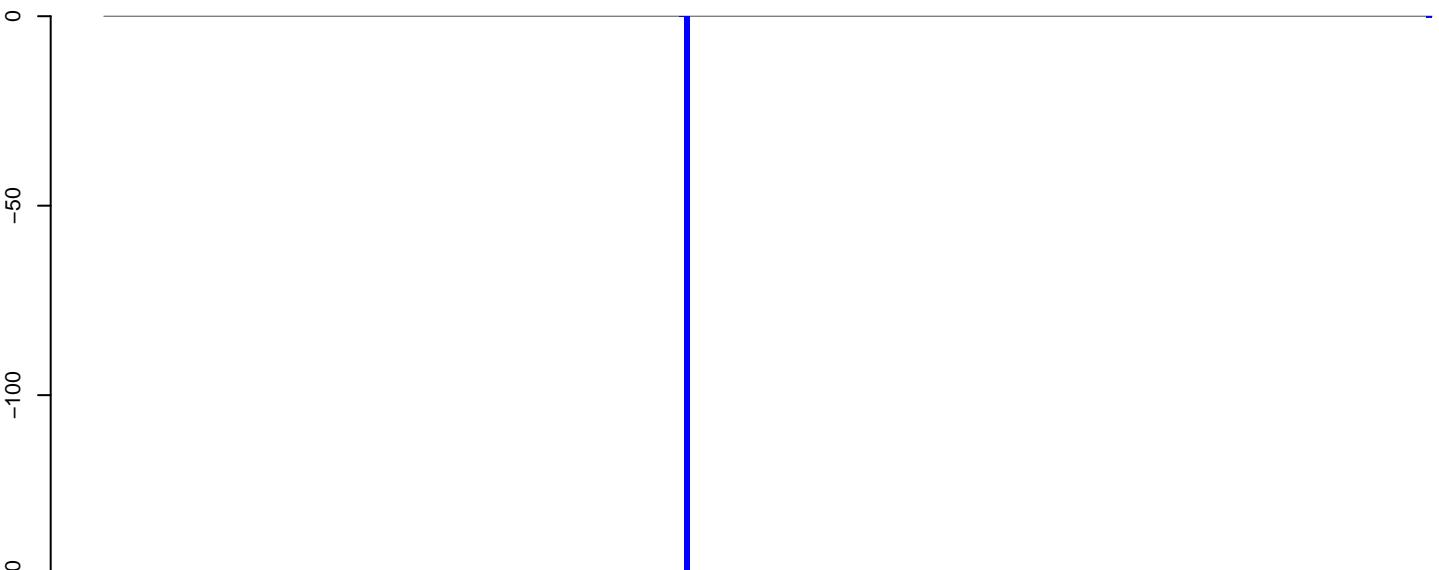
positive=0, negative=0, total=0

Window size=50, length=9264, virus=@NC\_001834.1-Drosophila\_C\_virus\_strain\_EB:1-9264

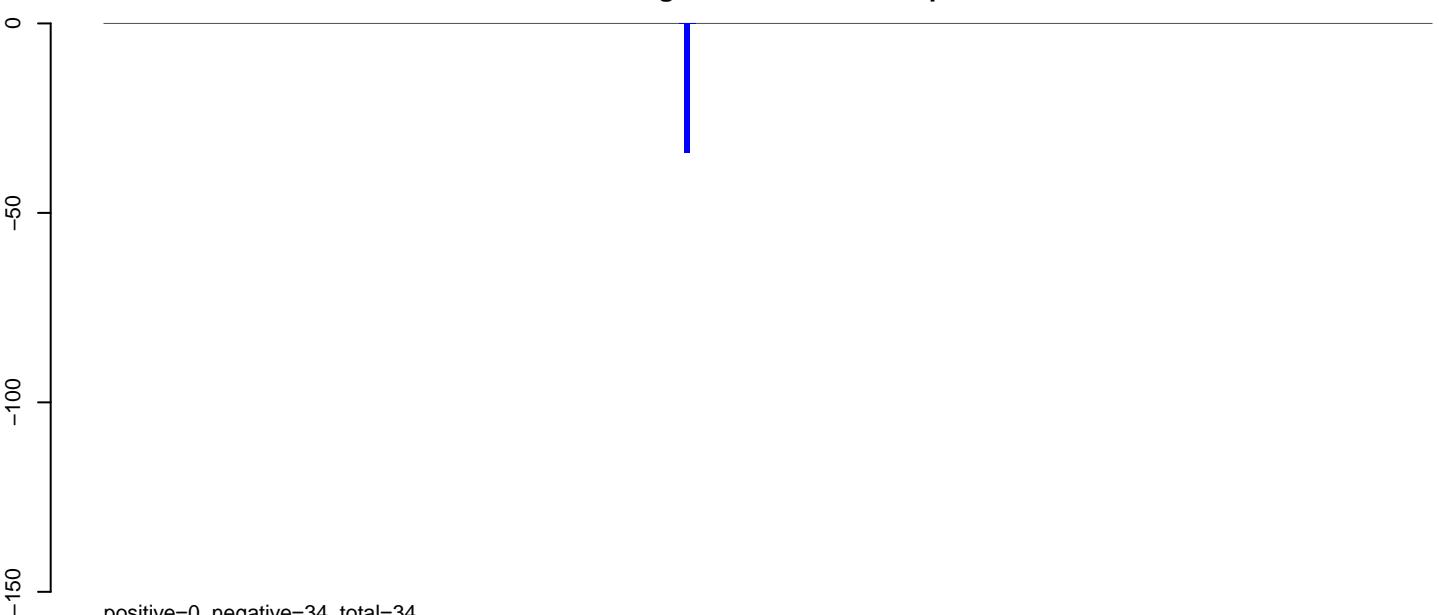


0 2000 4000 6000 8000

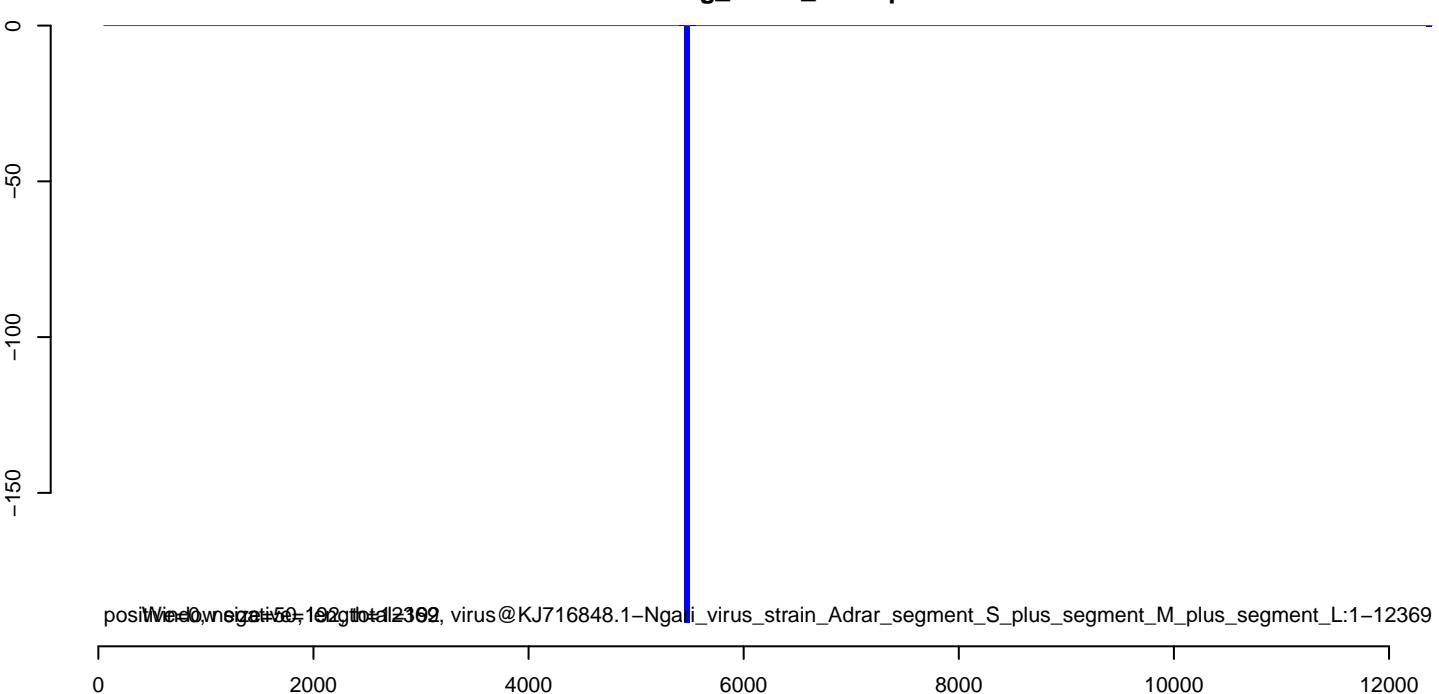
### AeAeg\_Fcanc\_GH.18\_23.rep



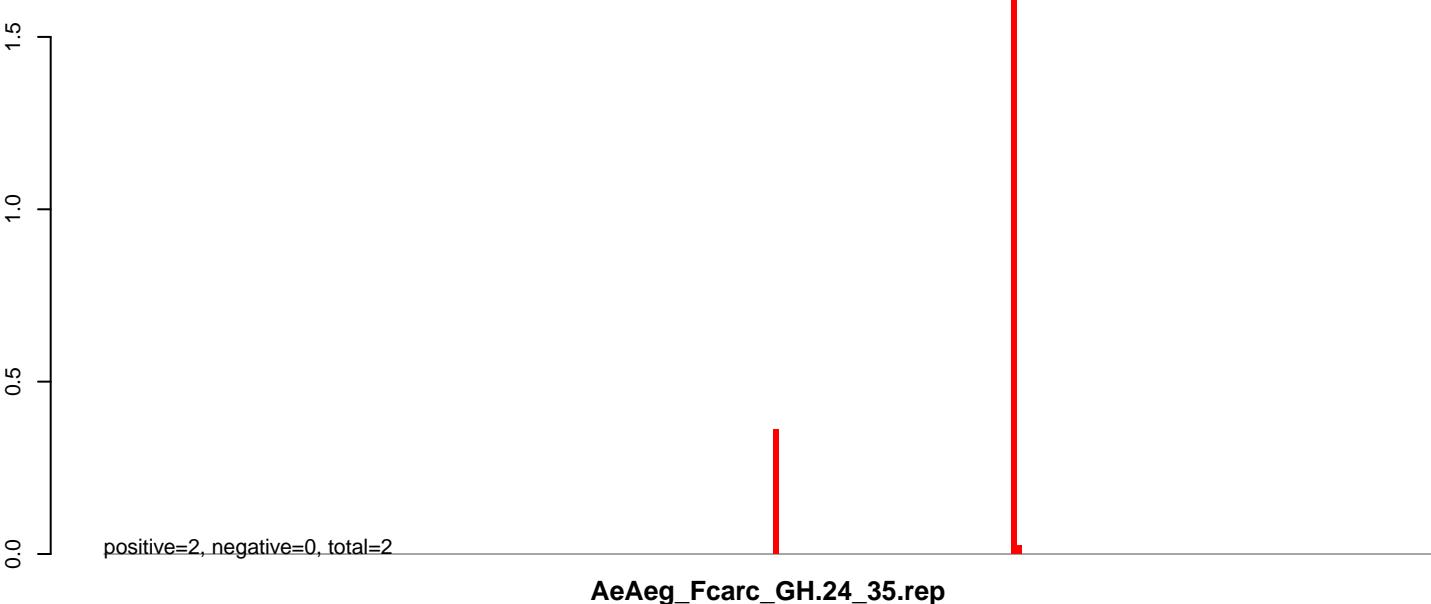
### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



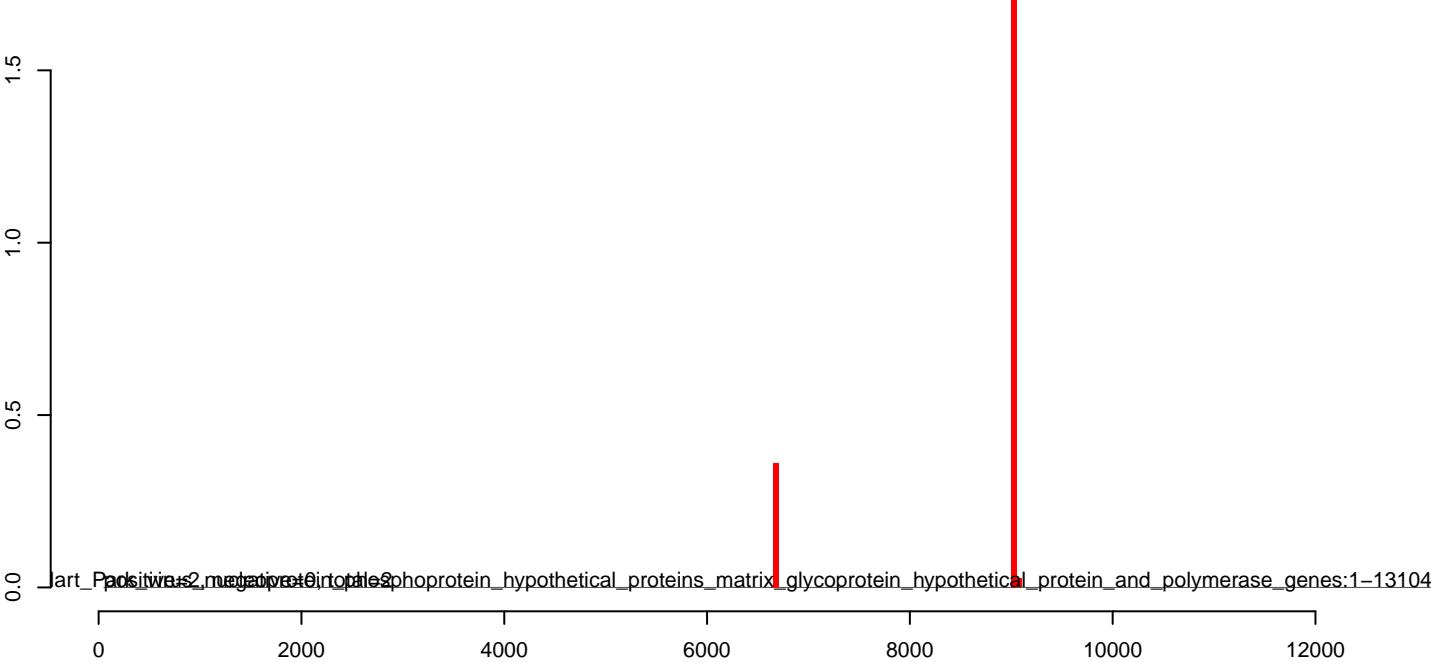
### AeAeg\_Fcanc\_GH.18\_23.rep



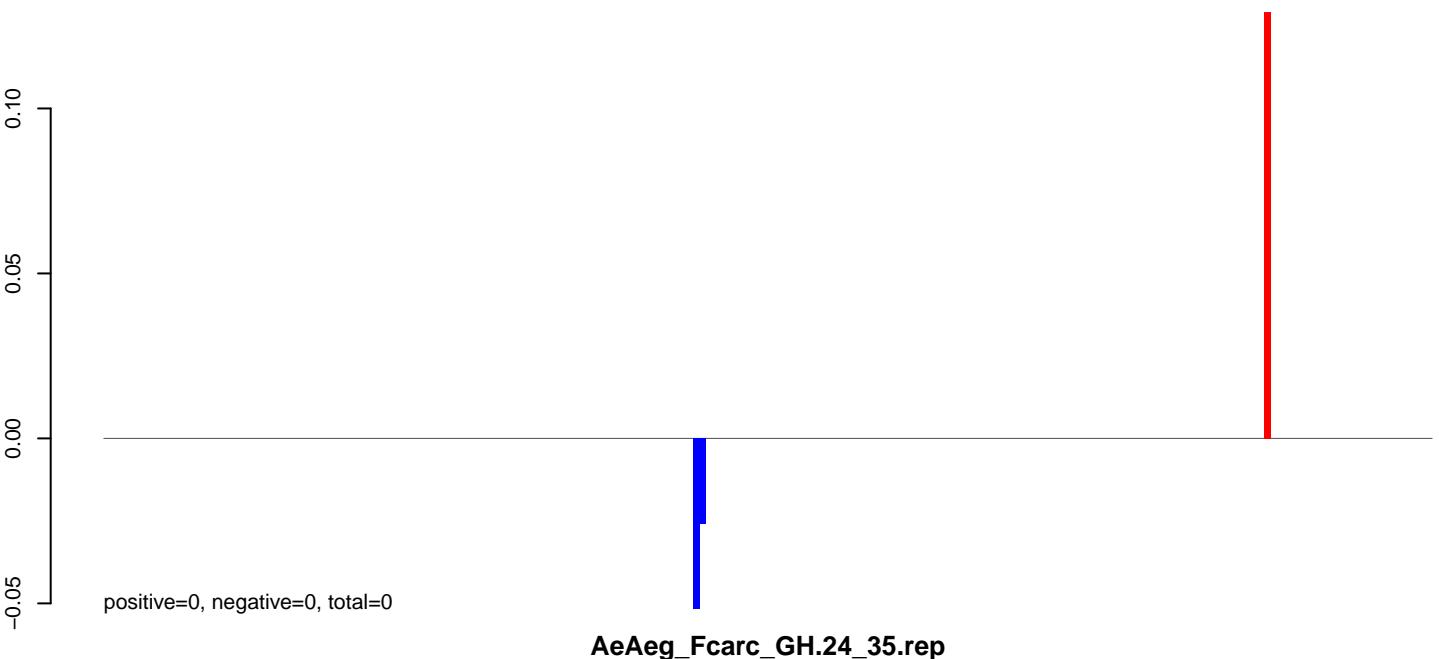
### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



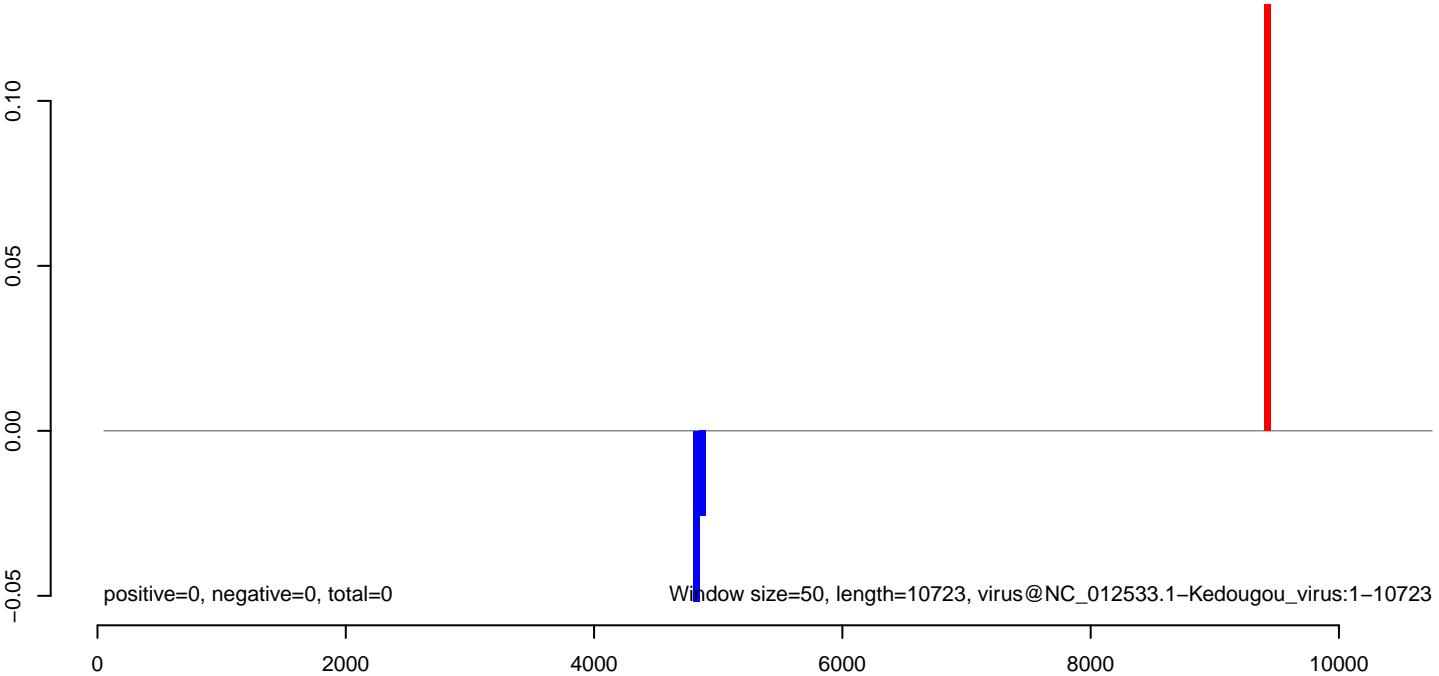
### AeAeg\_Fcanc\_GH.18\_23.rep



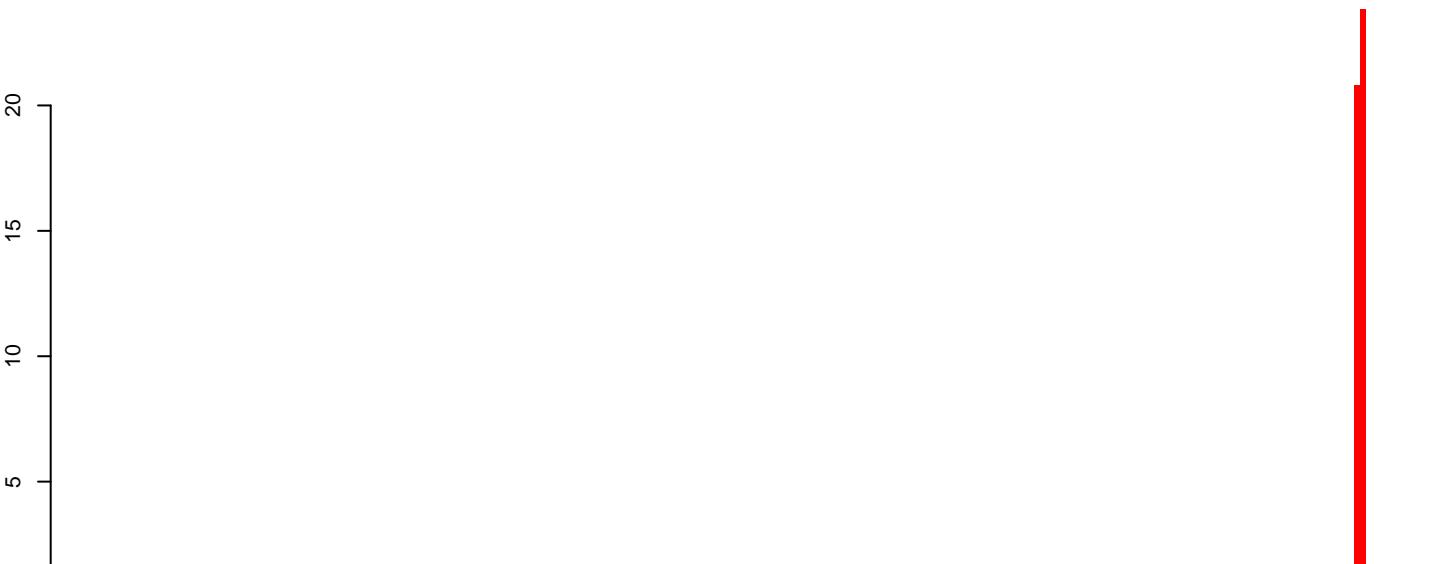
### AeAeg\_Fcanc\_GH.24\_35.rep



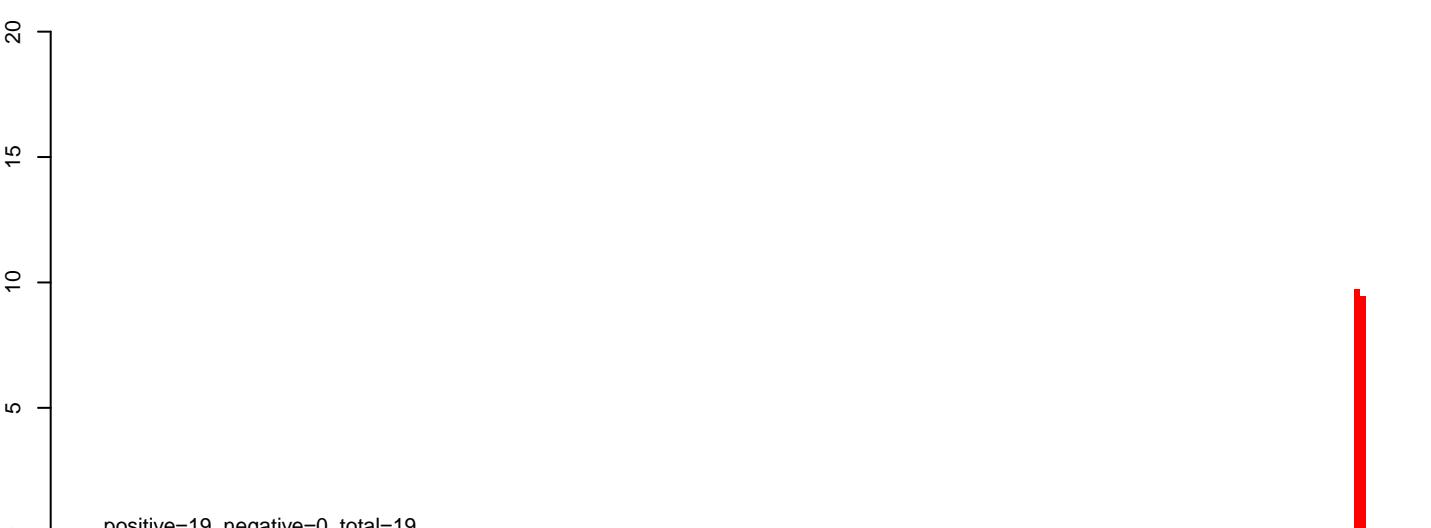
### AeAeg\_Fcanc\_GH.rep



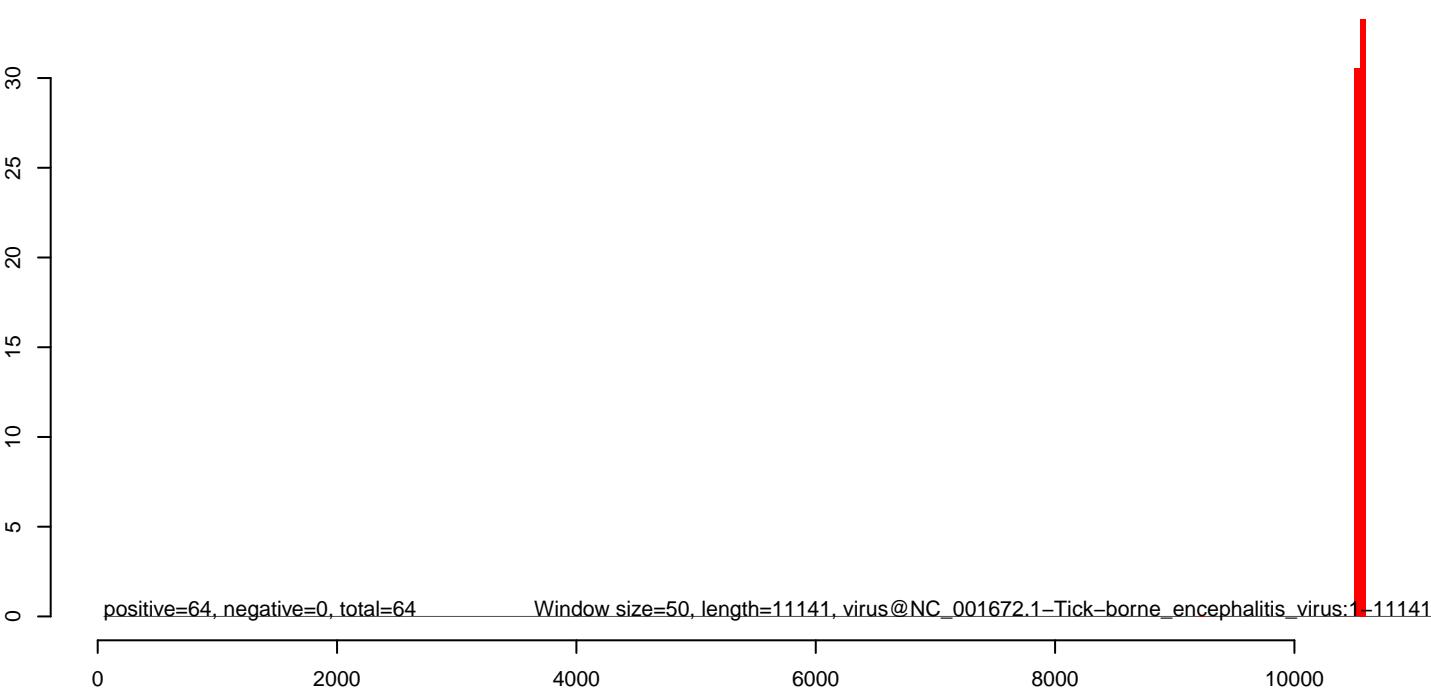
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep

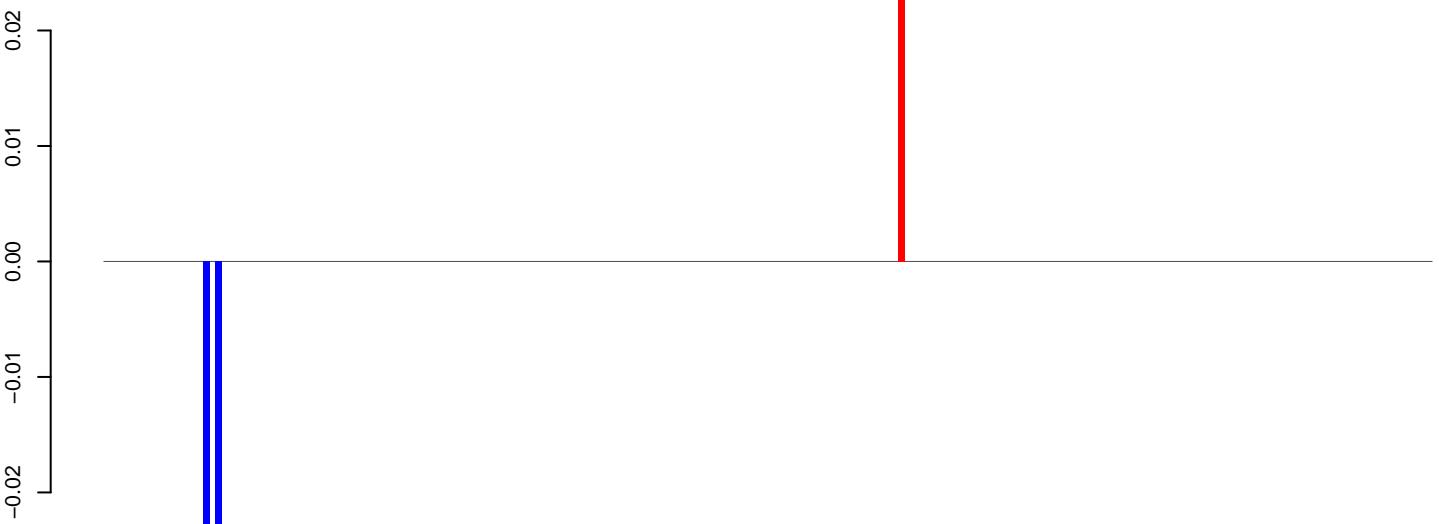


### AeAeg\_Fcanc\_GH.rep



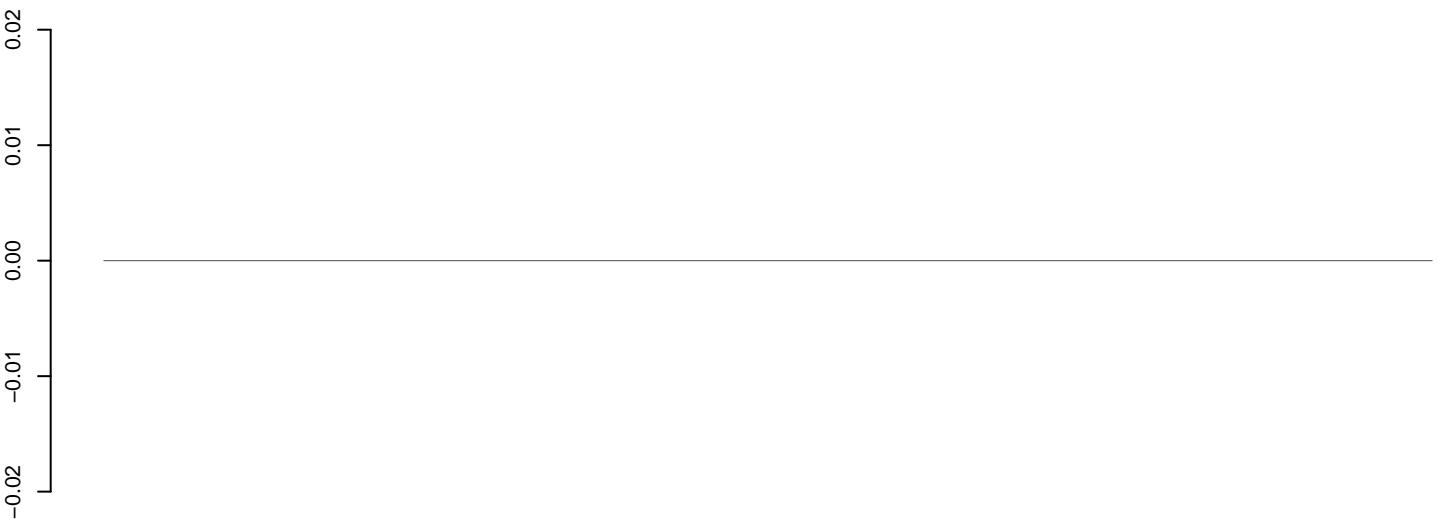
### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



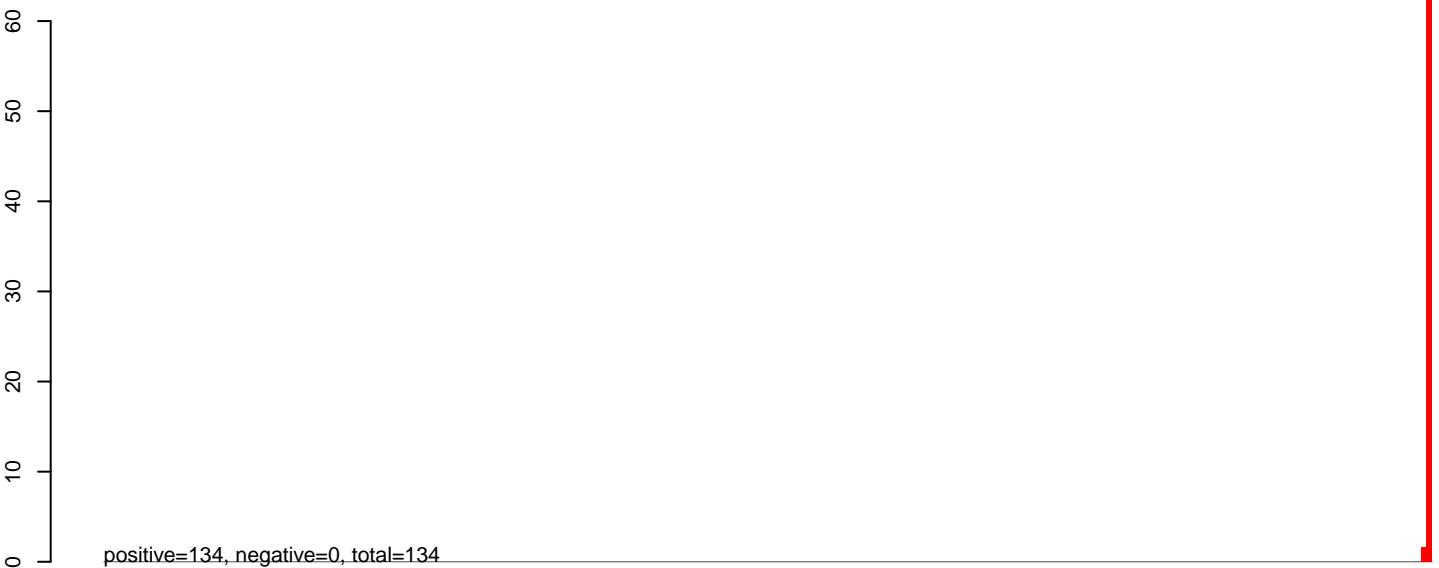
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10735, virus@NC\_001477.1-Dengue\_virus\_1:1-10735



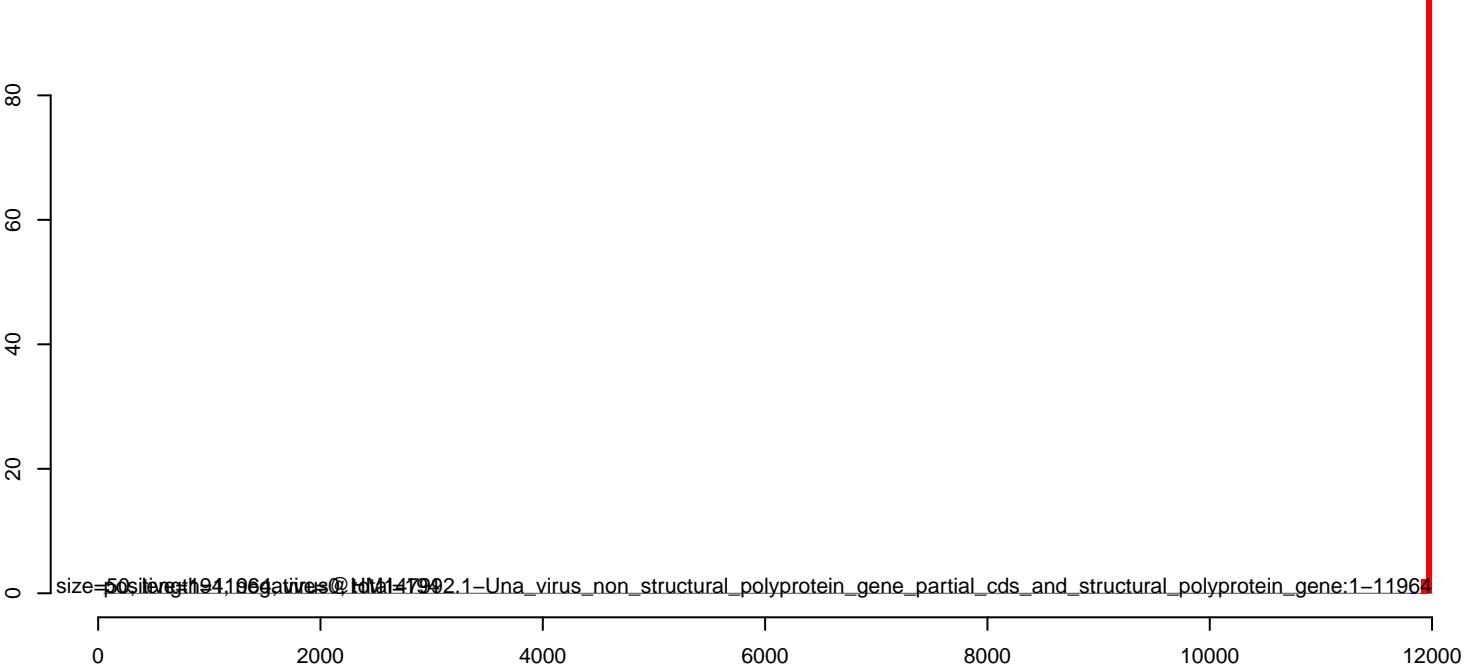
### AeAeg\_Fcanc\_GH.18\_23.rep



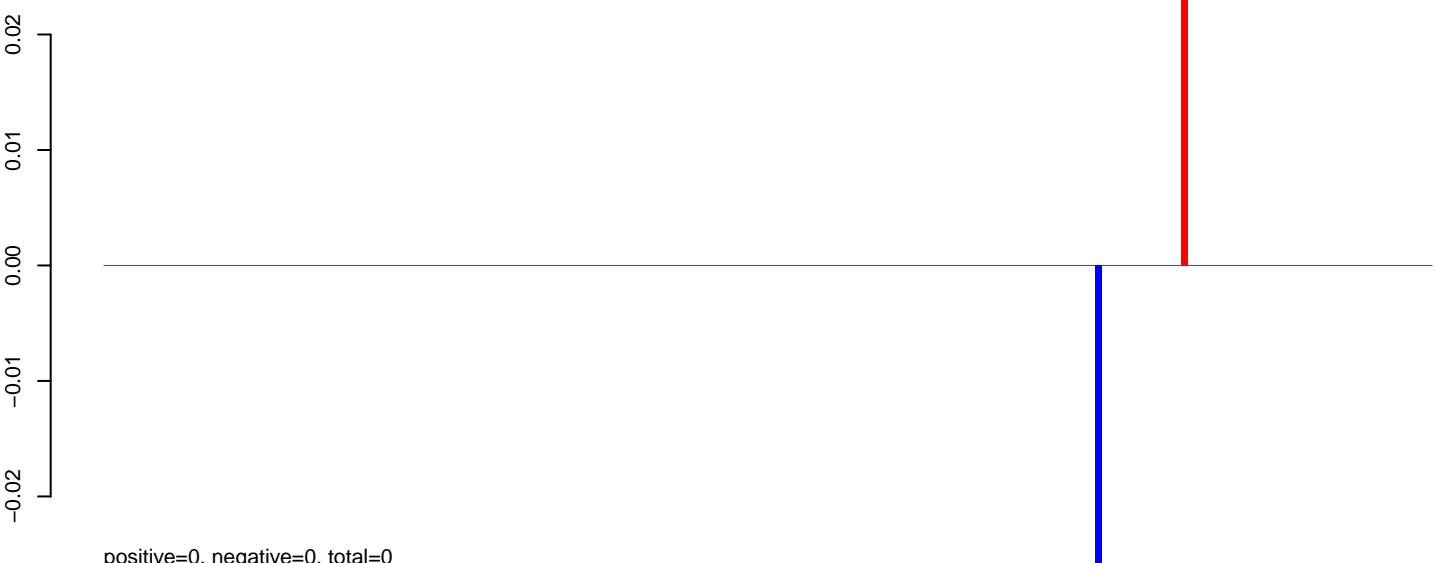
### AeAeg\_Fcanc\_GH.24\_35.rep



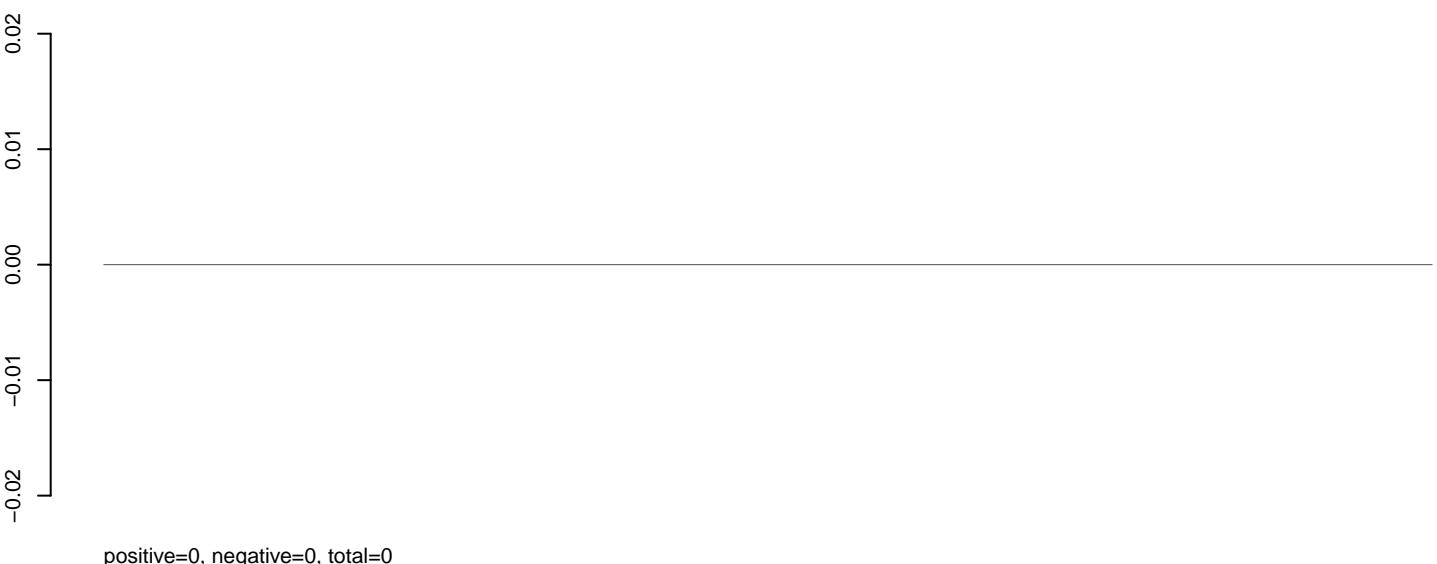
### AeAeg\_Fcanc\_GH.rep



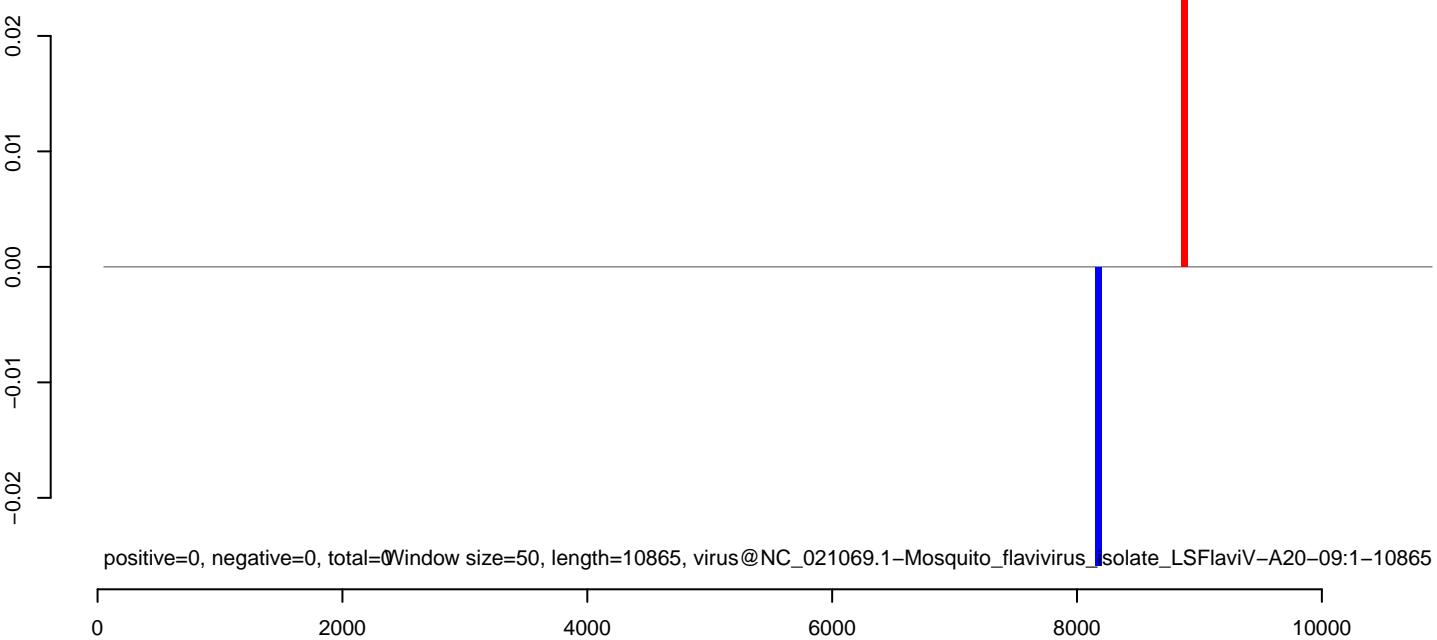
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep

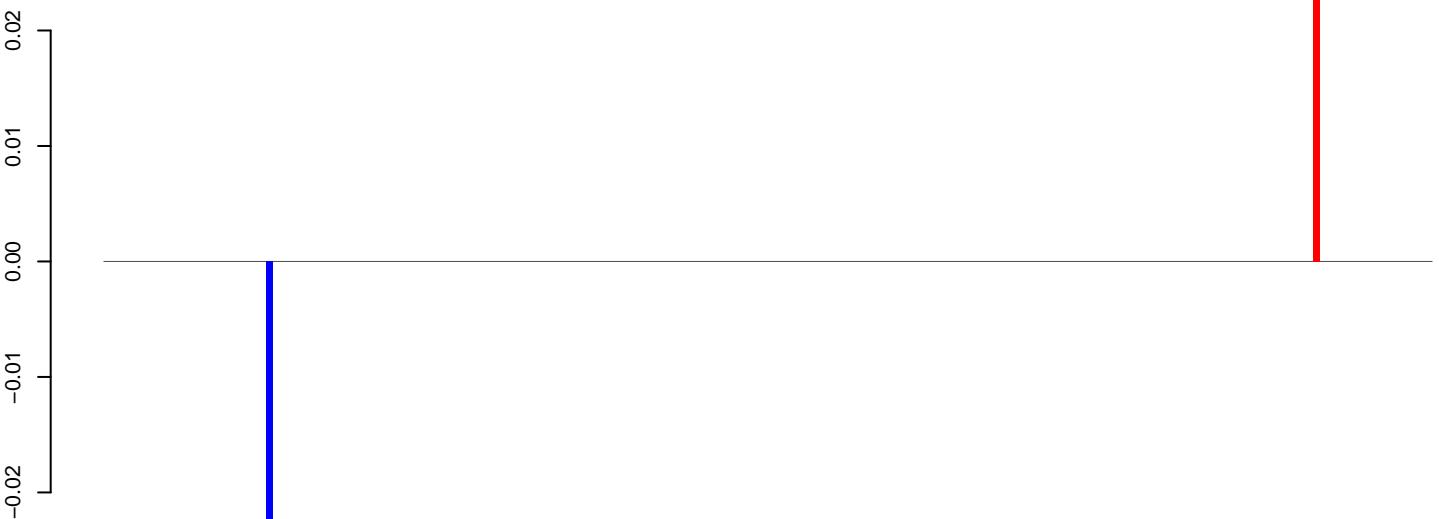


### AeAeg\_Fcanc\_GH.rep



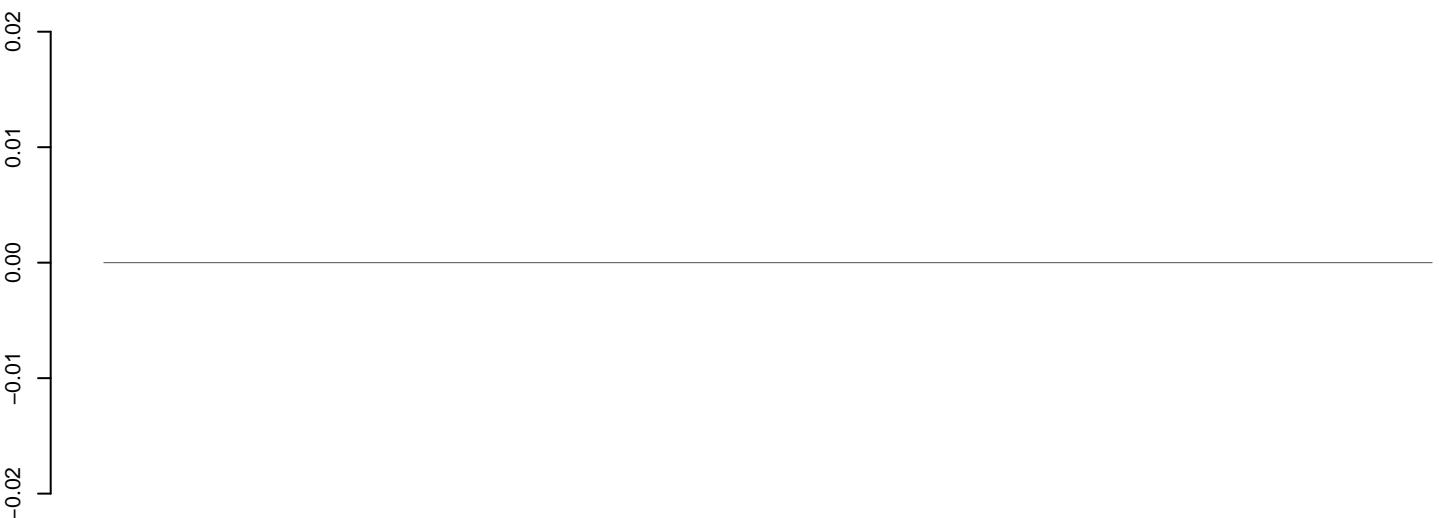
### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10648, virus@KM225264.1-Bainyik\_virus\_strain\_MK7979:1-10648



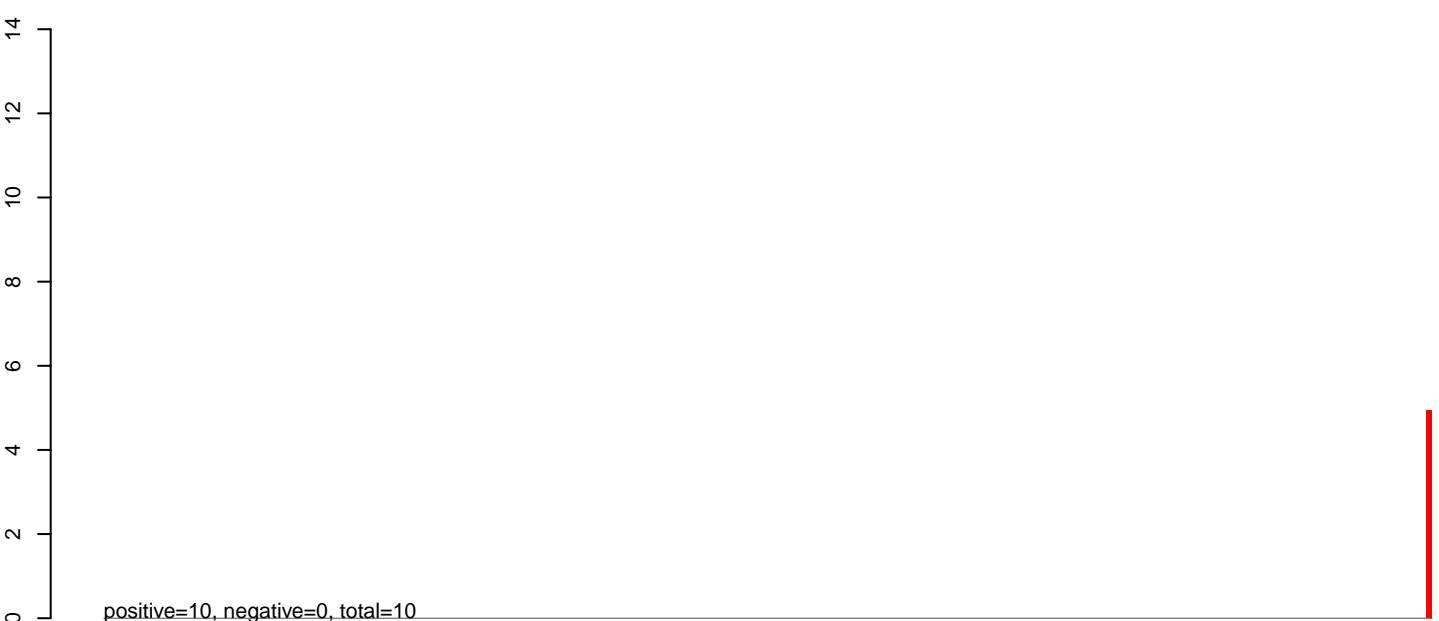
### AeAeg\_Fcanc\_GH.18\_23.rep

positive=28, negative=0, total=28



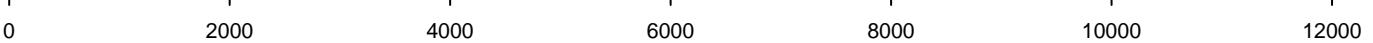
### AeAeg\_Fcanc\_GH.24\_35.rep

positive=10, negative=0, total=10

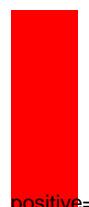


### AeAeg\_Fcanc\_GH.rep

positive=12052, negative=0, total=12052  
1-Trocara\_virus\_non\_structural\_polyprotein\_gene\_partial\_cds\_and\_structural\_polyprotein\_gene:1-12052



### AeAeg\_Fcarc\_GH.18\_23.rep

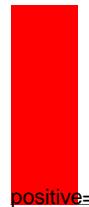


positive=0, negative=0, total=0

### AeAeg\_Fcarc\_GH.24\_35.rep

positive=0, negative=0, total=0

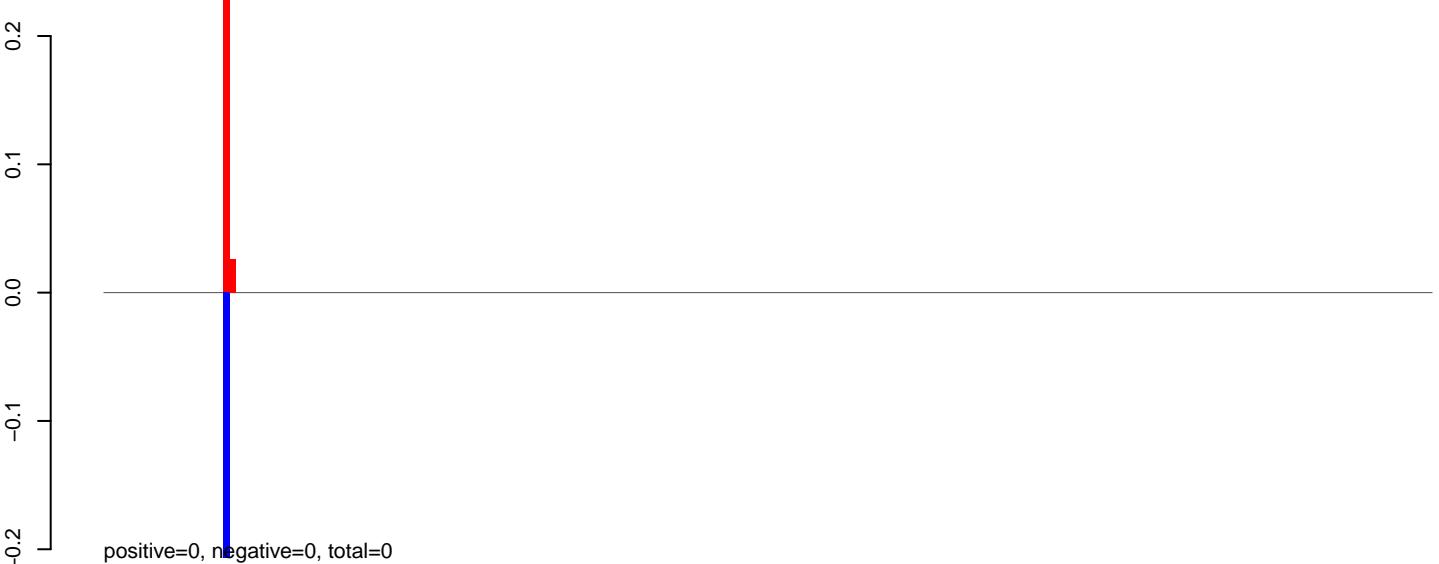
### AeAeg\_Fcarc\_GH.rep



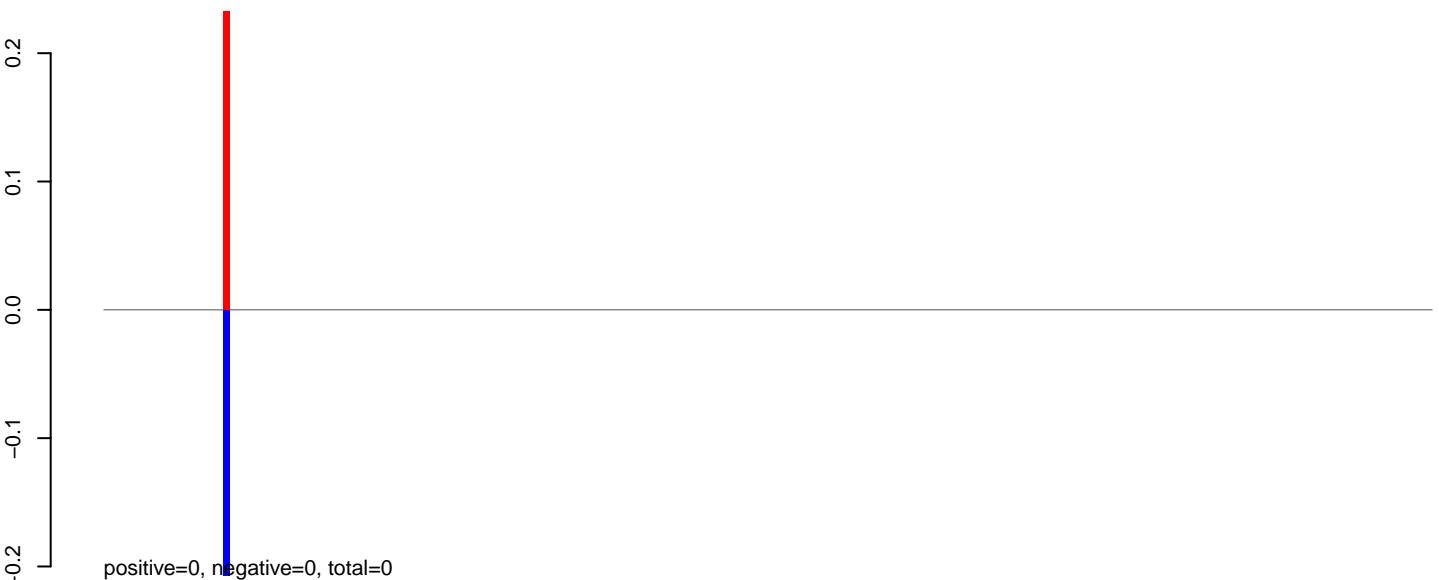
positive=0, negative=0, total=0, Length=1012, virus@AF411845.1-Elaeis\_guineensis\_MADS\_box\_transcription\_factor\_AGL2-3\_mRNA:1-1012

200 400 600 800 1000

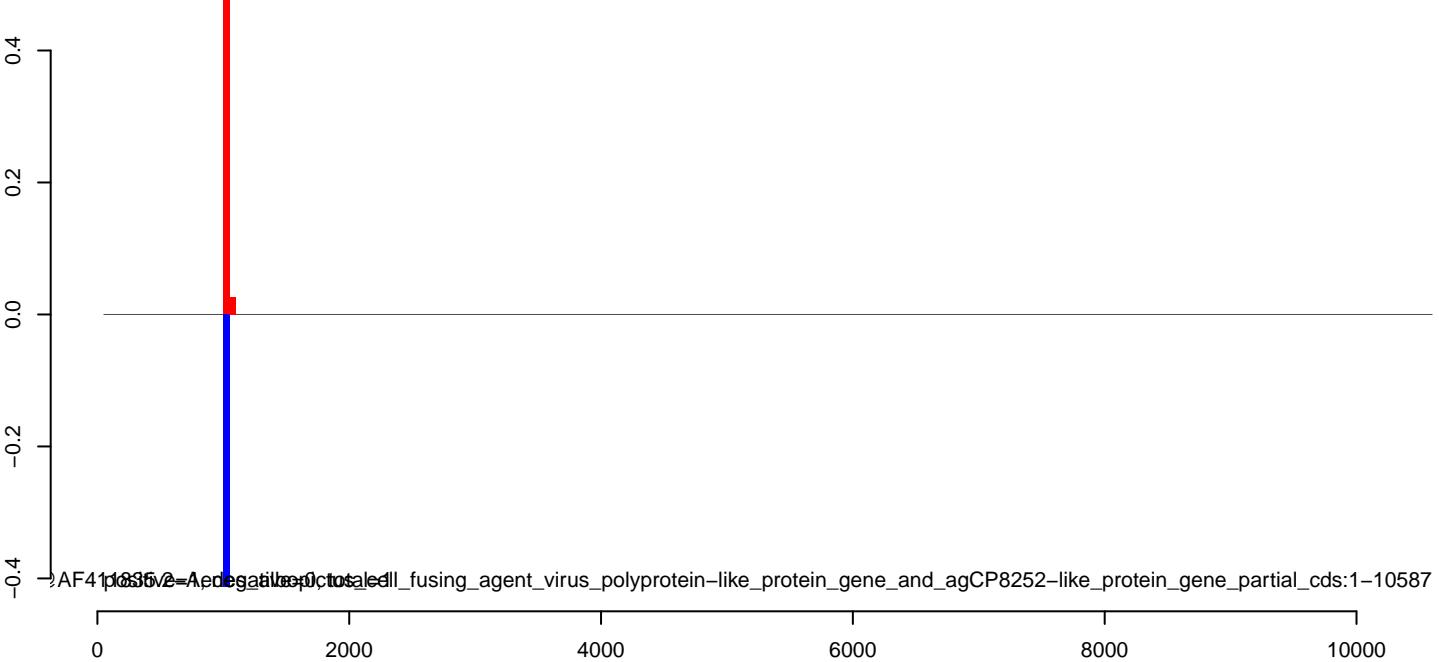
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcarc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcarc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcarc\_GH.rep

positive=0, negative=0, total=0

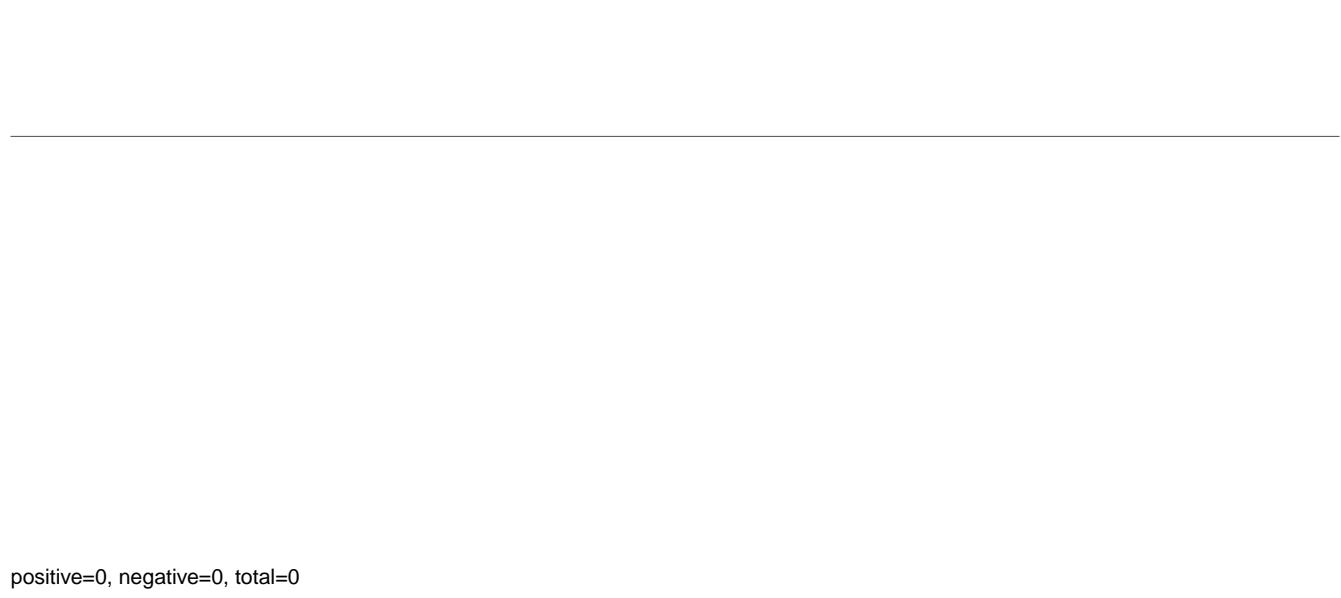
Window size=50, length=11494, virus@AF075258.1-Rio\_Negro\_virus\_strain\_AG80-663:1-11494

0 2000 4000 6000 8000 10000

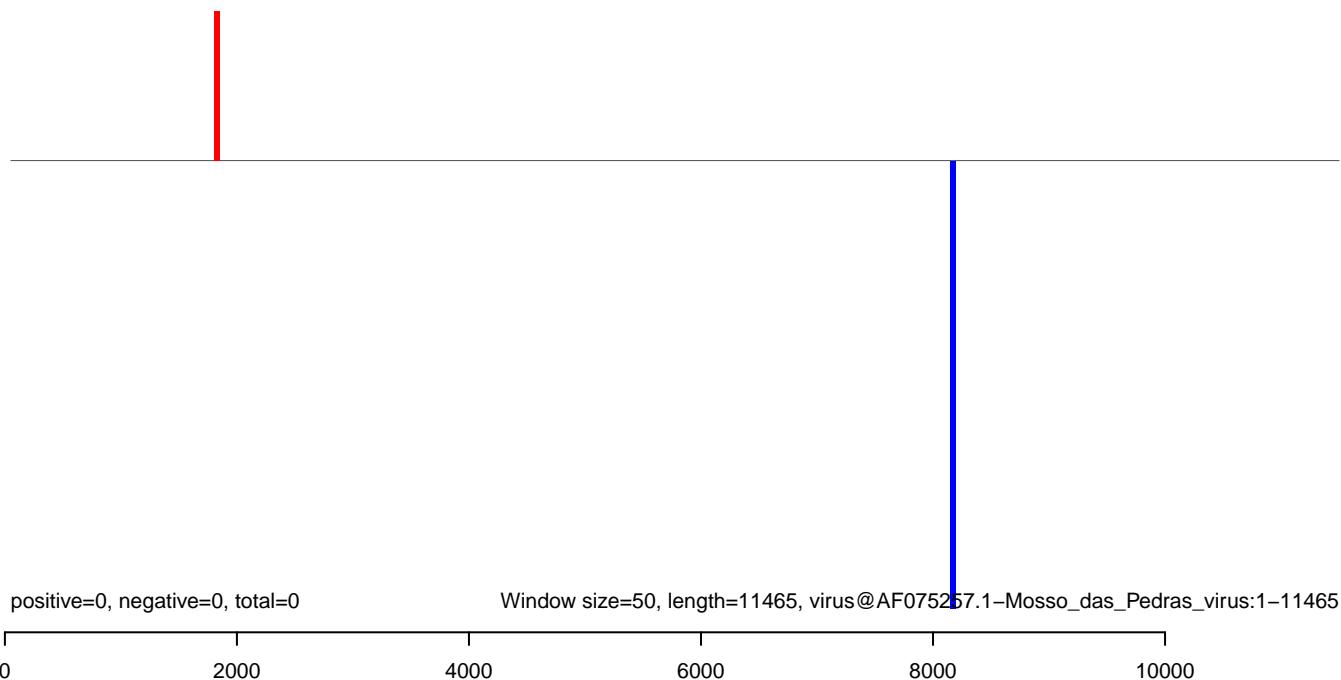
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

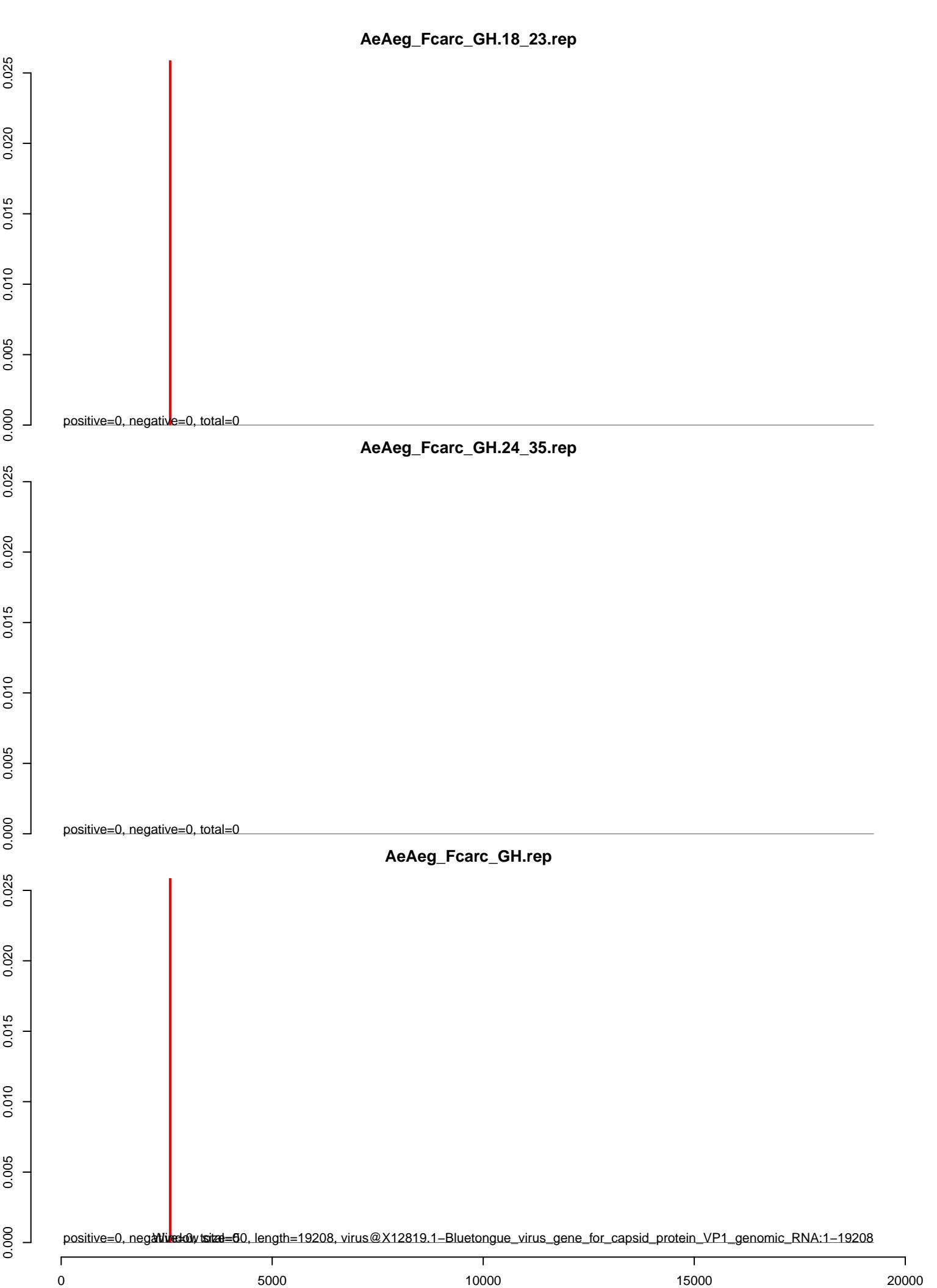
positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=60, length=19208, virus@X12819.1-Bluetongue\_virus\_gene\_for\_capsid\_protein\_VP1\_genomic\_RNA:1-19208



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

down 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

0 2000 4000 6000 8000 10000 12000 14000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

0 2000 4000 6000 8000 10000 12000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

-50, length=10968 negative=0 positive=0 NQ084639.1-Nkolbisson\_virus\_nucleoprotein\_phosphoprotein\_matrix\_glycoprotein\_and\_polymerase\_genes:1-10968

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

binjak positive=0, negative=0, total=0 phosphoprotein\_hypothetical\_protein\_matrix\_glycoprotein\_hypothetical\_proteins\_and\_polymerase\_genes:1-13155



Manipulusis0\_methyltransferase\_phoprotein\_hypothetical\_proteins\_matrix\_glycoprotein\_hypothetical\_protein\_and\_polymerase\_genes:1-13784

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

-Kam positive=0, negative=0, total=0 phosphoprotein\_hypothetical\_proteins\_matrix\_glycoprotein\_hypothetical\_protein\_and\_polymerase\_genes:1-13209

0 2000 4000 6000 8000 10000 12000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

rus@nos:026449:3 relativeData total nos\_nucleoprotein\_phosphoprotein\_matrix\_glycoprotein\_hypothetical\_protein\_and\_polymerase\_genes:1-11450

0 2000 4000 6000 8000 10000

### AeAeg\_Fcarc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcarc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcarc\_GH.rep

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

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

v size=50 itime=0.1 neg=28 virus=@NC\_0033327.1-Hubei\_chuvirus-like\_virus\_1\_strain\_QTM26249\_RNA-dependent\_RNA\_polymerase\_gene:1-6873

0 1000 2000 3000 4000 5000 6000 7000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

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



0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

incomplete positive=0 negative=0 total=0 X01-6\_ORF1\_ORF1\_ORF2\_ORF2\_glycoprotein\_G\_and\_RNA-dependent\_RNA\_polymerase\_L\_genes:1-12774

0 2000 4000 6000 8000 10000 12000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

Win positive=0 negative=0 total=0 @ NC\_030115.1-Anopheles\_C\_virus\_non-structural\_polyprotein\_and\_structural\_polyprotein\_genes:1-8919

0 2000 4000 6000 8000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

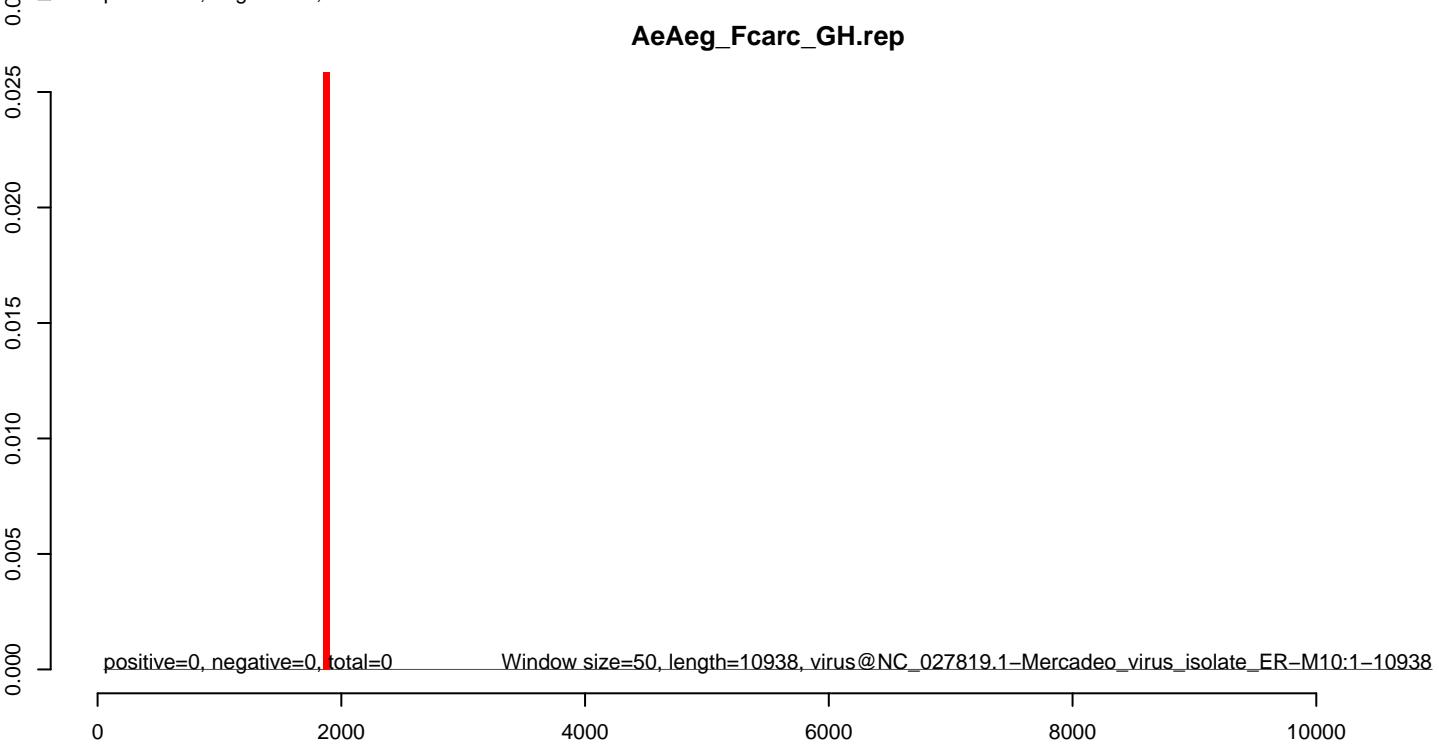
positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10938, virus@NC\_027819.1-Mercadeo\_virus\_isolate\_ER-M10:1-10938



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.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

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

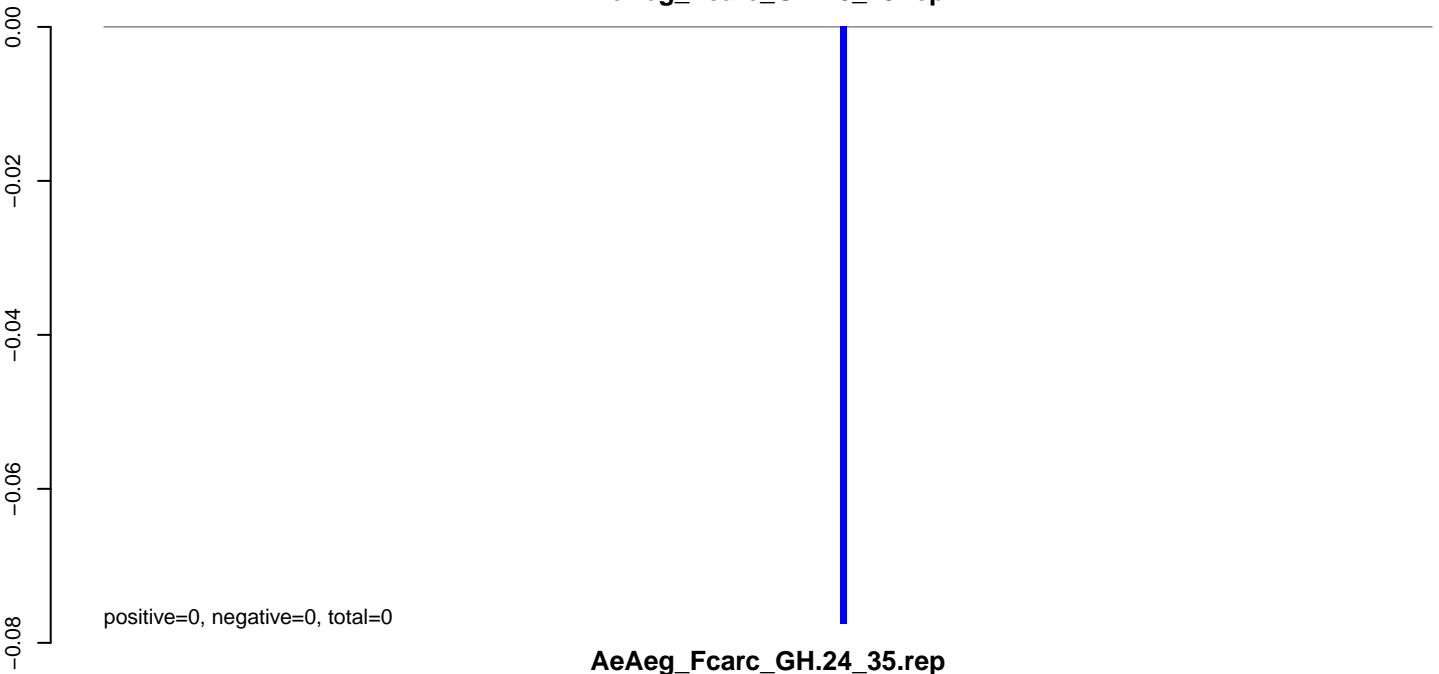


### AeAeg\_Fcanc\_GH.rep

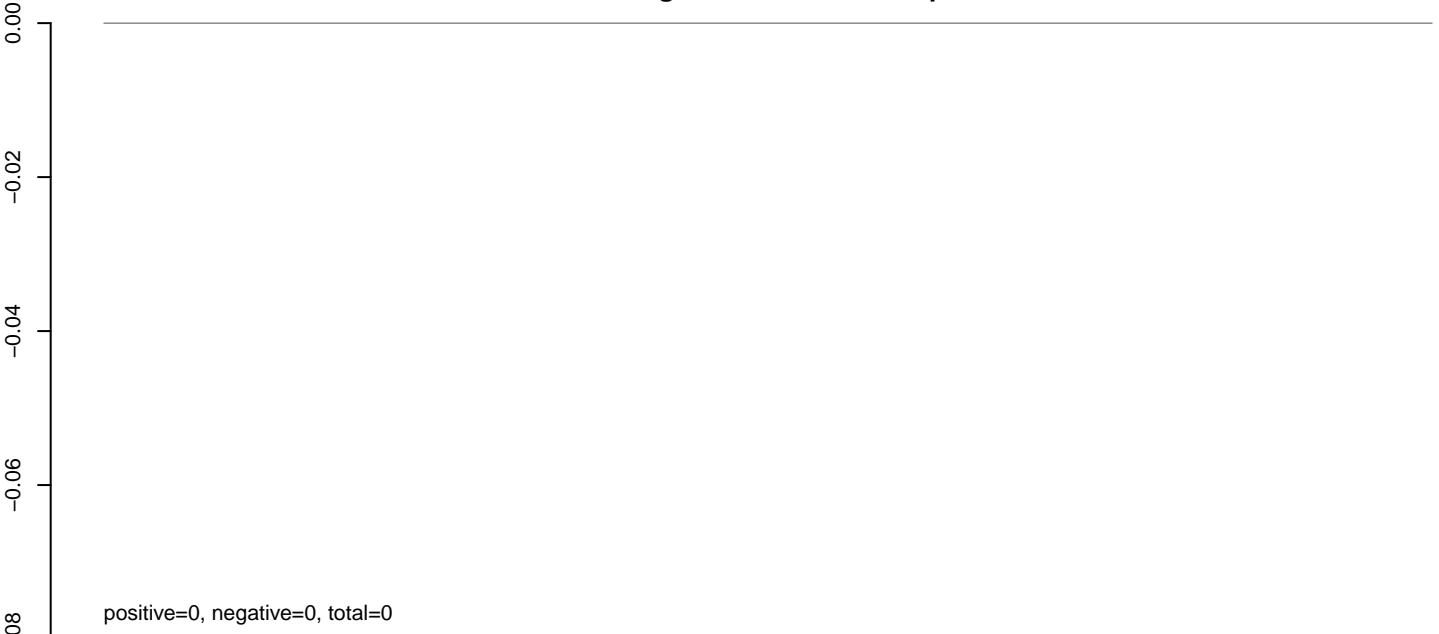
positive=0, negative=0, total=0



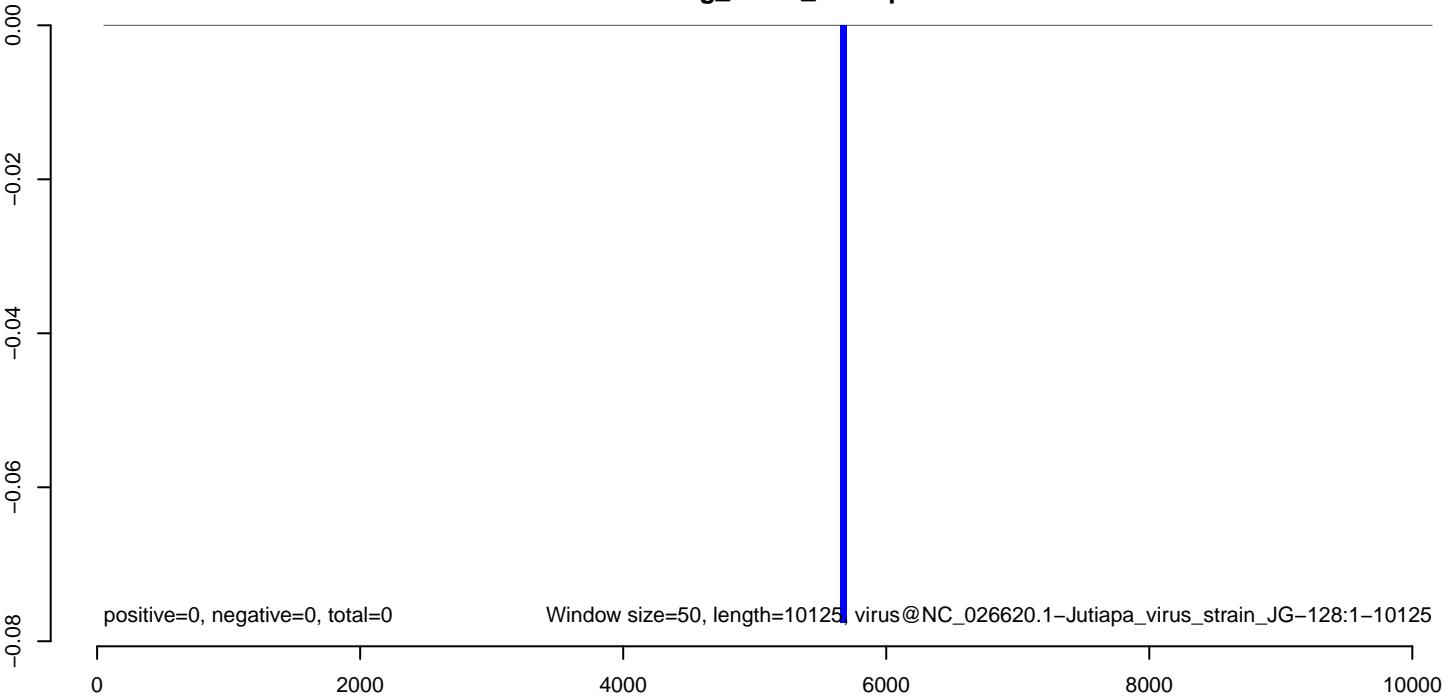
### AeAeg\_Fcanc\_GH.18\_23.rep



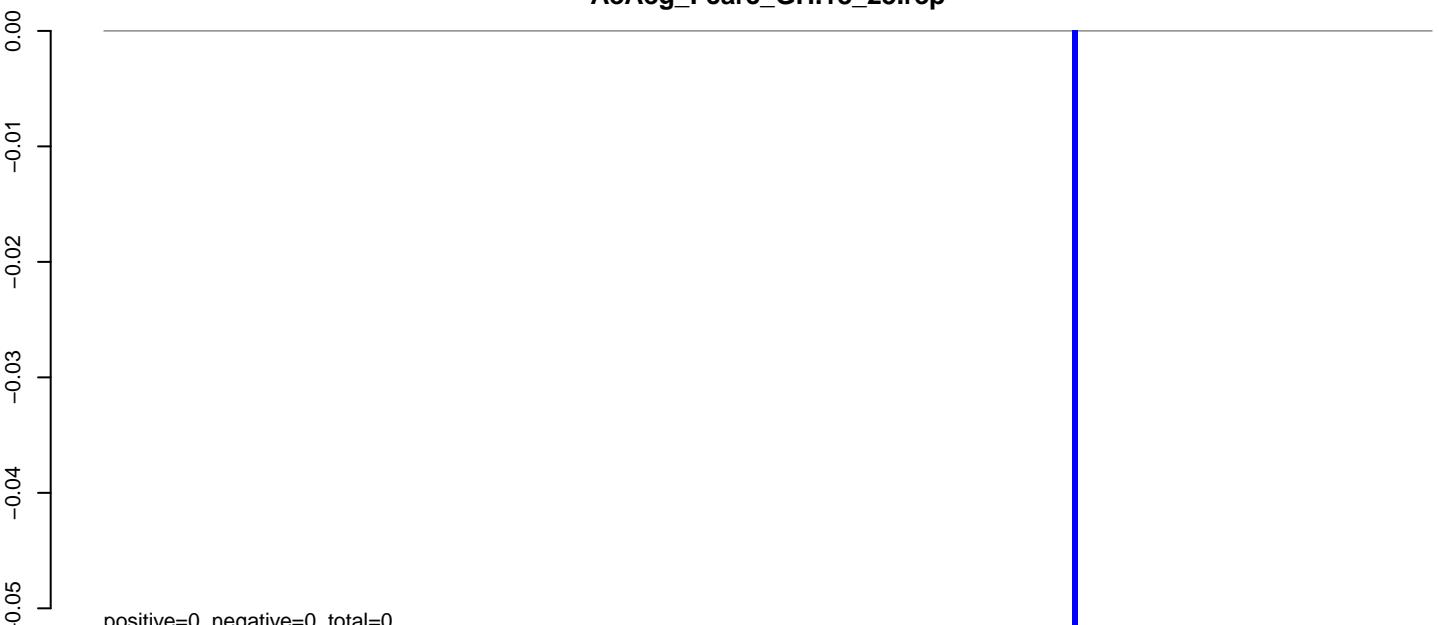
### AeAeg\_Fcanc\_GH.24\_35.rep



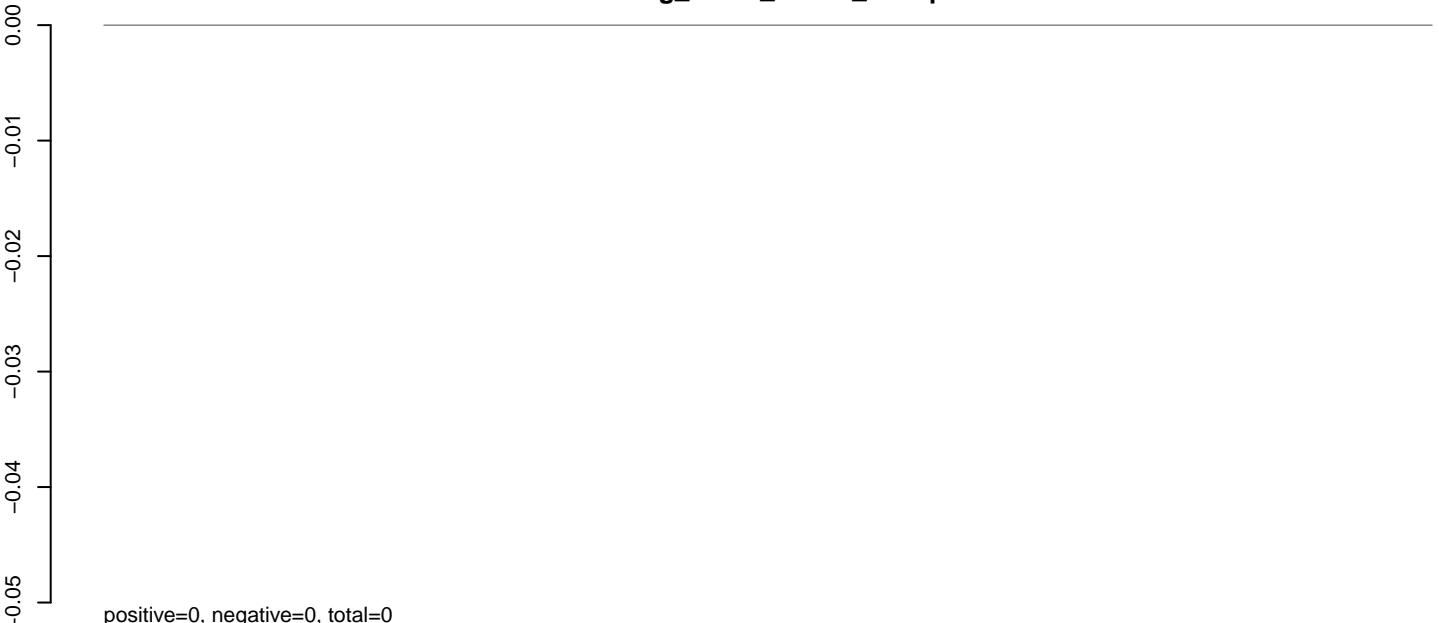
### AeAeg\_Fcanc\_GH.rep



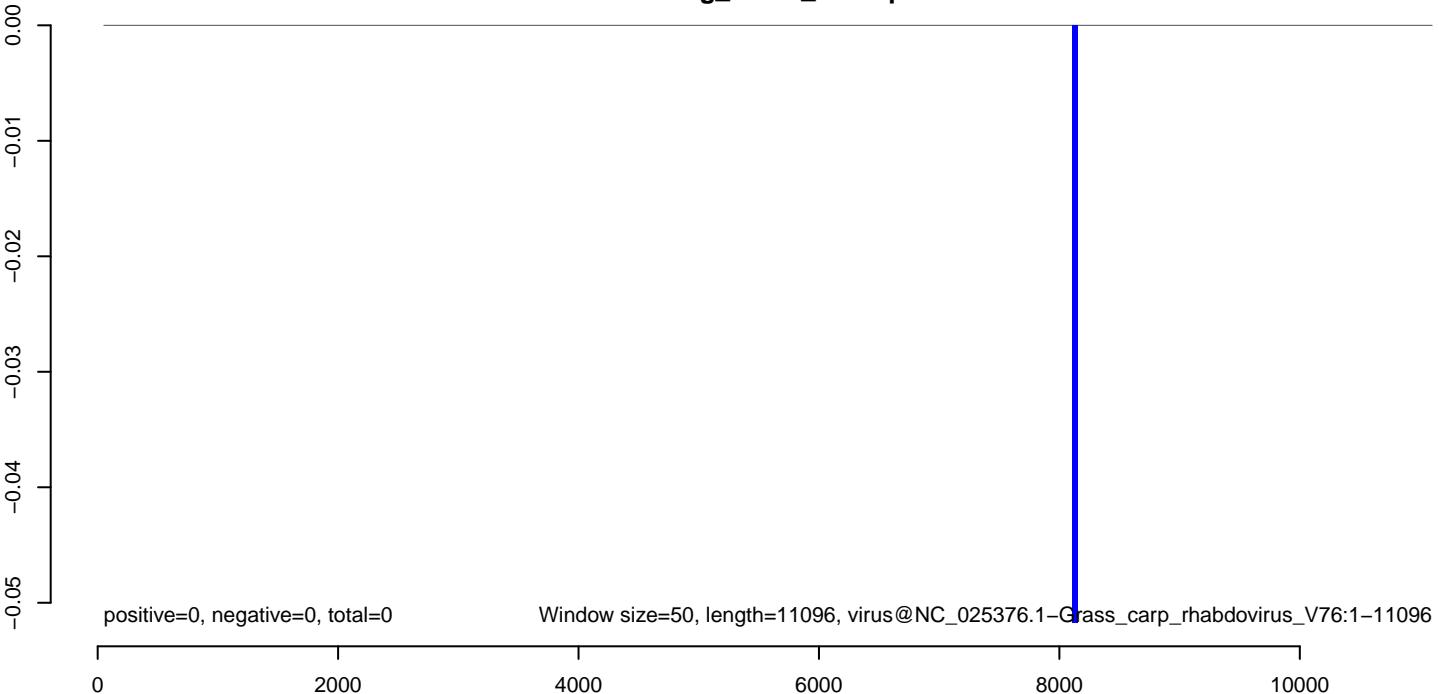
### AeAeg\_Fcarc\_GH.18\_23.rep



### AeAeg\_Fcarc\_GH.24\_35.rep



### AeAeg\_Fcarc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0      Window size=50, length=11019, virus@NC\_025364.1-Malpais\_Spring\_virus\_strain\_85-488NM:1-11019

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0, window size=50, length=12240, virus@NC\_025362.1-Xiburema\_virus\_isolate\_XIBV\_BE\_AR\_362159:1-12240

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=11550, virus@NC\_024887.1-Middleburg\_virus\_isolate\_ArB-8422:1-11550

0 2000 4000 6000 8000 10000 12000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10878, virus@NC\_024299.2-Nienokoue\_virus\_isolate\_B51\_CI\_2004:1-10878



0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

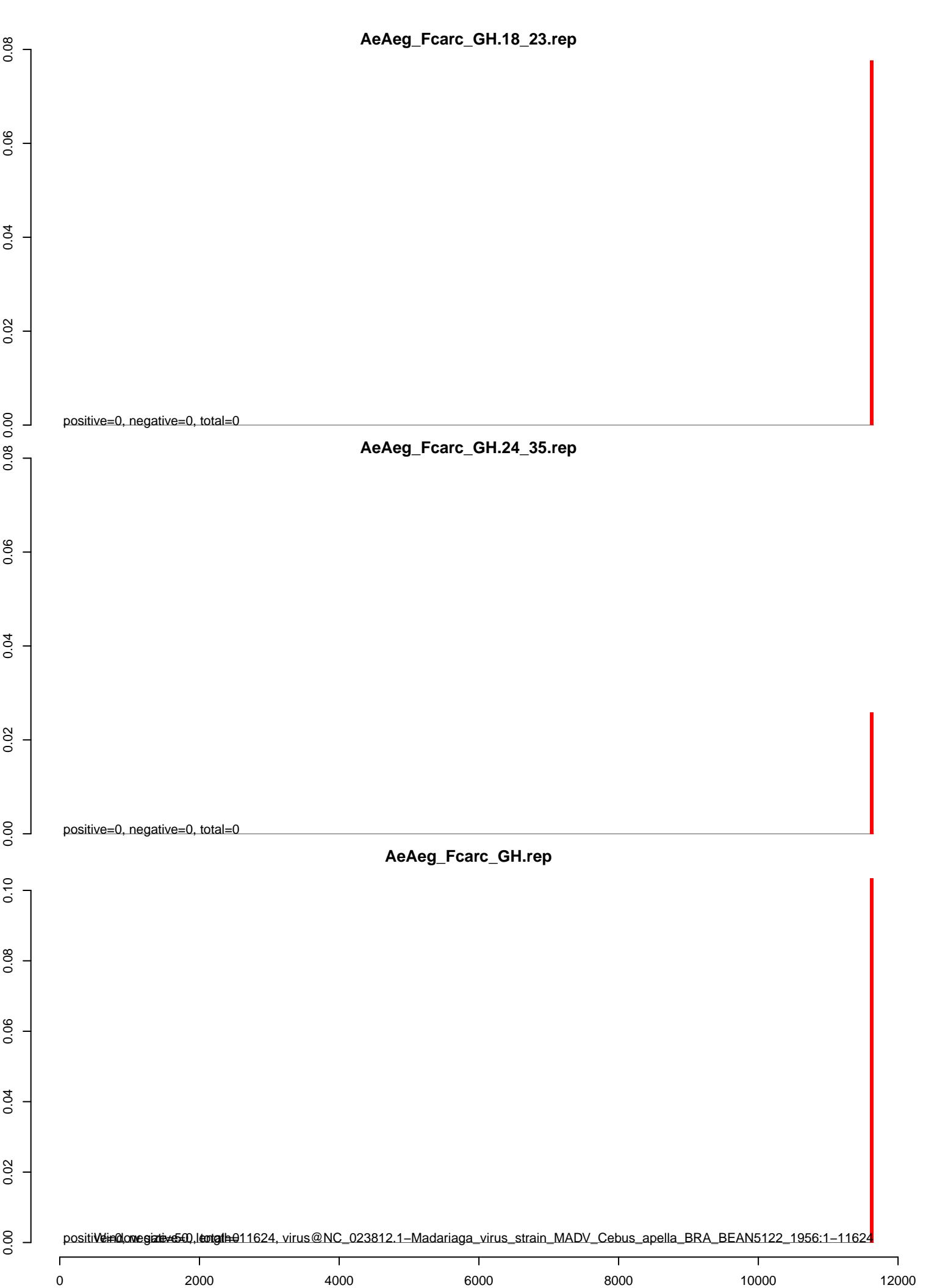
positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=500, total=11624, virus@NC\_023812.1-Madariaga\_virus\_strain\_MADV\_Cebus\_apella\_BRA\_BEAN5122\_1956:1-11624



### AeAeg\_Fcarc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcarc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcarc\_GH.rep

positive=0, negative=0, total=0, window size=50, length=10673, virus@NC\_023424.1-Tyuleny\_virus\_strain\_LEIV-6C\_polyprotein\_gene1-10673

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=13298, virus@NC\_020804.1-Tibrogargan\_virus\_strain\_CS132:1-13298

0 2000 4000 6000 8000 10000 12000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=11634, virus@NC\_018615.1-Eilat\_virus\_isolate\_EO329:1-11634



0 2000 4000 6000 8000 10000 12000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10791, virus@NC\_016997.1-Donggang\_virus:1-10791

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=6, negative=0, total=6

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=3, negative=0, total=3

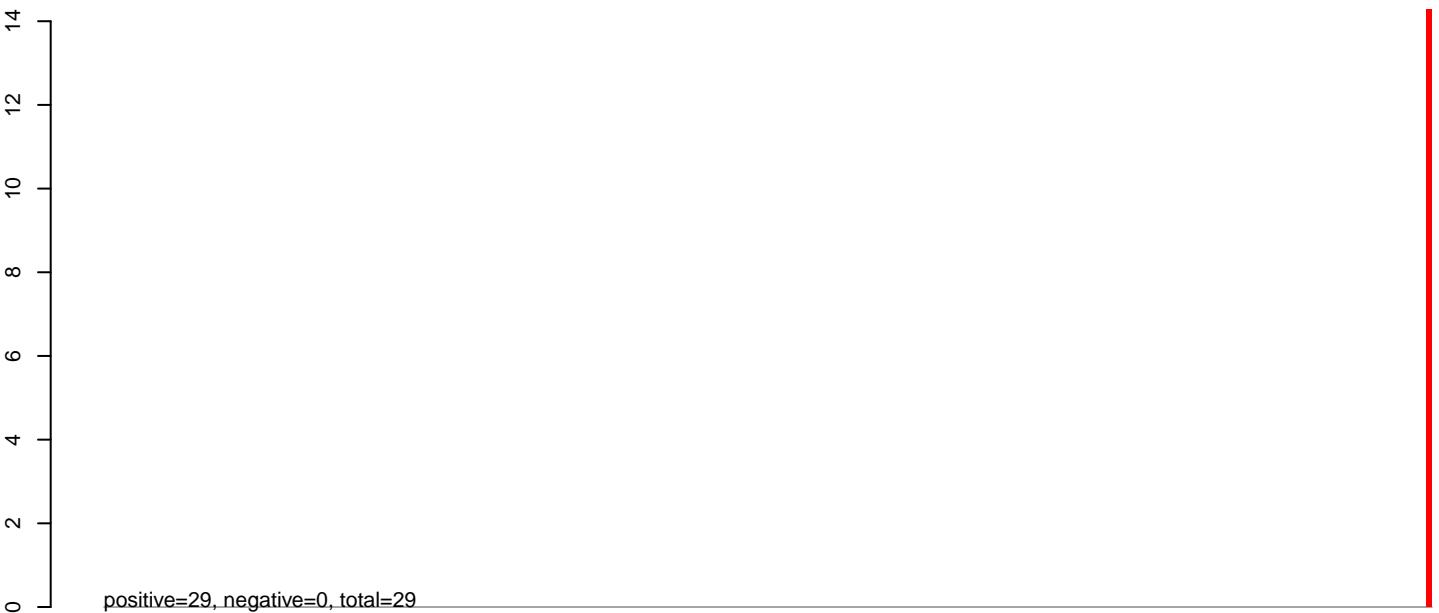
### AeAeg\_Fcanc\_GH.rep

positive=9, negative=0, total=9

Window size=50, length=11877, virus@NC\_016962.1-Bebaru\_virus:1-11877

0 2000 4000 6000 8000 10000 12000

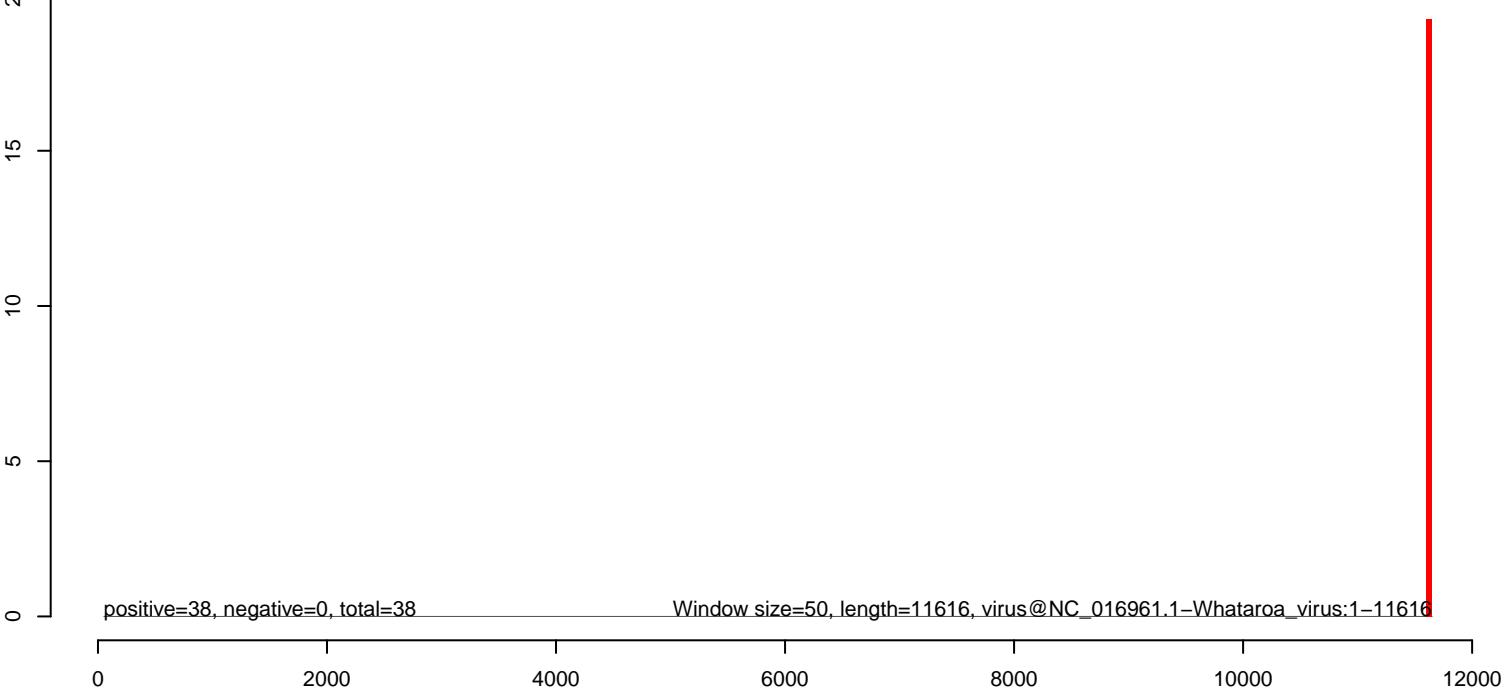
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=10990, virus@NC\_015843.2-Tembusu\_virus\_flavivirus\_polyprotein\_flavivirus\_polyprotein\_gene\_gene:1-10990



0 2000 4000 6000 8000 10000

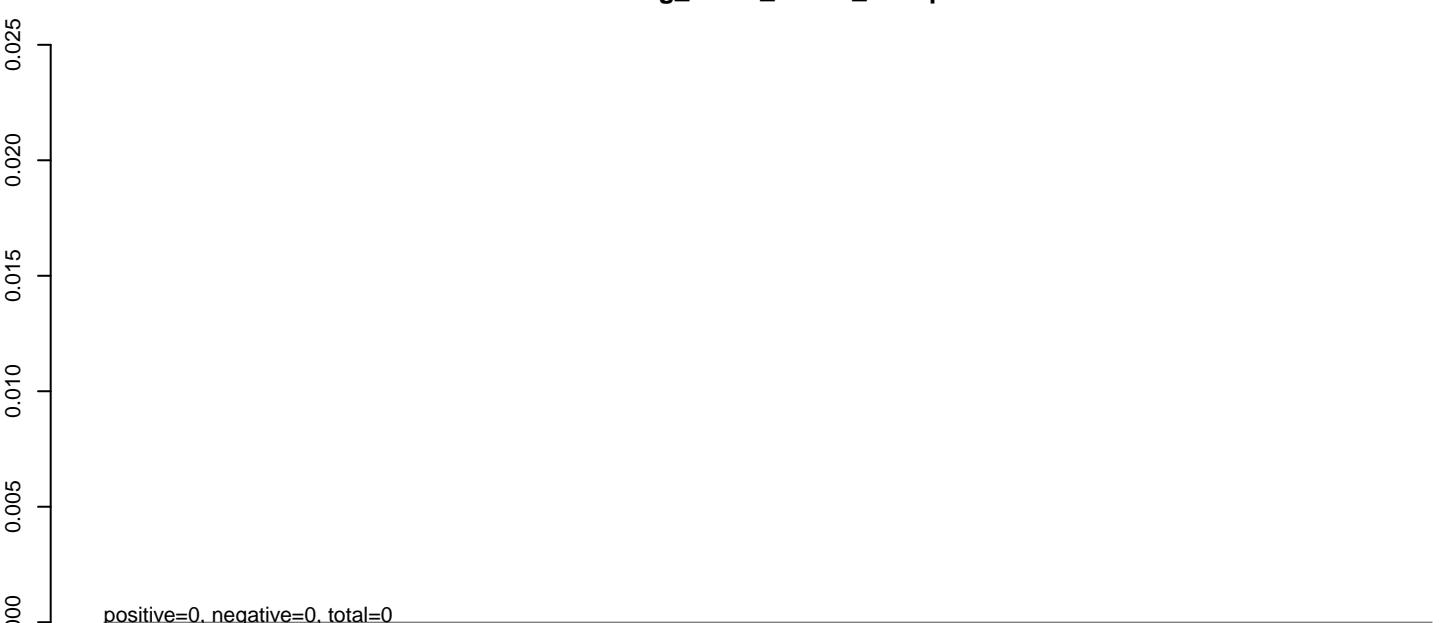
### AeAeg\_Fcarc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcarc\_GH.24\_35.rep

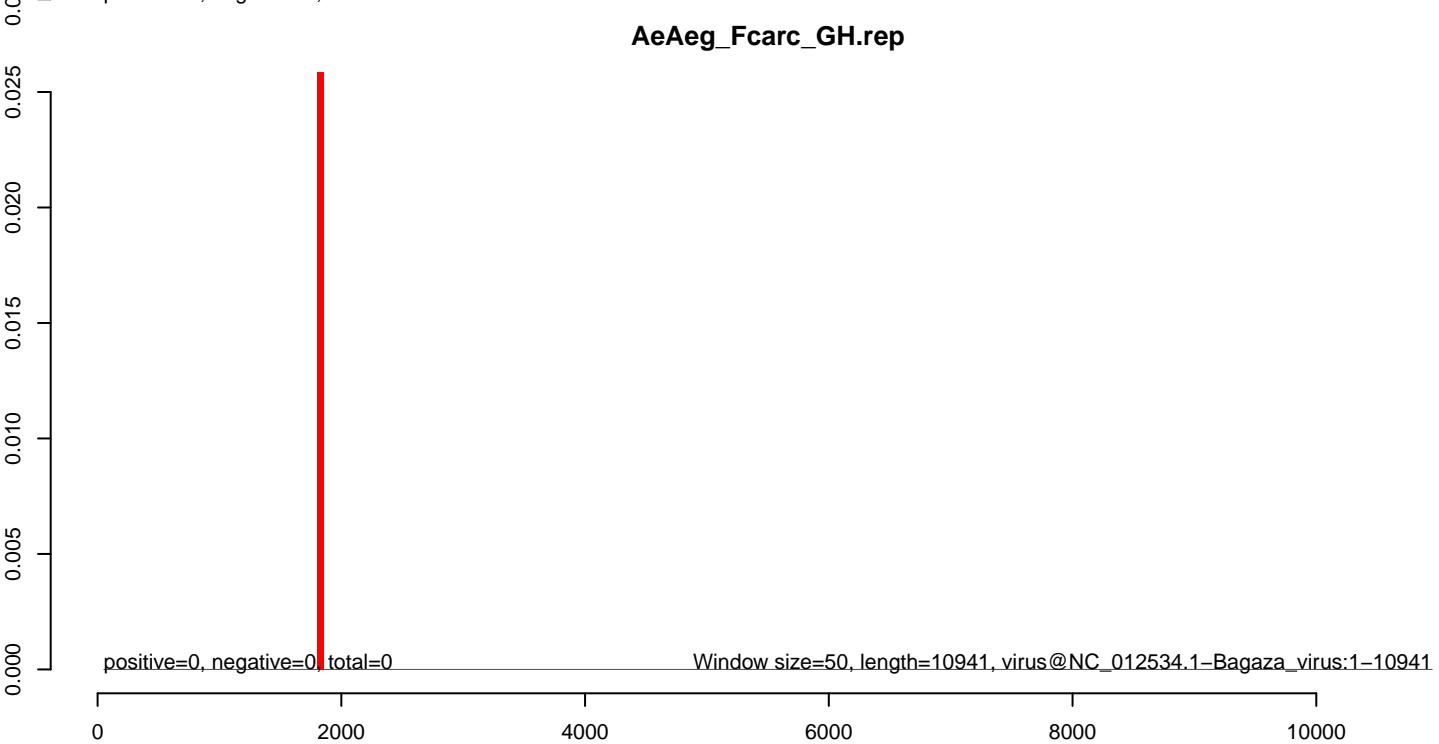
positive=0, negative=0, total=0



### AeAeg\_Fcarc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10941, virus@NC\_012534.1-Bagaza\_virus:1-10941



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=11029, virus@NC\_009942.1-West\_Nile\_virus\_lineage\_1:1-11029



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10874, virus@NC\_009029.2-Kokobera\_virus:1-10874



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

positive=0, negative=0, total=0

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

Window size=50, length=10815, virus@NC\_009026.2-Bussuquara\_virus:1-10815

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

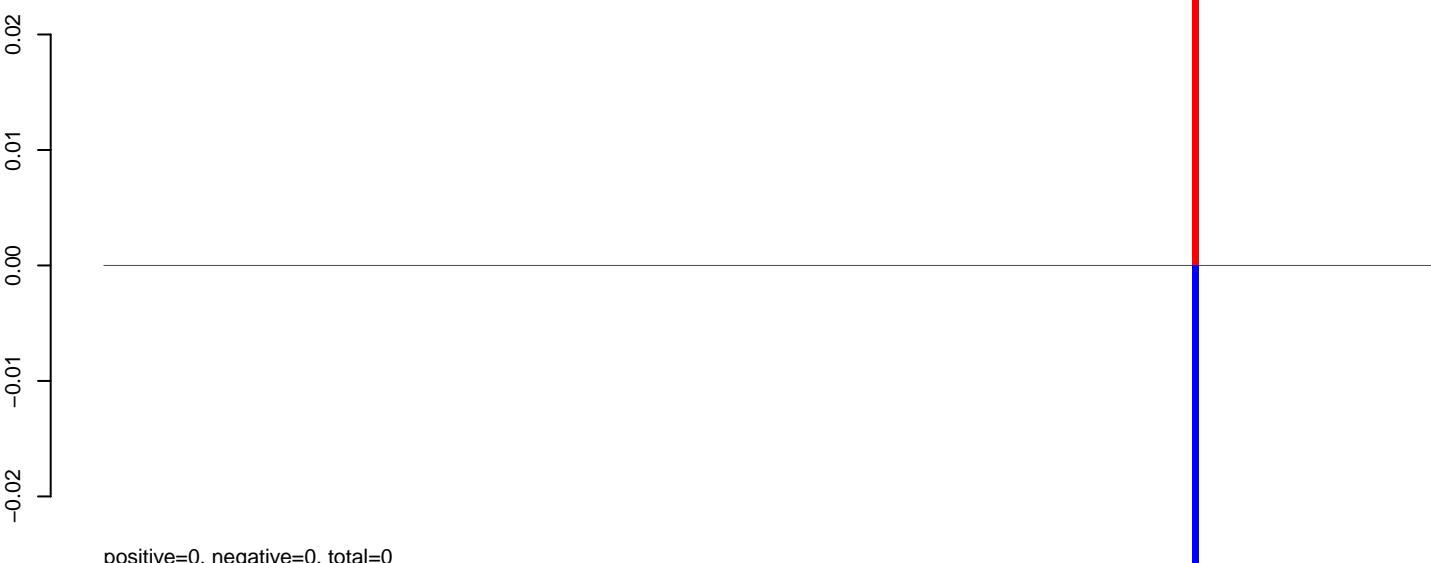
positive=0, negative=0, total=0

Window size=50, length=10510, virus@NC\_008718.1-Entebbe\_bat\_virus:1-10510

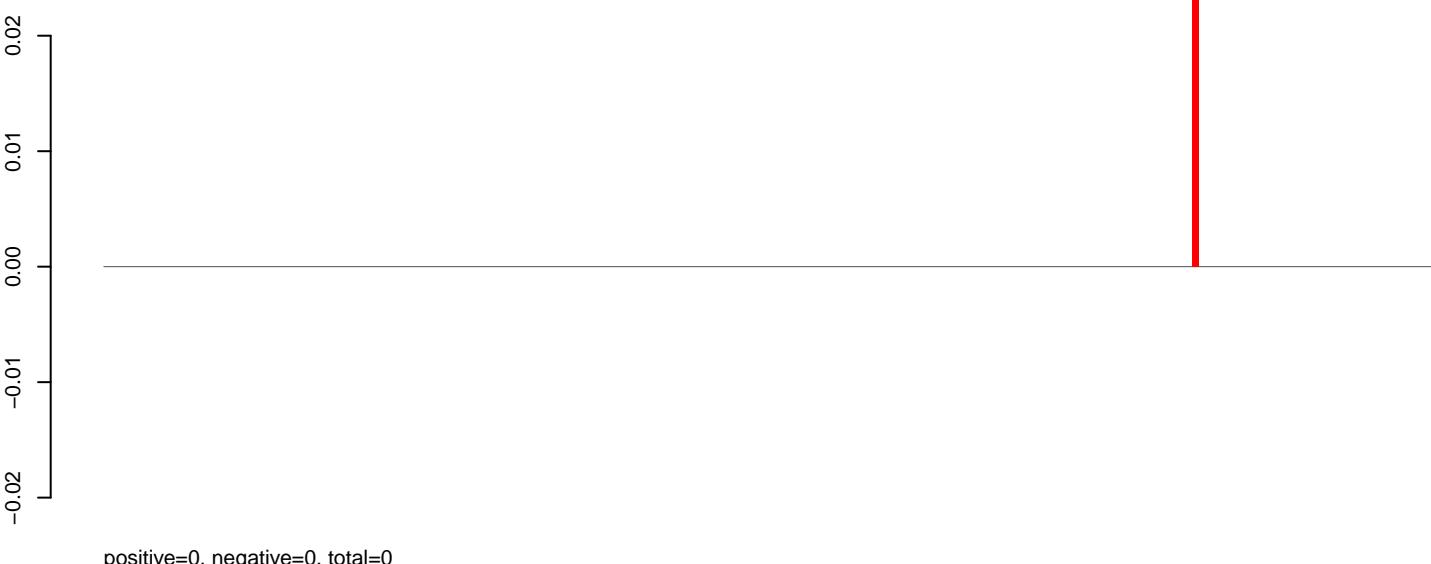


0 2000 4000 6000 8000 10000

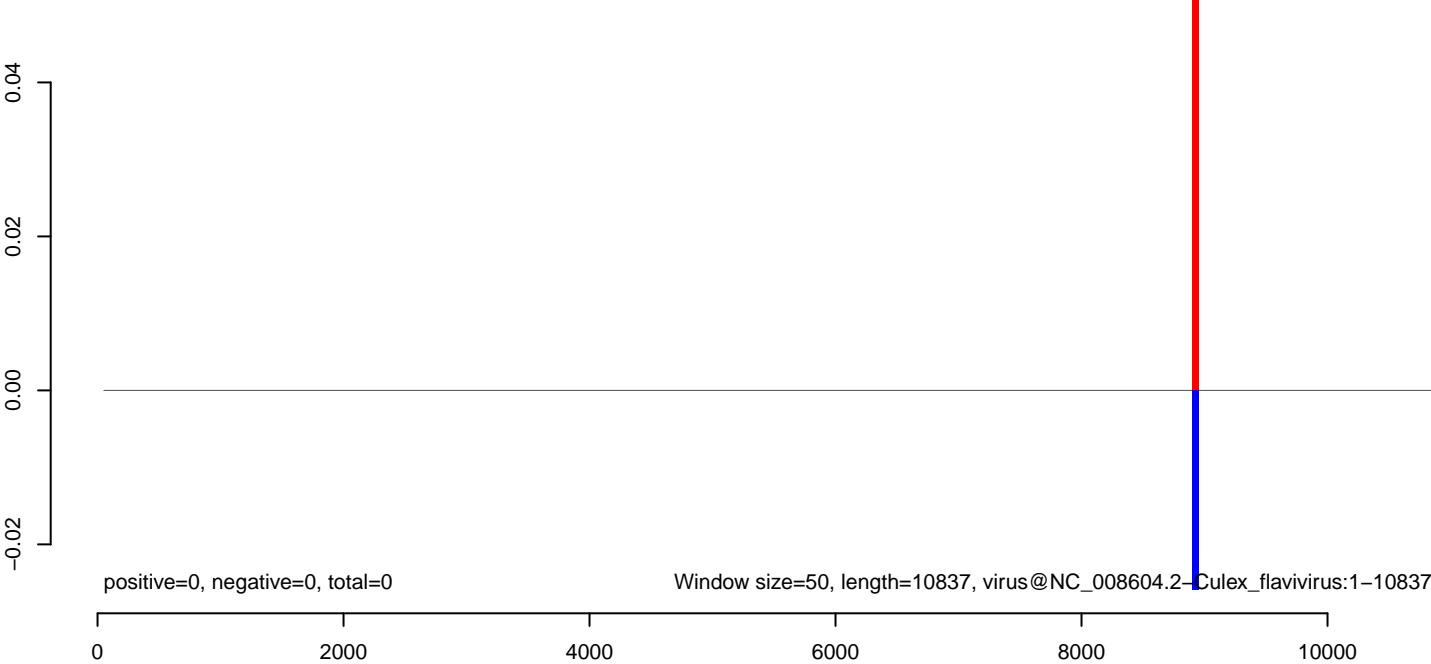
### AeAeg\_Fcarc\_GH.18\_23.rep



### AeAeg\_Fcarc\_GH.24\_35.rep



### AeAeg\_Fcarc\_GH.rep



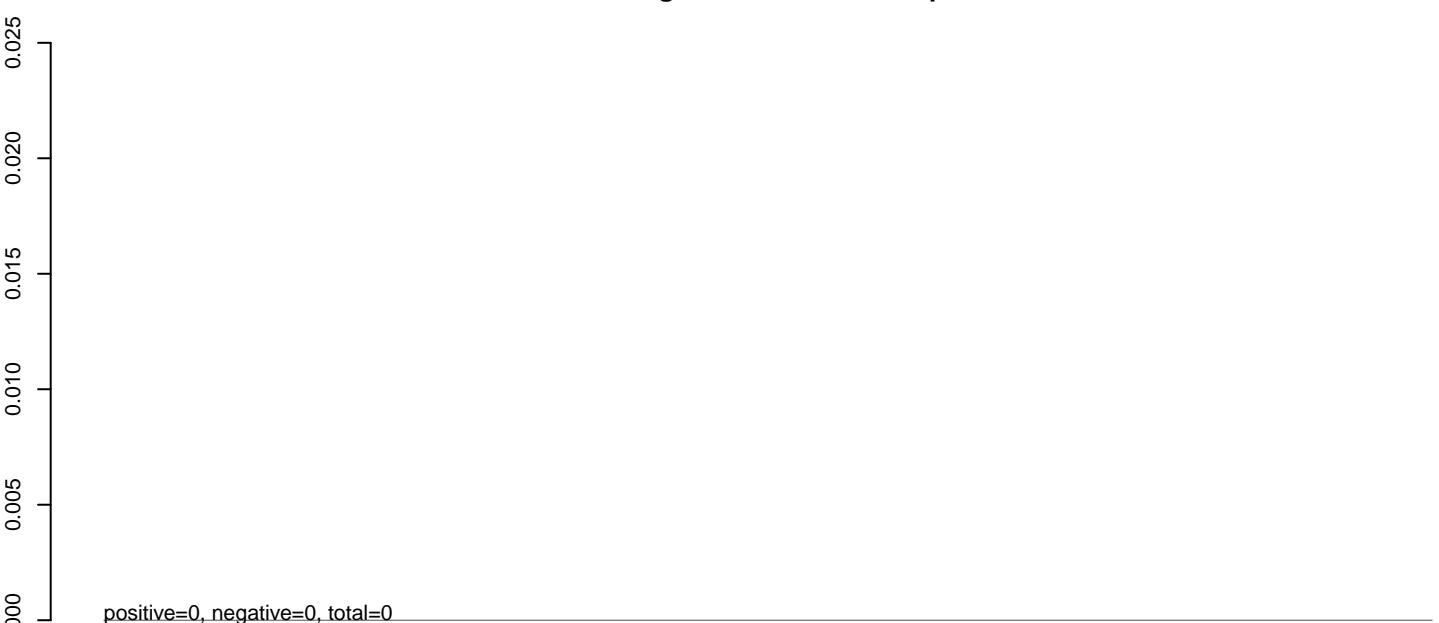
### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

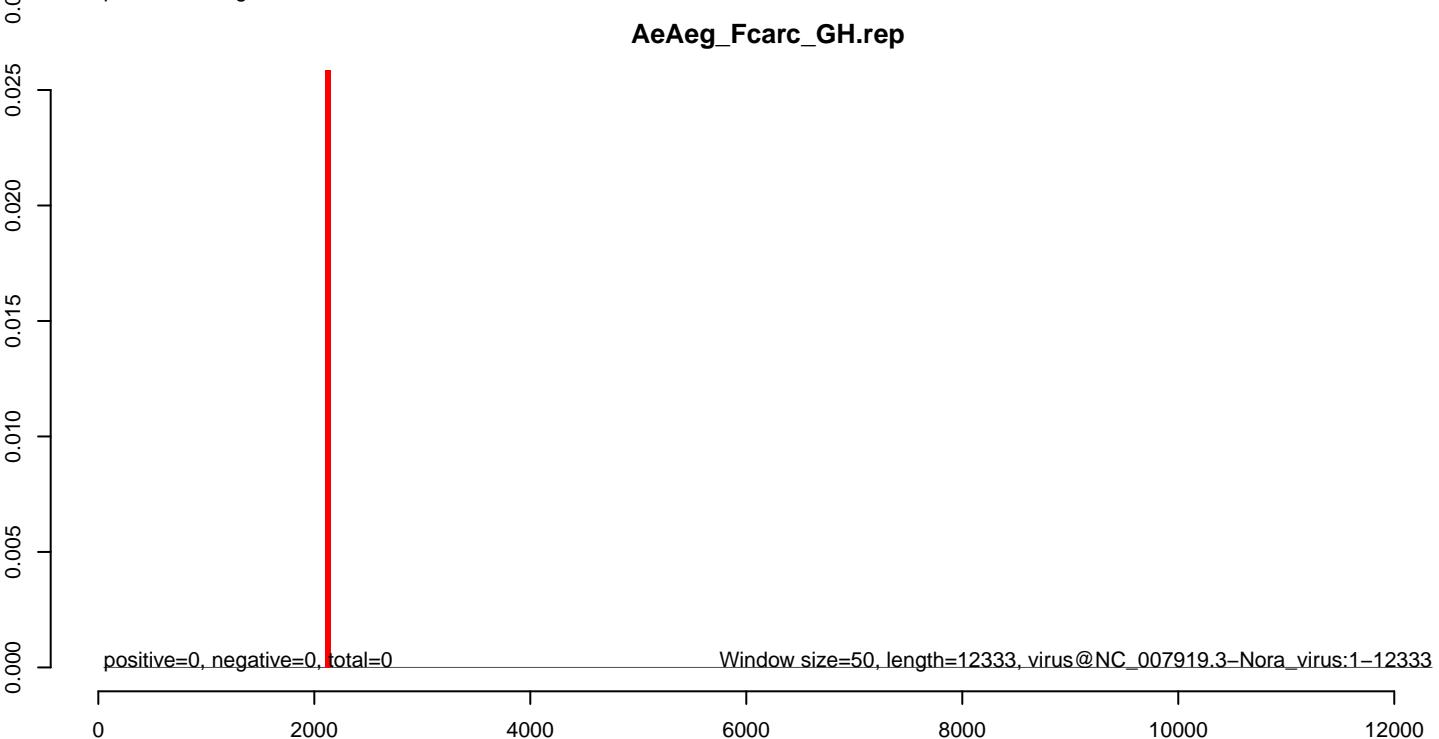
positive=0, negative=0, total=0



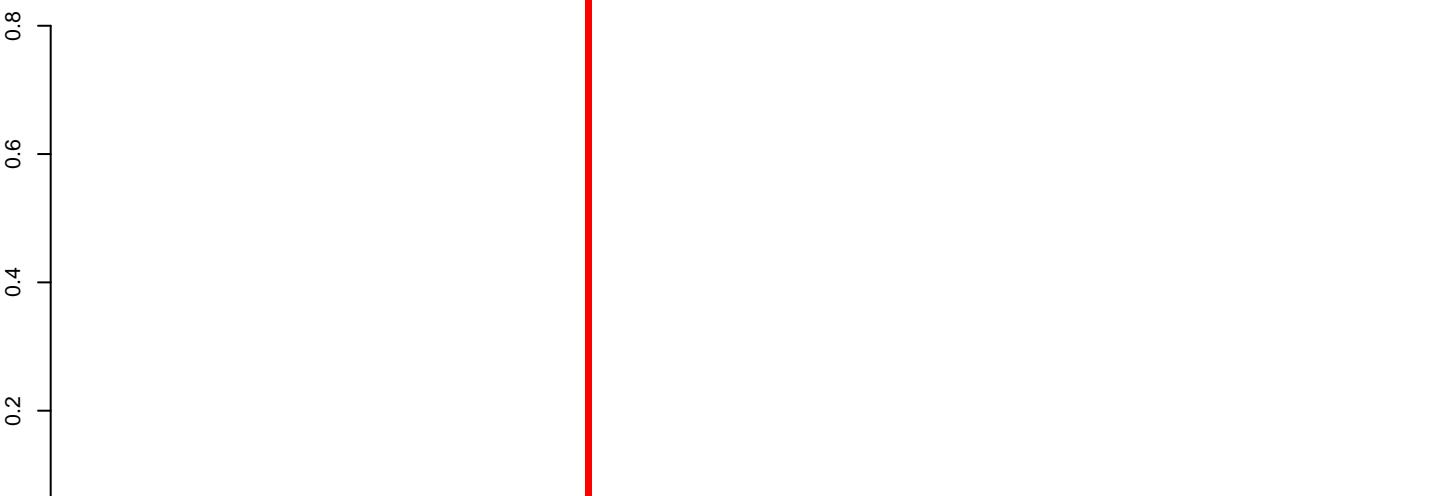
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=12333, virus@NC\_007919.3-Nora\_virus:1-12333



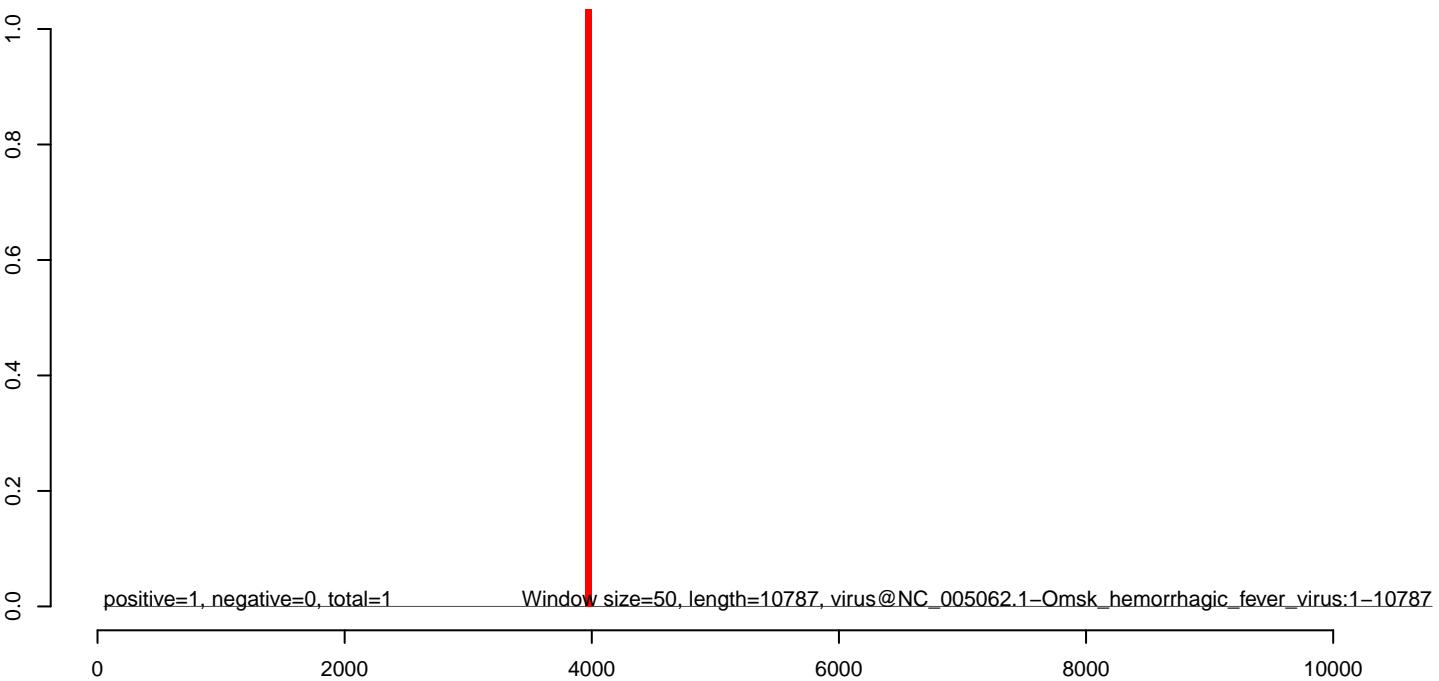
### AeAeg\_Fcanc\_GH.18\_23.rep



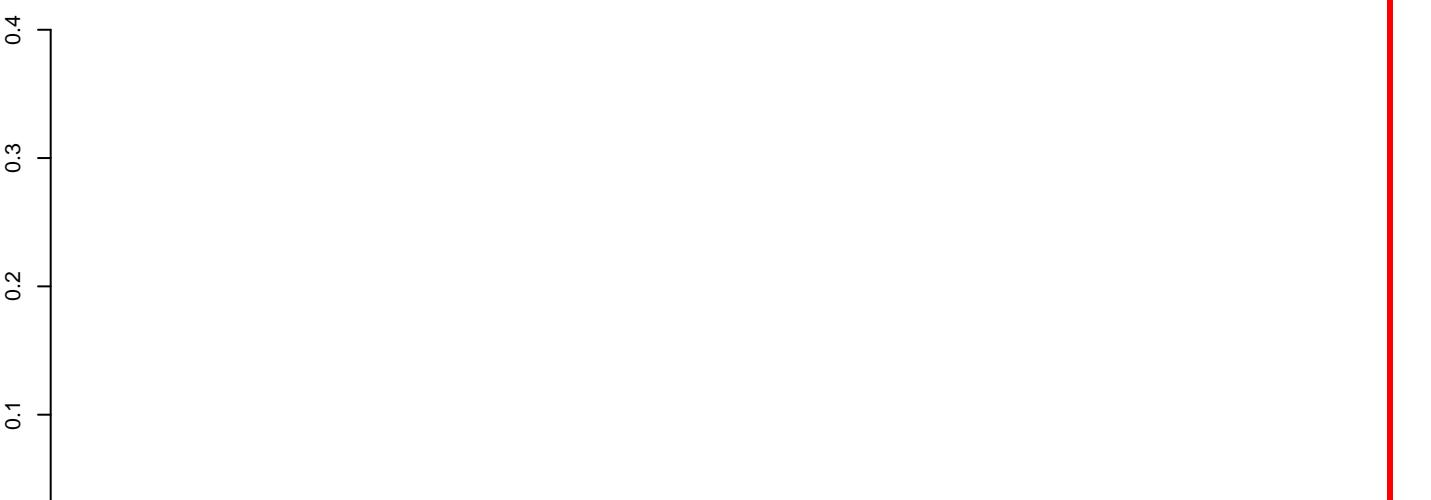
### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



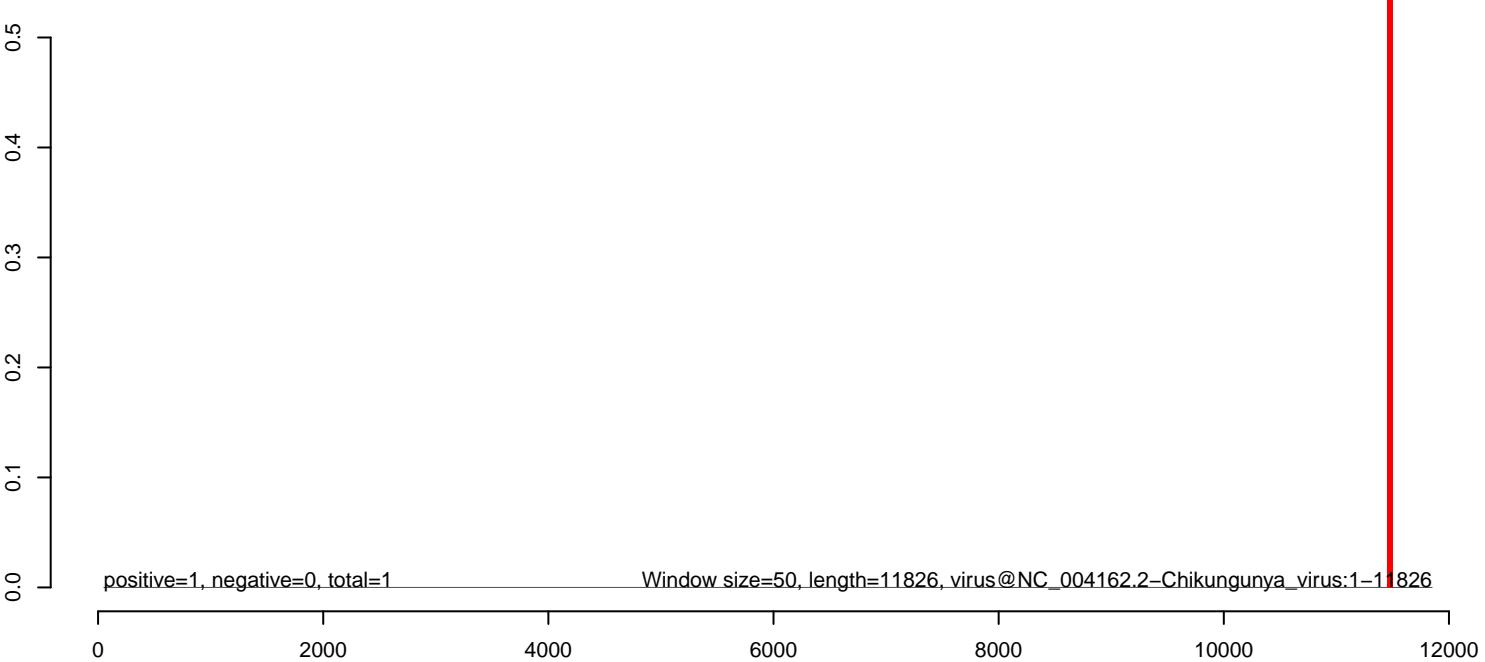
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=4507, virus@NC\_004146.1-Flock\_house\_virus\_RNA\_1\_and\_RNA2:1-4507

0 1000 2000 3000 4000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10690, virus@NC\_004119.1-Montana\_myotis\_leukoencephalitis\_virus:1-10690



0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0 length=12491, virus@NC\_004110.1-La\_Crosse\_virus\_segment\_S\_plus\_segment\_M\_plus\_segment\_L:1-12491



0 2000 4000 6000 8000 10000 12000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

Vind positive=50, negative=0, total=0 @ NC\_003924.1-Cricket\_paralysis\_virus\_nonstructural\_polyprotein\_and\_structural\_polyprotein\_genes:1-9185

0 2000 4000 6000 8000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10976, virus @NC\_001437.1–Japanese\_encephalitis\_virus\_genome:1–10976

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

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

0 1000 2000 3000 4000

### AeAeg\_Fcarc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcarc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcarc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=11014, virus@NC\_000943.1–Murray\_Valley\_encephalitis\_virus:1–11014

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0, length=5982, virus@MF176317.1-Culex\_mononegative-like\_virus\_1\_strain\_mos191X45127\_segment\_2:1-5982

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0, window size=50, length=11835, virus@M20303.1-O-Nyong-nyong\_virus\_Gulu\_strain\_complete\_genome:1-11835



0 2000 4000 6000 8000 10000 12000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0, size=50, length=10749, virus@KX669682.1-Calbertado\_virus\_isolate\_Cal\_mos\_2\_polyprotein\_gene:1-10749

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10560, virus@KM225263.1-Stratford\_virus\_strain\_C338:1-10560

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



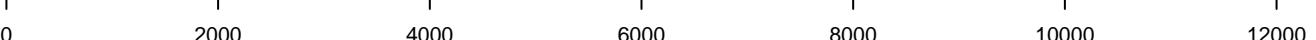
### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

50, length=12580 negative@KM201990.1-Muir\_Springs\_virus\_nucleoprotein\_phosphoprotein\_matrix\_glycoprotein\_and\_polymerase\_genes:1-12580



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

Wip positive=0, negative=0, total=0  
length=12297, Virus@KC608151.1-llesha\_virus\_strain\_ILESCHA\_8e\_segment\_S\_plus\_segment\_M\_plus\_segment\_L:1-12297

0 2000 4000 6000 8000 10000 12000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

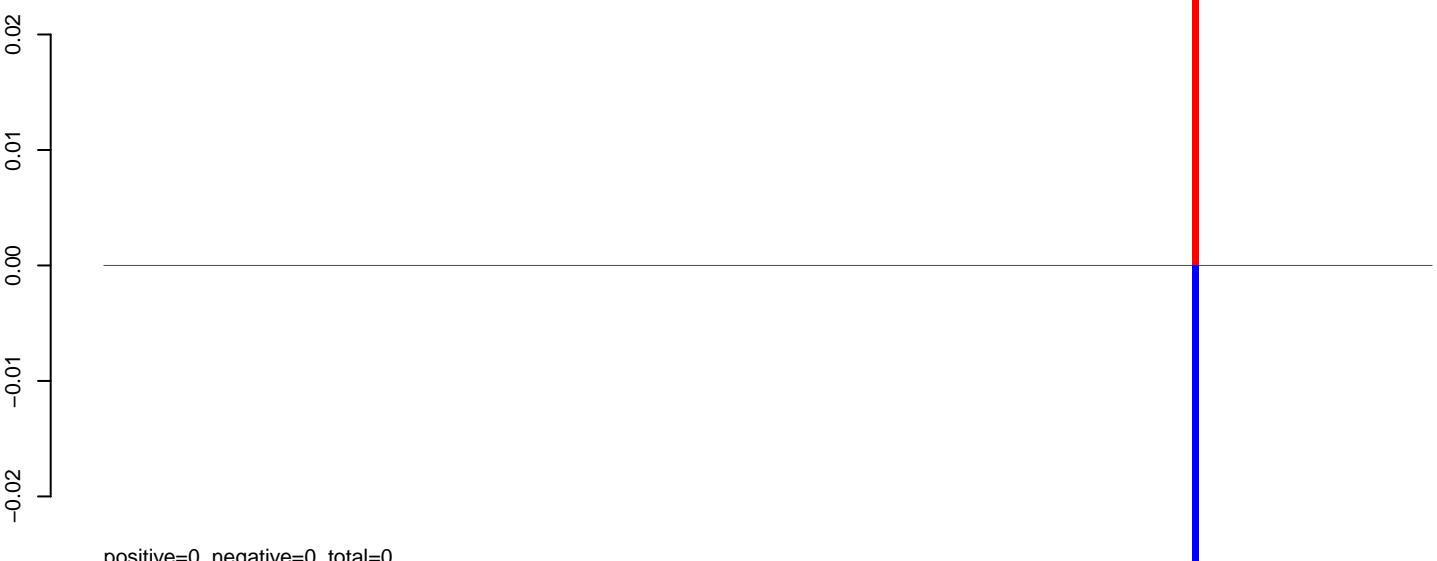


### AeAeg\_Fcanc\_GH.rep

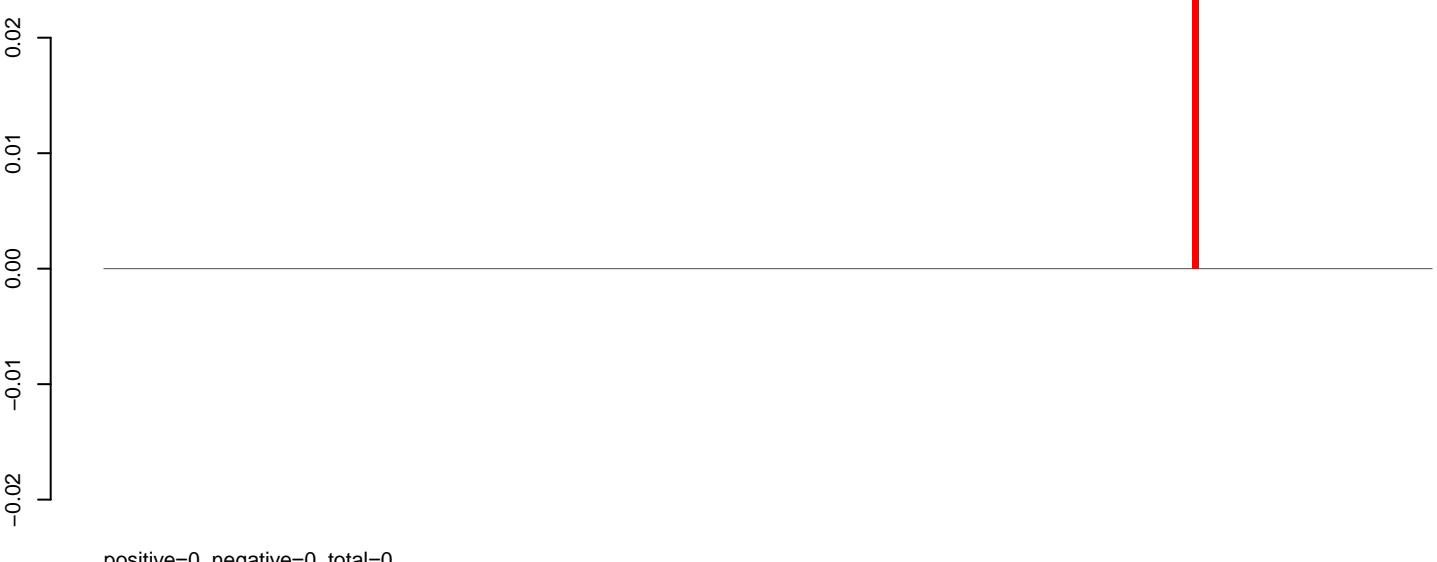
ze=500 density=0.529gavivs@Goma12965.1-Drosophila\_melanogaster\_American\_nodavirus\_ANV\_SW-2009a\_segment\_RNA1\_and\_RNA2:1-4523

0 1000 2000 3000 4000

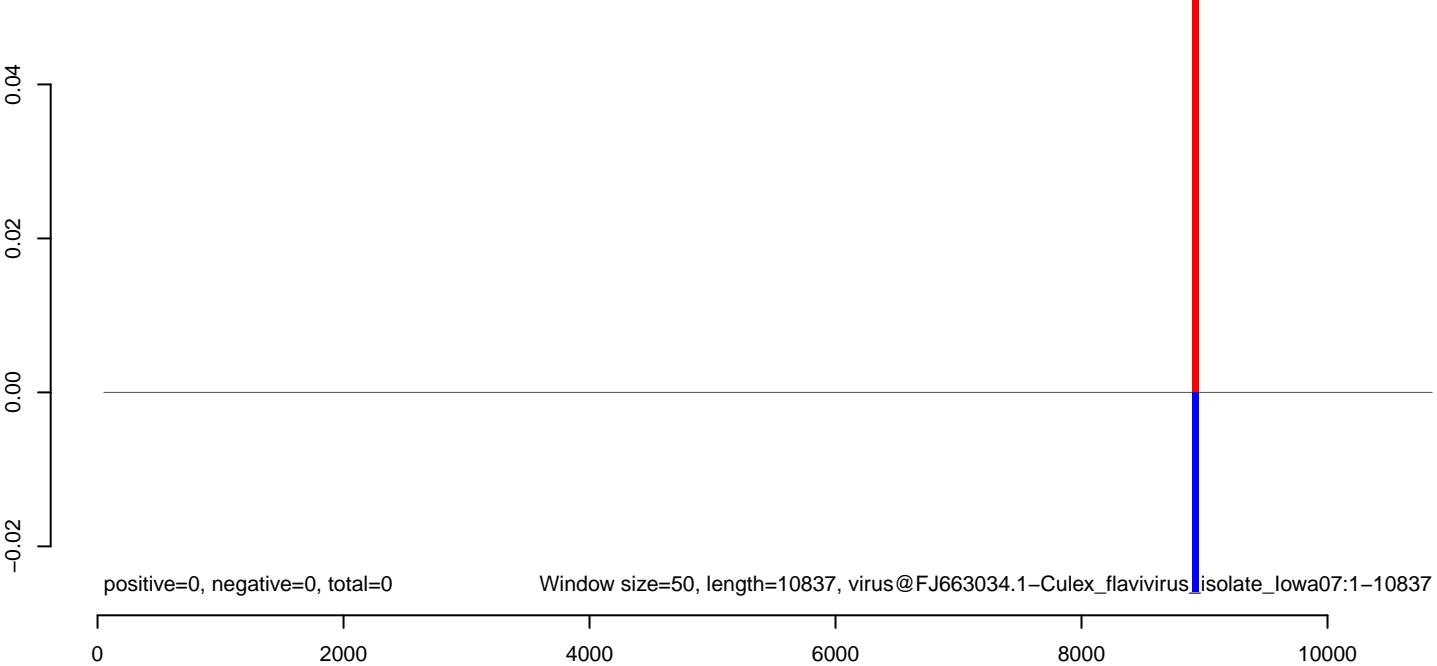
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0, size=50, length=10254, virus@DQ235149.1-Royal\_Farm\_virus\_from\_Afghanistan\_polyprotein gene:1-10254



0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10922, virus@AY632538.4–Iguape\_virus\_strain\_SPAn\_71686:1–10922

0 2000 4000 6000 8000 10000

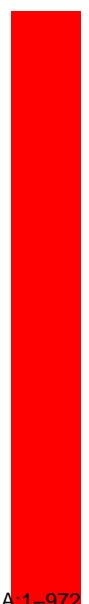
### AeAeg\_Fcarc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcarc\_GH.24\_35.rep

positive=0, negative=0, total=0



### AeAeg\_Fcarc\_GH.rep

positive=0, negative=0, total=50, length=972, virus@AF411844.1-Elaeis\_guineensis\_MADS\_box\_transcription\_factor\_AGL2-2\_mRNA-1-972



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0



### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0



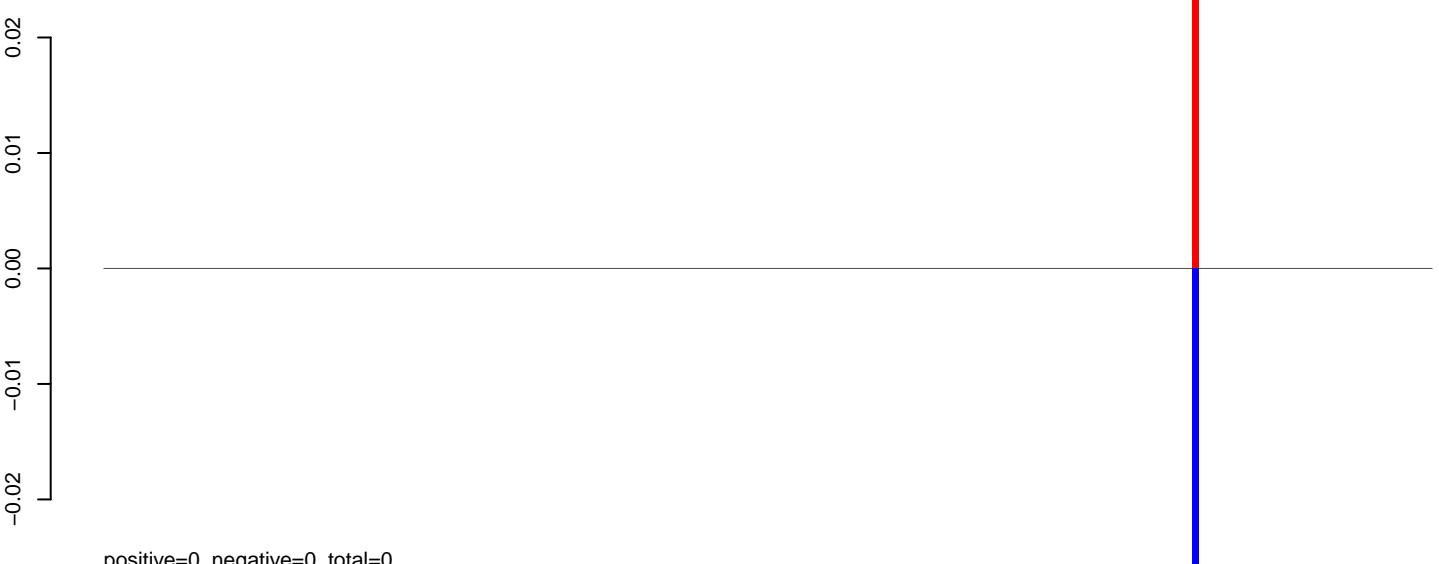
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

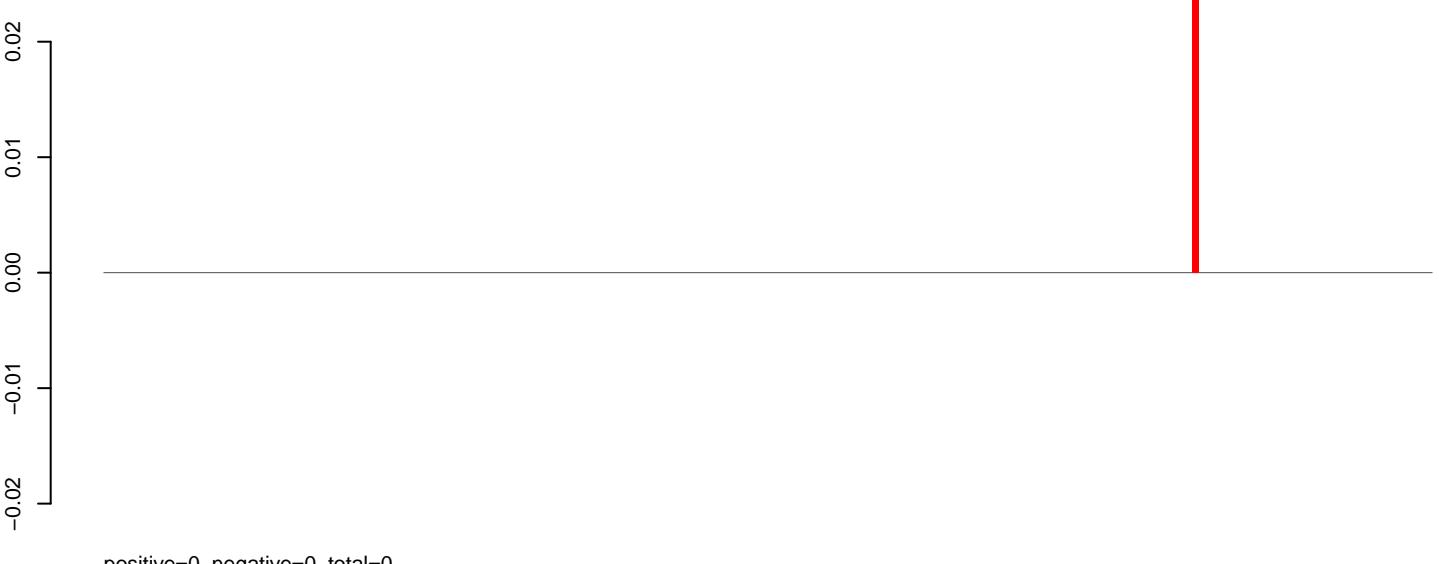
Window size=50, length=11385, virus@AF075259.1-Cabassou\_virus\_strain\_CaAr\_508:1-11385

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep



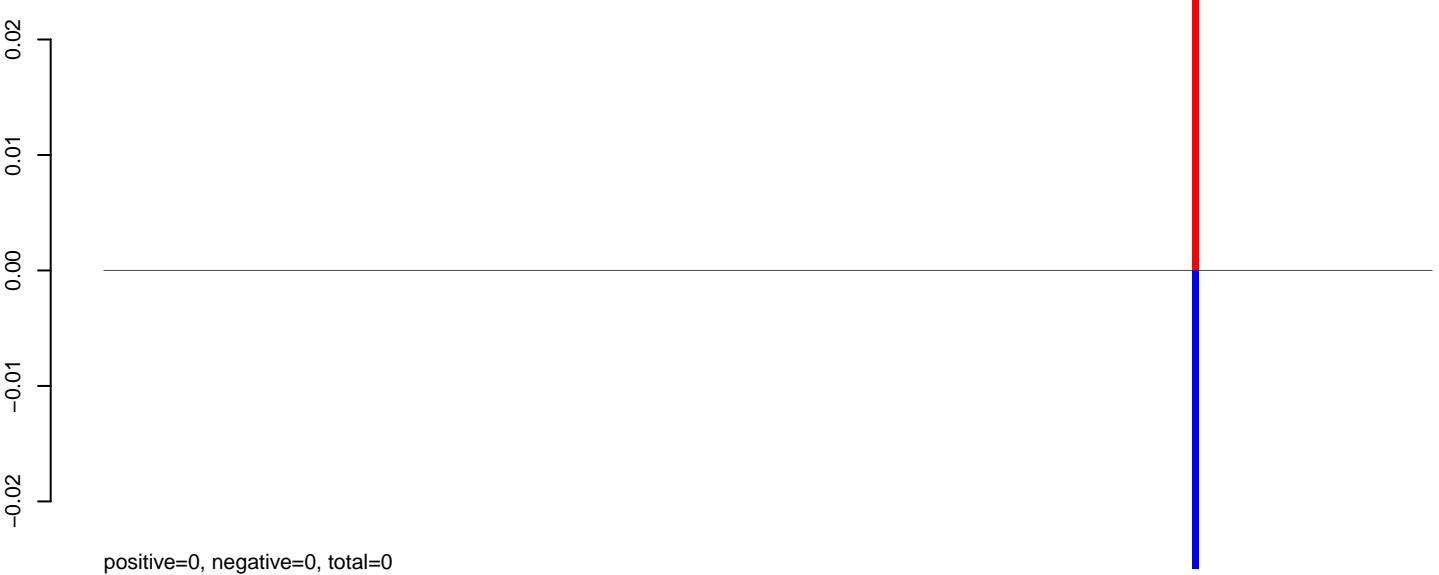
### AeAeg\_Fcanc\_GH.24\_35.rep



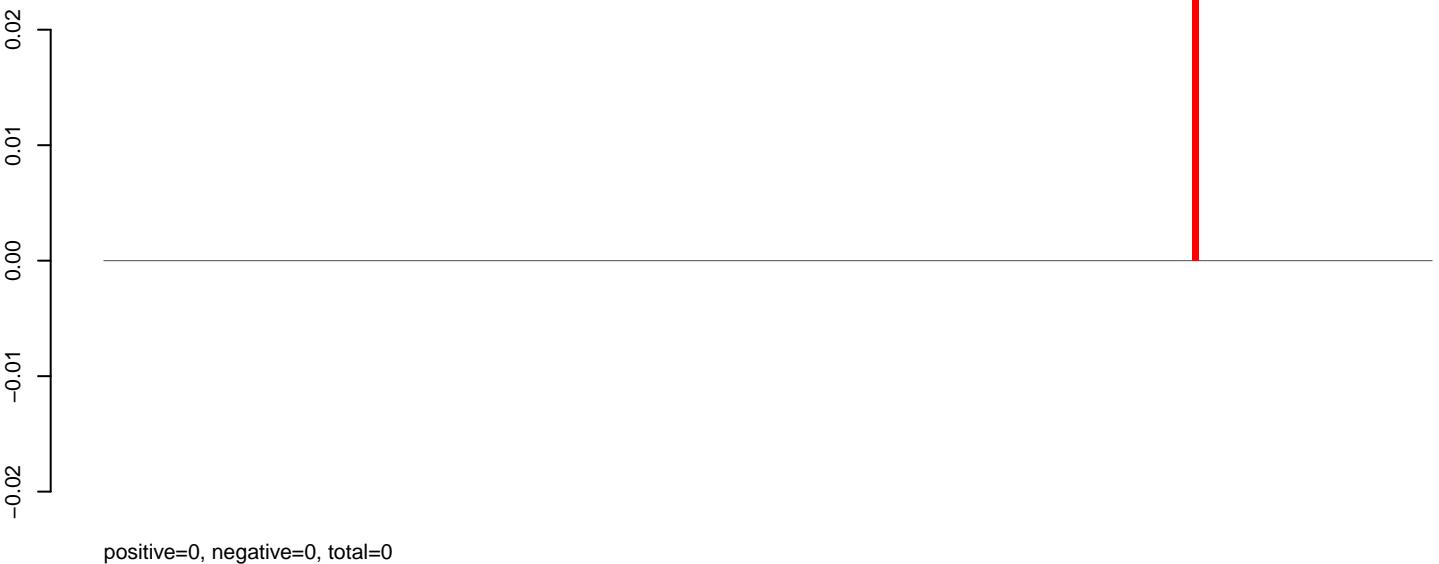
### AeAeg\_Fcanc\_GH.rep



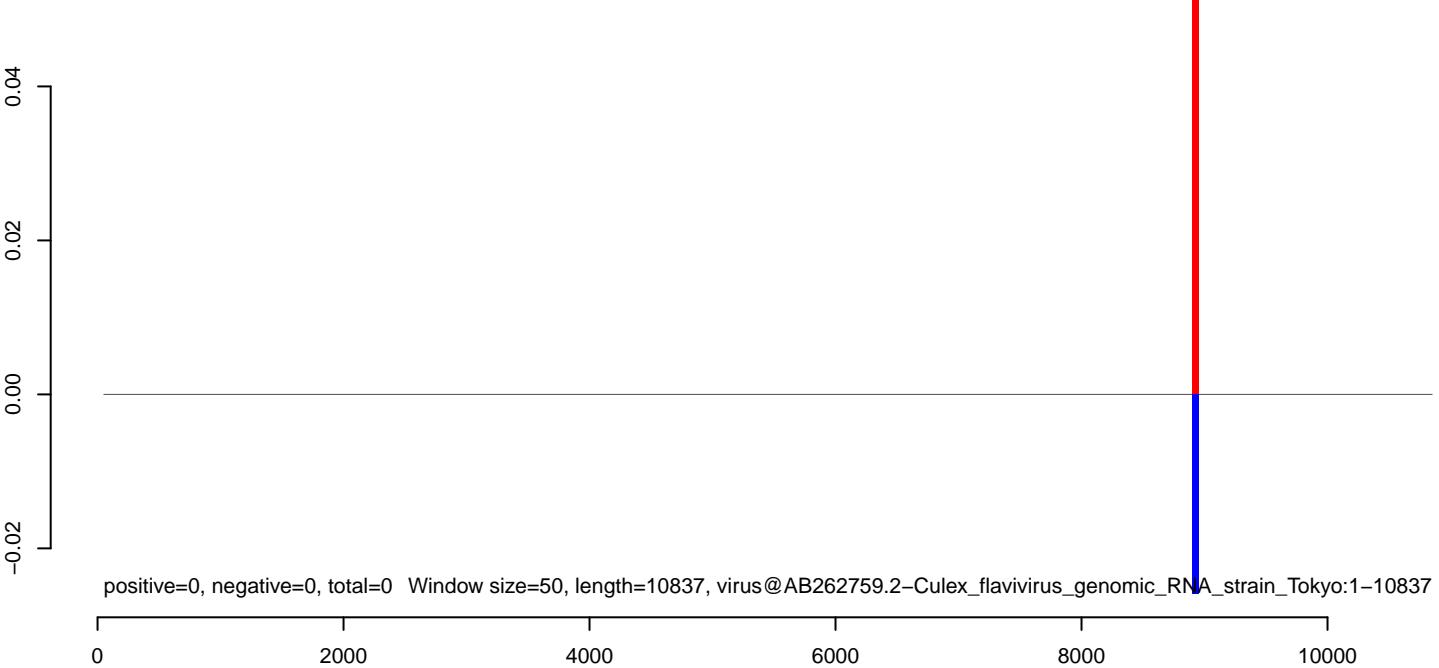
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

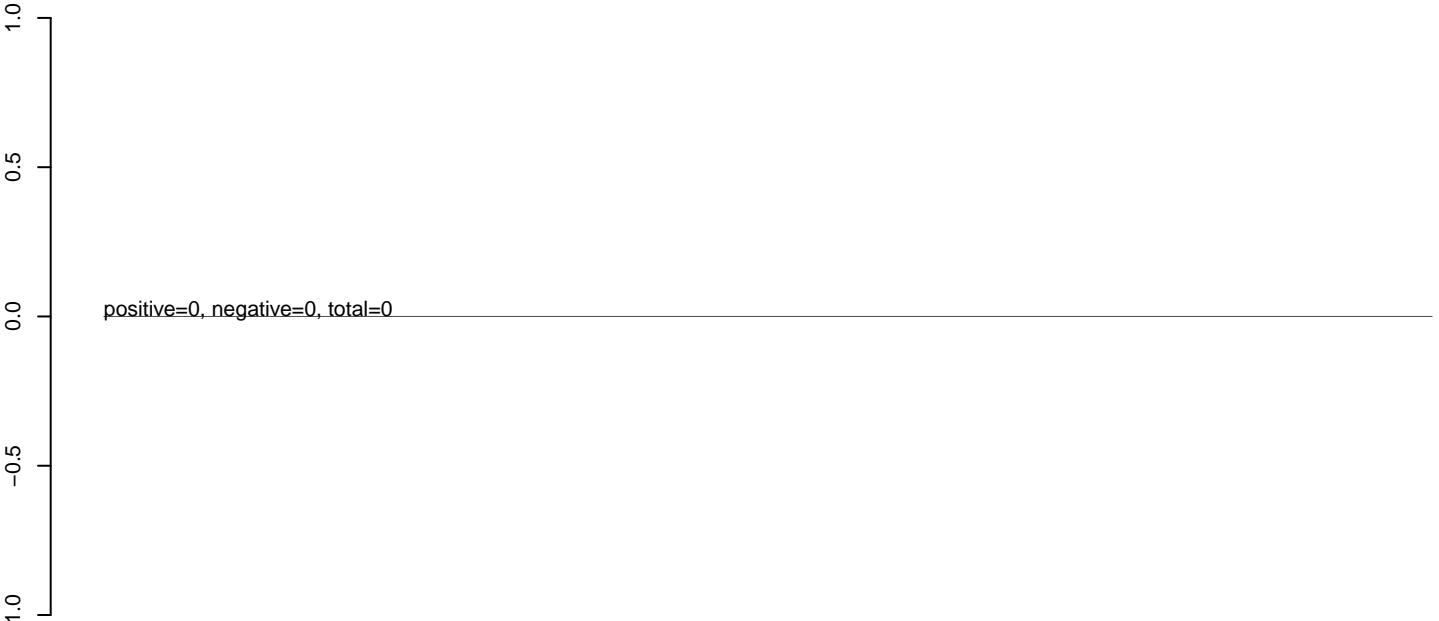
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

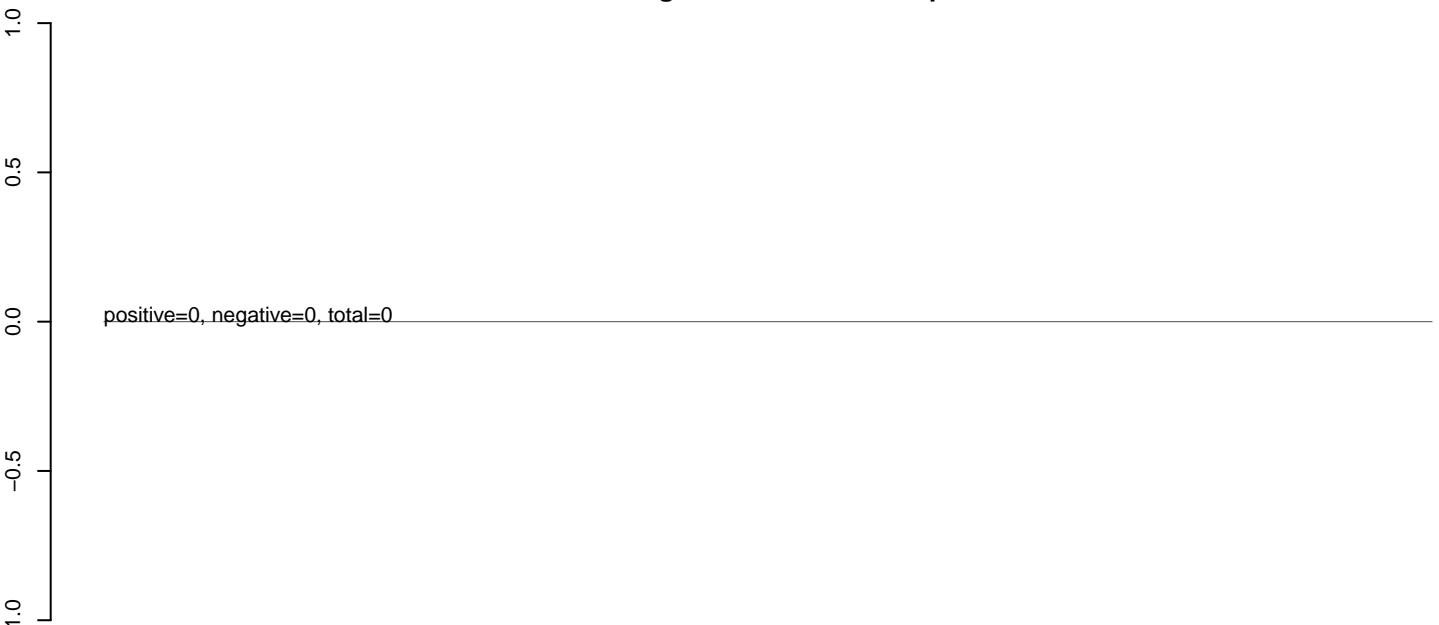
Window size=50, length=3893, Virus@GU452720.1-Mosquito\_densovirus\_BR\_07:1-3893

0 1000 2000 3000 4000

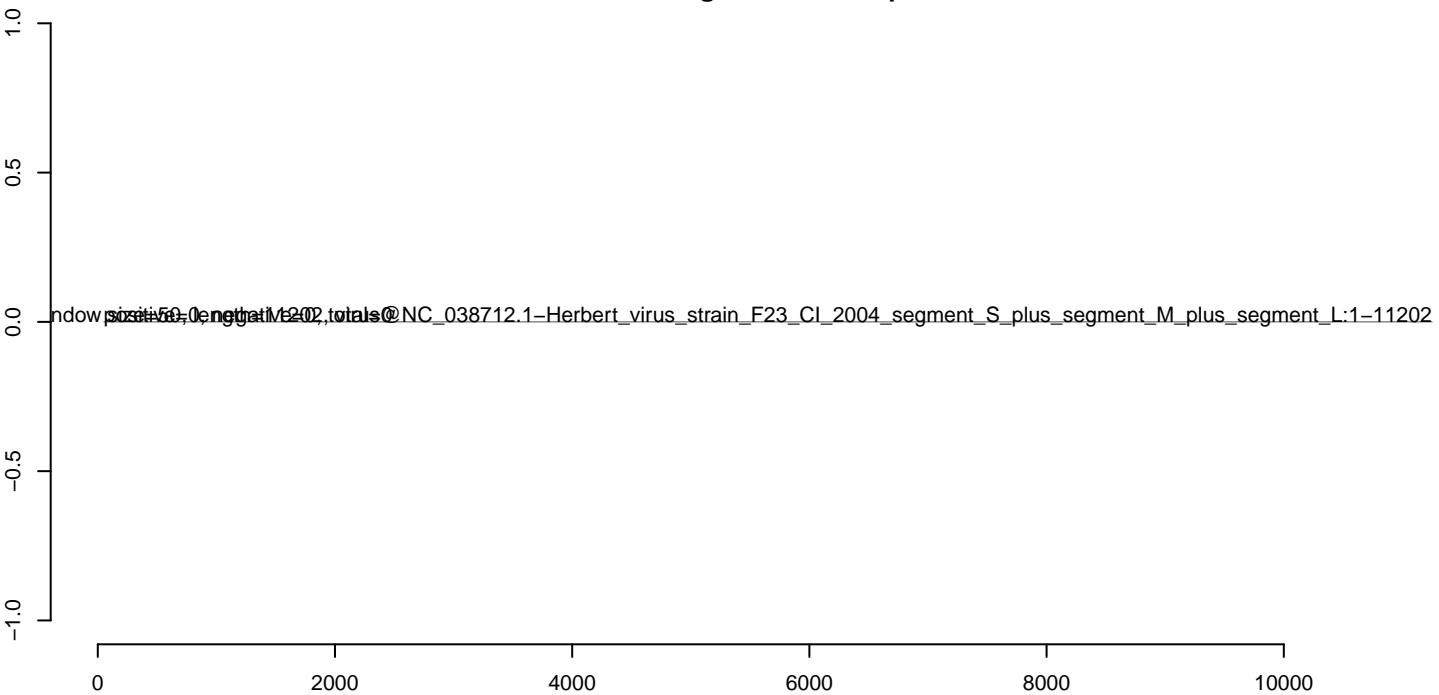
### AeAeg\_Fcanc\_GH.18\_23.rep



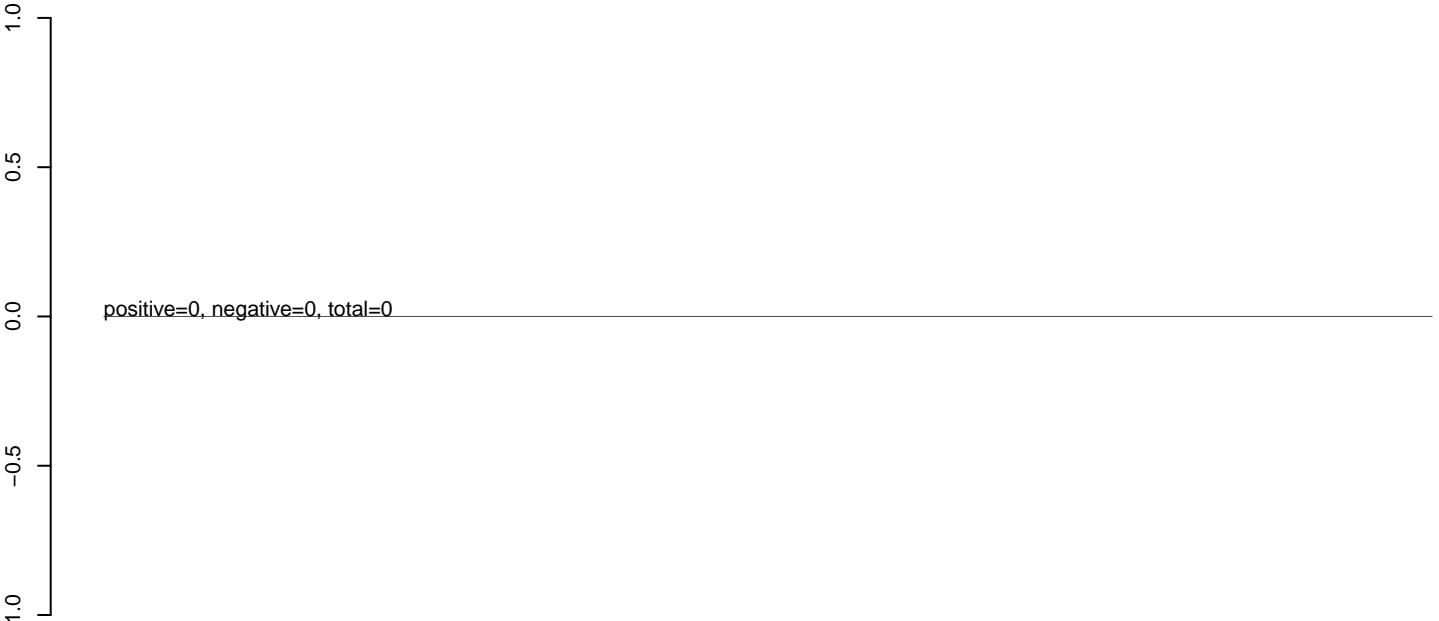
### AeAeg\_Fcanc\_GH.24\_35.rep



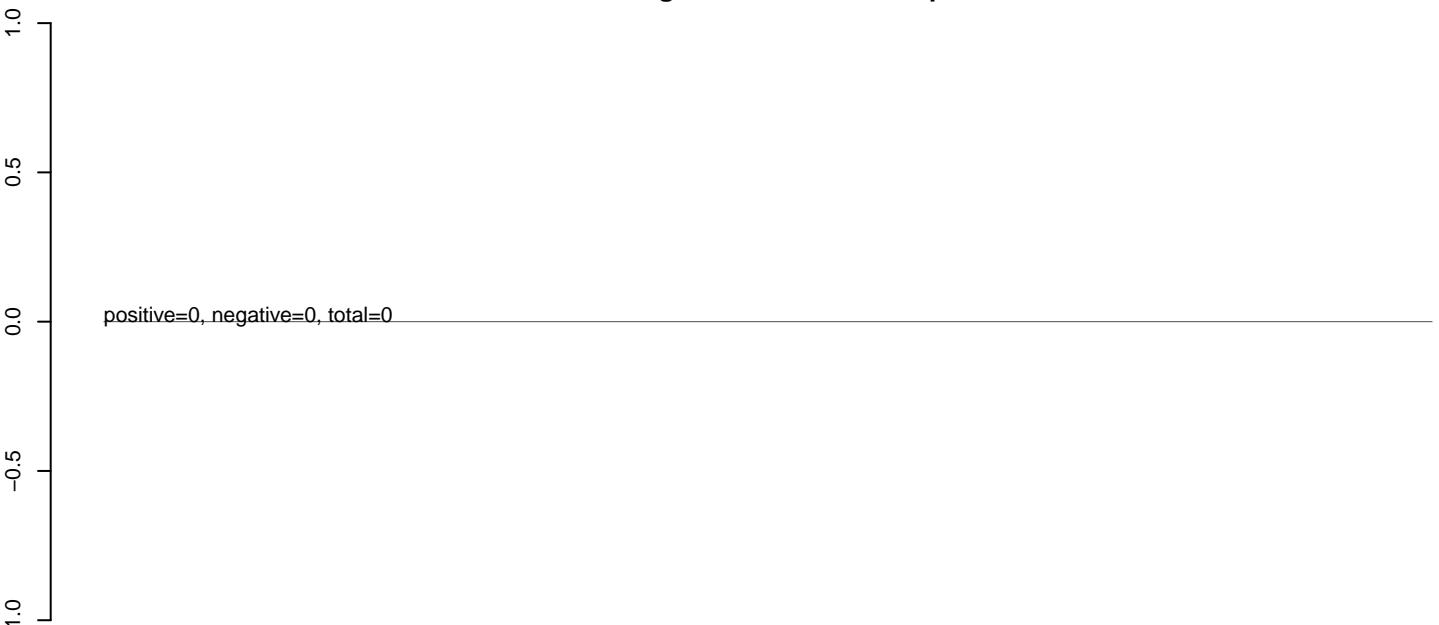
### AeAeg\_Fcanc\_GH.rep



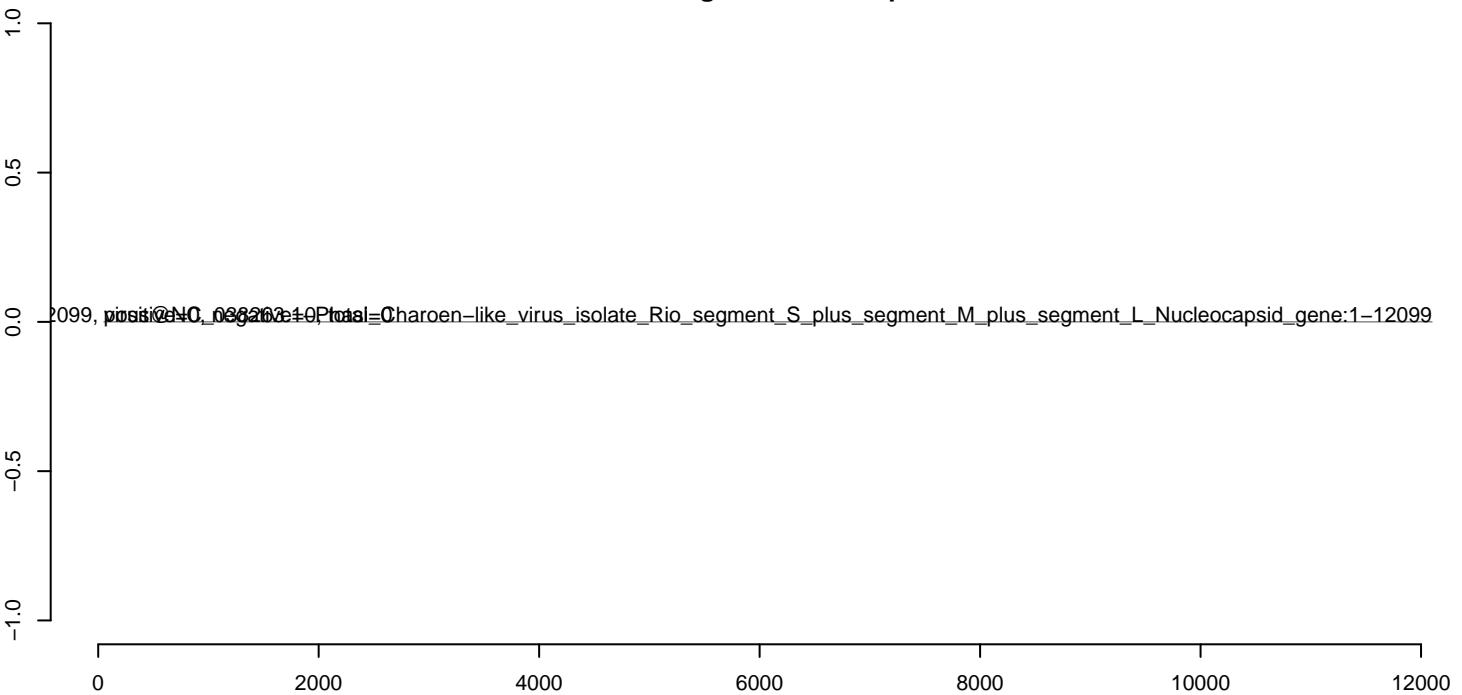
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep





positive=0, negative=0, total=0

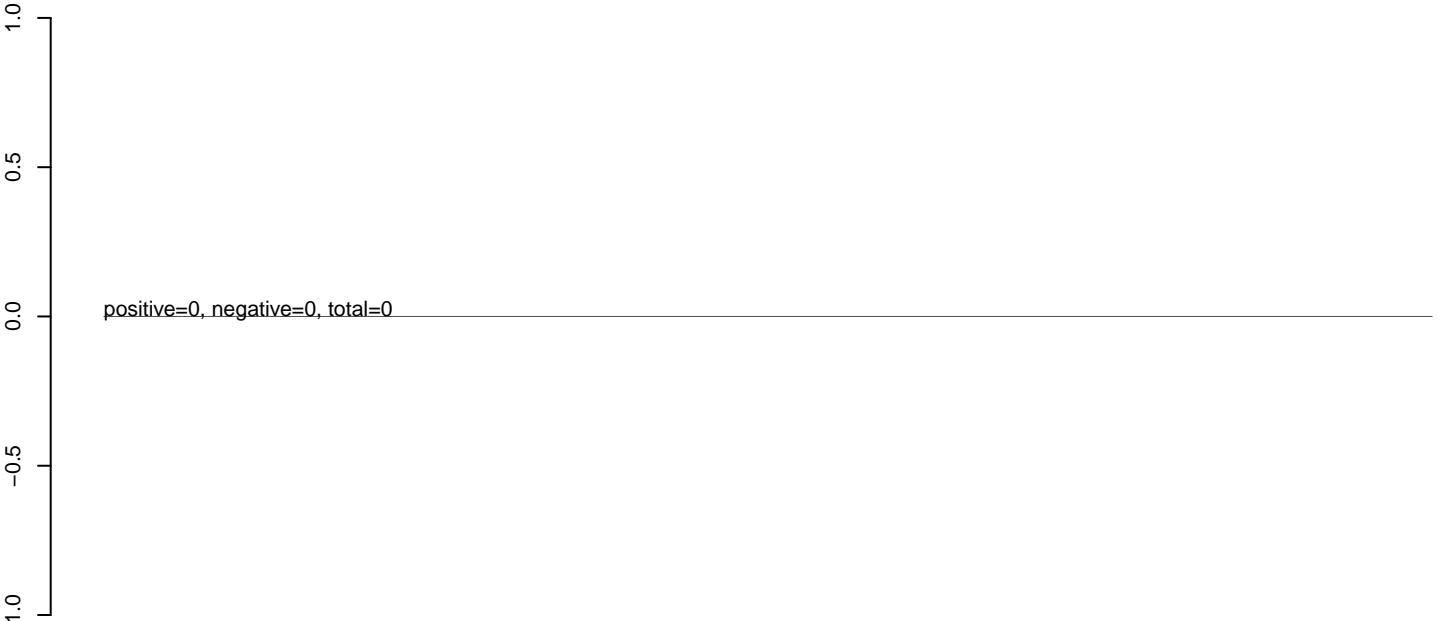


positive=0, negative=0, total=0

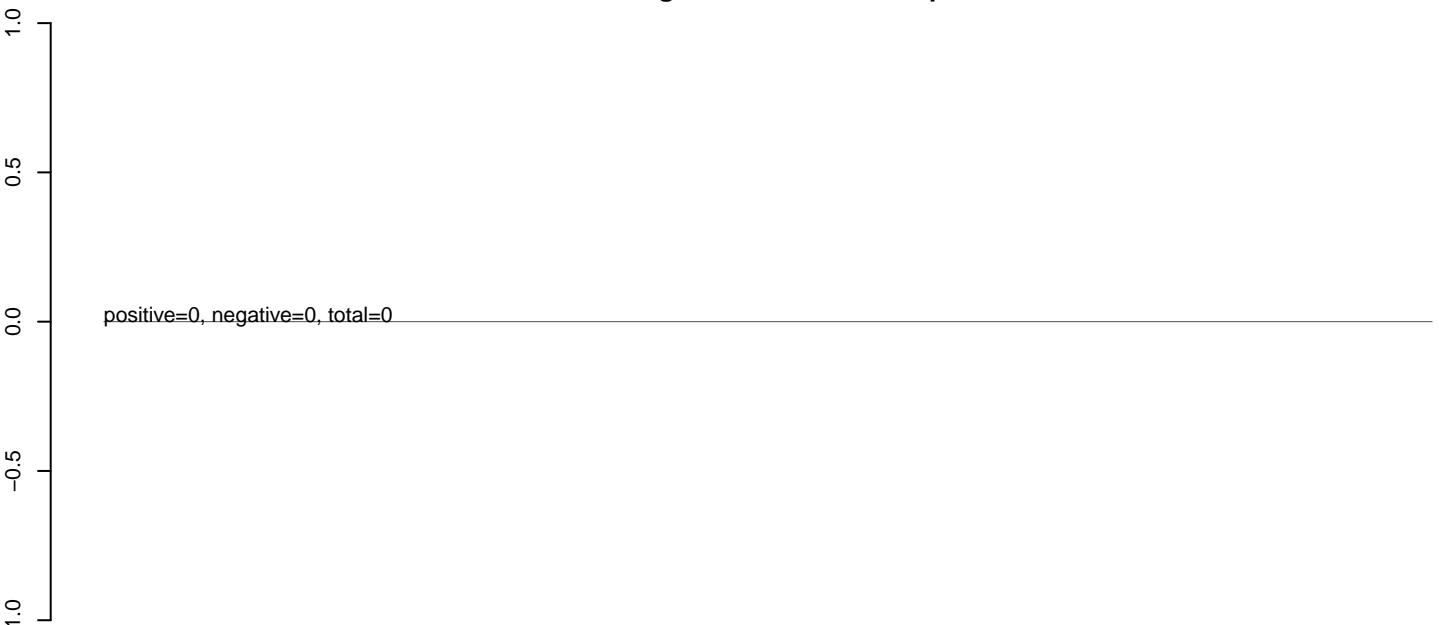


Virion positive for SARS-CoV-2, isolate TS6347, virus@NC\_038259.1-Badu\_virus\_isolate\_TS6347\_segment\_S\_plus\_segment\_M\_plus\_segment\_L:1-12247

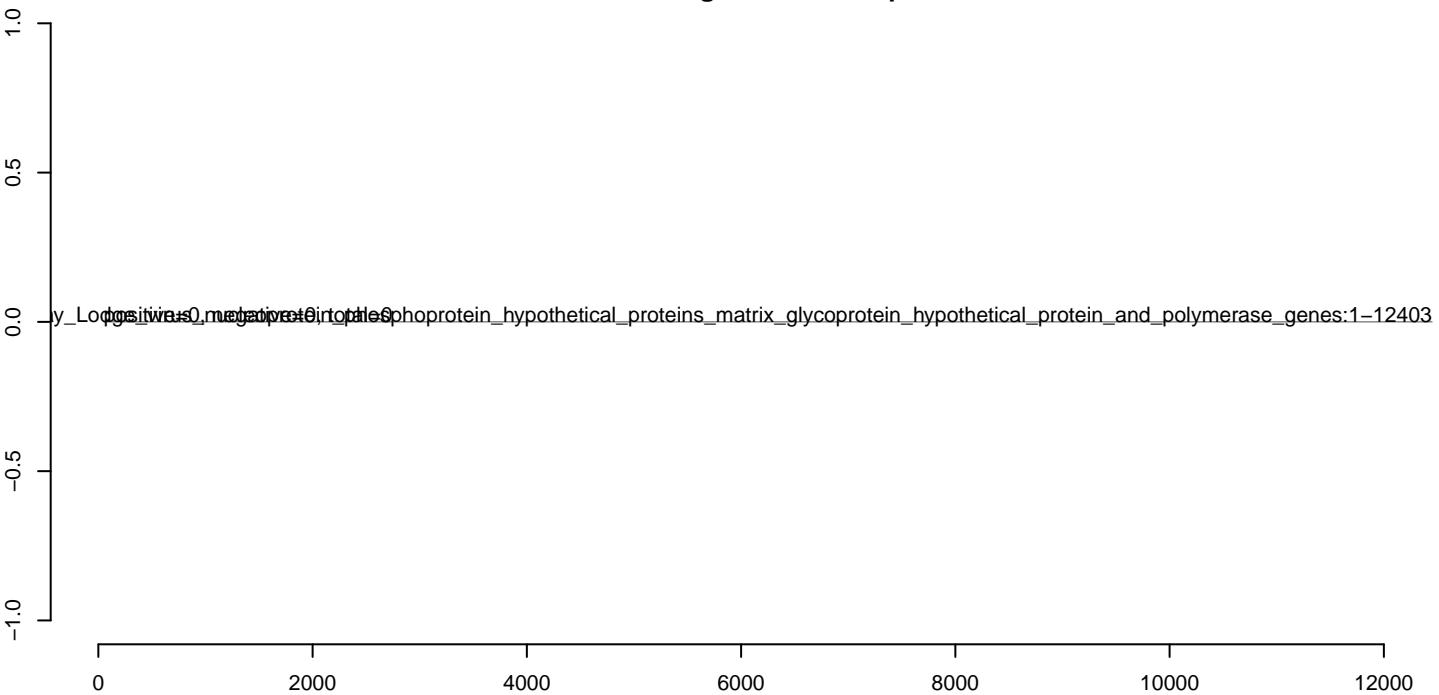
### AeAeg\_Fcanc\_GH.18\_23.rep



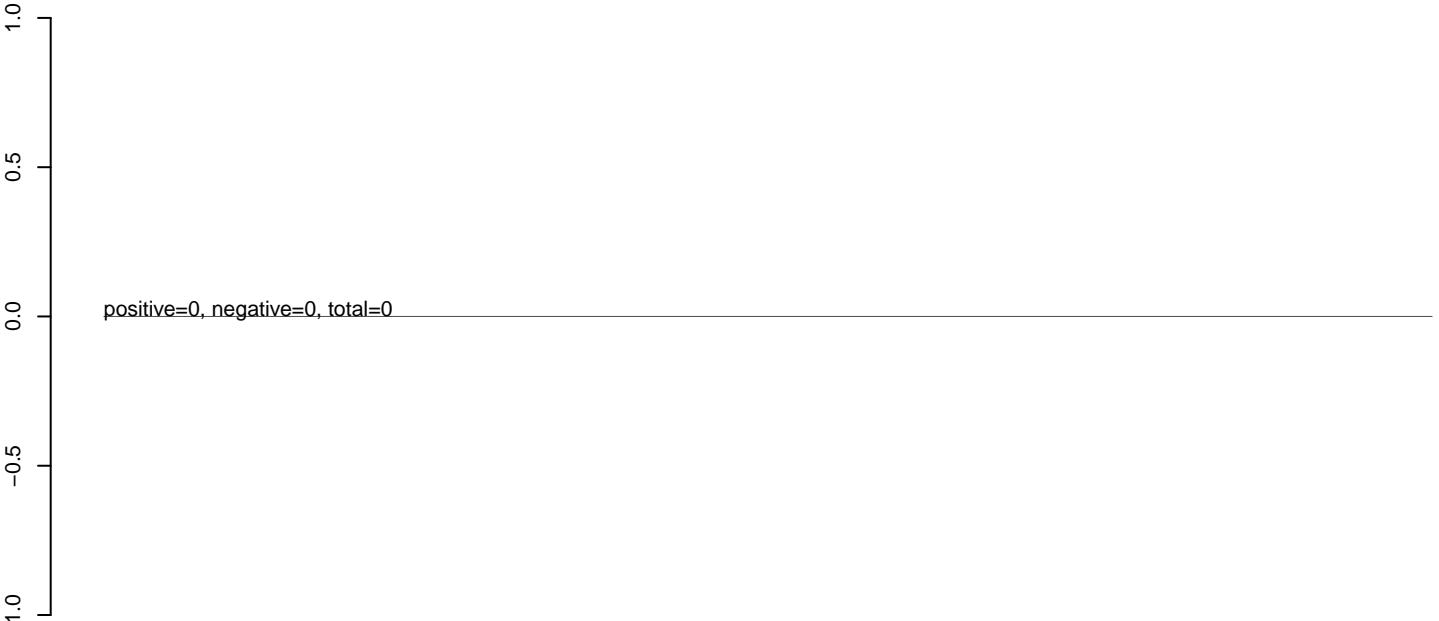
### AeAeg\_Fcanc\_GH.24\_35.rep



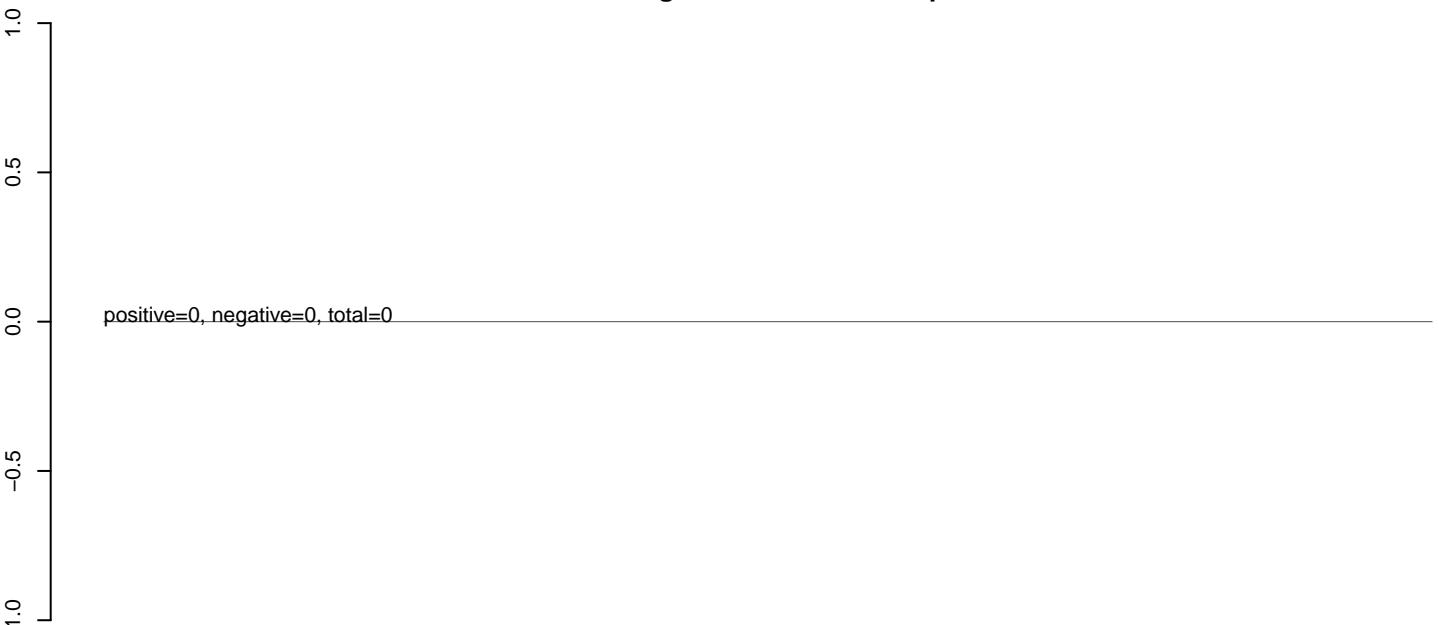
### AeAeg\_Fcanc\_GH.rep



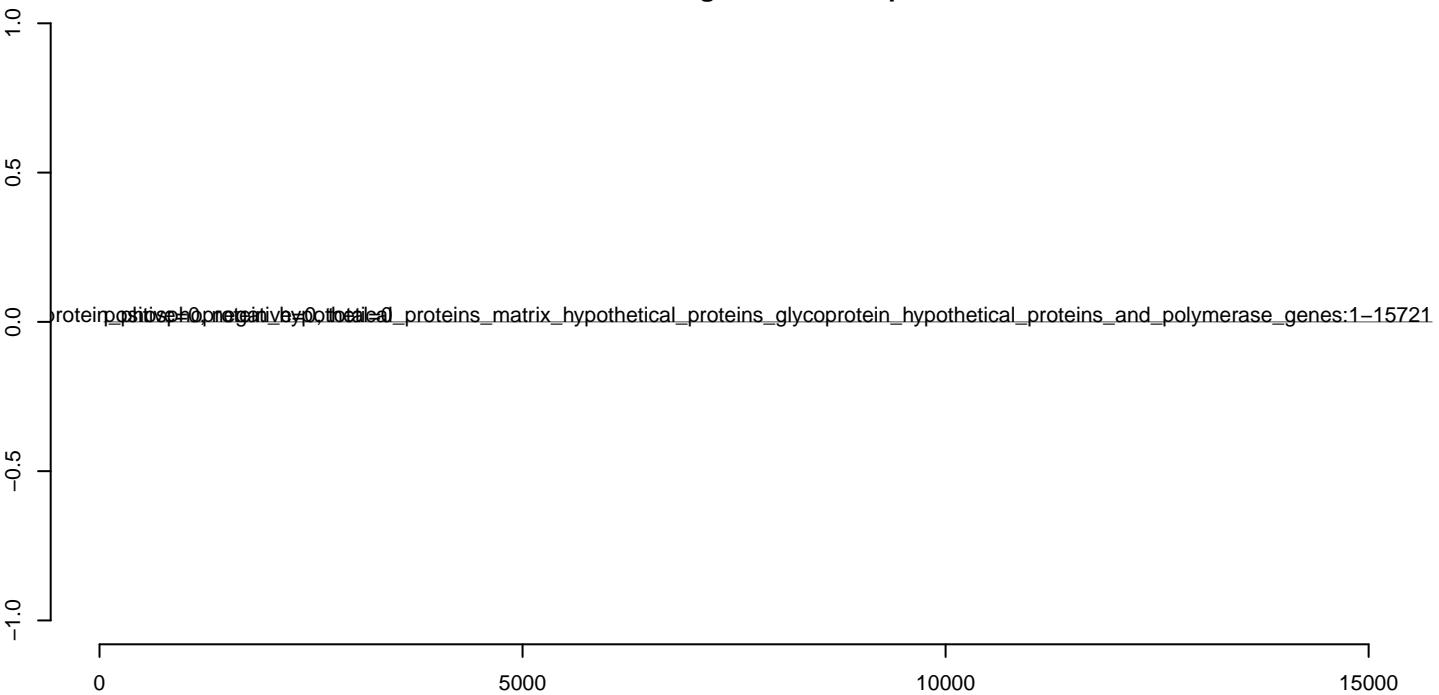
### AeAeg\_Fcanc\_GH.18\_23.rep



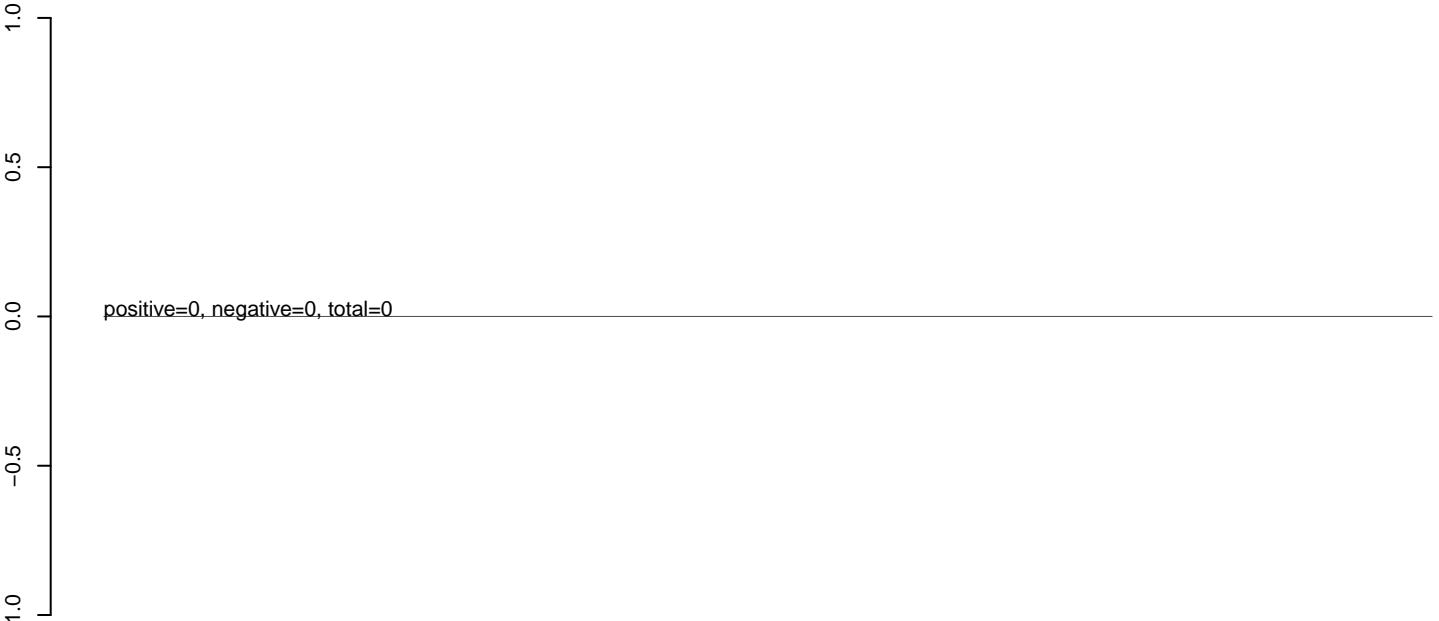
### AeAeg\_Fcanc\_GH.24\_35.rep



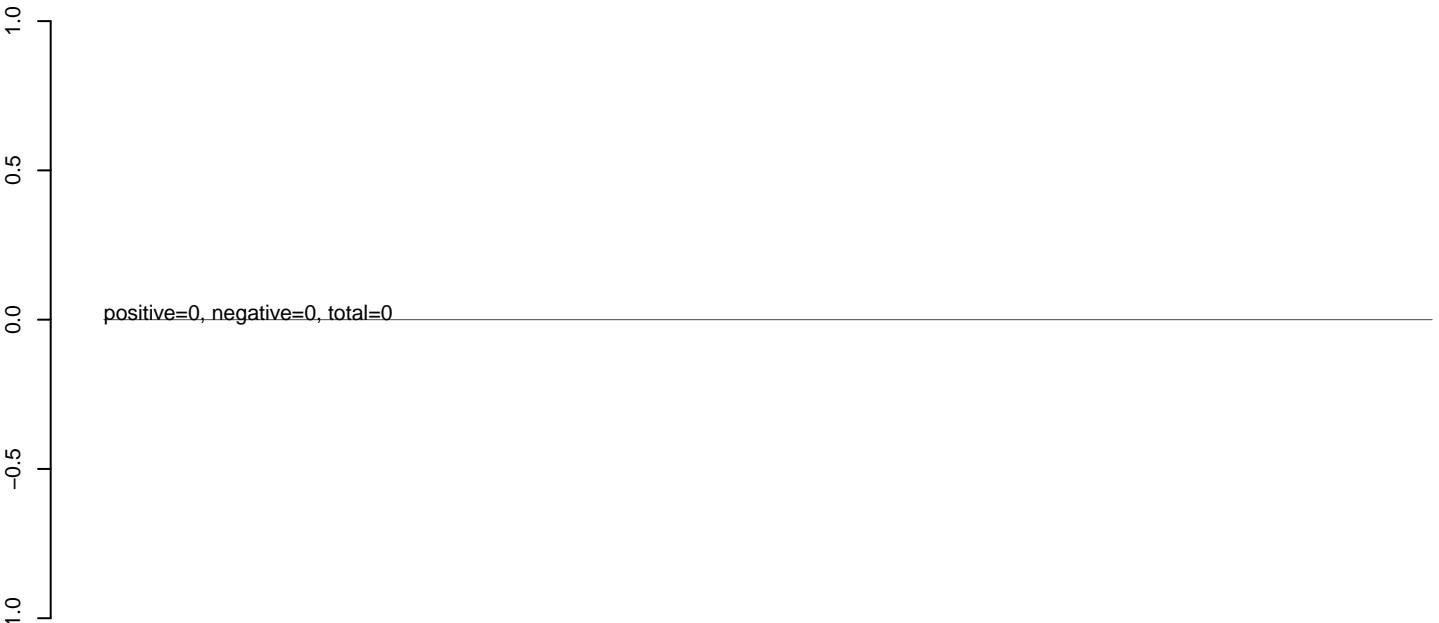
### AeAeg\_Fcanc\_GH.rep



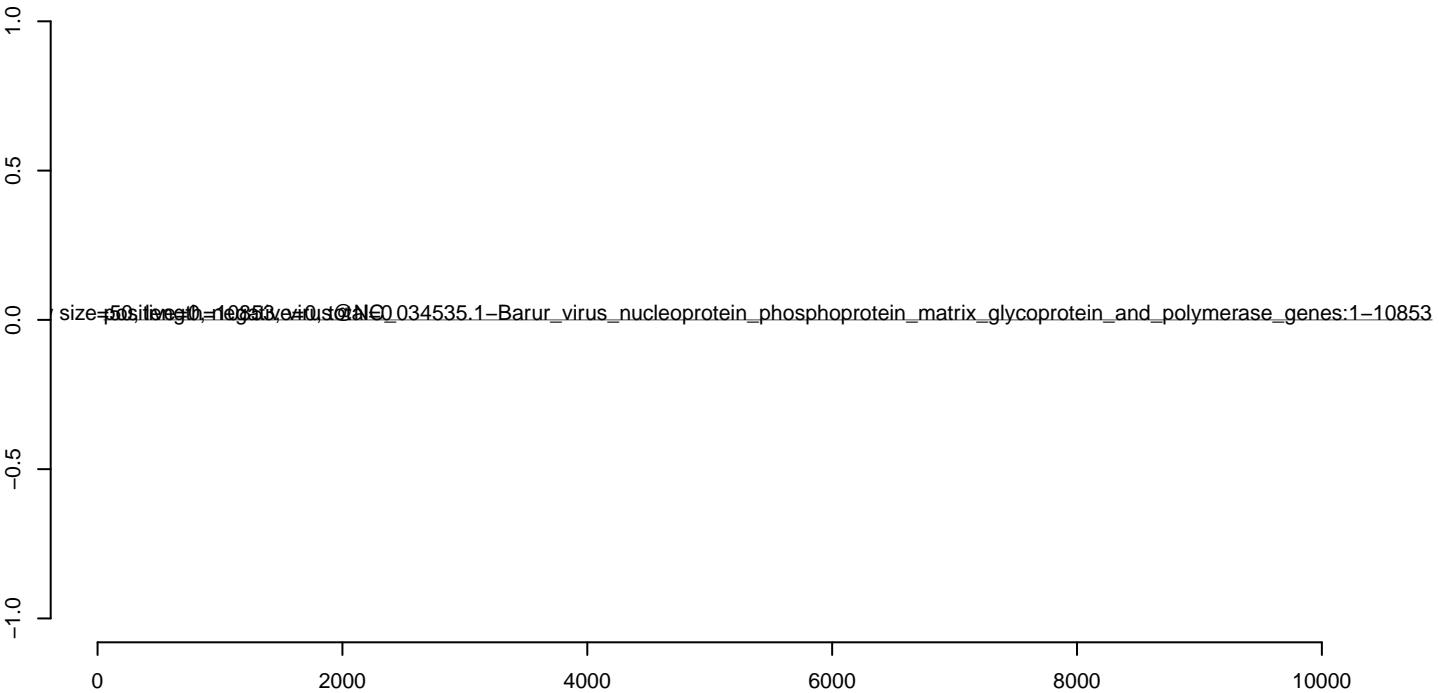
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep





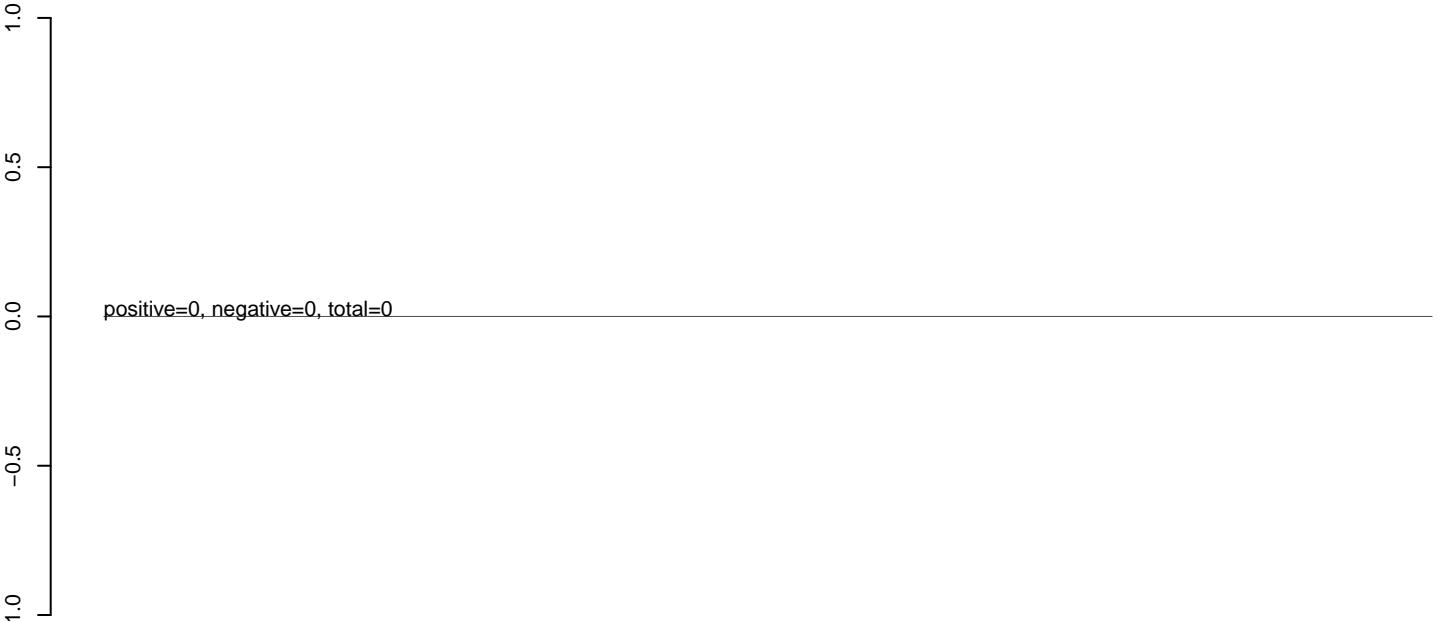
positive=0, negative=0, total=0



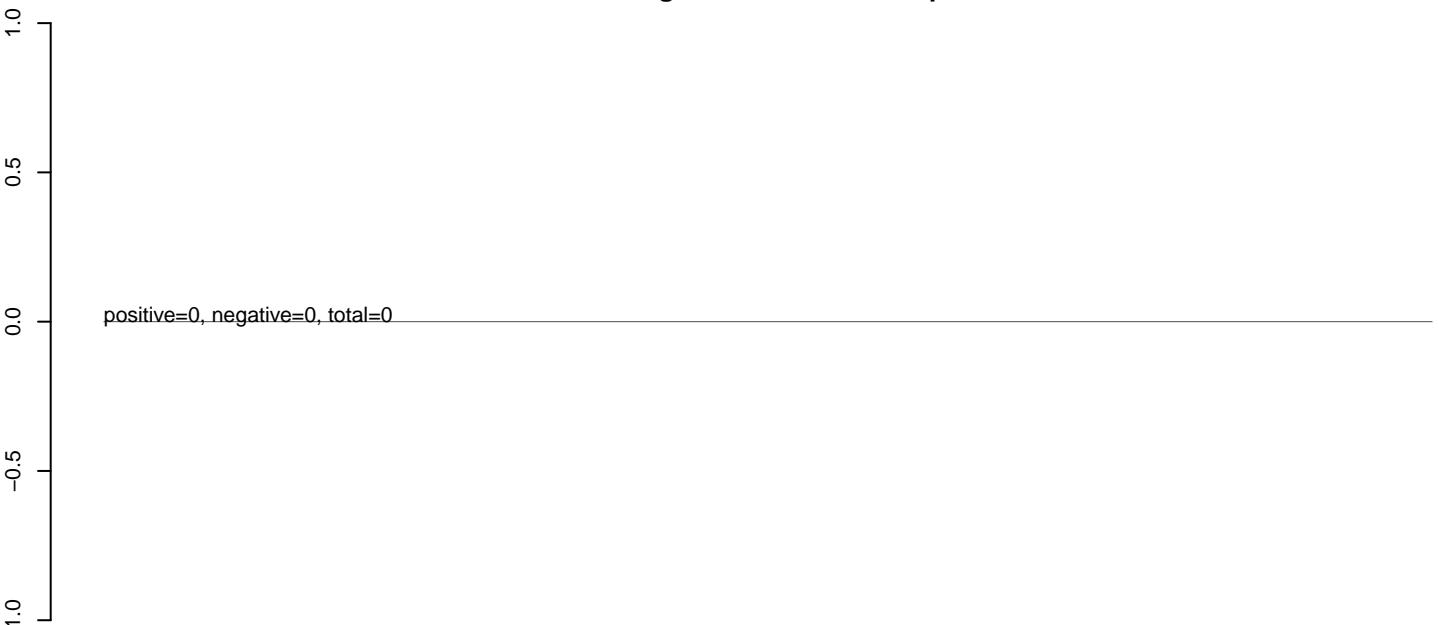
positive=0, negative=0, total=0



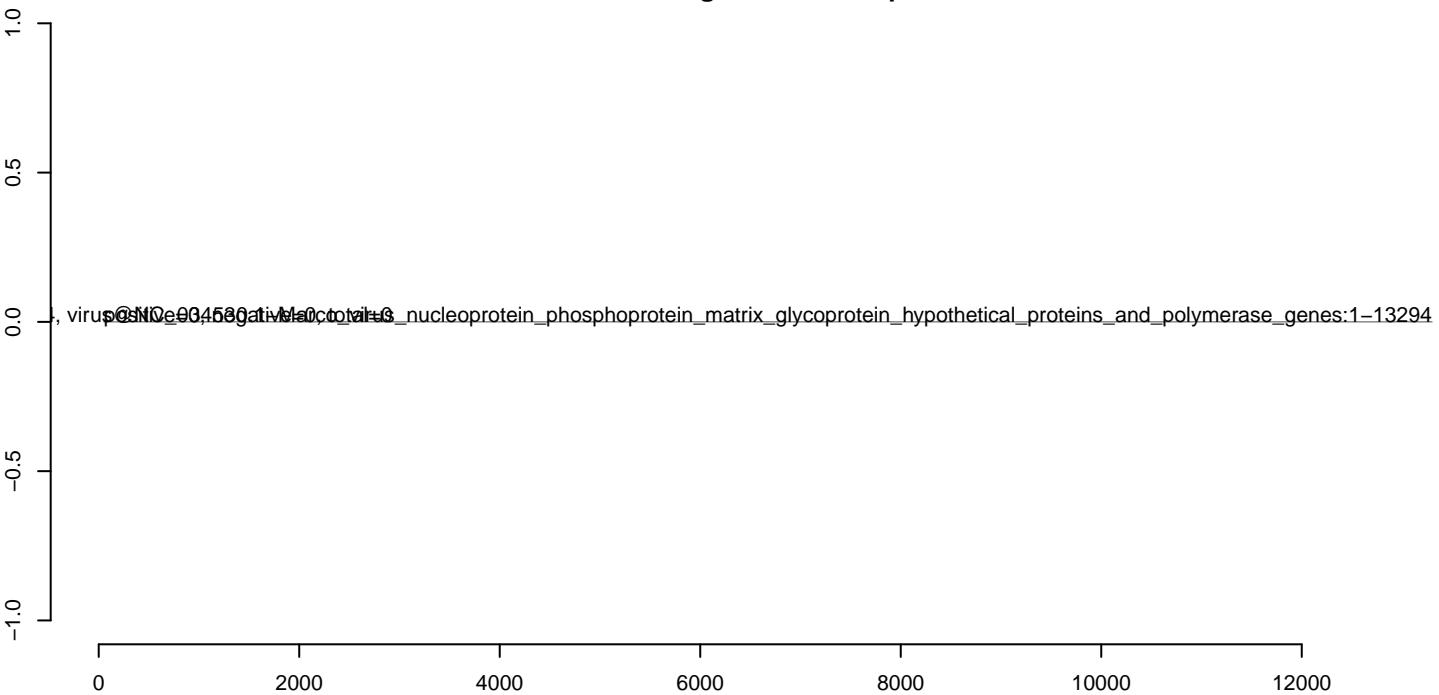
### AeAeg\_Fcanc\_GH.18\_23.rep



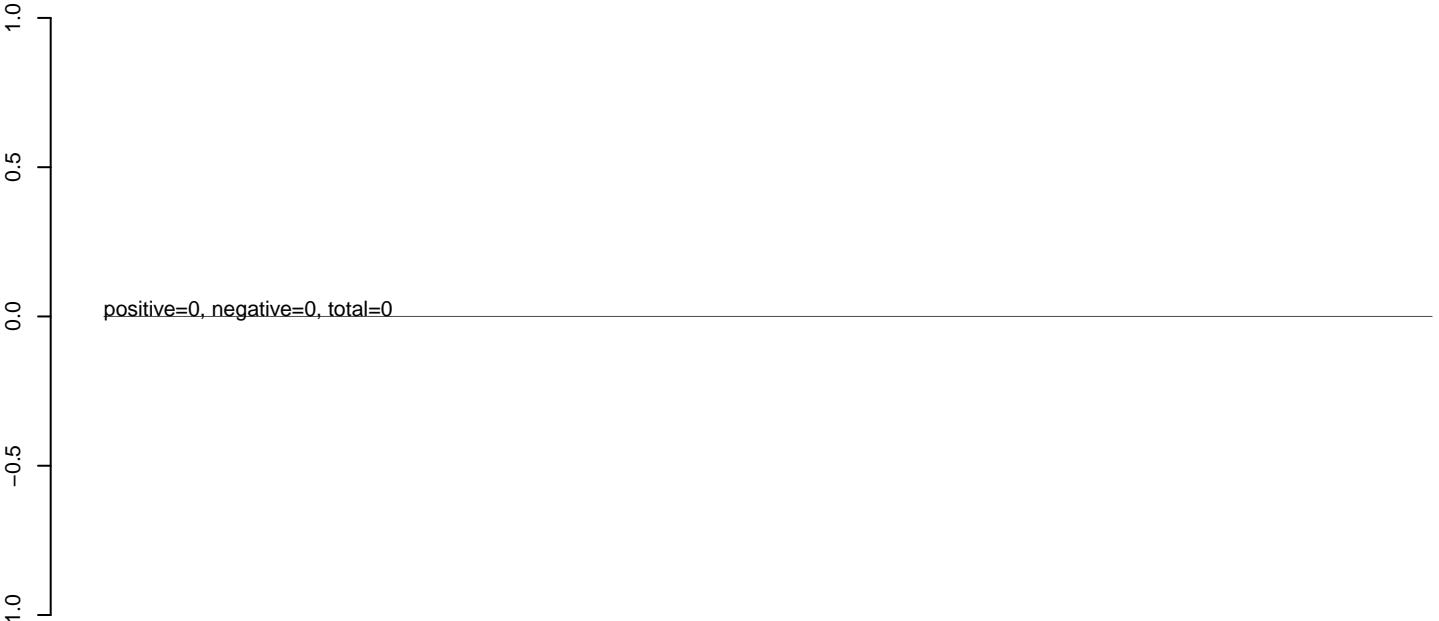
### AeAeg\_Fcanc\_GH.24\_35.rep



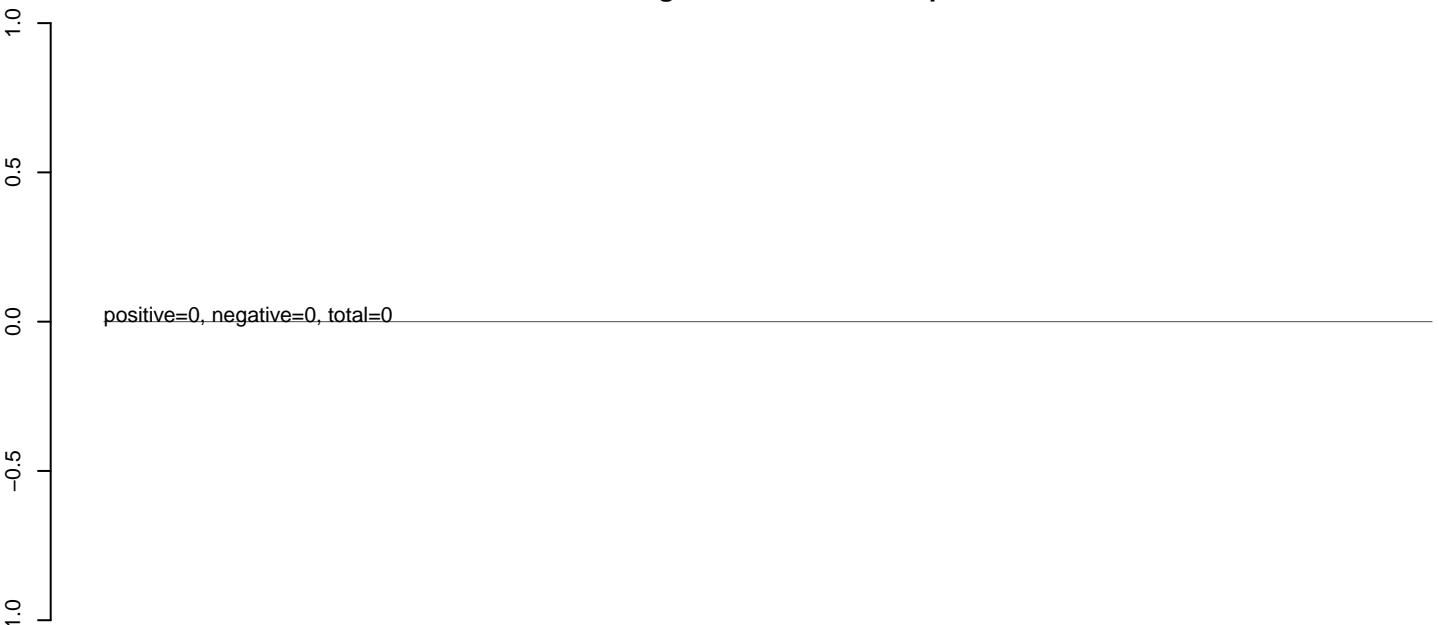
### AeAeg\_Fcanc\_GH.rep



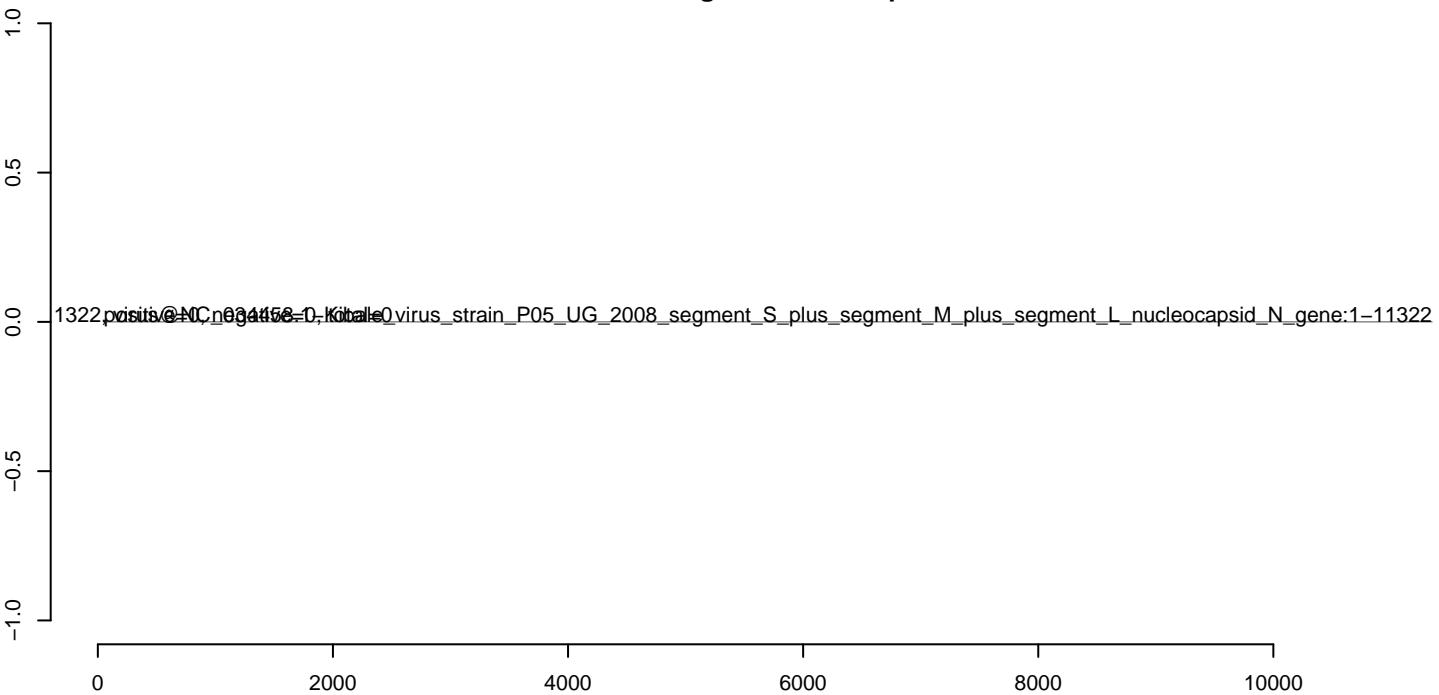
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

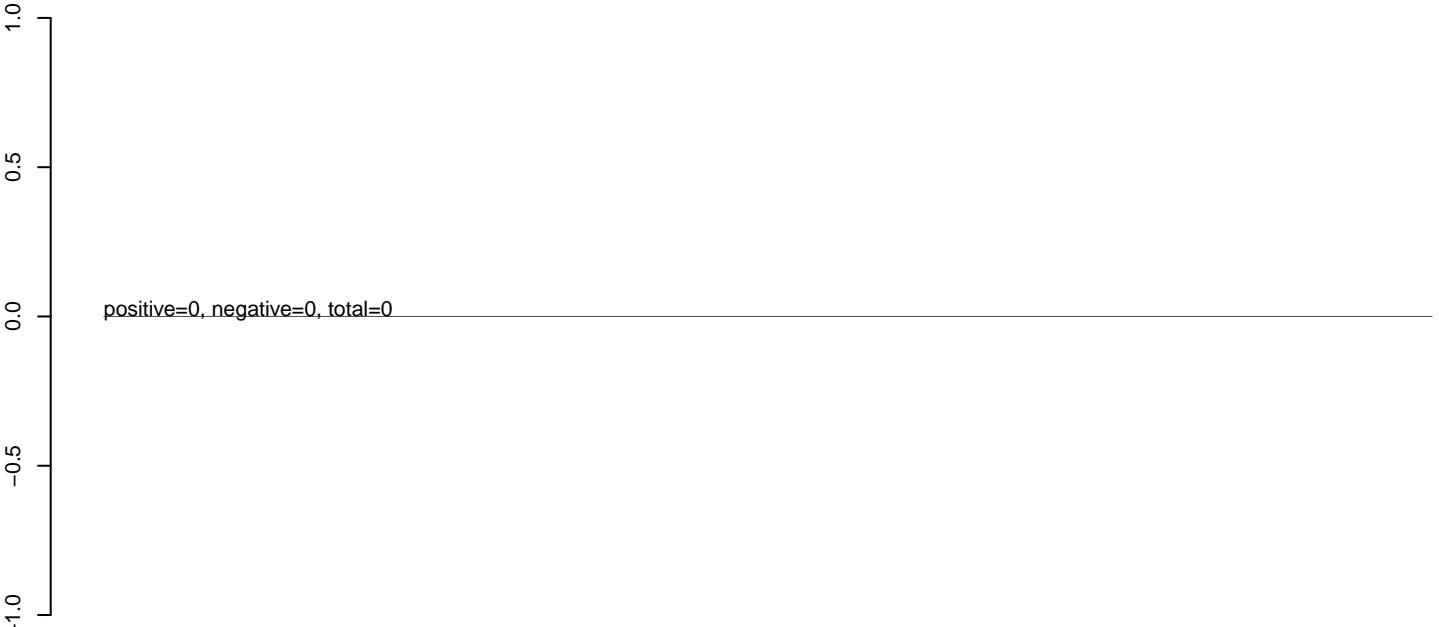
### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

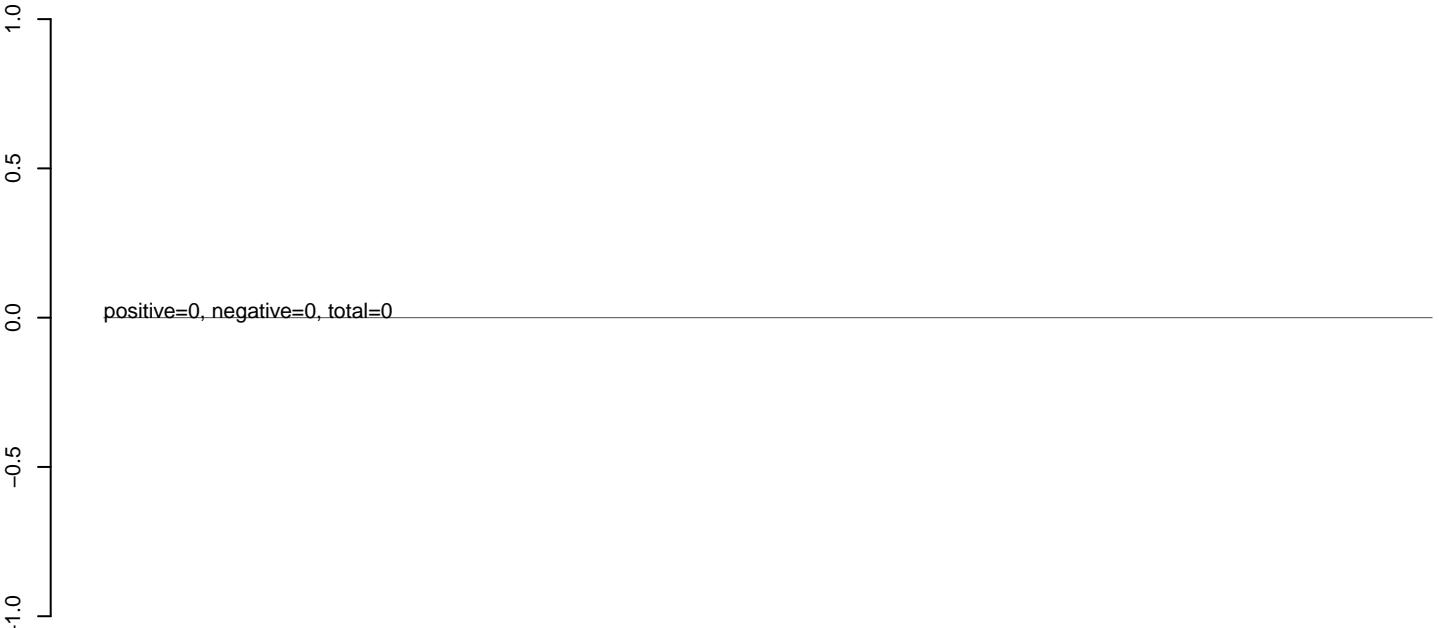
### AeAeg\_Fcanc\_GH.rep

length=11728, virus=@AeAeg, total=11728, virus\_strain=F47\_CI\_2004\_segment\_S\_plus\_segment\_M\_plus\_segment\_L\_nucleocapsid\_N\_gene:1-11728

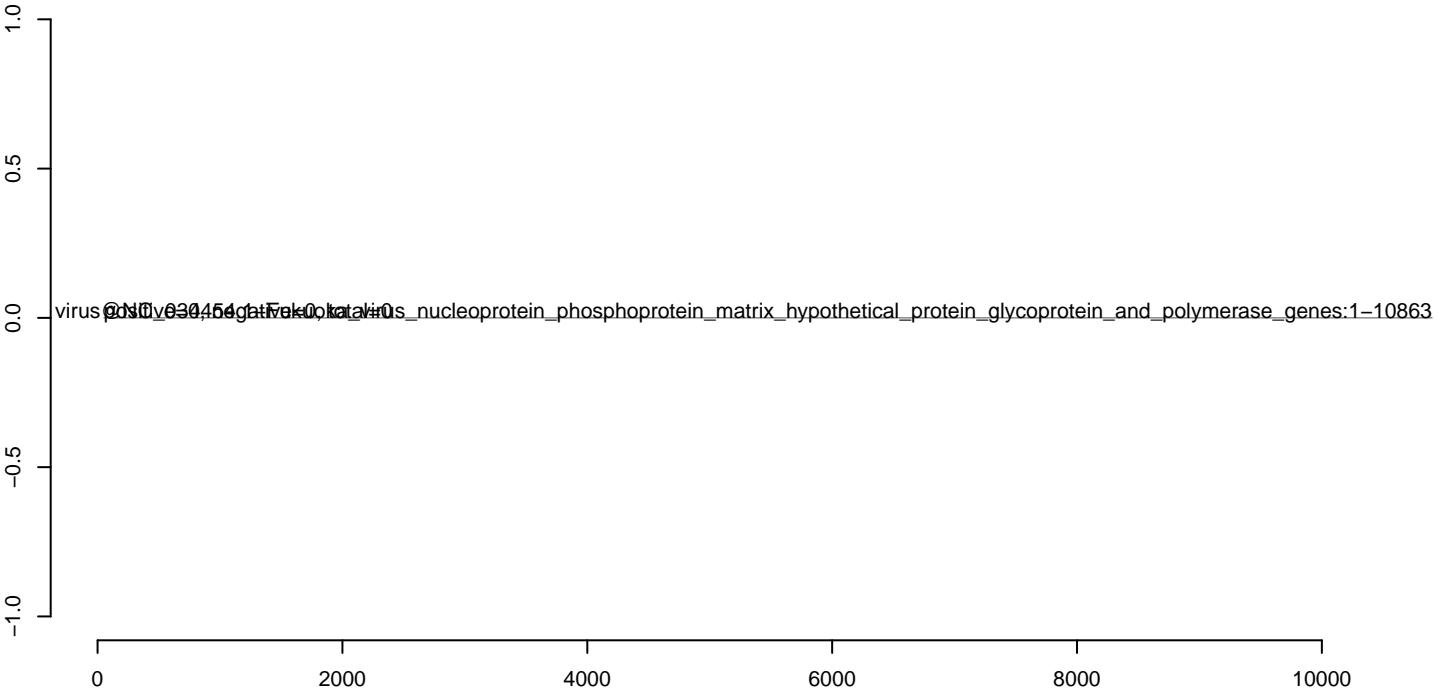
### AeAeg\_Fcanc\_GH.18\_23.rep



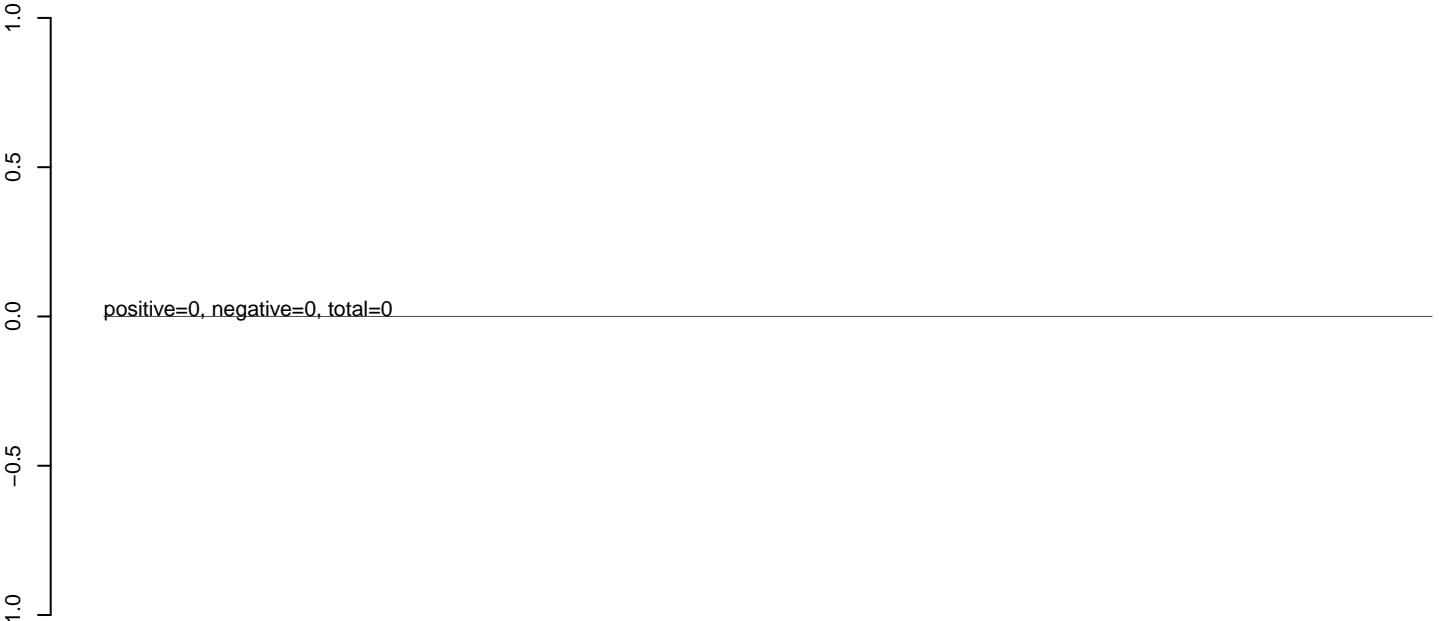
### AeAeg\_Fcanc\_GH.24\_35.rep



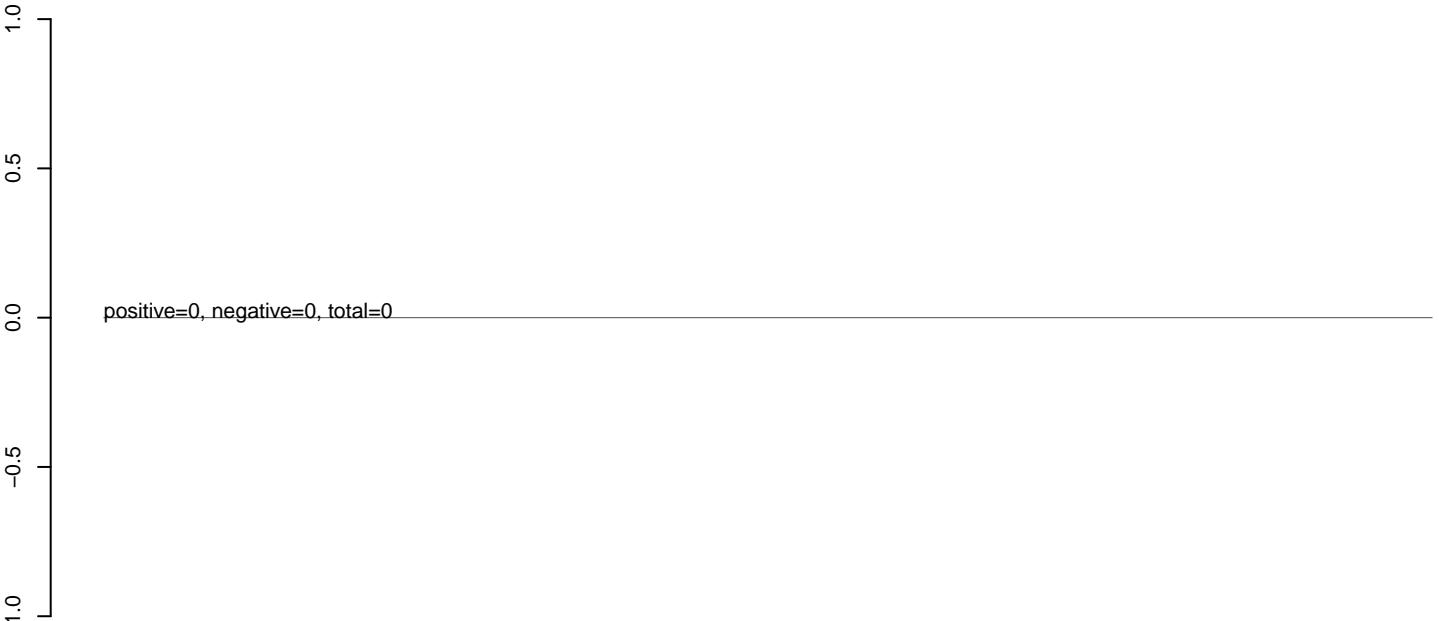
### AeAeg\_Fcanc\_GH.rep



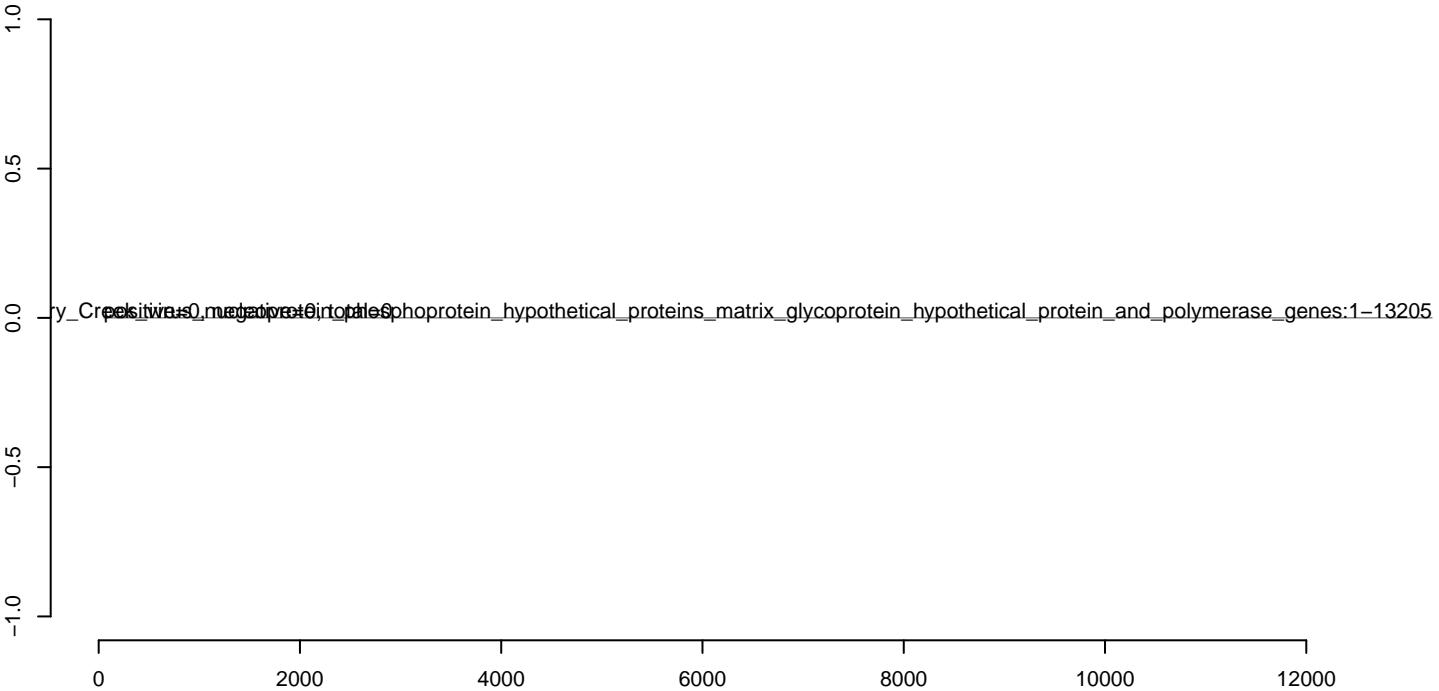
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep





positive=0, negative=0, total=0



positive=0, negative=0, total=0



positive=0 mutations in the phosphoprotein hypothetical proteins matrix glycoprotein hypothetical protein and polymerase genes:1-12957

Finals, Wrestling, Mathematics, Updates

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

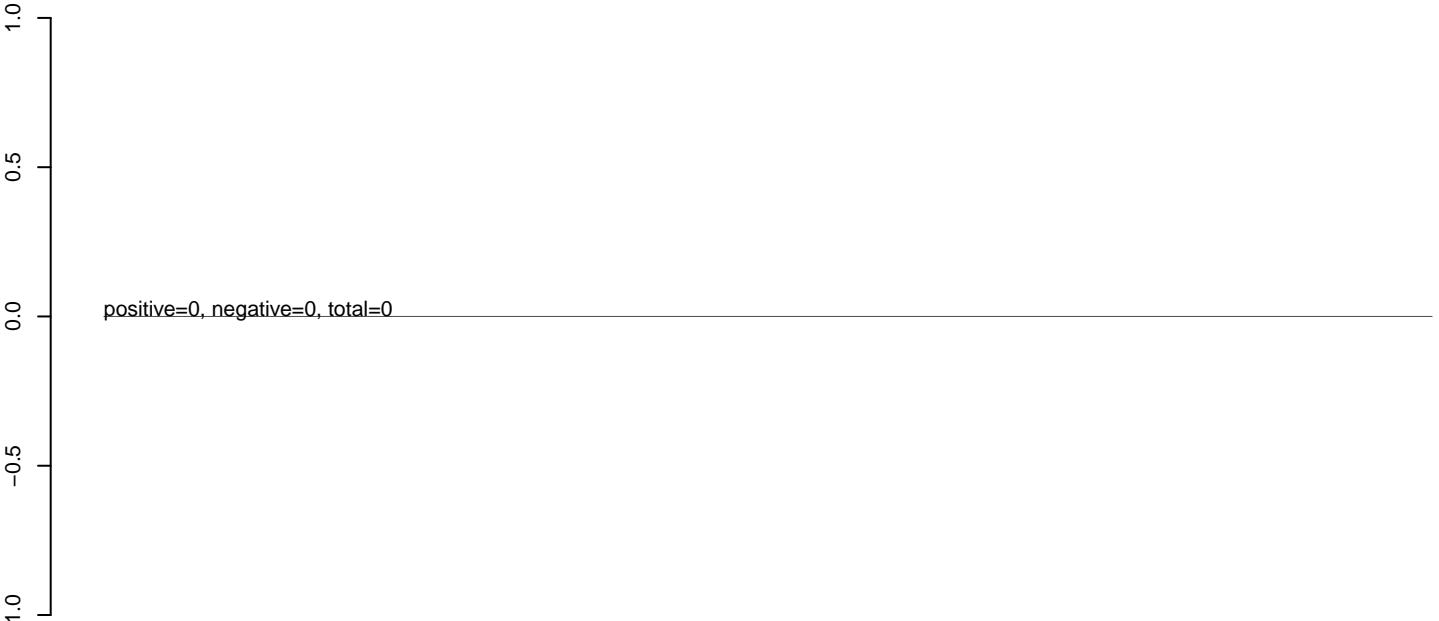
### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

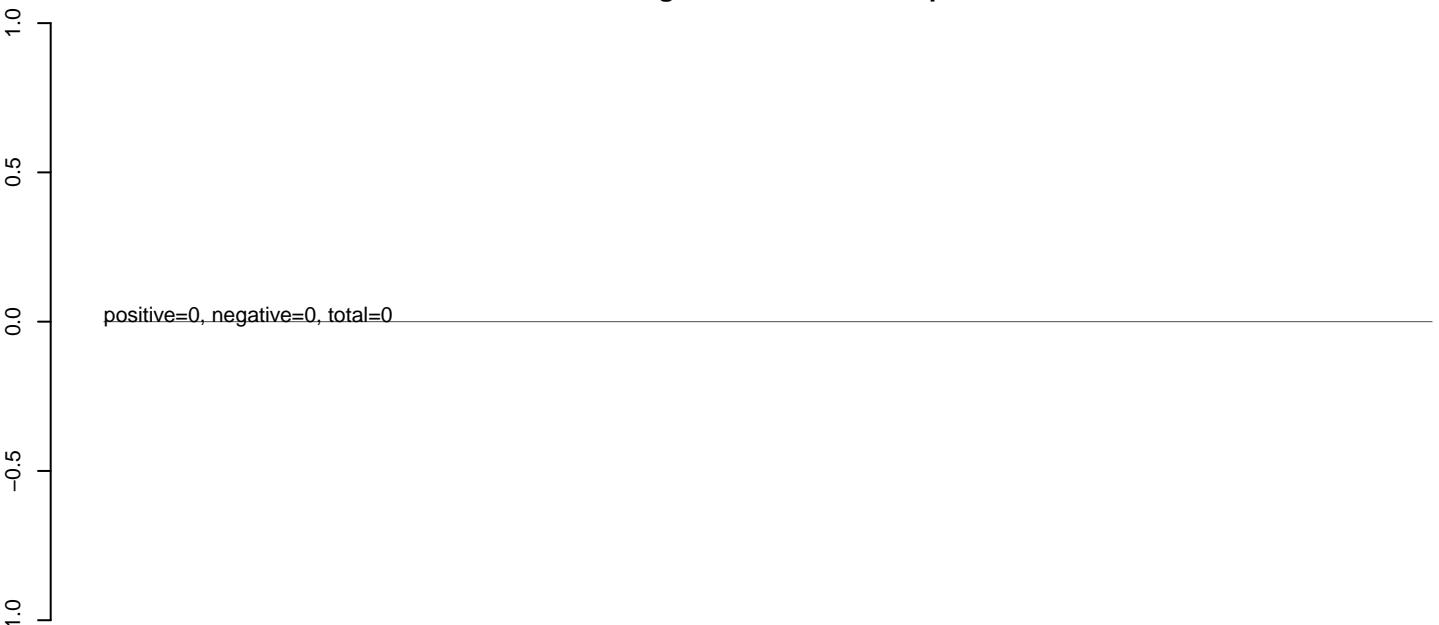
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, window\_size=50, length=10293, virus@NC\_034018.1-Yaounde\_virus\_strain\_Dak\_Ar\_Y276\_polyprotein\_gene:1-10293

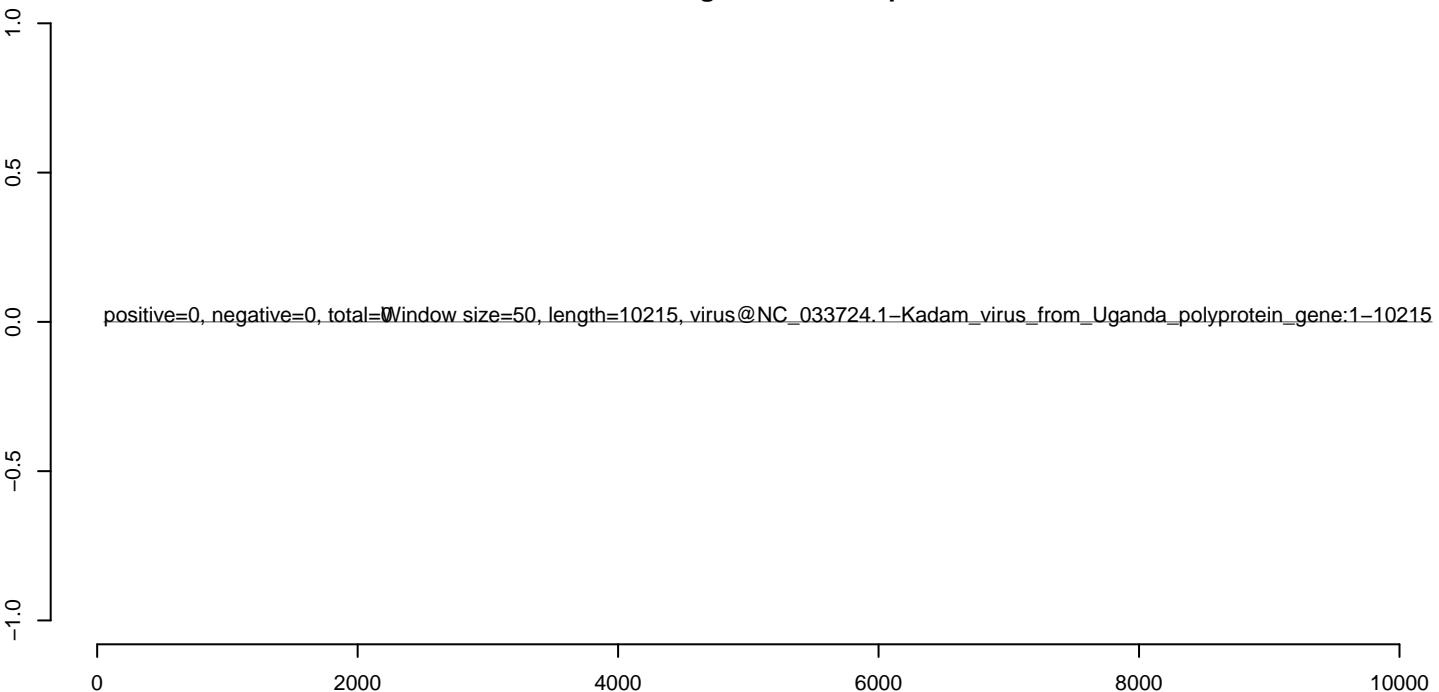
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

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

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0, window size=50, length=10173, virus@NC\_033699.1-Jugra\_virus\_strain\_P-9-314\_polyprotein\_gene:1-10173

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10182, virus@NC\_033698.1-Uganda\_S\_virus\_polyprotein\_gene:1-10182

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0, size=50, length=10173, virus@NC\_033697.1-Saboya\_virus\_strain\_Dak\_AR\_D4600\_polyprotein\_gene:1-10173

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0, size=50, length=10173, virus@NC\_033693.1-Bouboui\_virus\_strain\_DAK\_AR\_B490\_polyprotein\_gene:1-10173



positive=0, negative=0, total=0



positive=0, negative=0, total=0



Open Access  CC-BY  M26608 RNA-dependent RNA polymerase, putative glycoprotein and hypothetical protein genes; 1\_11280

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0, window size=50, length=10869, virus@NC\_031957.1-UNVERIFIED\_Coot\_Bay\_virus\_strain\_EVG5-53:1-10869

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

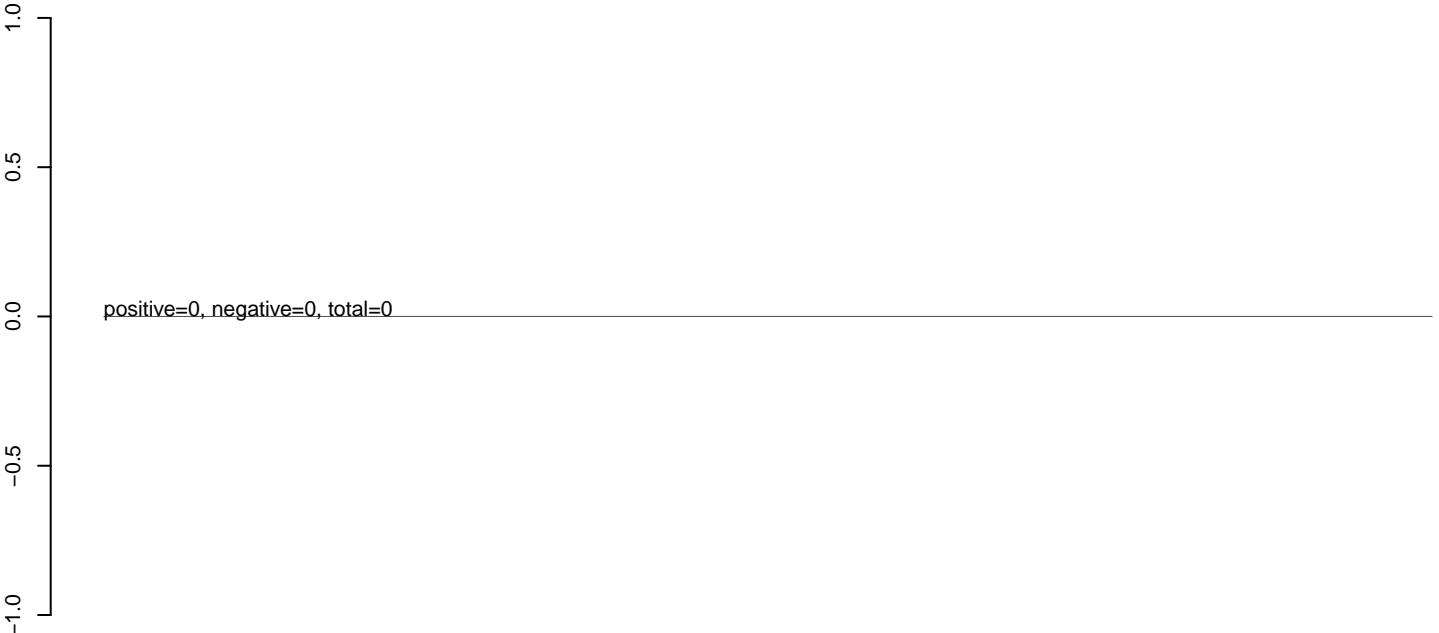
### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

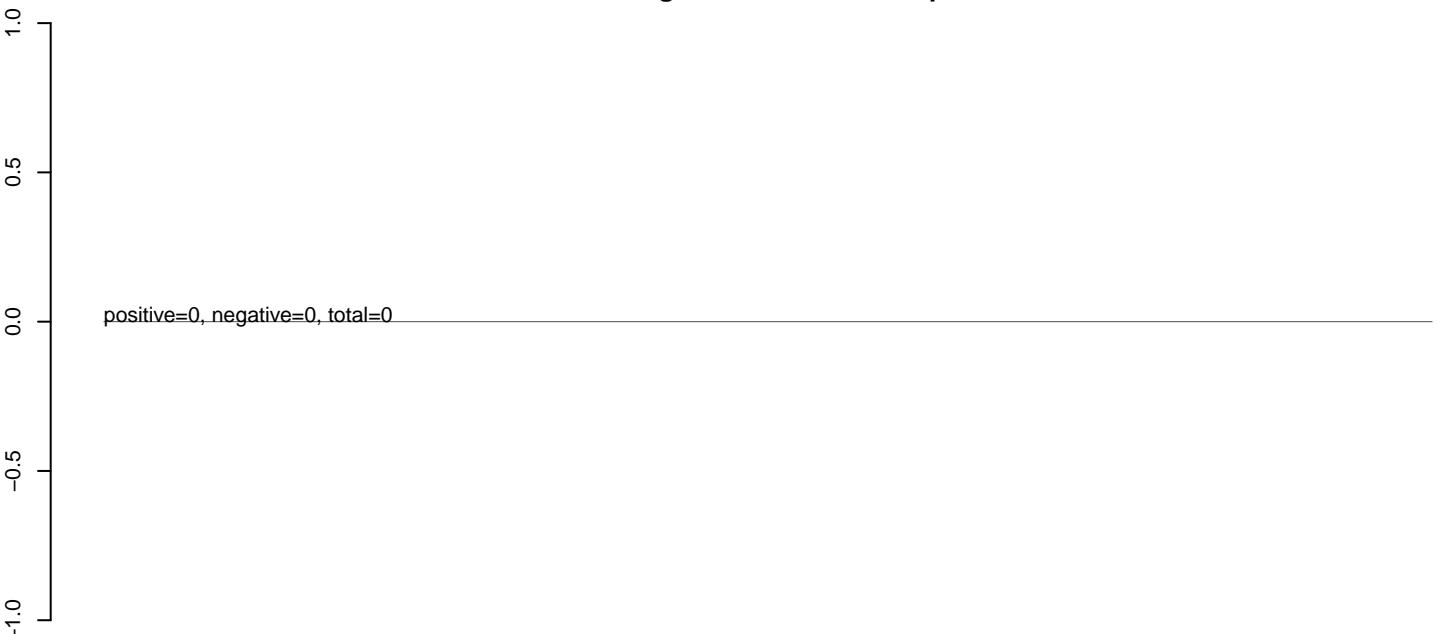
### AeAeg\_Fcanc\_GH.rep

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

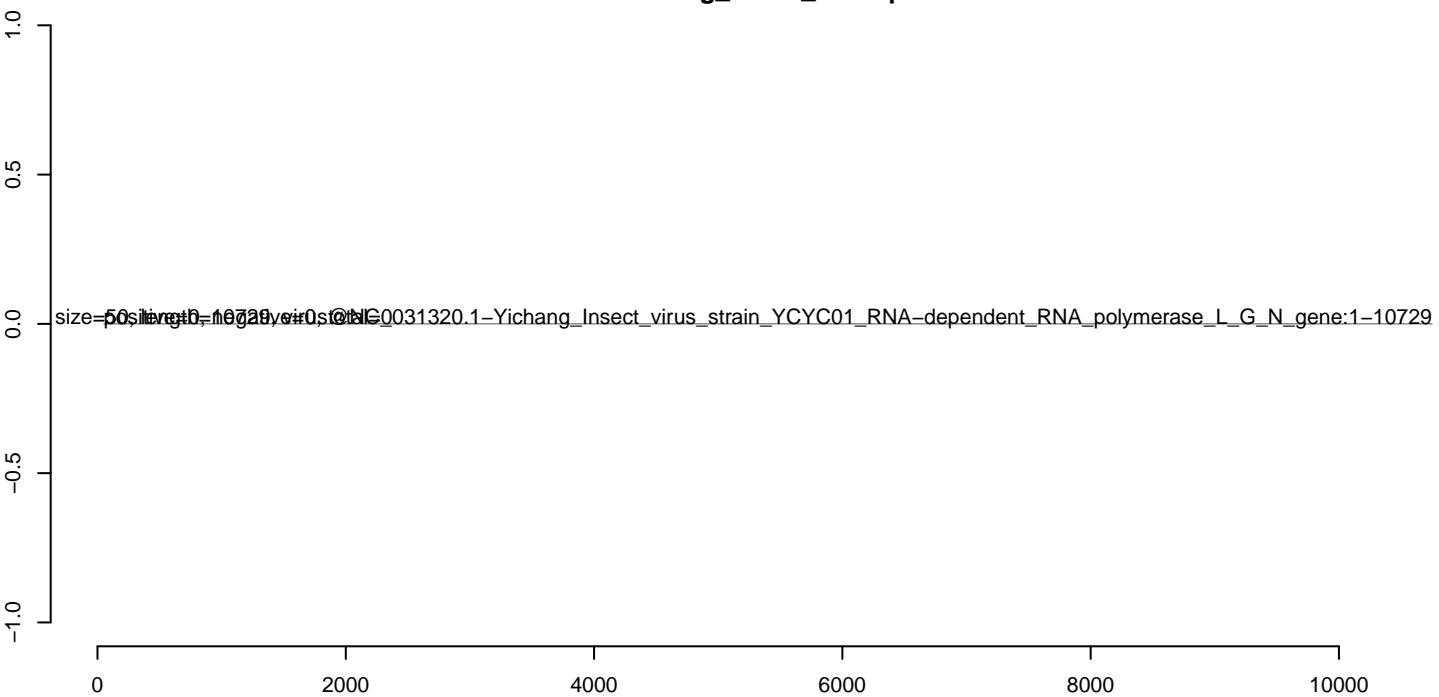
### AeAeg\_Fcanc\_GH.18\_23.rep



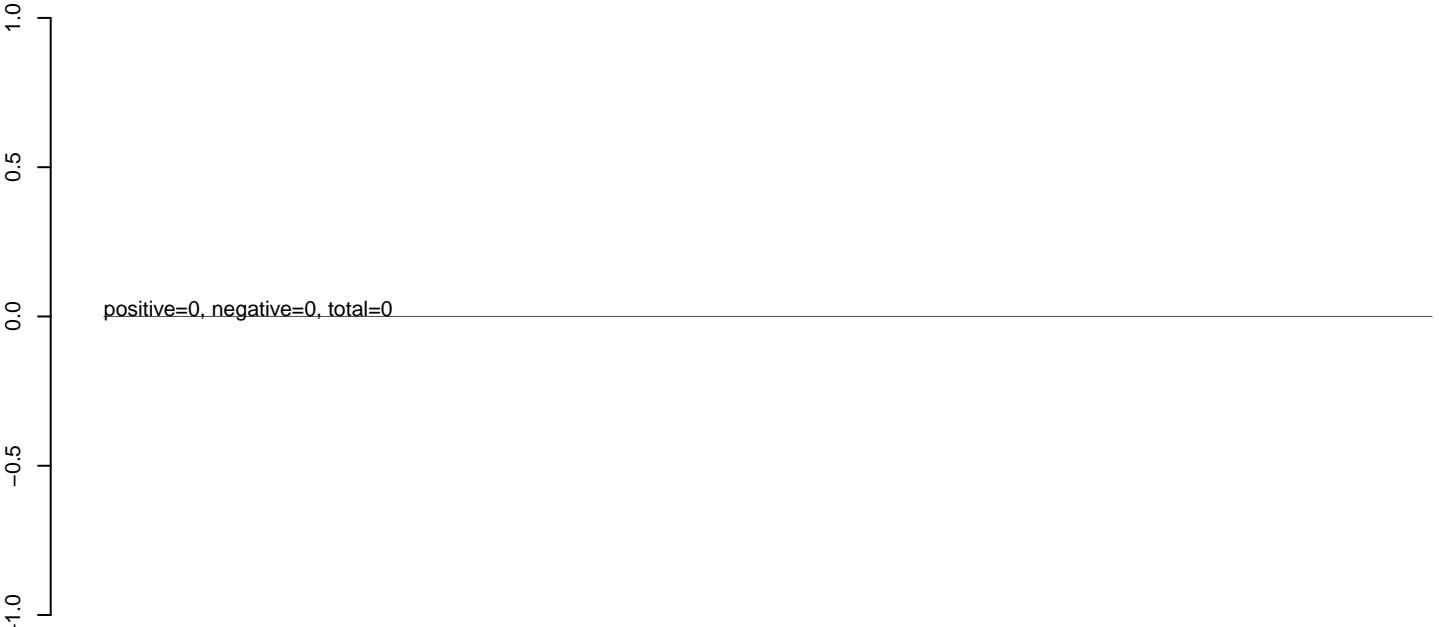
### AeAeg\_Fcanc\_GH.24\_35.rep



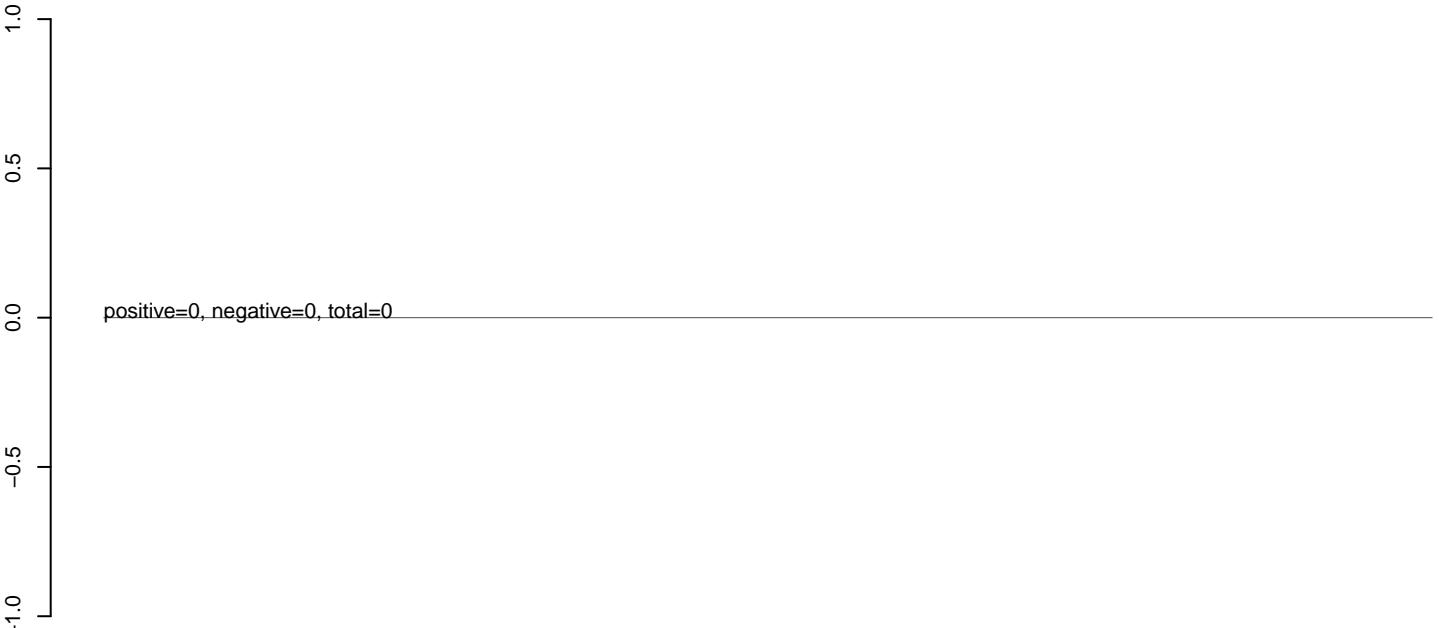
### AeAeg\_Fcanc\_GH.rep



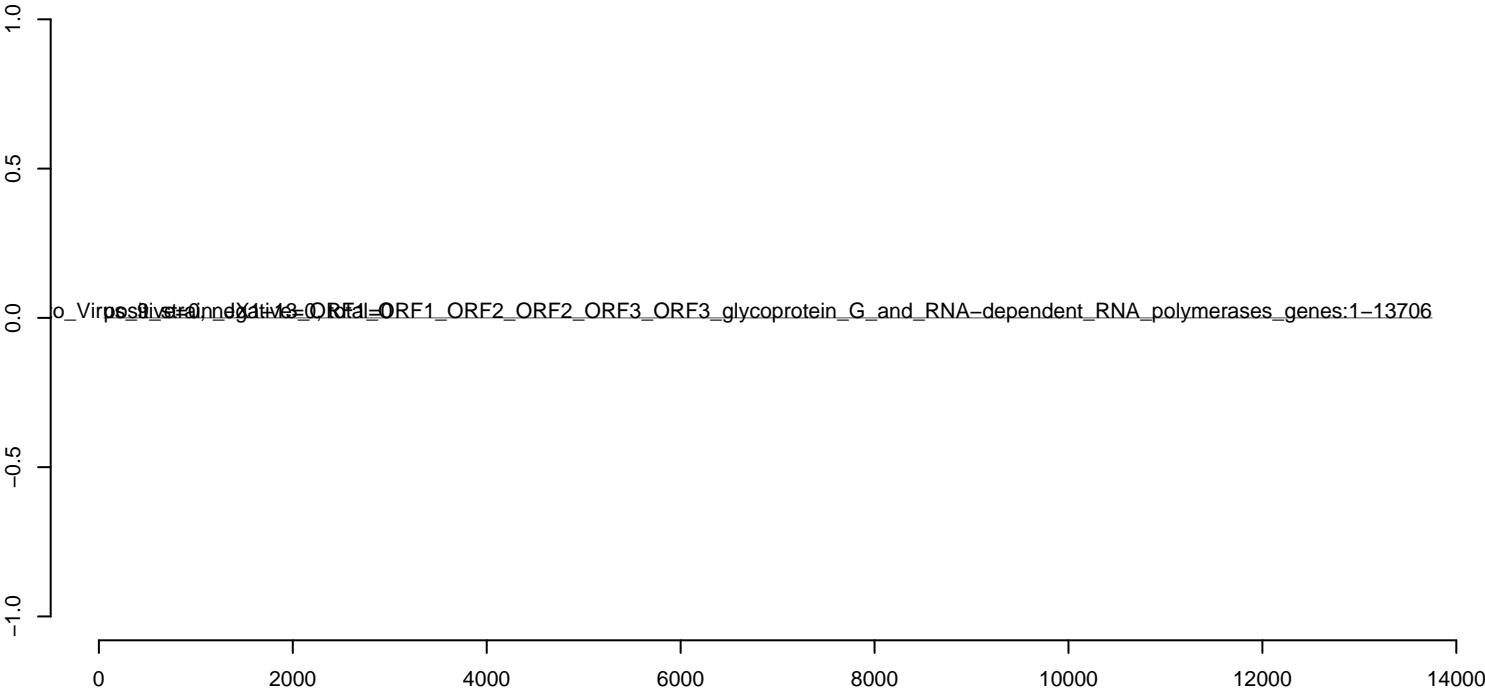
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10158, virus@NC\_030401.1-Hanko\_virus\_polyprotein\_gene:1-10158

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

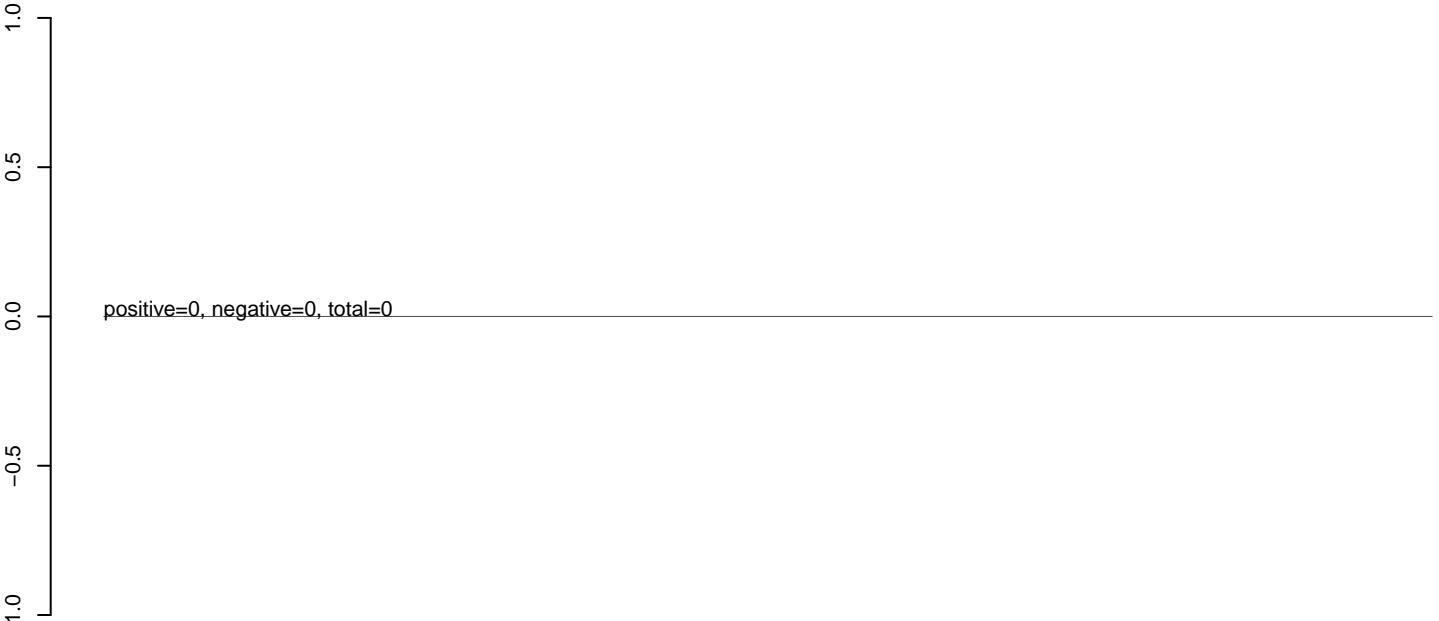
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

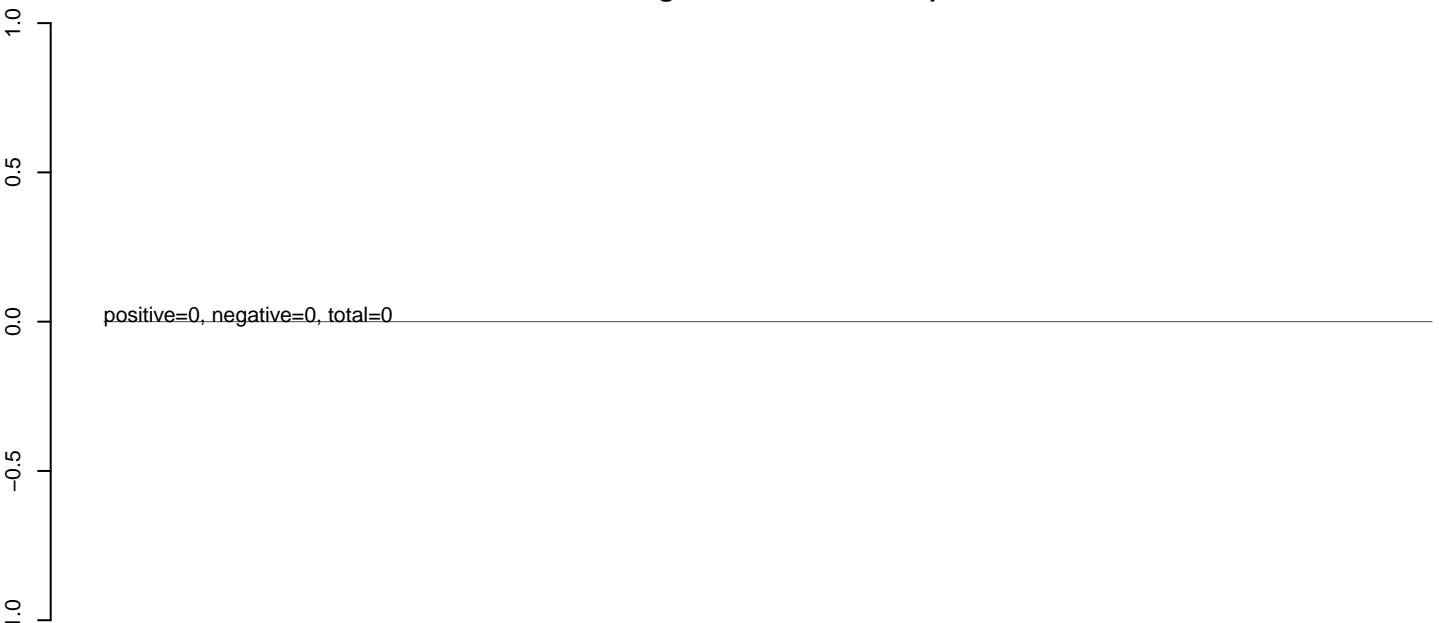
Window size=50, length=10864, virus@NC\_030290.1-Culiseta\_flavivirus\_strain\_502-13:1-10864

0 2000 4000 6000 8000 10000

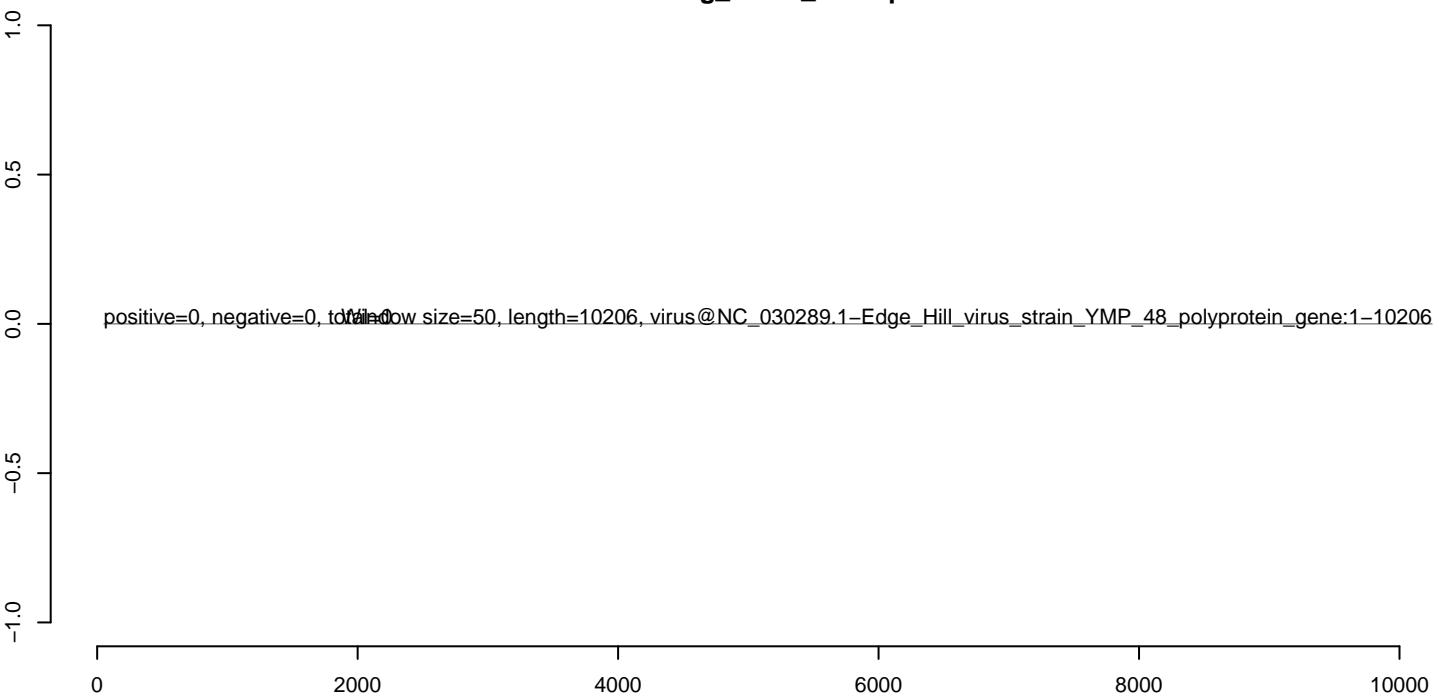
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=14478, virus@NC\_028241.1-Yata\_virus\_isolate\_DakArB\_2181:1-14478

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=15444, virus@NC\_025400.1-Malakal\_virus\_isolate\_SudAr\_1169-64:1-15444

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=11220, virus@NC\_025399.1-Oak-Vale\_virus\_strain\_CSIRO\_1342:1-11220

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

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

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0 size=50, length=11190, virus@NC\_025384.1-Culex\_tritaeniorhynchus\_rhabdovirus\_RNA\_strain\_TY:1-11190

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.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

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

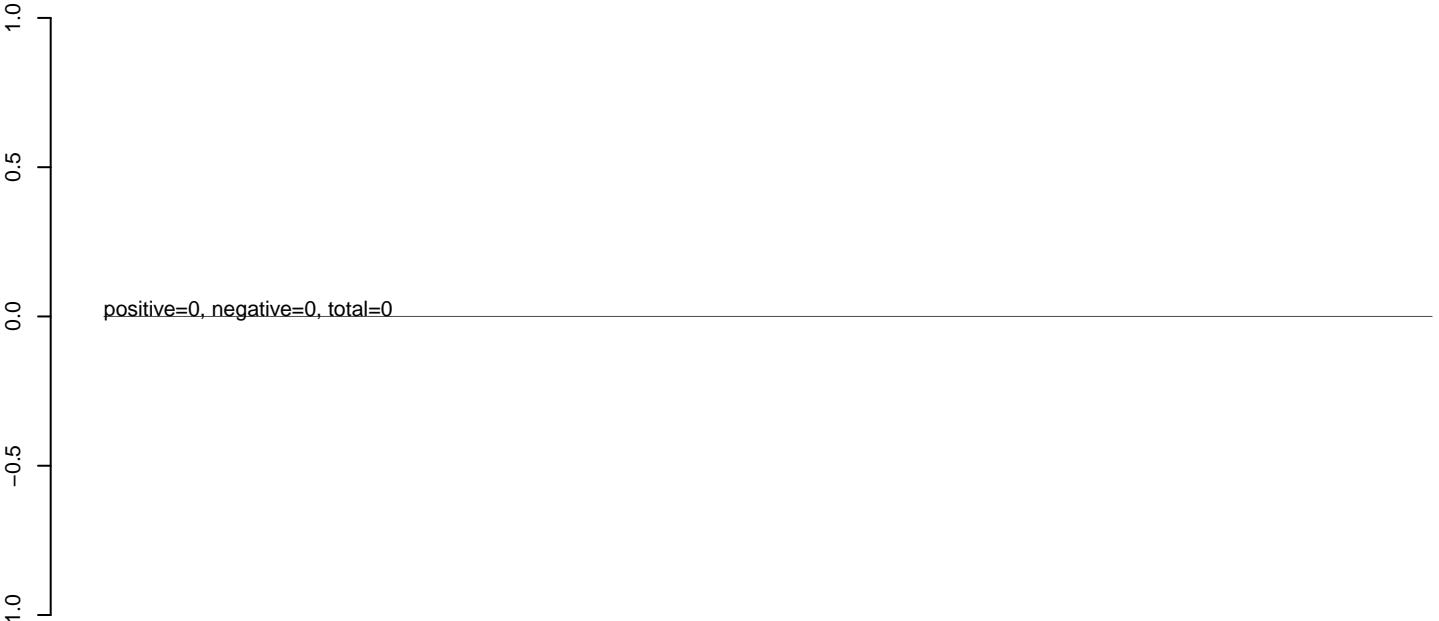
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

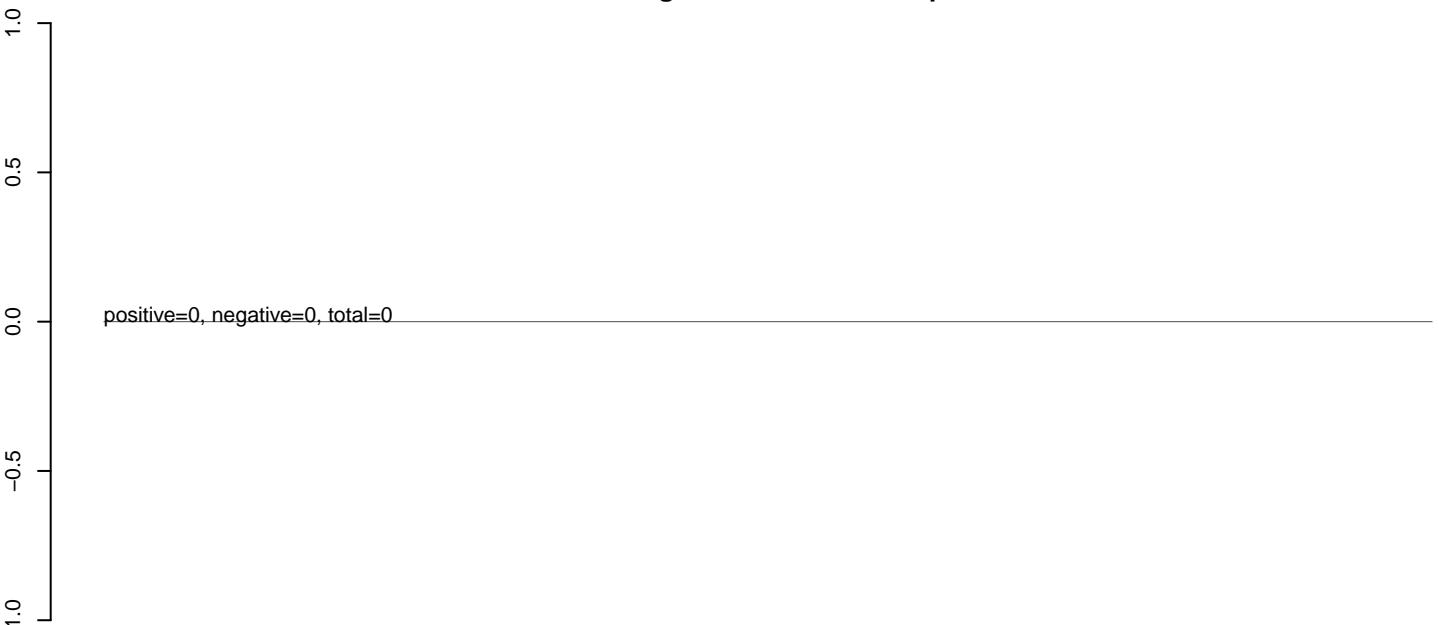
Window size=50, length=11097, virus@NC\_025356.1–Pike\_fry\_rhabdovirus\_isolate\_F4:1–11097

0 2000 4000 6000 8000 10000

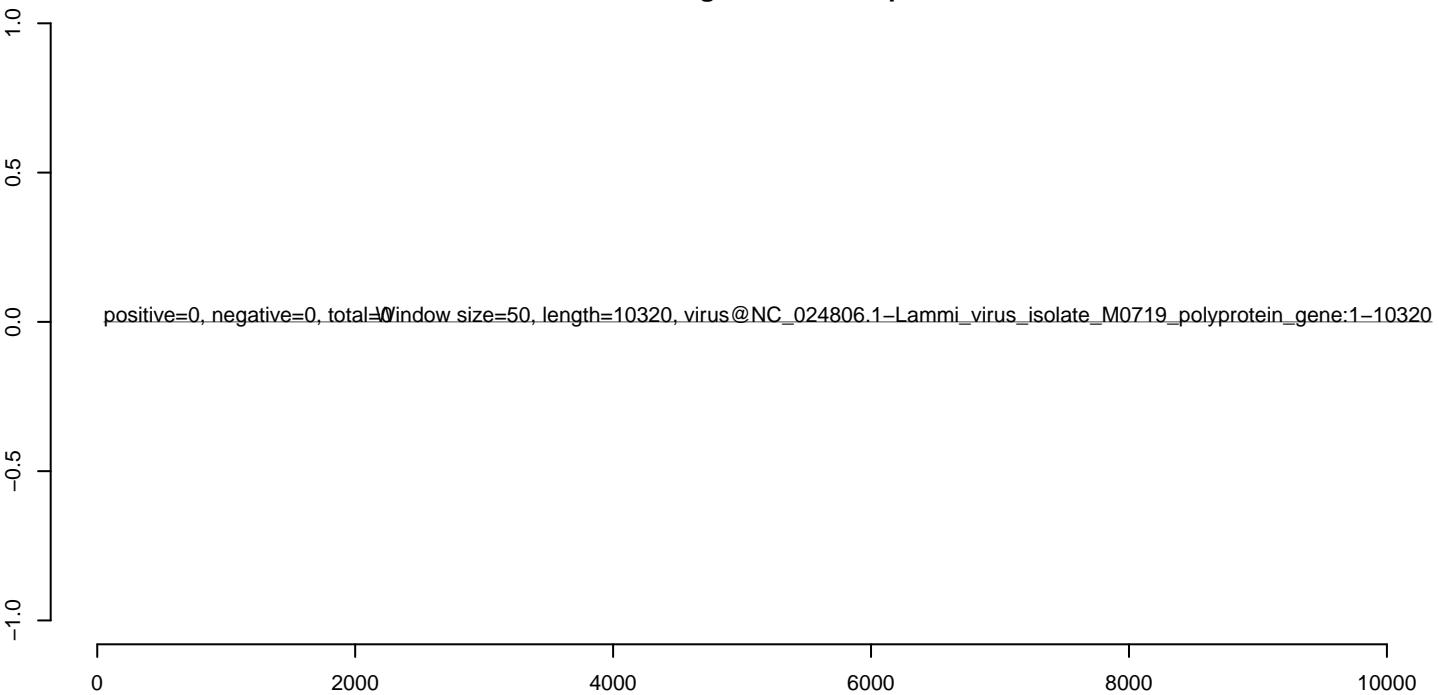
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

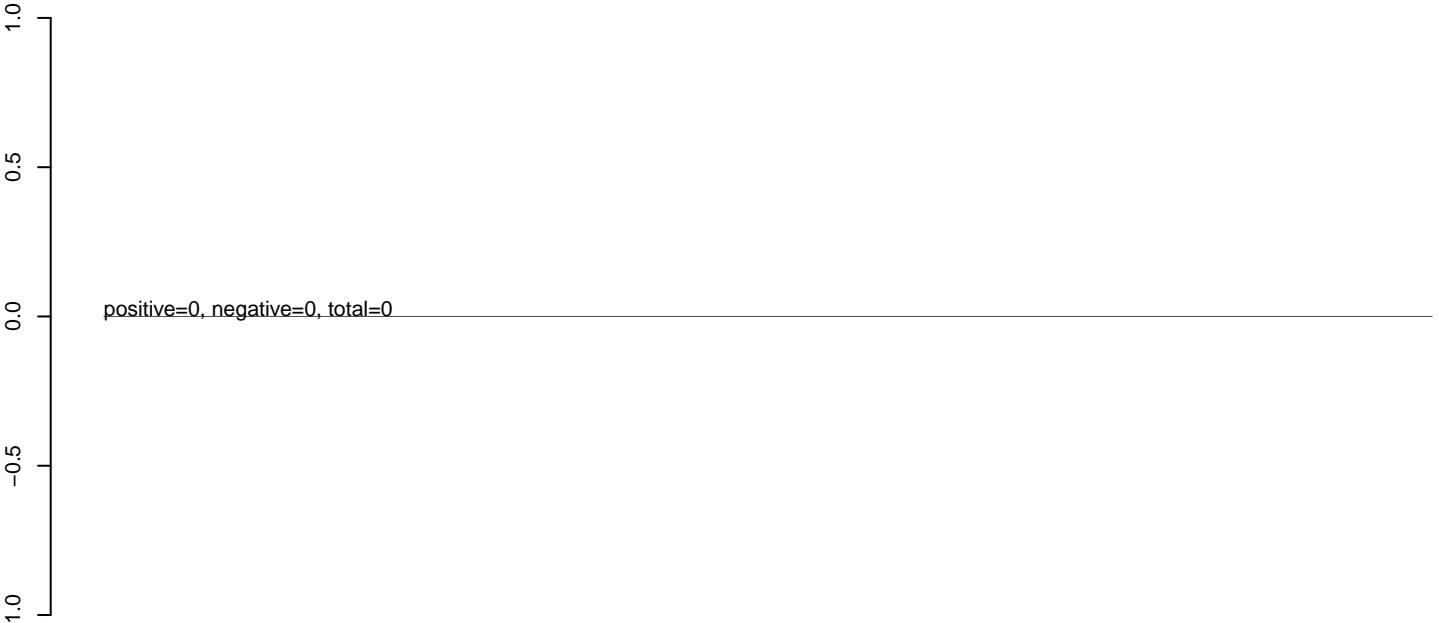
positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

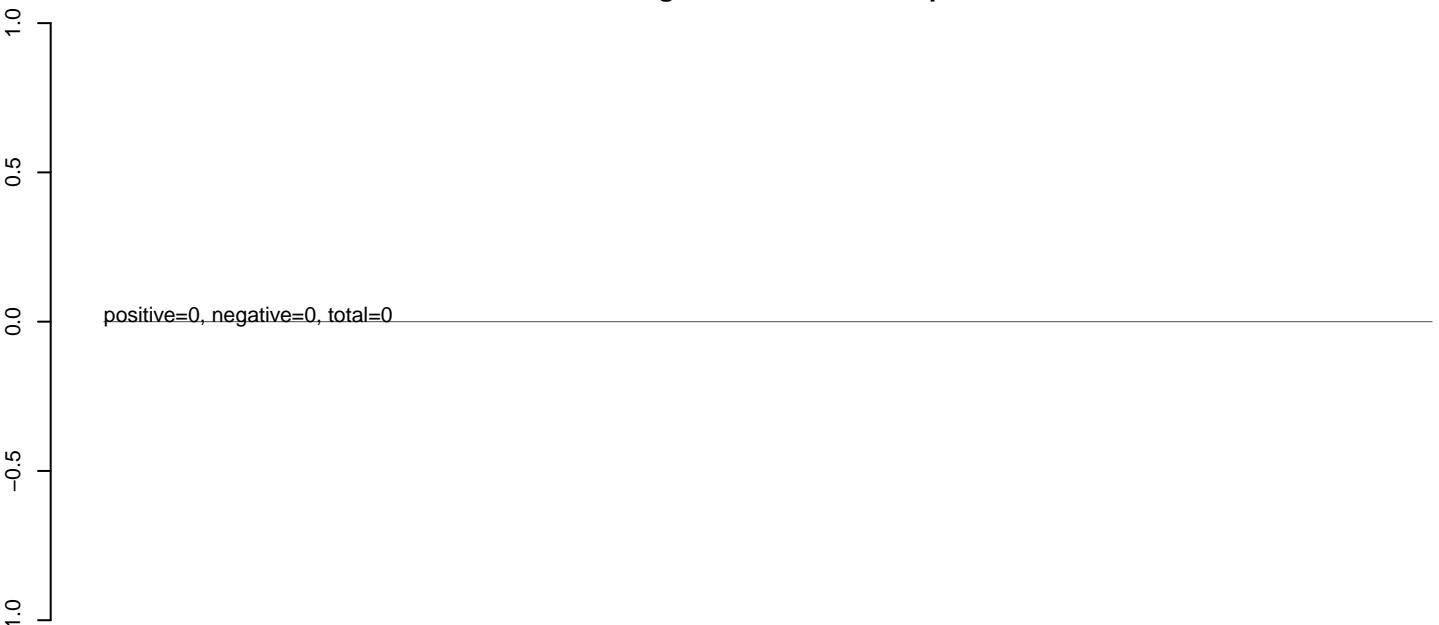
positive=0, negative=0, total=140 window size=50, length=10365, virus@NC\_024805.1–Ijomantsi\_virus\_isolate\_M0724\_polyprotein\_gene:1–10365

0 2000 4000 6000 8000 10000

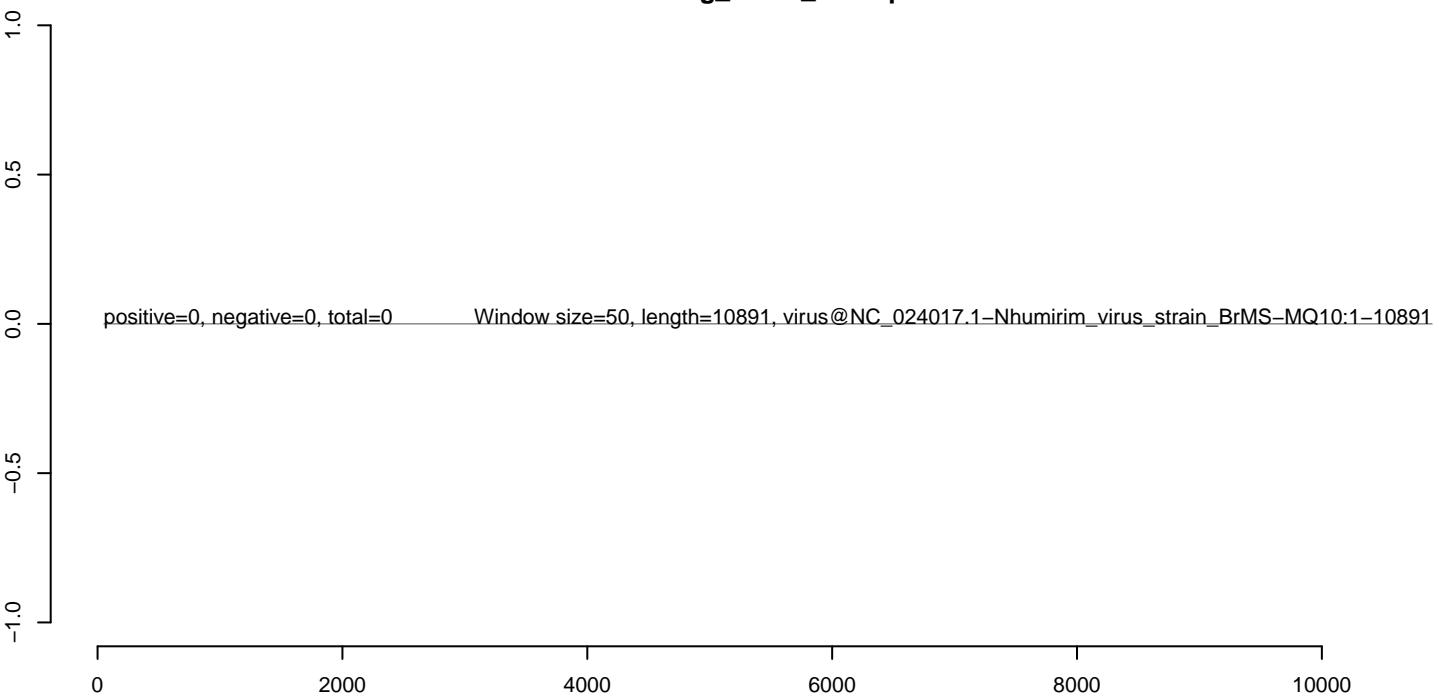
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

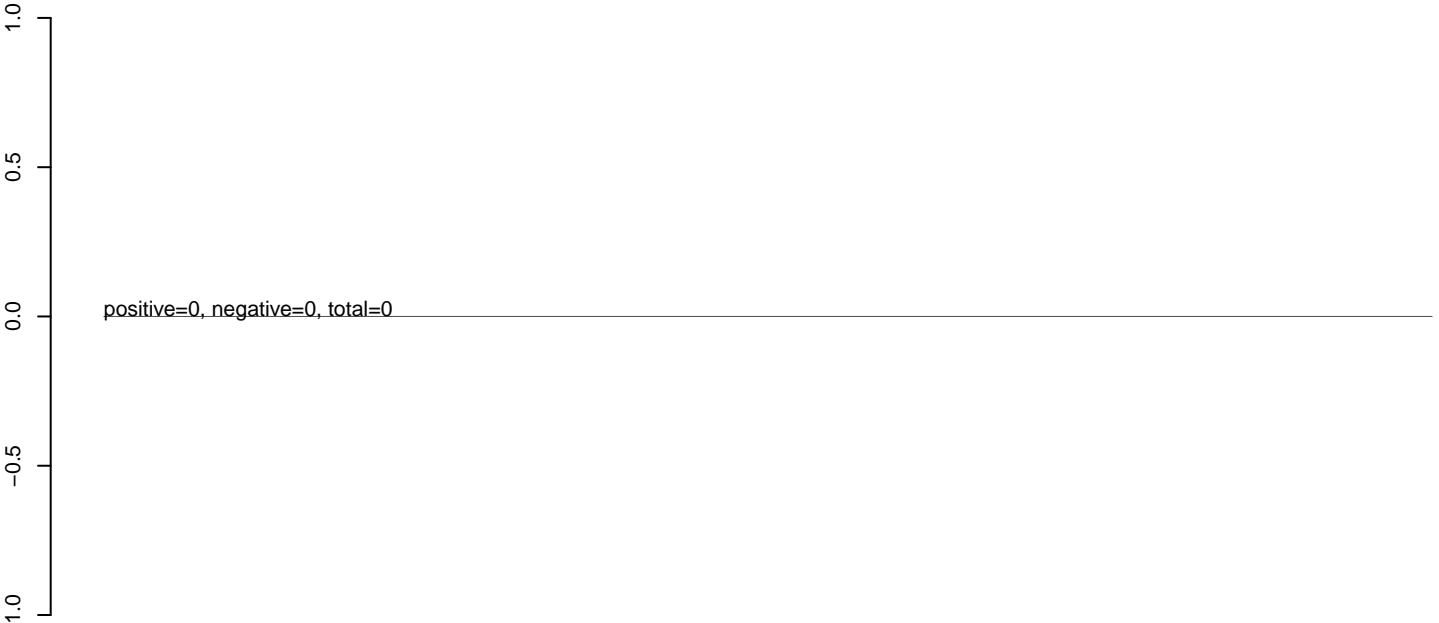
positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

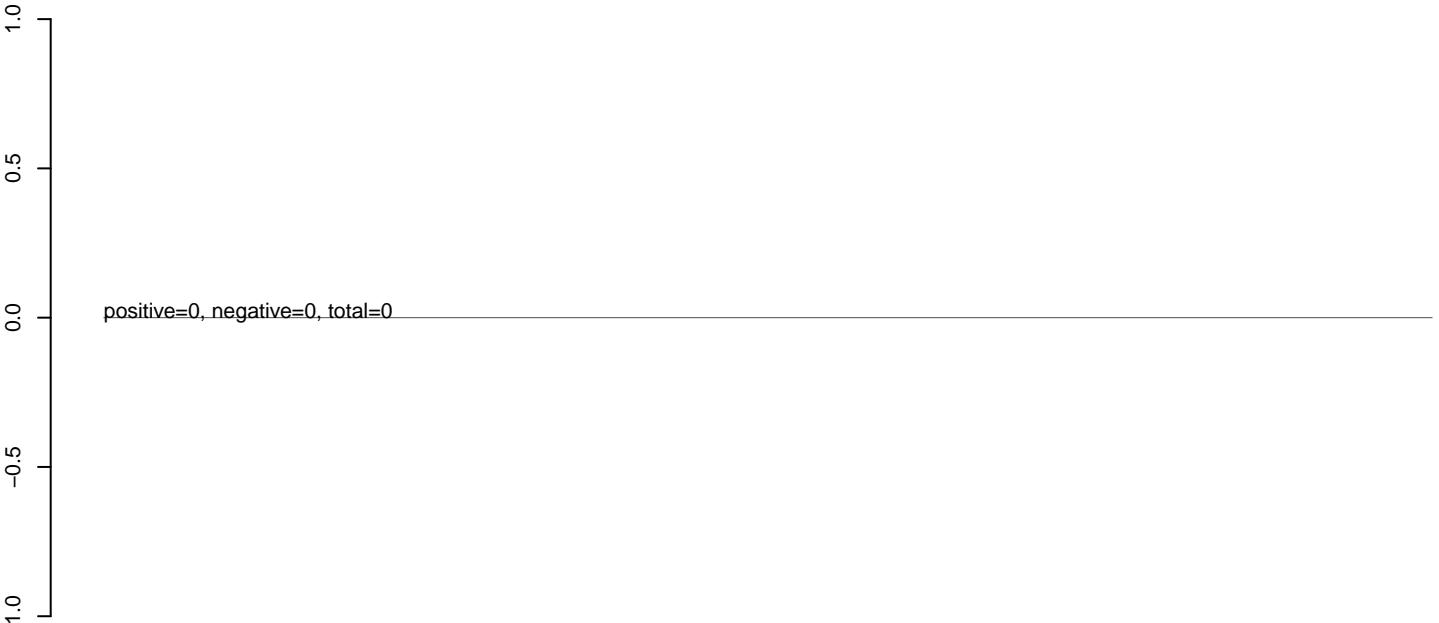
positive=0, negative=0, total=0 size=50, length=10688, virus@NC\_023439.1-Kama\_virus\_strain\_LEIV-20776Tat\_polyprotein\_gene:1-10688

0 2000 4000 6000 8000 10000

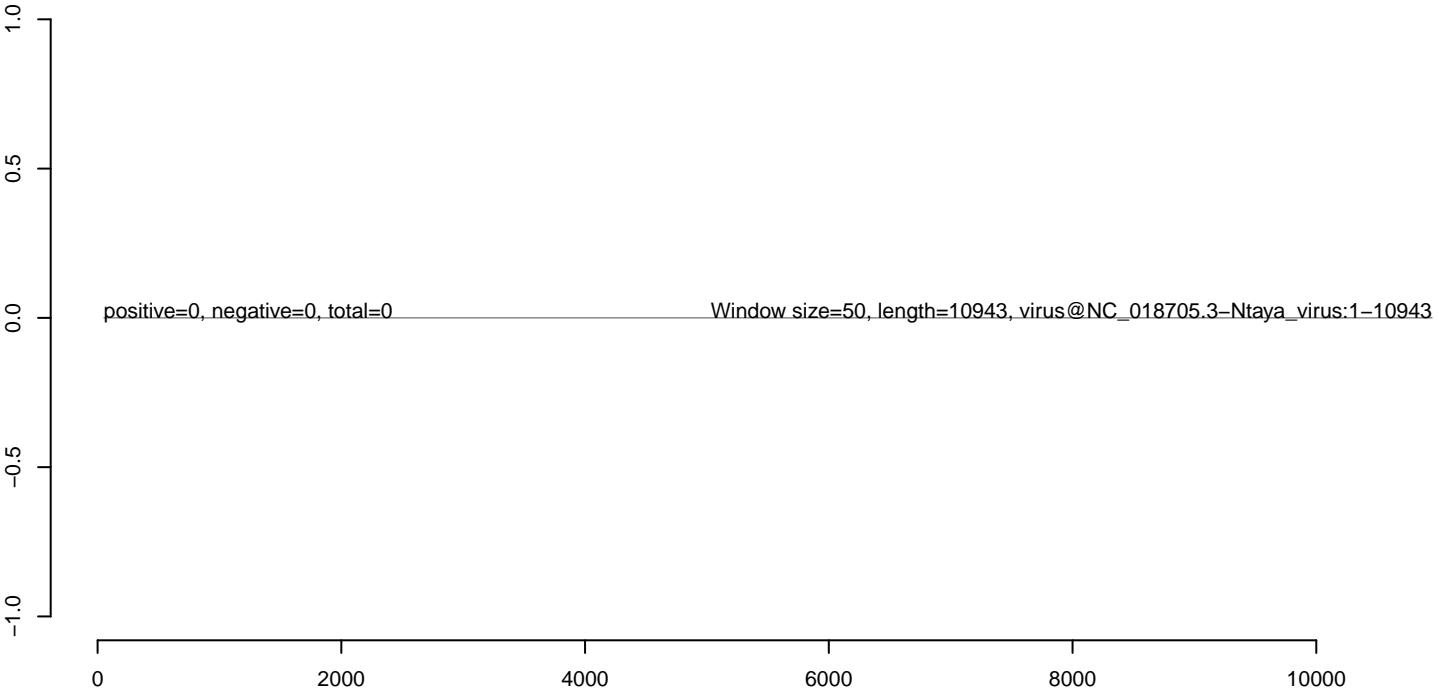
### AeAeg\_Fcanc\_GH.18\_23.rep



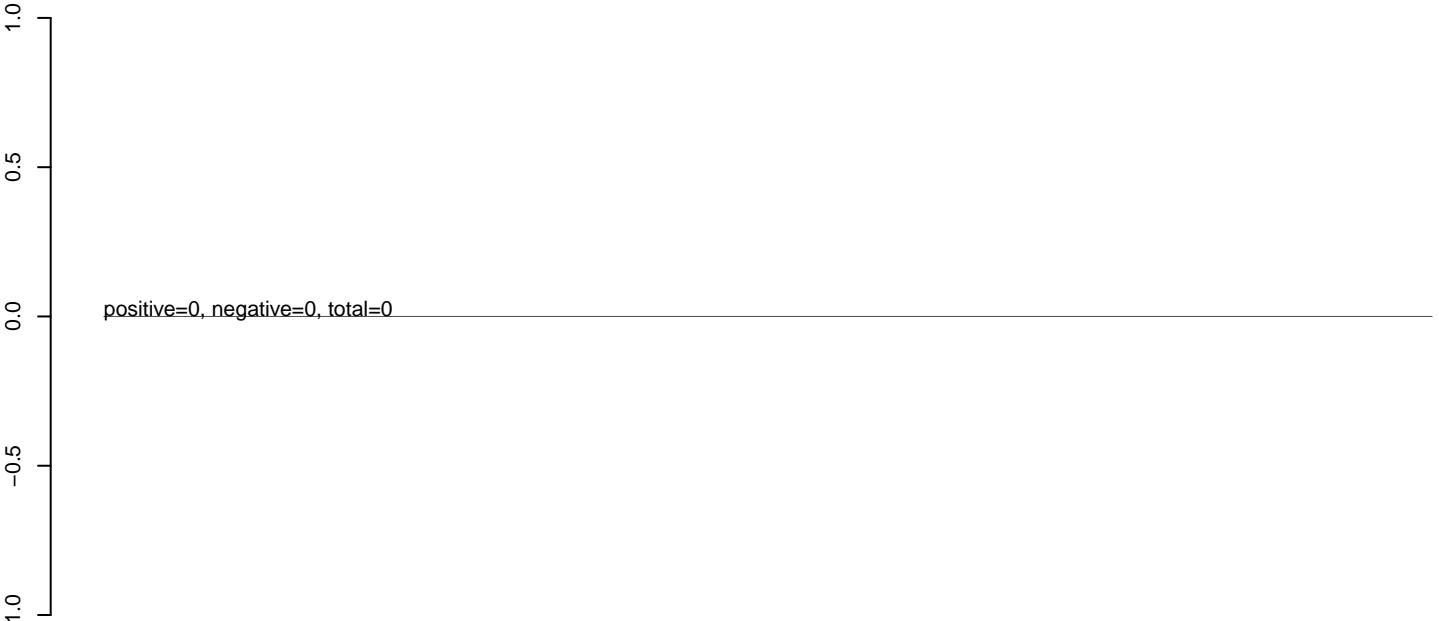
### AeAeg\_Fcanc\_GH.24\_35.rep



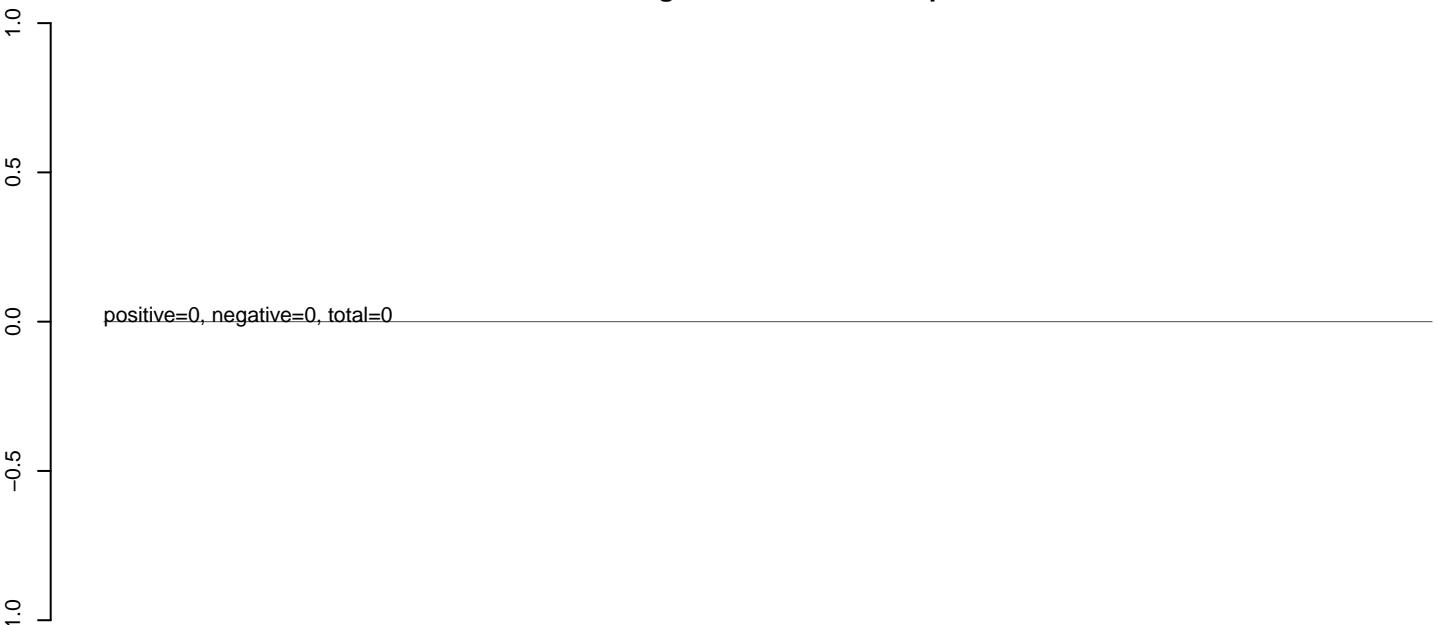
### AeAeg\_Fcanc\_GH.rep



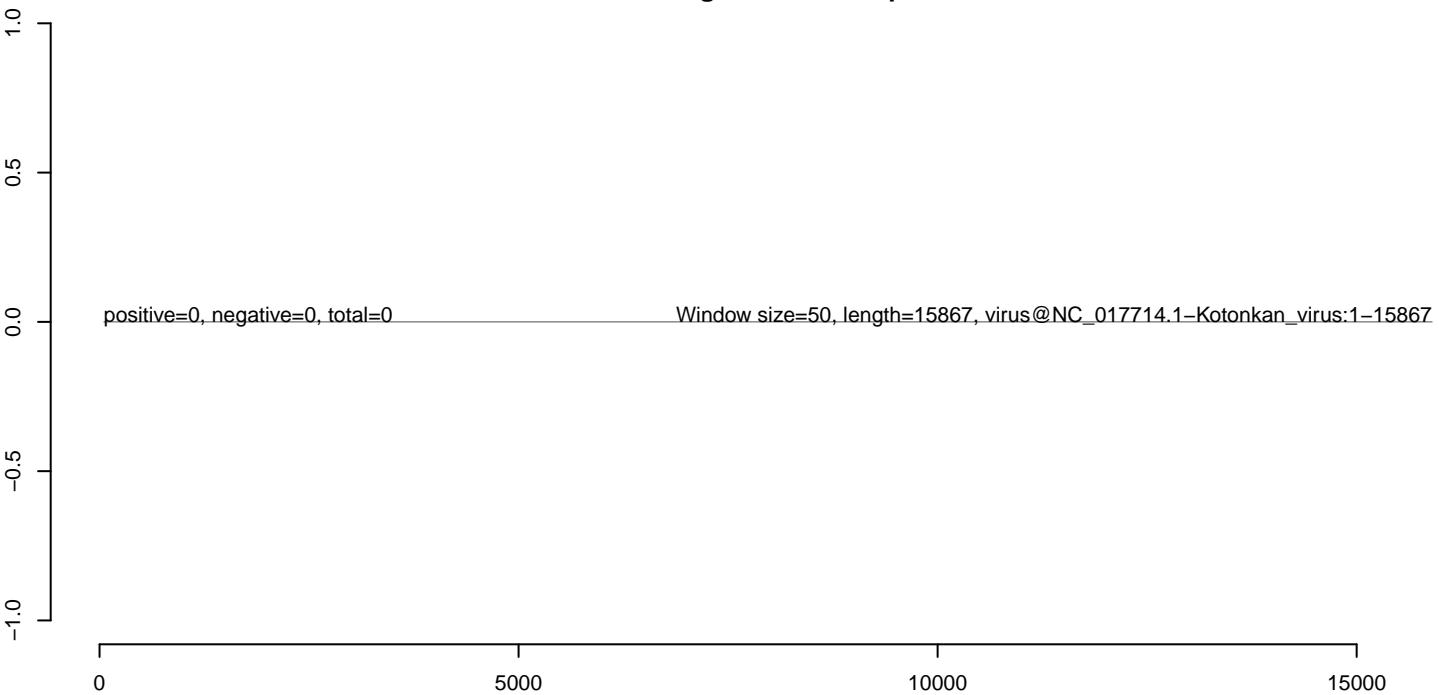
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=14717, virus@NC\_017685.1–Obodhiang\_virus:1–14717

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window.size=50, length=10733, virus@NC\_017086.1-Chaoyang\_virus:1-10733

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=11688, virus@NC\_016959.1–Ndumu\_virus:1–11688

0 2000 4000 6000 8000 10000 12000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0 1979, virus@NC\_014395.1-Rift\_Valley\_fever\_virus\_segment\_S\_plus\_segment\_M\_plus\_segment\_L:1-11979

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

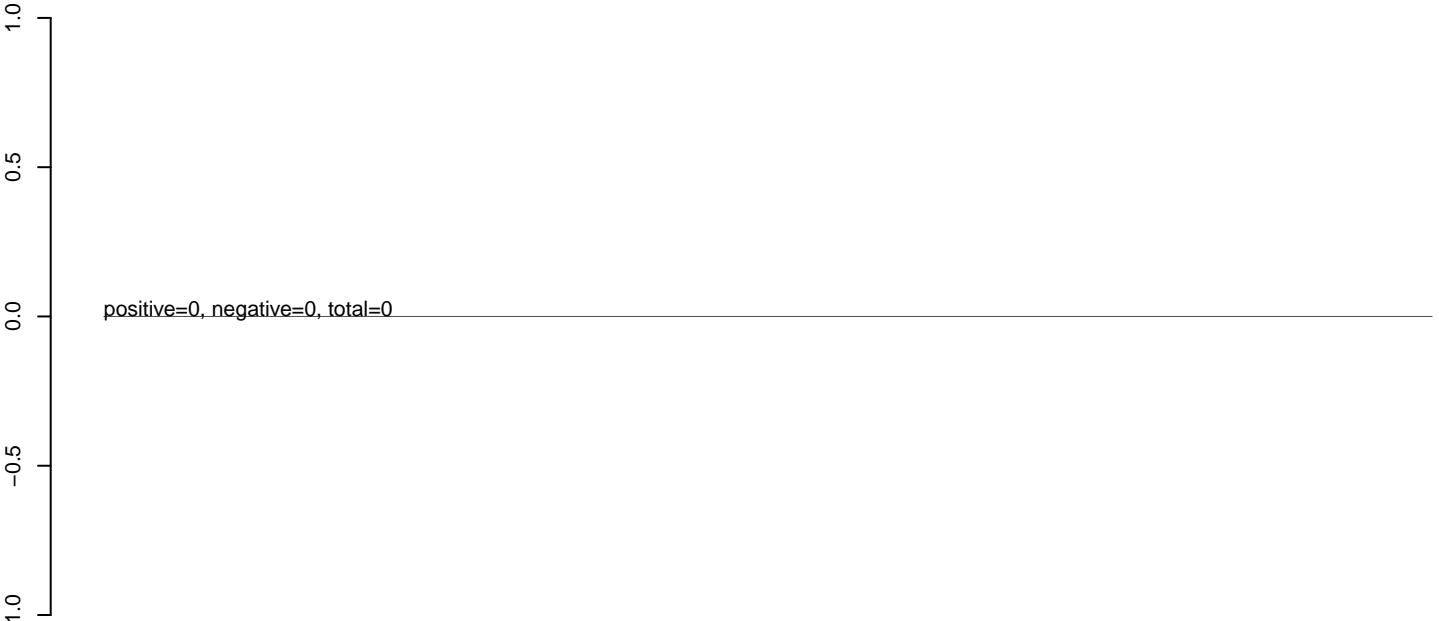
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

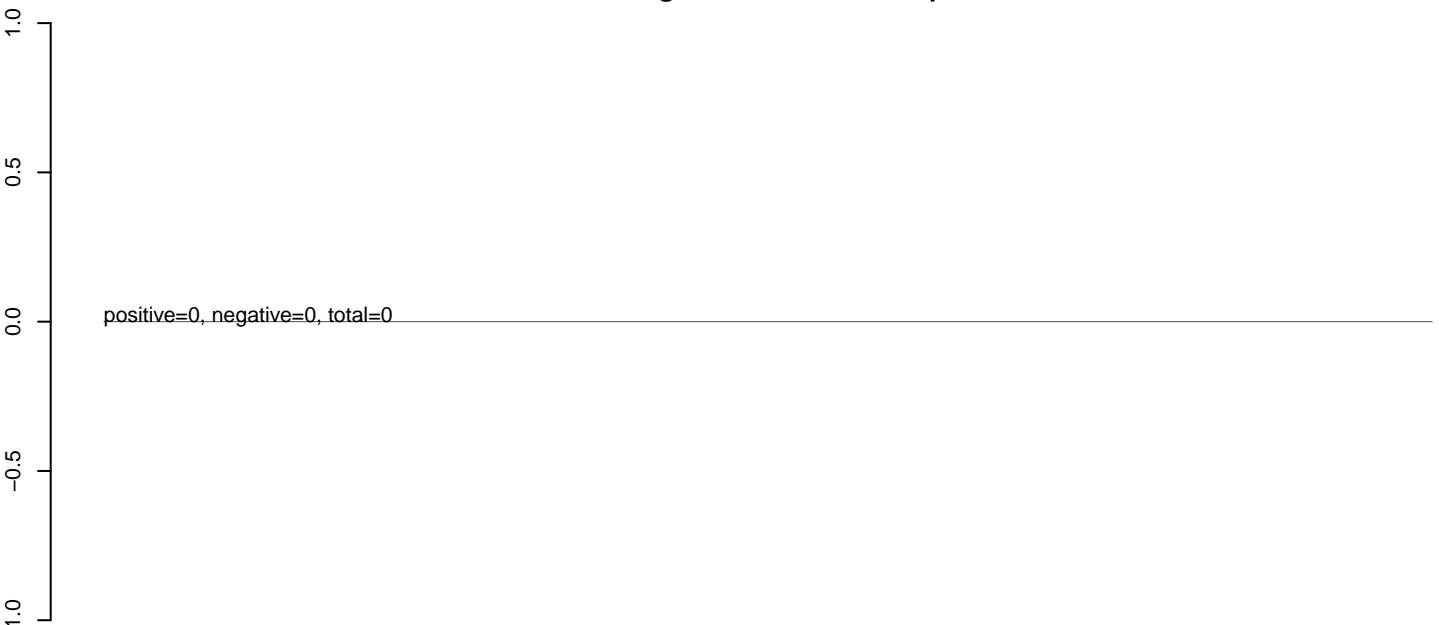
Window size=50, length=11381, virus@NC\_013528.1–Fort\_Morgan\_virus:1–11381

0 2000 4000 6000 8000 10000

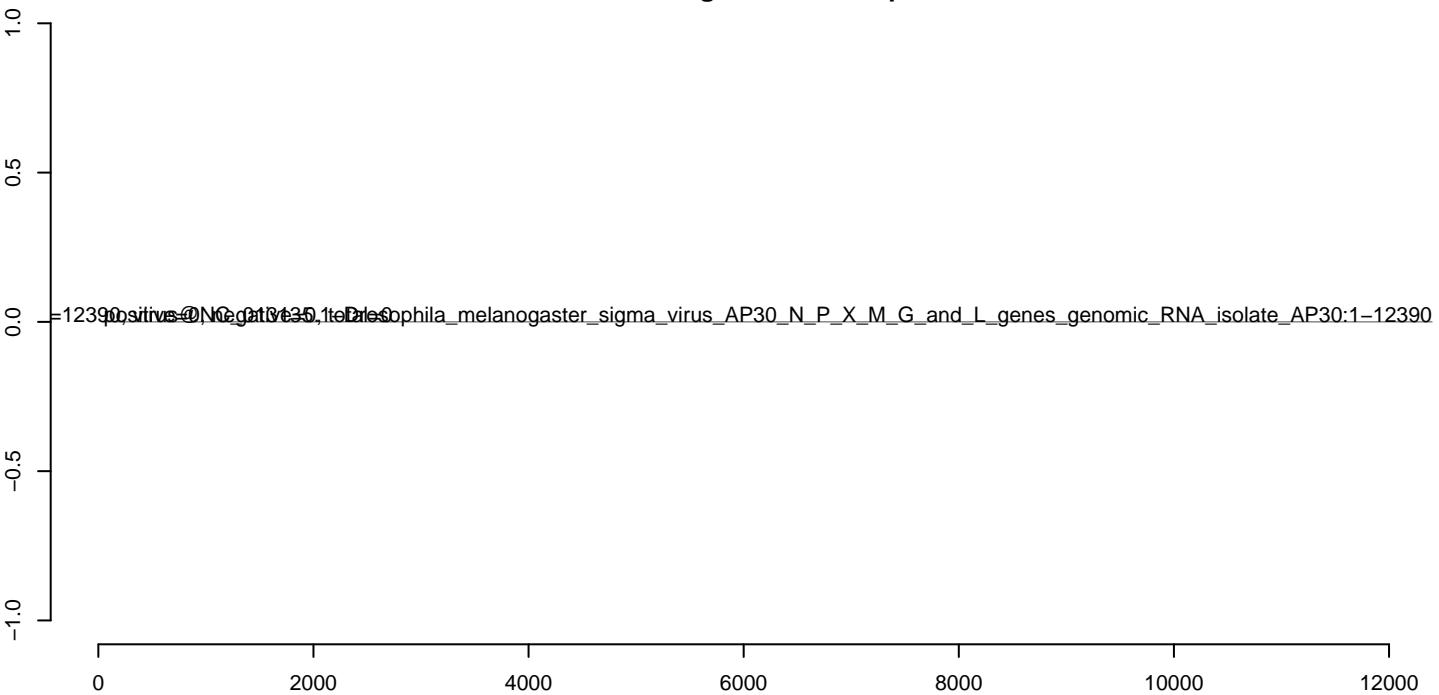
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10814, virus@NC\_012735.1-Wesselsbron\_virus:1-10814

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

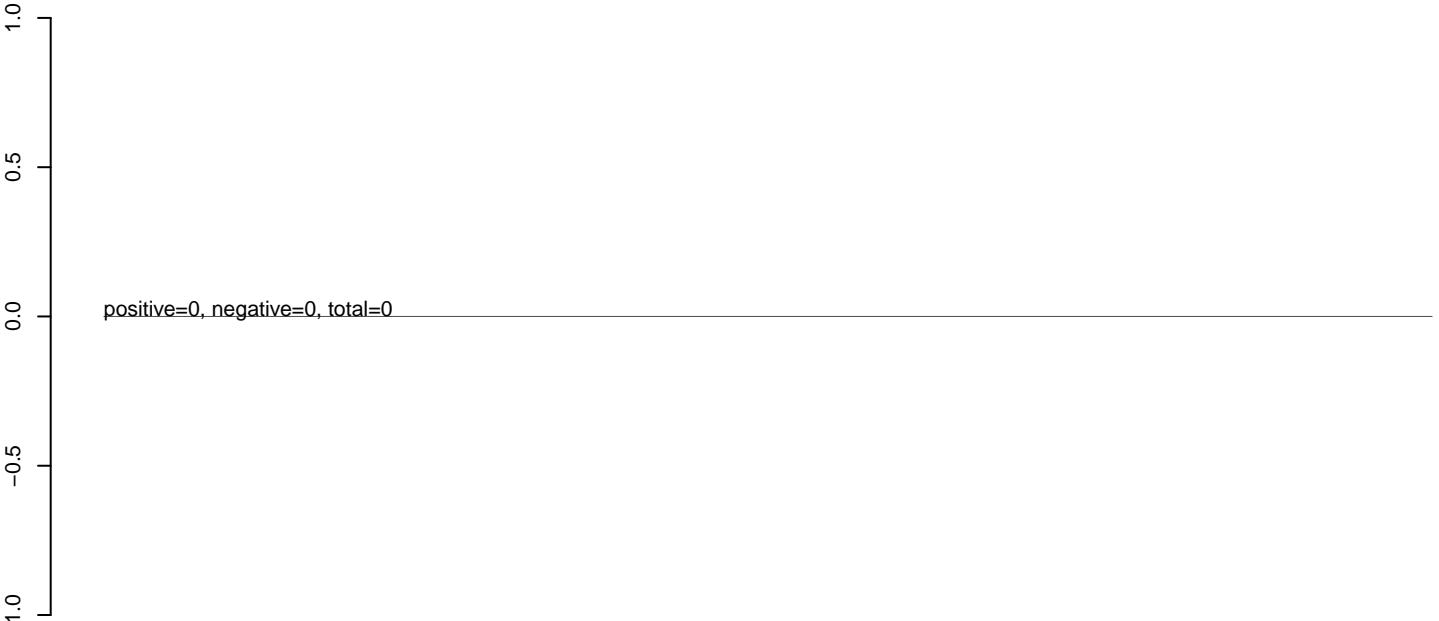
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

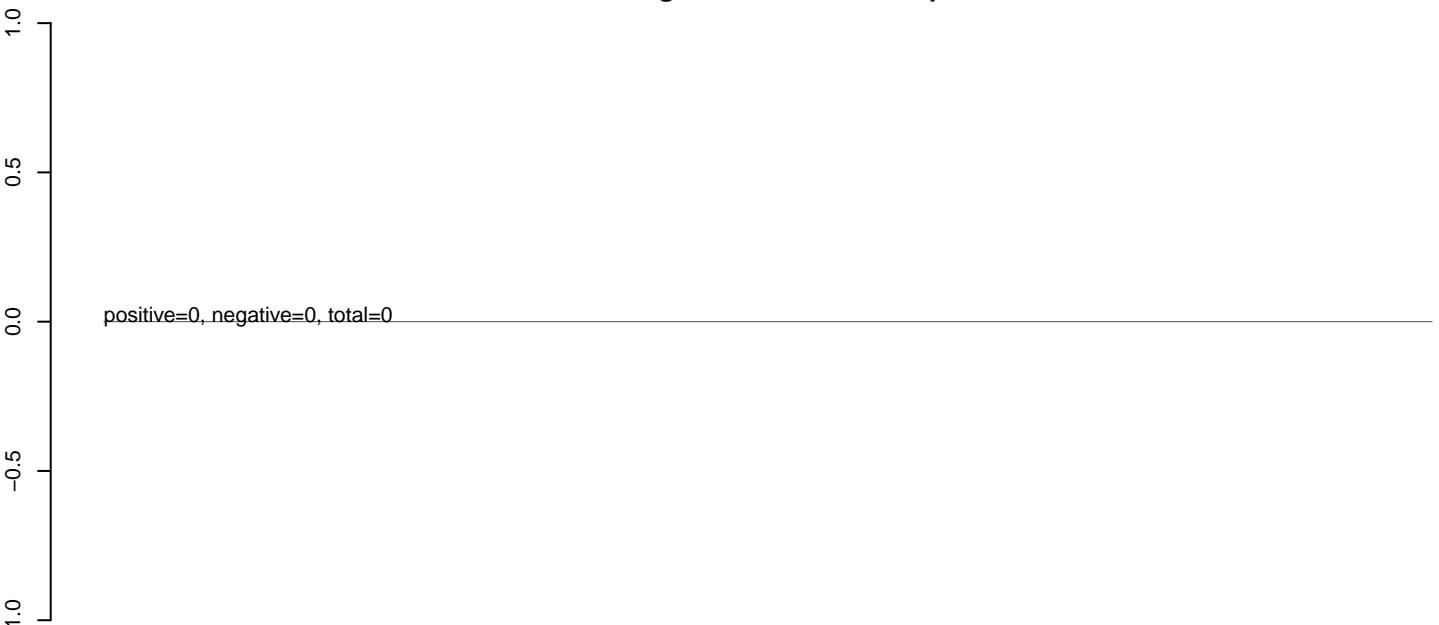
Window size=50, length=10865, virus@NC\_012671.1–Quang\_Binh\_virus:1–10865

0 2000 4000 6000 8000 10000

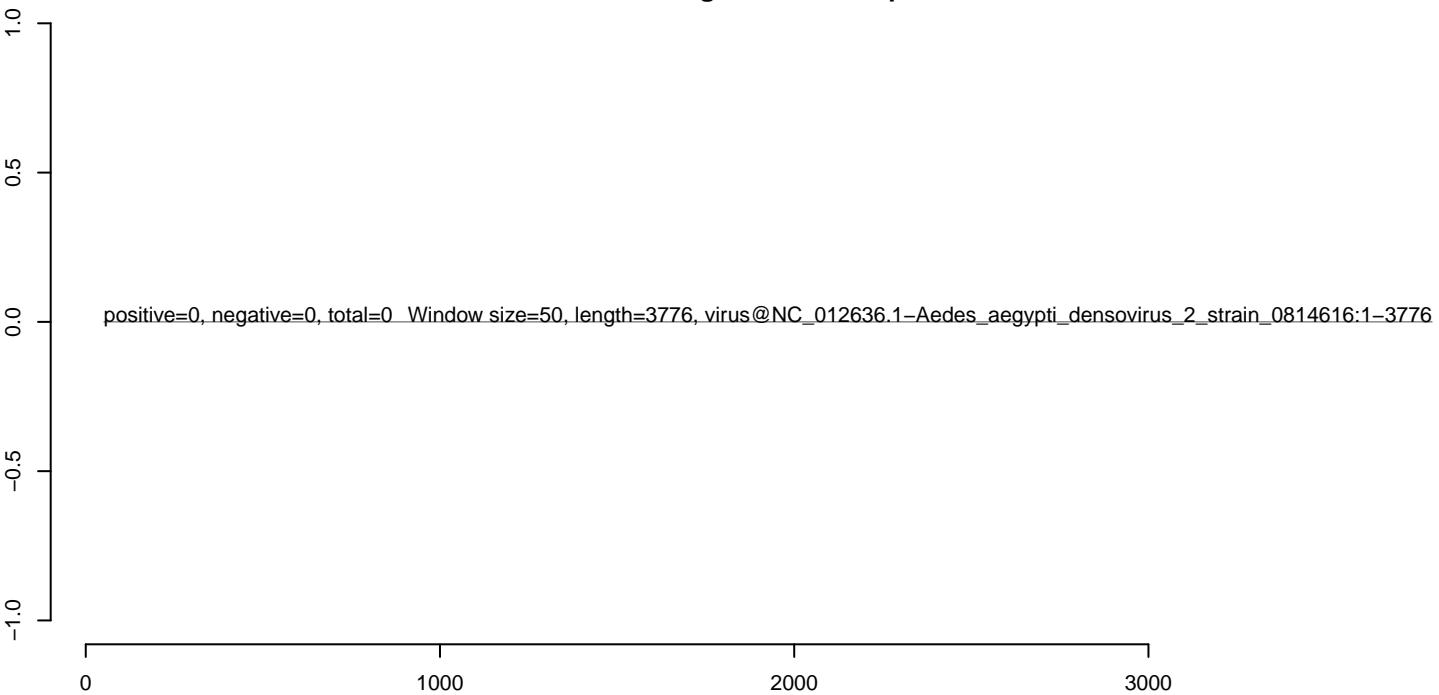
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=11526, virus@NC\_012561.1-Highlands\_J\_virus:1-11526

0 2000 4000 6000 8000 10000 12000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

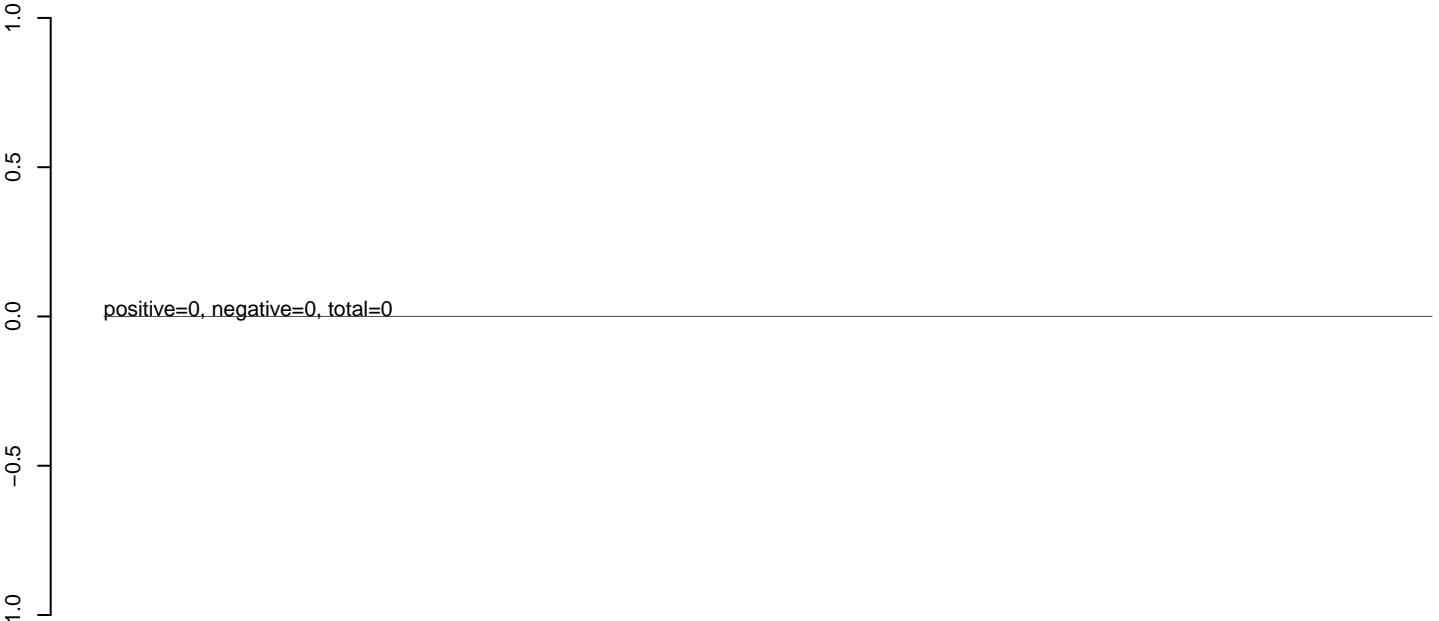
### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

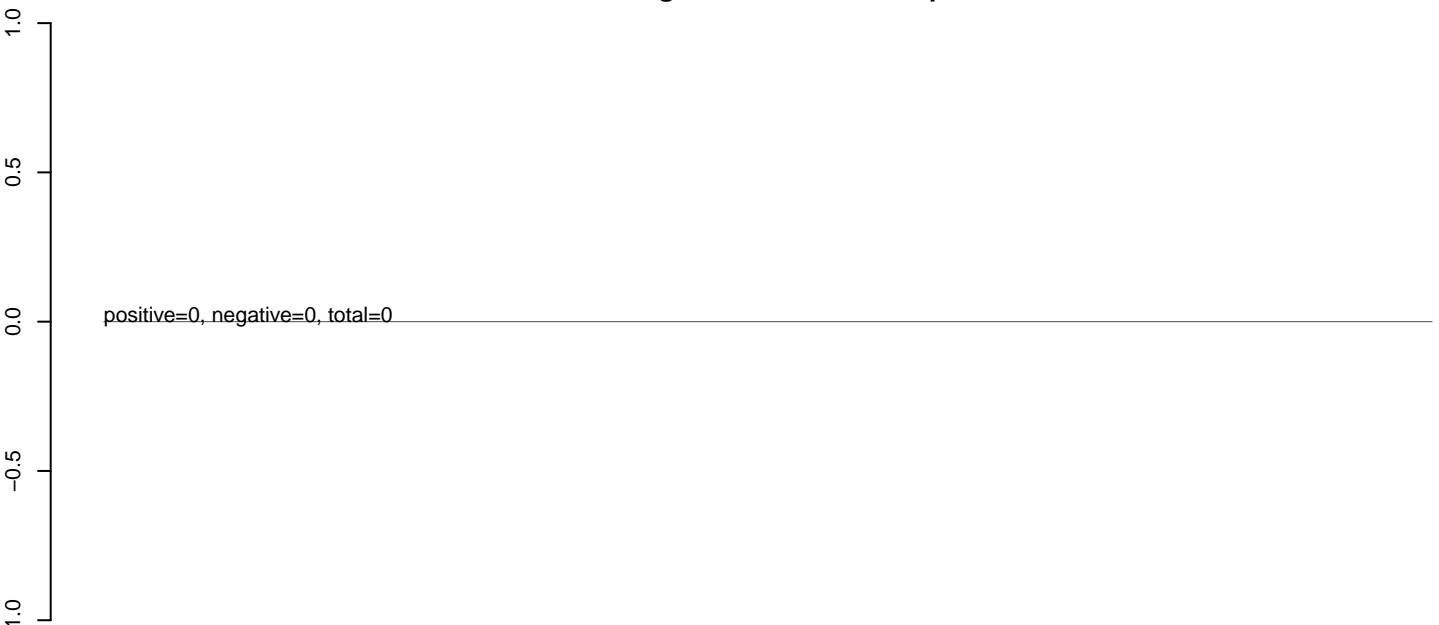
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0, length=10794, virus@NC\_012532.1-Zika\_virus\_isolate\_ZIKV\_Monkey\_Uganda\_MR766\_1947:1-10794

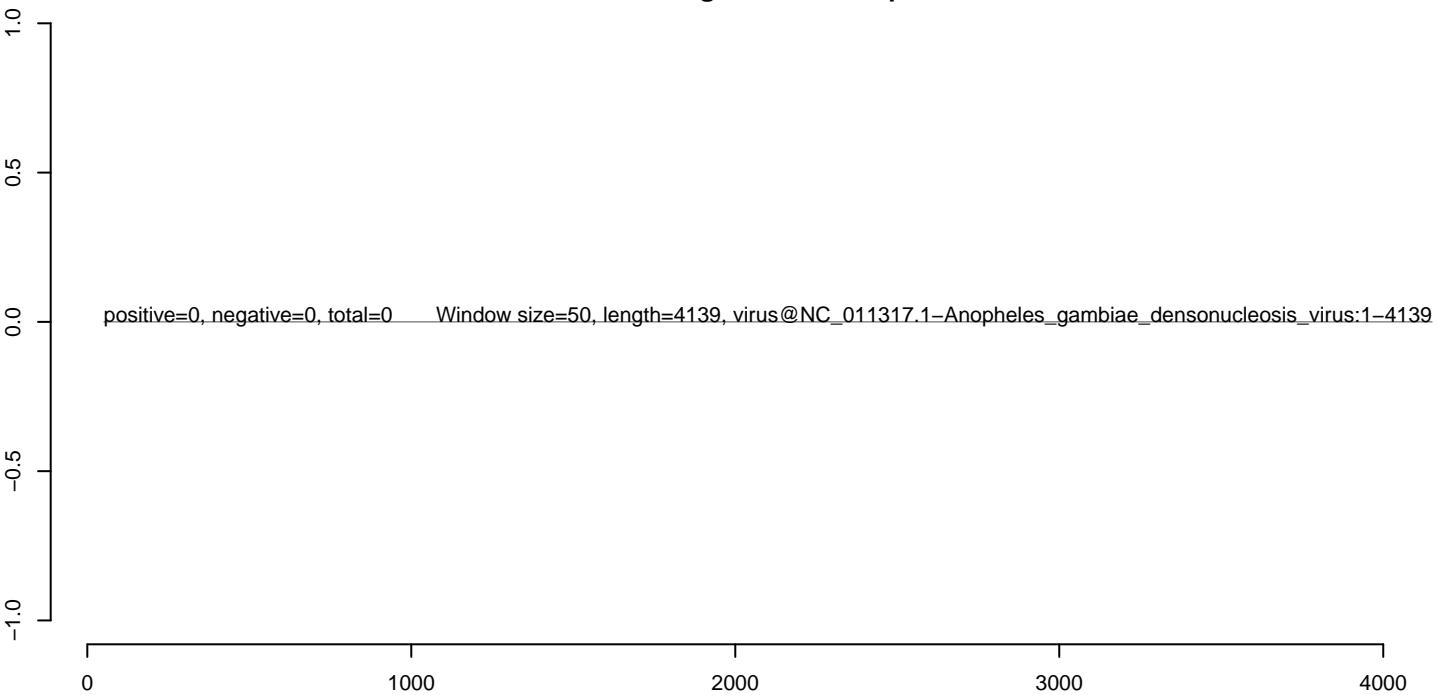
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10755, virus@NC\_009028.2-llheus\_virus:1-10755

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10793, virus@NC\_008719.1–Sepik\_virus:1–10793

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10940, virus@NC\_007580.2-St\_Louis\_encephalitis\_virus:1-10940

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=11440, virus@NC\_007020.1-Tupaia\_virus:1-11440

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10653, virus@NC\_006947.1-Karshi\_virus:1-10653

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

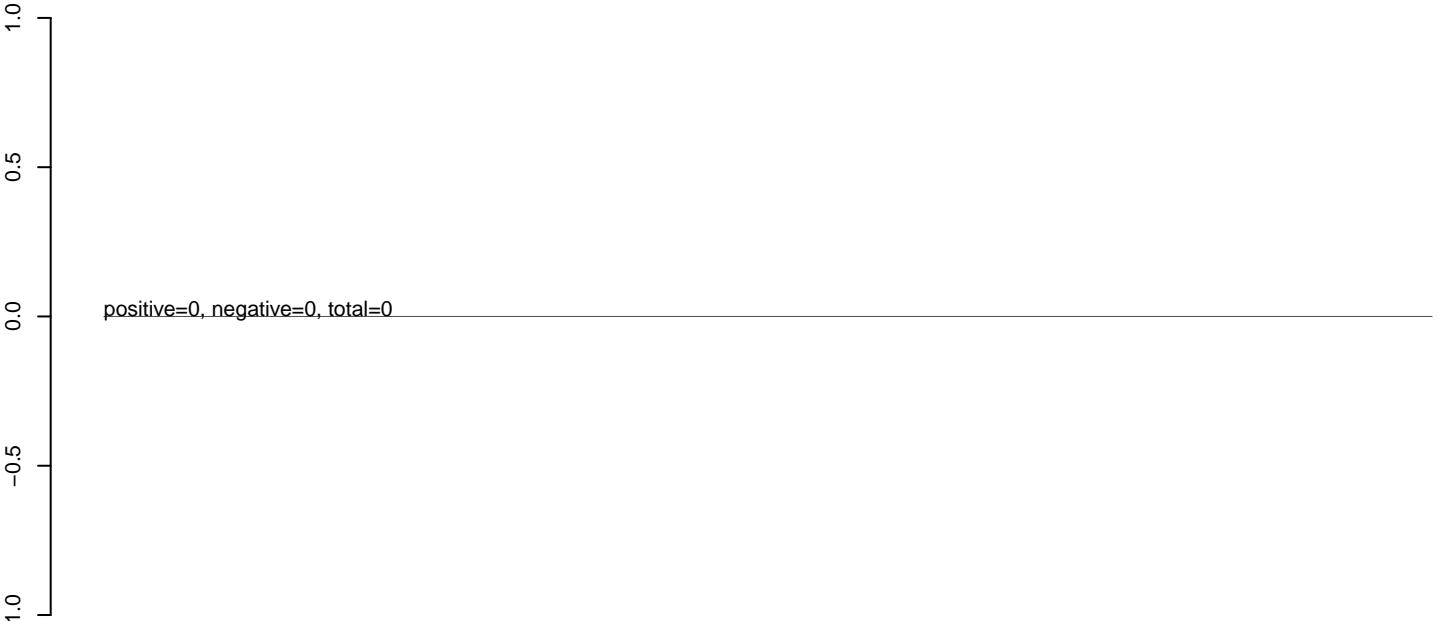
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

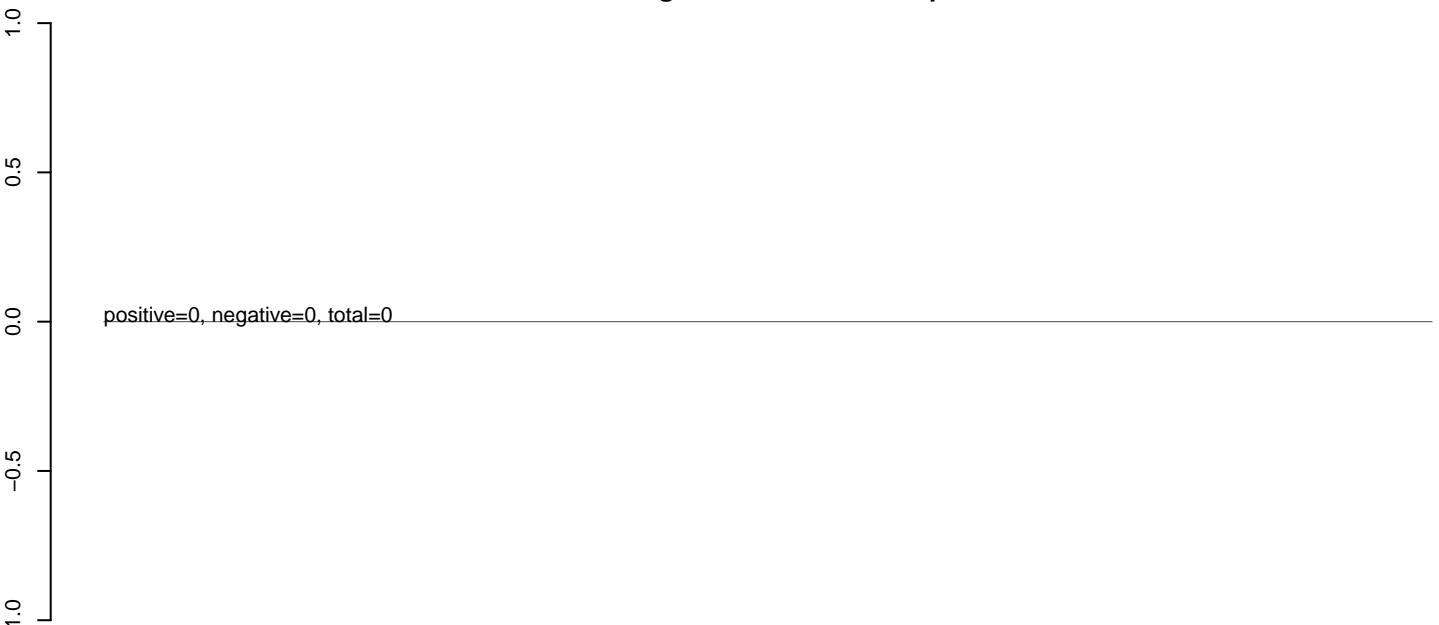
Window size=50, length=11597, virus@NC\_006558.1-Getah\_virus:1-11597

0 2000 4000 6000 8000 10000 12000

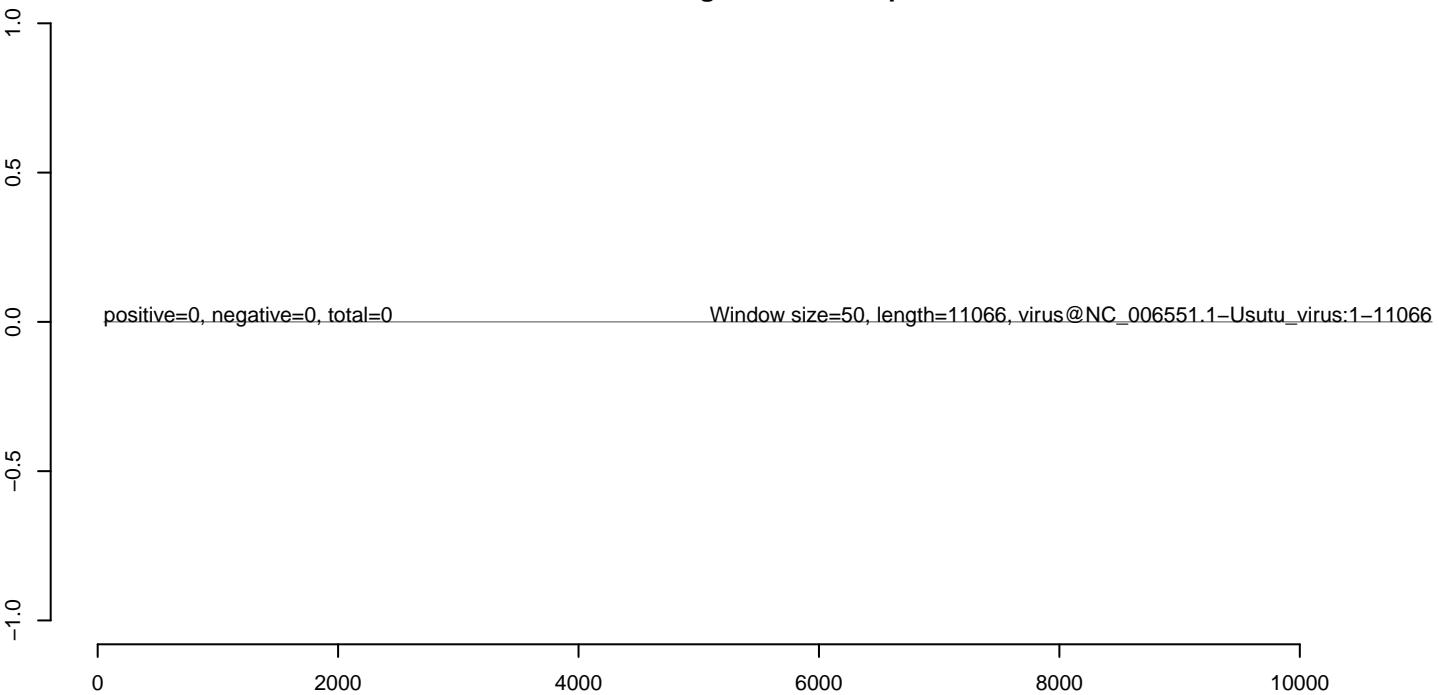
### AeAeg\_Fcanc\_GH.18\_23.rep



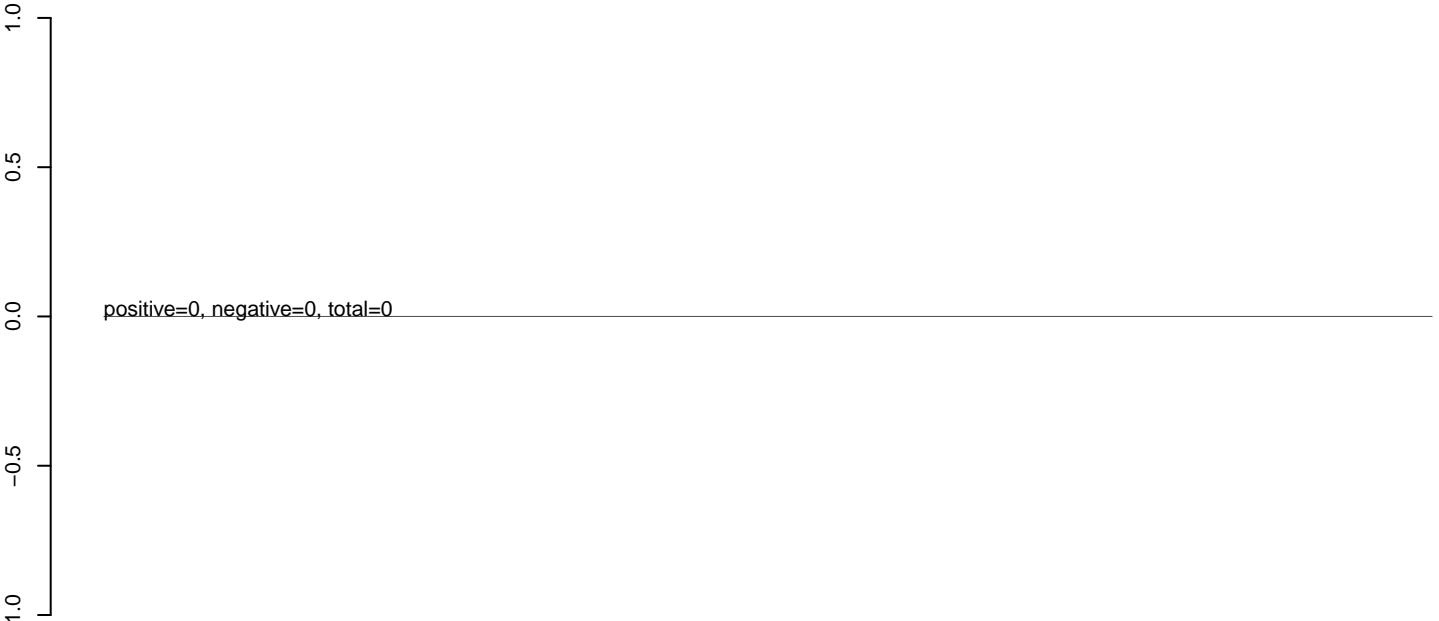
### AeAeg\_Fcanc\_GH.24\_35.rep



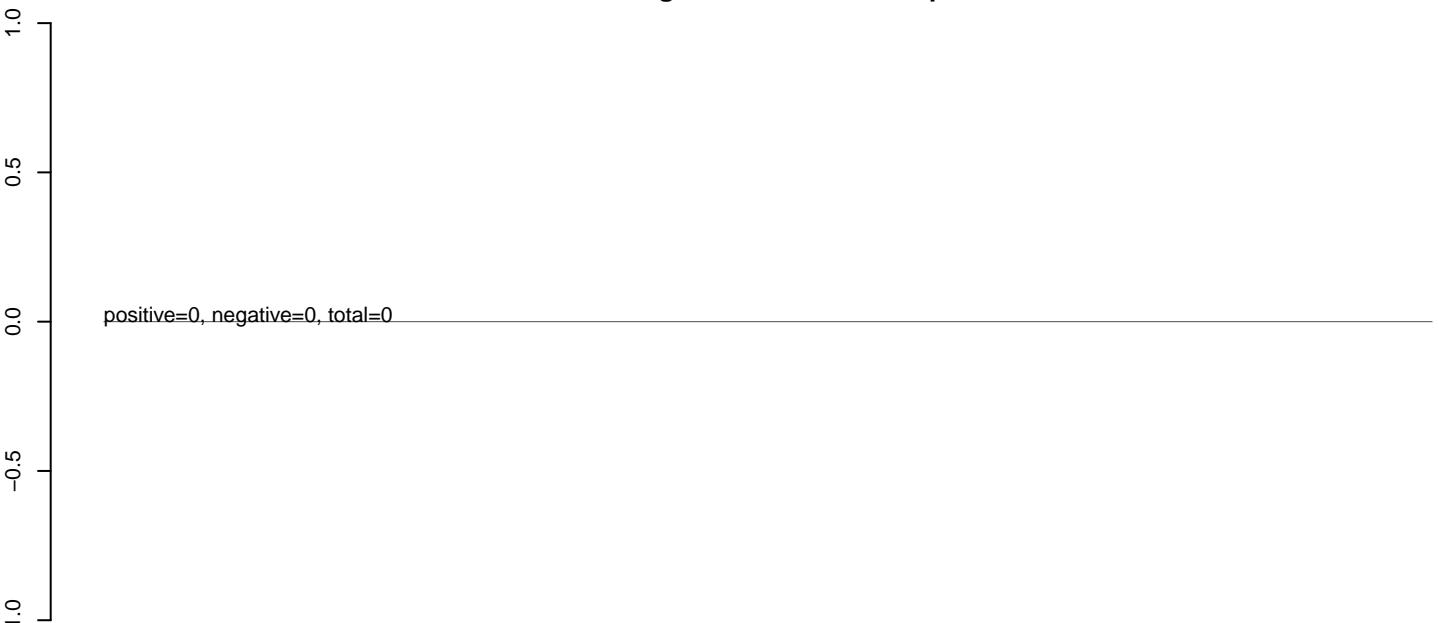
### AeAeg\_Fcanc\_GH.rep



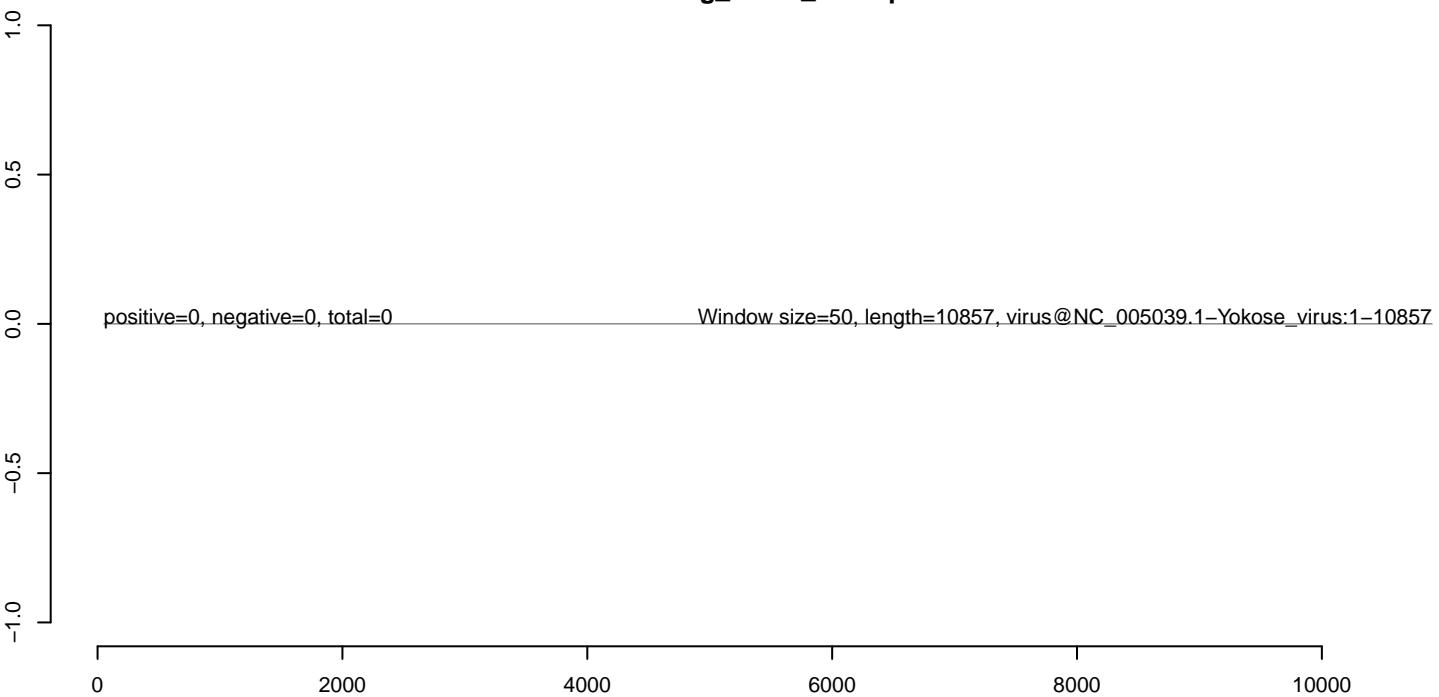
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10685, virus@NC\_004355.1-Alkhurma\_virus:1-10685

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

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

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0    Window size=50, length=11484, virus@NC\_003908.1-Western\_equine\_encephalomyelitis\_virus:1-11484

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.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

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

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

0 2000 4000 6000 8000 10000 12000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

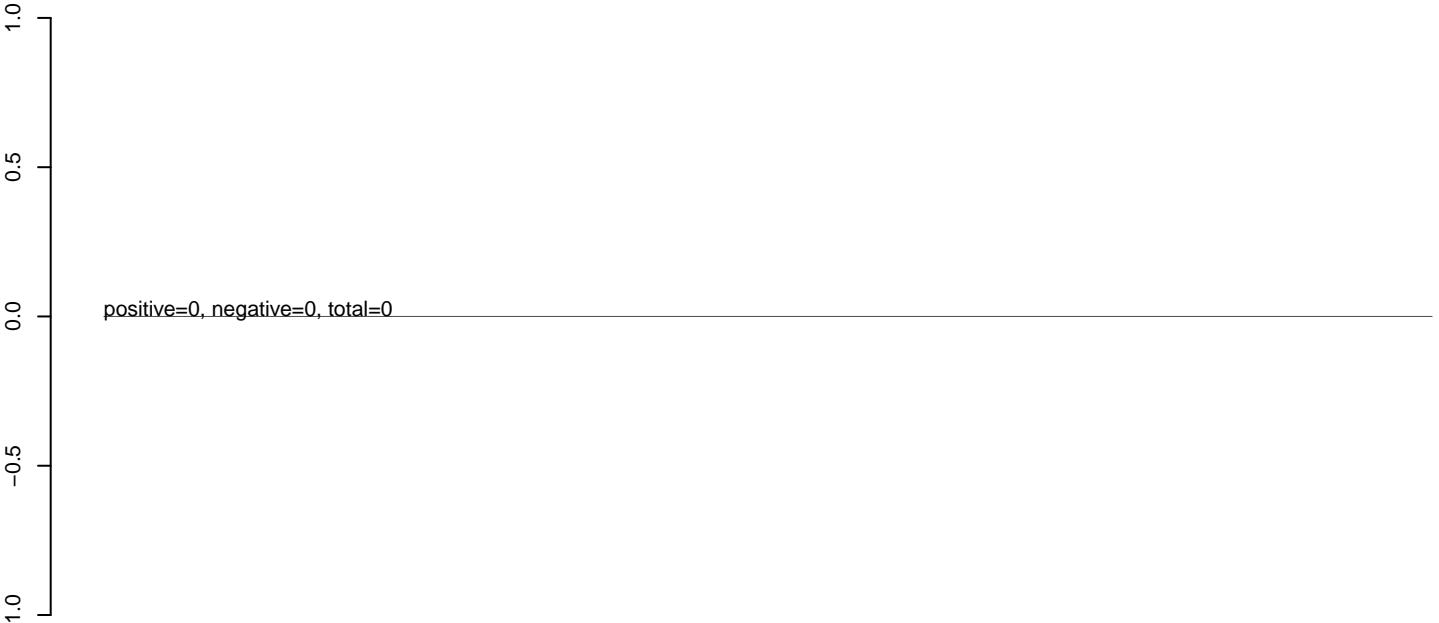
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

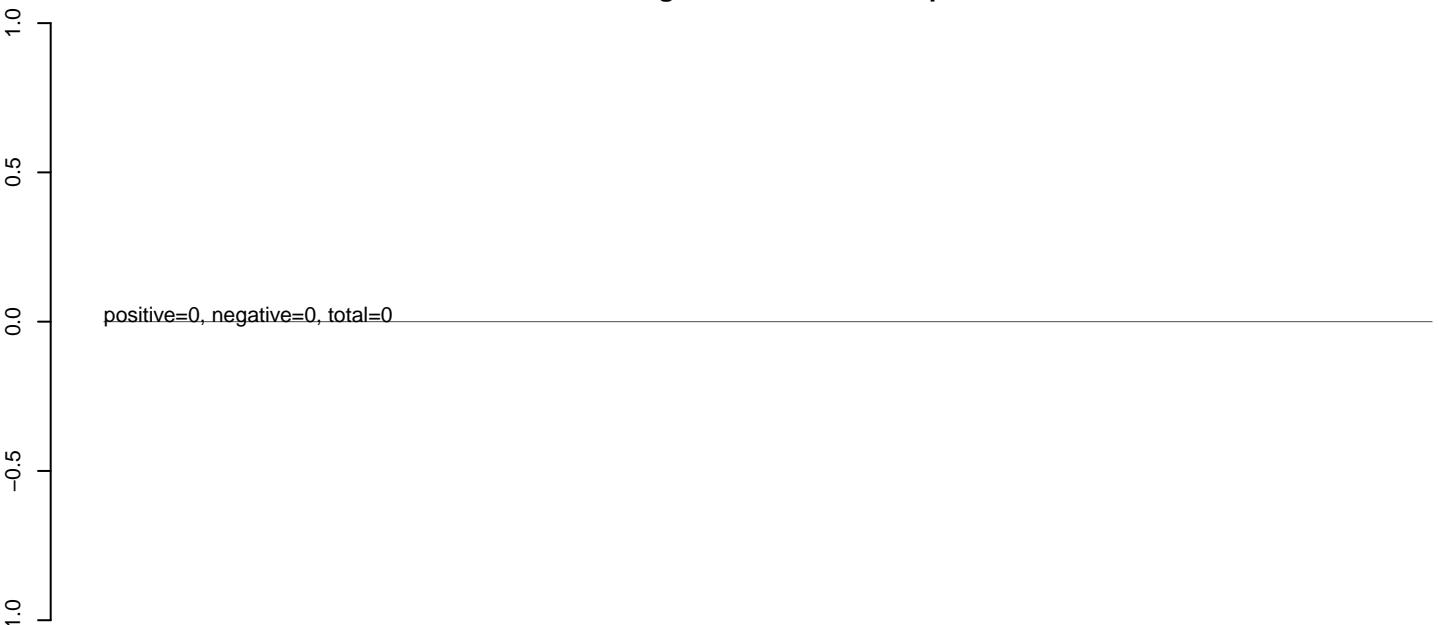
Window size=50, length=10943, virus@NC\_003690.1-Langat\_virus:1-10943

0 2000 4000 6000 8000 10000

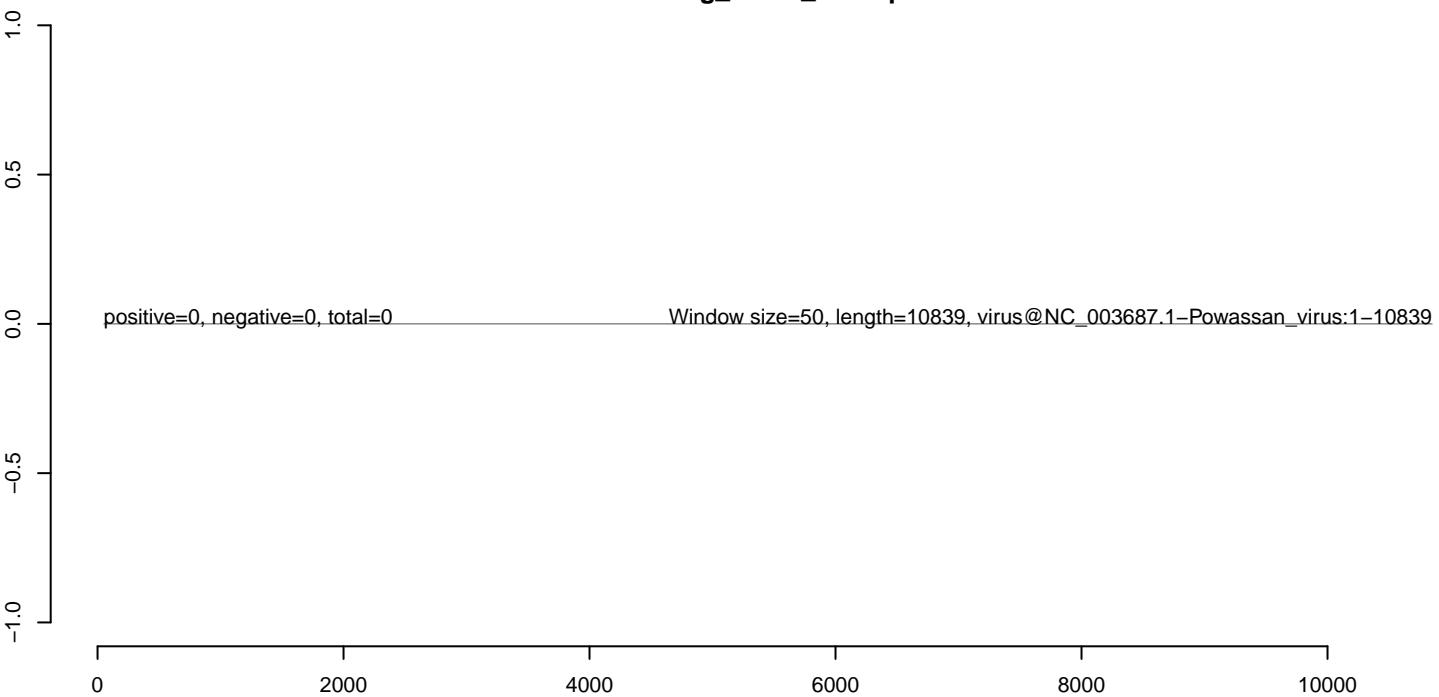
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.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

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

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

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

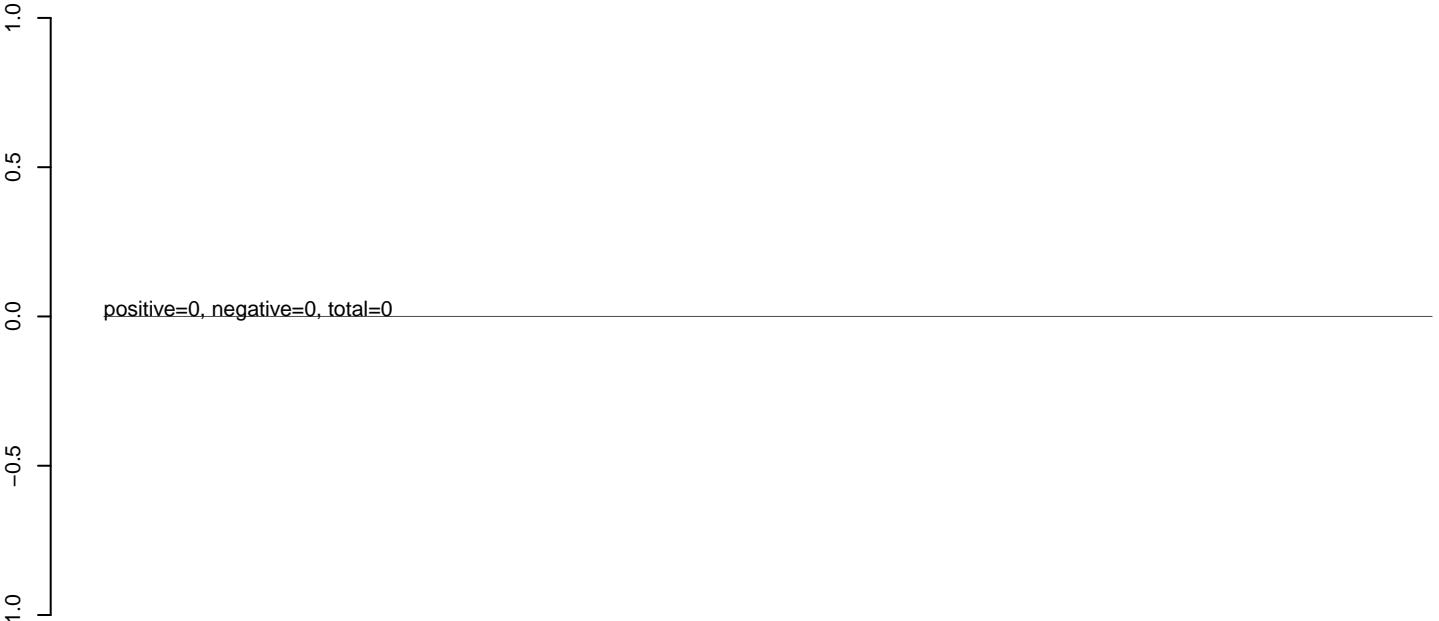
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

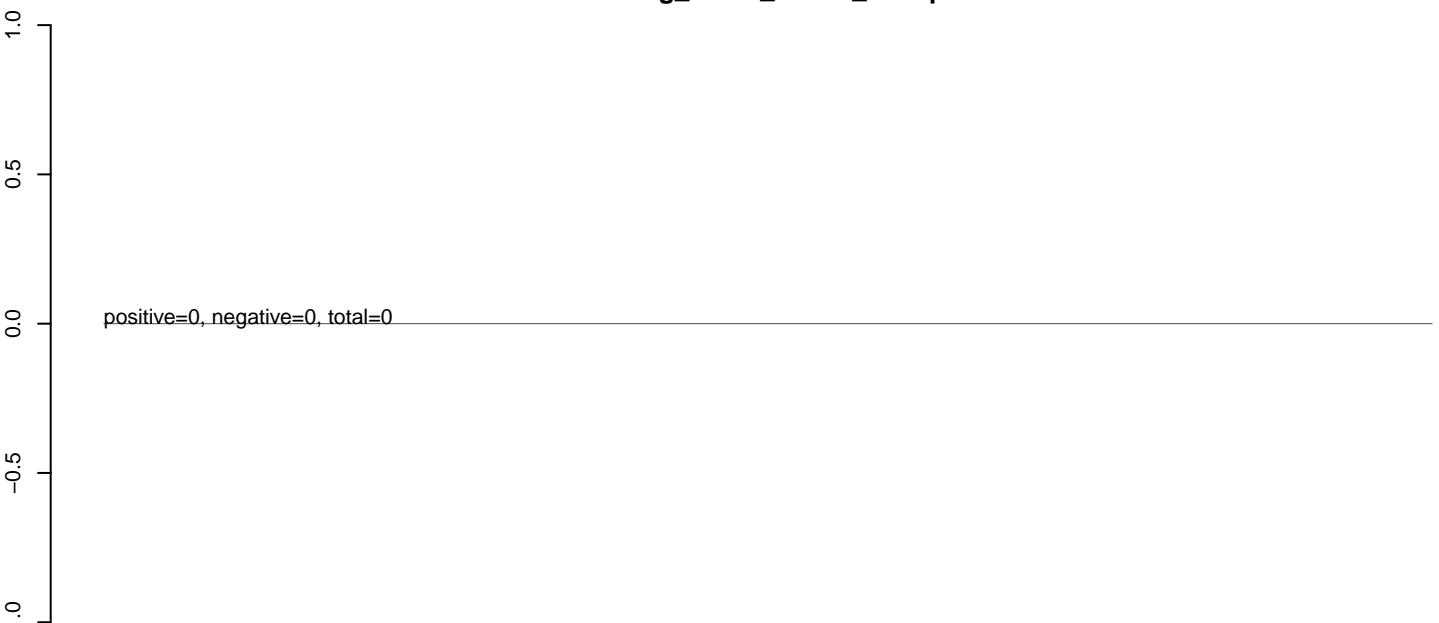
Window size=50, length=10600, virus@NC\_003635.1-Modoc\_virus:1-10600

0 2000 4000 6000 8000 10000

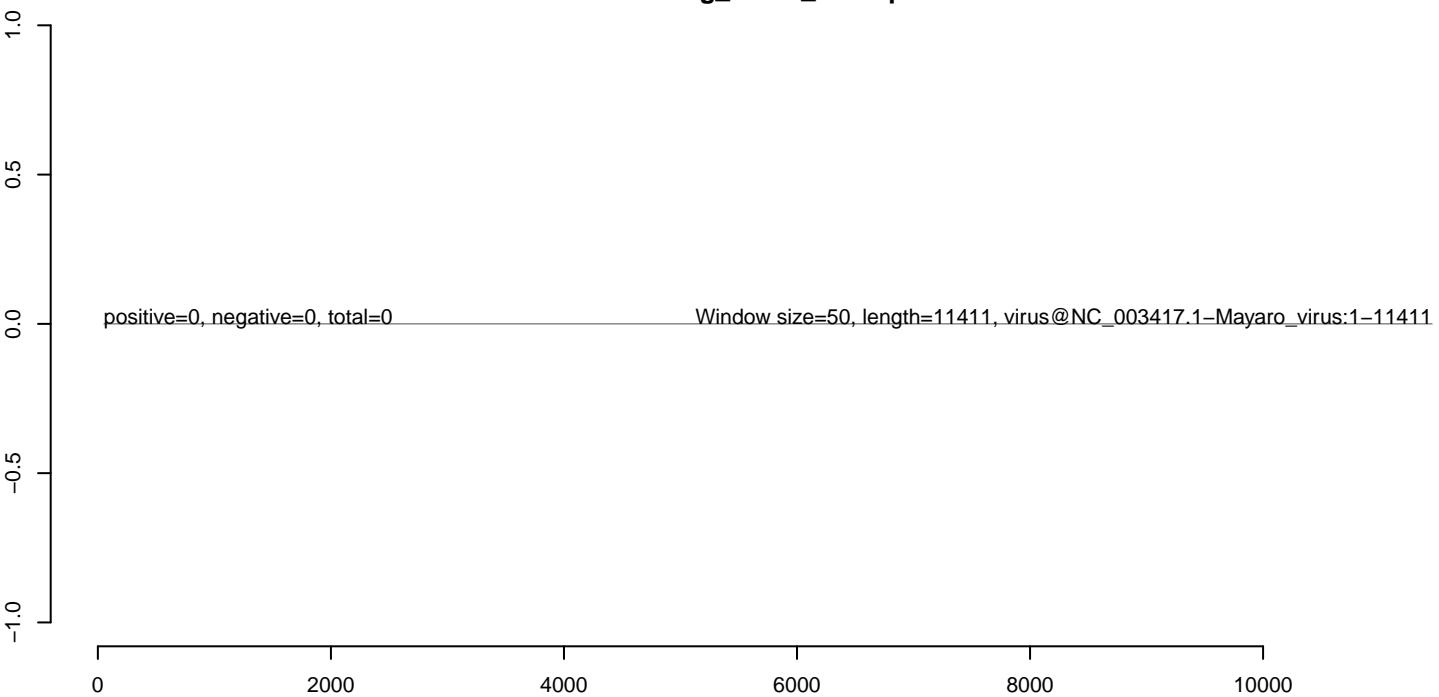
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=11442, virus@NC\_003215.1-Semliki\_forest\_virus:1-11442

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=11019, virus@NC\_002803.1-Spring\_viraemia\_of\_carp\_virus:1-11019

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

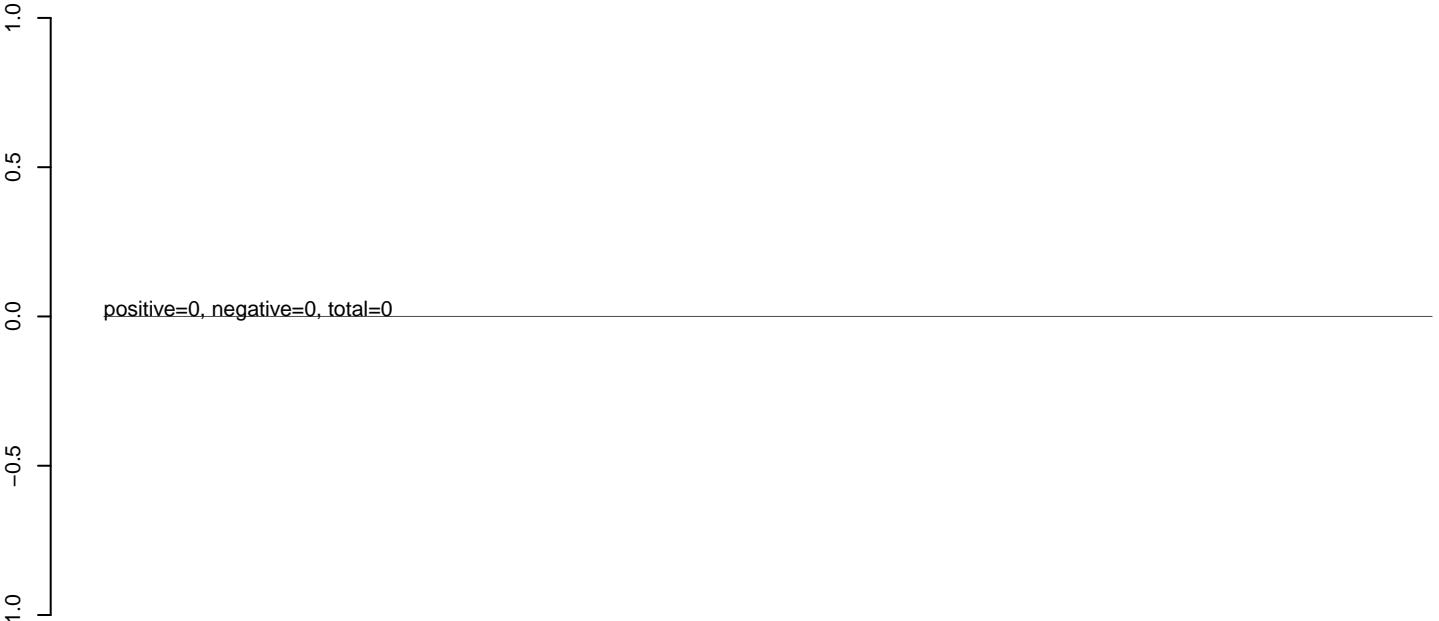
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

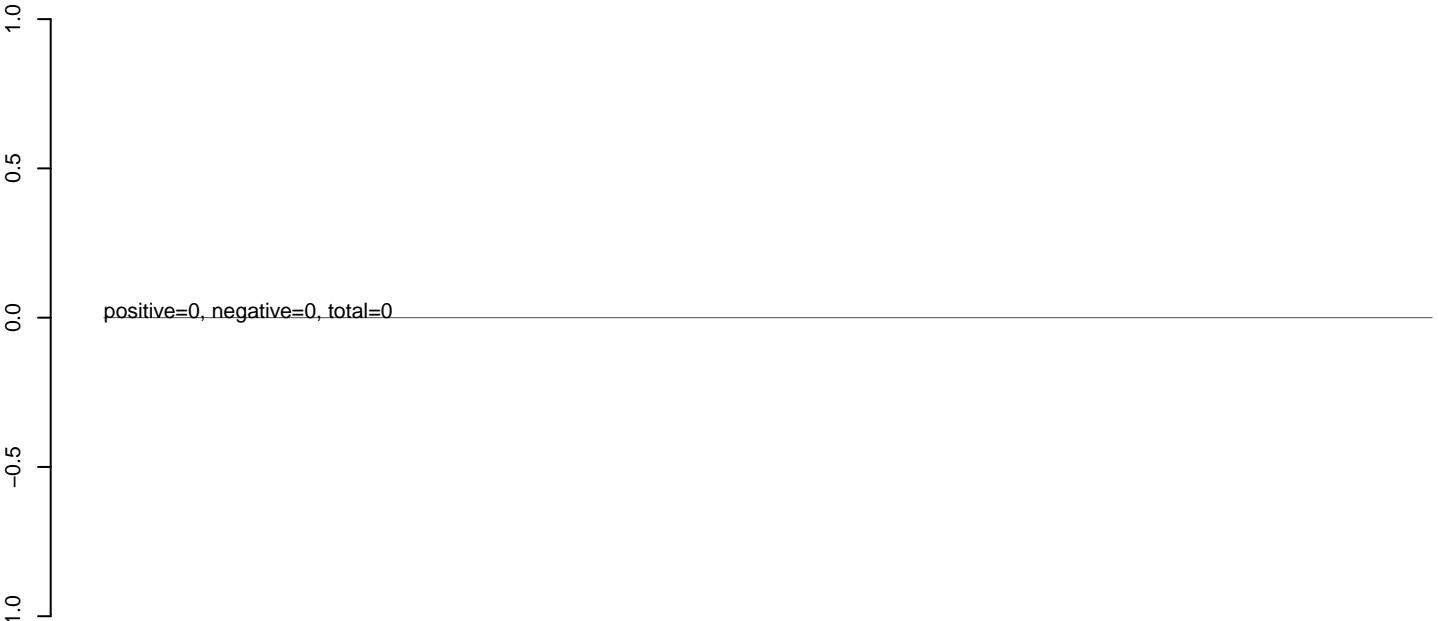
Window size=50, length=10649, virus@NC\_002640.1-Dengue\_virus\_4:1-10649

0 2000 4000 6000 8000 10000

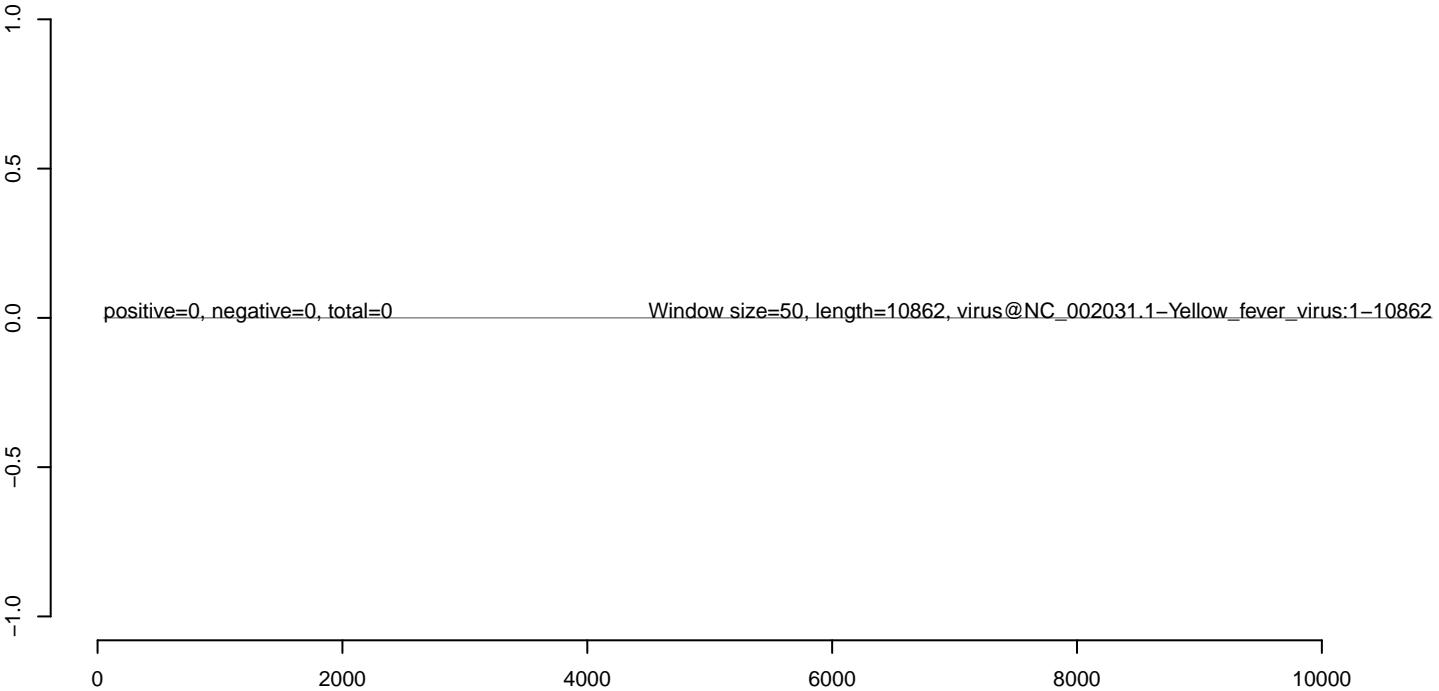
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

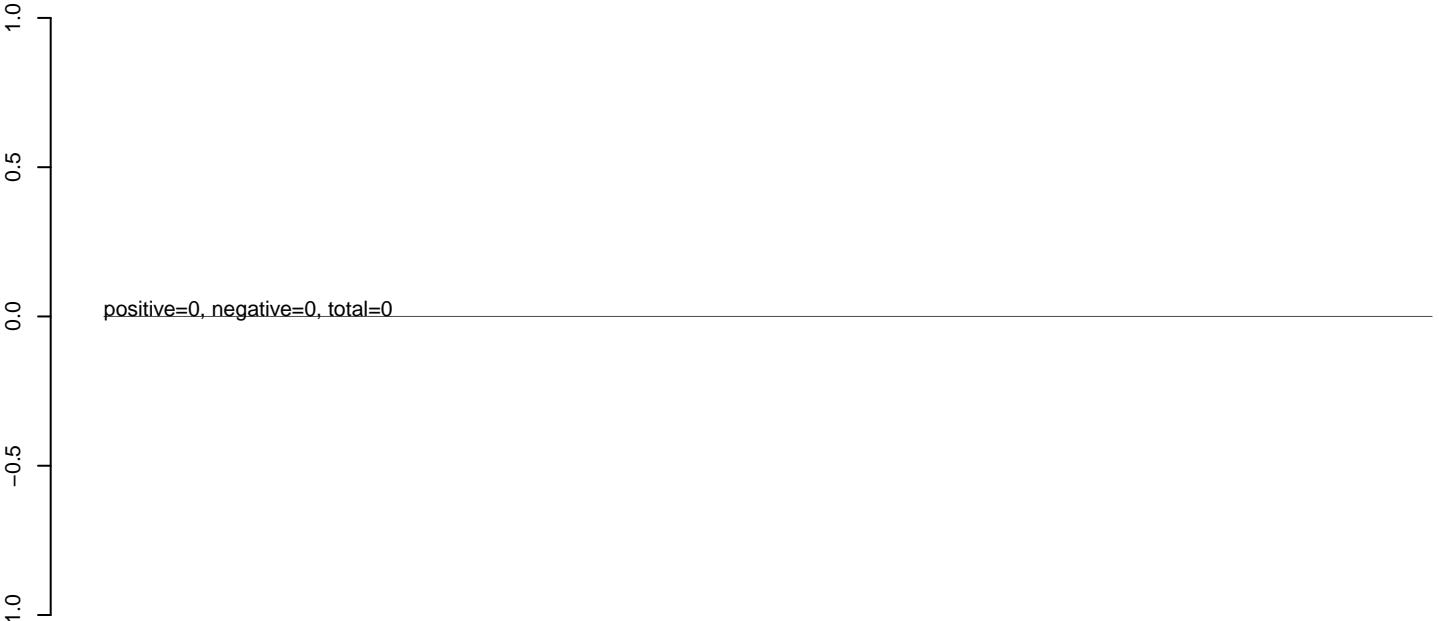
### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

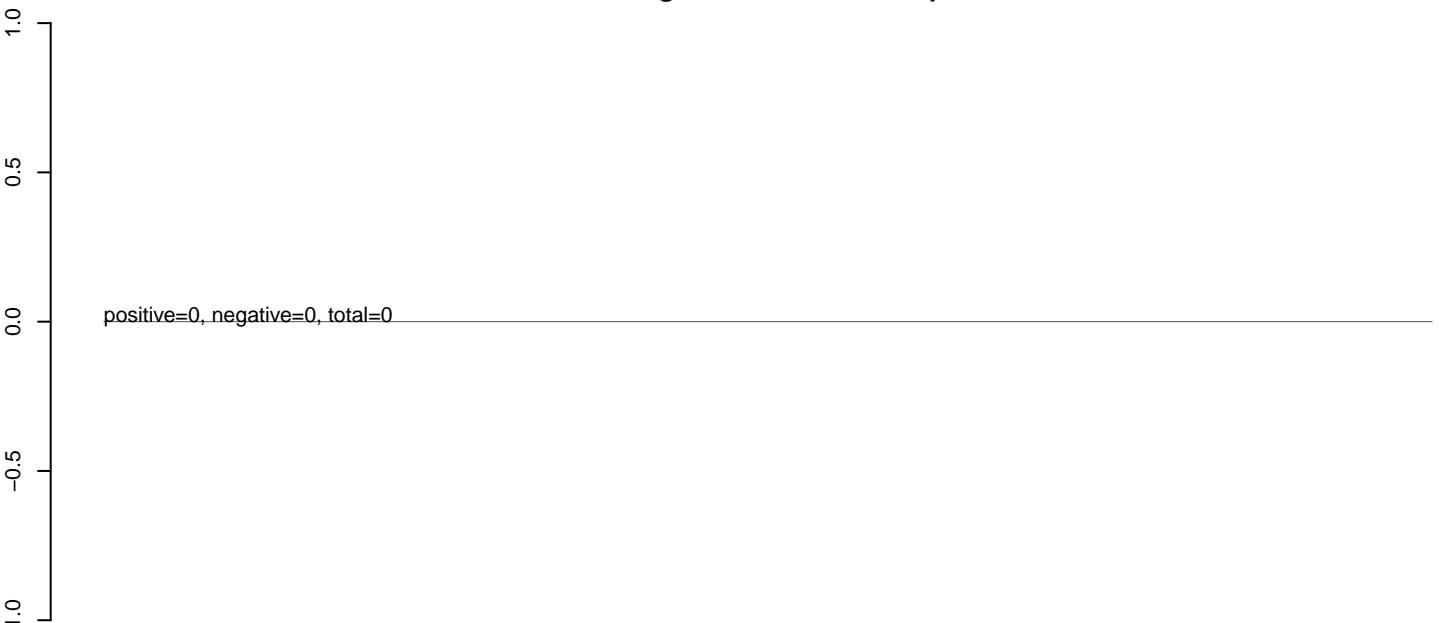
### AeAeg\_Fcanc\_GH.rep

positive=45, negative=50, total=95, length=12294, virus@NC\_001927.1-Bunyamwera\_virus\_segment\_S\_plus\_segment\_M\_plus\_segment\_L:1-12294

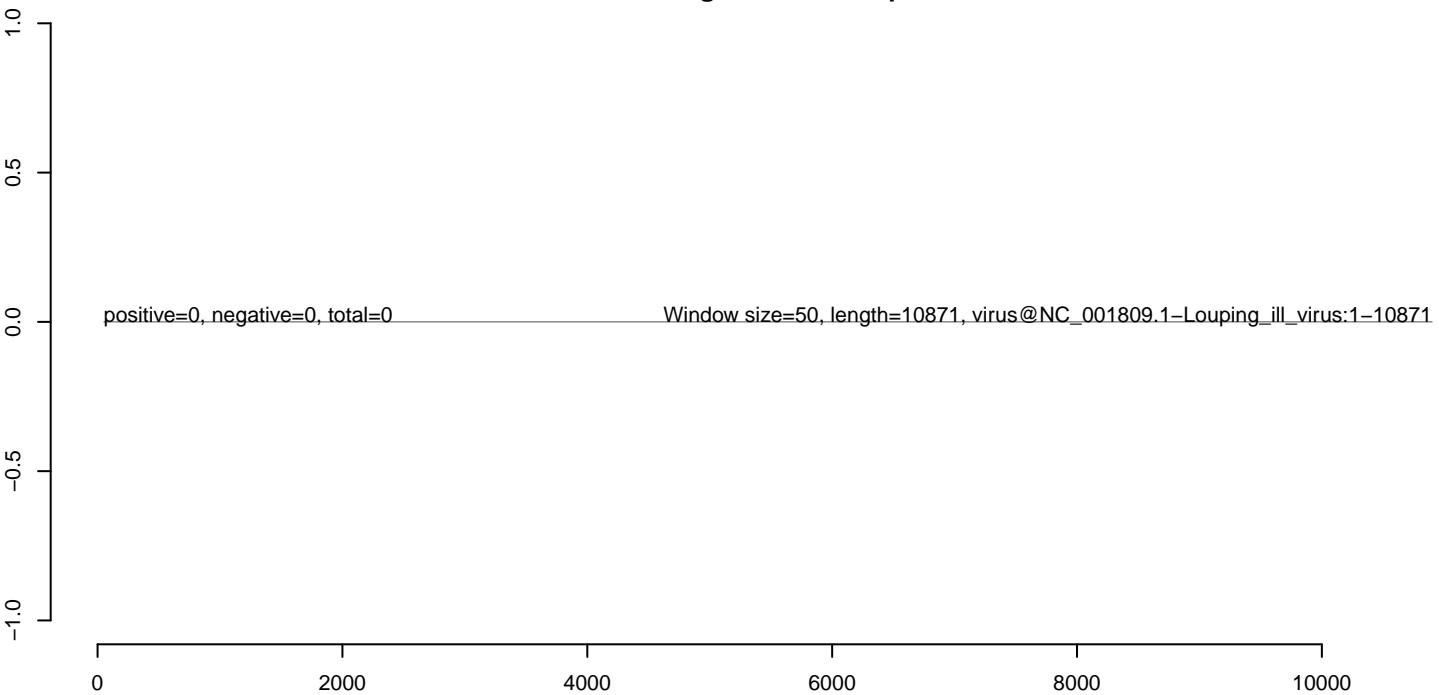
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.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

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.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

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10707, virus@NC\_001475.2-Dengue\_virus\_3:1-10707

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10723, virus@NC\_001474.2-Dengue\_virus\_2:1-10723

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

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

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0 window size=50, length=6065, virus@MH188054.1-Culex\_Y\_virus\_strain\_CYV\_Kern\_partial\_genome:1-6065

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0, size=50, length=13297, virus@MF176298.1-Culex\_mononega-like\_virus\_2\_strain\_mos191gb23464:1-13297

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

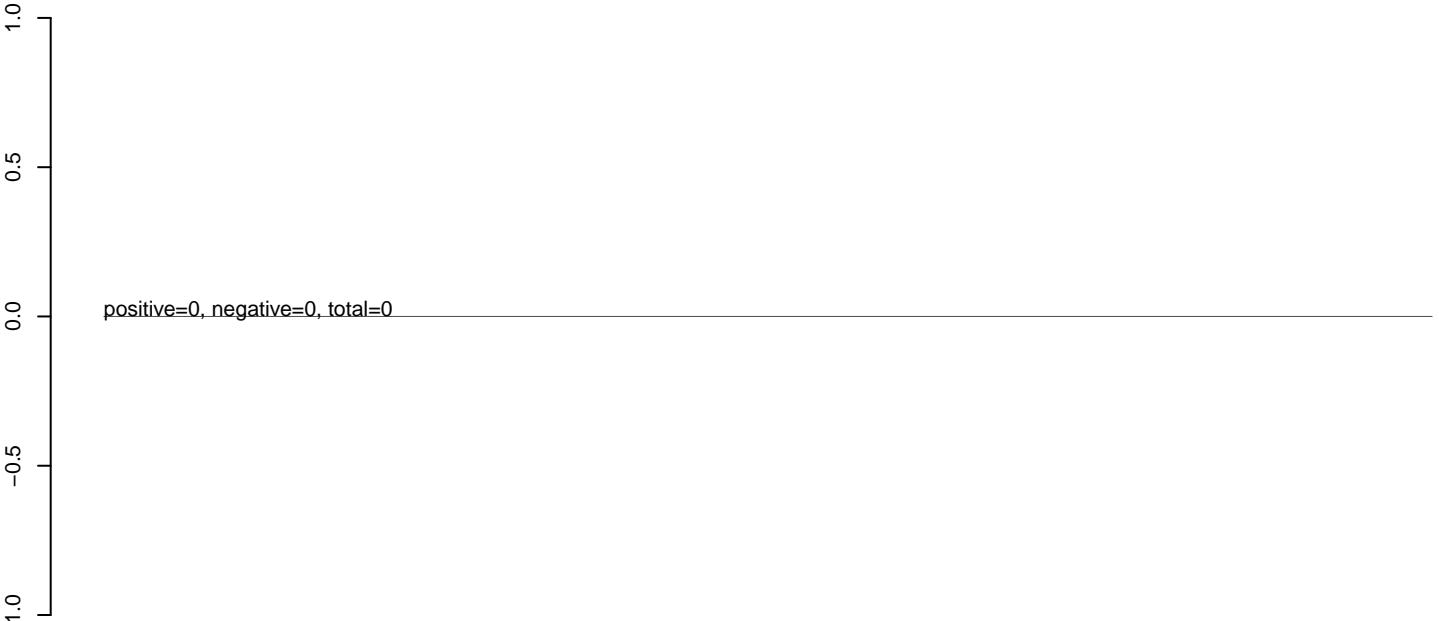
### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

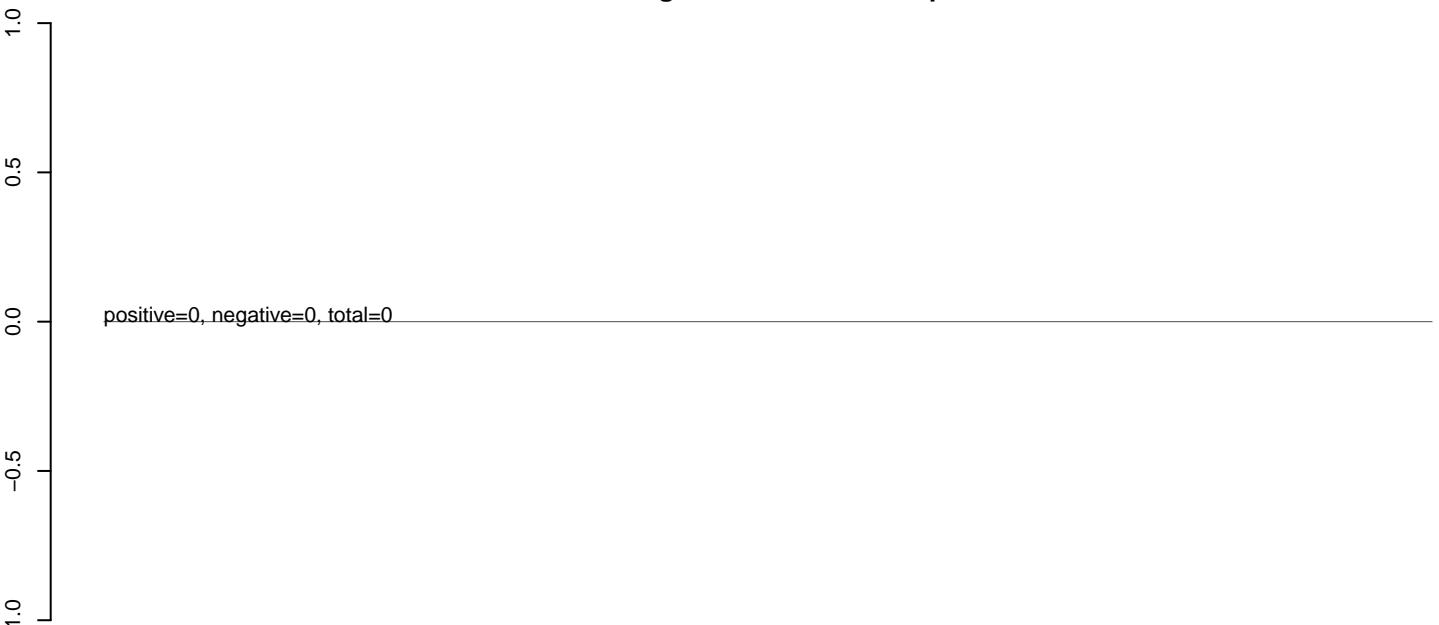
### AeAeg\_Fcanc\_GH.rep

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

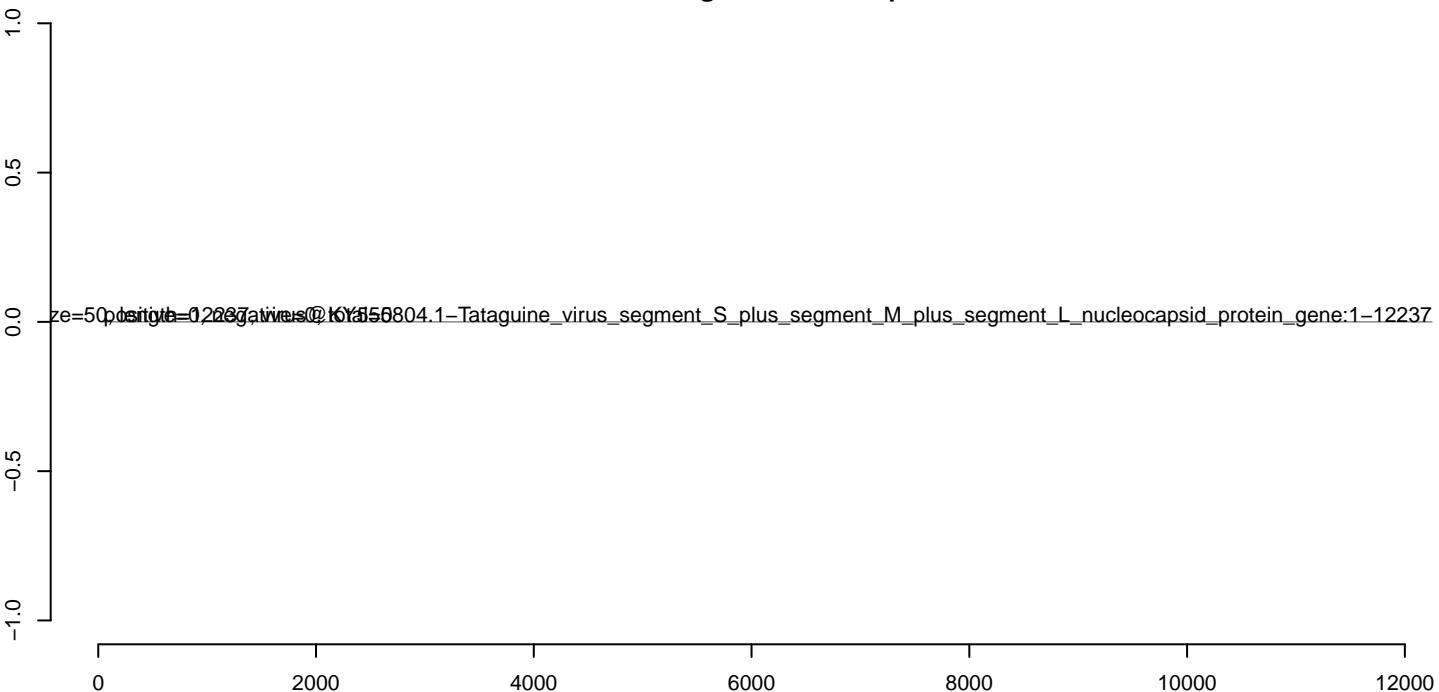
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

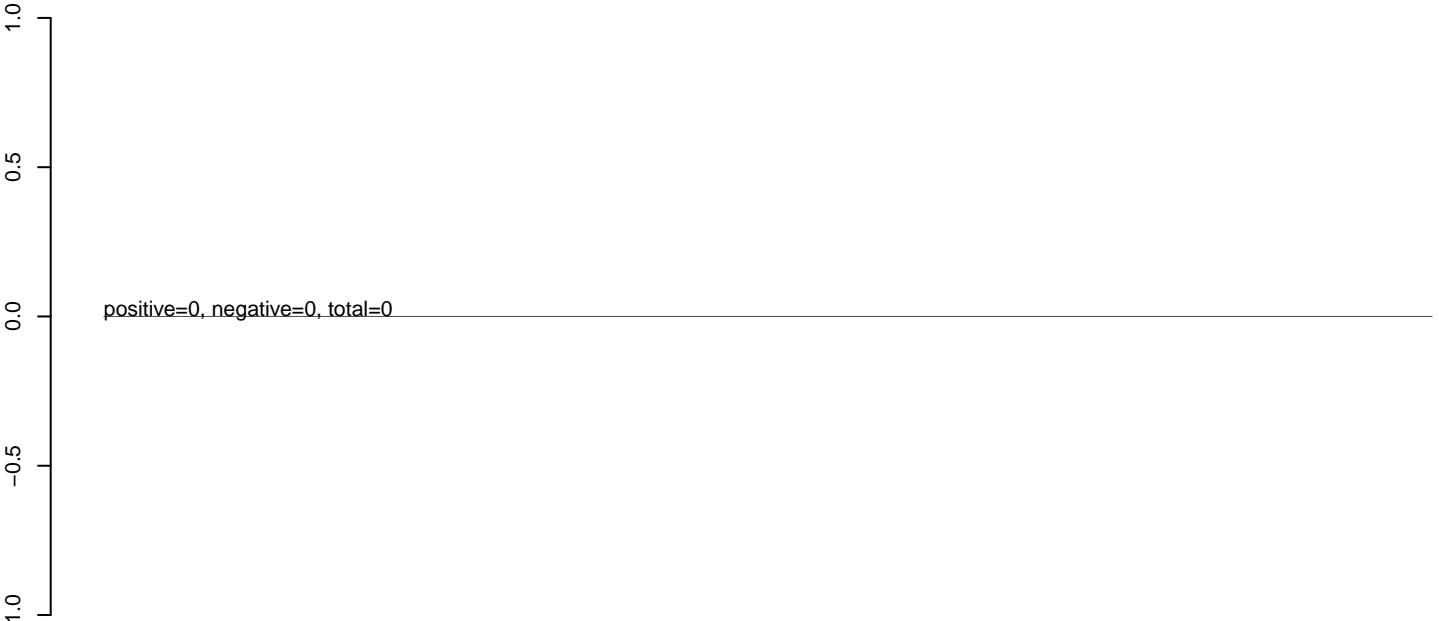
### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

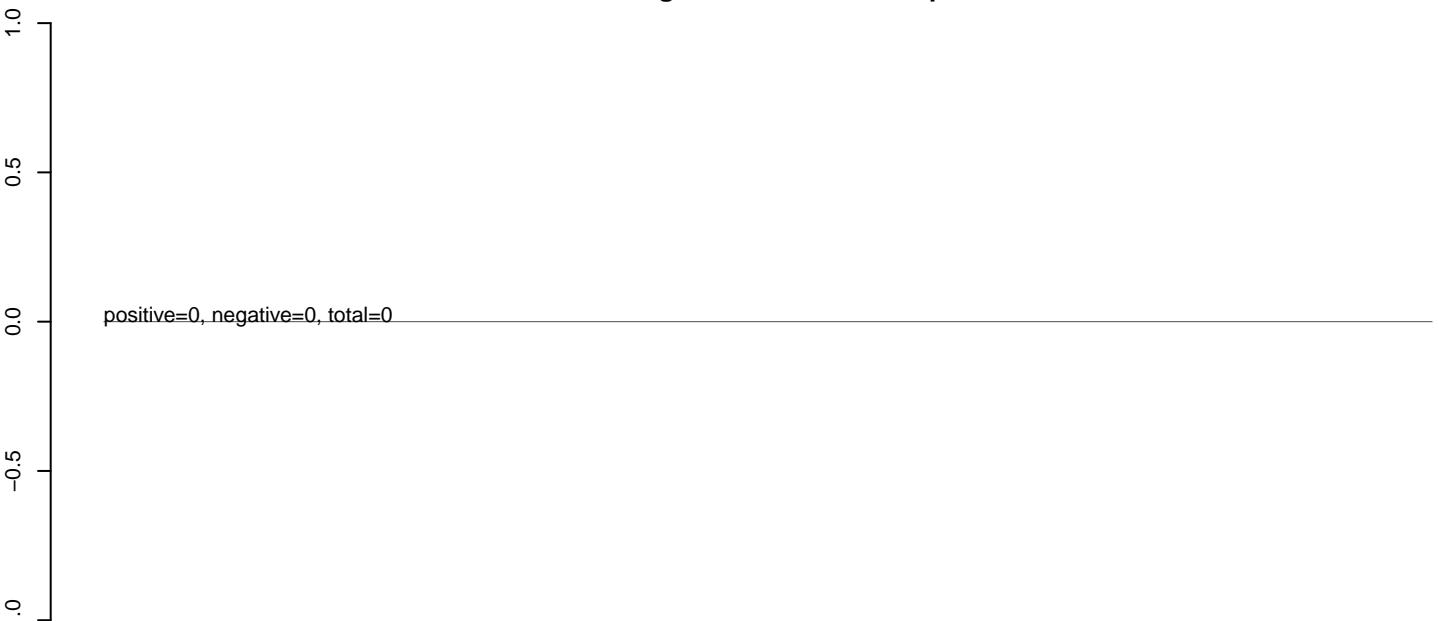
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=6 small\_transmembrane\_protein\_putative\_glycoprotein\_putative\_small\_ZnF\_protein\_and\_RdRp\_genes:1-11652

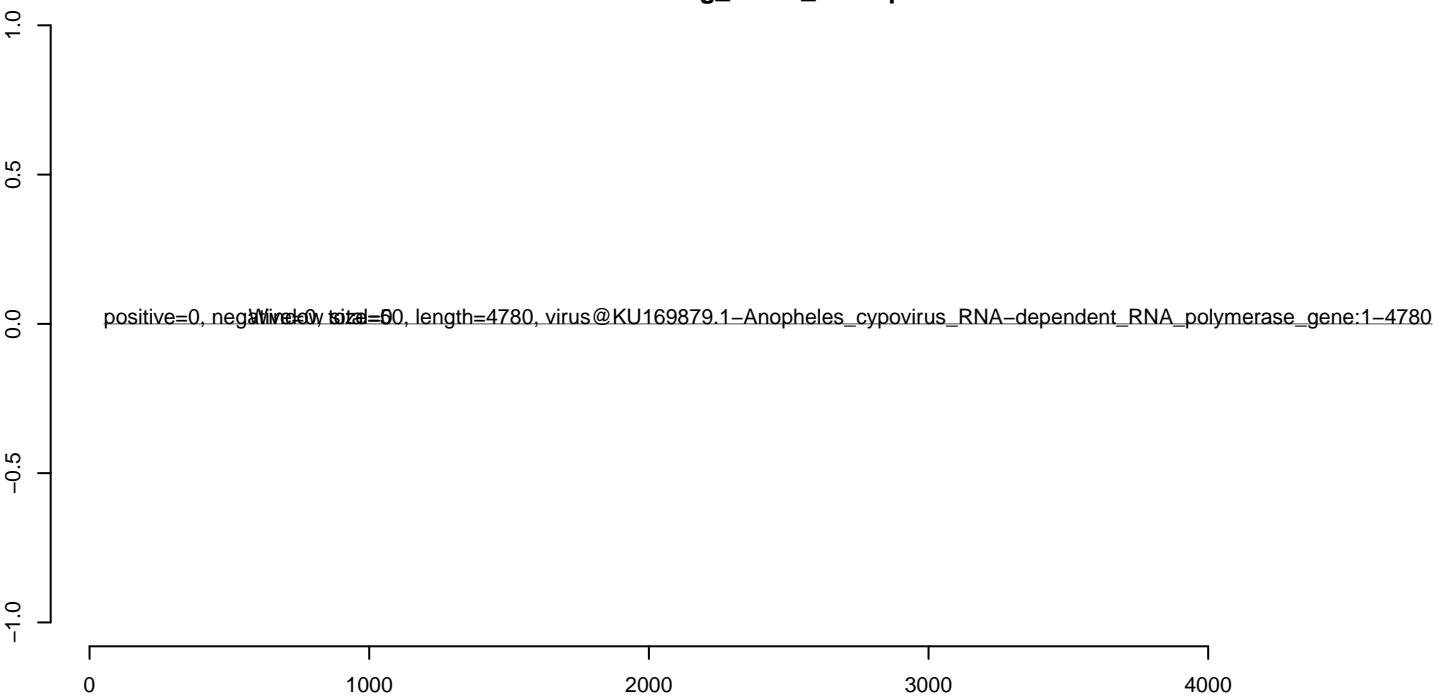
### AeAeg\_Fcanc\_GH.18\_23.rep



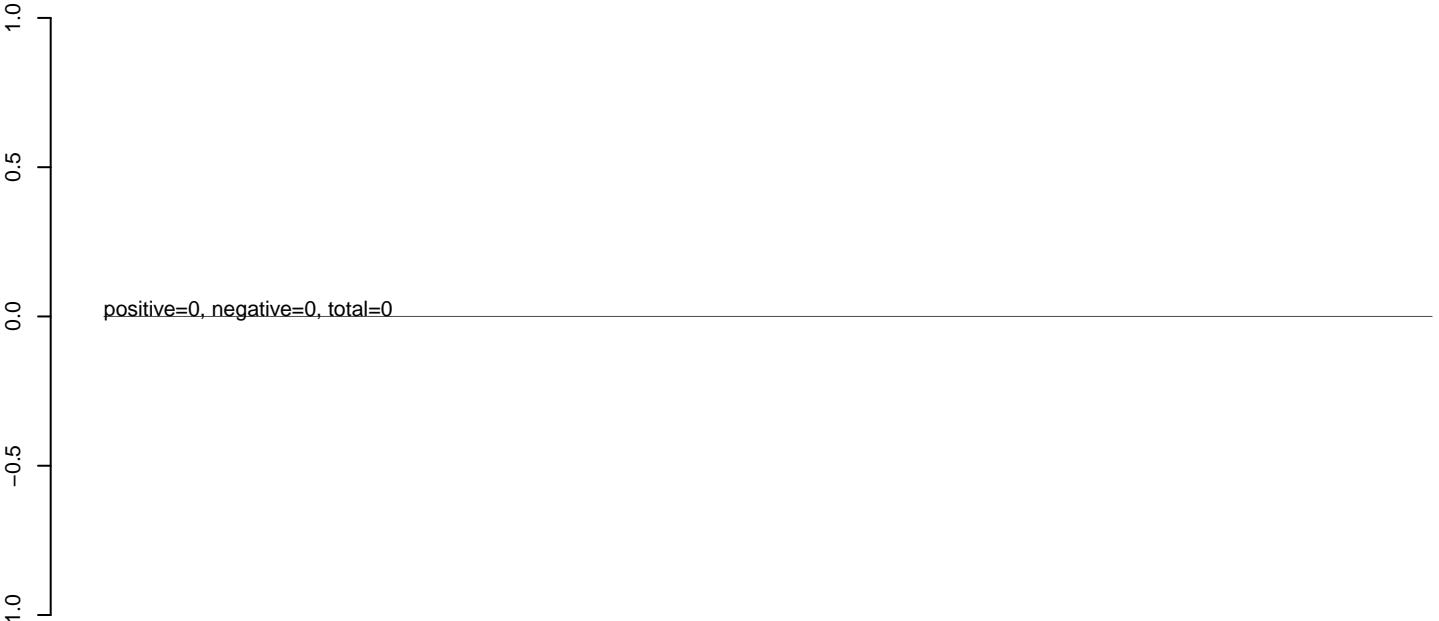
### AeAeg\_Fcanc\_GH.24\_35.rep



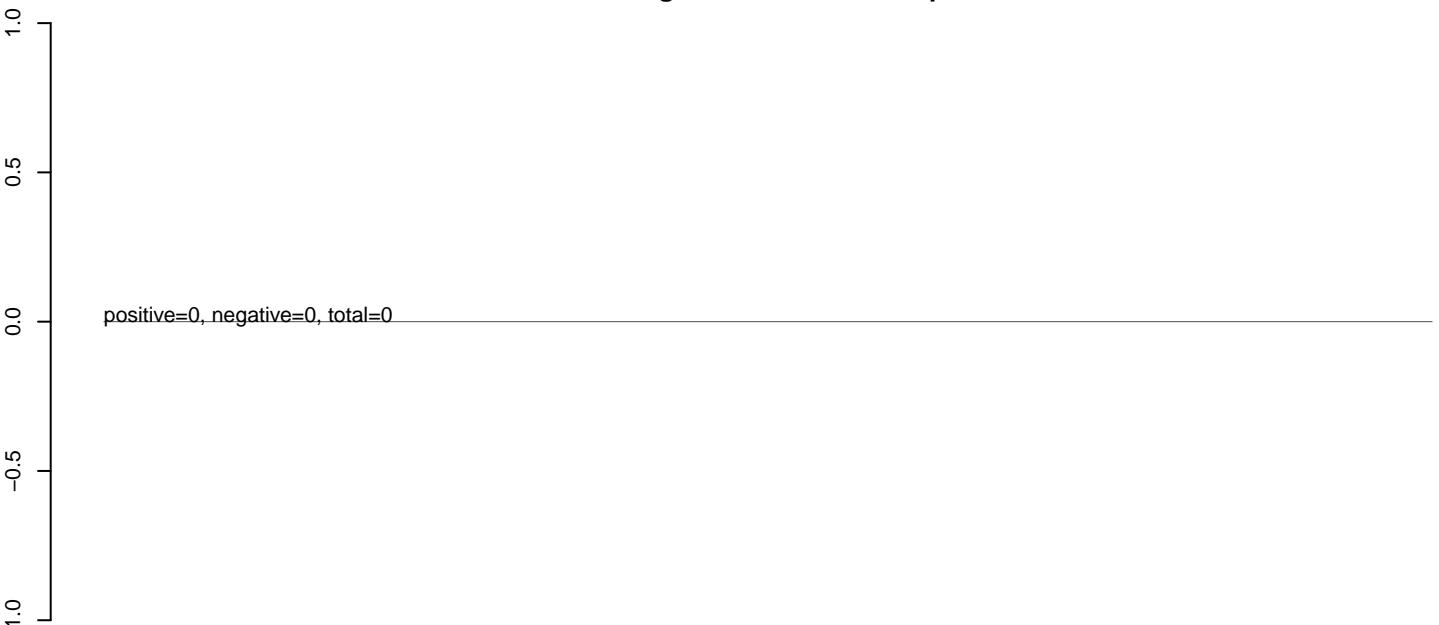
### AeAeg\_Fcanc\_GH.rep



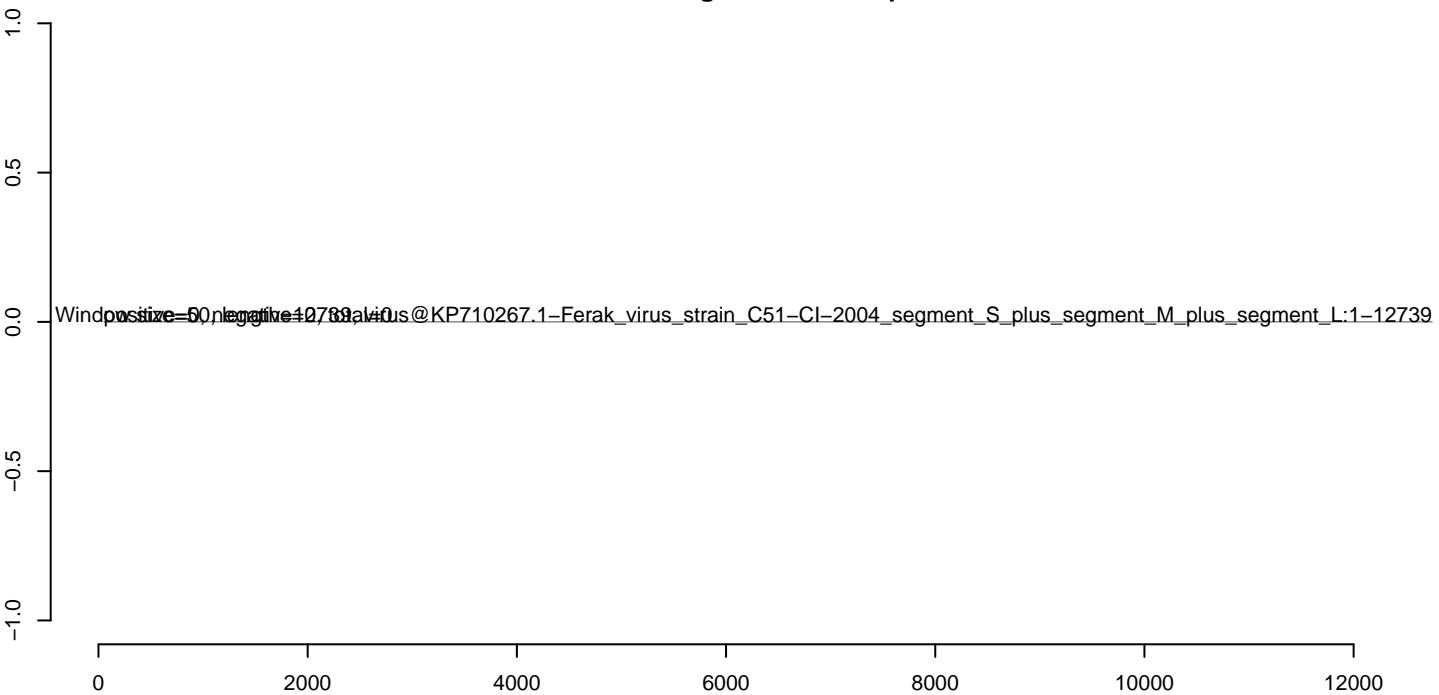
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10735, virus@KM225265.1-Torres\_virus\_strain\_TS5273:1-10735

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

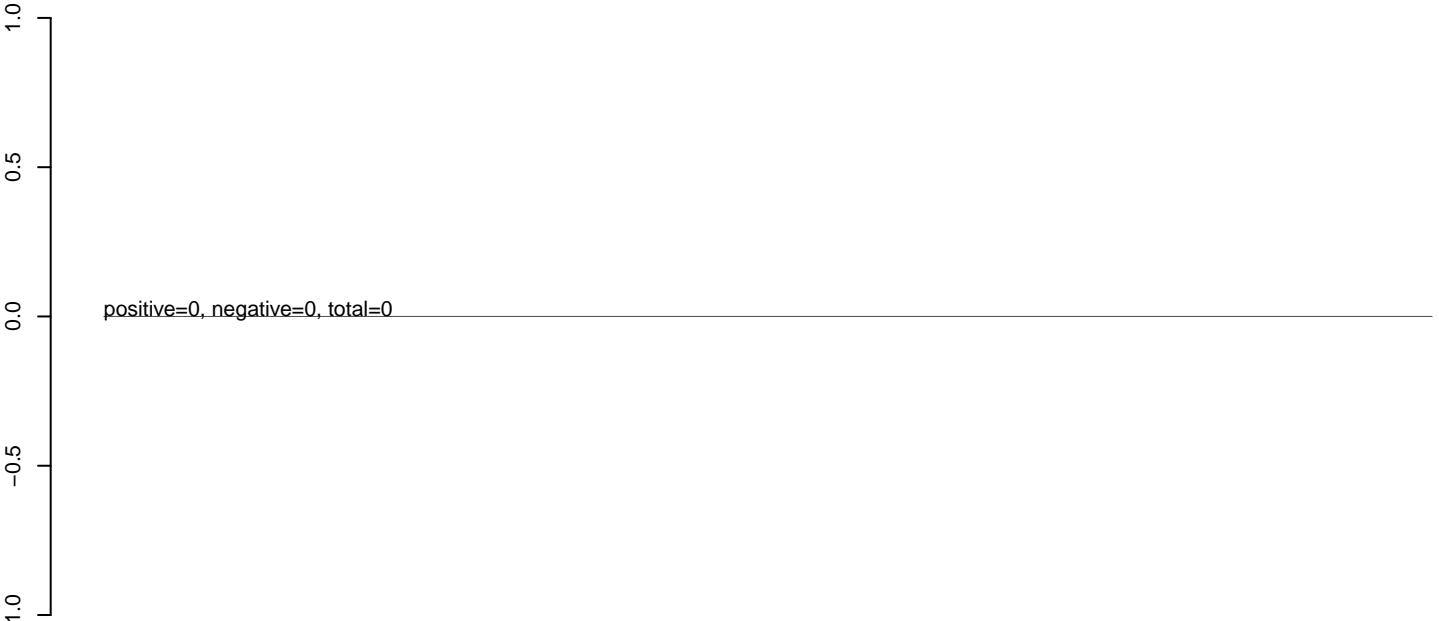
### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

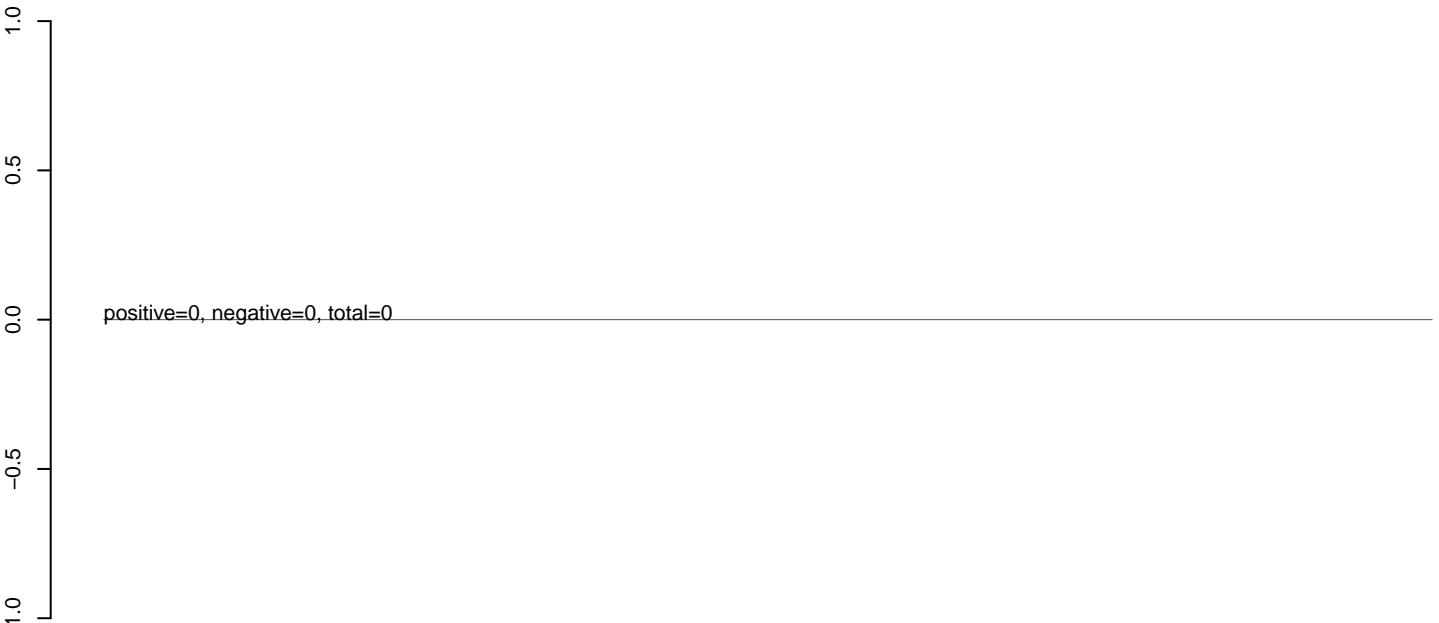
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, size=50, length=4773, virus@KJ632942.1-Mosinovirus\_strain\_C36\_Cl-2004\_segment\_RNA\_1\_and\_RNA\_2:1-4773

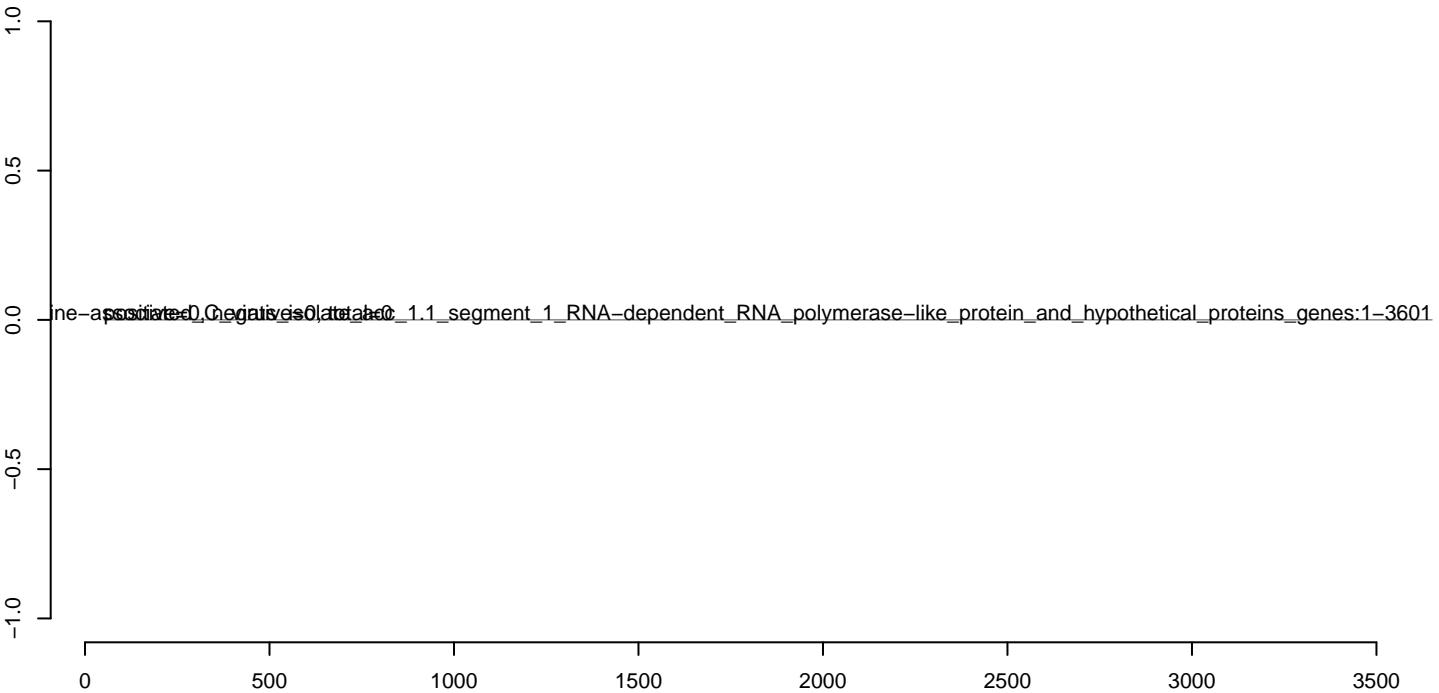
### AeAeg\_Fcanc\_GH.18\_23.rep



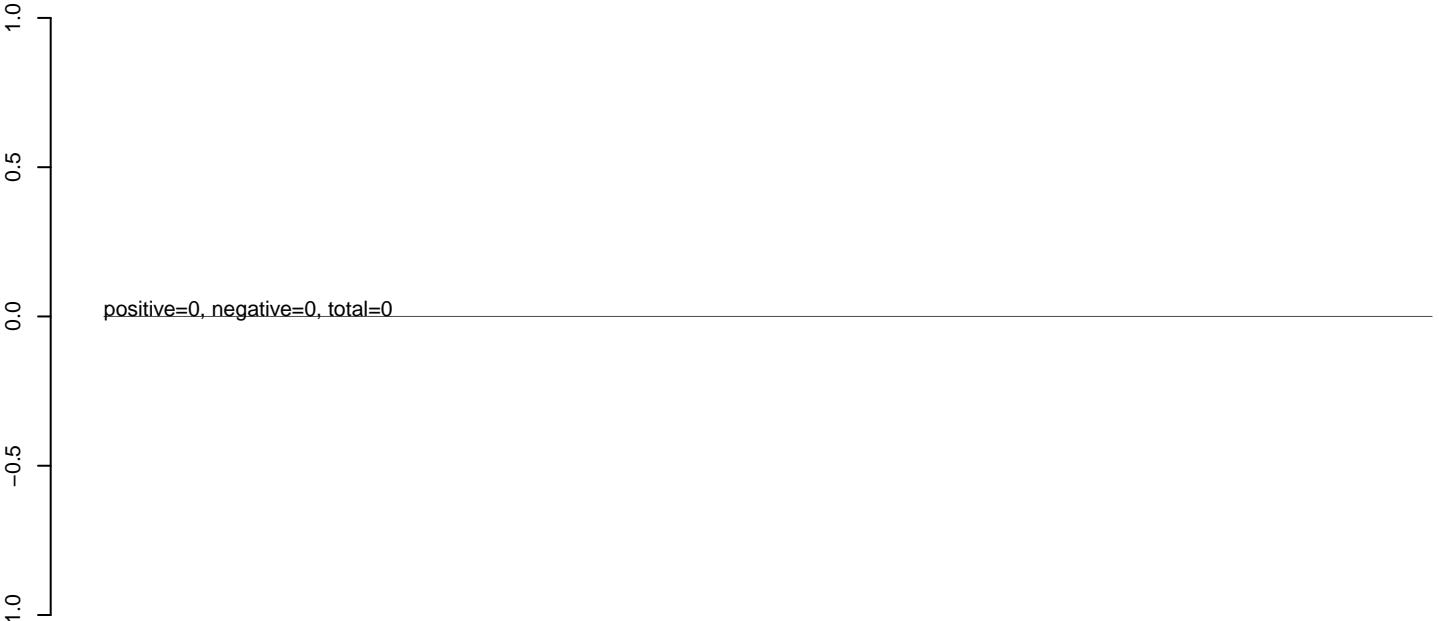
### AeAeg\_Fcanc\_GH.24\_35.rep



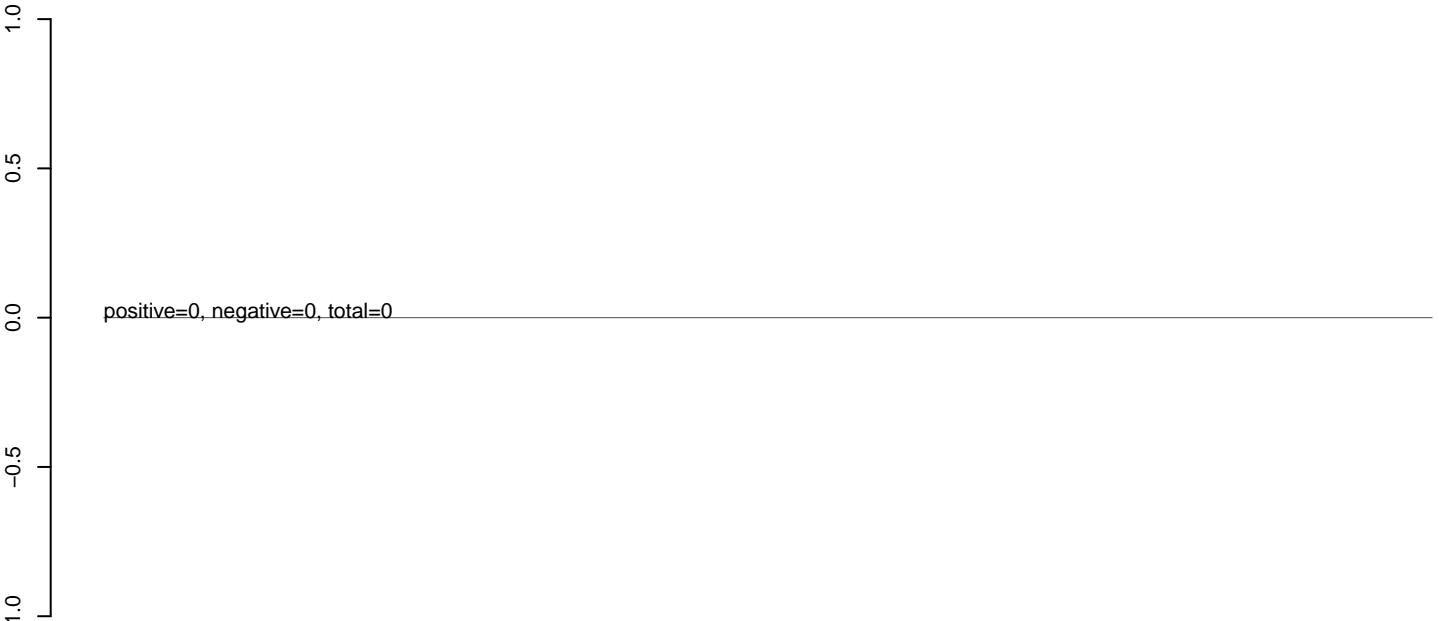
### AeAeg\_Fcanc\_GH.rep



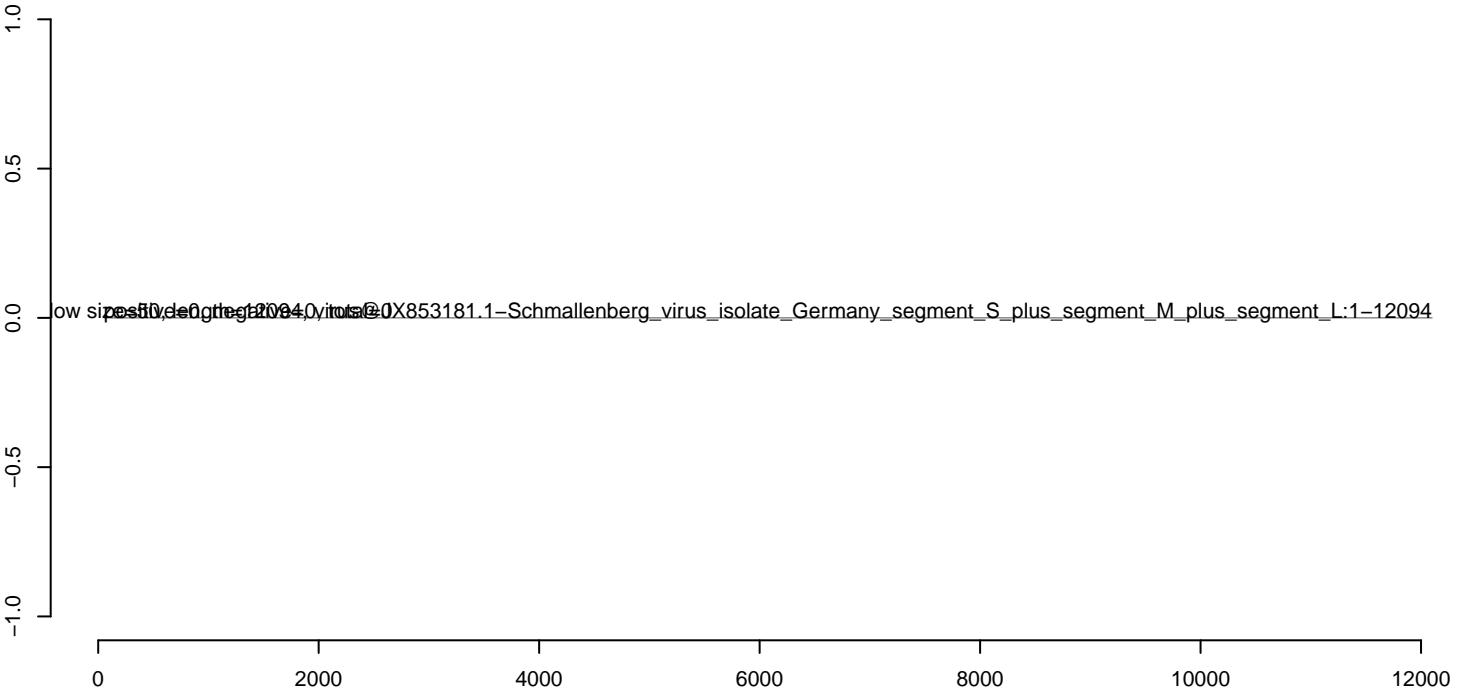
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10841, virus@JQ308187.1–Culex\_flavivirus\_isolate\_HLD105:1–10841

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10841, virus@HQ678513.1-Culex\_flavivirus\_isolate\_H0901:1-10841

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

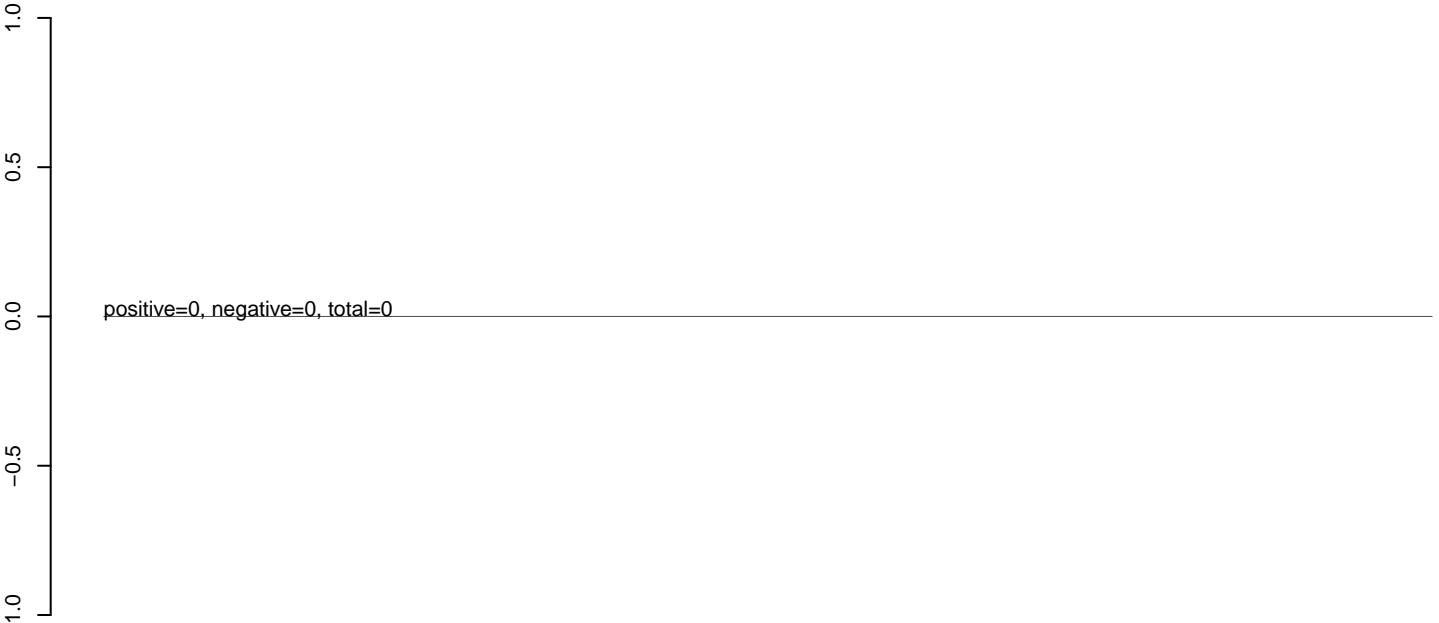
### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

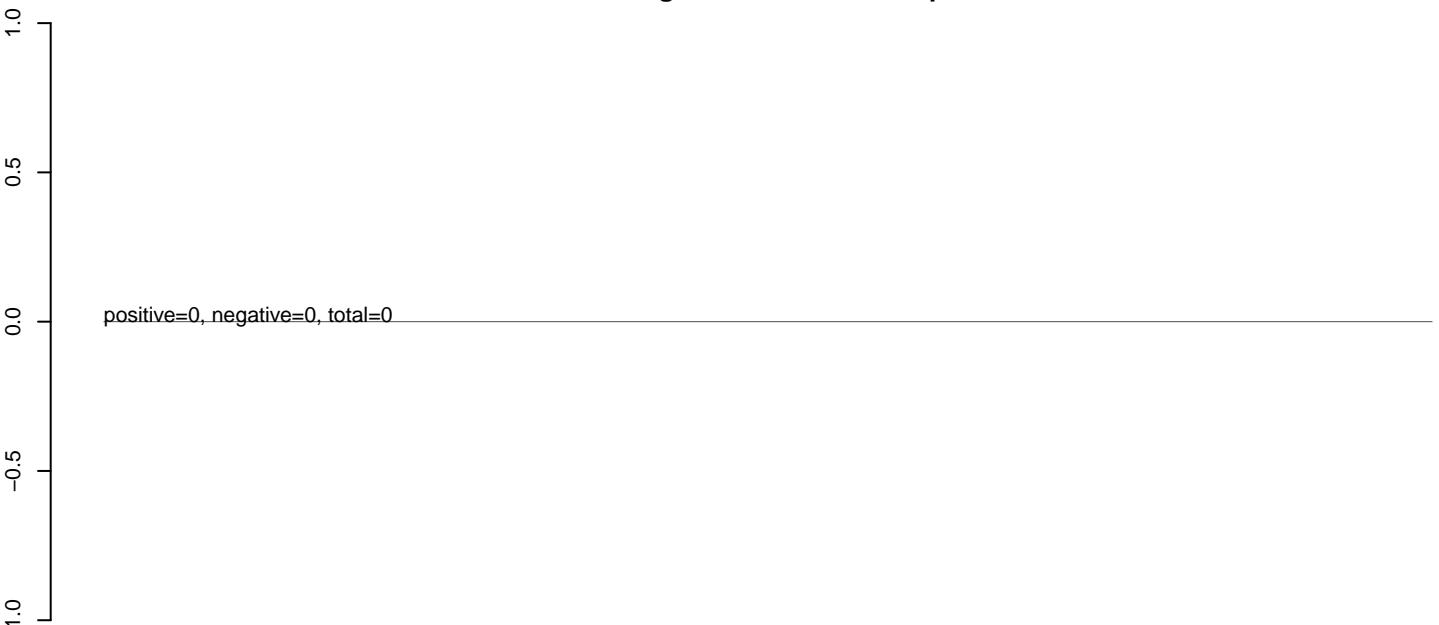
### AeAeg\_Fcanc\_GH.rep

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

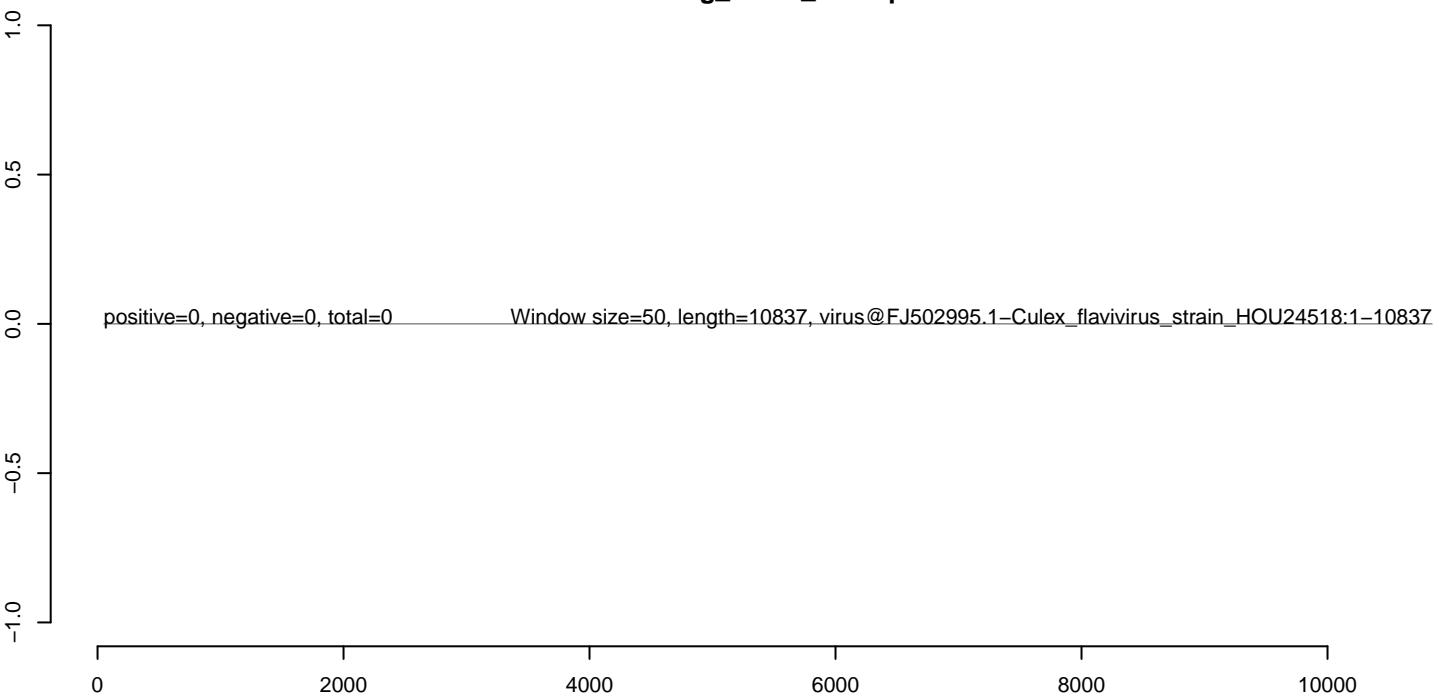
### AeAeg\_Fcanc\_GH.18\_23.rep



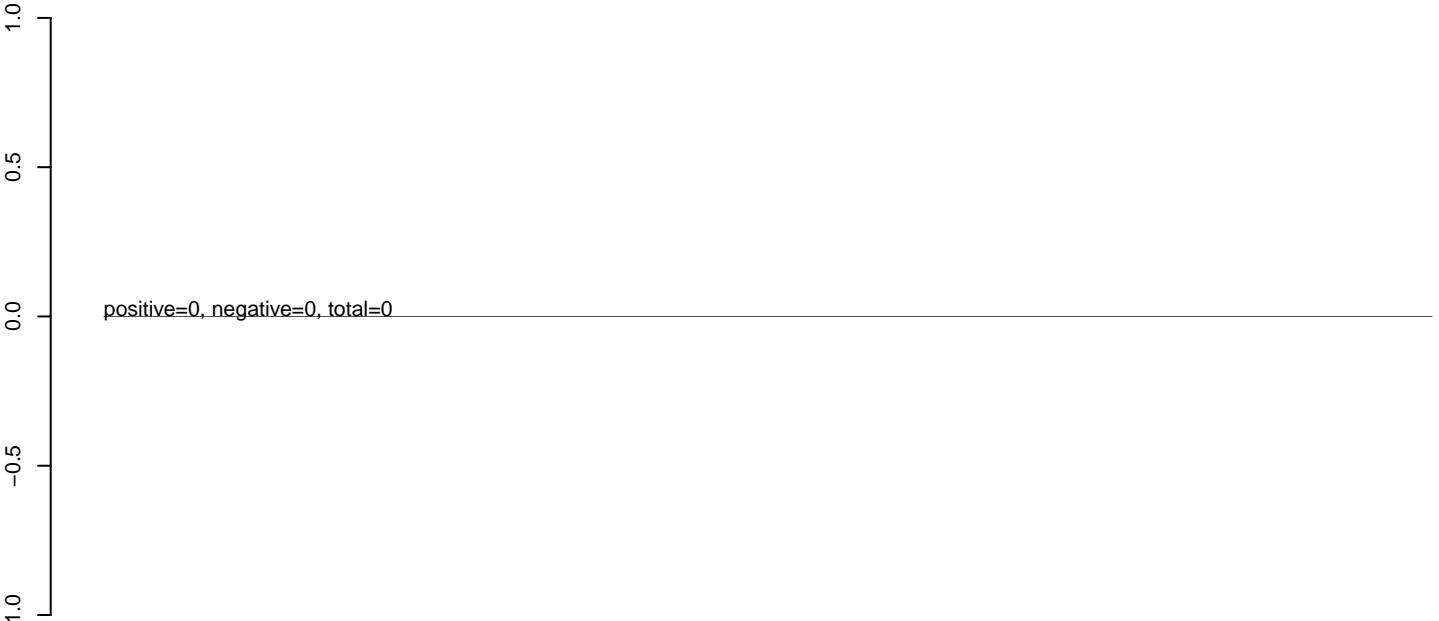
### AeAeg\_Fcanc\_GH.24\_35.rep



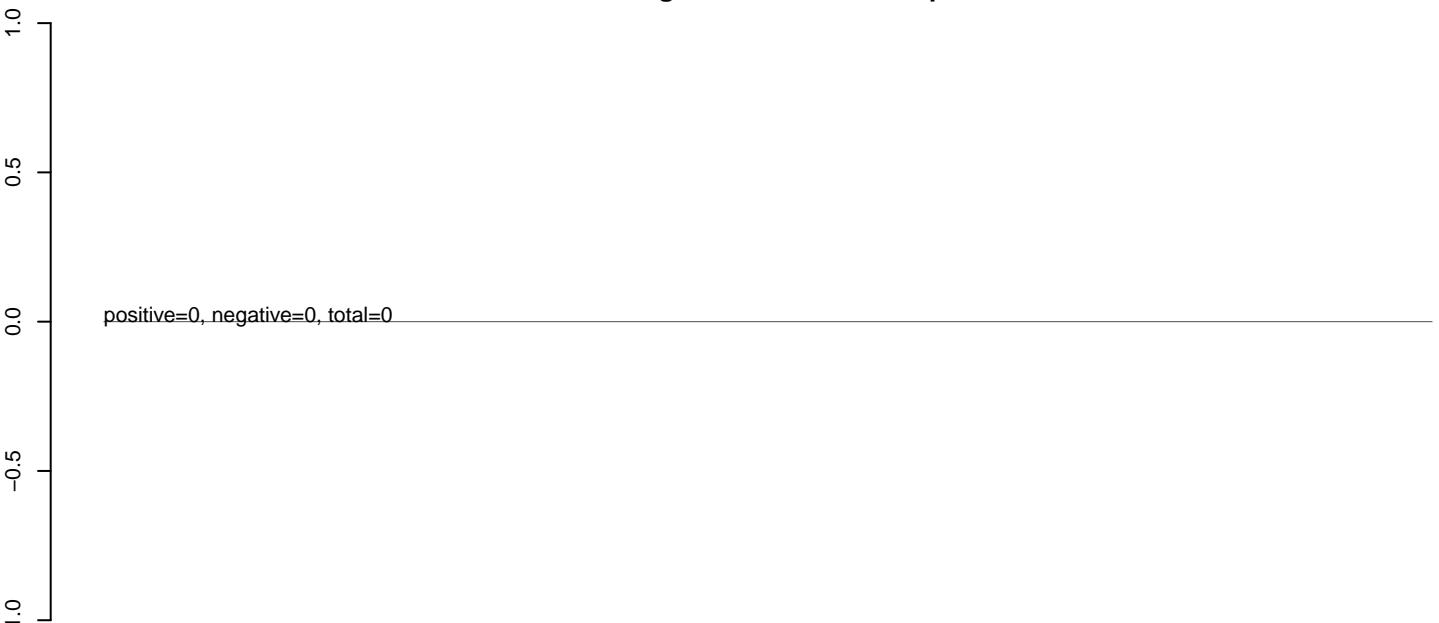
### AeAeg\_Fcanc\_GH.rep



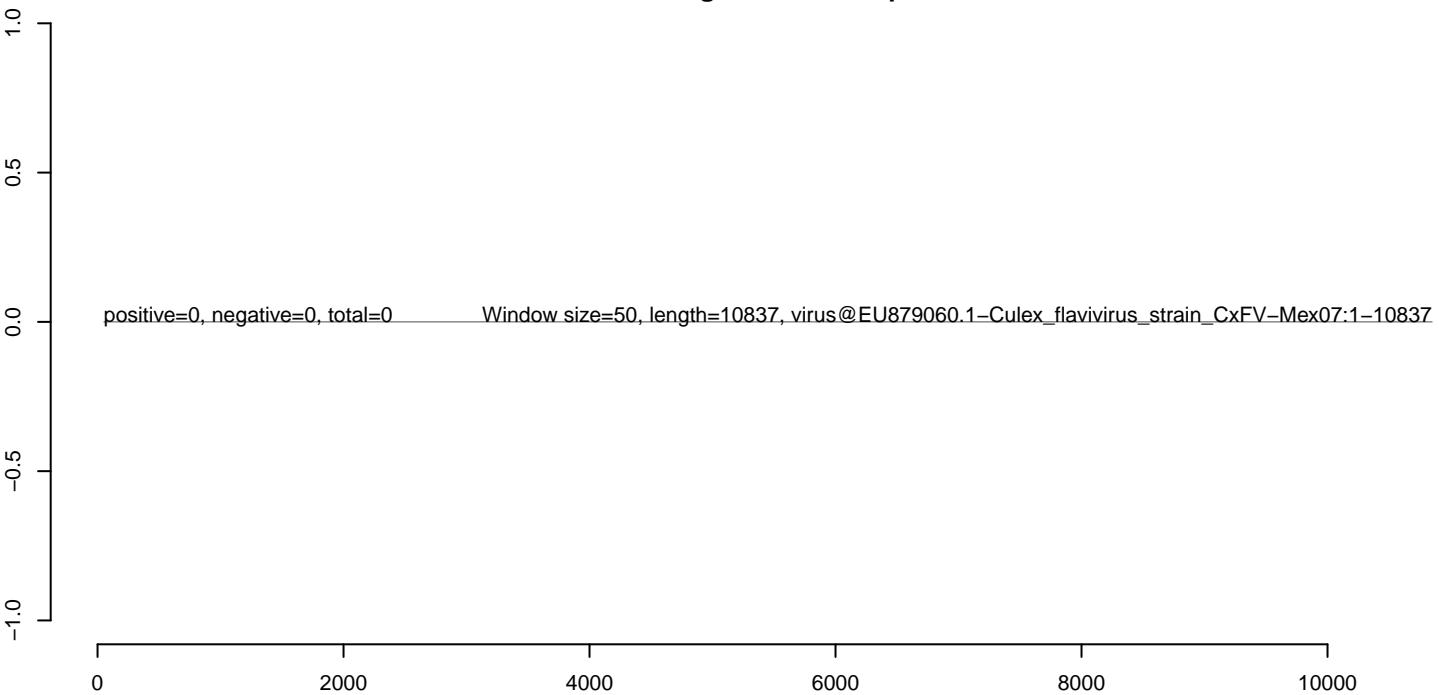
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

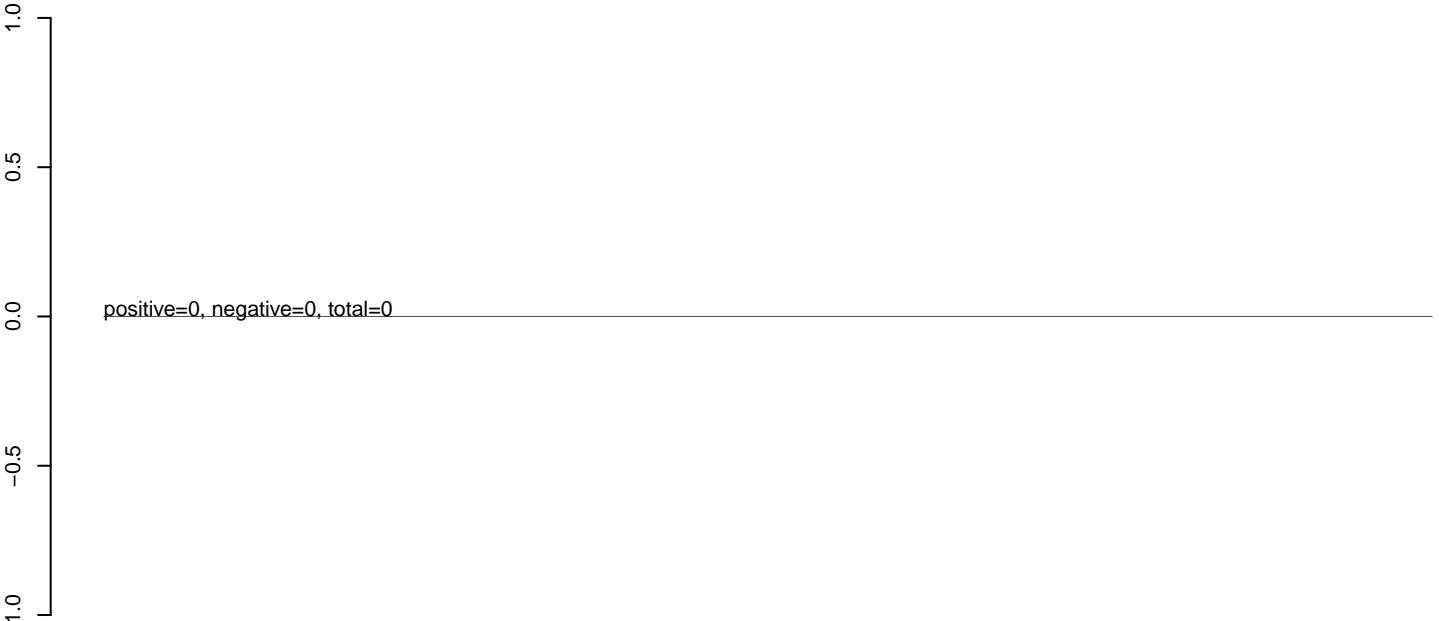
positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

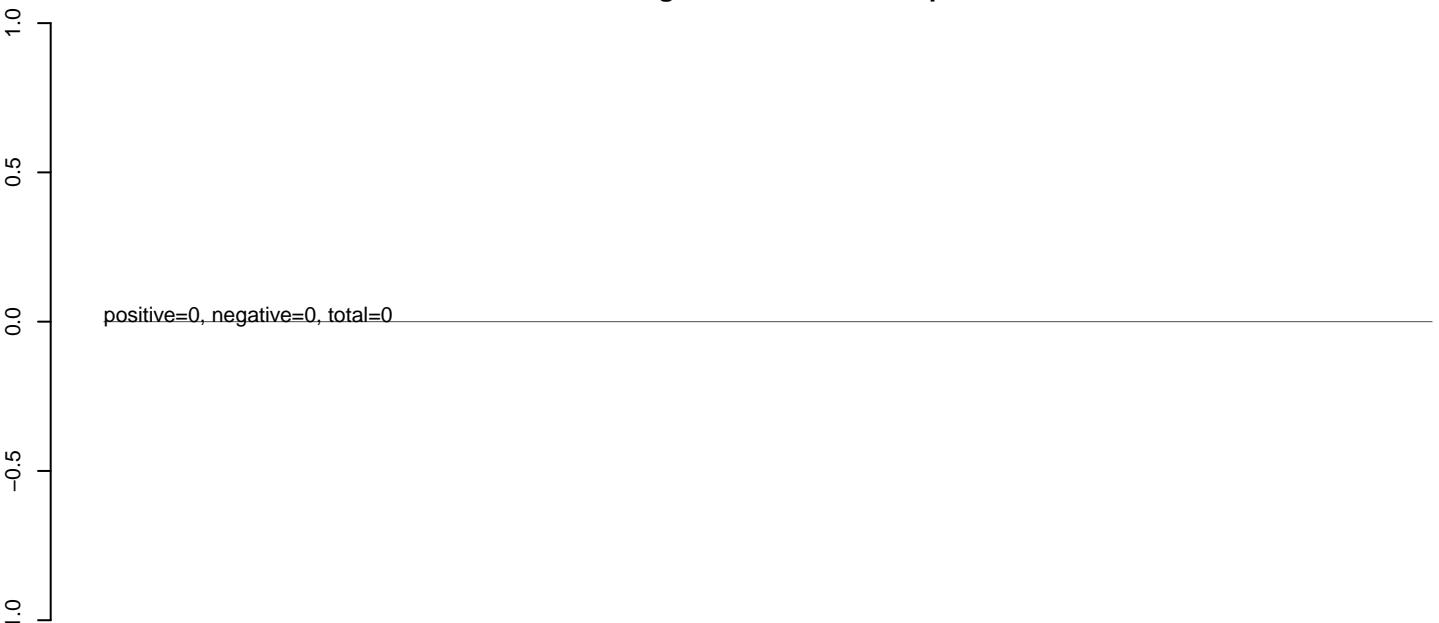
positive=0, negative=0, total=0, length=10302, virus@EU082200.2-Koutango\_virus\_strain\_Dak\_Ar\_D\_5443\_polyprotein\_gene:1-10302

0 2000 4000 6000 8000 10000

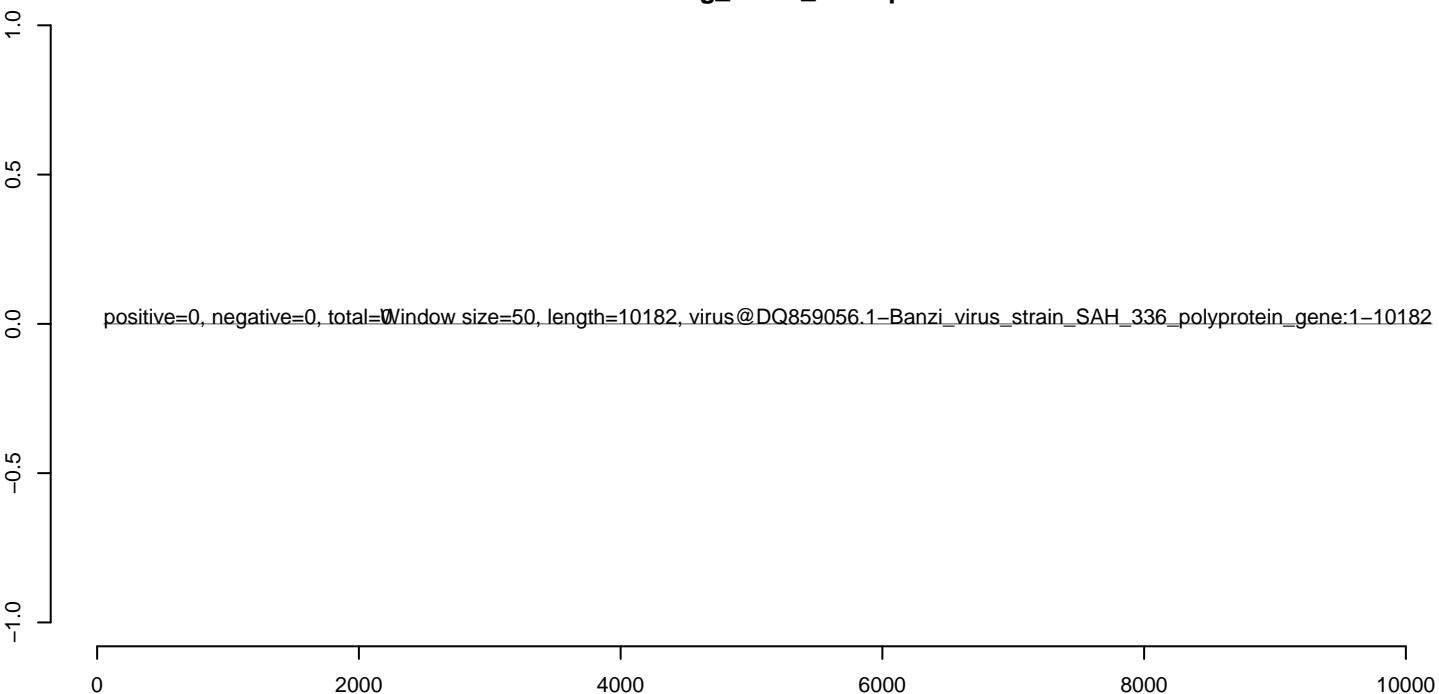
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

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

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=10794, virus@AY632542.4–Rocio\_virus\_strain\_SPH\_34675:1–10794

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=11344, virus@AF075256.1-Pixuna\_virus\_strain\_Pixuna\_BeAr\_35645:1-11344

0 2000 4000 6000 8000 10000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0

Window size=50, length=11530, virus@AF075254.1-Tonate\_virus\_strain\_Tonate\_CaAn\_410d:1-11530

0 2000 4000 6000 8000 10000 12000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

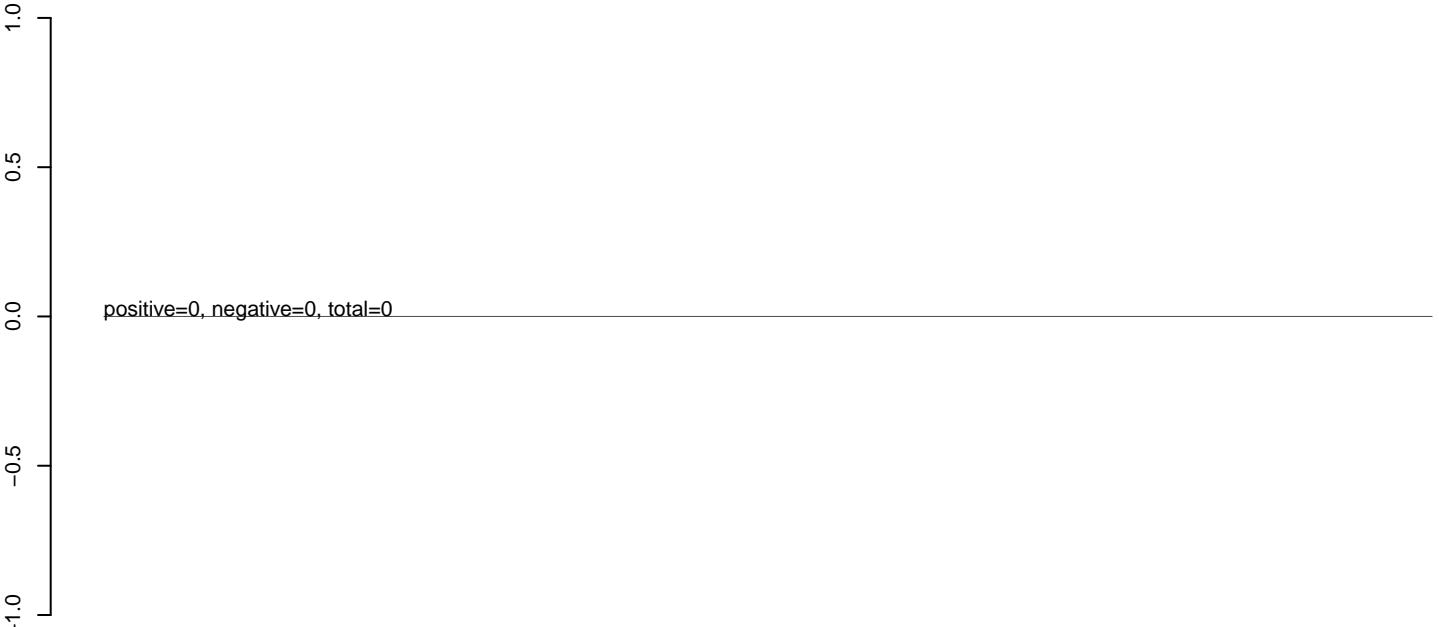
positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

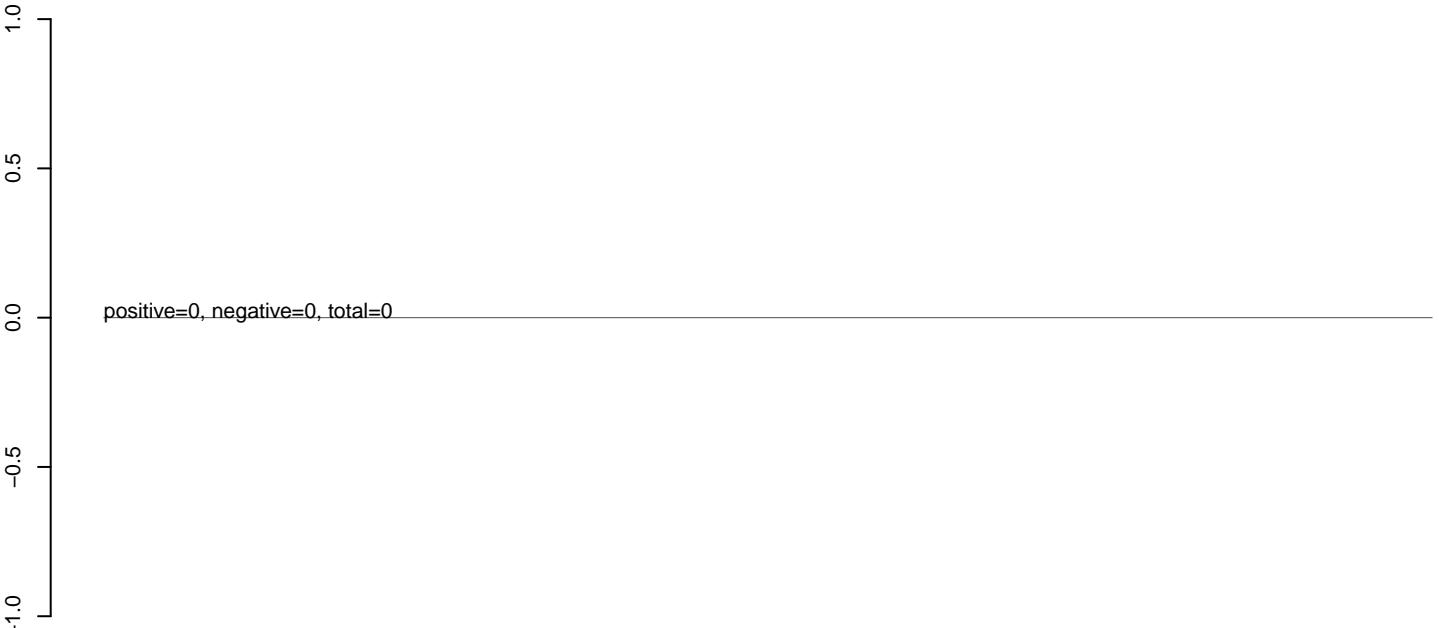
positive=0, negative=0, total=0    Window size=50, length=11391, virus@AF075253.1-Mucambo\_virus\_strain\_Mucambo\_BeAn\_8:1-11391

0 2000 4000 6000 8000 10000

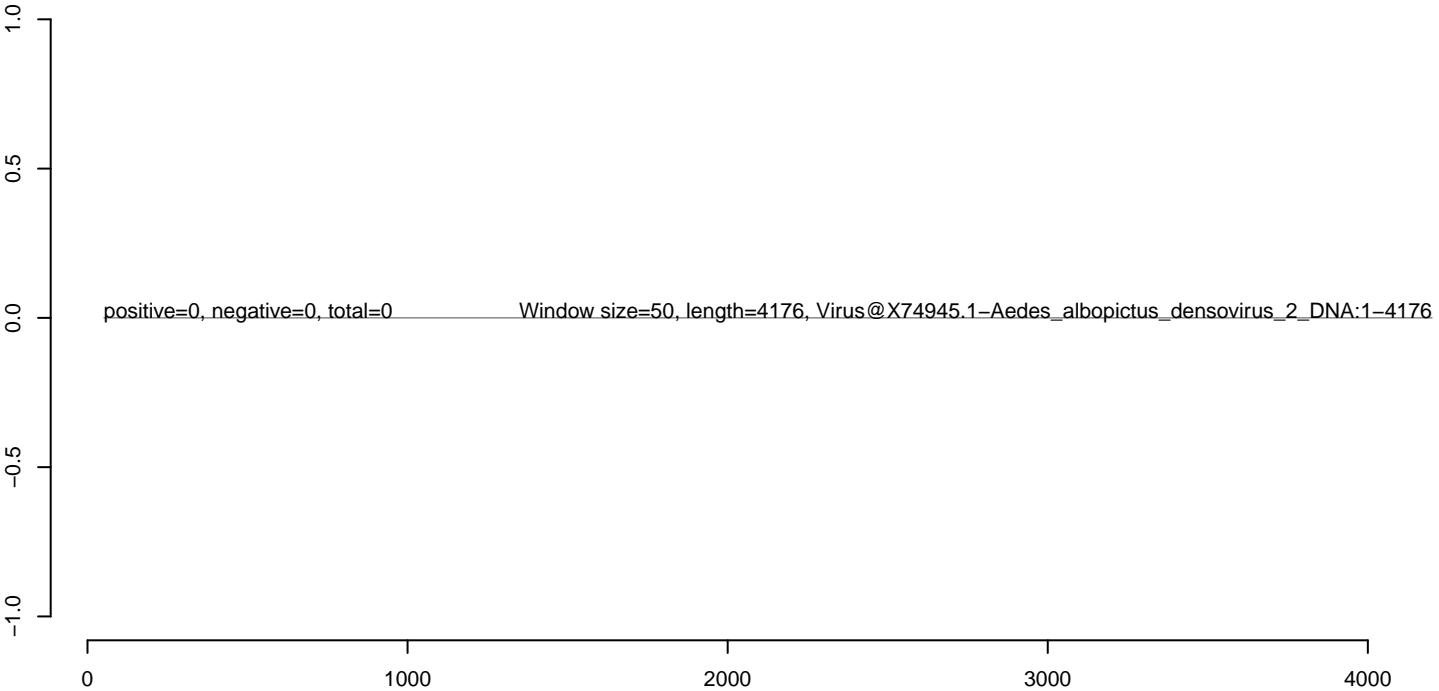
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

Window size: 50, given length: 3937, Virus@MH550148.1-Aedes\_albopictus\_densovirus\_isolate\_NC01\_L3\_1\_trimmed\_contig\_5889-RC:1-3937

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0    Window.size=50, length=3805, Virus@MH188045.1-Culex\_densovirus\_strain\_CDenv\_Madera-4:1-3805

0 1000 2000 3000 4000

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0    Window.size=50, length=3991, Virus@MH188044.1-Culex\_densovirus\_strain\_CDenv\_Madera-3:1-3991

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.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

### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

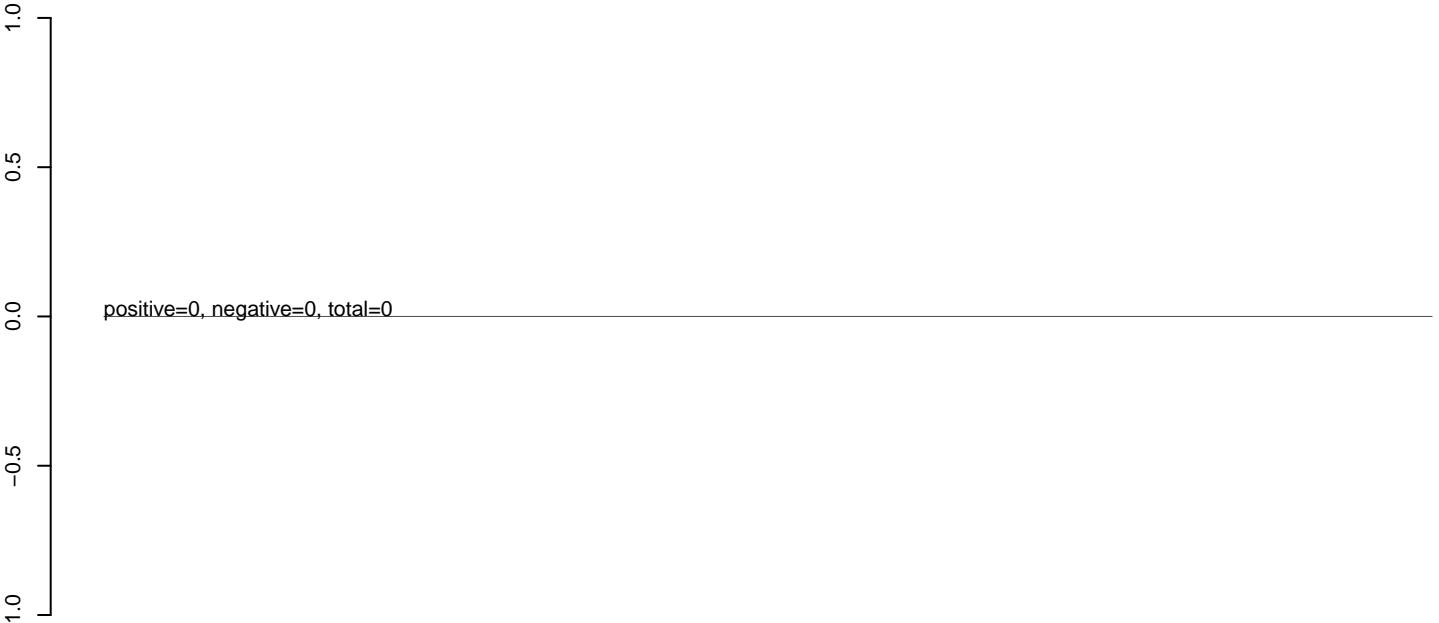
### AeAeg\_Fcanc\_GH.24\_35.rep

positive=0, negative=0, total=0

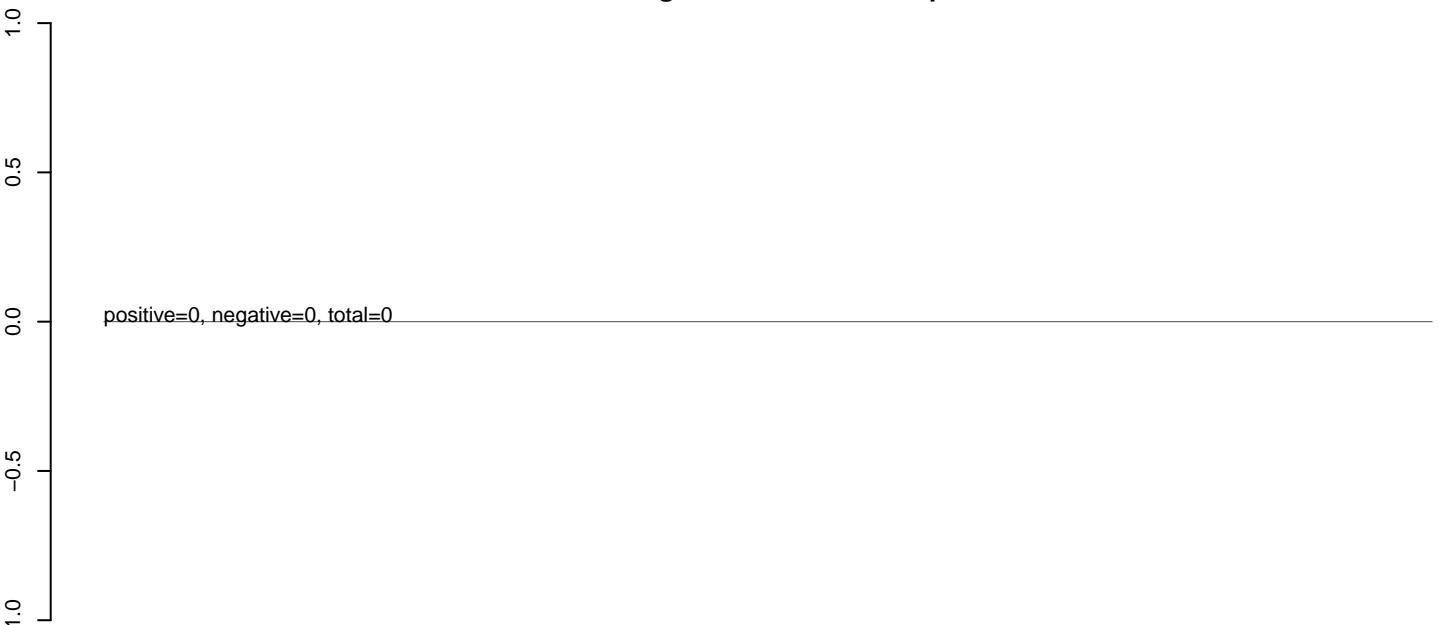
### AeAeg\_Fcanc\_GH.rep

positive=0, negative=0, total=0 Window.size=50, length=3978, Virus@M37899.1-Aedes\_aegypti\_densovirus\_1\_genomic\_sequence:1-3978

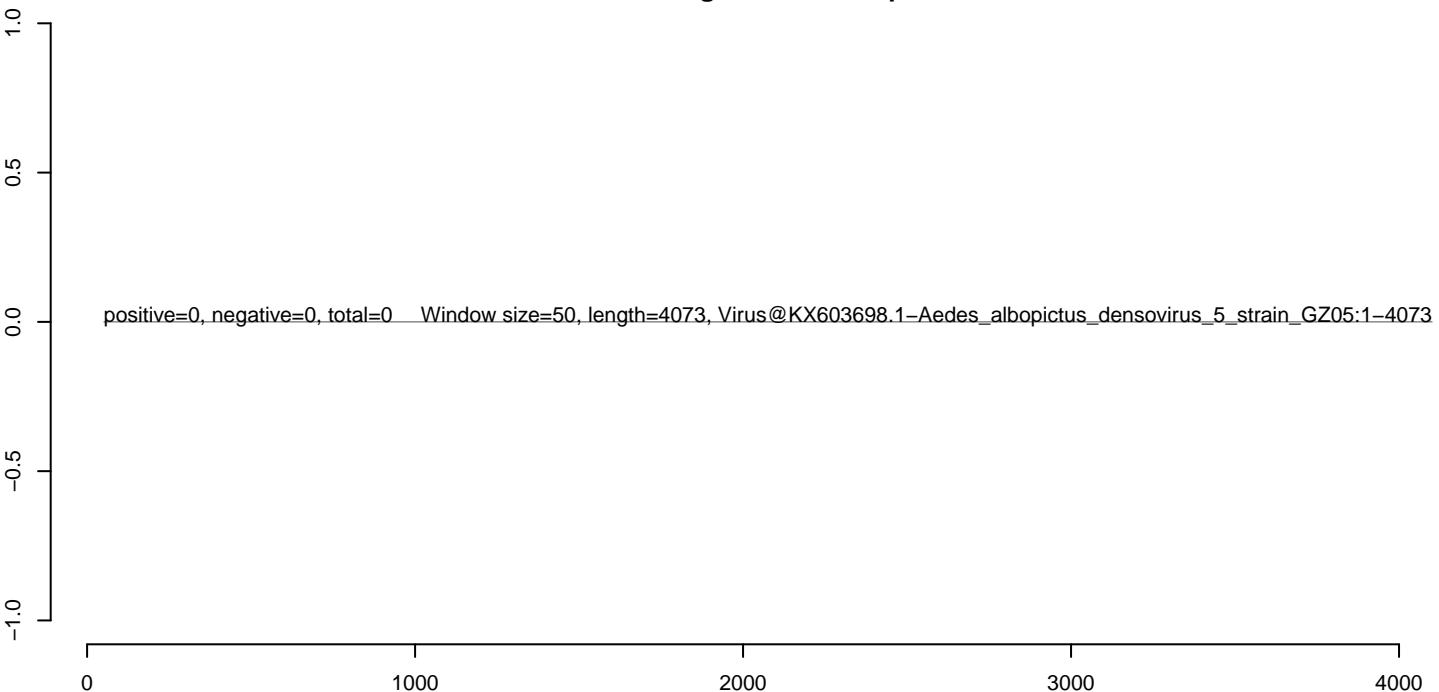
### AeAeg\_Fcanc\_GH.18\_23.rep



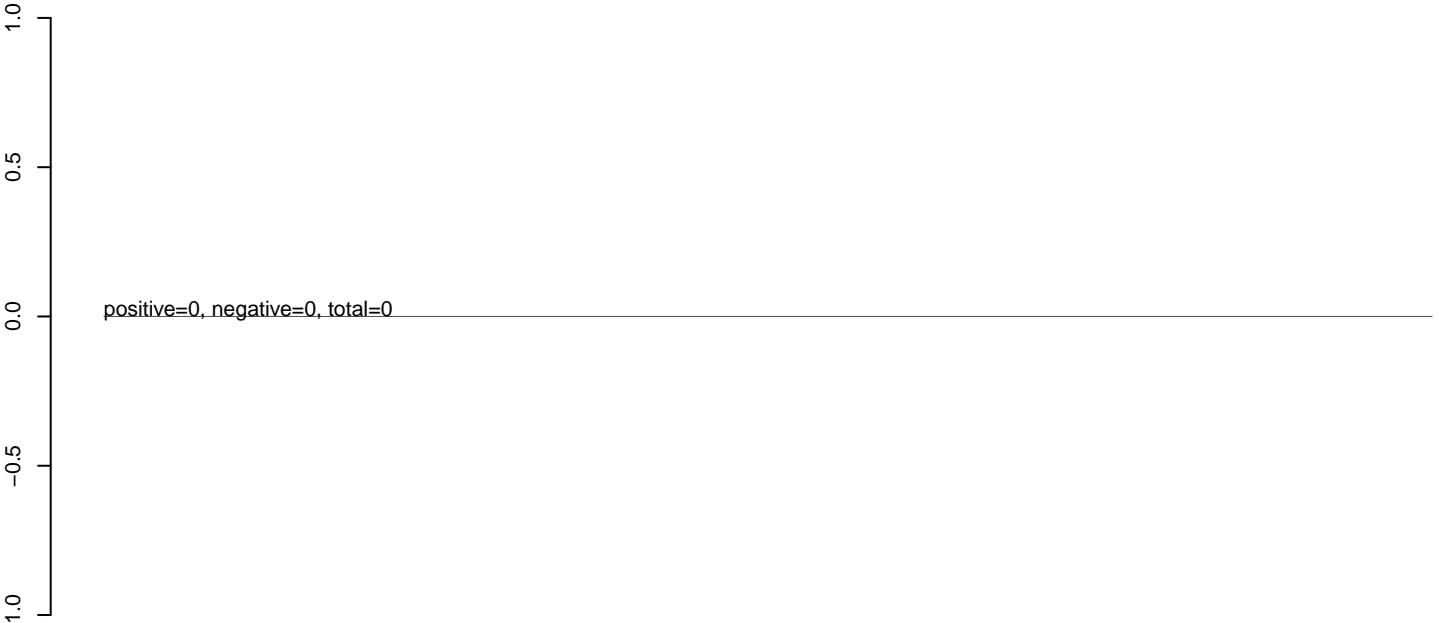
### AeAeg\_Fcanc\_GH.24\_35.rep



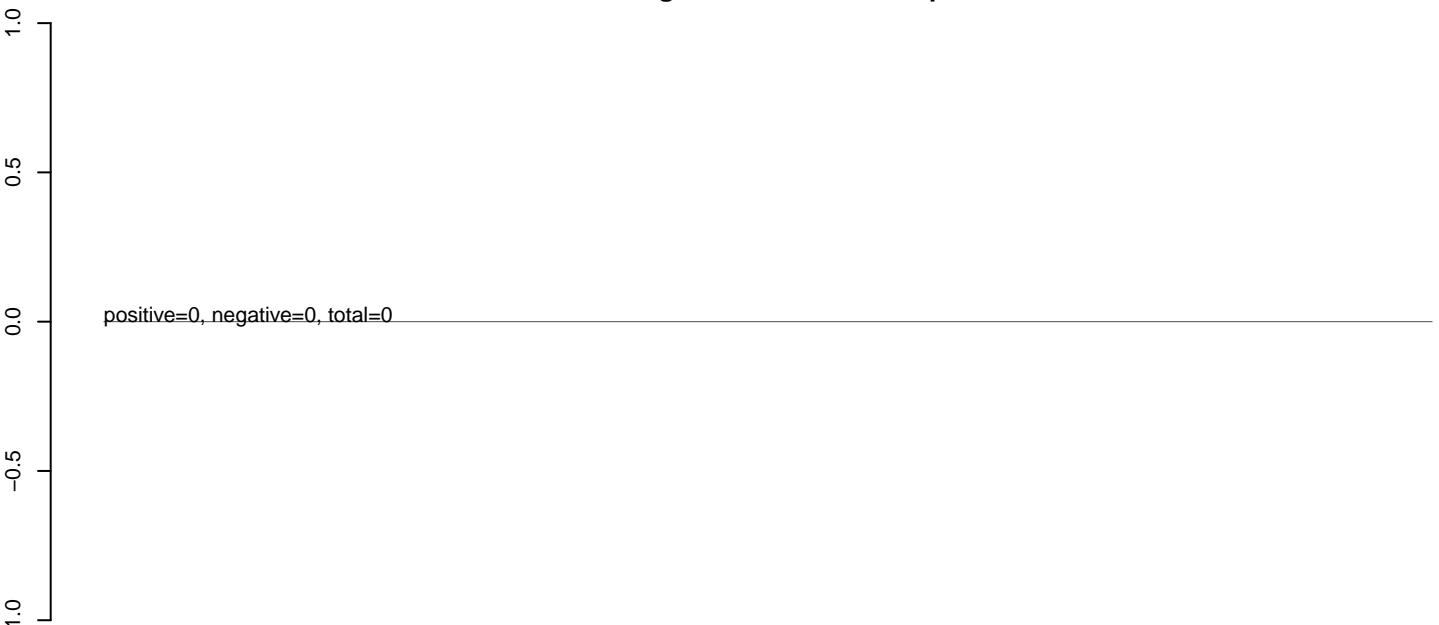
### AeAeg\_Fcanc\_GH.rep



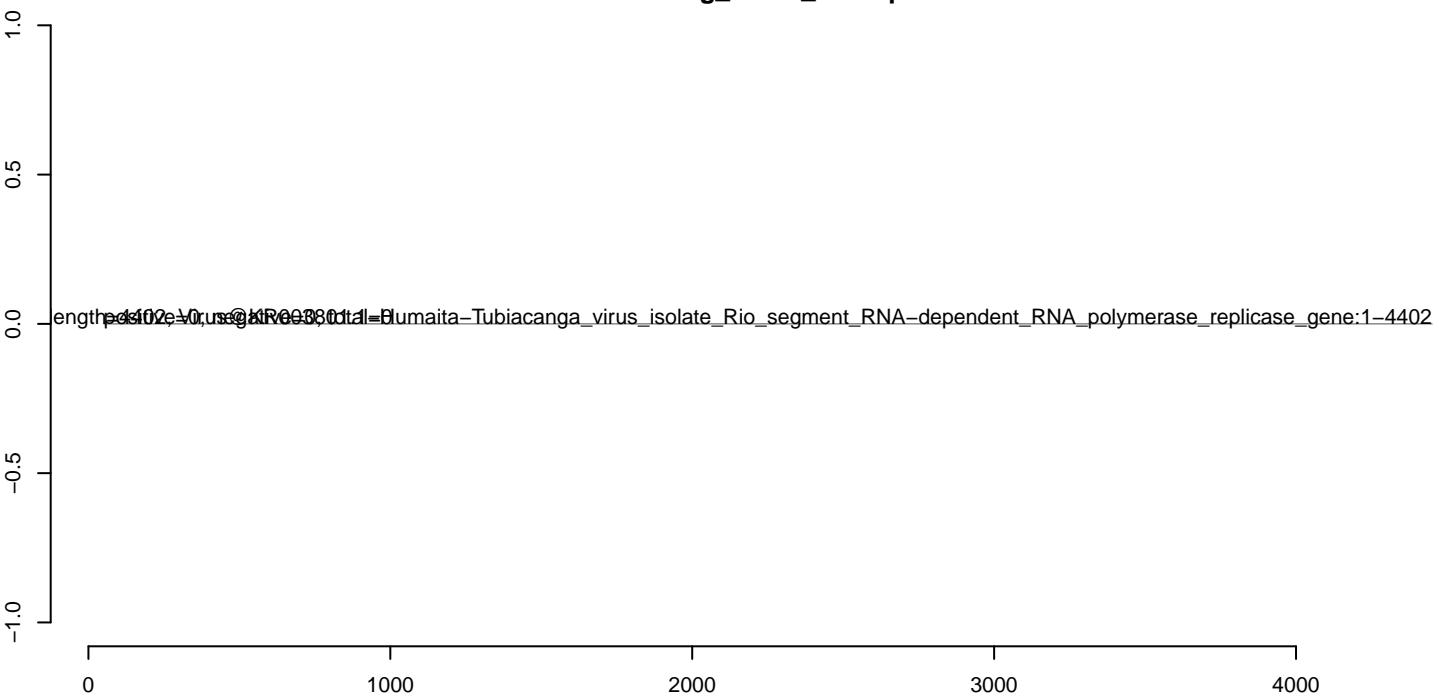
### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep

positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.24\_35.rep

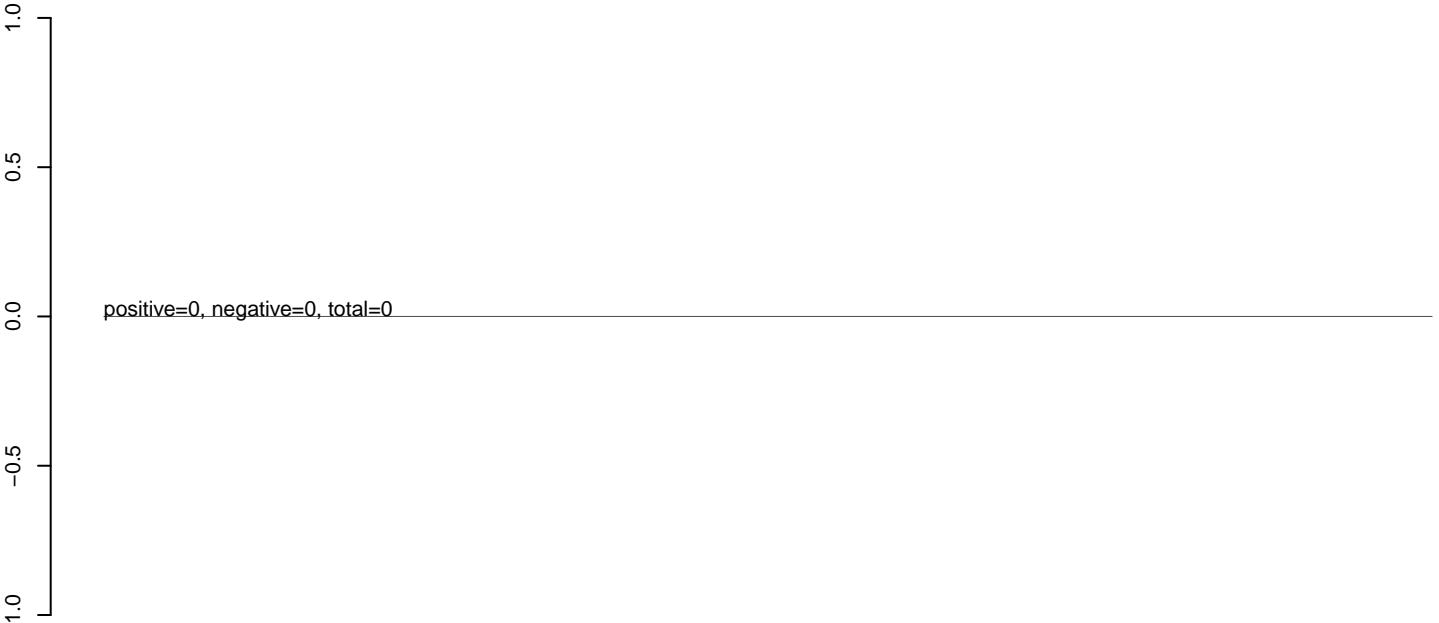
positive=0, negative=0, total=0

### AeAeg\_Fcanc\_GH.rep

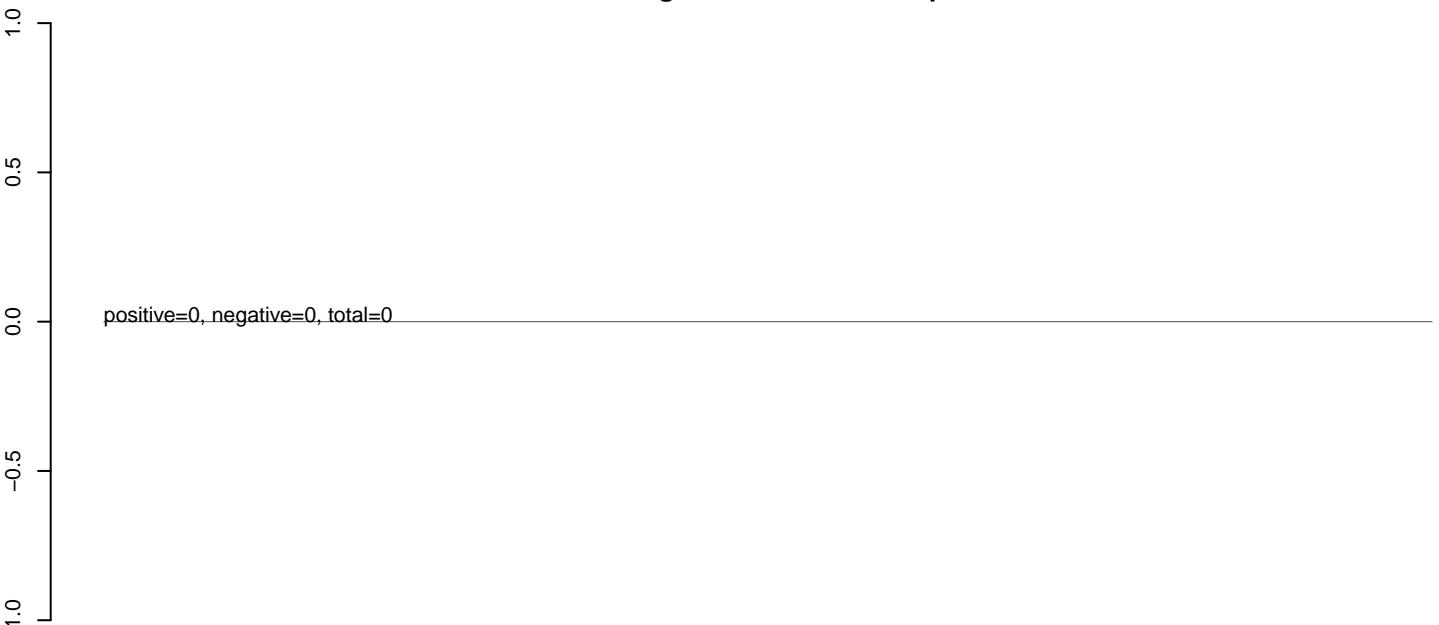
positive=0, negative=0, total=0 Window size=50, length=3776, Virus@FJ360744.1-Aedes\_aegypti\_densovirus\_2\_strain\_0814616:1-3776

0 1000 2000 3000

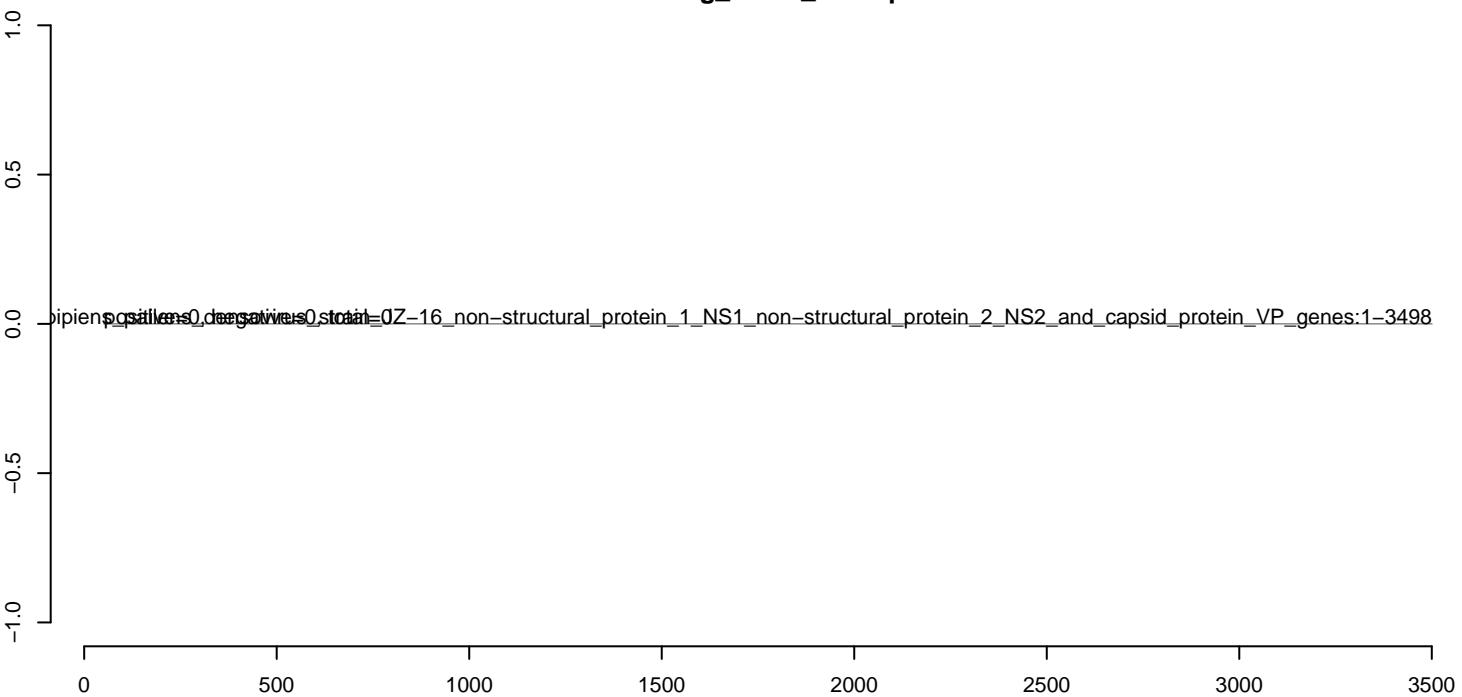
### AeAeg\_Fcanc\_GH.18\_23.rep



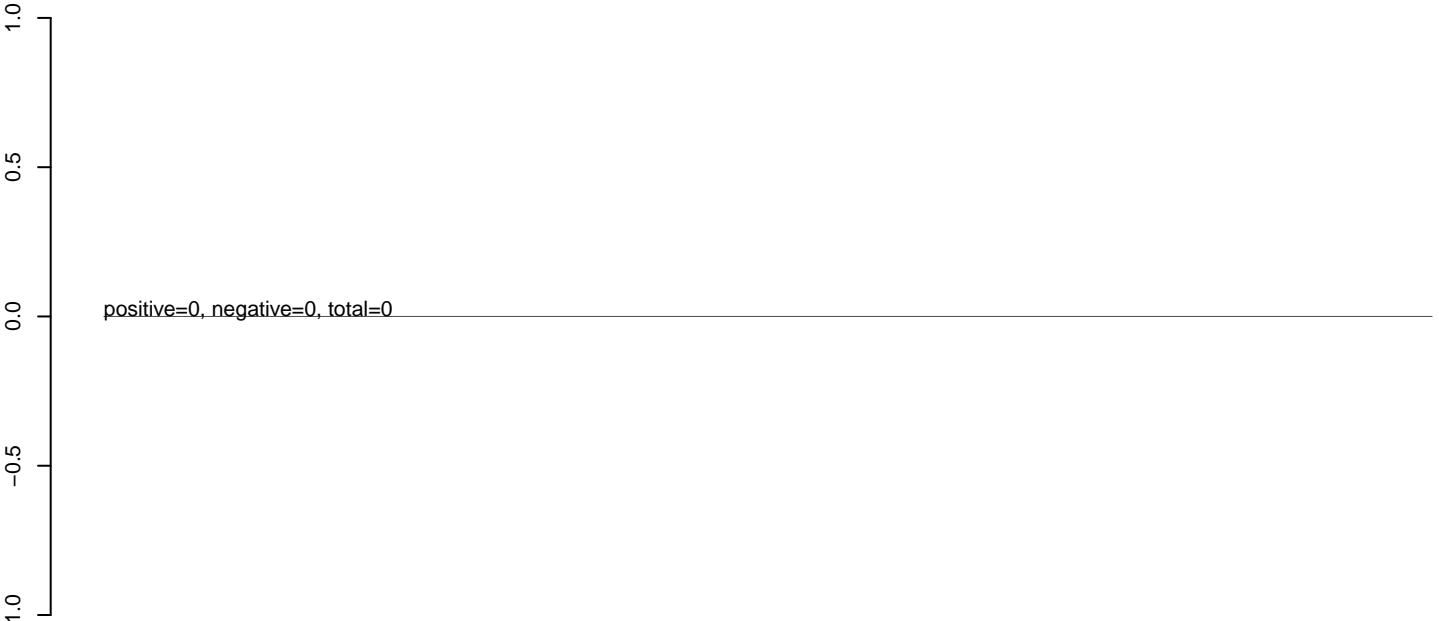
### AeAeg\_Fcanc\_GH.24\_35.rep



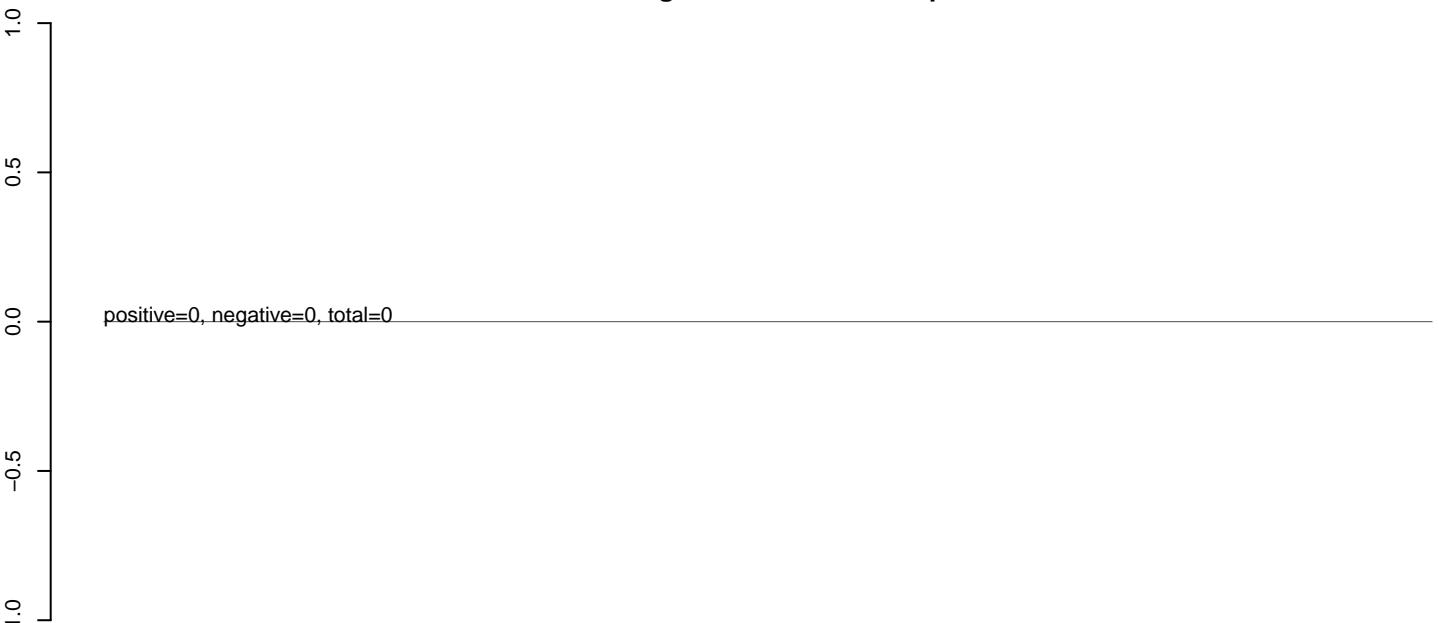
### AeAeg\_Fcanc\_GH.rep



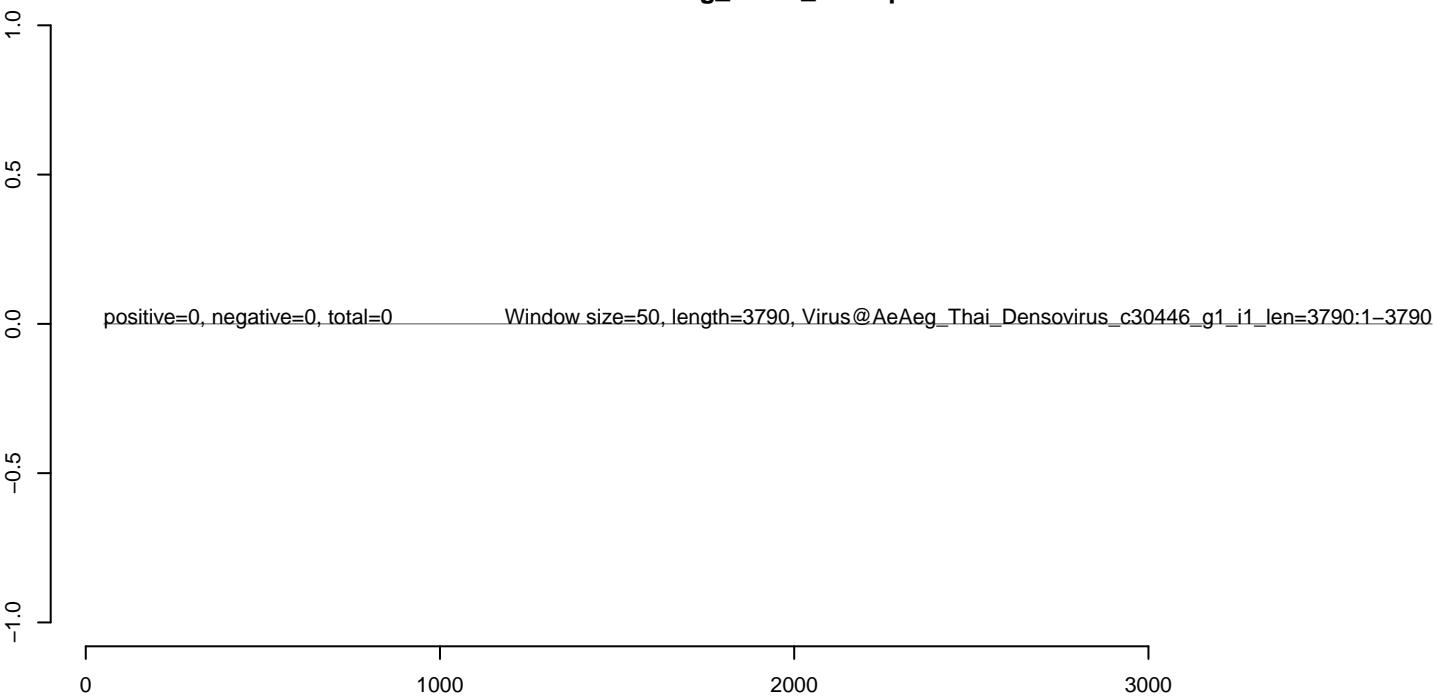
### AeAeg\_Fcanc\_GH.18\_23.rep



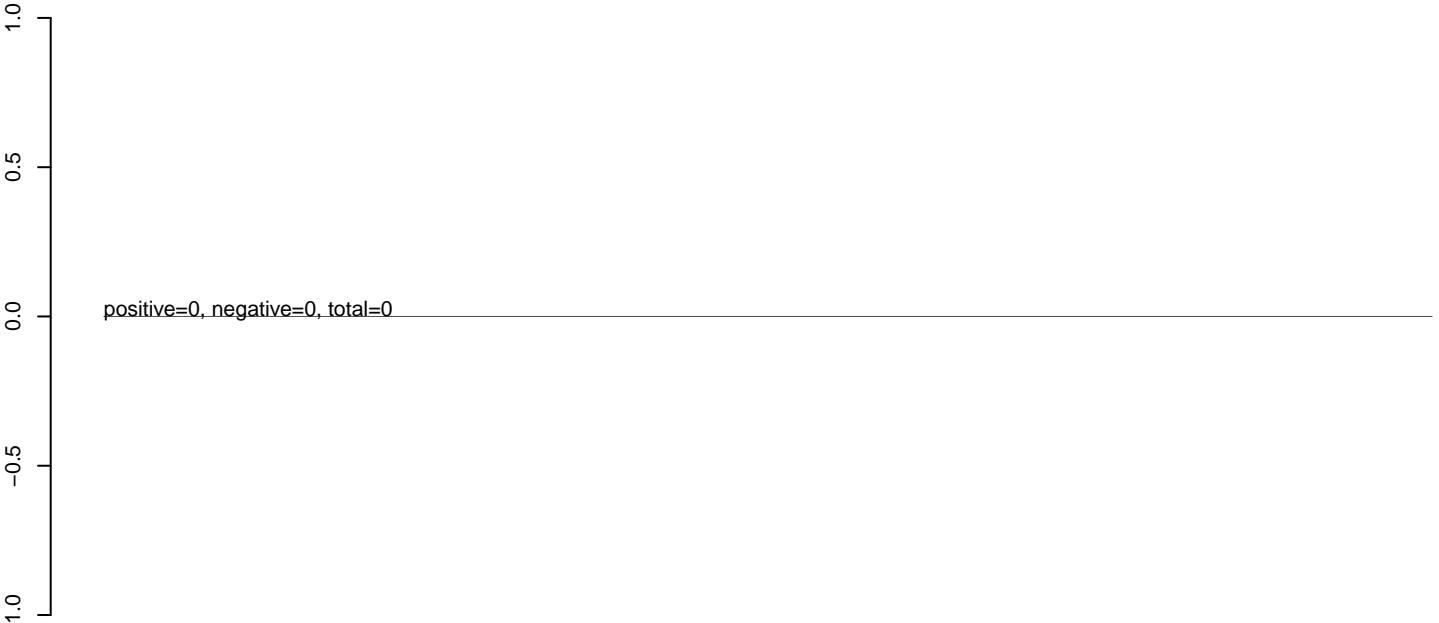
### AeAeg\_Fcanc\_GH.24\_35.rep



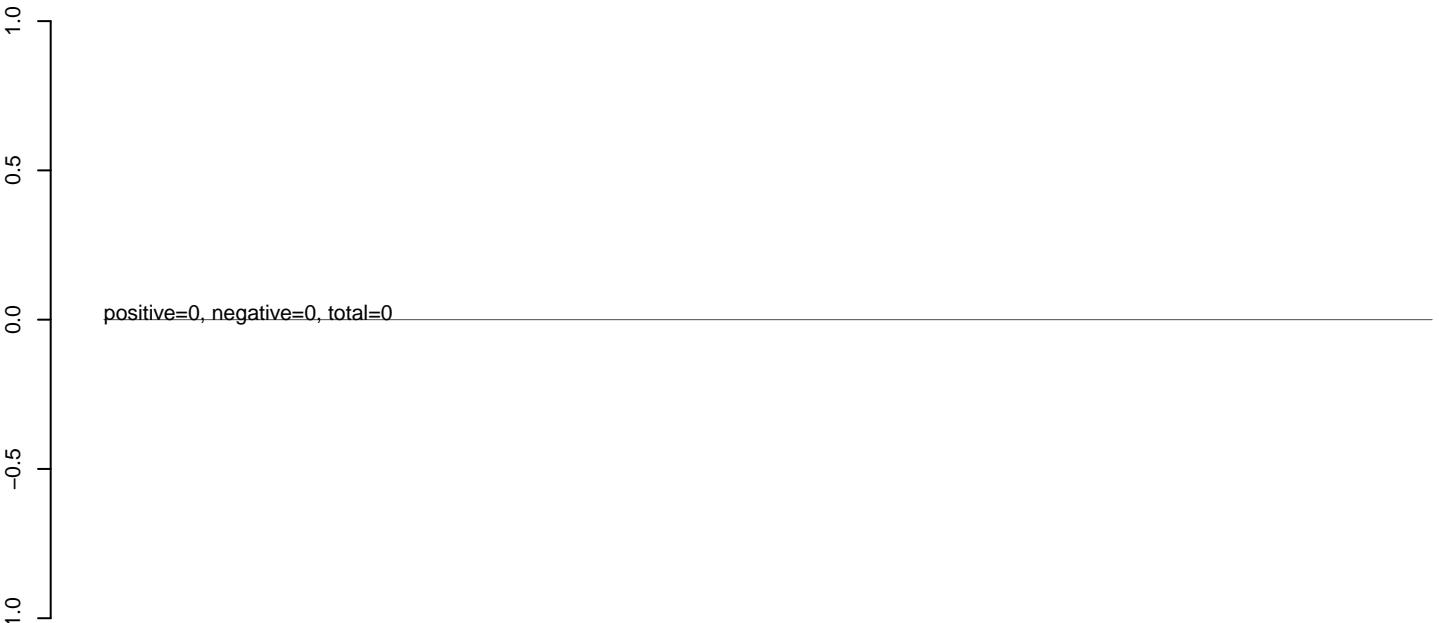
### AeAeg\_Fcanc\_GH.rep



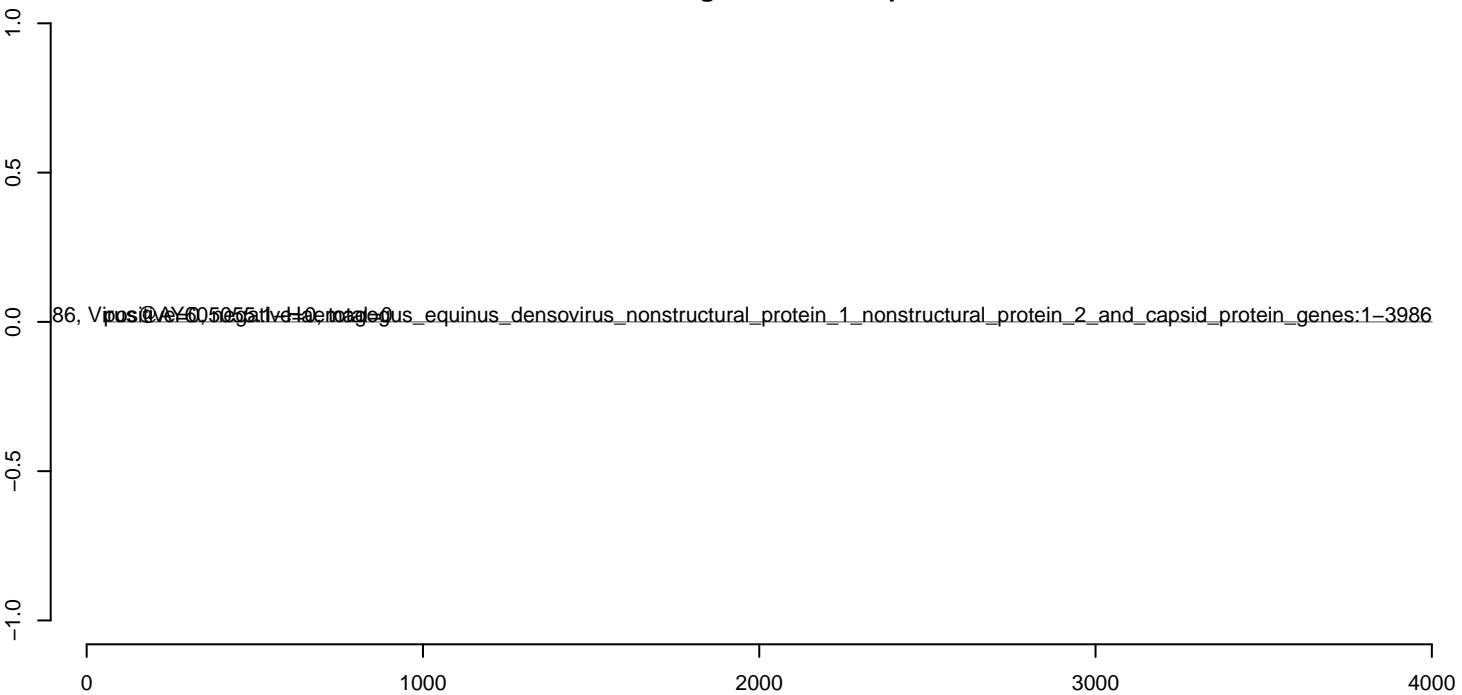
### AeAeg\_Fcanc\_GH.18\_23.rep



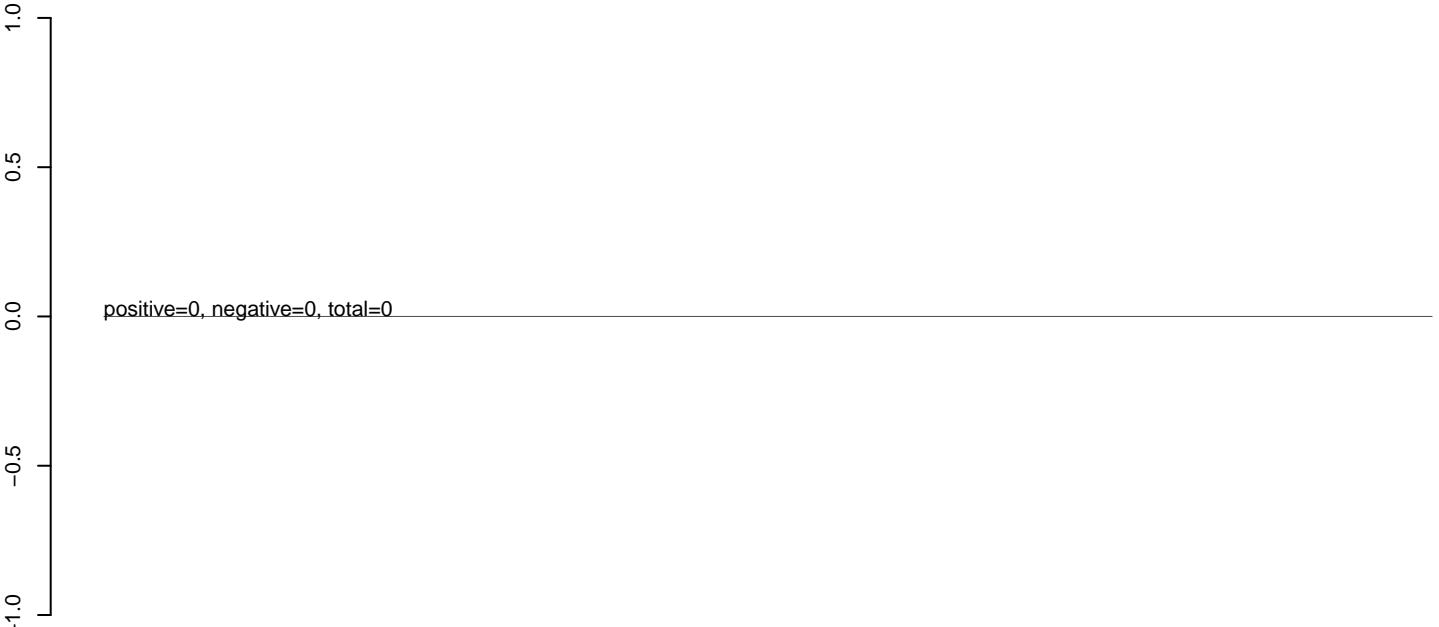
### AeAeg\_Fcanc\_GH.24\_35.rep



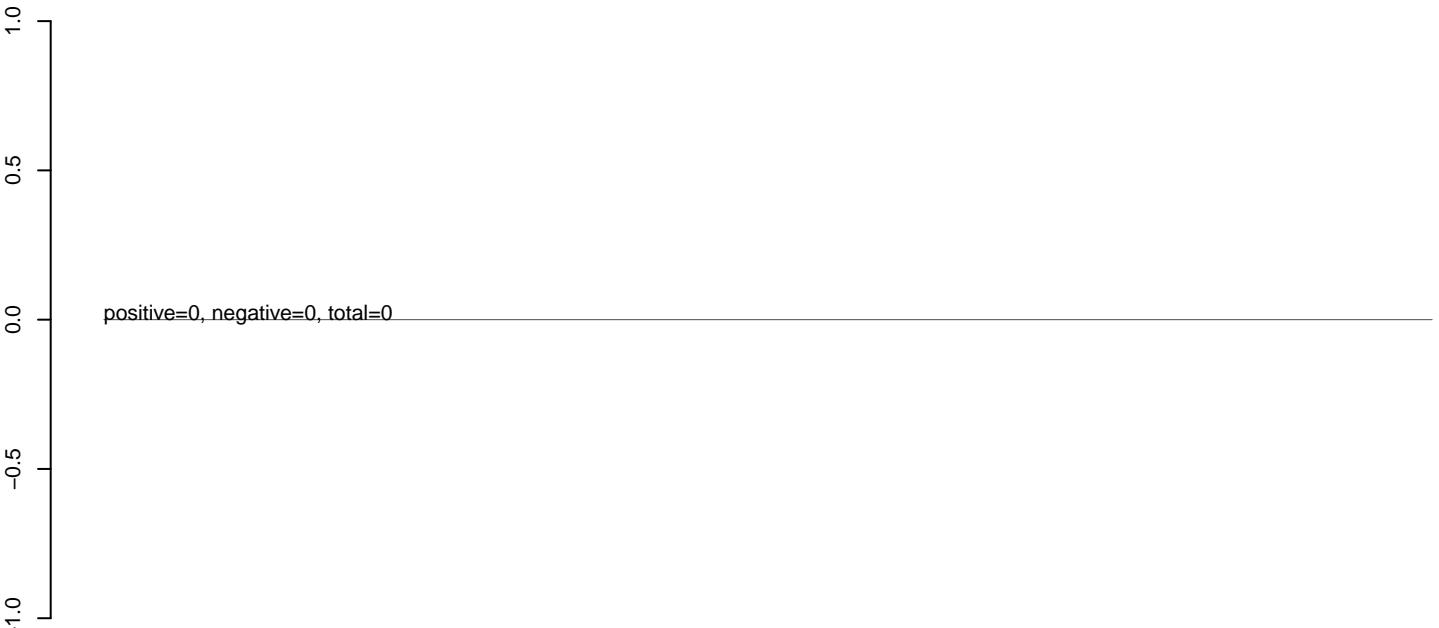
### AeAeg\_Fcanc\_GH.rep



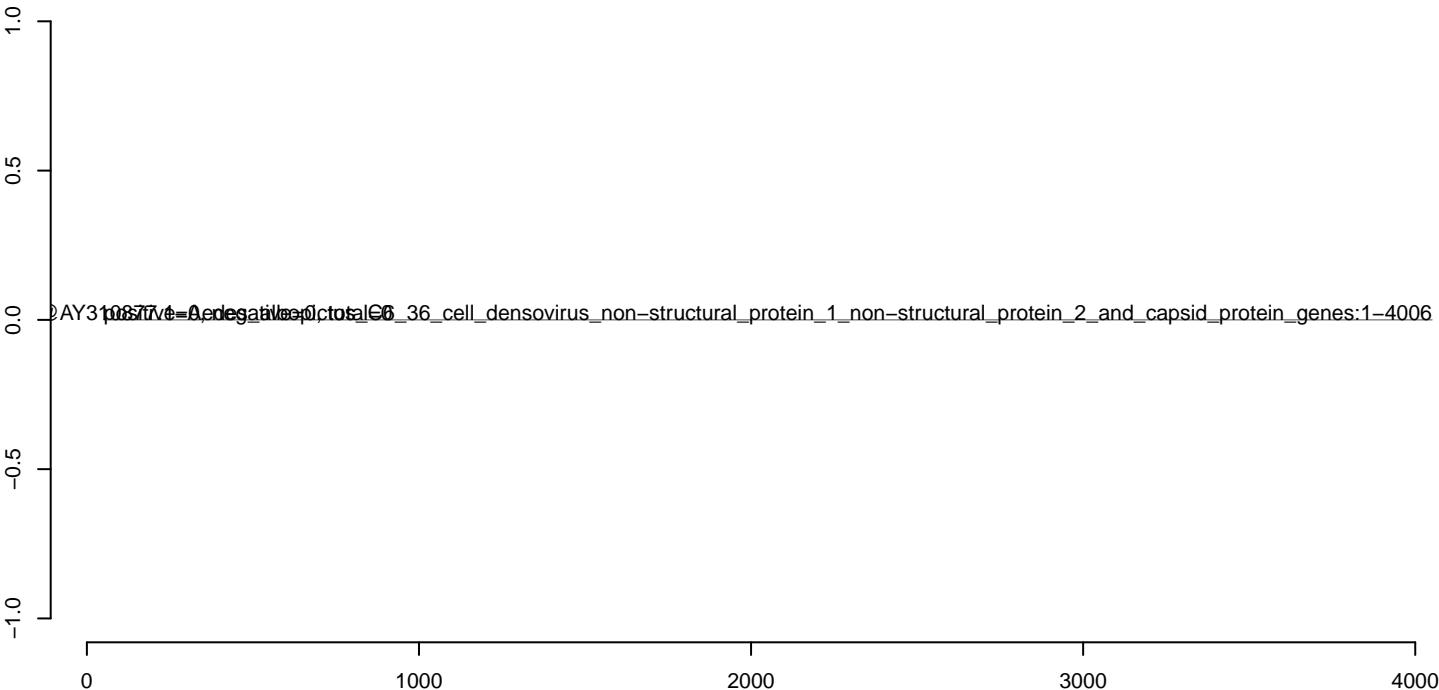
### AeAeg\_Fcanc\_GH.18\_23.rep



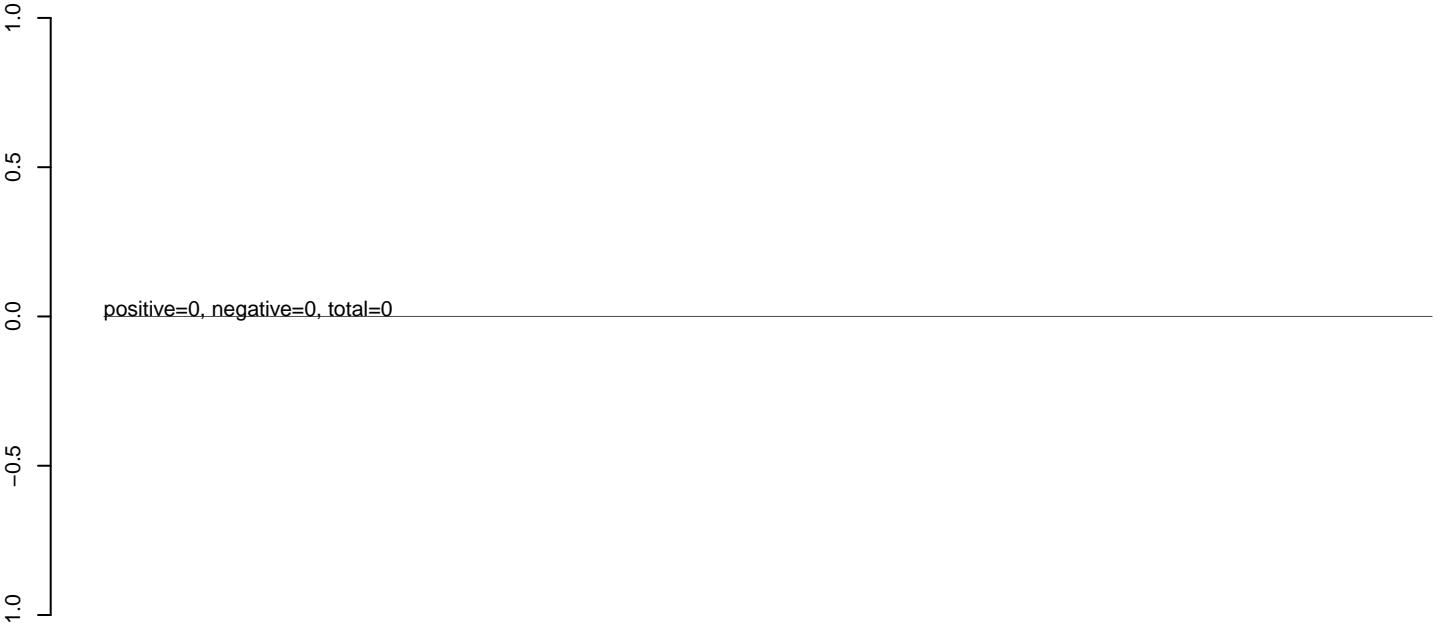
### AeAeg\_Fcanc\_GH.24\_35.rep



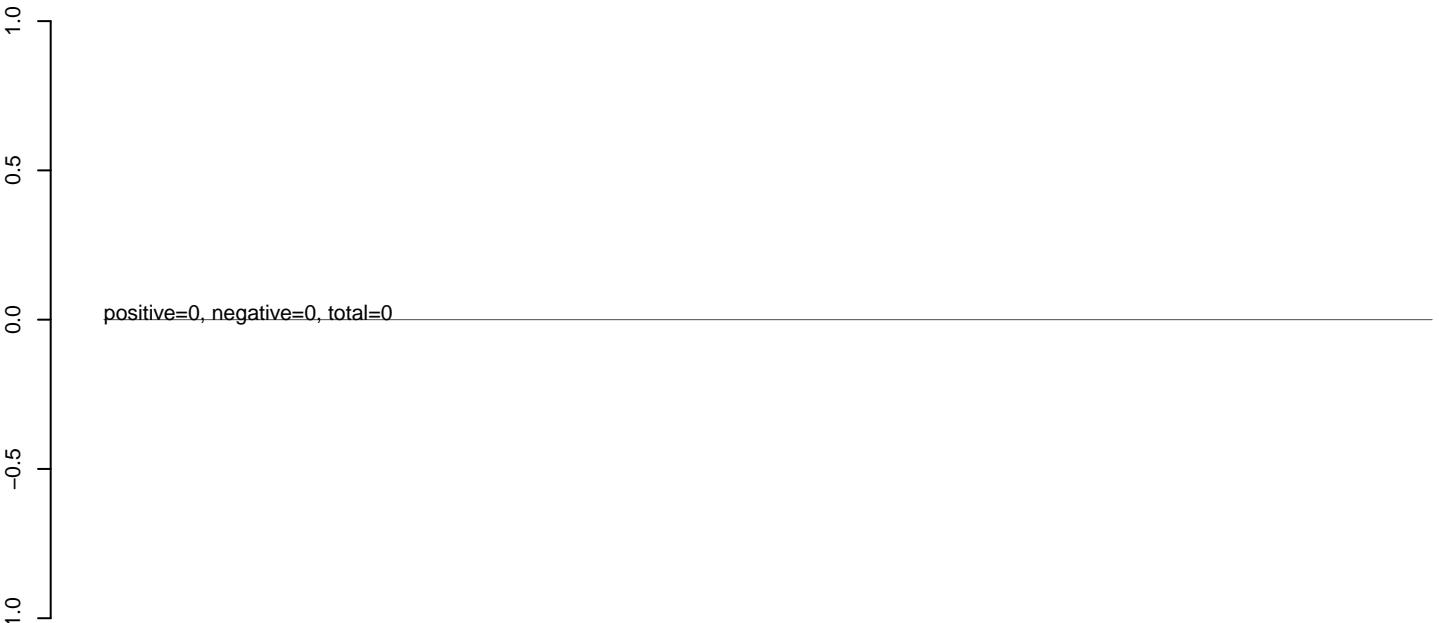
### AeAeg\_Fcanc\_GH.rep



### AeAeg\_Fcanc\_GH.18\_23.rep



### AeAeg\_Fcanc\_GH.24\_35.rep



### AeAeg\_Fcanc\_GH.rep

