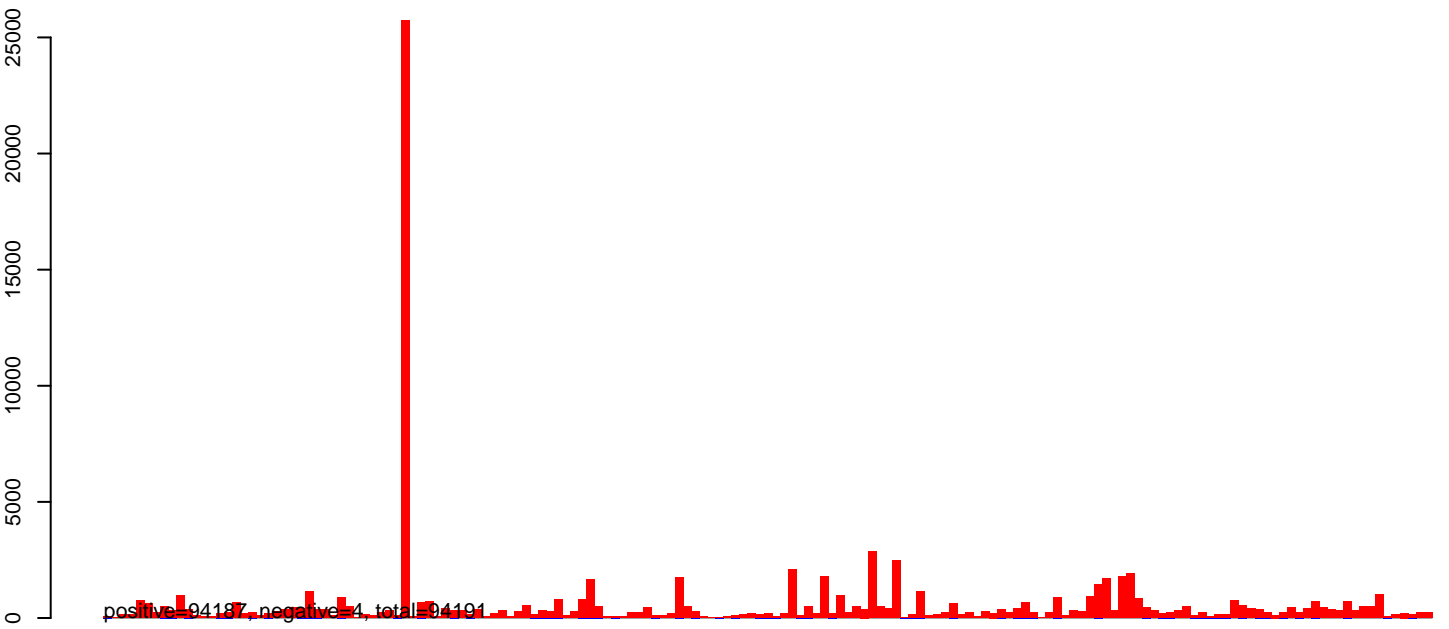
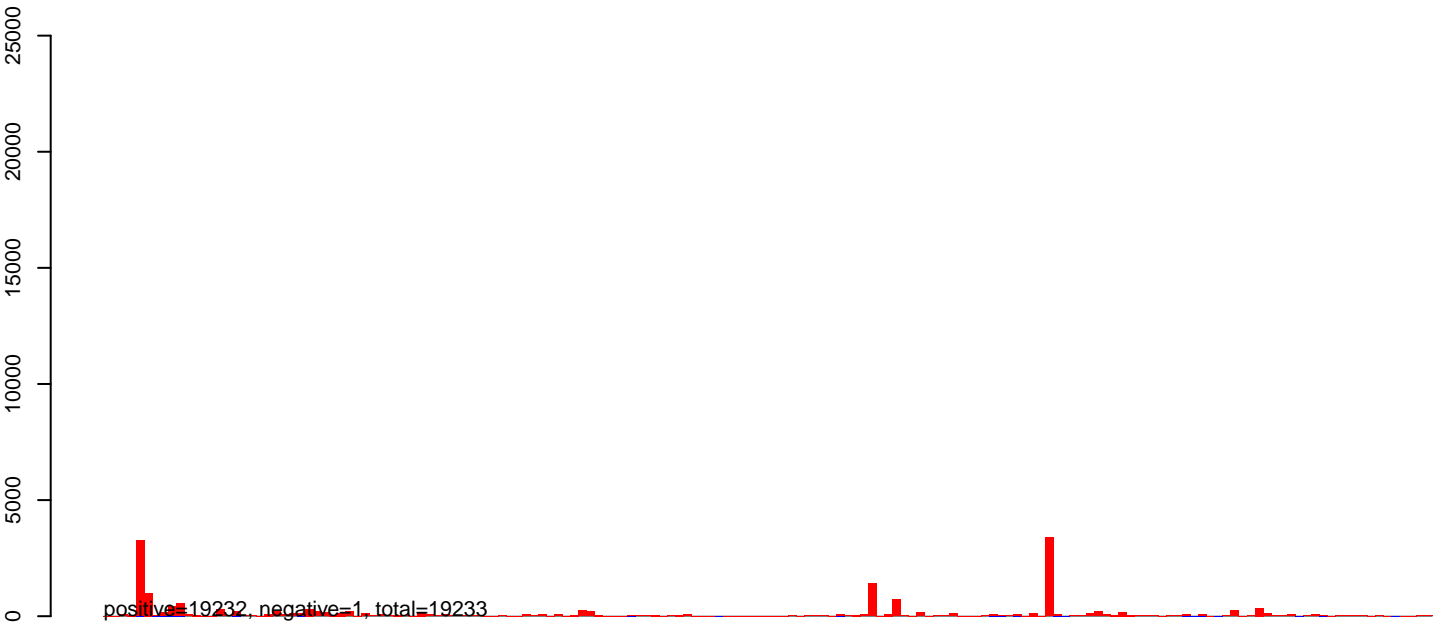


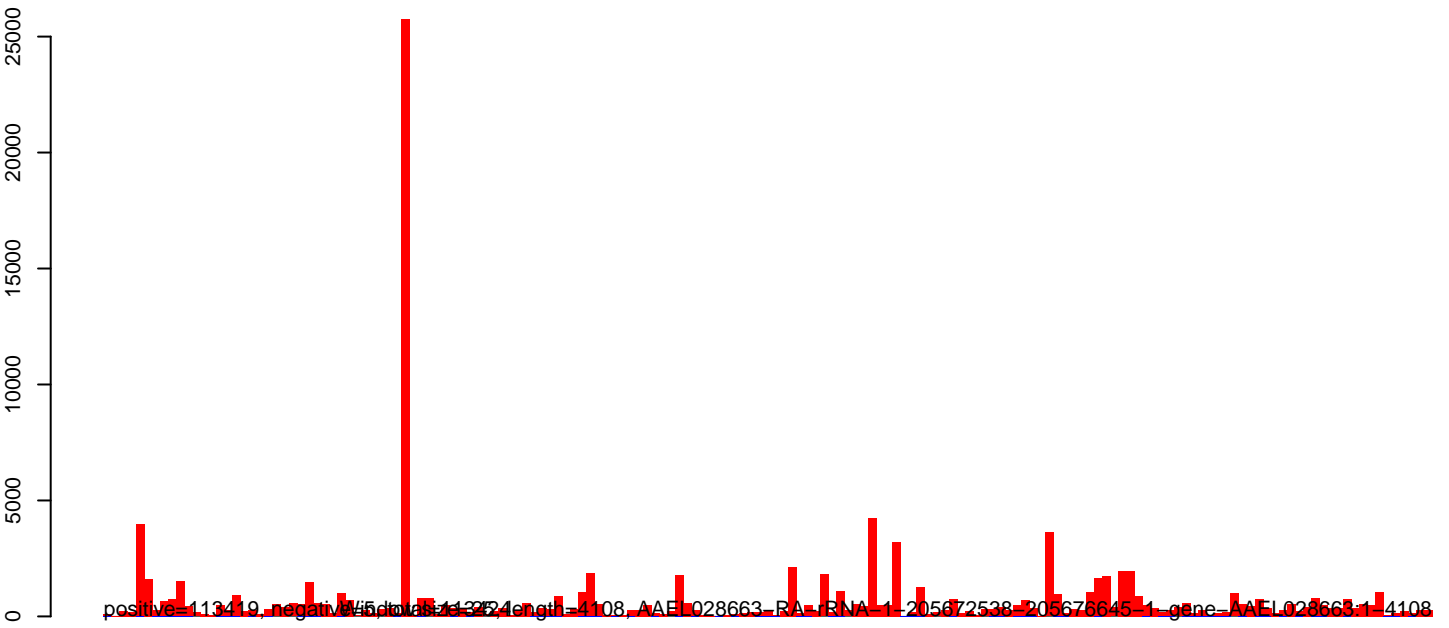
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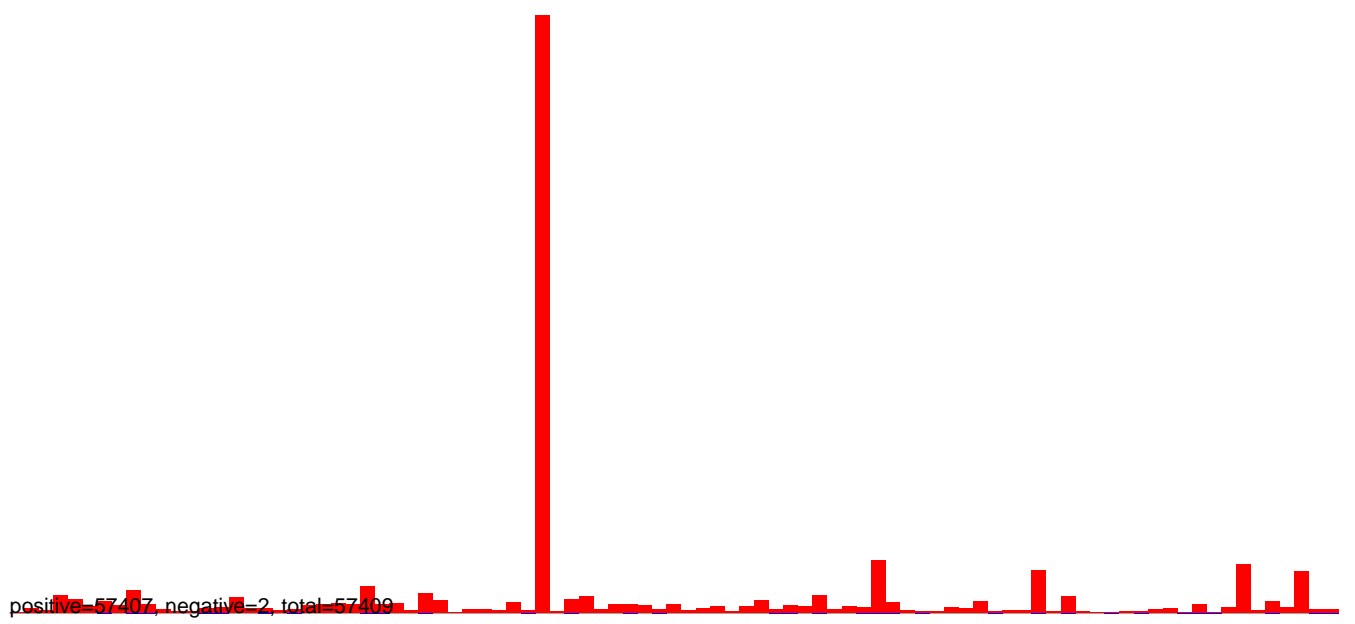
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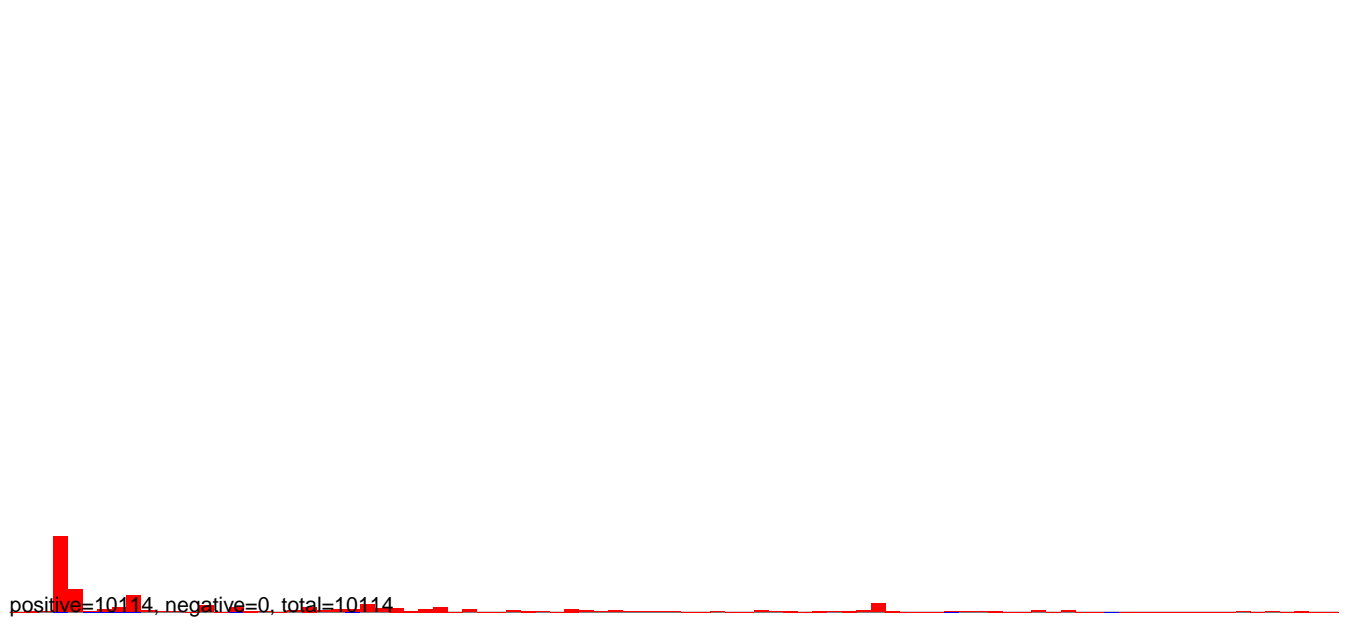
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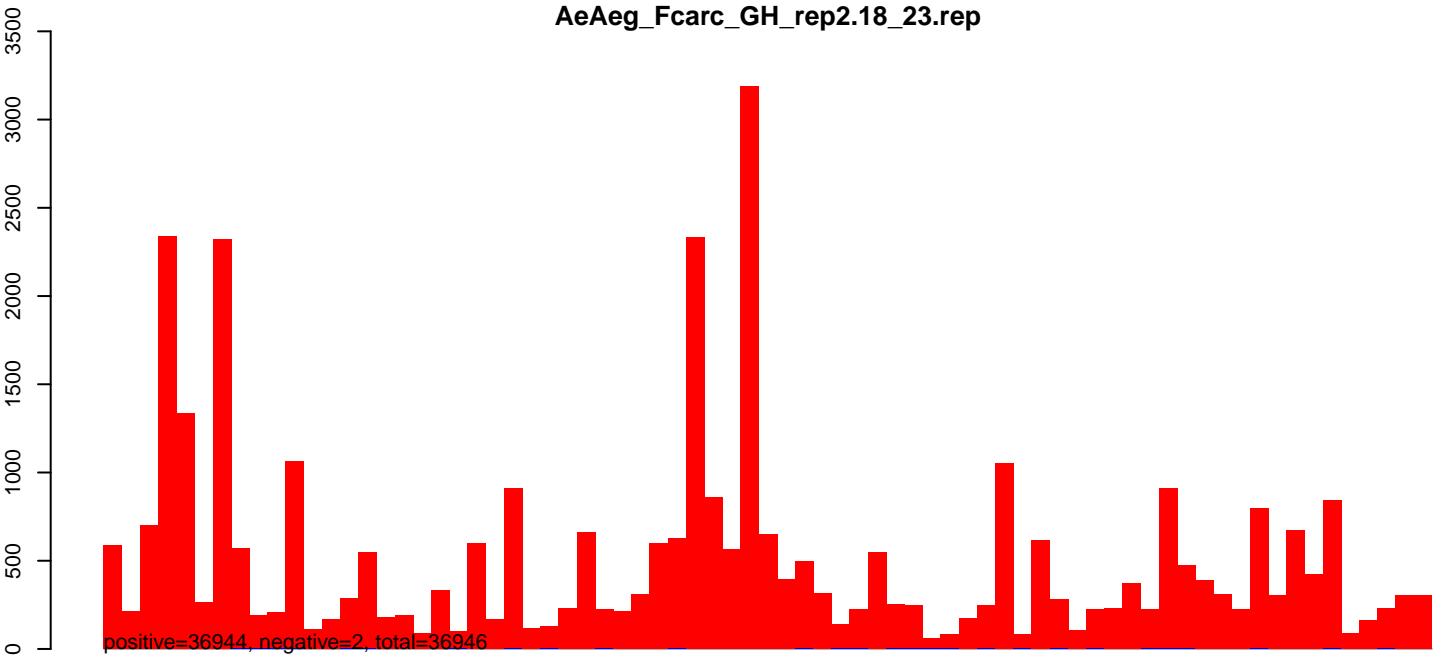
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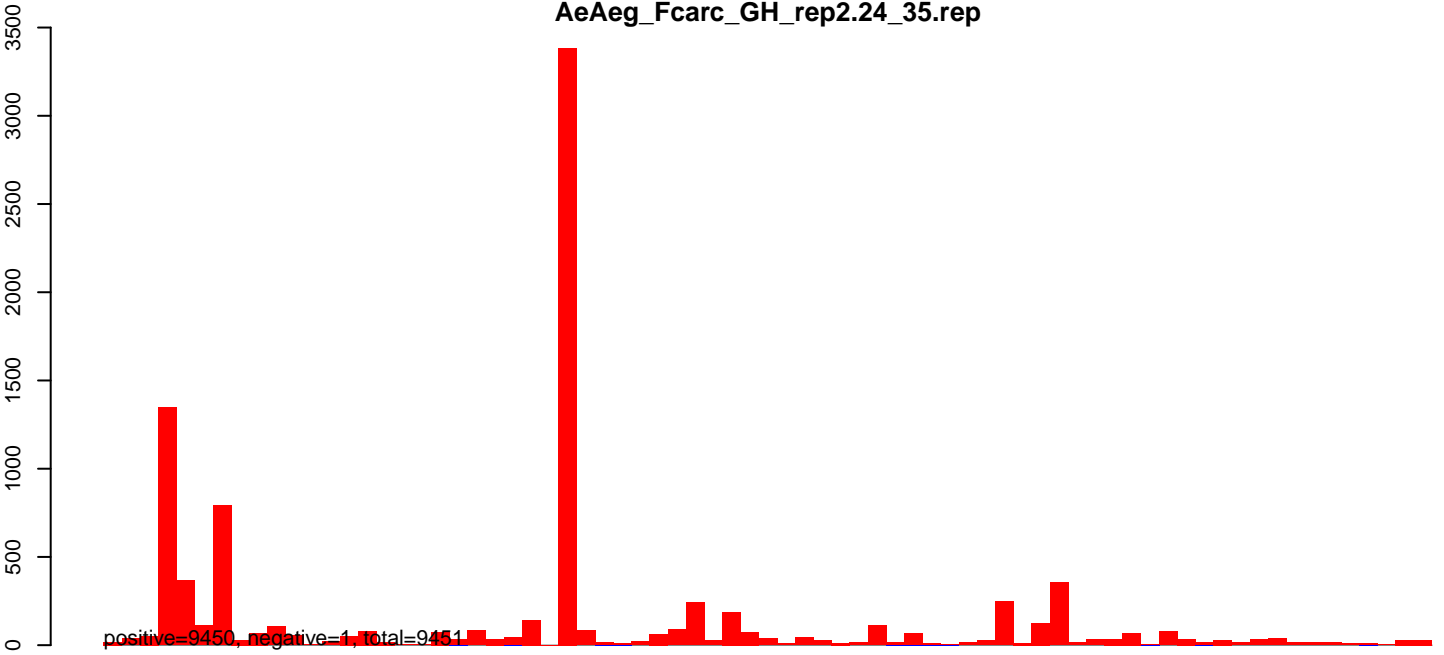
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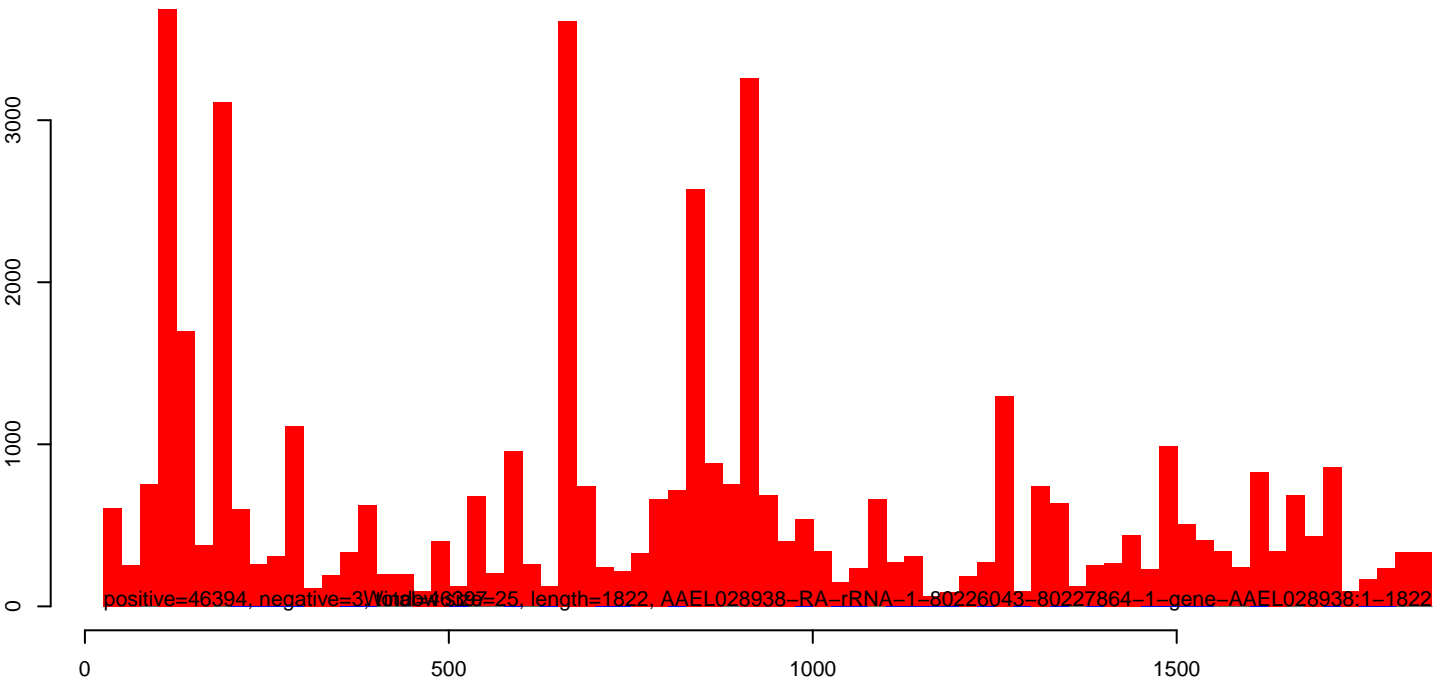
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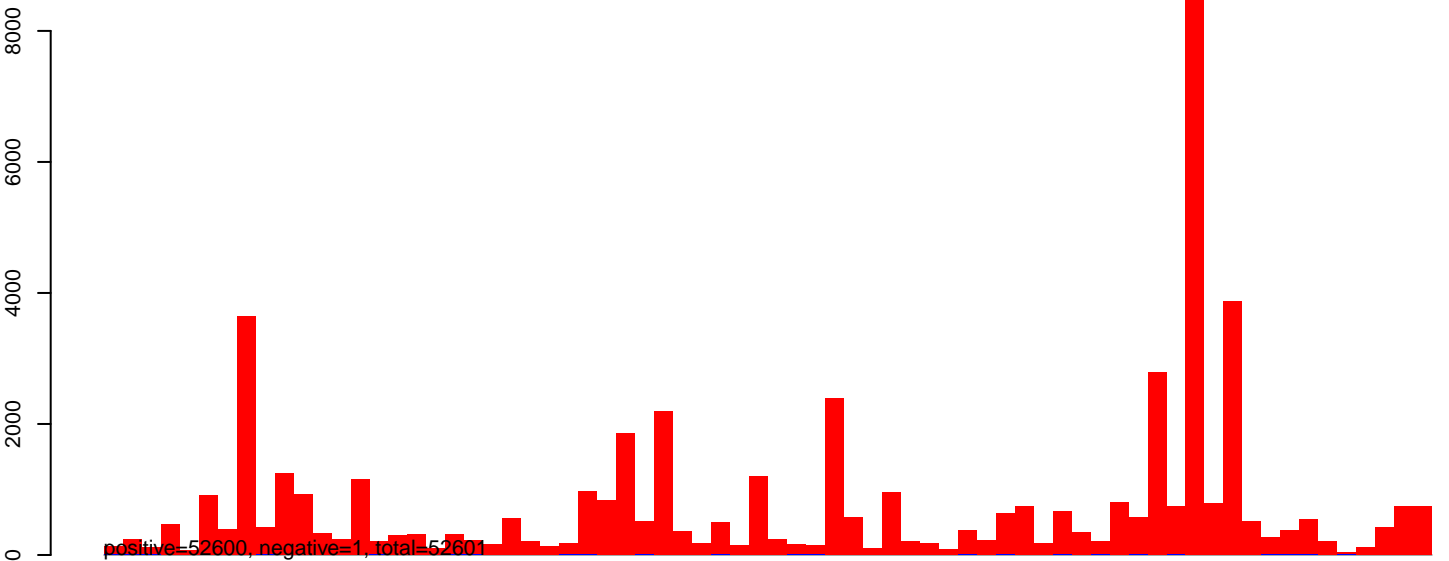
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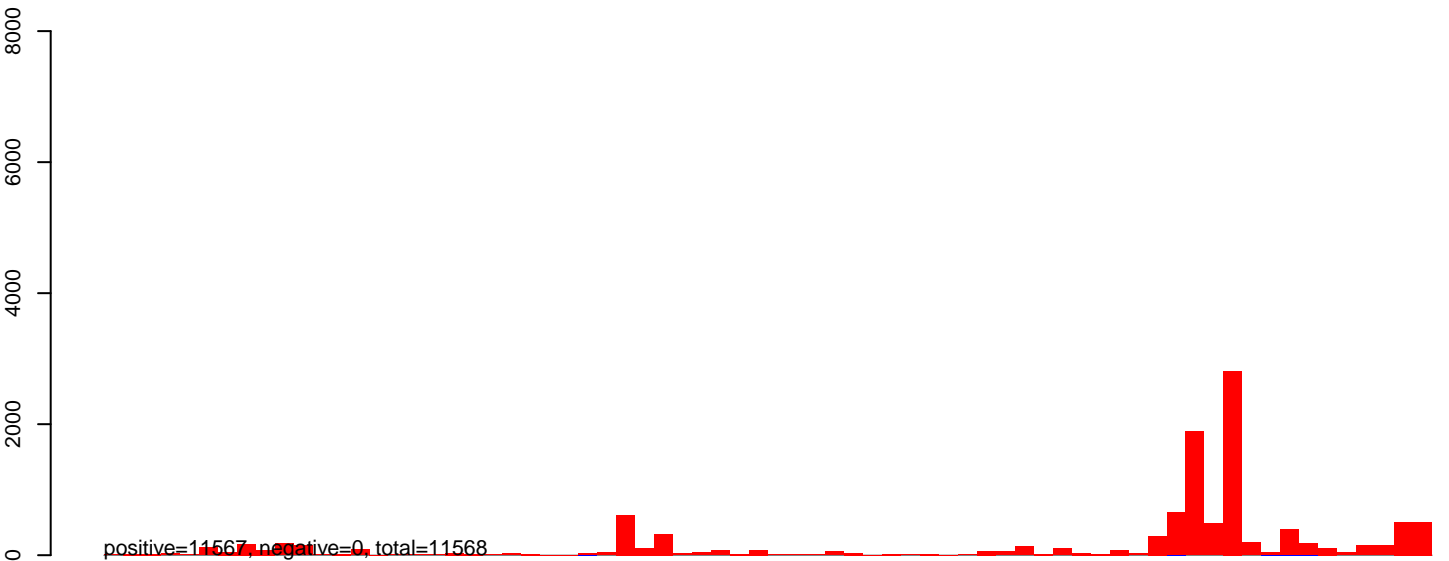
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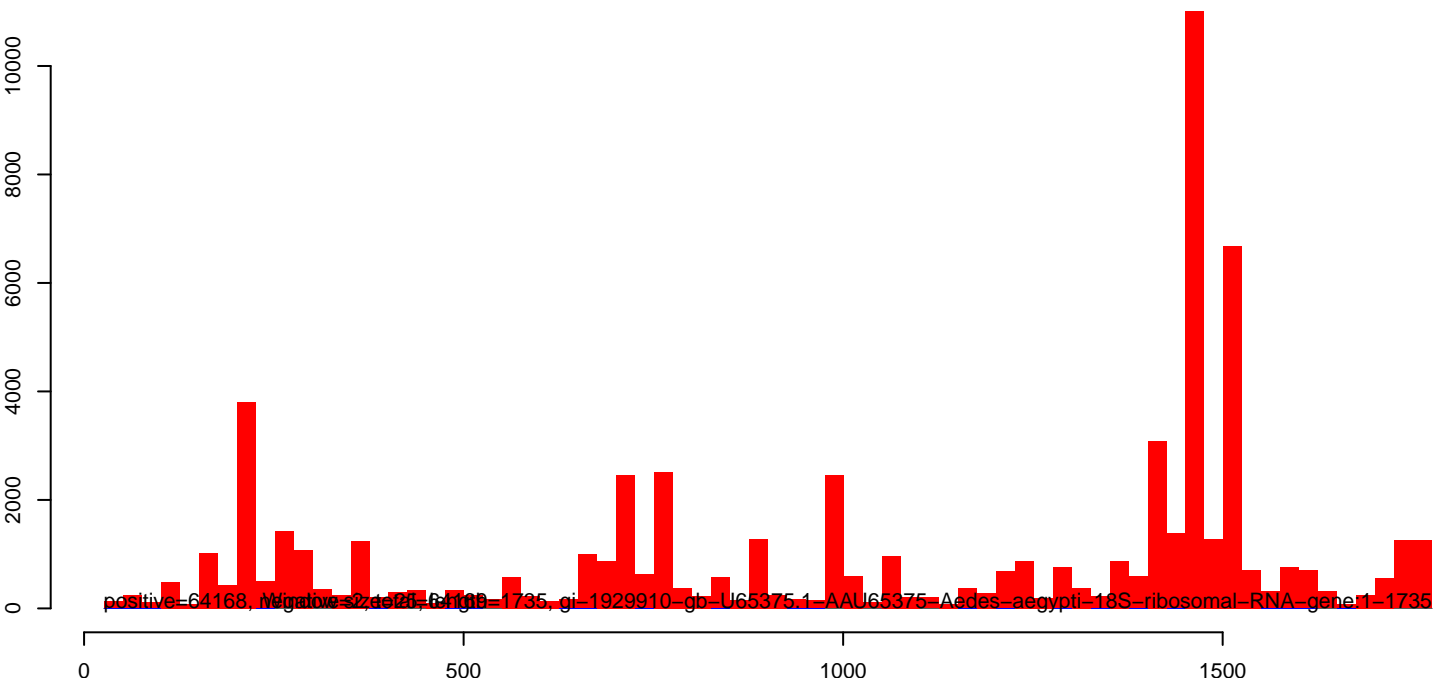
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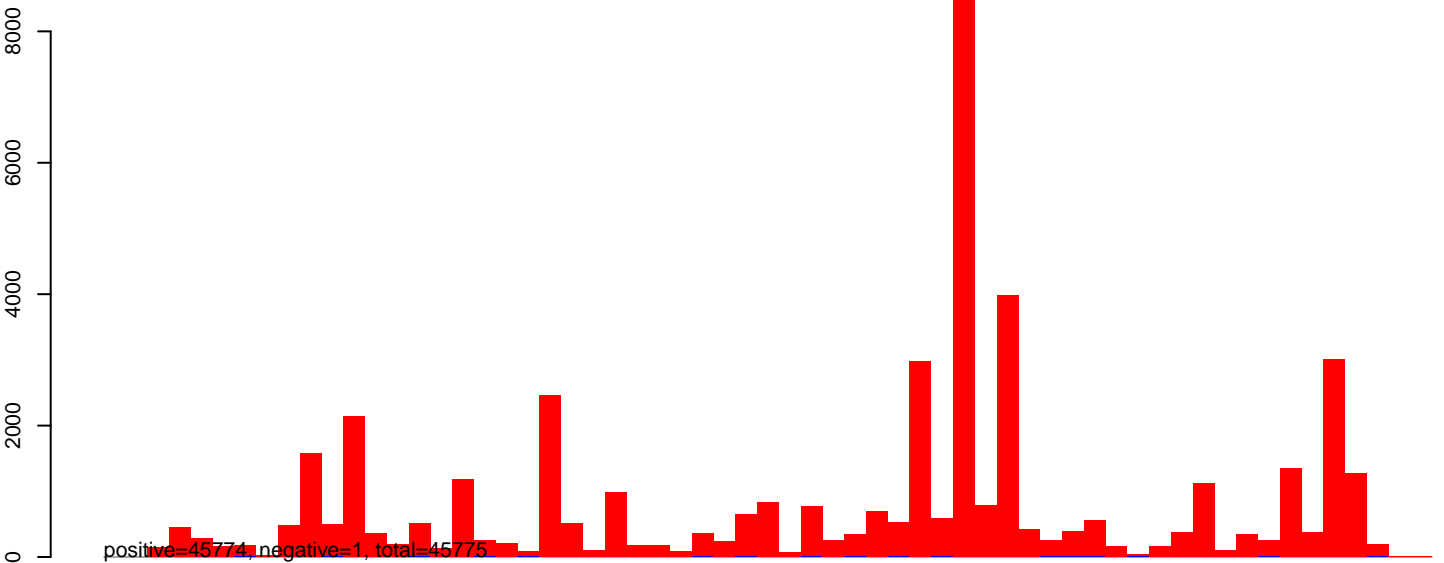
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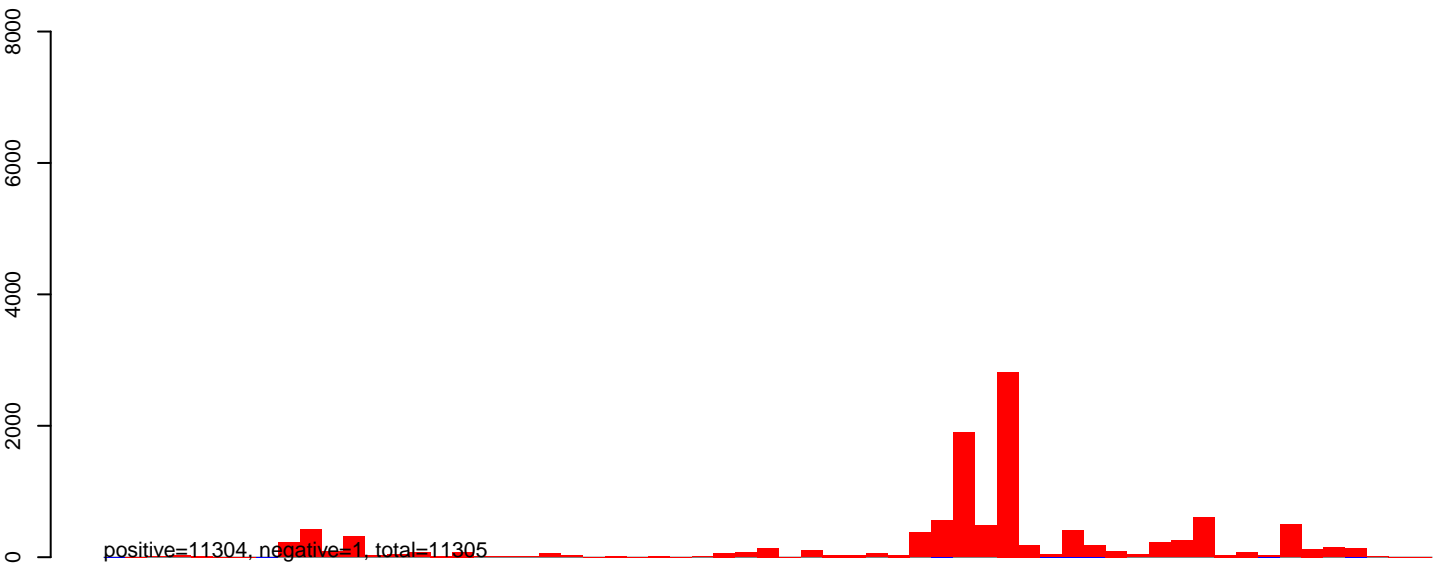
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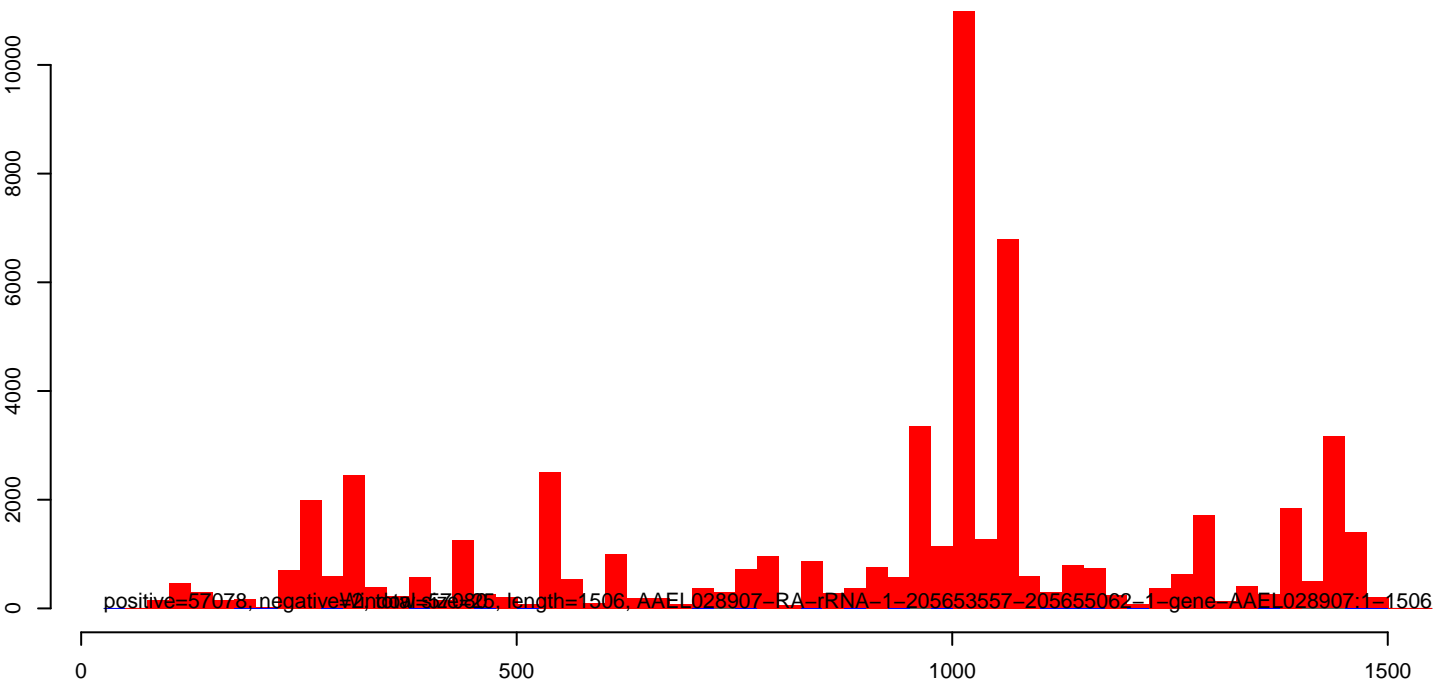
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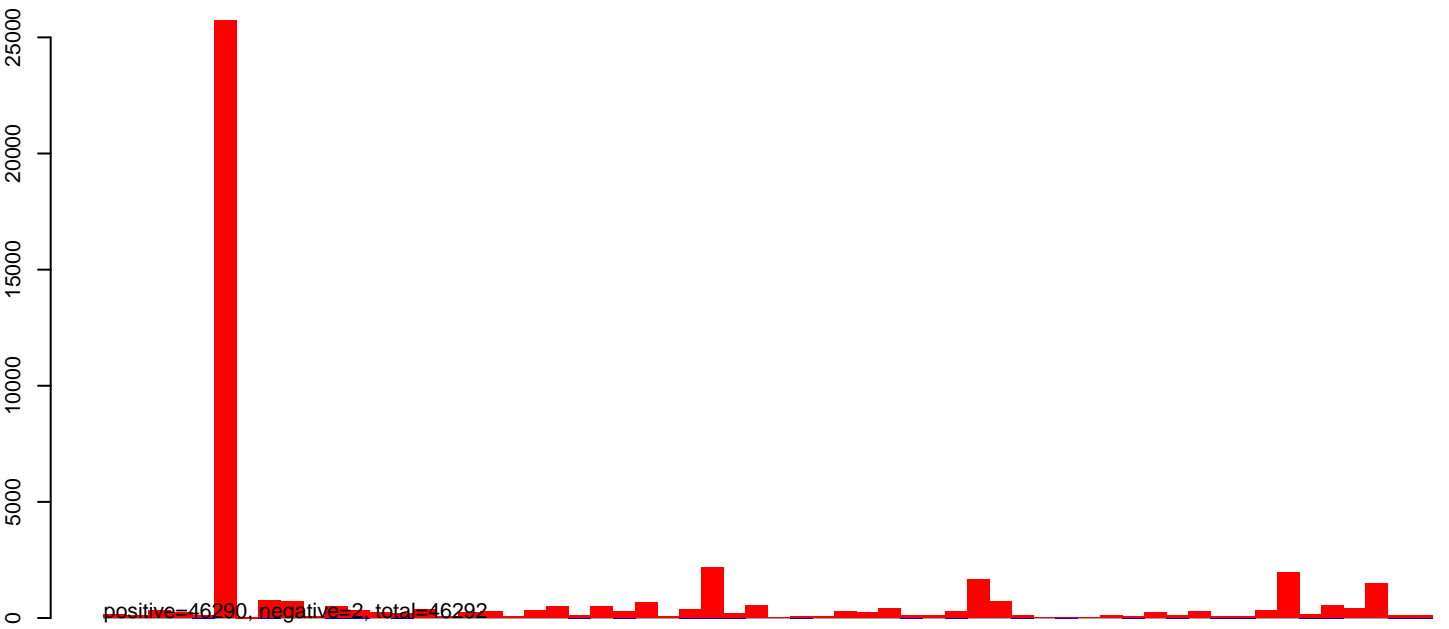
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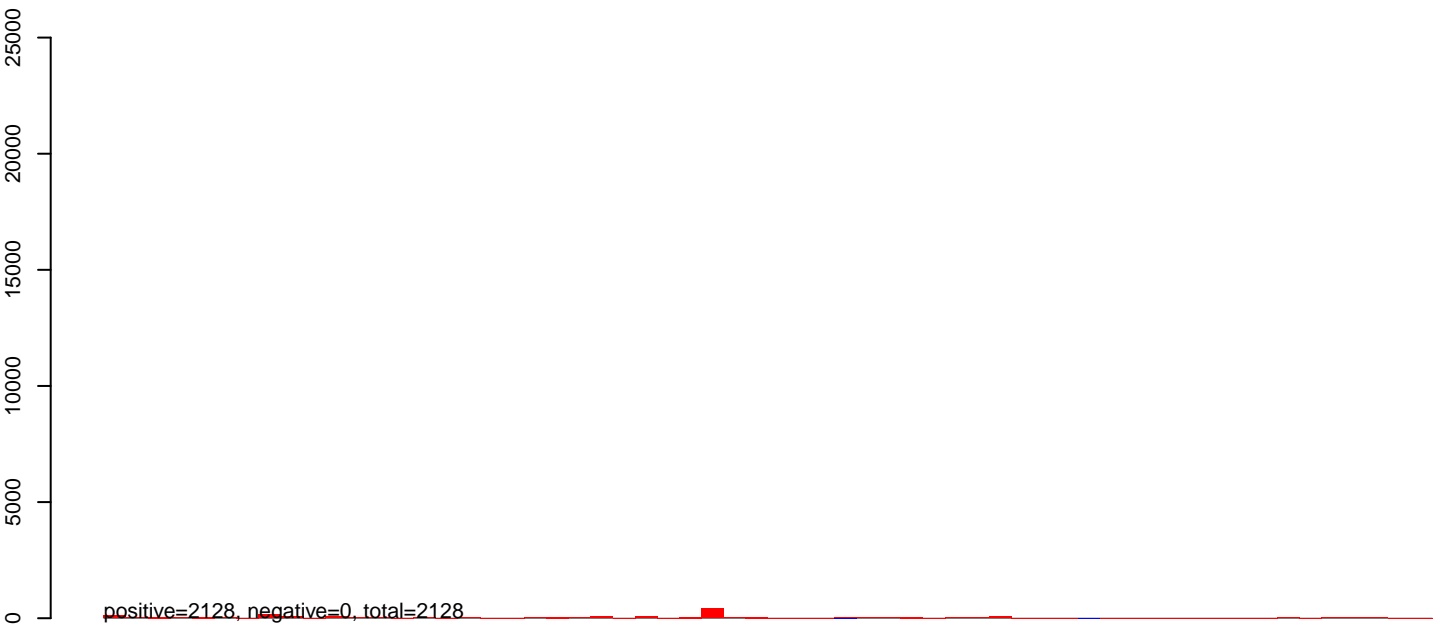
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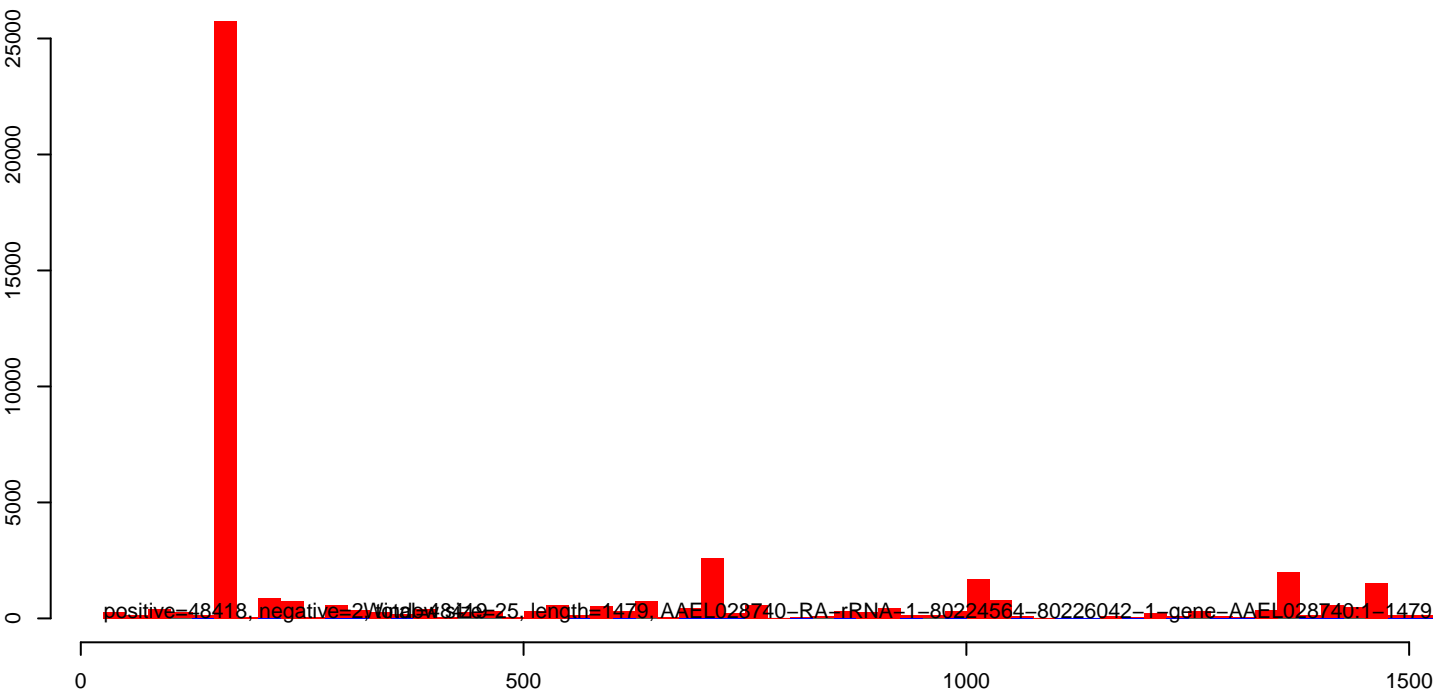
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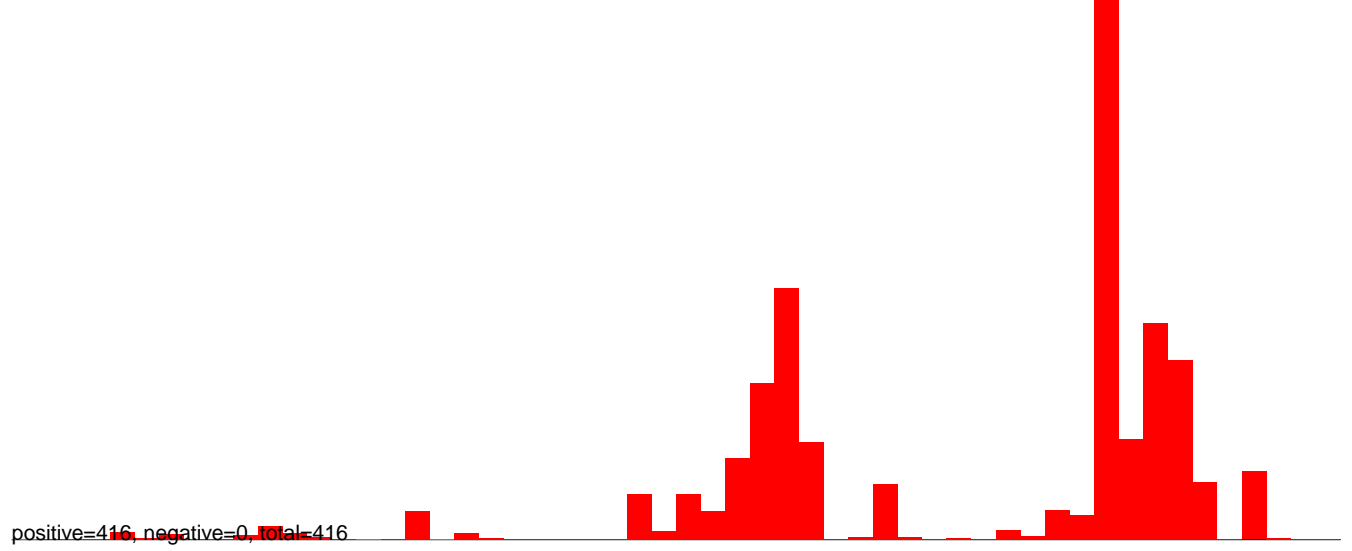
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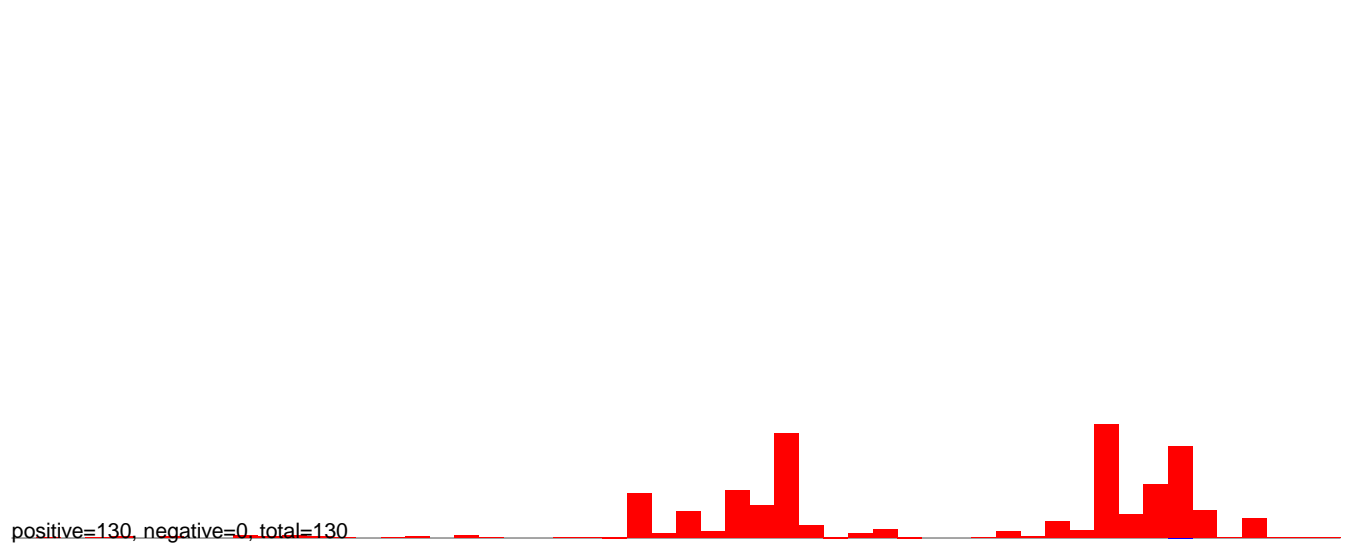
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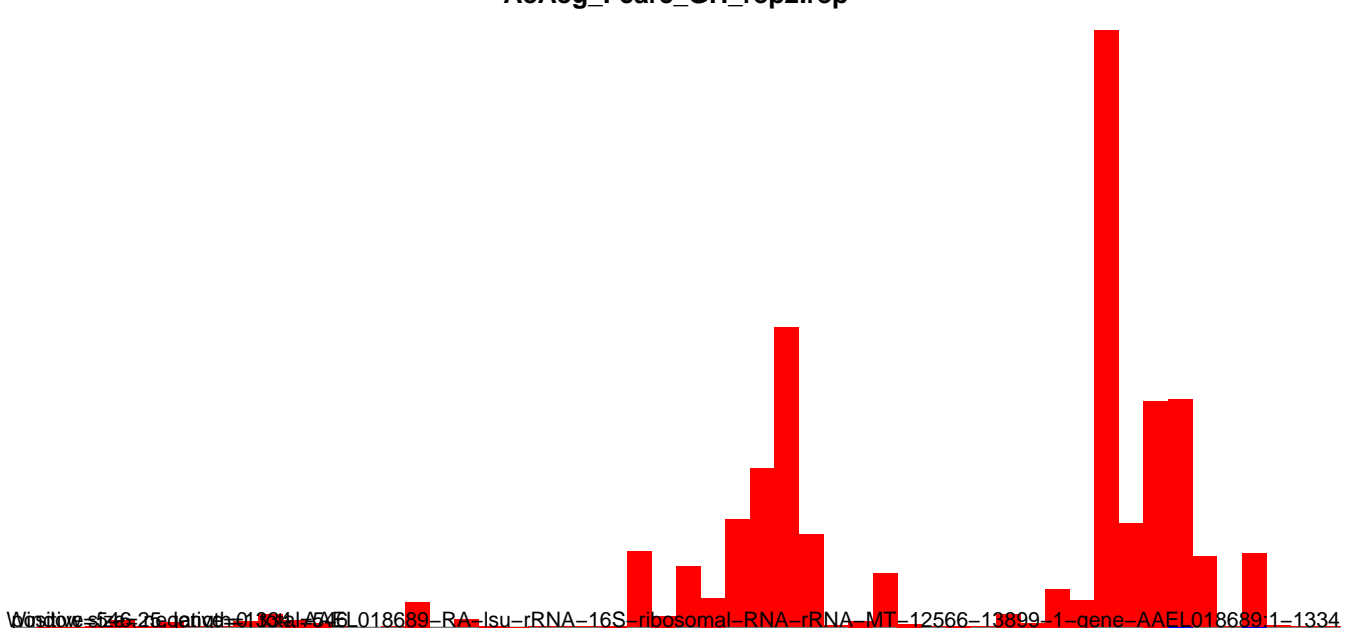
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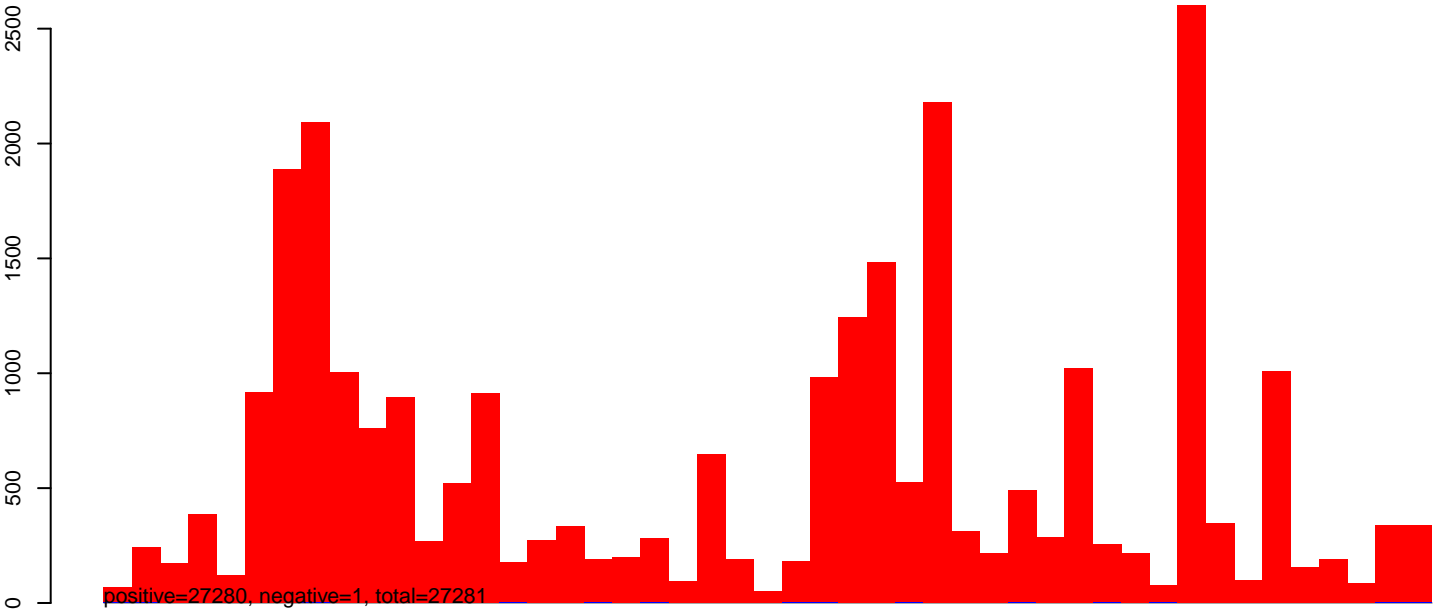


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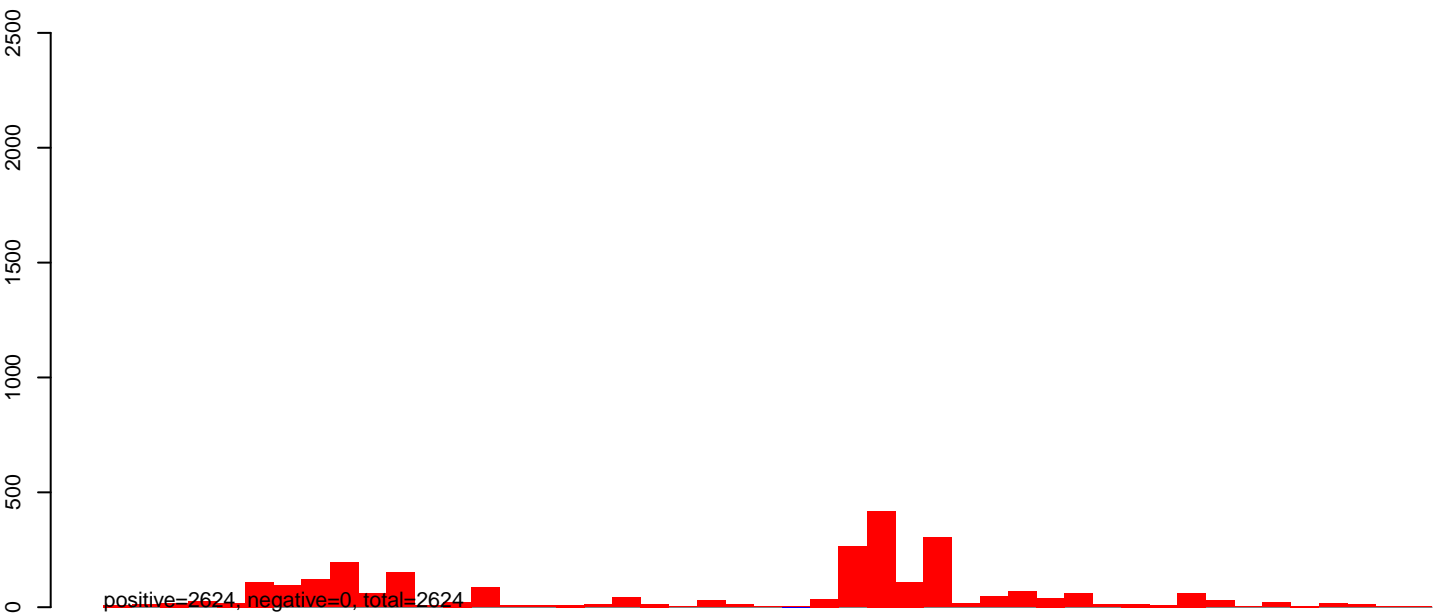


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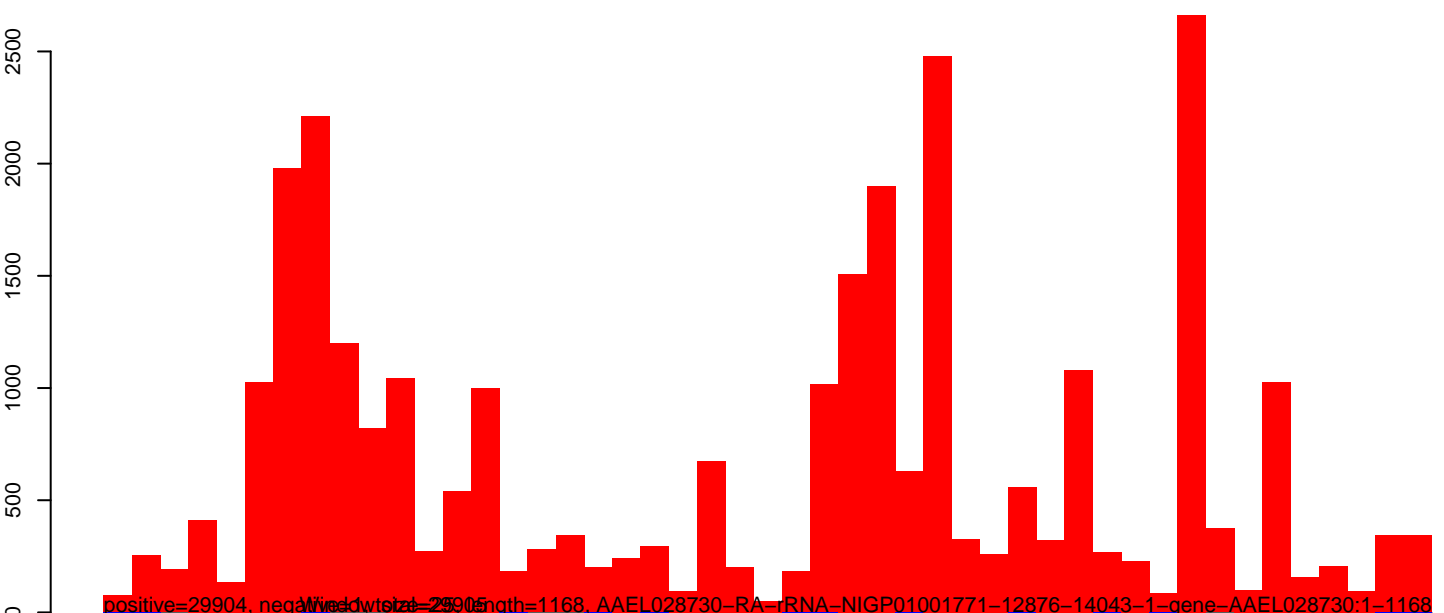
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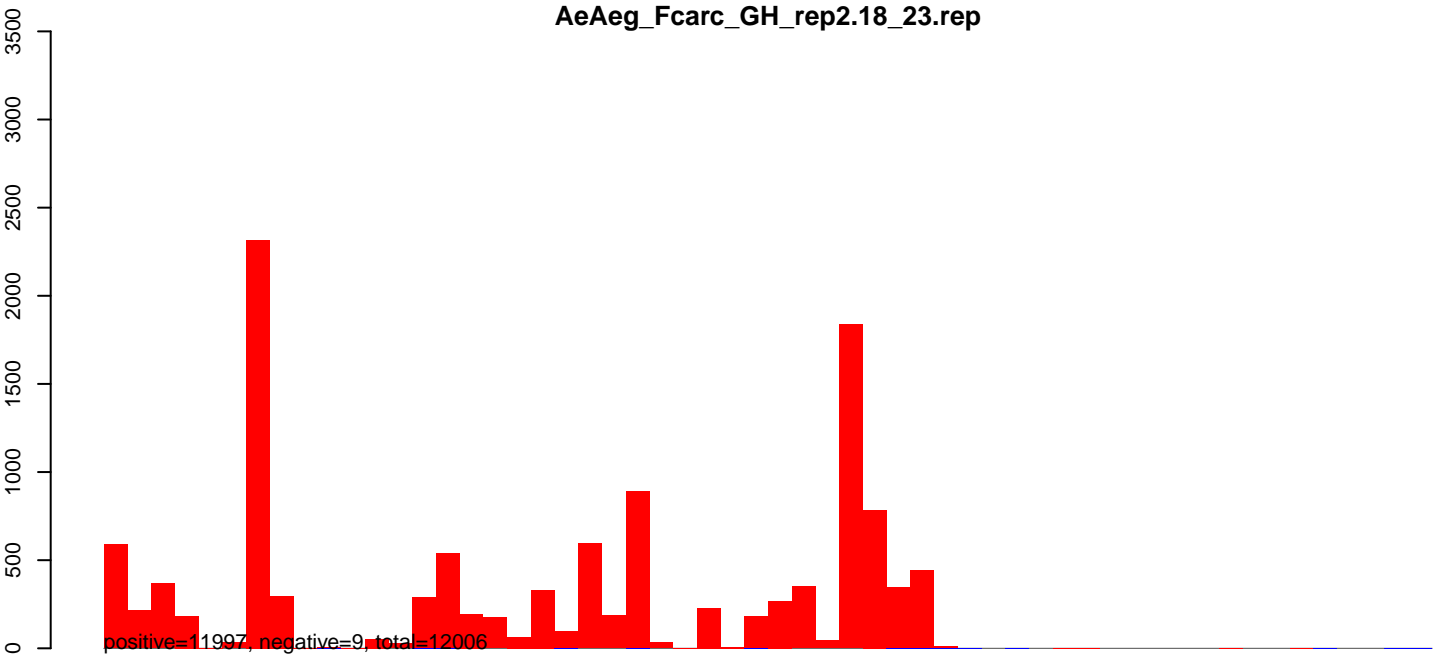


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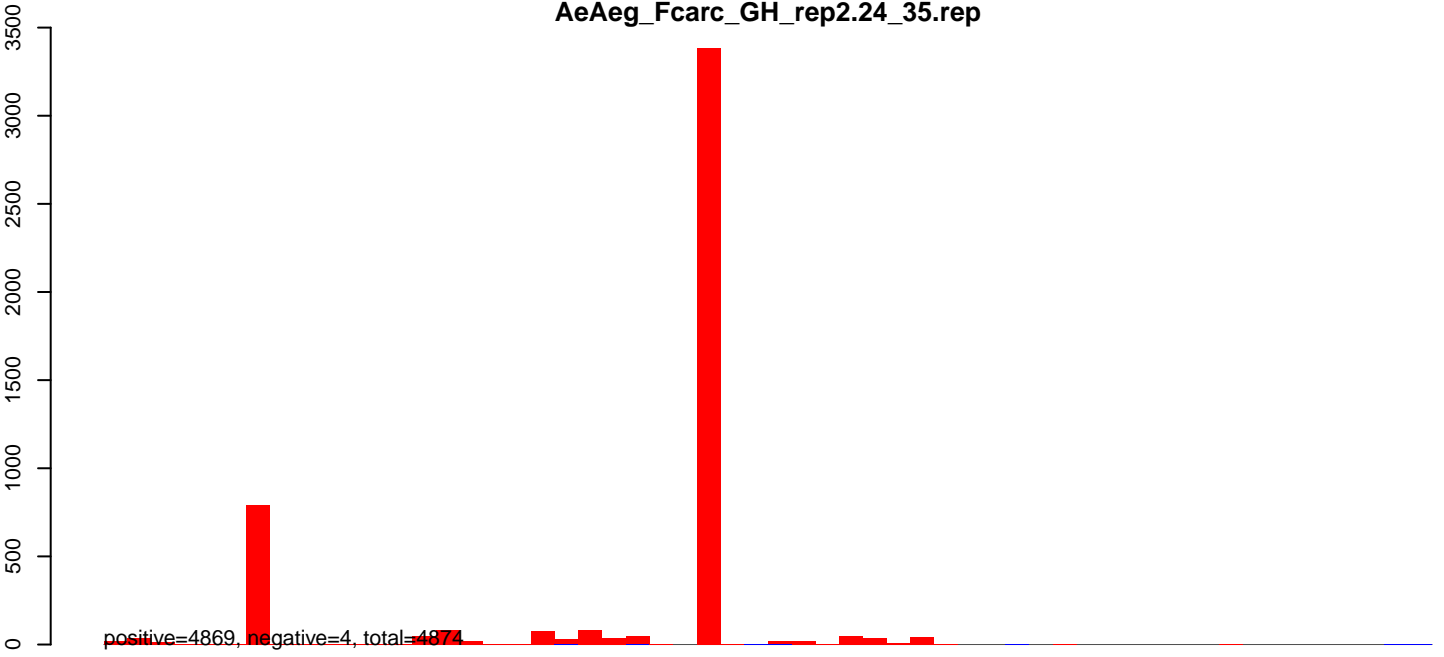


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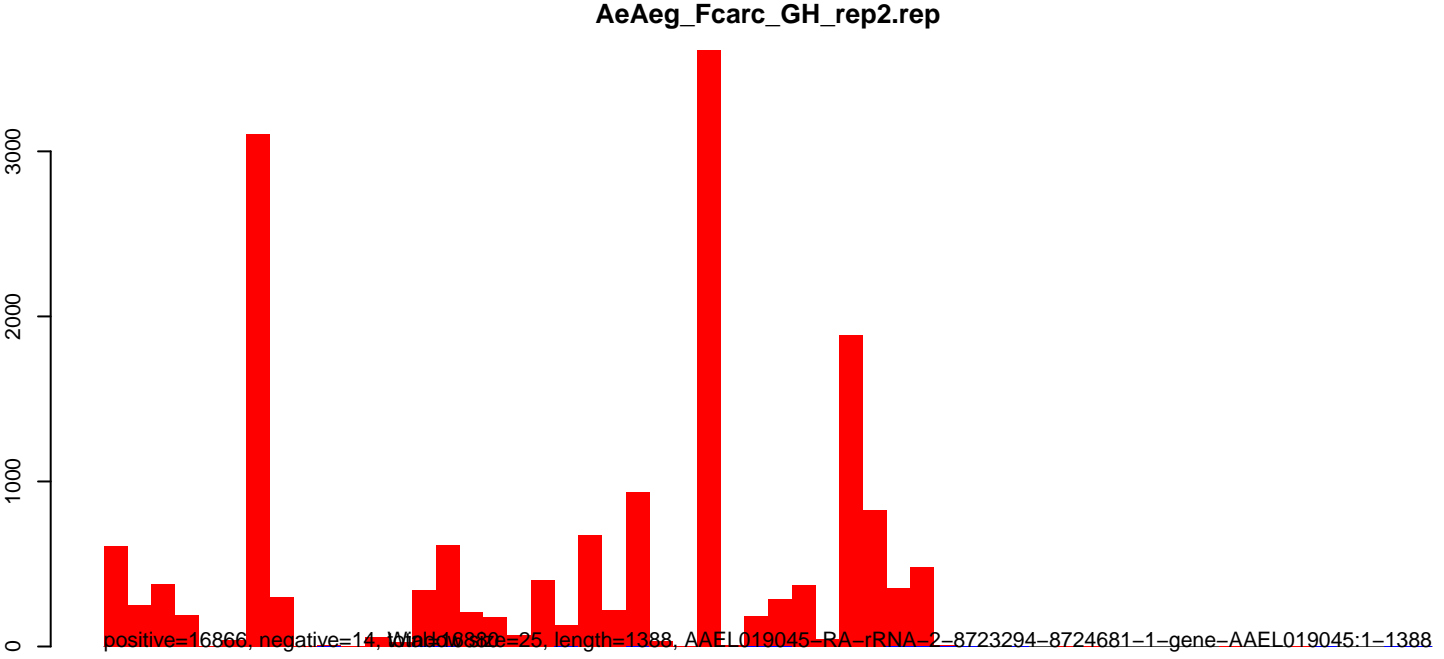
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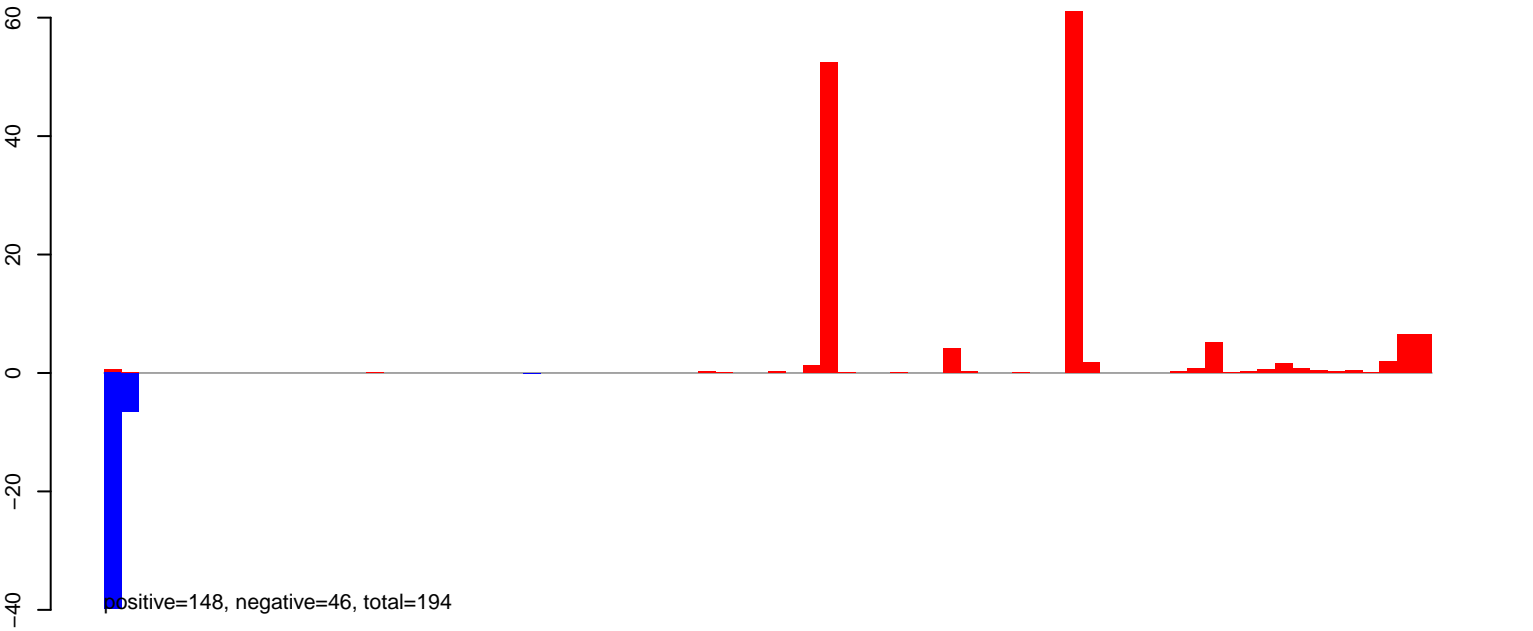
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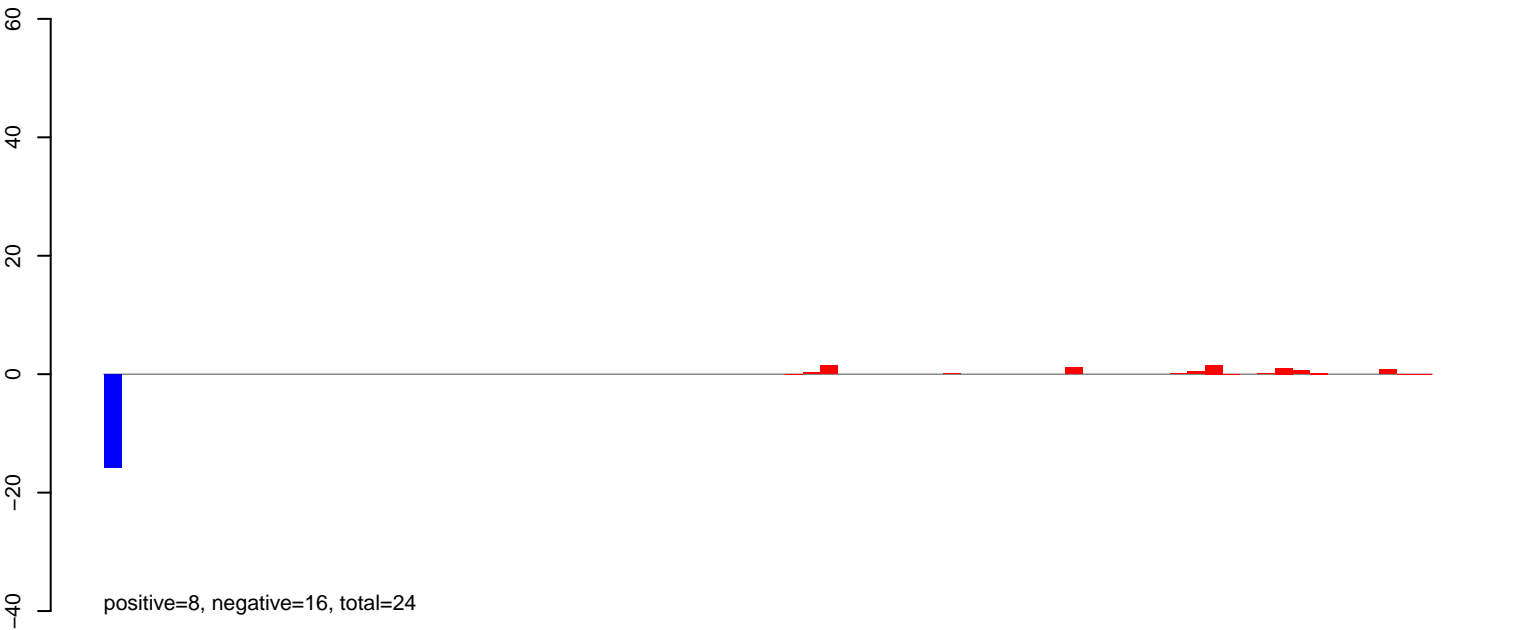
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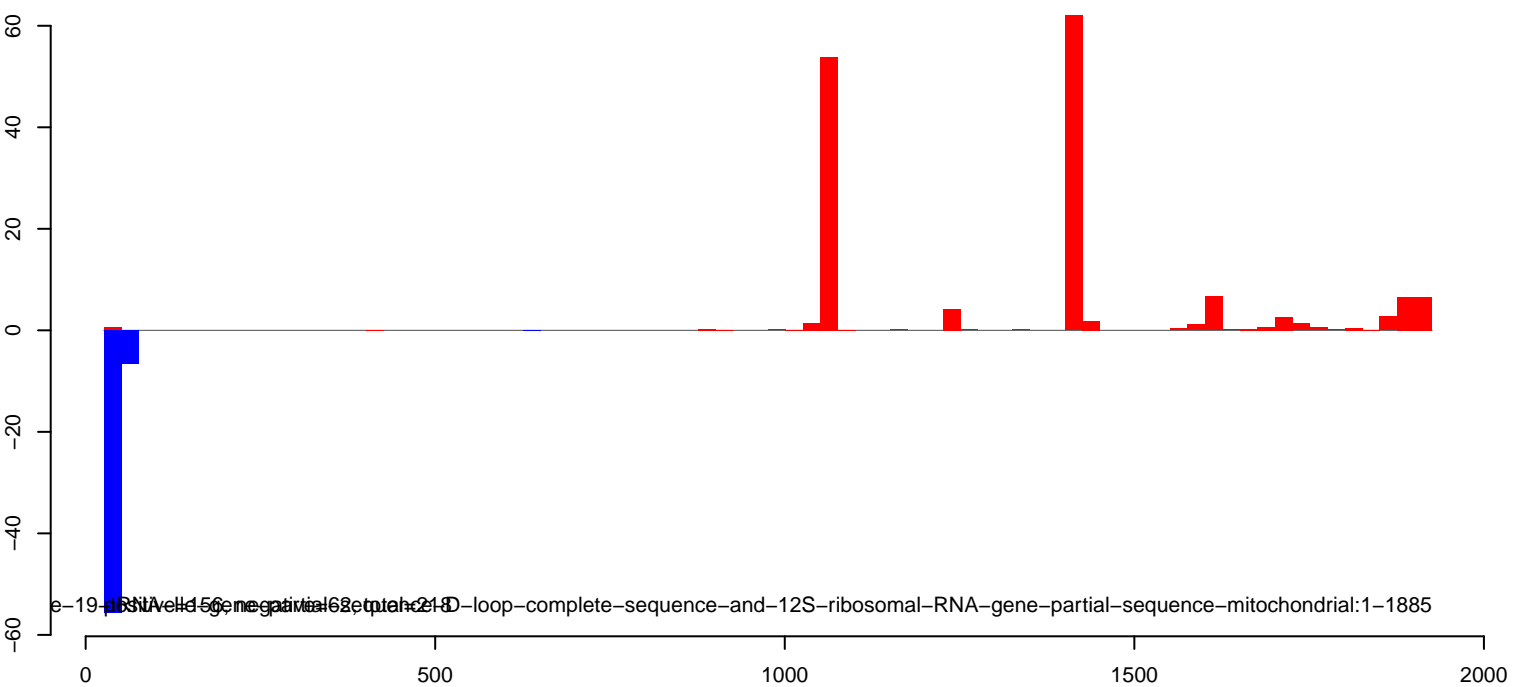
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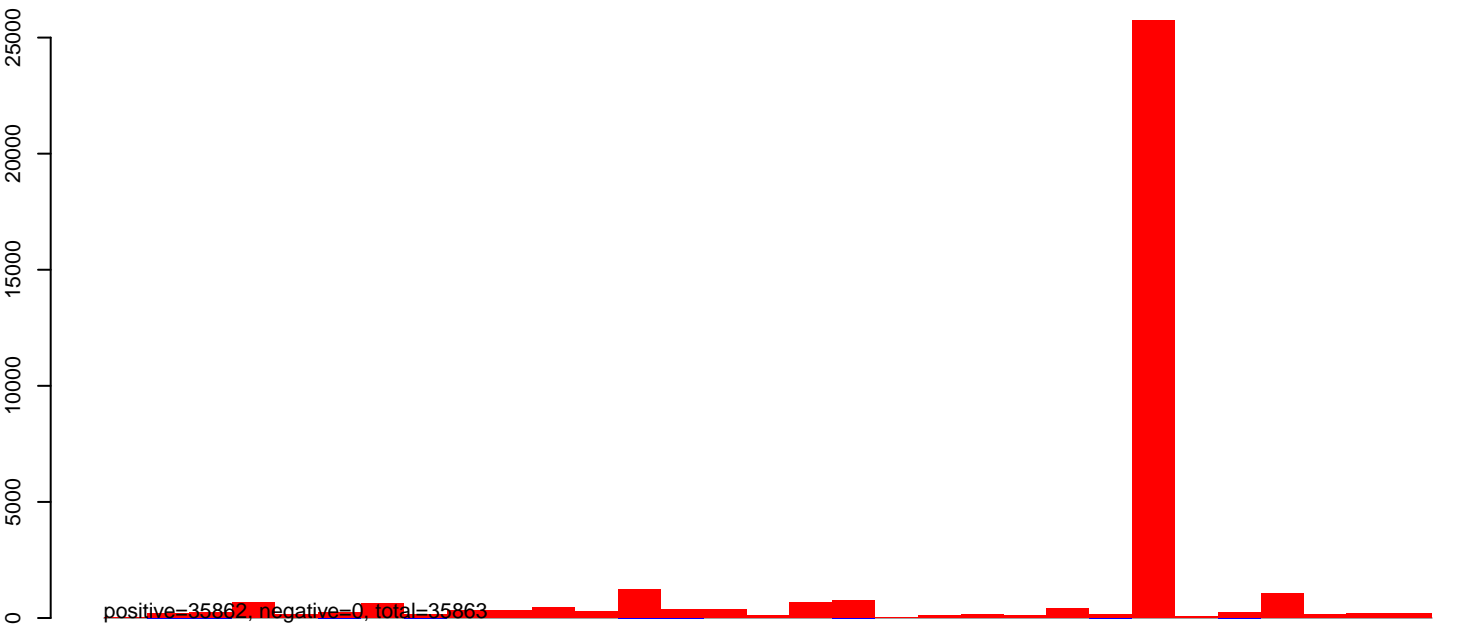


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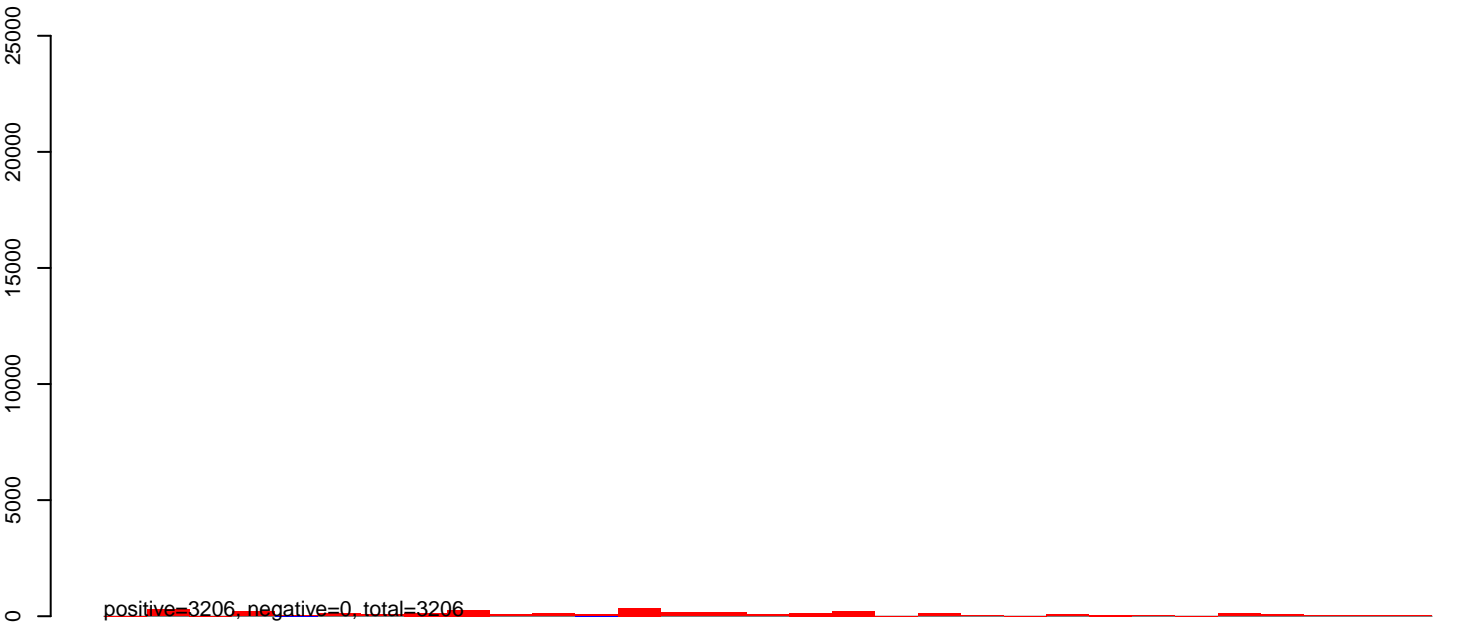


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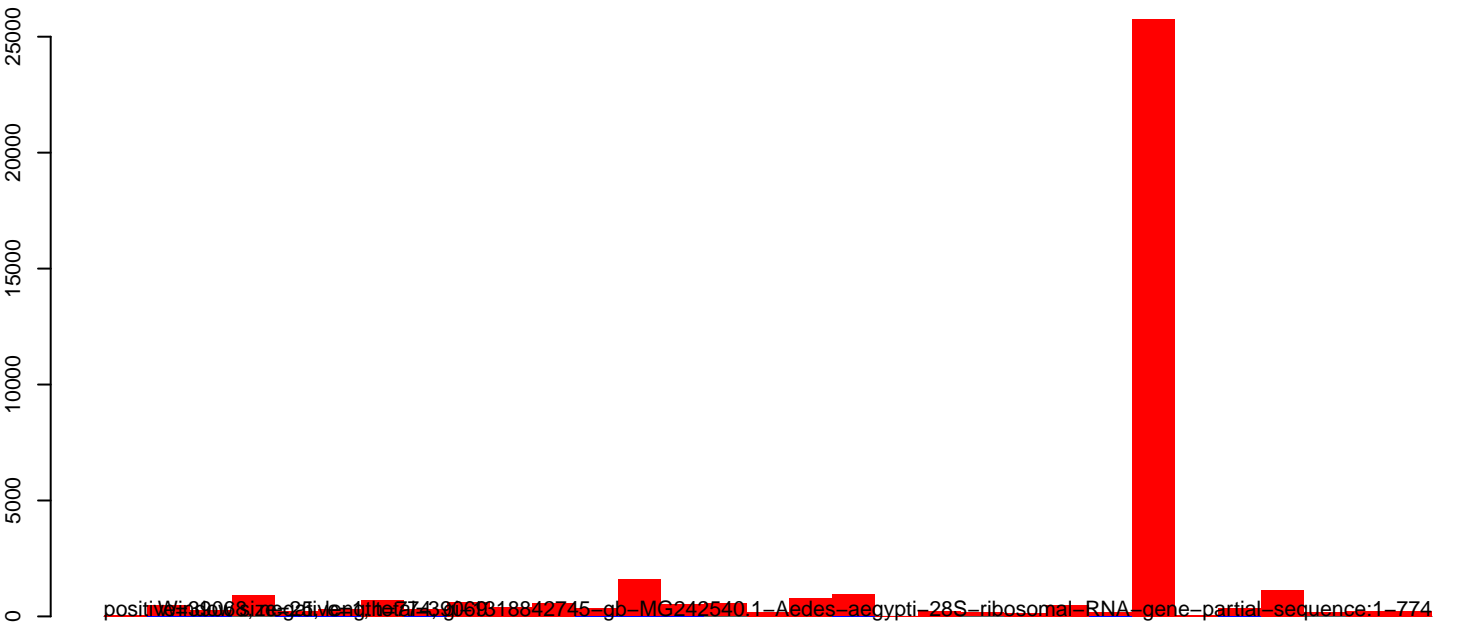
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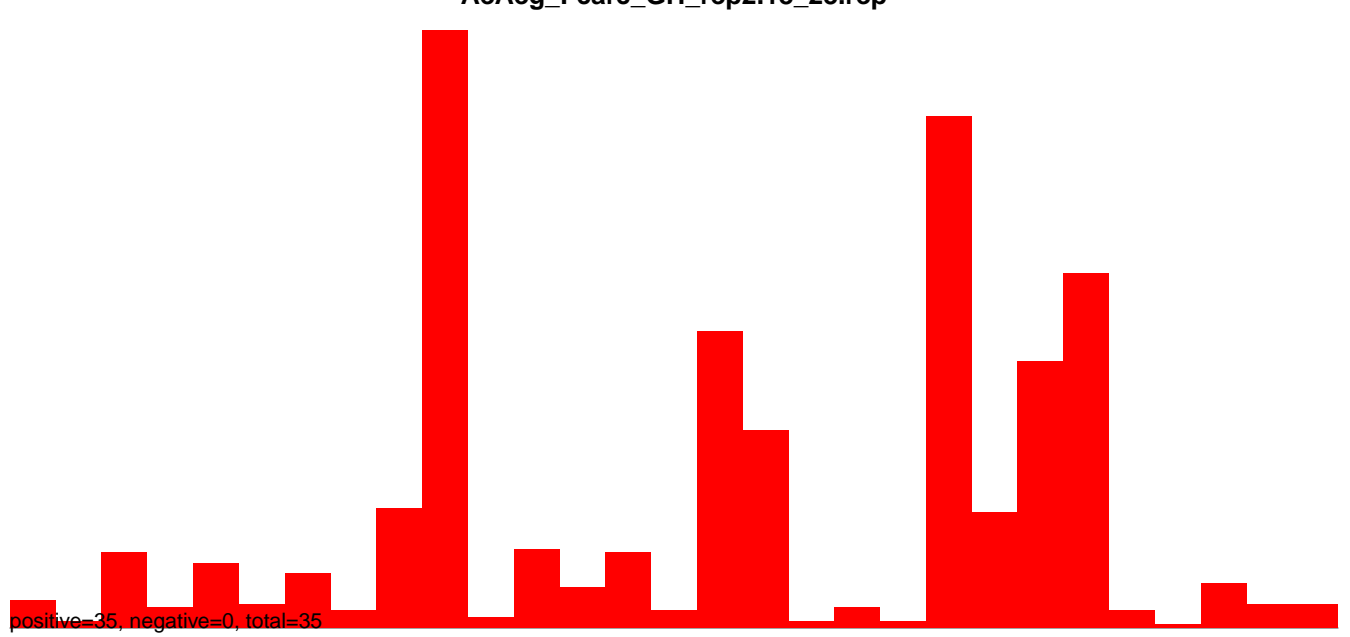
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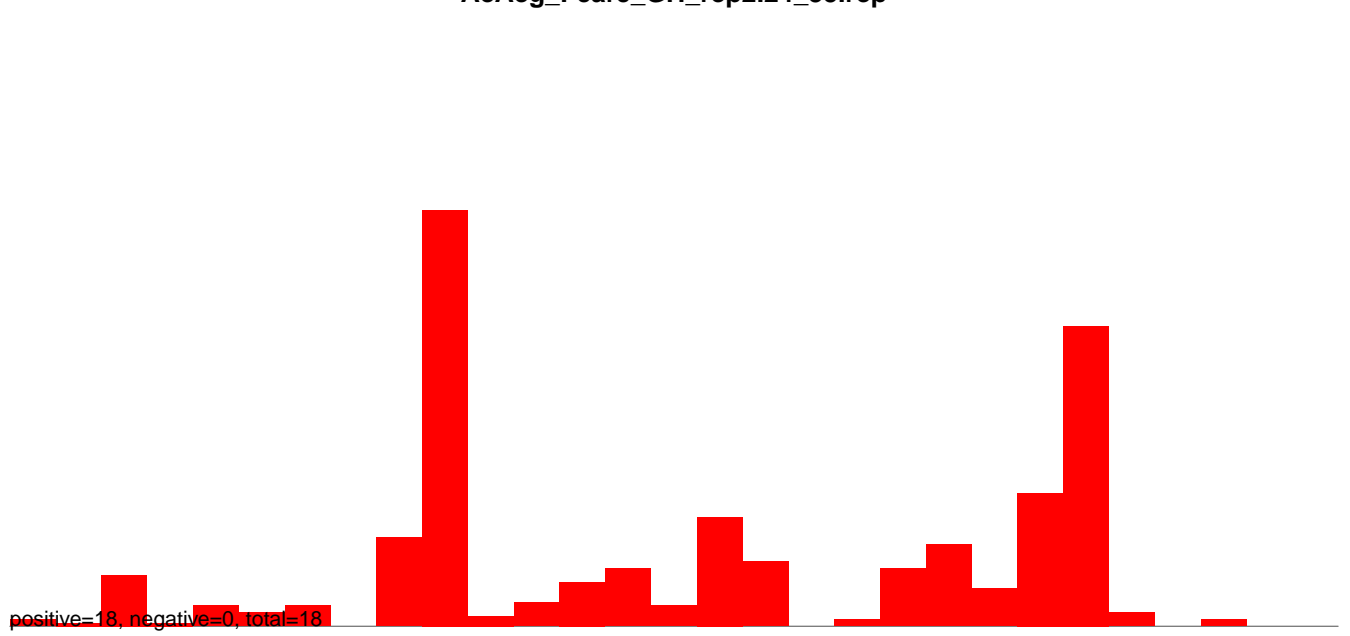
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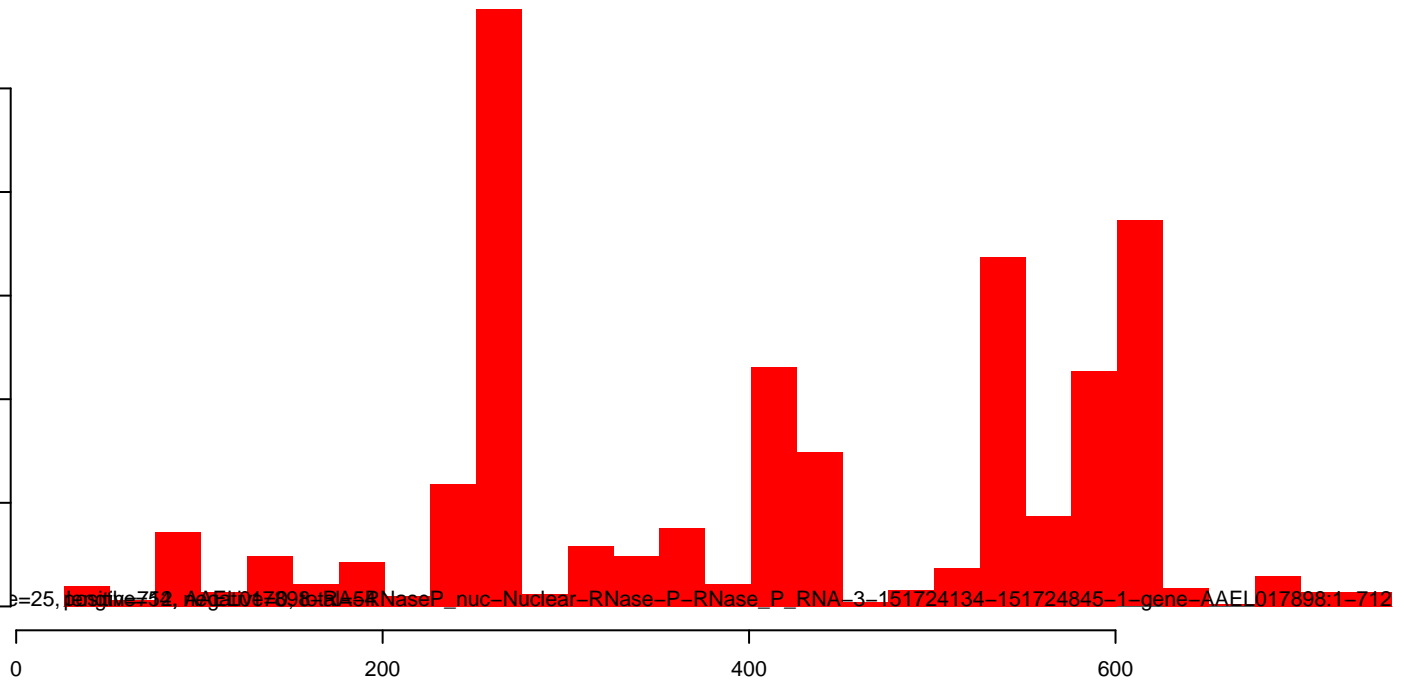
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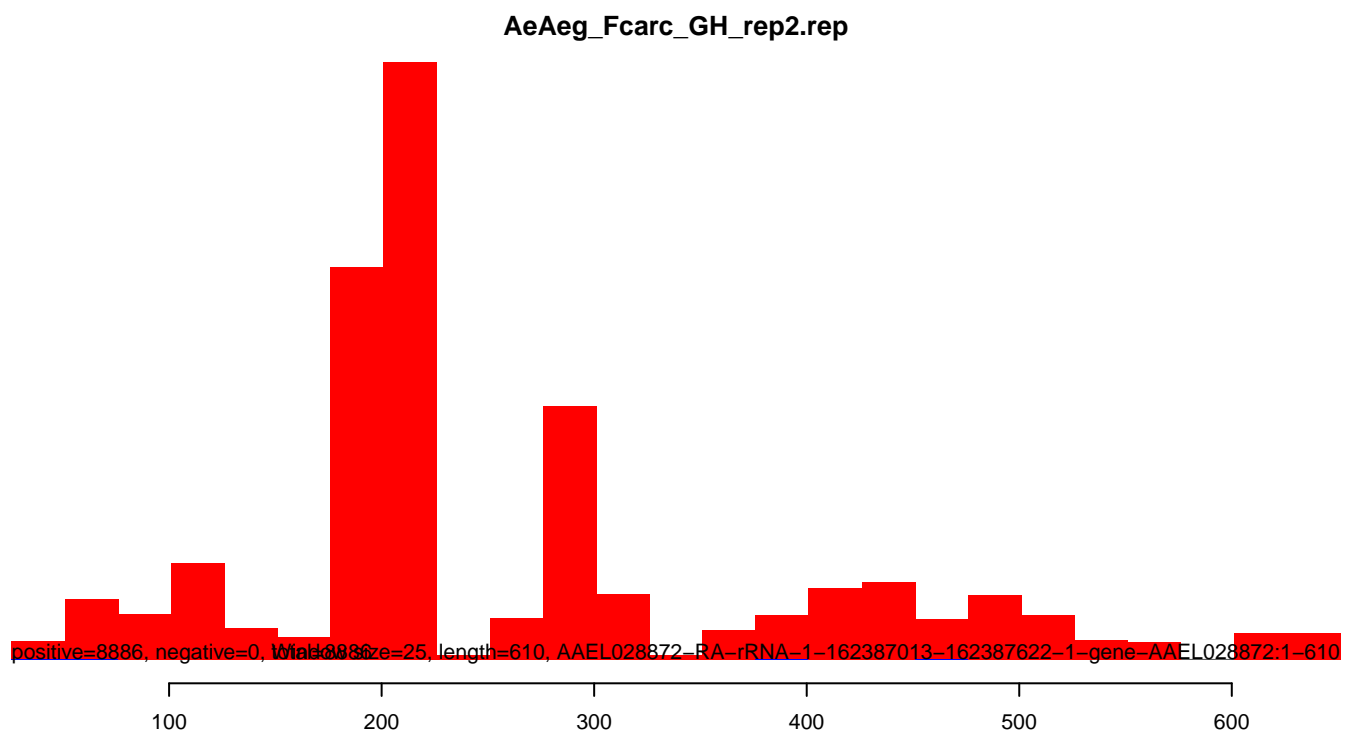
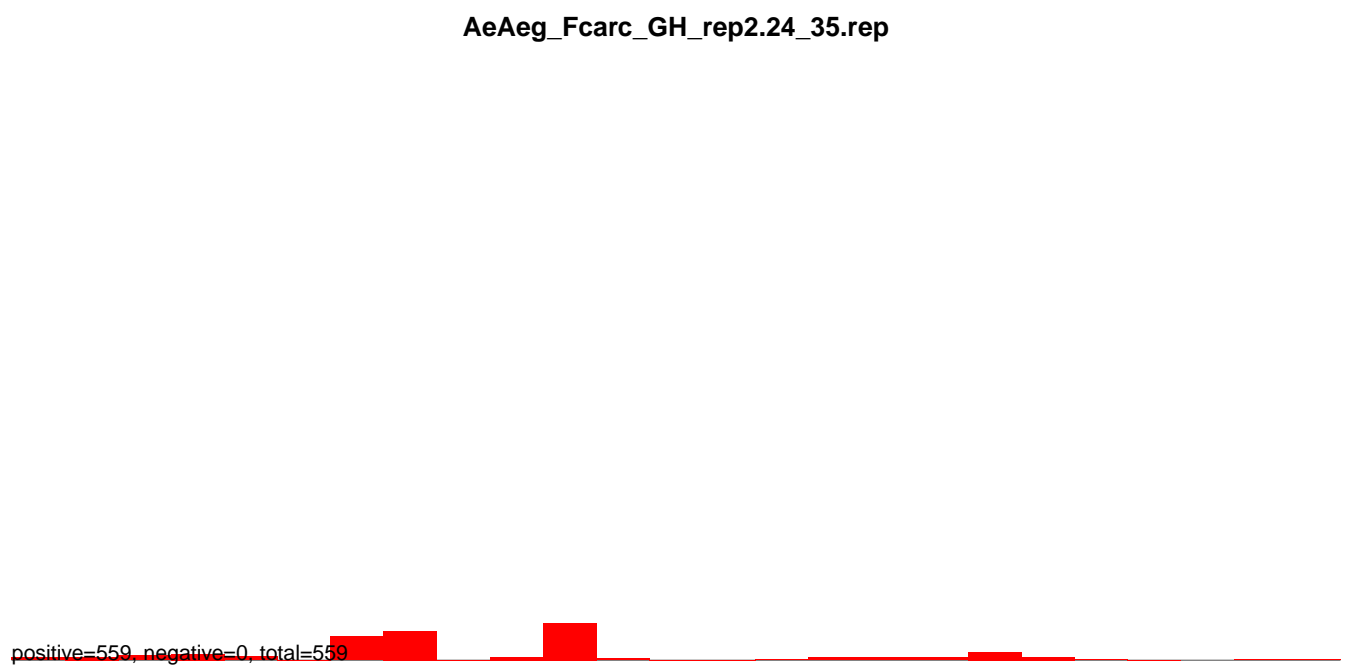
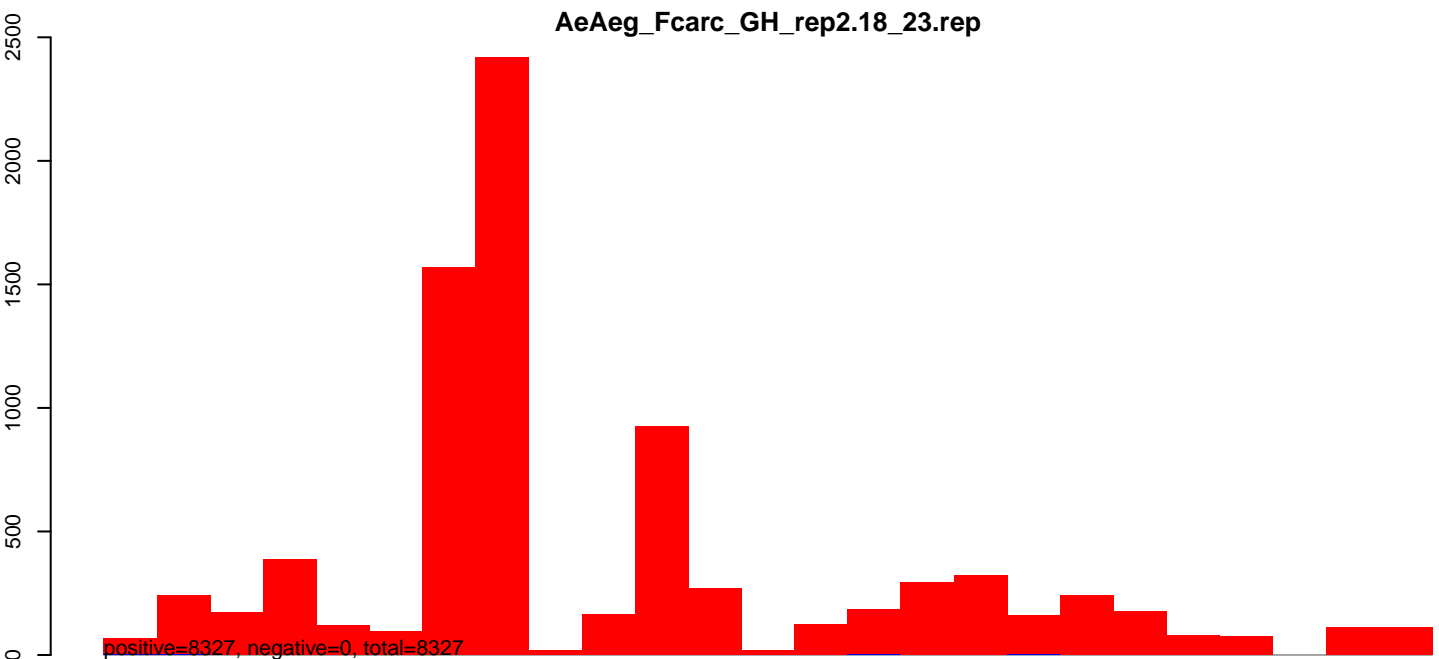


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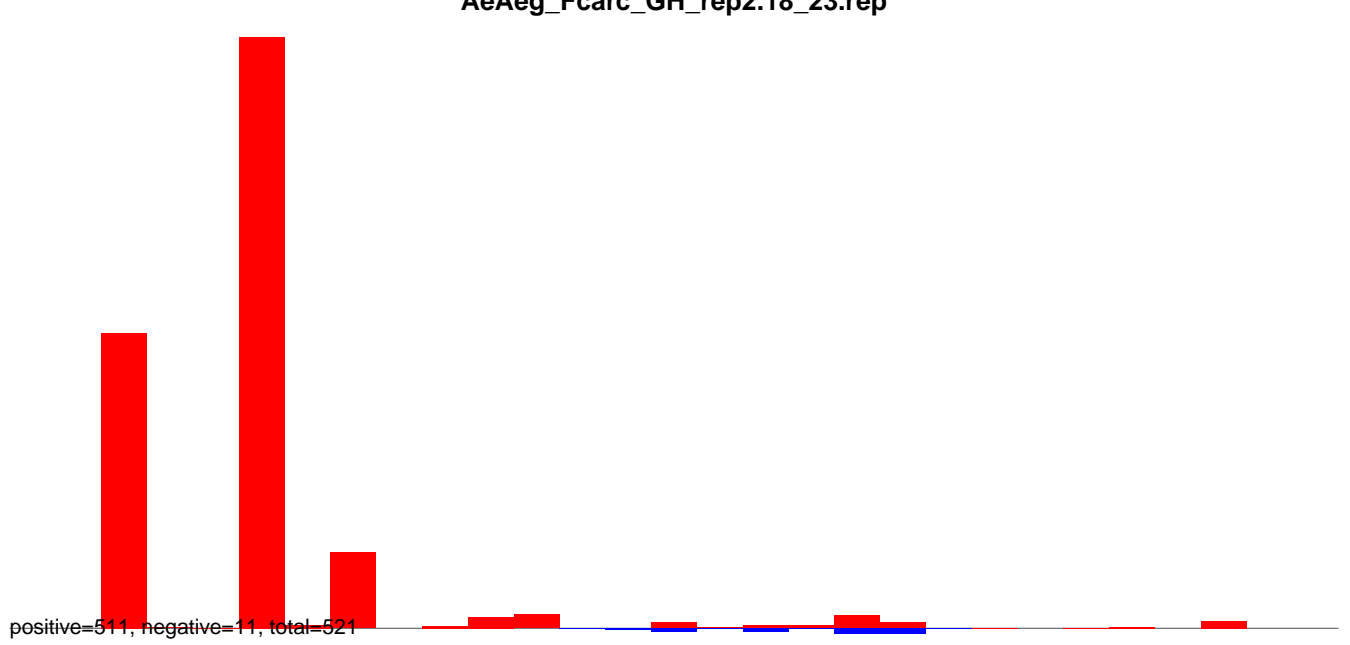


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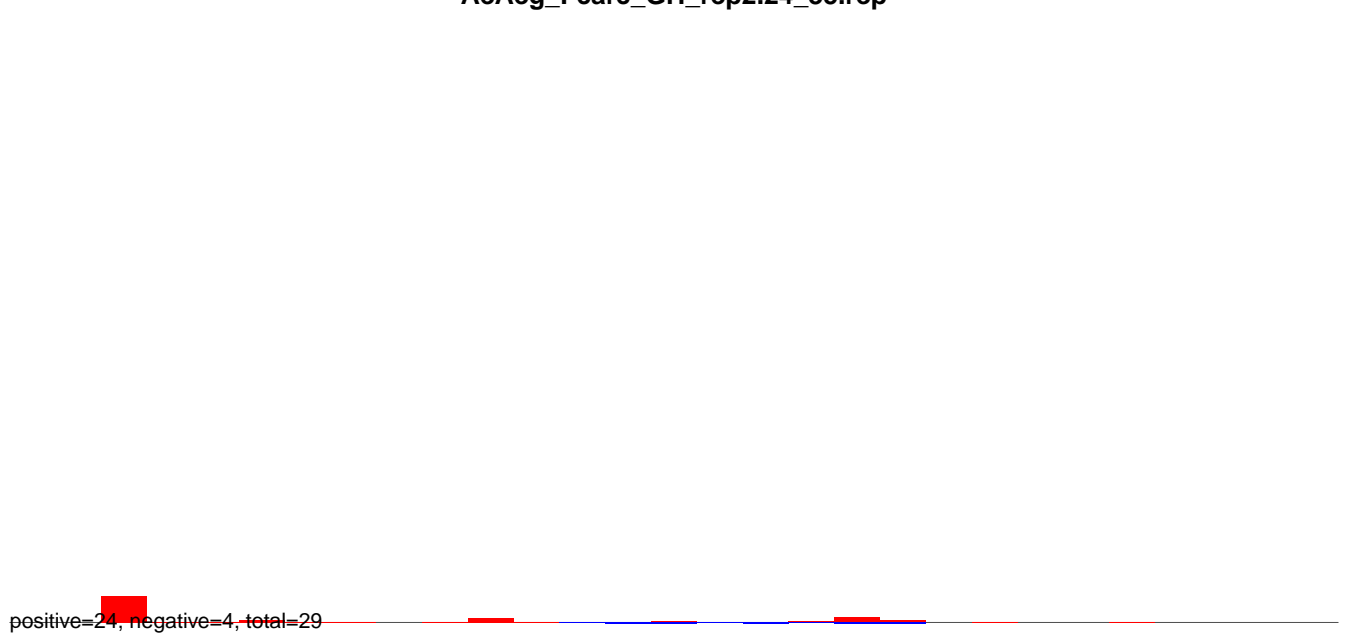




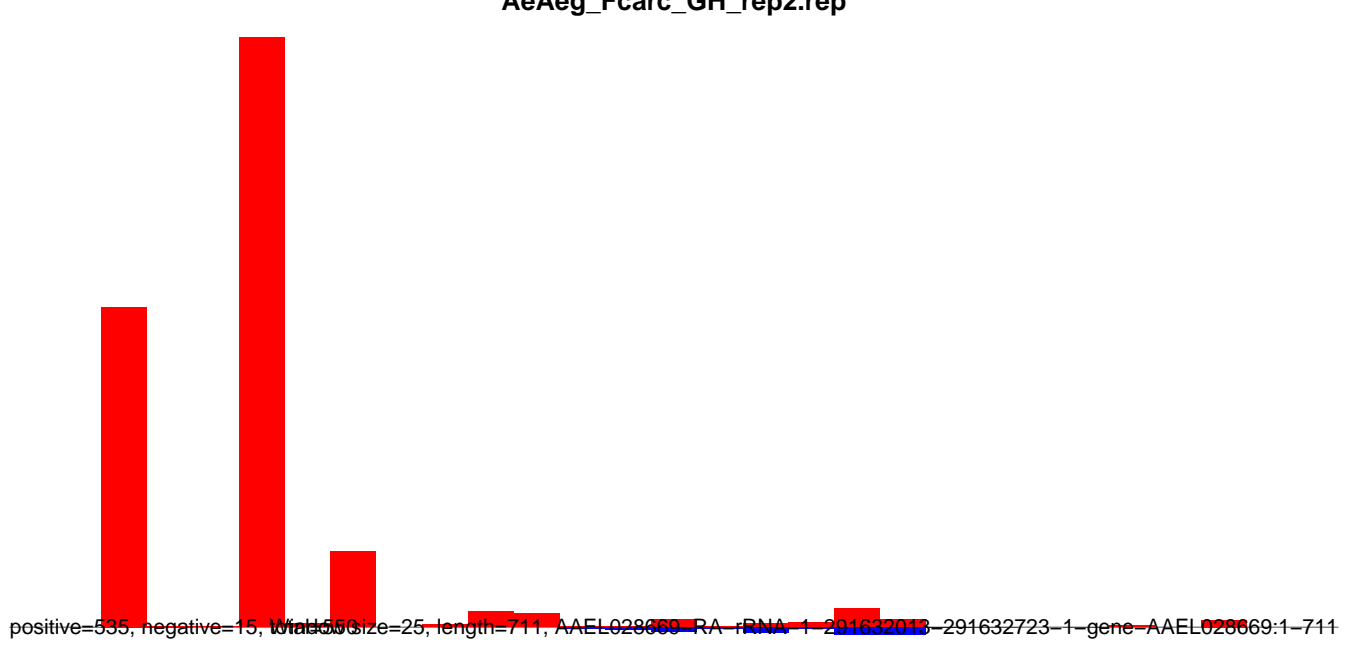
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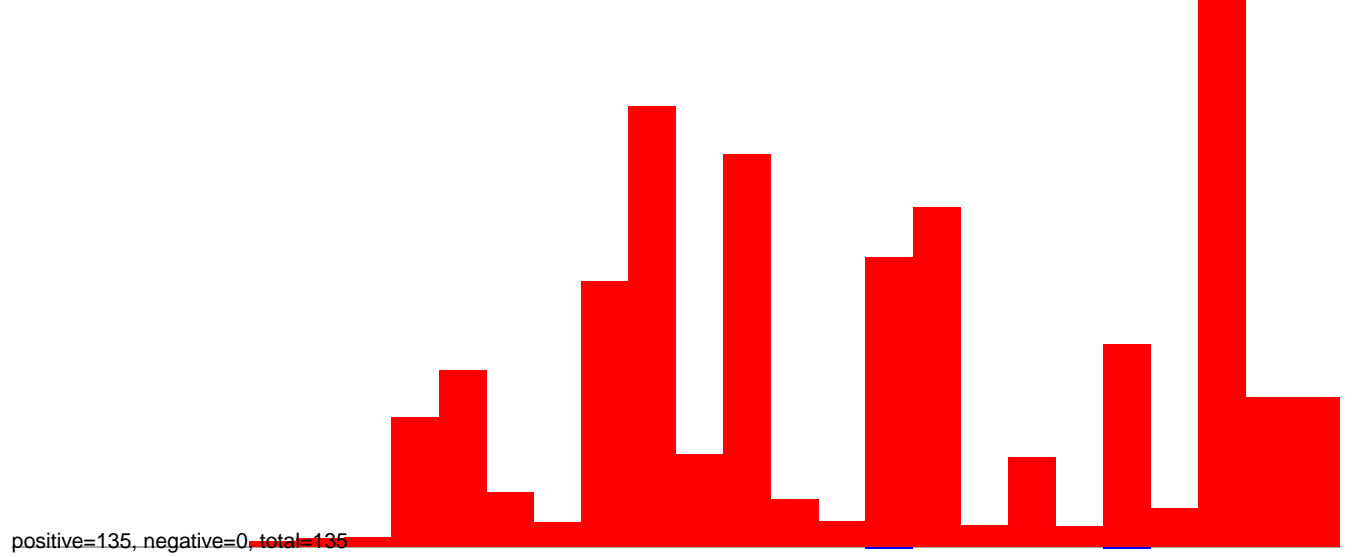
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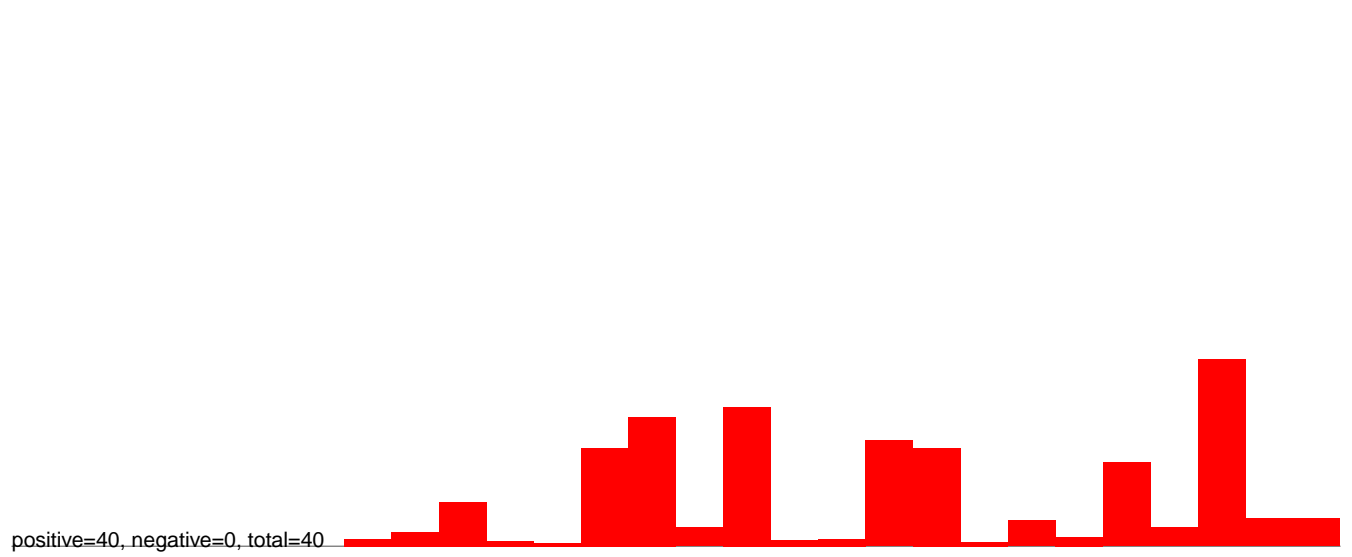
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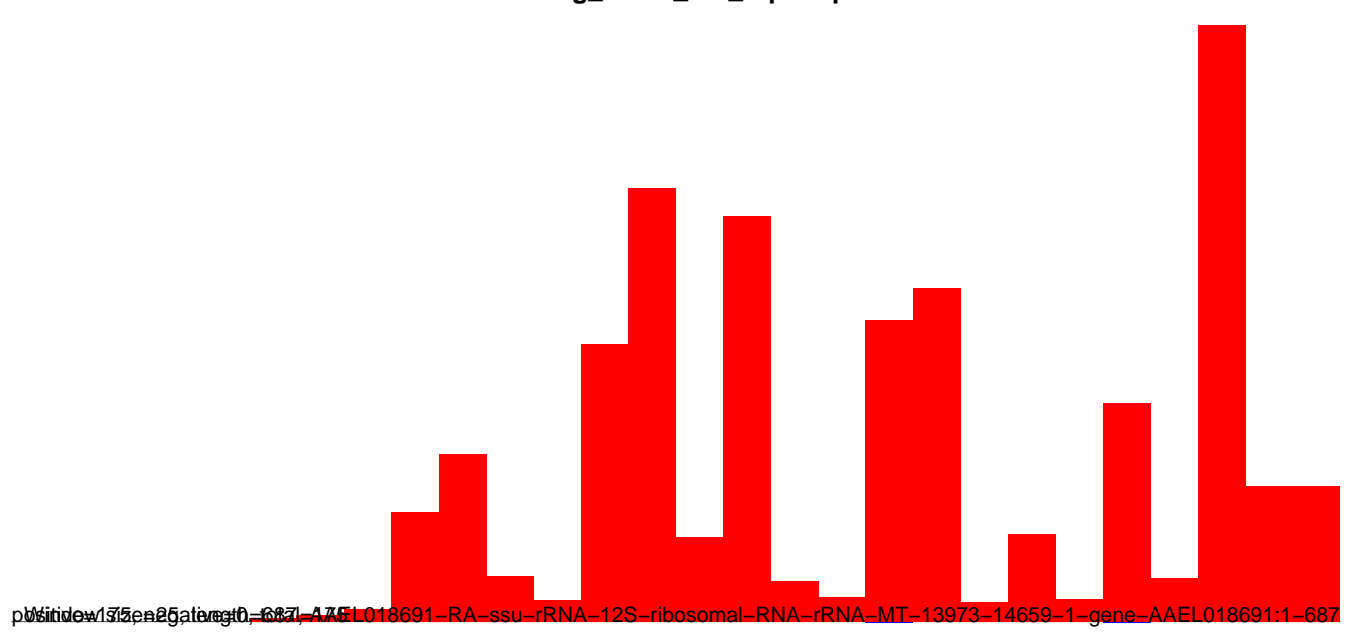
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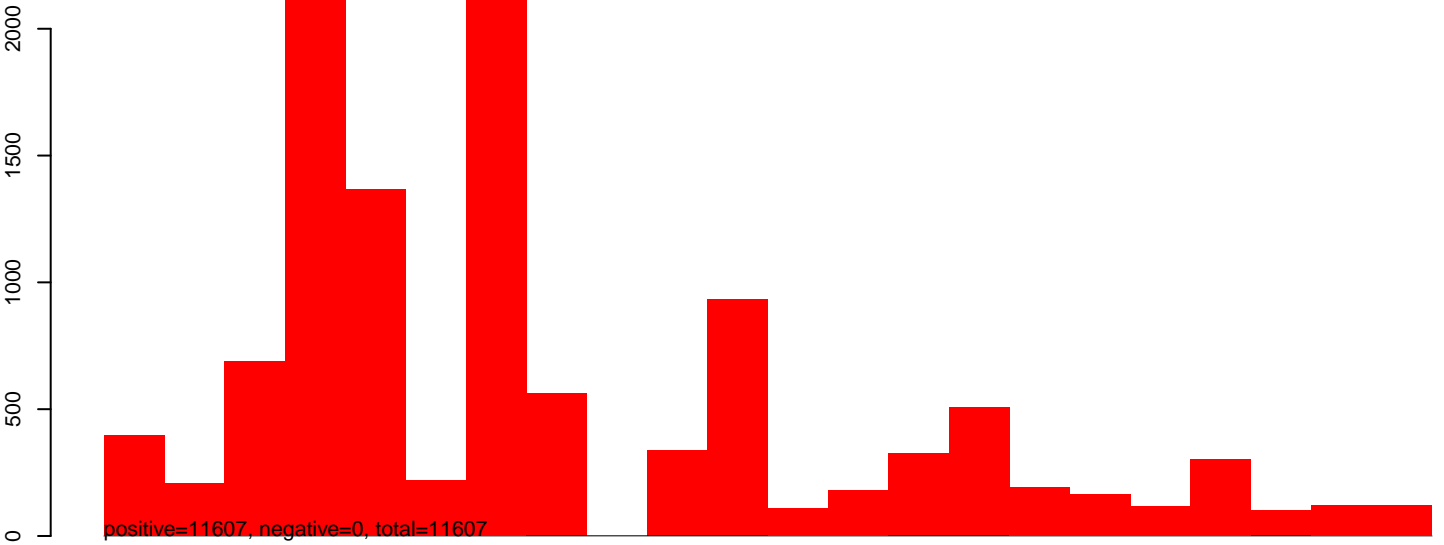
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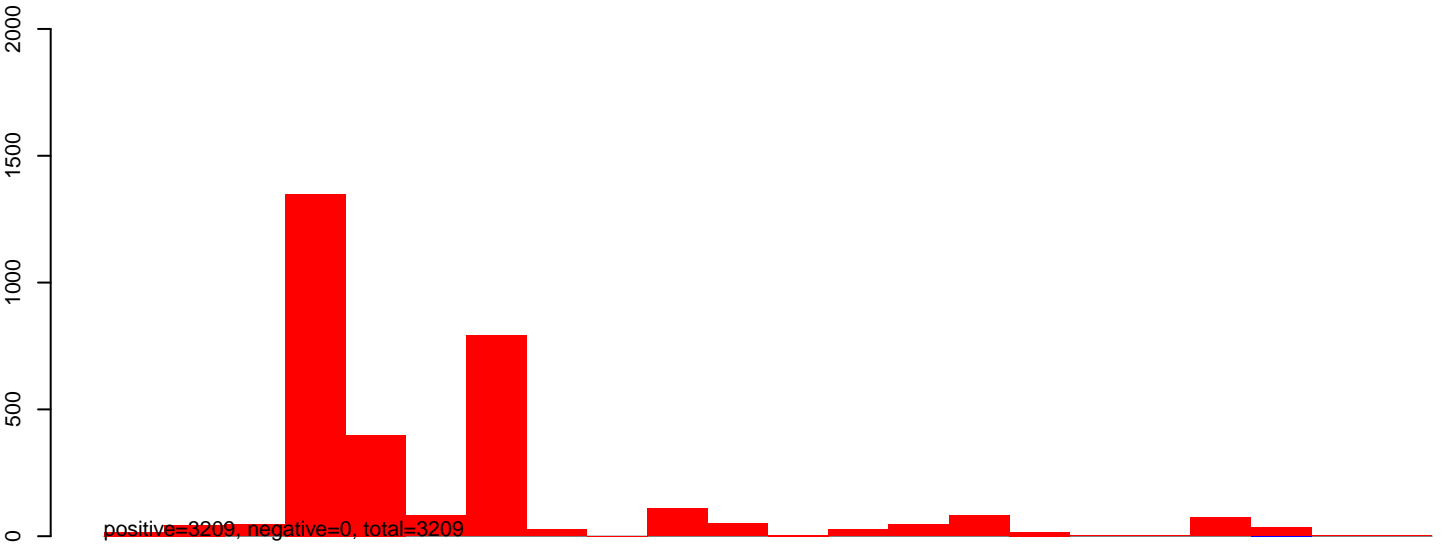
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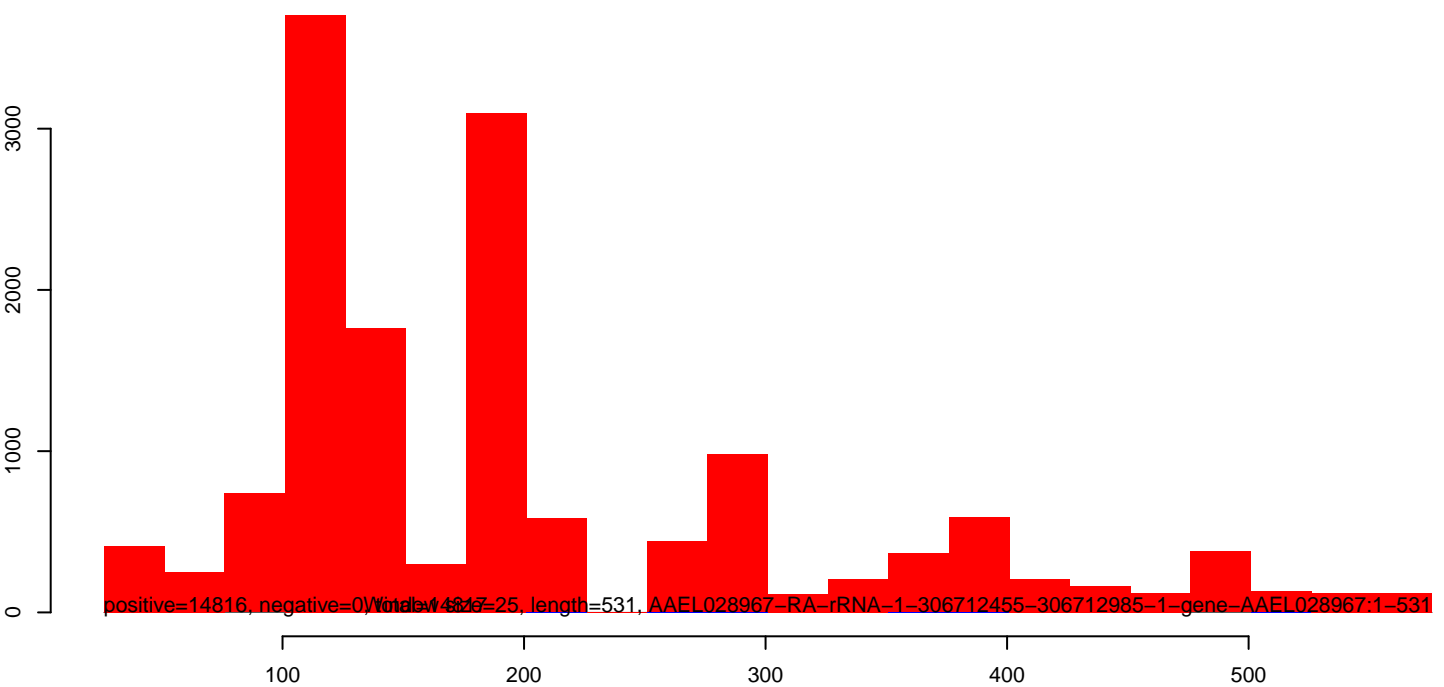
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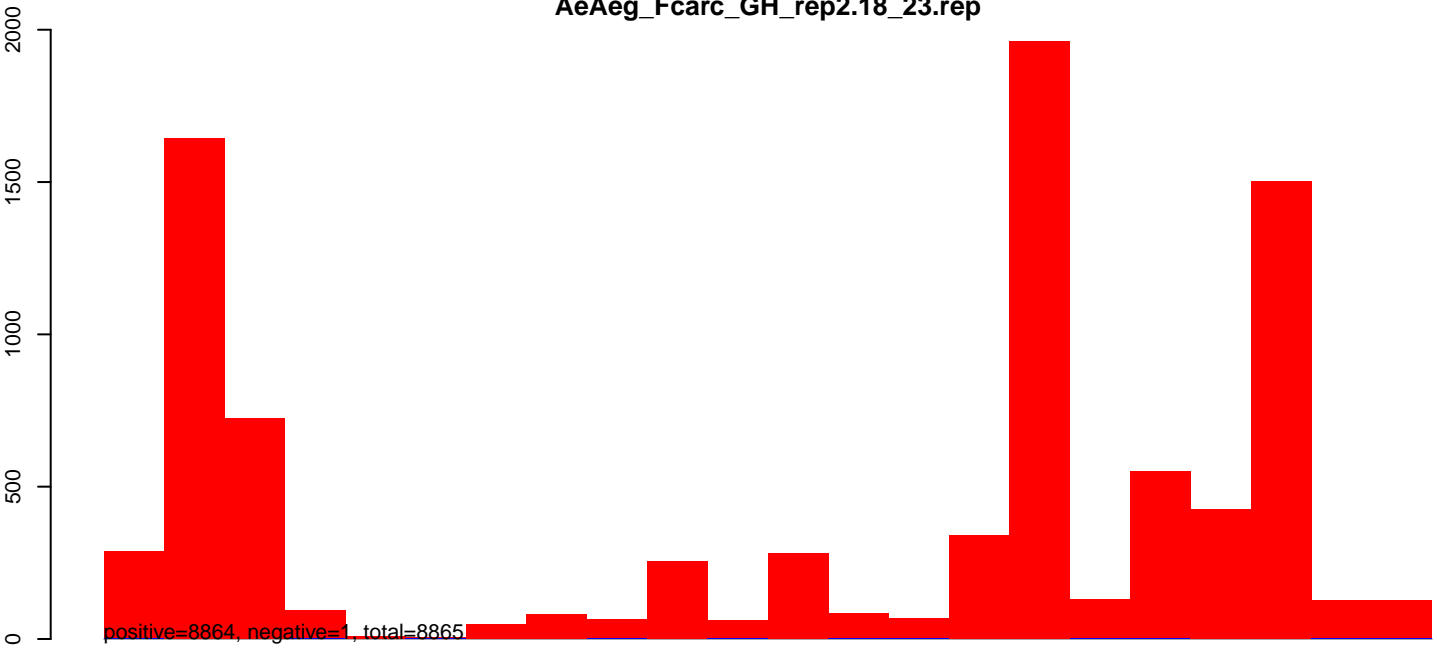
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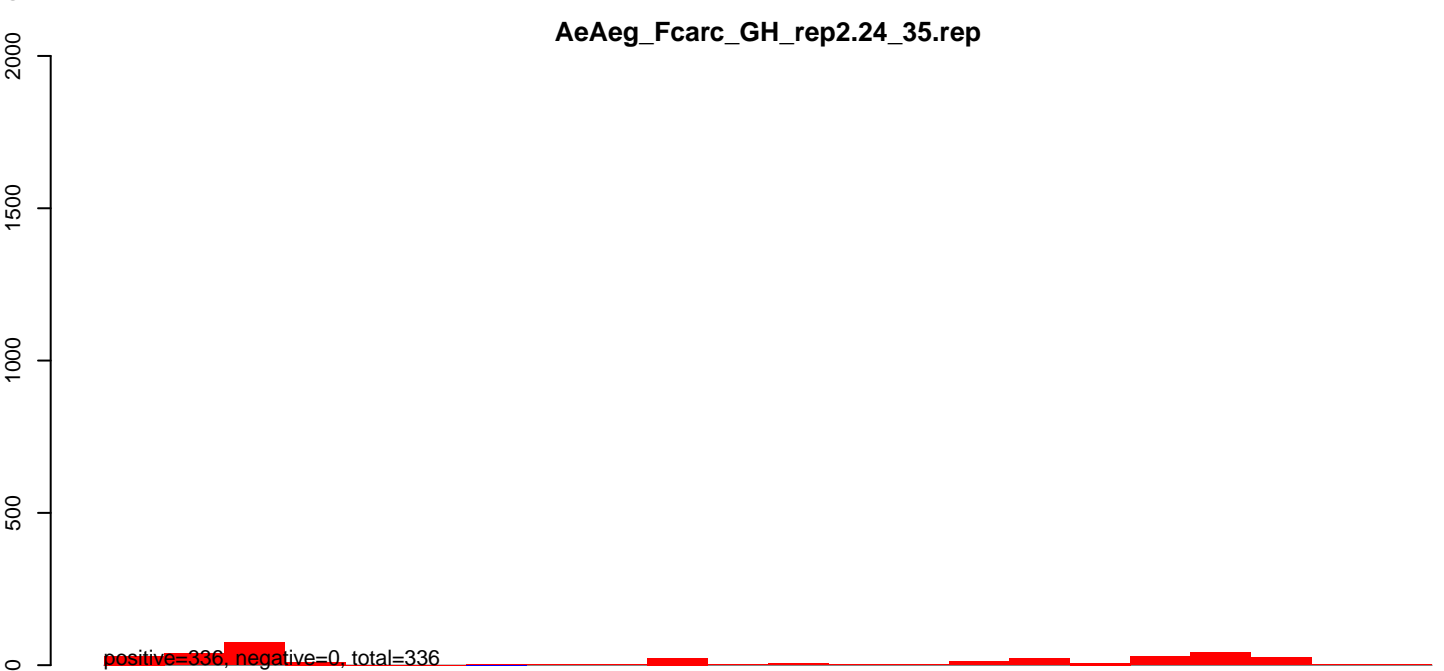
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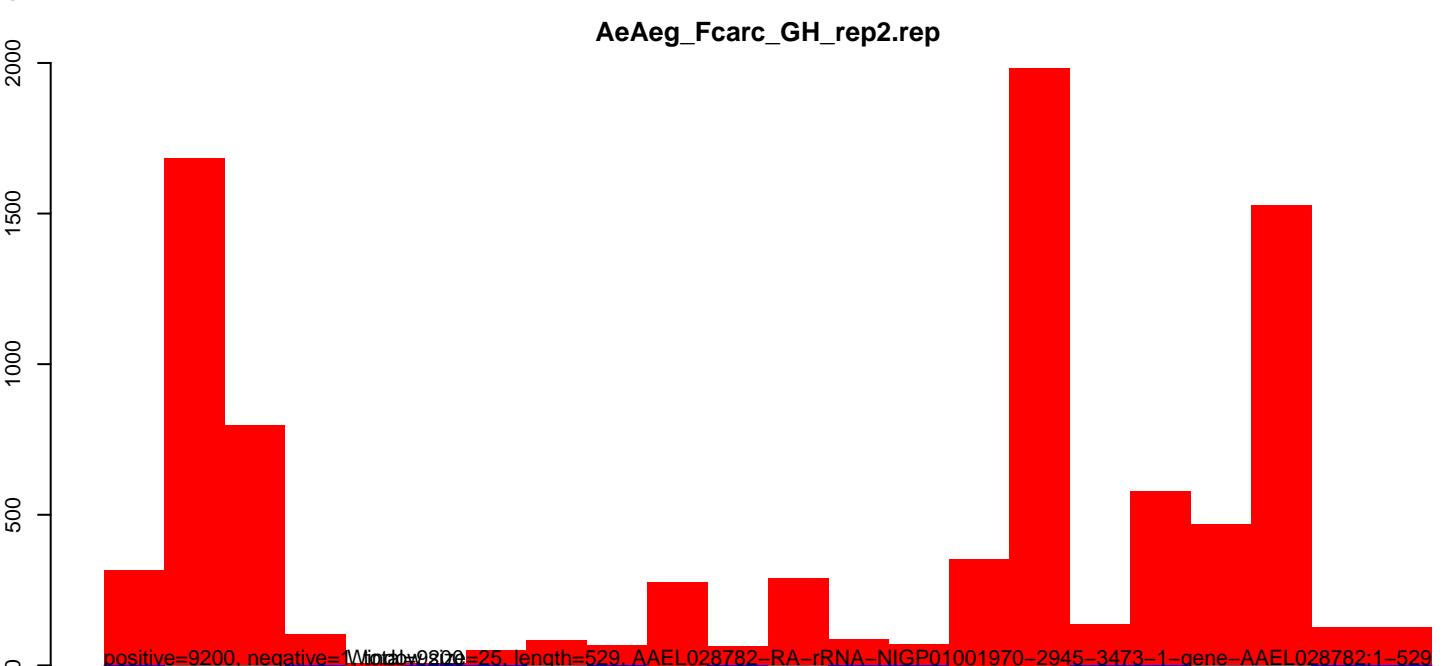
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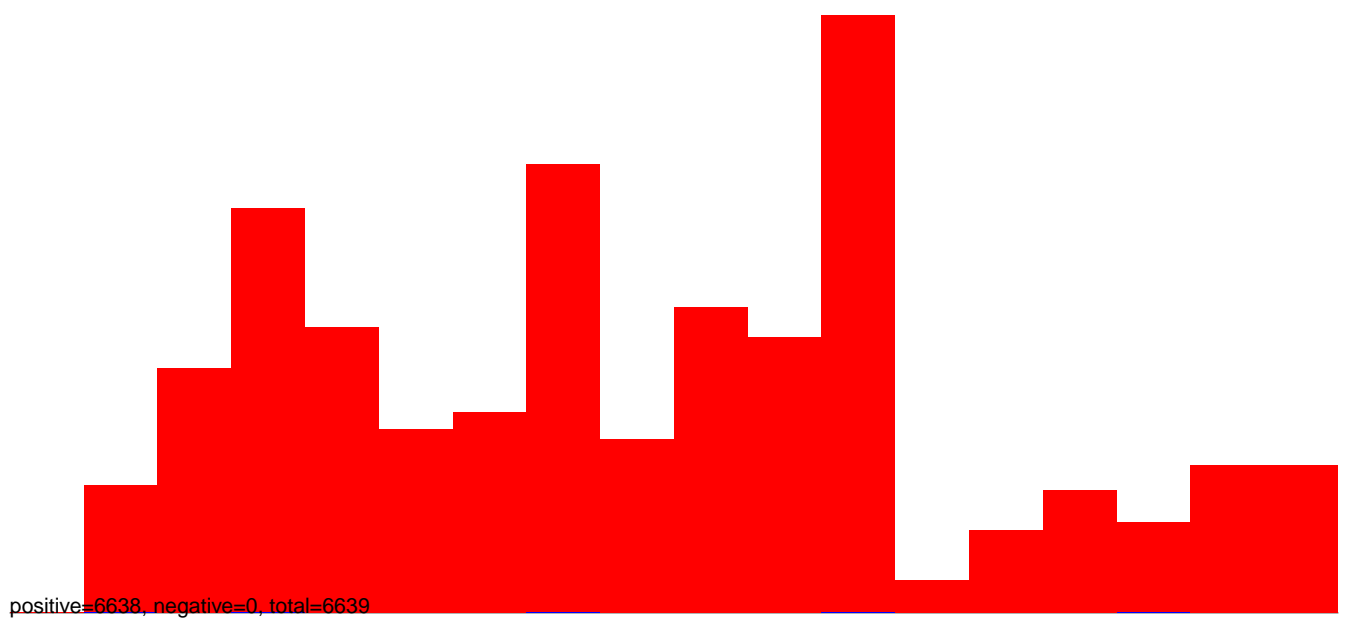
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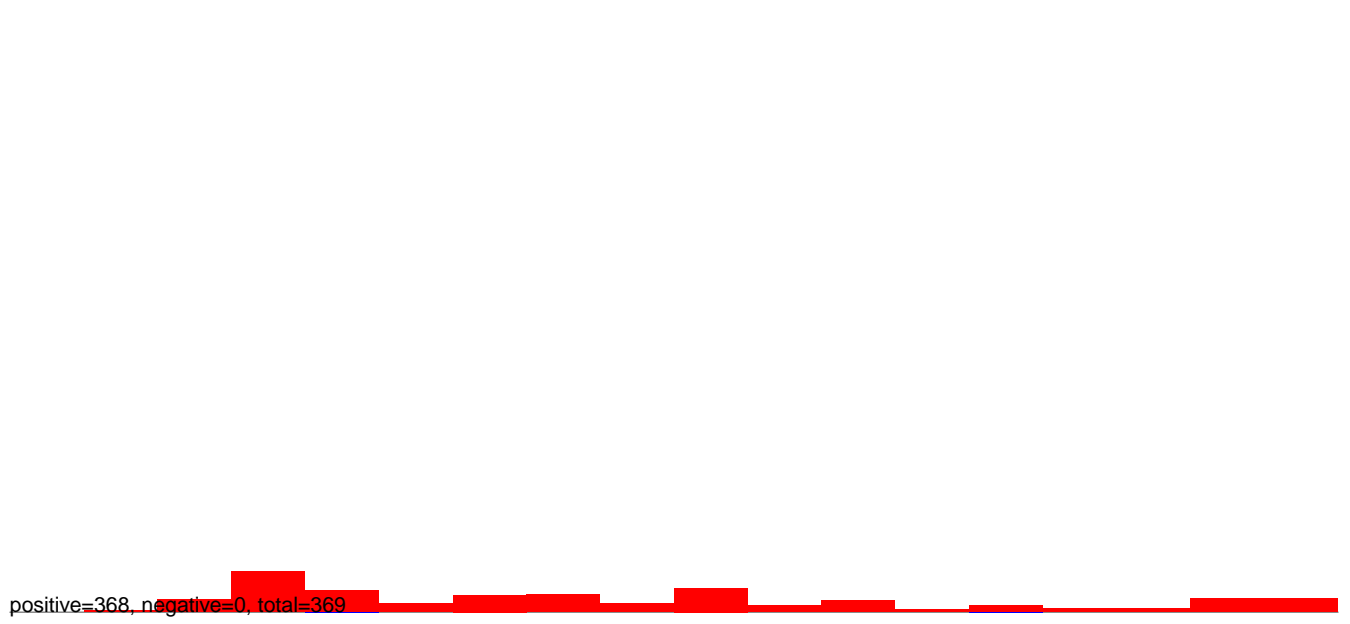
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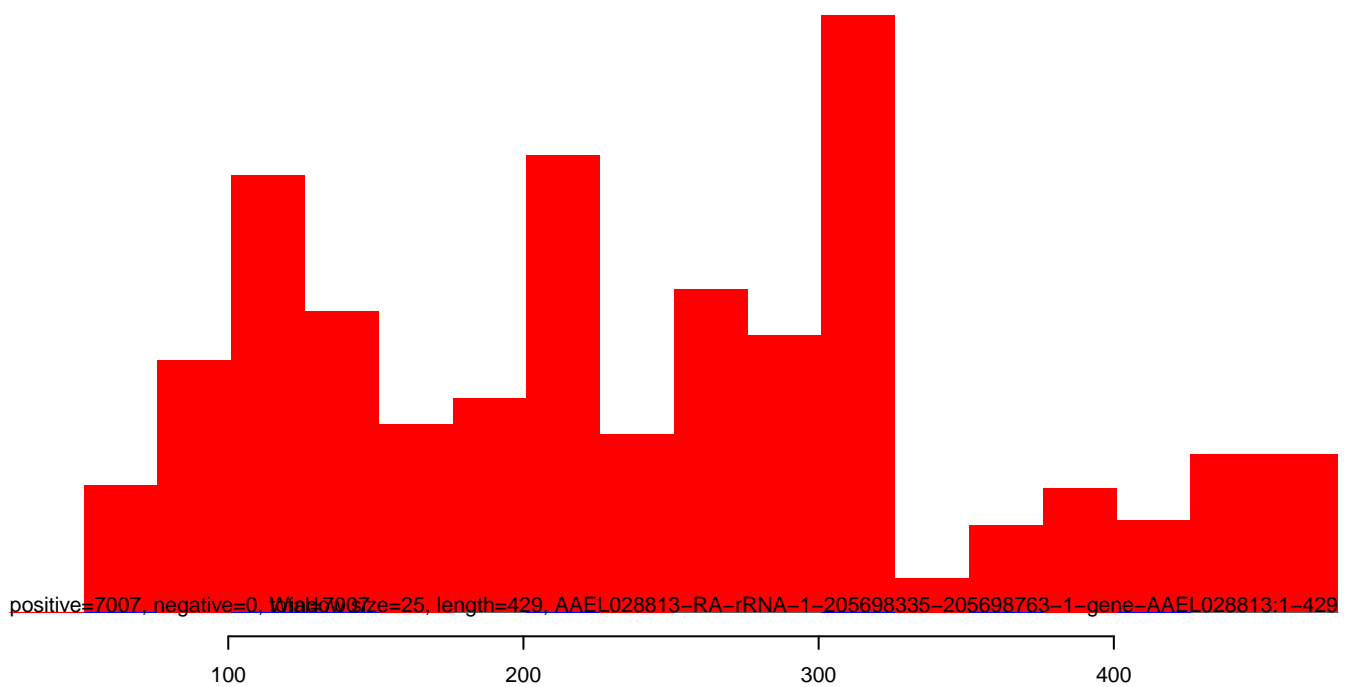
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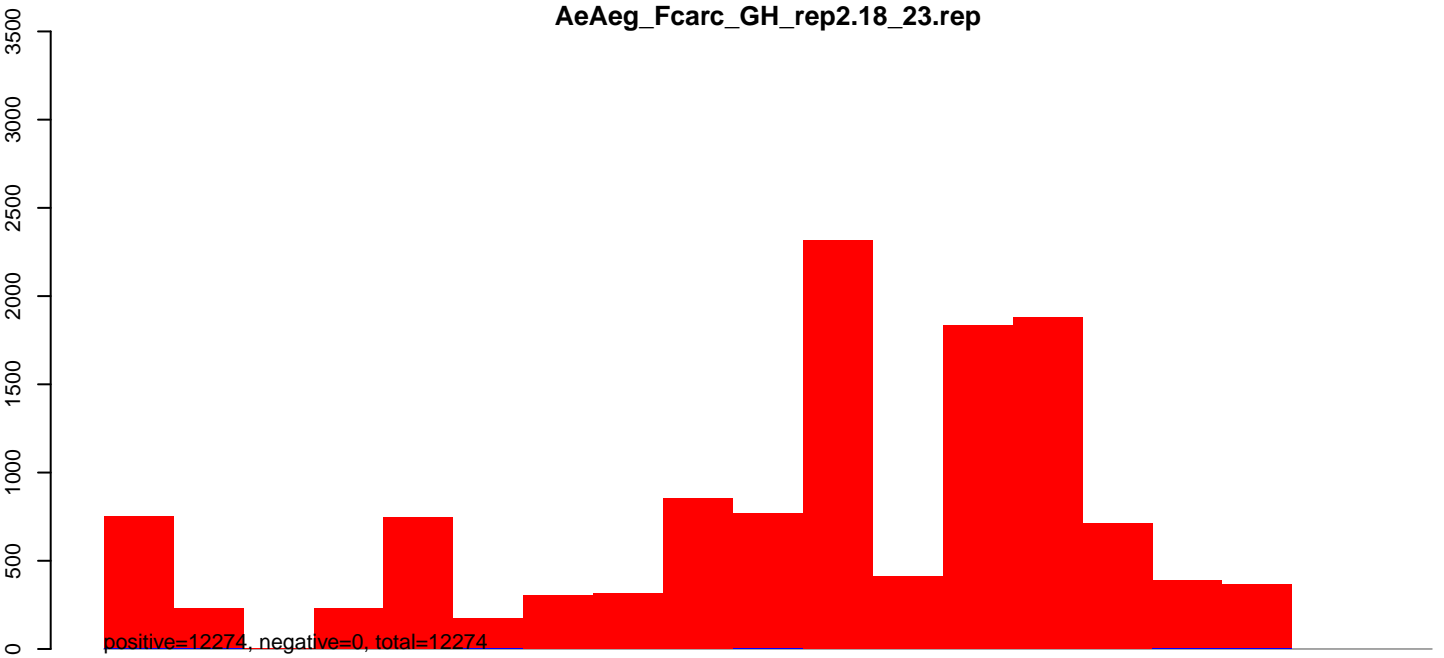
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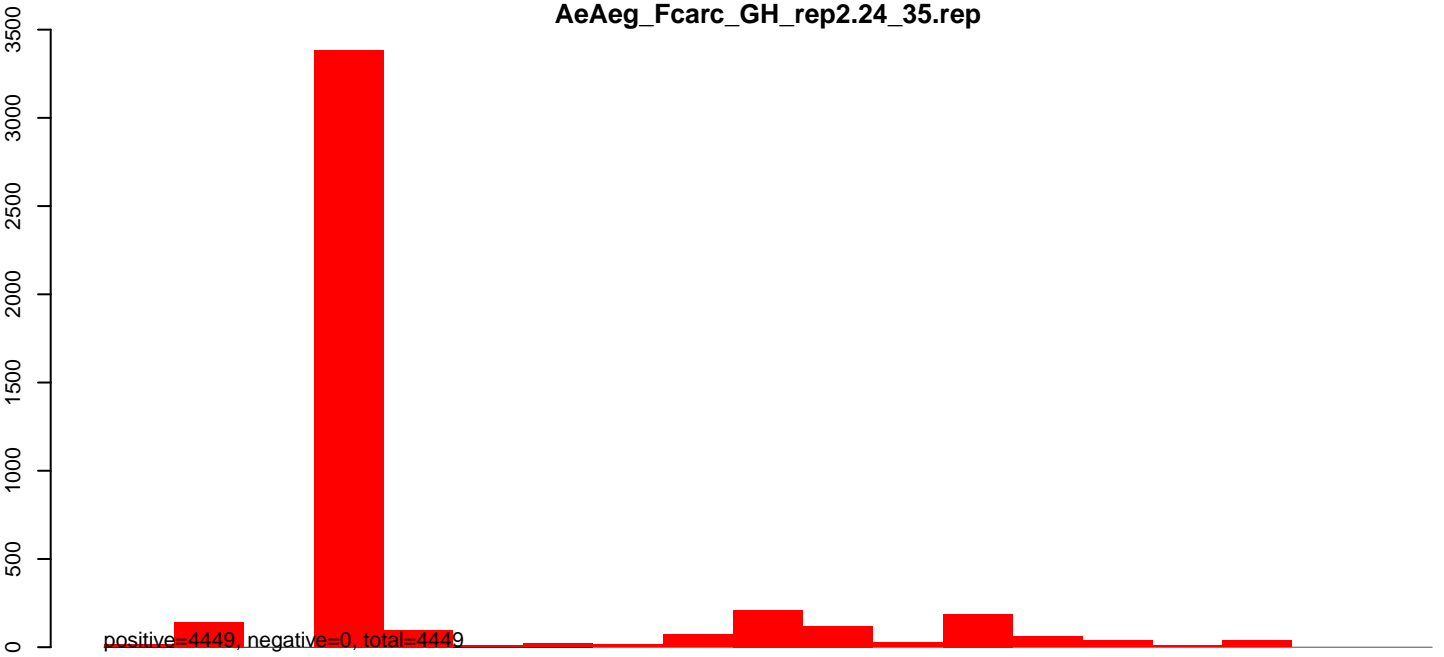
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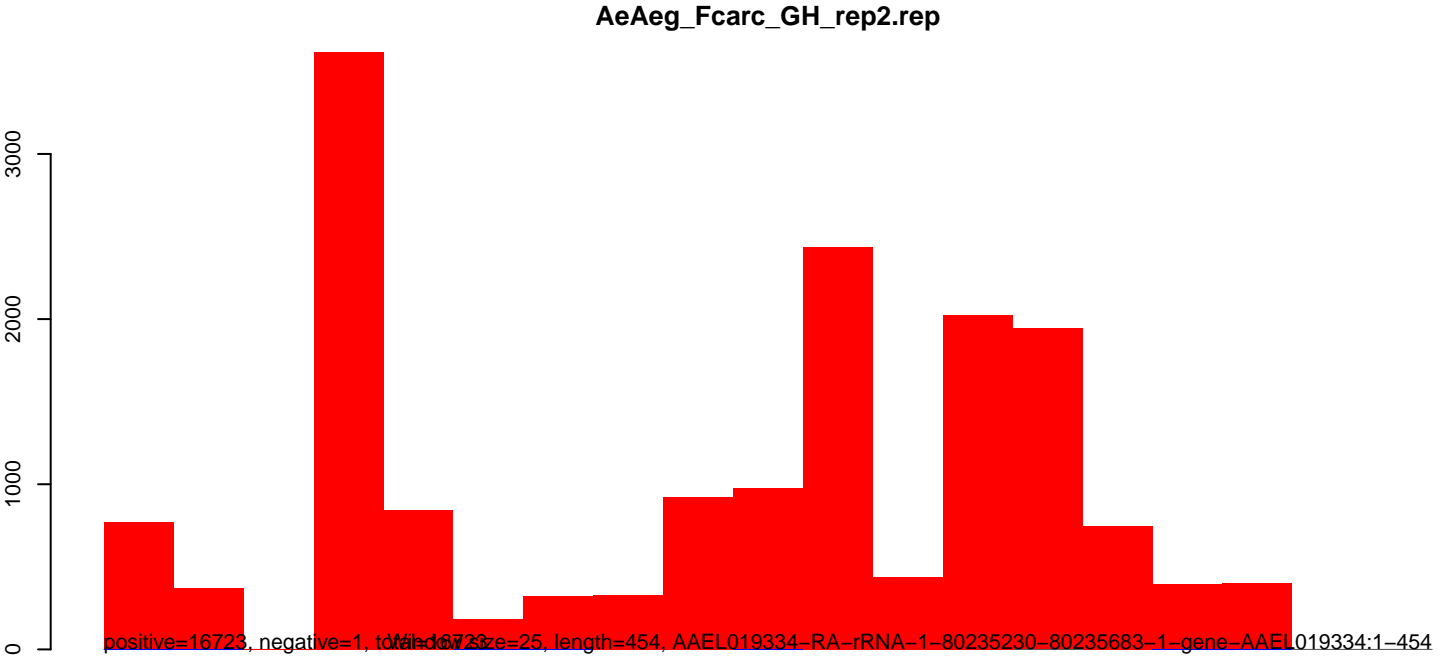
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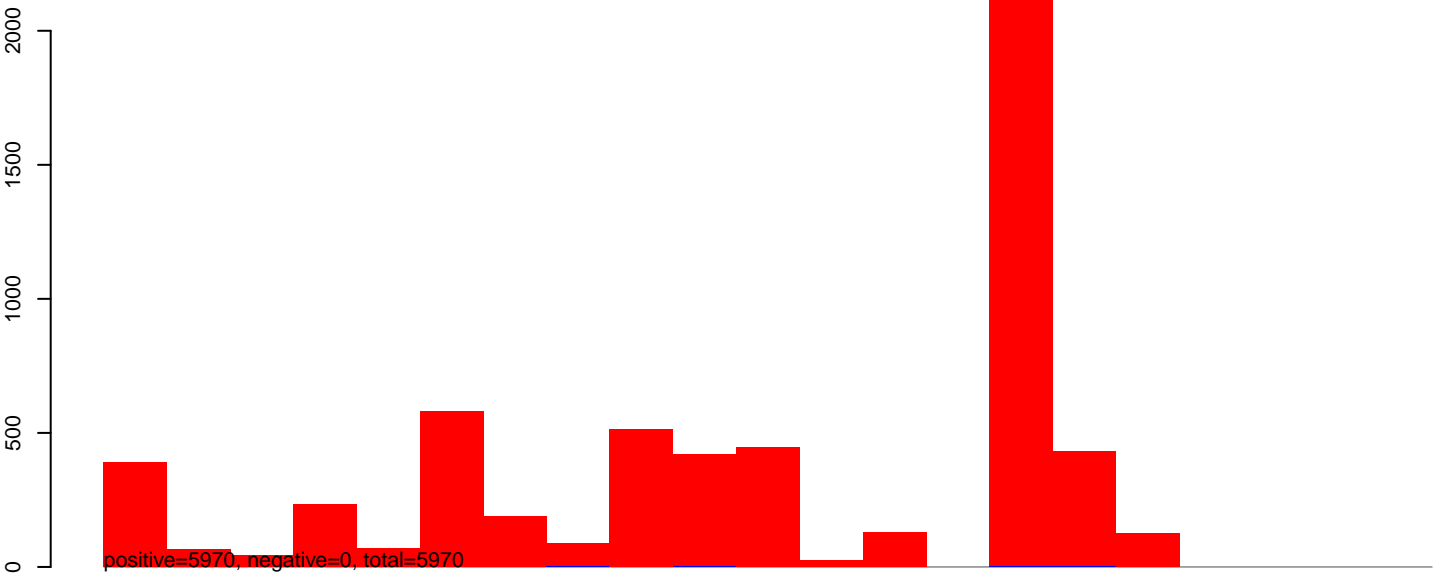
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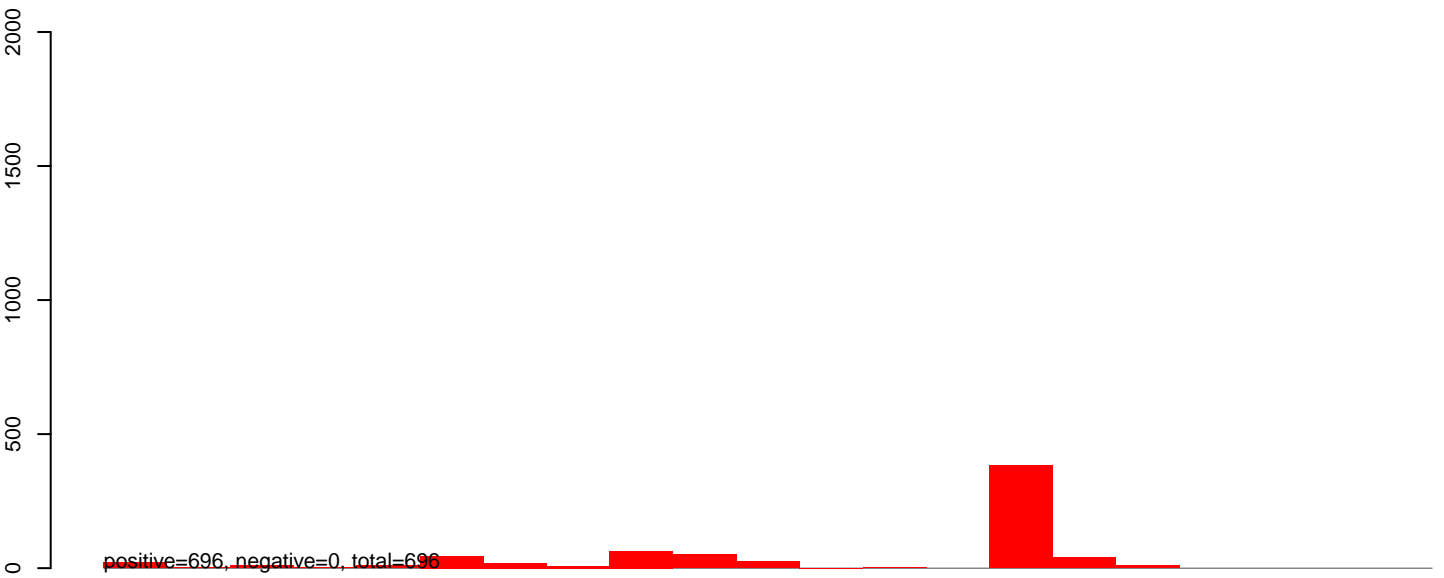
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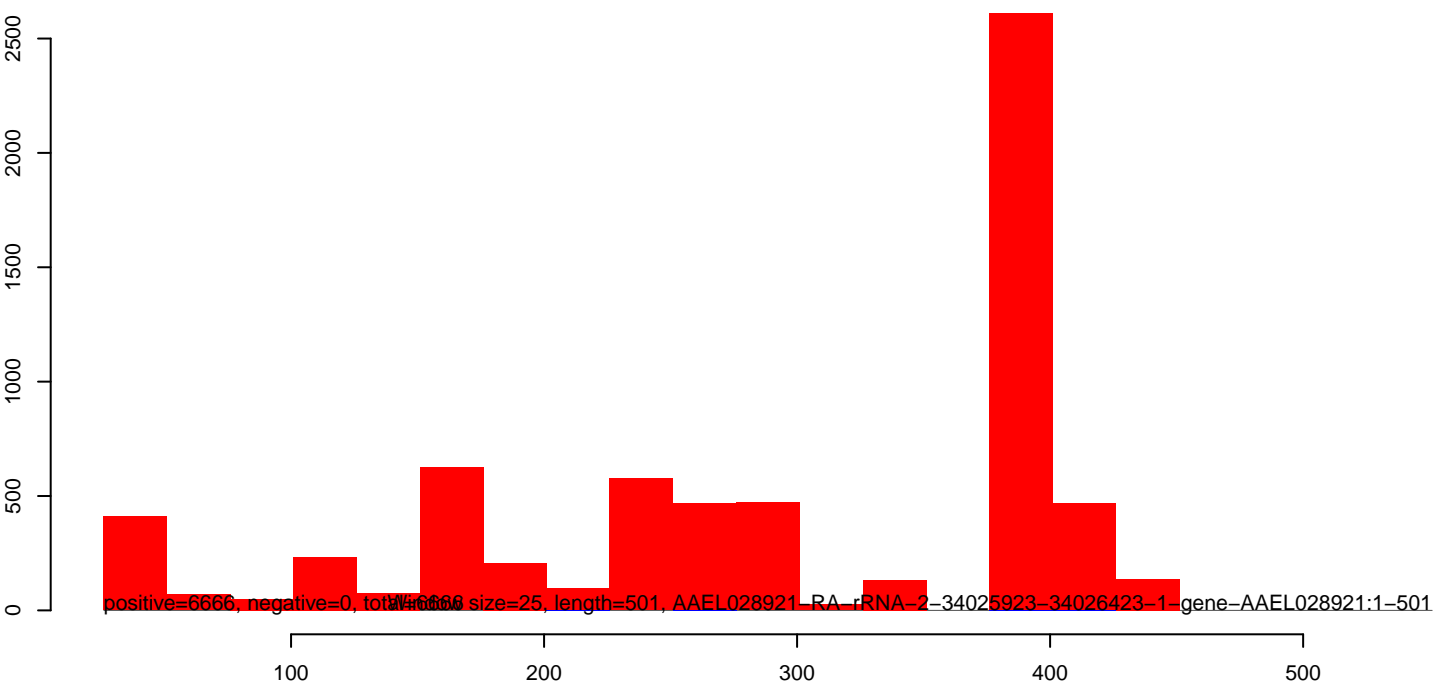
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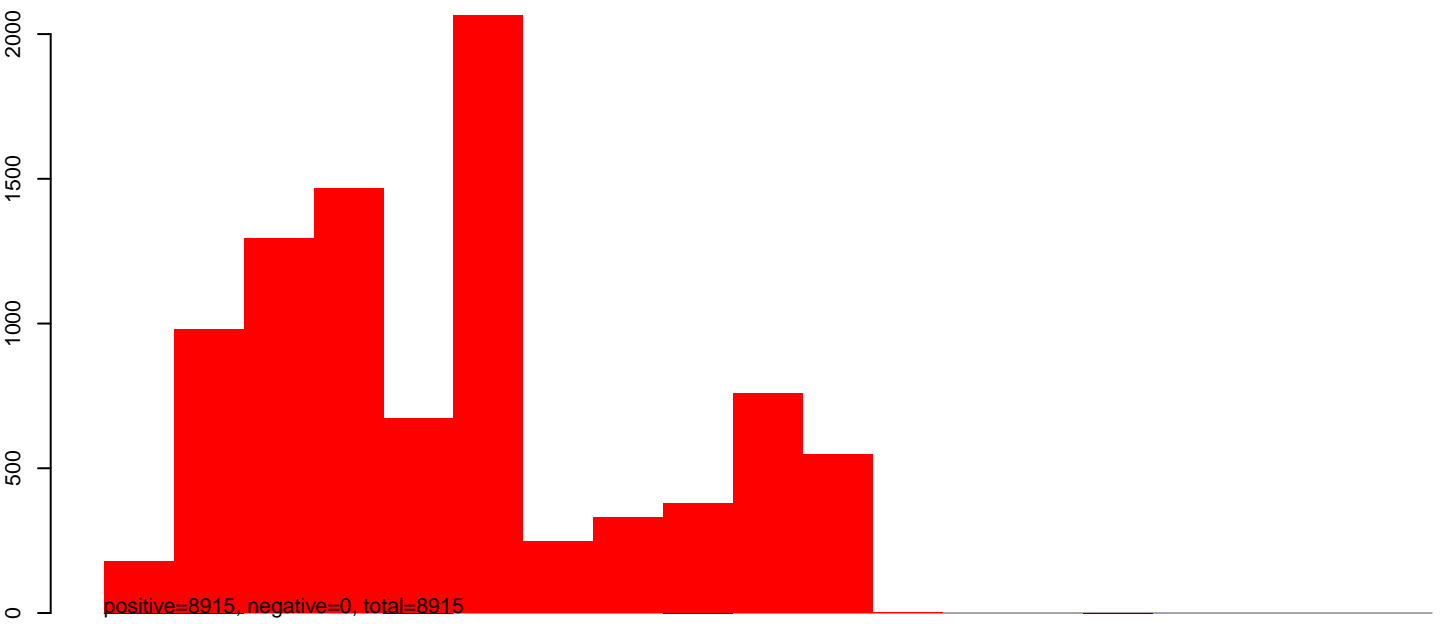
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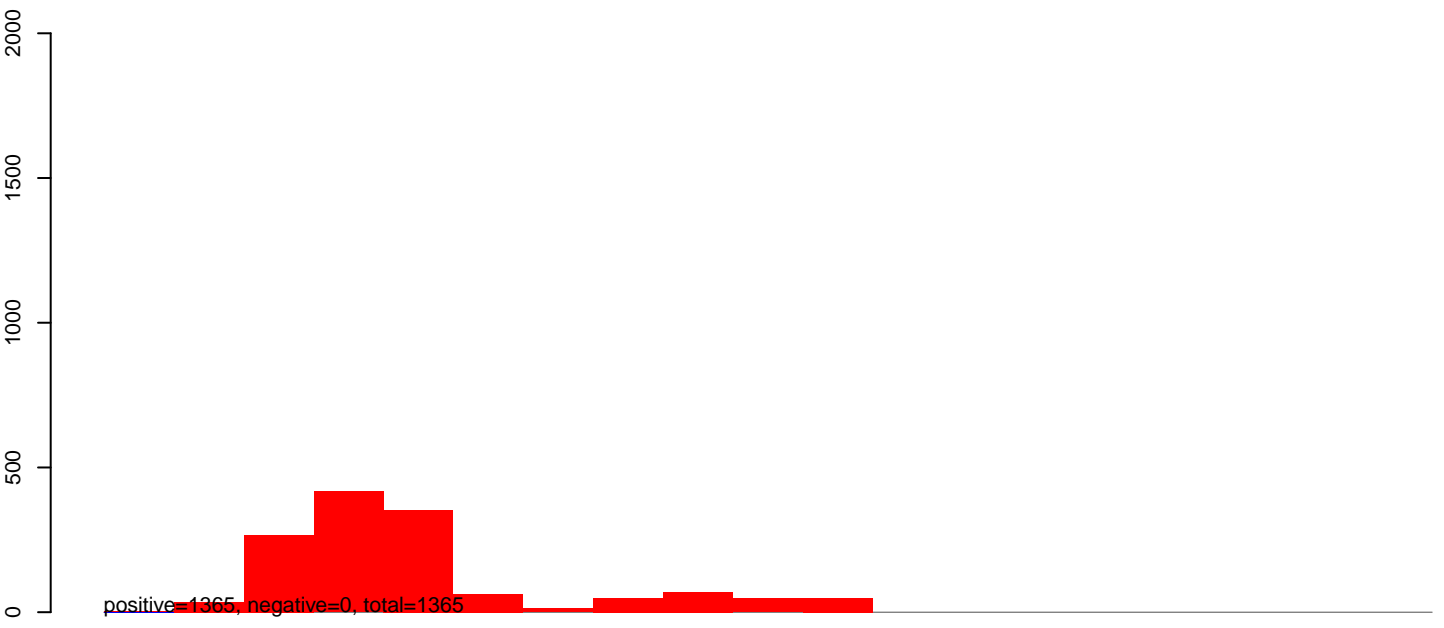
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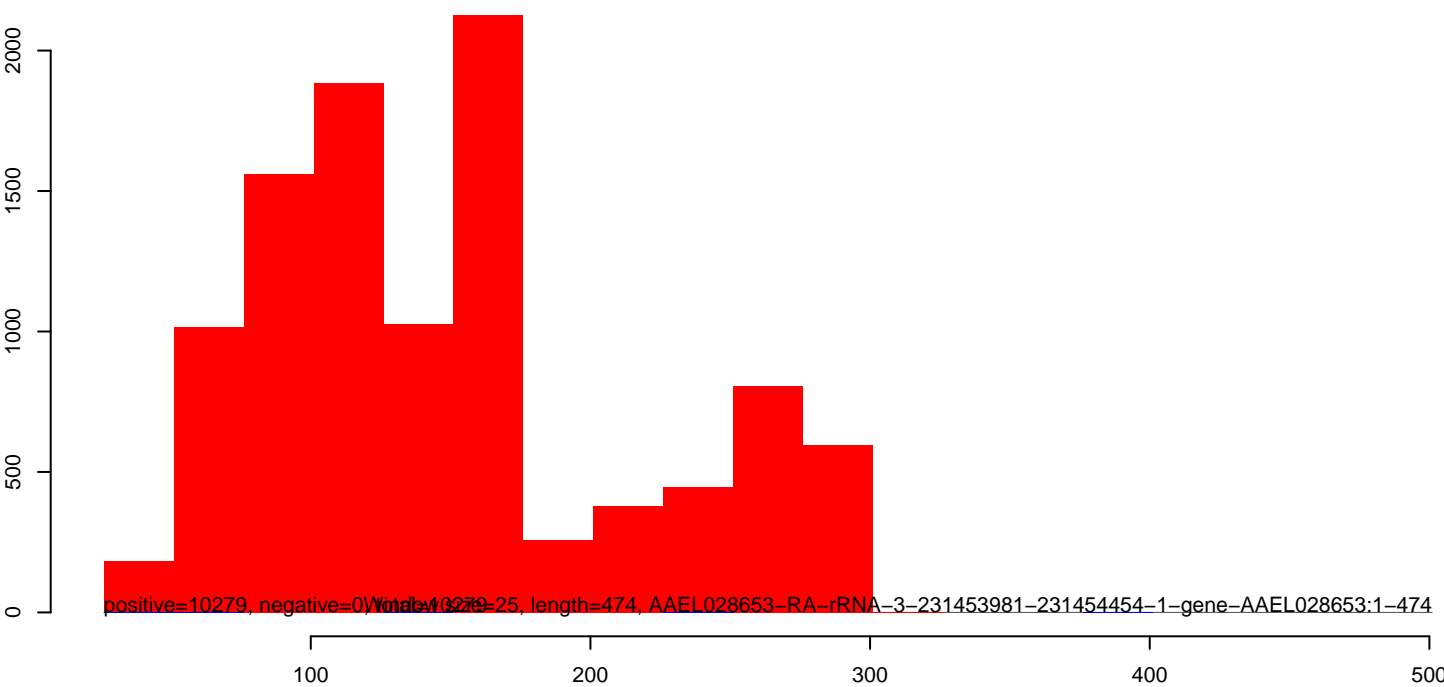
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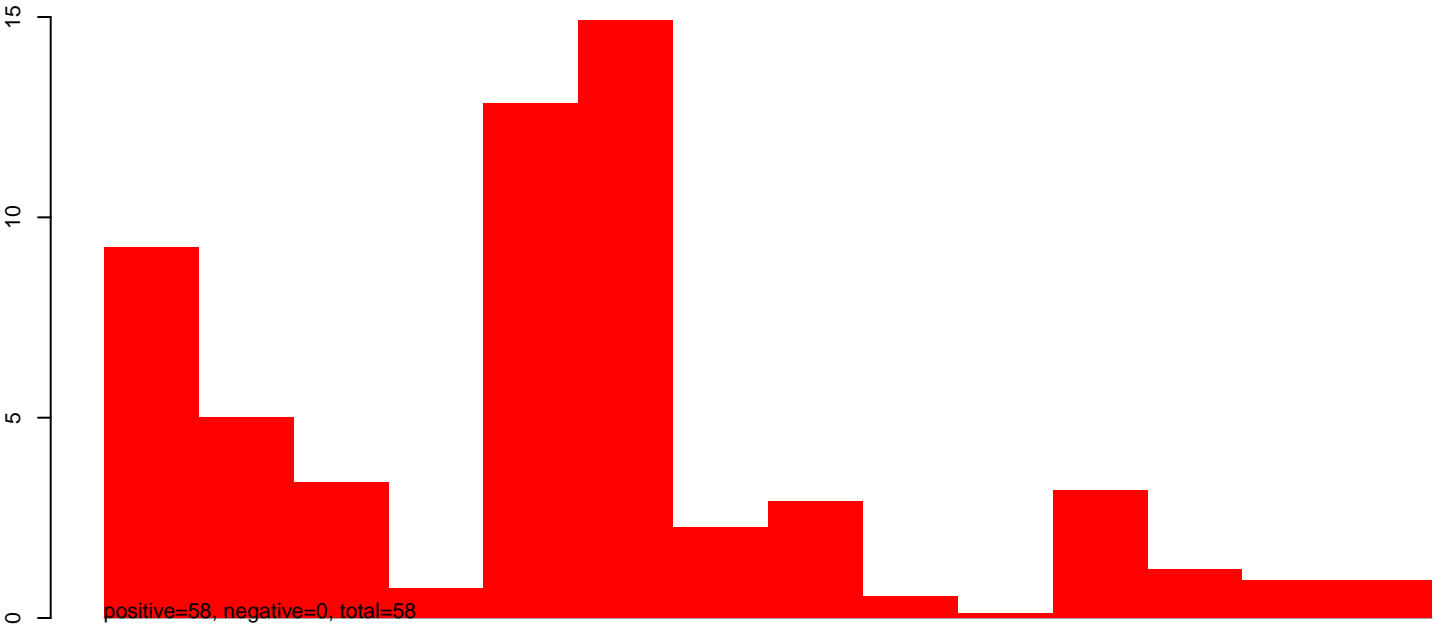
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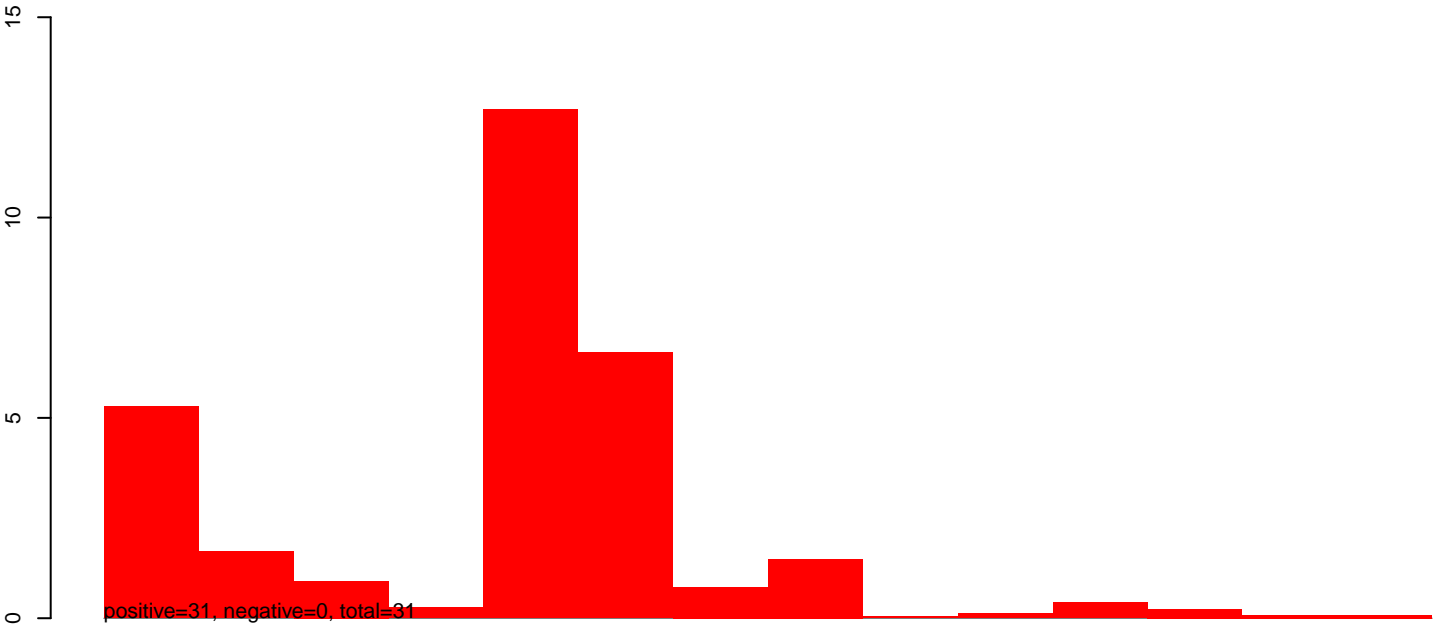
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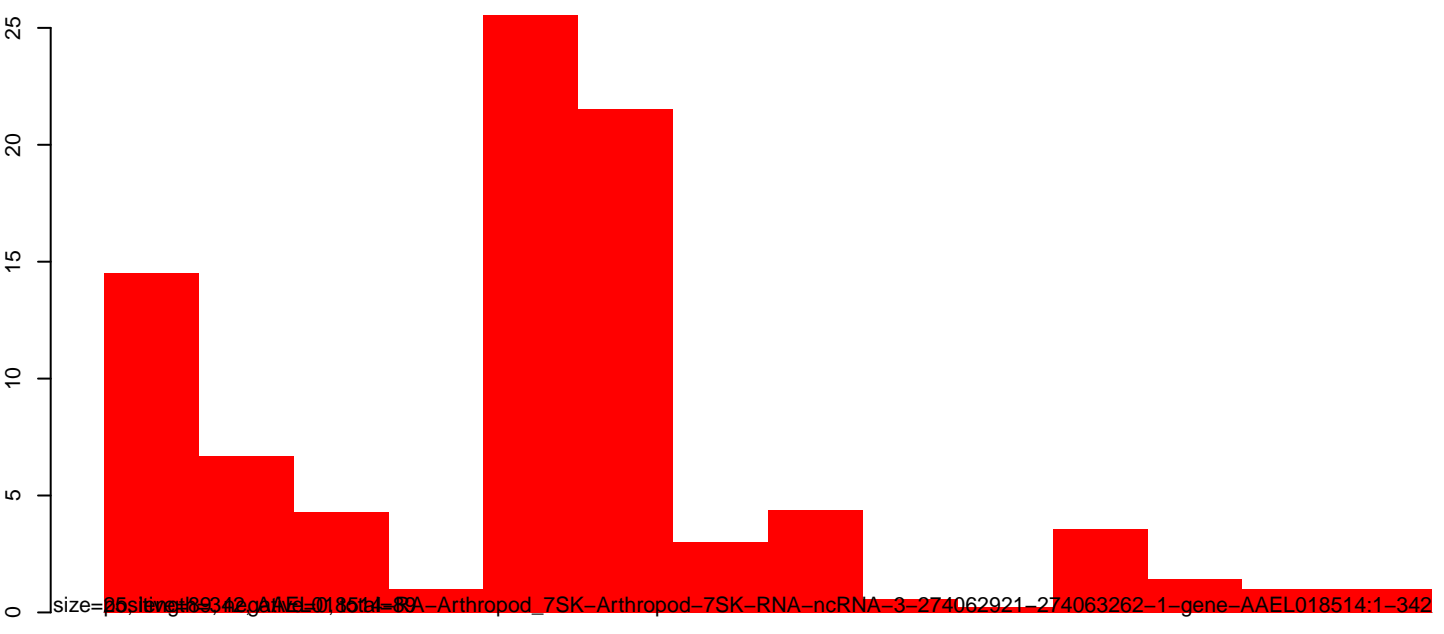
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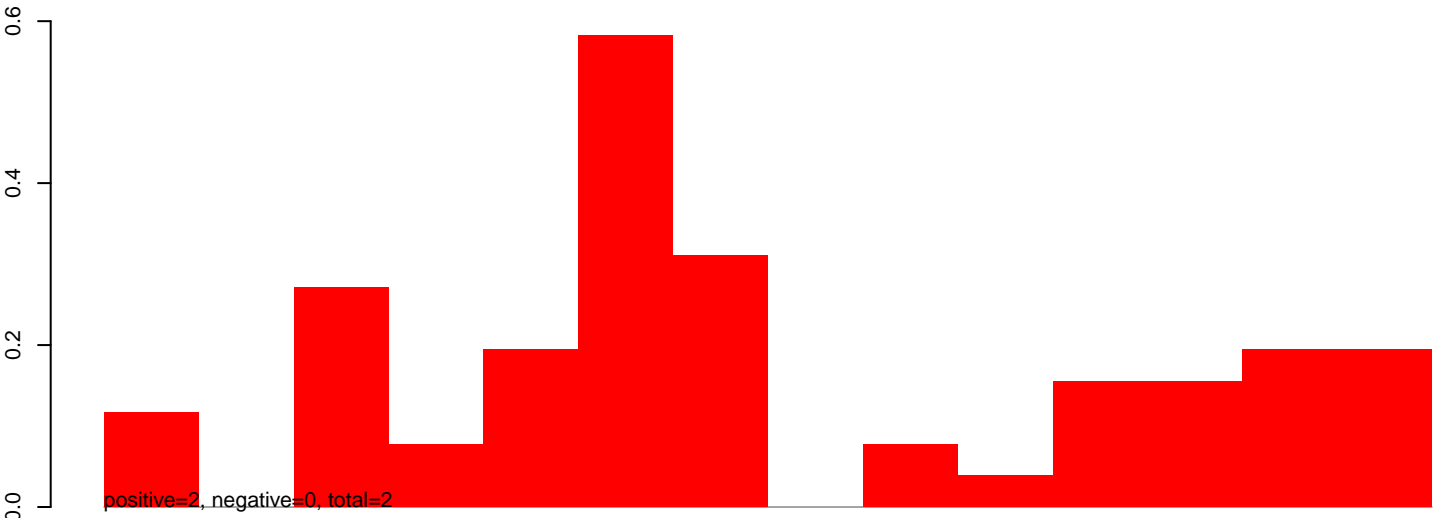
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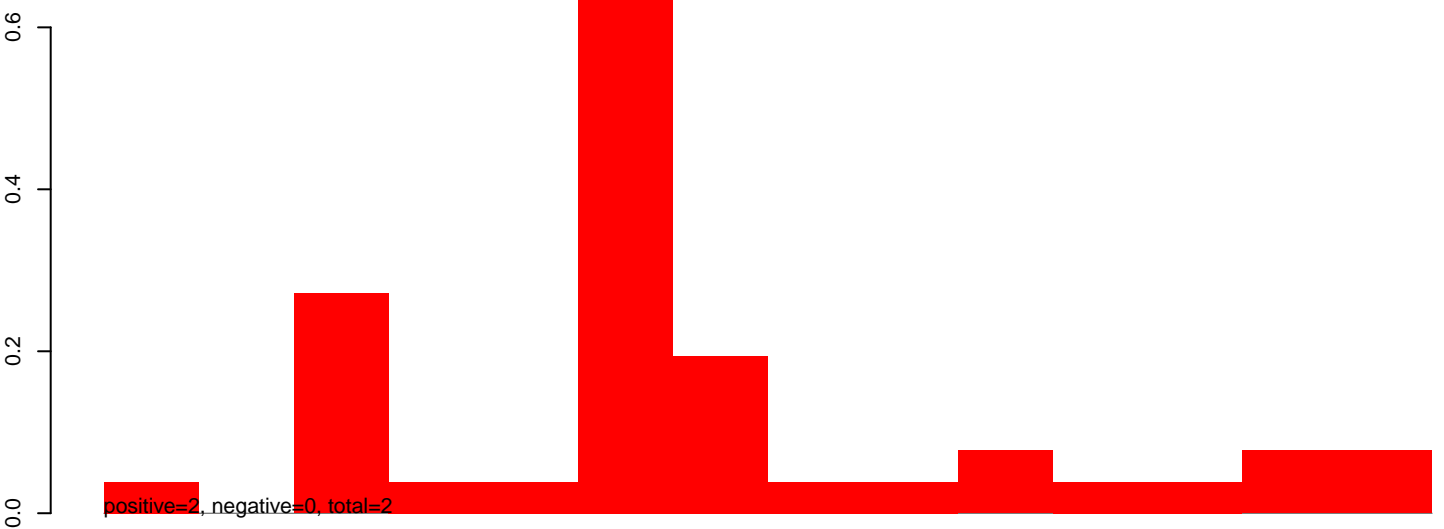
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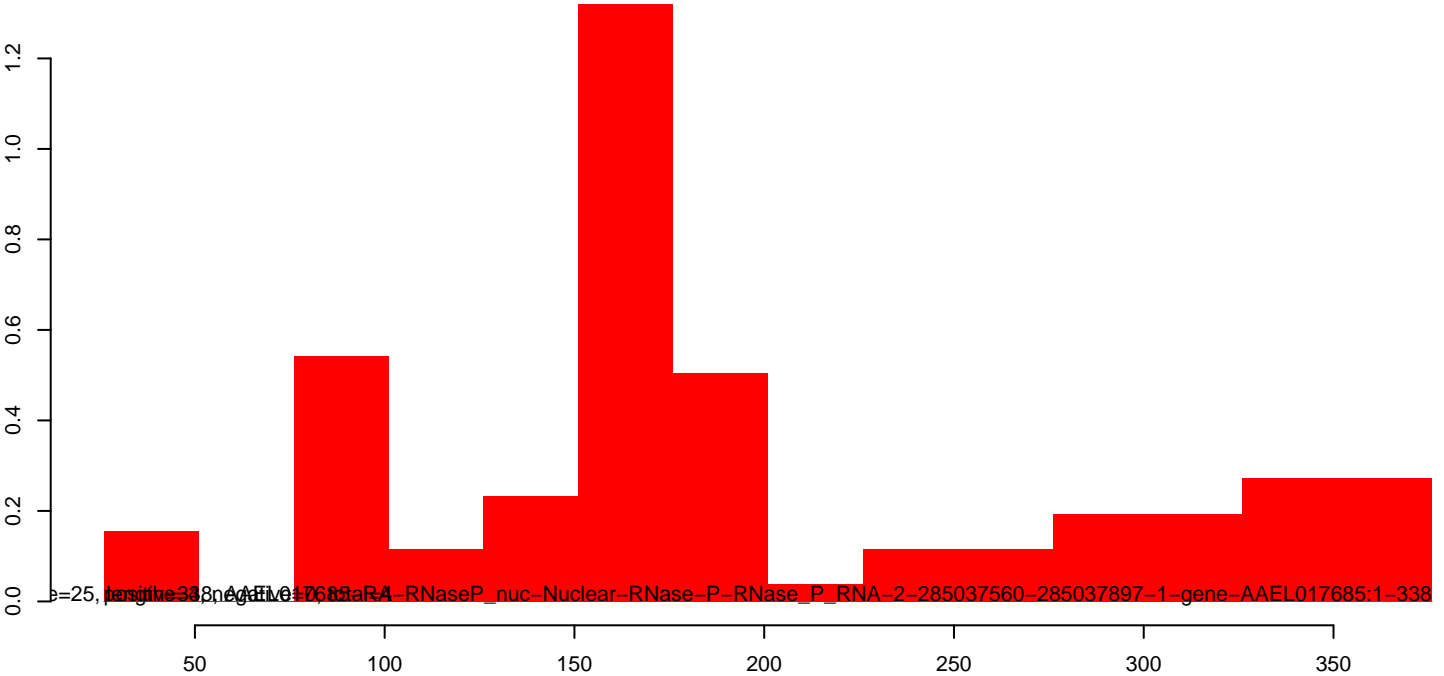
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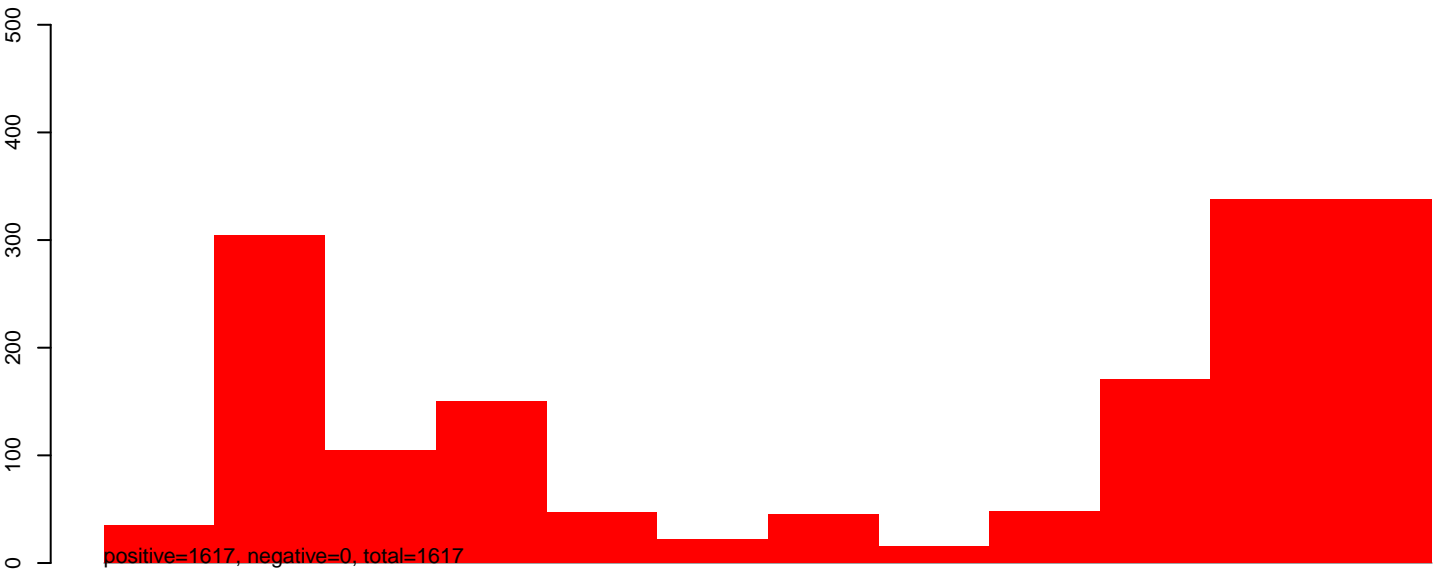
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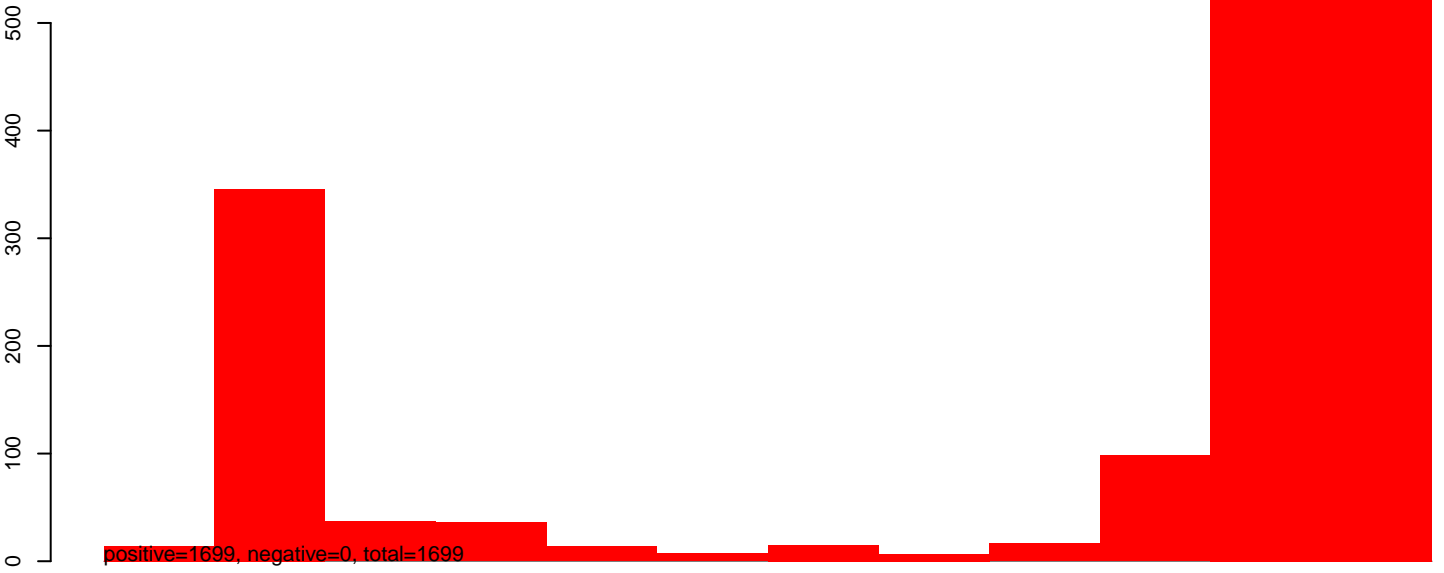
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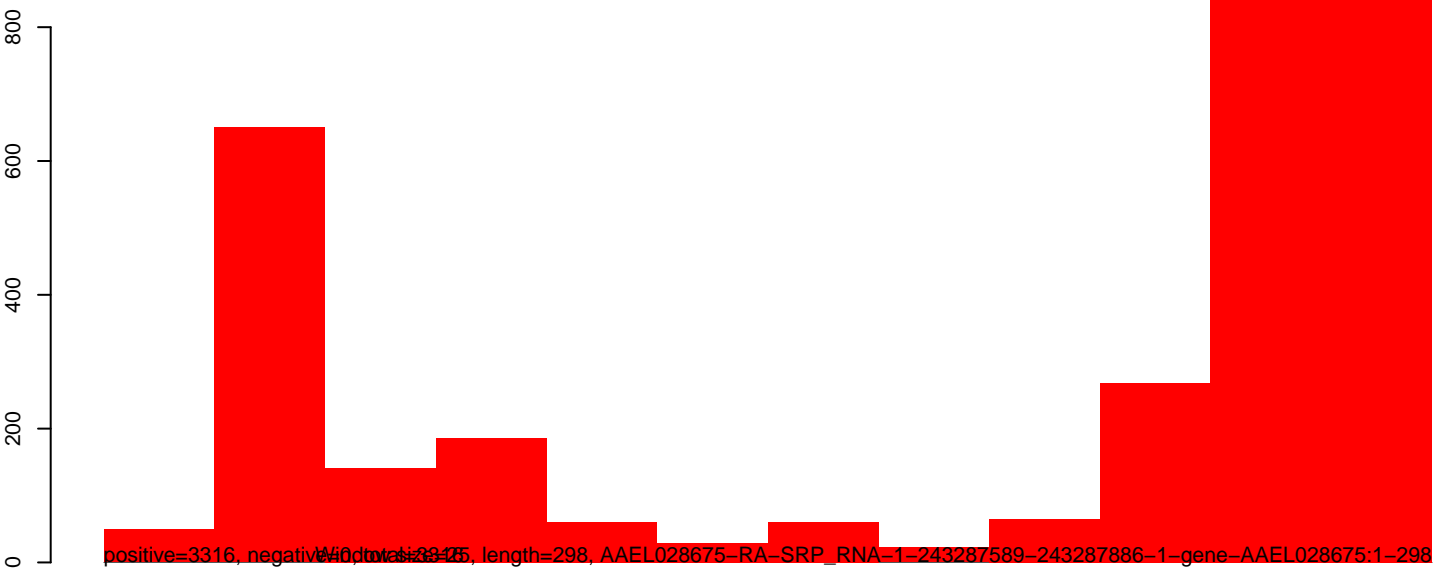
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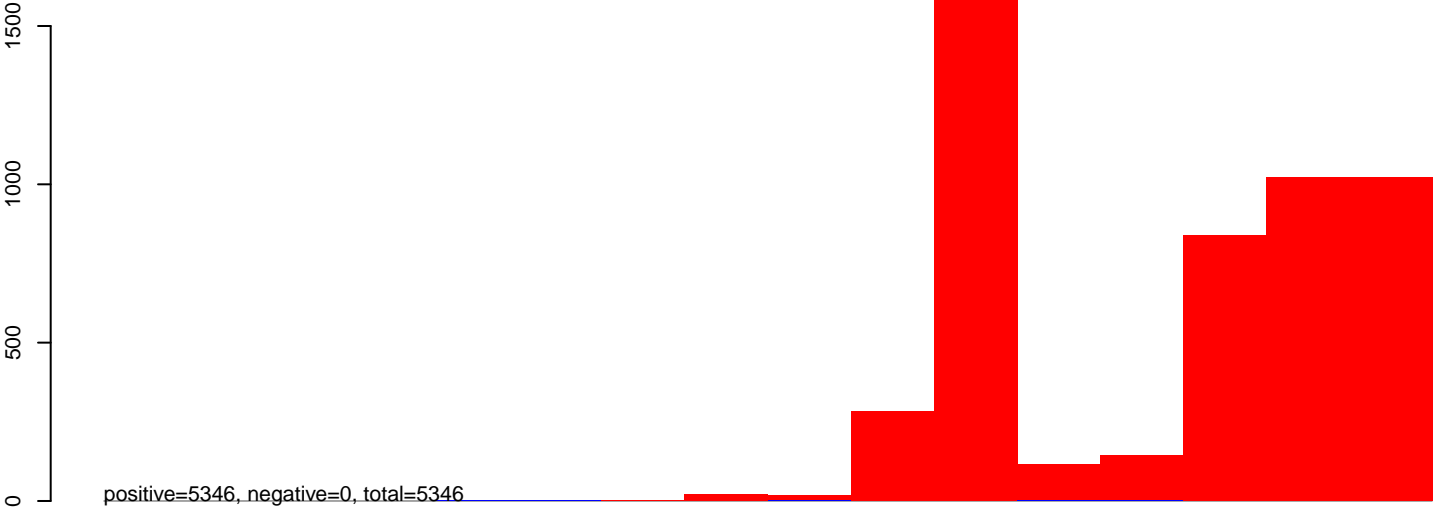
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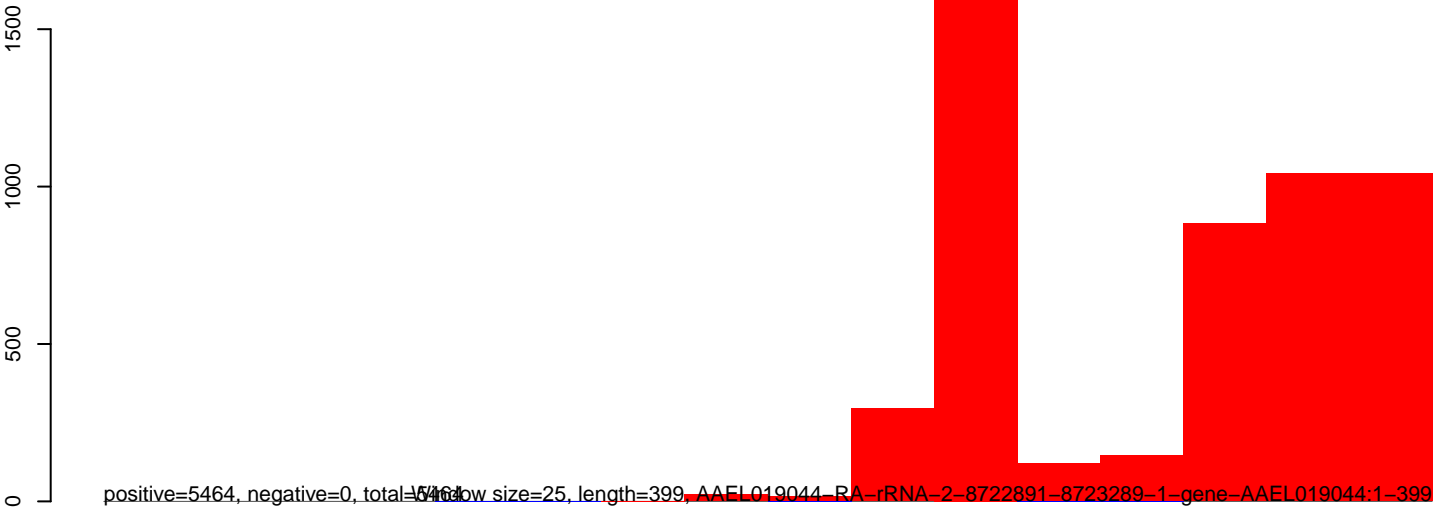
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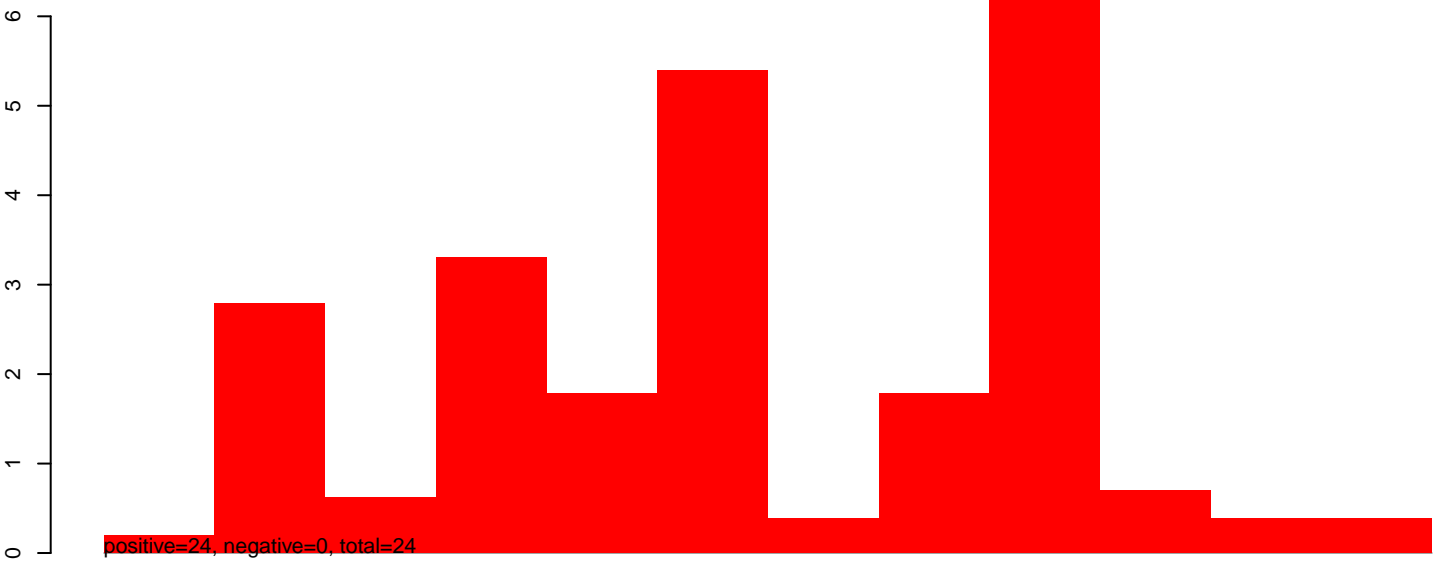


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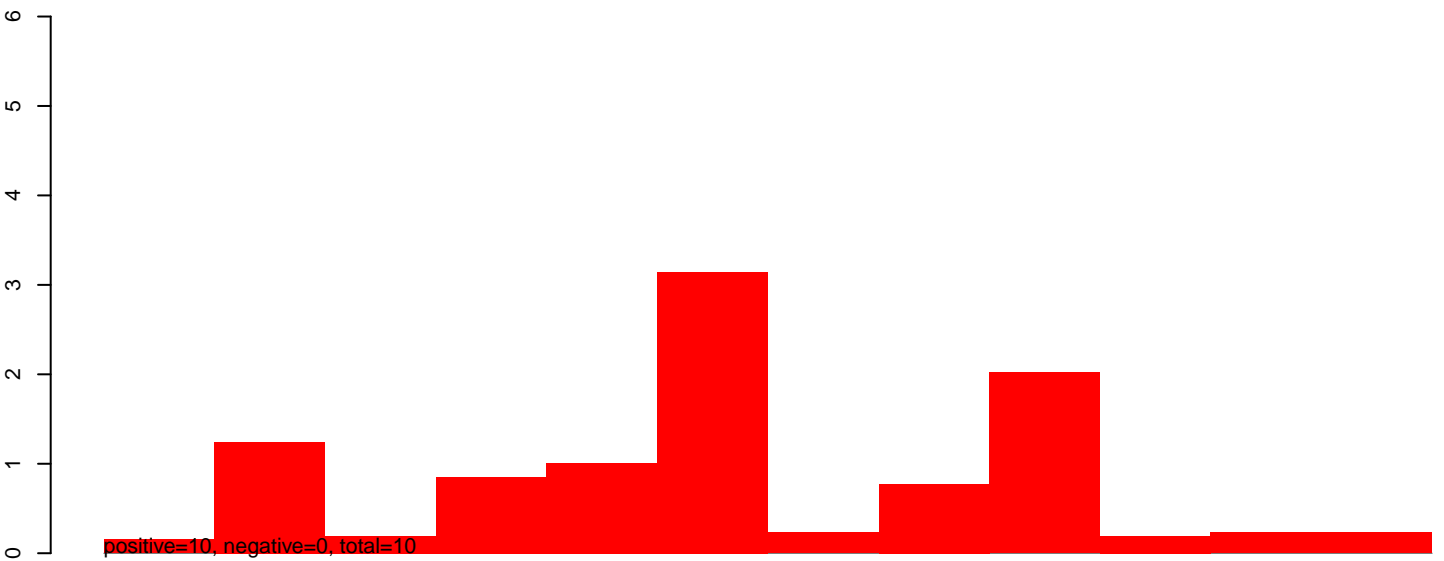


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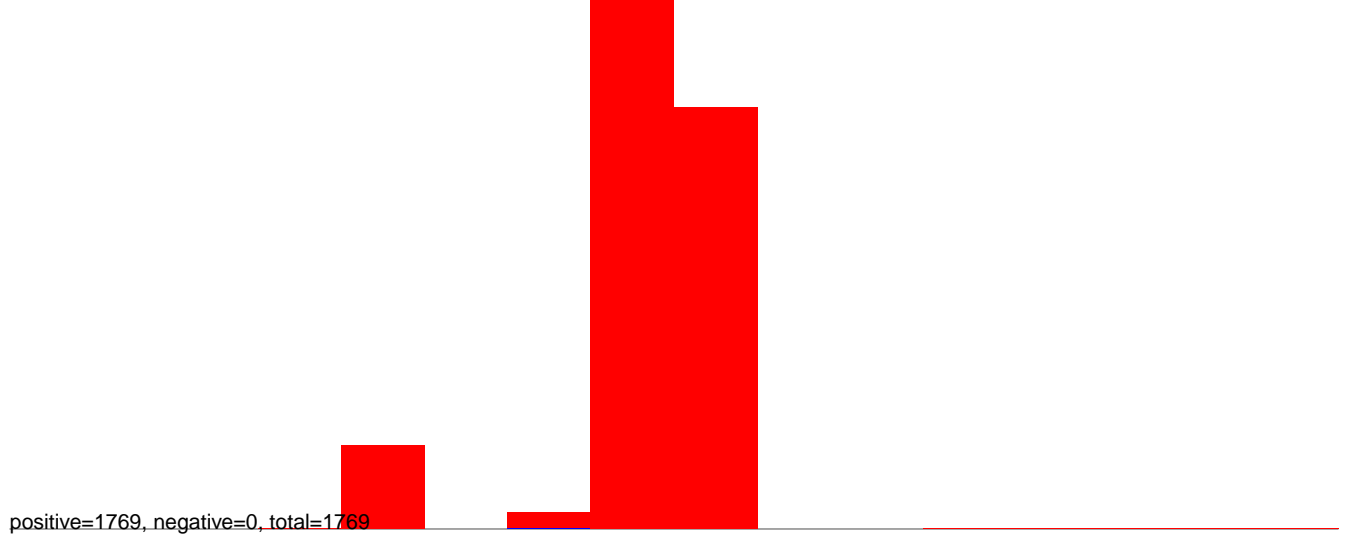
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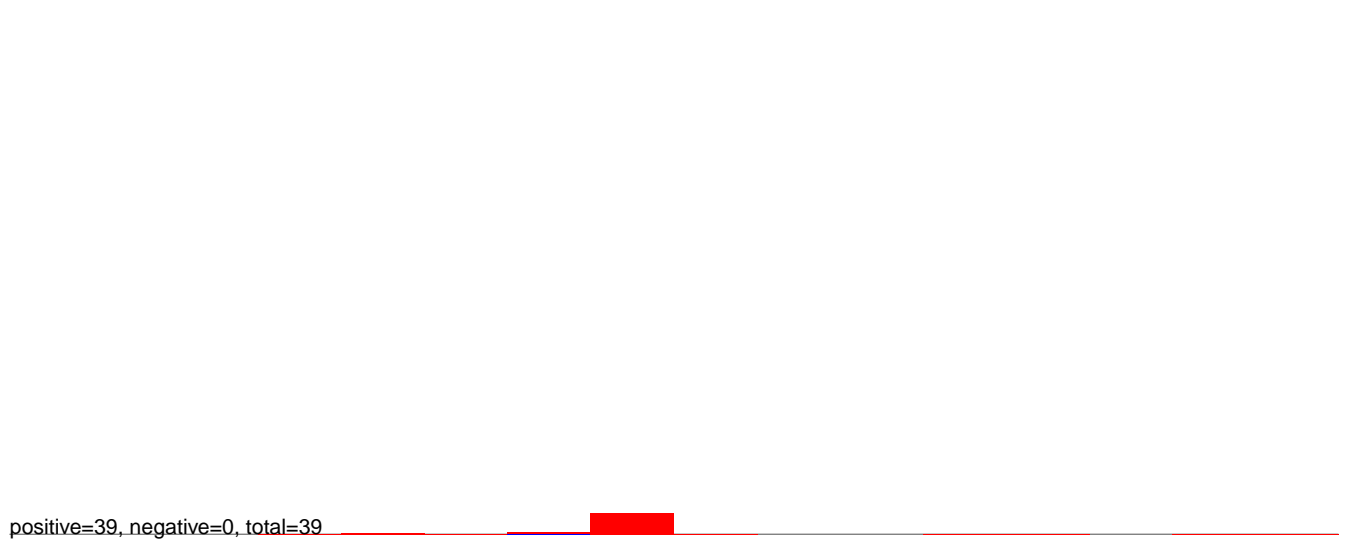
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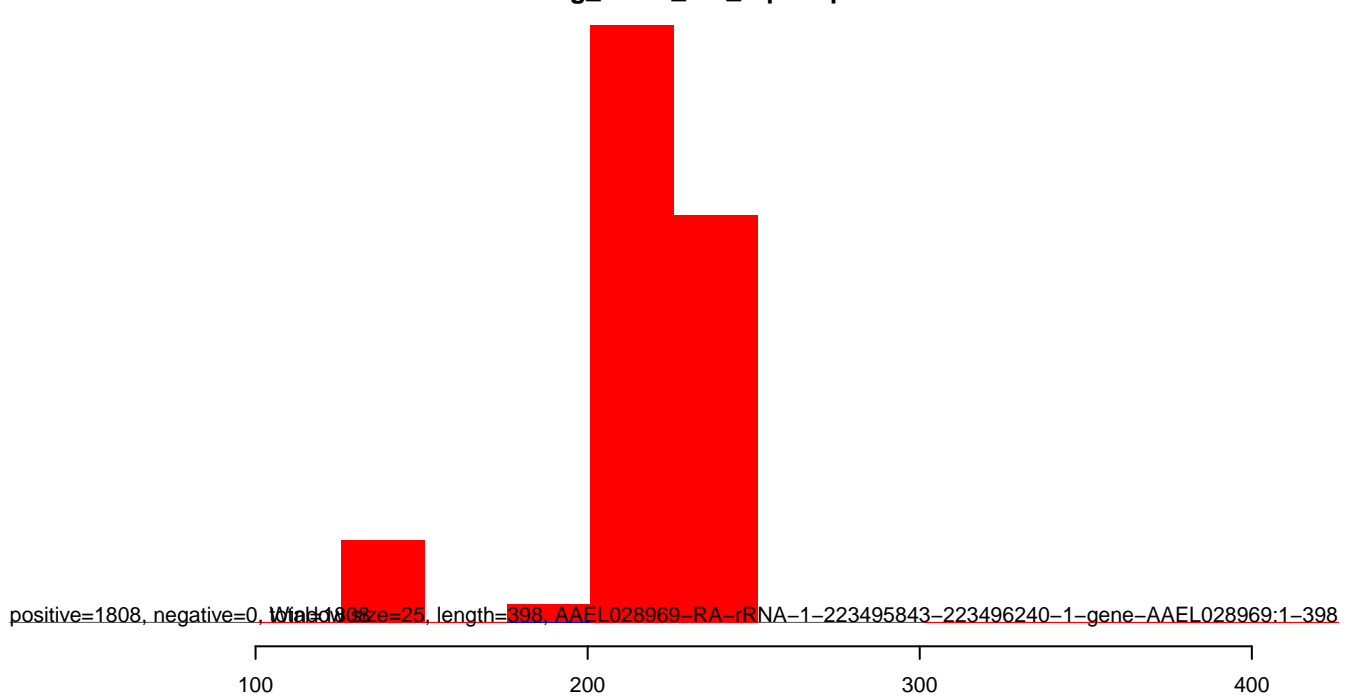
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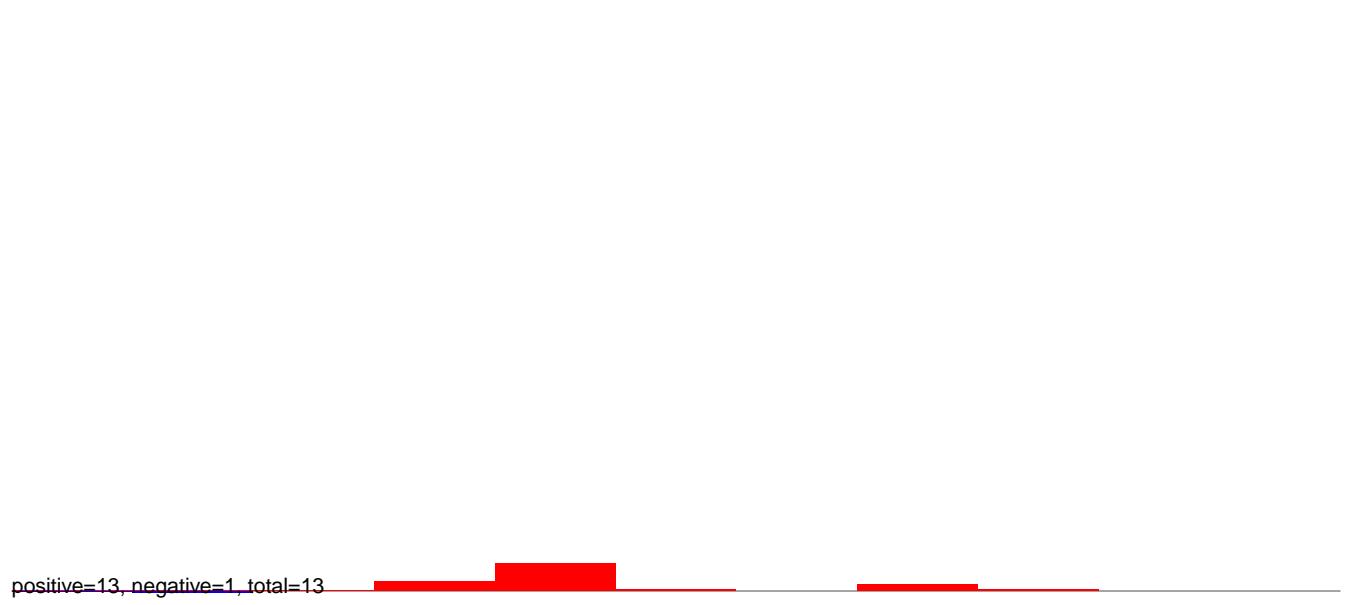
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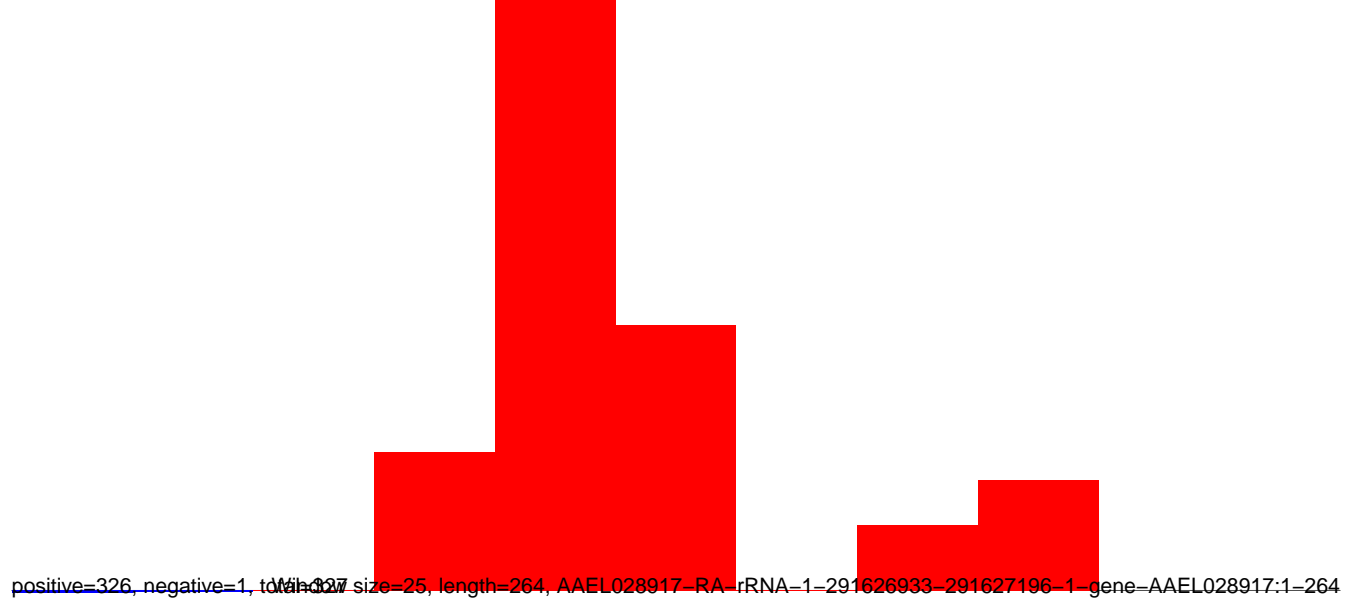
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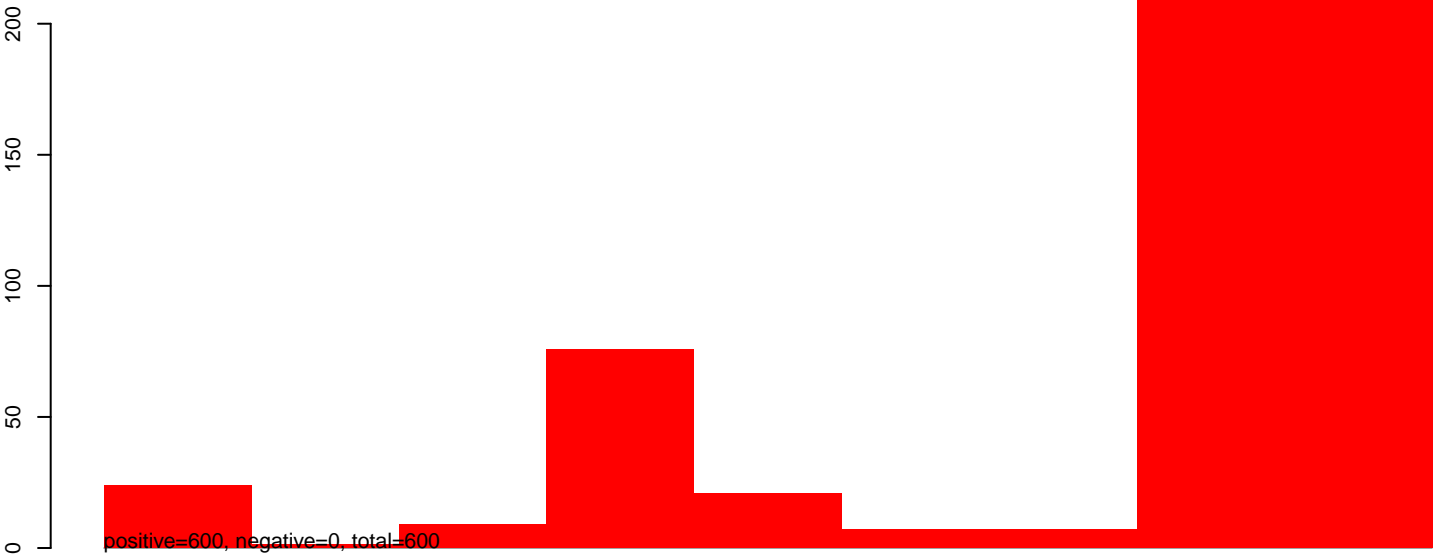


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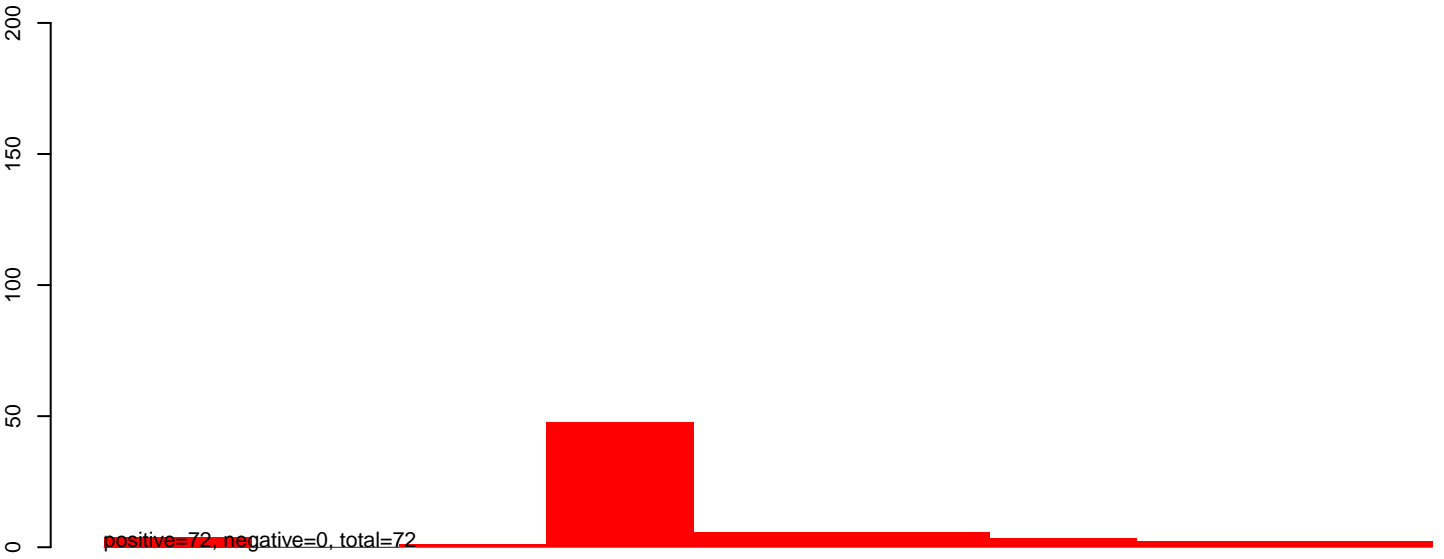


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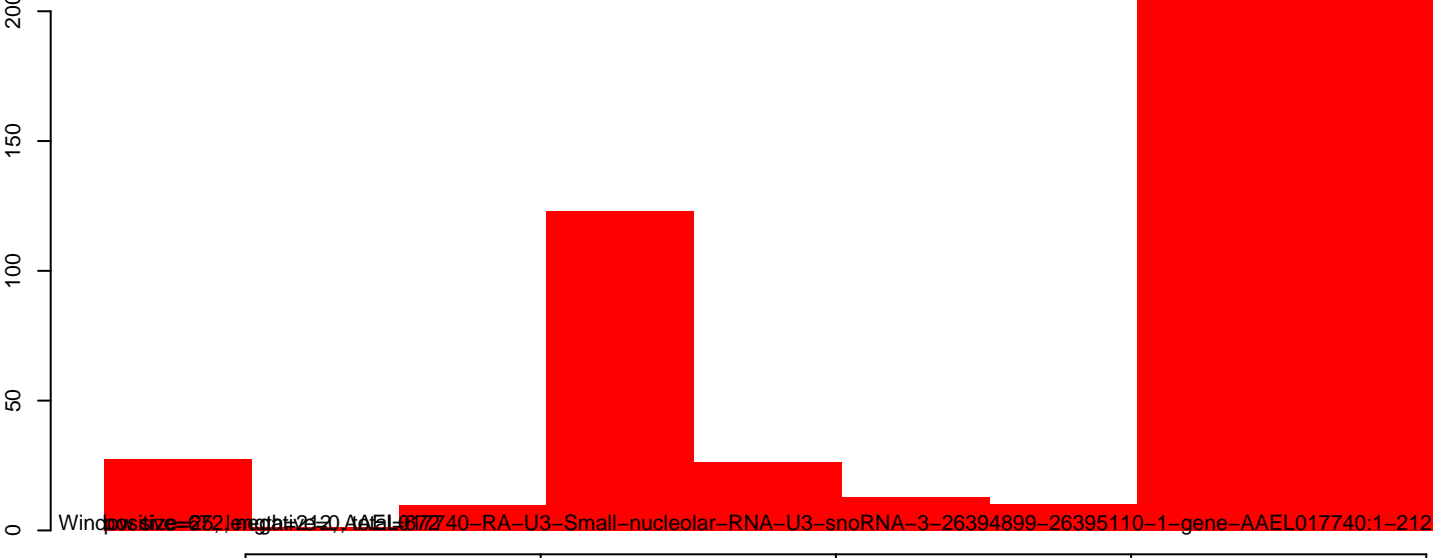
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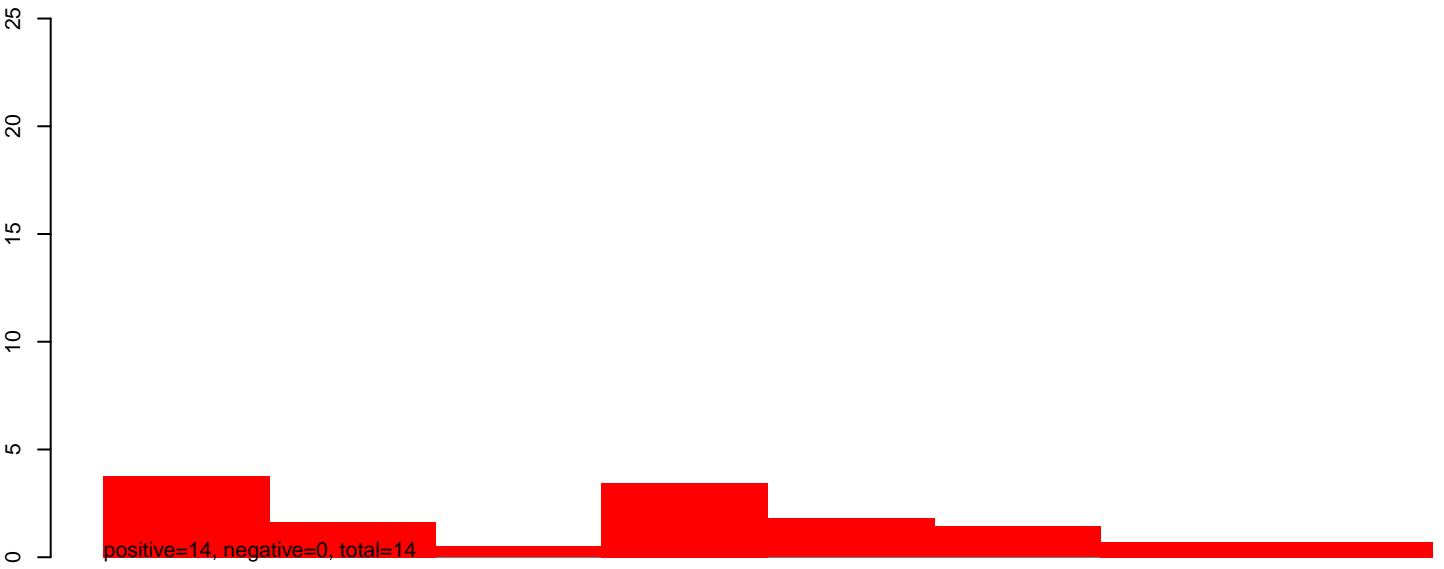
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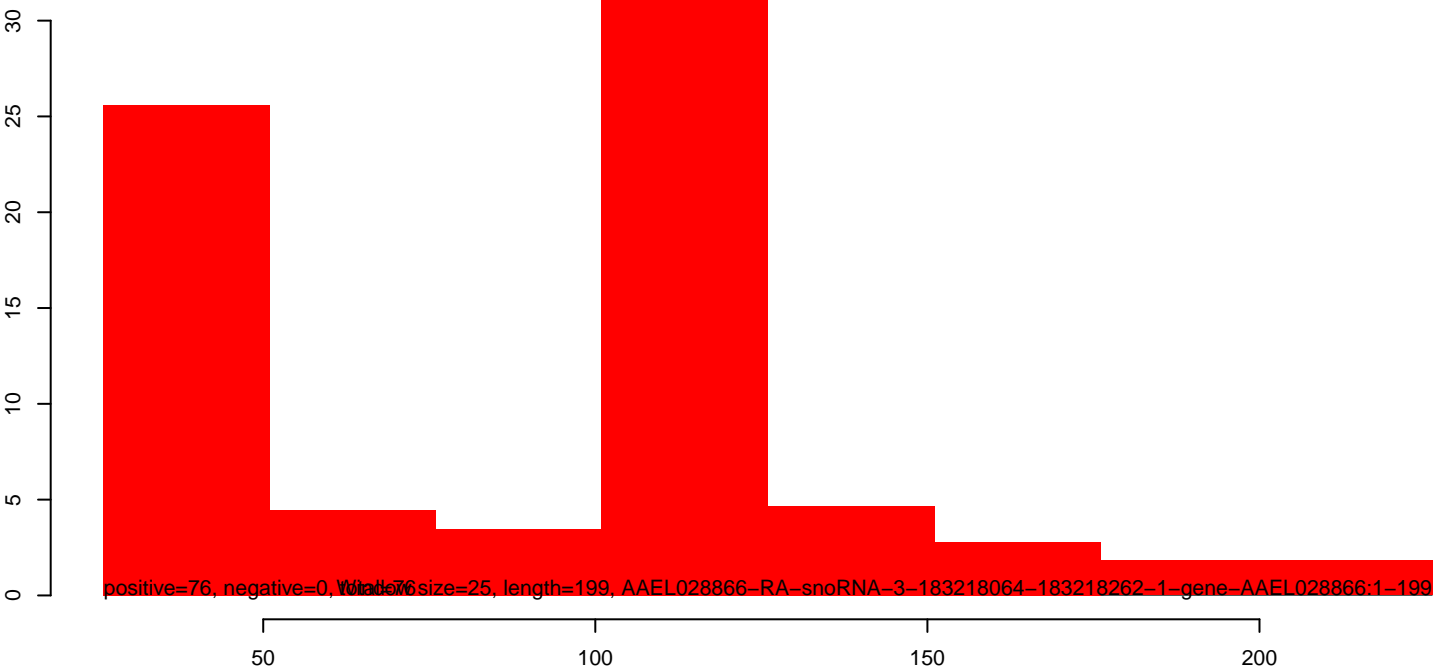
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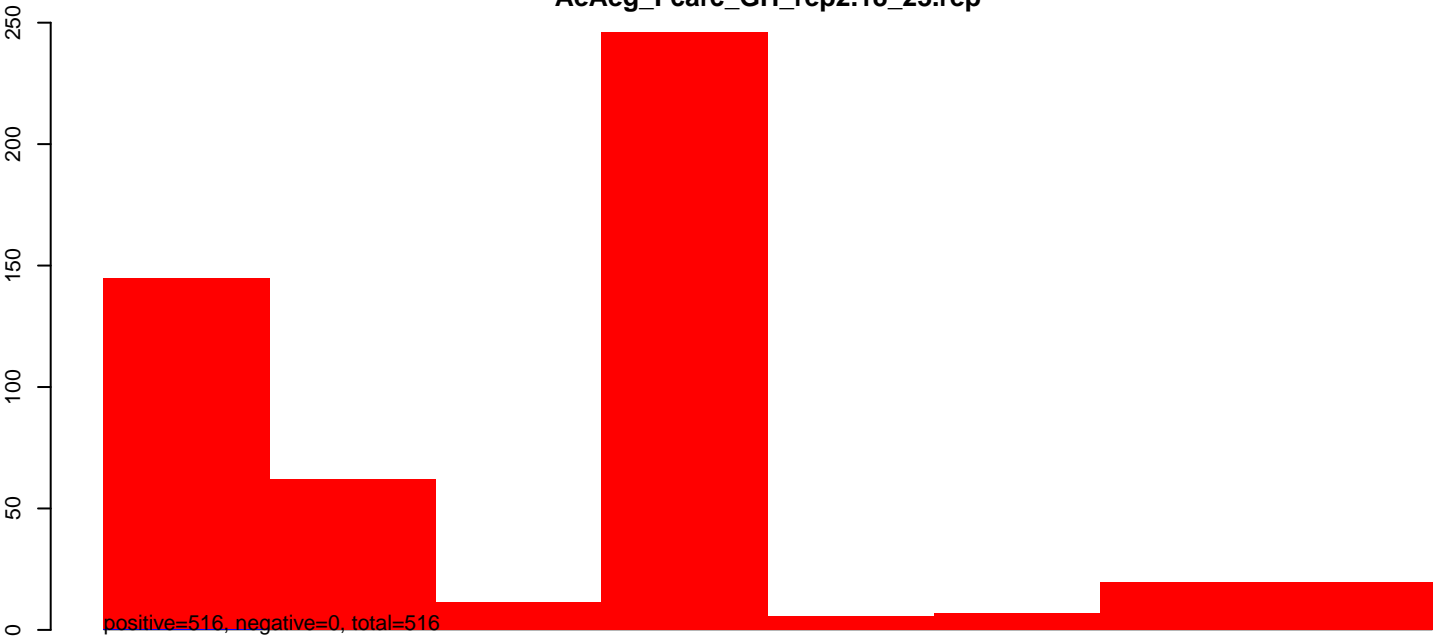
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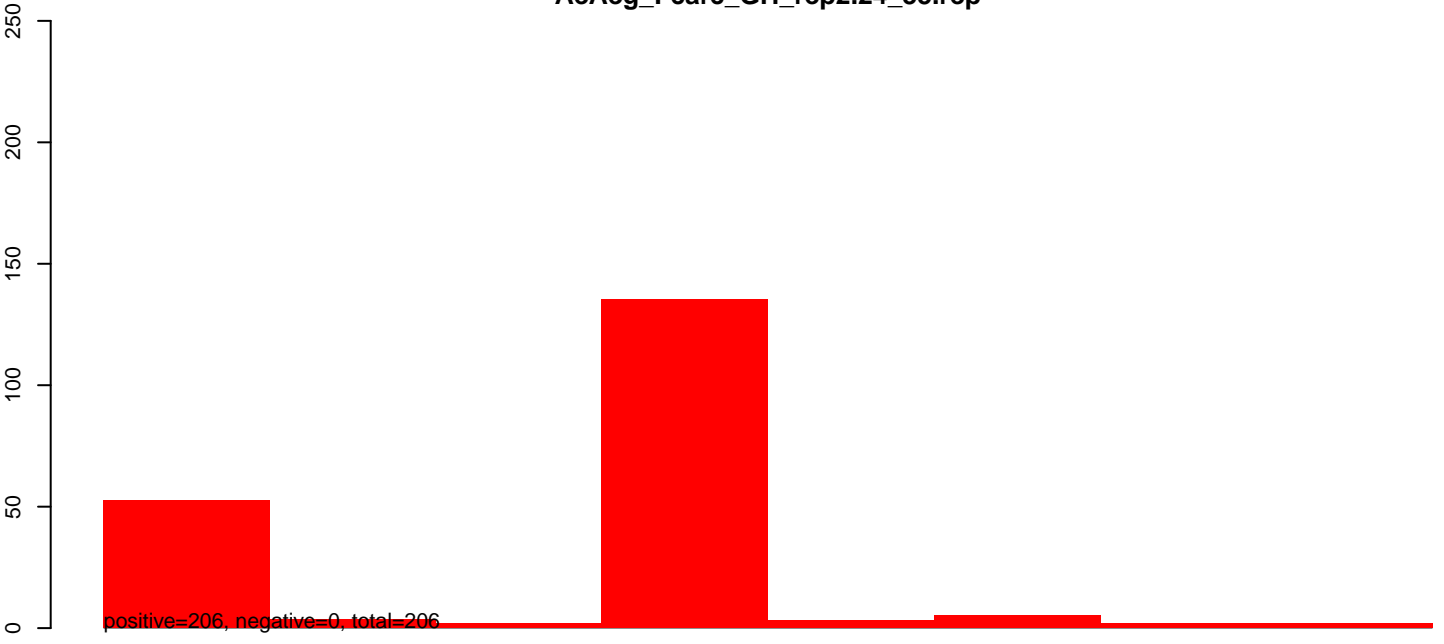
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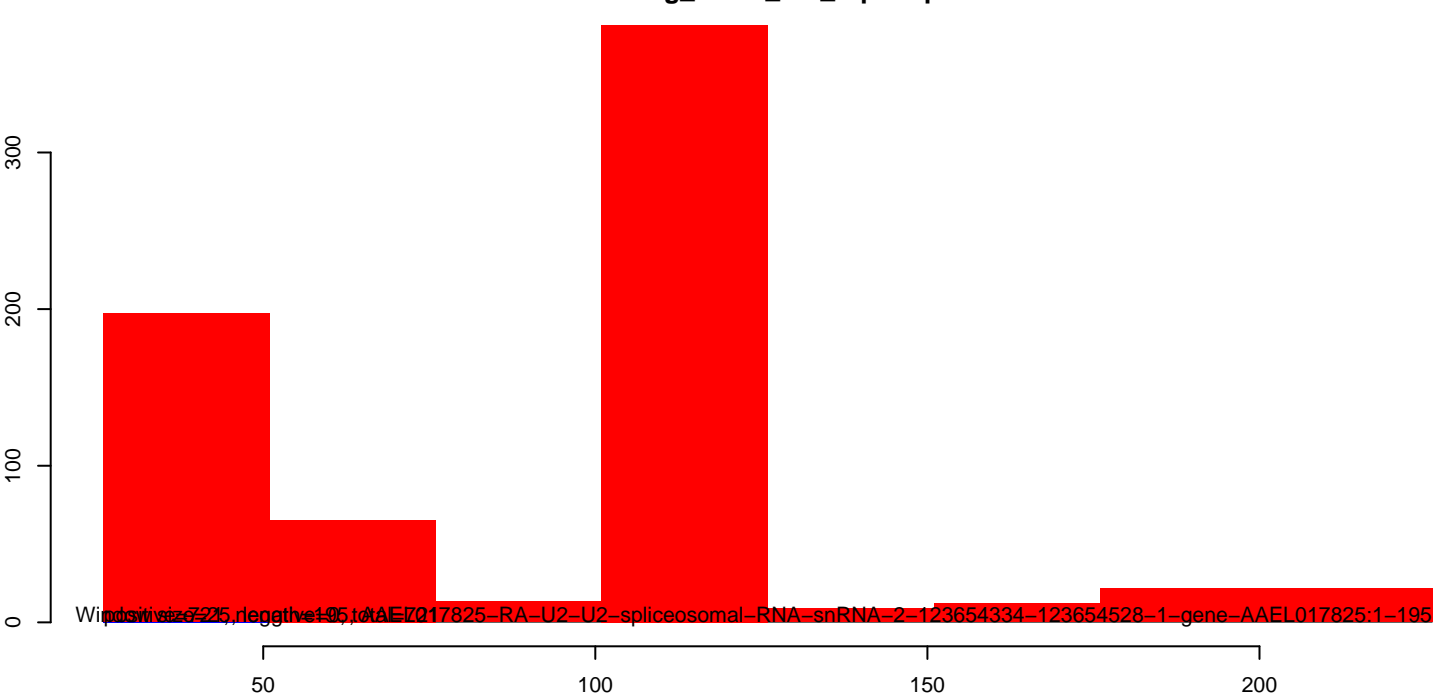
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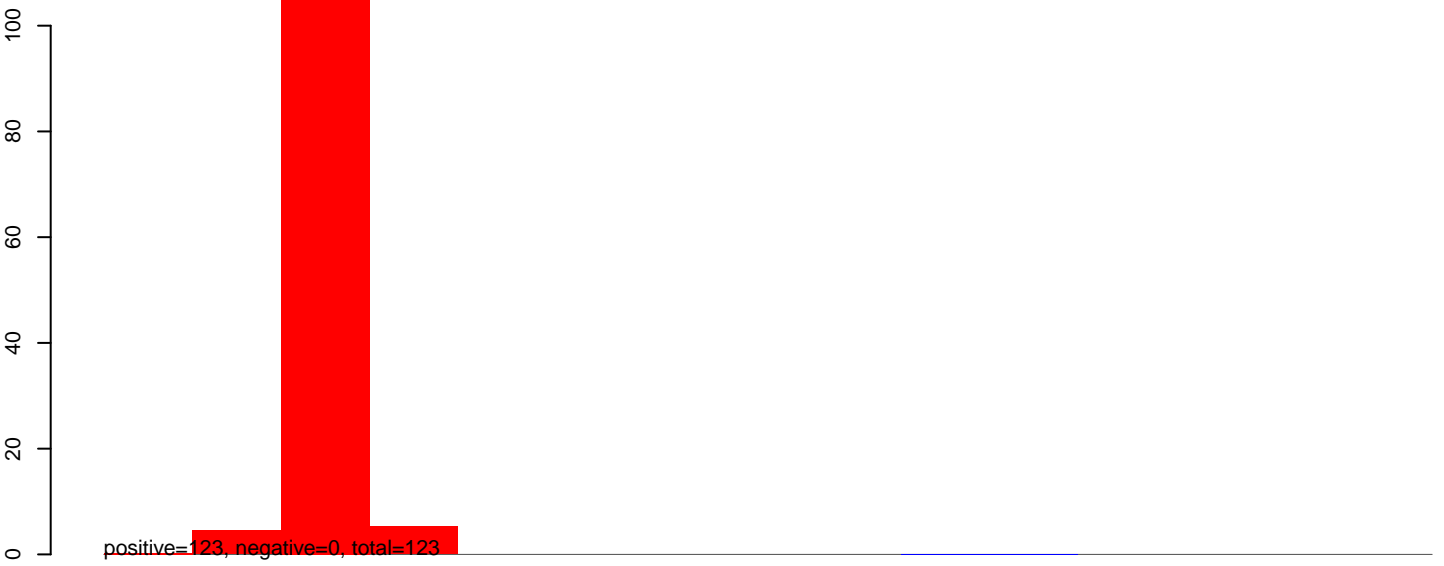
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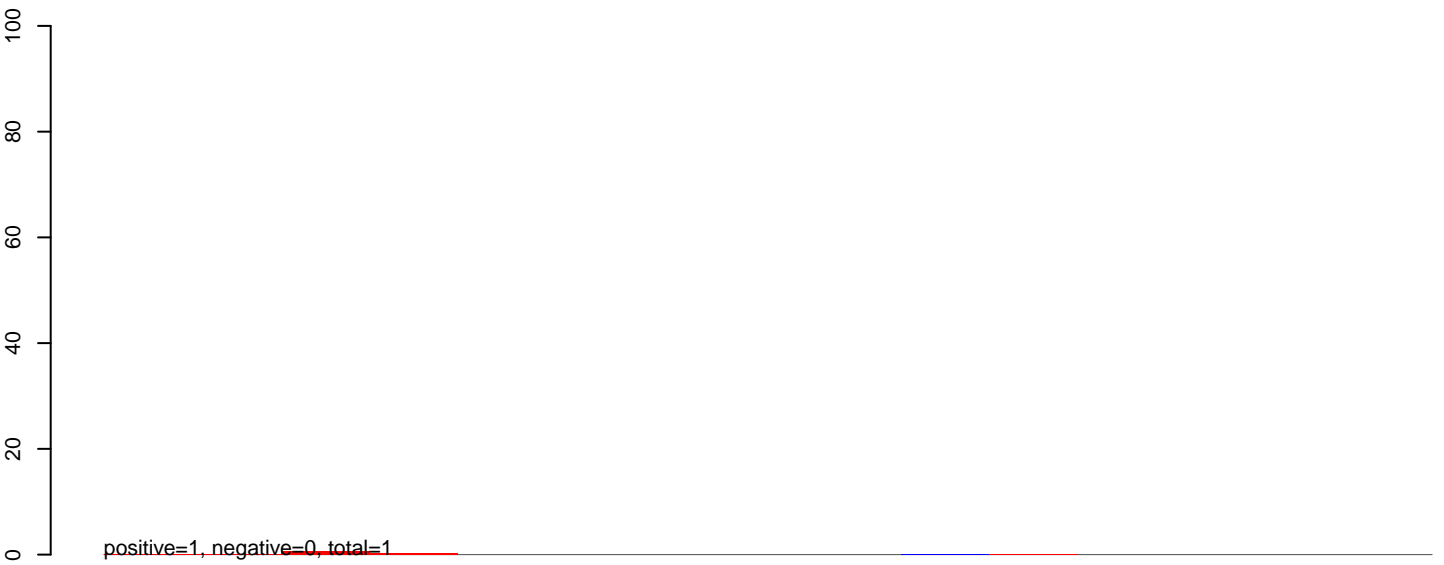
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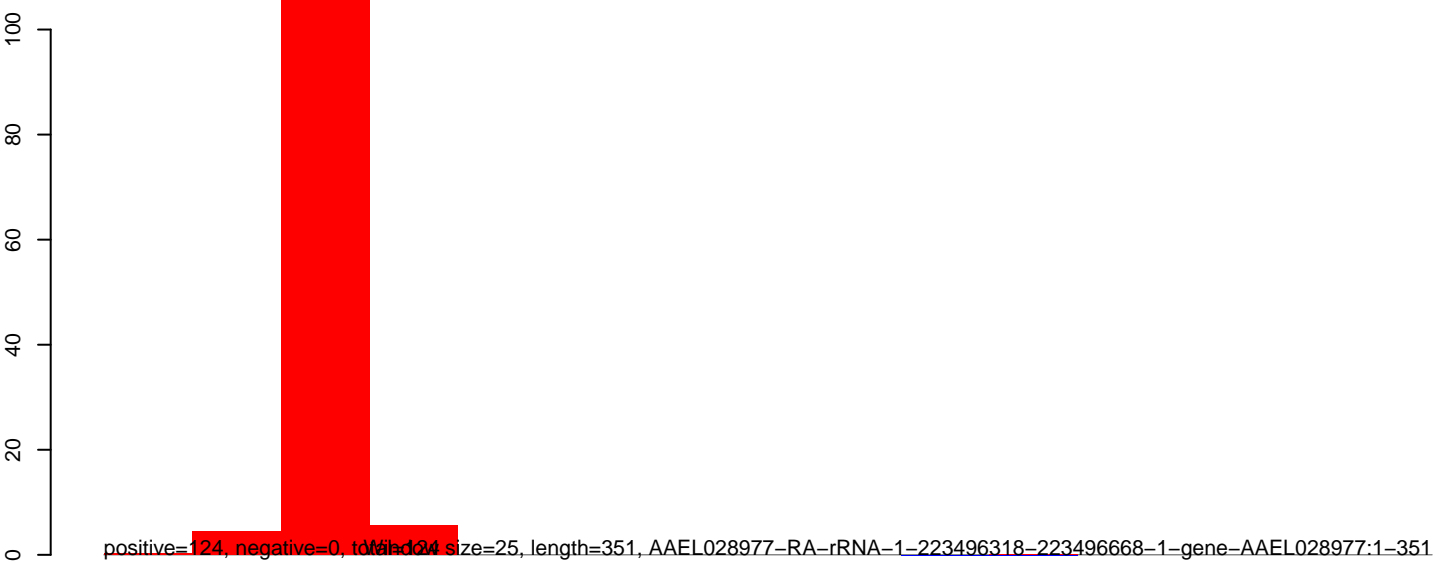
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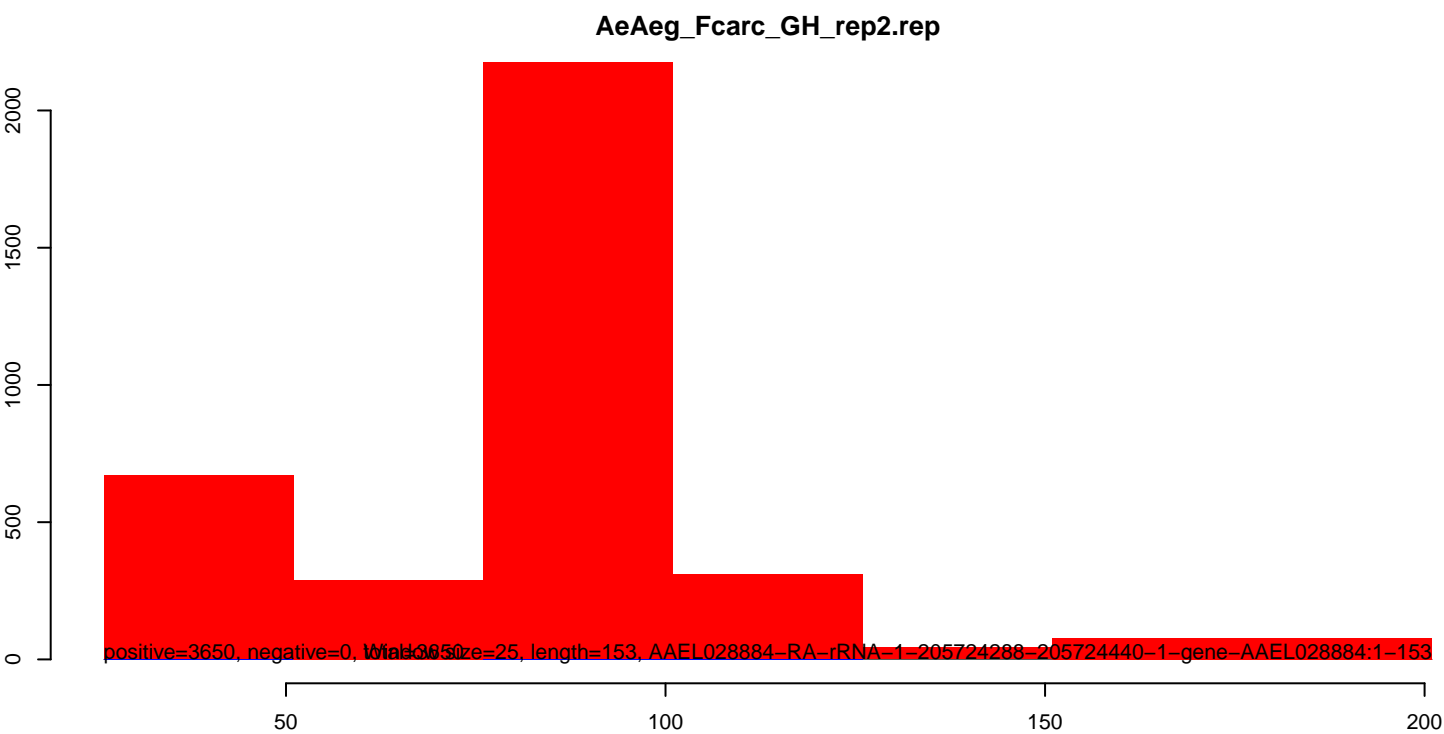
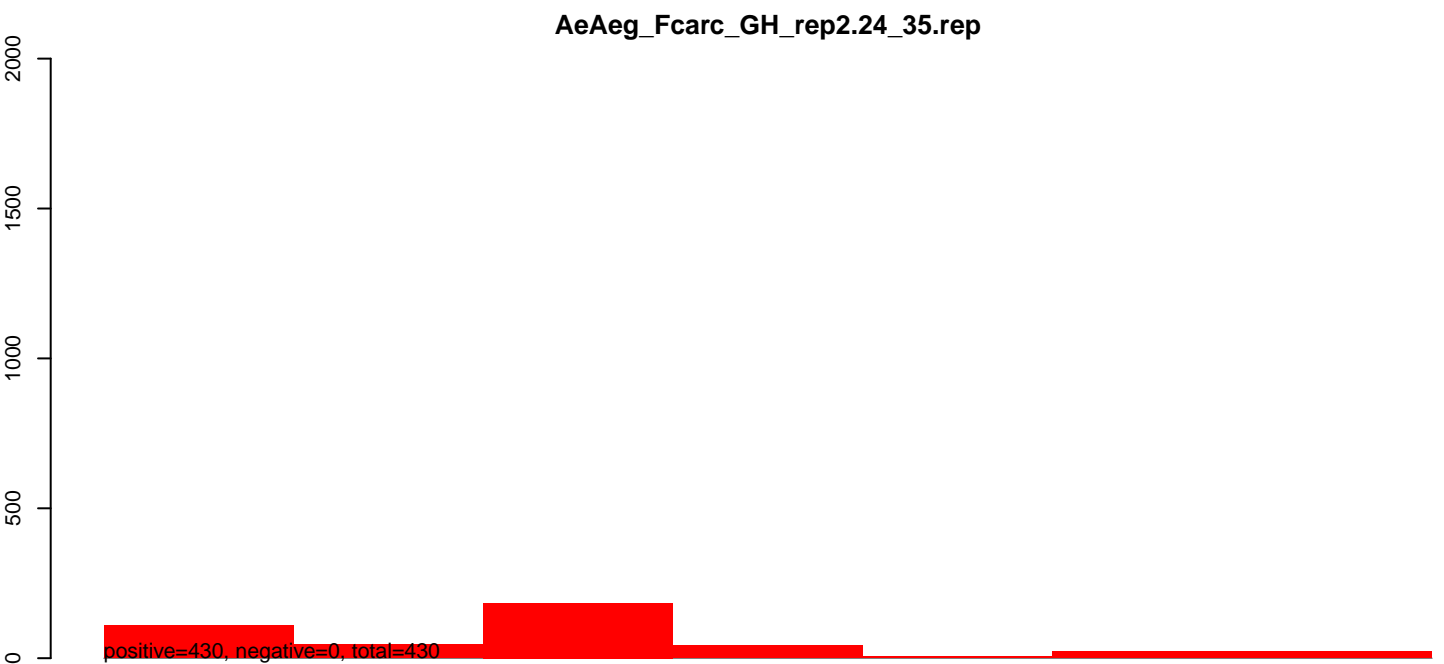
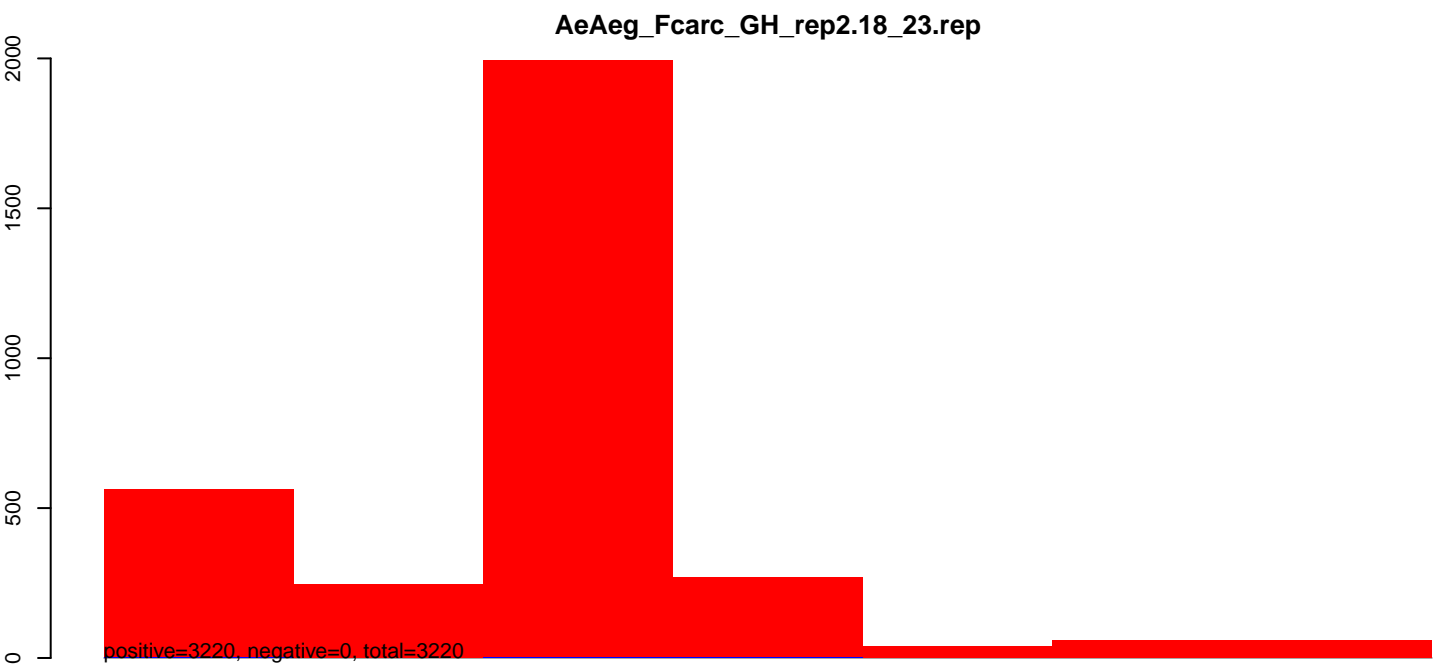


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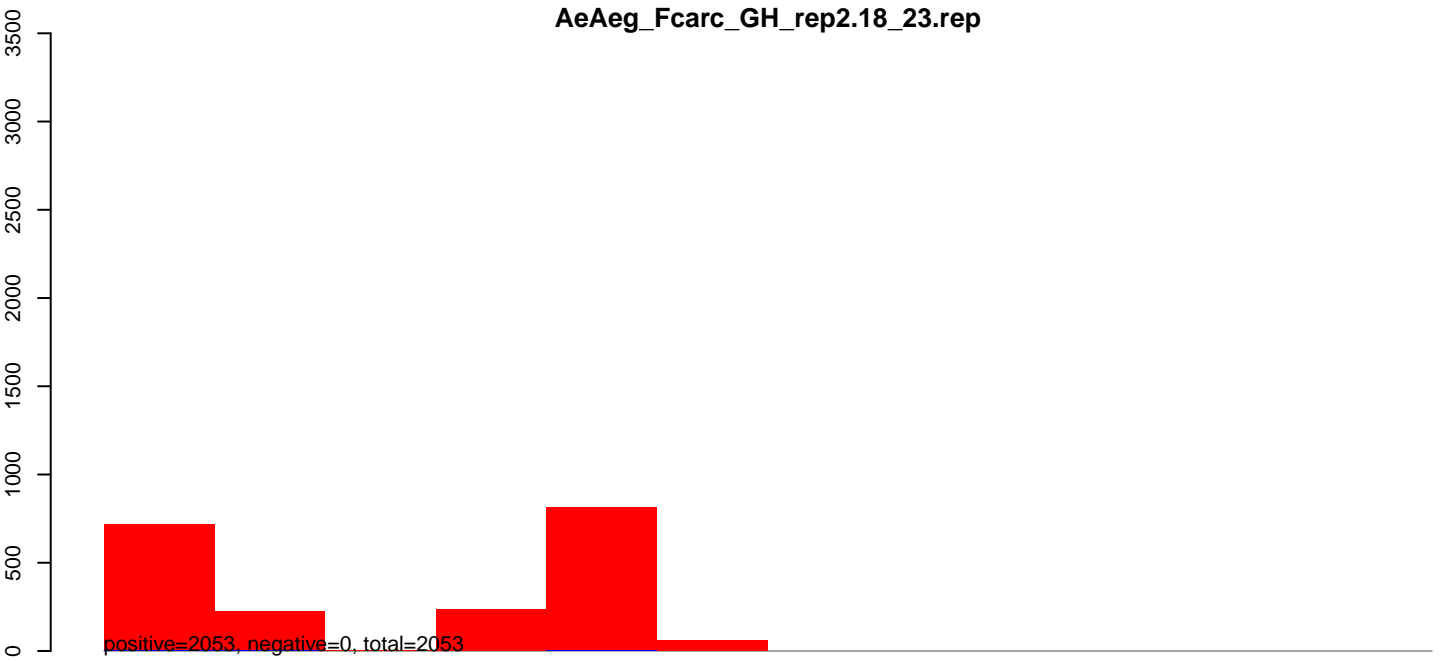
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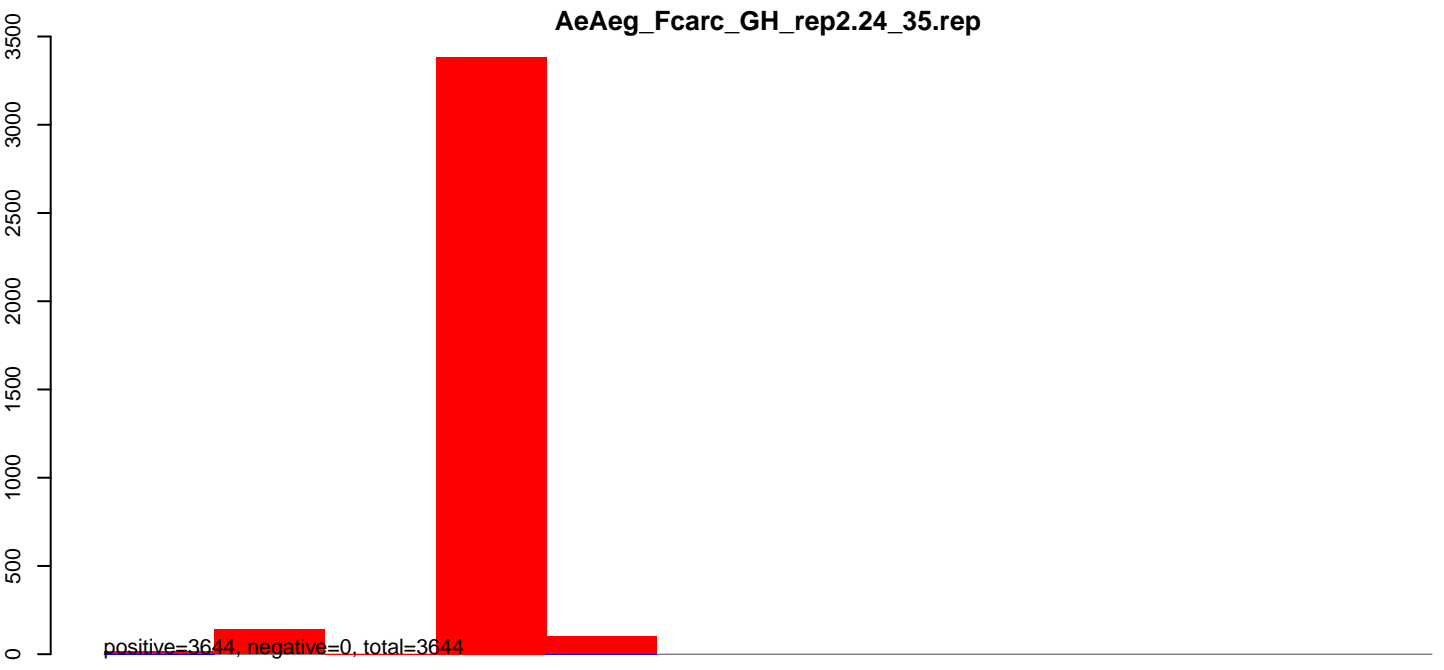
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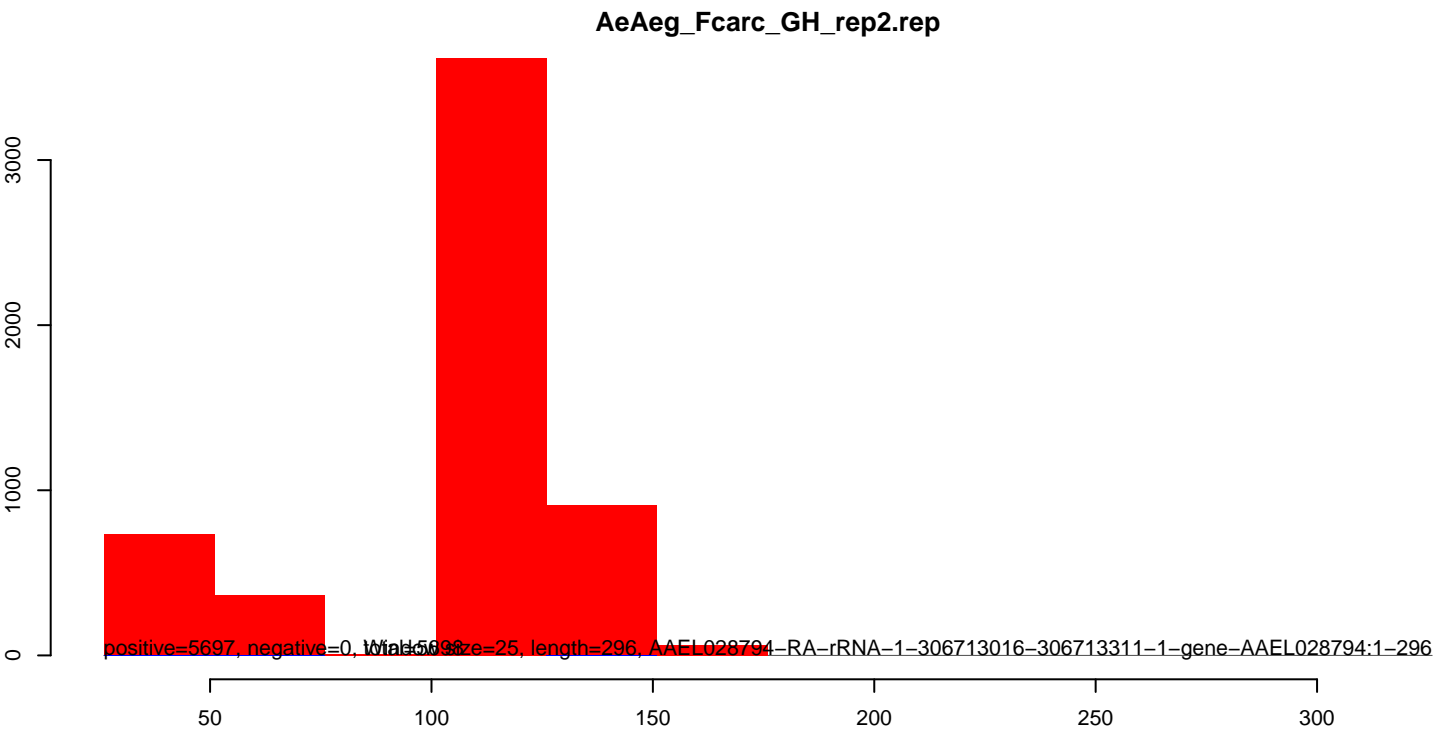
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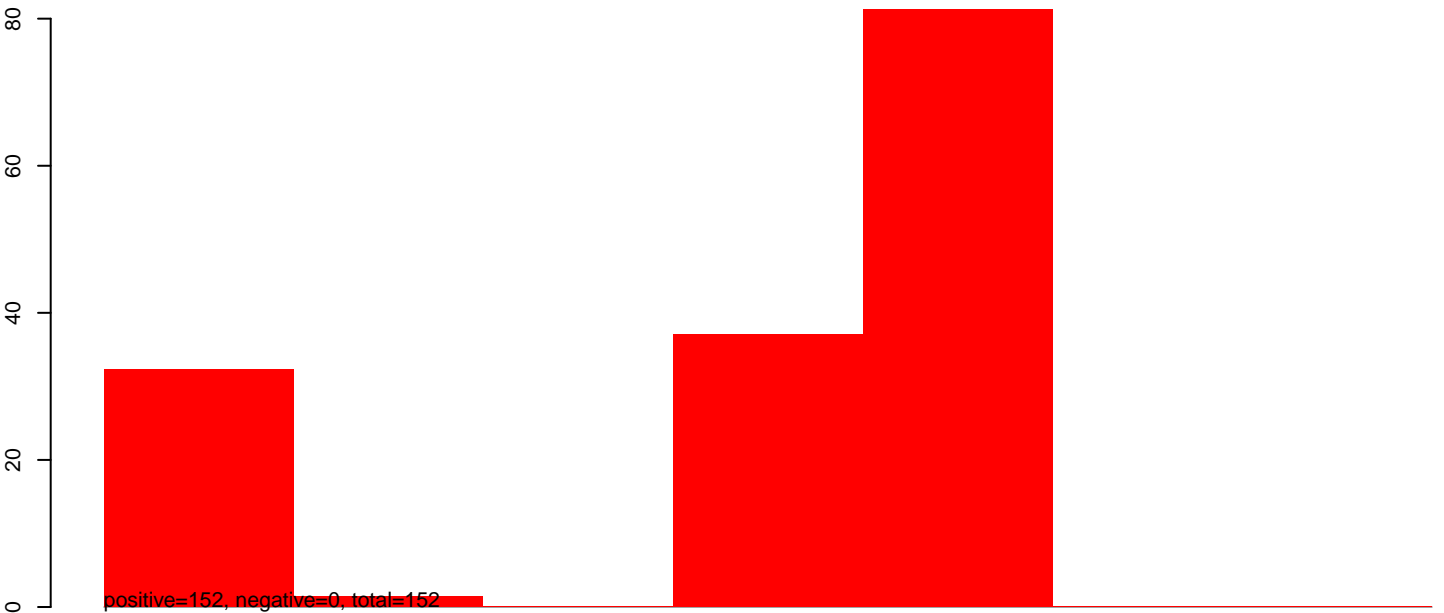
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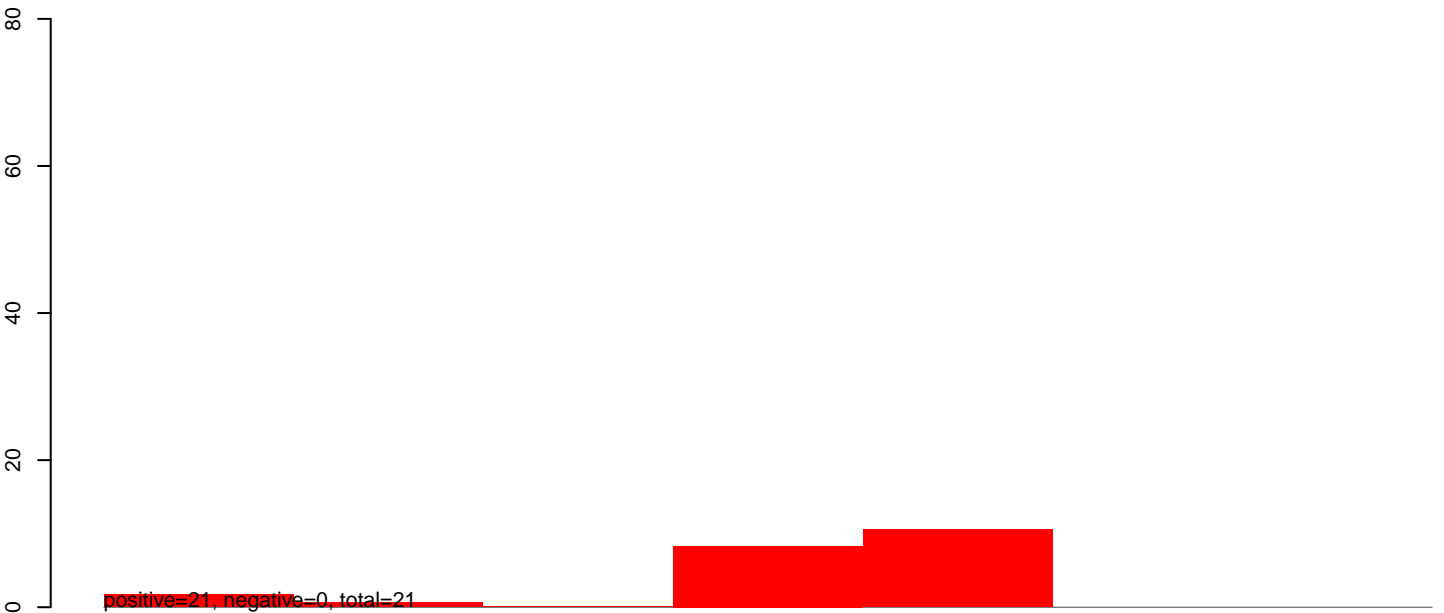
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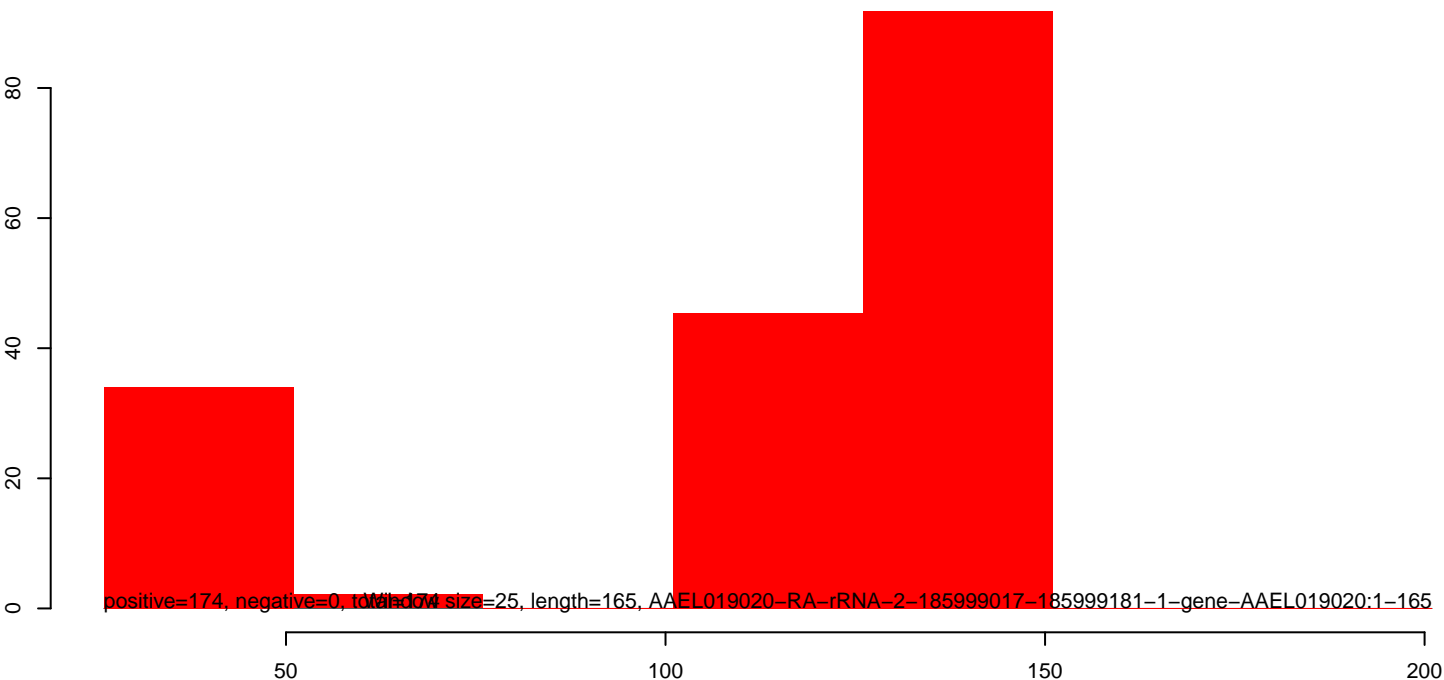
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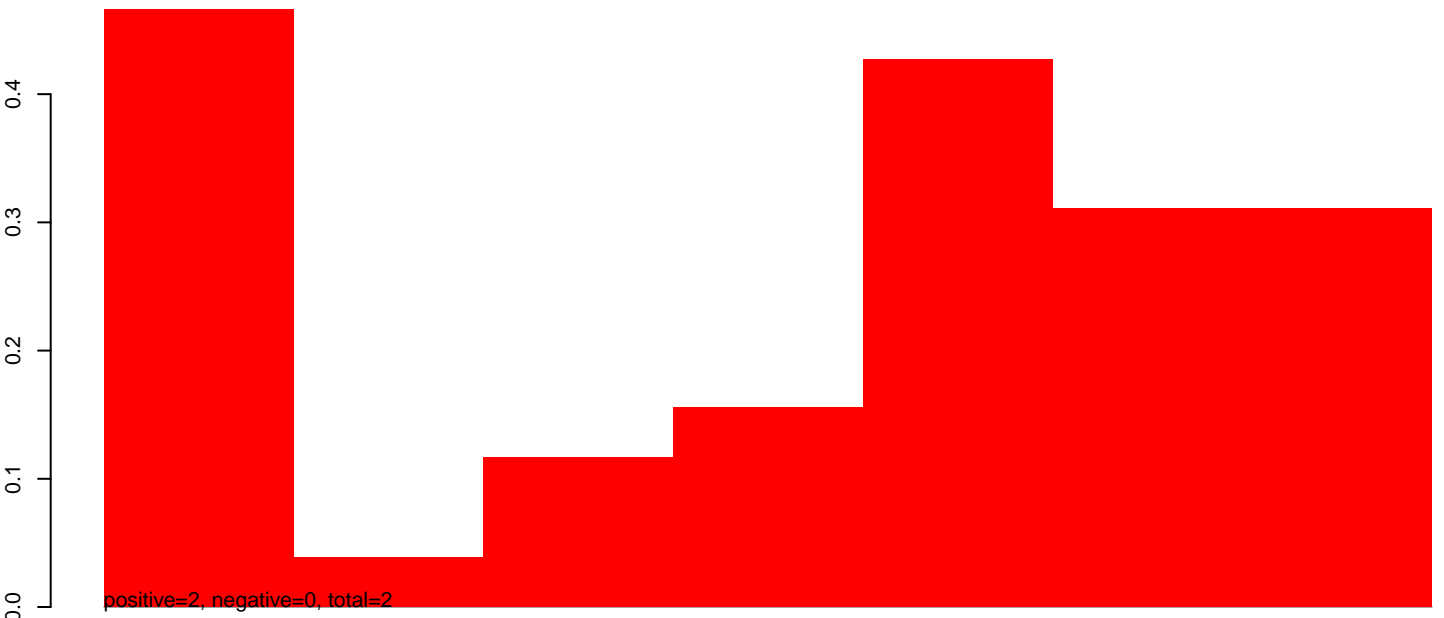
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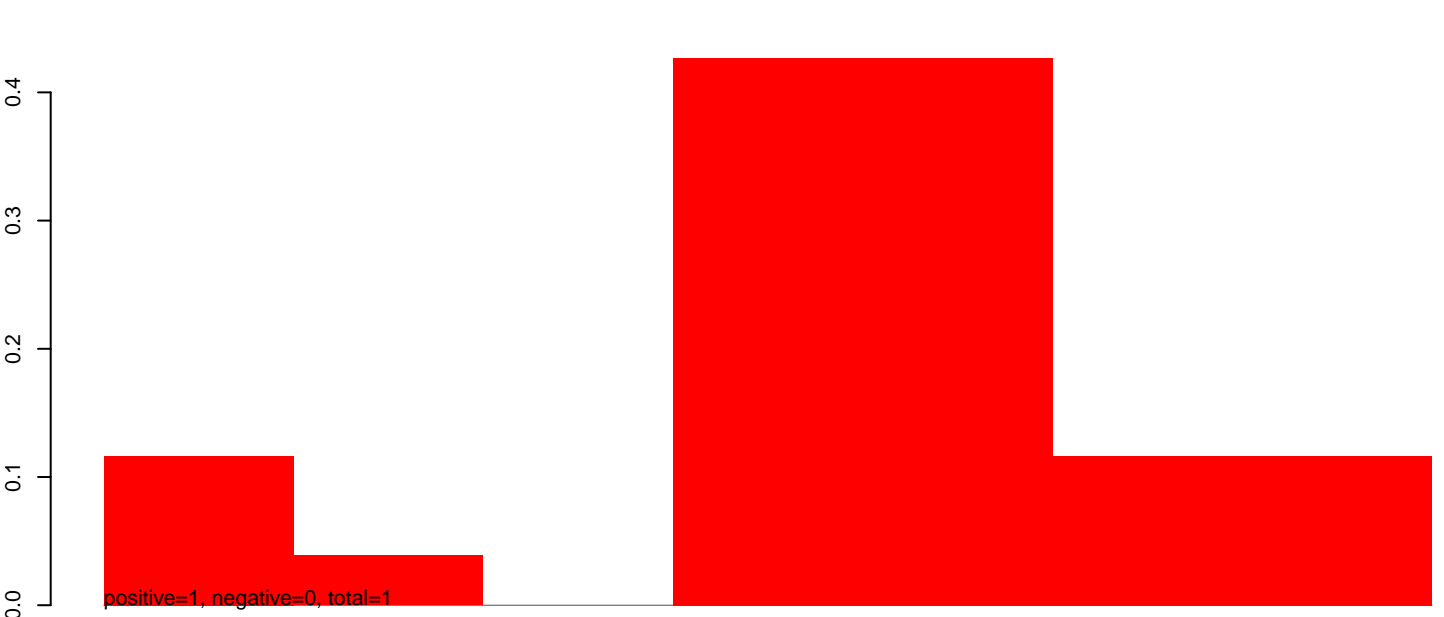
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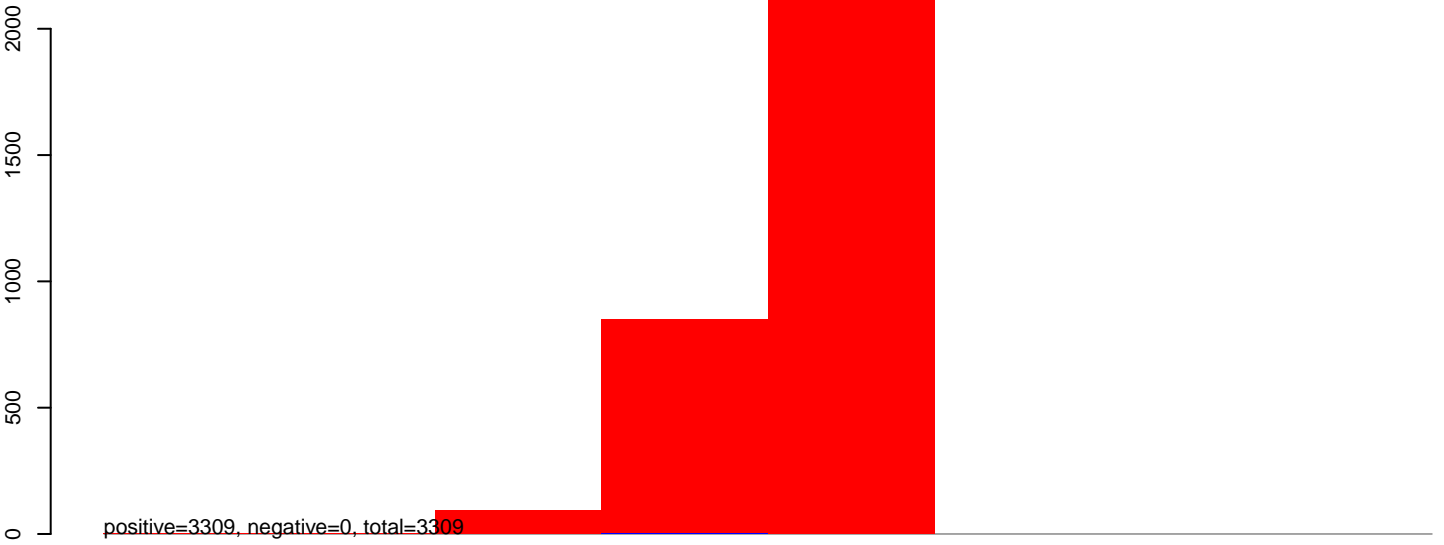
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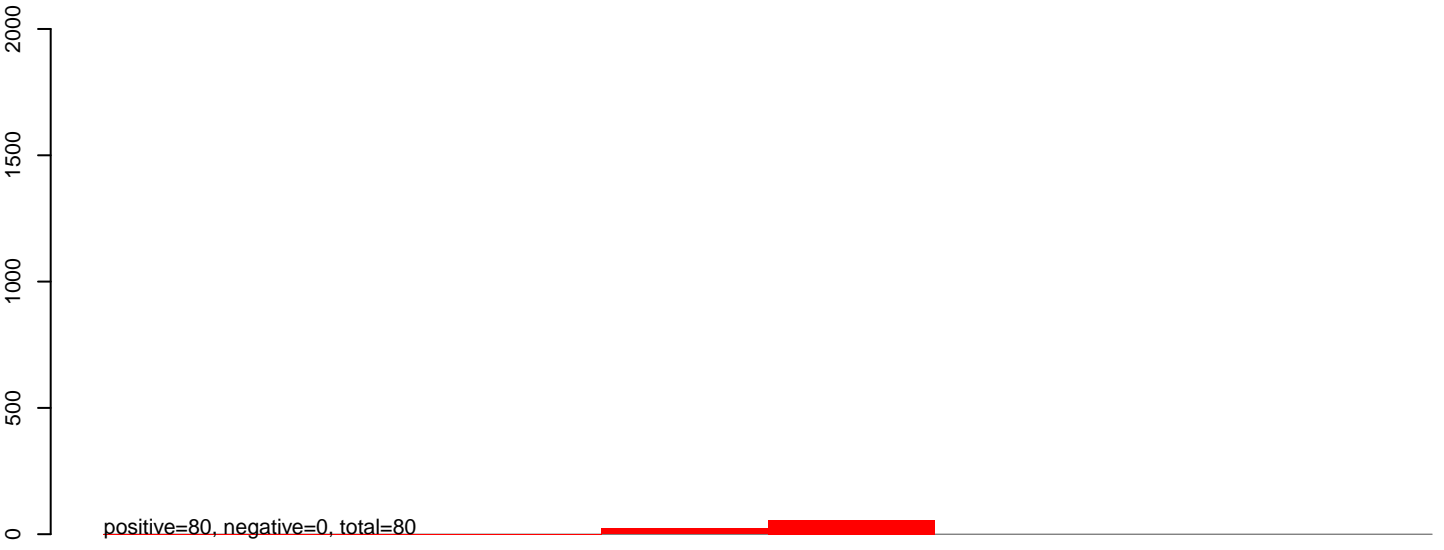
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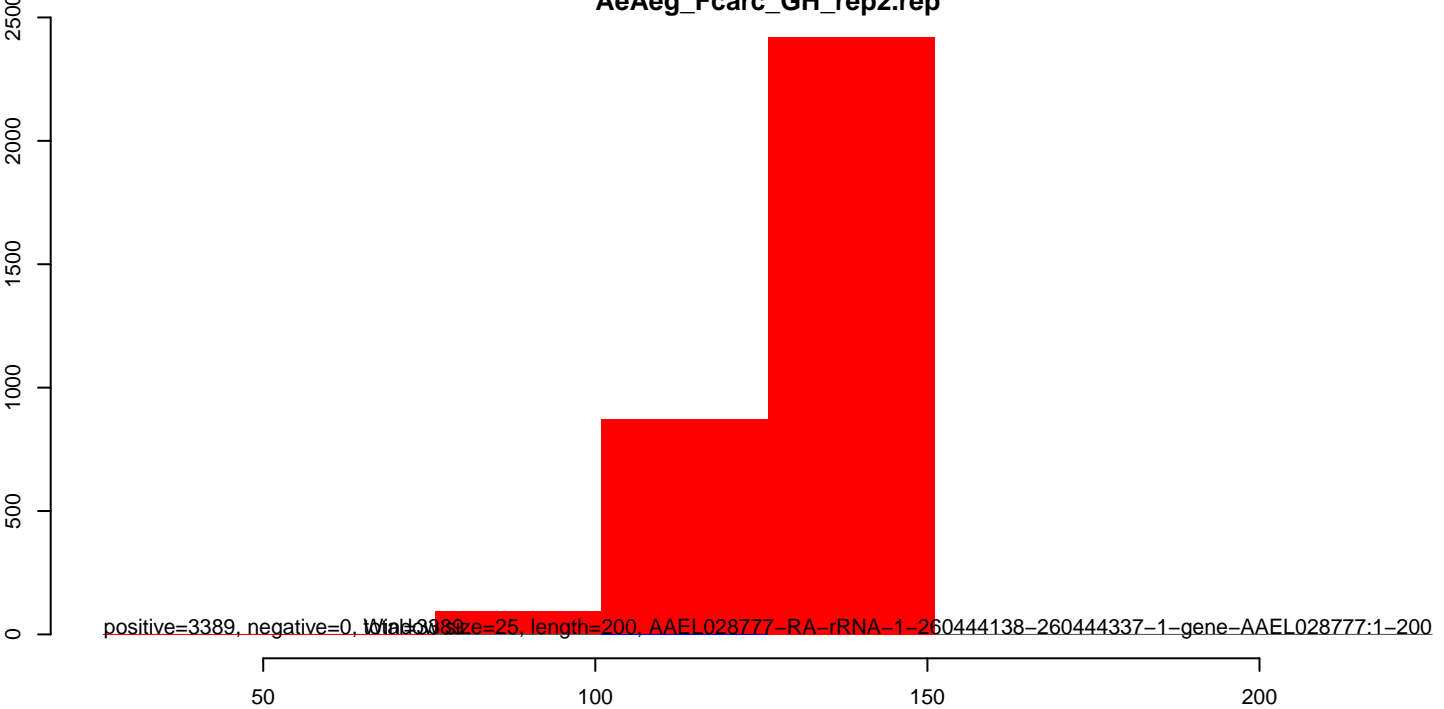
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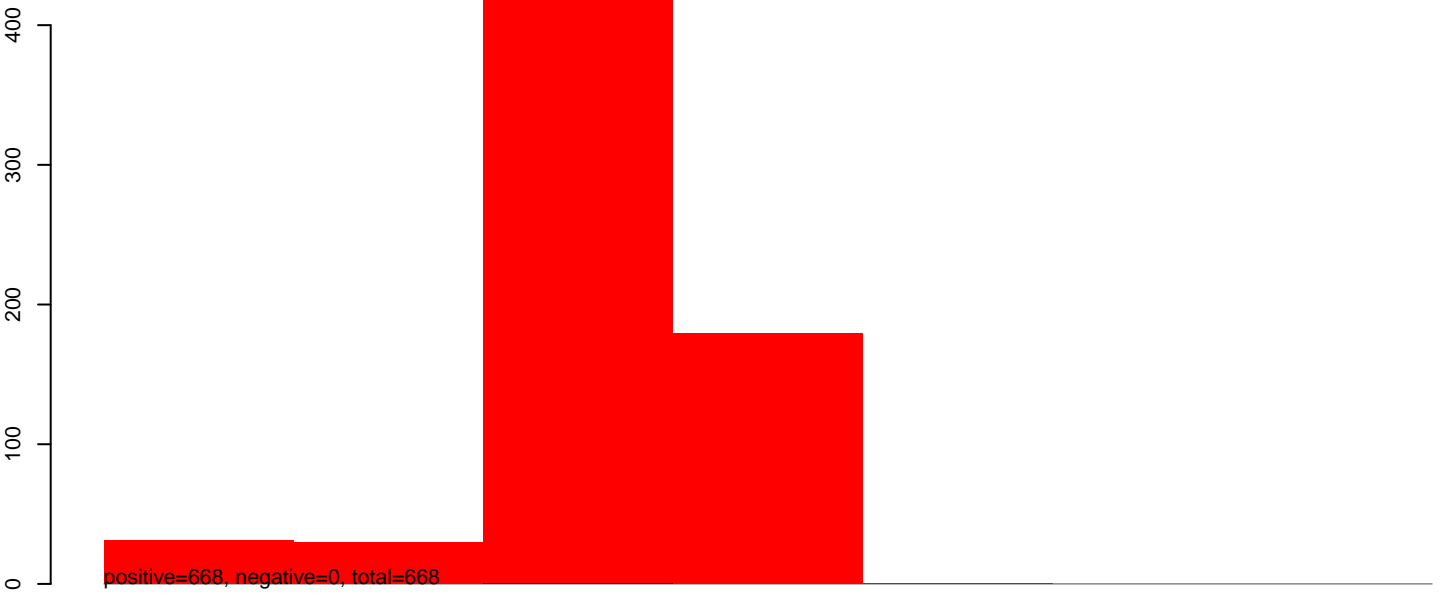
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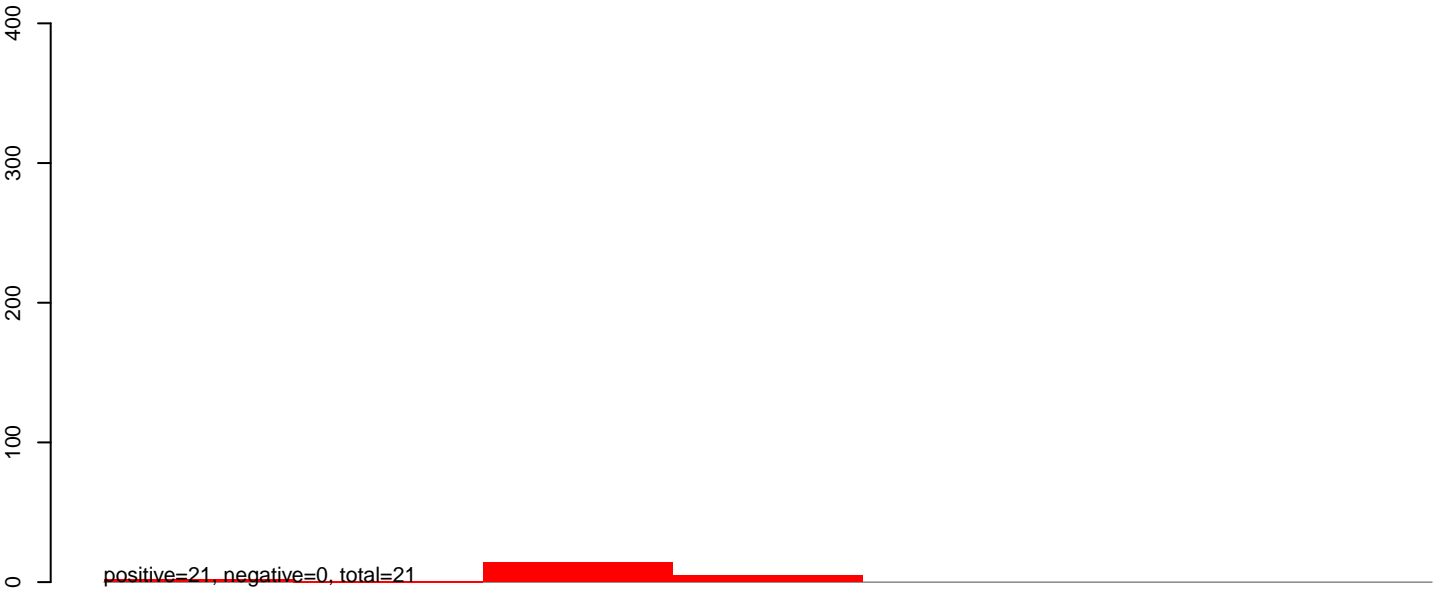
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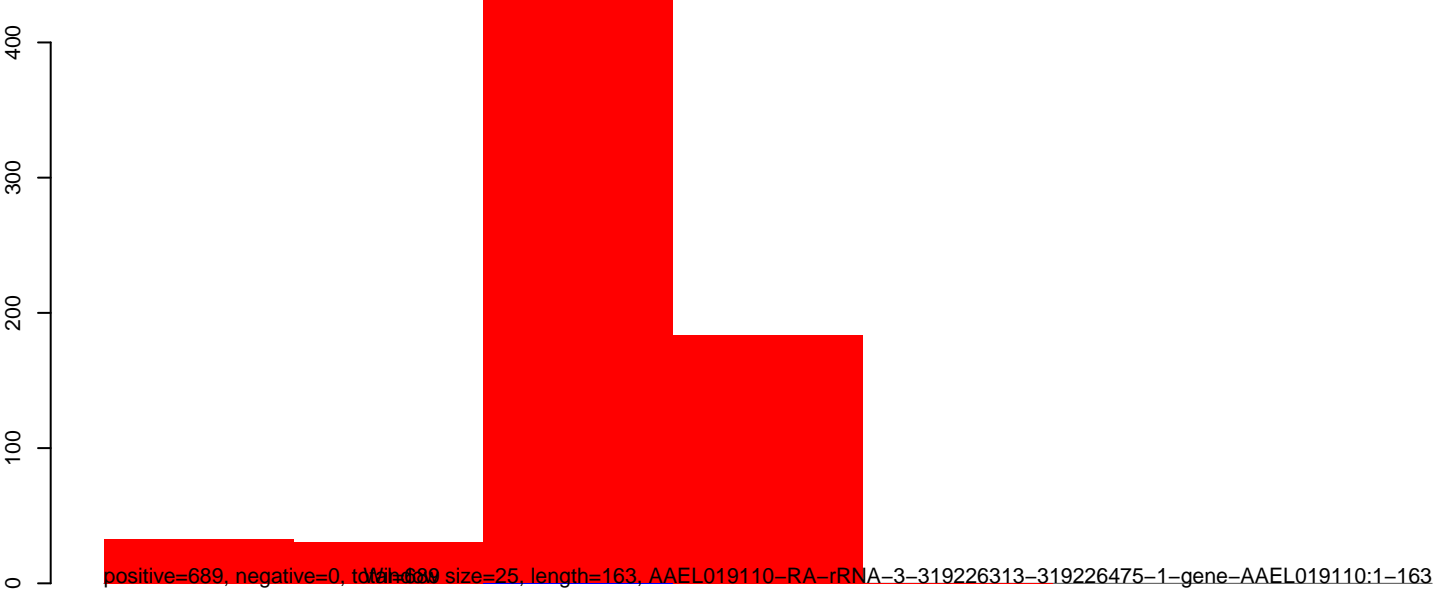
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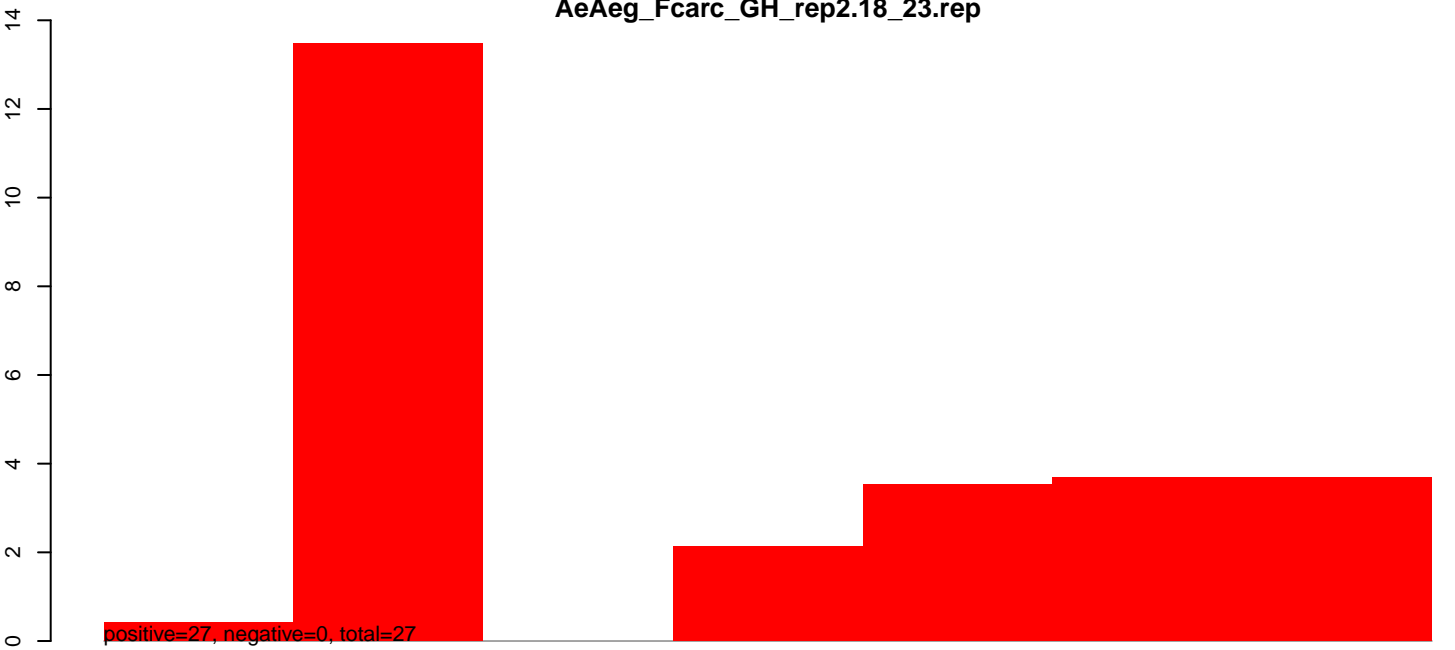
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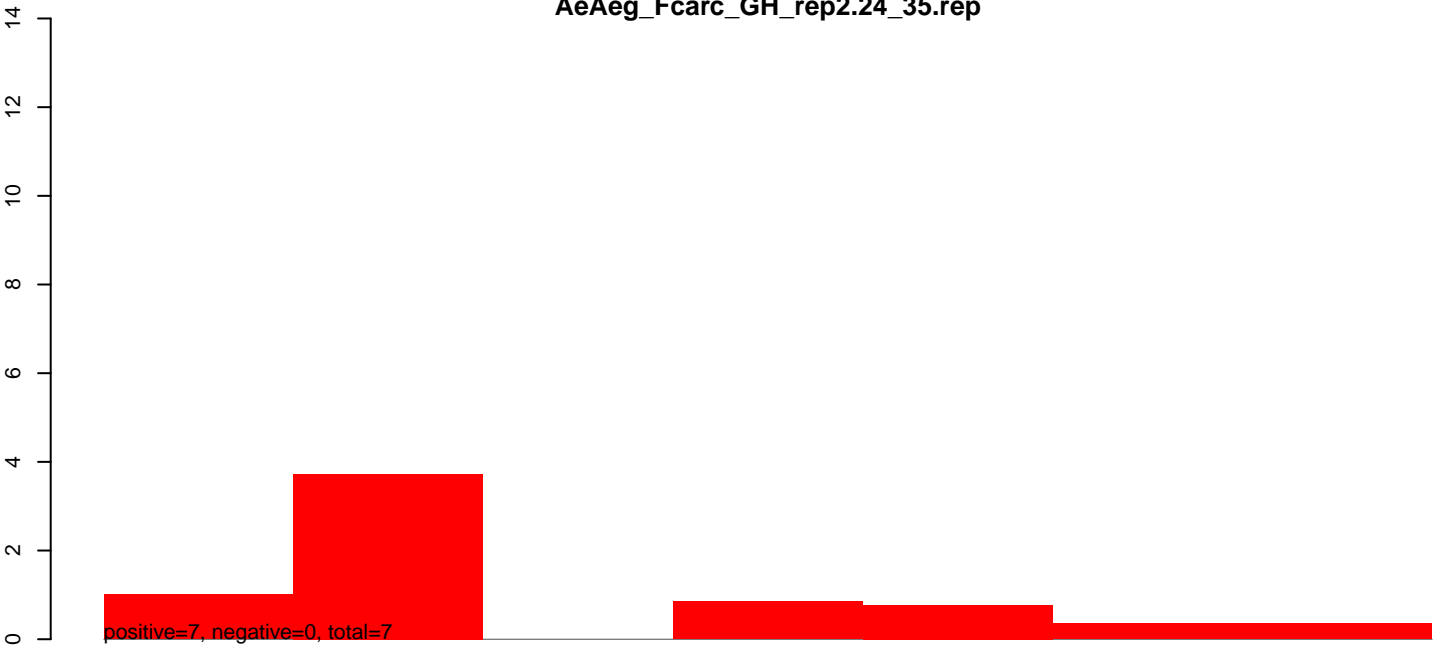
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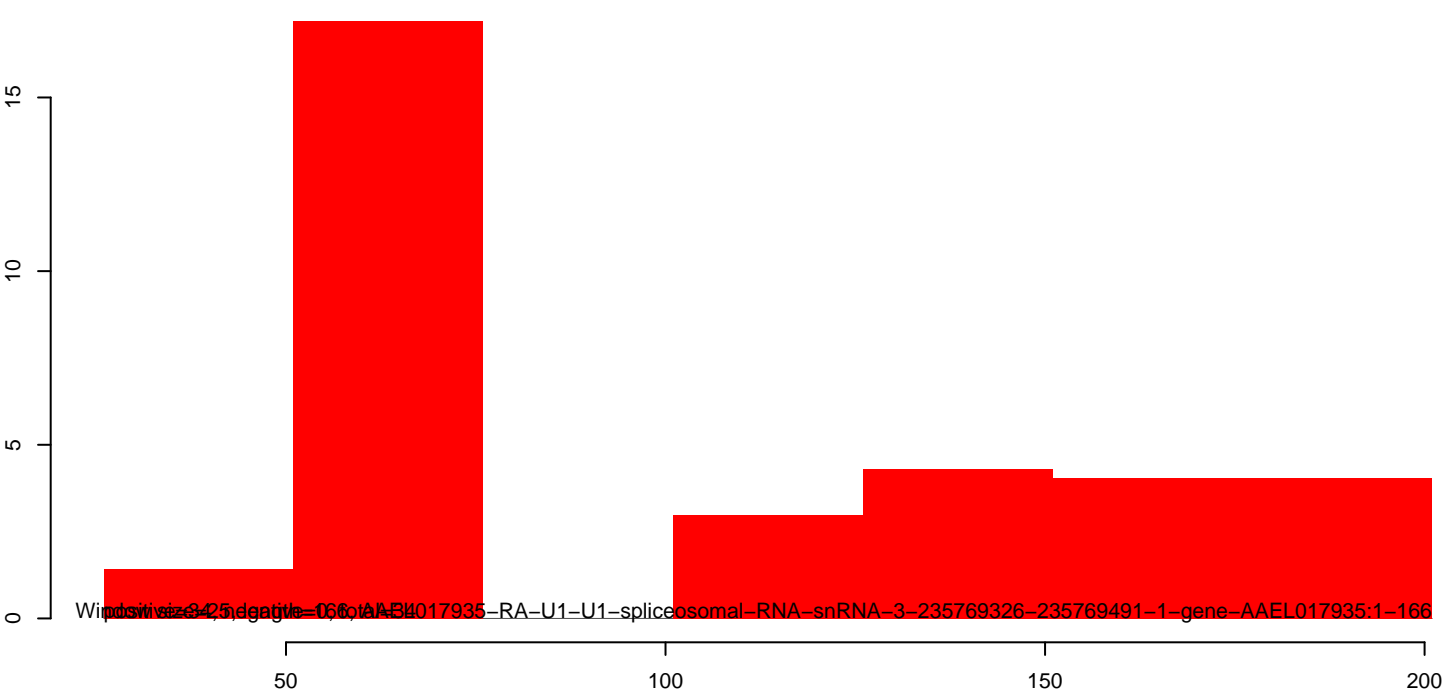
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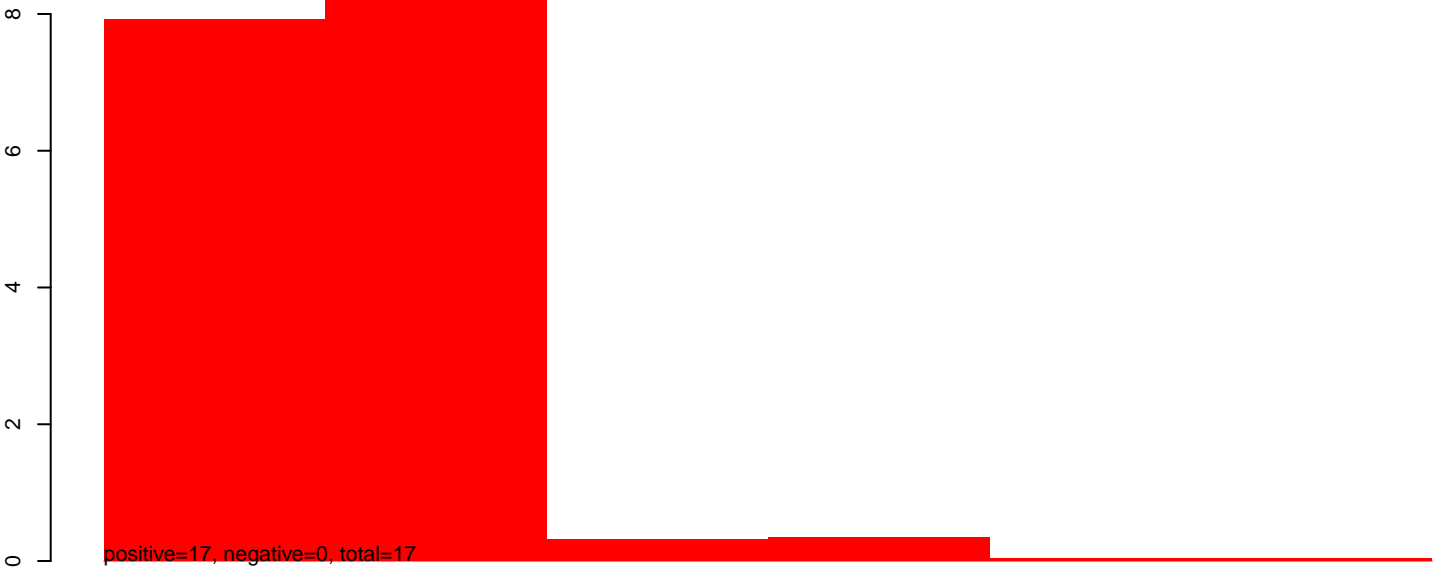
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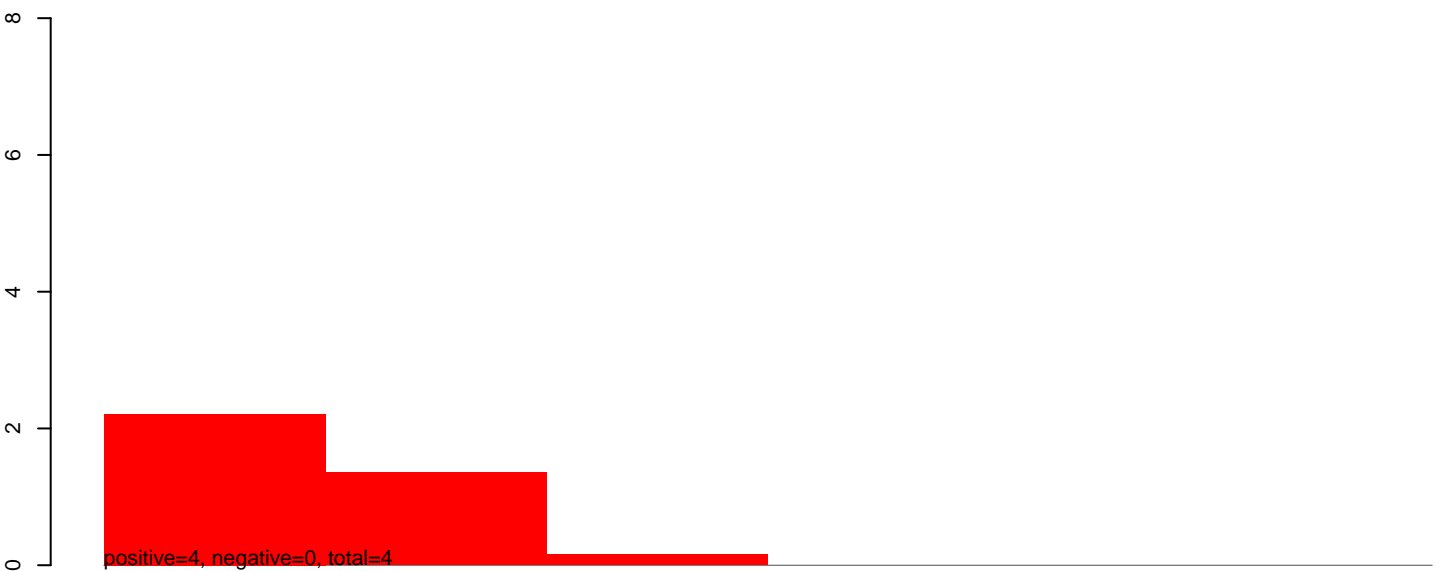
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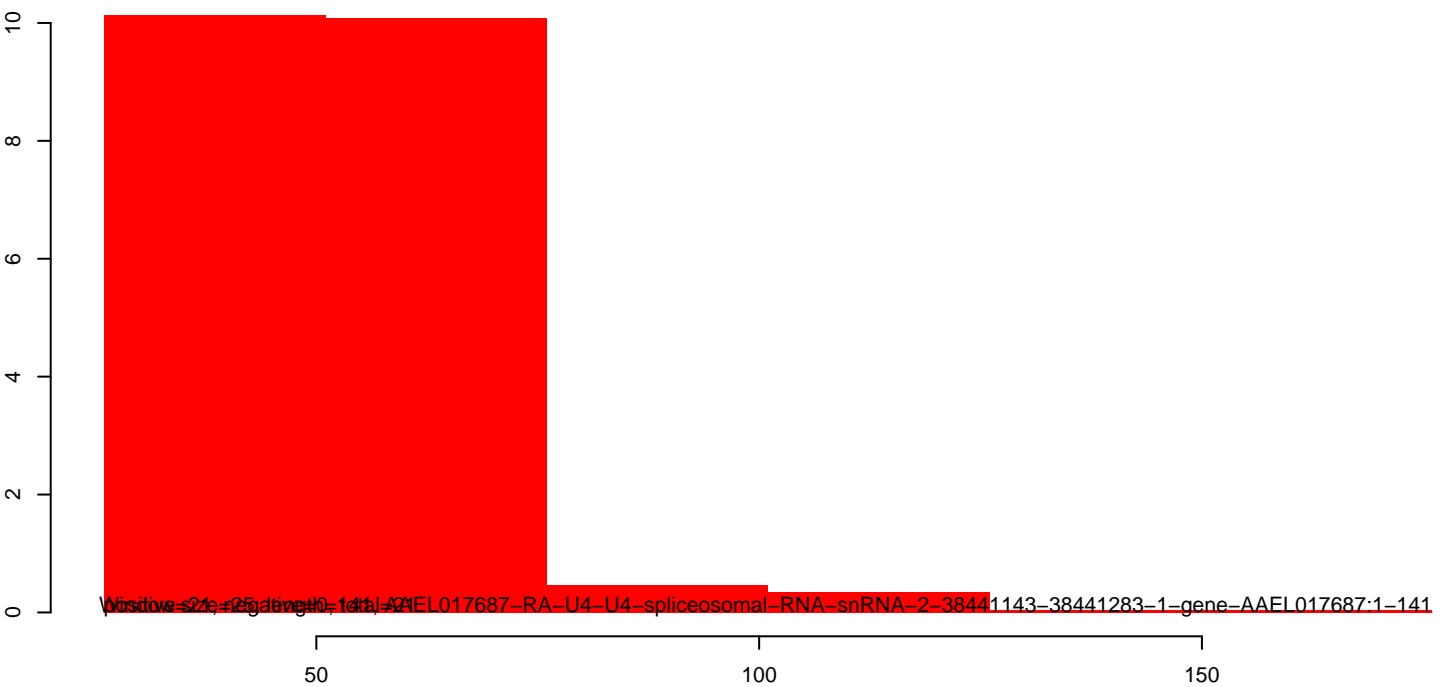
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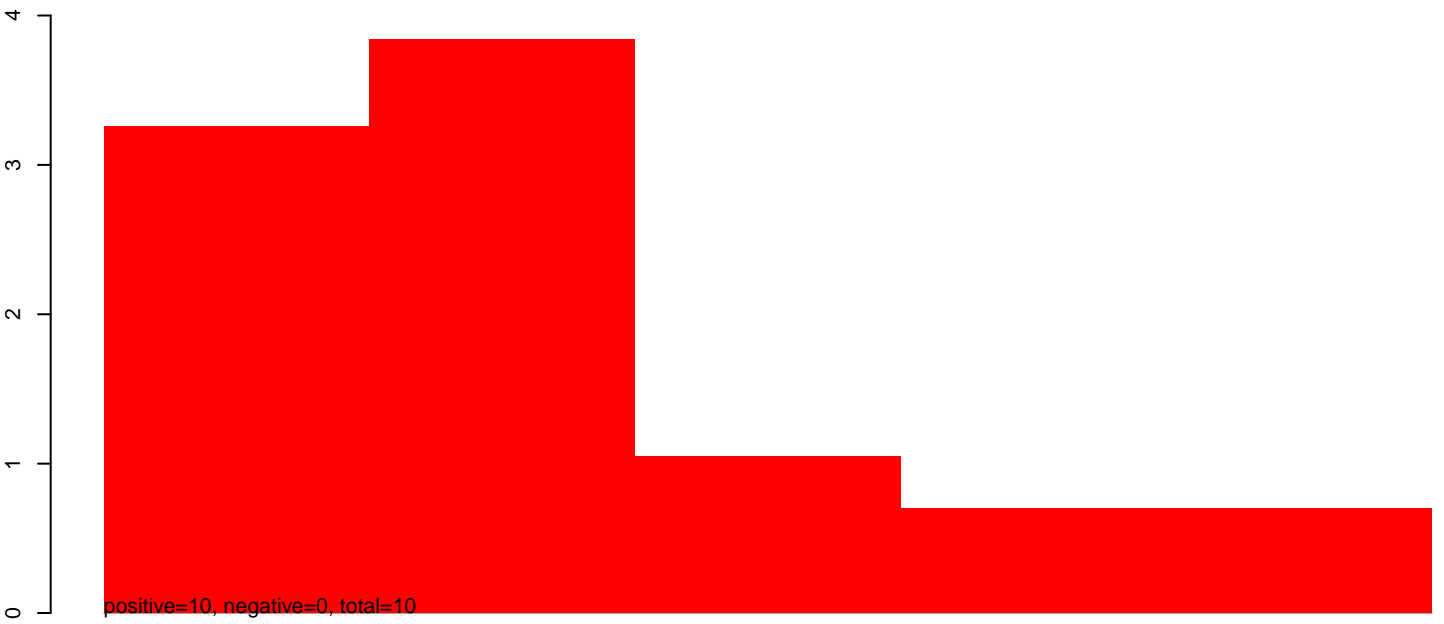
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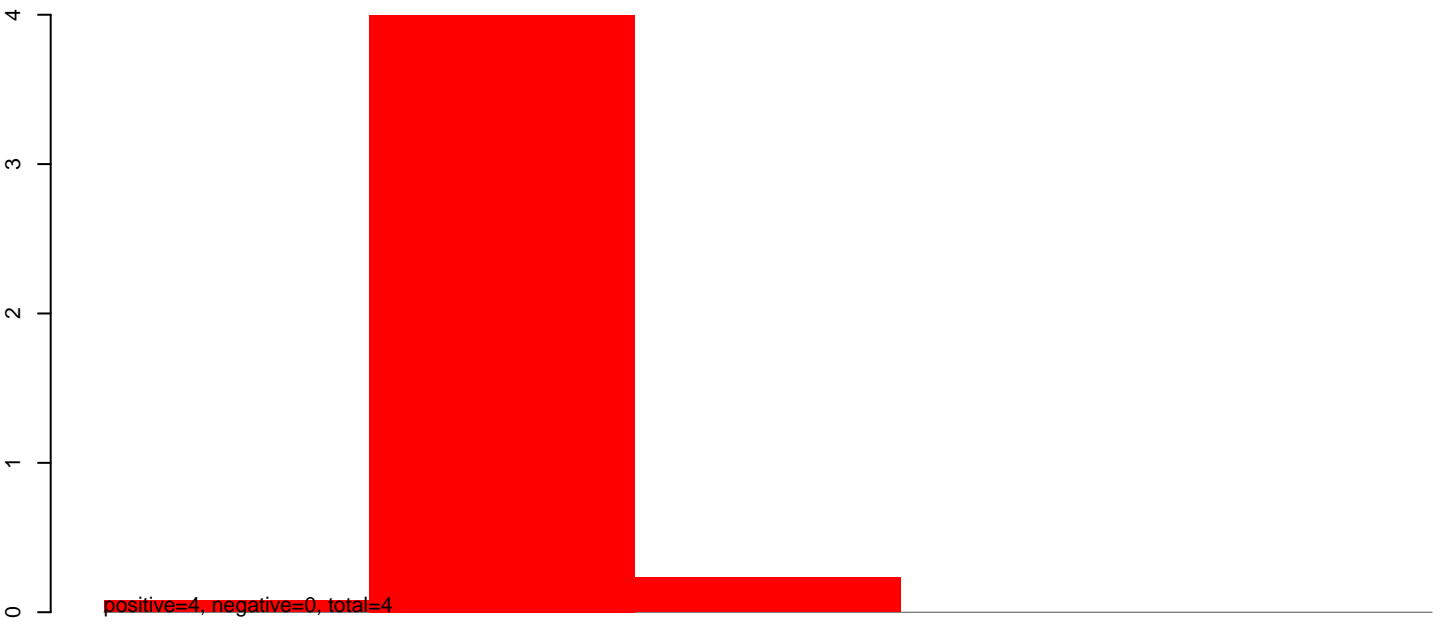
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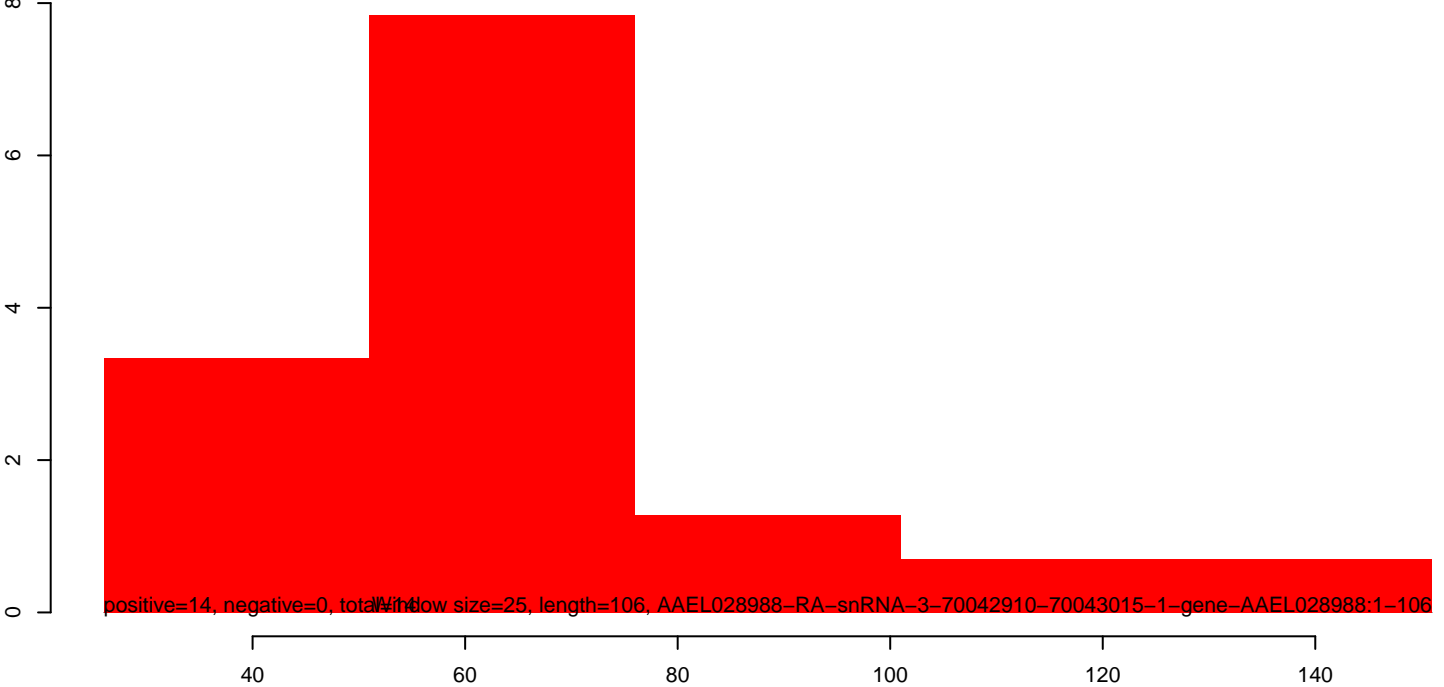
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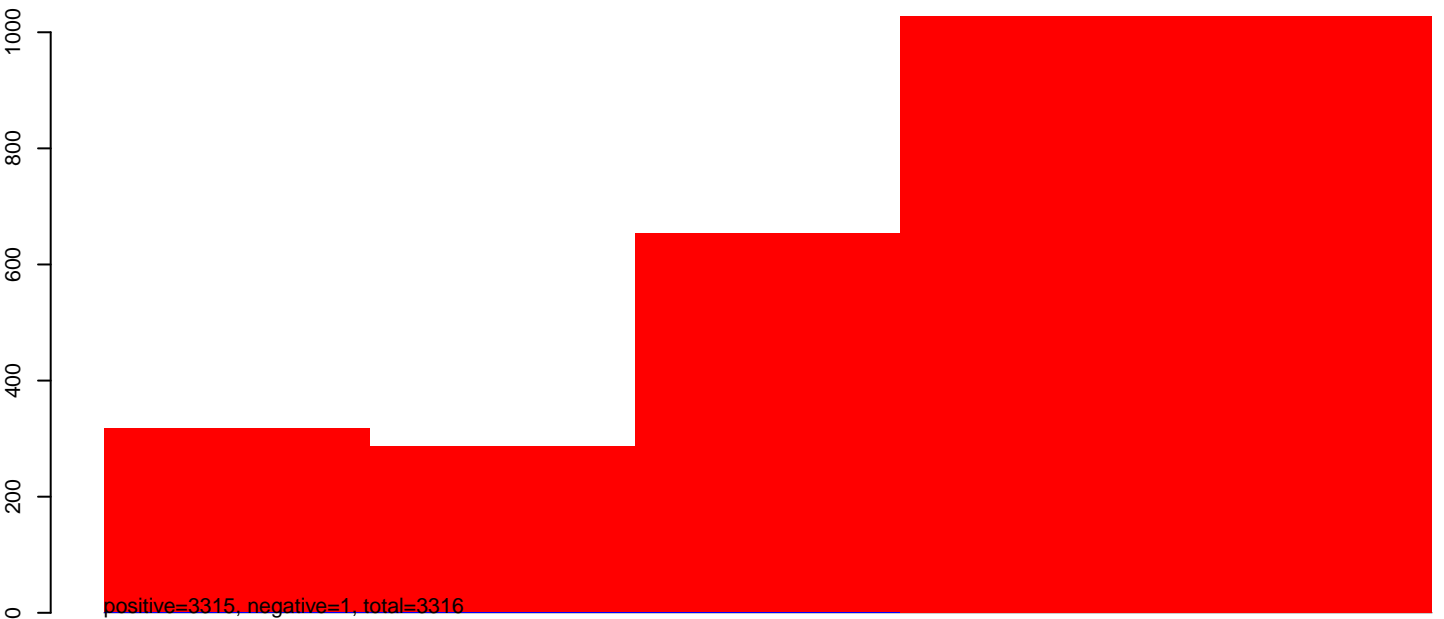
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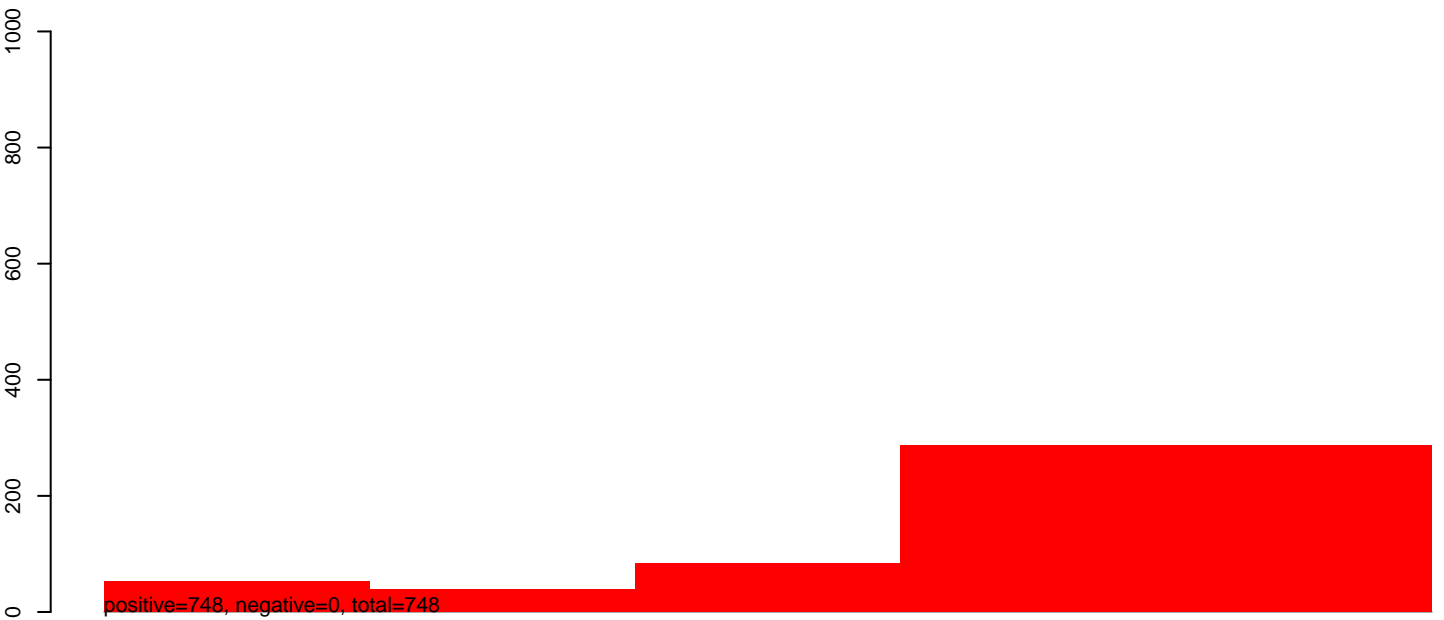
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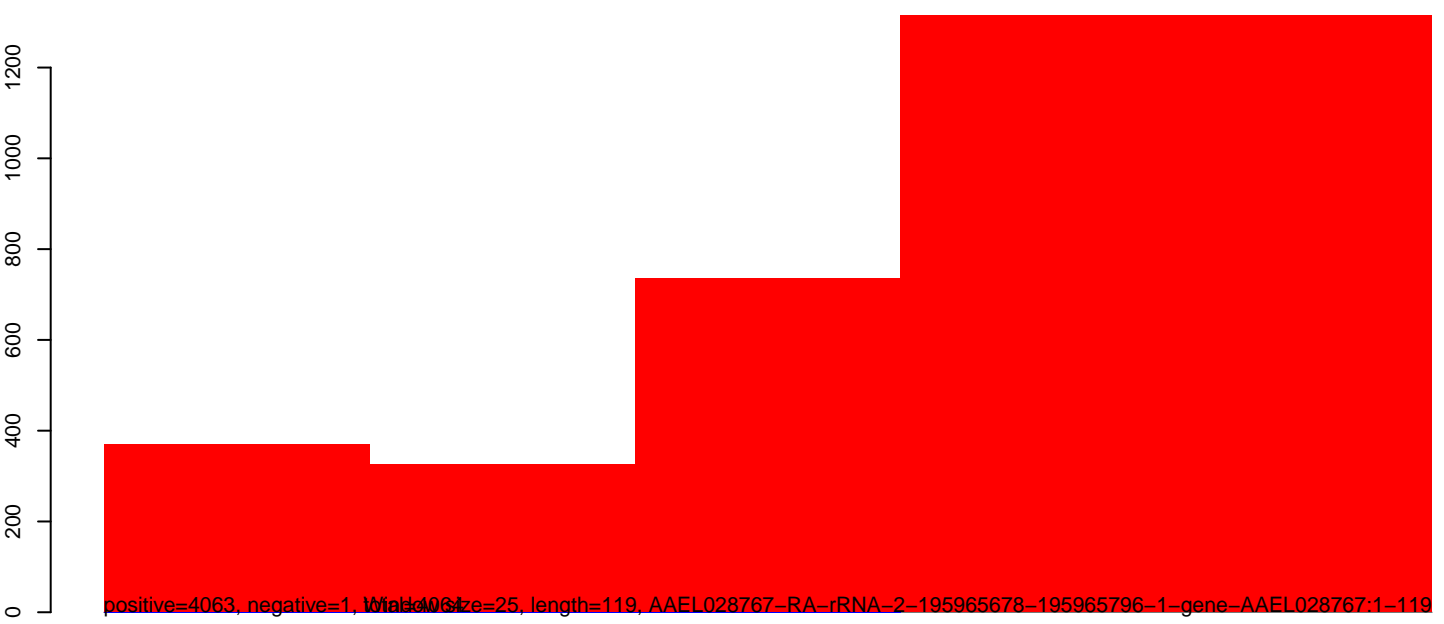
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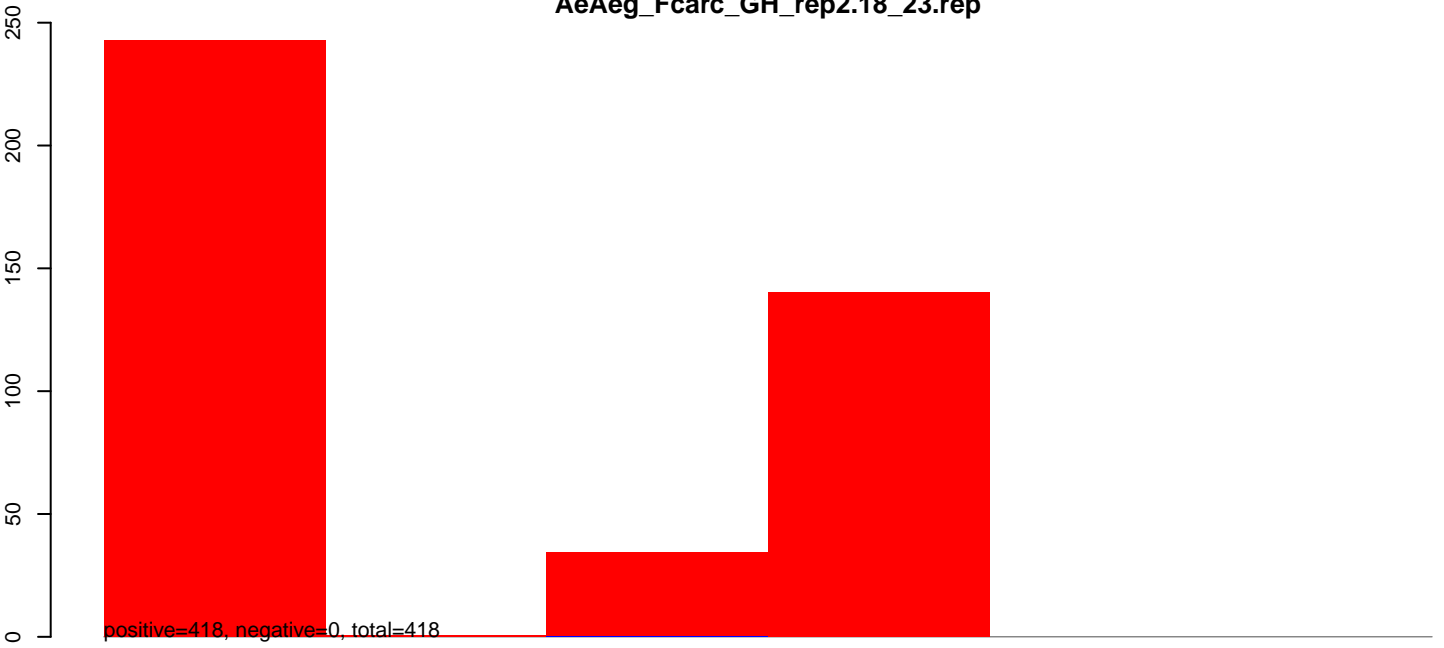
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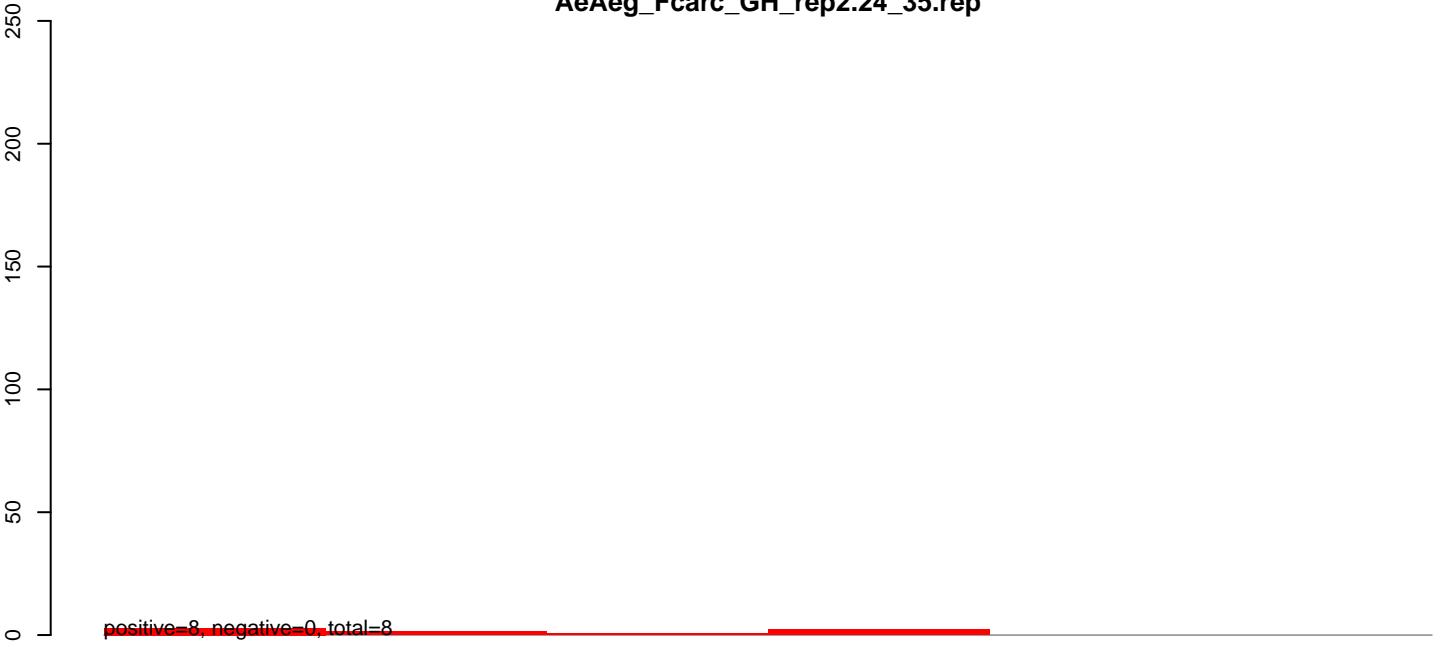
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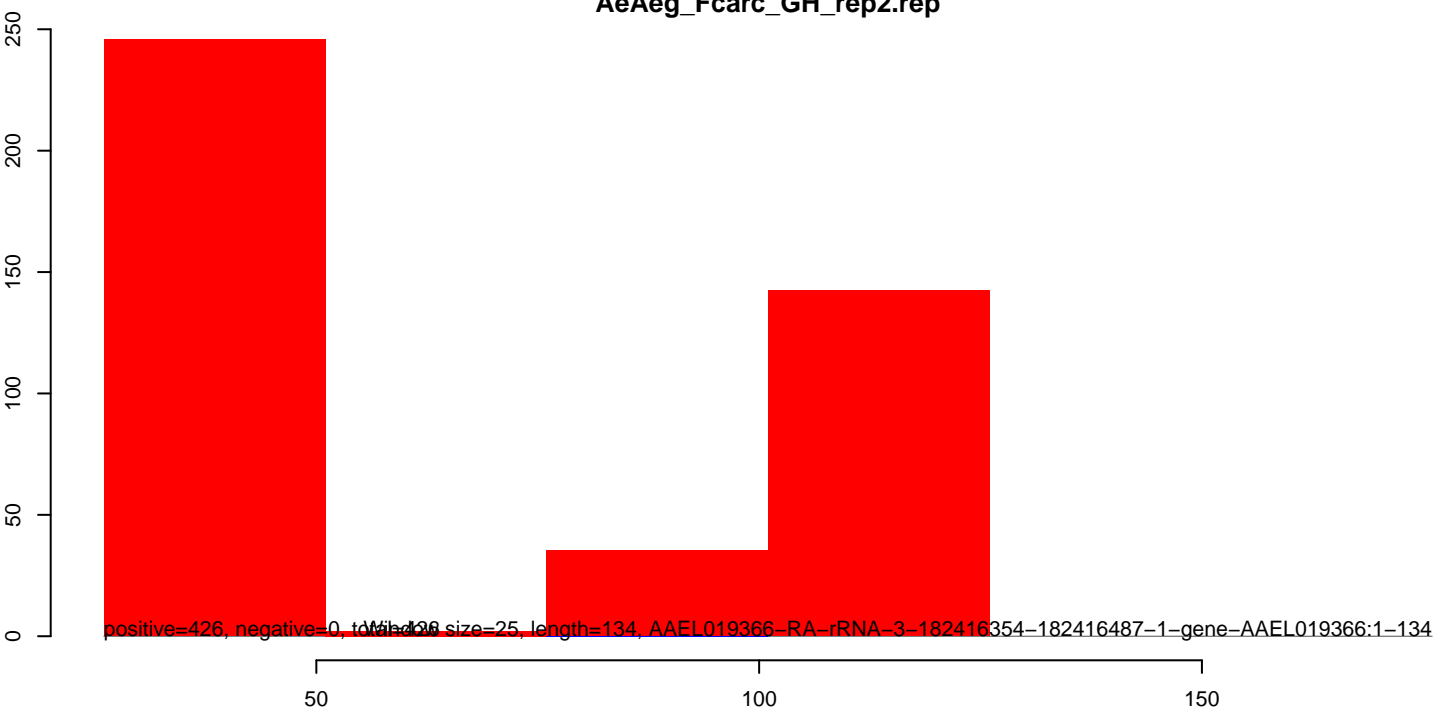
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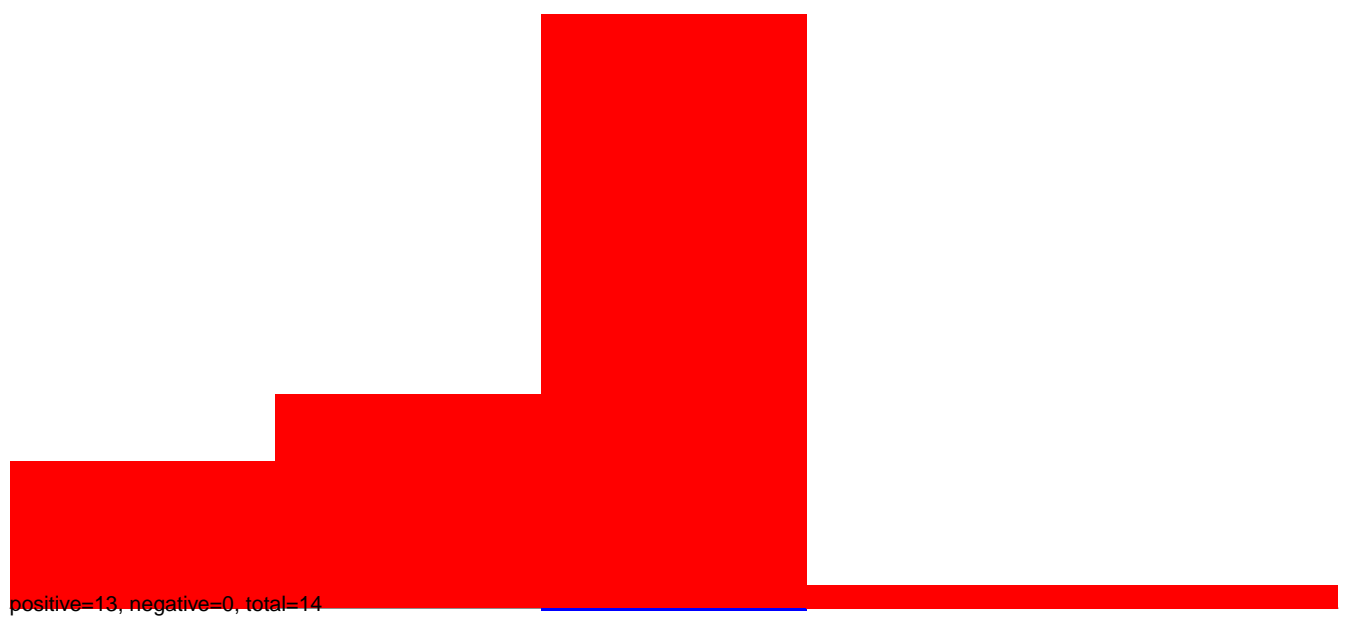
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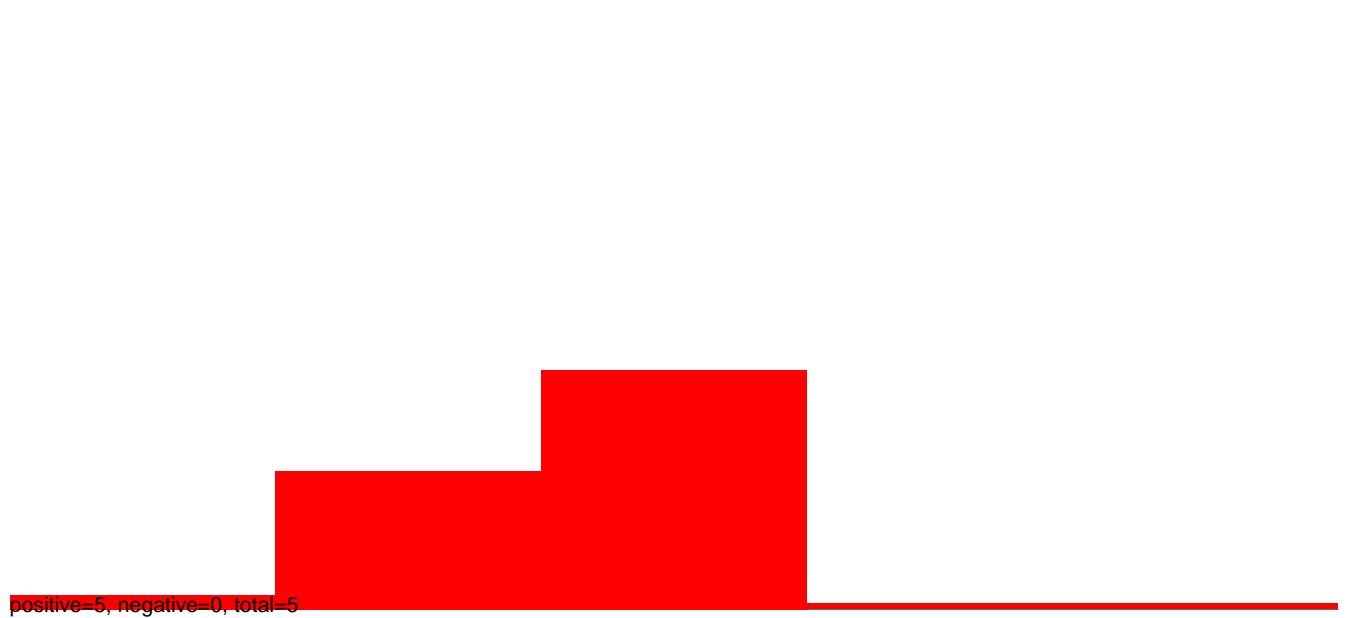
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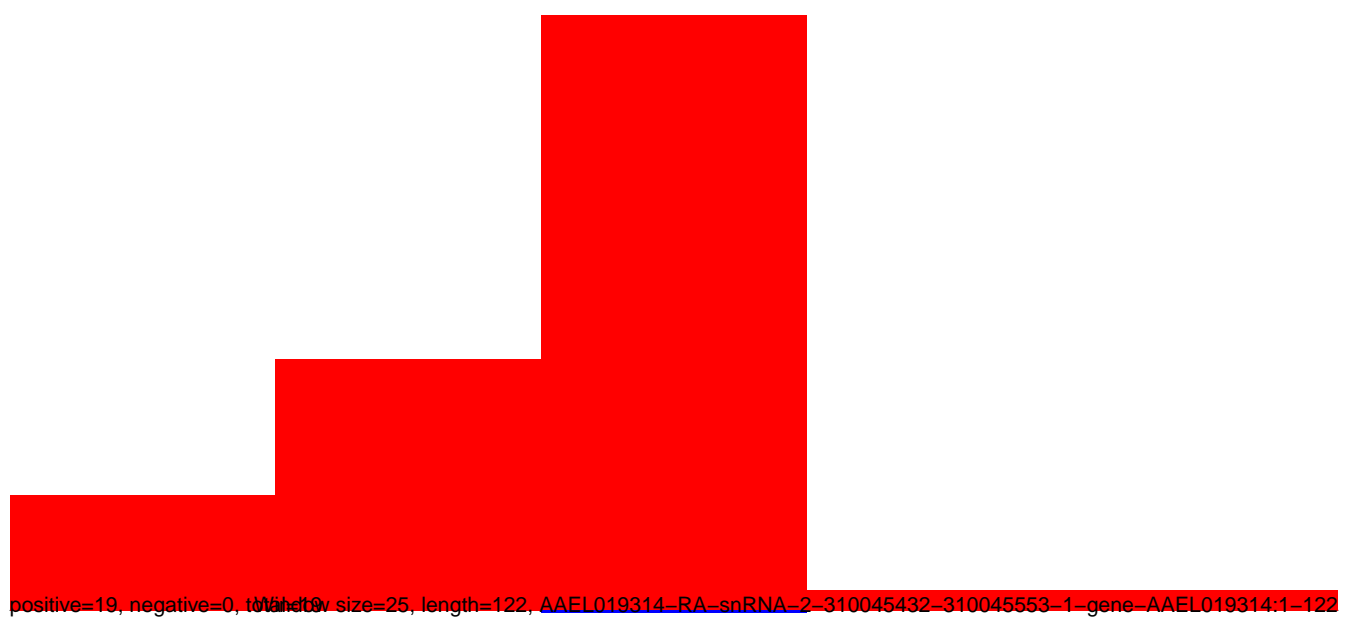
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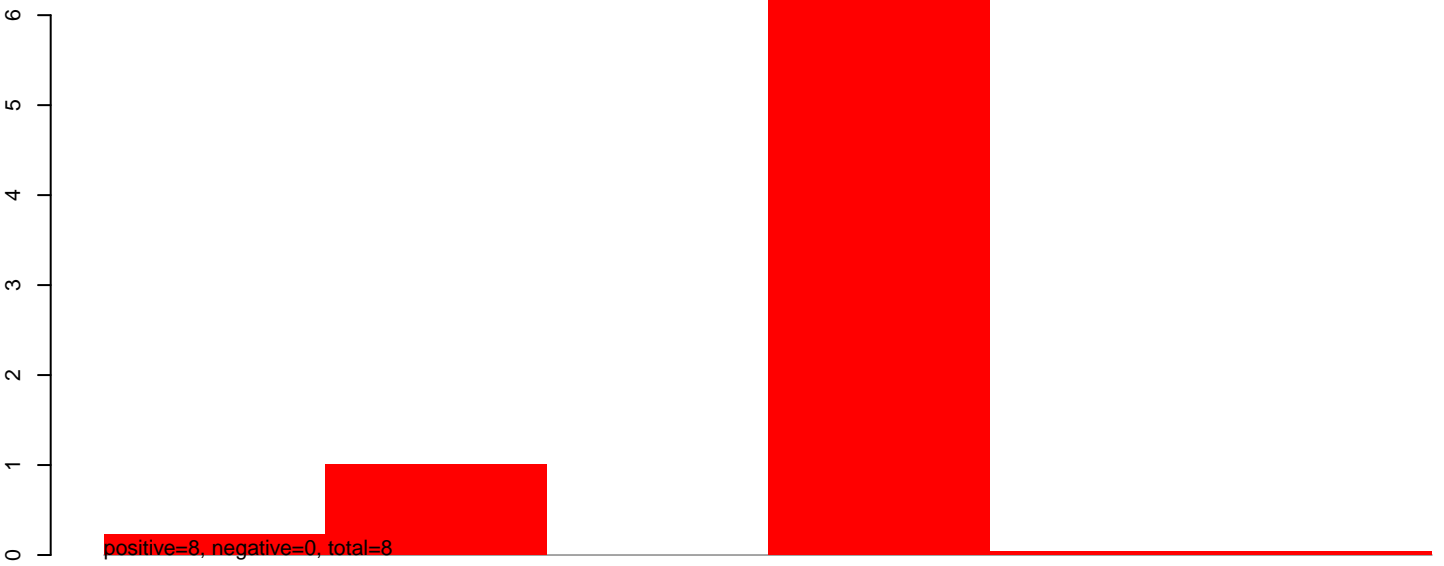


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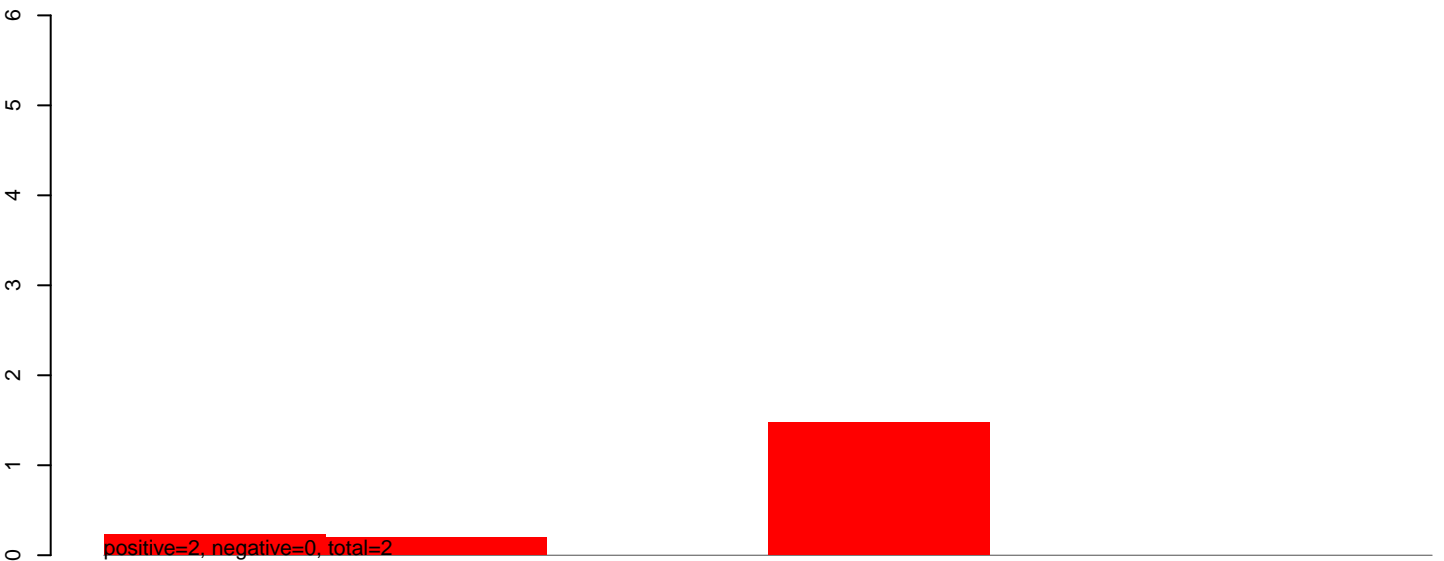


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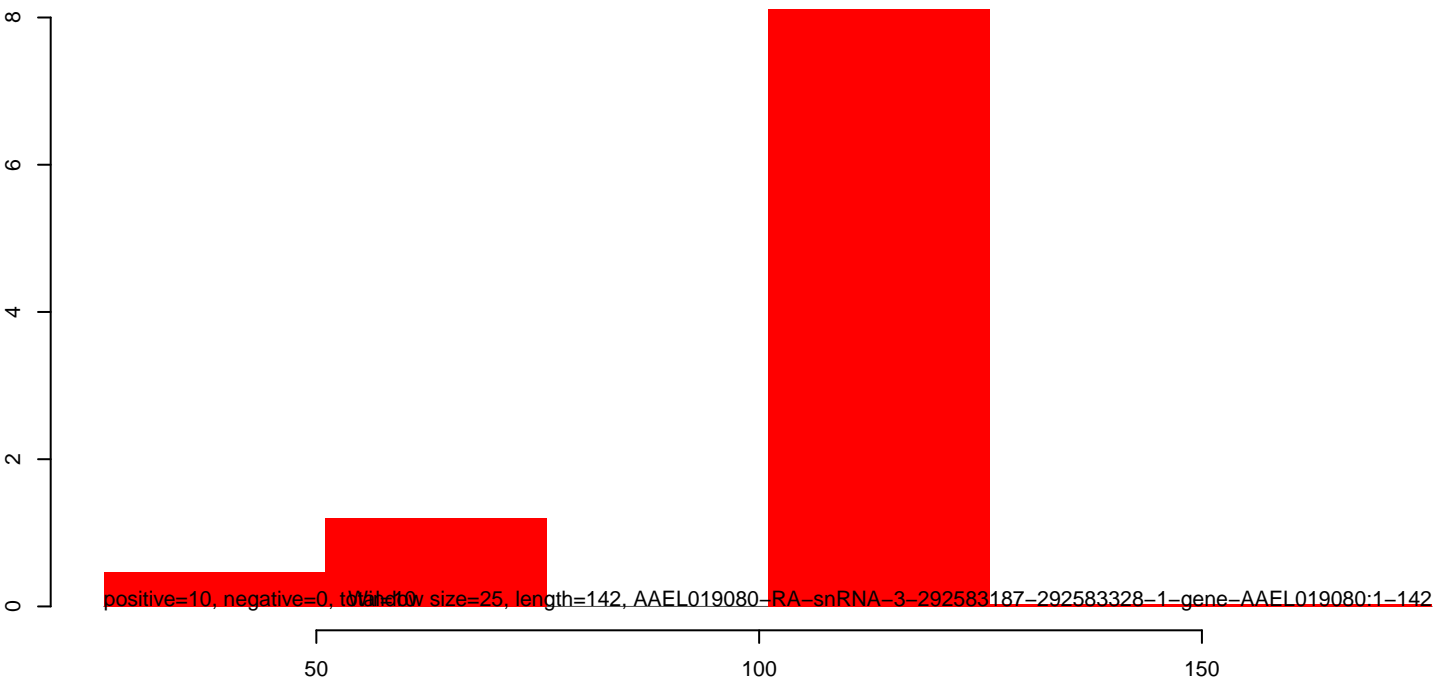
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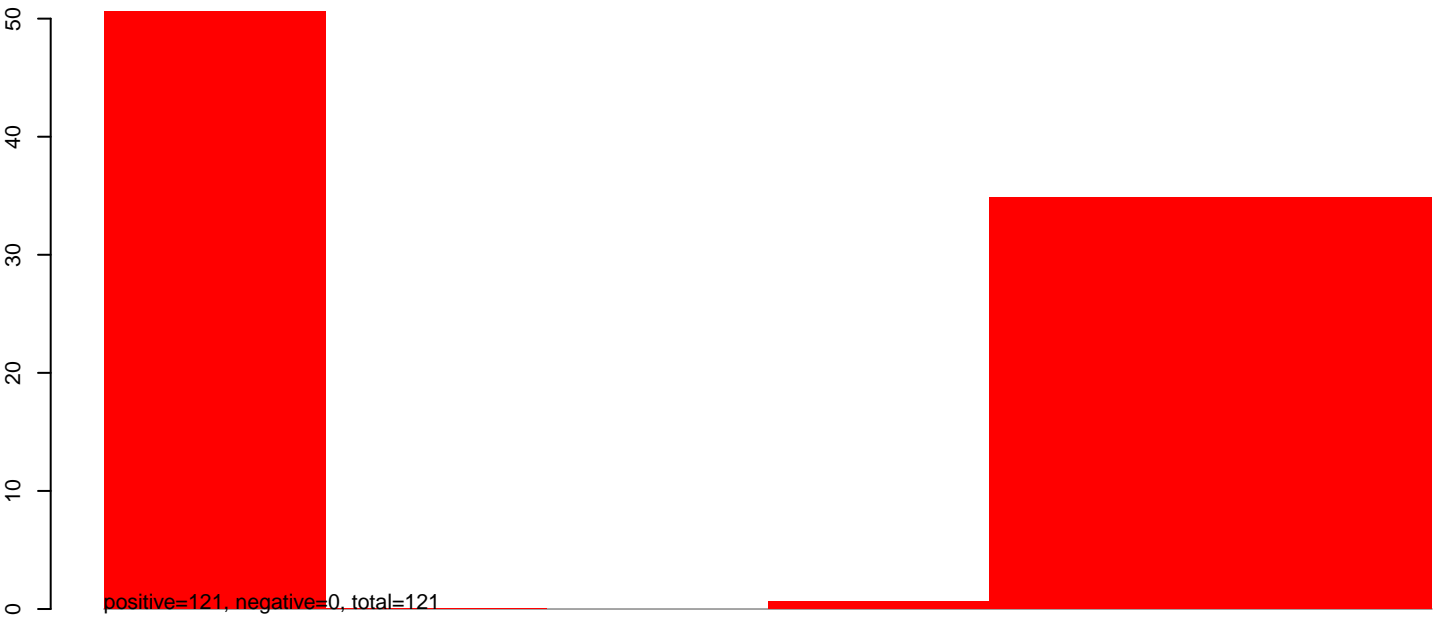
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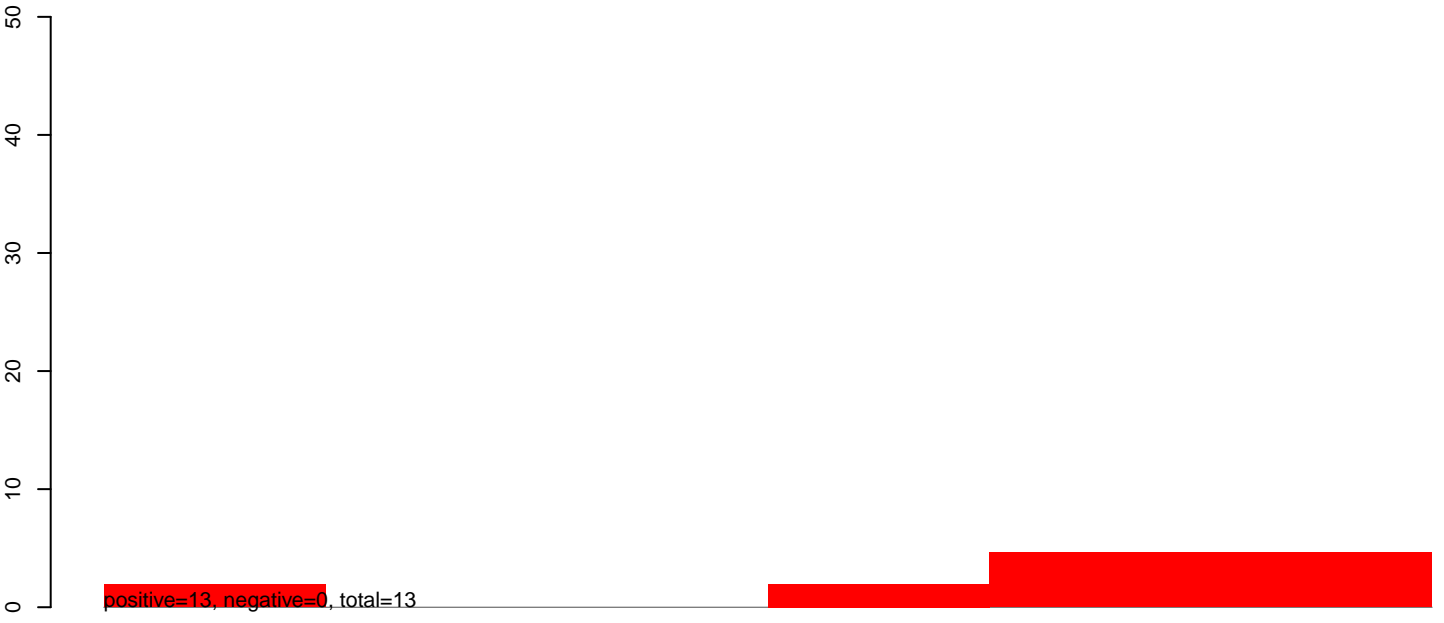
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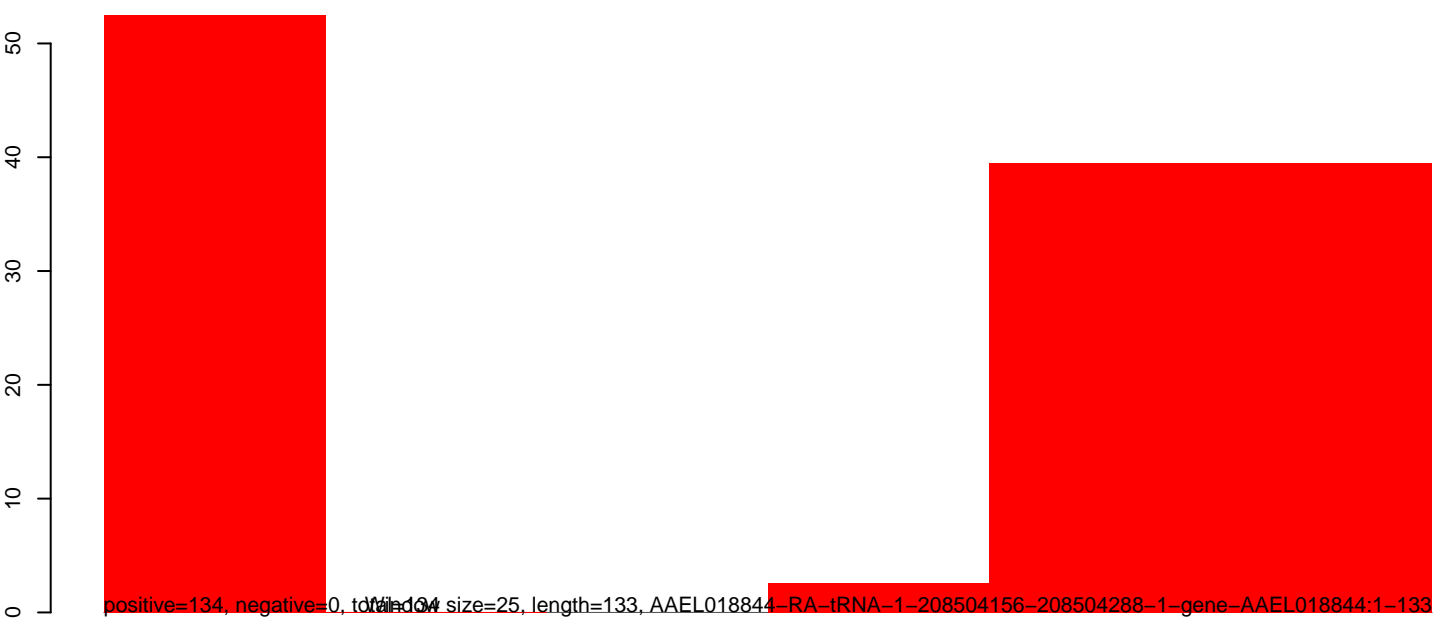
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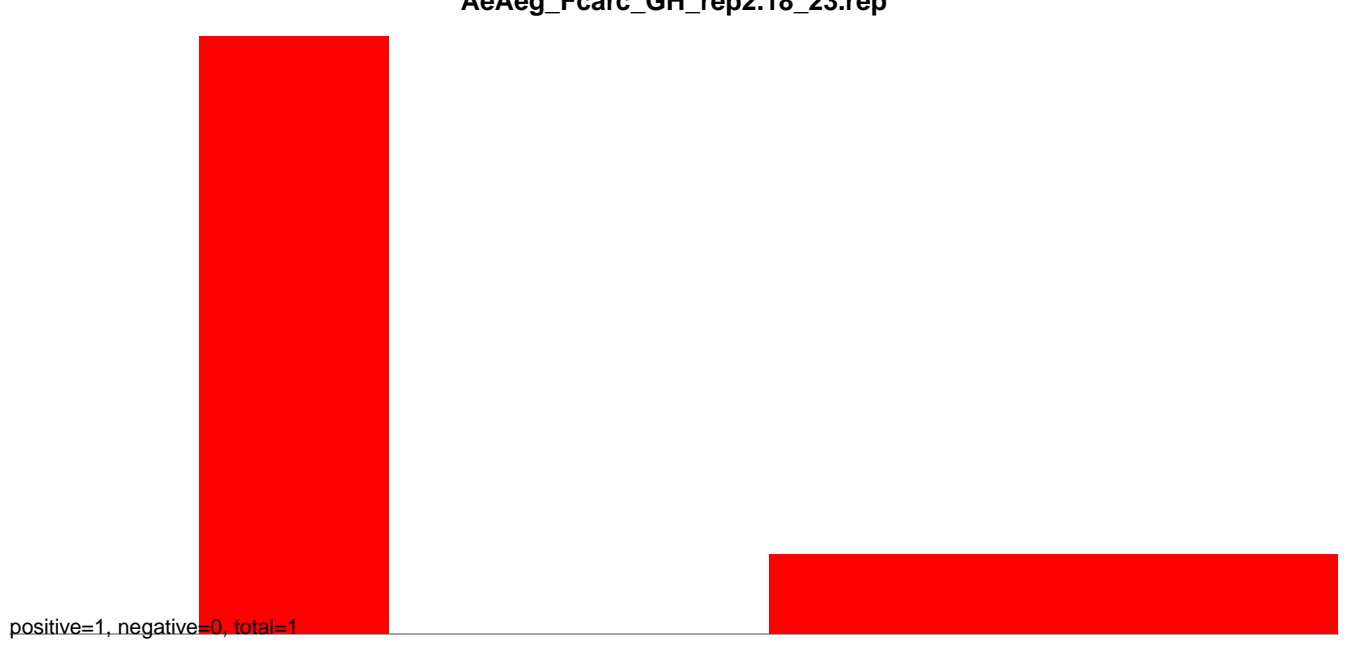
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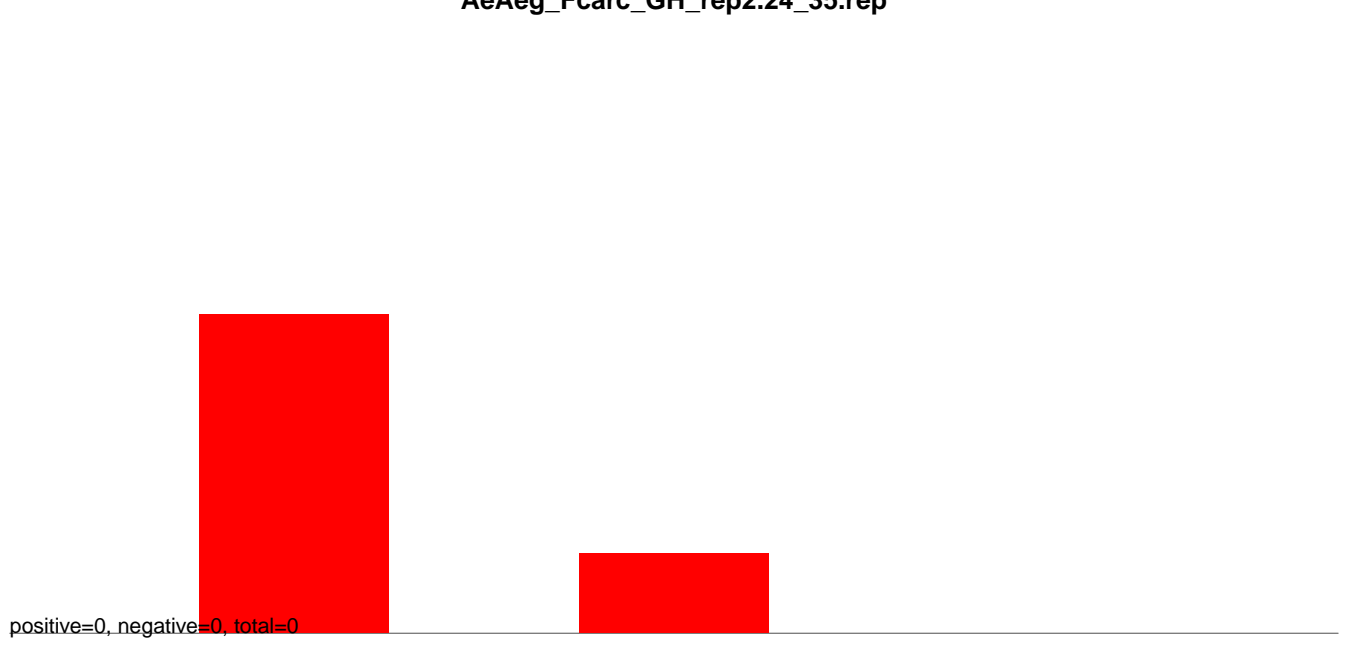
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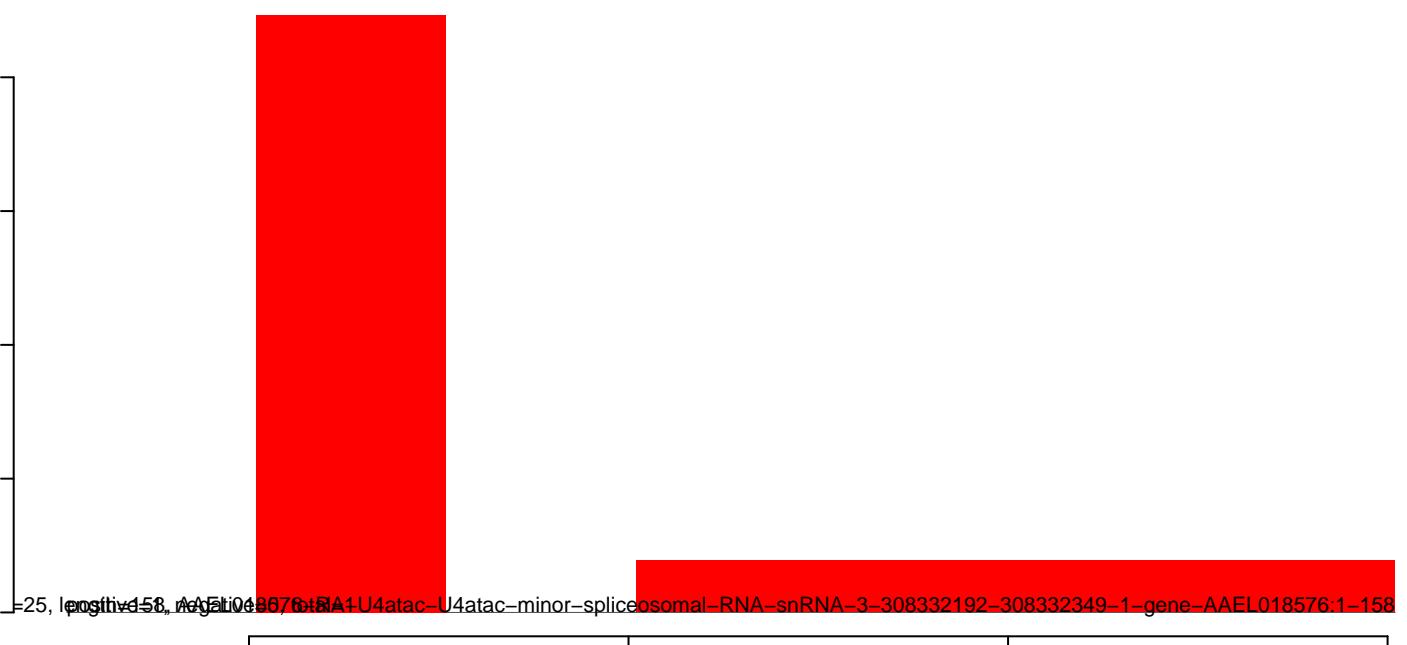
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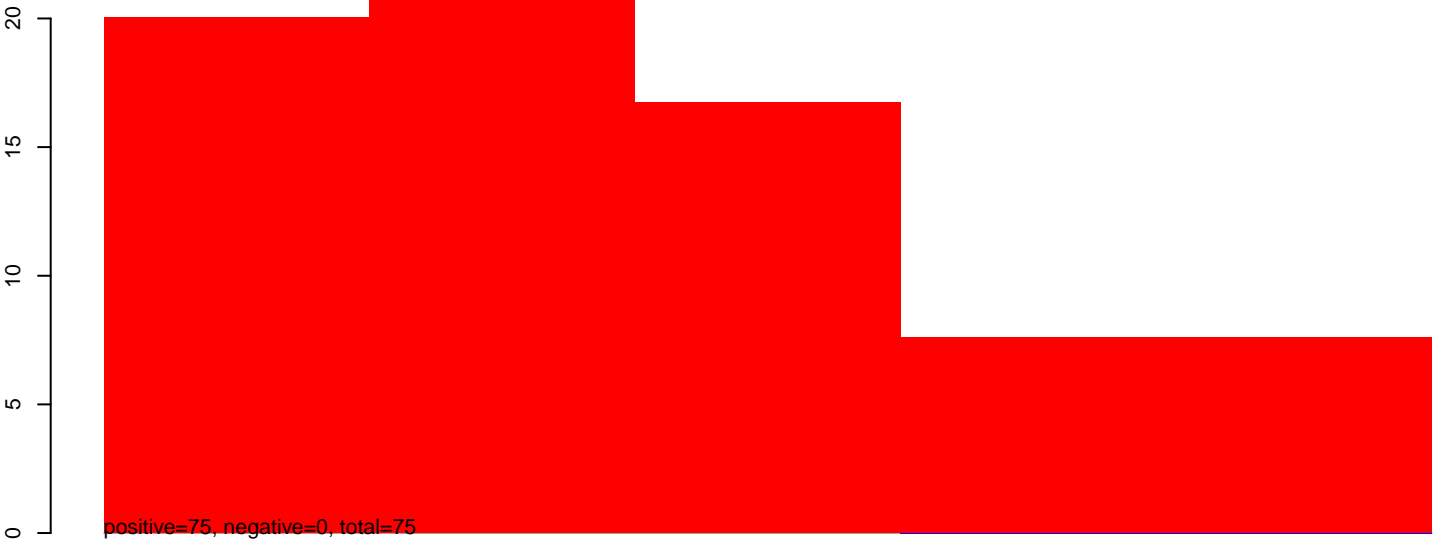
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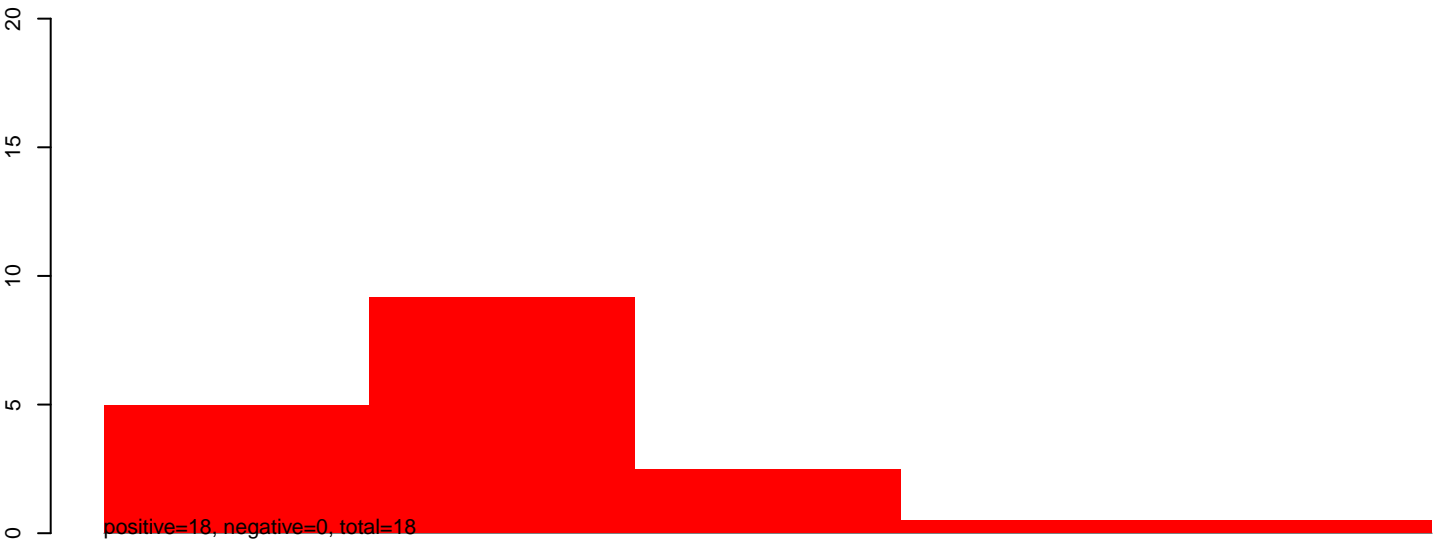
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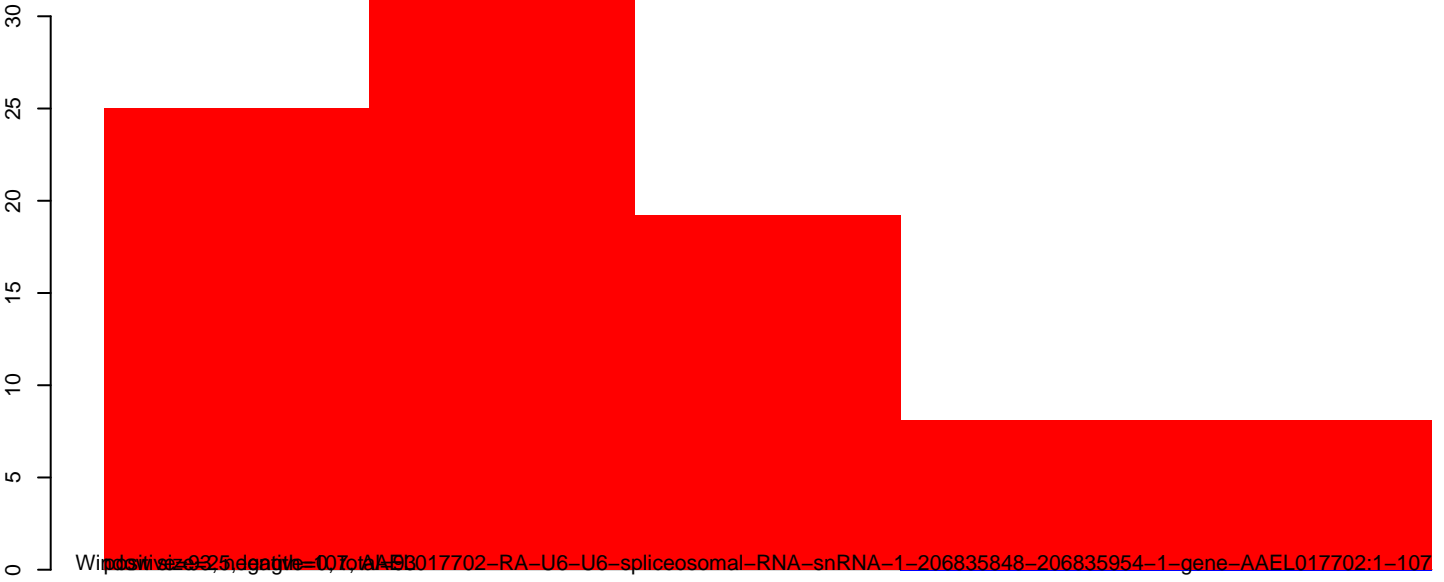
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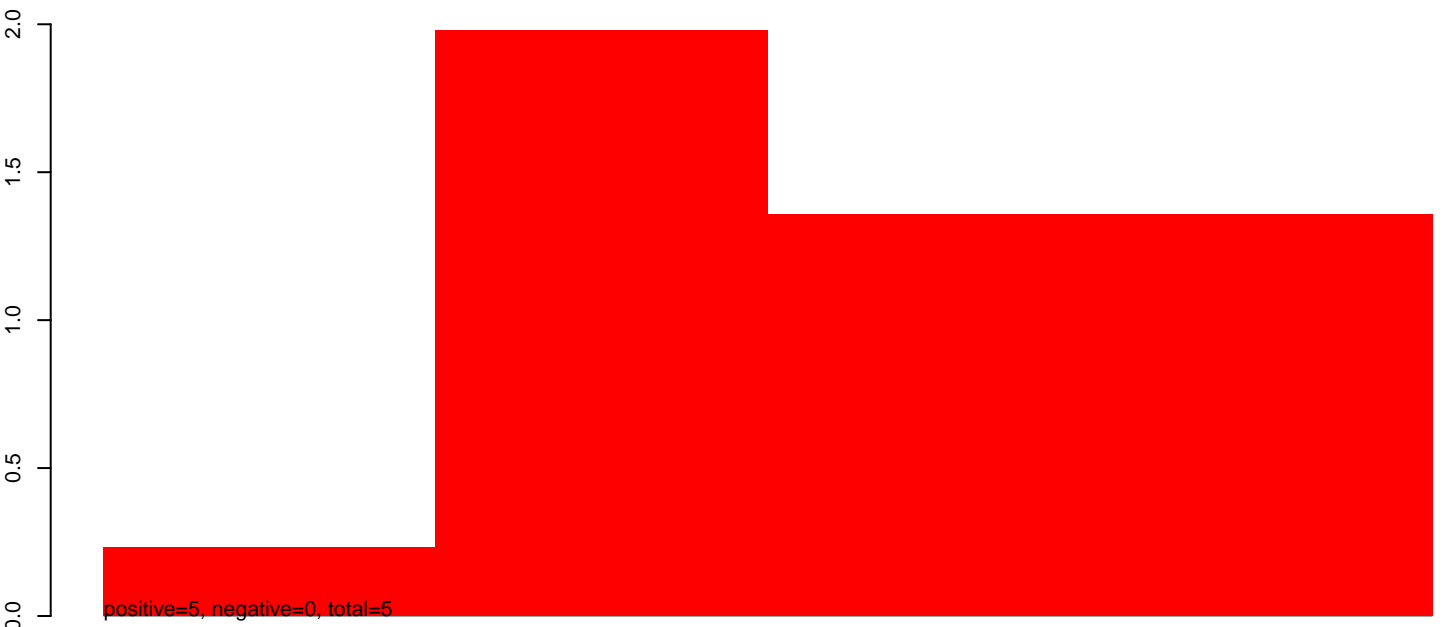


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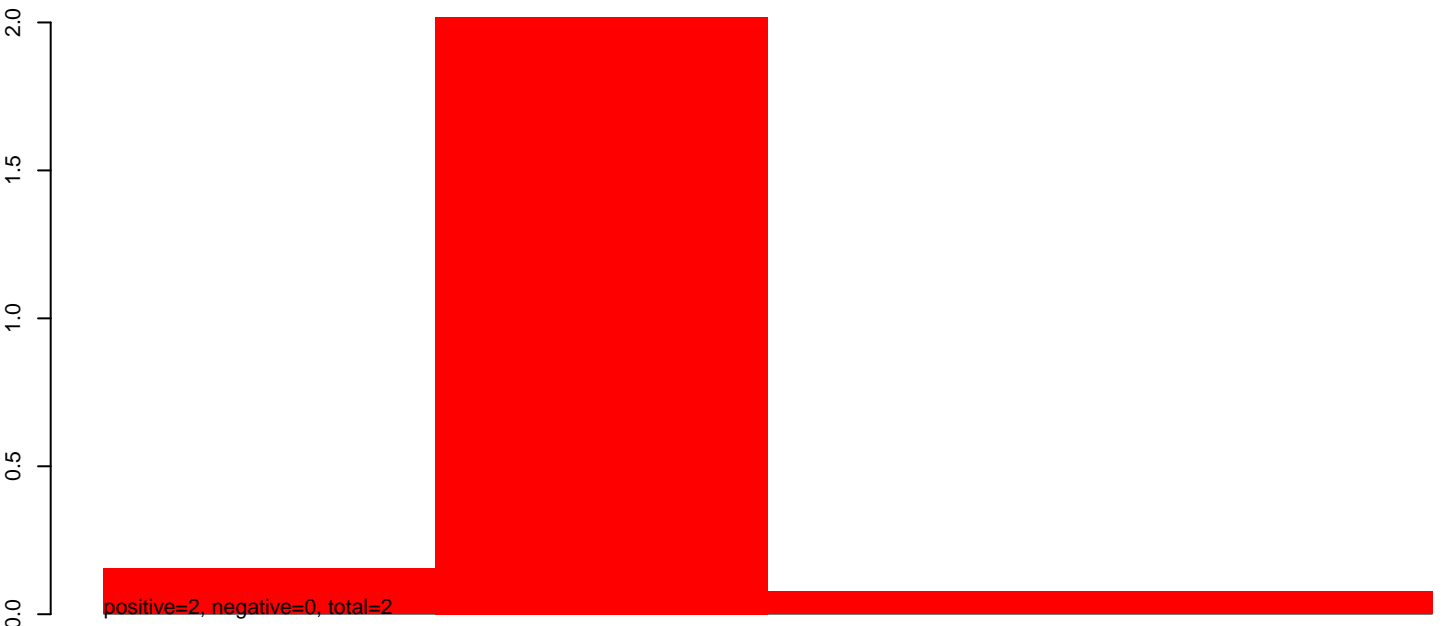


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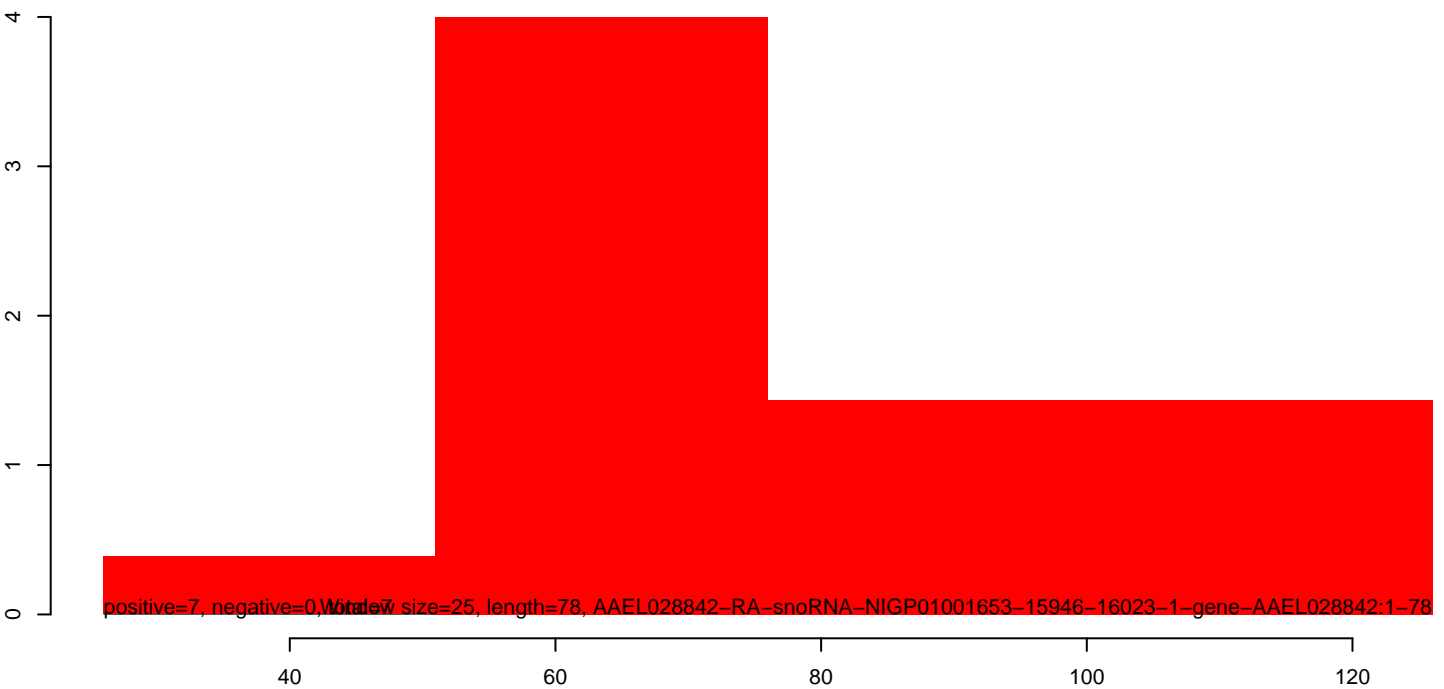
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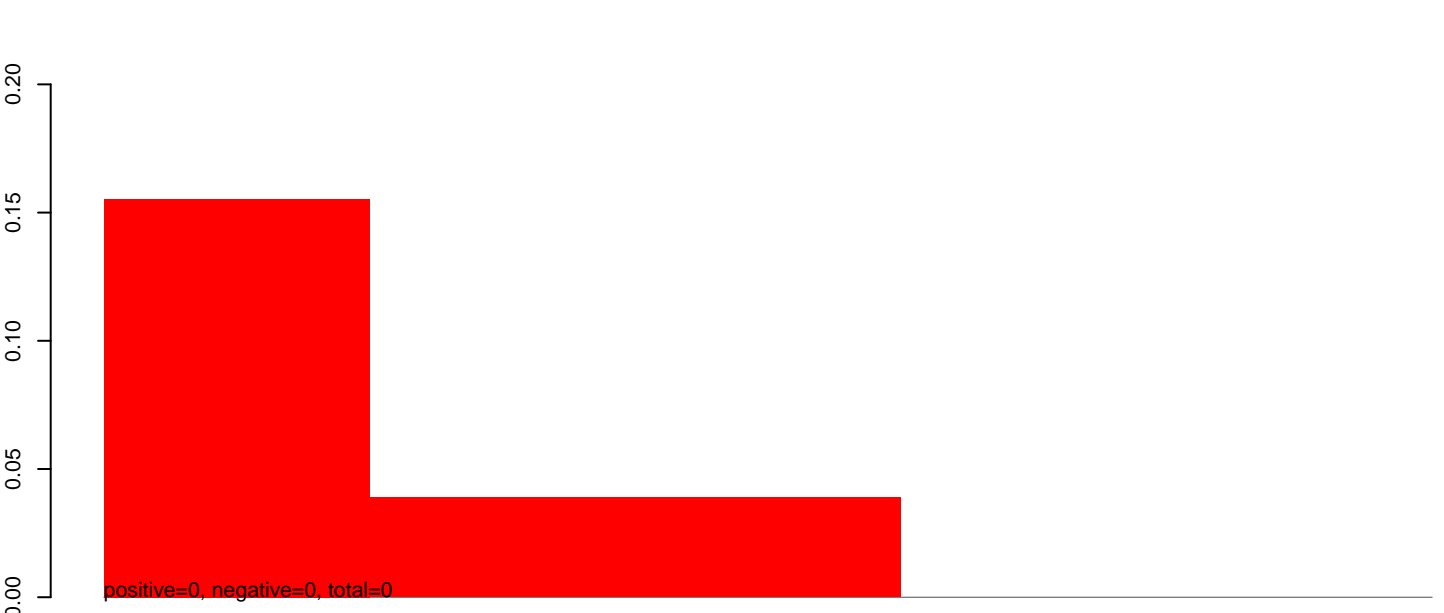
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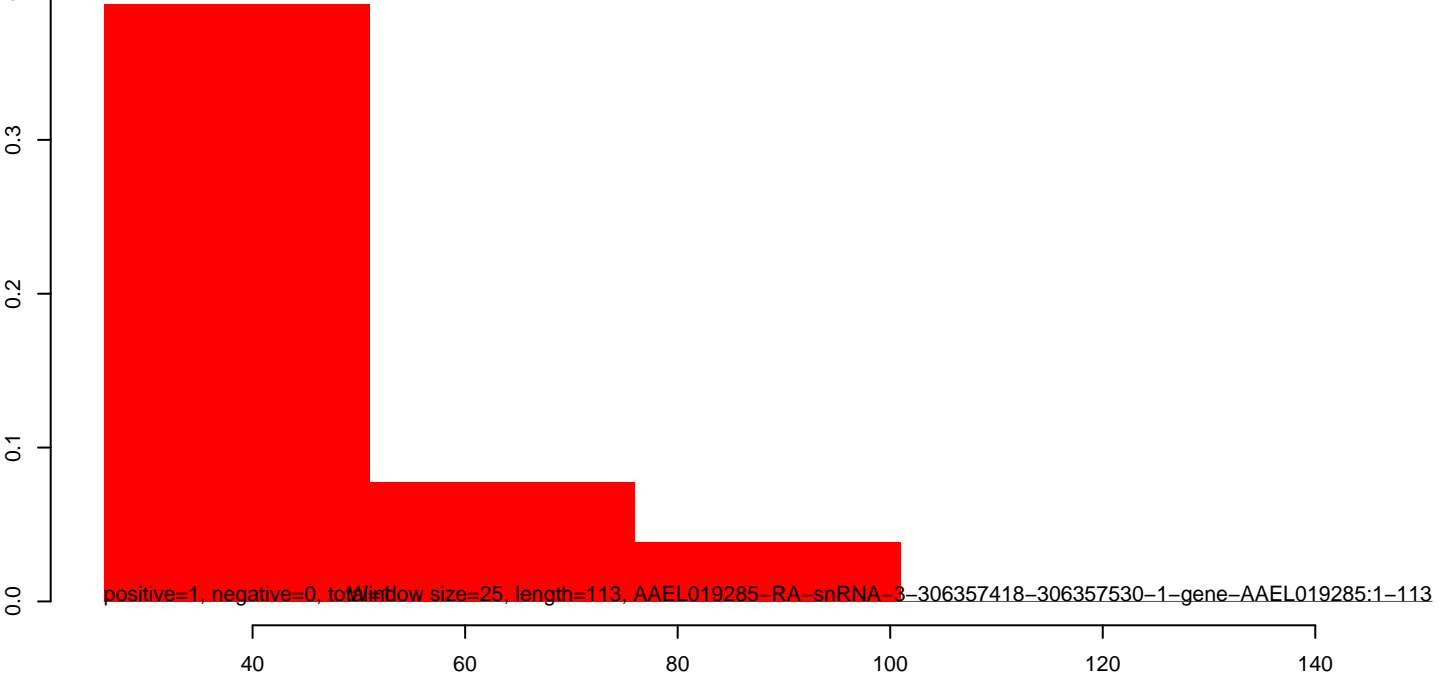
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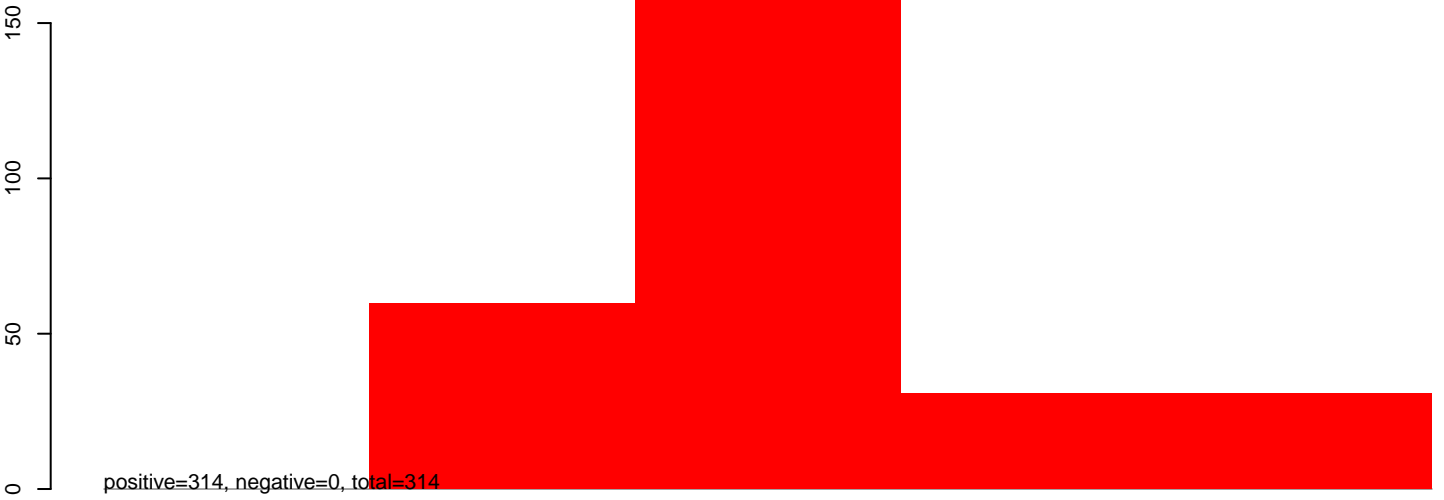
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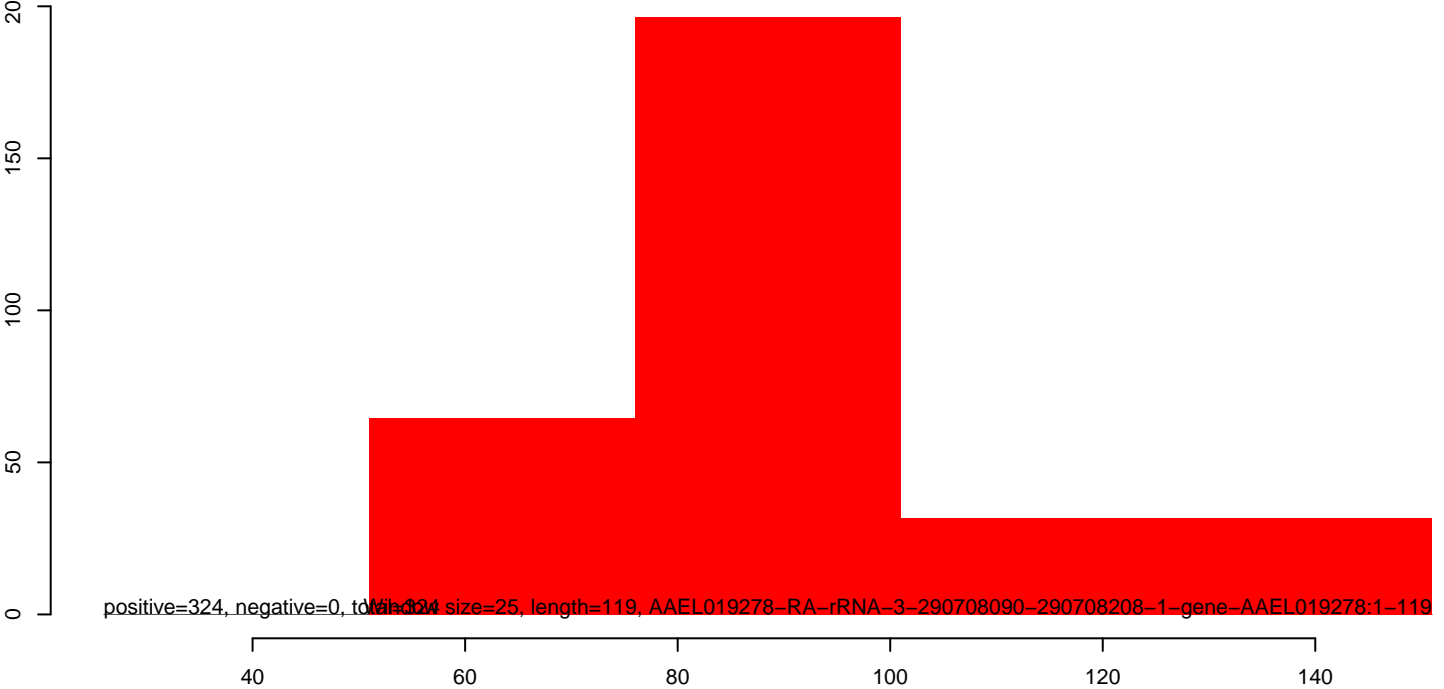
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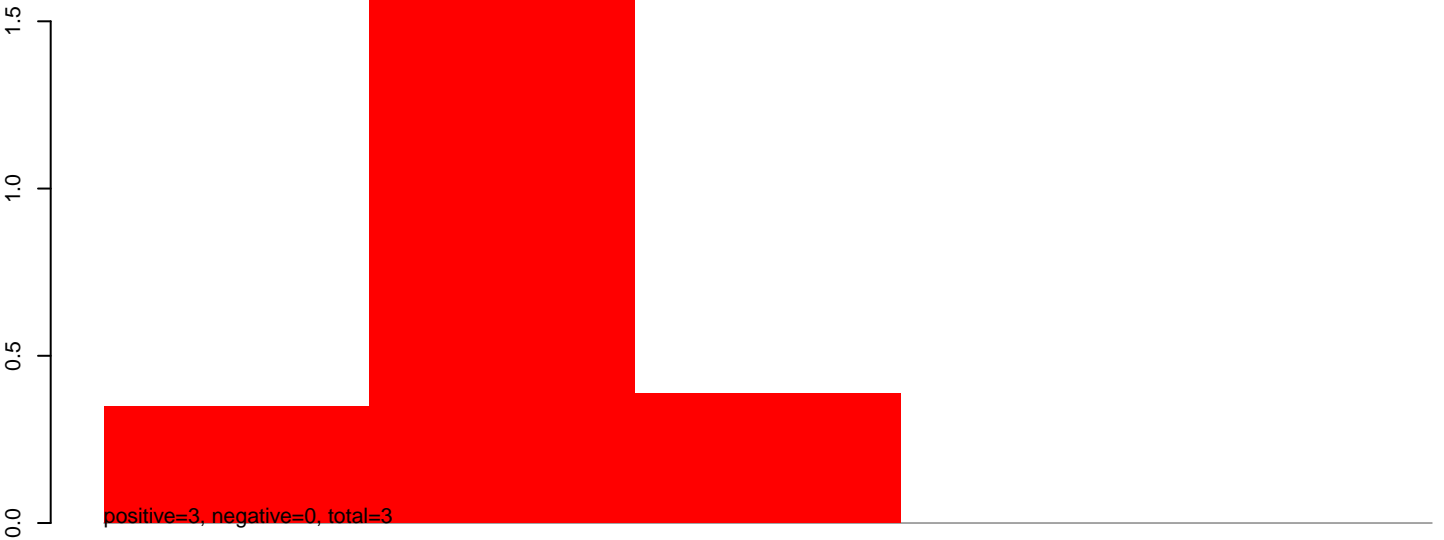
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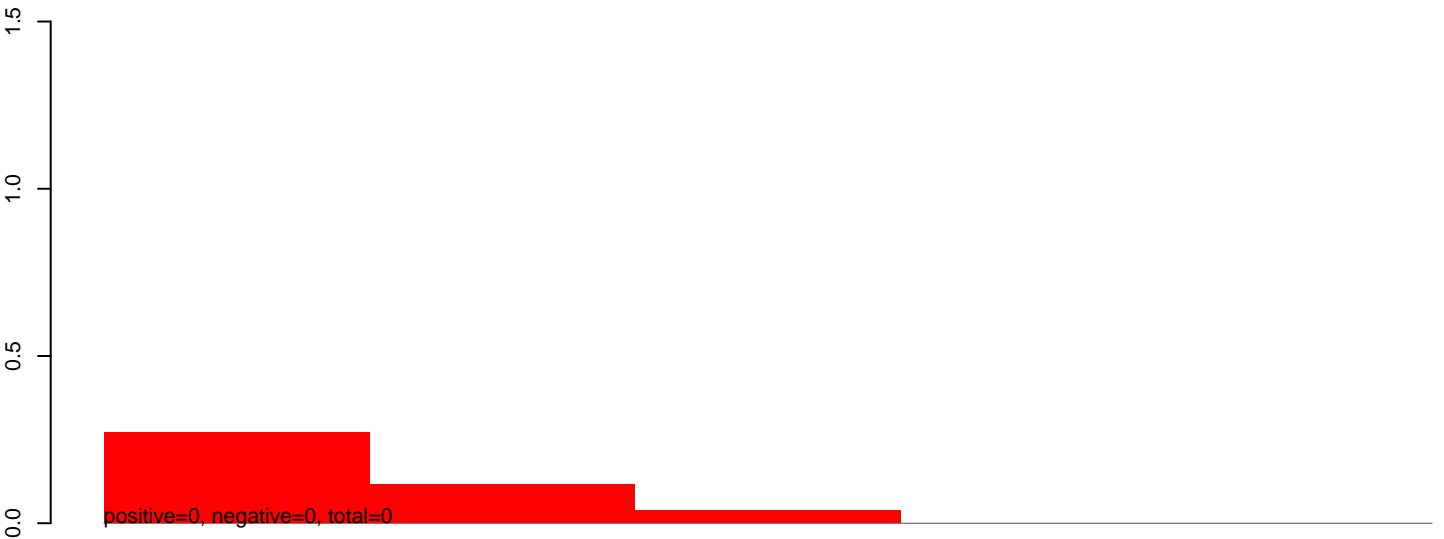
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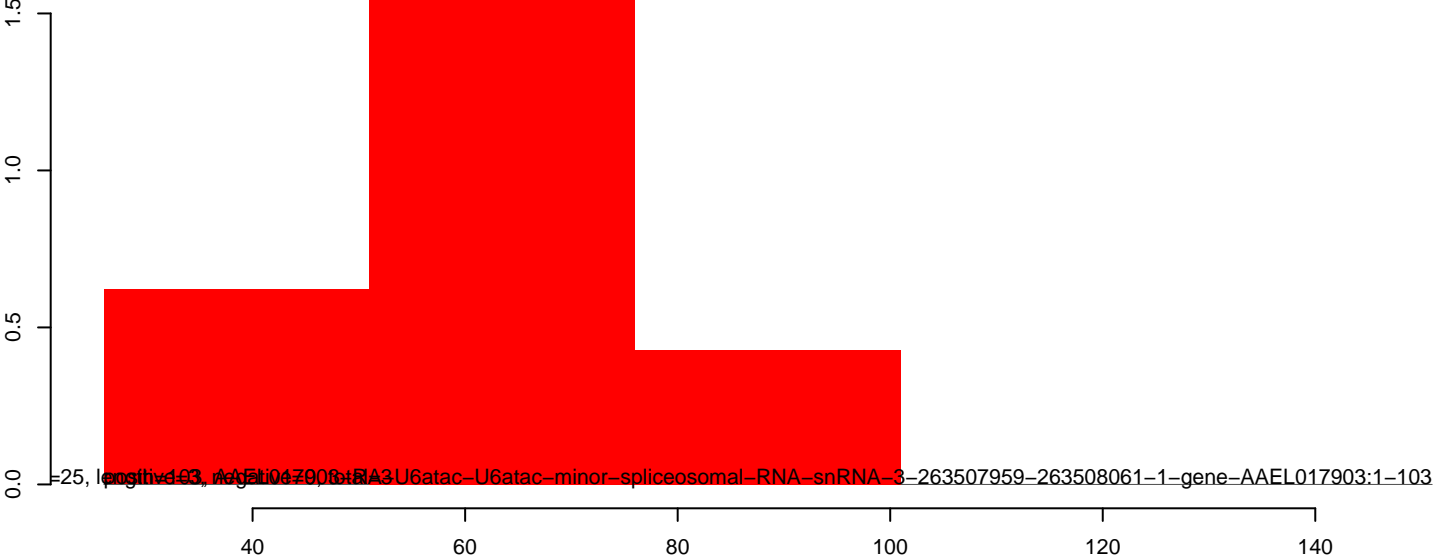
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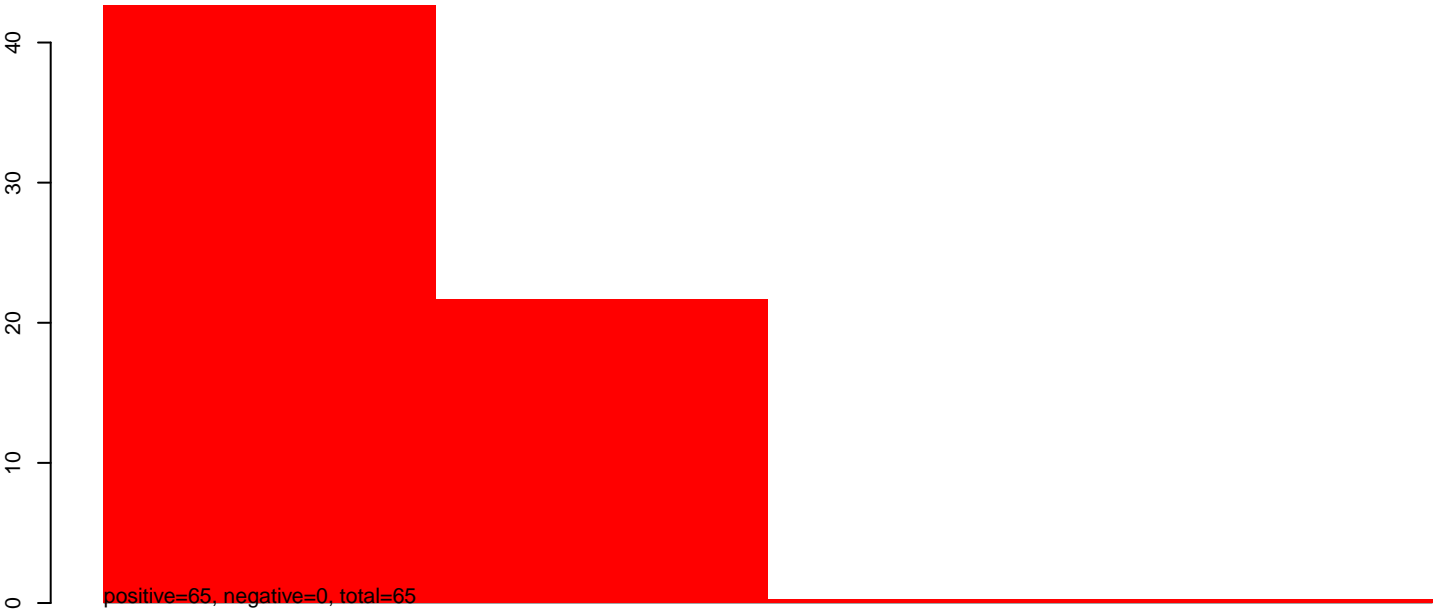


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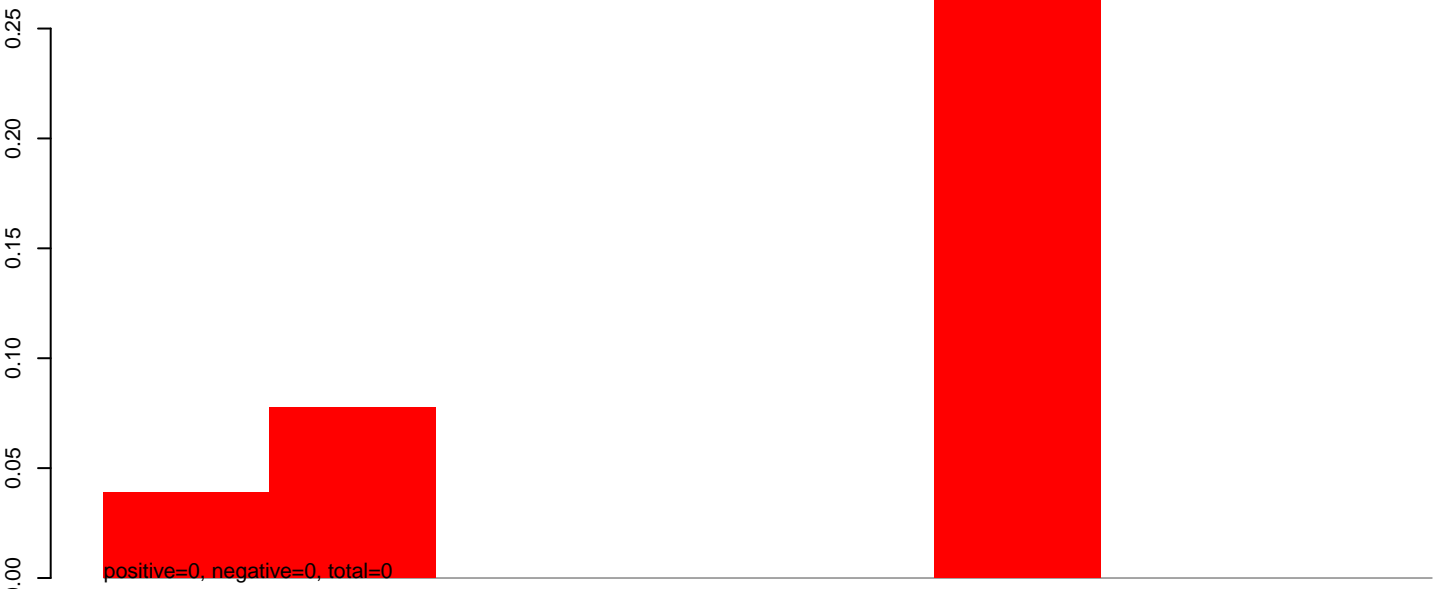
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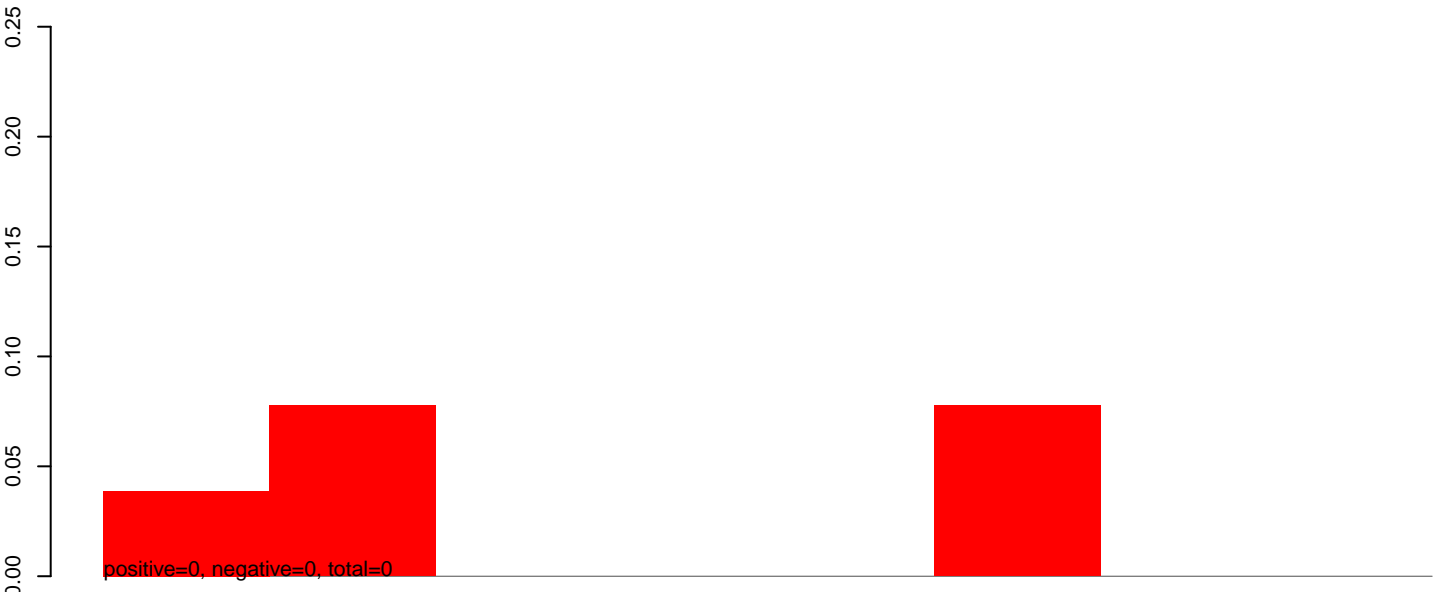
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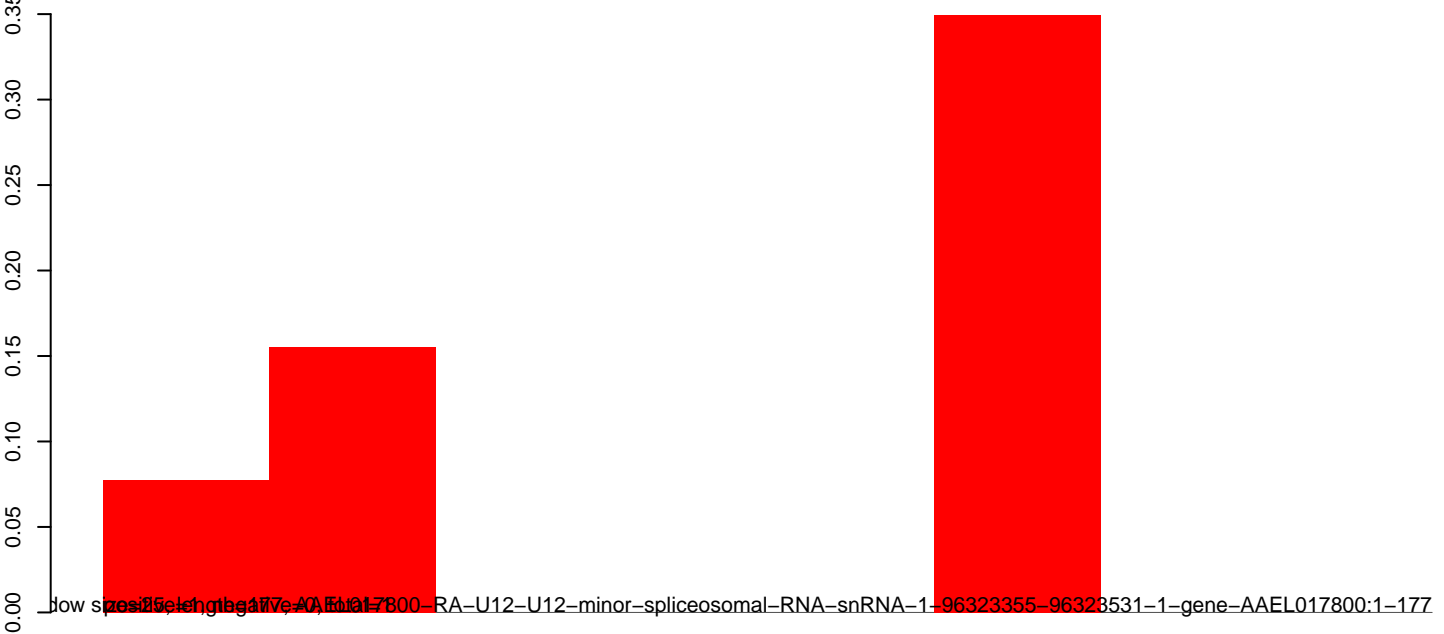
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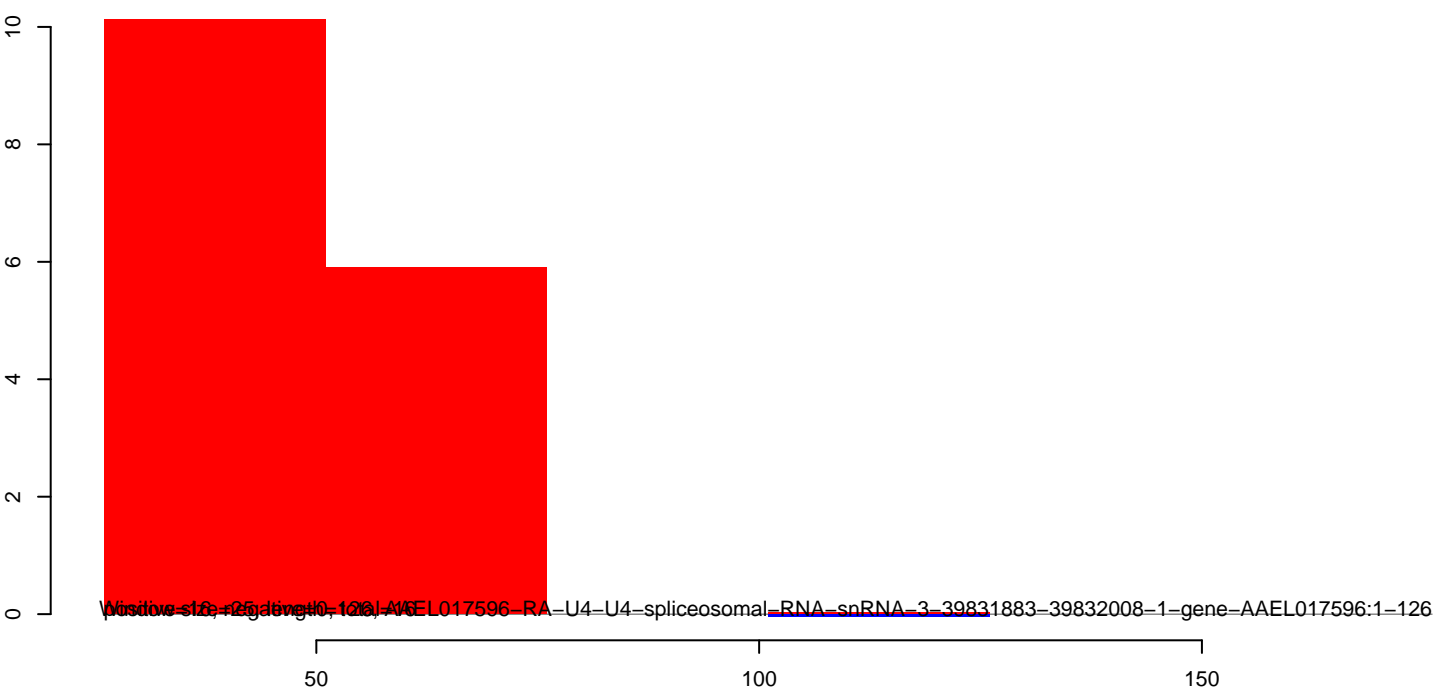
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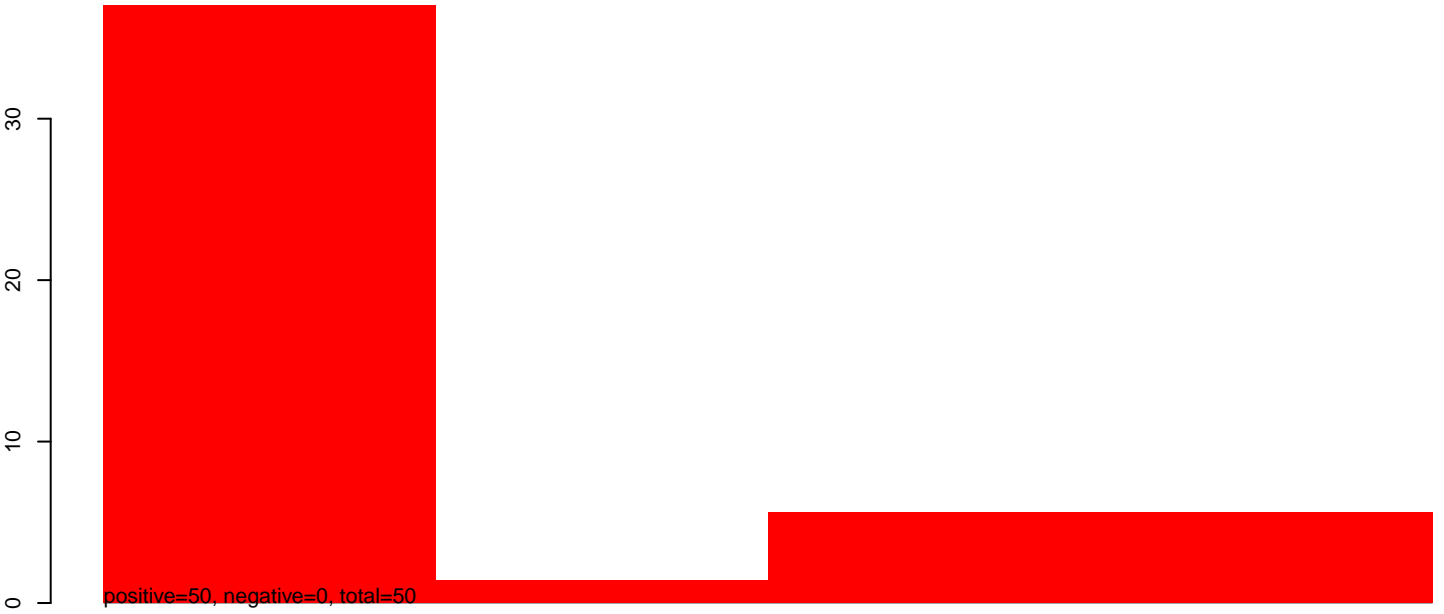
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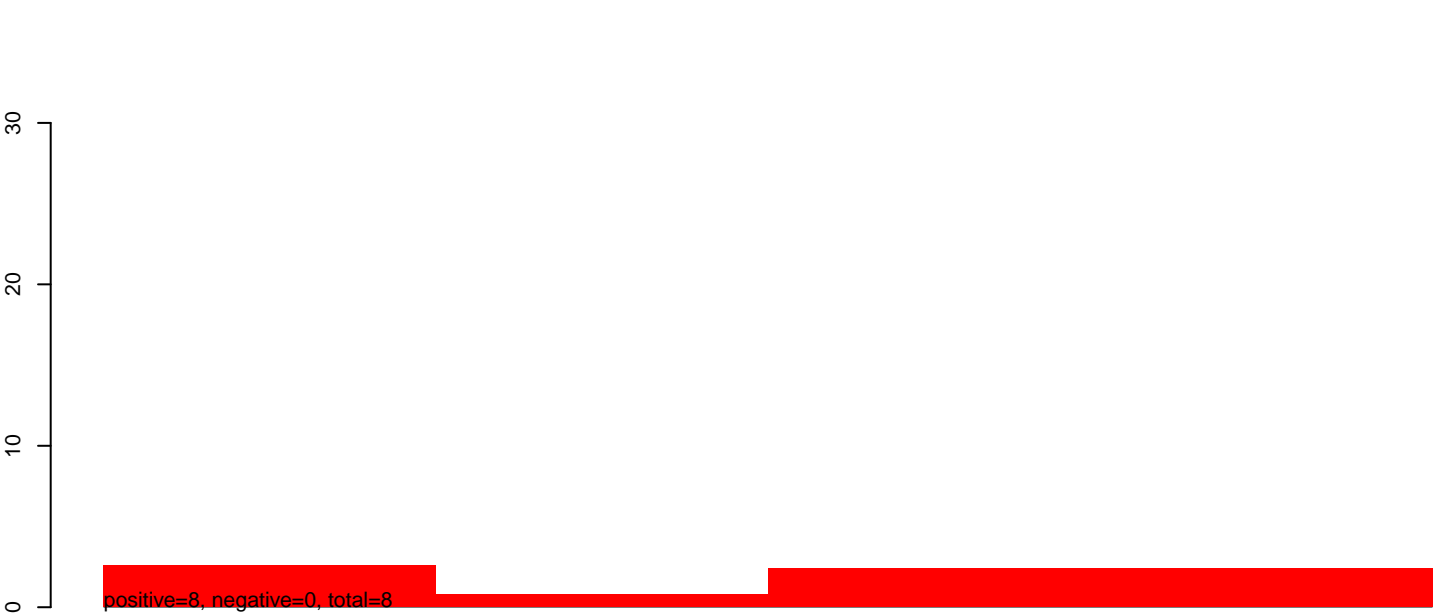
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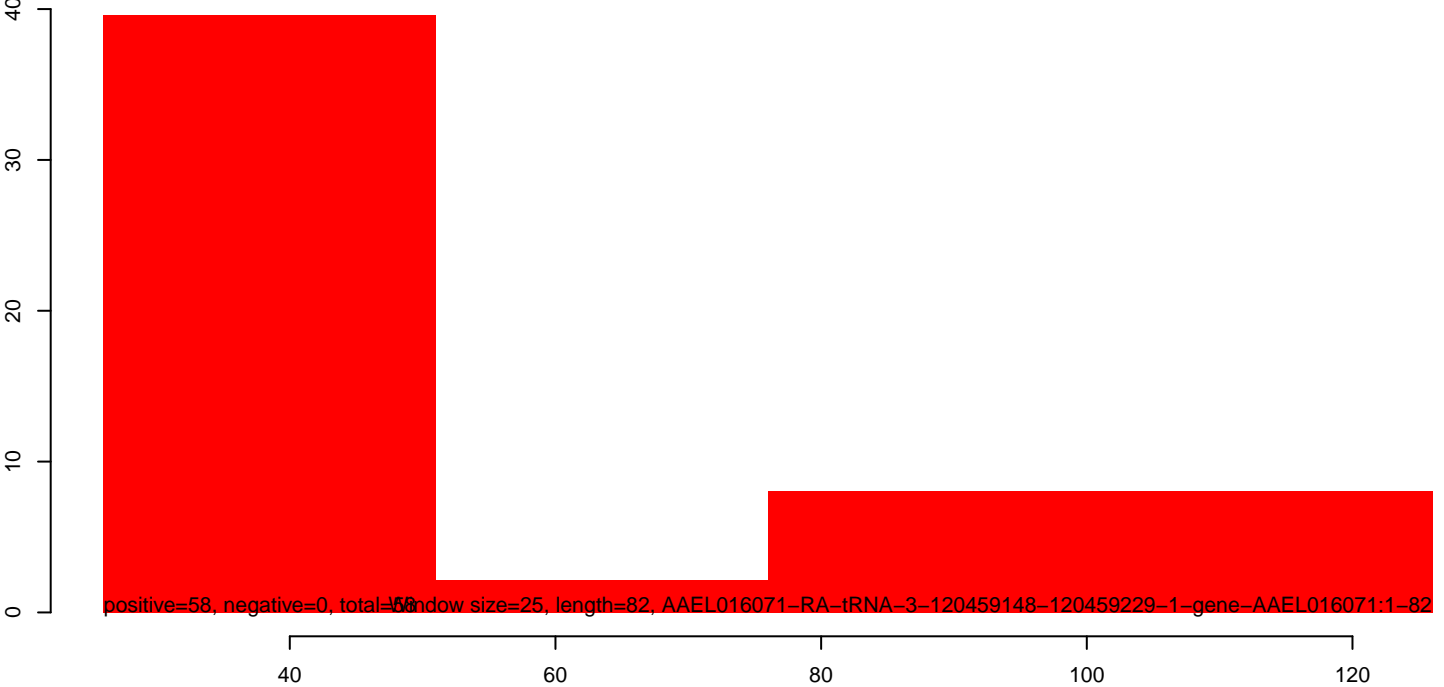
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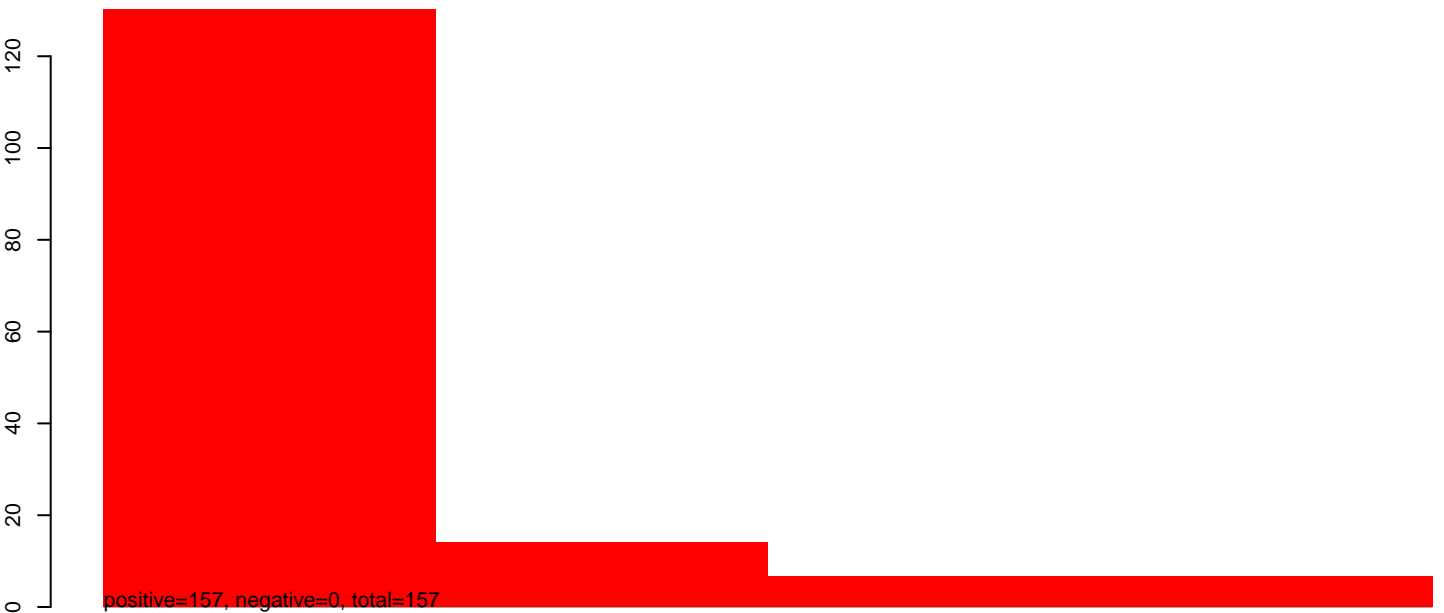
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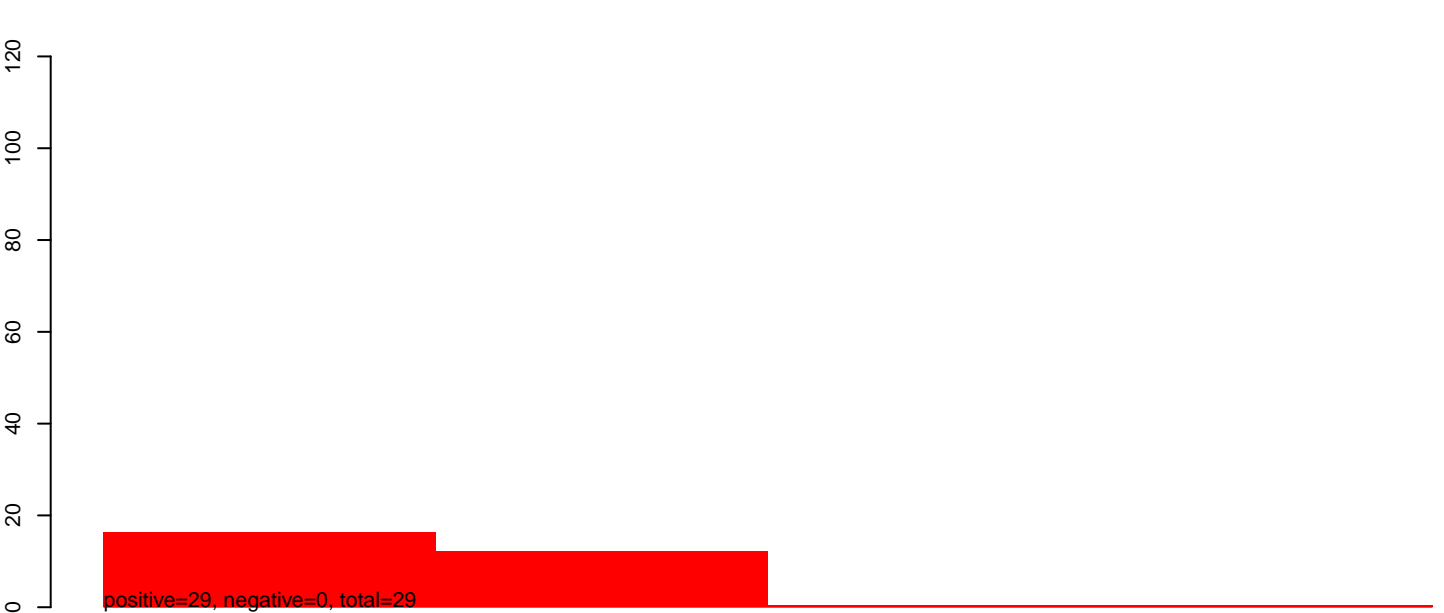
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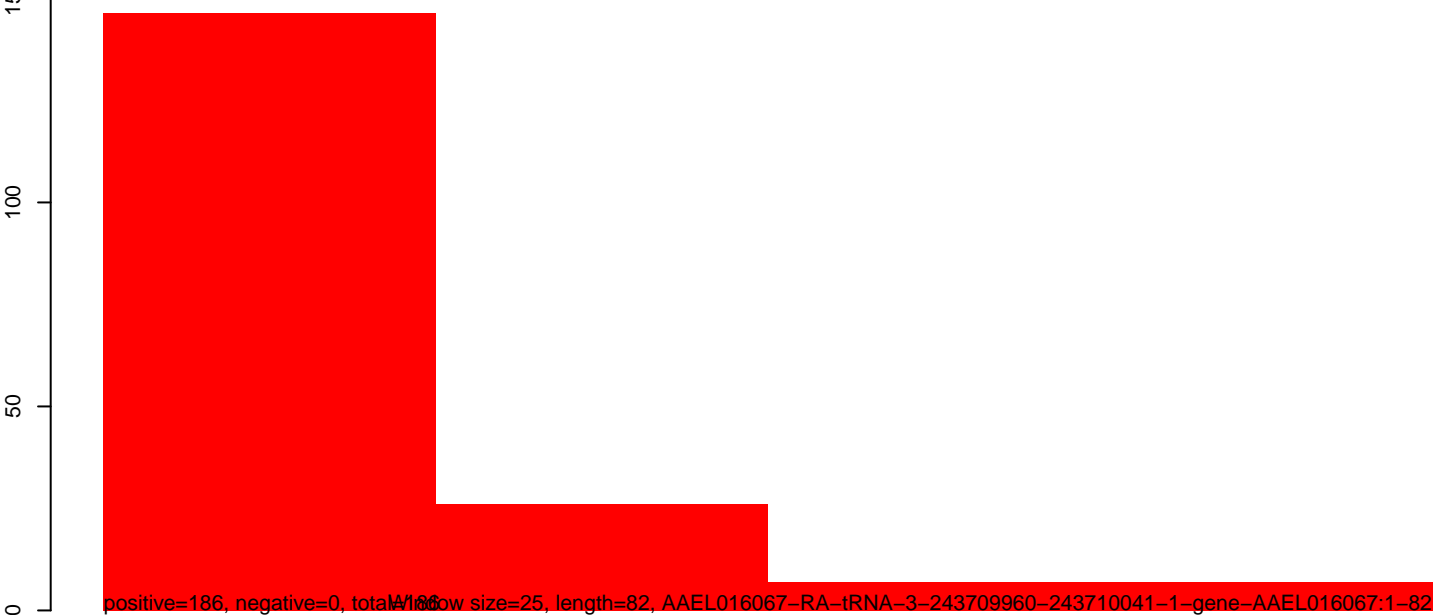
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AeAeg_Fcarc_GH_rep2.24_35.rep

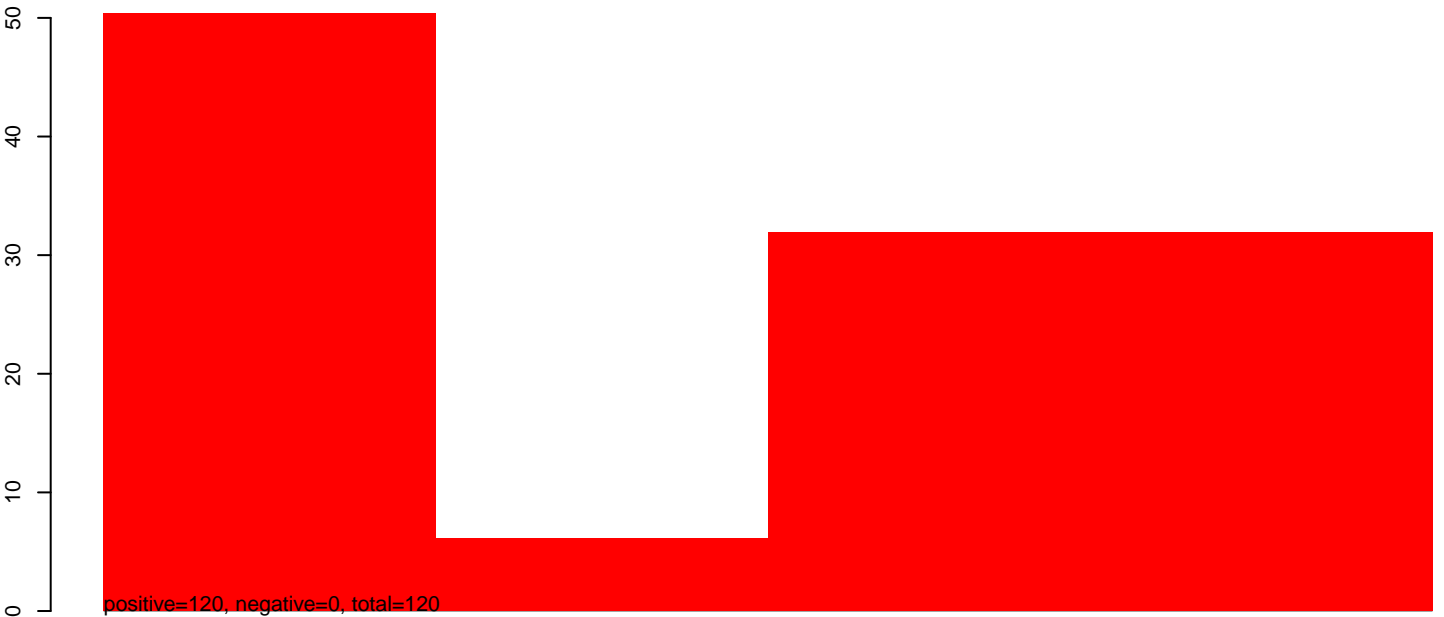


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40 60 80 100 120

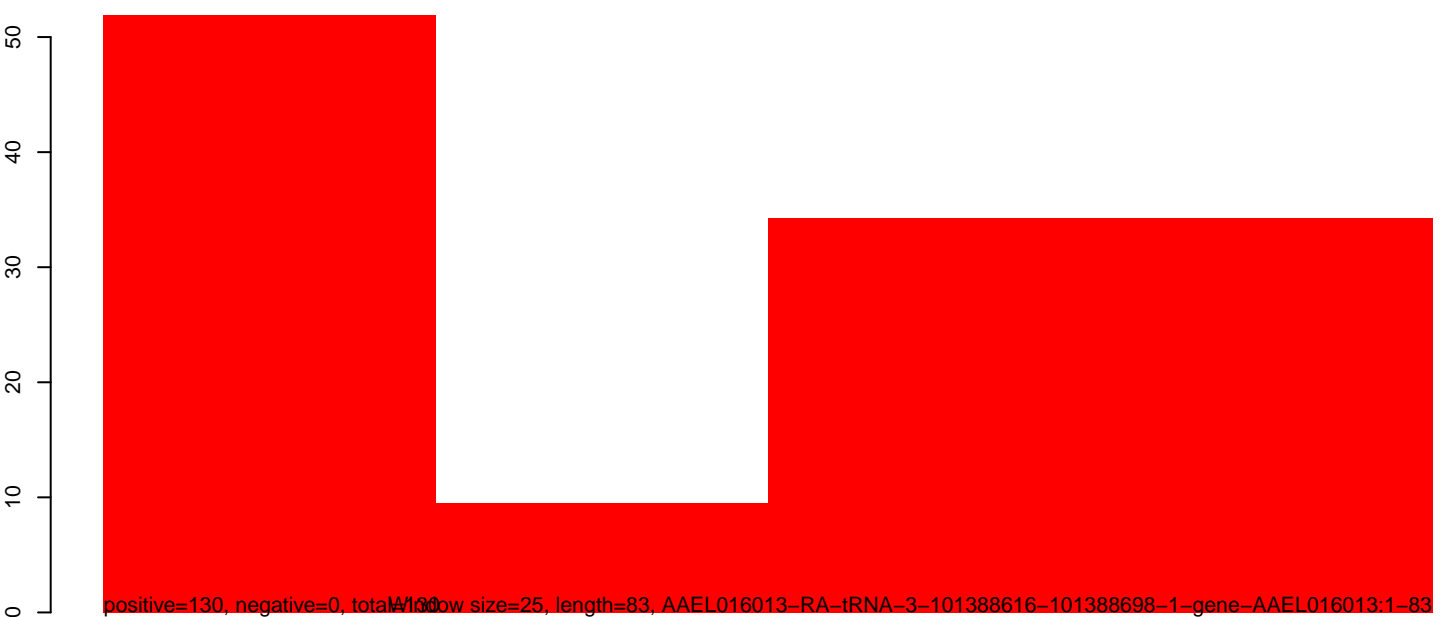
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AeAeg_Fcarc_GH_rep2.24_35.rep

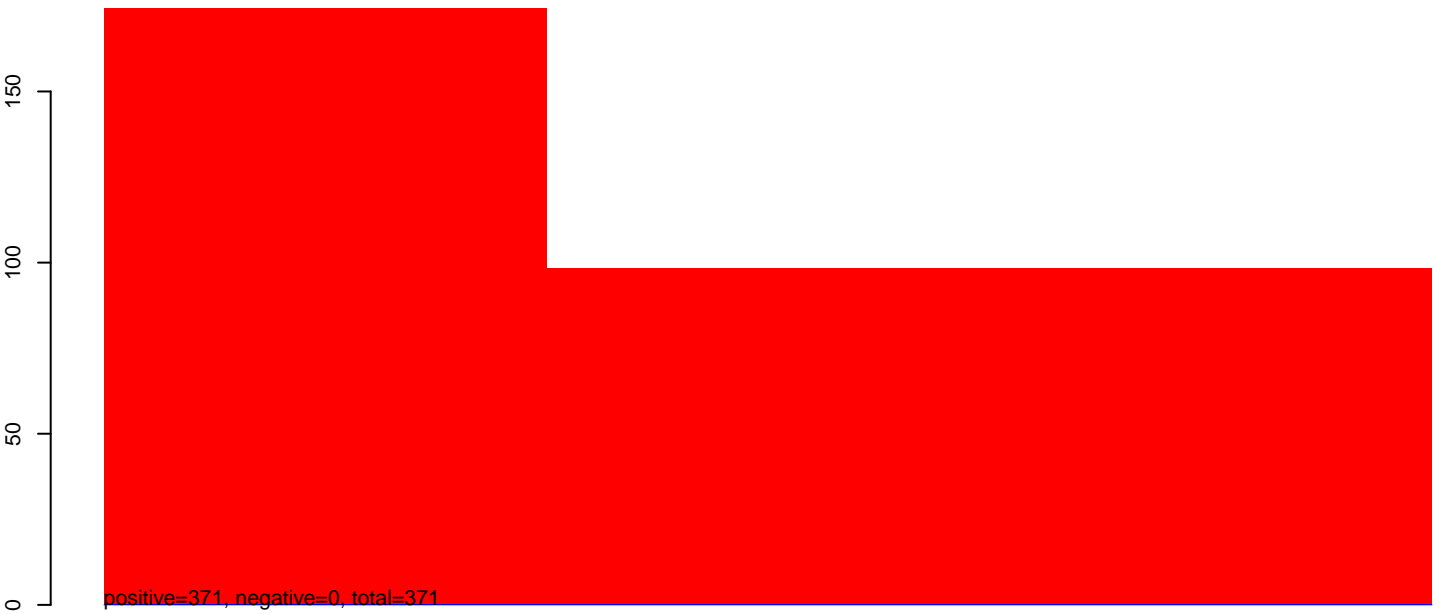


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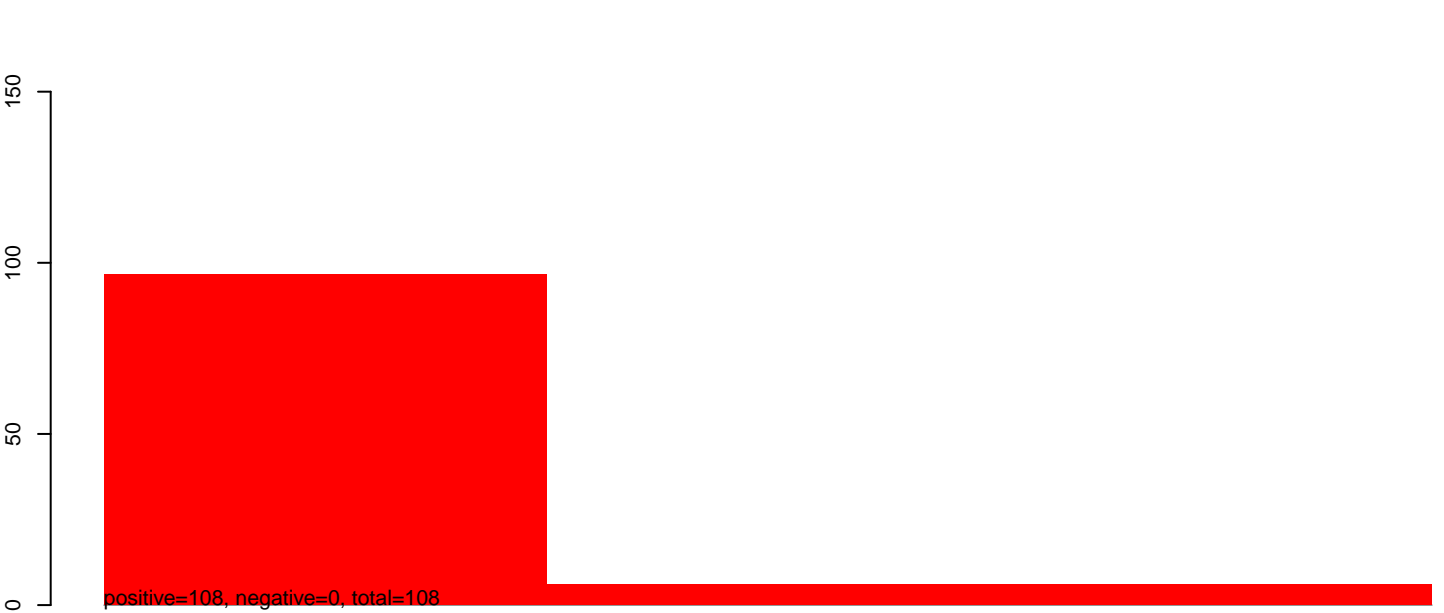


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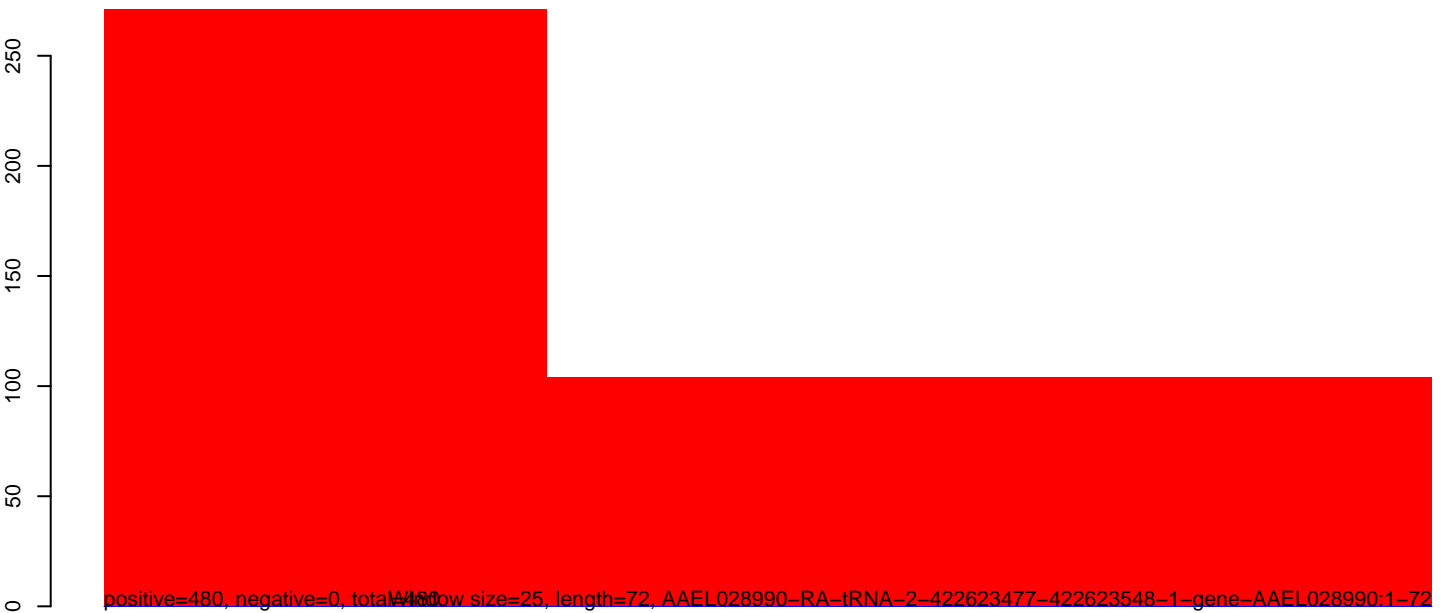
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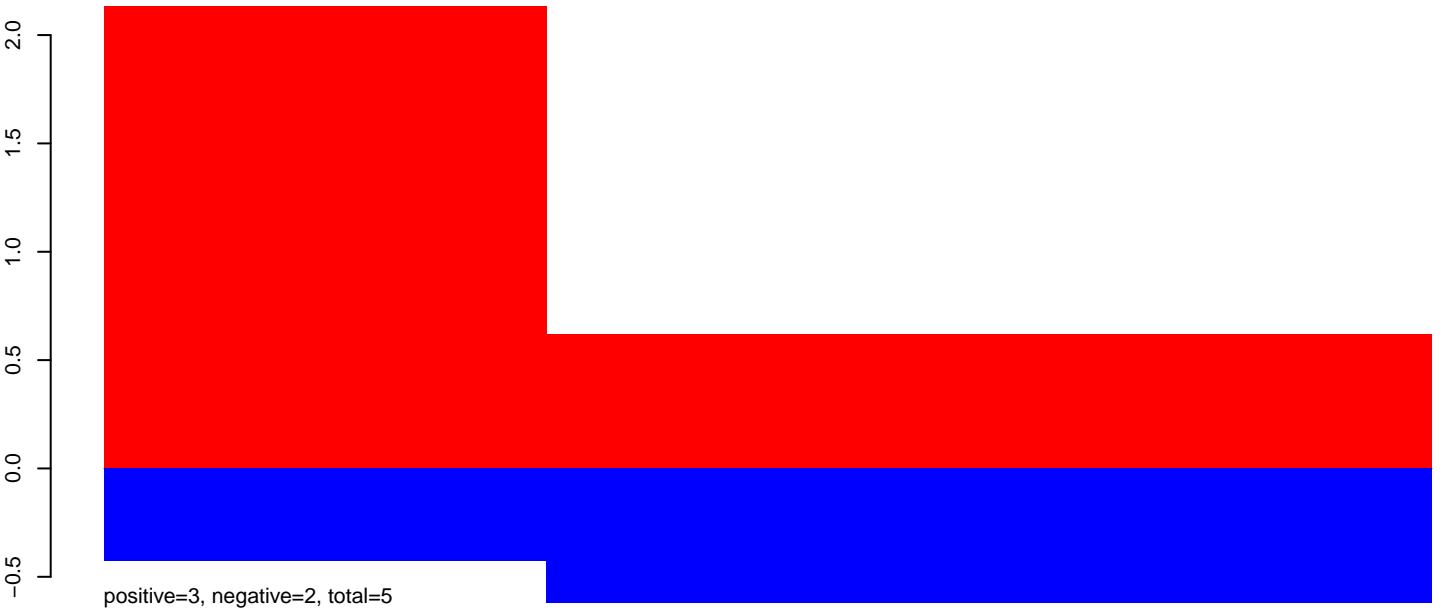


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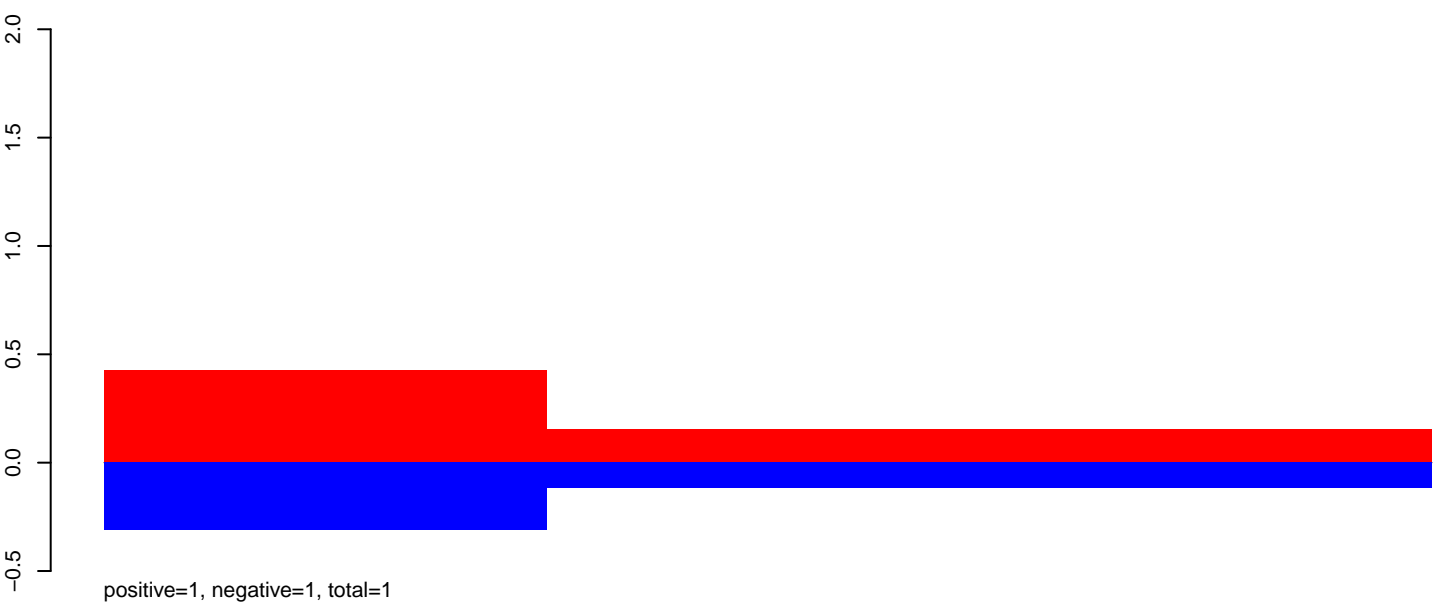


40 60 80 100
AAEL028990-RA-tRNA-2-422623477-422623548-1-gene-AAEL028990:1-72
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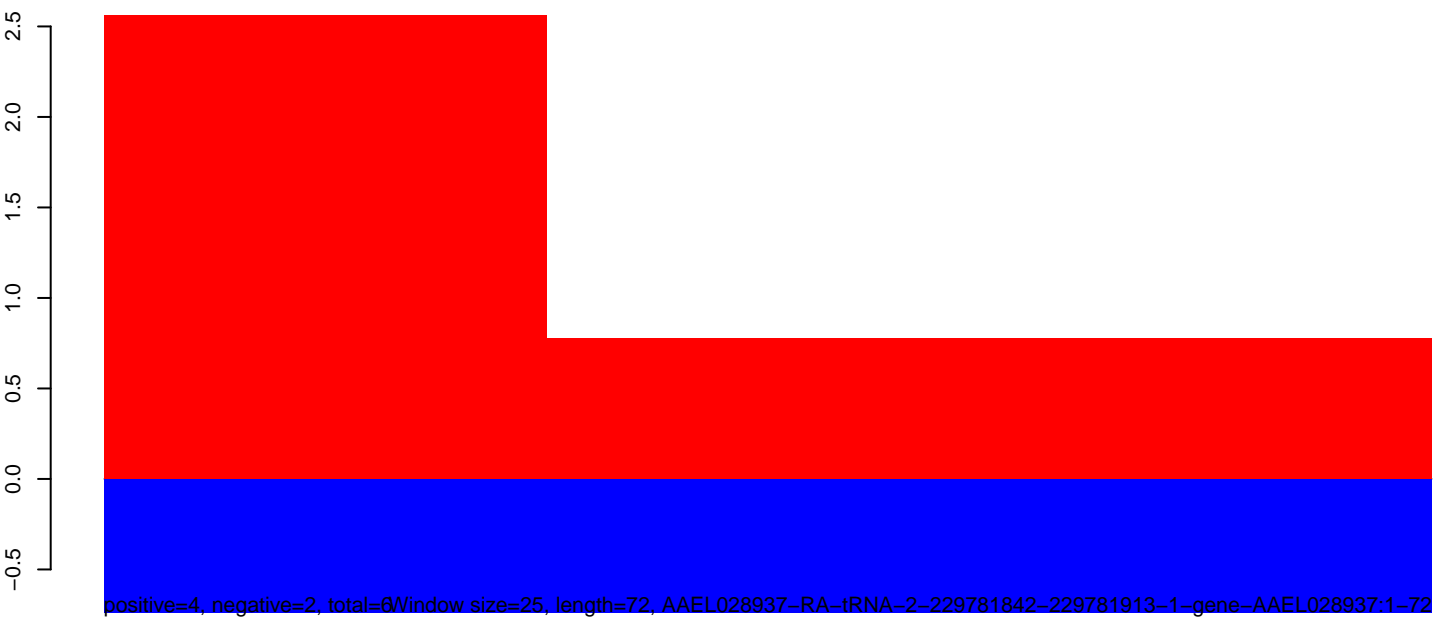
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AeAeg_Fcarc_GH_rep2.24_35.rep



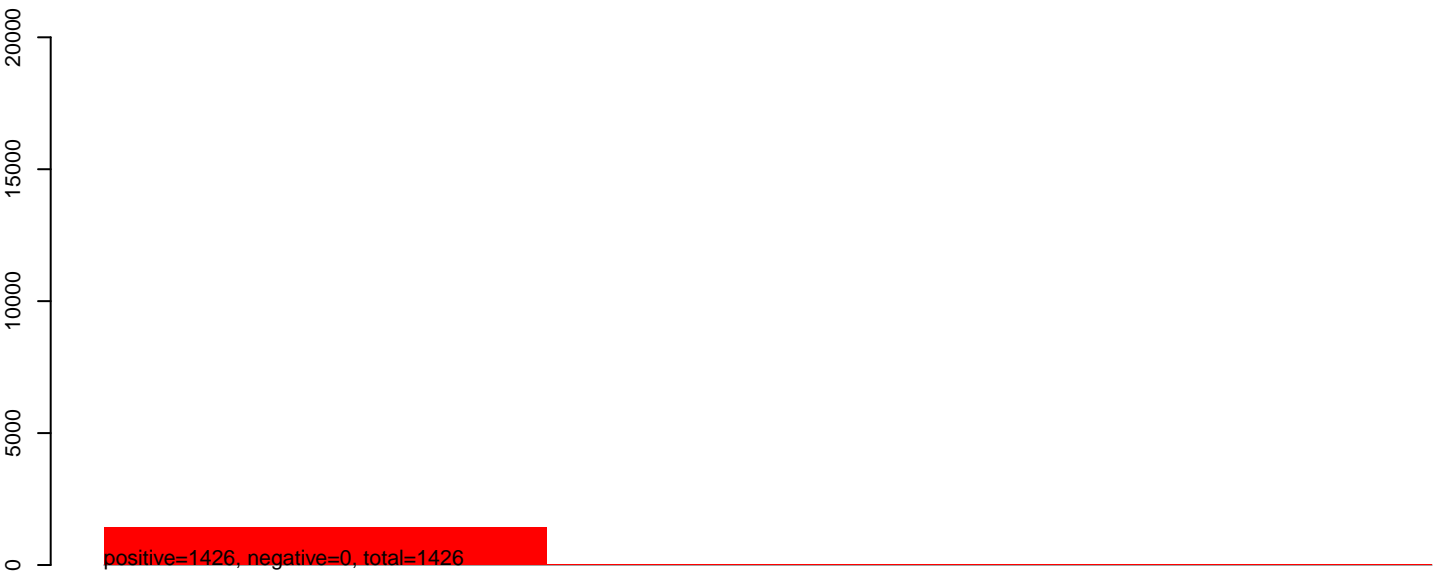
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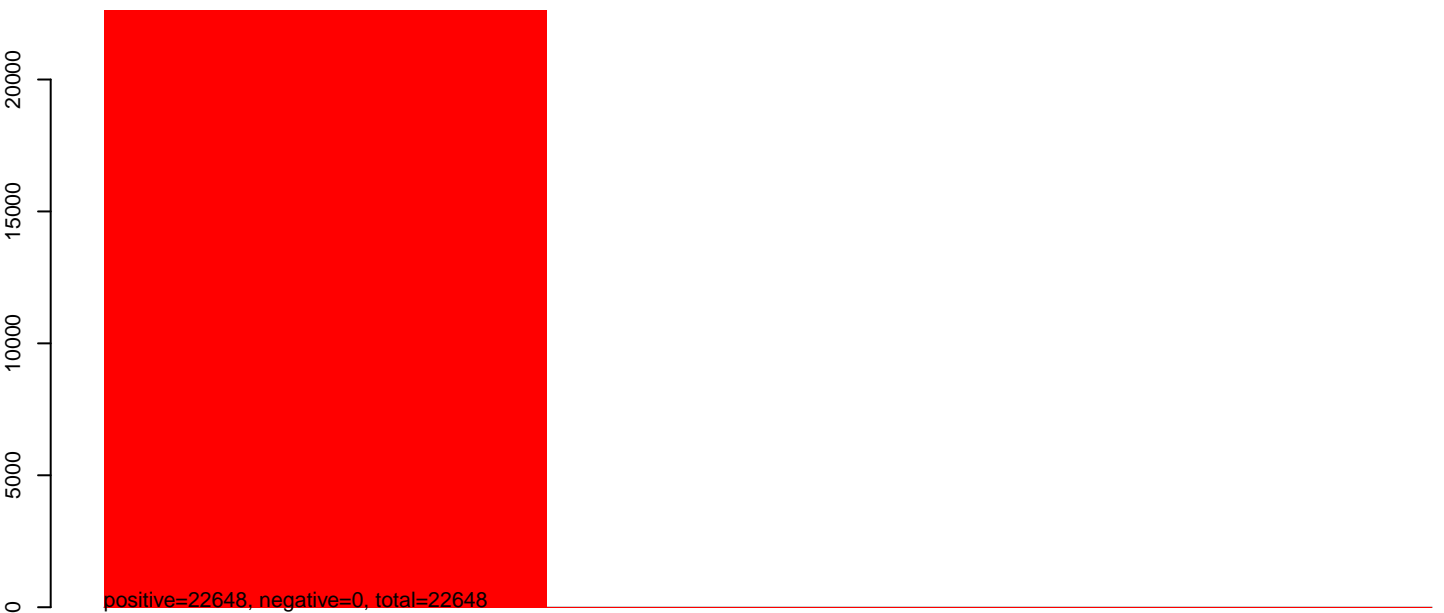
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40 60 80 100

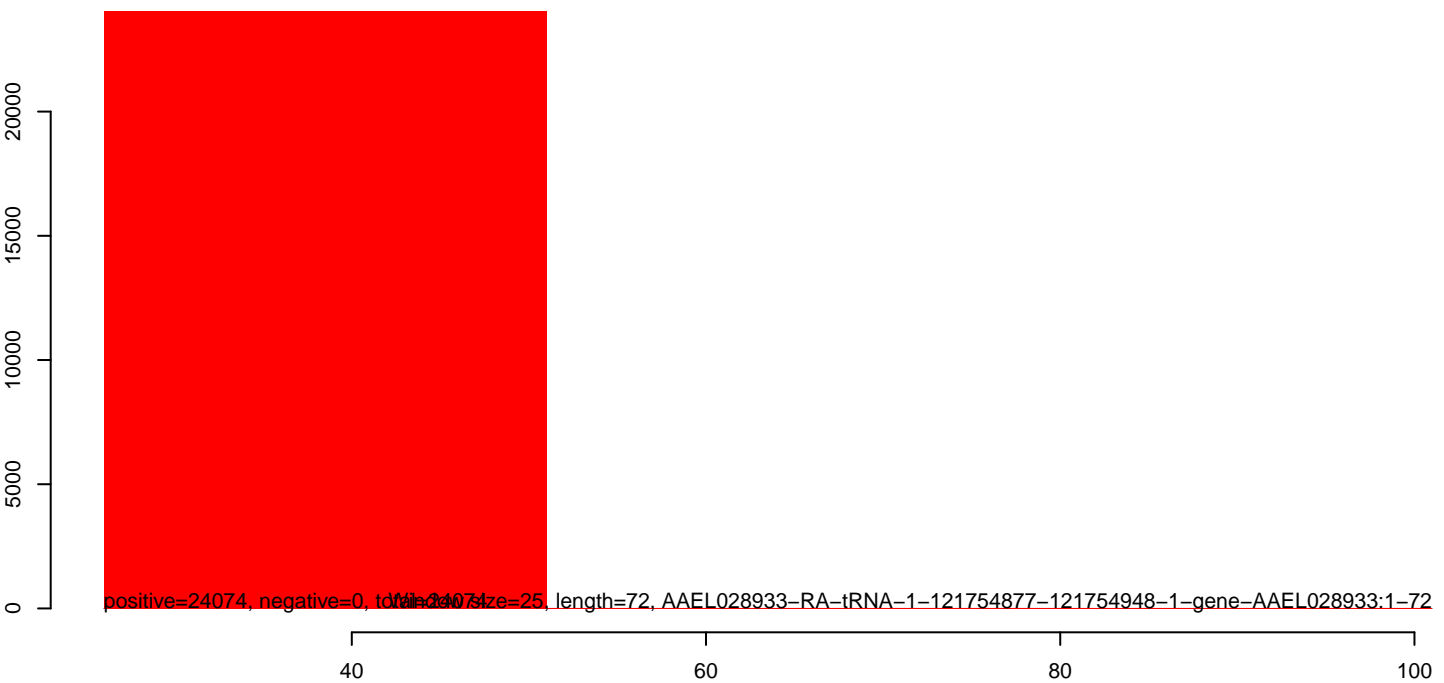
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AeAeg_Fcarc_GH_rep2.24_35.rep



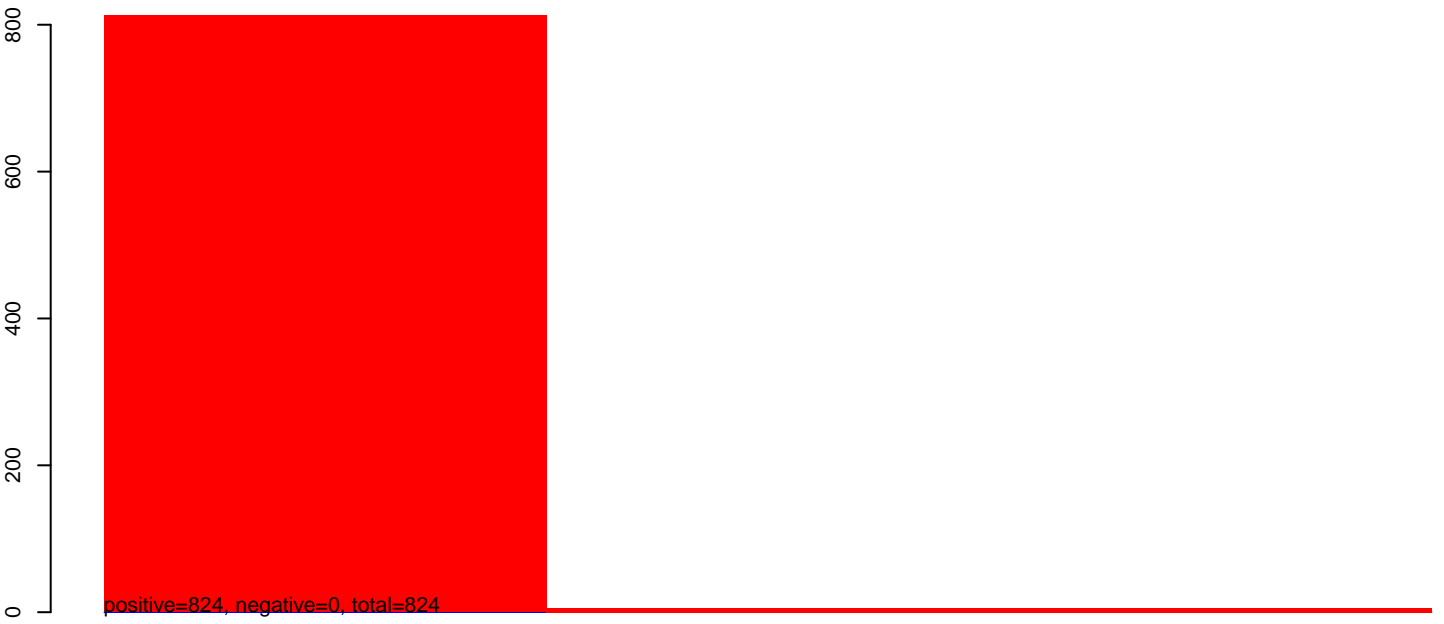
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AeAeg_Fcarc_GH_rep2.18_23.rep



AeAeg_Fcarc_GH_rep2.24_35.rep

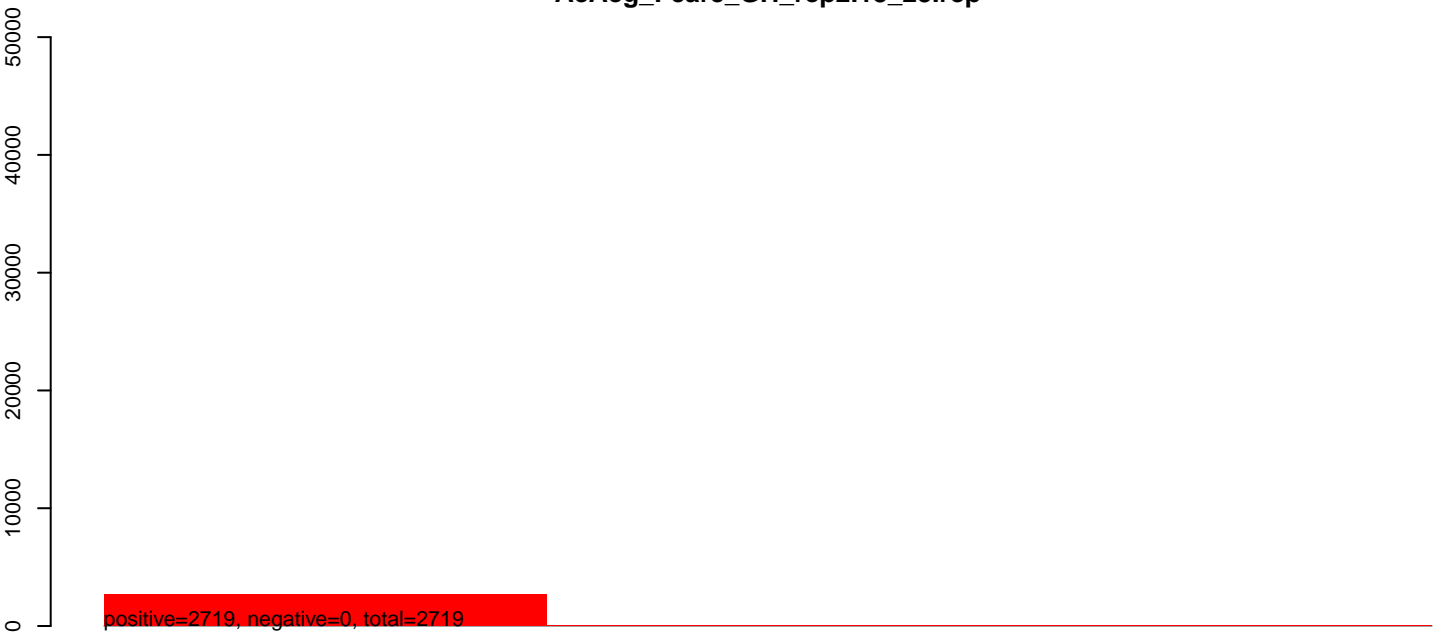


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40 60 80 100

AeAeg_Fcarc_GH_rep2.18_23.rep



AeAeg_Fcarc_GH_rep2.24_35.rep

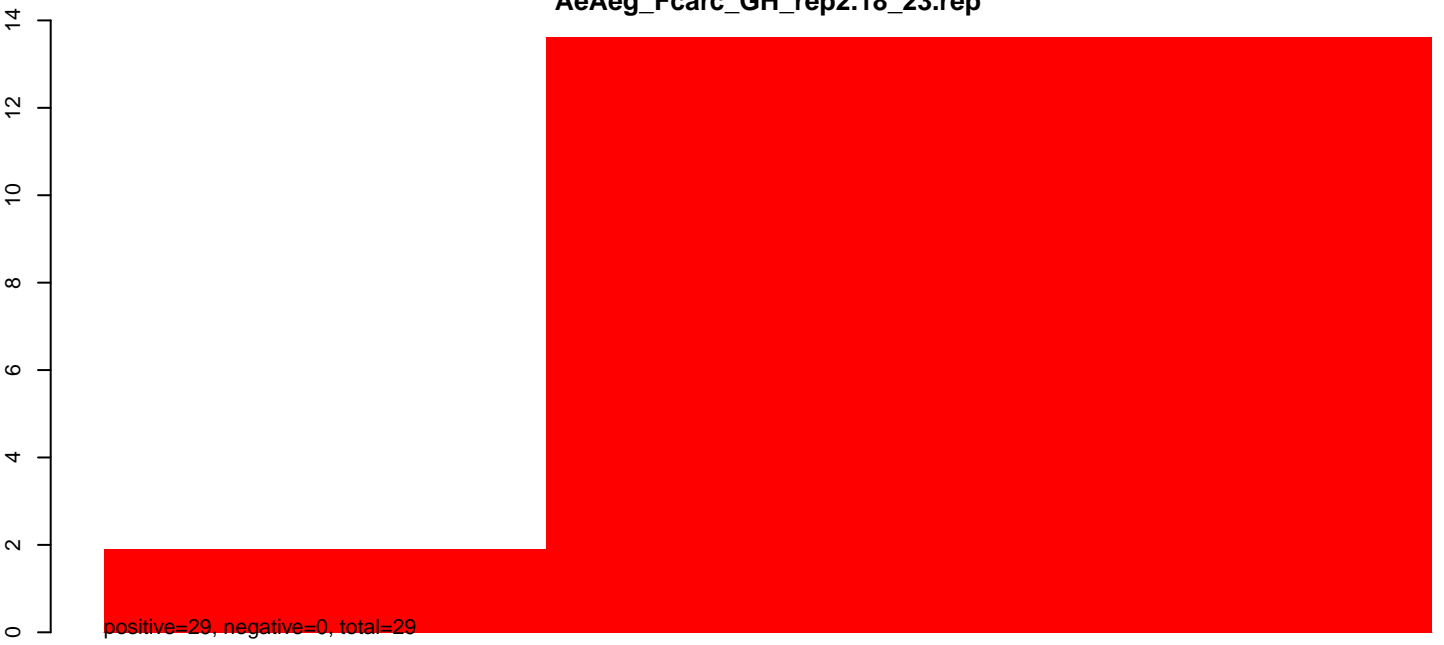


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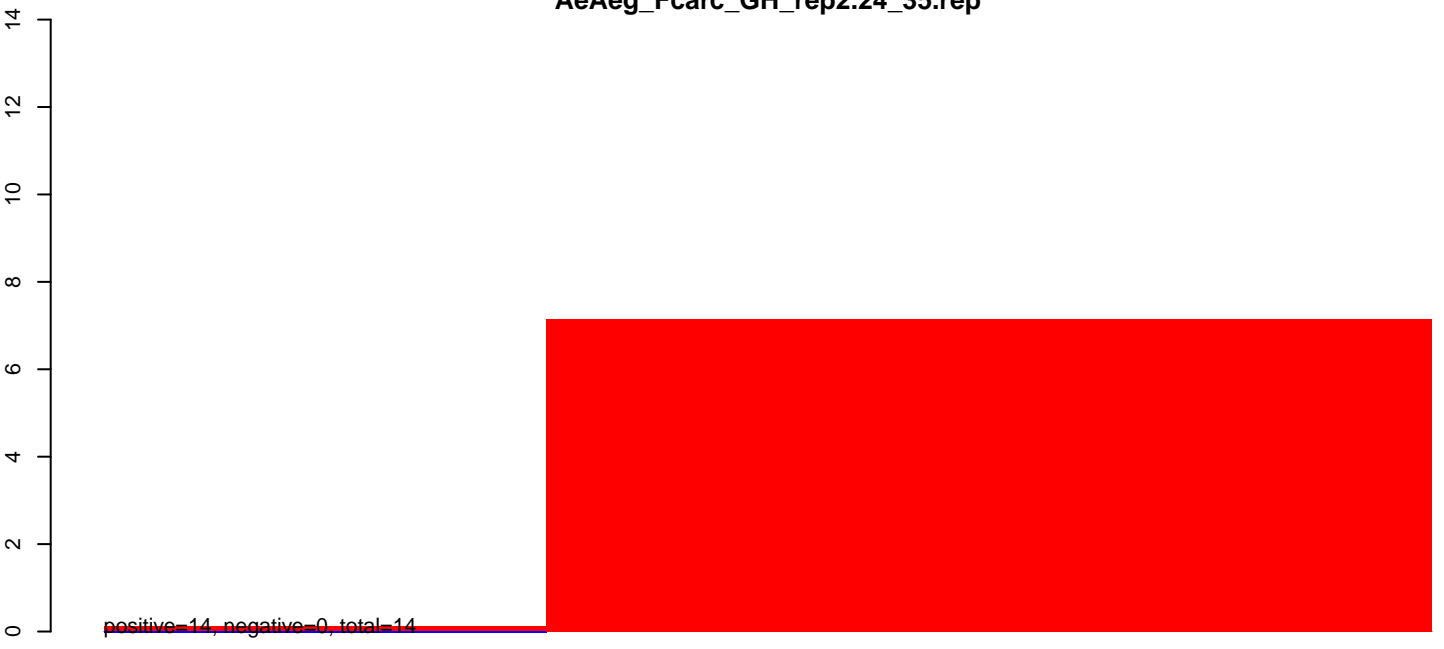


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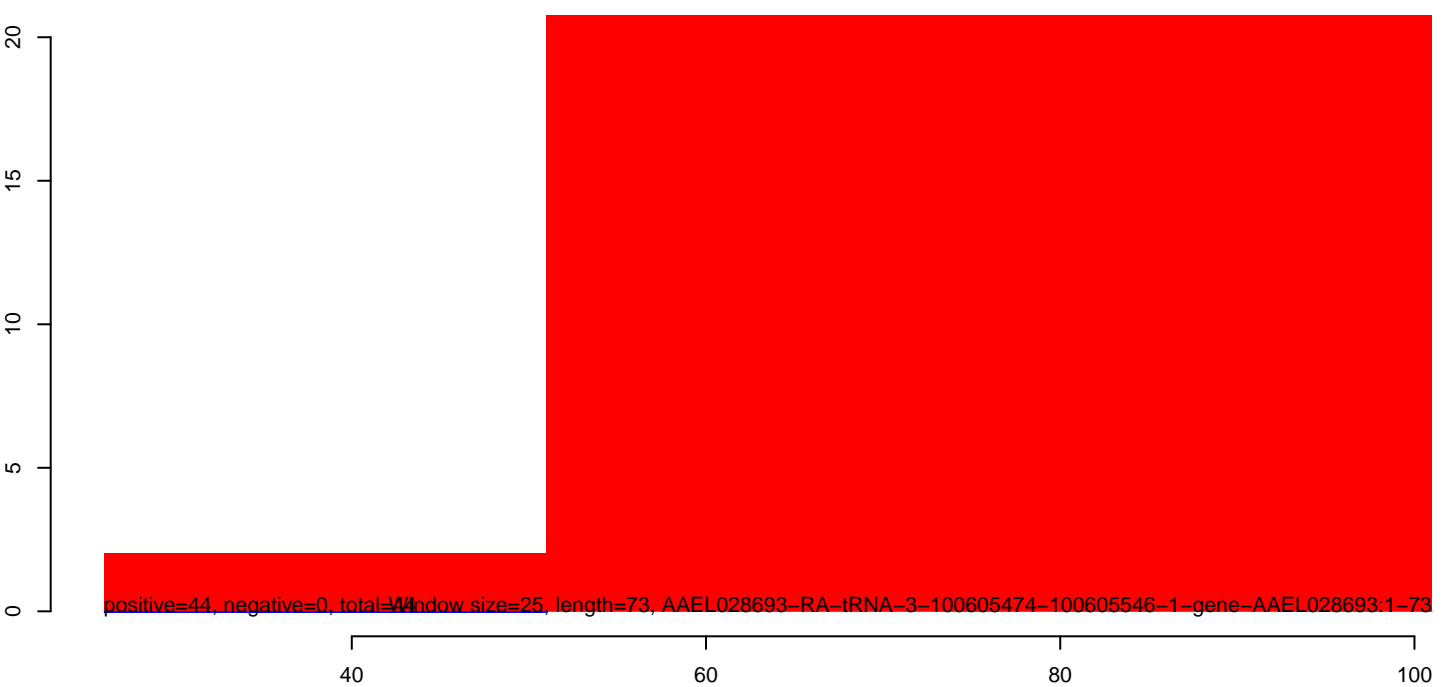
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AeAeg_Fcarc_GH_rep2.24_35.rep



AeAeg_Fcarc_GH_rep2.rep

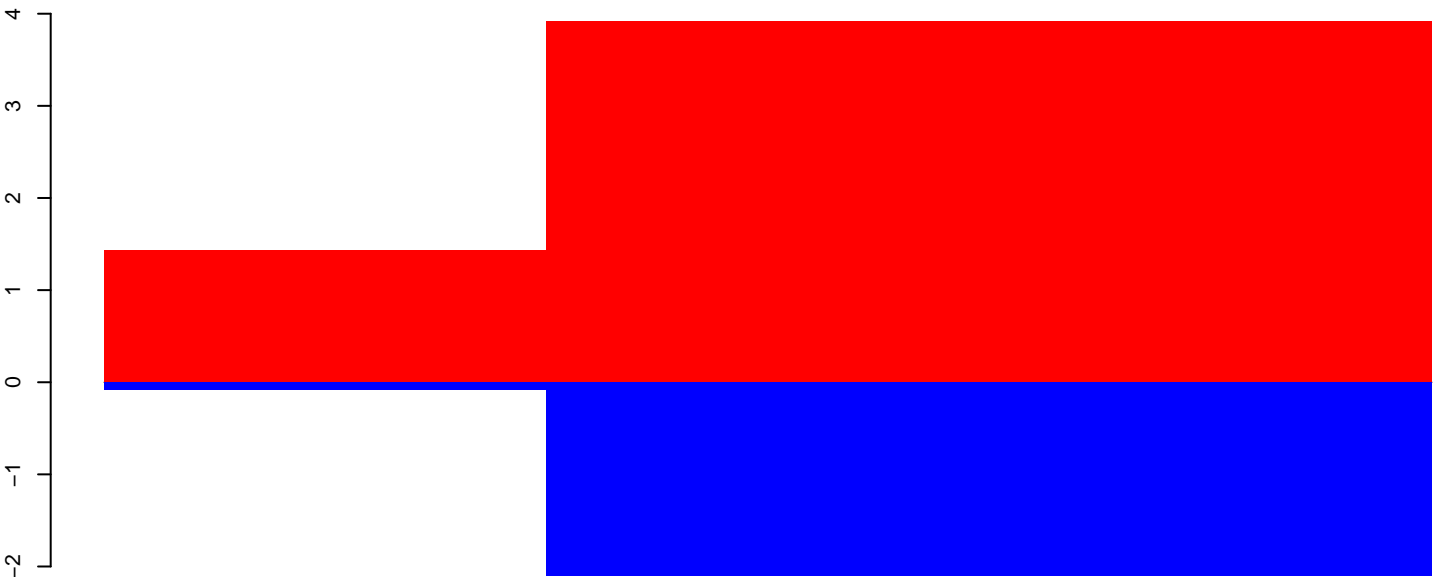


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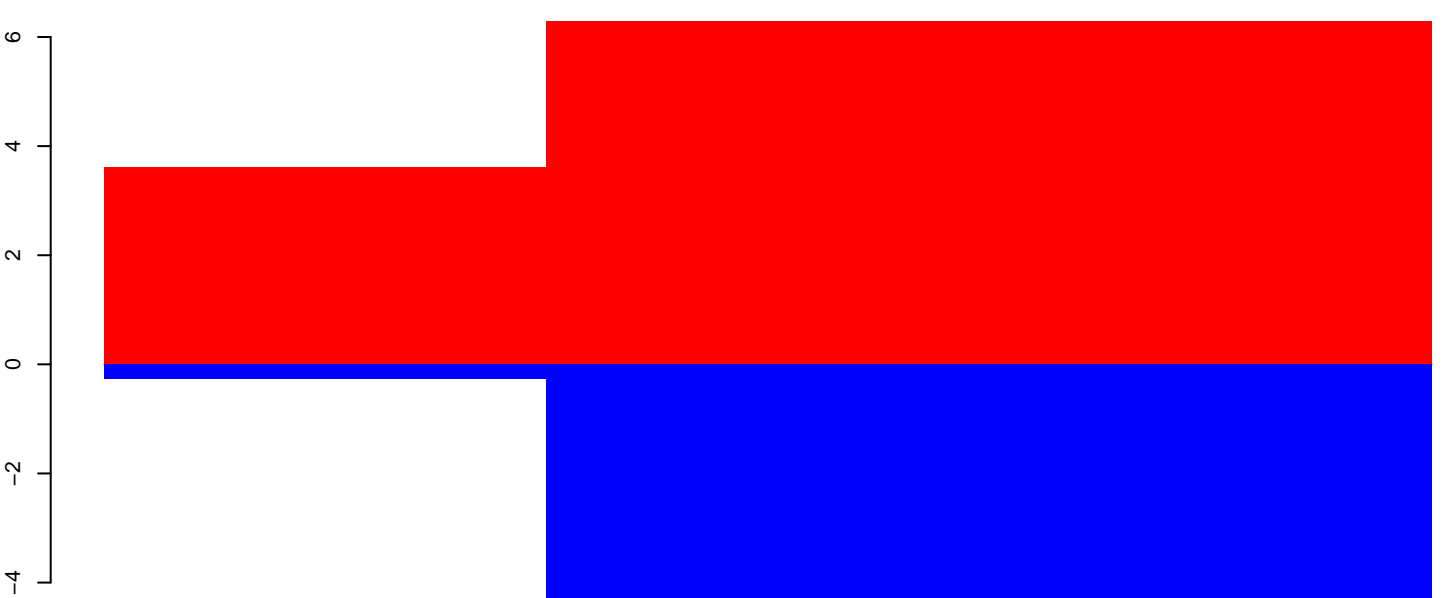
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AeAeg_Fcarc_GH_rep2.24_35.rep



positive=9, negative=4, total=14

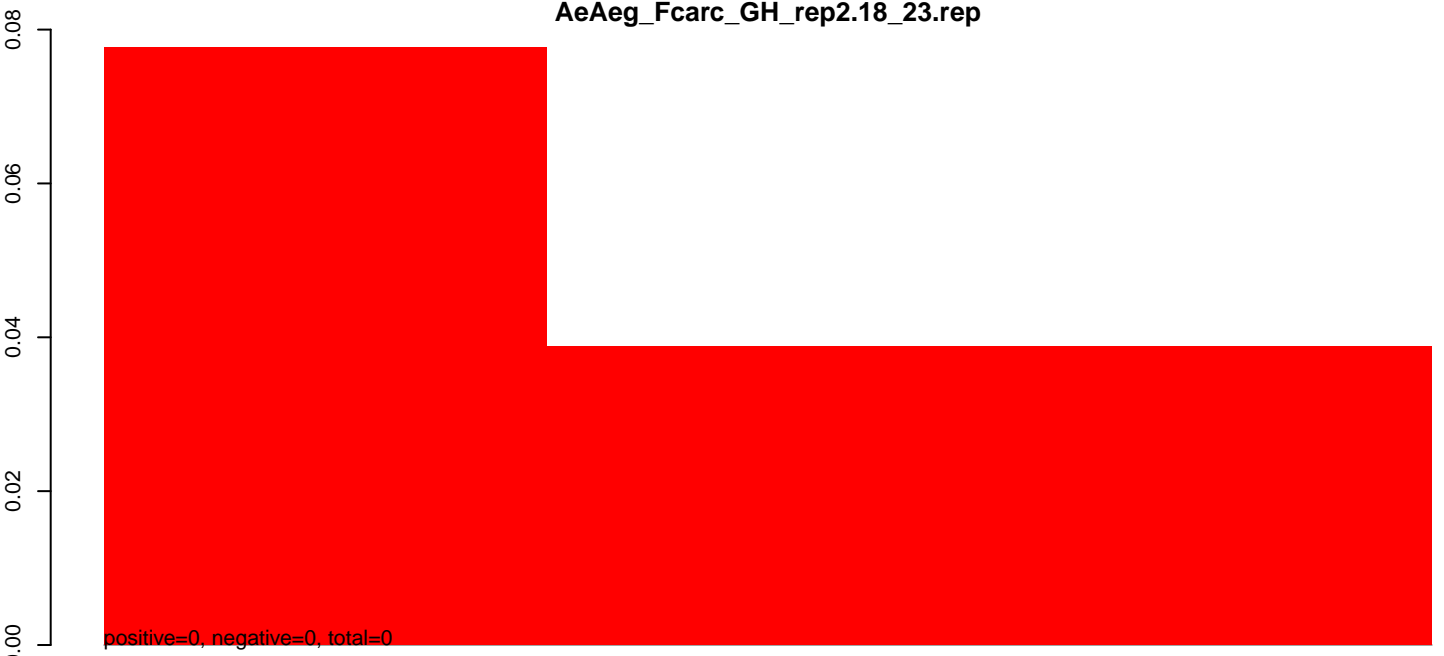
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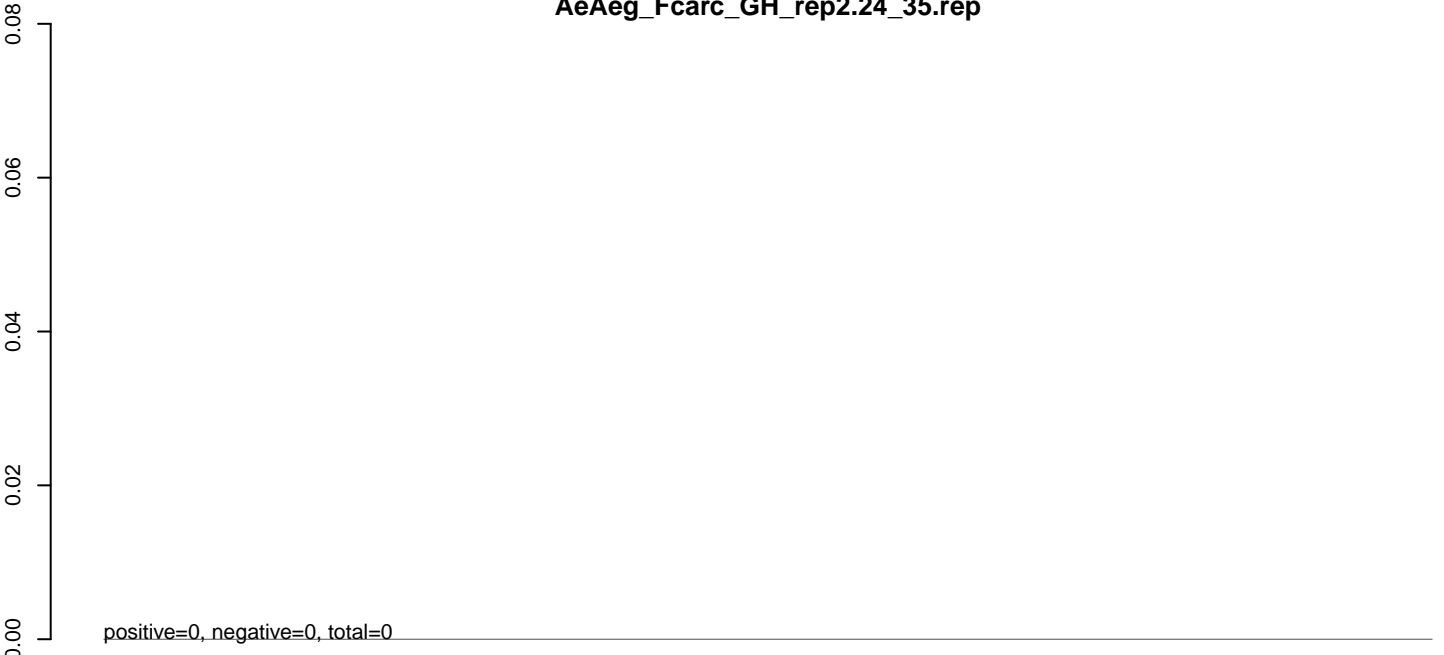
positive=16, negative=10, total=26, window size=25, length=73, AAEL028646-RA-tRNA-3-103003915-103003987-1-gene-AAEL028646.1-73

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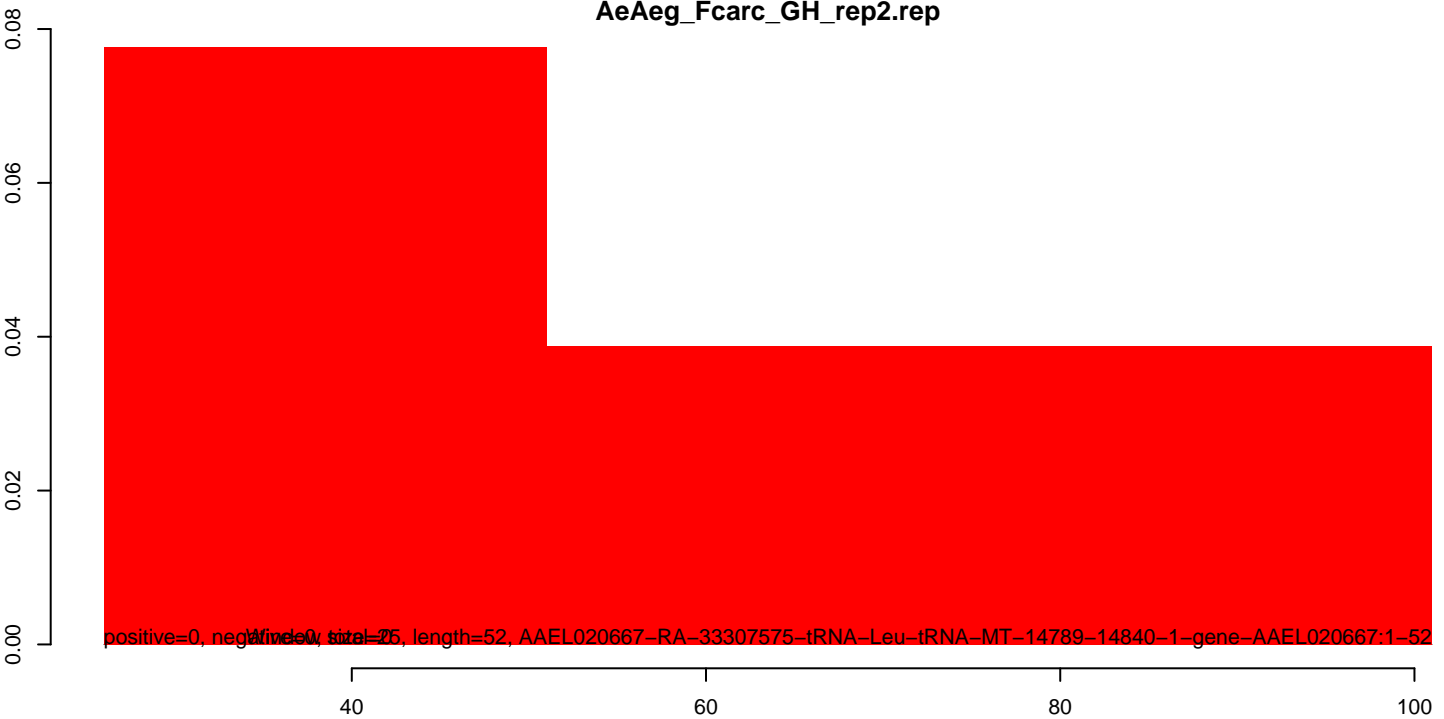
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AeAeg_Fcarc_GH_rep2.24_35.rep



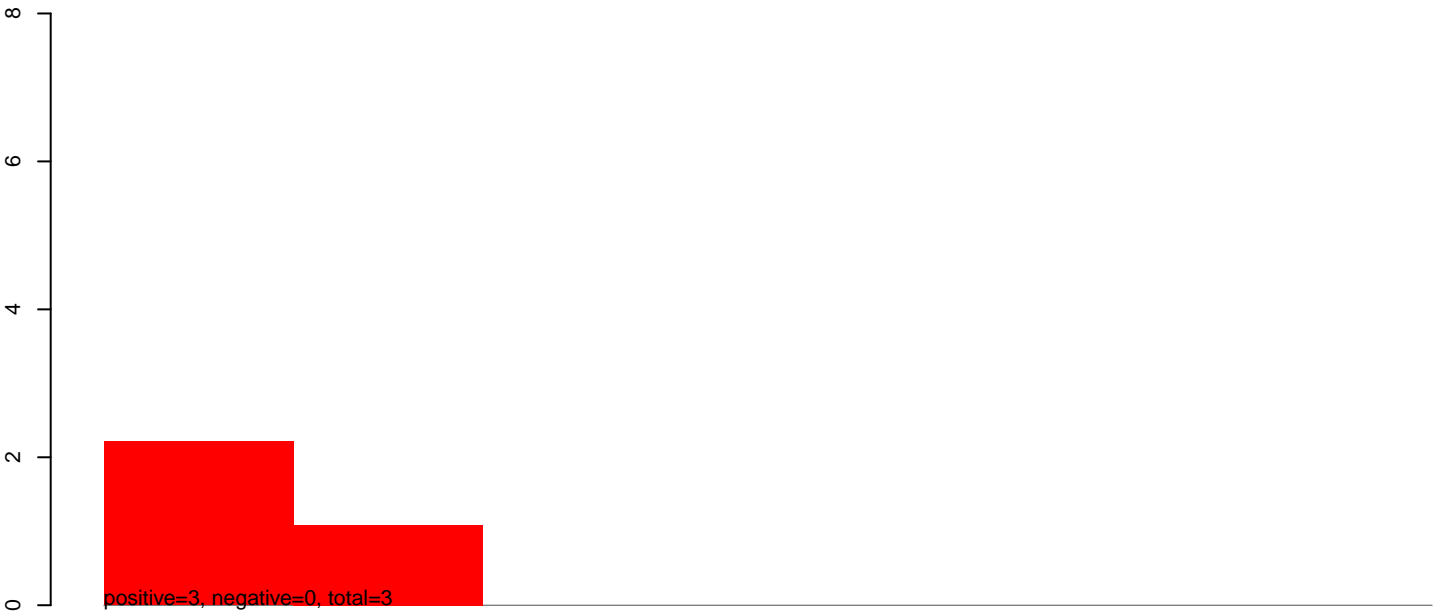
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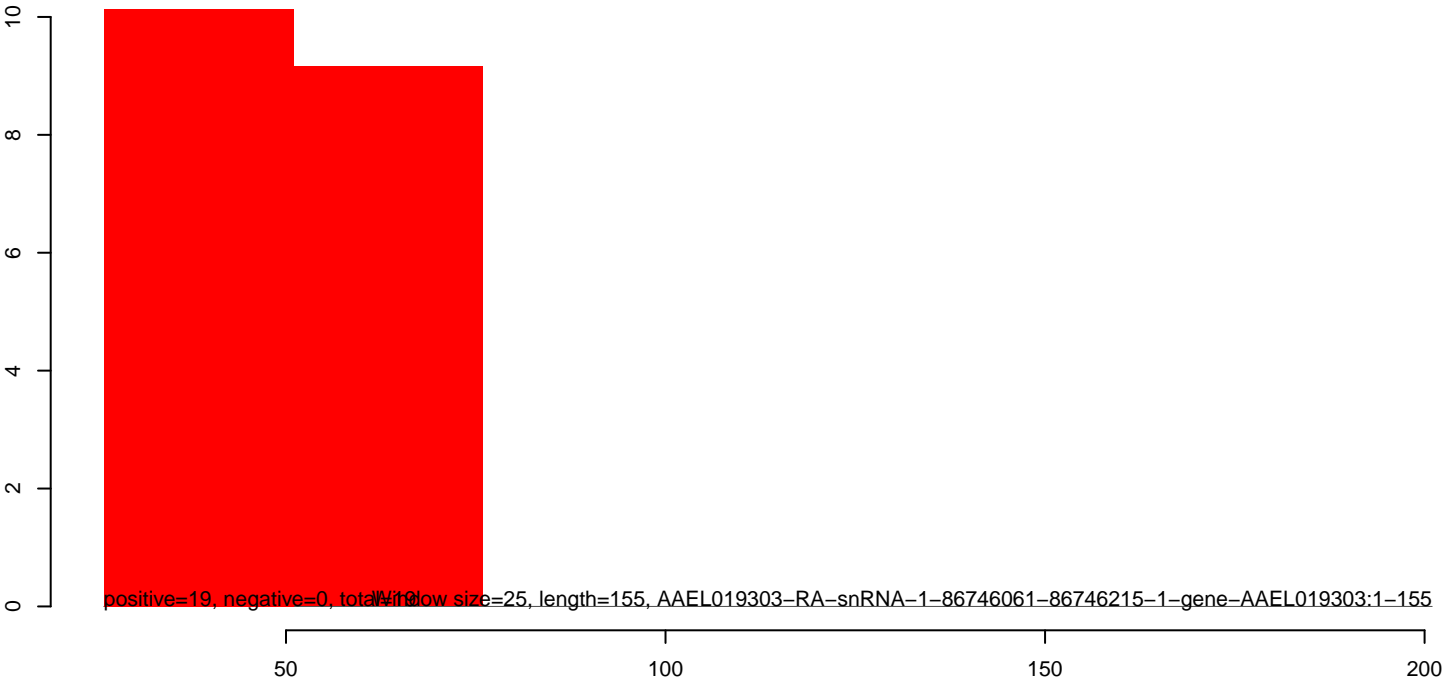
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AeAeg_Fcarc_GH_rep2.24_35.rep



AeAeg_Fcarc_GH_rep2.rep



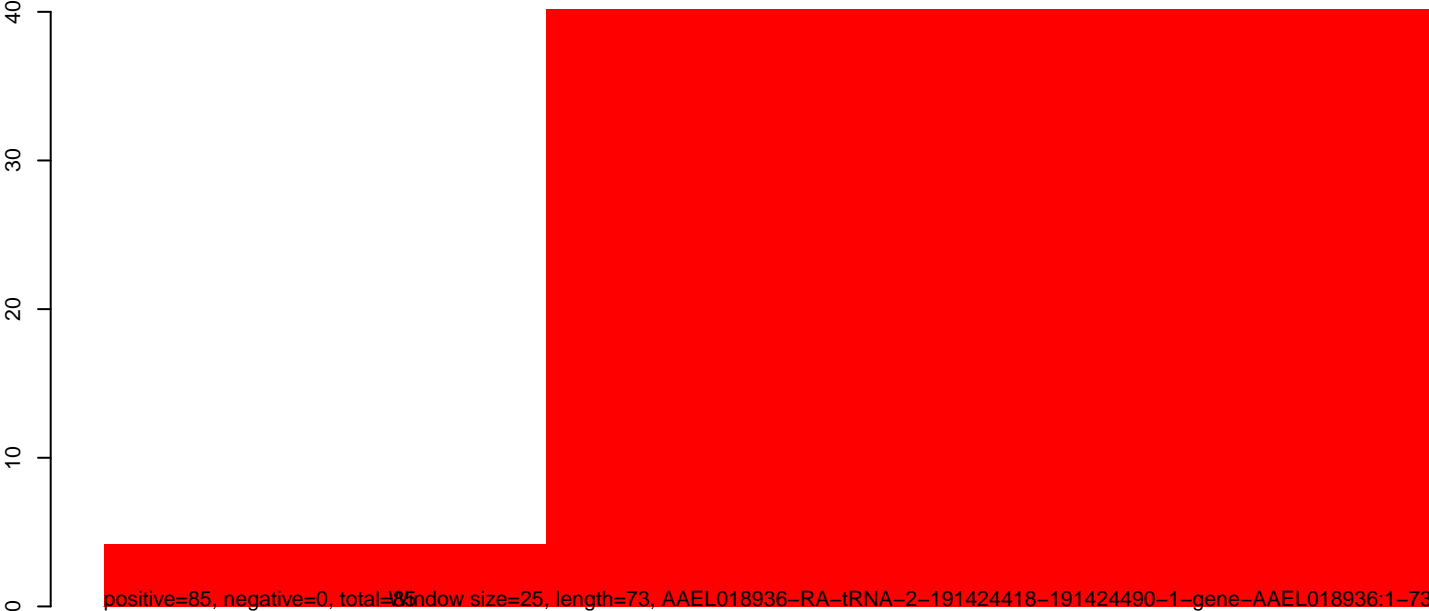
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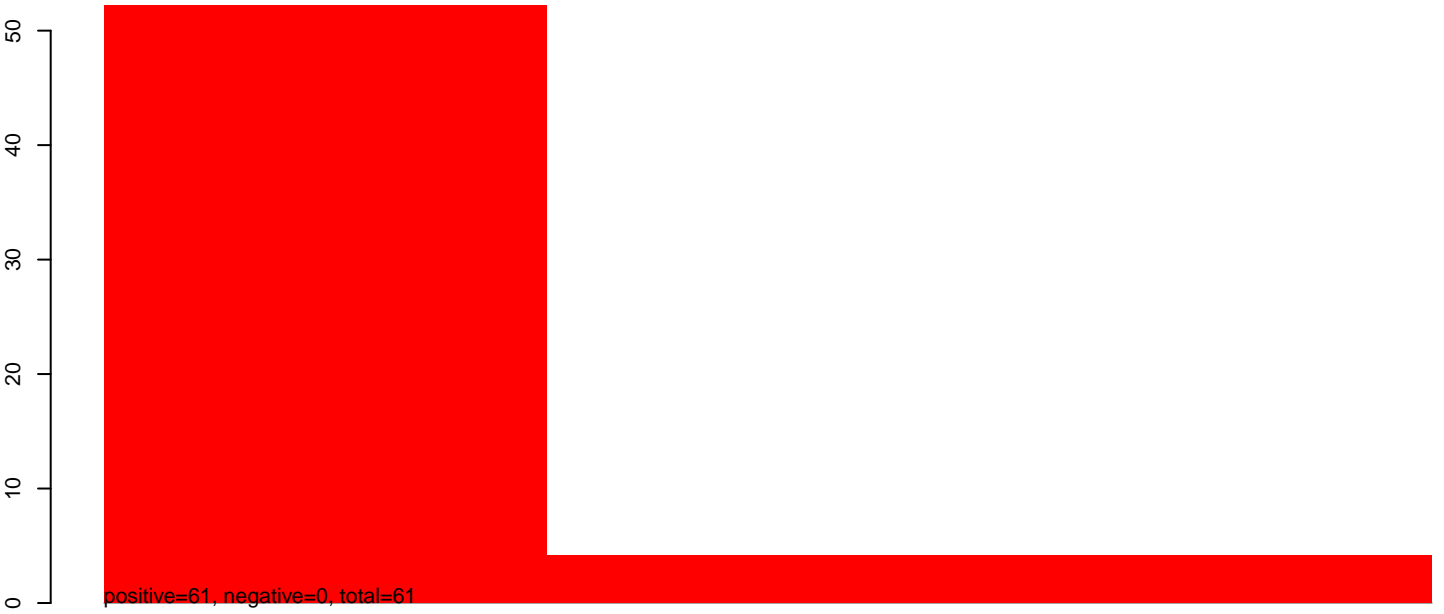
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AeAeg_Fcarc_GH_rep2.rep



AeAeg_Fcarc_GH_rep2.18_23.rep



AeAeg_Fcarc_GH_rep2.24_35.rep



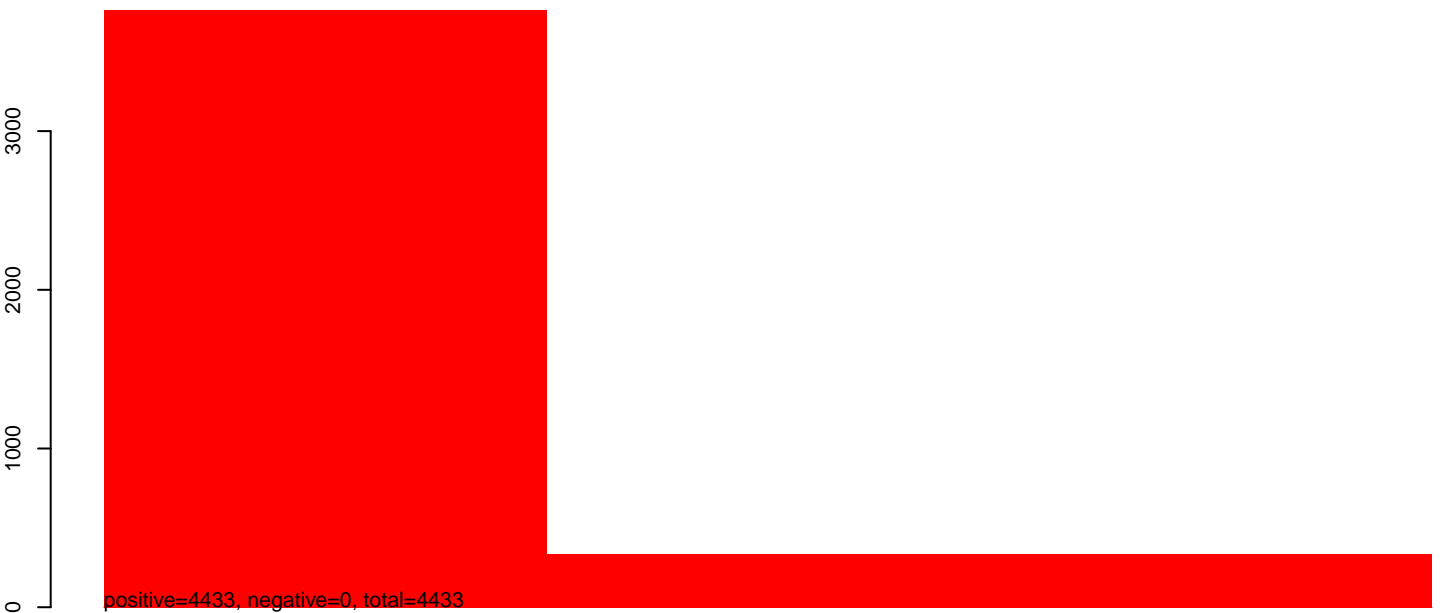
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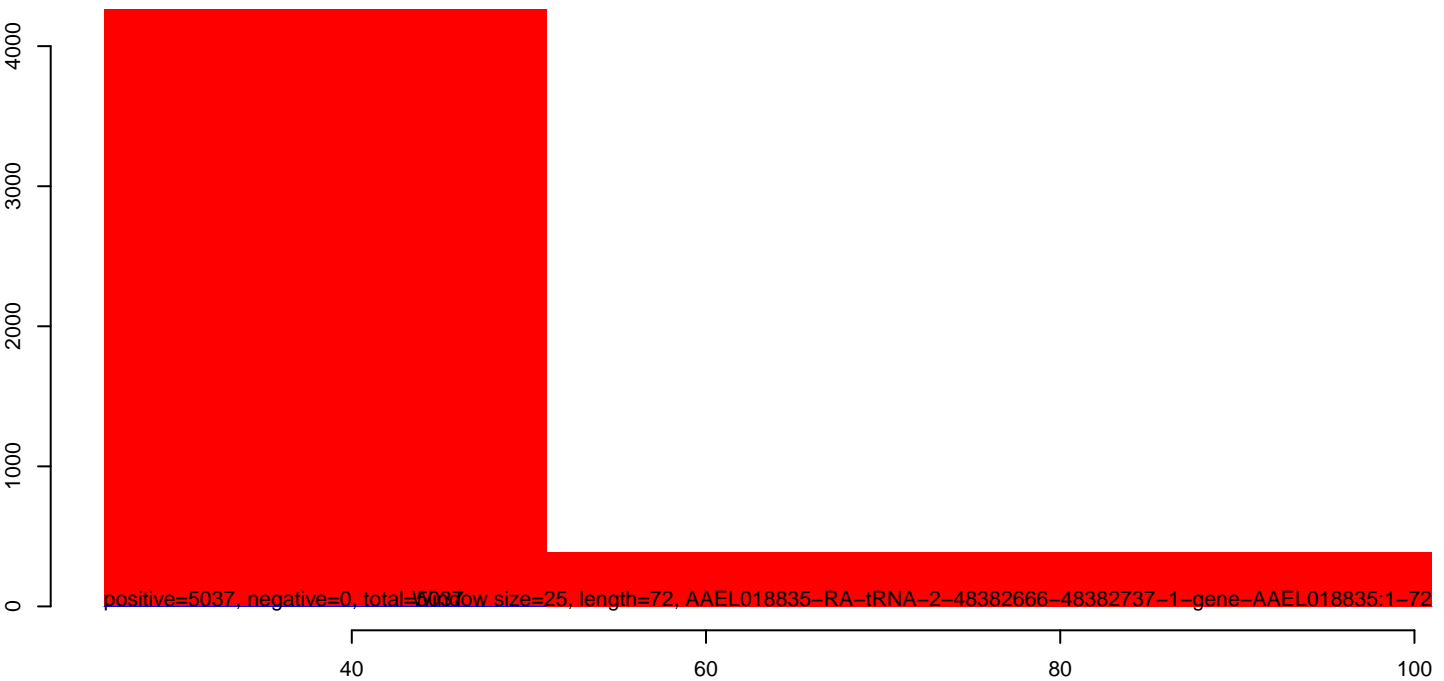
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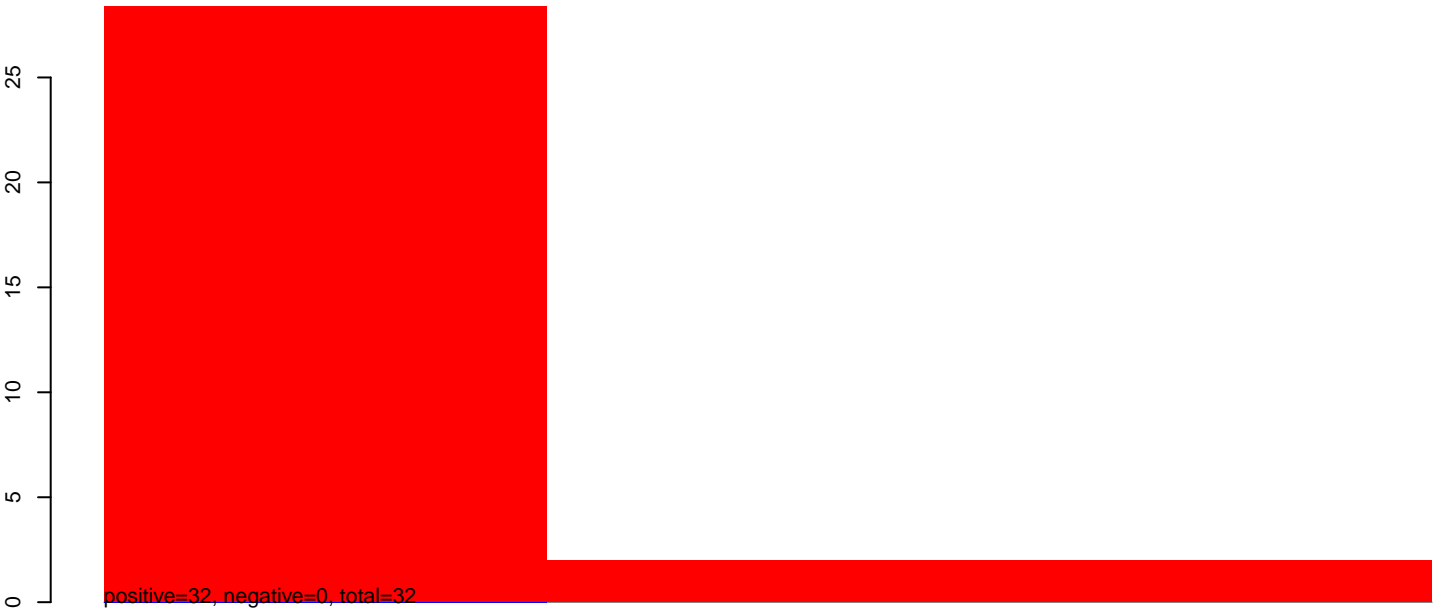
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AeAeg_Fcarc_GH_rep2.rep



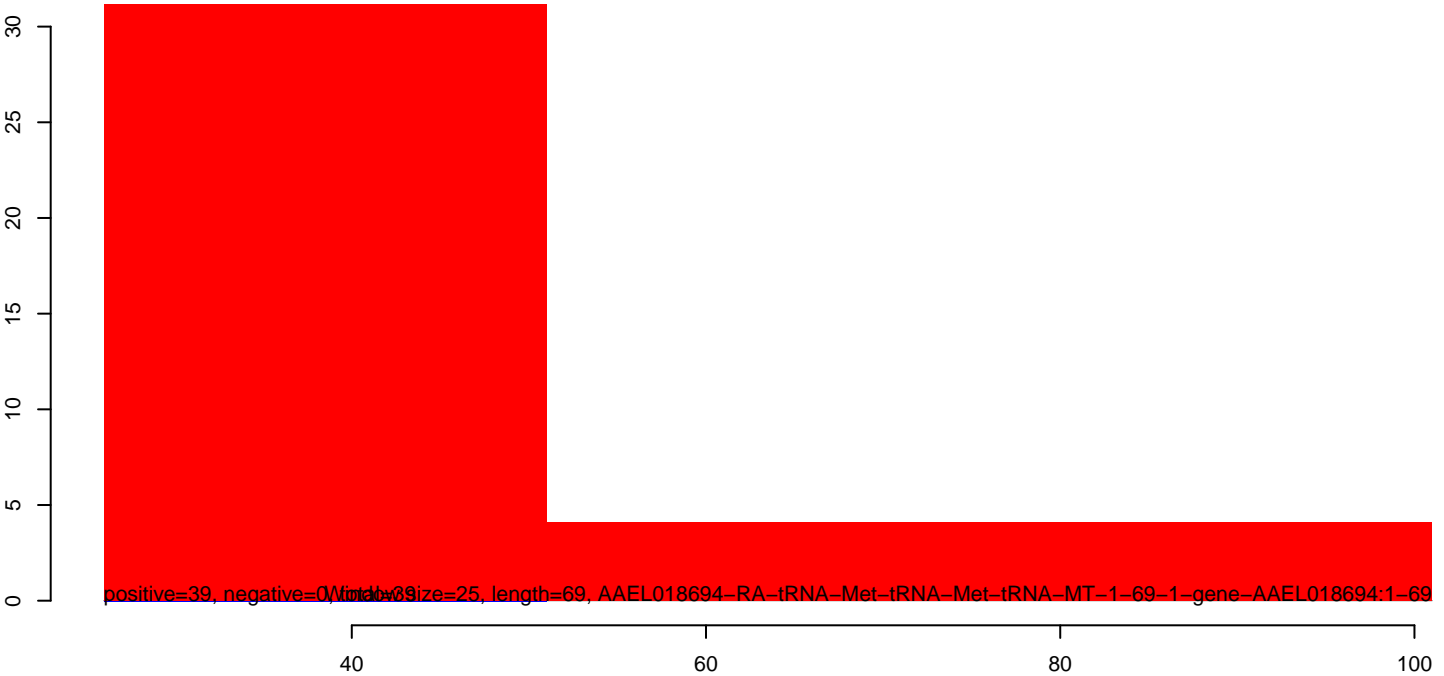
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AeAeg_Fcarc_GH_rep2.24_35.rep



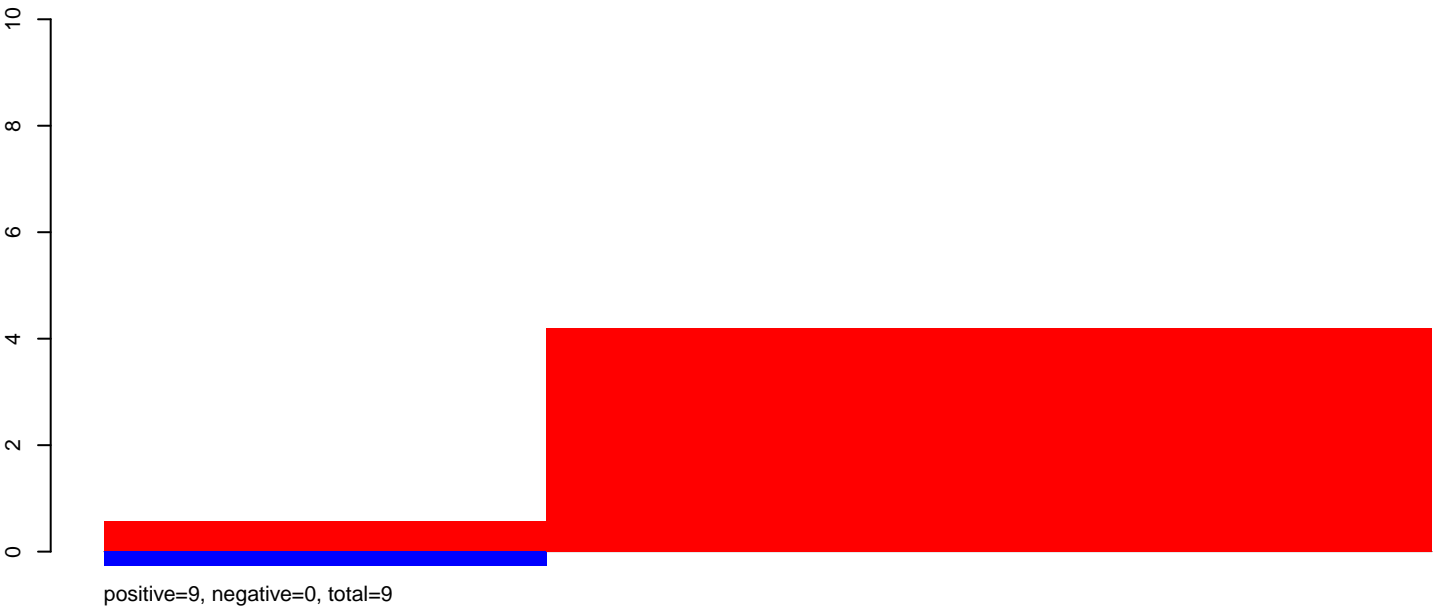
AeAeg_Fcarc_GH_rep2.rep



AeAeg_Fcarc_GH_rep2.18_23.rep



AeAeg_Fcarc_GH_rep2.24_35.rep



AeAeg_Fcarc_GH_rep2.rep



Window size=25, length=69, AAEL018693-RA-tRNA-Glu-tRNA-Gln-tRNA-MT-16677-16745-1-gene-AAEL018693:1-69

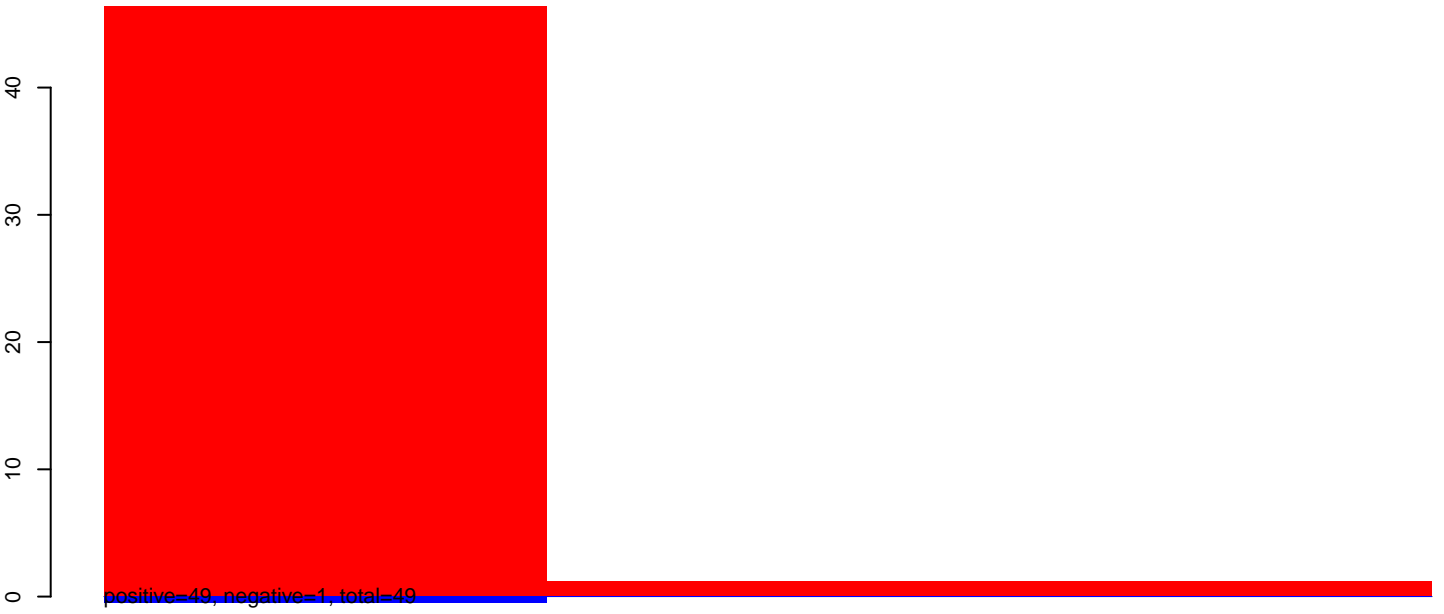
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60

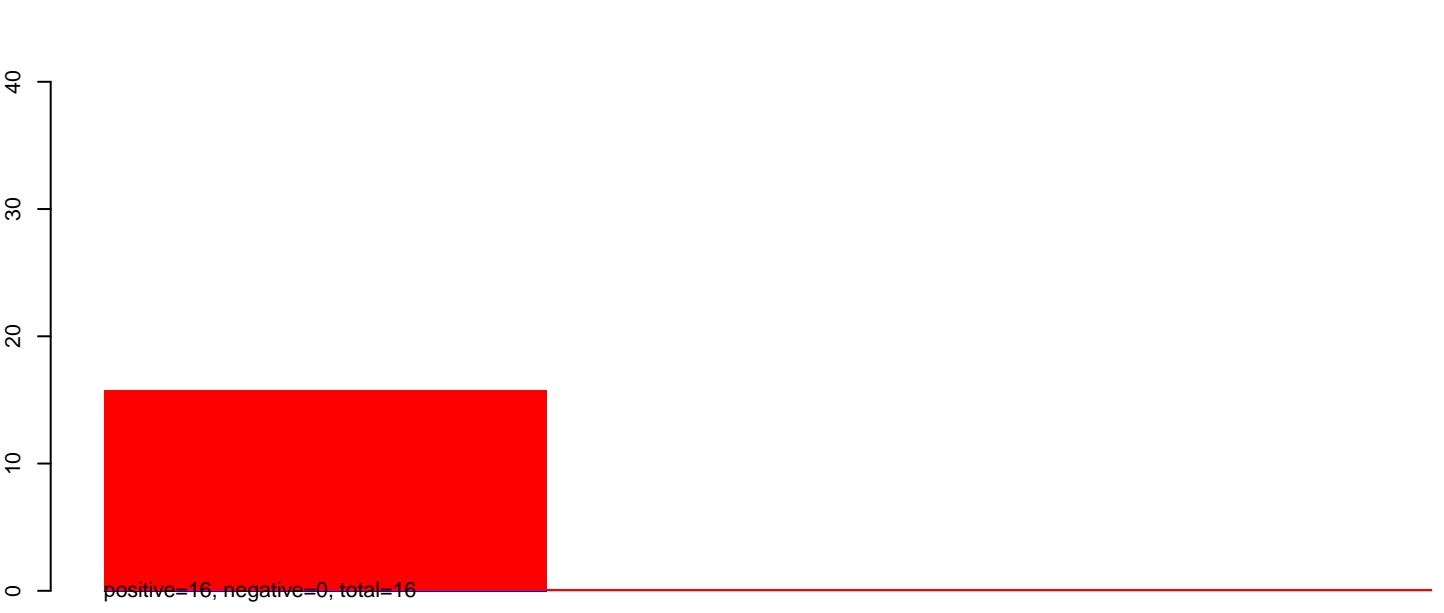
80

100

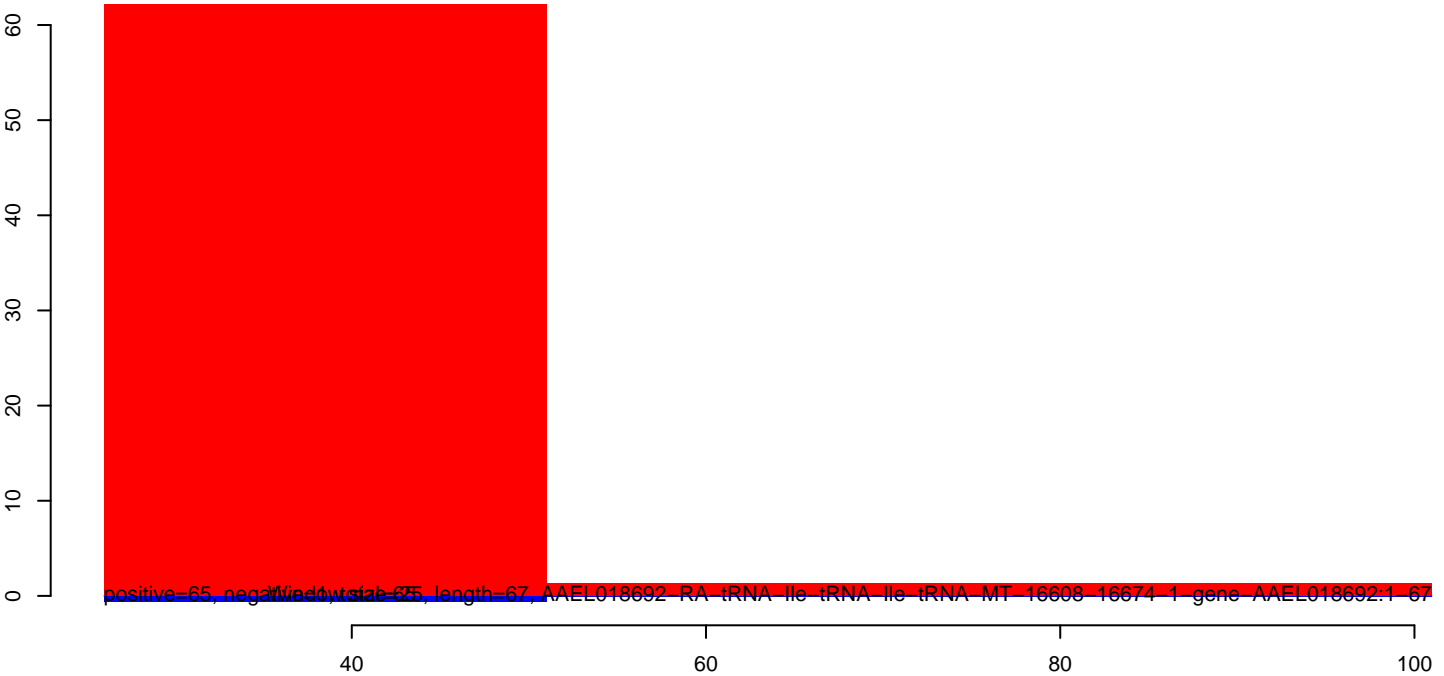
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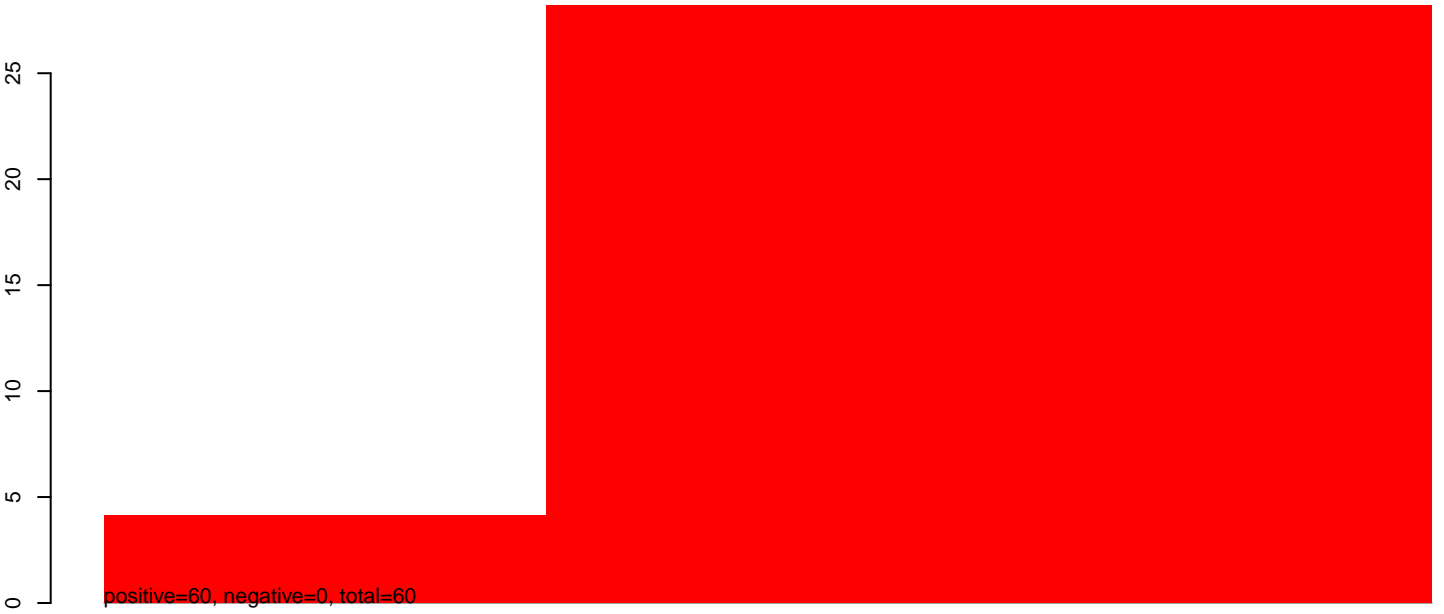
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AeAeg_Fcarc_GH_rep2.rep



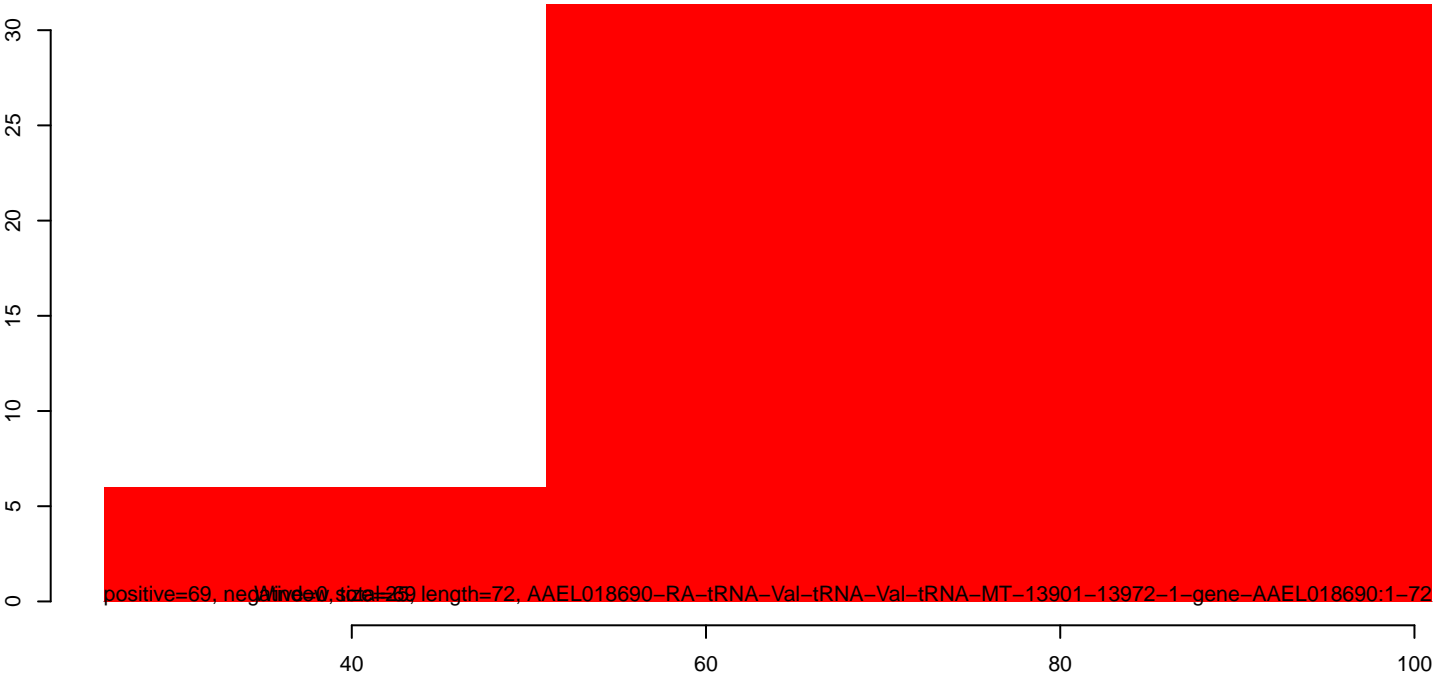
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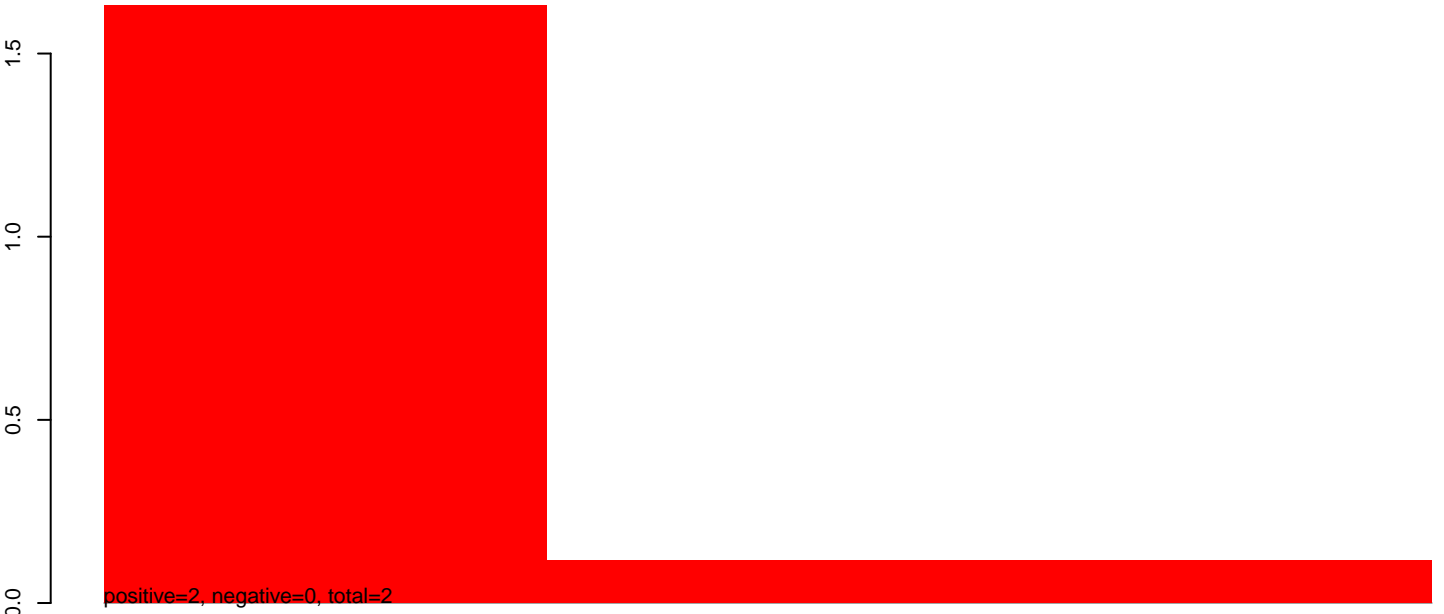
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AeAeg_Fcarc_GH_rep2.rep



AeAeg_Fcarc_GH_rep2.18_23.rep



AeAeg_Fcarc_GH_rep2.24_35.rep

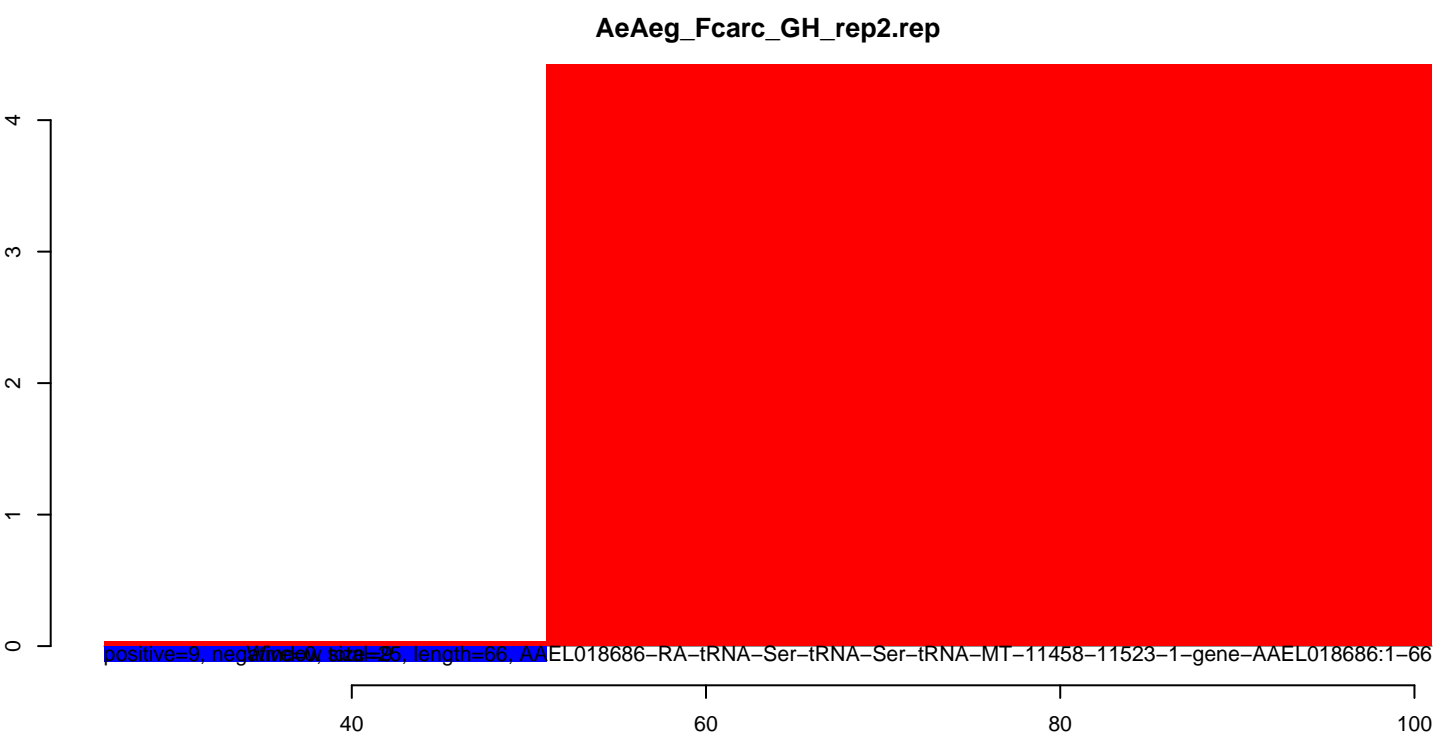
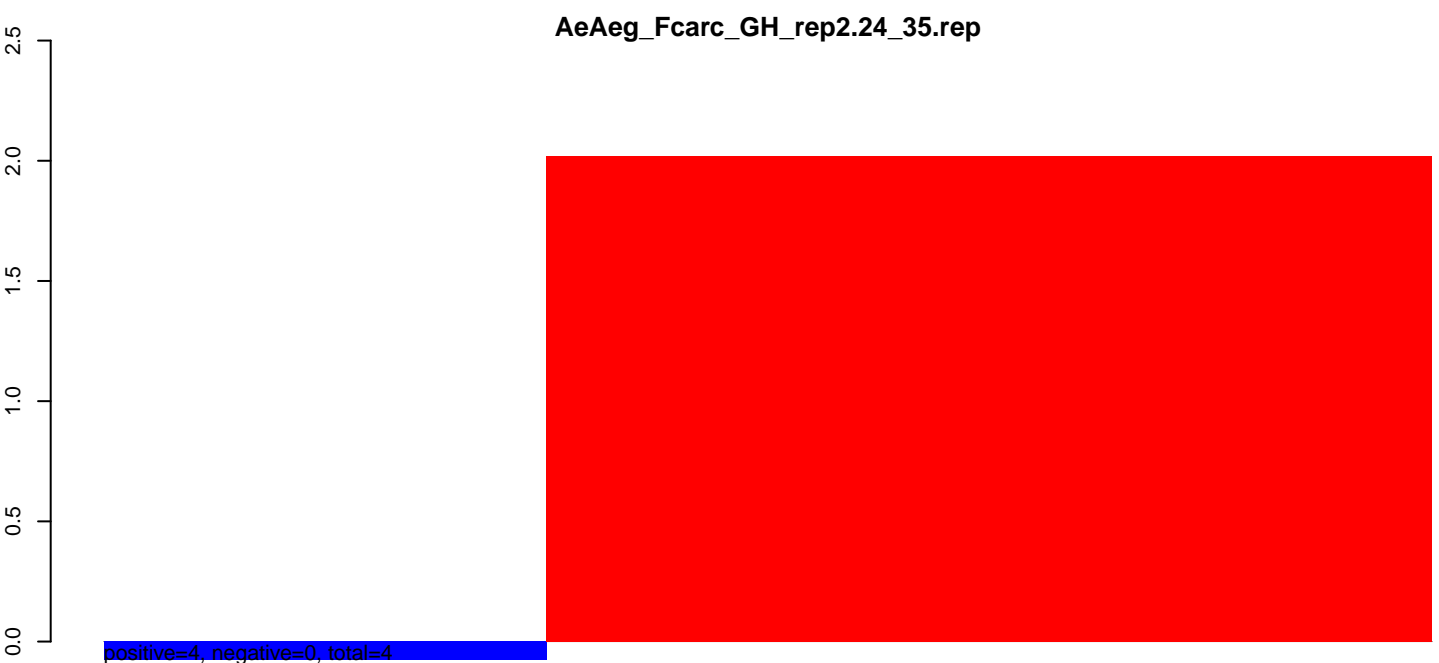
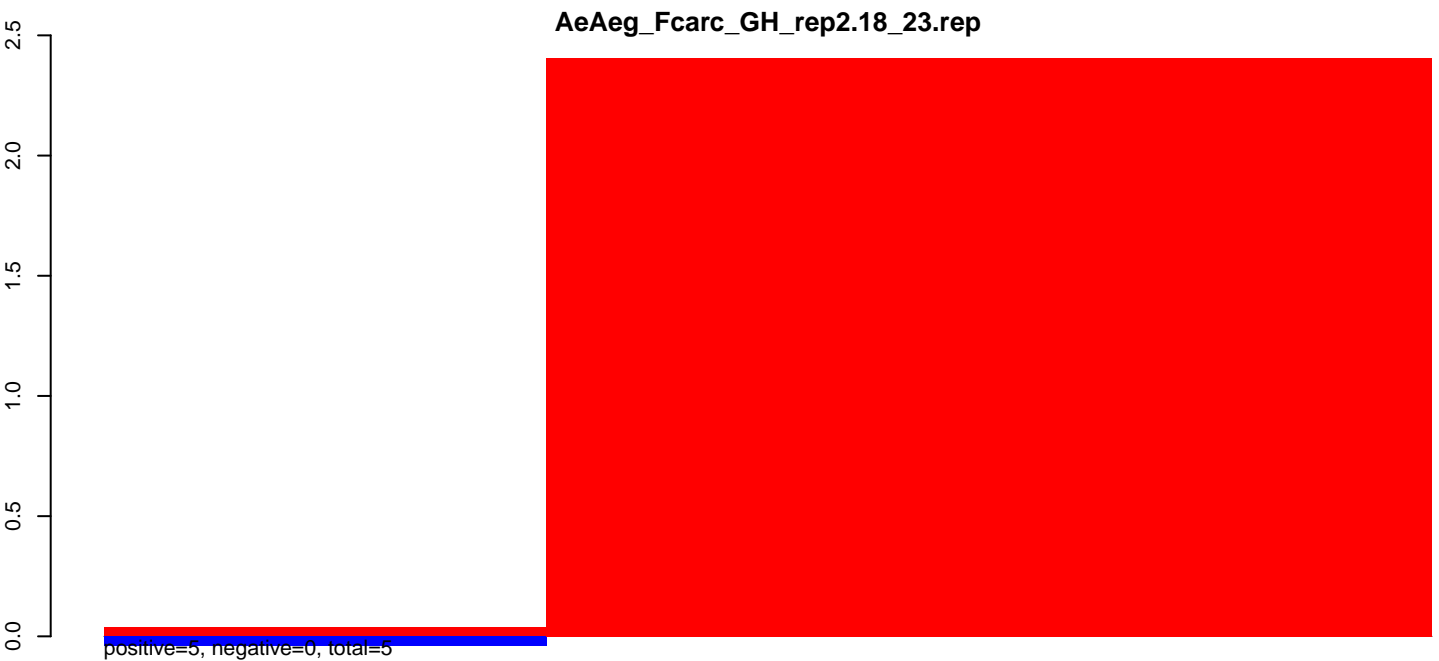


AeAeg_Fcarc_GH_rep2.rep

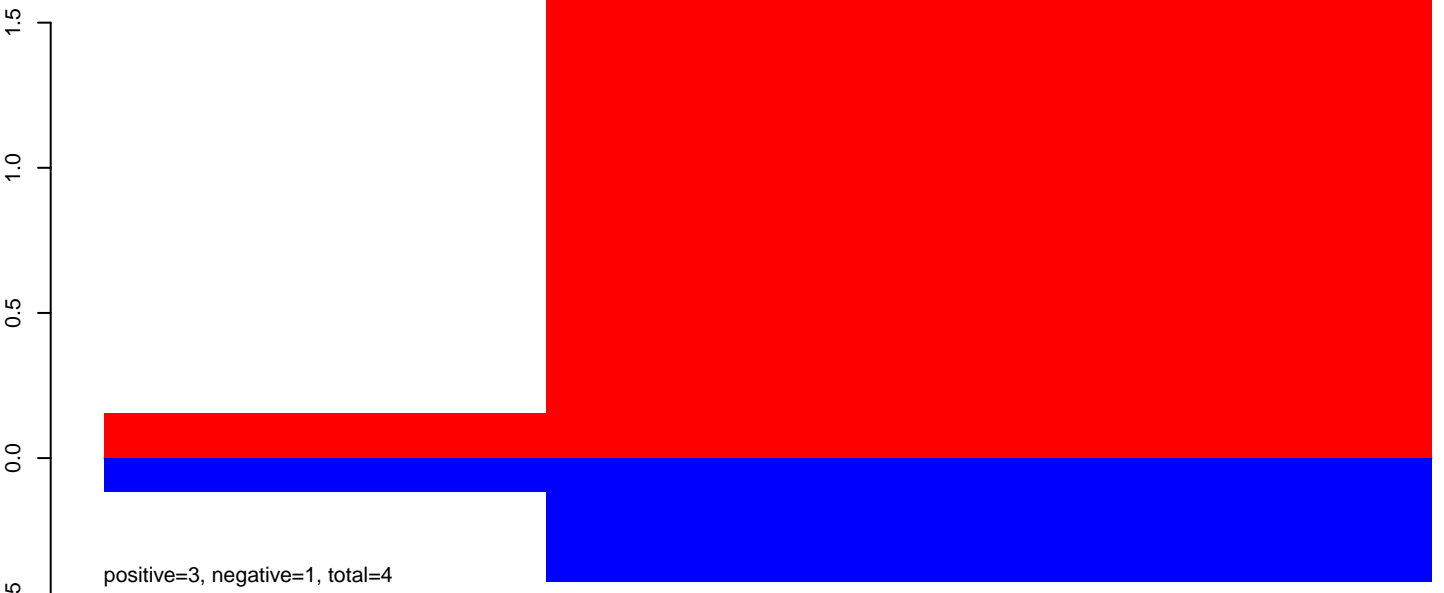


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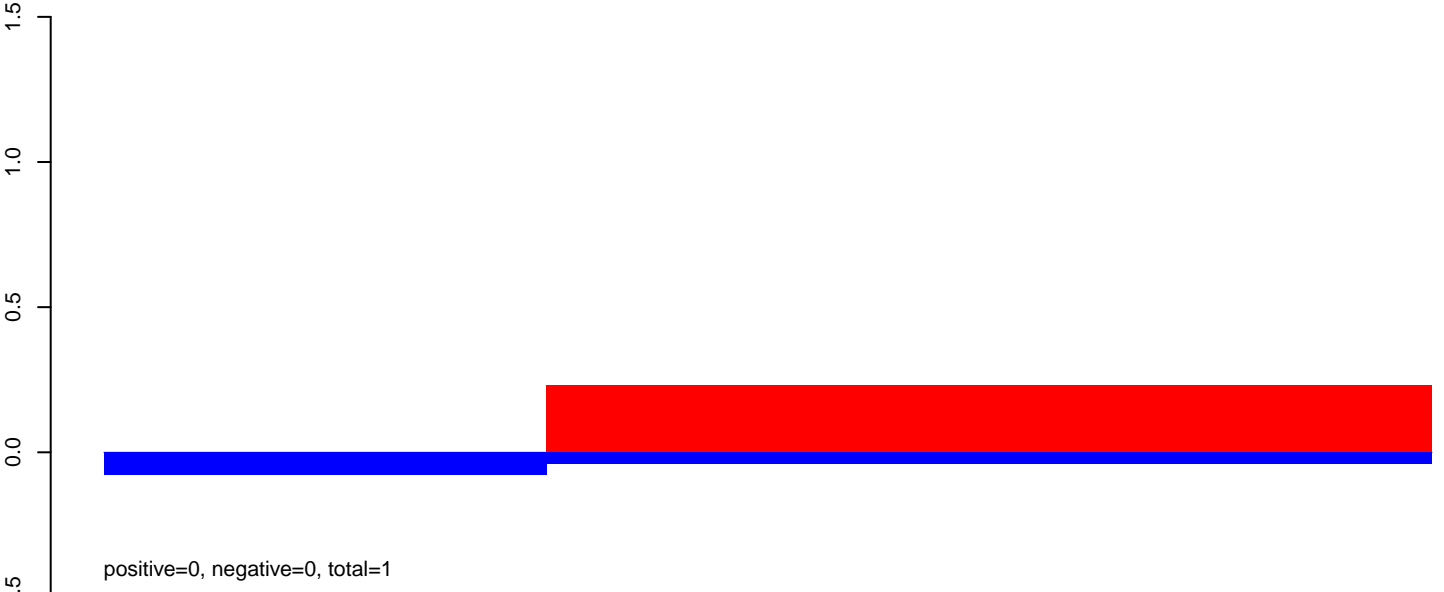
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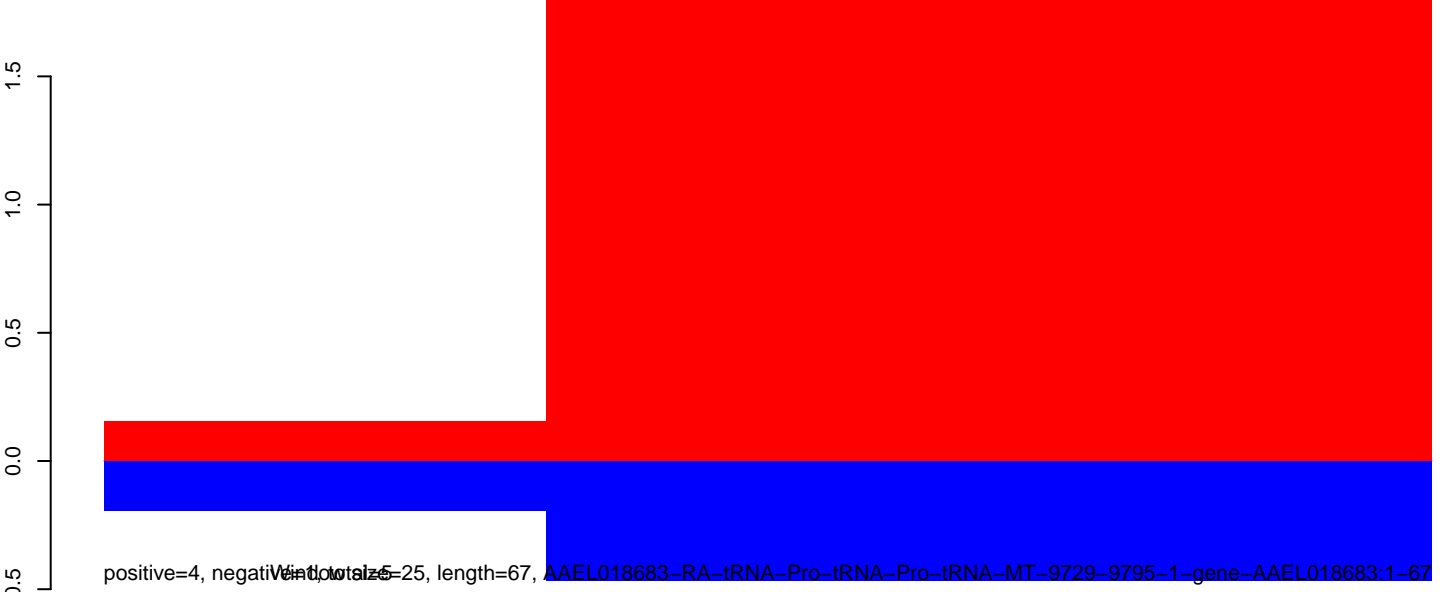
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AeAeg_Fcarc_GH_rep2.24_35.rep



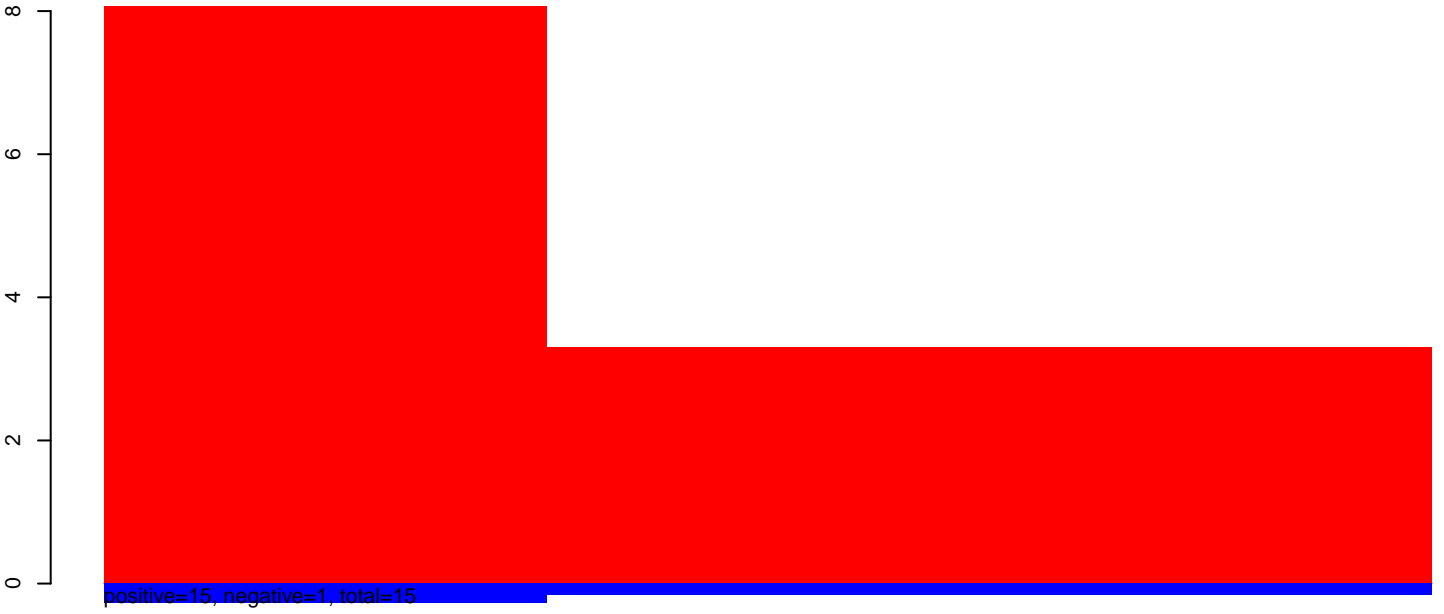
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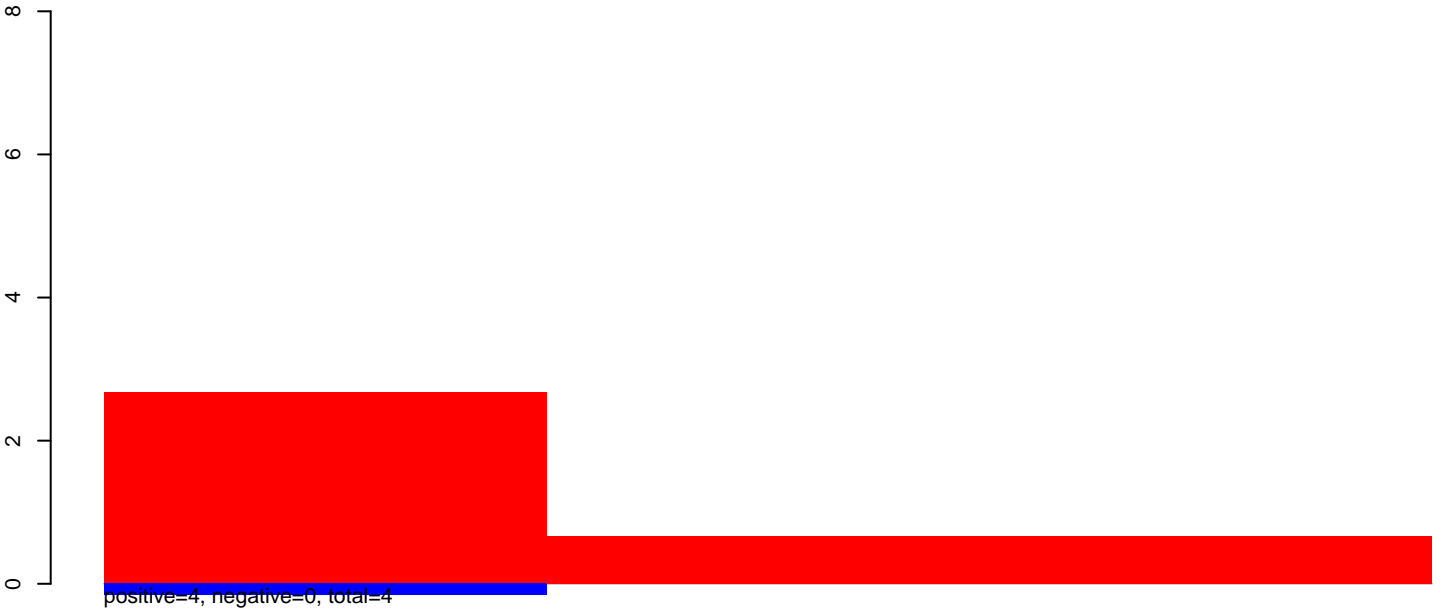
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40 60 80 100

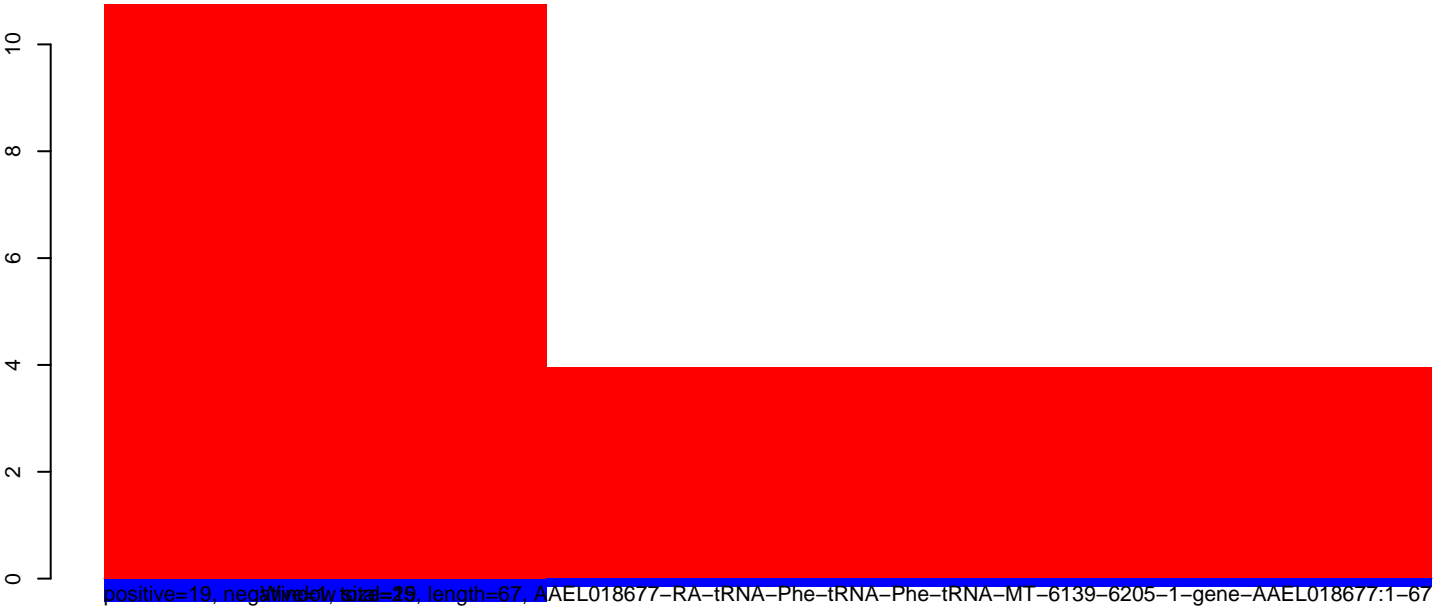
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AeAeg_Fcarc_GH_rep2.24_35.rep

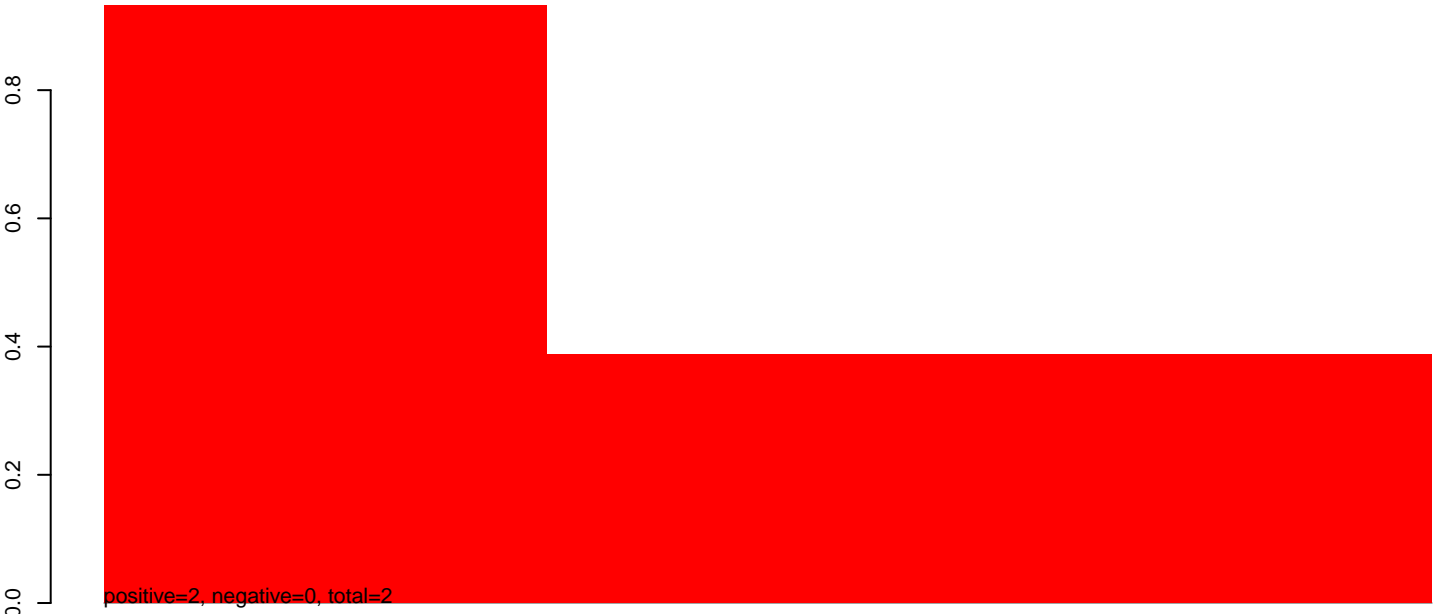


AeAeg_Fcarc_GH_rep2.rep



40 60 80 100

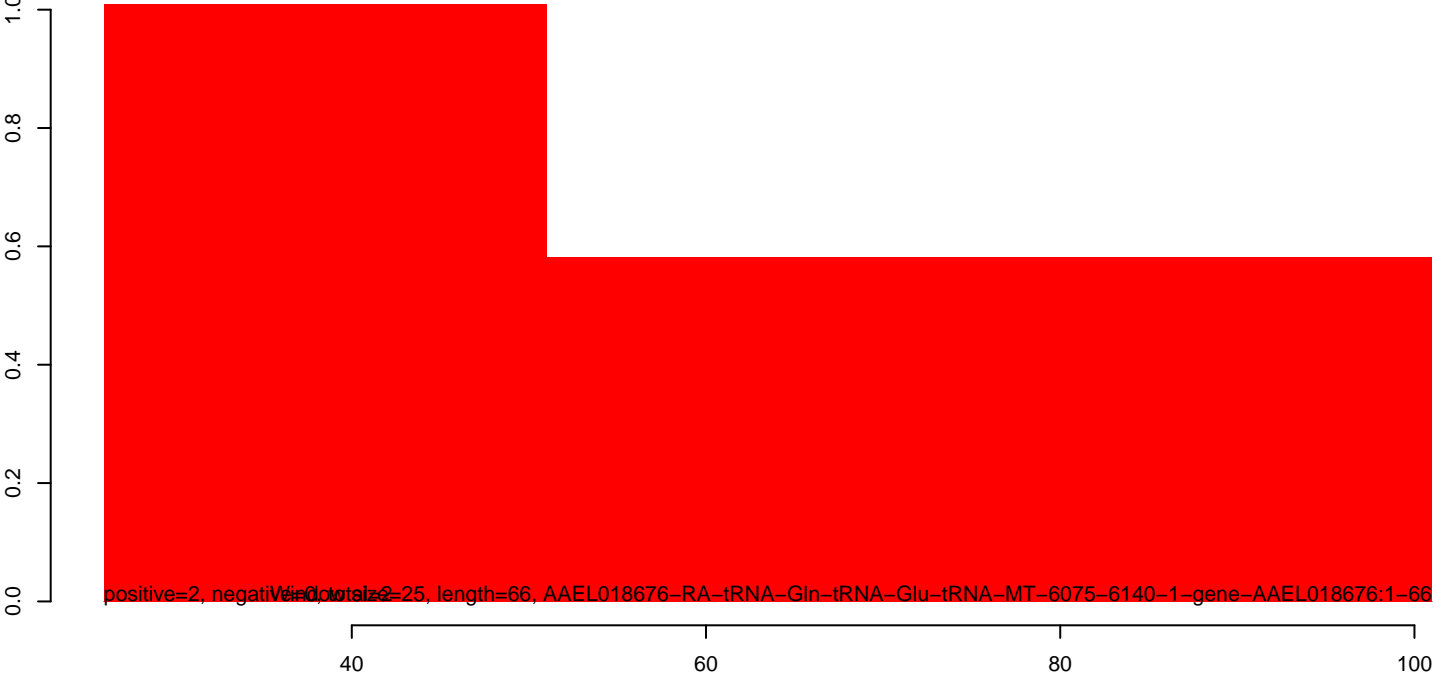
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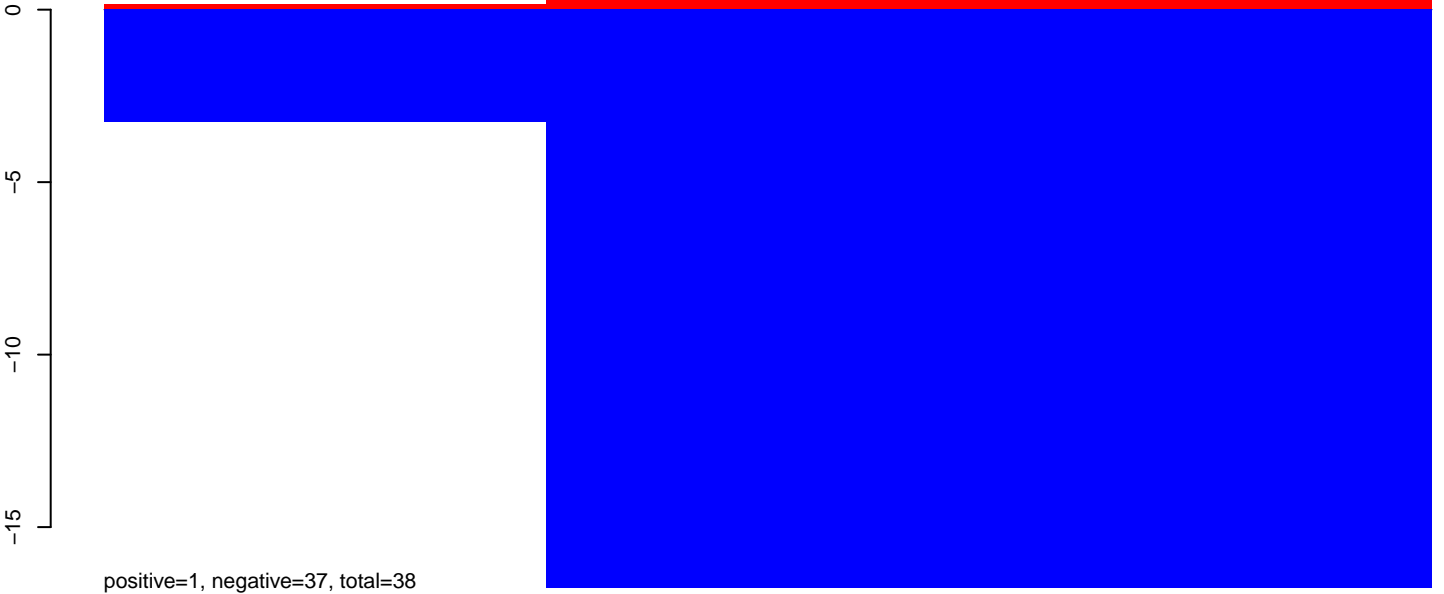
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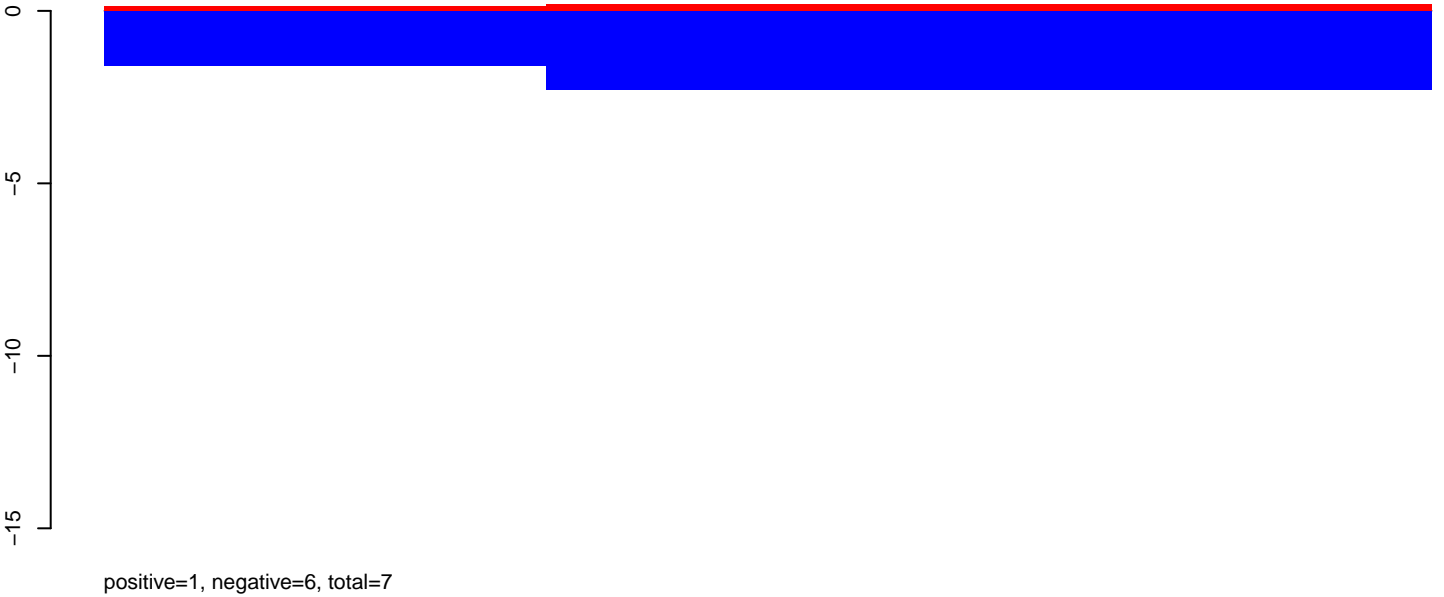
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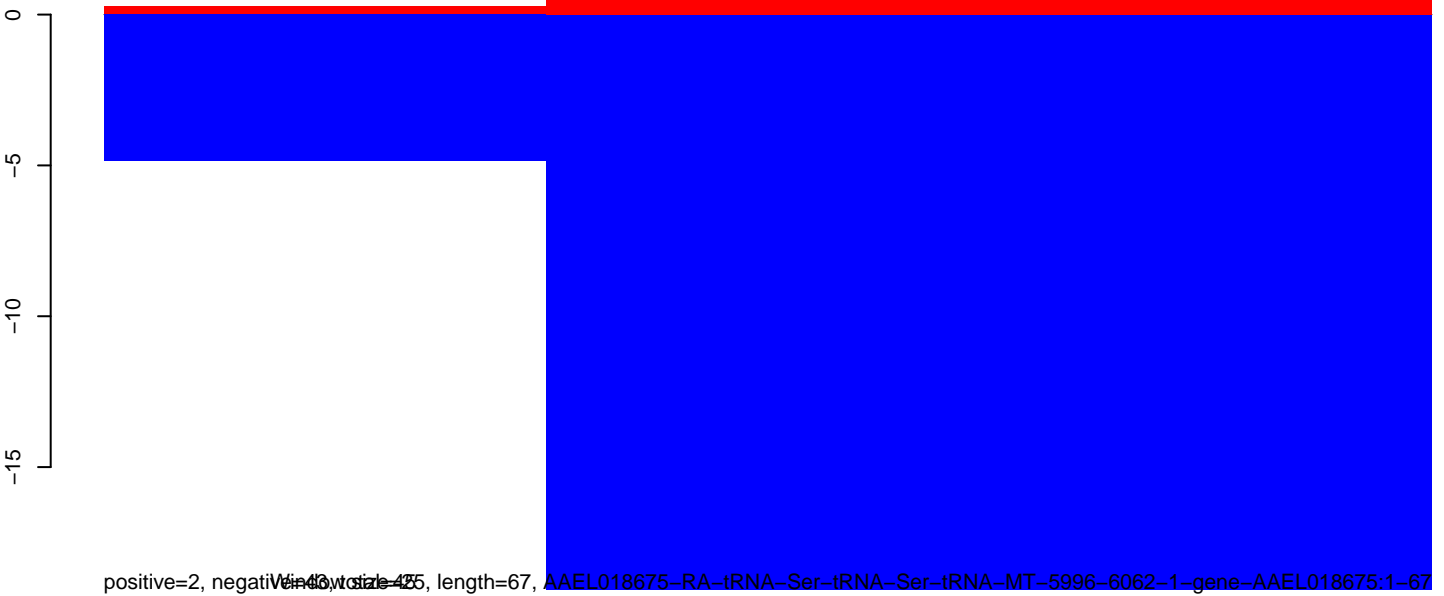
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AeAeg_Fcarc_GH_rep2.24_35.rep



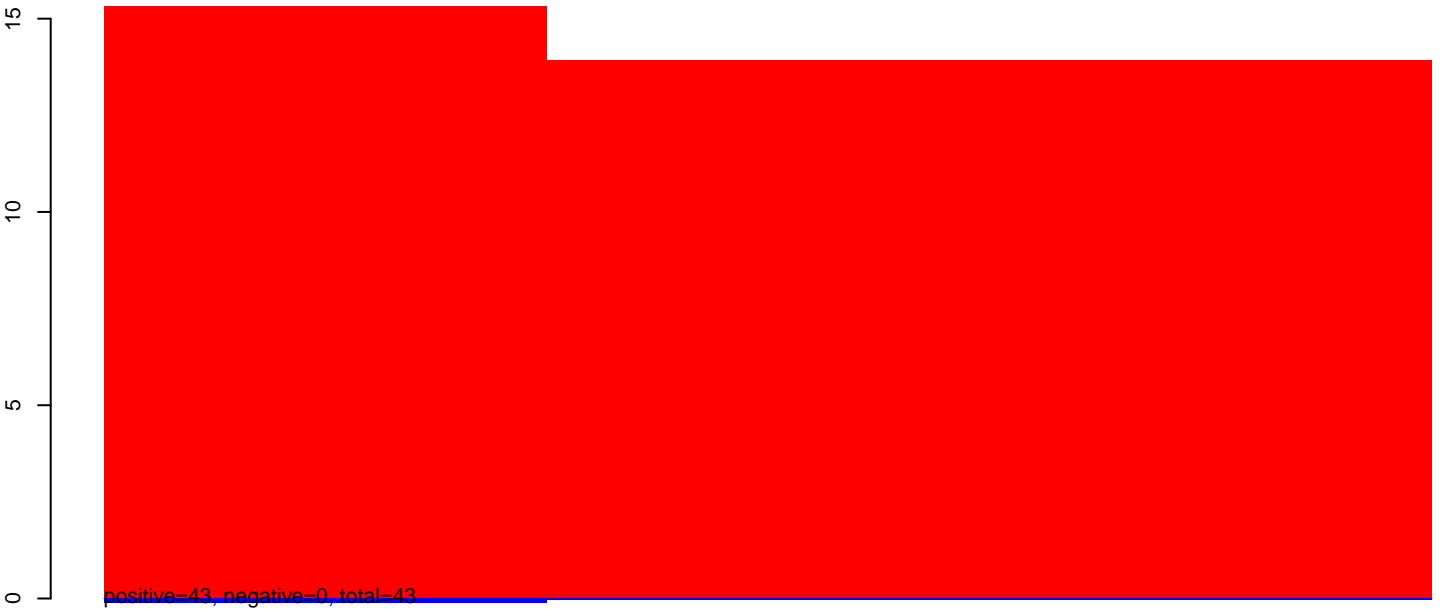
AeAeg_Fcarc_GH_rep2.rep



AAEL018675-RA-tRNA-Ser-tRNA-Ser-tRNA-MT-5996-6062-1-gene-AAEL018675:1-67

40 60 80 100

AeAeg_Fcarc_GH_rep2.18_23.rep



AeAeg_Fcarc_GH_rep2.24_35.rep

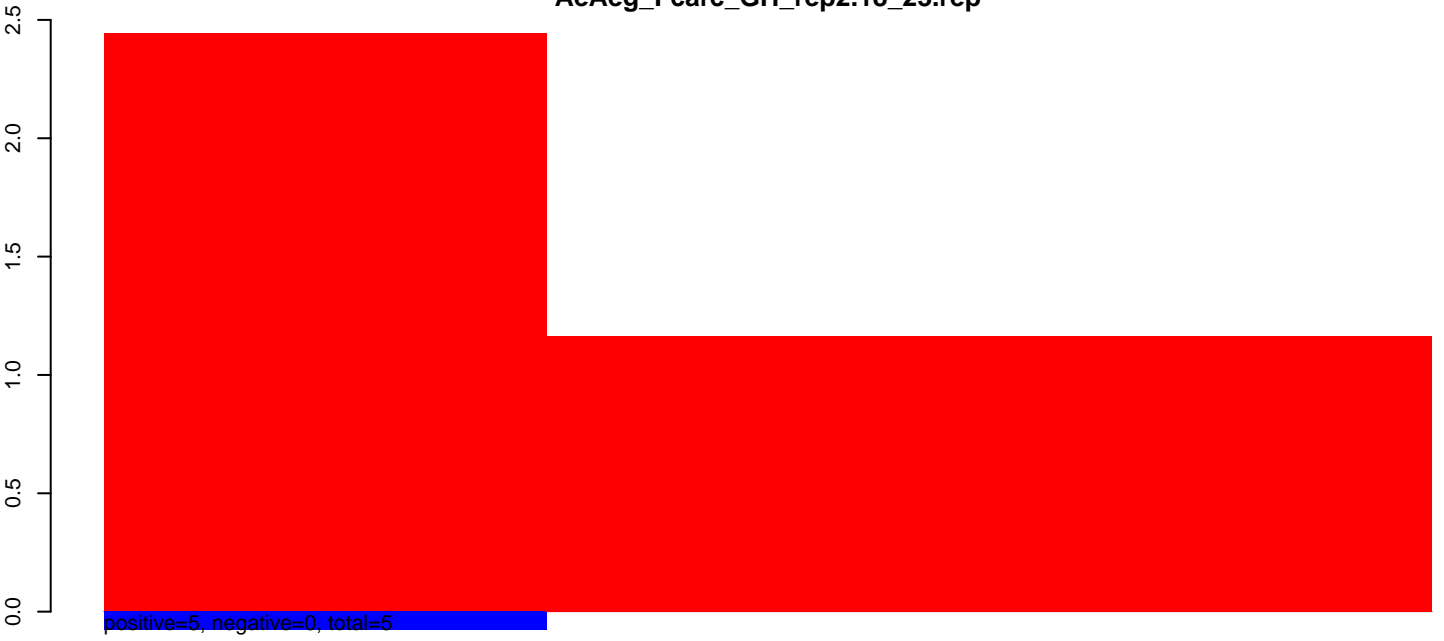


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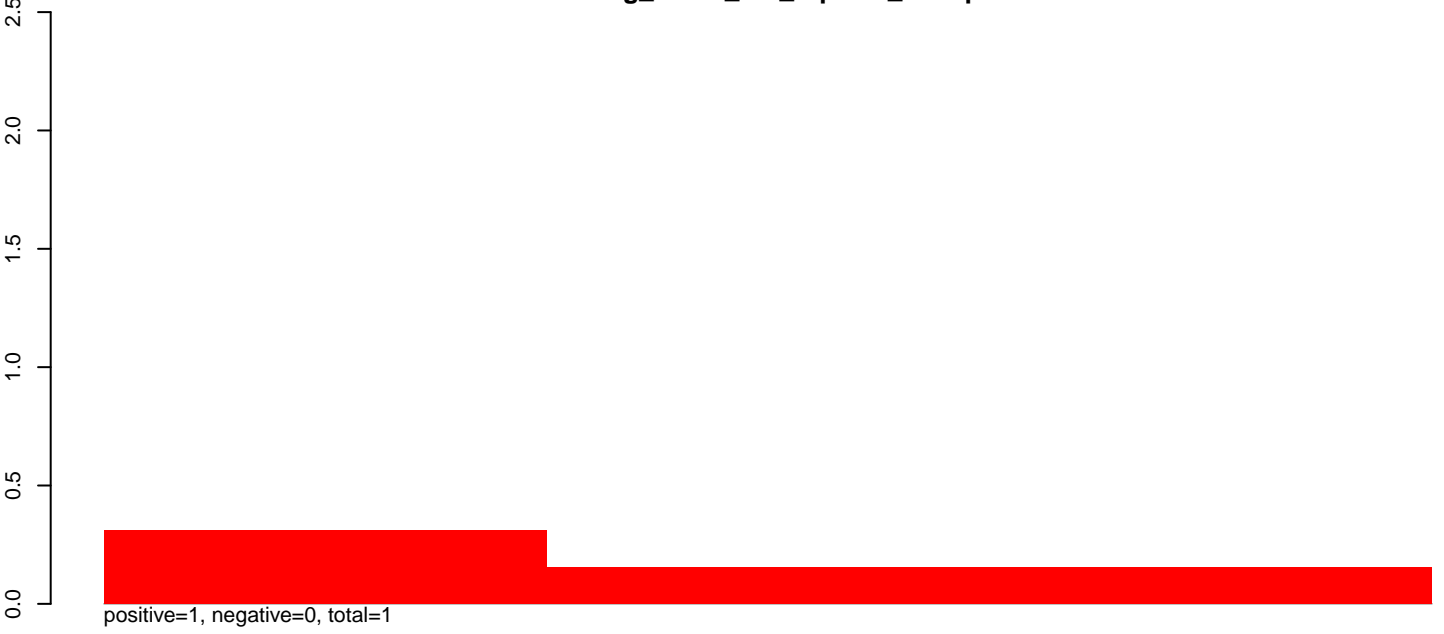


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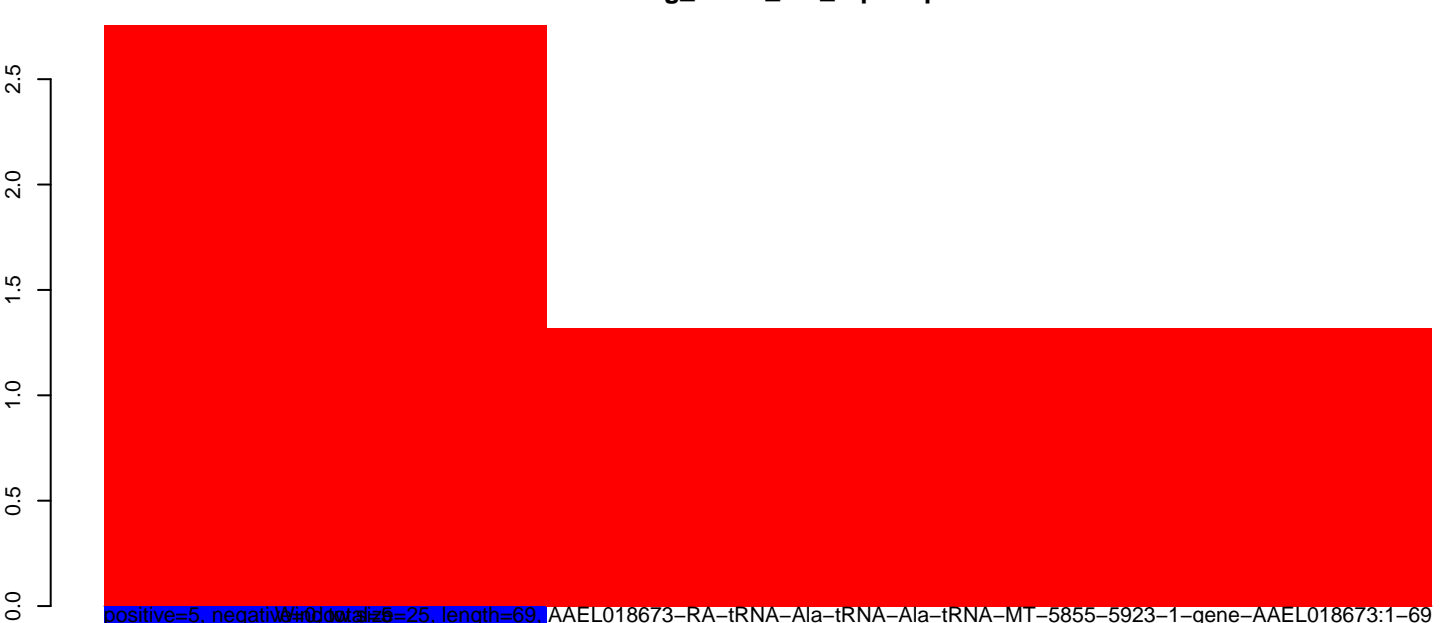
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AeAeg_Fcarc_GH_rep2.24_35.rep

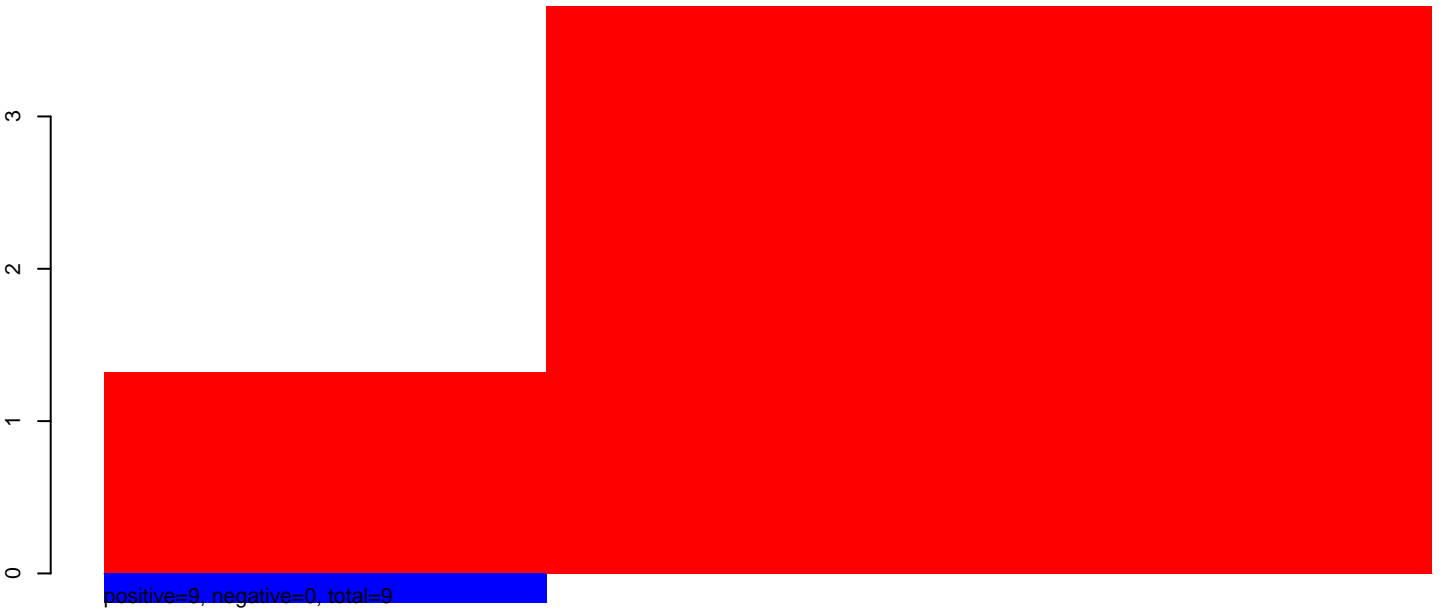


AeAeg_Fcarc_GH_rep2.rep



40 60 80 100

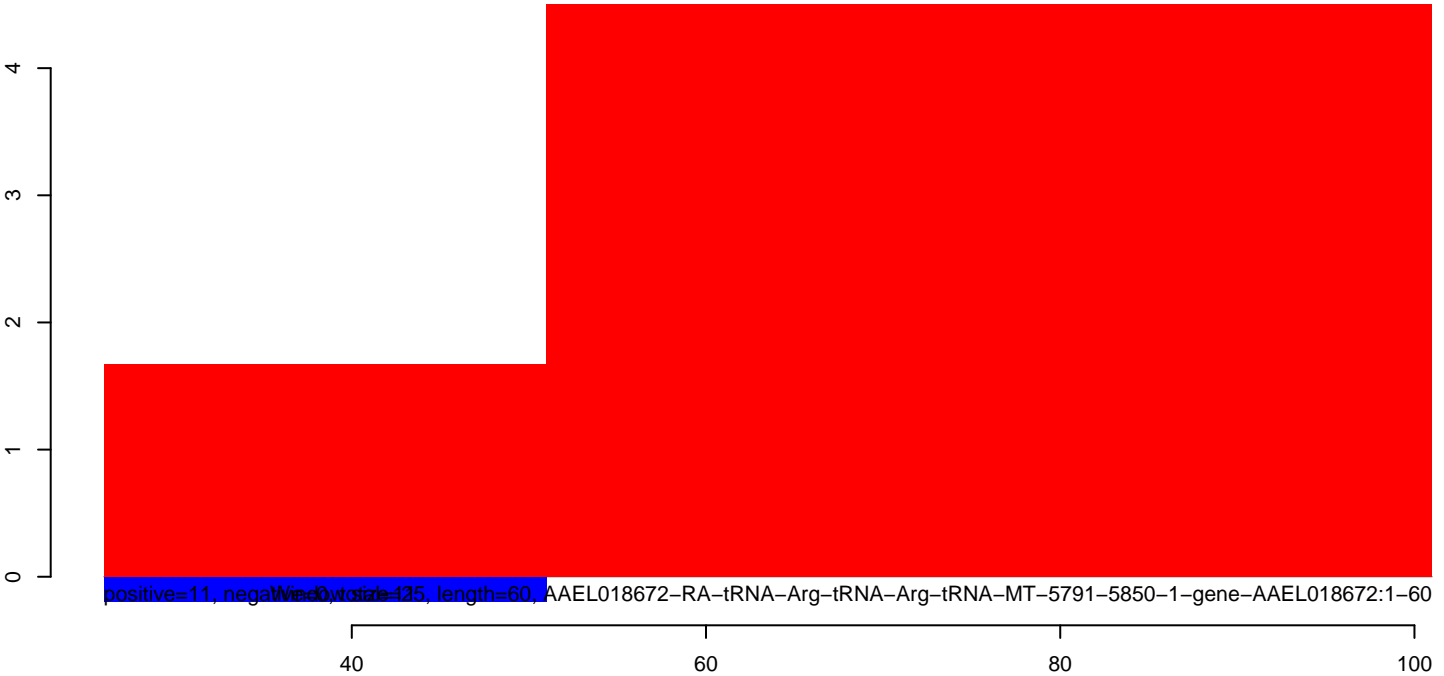
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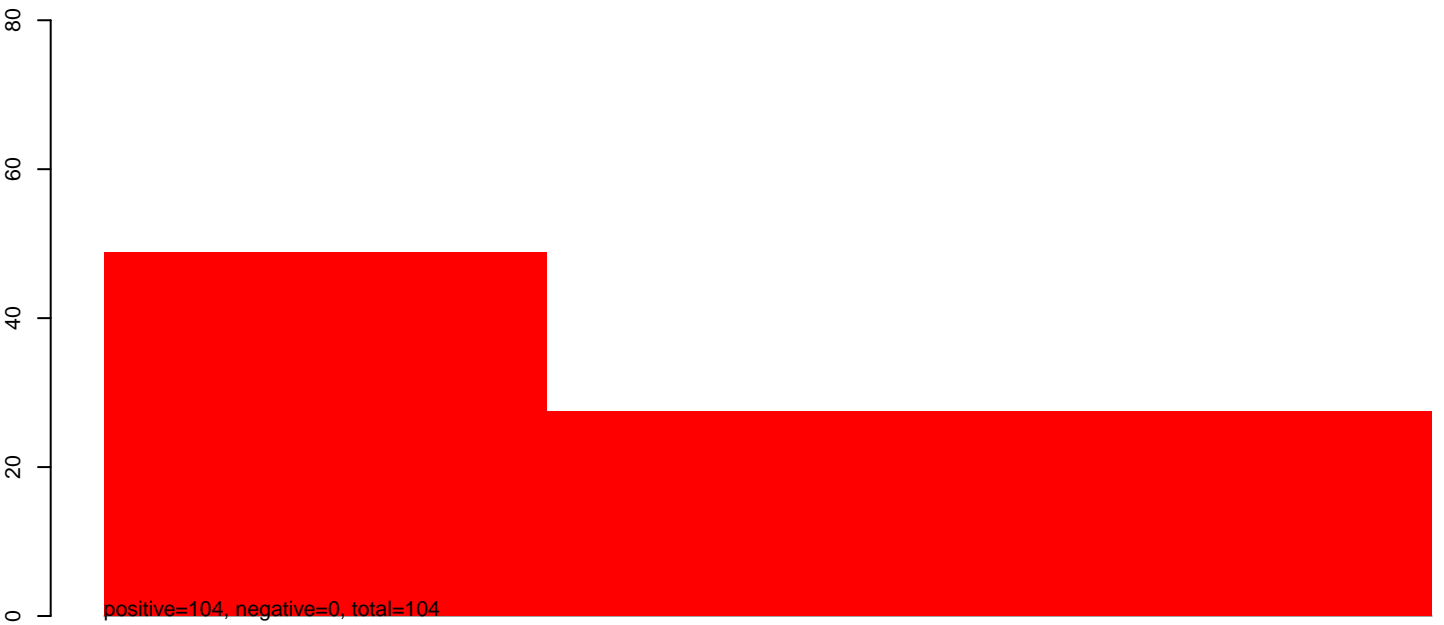
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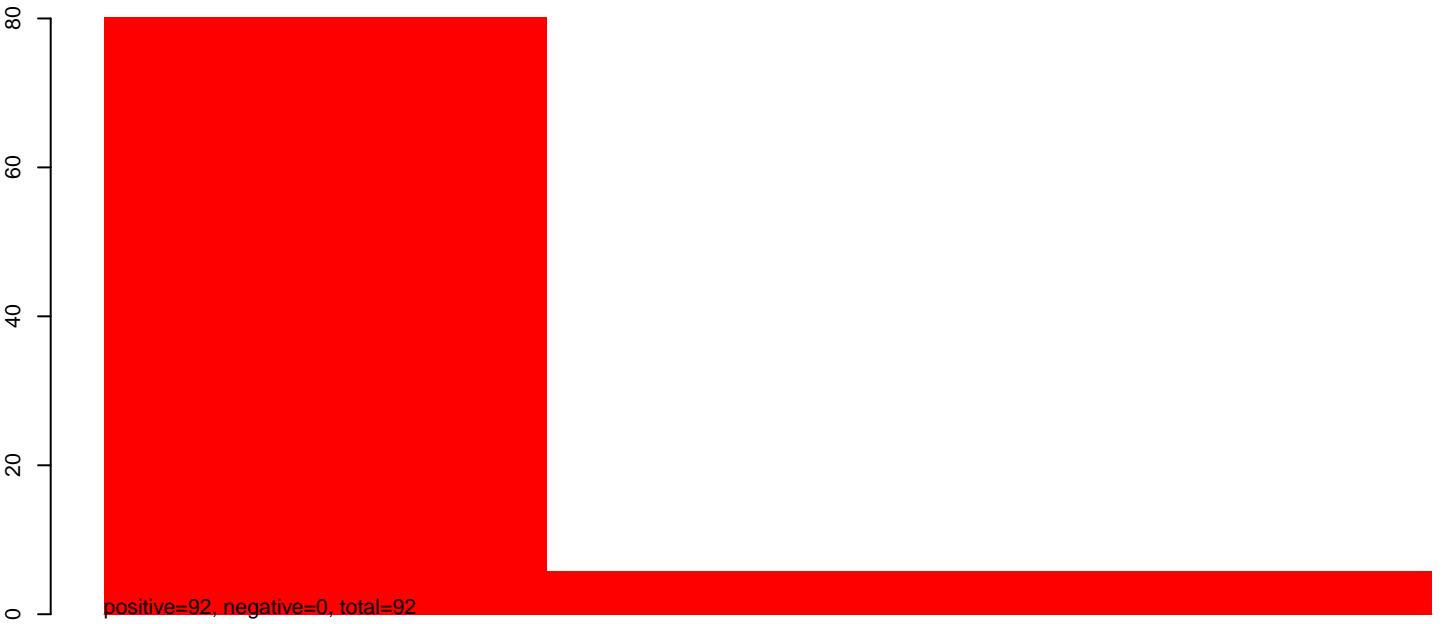
AeAeg_Fcarc_GH_rep2.rep



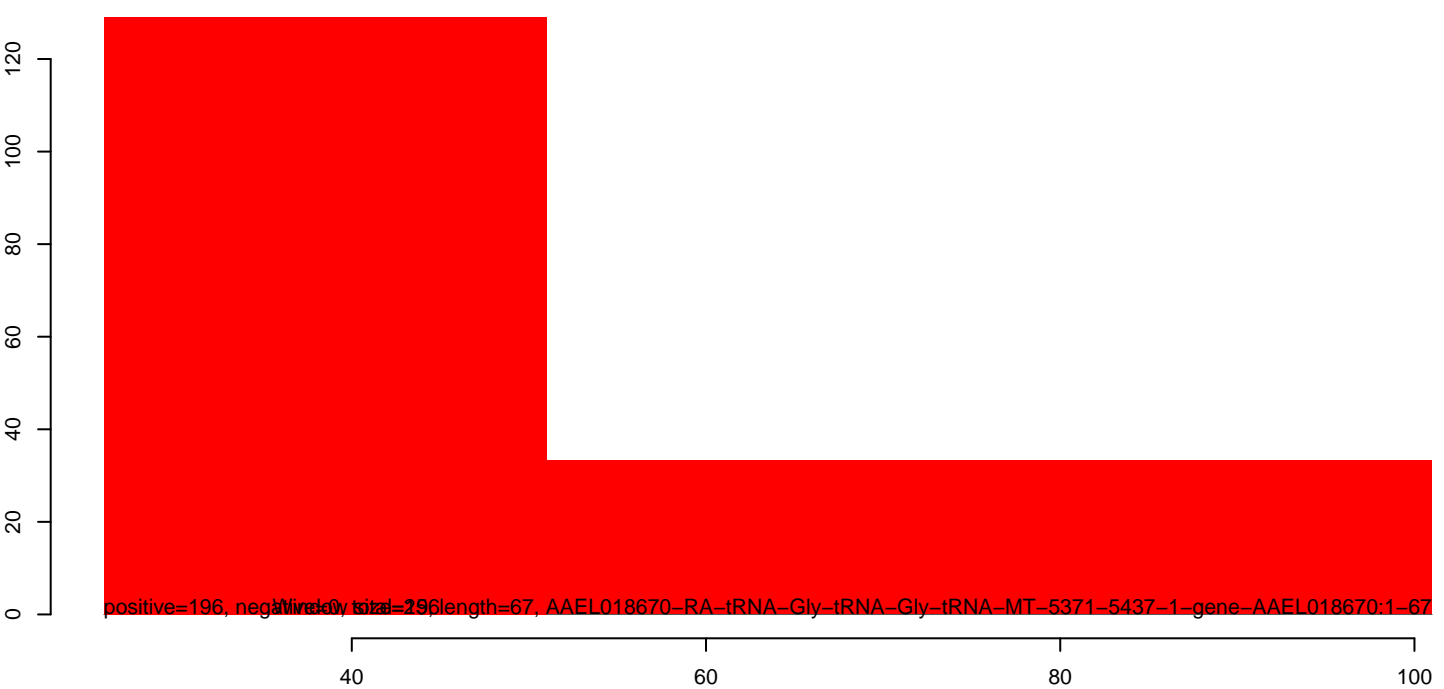
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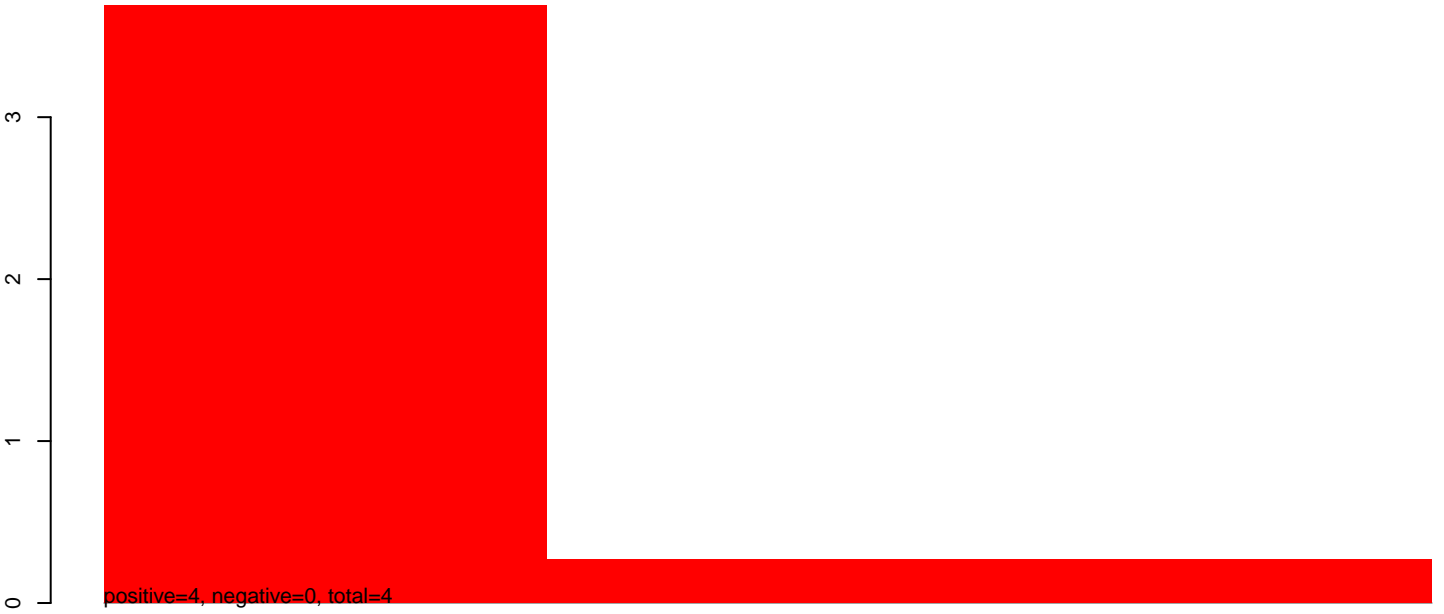
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AeAeg_Fcarc_GH_rep2.rep



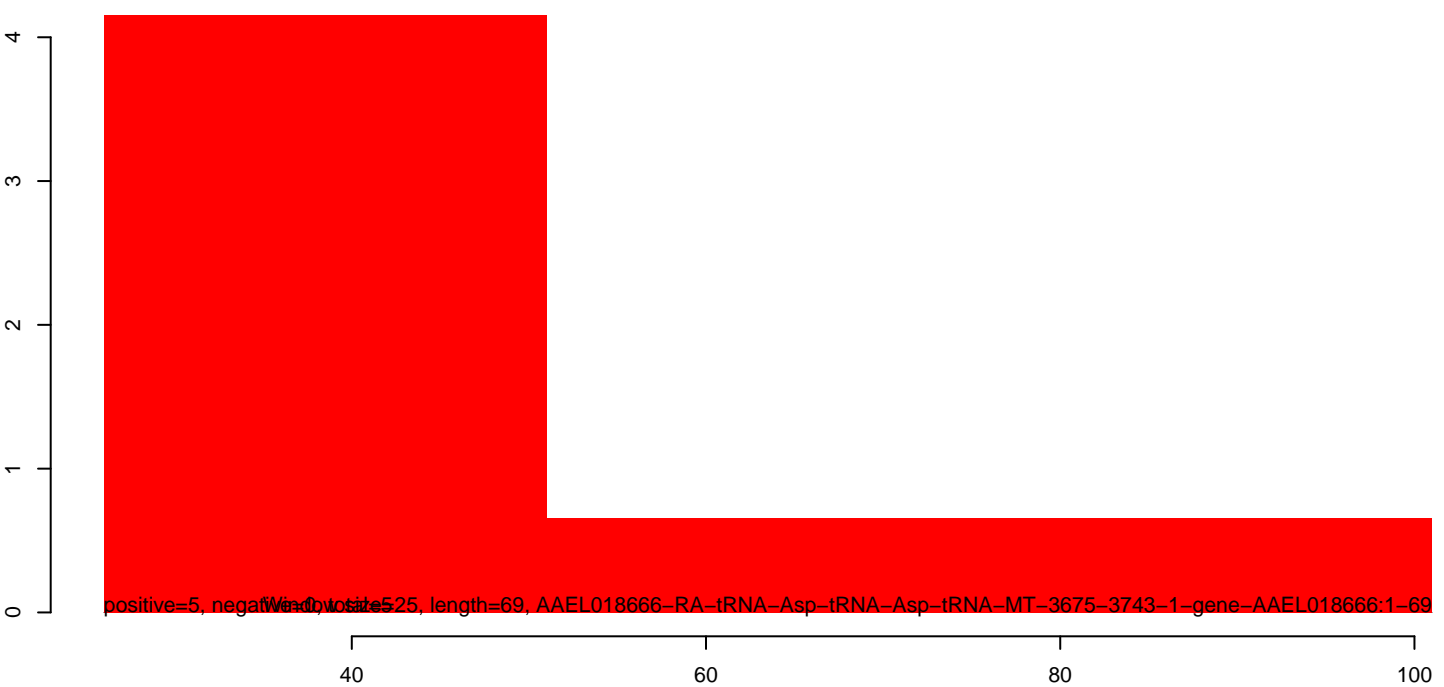
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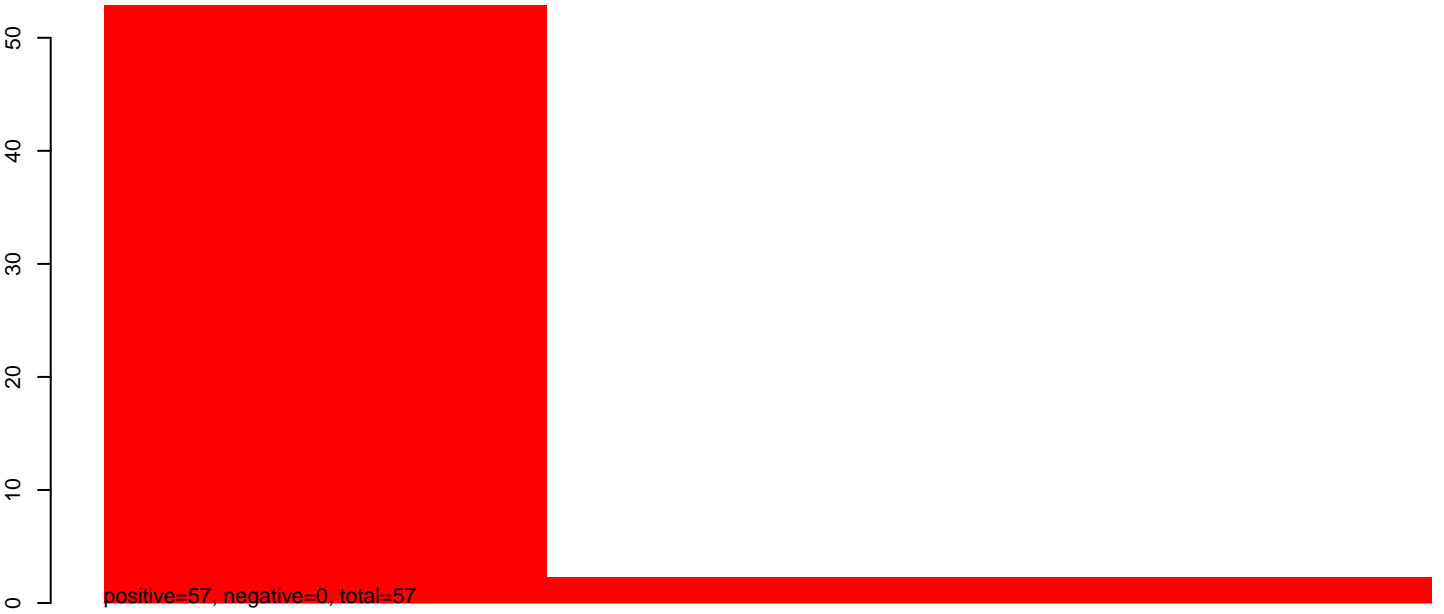
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AeAeg_Fcarc_GH_rep2.rep



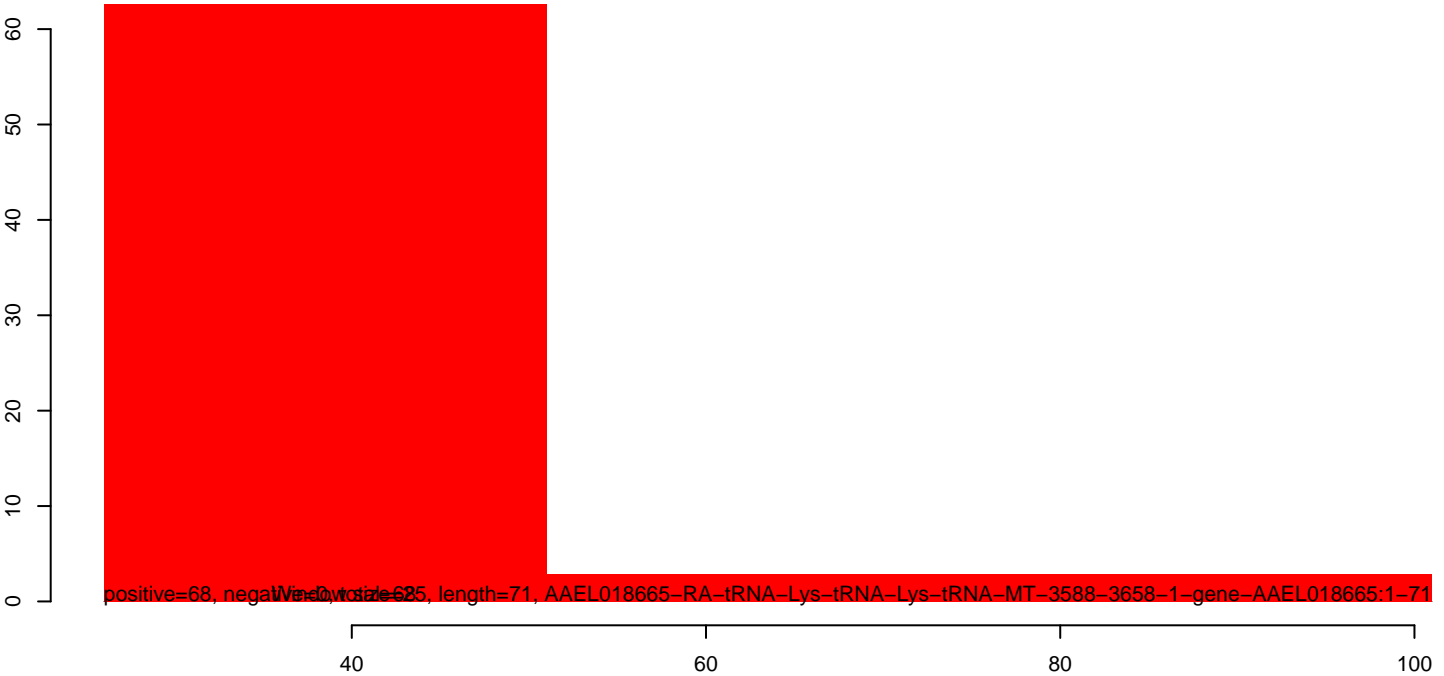
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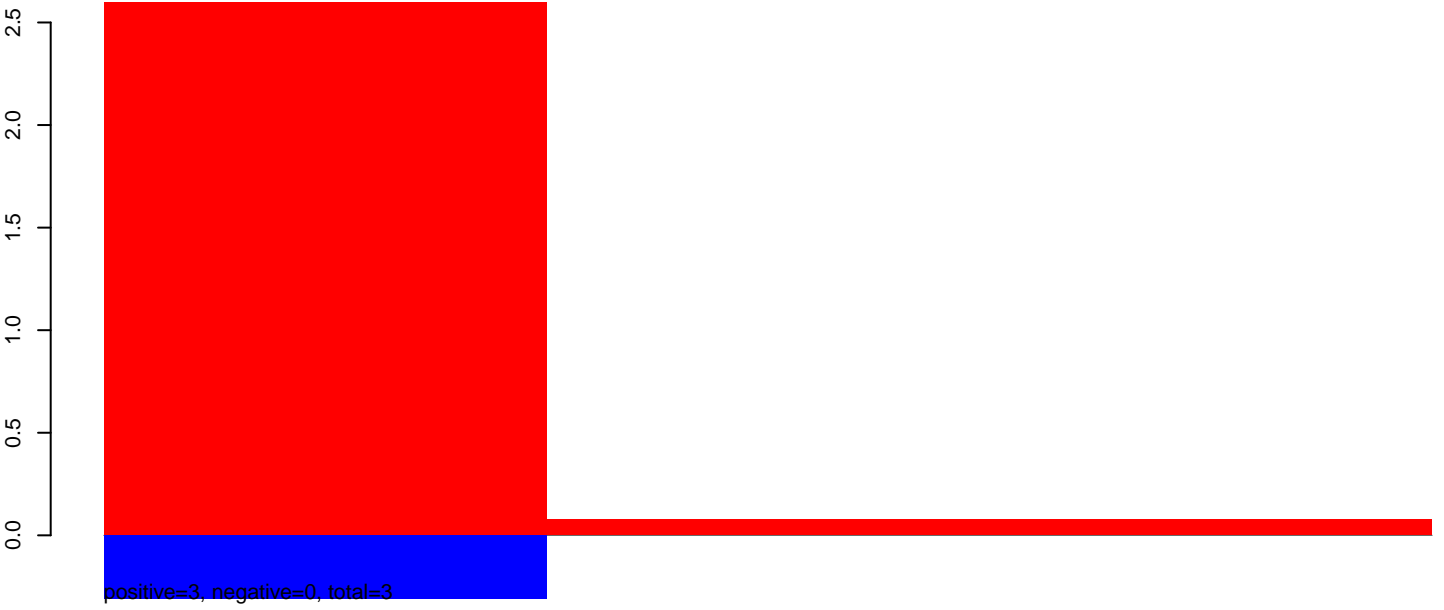
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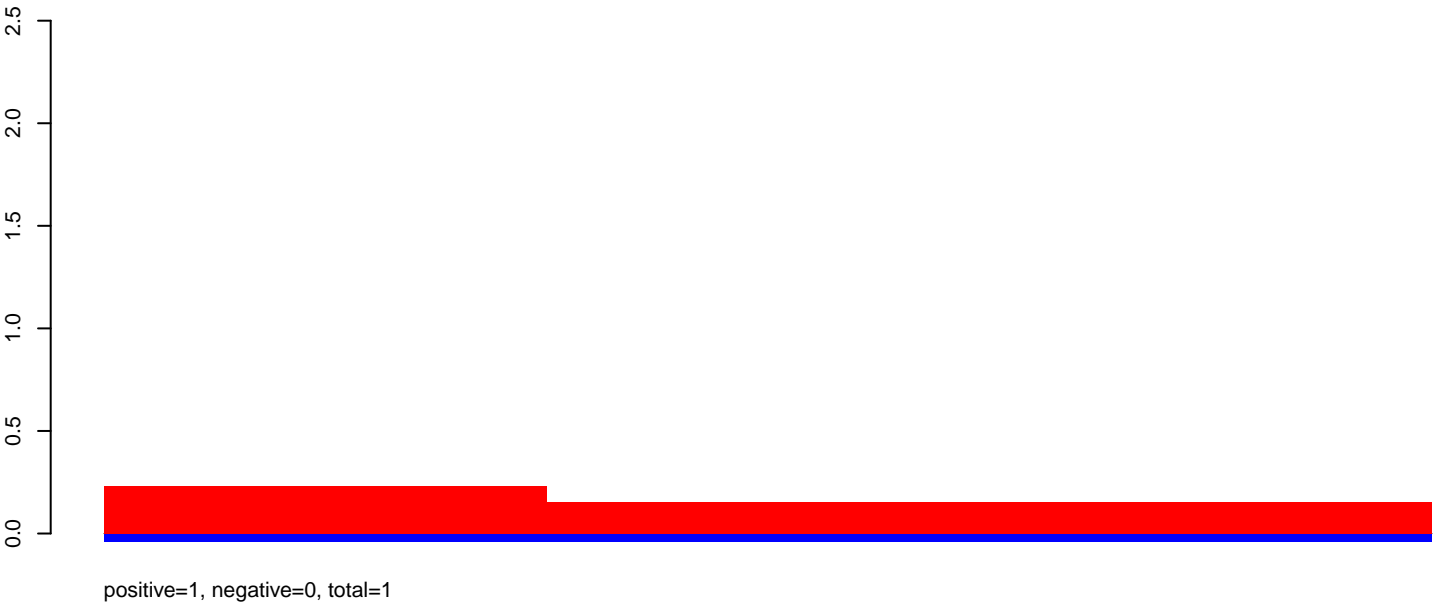
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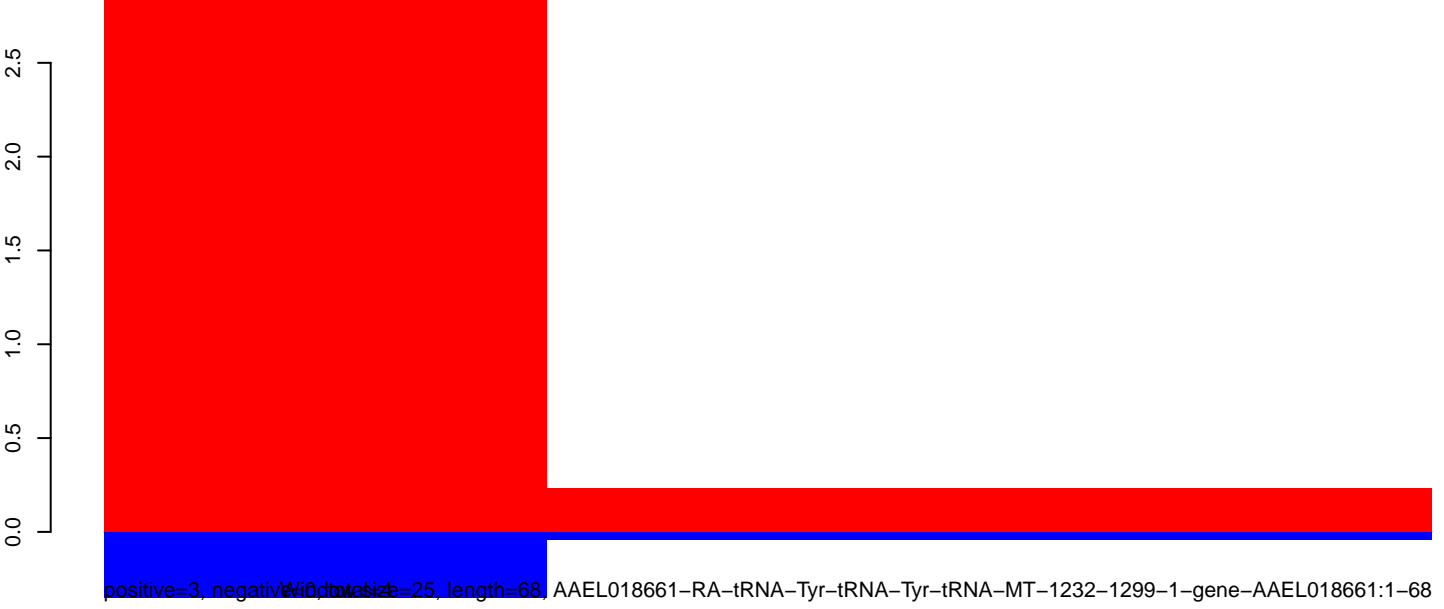
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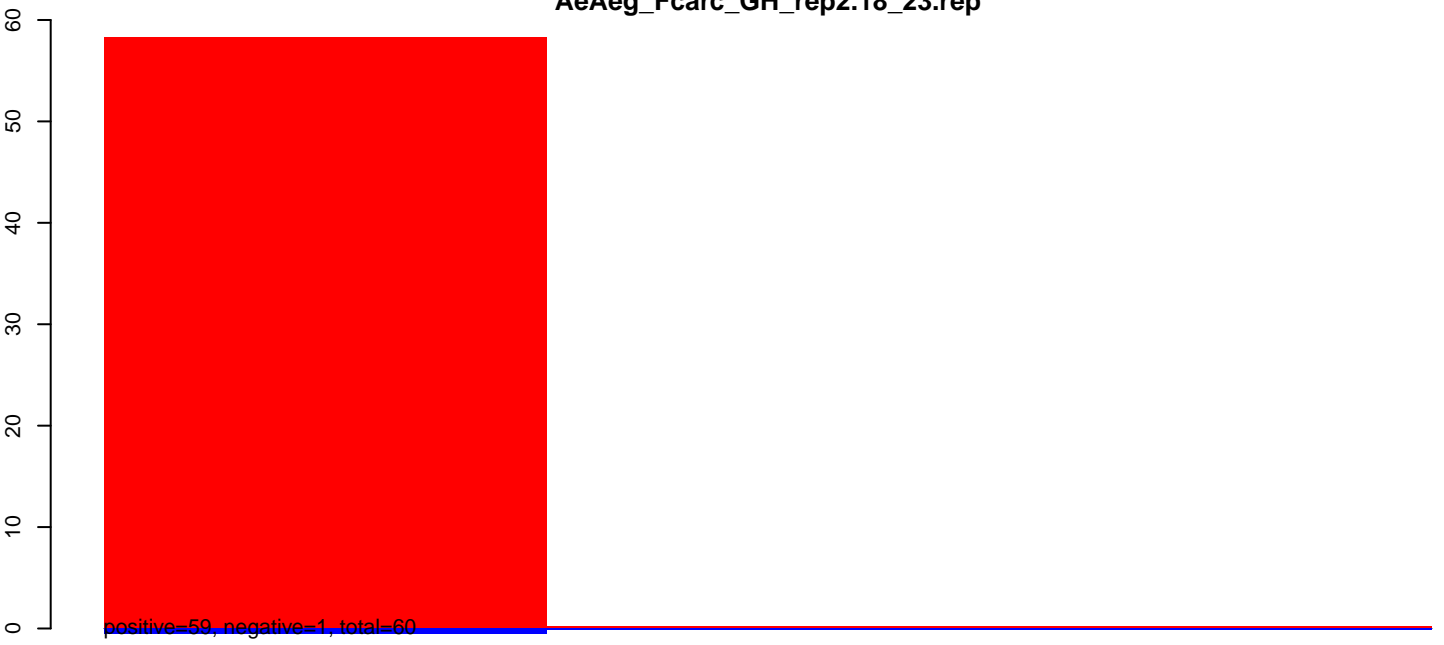


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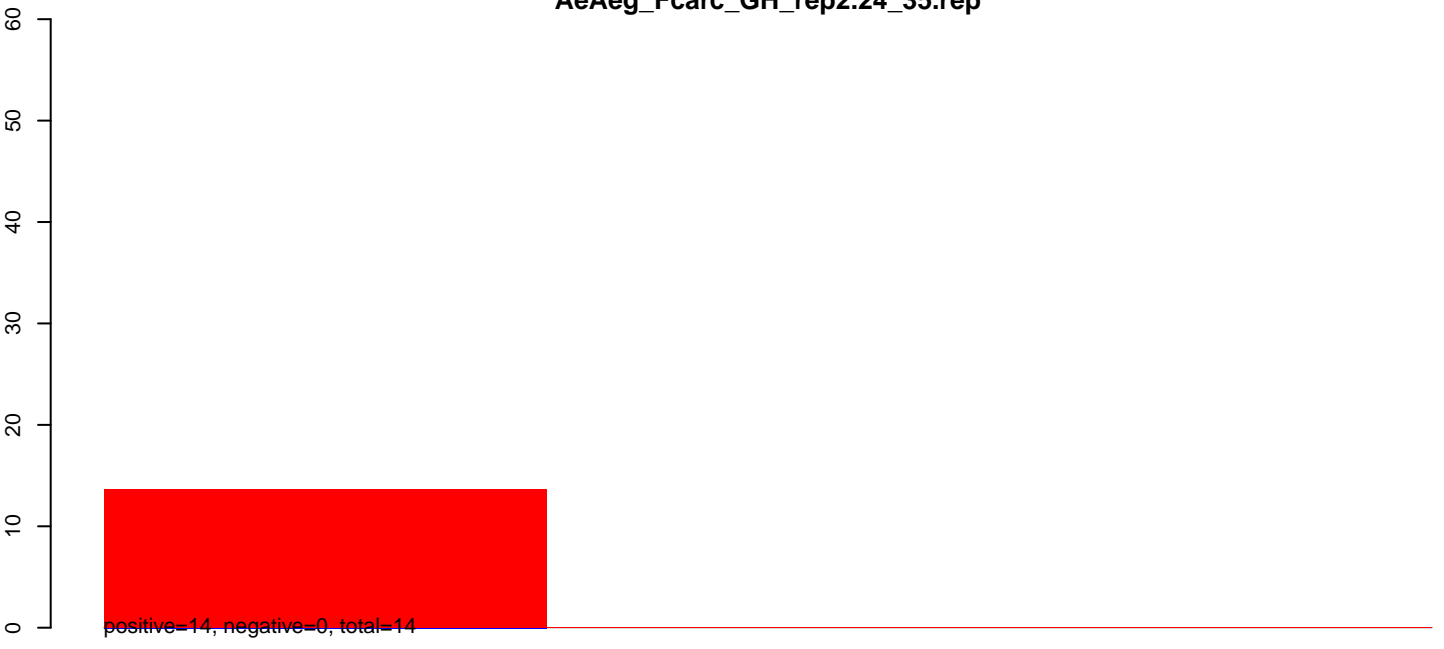


40 60 80 100
AAEL018661-RA-tRNA-Tyr-tRNA-Tyr-tRNA-MT-1232-1299-1-gene-AAEL018661:1-68

AeAeg_Fcarc_GH_rep2.18_23.rep



AeAeg_Fcarc_GH_rep2.24_35.rep



AeAeg_Fcarc_GH_rep2.rep



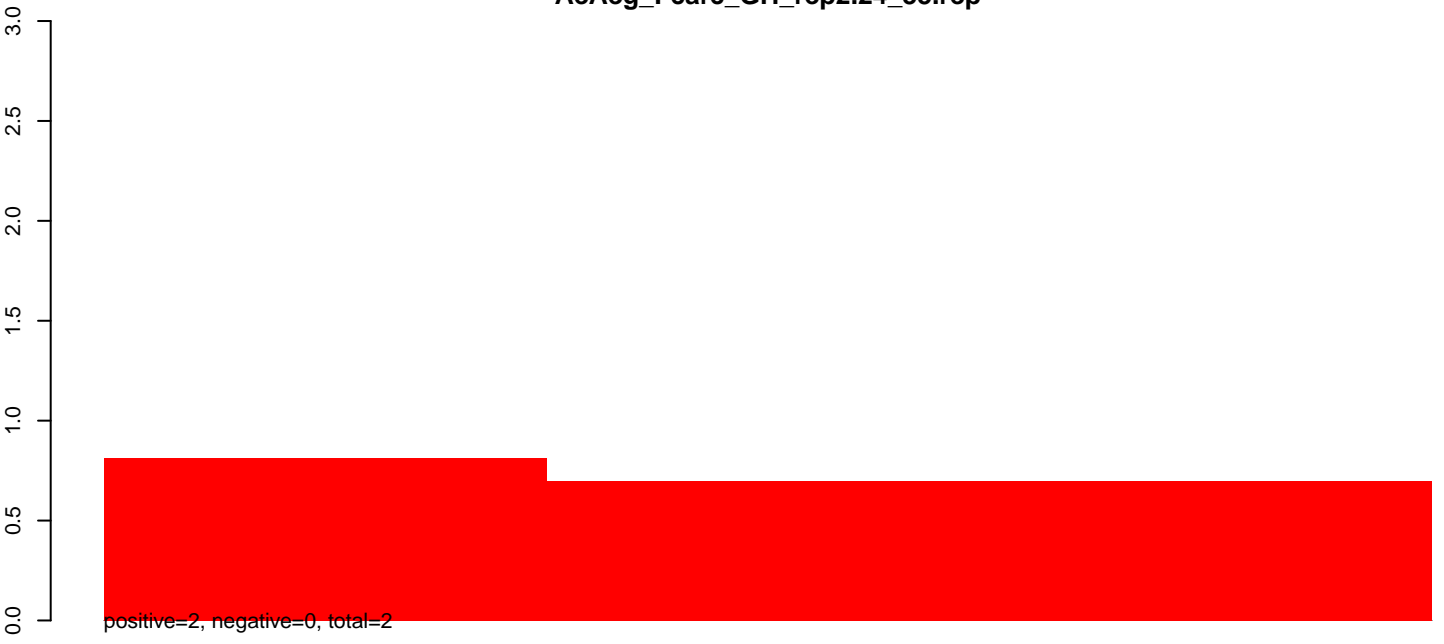
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40 60 80 100

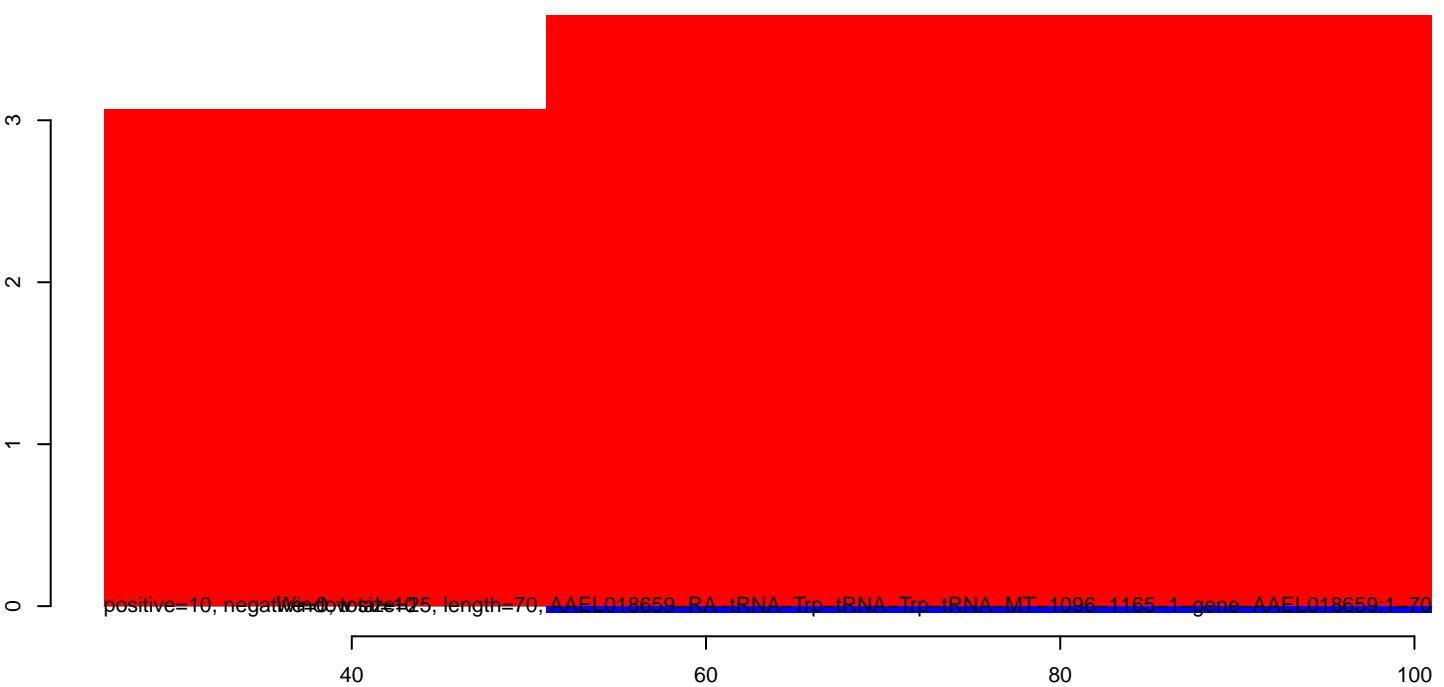
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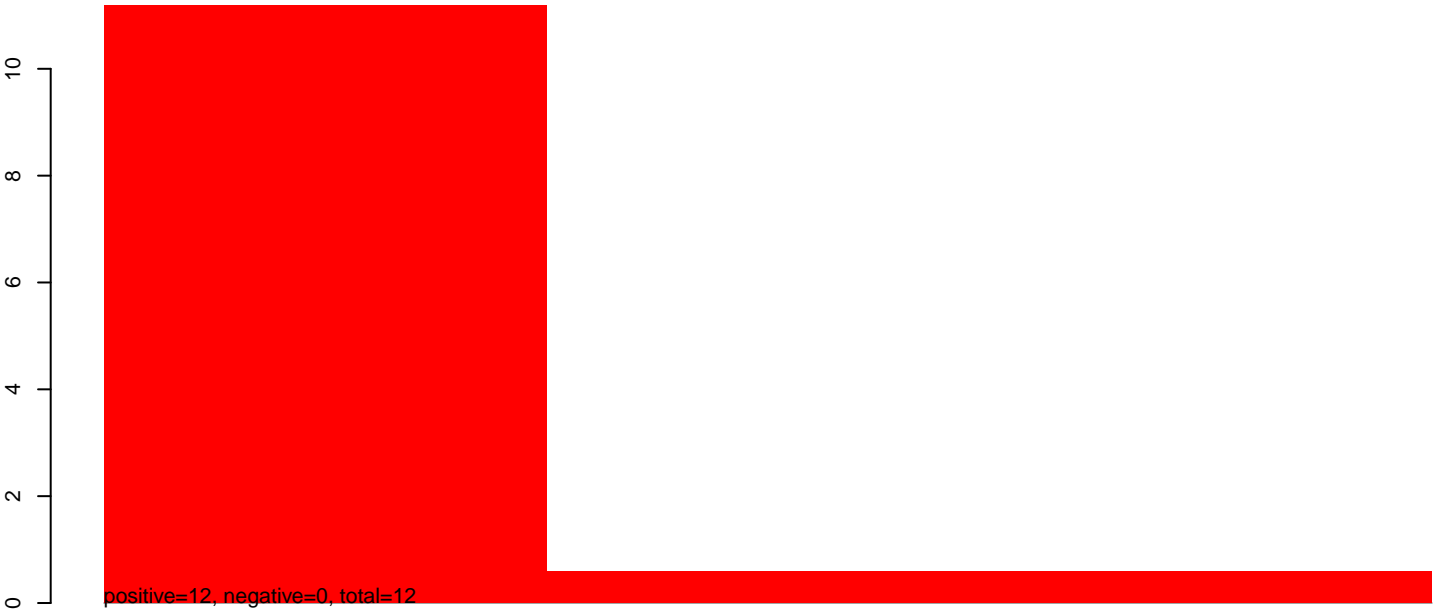
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AeAeg_FcArc_GH_rep2.rep



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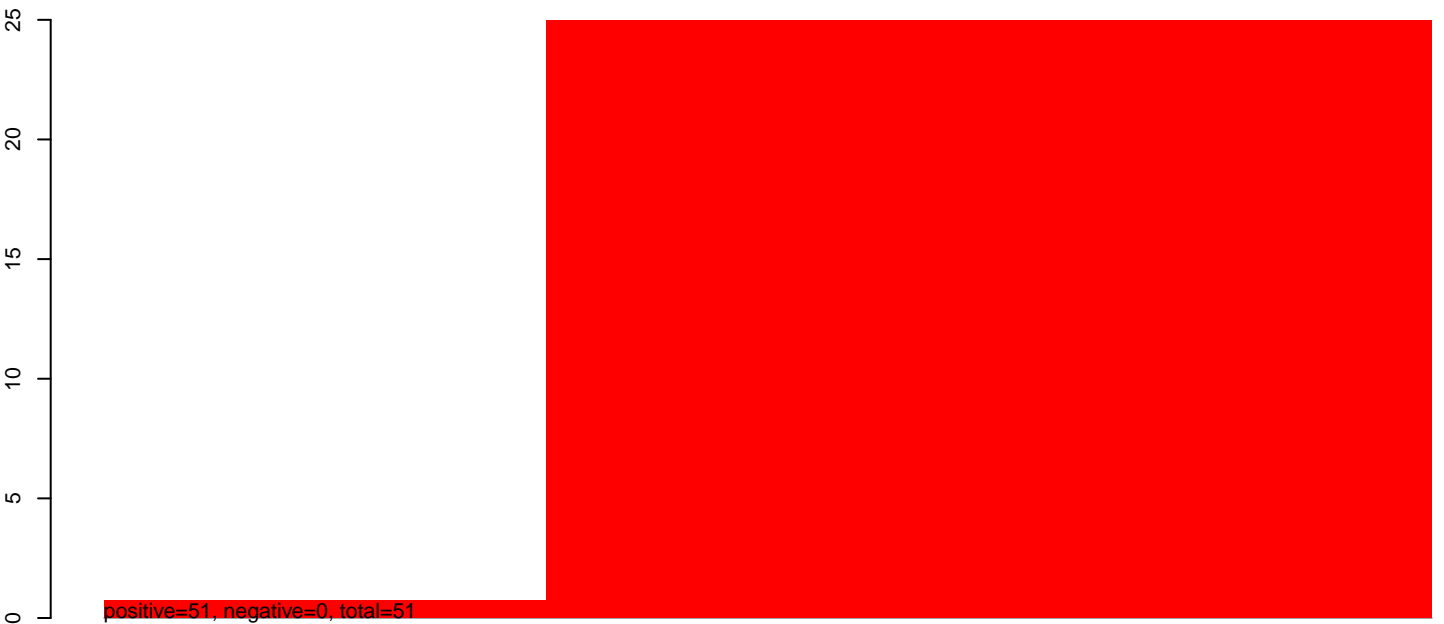
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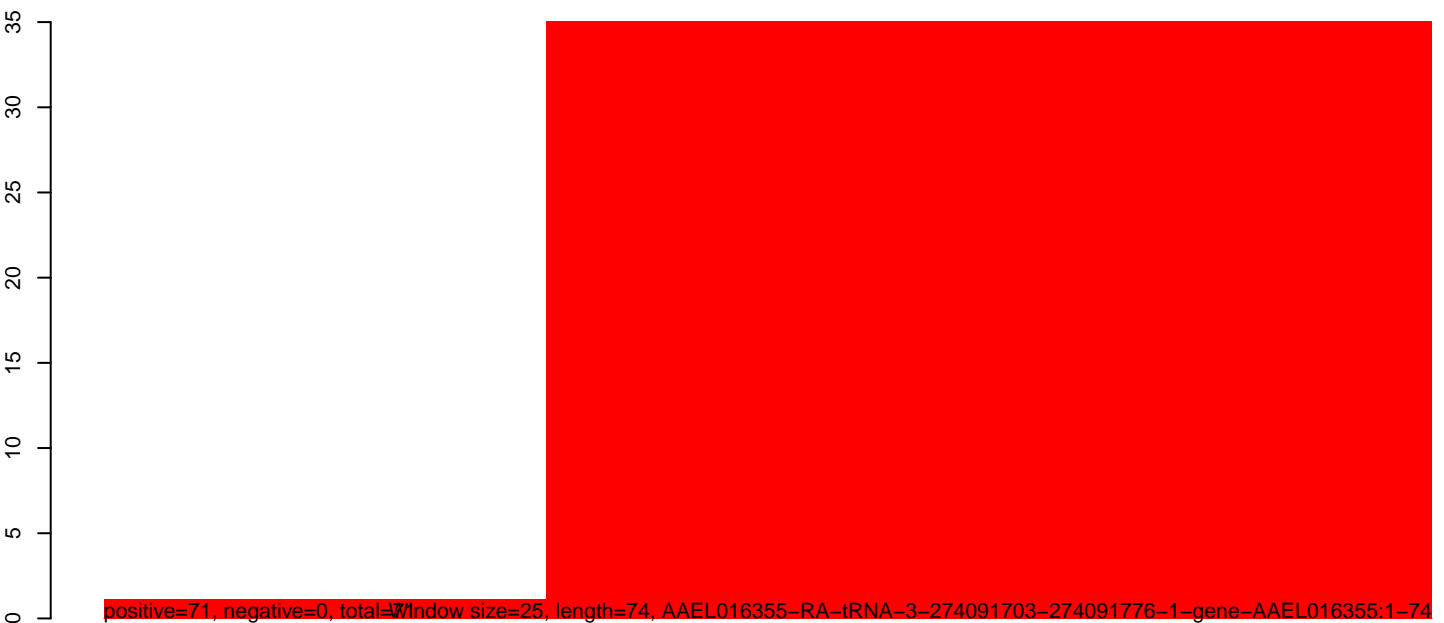
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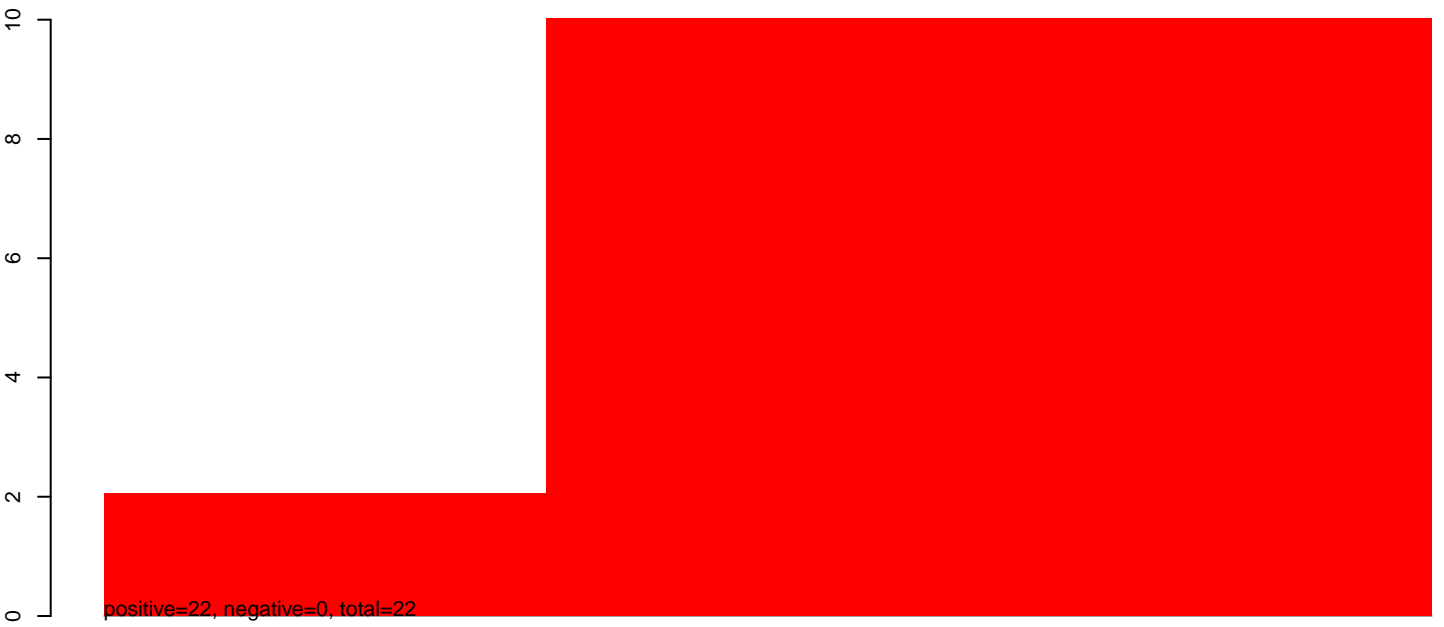


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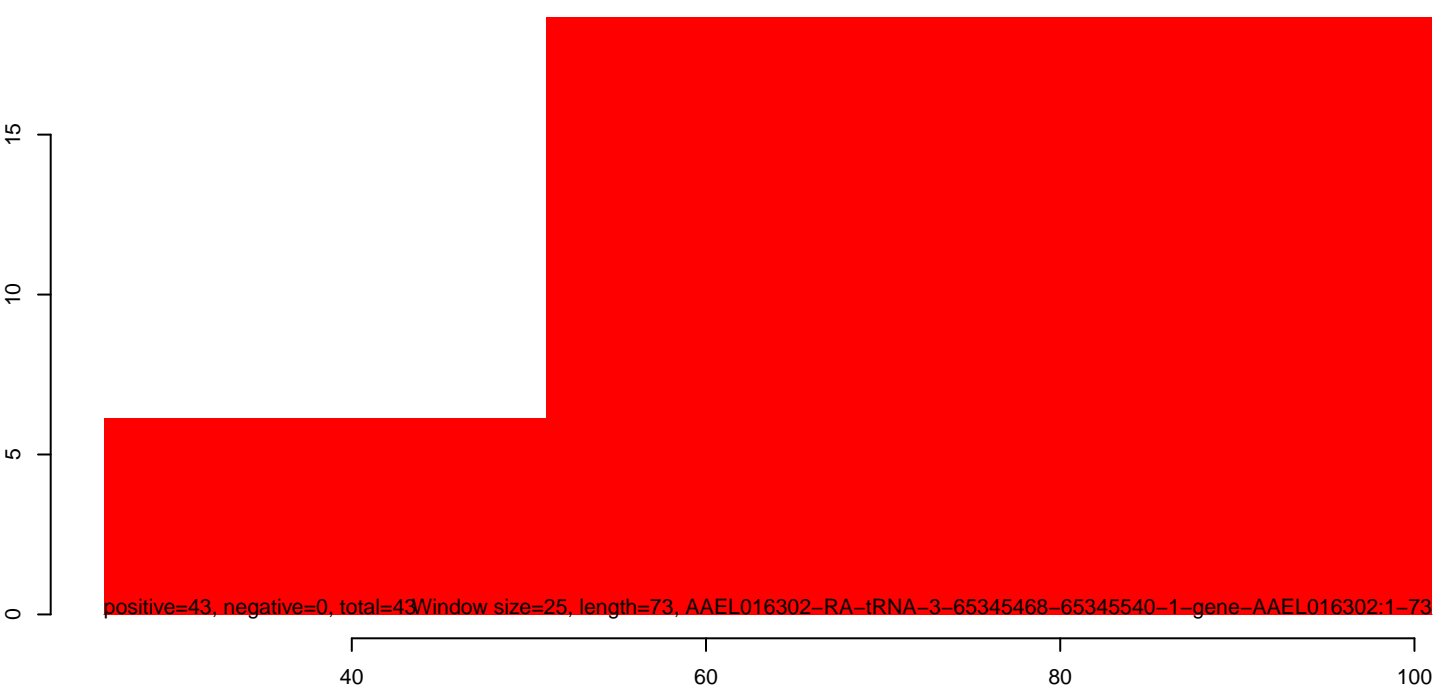
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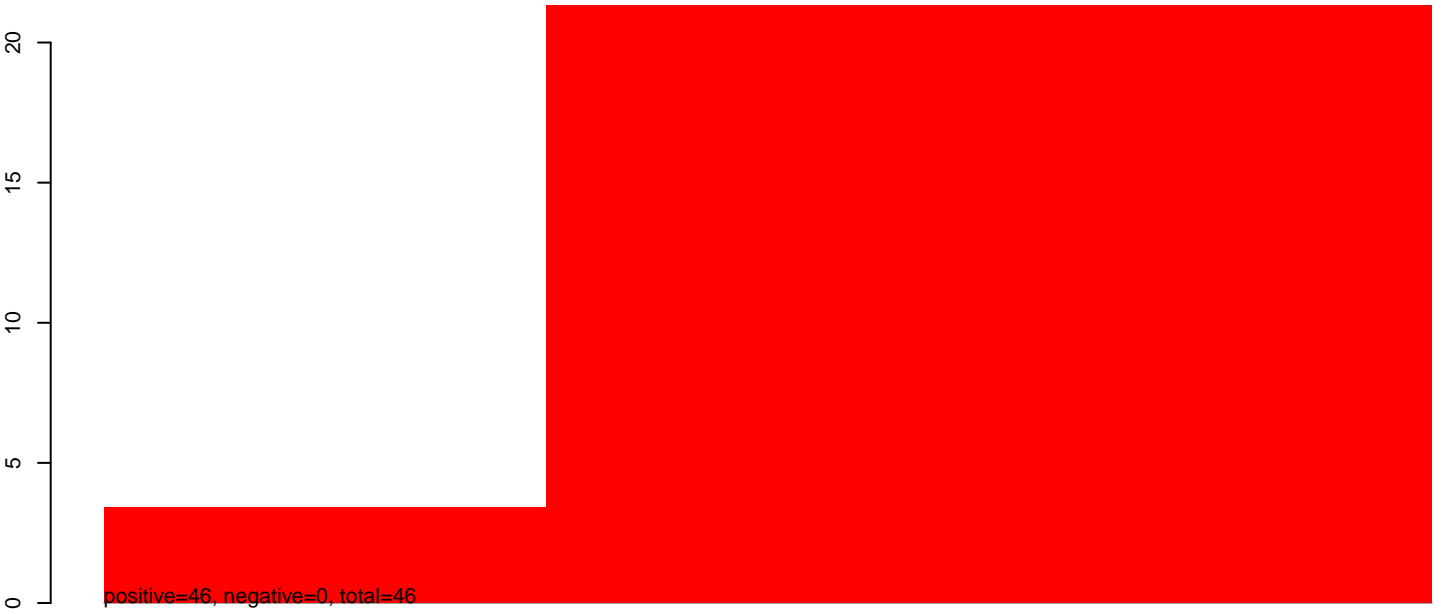
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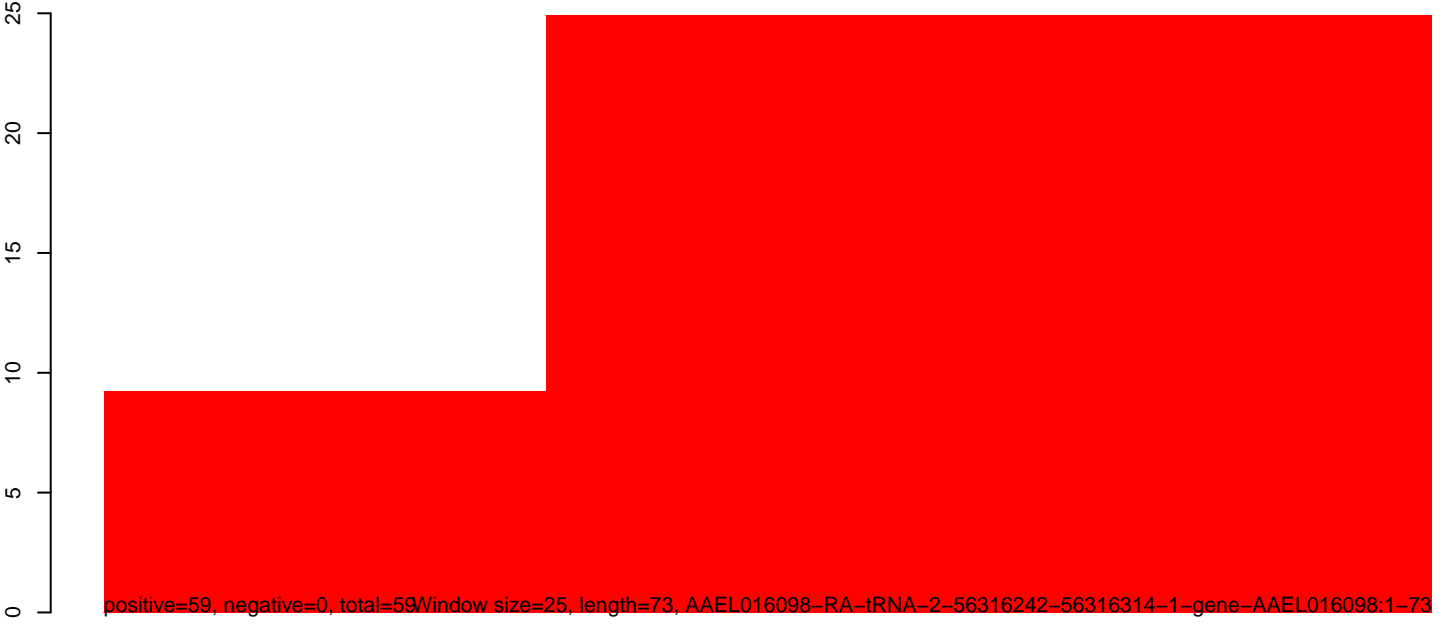
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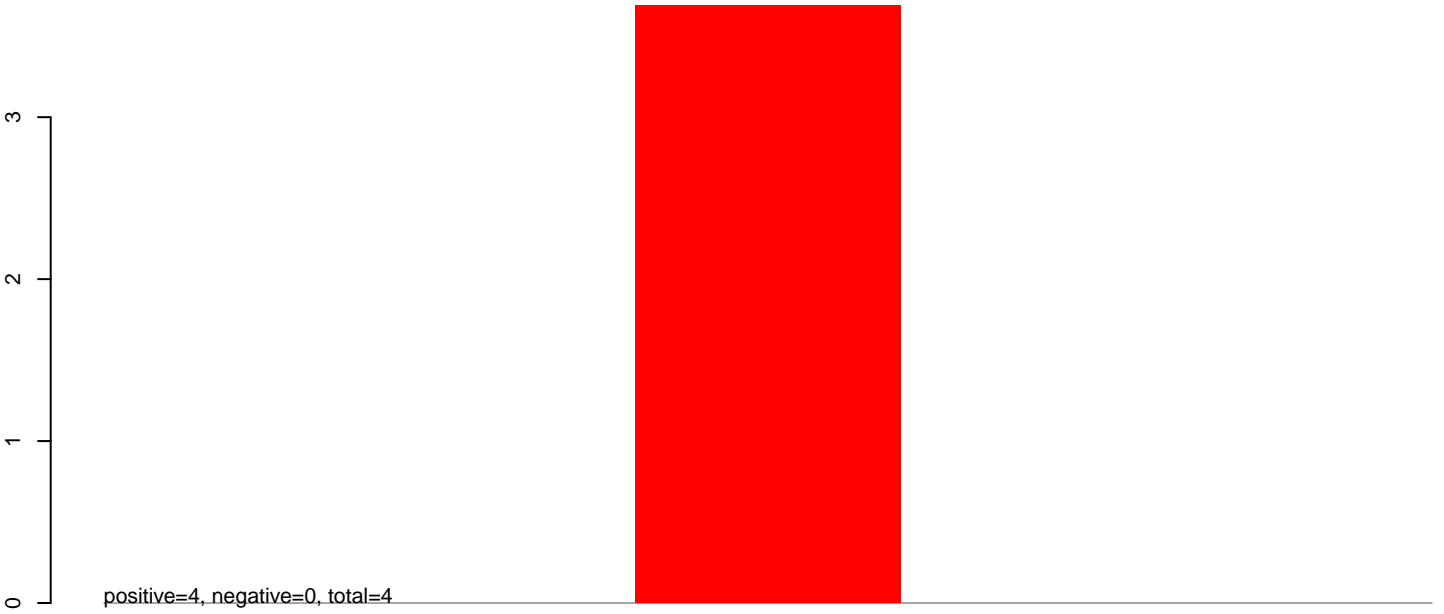


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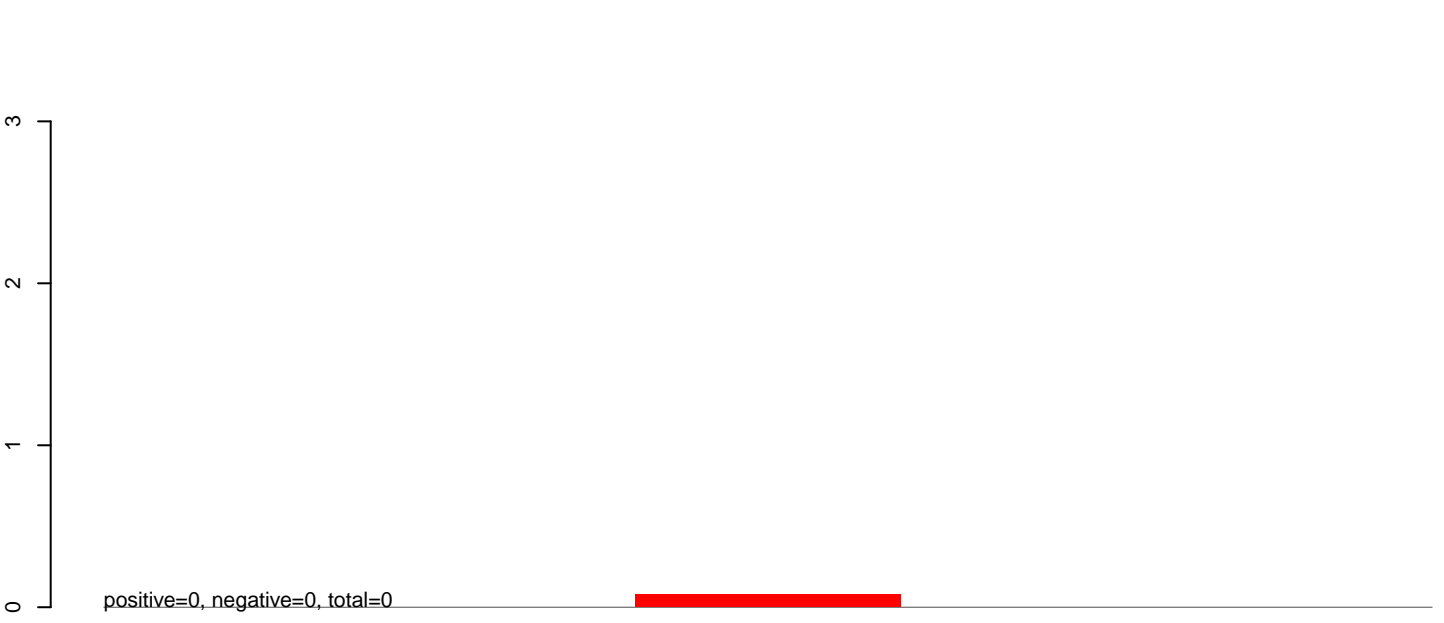


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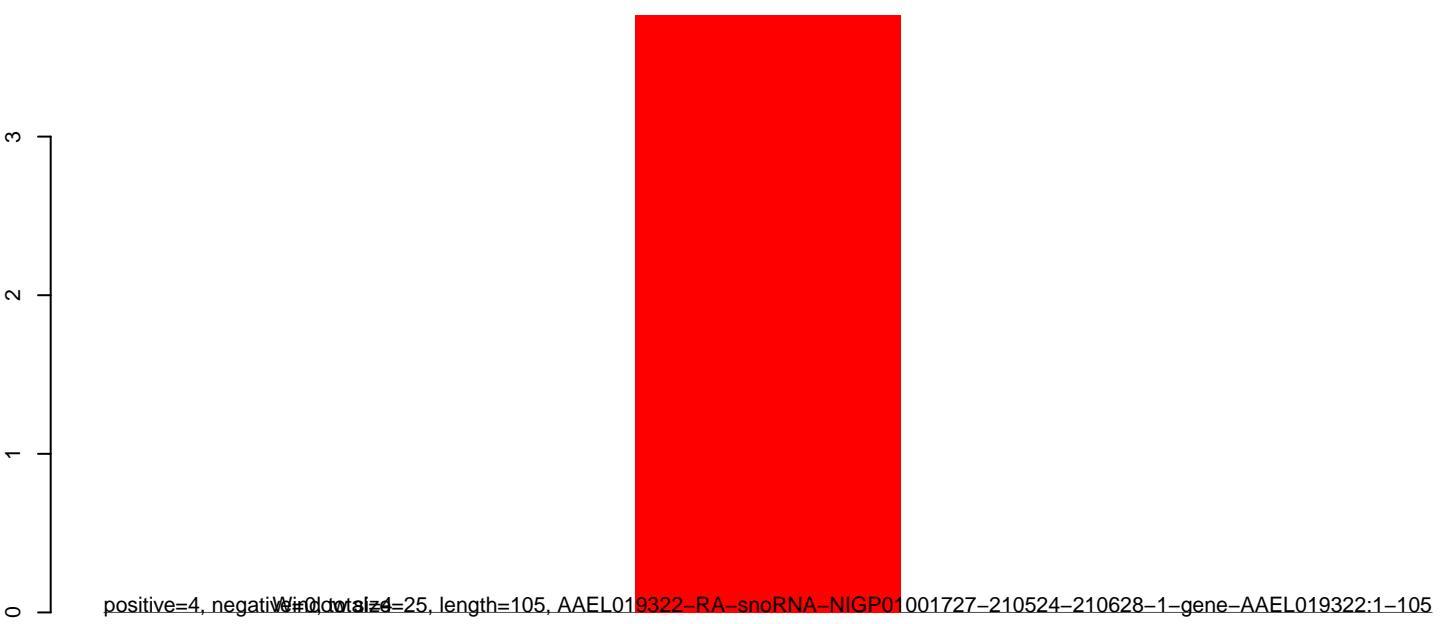
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AeAeg_Fcarc_GH_rep2.24_35.rep



AeAeg_Fcarc_GH_rep2.rep

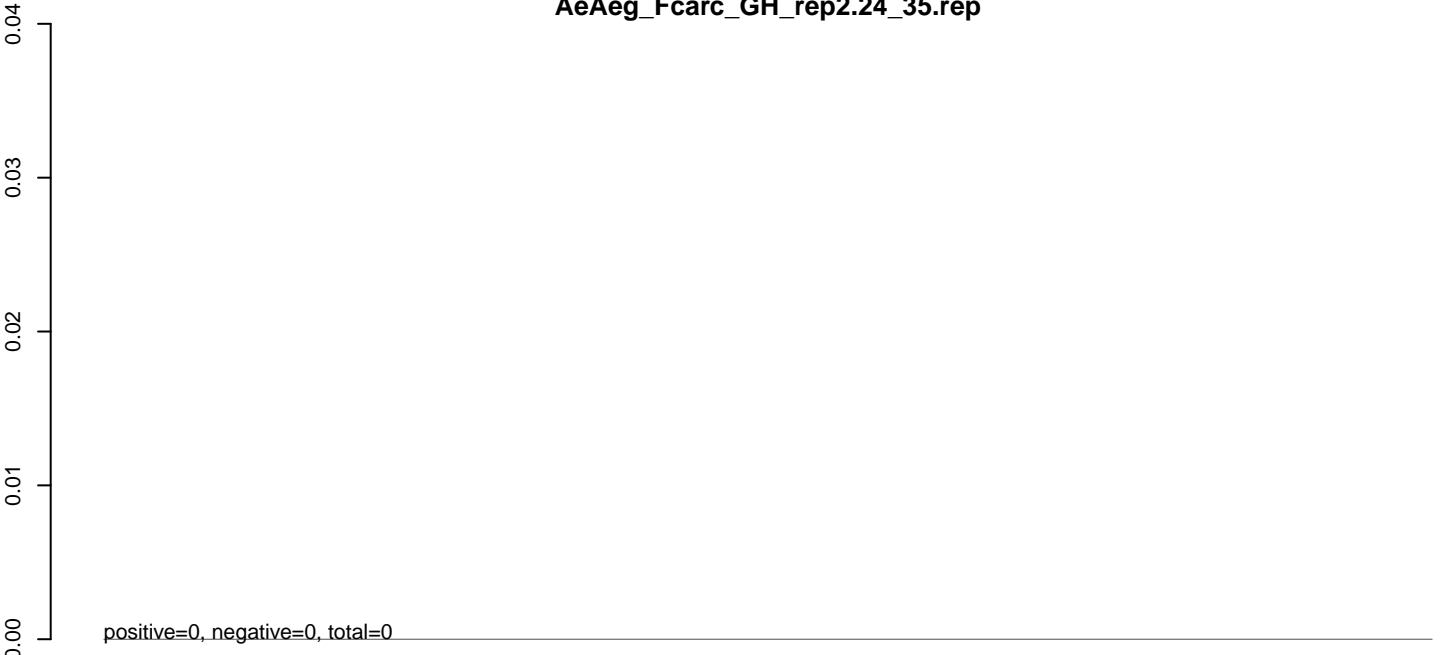


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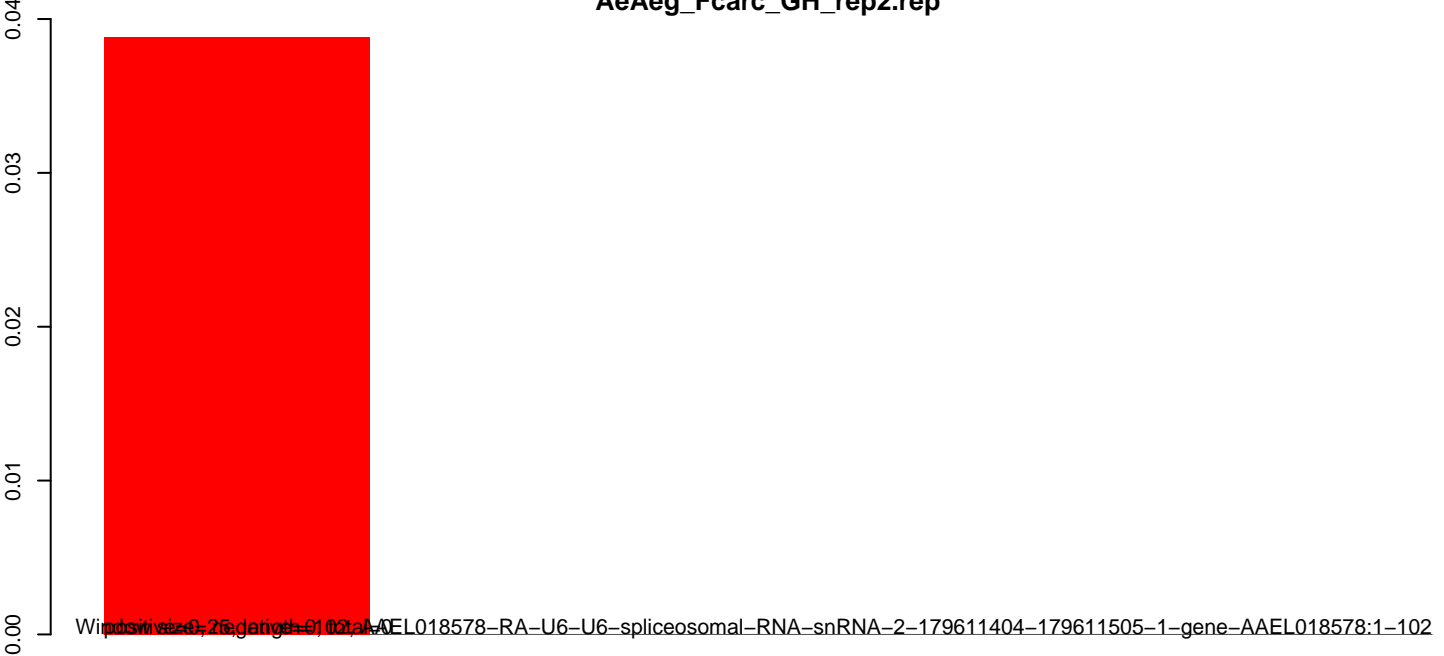
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AeAeg_Fcarc_GH_rep2.24_35.rep



AeAeg_Fcarc_GH_rep2.rep



Wings: size=6, range=0-102, AAEL018578-RA-U6-U6-spliceosomal-RNA-snRNA-2-179611404-179611505-1-gene-AAEL018578:1-102

40 60 80 100 120 140

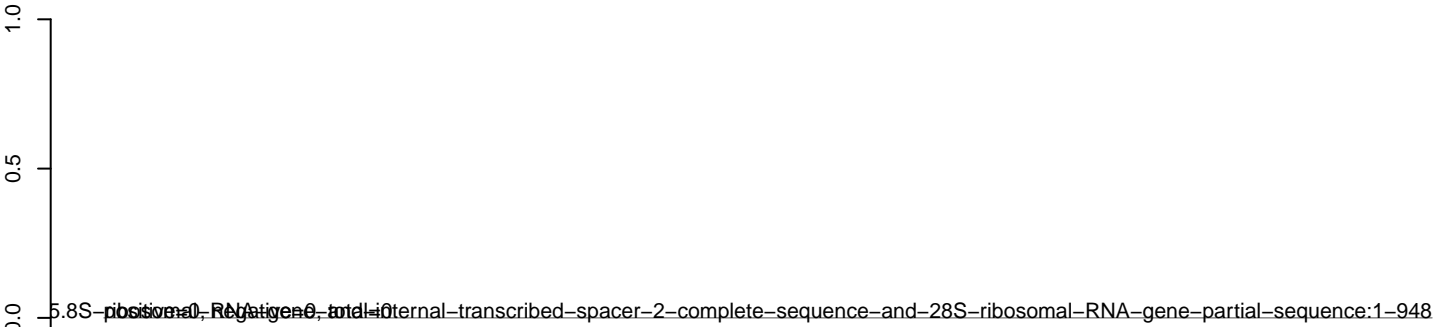
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AeAeg_Fcarc_GH_rep2.24_35.rep



AeAeg_Fcarc_GH_rep2.rep



0 200 400 600 800 1000