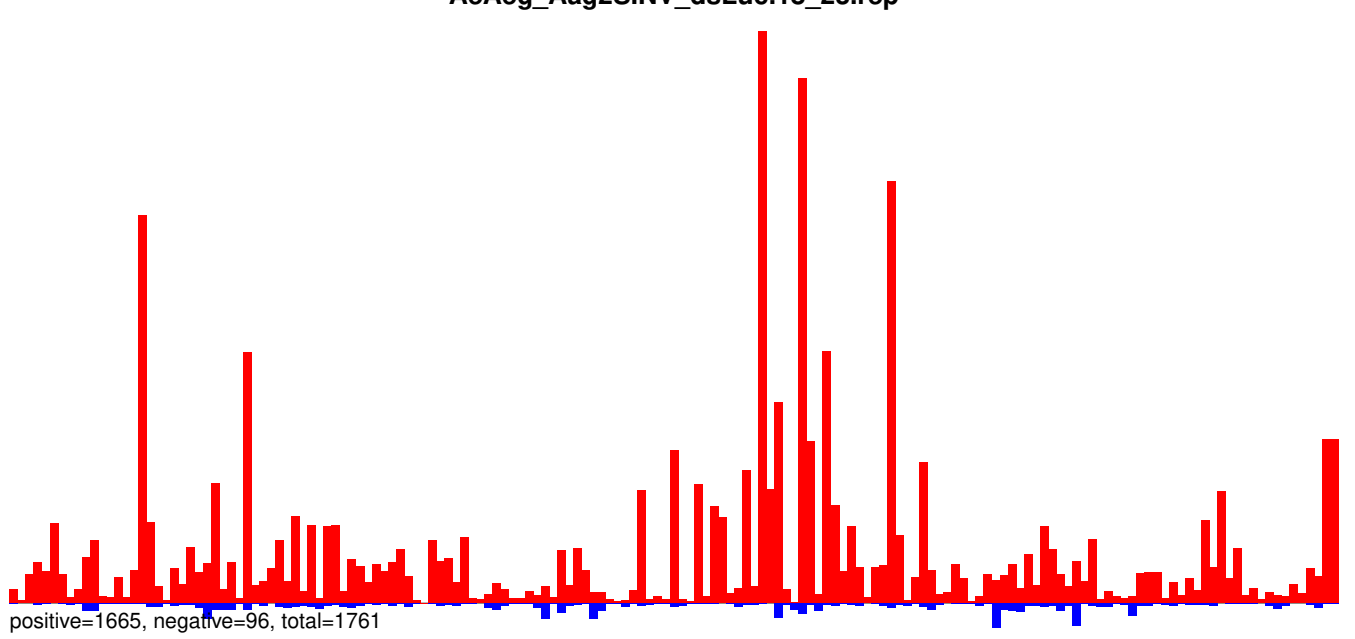
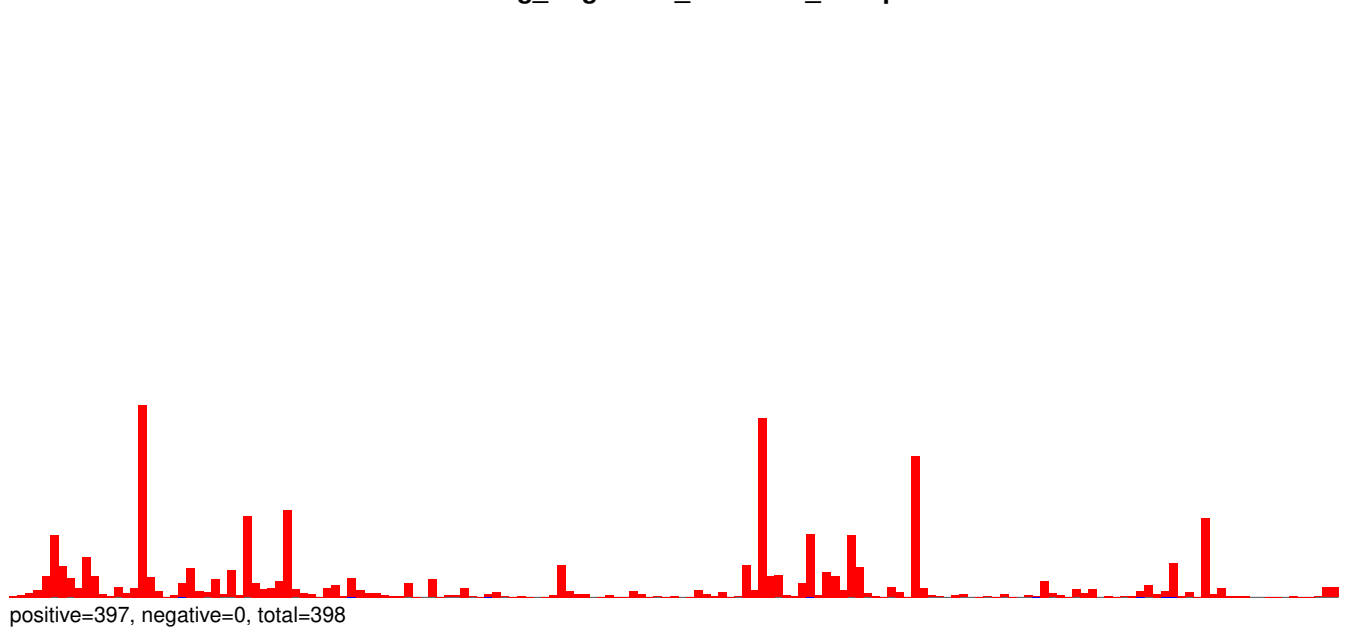


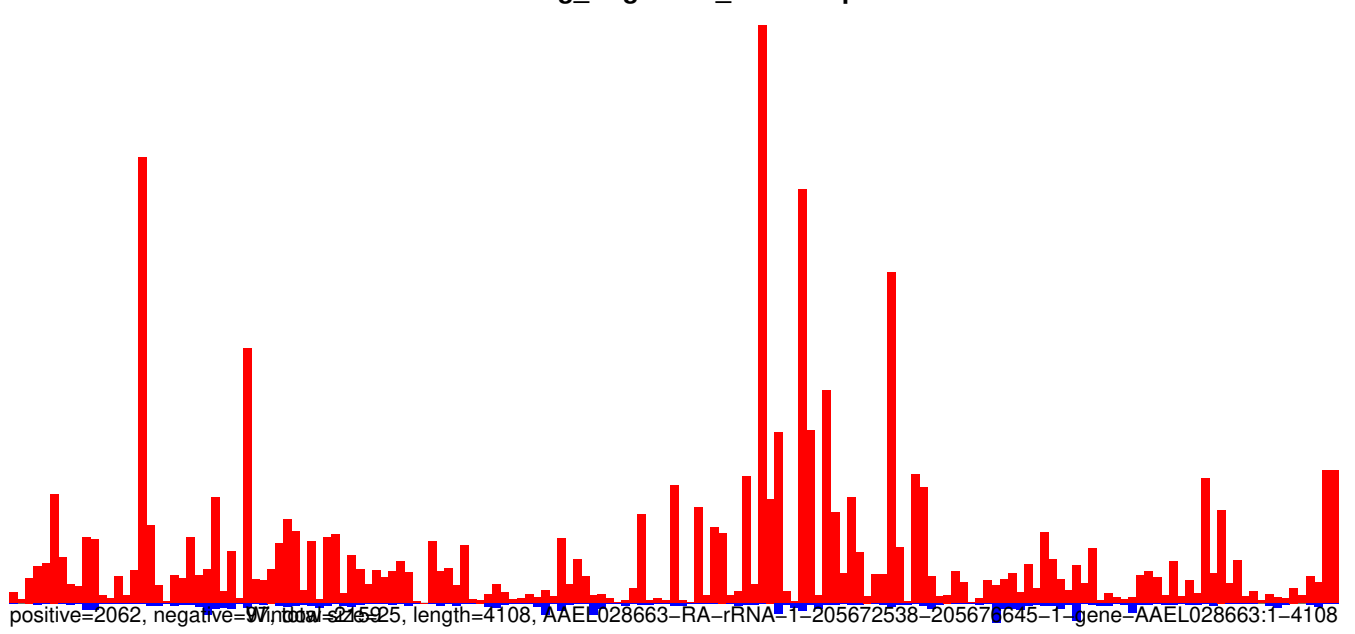
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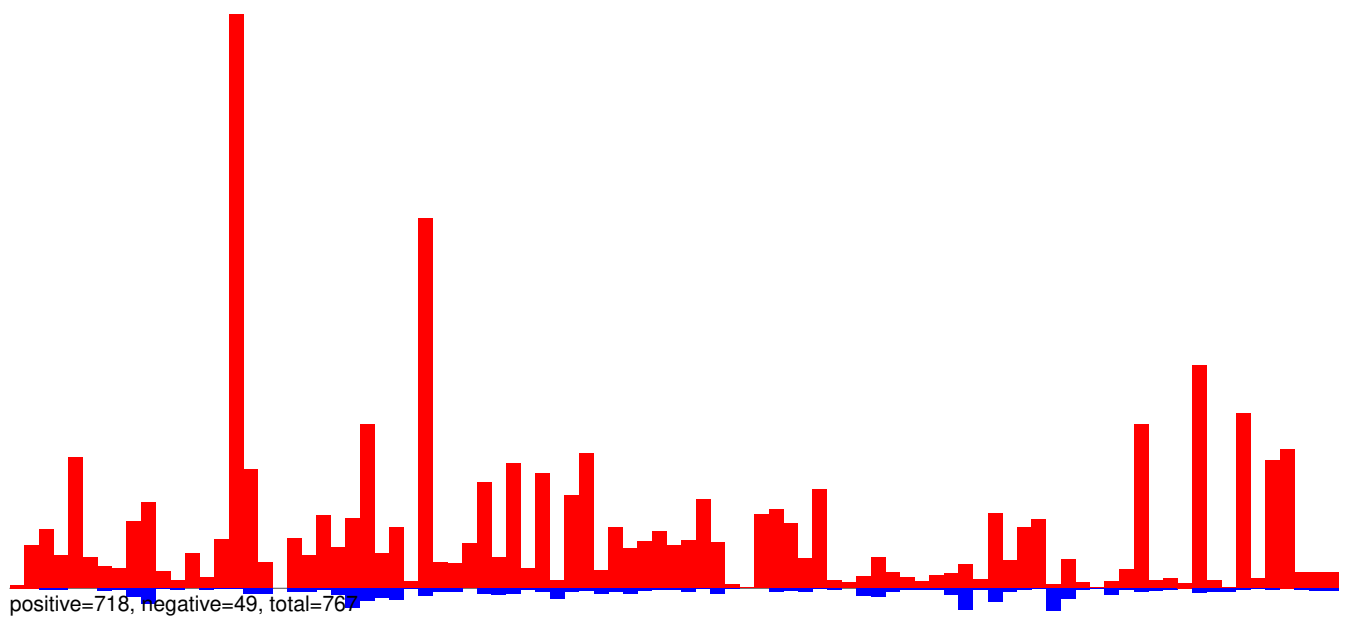
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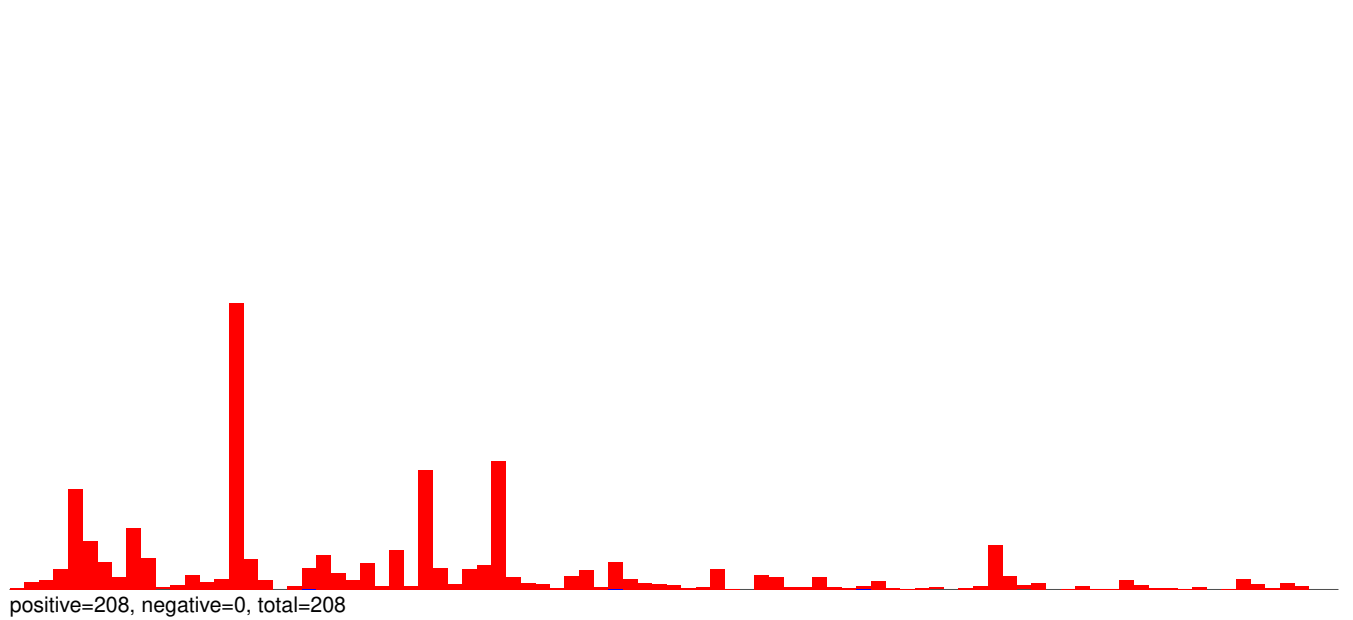
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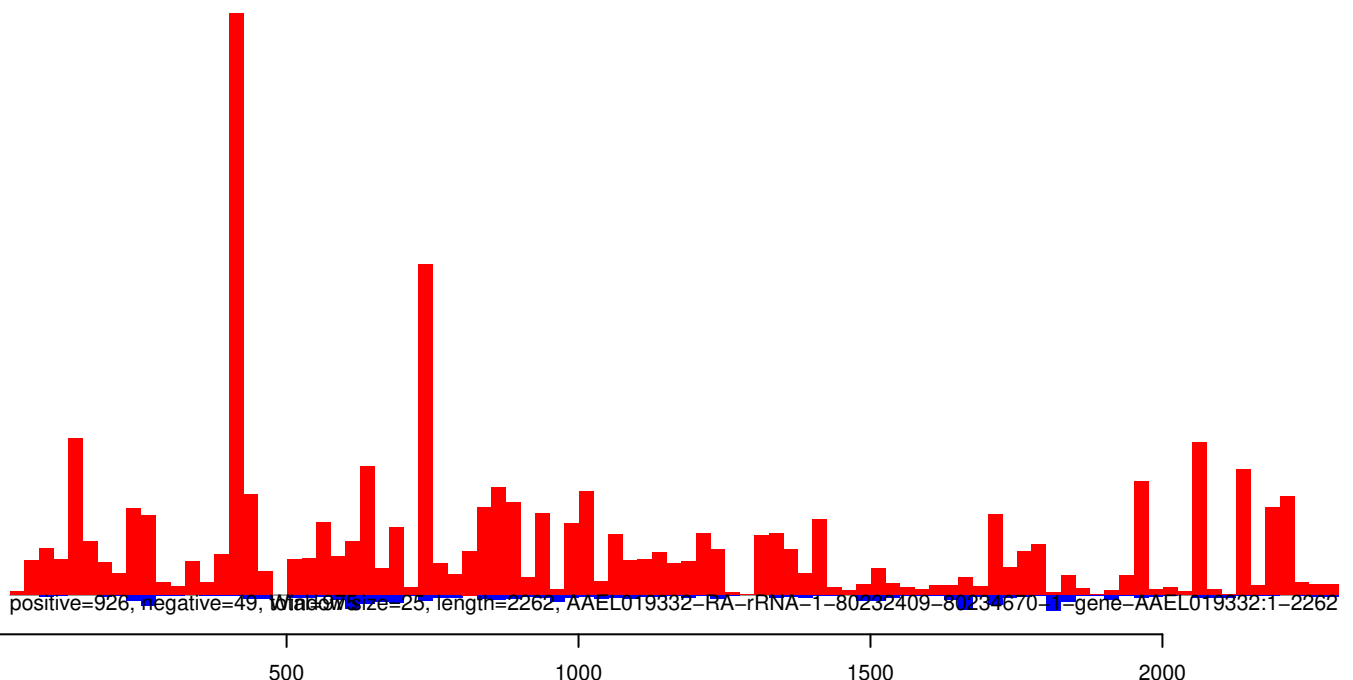
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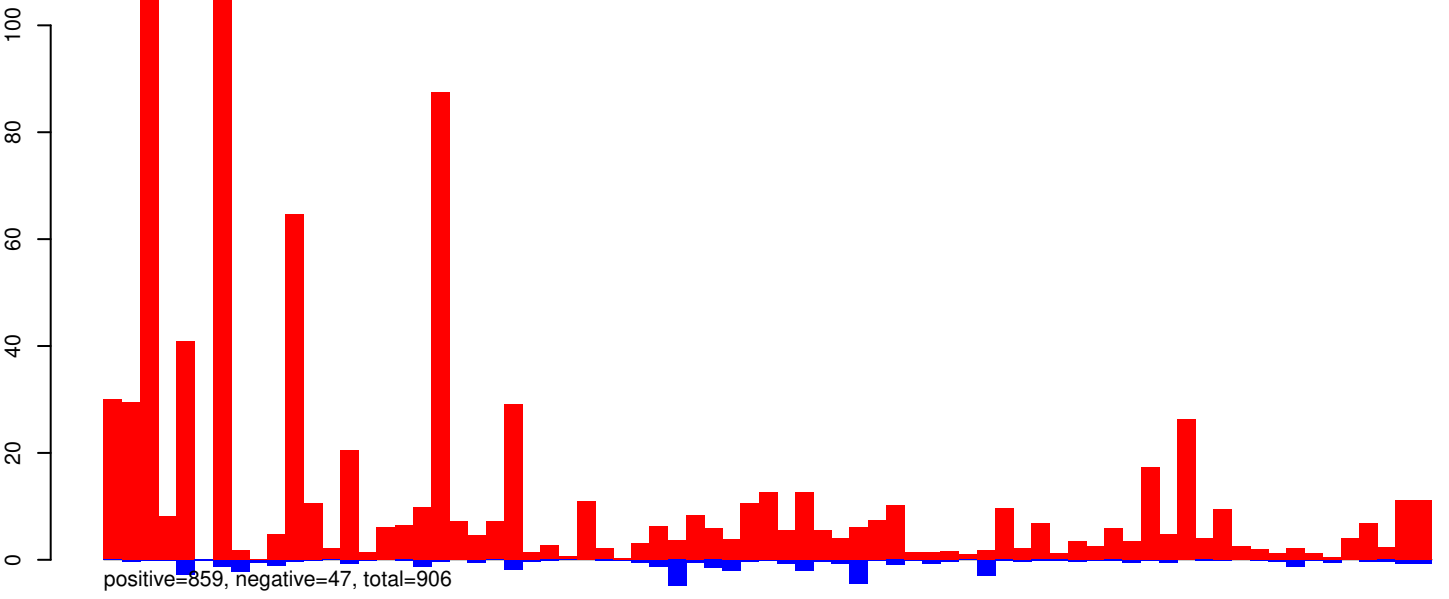
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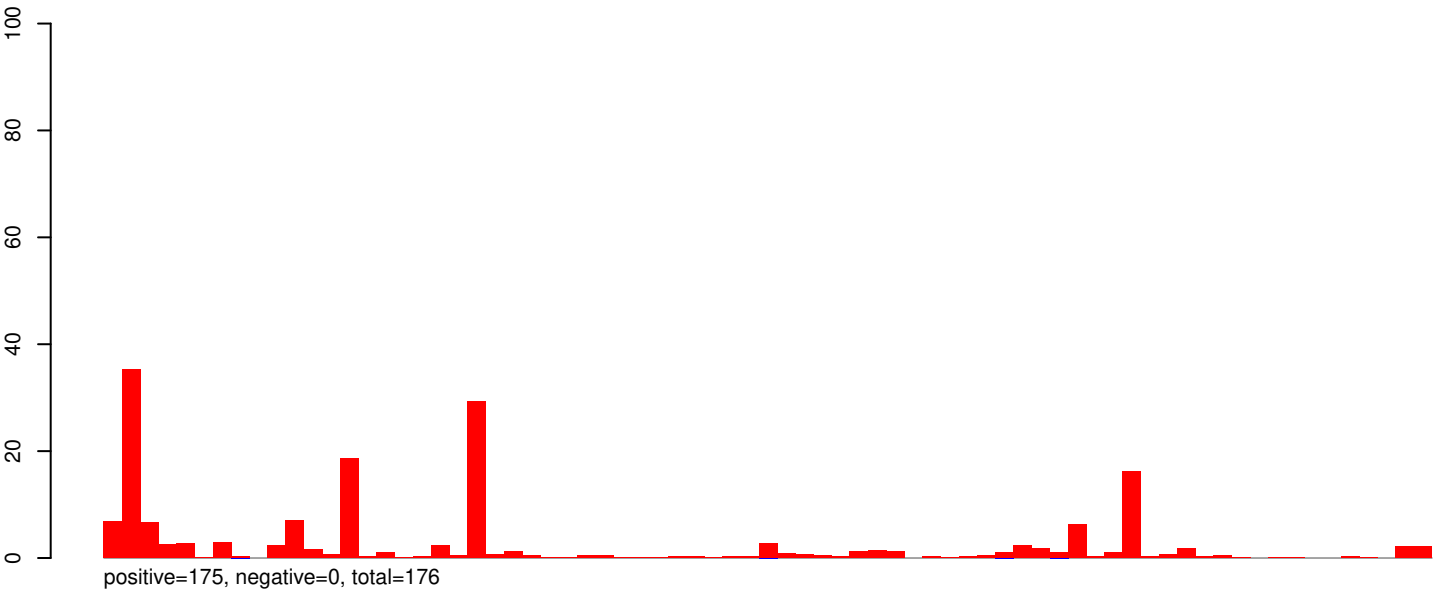
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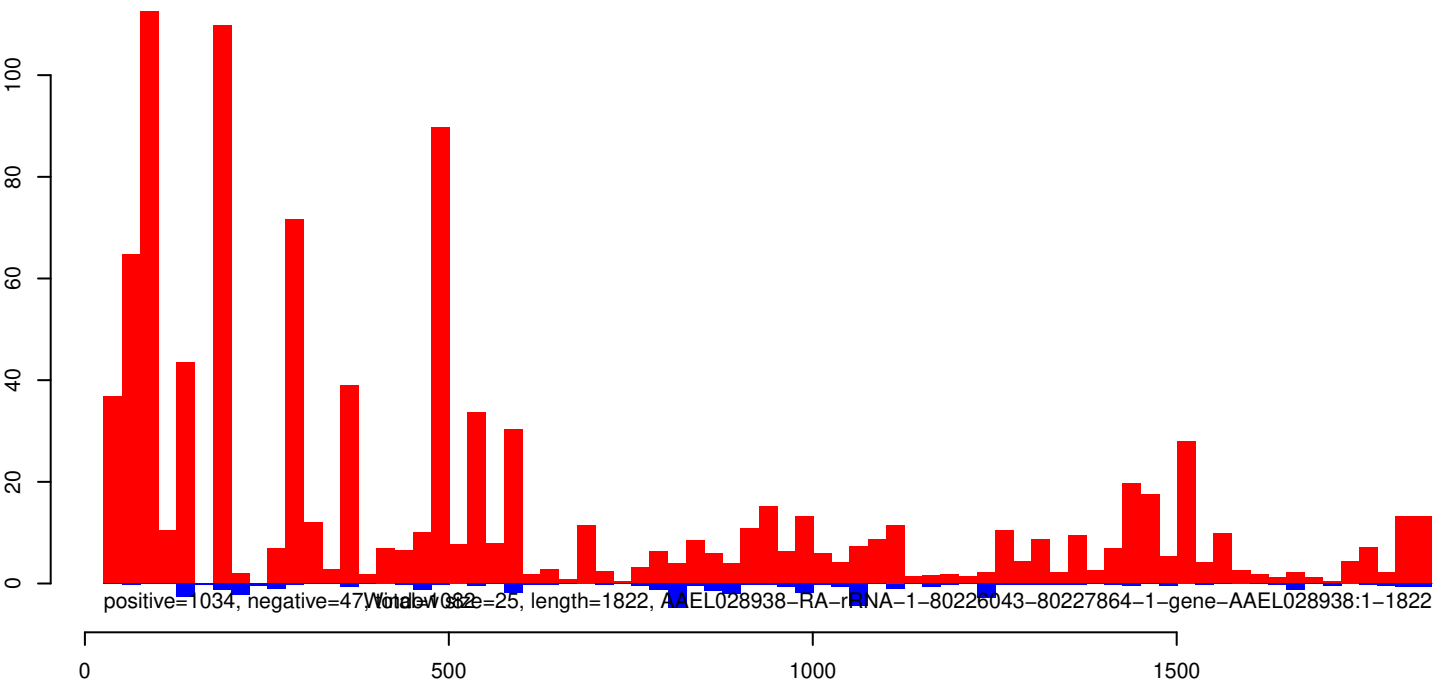
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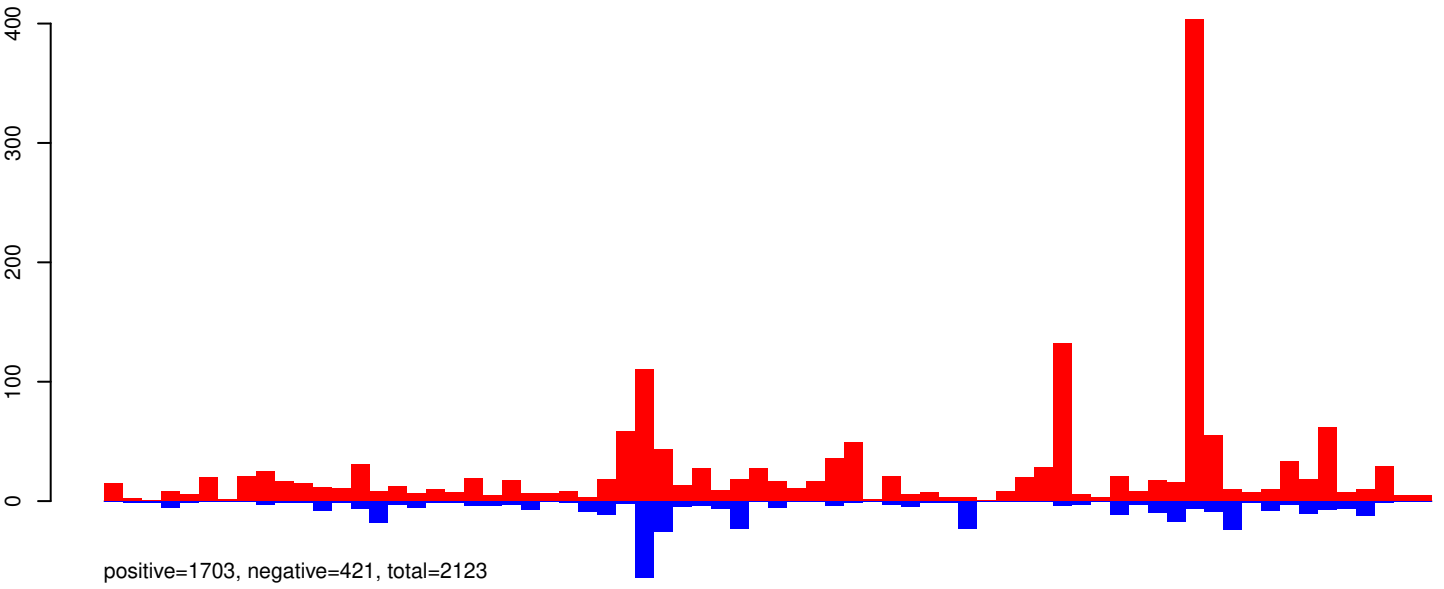
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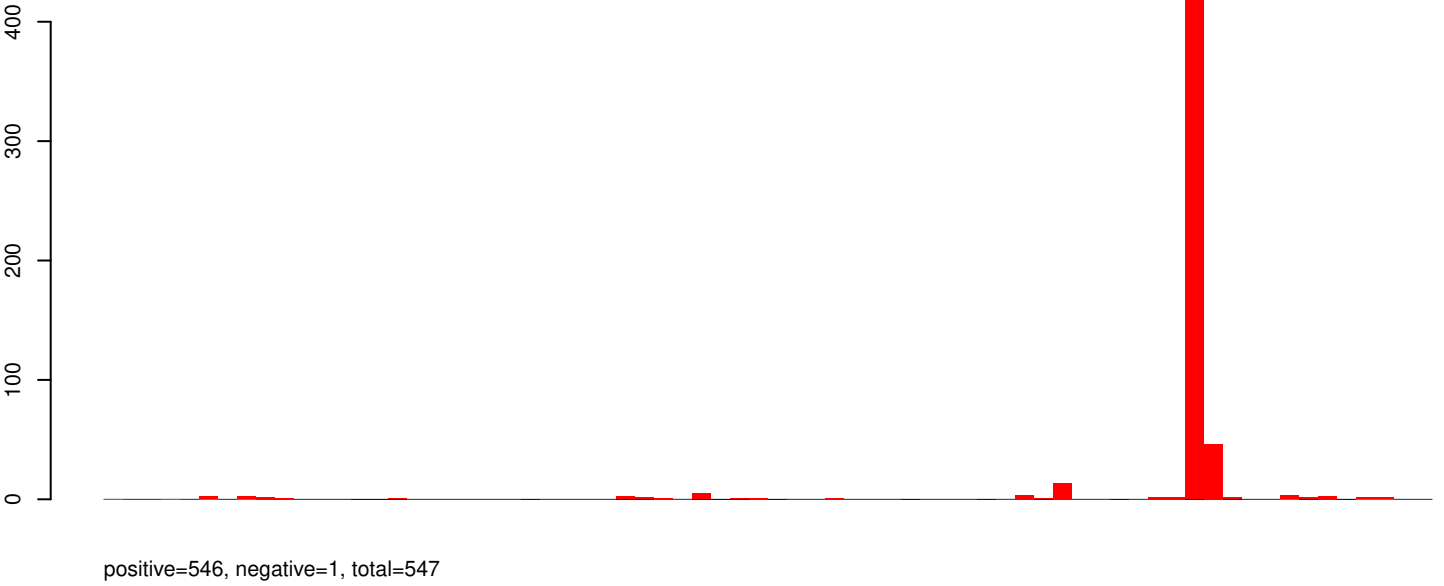
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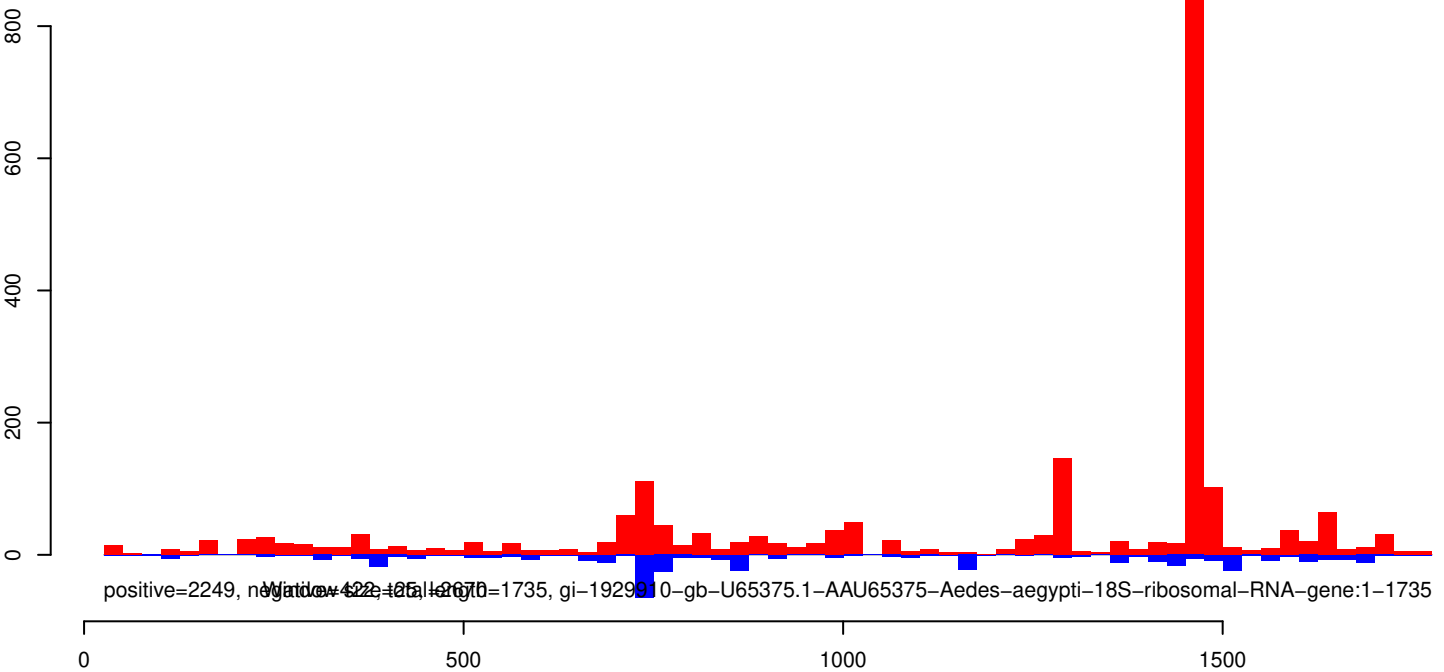
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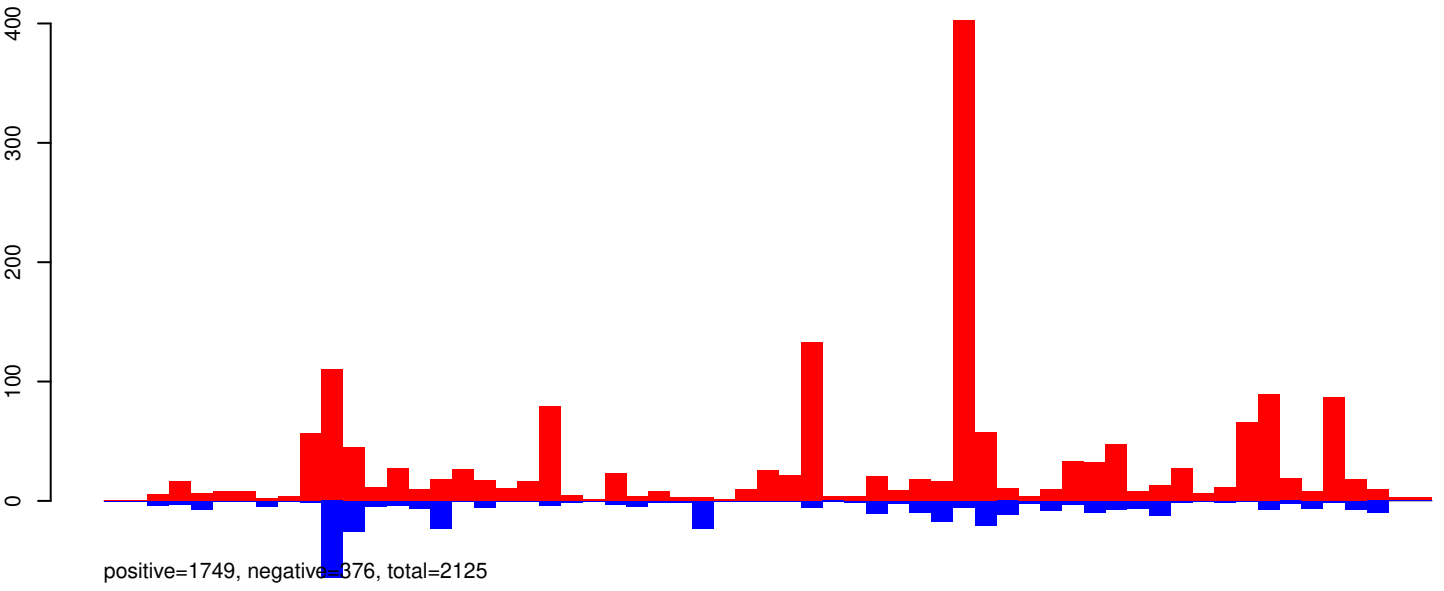
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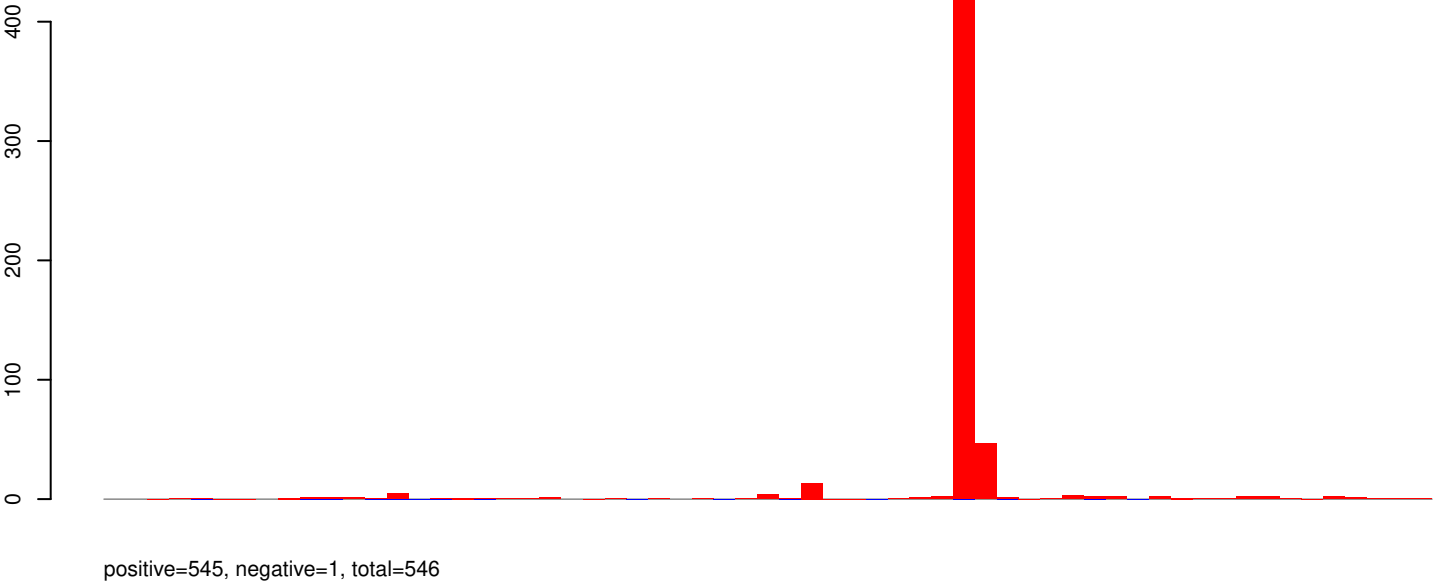
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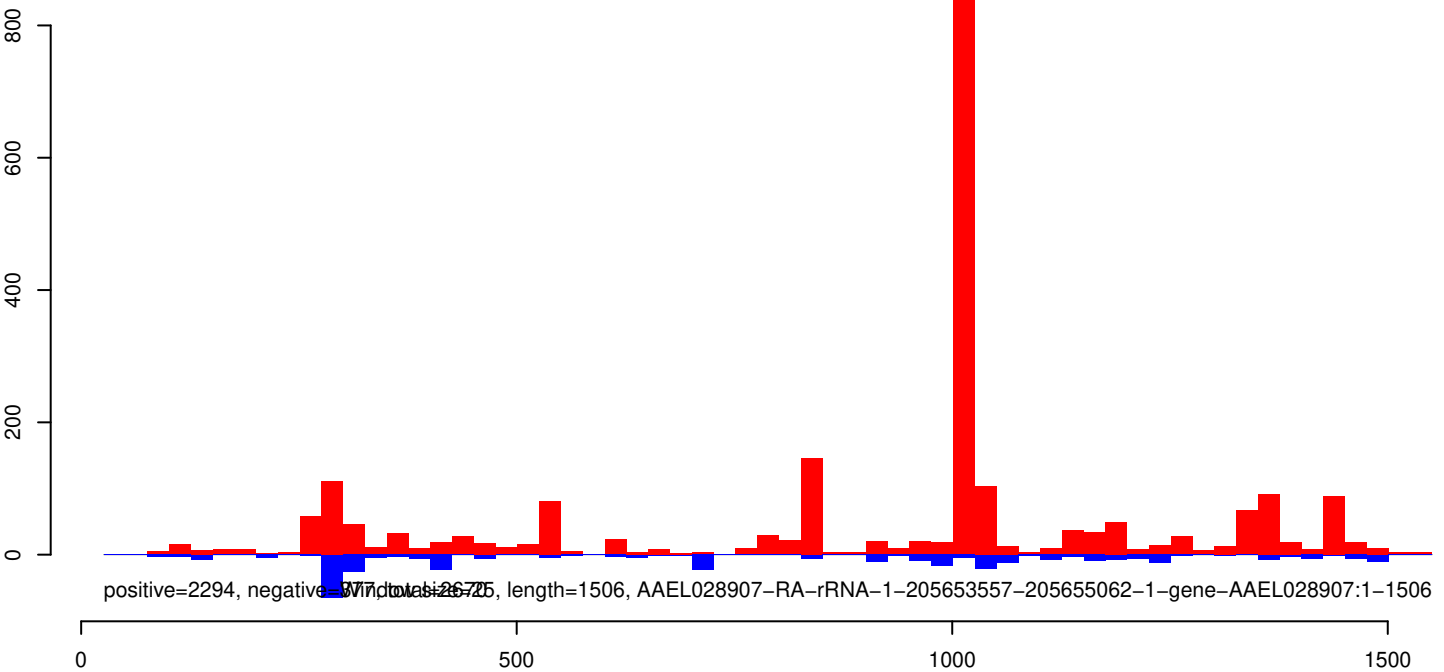
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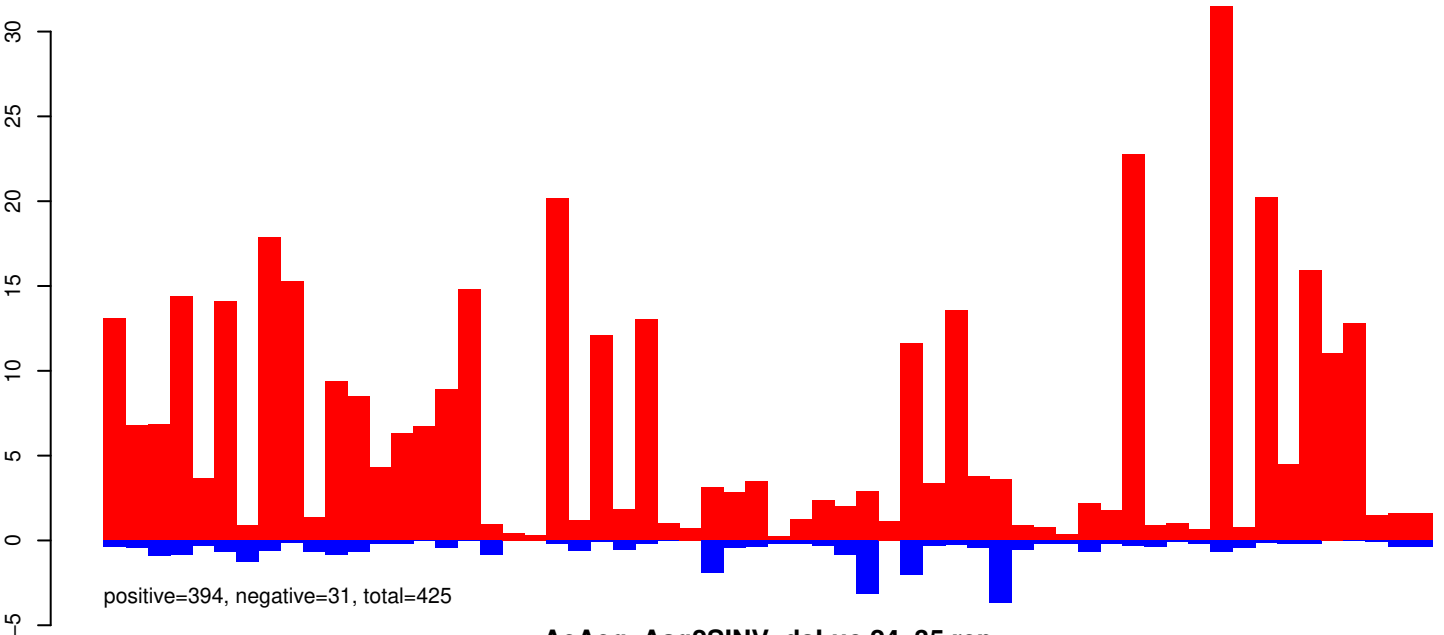
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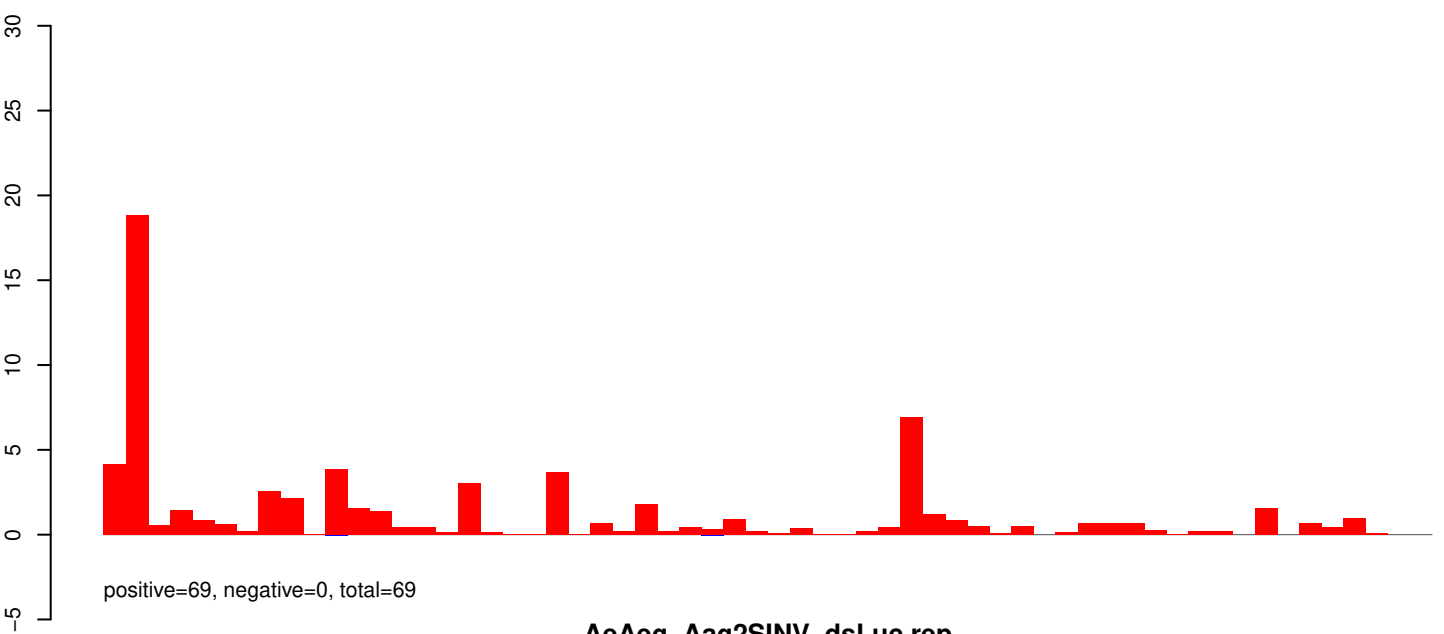
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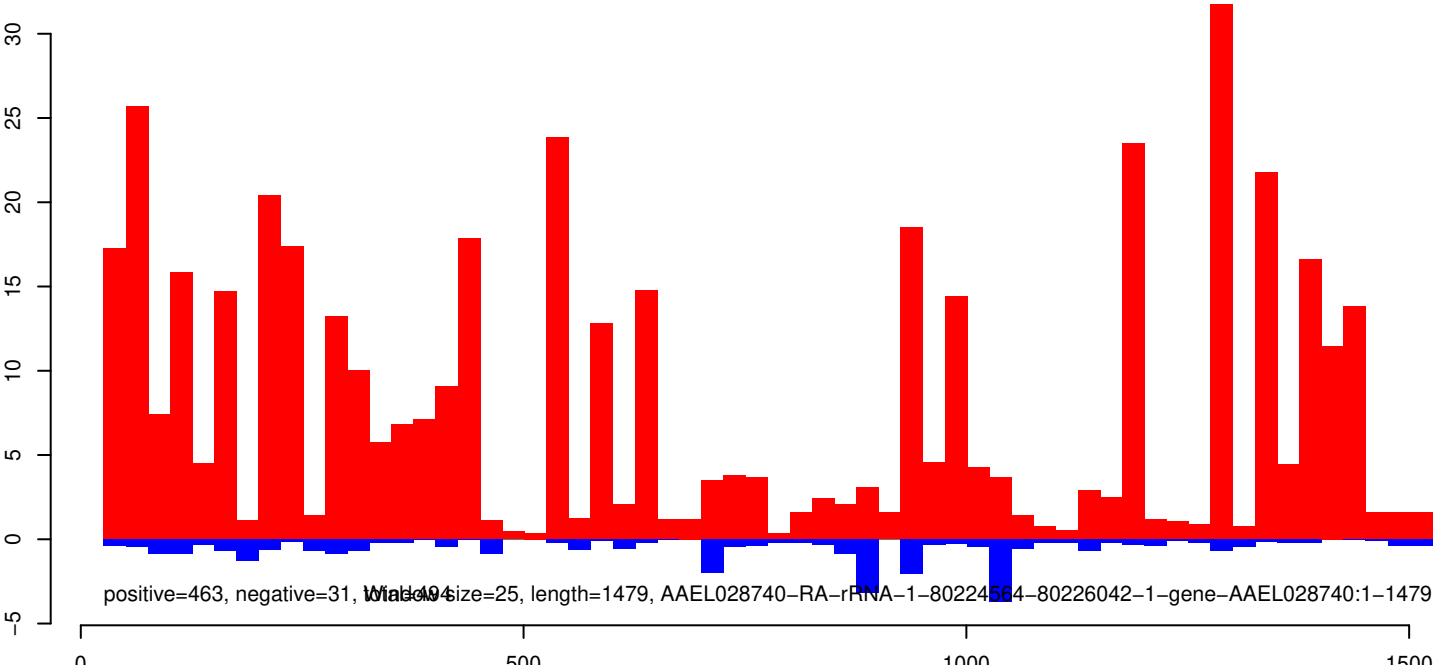
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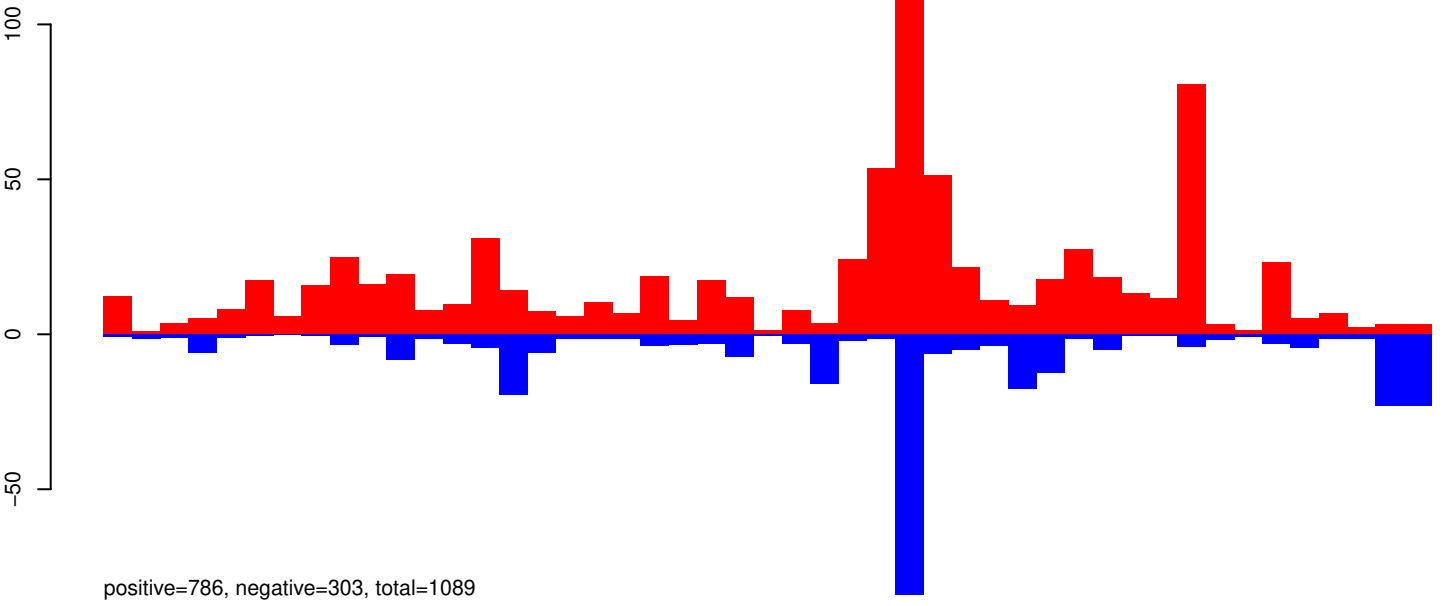
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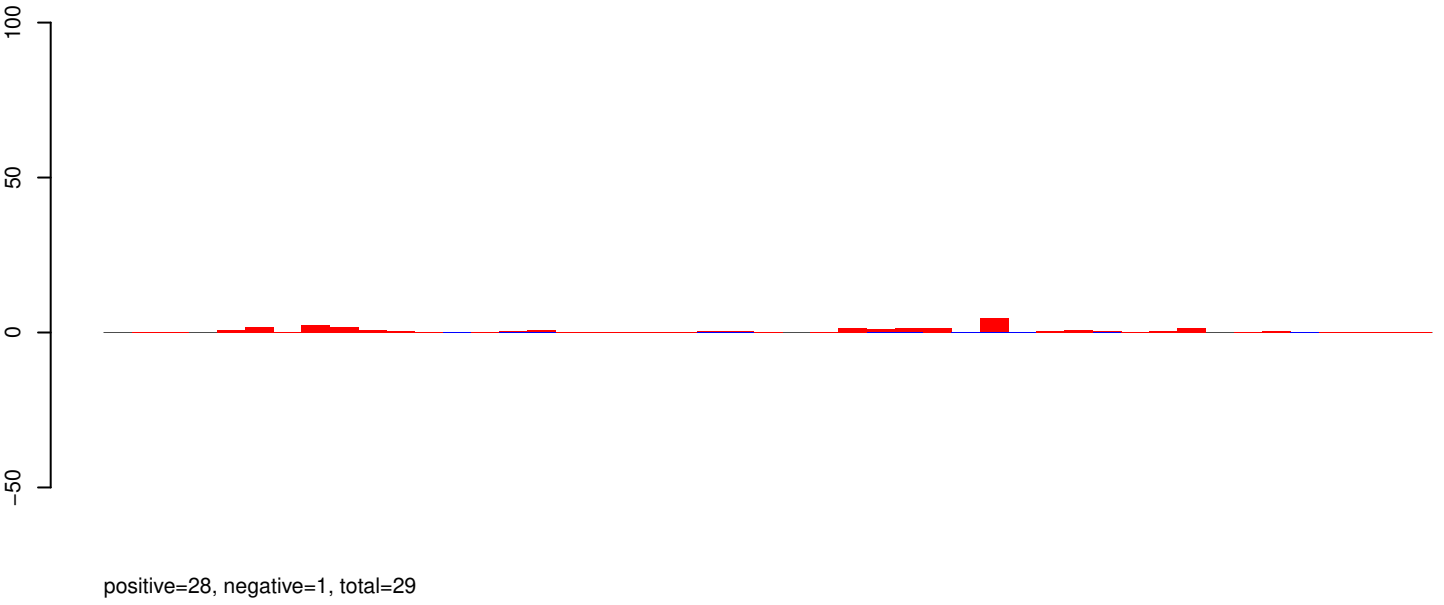
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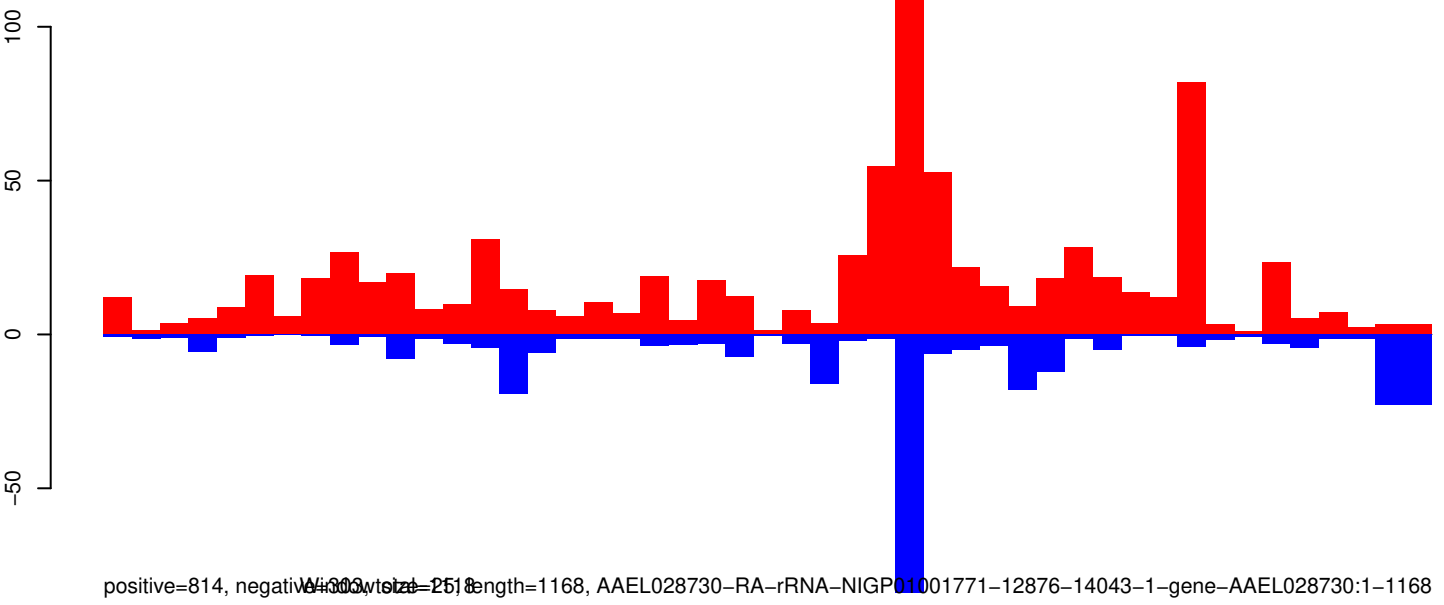
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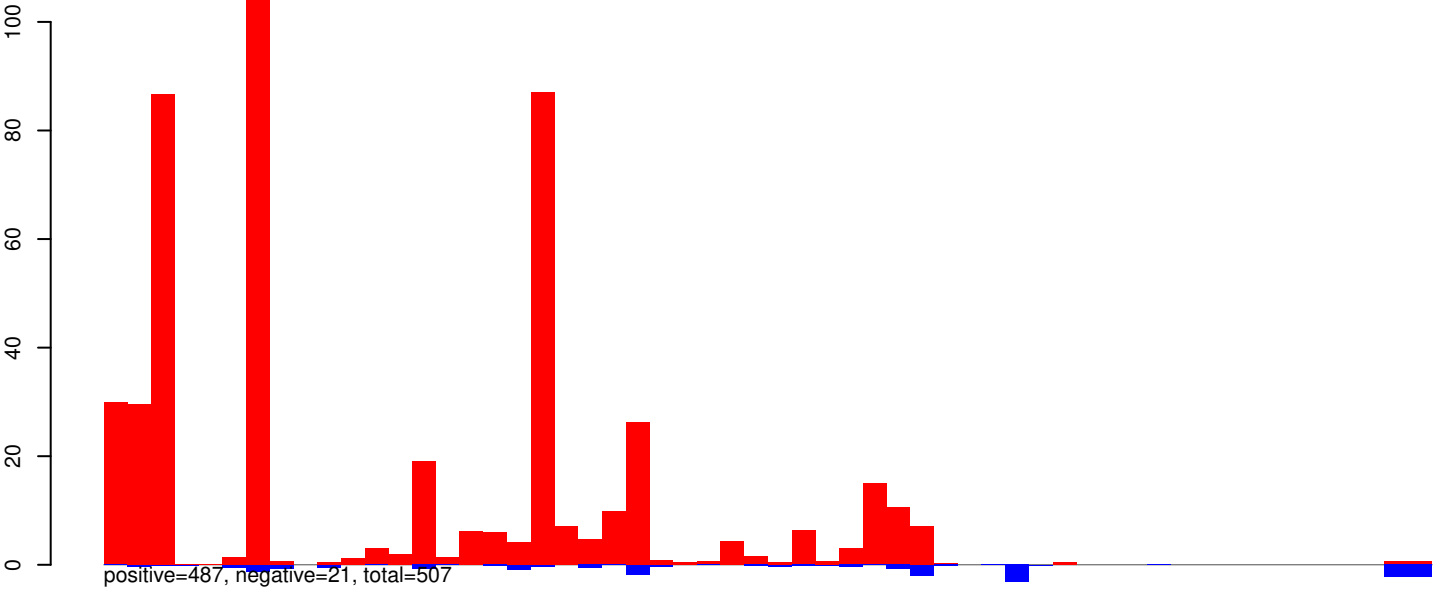
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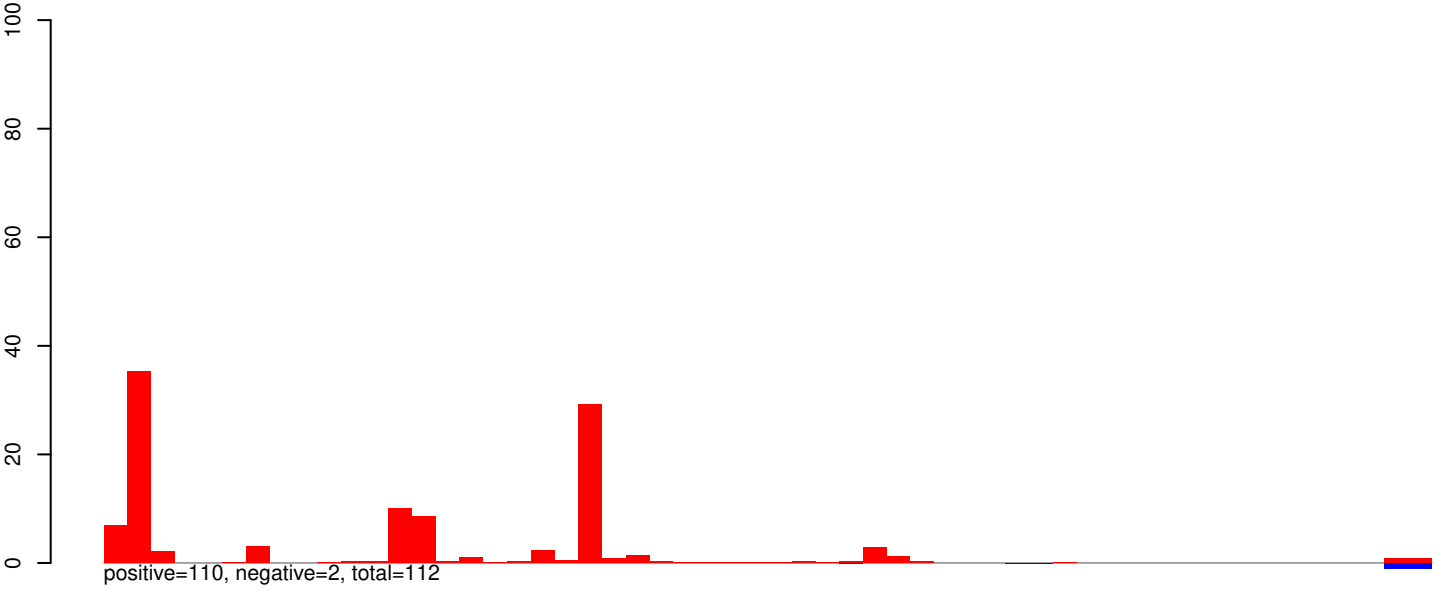
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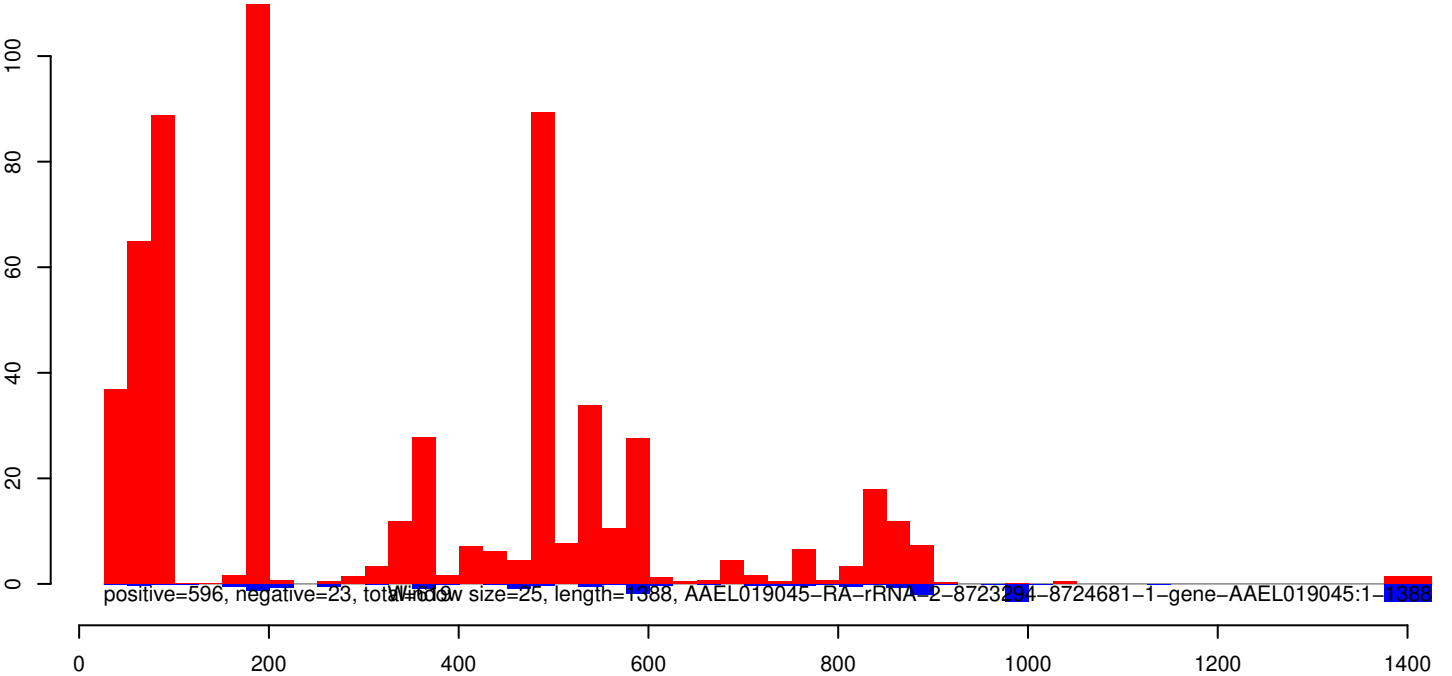
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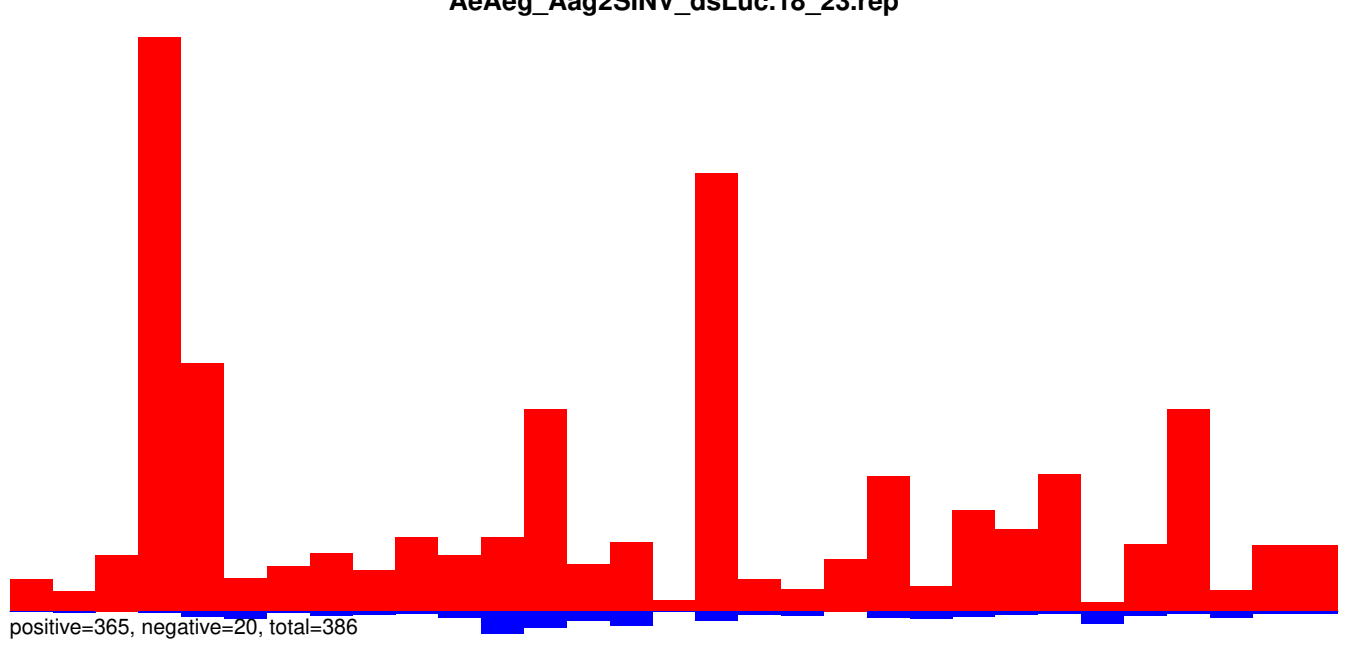
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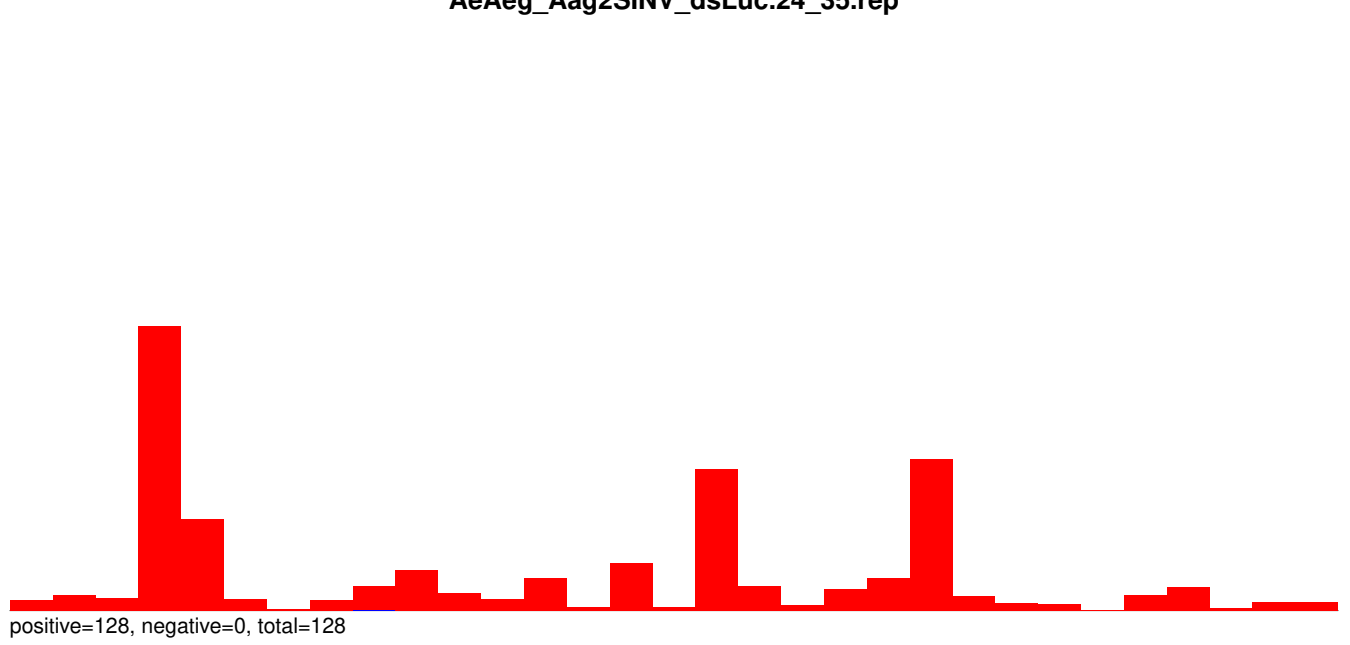
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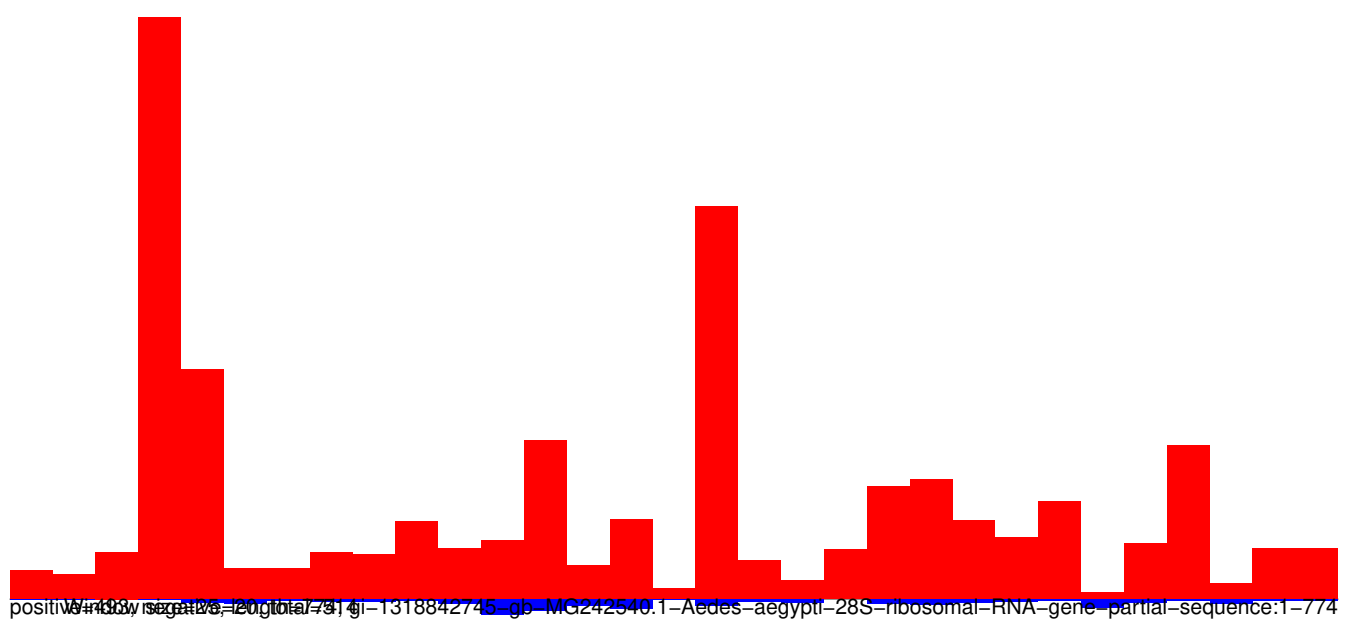
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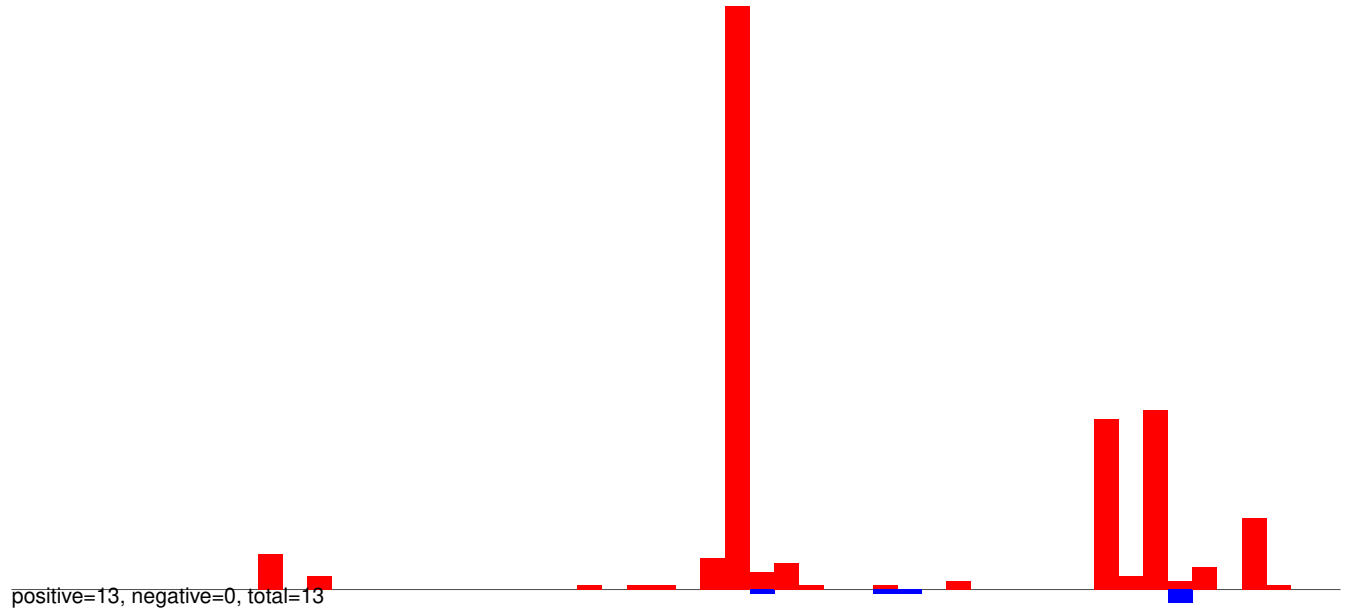
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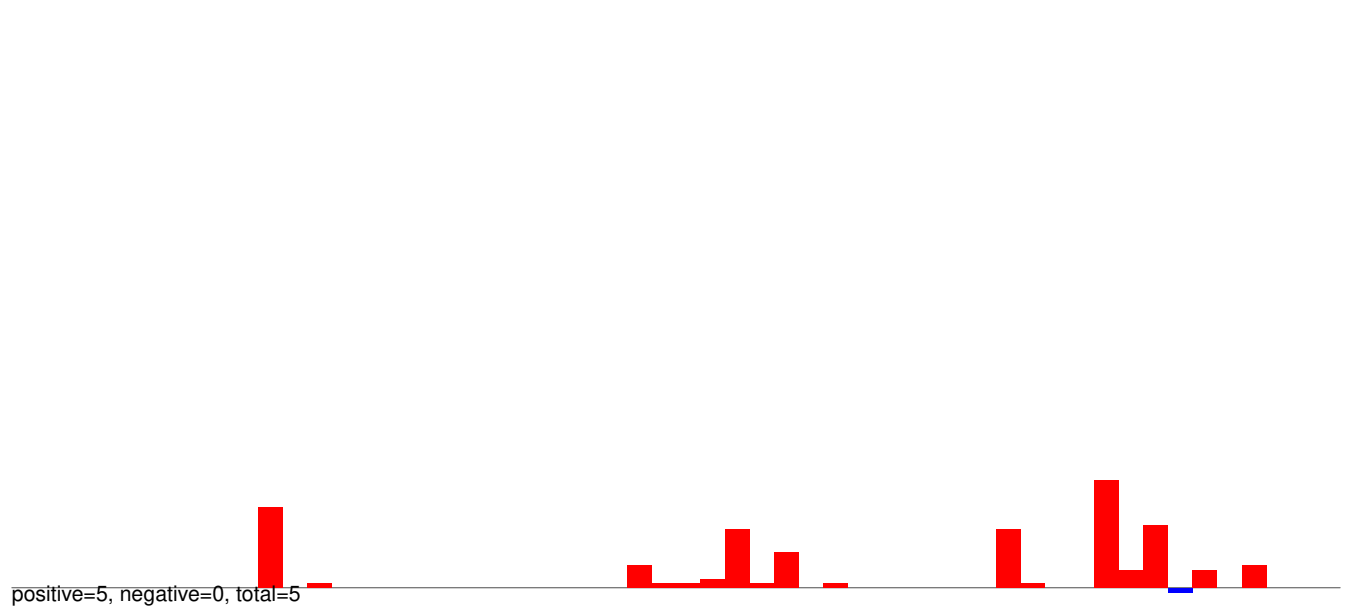
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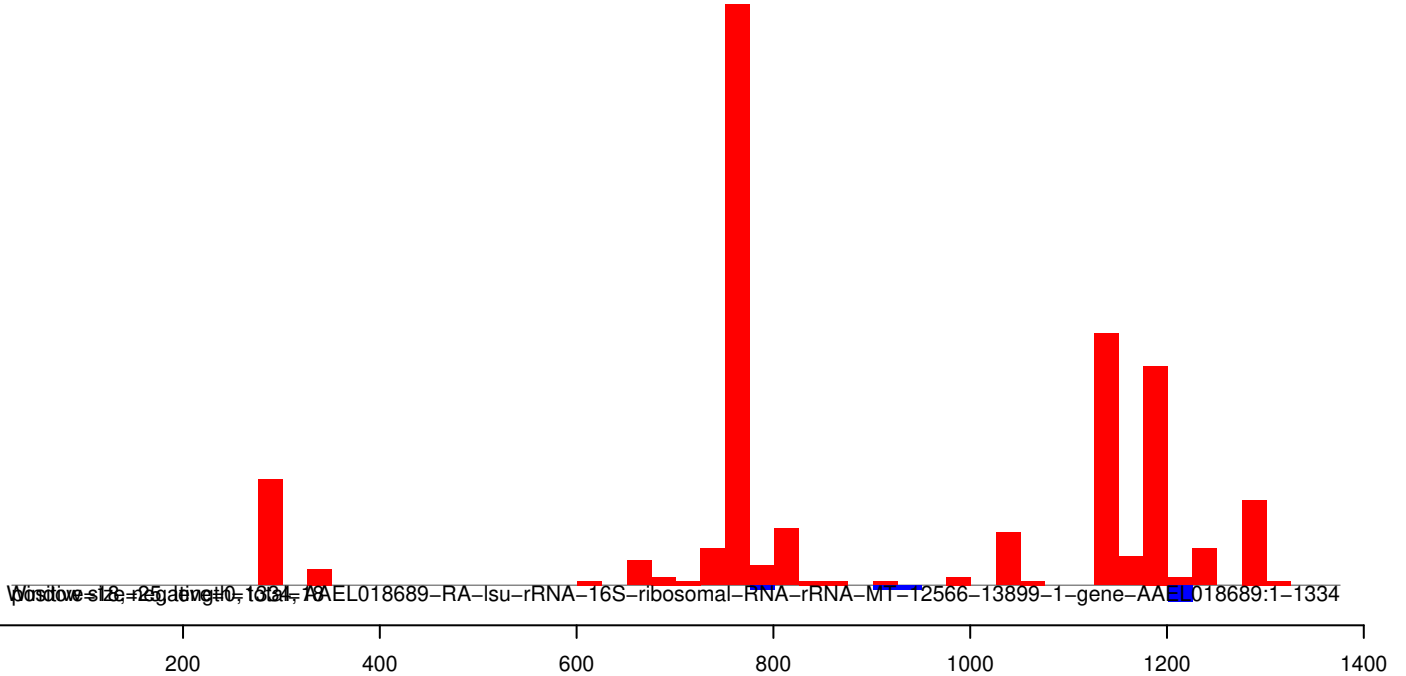
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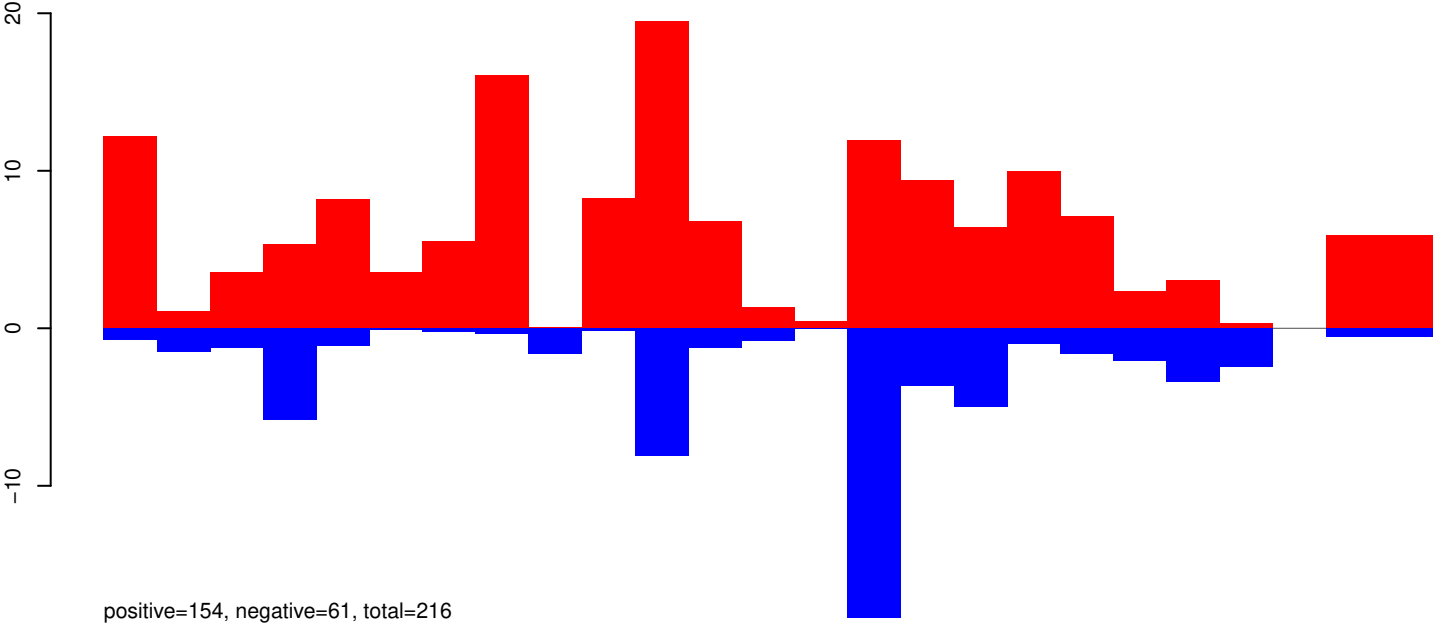
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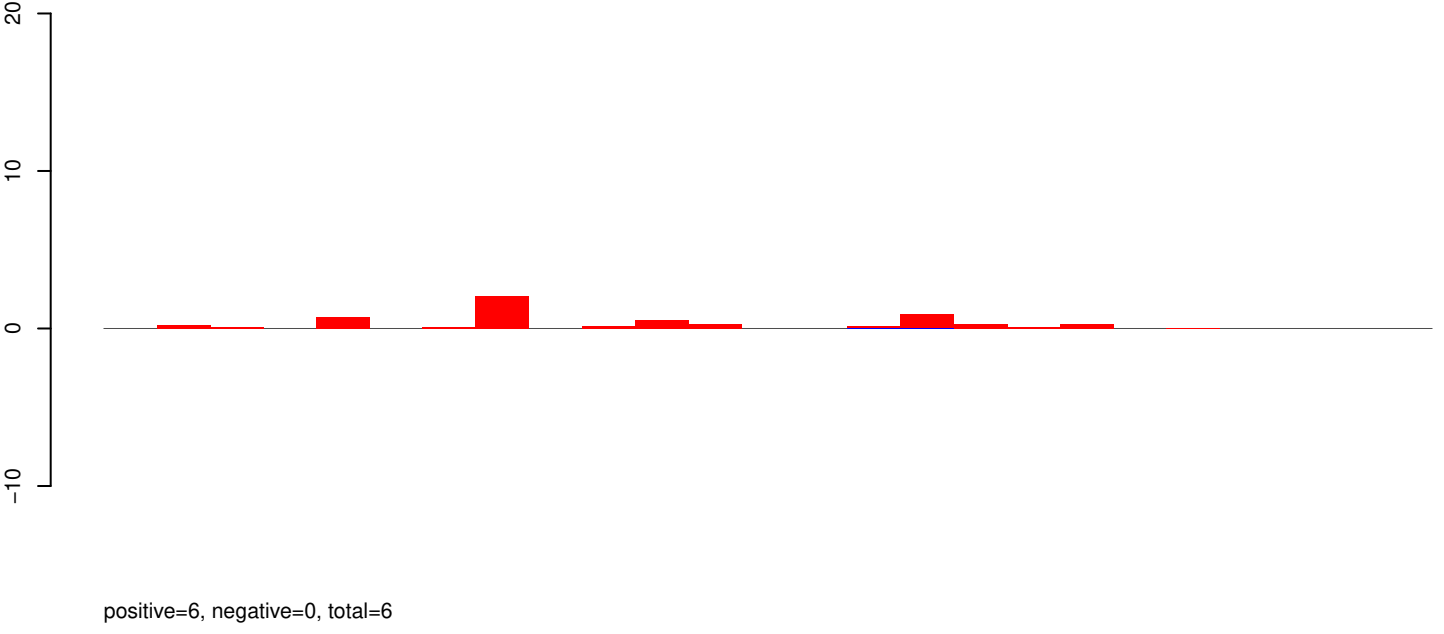
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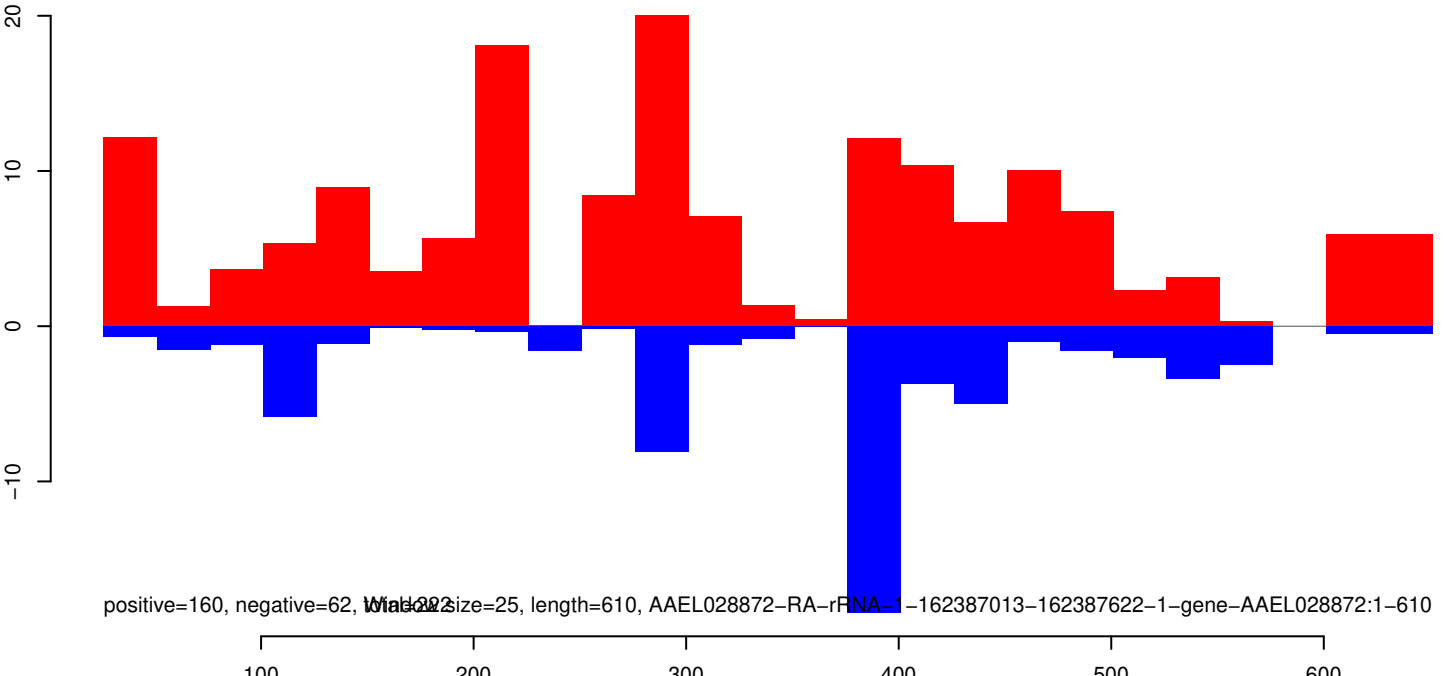
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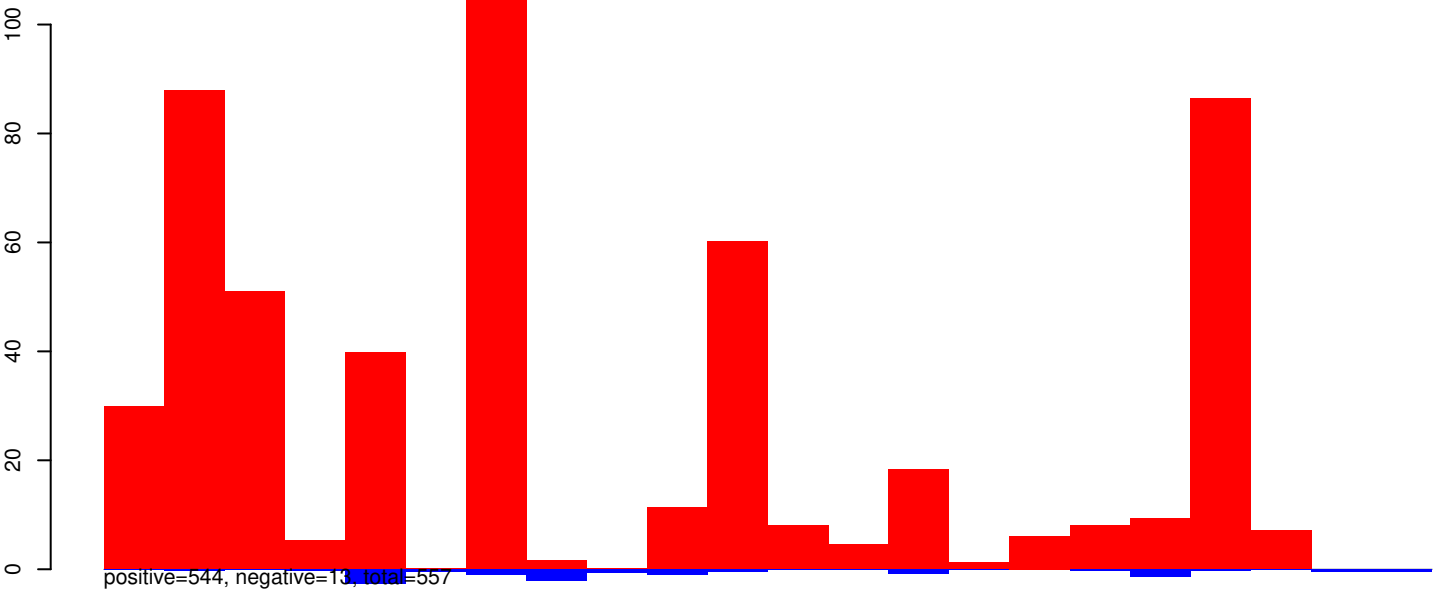
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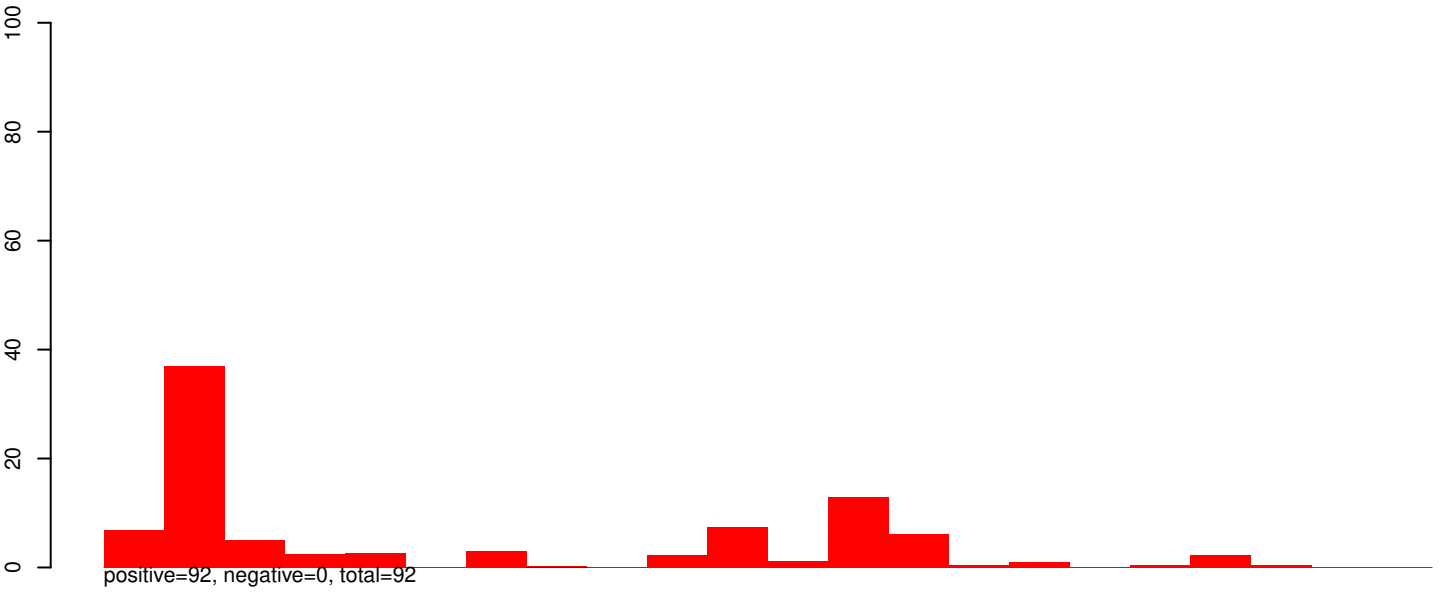
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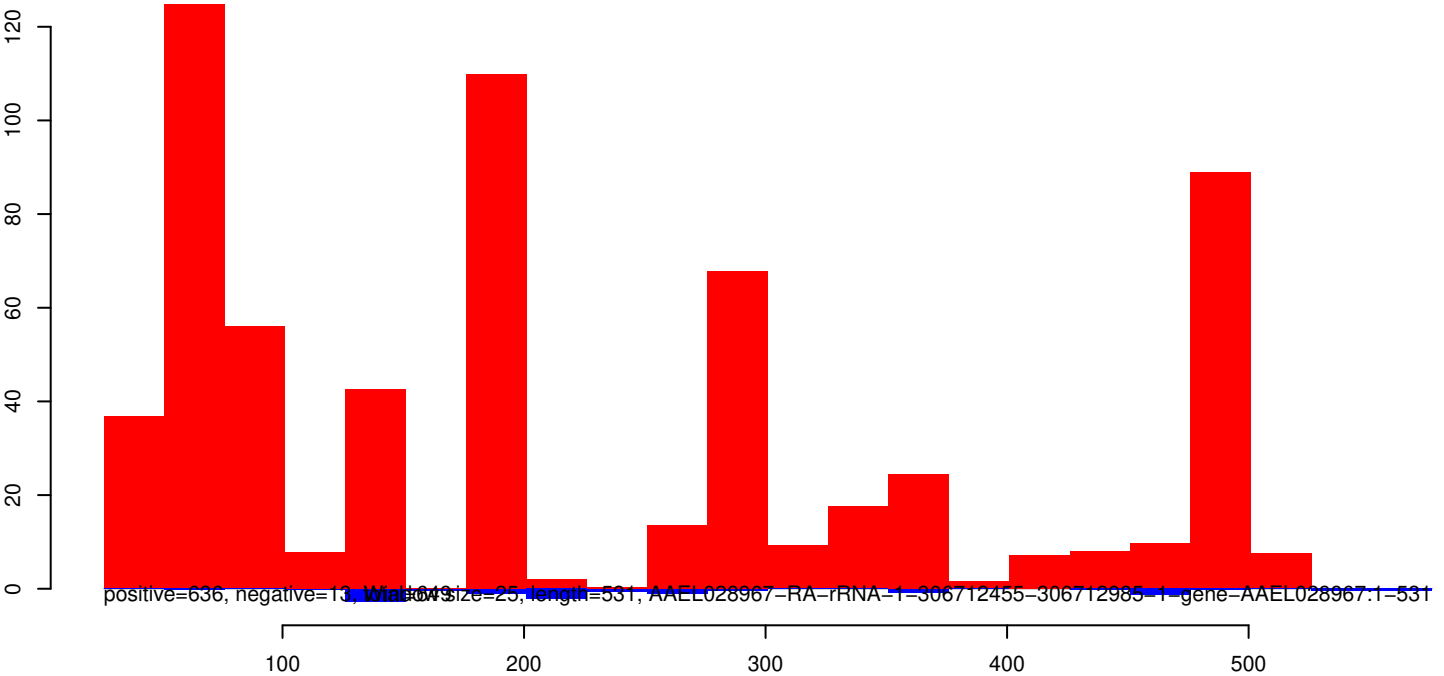
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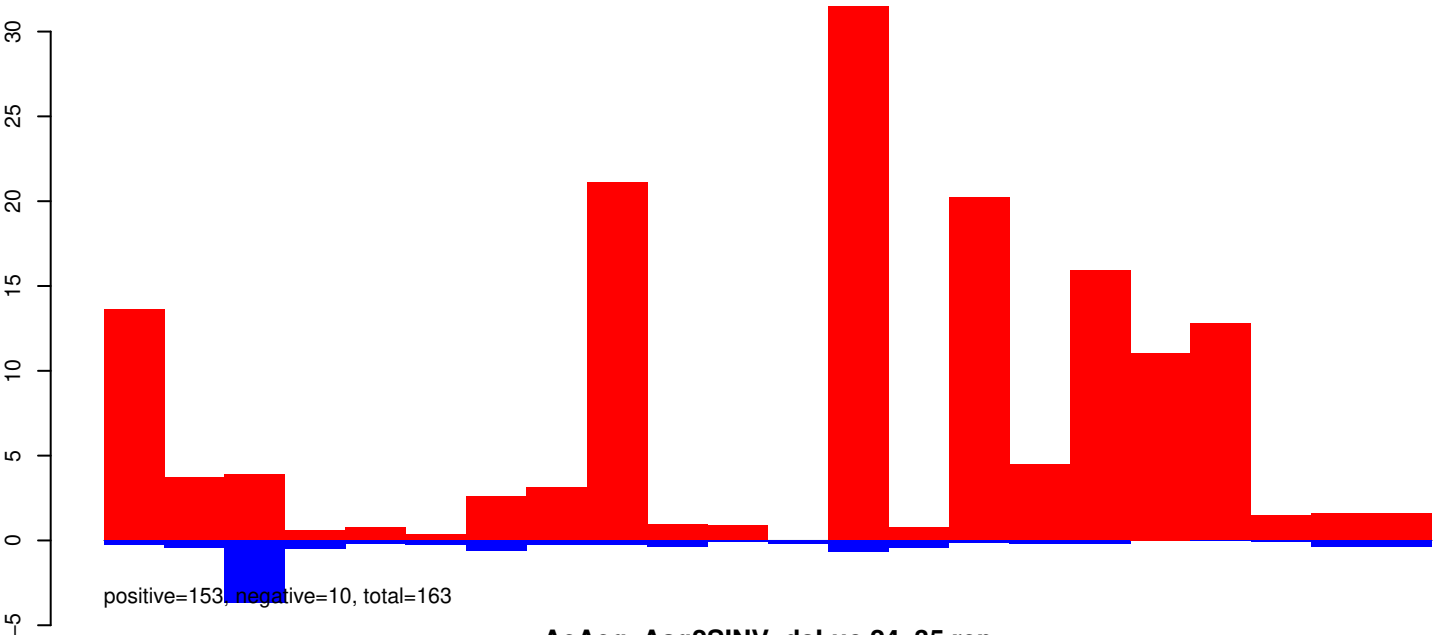
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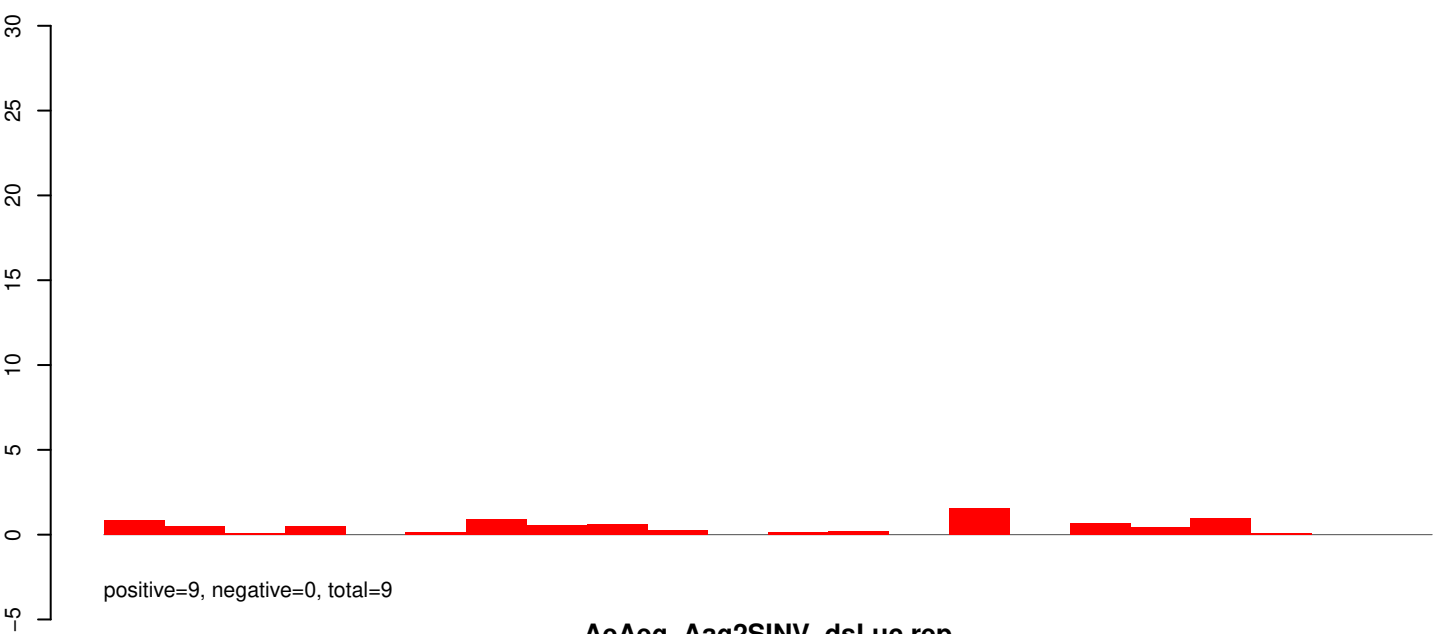
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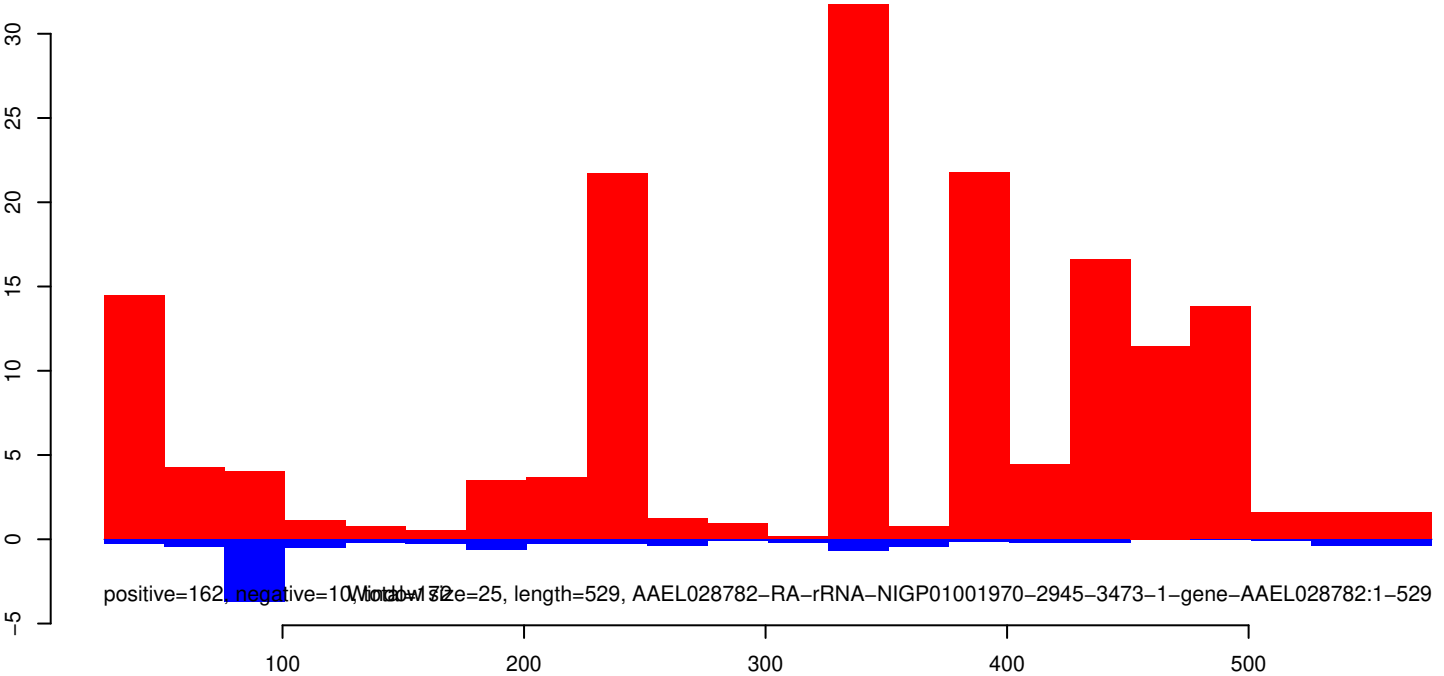
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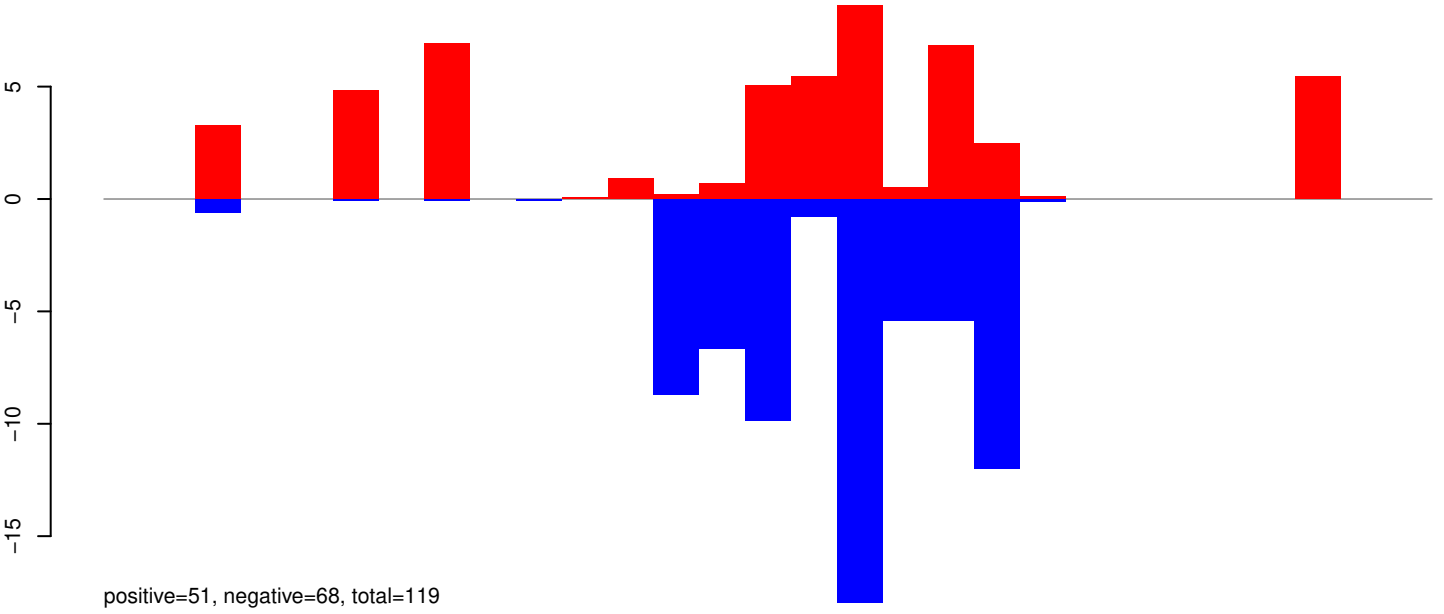
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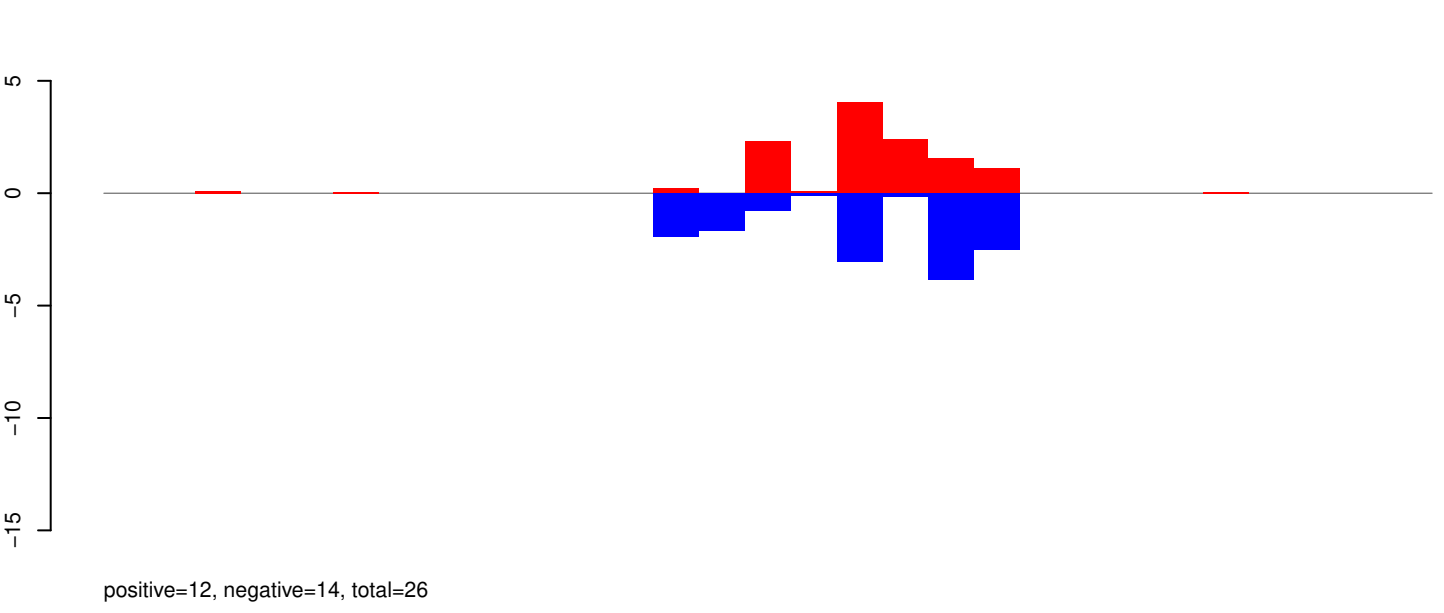
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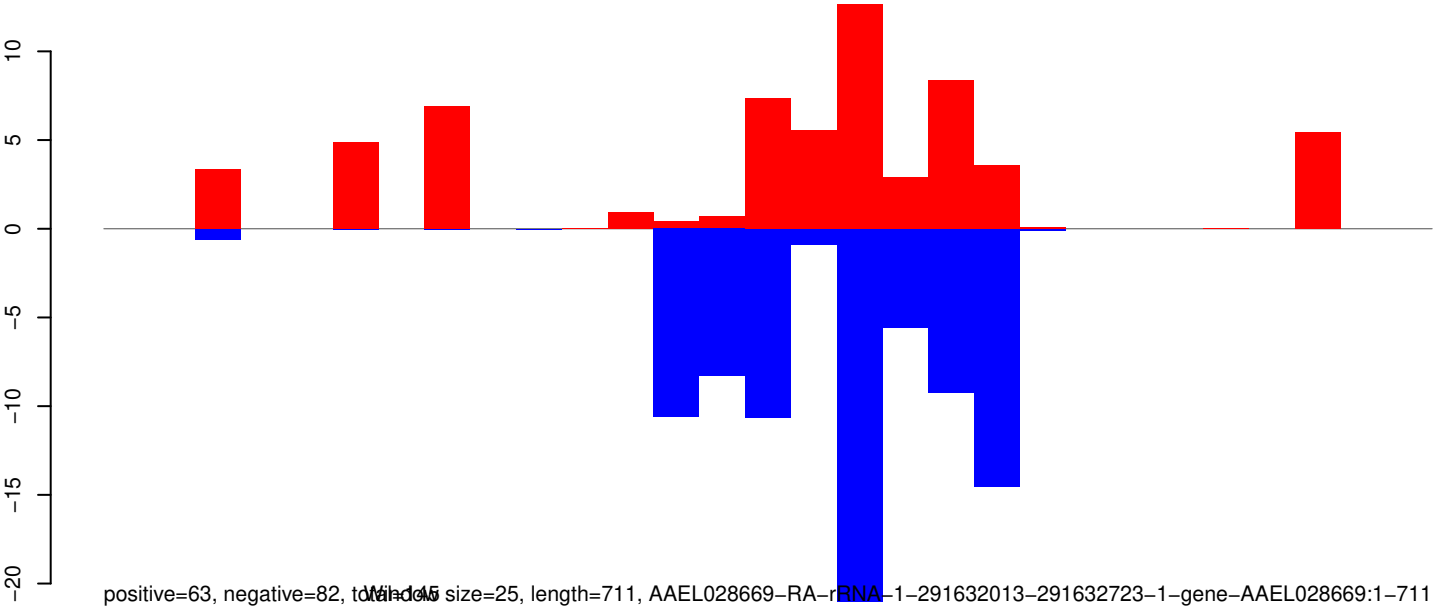
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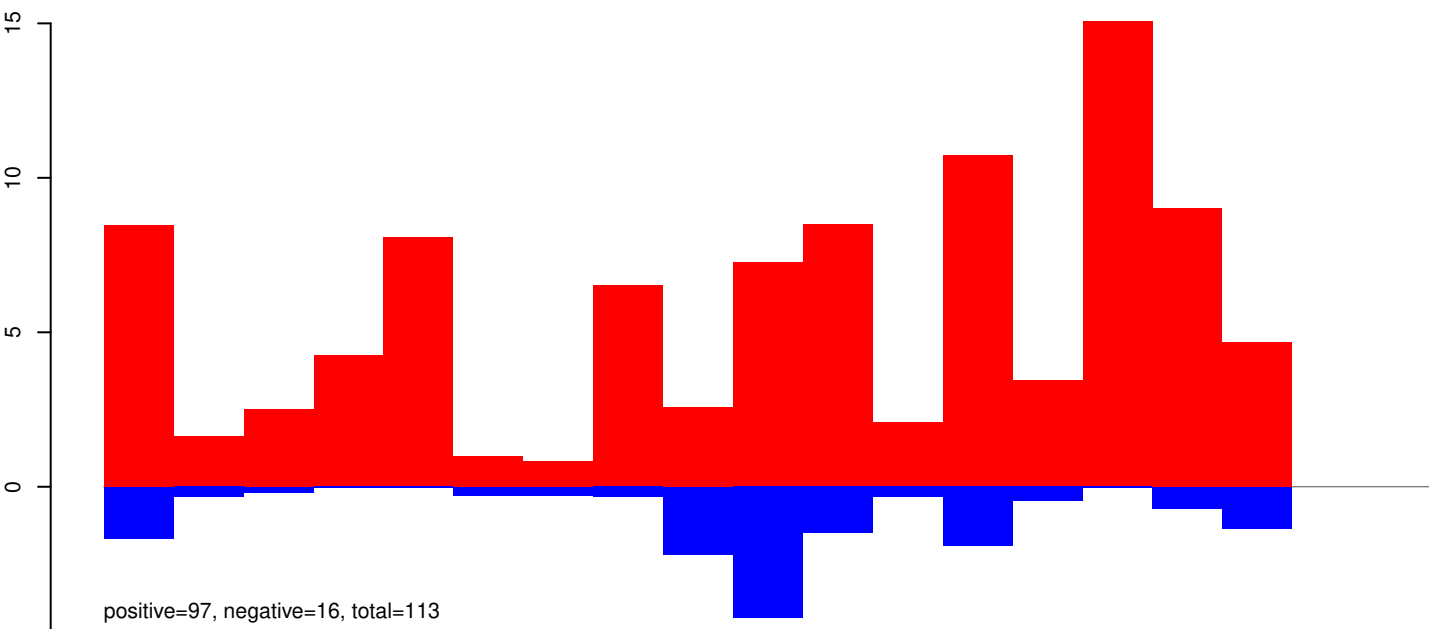
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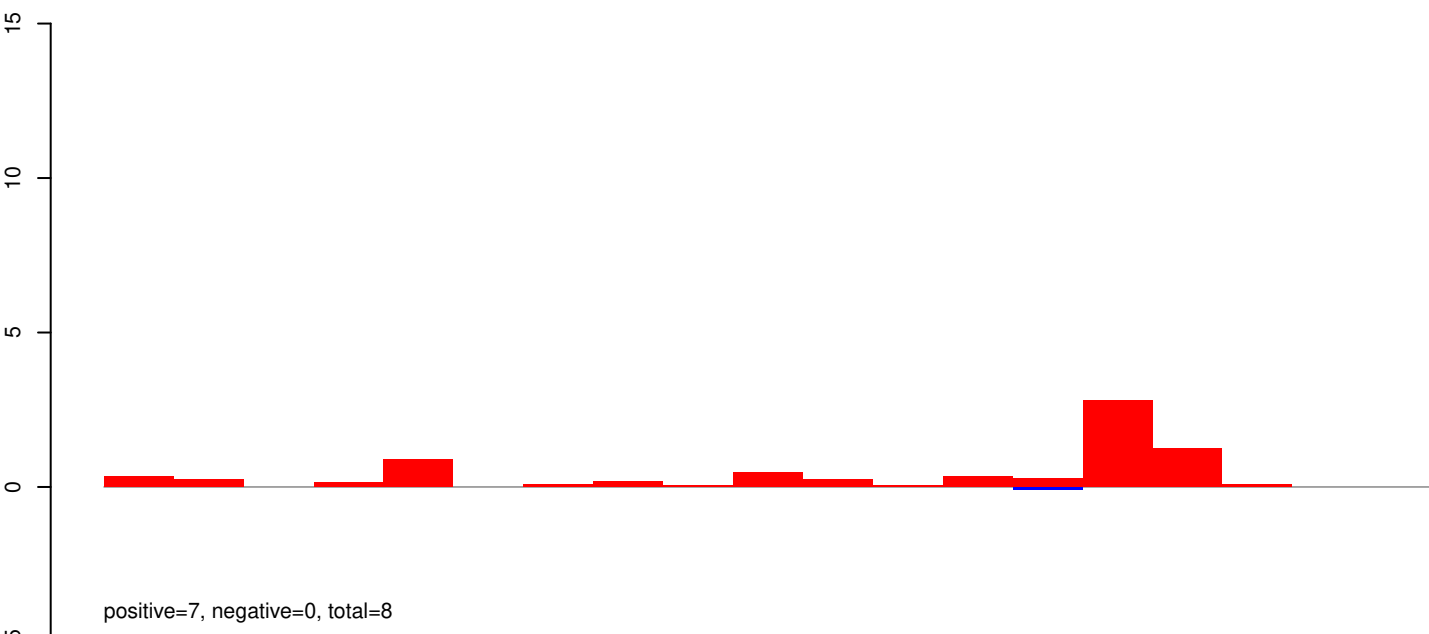
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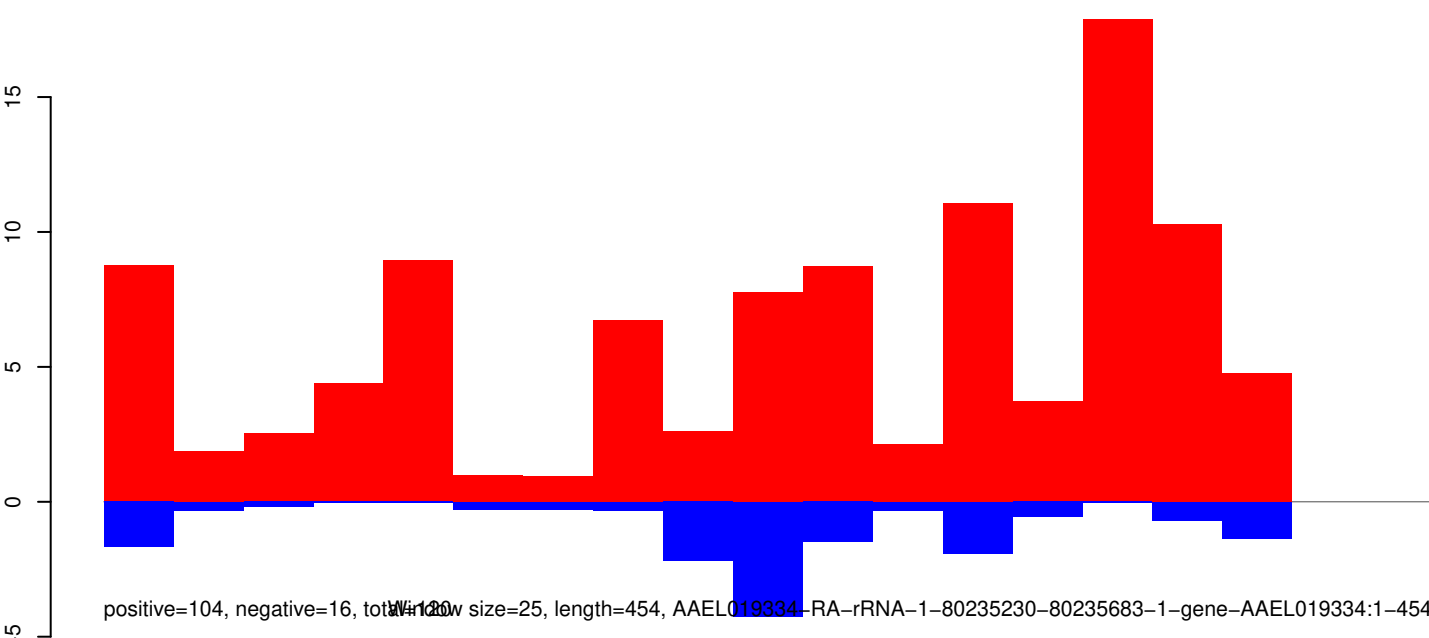
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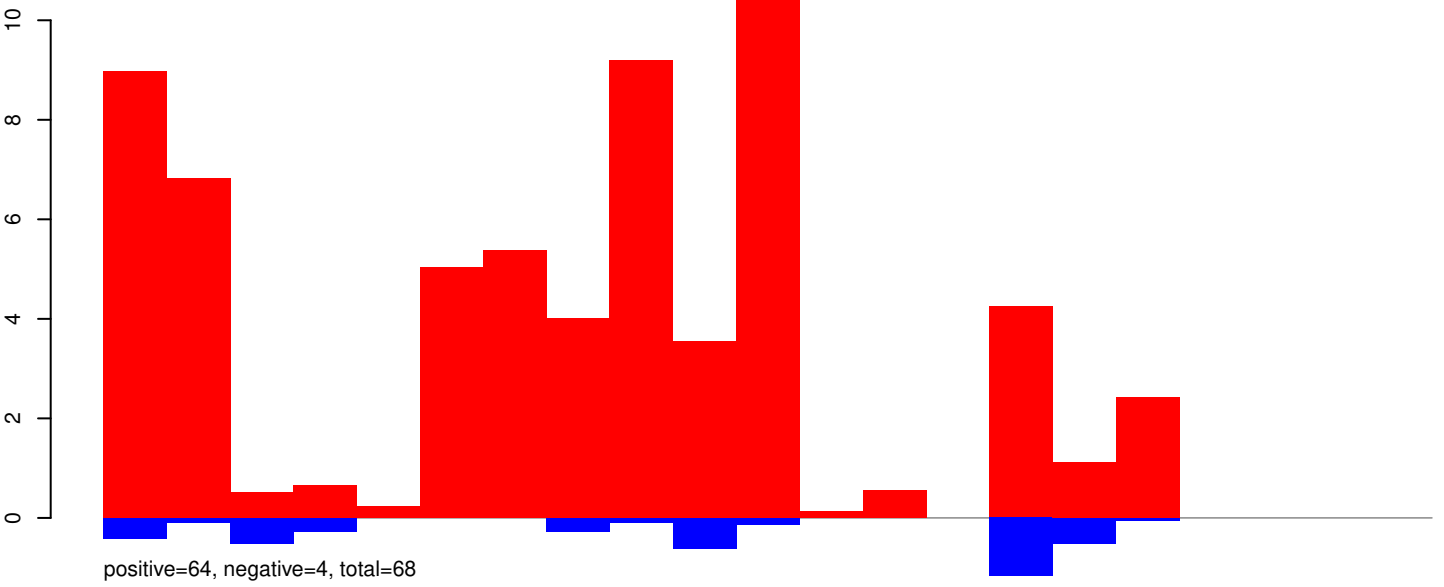


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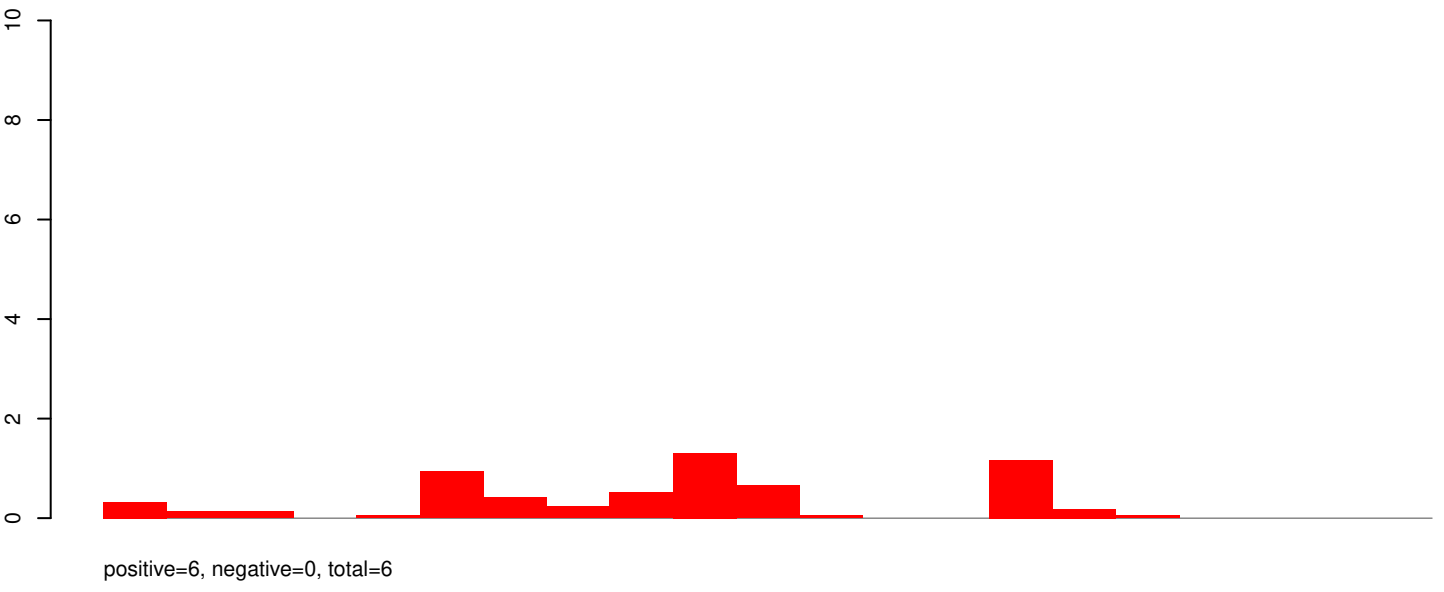


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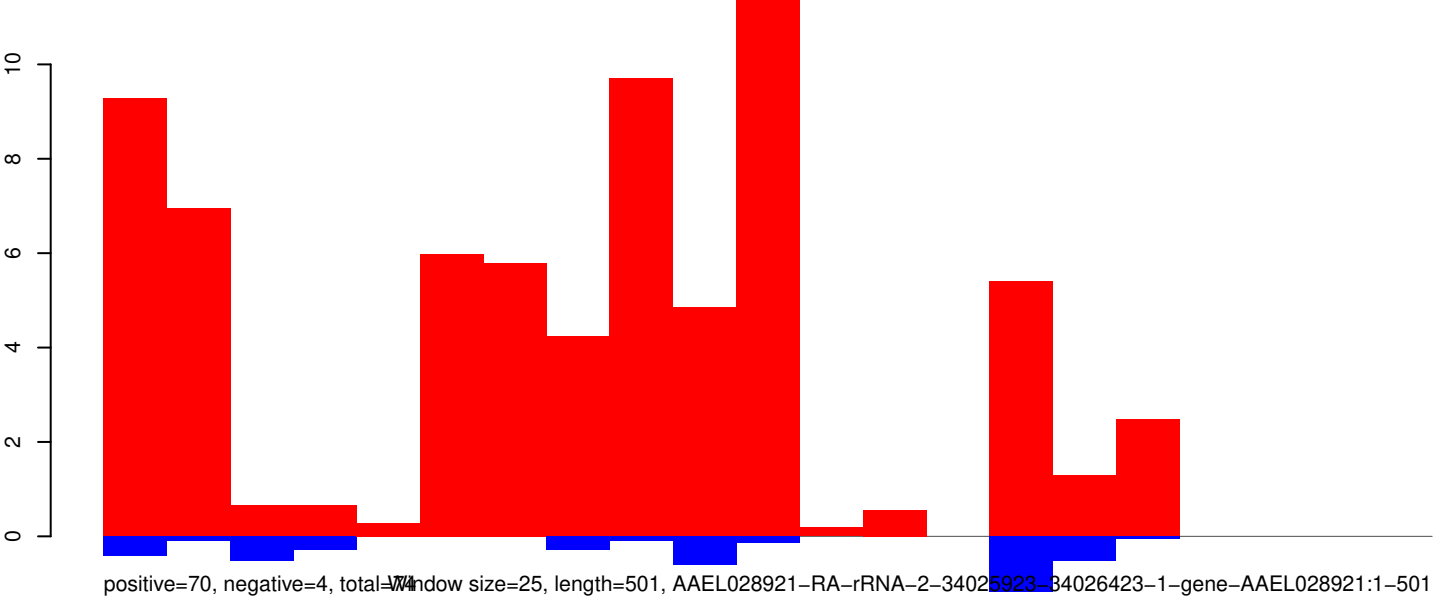
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AeAeg_Aag2SINV_dsLuc.24_35.rep

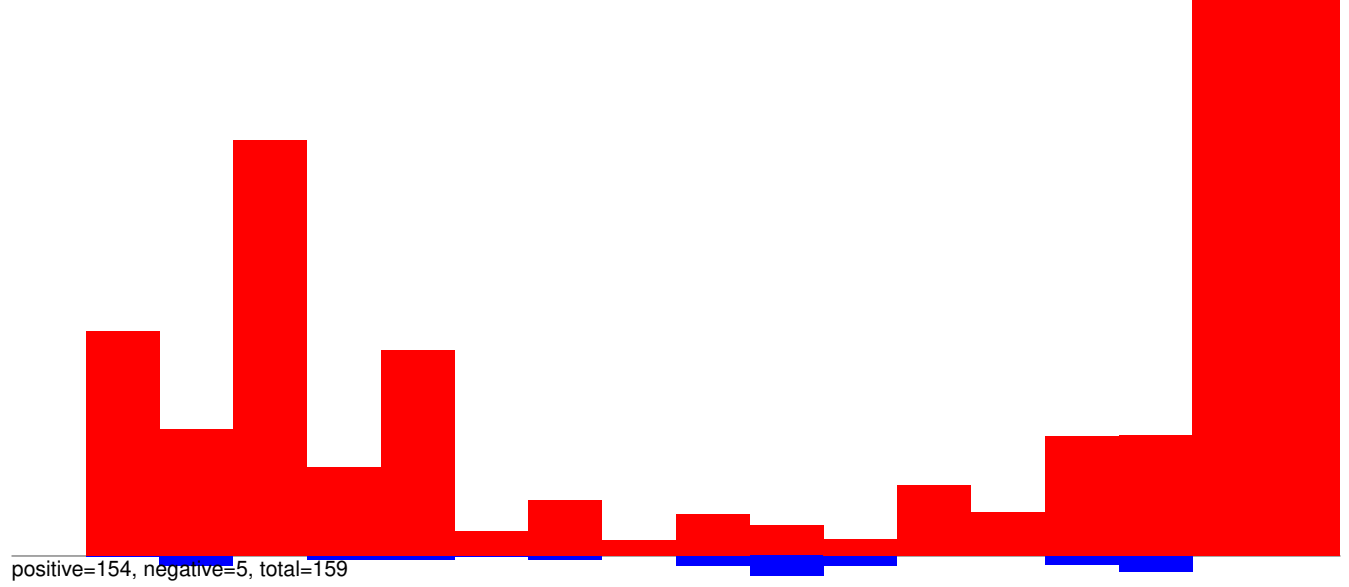


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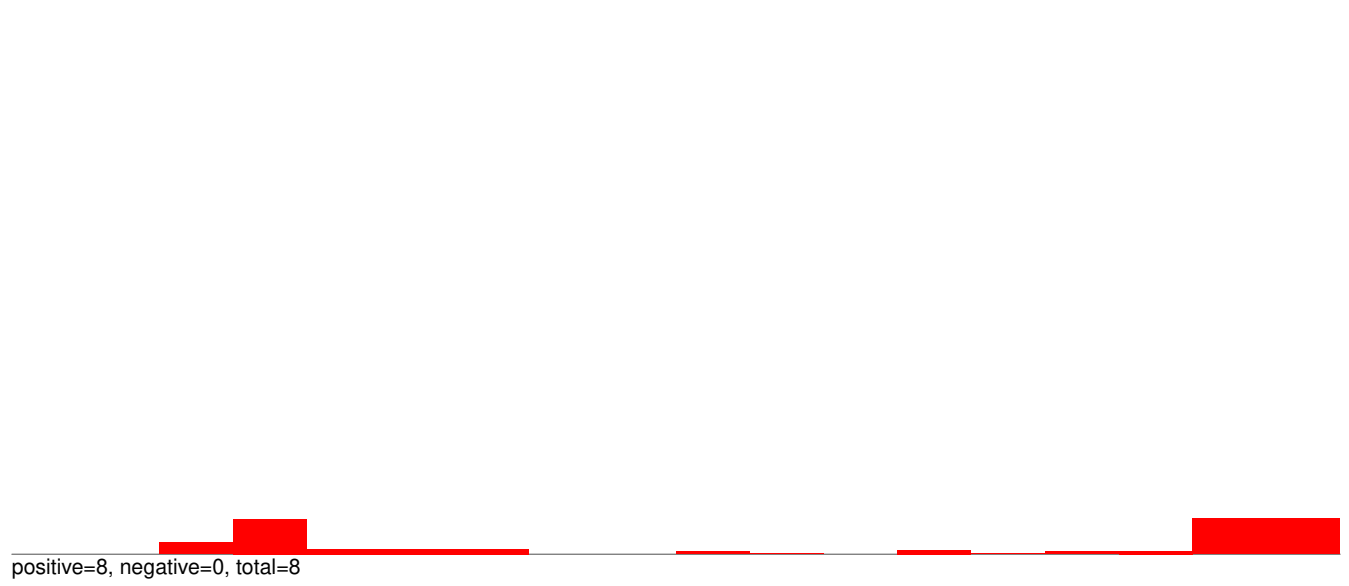


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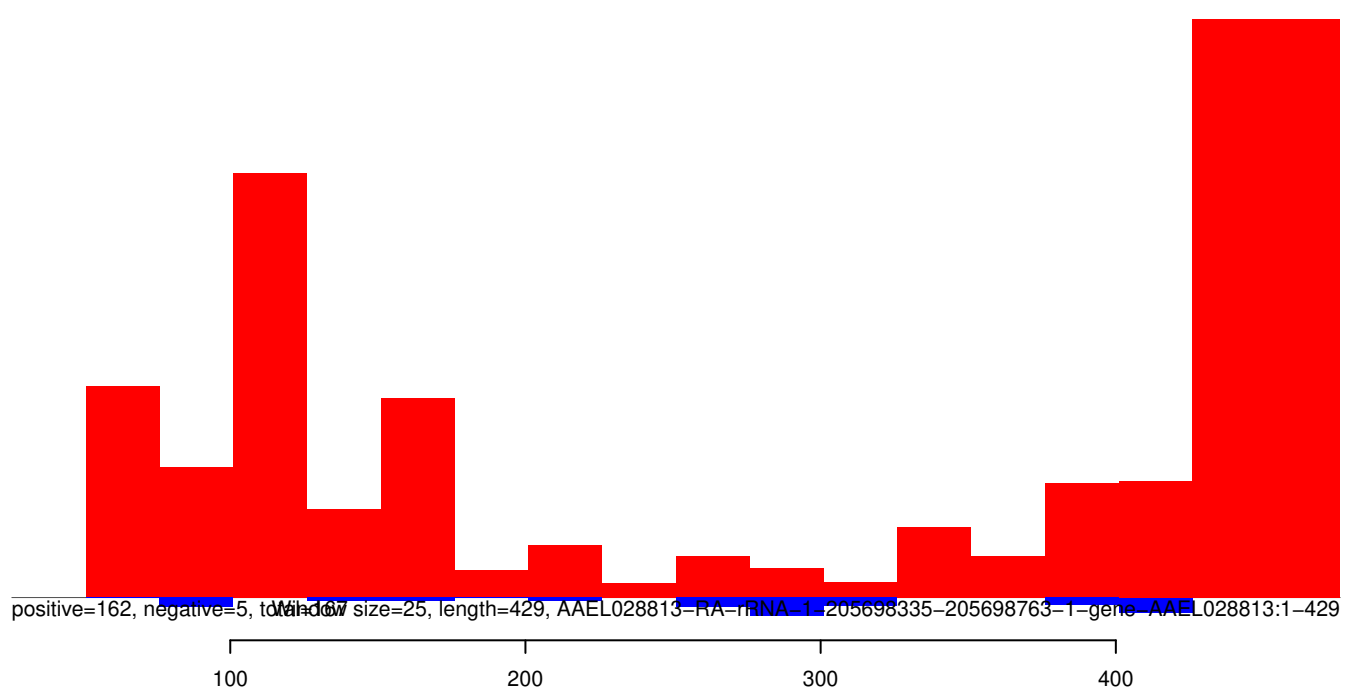
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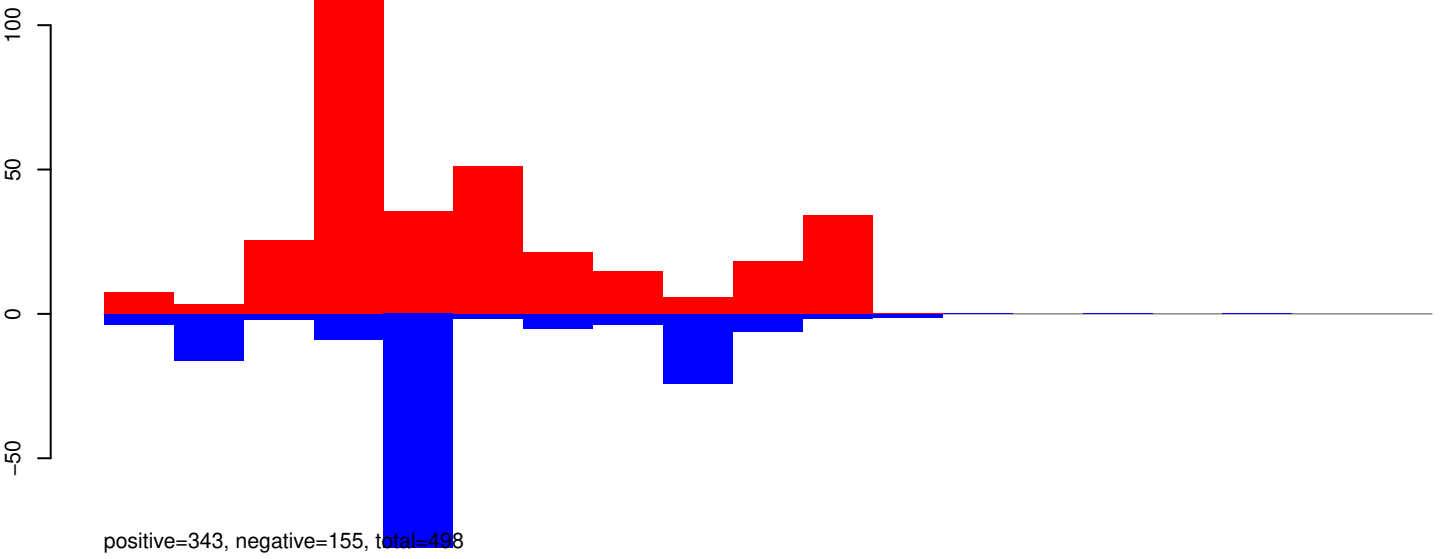
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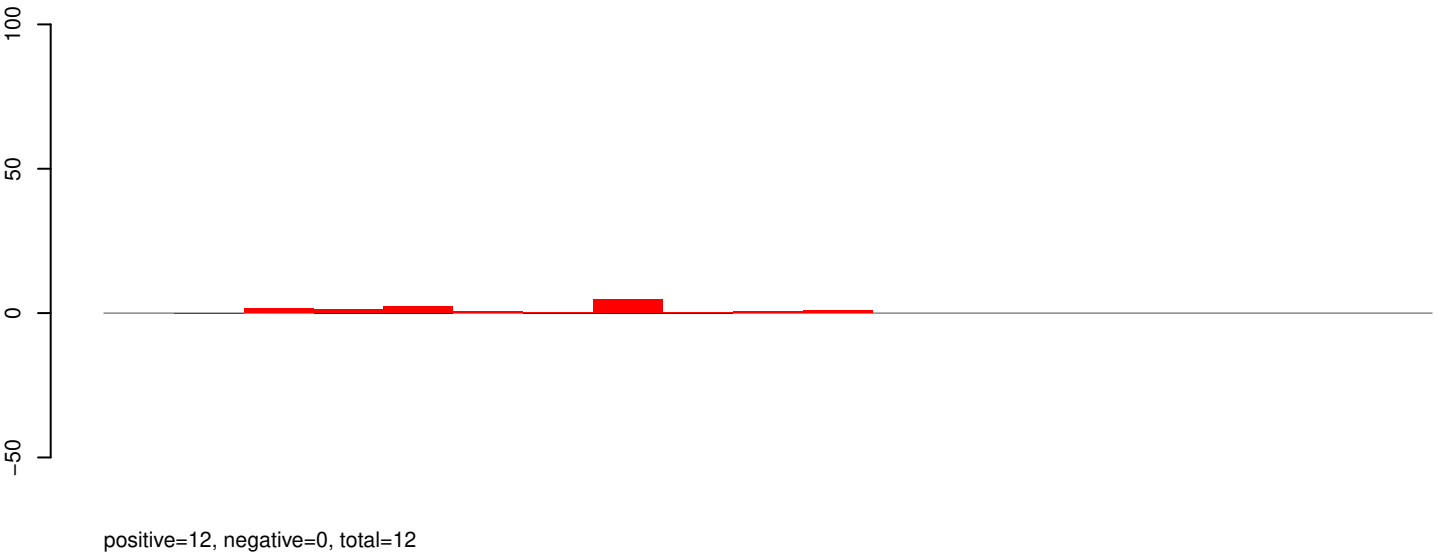
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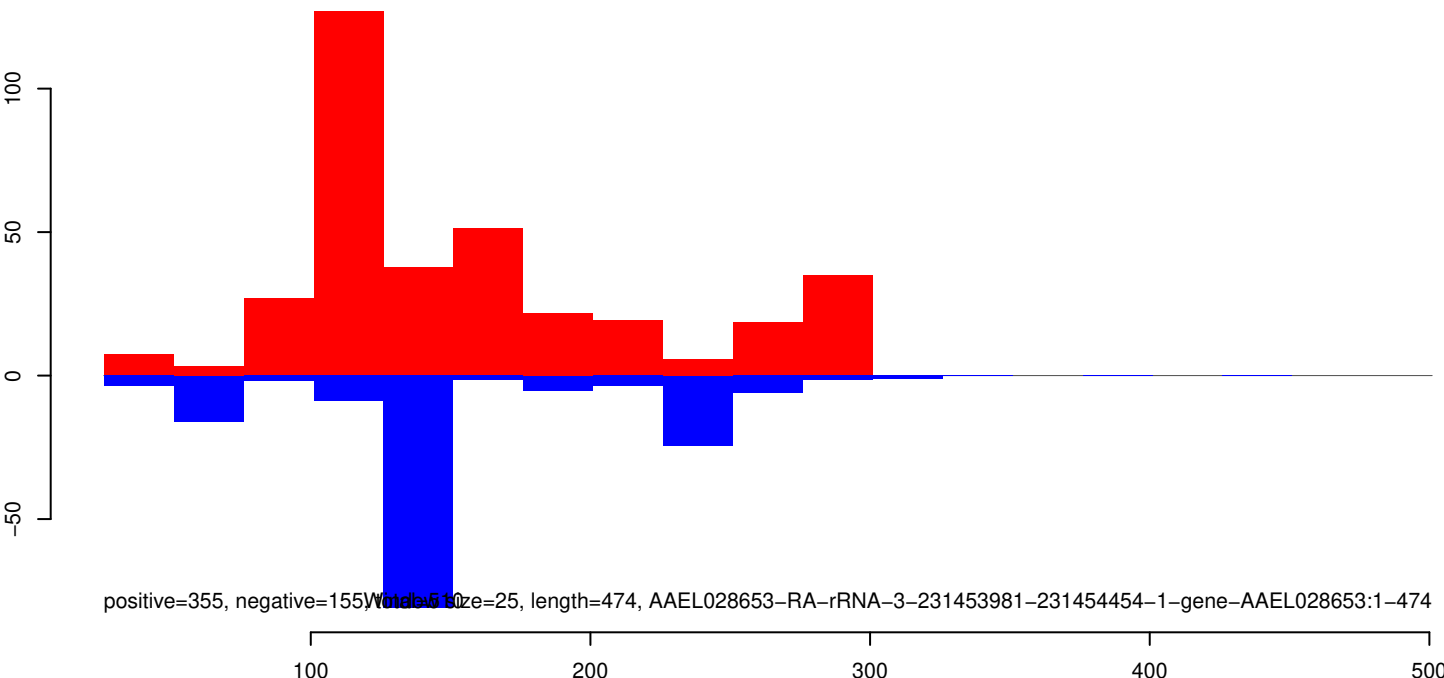
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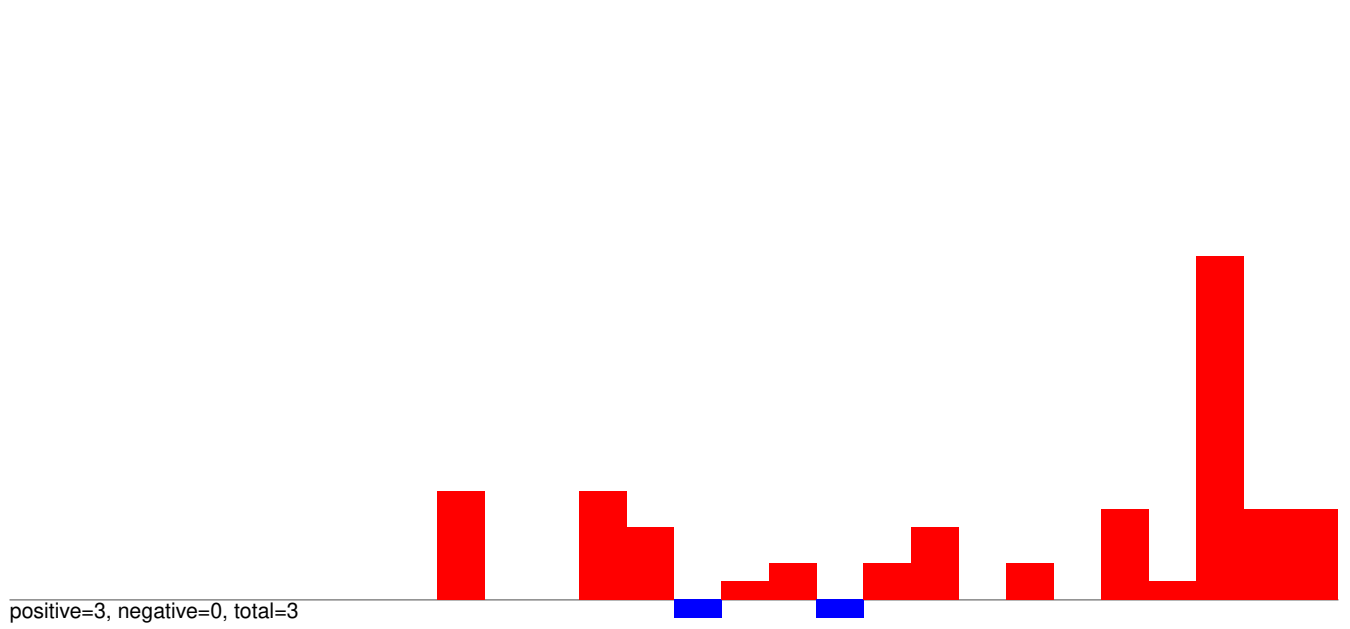
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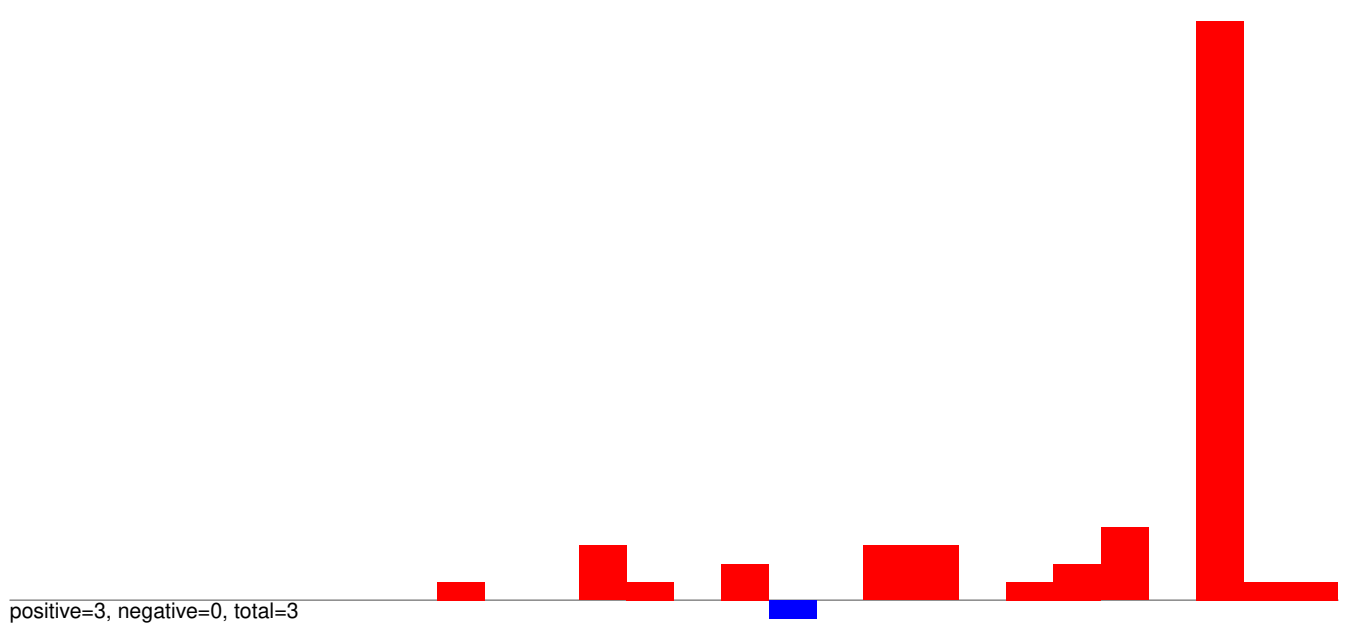
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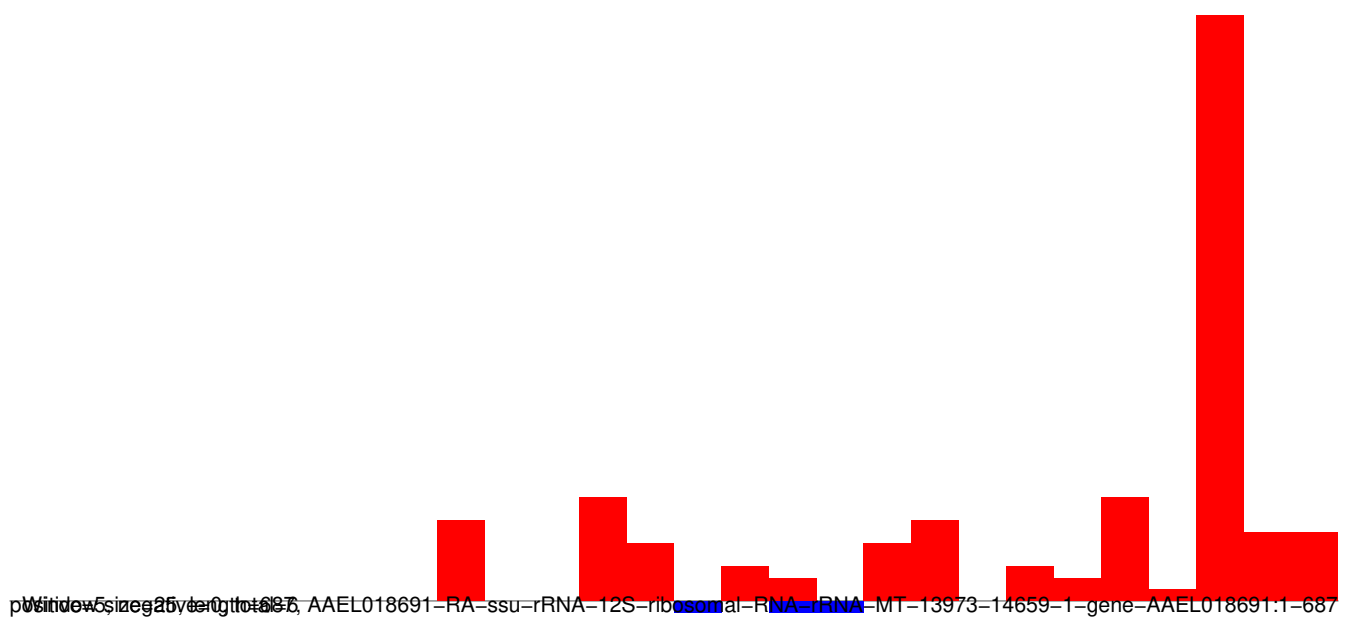
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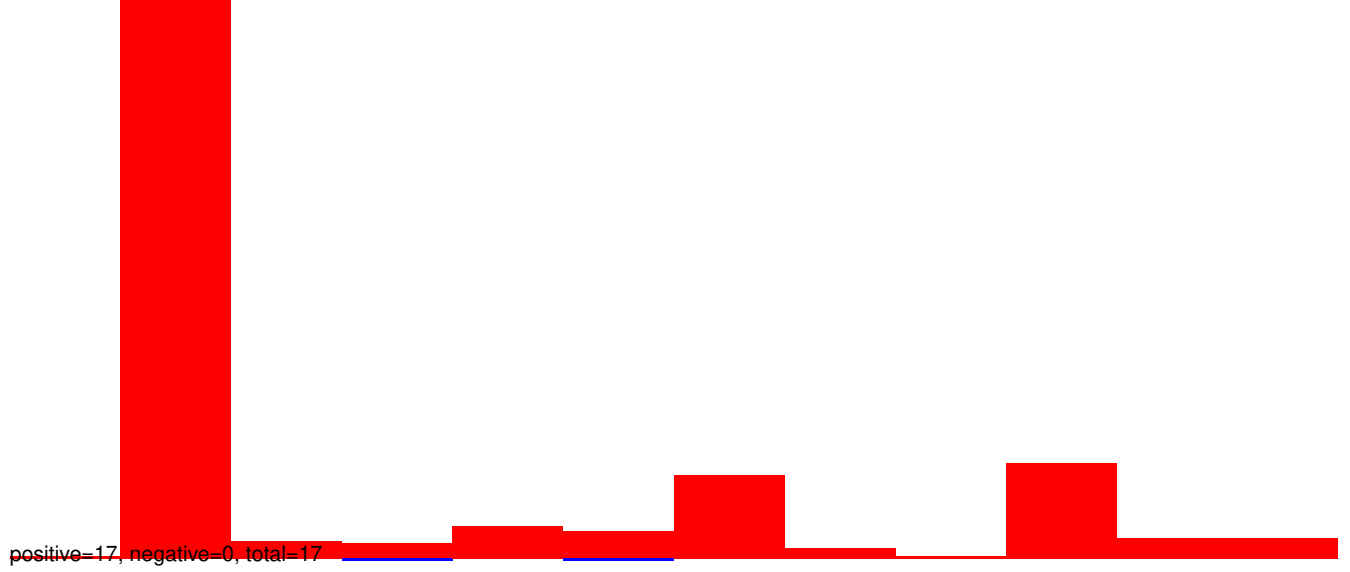
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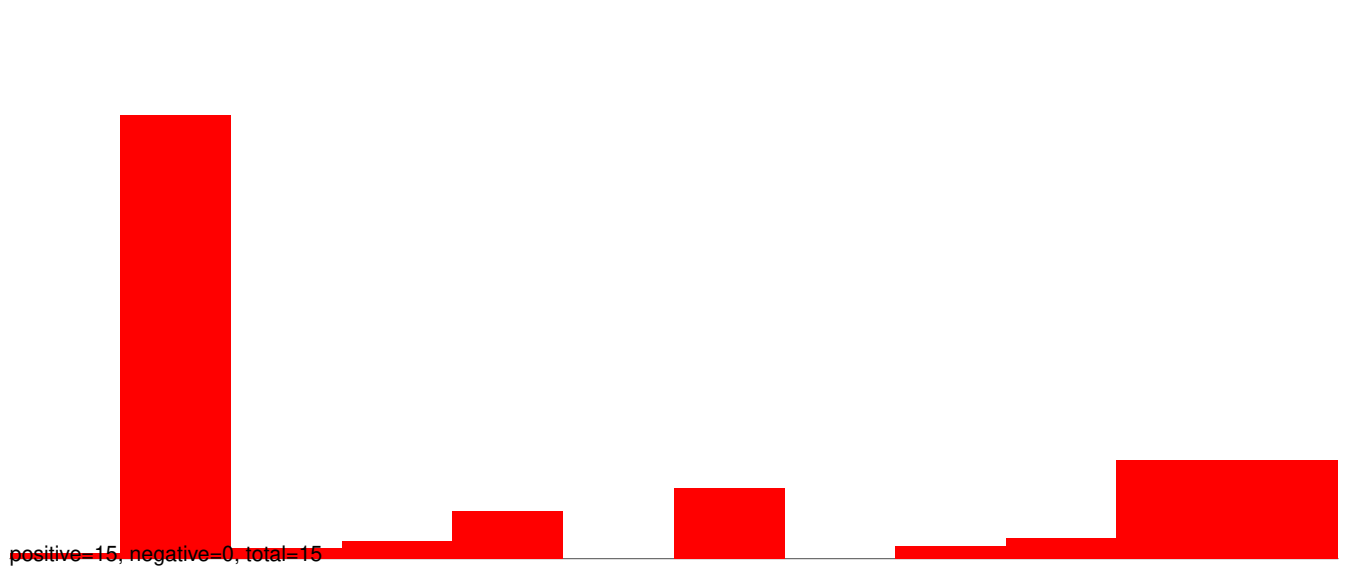
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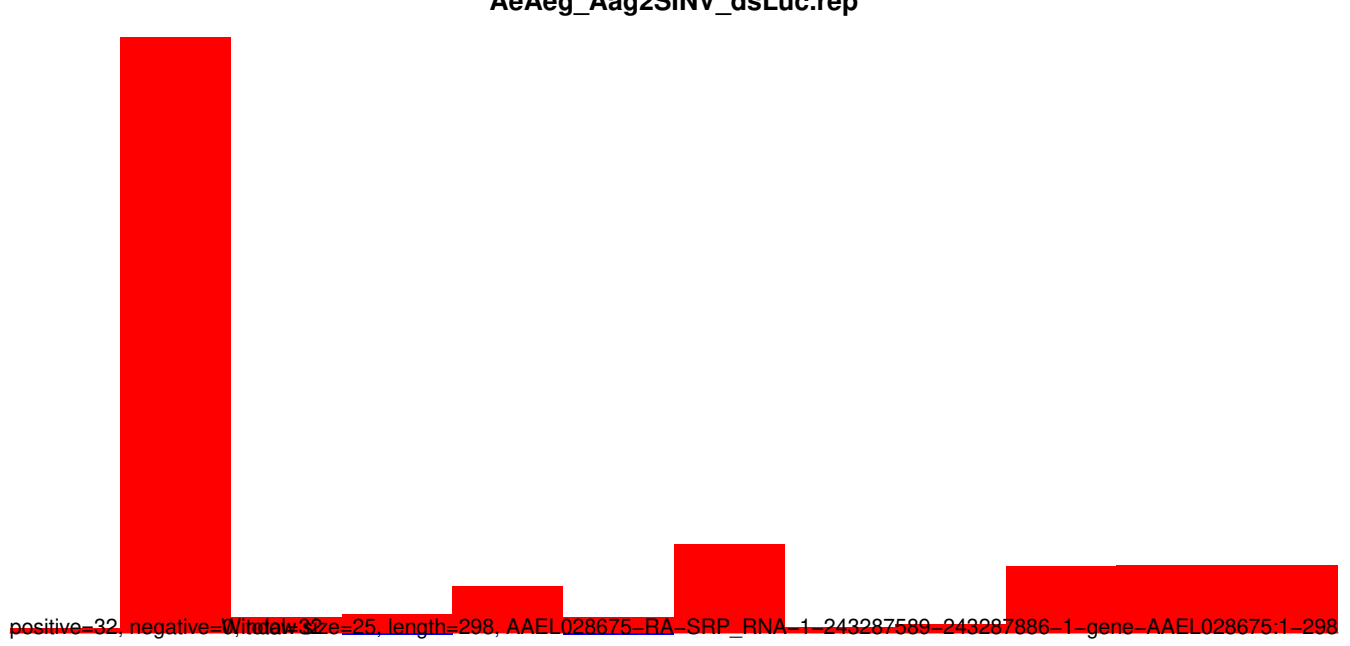
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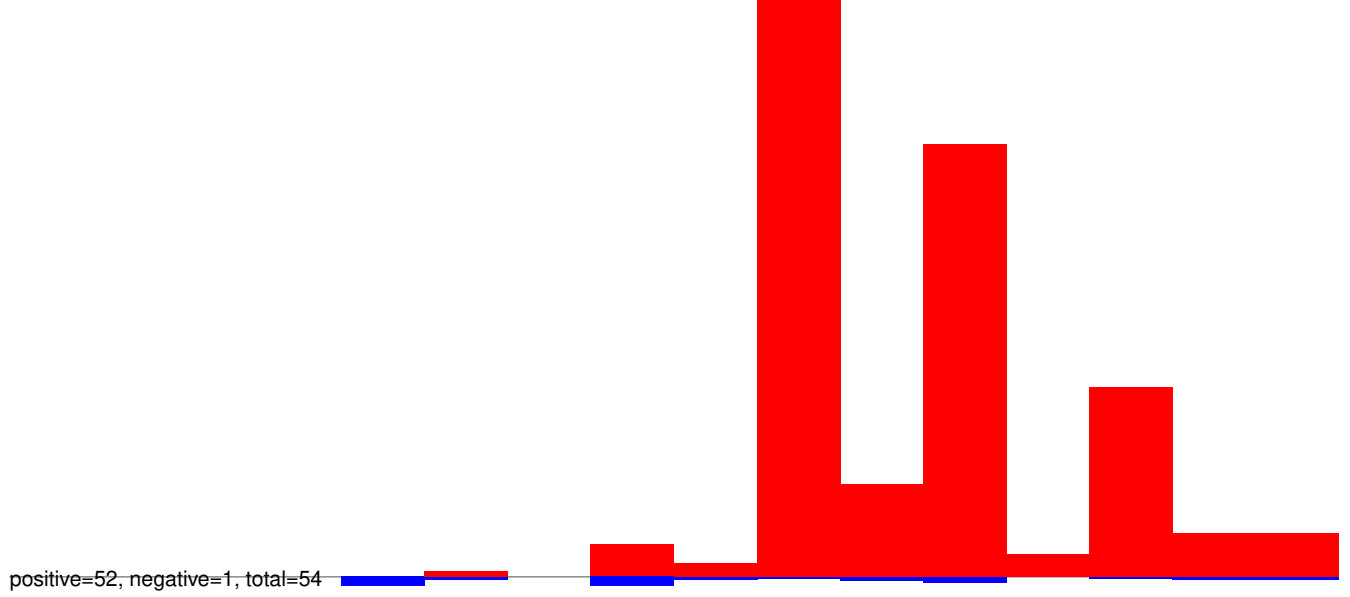
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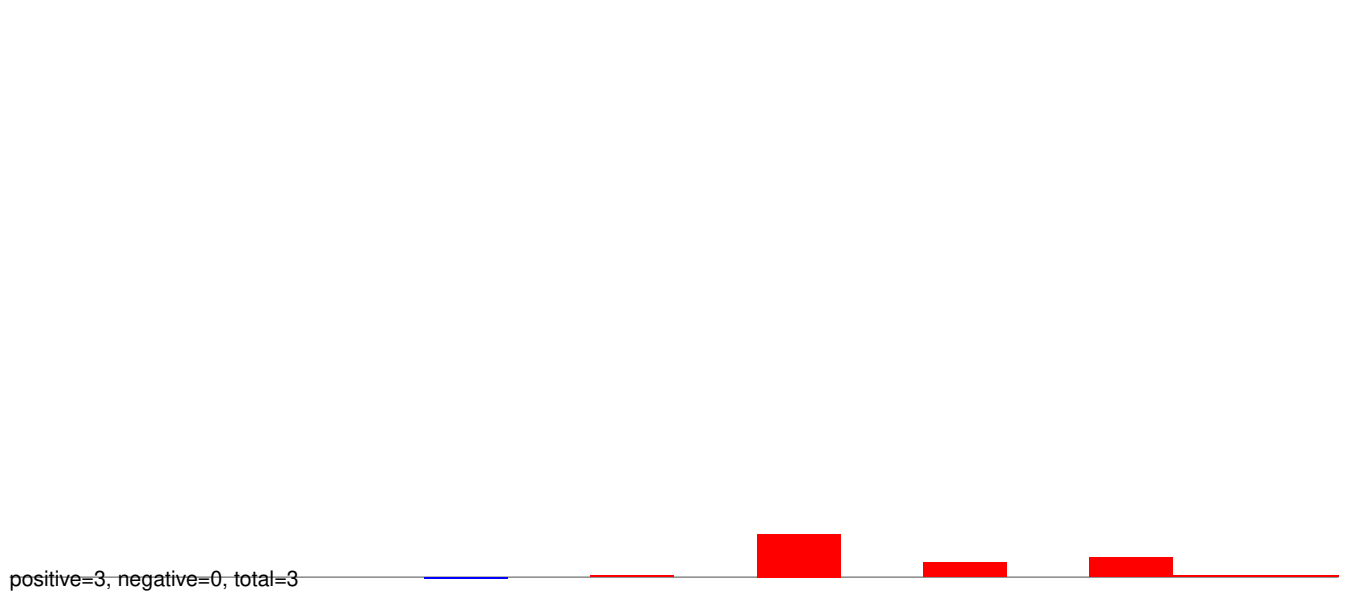
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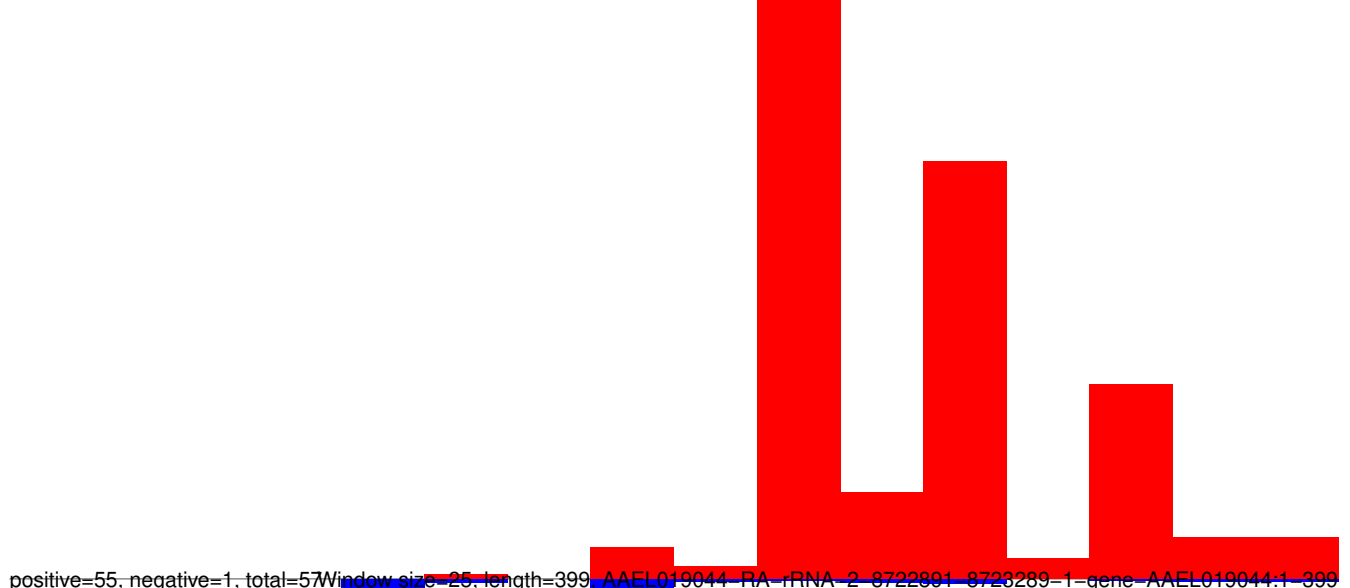
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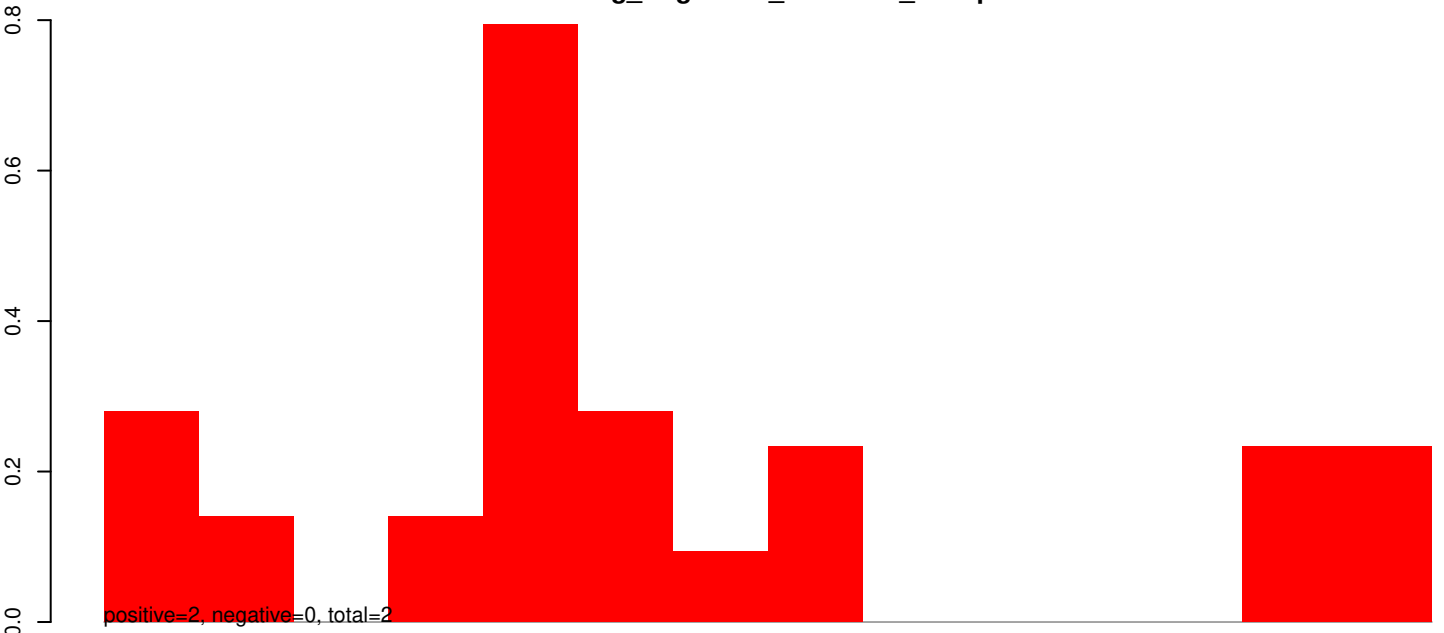


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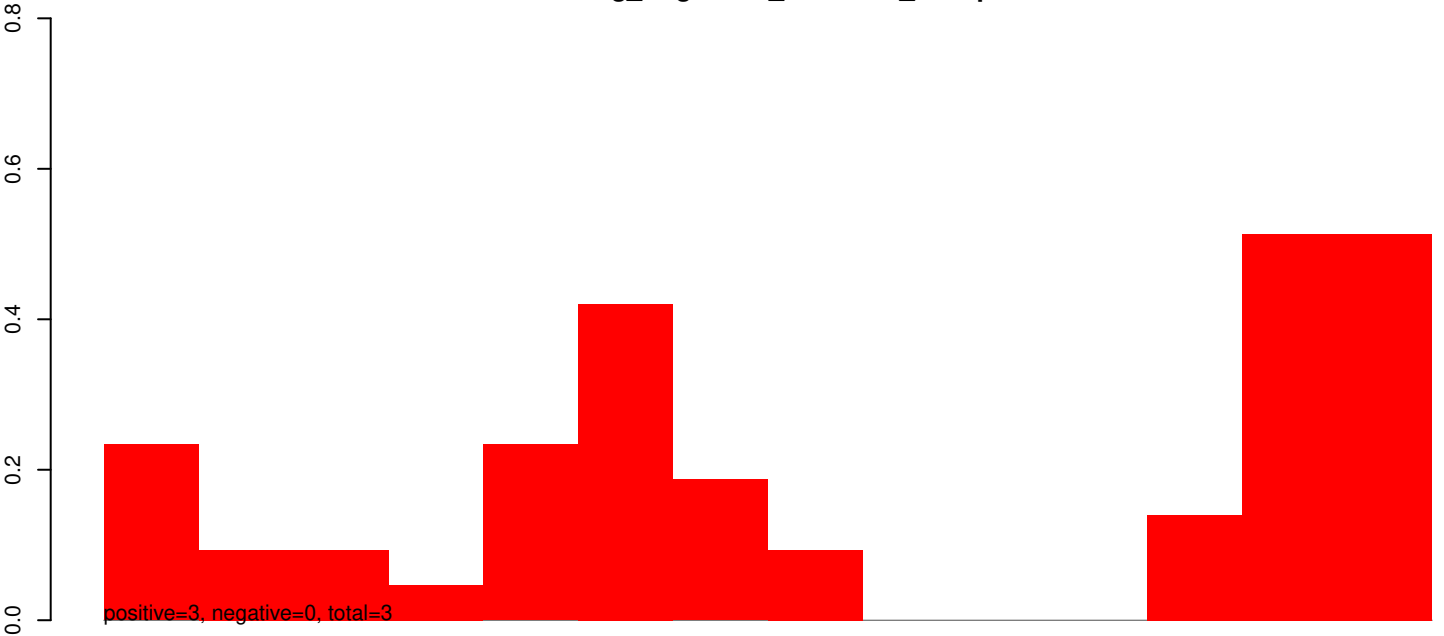


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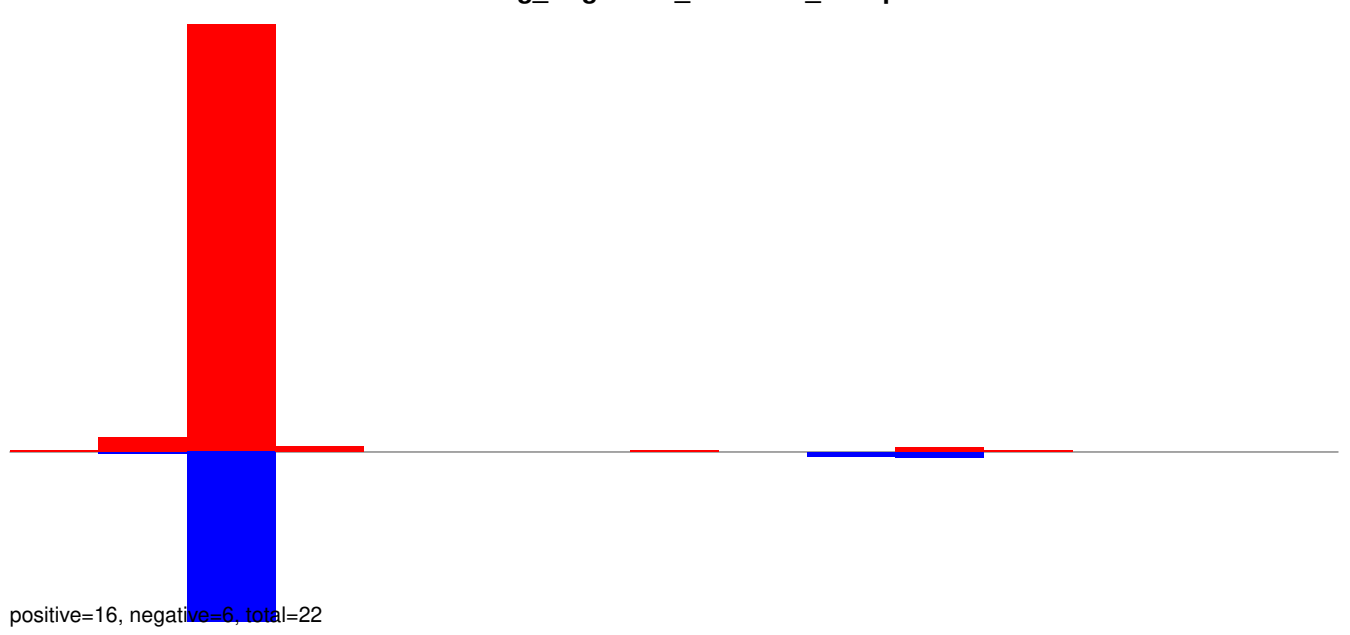
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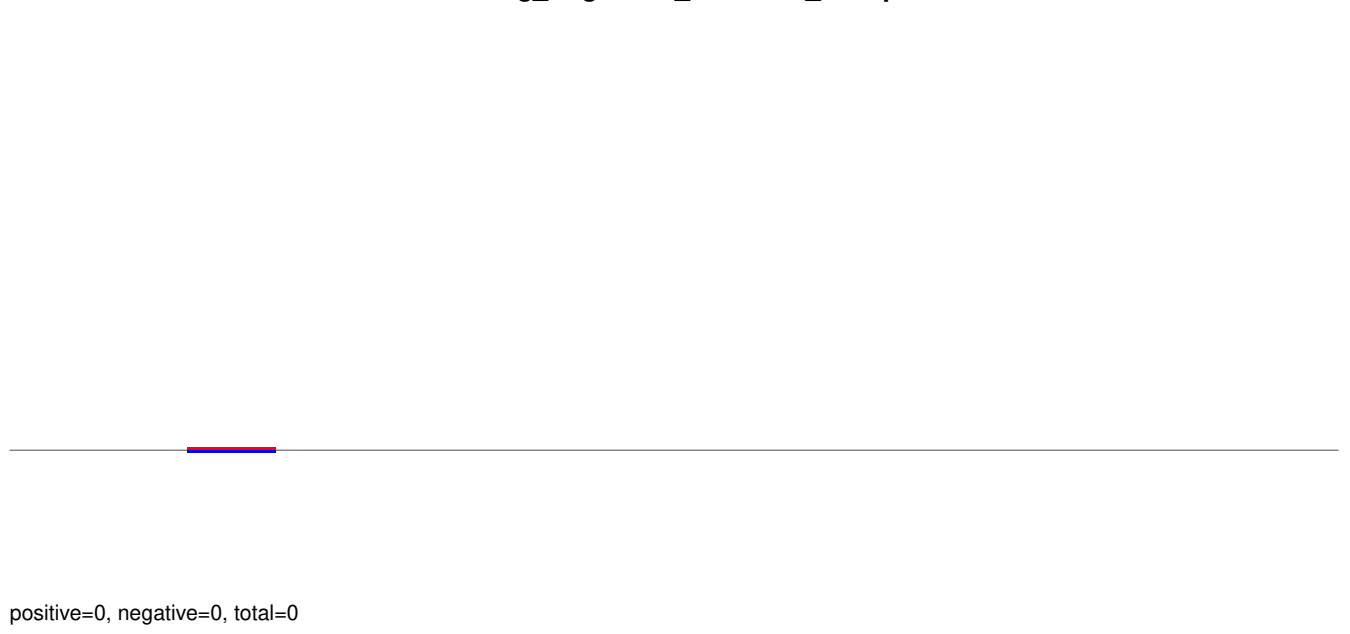
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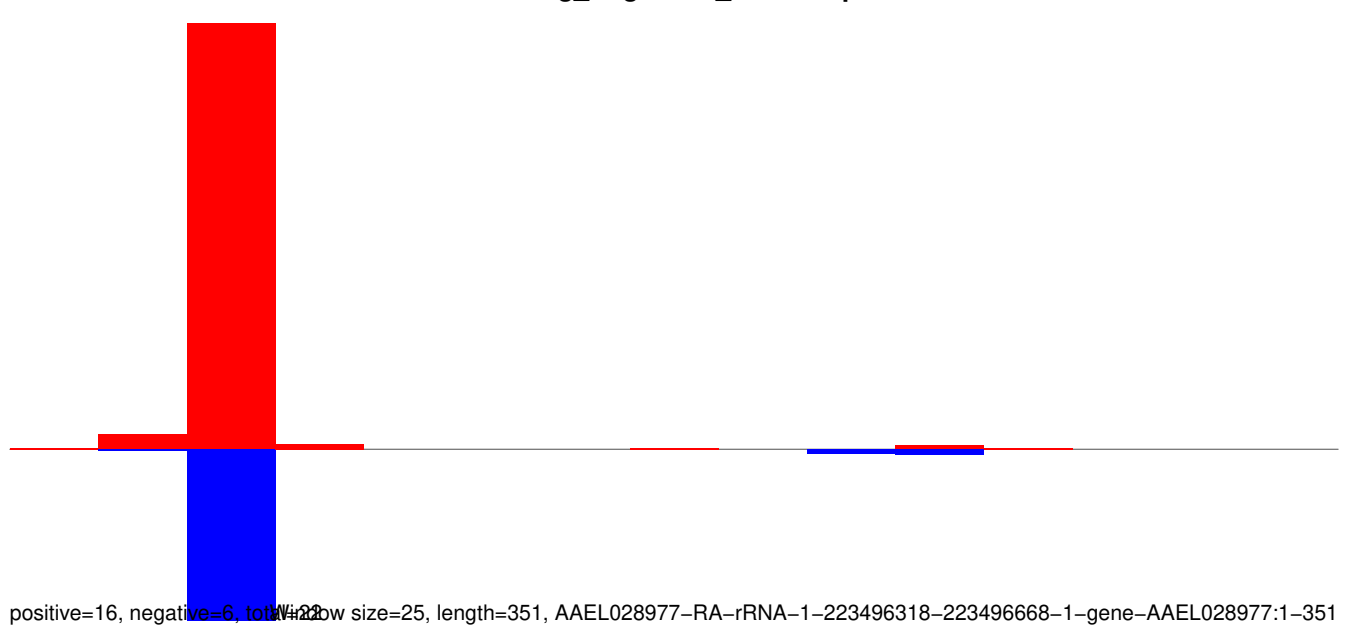
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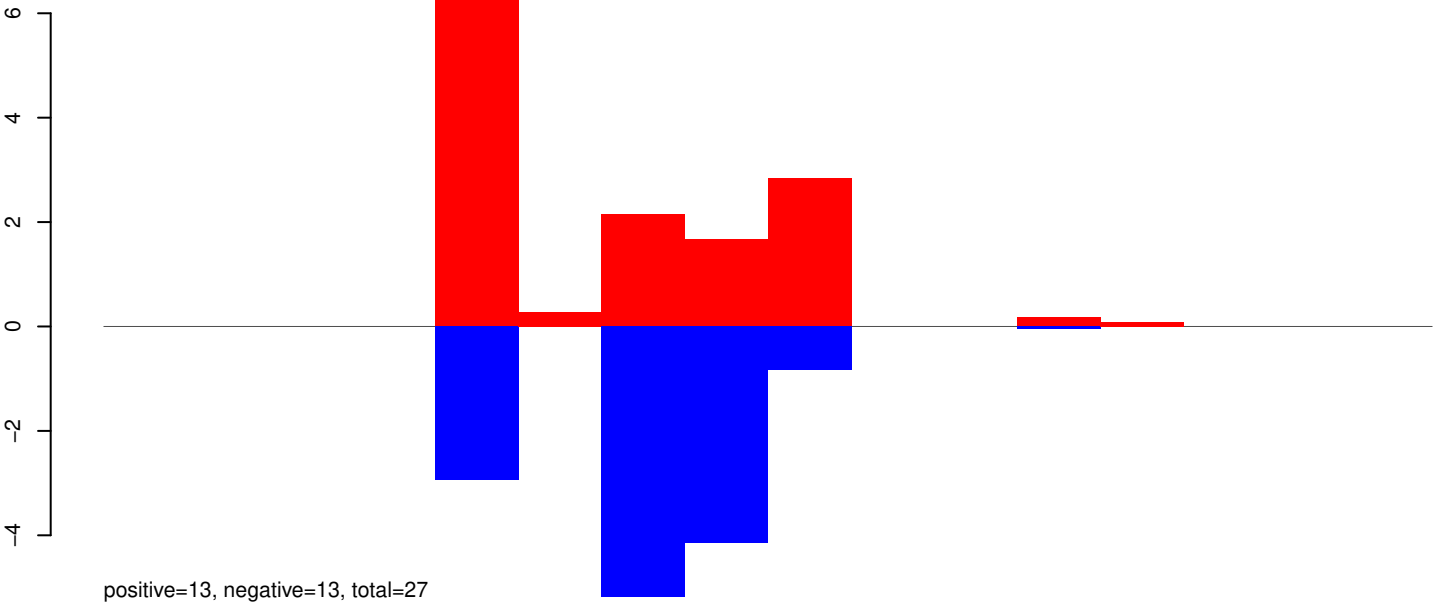
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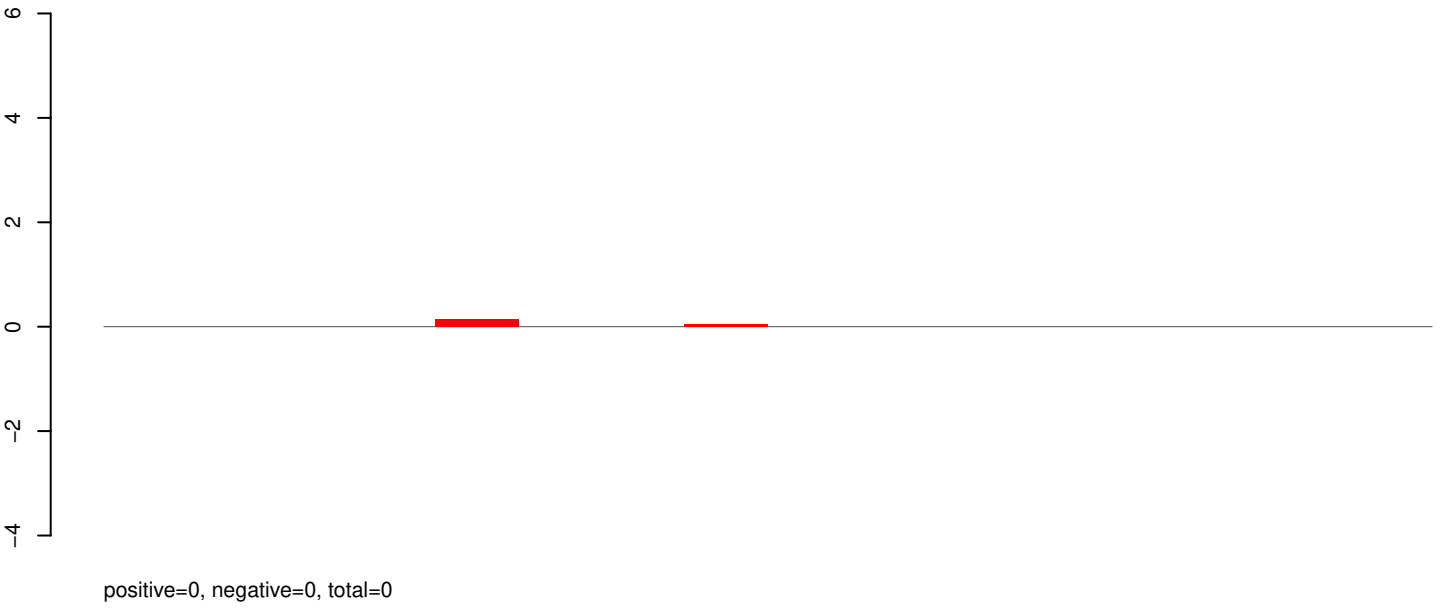
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100 200 300 400

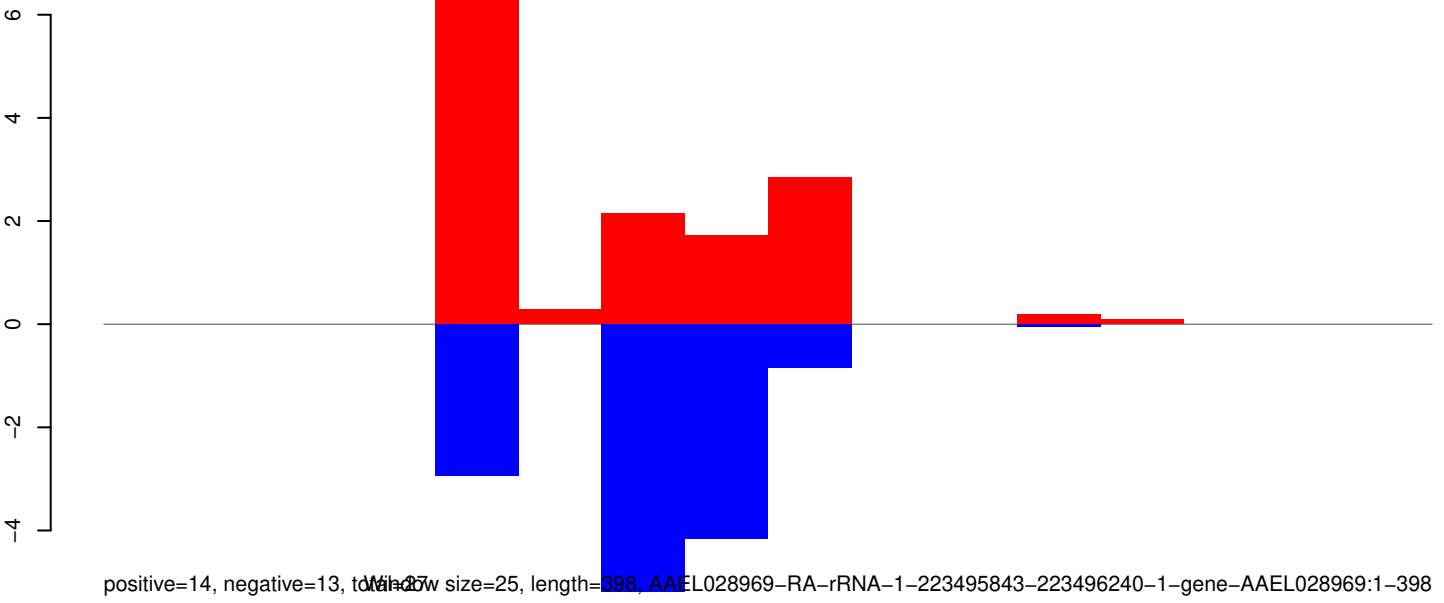
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AeAeg_Aag2SINV_dsLuc.24_35.rep

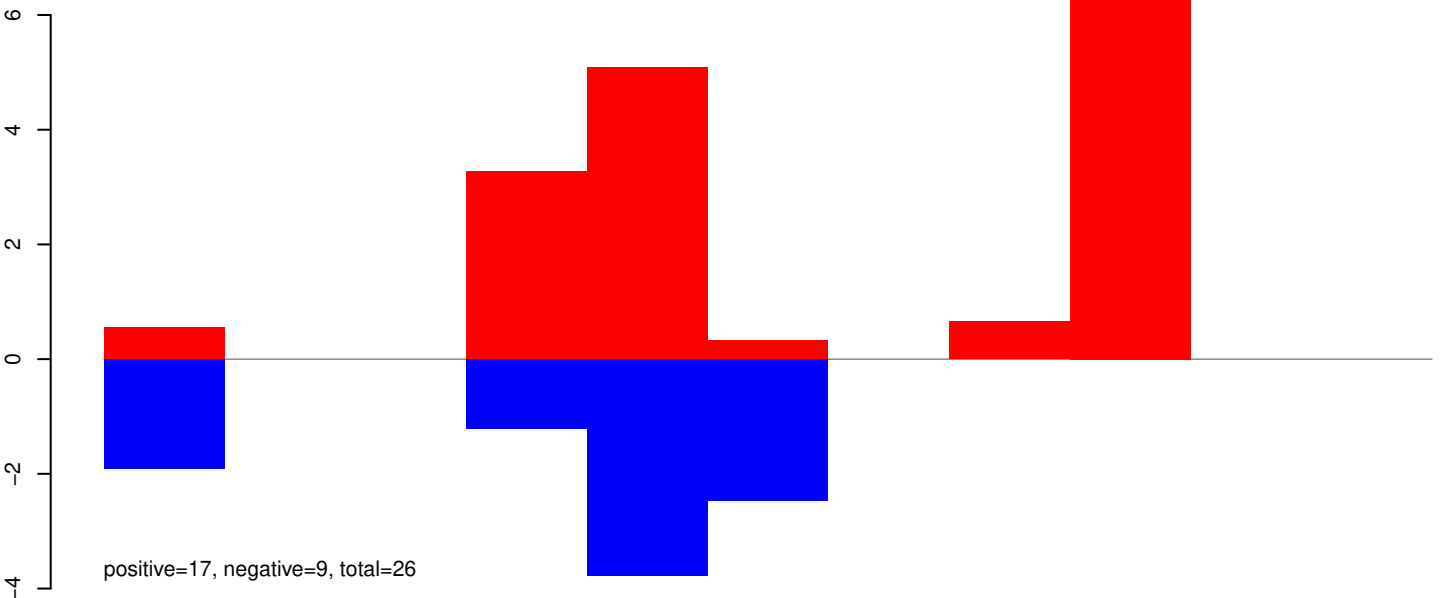


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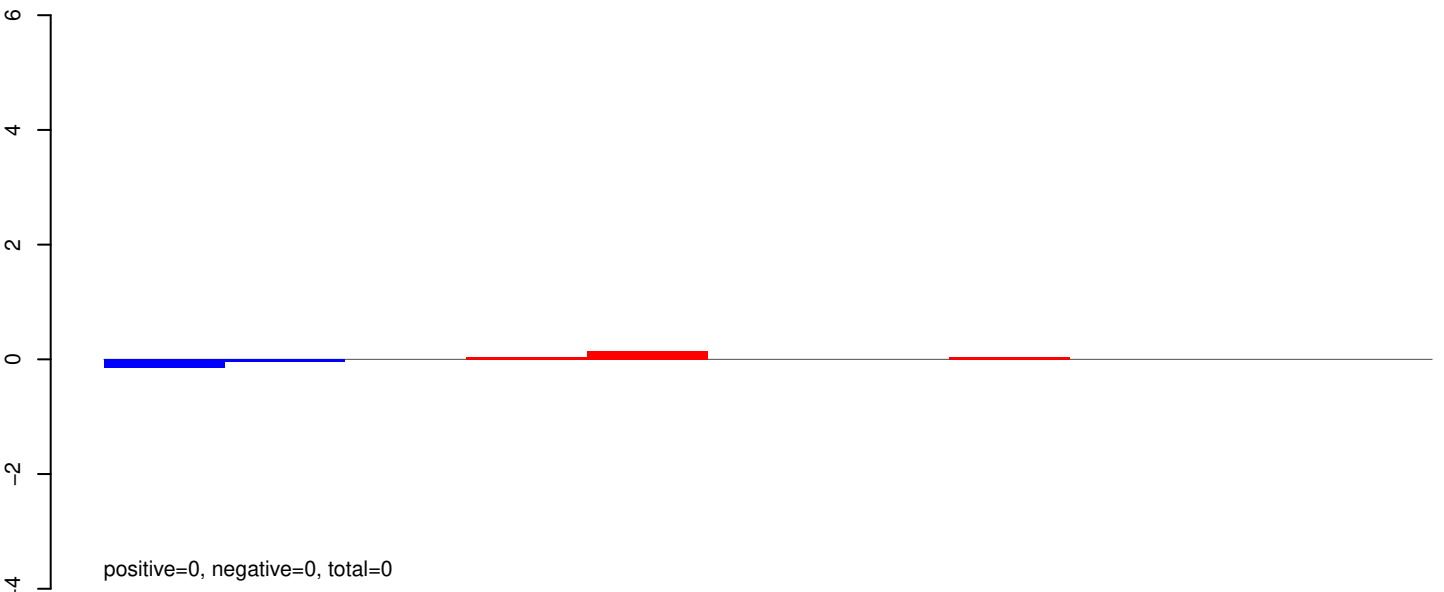


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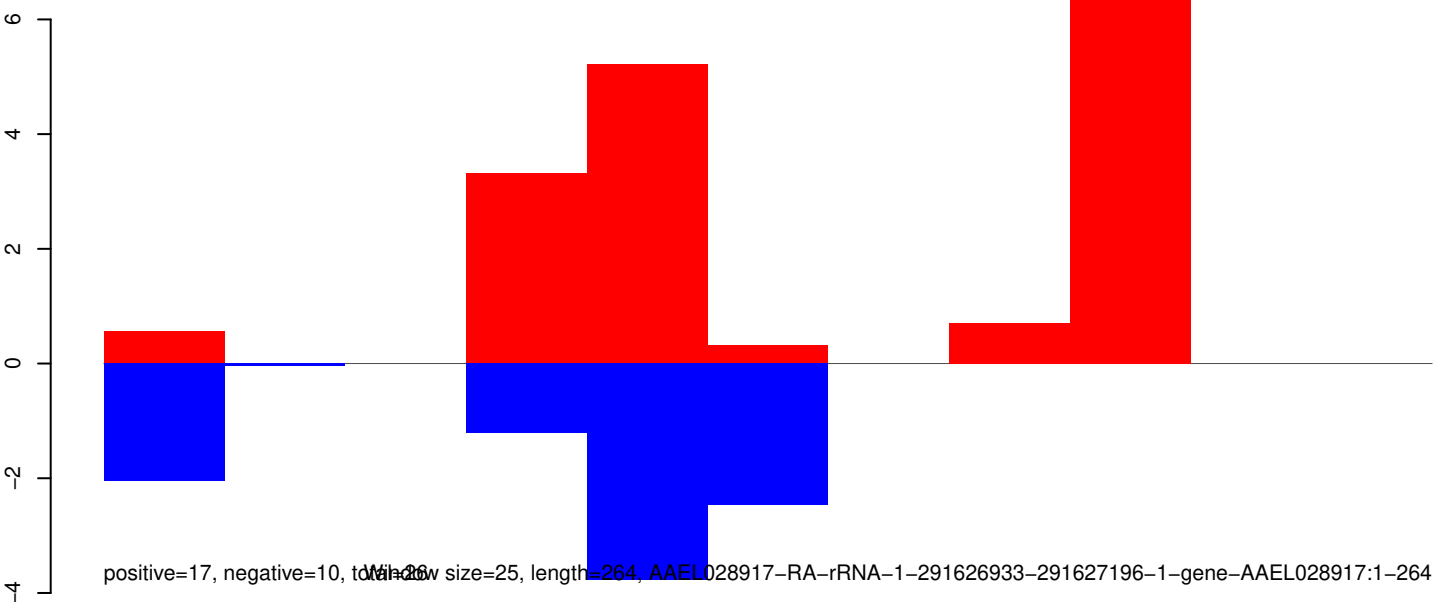
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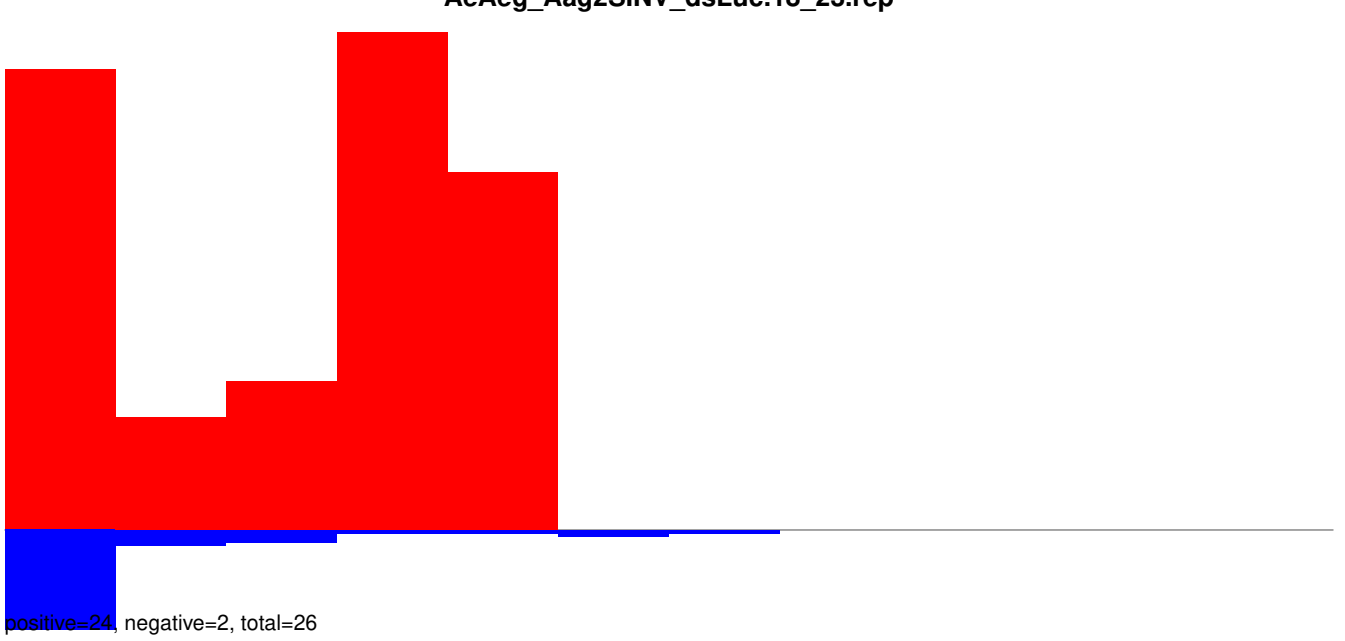


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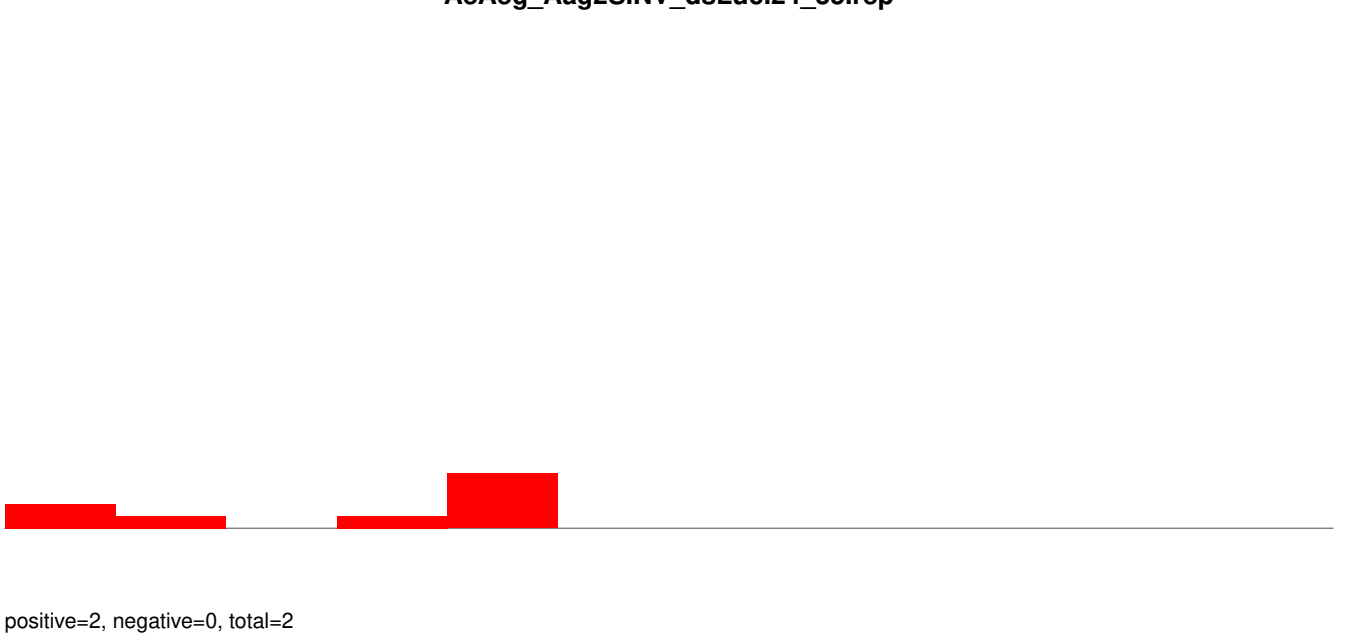


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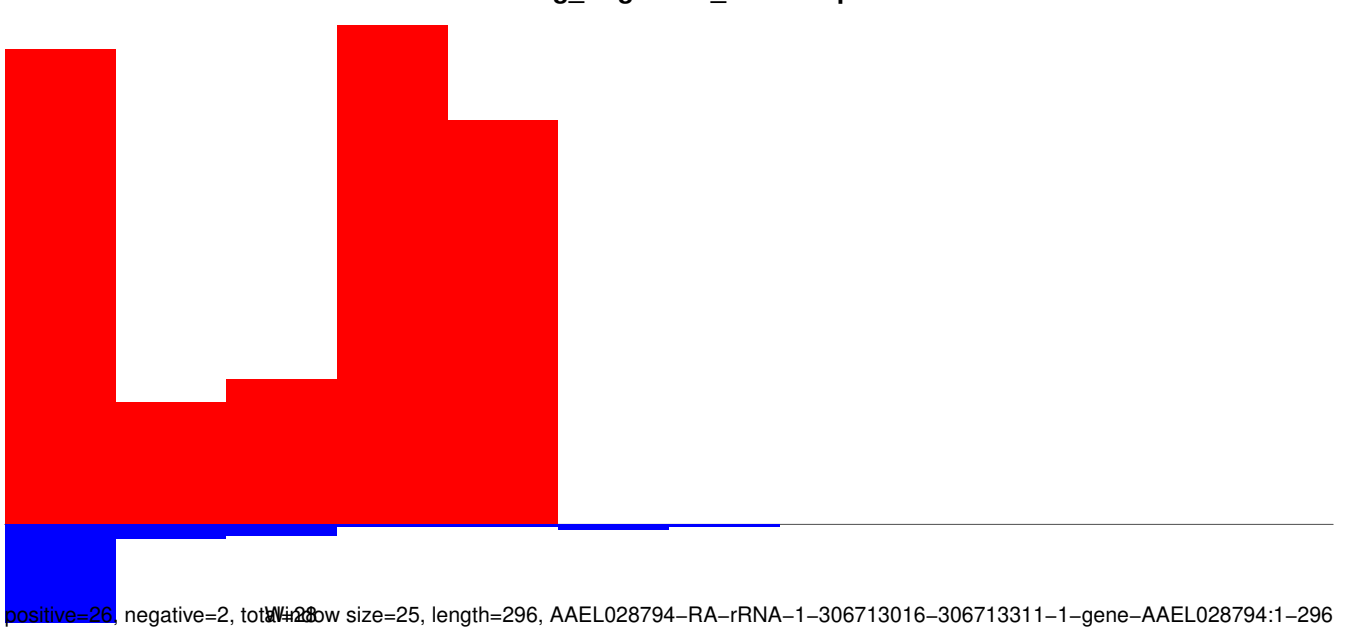
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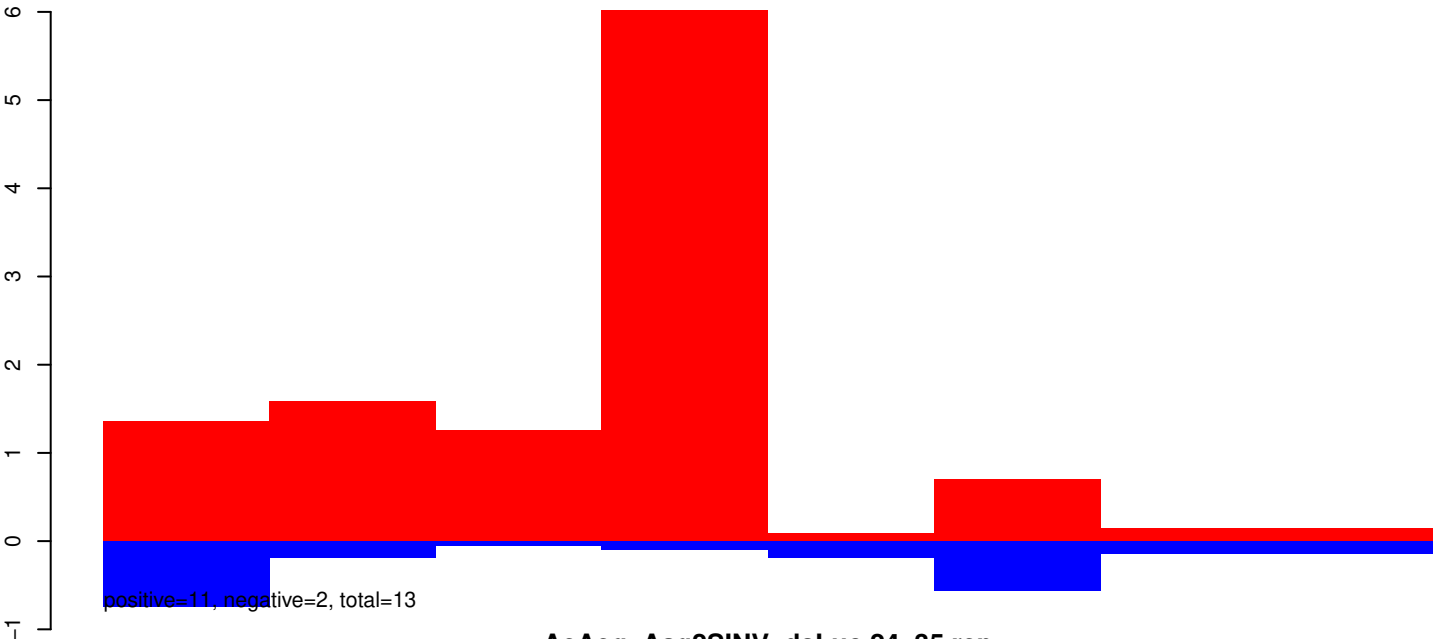
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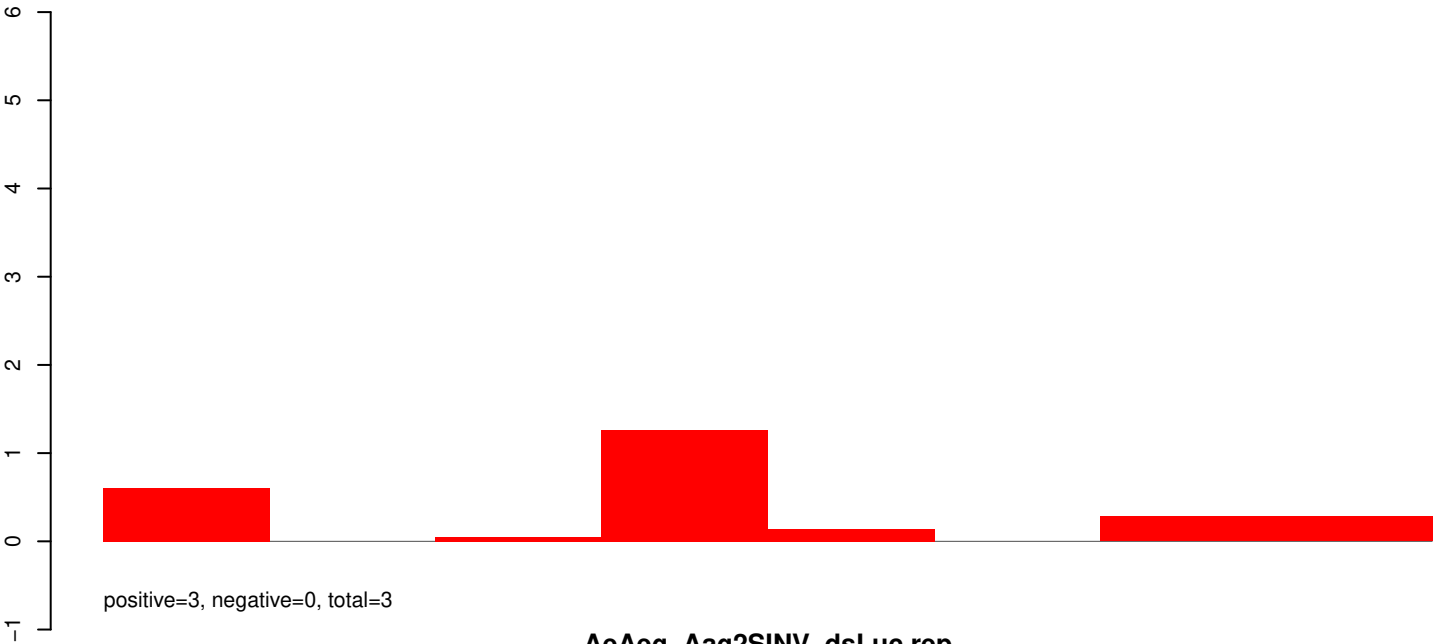
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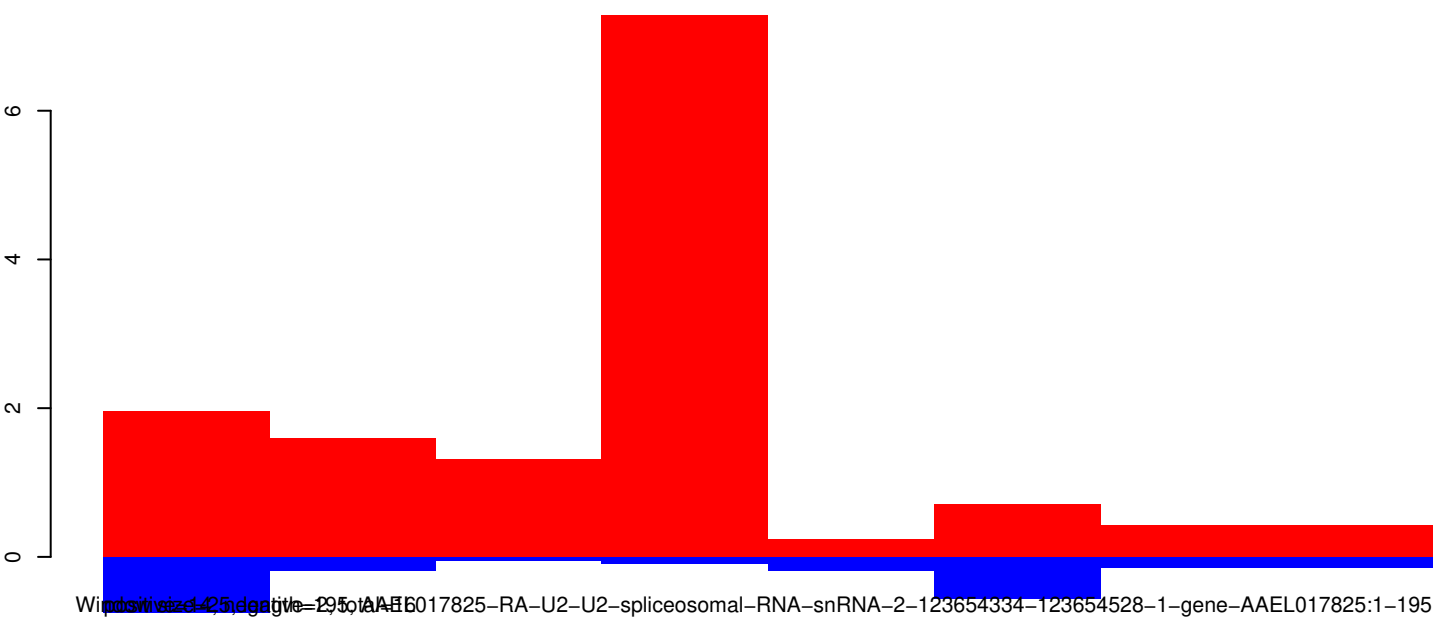
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AeAeg_Aag2SINV_dsLuc.24_35.rep

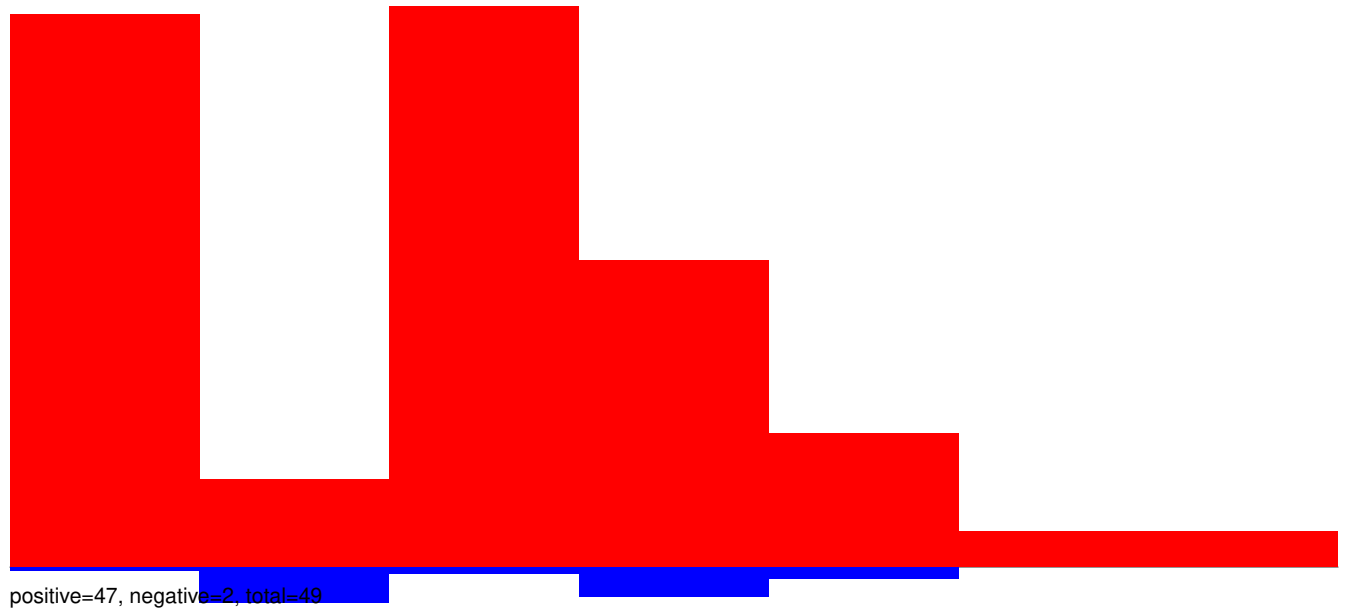


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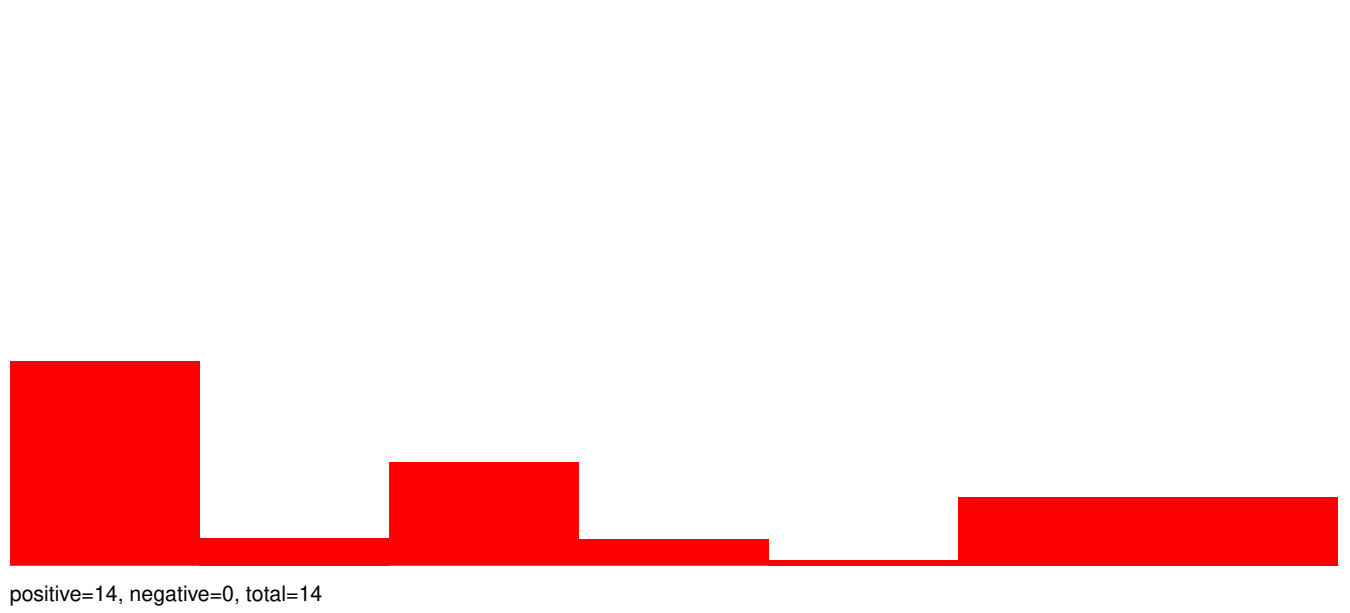


Window size=25, gene=AAE017825-RA-U2-U2-spliceosomal-RNA-snRNA-2-123654334-123654528-1-gene-AAEL017825:1-195

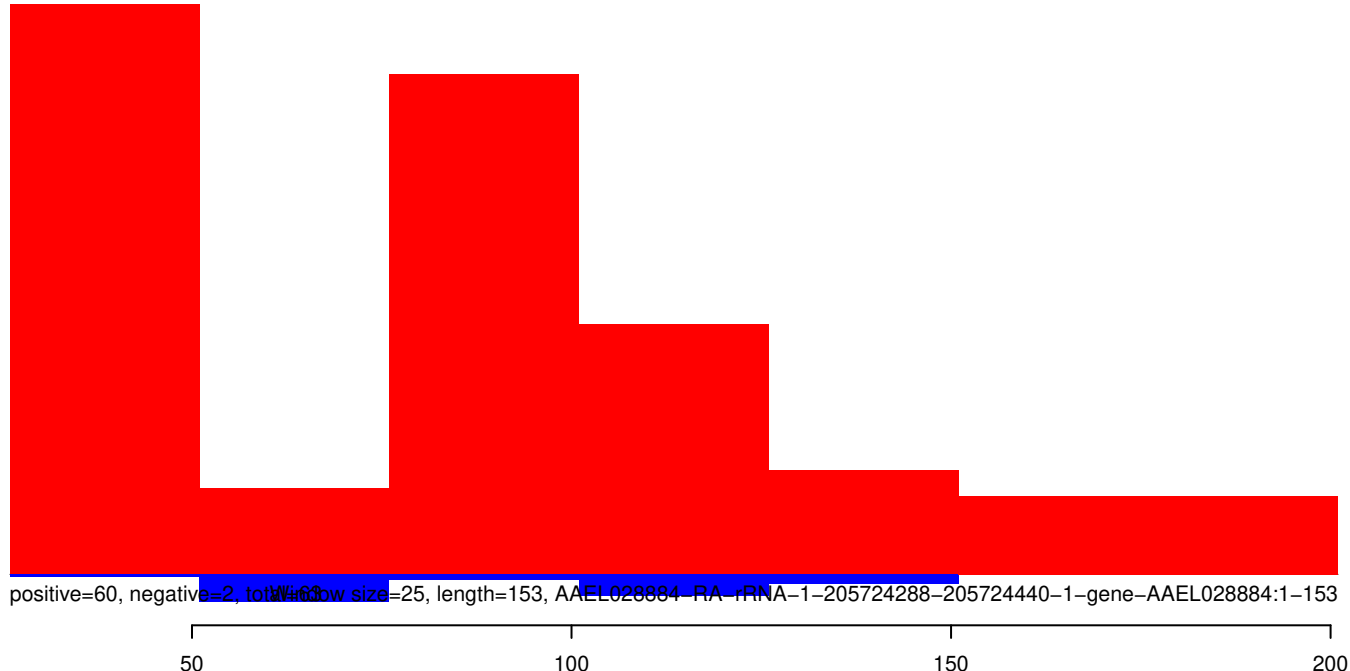
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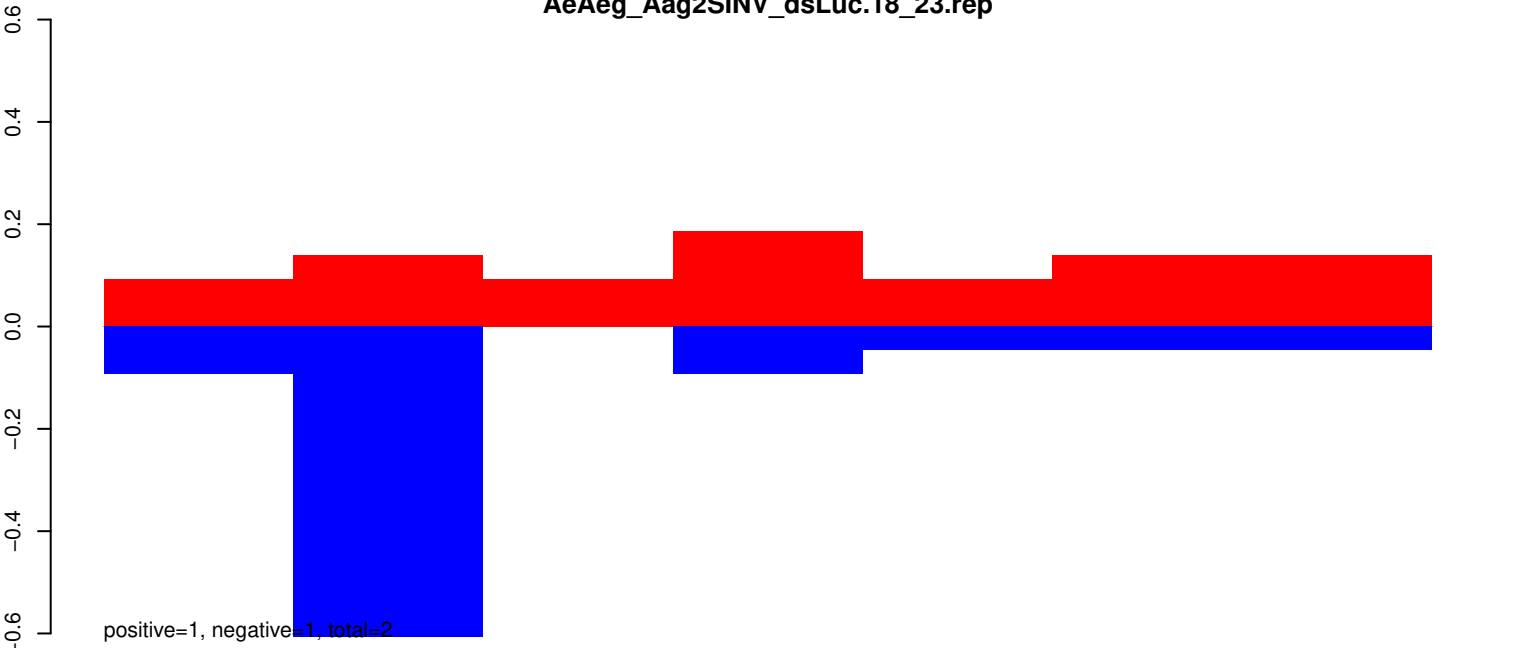
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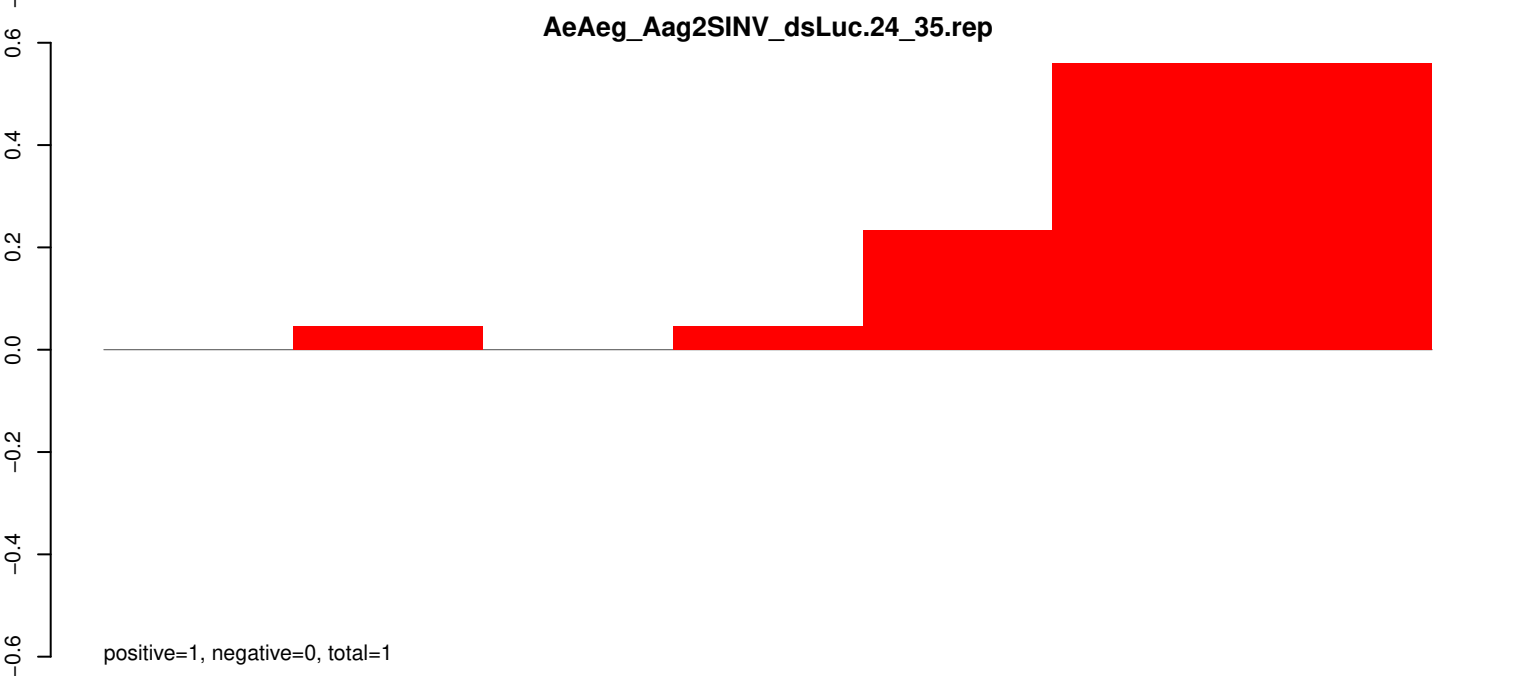
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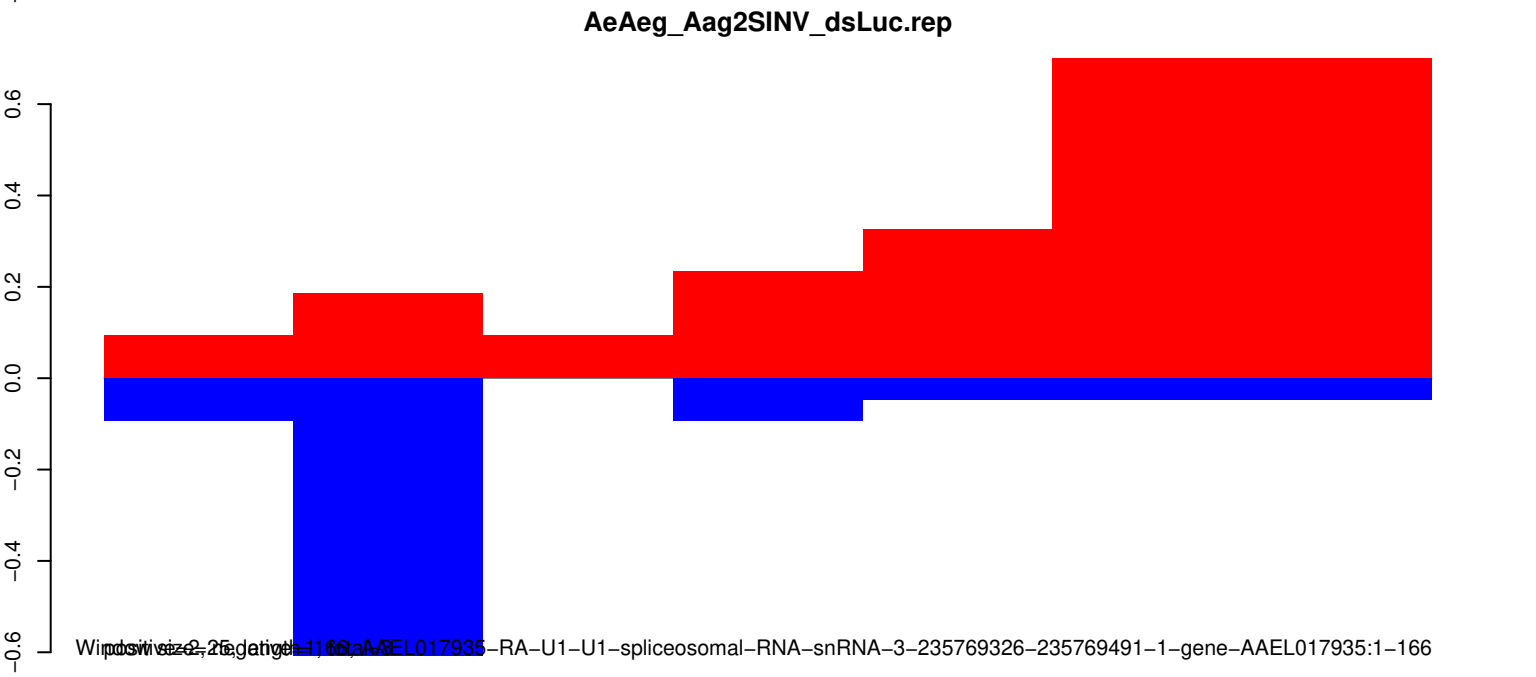
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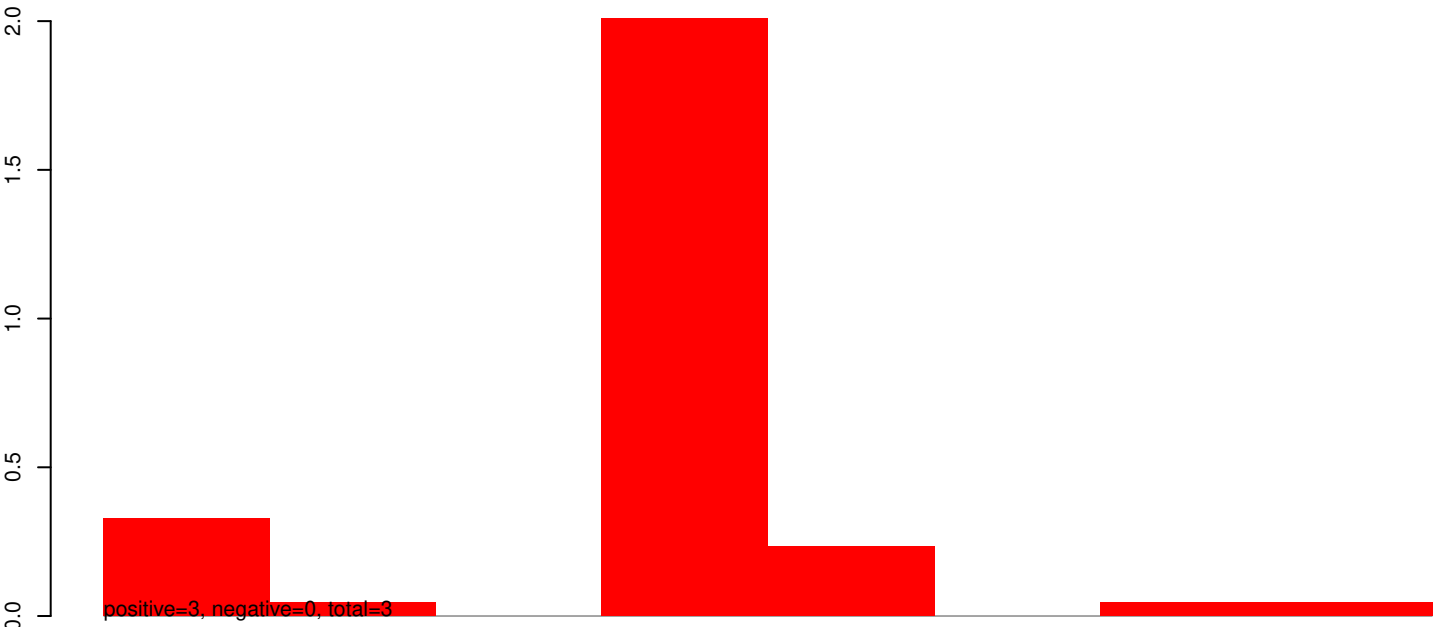
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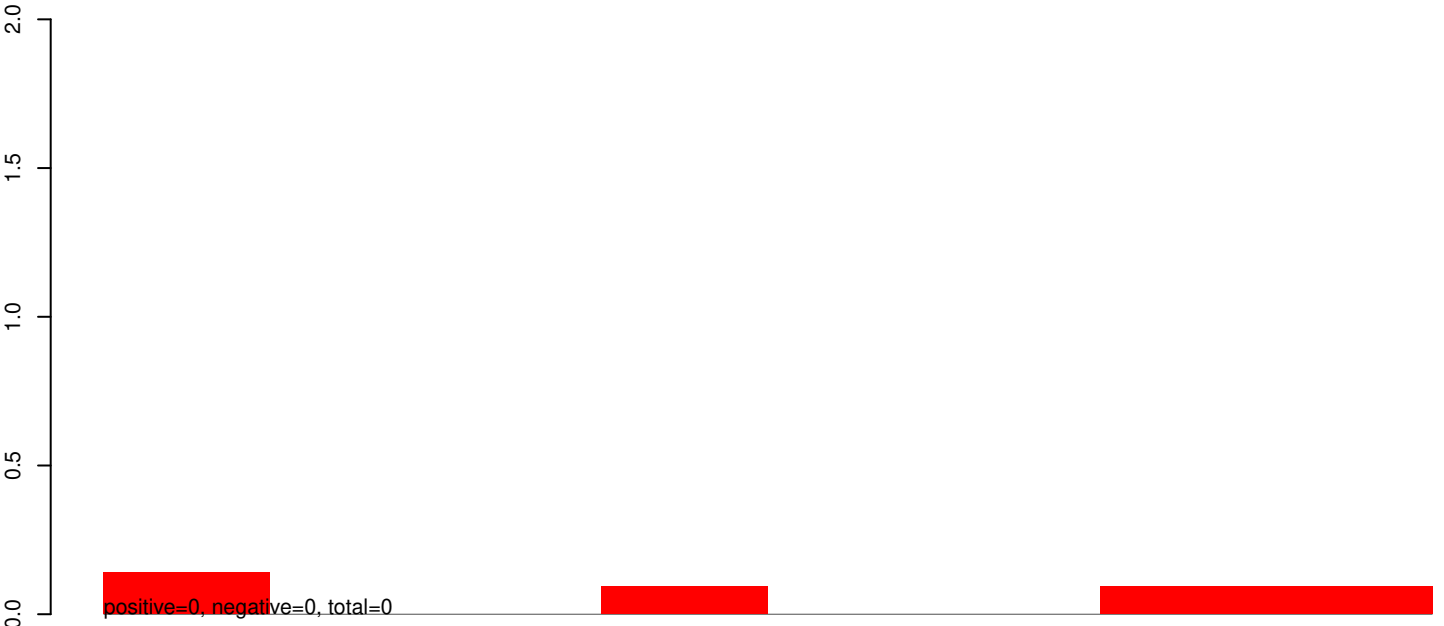
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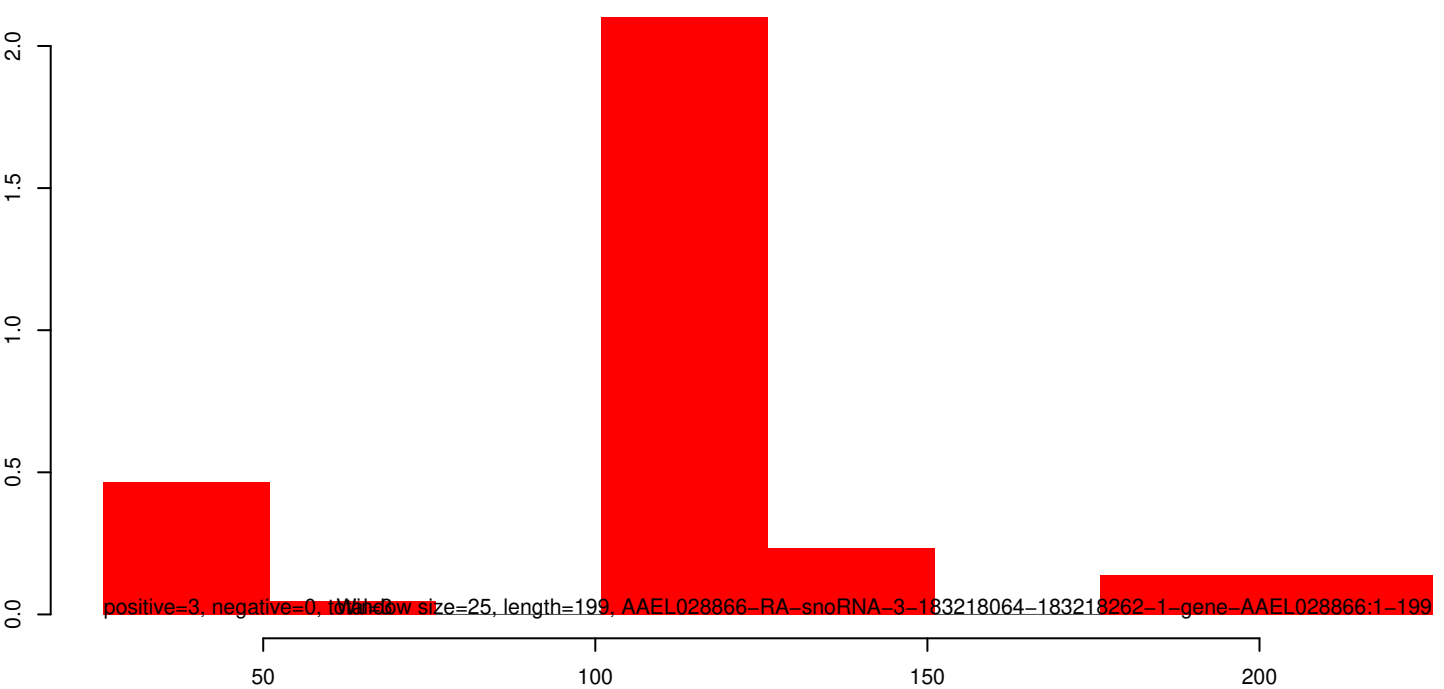
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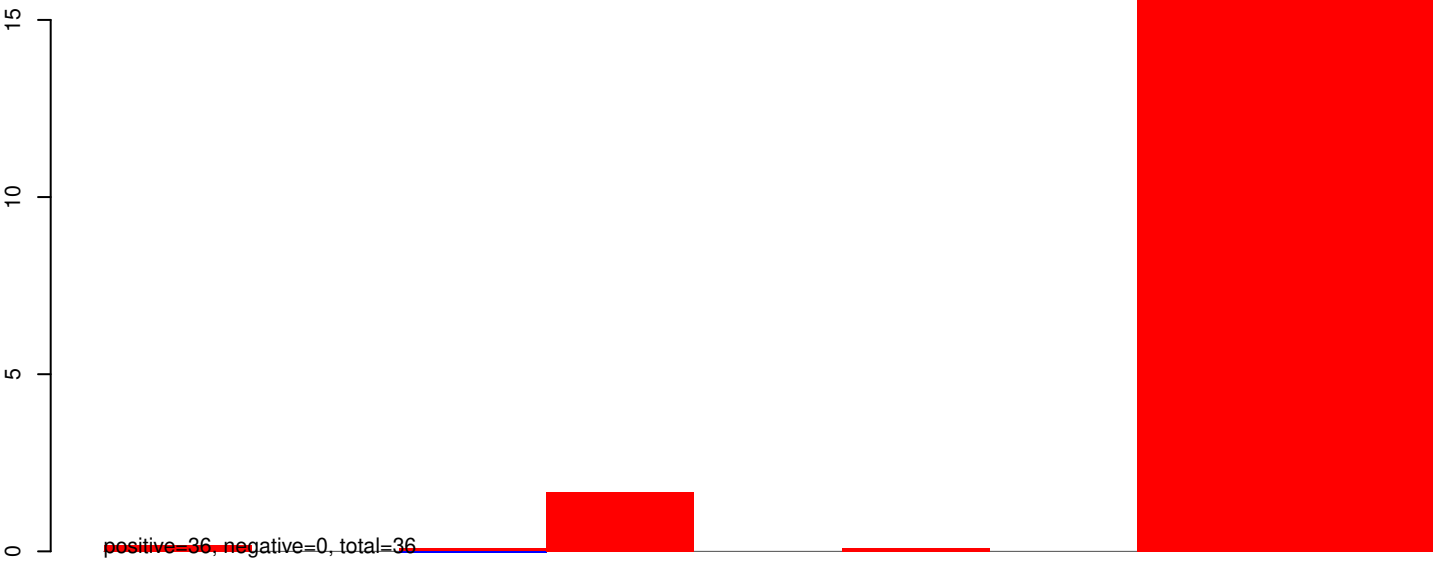
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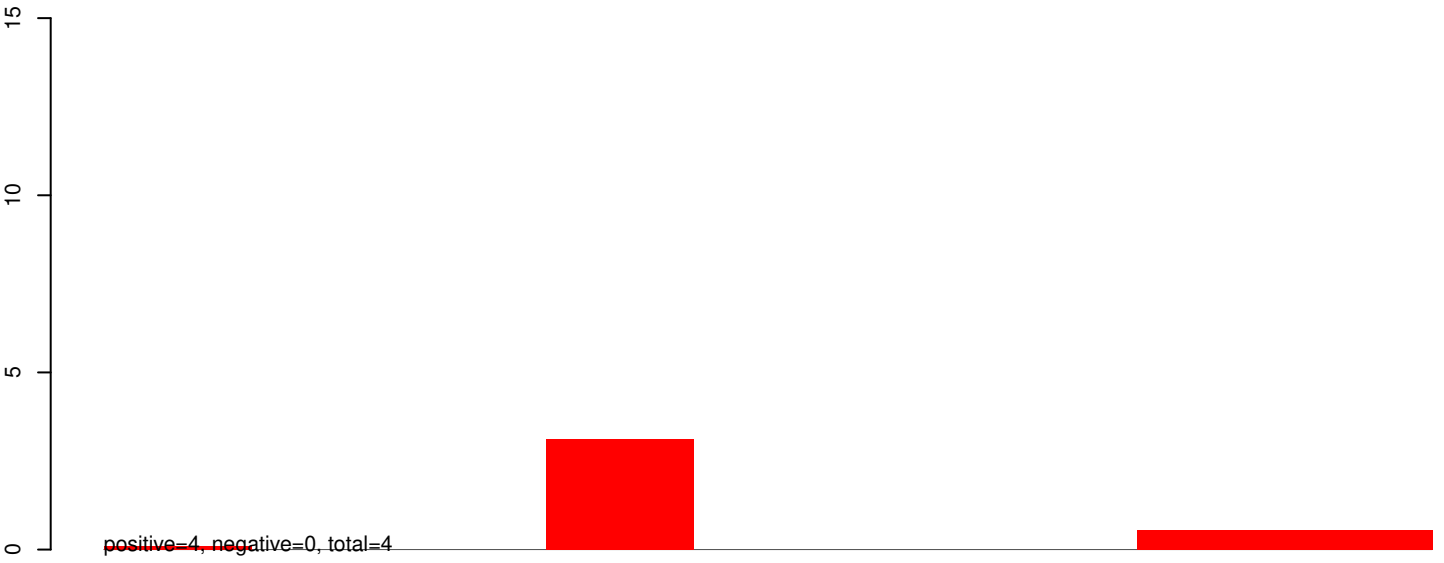
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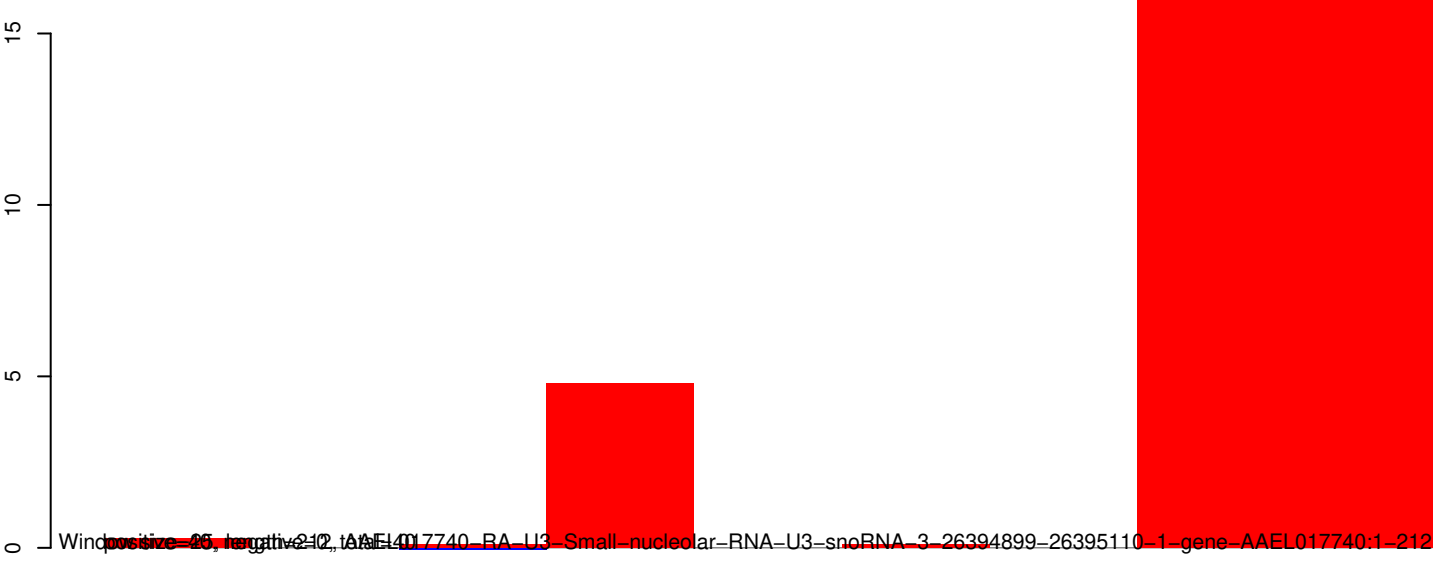
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AeAeg_Aag2SINV_dsLuc.24_35.rep

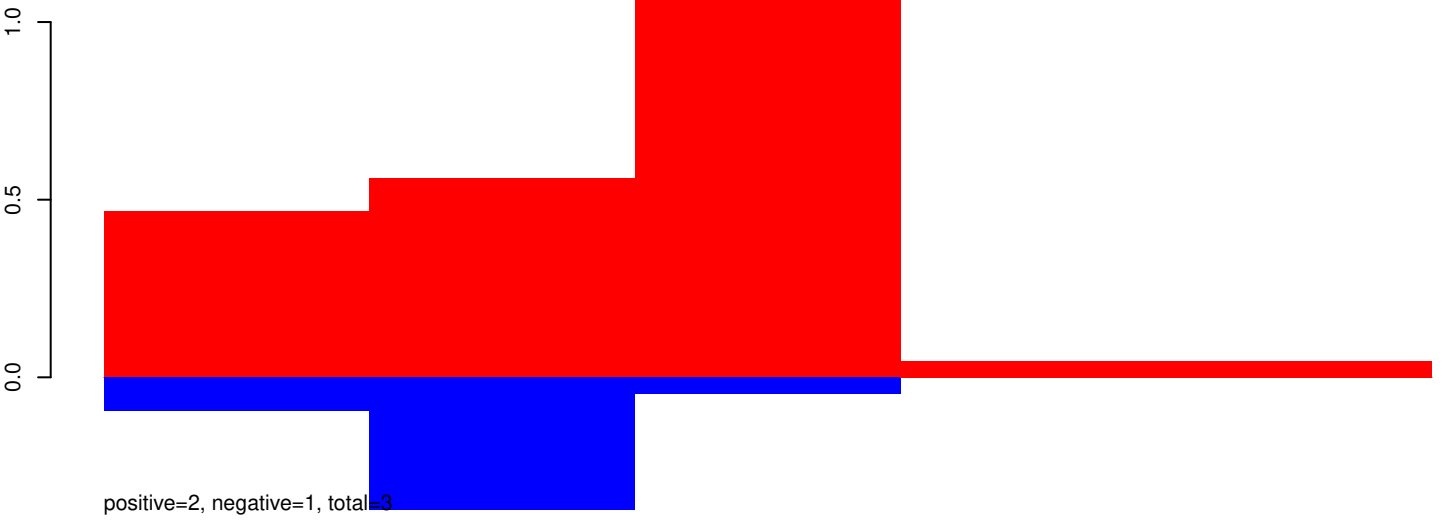


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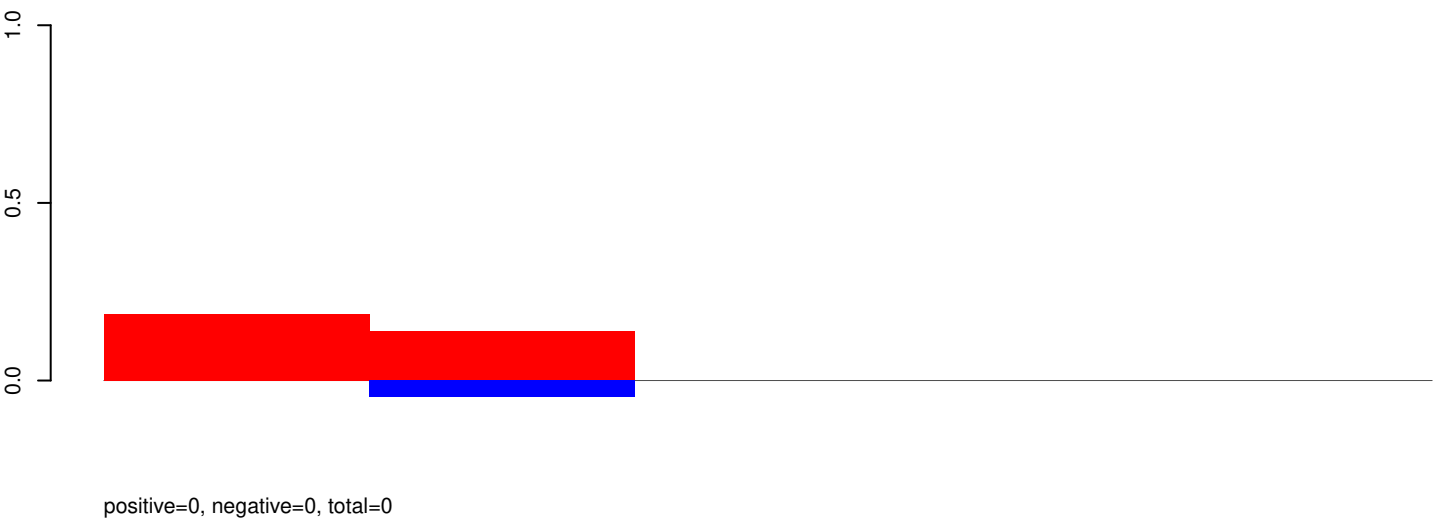


Windos positive=25, negative=10, total=35, gene=AAEL017740-RA-U3-Small-nucleolar-RNA-U3-snoRNA-3-26394899-26395110-1-gene-AAEL017740:1-212

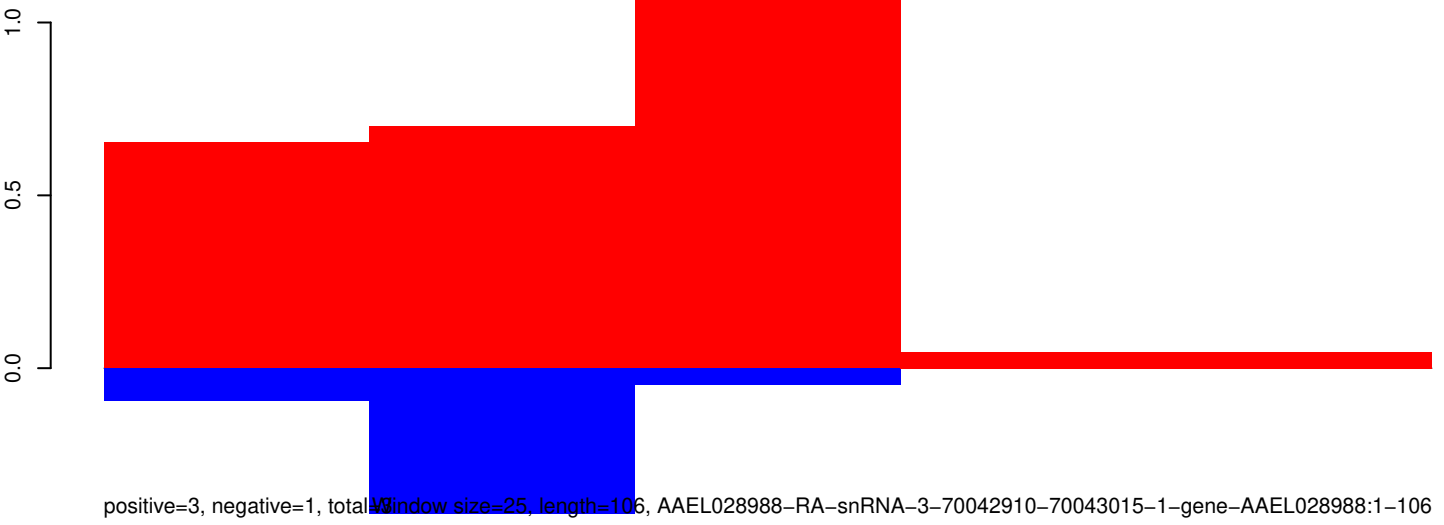
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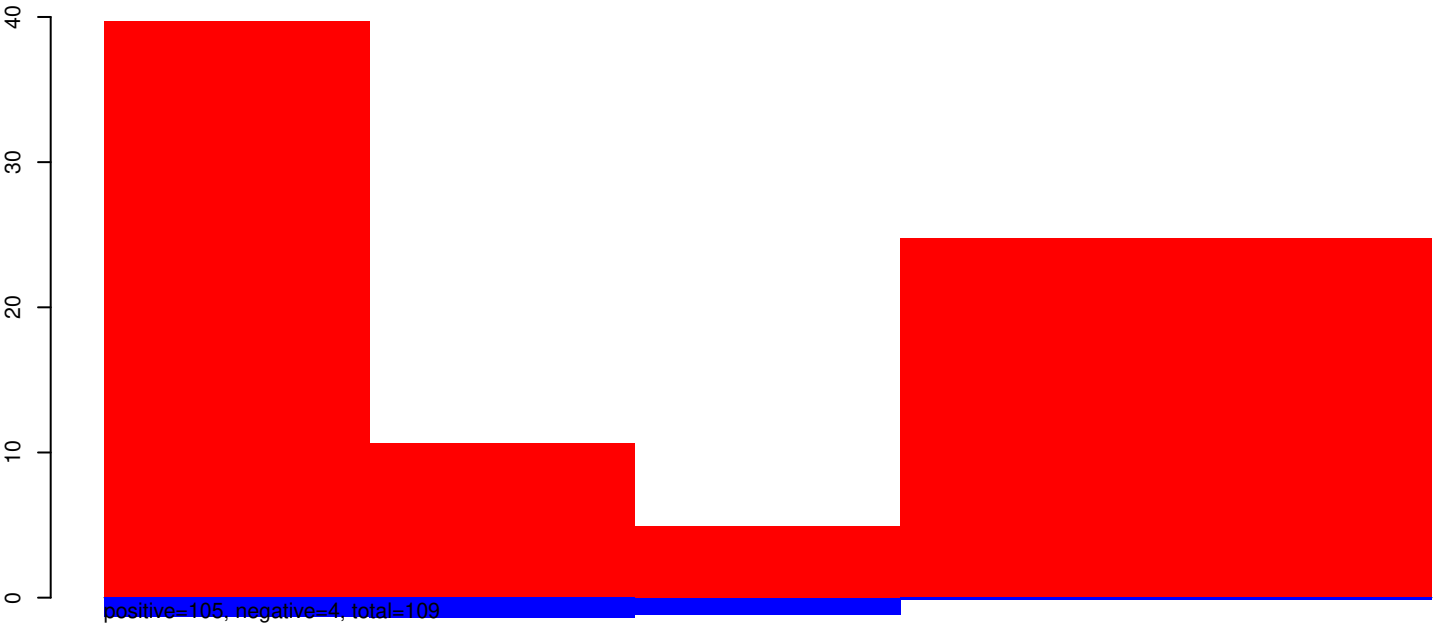
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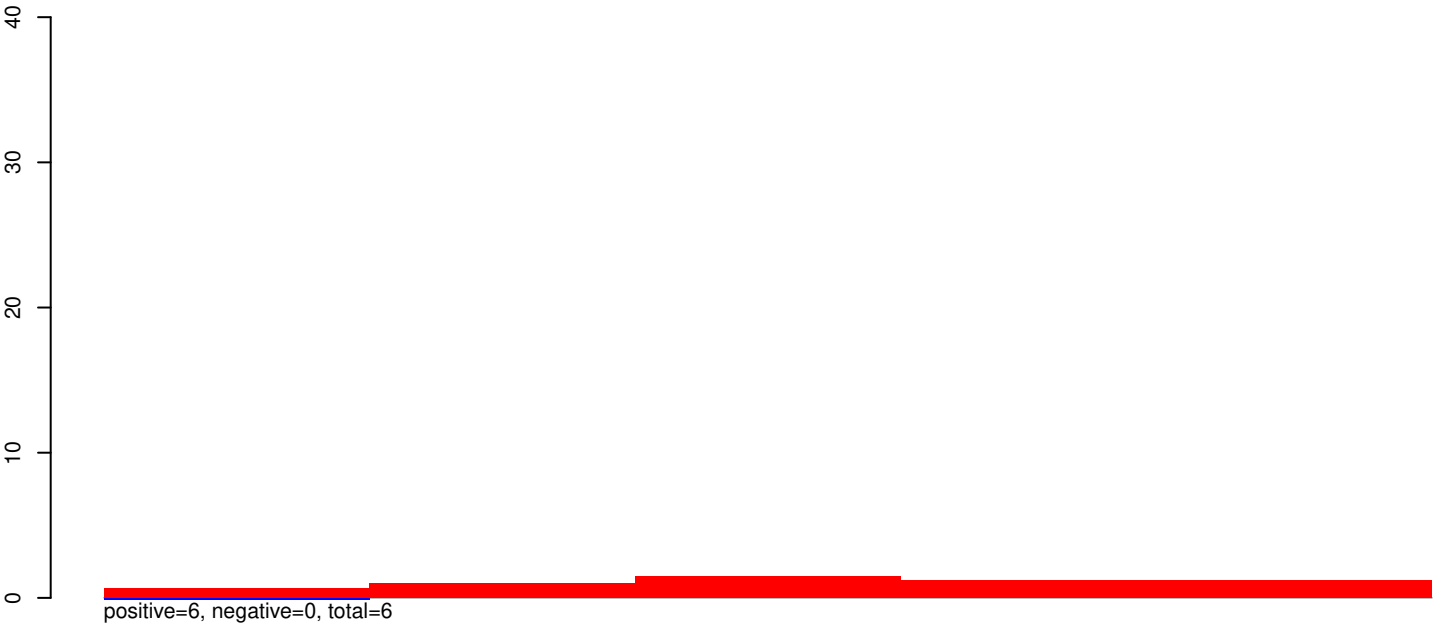
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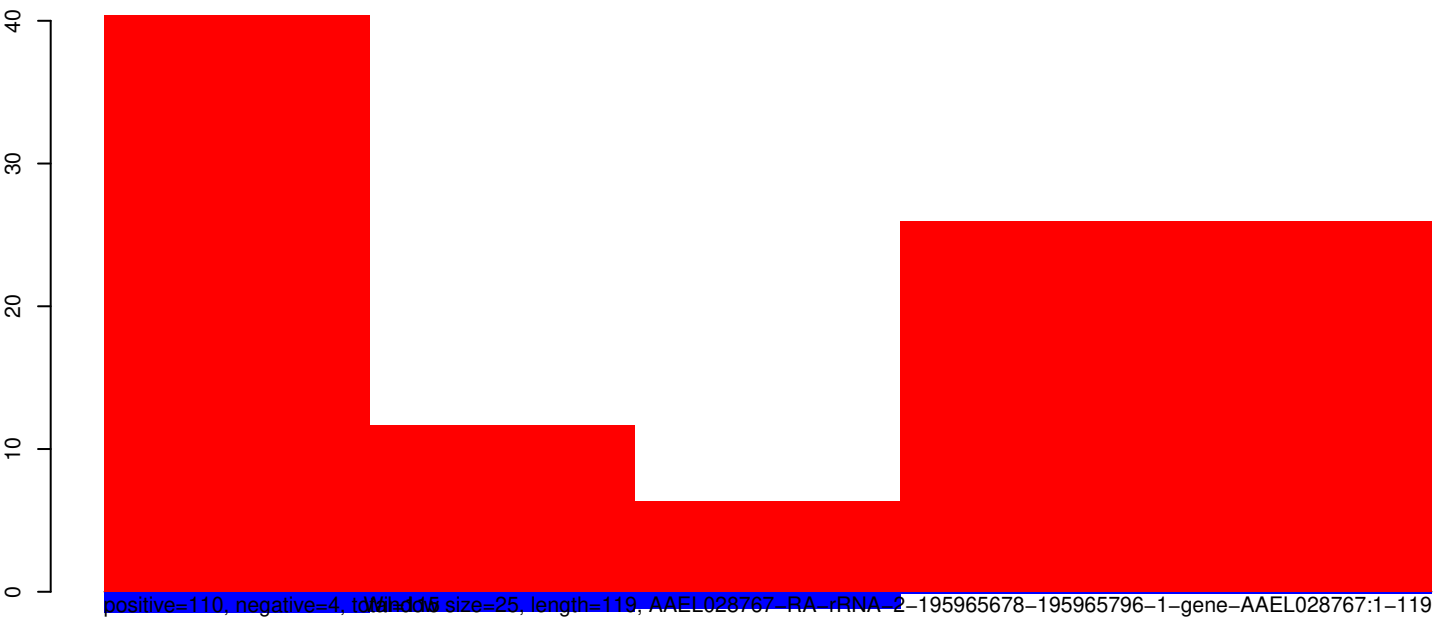
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AeAeg_Aag2SINV_dsLuc.24_35.rep



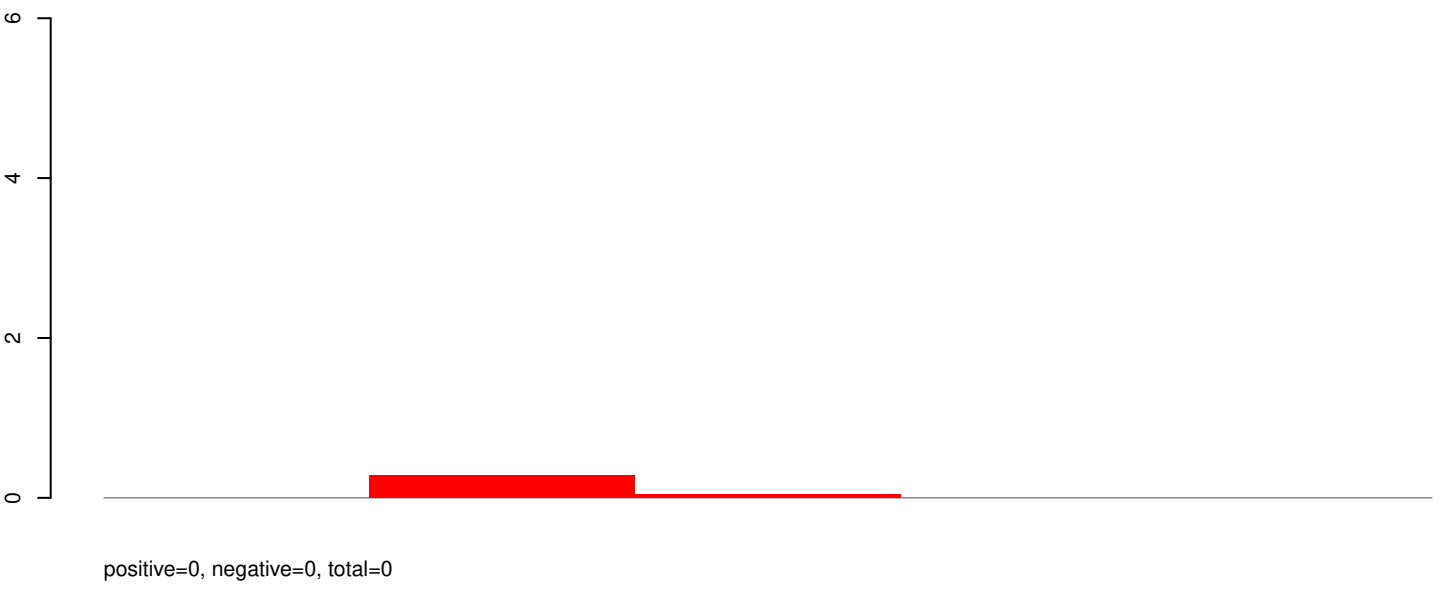
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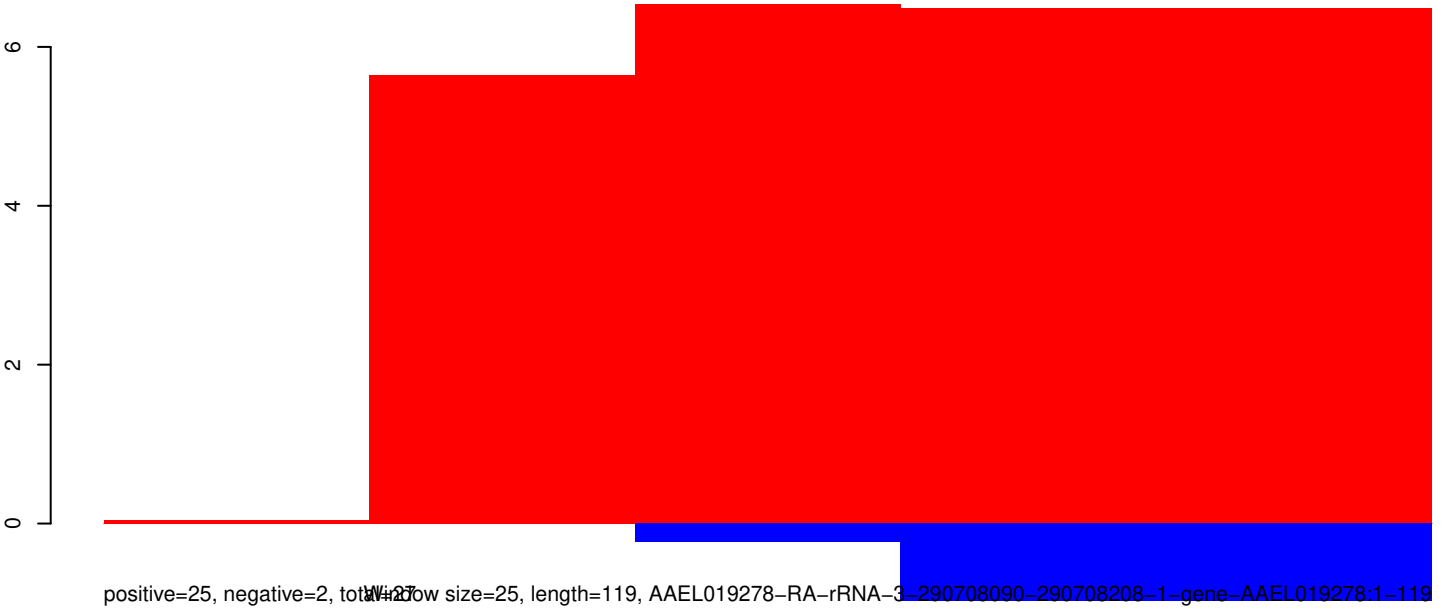
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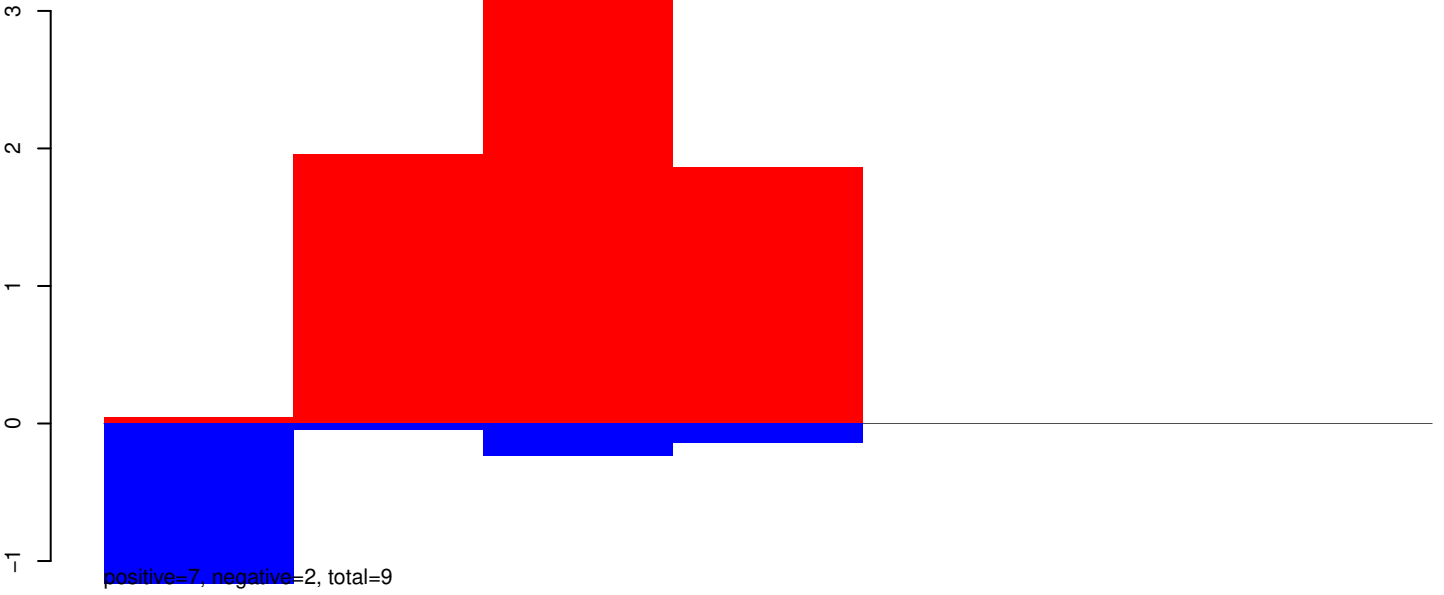
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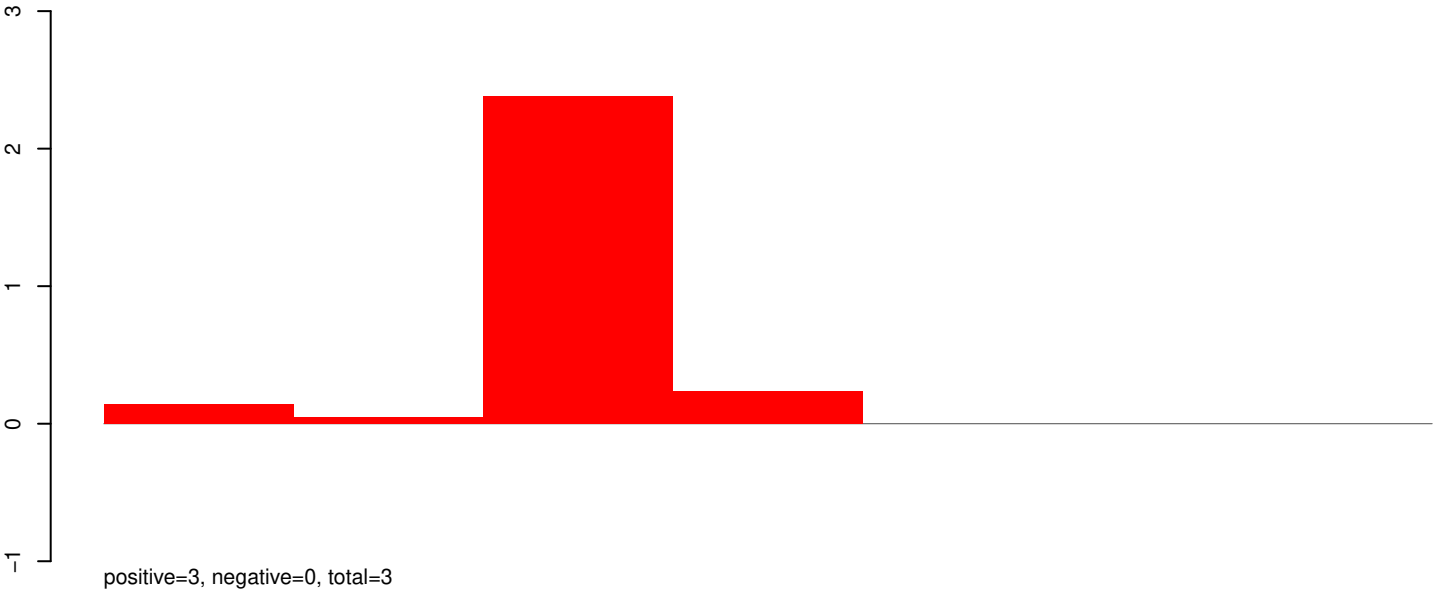
40 60 80 100 120 140

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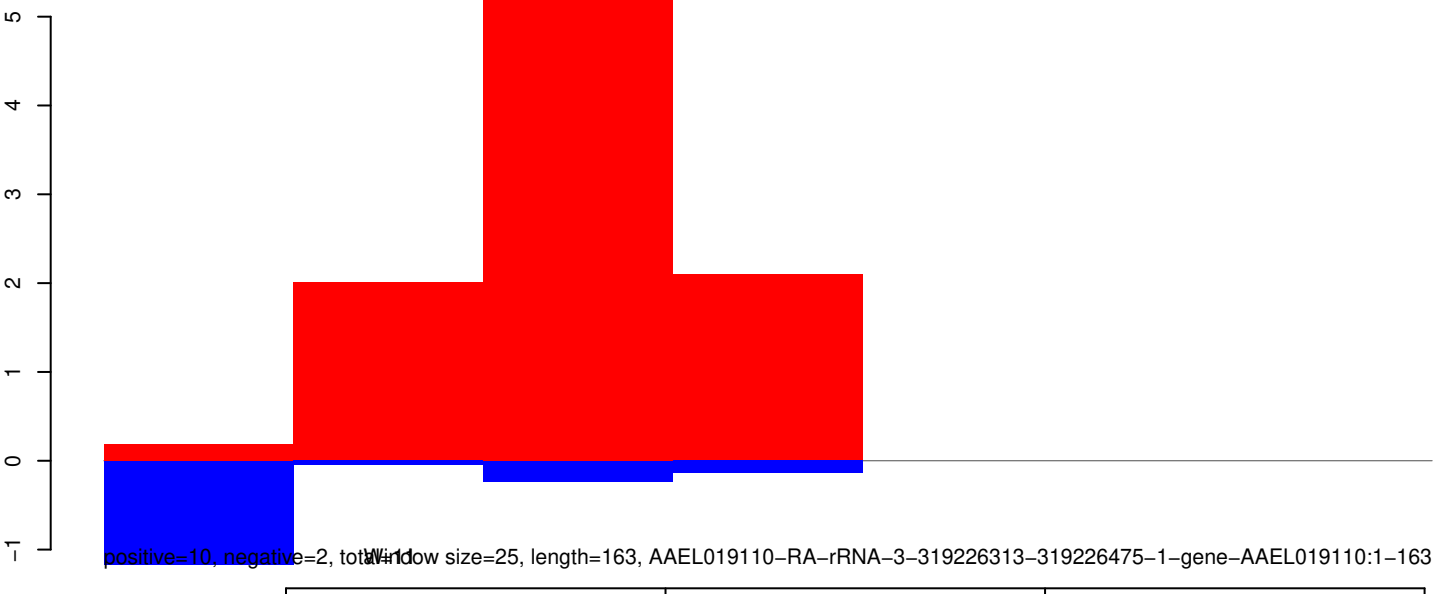
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AeAeg_Aag2SINV_dsLuc.24_35.rep

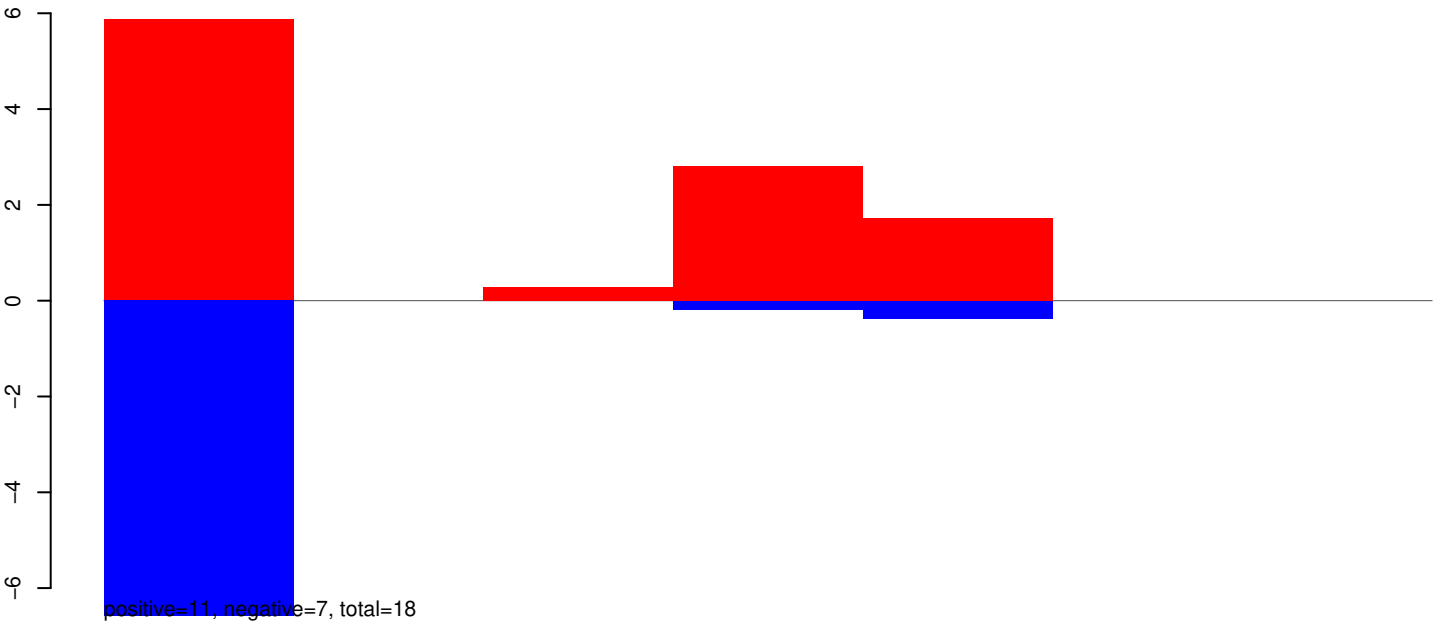


AeAeg_Aag2SINV_dsLuc.rep



Window size=25, length=163, AAEL019110-RA-rRNA-3-319226313-319226475-1-gene-AAEL019110:1-163

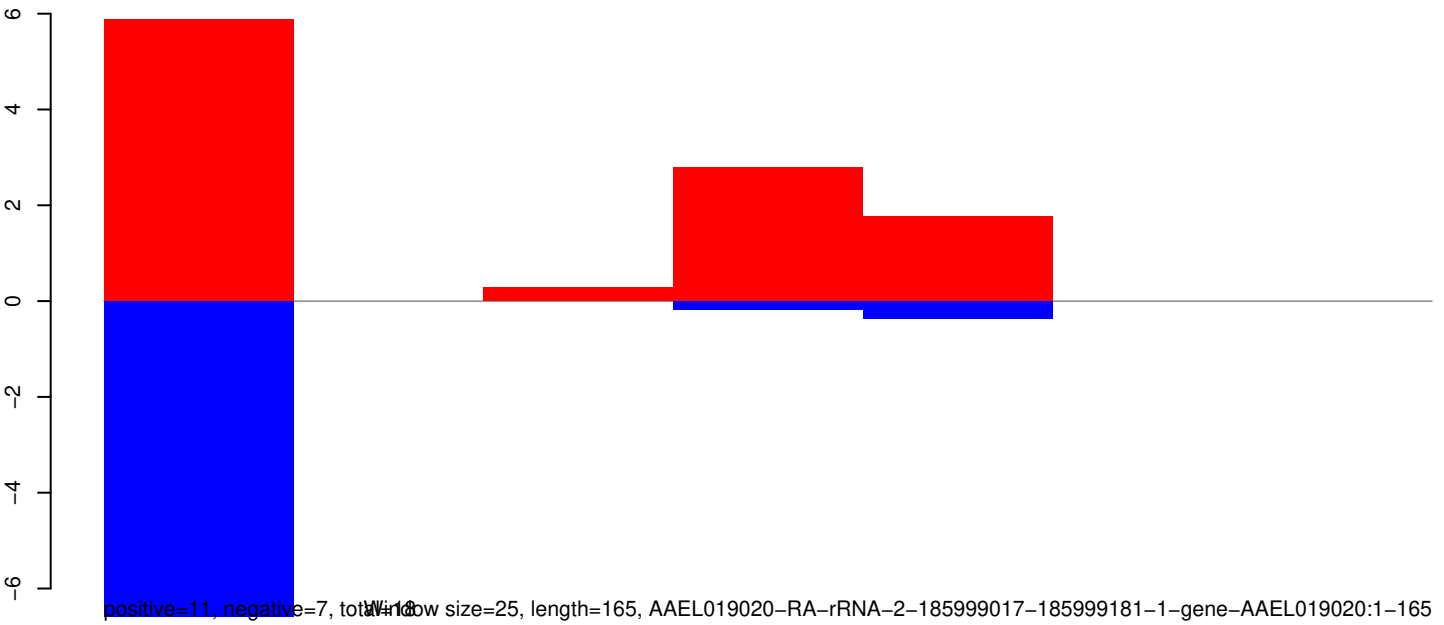
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AeAeg_Aag2SINV_dsLuc.24_35.rep

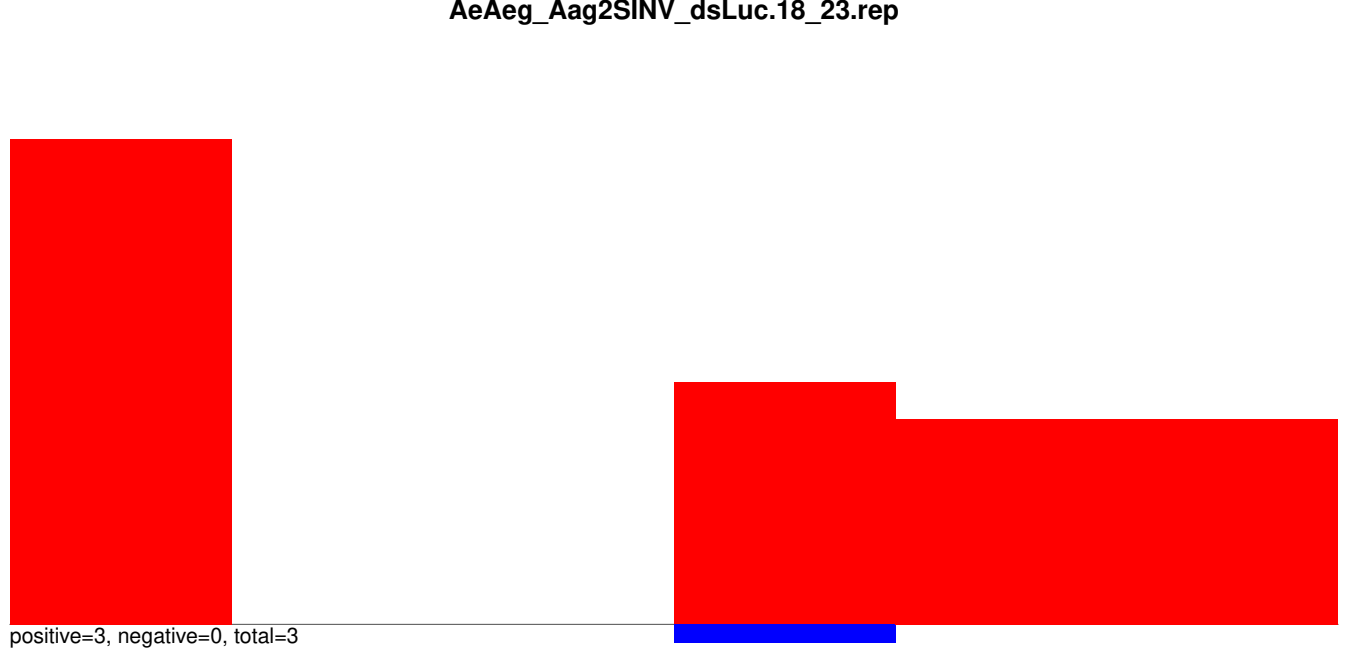


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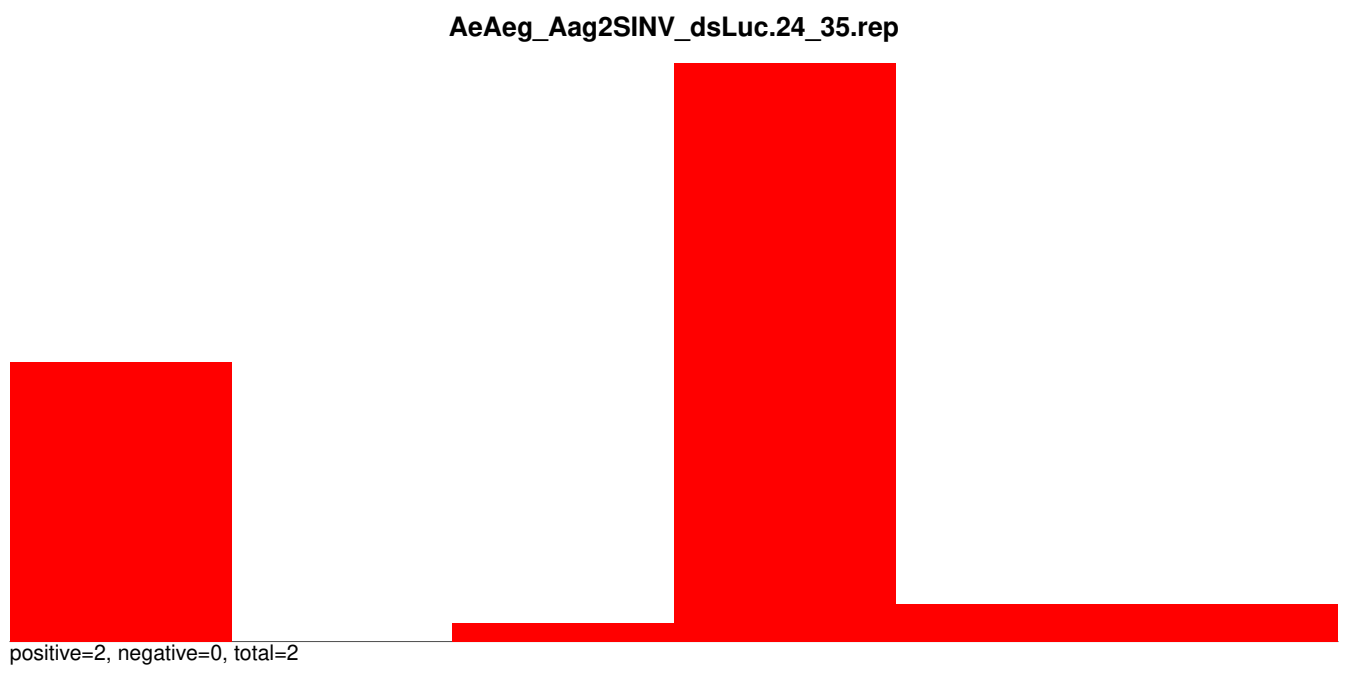


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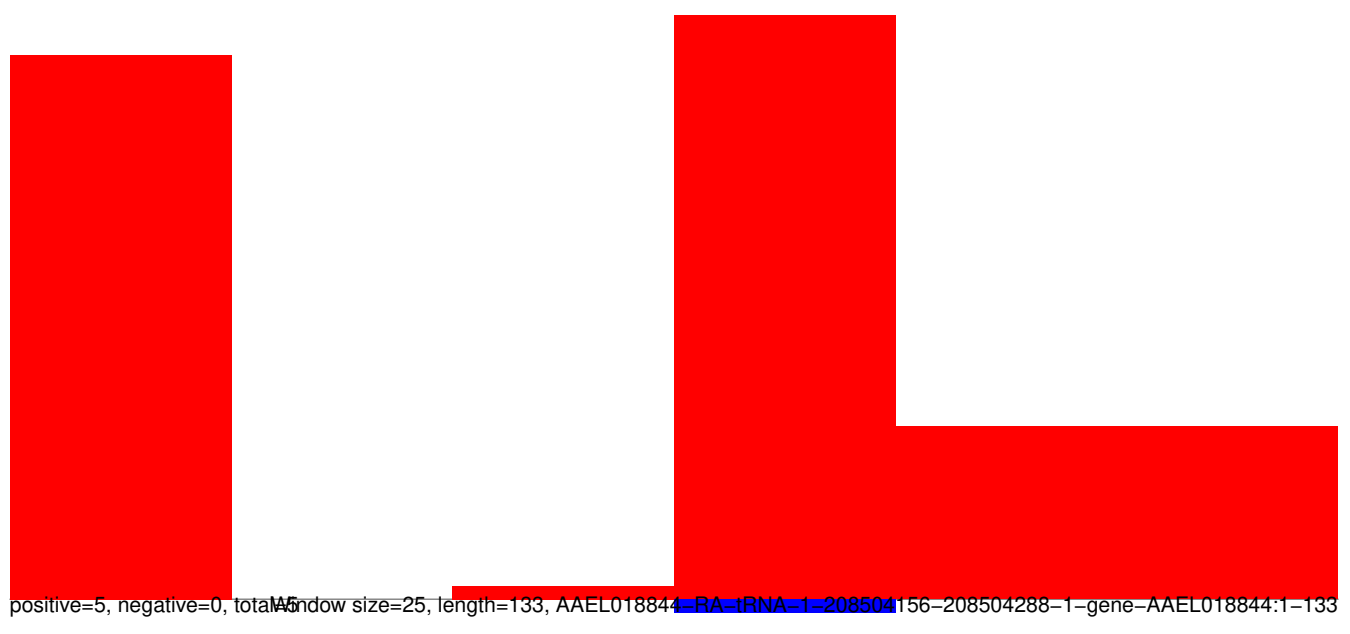
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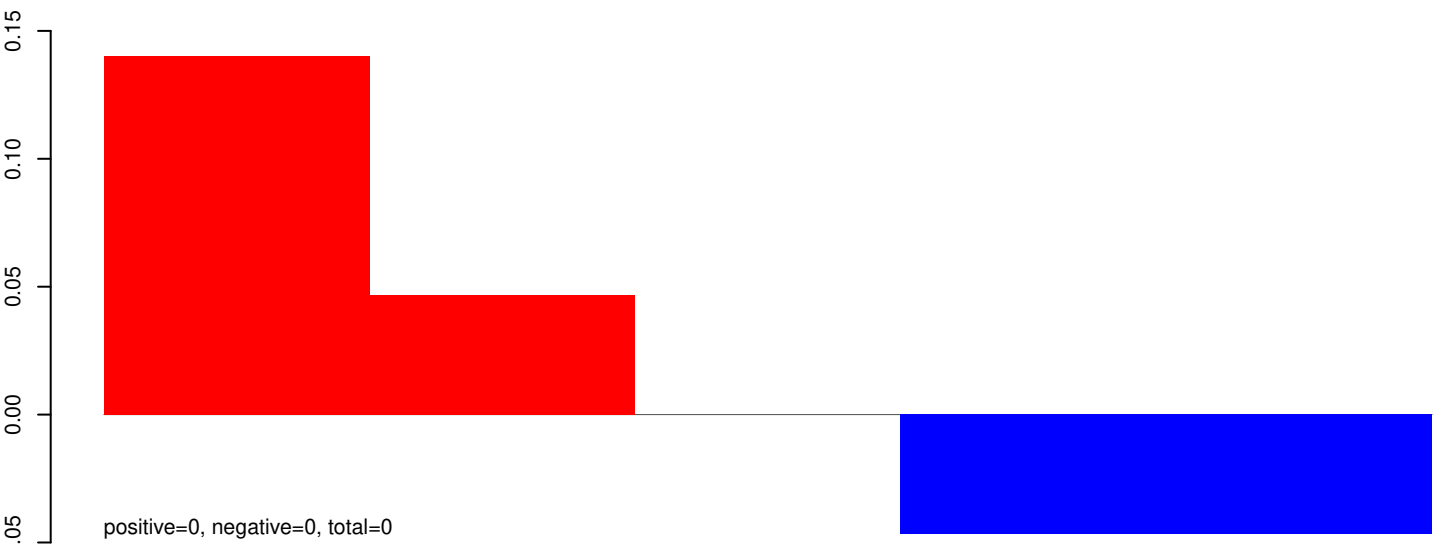
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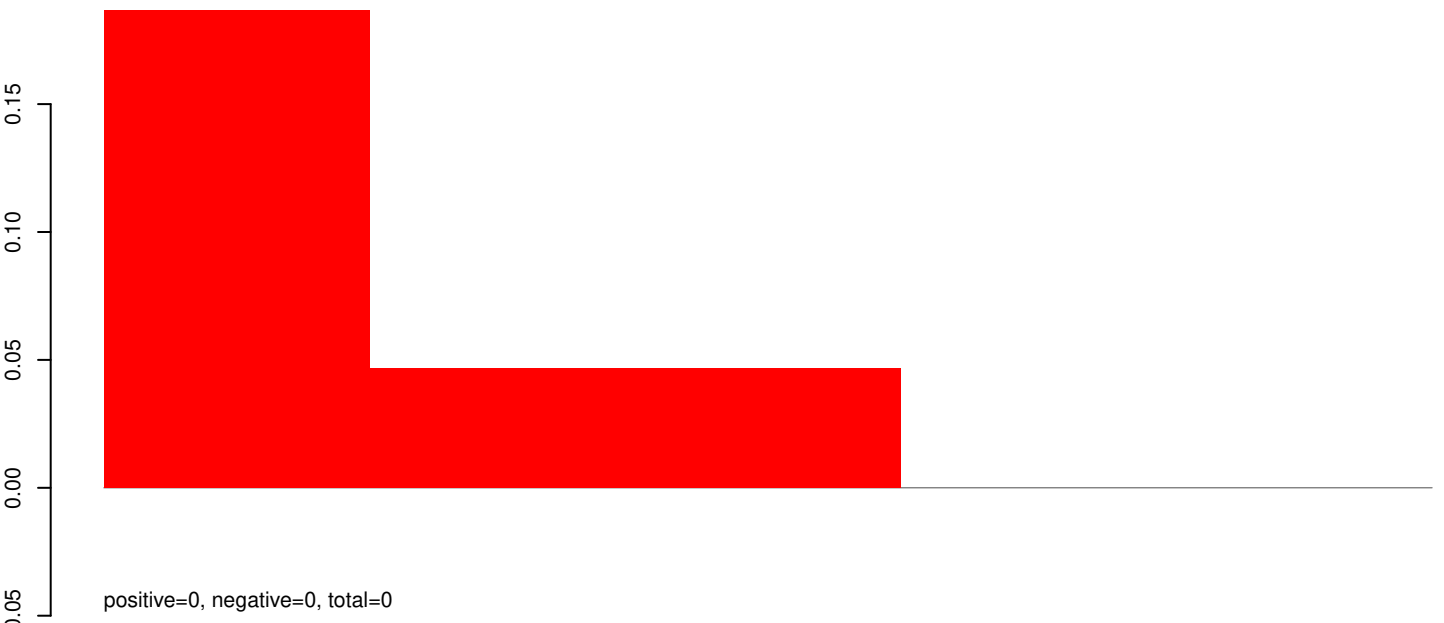
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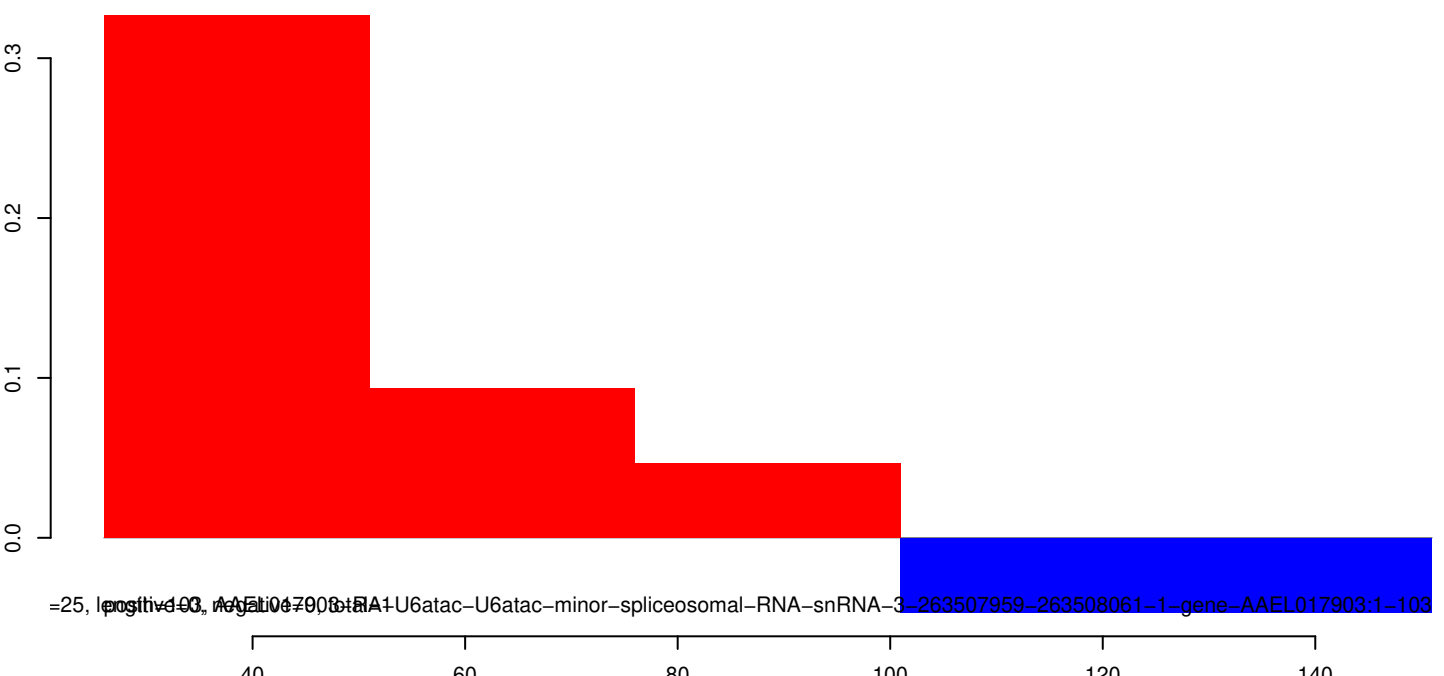
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



=25, length=103, ref=1047906+AA4U6atac-U6atac-minor-spliceosomal-RNA-snRNA-3-263507959-263508061-1-gene-AAEL017903:1-103

40 60 80 100 120 140

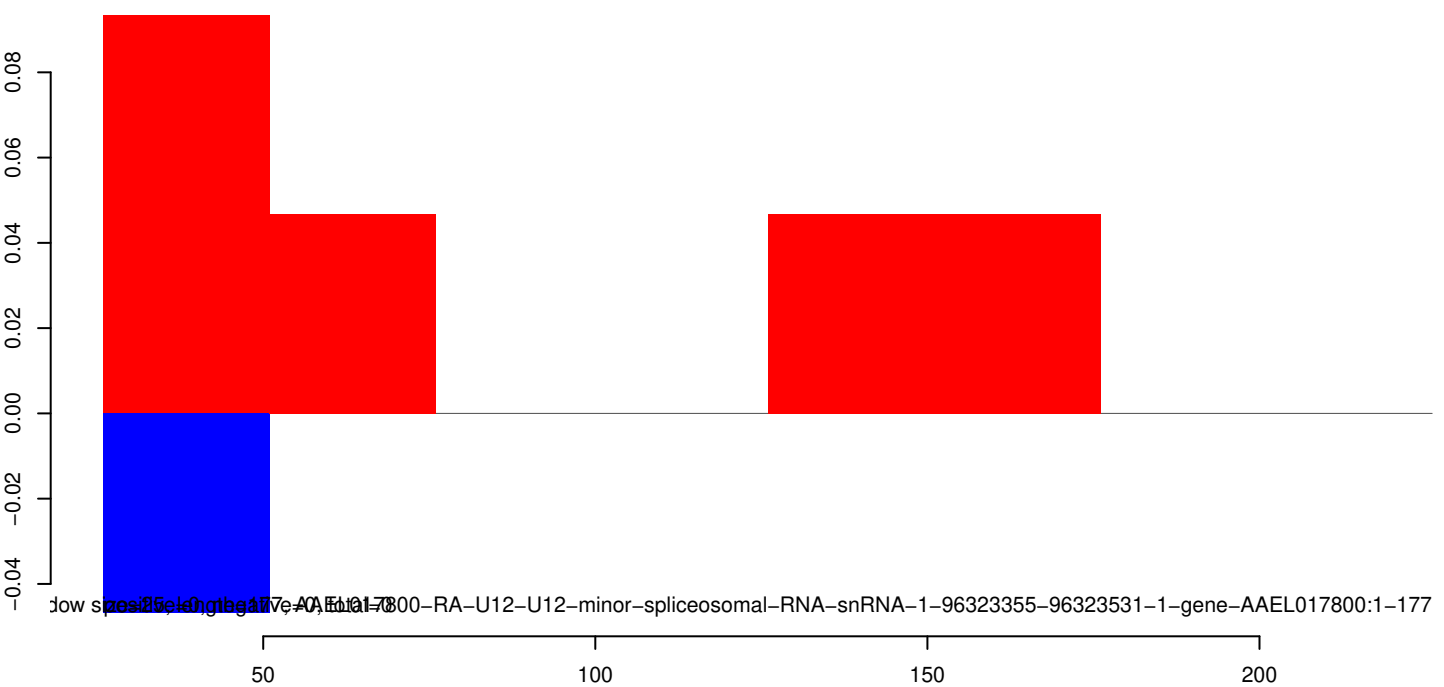
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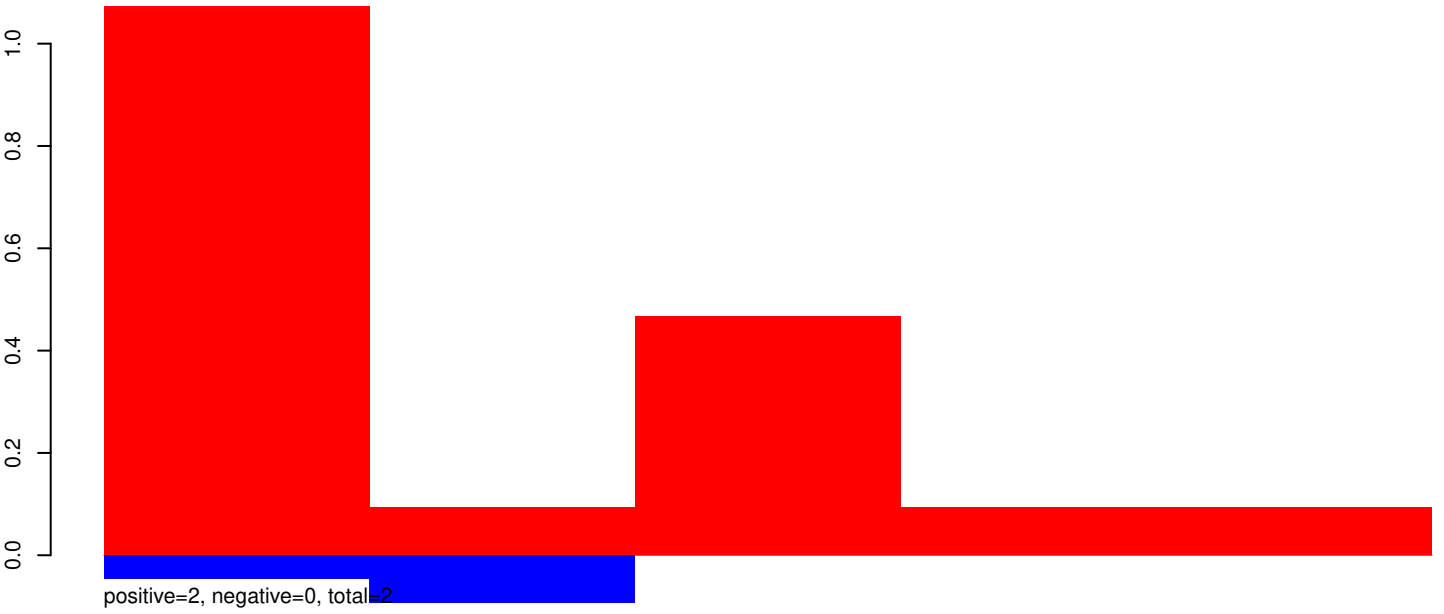


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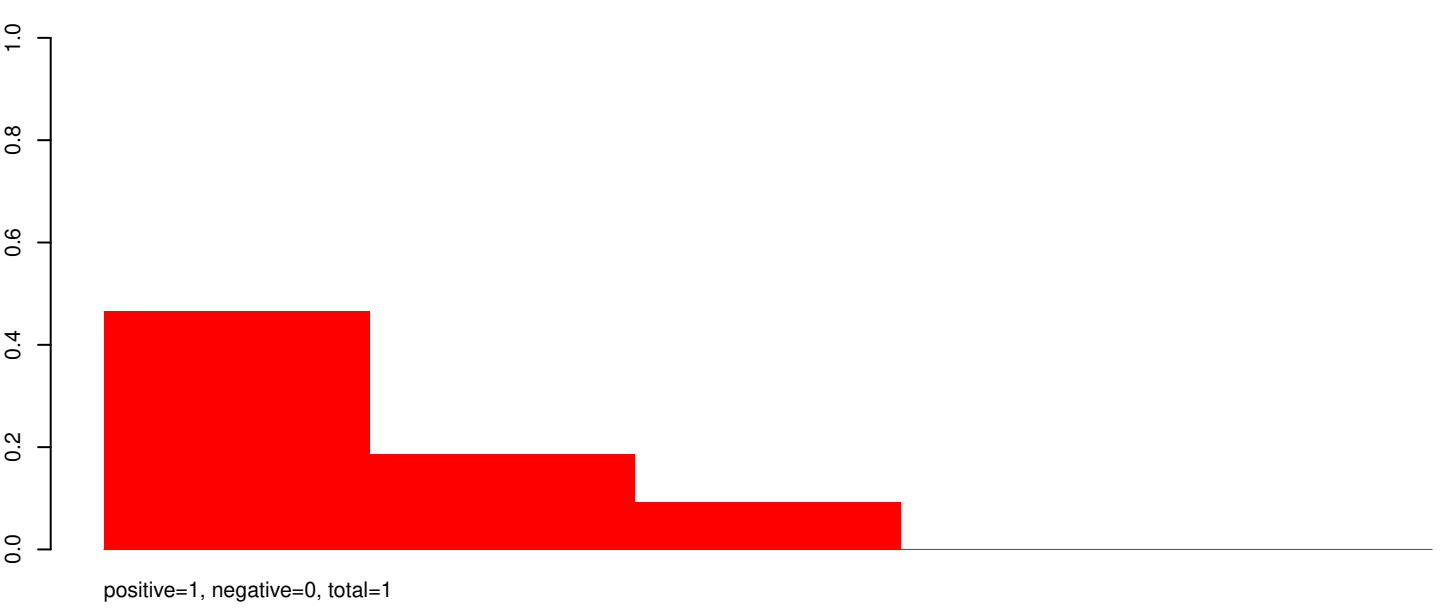


low speed single gene: AEL01780-RA-U12-U12-minor-spliceosomal-RNA-snrRNA-1-96323355-96323531-1-gene-AAEL017800:1-177

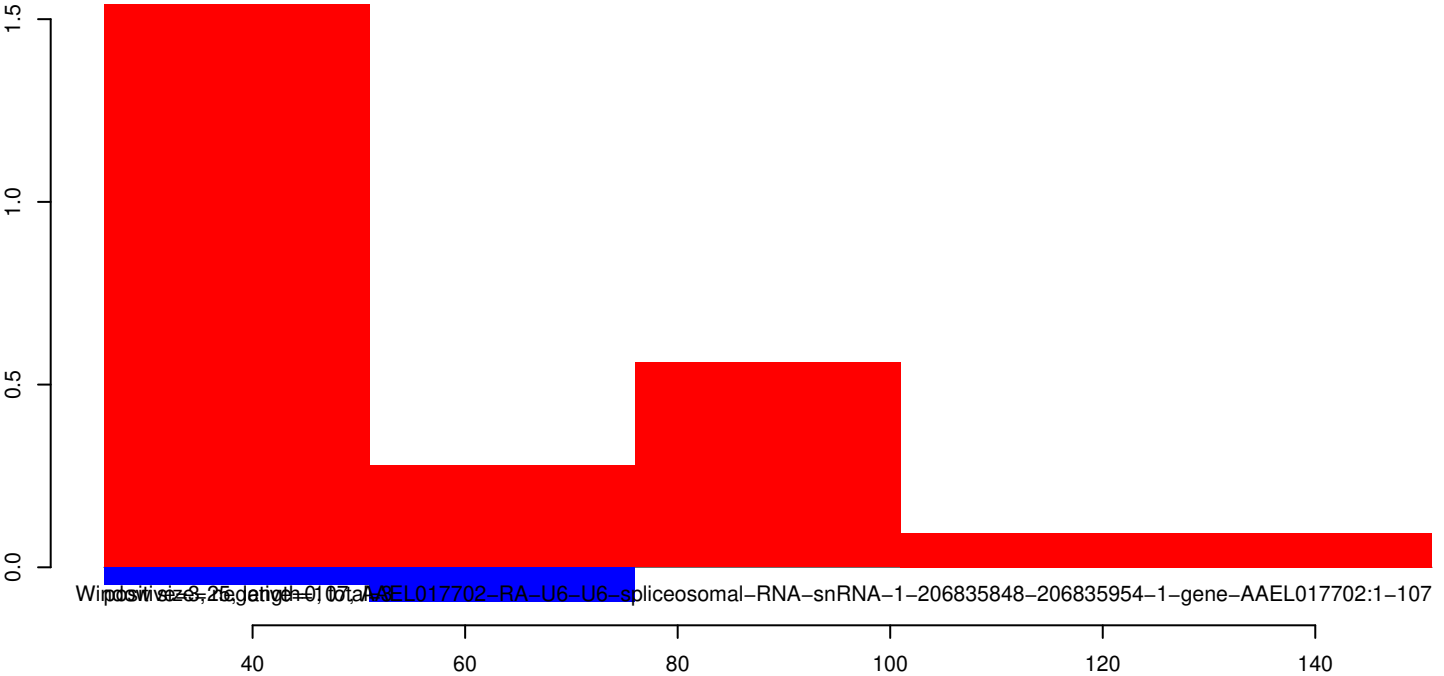
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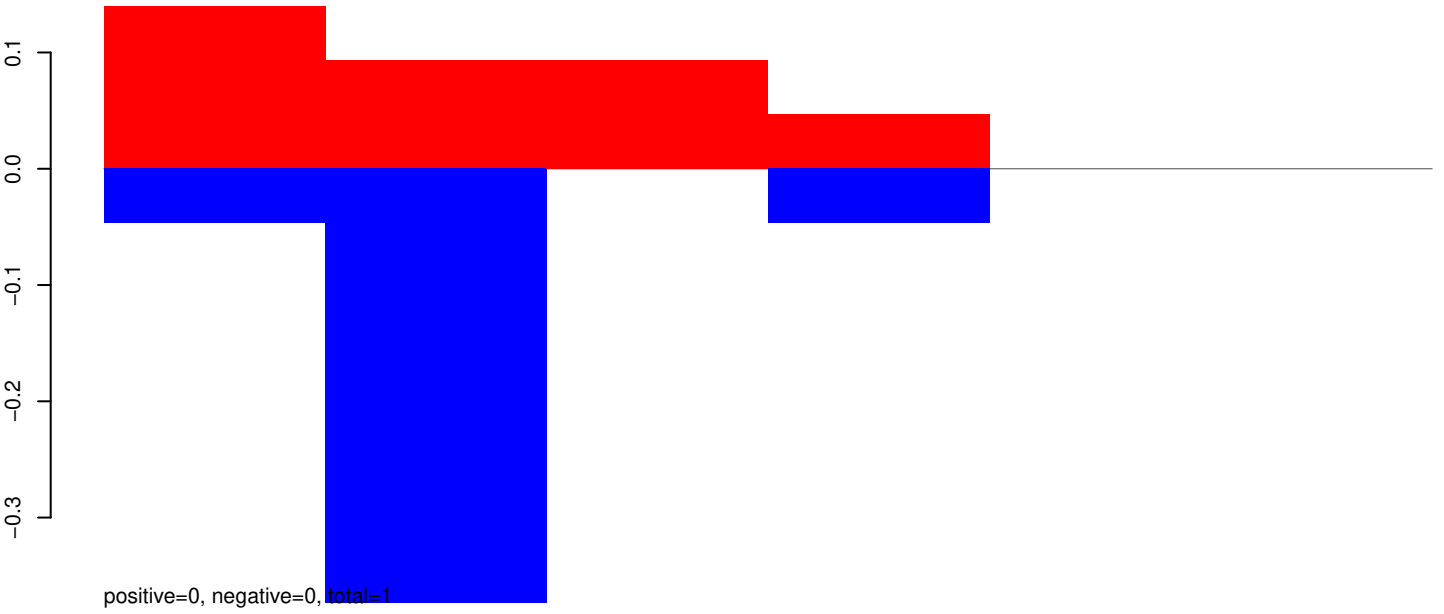
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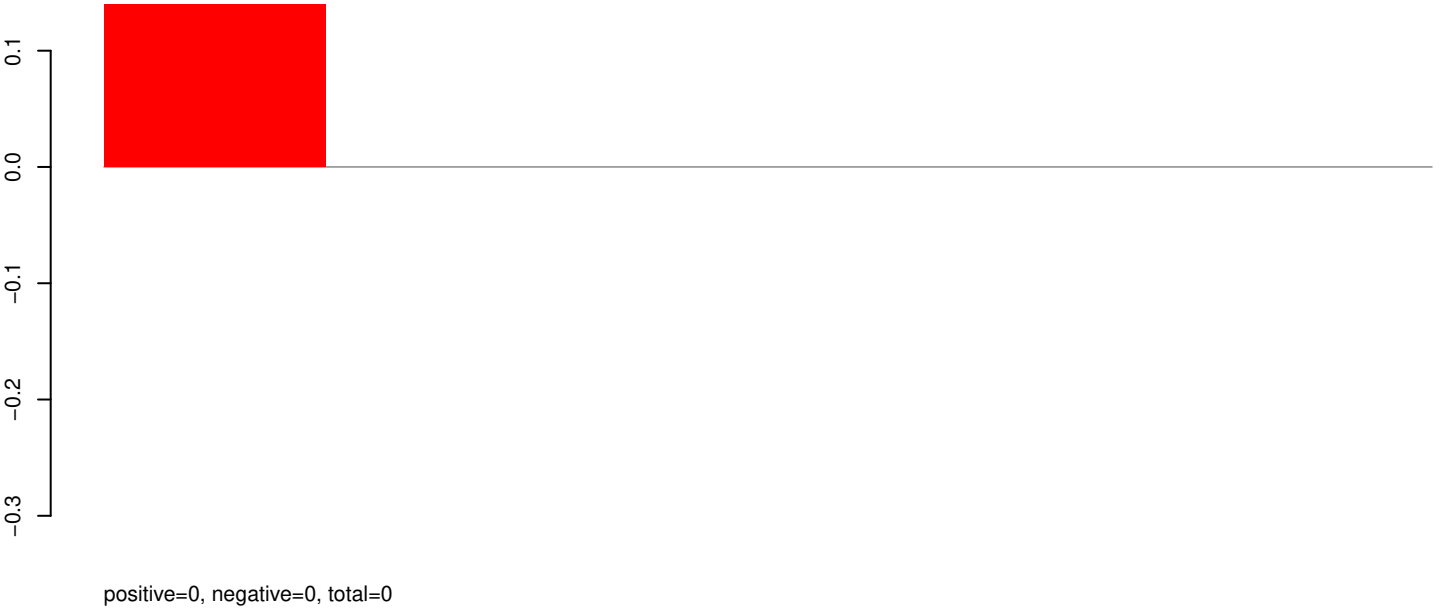
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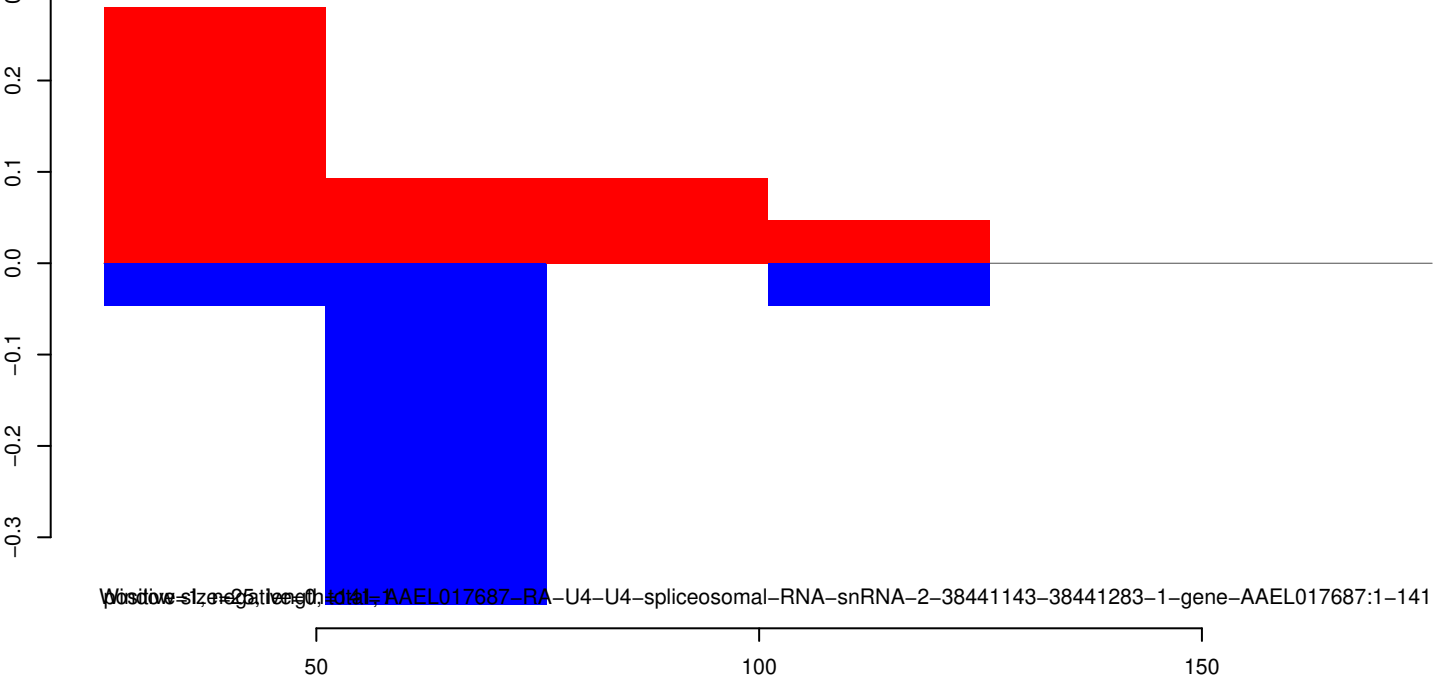
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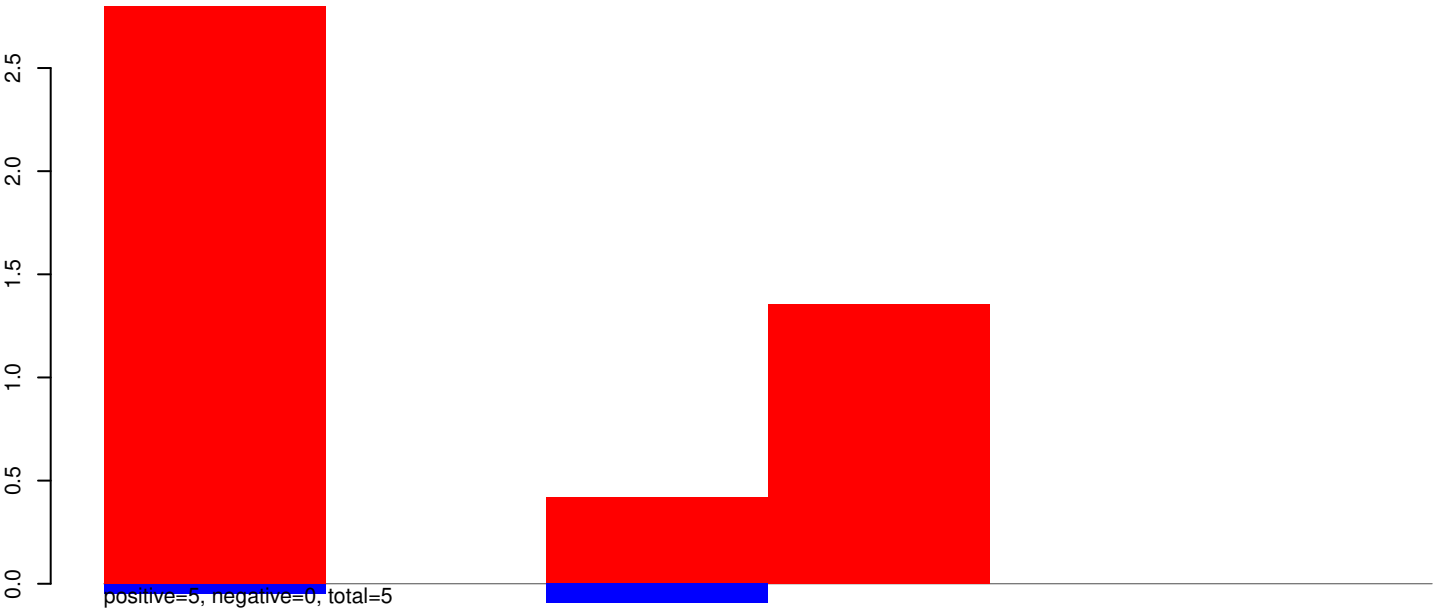
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AeAeg_Aag2SINV_dsLuc.rep



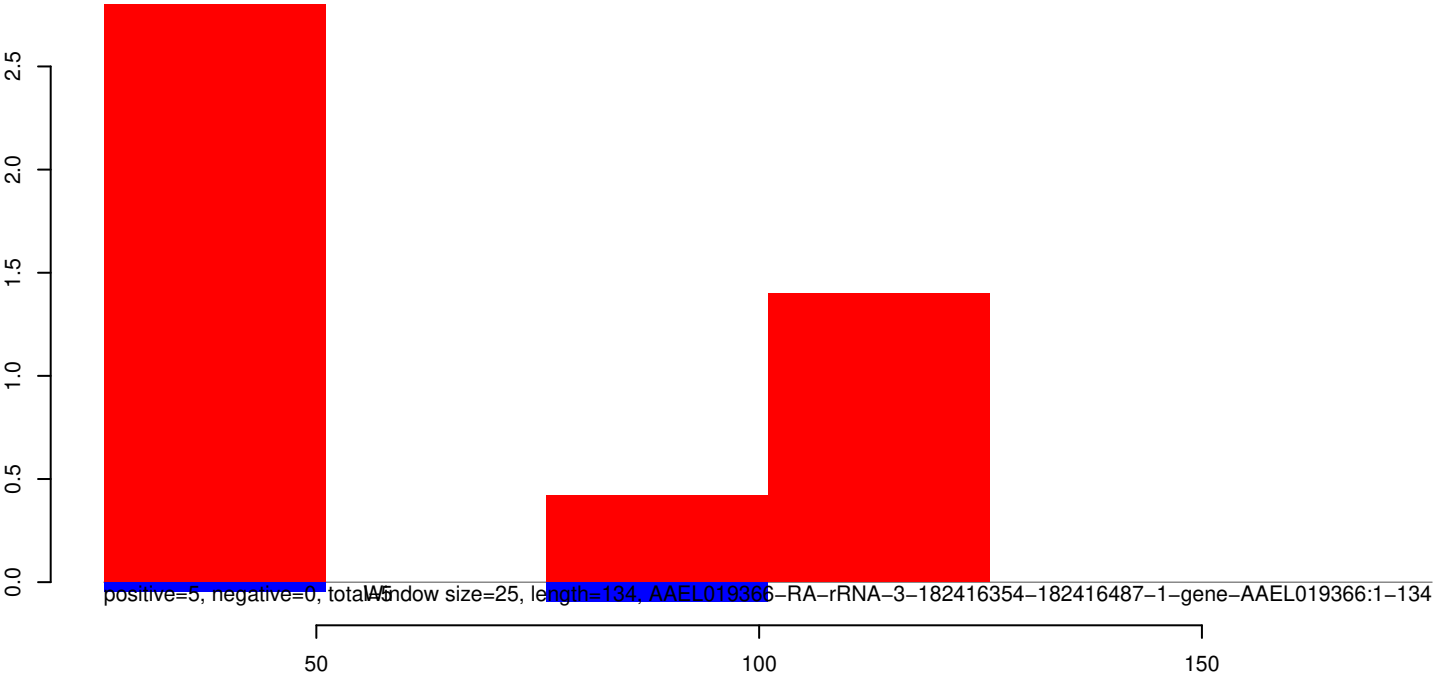
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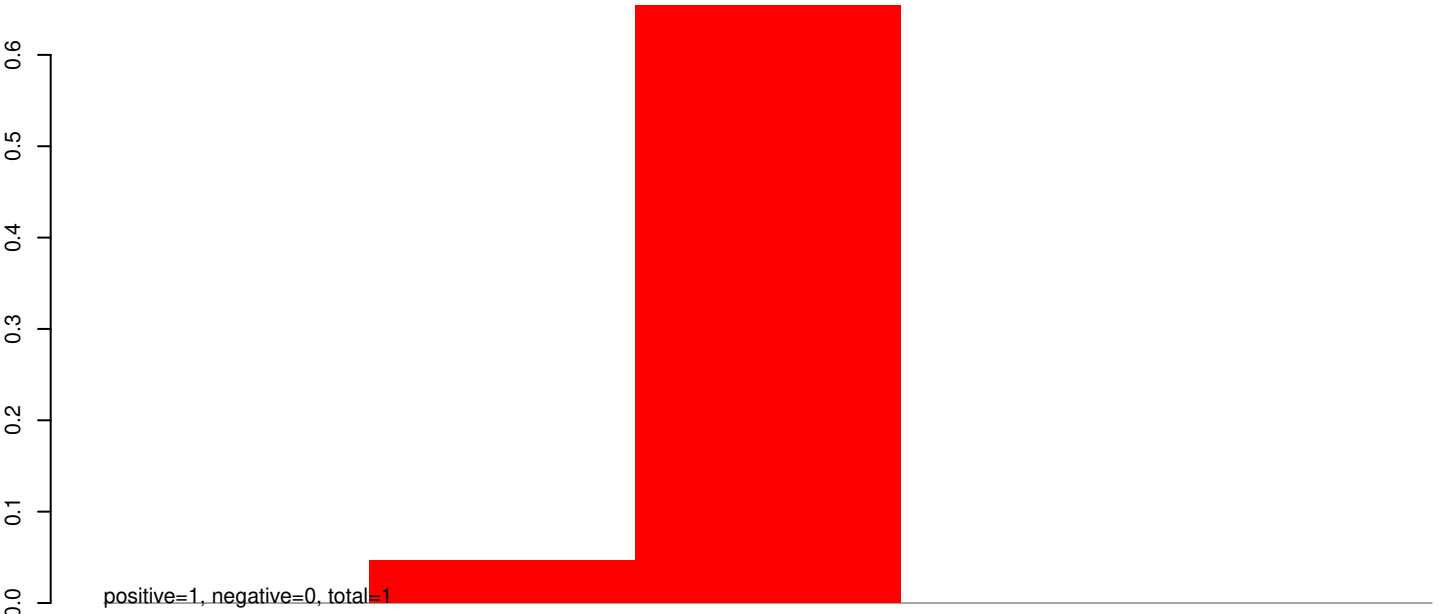
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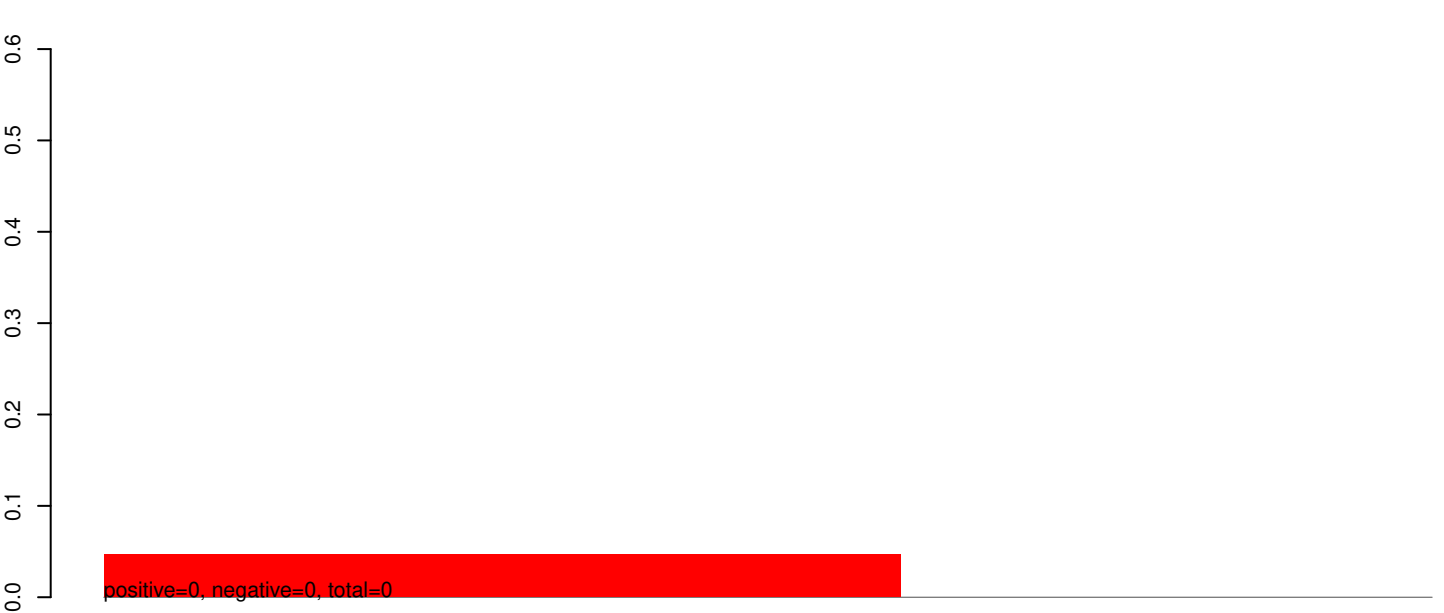
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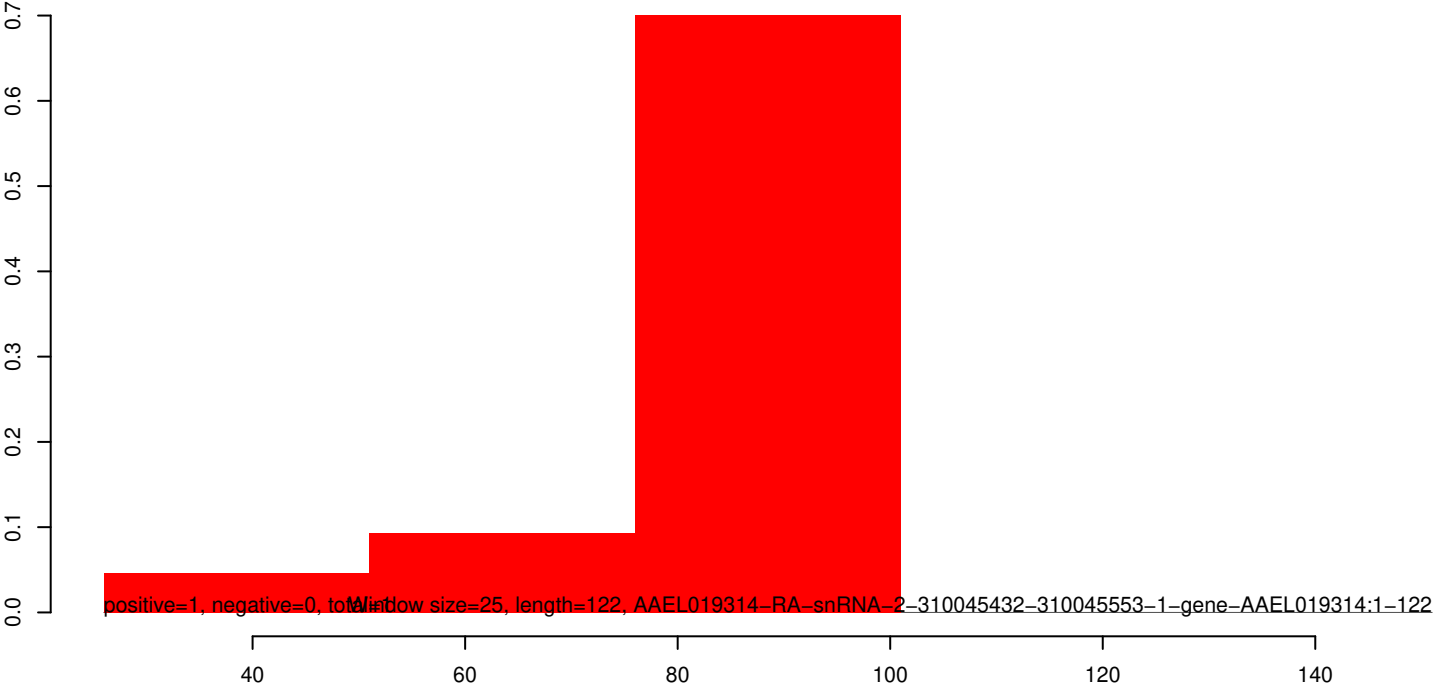
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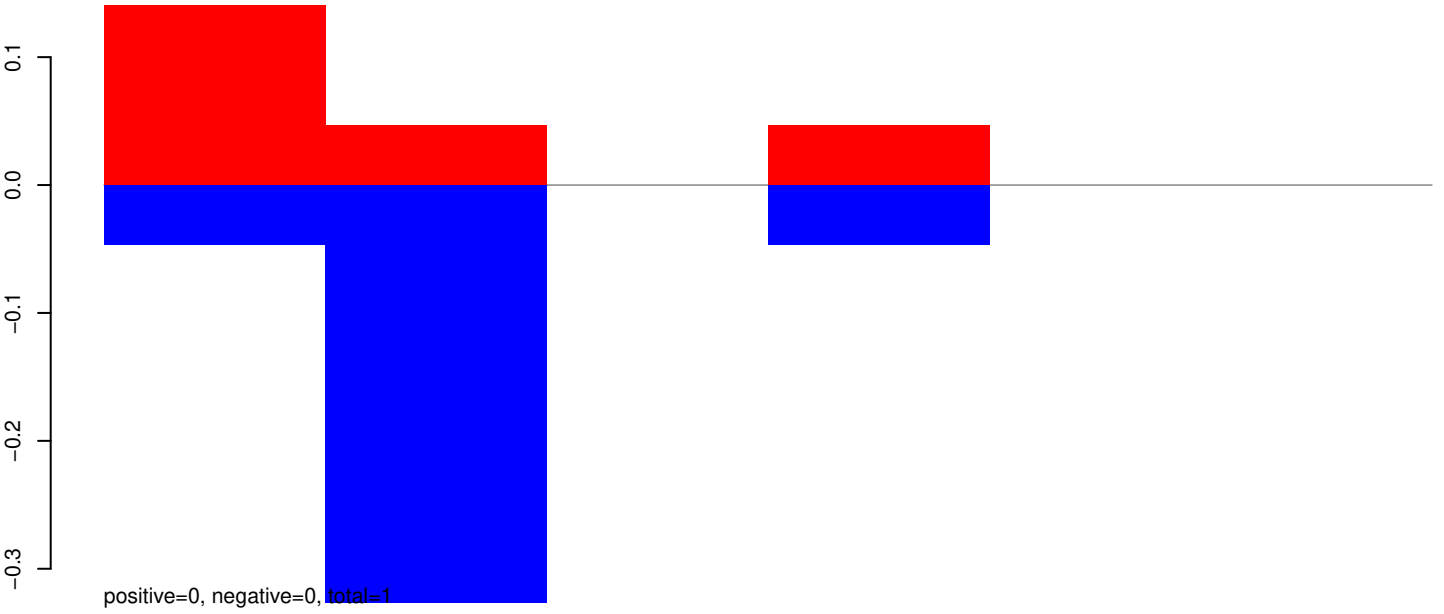
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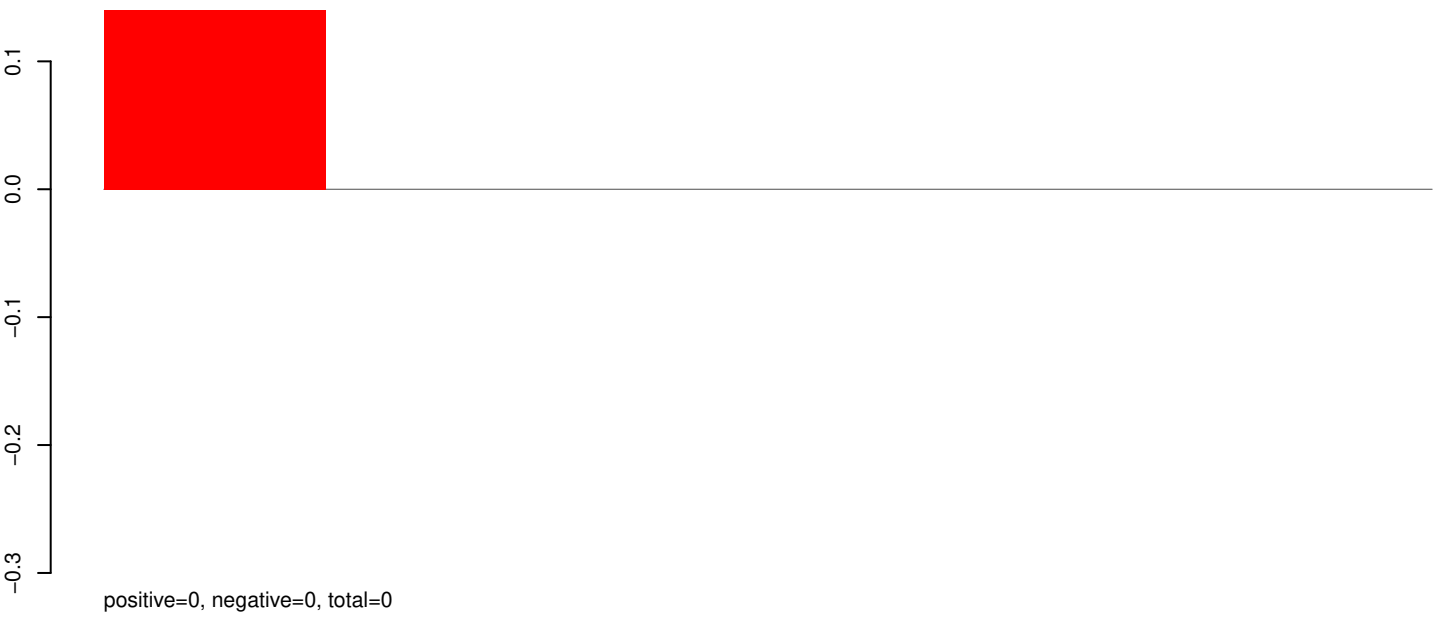
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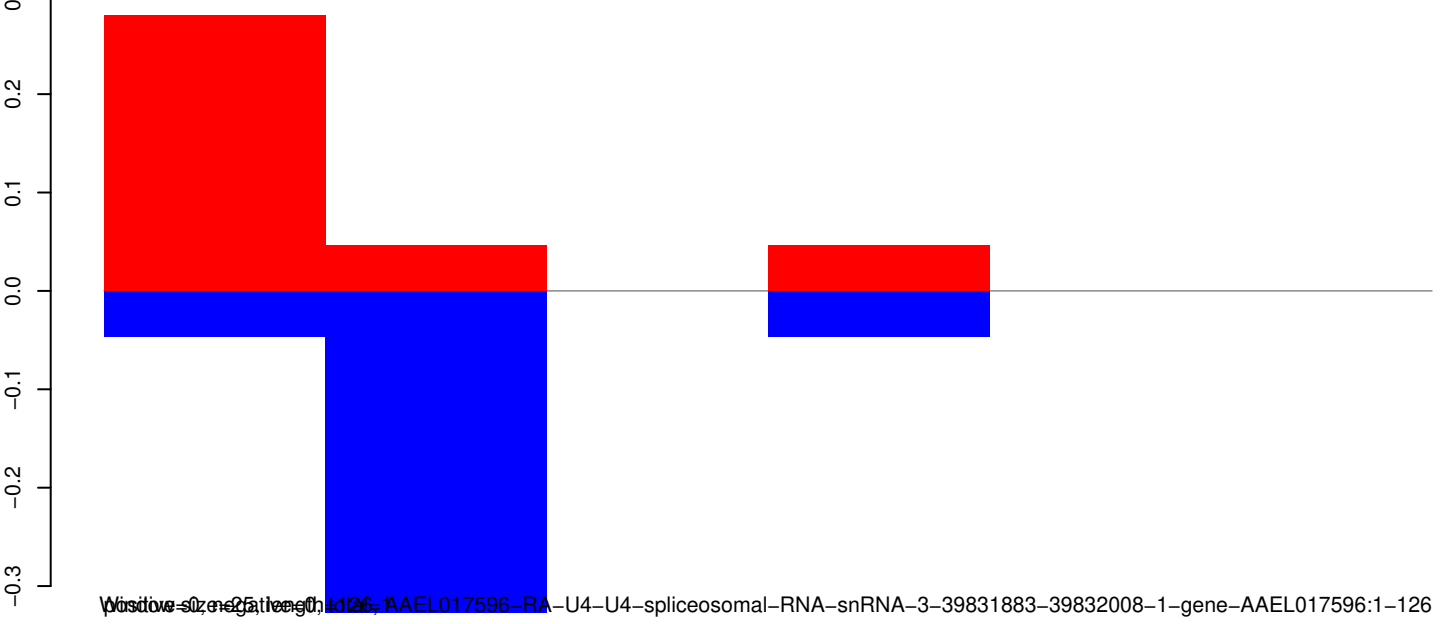
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AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



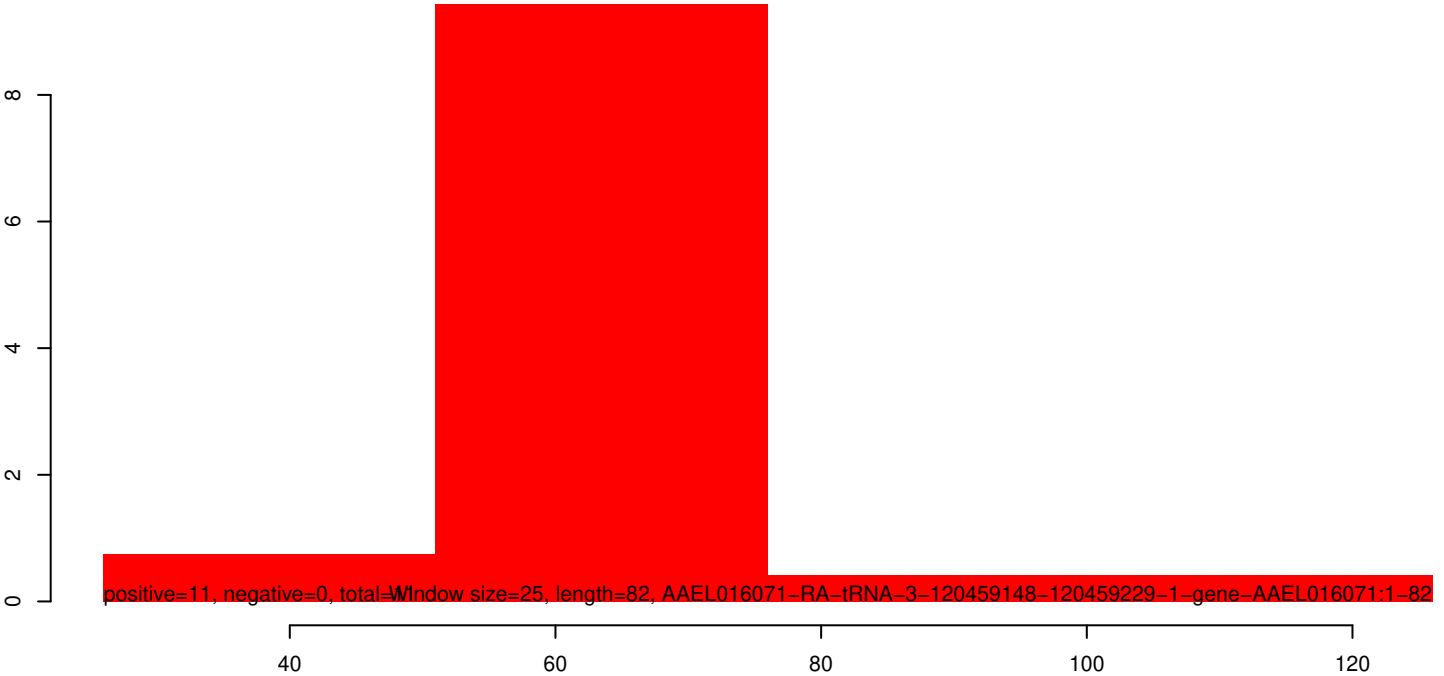
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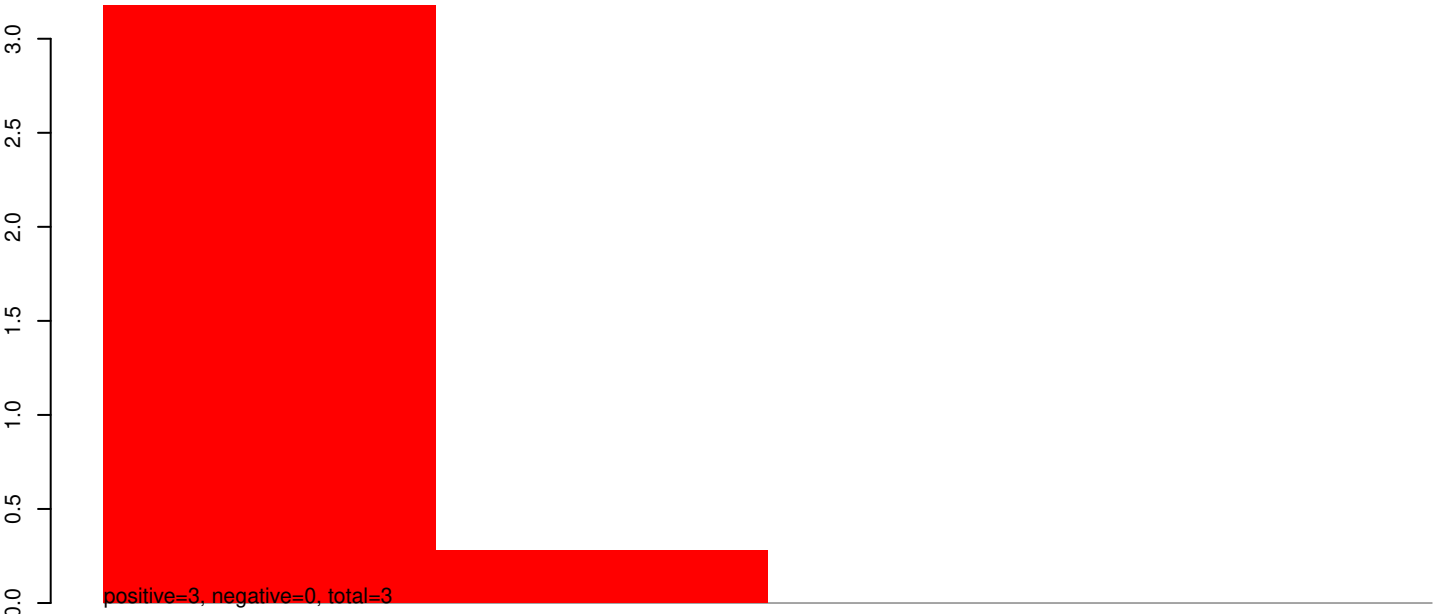
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AeAeg_Aag2SINV_dsLuc.rep



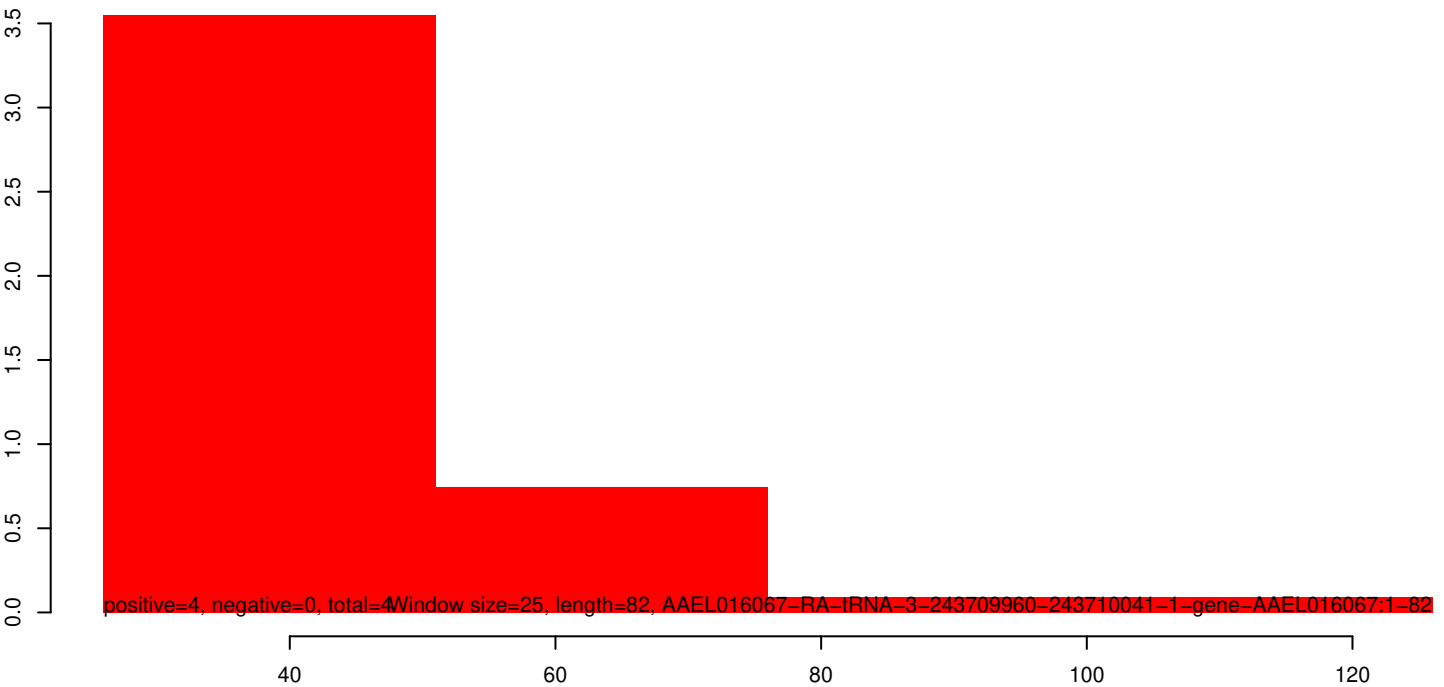
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AeAeg_Aag2SINV_dsLuc.24_35.rep



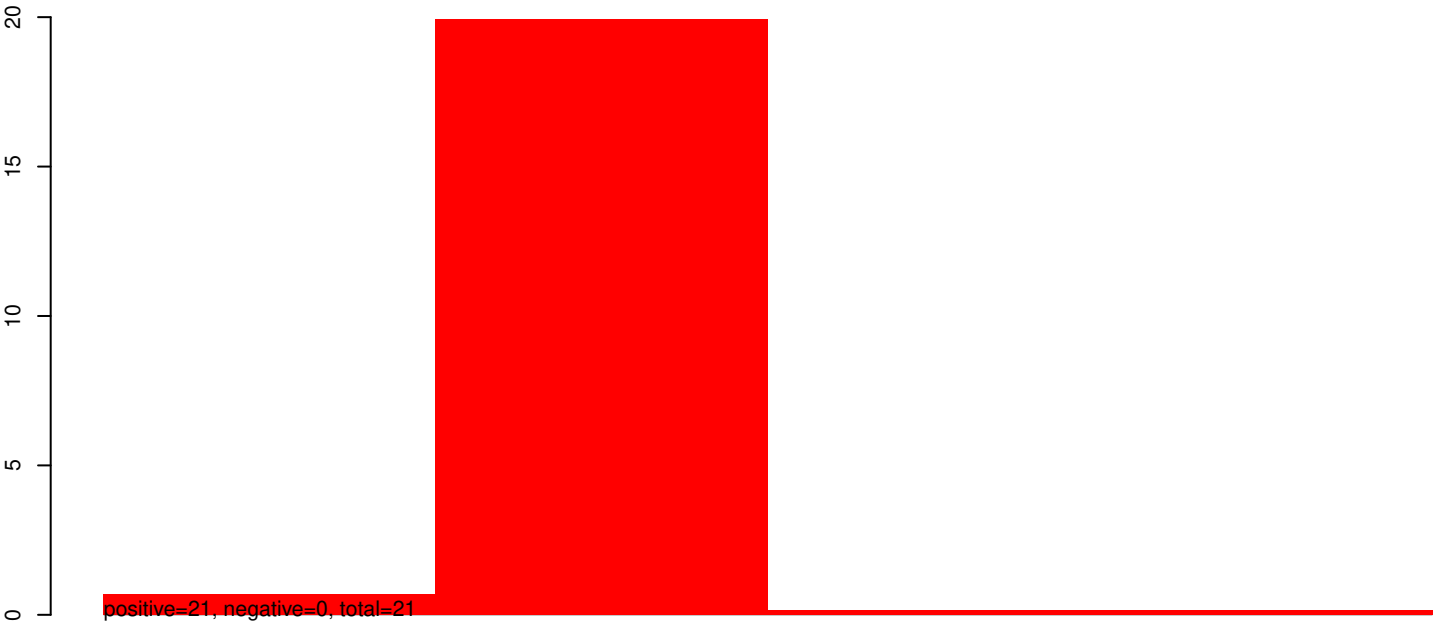
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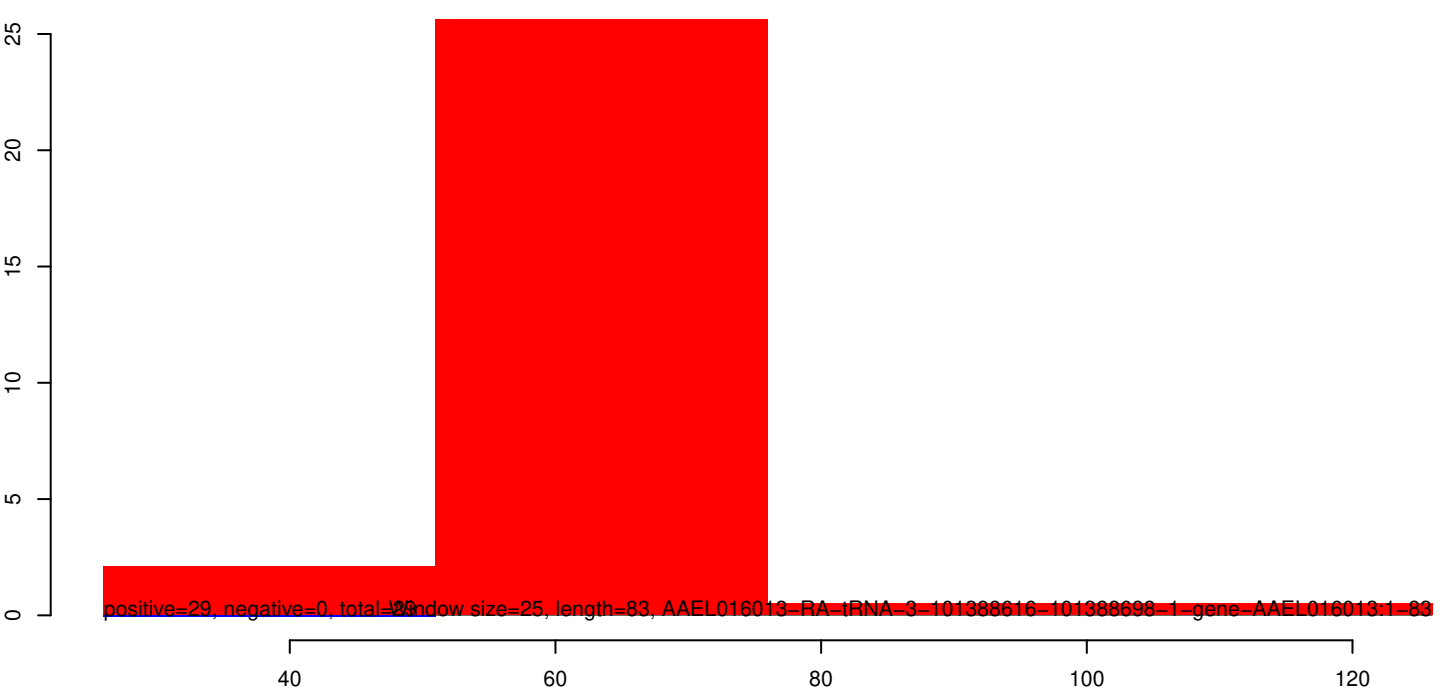
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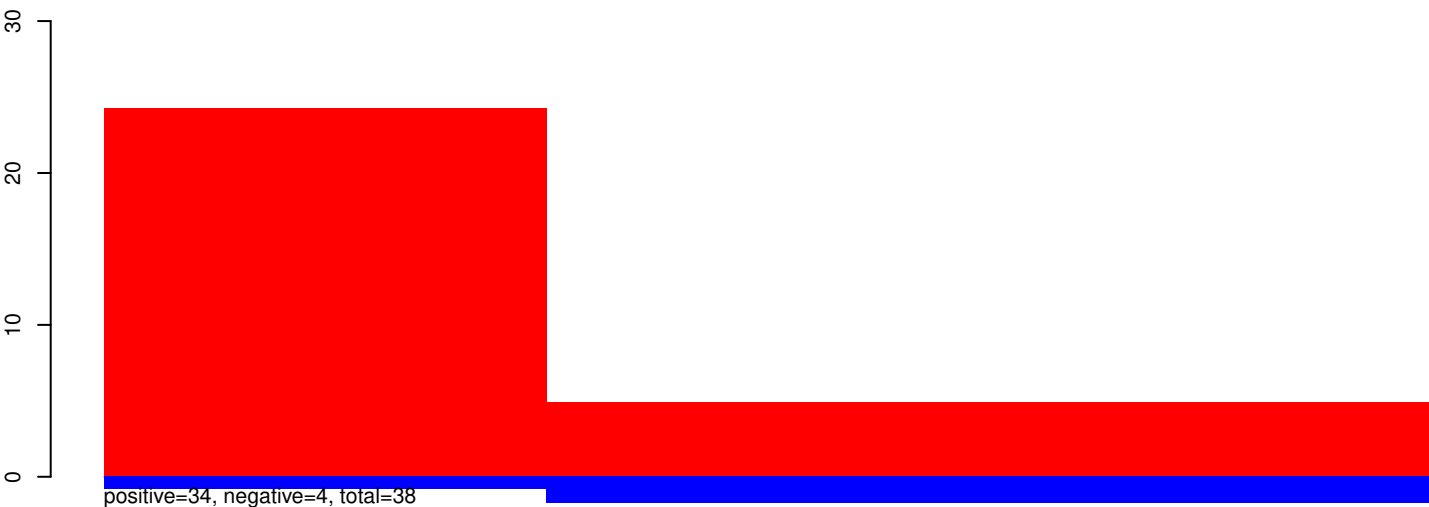
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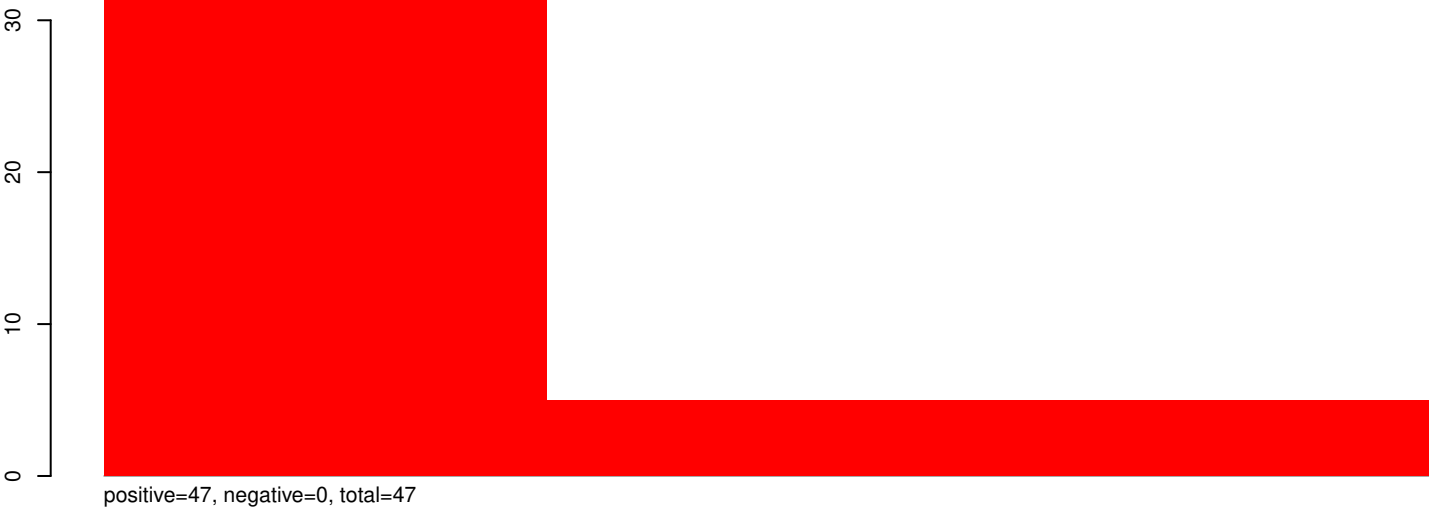
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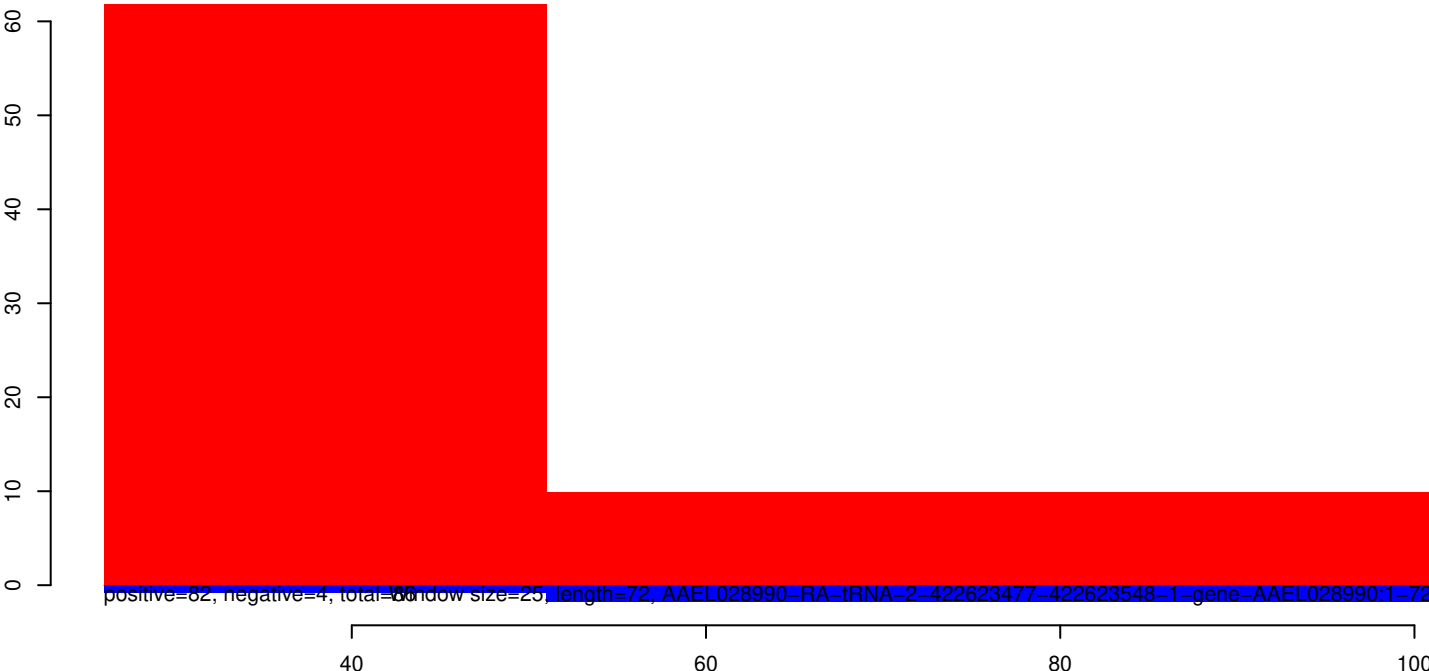
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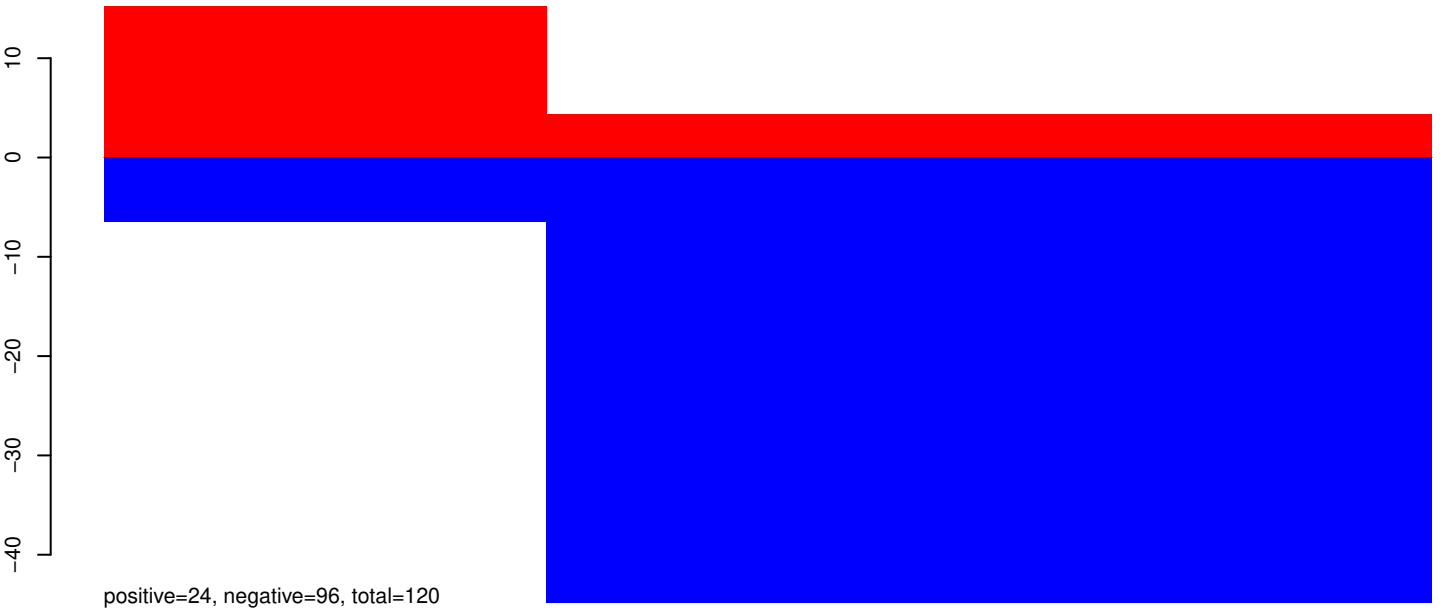
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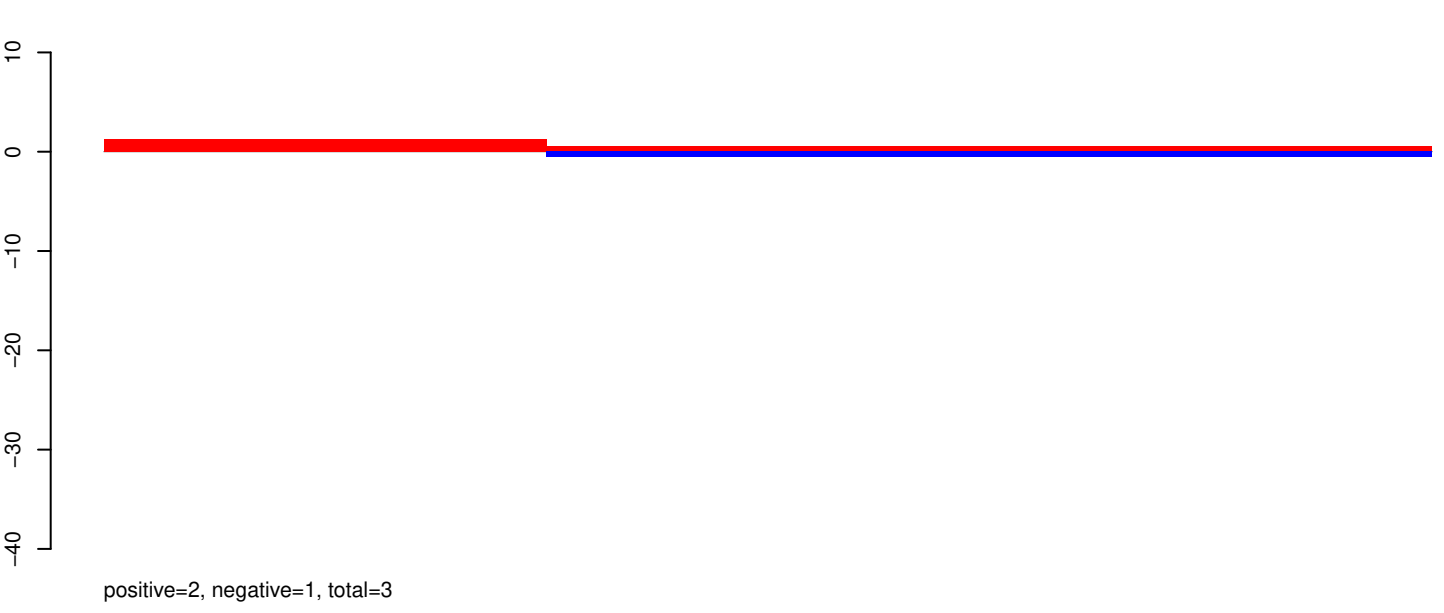
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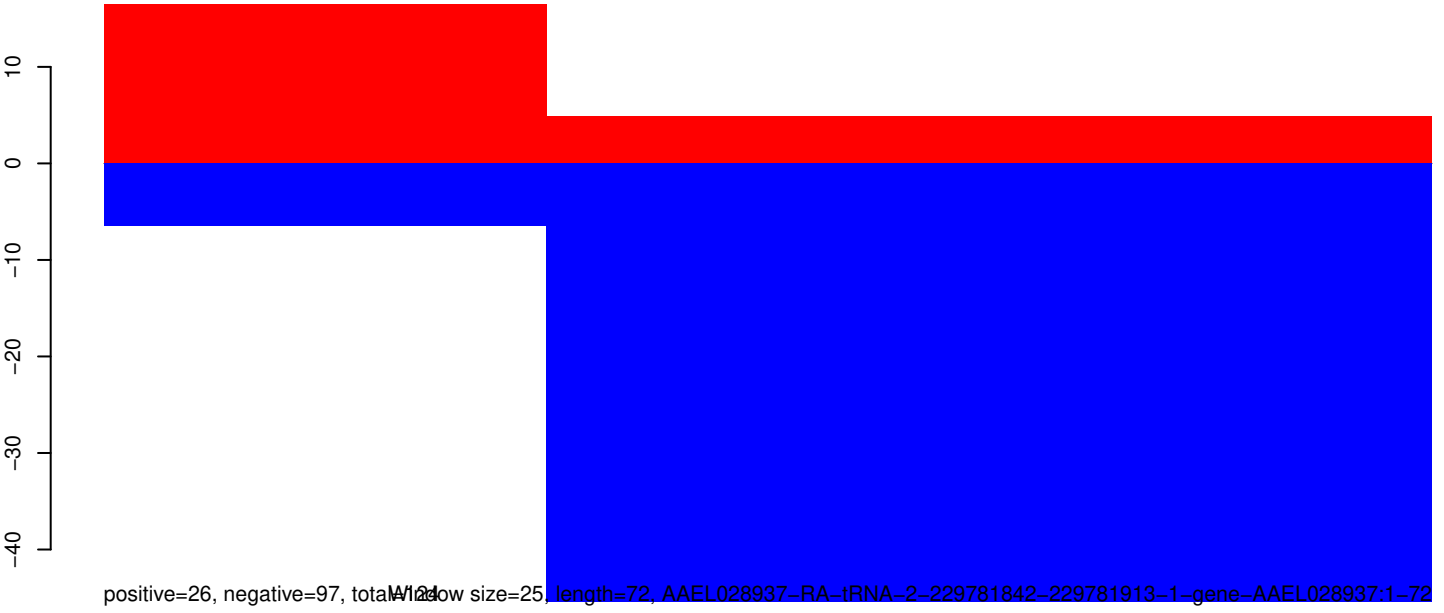
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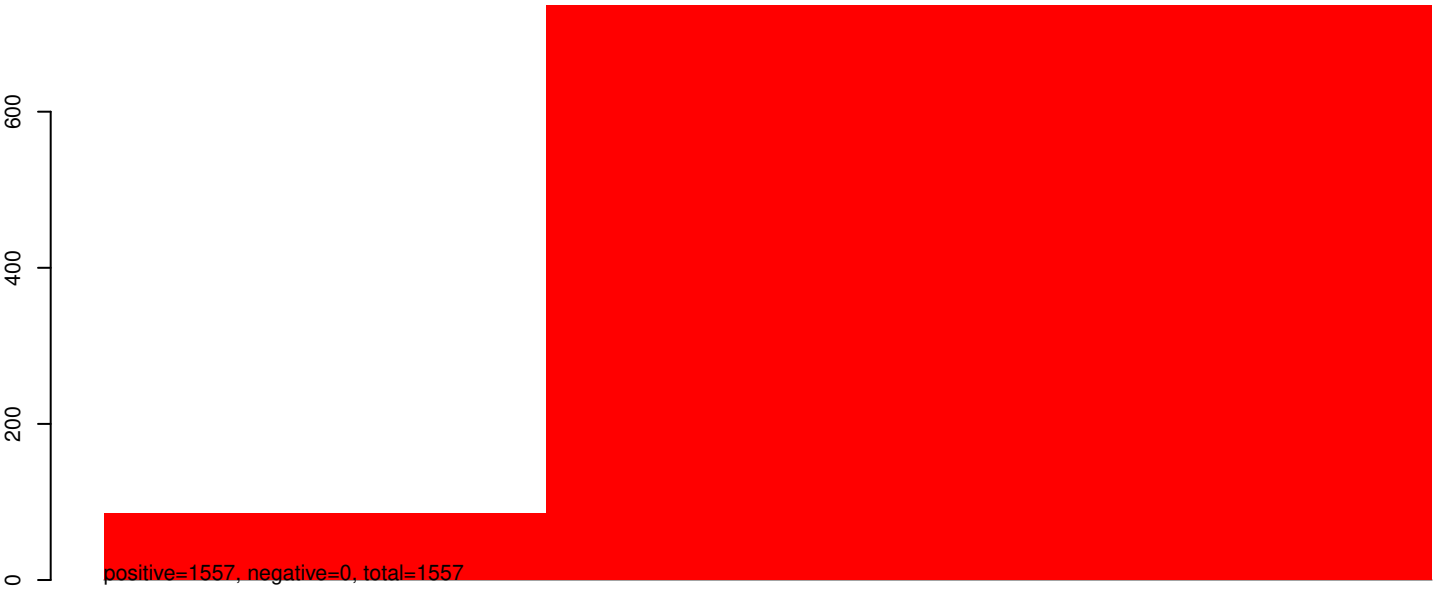
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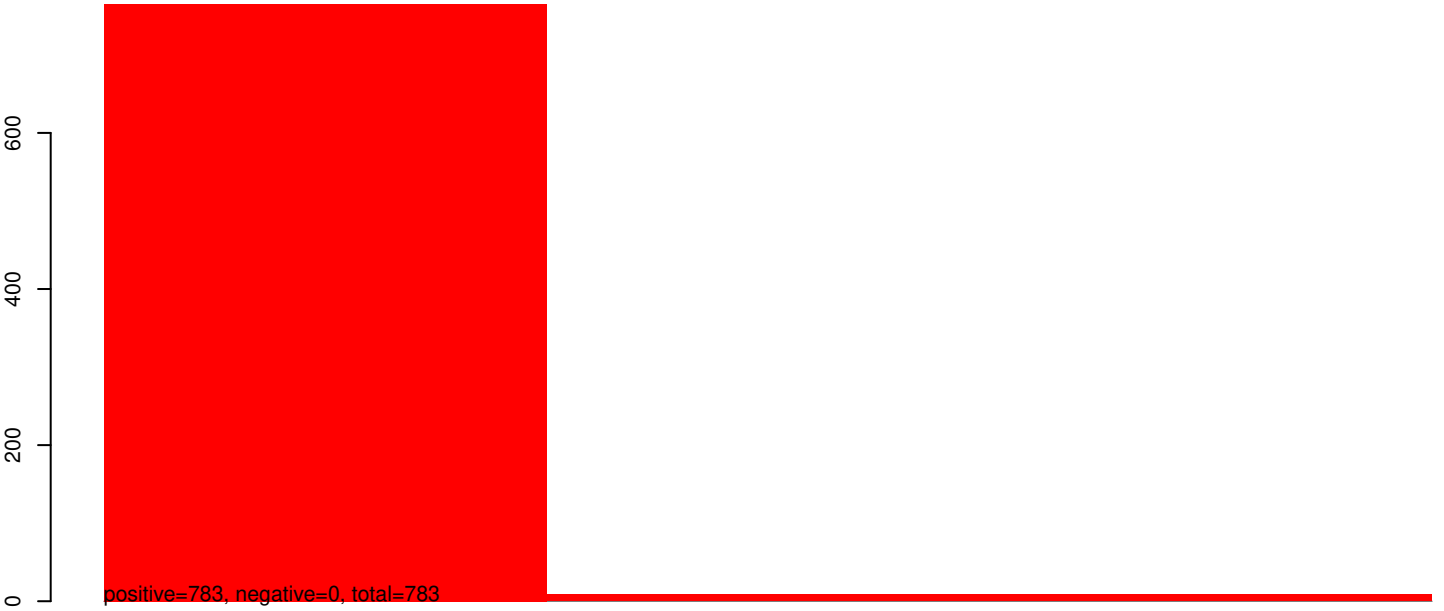
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40 60 80 100

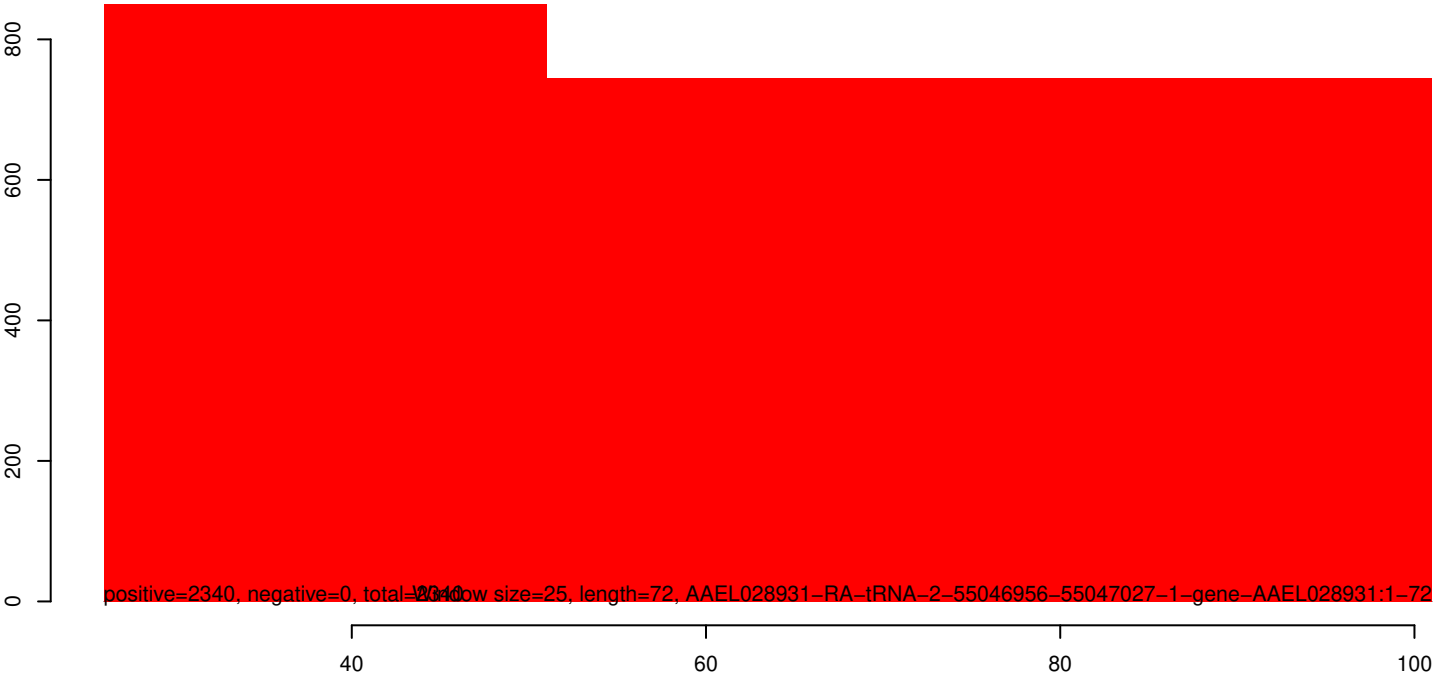
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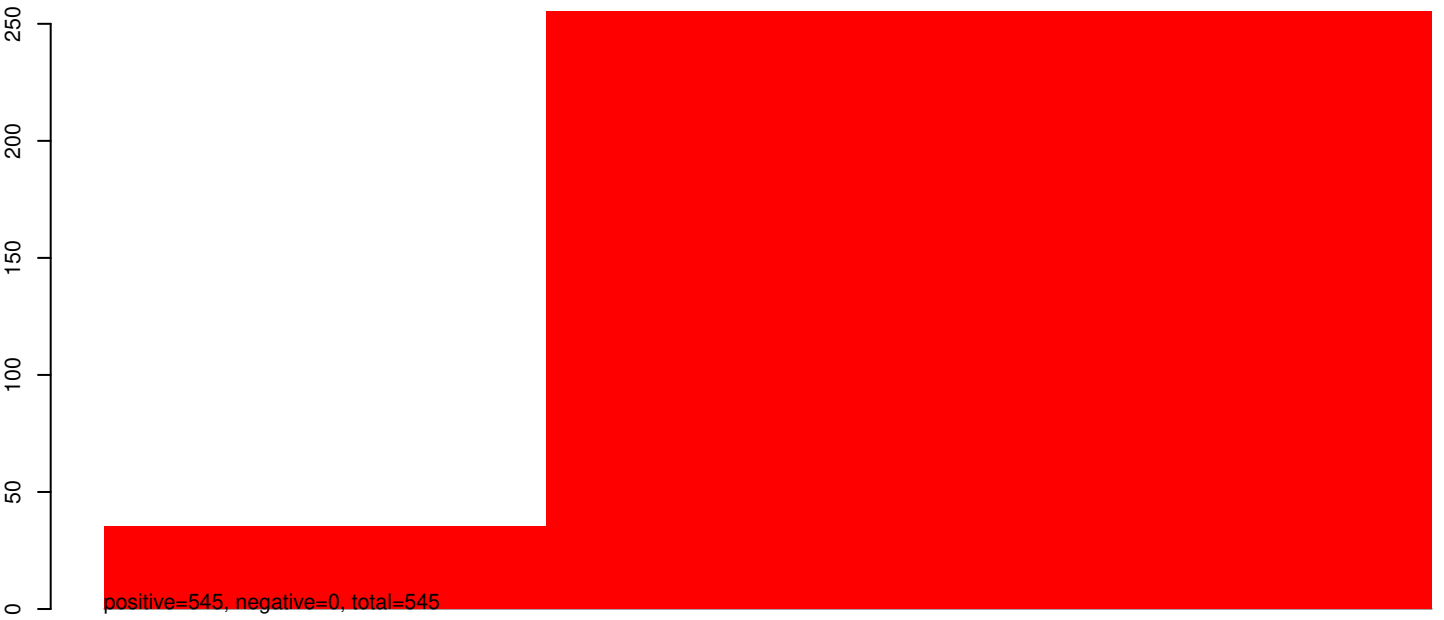
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AeAeg_Aag2SINV_dsLuc.rep



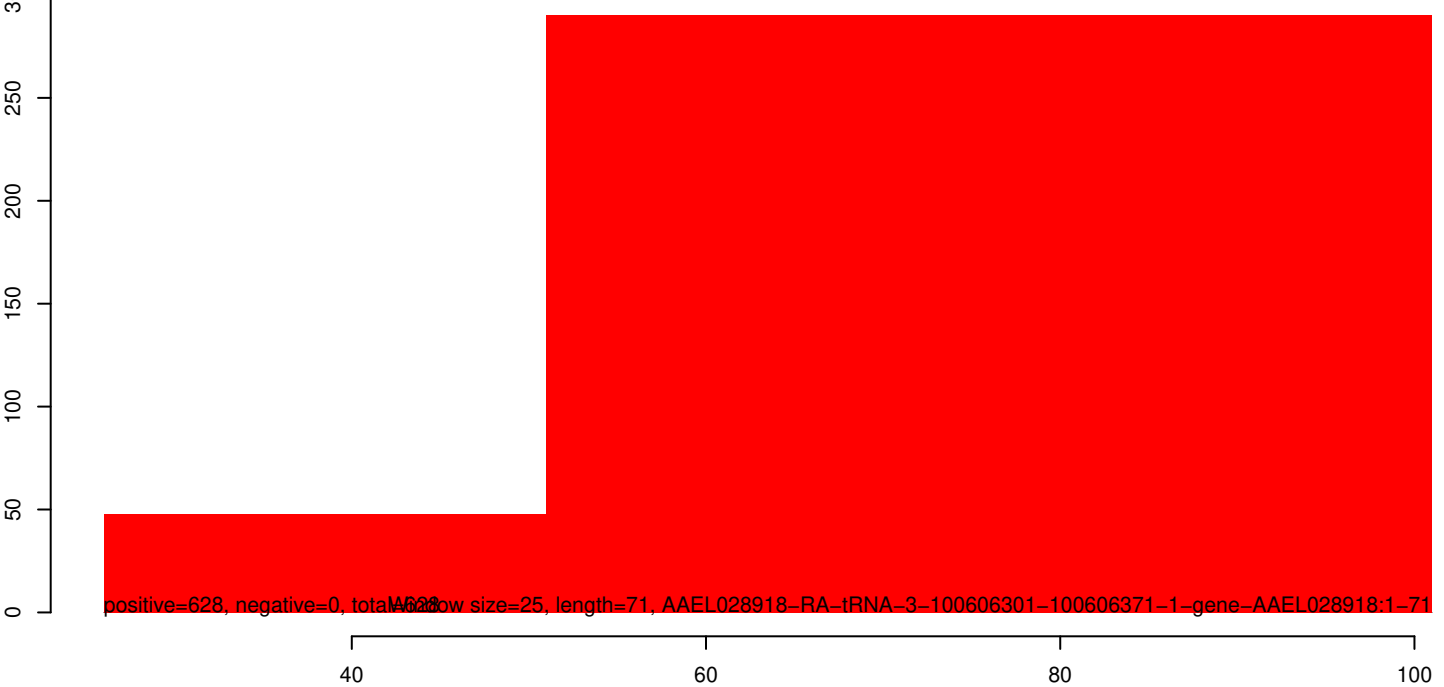
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AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



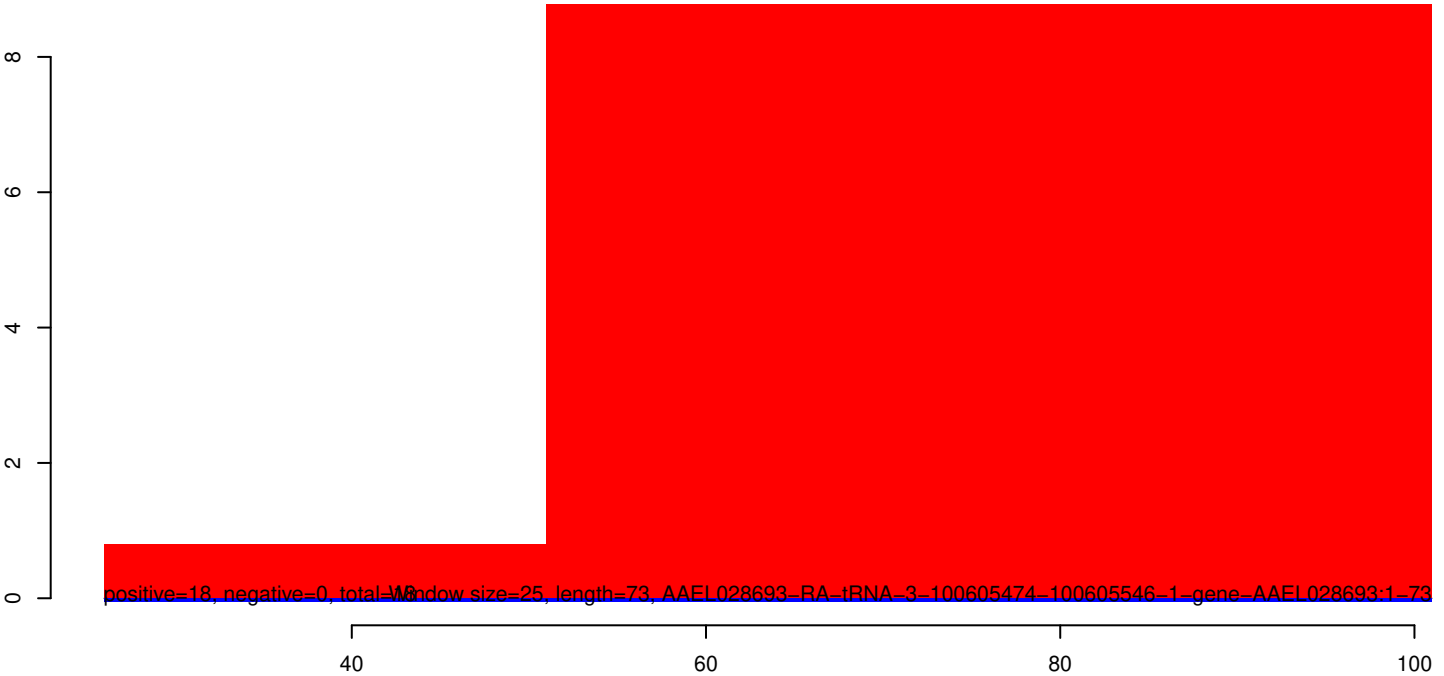
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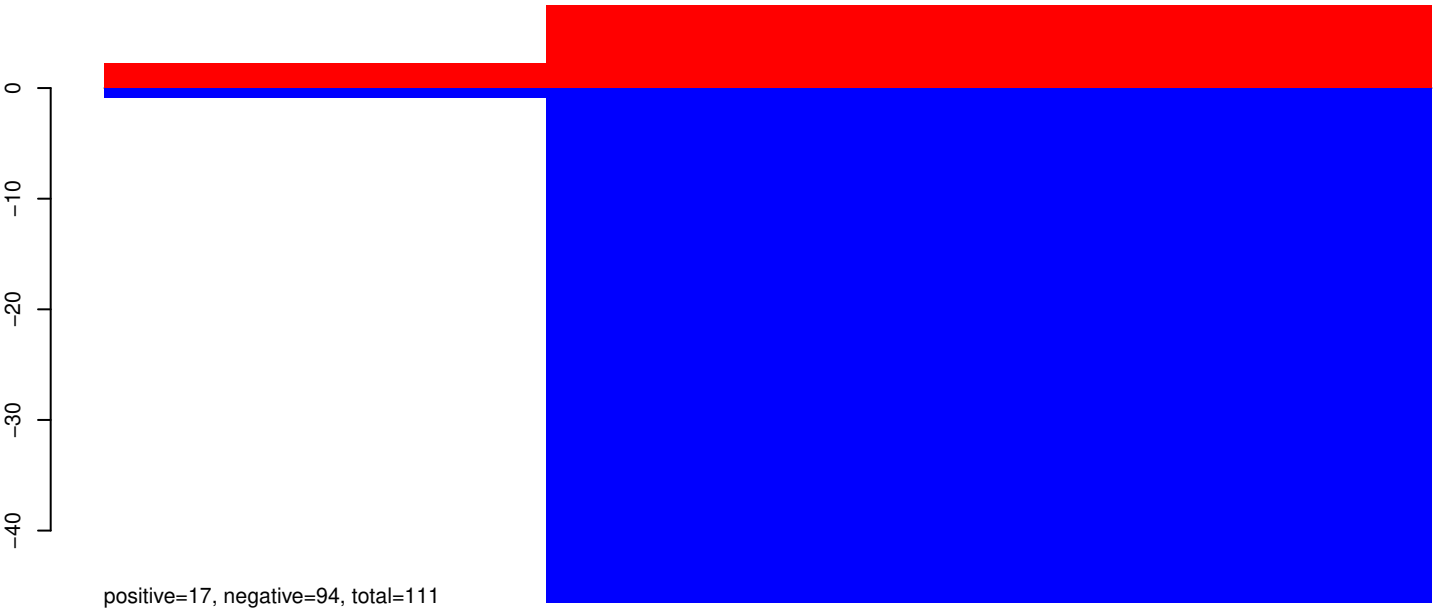
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AeAeg_Aag2SINV_dsLuc.rep



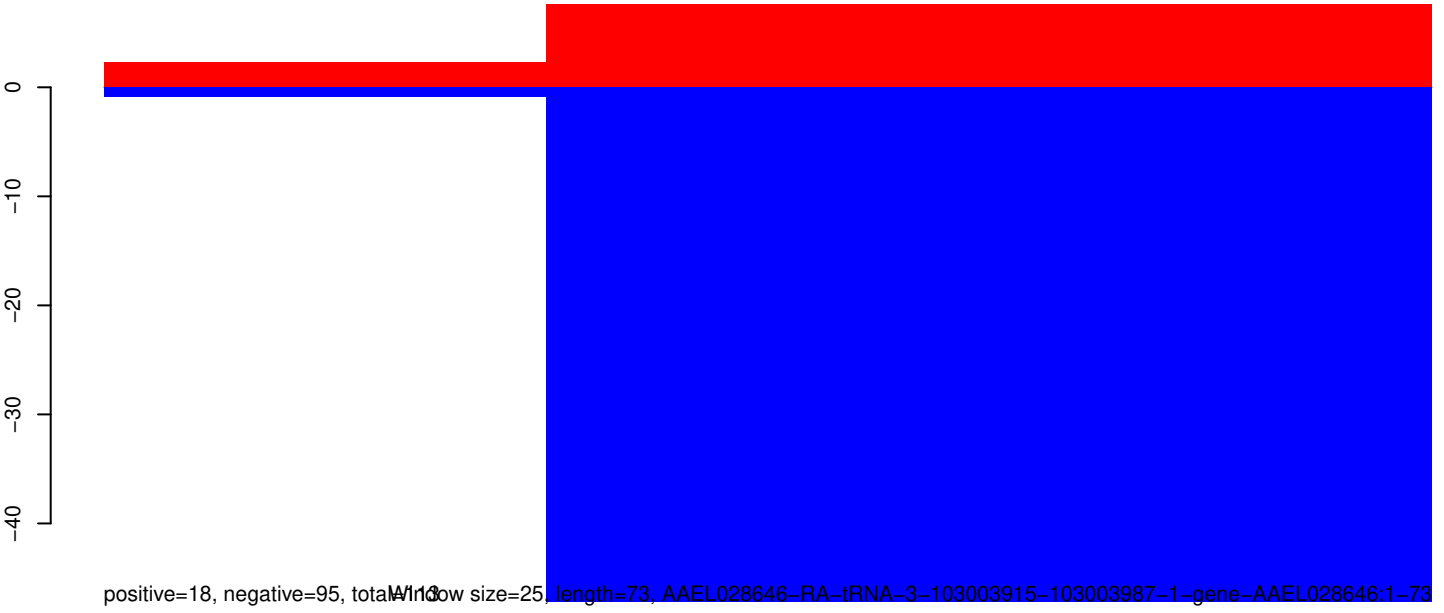
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AeAeg_Aag2SINV_dsLuc.24_35.rep



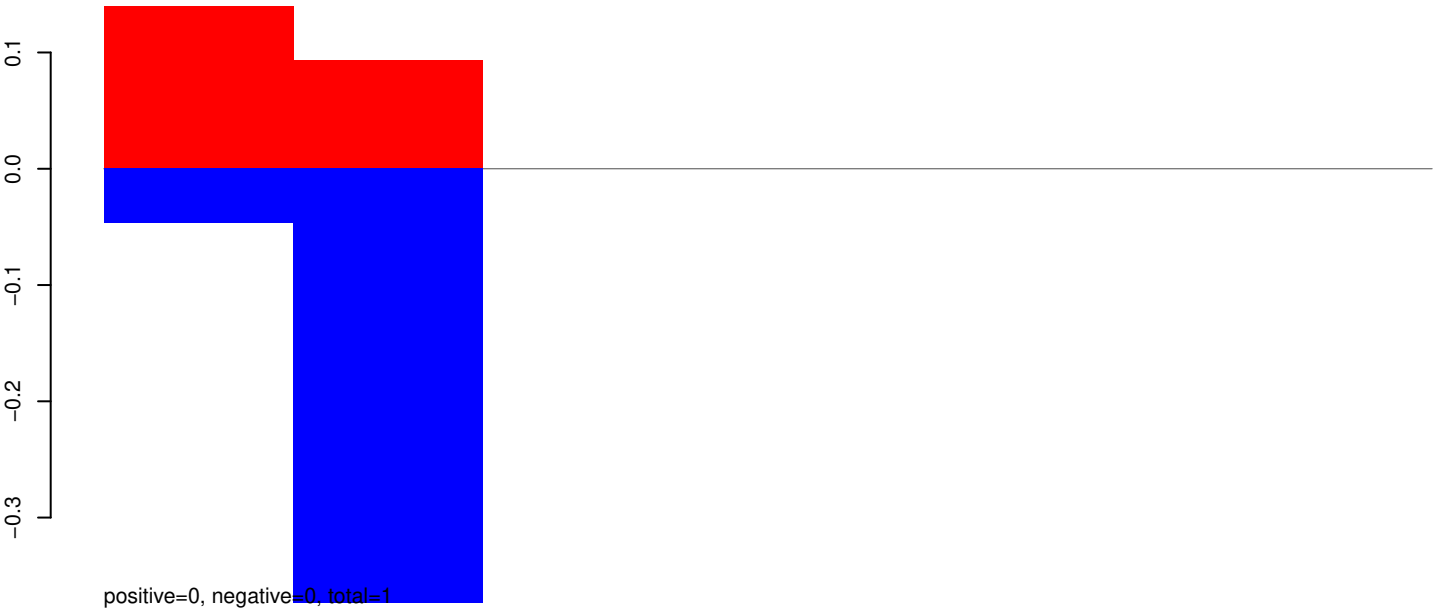
AeAeg_Aag2SINV_dsLuc.rep



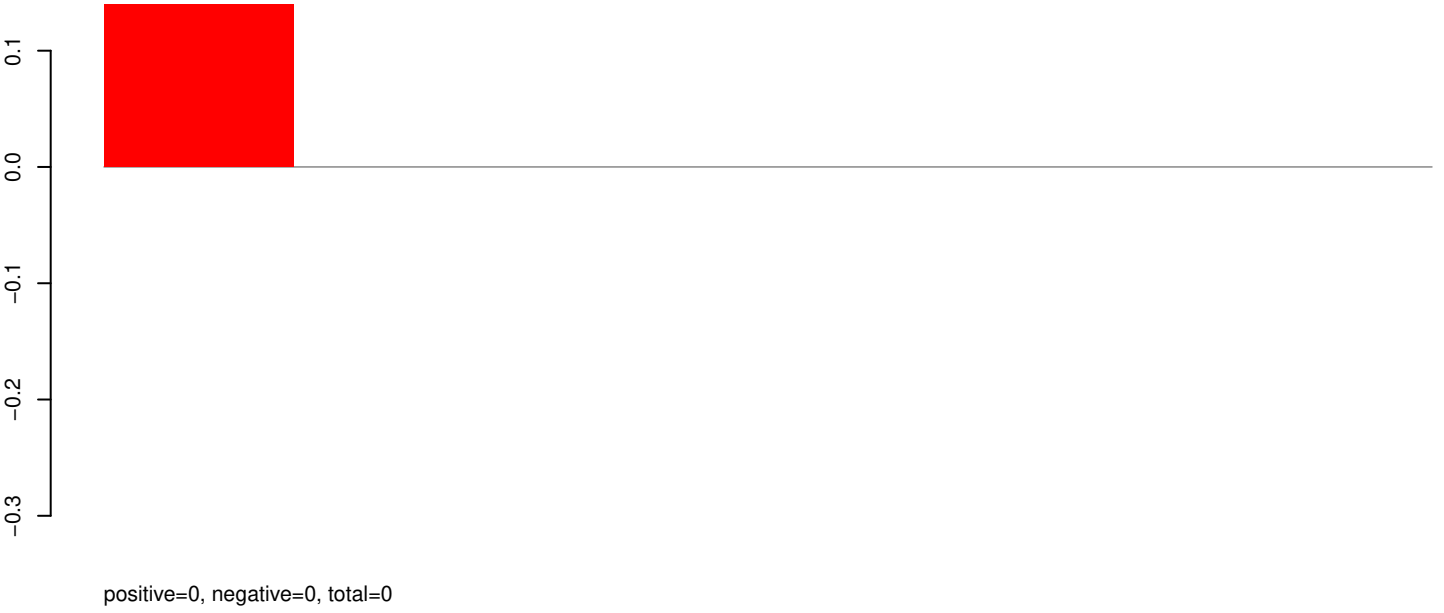
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40 60 80 100

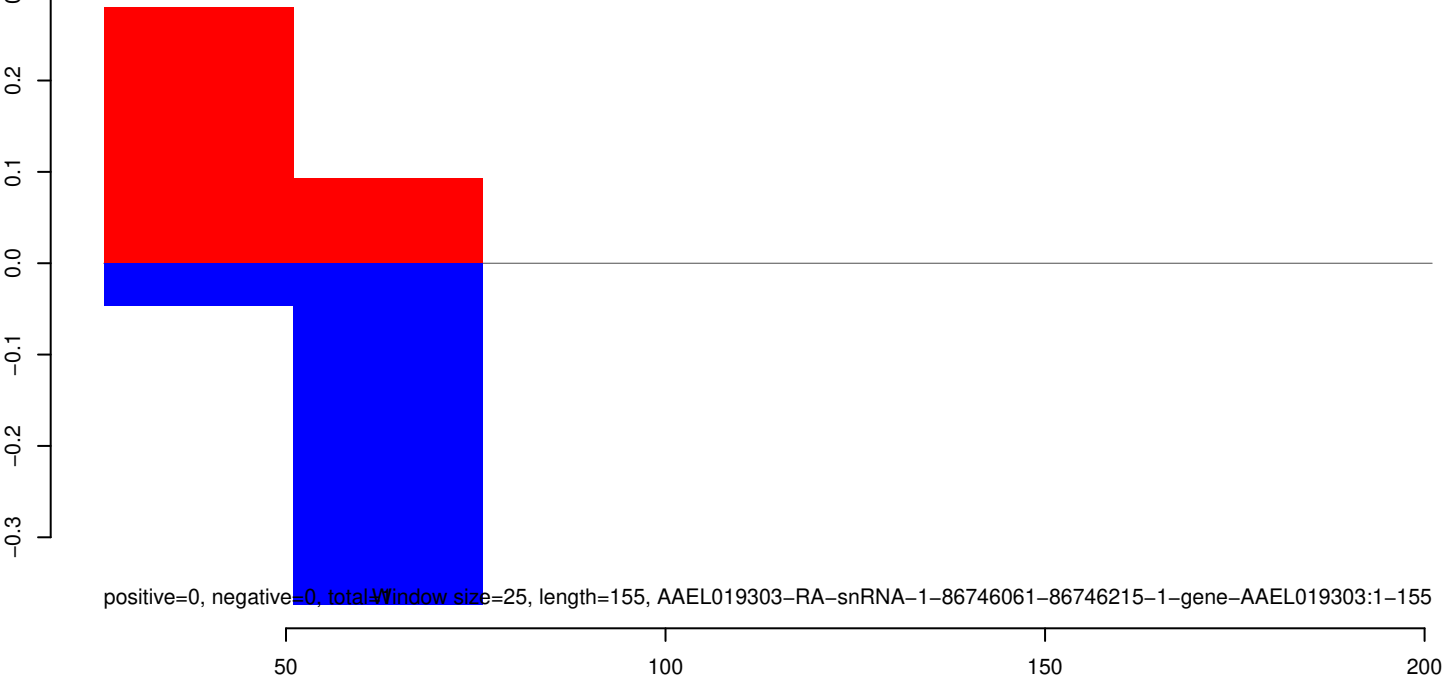
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



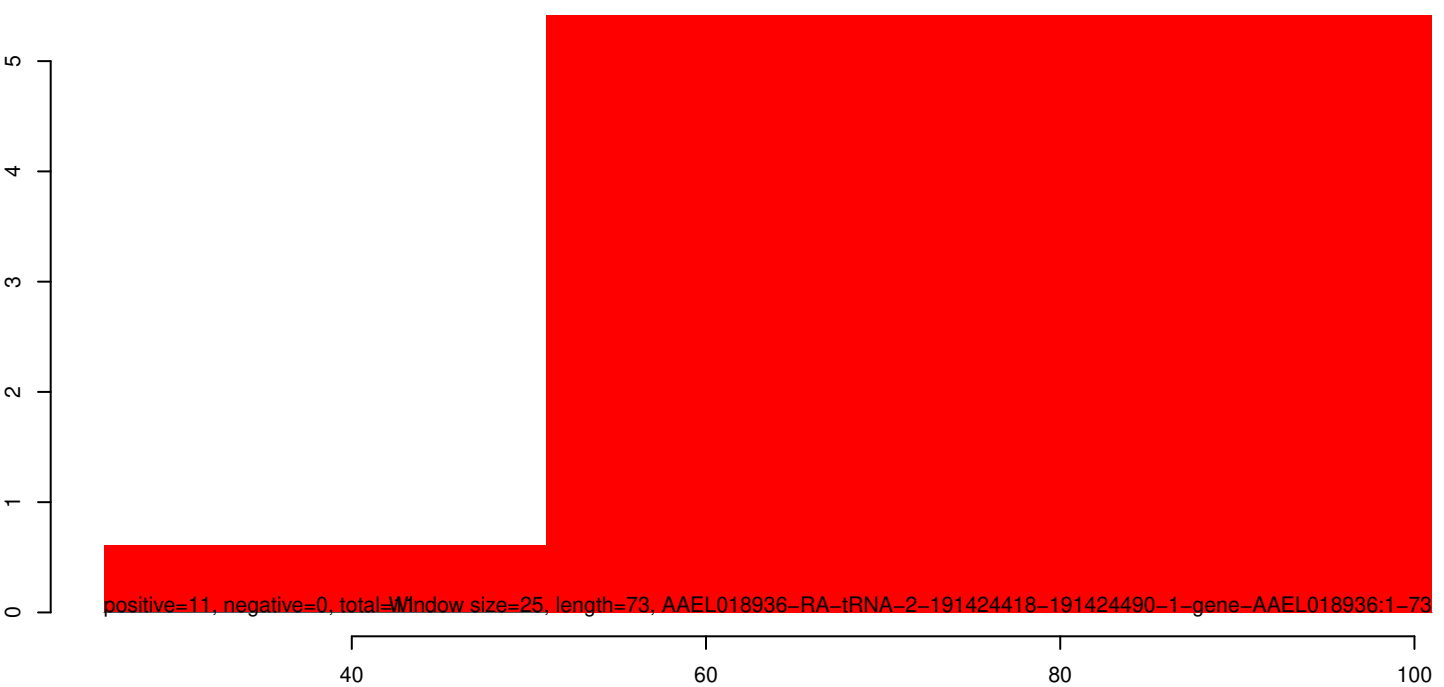
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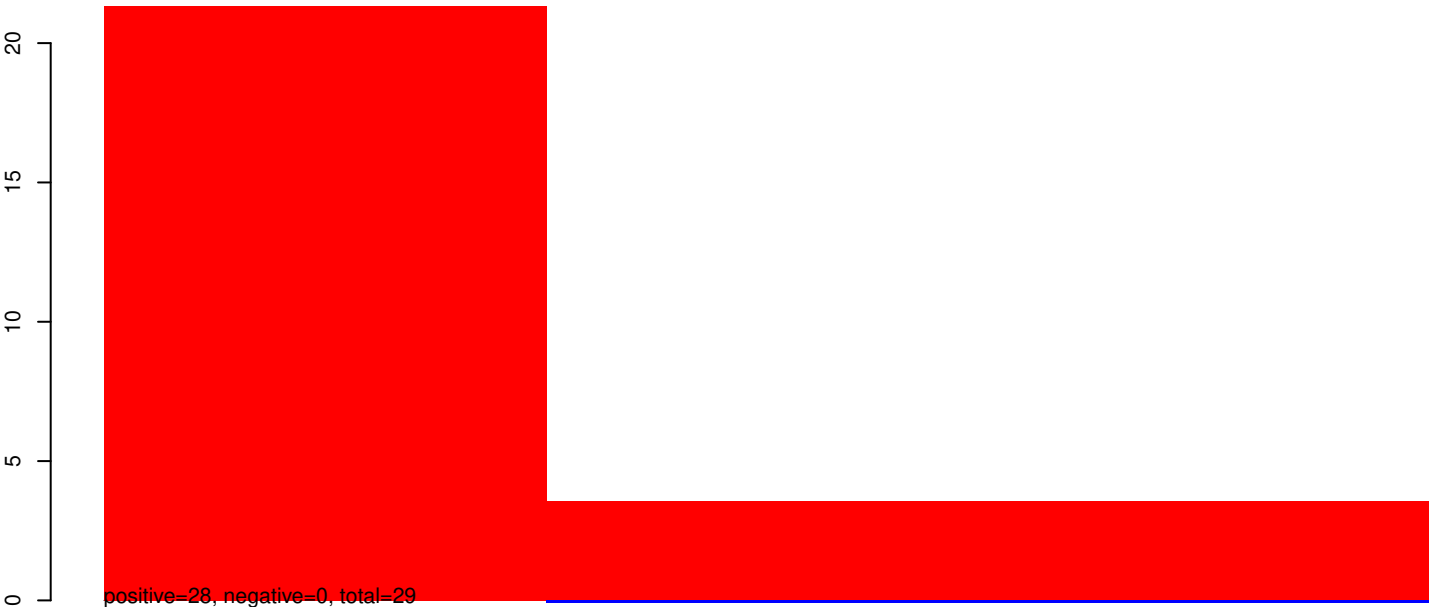
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AeAeg_Aag2SINV_dsLuc.rep



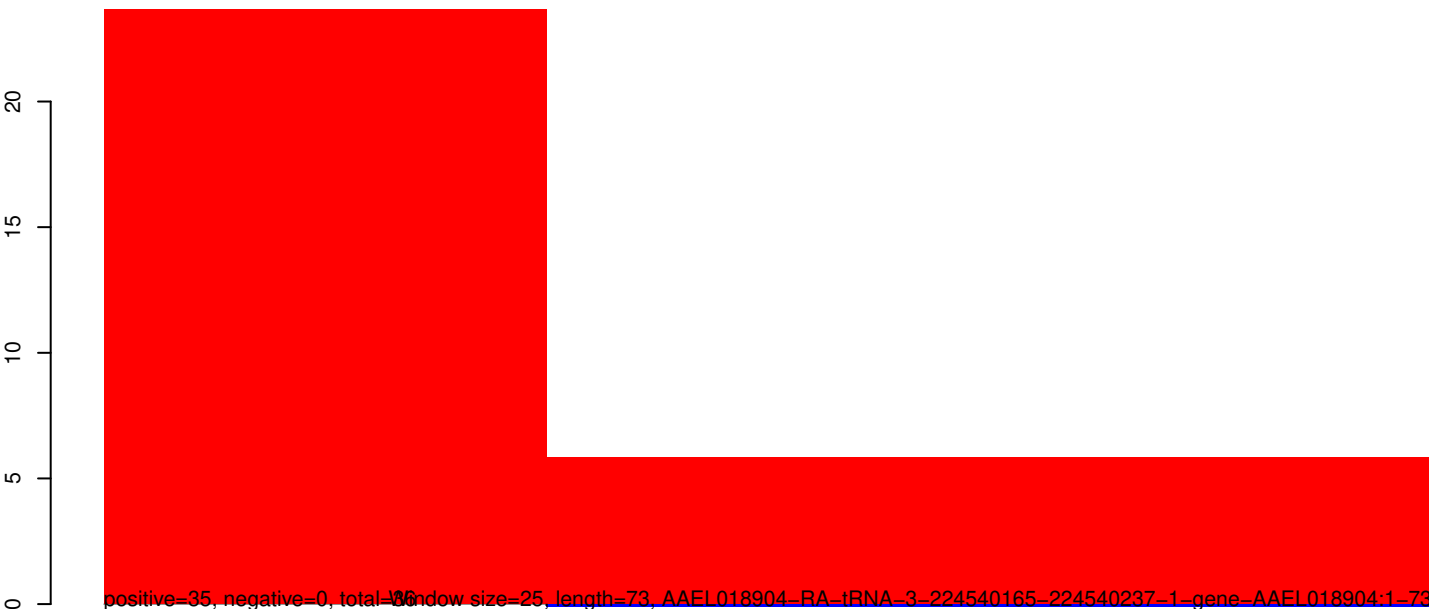
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AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



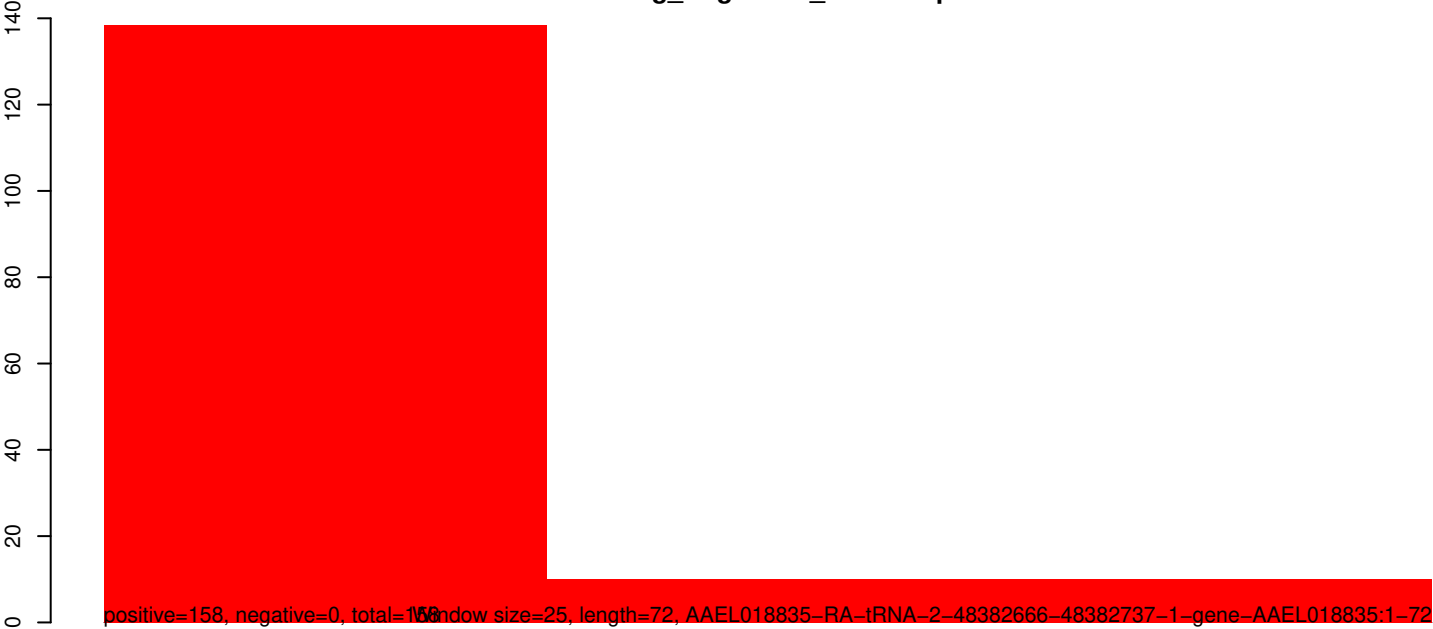
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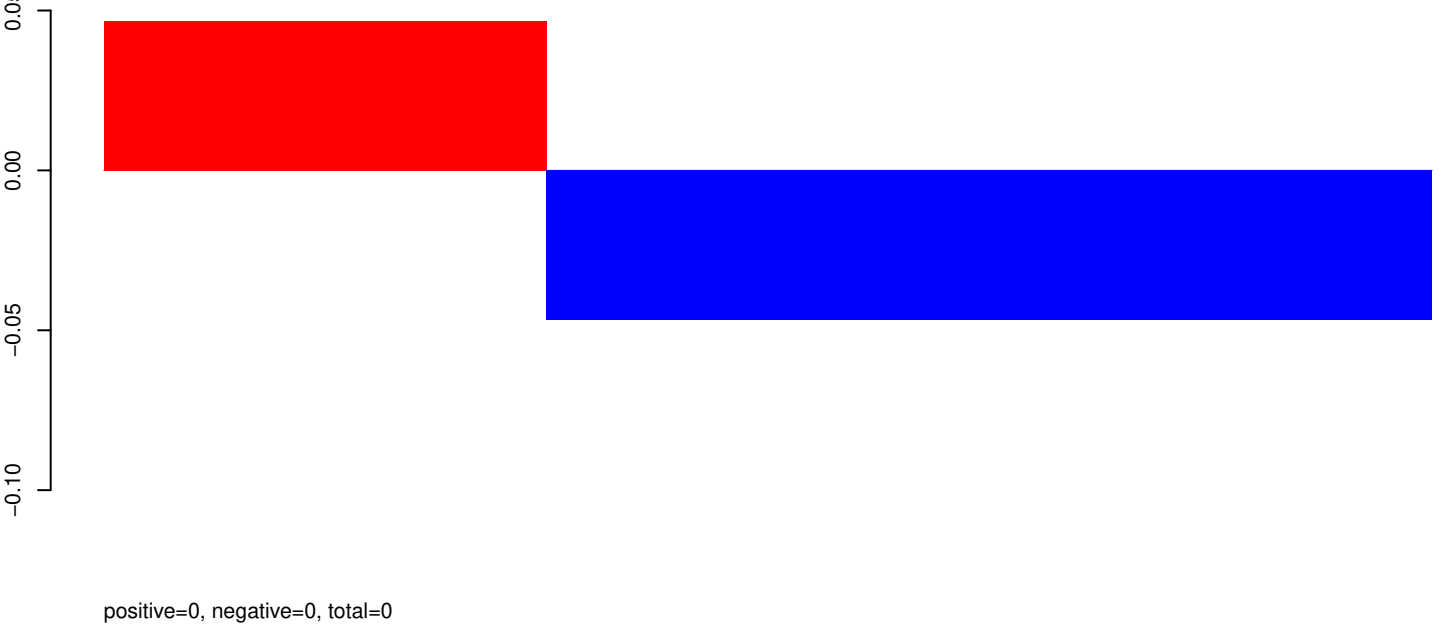
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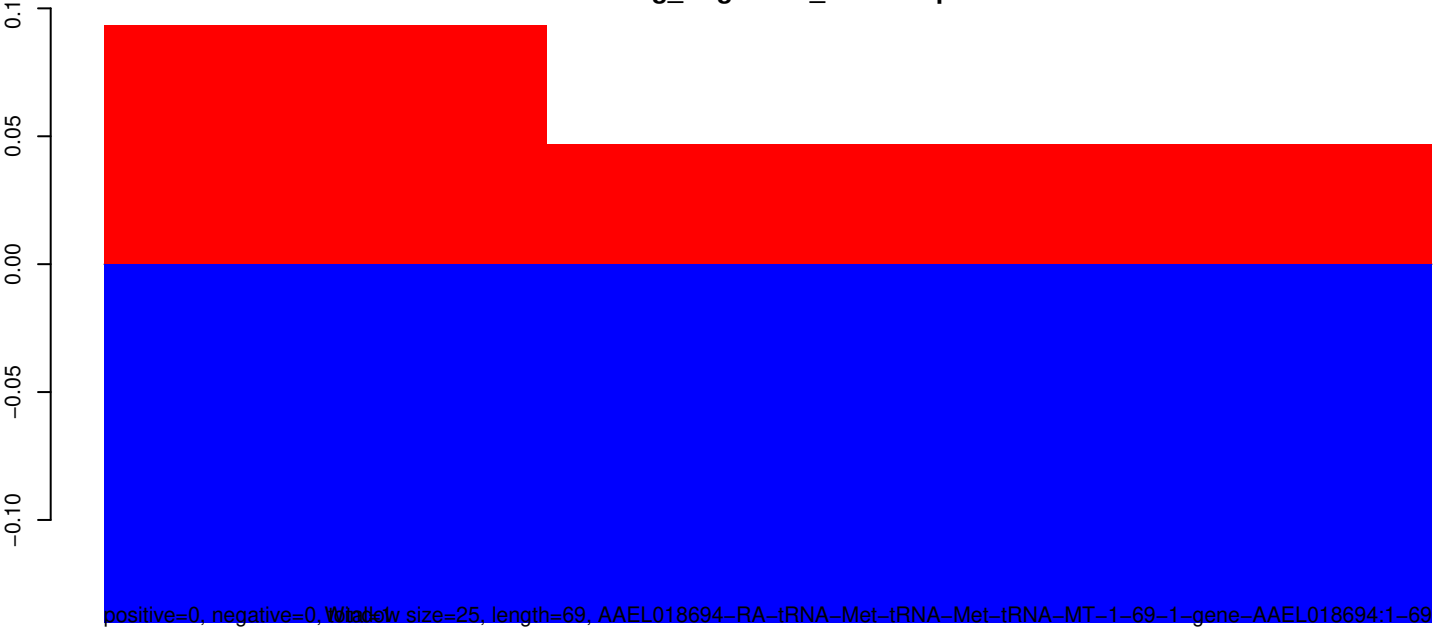
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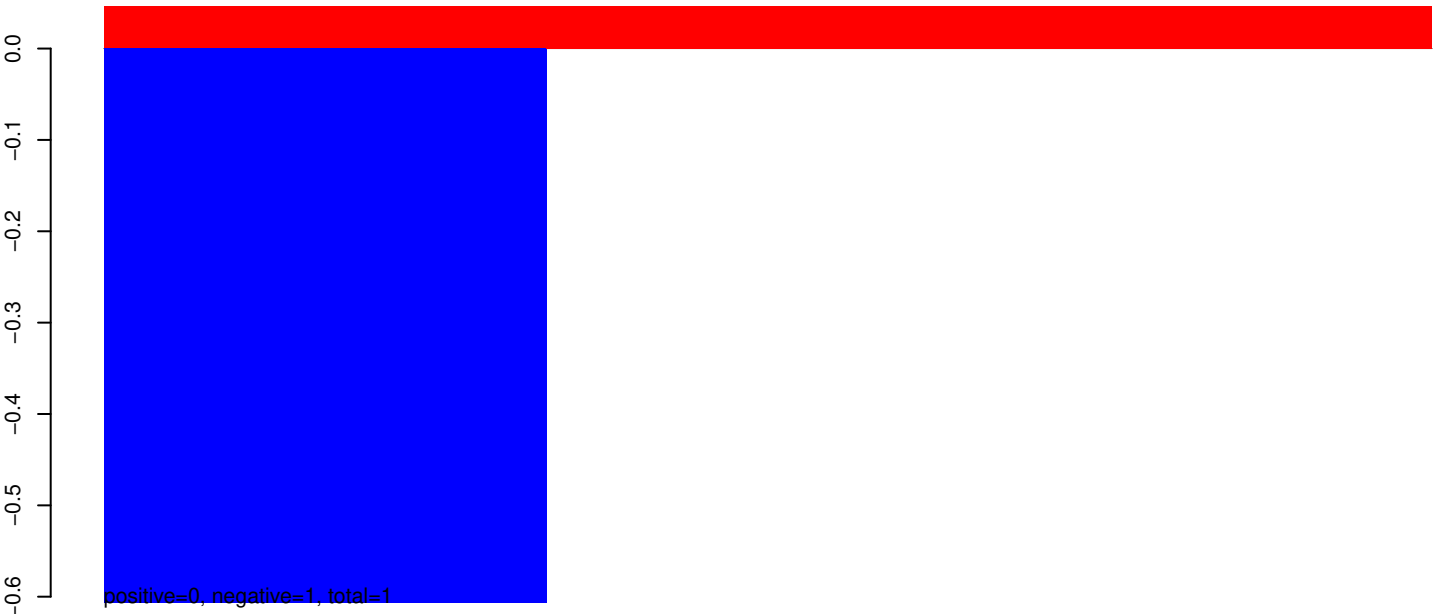
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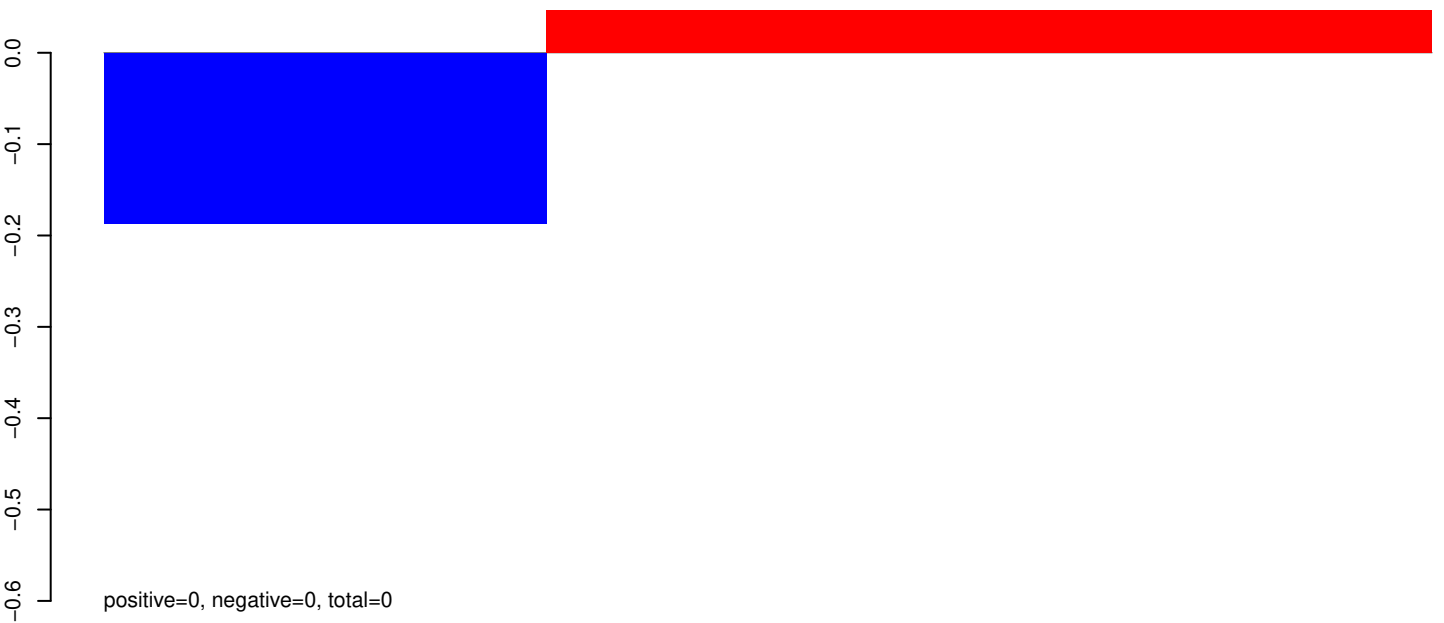
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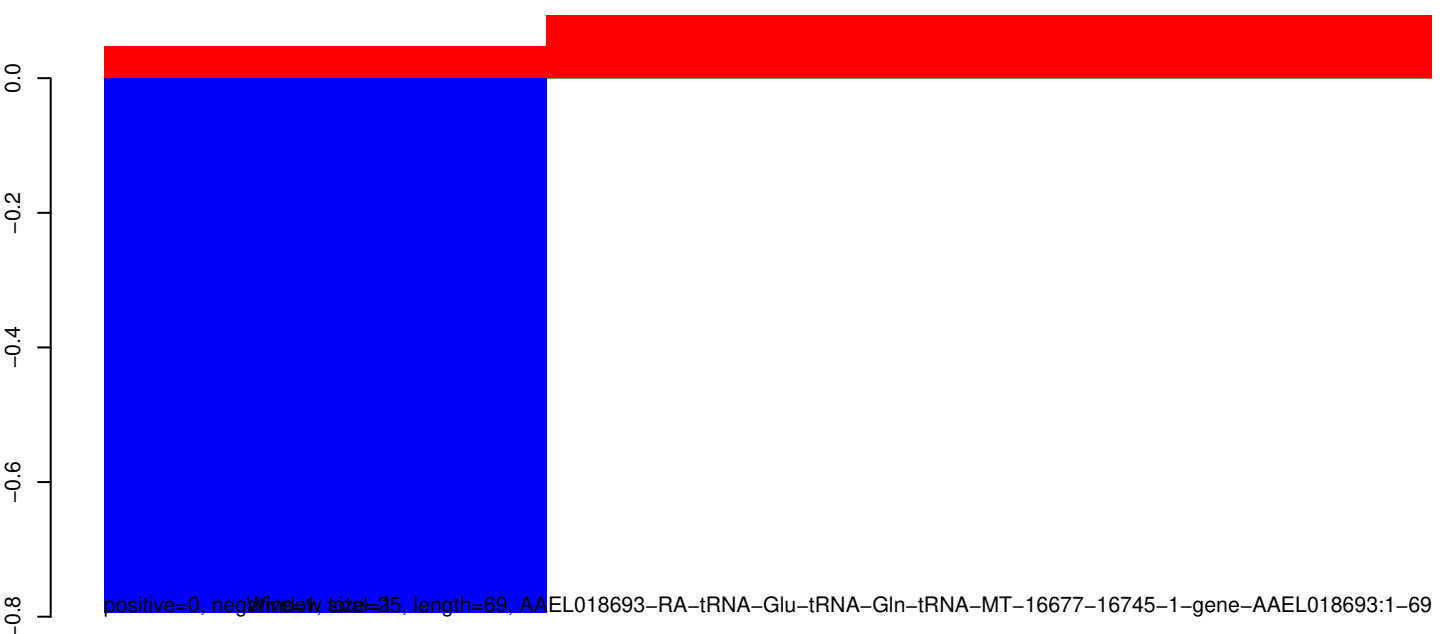
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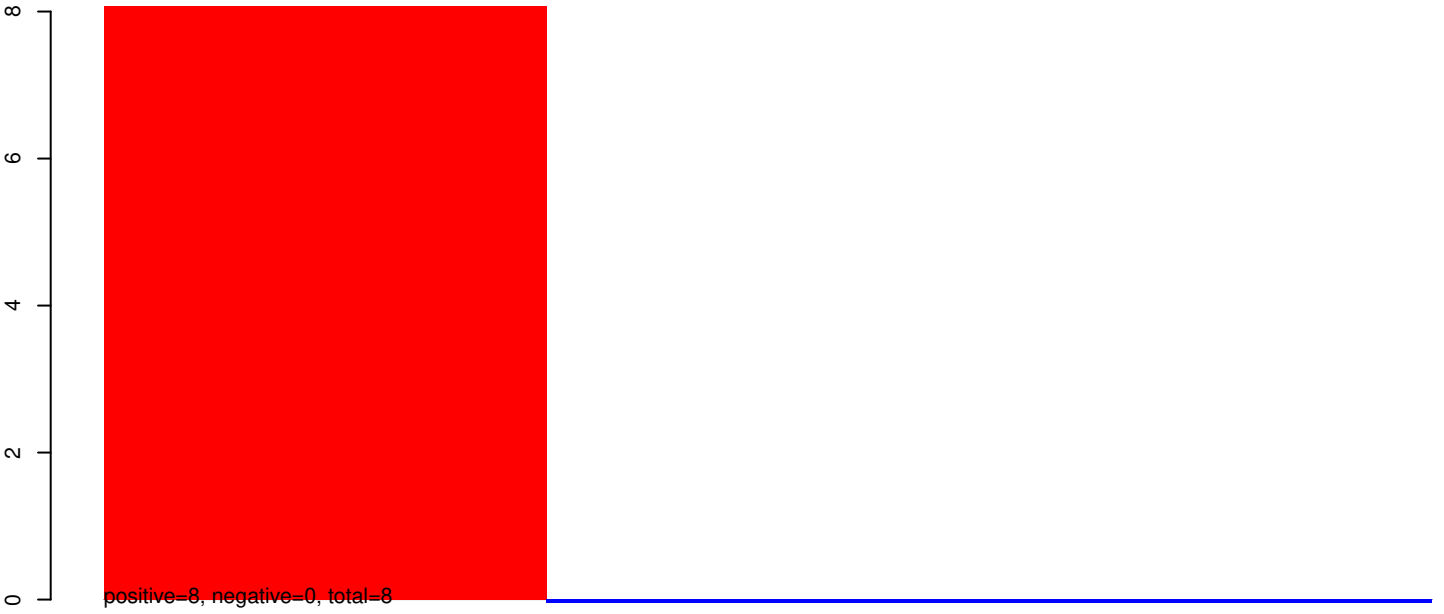
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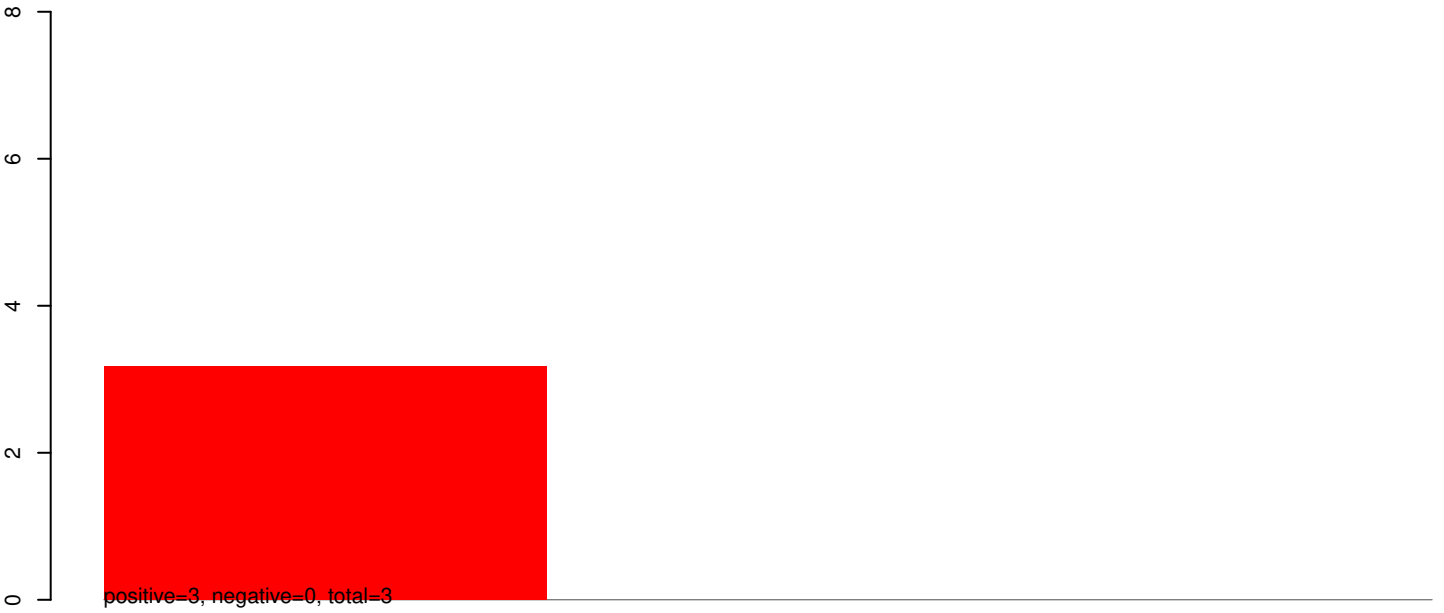
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AeAeg_Aag2SINV_dsLuc.18_23.rep



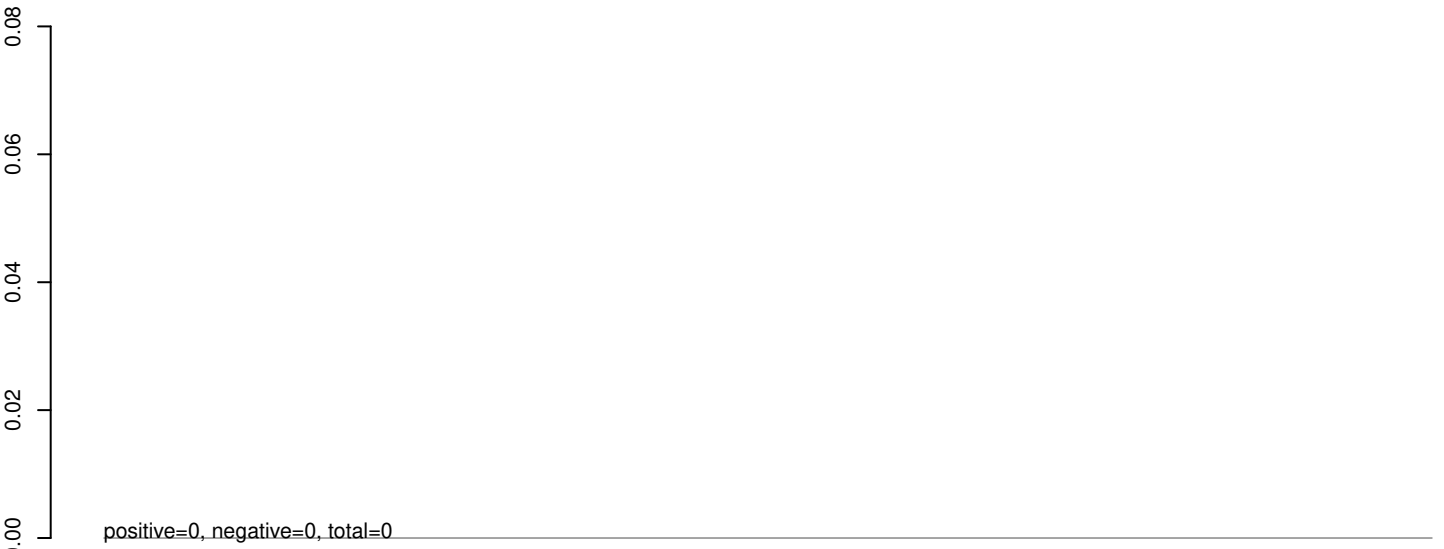
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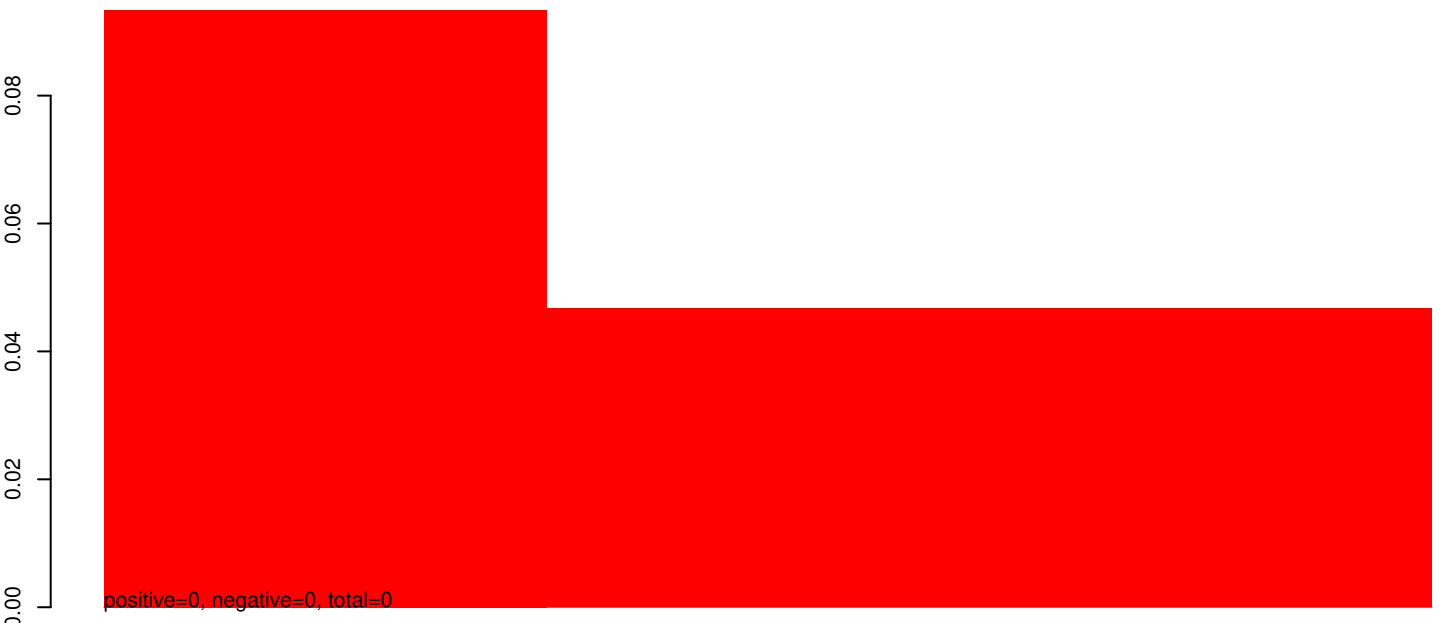
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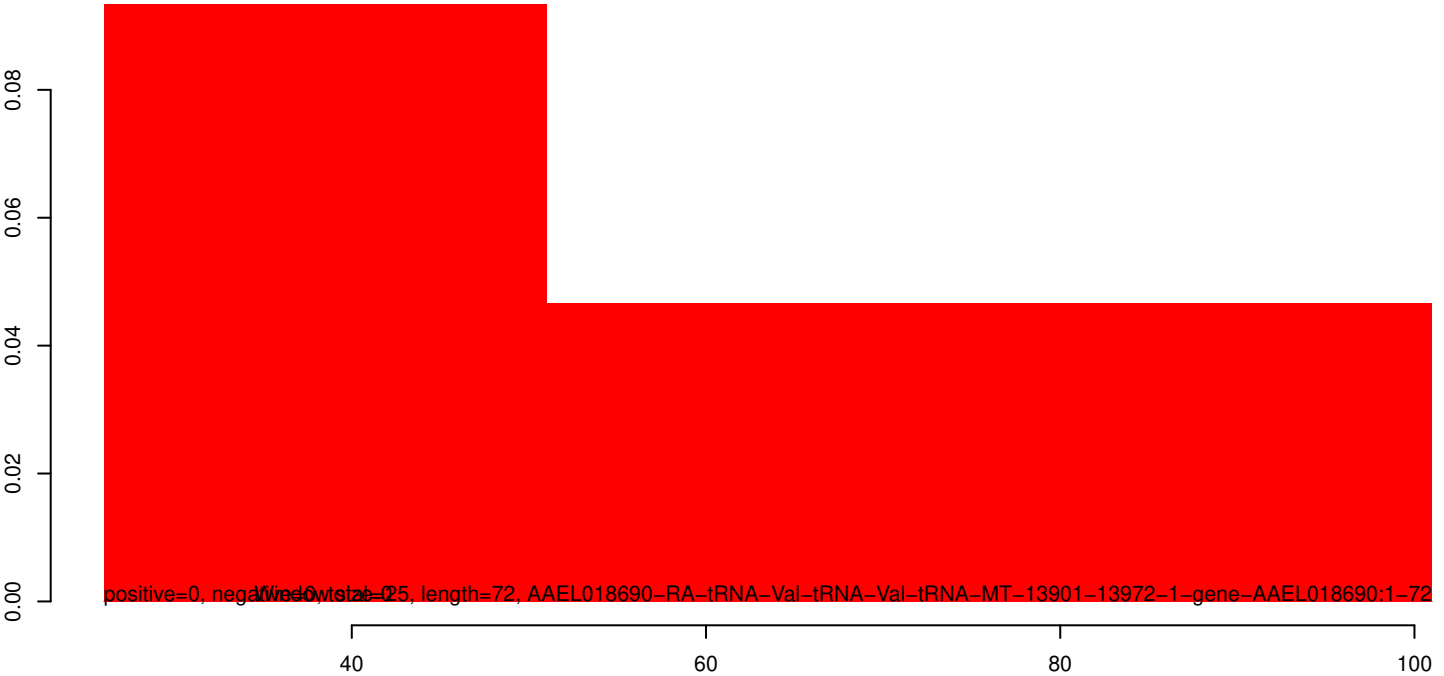
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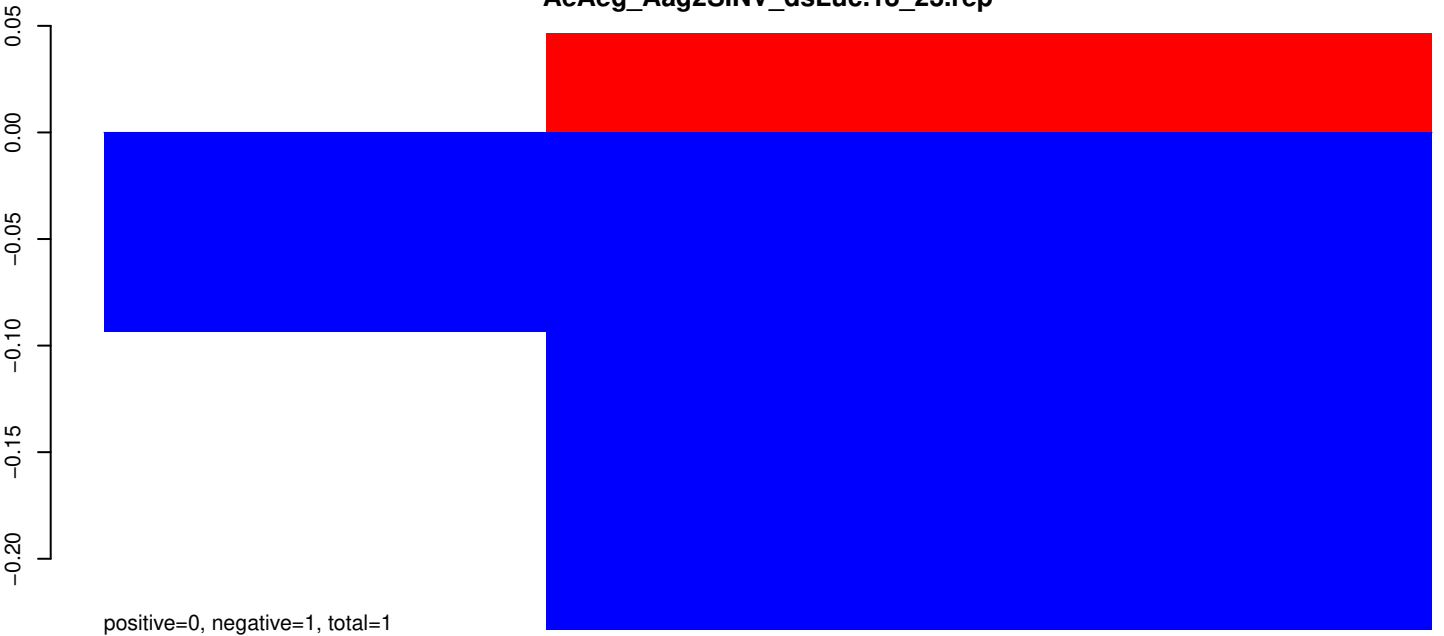
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AeAeg_Aag2SINV_dsLuc.rep



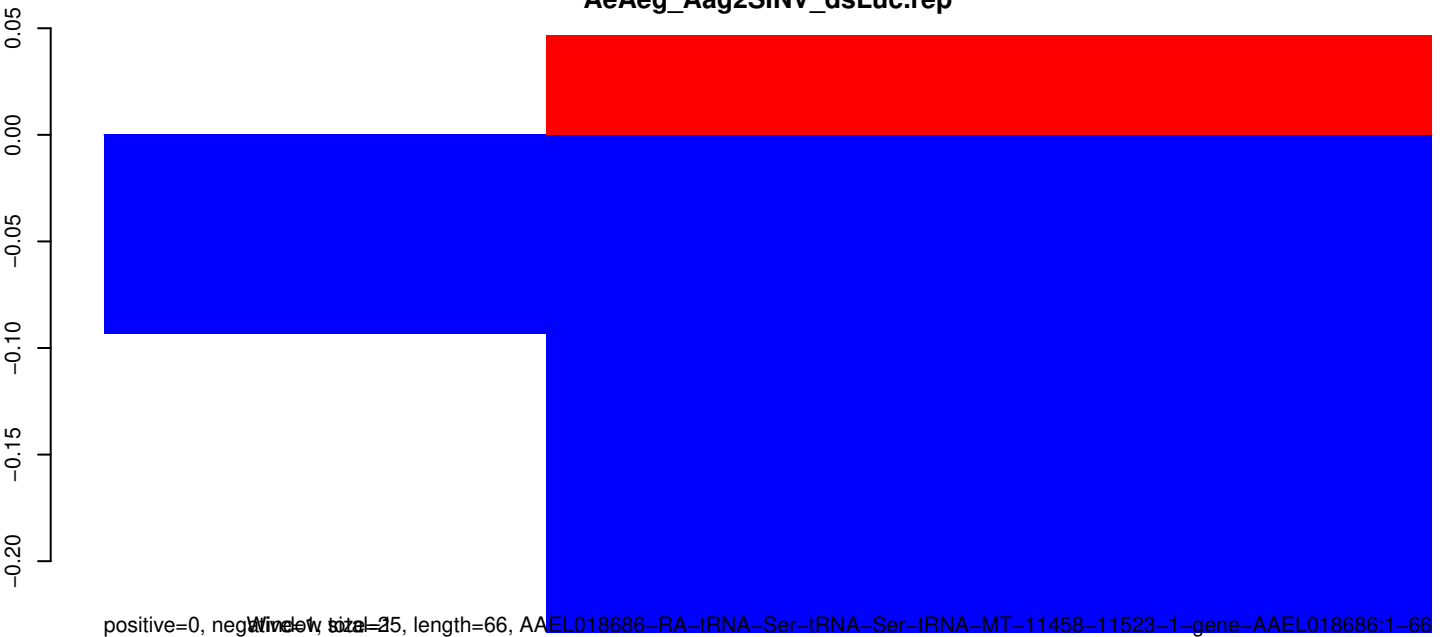
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AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



Window size=25, length=66, AAEL018686-RA-tRNA-Ser-tRNA-Ser-tRNA-MT-11458-11523-1-gene-AAEL018686:1-66

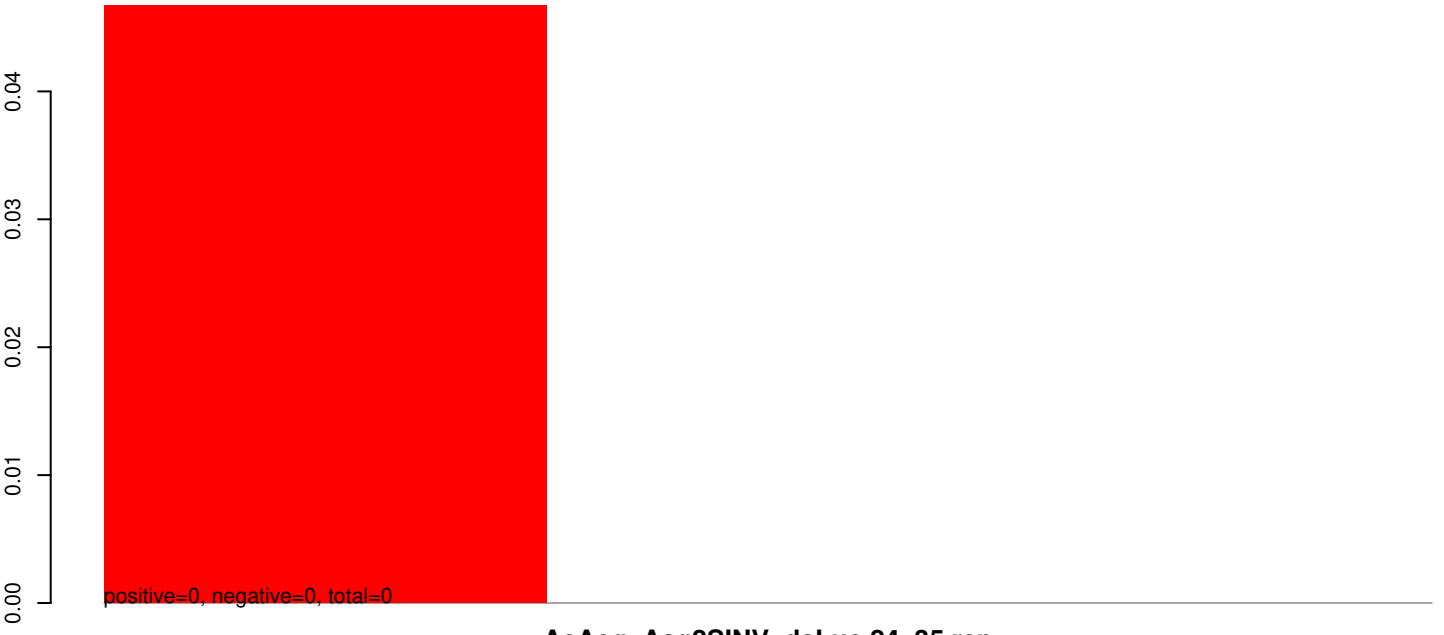
40

60

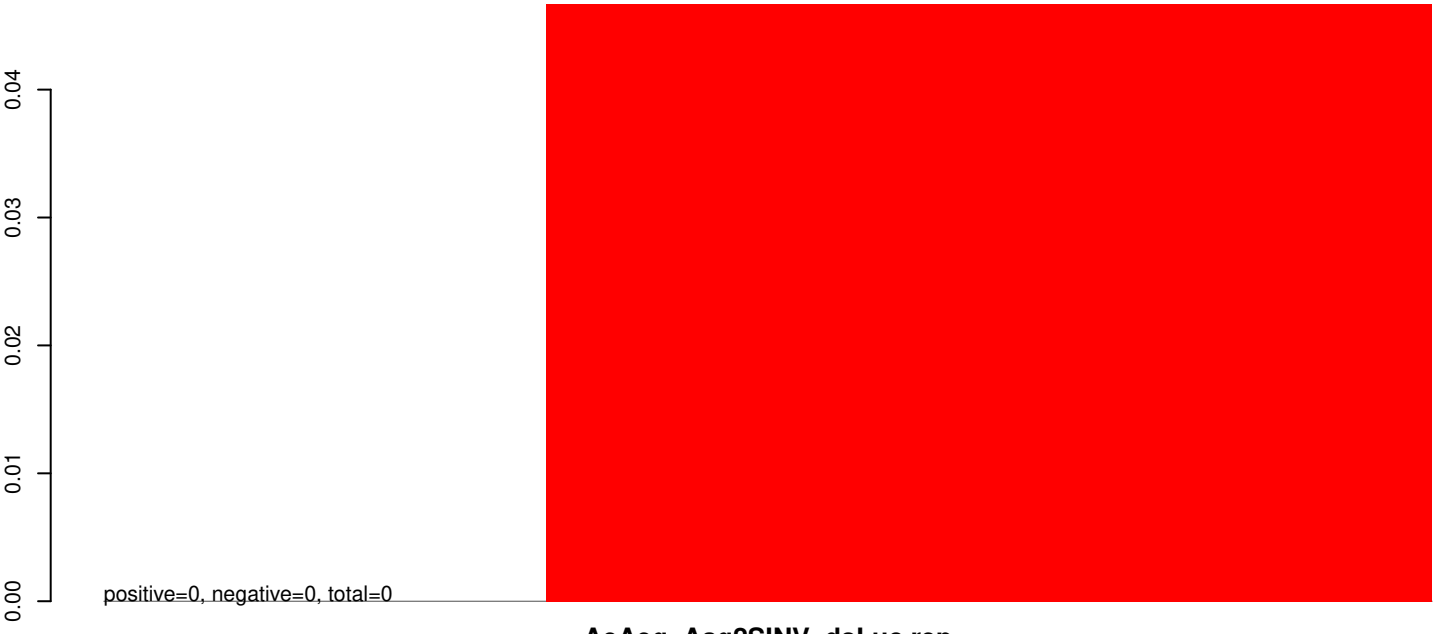
80

100

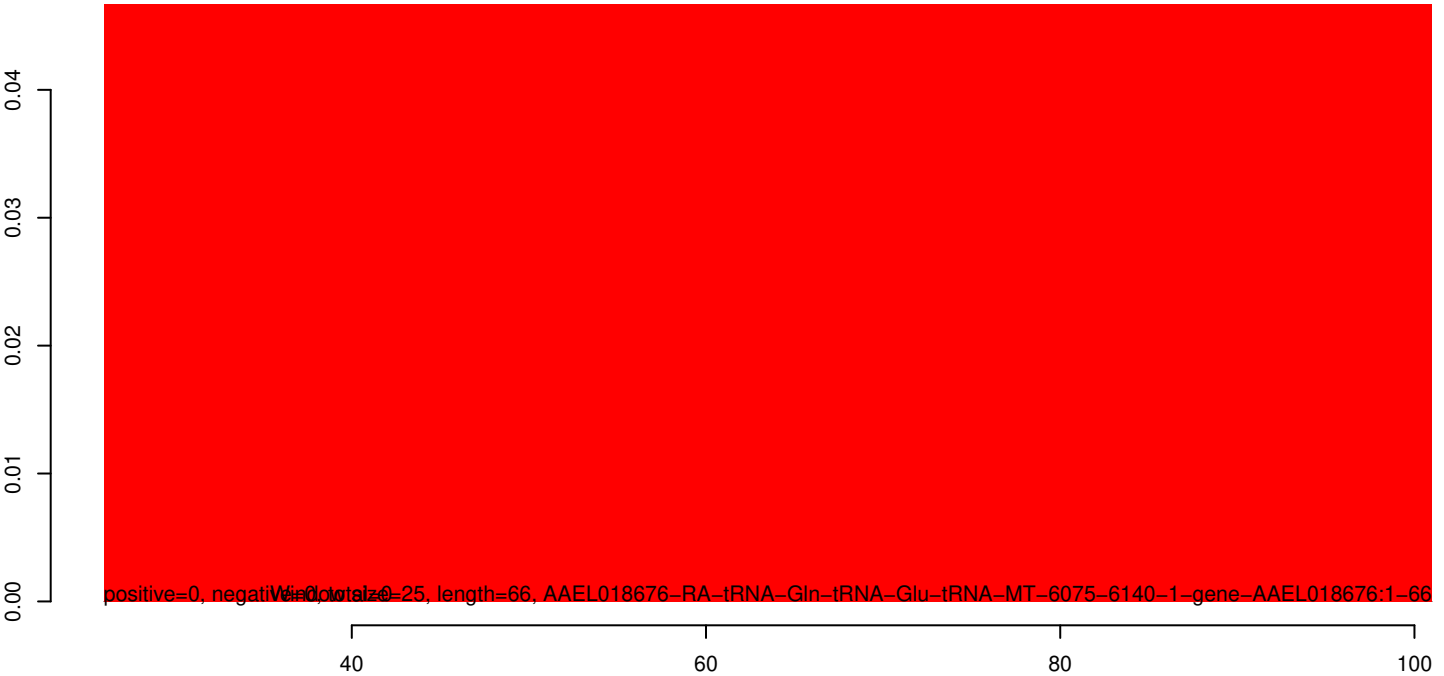
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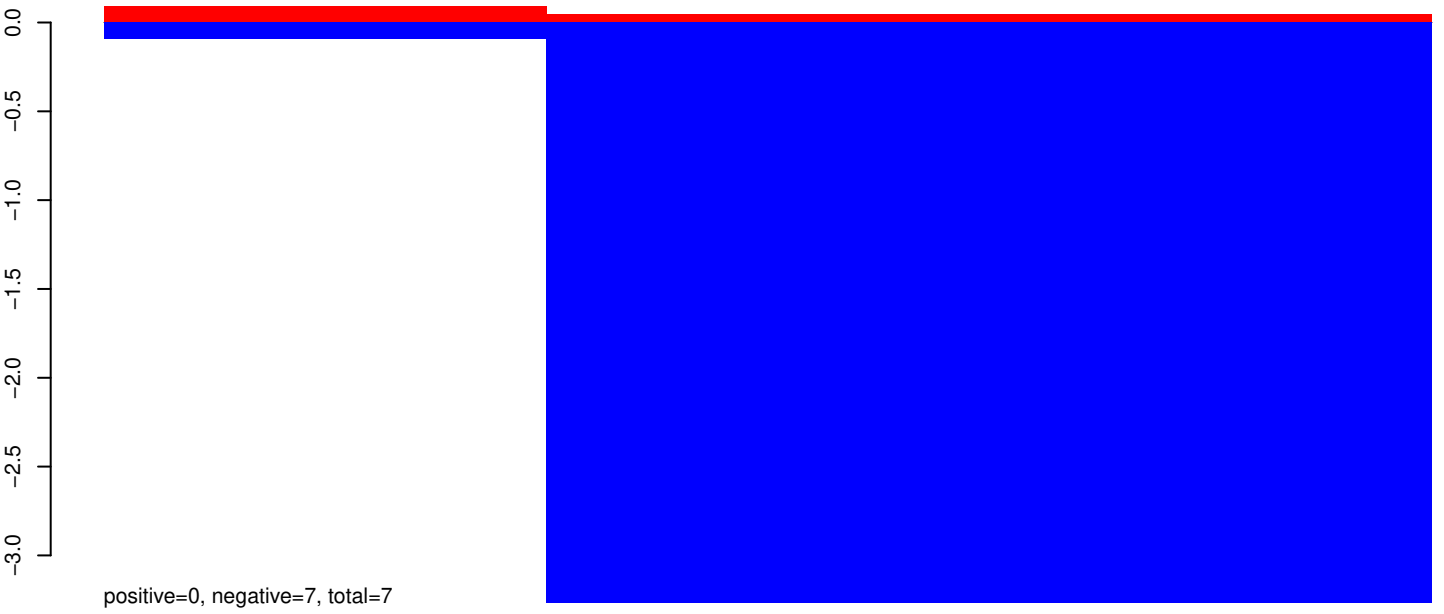
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AeAeg_Aag2SINV_dsLuc.rep

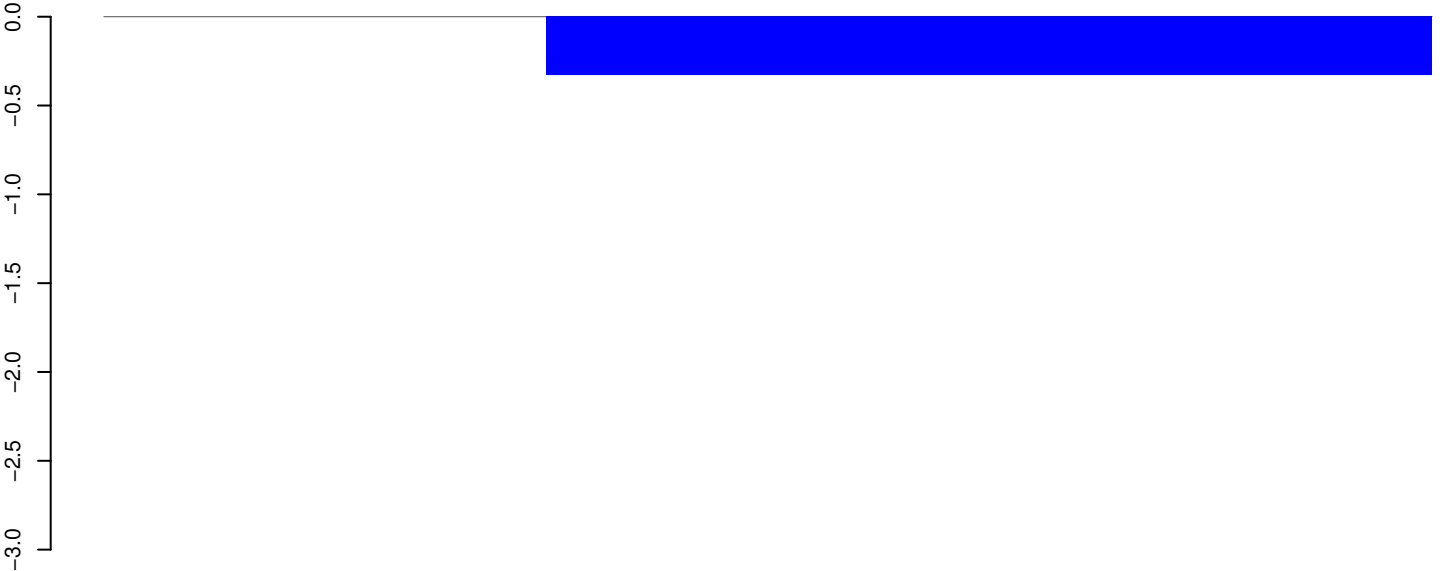


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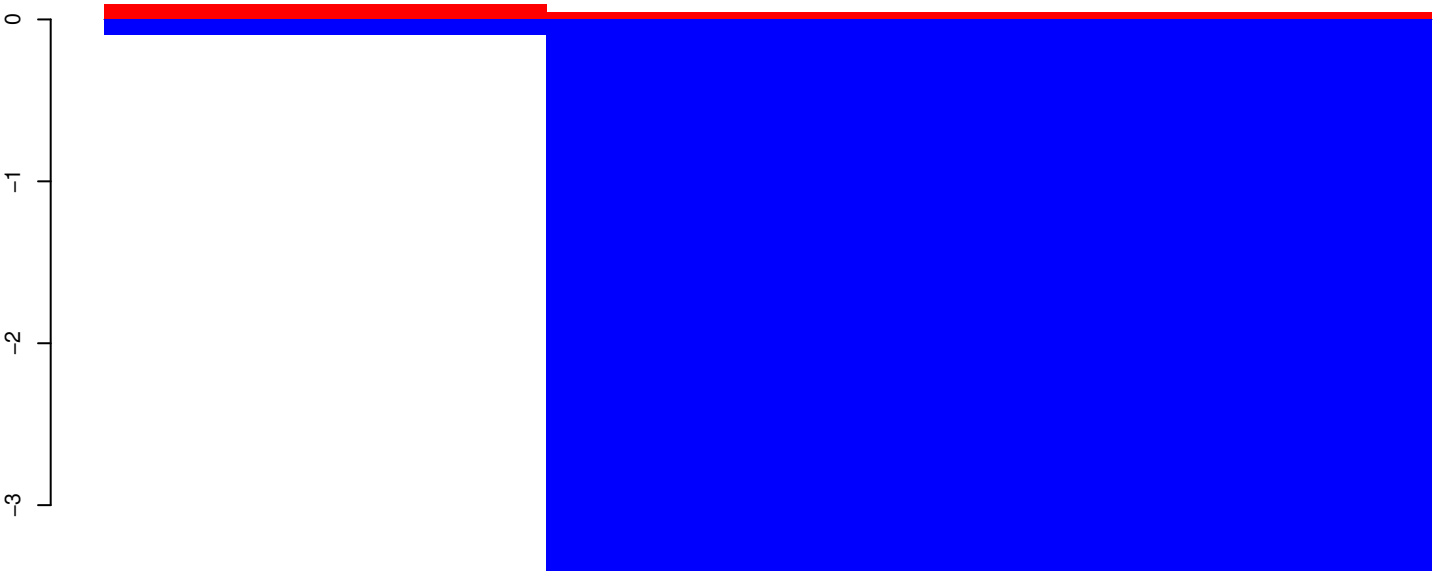
positive=0, negative=7, total=7

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positive=0, negative=1, total=1

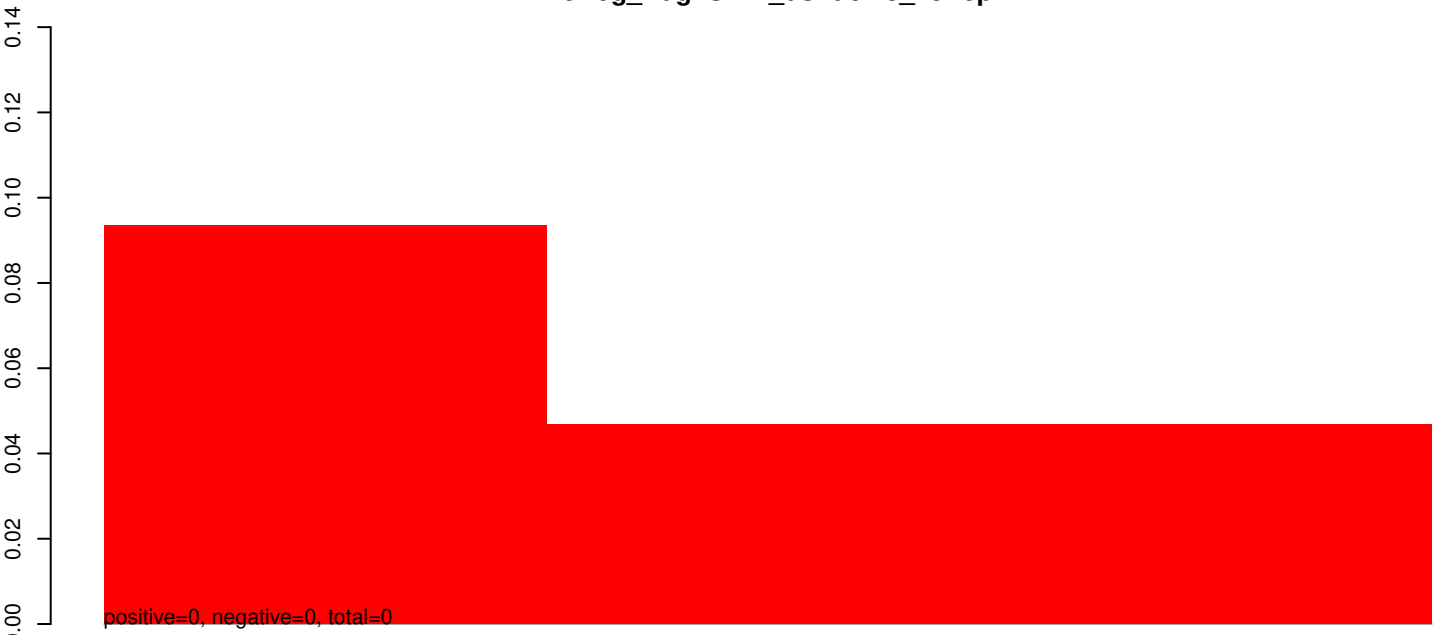
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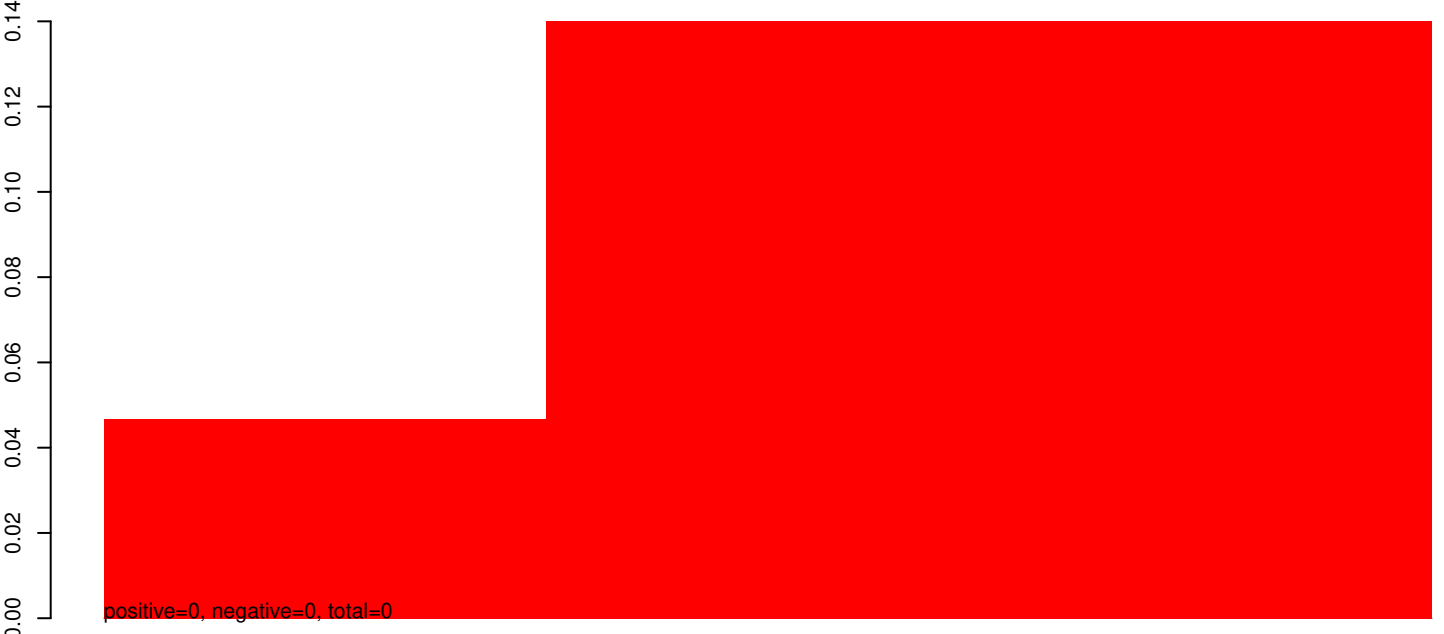
positive=0, negative=25, total=25, length=67, AAEL018675-RA-tRNA-Ser-tRNA-Ser-tRNA-MT-5996-6062-1-gene-AAEL018675:1-67

40 60 80 100

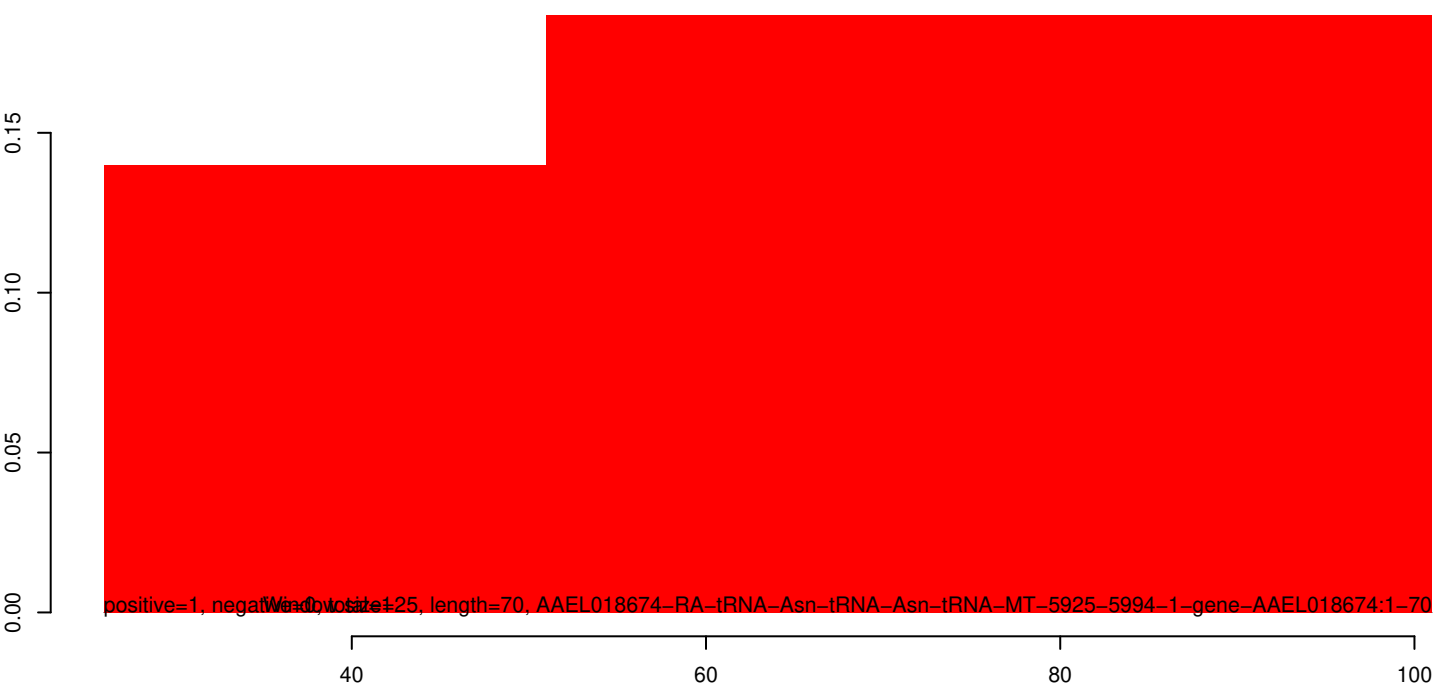
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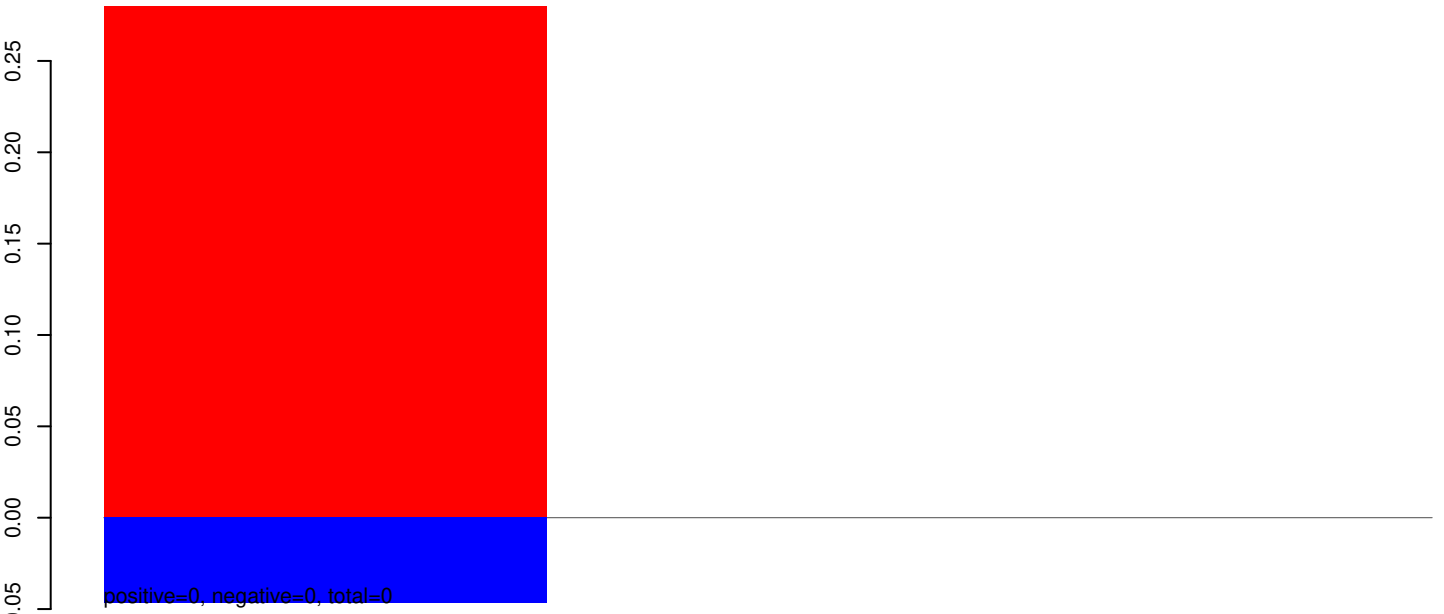
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AeAeg_Aag2SINV_dsLuc.rep



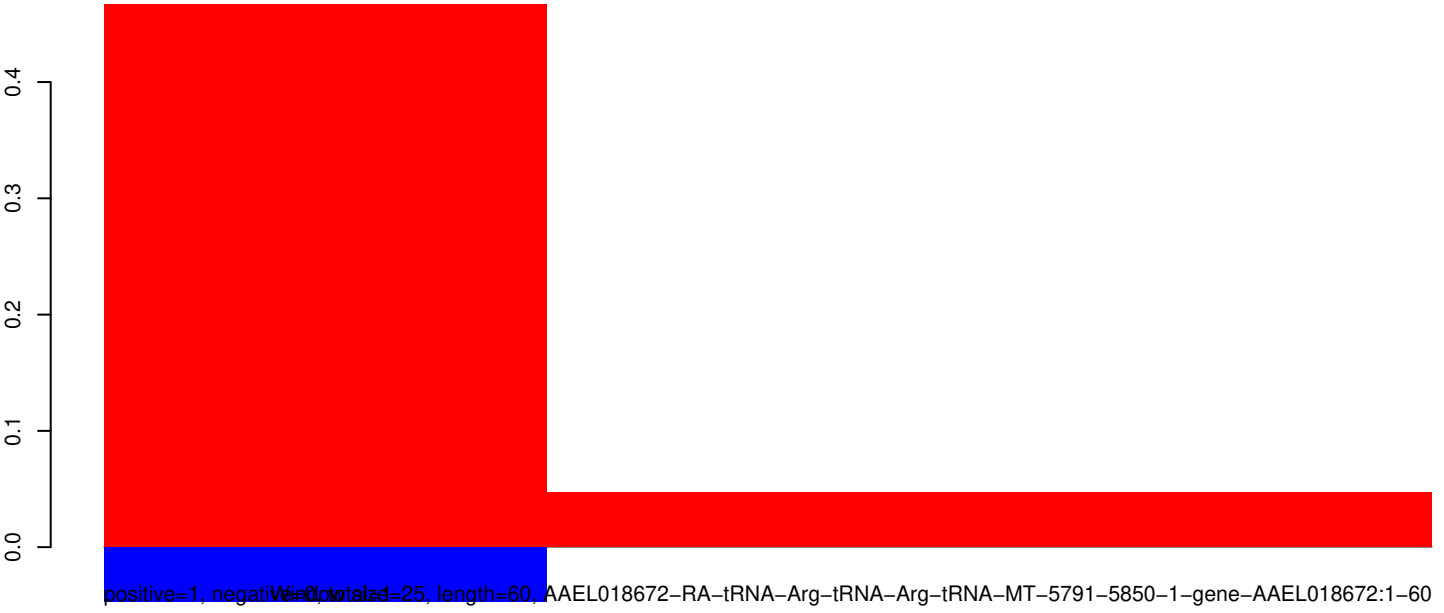
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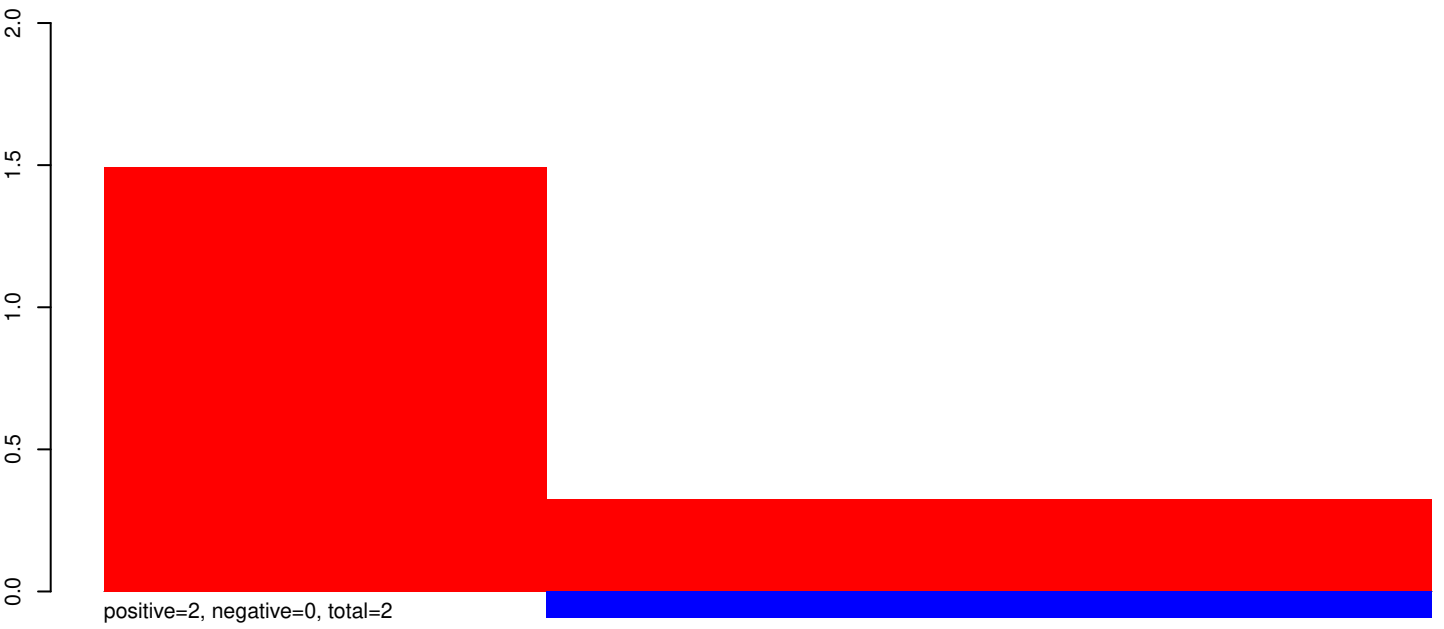
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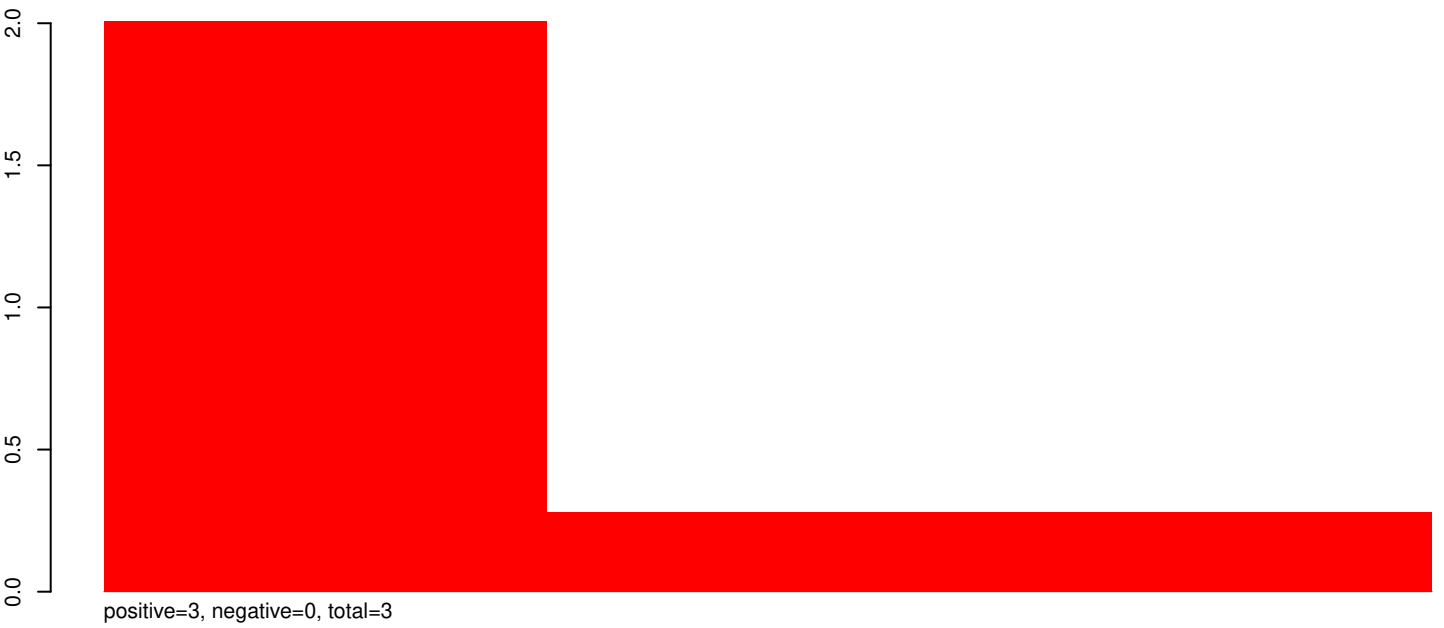
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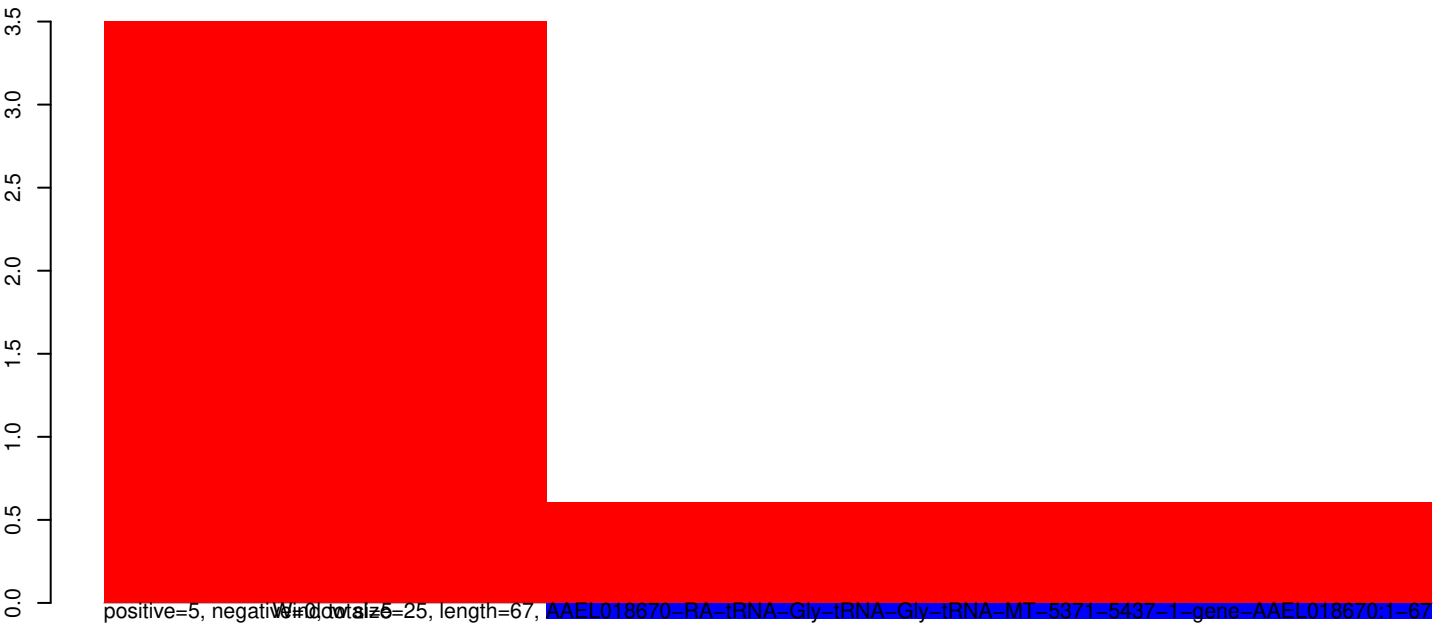
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AeAeg_Aag2SINV_dsLuc.24_35.rep



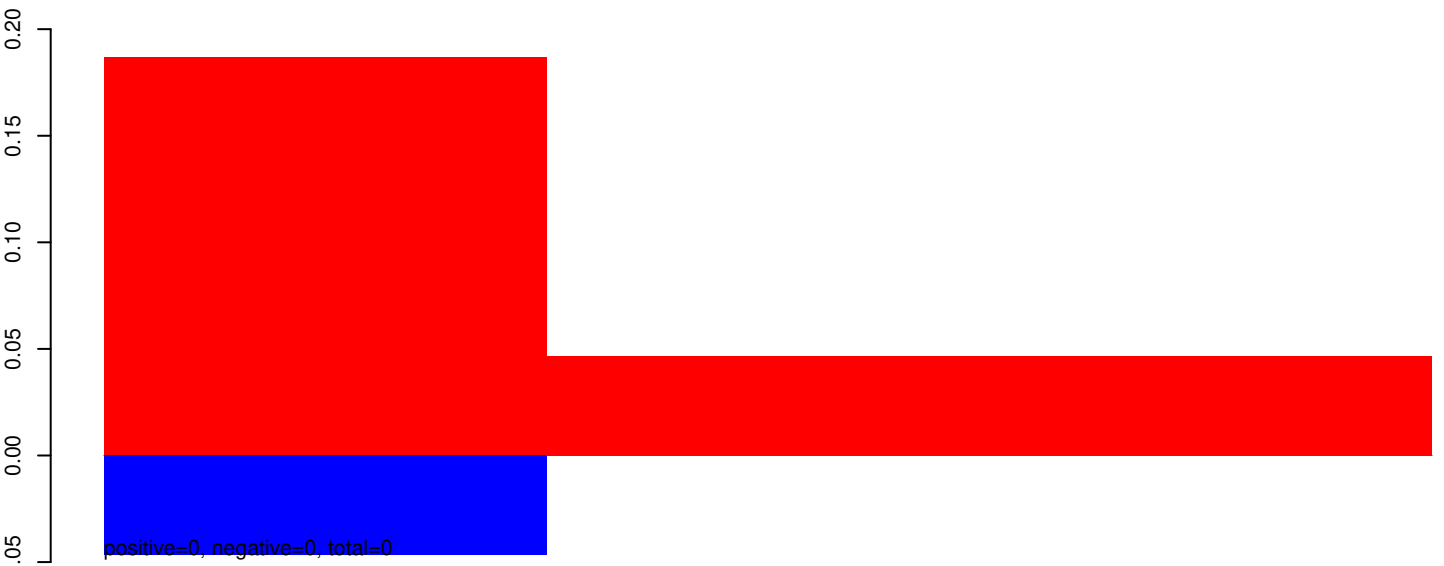
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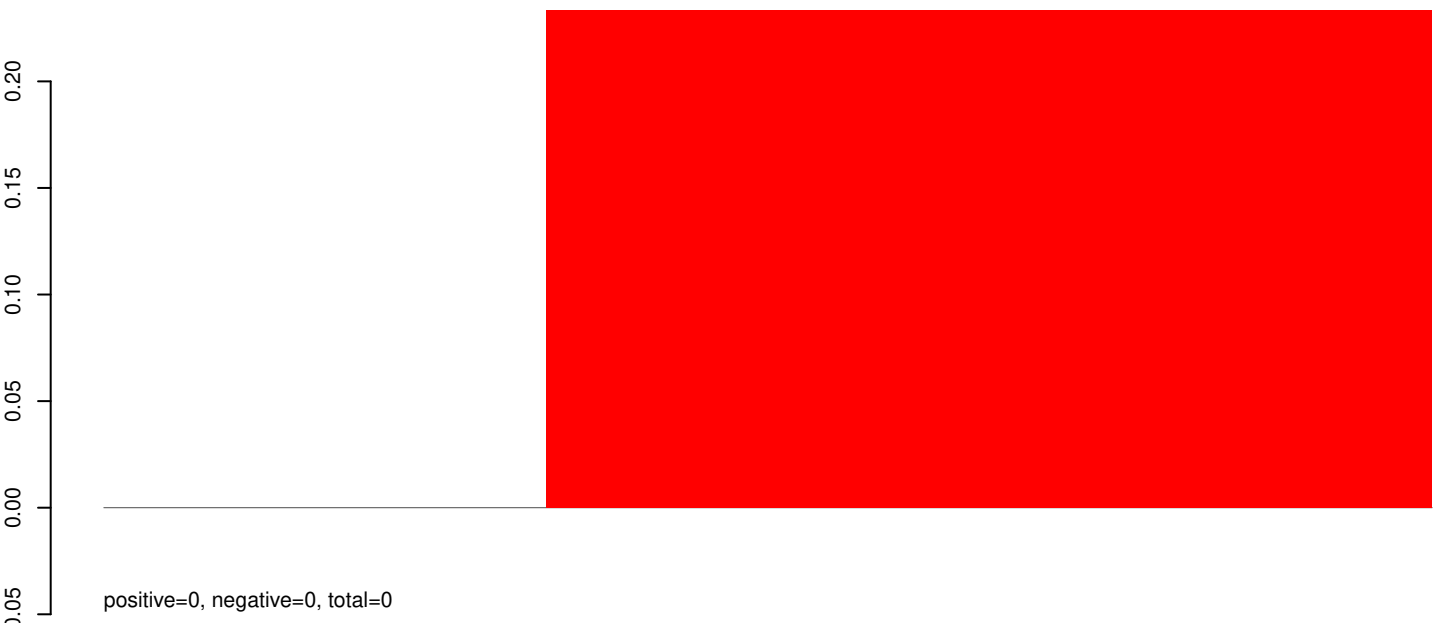
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40 60 80 100

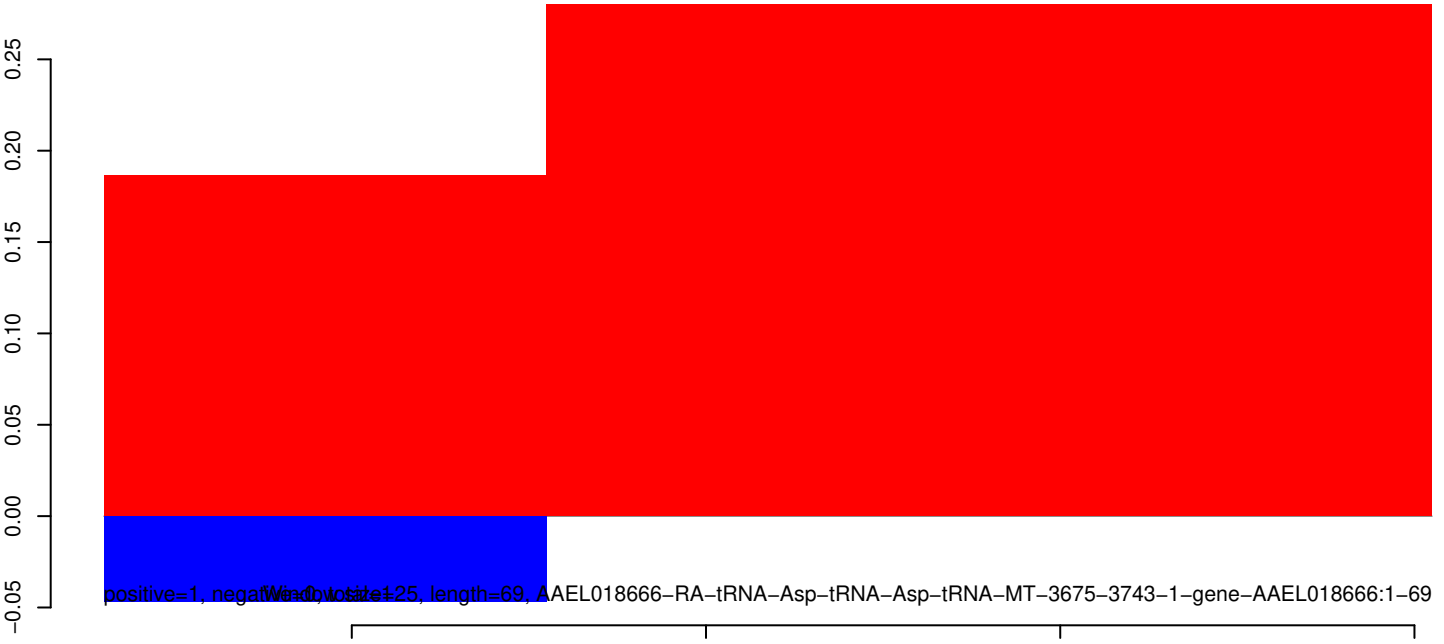
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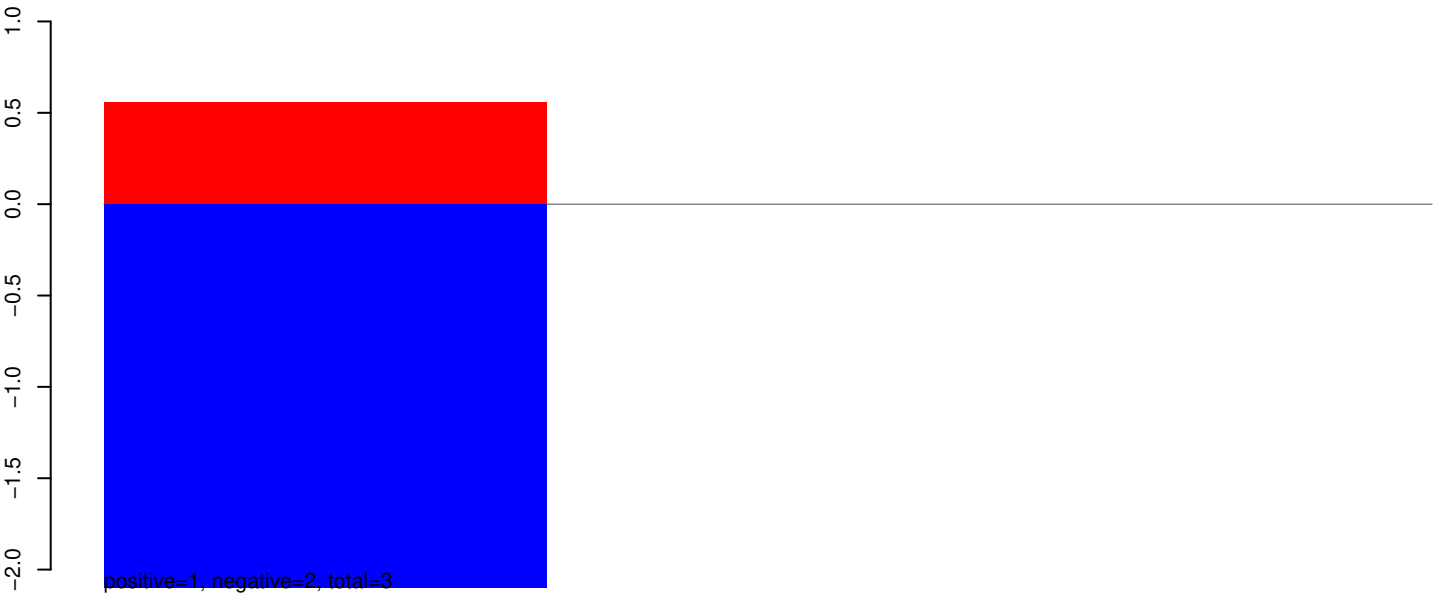
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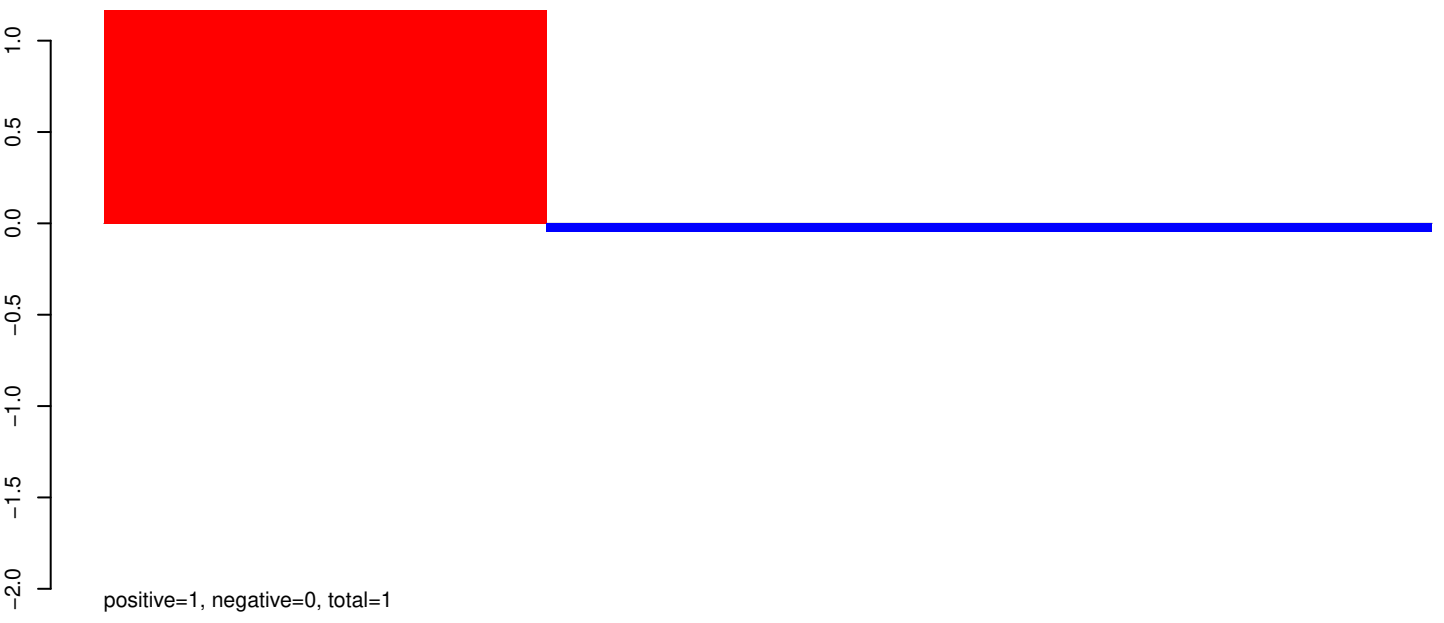
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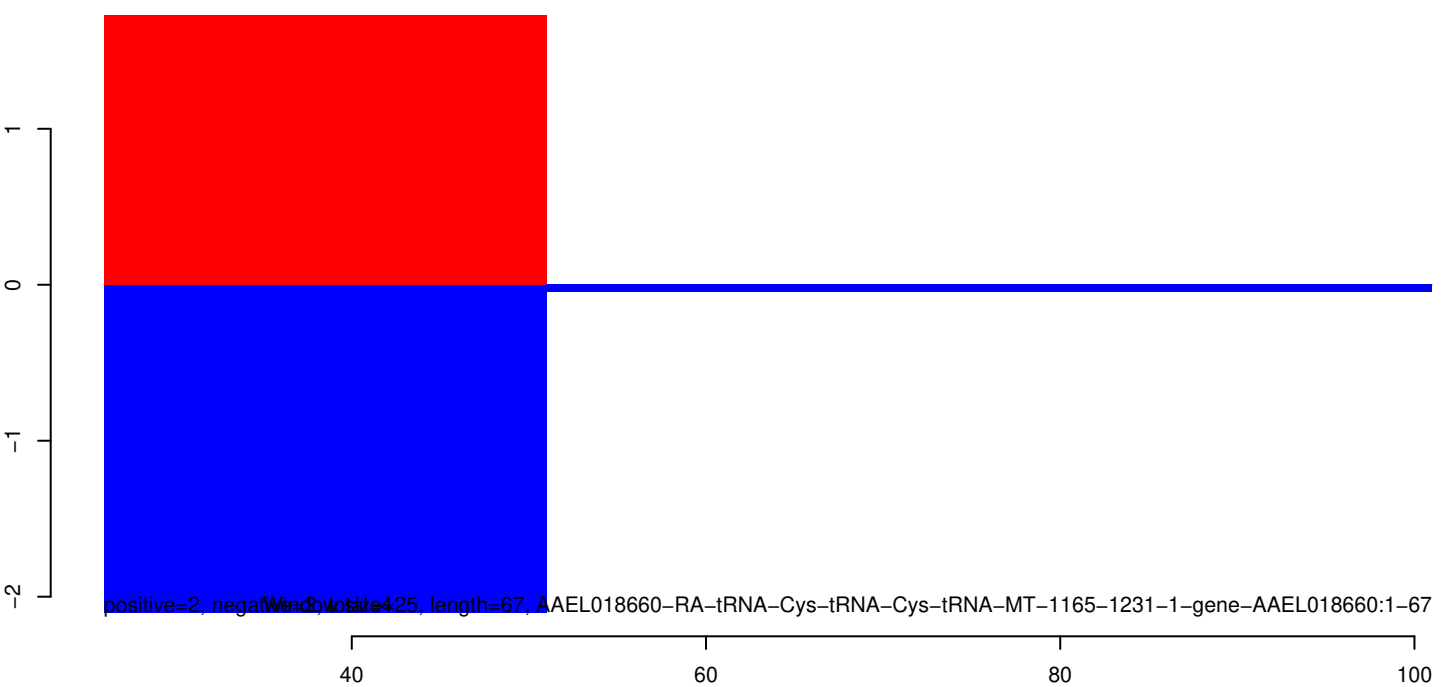
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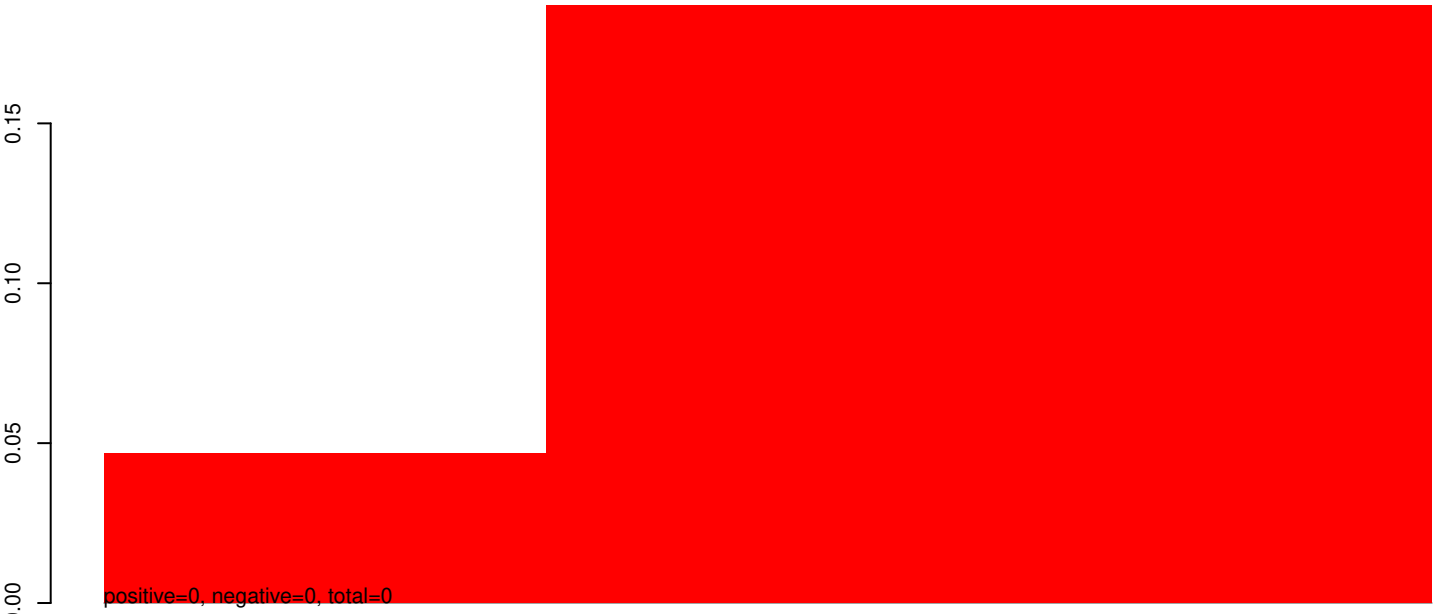
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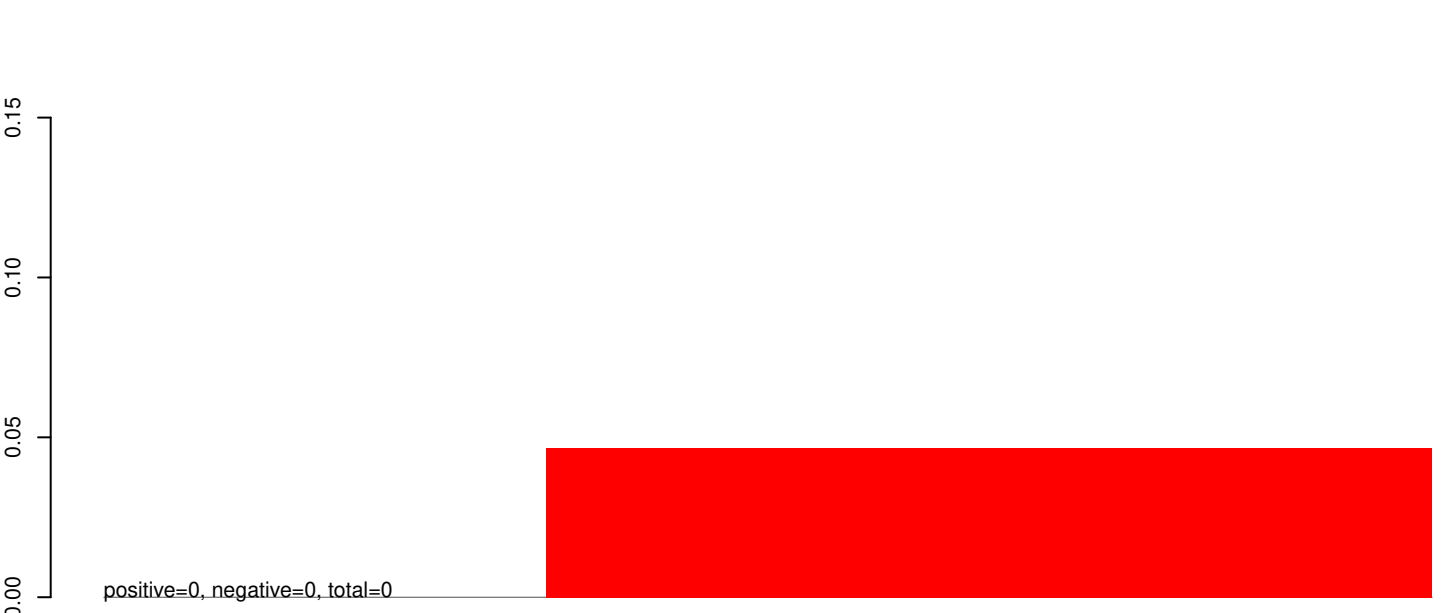
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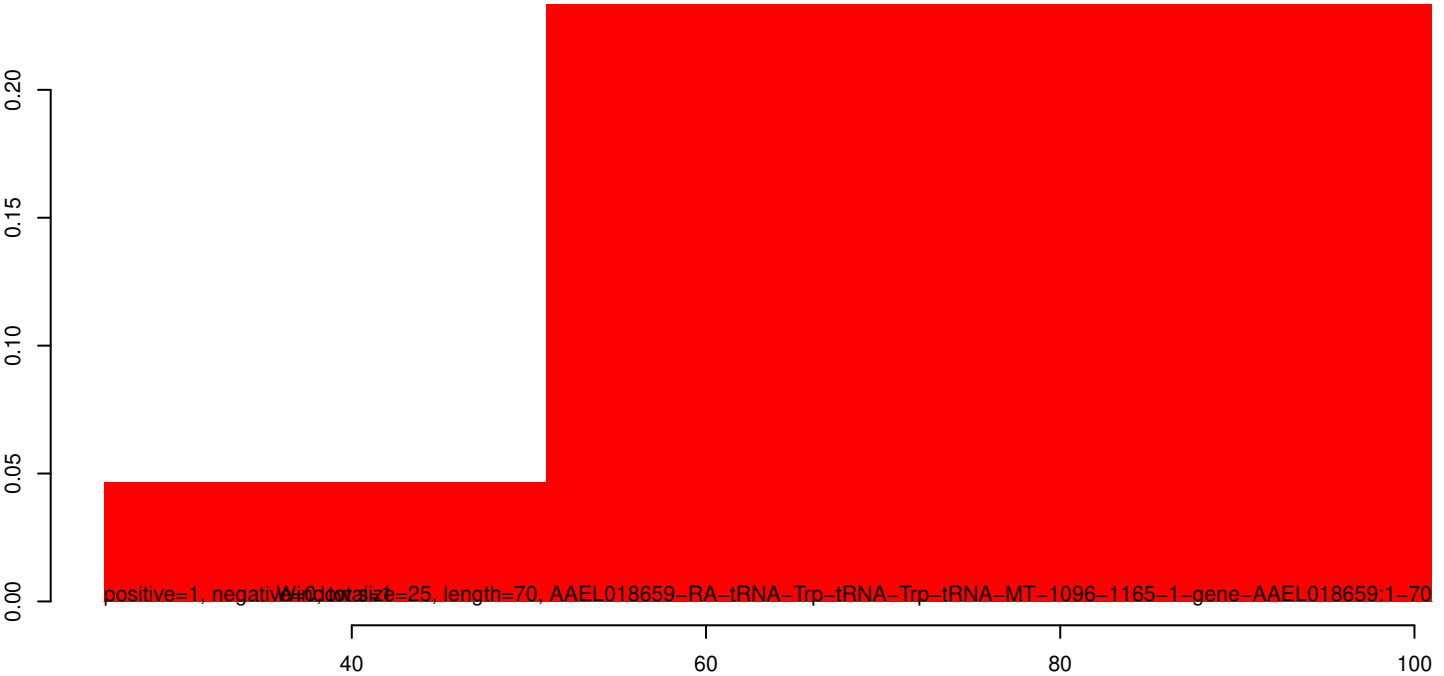
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AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep



AeAeg_Aag2SINV_dsLuc.18_23.rep



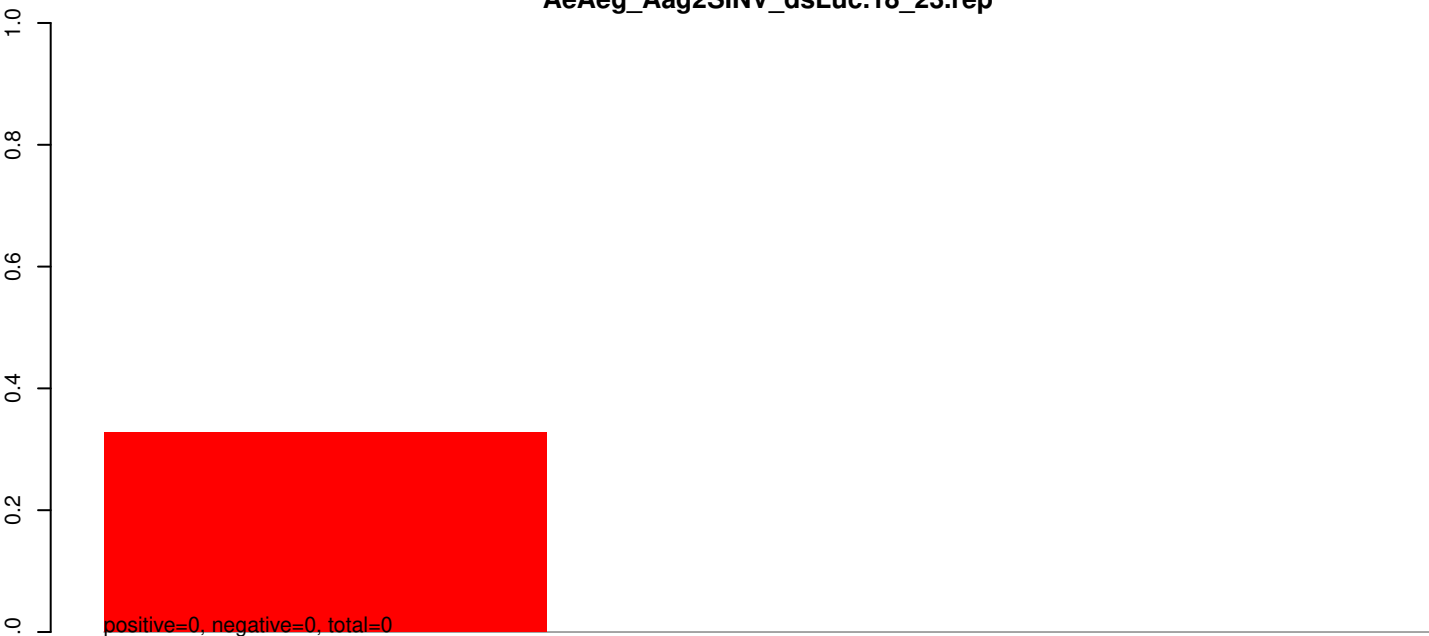
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AeAeg_Aag2SINV_dsLuc.rep



AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep



AeAeg_Aag2SINV_dsLuc.rep

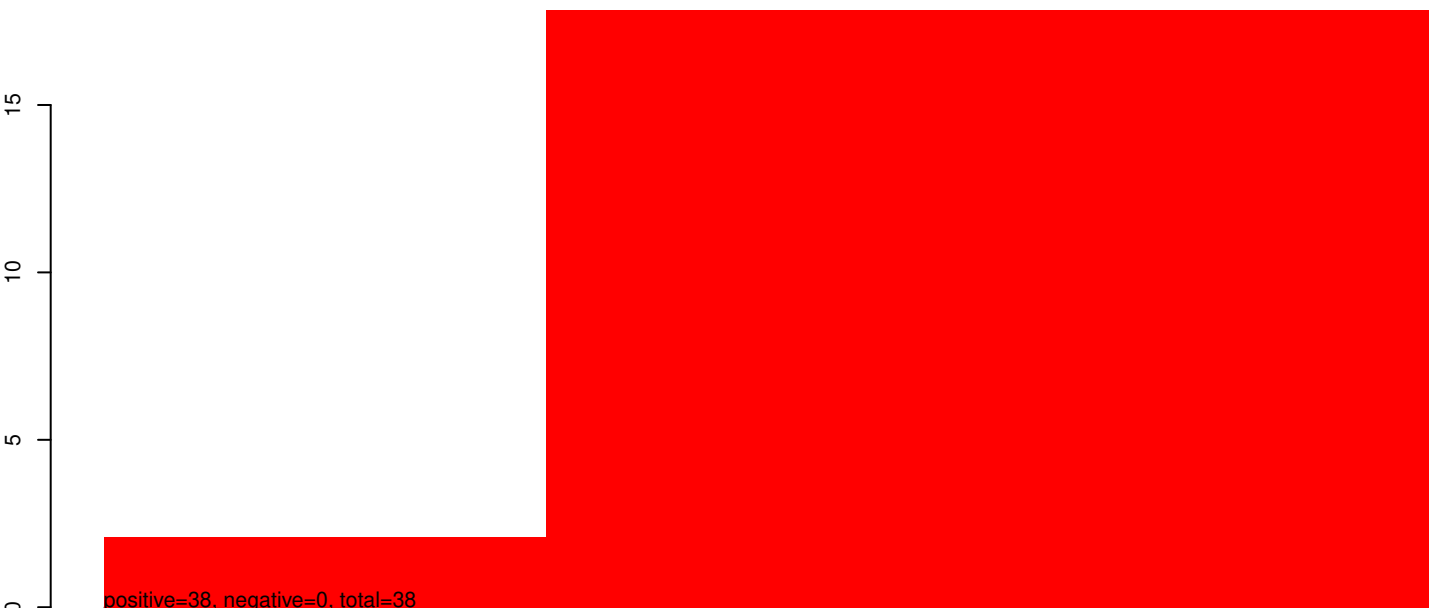


40 60 80 100

AeAeg_Aag2SINV_dsLuc.18_23.rep



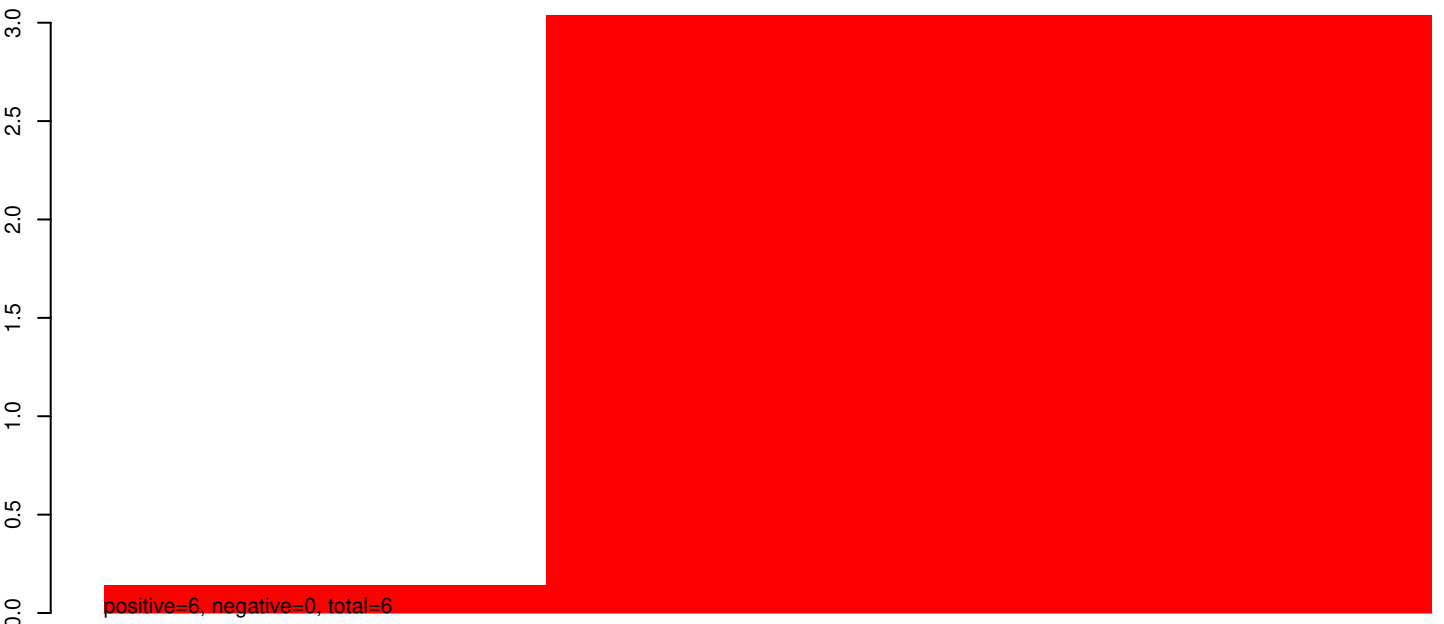
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AeAeg_Aag2SINV_dsLuc.rep



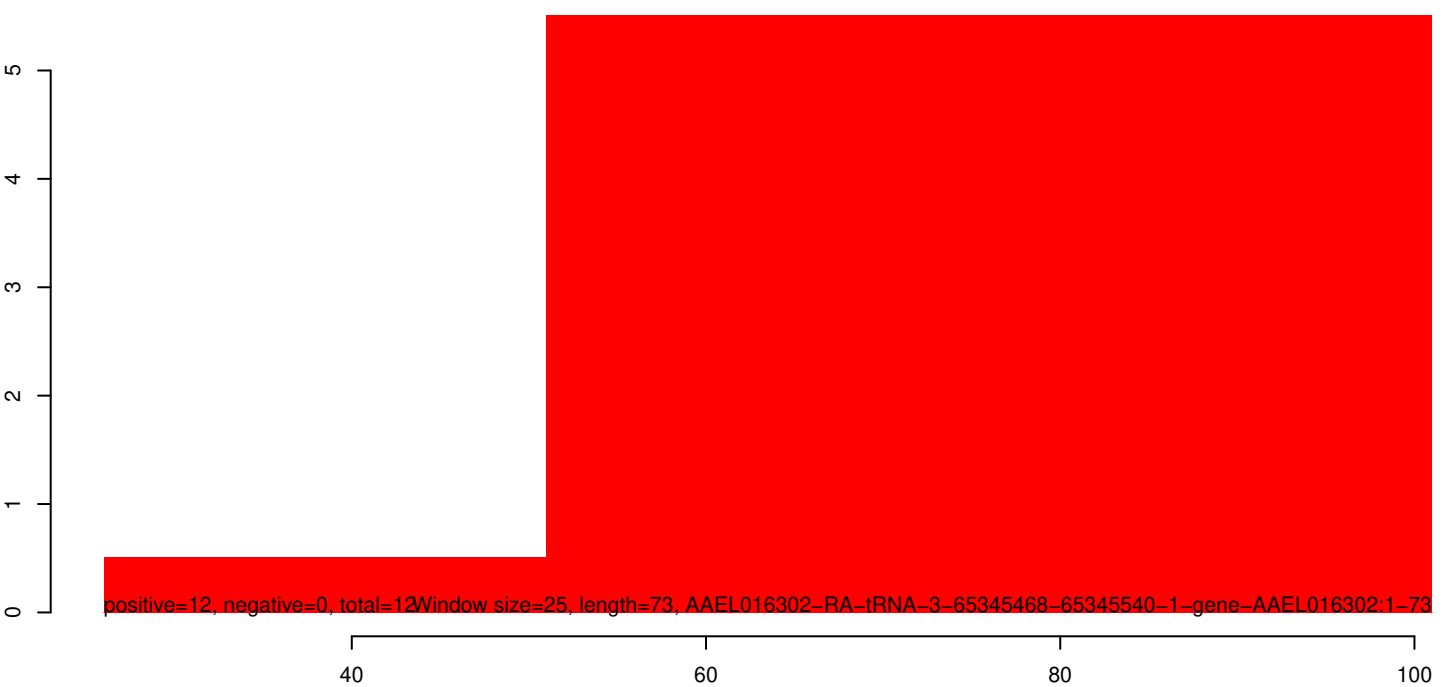
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AeAeg_Aag2SINV_dsLuc.24_35.rep



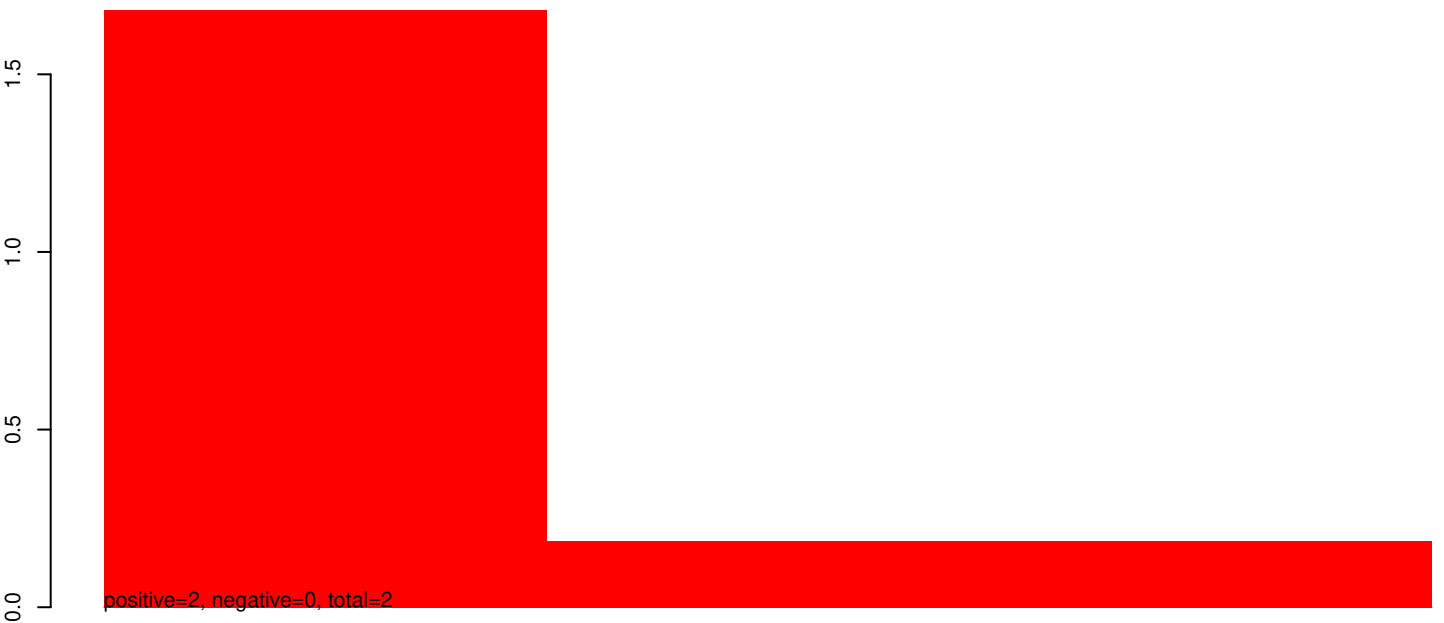
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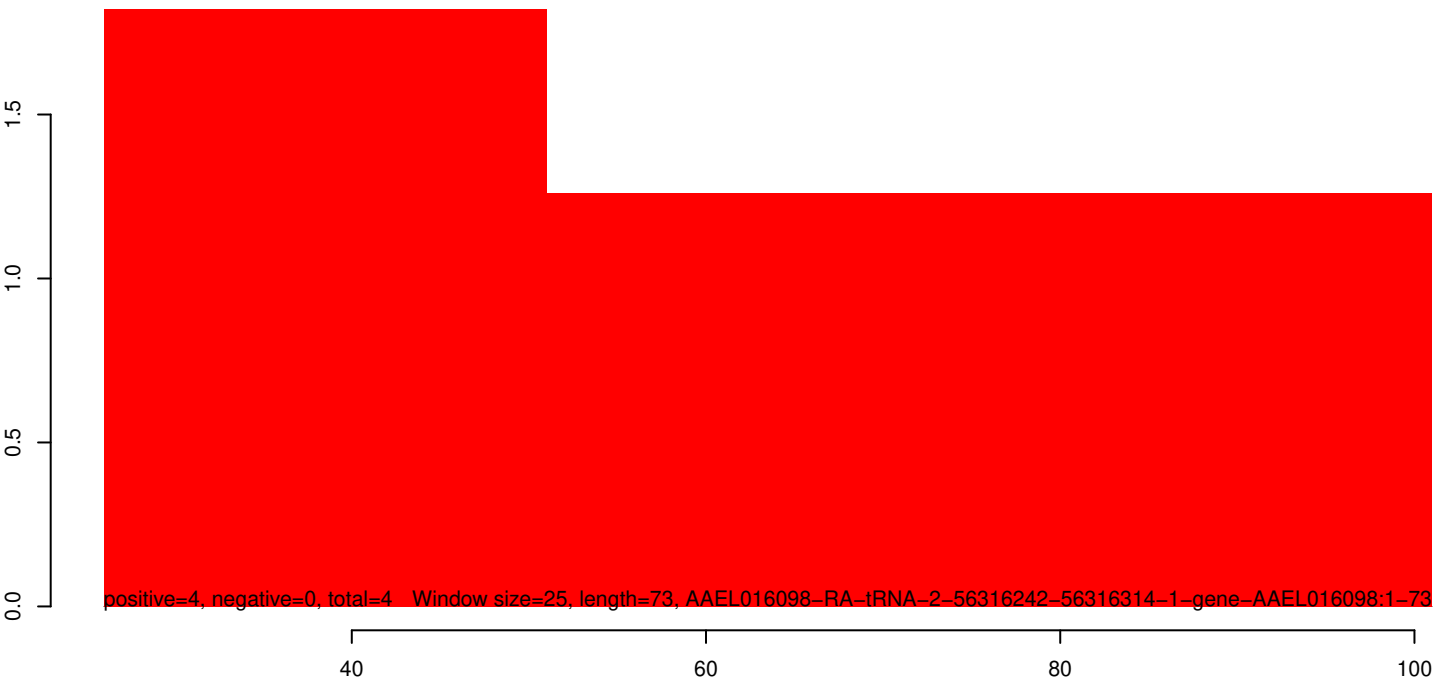
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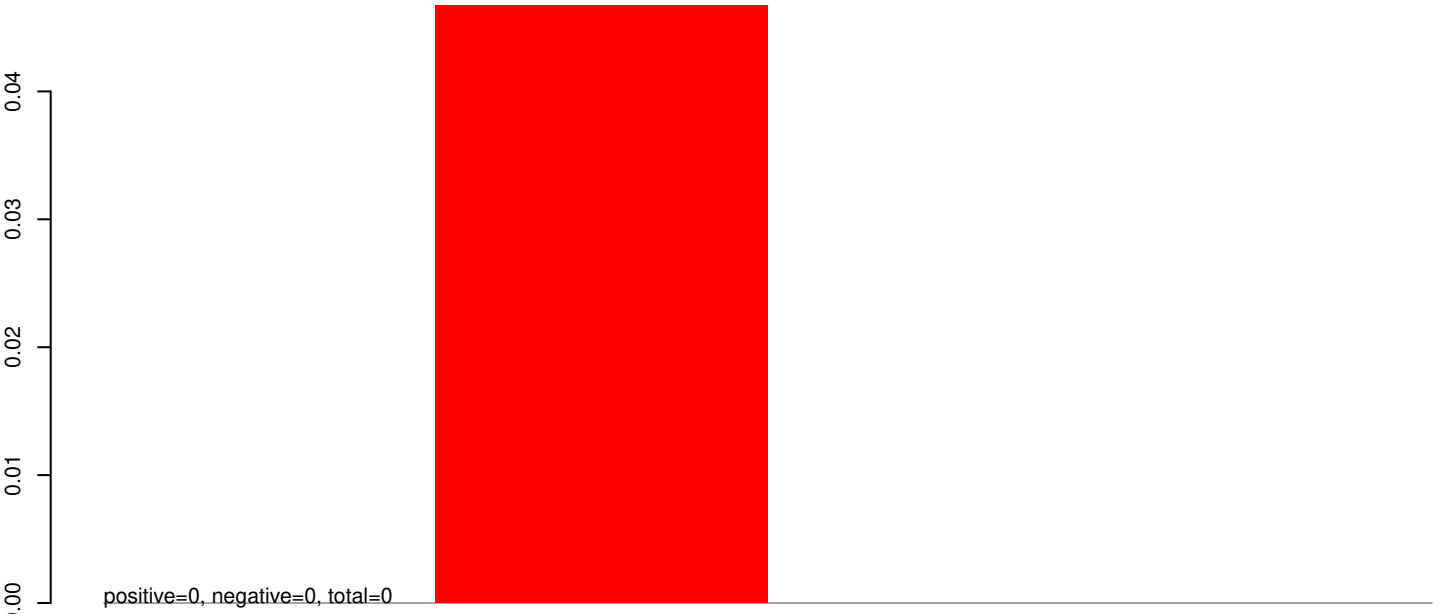
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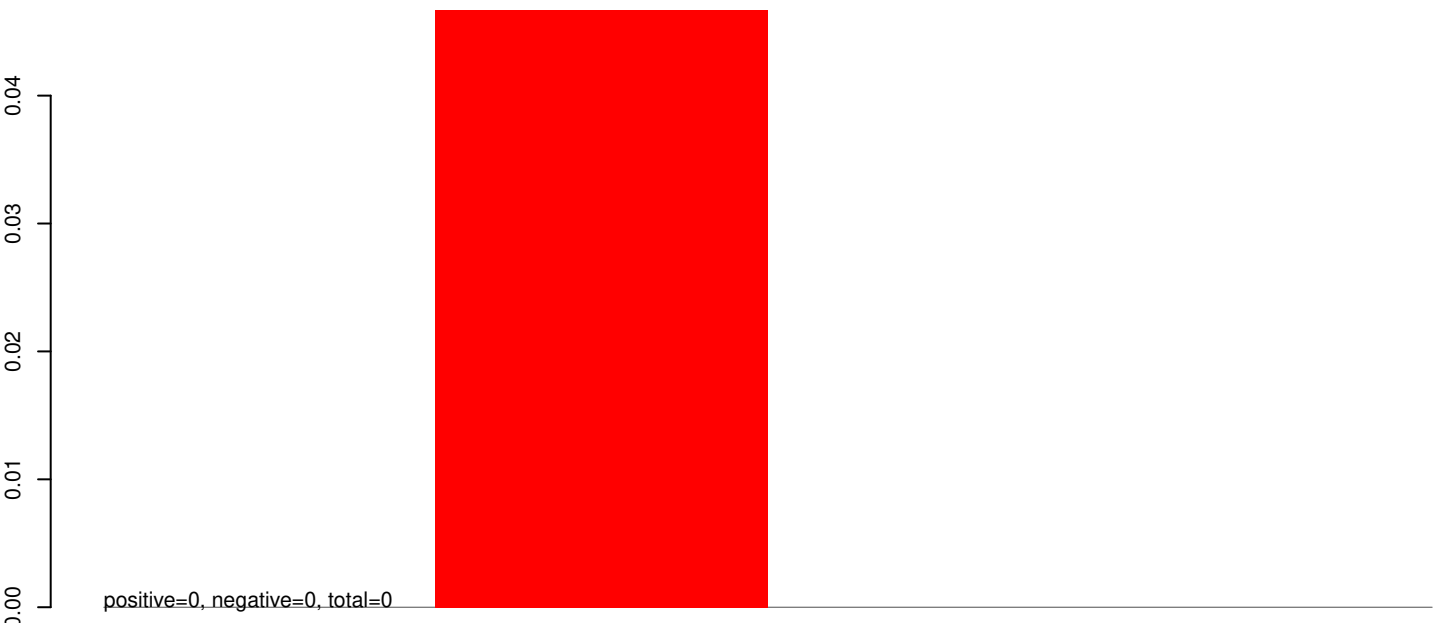
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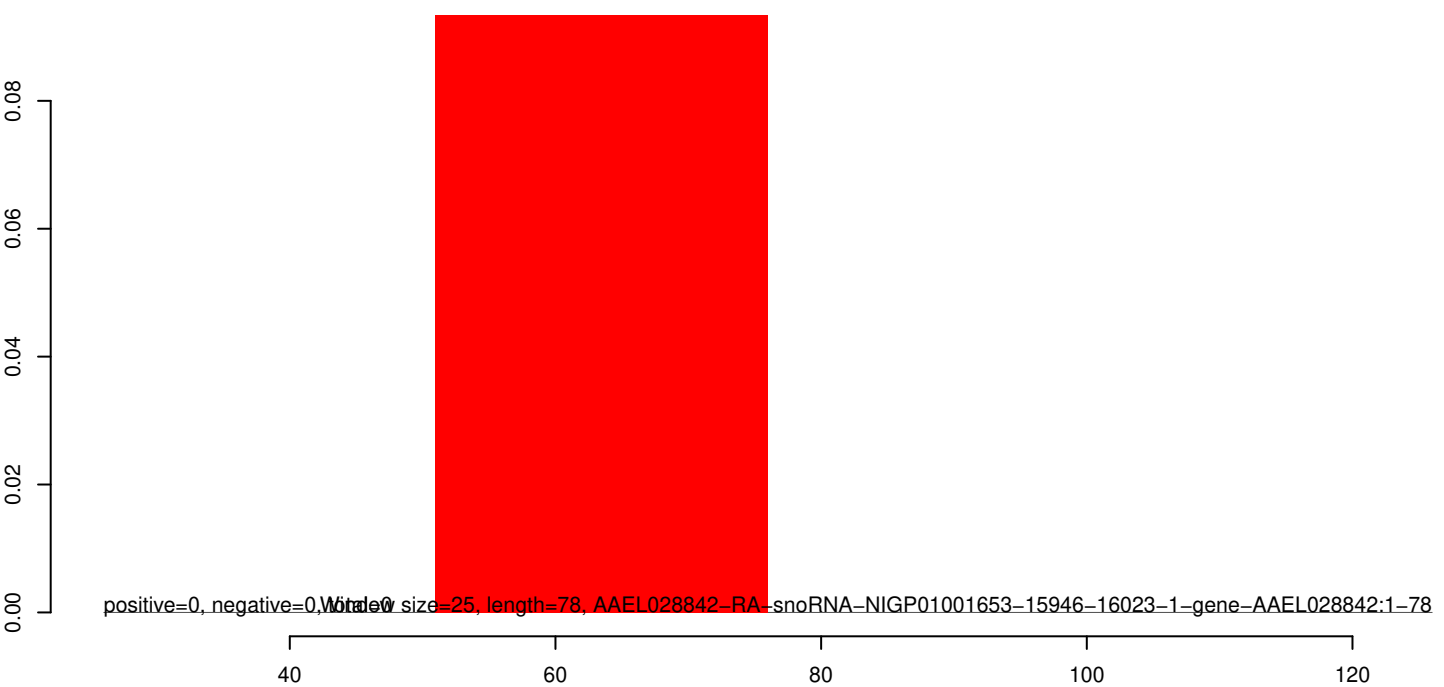
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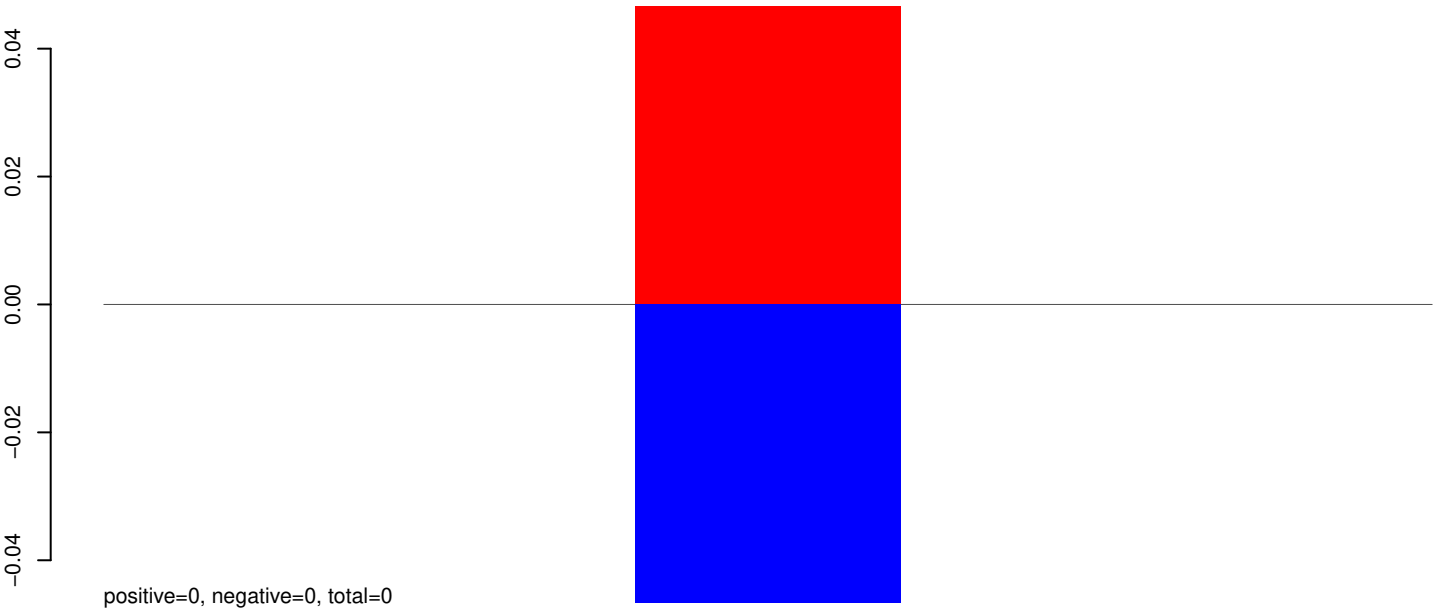
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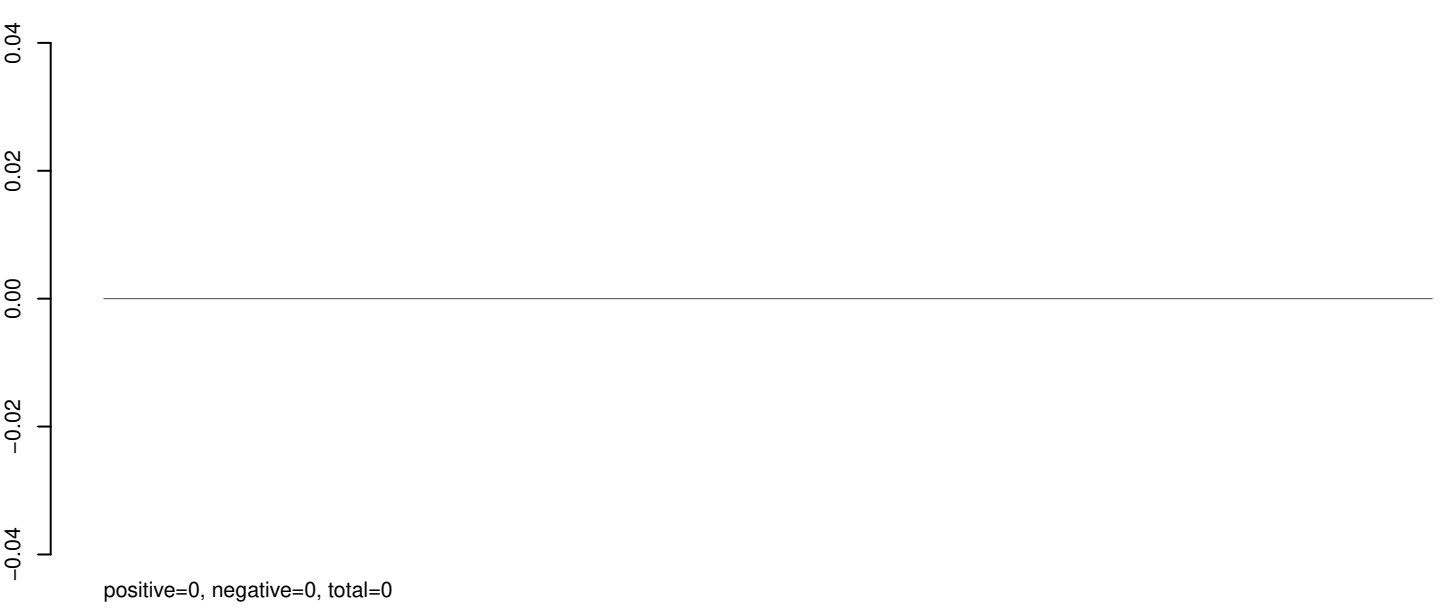
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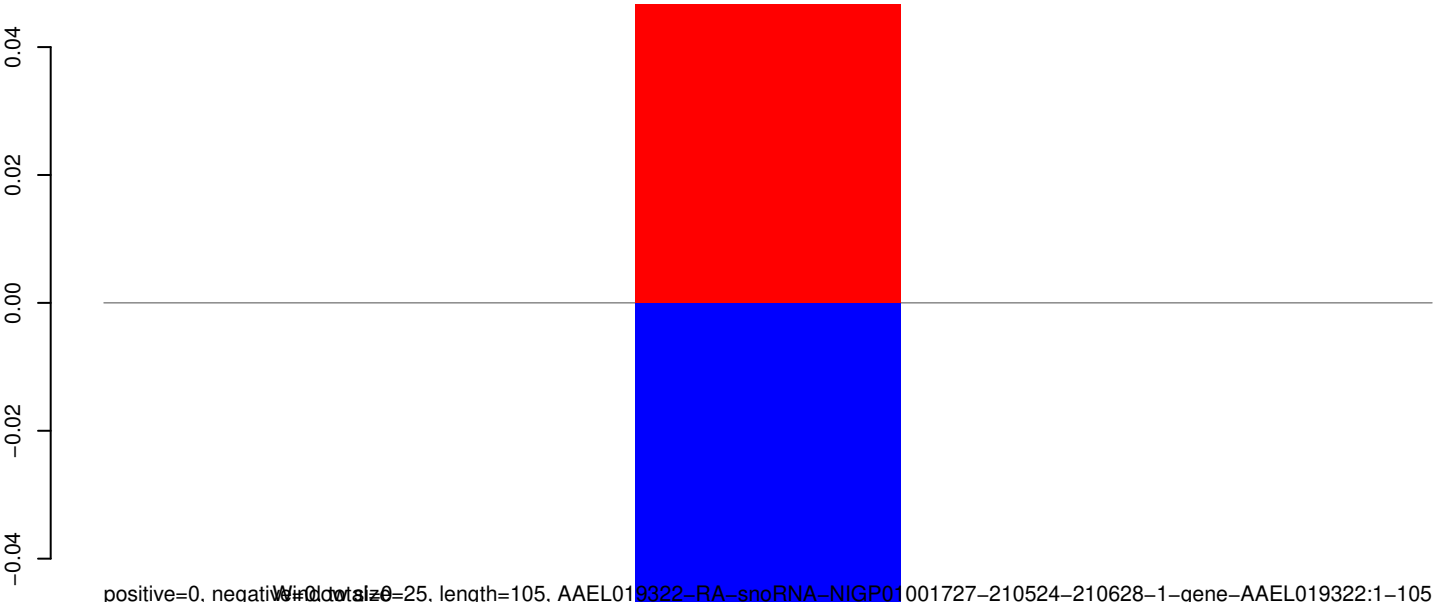
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AeAeg_Aag2SINV_dsLuc.24_35.rep



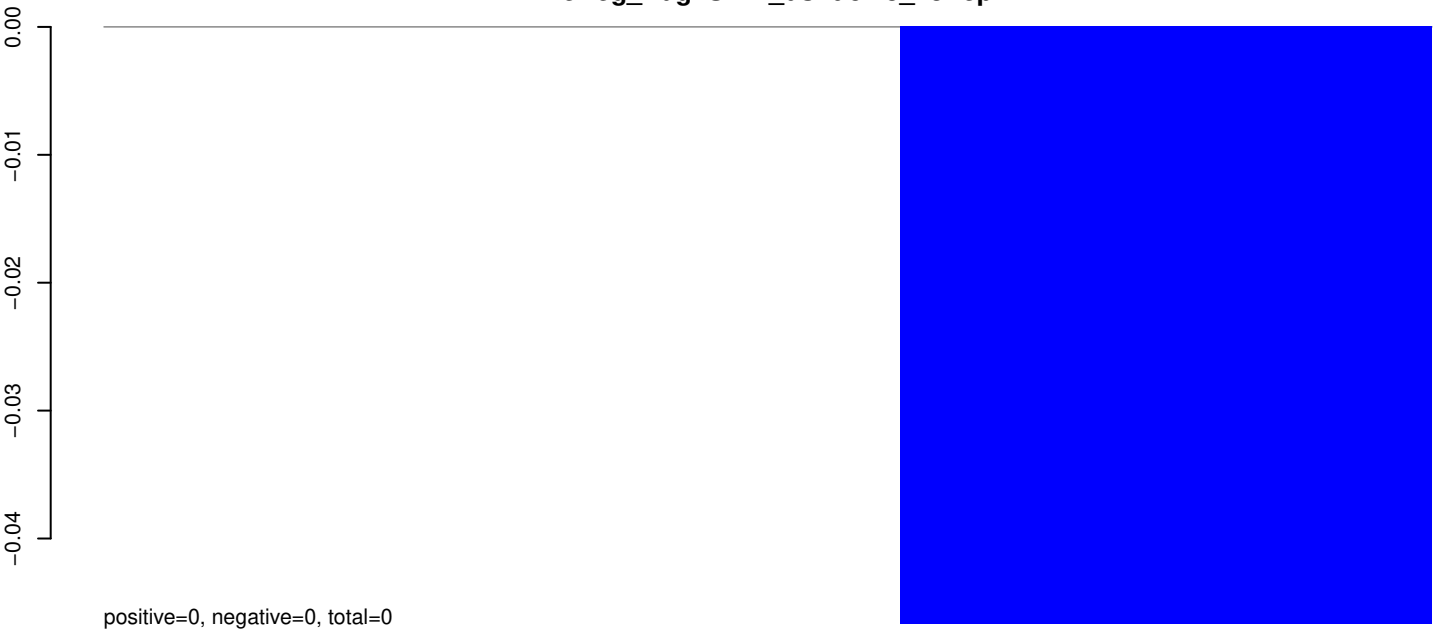
AeAeg_Aag2SINV_dsLuc.rep



Window size=25, length=105, AAEL019322-RA-snoRNA-NIGP01001727-210524-210628-1-gene-AAEL019322:1-105

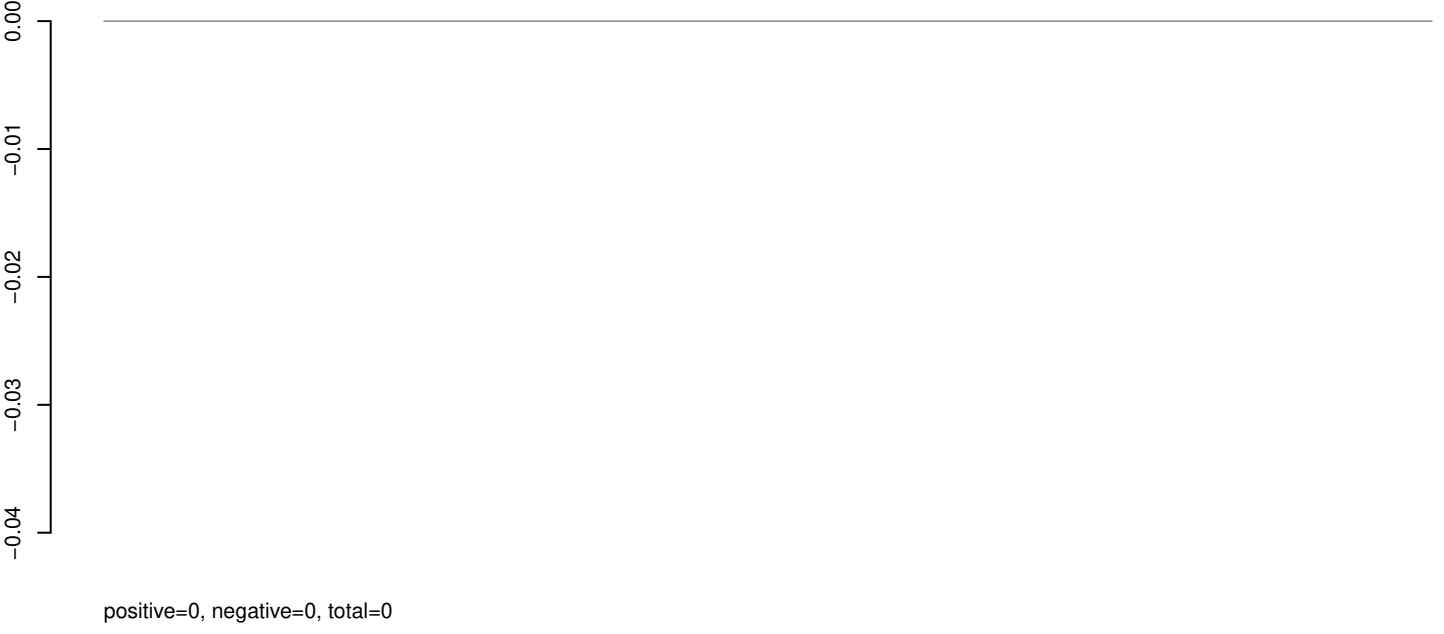
40 60 80 100 120 140

AeAeg_Aag2SINV_dsLuc.18_23.rep



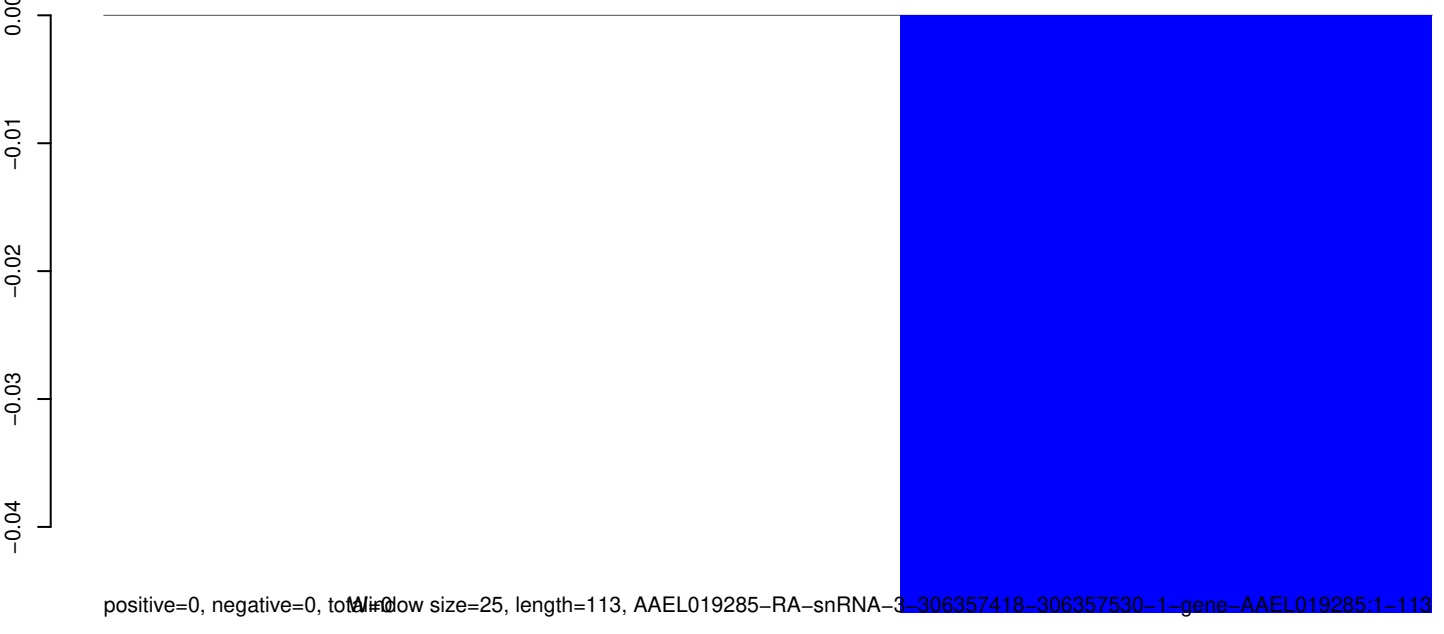
positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.24_35.rep



positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsLuc.rep



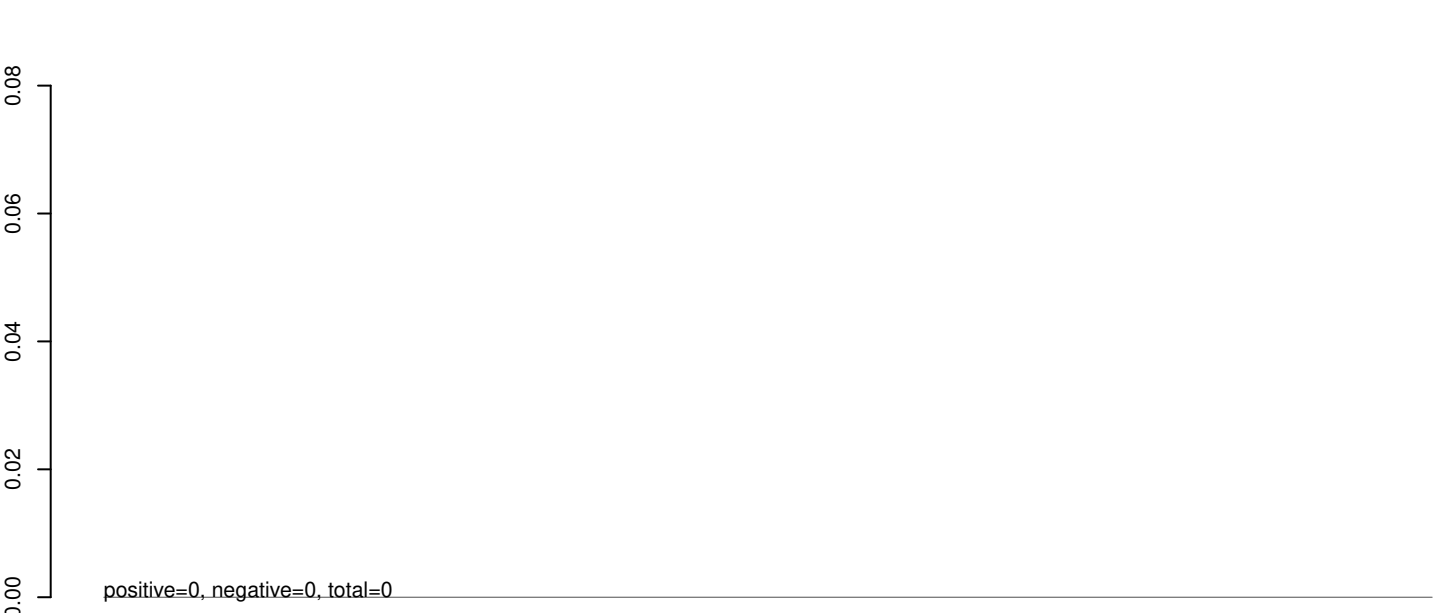
positive=0, negative=0, total=0 window size=25, length=113, AAEL019285-RA-snrRNA-3-306357418-306357530-1-gene-AAEL019285:1-113

40 60 80 100 120 140

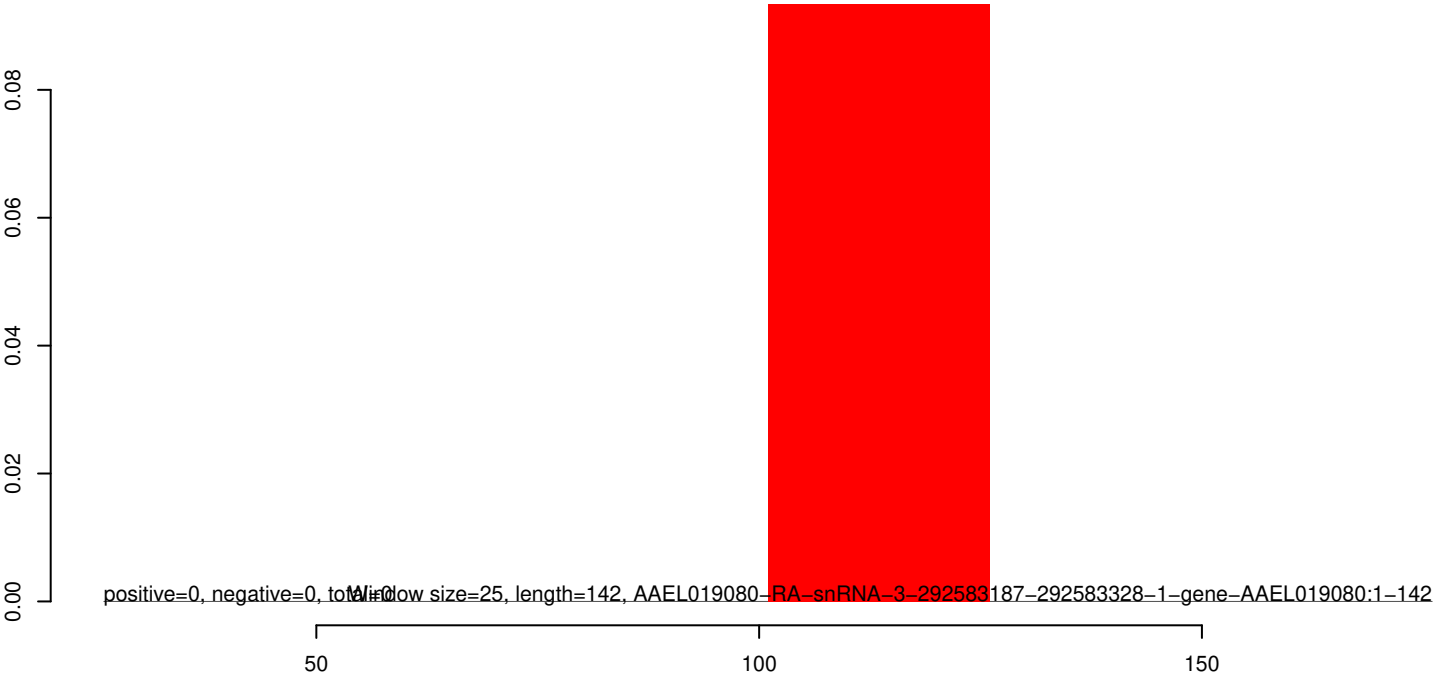
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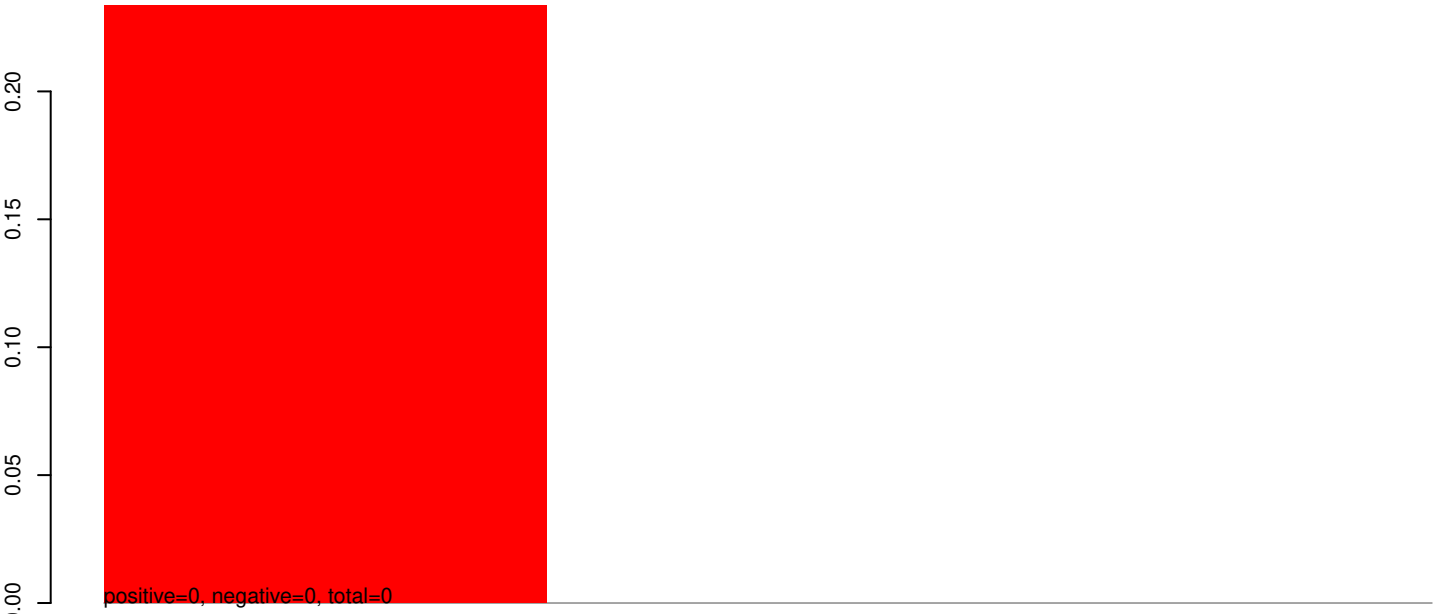
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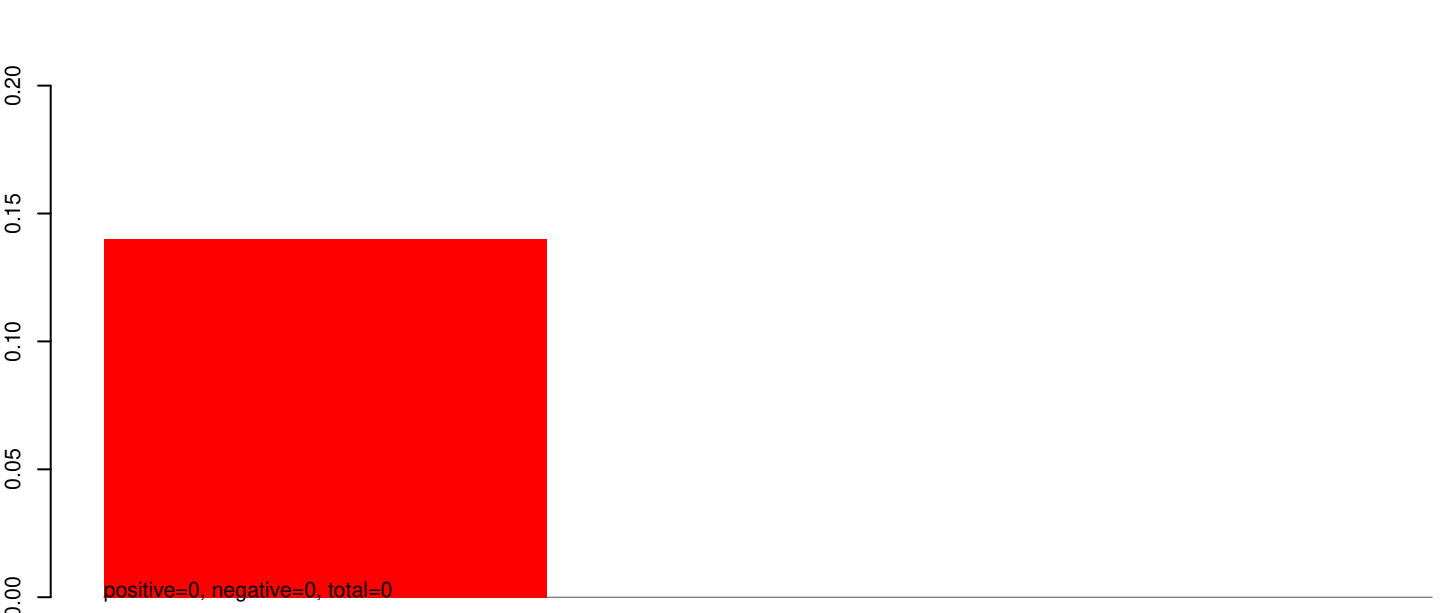
AeAeg_Aag2SINV_dsLuc.rep



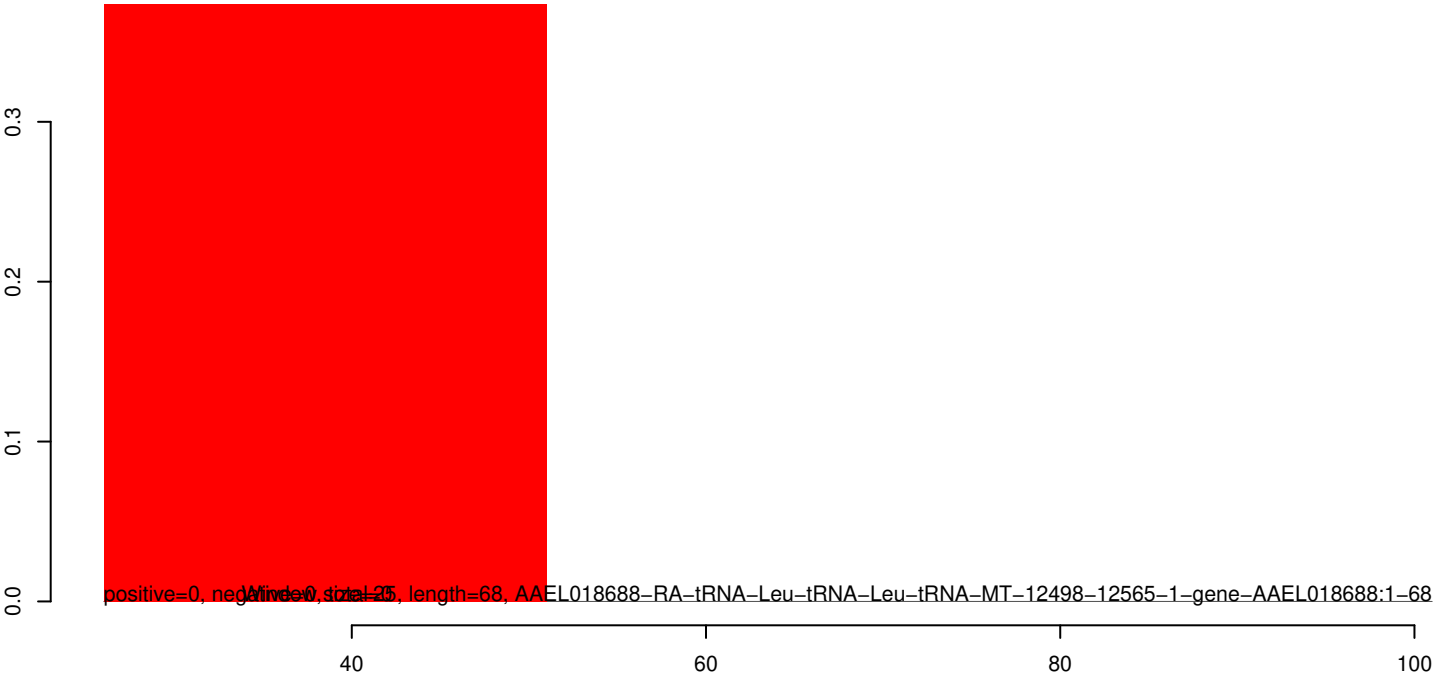
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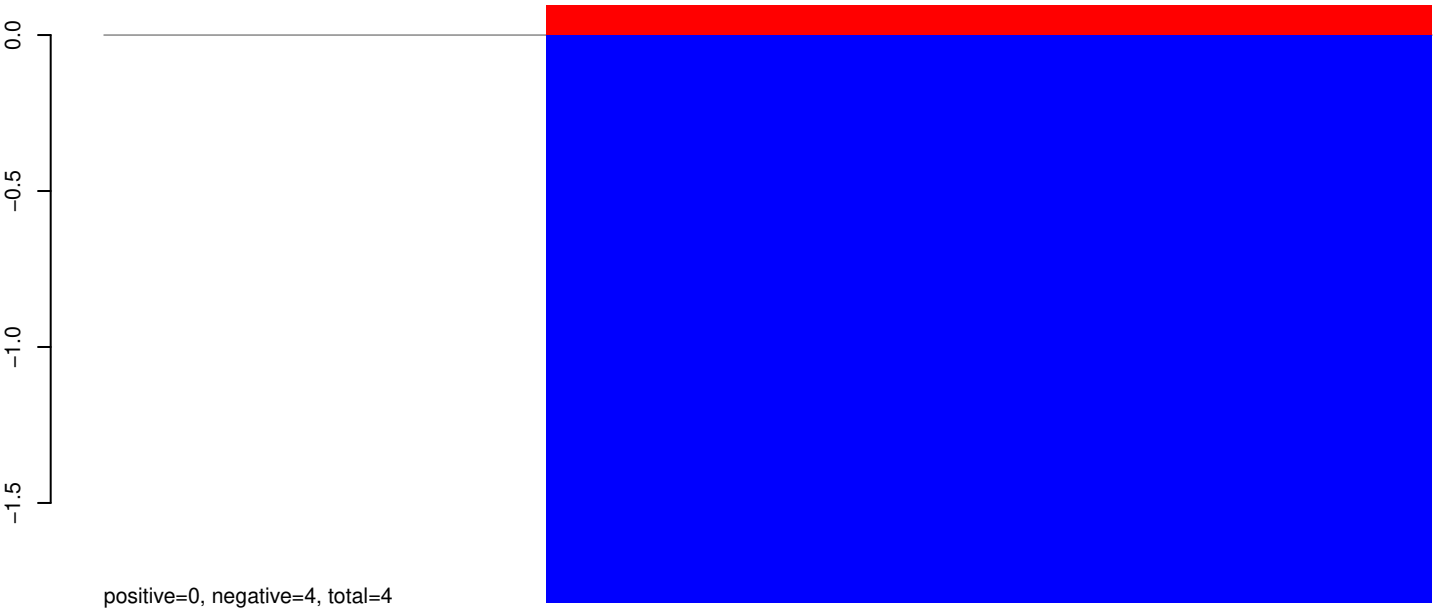
AeAeg_Aag2SINV_dsLuc.24_35.rep



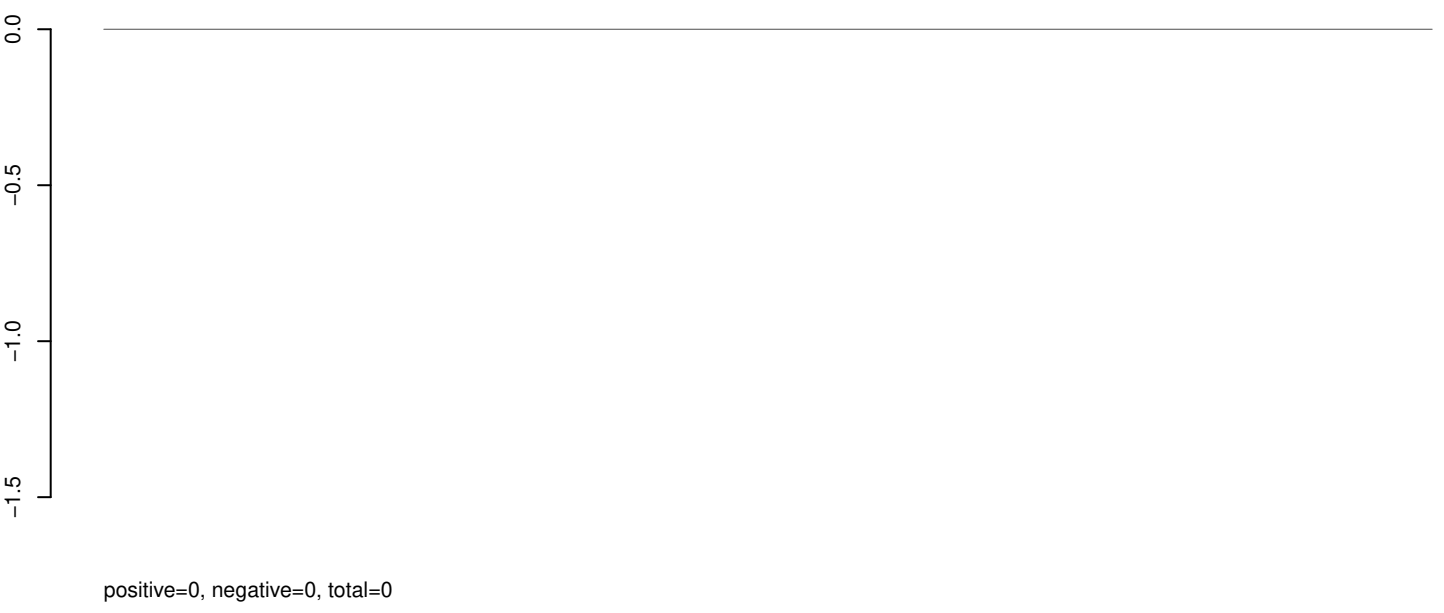
AeAeg_Aag2SINV_dsLuc.rep



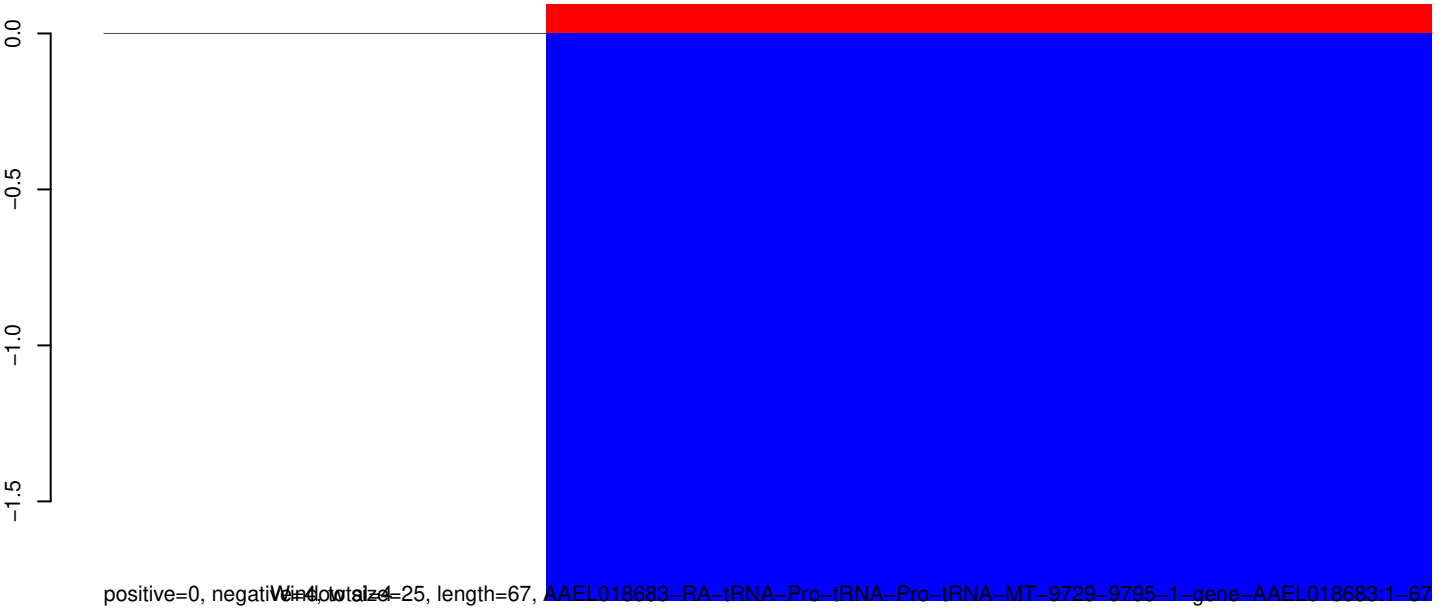
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AeAeg_Aag2SINV_dsLuc.24_35.rep

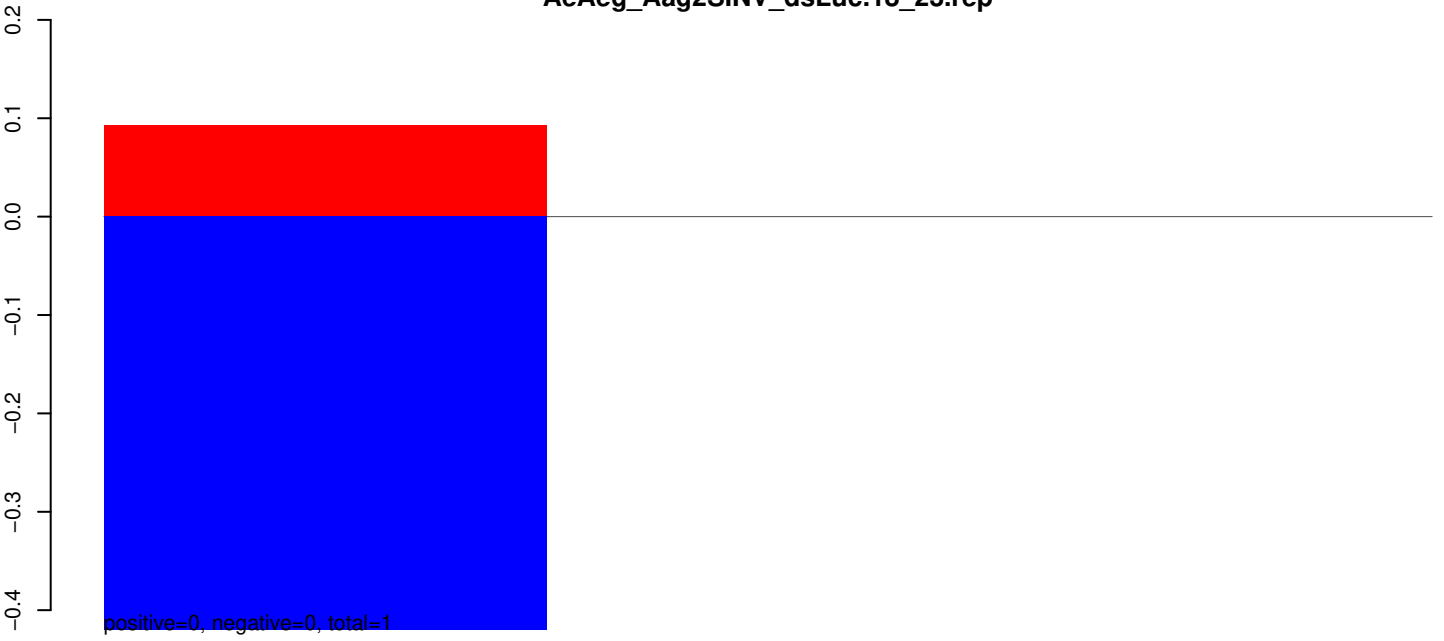


AeAeg_Aag2SINV_dsLuc.rep

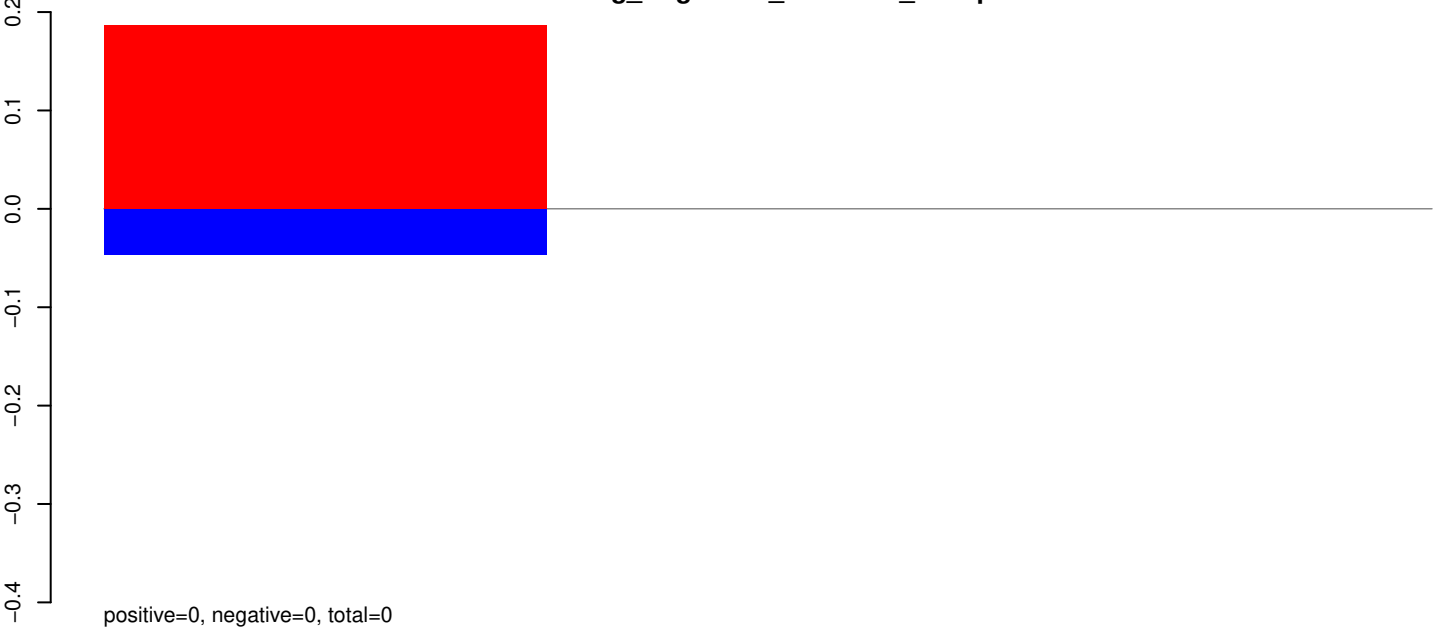


40 60 80 100

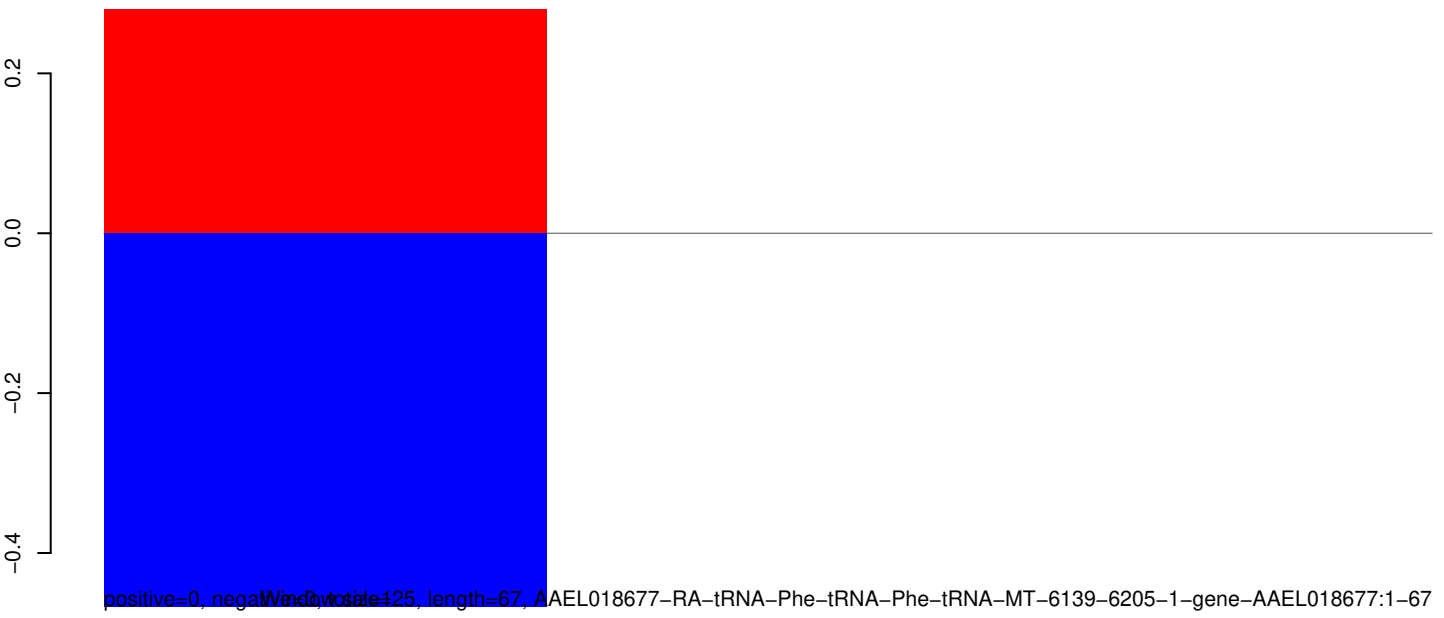
AeAeg_Aag2SINV_dsLuc.18_23.rep



AeAeg_Aag2SINV_dsLuc.24_35.rep

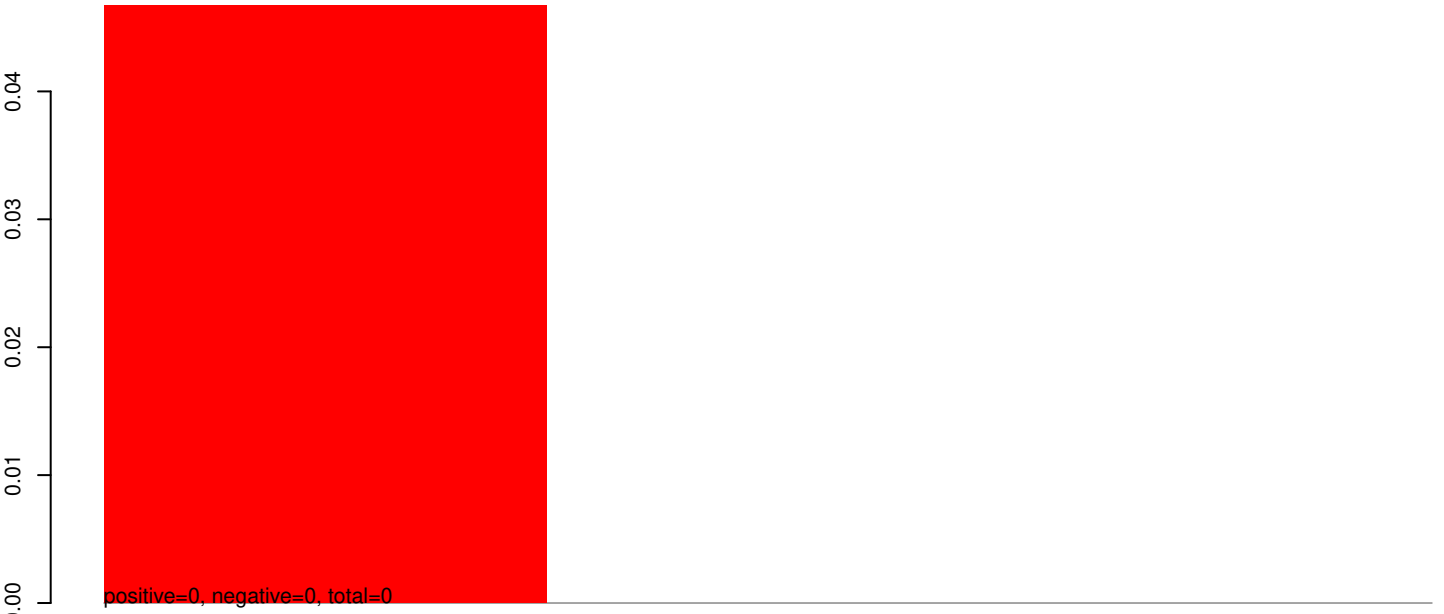


AeAeg_Aag2SINV_dsLuc.rep

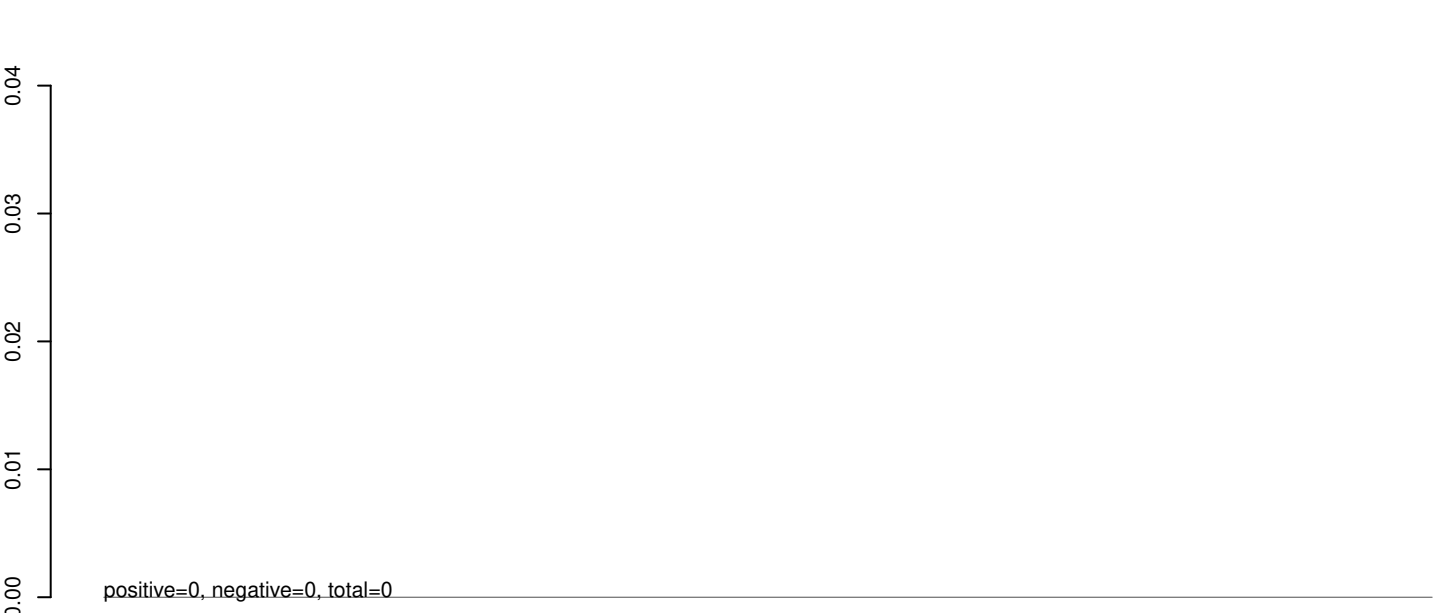


Window size=425, length=67, AAEL018677-RA-tRNA-Phe-tRNA-Phe-tRNA-MT-6139-6205-1-gene-AAEL018677:1-67

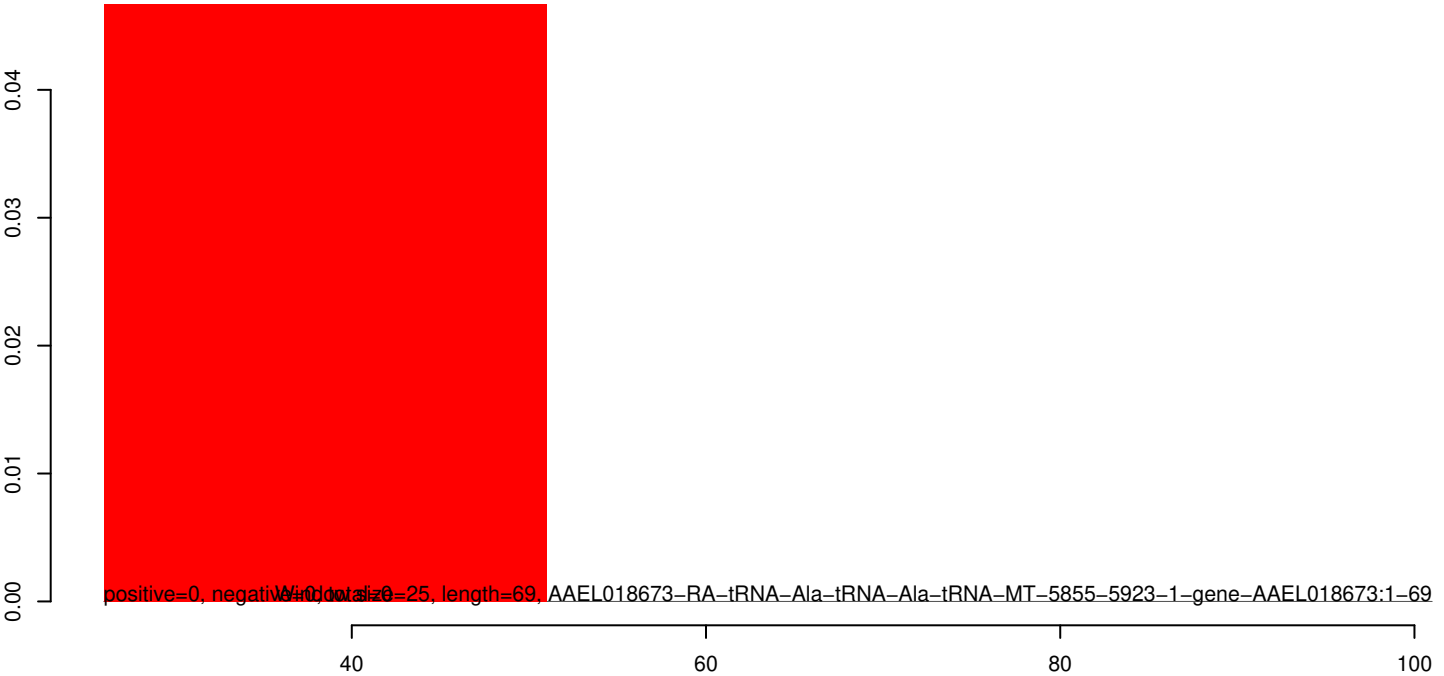
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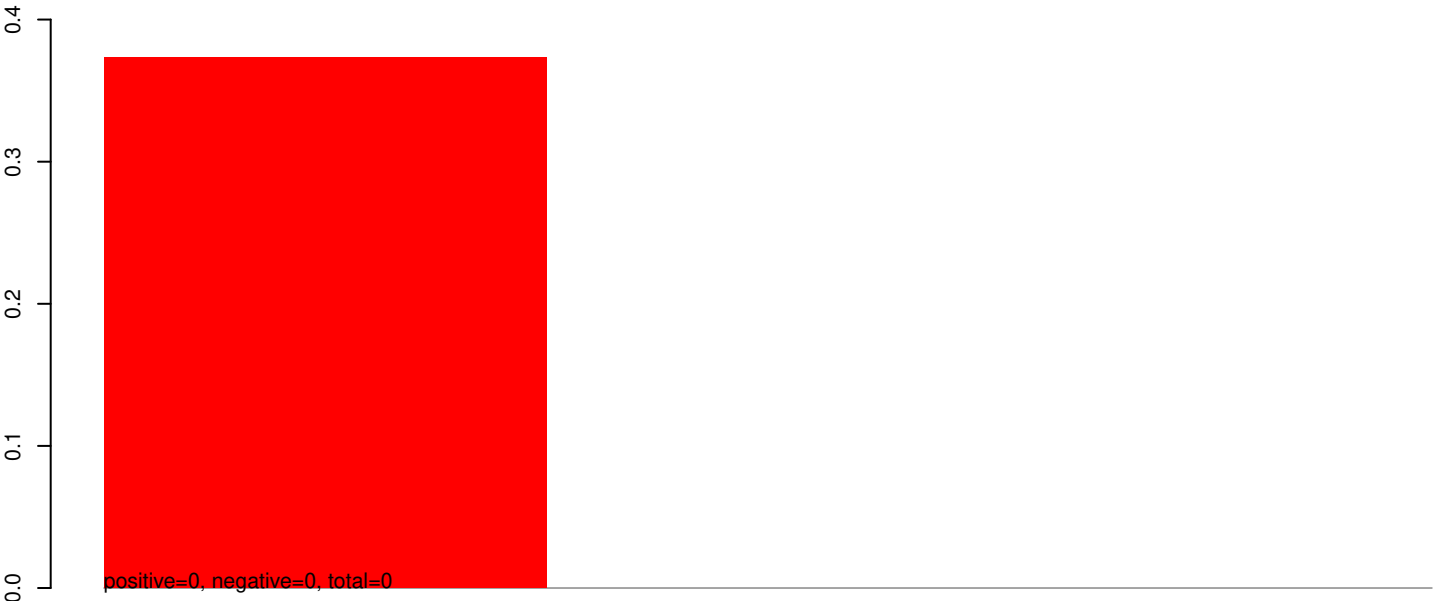
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AeAeg_Aag2SINV_dsLuc.rep



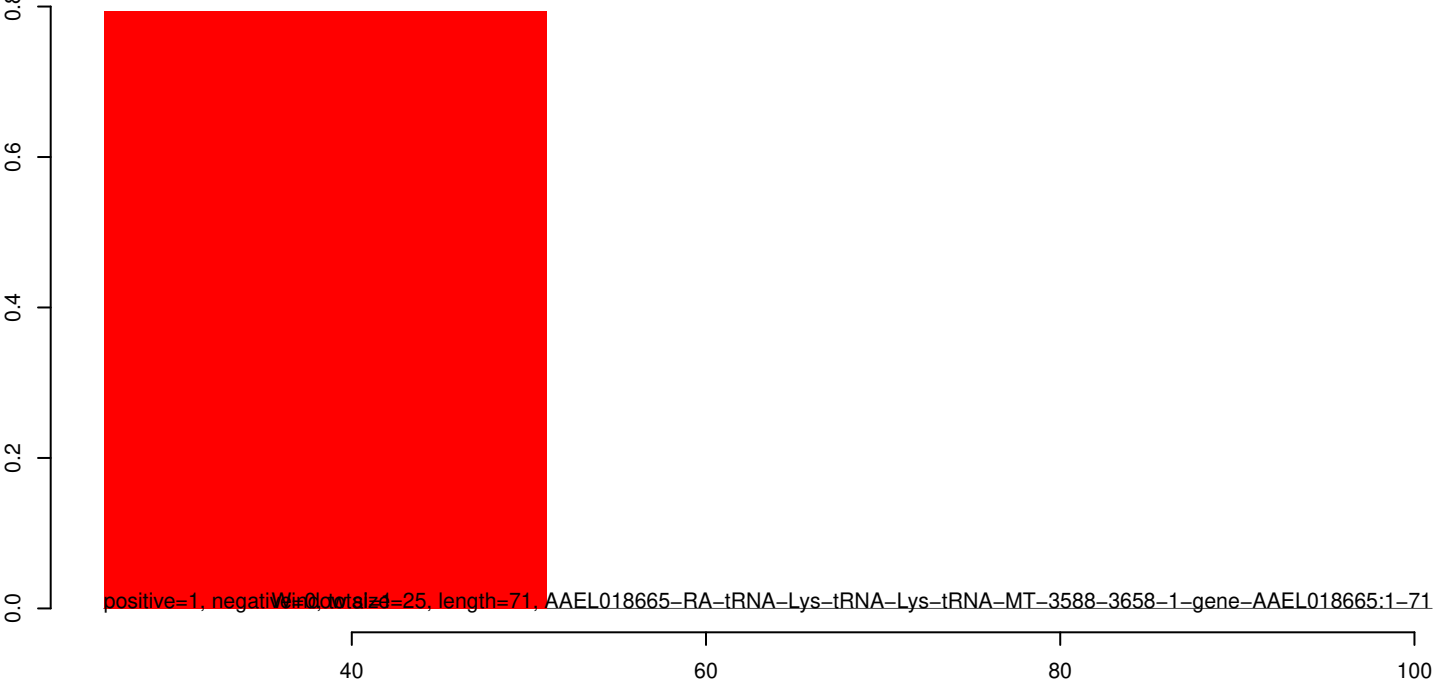
AeAeg_Aag2SINV_dsLuc.18_23.rep



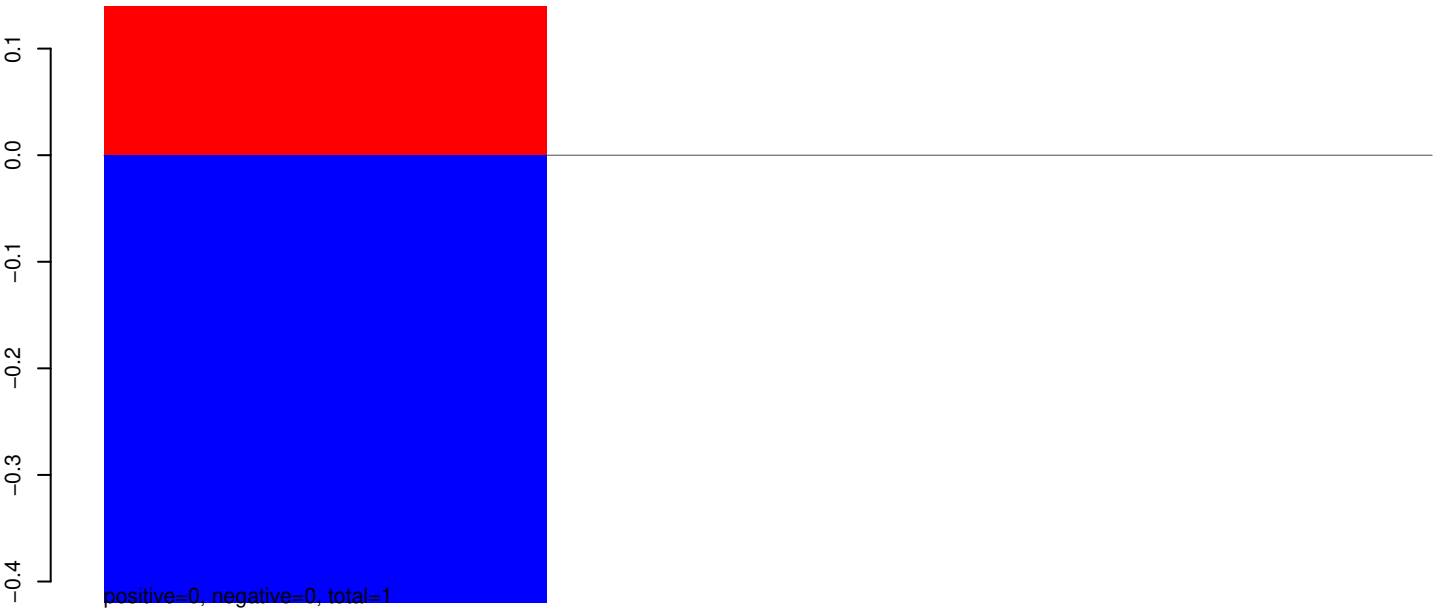
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AeAeg_Aag2SINV_dsLuc.rep



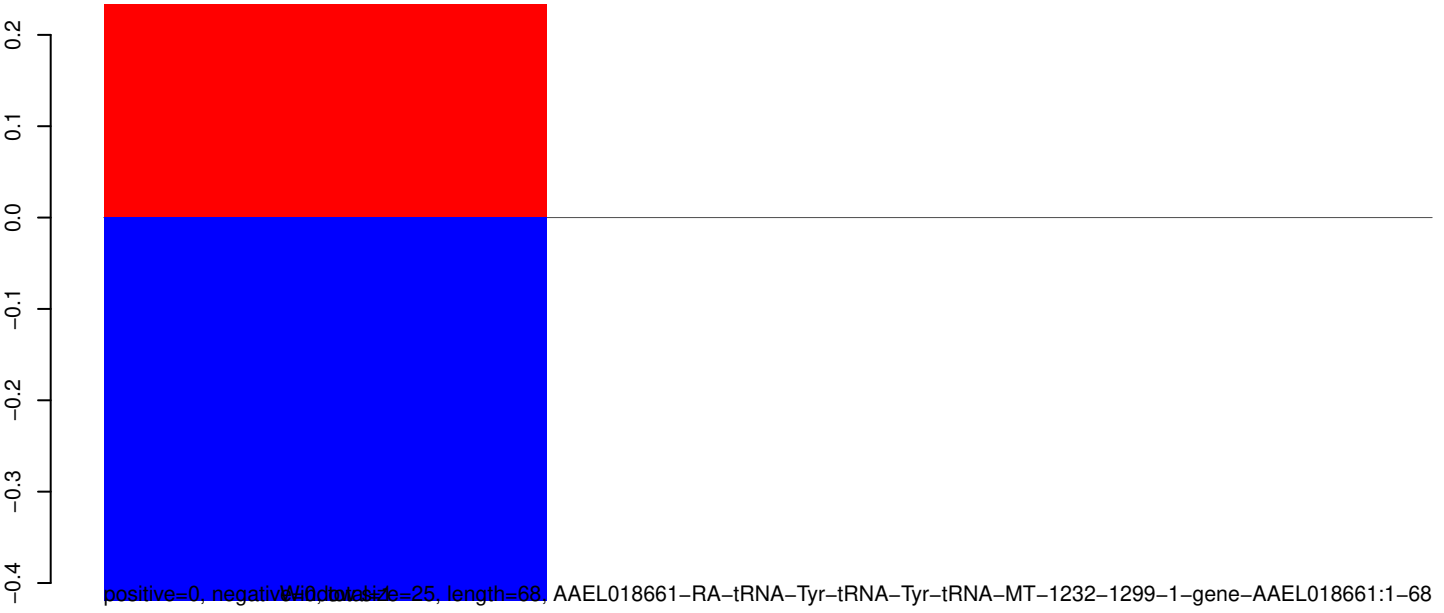
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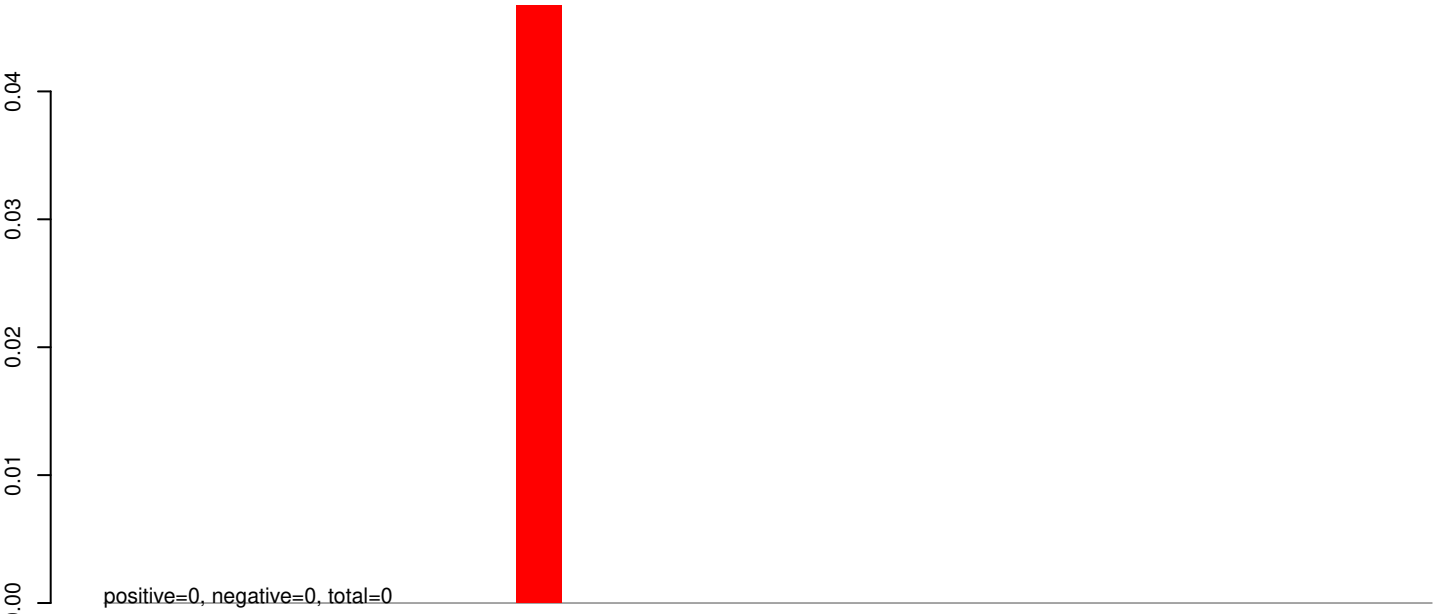
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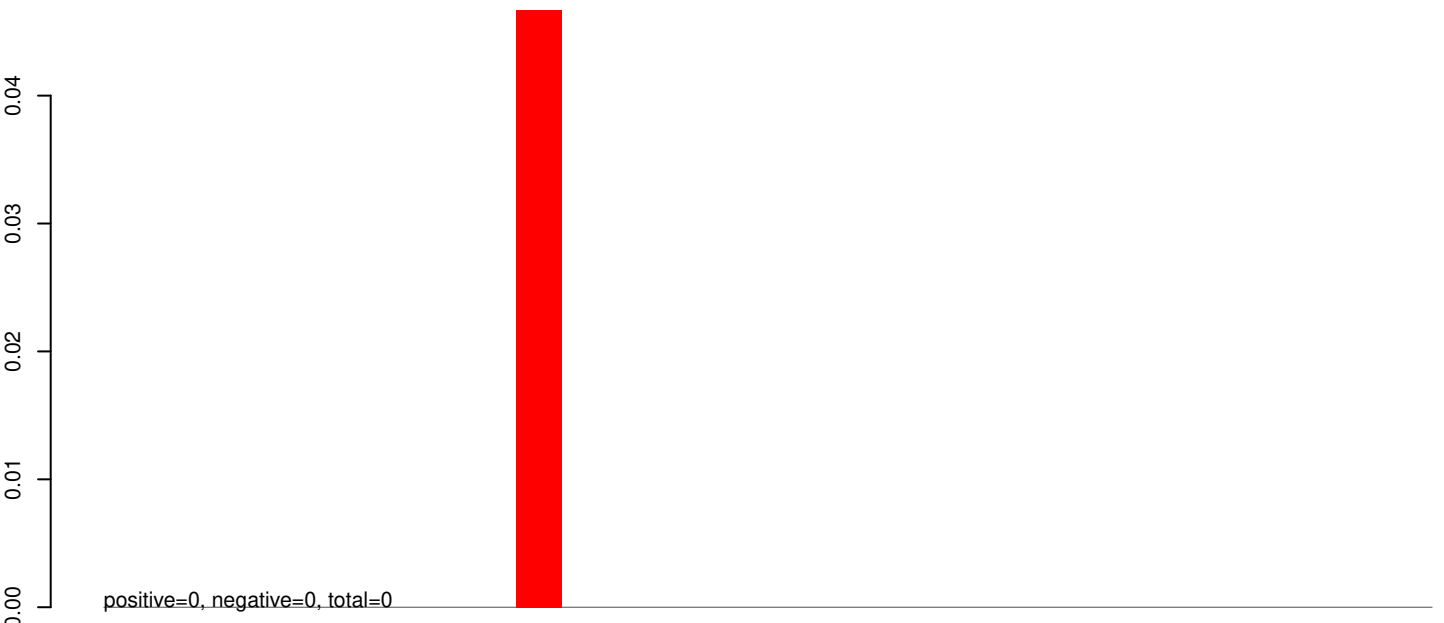
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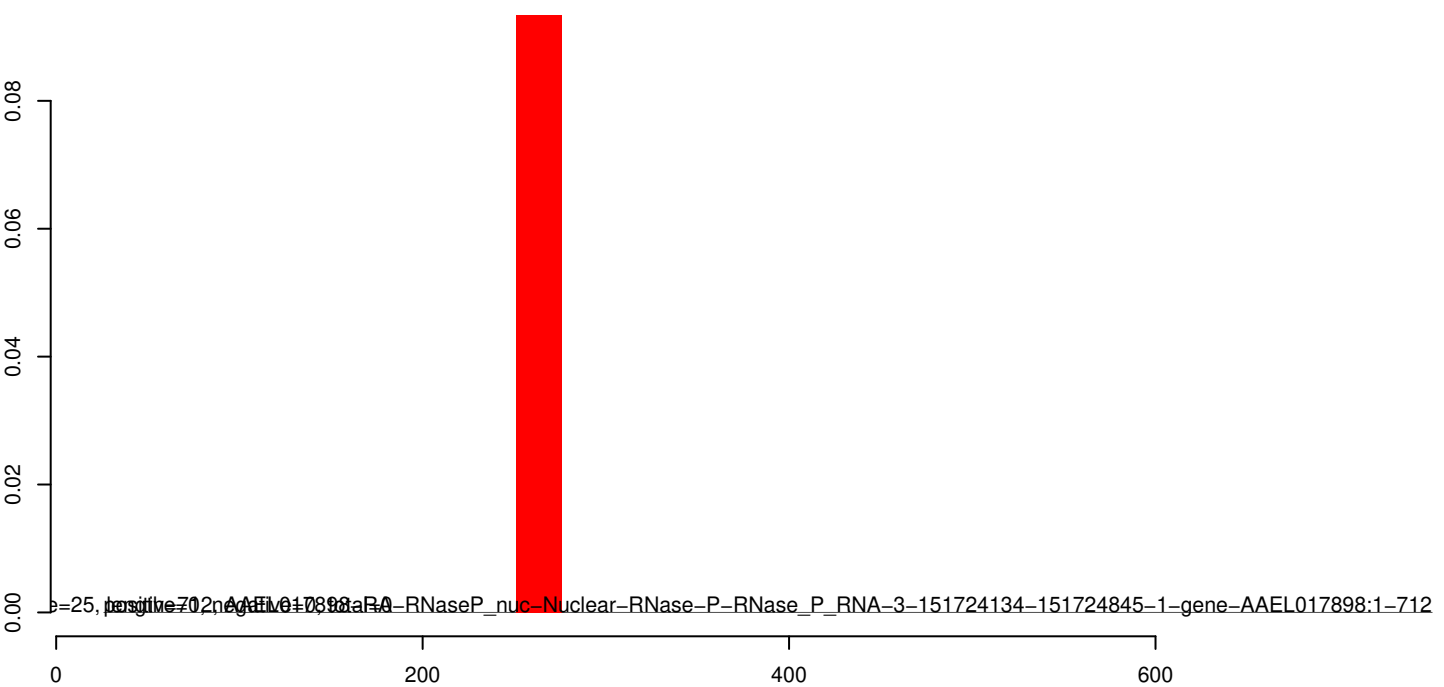
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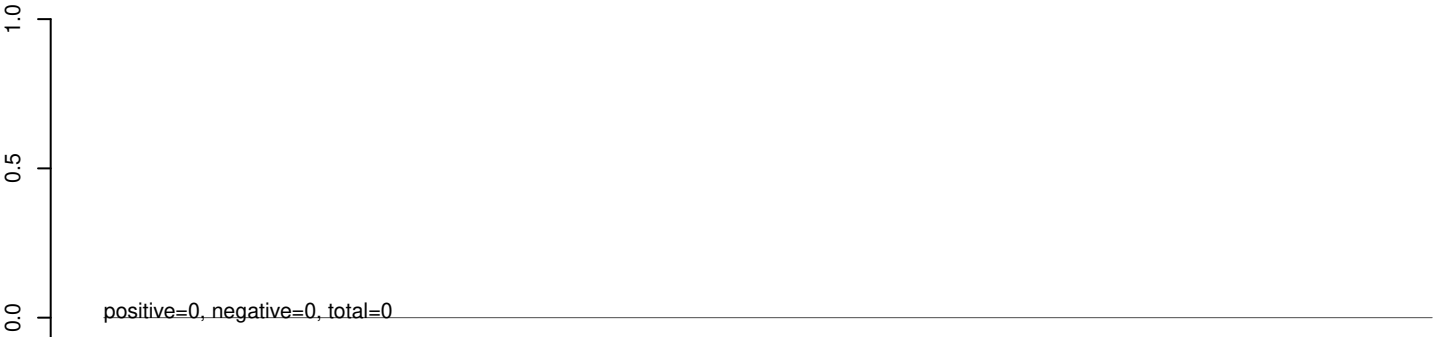
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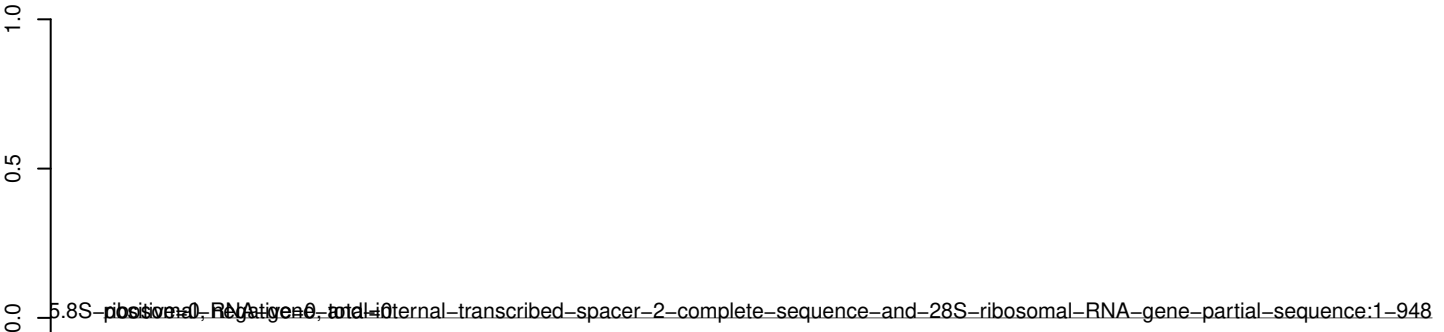
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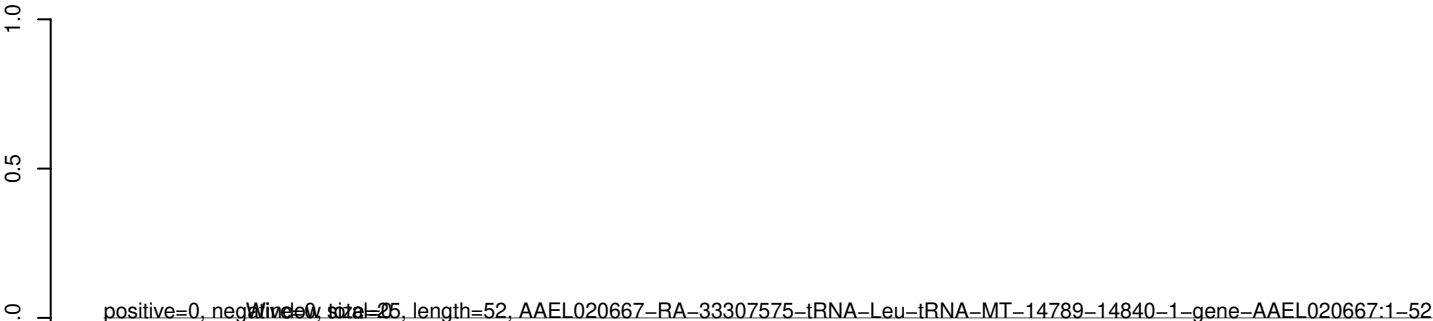
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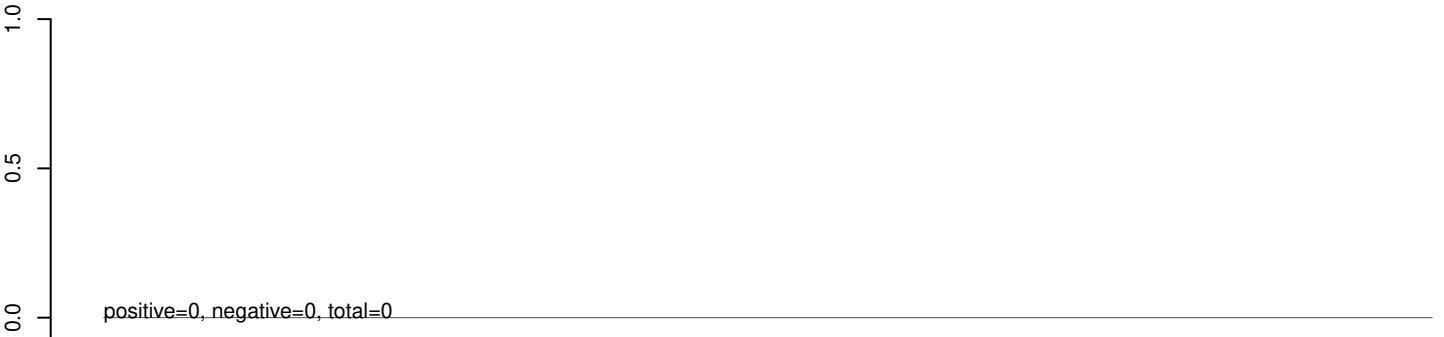
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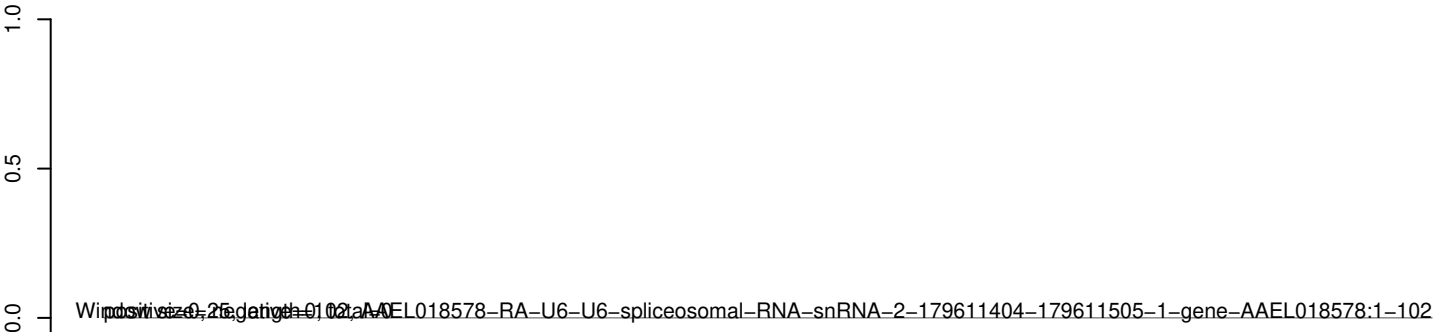
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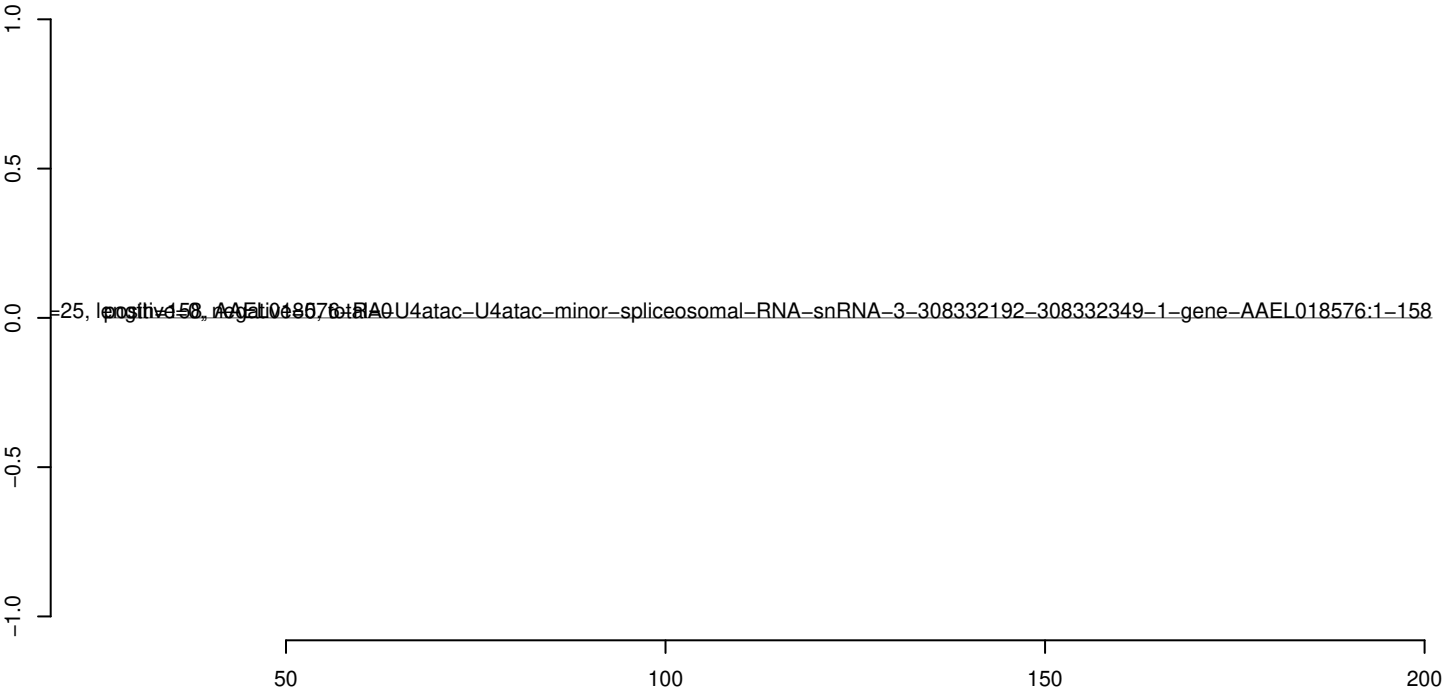
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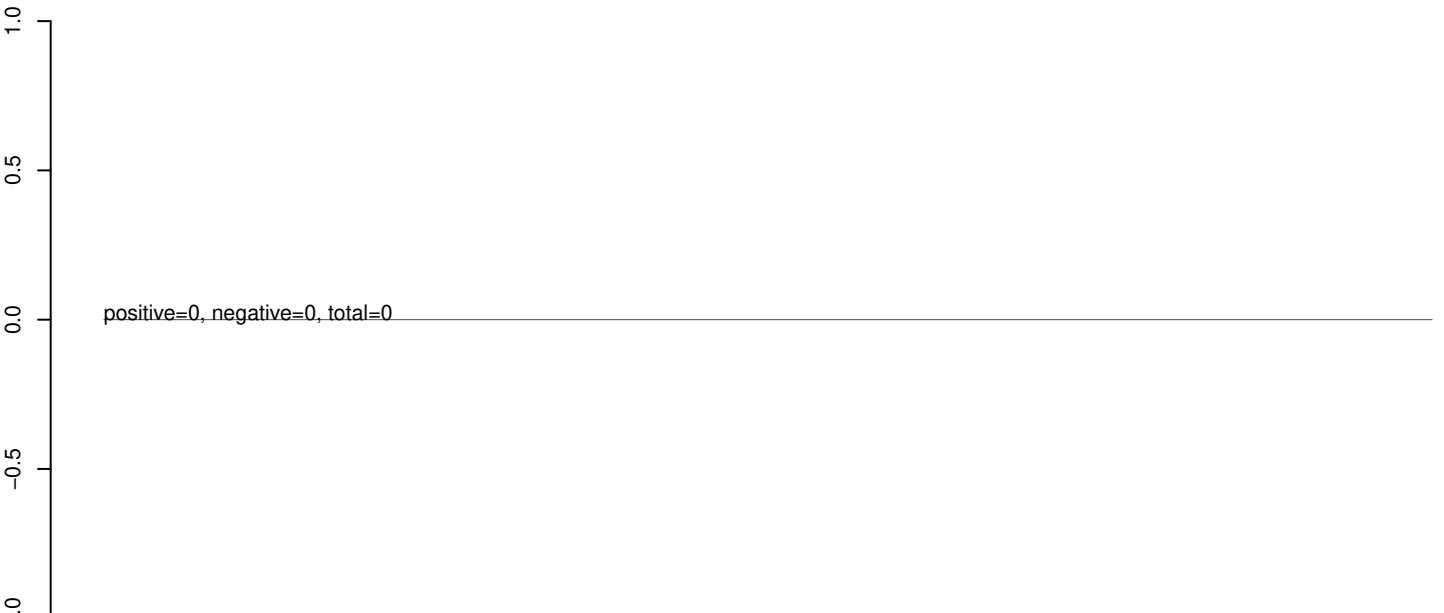
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AeAeg_Aag2SINV_dsLuc.rep

