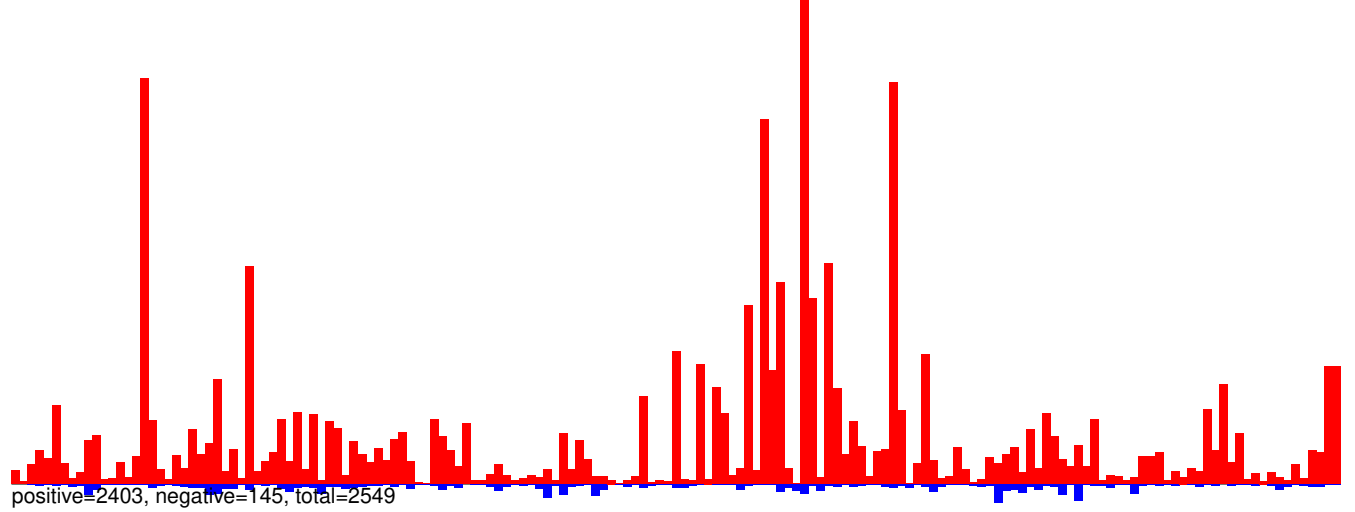
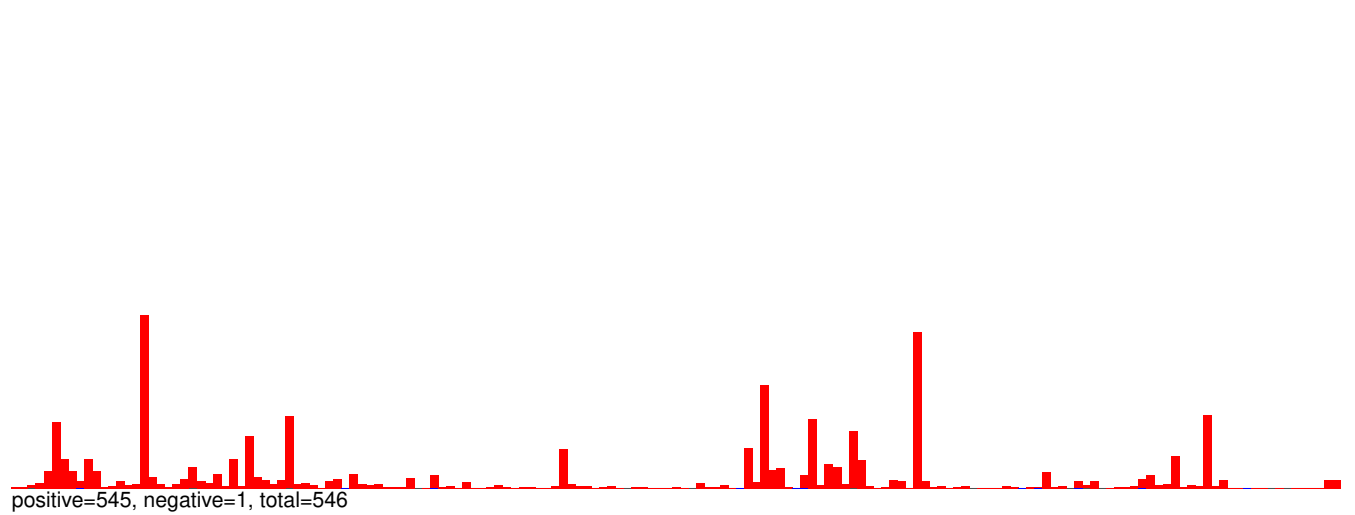


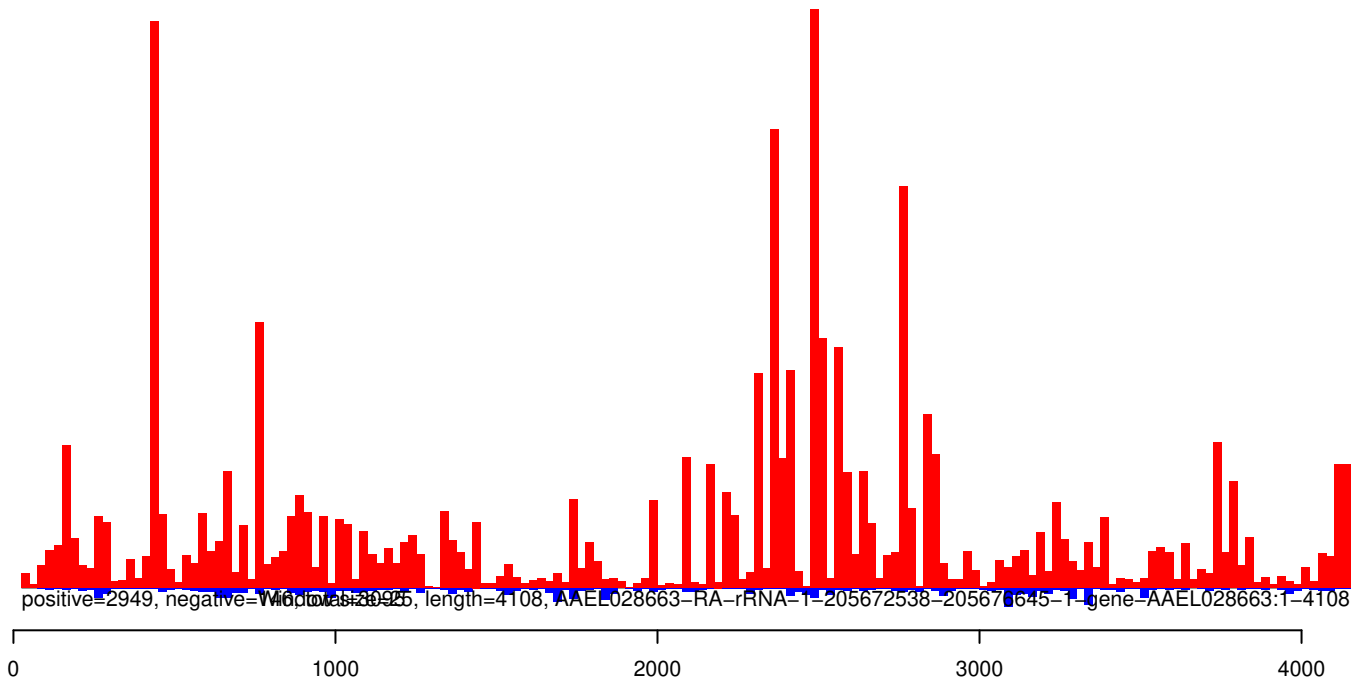
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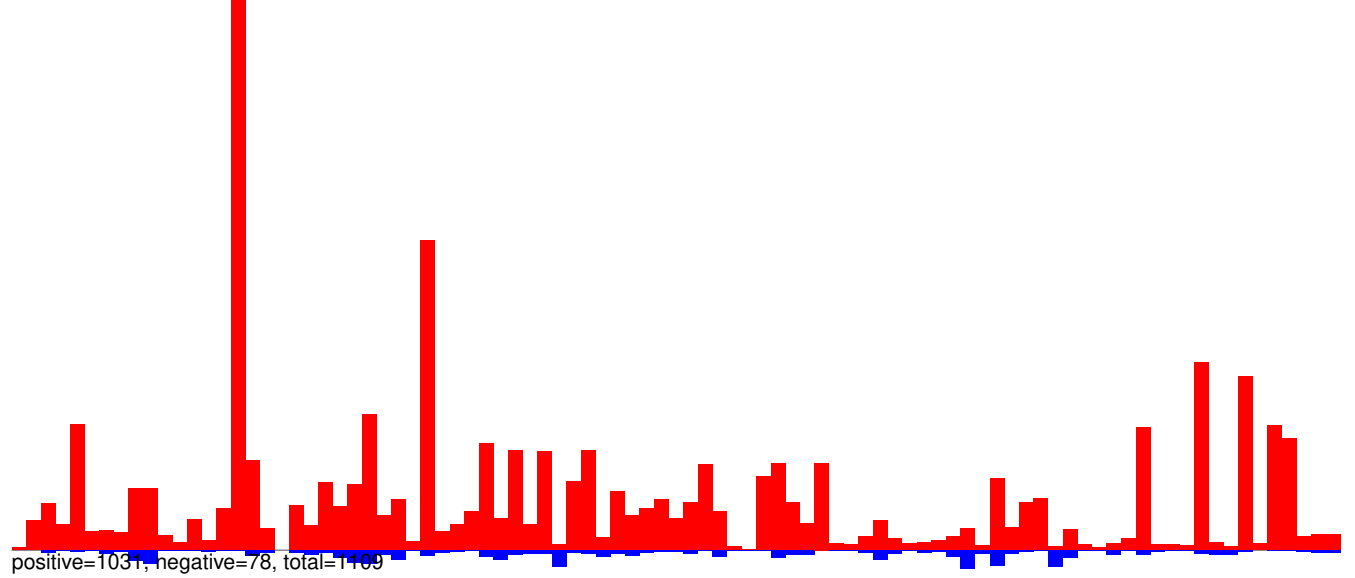
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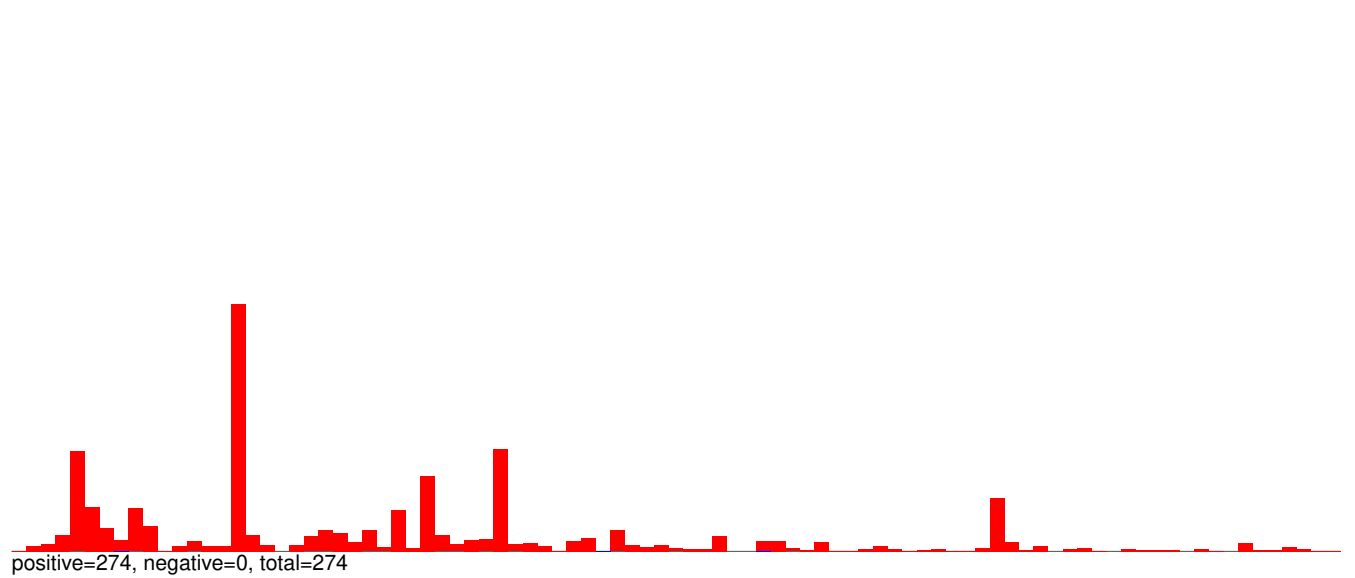
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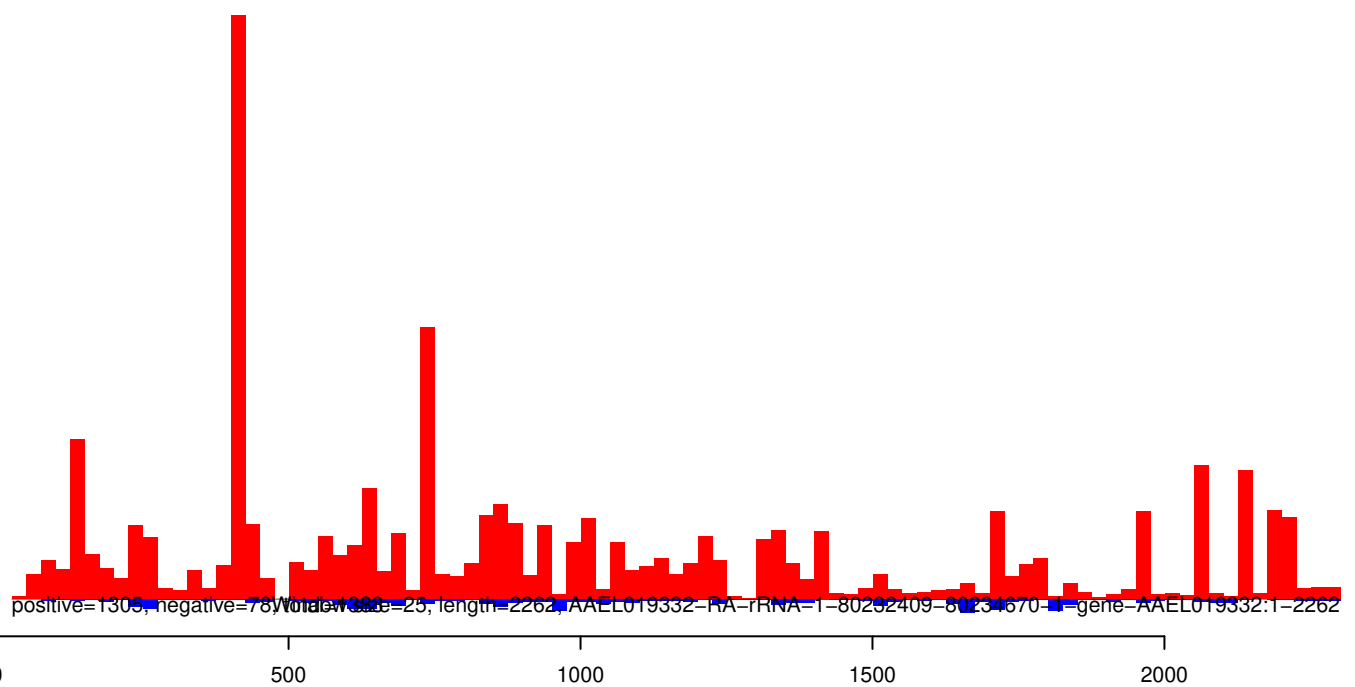
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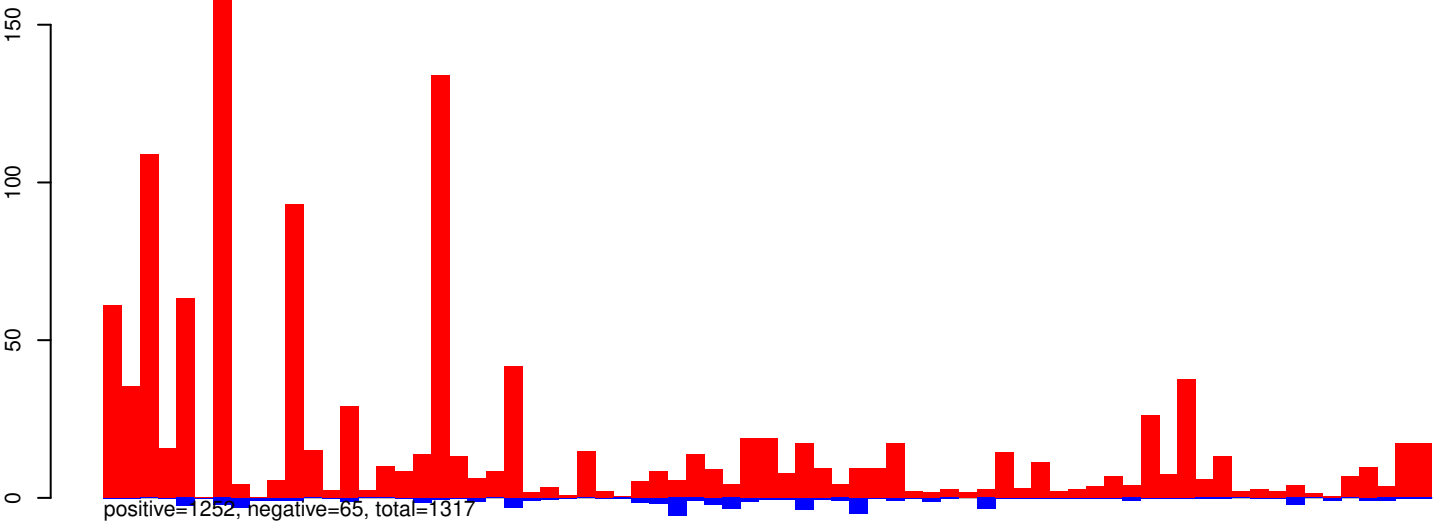
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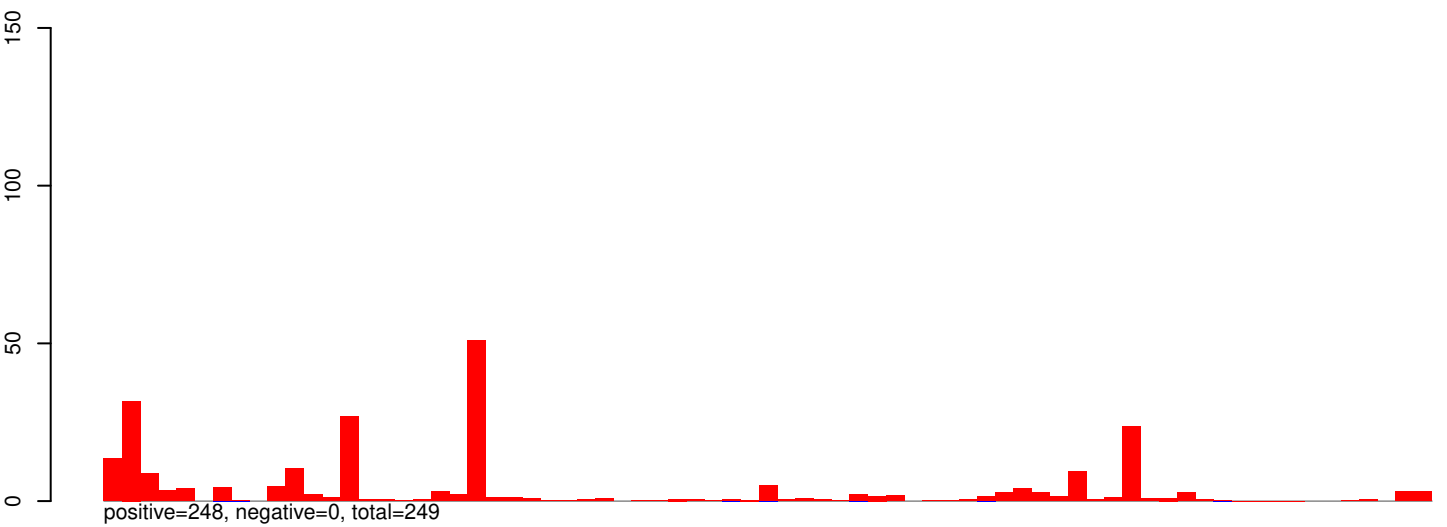
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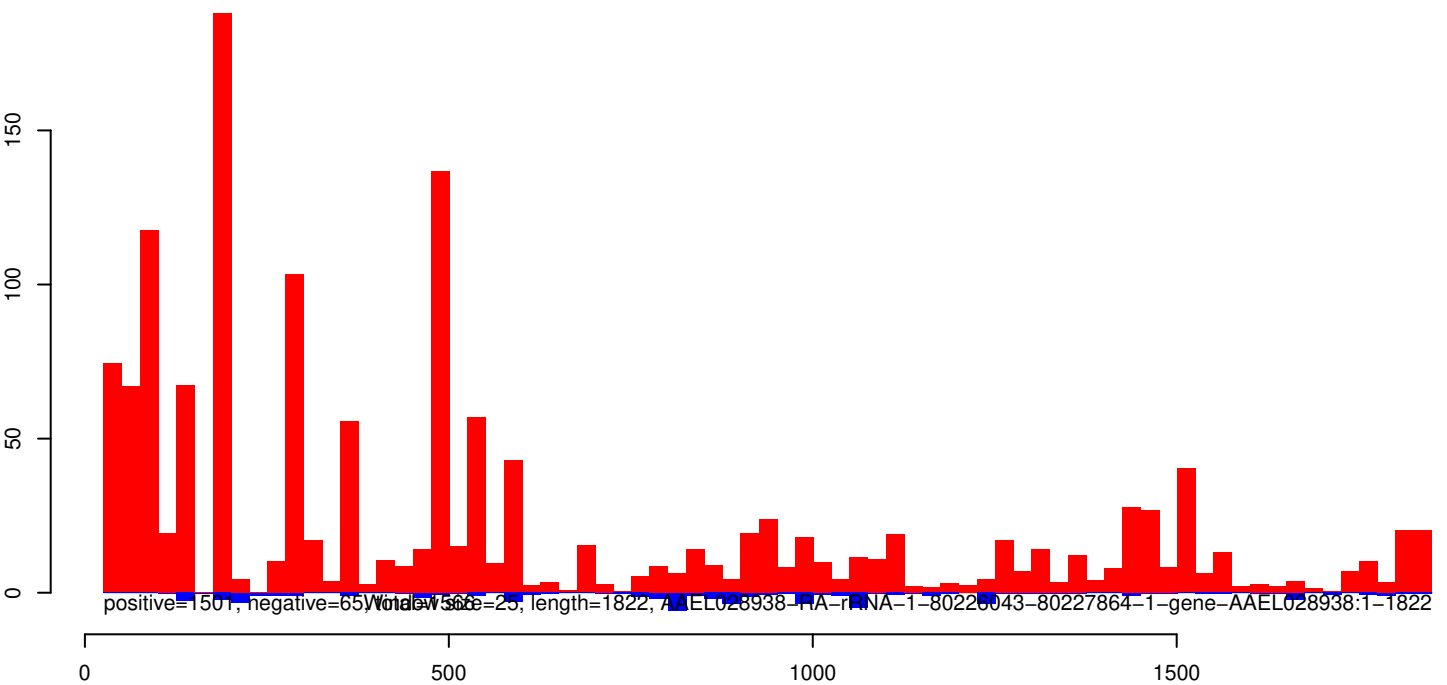
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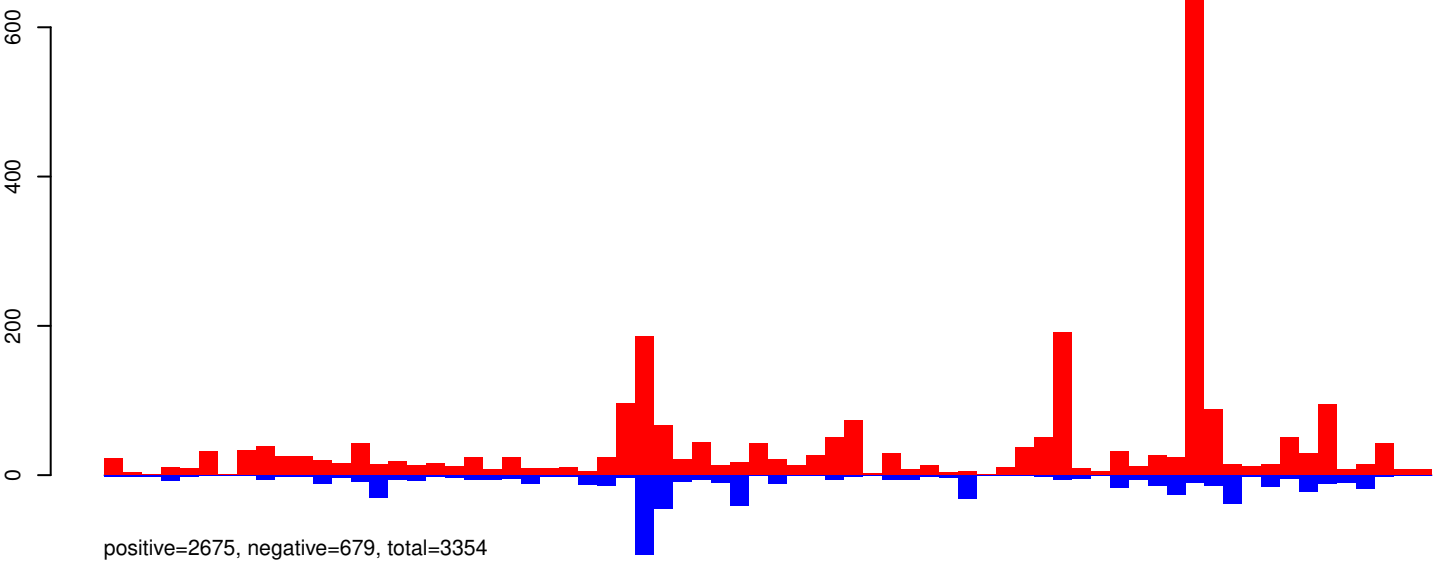
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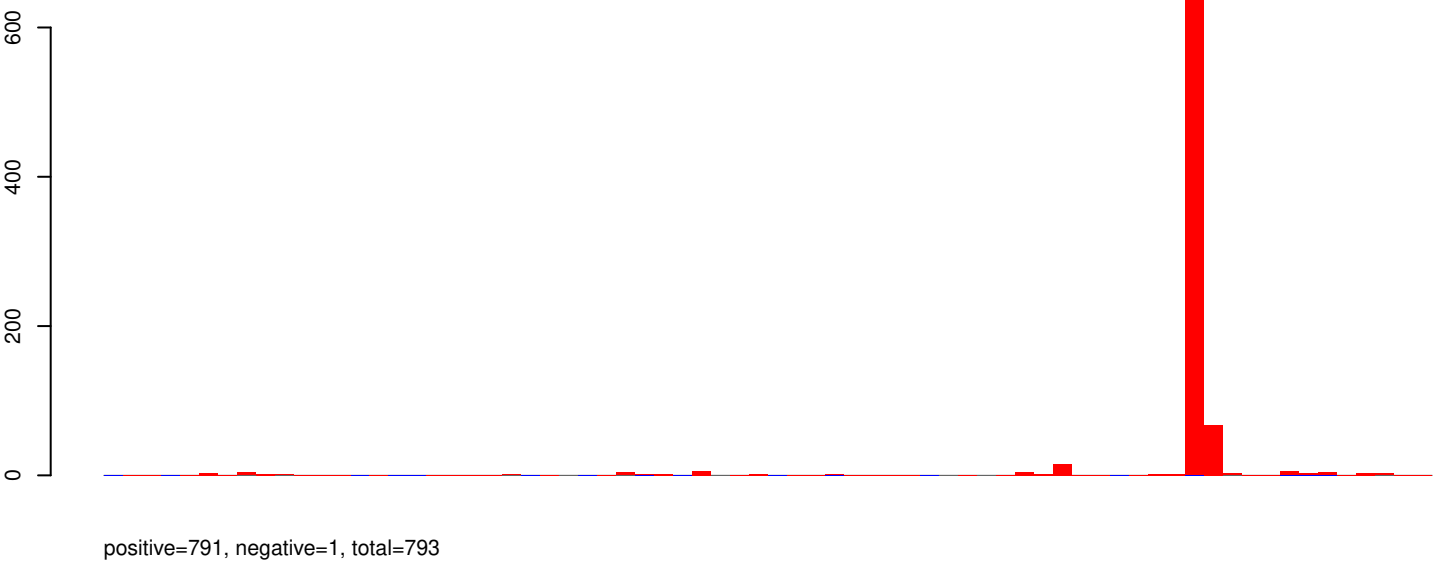
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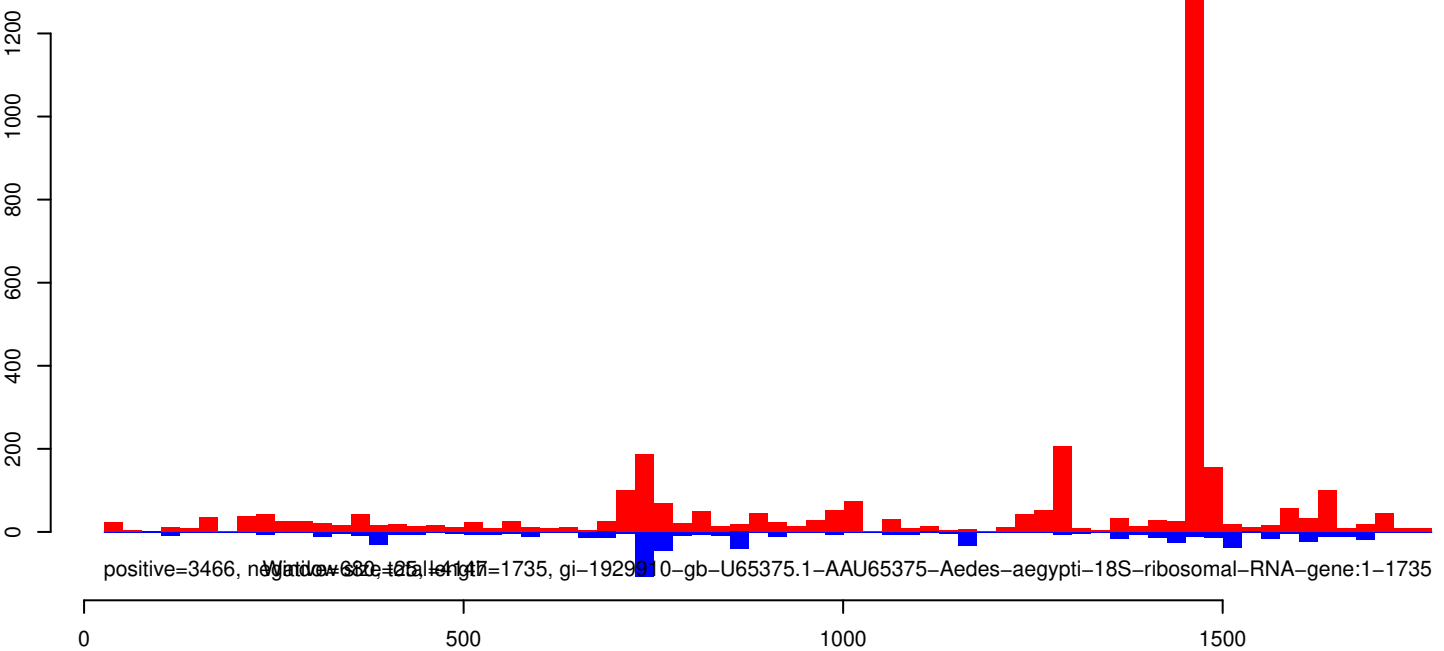
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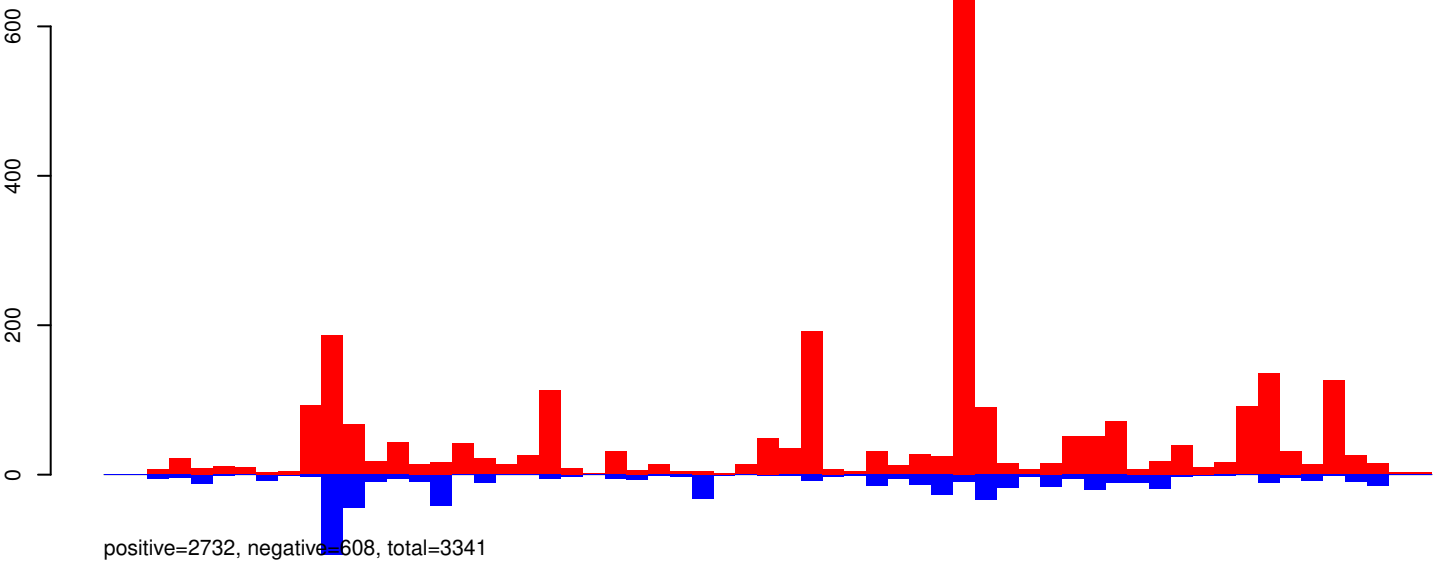
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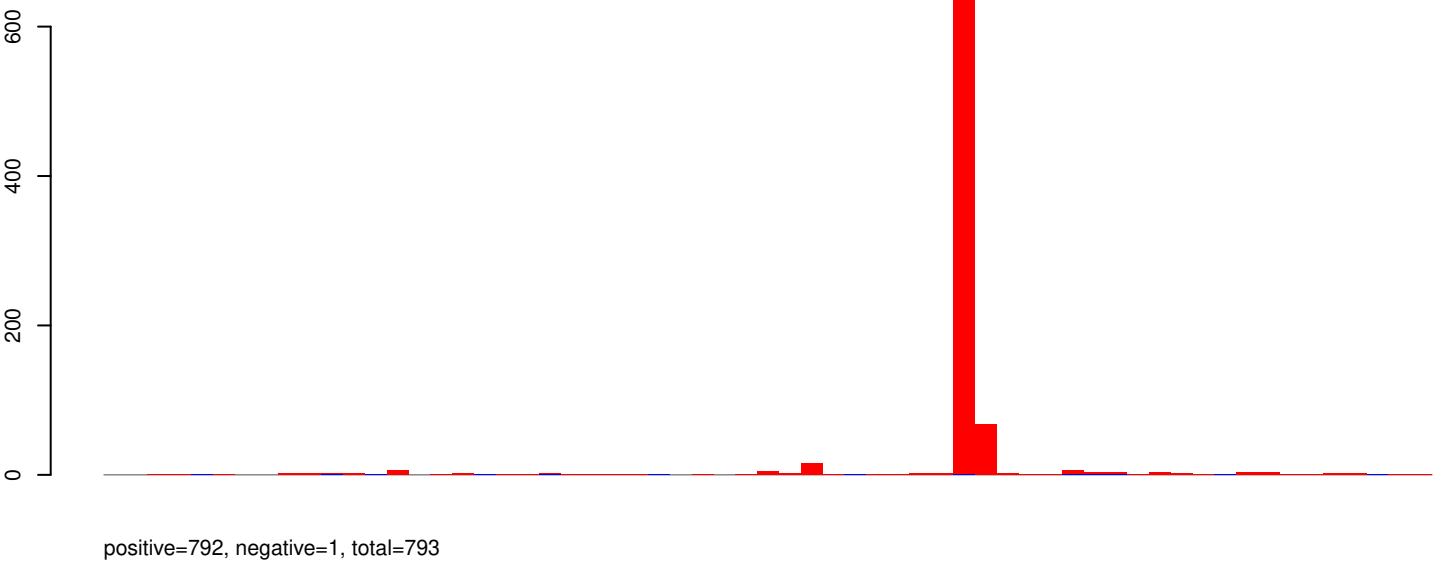
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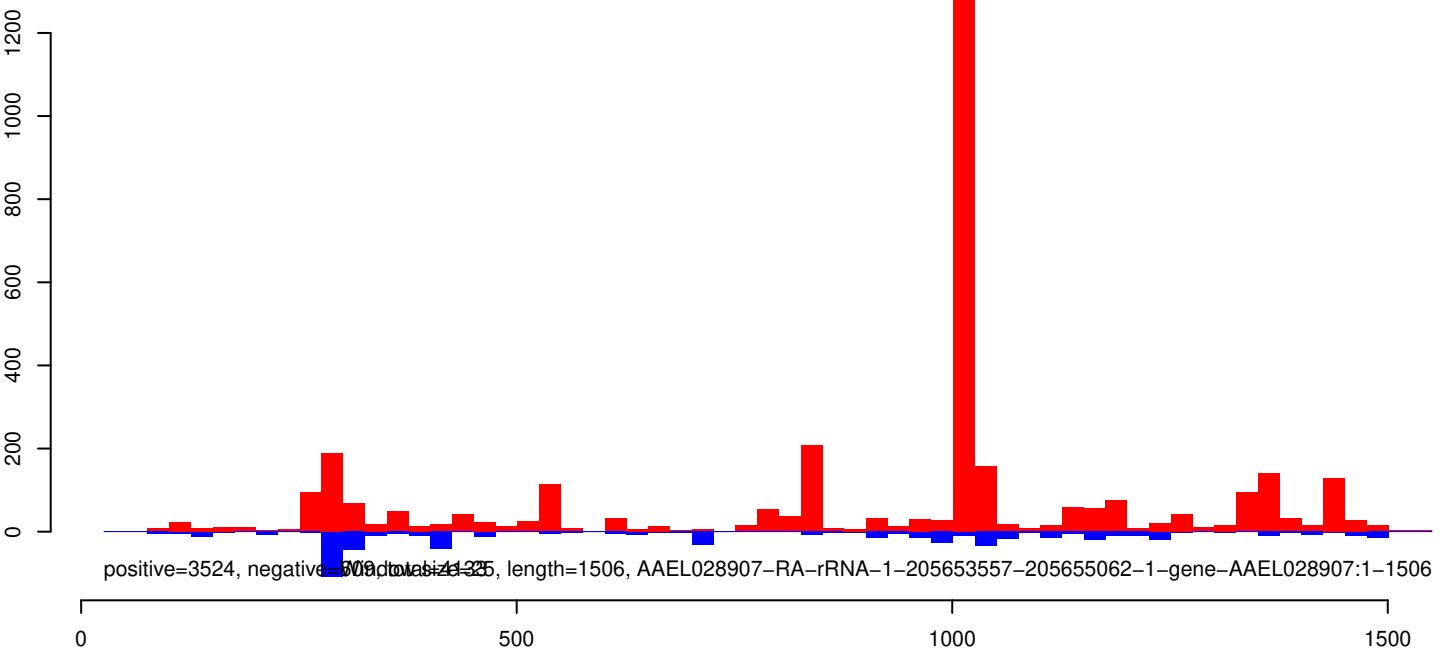
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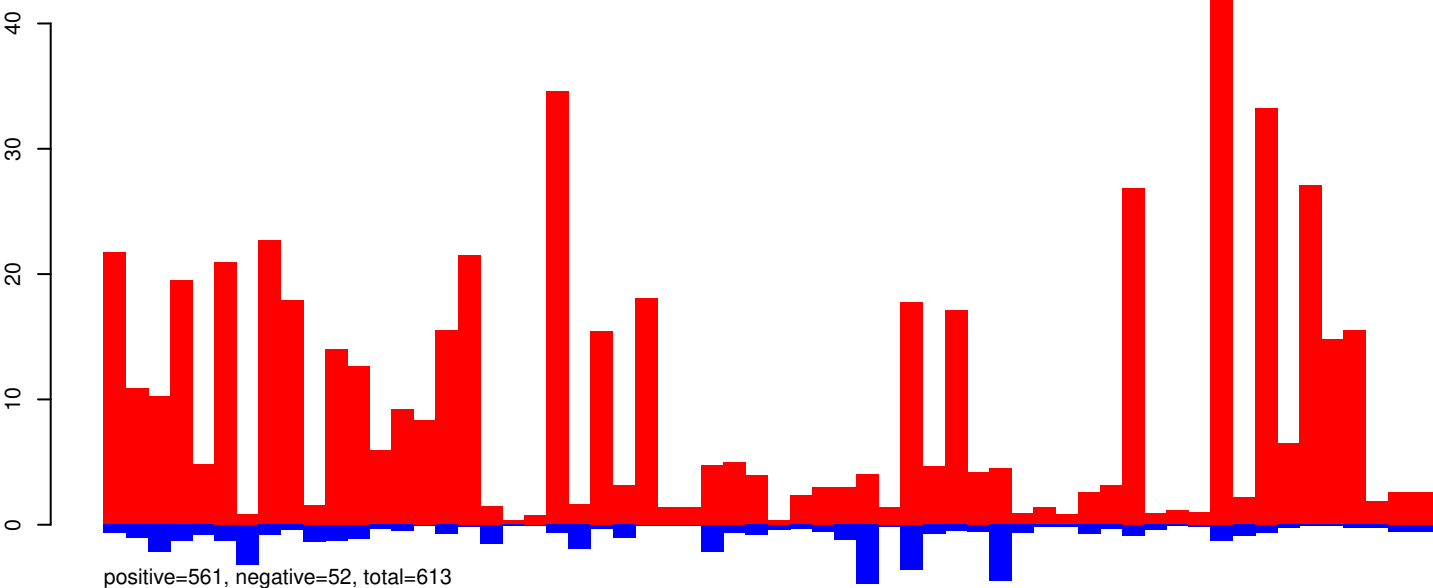
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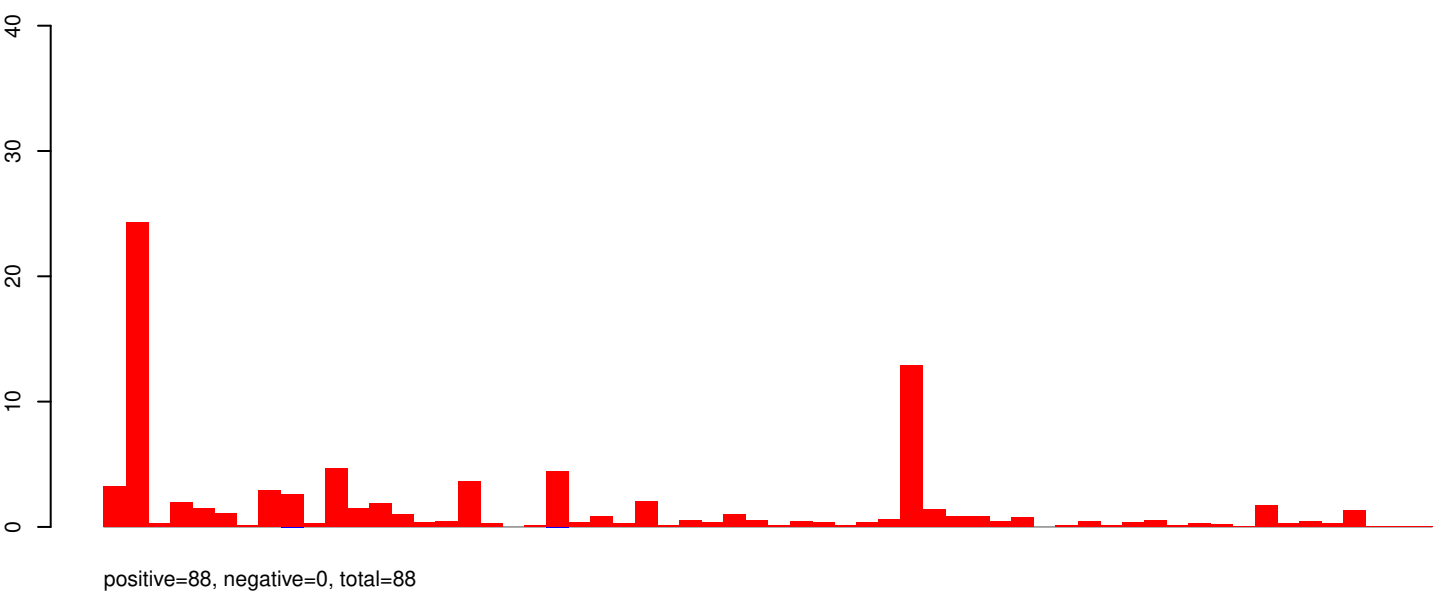
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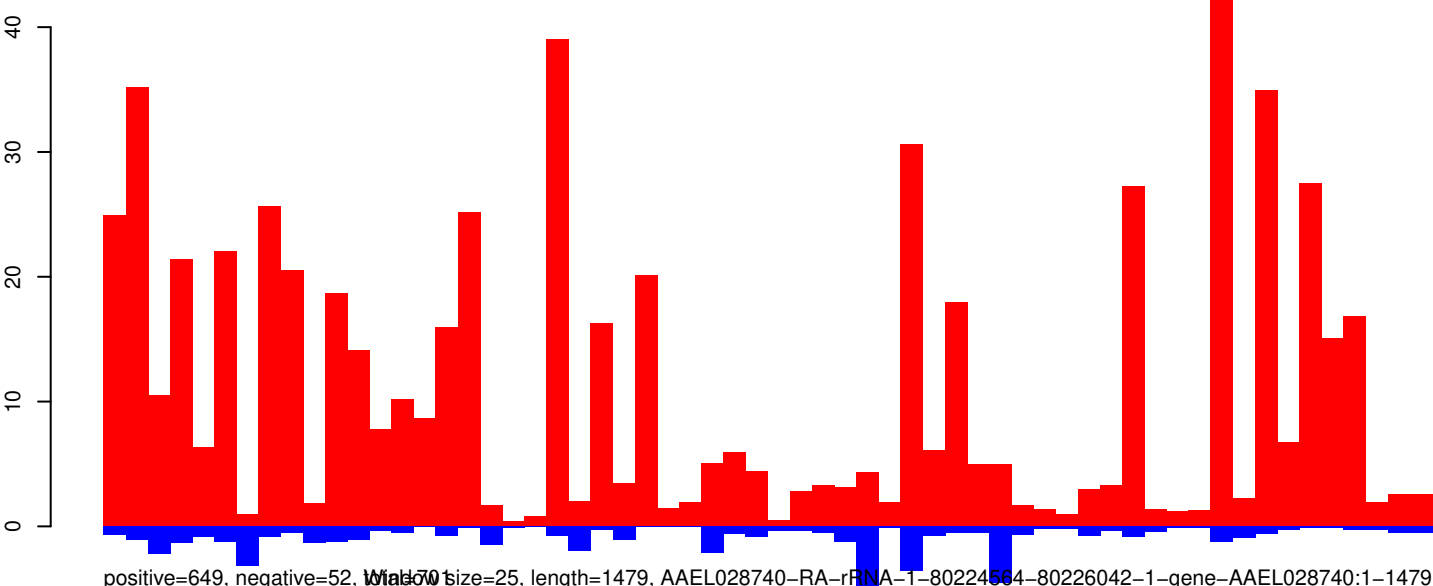
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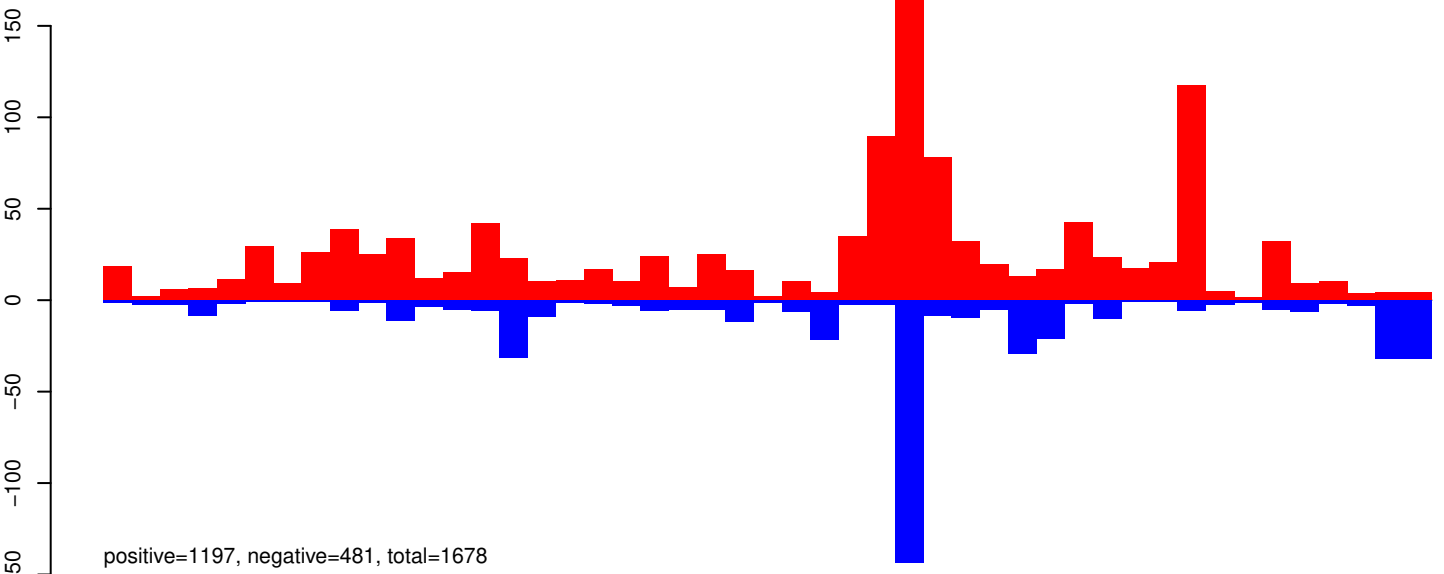


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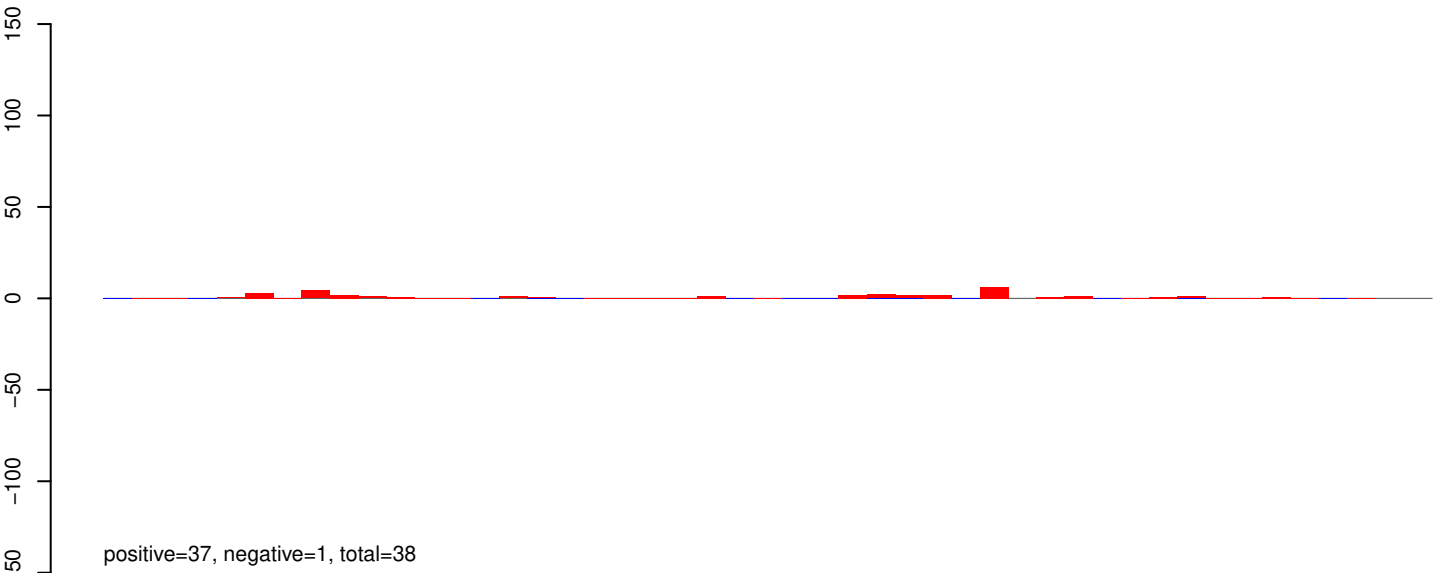


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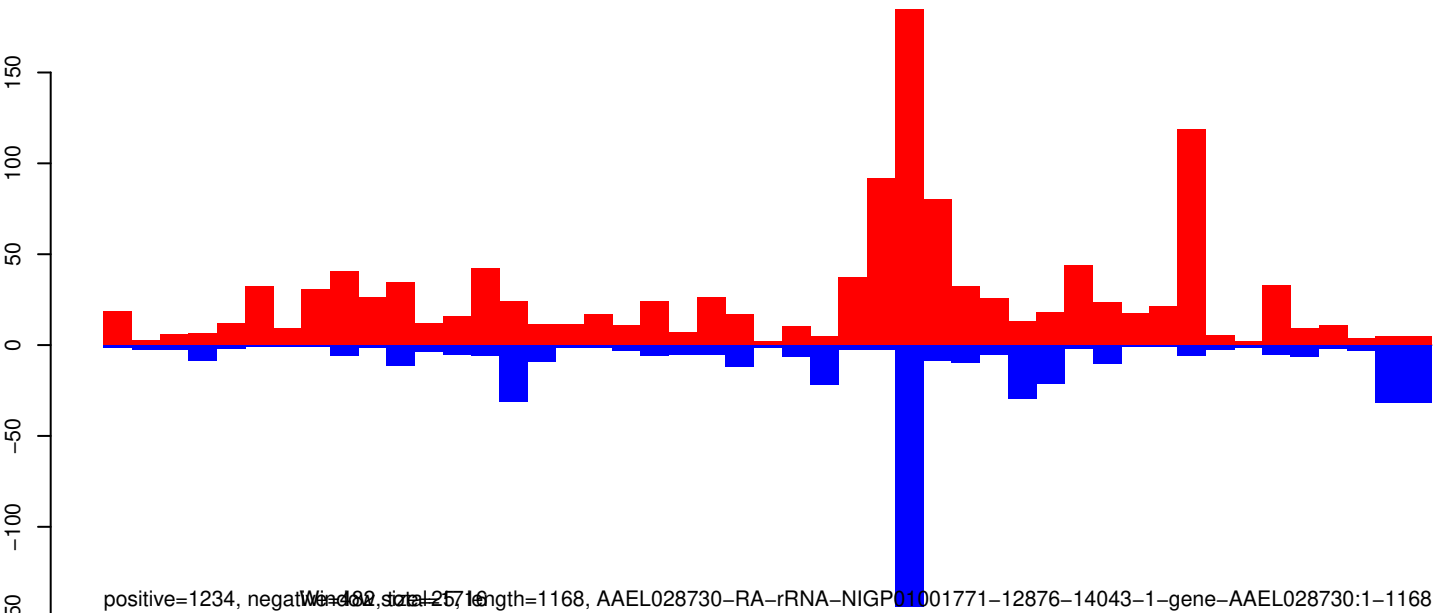
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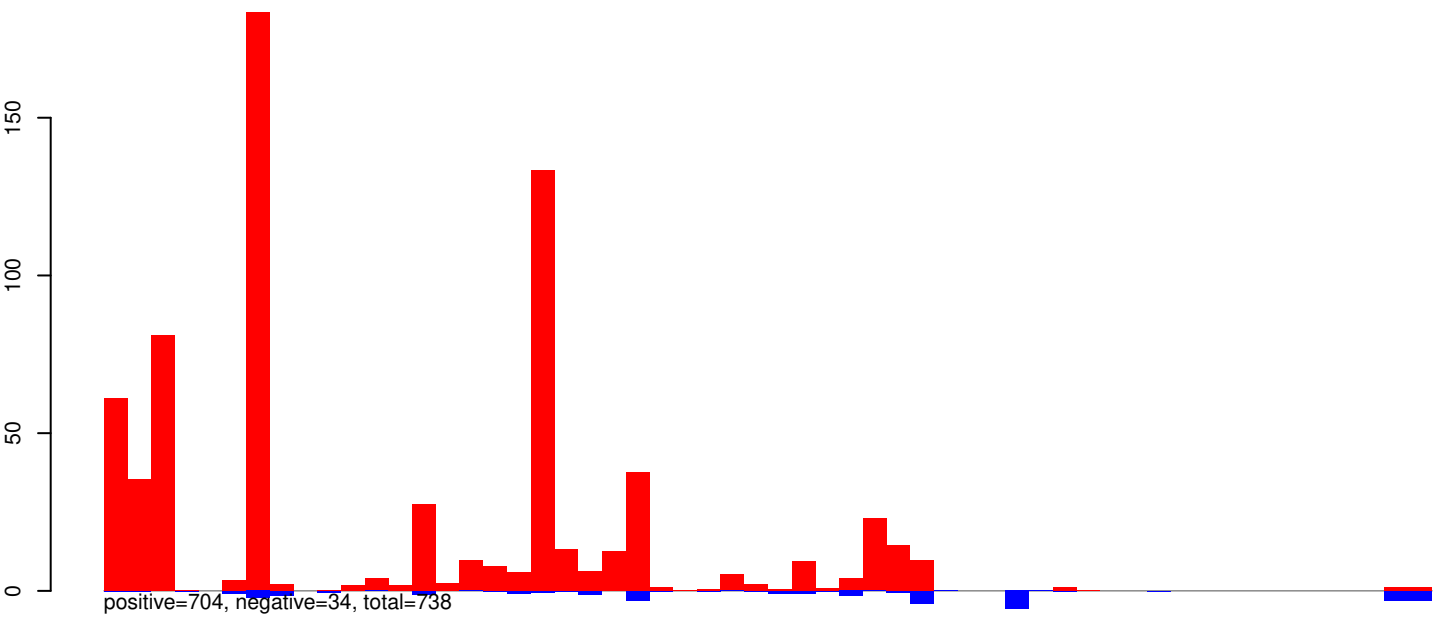


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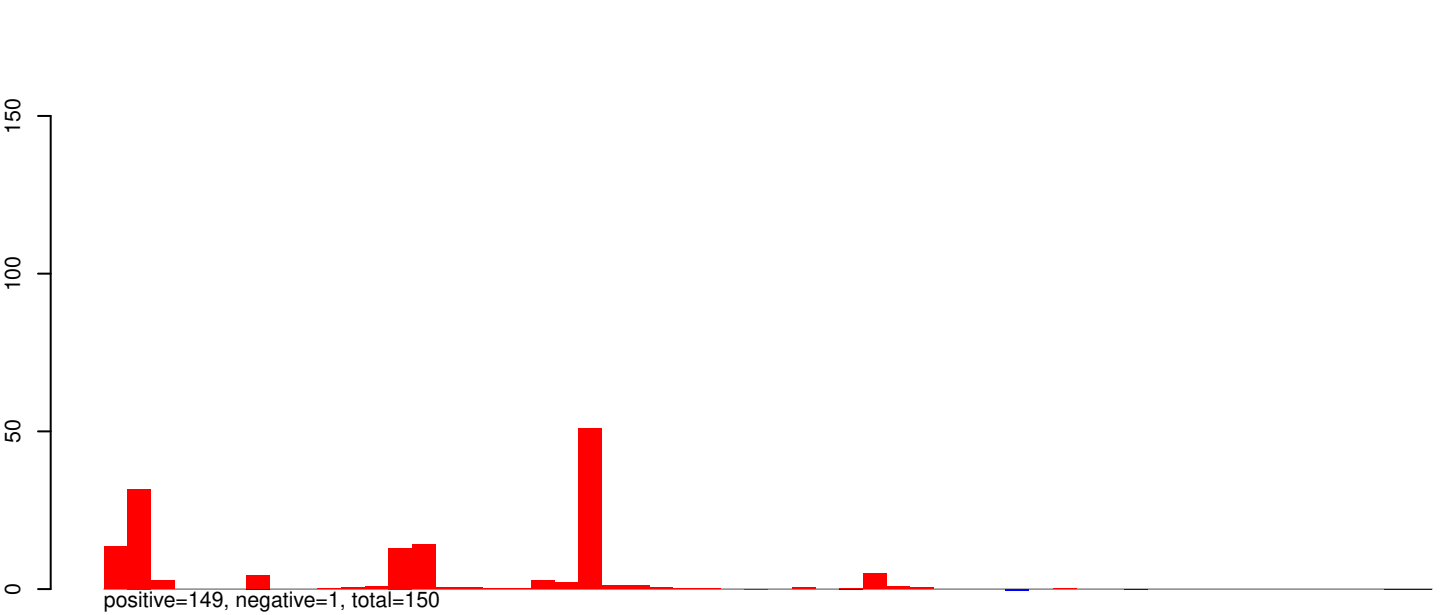


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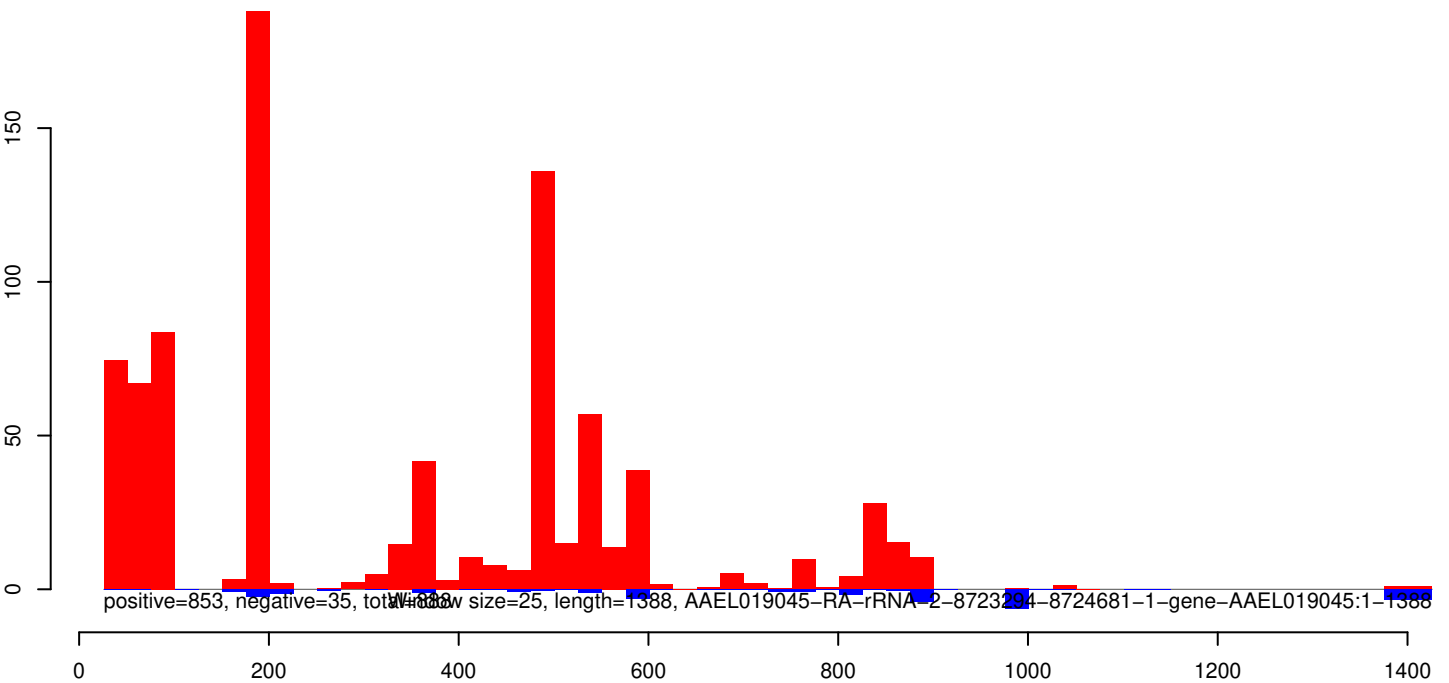
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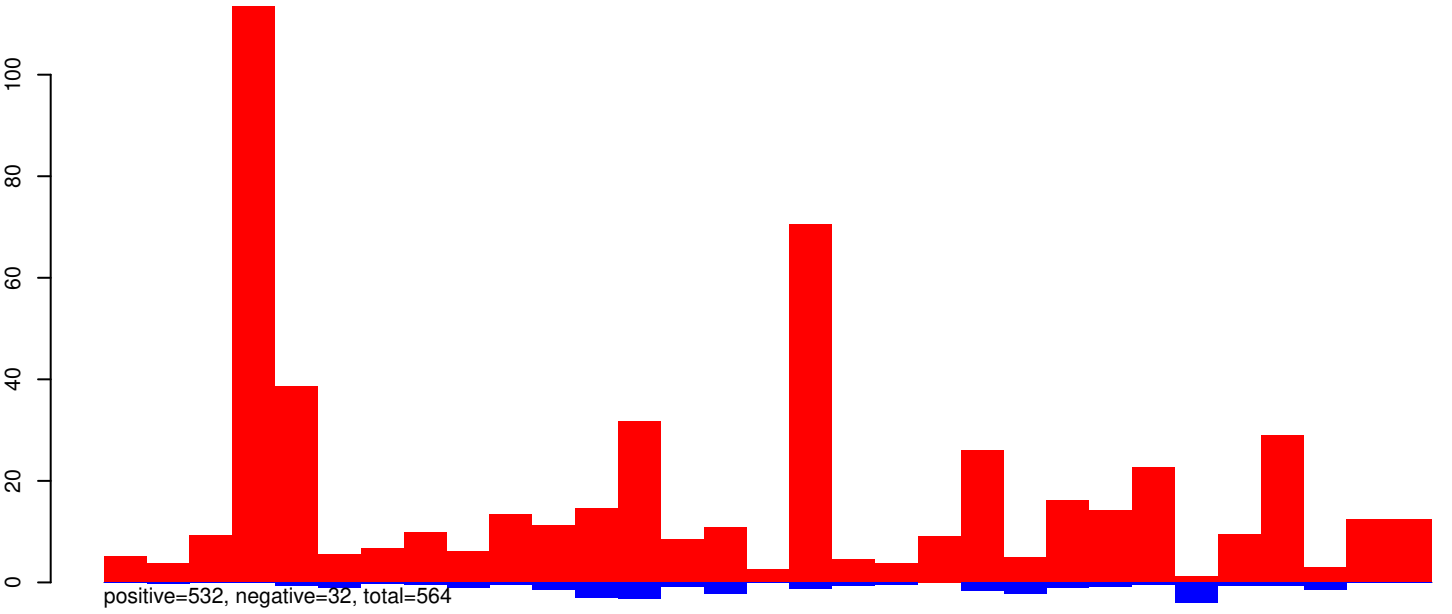
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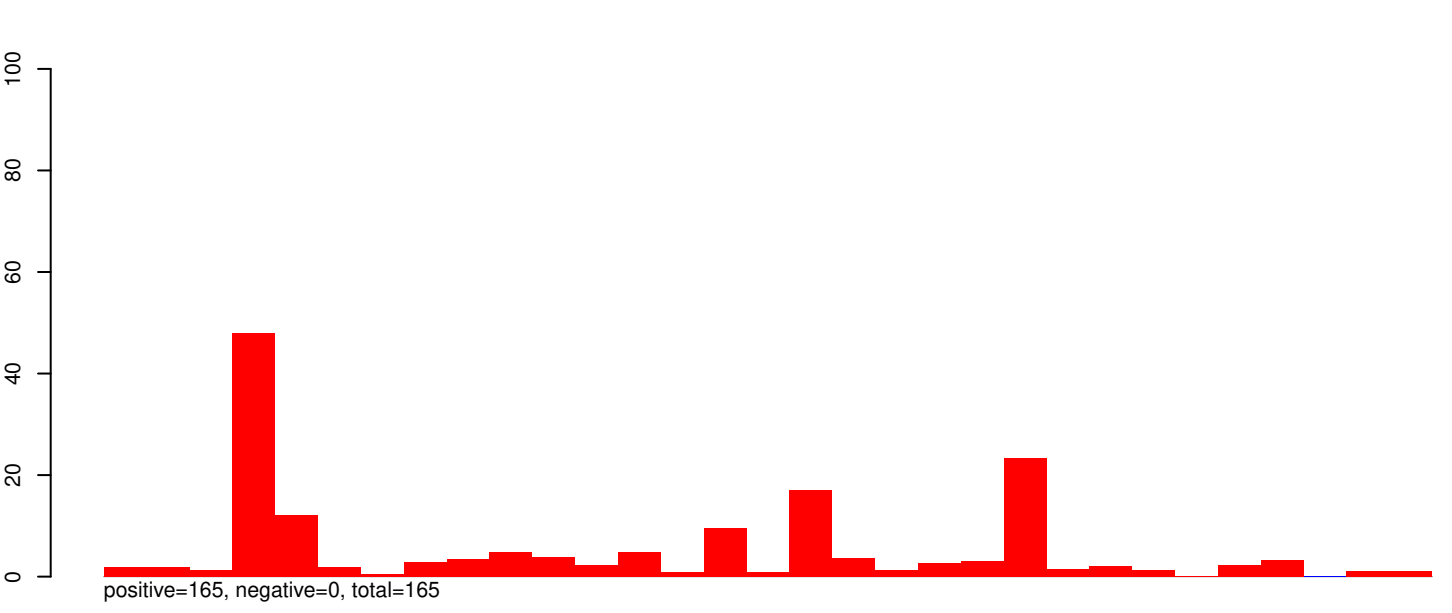
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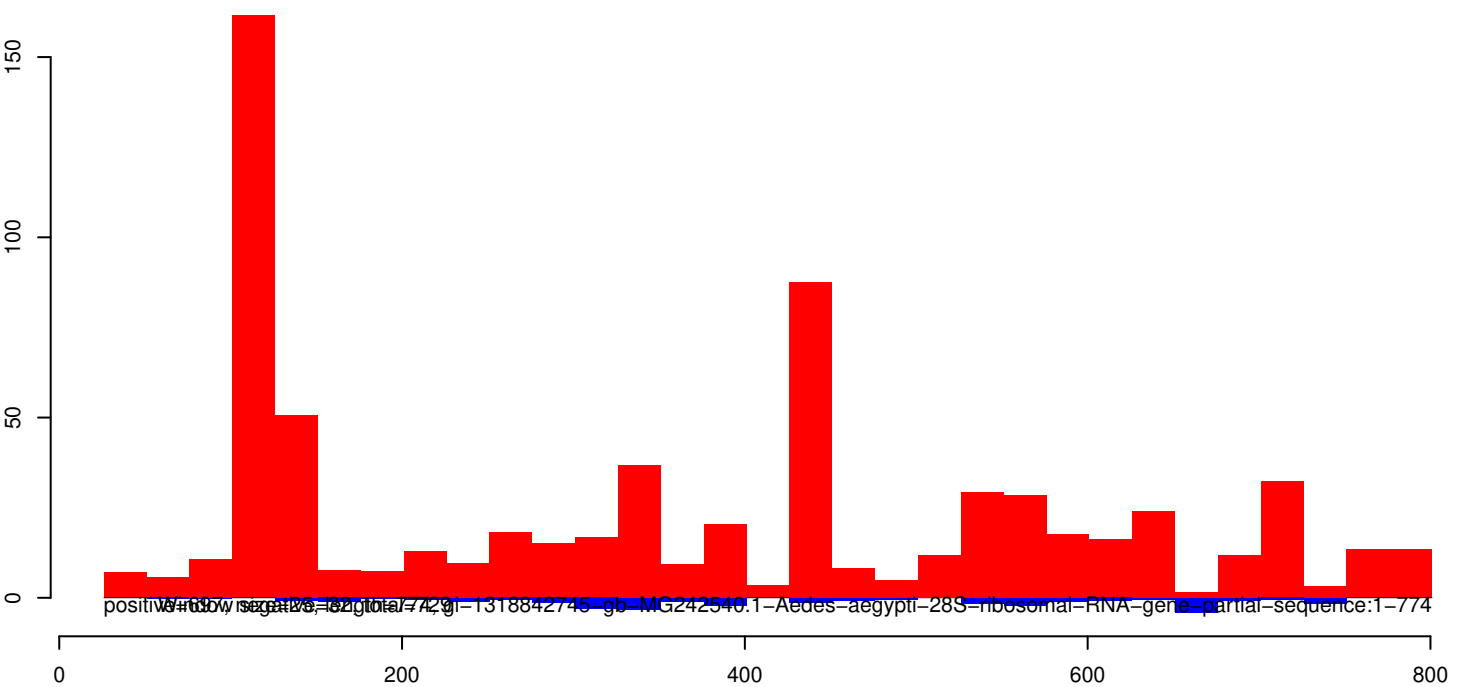
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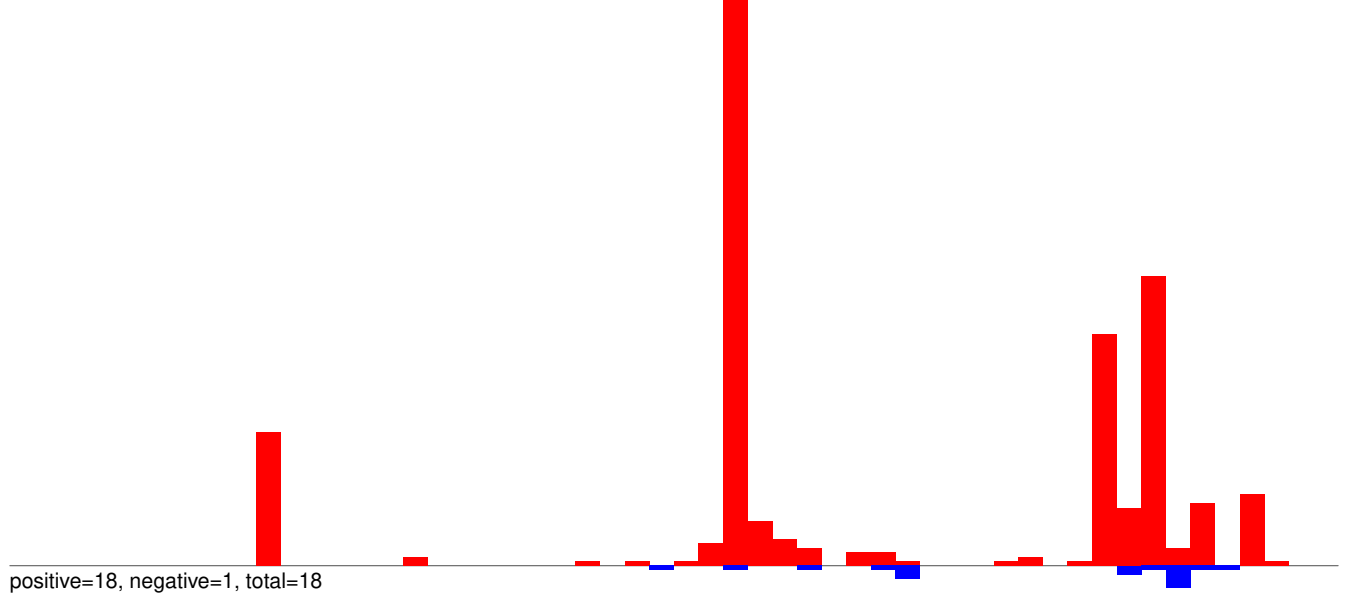
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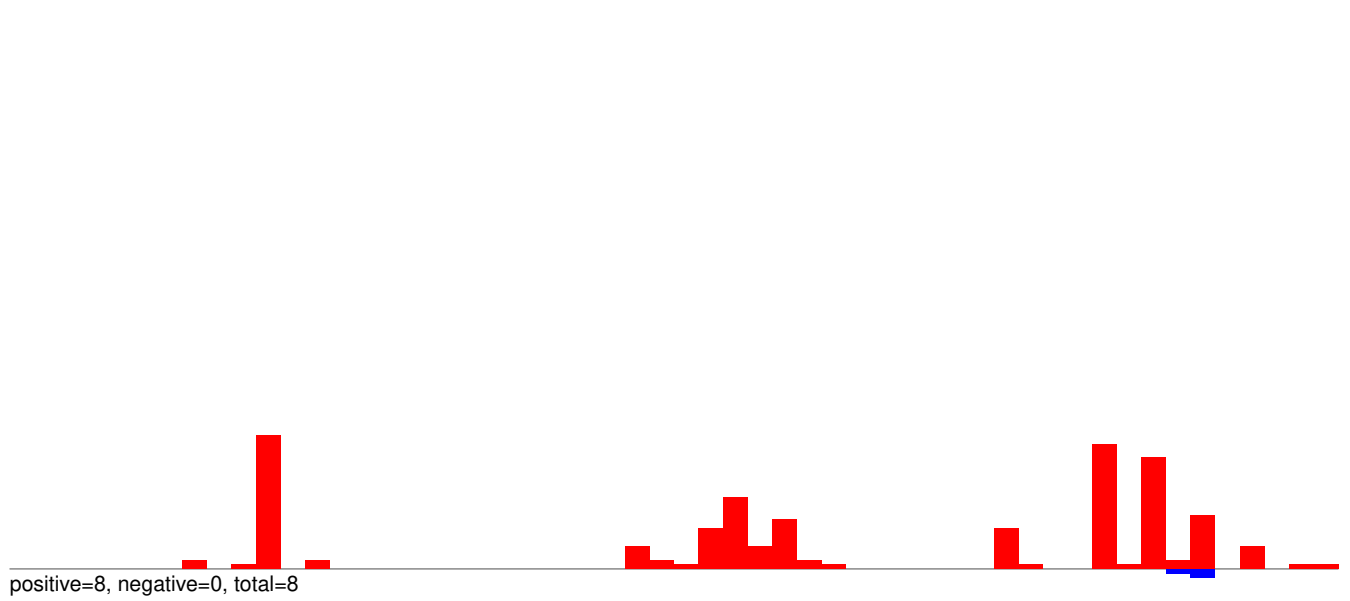
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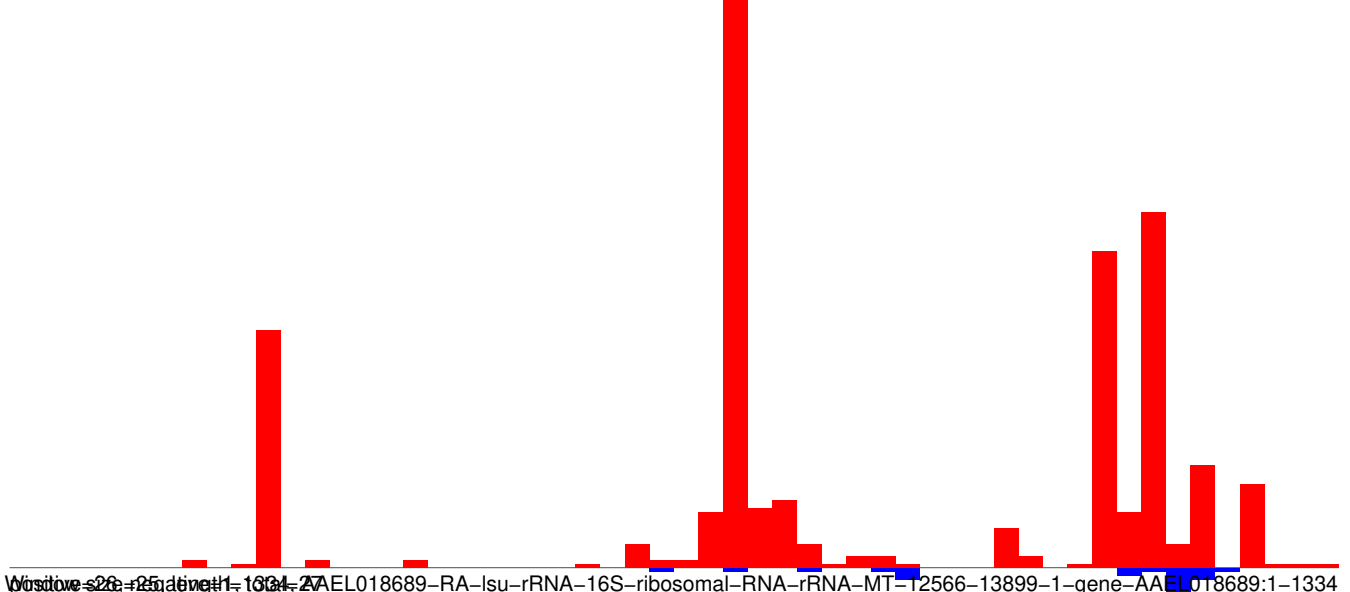
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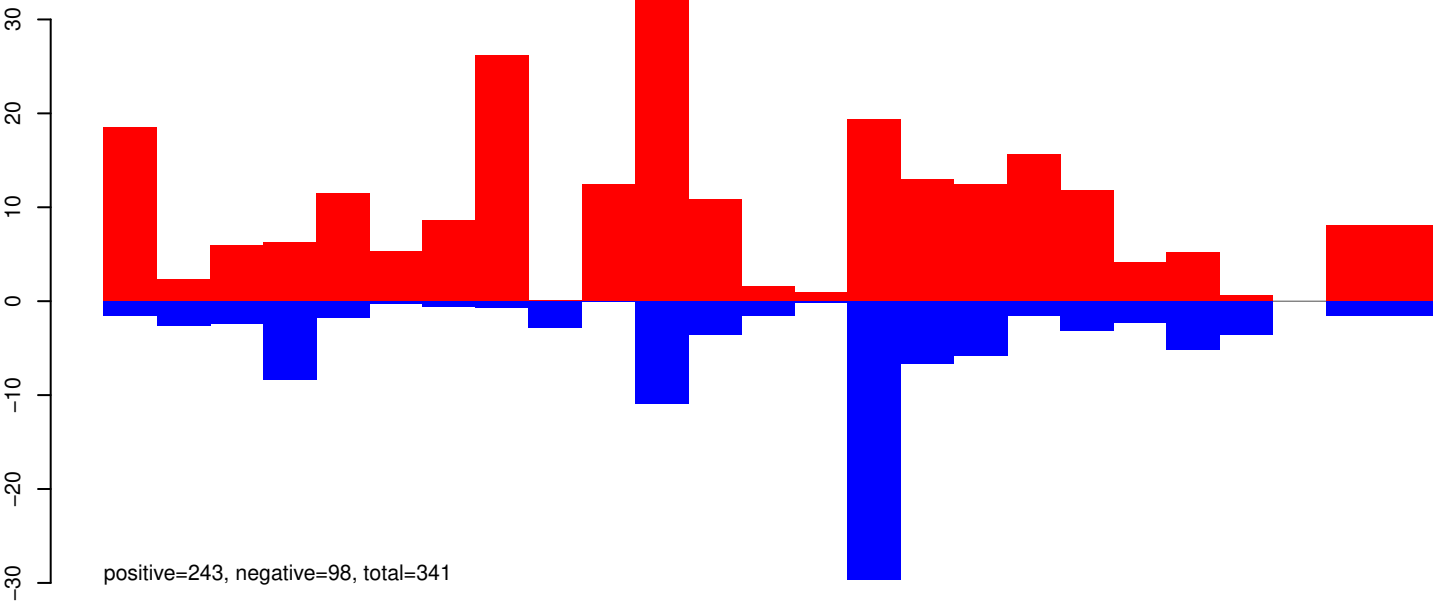
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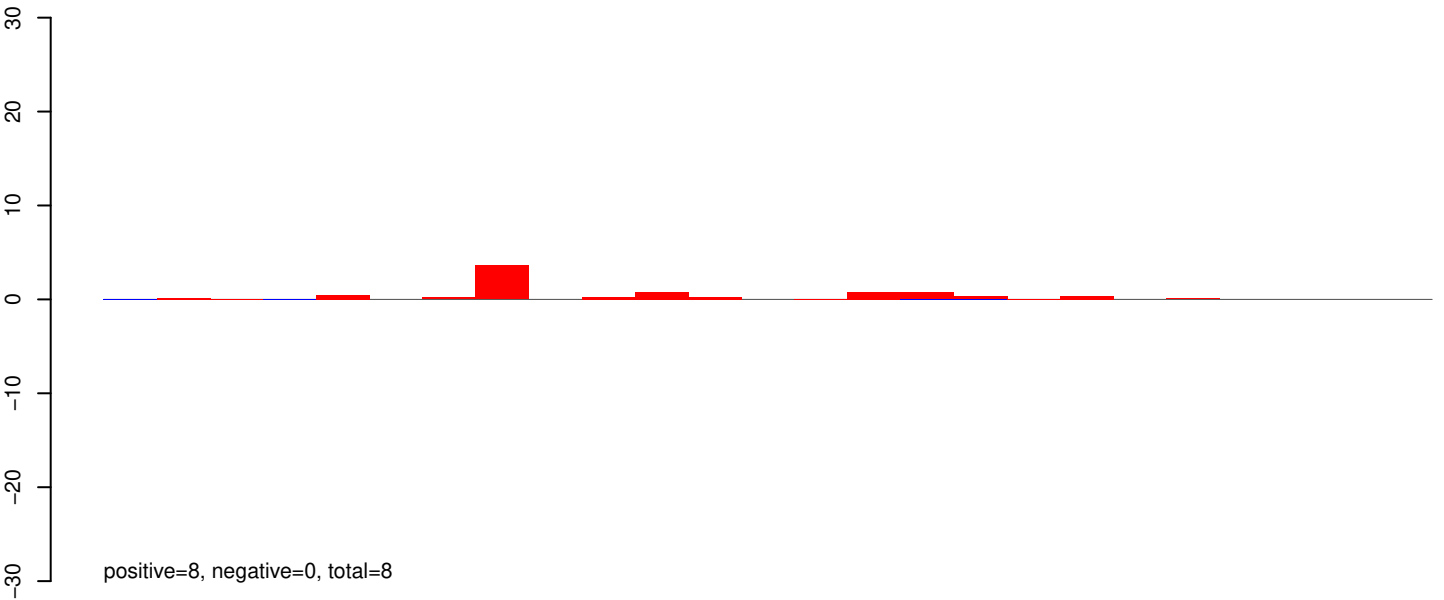
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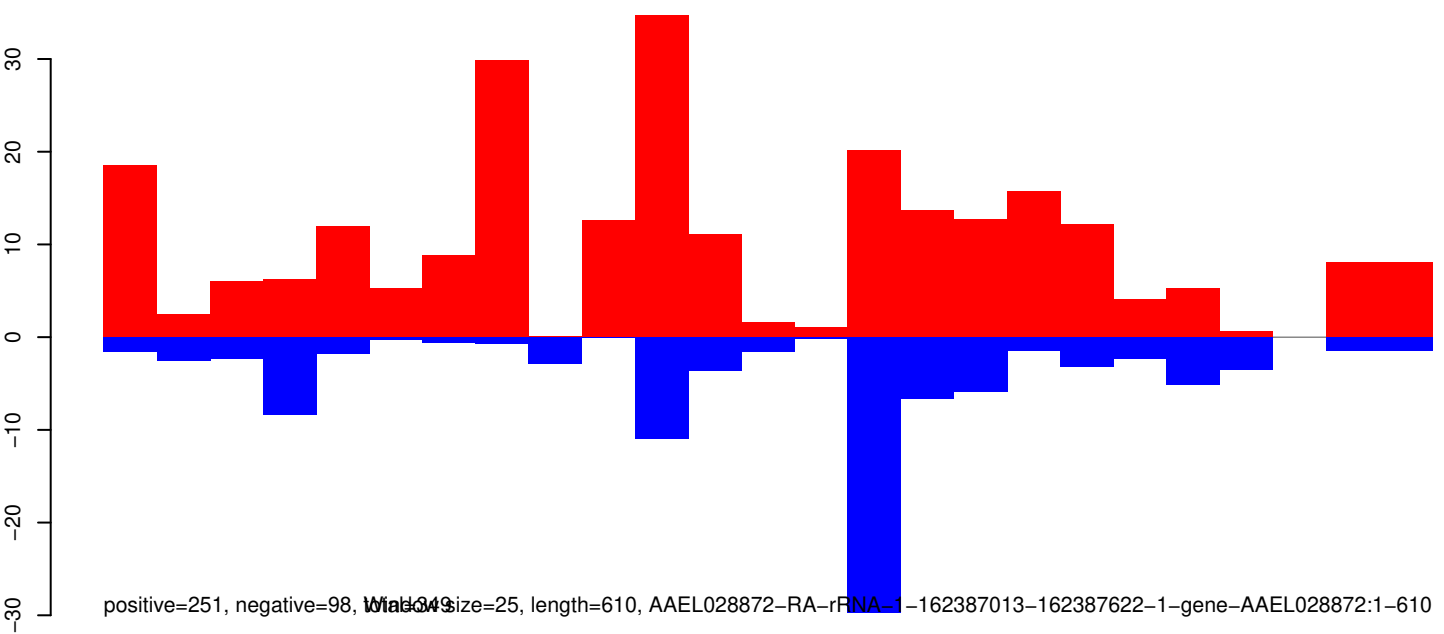
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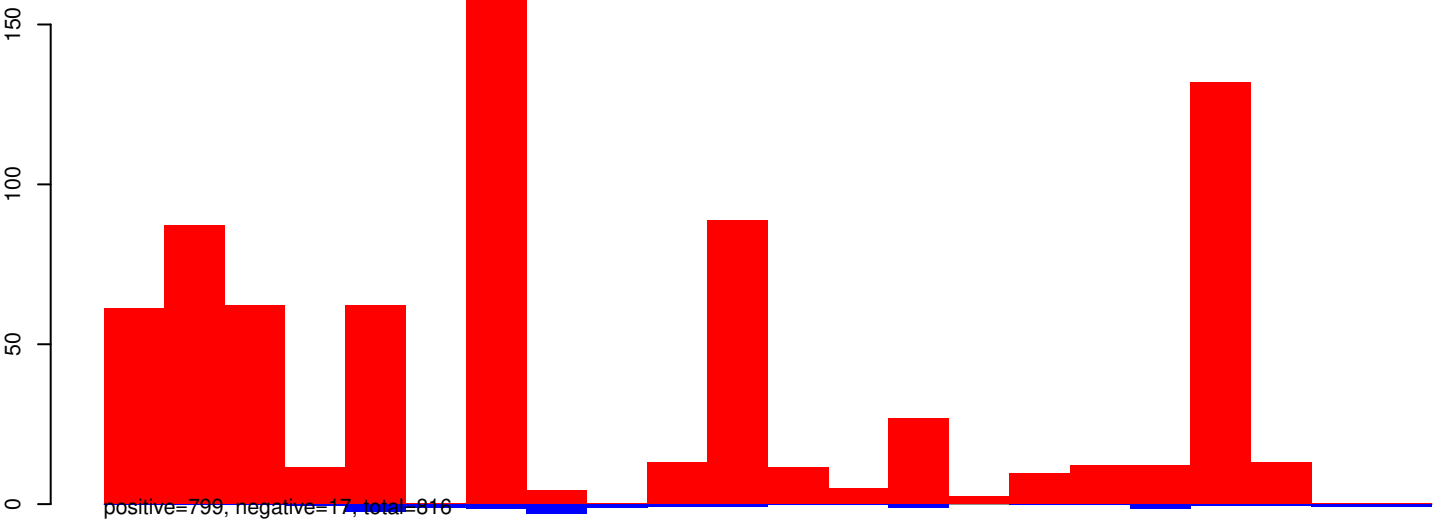
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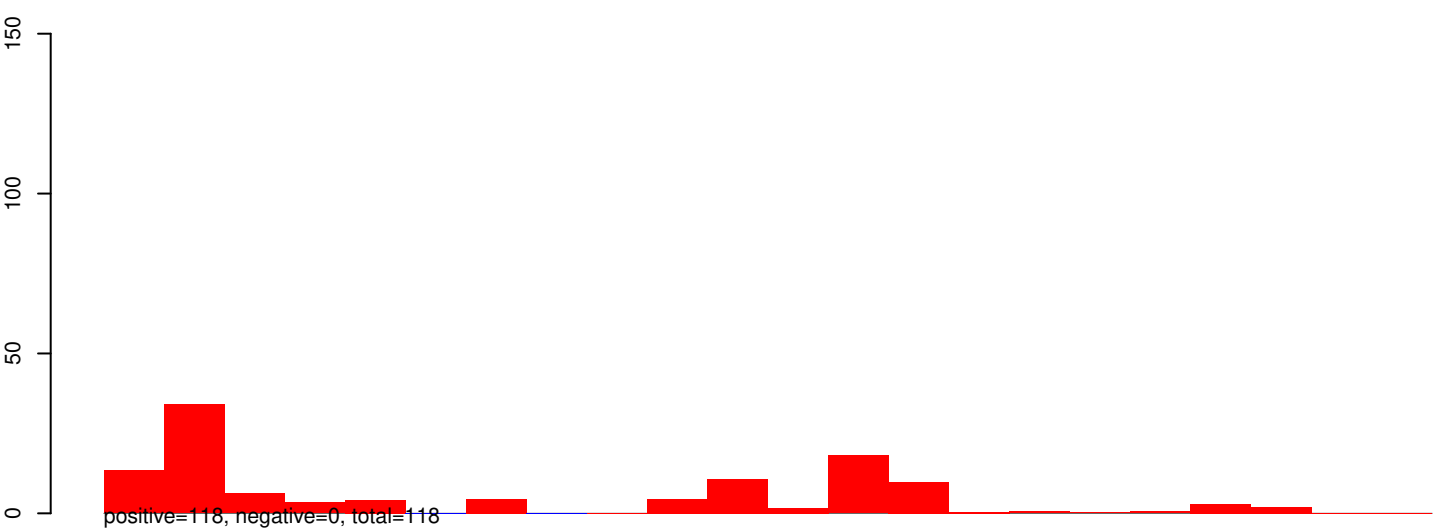
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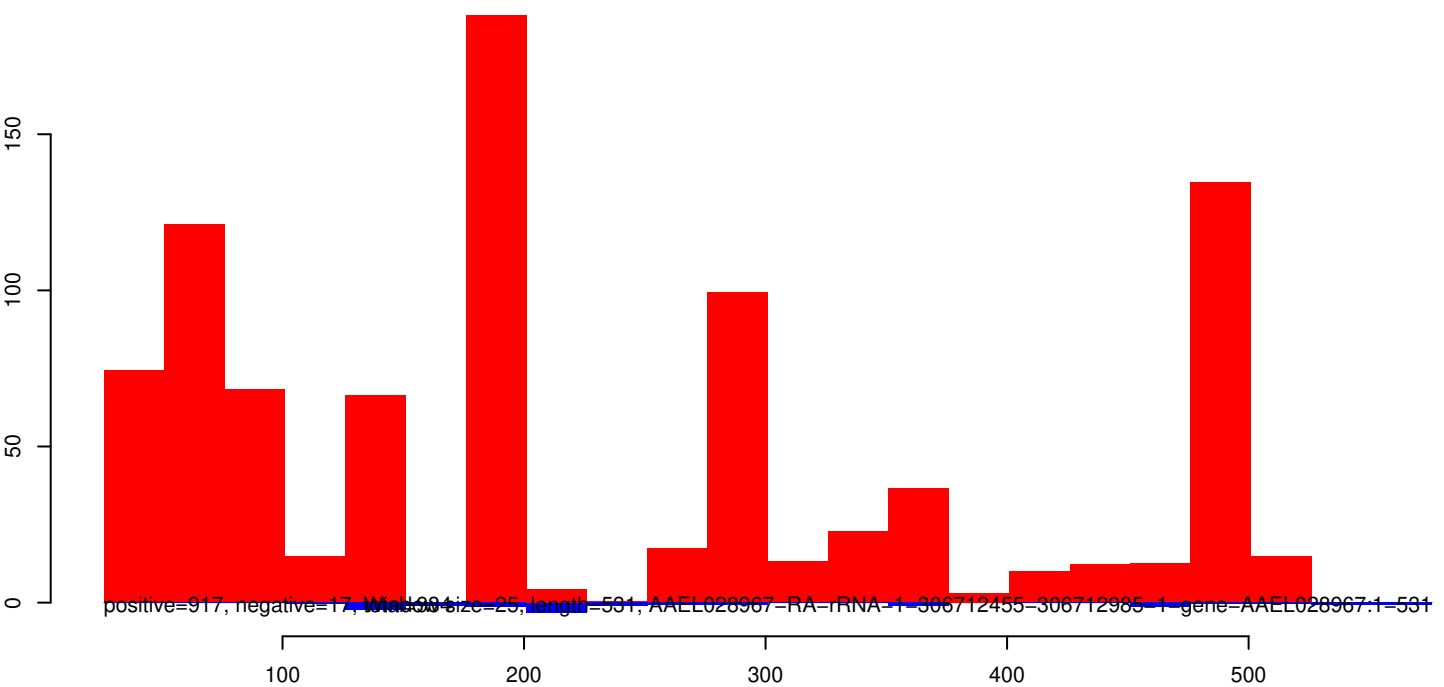
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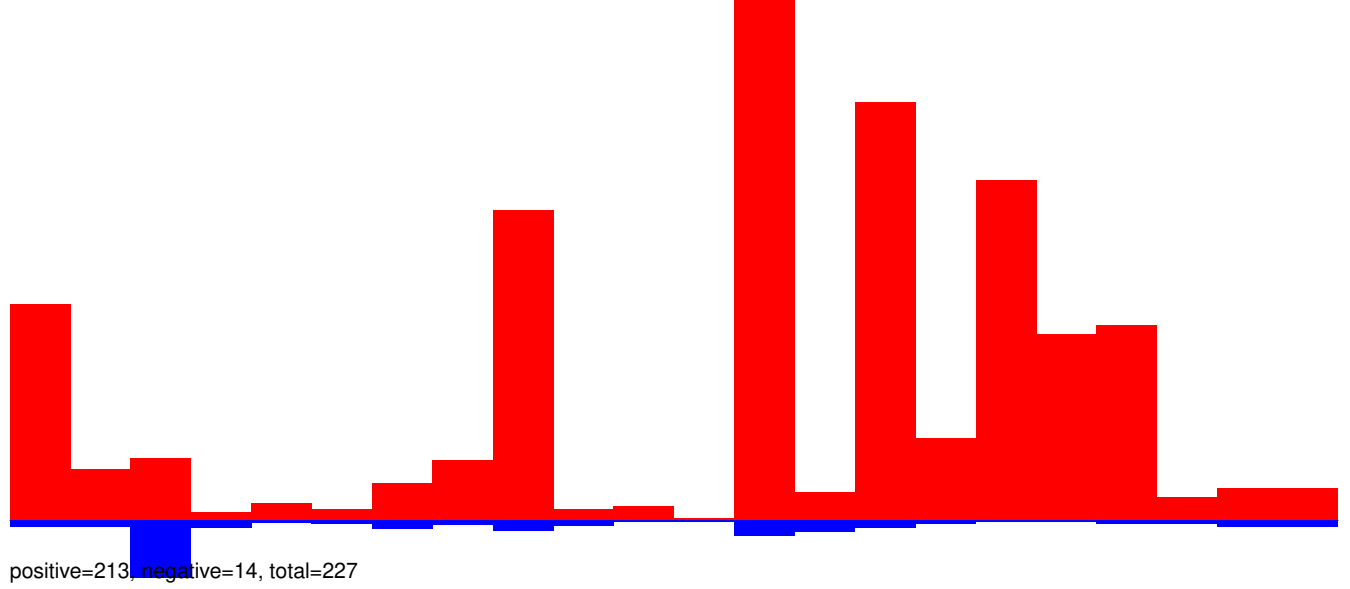
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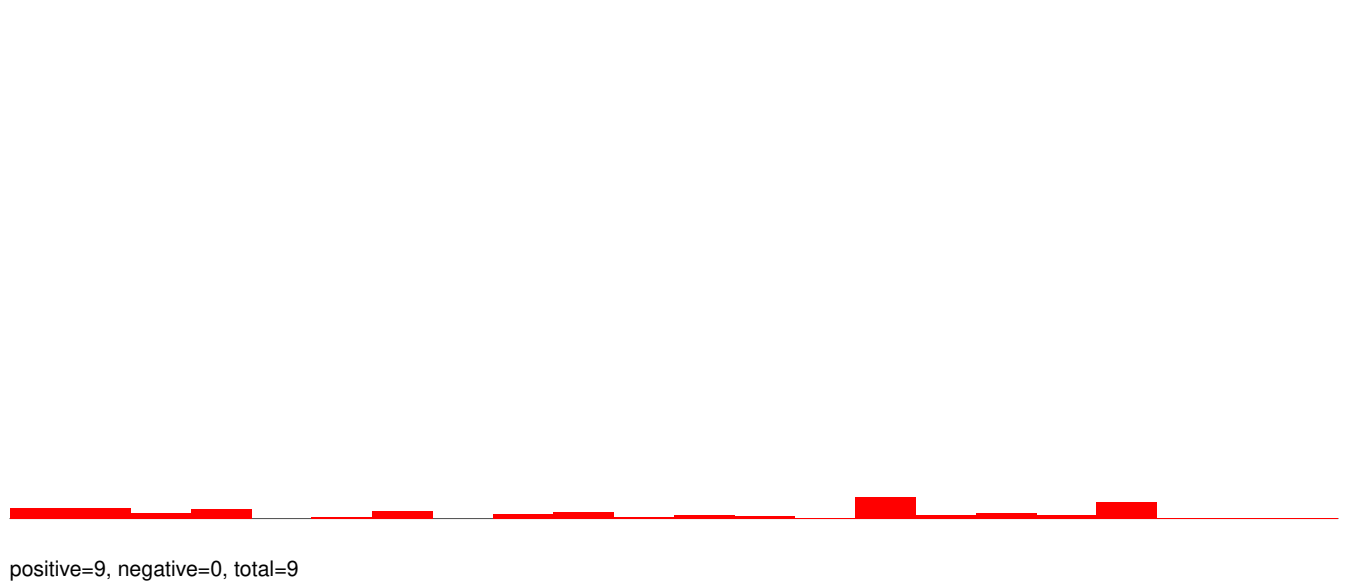
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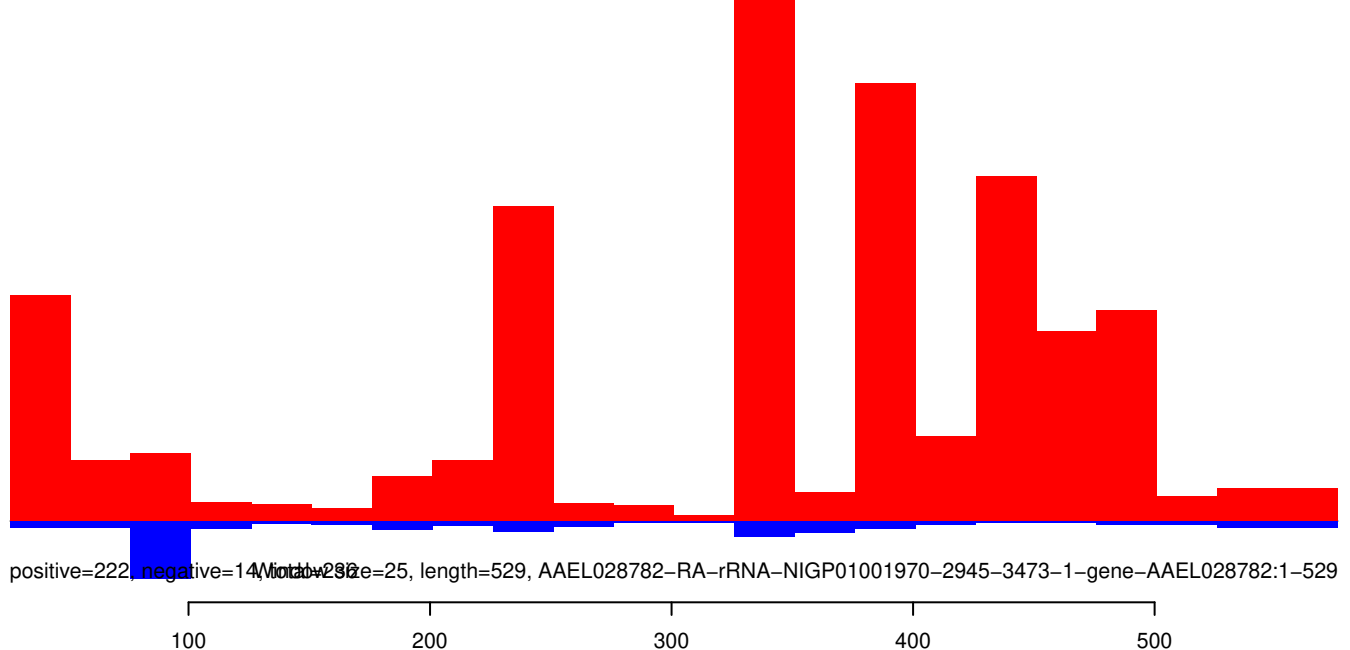
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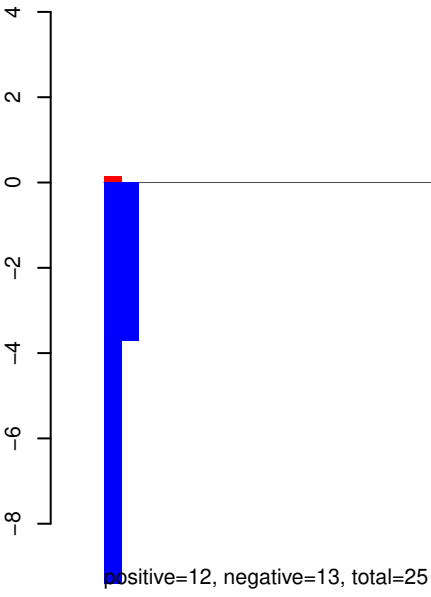
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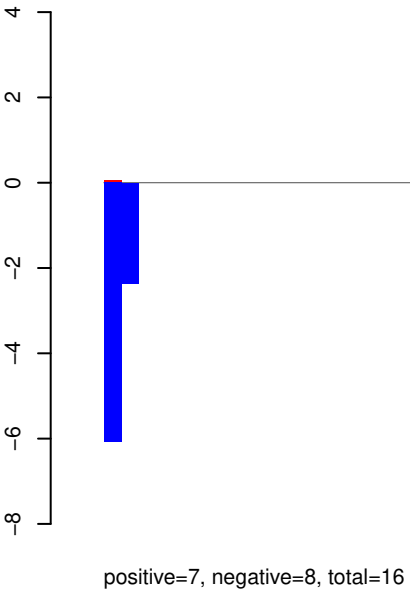
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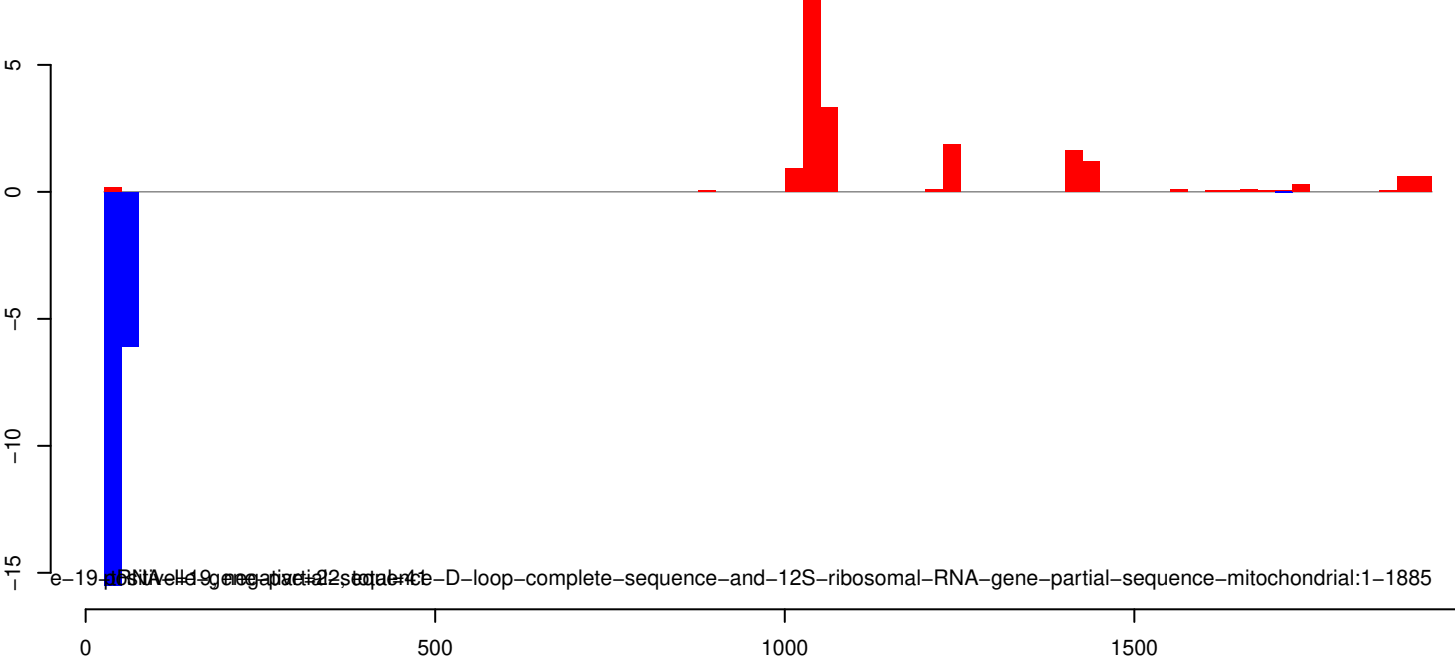
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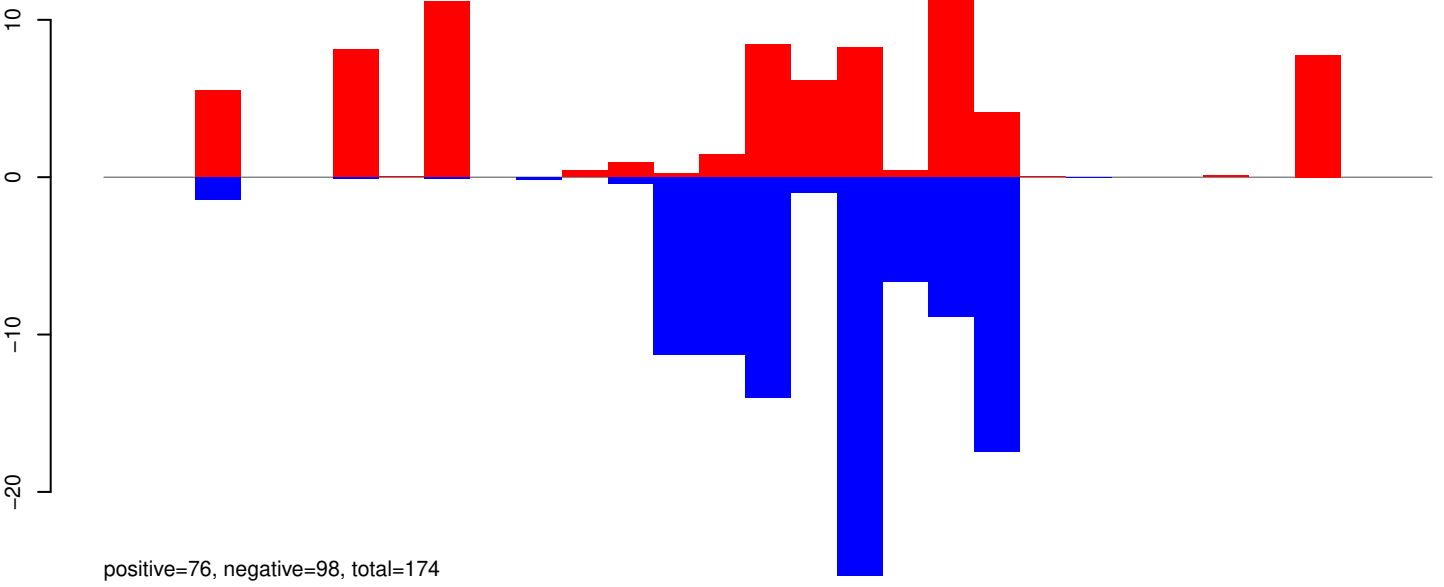
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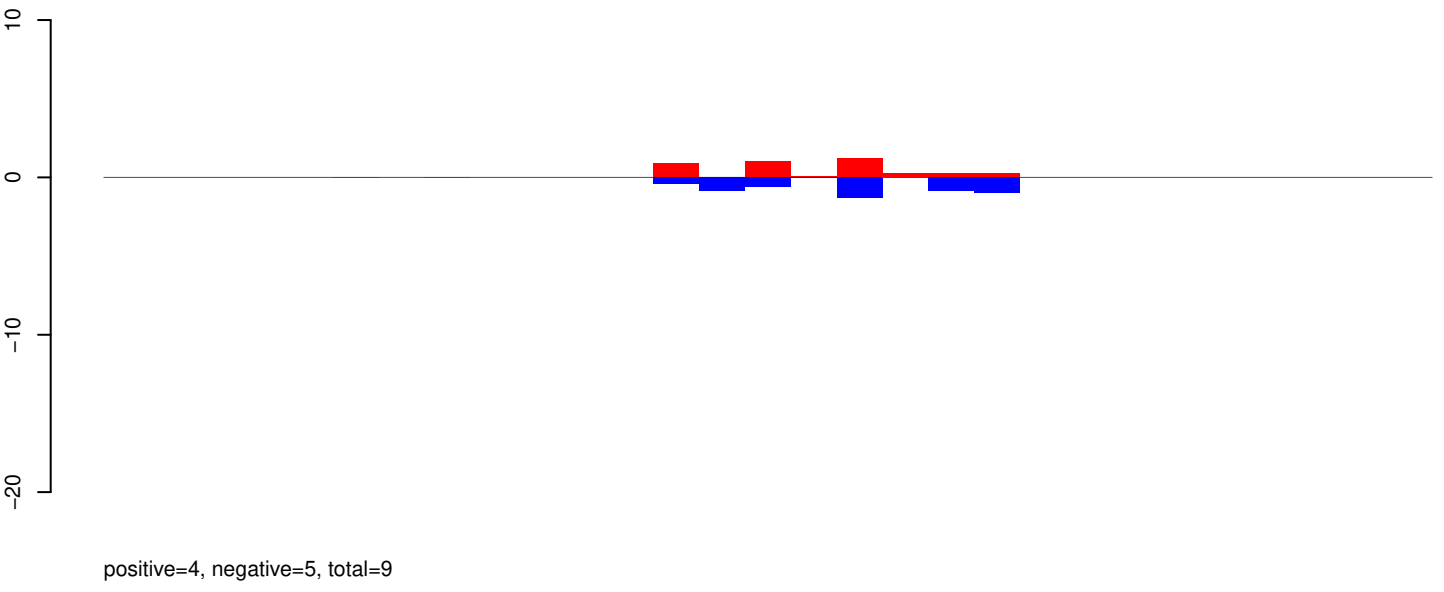
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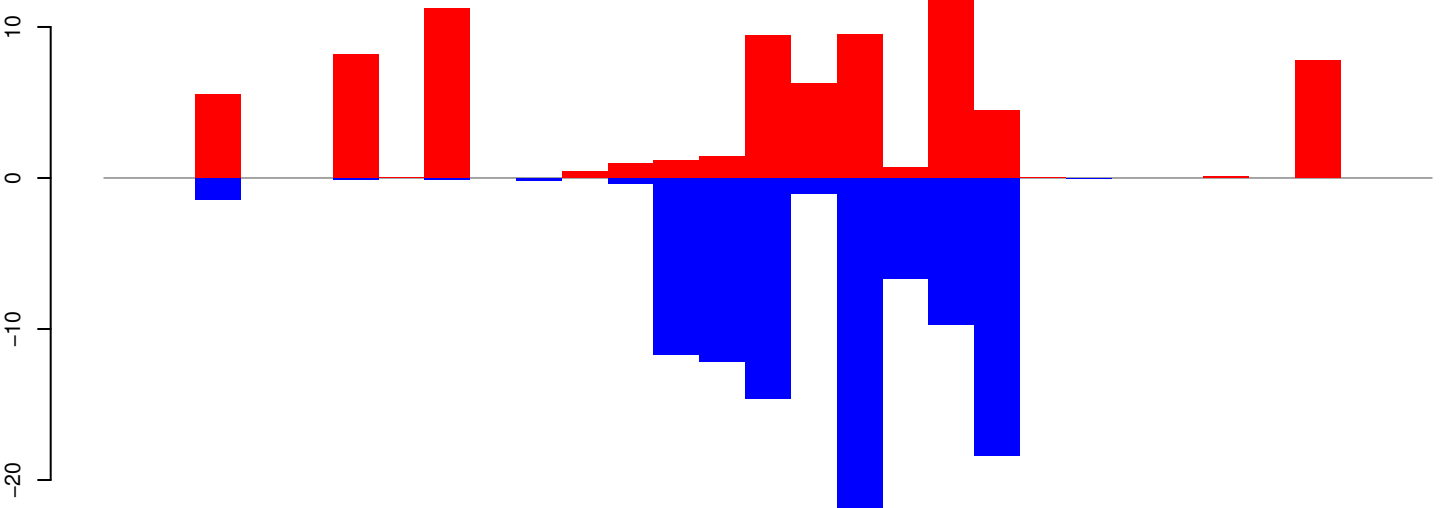
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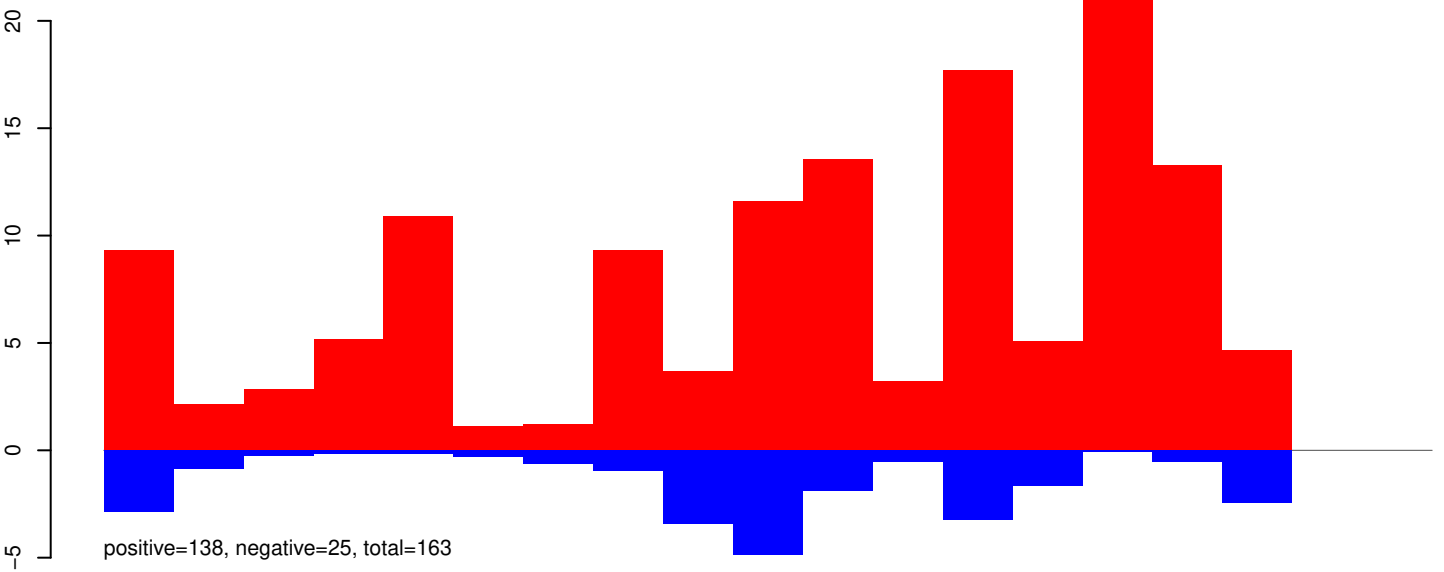
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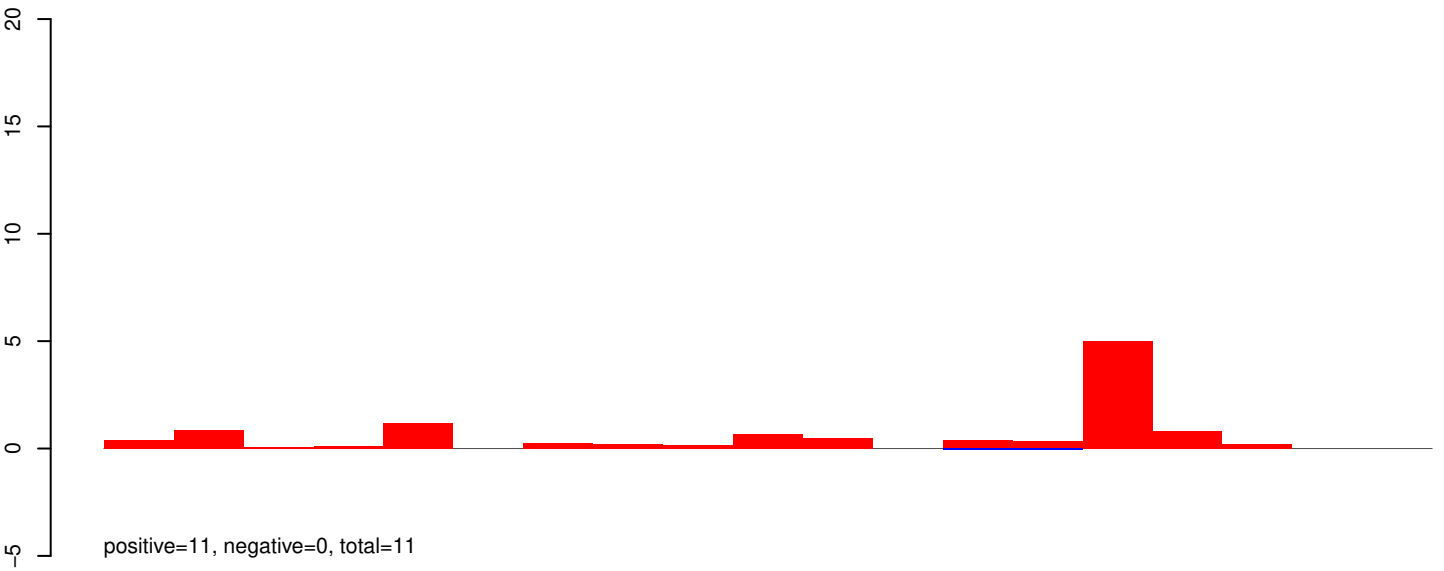
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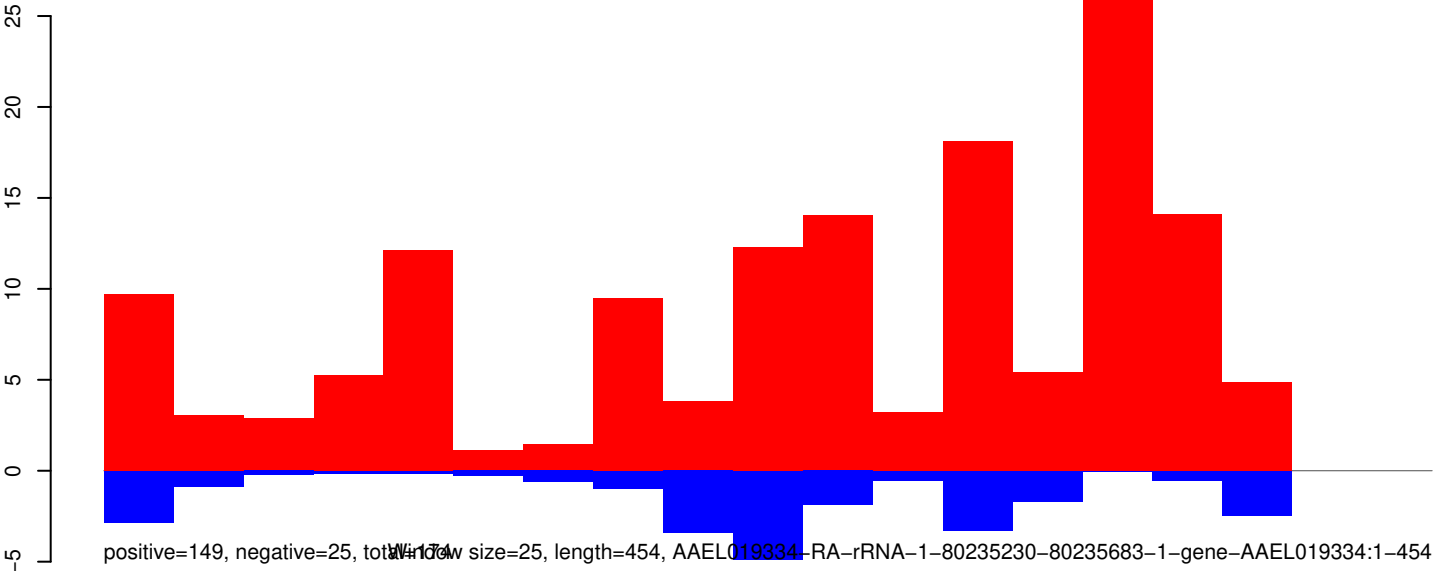
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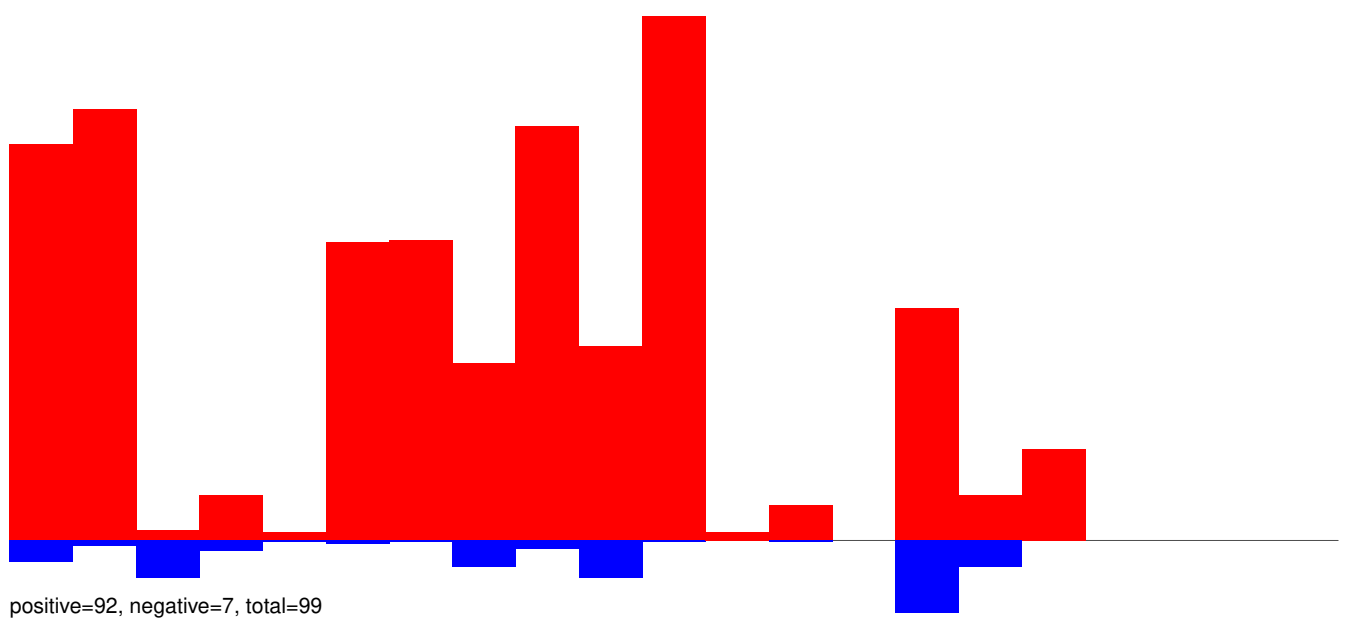
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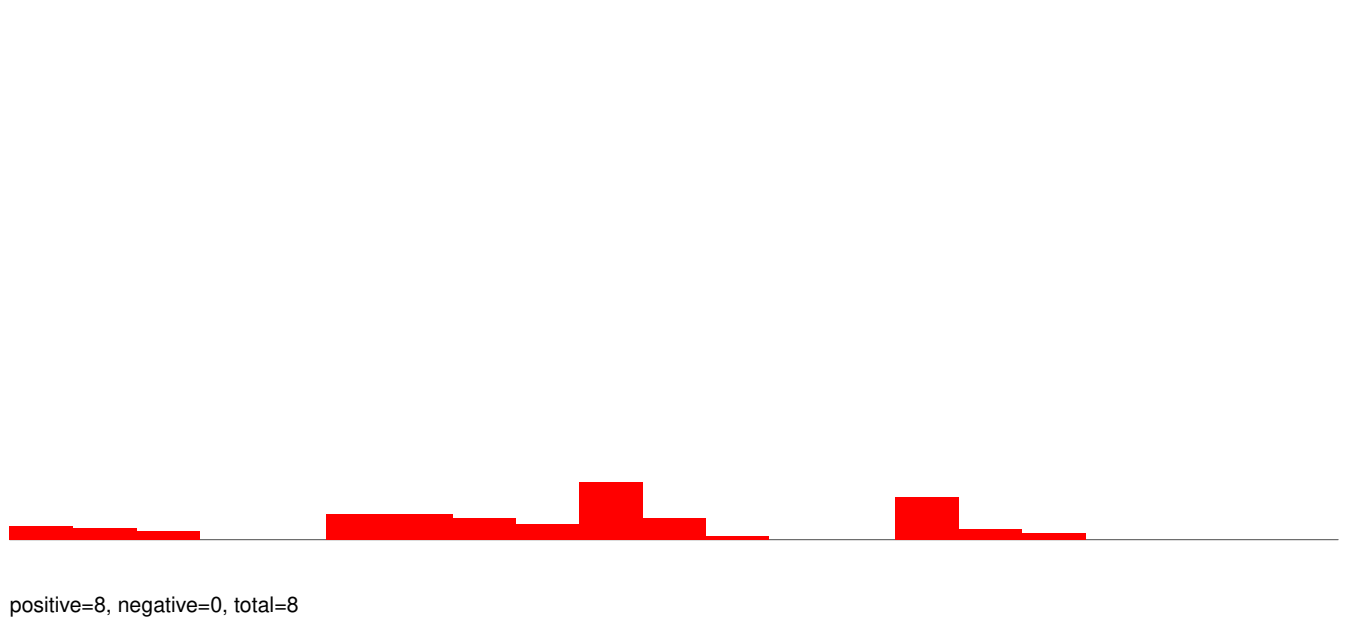
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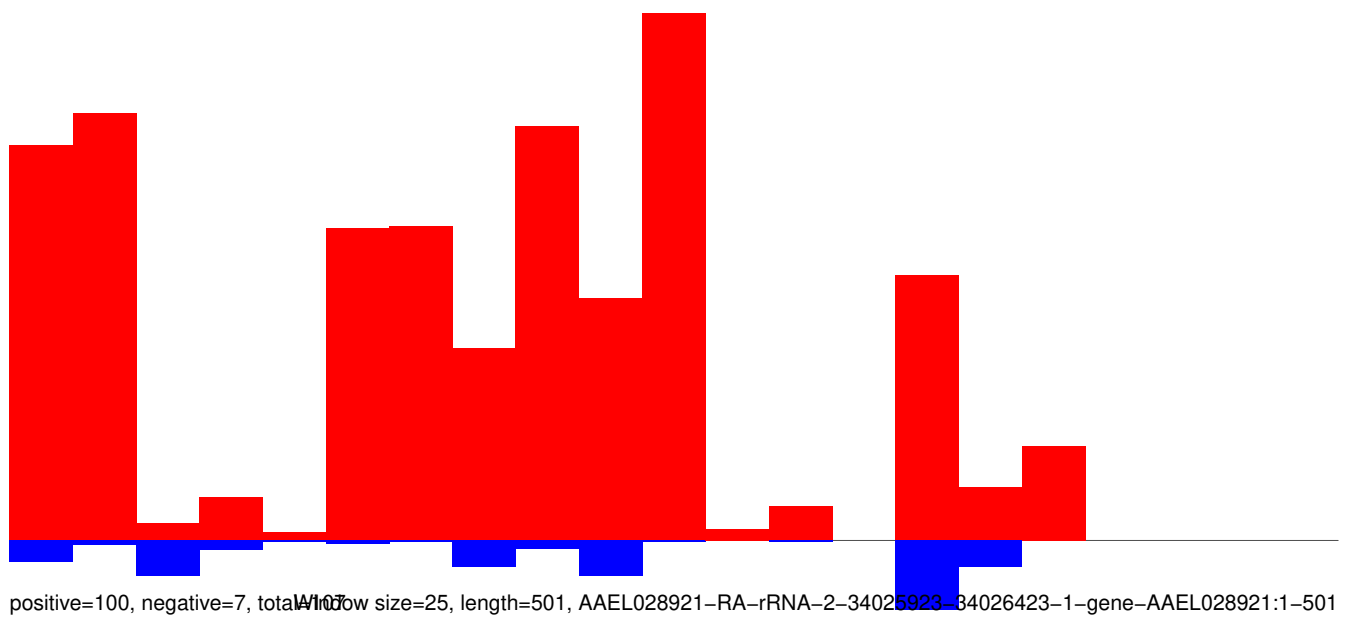
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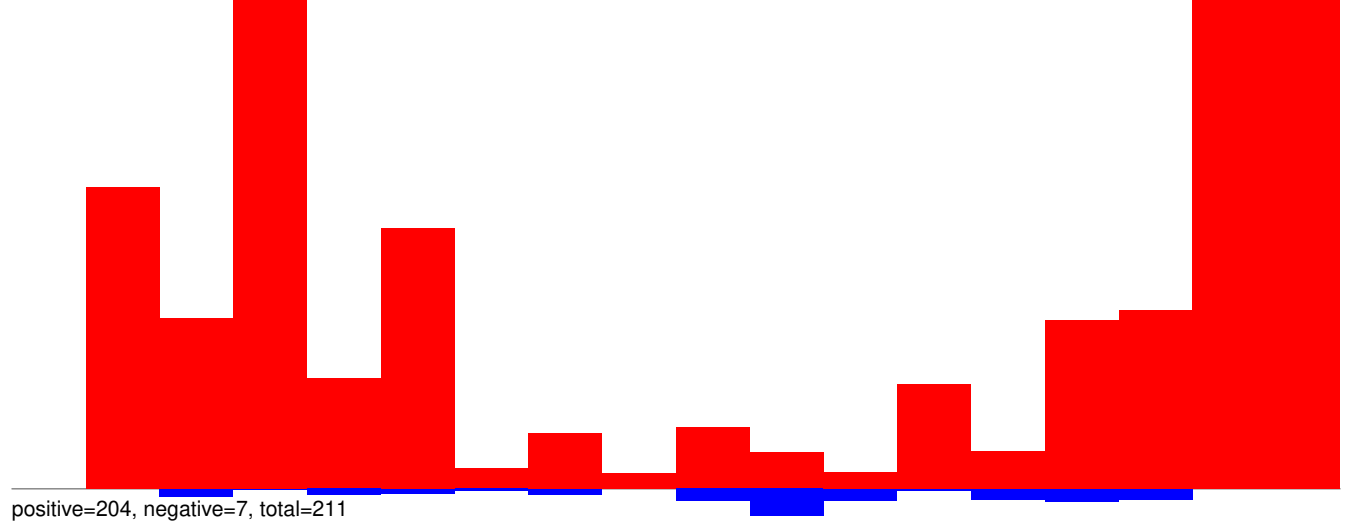
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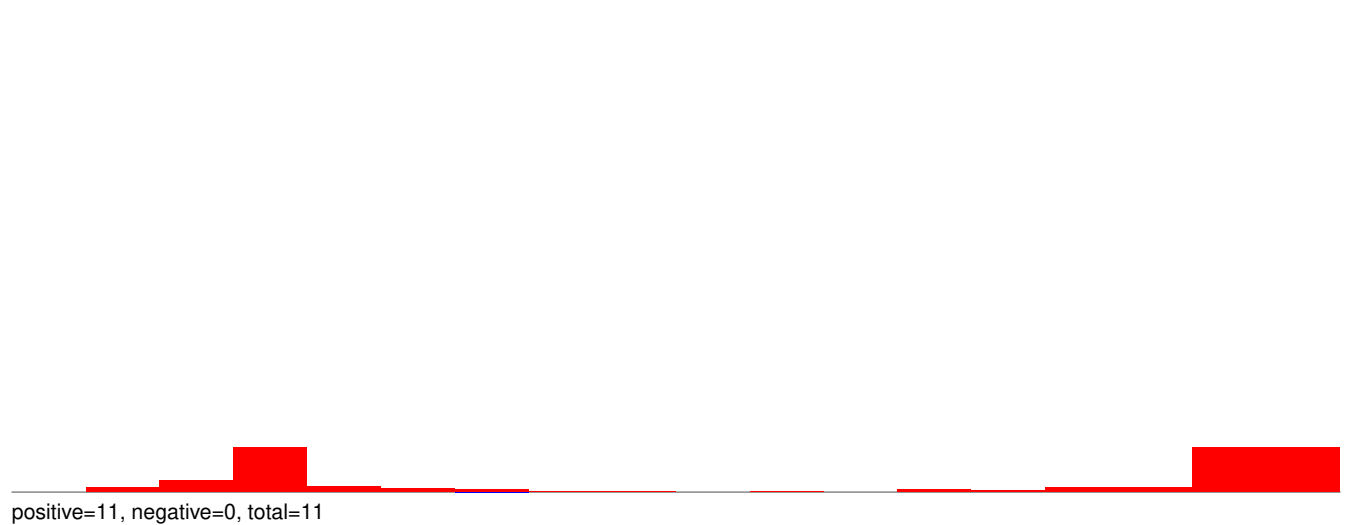
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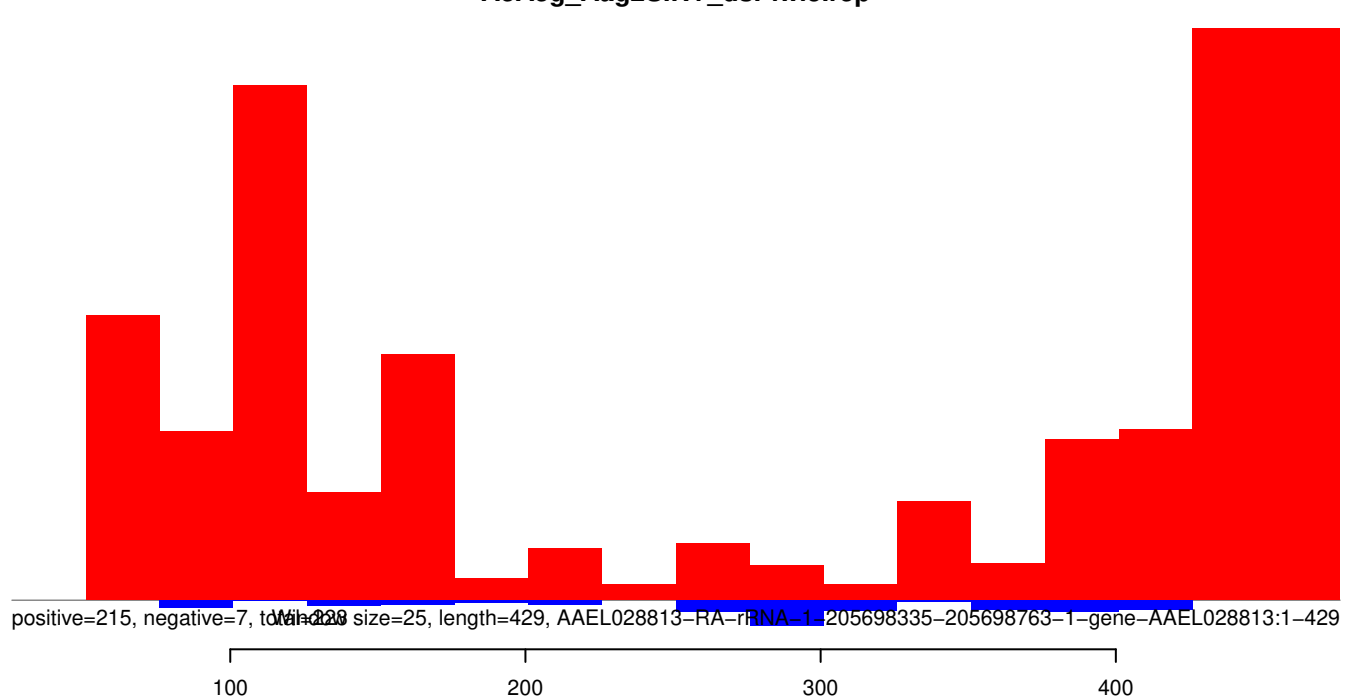
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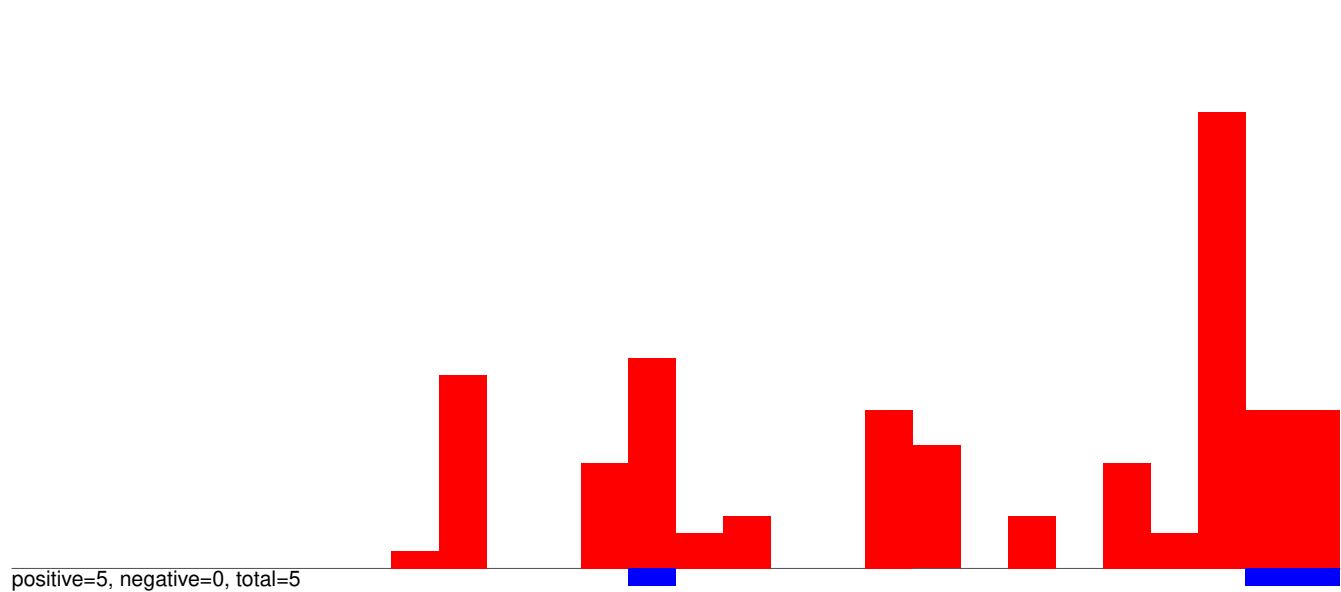
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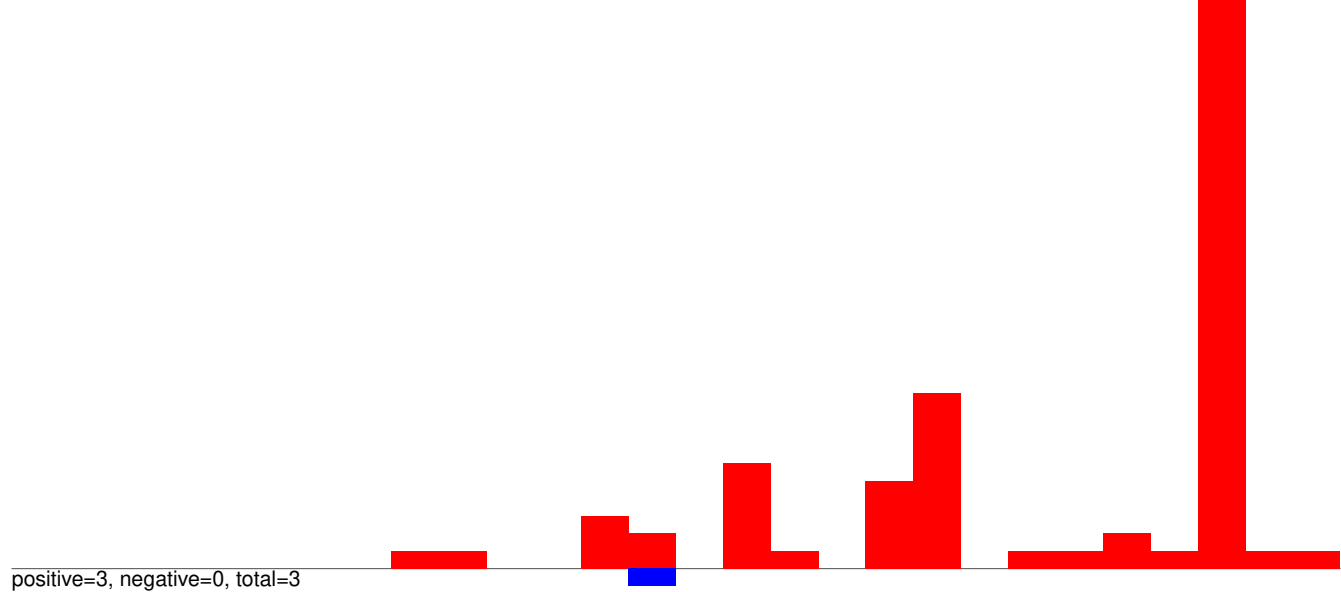
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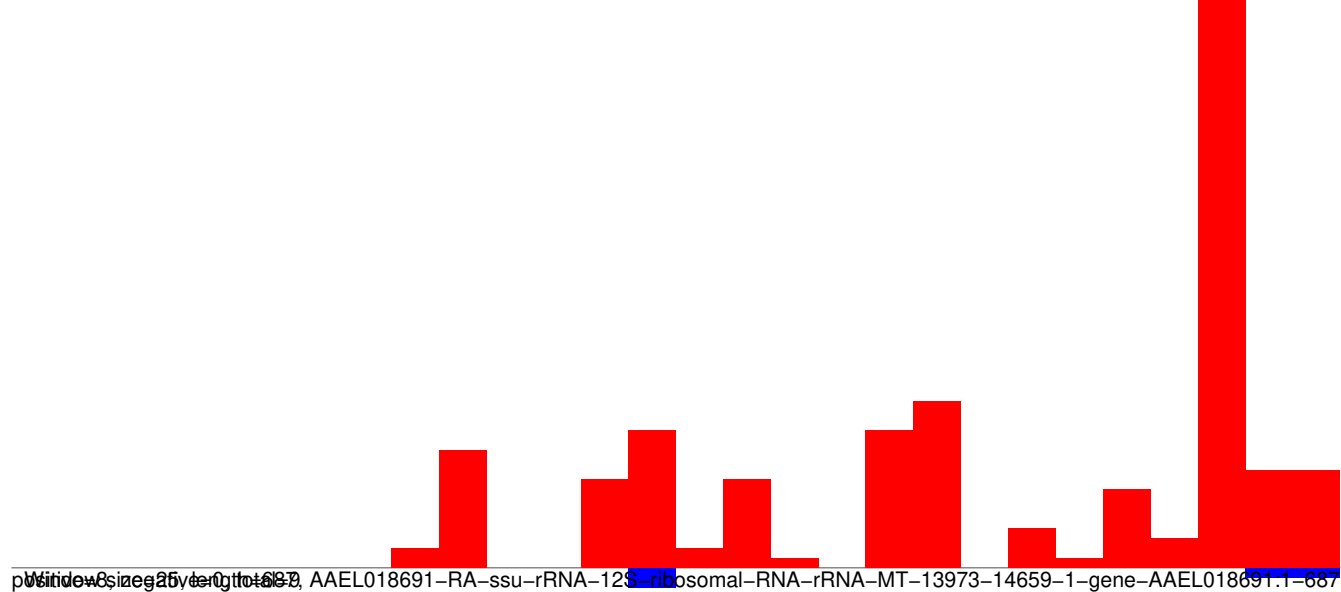
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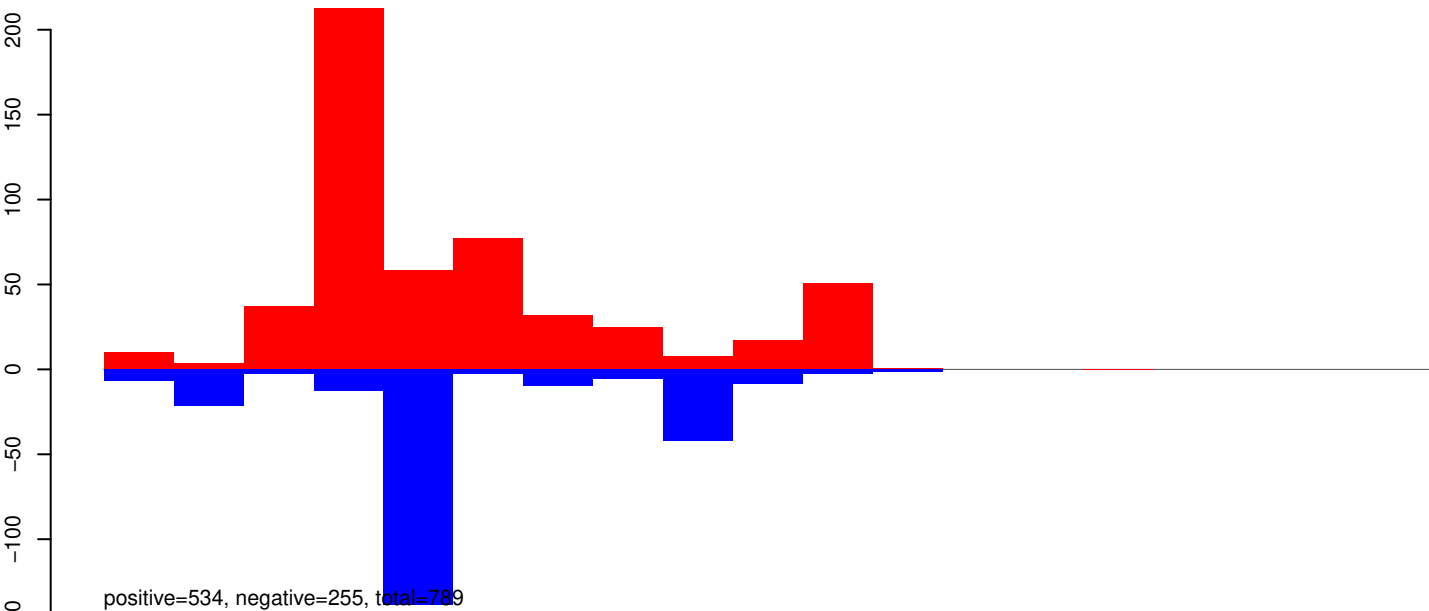
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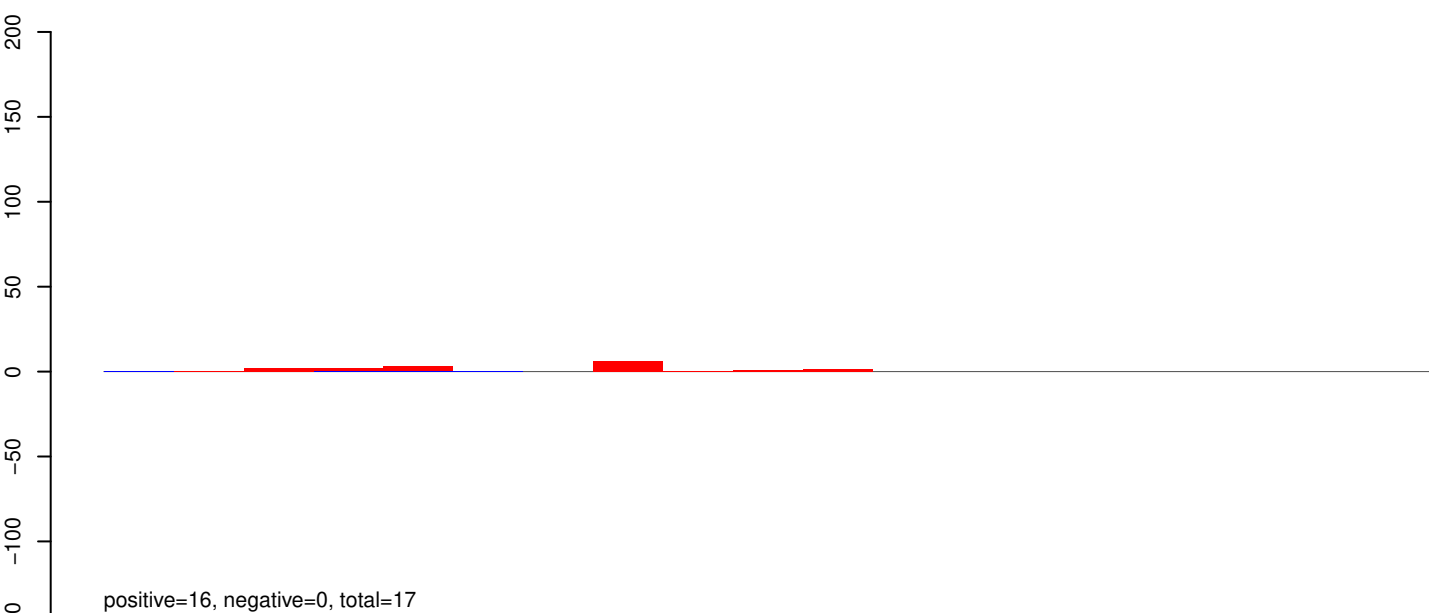
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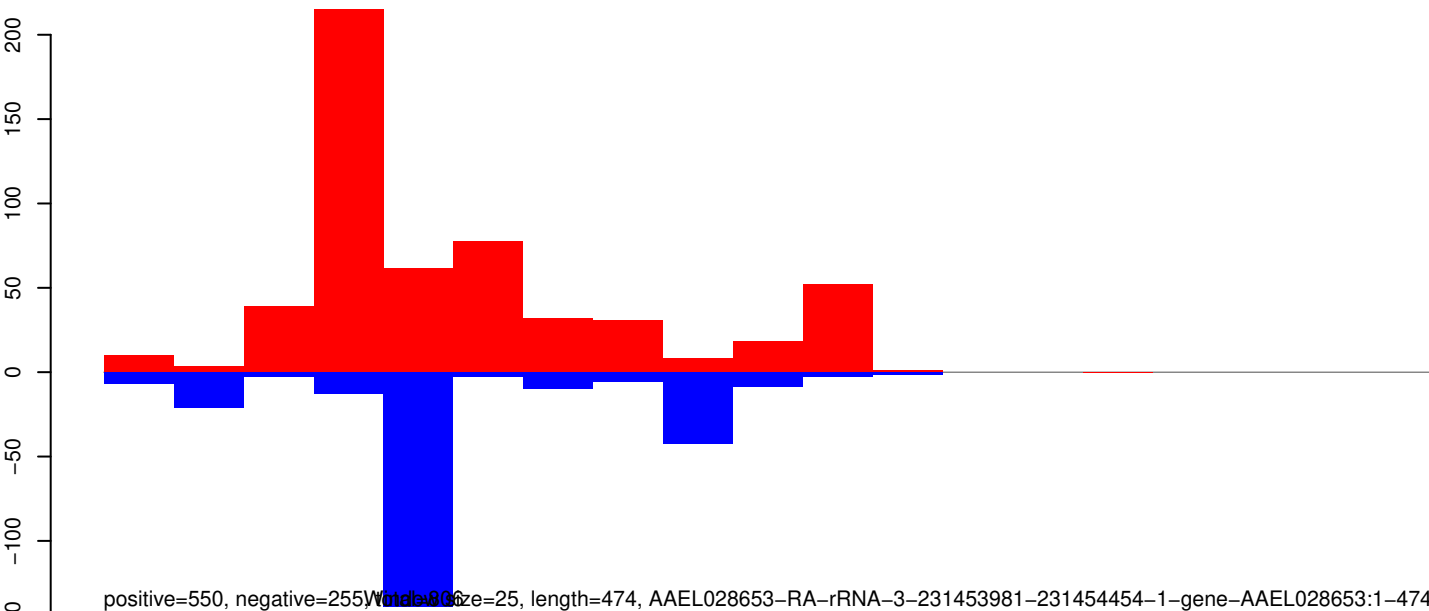
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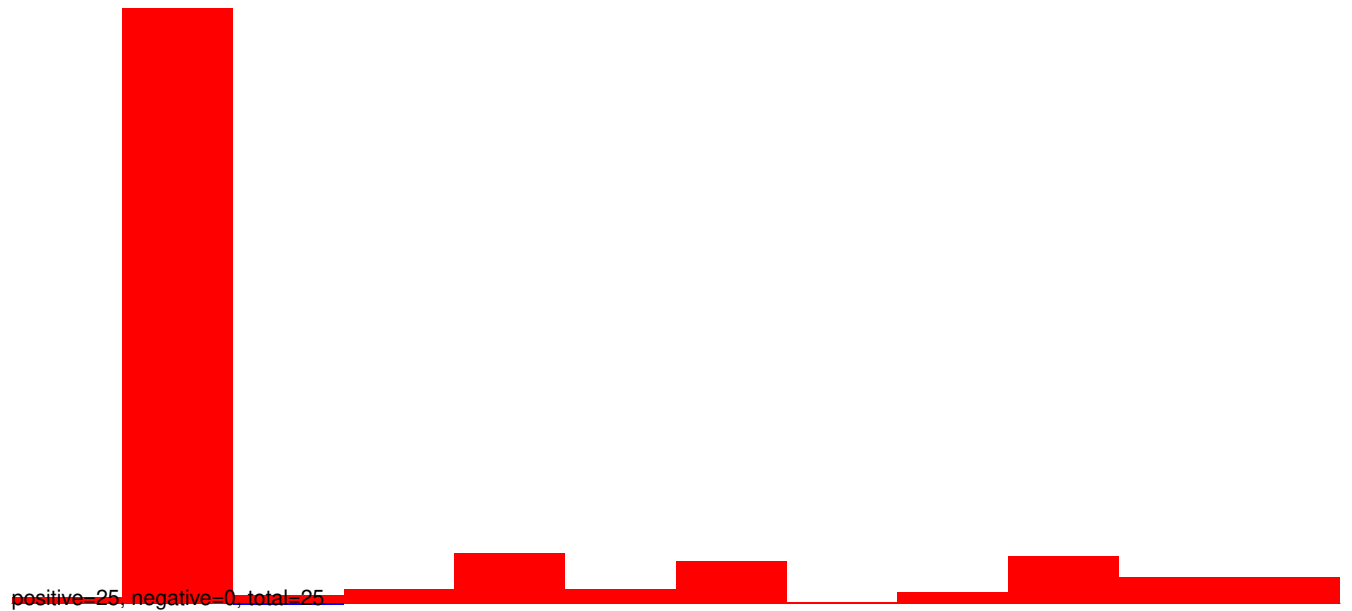
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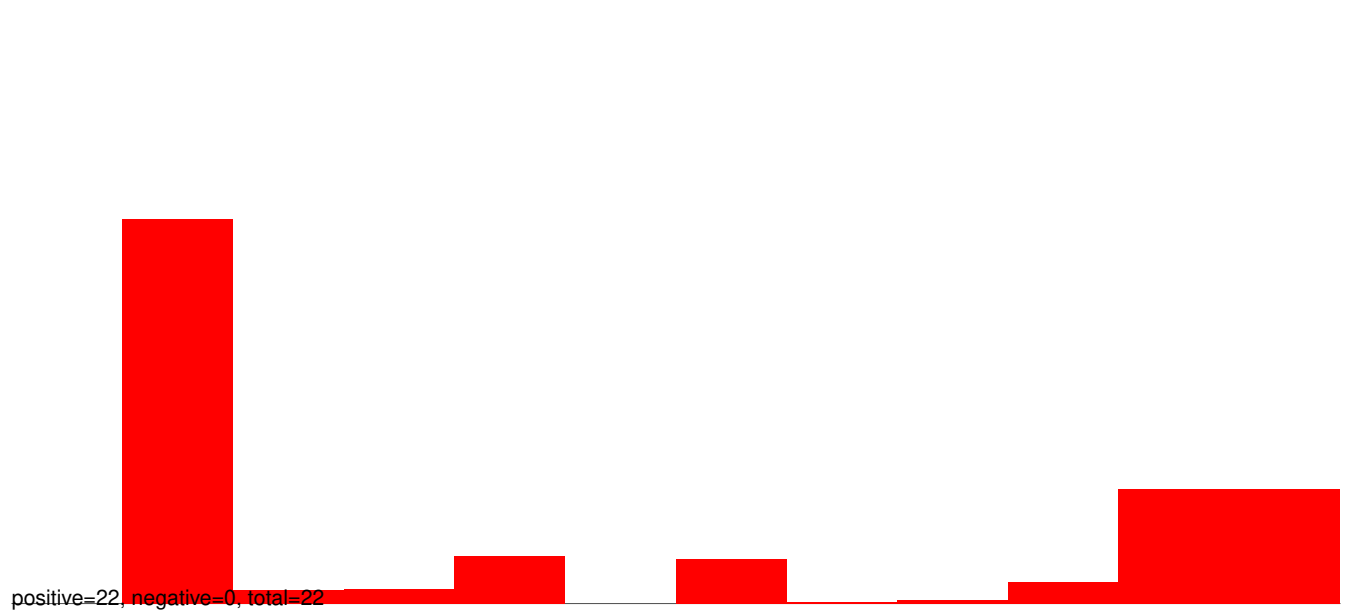
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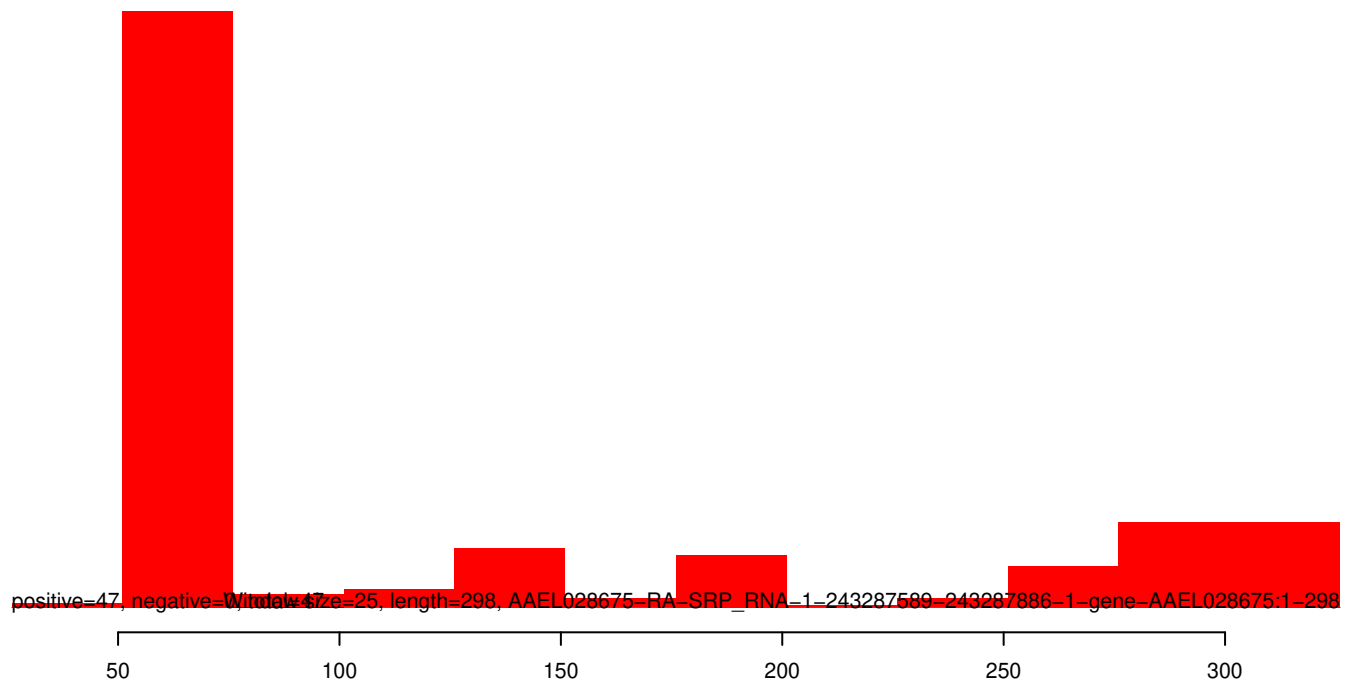
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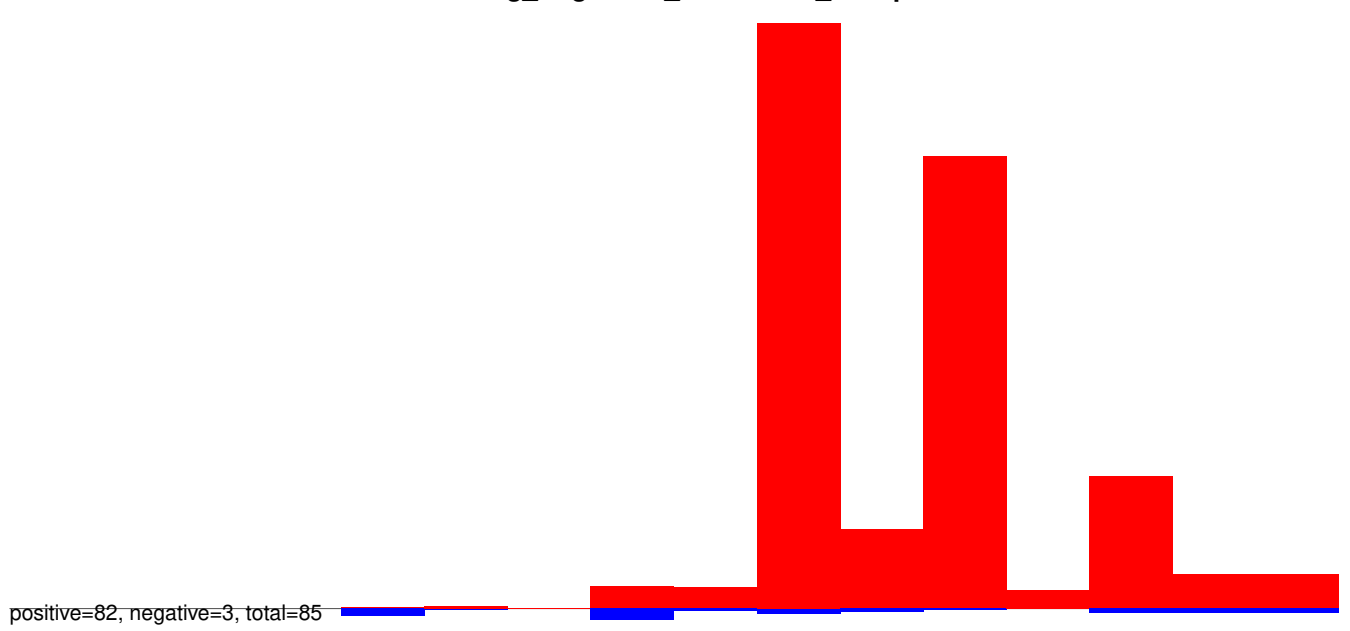
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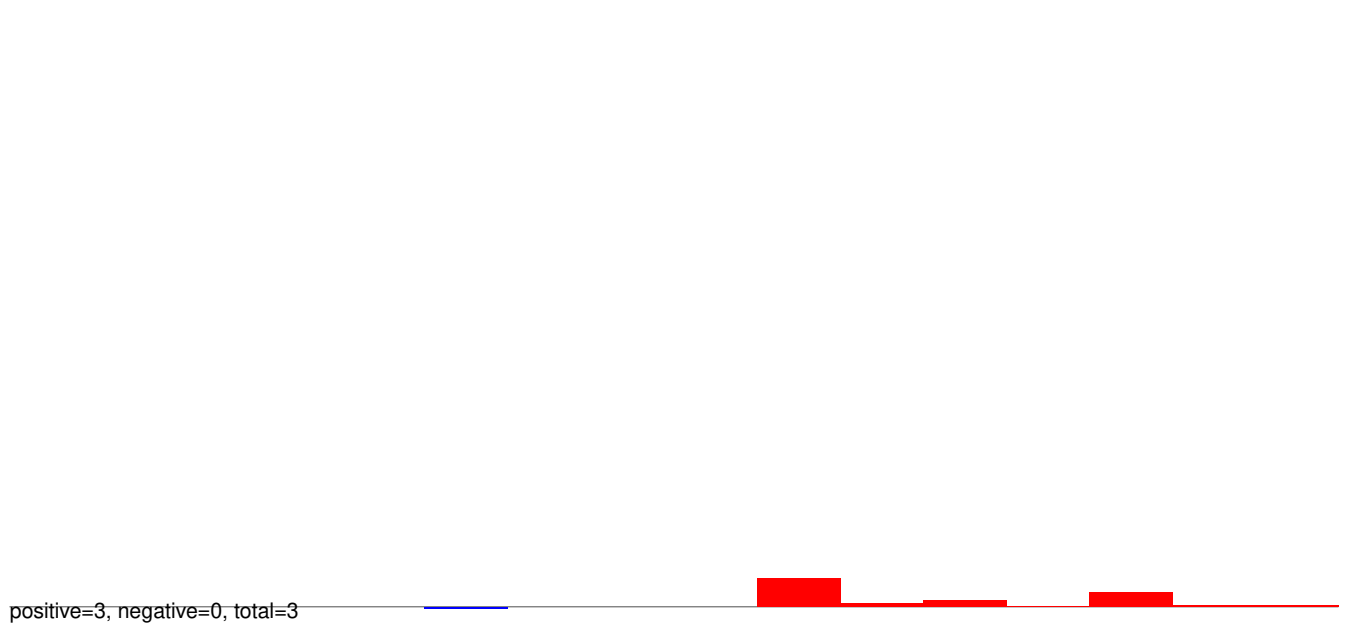
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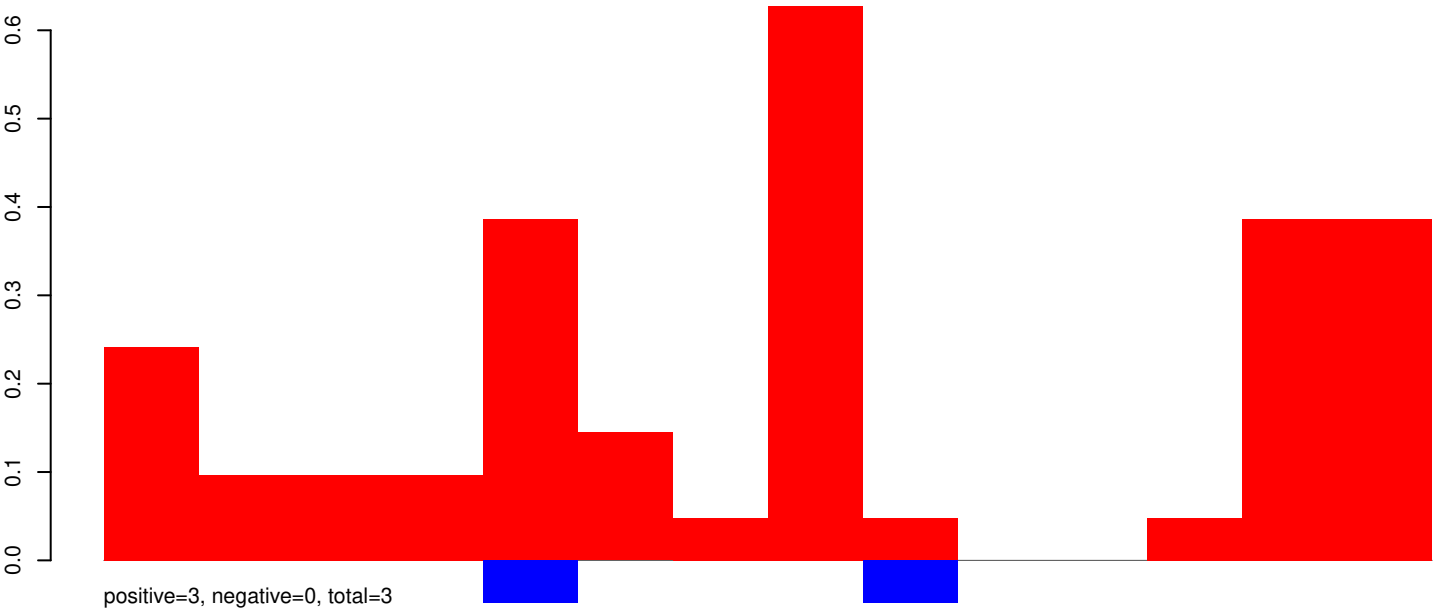
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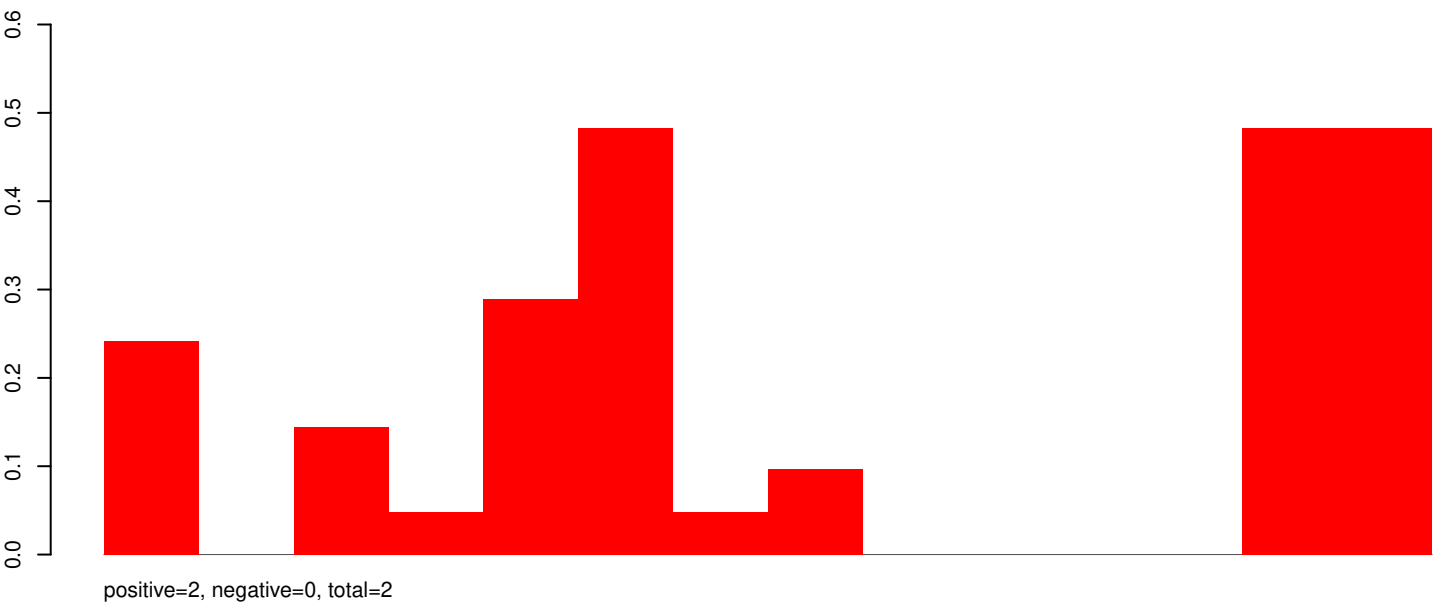
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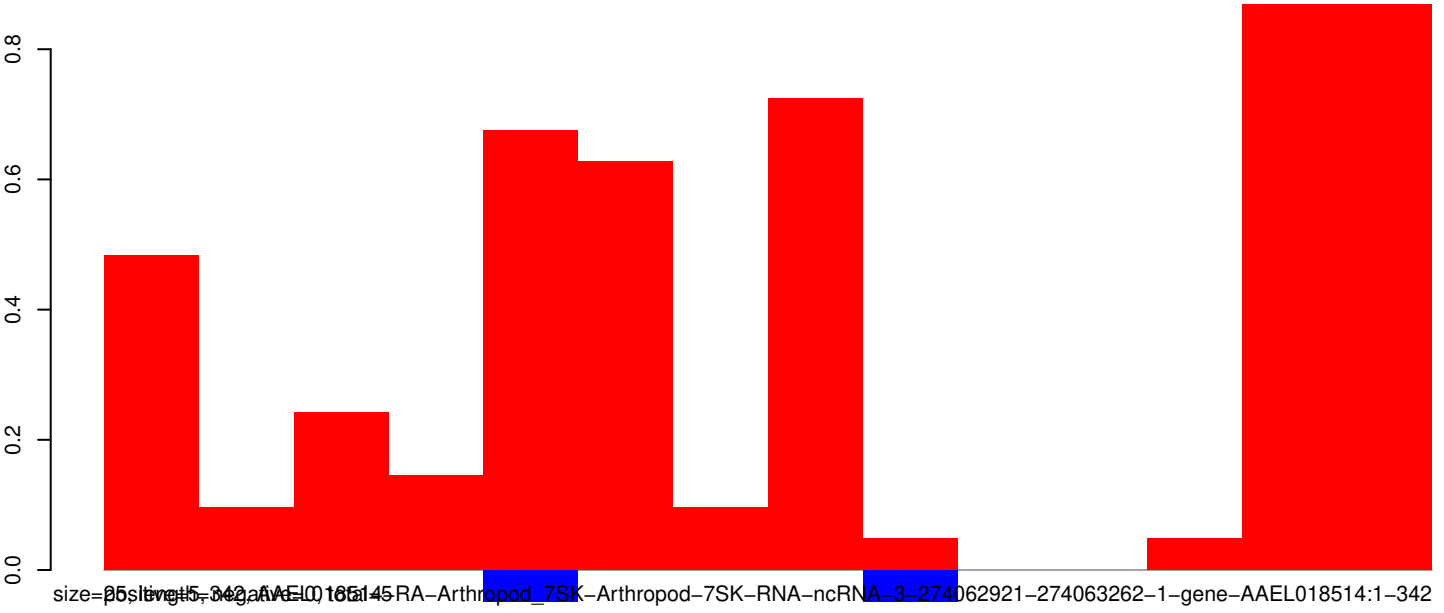
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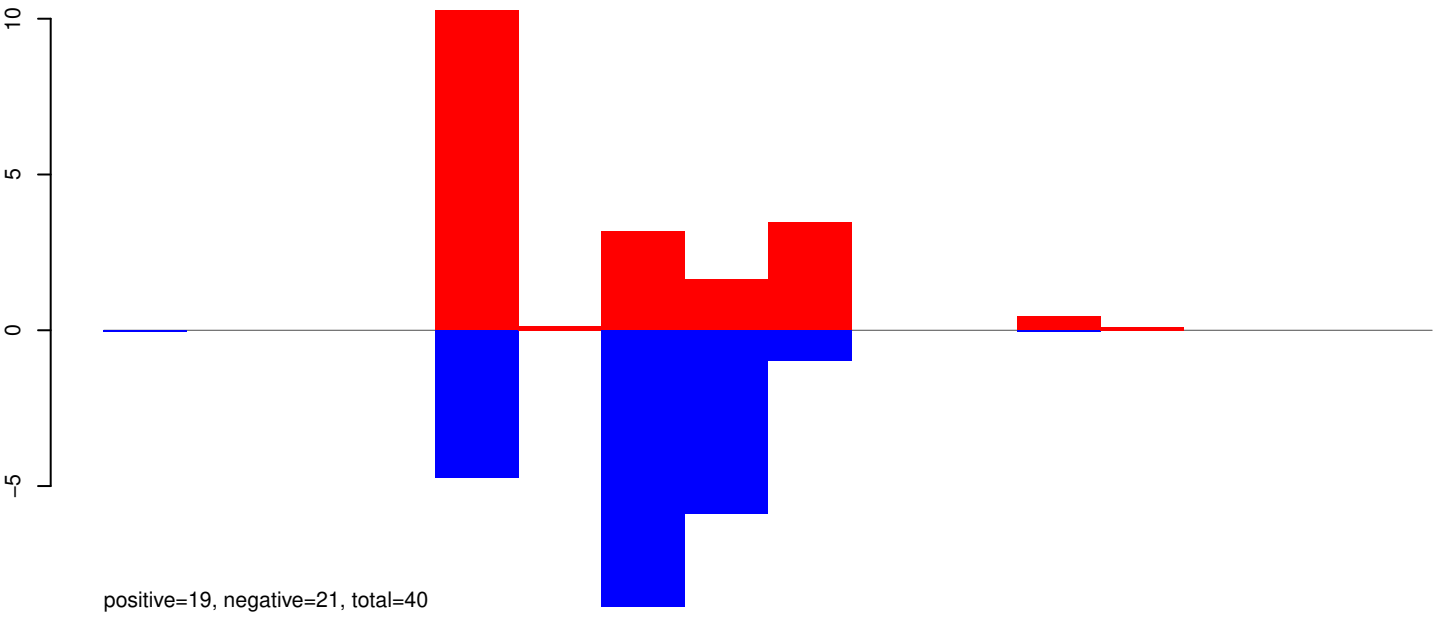


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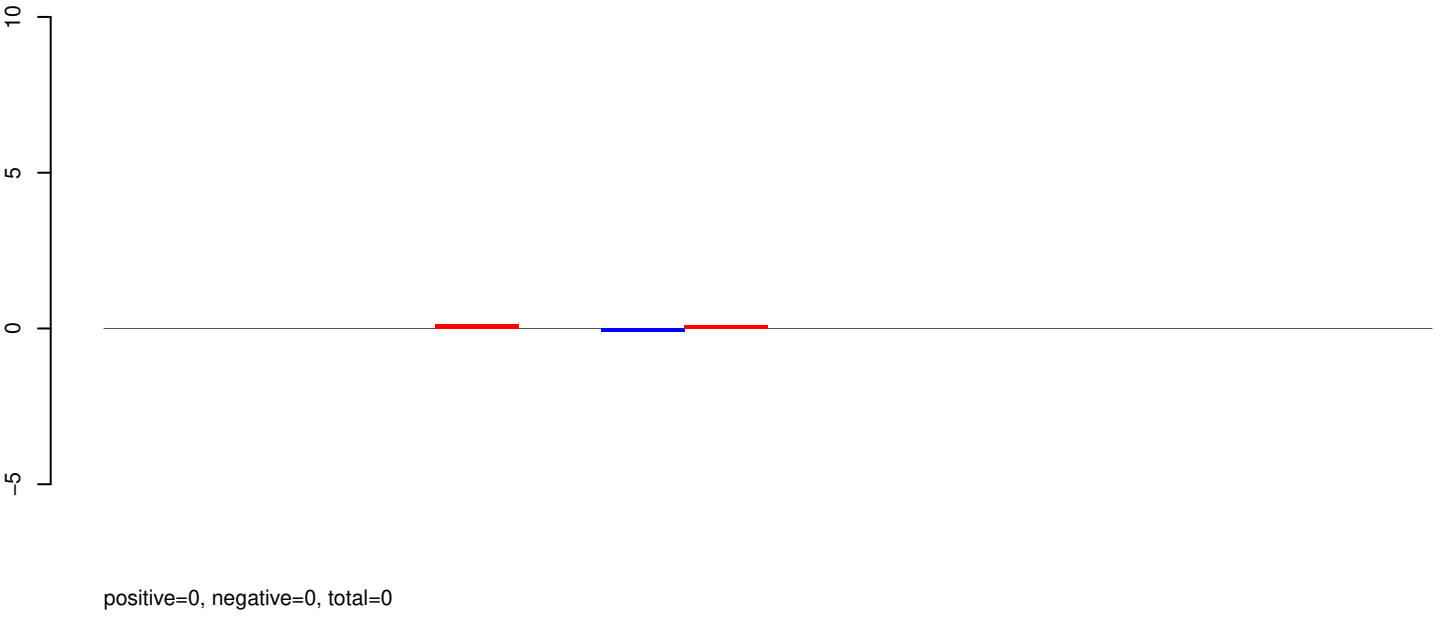


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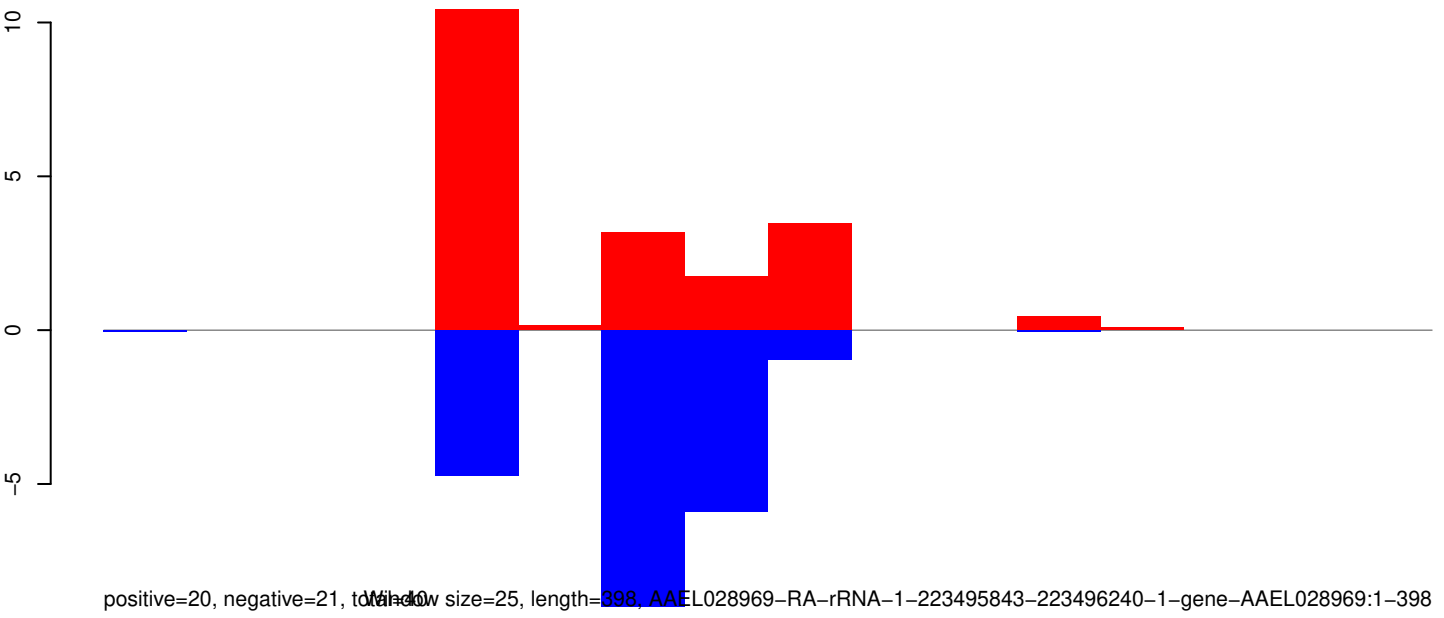
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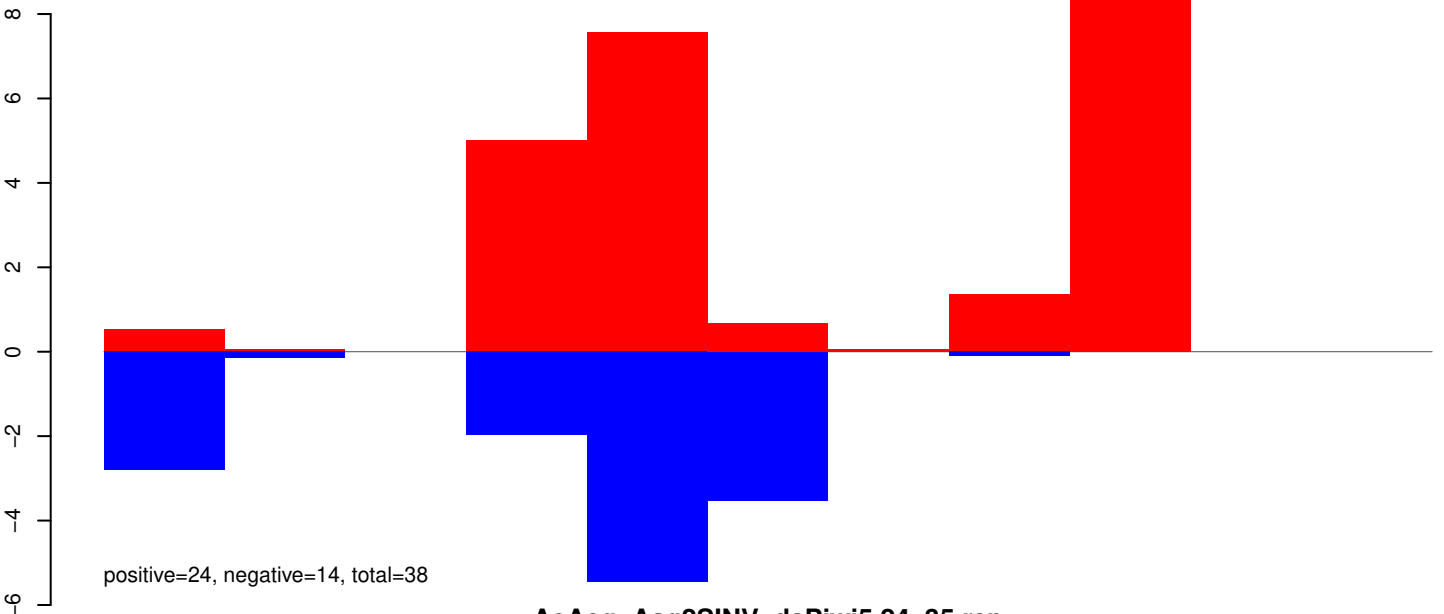
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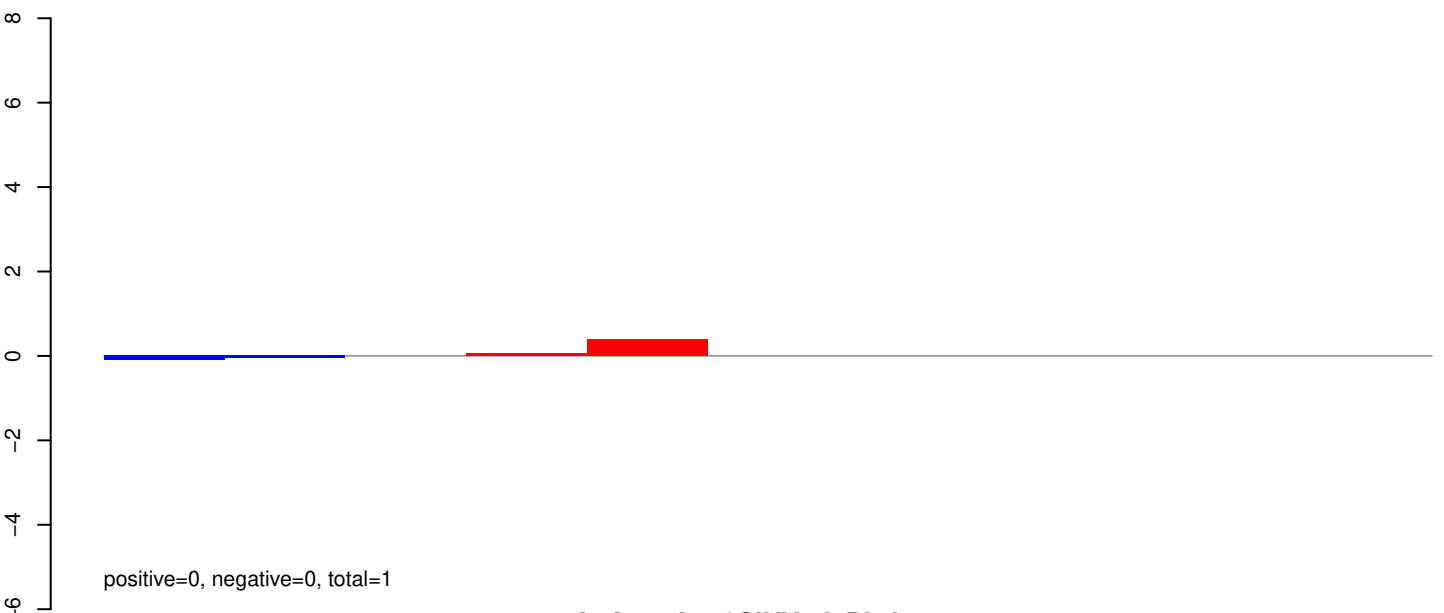
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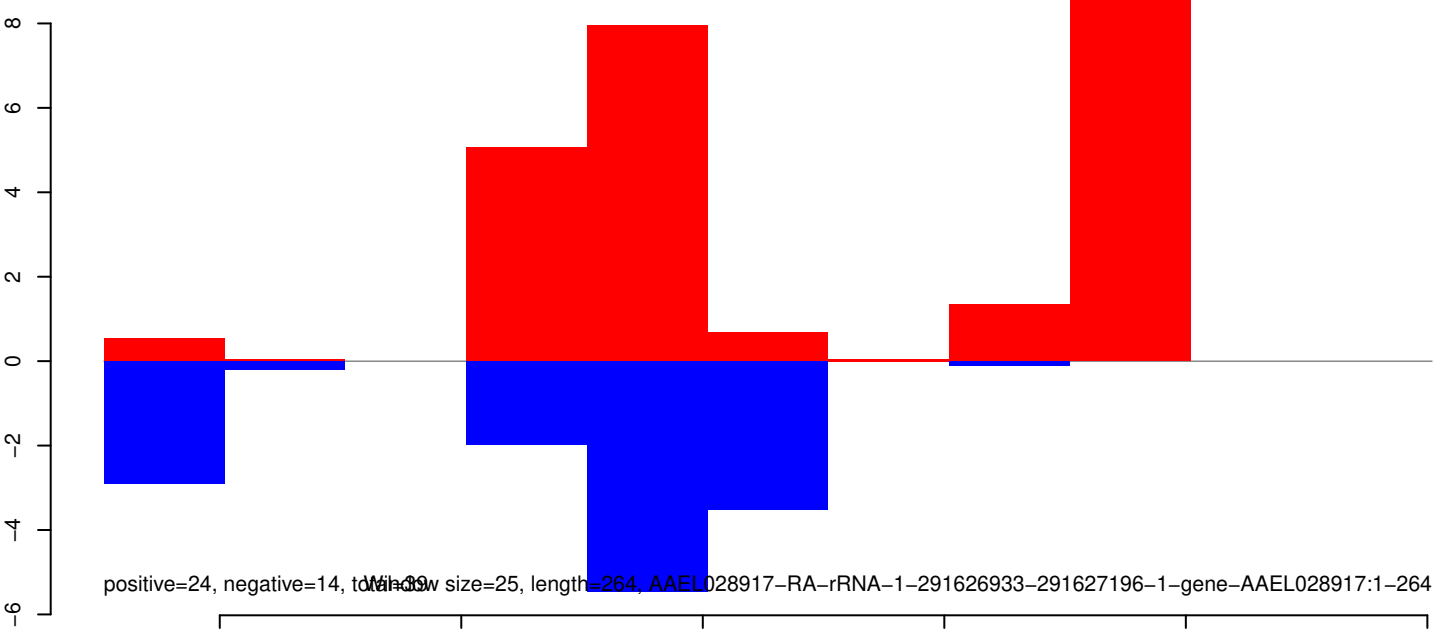
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AeAeg_Aag2SINV_dsPiwi5.24_35.rep

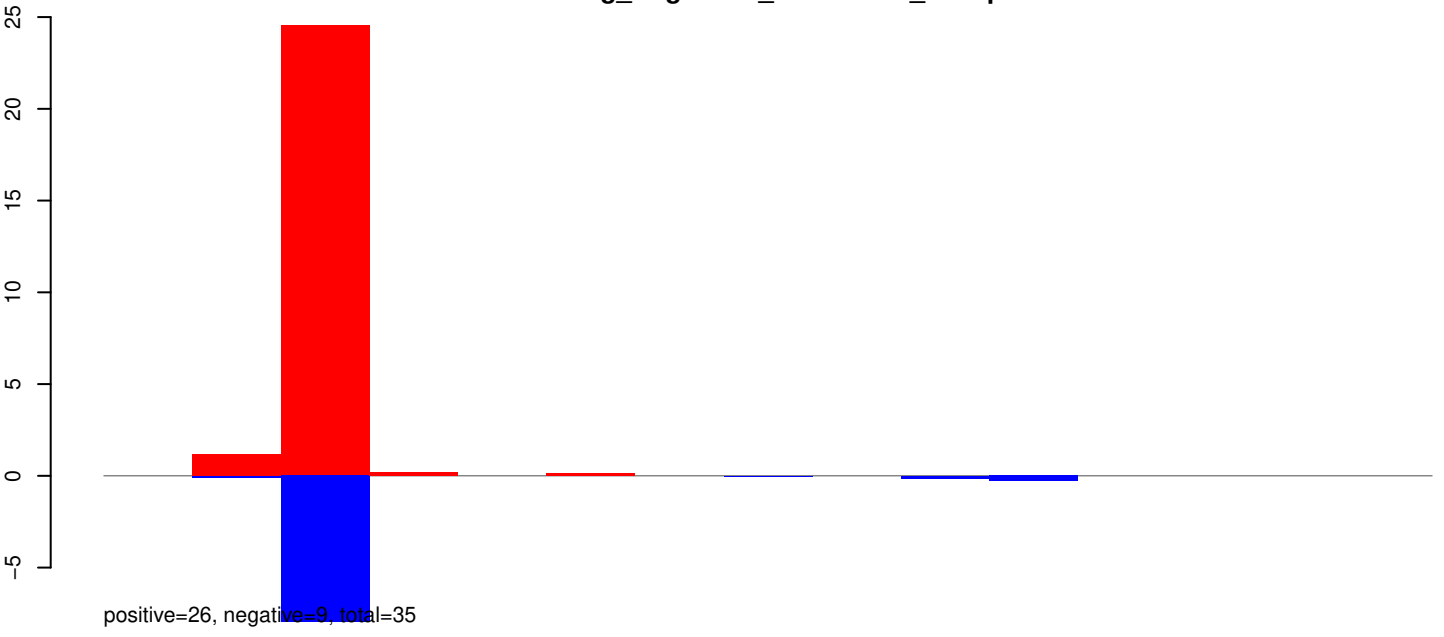


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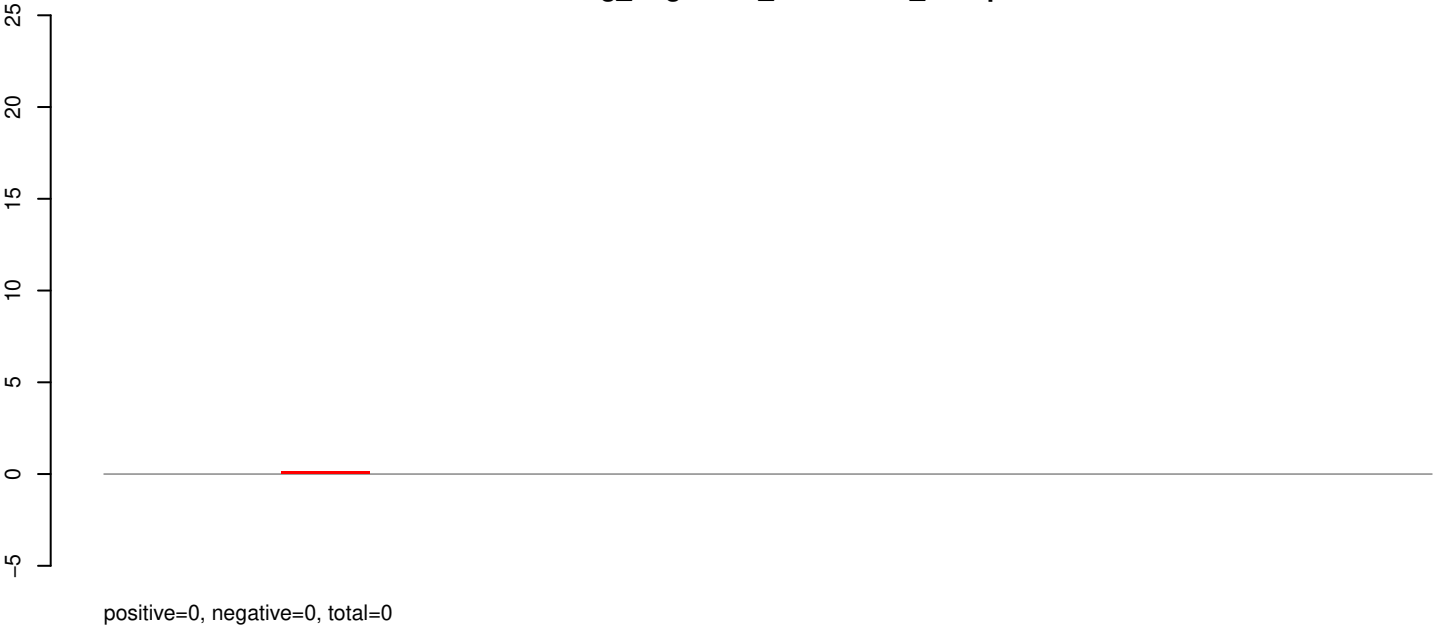


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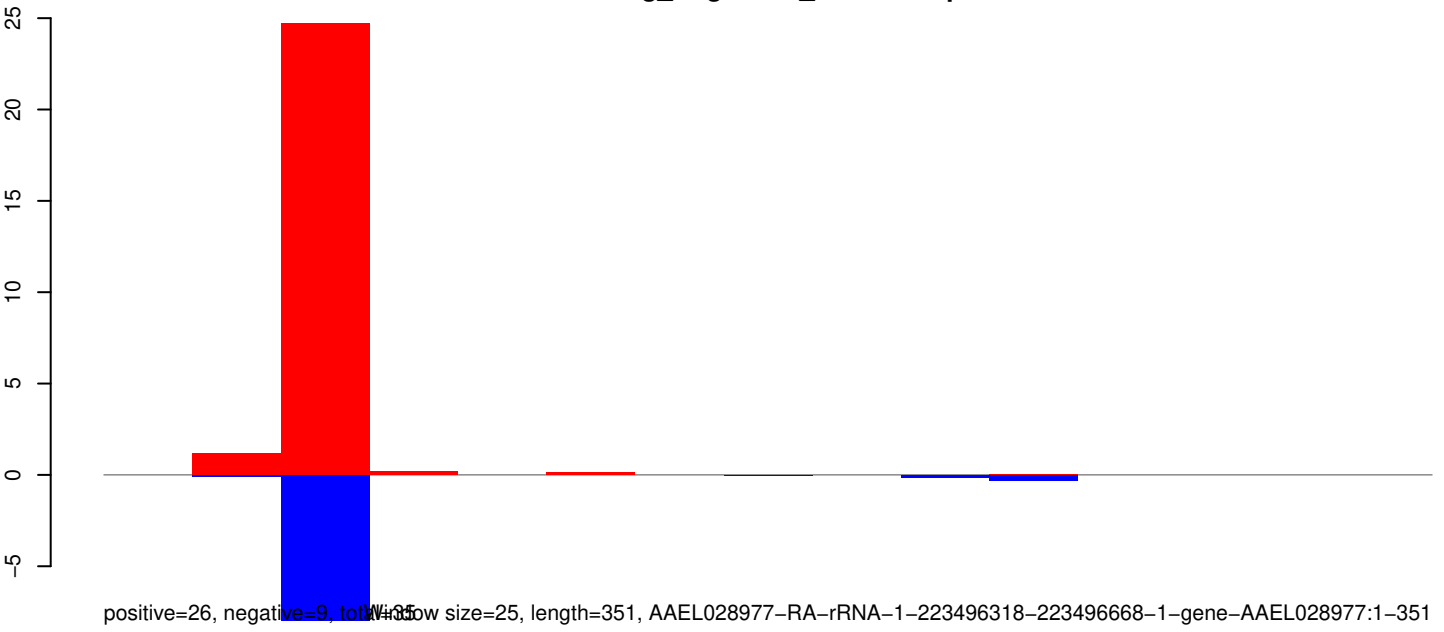
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AeAeg_Aag2SINV_dsPiwi5.24_35.rep



AeAeg_Aag2SINV_dsPiwi5.rep



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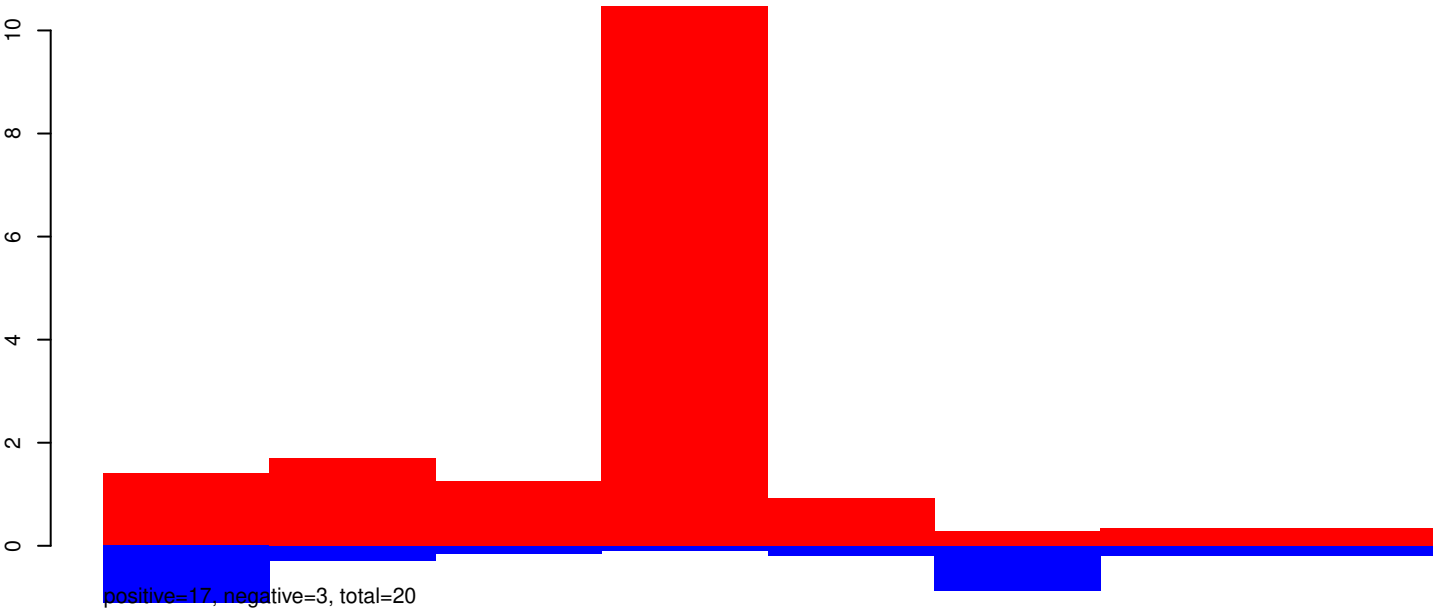
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200

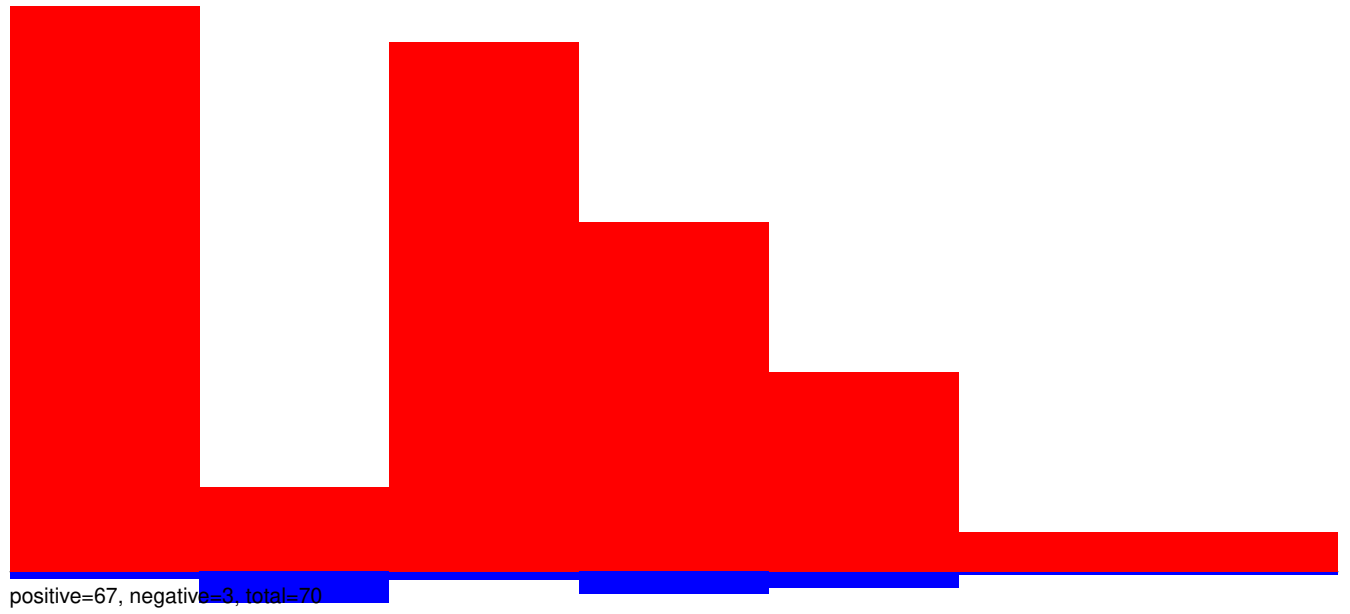
300

400

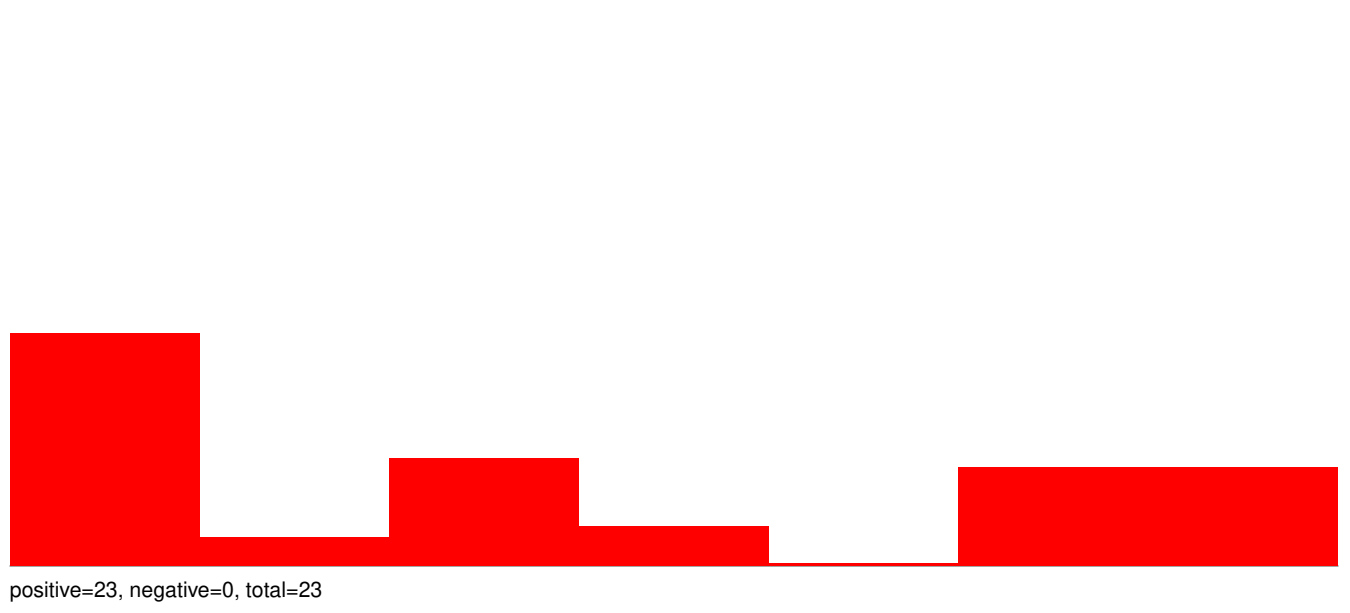
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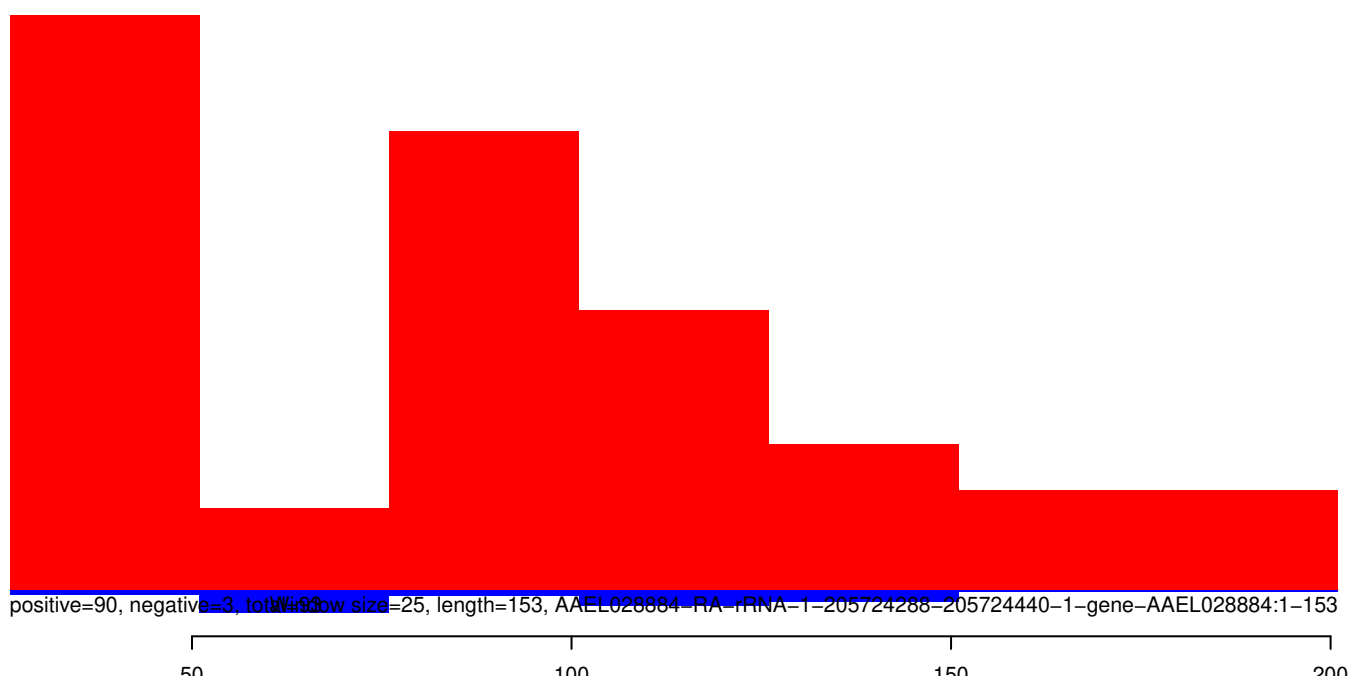
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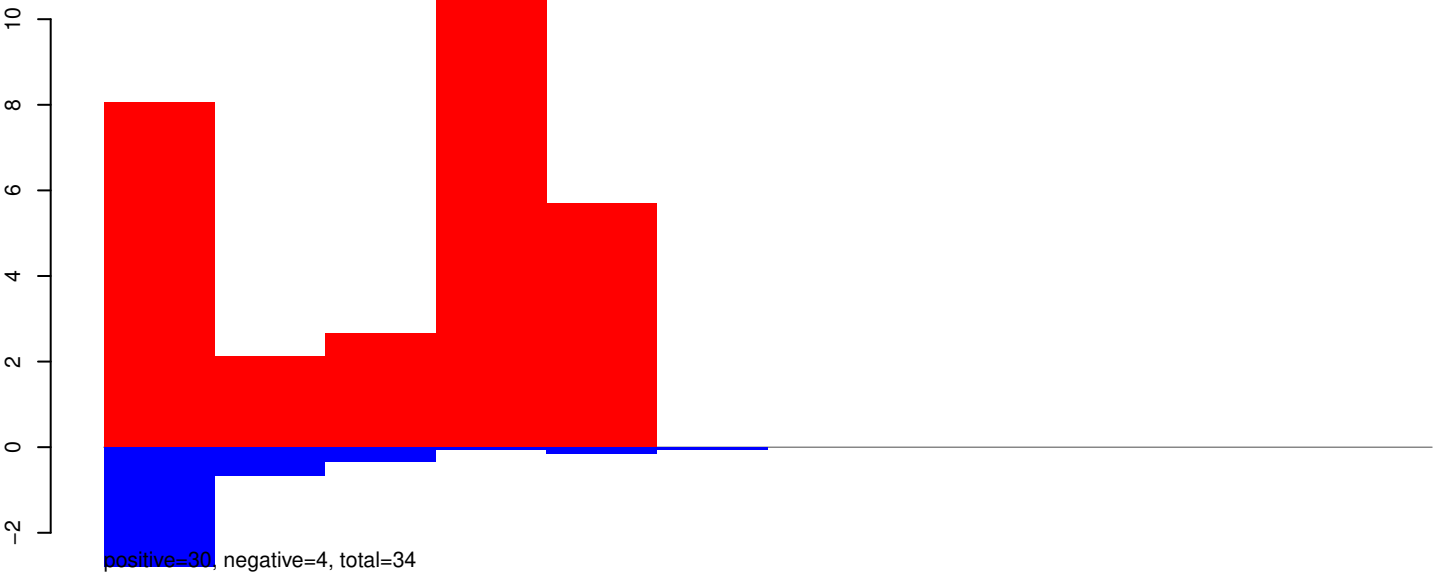
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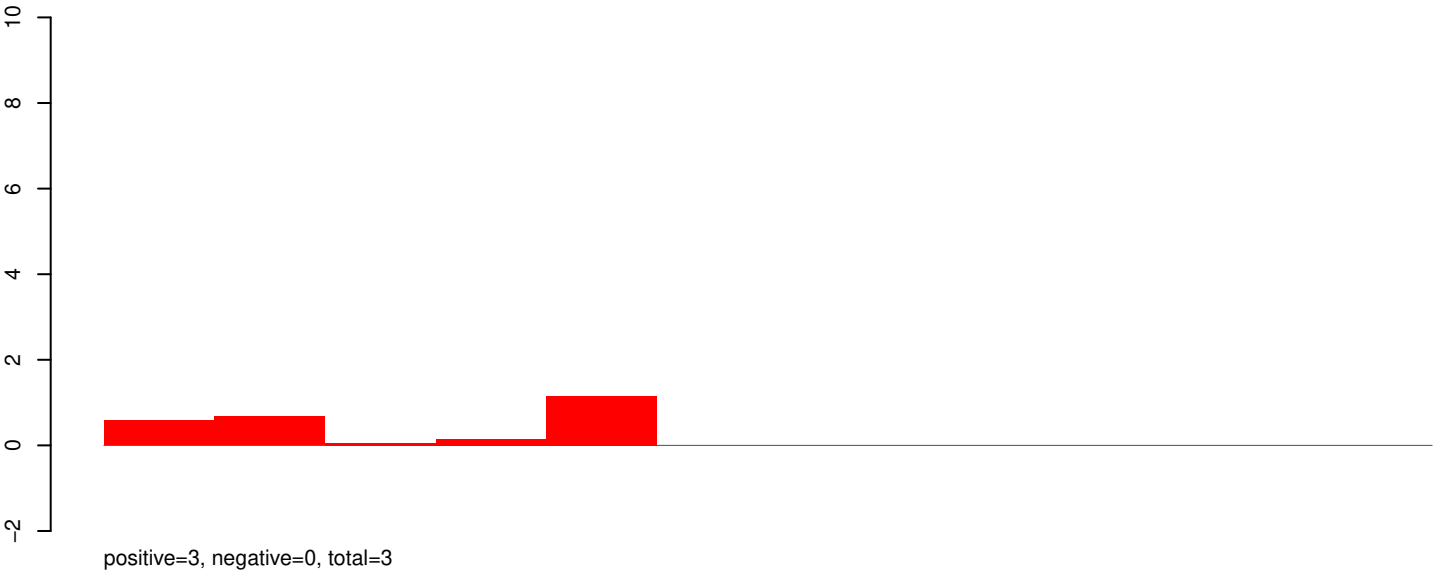
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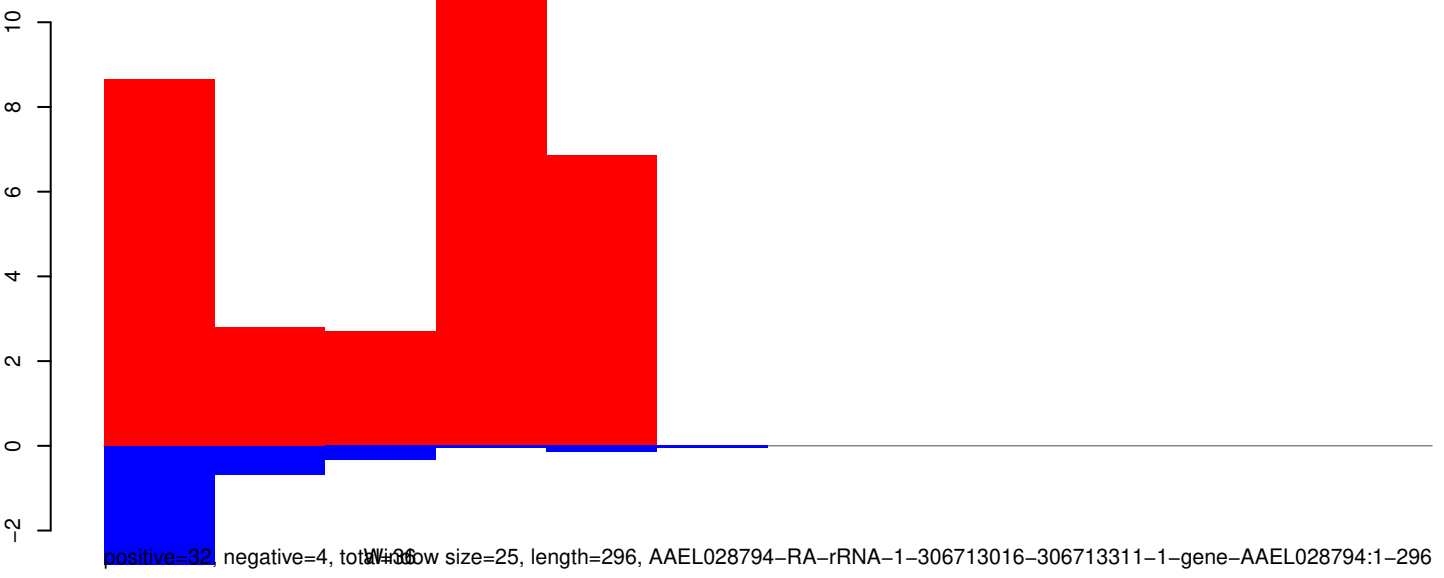
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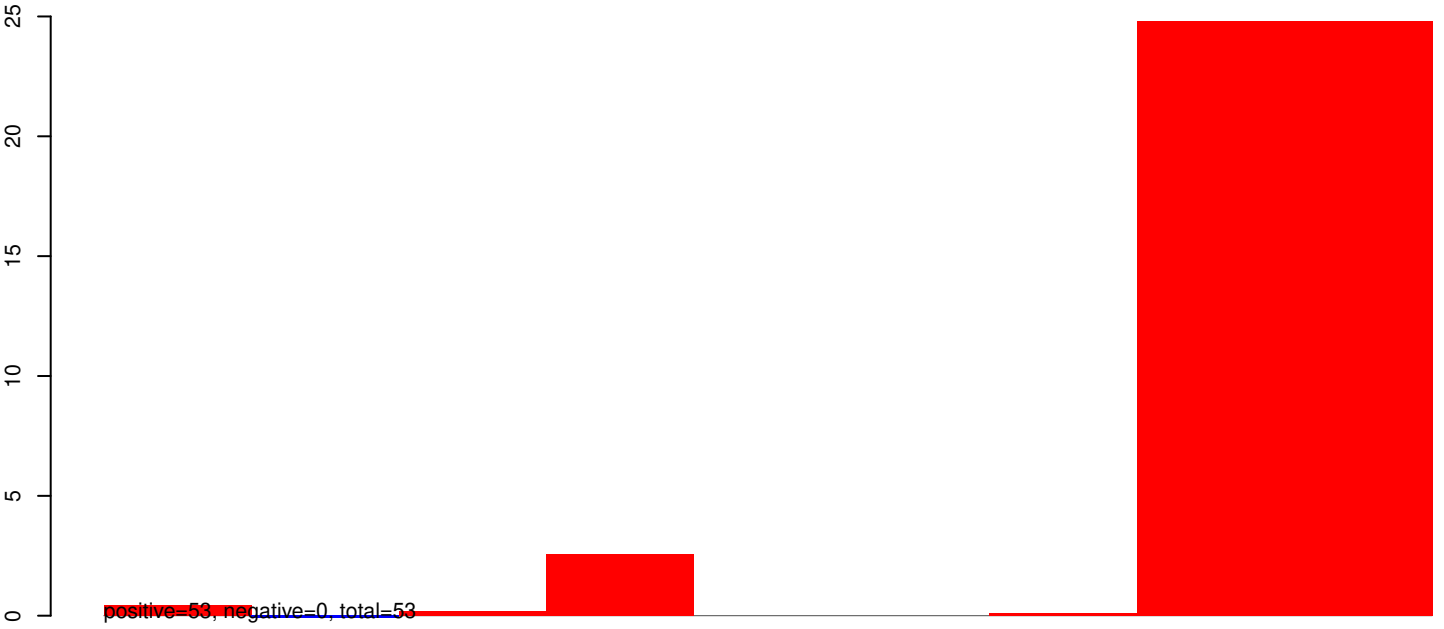


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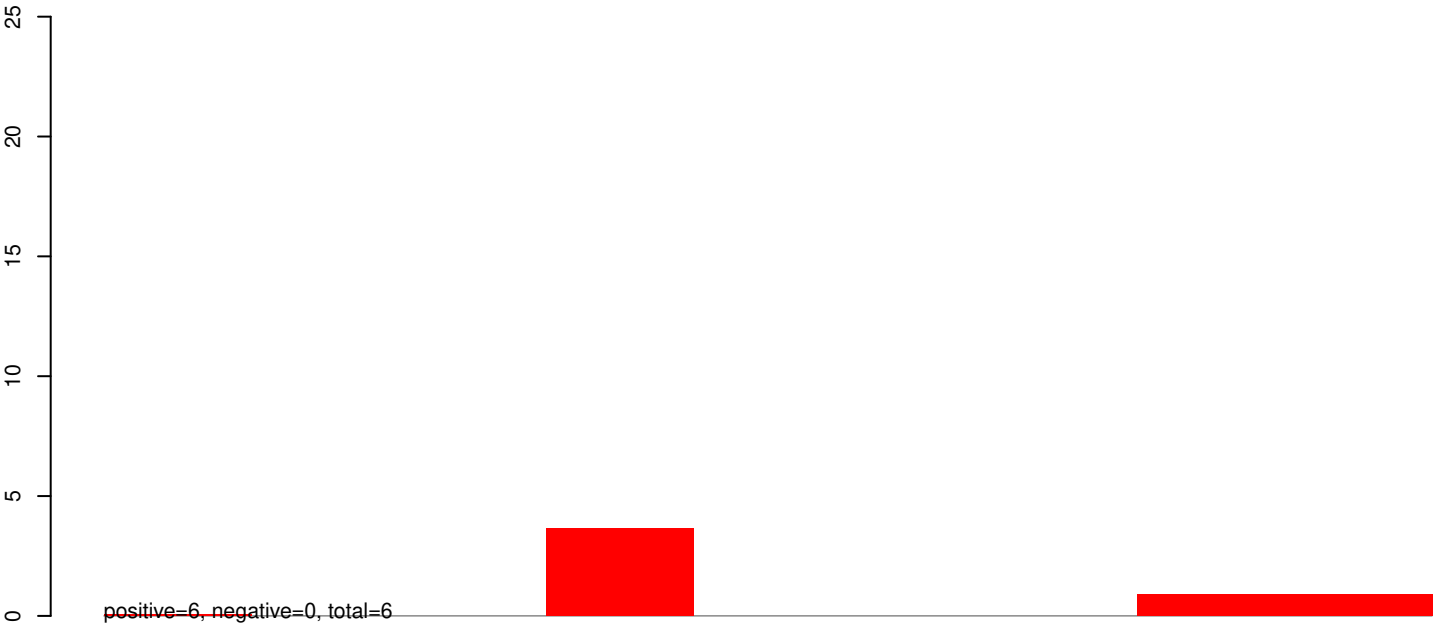


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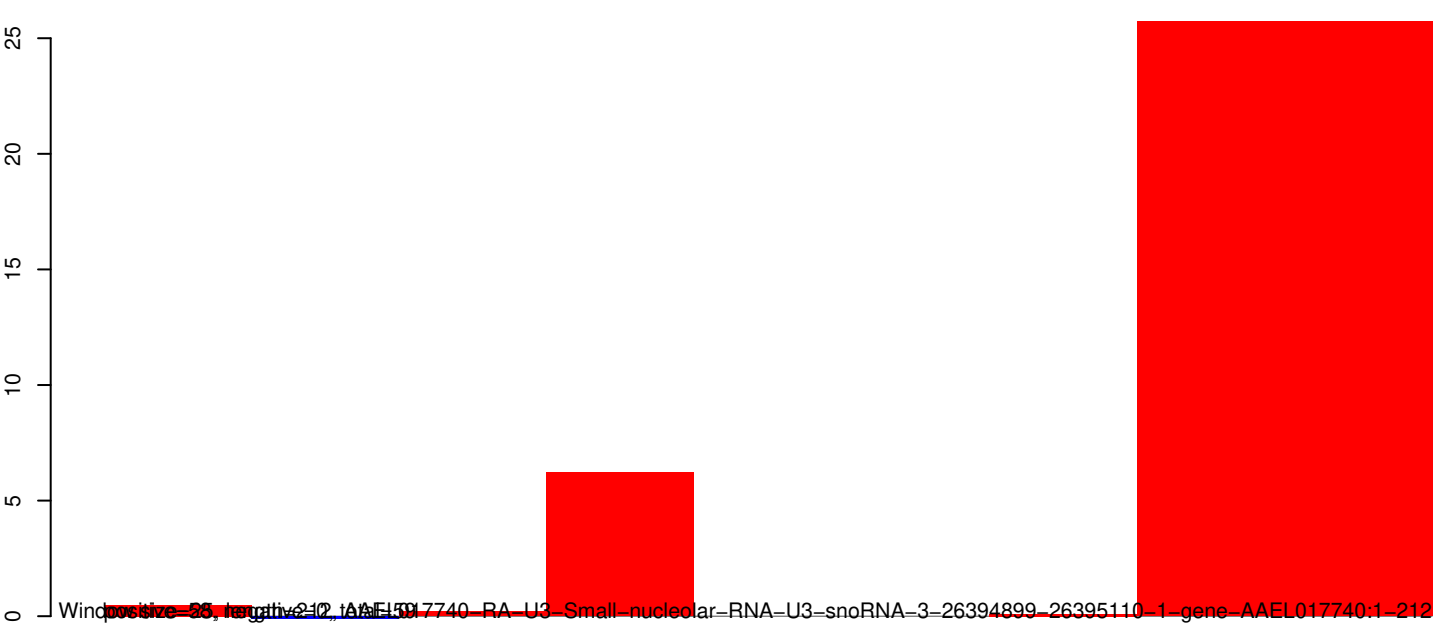
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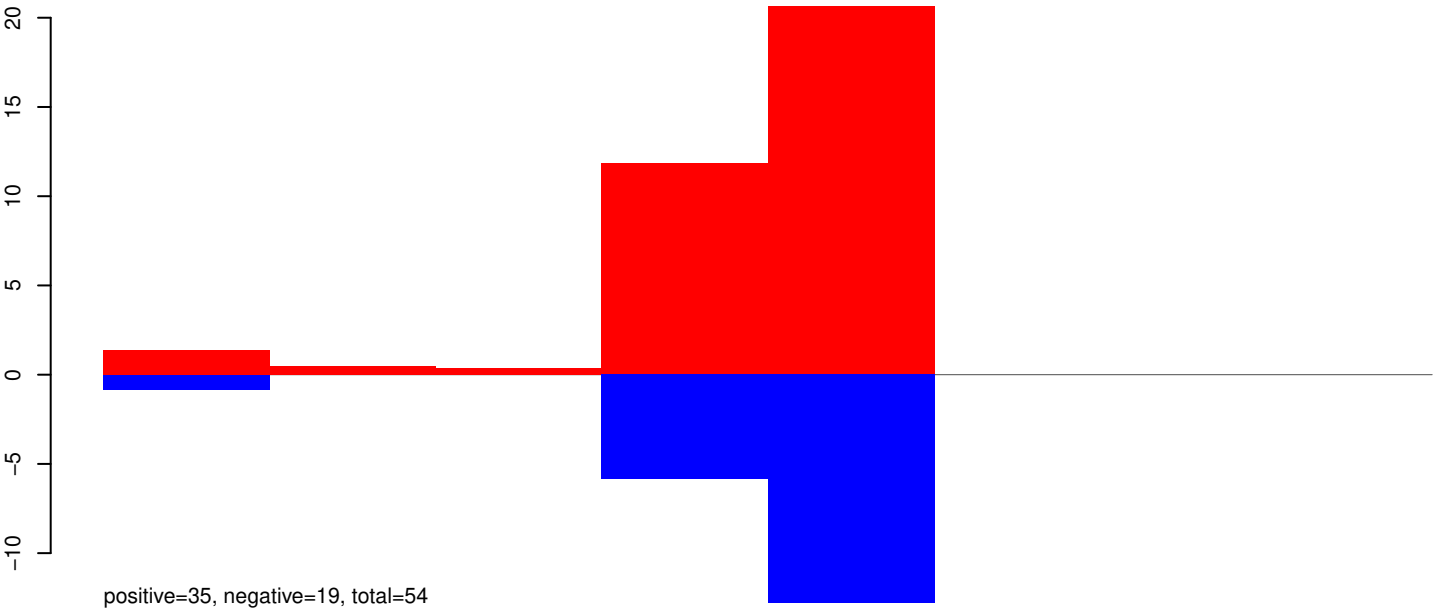
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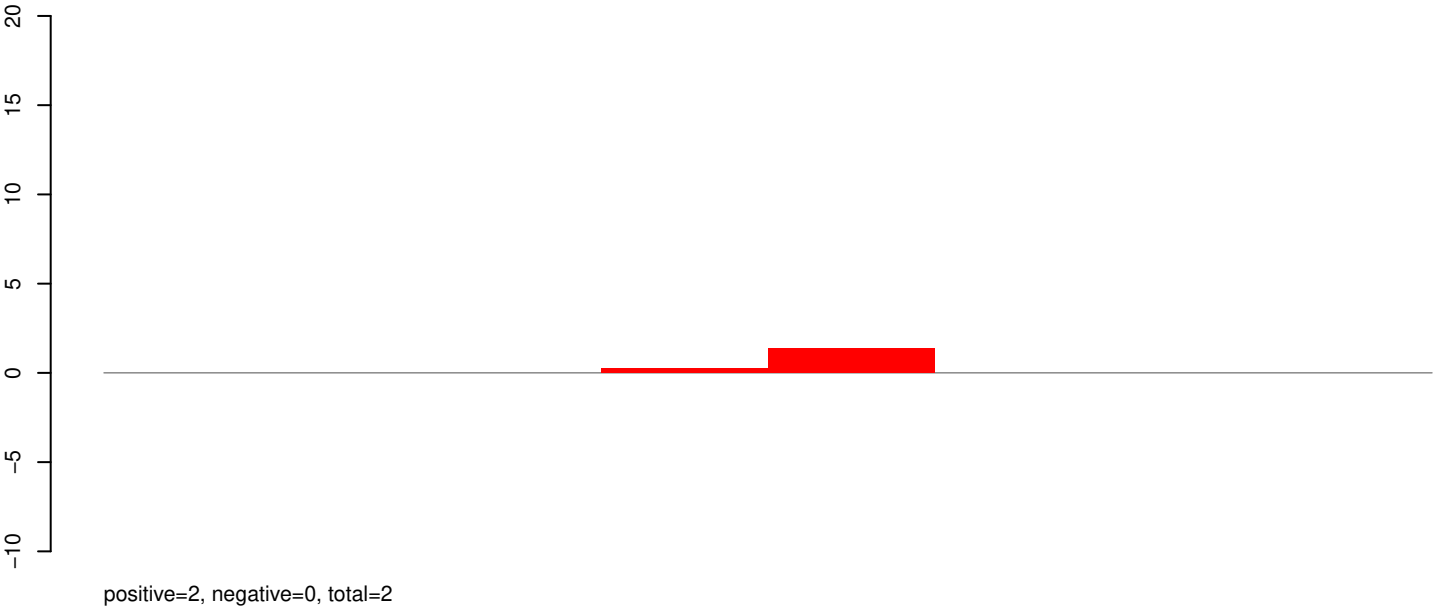
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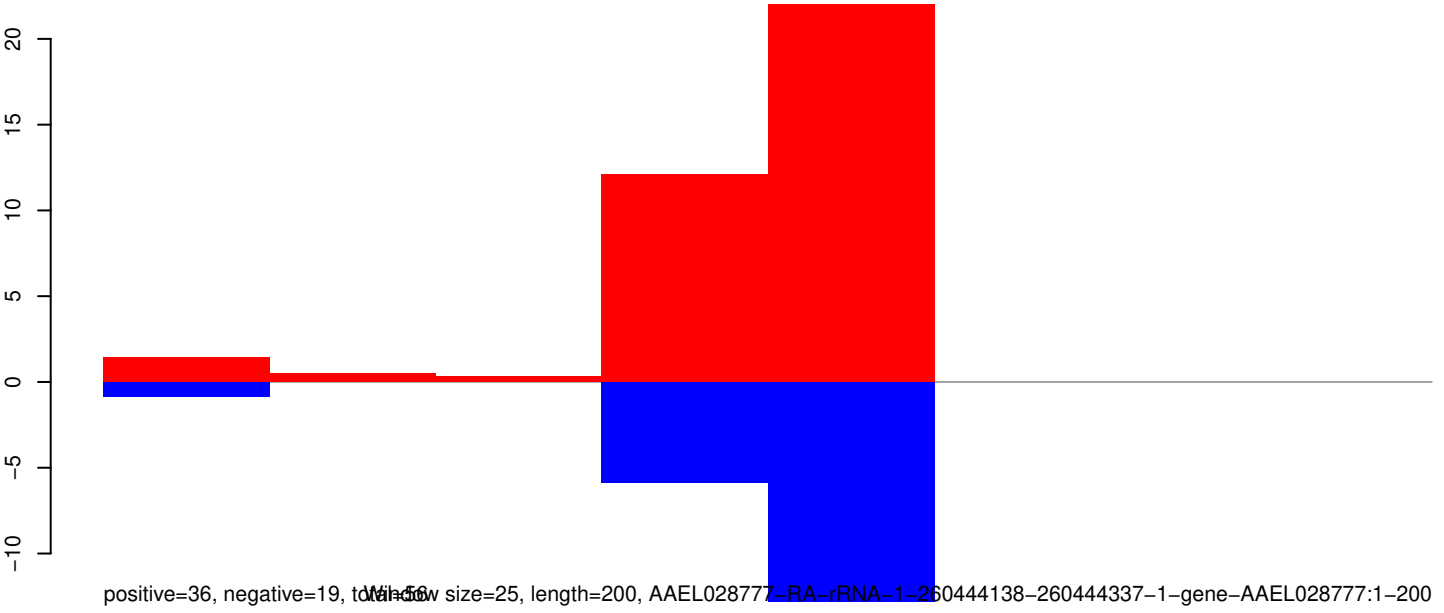
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AeAeg_Aag2SINV_dsPiwi5.24_35.rep



AeAeg_Aag2SINV_dsPiwi5.rep



Window size=25, length=200, AAEL028777-RA-rRNA-1-260444138-260444337-1-gene-AAEL028777:1-200

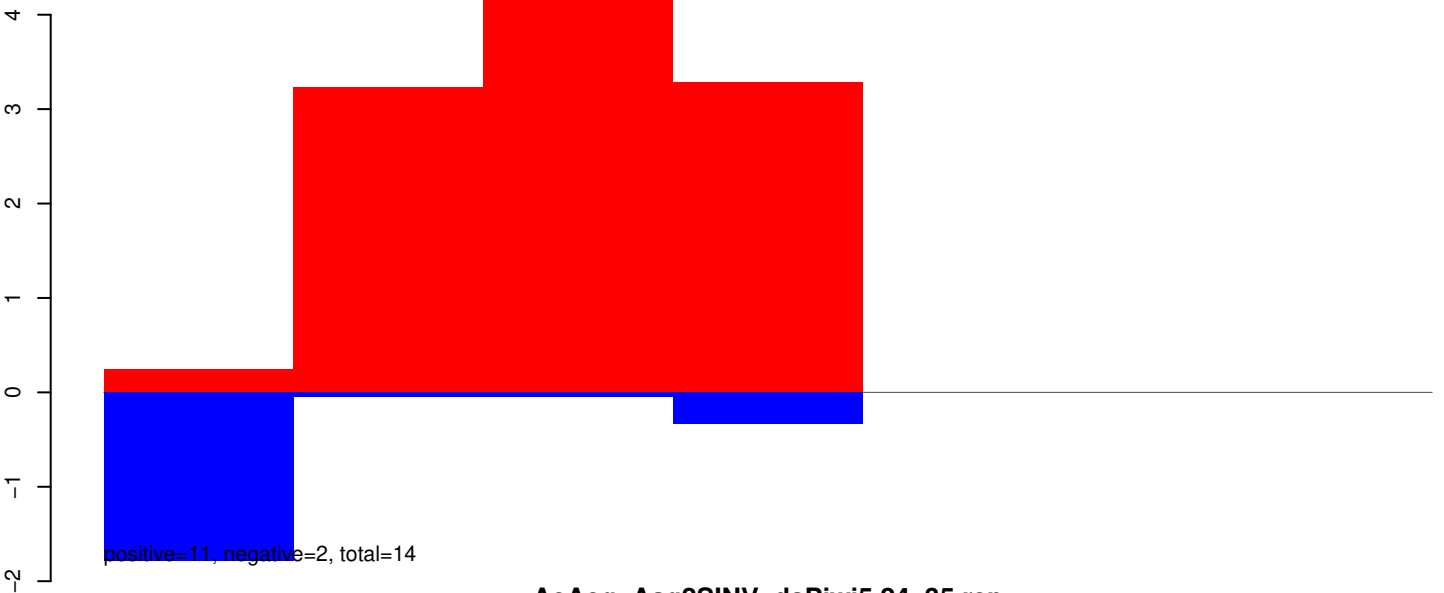
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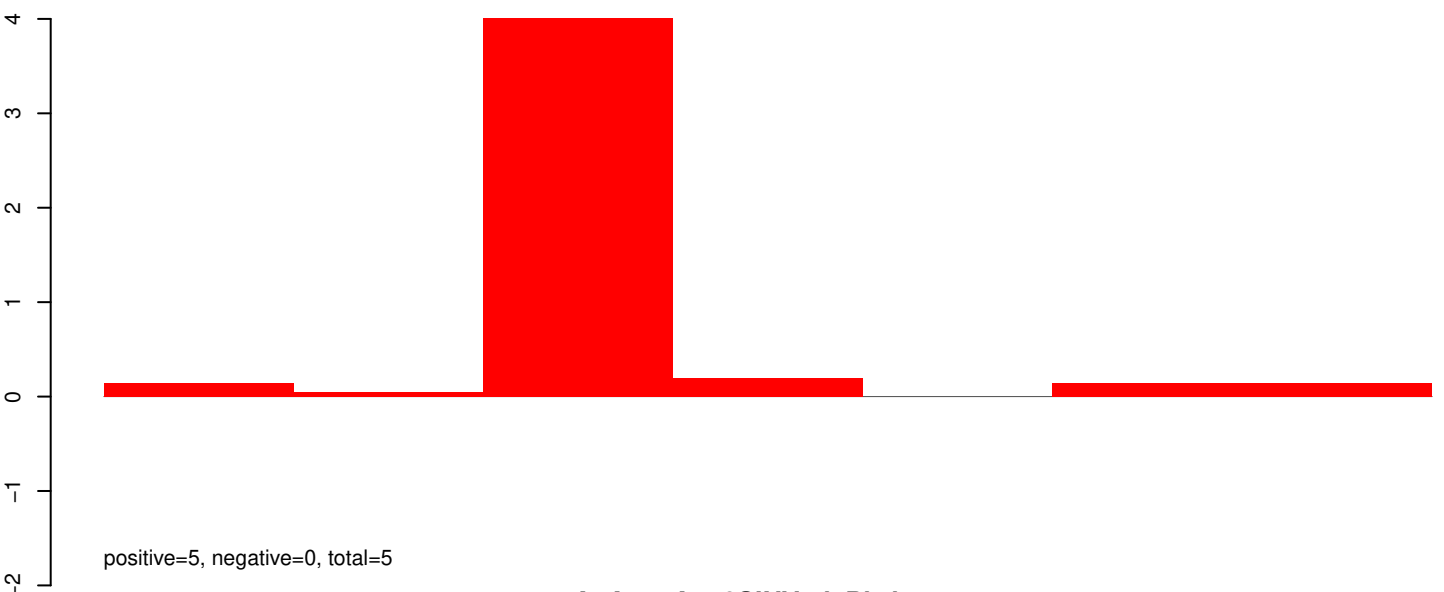
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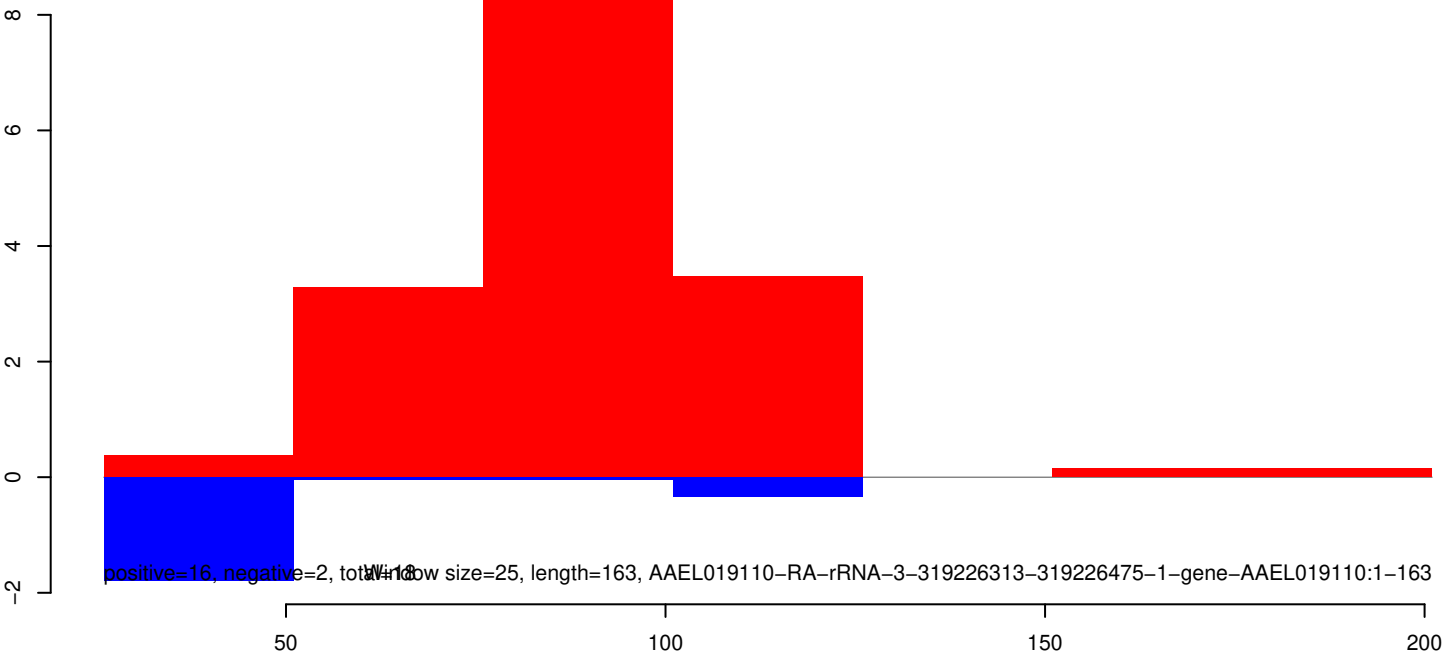
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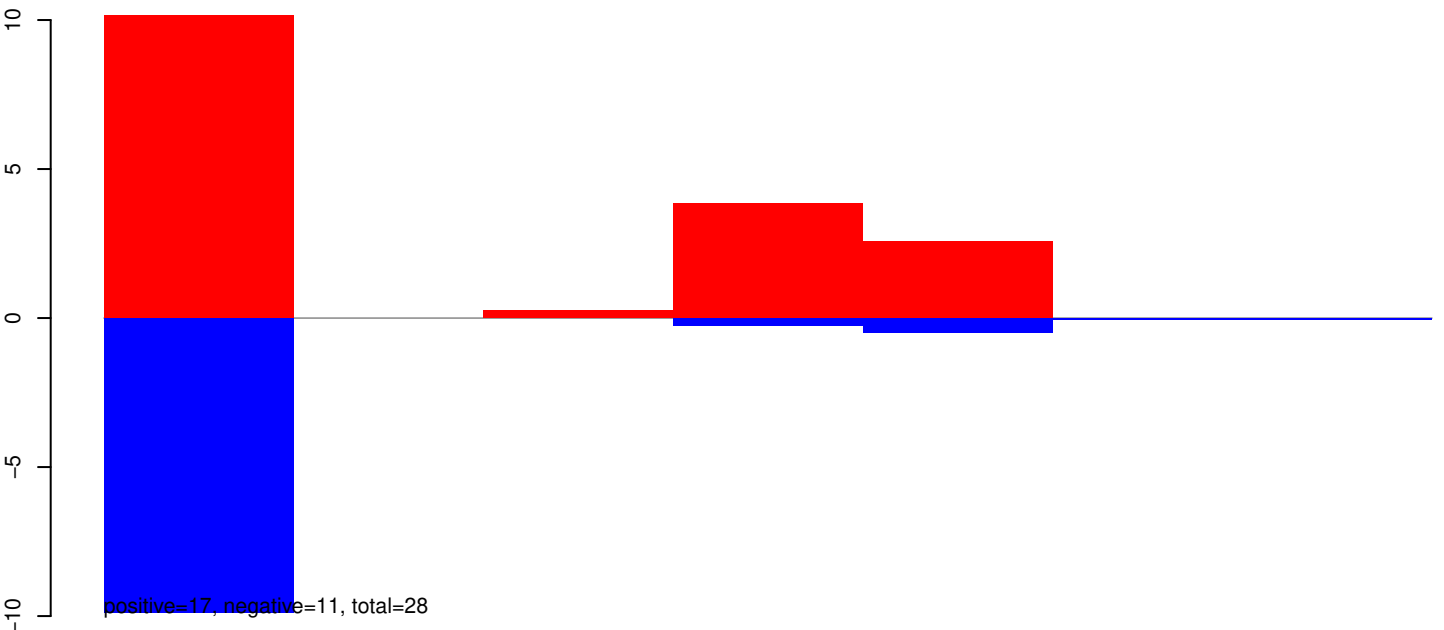


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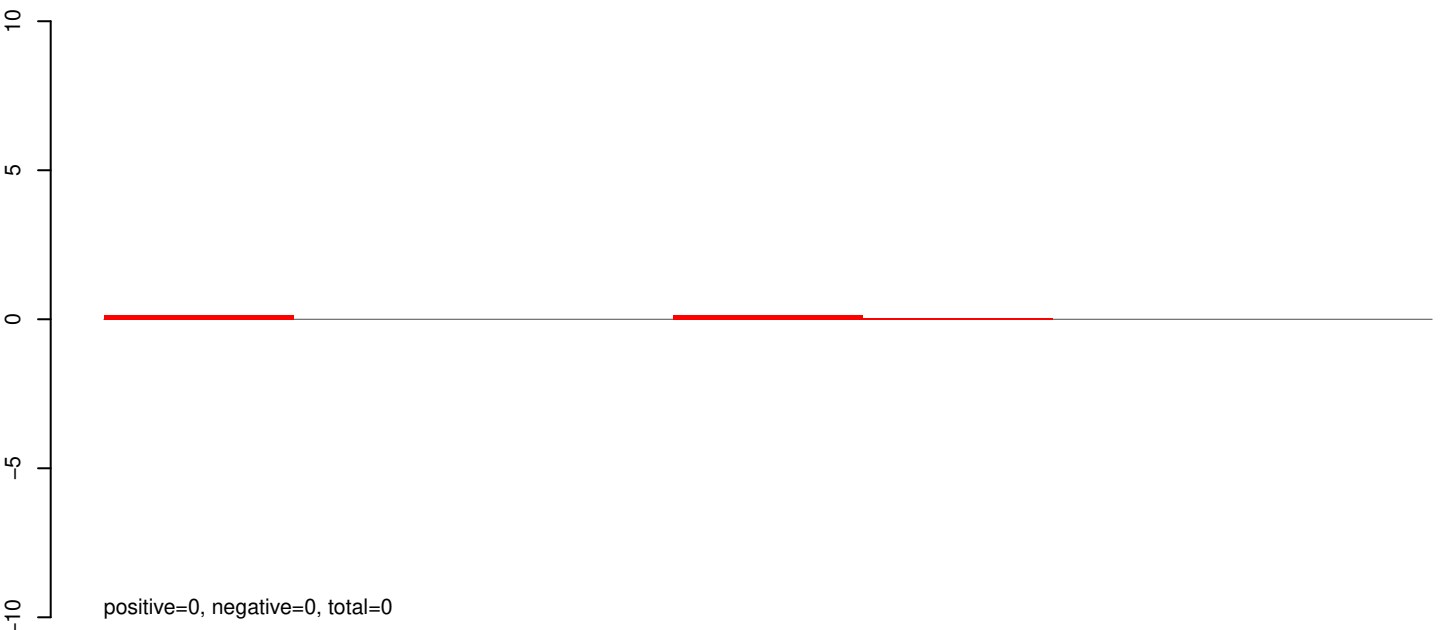


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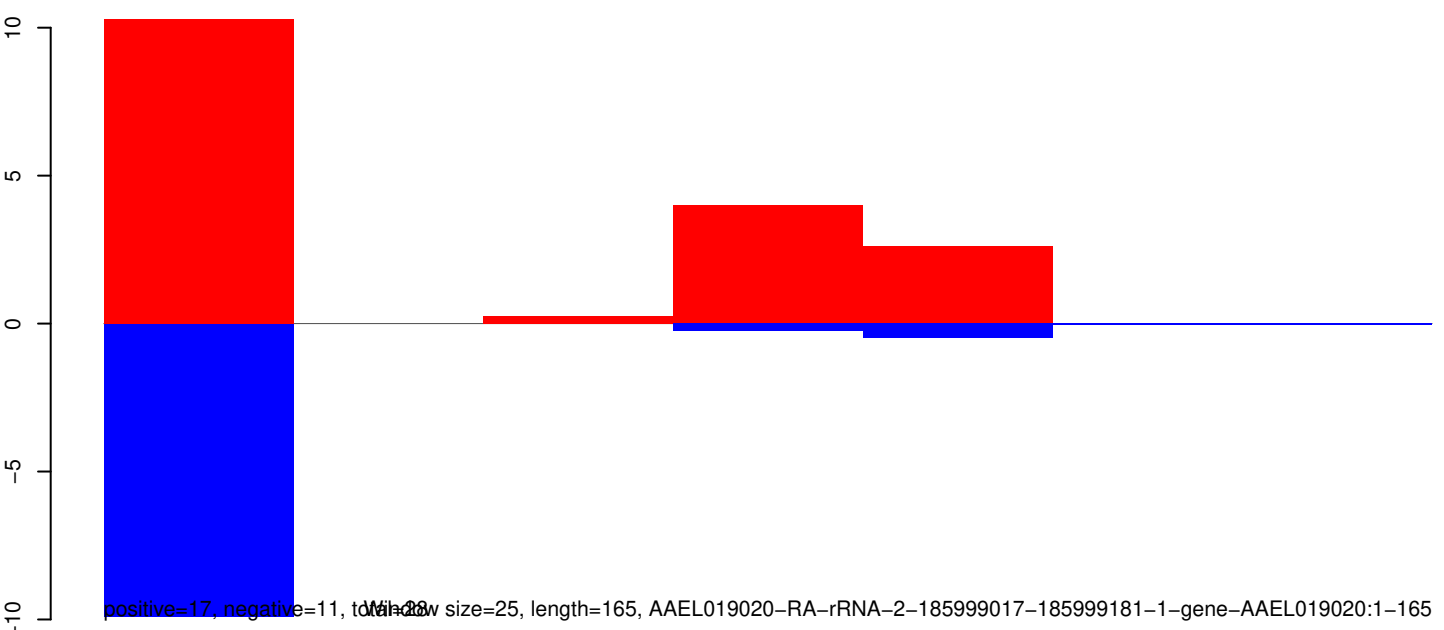
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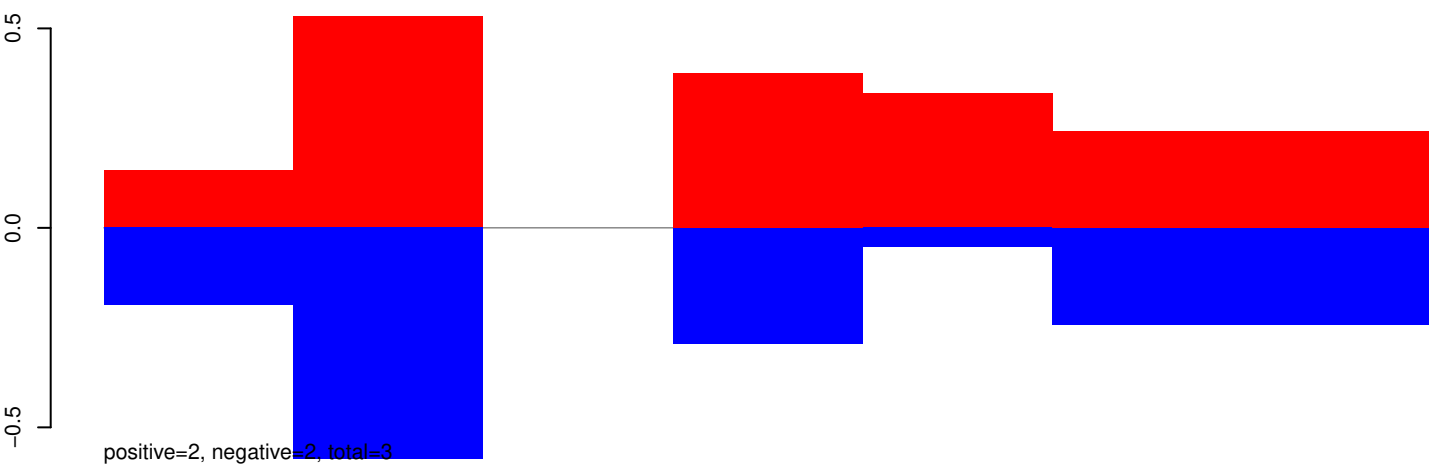
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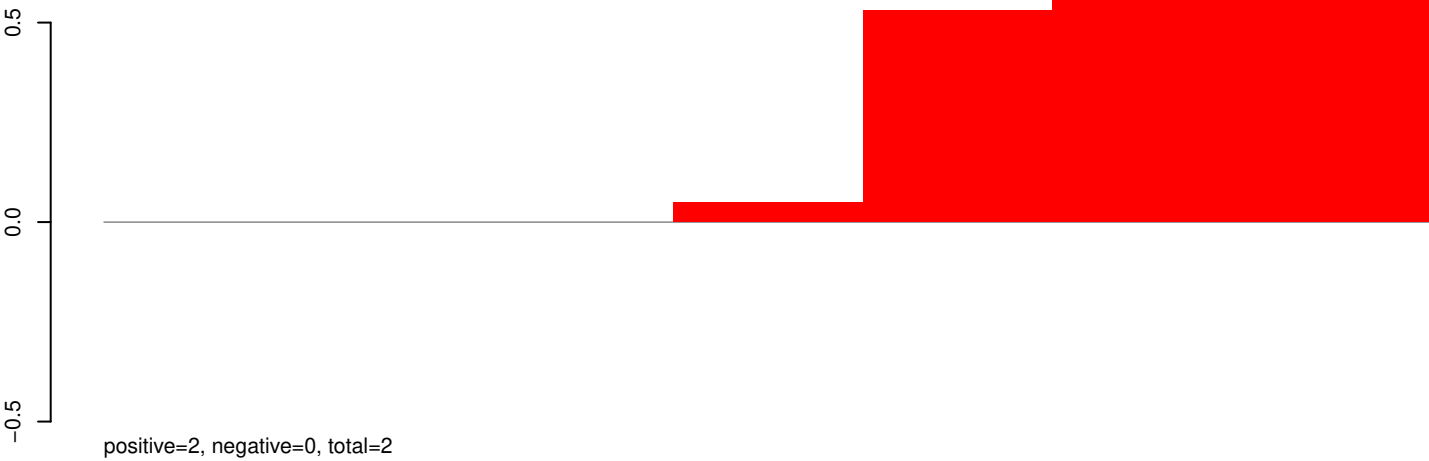
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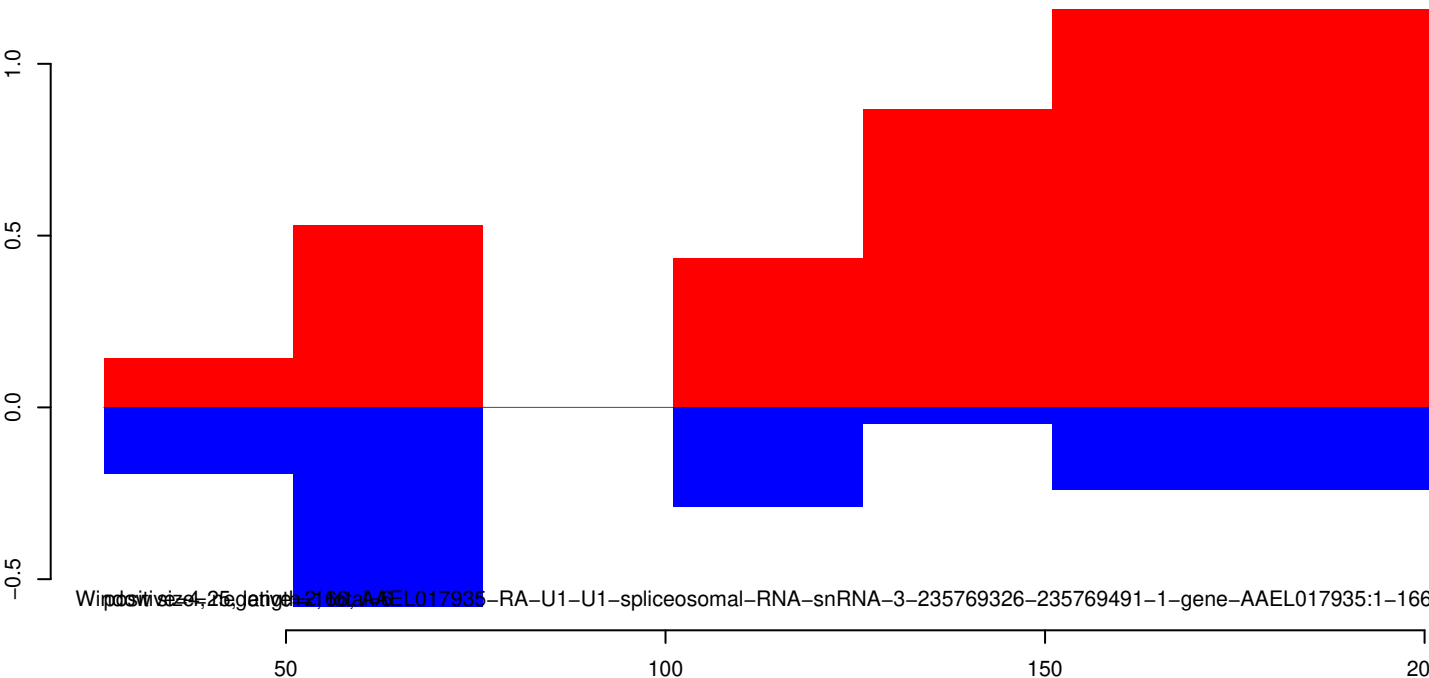
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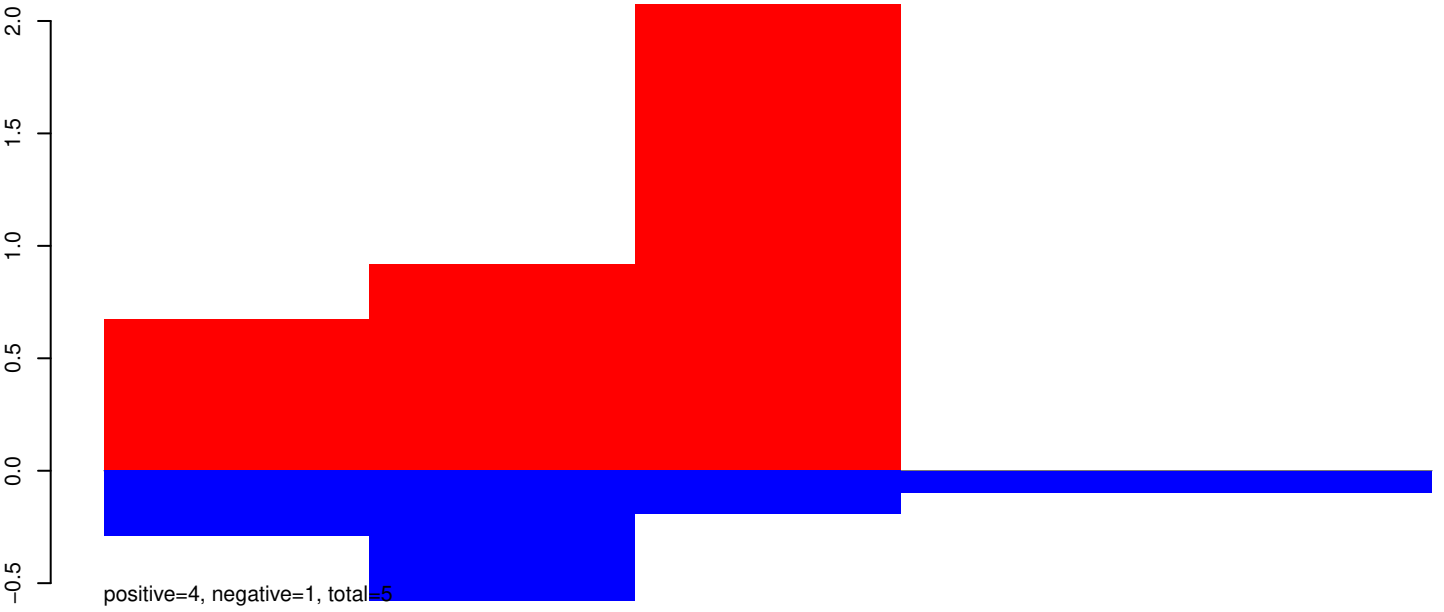
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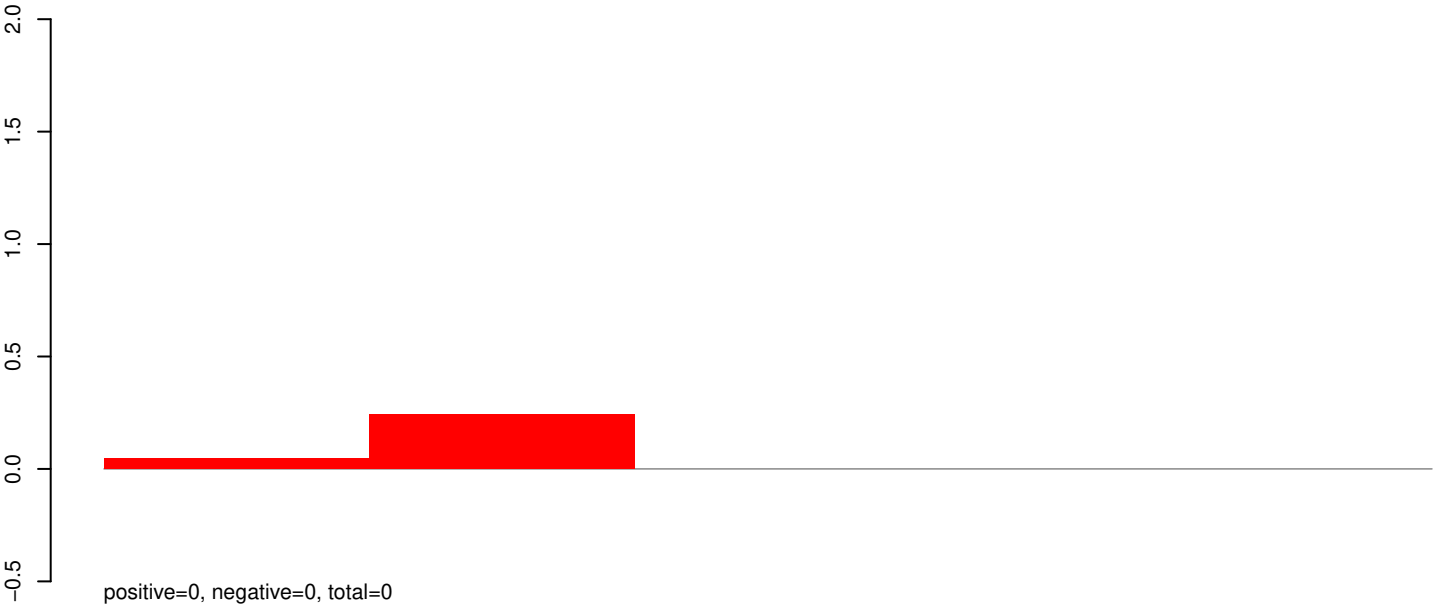
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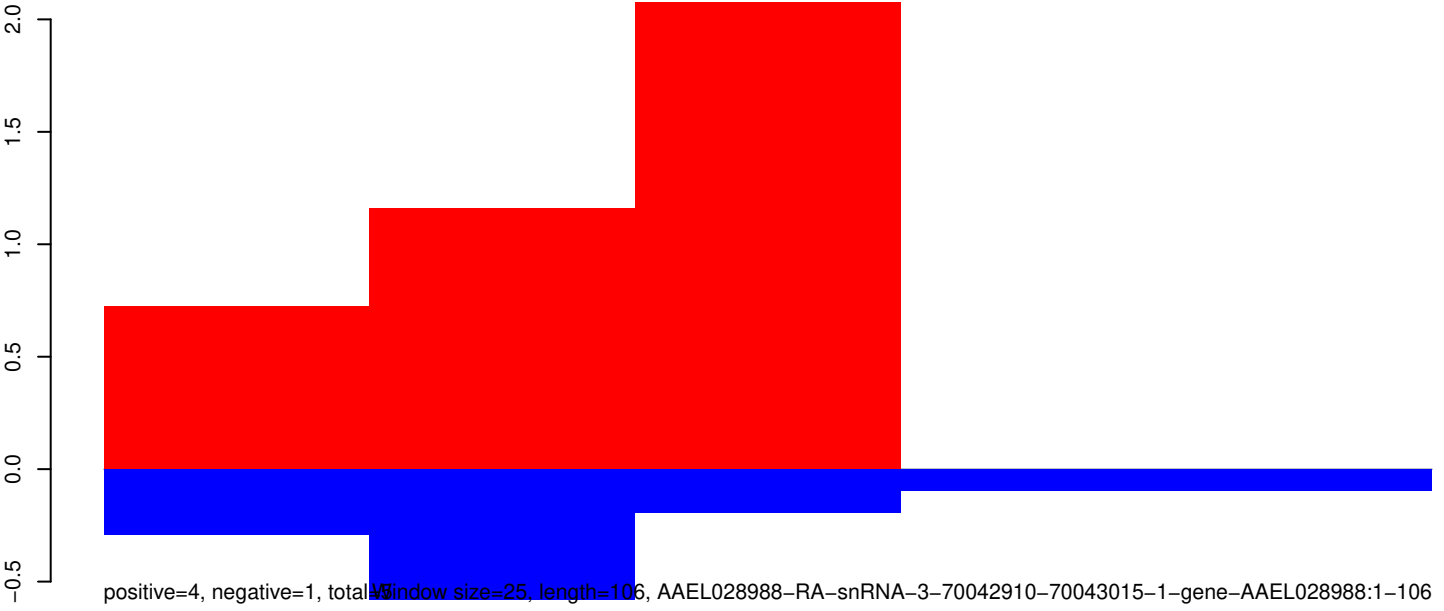
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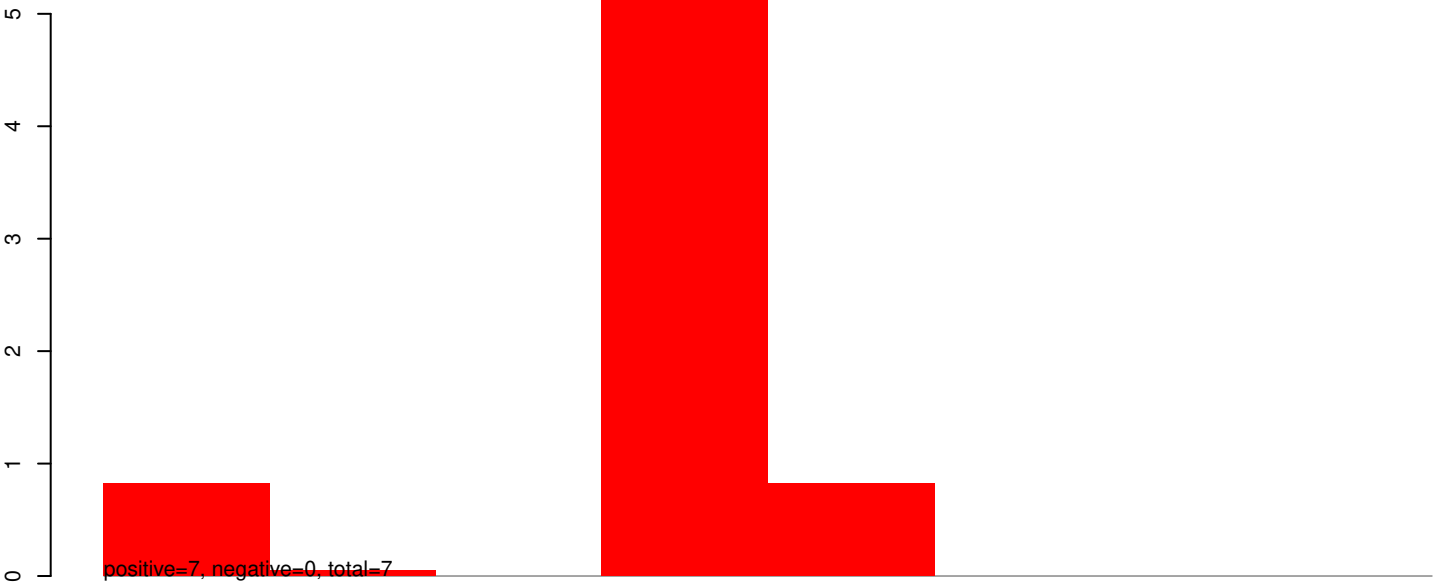
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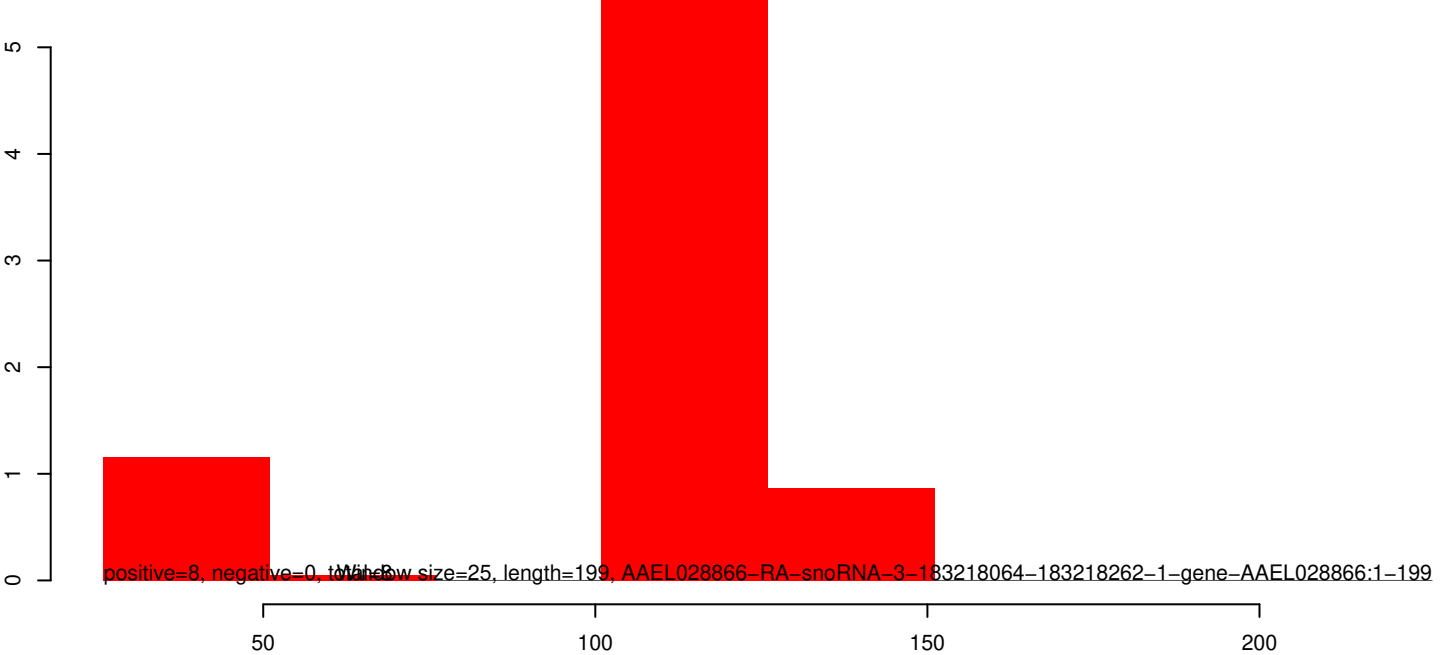
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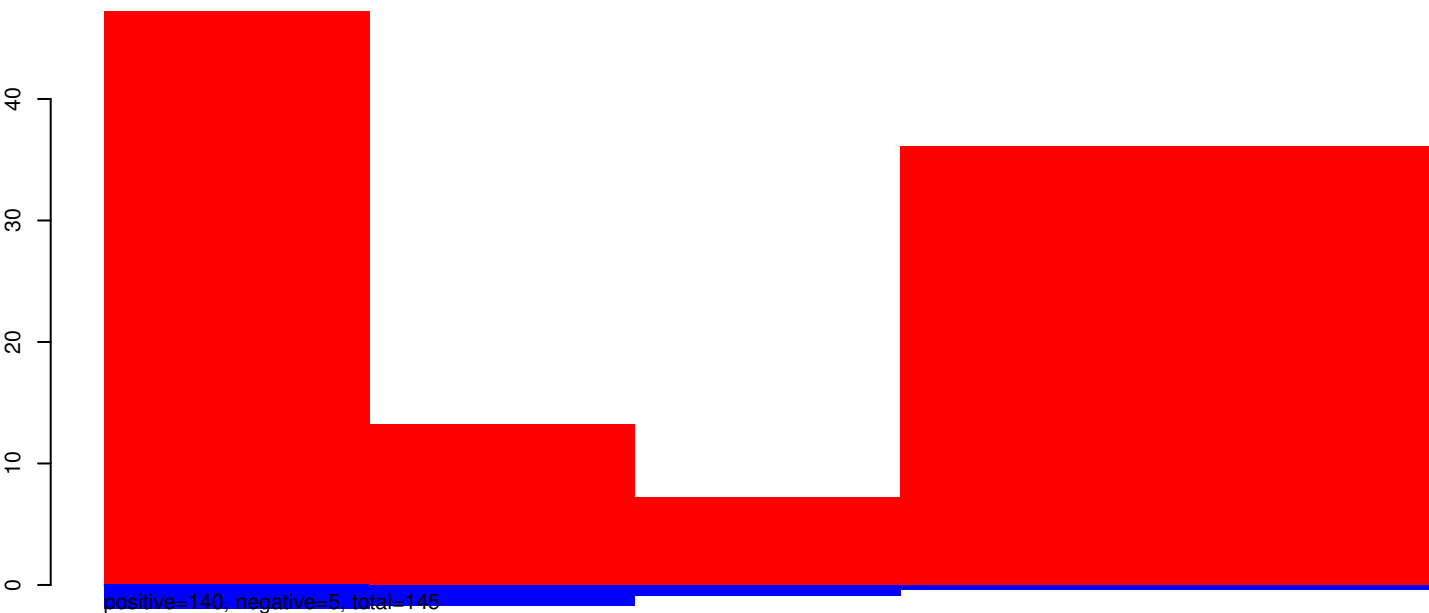
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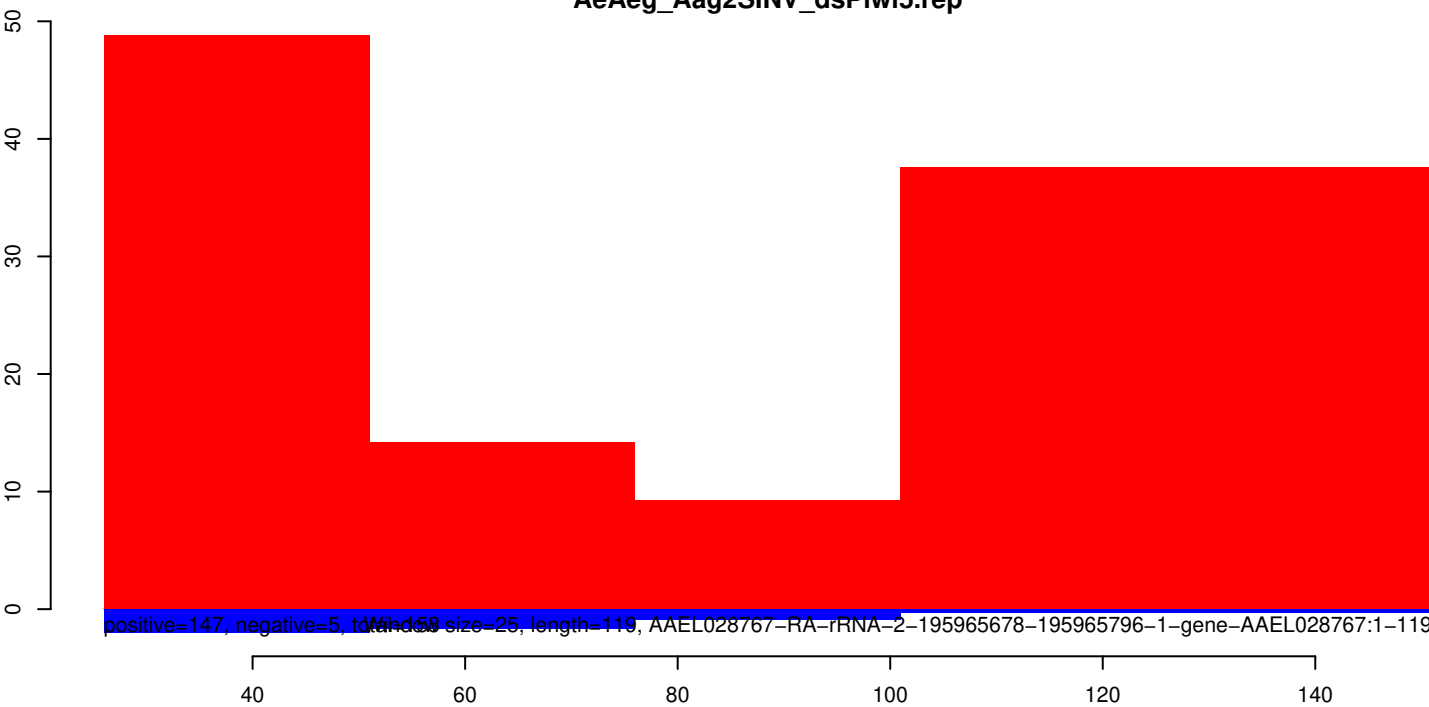
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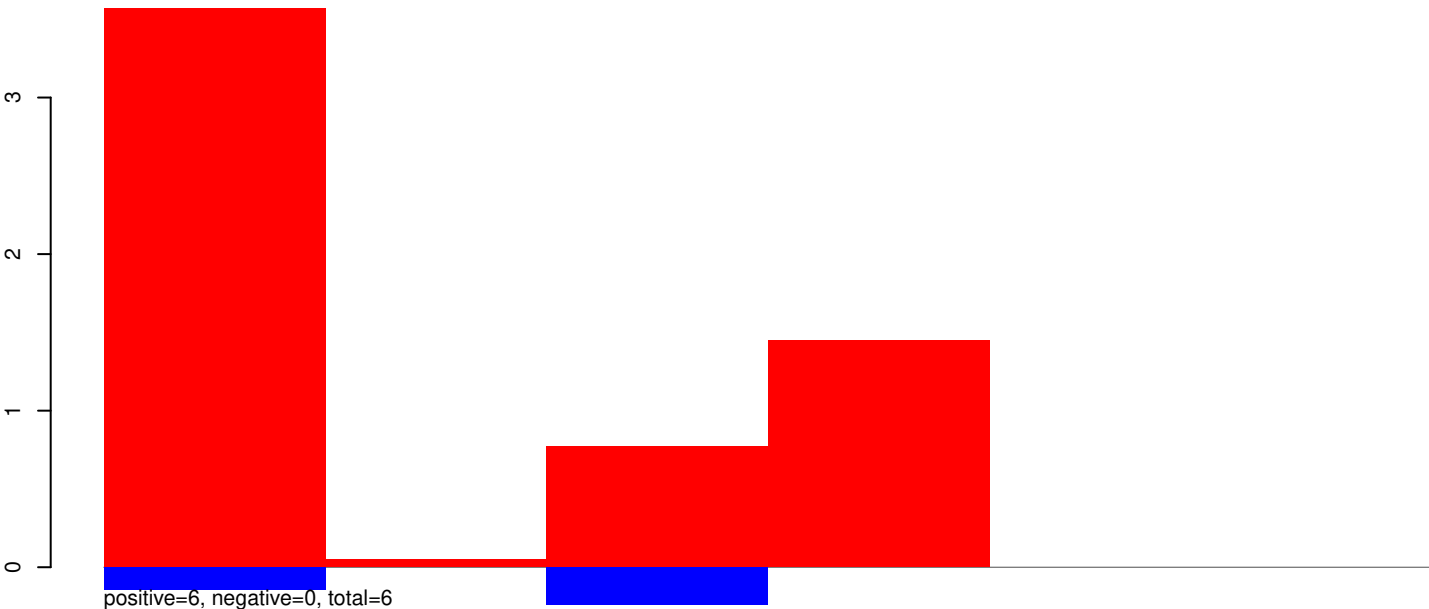
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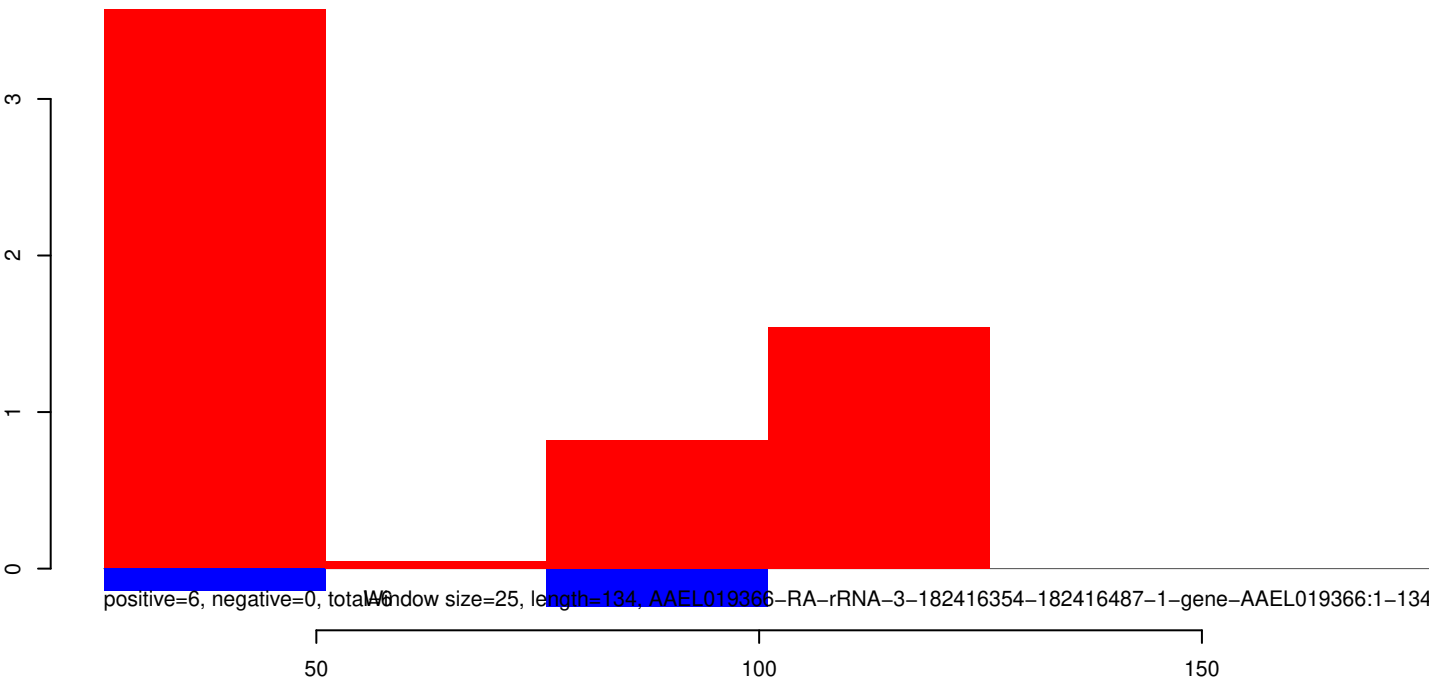
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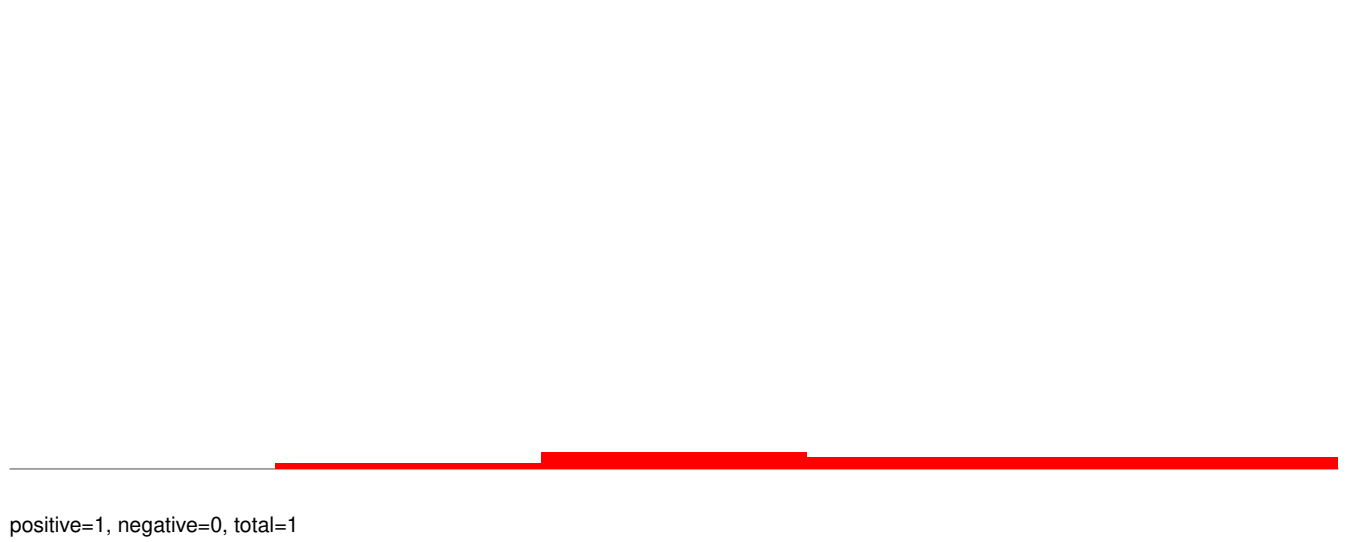
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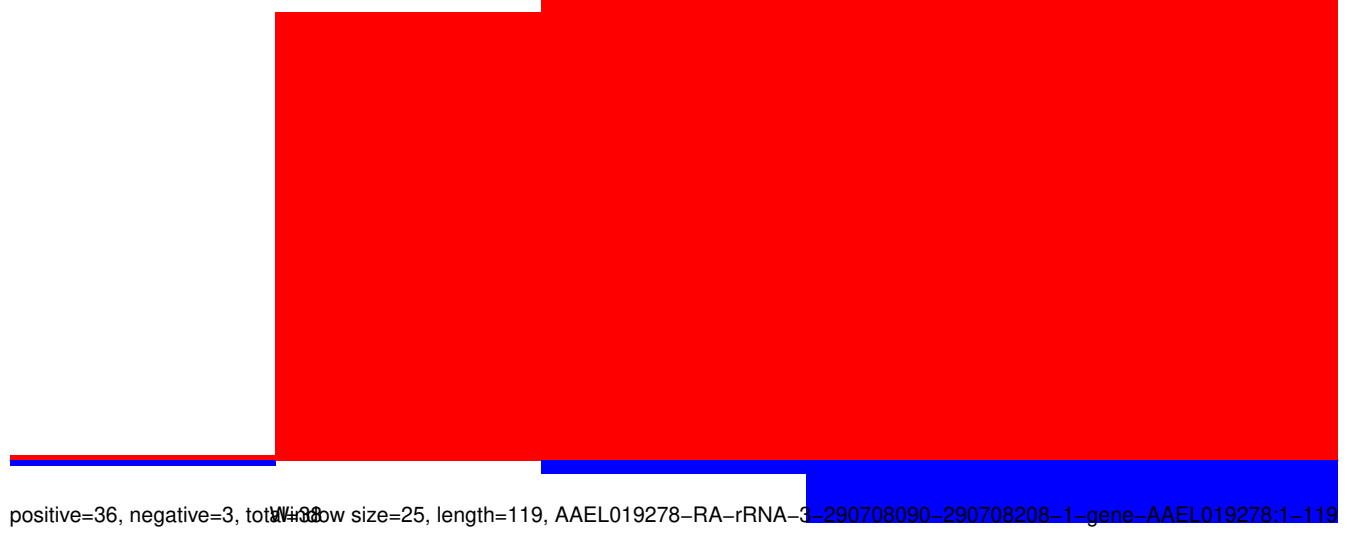
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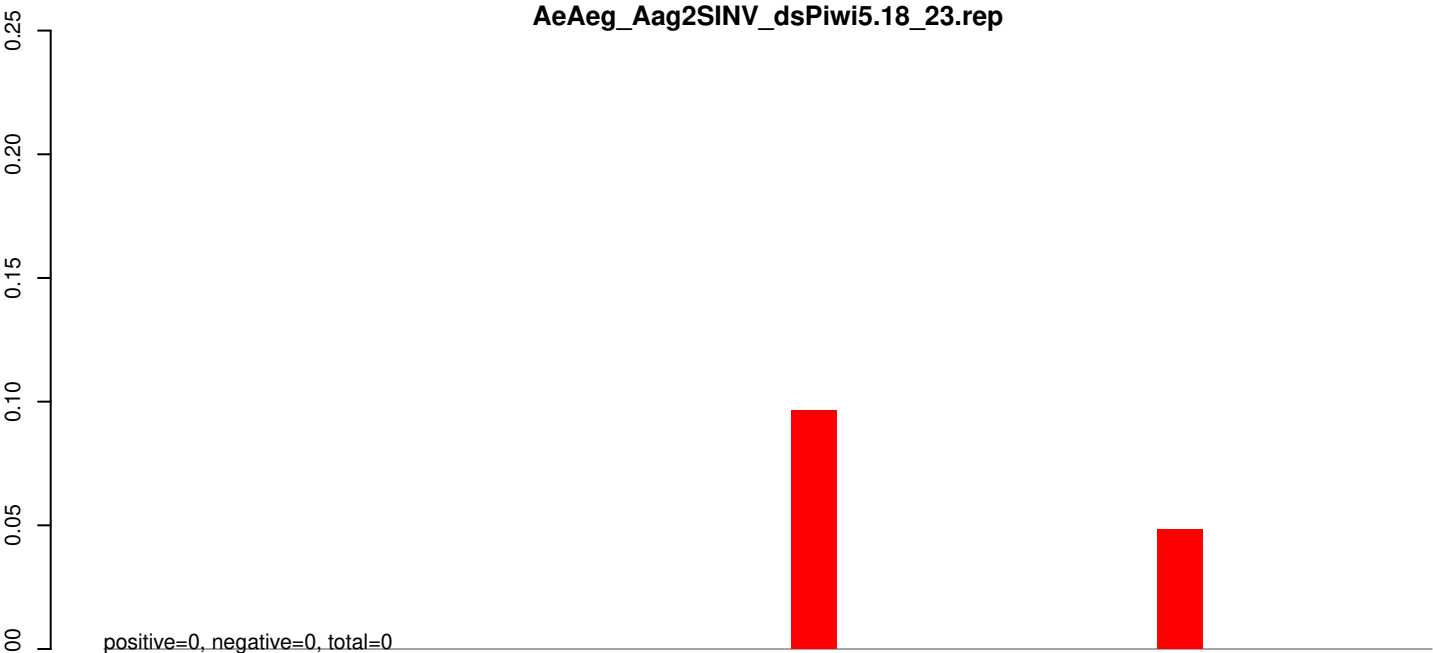
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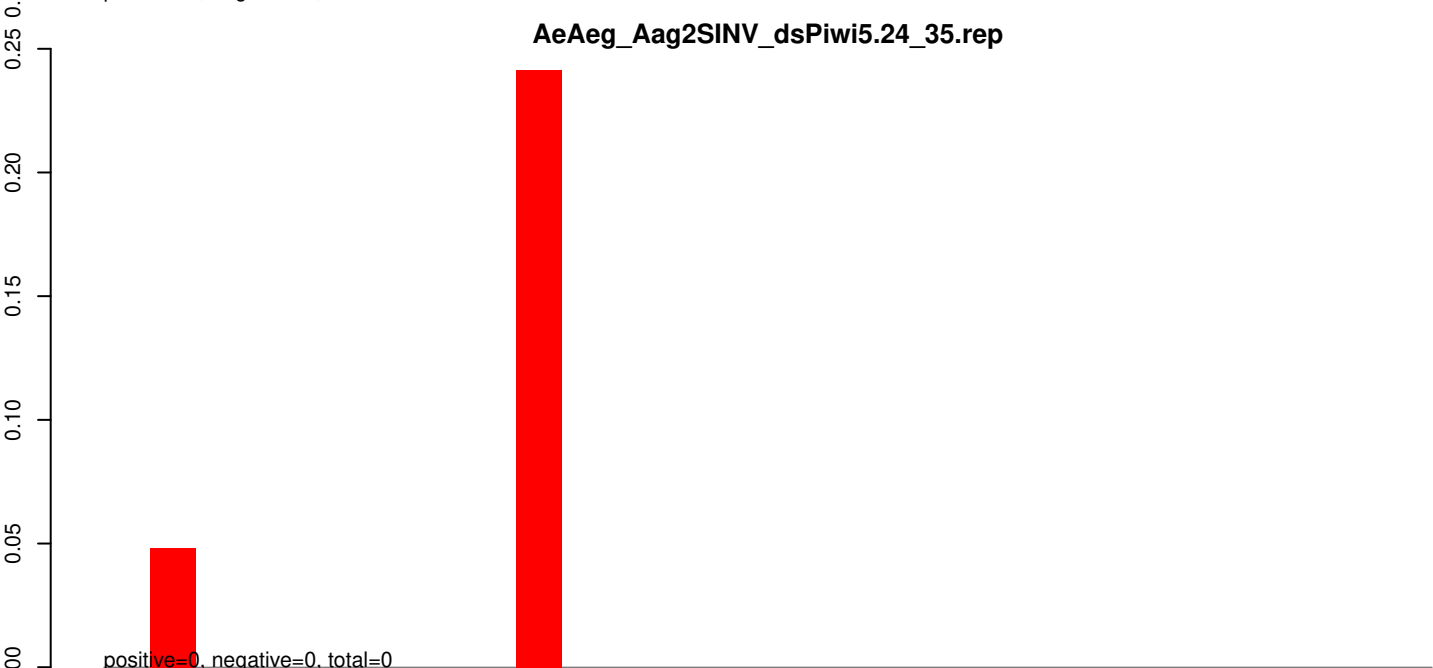
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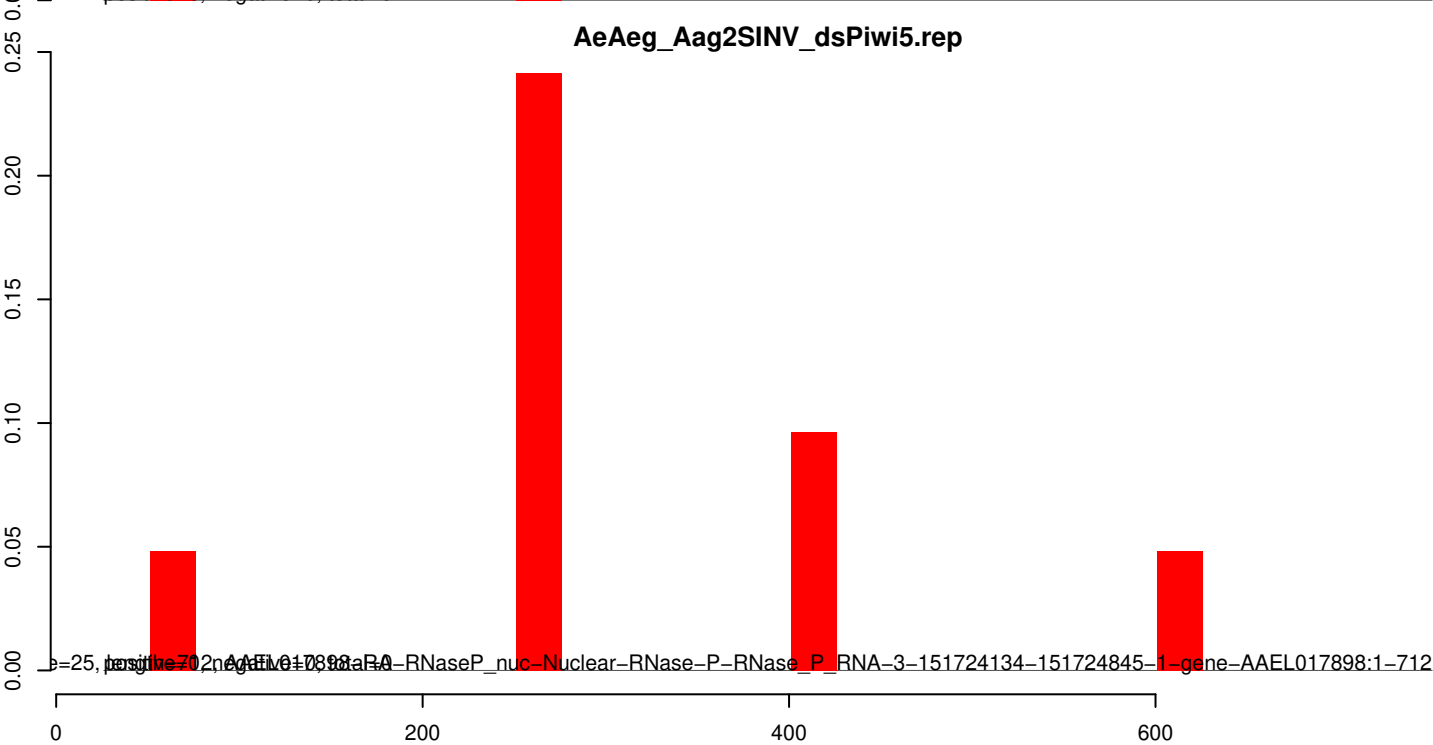
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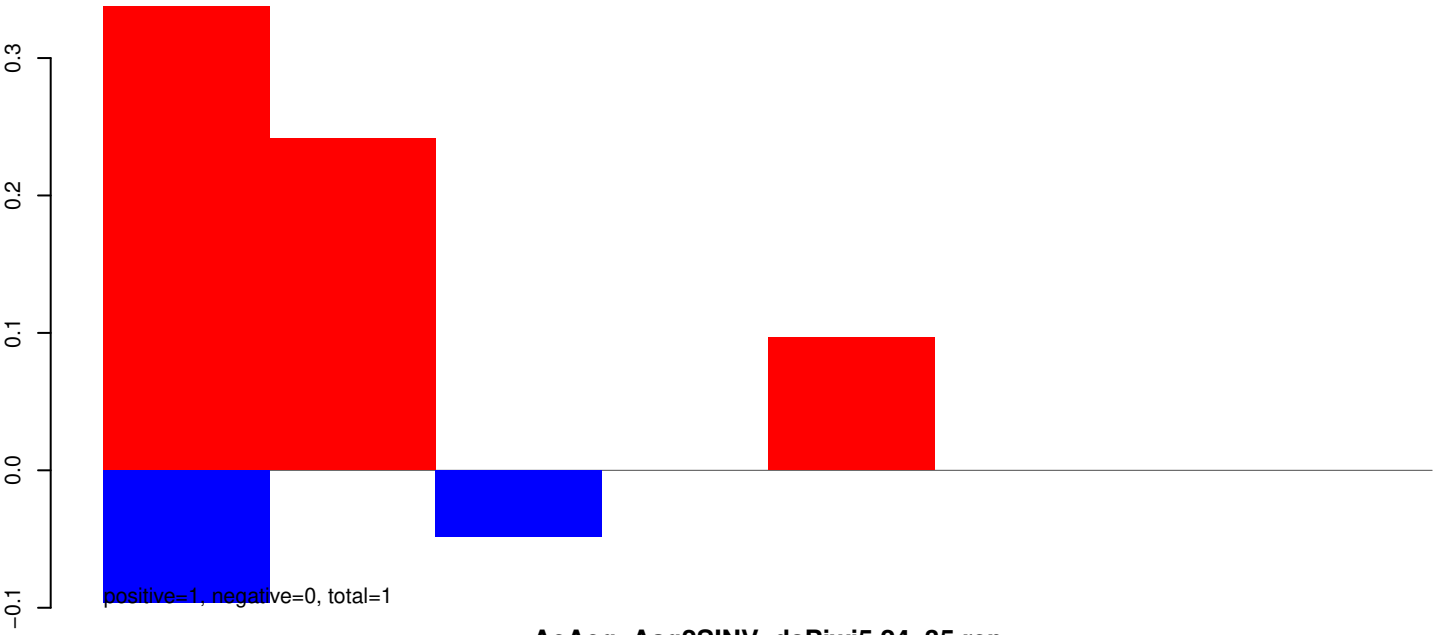
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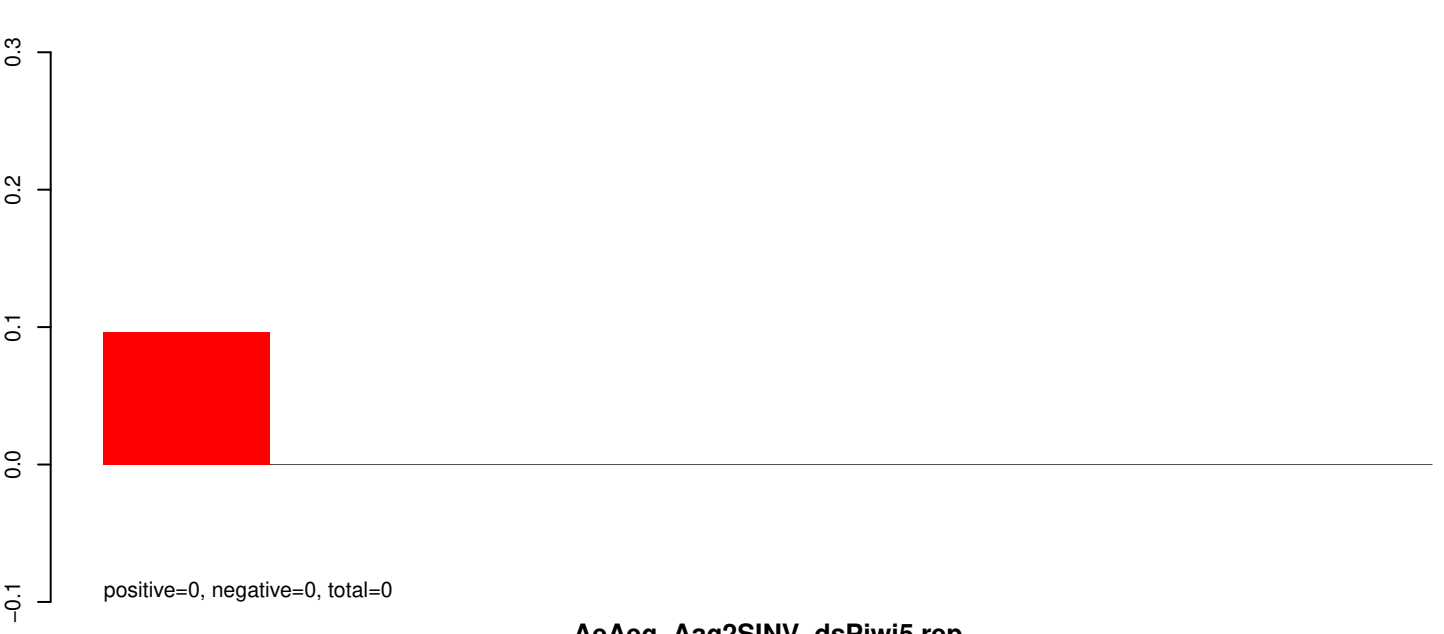
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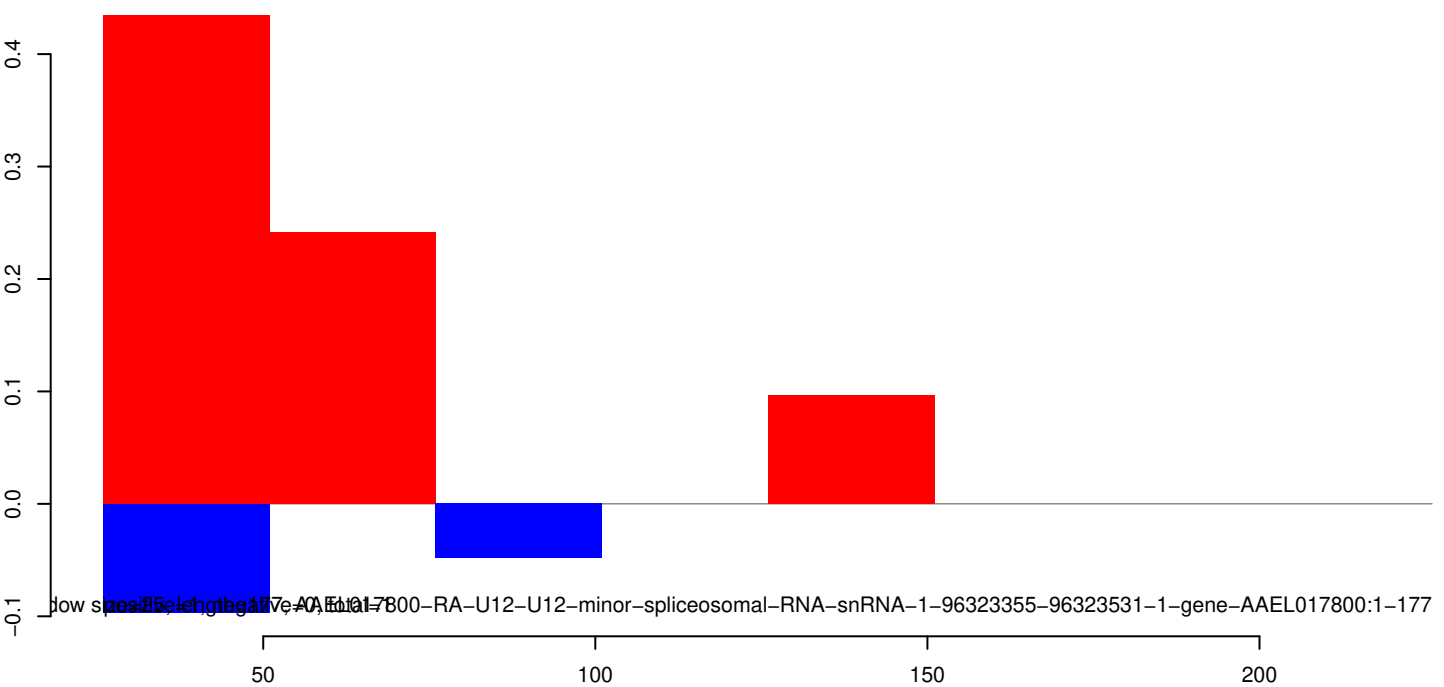
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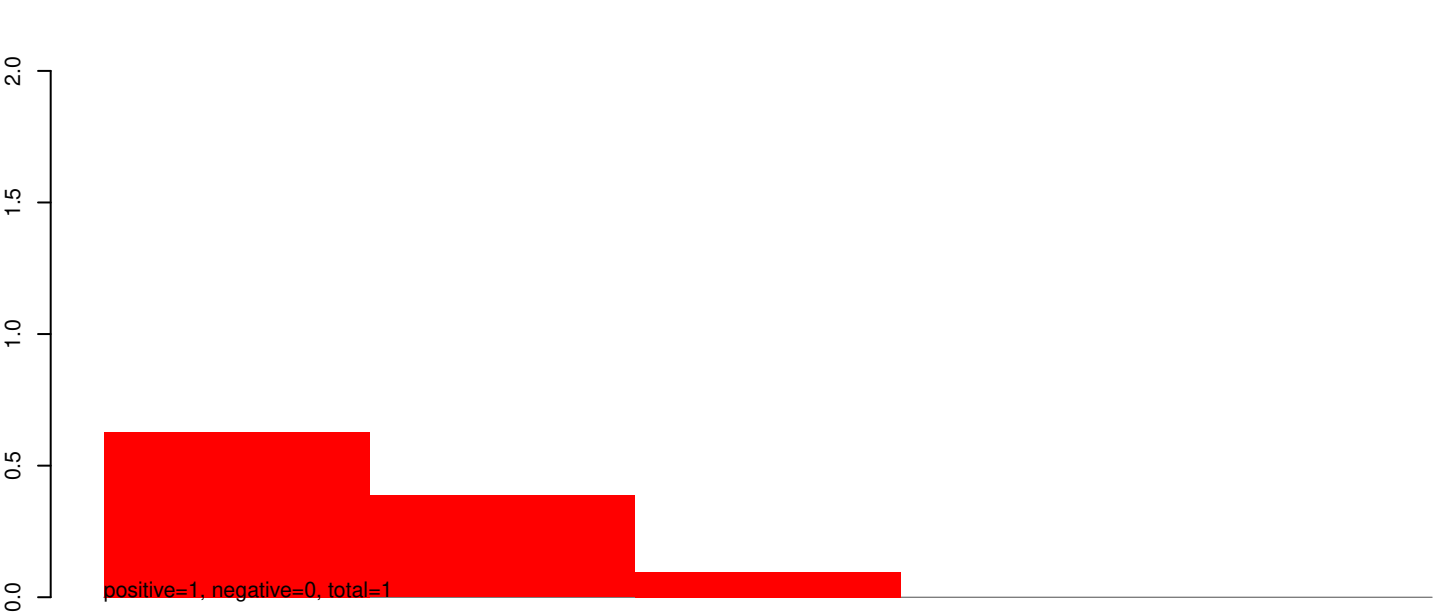
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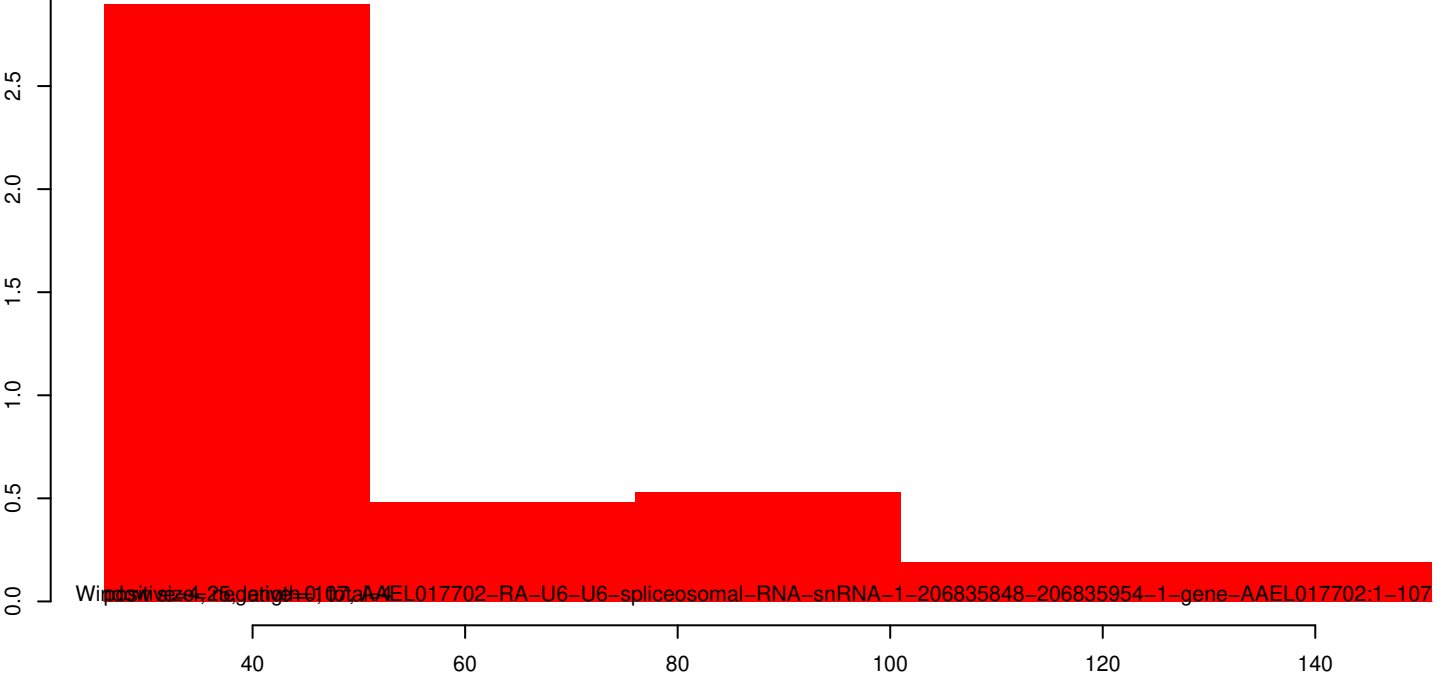
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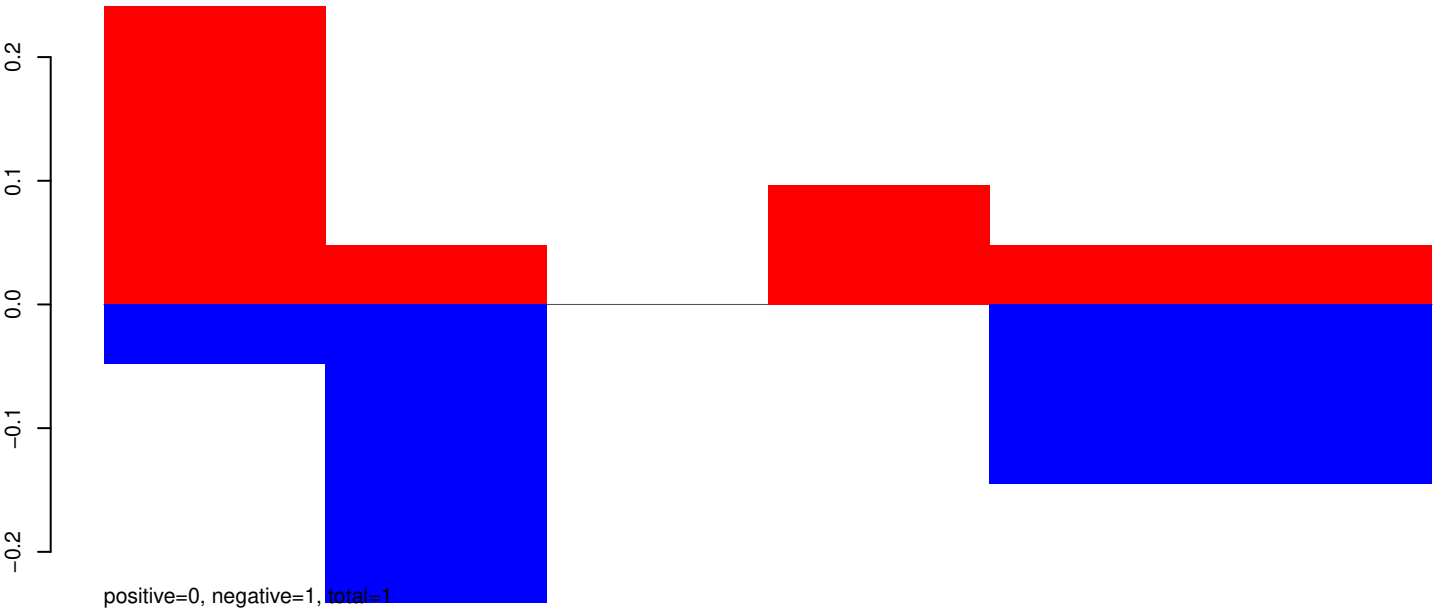
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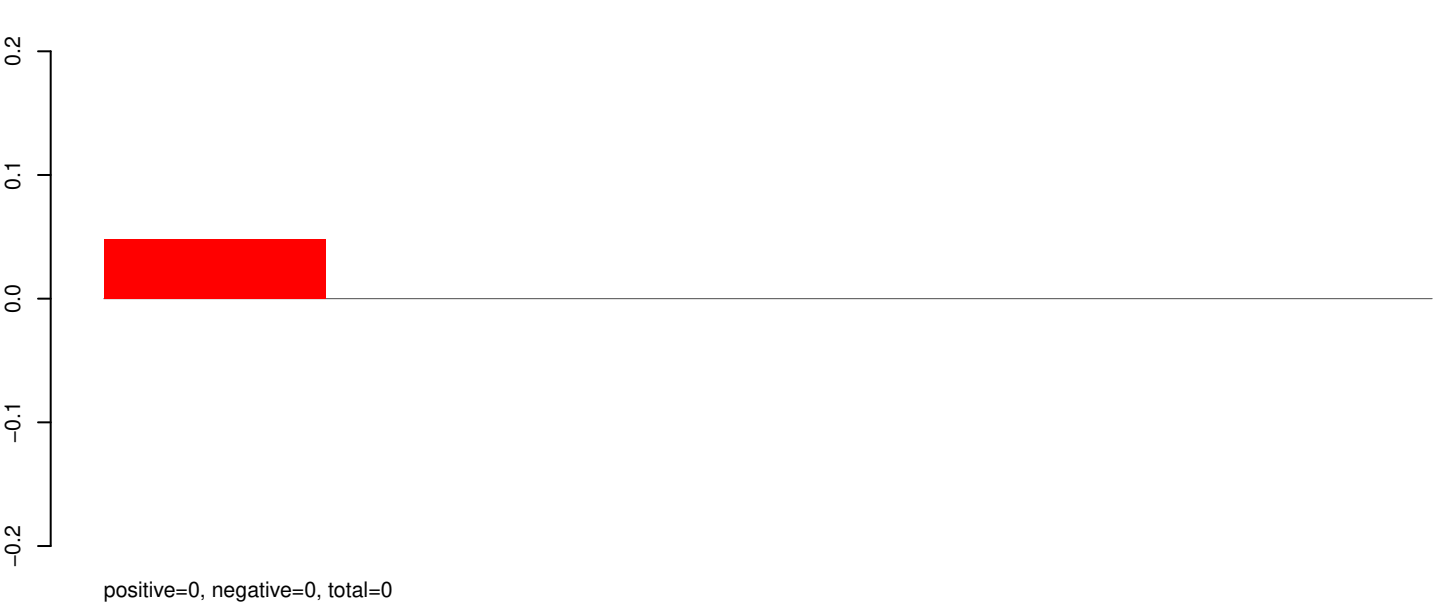
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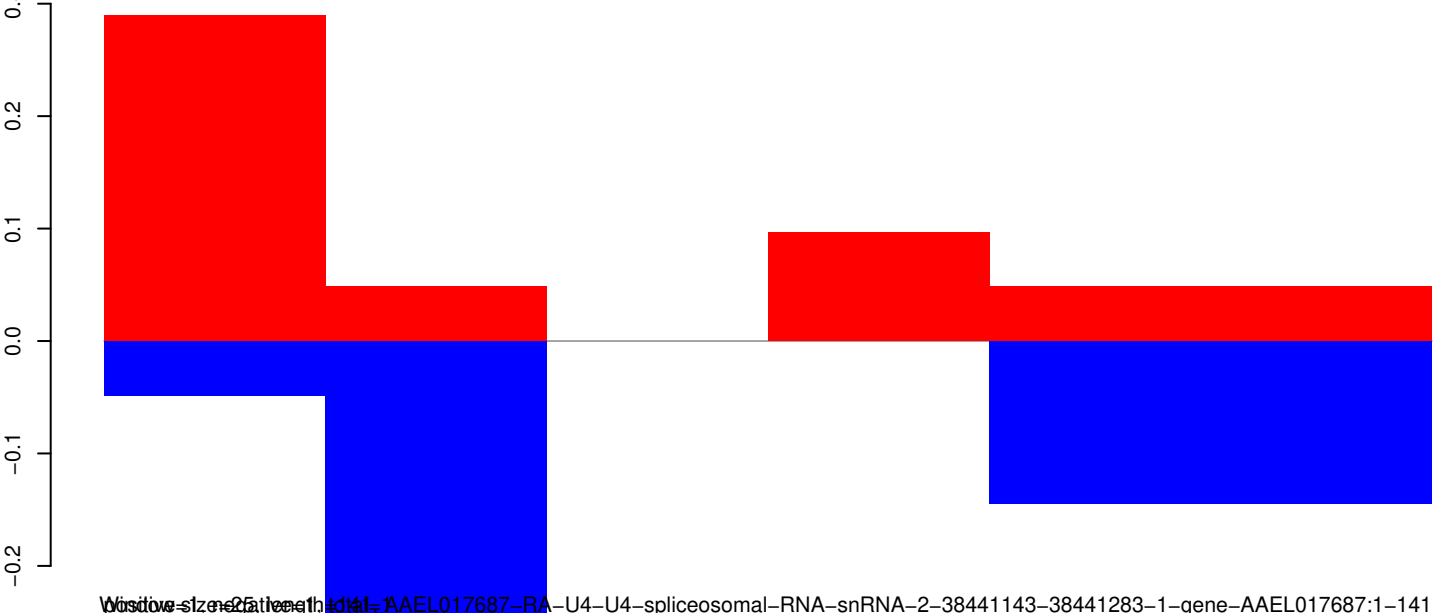
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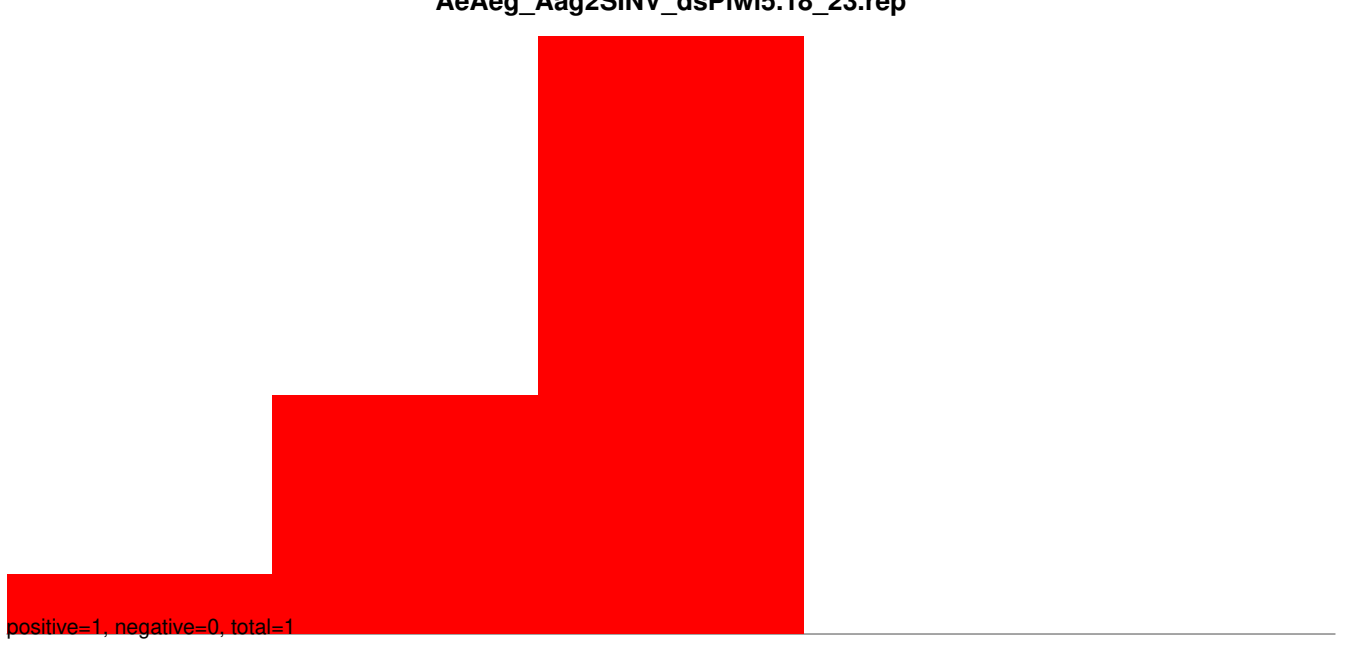
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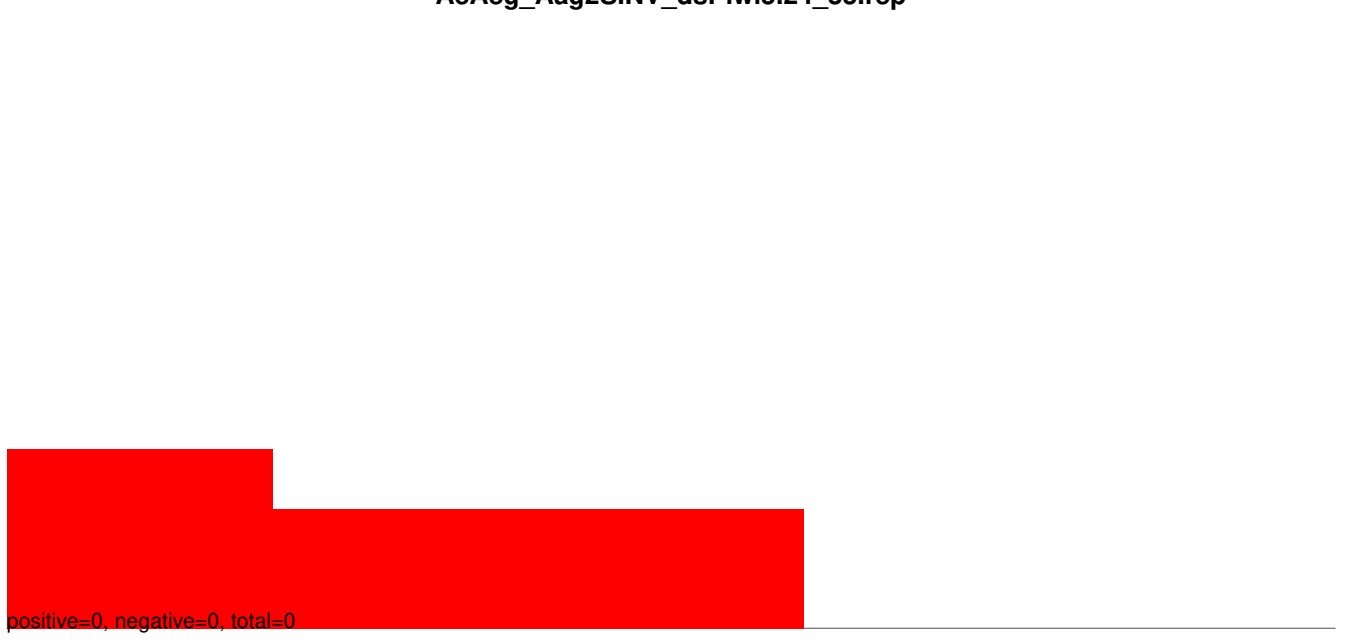
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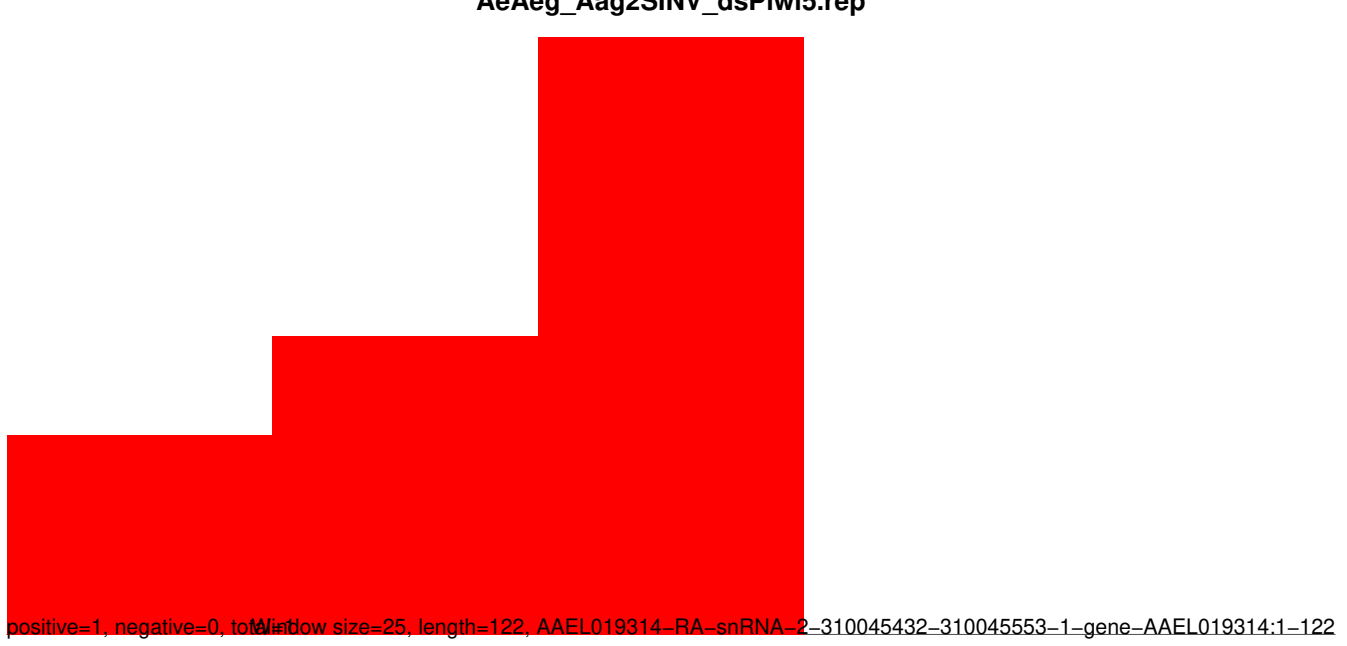
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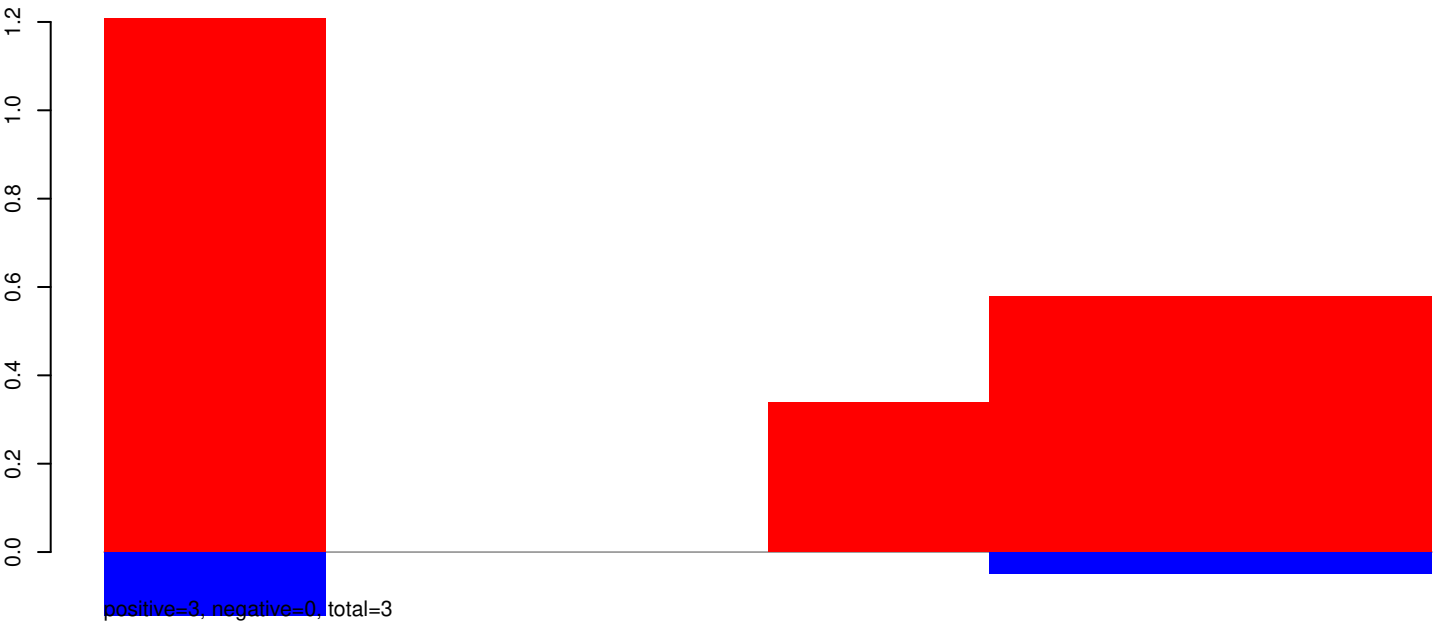


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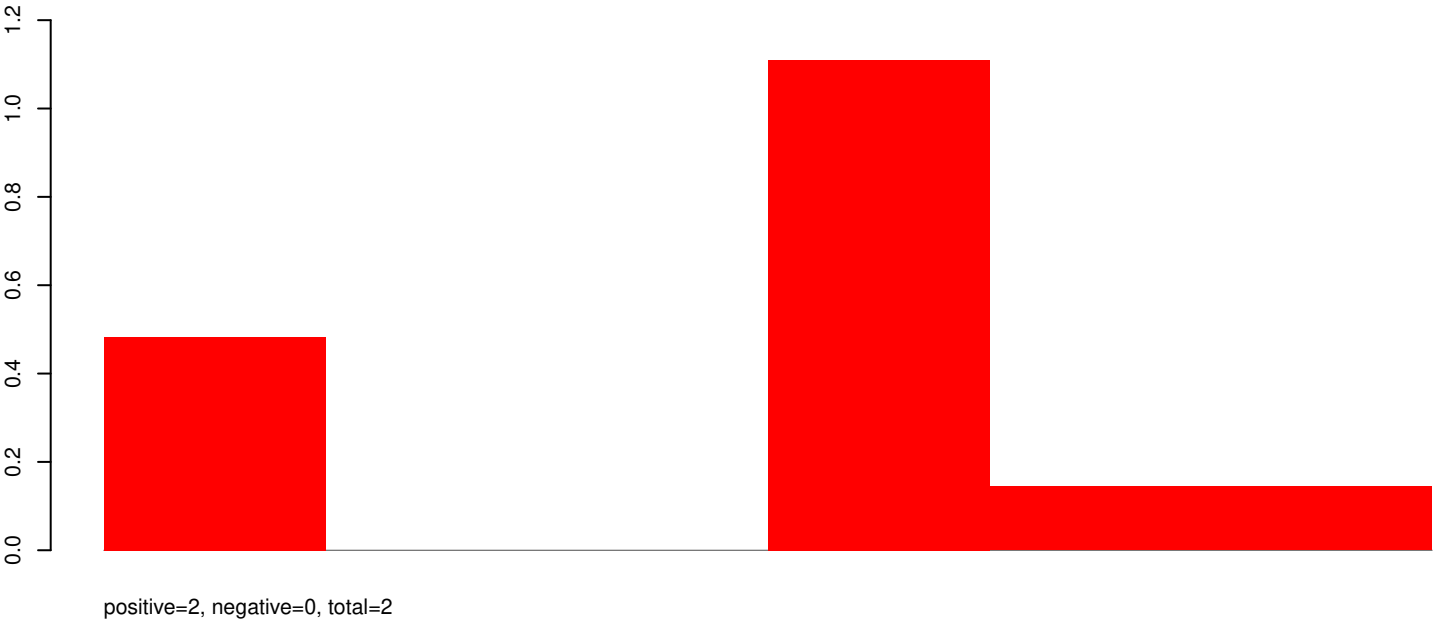


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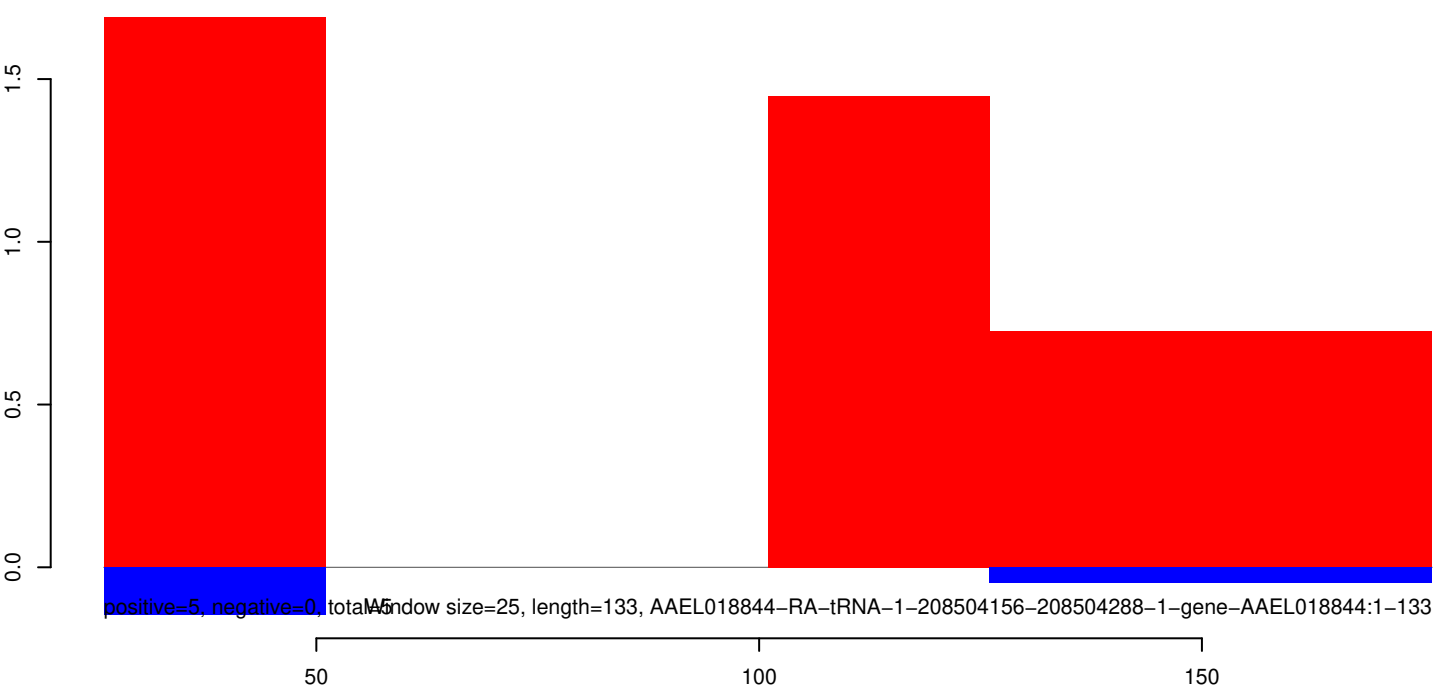
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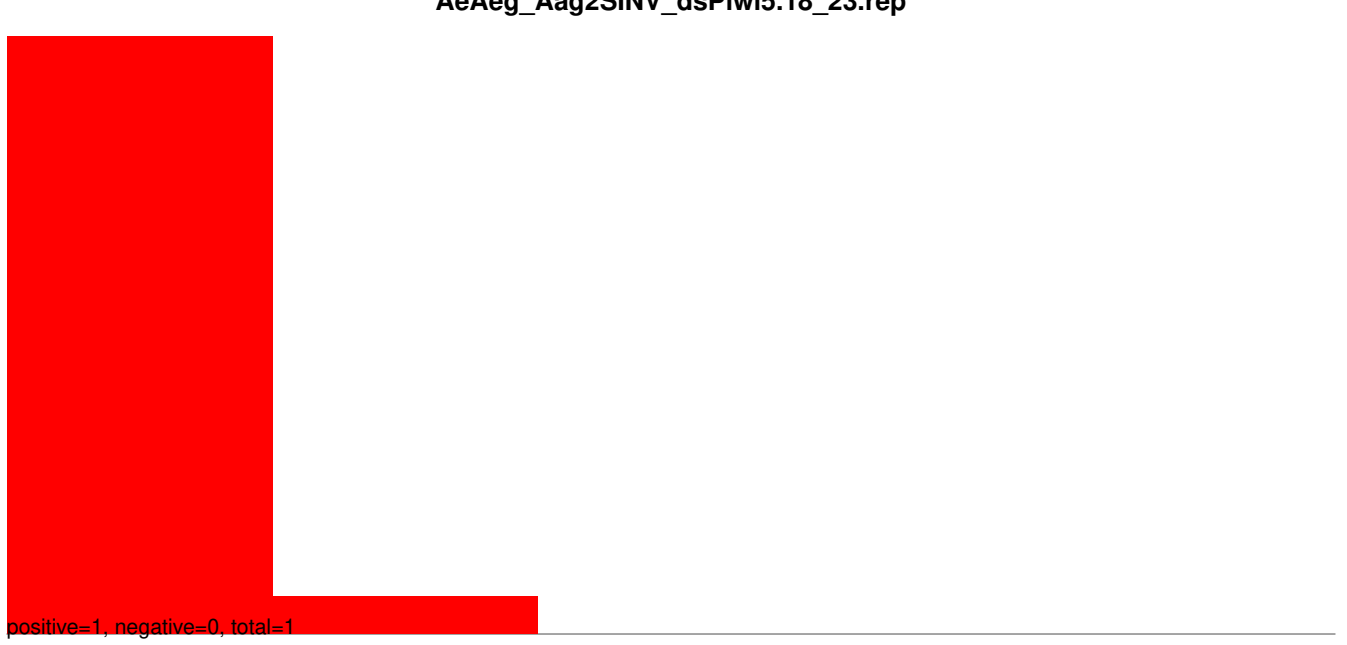
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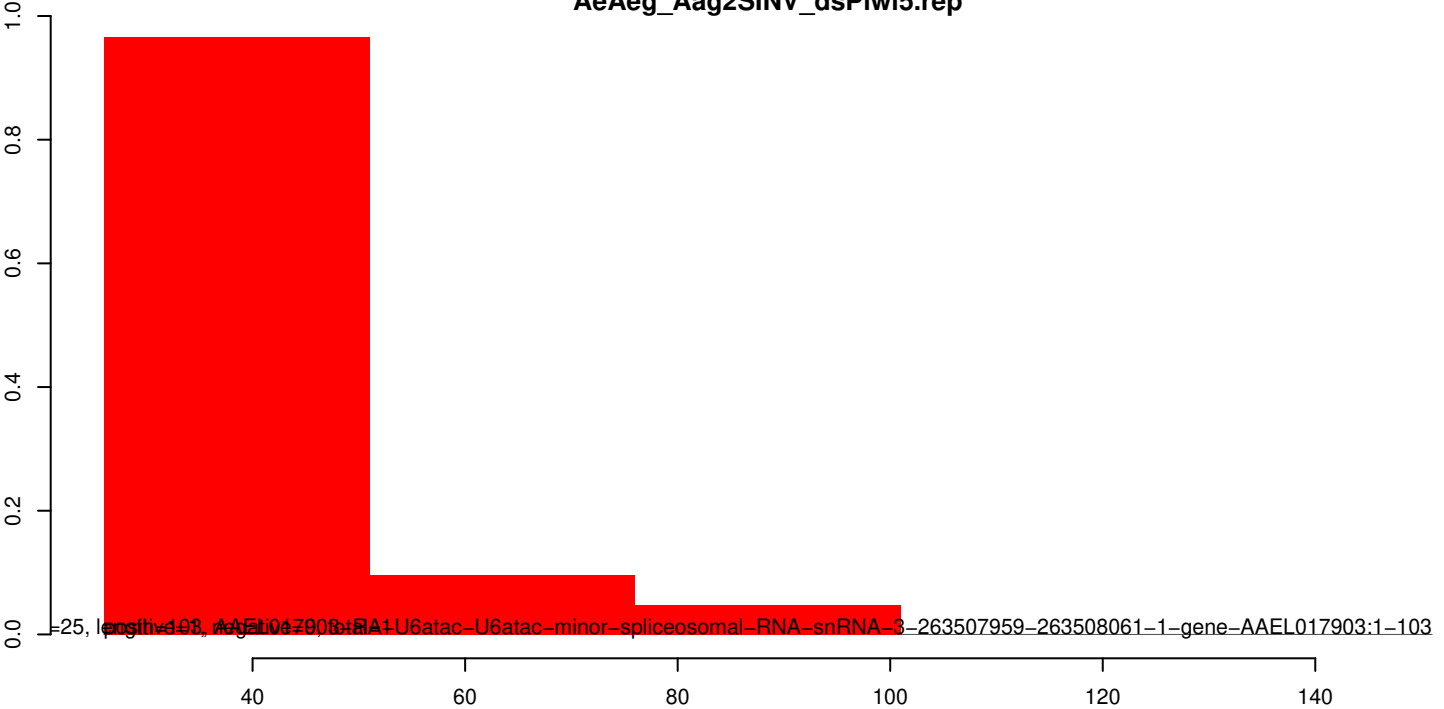
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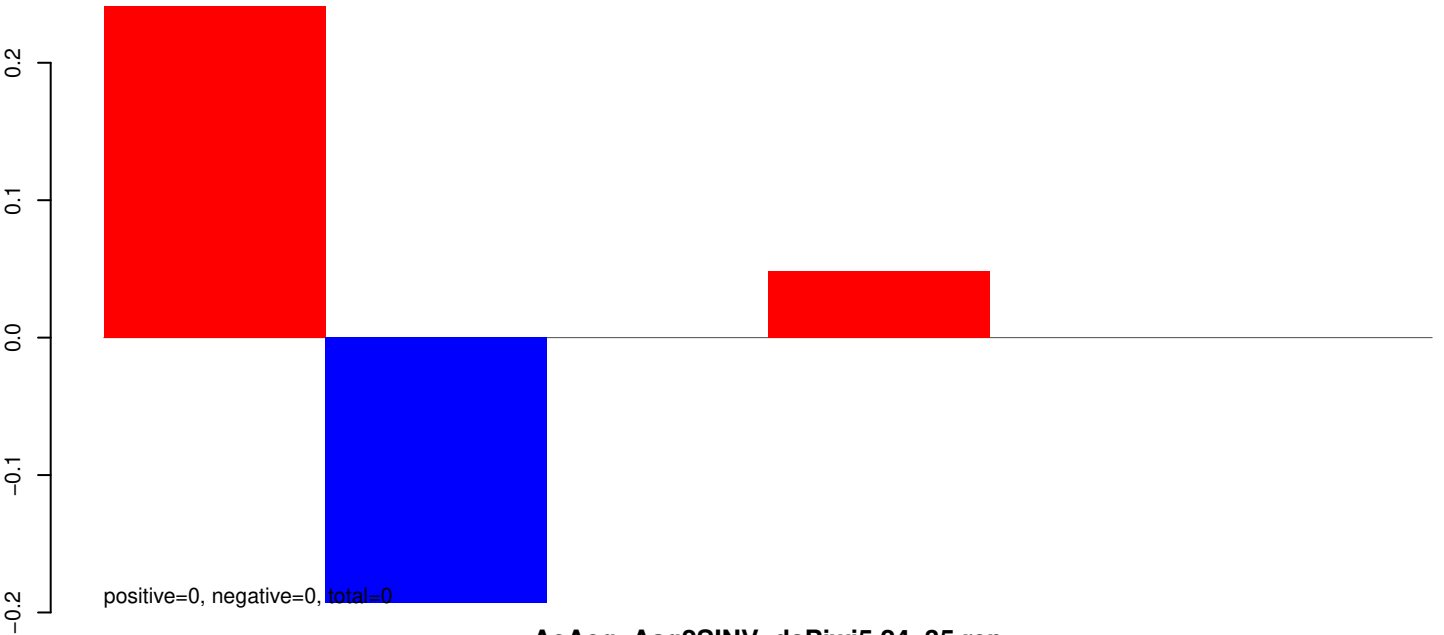
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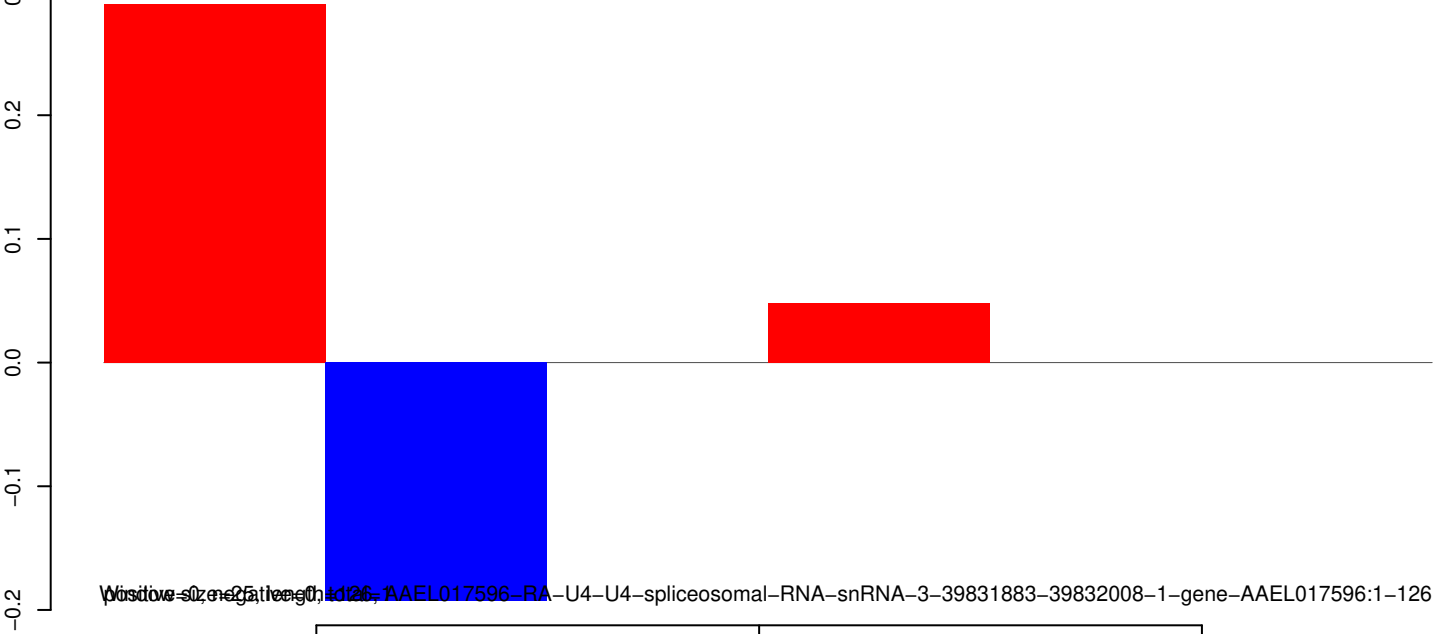
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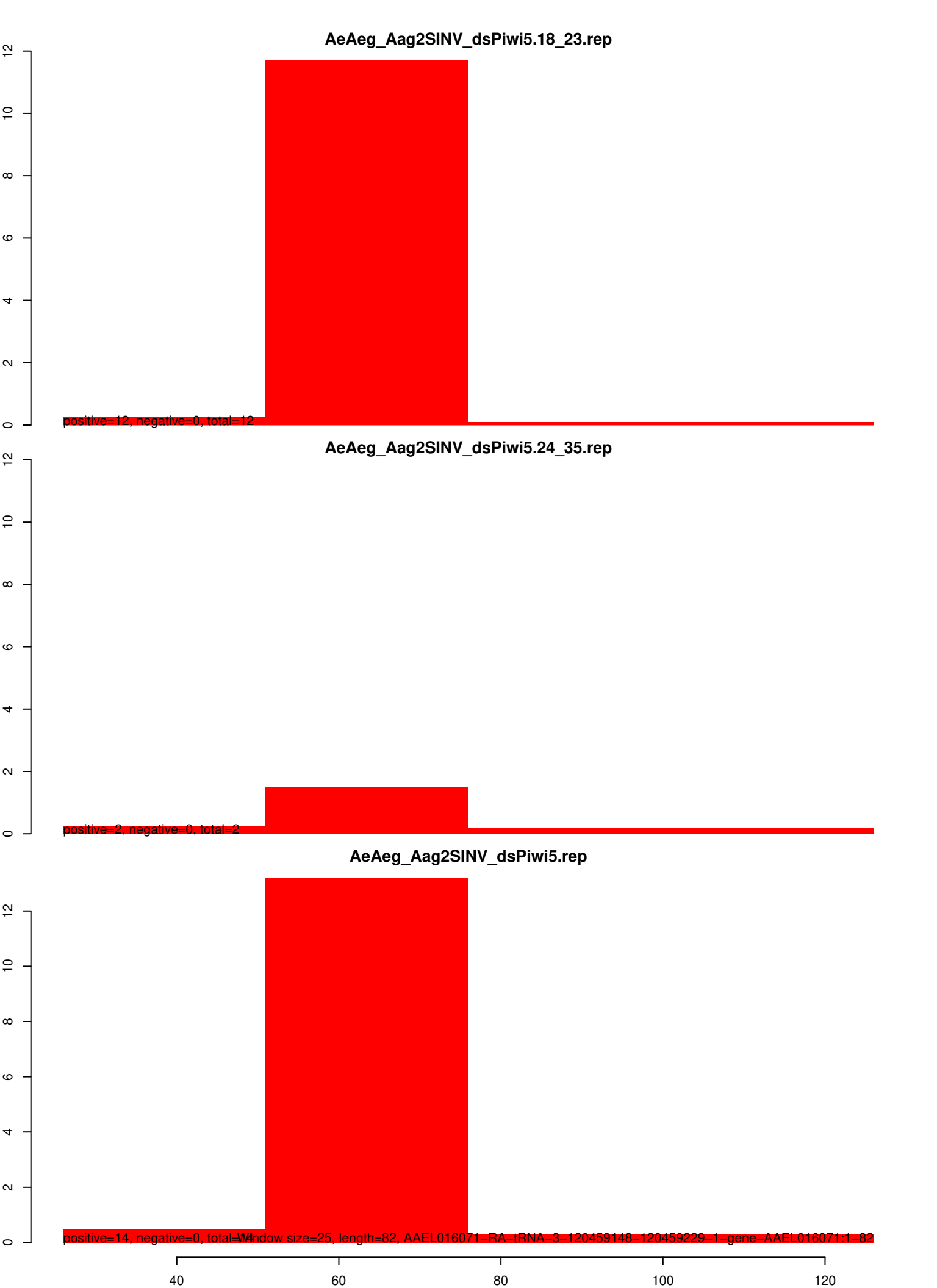


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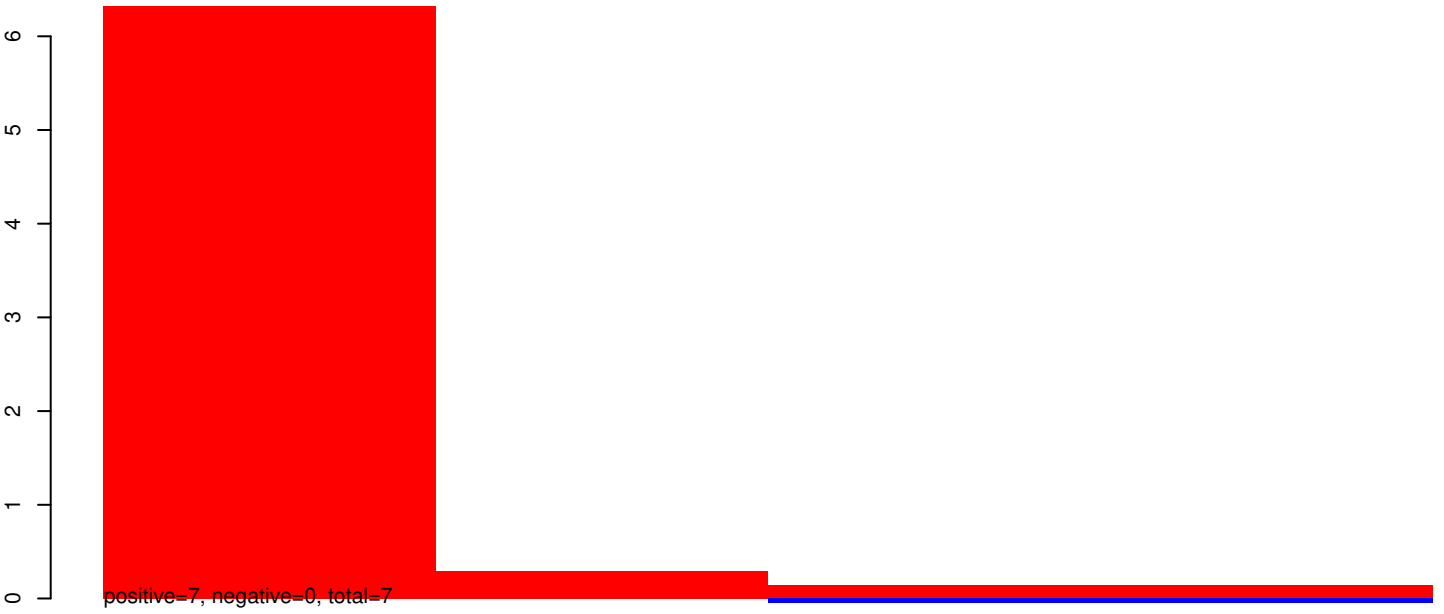


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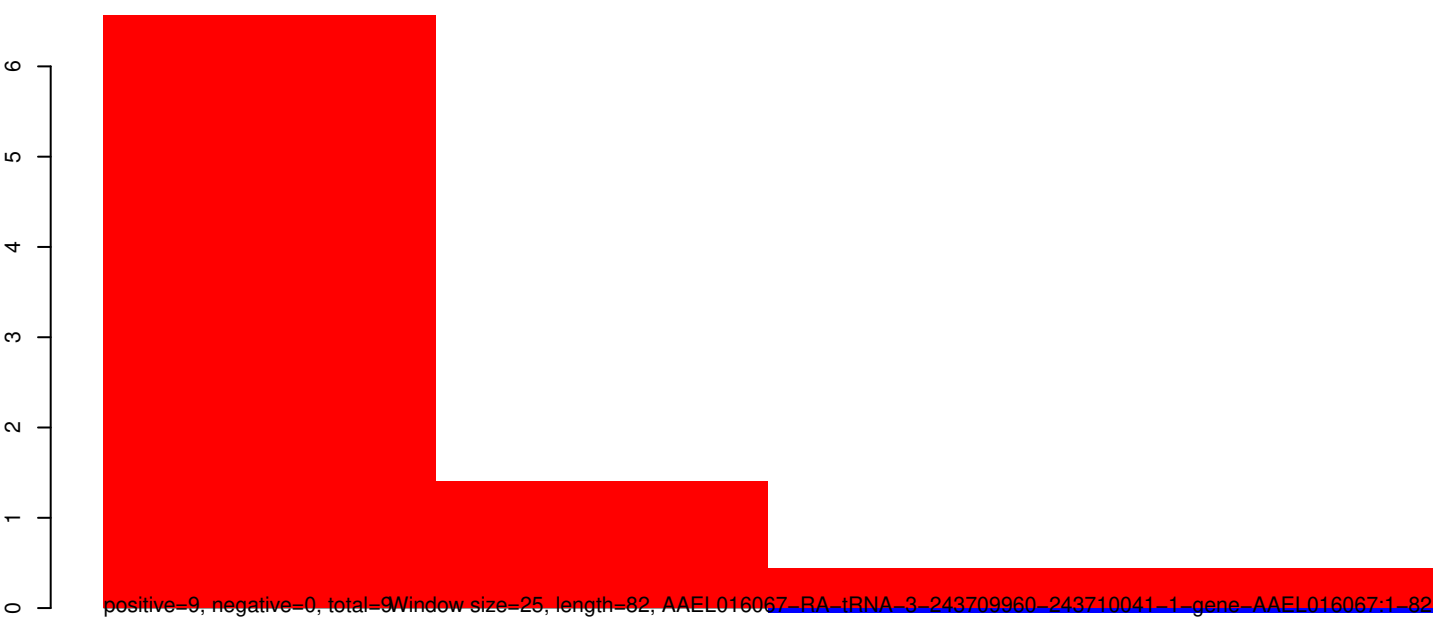
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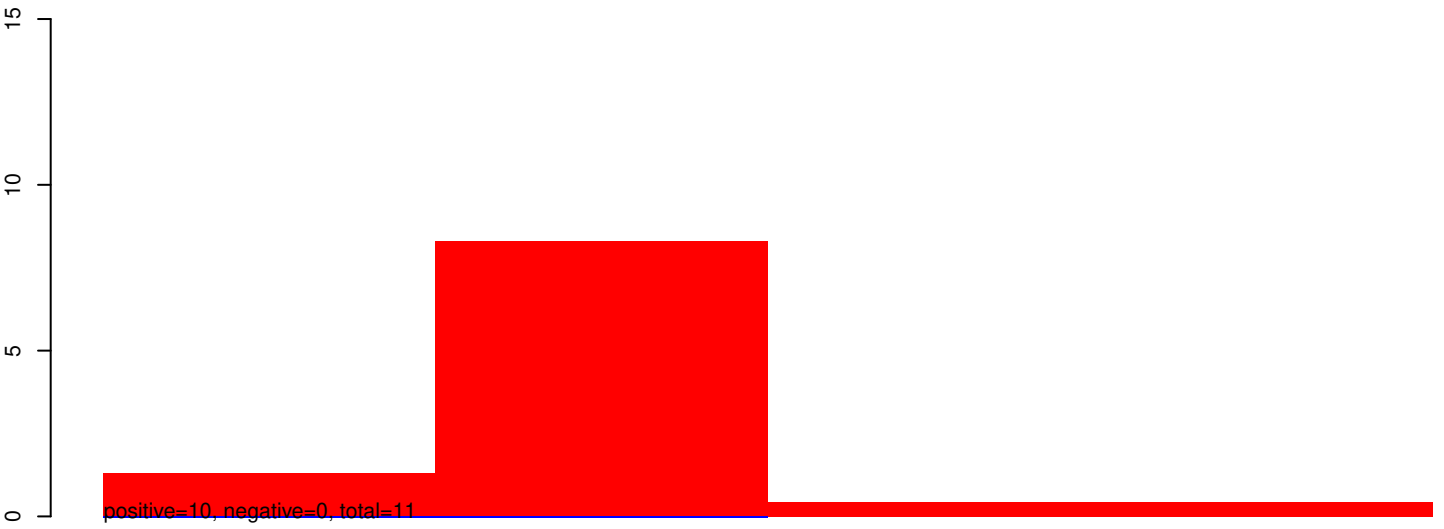
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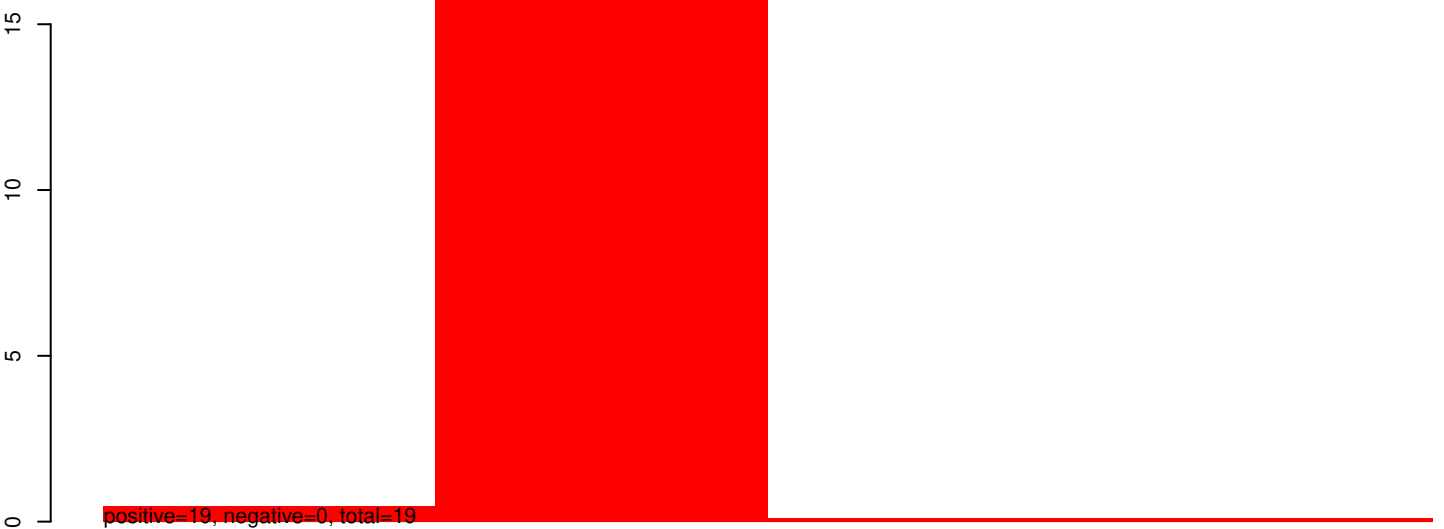
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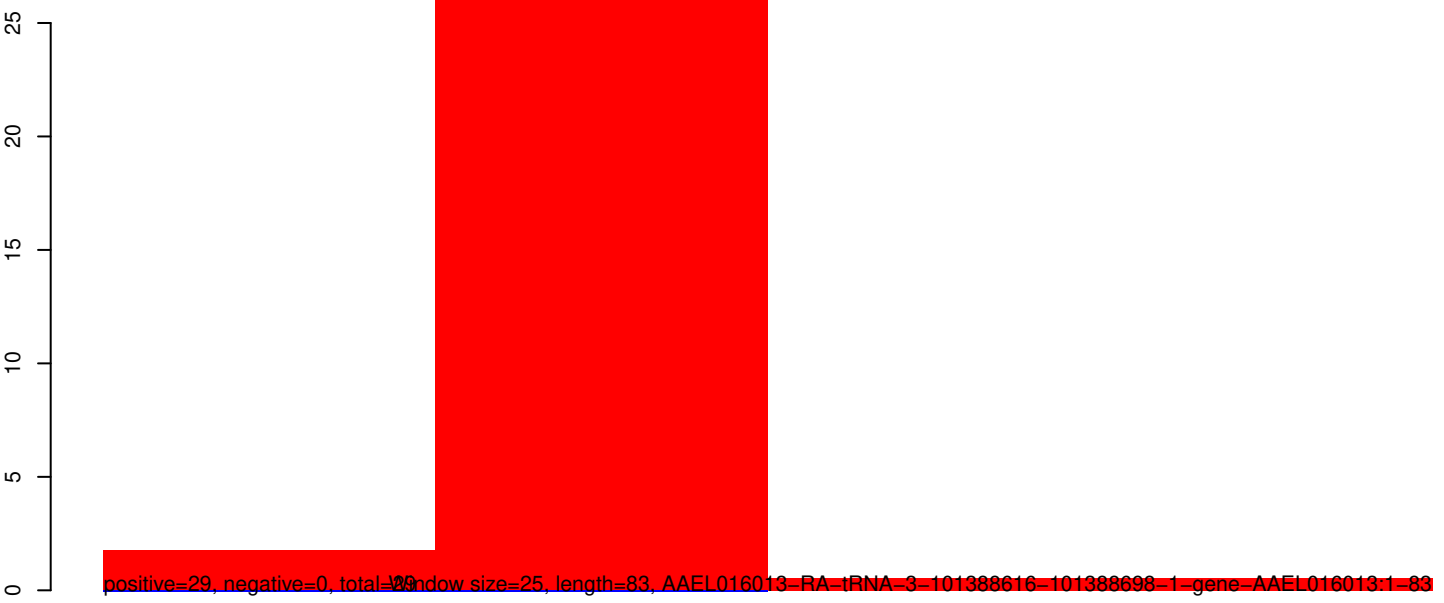
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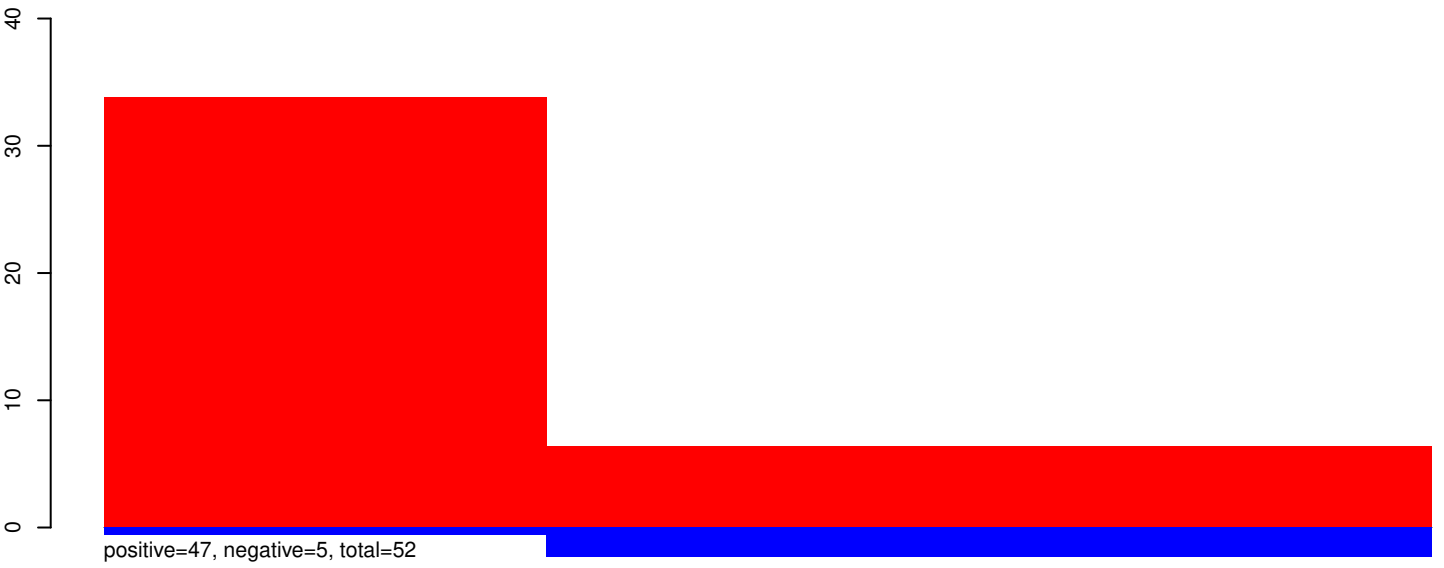


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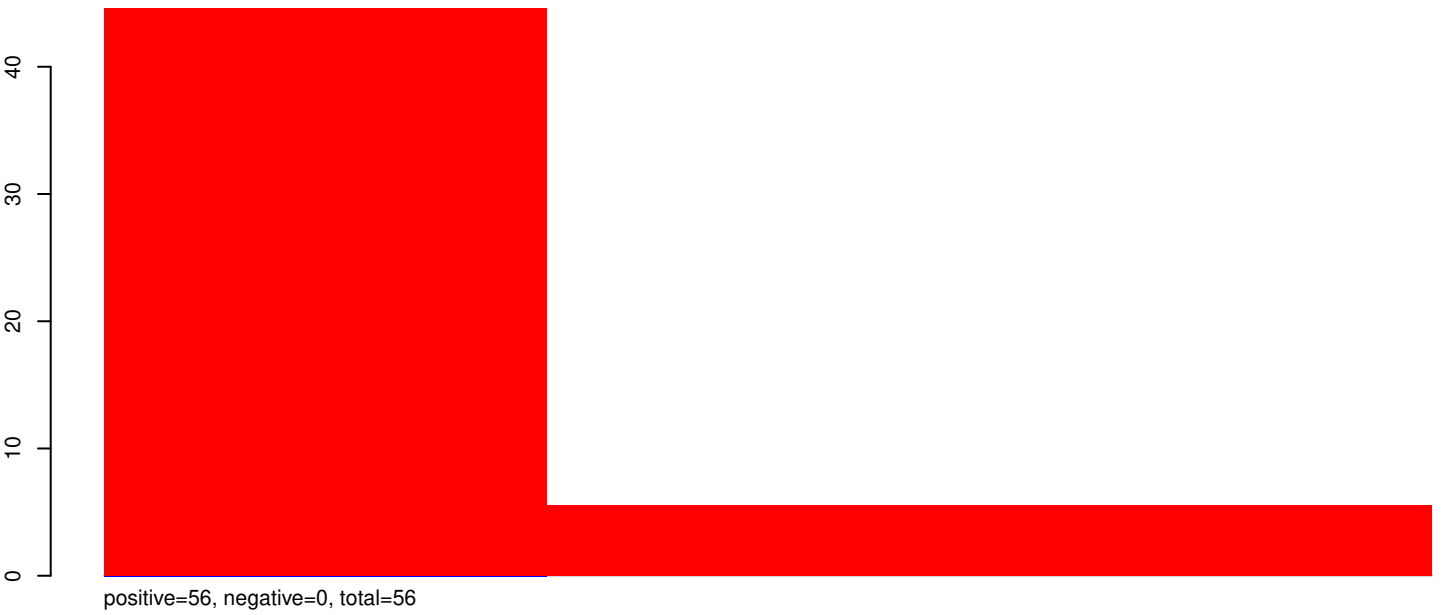


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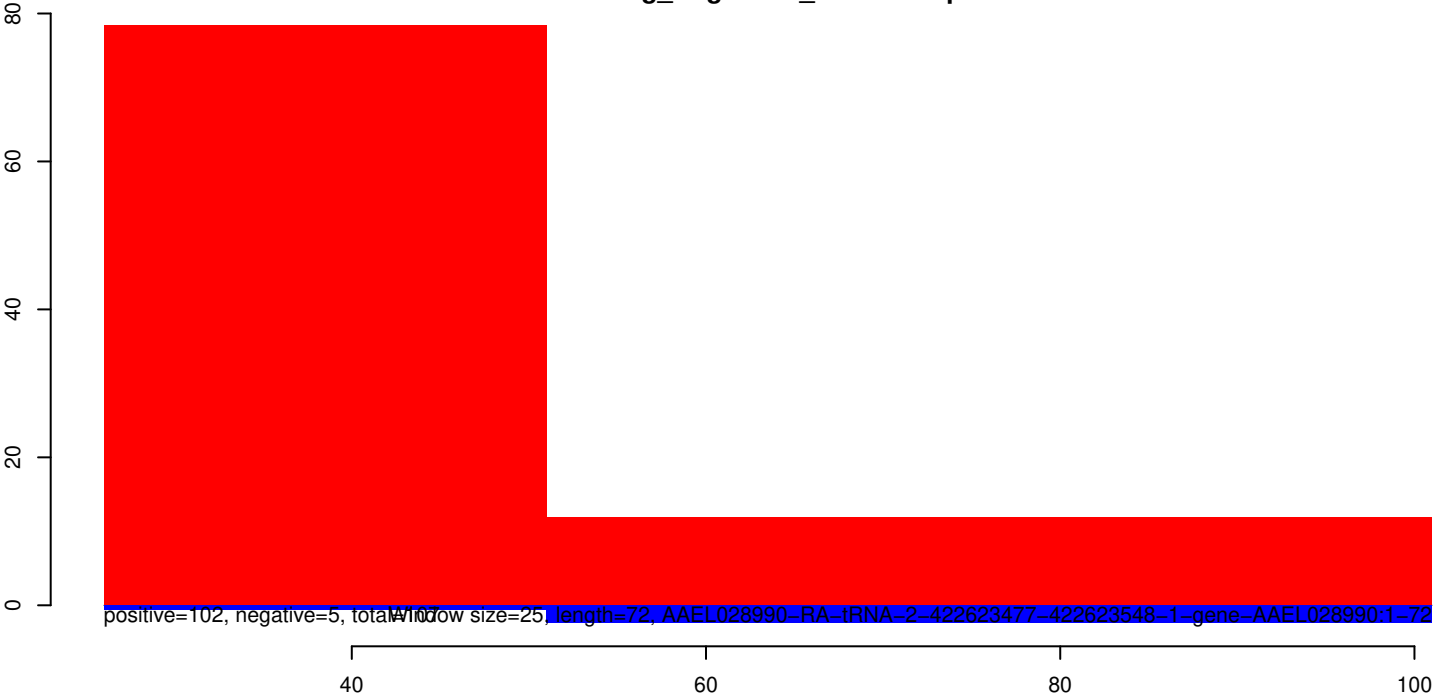
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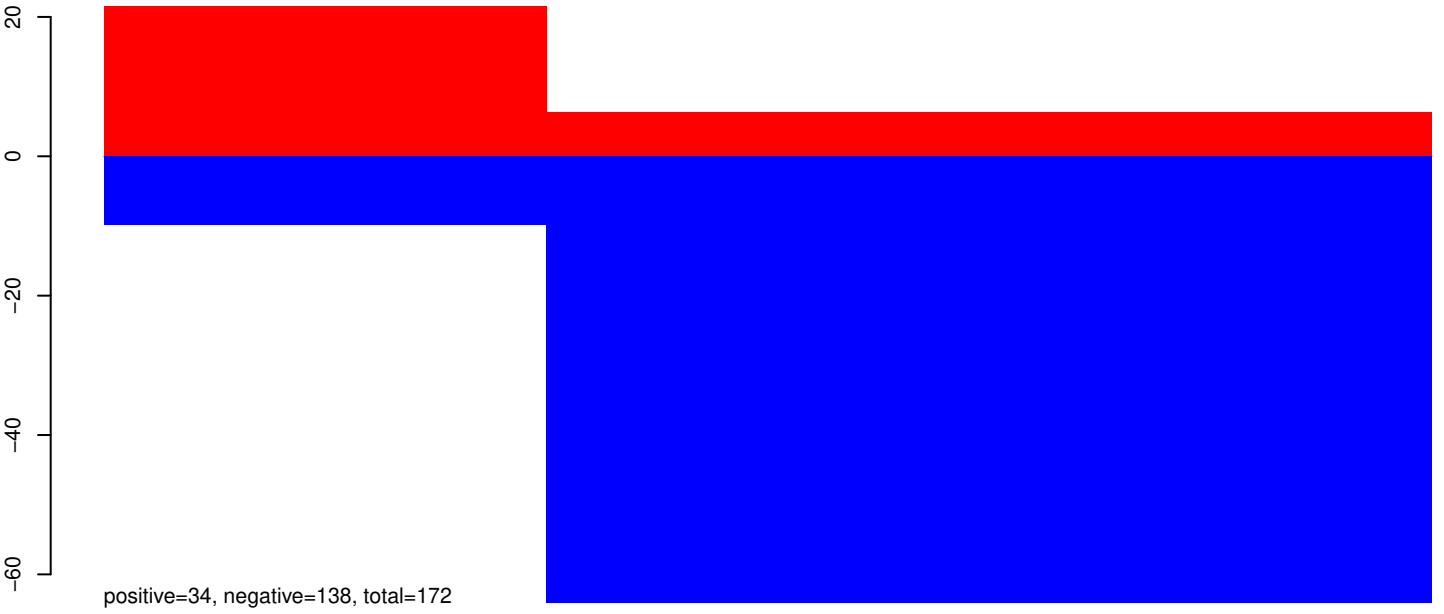
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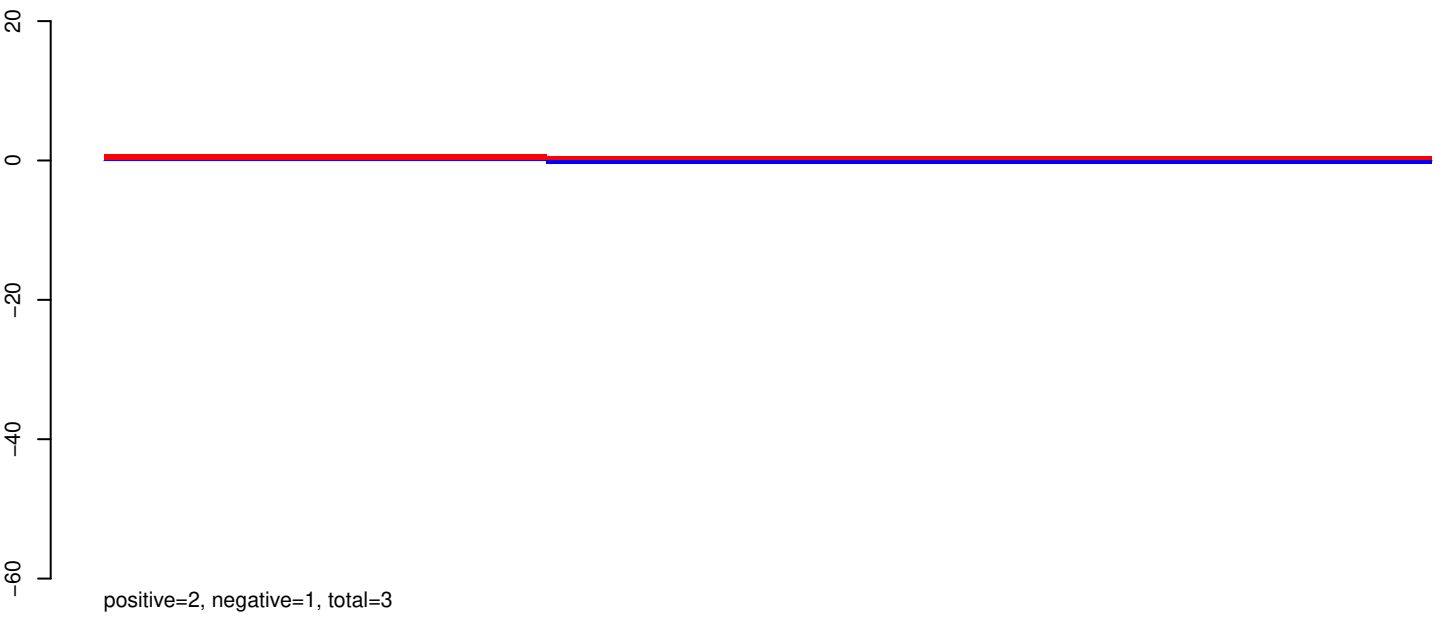
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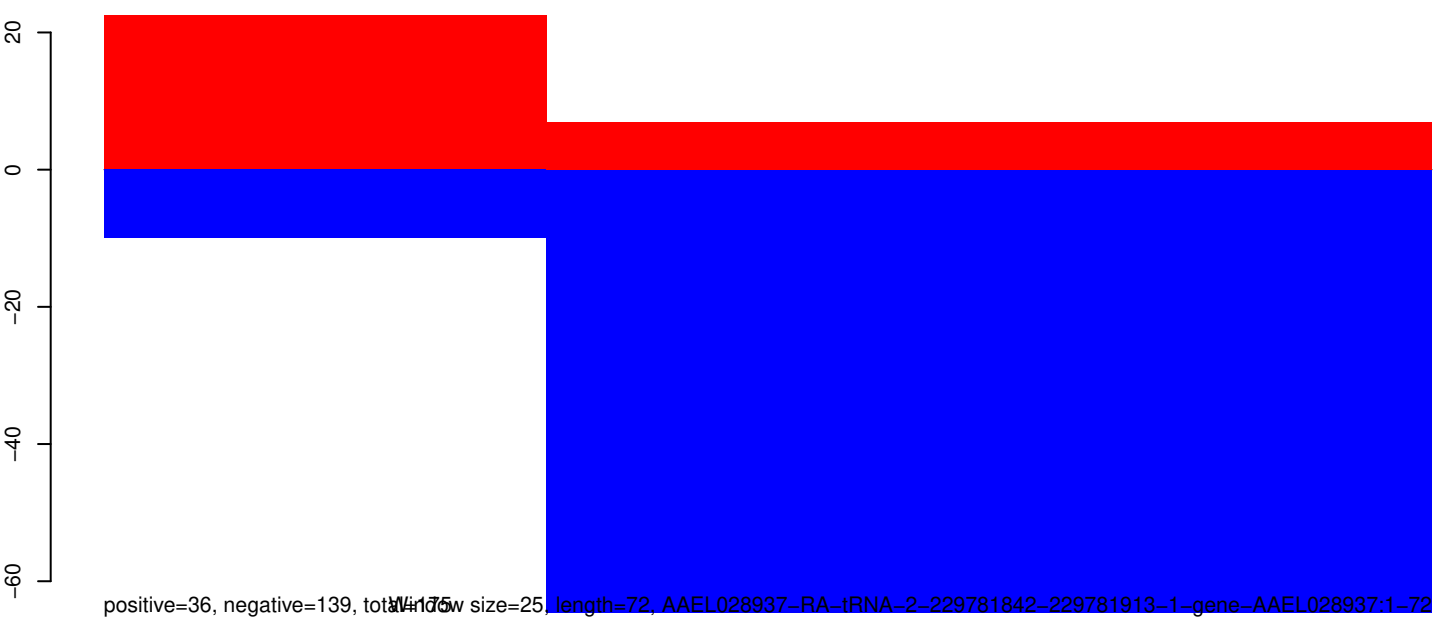
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AeAeg_Aag2SINV_dsPiwi5.rep



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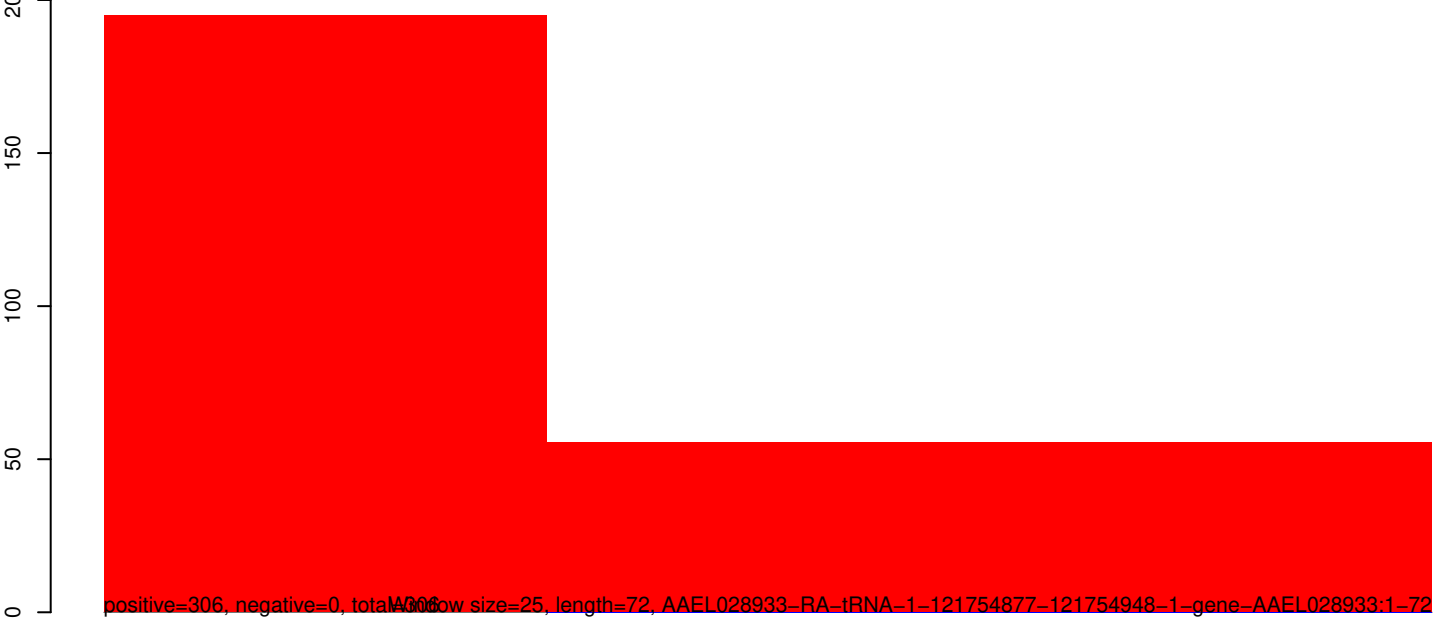
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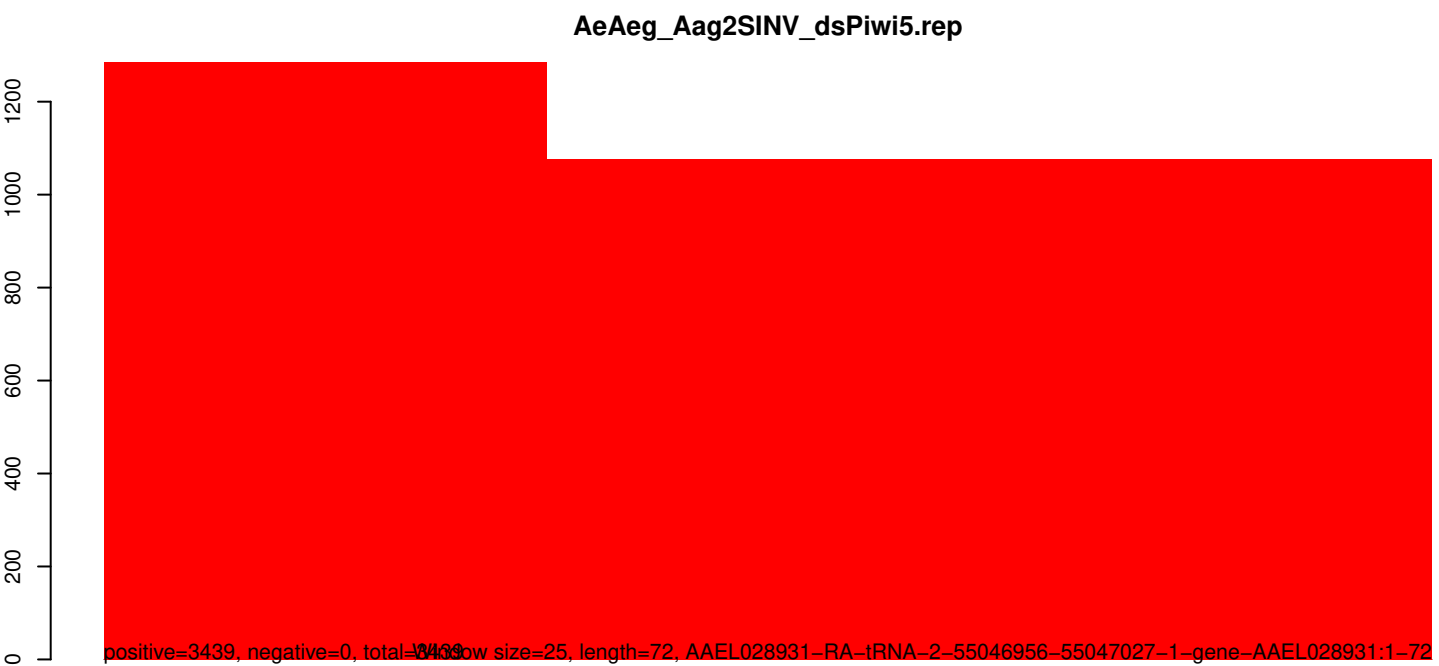
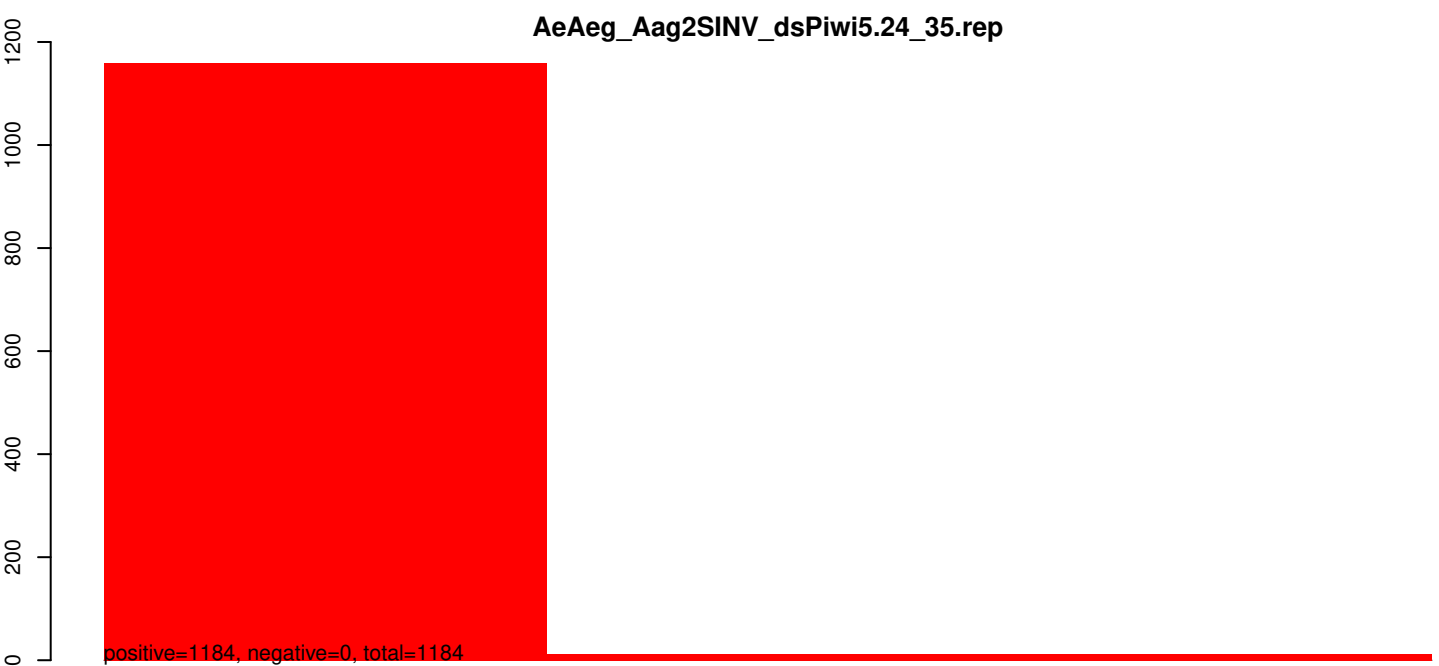
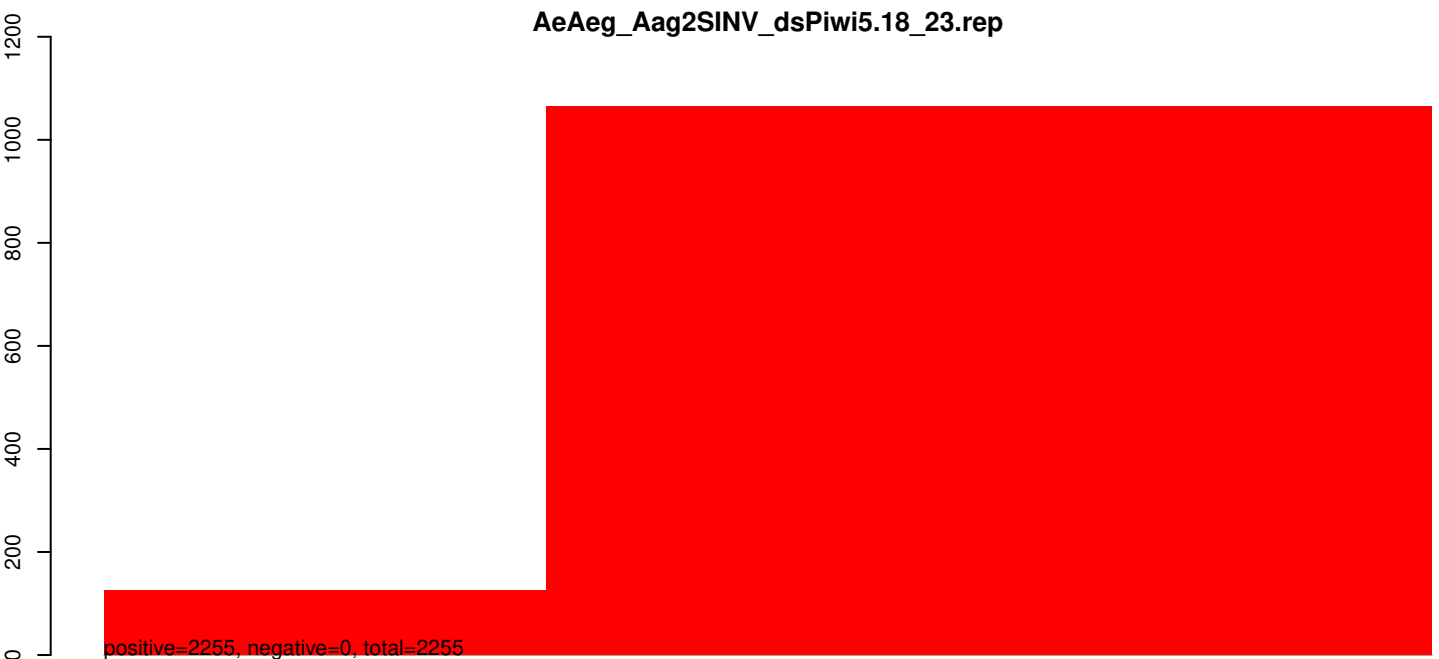


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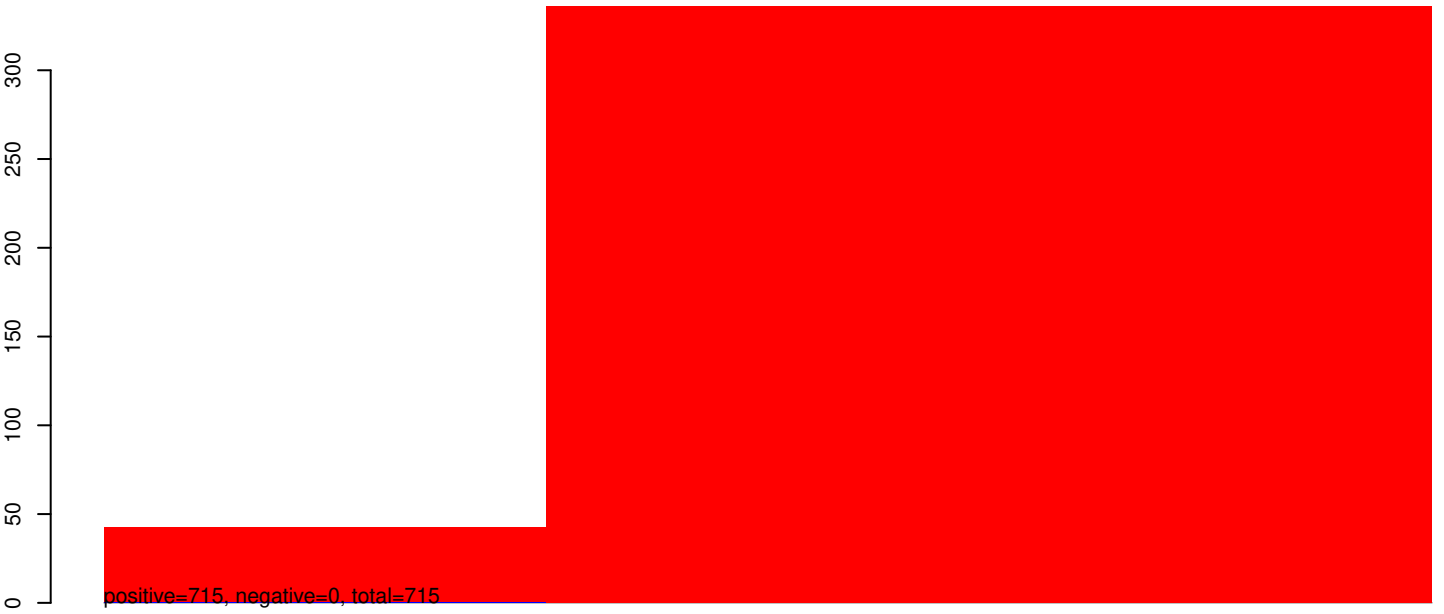
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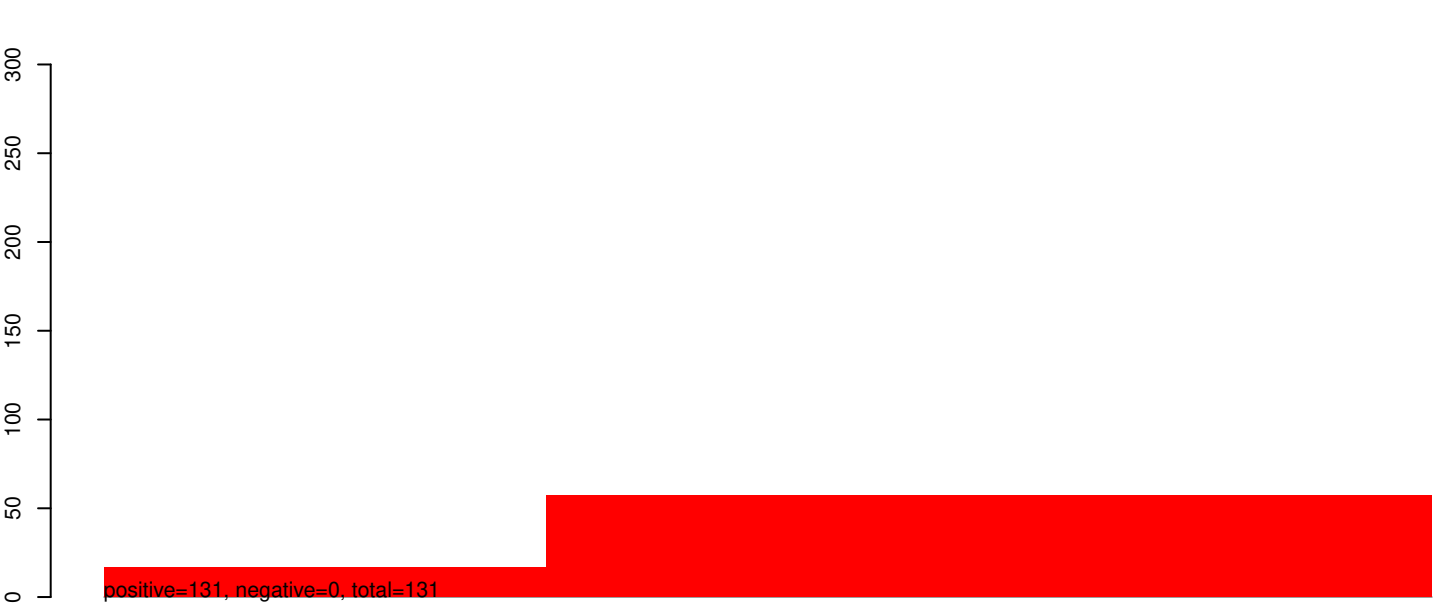


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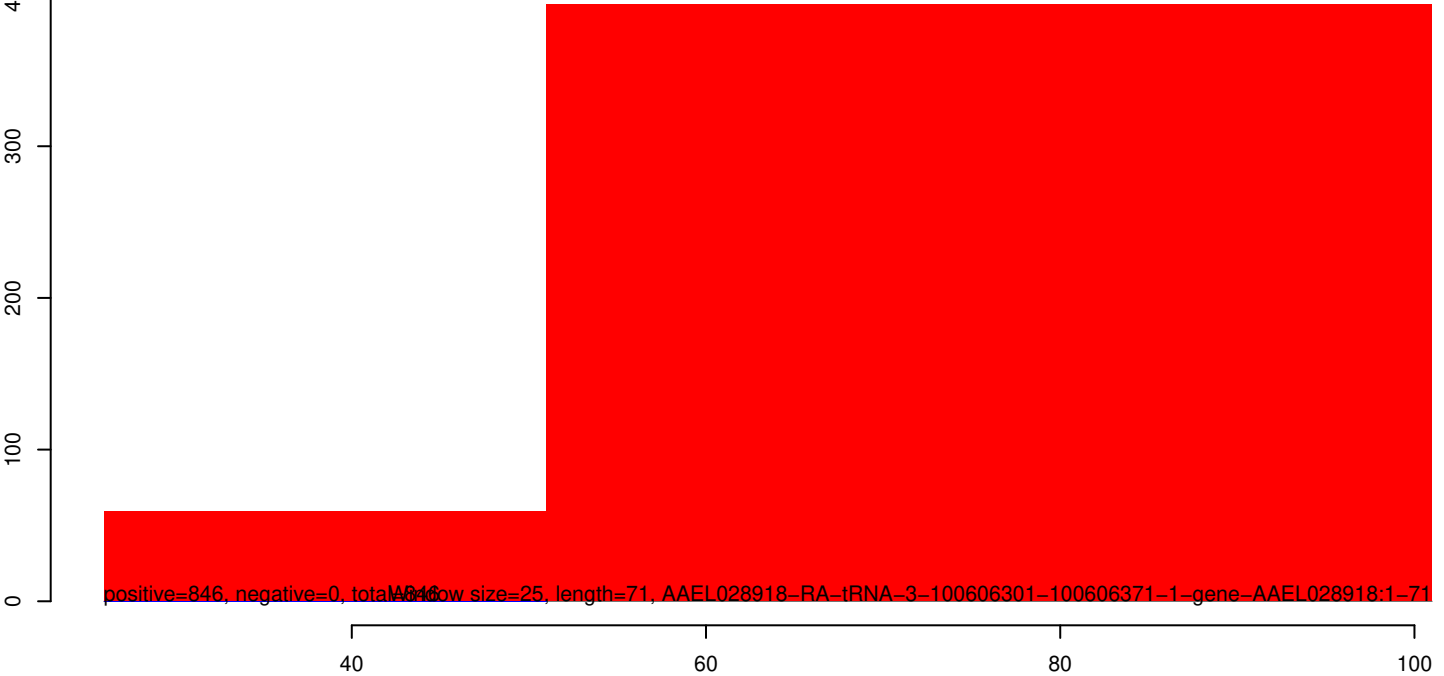
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AeAeg_Aag2SINV_dsPiwi5.24_35.rep



AeAeg_Aag2SINV_dsPiwi5.rep



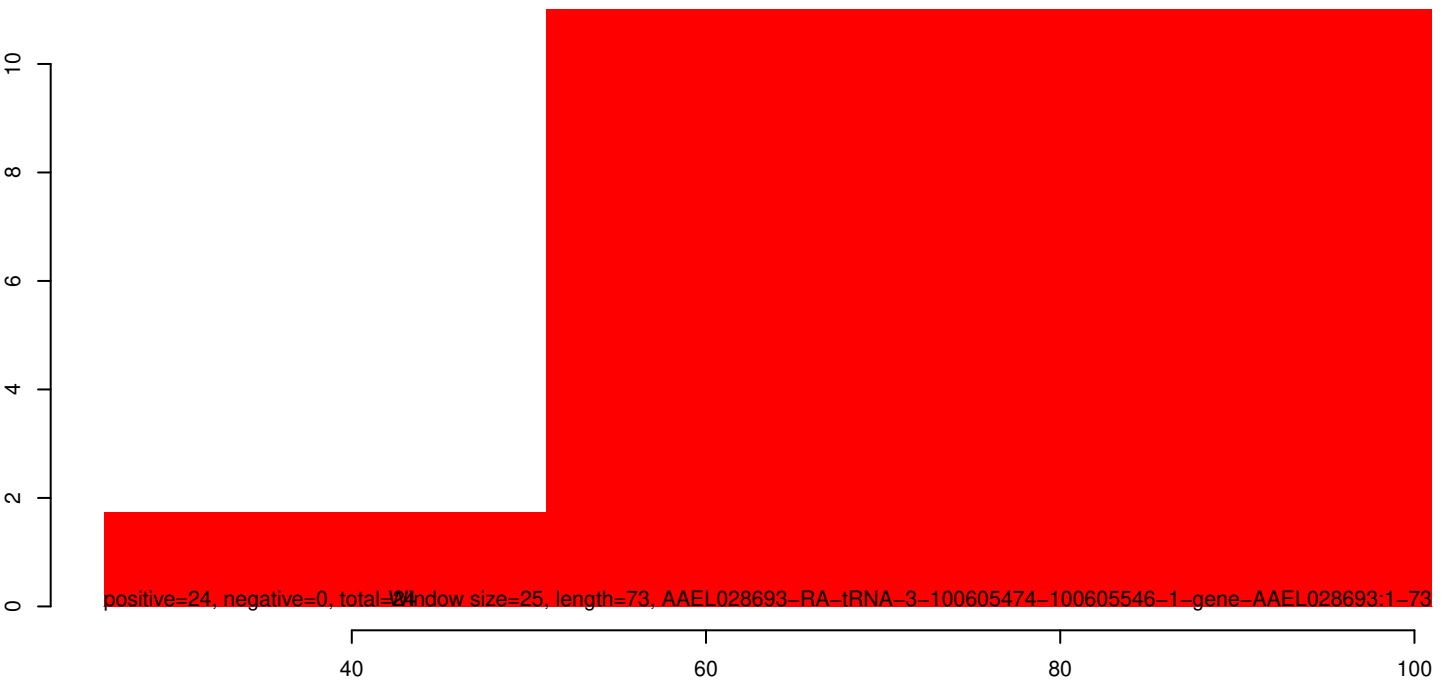
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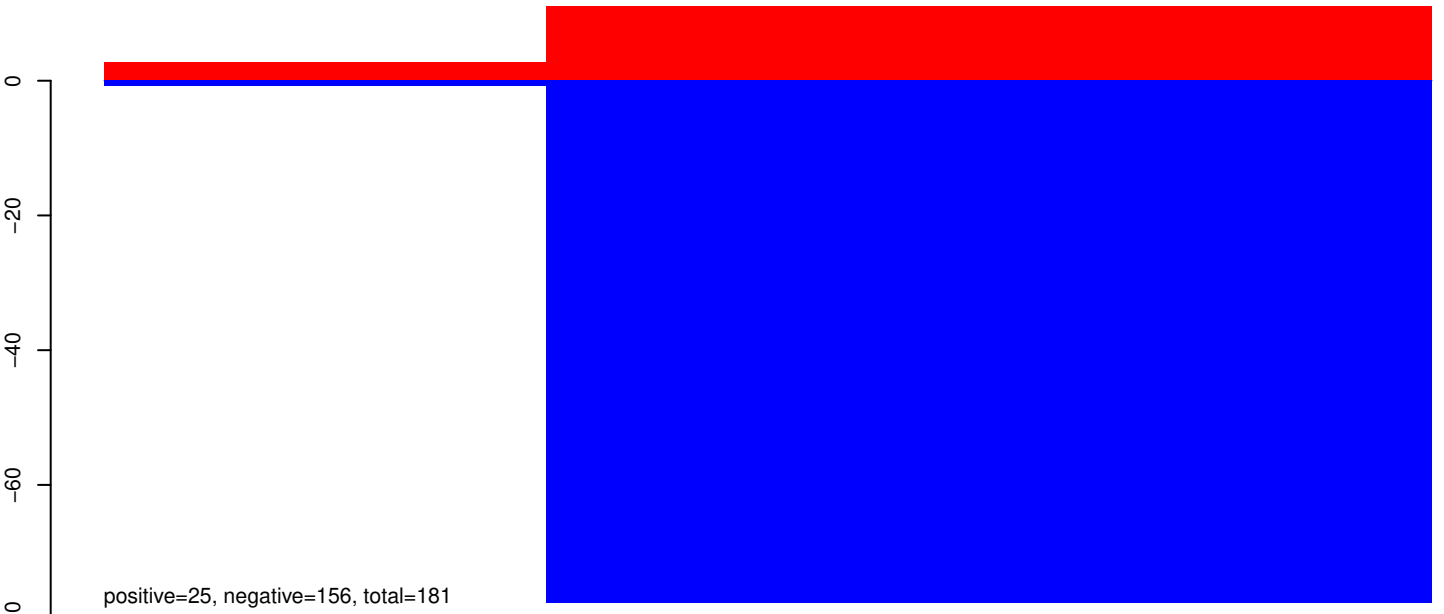
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AeAeg_Aag2SINV_dsPiwi5.rep



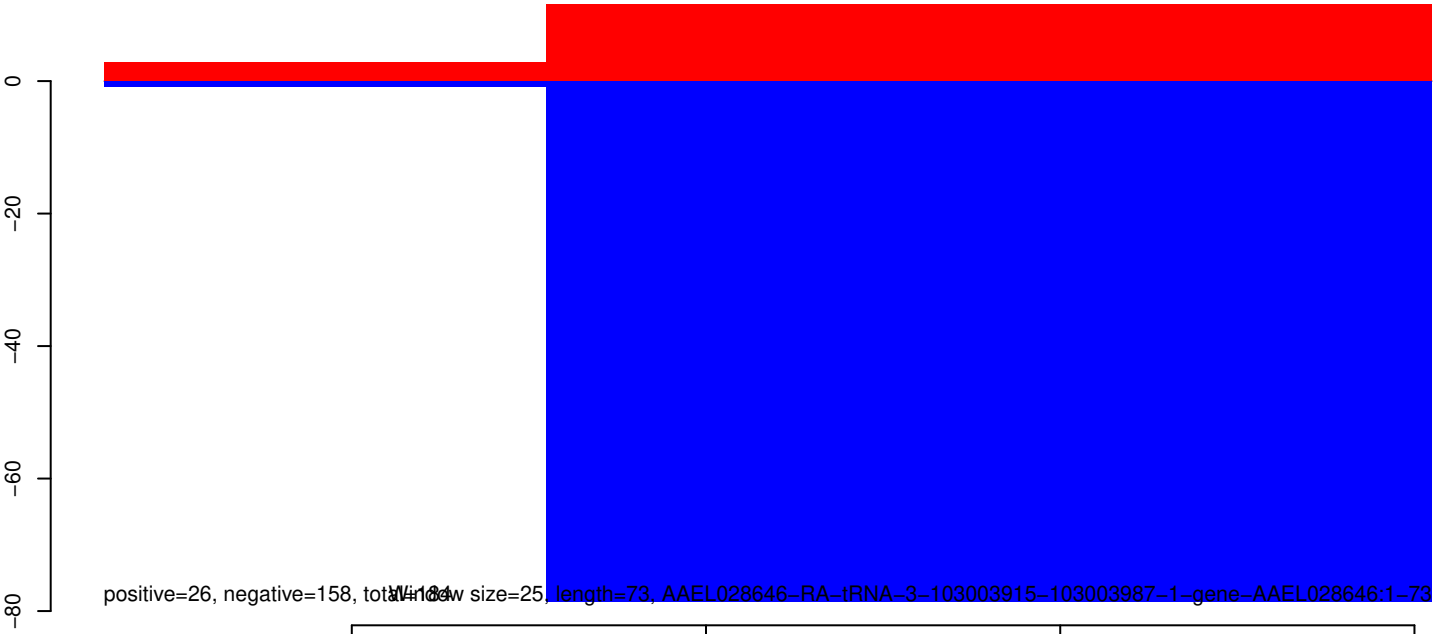
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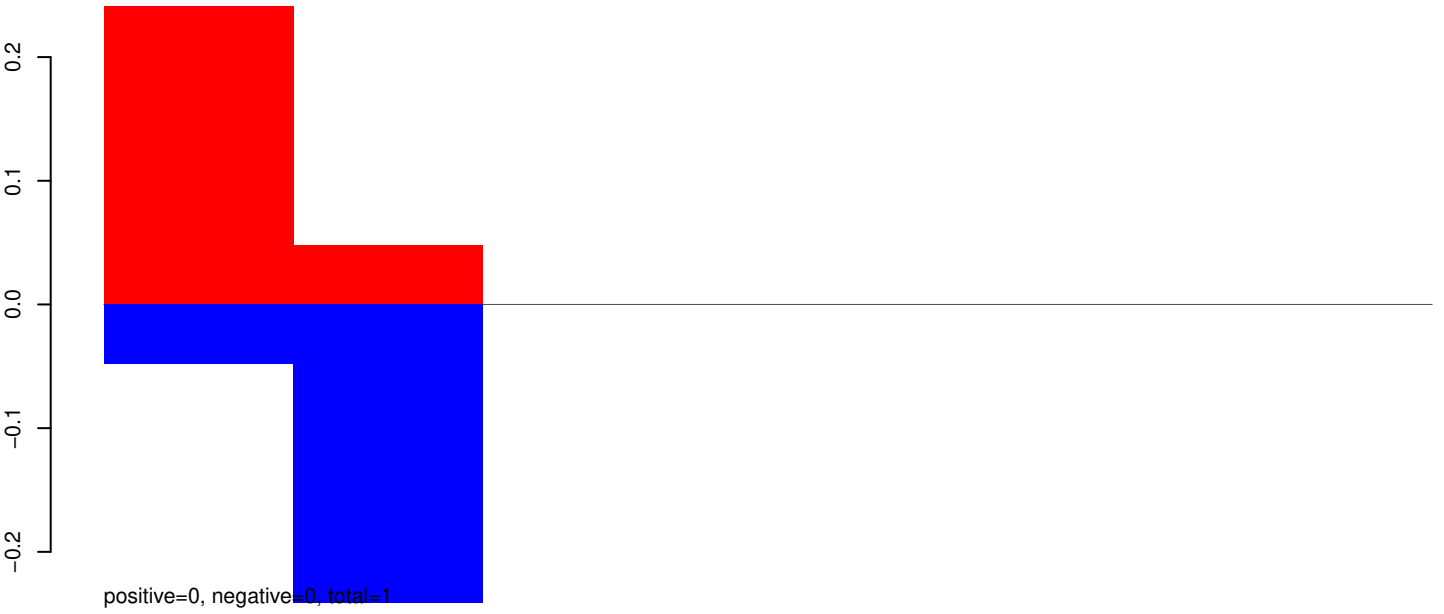
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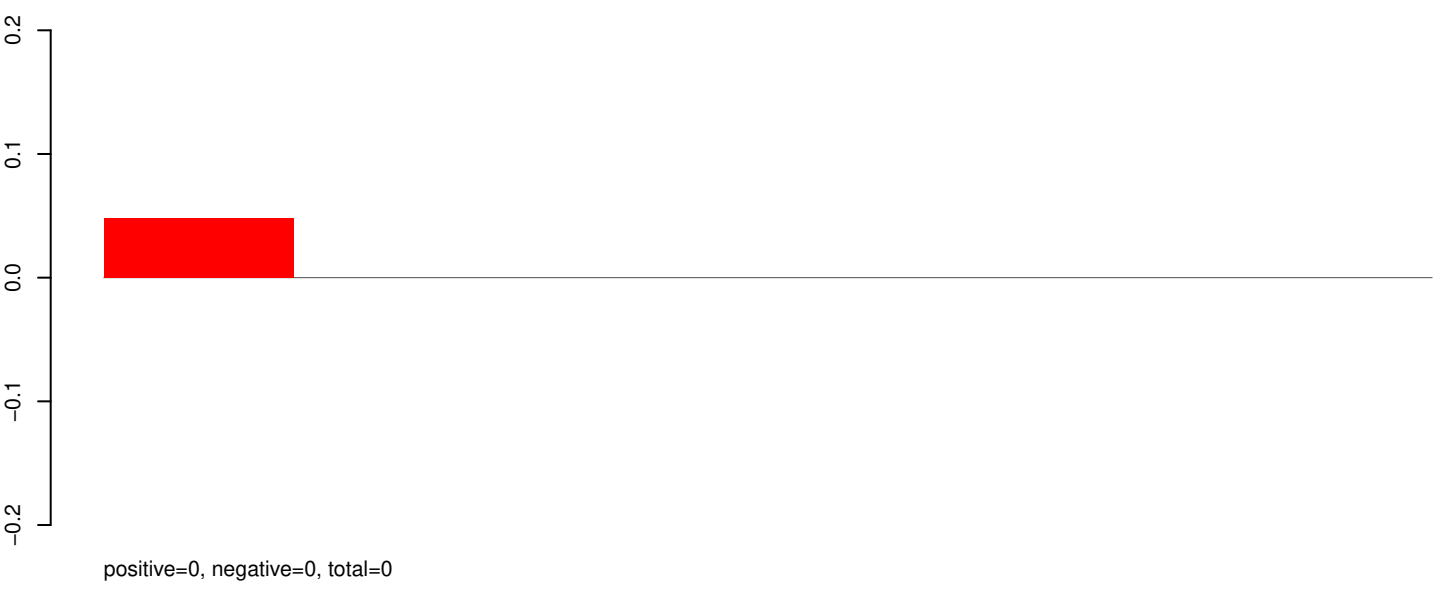
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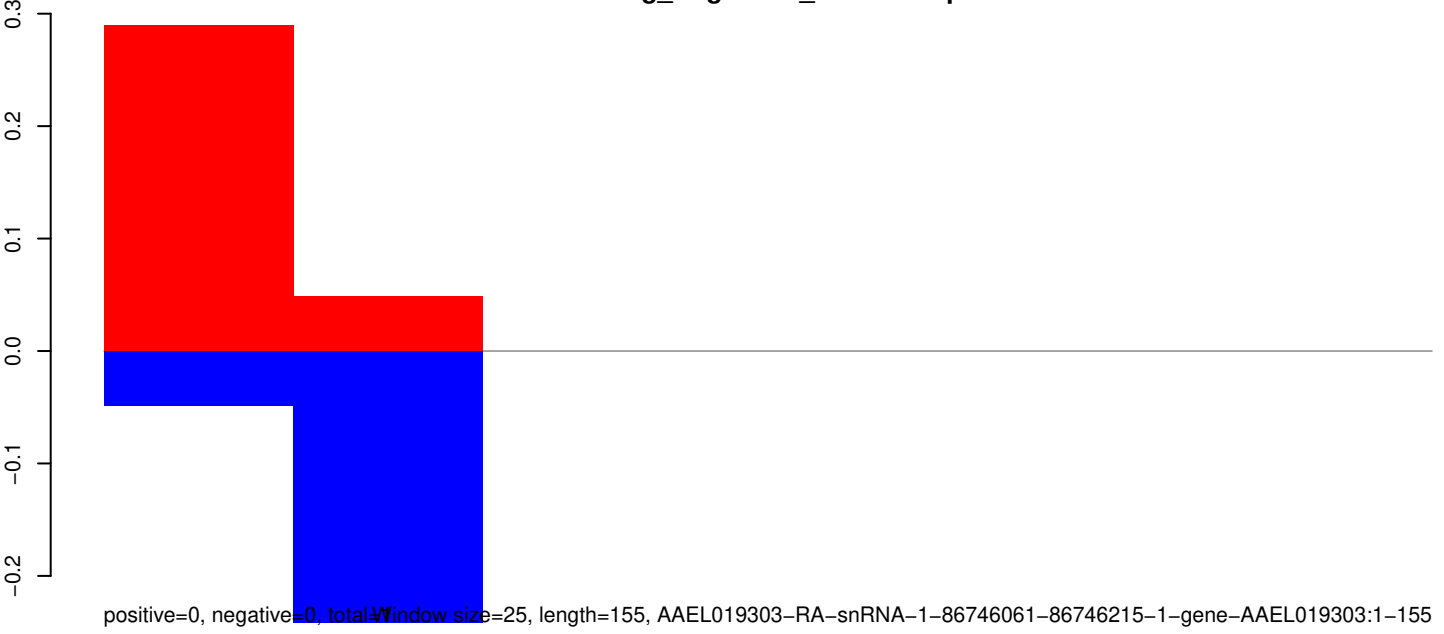
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AeAeg_Aag2SINV_dsPiwi5.rep



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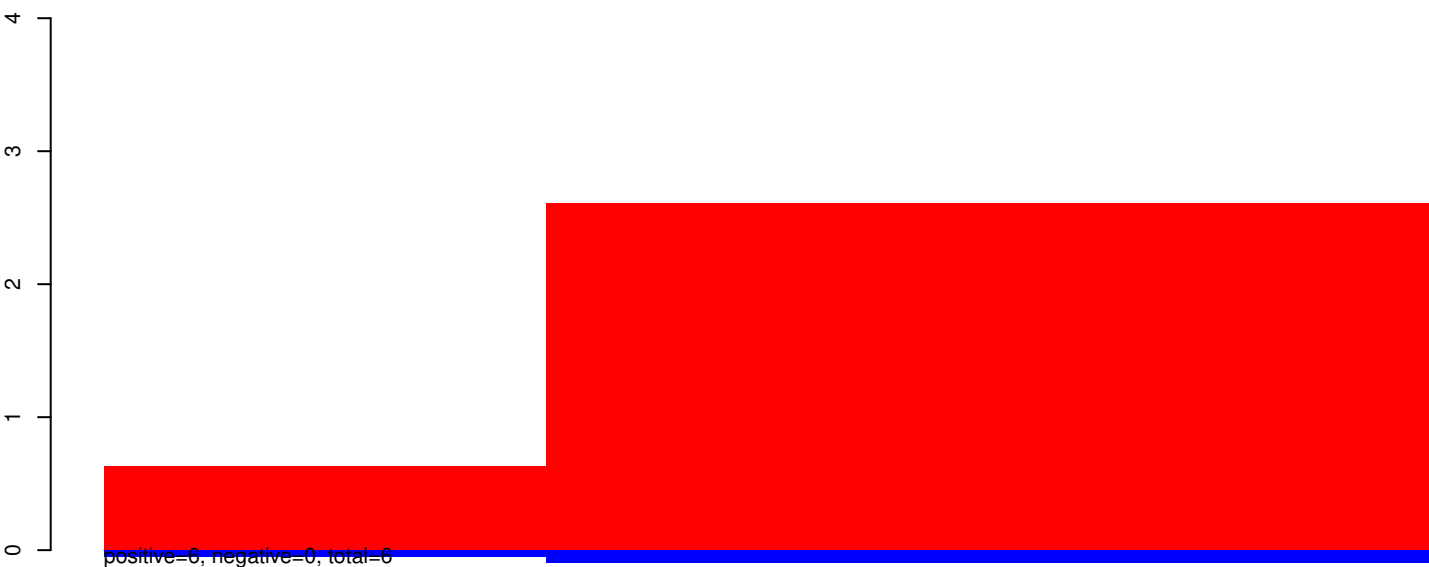
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150

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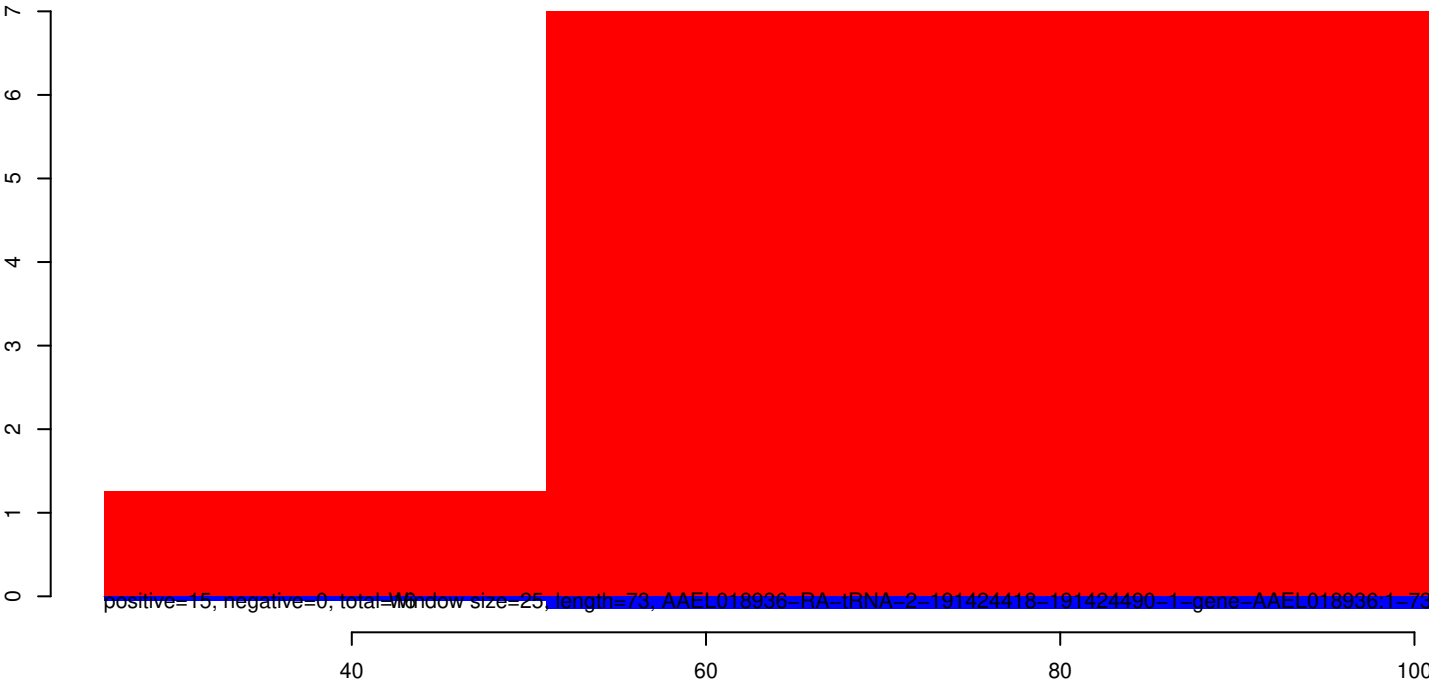
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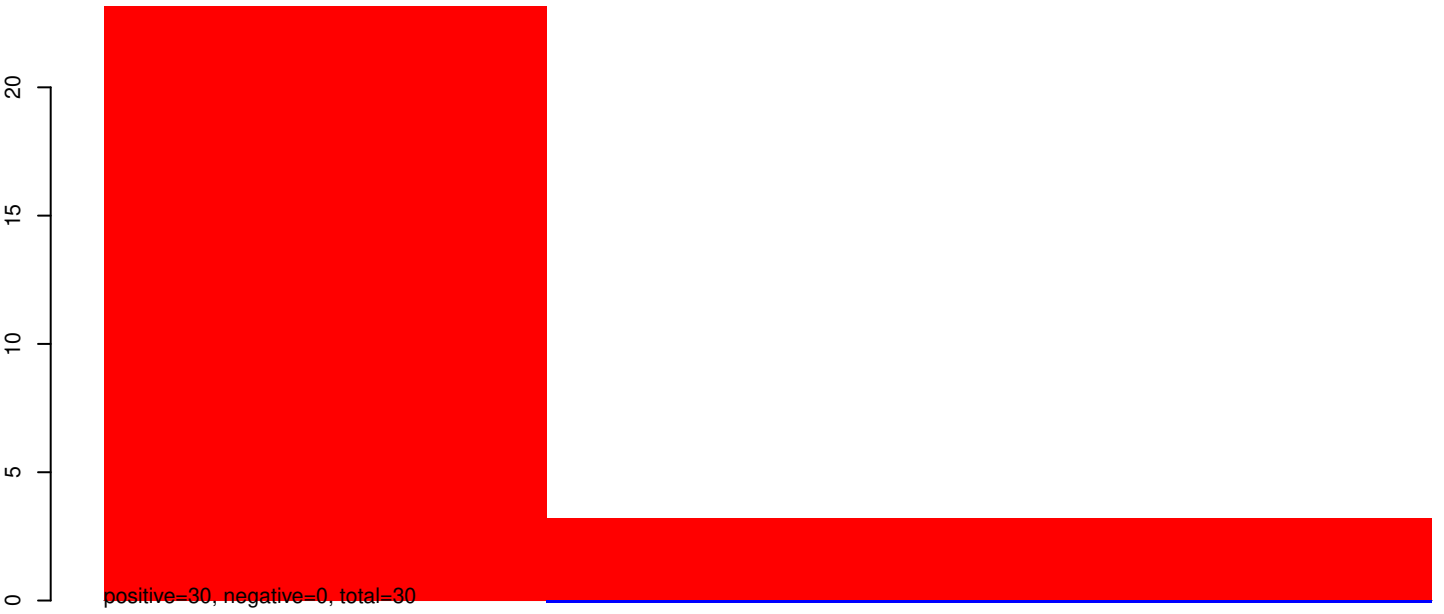
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AeAeg_Aag2SINV_dsPiwi5.rep



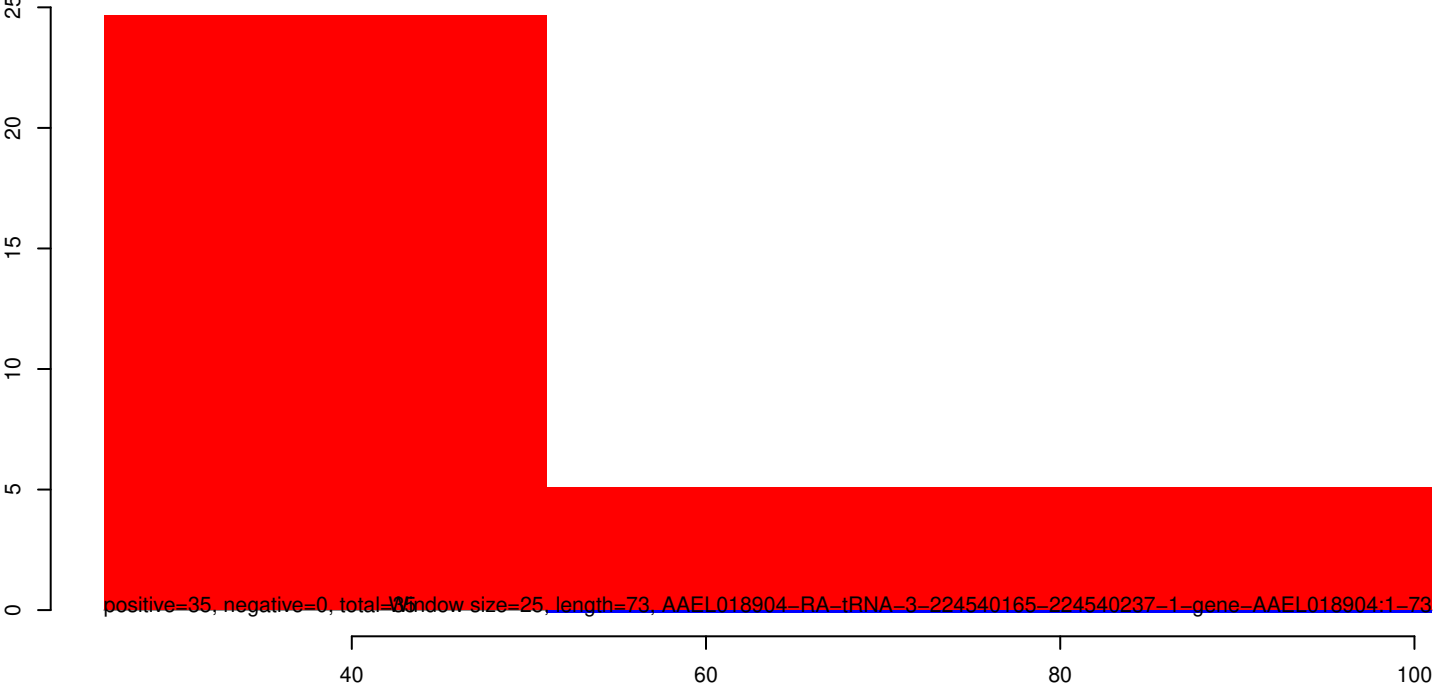
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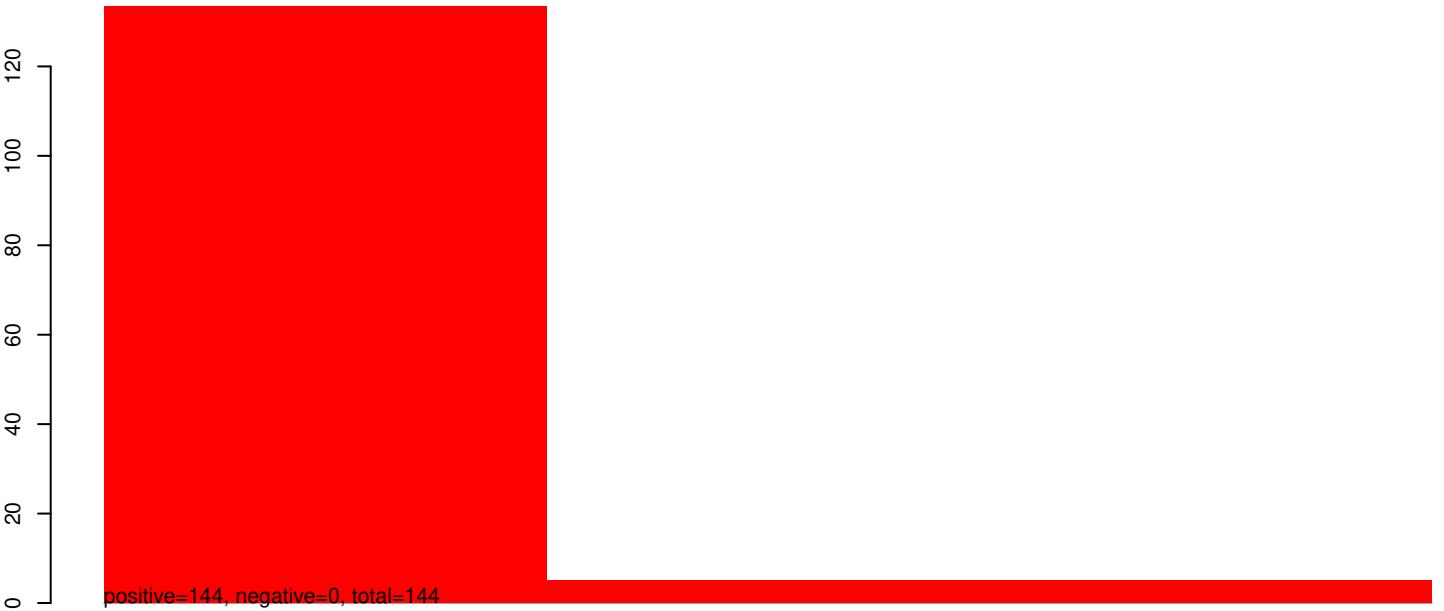


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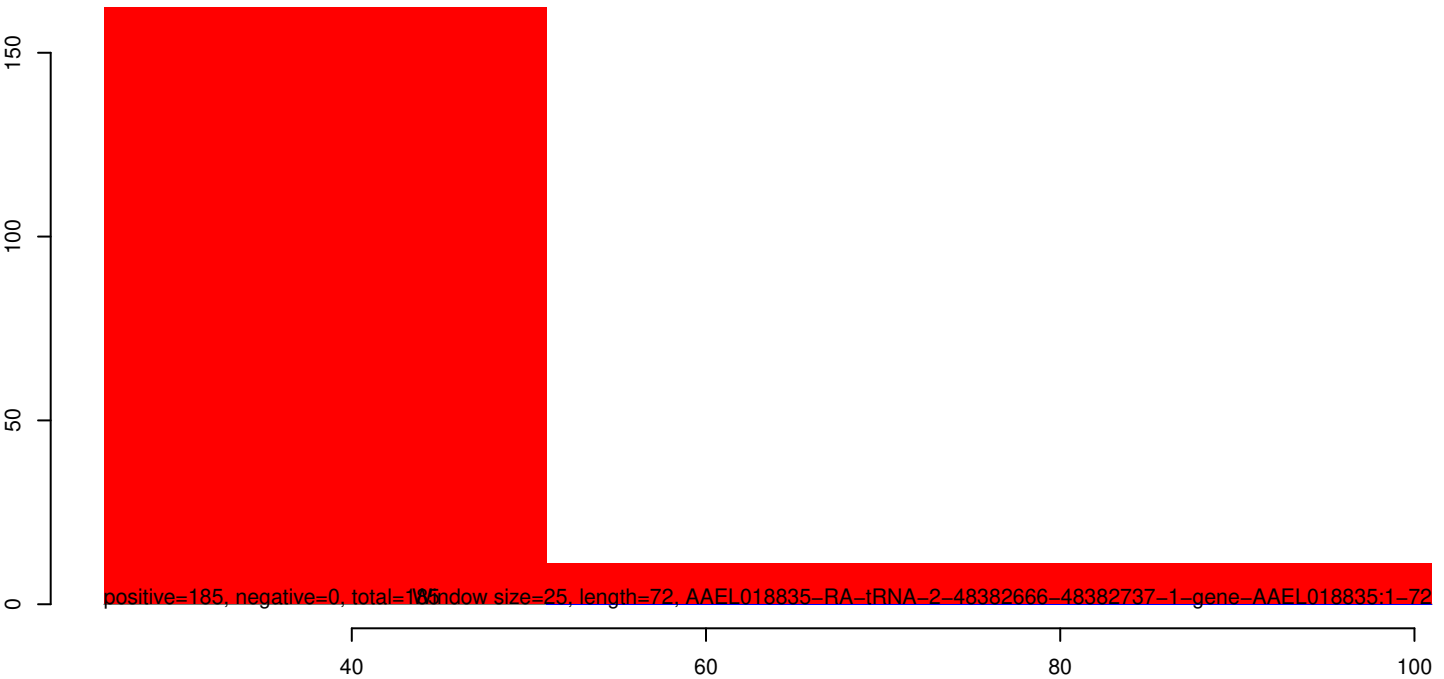
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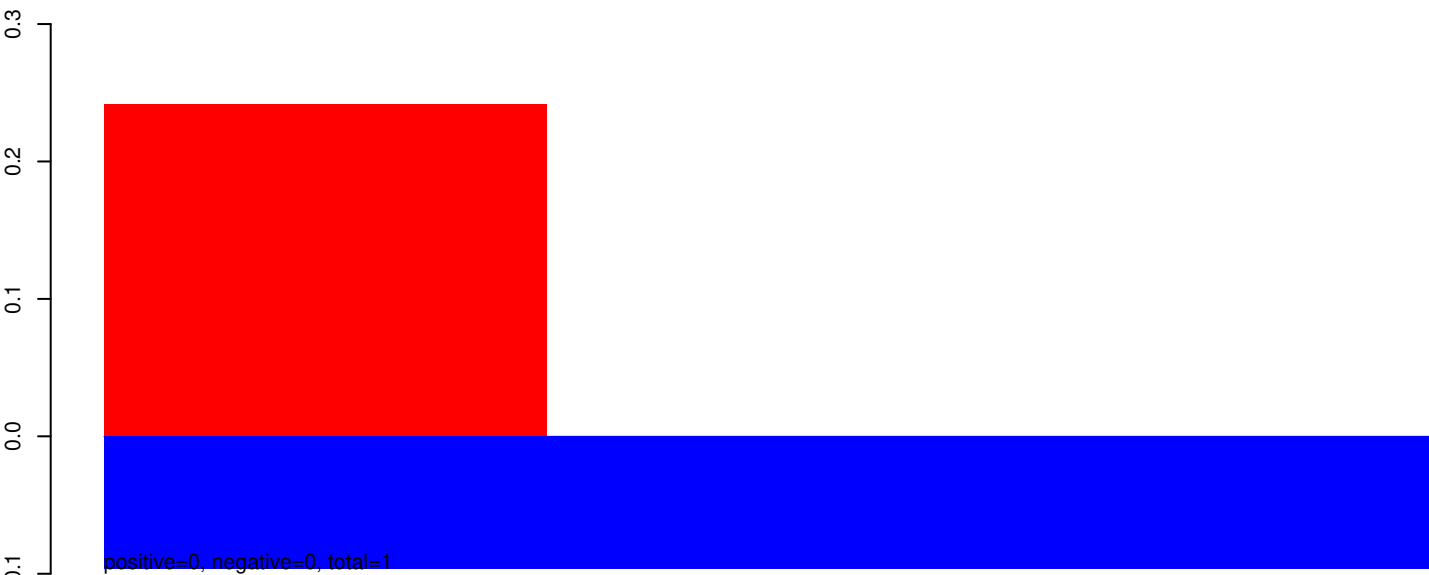
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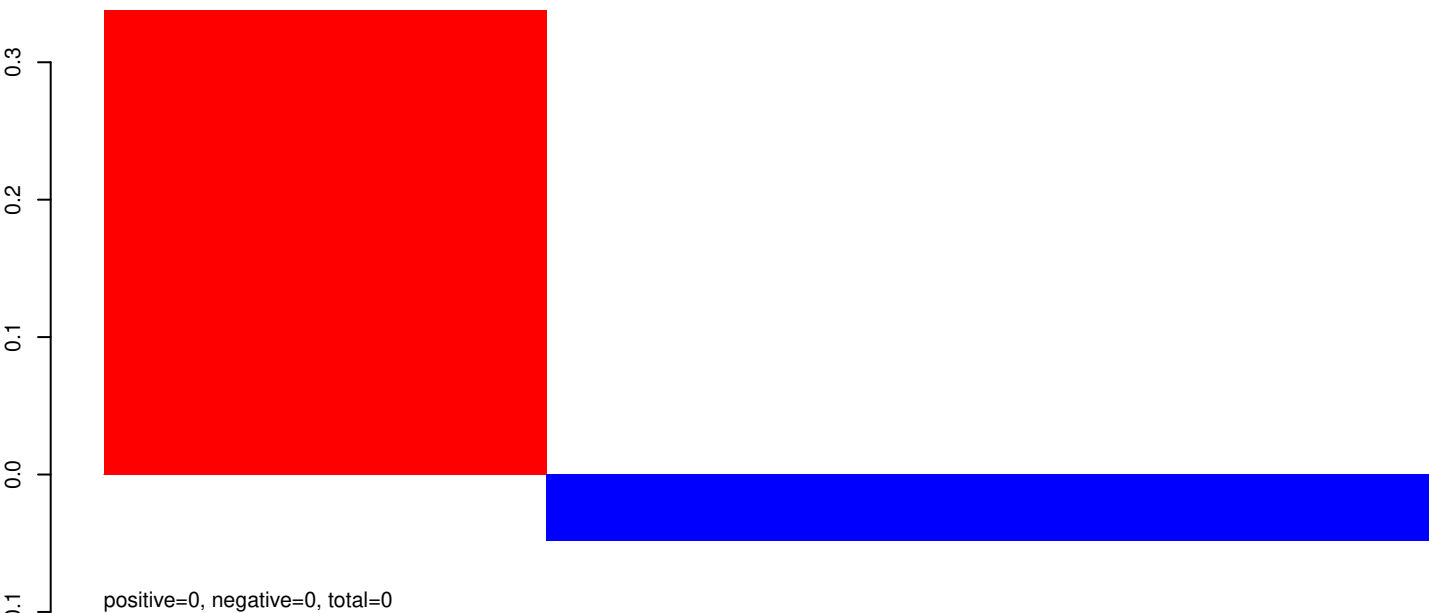
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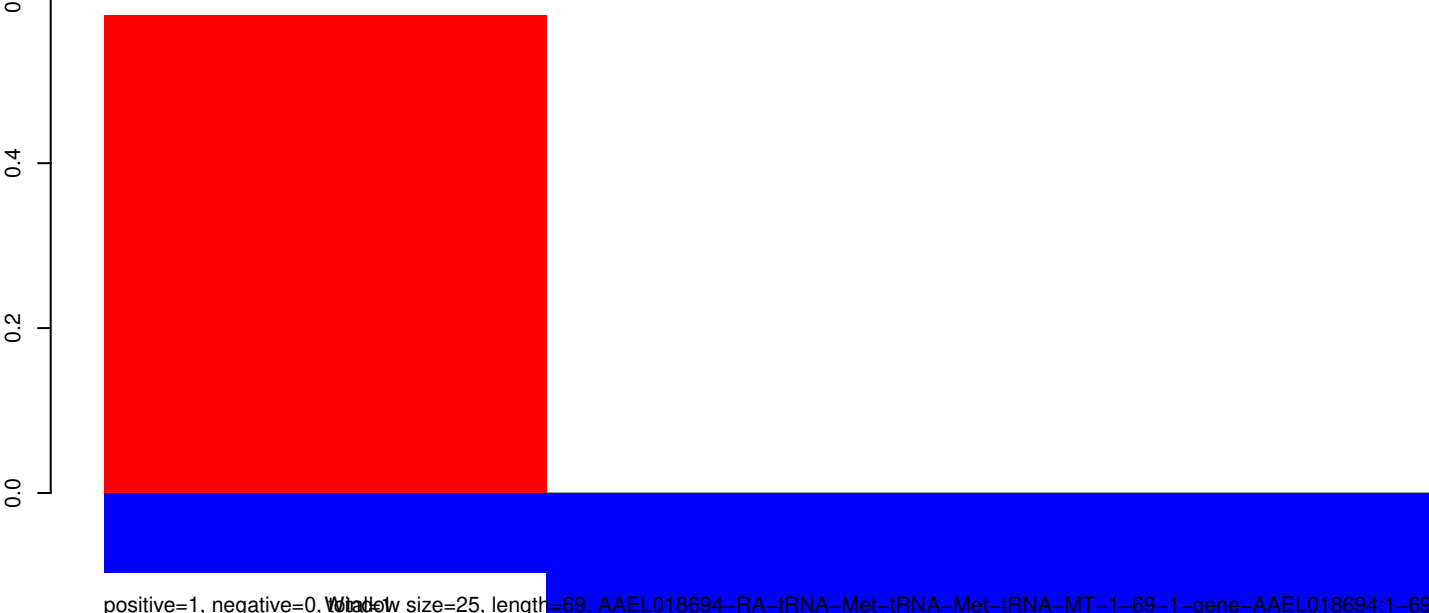
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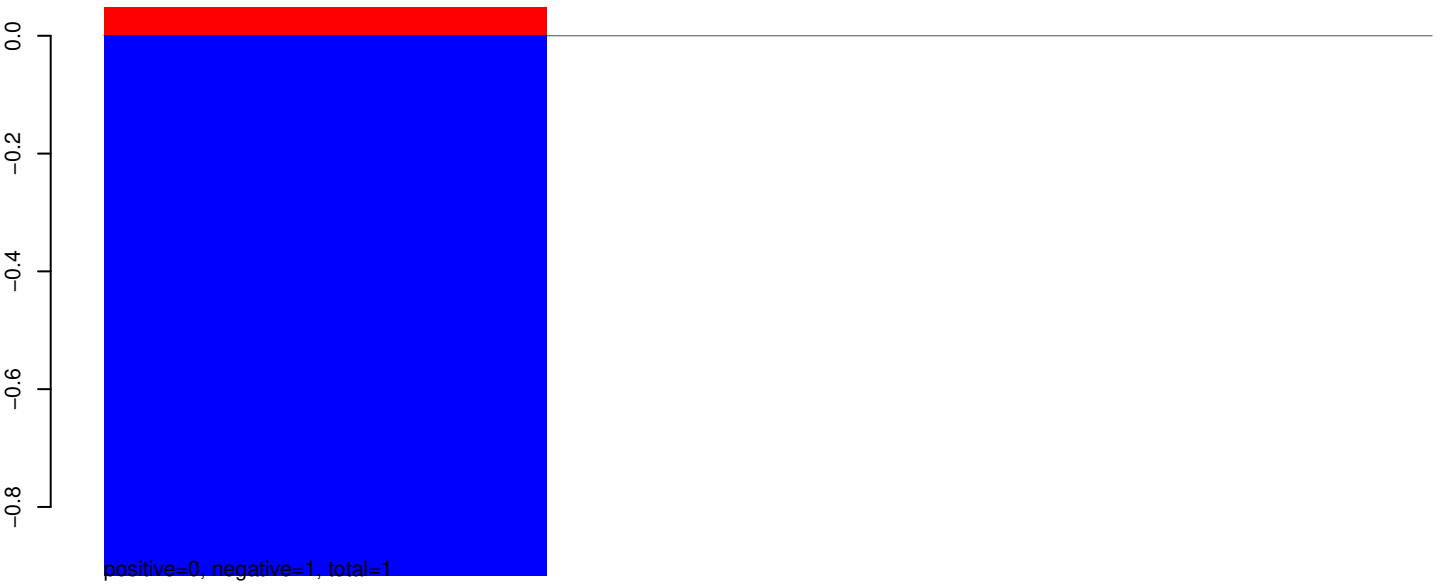
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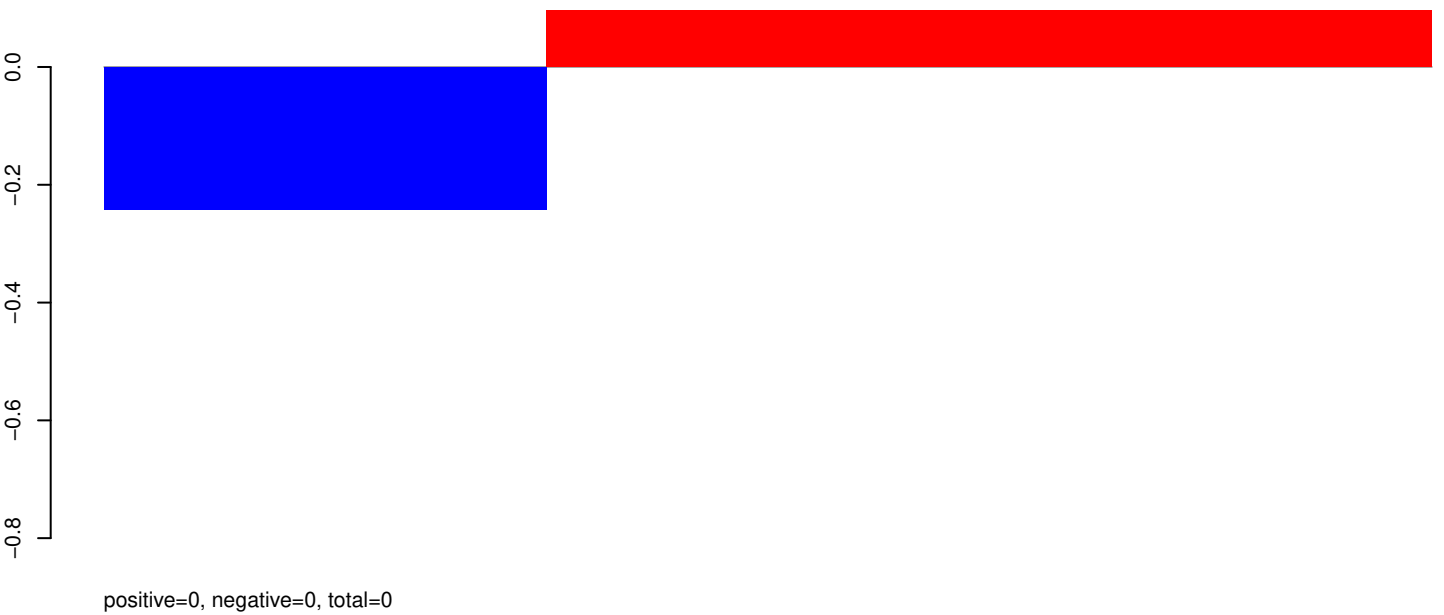
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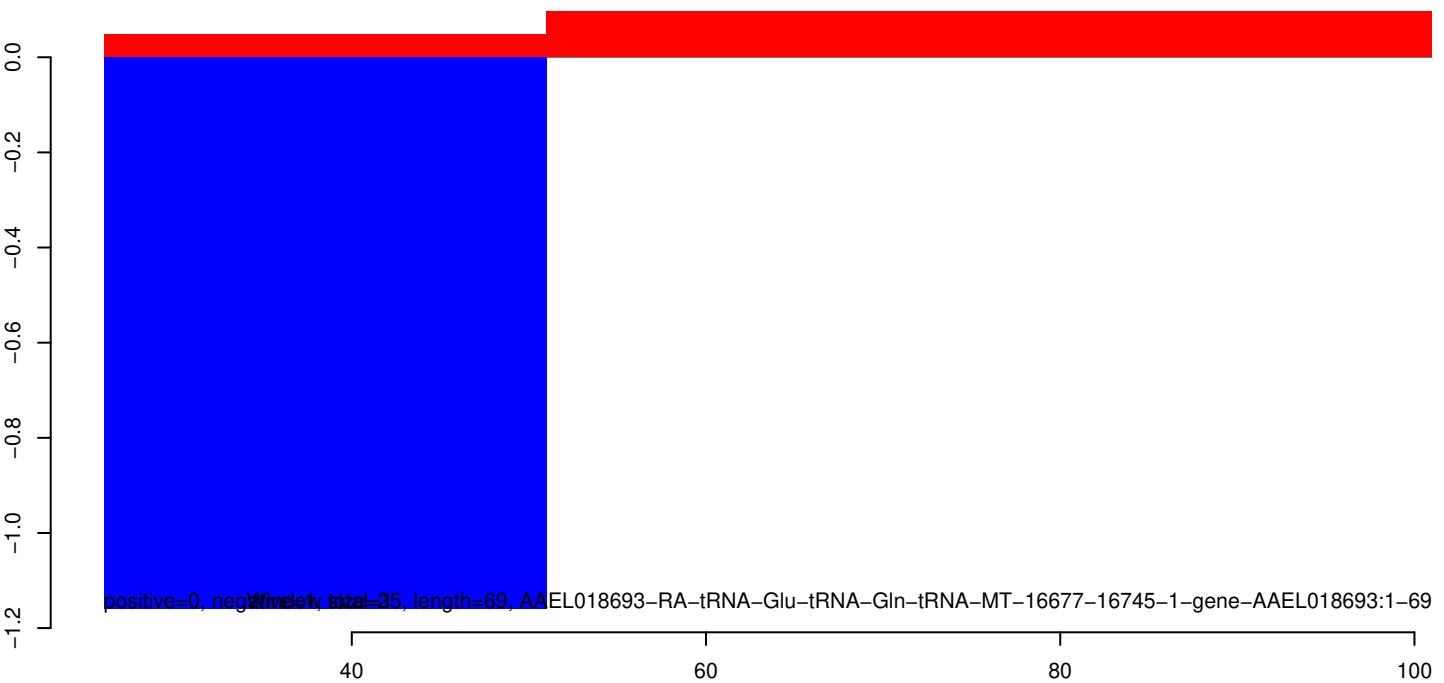
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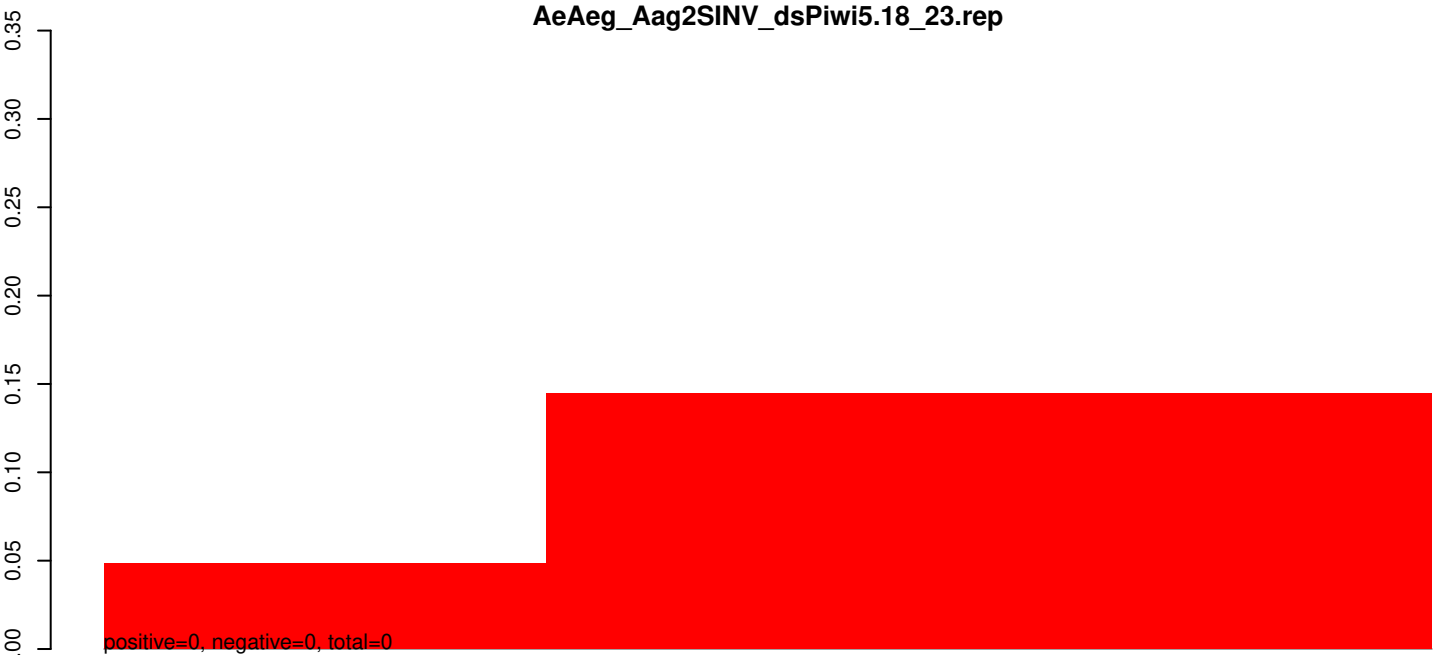
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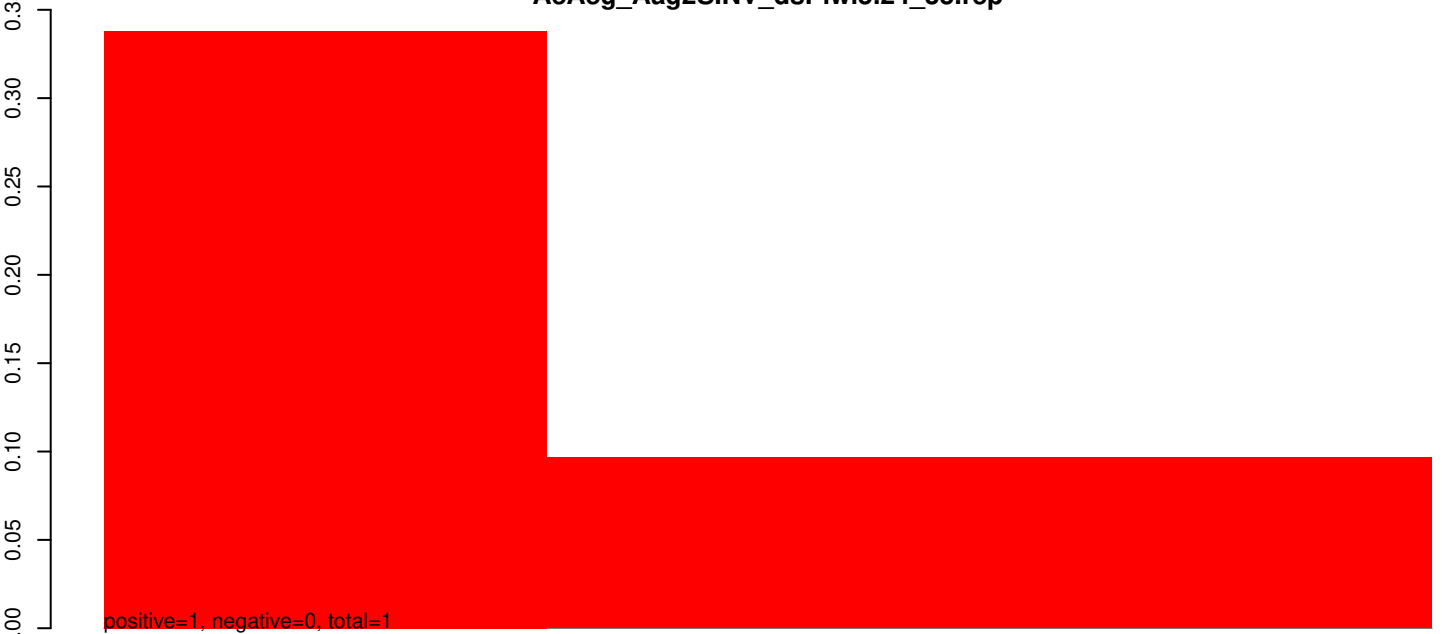
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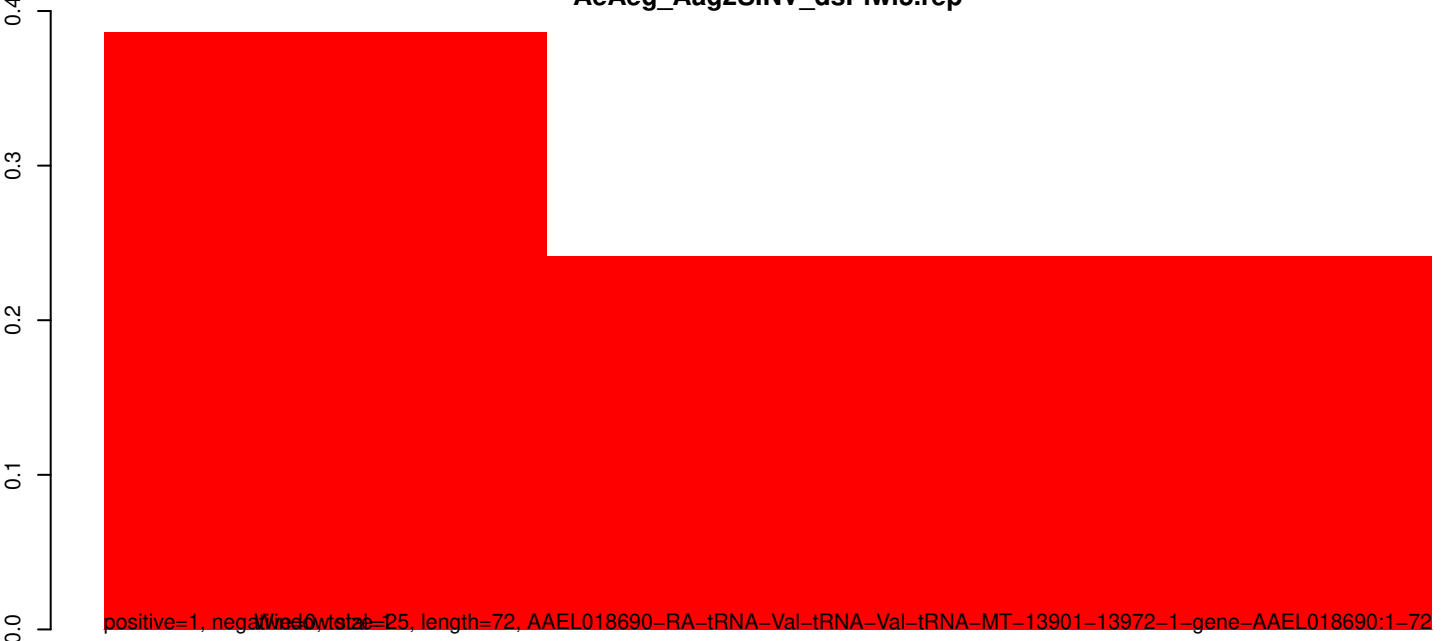
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AeAeg_Aag2SINV_dsPiwi5.24_35.rep



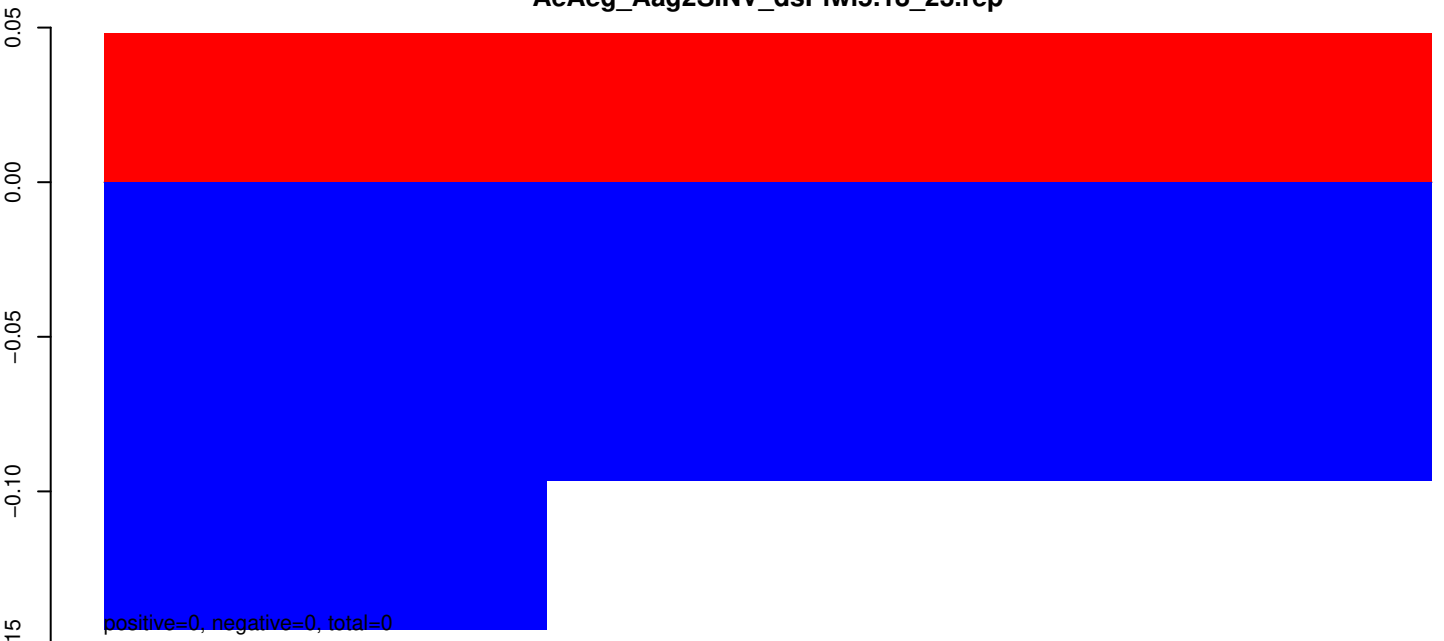
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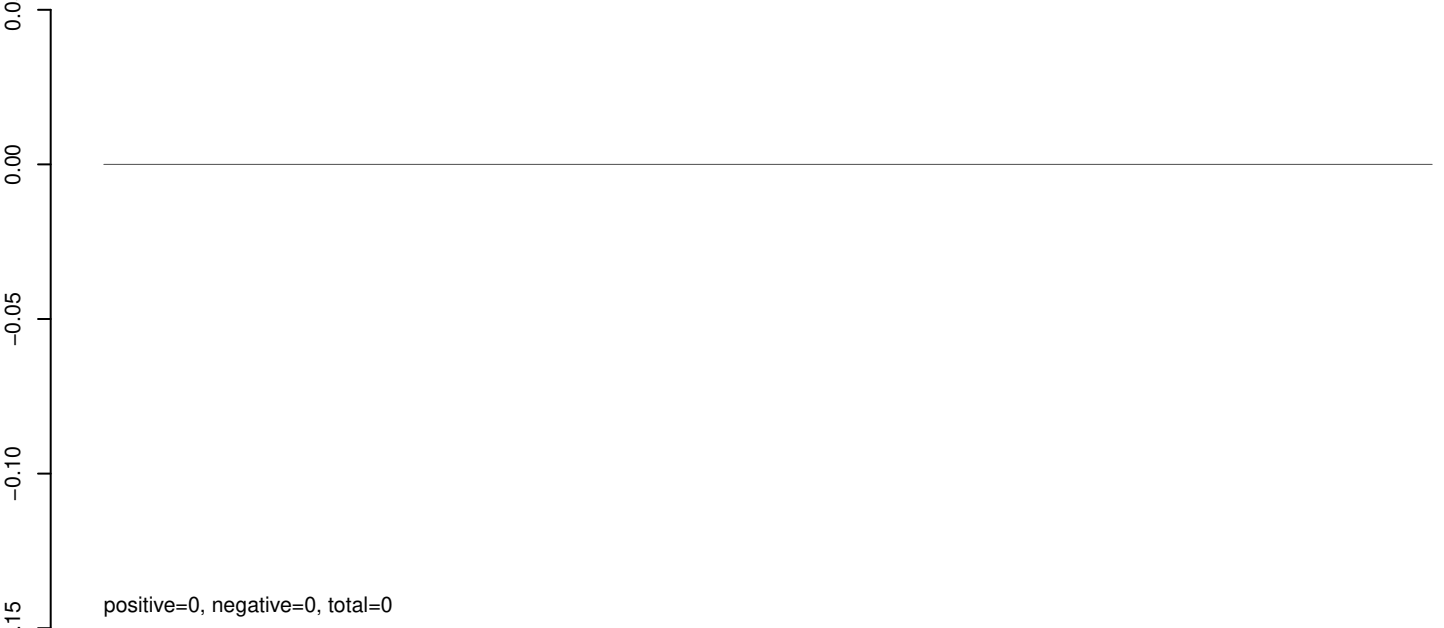
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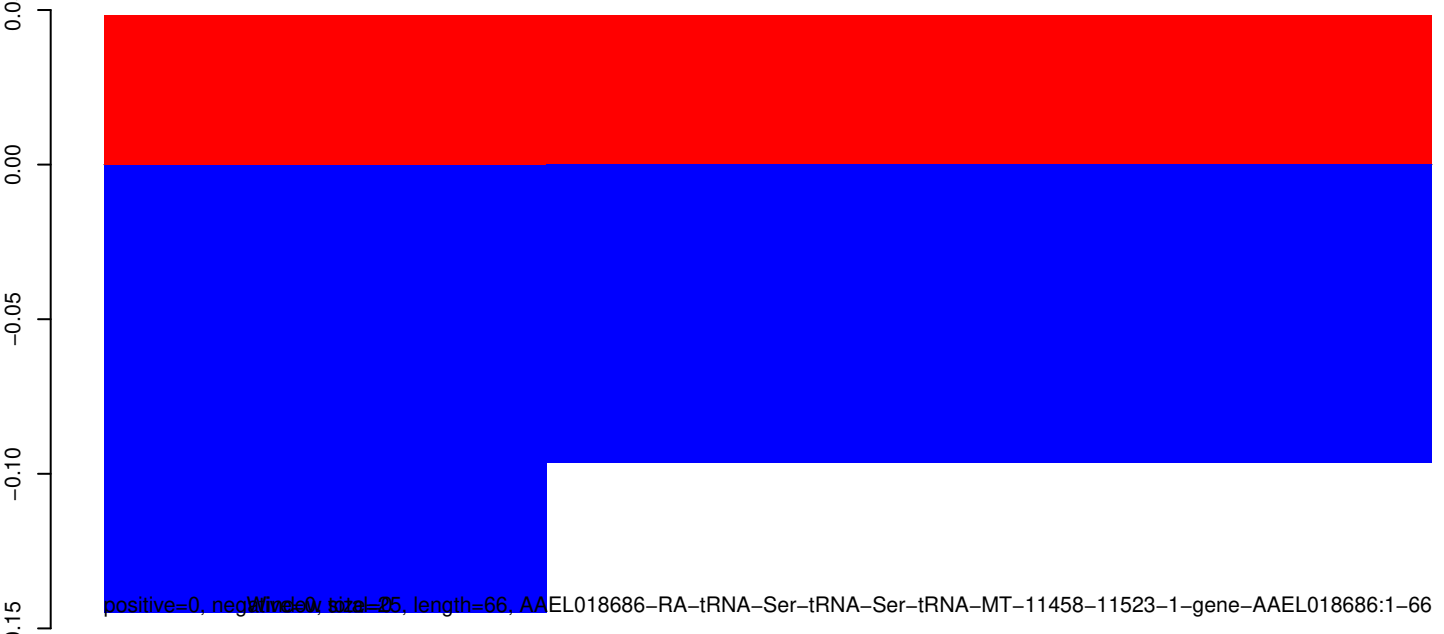
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AeAeg_Aag2SINV_dsPiwi5.24_35.rep

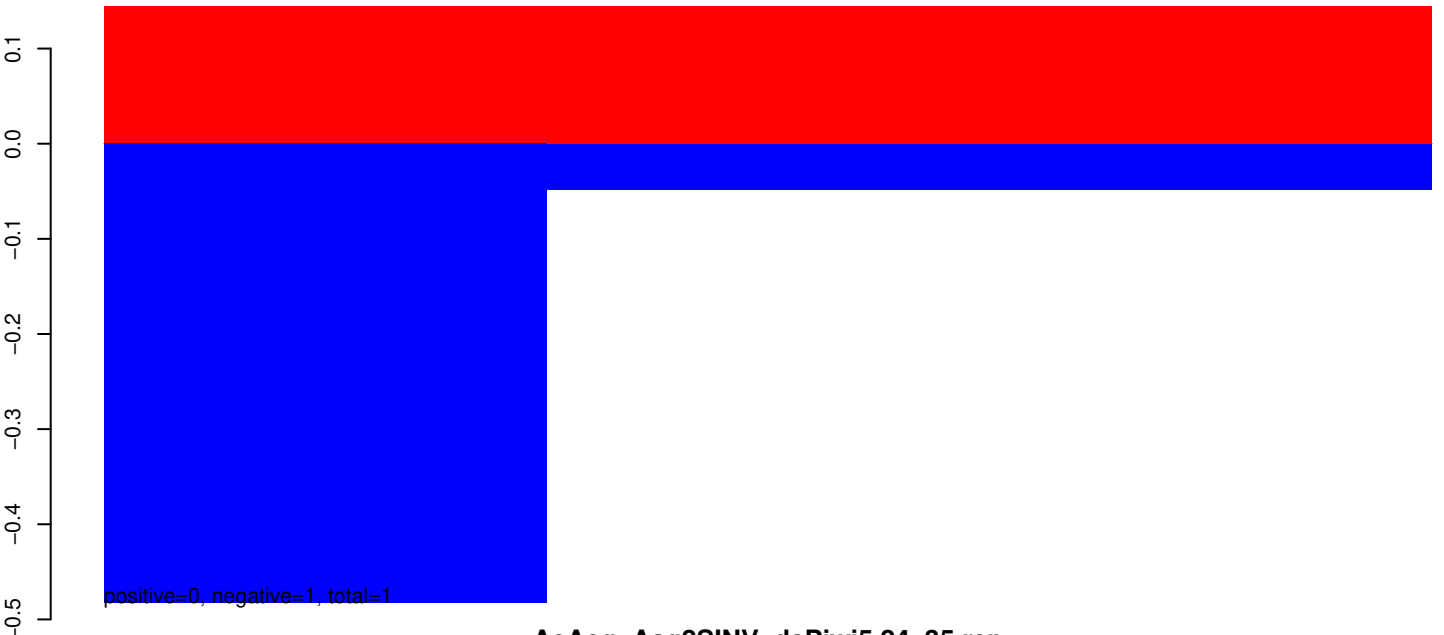


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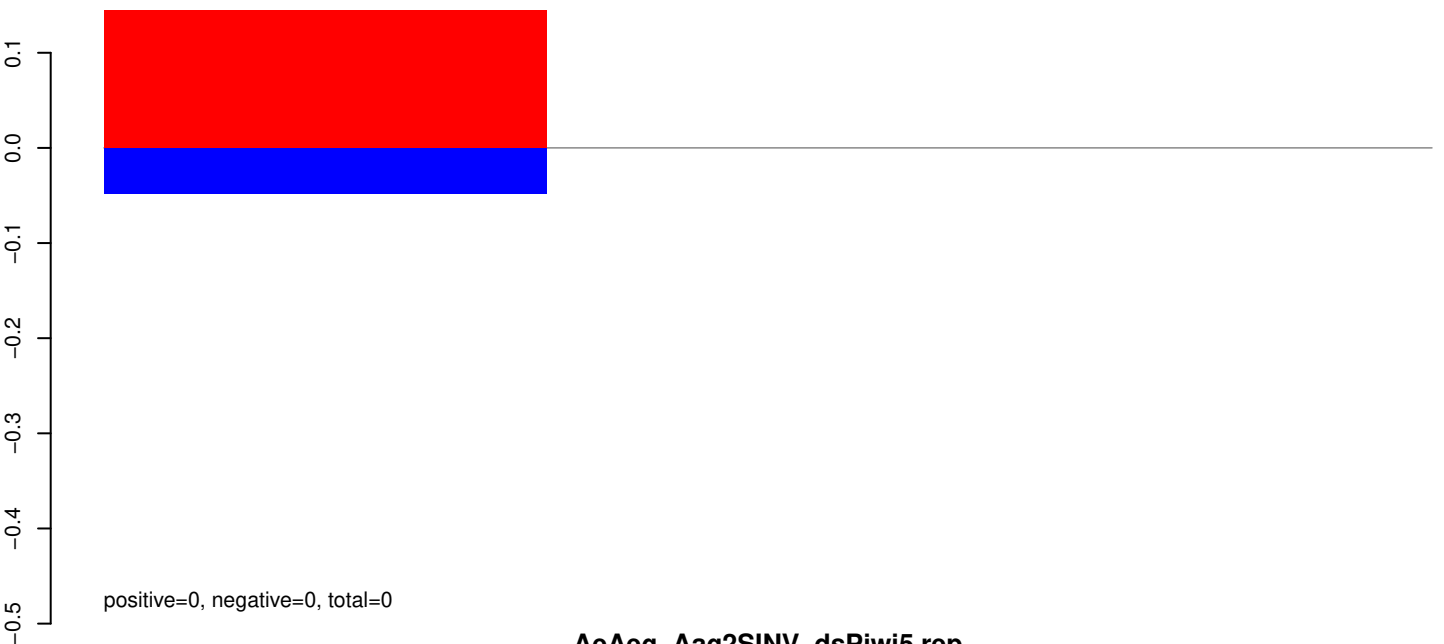


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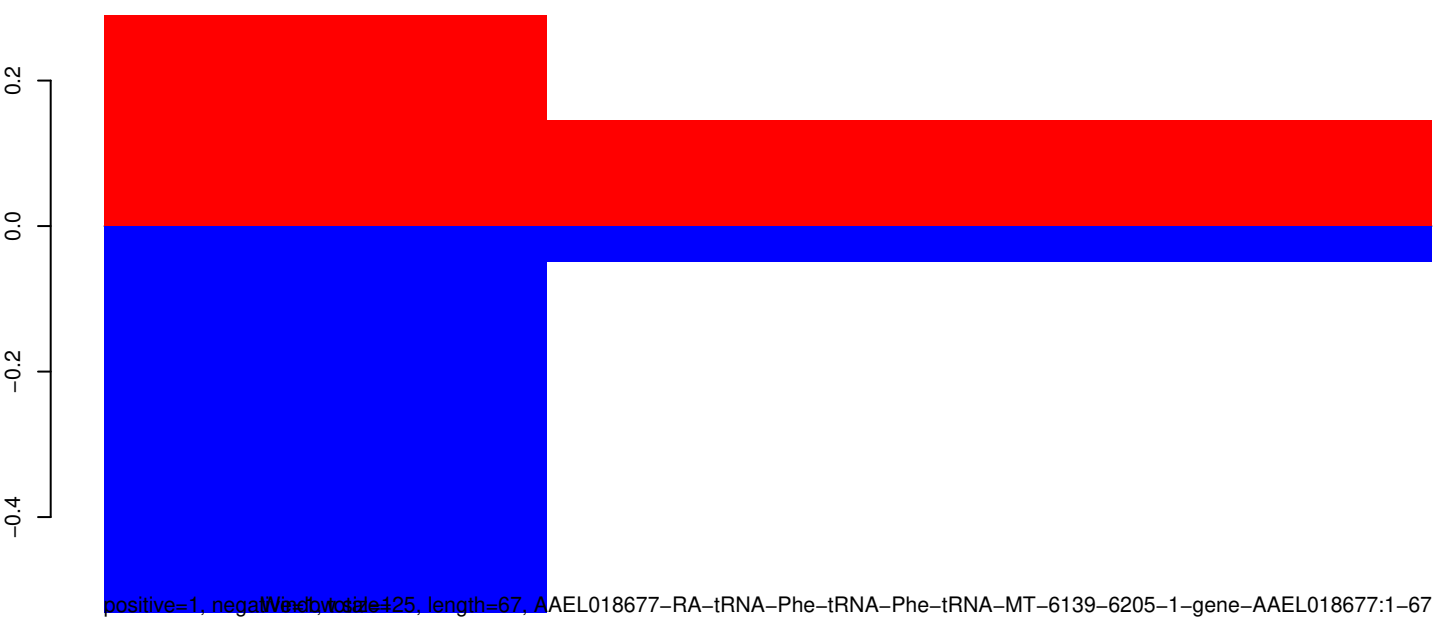
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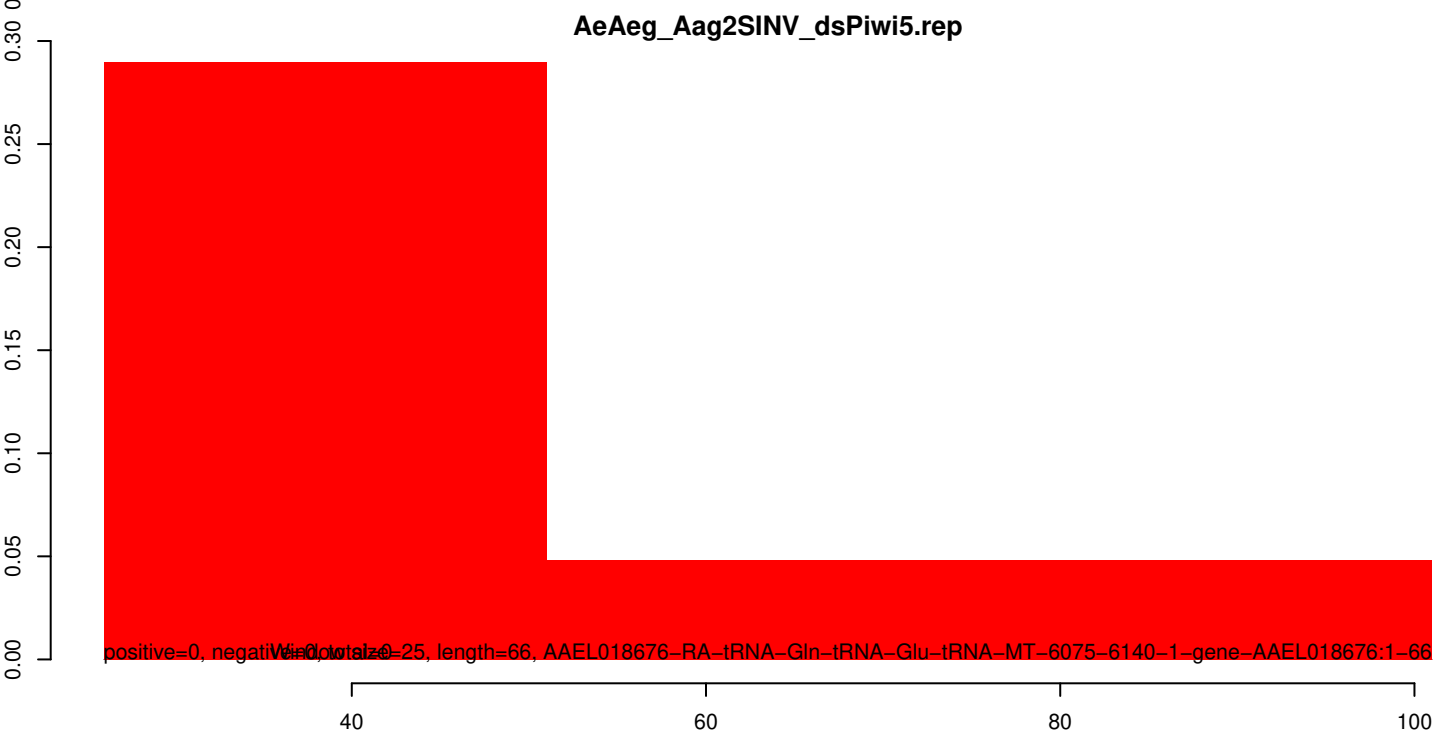
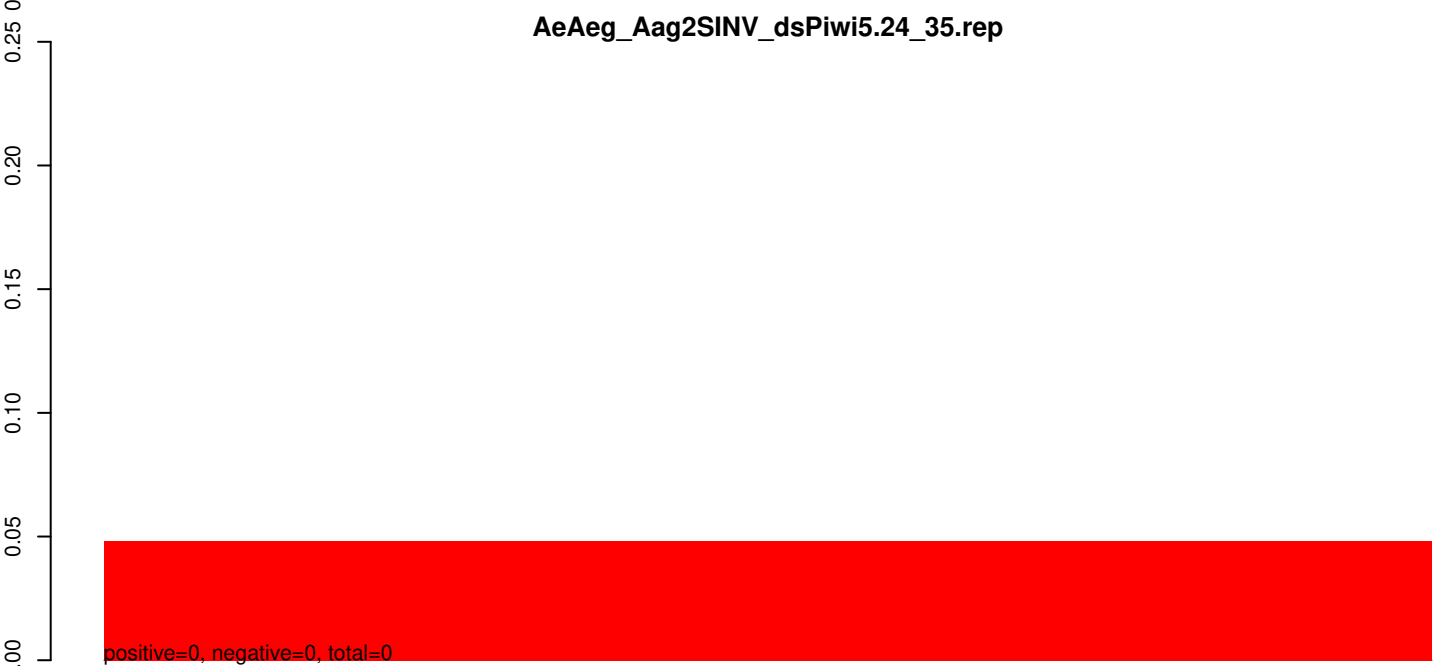
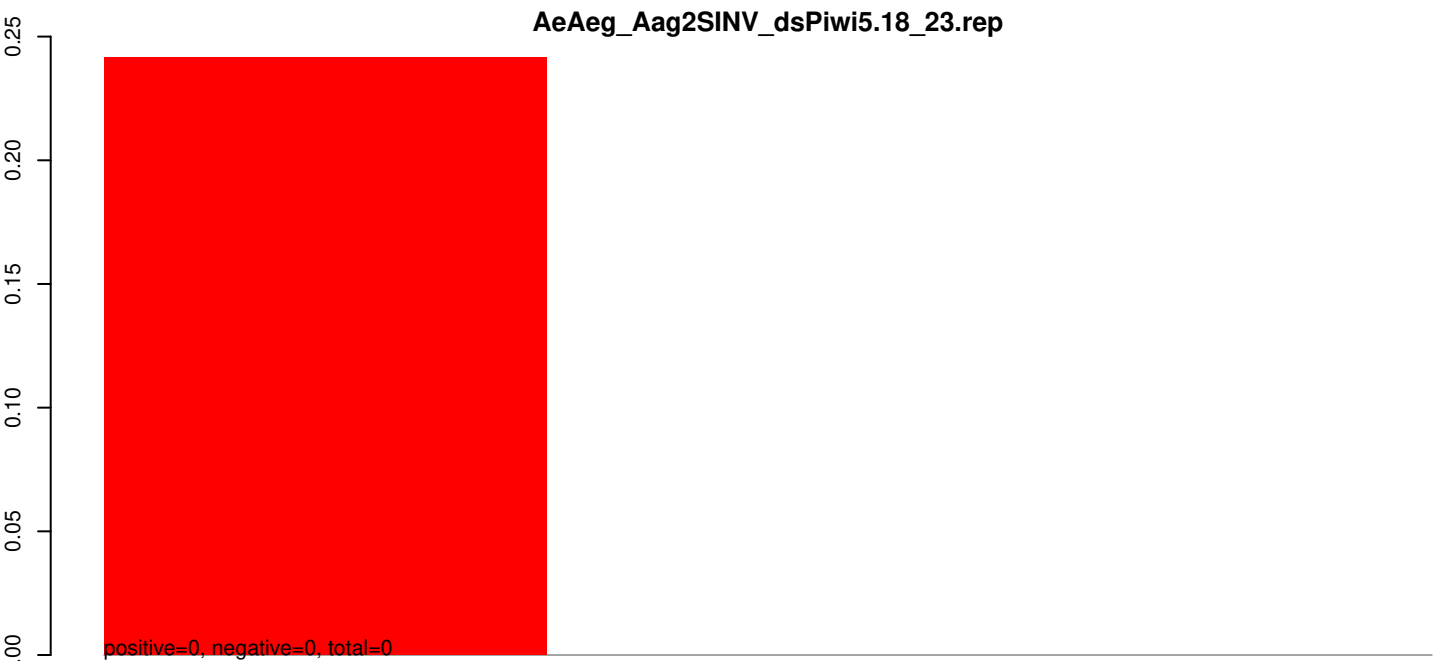


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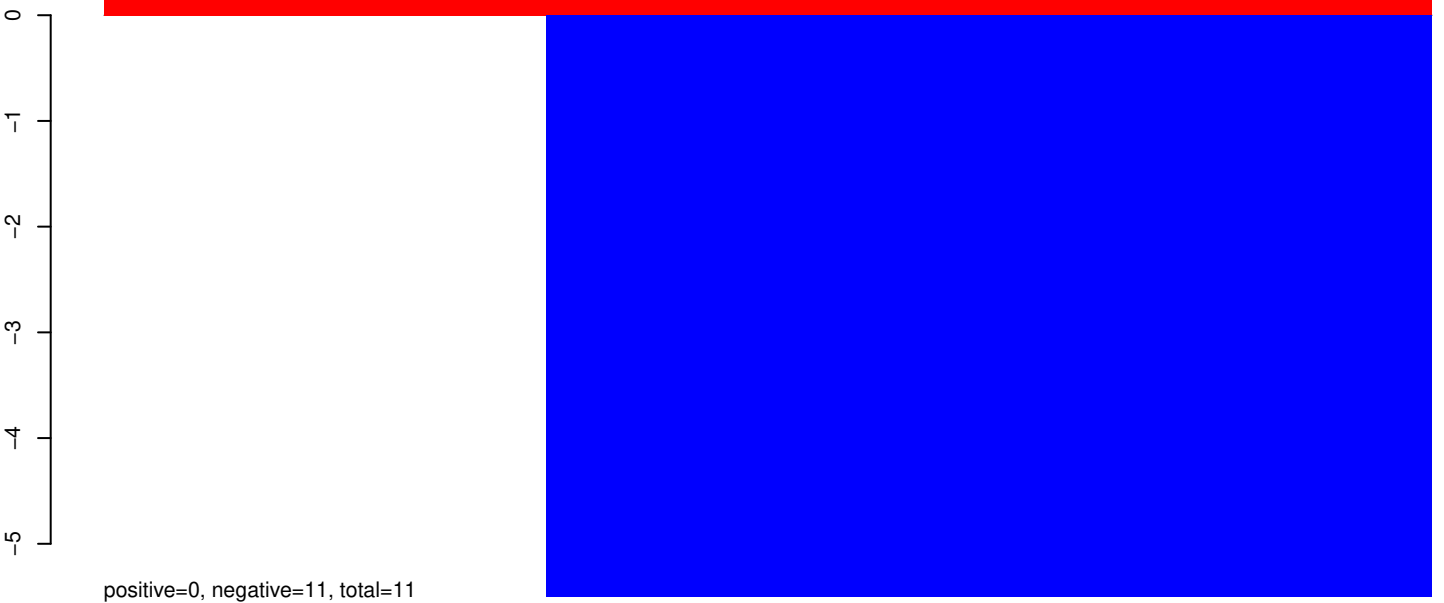


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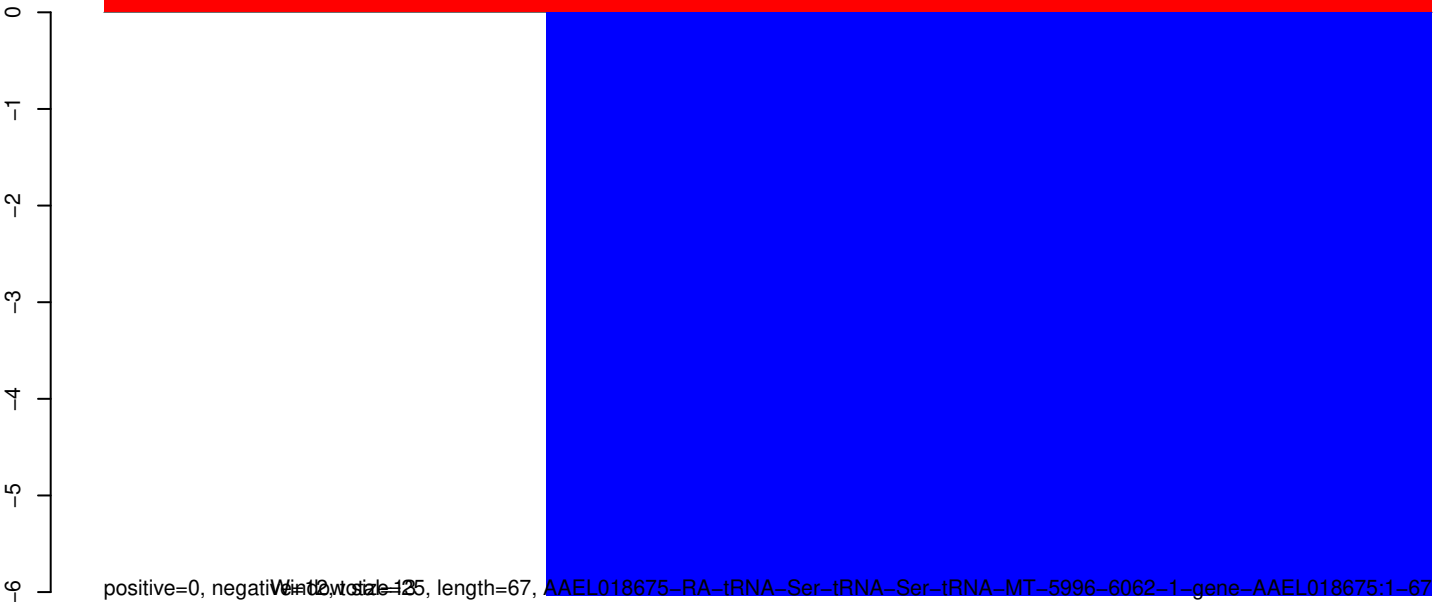
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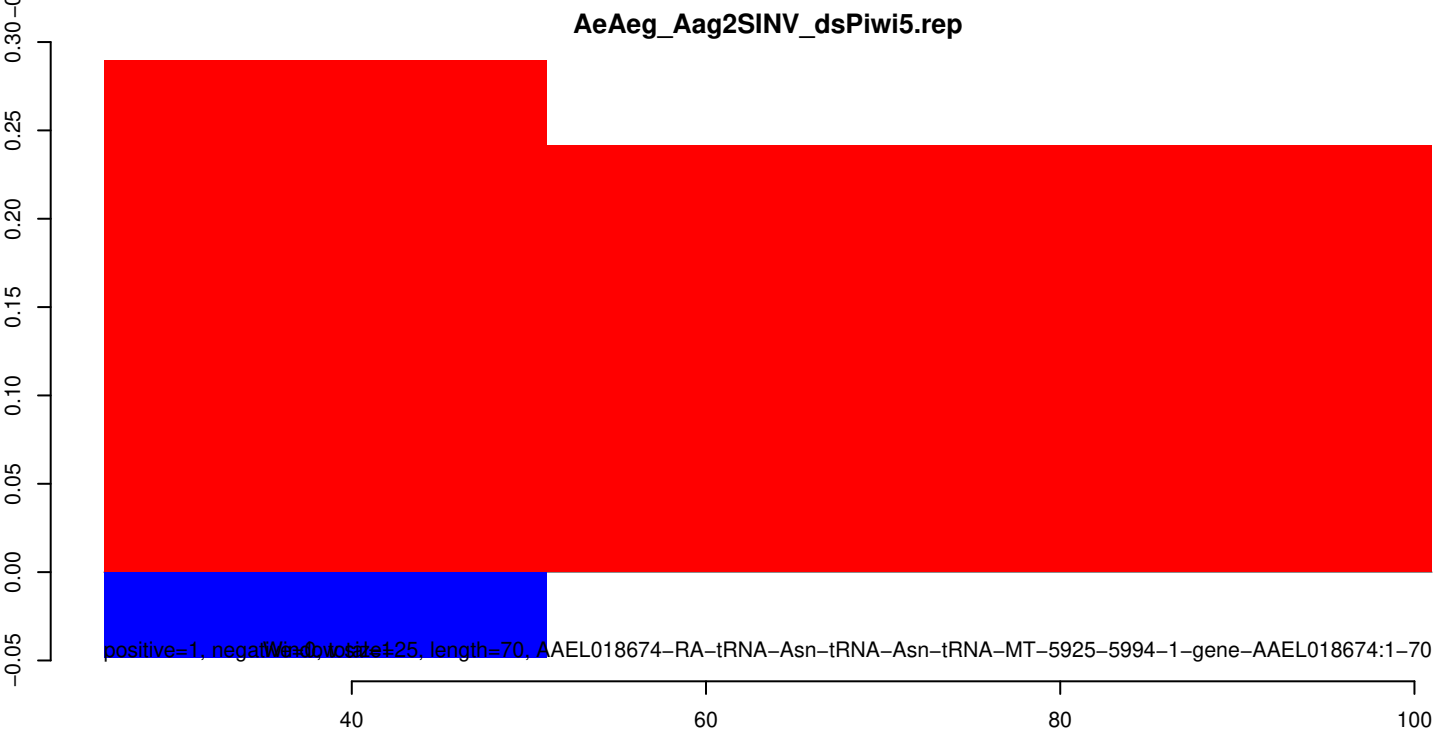
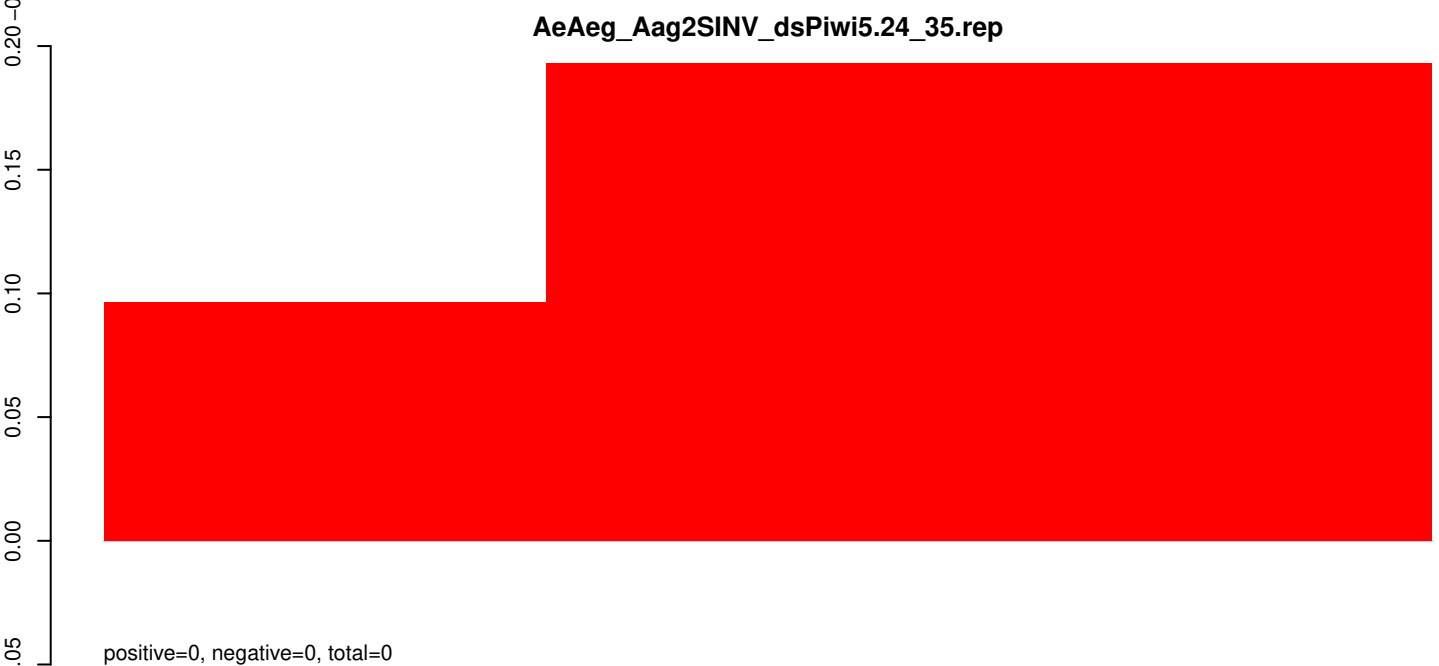
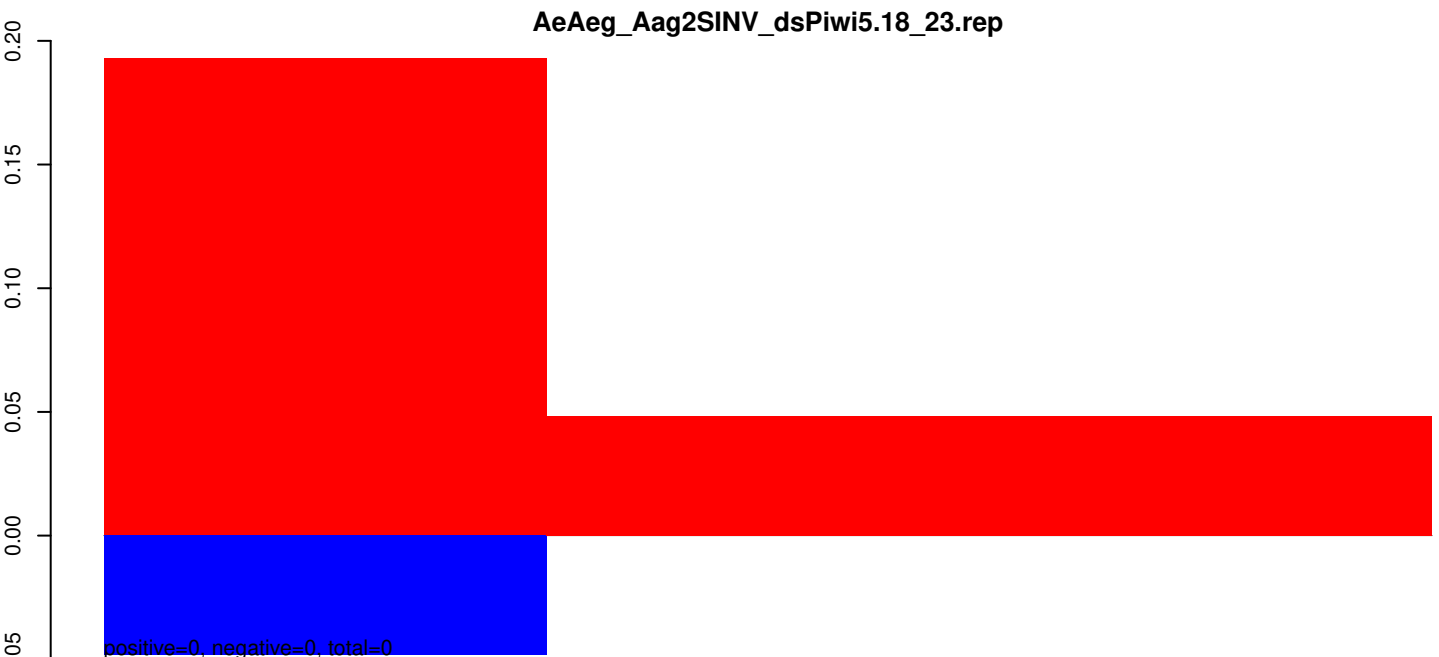


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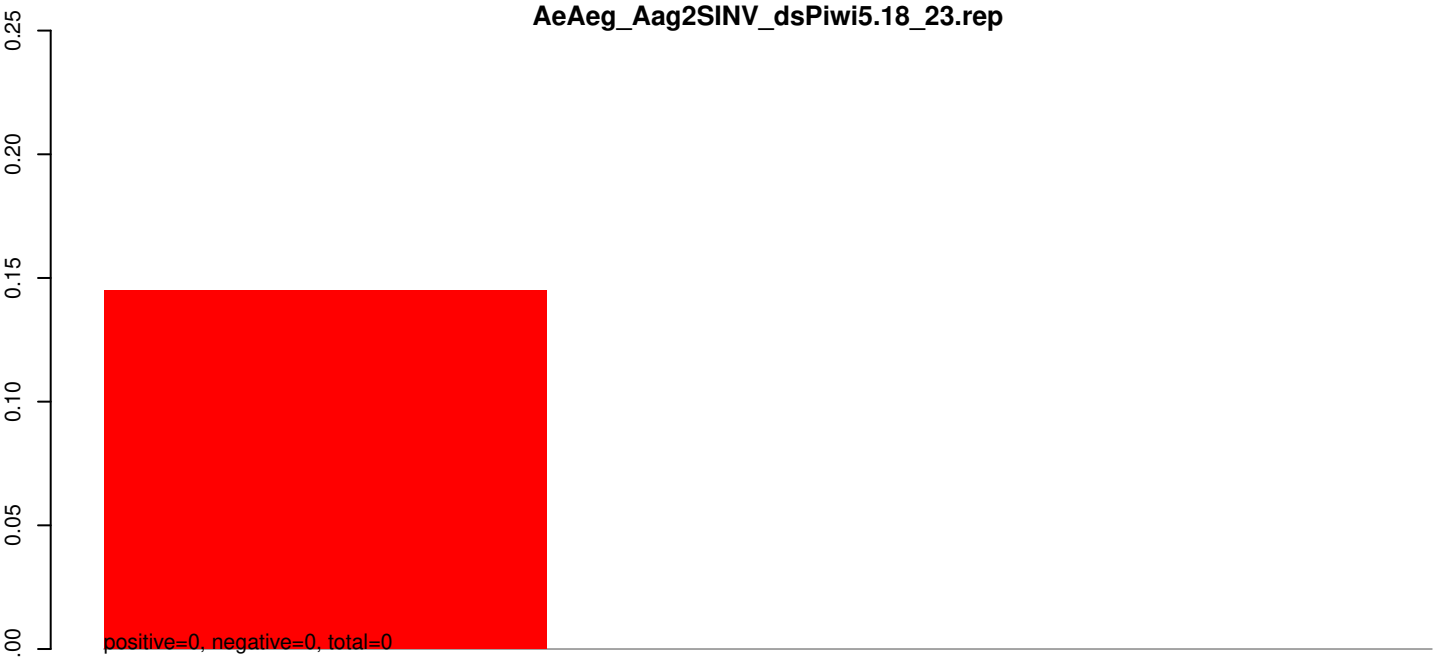


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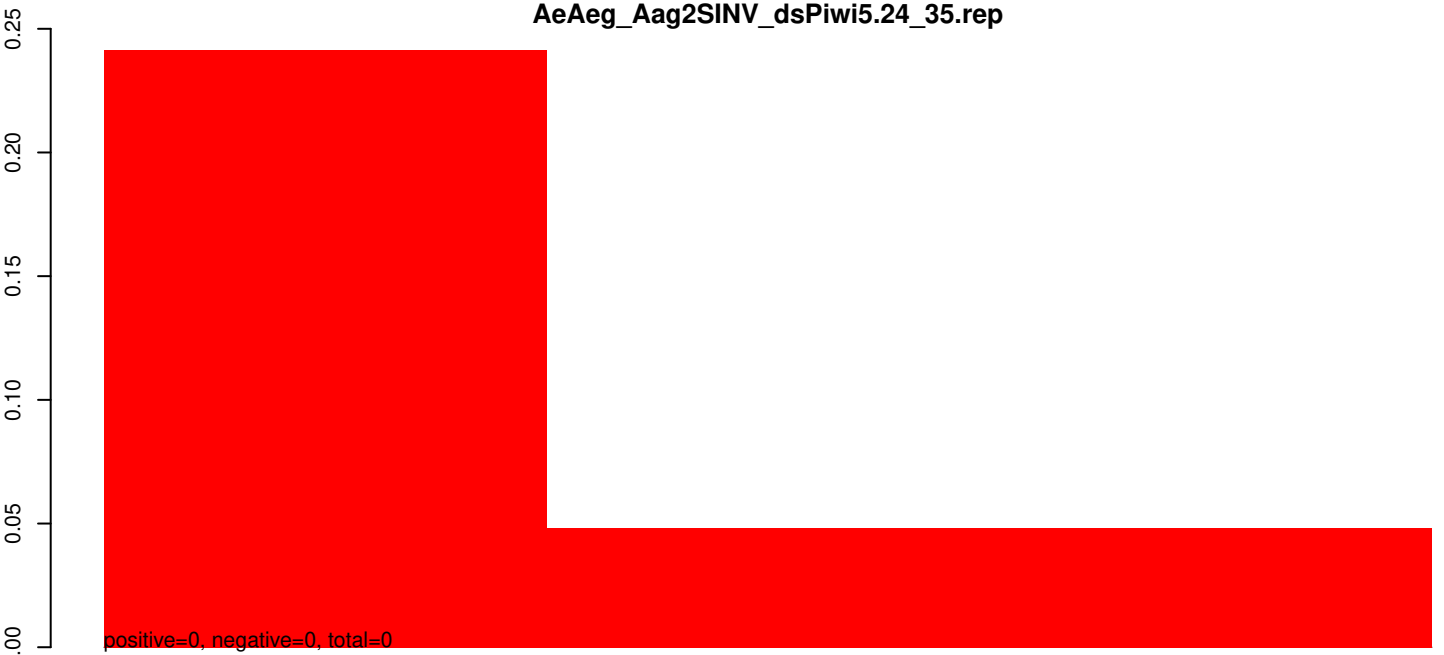
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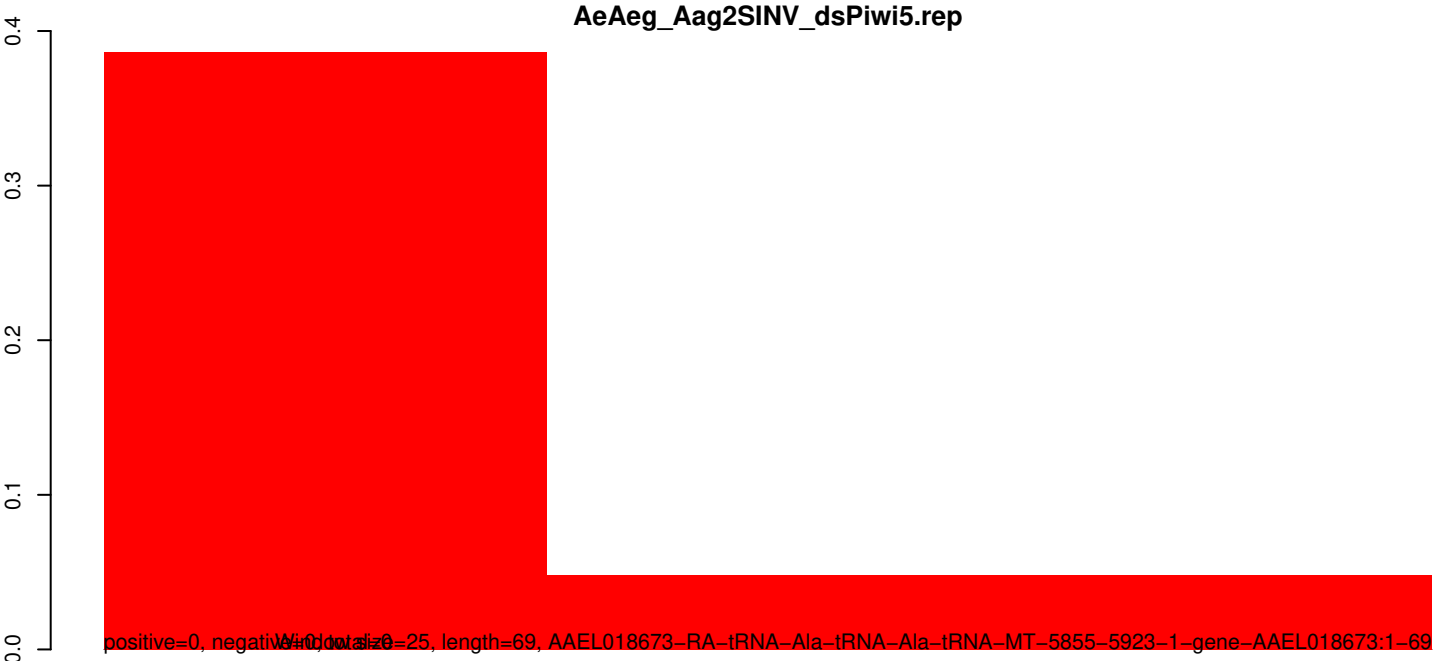
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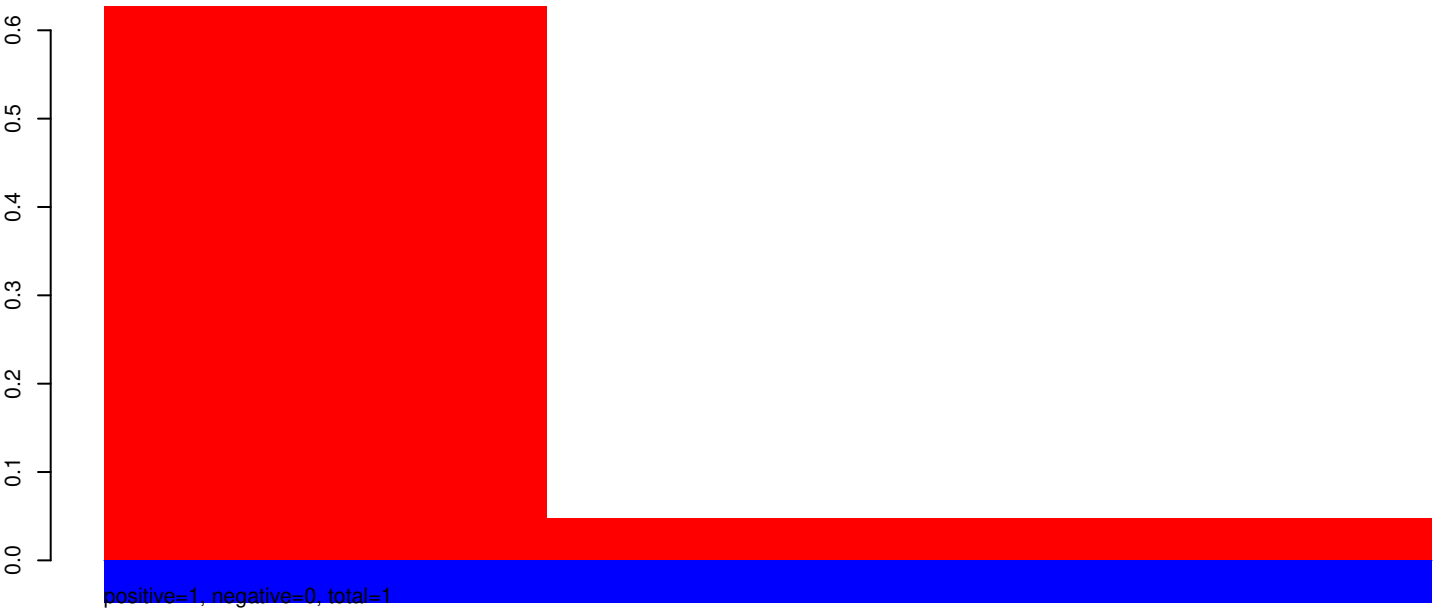


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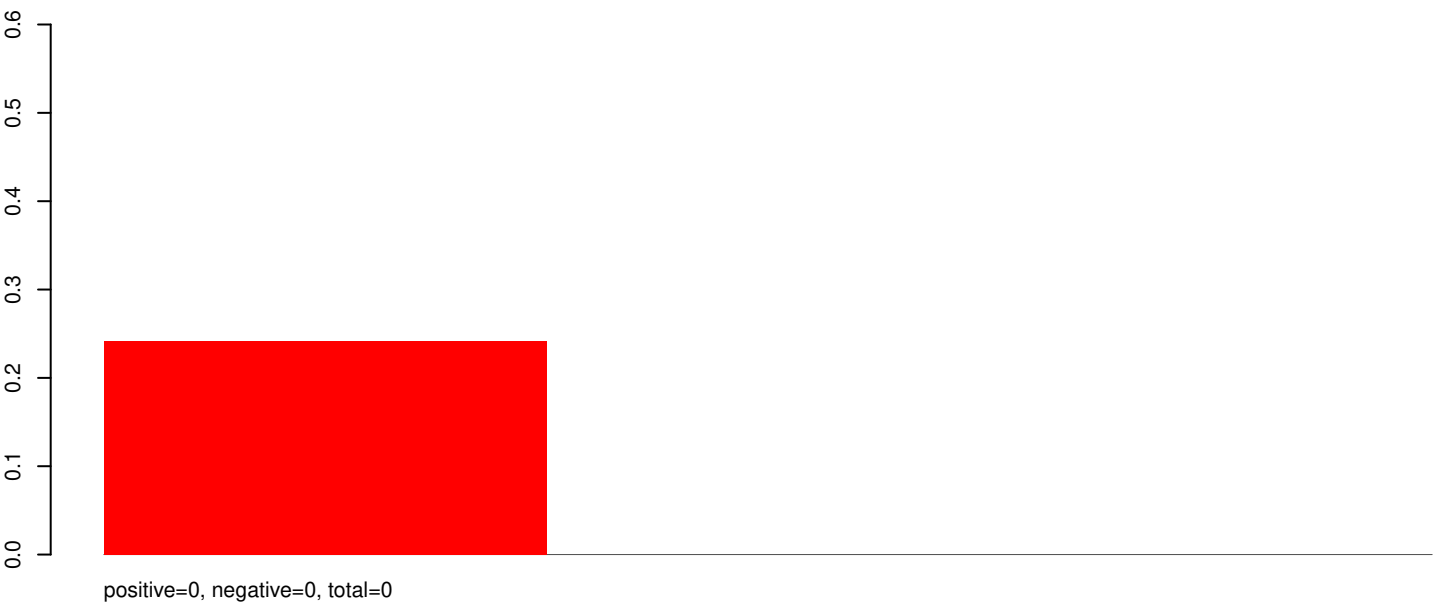


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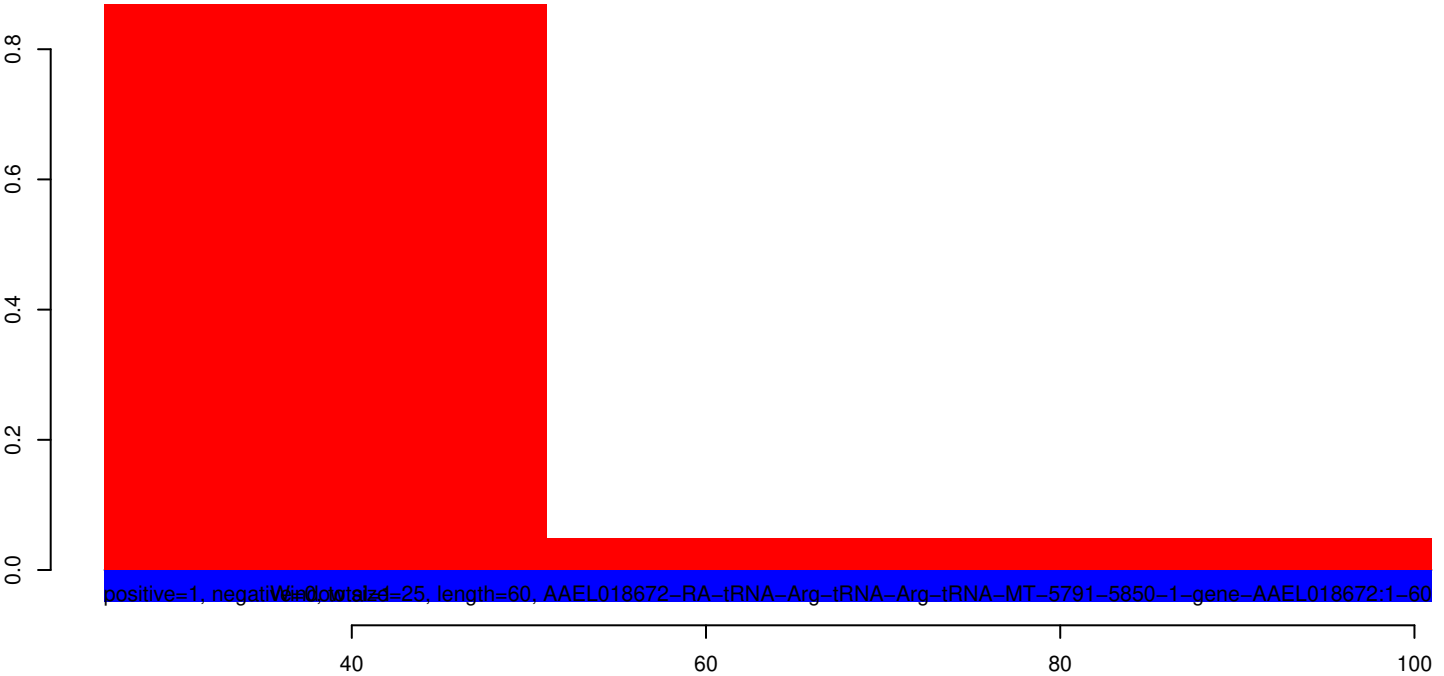
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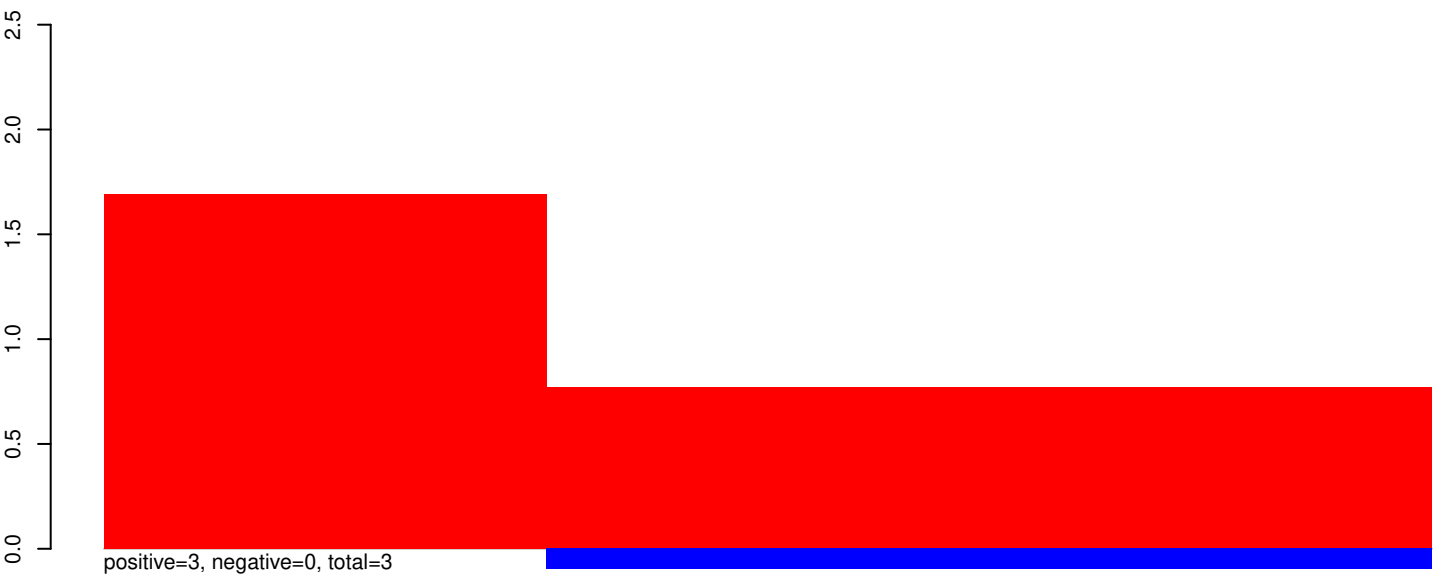
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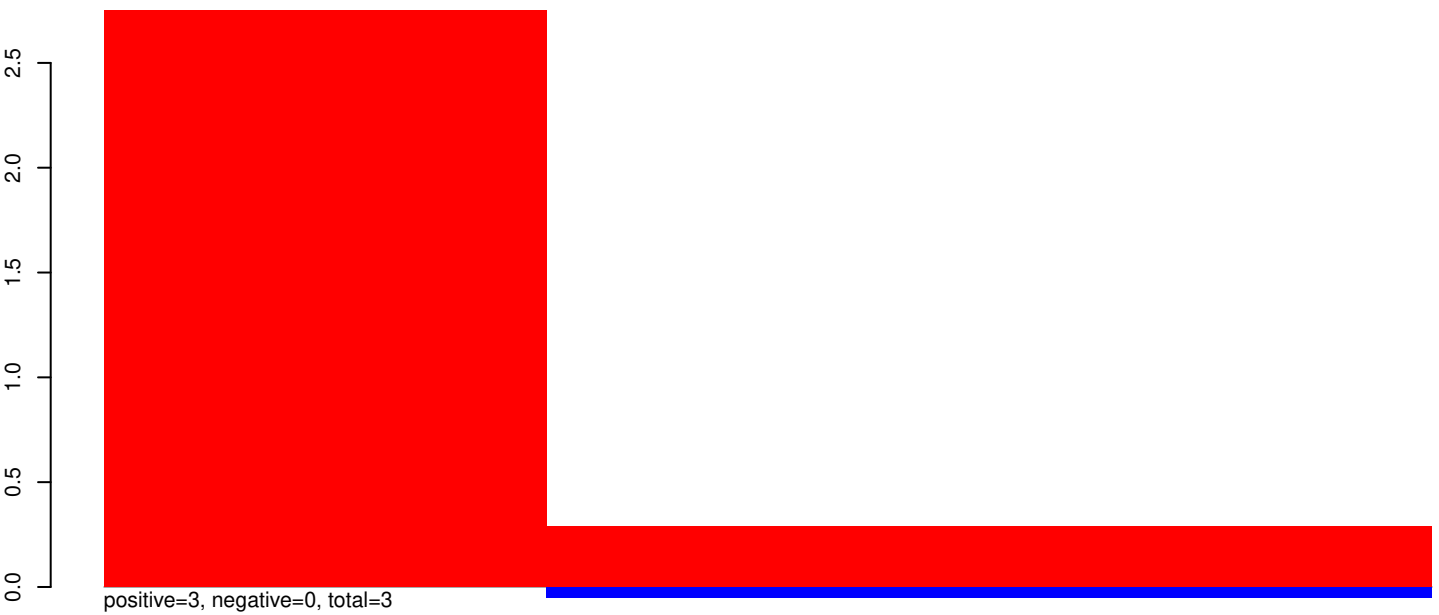
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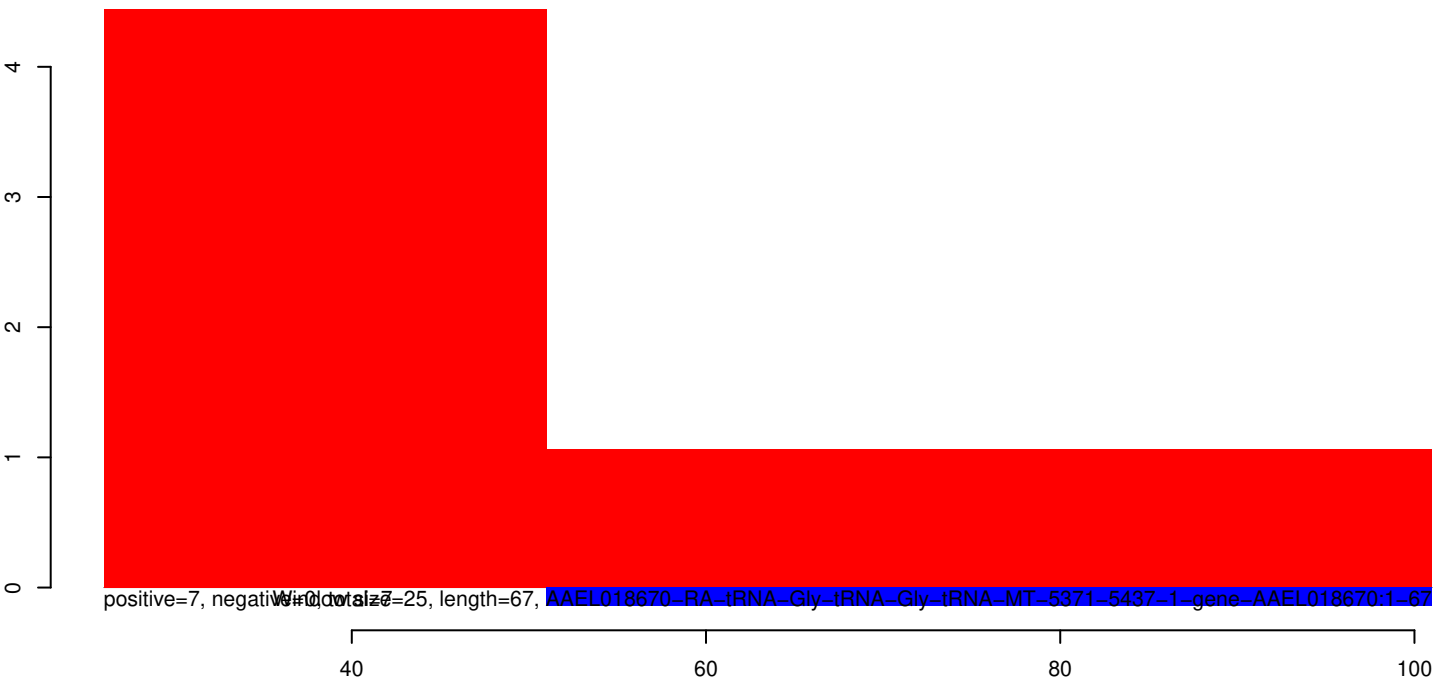
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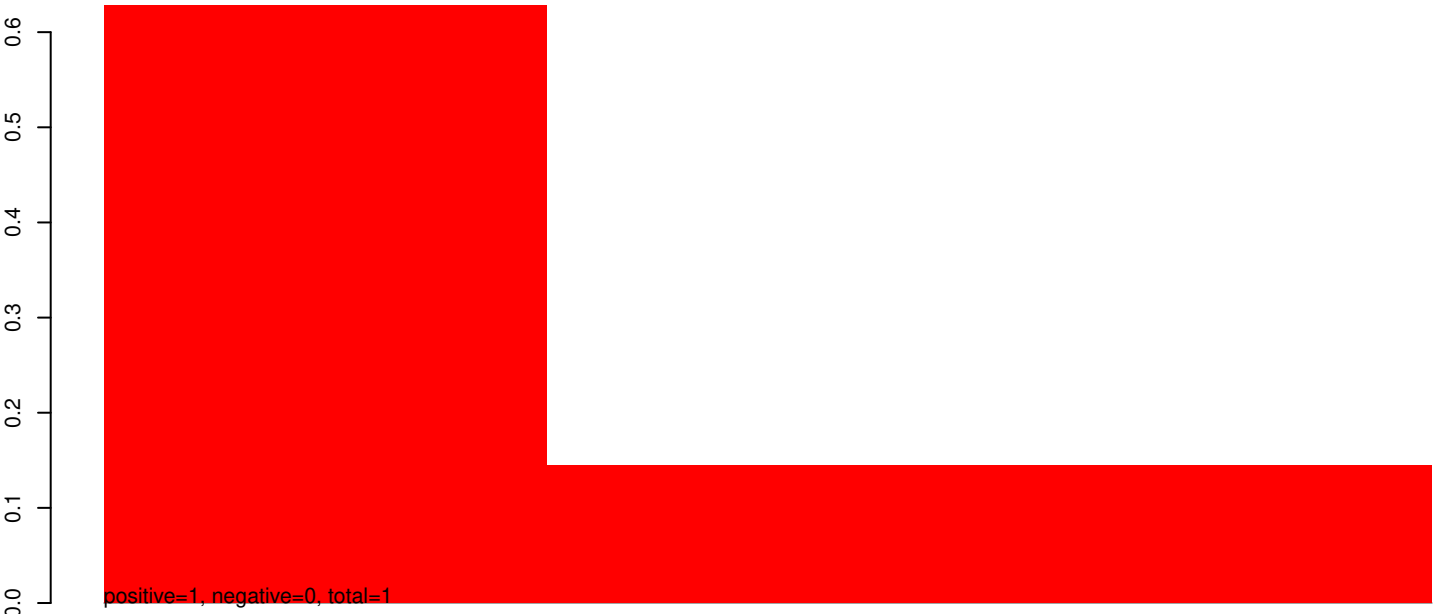
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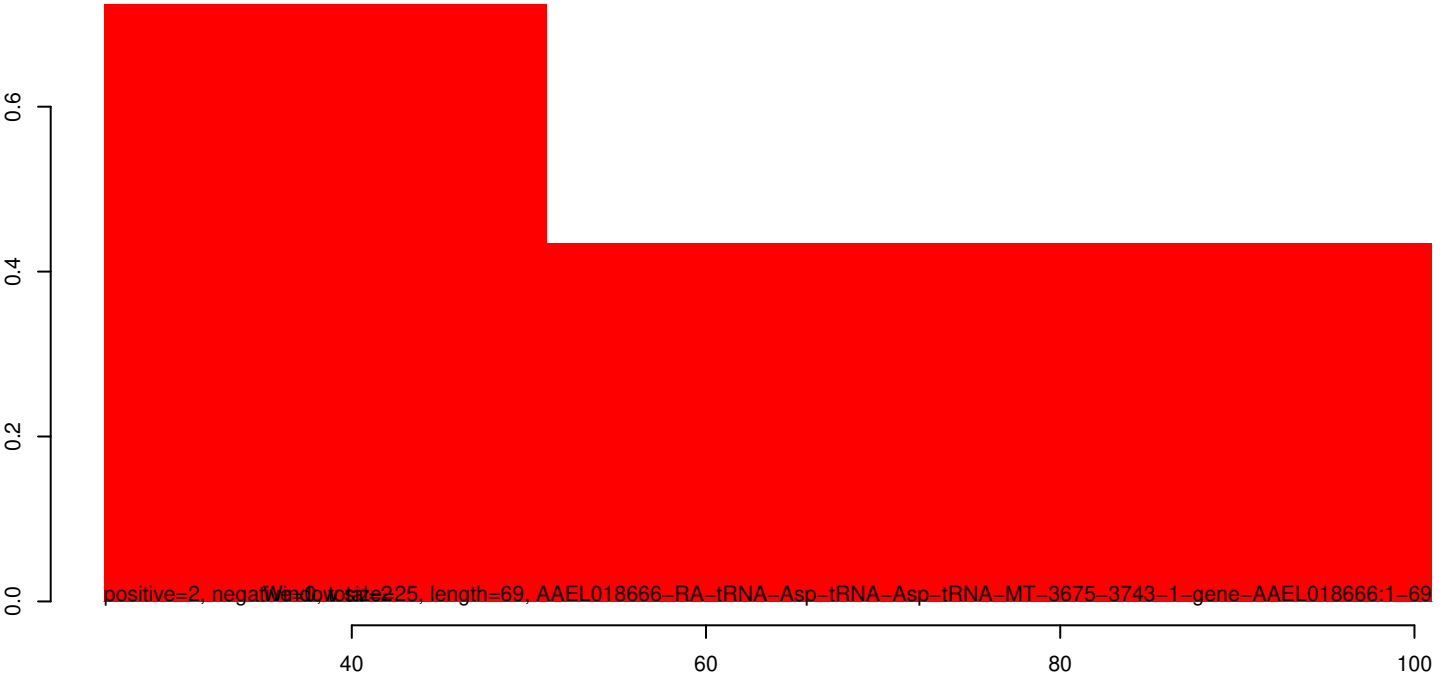
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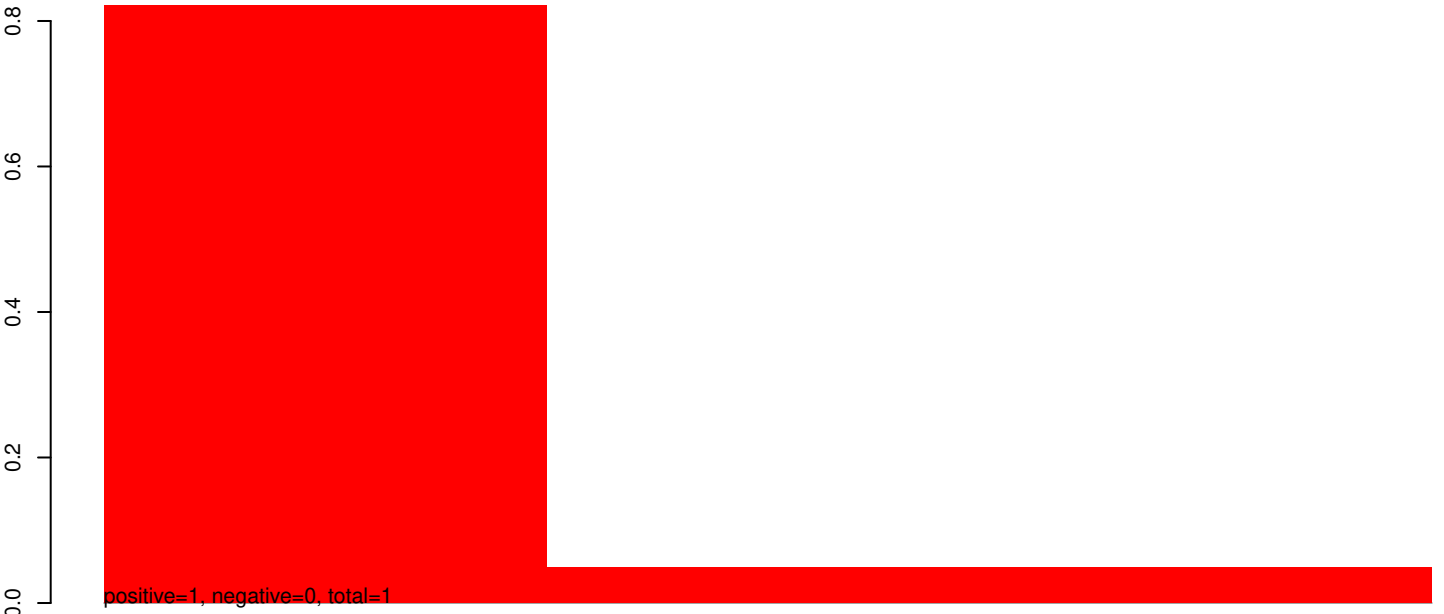
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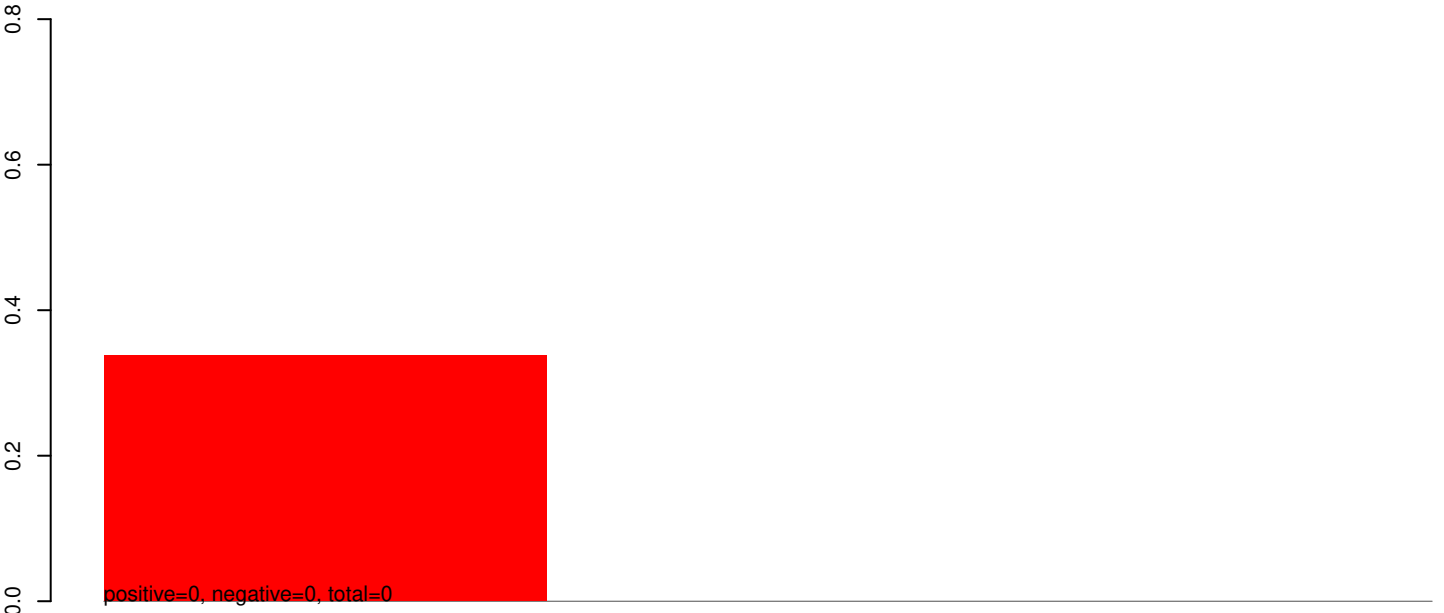
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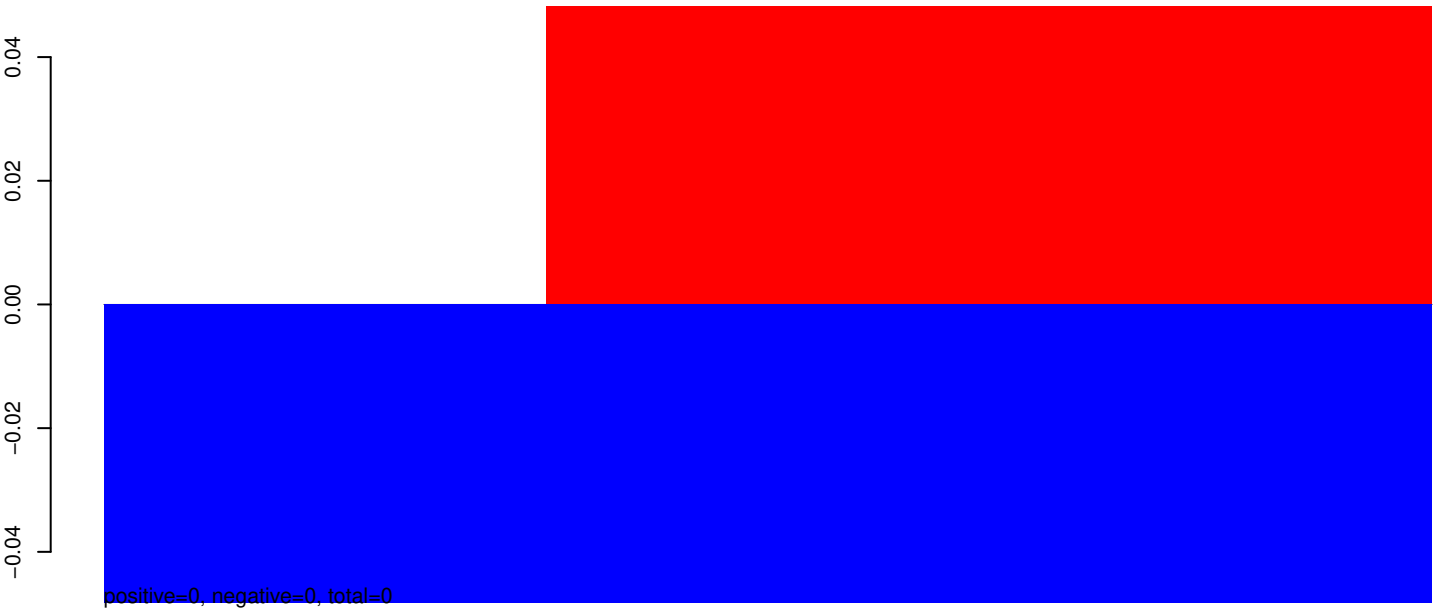


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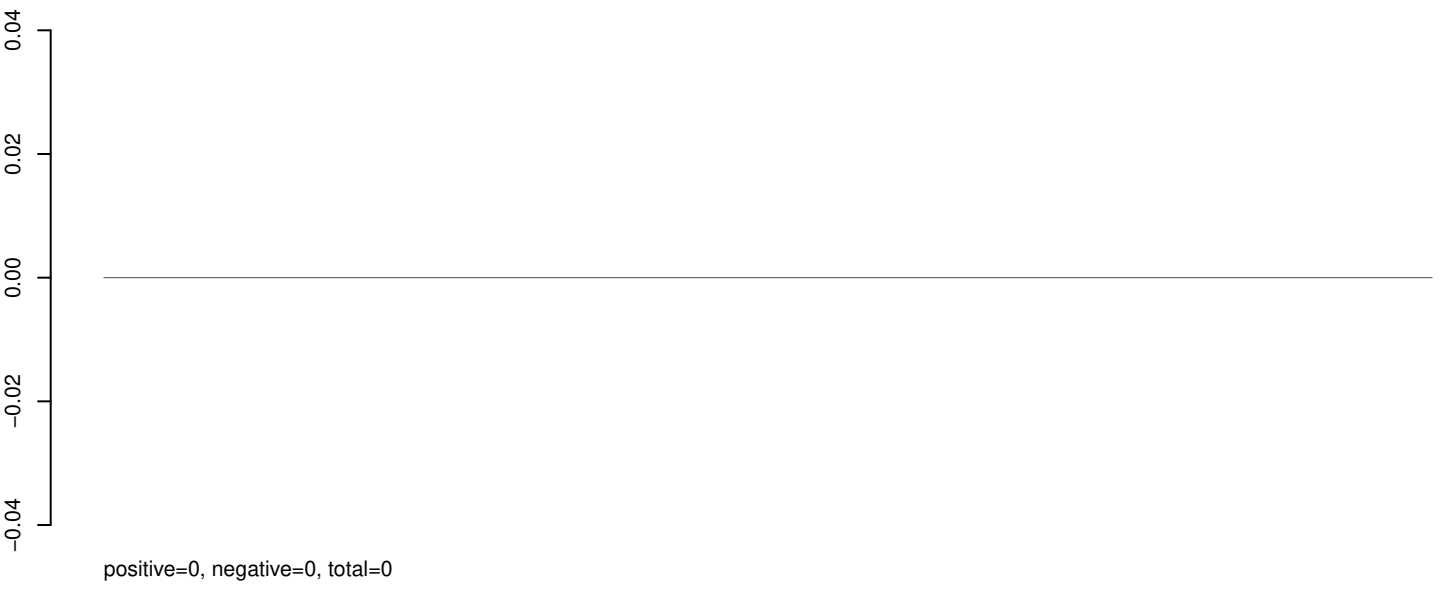


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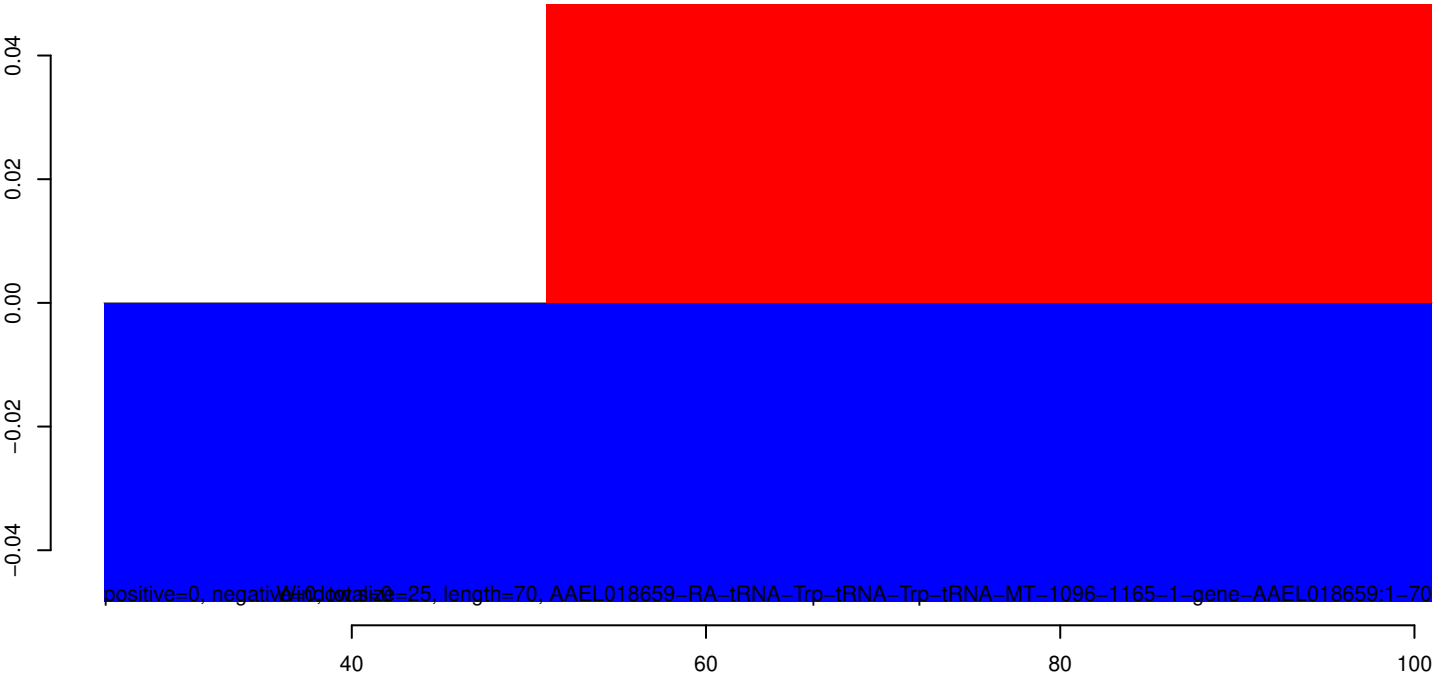
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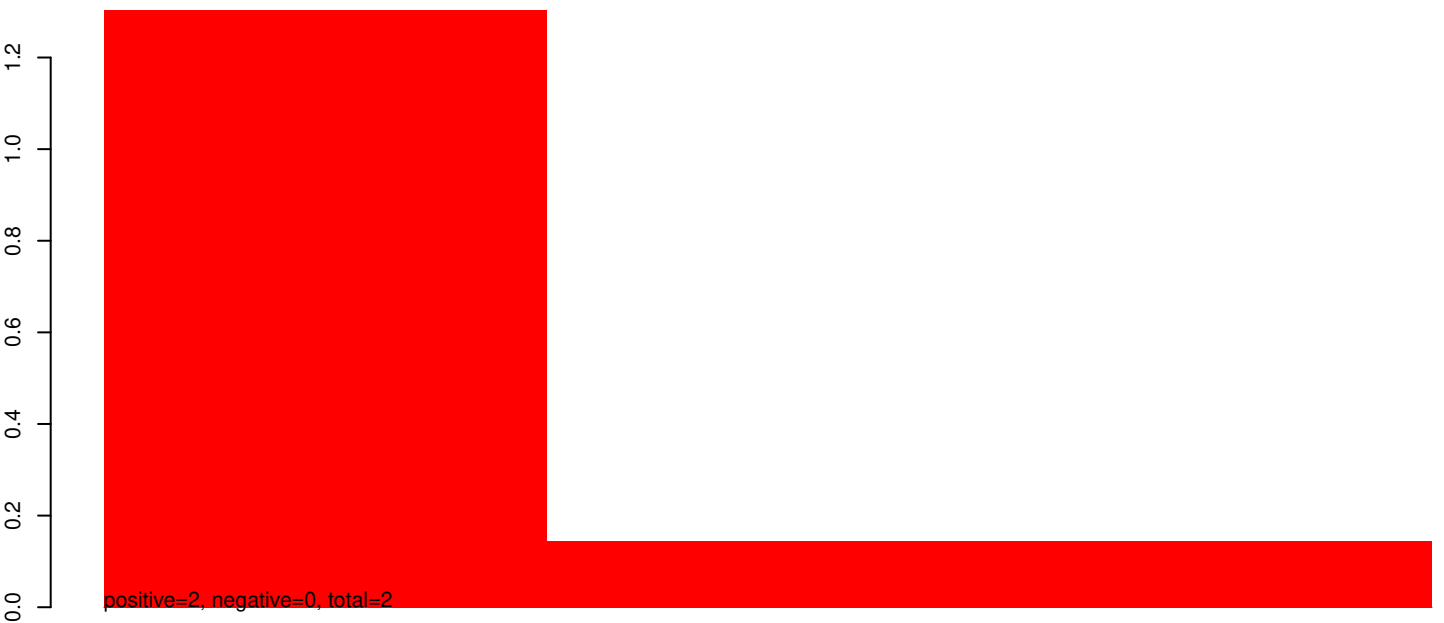
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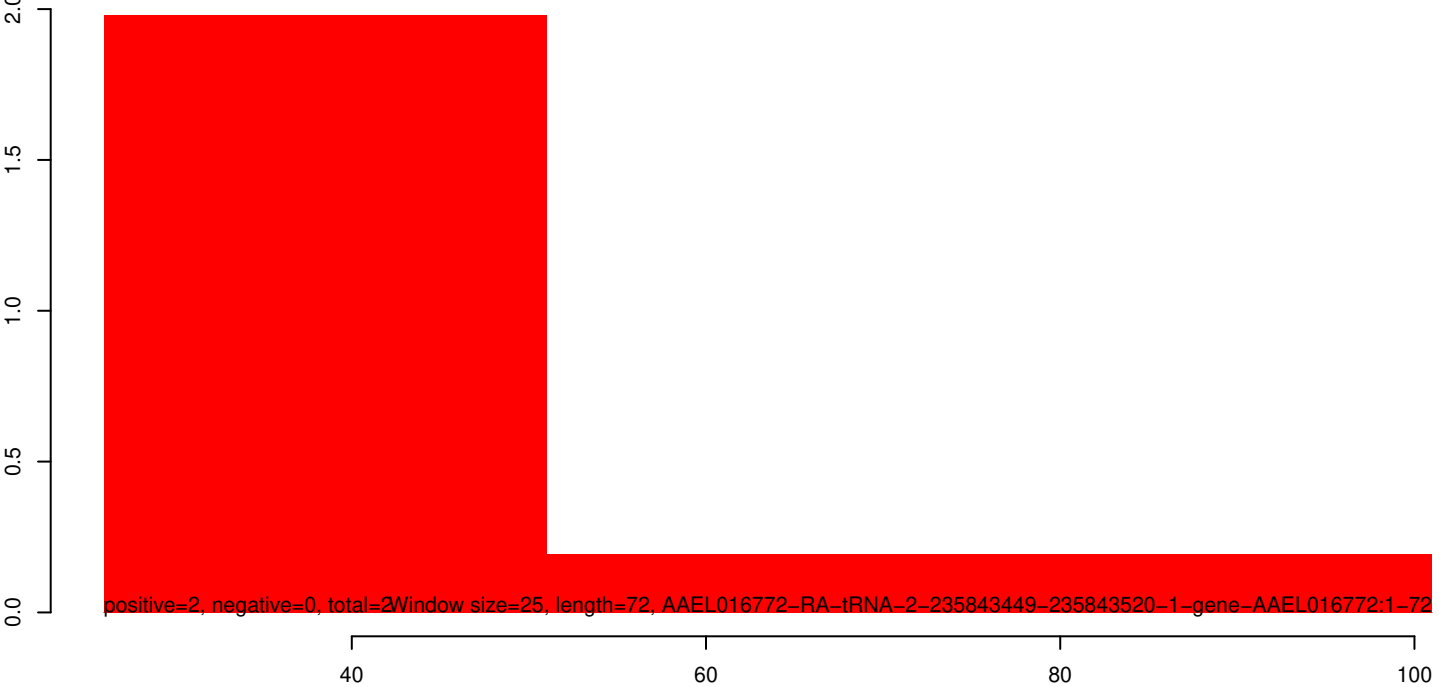
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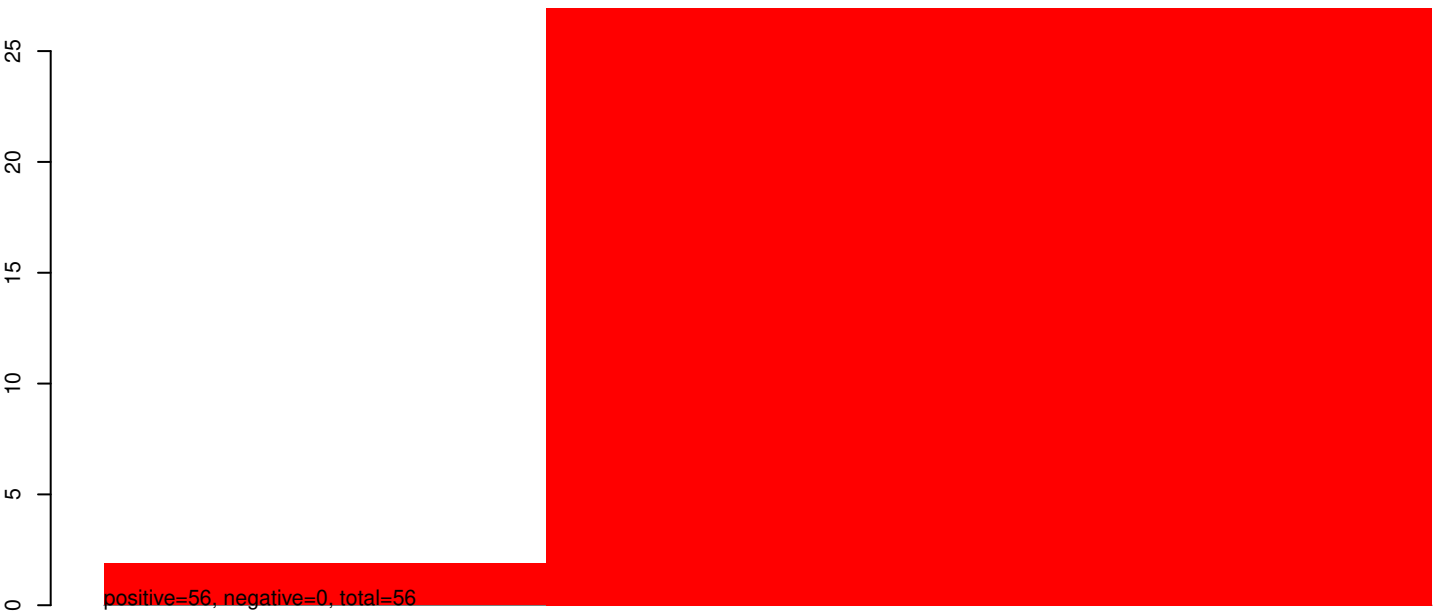
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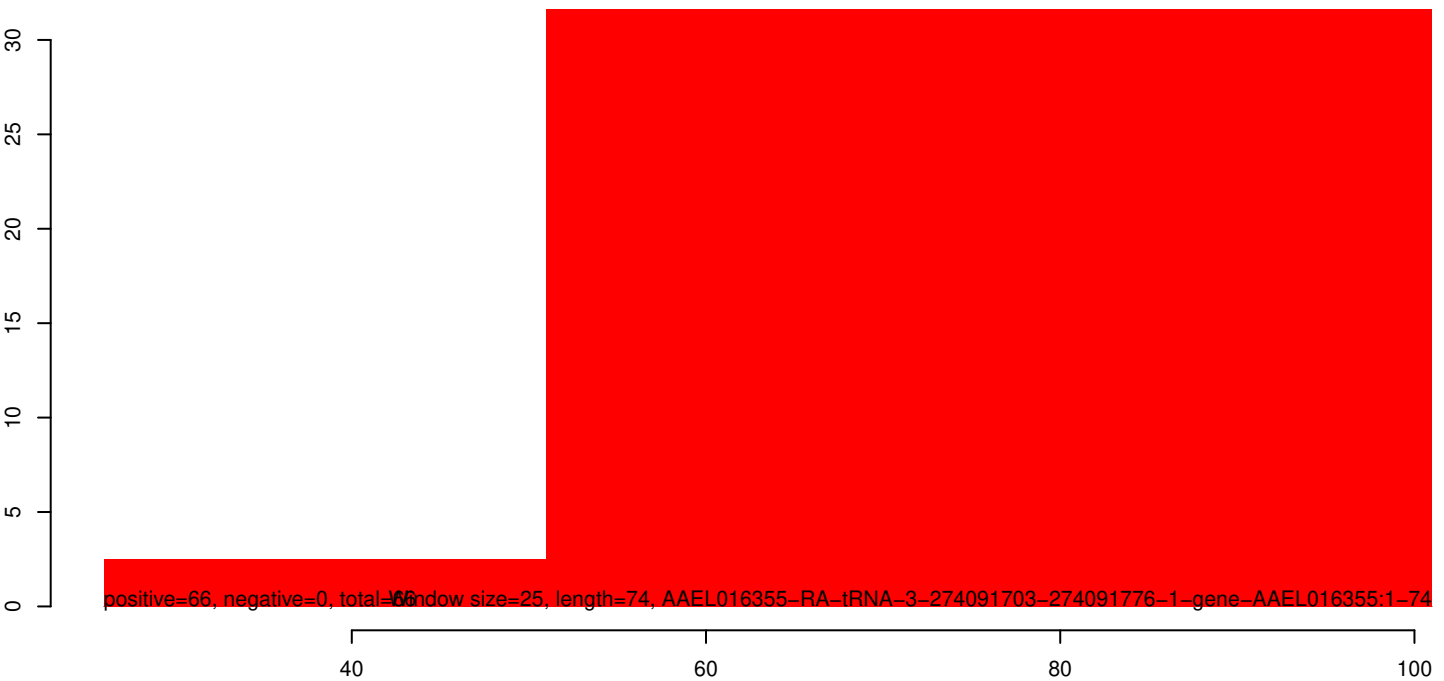
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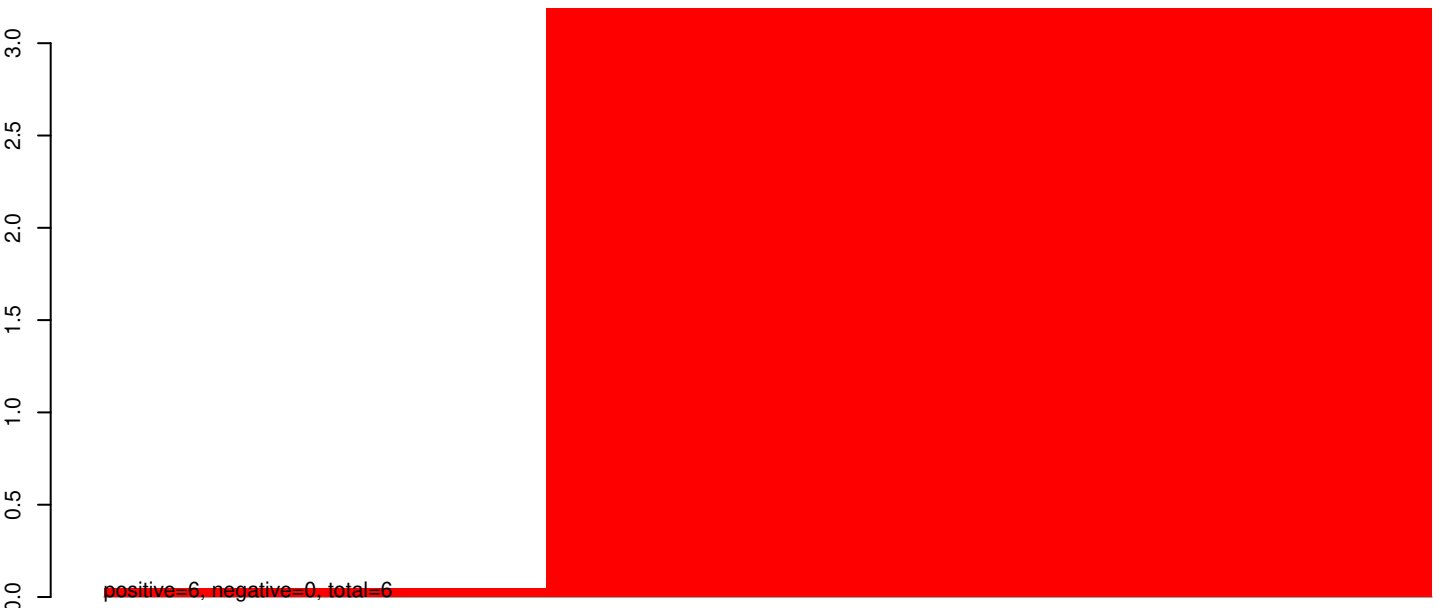
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AeAeg_Aag2SINV_dsPiwi5.24_35.rep



AeAeg_Aag2SINV_dsPiwi5.rep

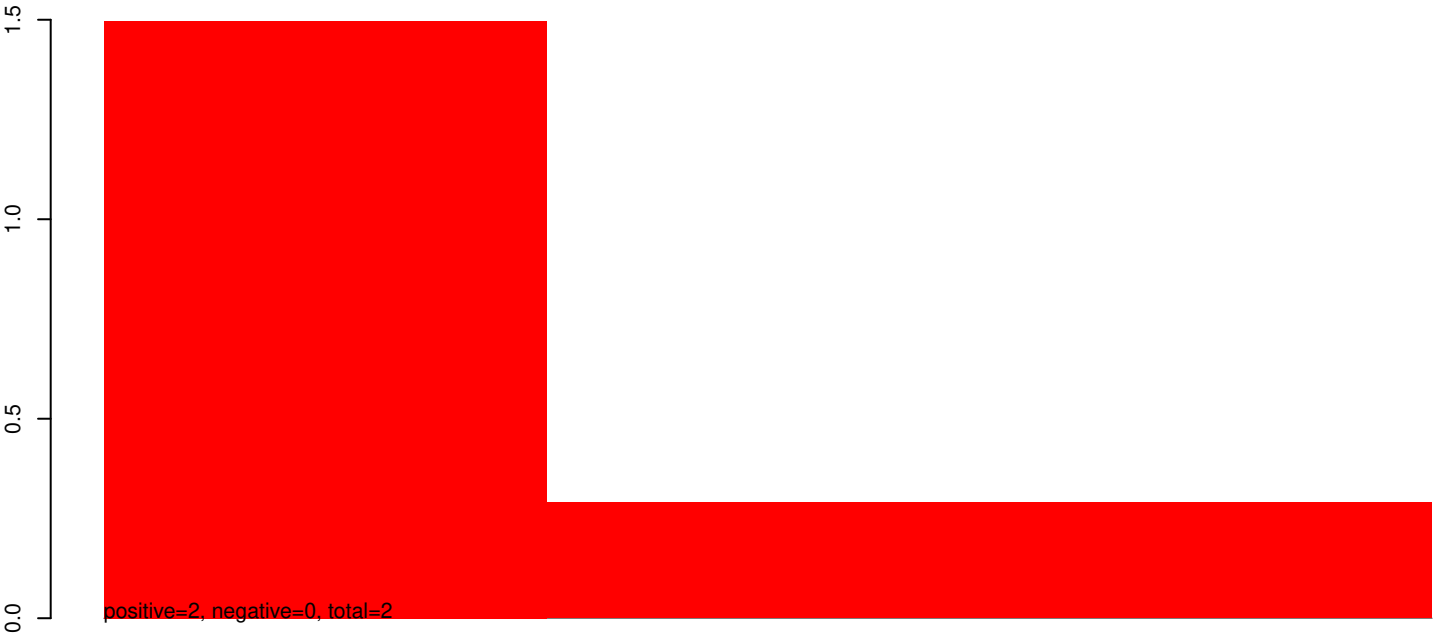


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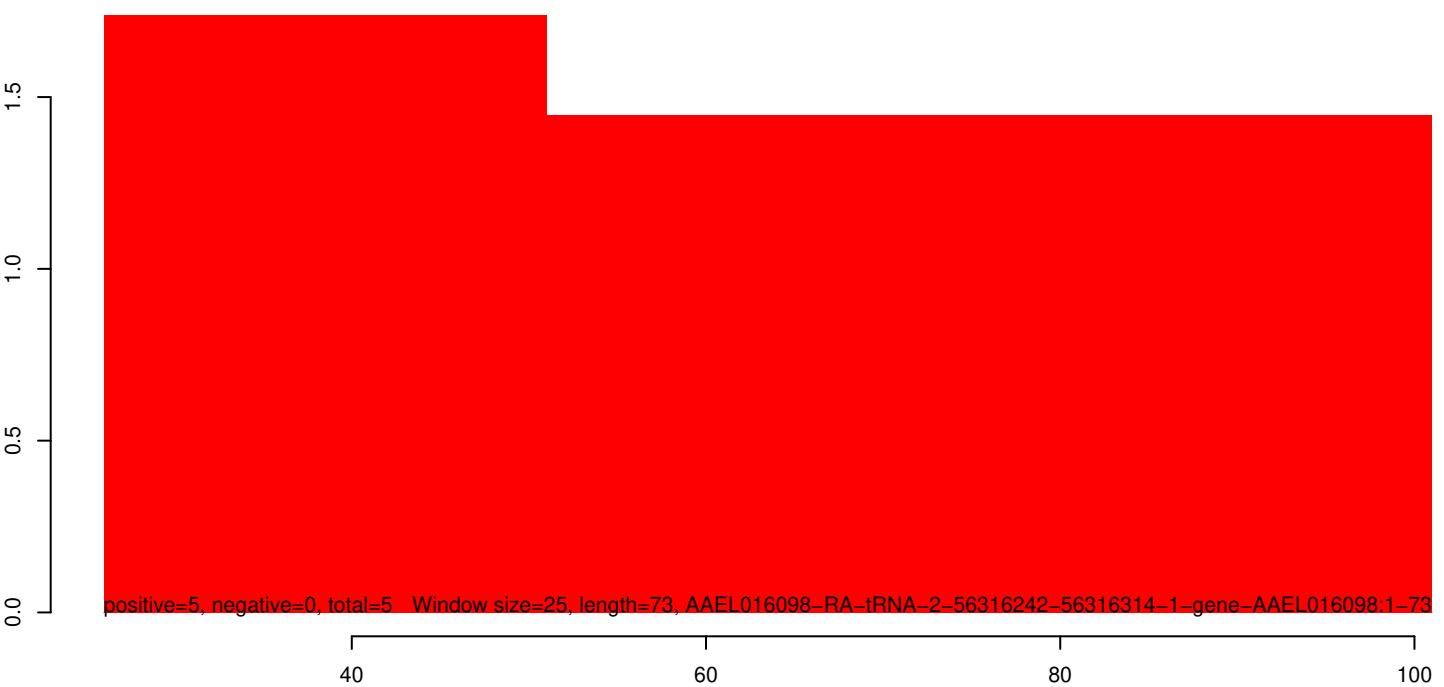
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AeAeg_Aag2SINV_dsPiwi5.24_35.rep



AeAeg_Aag2SINV_dsPiwi5.rep



AeAeg_Aag2SINV_dsPiwi5.18_23.rep

0.15
0.10
0.05
0.00

positive=0, negative=0, total=0

AeAeg_Aag2SINV_dsPiwi5.24_35.rep

0.15
0.10
0.05
0.00

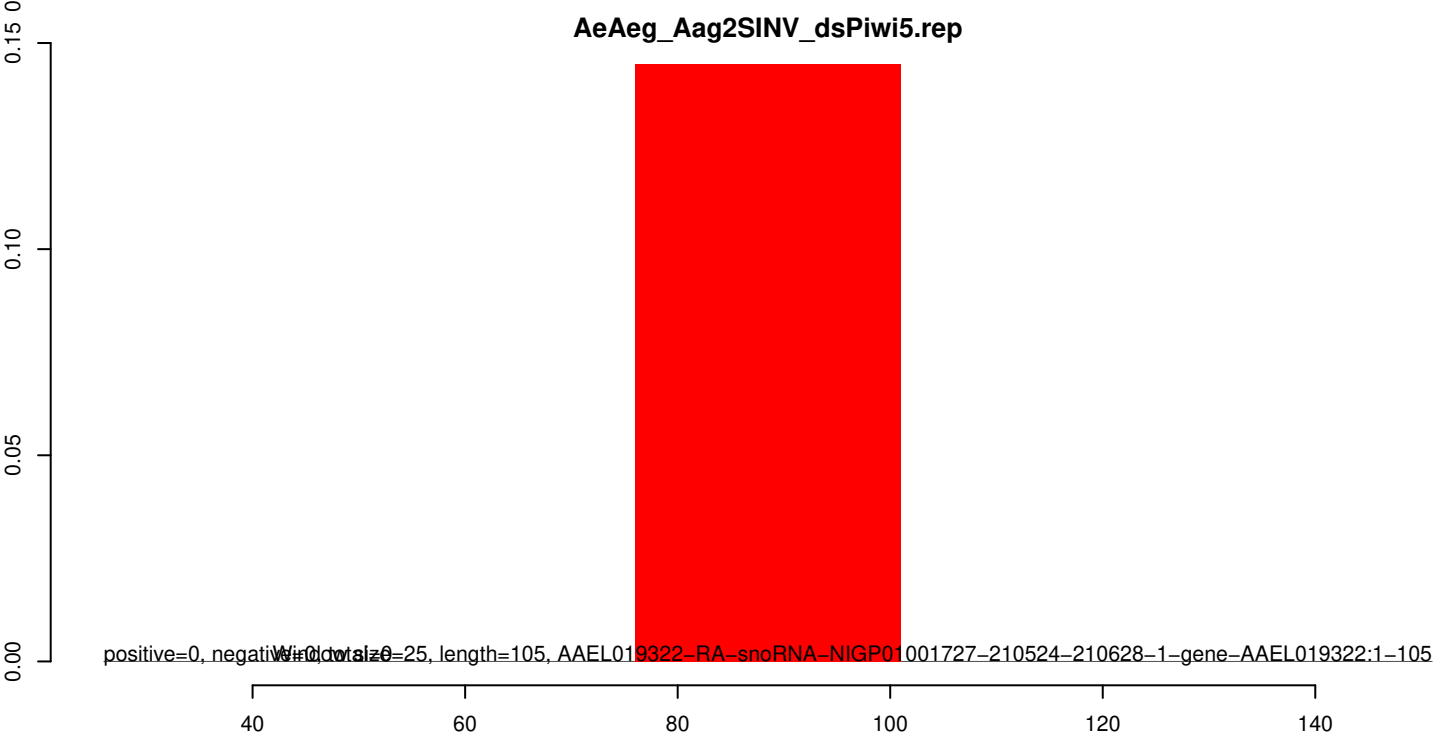
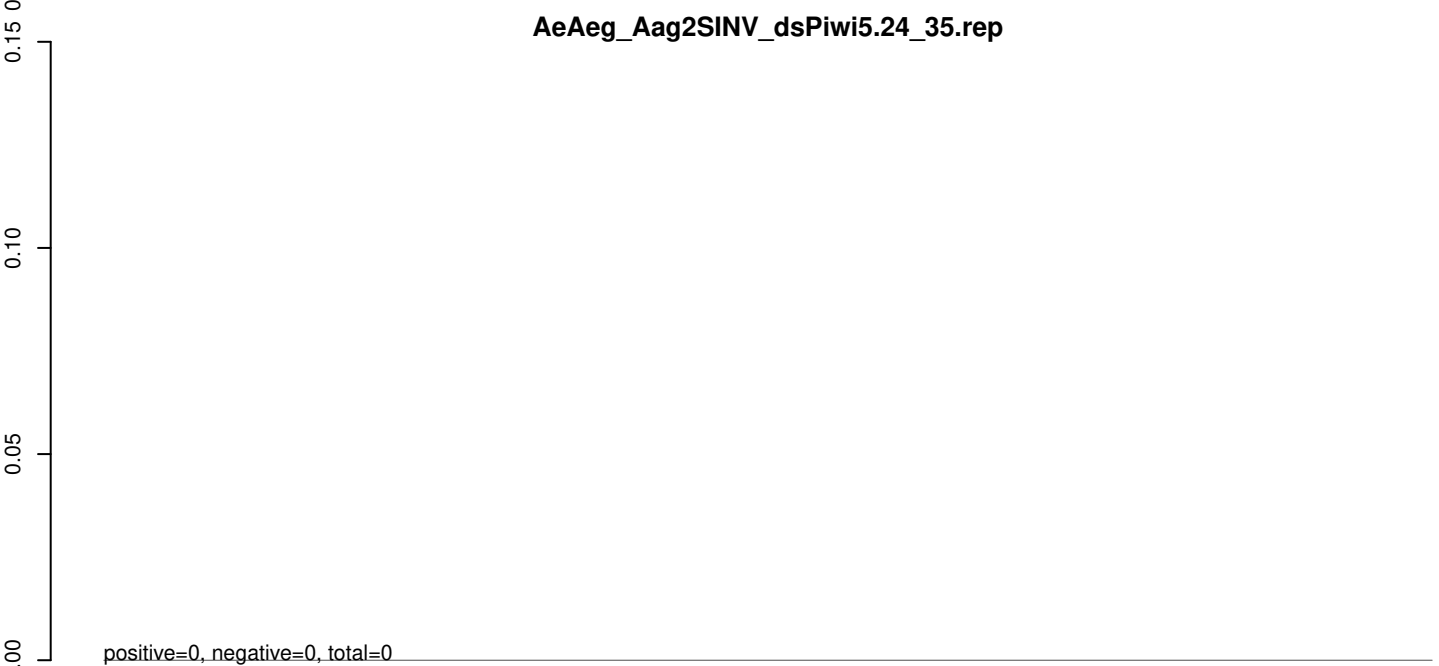
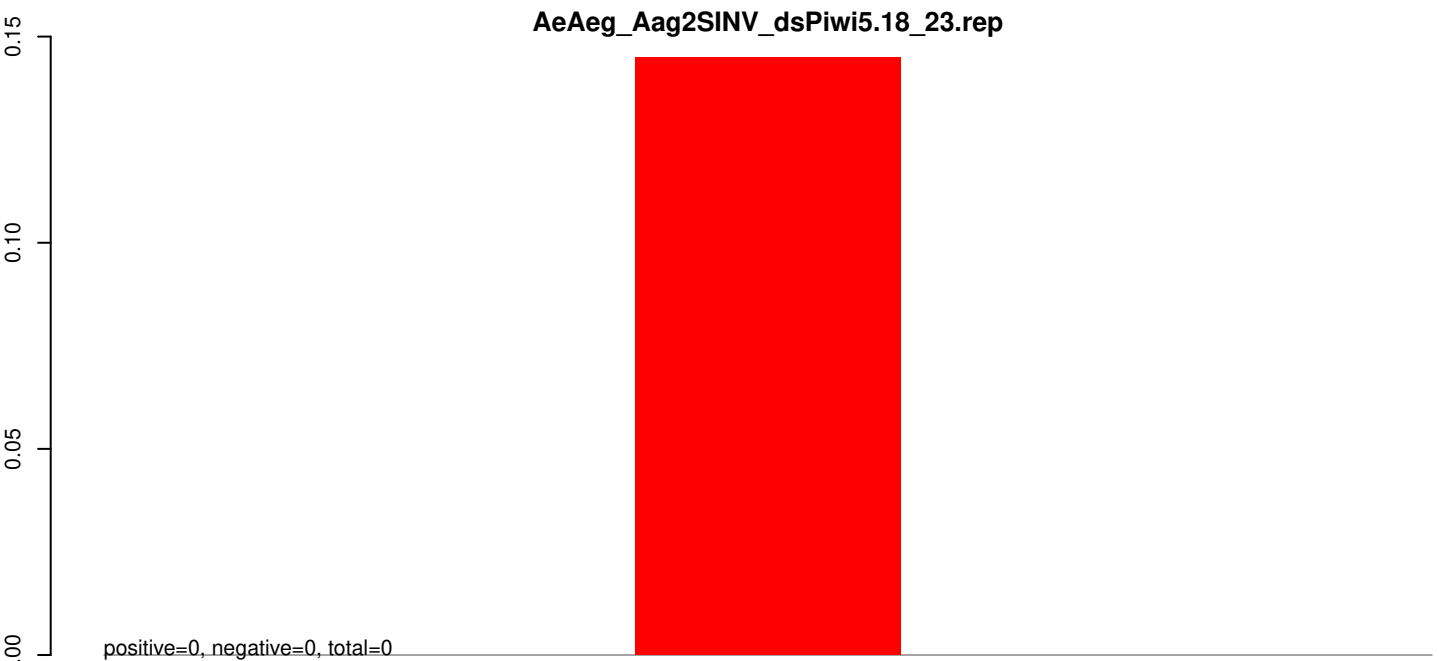
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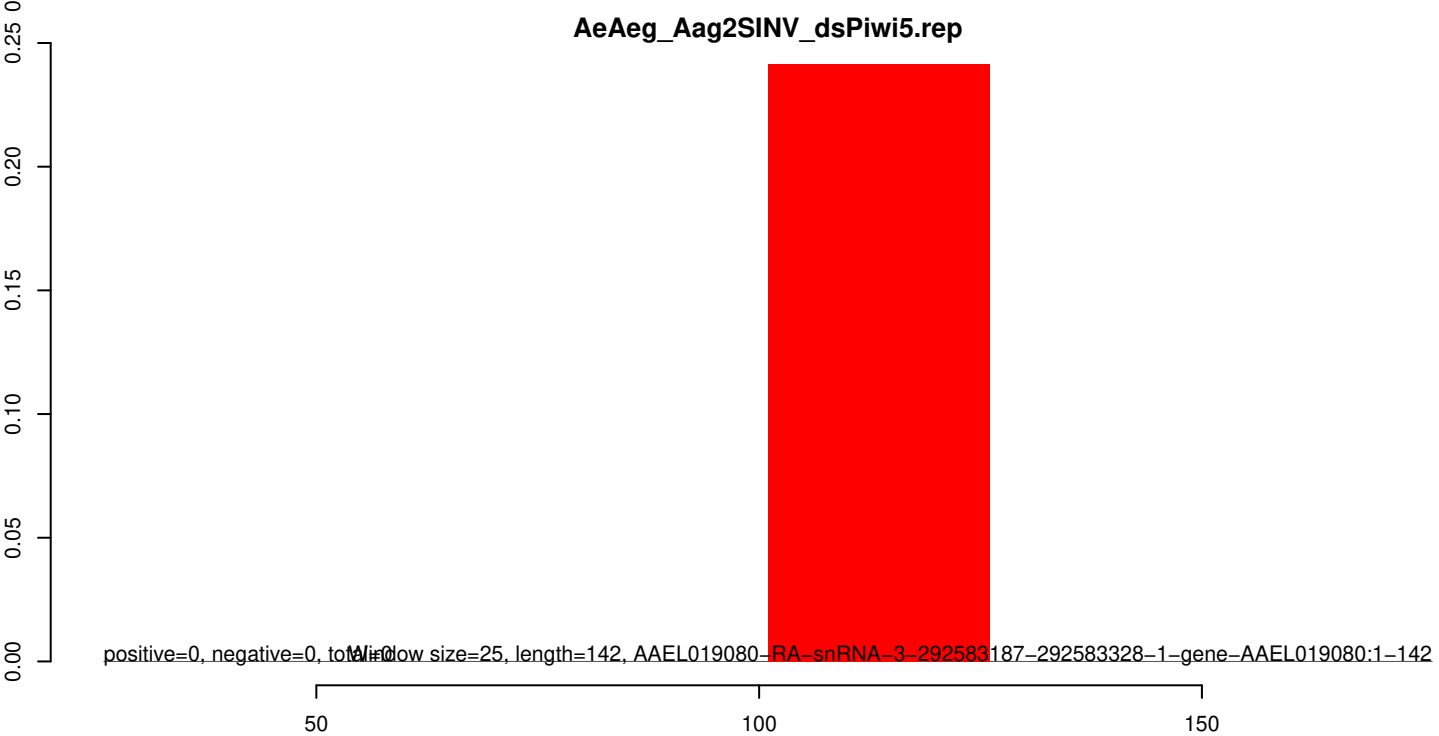
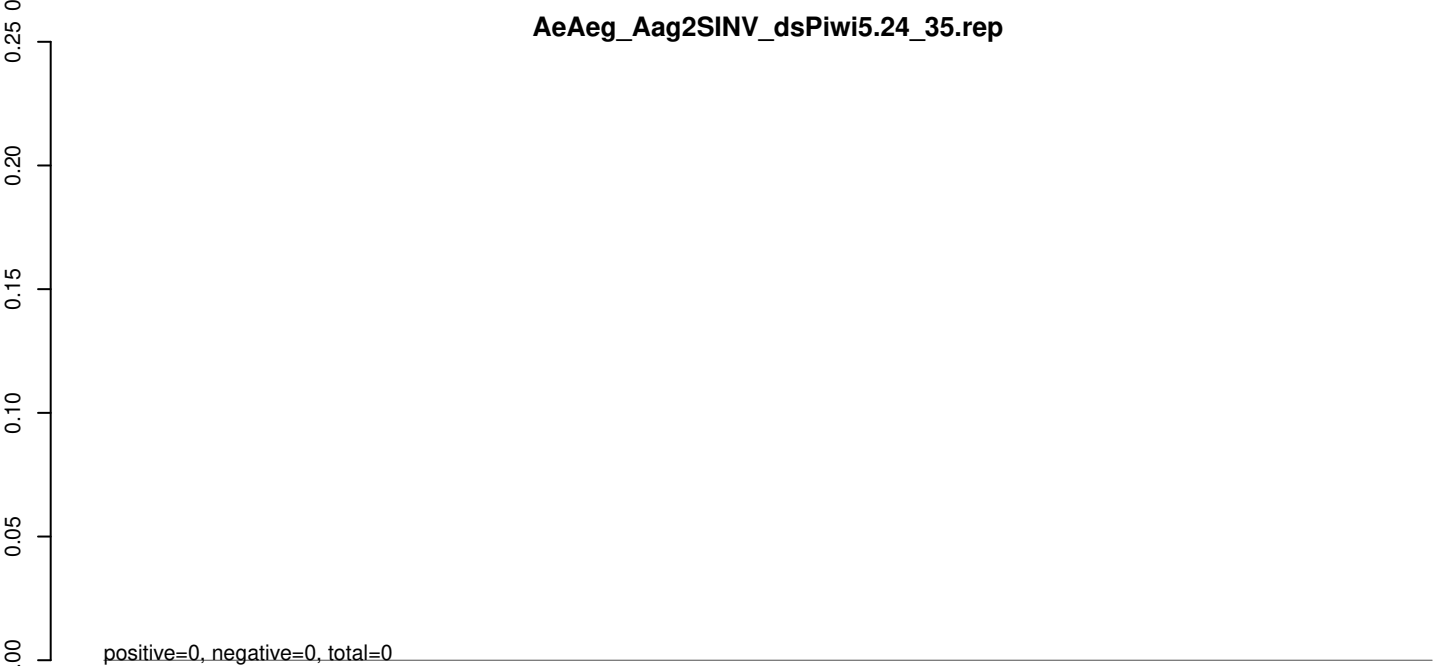
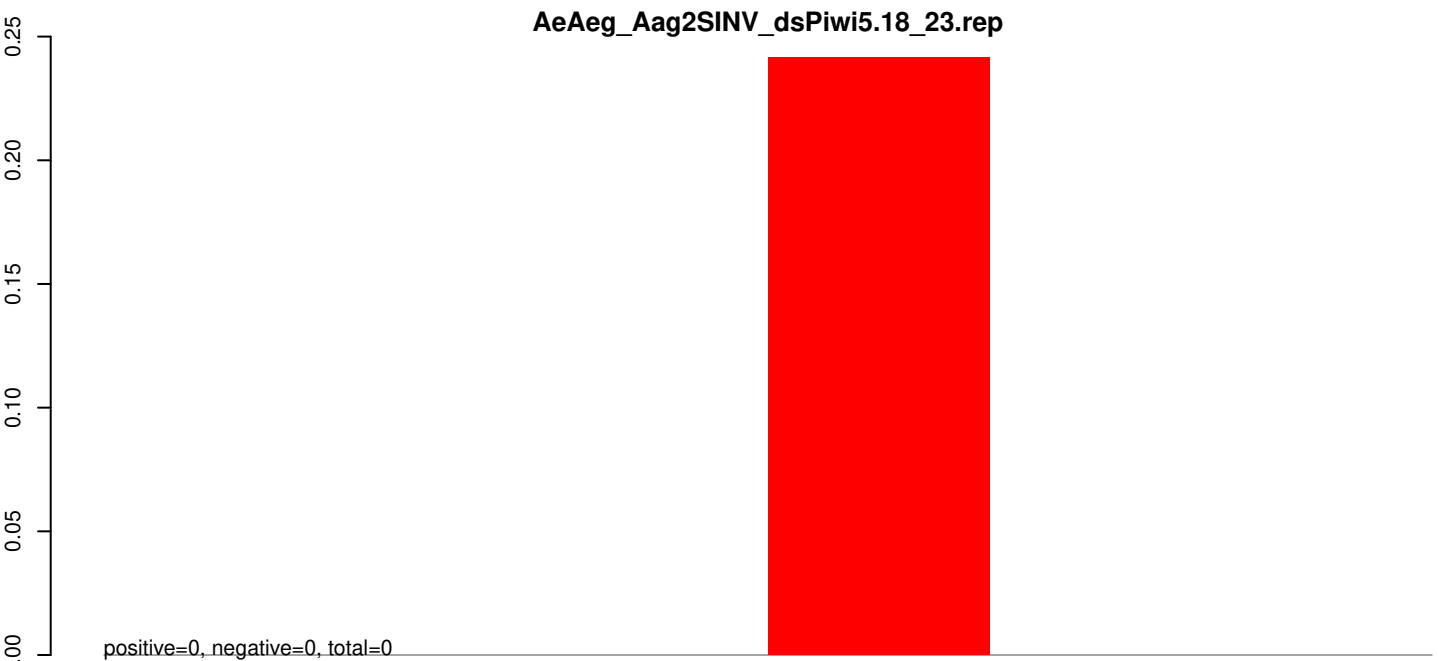
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0.15
0.10
0.05
0.00

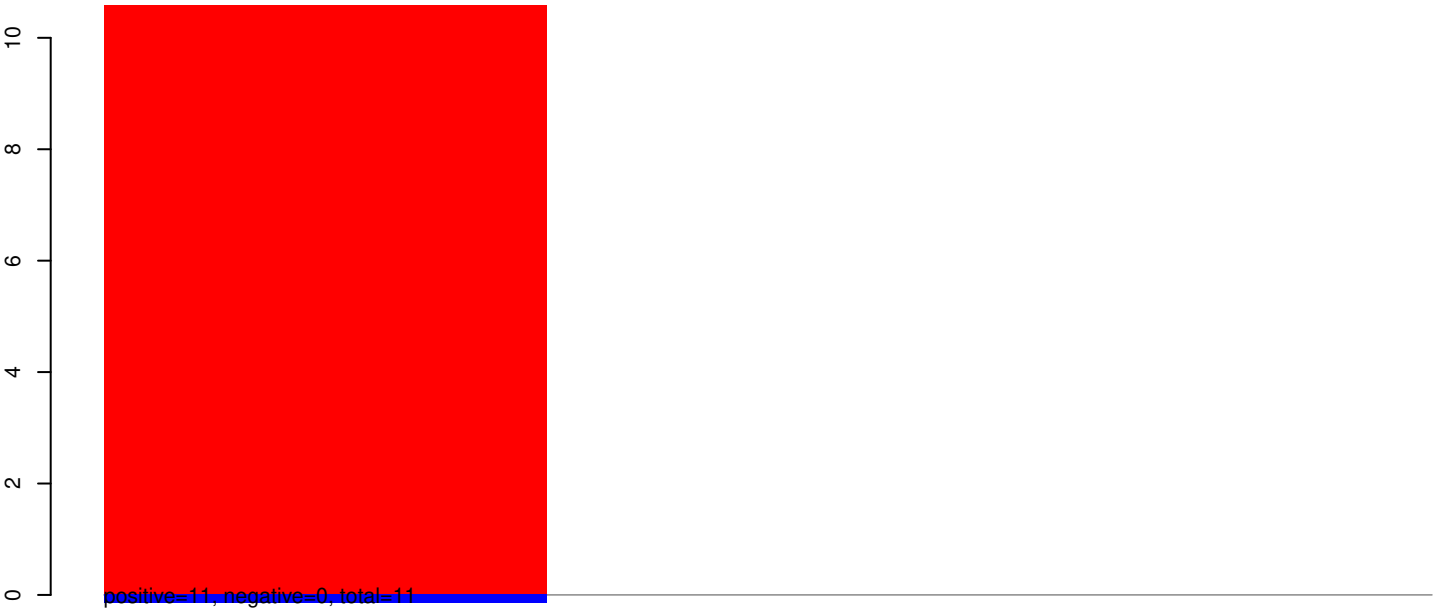
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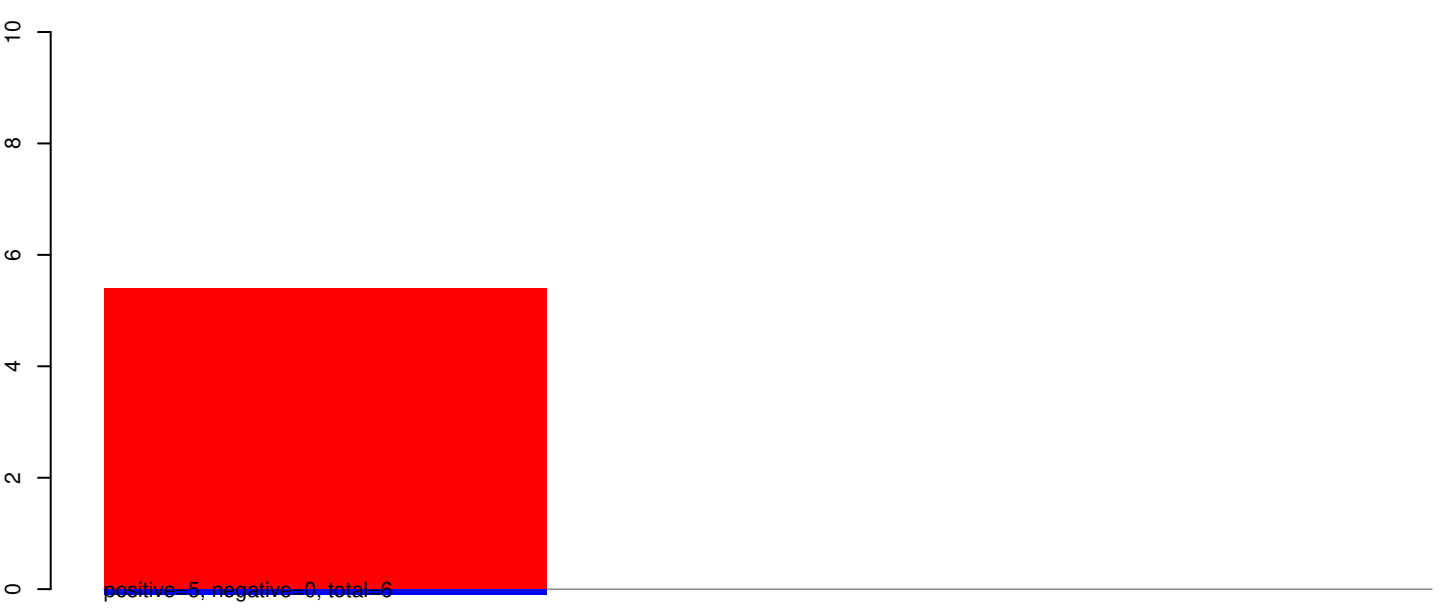




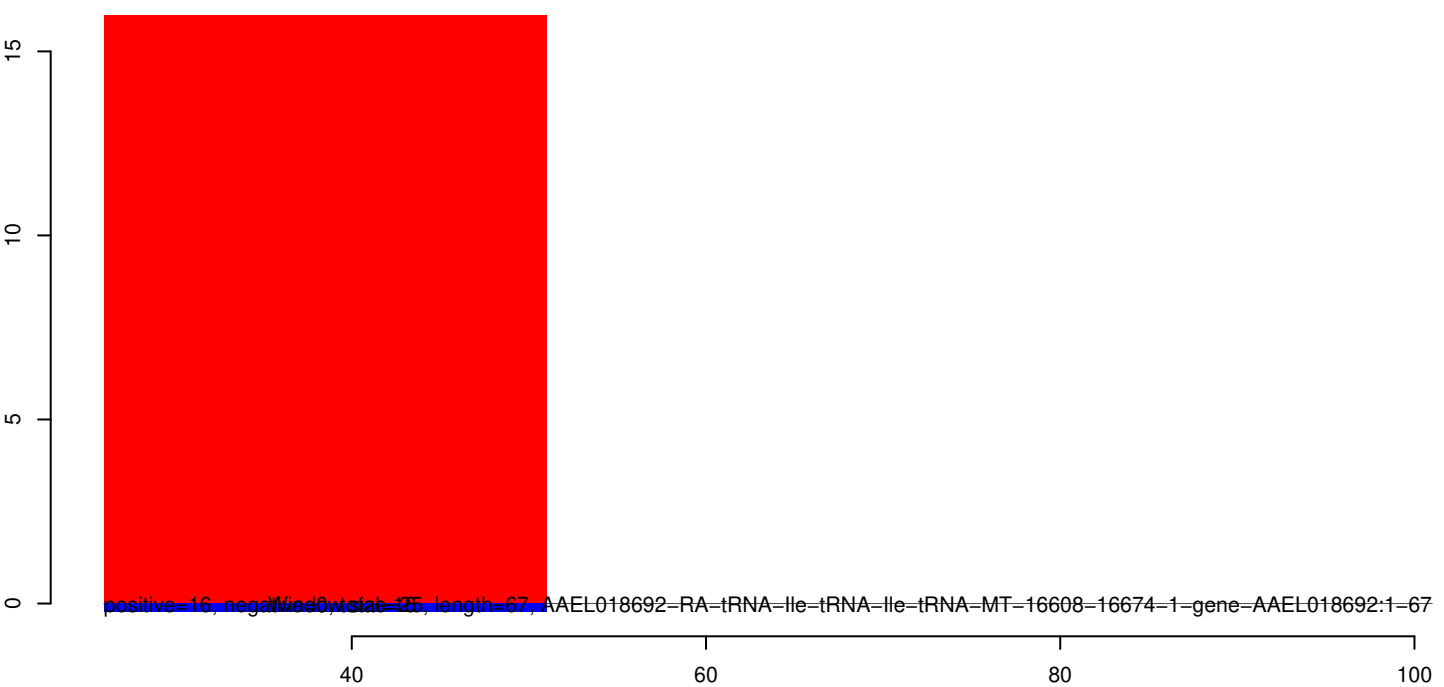
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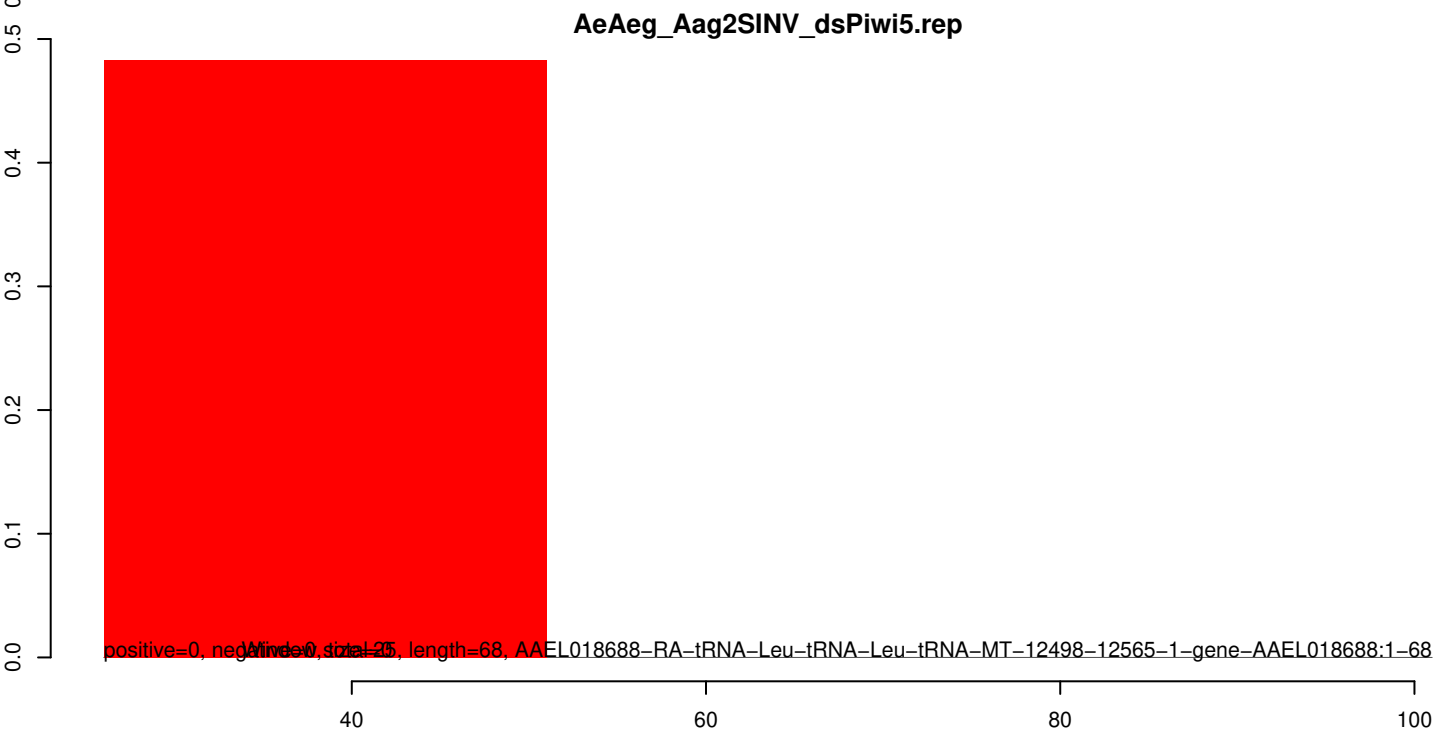
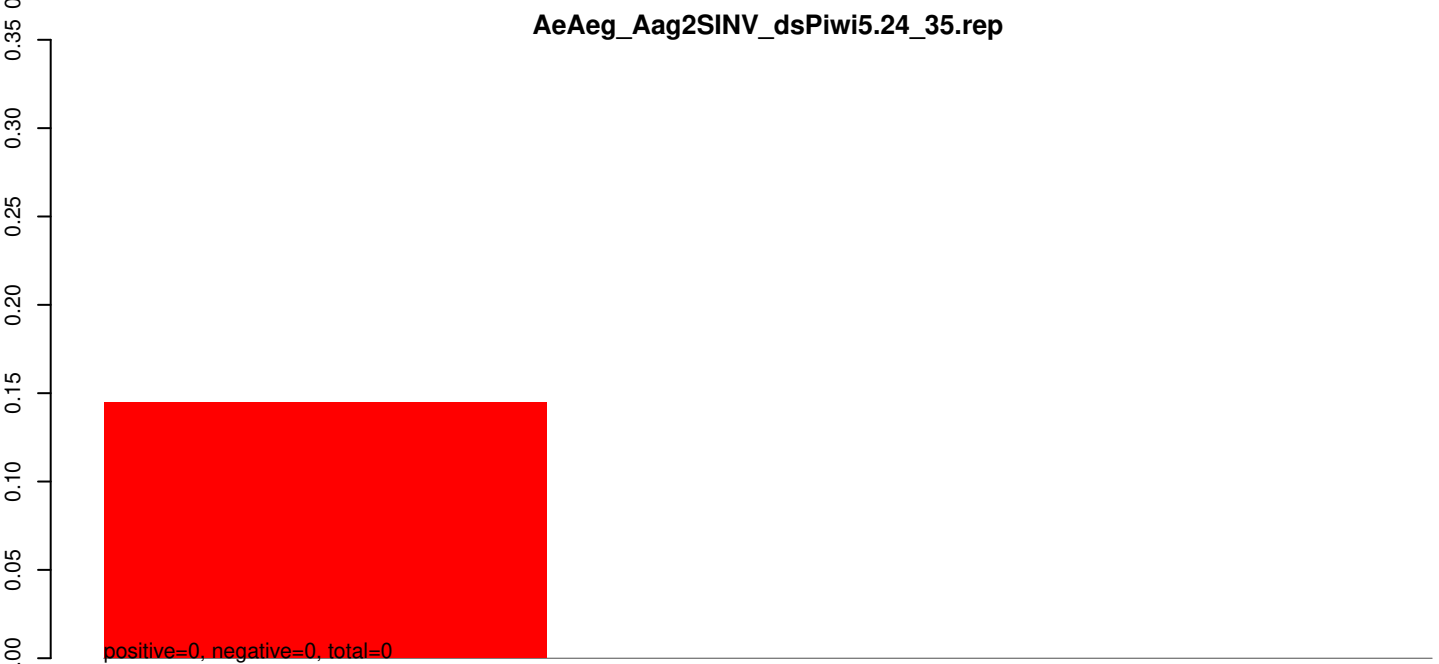
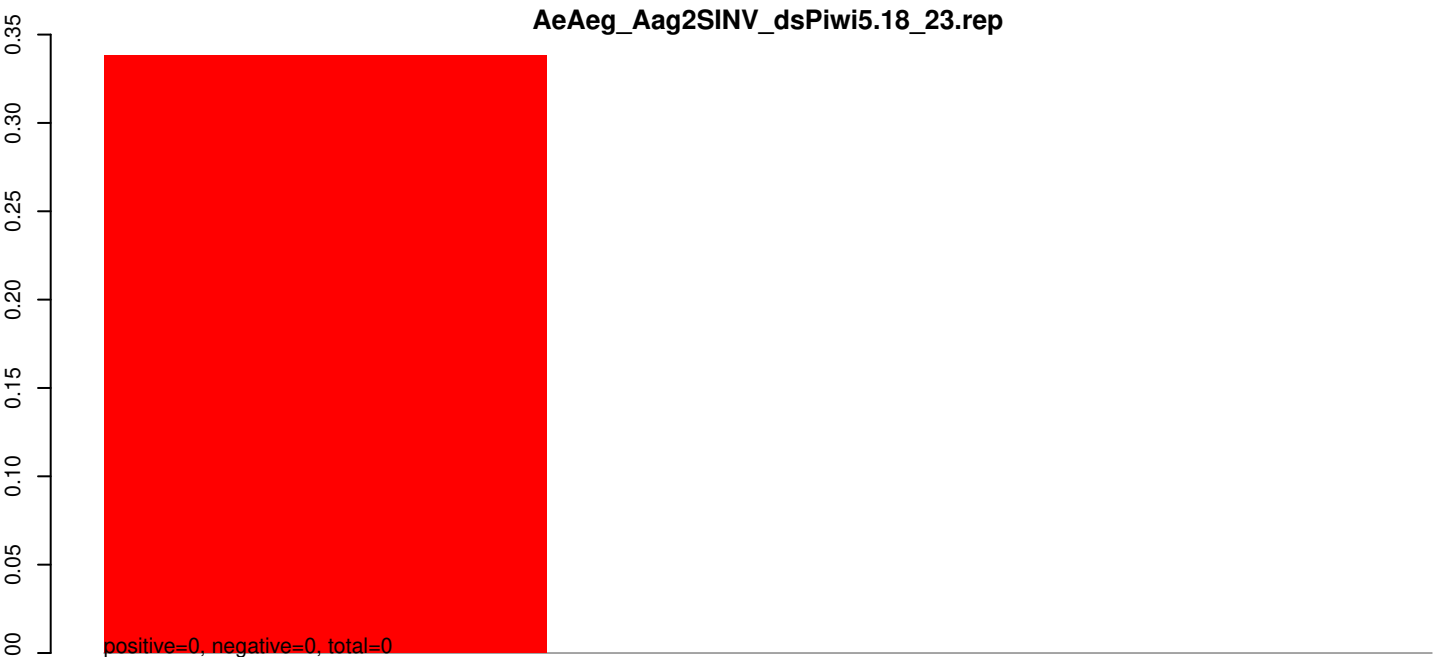


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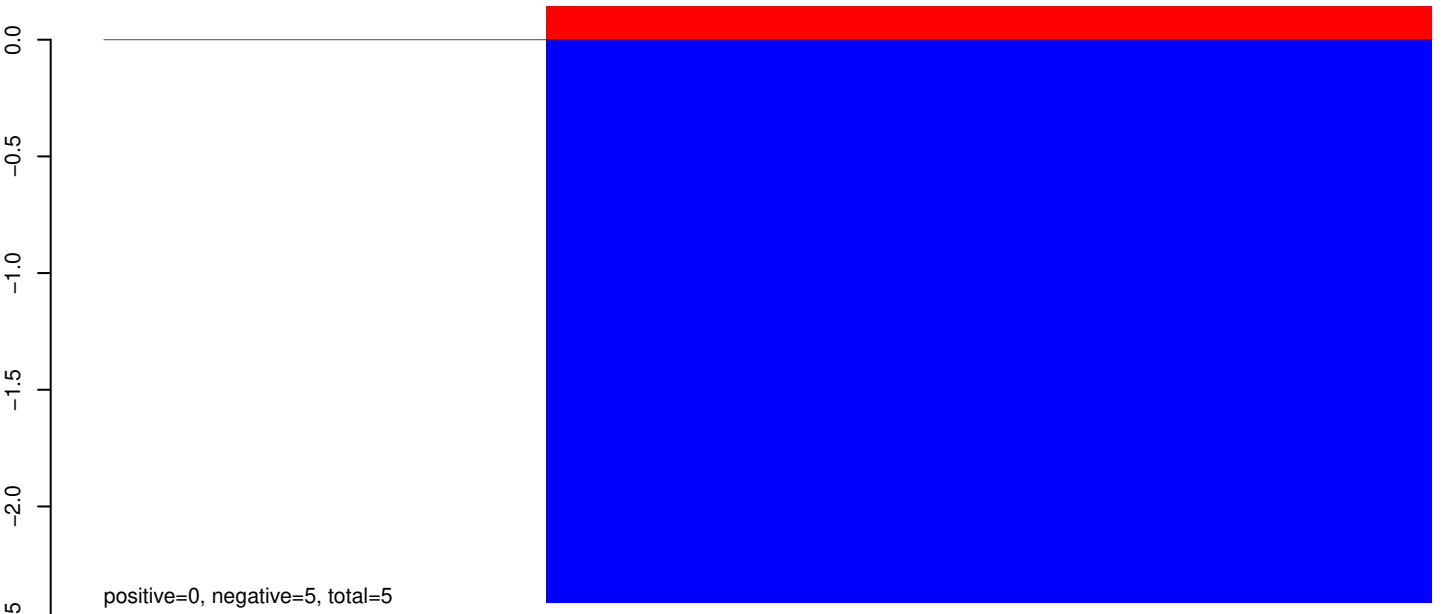


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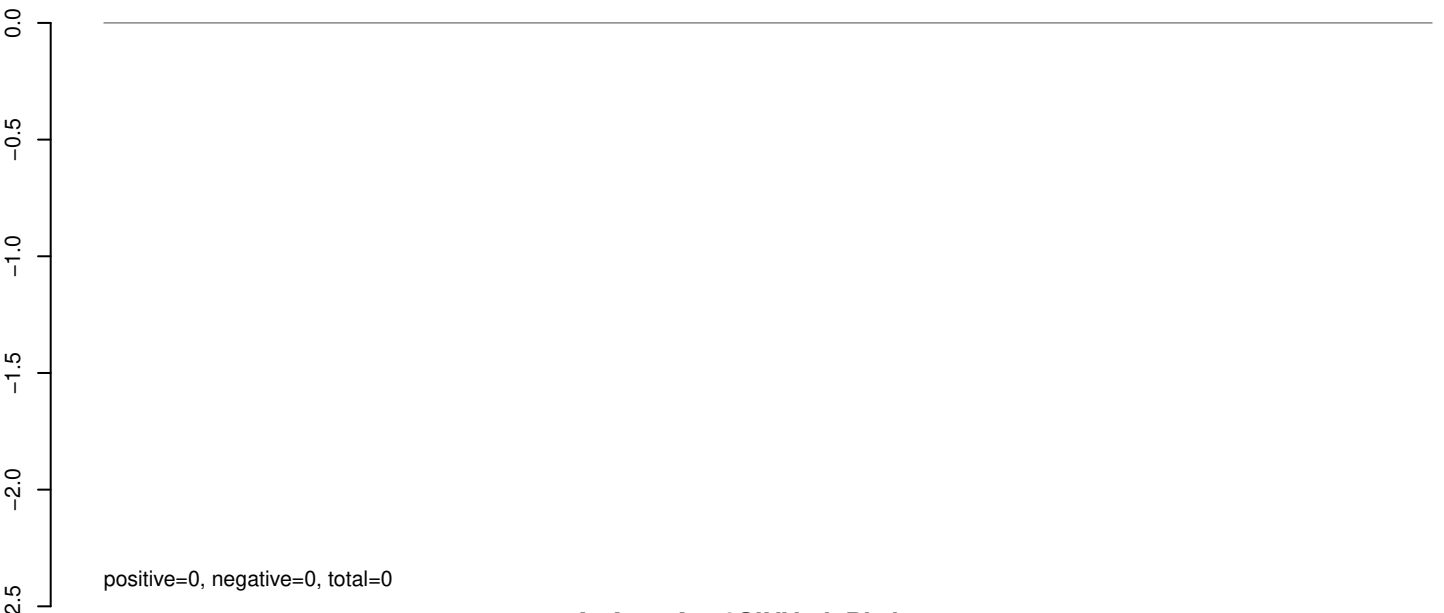


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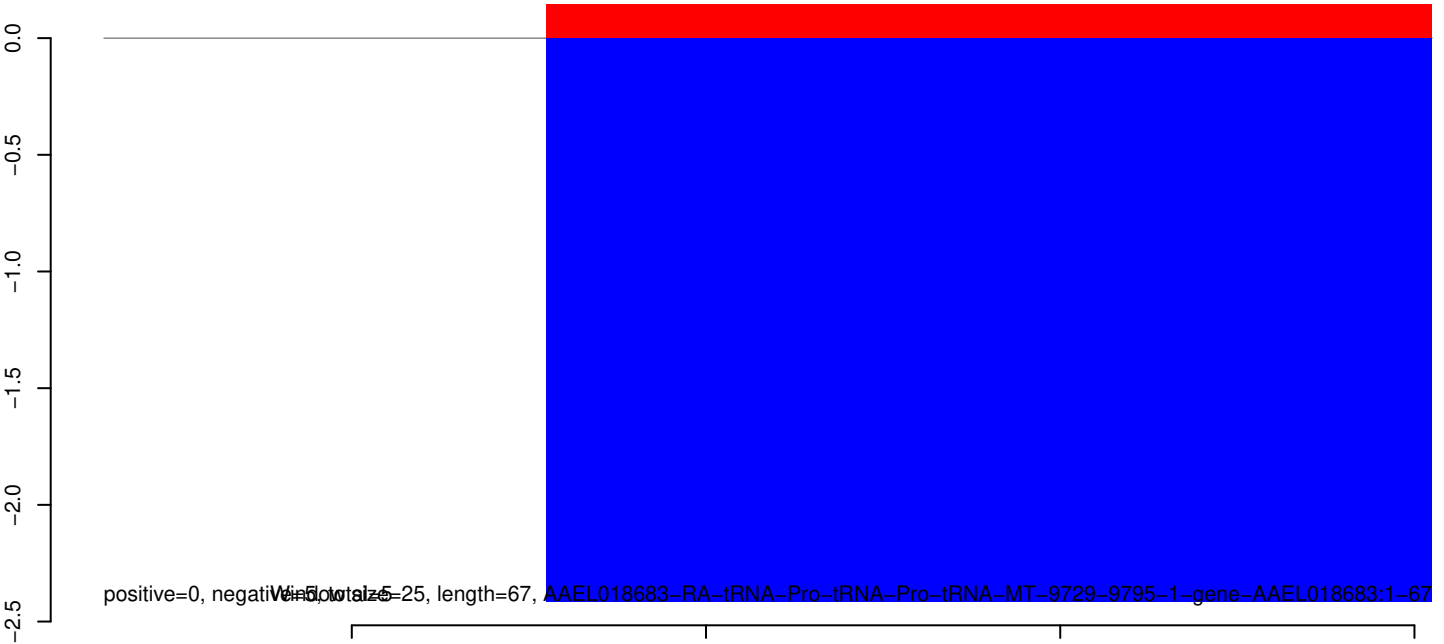
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positive=0, negative=0, total=0

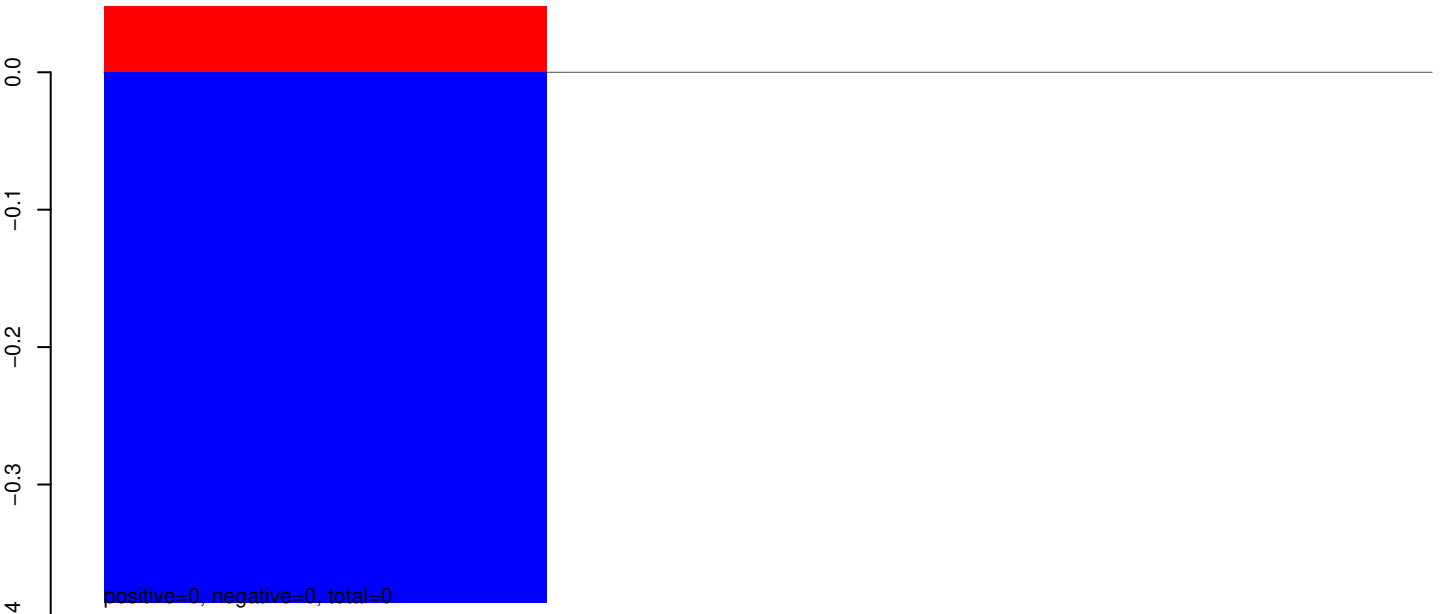
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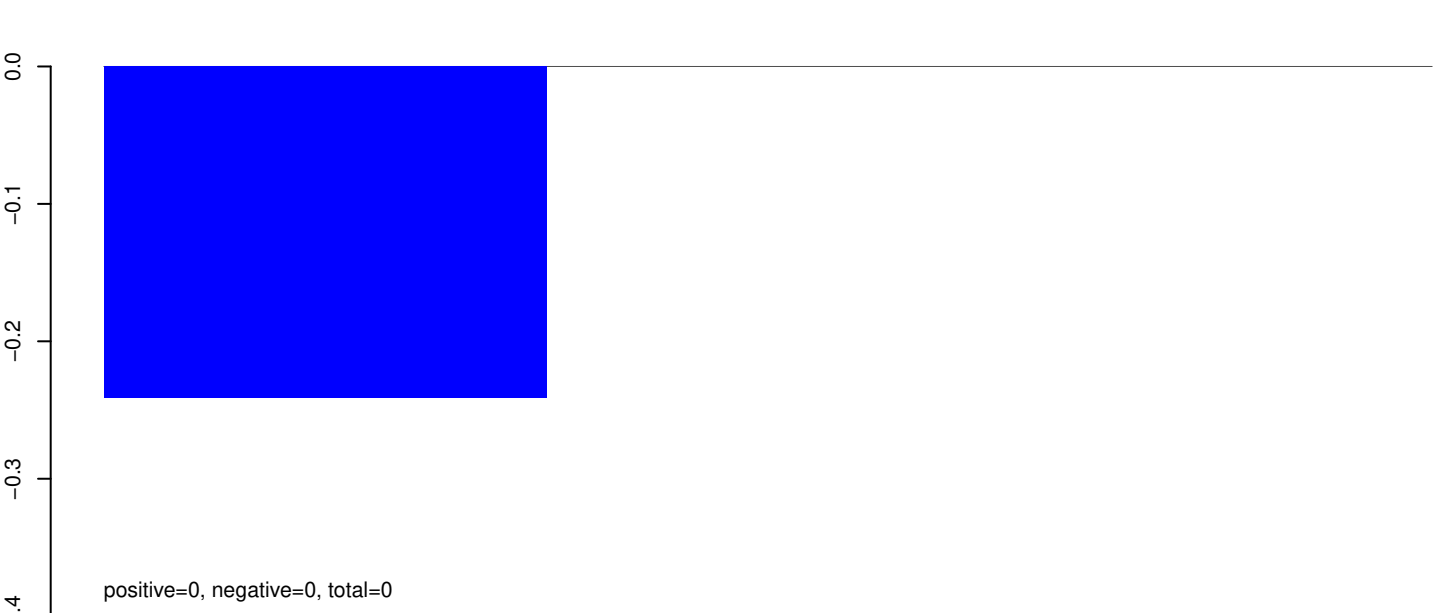
positive=0, negative=5, total=5, length=67, AAEL018683-RA-tRNA-Pro-tRNA-Pro-tRNA-MT-9729-9795-1-gene-AAEL018683:1-67

40 60 80 100

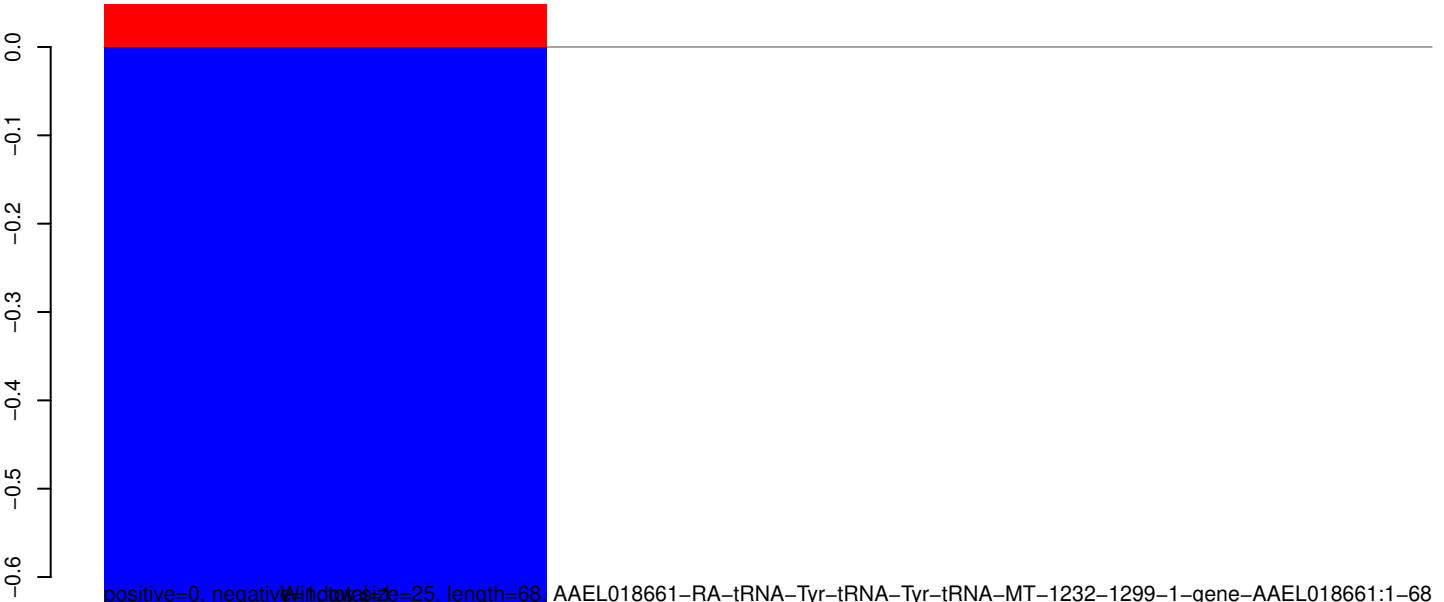
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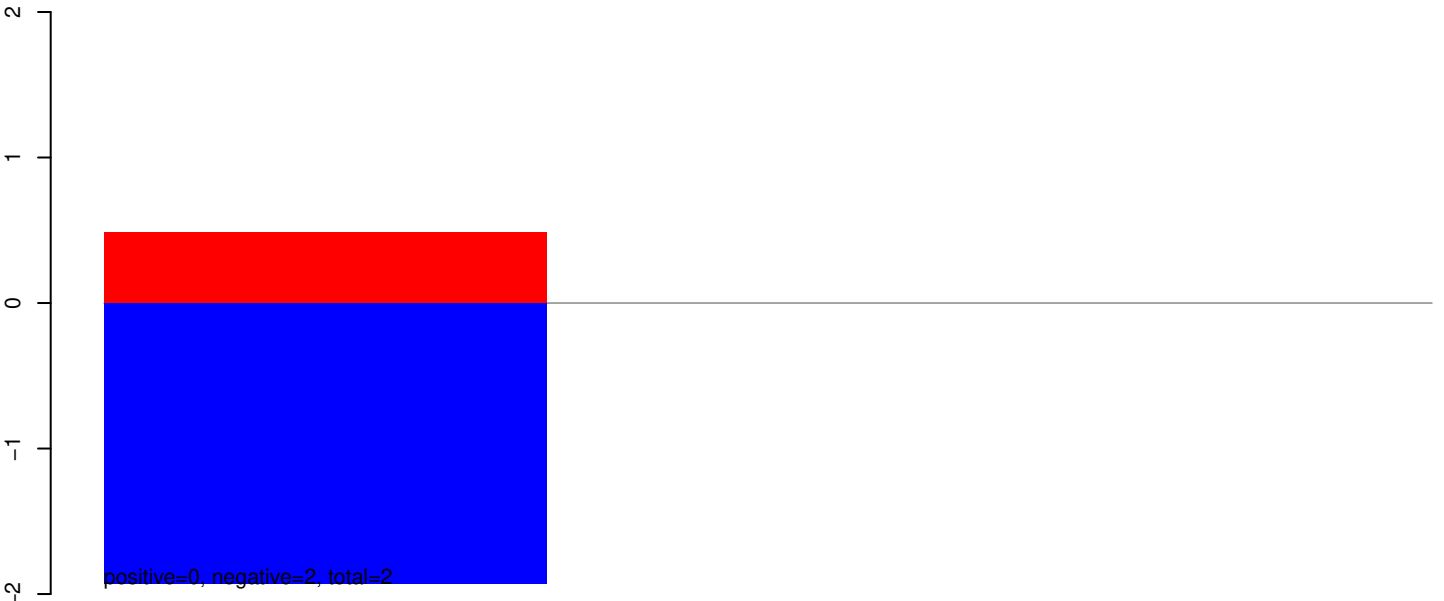


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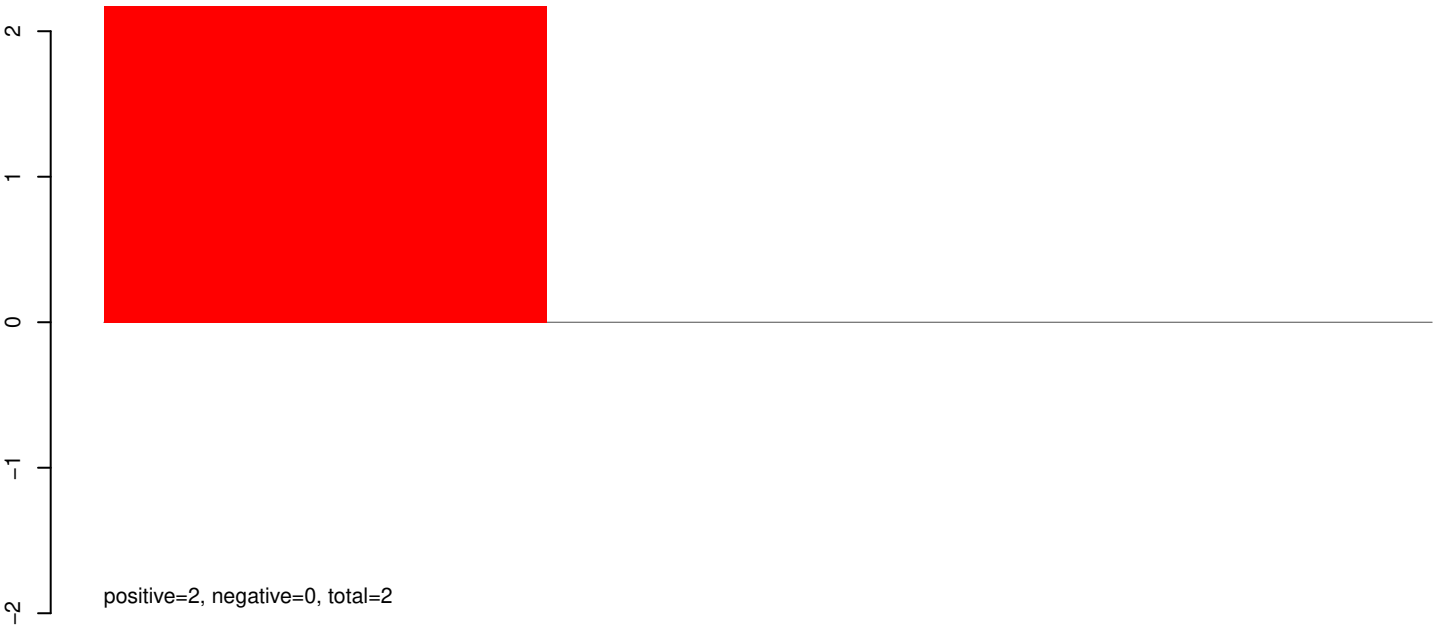


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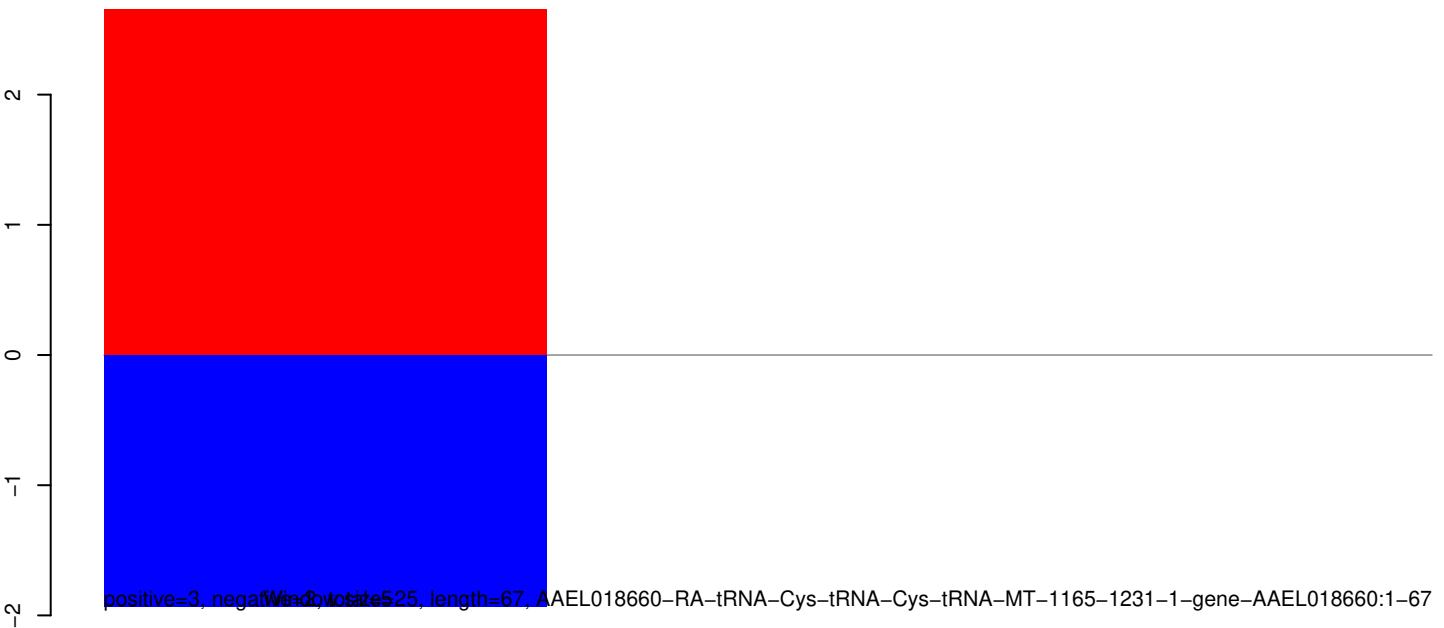
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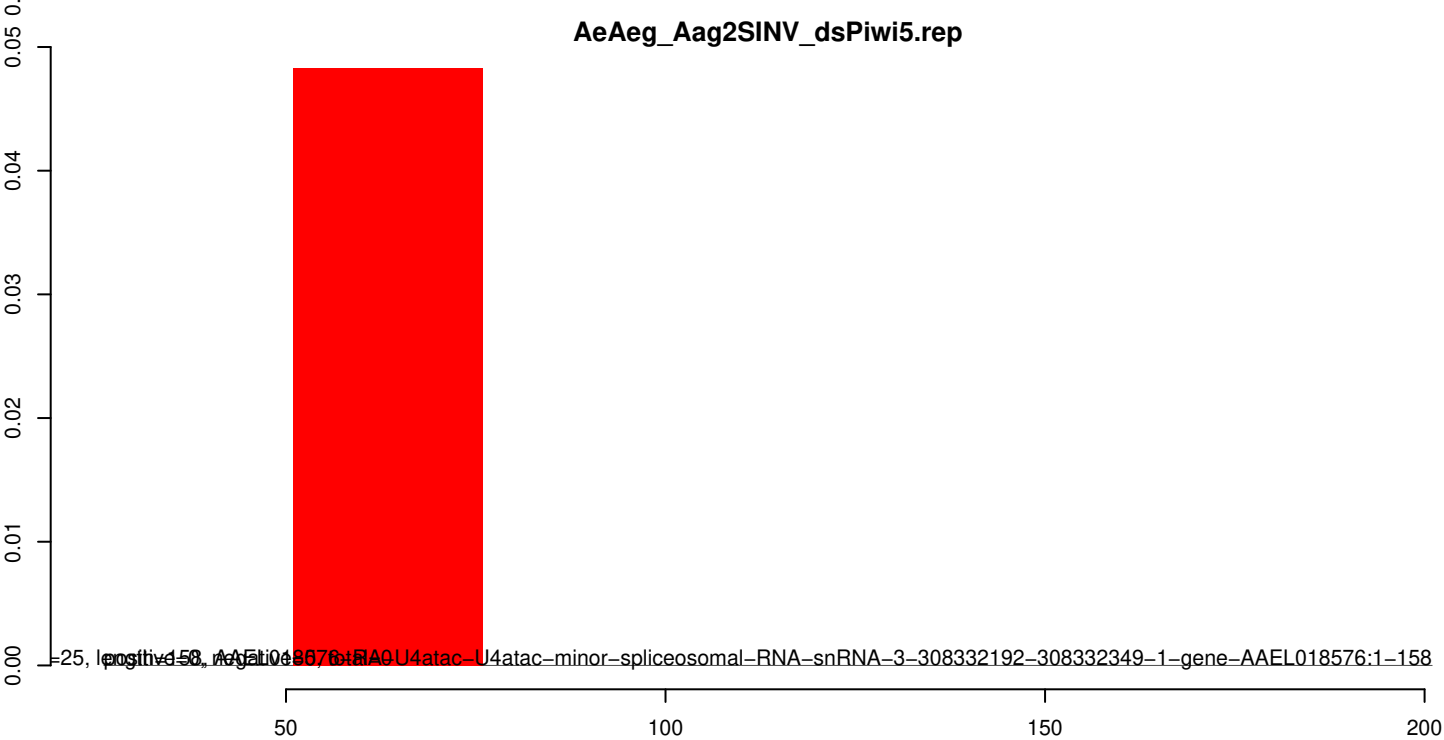
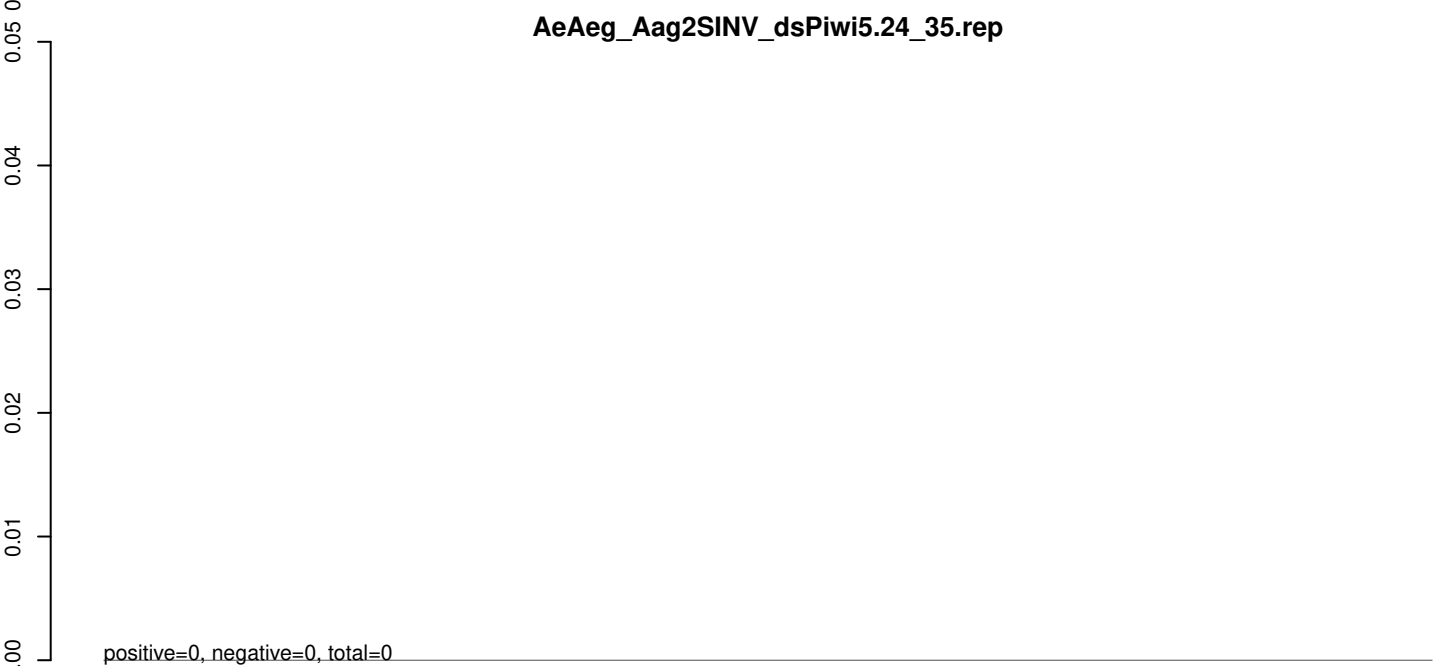
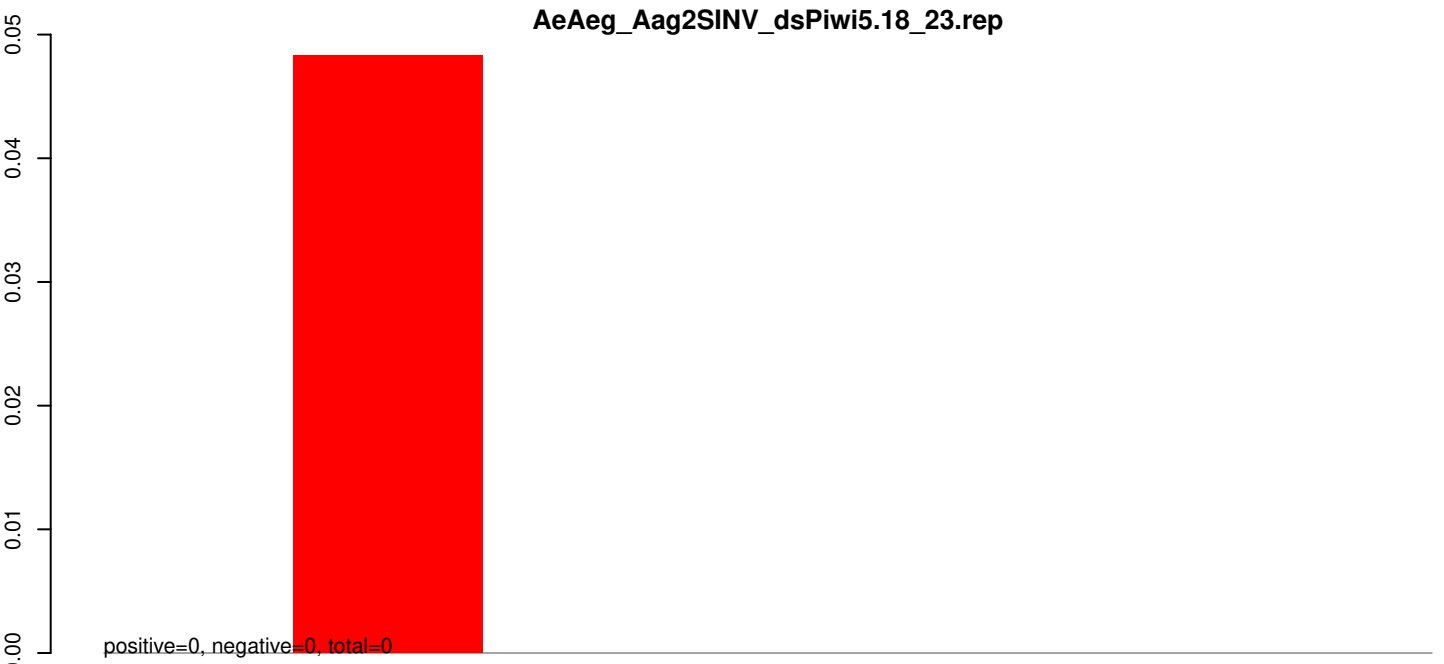


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AeAeg_Aag2SINV_dsPiwi5.rep





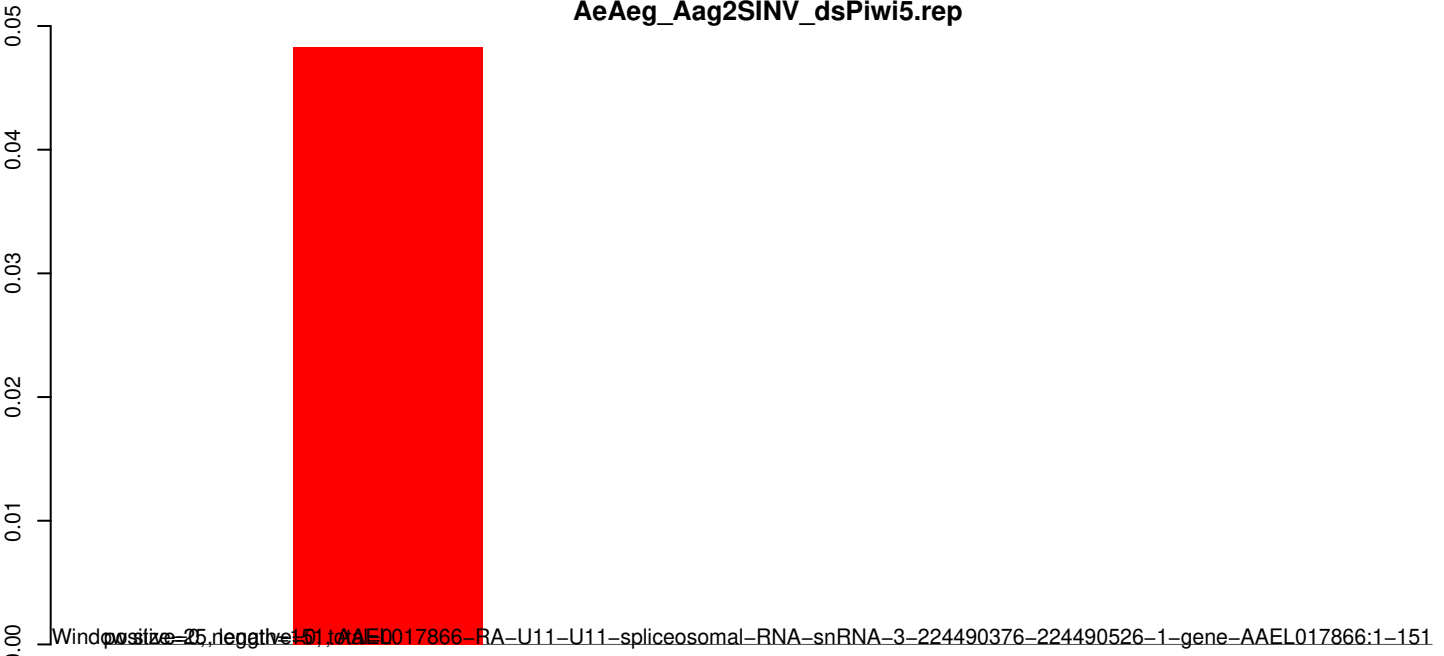
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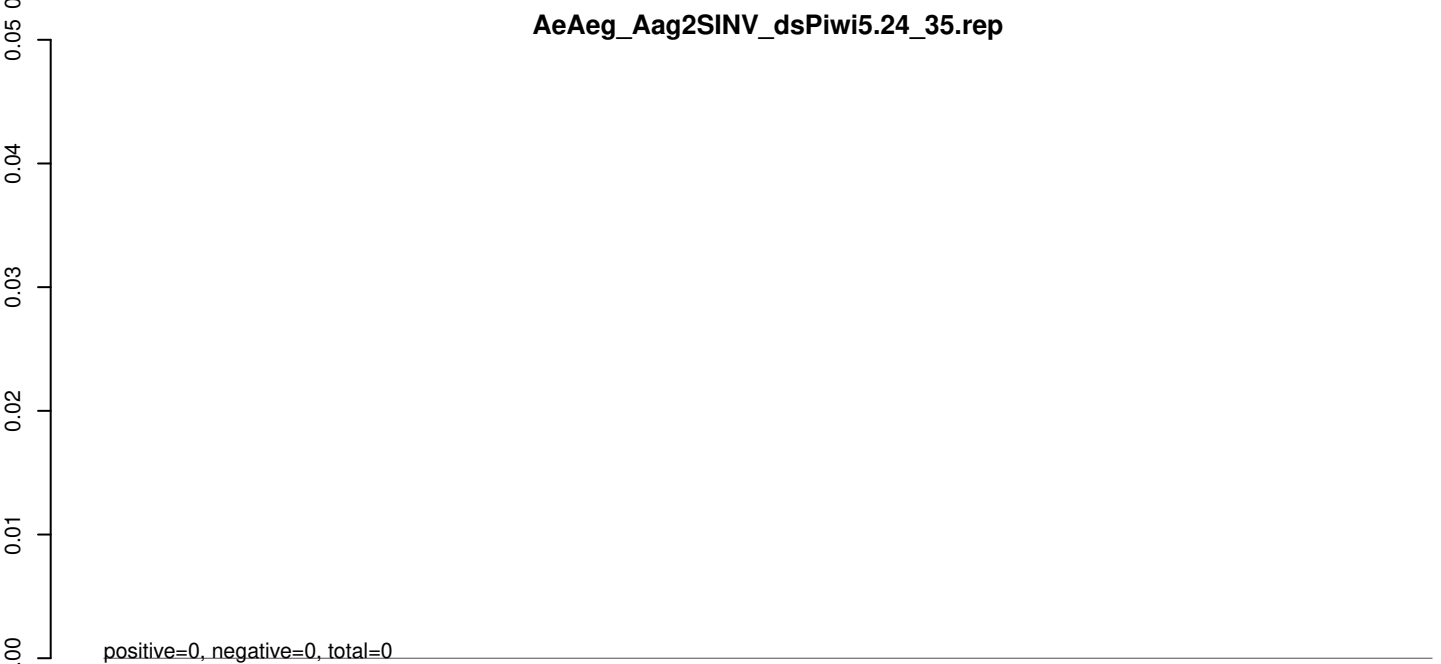
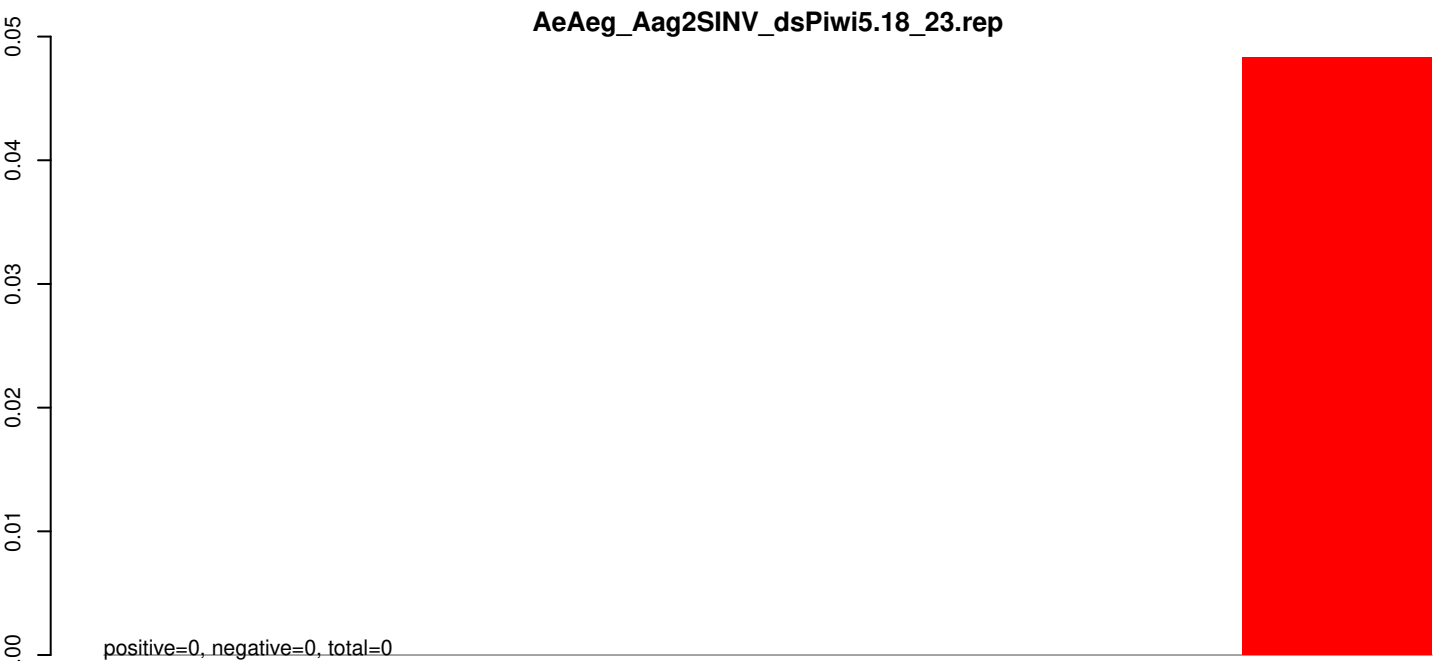


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AeAeg_Aag2SINV_dsPiwi5.rep





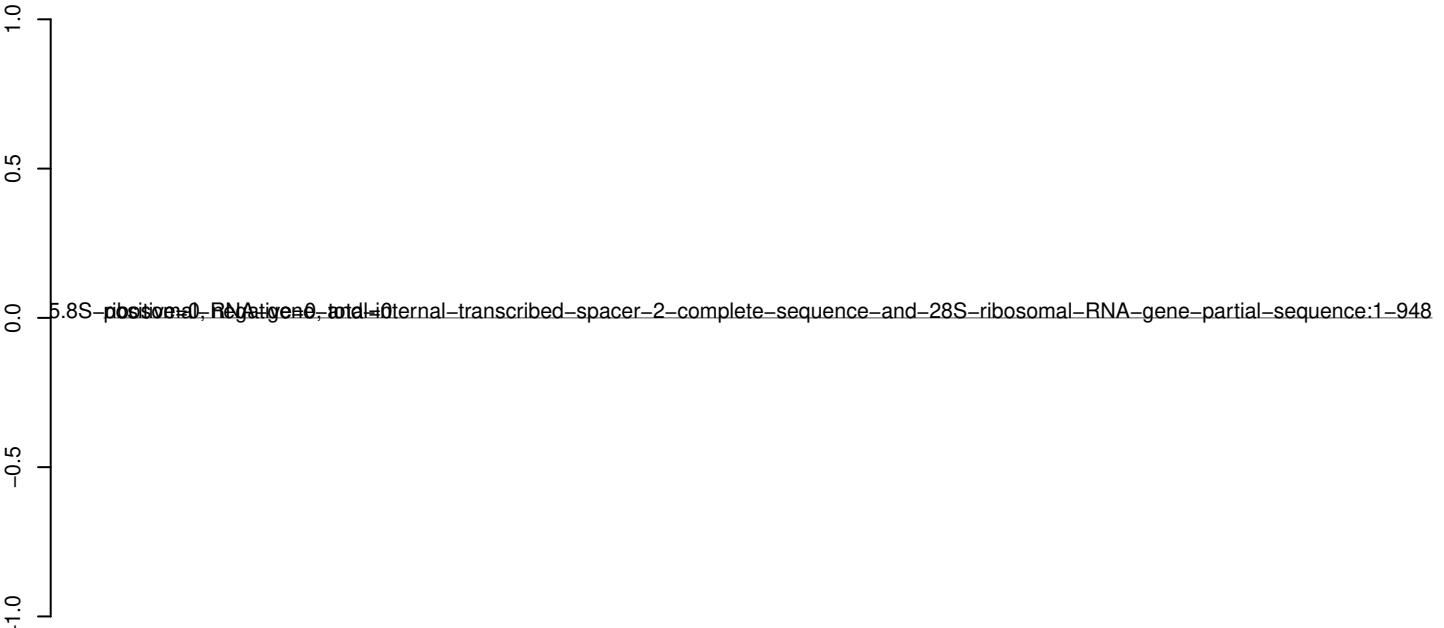
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AeAeg_Aag2SINV_dsPiwi5.rep



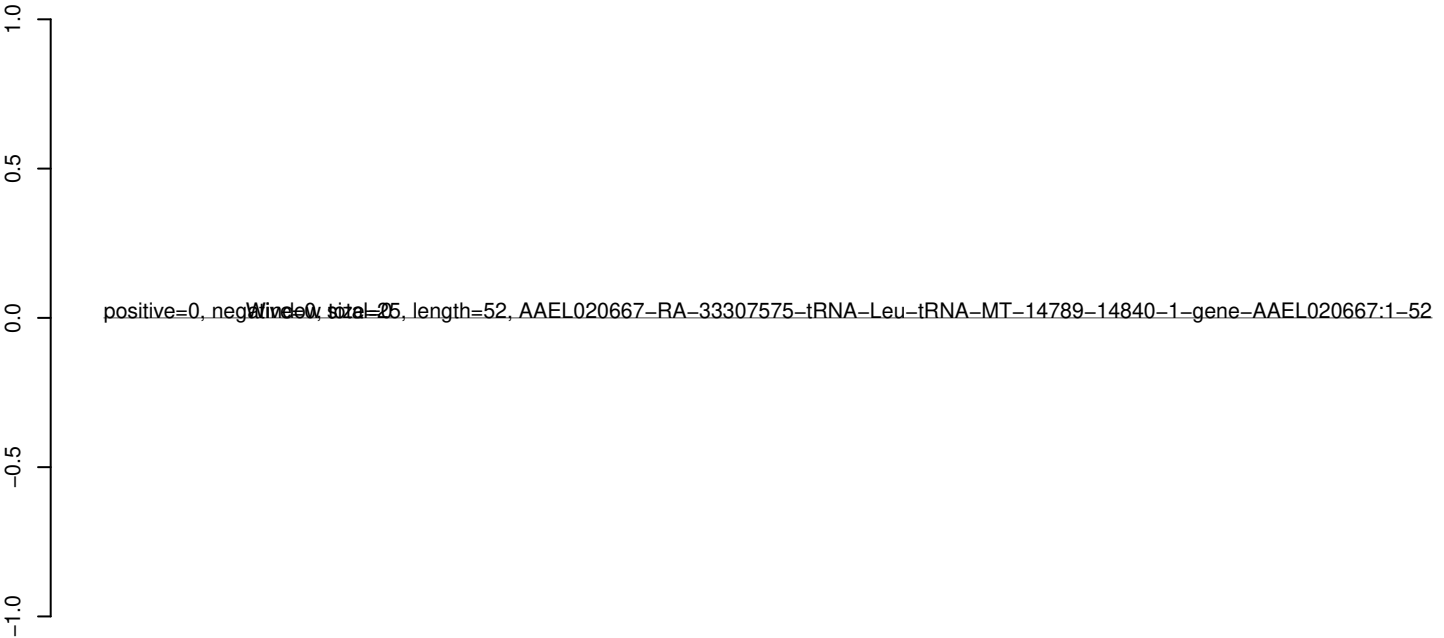
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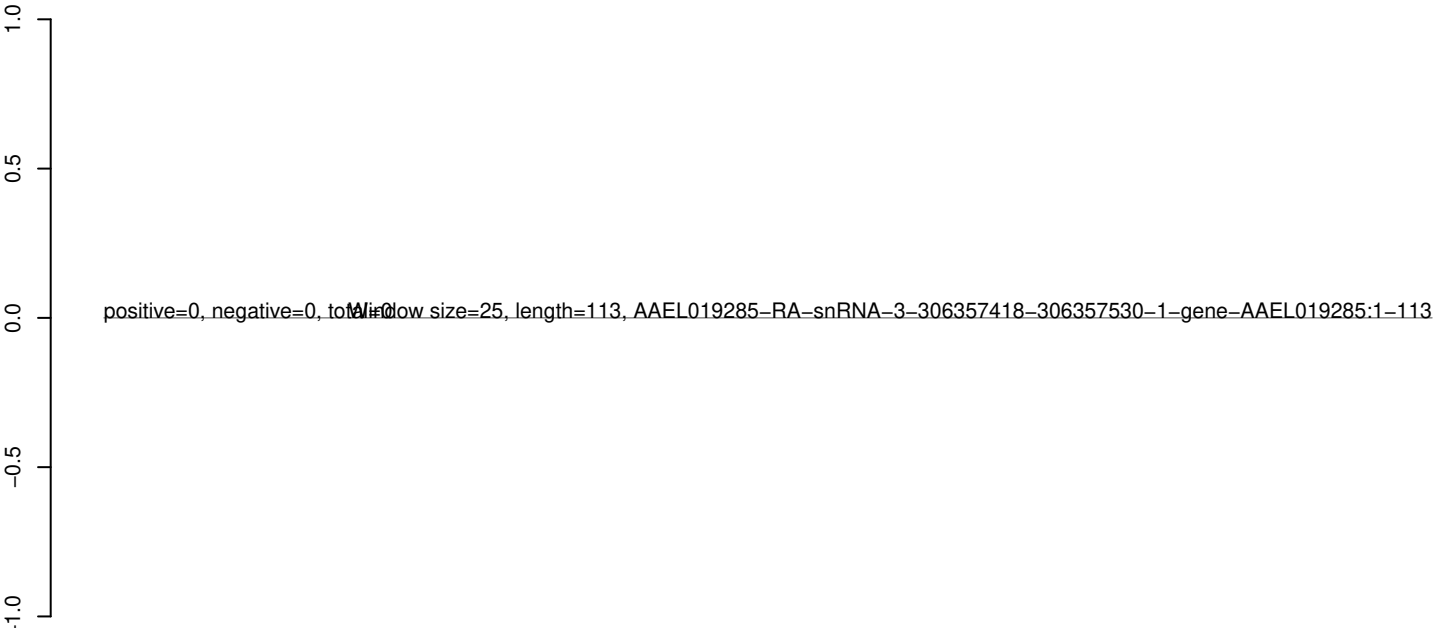
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AeAeg_Aag2SINV_dsPiwi5.rep



40 60 80 100 120 140

AeAeg_Aag2SINV_dsPiwi5.18_23.rep



AeAeg_Aag2SINV_dsPiwi5.24_35.rep



AeAeg_Aag2SINV_dsPiwi5.rep

