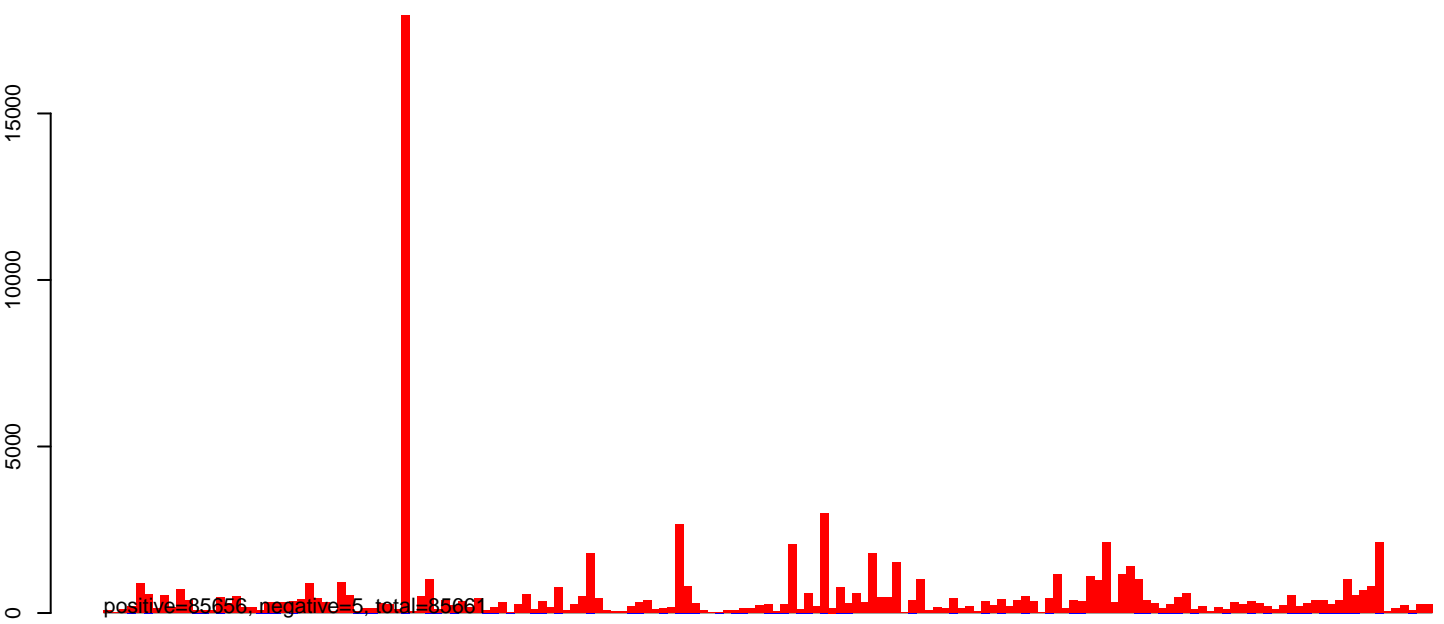


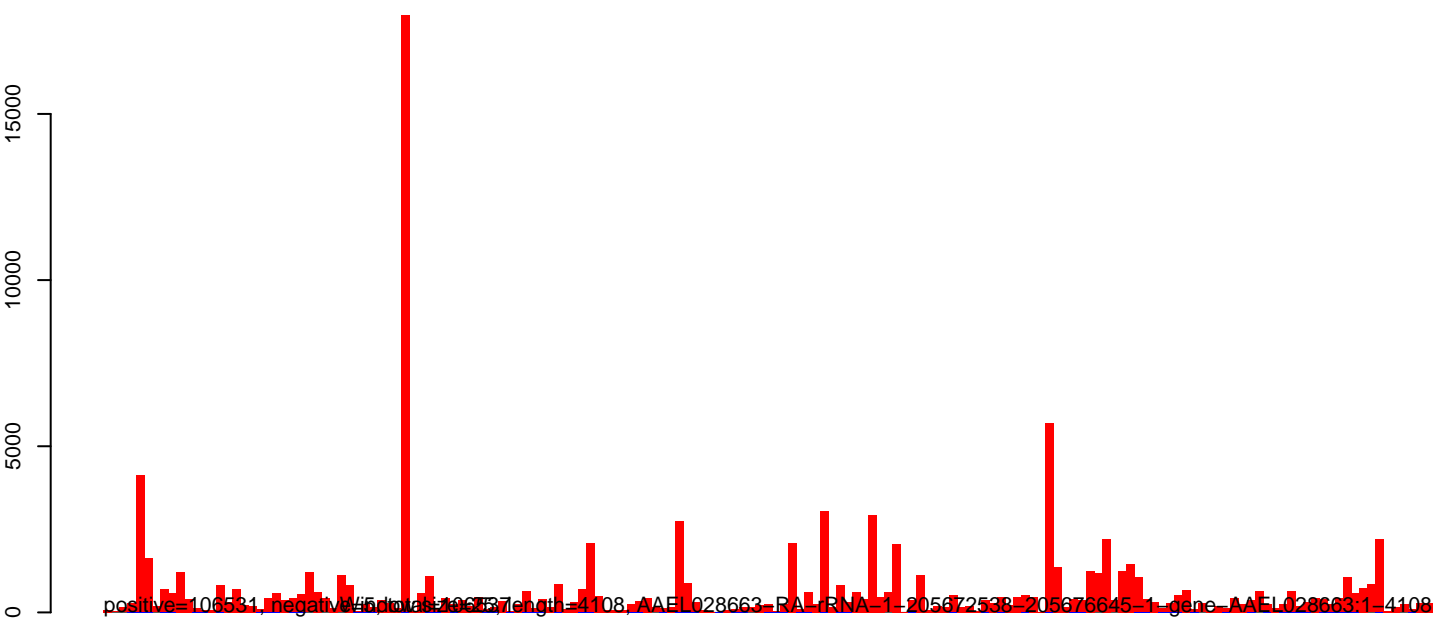
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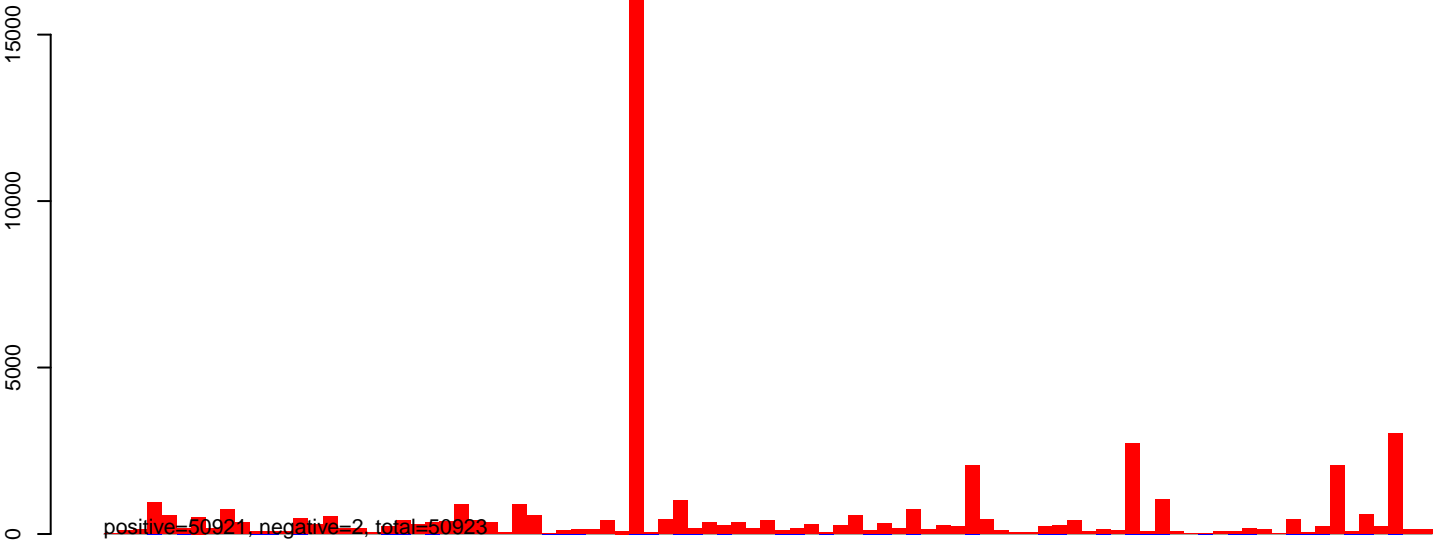
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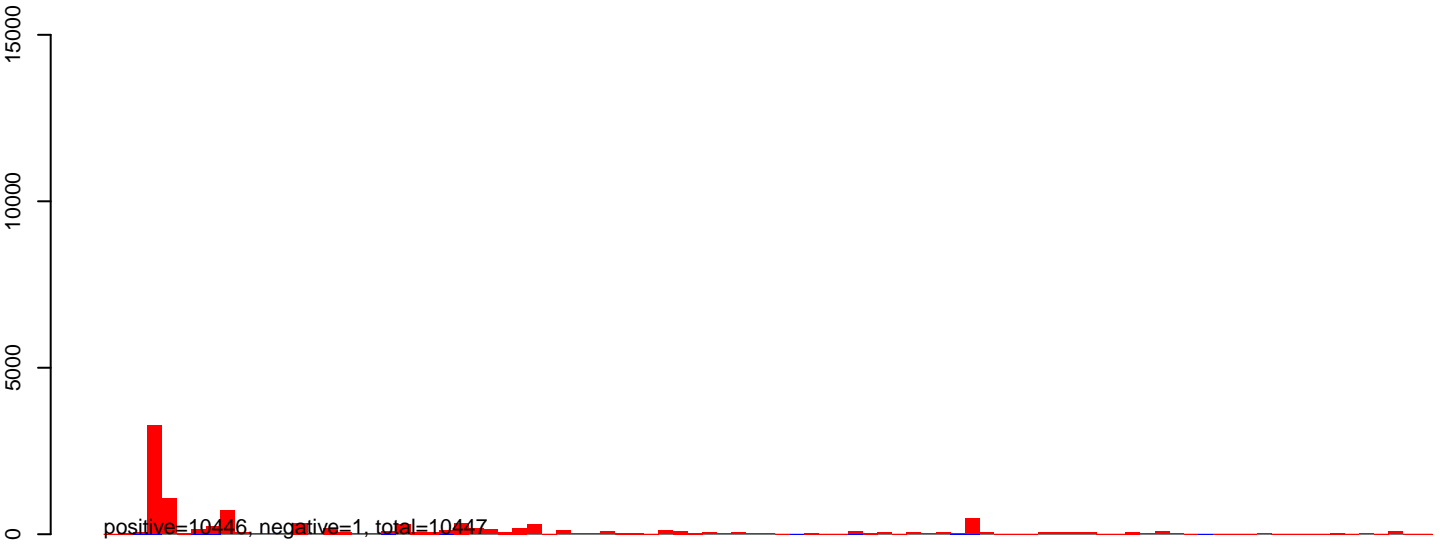
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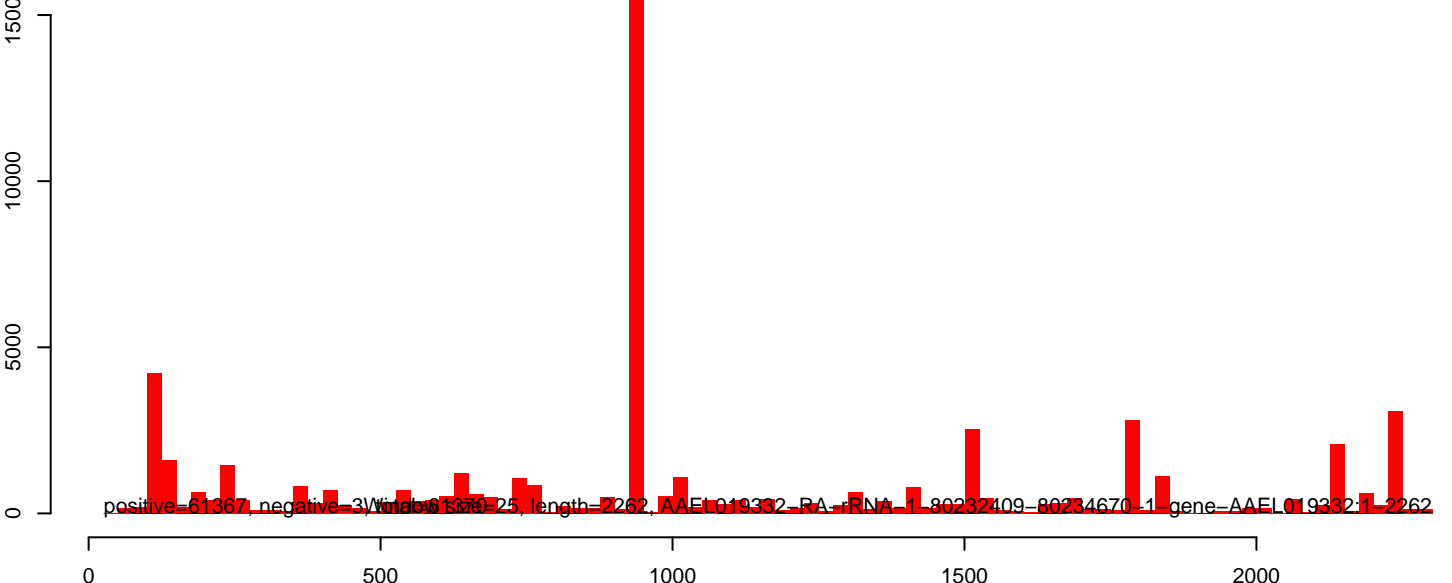
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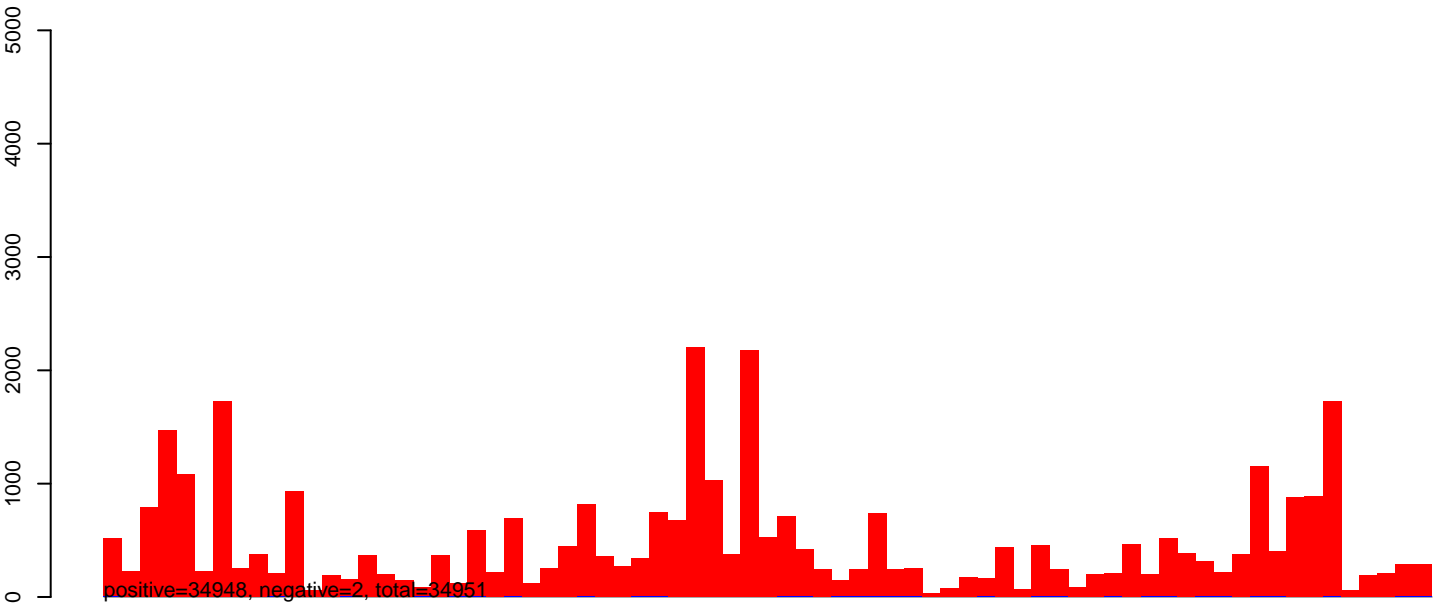
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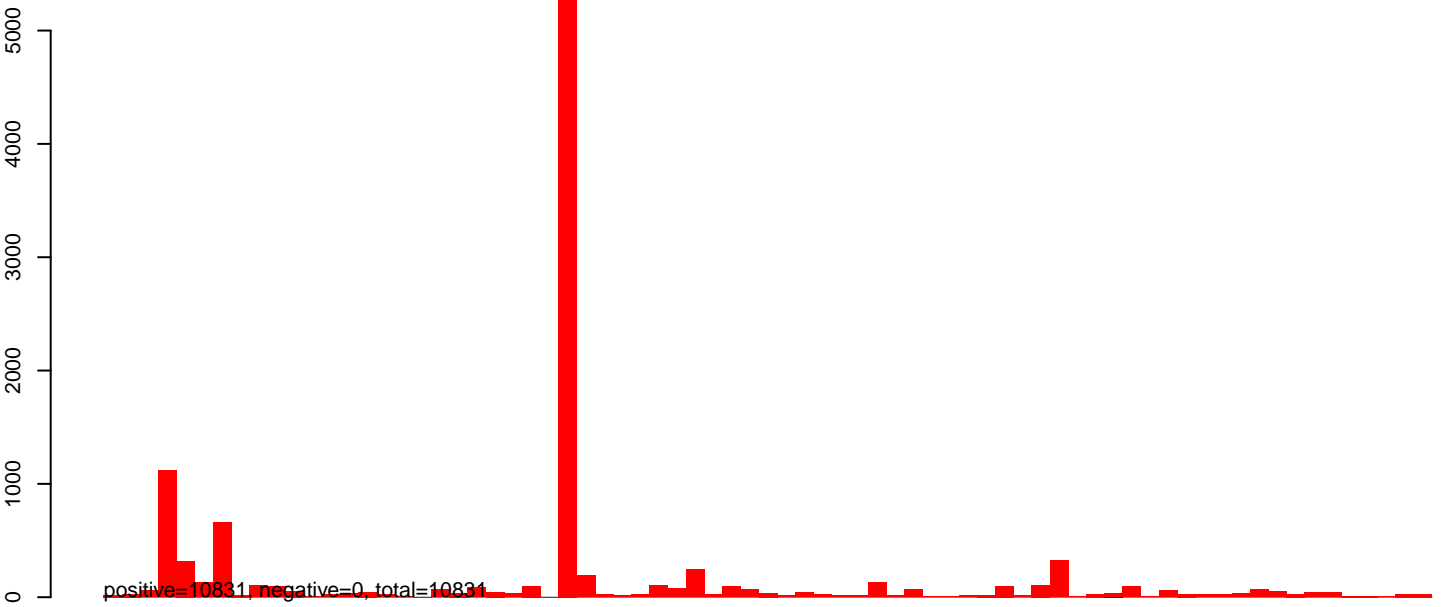
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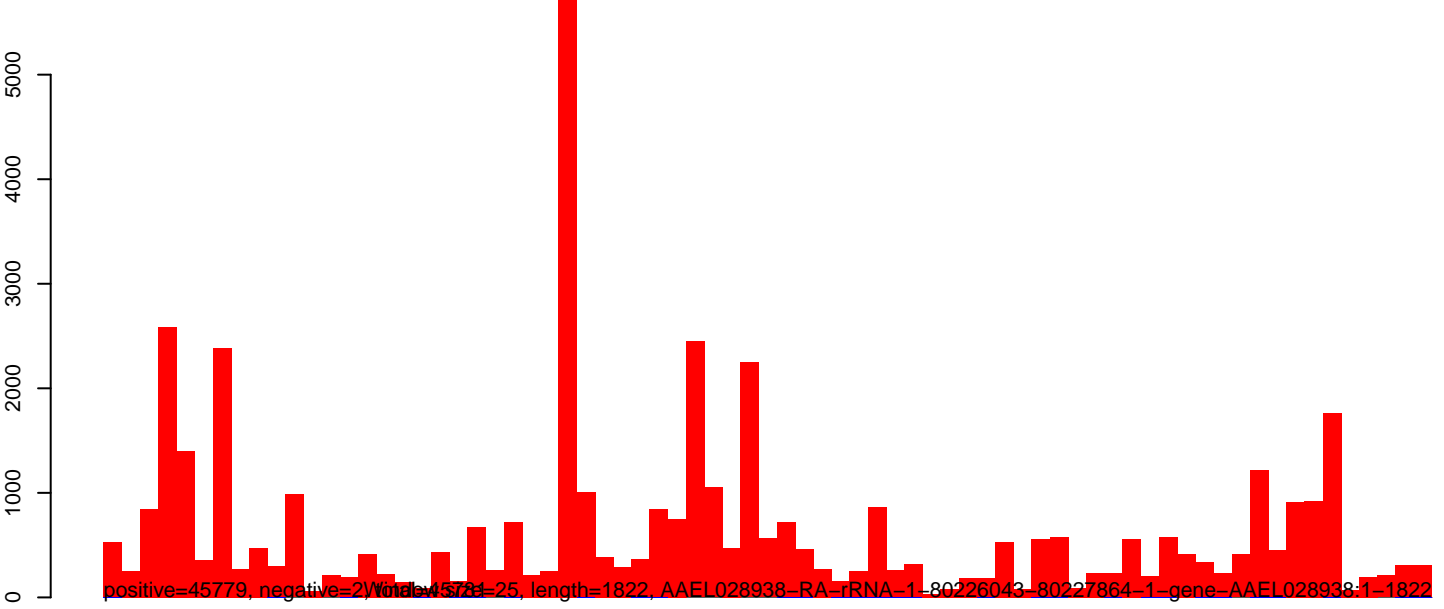
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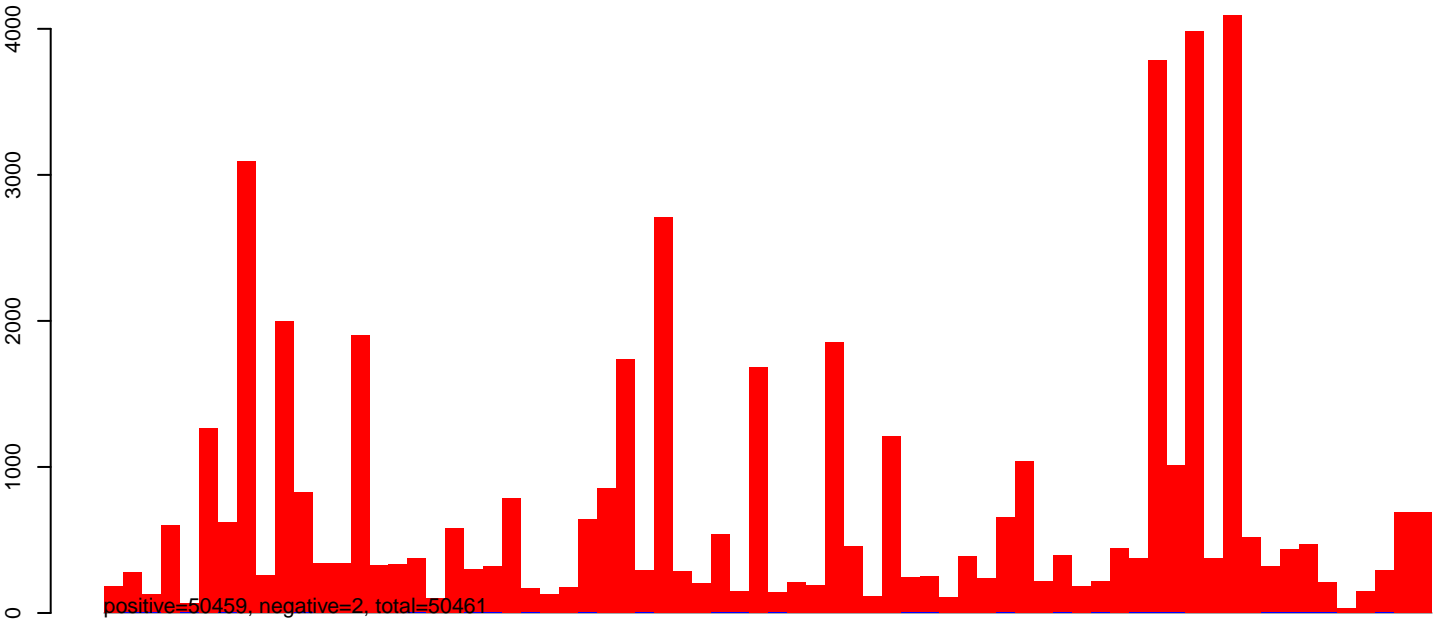
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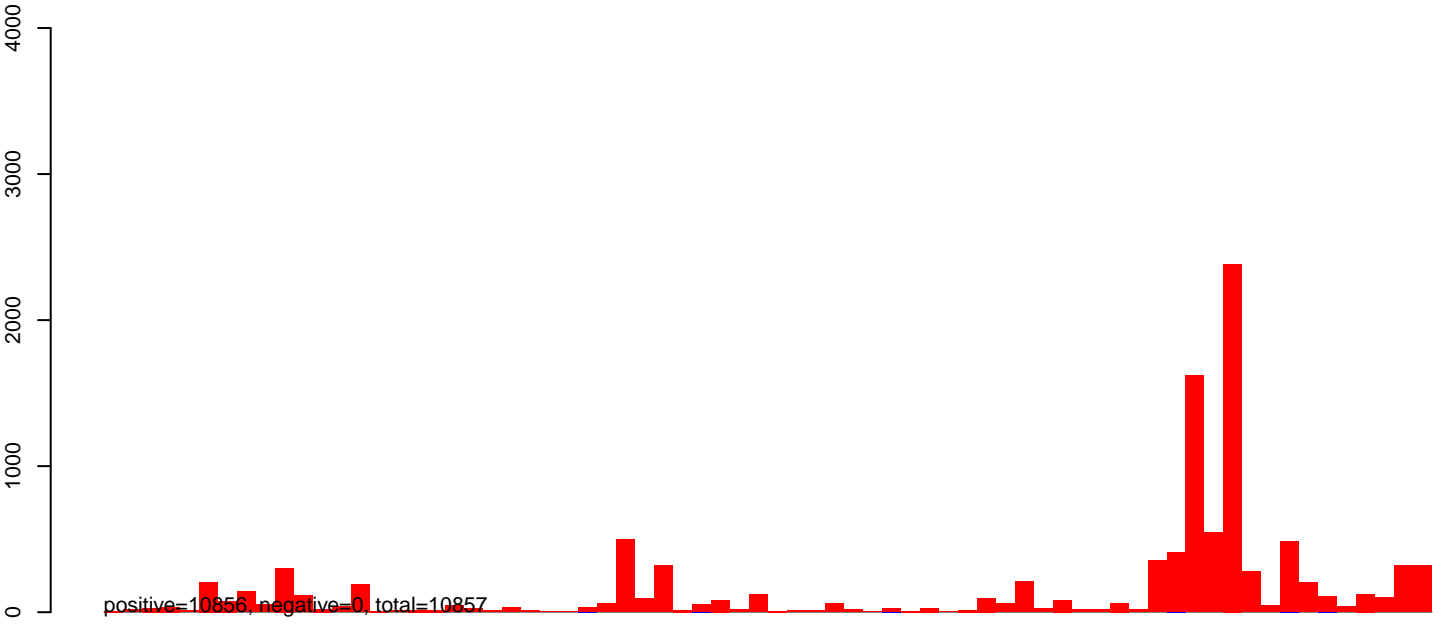
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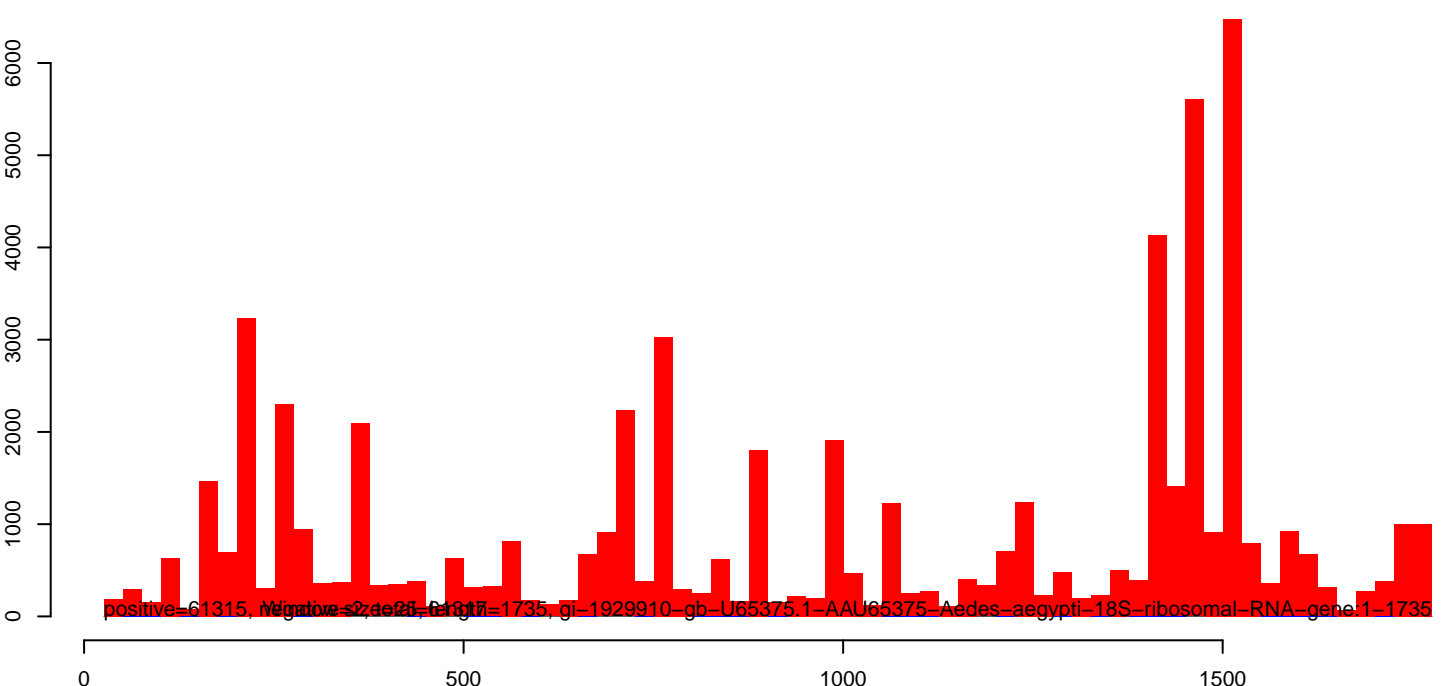
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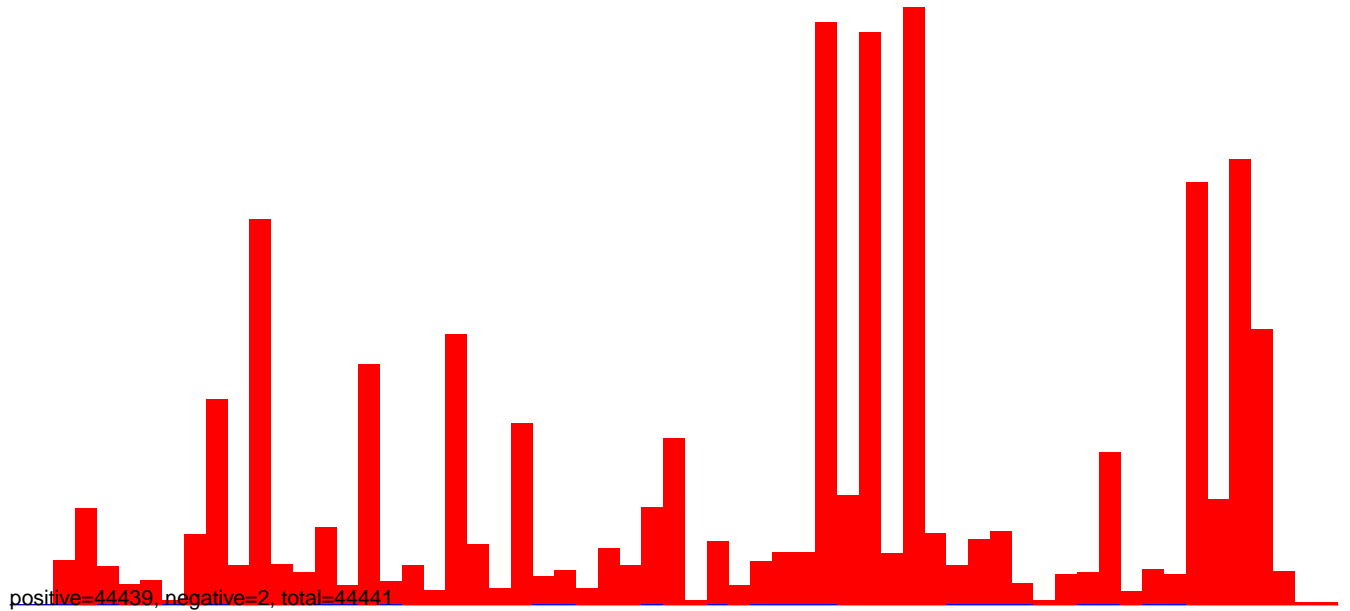
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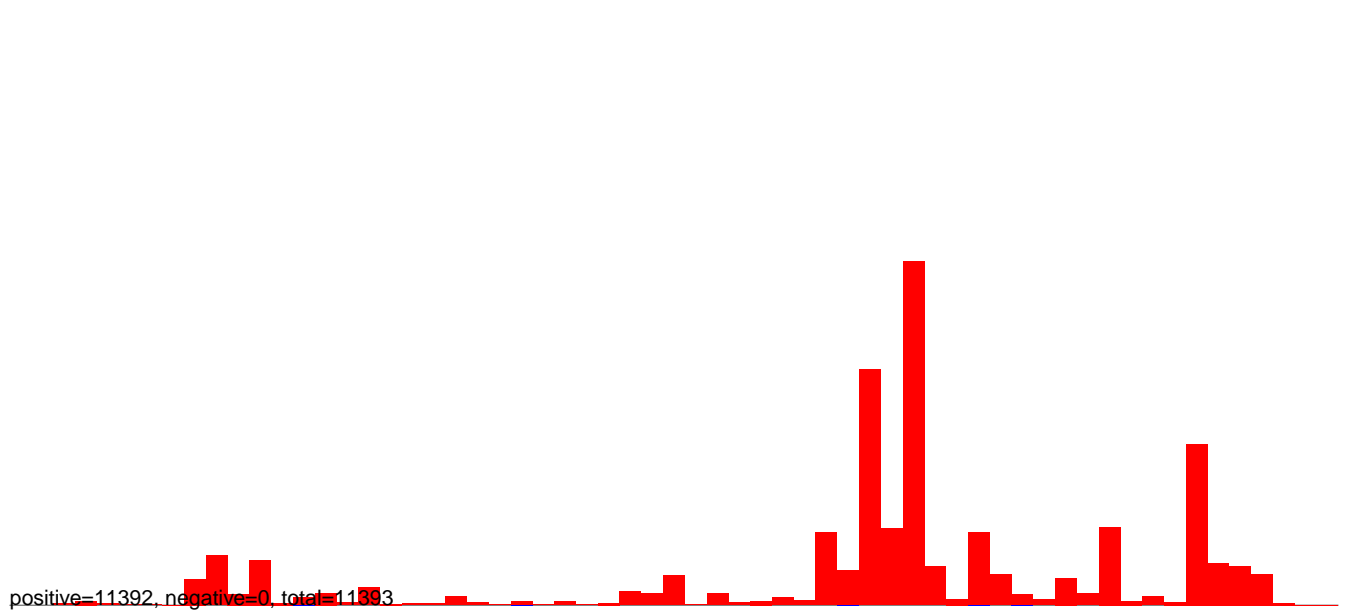
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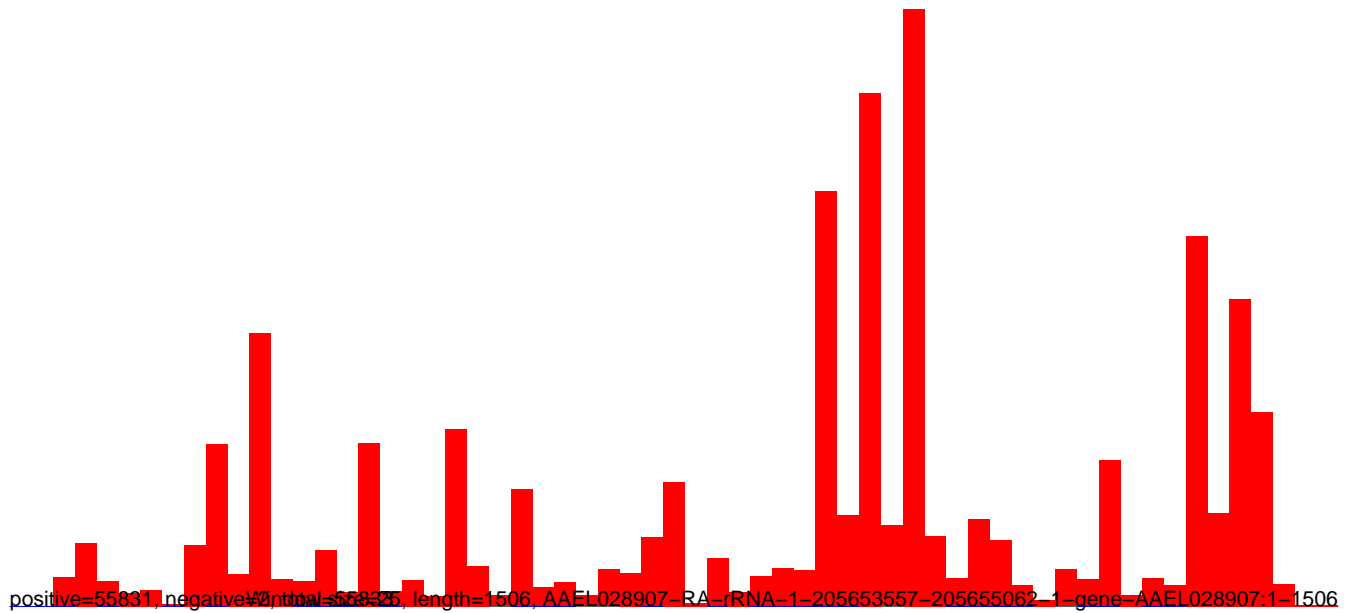
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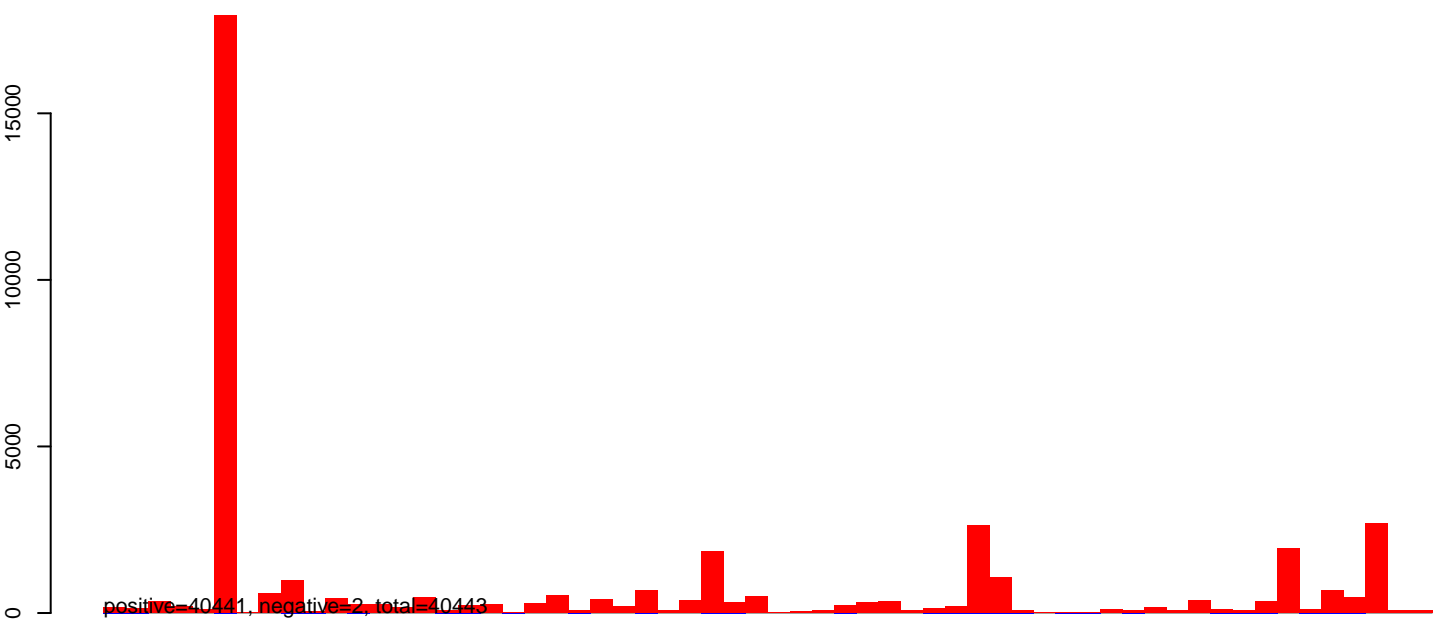
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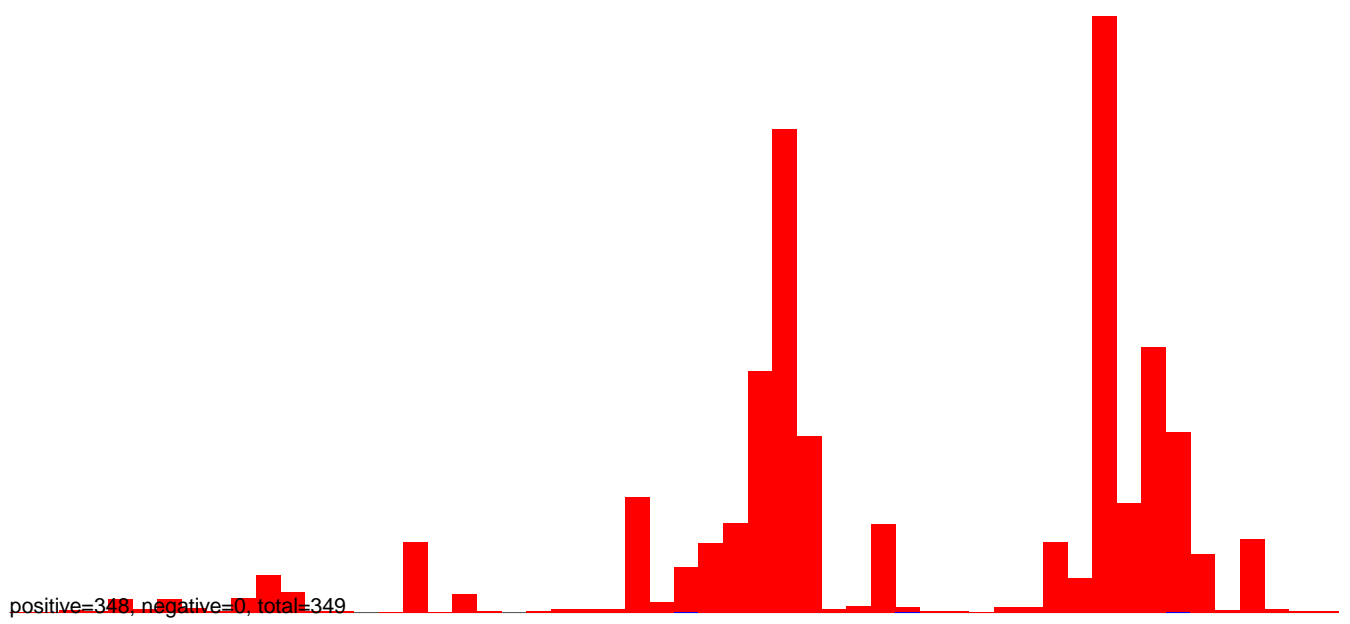
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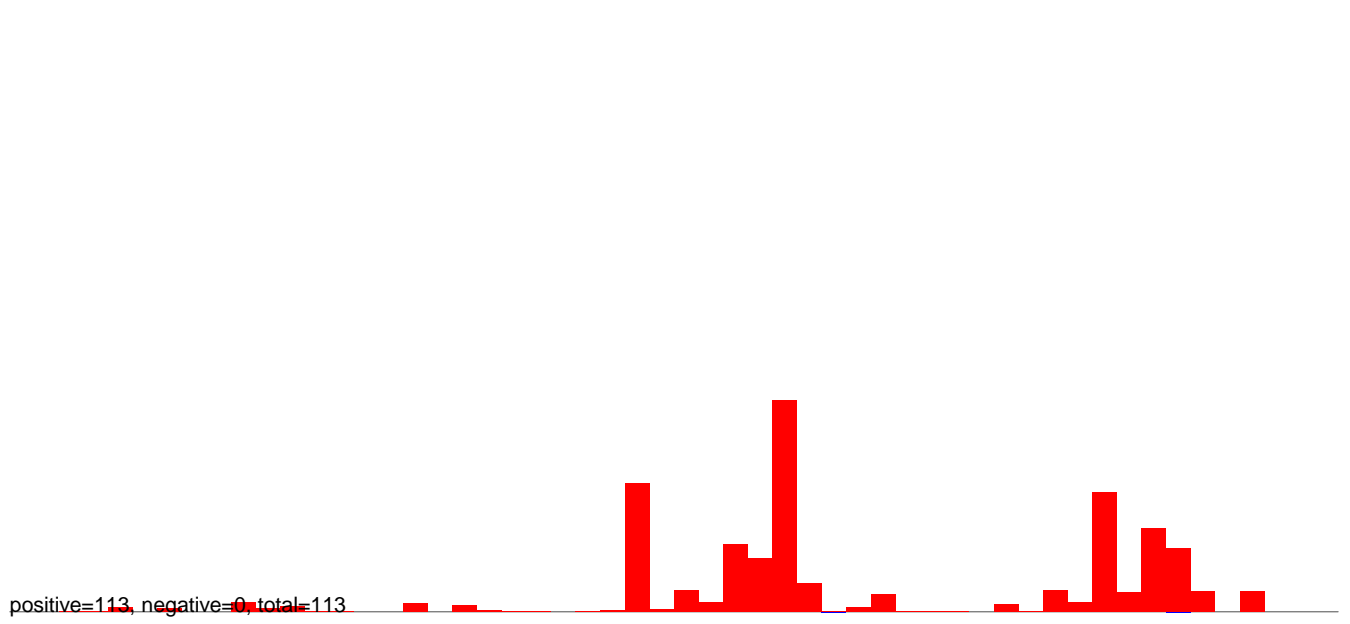
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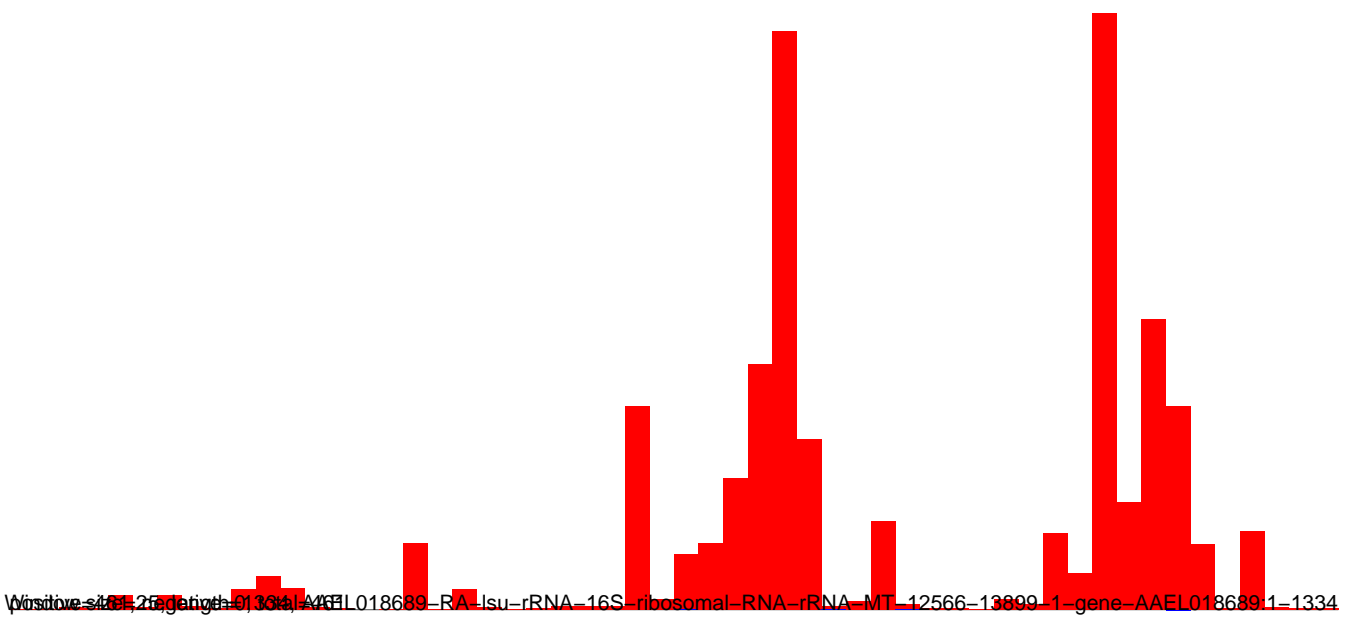
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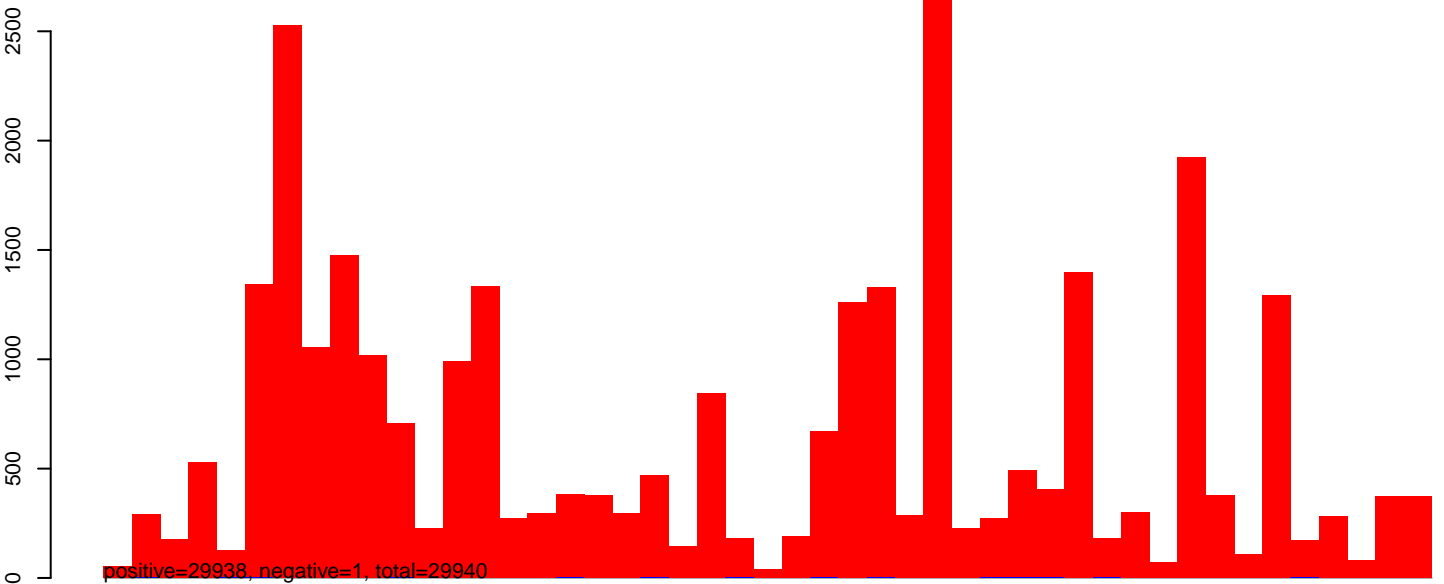


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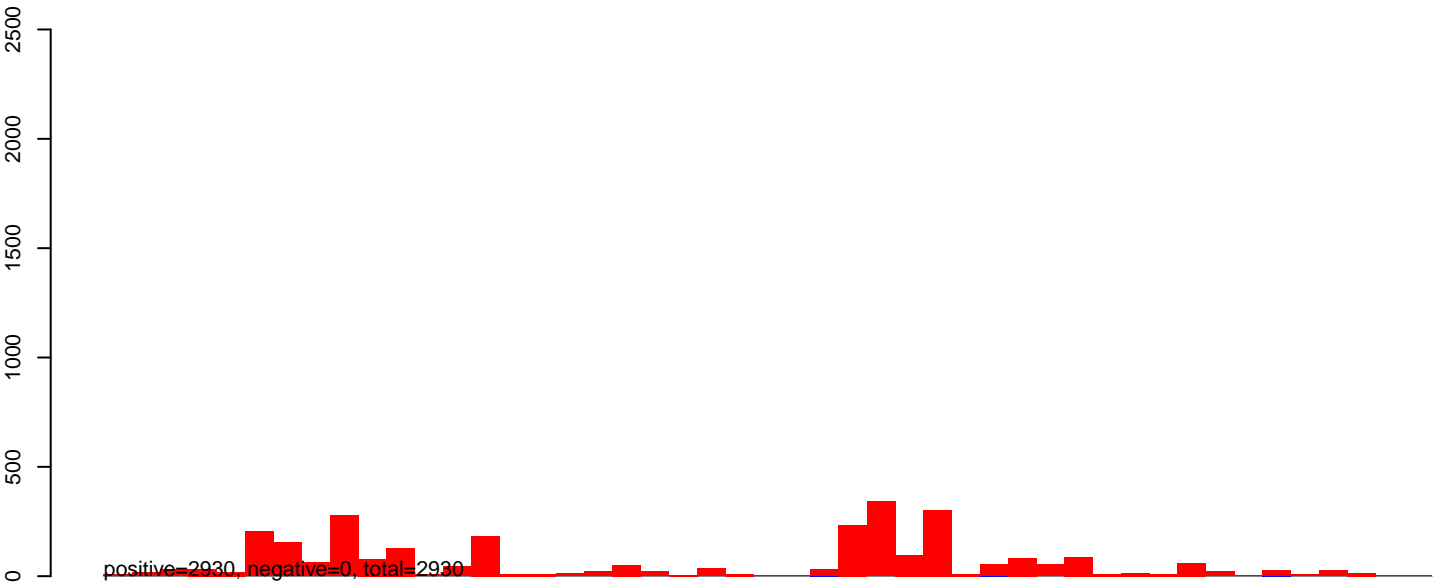


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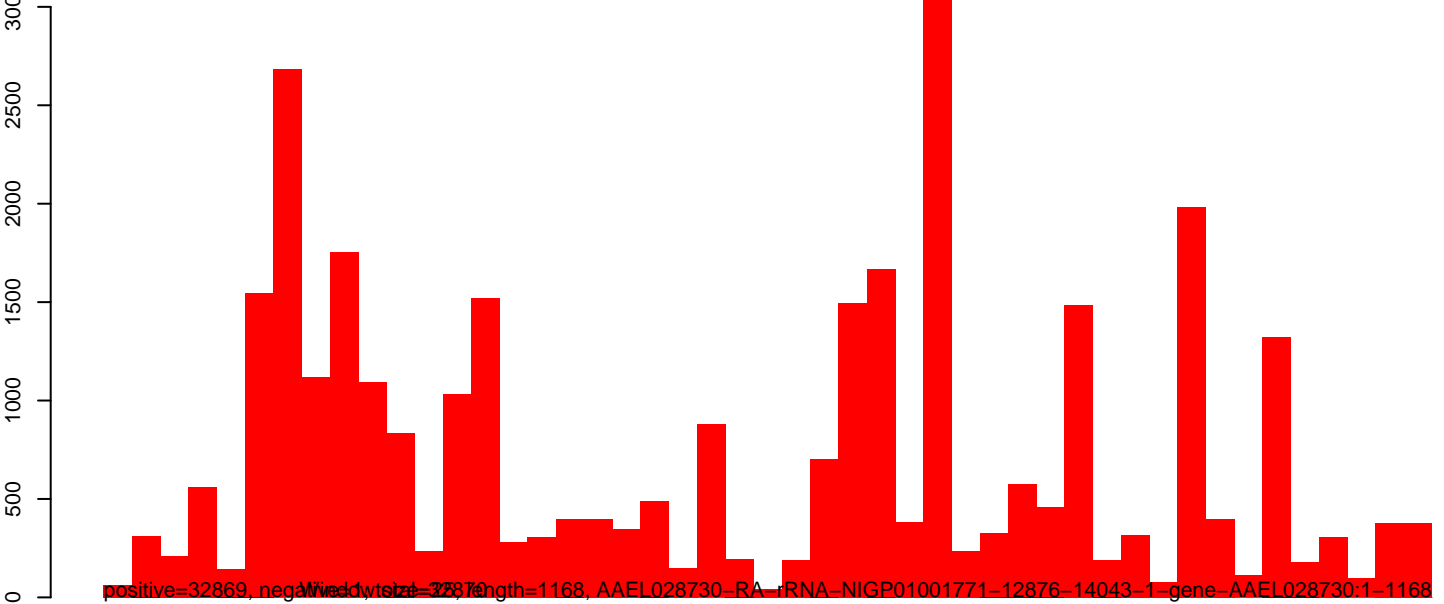
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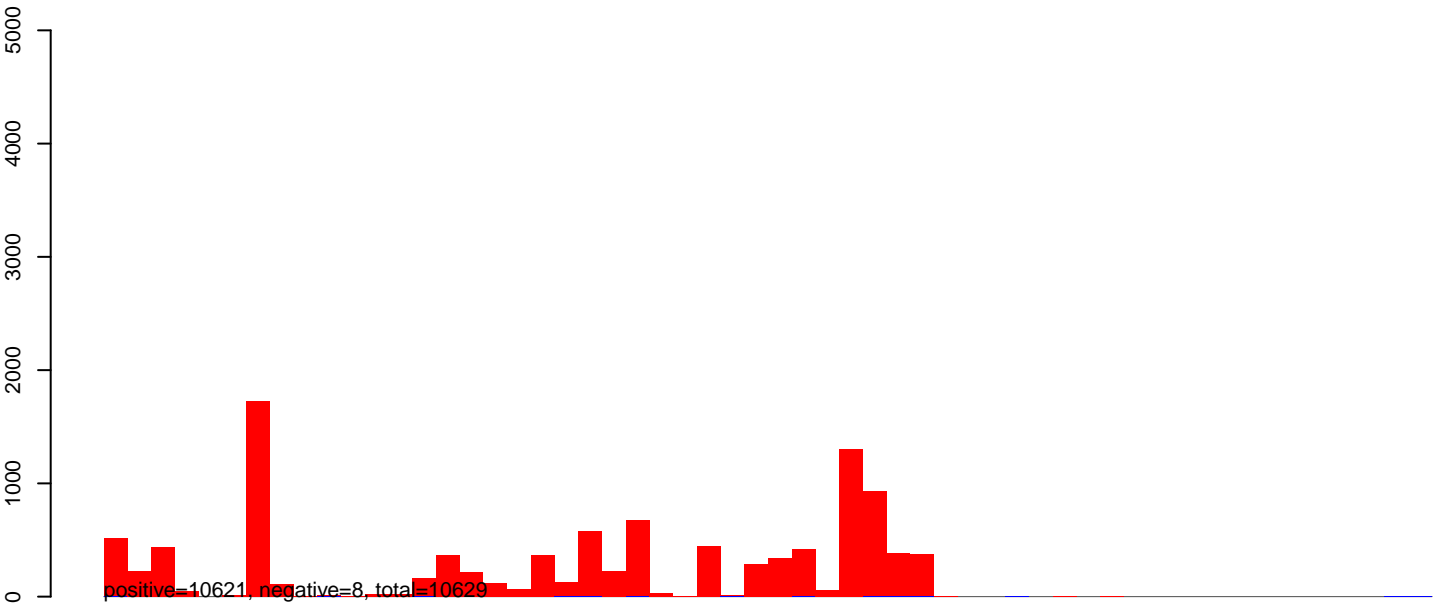


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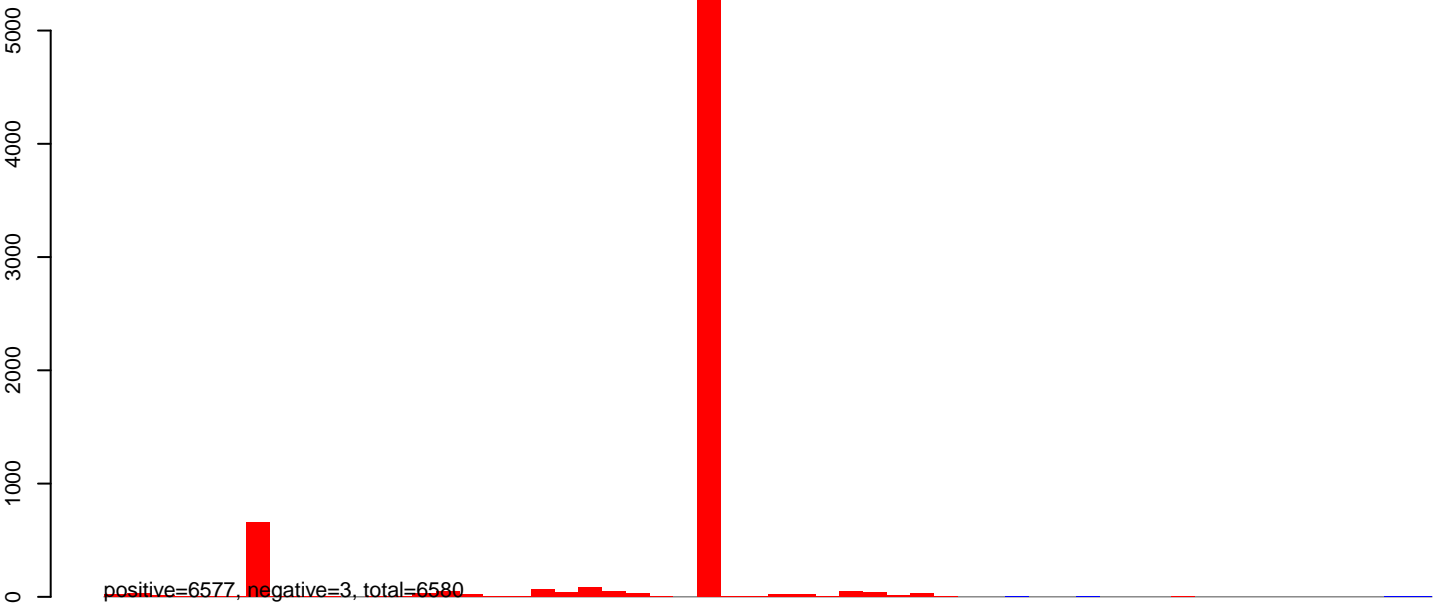


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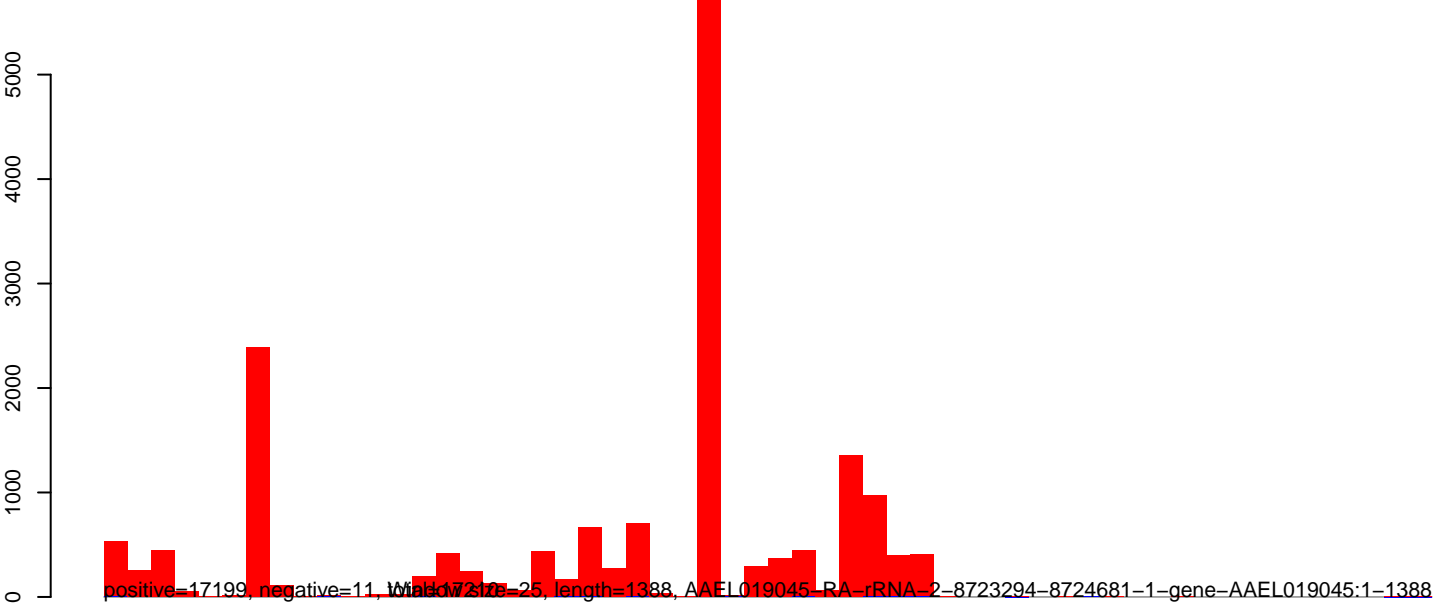
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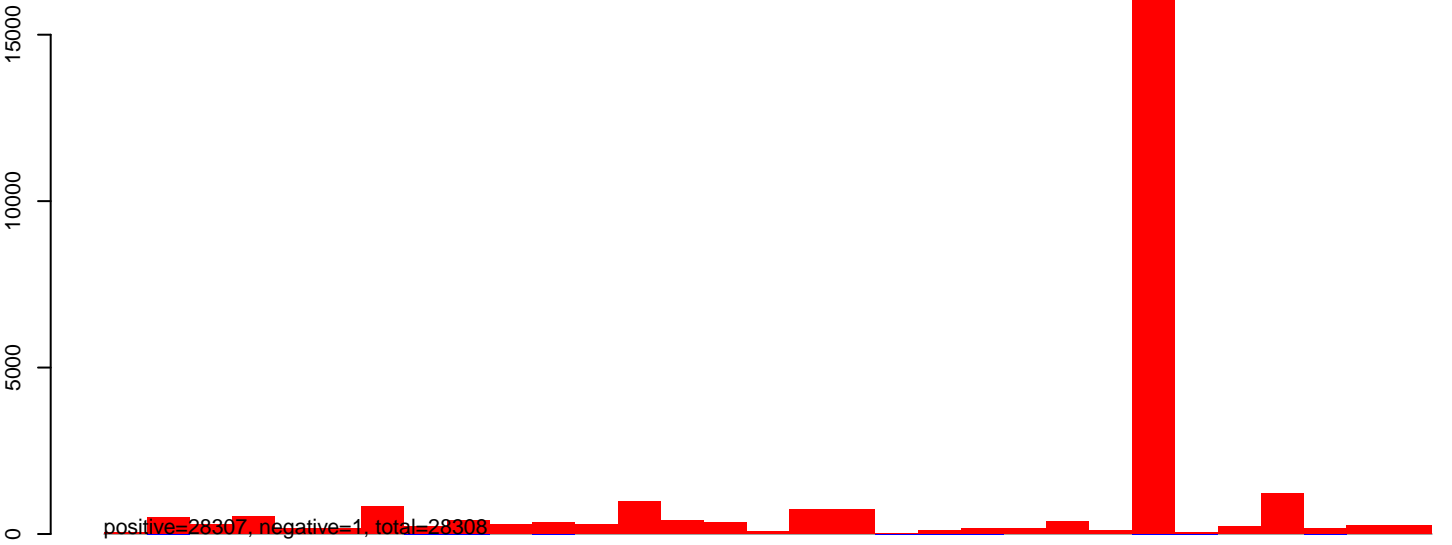
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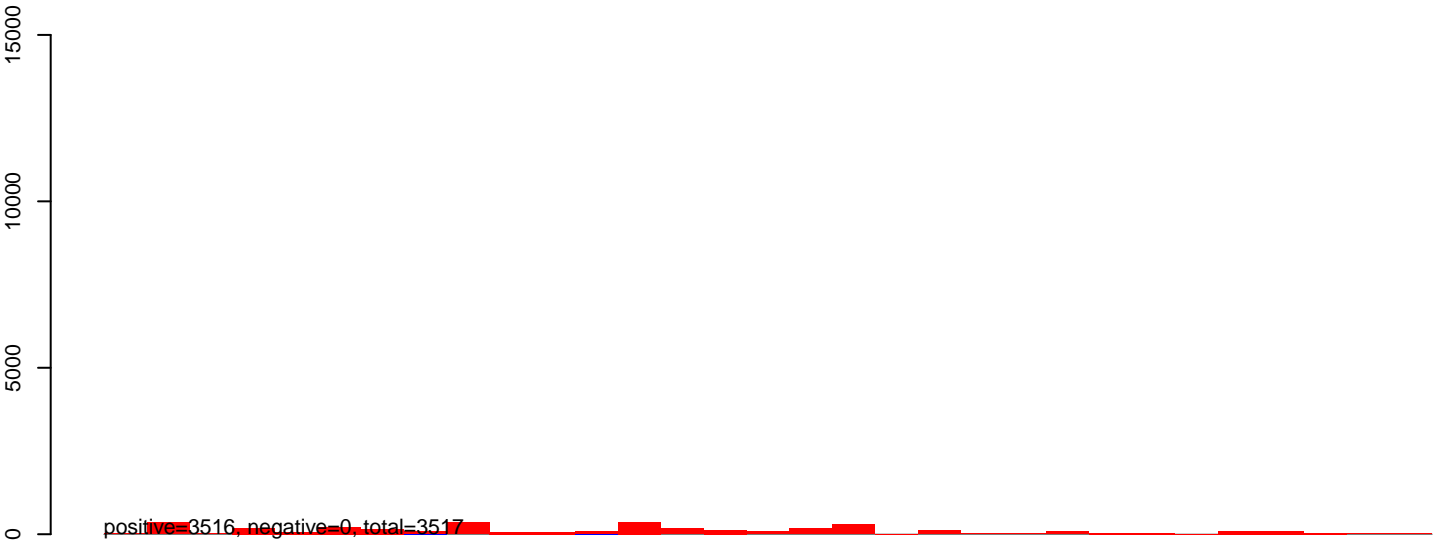
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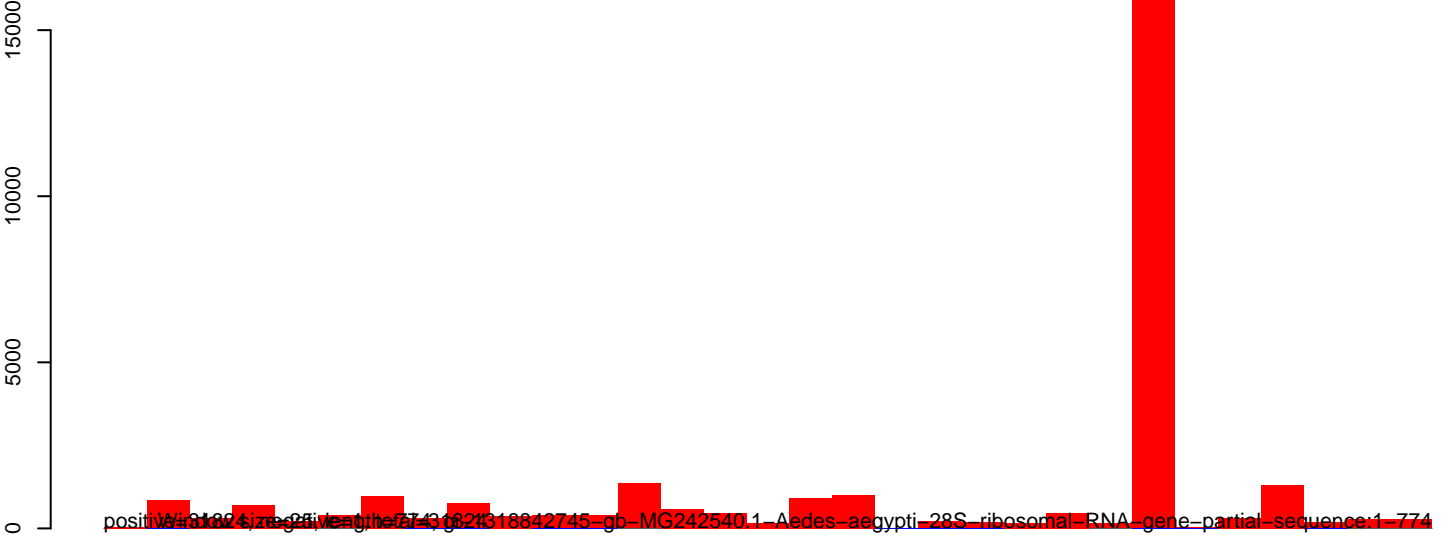
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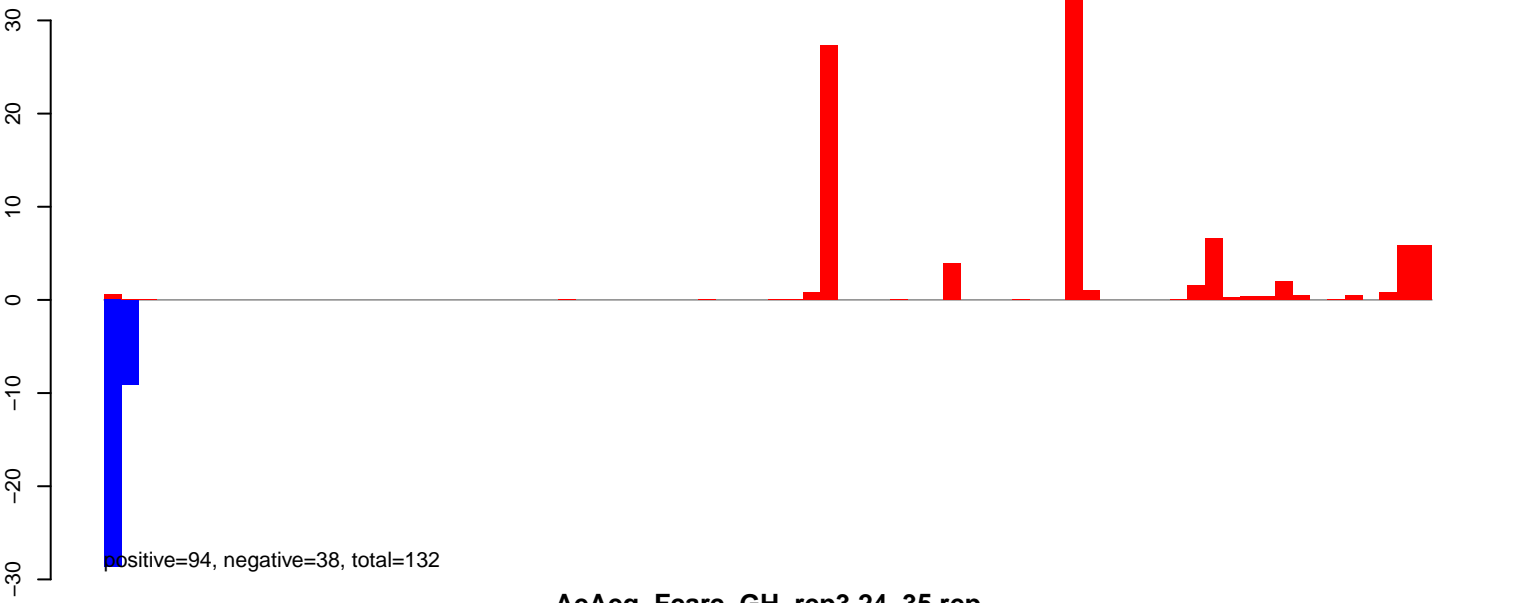
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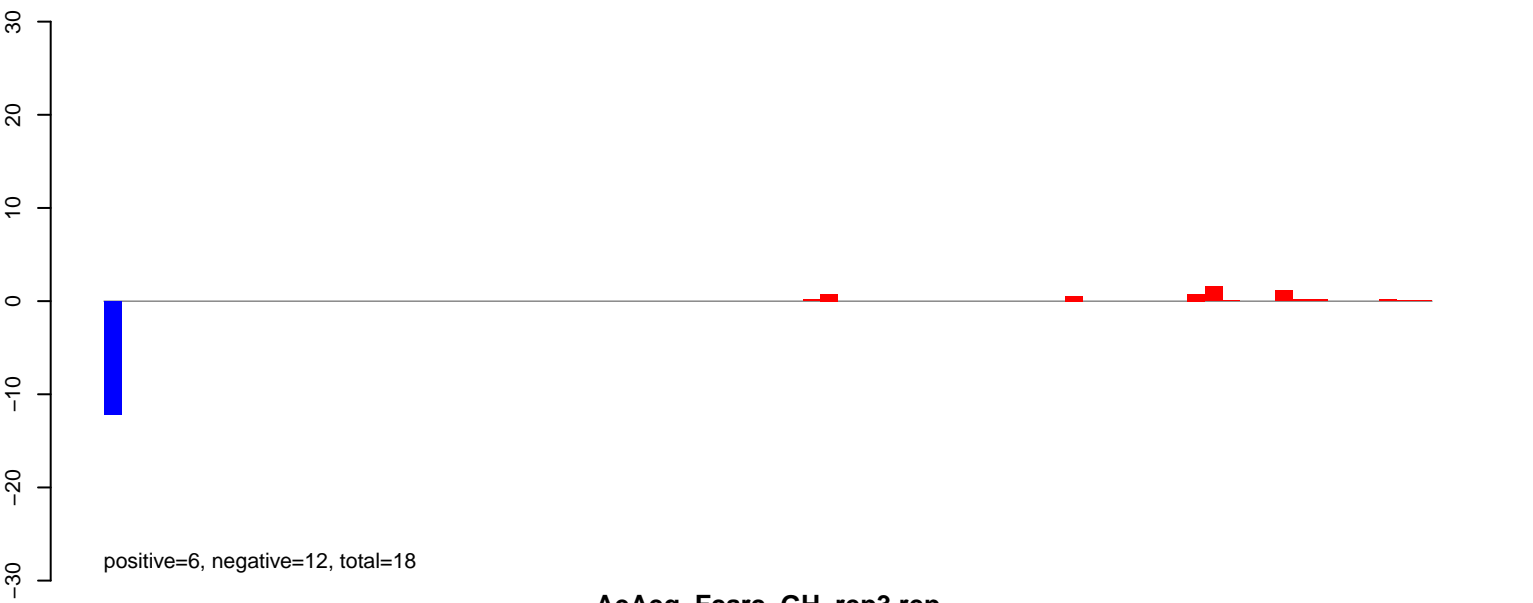
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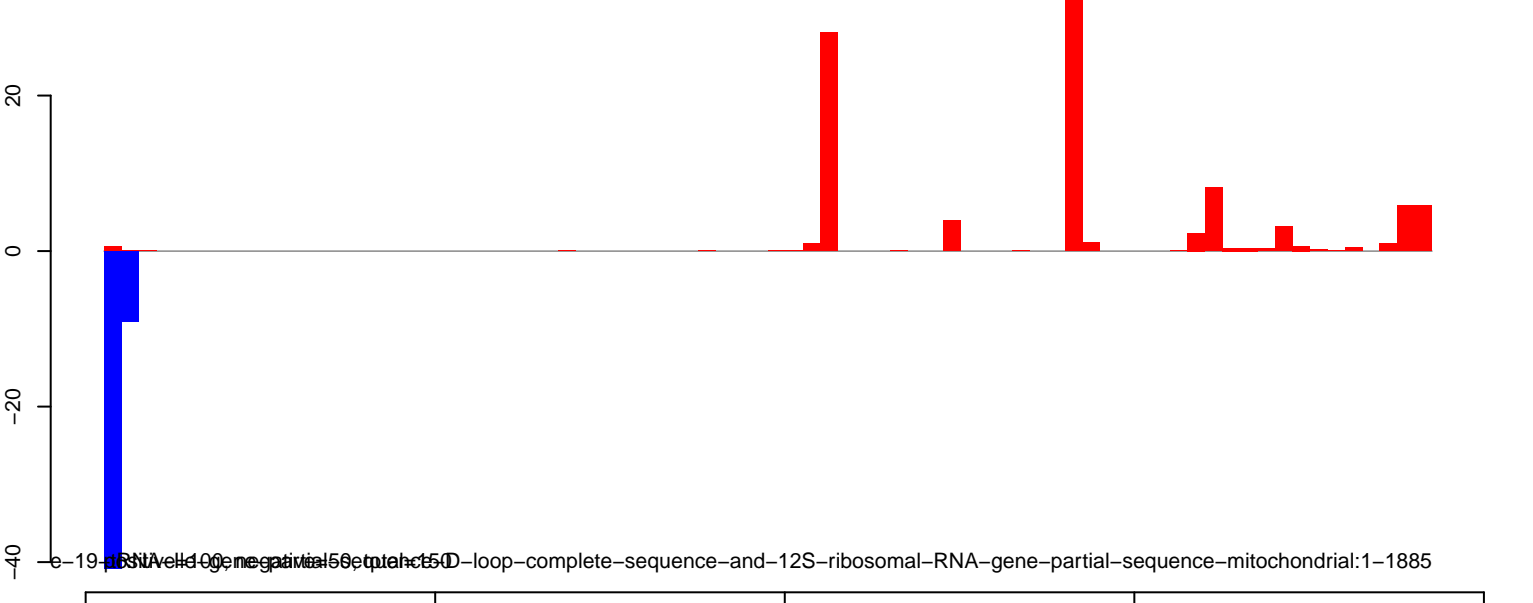
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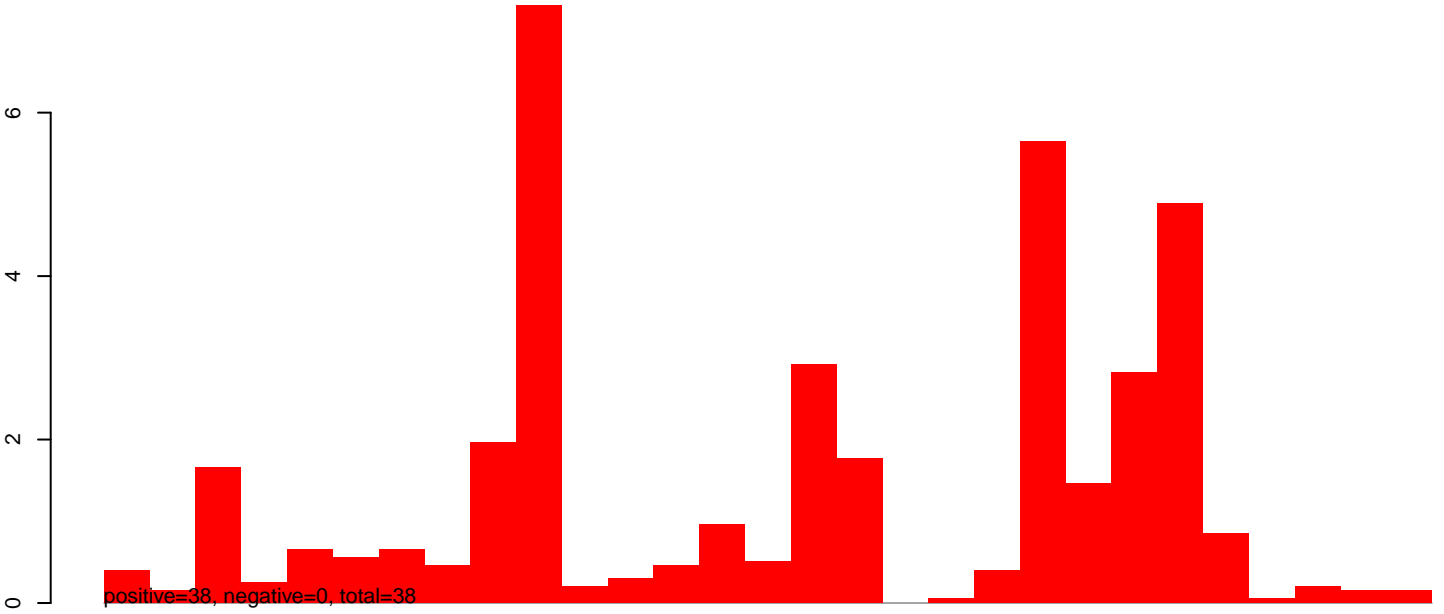


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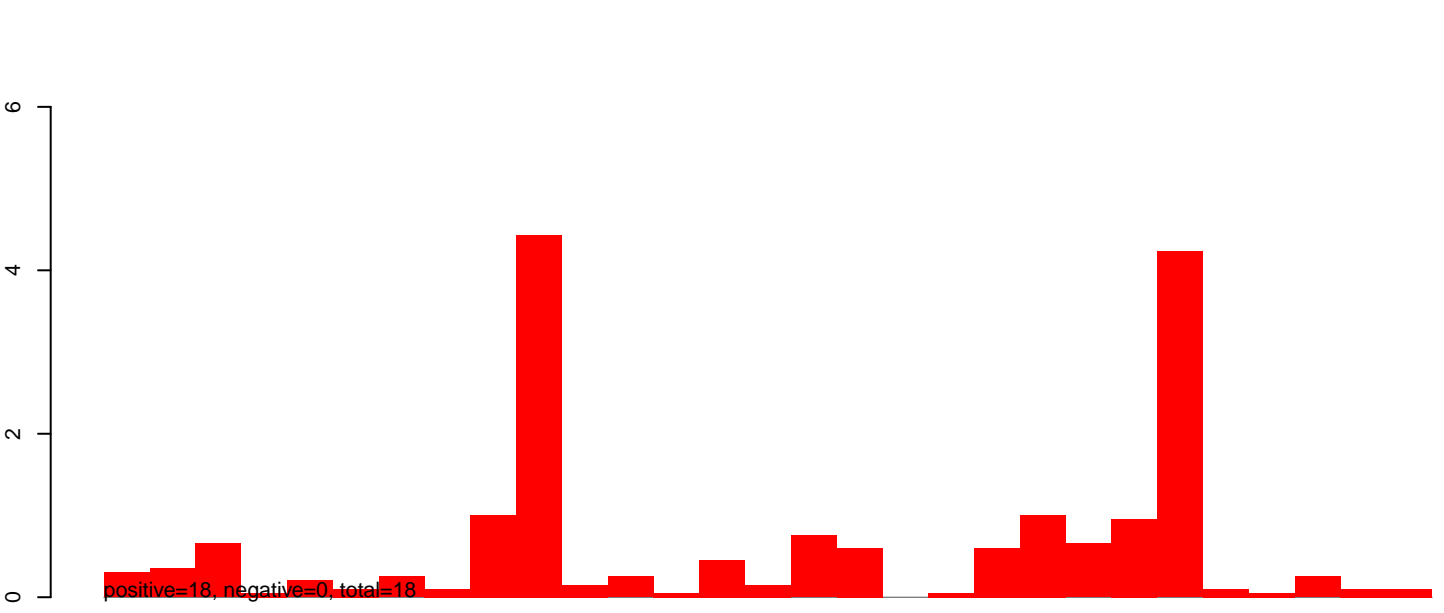


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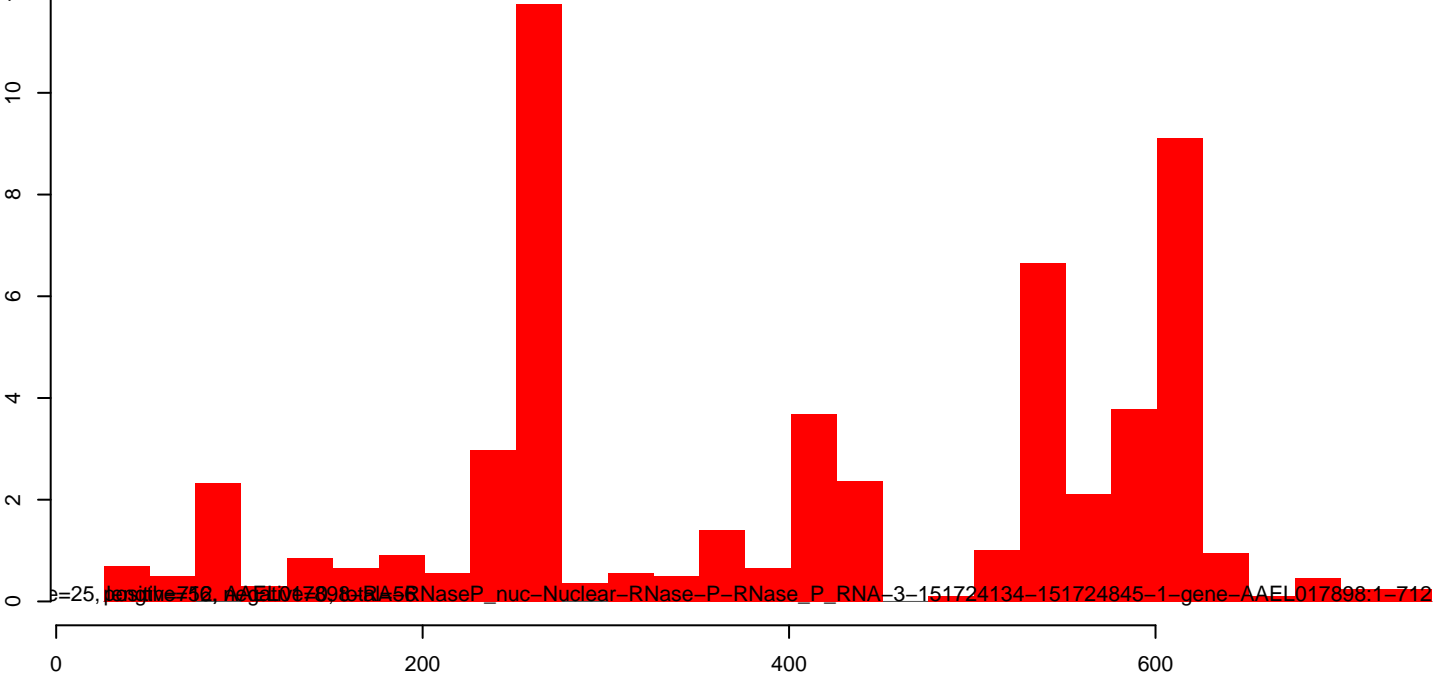
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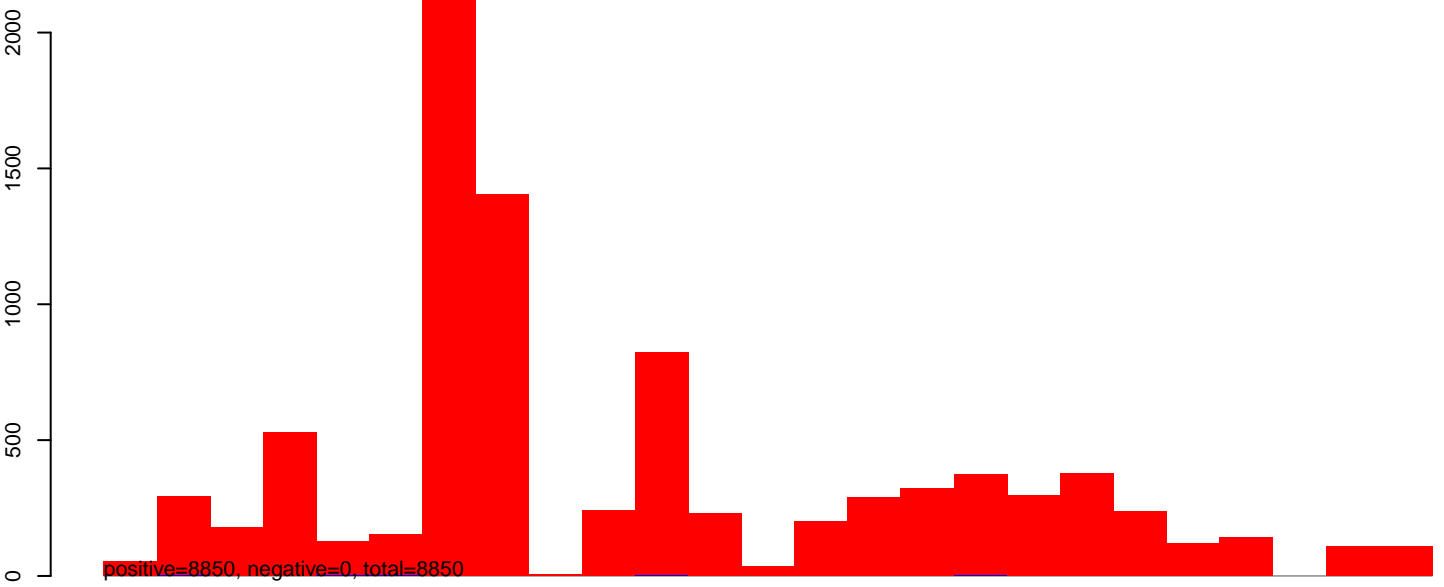
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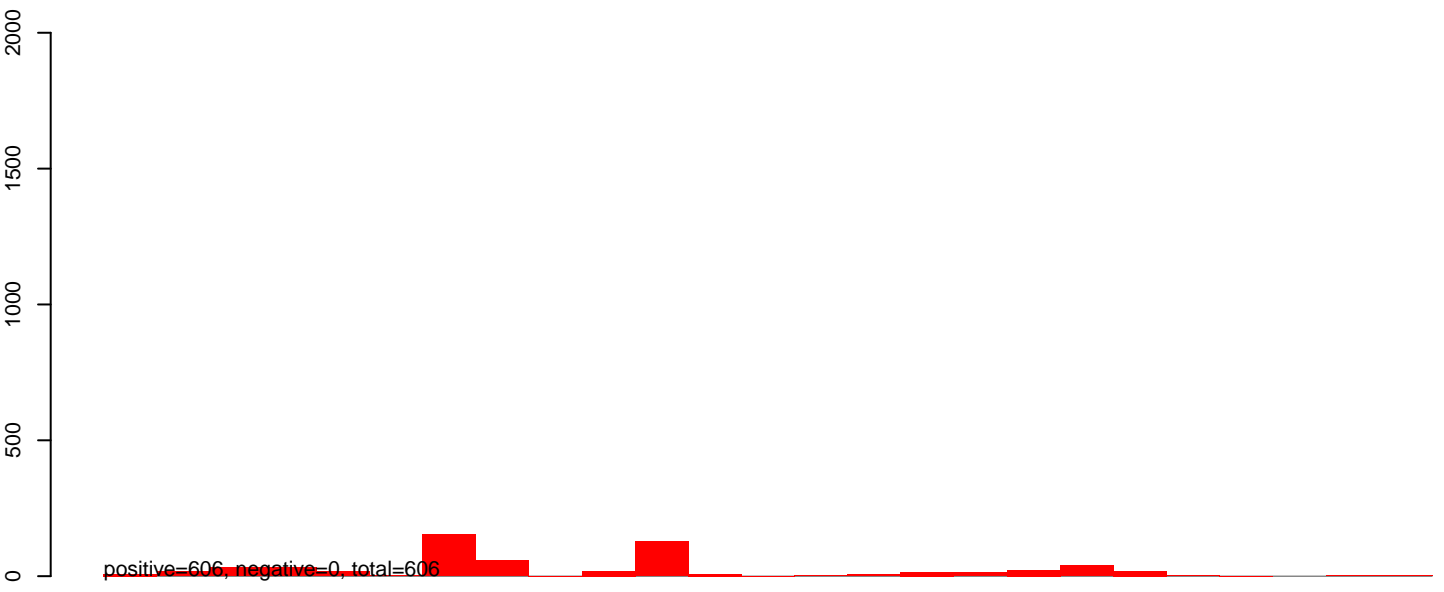
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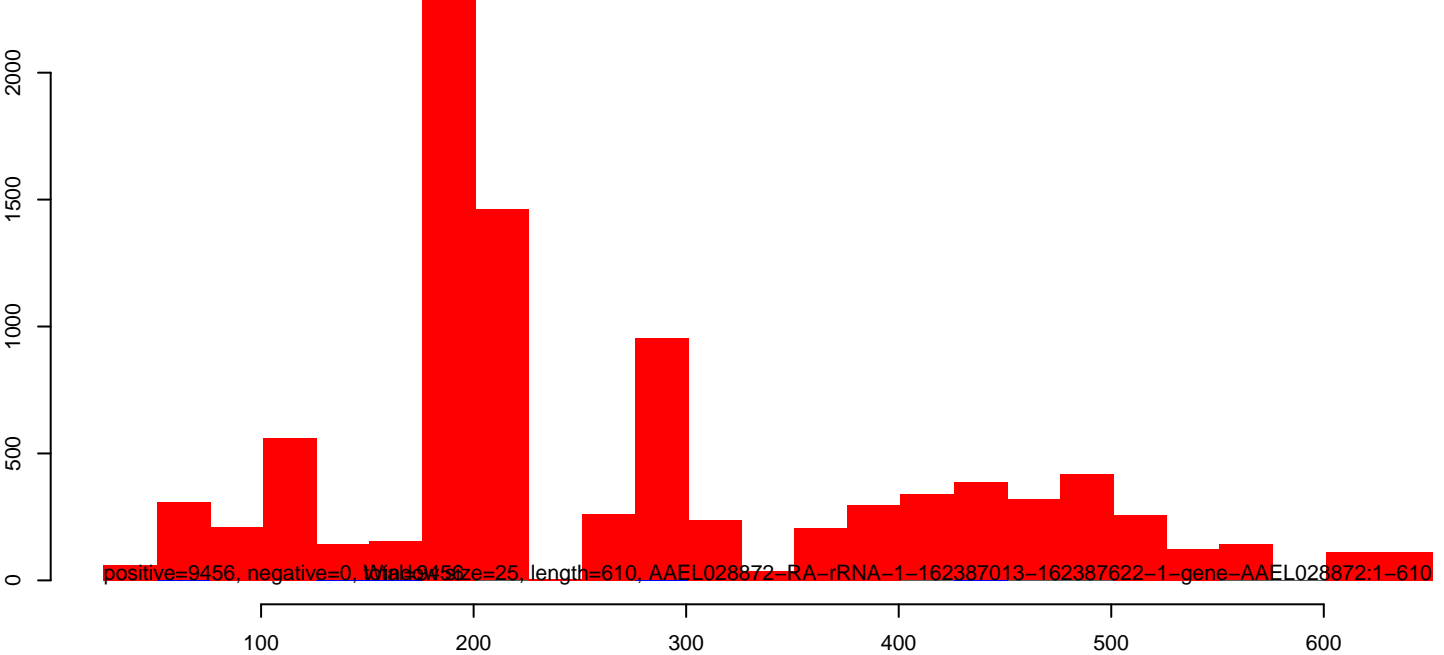
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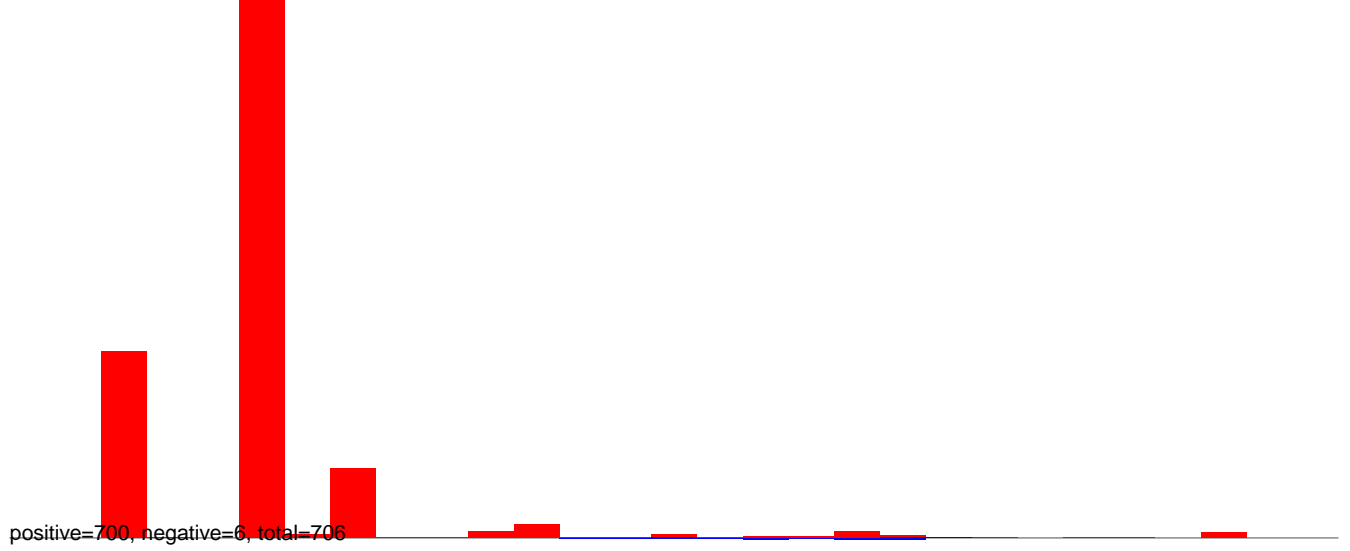
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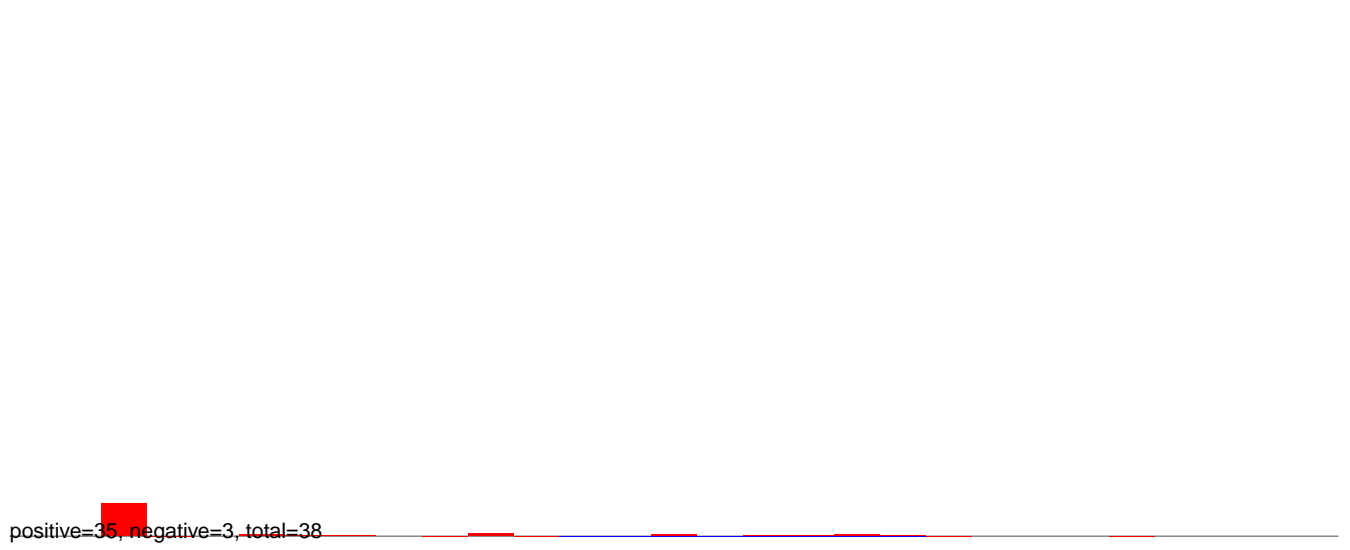
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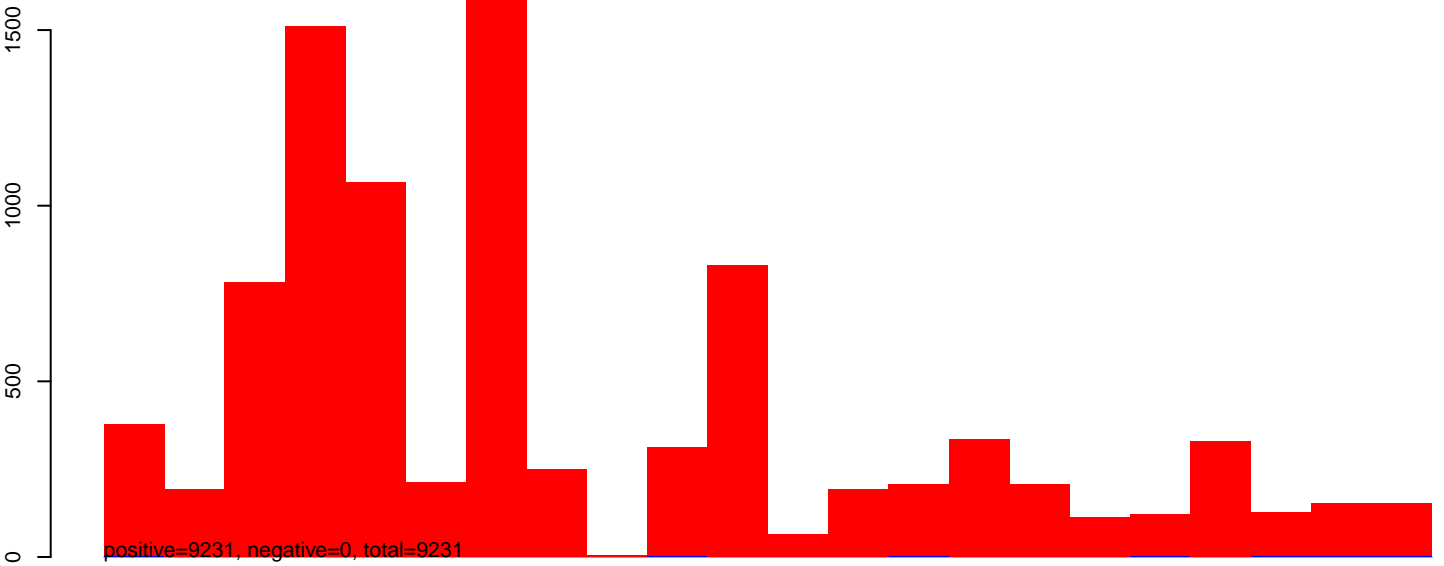
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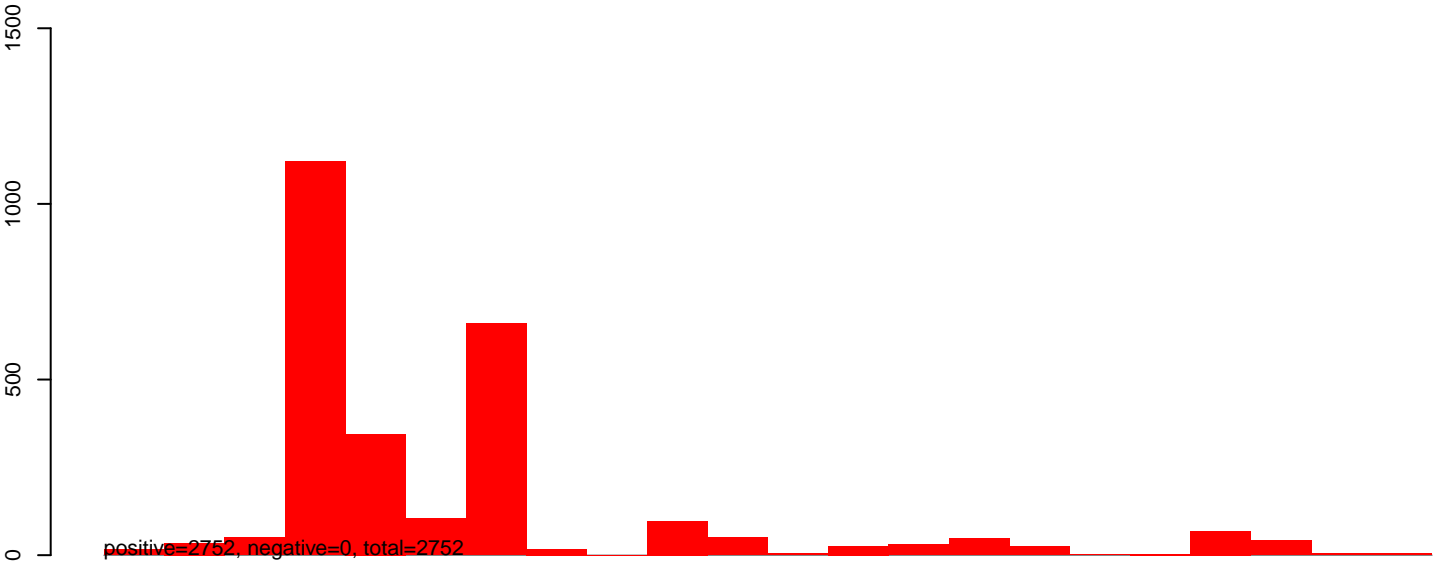
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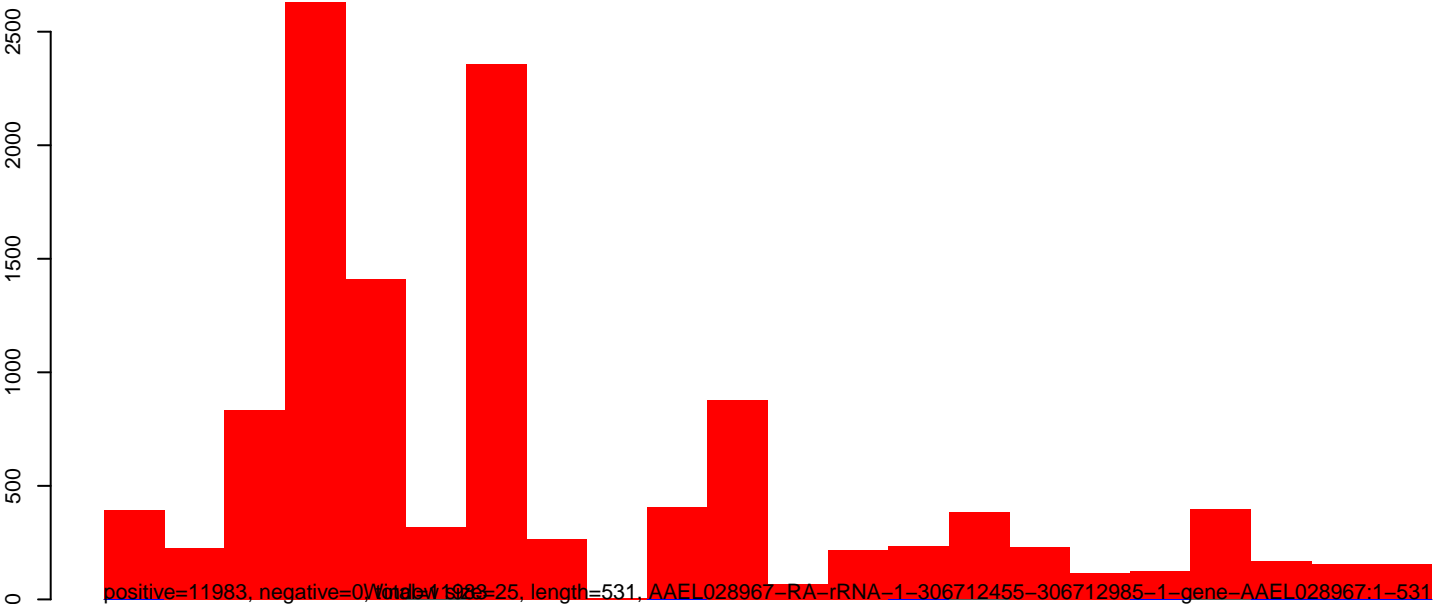
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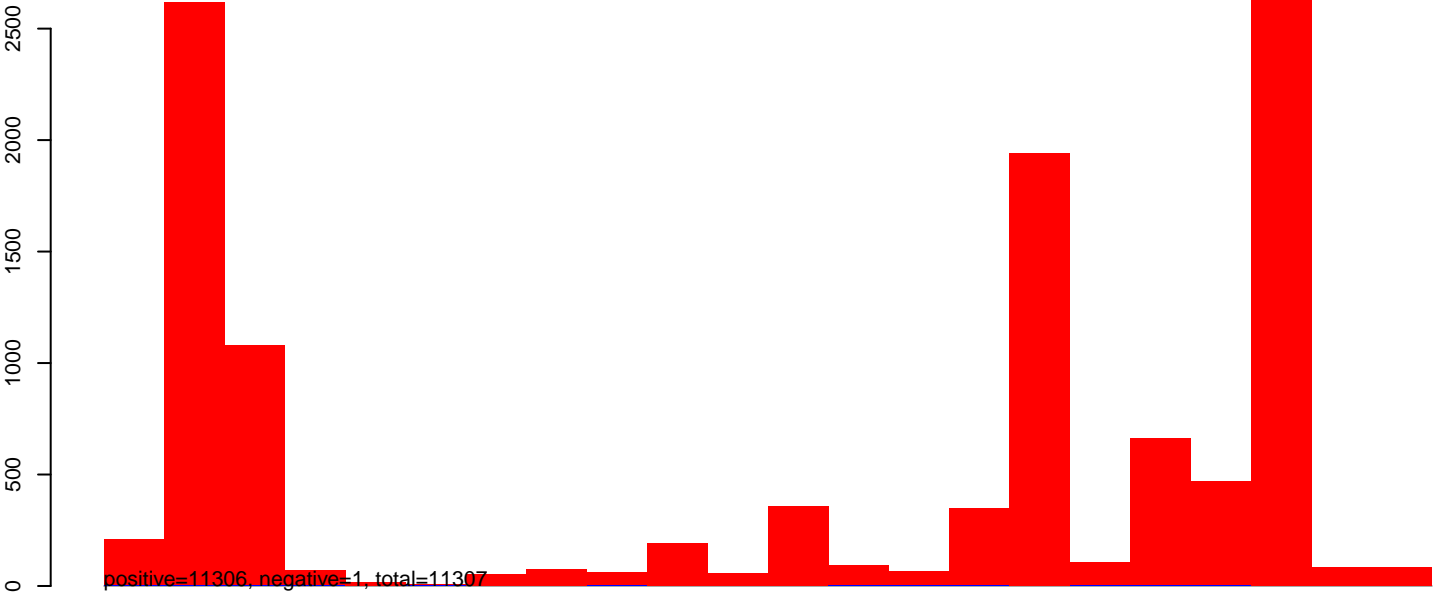
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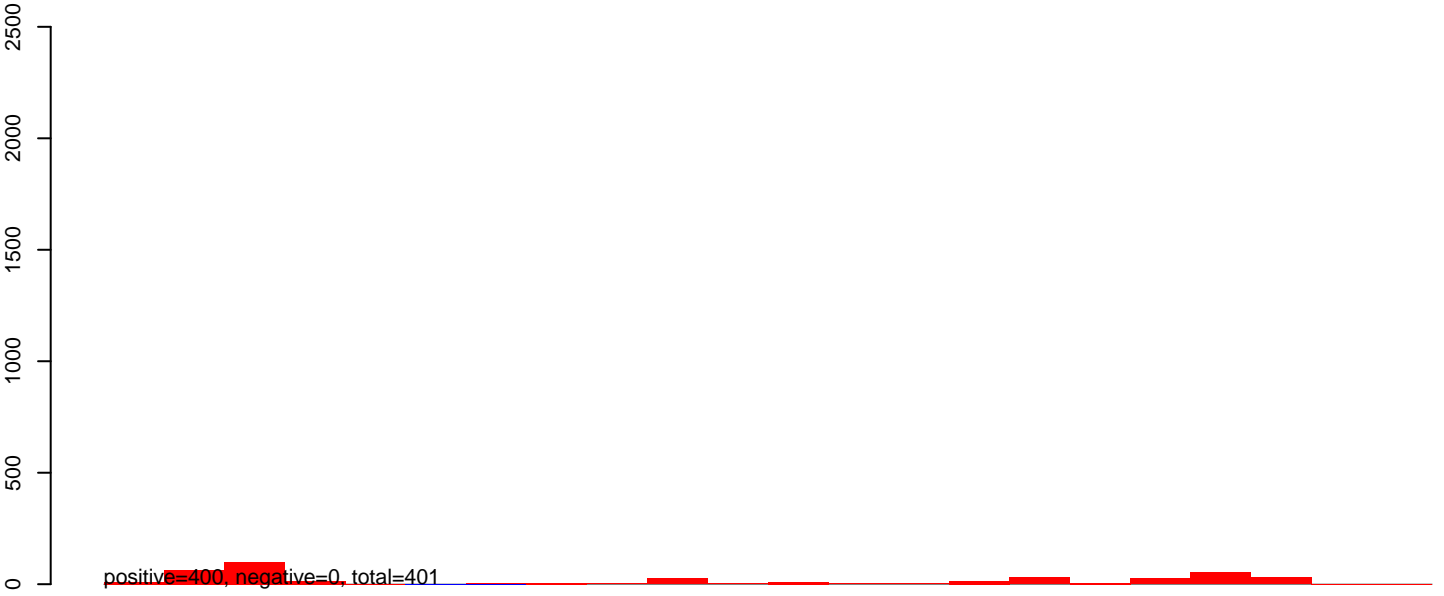
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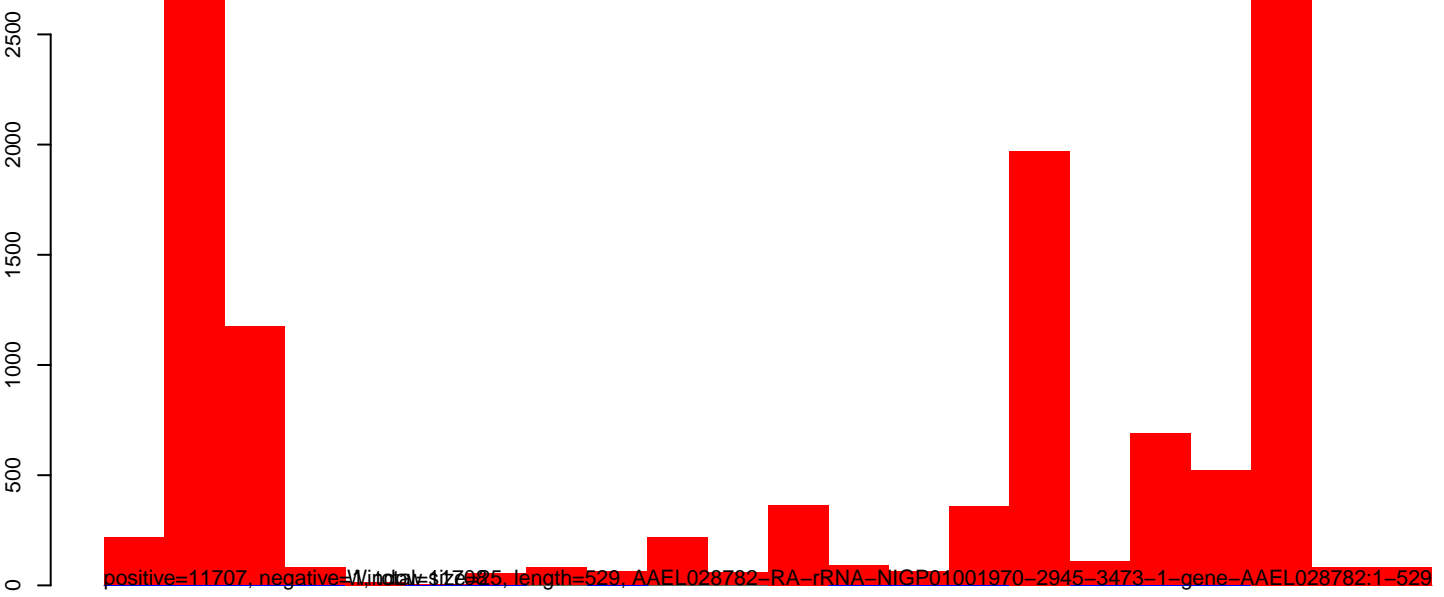
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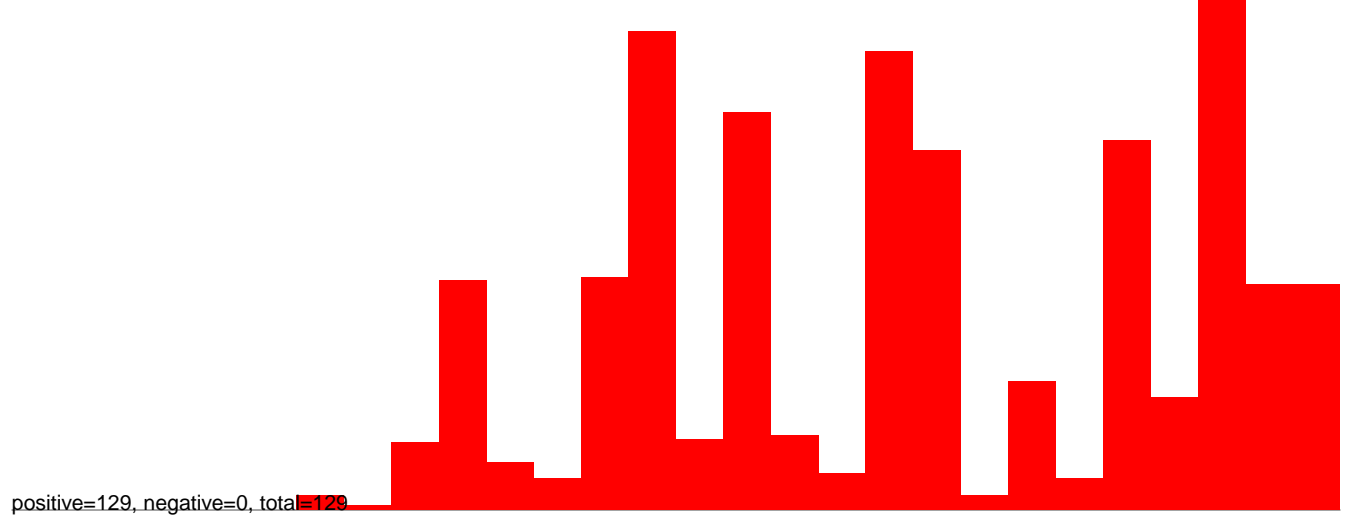
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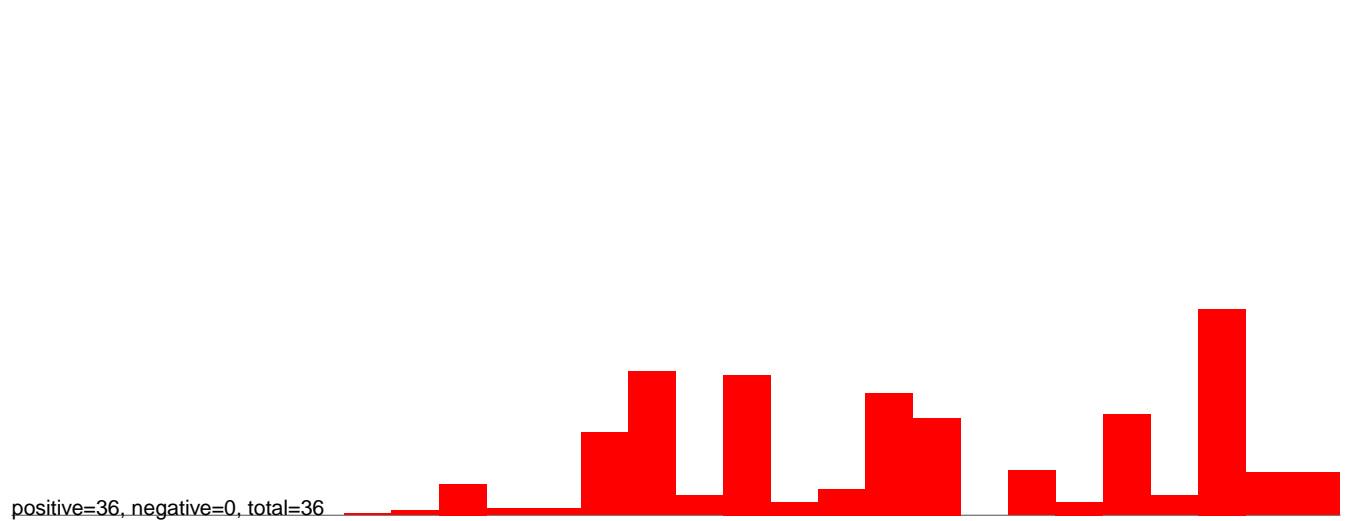
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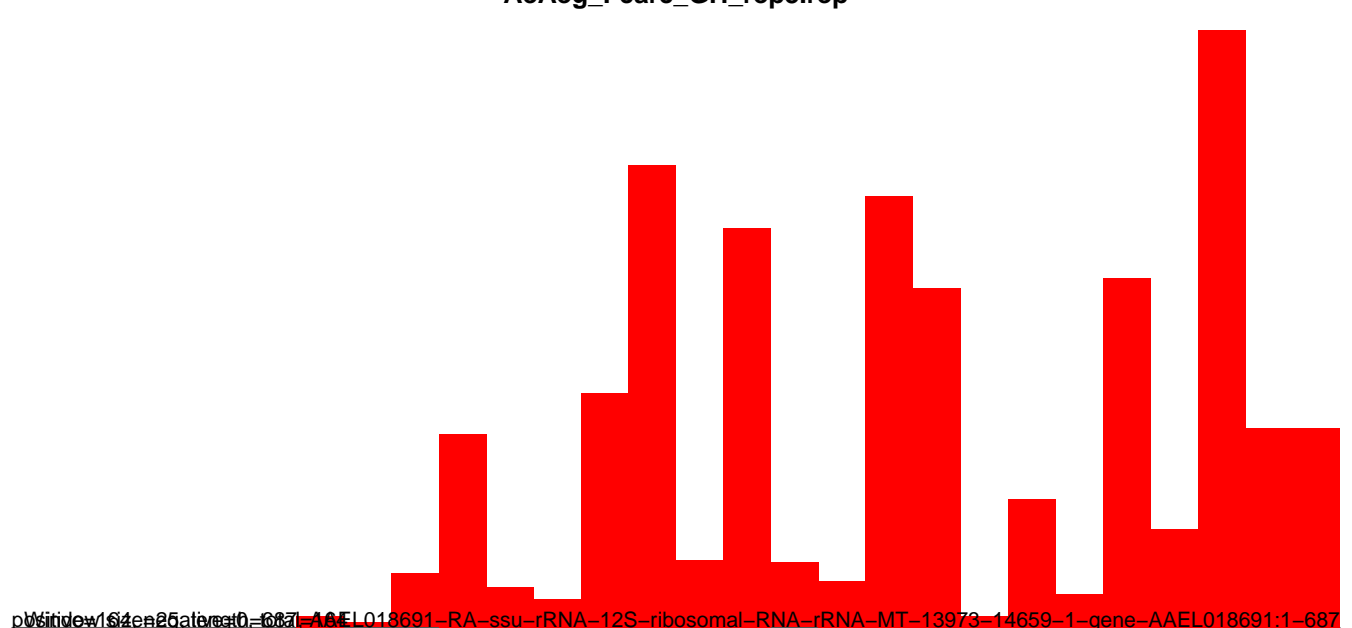
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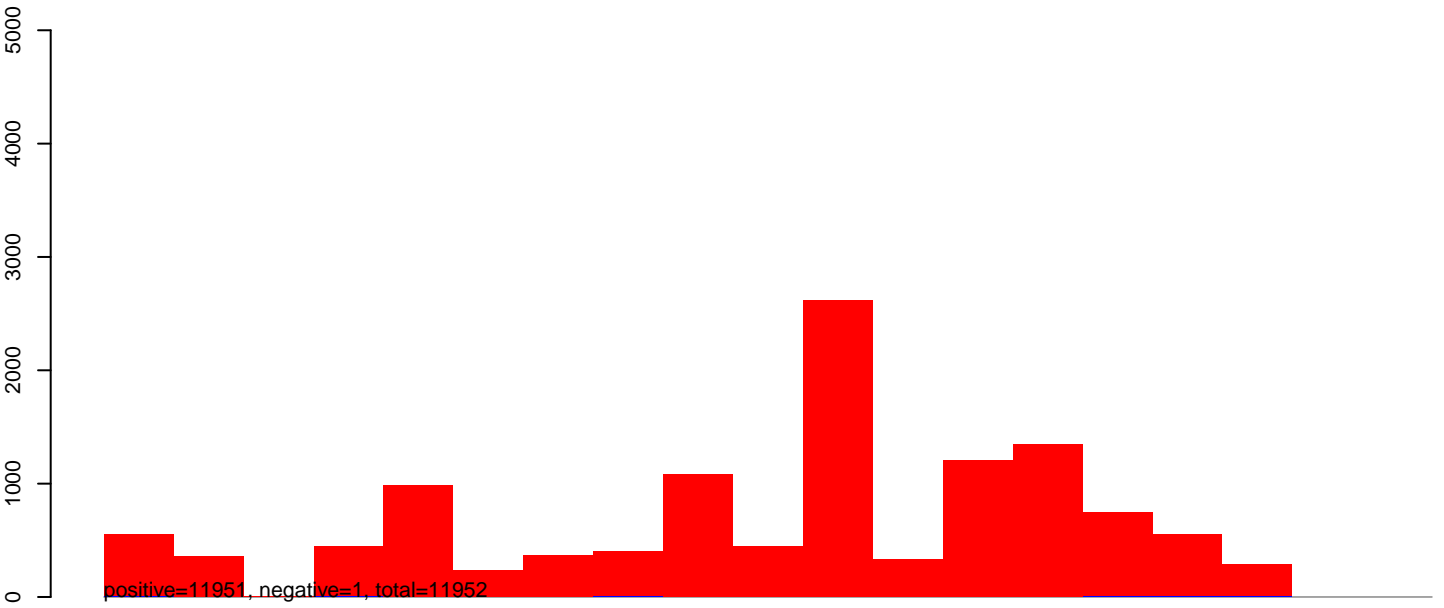
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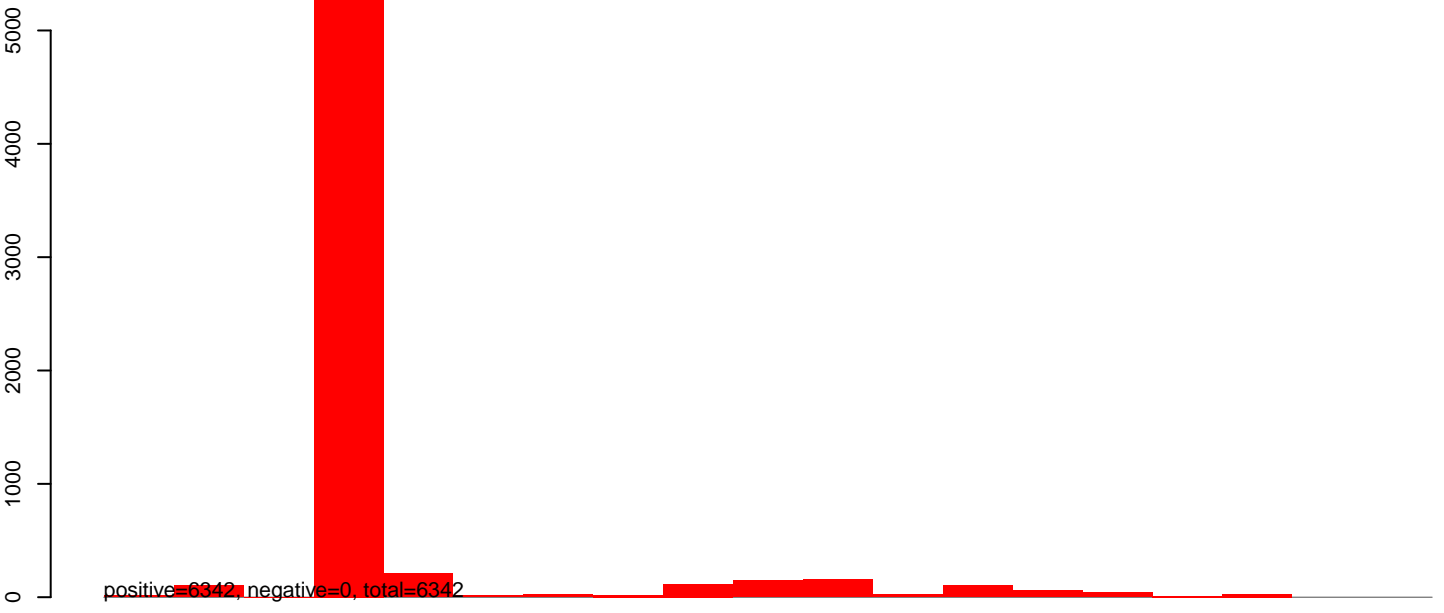
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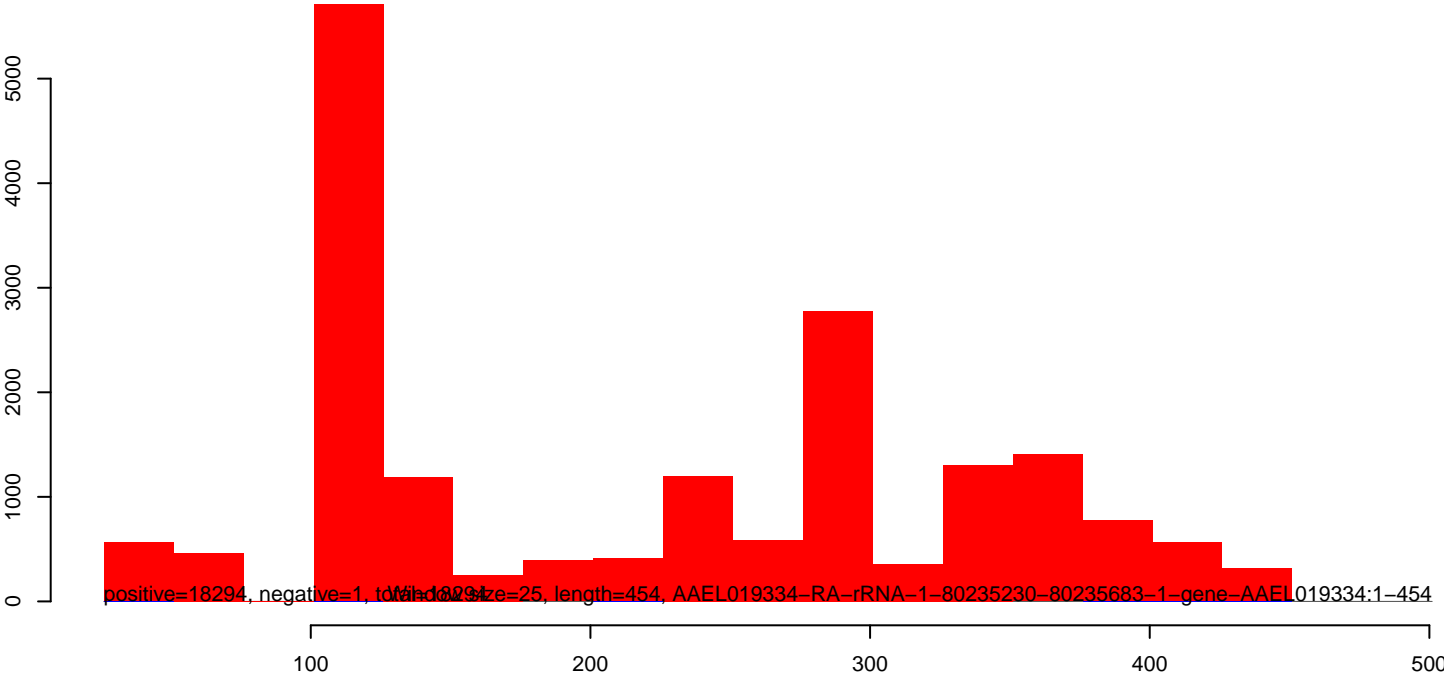
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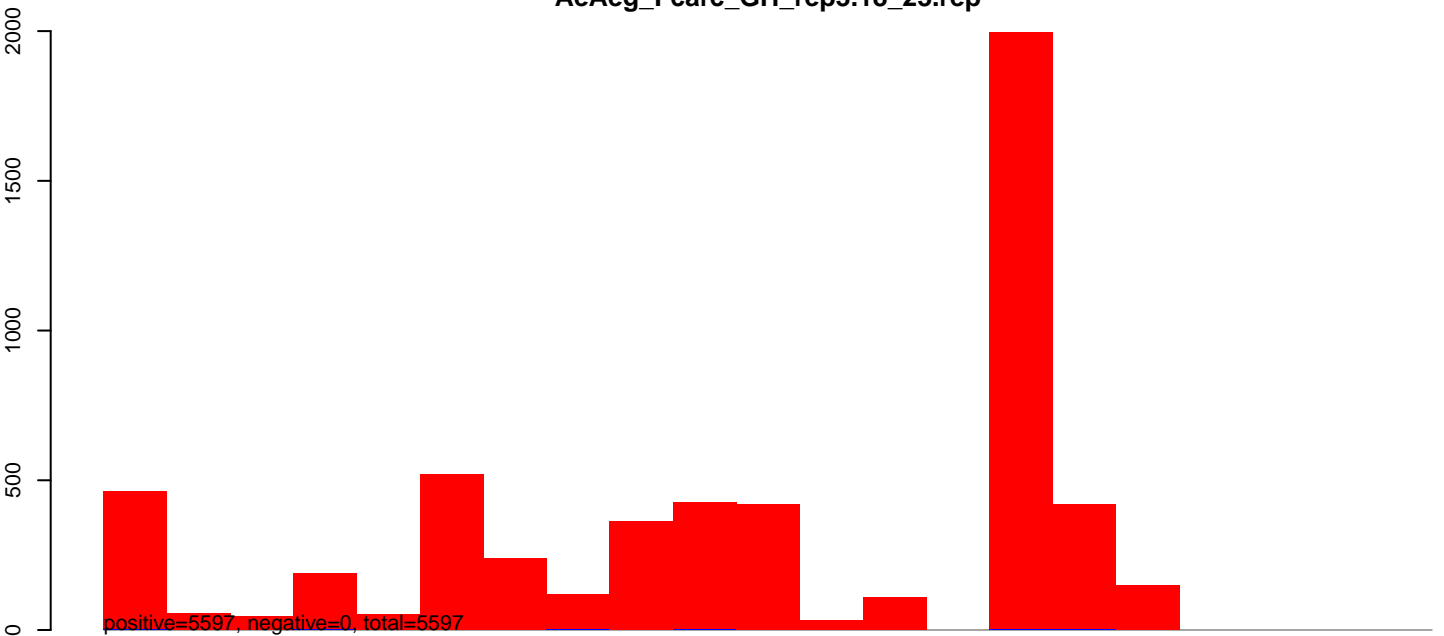
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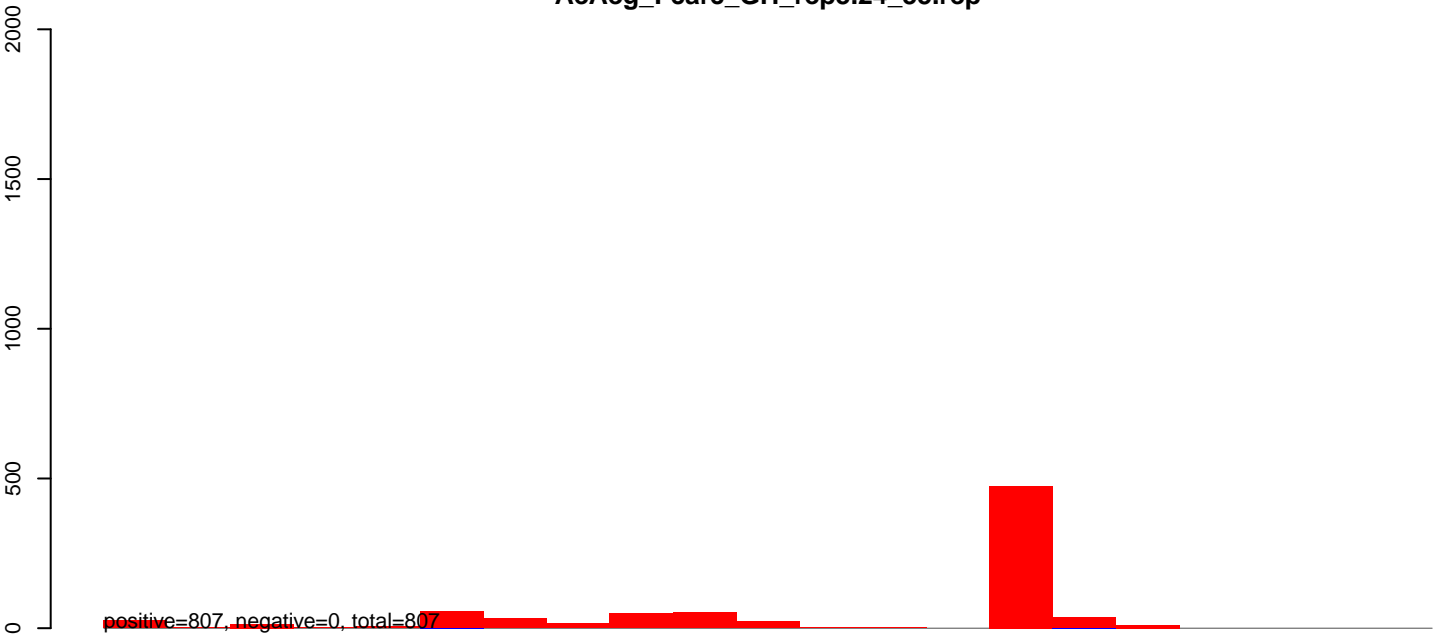
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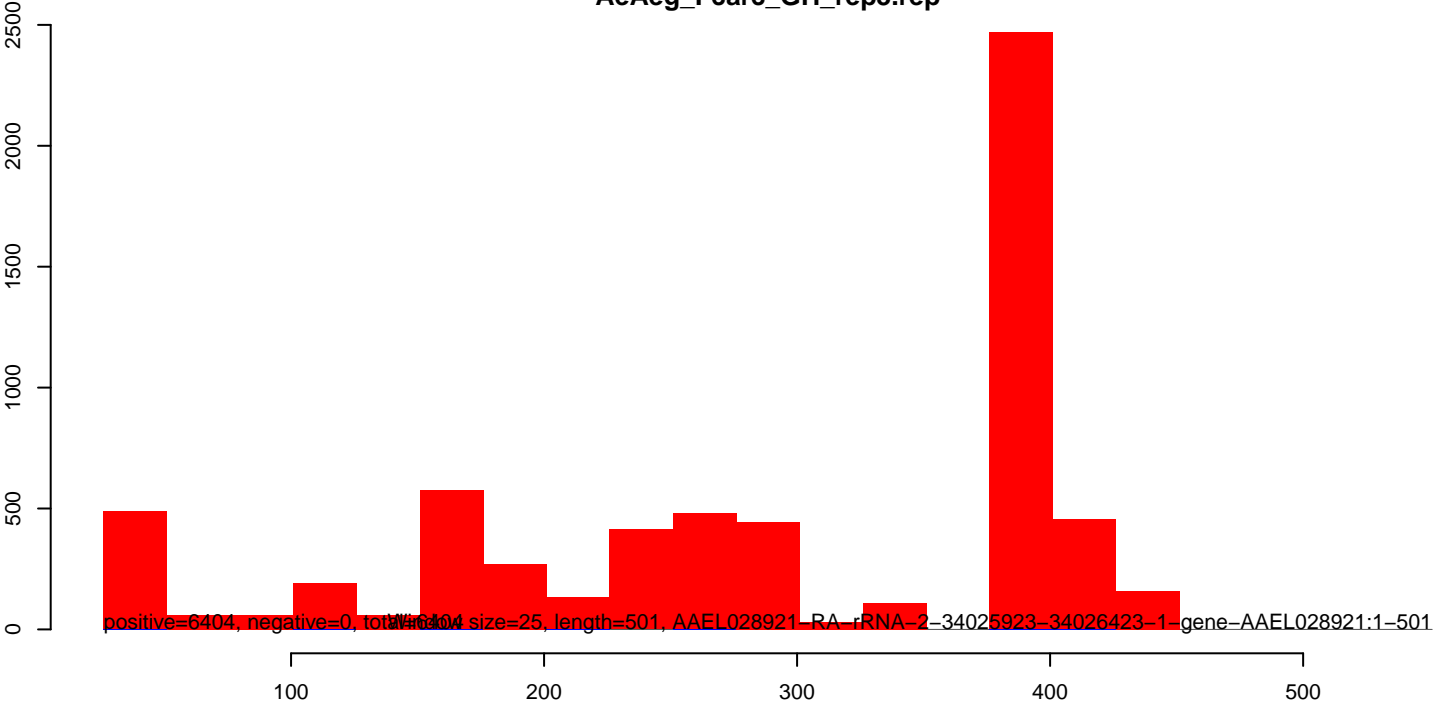
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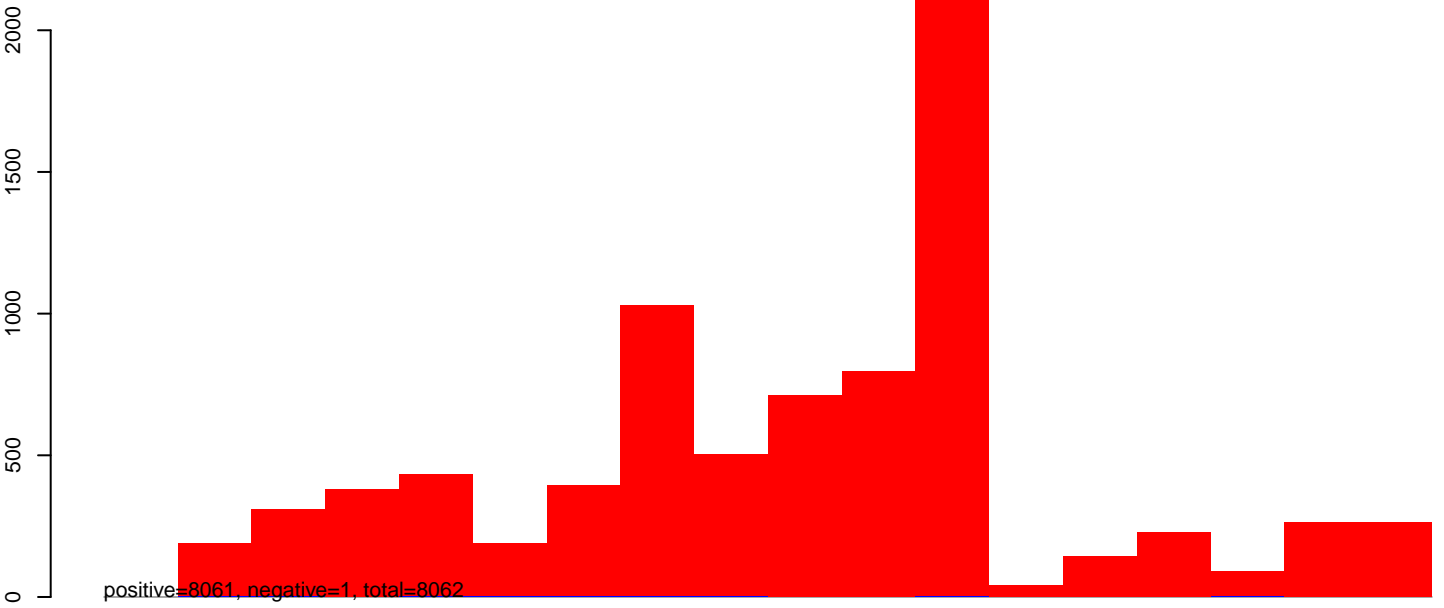
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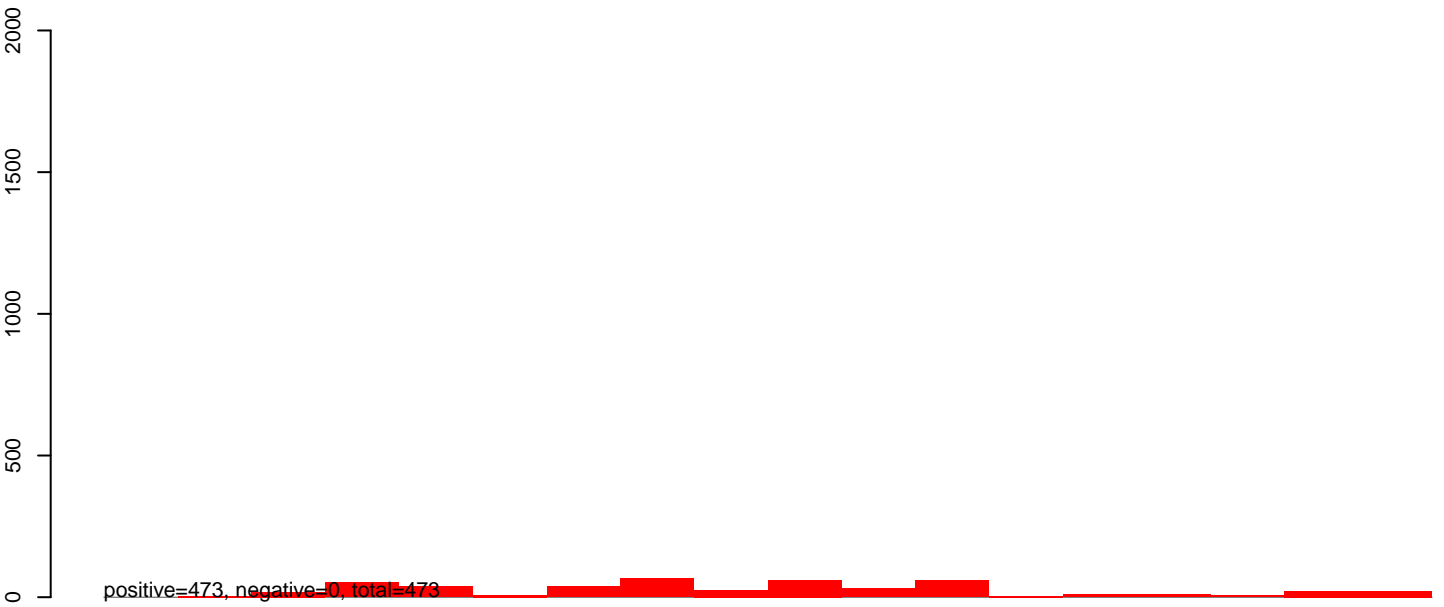
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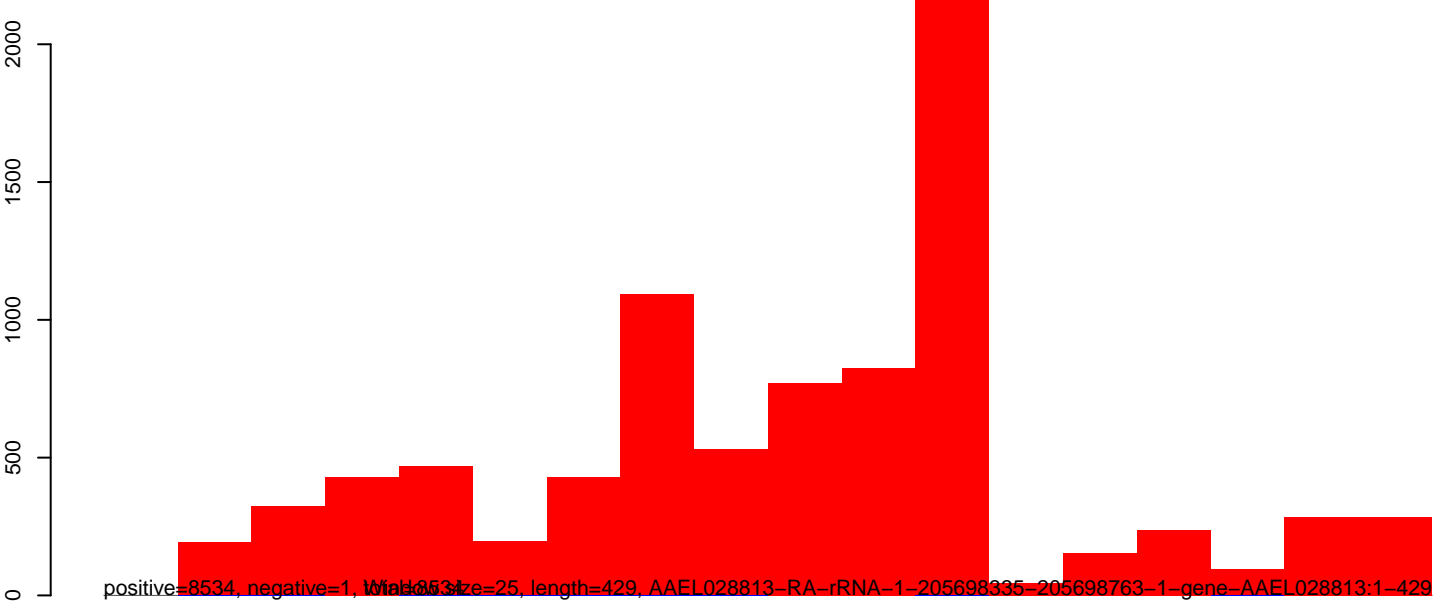
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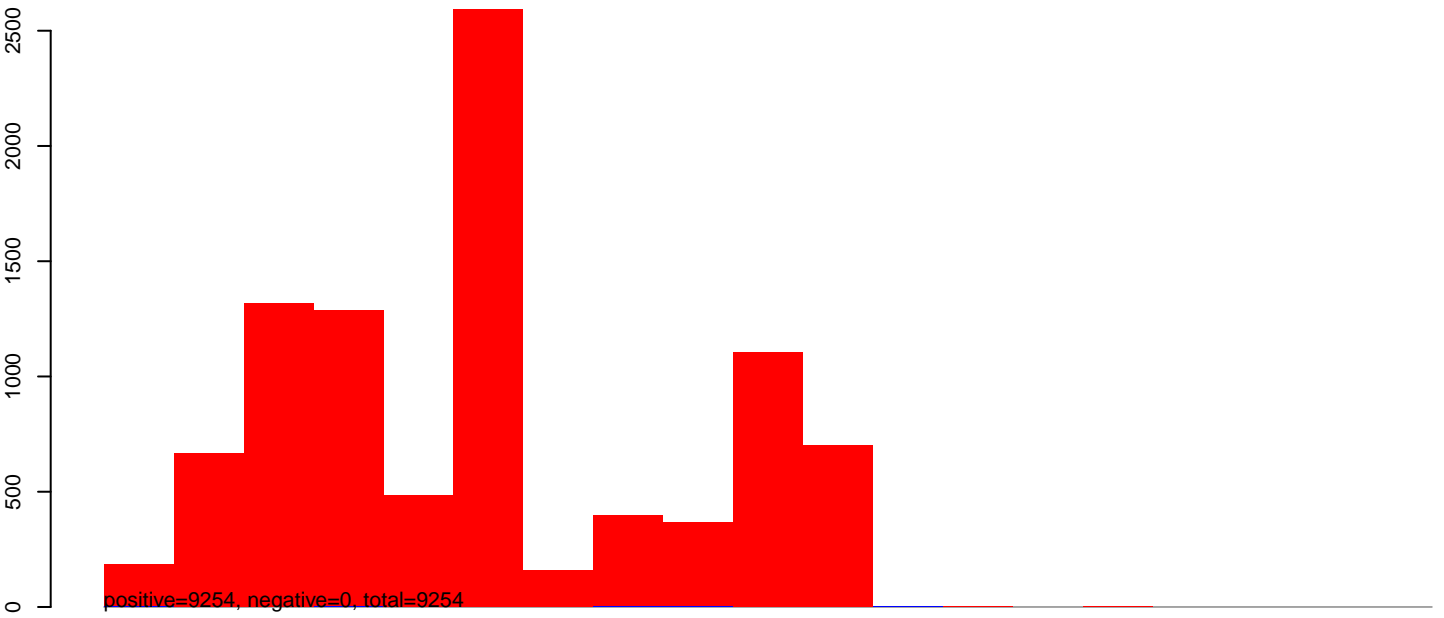
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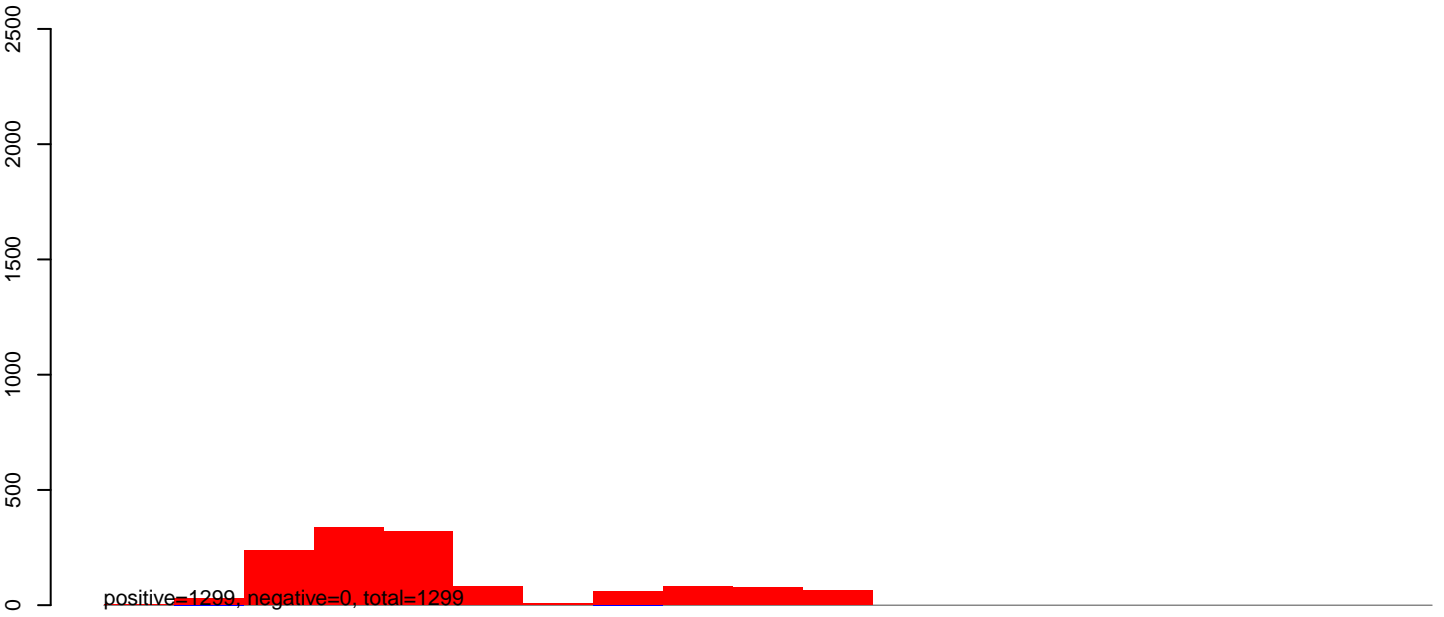
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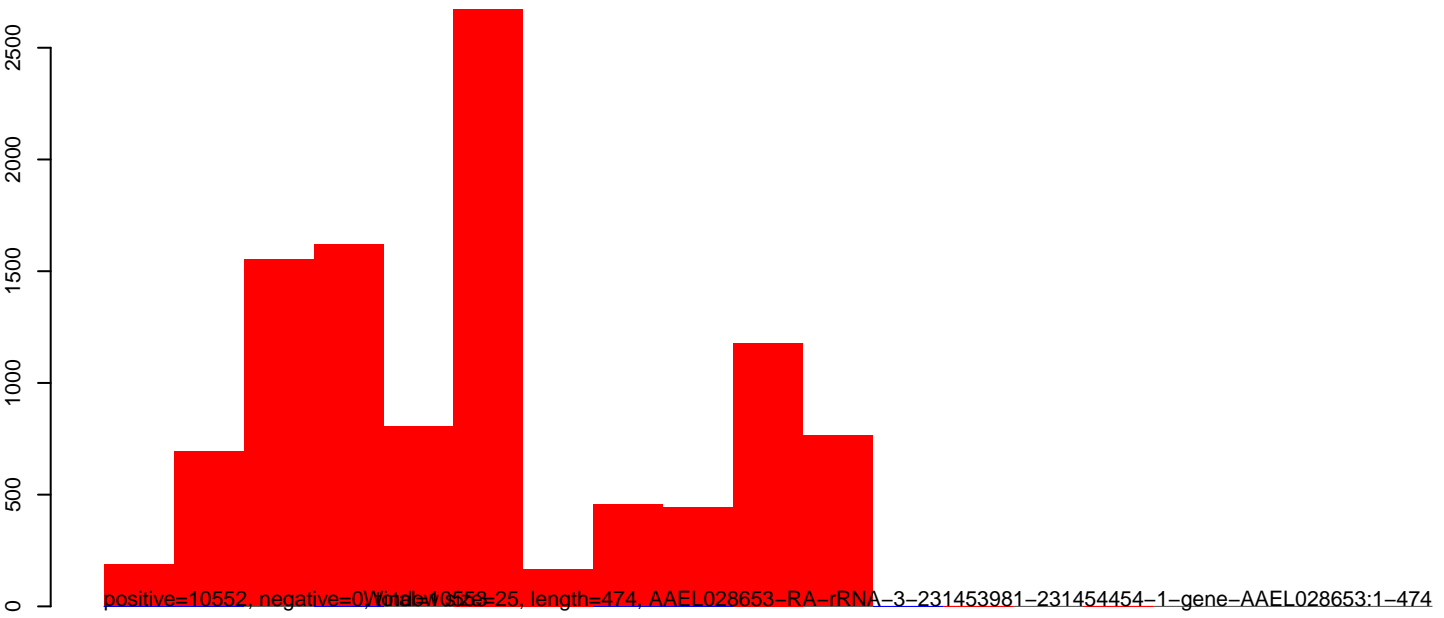
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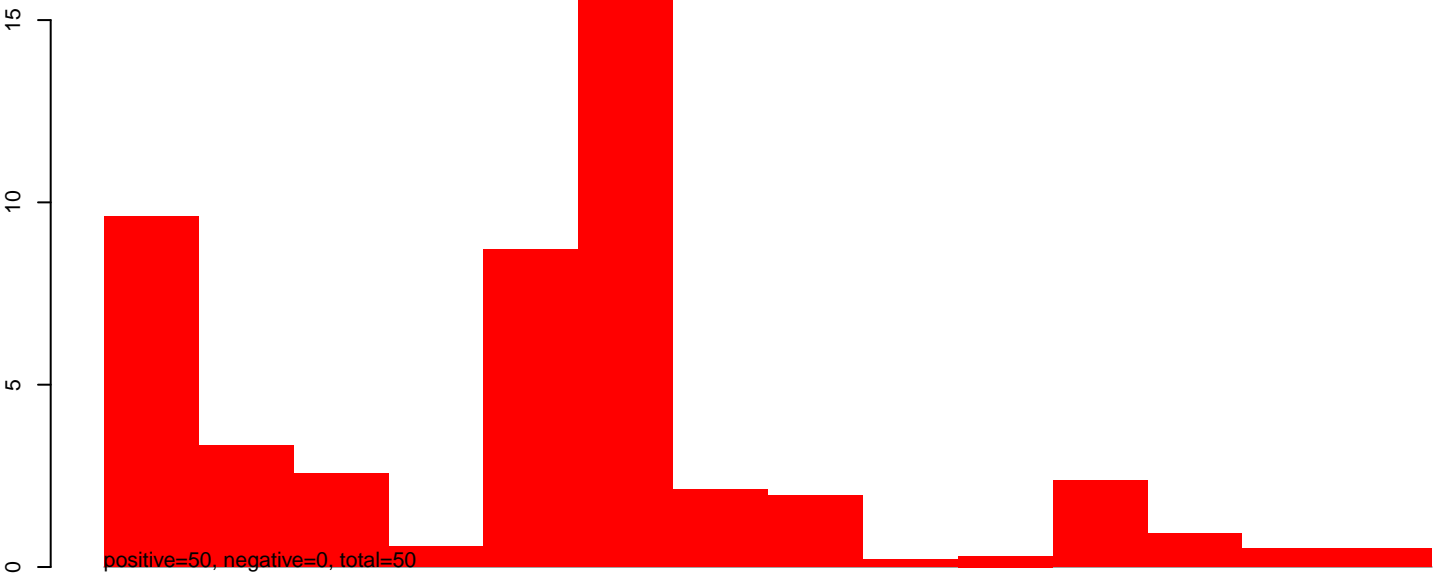


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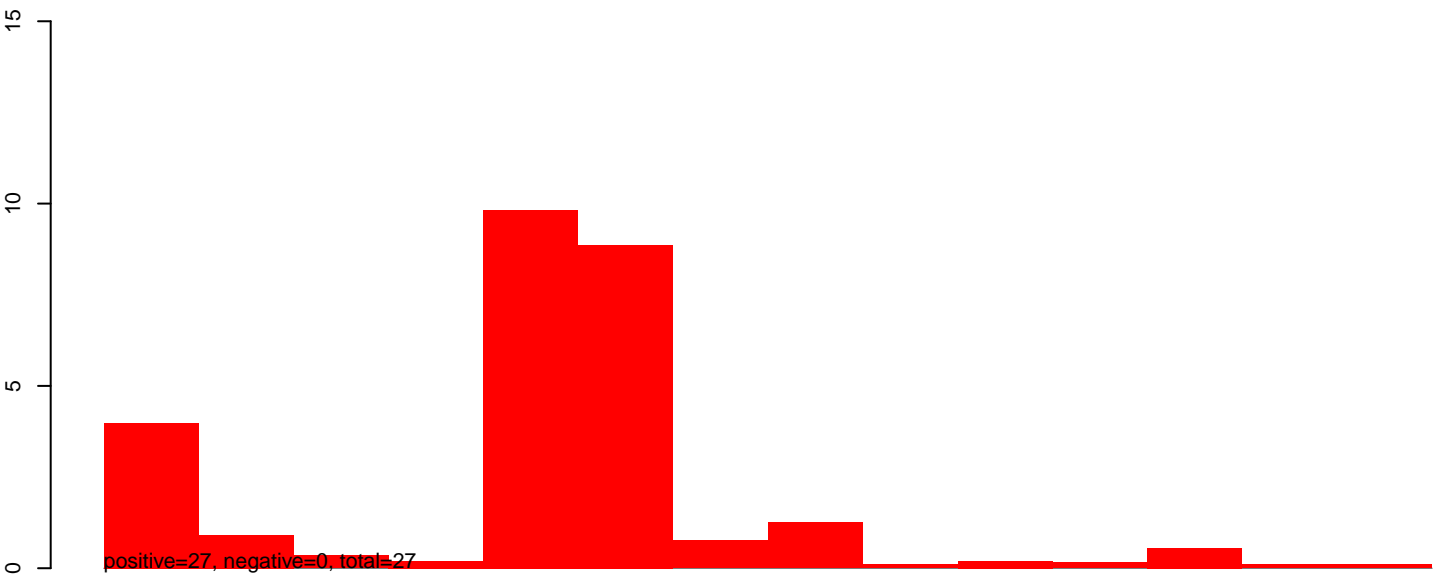


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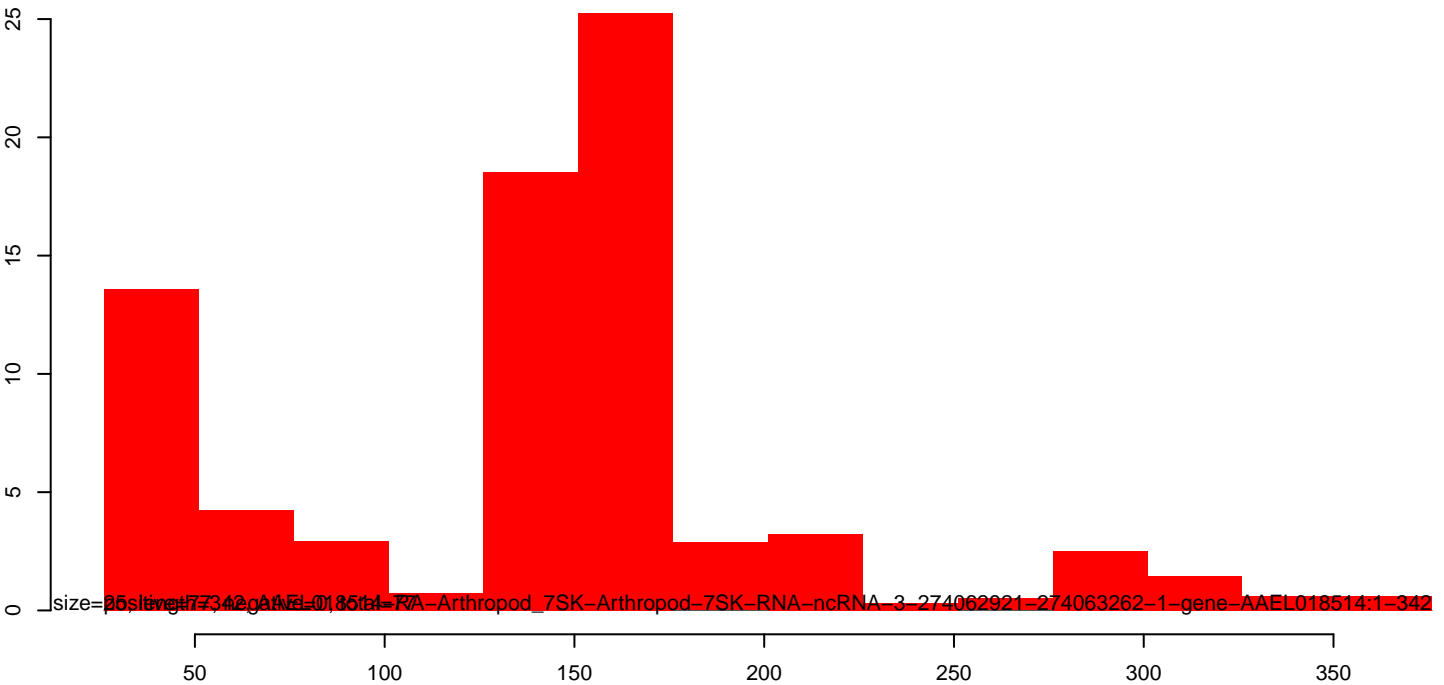
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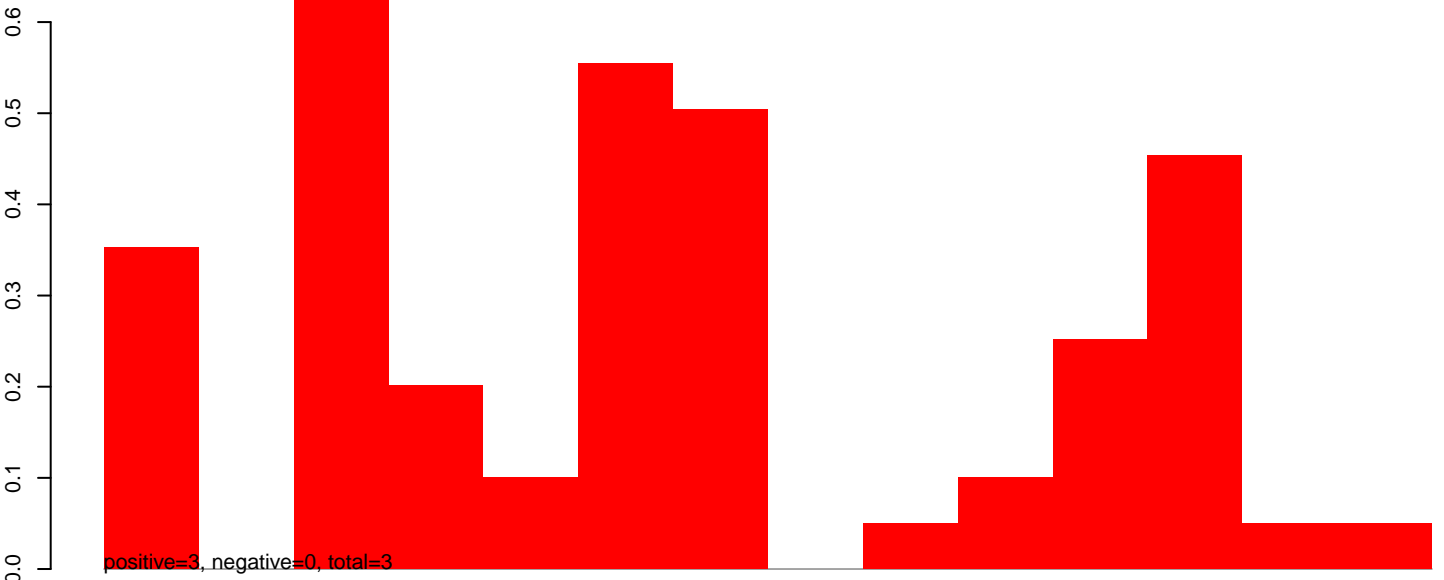
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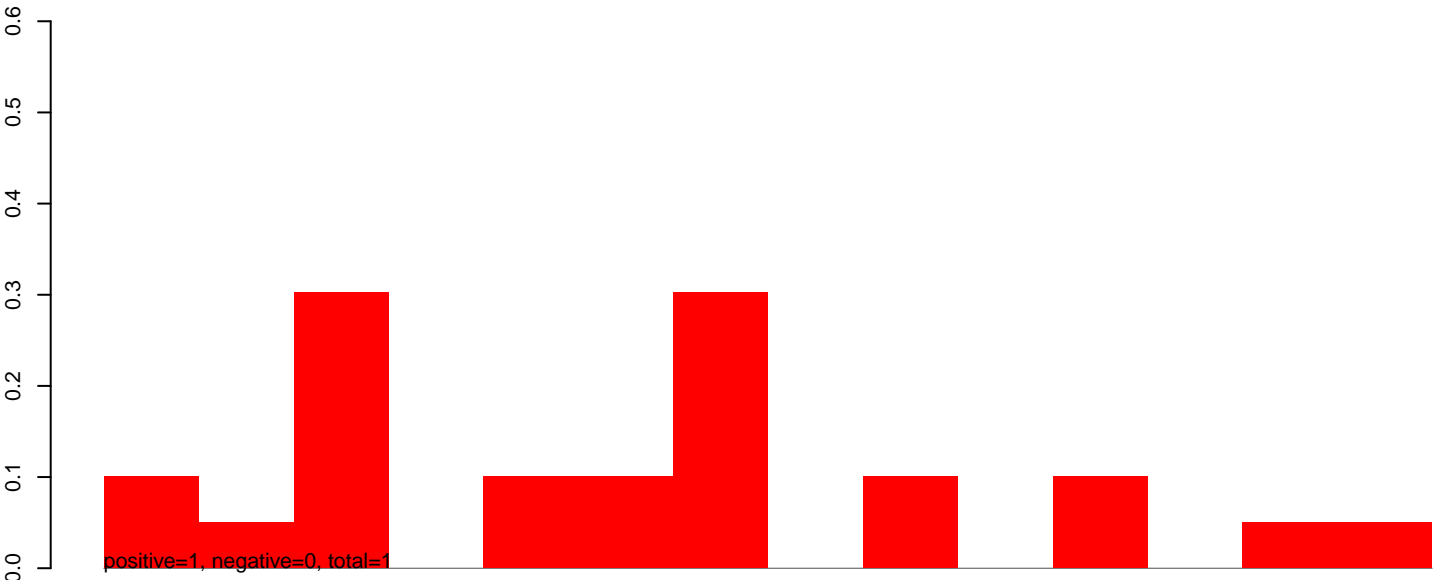
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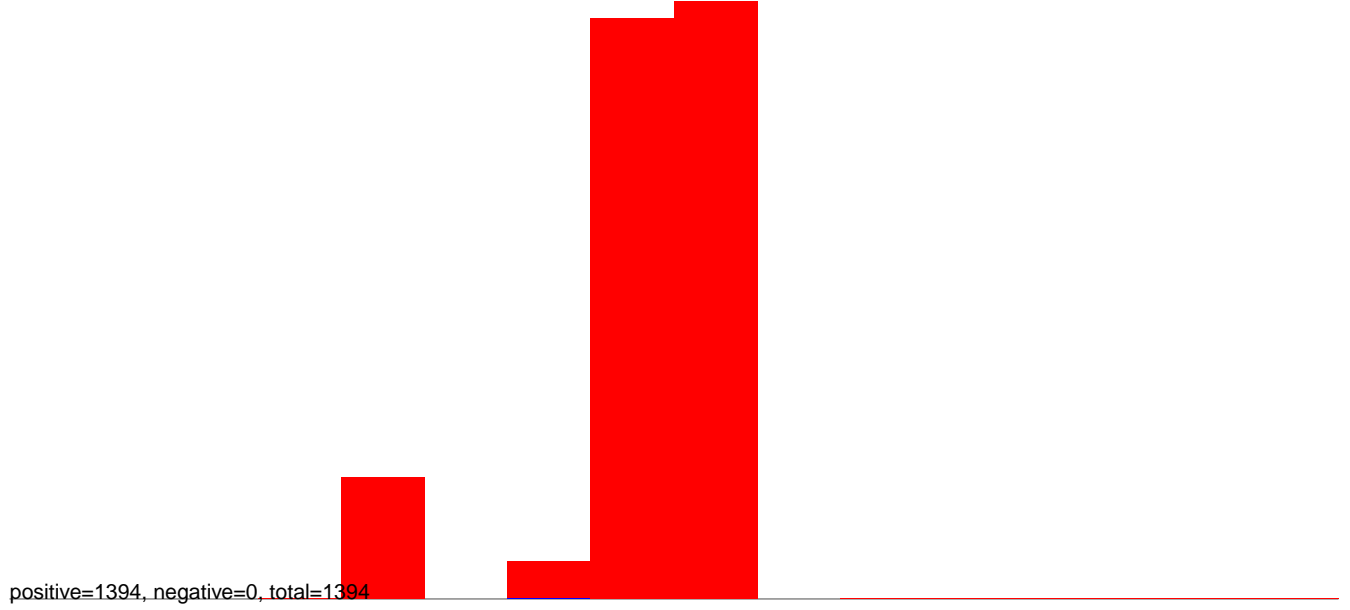
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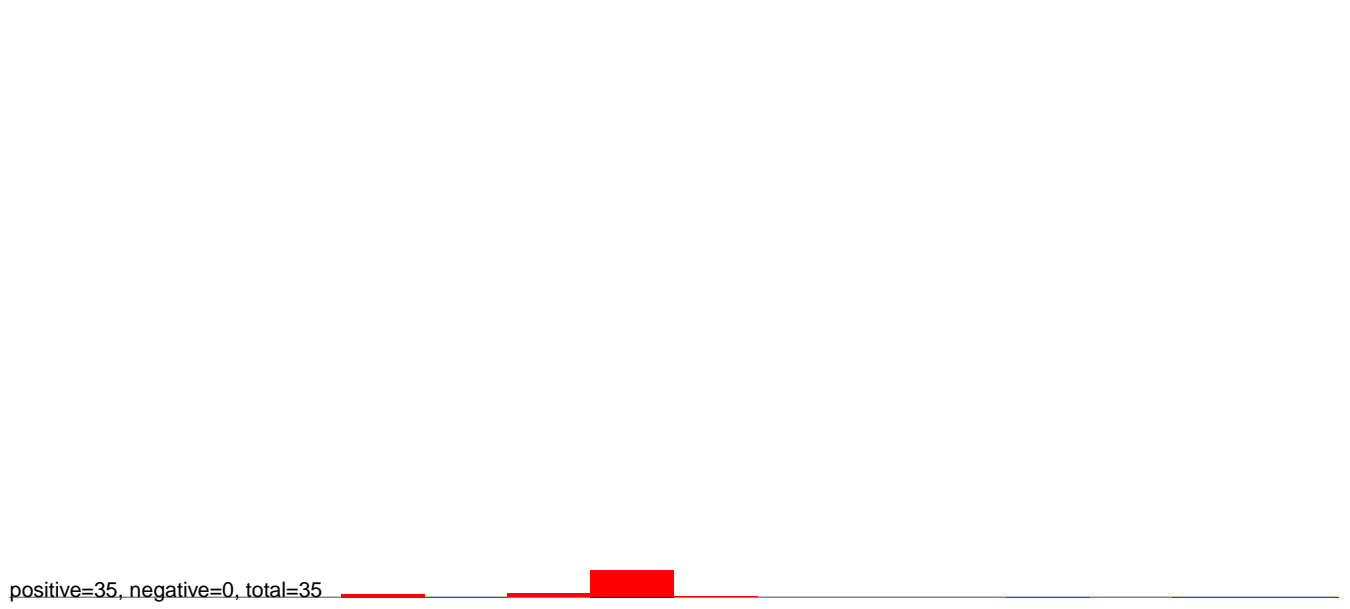
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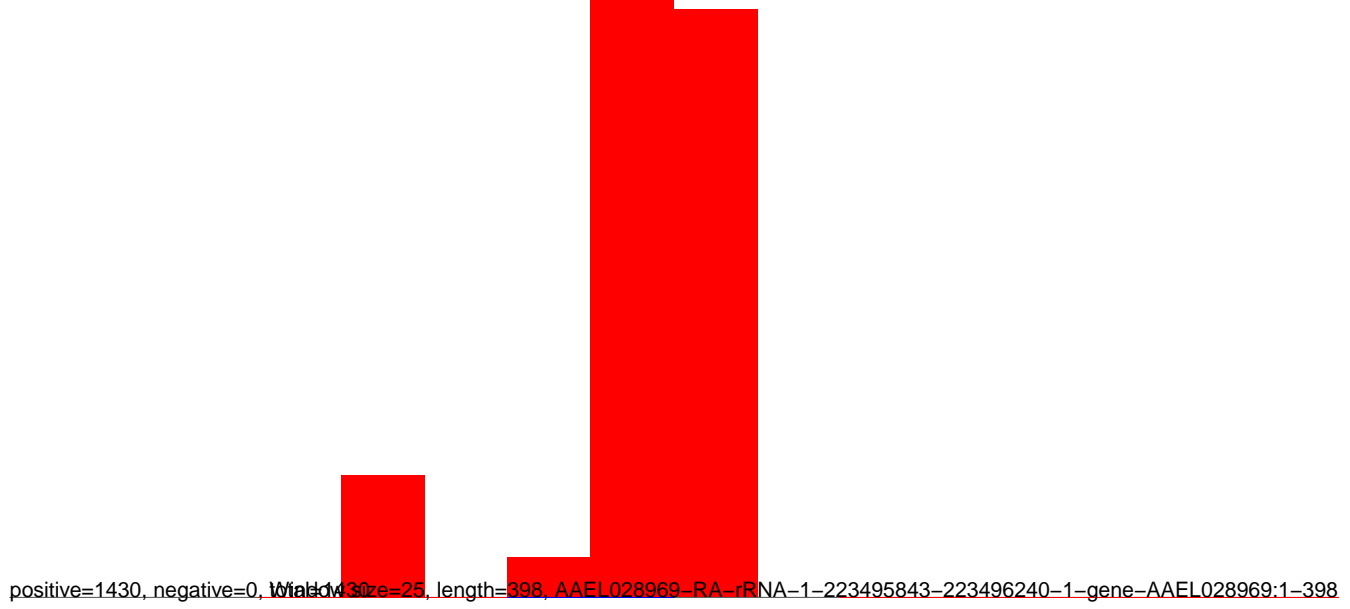
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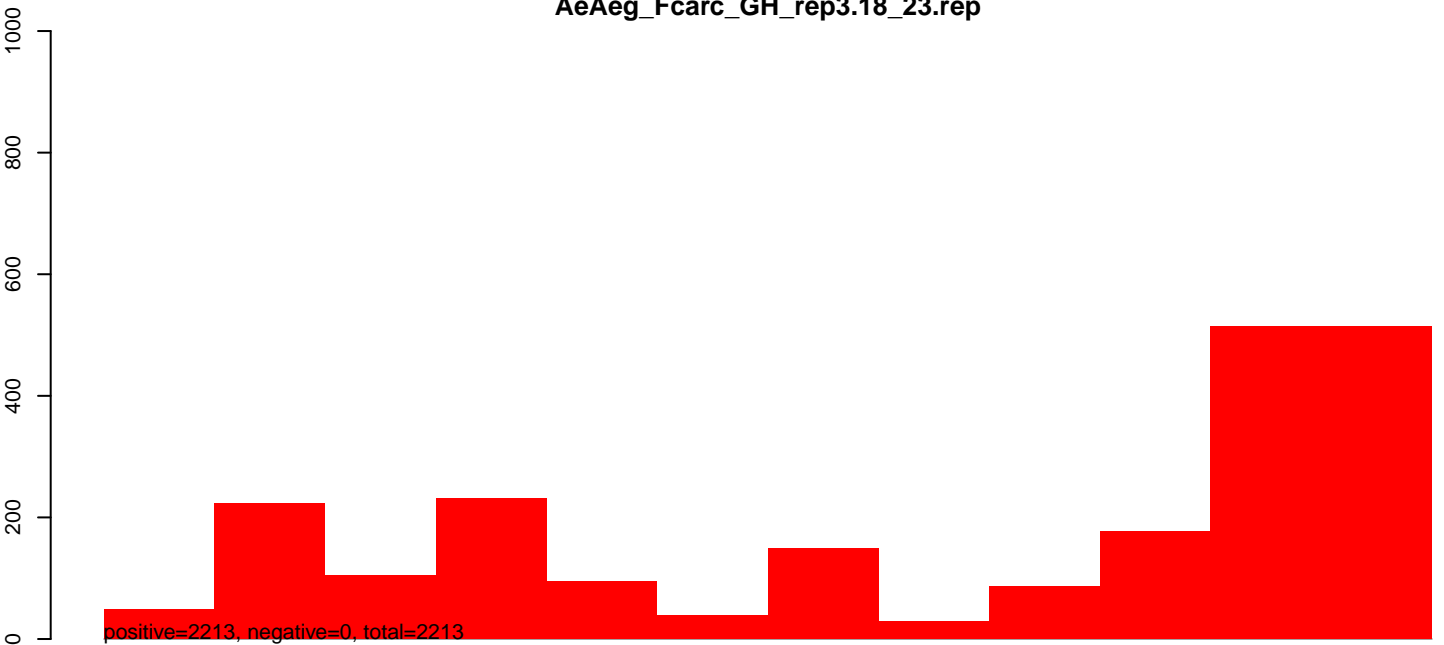


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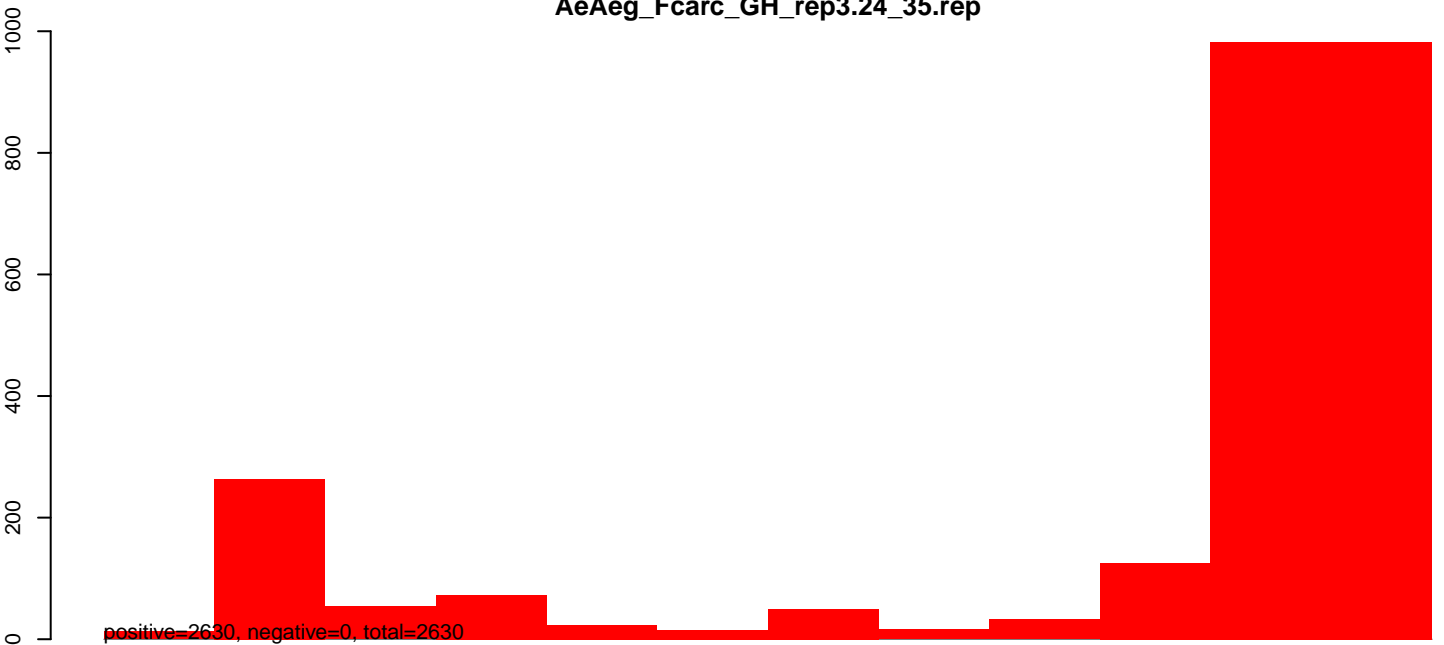


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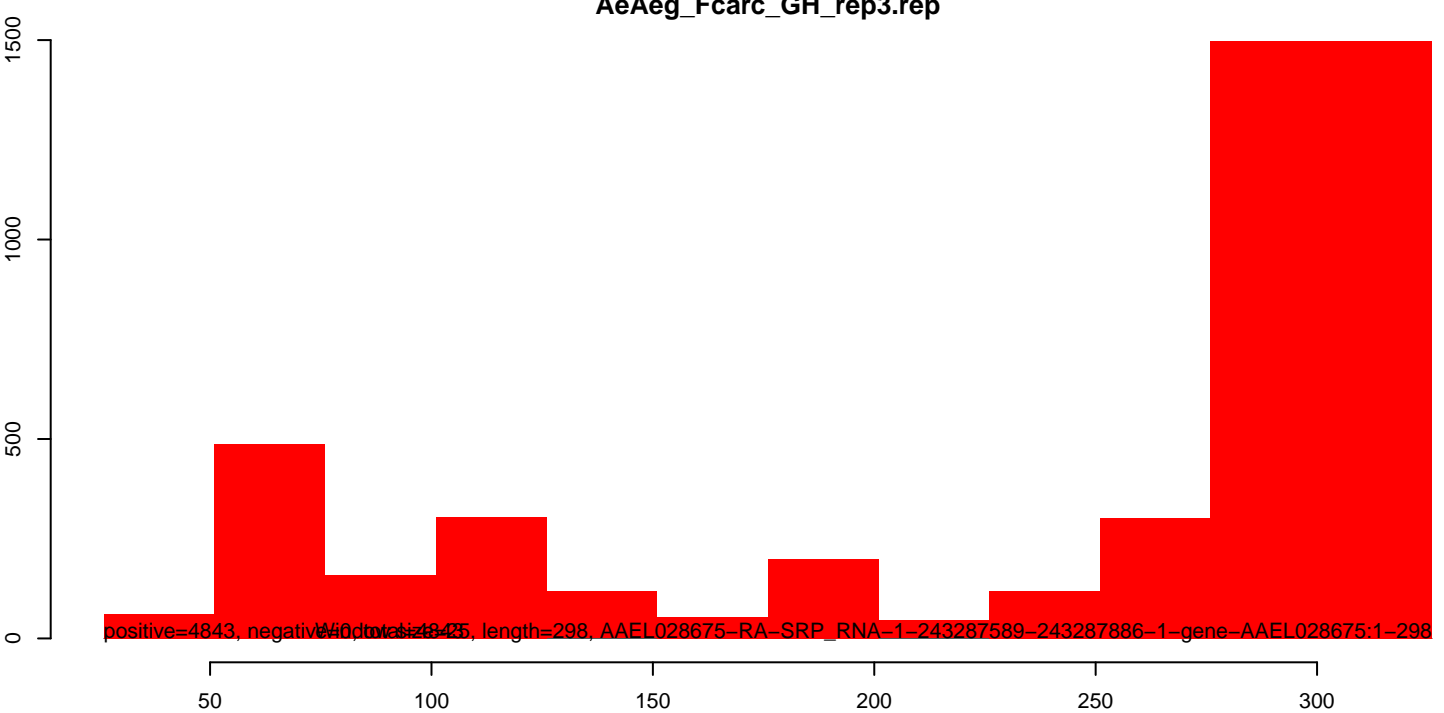
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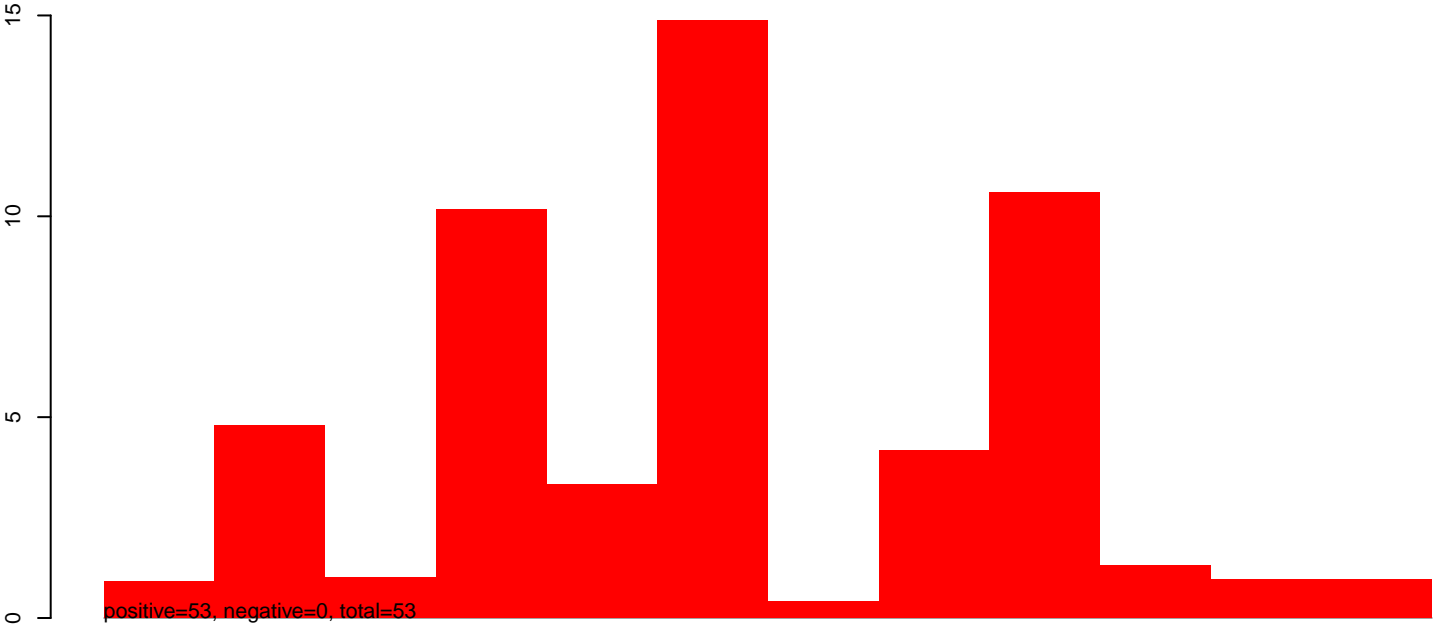
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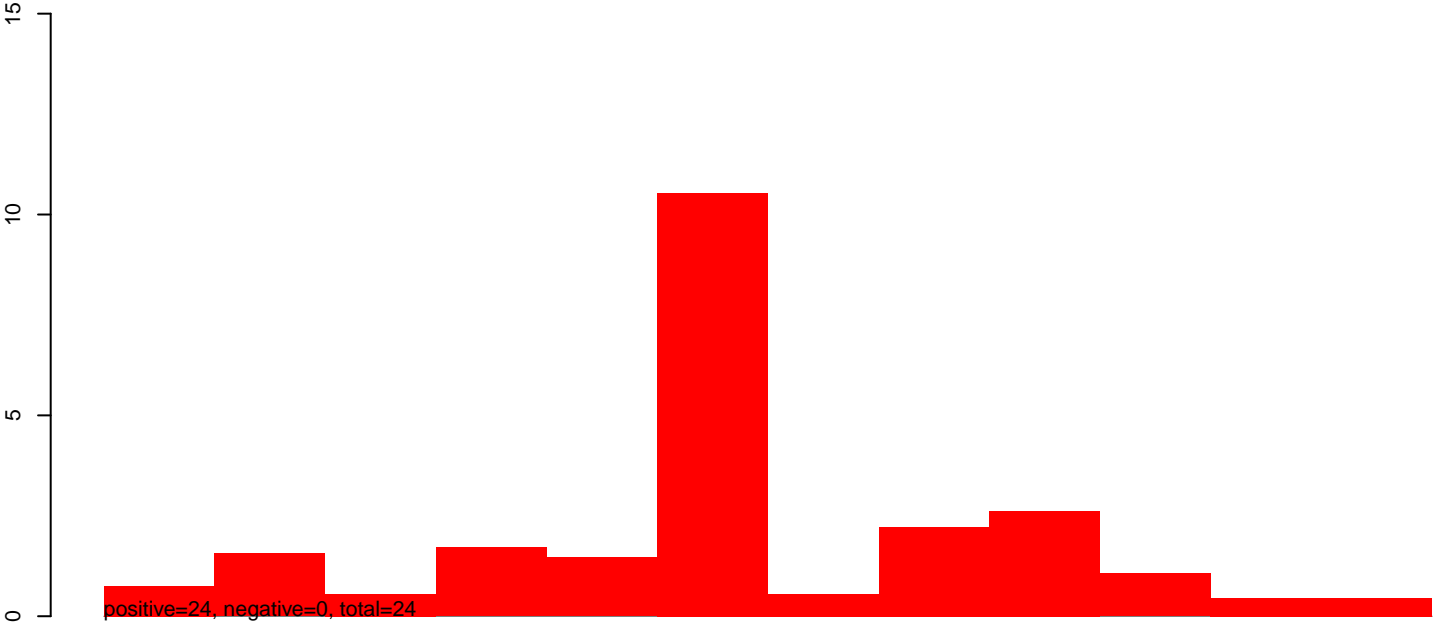
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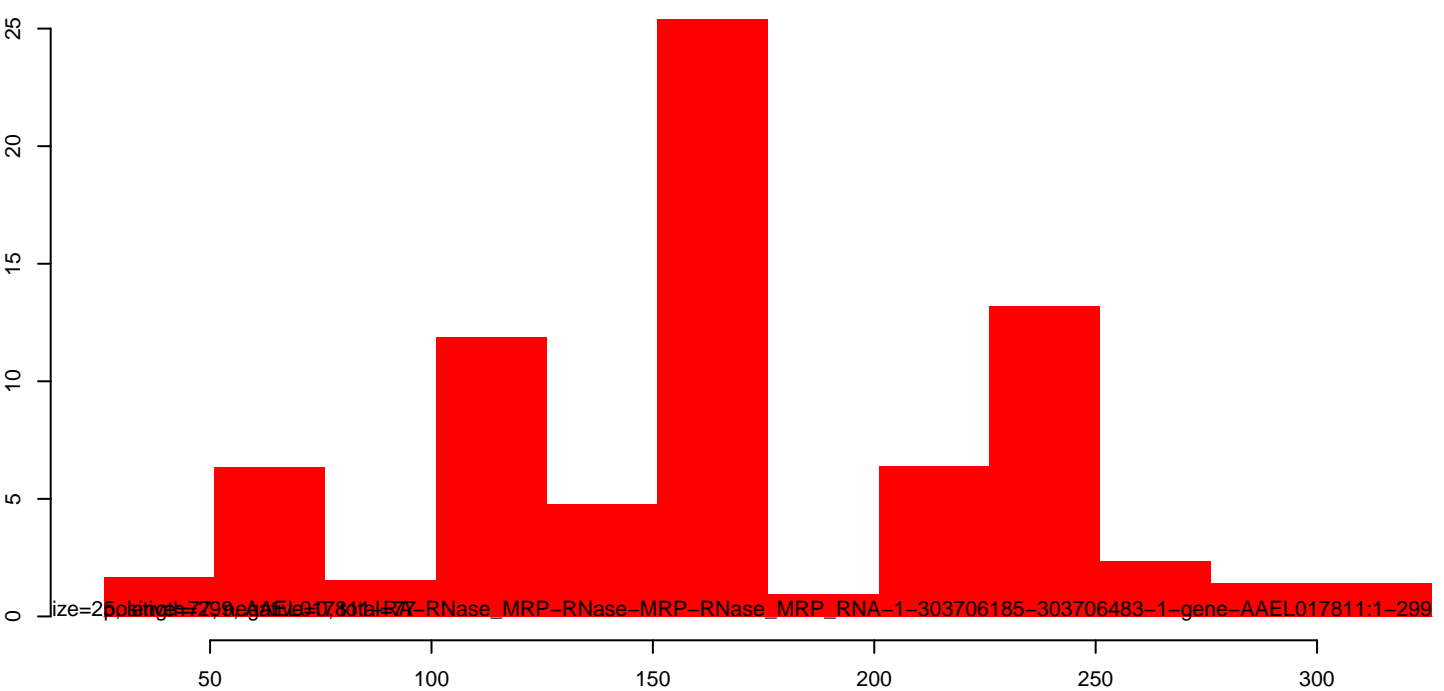
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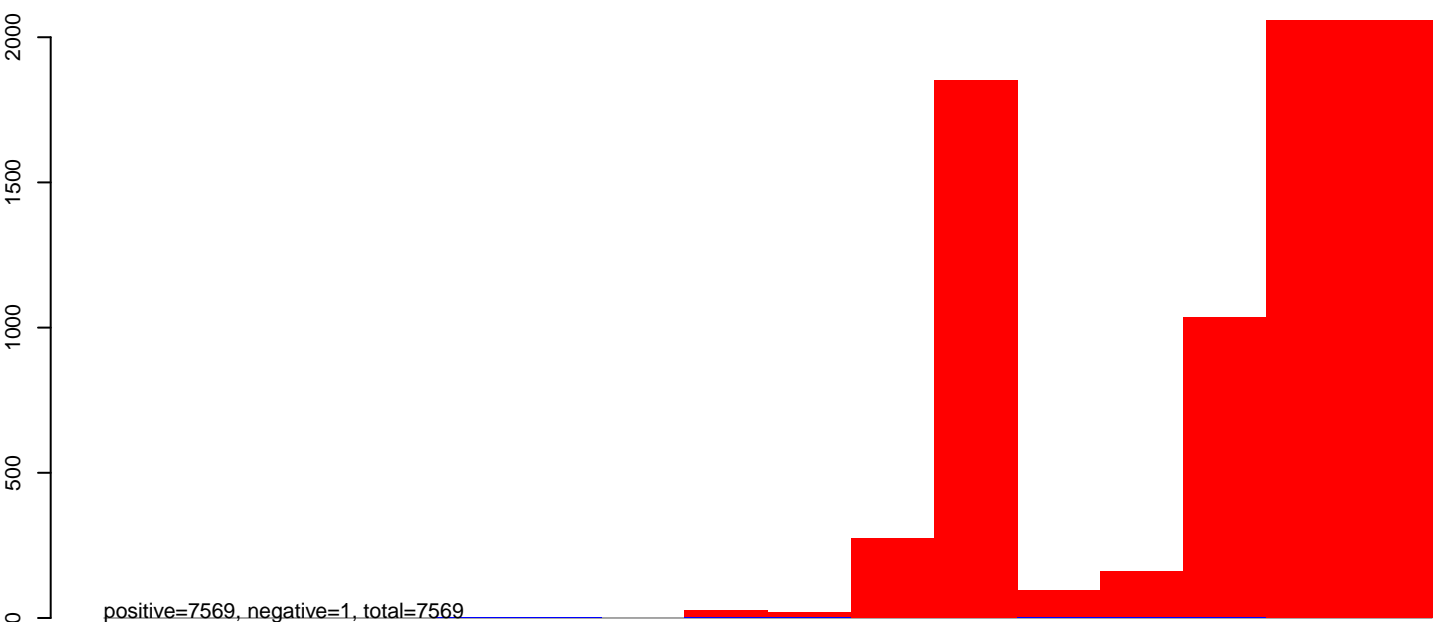
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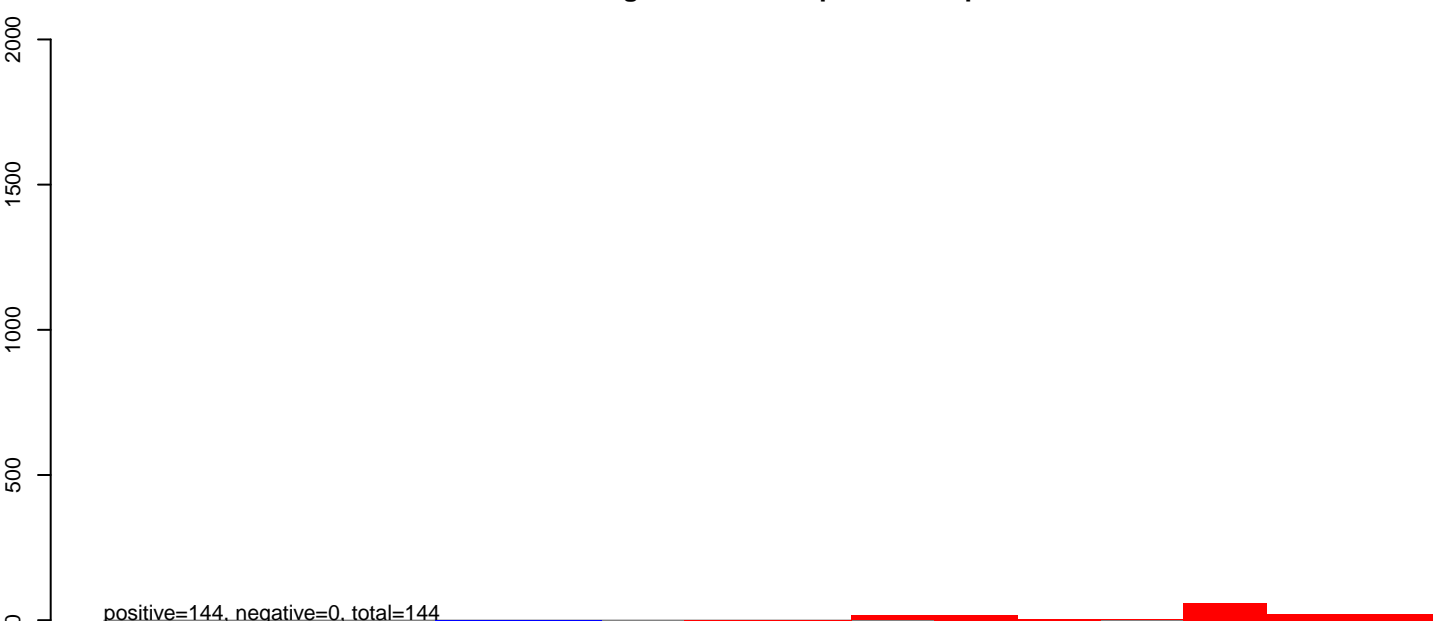
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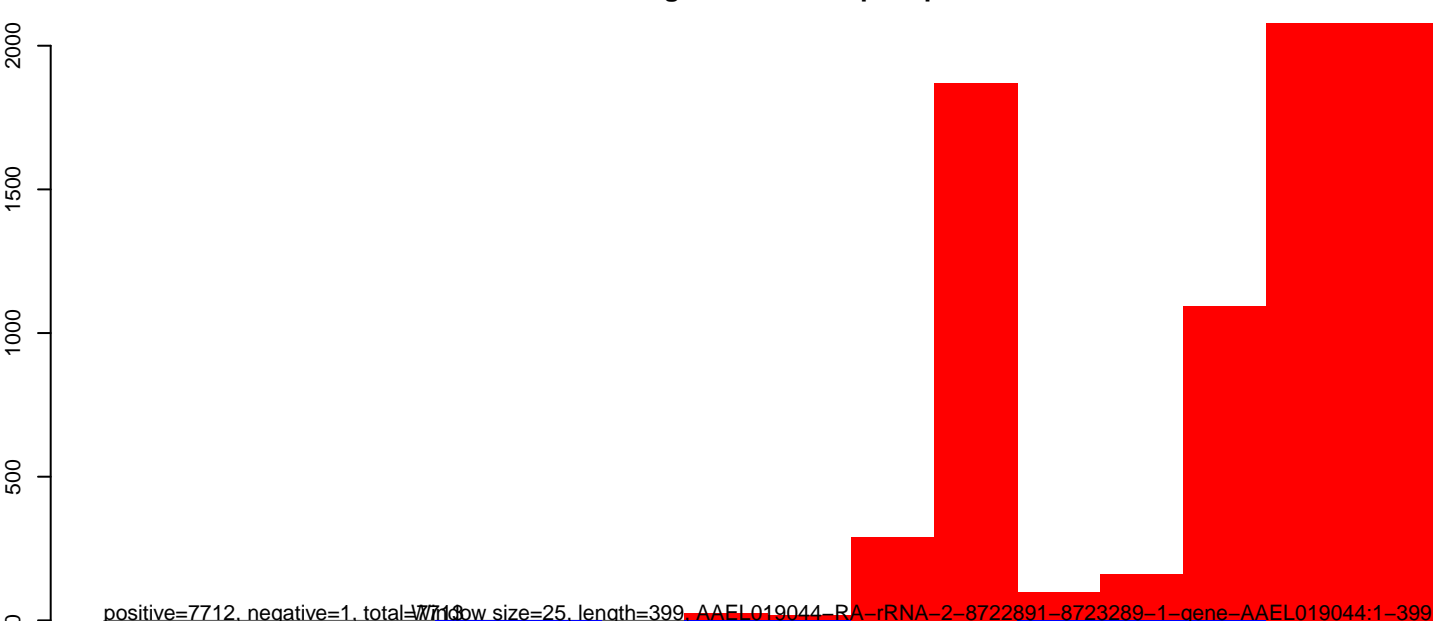
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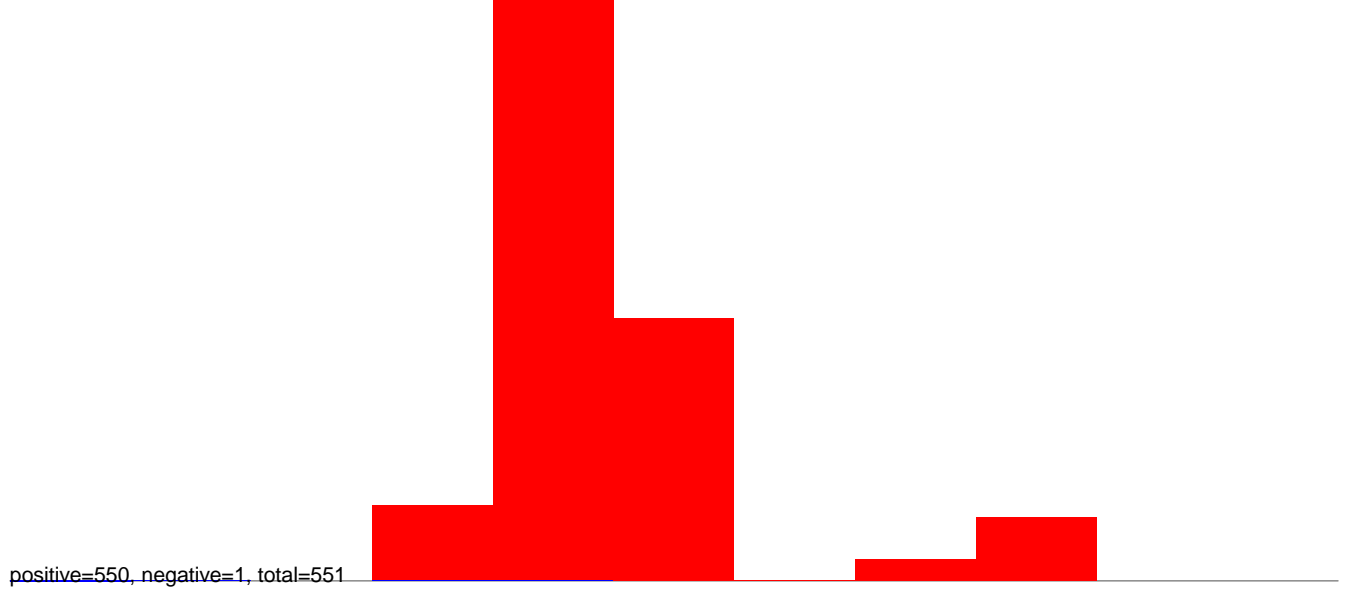
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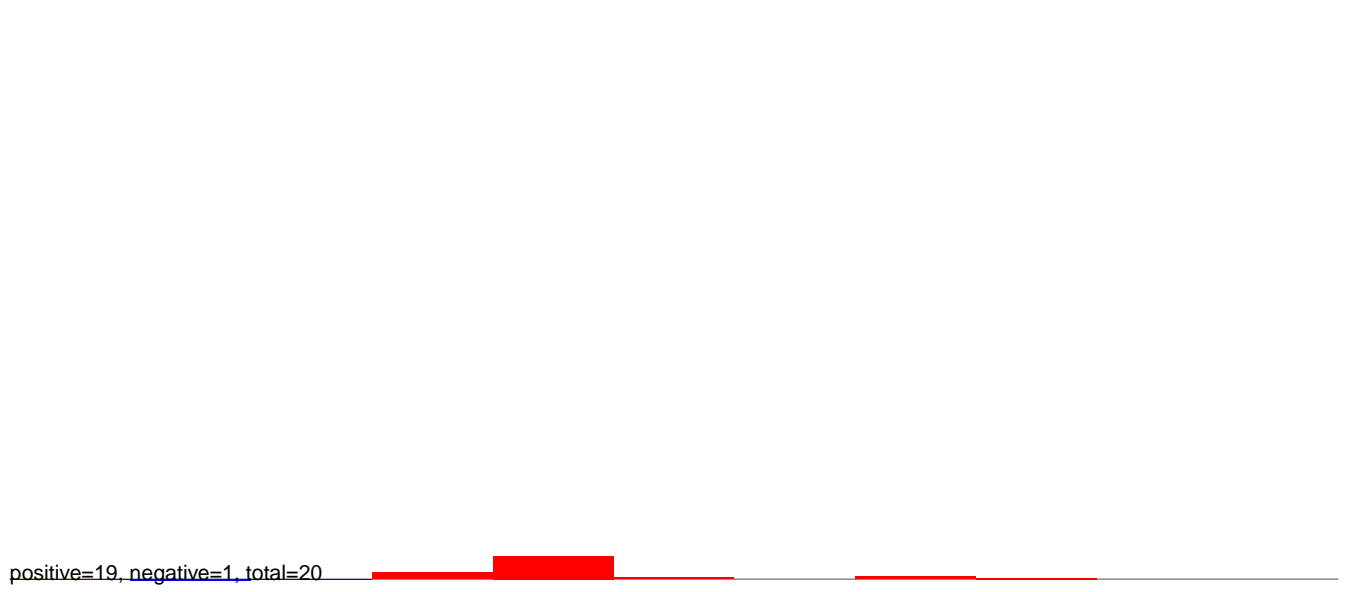
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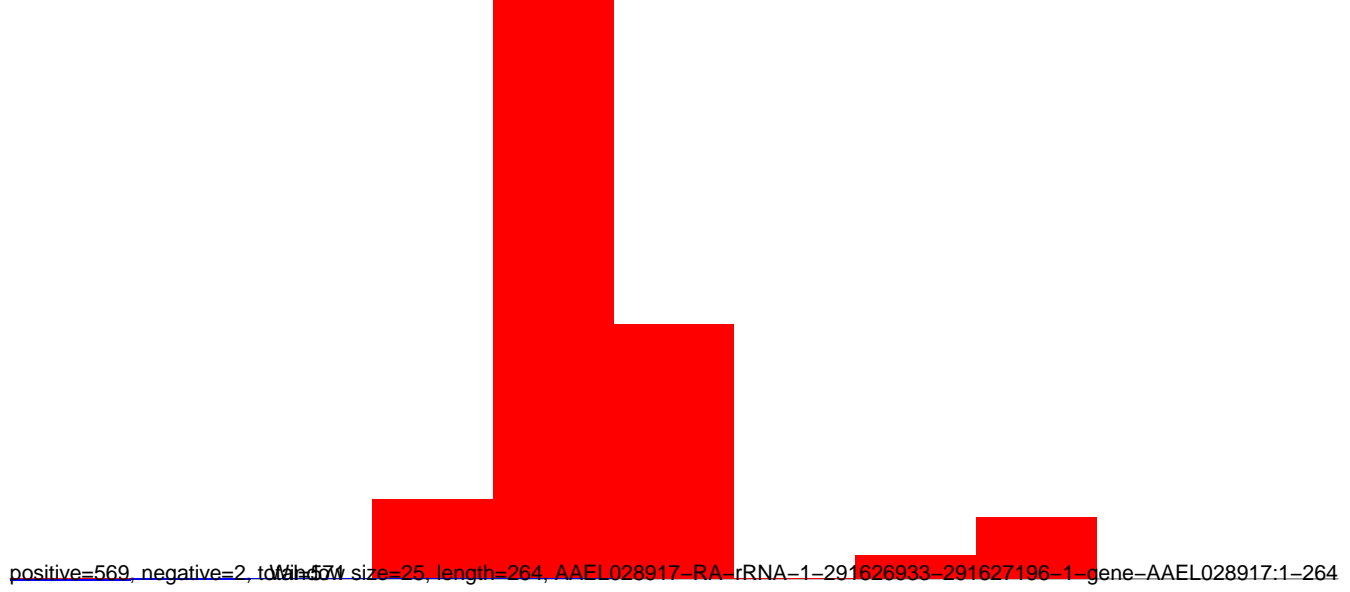
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AeAeg_Fcarc_GH_rep3.24_35.rep

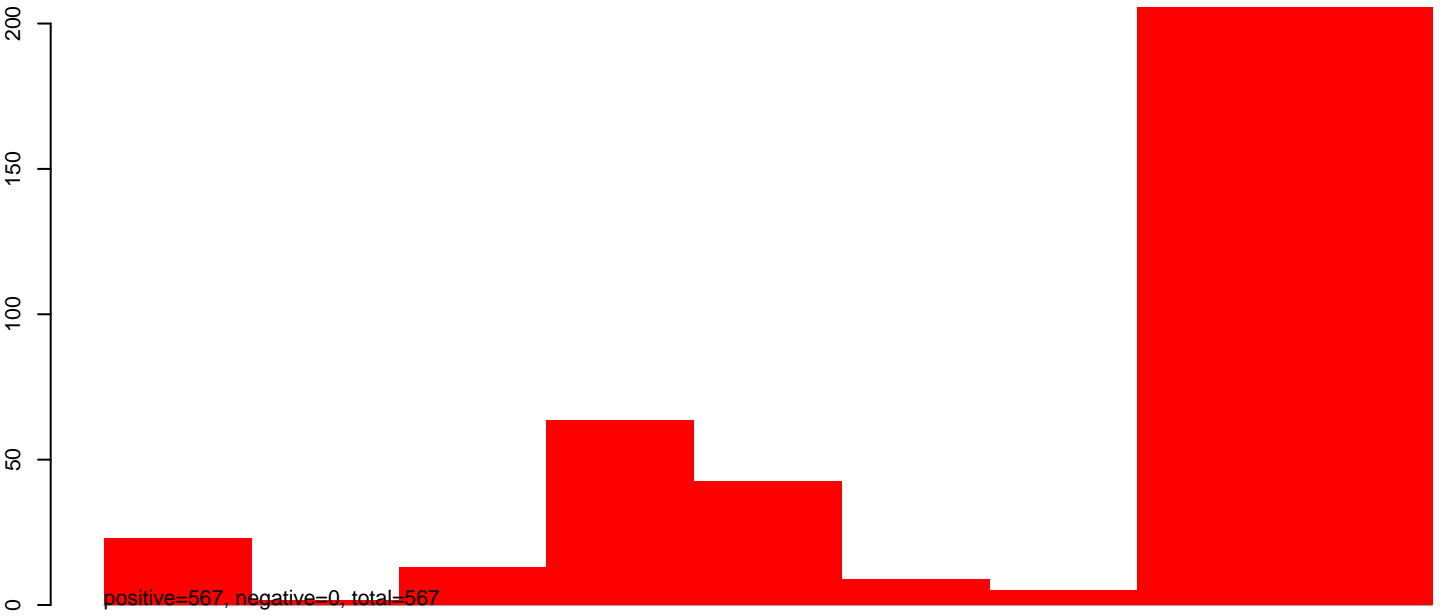


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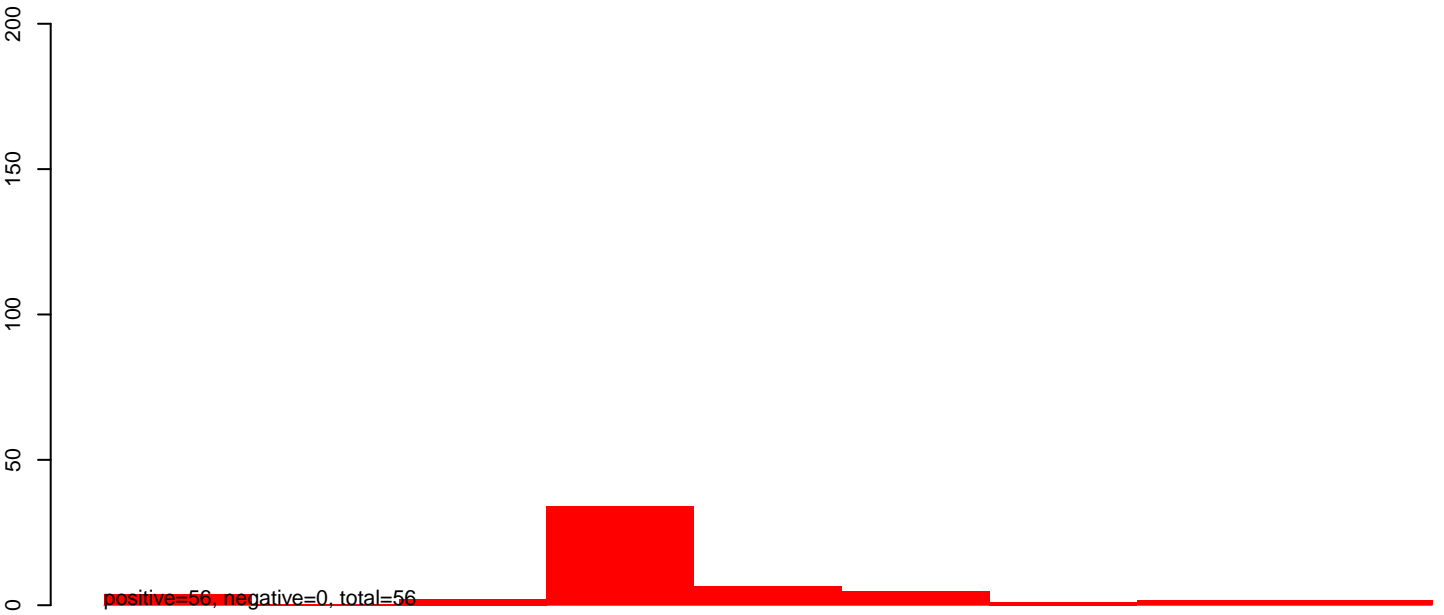


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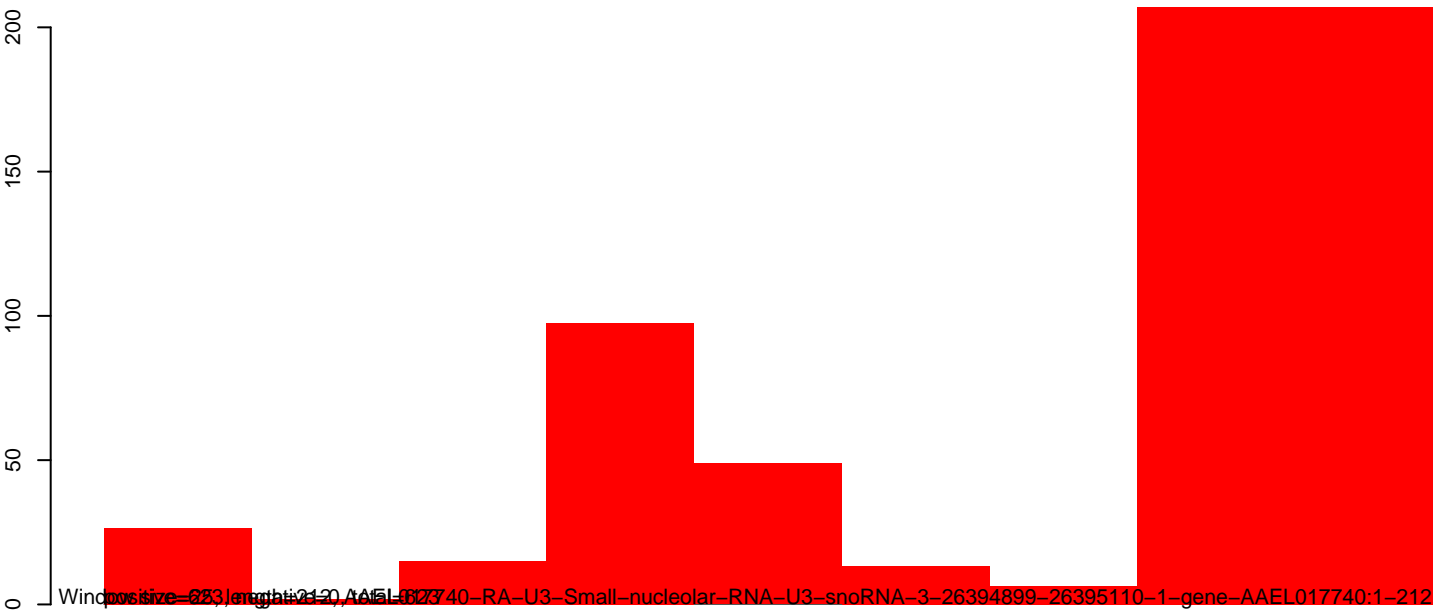
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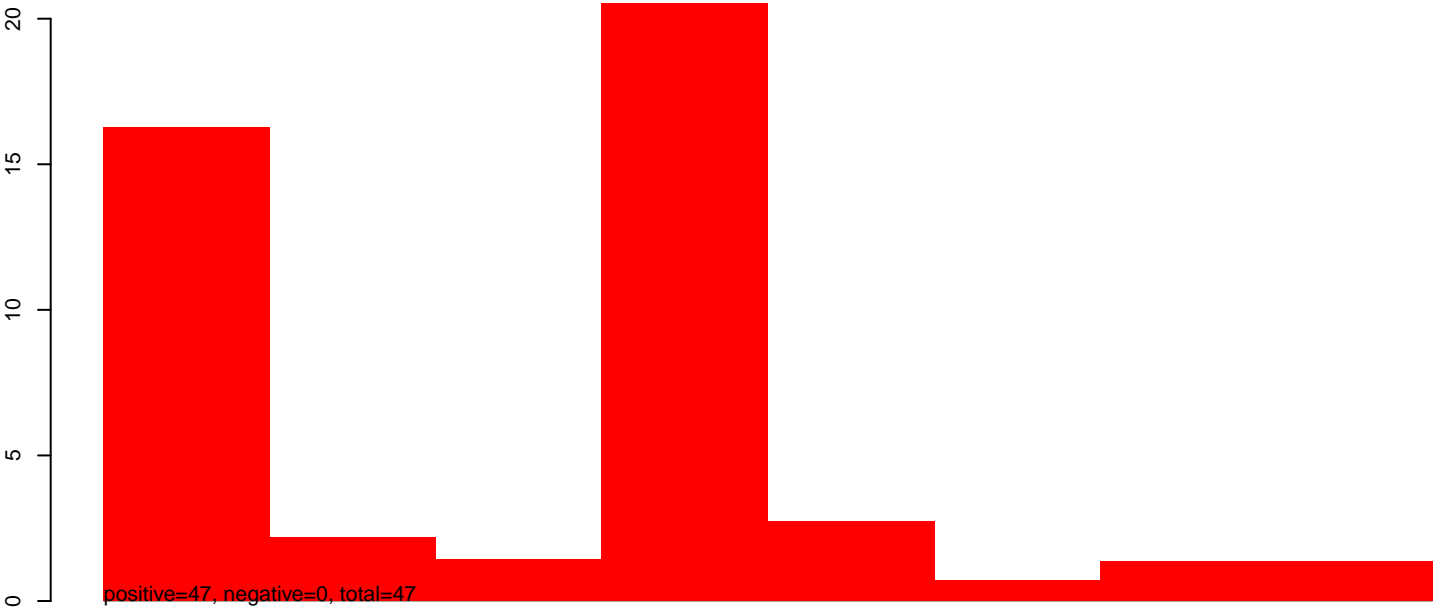
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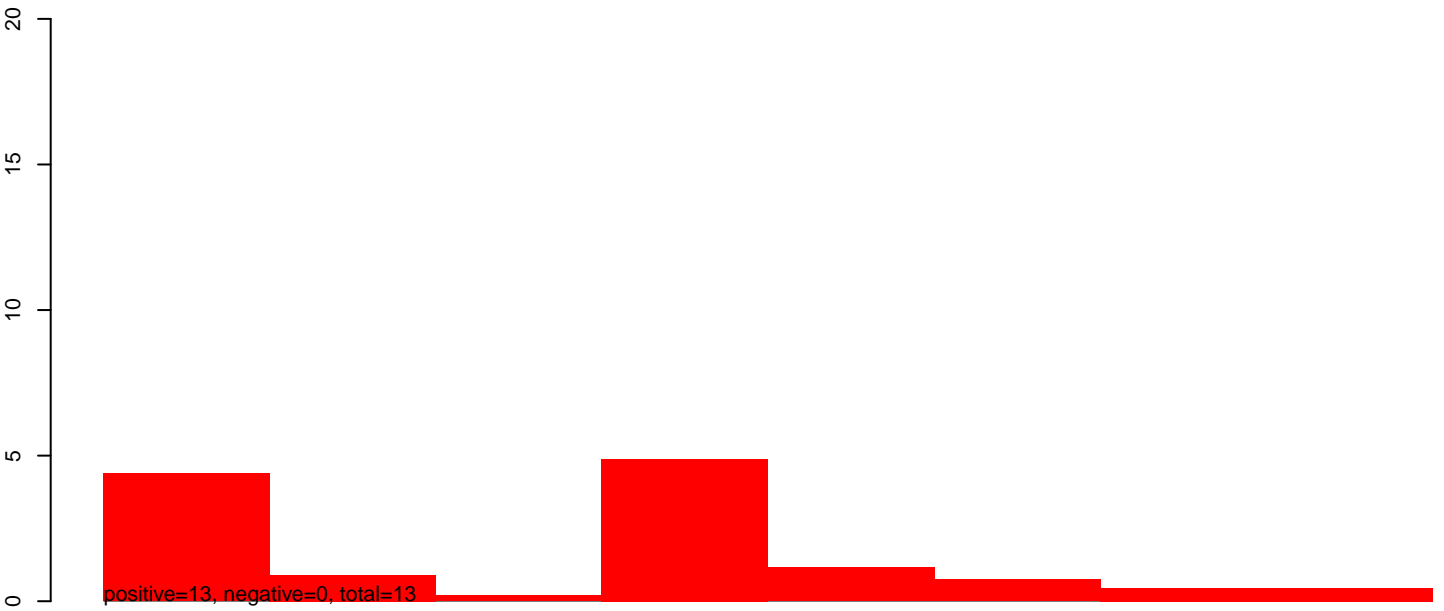
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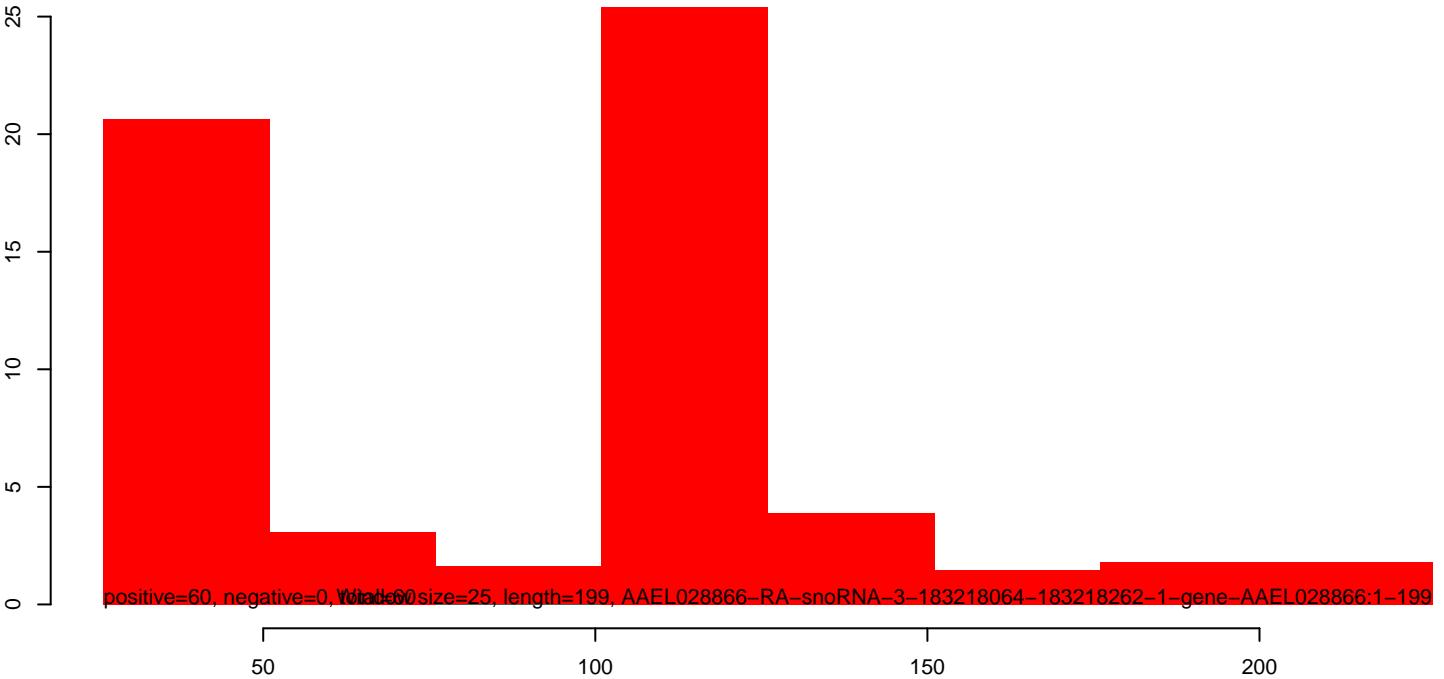
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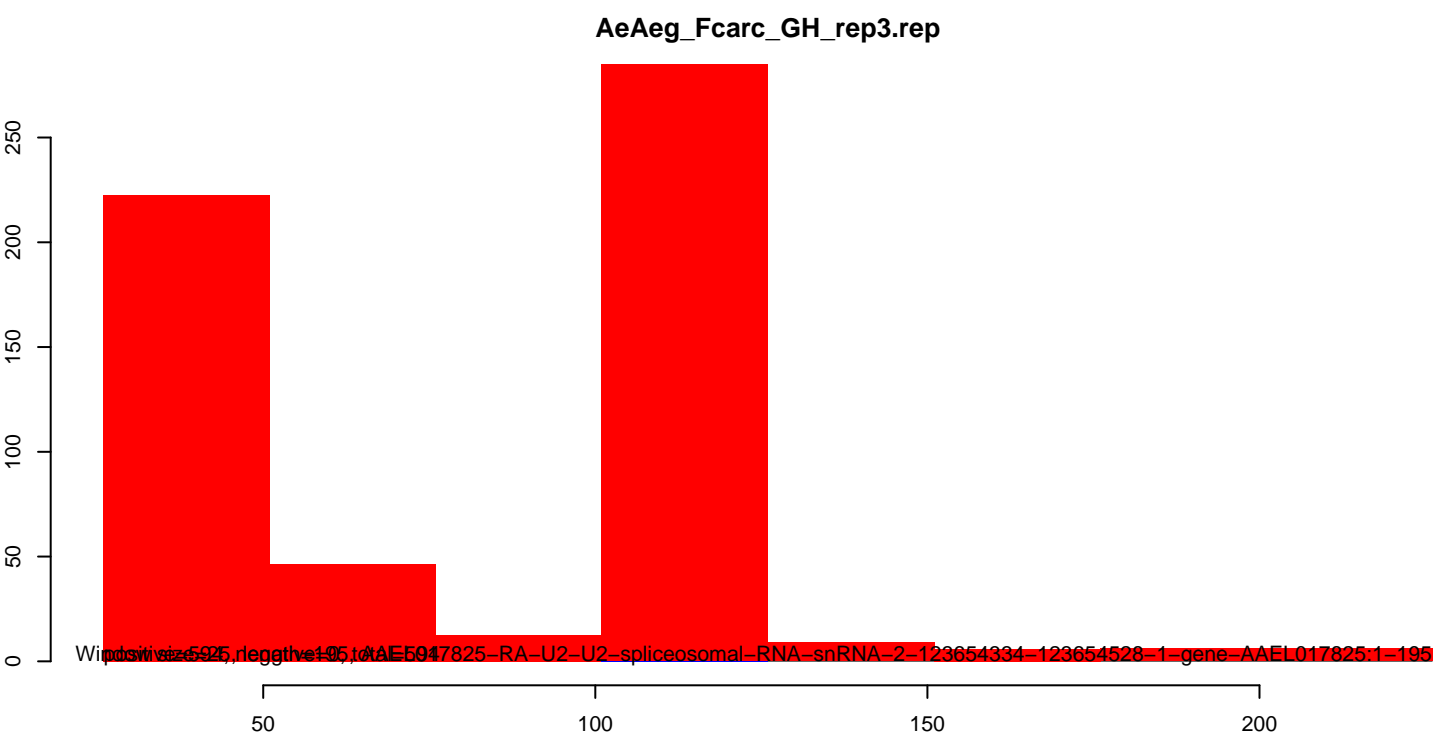
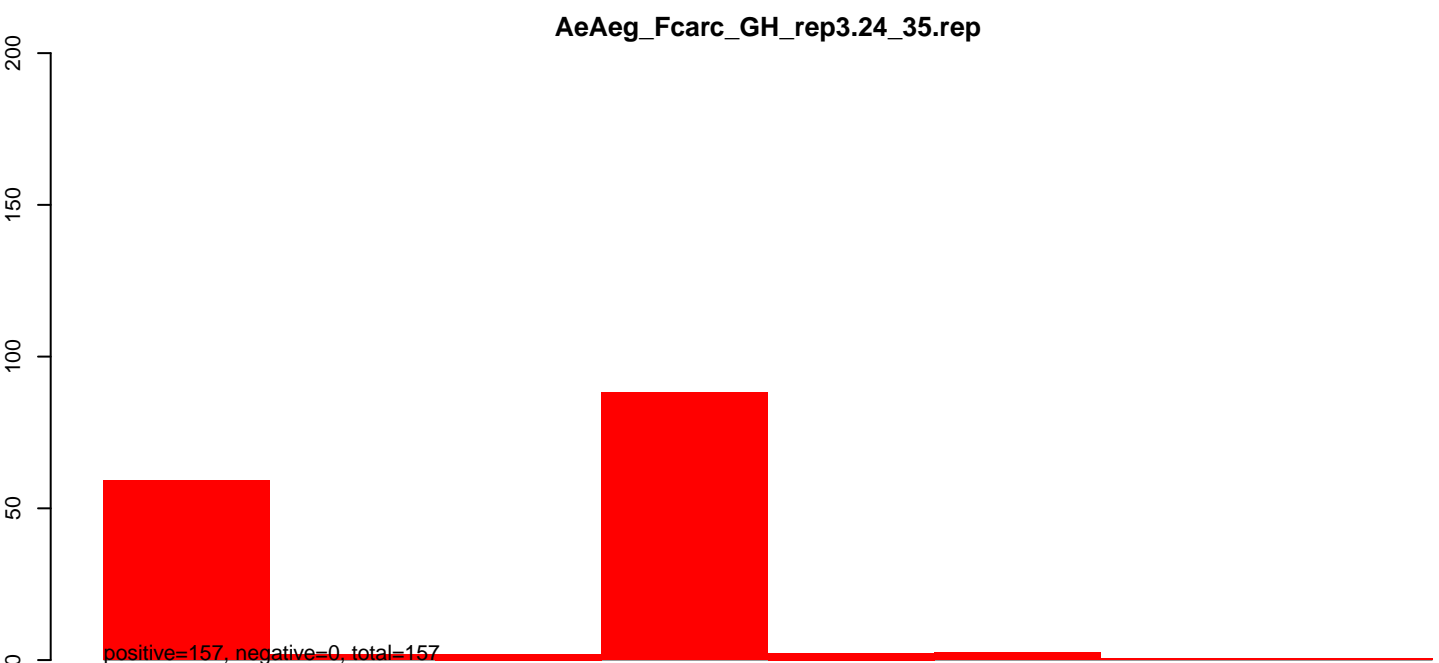
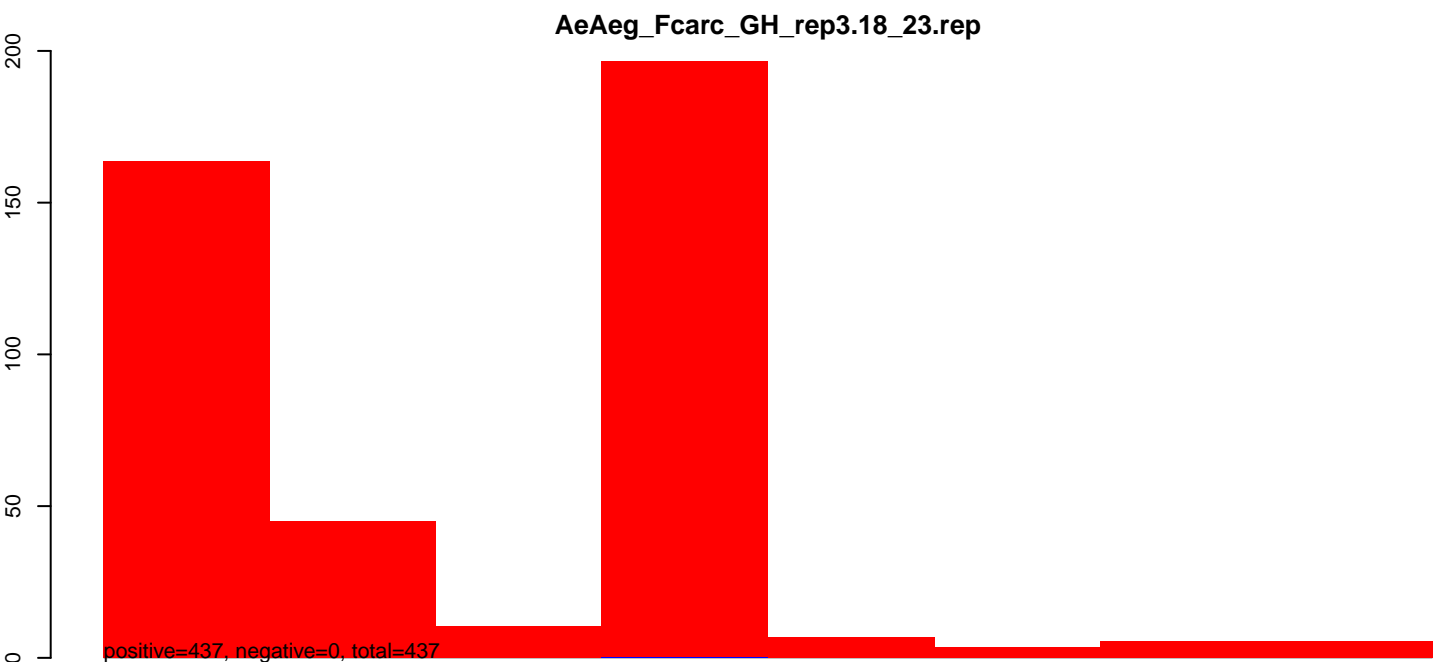


AeAeg_Fcarc_GH_rep3.24_35.rep



AeAeg_Fcarc_GH_rep3.rep





AeAeg_Fcarc_GH_rep3.18_23.rep

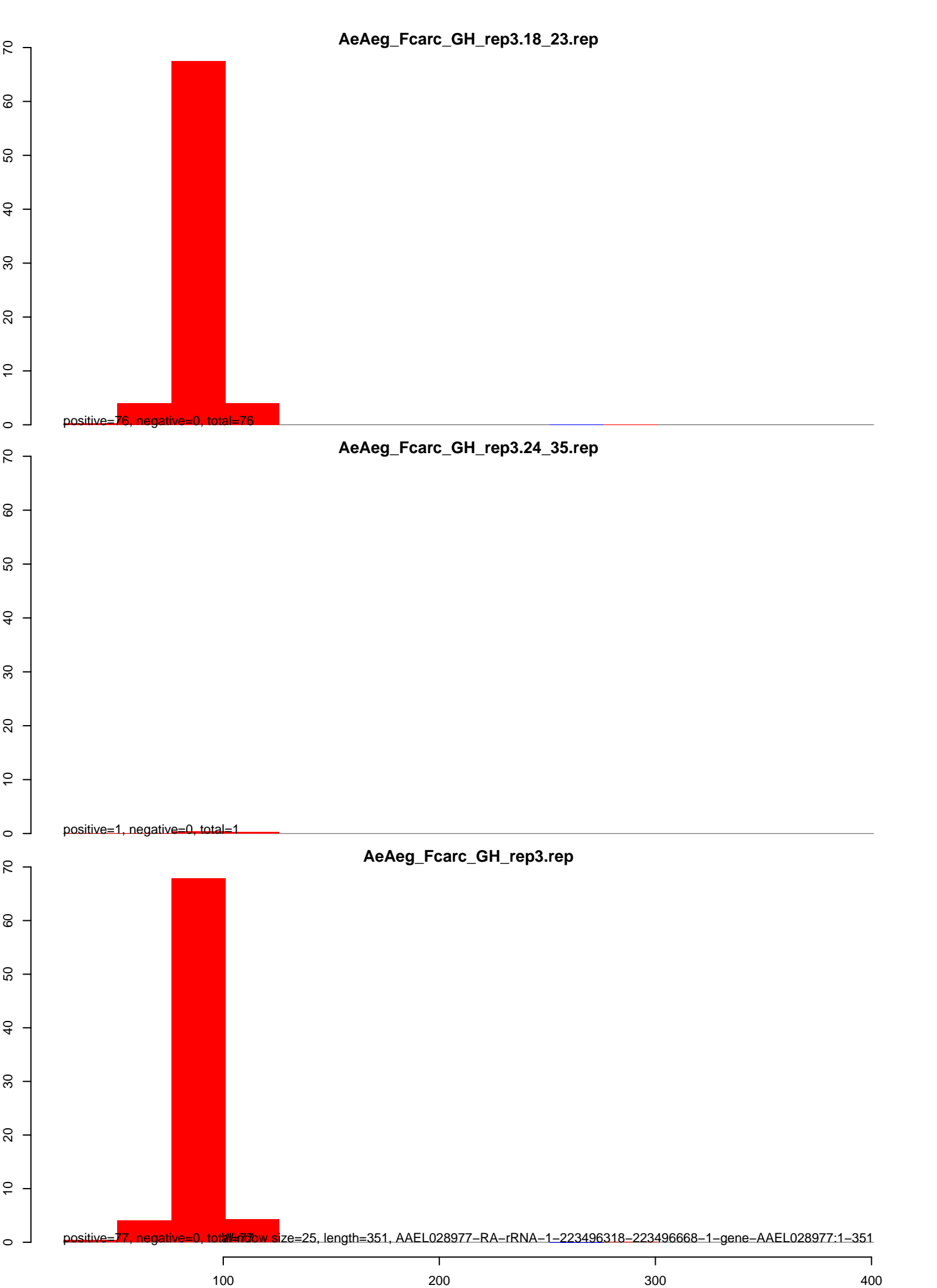
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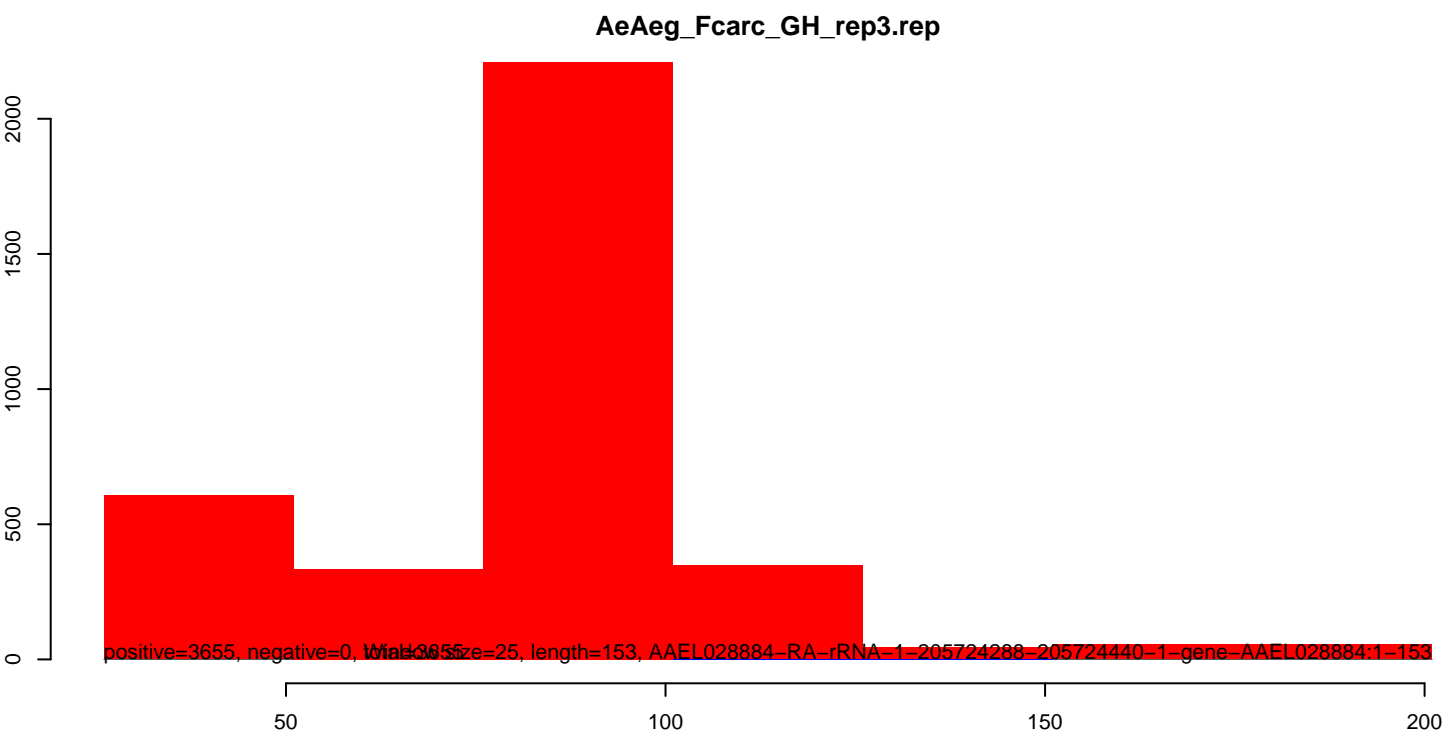
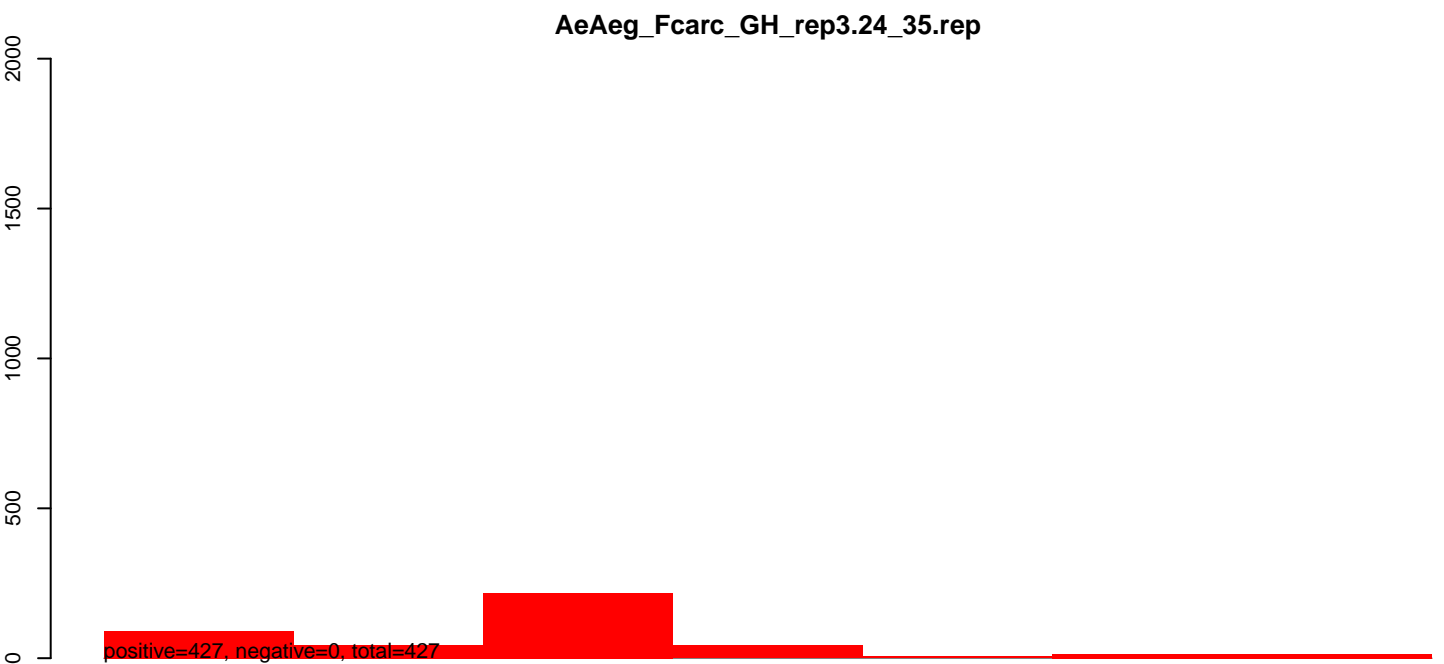
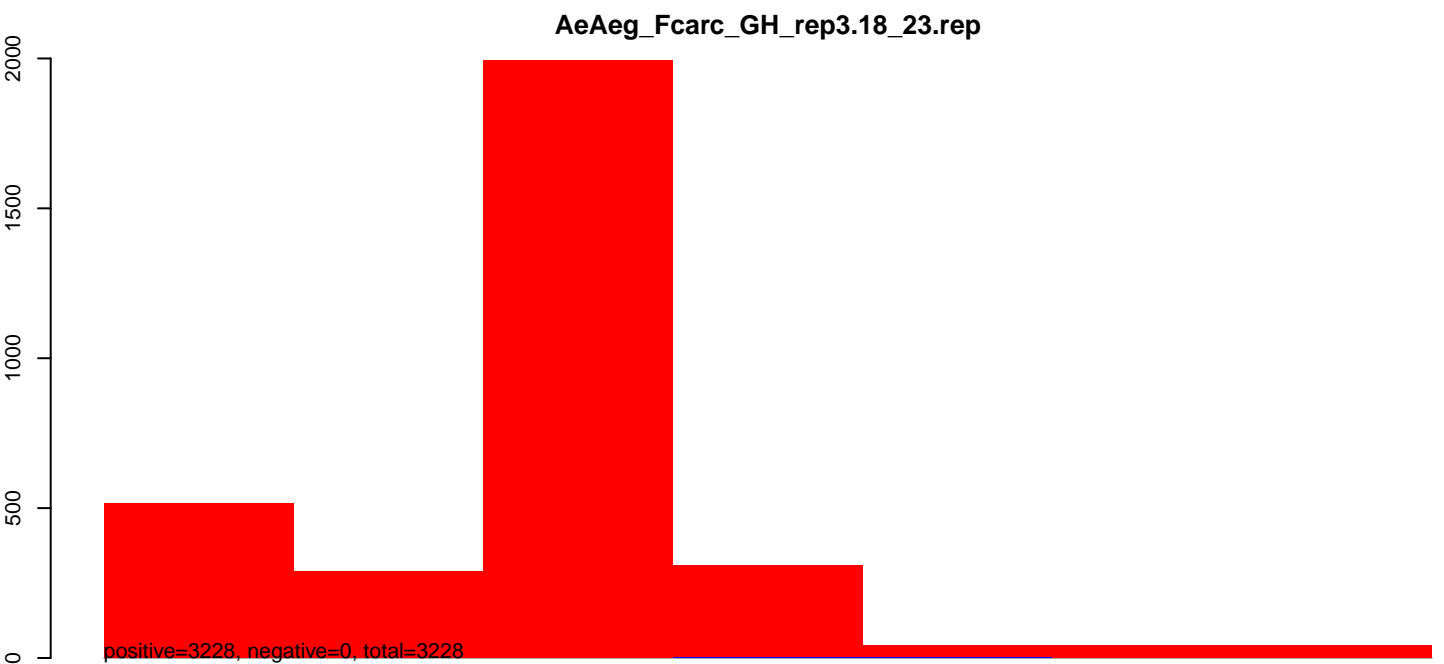
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positive=1, negative=0, total=1

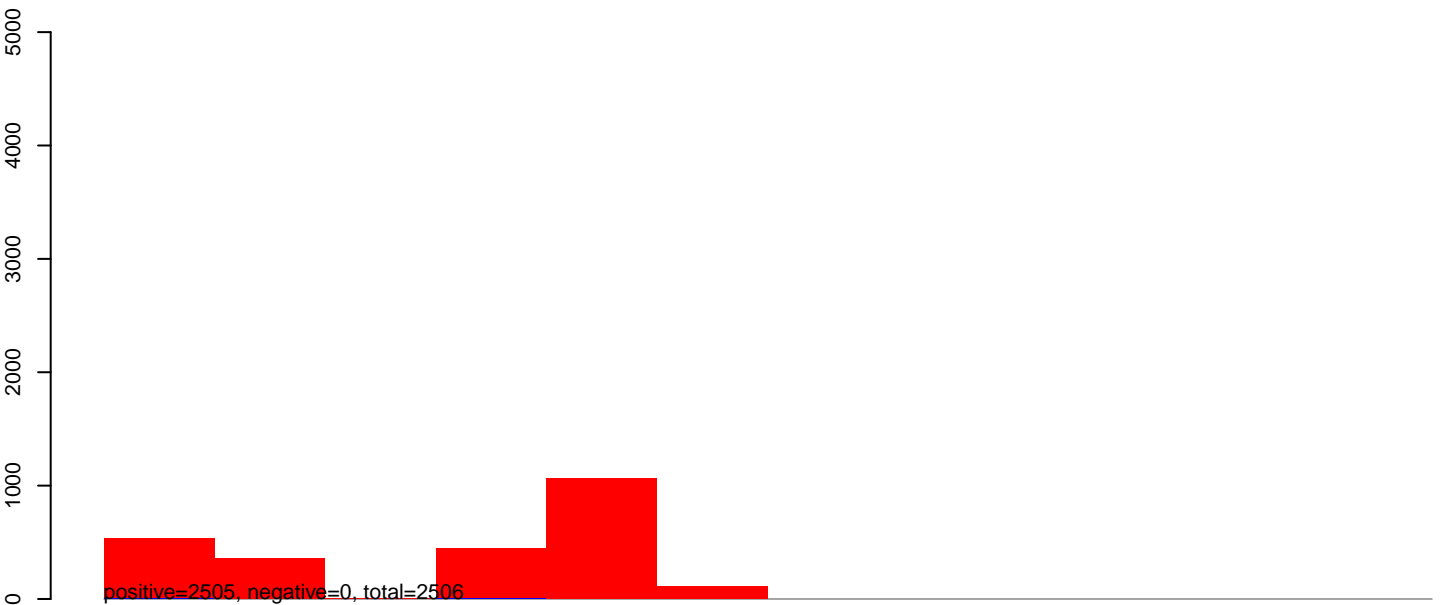
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positive=77, negative=0, total=77, window size=25, length=351, AAEL028977-RA-rRNA-1-223496318-223496668-1-gene-AAEL028977:1-351

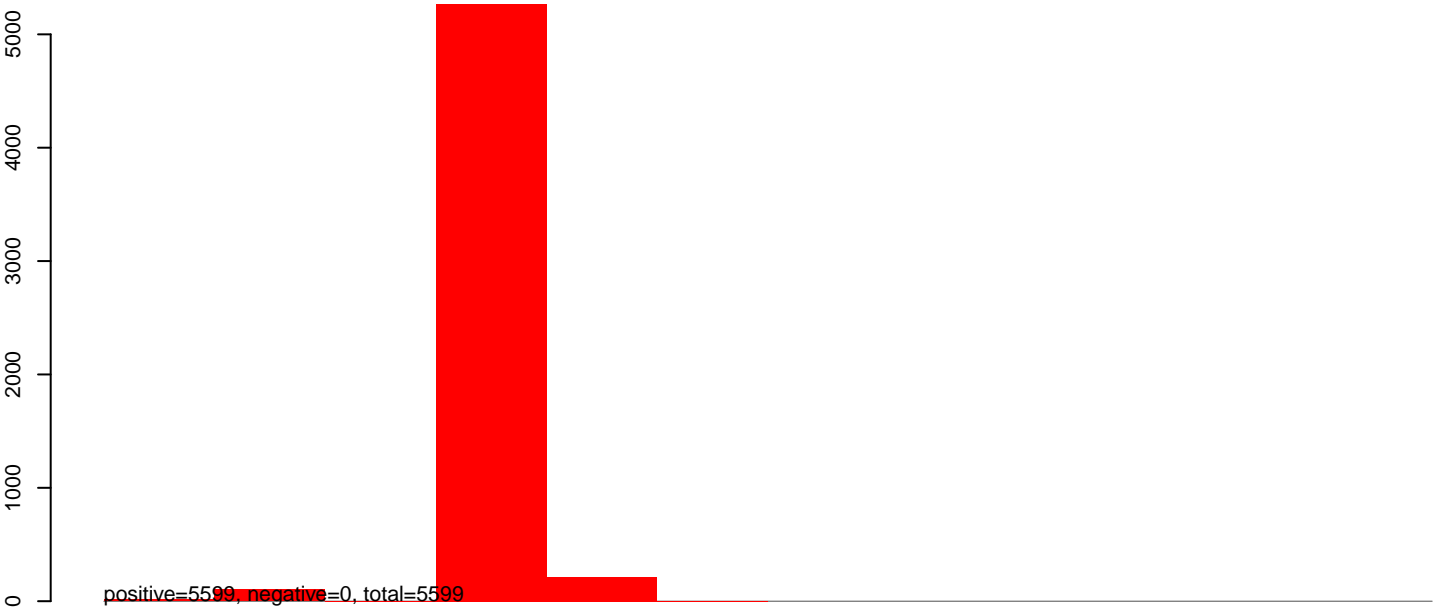




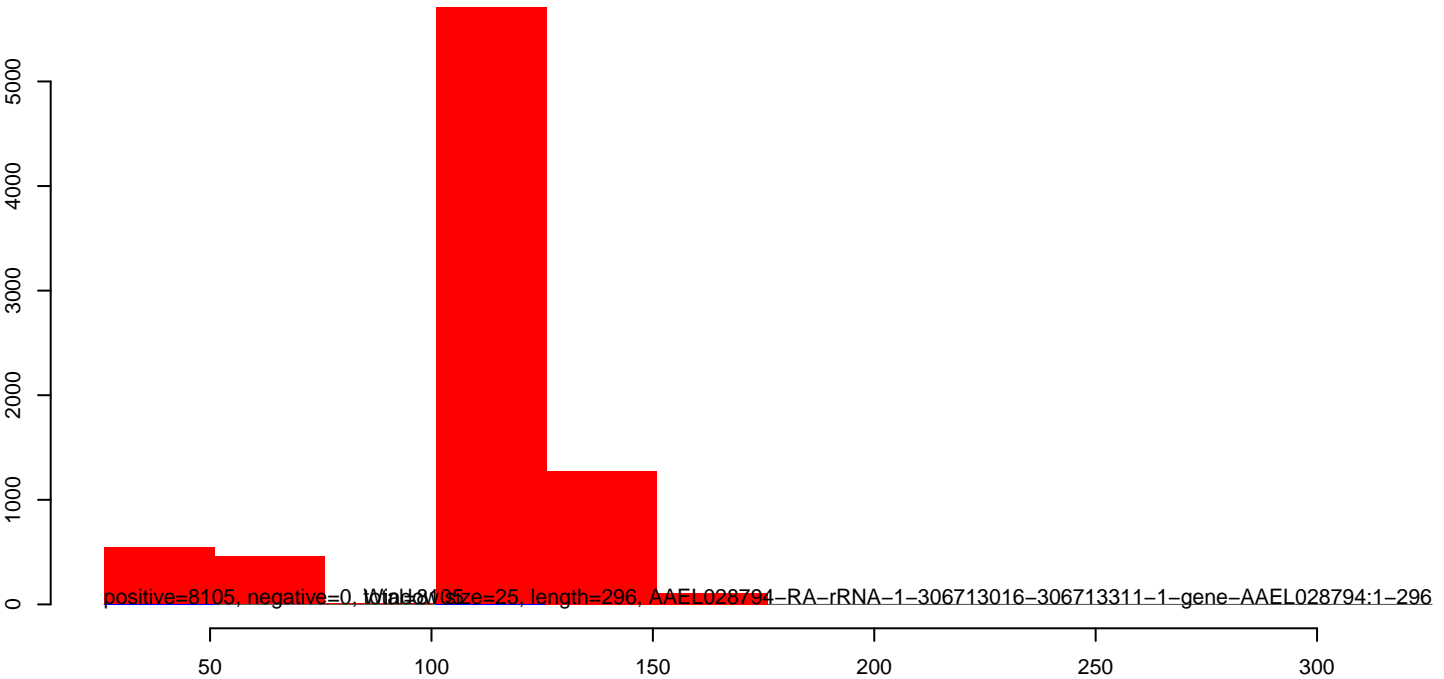
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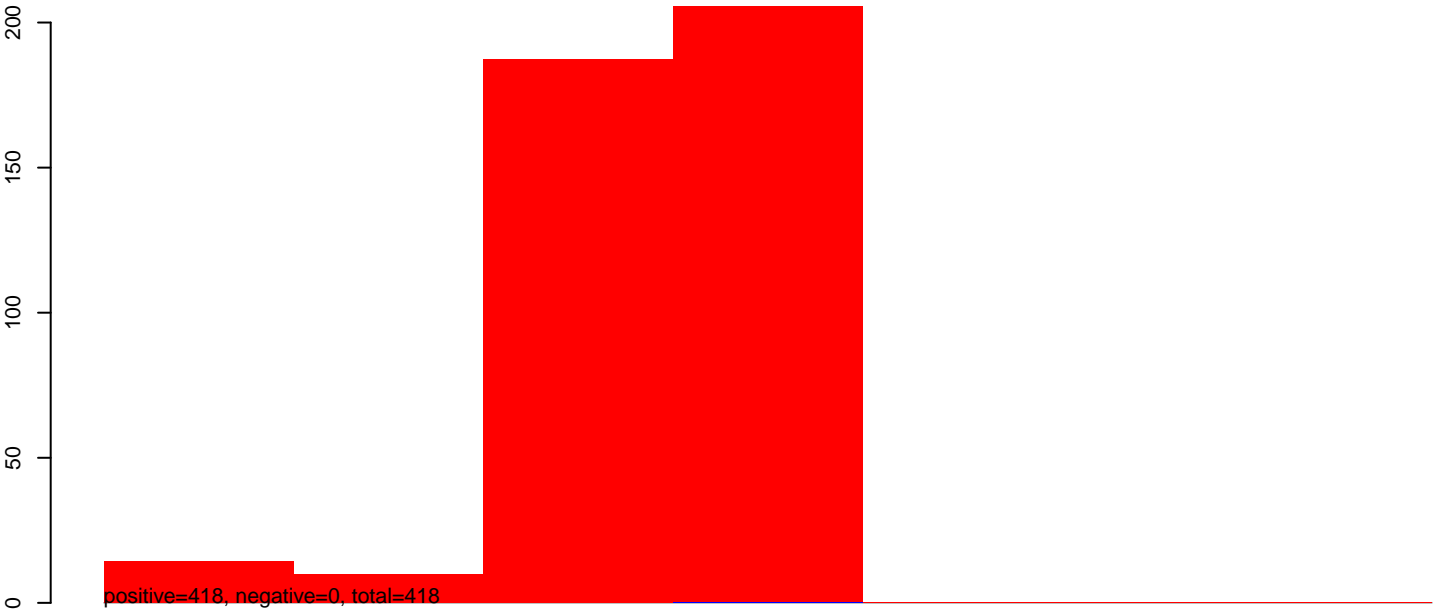
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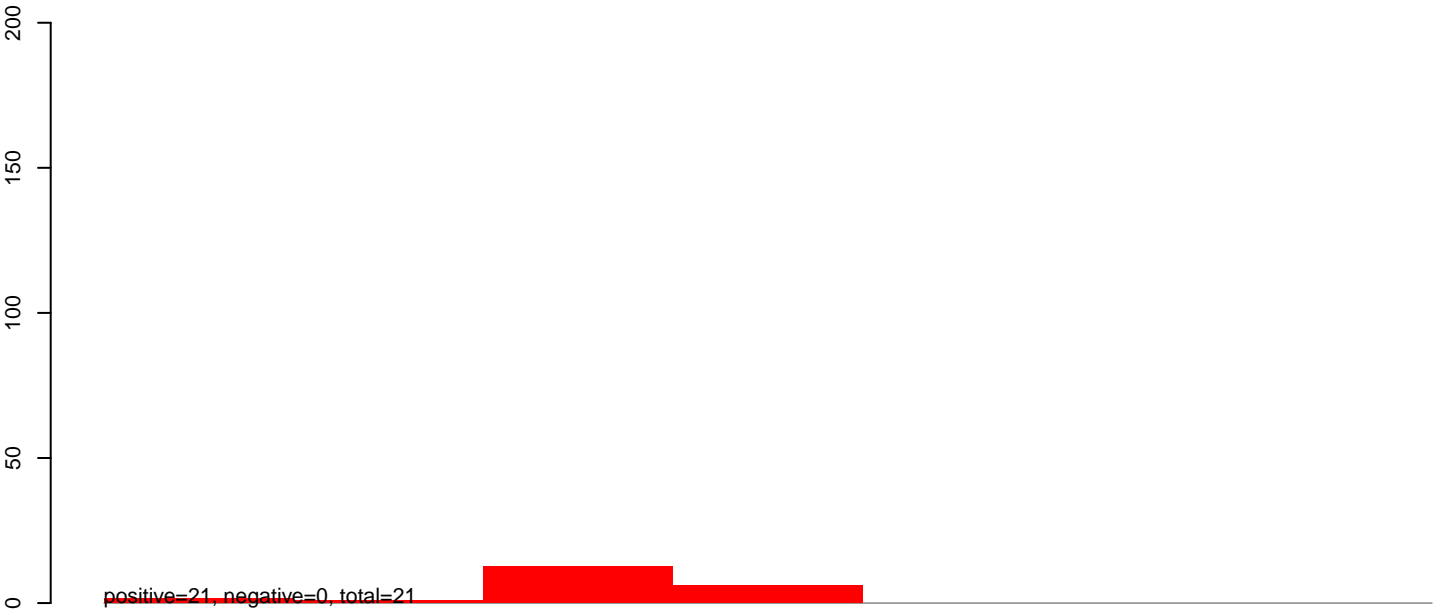
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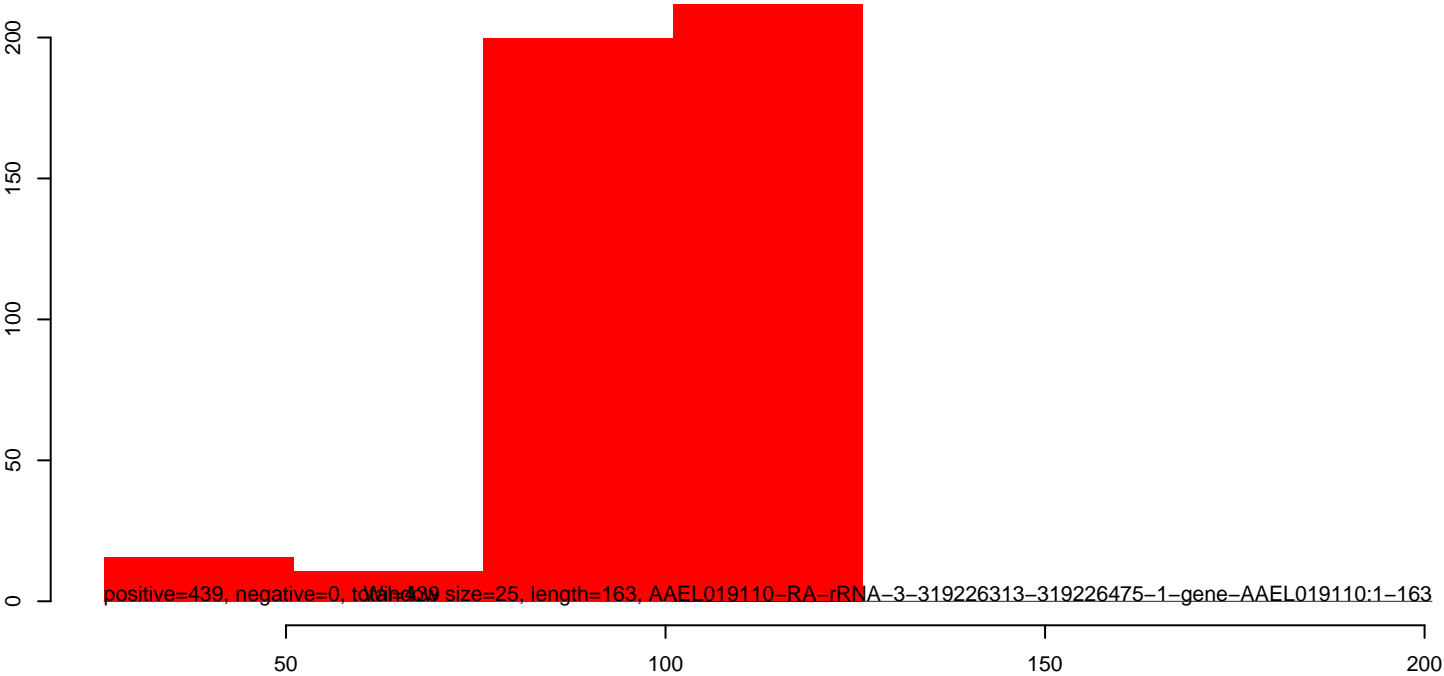
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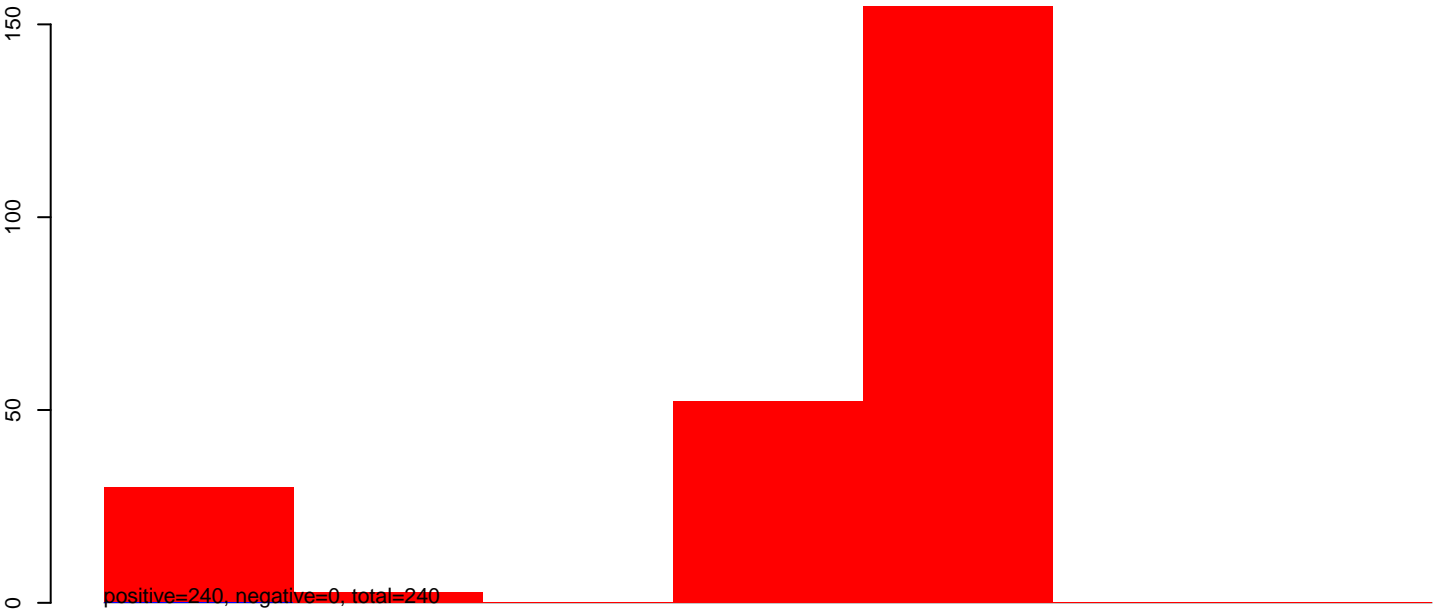
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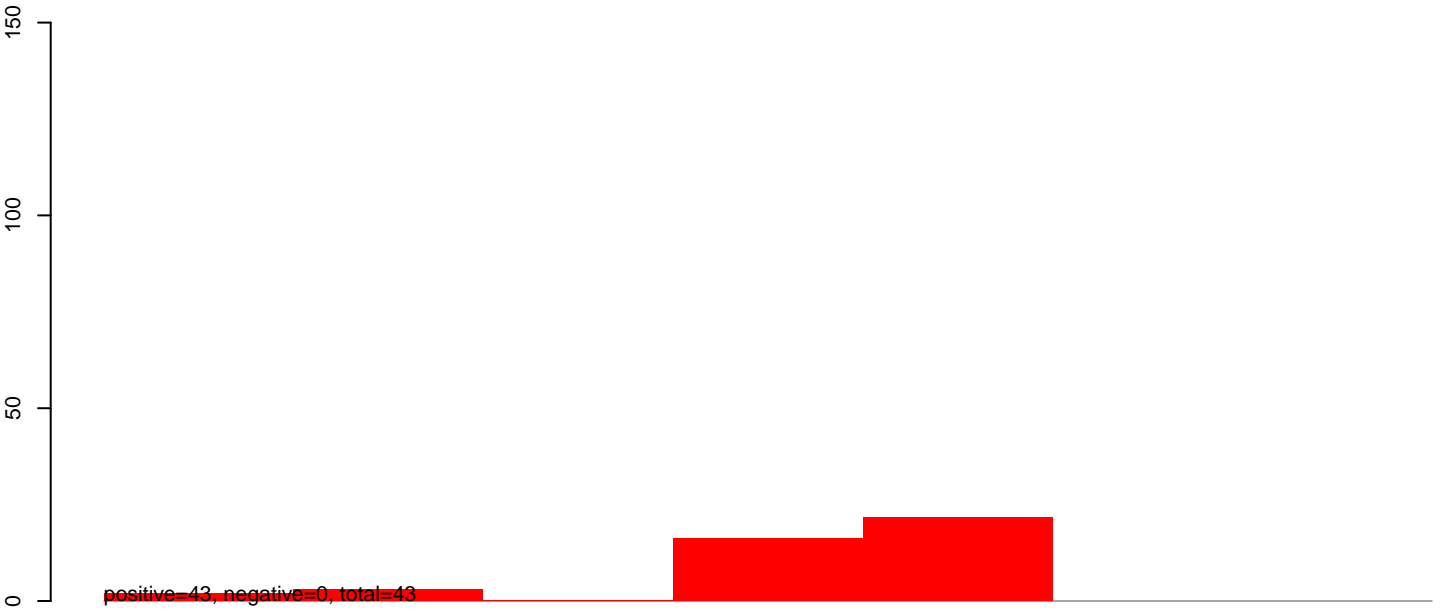
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AeAeg_Fcarc_GH_rep3.18_23.rep



AeAeg_Fcarc_GH_rep3.24_35.rep

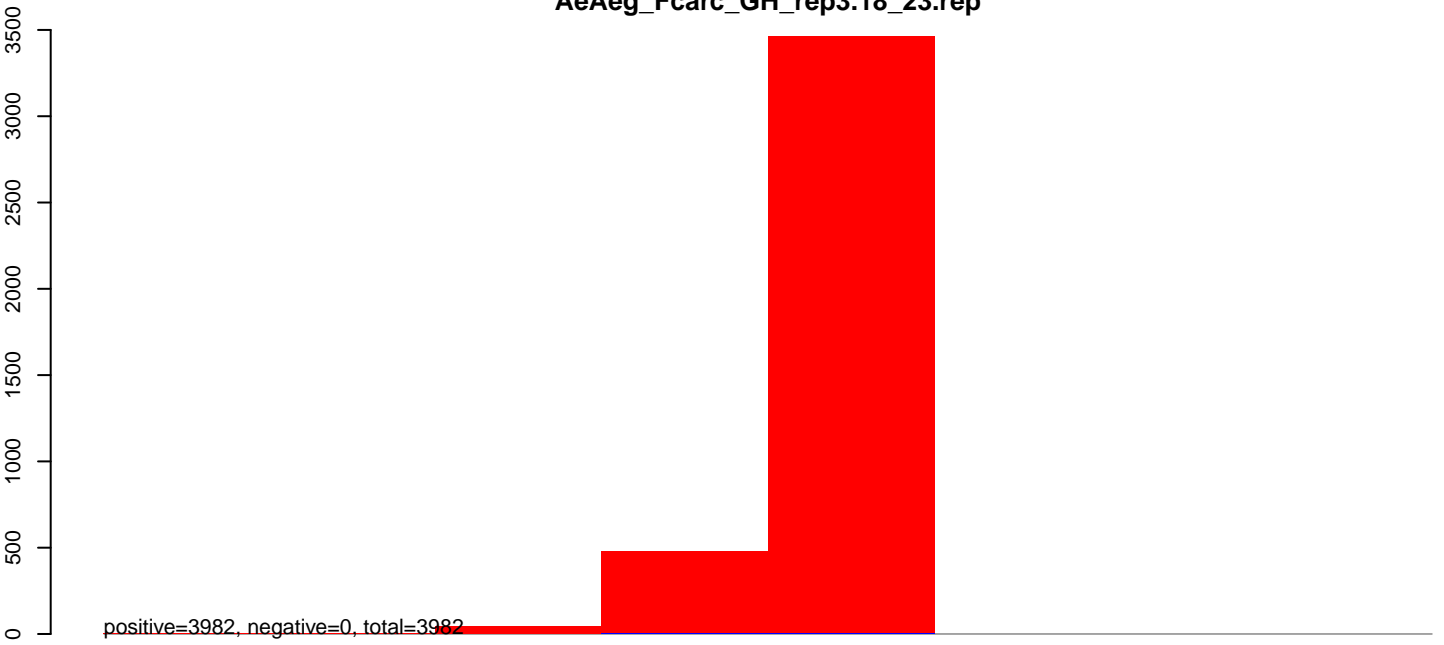


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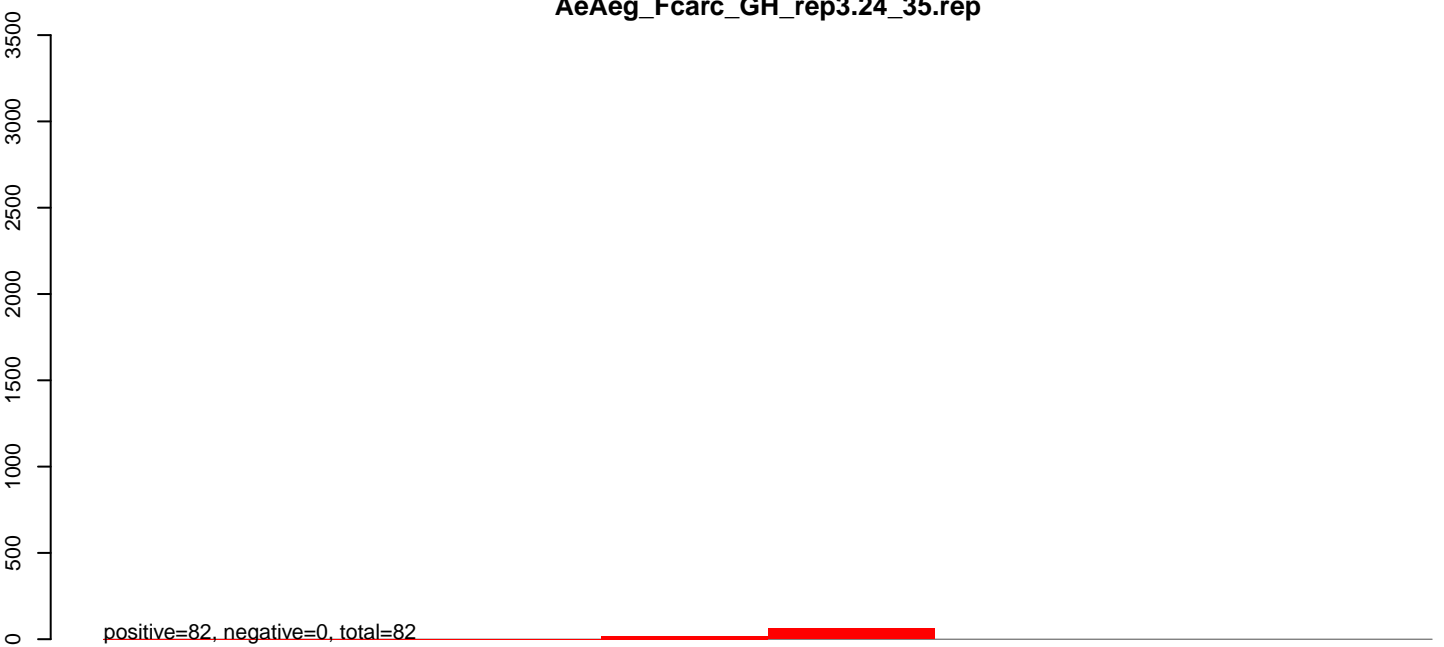
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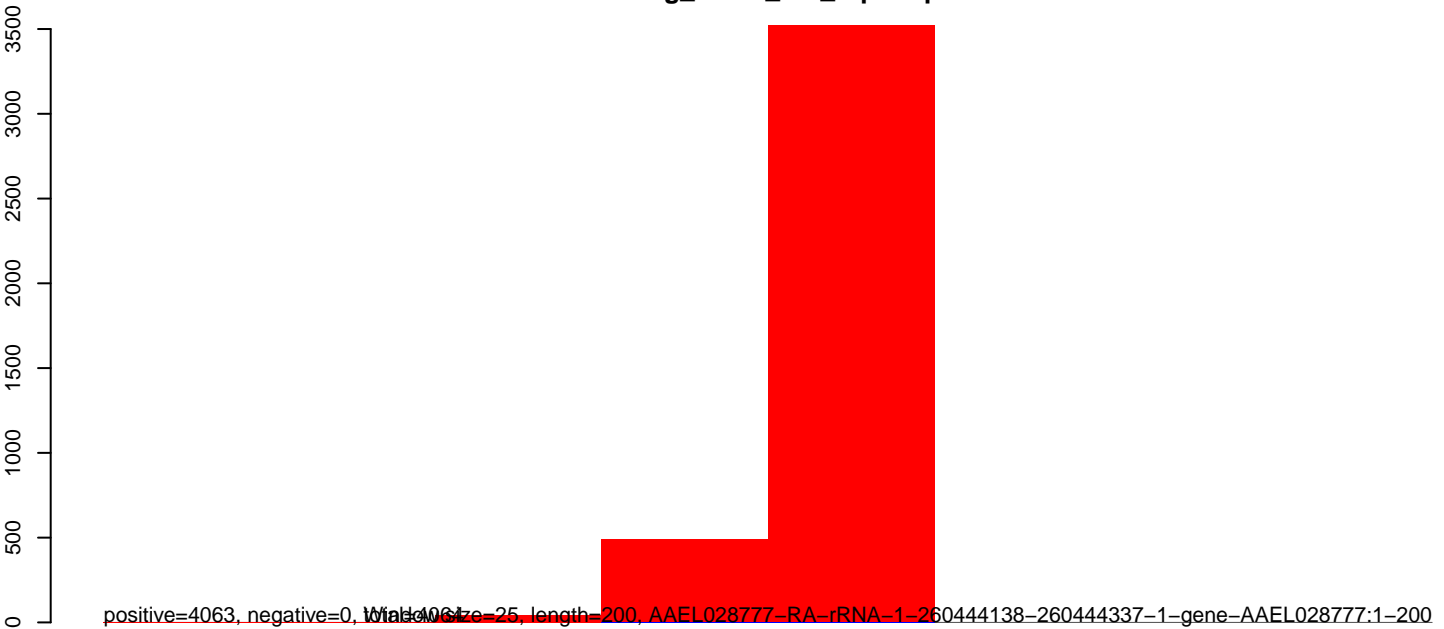
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positive=82, negative=0, total=82



AeAeg_FcArc_GH_rep3.rep

positive=4063, negative=0, total=4063, Window Size=25, length=200, AAEL028777-RA-rRNA-1-260444138-260444337-1-gene-AAEL028777:1-200



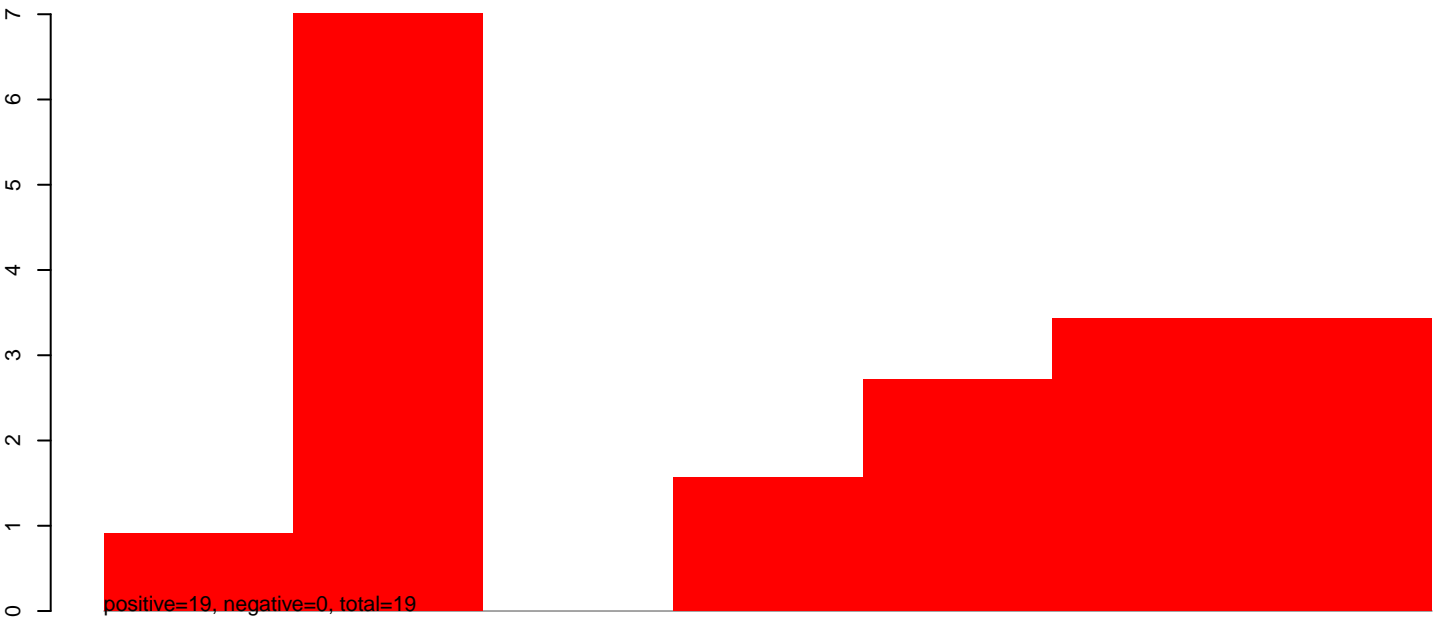
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100

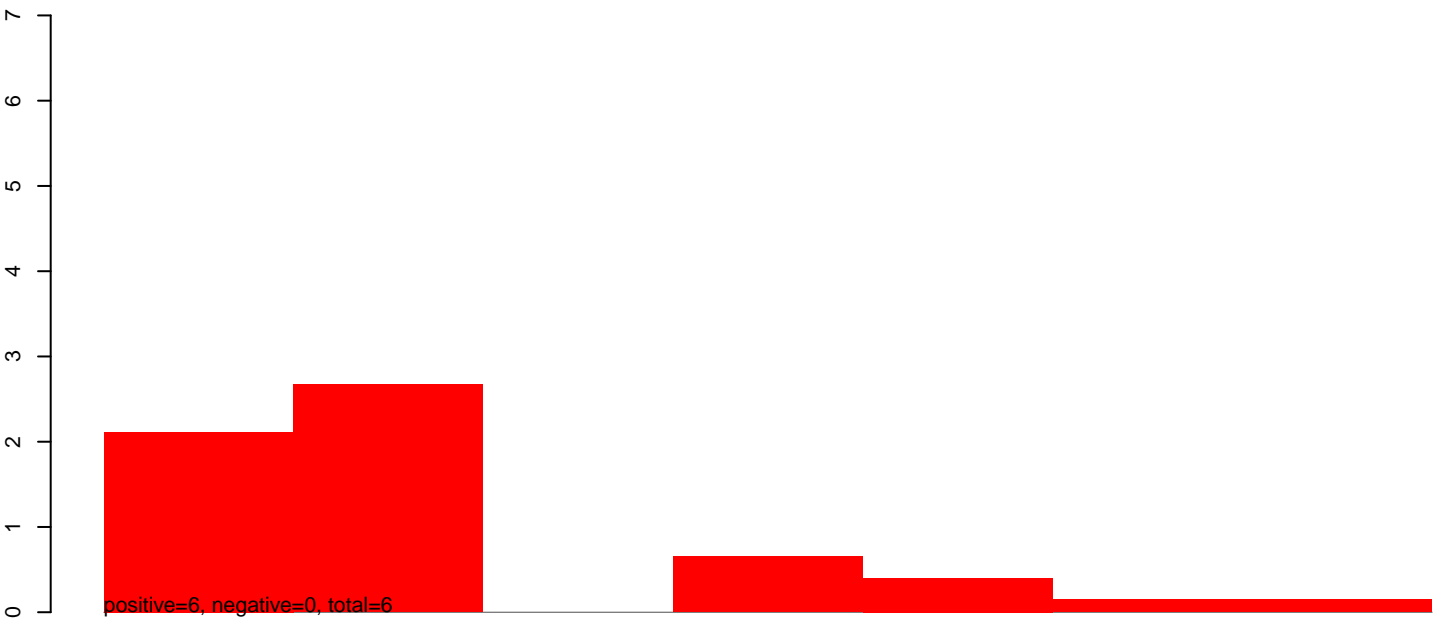
150

200

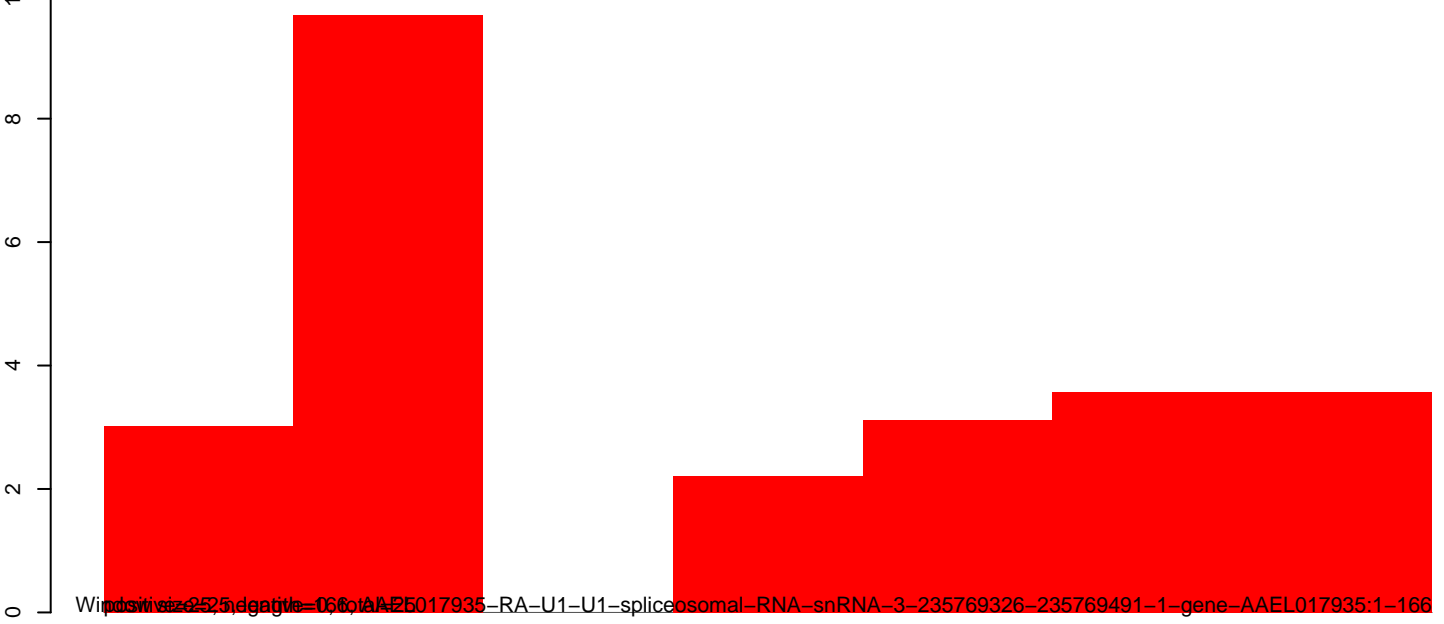
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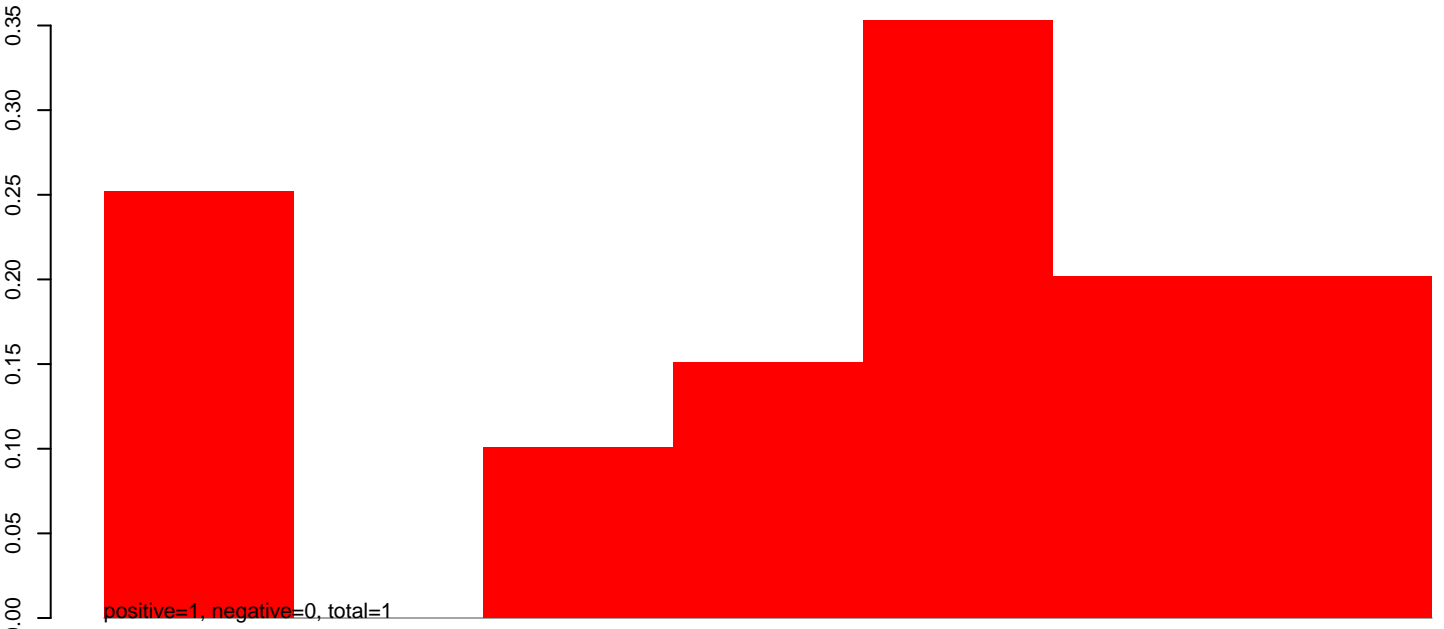
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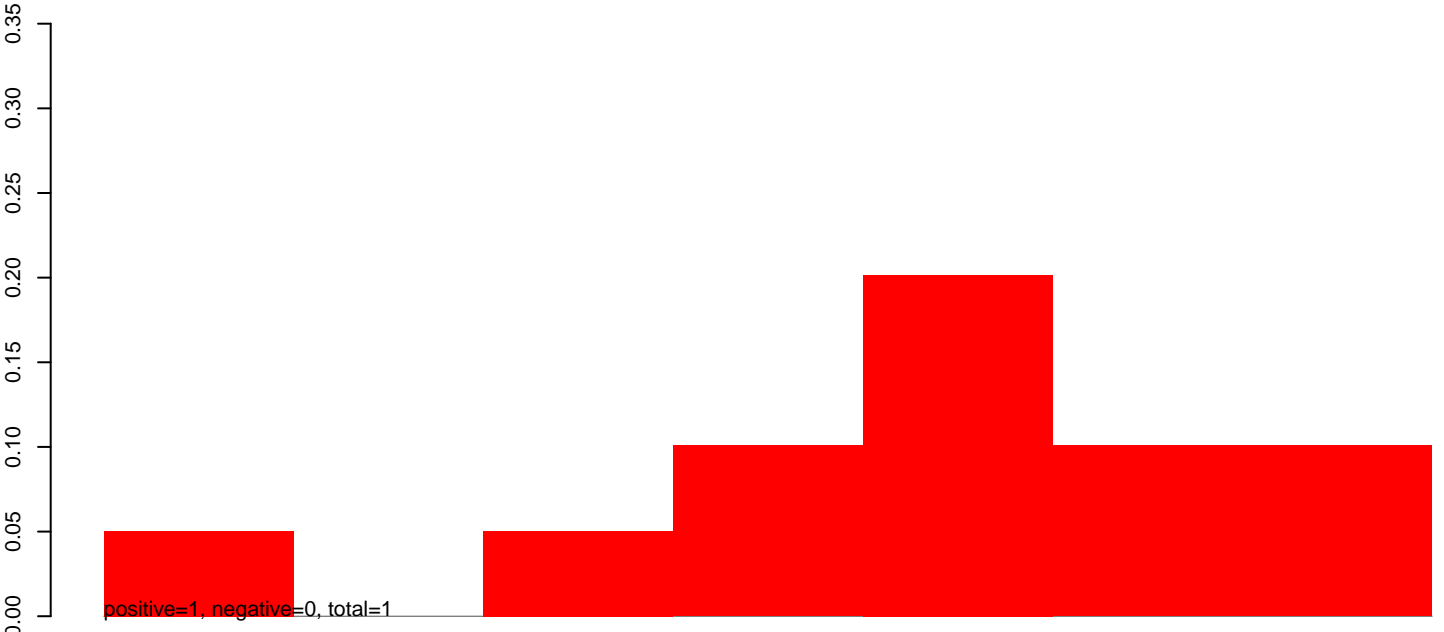
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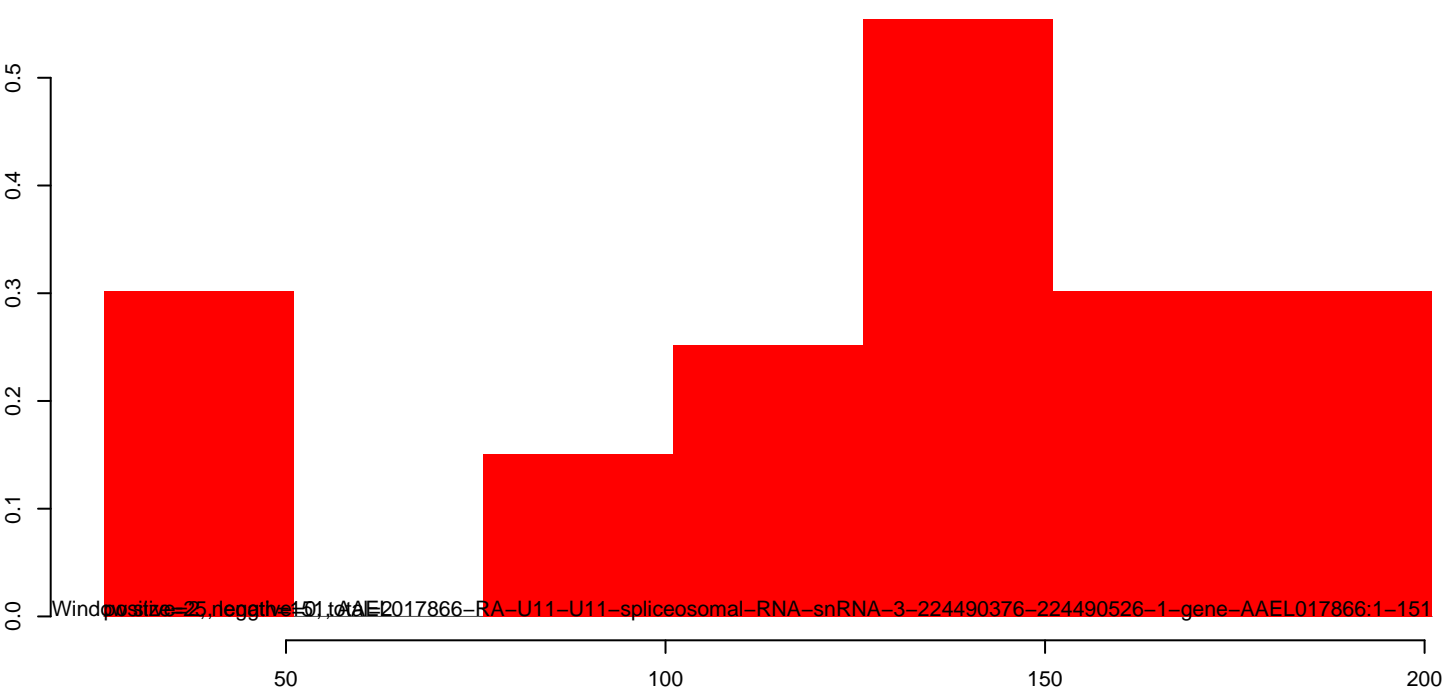
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AeAeg_Fcarc_GH_rep3.24_35.rep

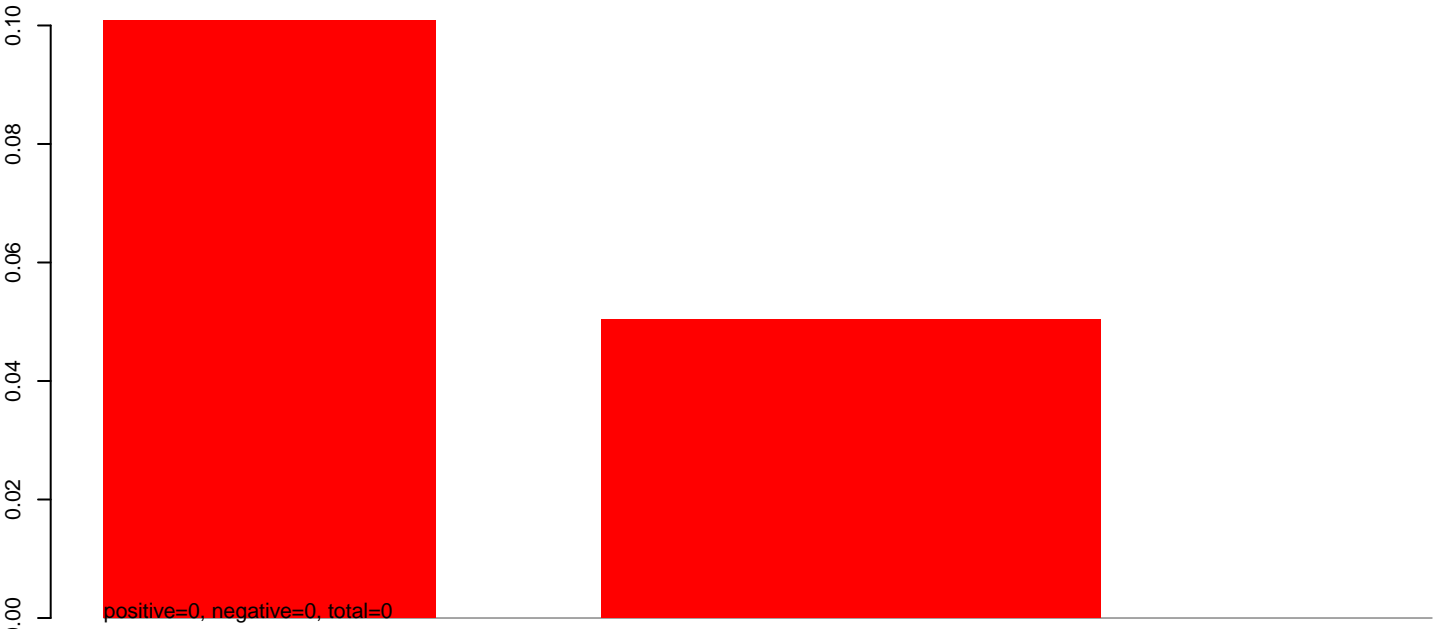


AeAeg_Fcarc_GH_rep3.rep

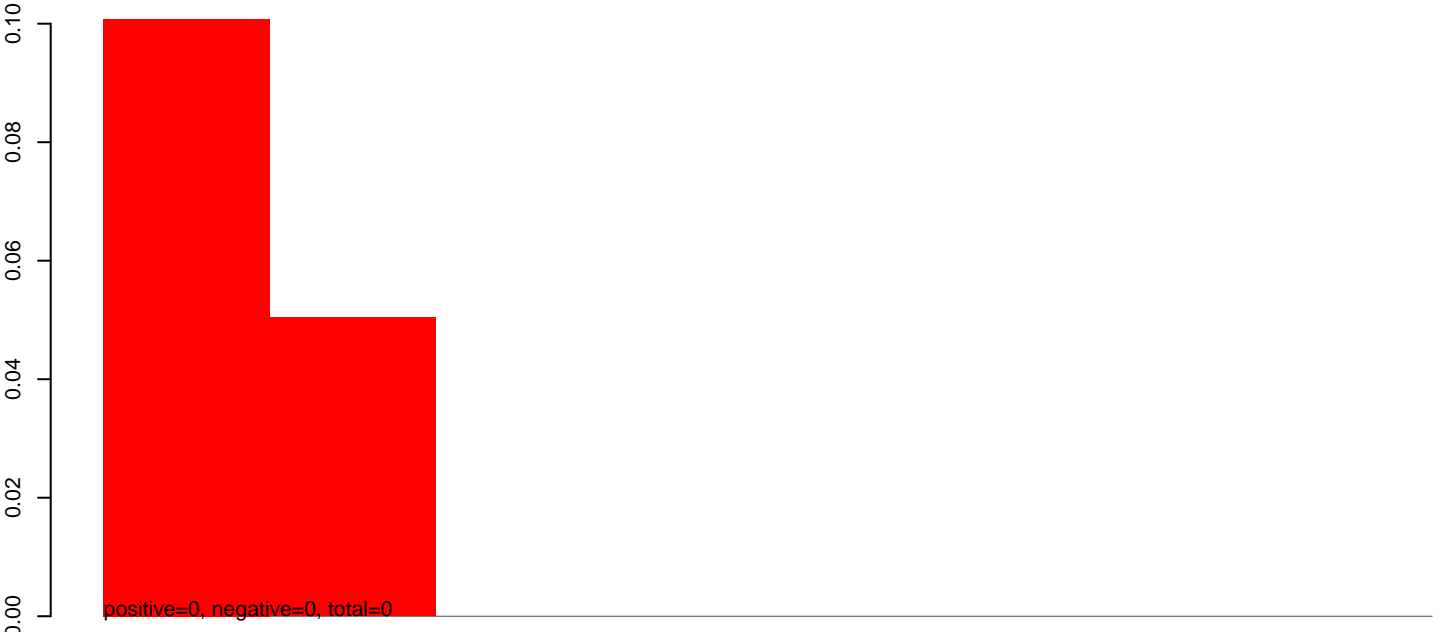


Window=25, negative=0, total=1
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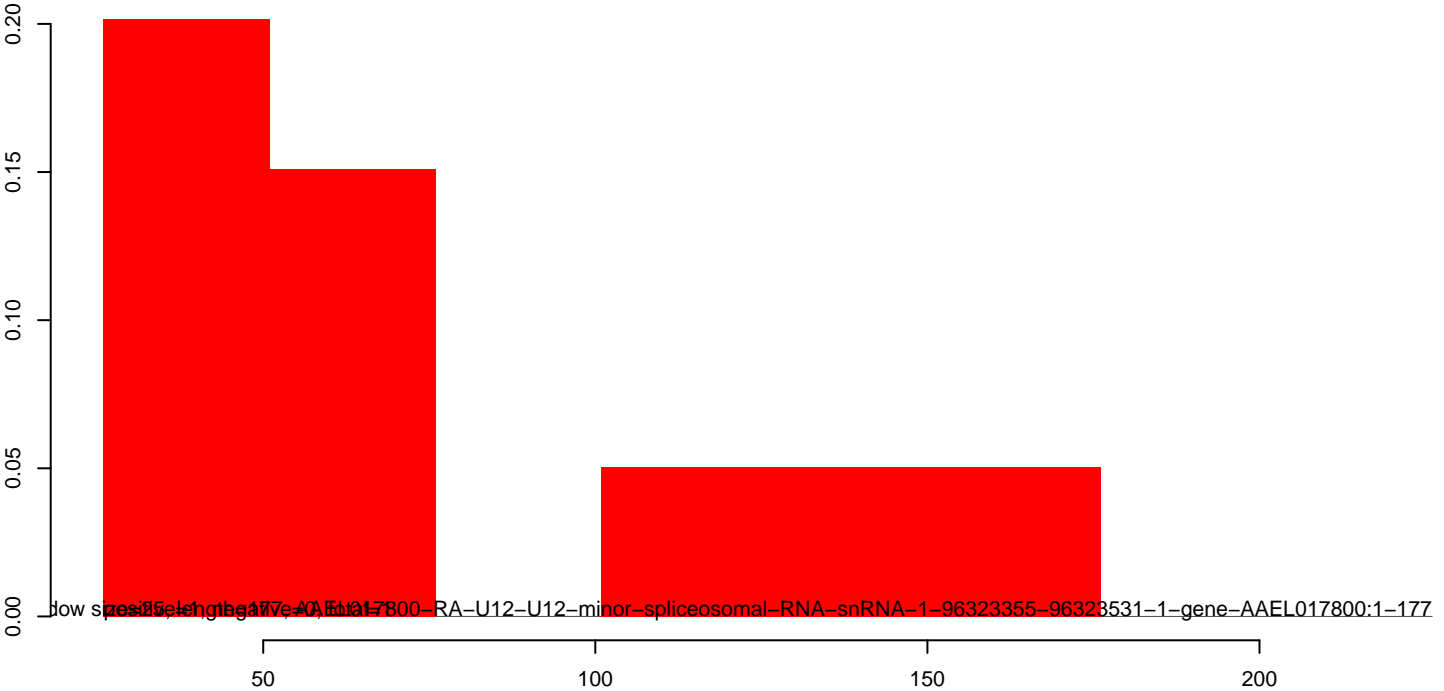
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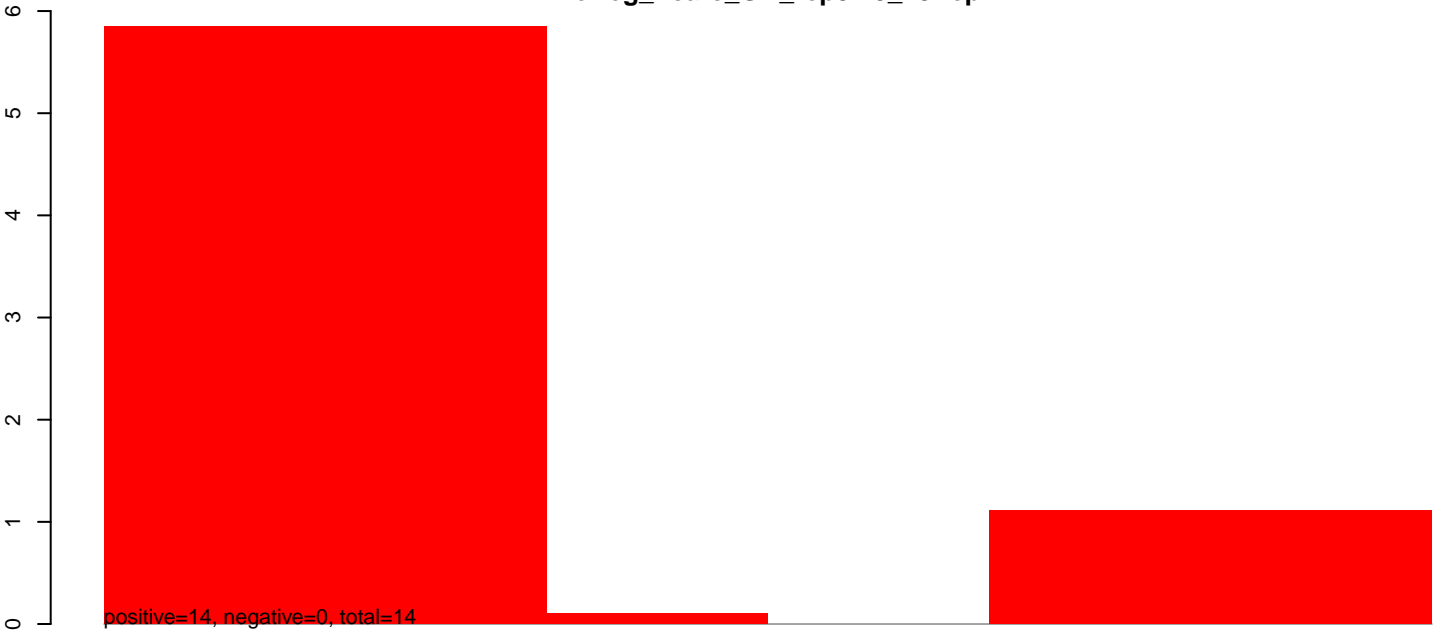
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AeAeg_Fcarc_GH_rep3.rep



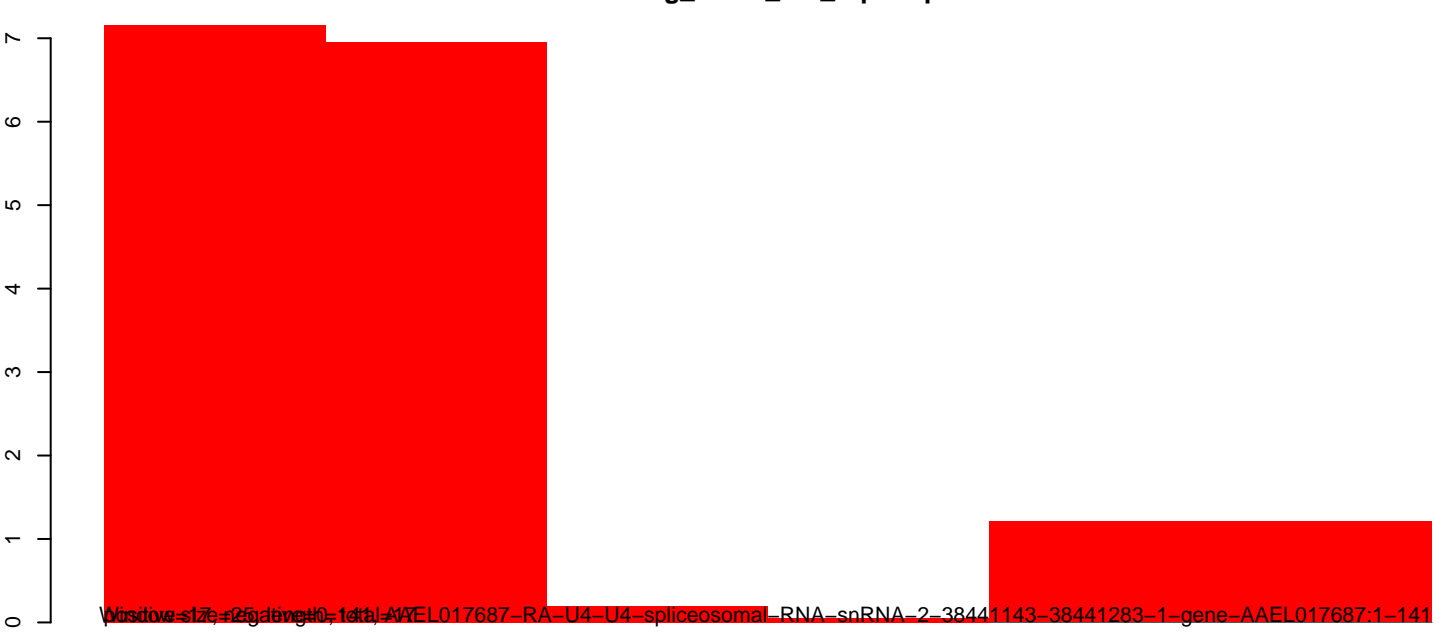
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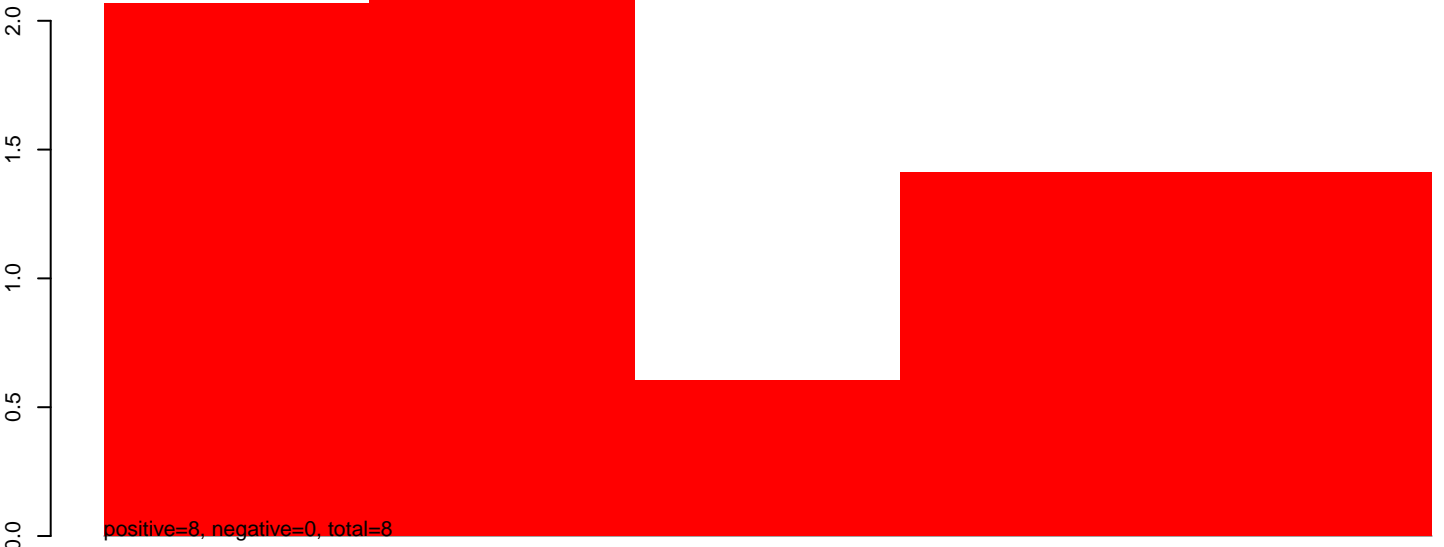
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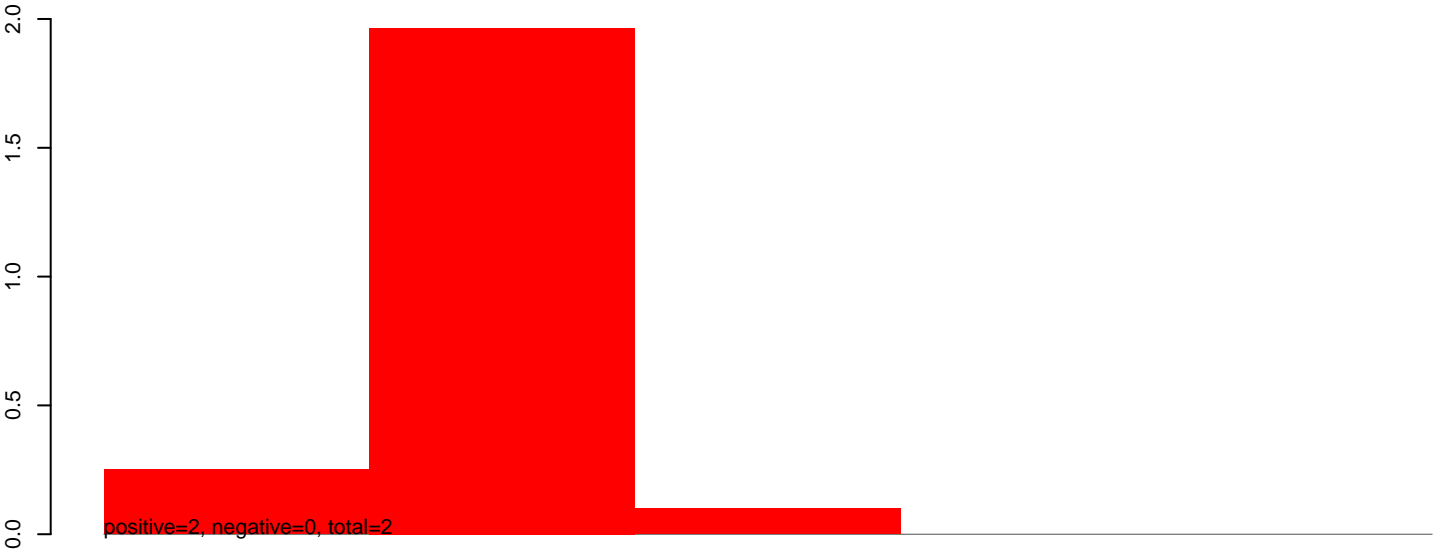
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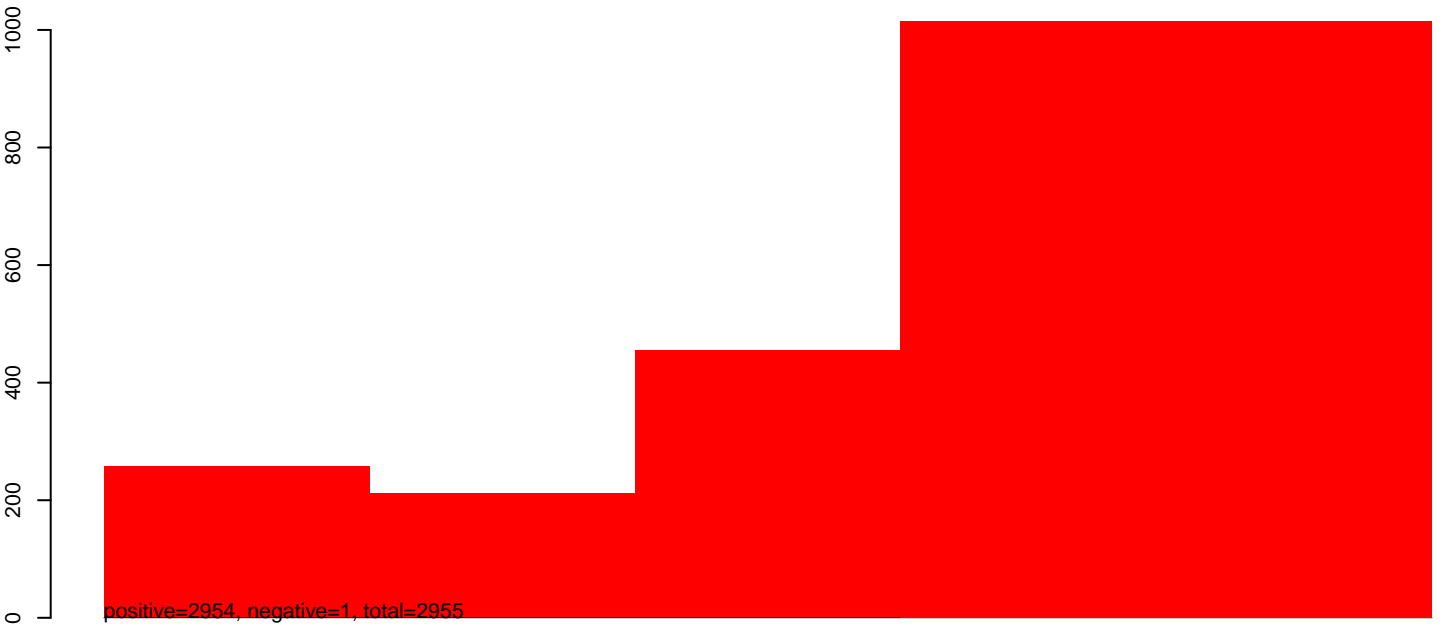
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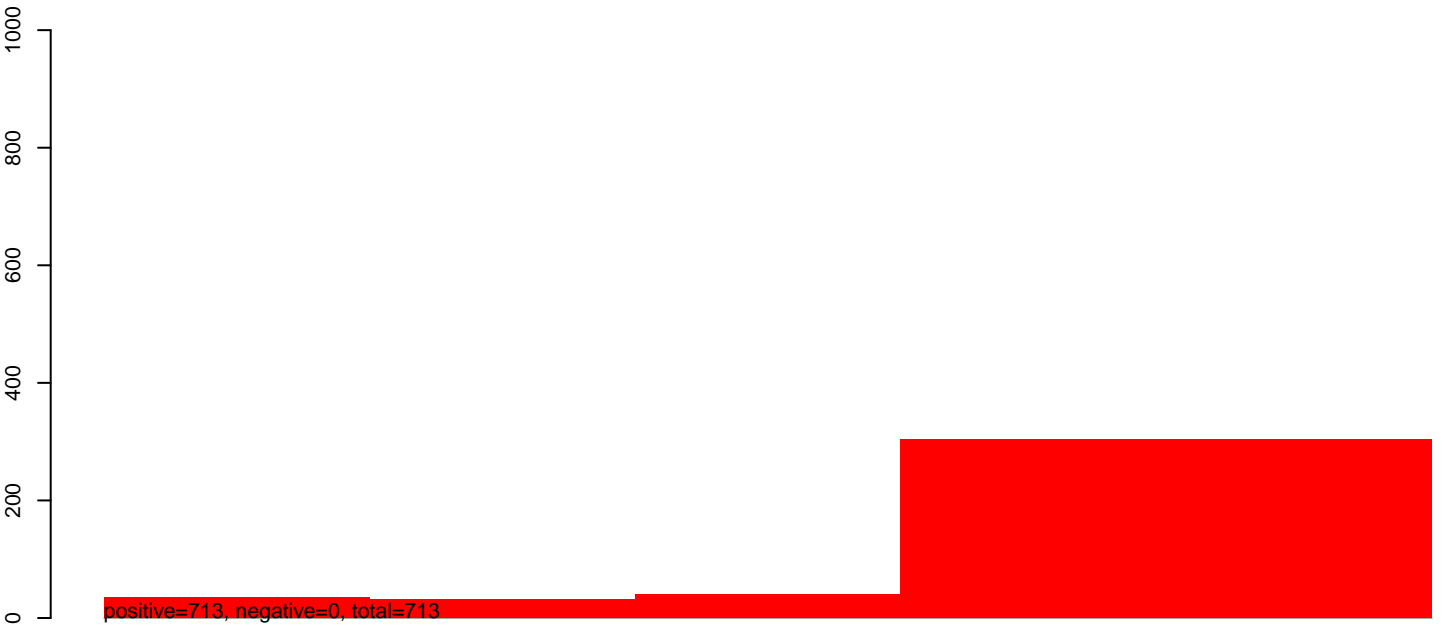
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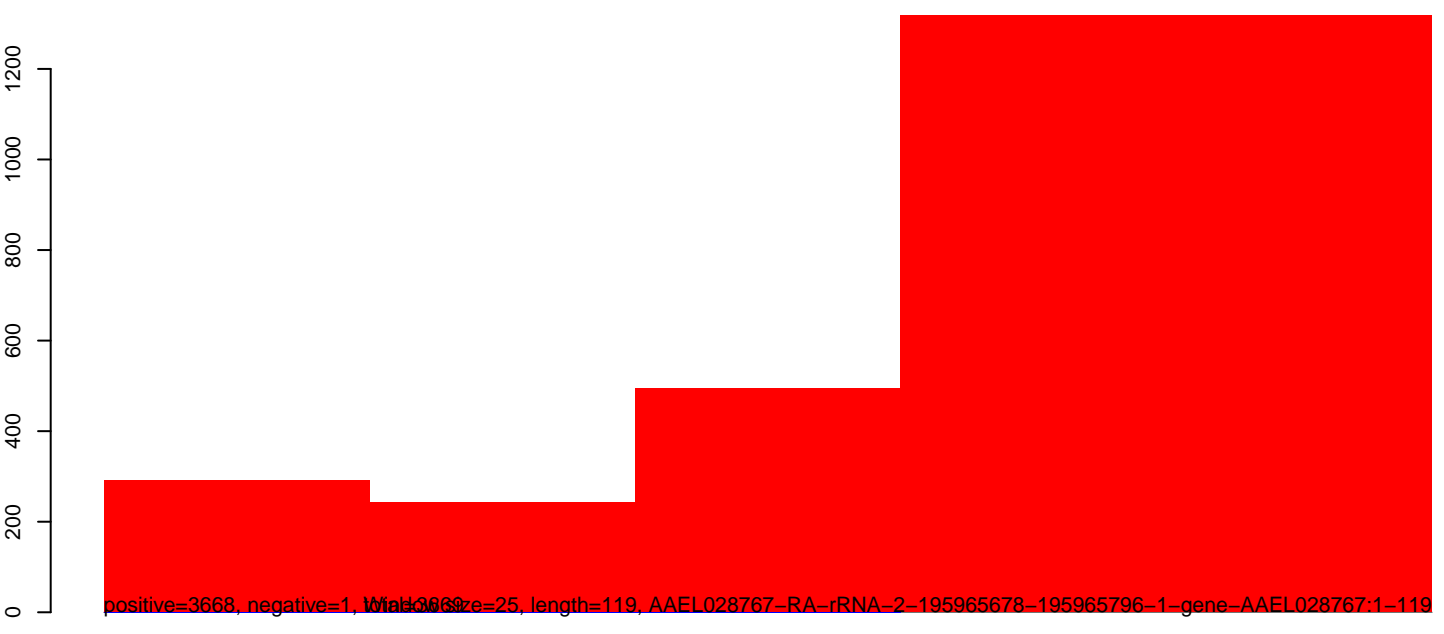
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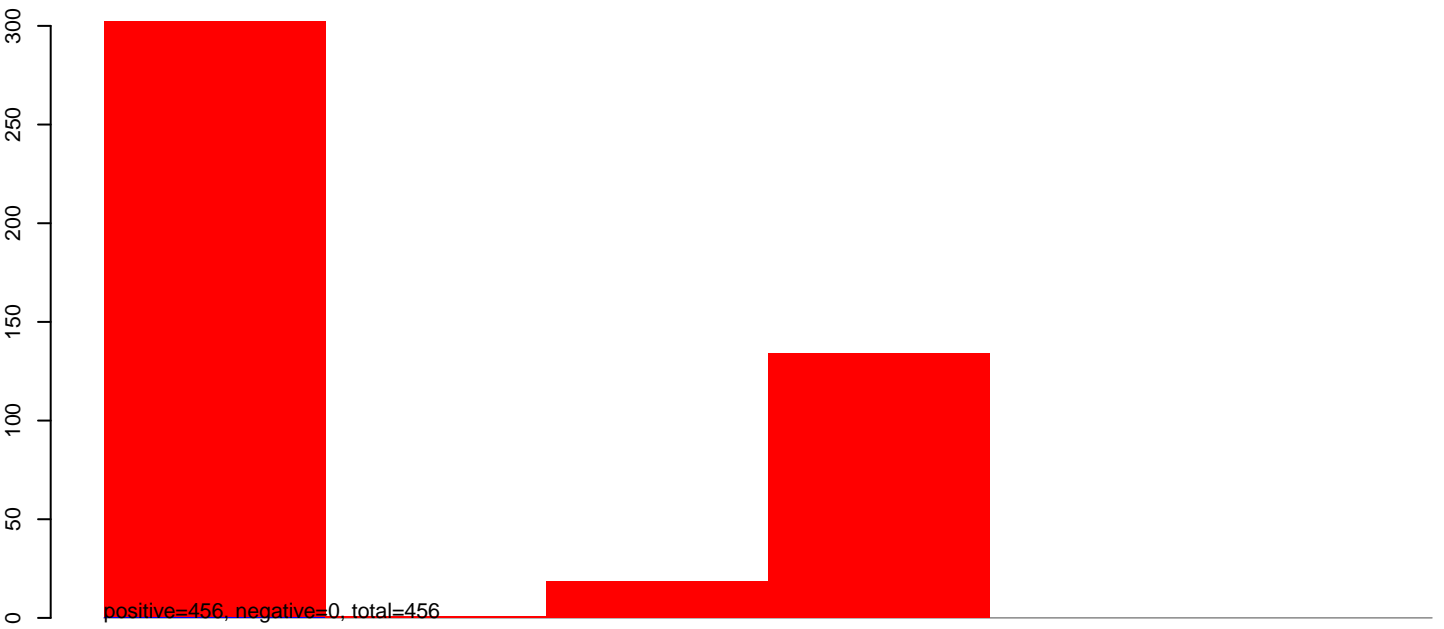
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AeAeg_Fcarc_GH_rep3.rep



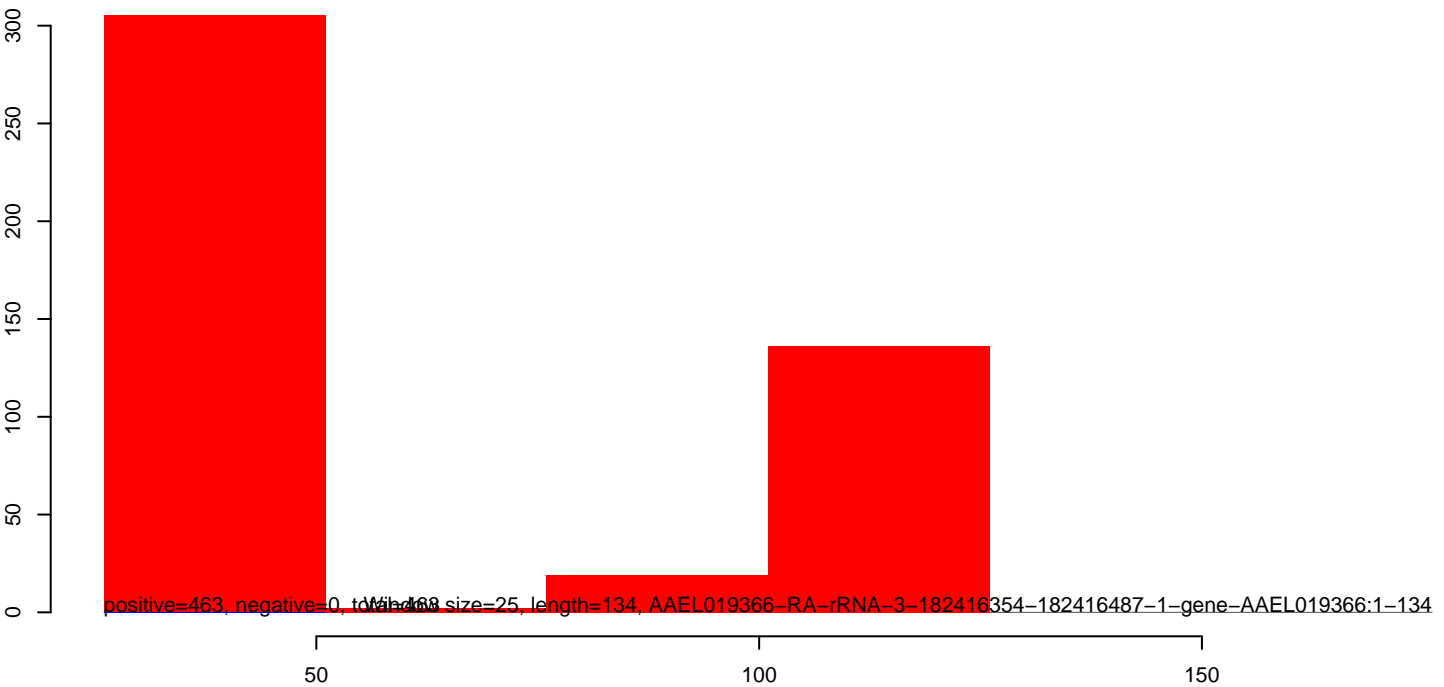
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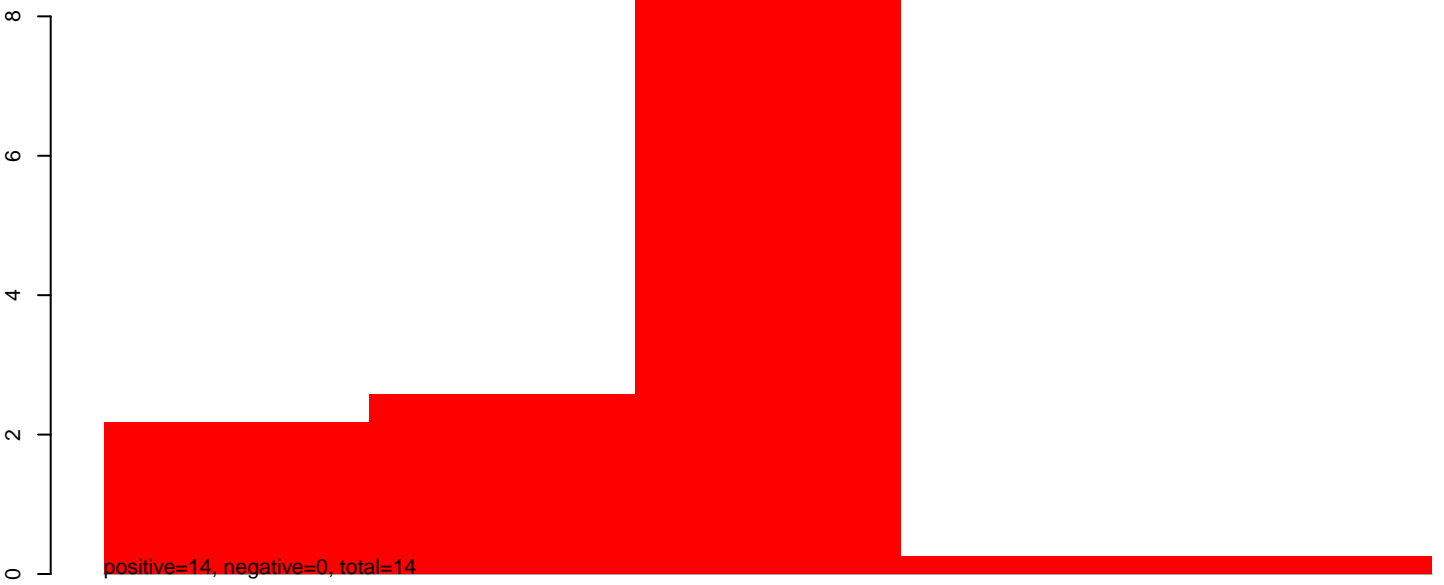
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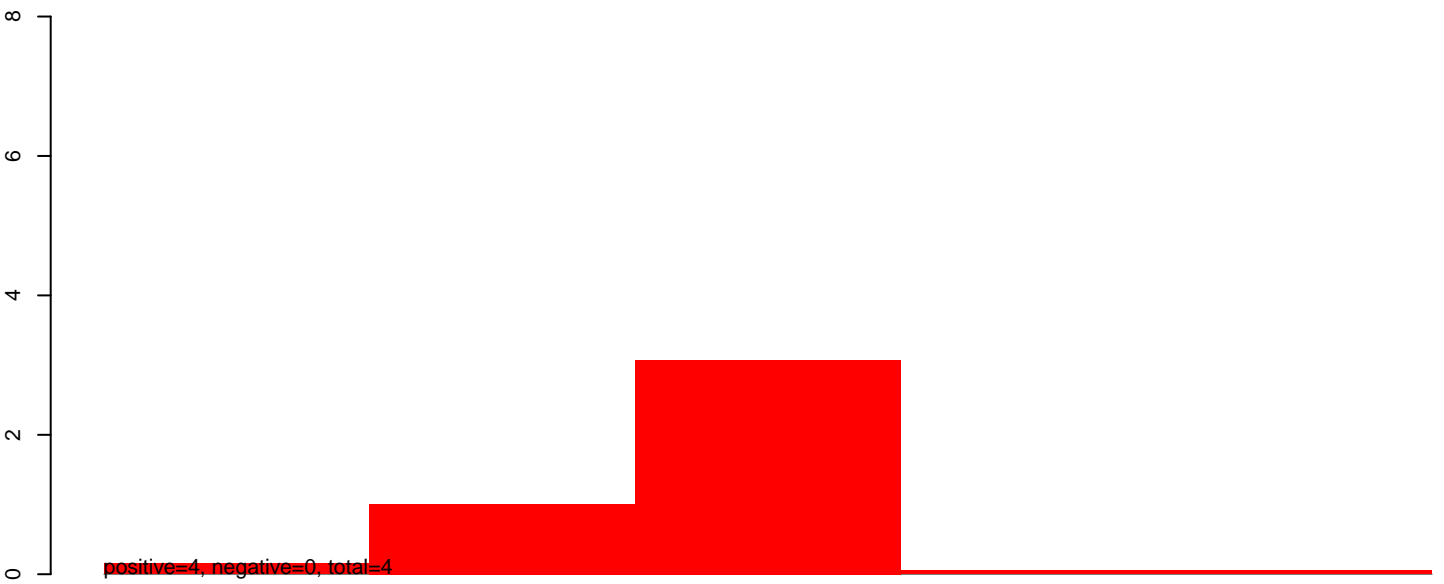
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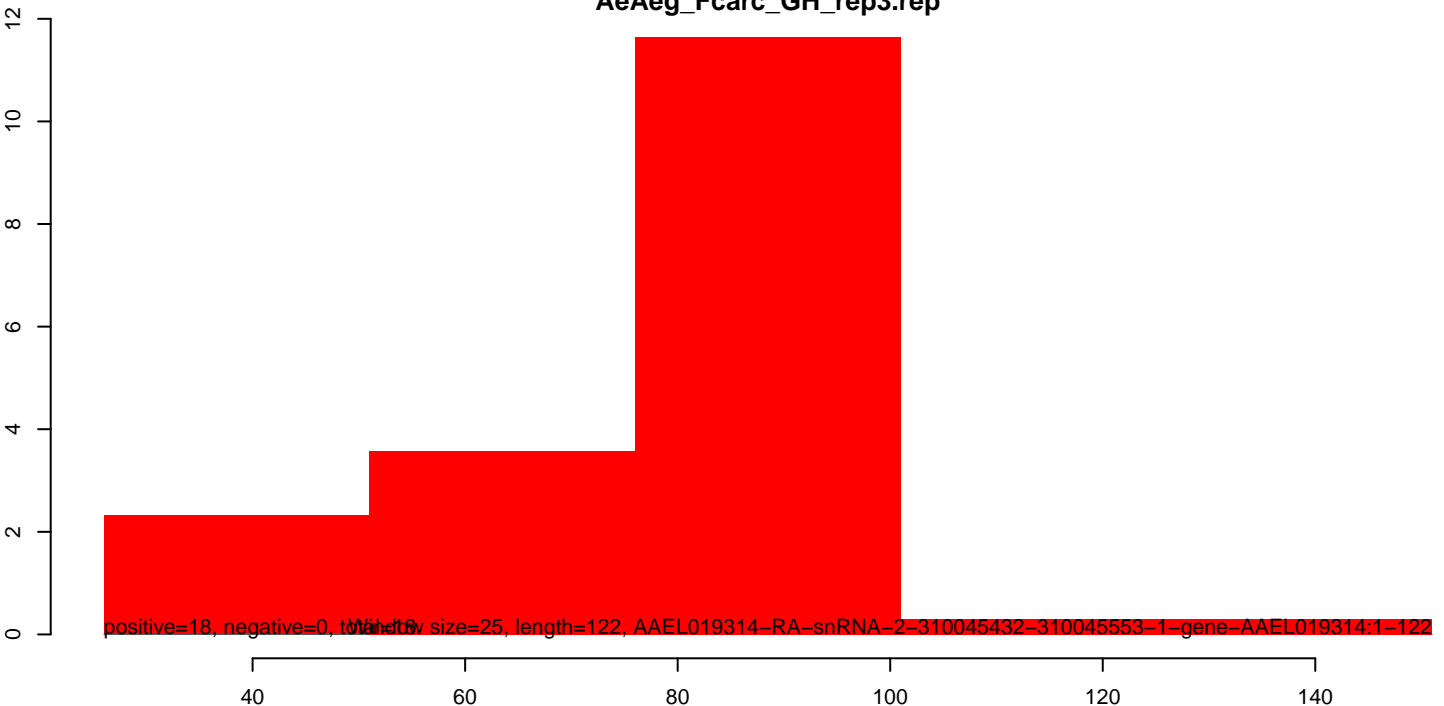
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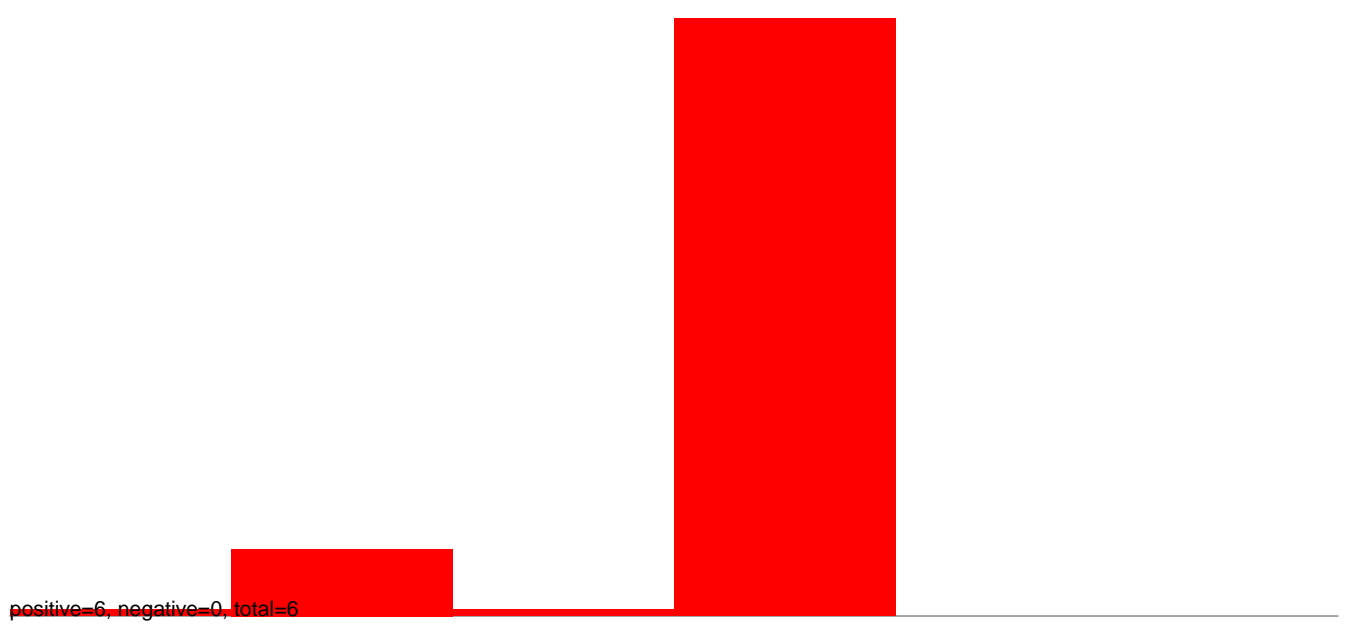
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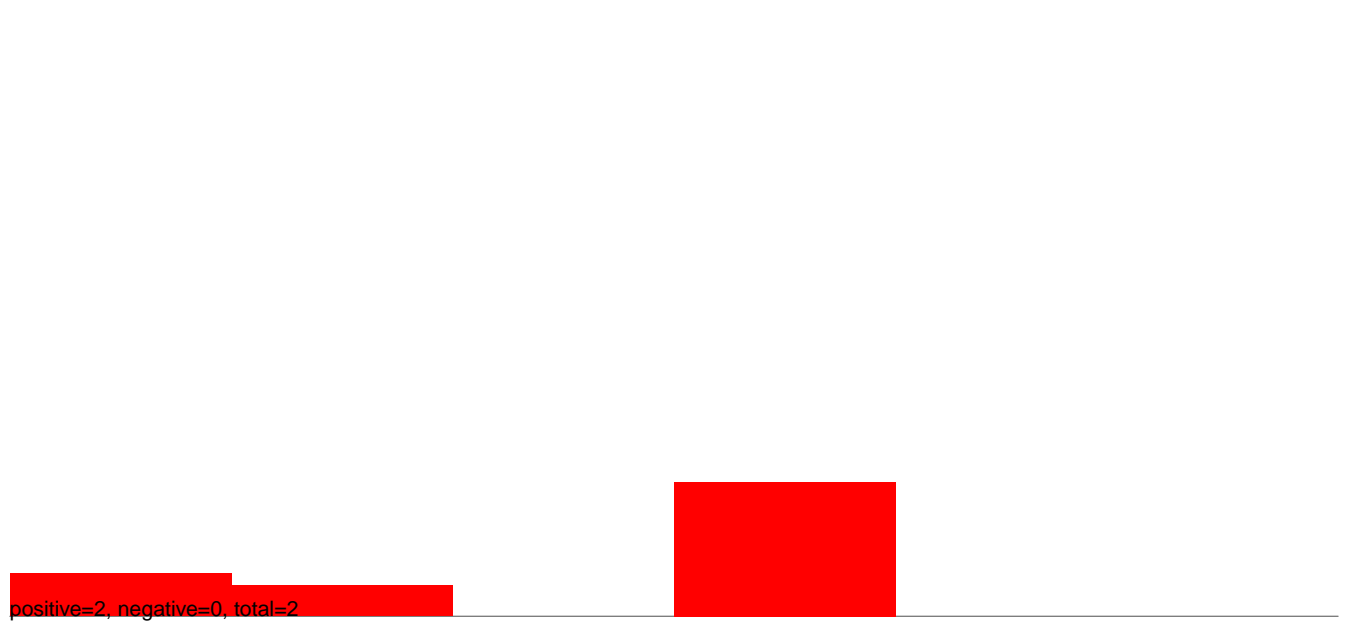
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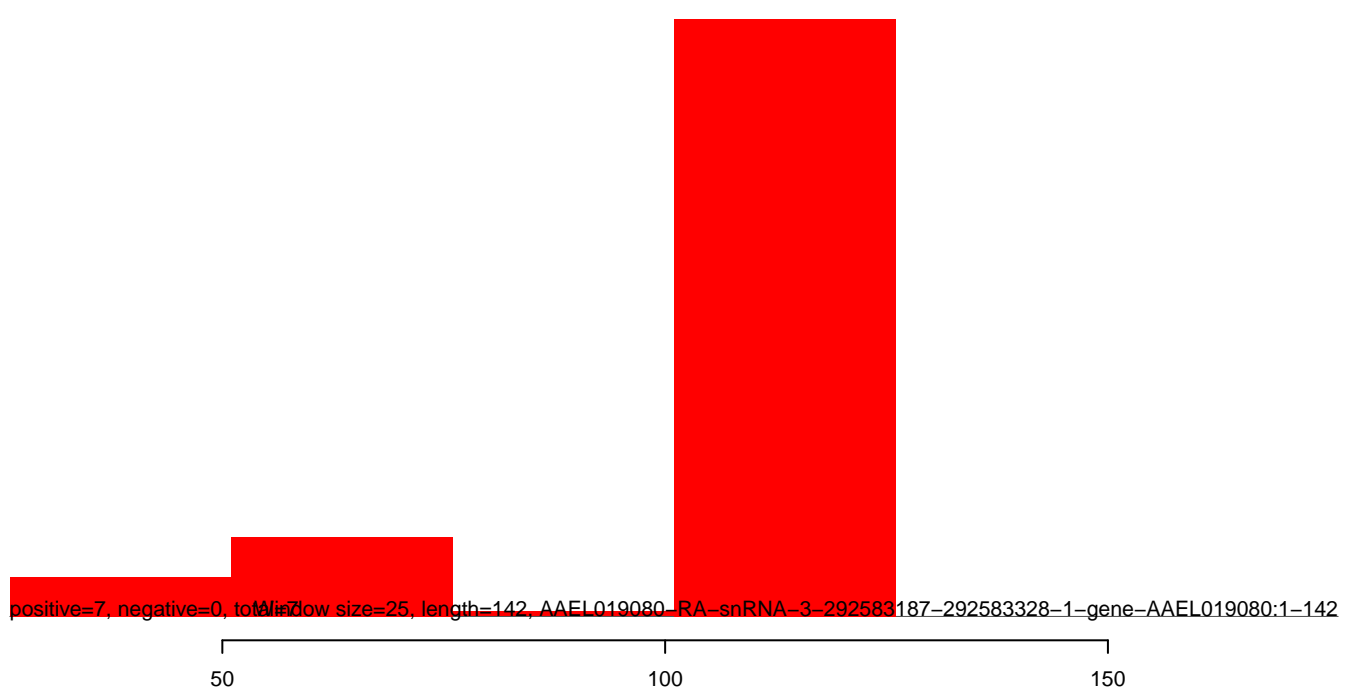
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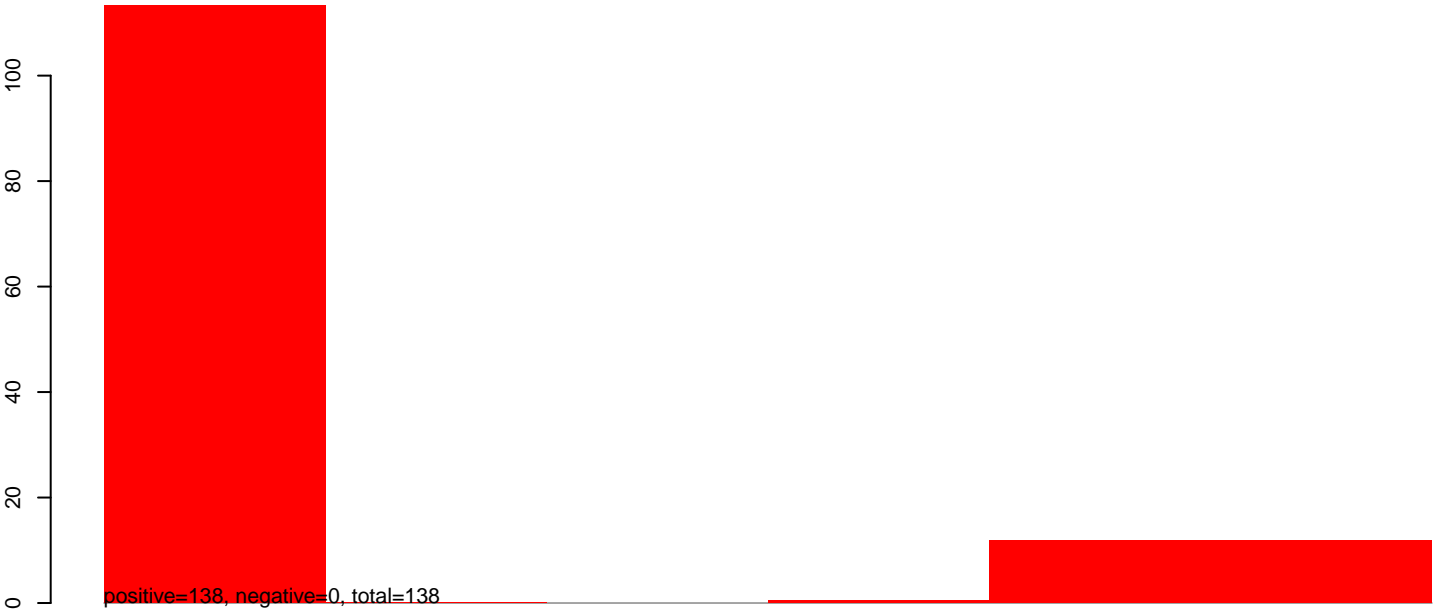
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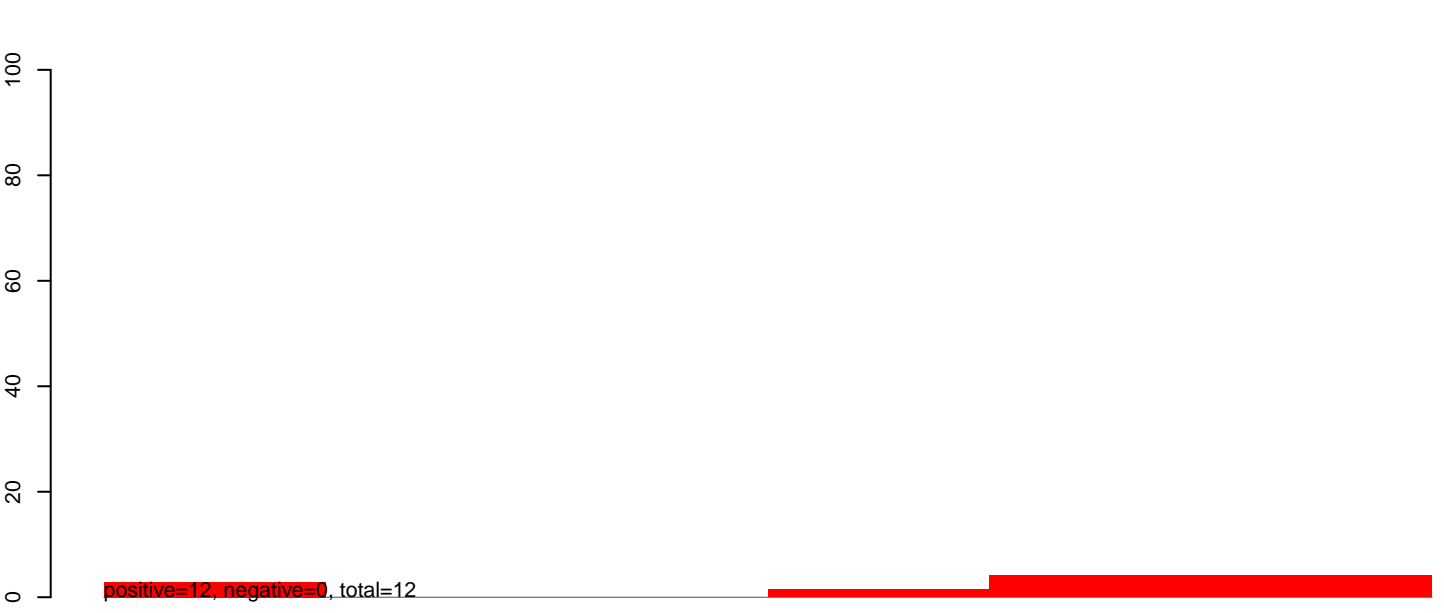
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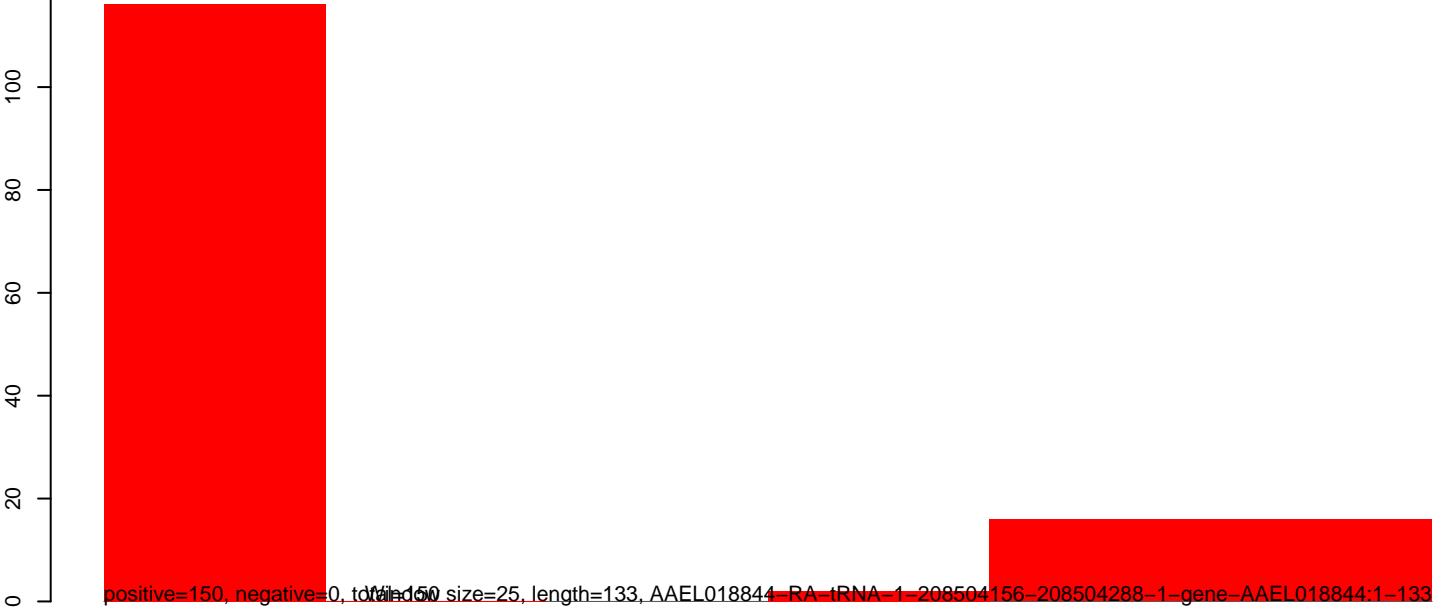
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AeAeg_Fcarc_GH_rep3.24_35.rep

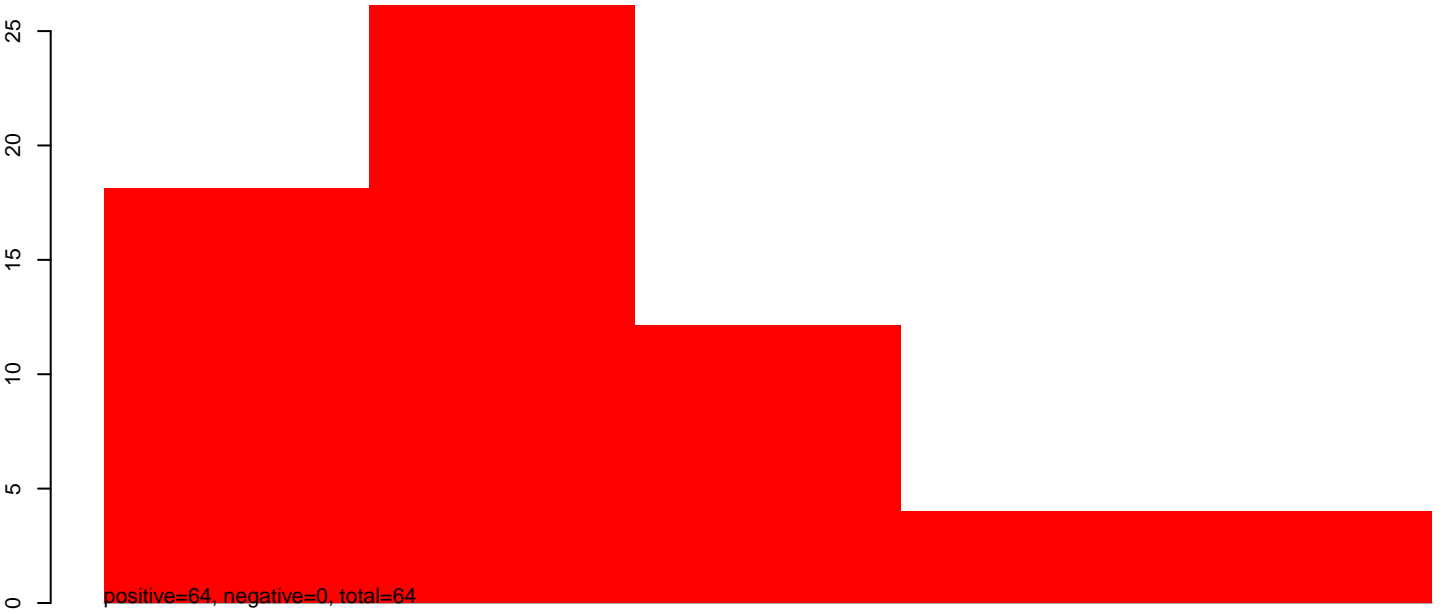


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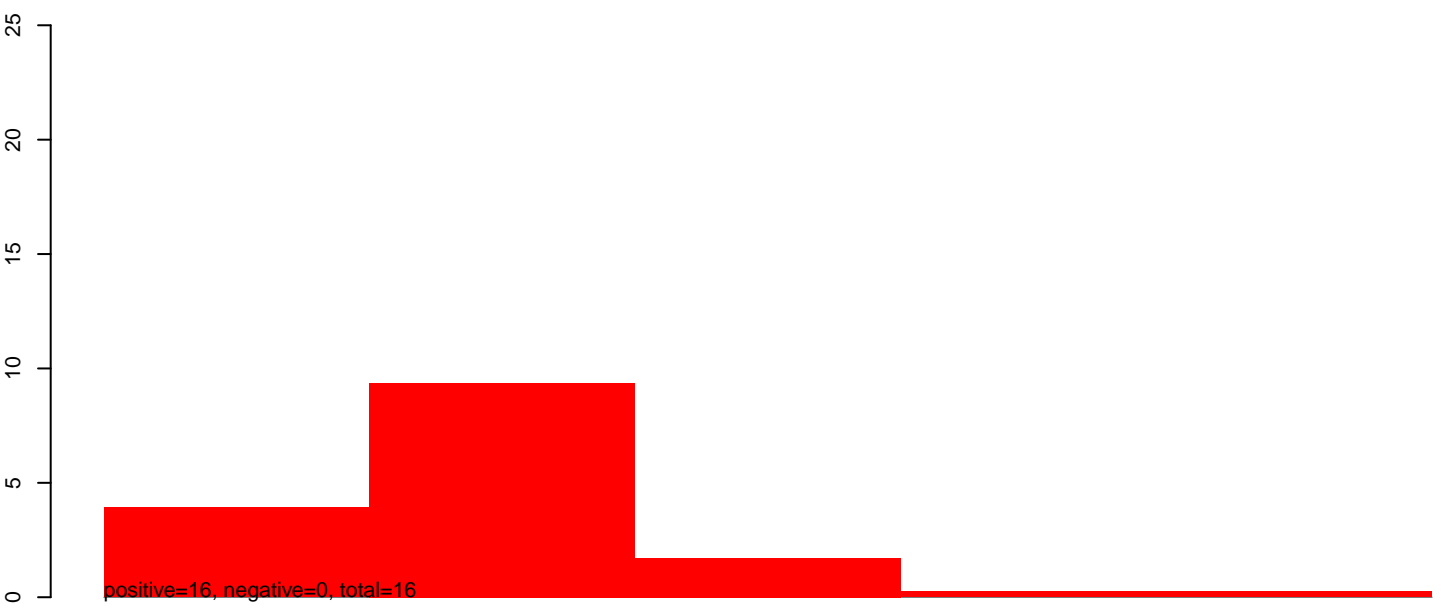


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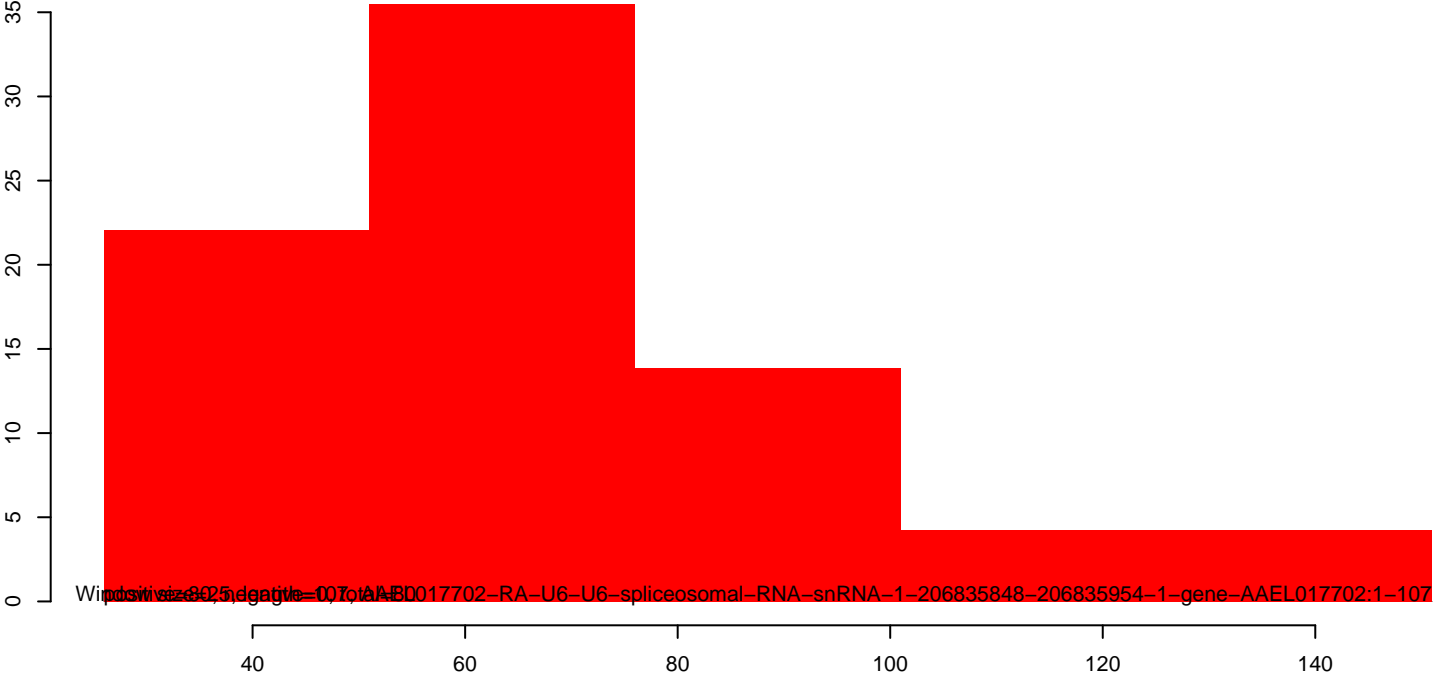
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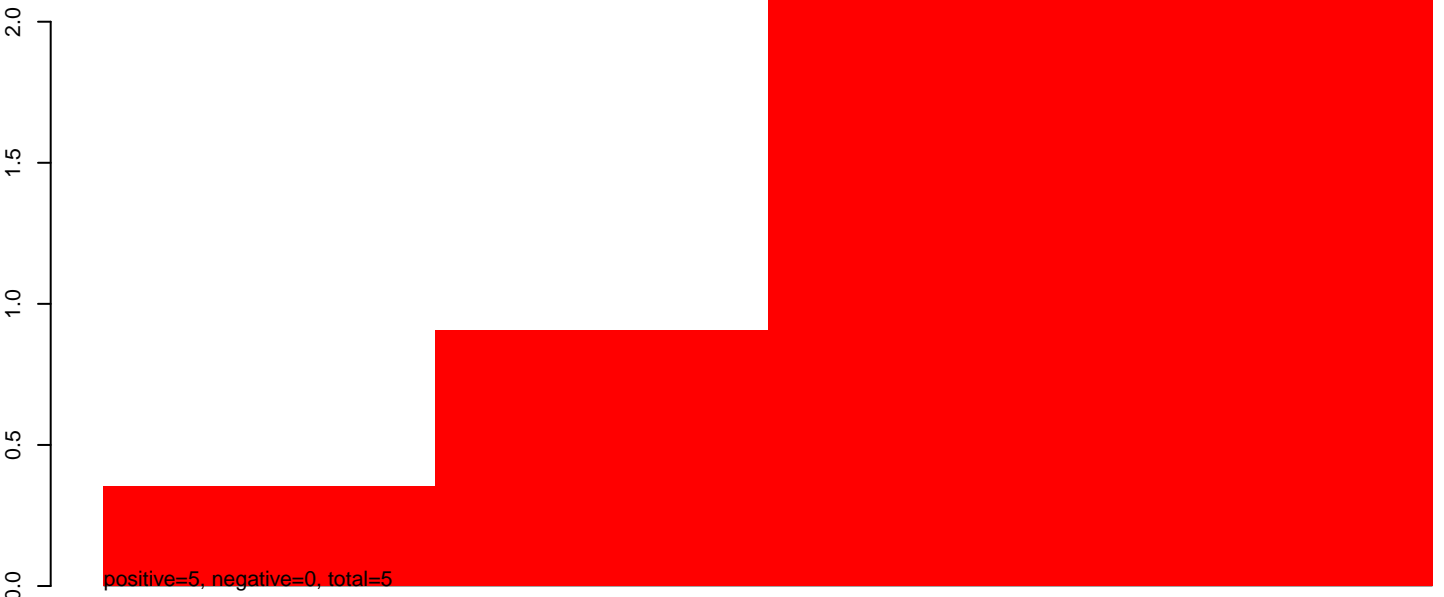
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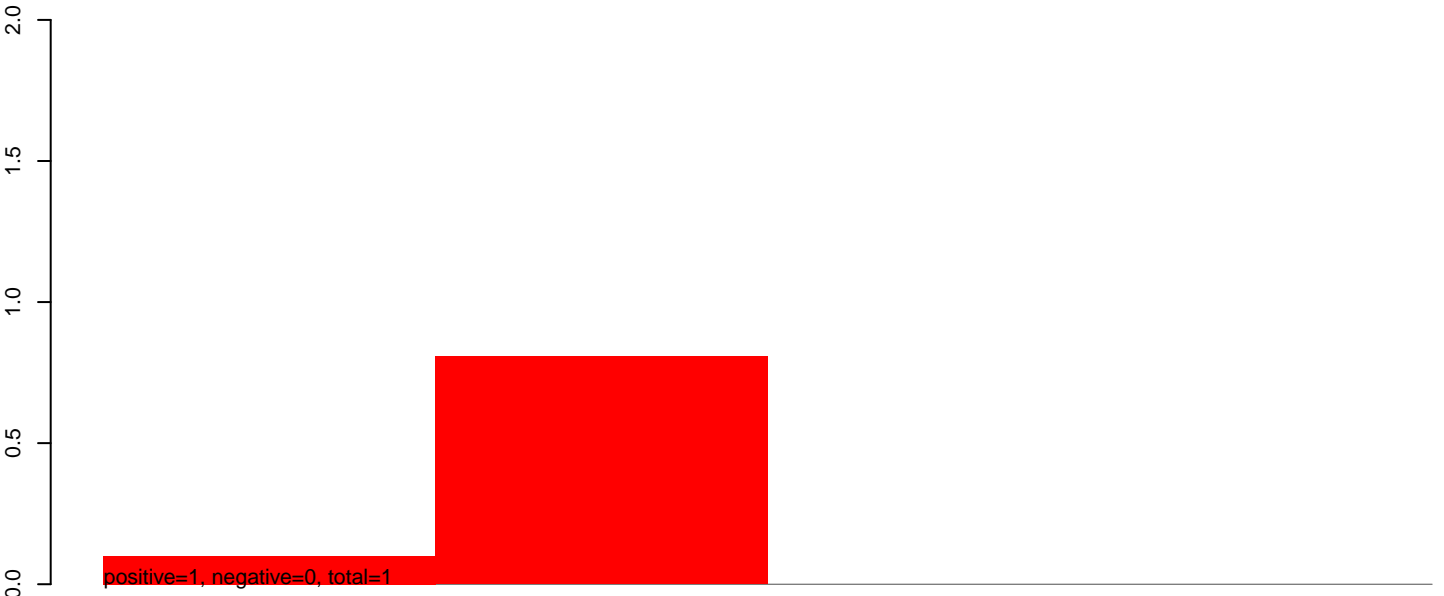
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AeAeg_Fcarc_GH_rep3.24_35.rep

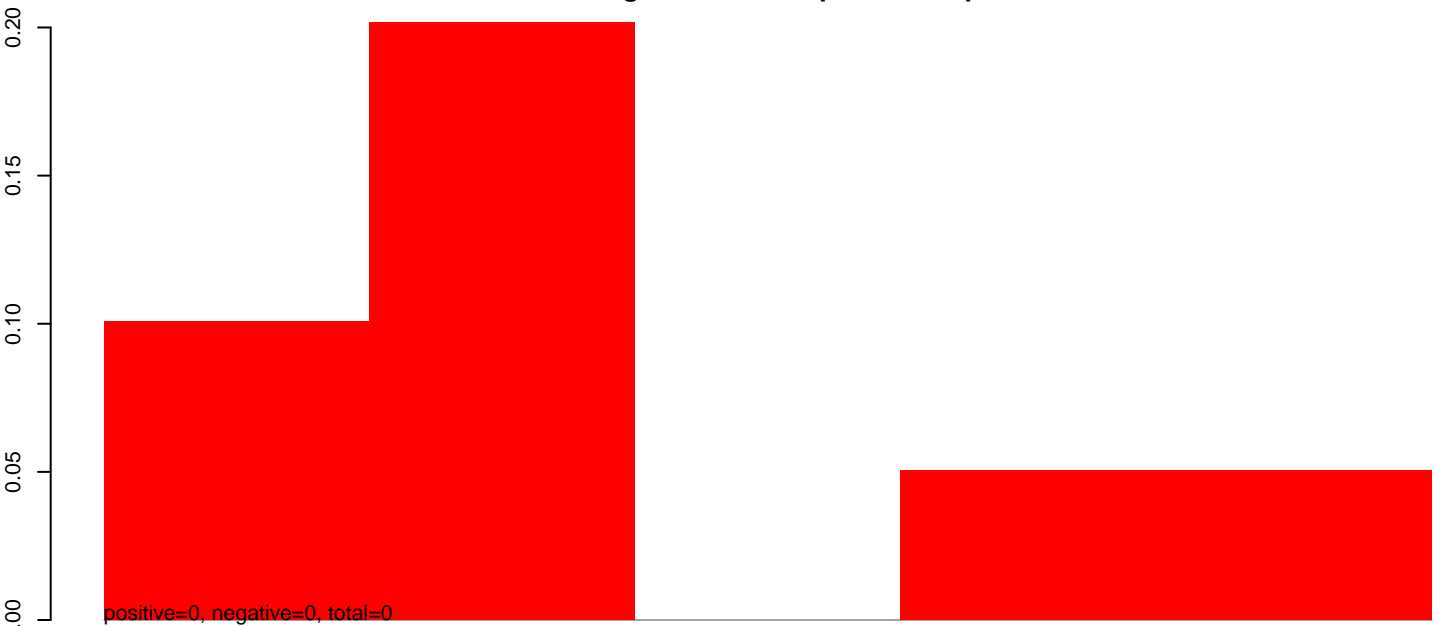


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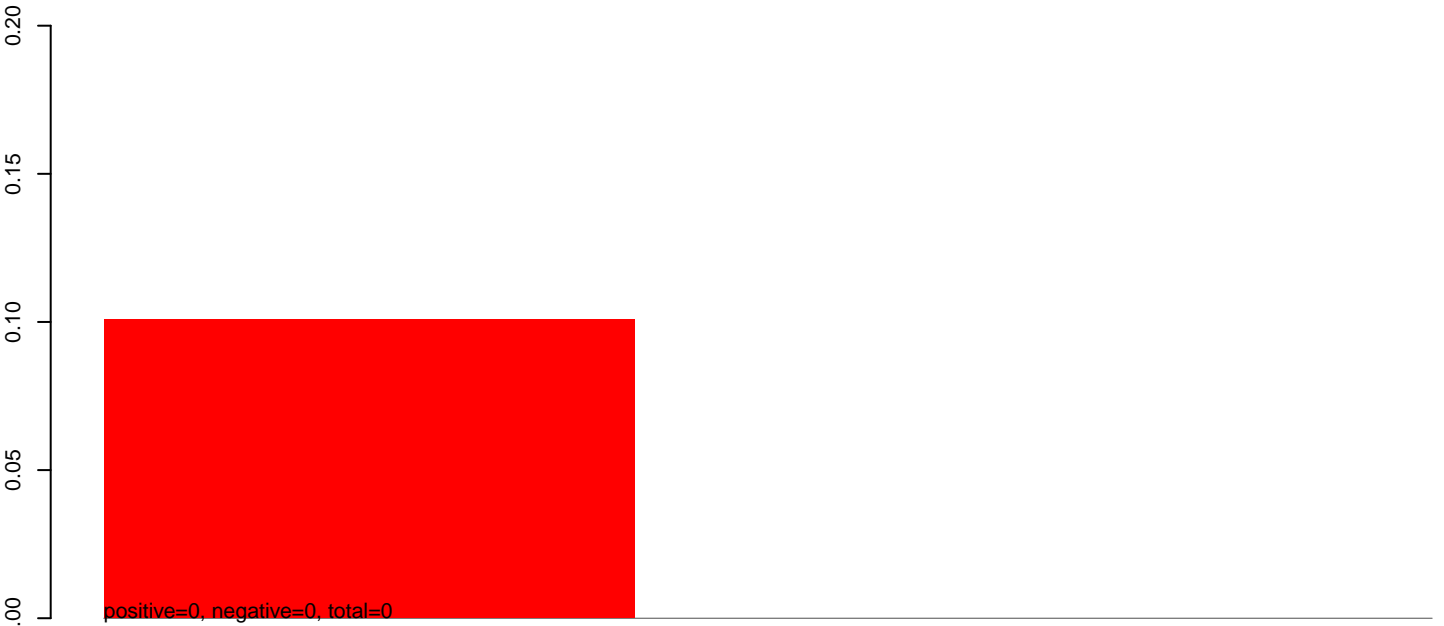


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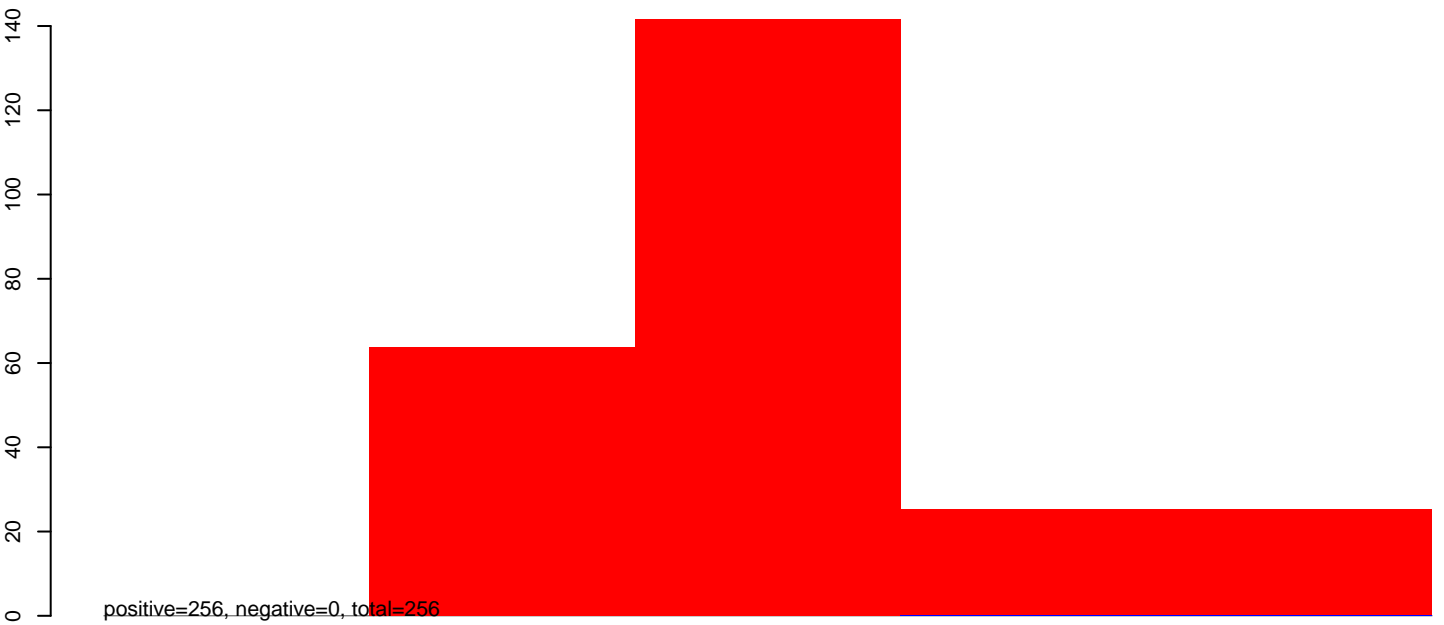
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AeAeg_Fcarc_GH_rep3.rep



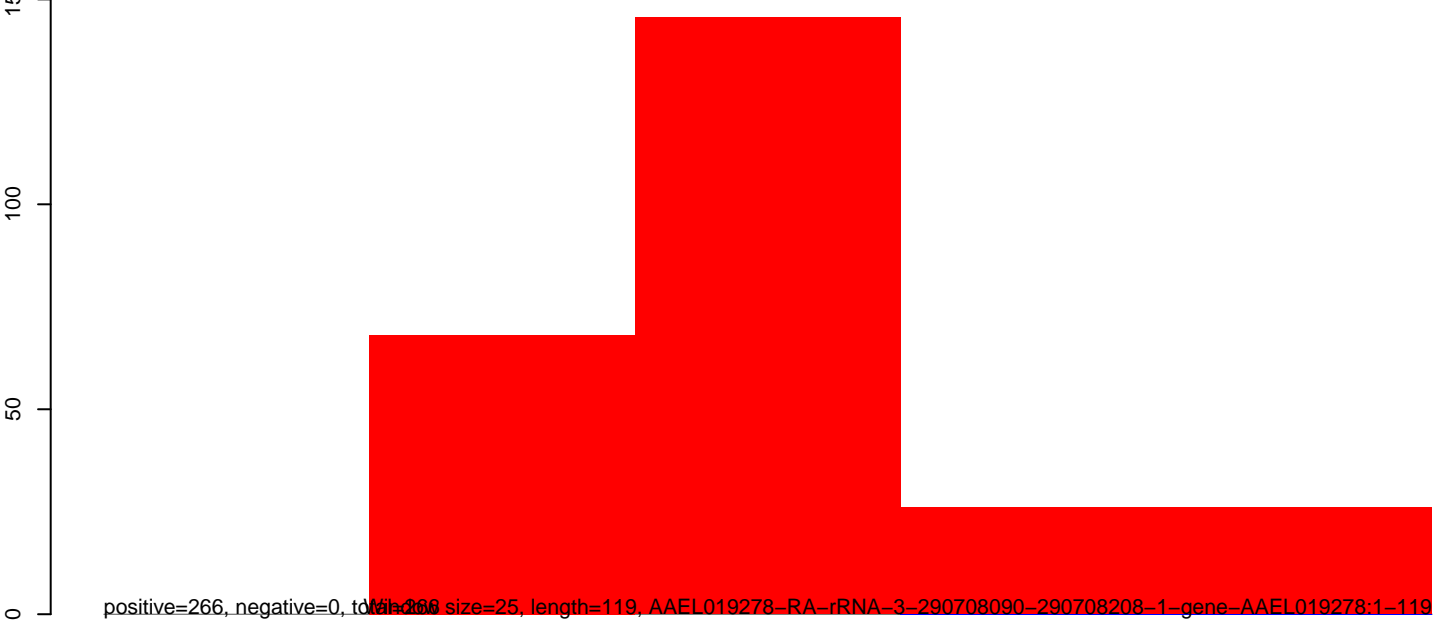
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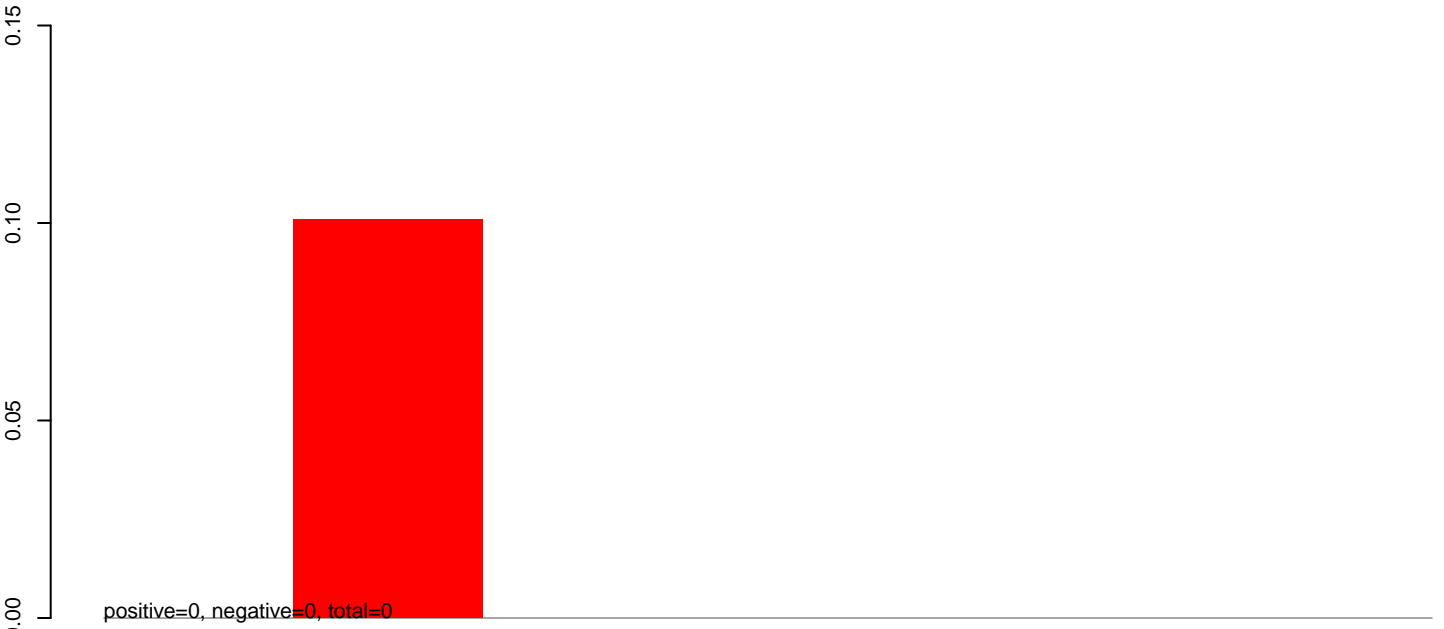


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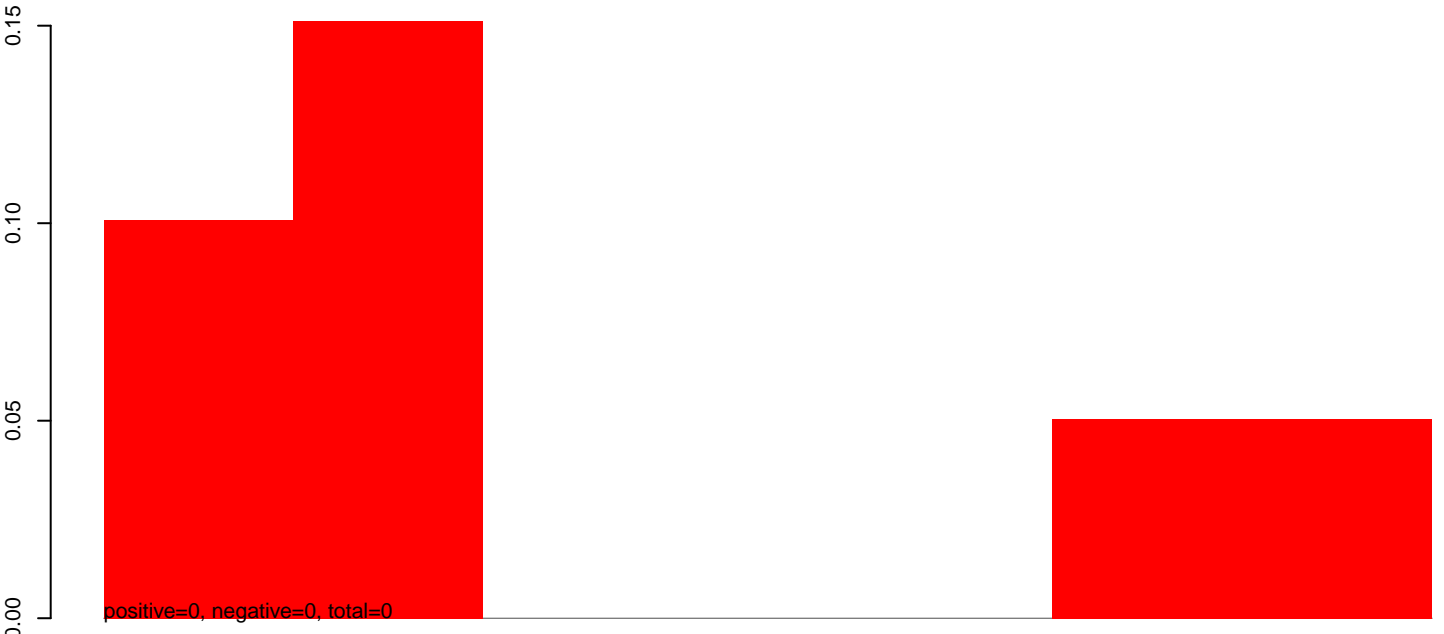


40 60 80 100 120 140

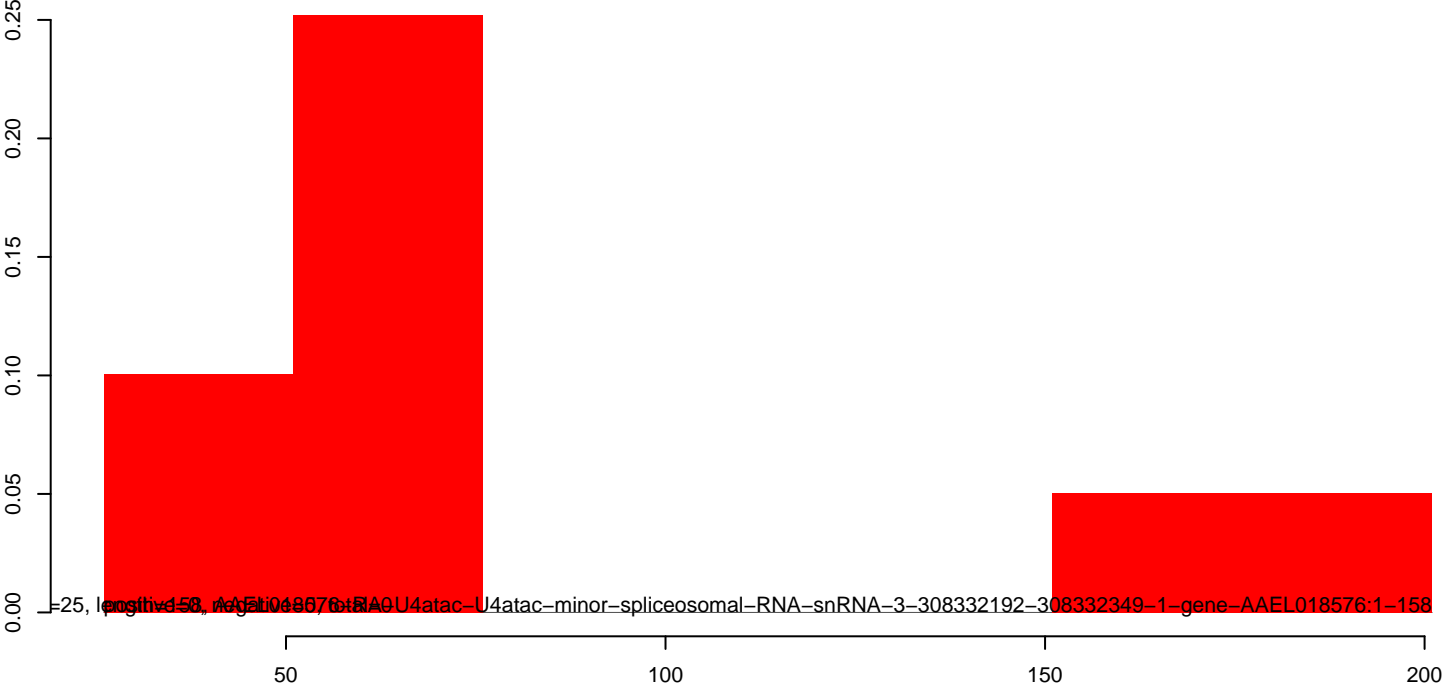
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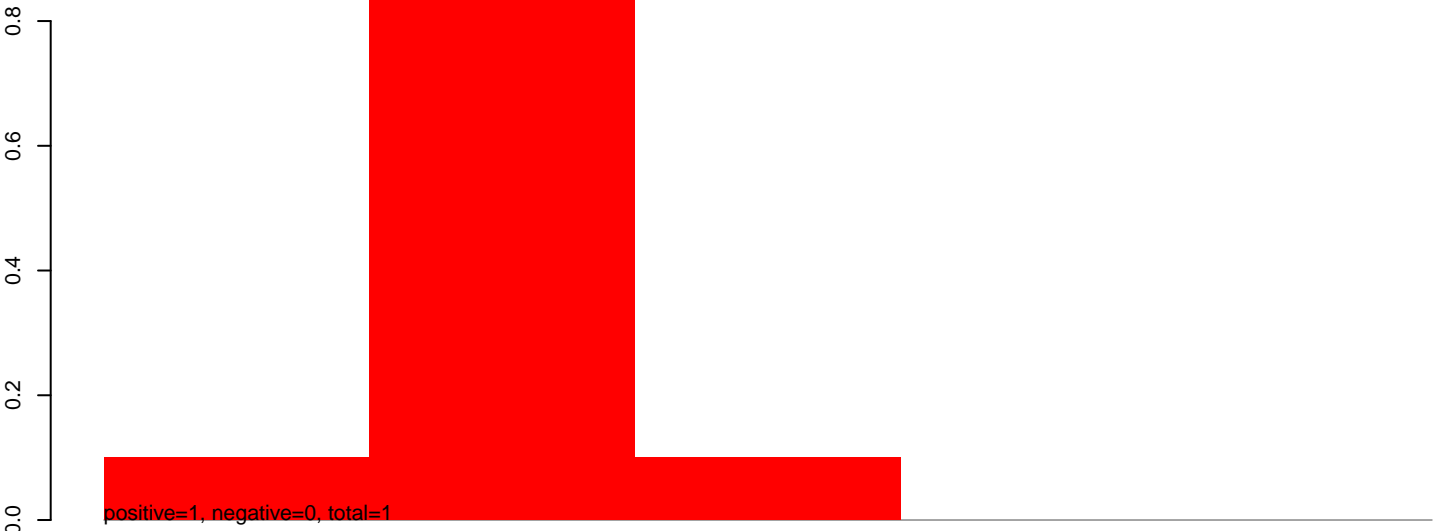
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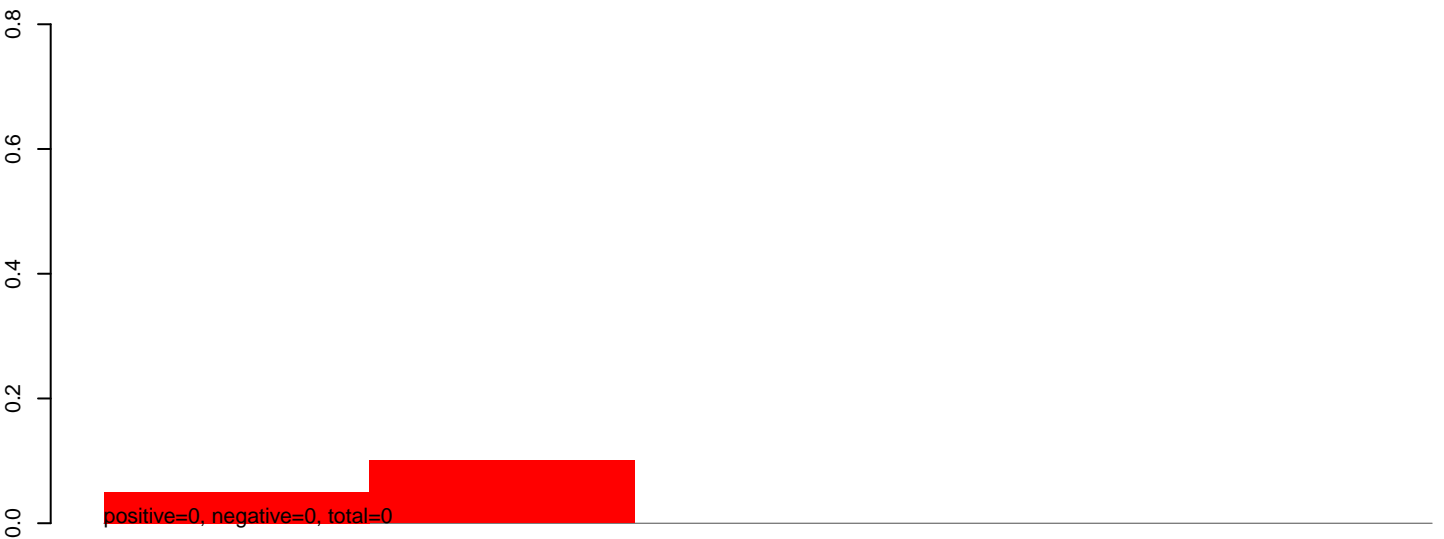
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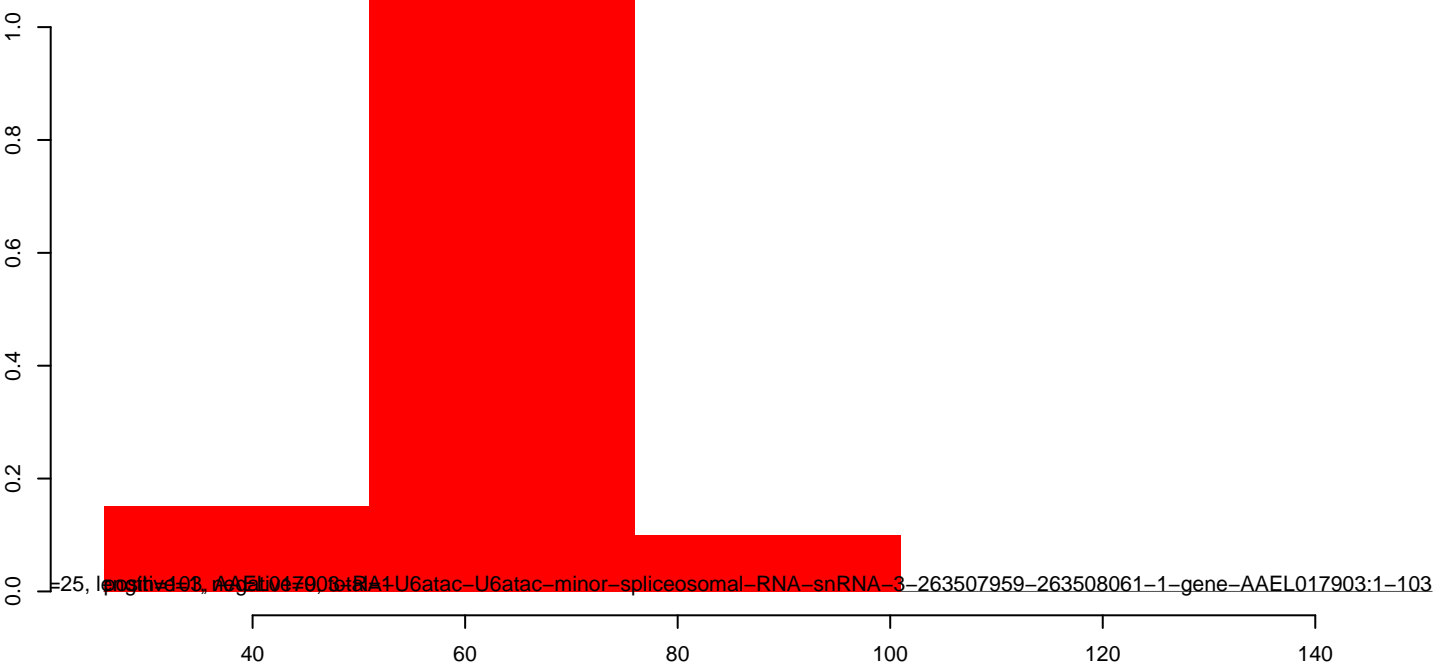
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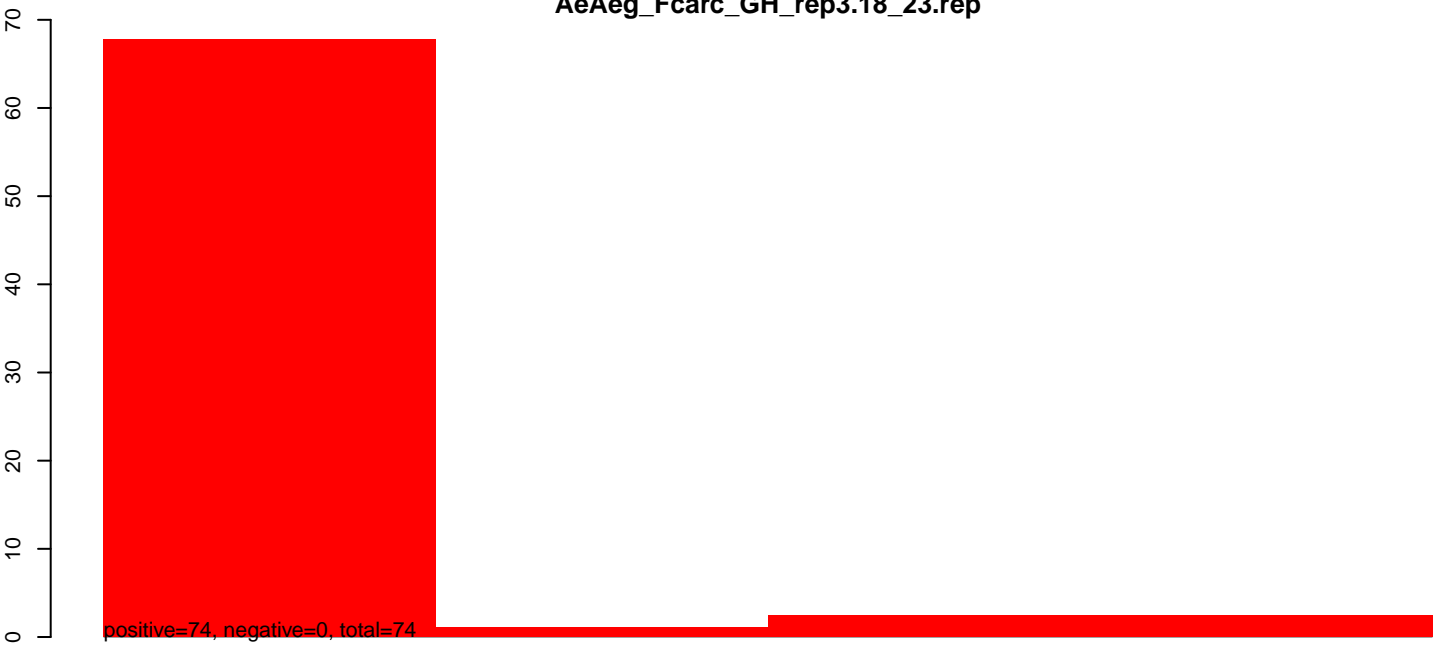
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AeAeg_Fcarc_GH_rep3.rep



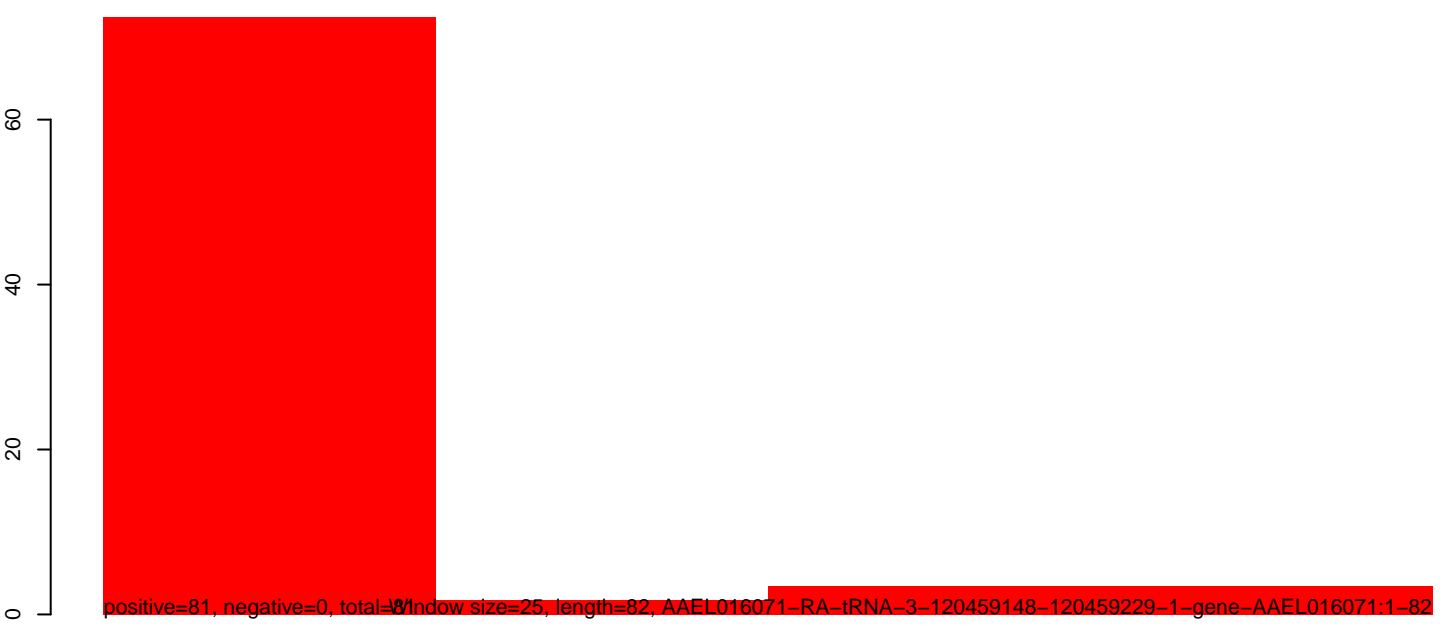
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AeAeg_Fcarc_GH_rep3.24_35.rep

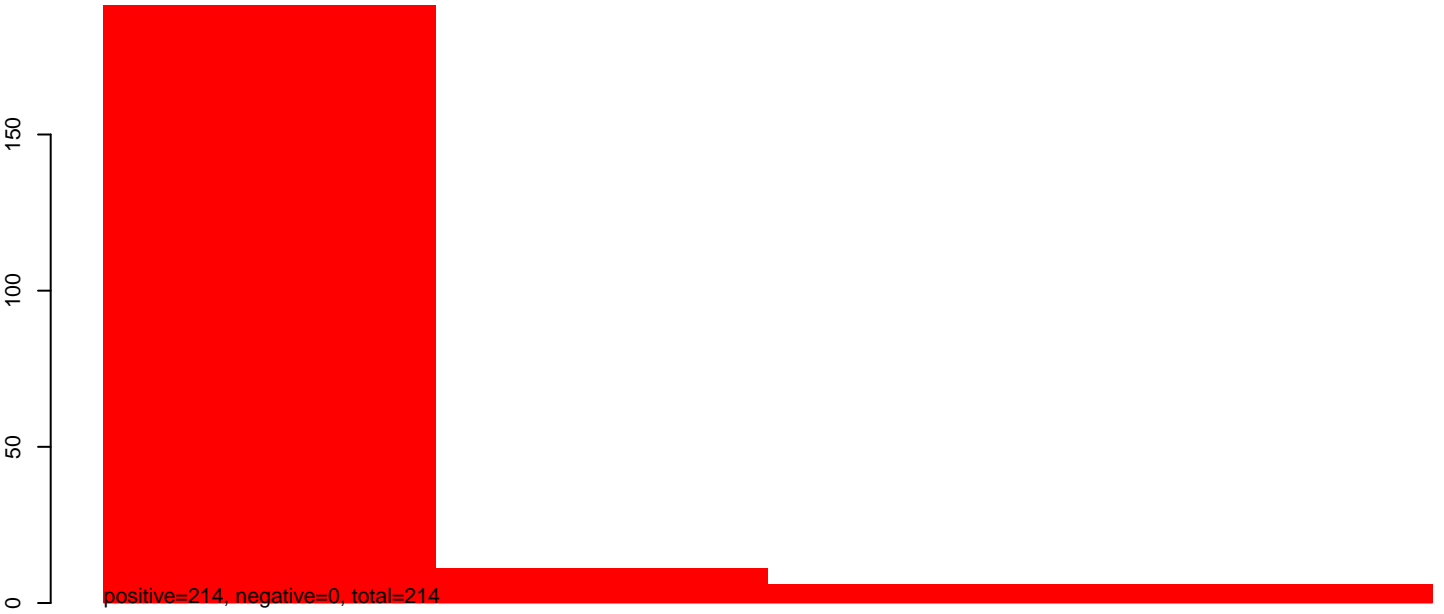


AeAeg_Fcarc_GH_rep3.rep

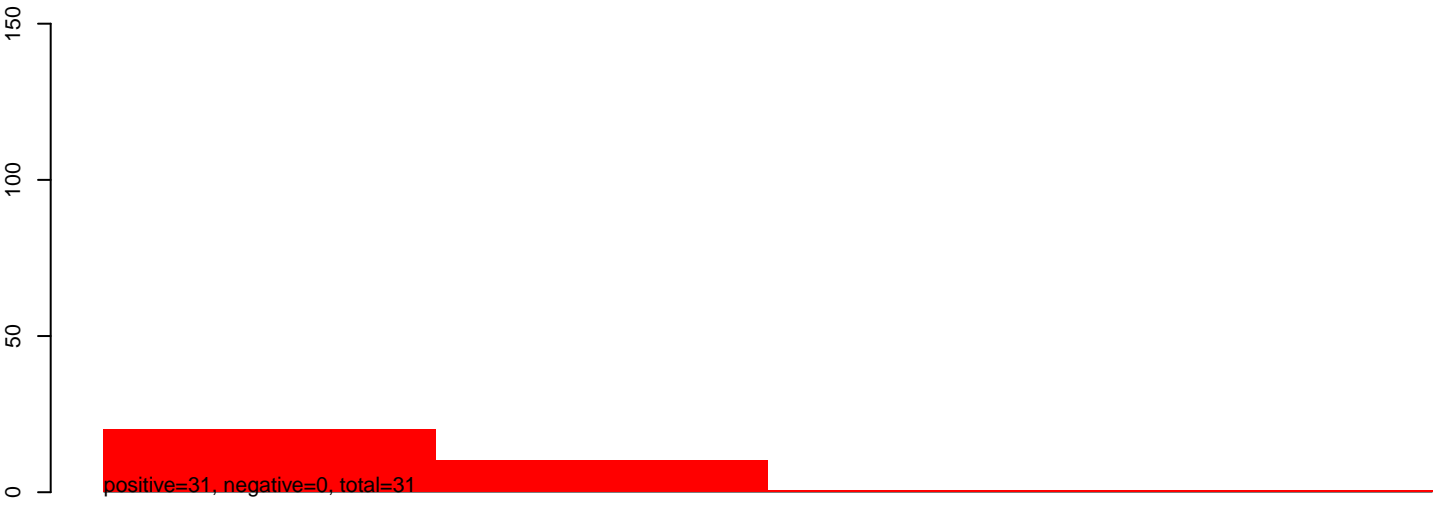


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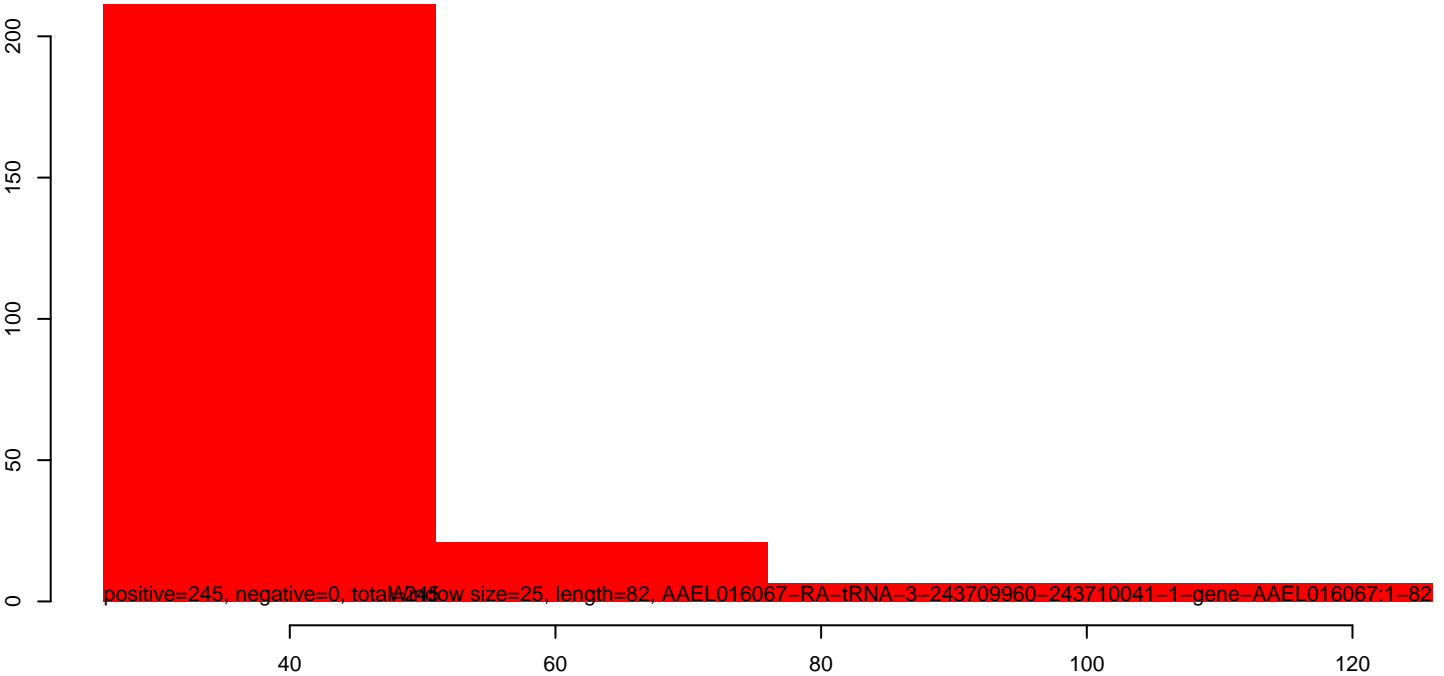
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AeAeg_Fcarc_GH_rep3.24_35.rep

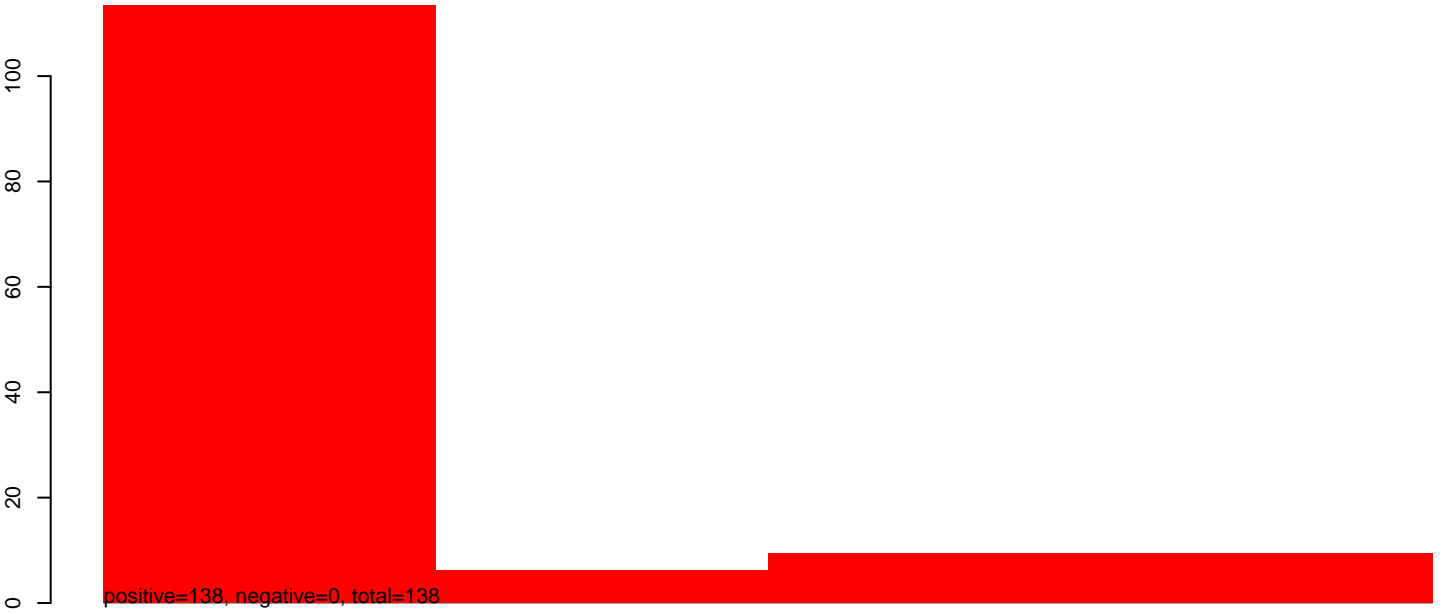


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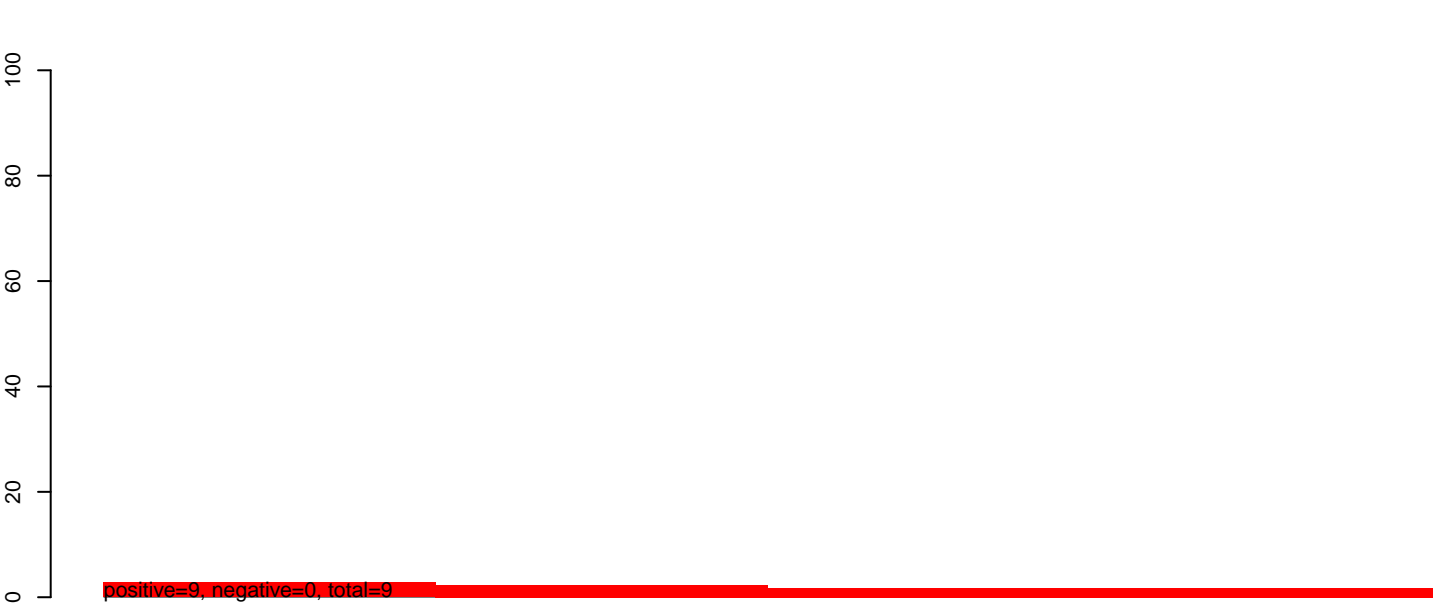


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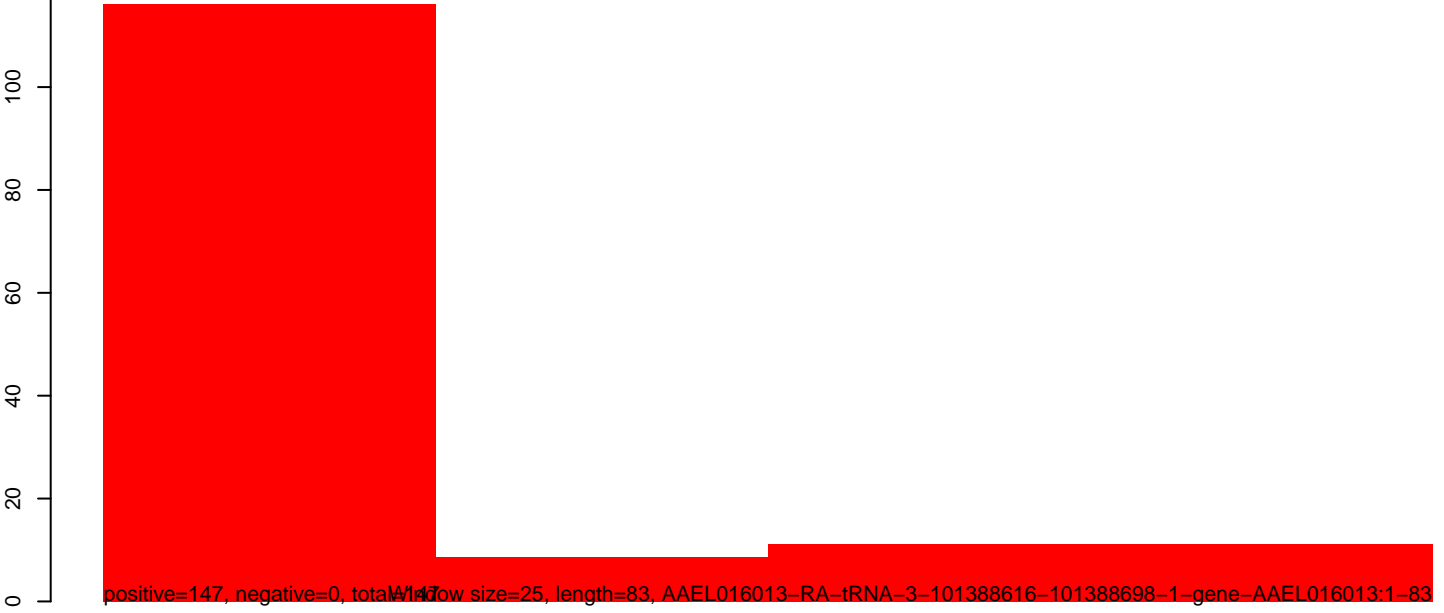
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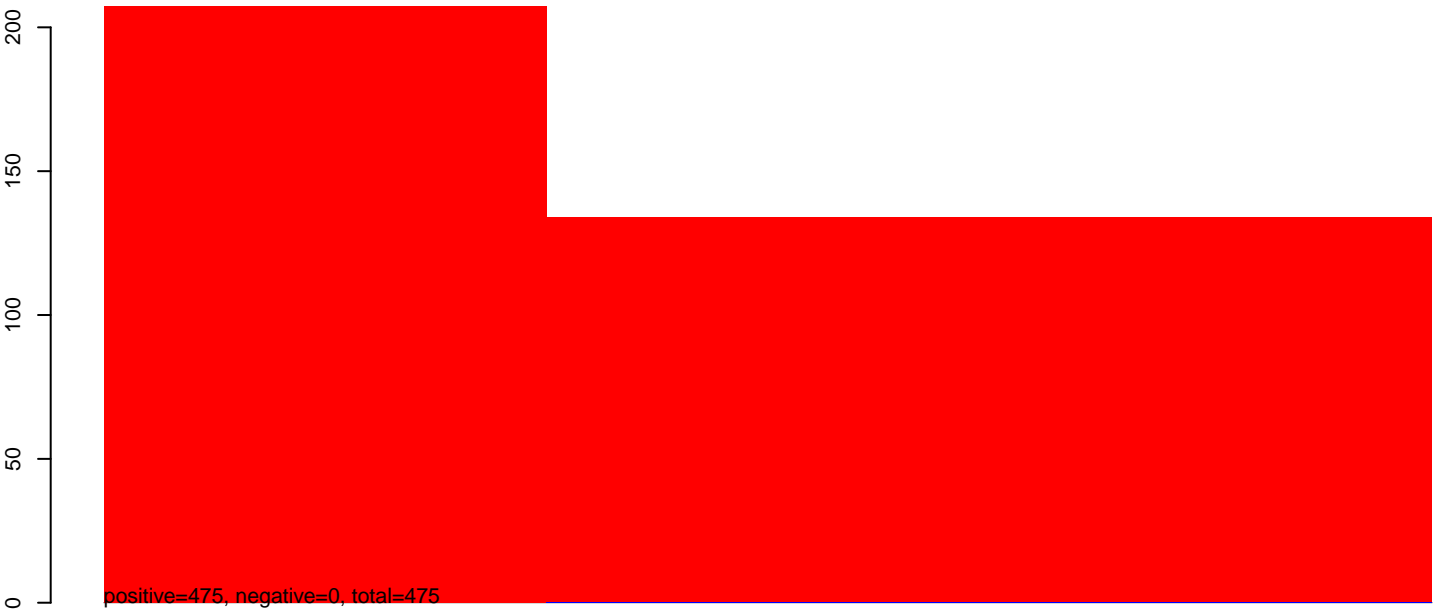


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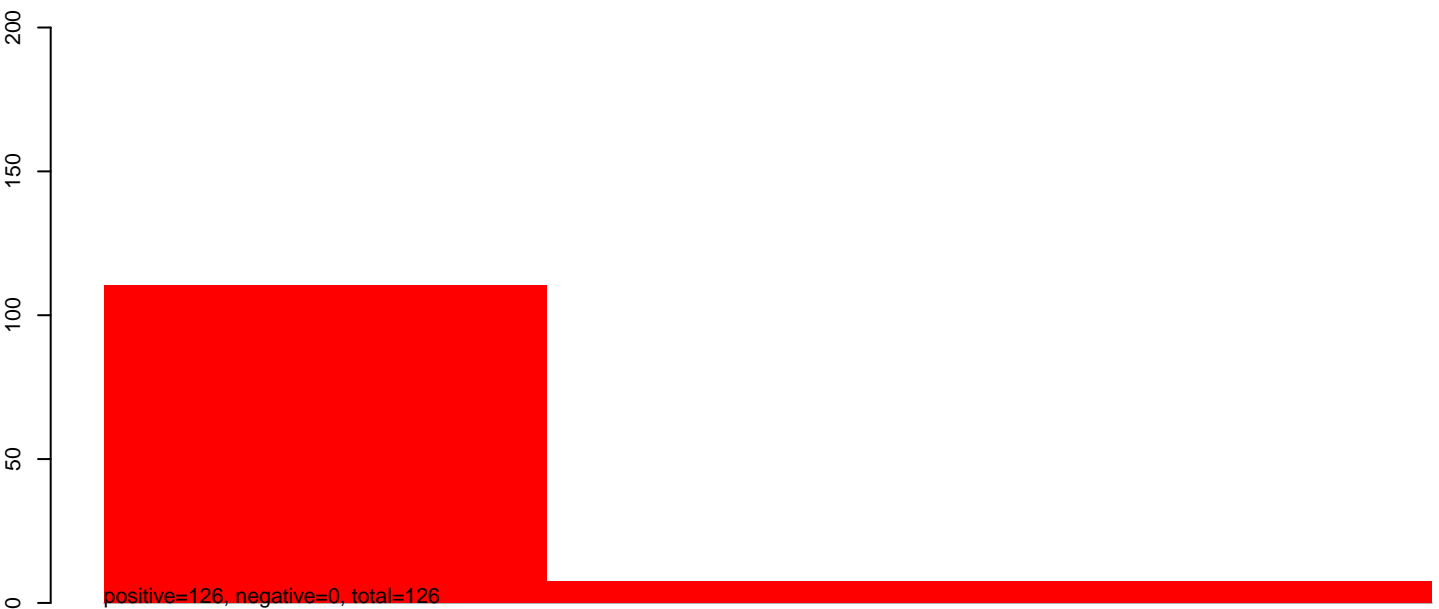


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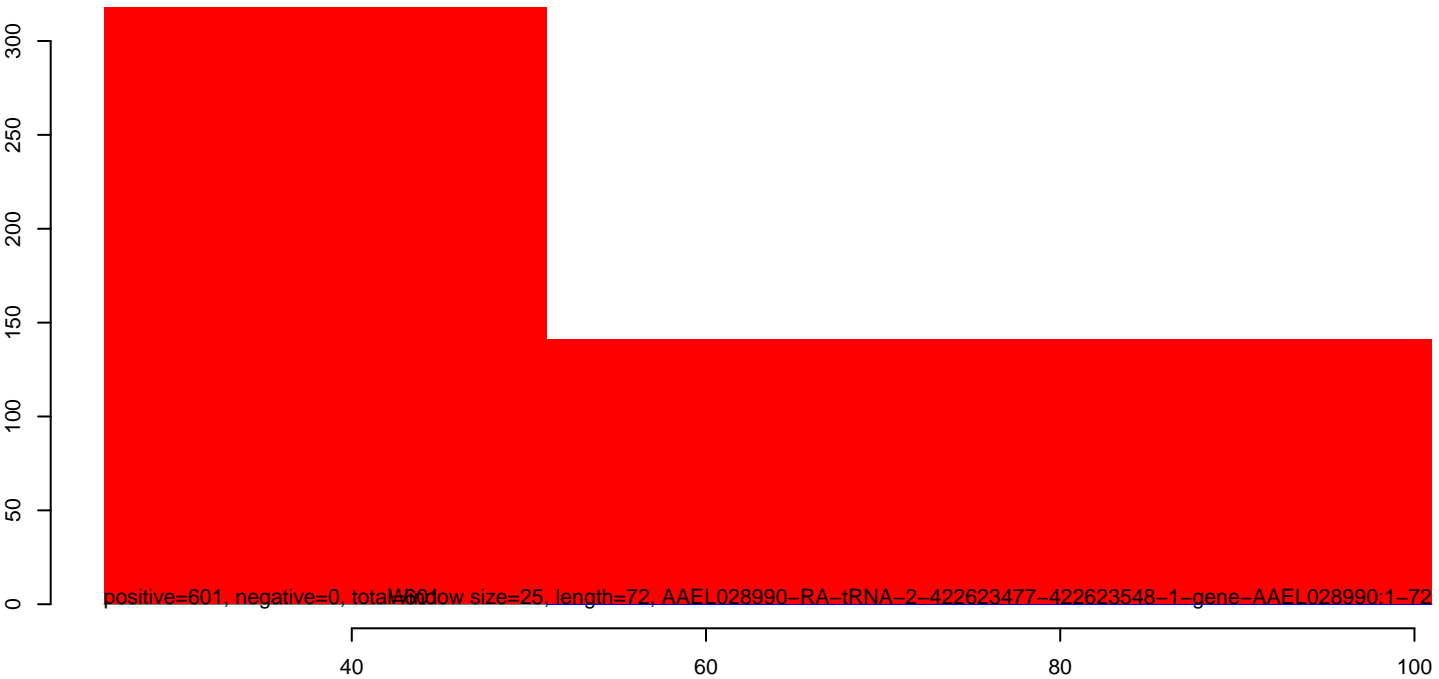
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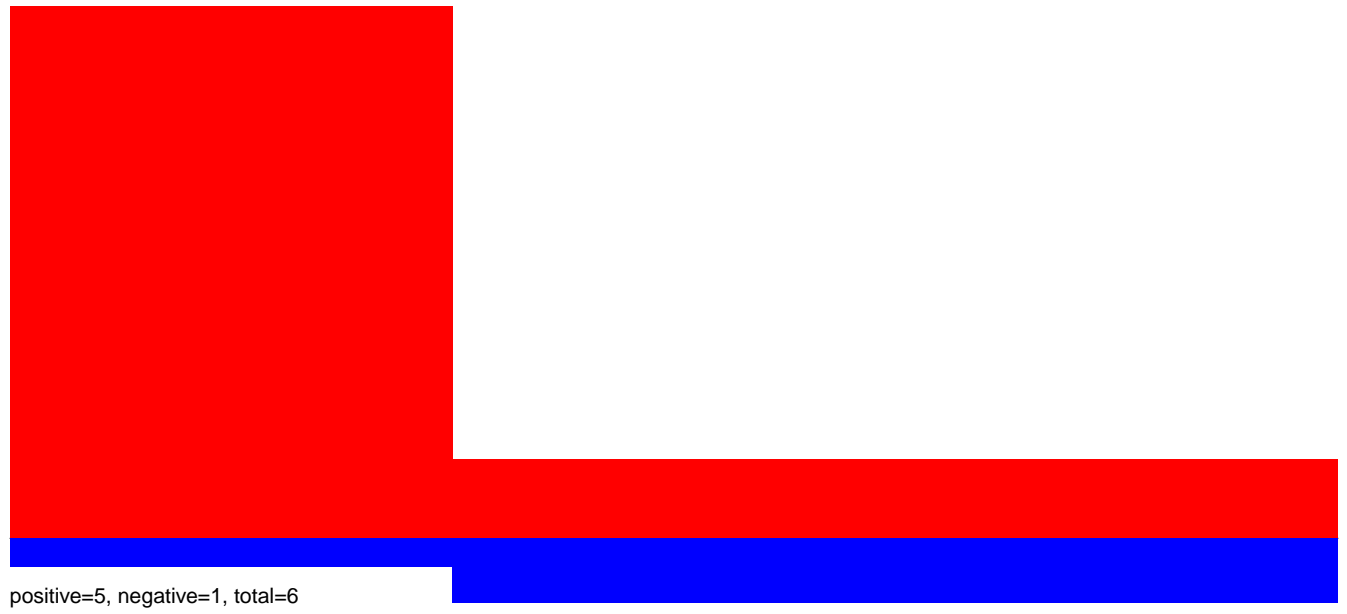
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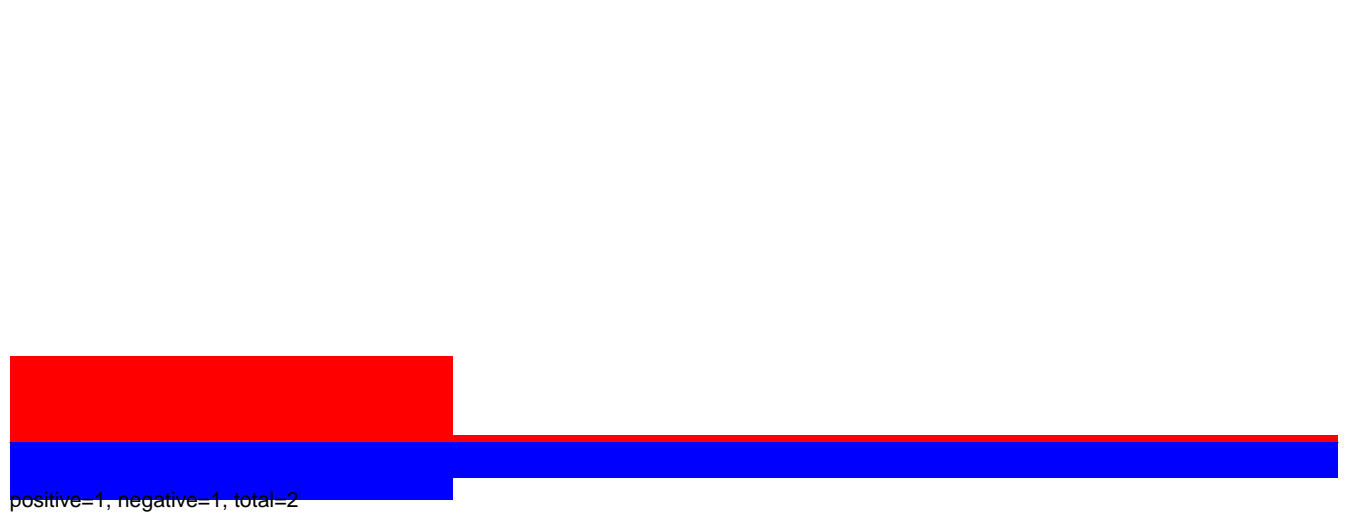
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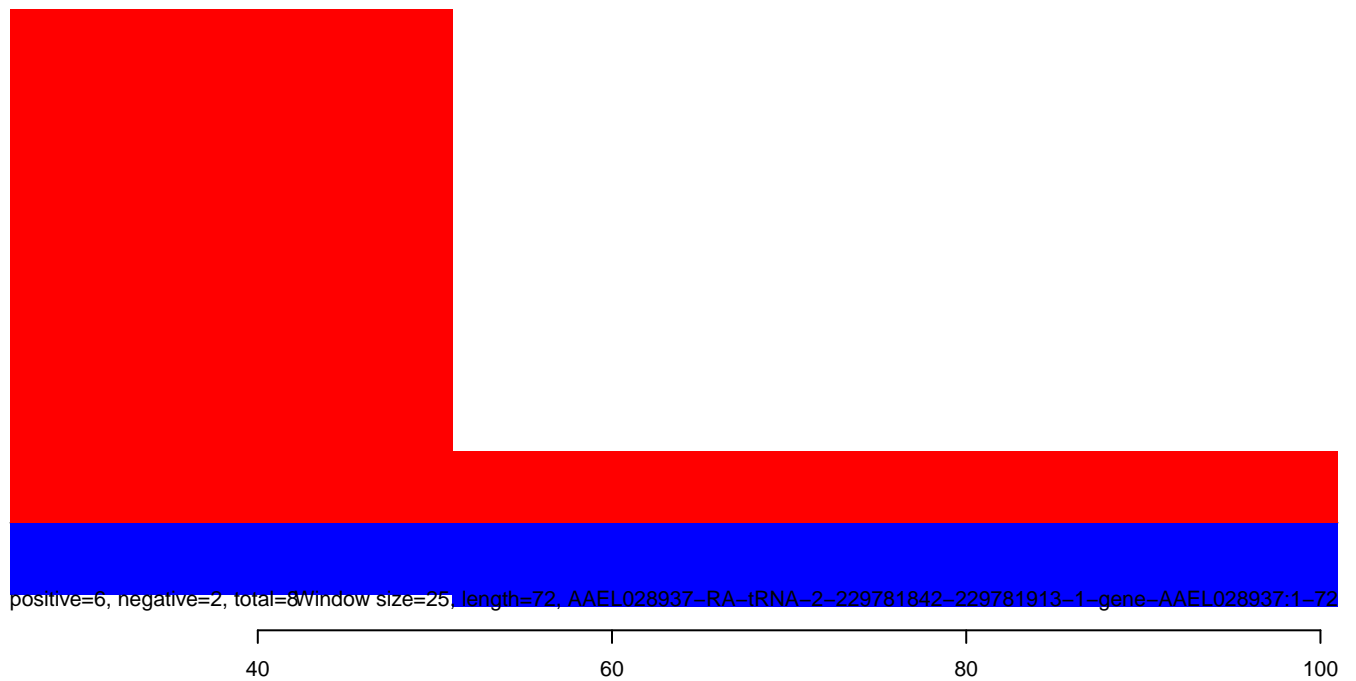
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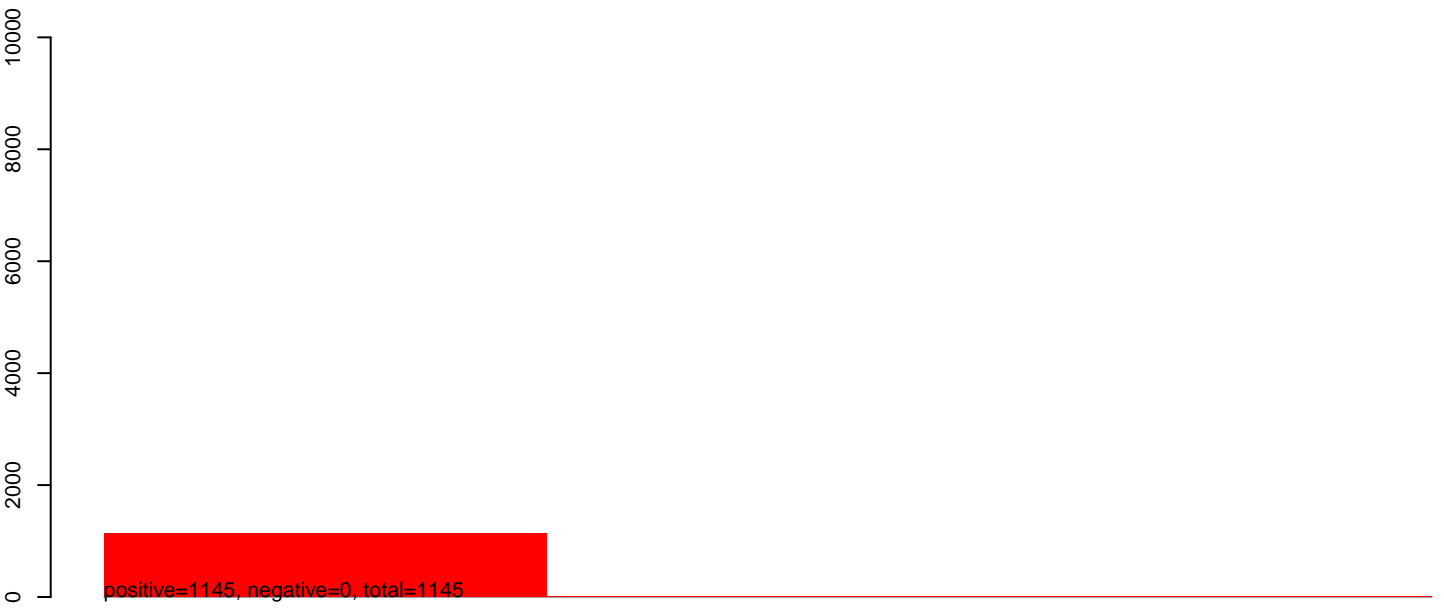
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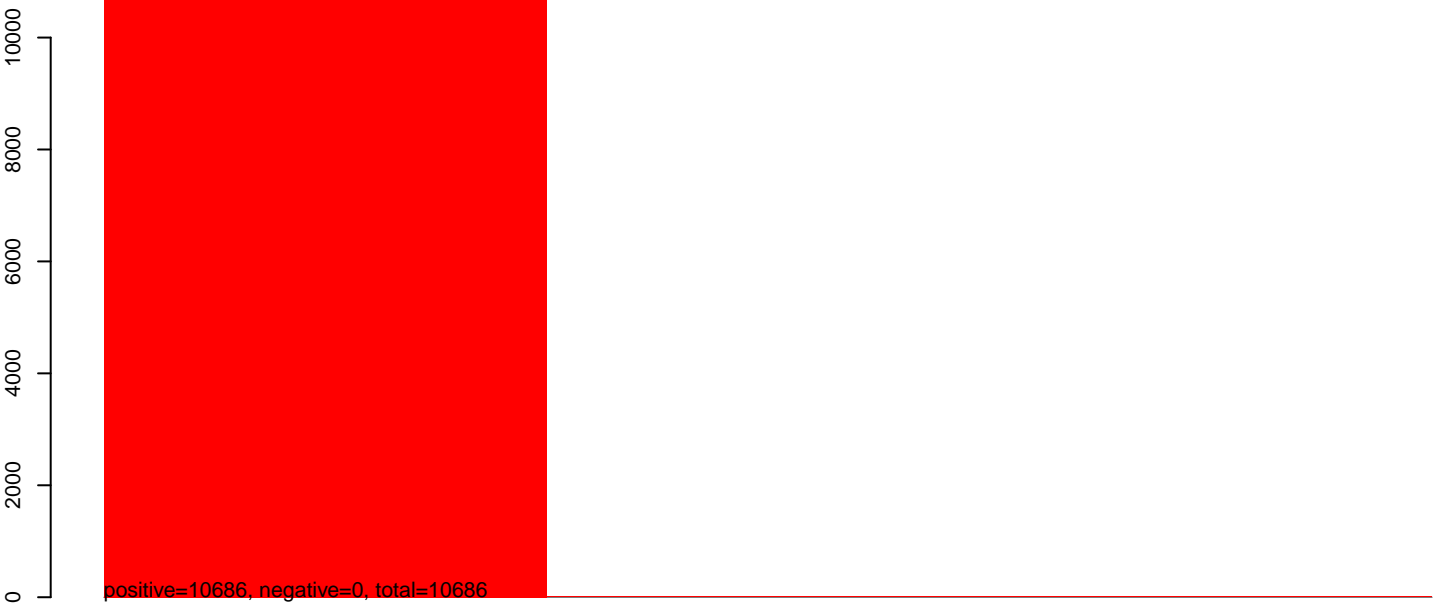
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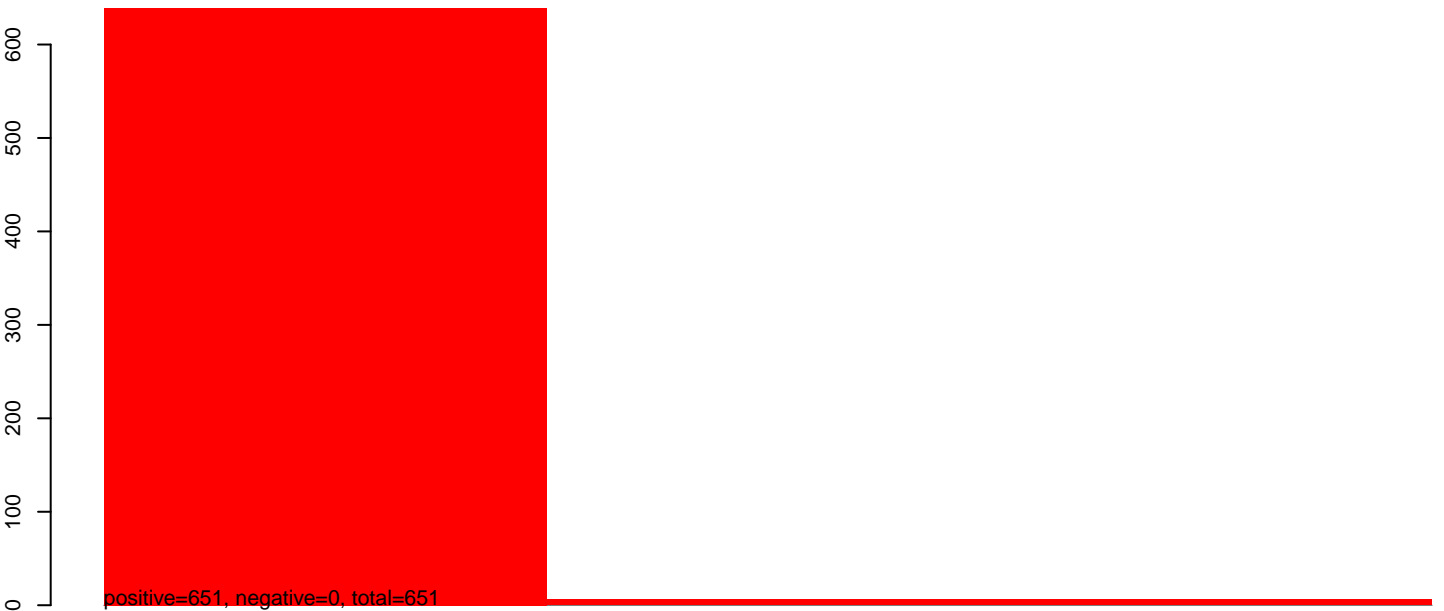
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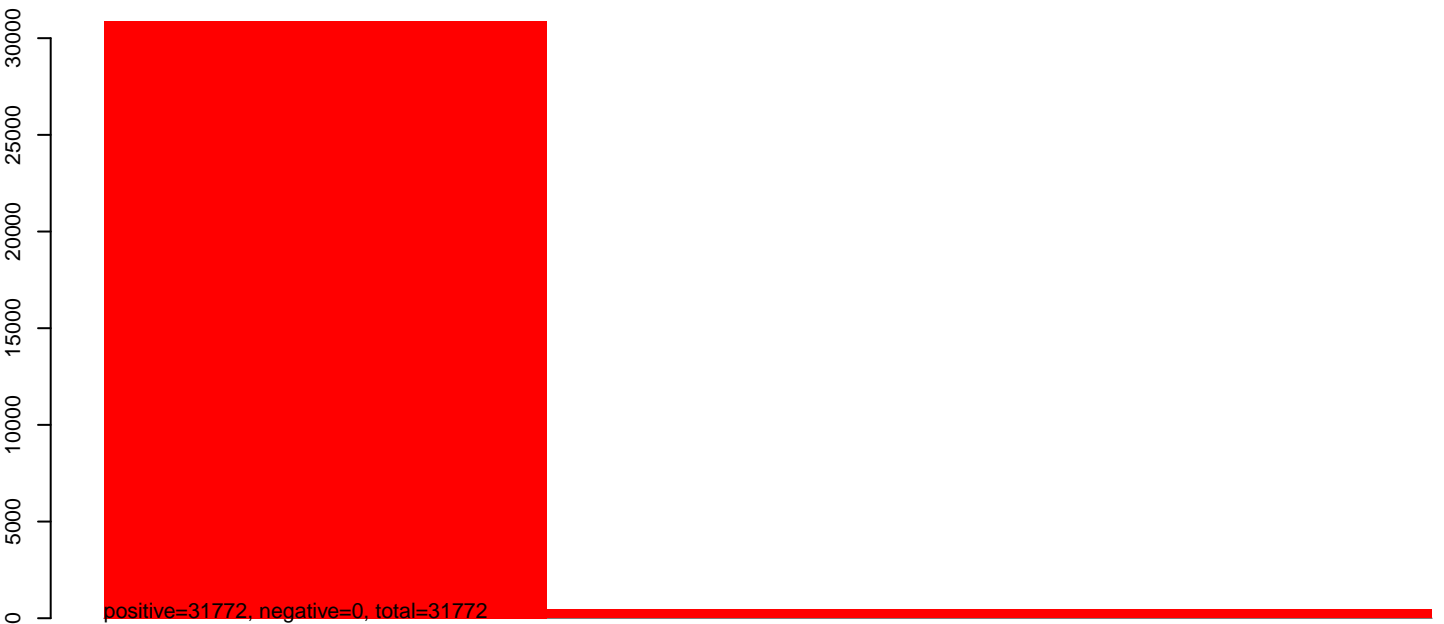
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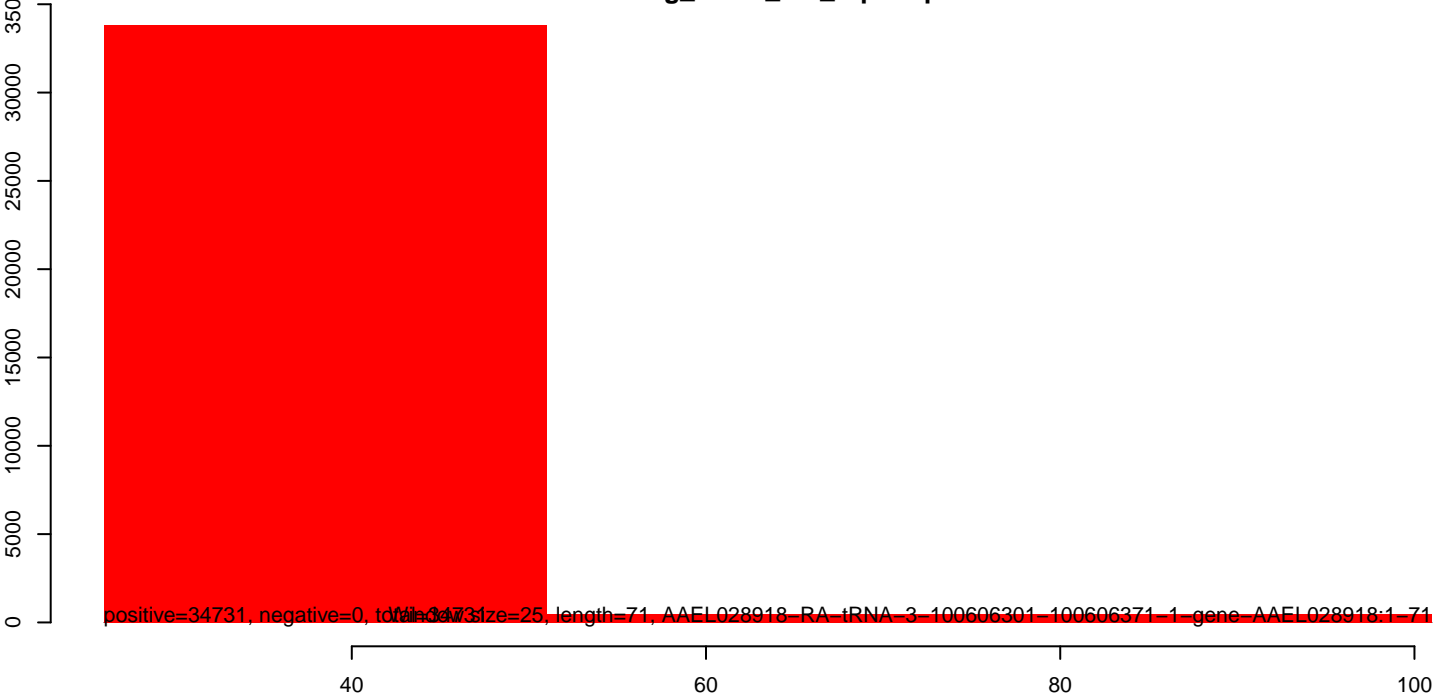
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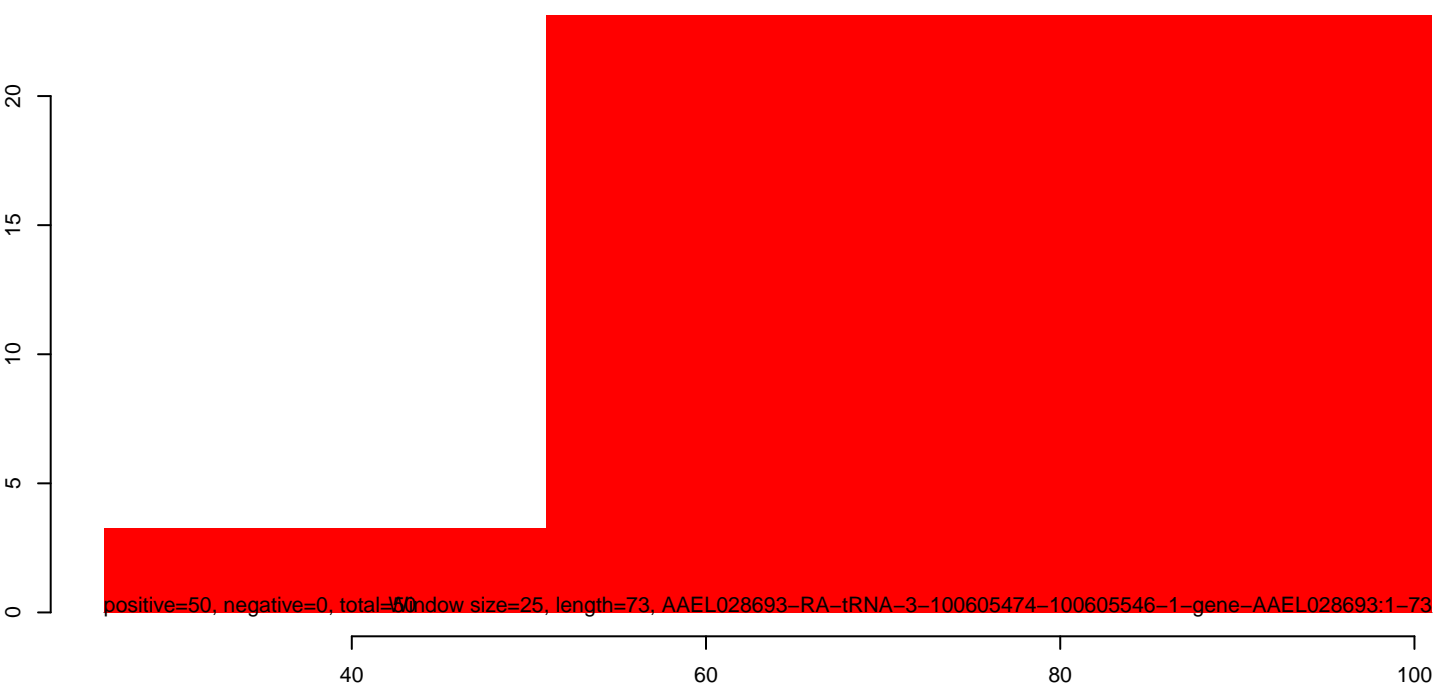
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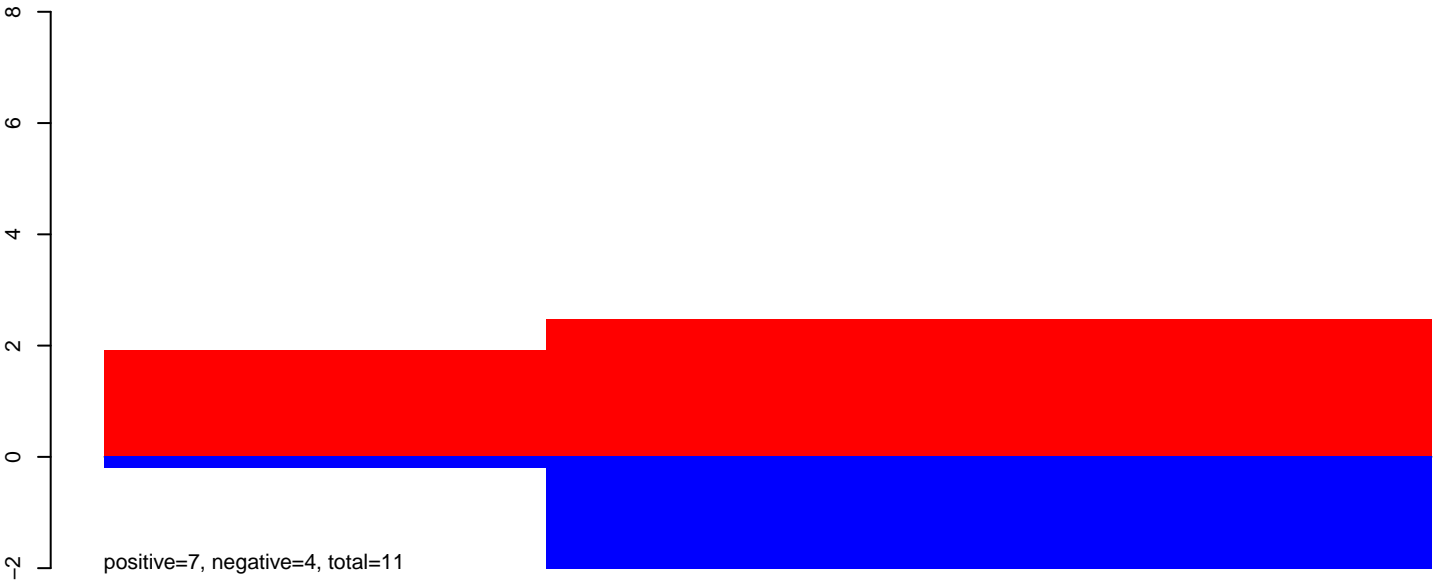
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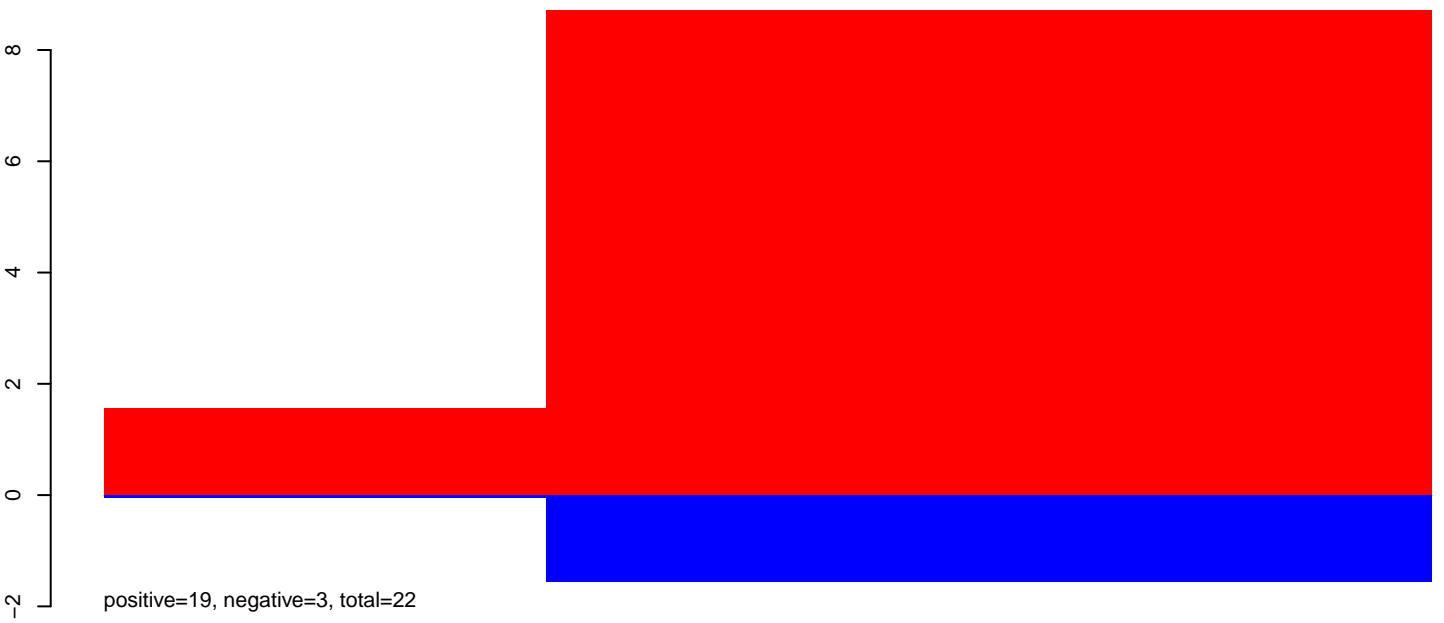
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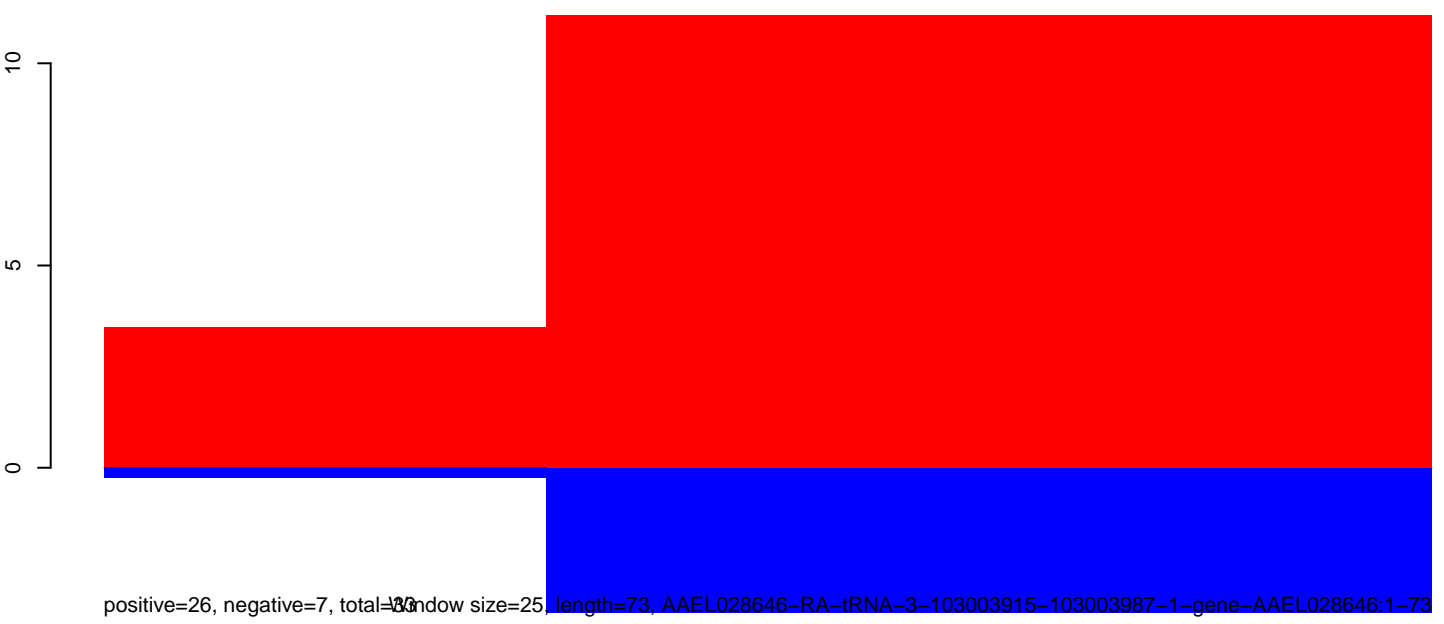
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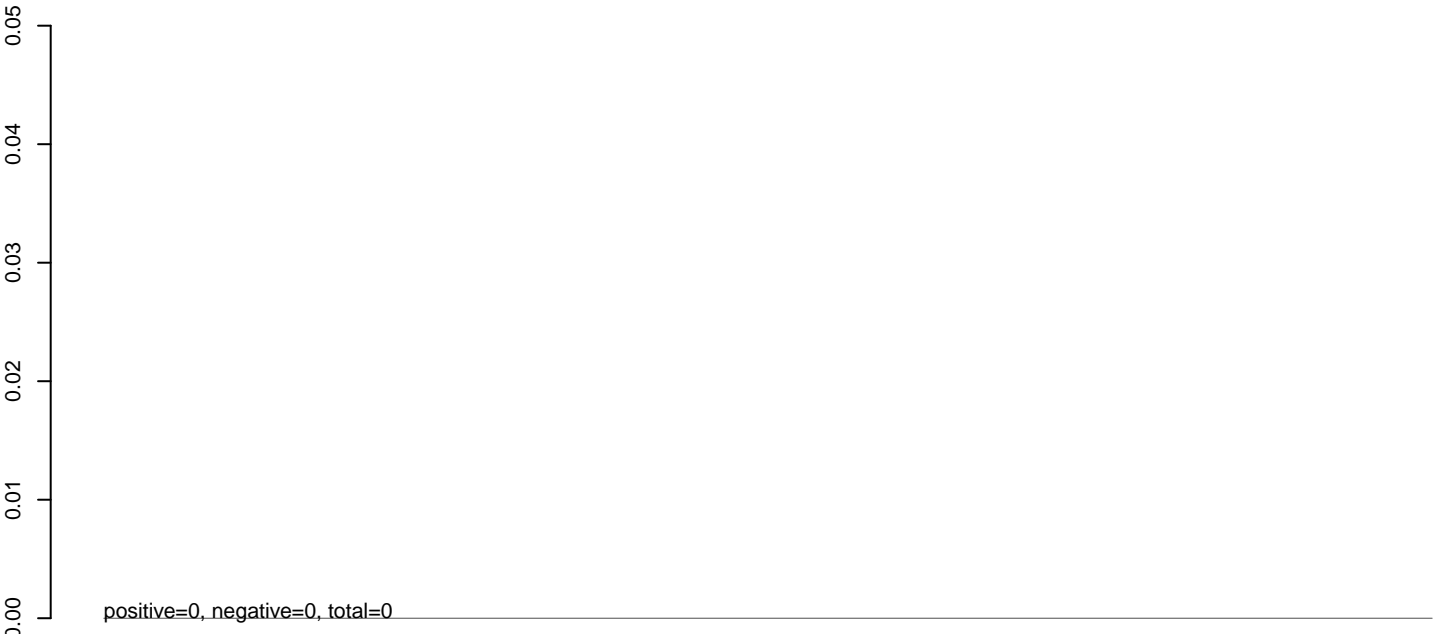
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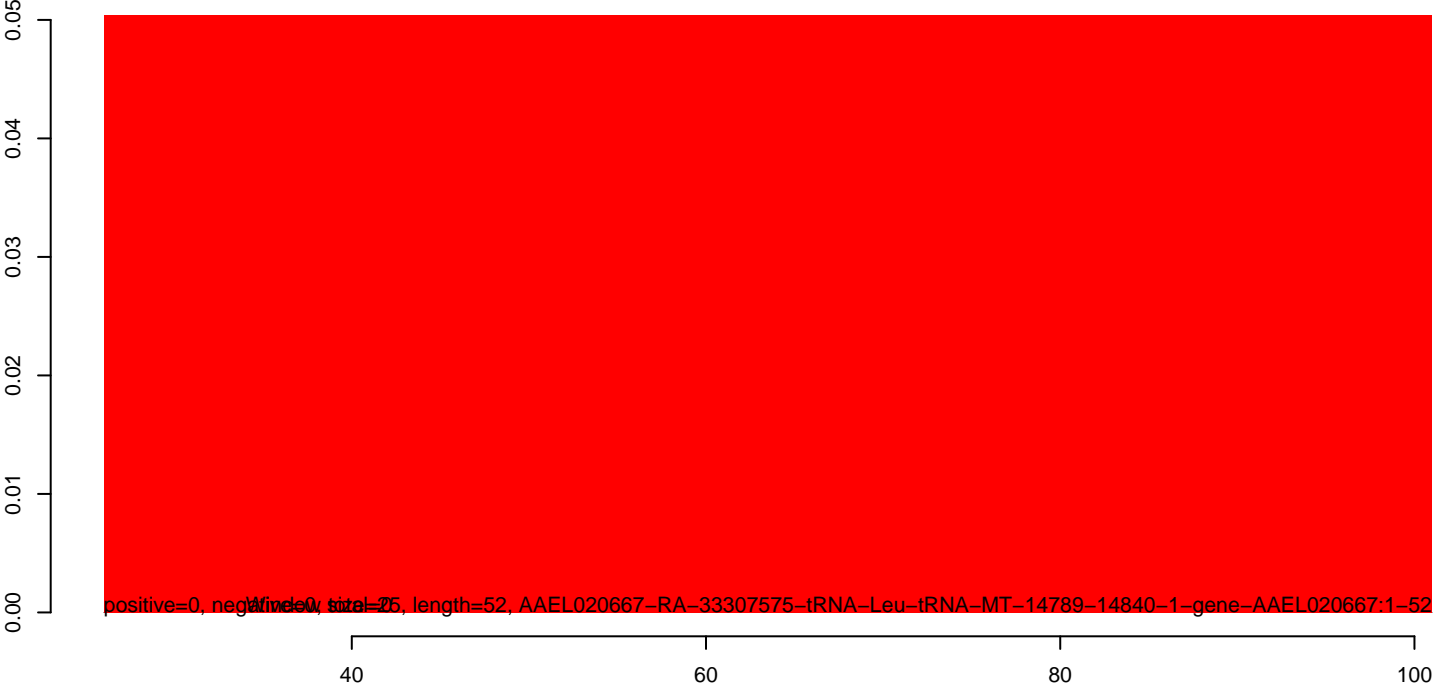
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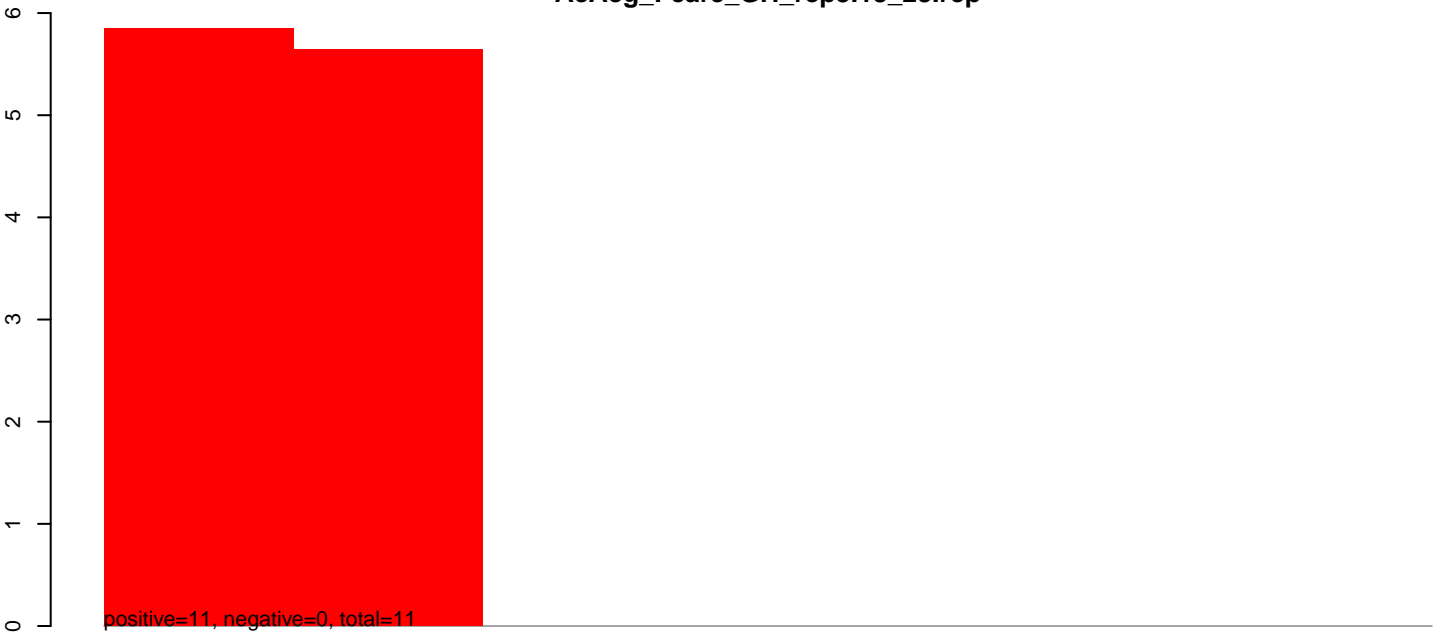
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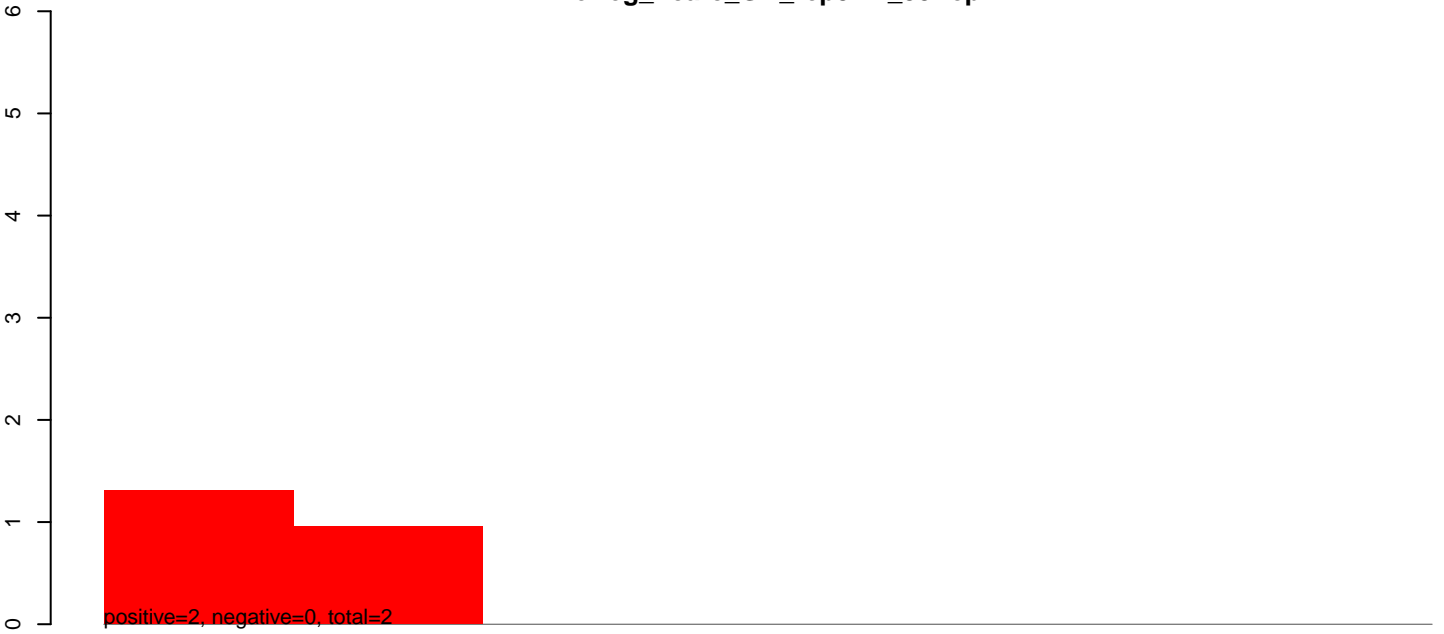
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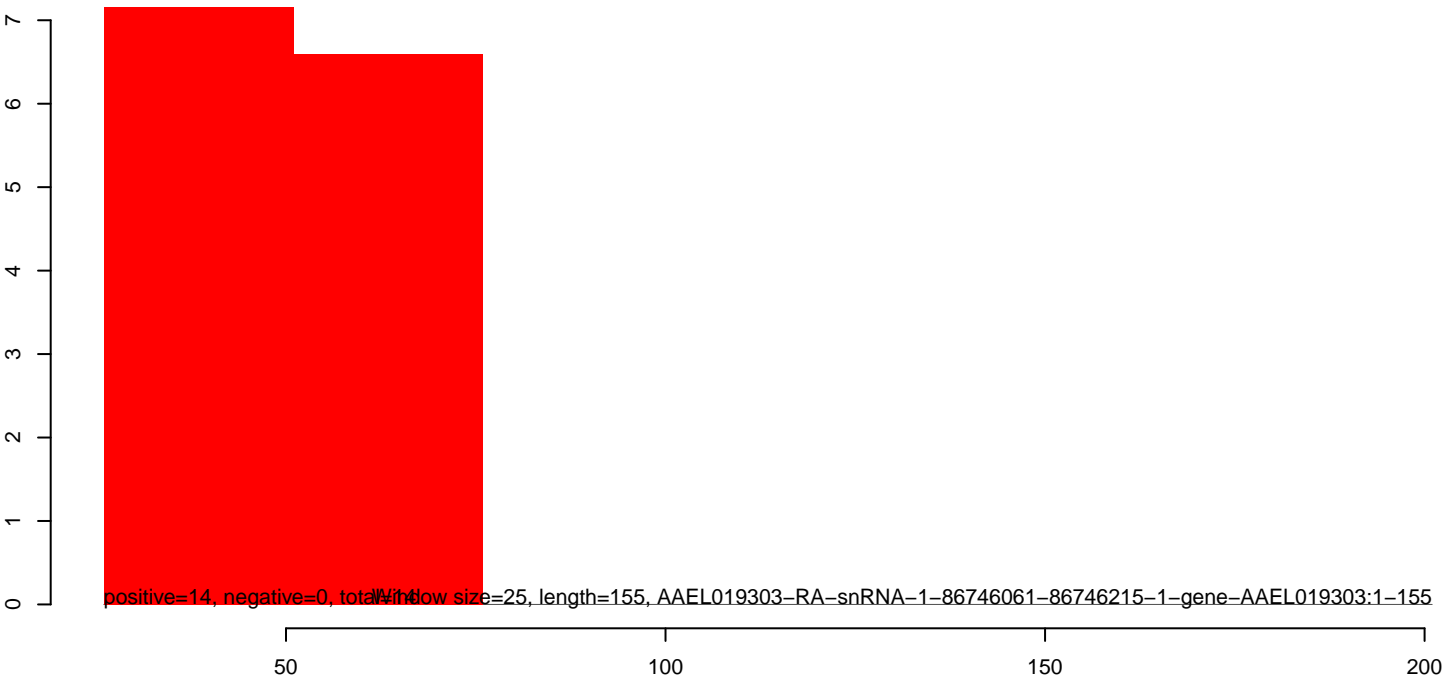
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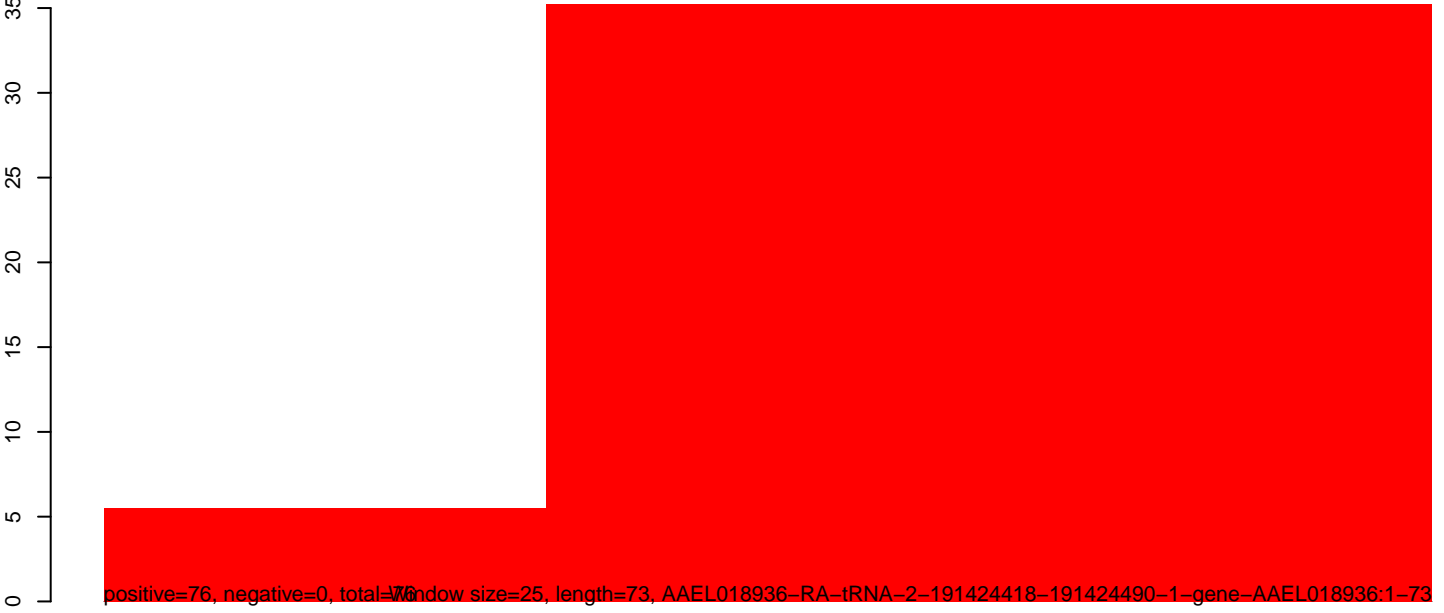
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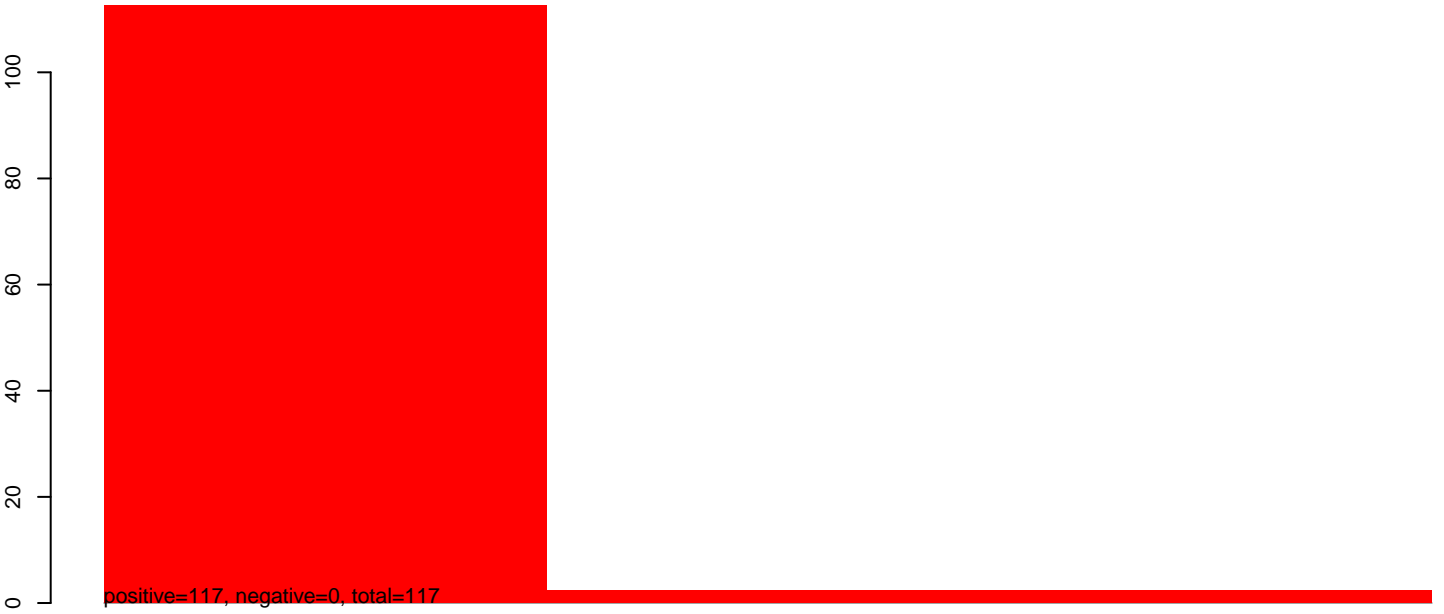


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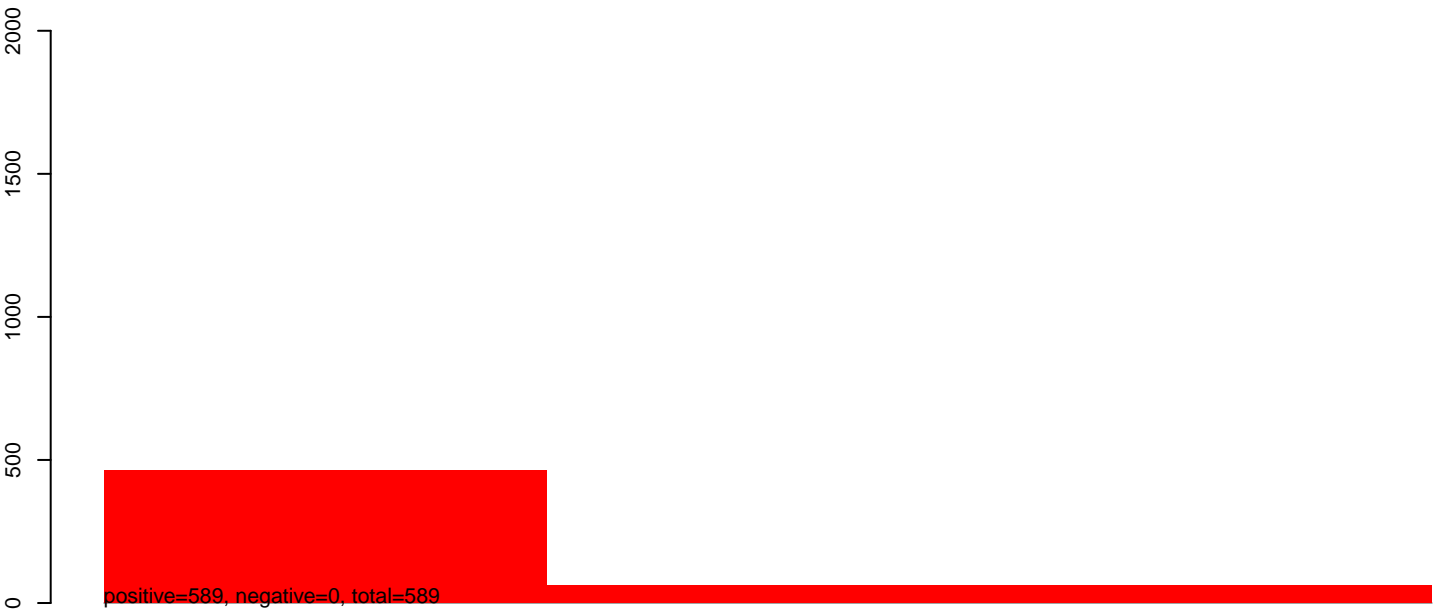


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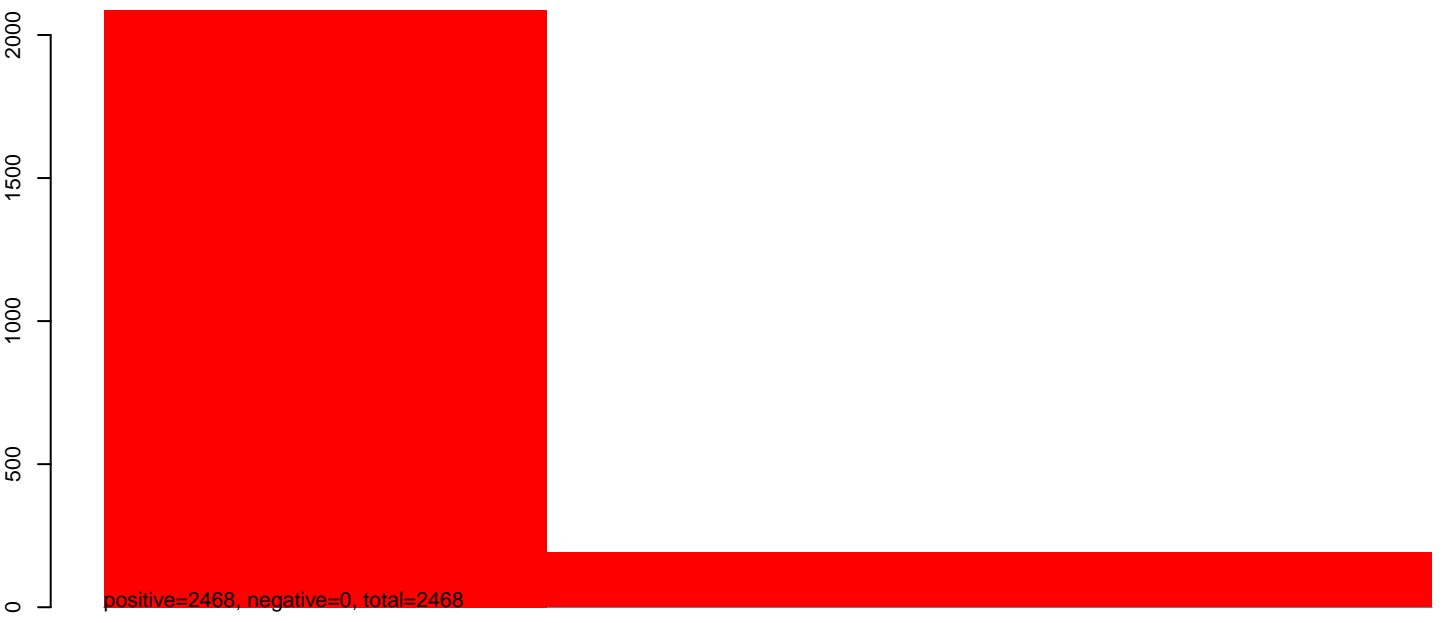


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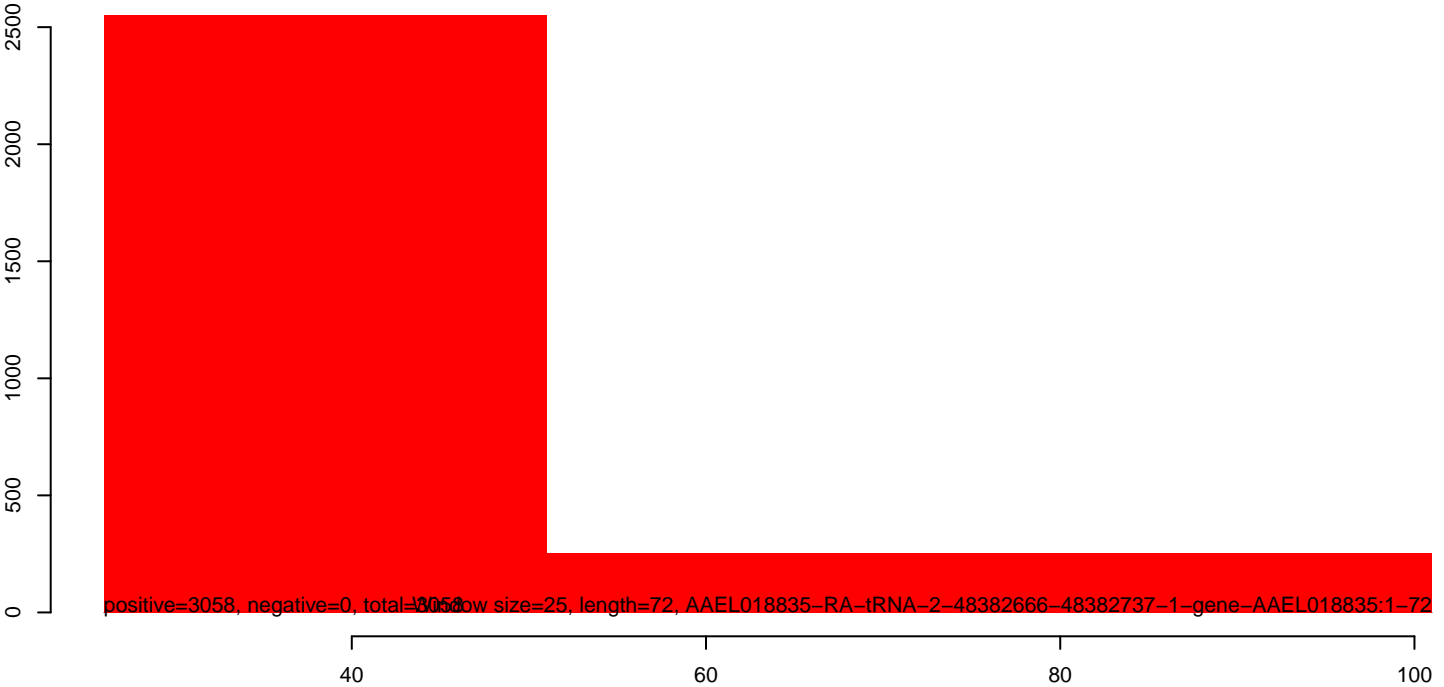
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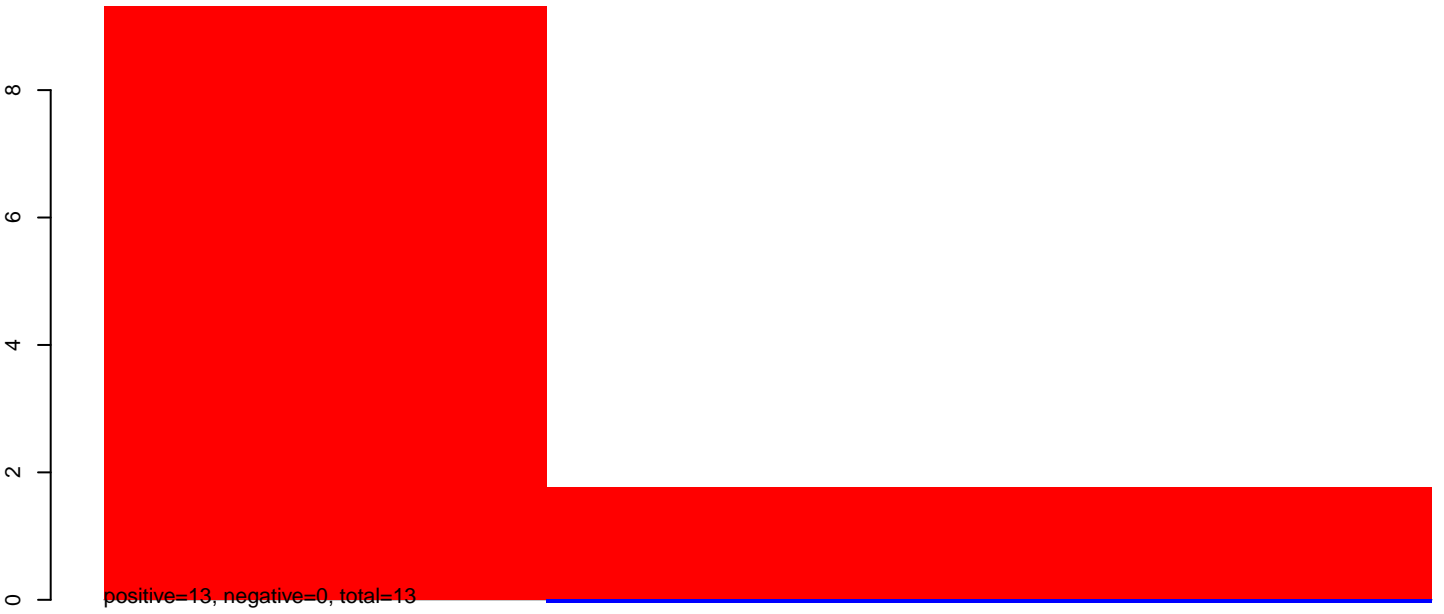
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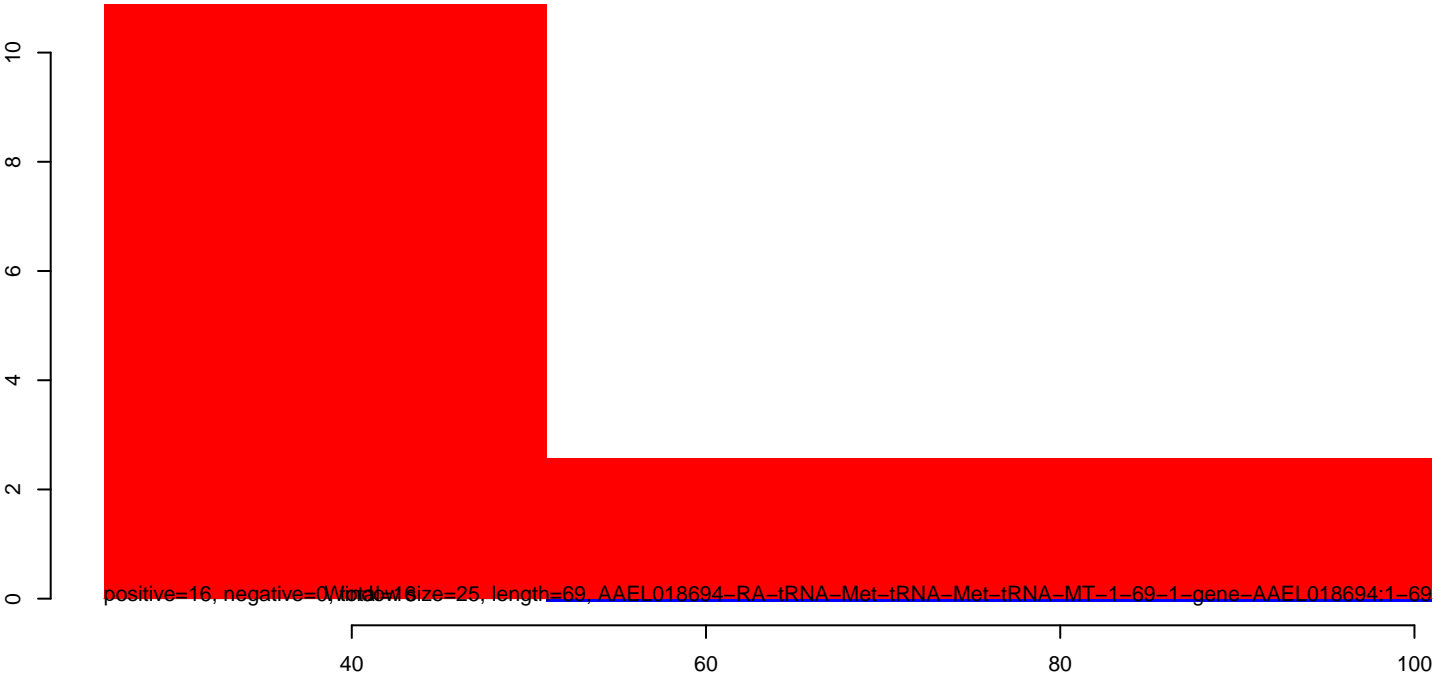
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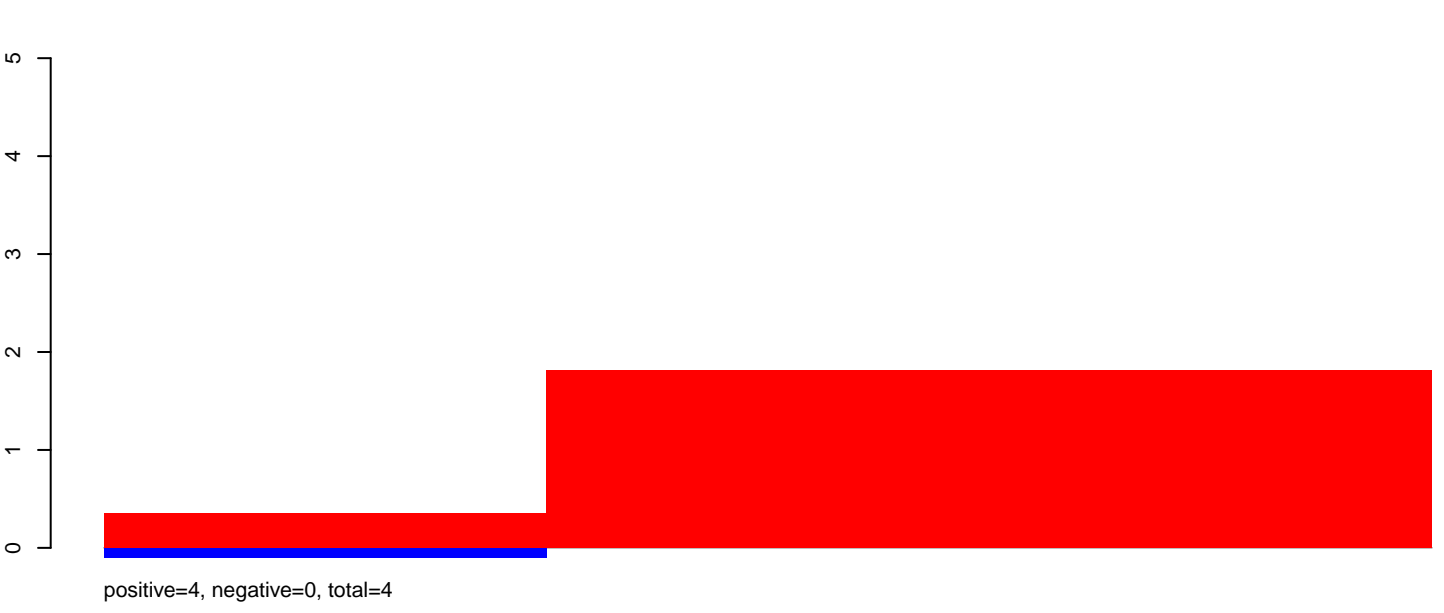
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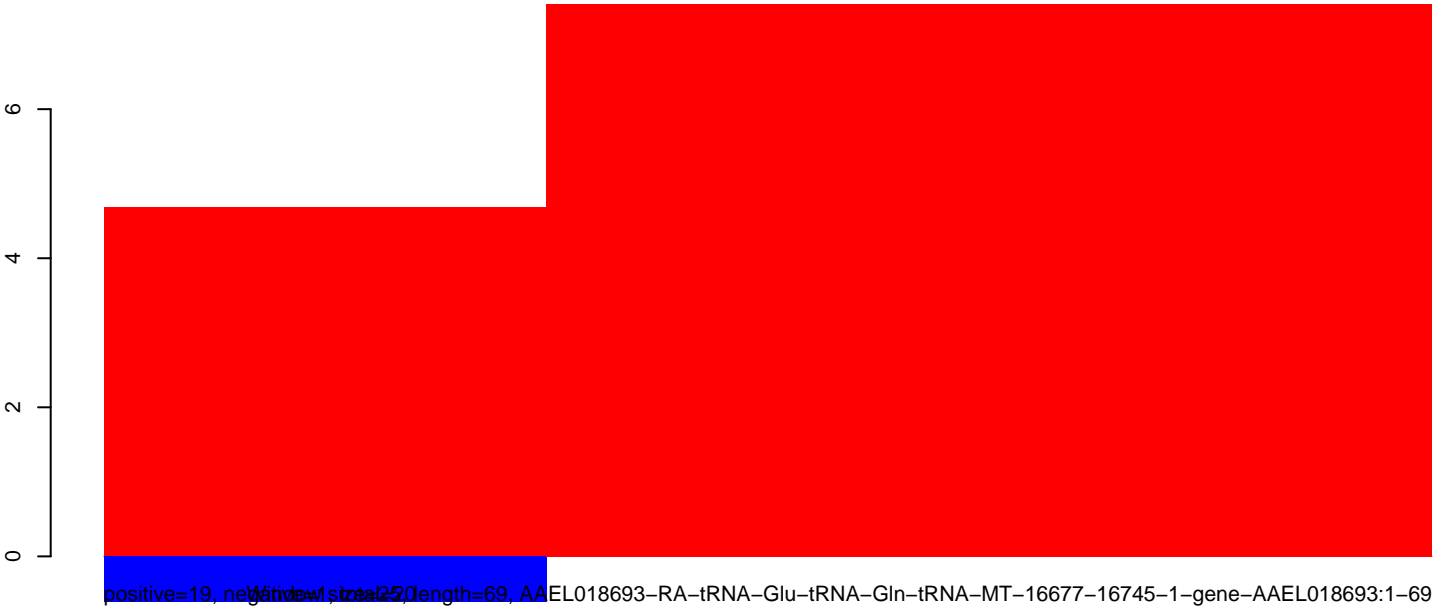
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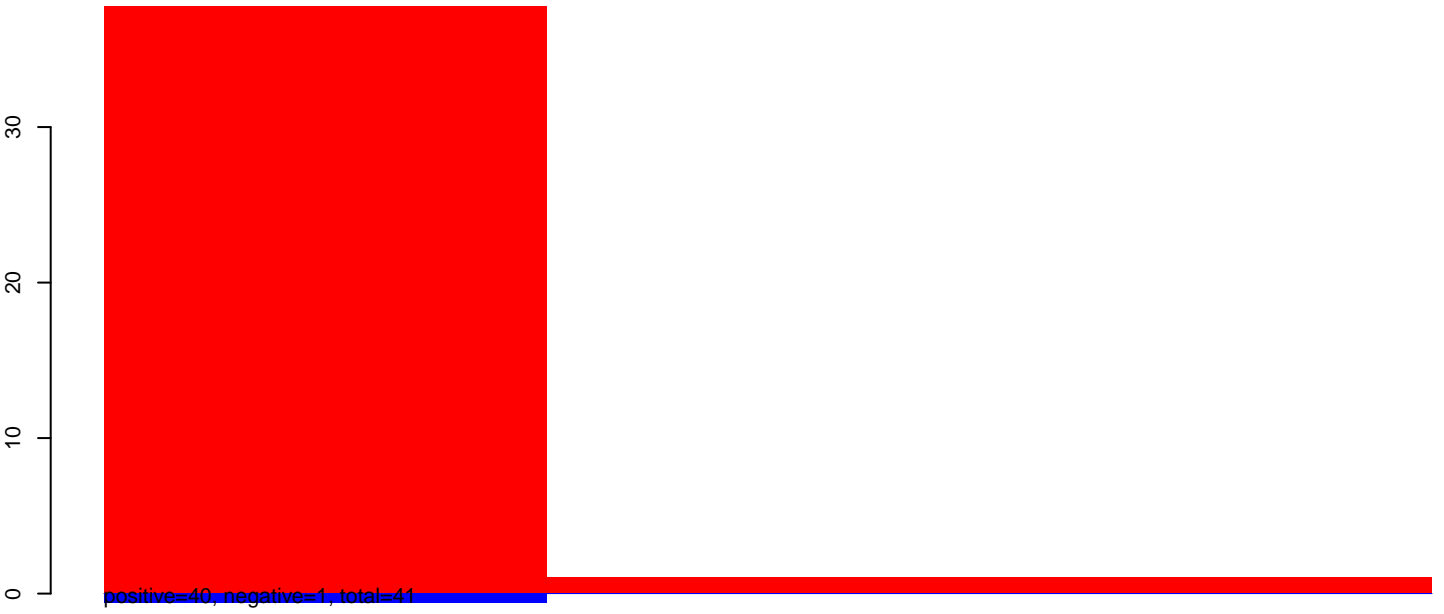
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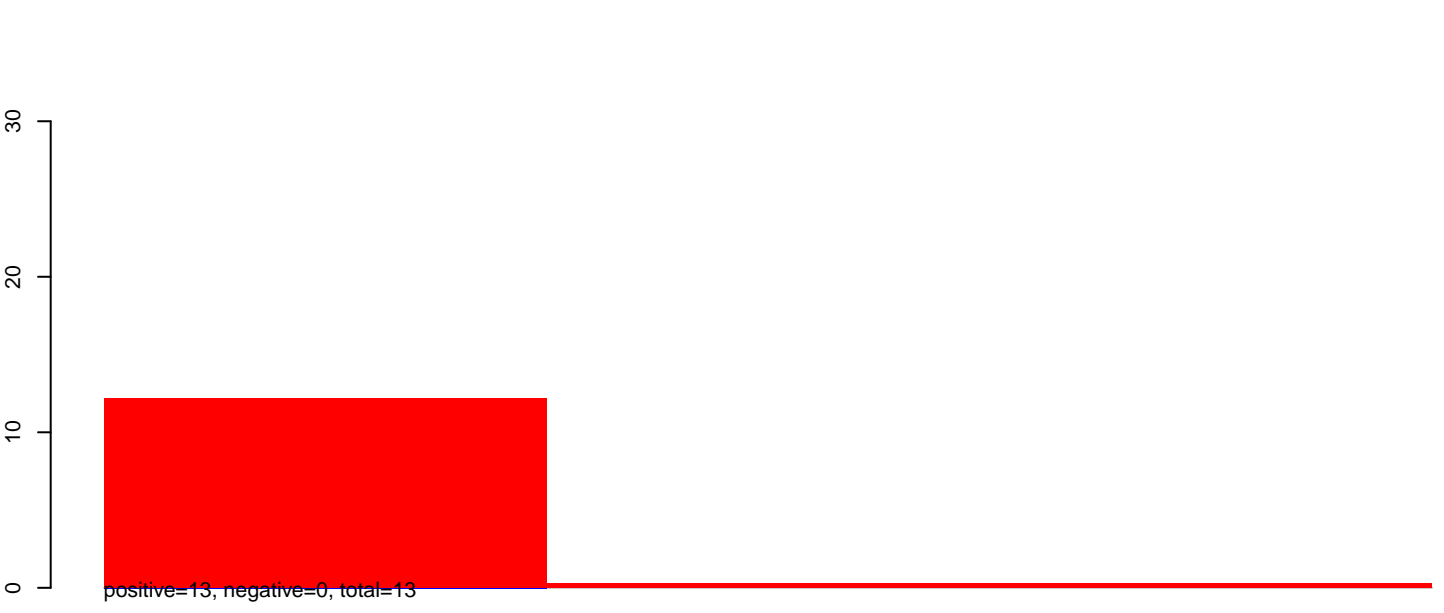
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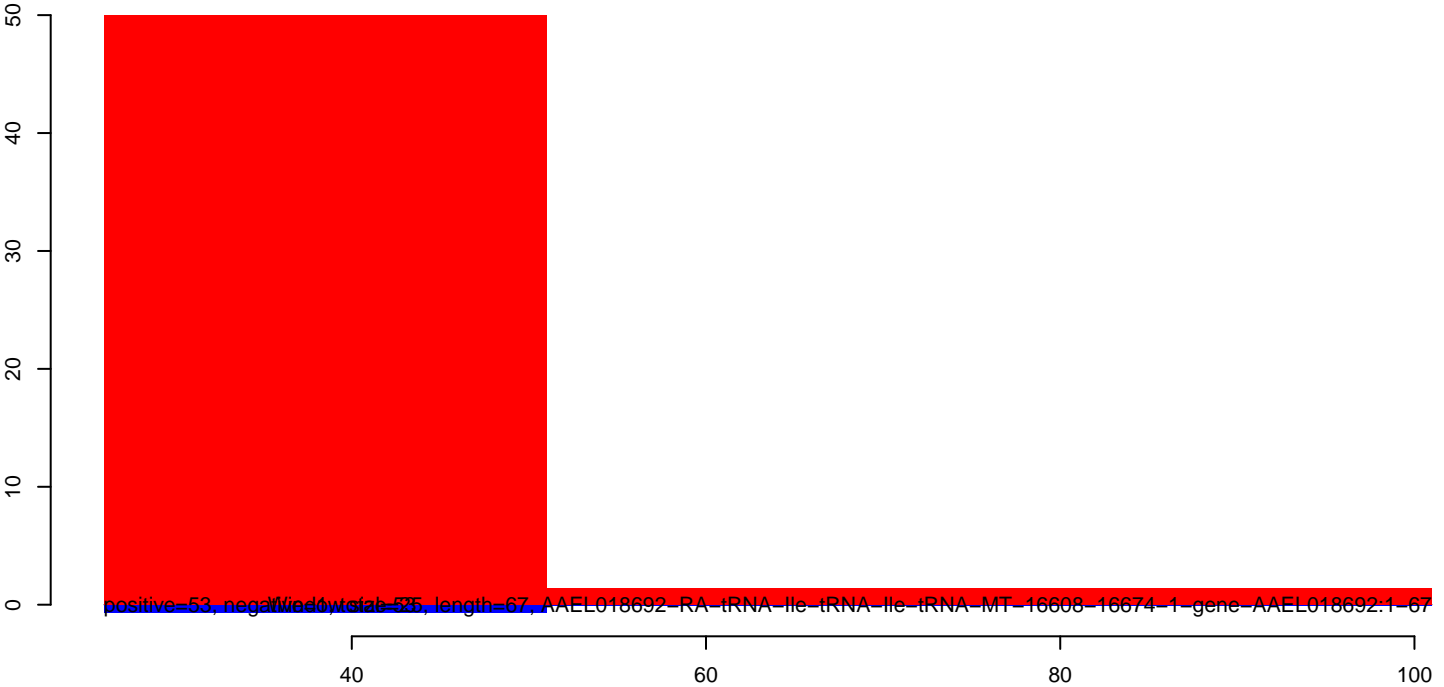
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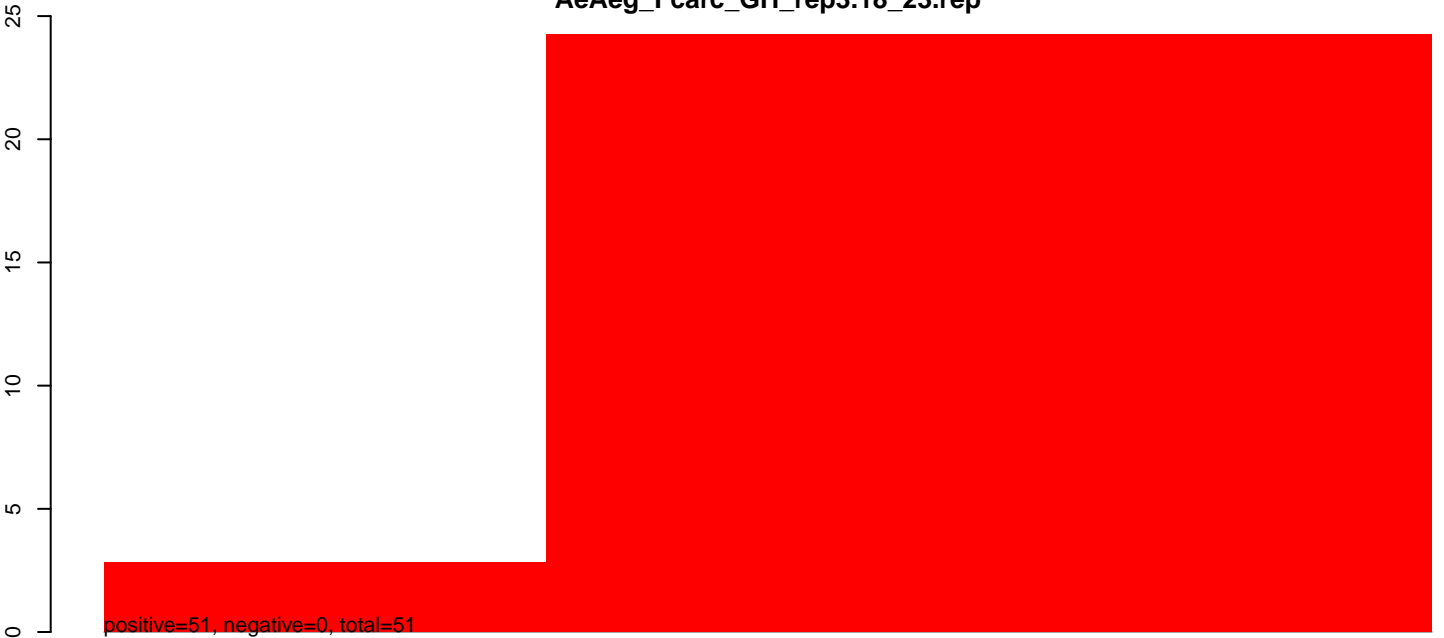
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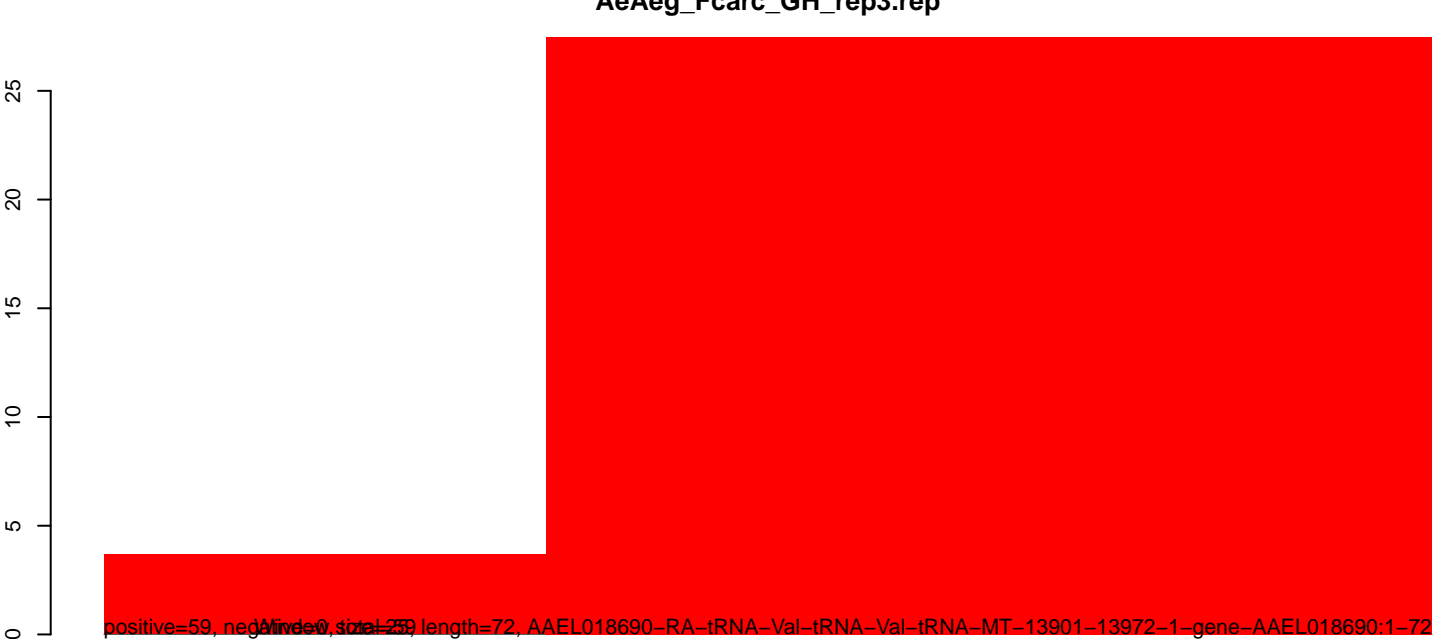
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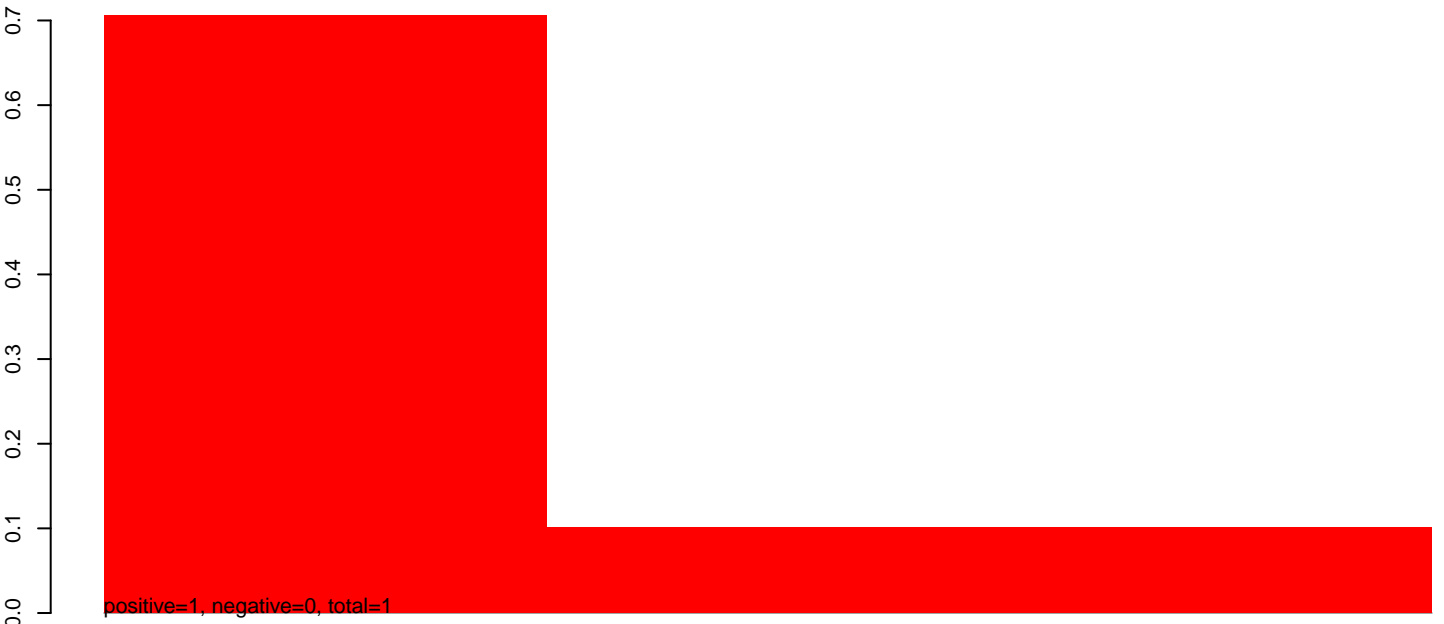


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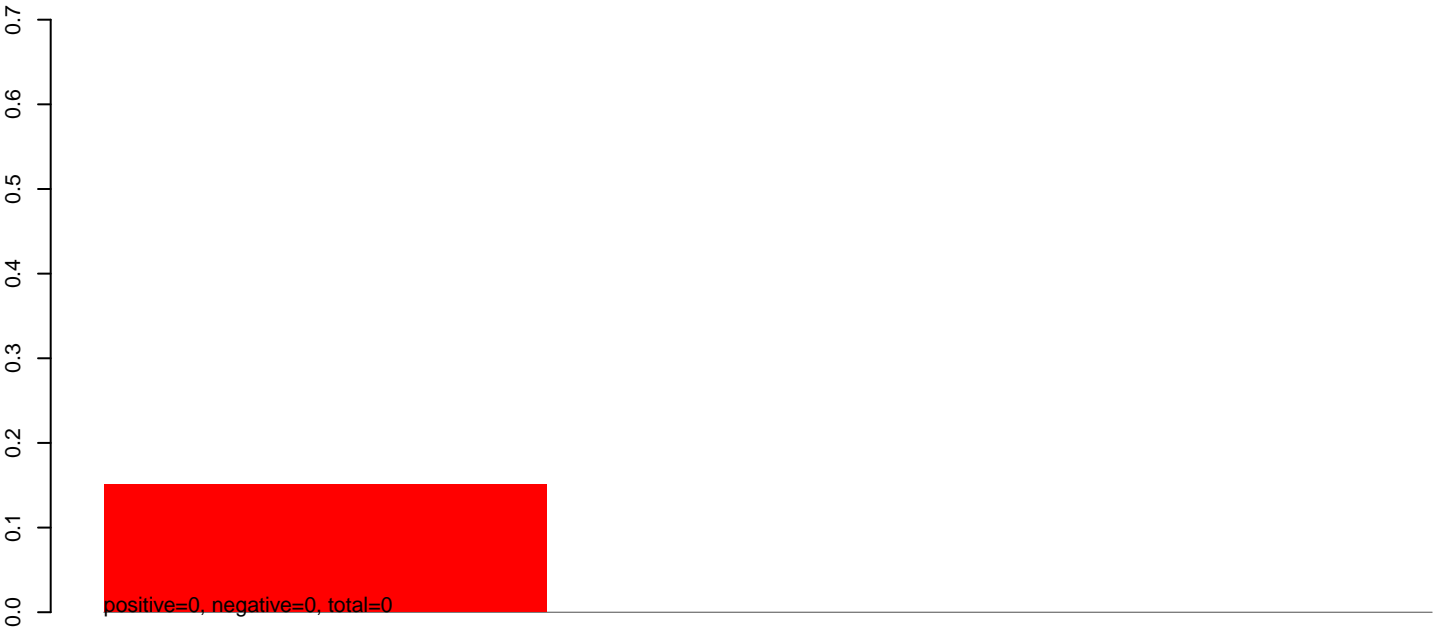


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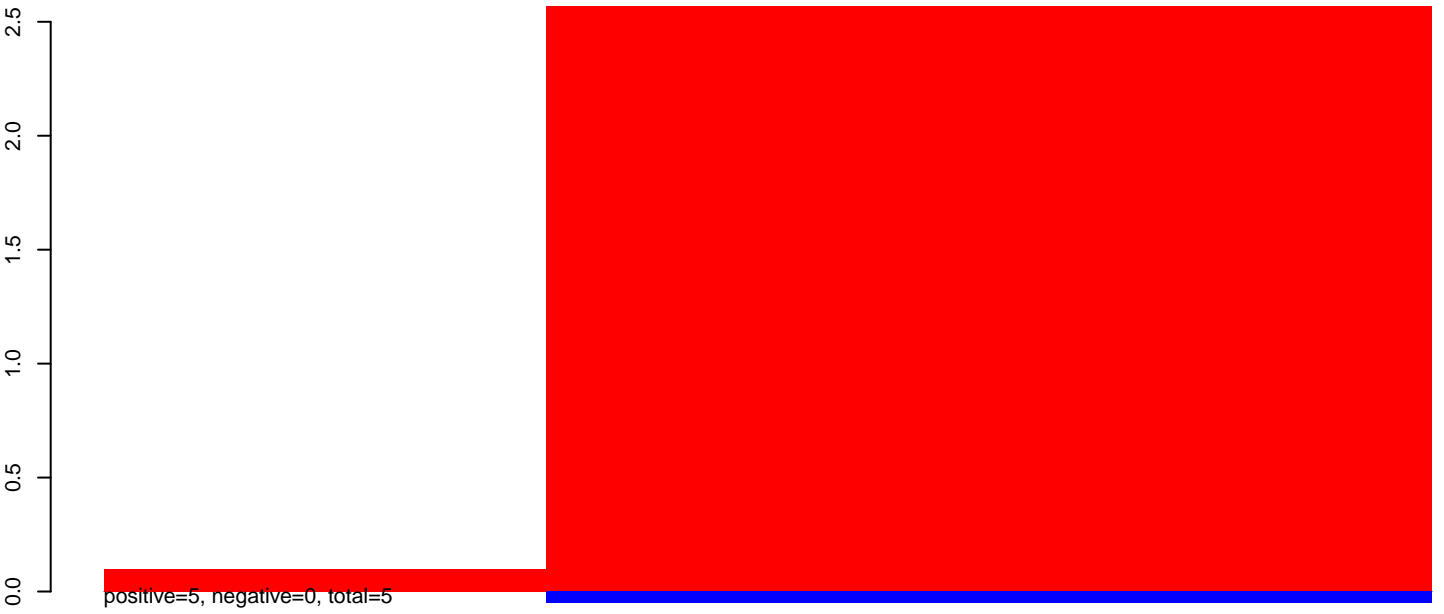
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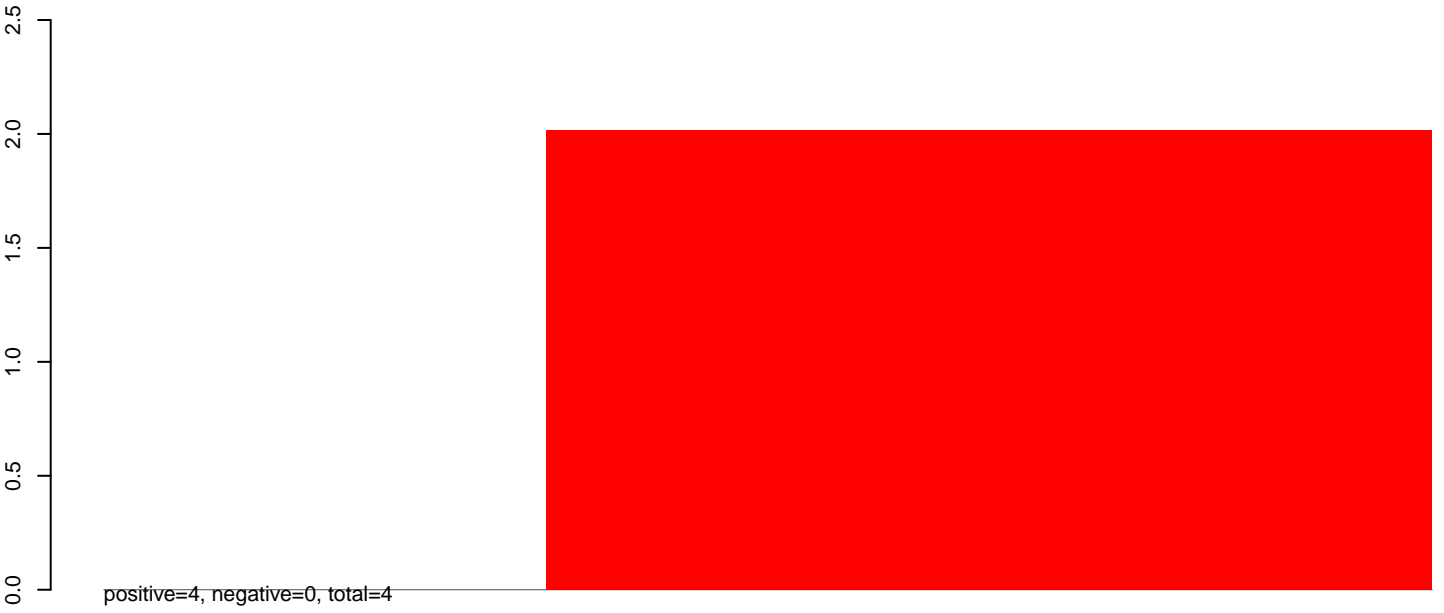
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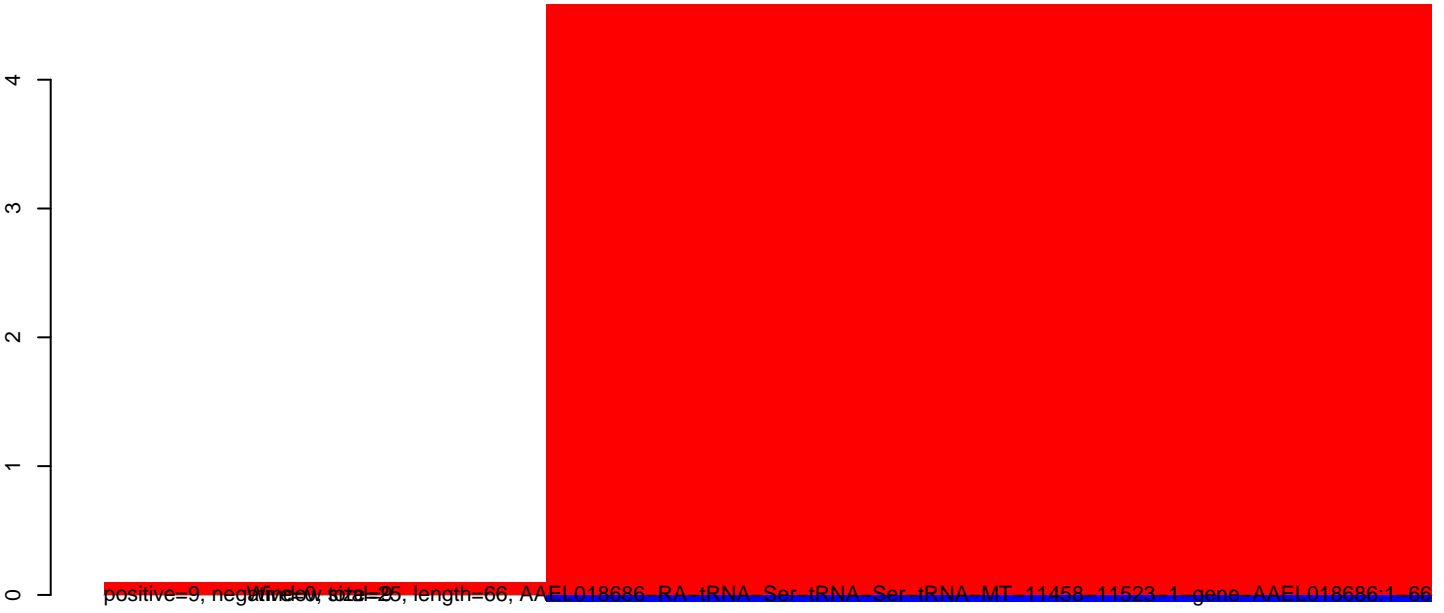
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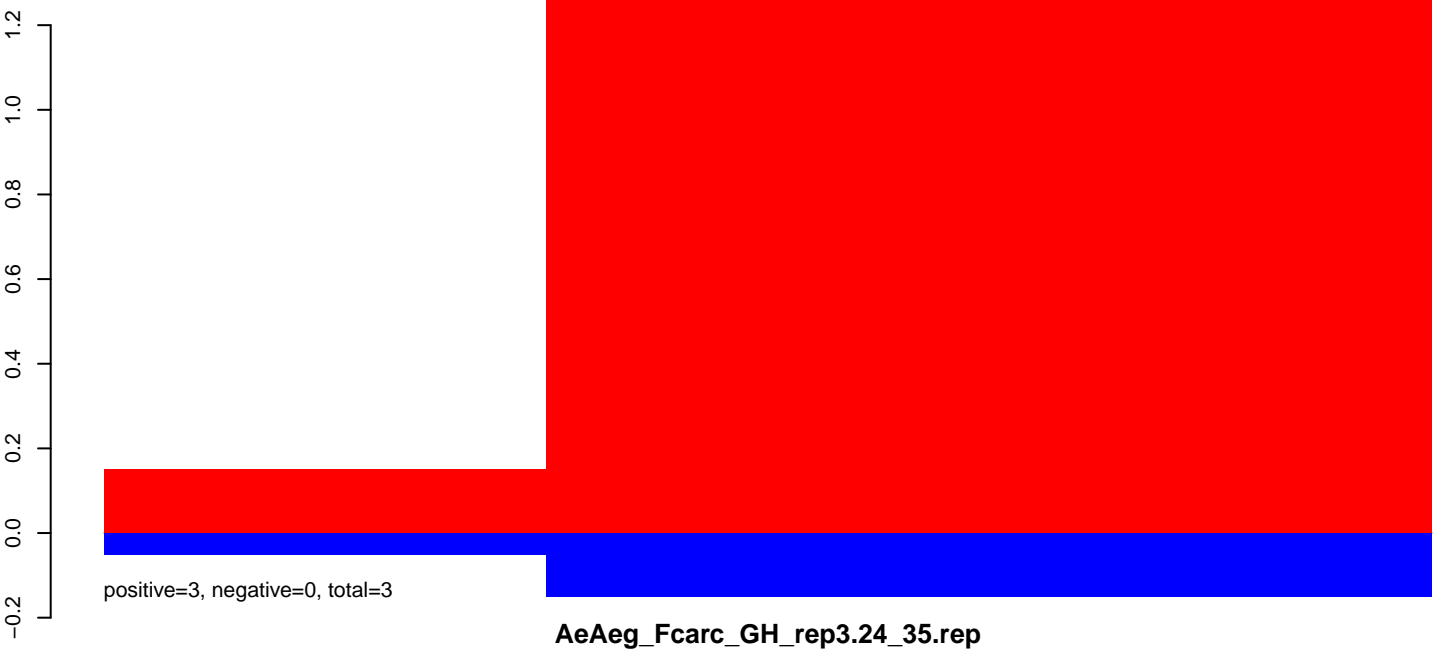


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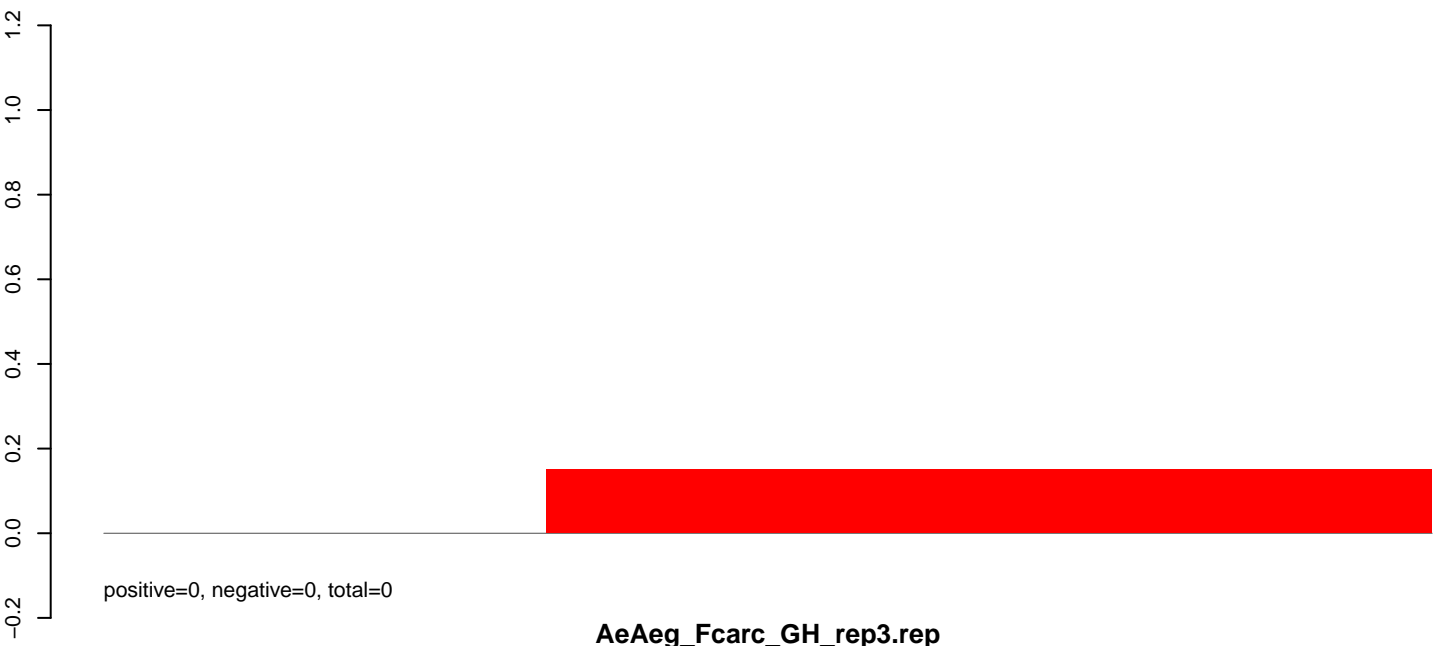


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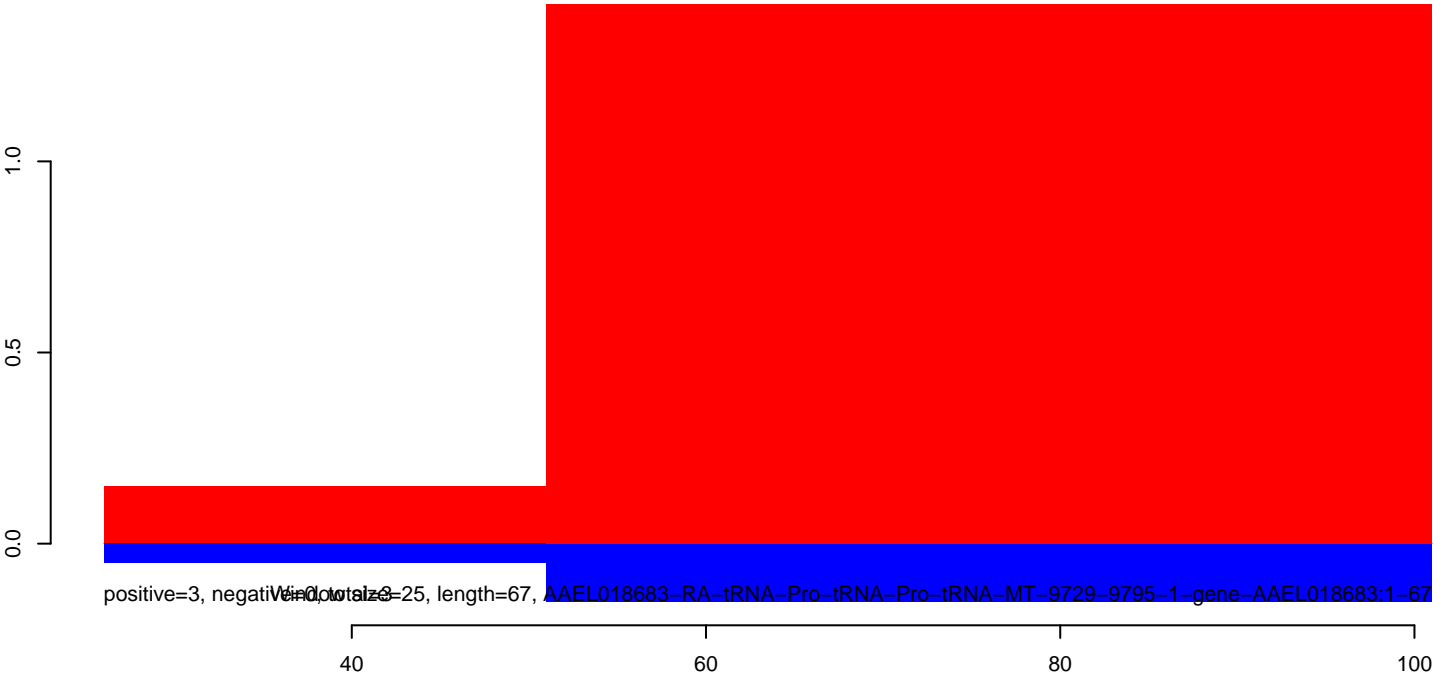
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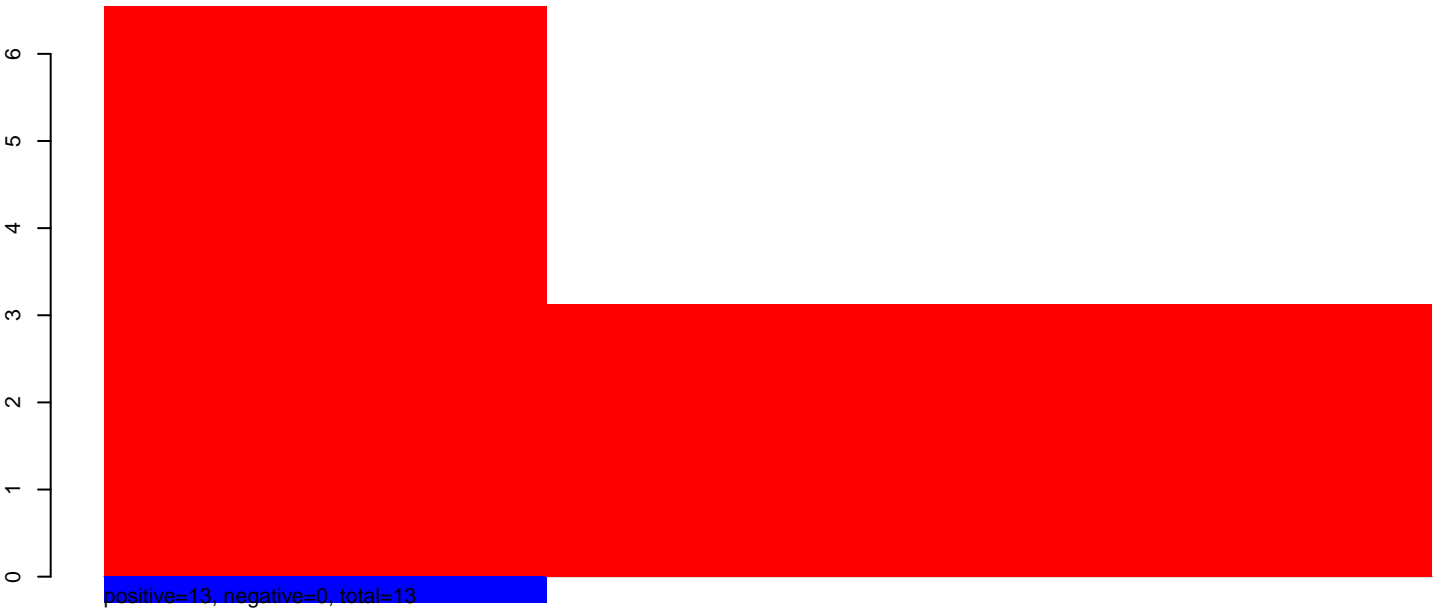
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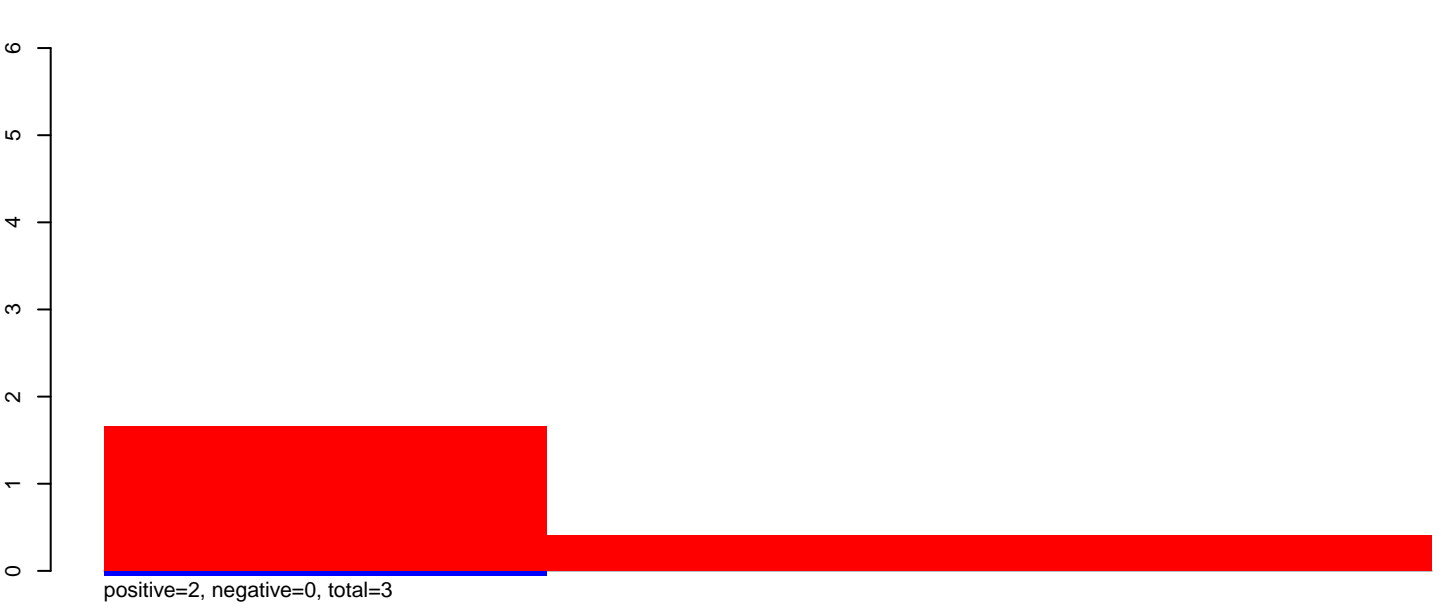
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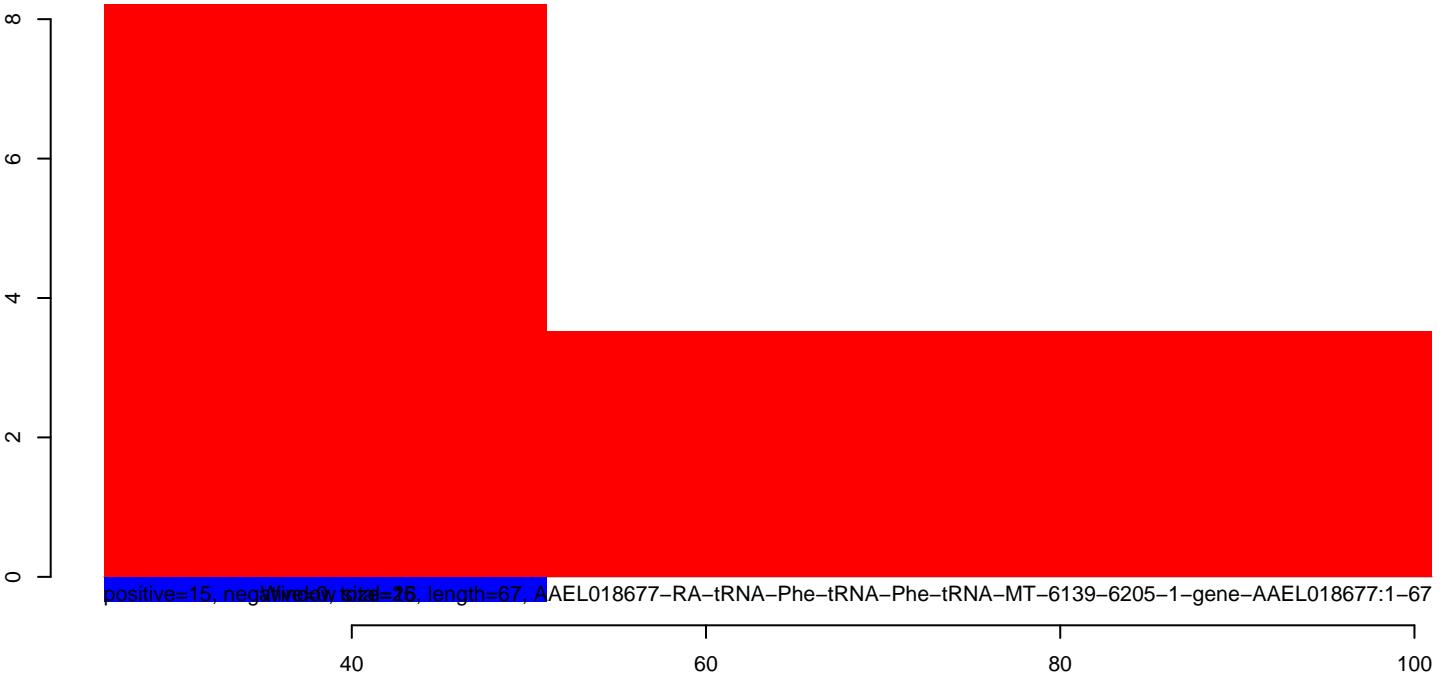
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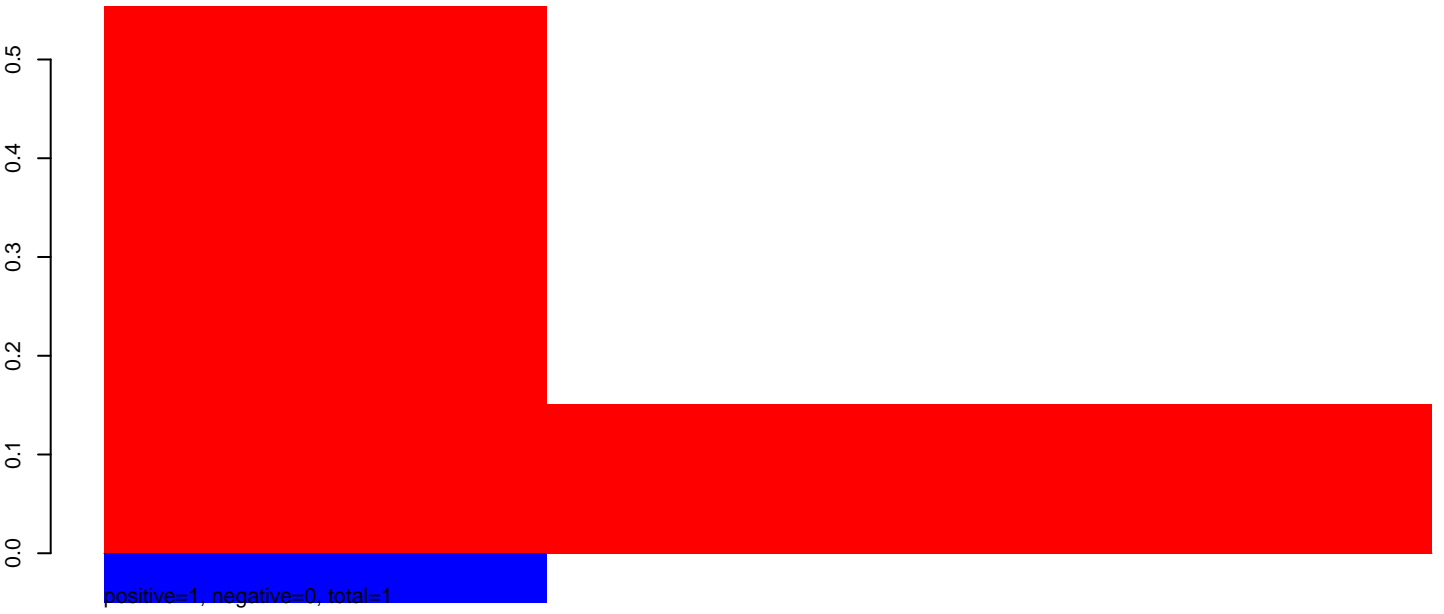


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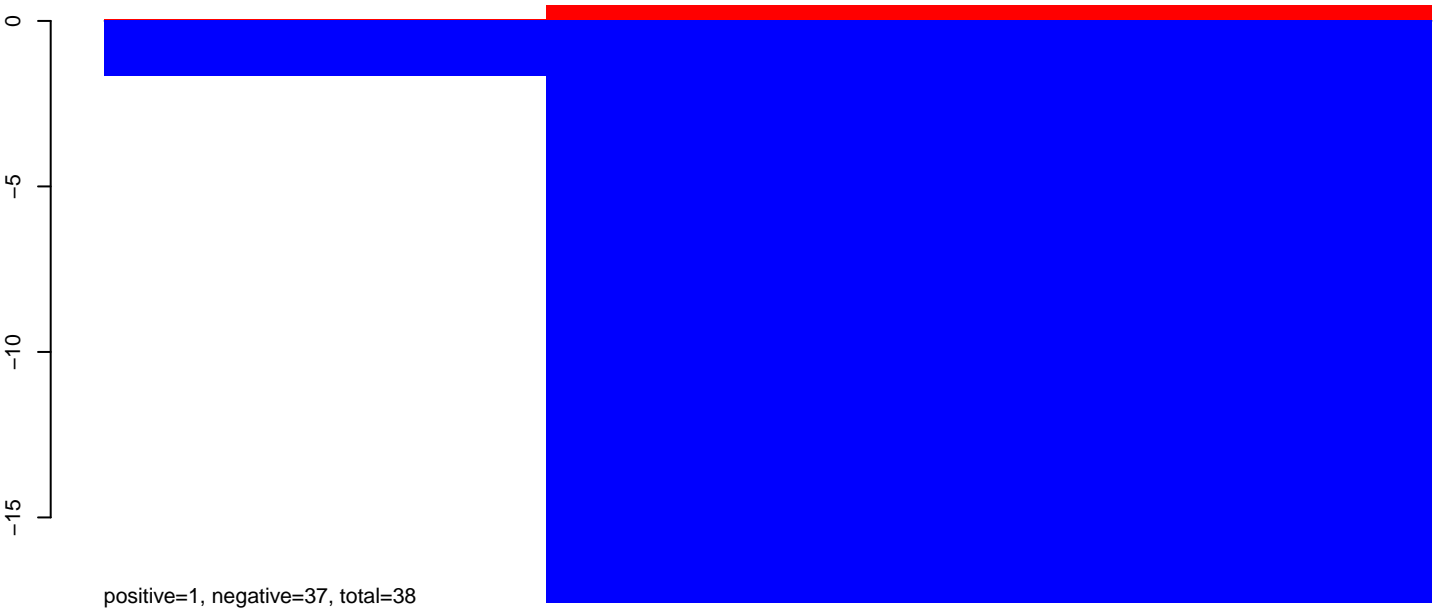
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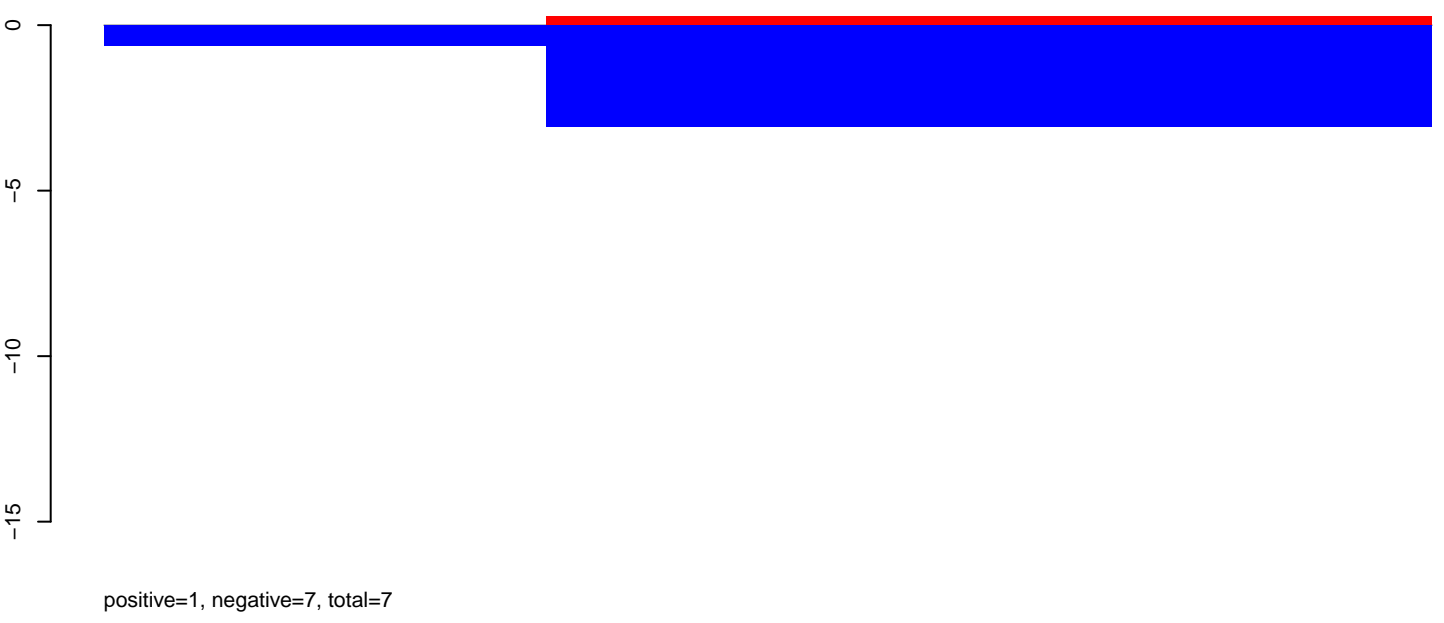
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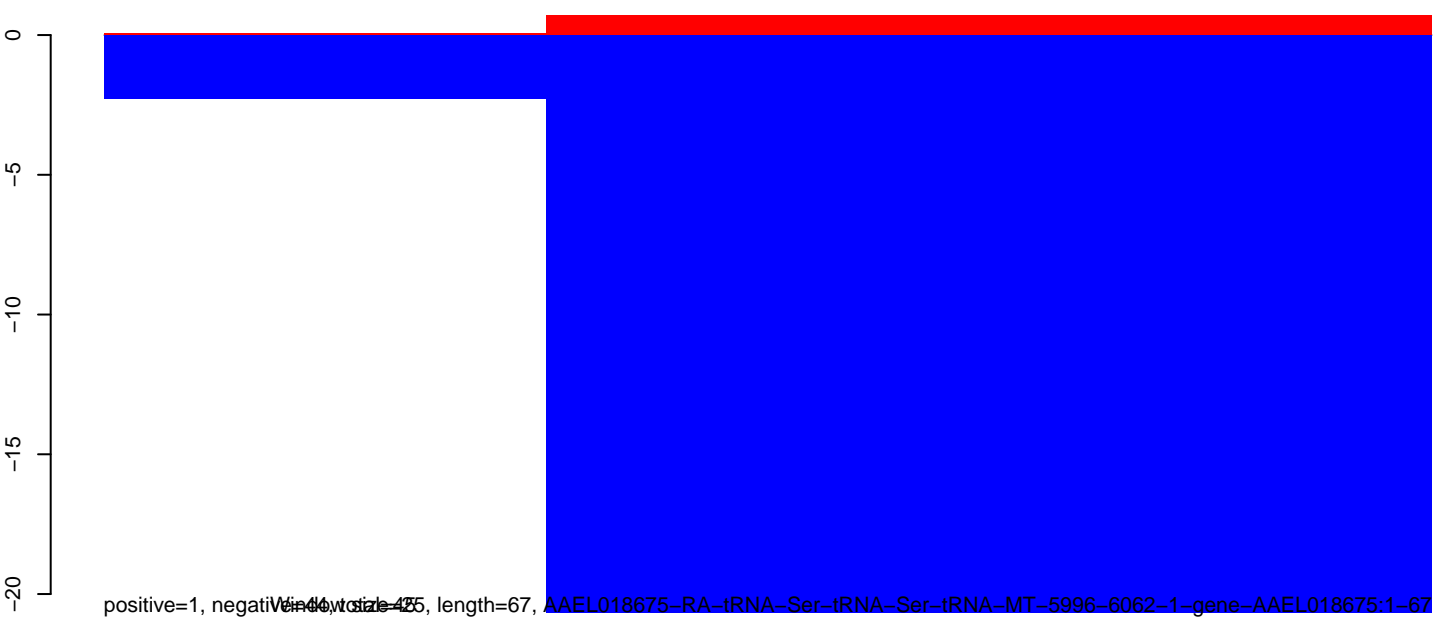
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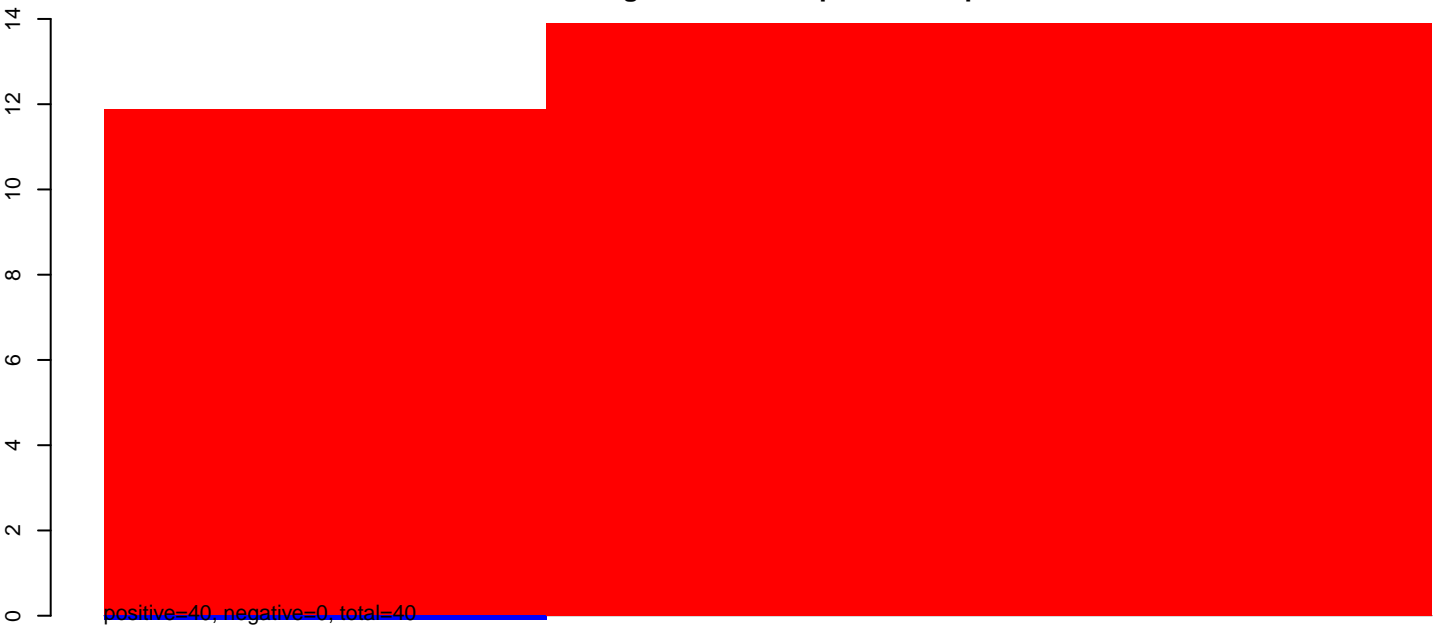


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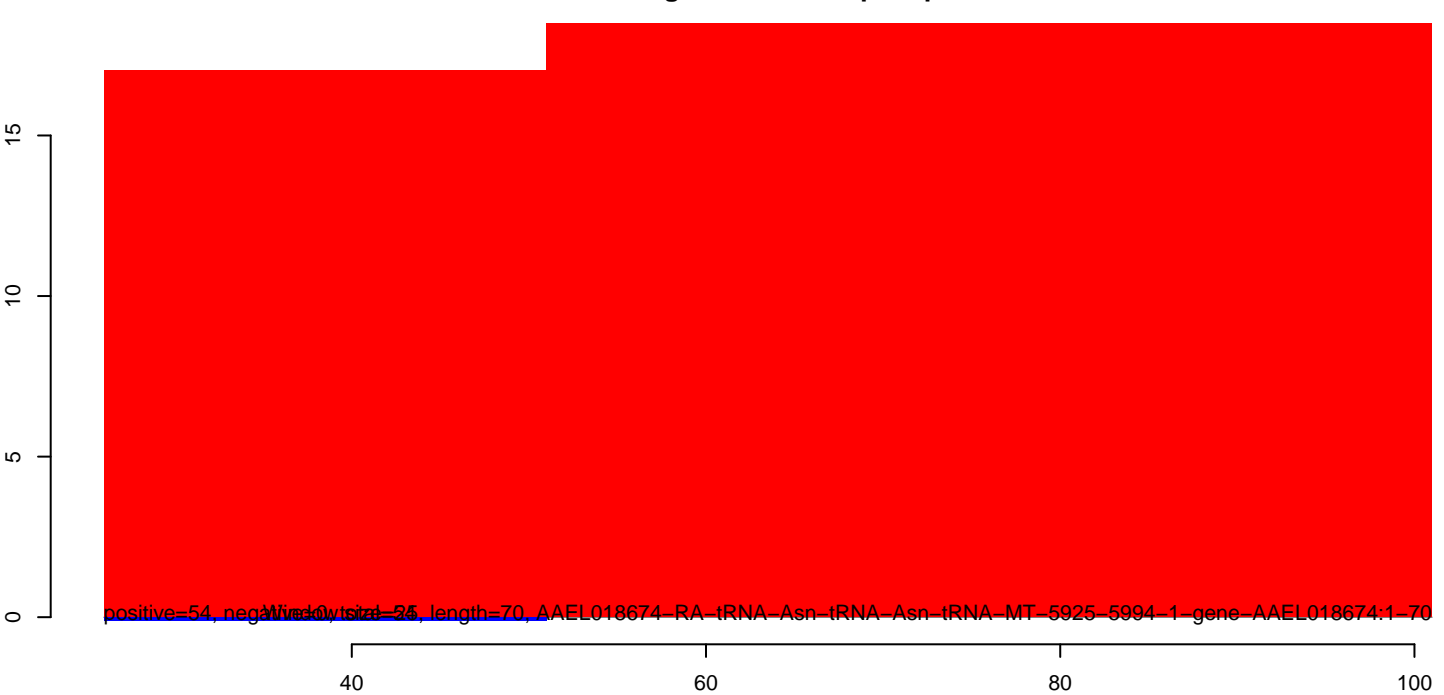
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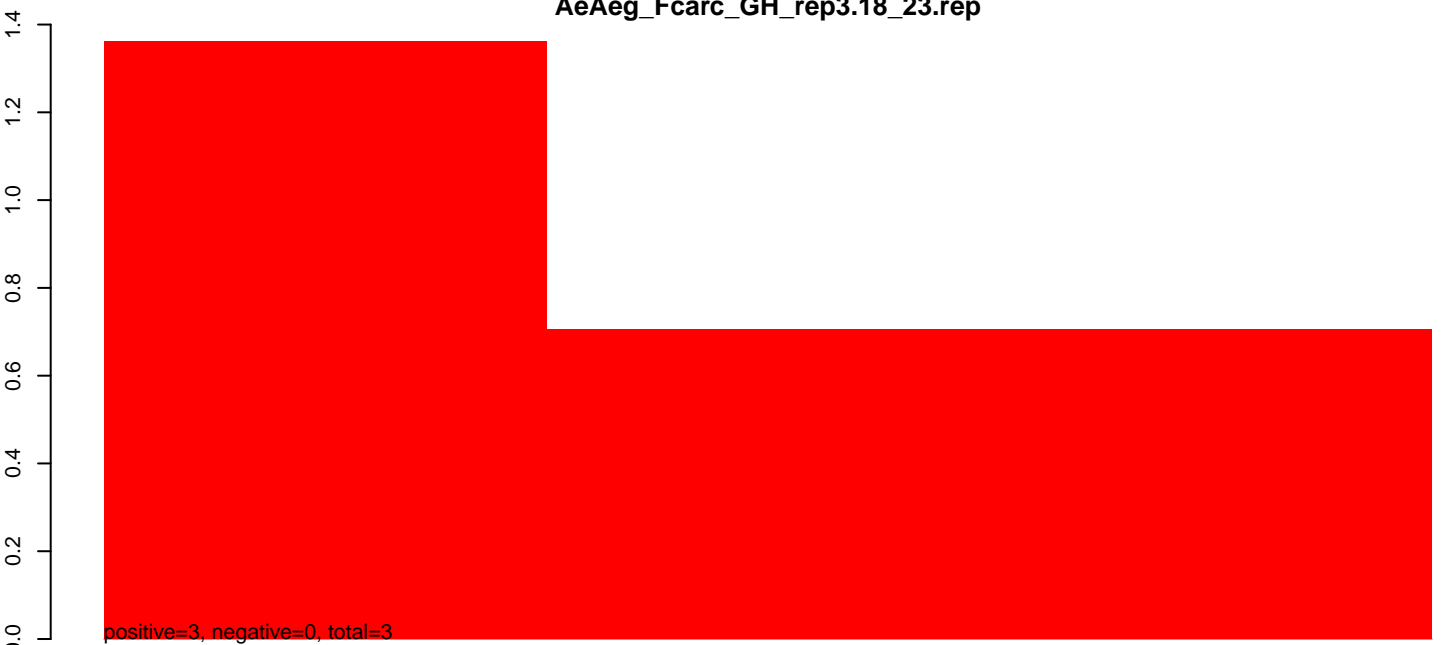
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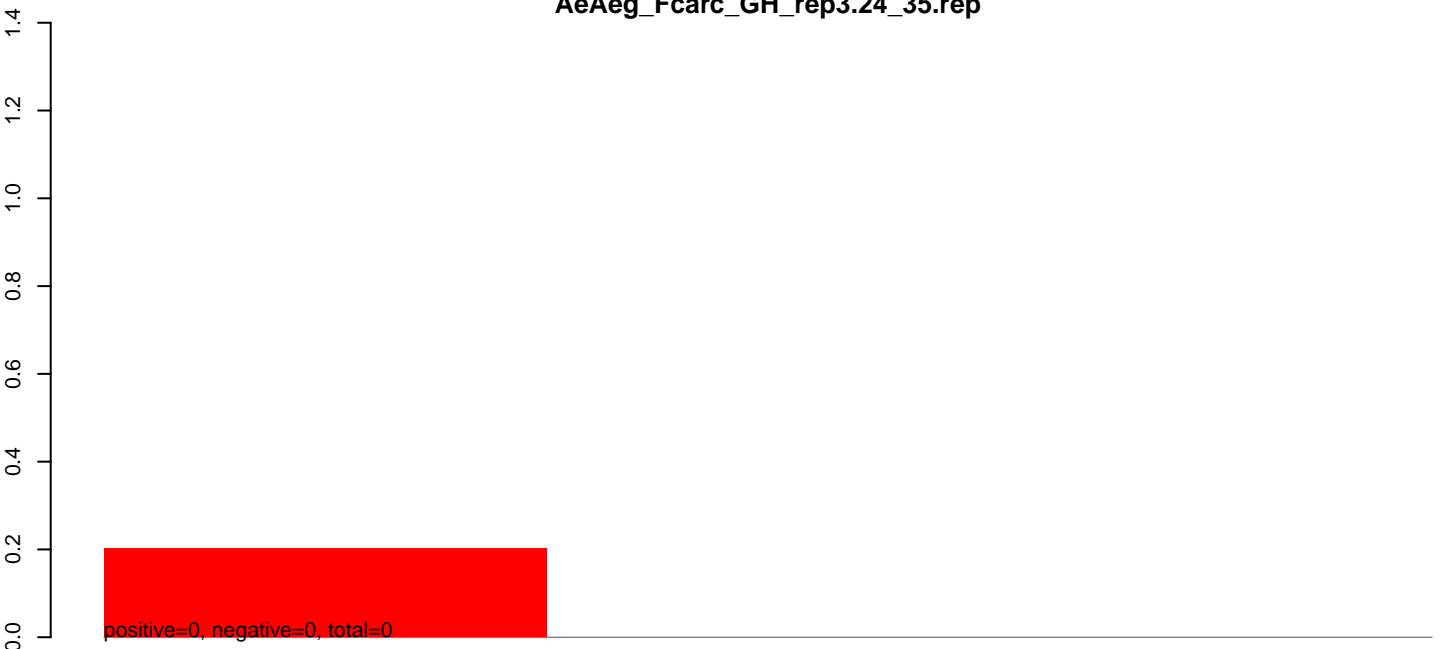
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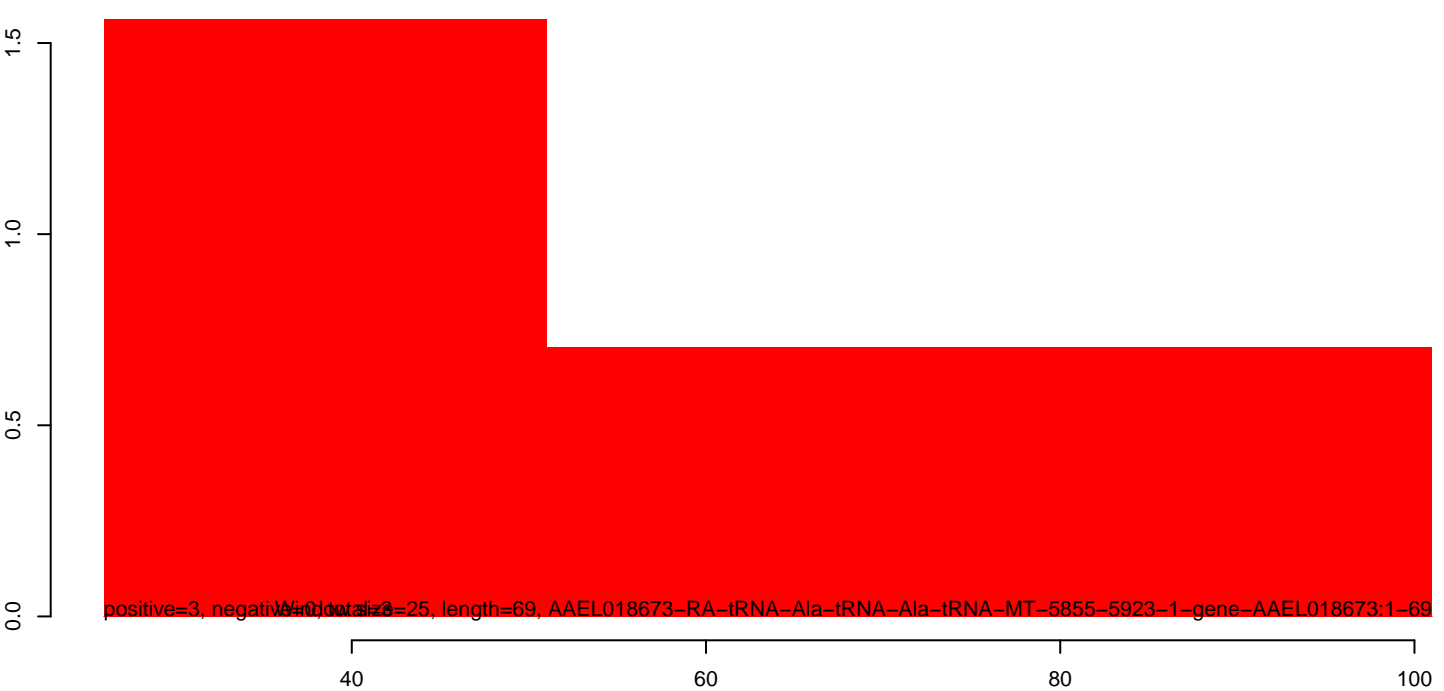
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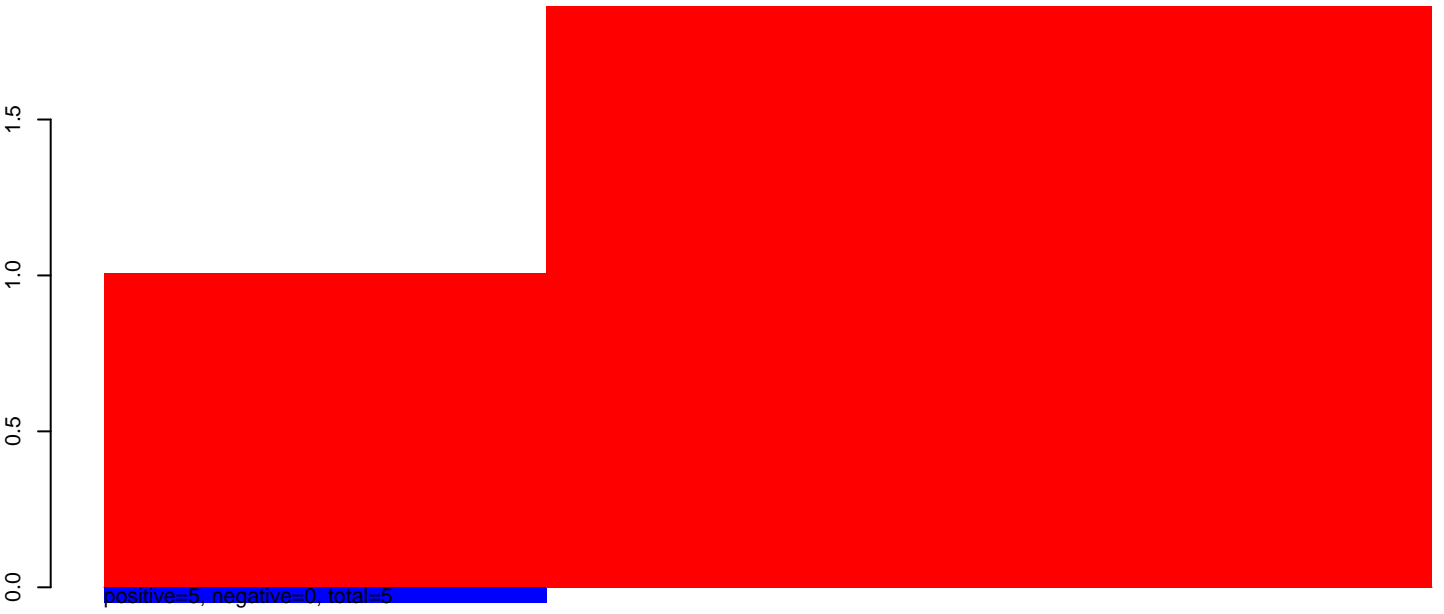
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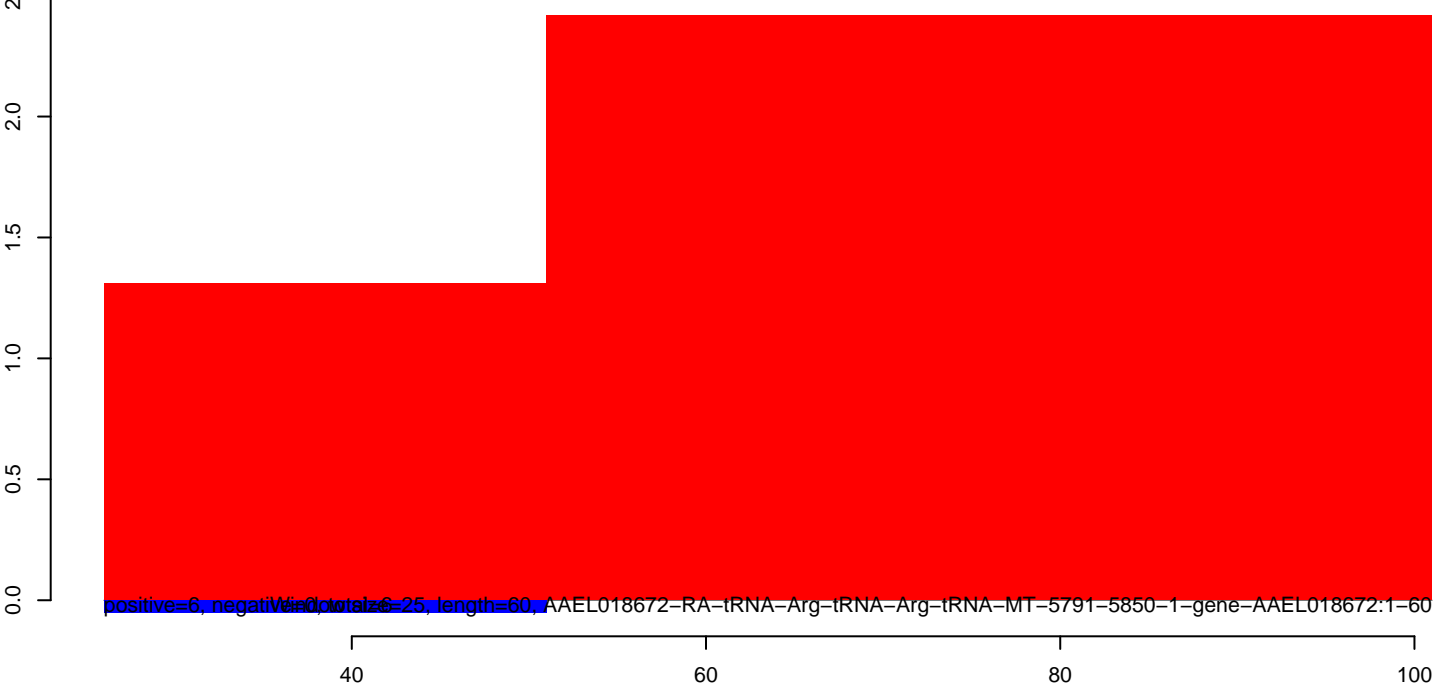
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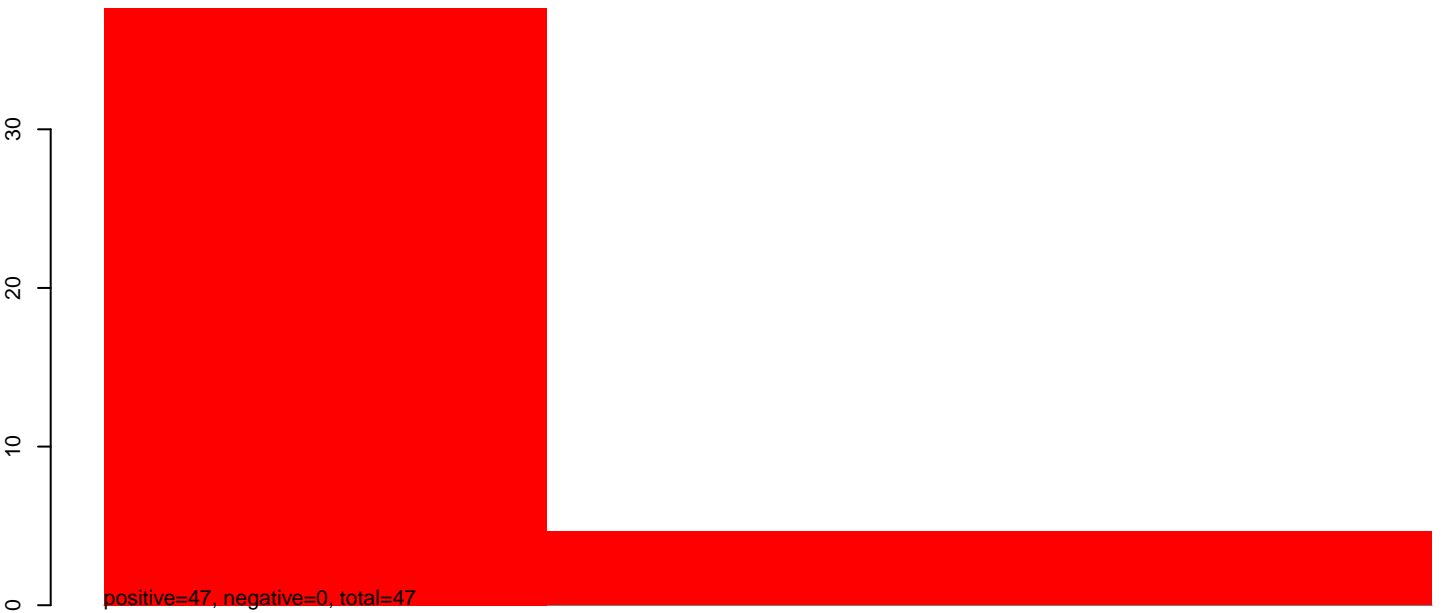
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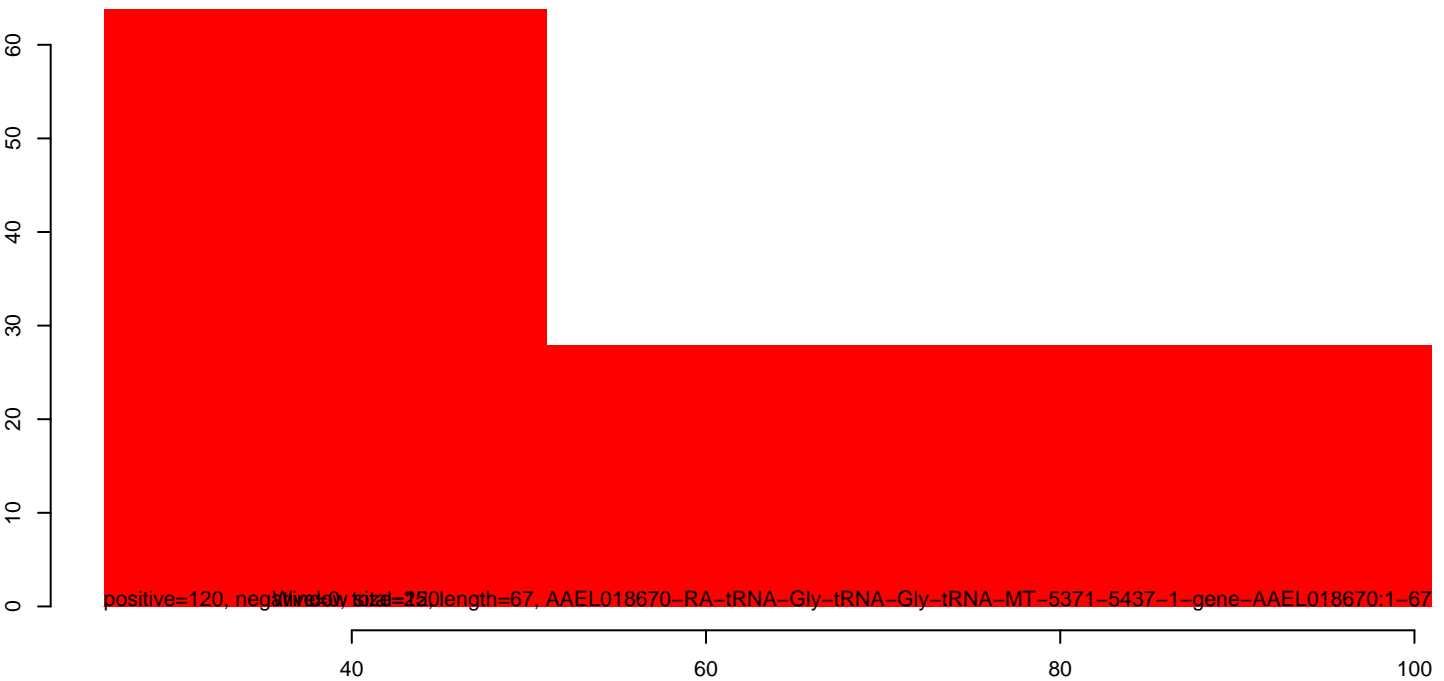
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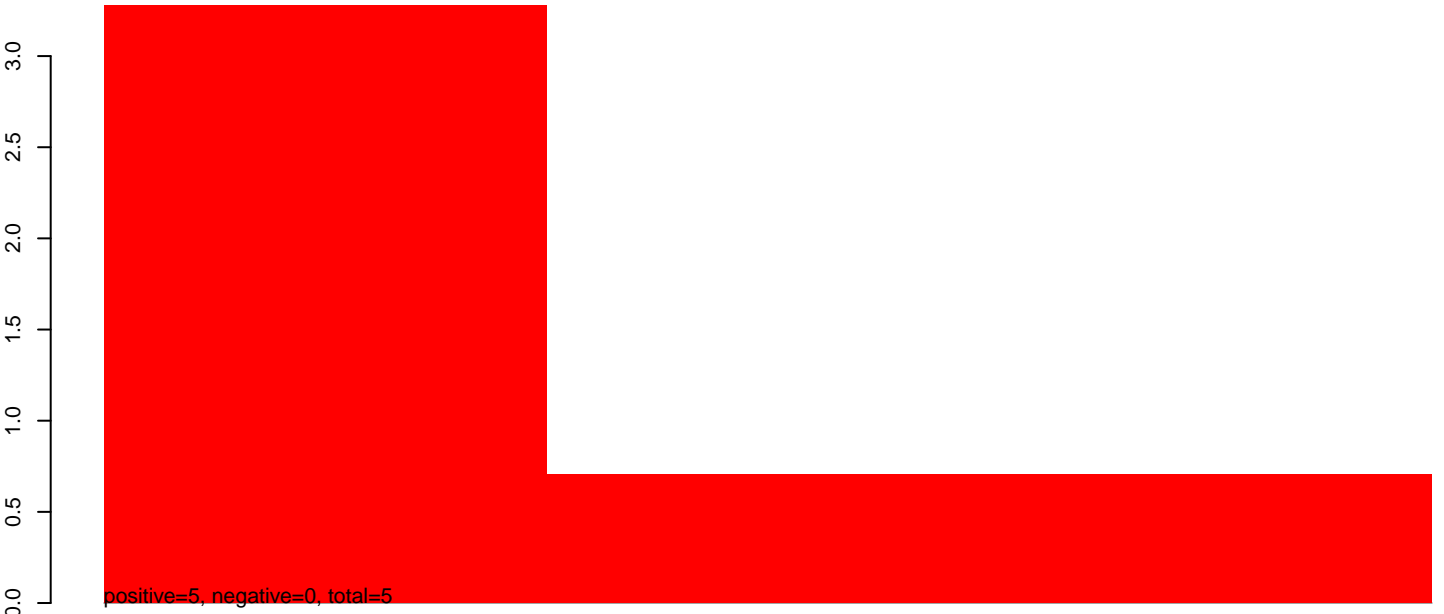
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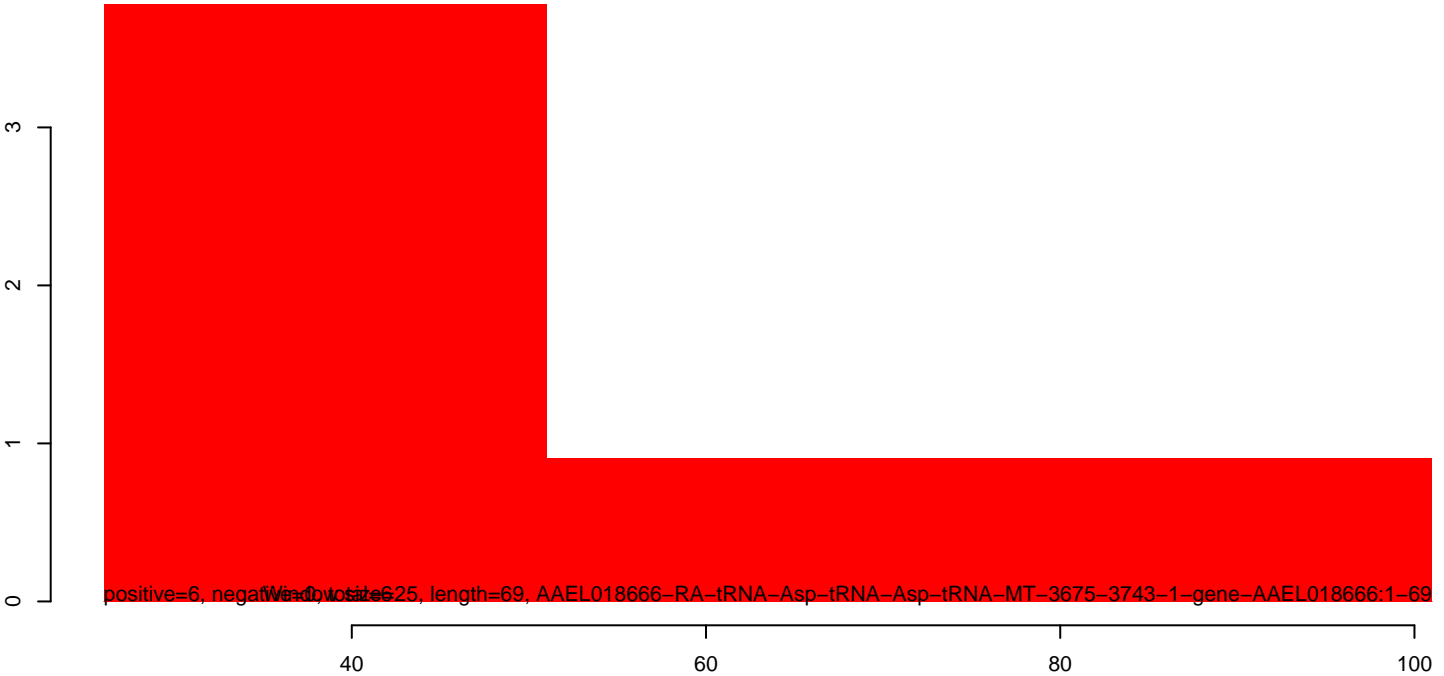
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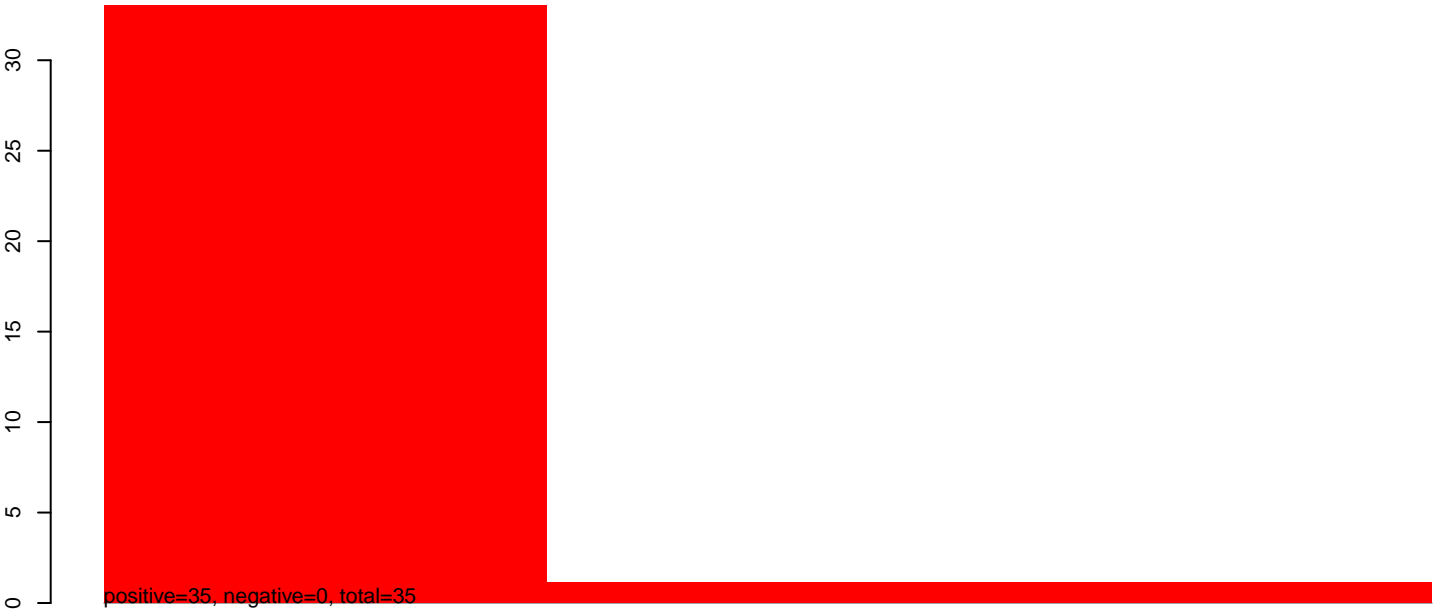
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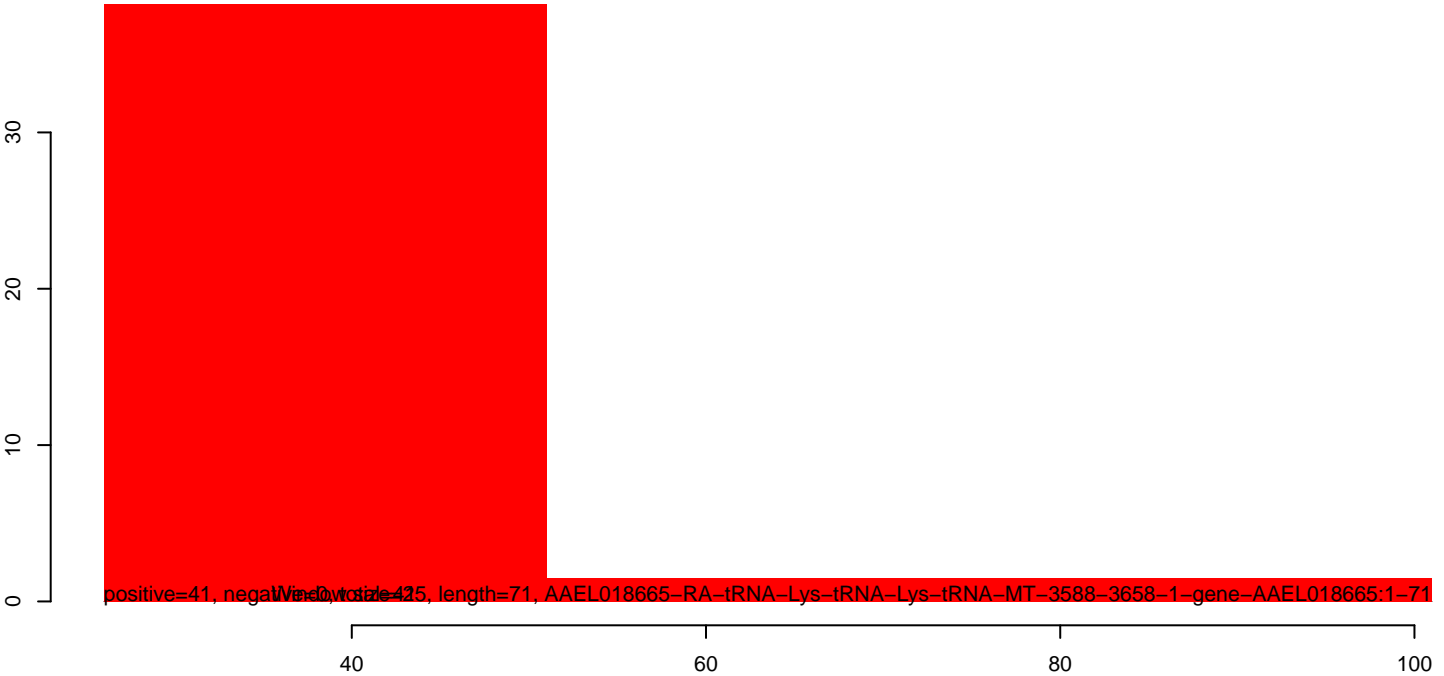
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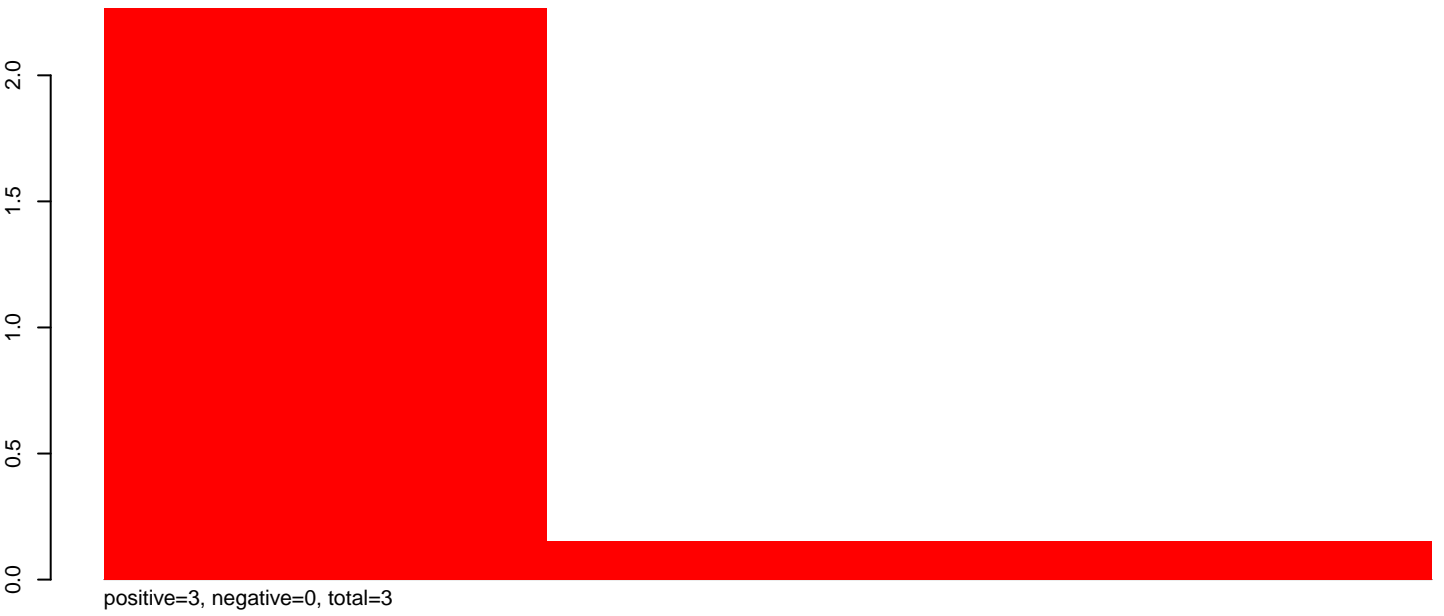
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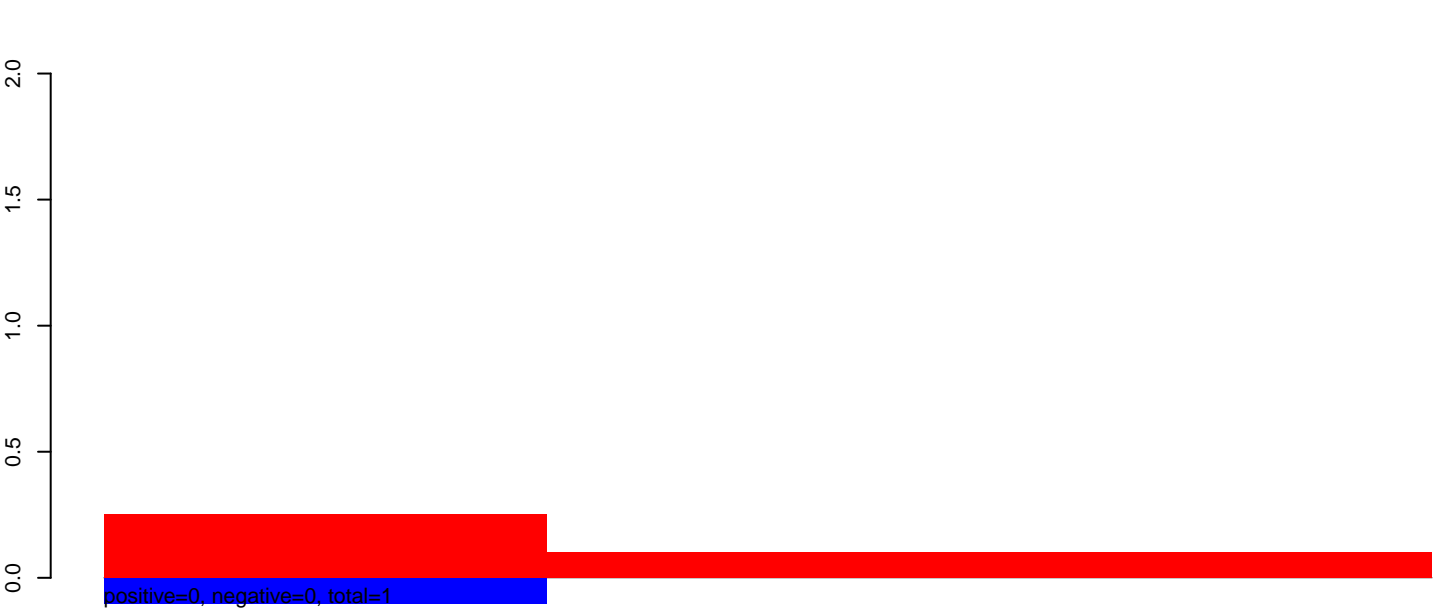
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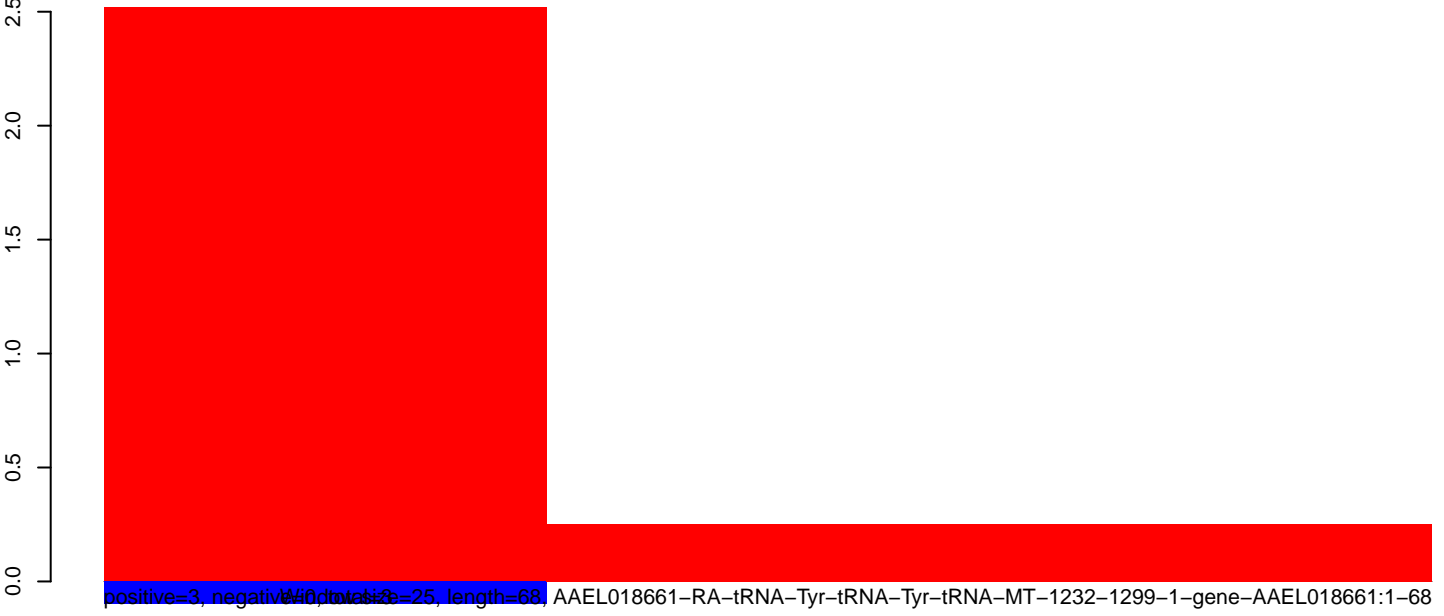
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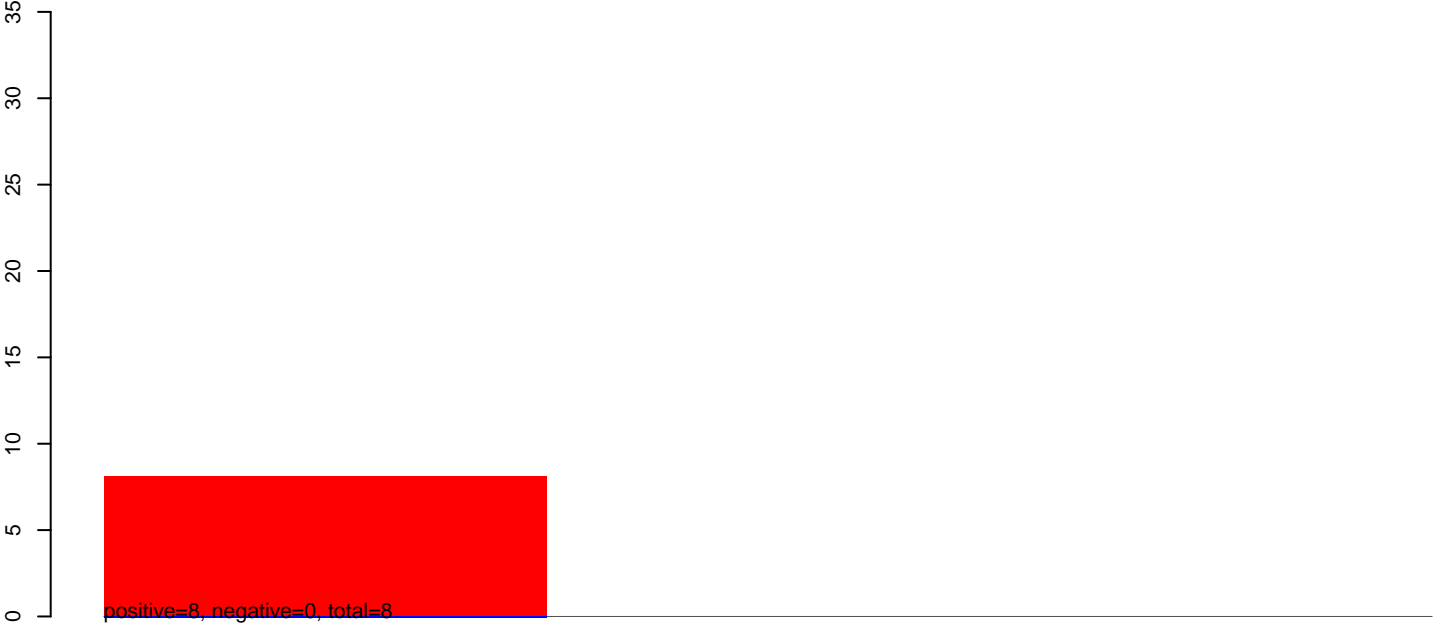


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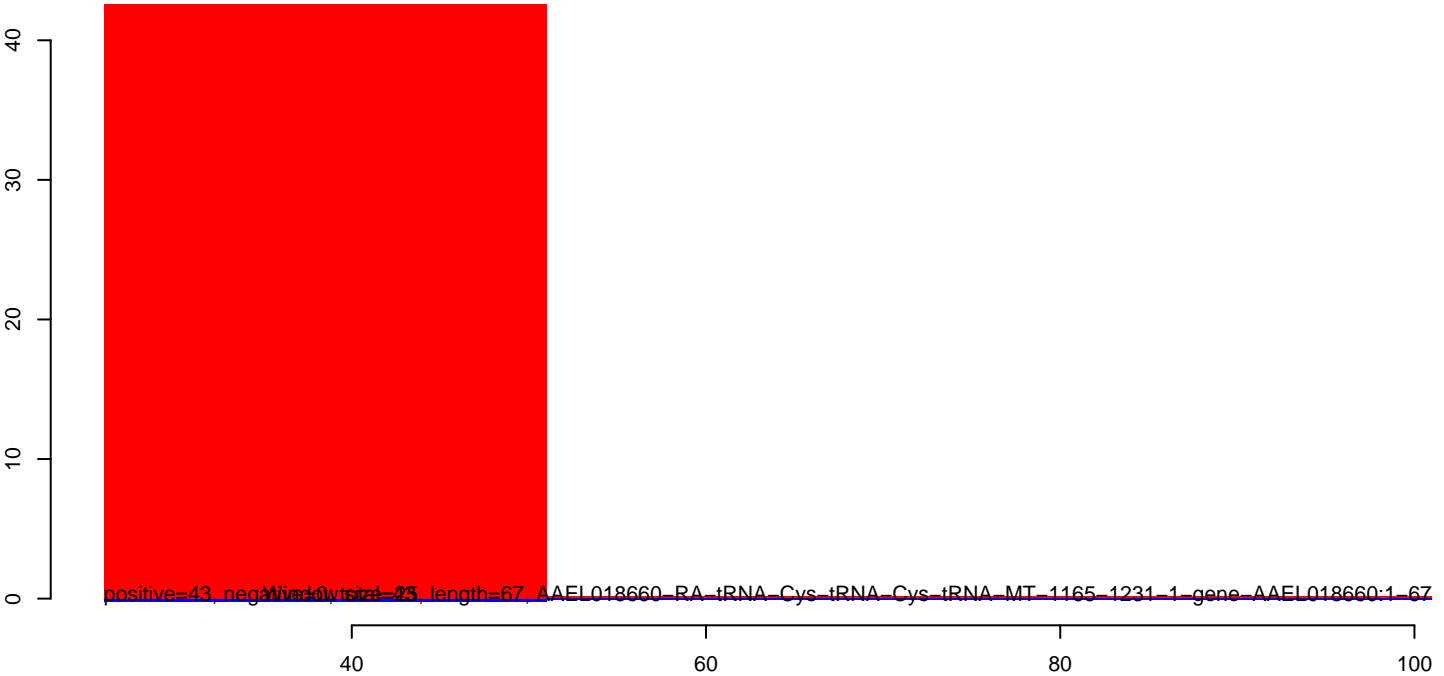
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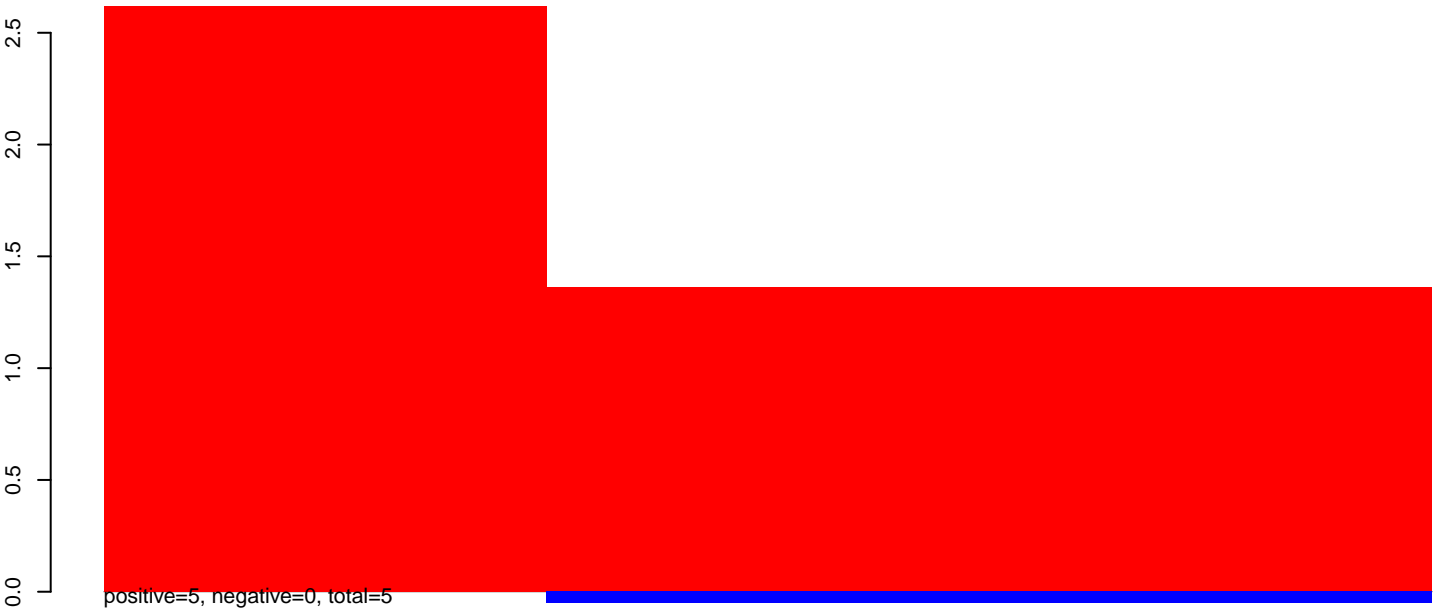
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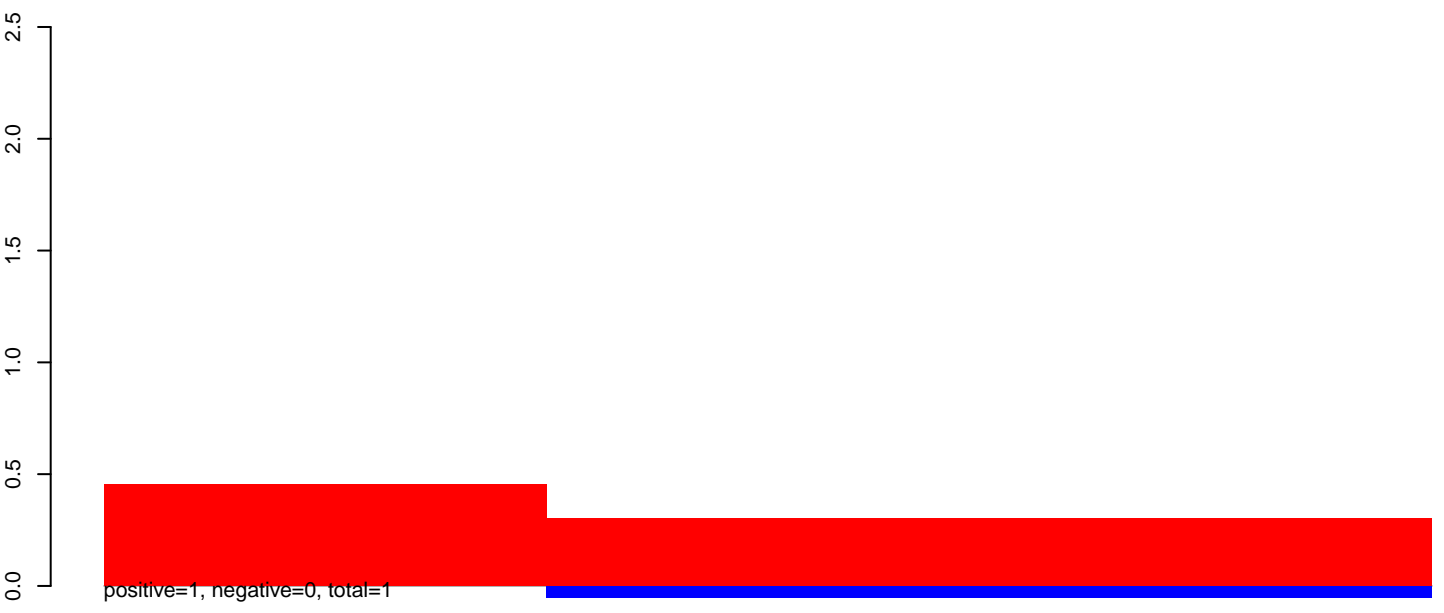
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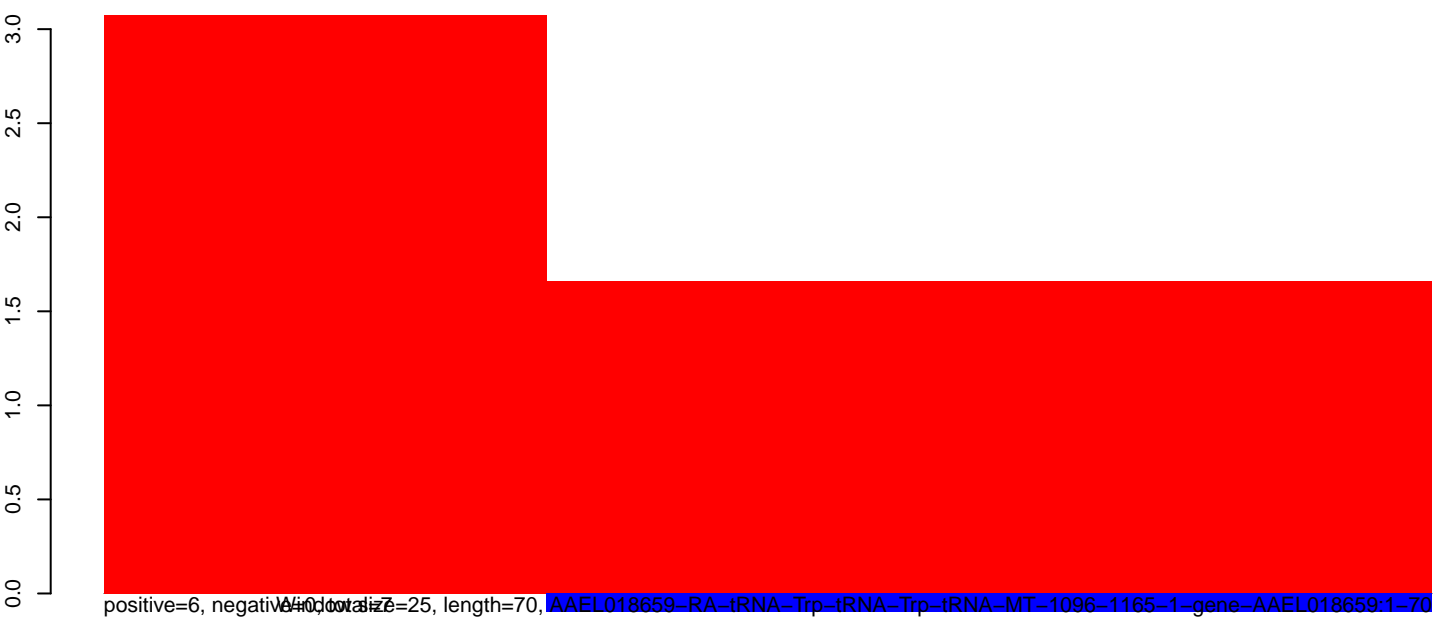
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AeAeg_Fcarc_GH_rep3.24_35.rep



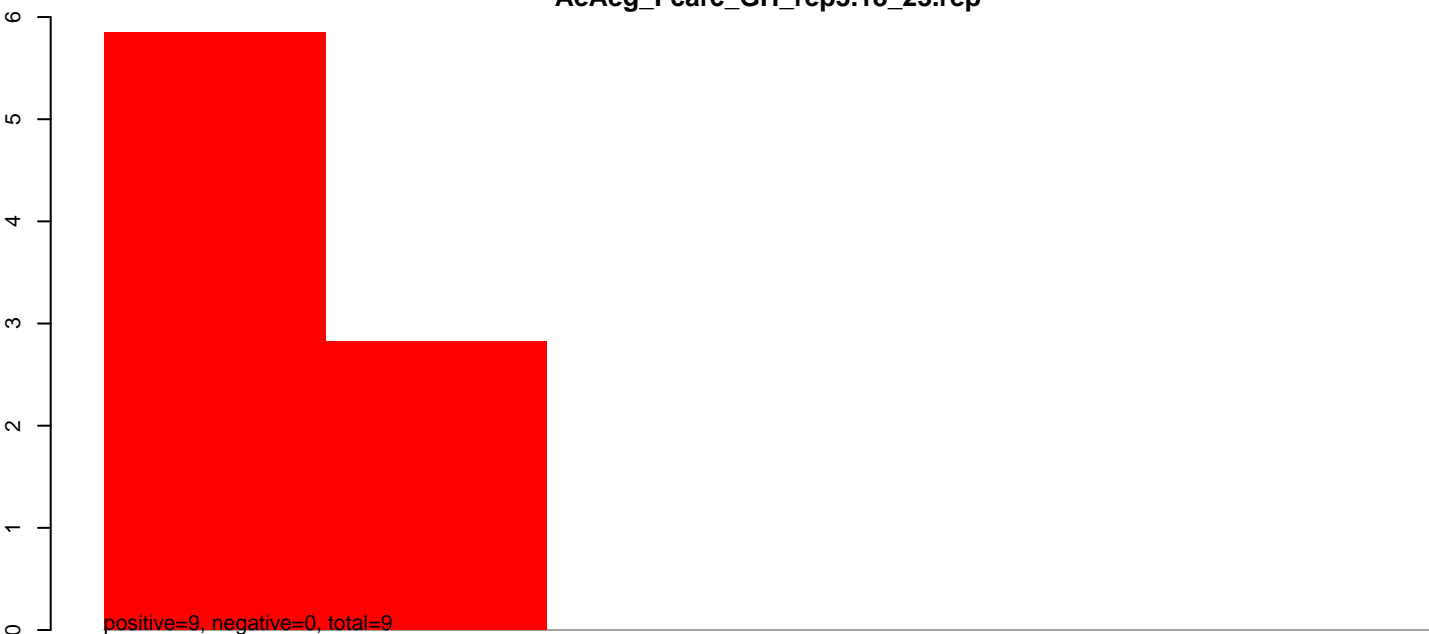
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AAEL018659-RA-tRNA-Trp-tRNA-Trp-tRNA-MT-1096-1165-1-gene-AAEL018659-1-70

40 60 80 100

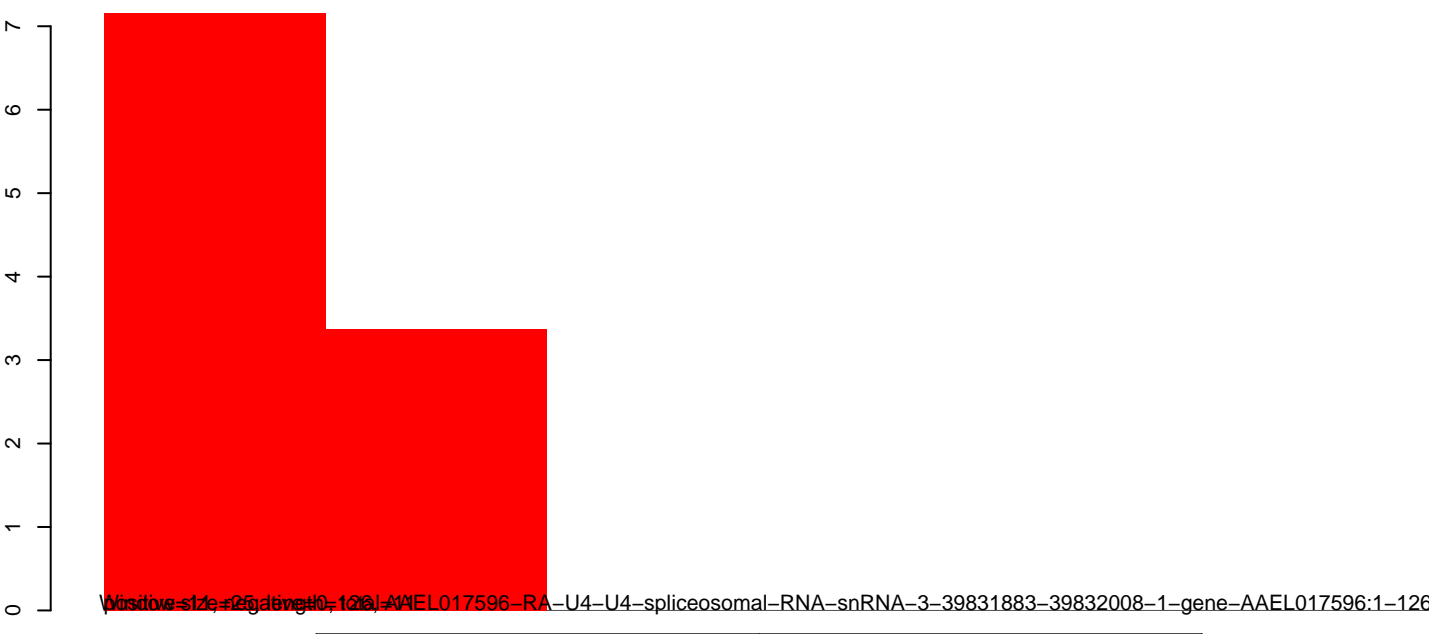
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AeAeg_Fcirc_GH_rep3.24_35.rep



AeAeg_Fcirc_GH_rep3.rep



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AeAeg_Fcarc_GH_rep3.24_35.rep

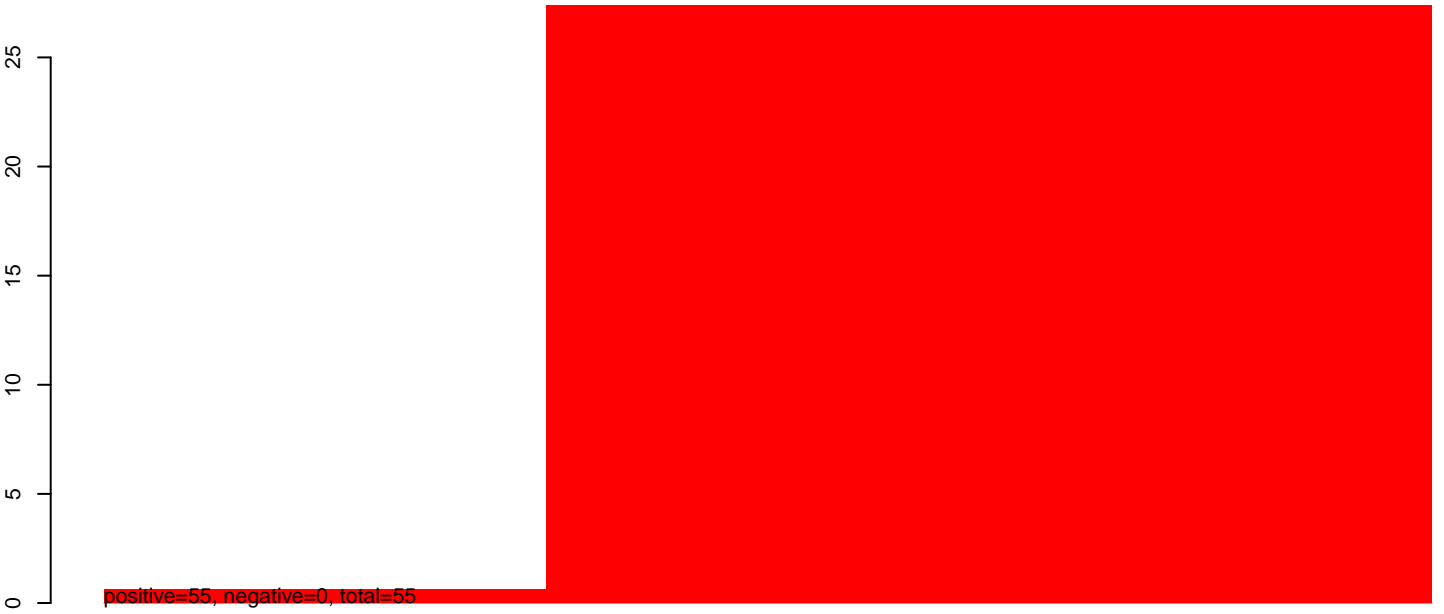


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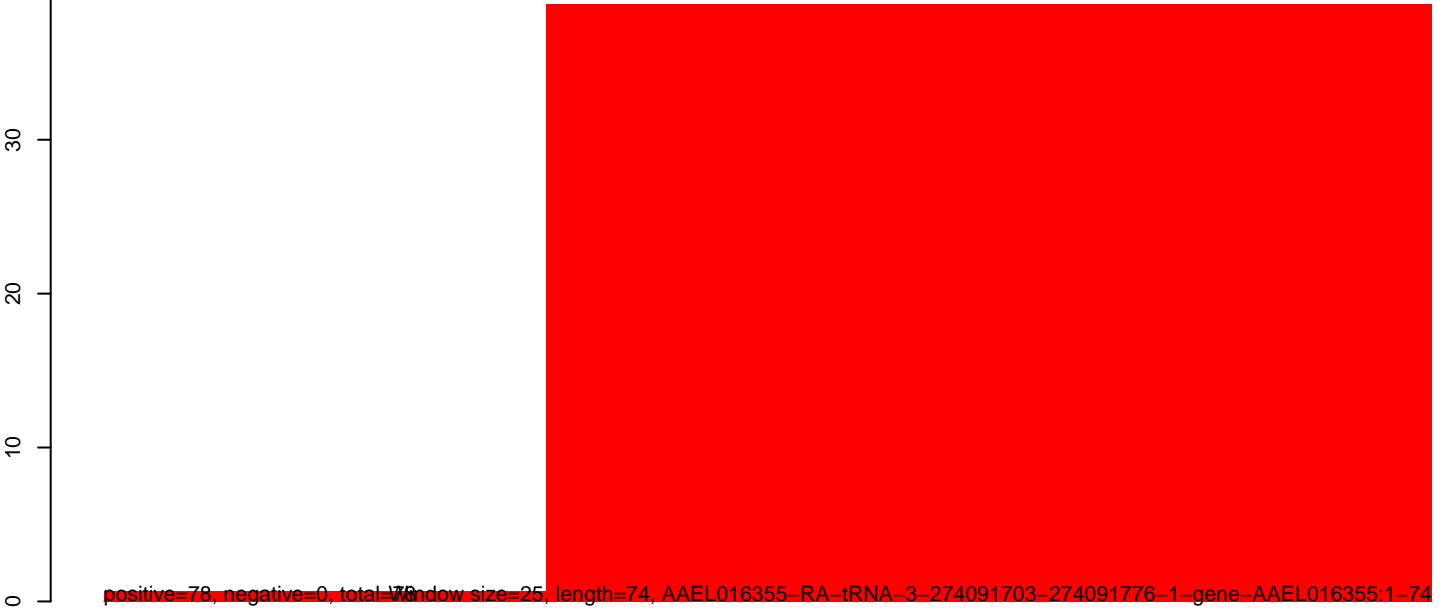
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AeAeg_Fcarc_GH_rep3.24_35.rep



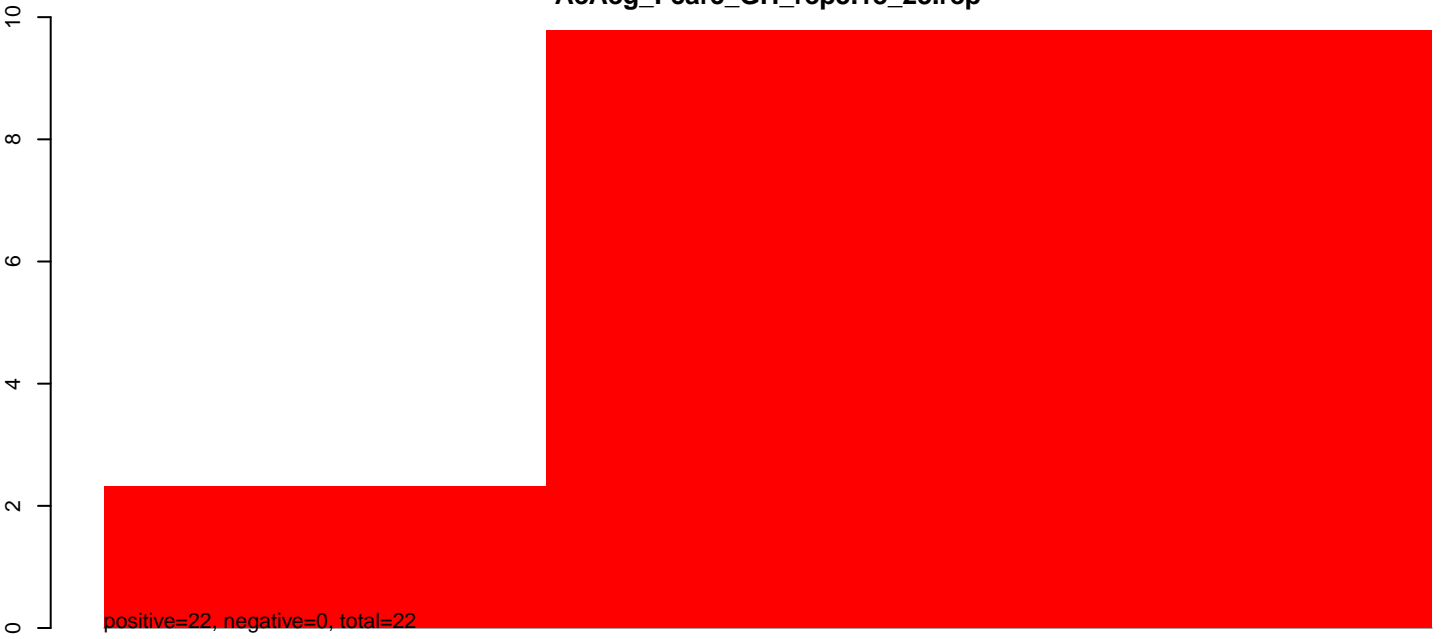
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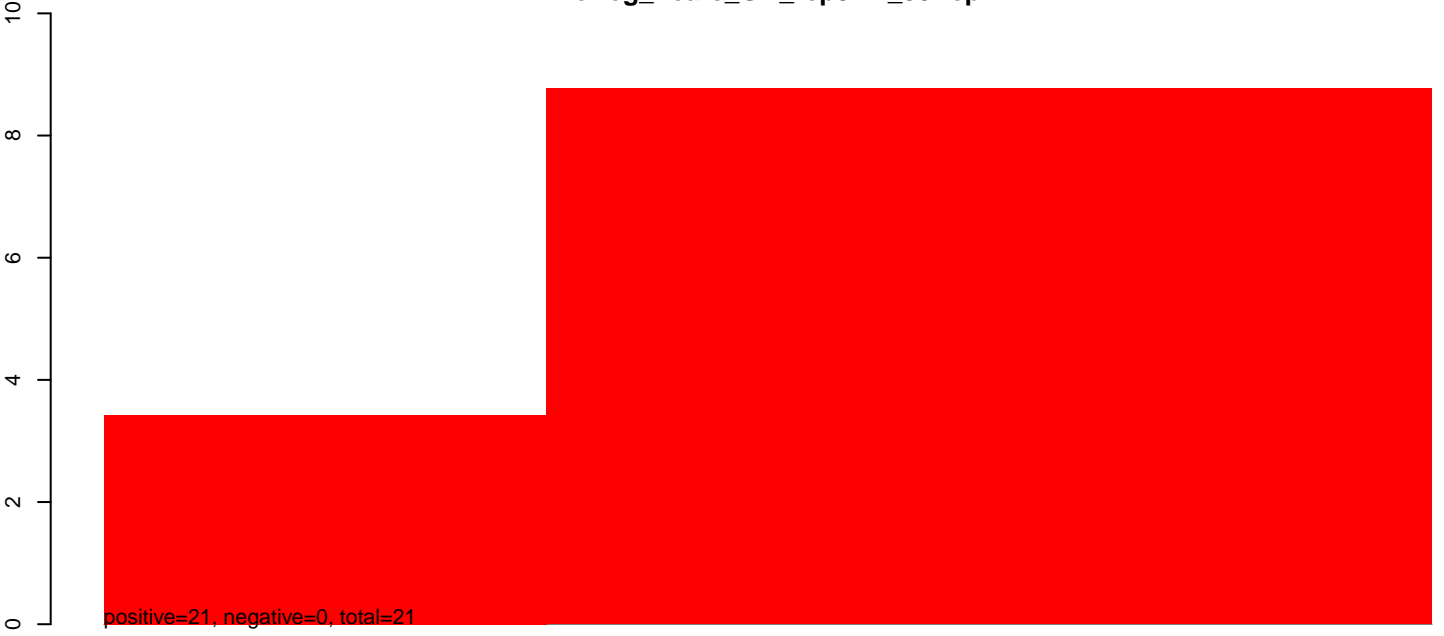
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40 60 80 100

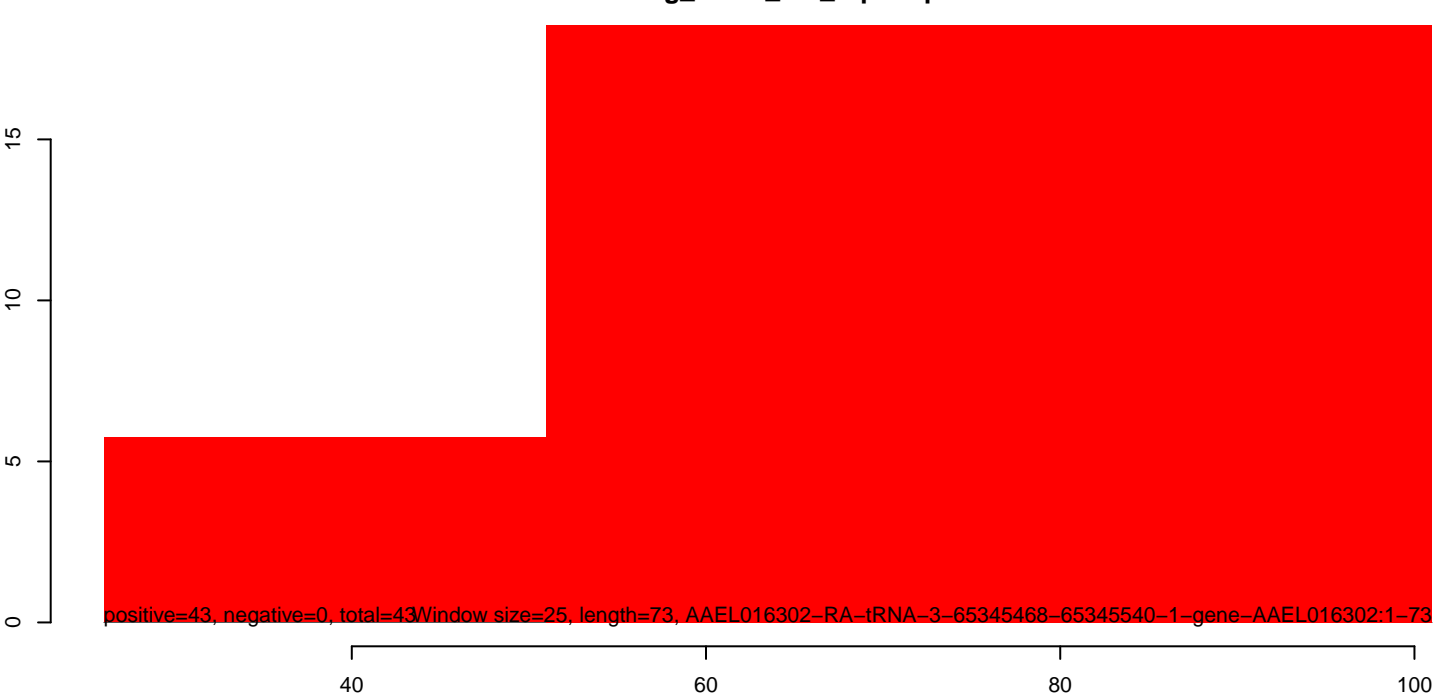
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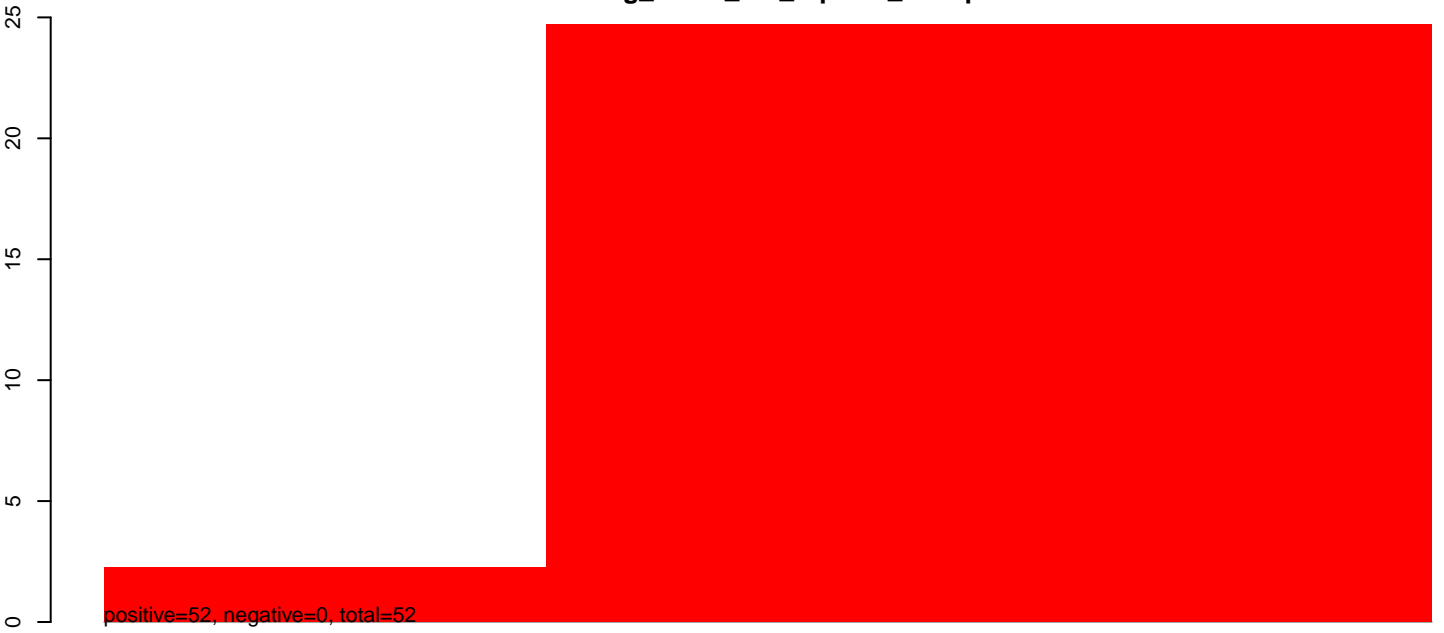
AeAeg_Fcarc_GH_rep3.24_35.rep



AeAeg_Fcarc_GH_rep3.rep



AeAeg_Fcarc_GH_rep3.18_23.rep



AeAeg_Fcarc_GH_rep3.24_35.rep



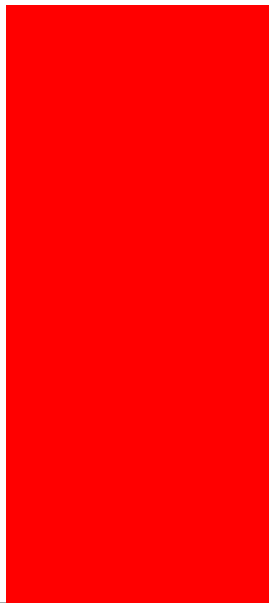
AeAeg_Fcarc_GH_rep3.rep



Window size=25, length=73, AAEL016098-RA-tRNA-2-56316242-56316314-1-gene-AAEL016098:1-73

AeAeg_Fcarc_GH_rep3.18_23.rep

positive=7, negative=0, total=7



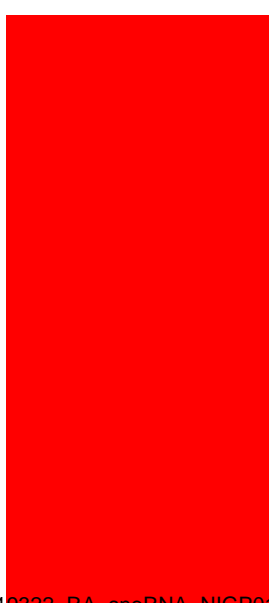
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positive=0, negative=0, total=0



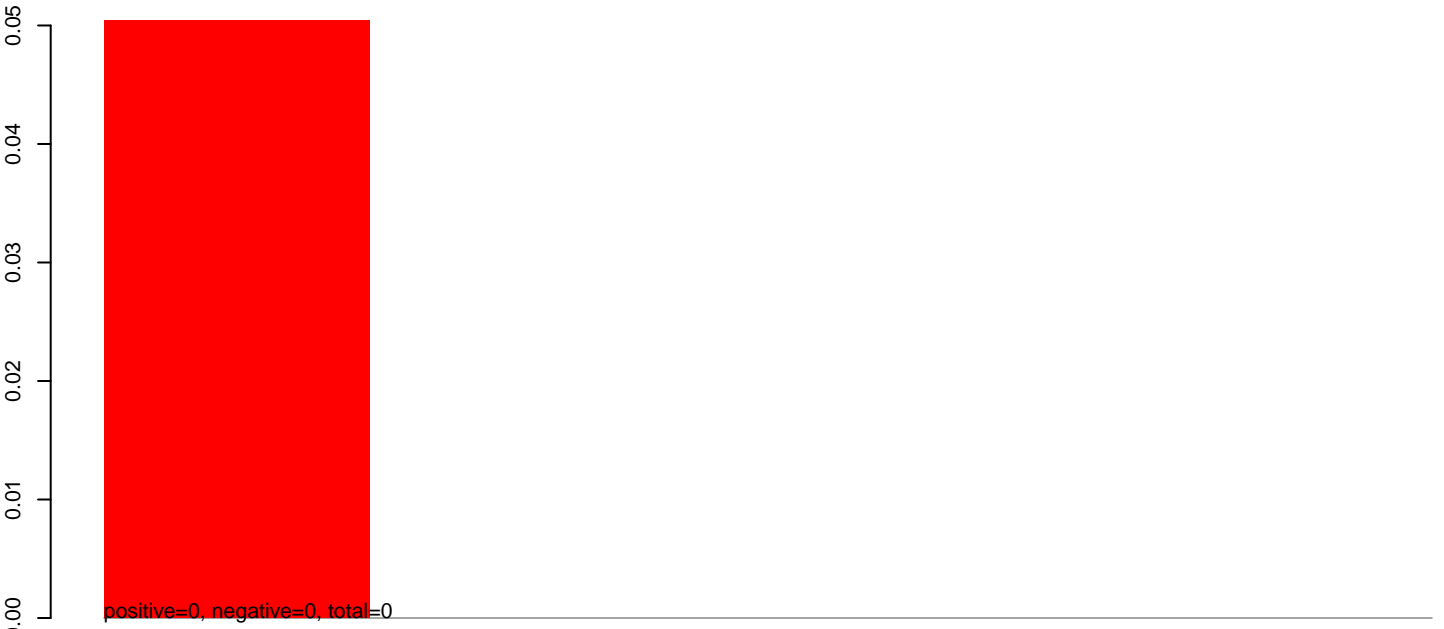
AeAeg_Fcarc_GH_rep3.rep

positive=7, negative=0, total=7
Window size=25, length=105, AAEL019322-RA-snoRNA-NIGP01001727-210524-210628-1-gene-AAEL019322:1-105

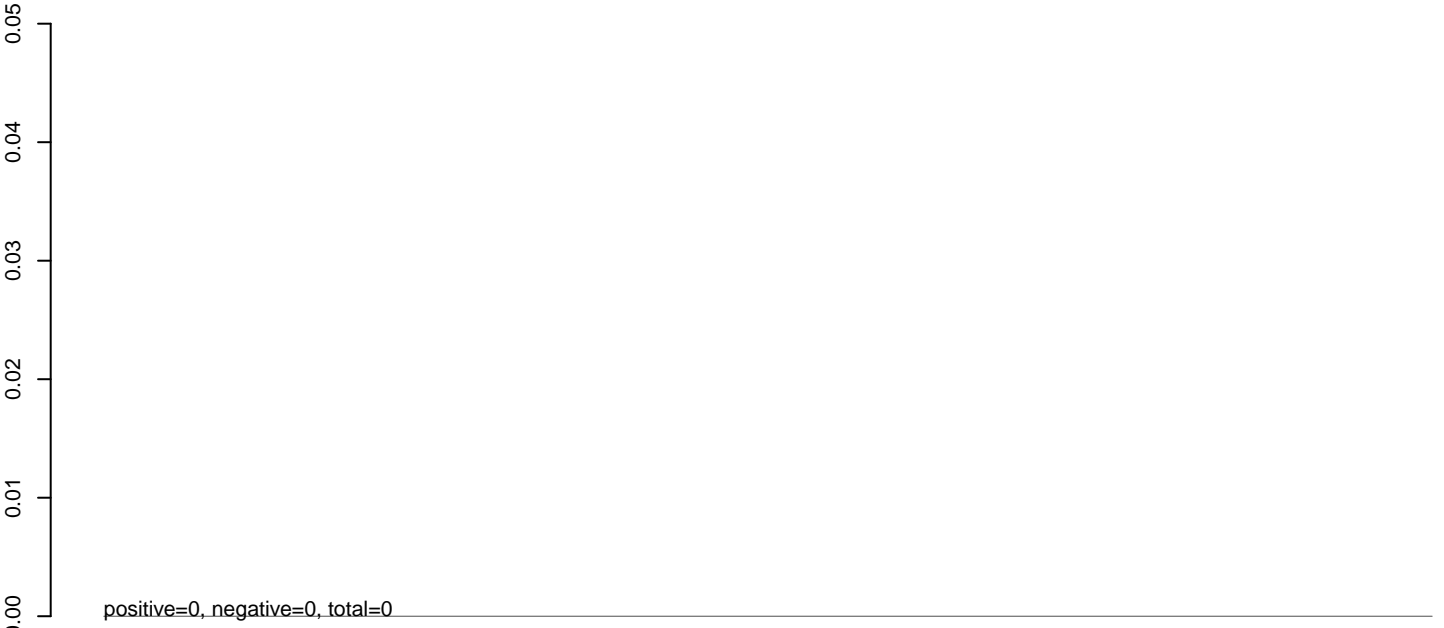


40 60 80 100 120 140

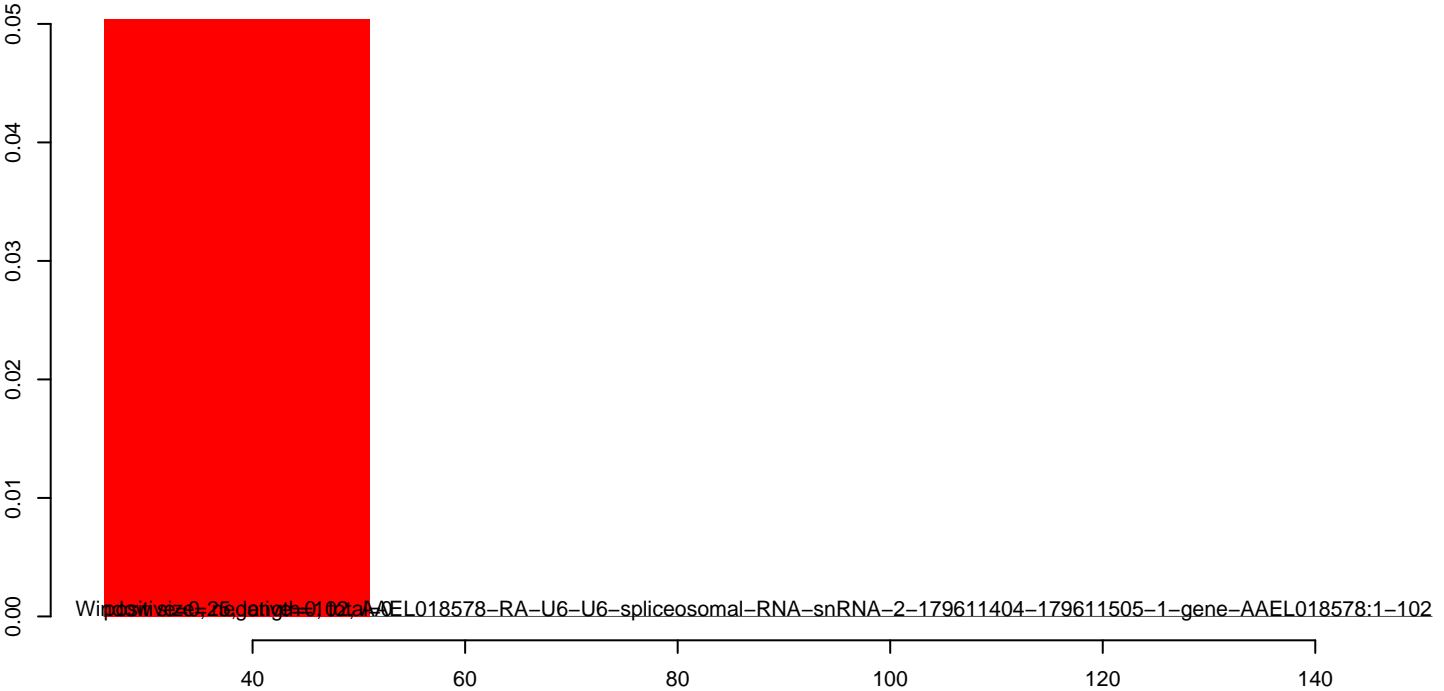
AeAeg_Fcarc_GH_rep3.18_23.rep



AeAeg_Fcarc_GH_rep3.24_35.rep



AeAeg_Fcarc_GH_rep3.rep



AeAeg_FcArc_GH_rep3.18_23.rep



AeAeg_FcArc_GH_rep3.24_35.rep



AeAeg_FcArc_GH_rep3.rep

