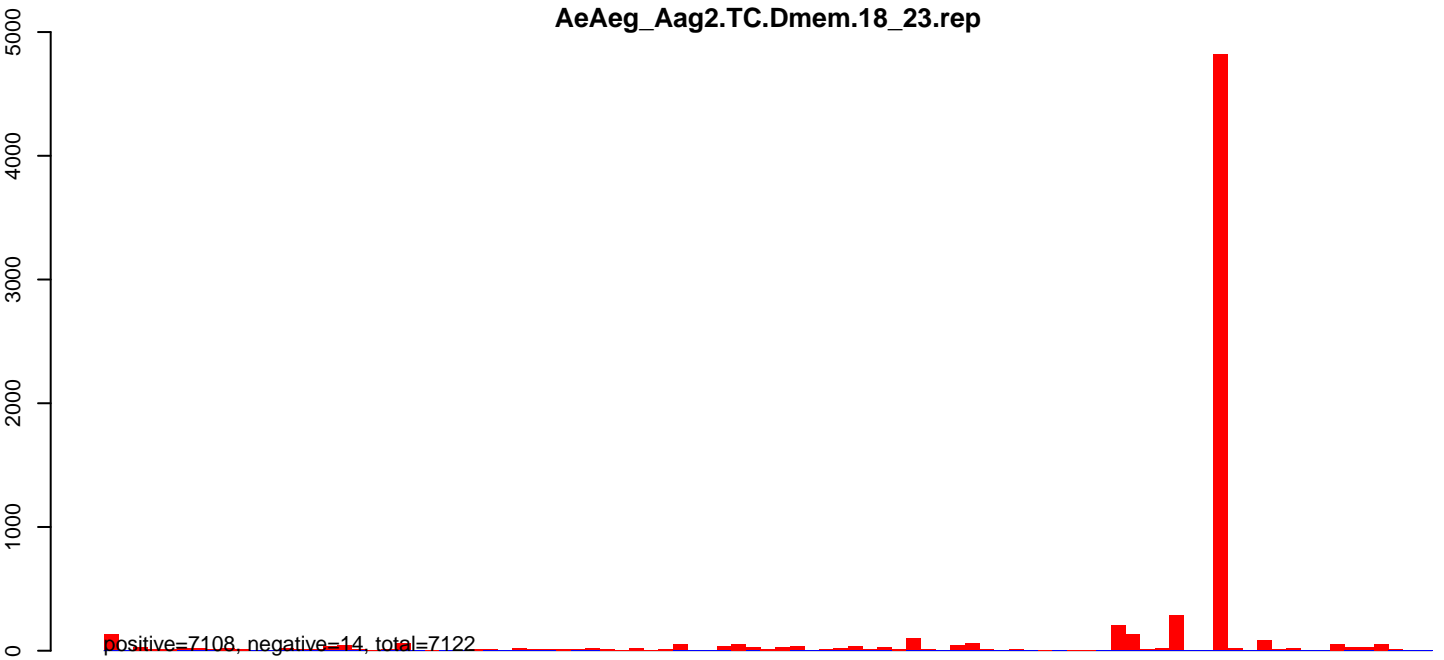
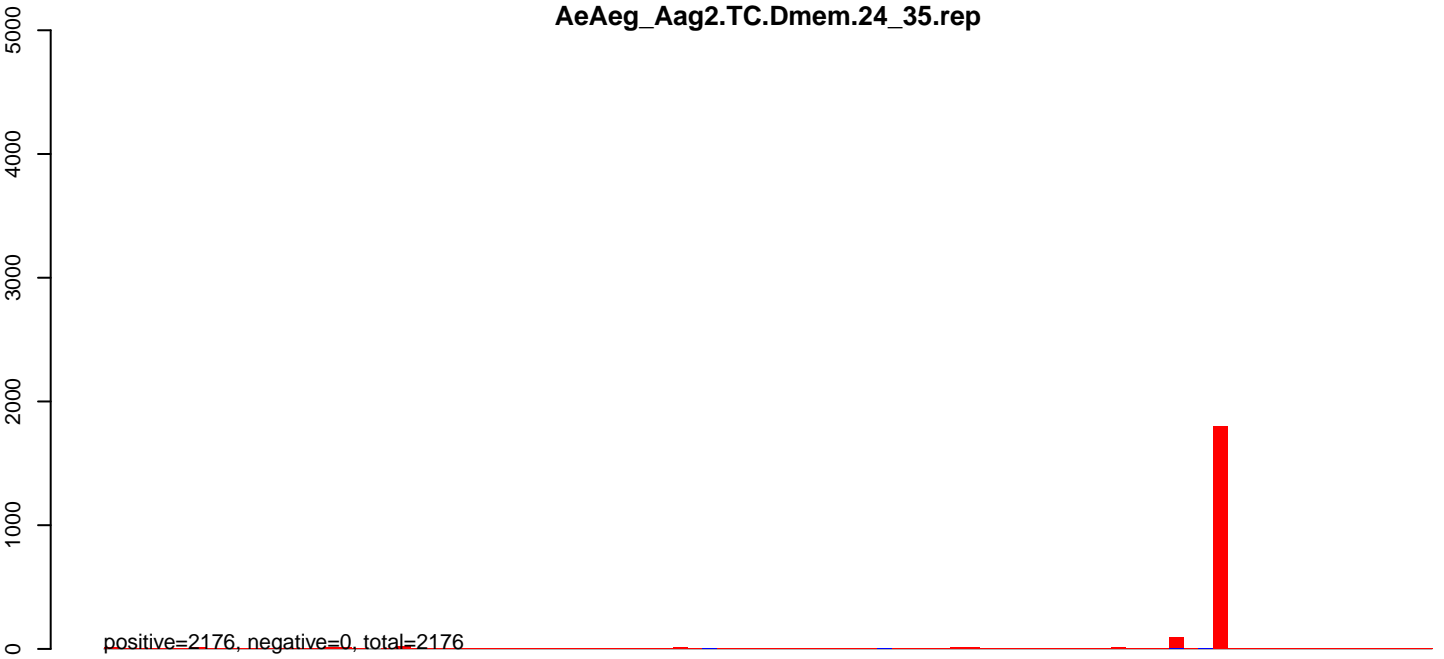


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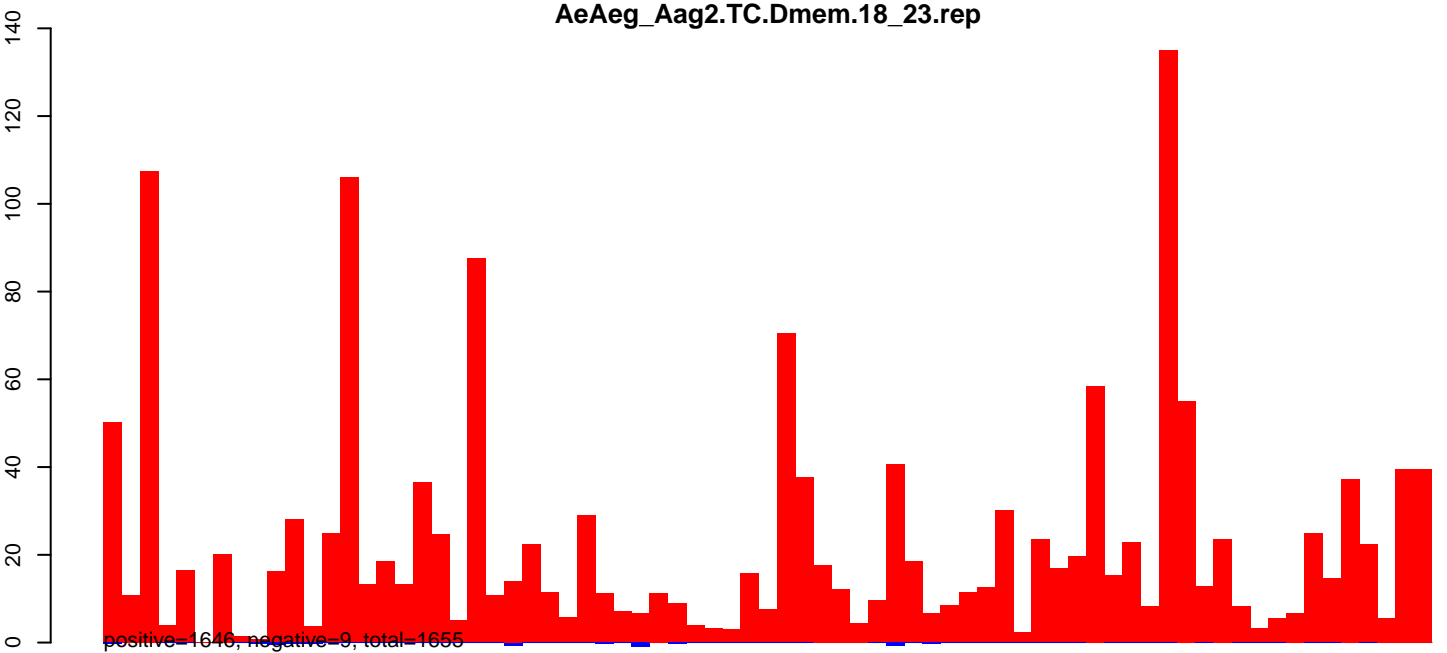
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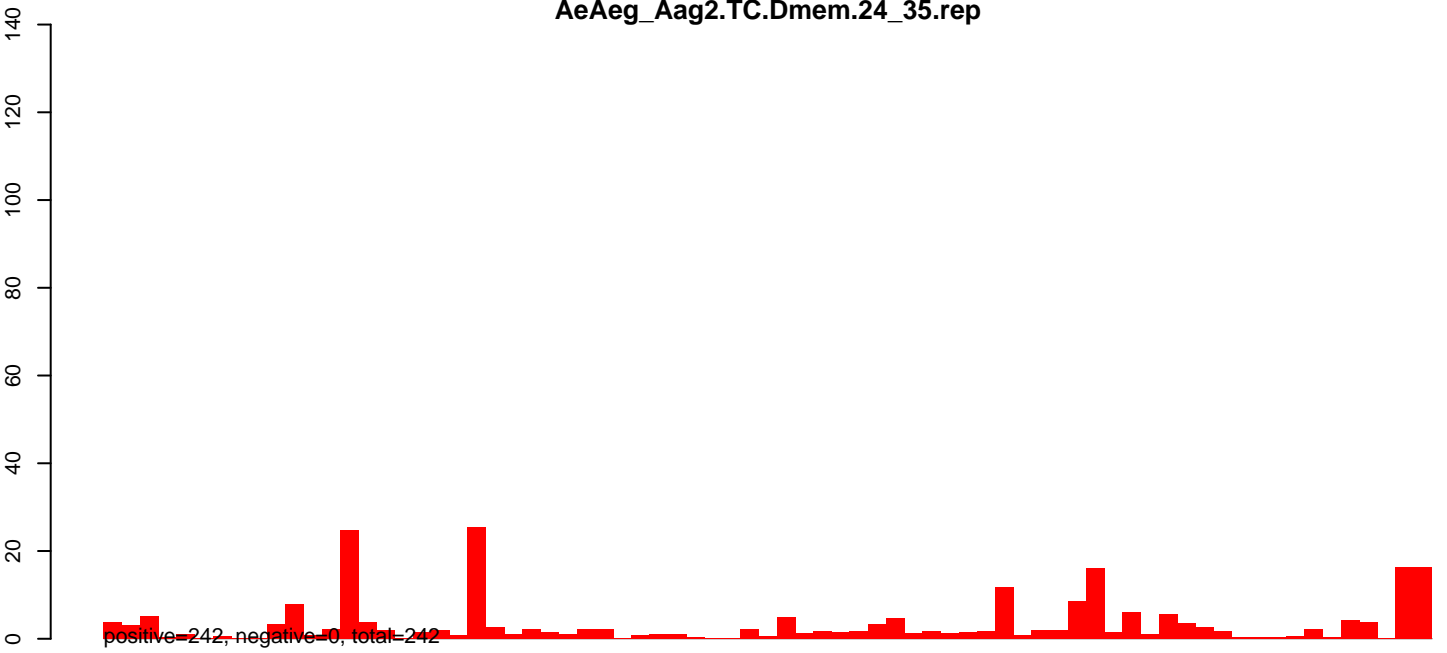
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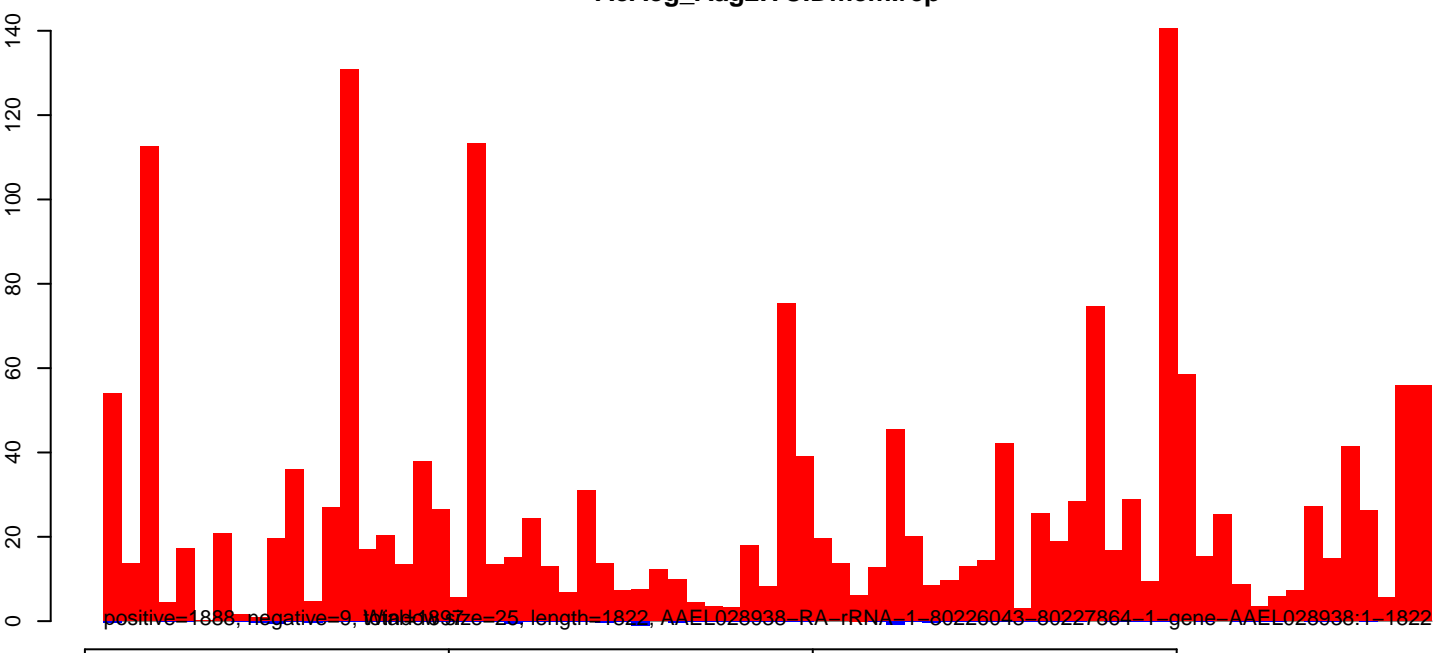
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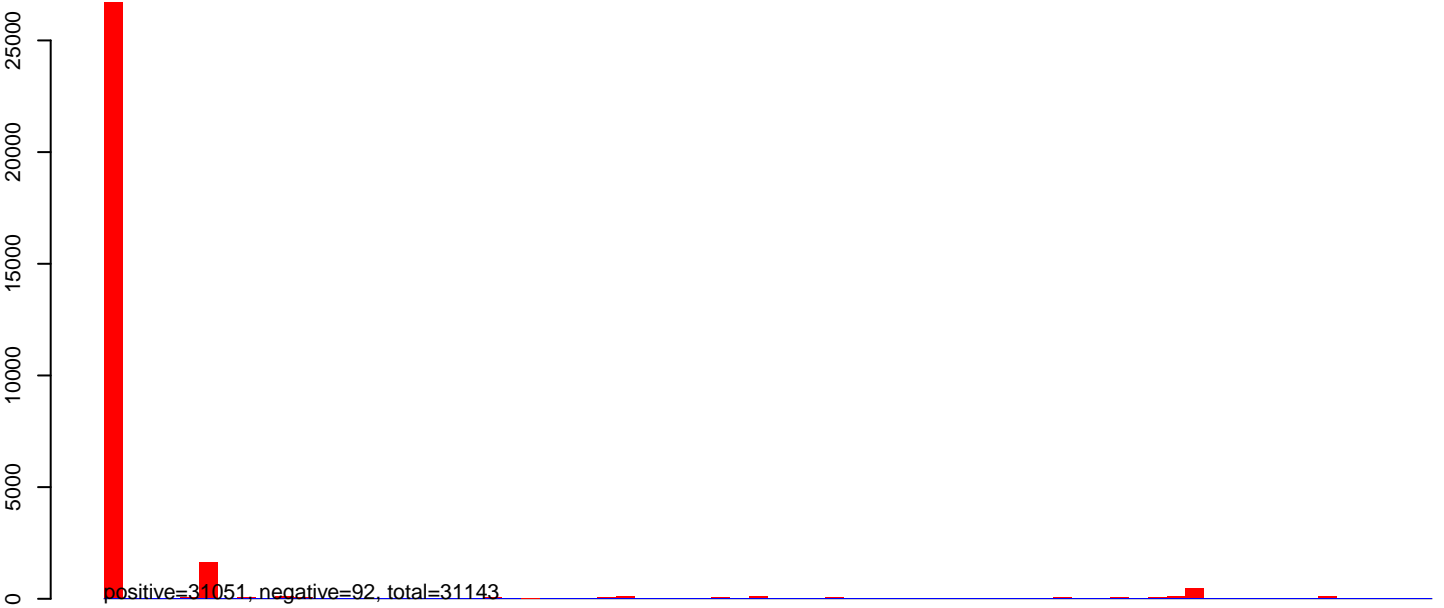
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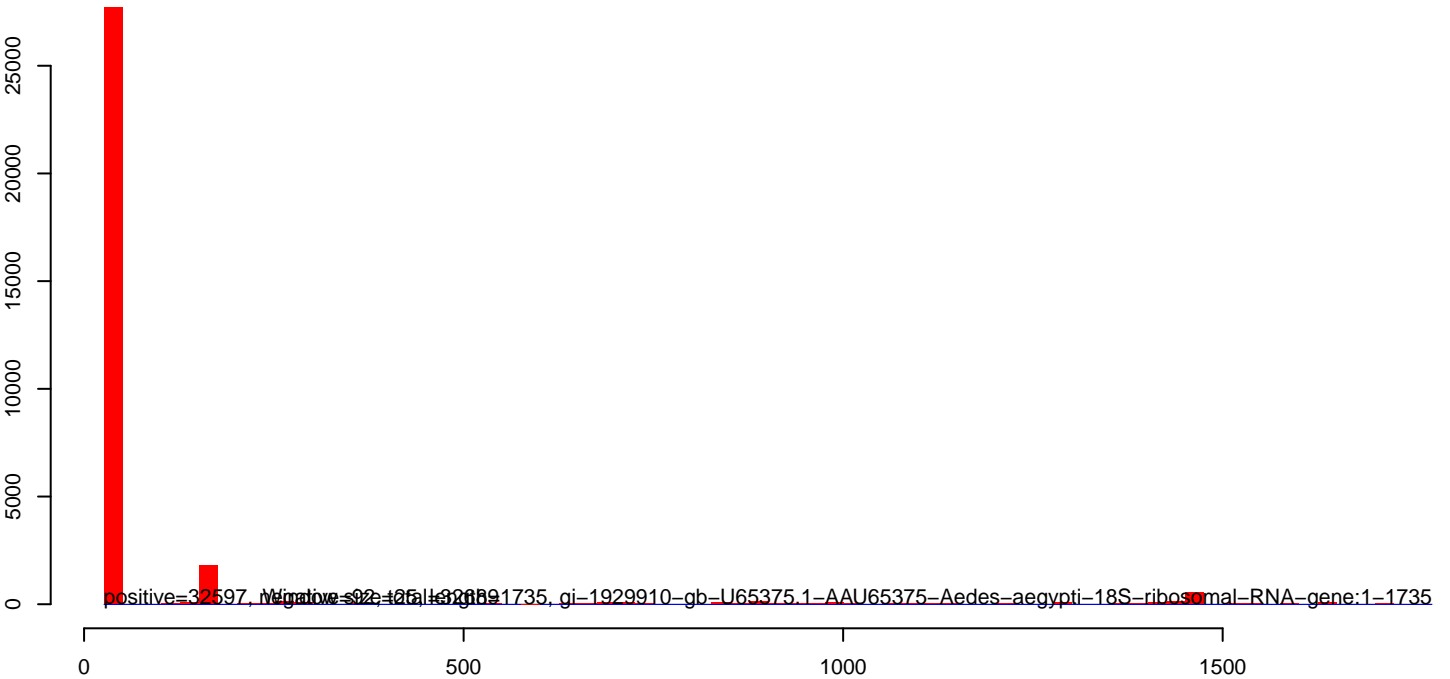
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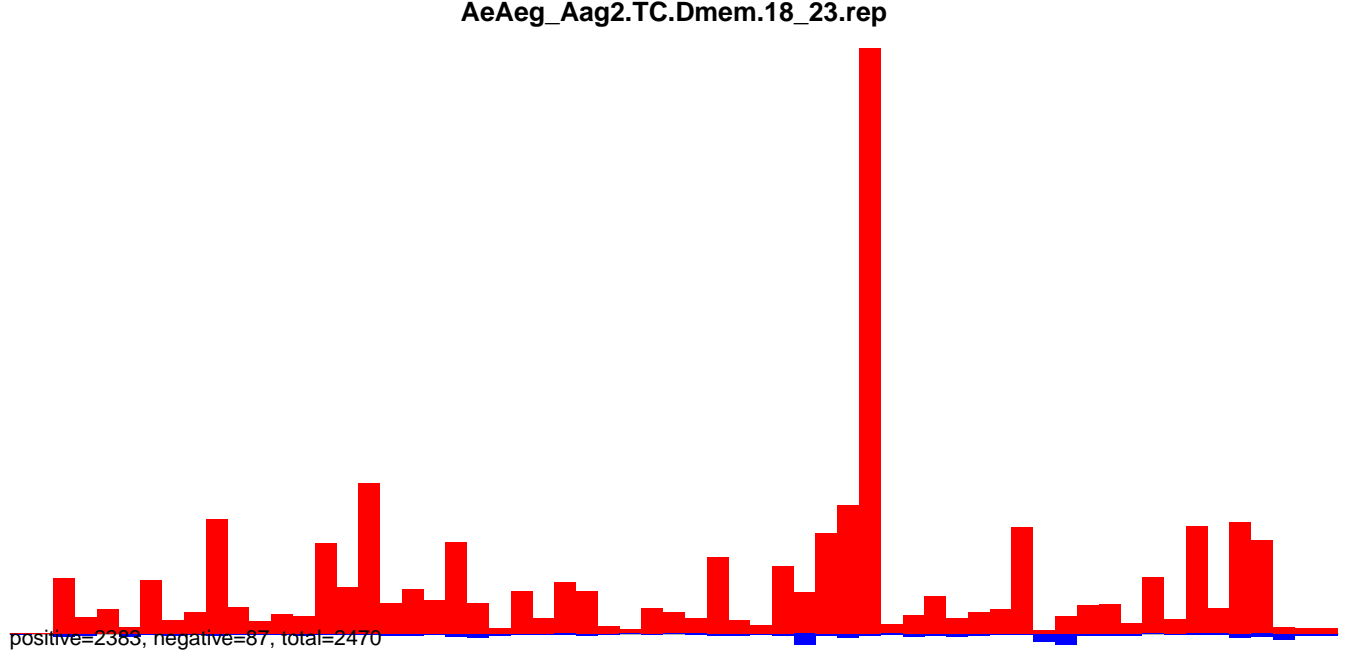
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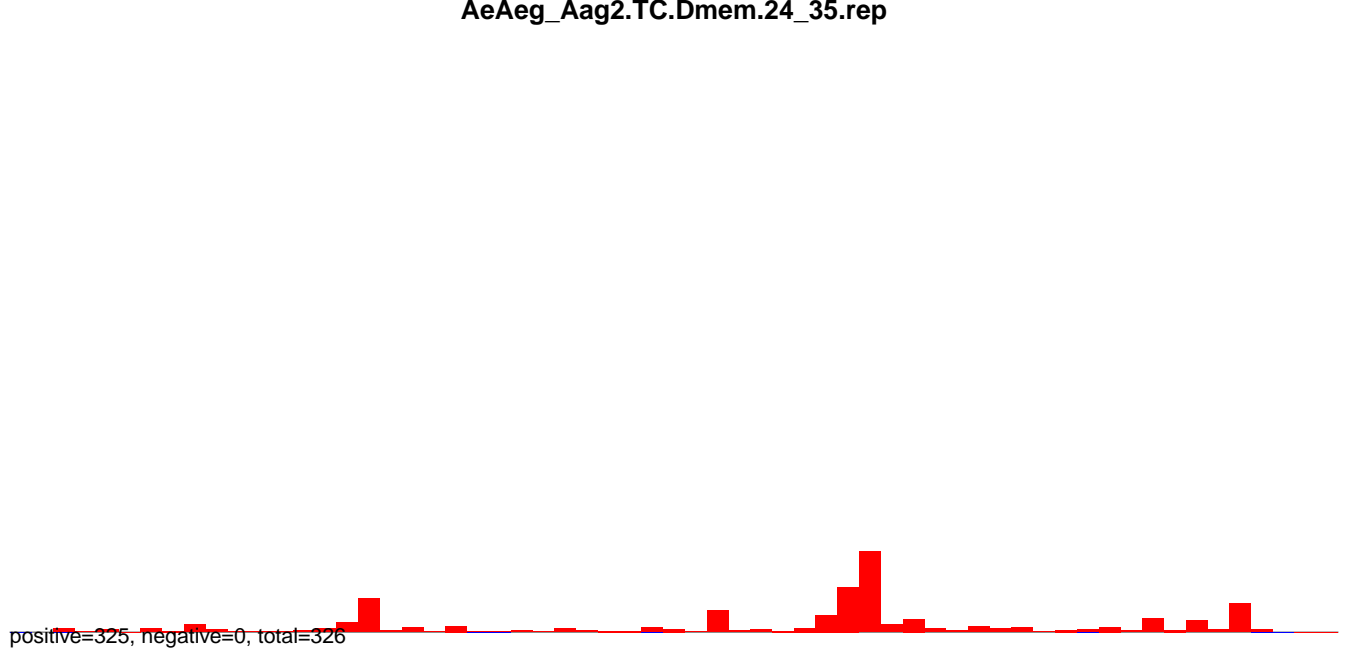
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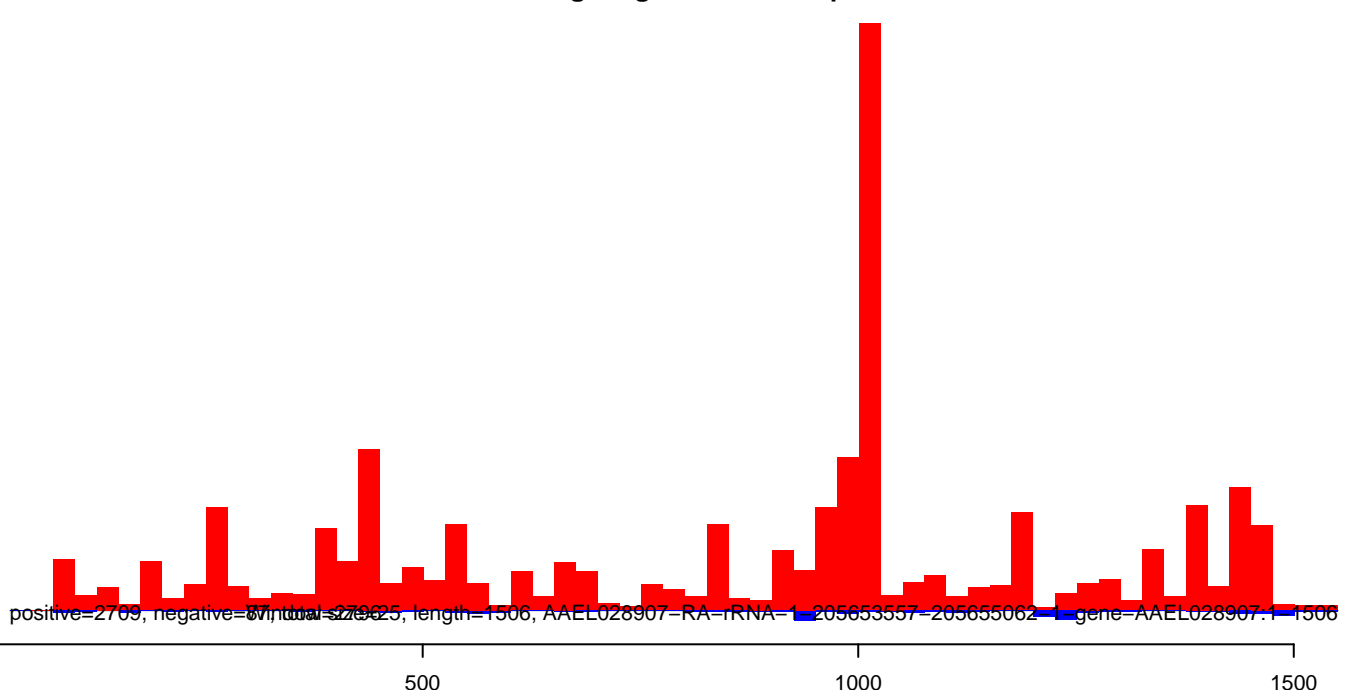
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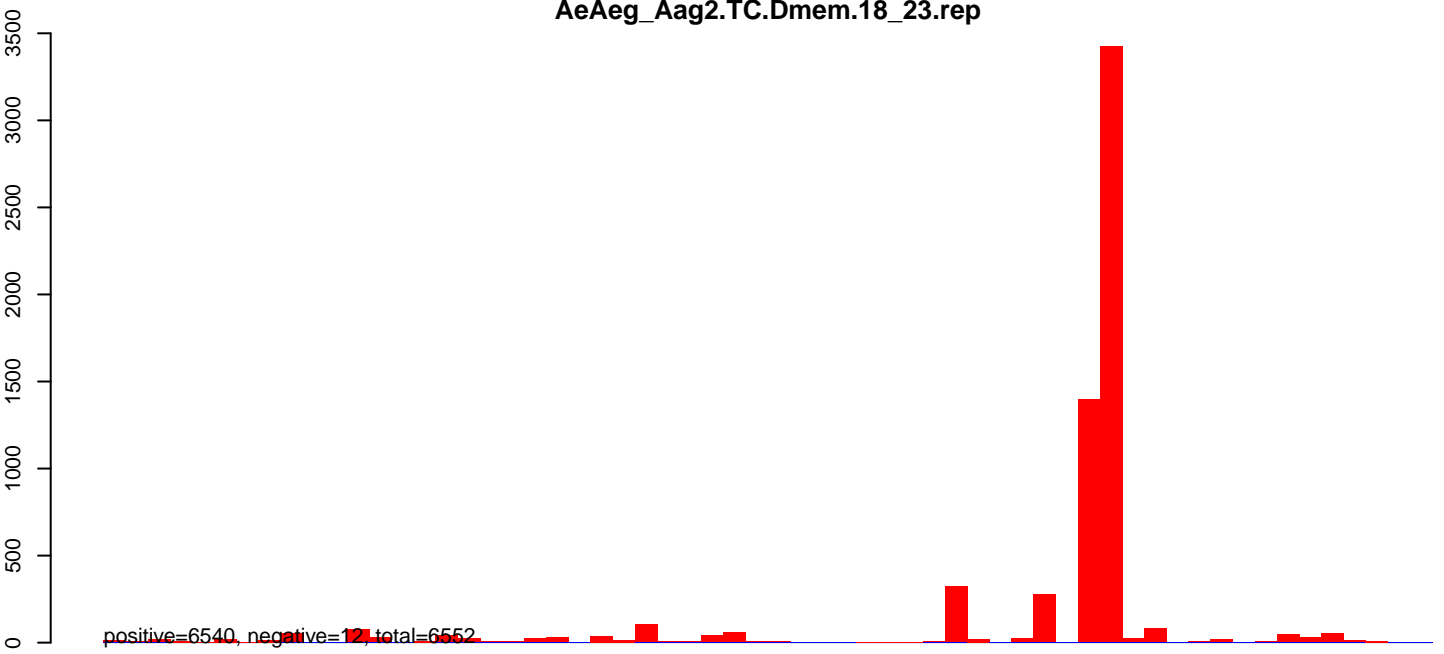
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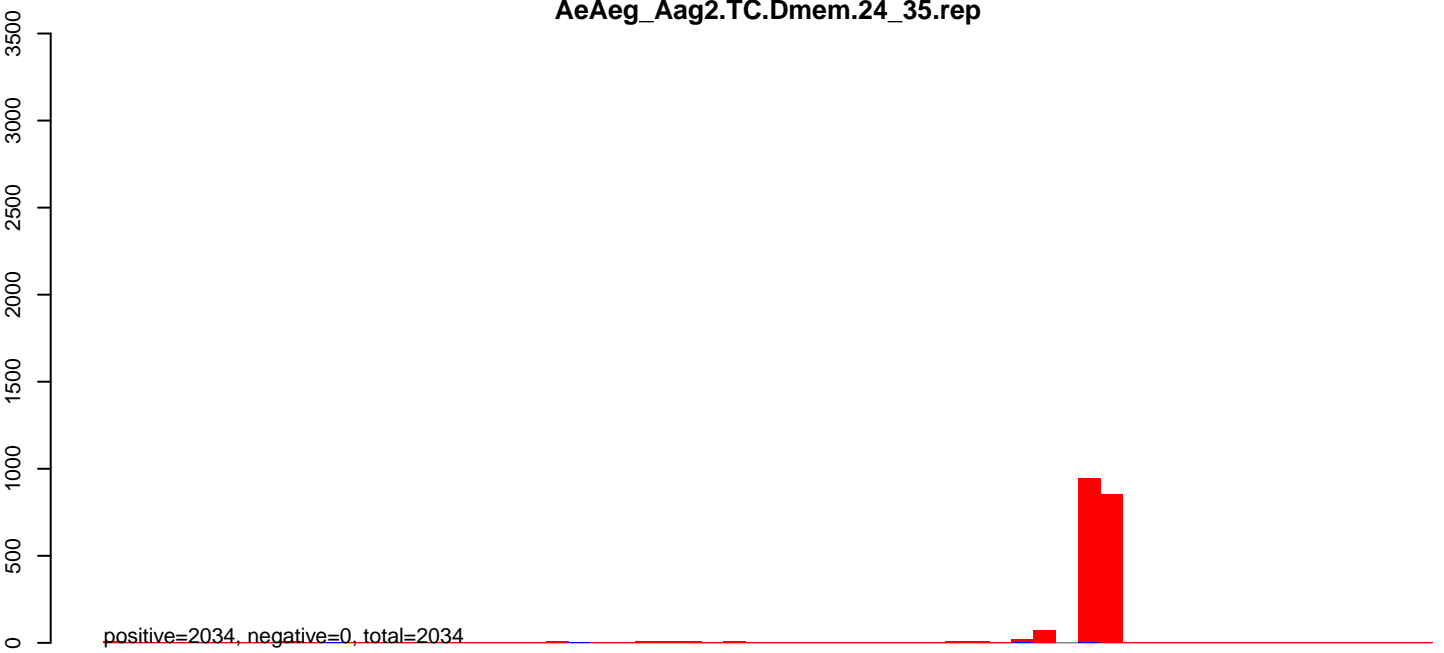
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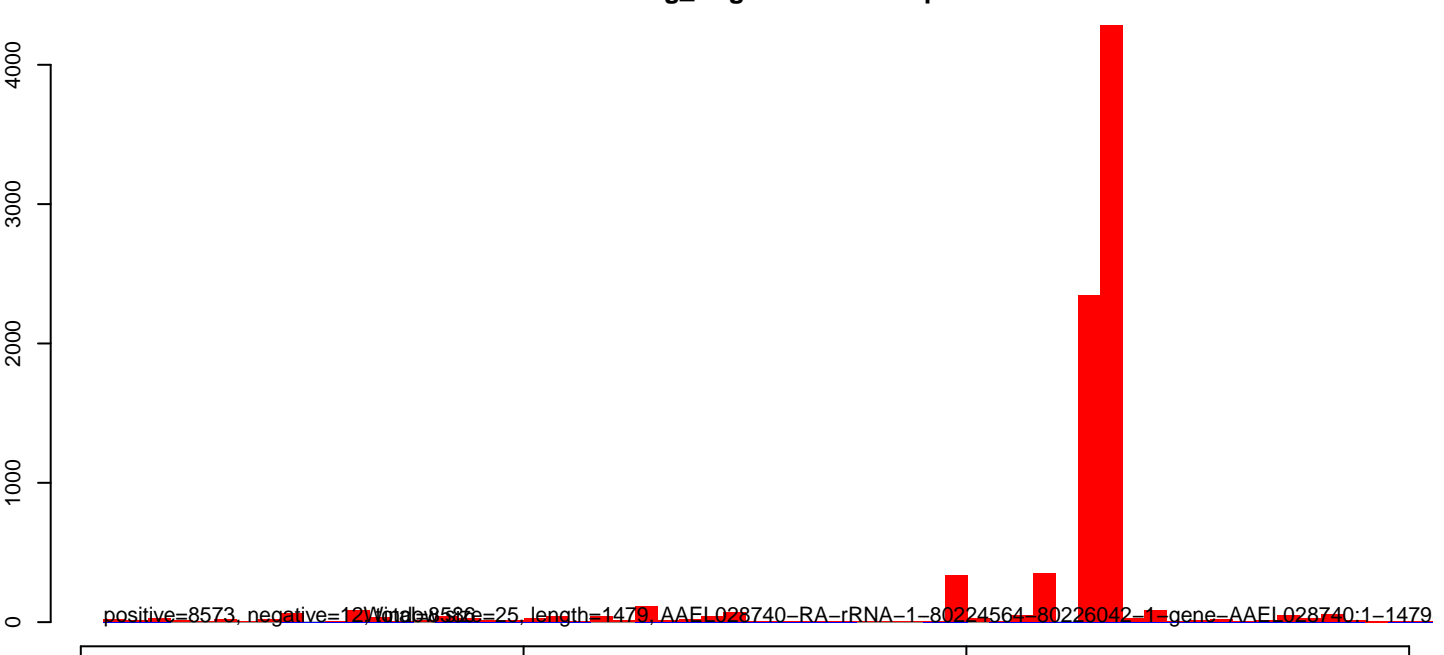
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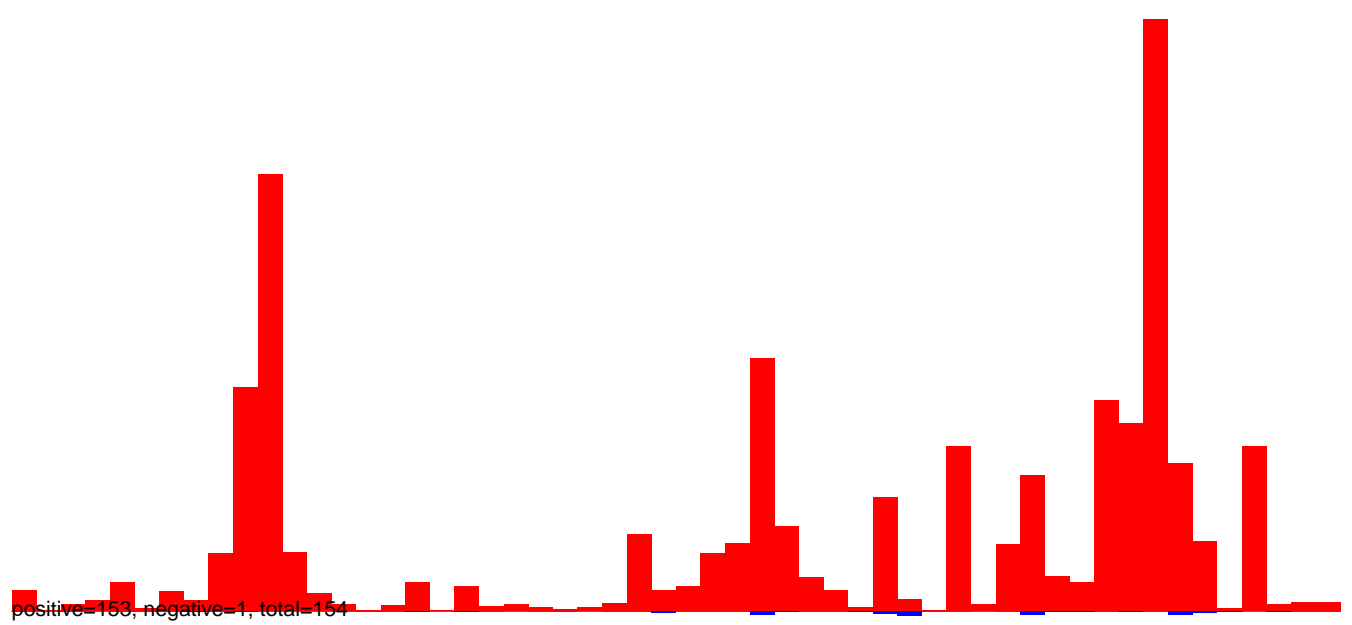
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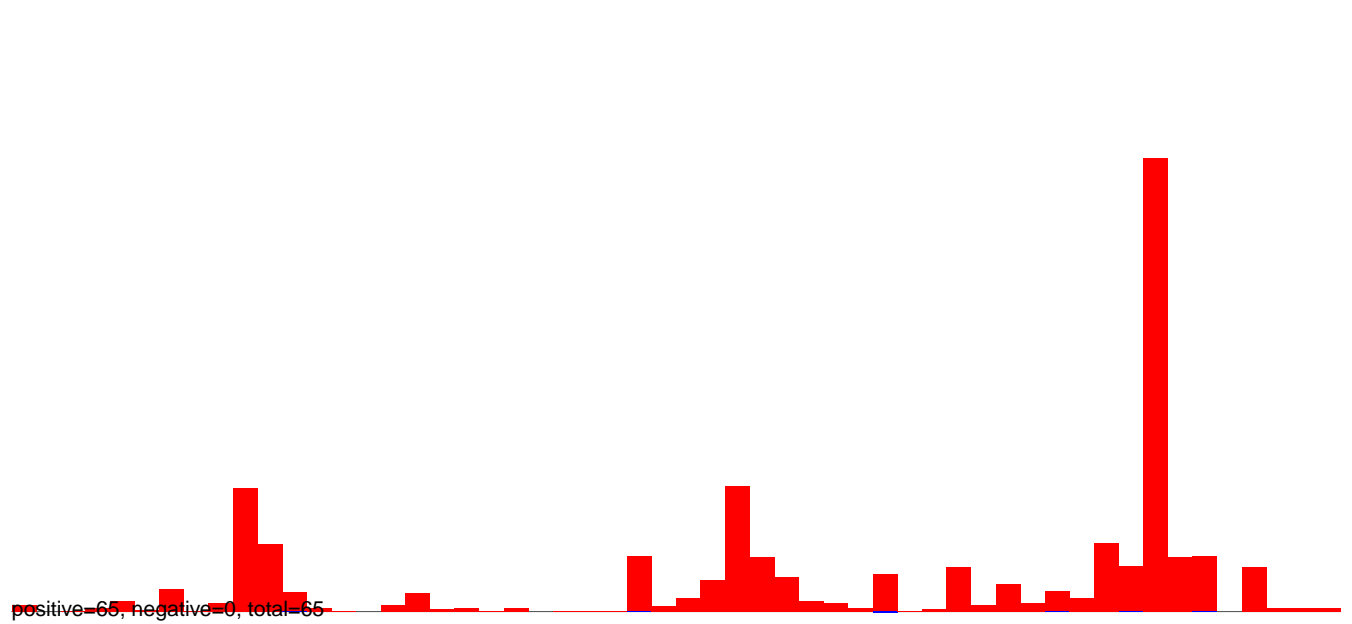
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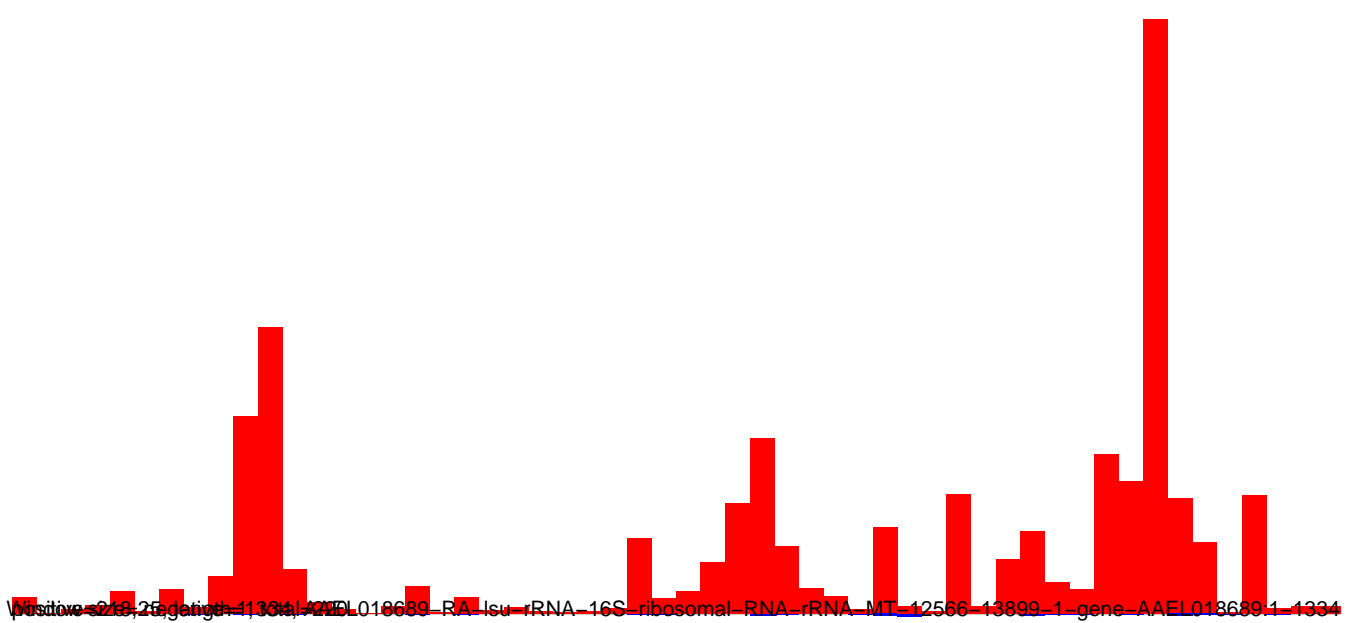
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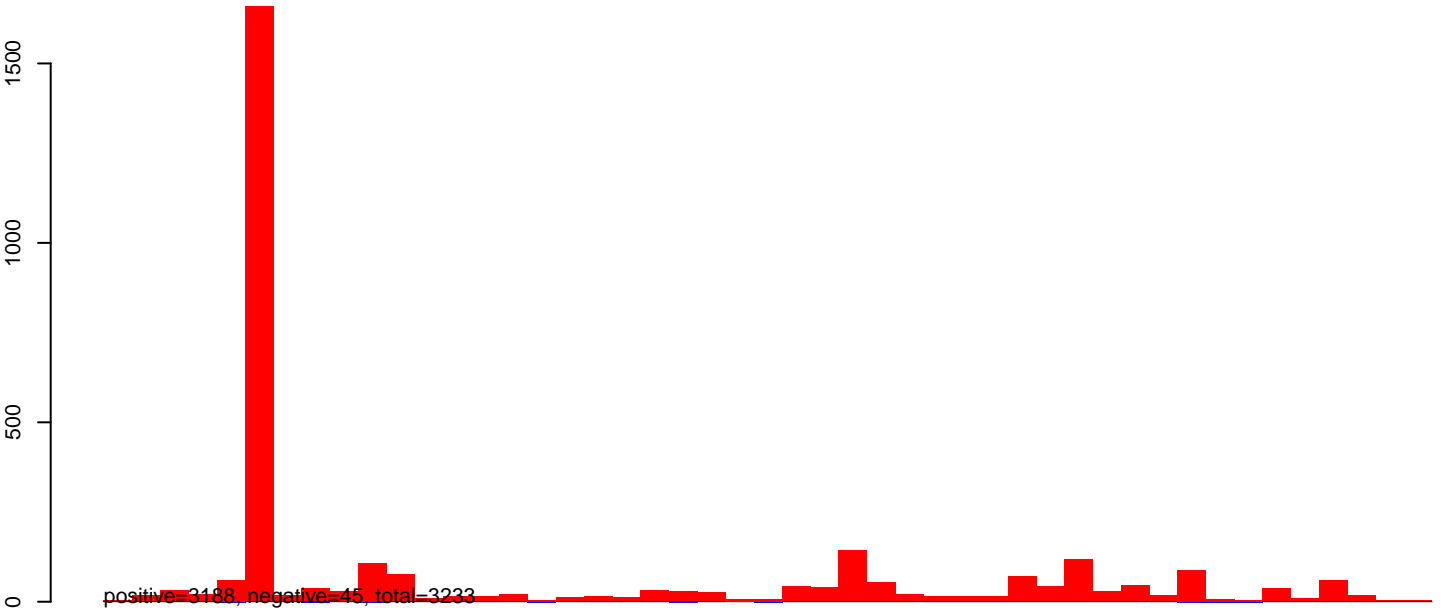
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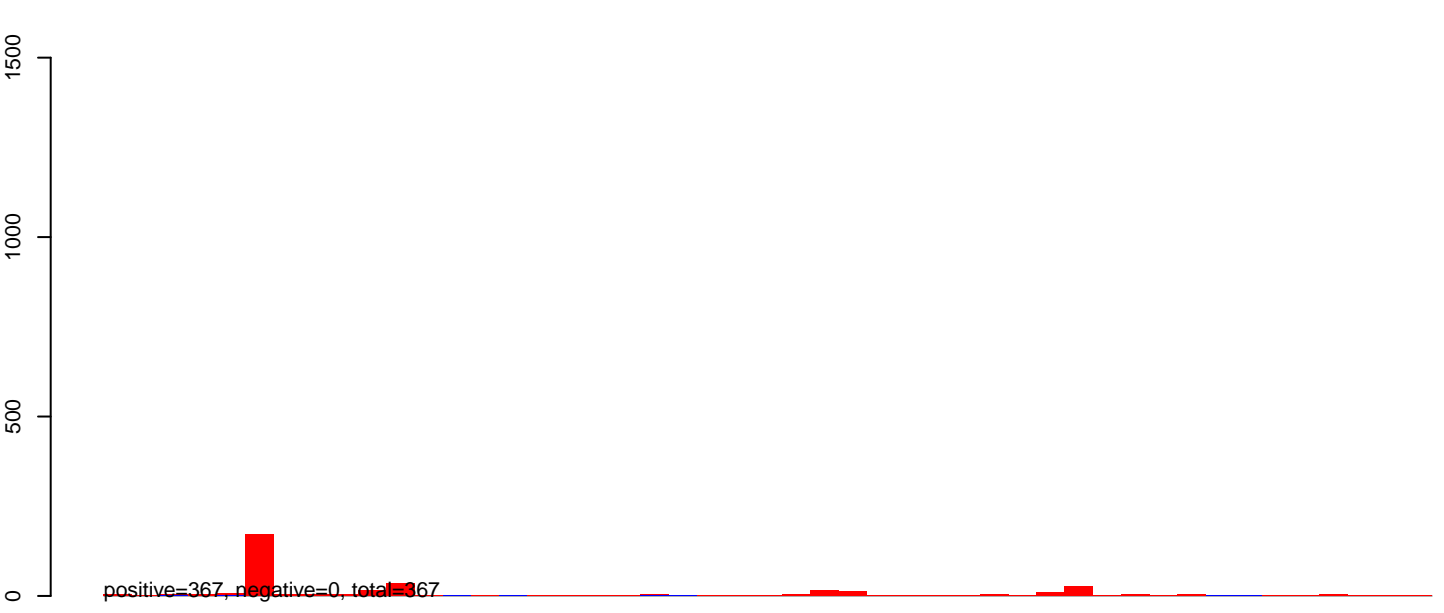
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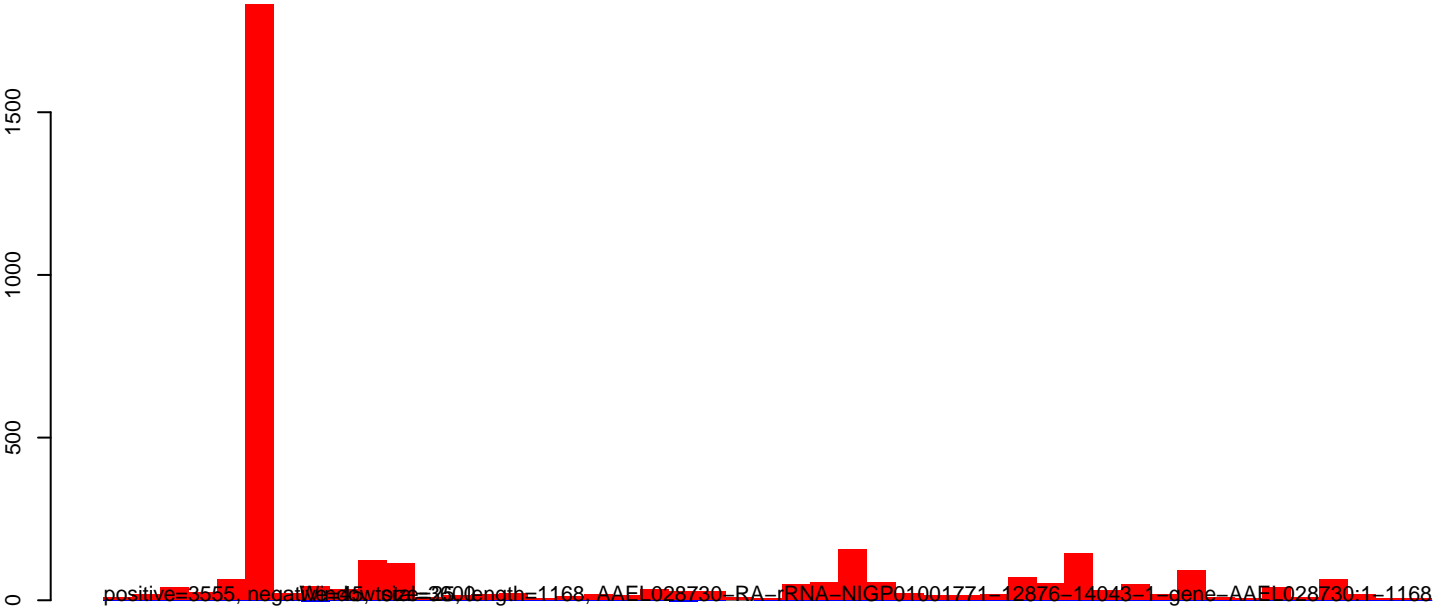
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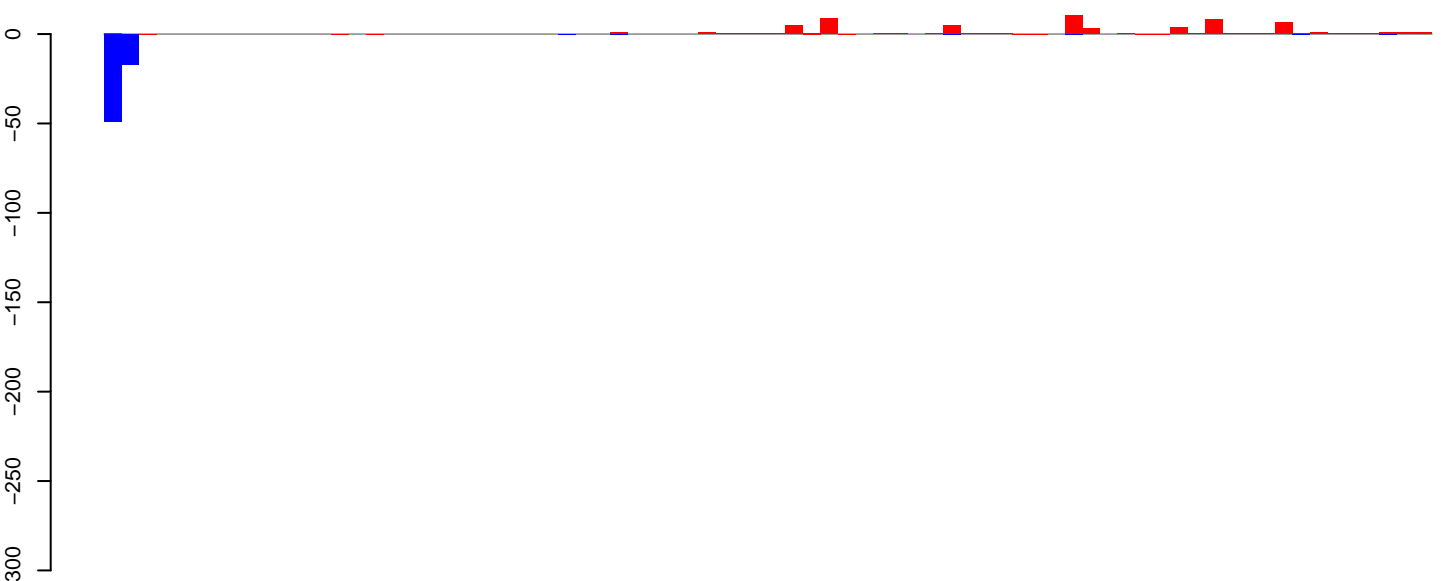


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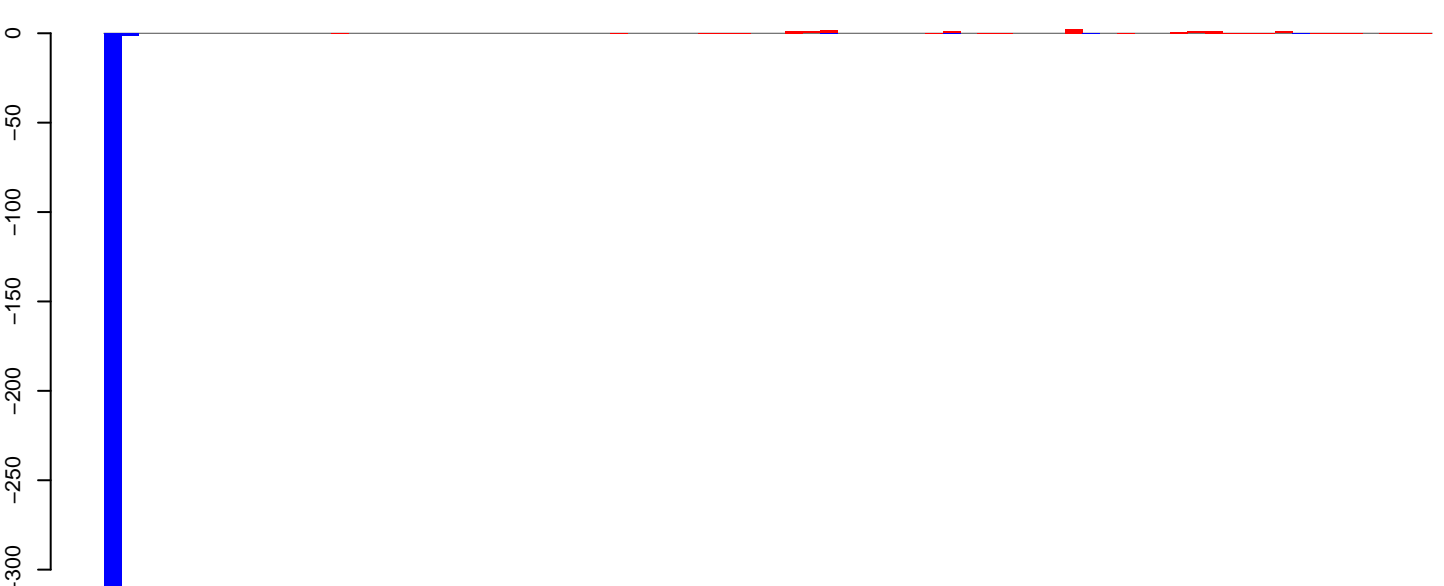
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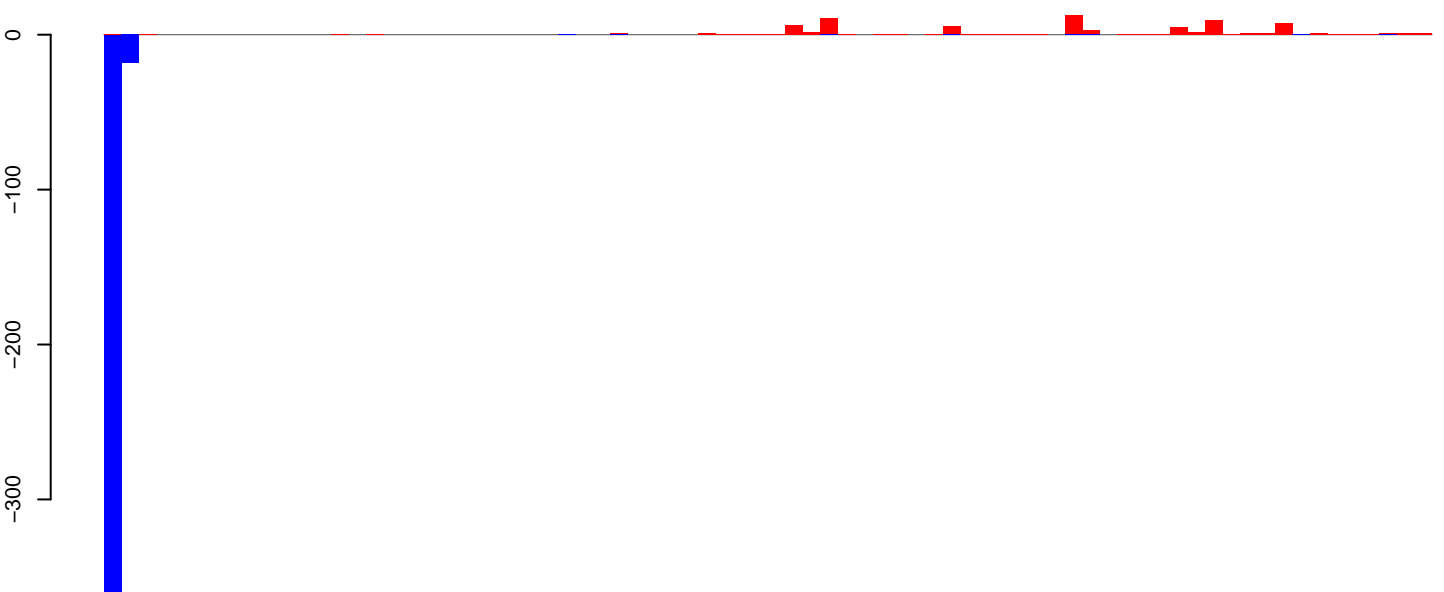
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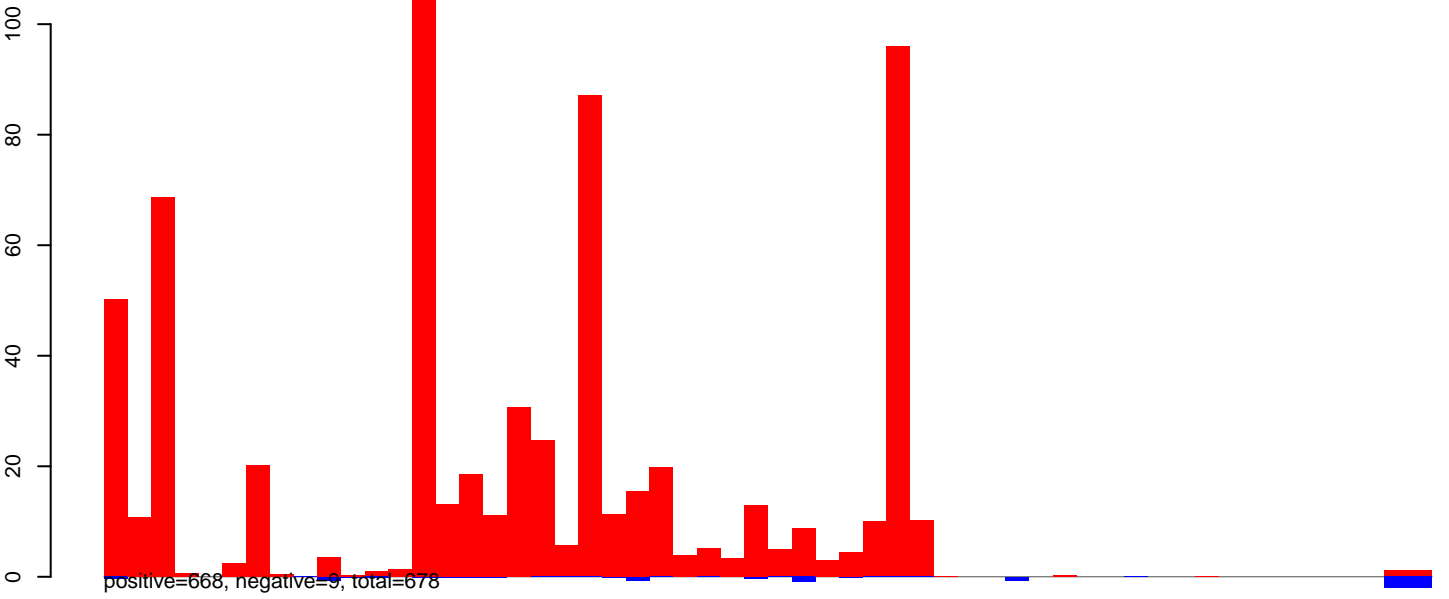
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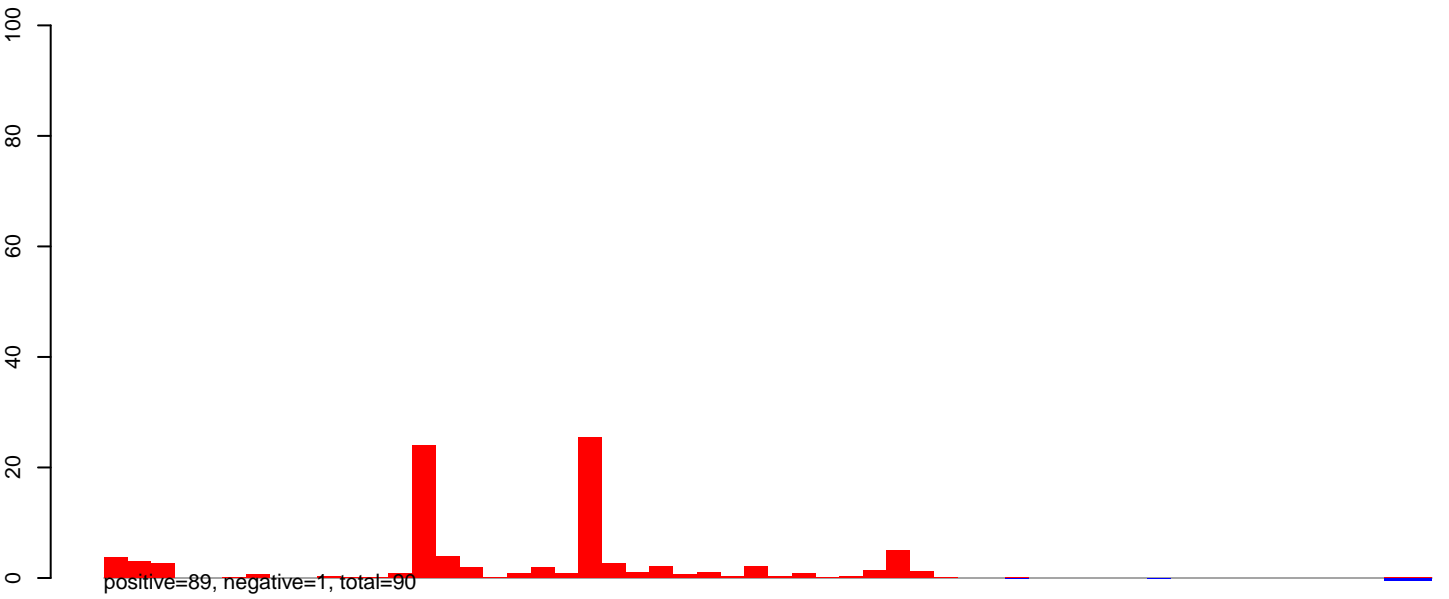


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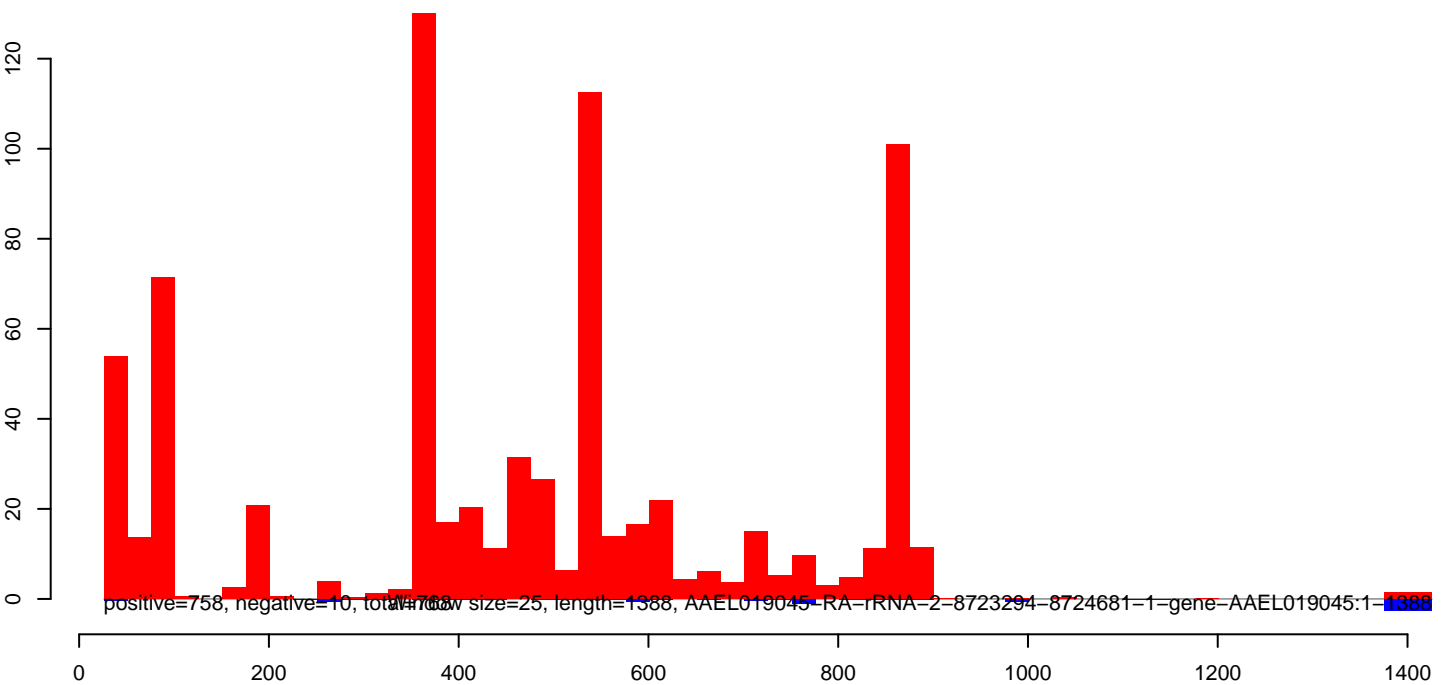
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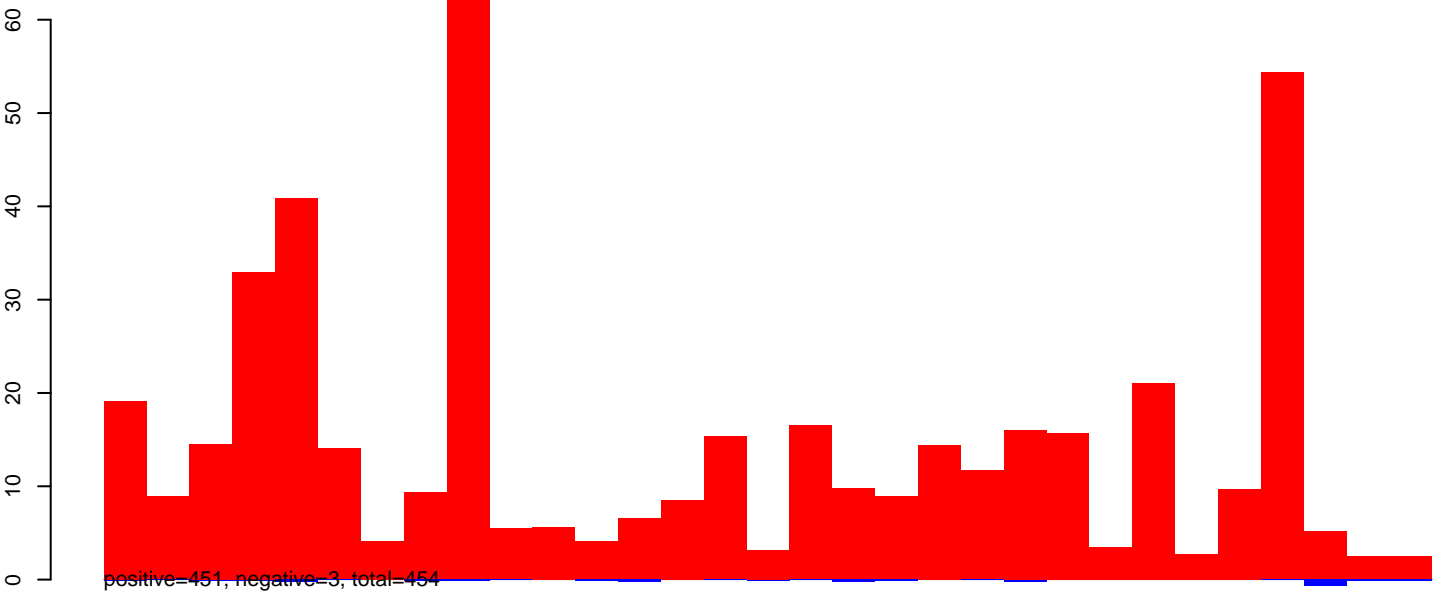
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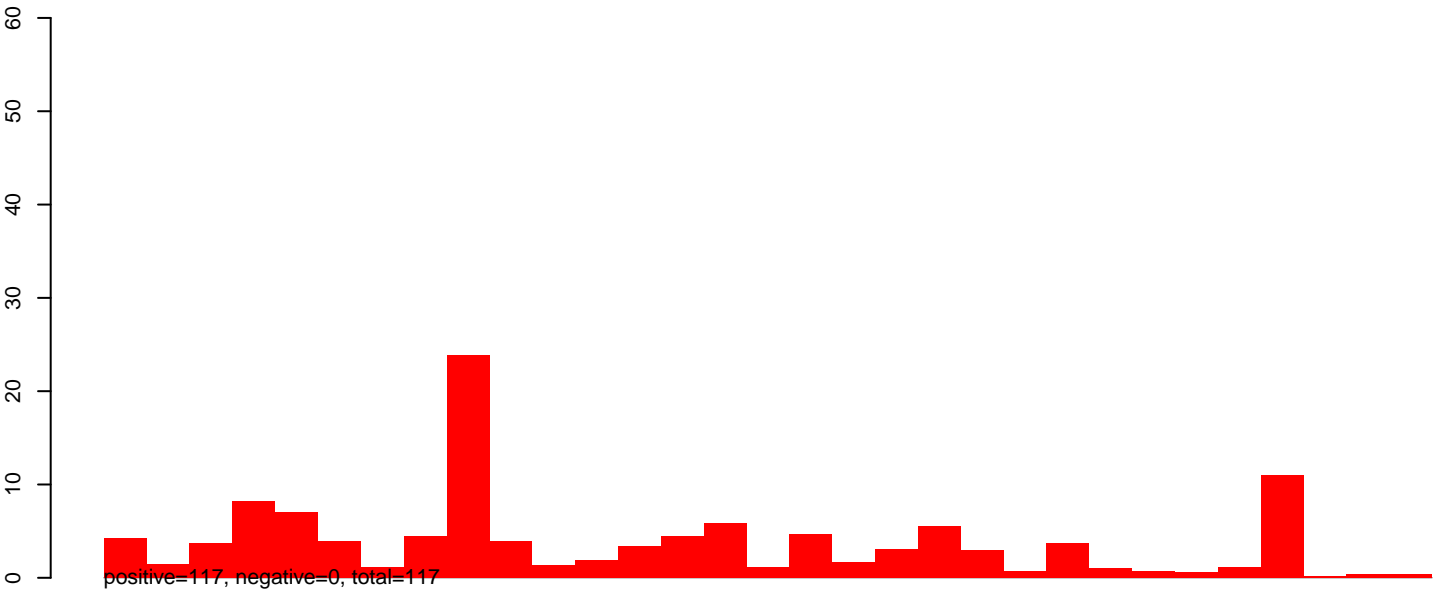
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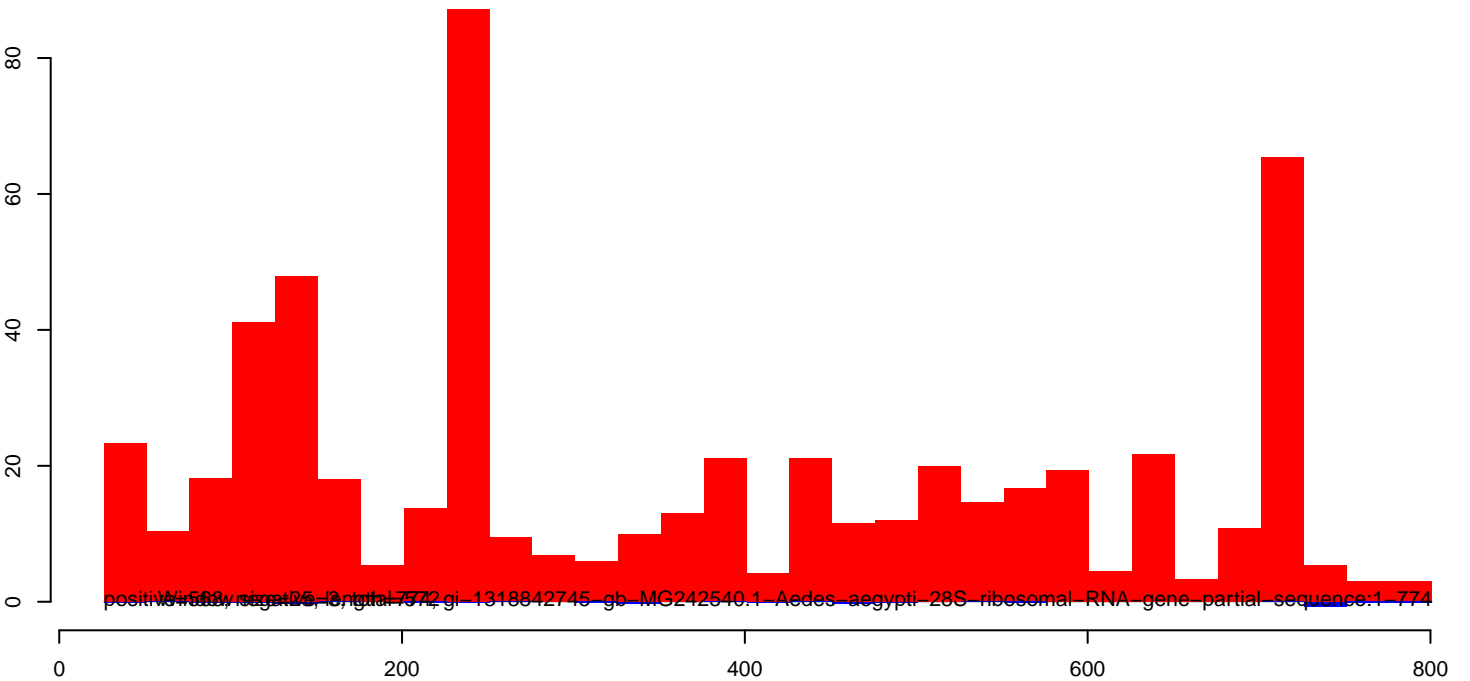
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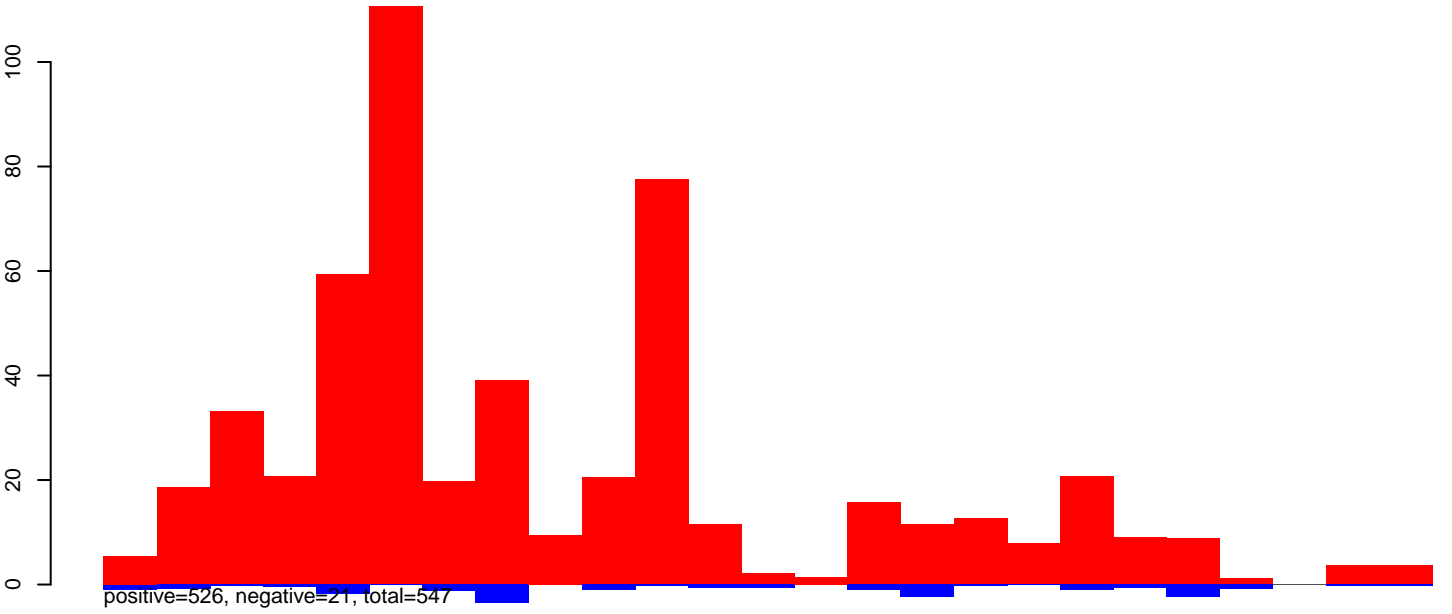
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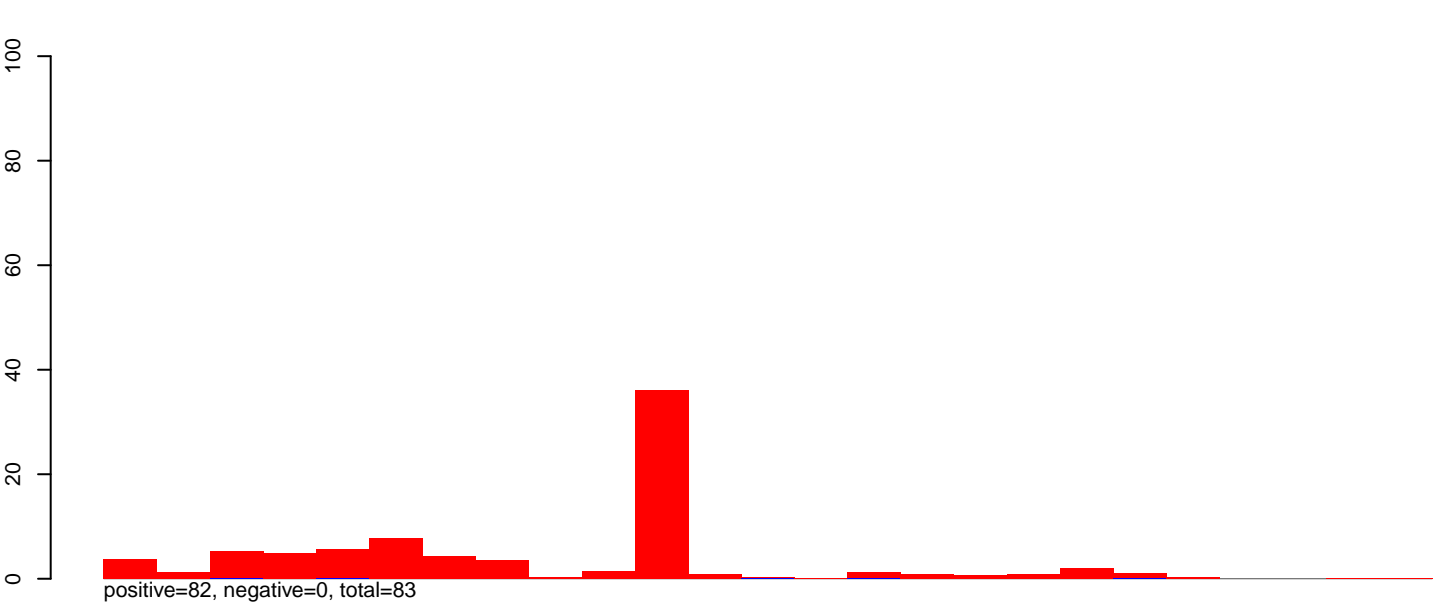
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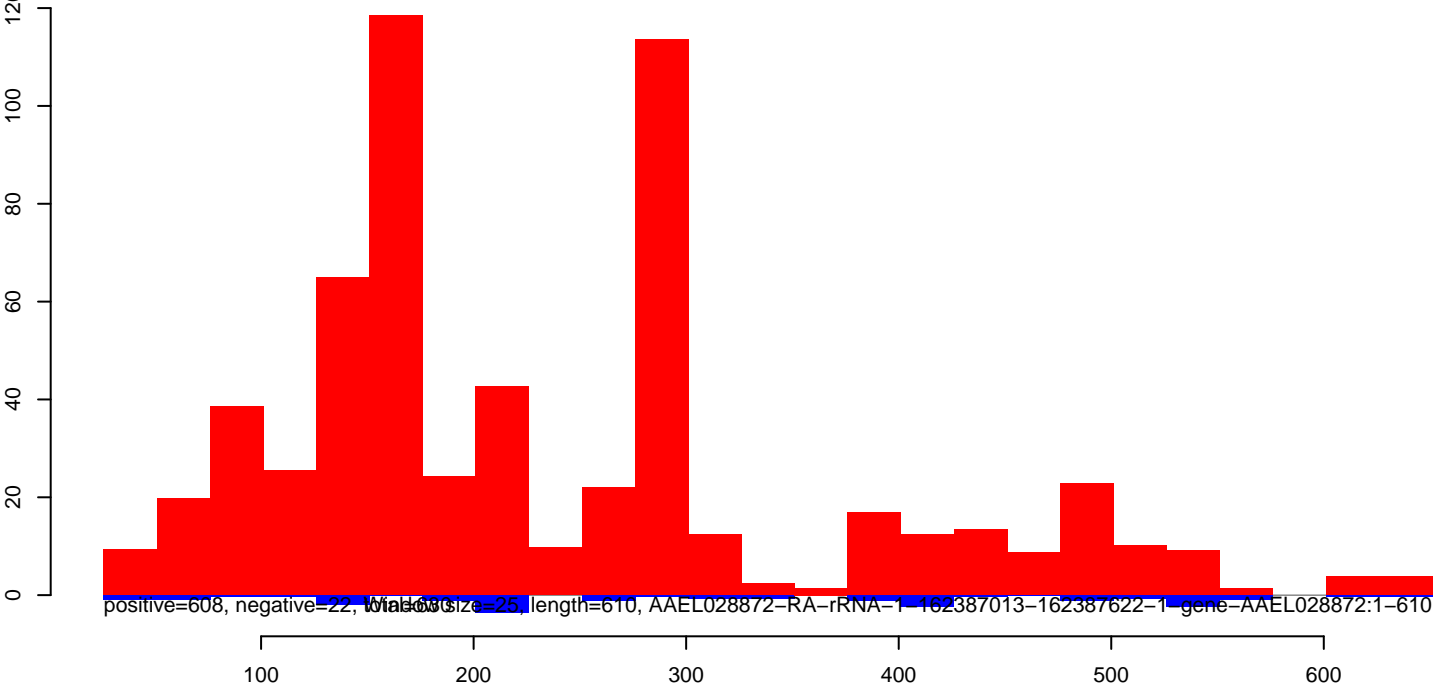
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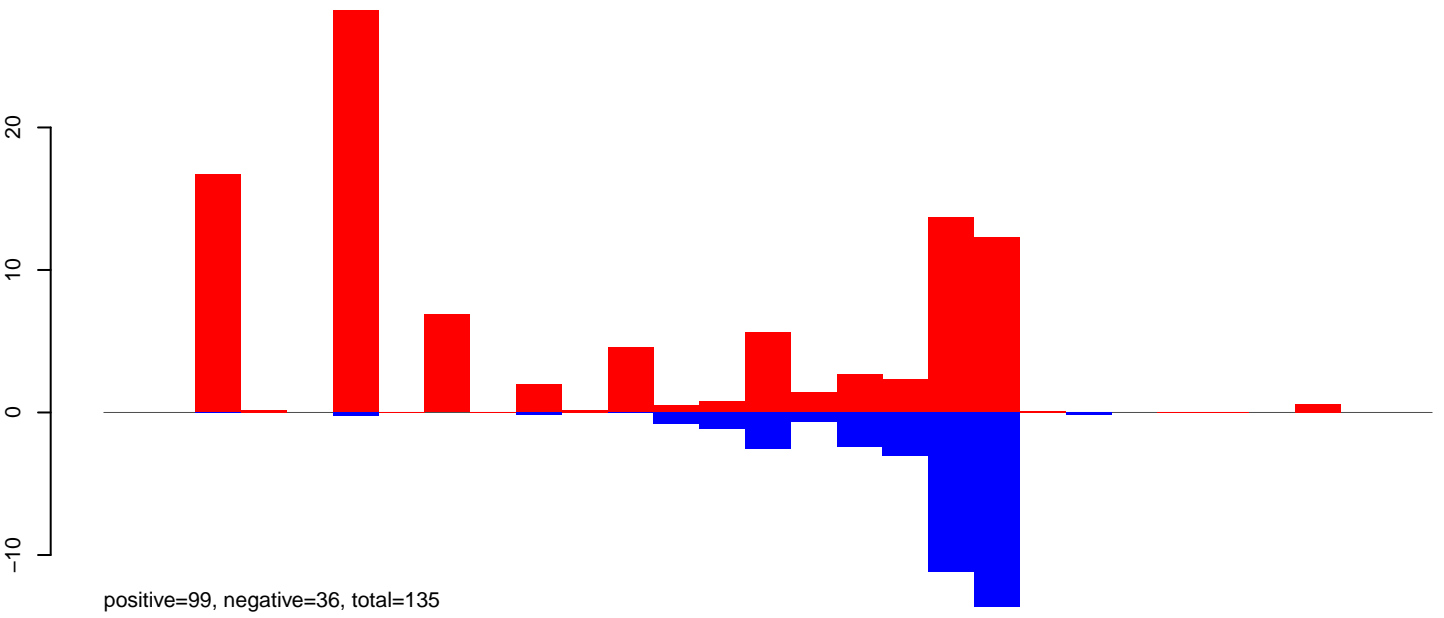
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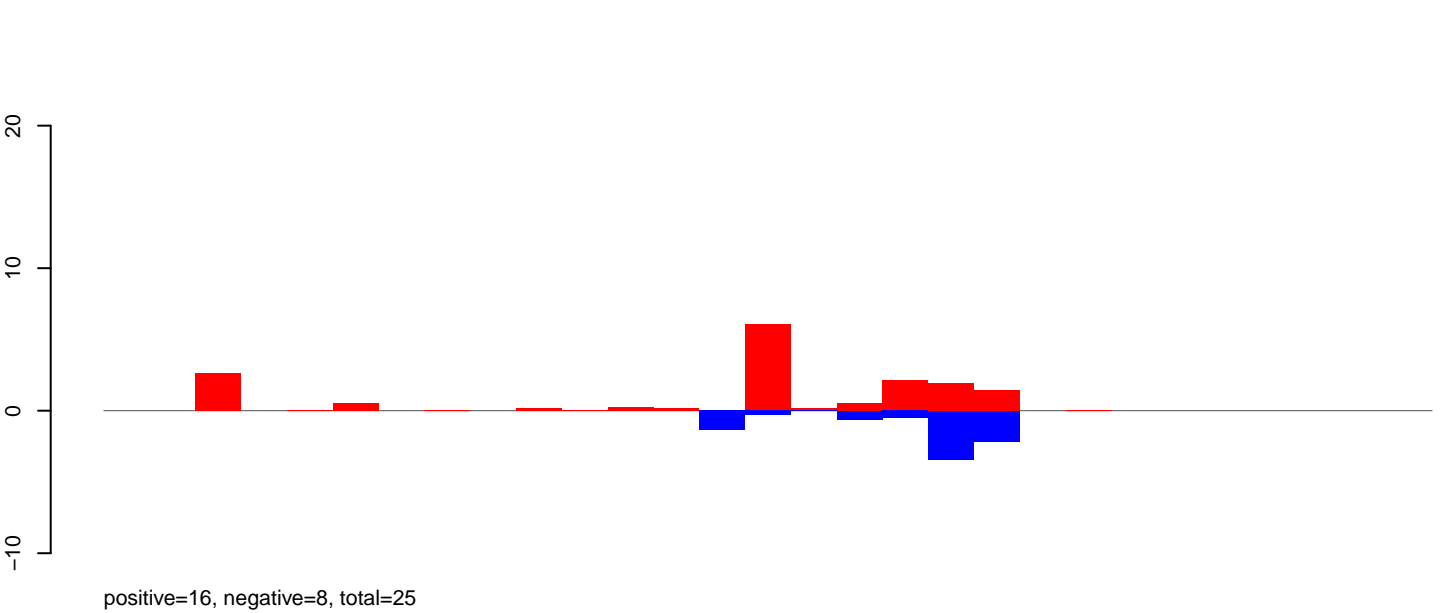
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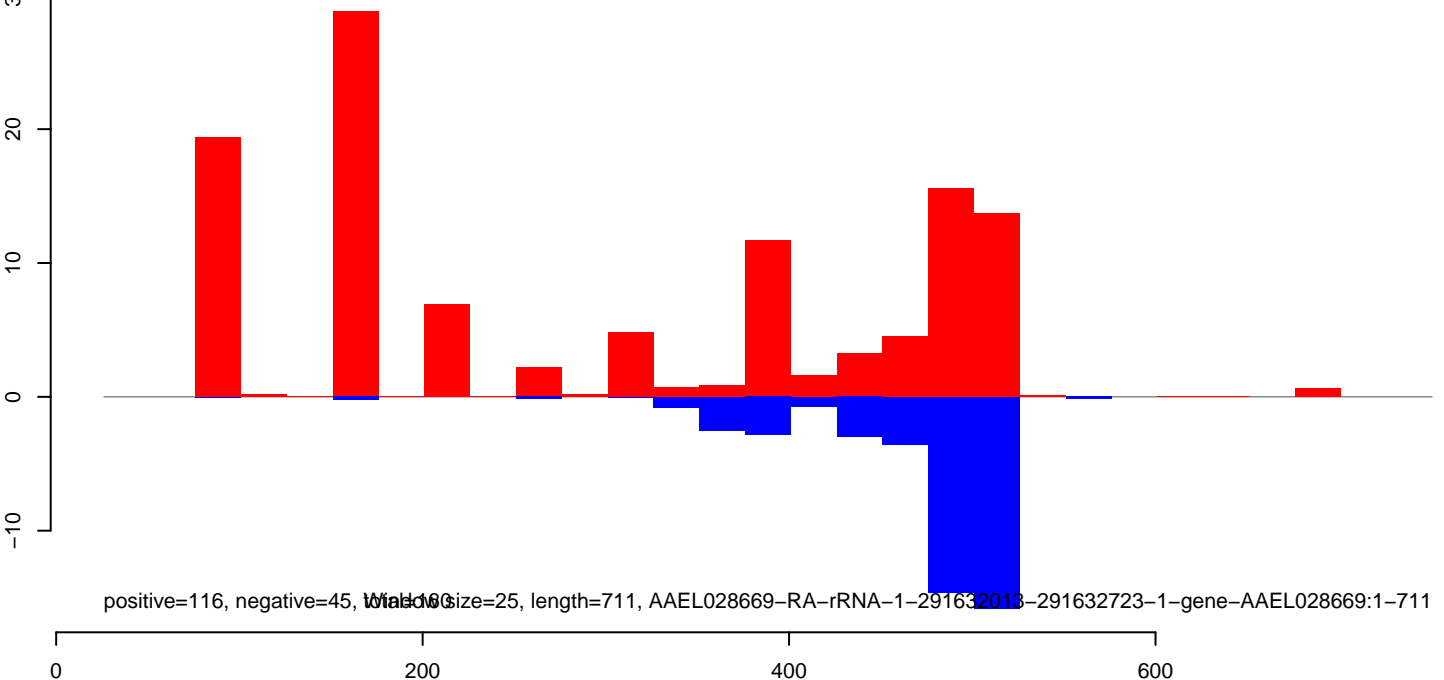
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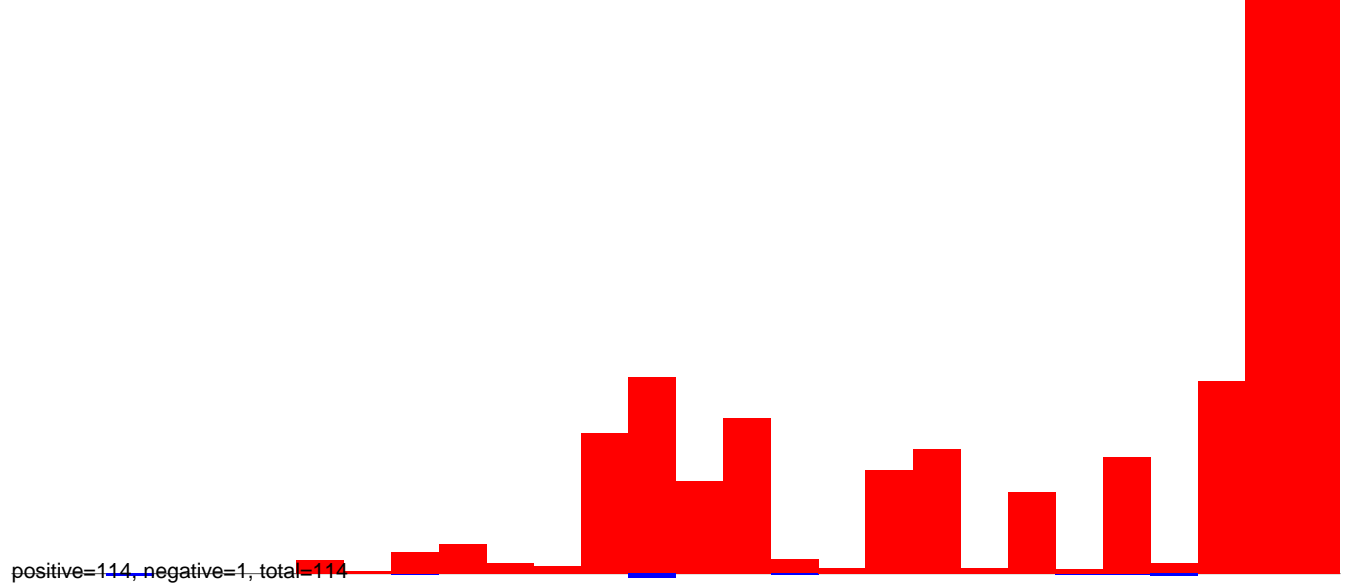
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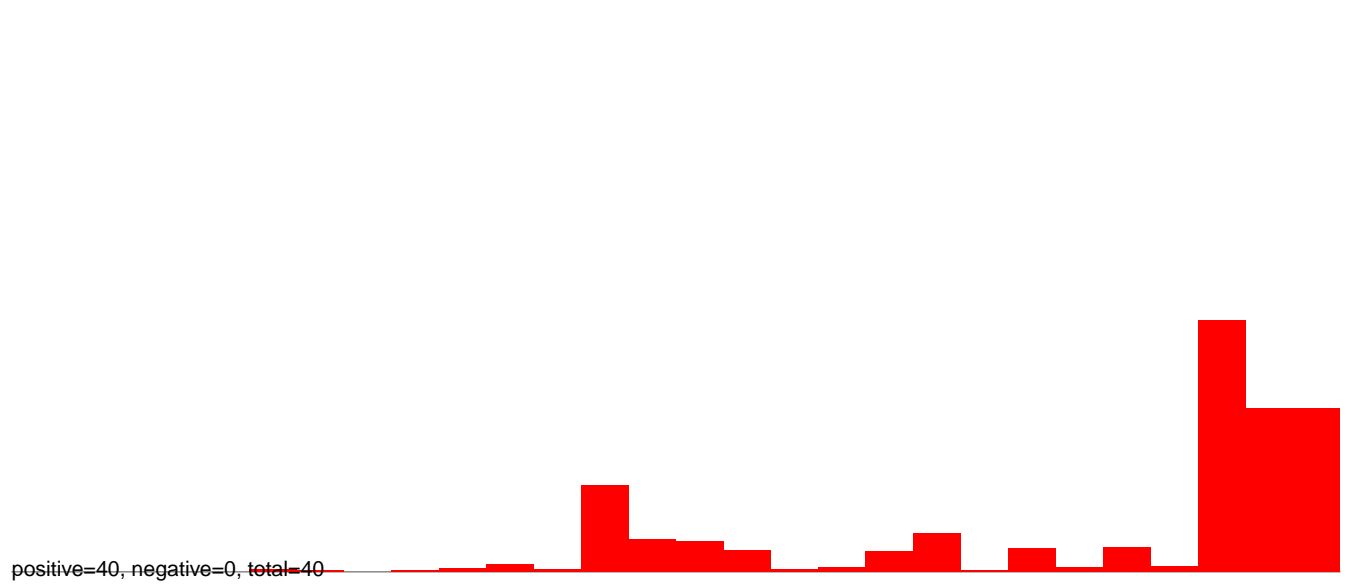
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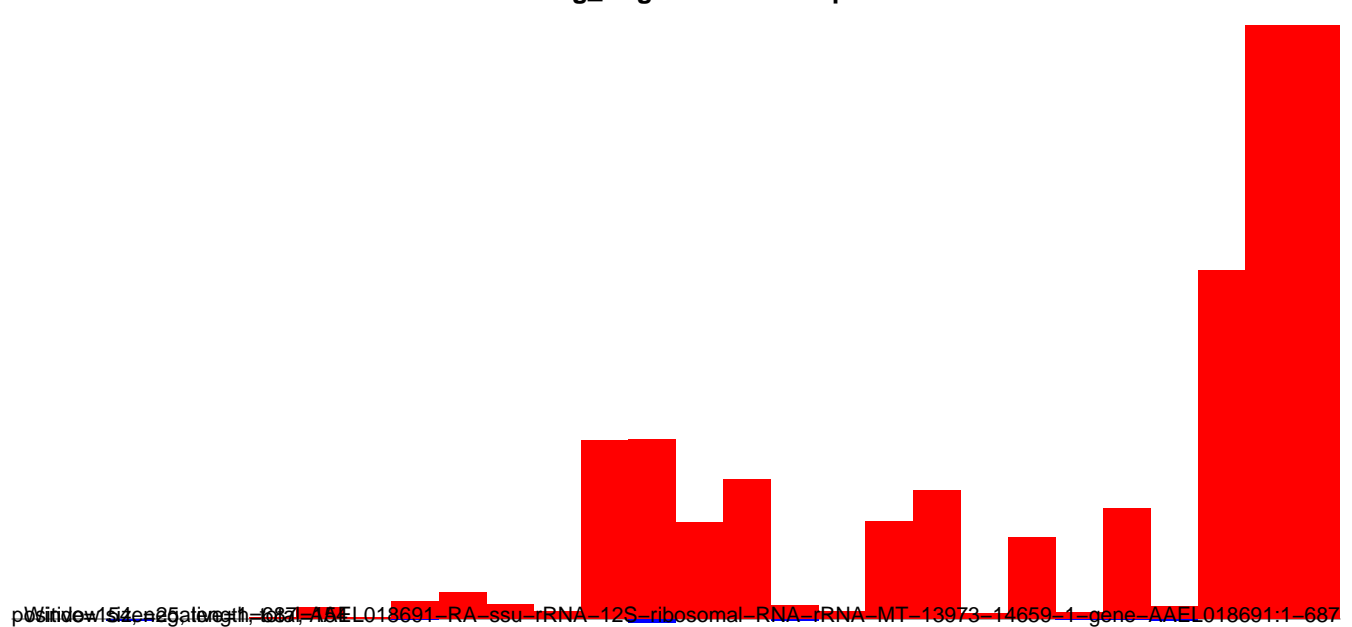
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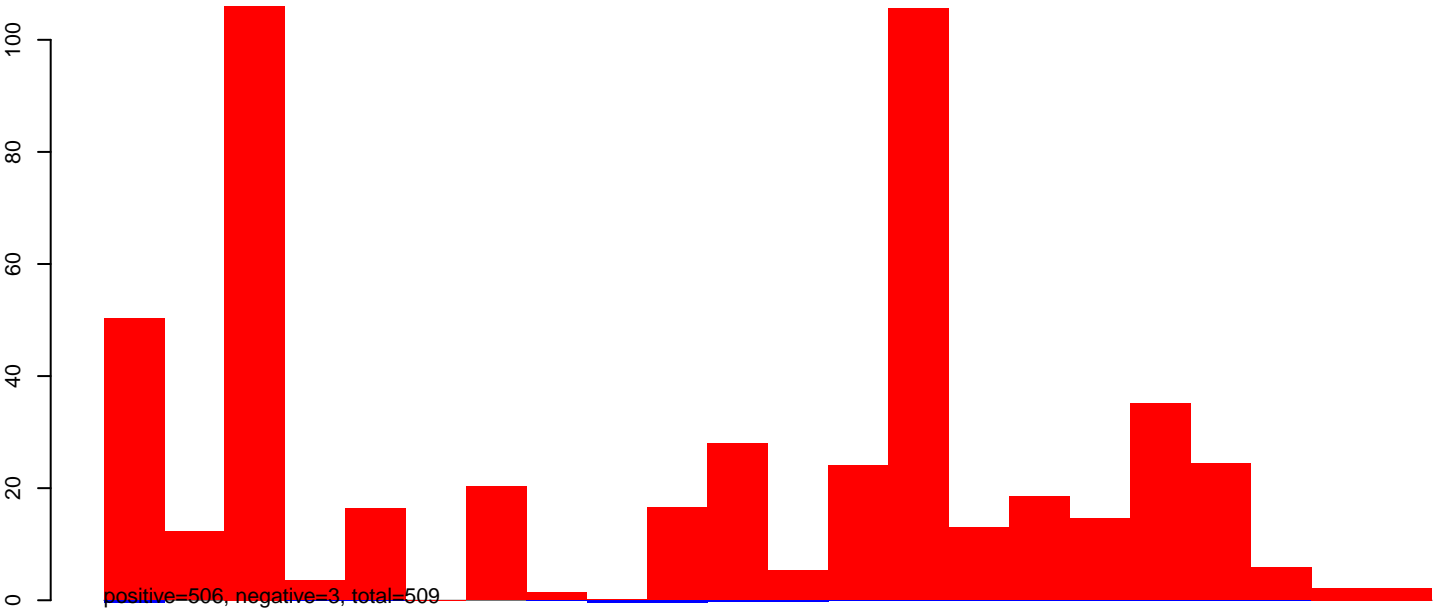
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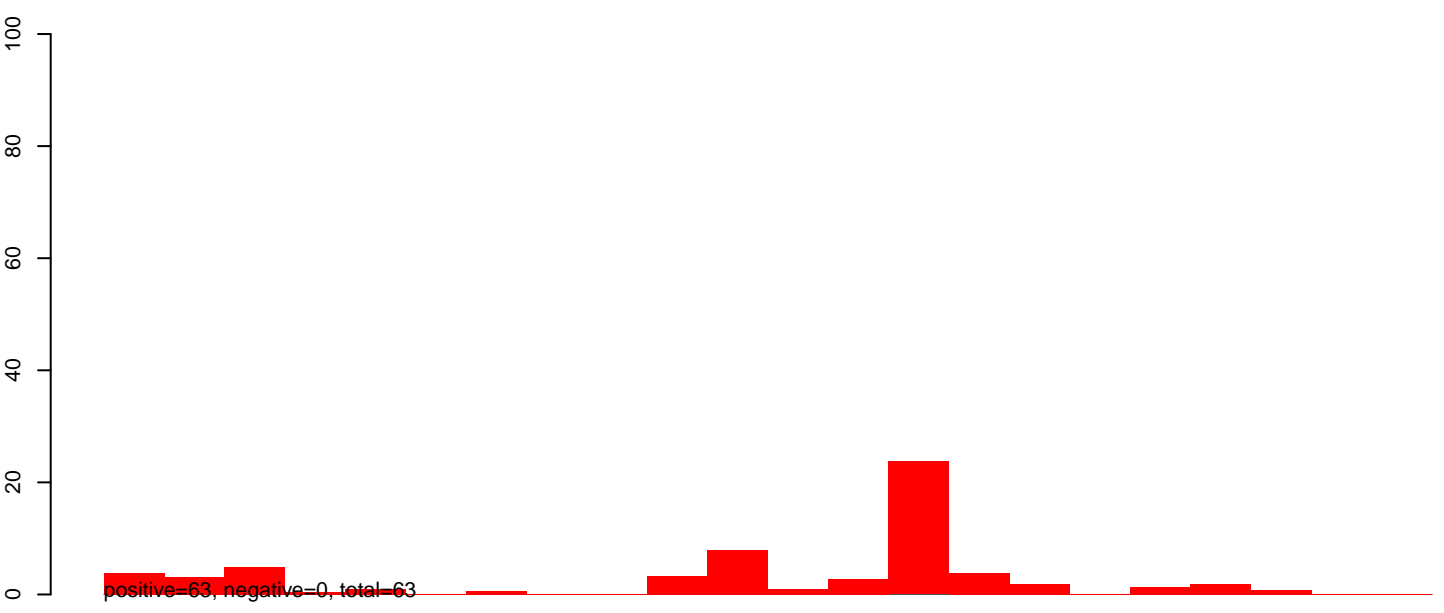
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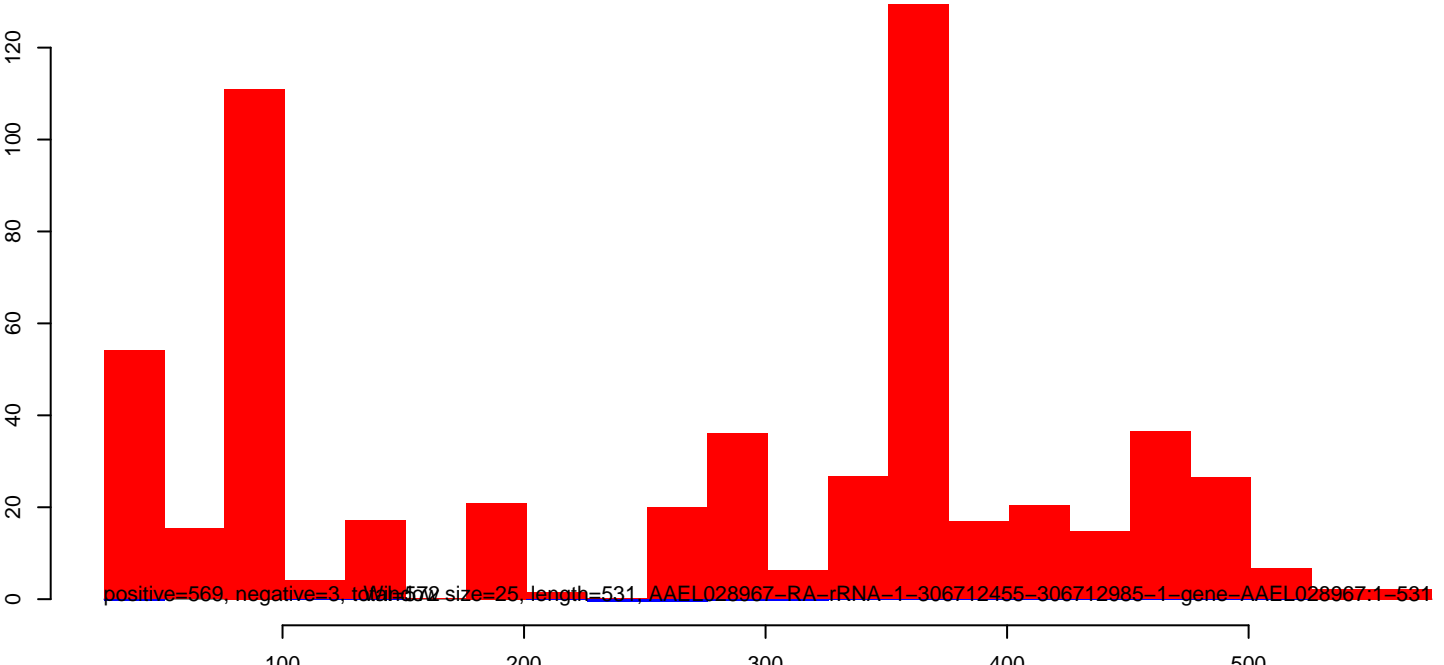
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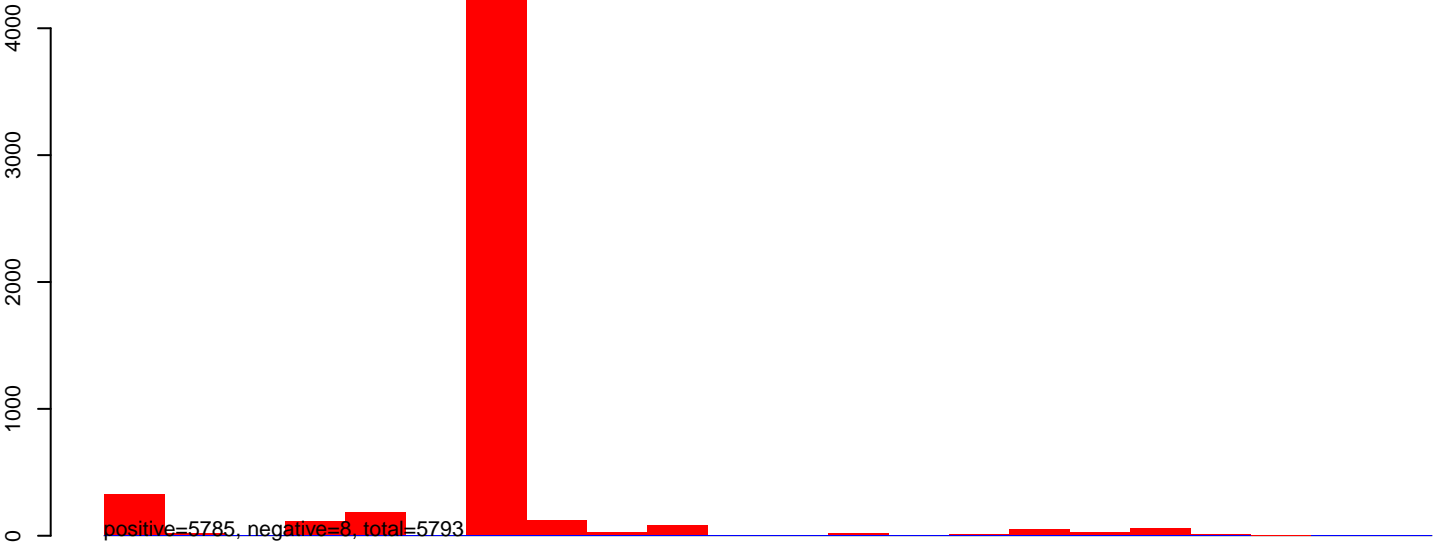
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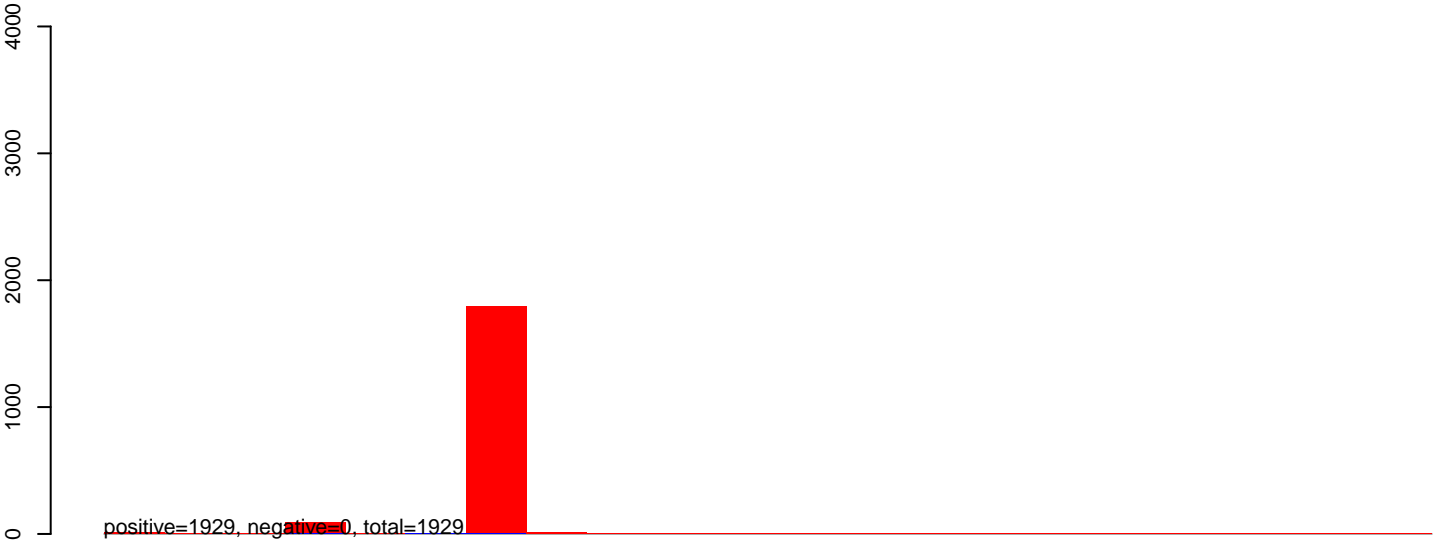
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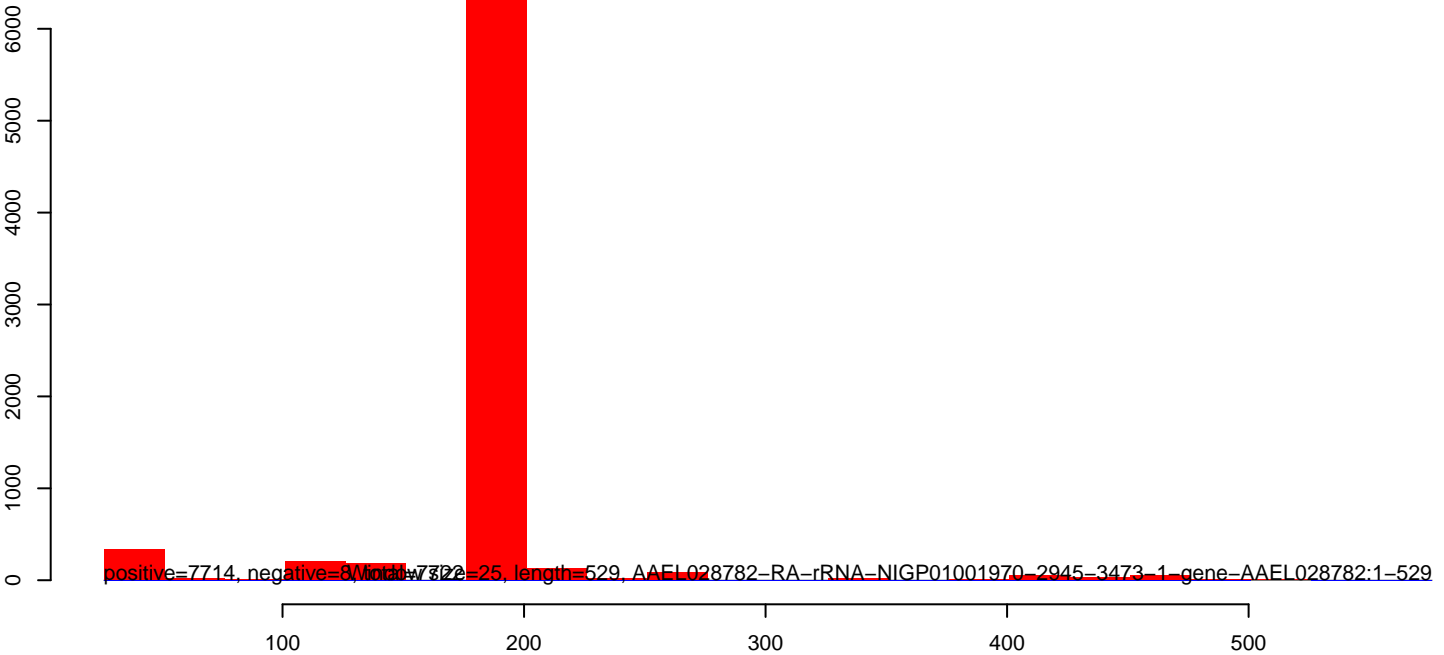
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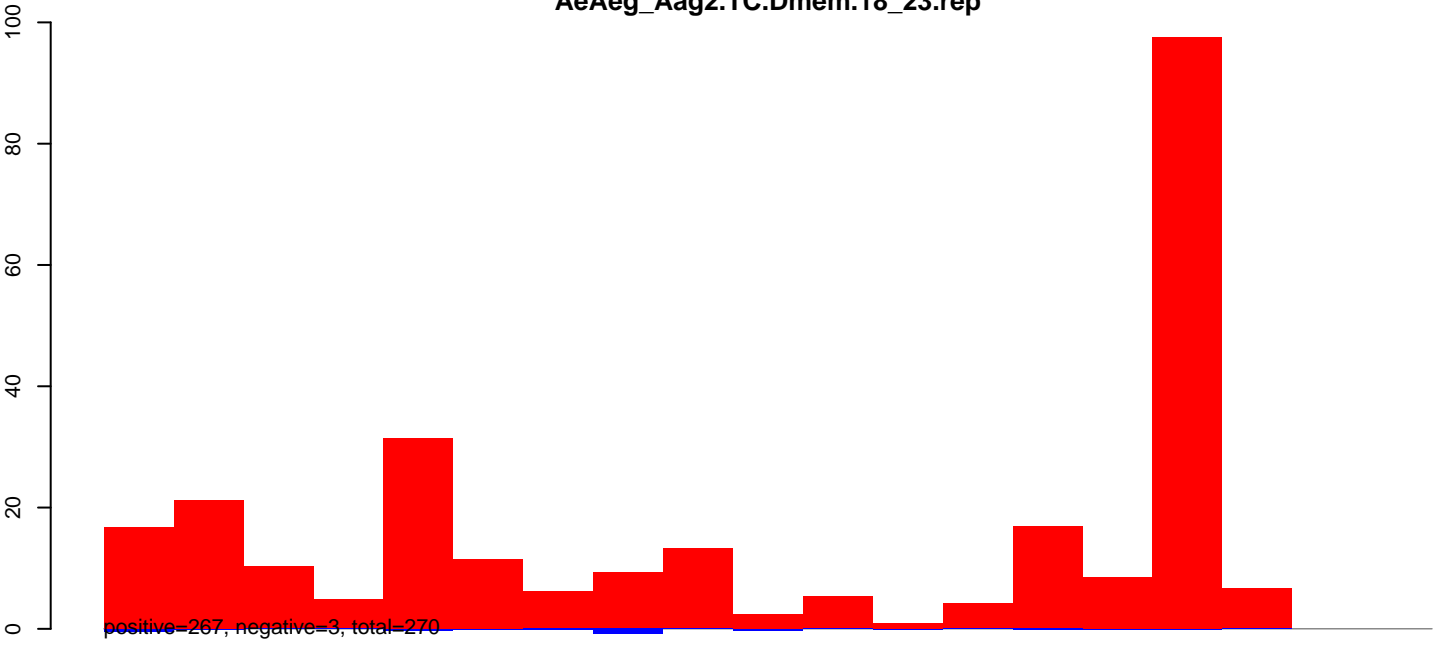
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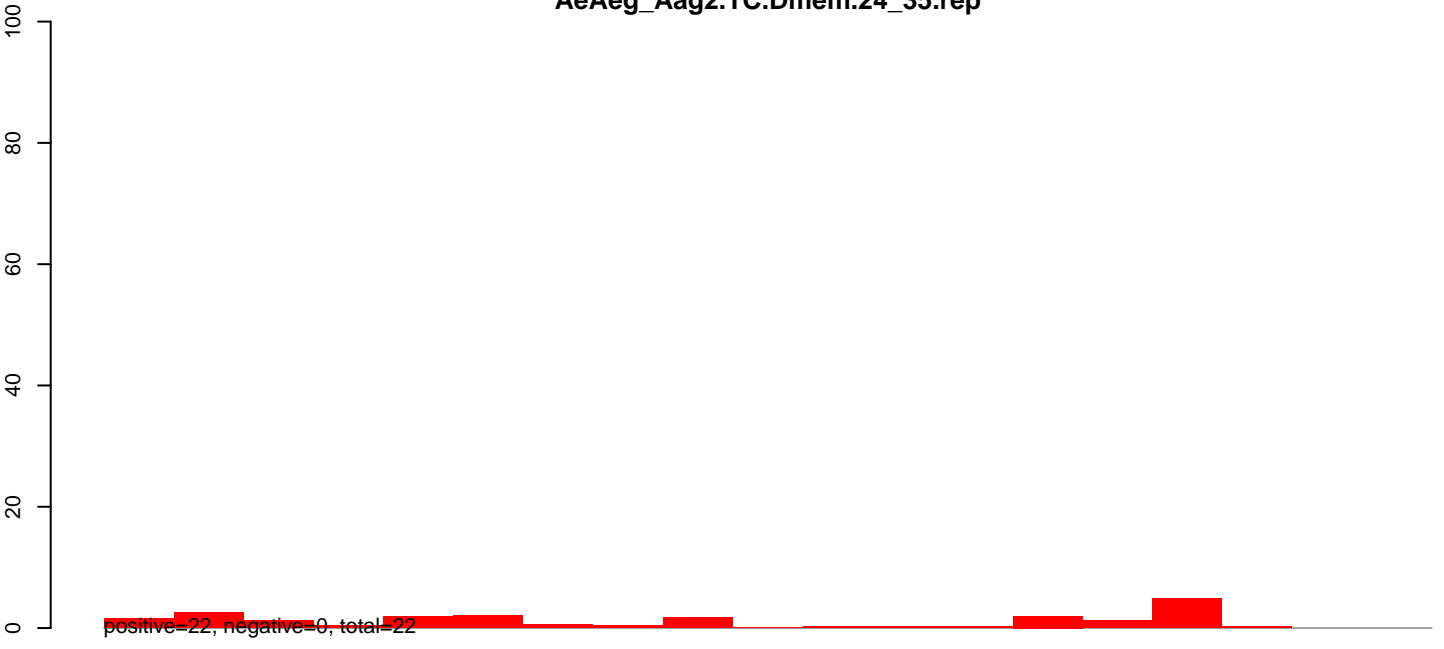
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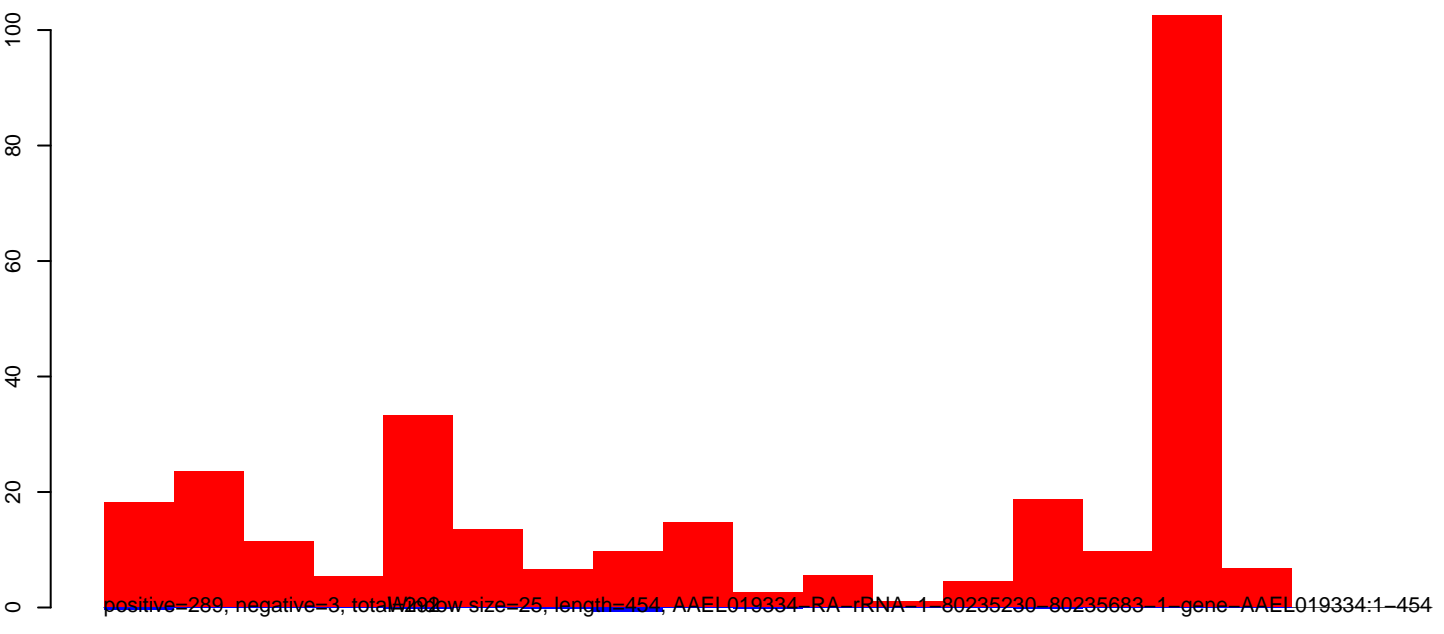
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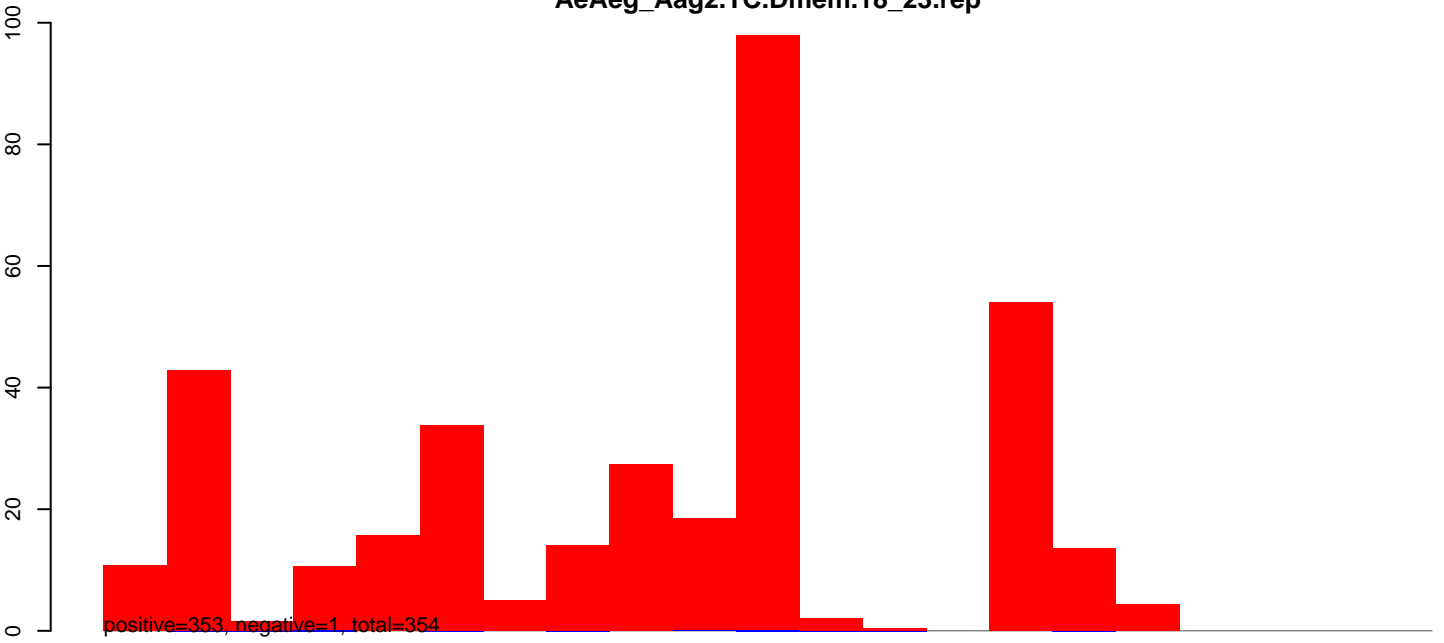
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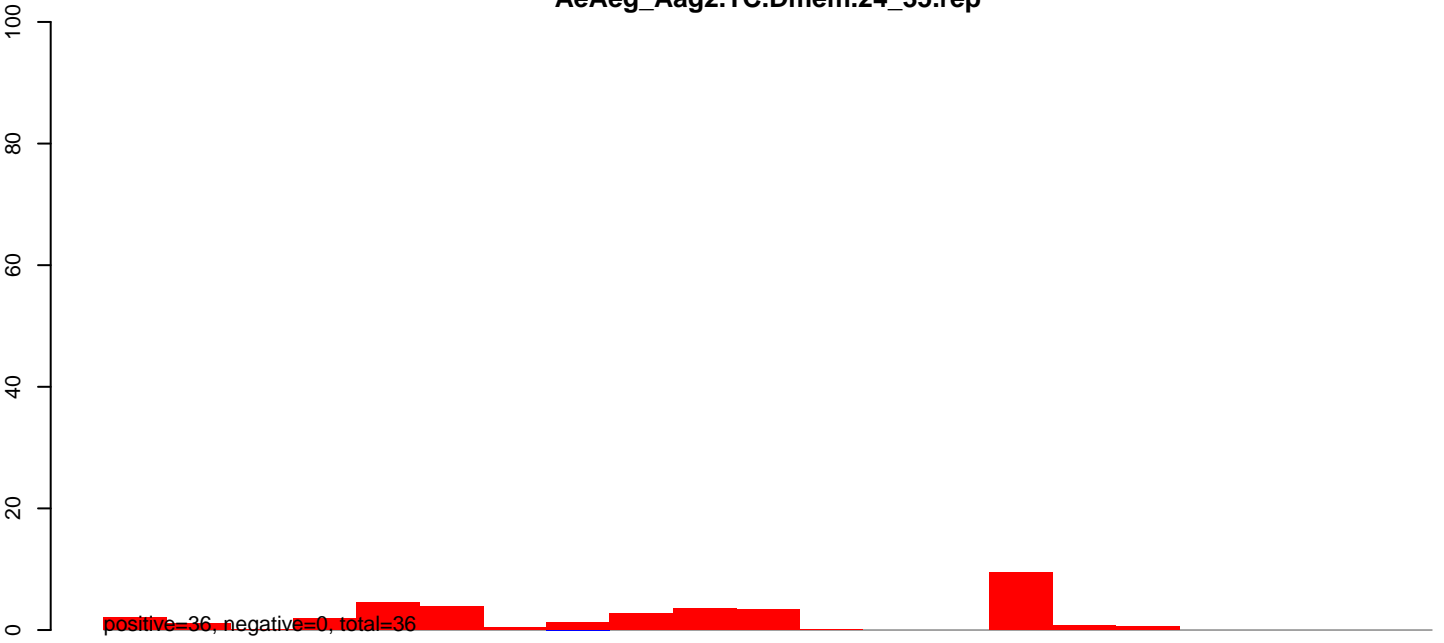
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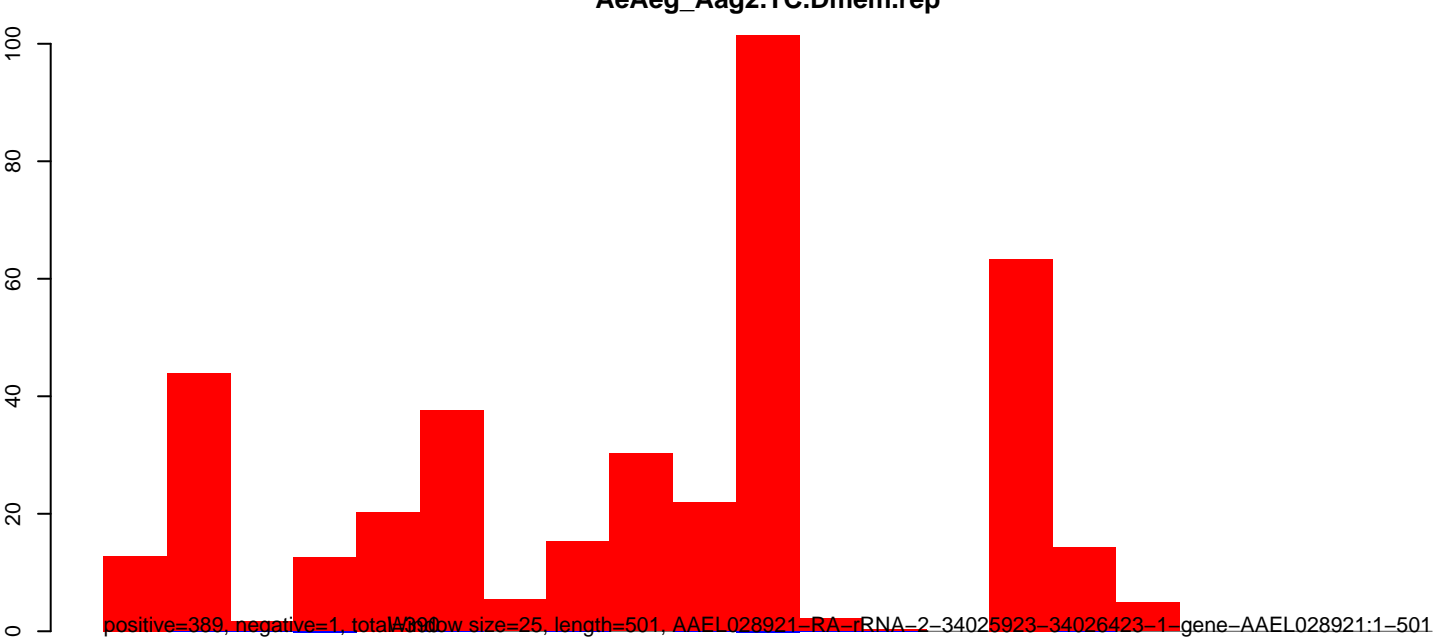
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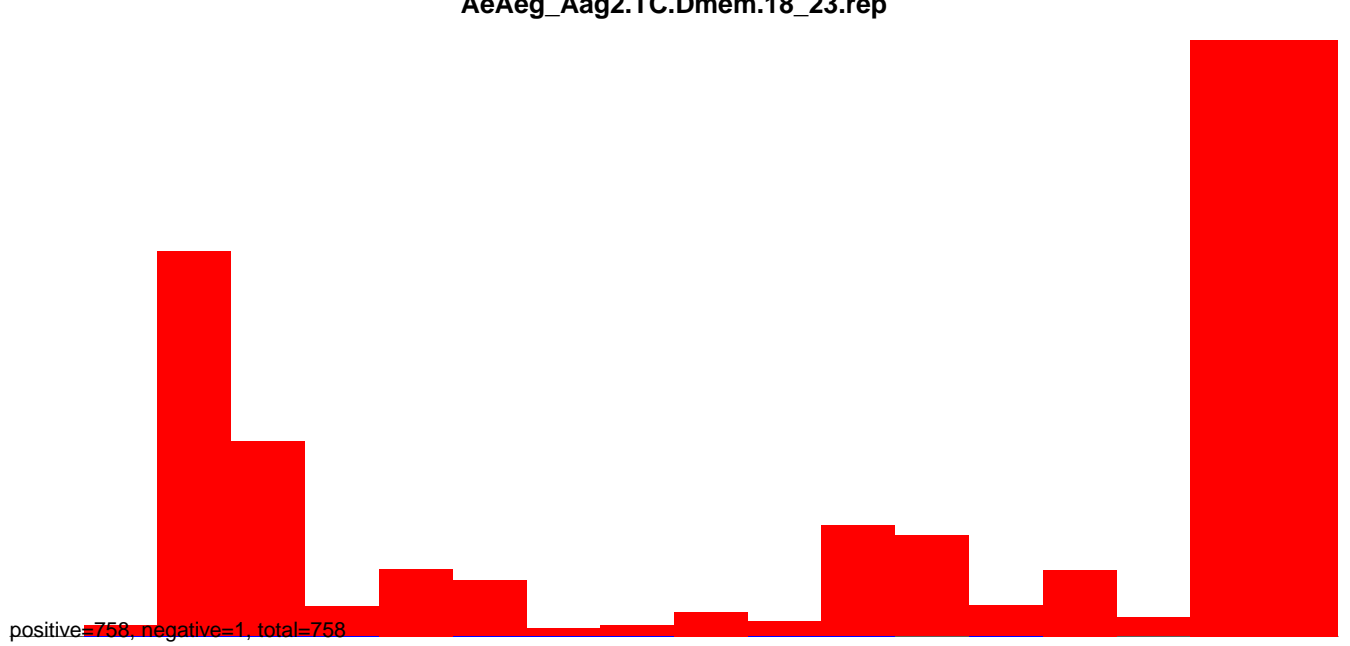
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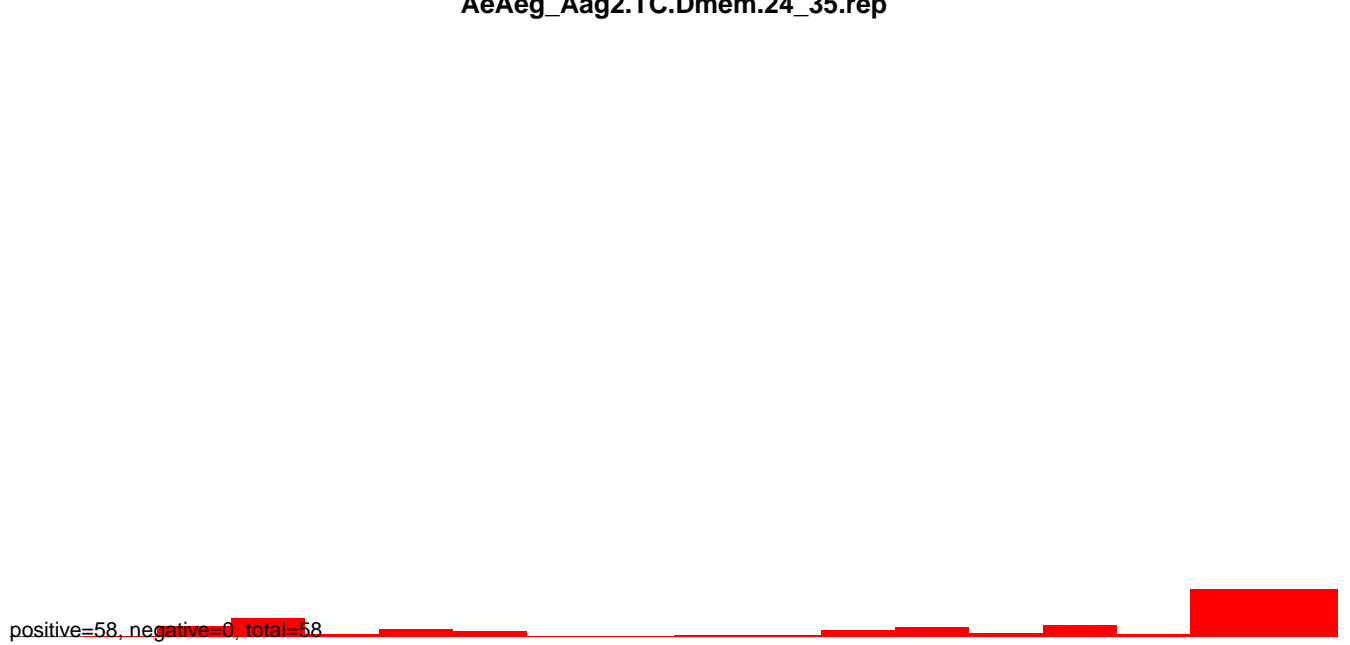
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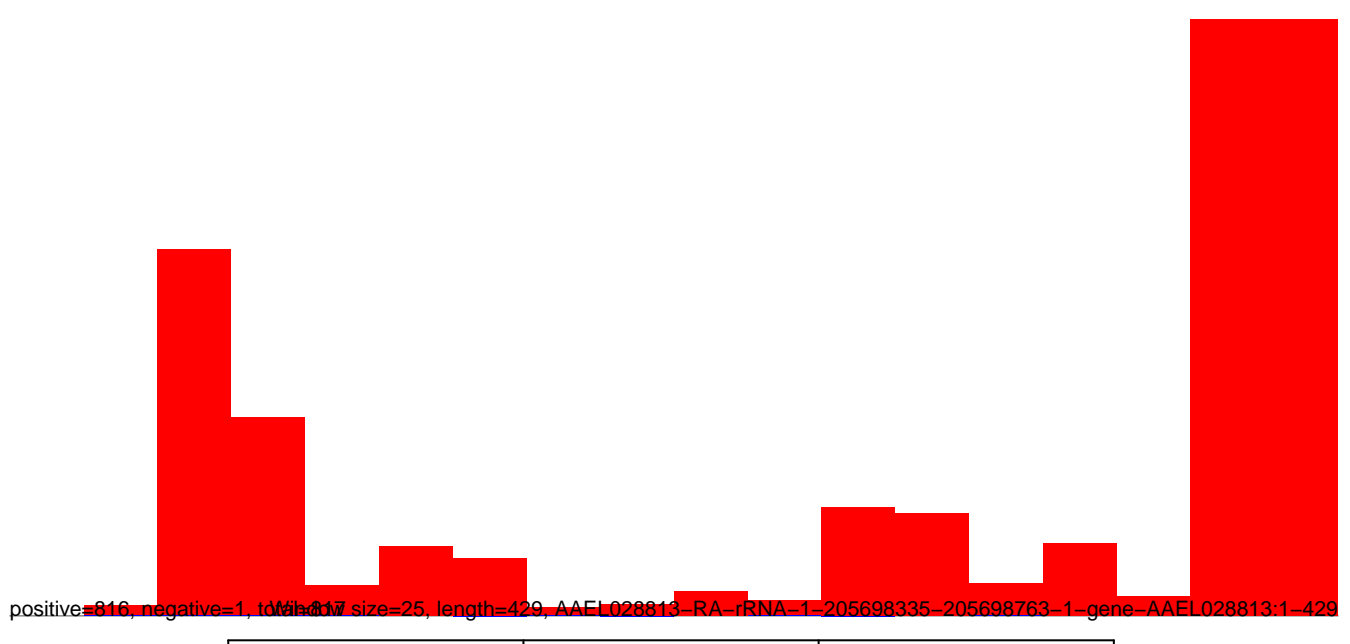
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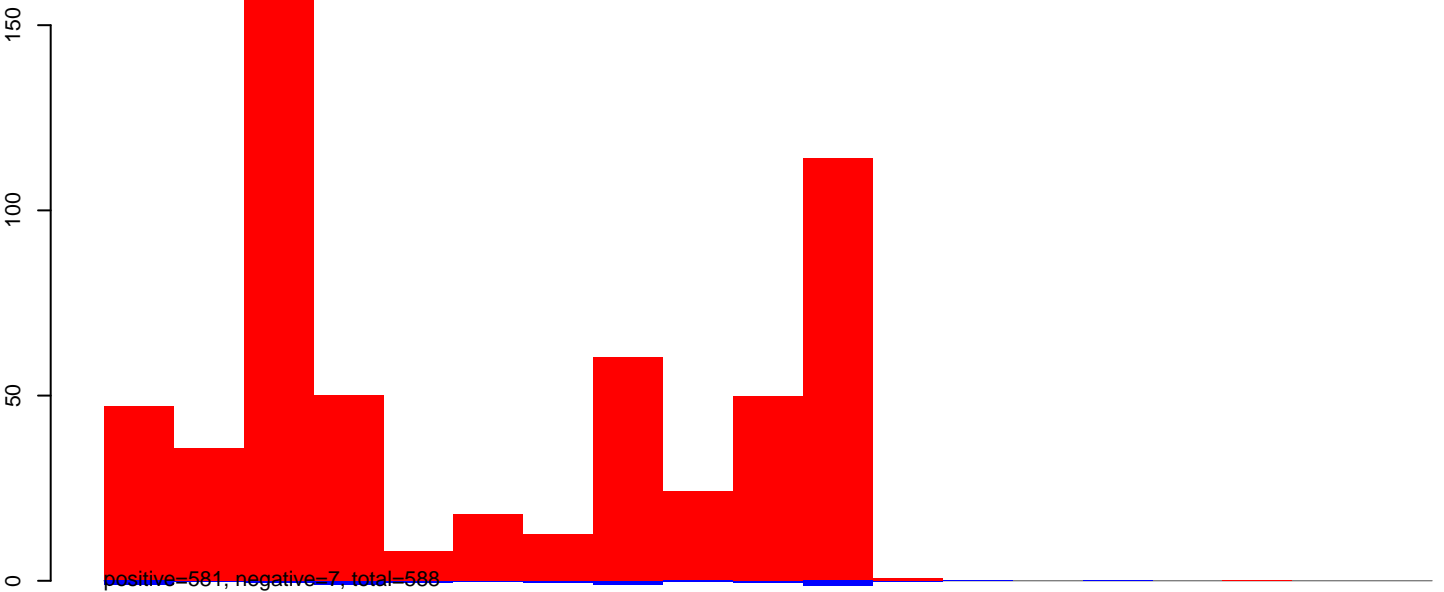
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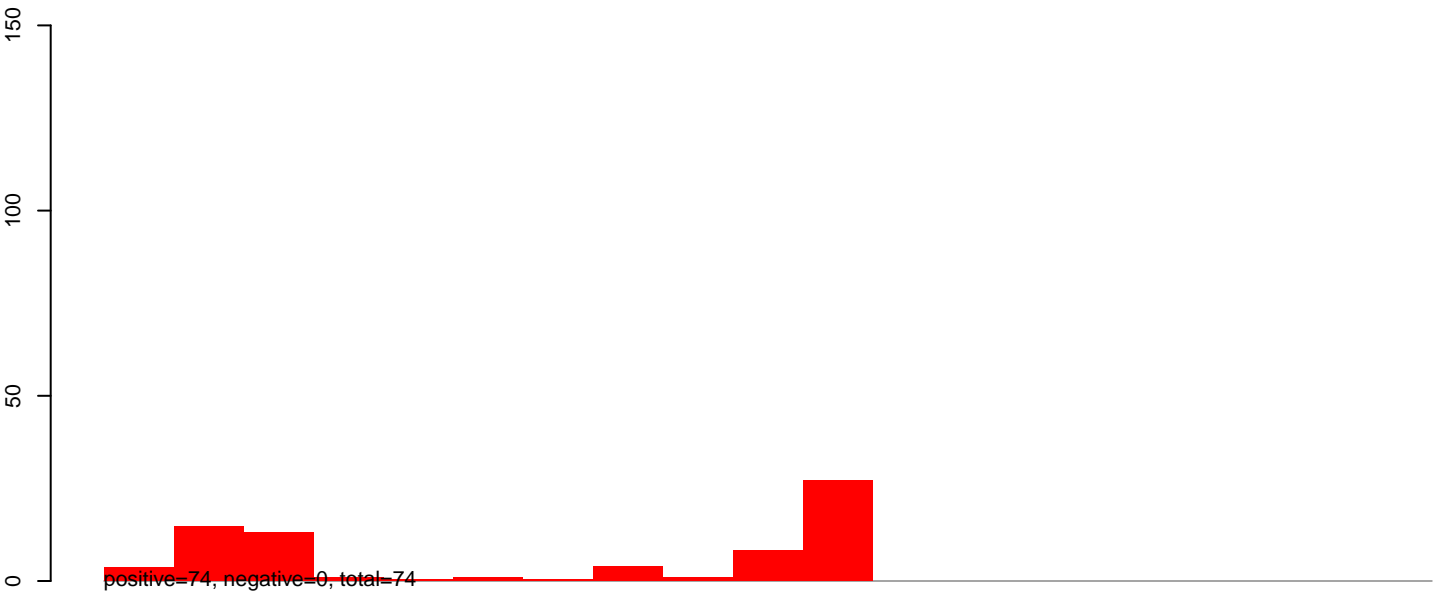
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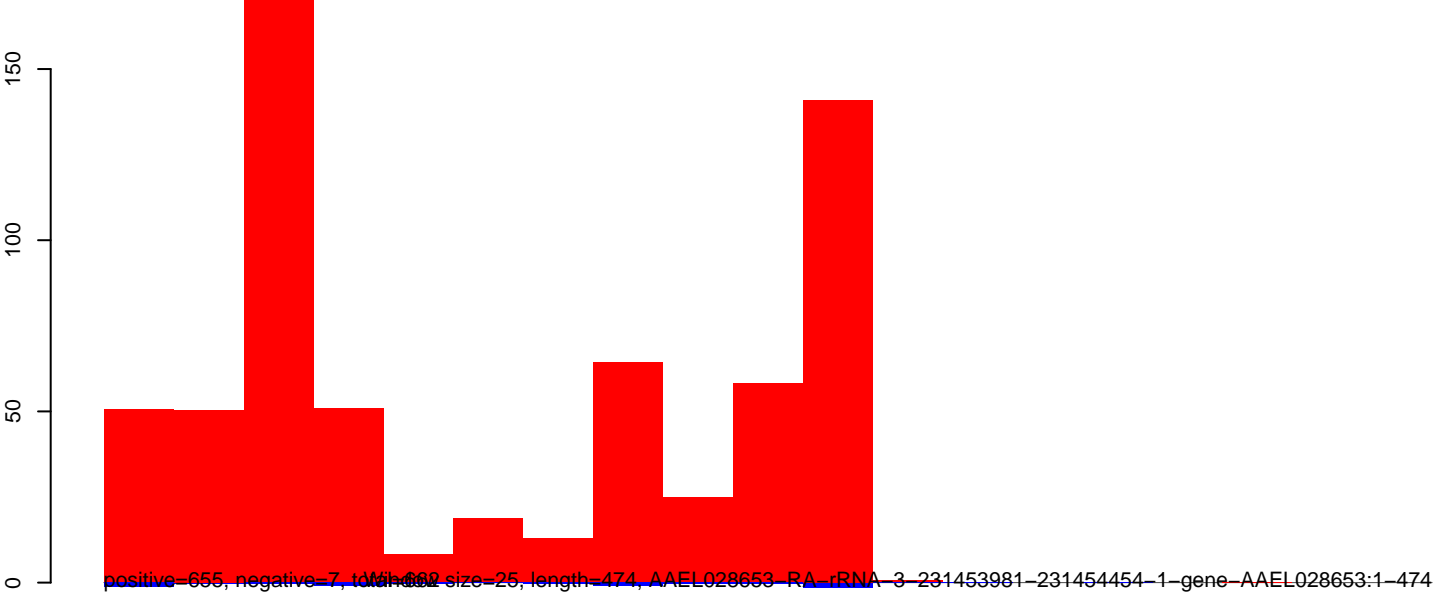
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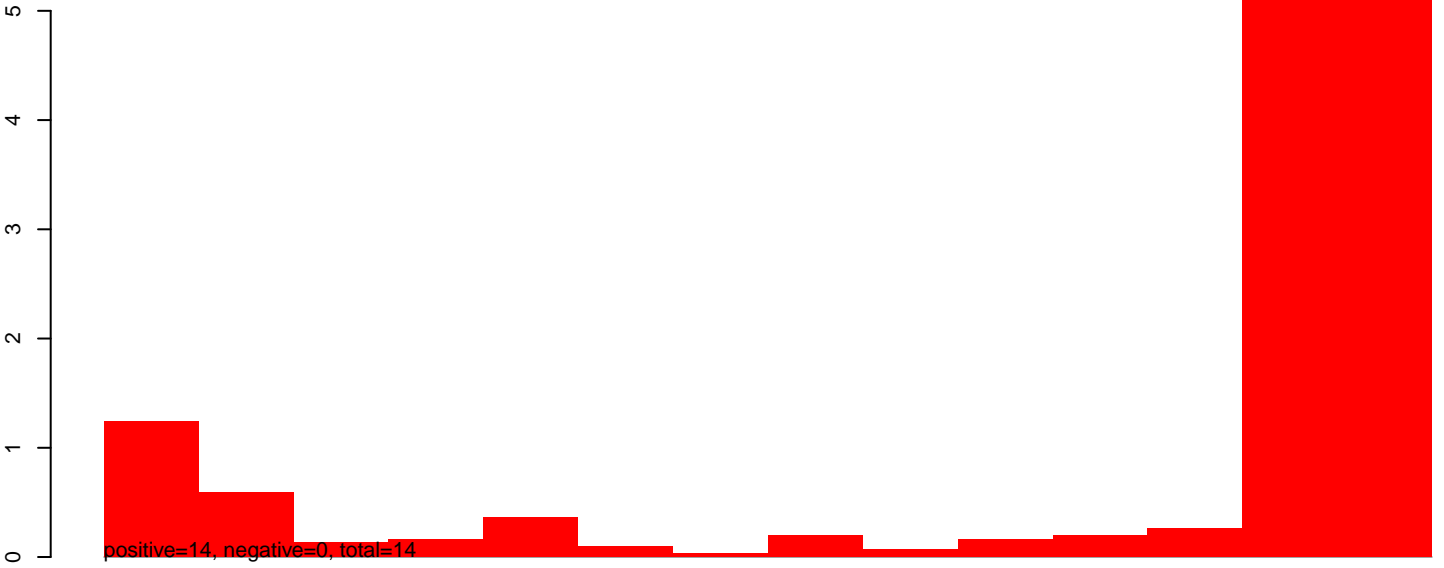
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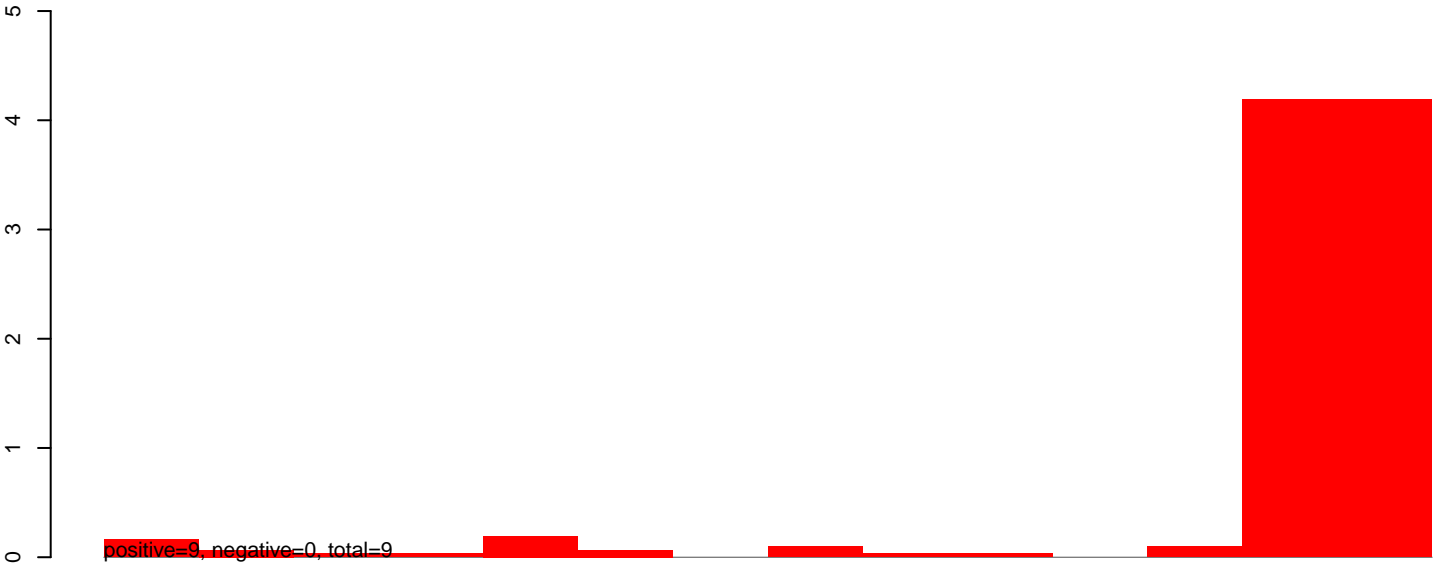
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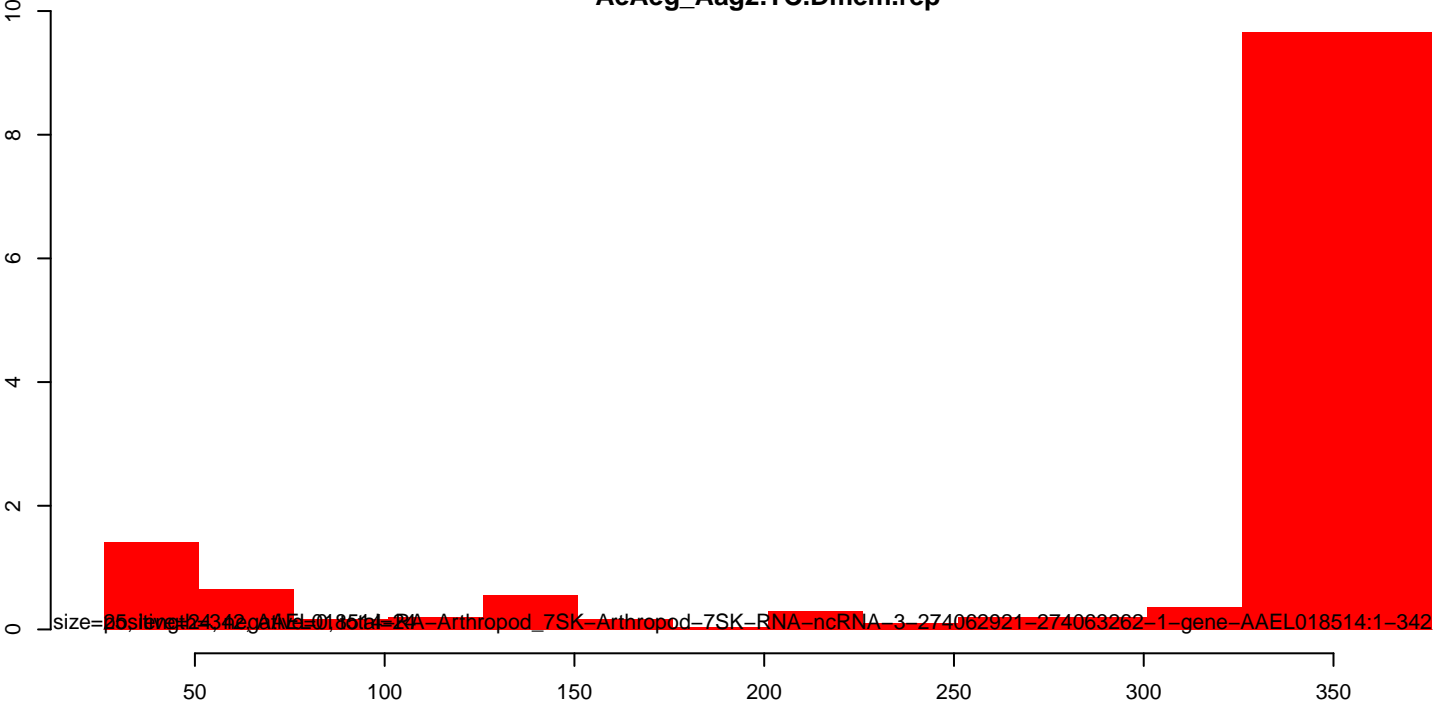
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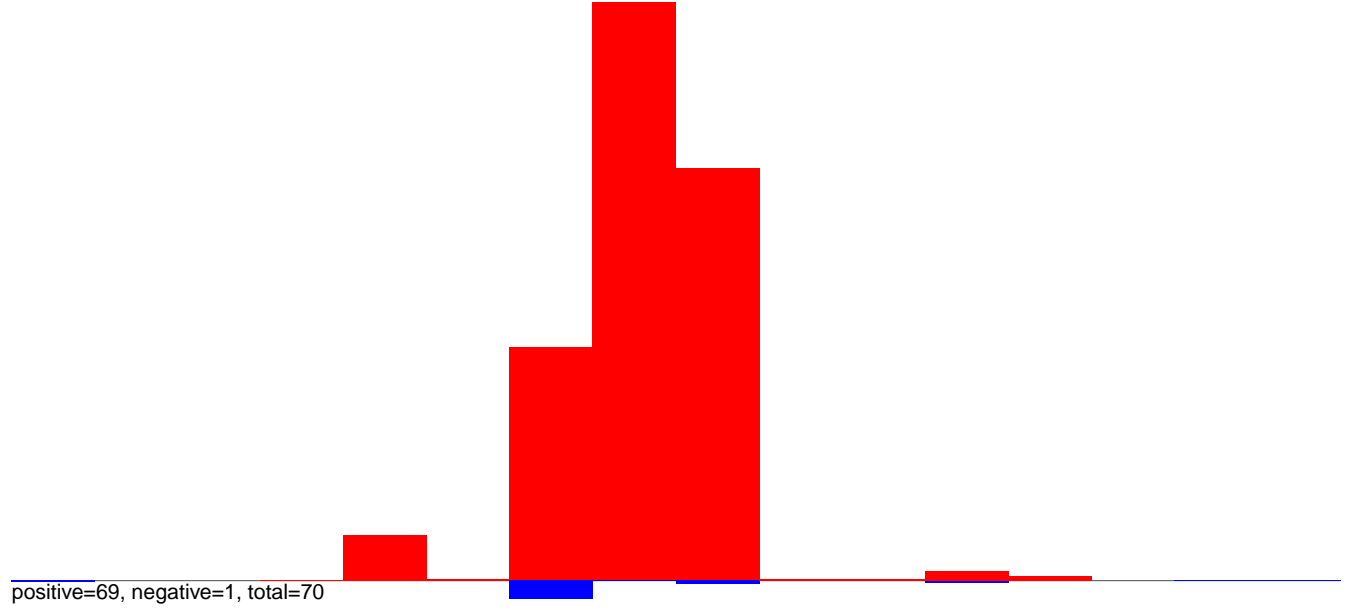
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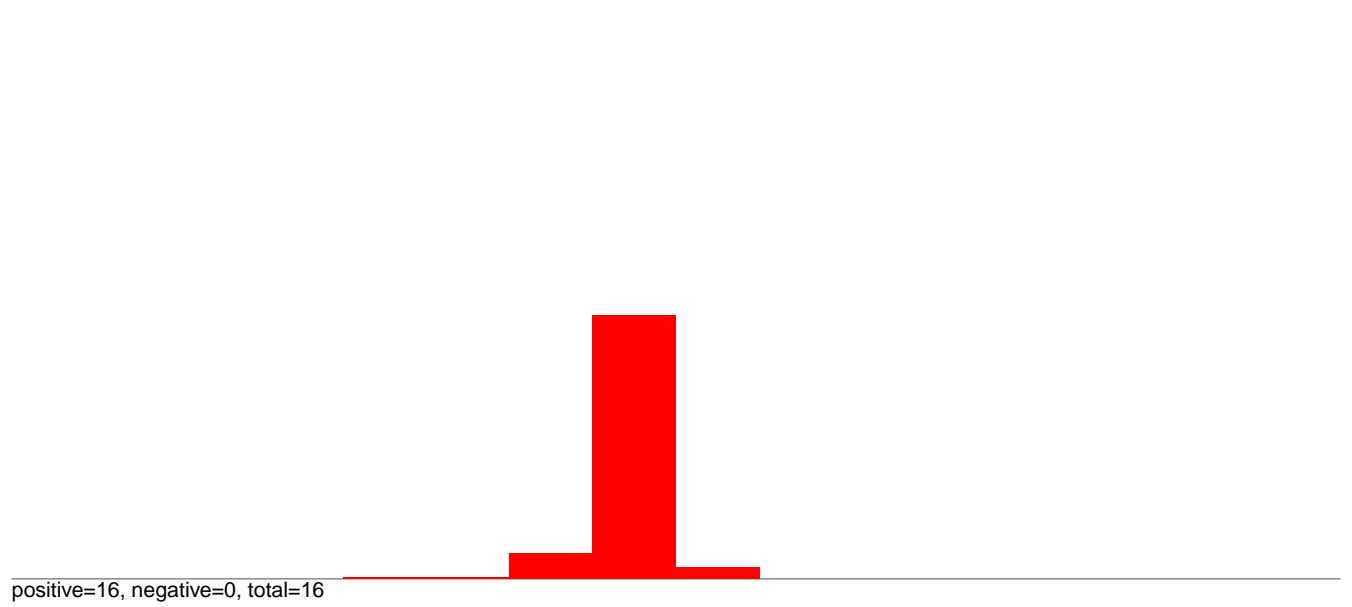
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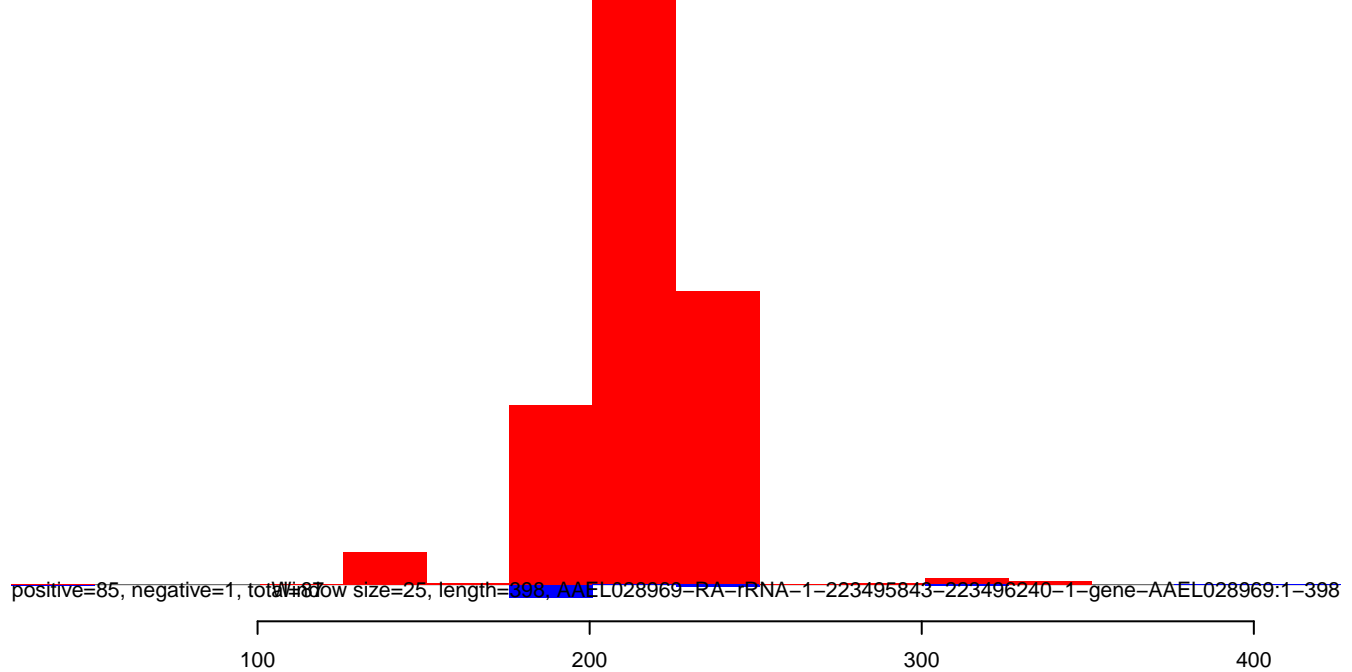
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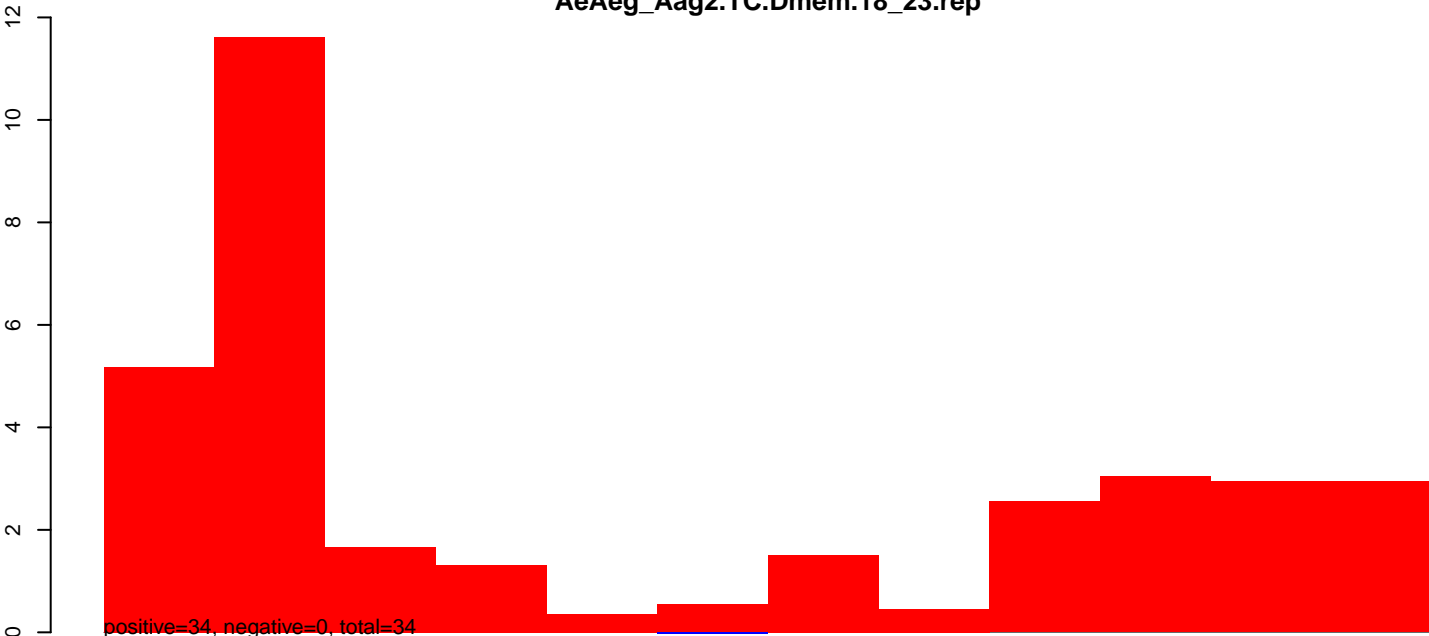
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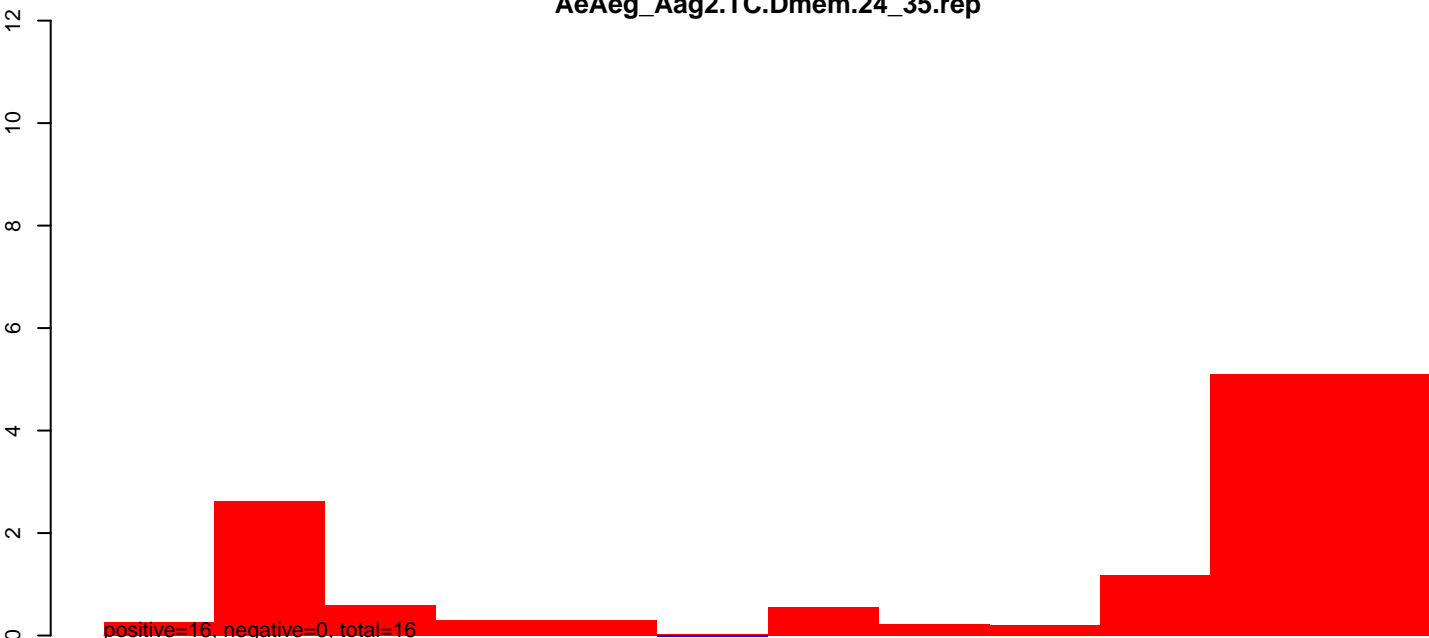
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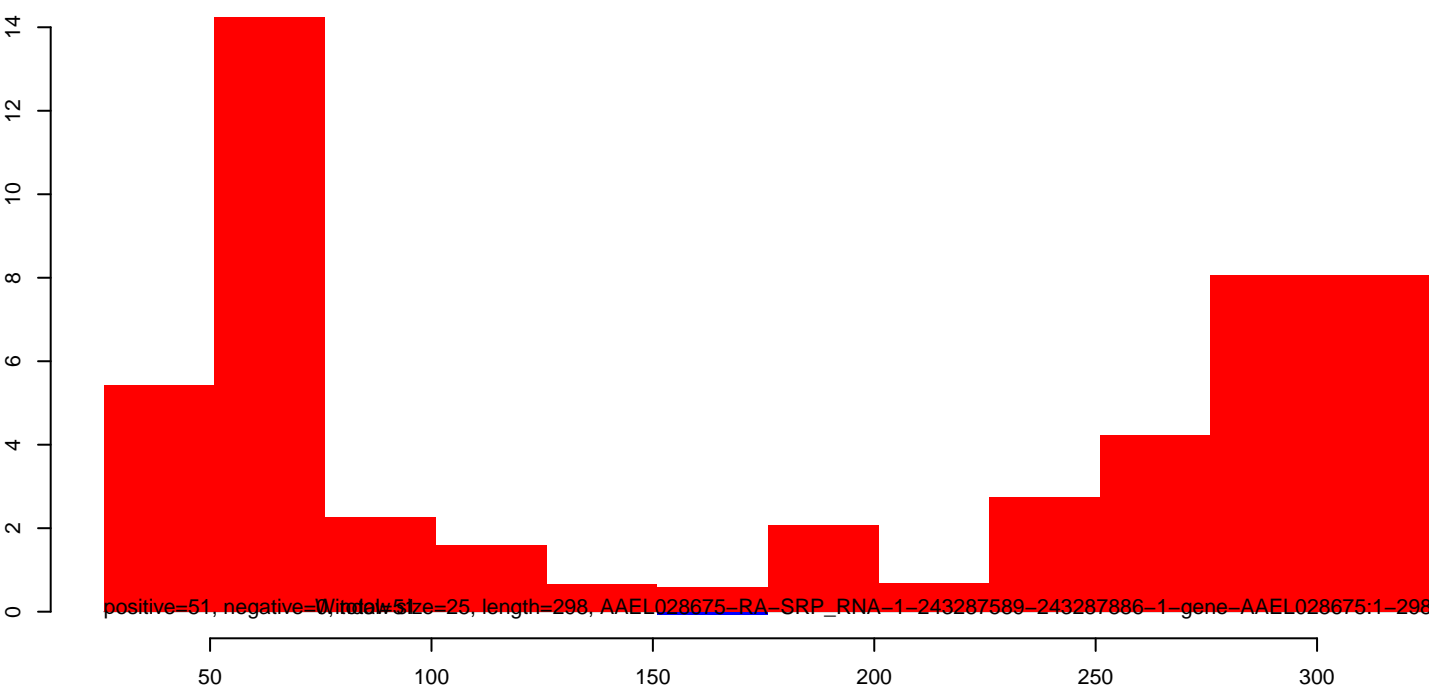
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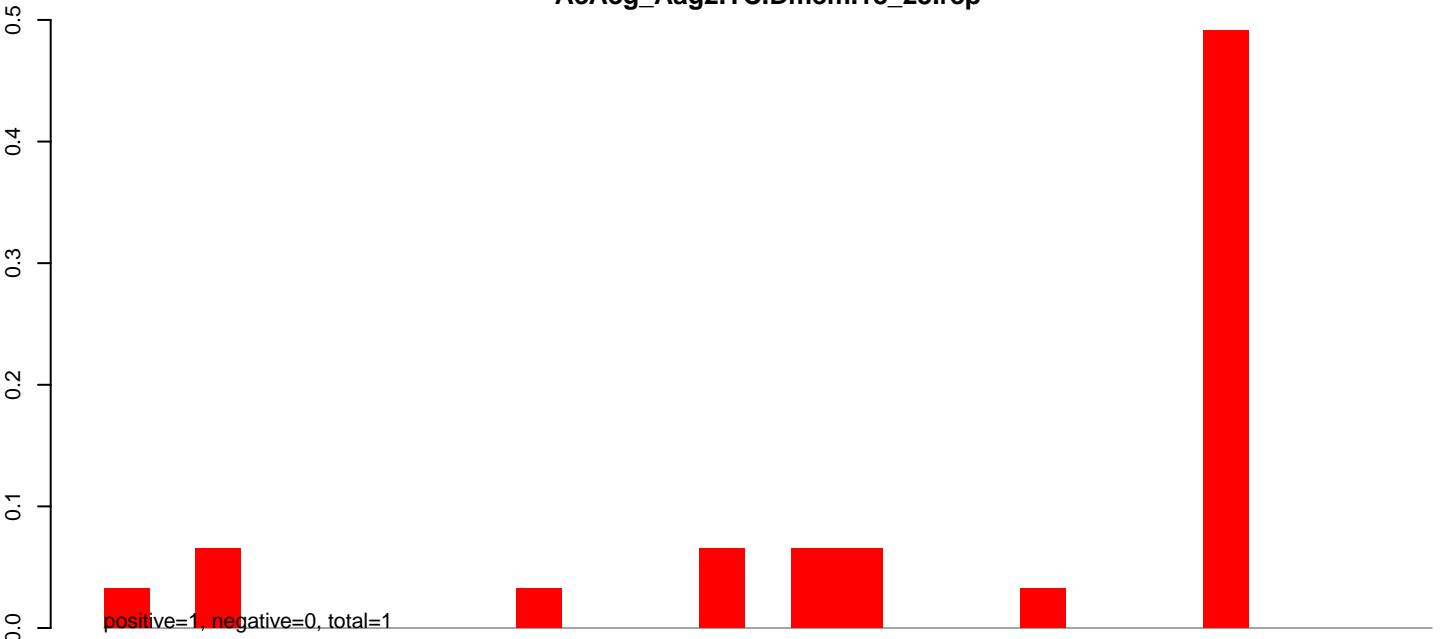
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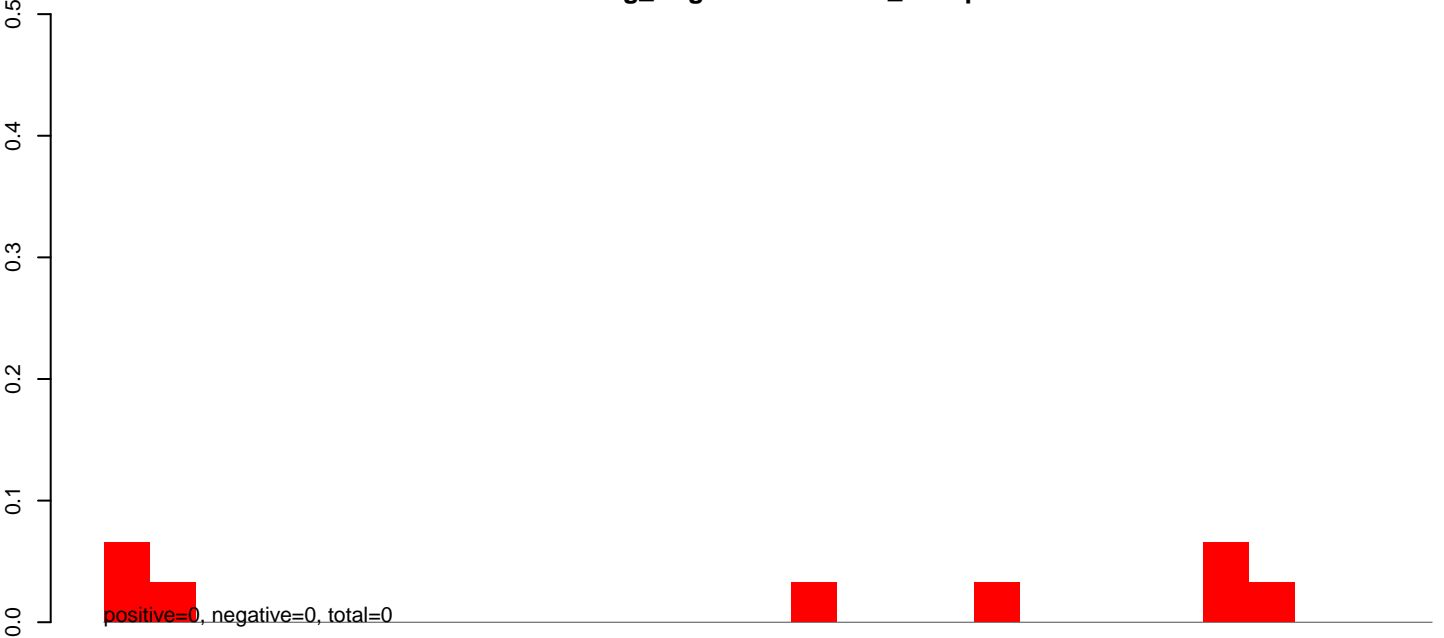
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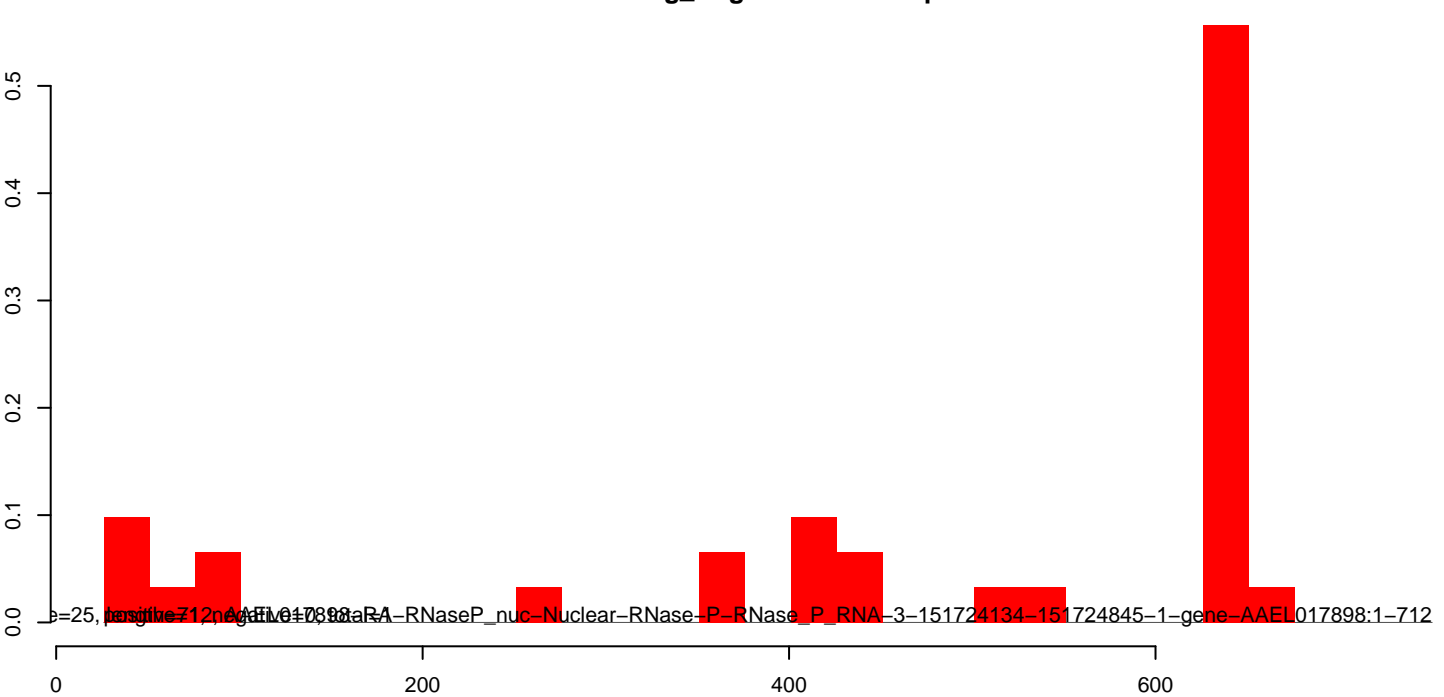
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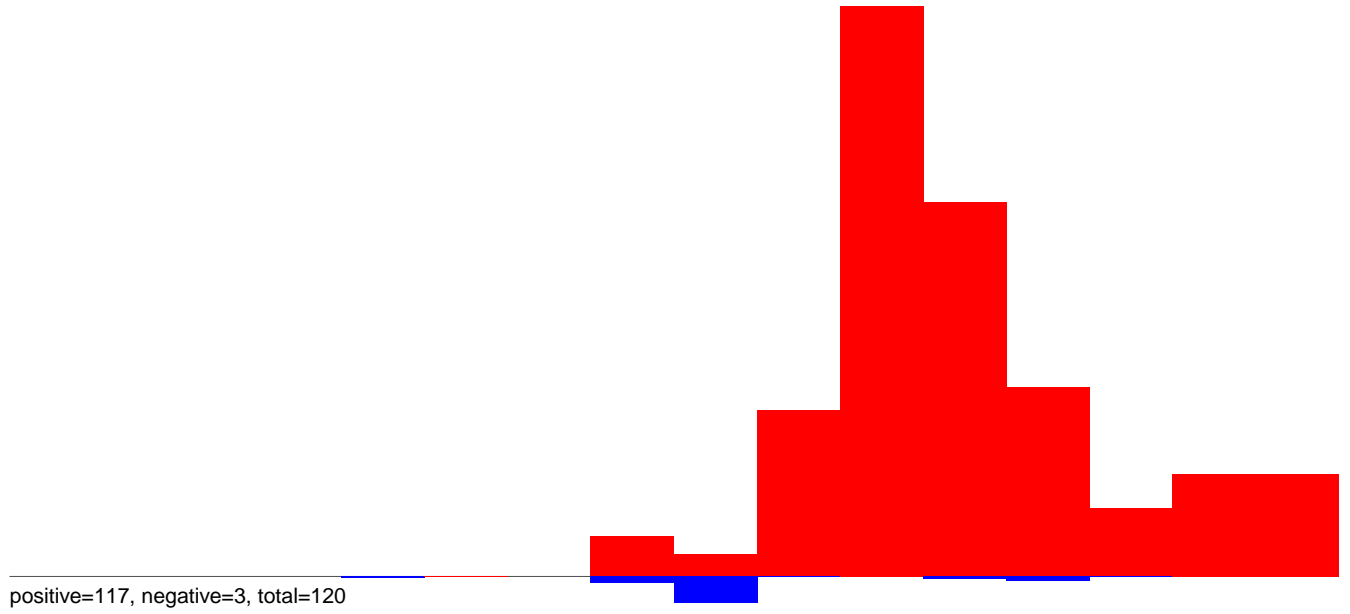
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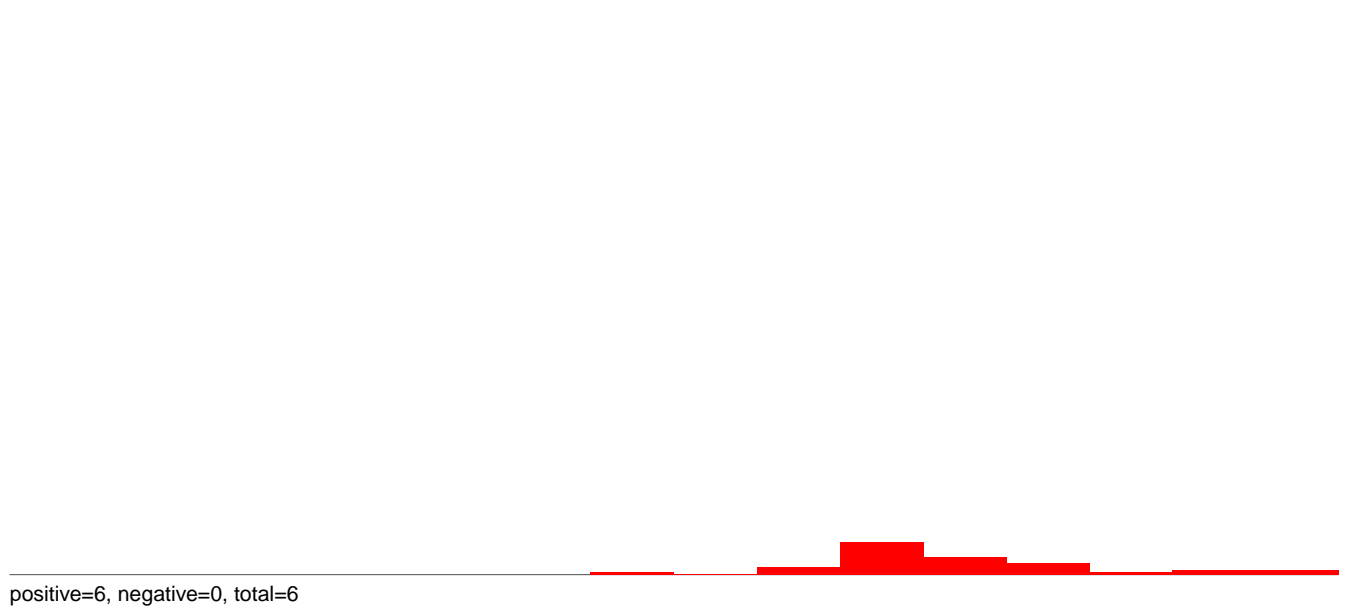
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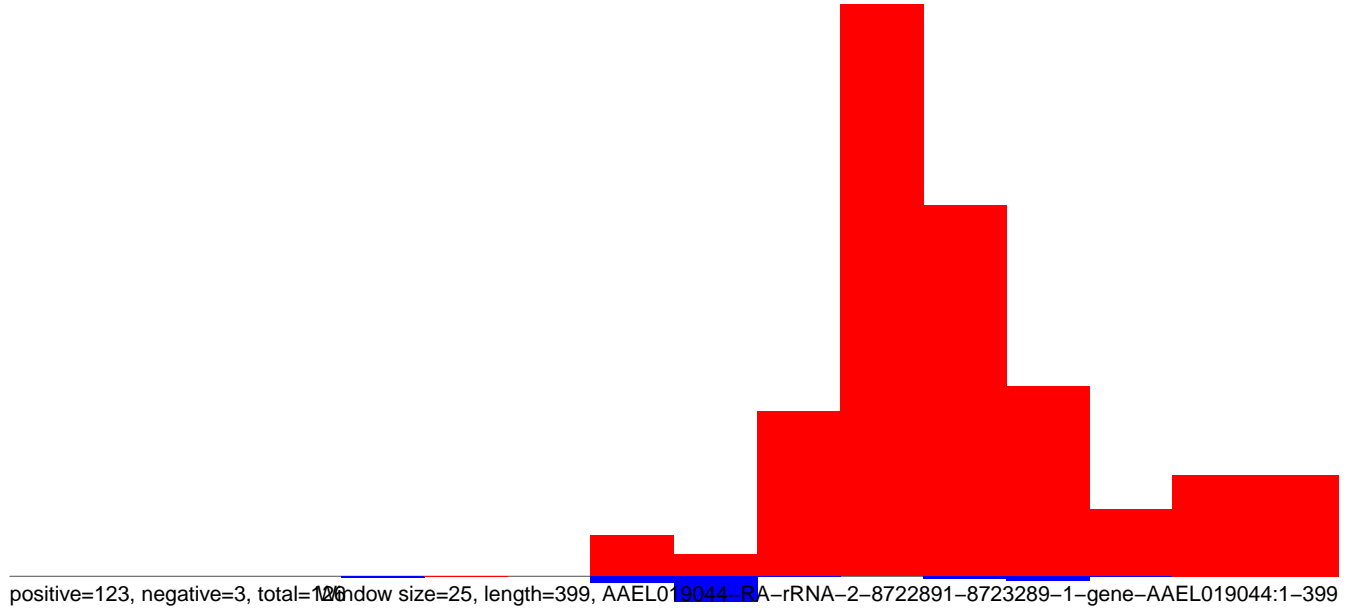
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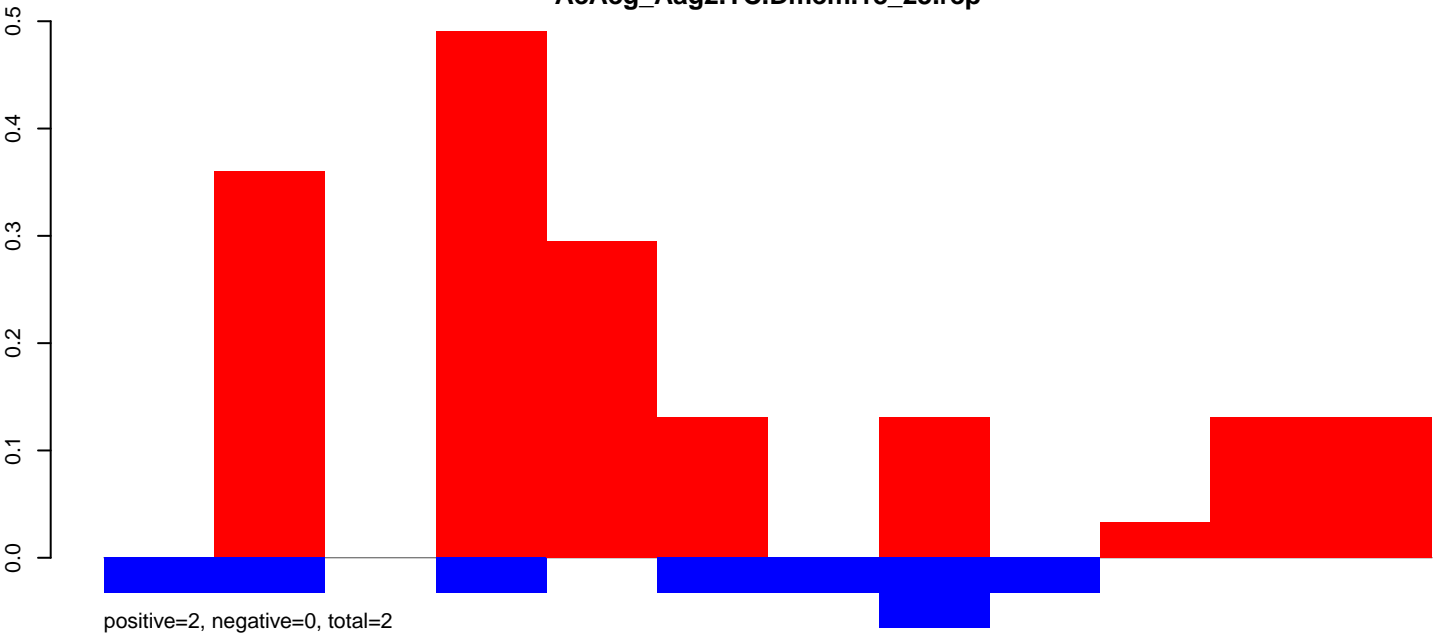


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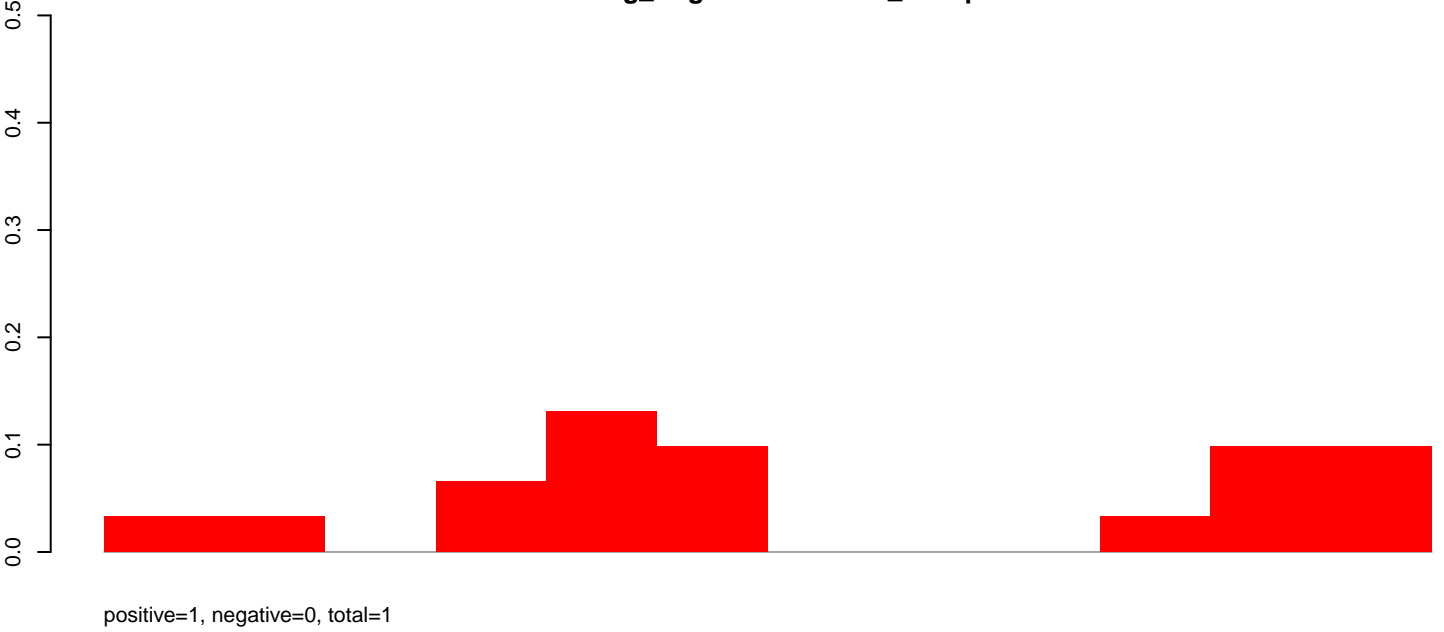


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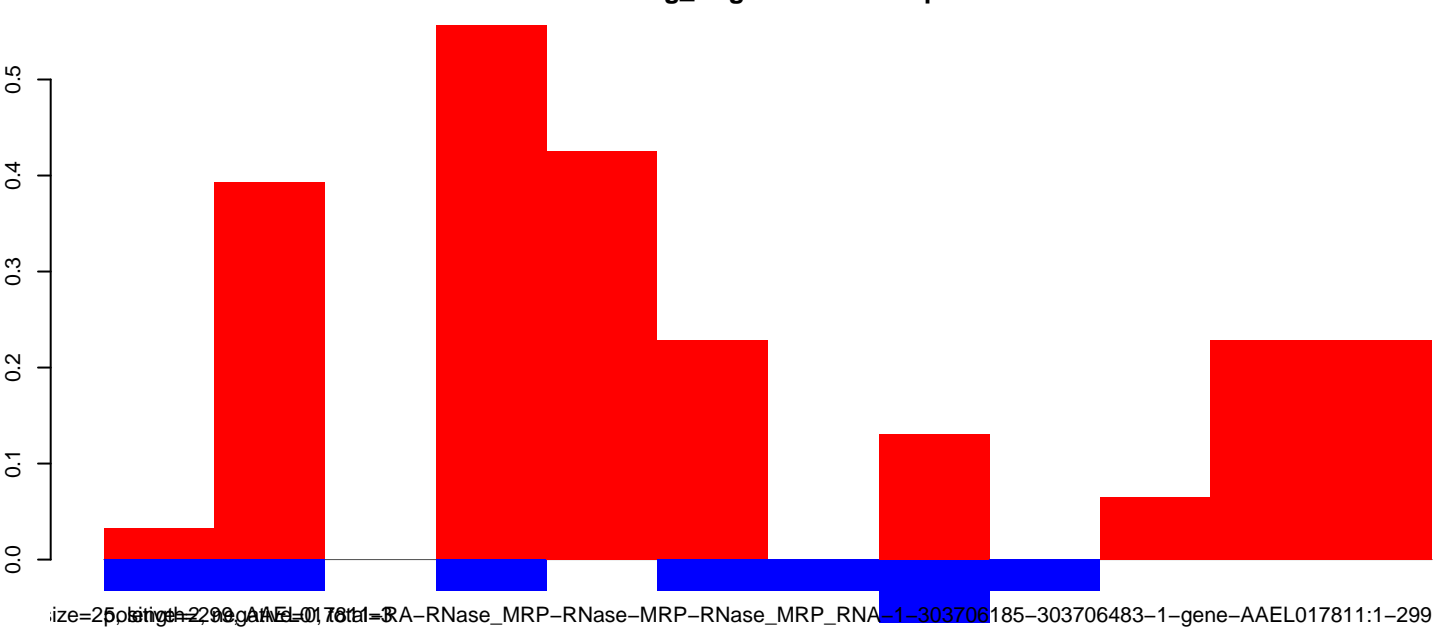
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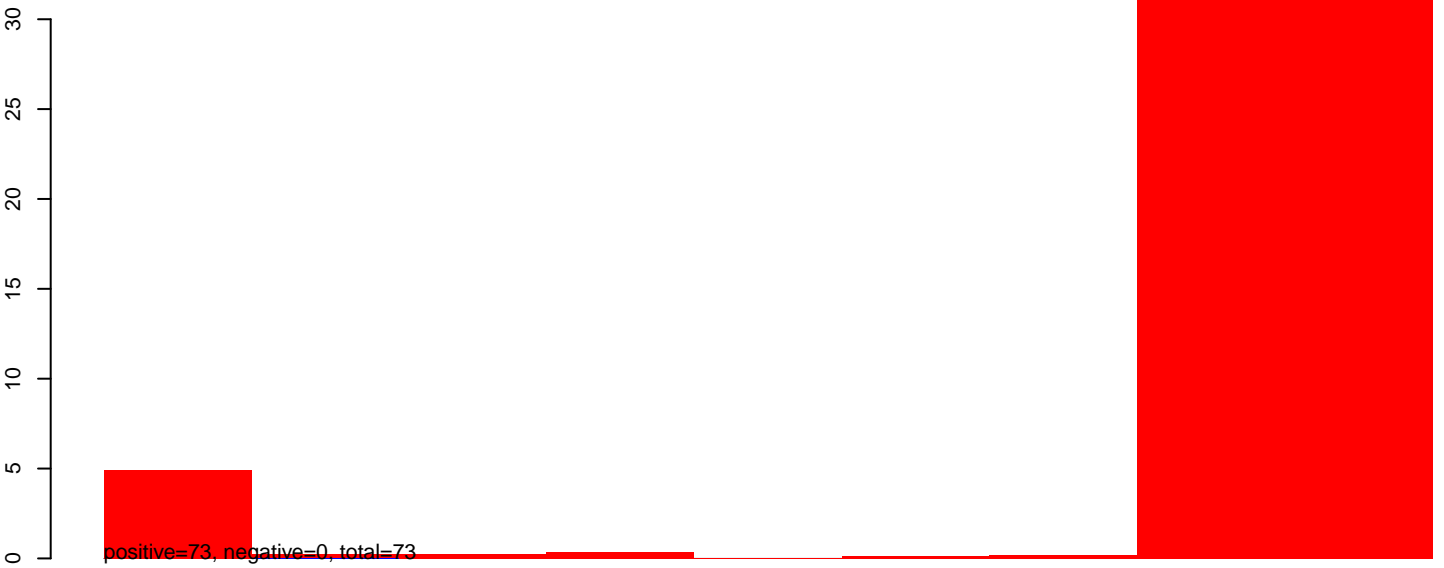
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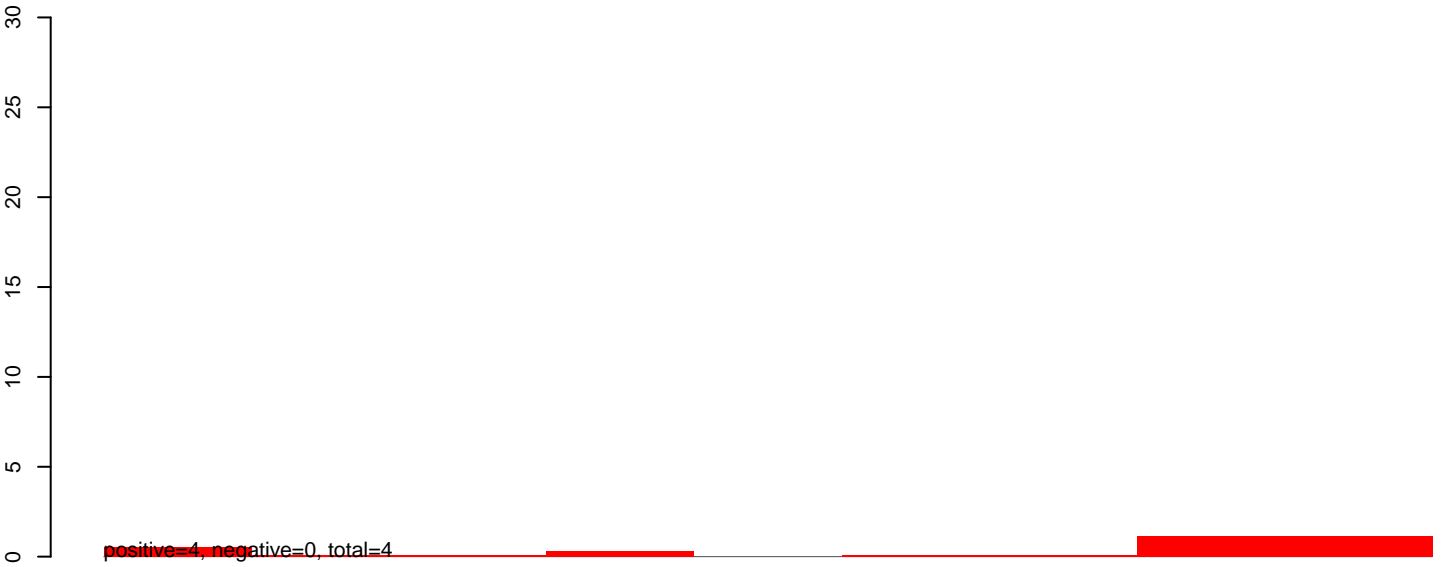
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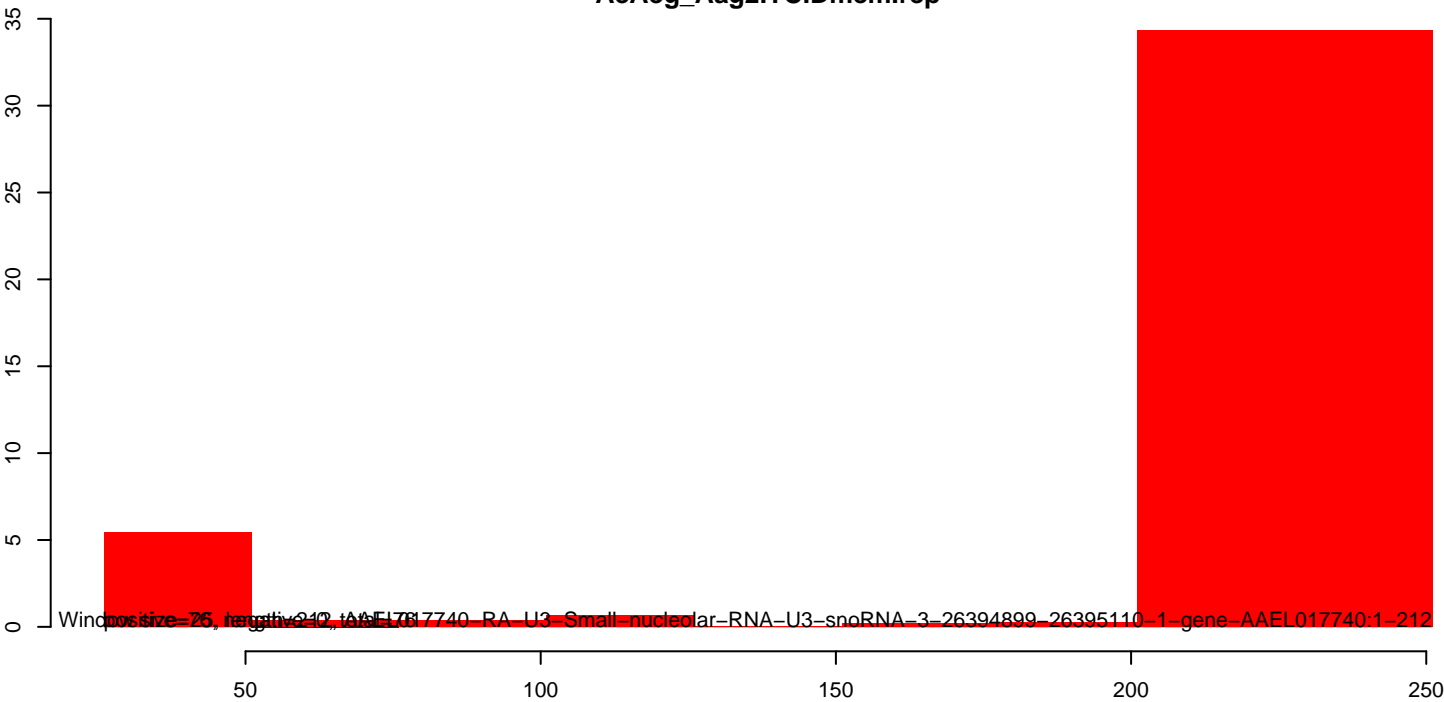
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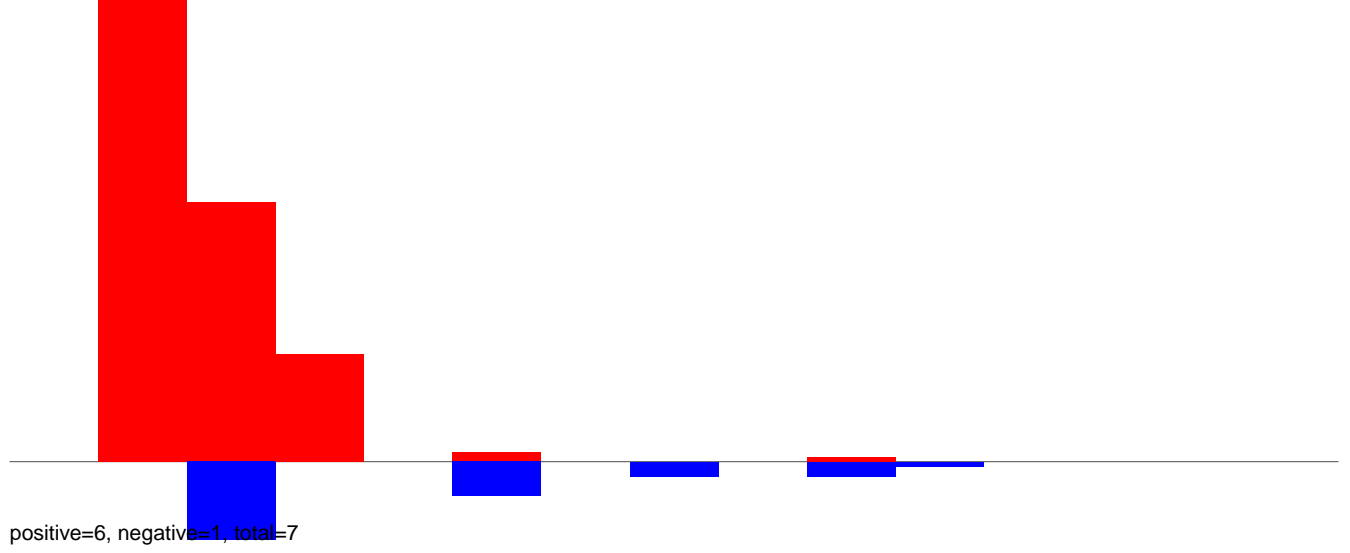
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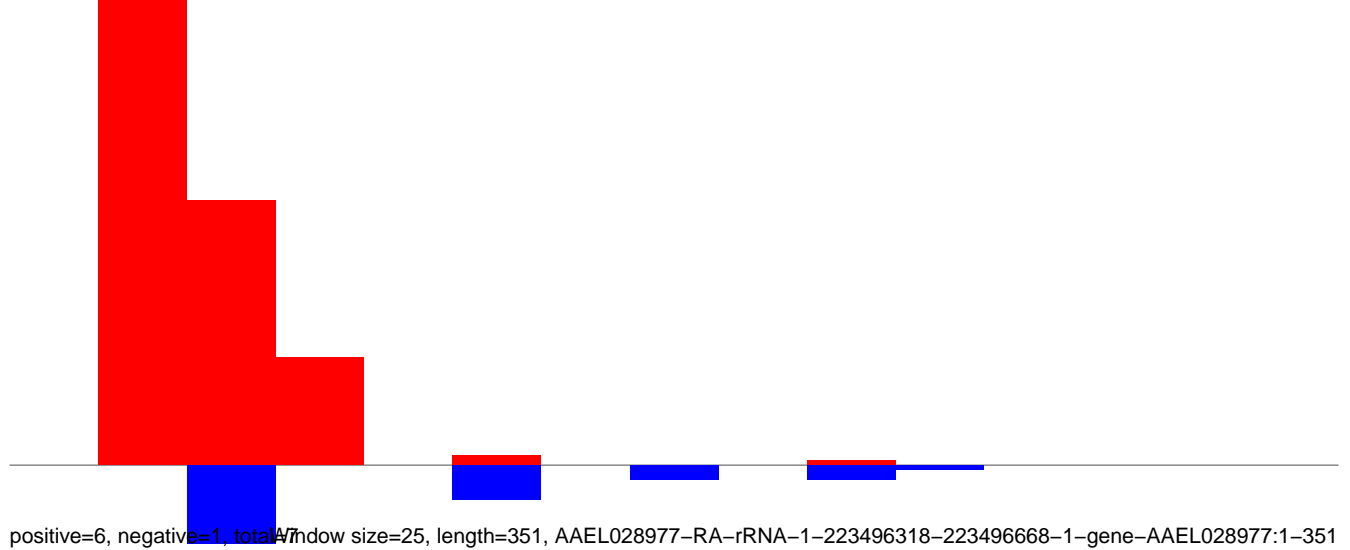
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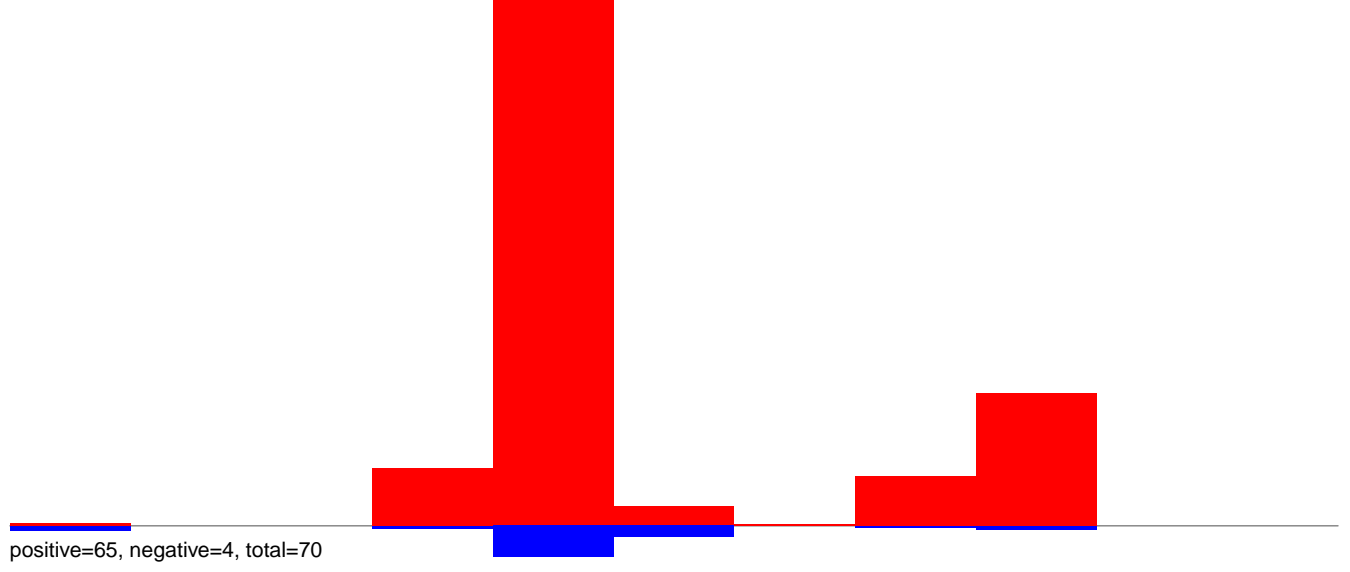


AeAeg_Aag2.TC.Dmem.rep

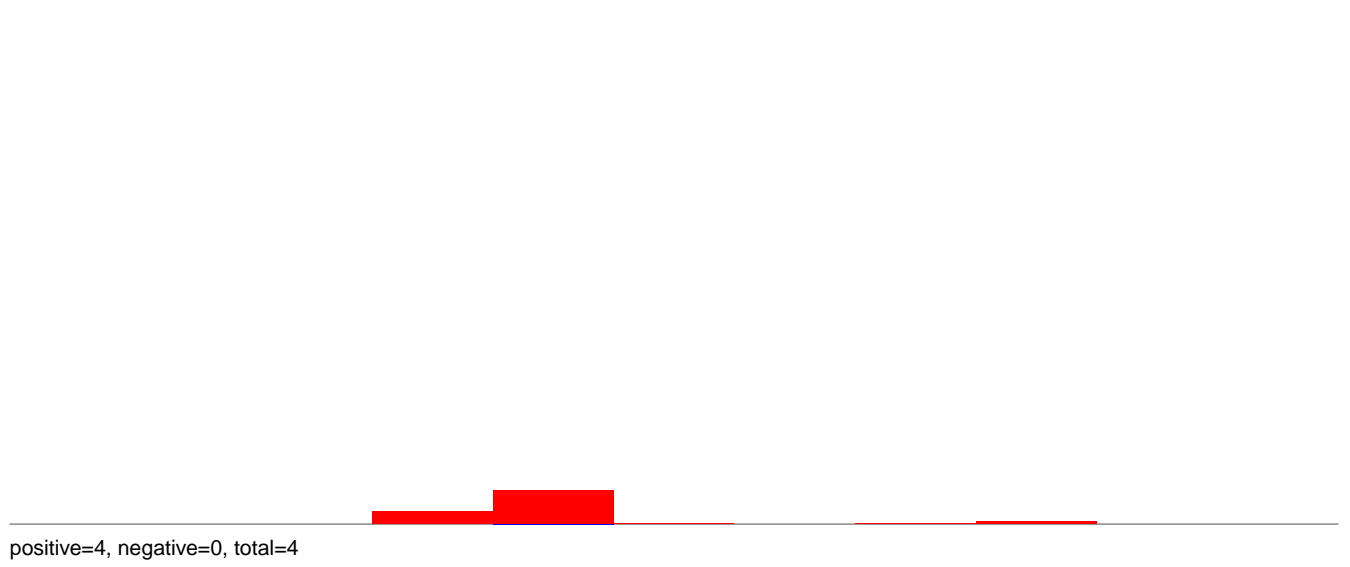


Window size=25, length=351, AAEL028977-RA-rRNA-1-223496318-223496668-1-gene-AAEL028977:1-351

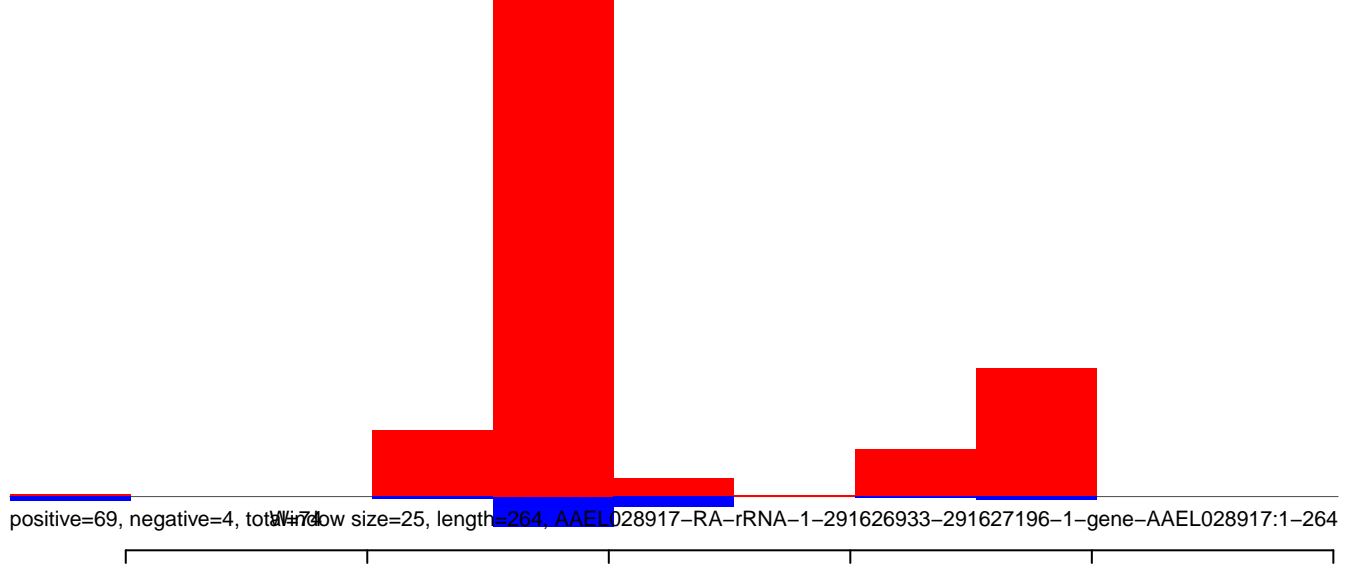
AeAeg_Aag2.TC.Dmem.18_23.rep



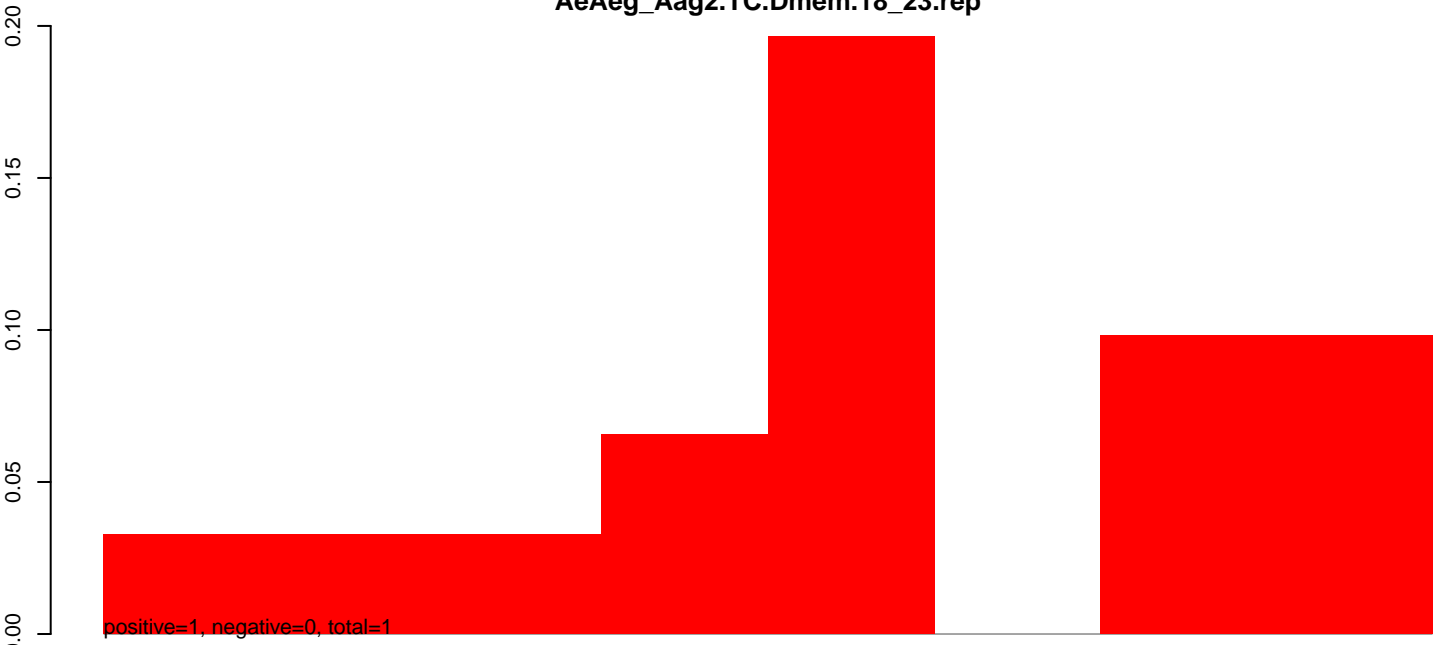
AeAeg_Aag2.TC.Dmem.24_35.rep



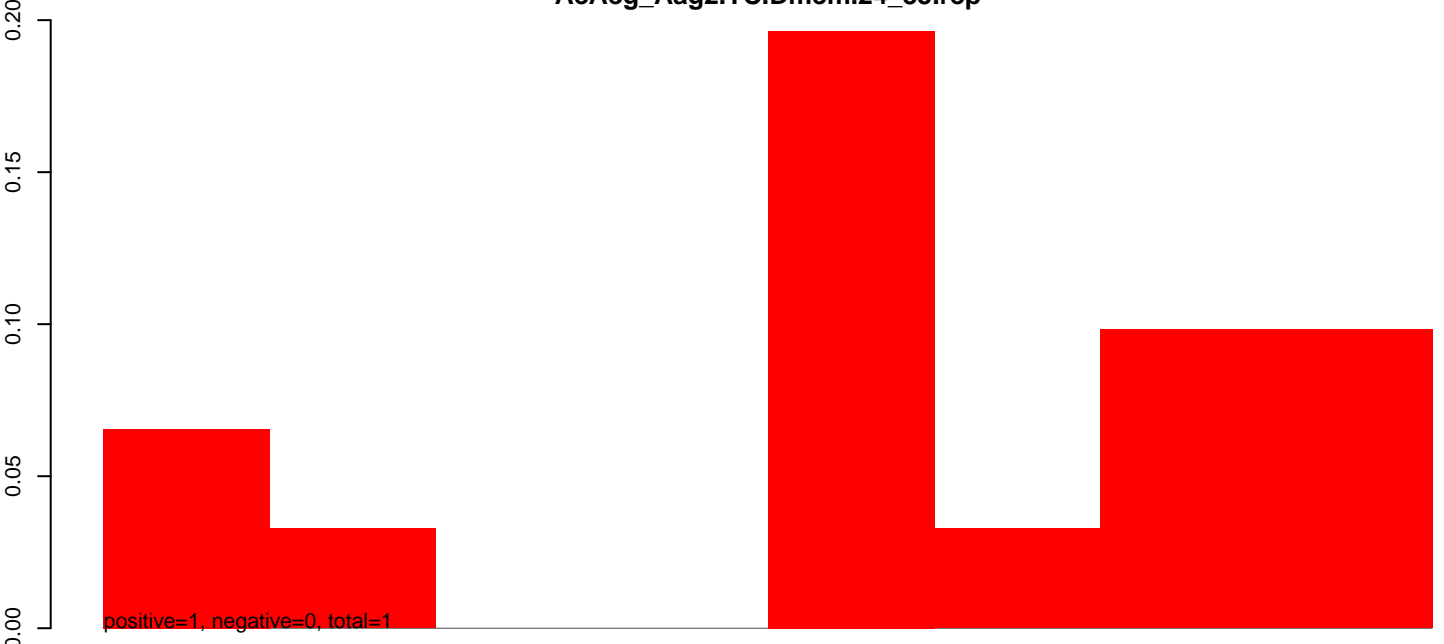
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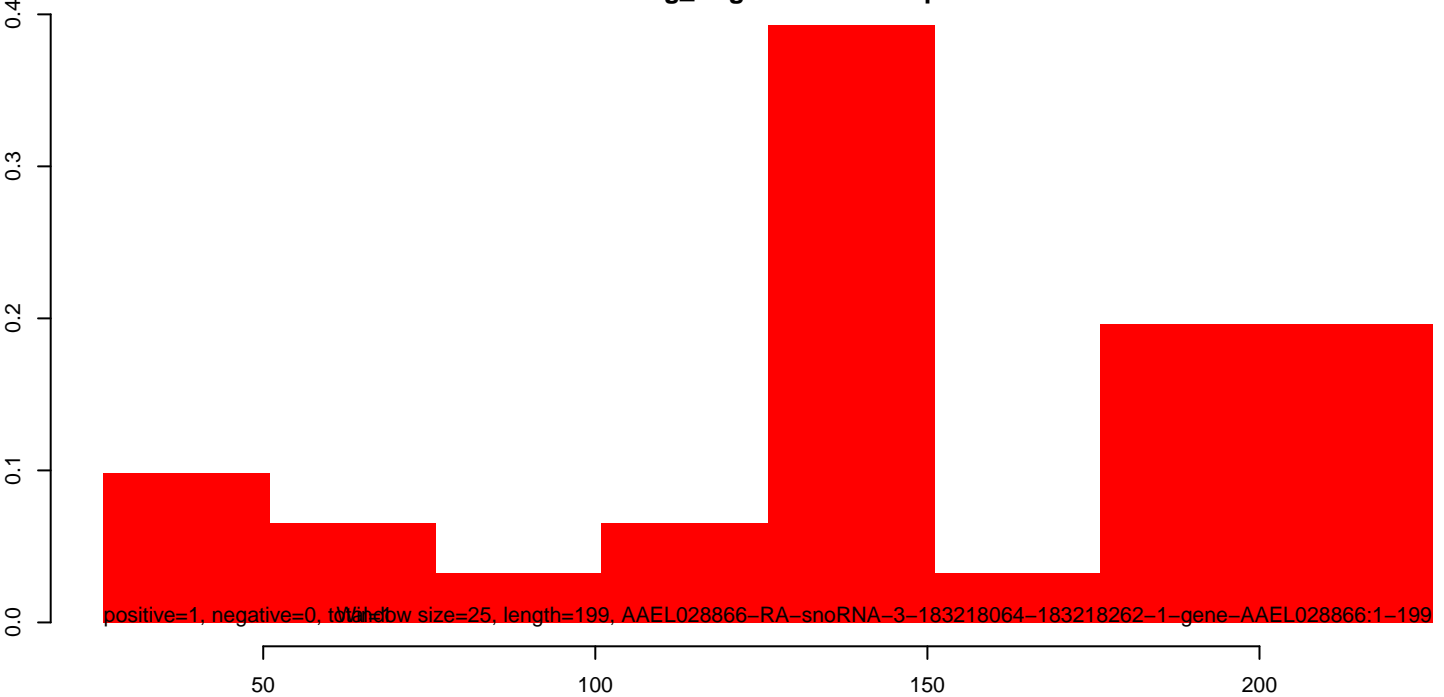
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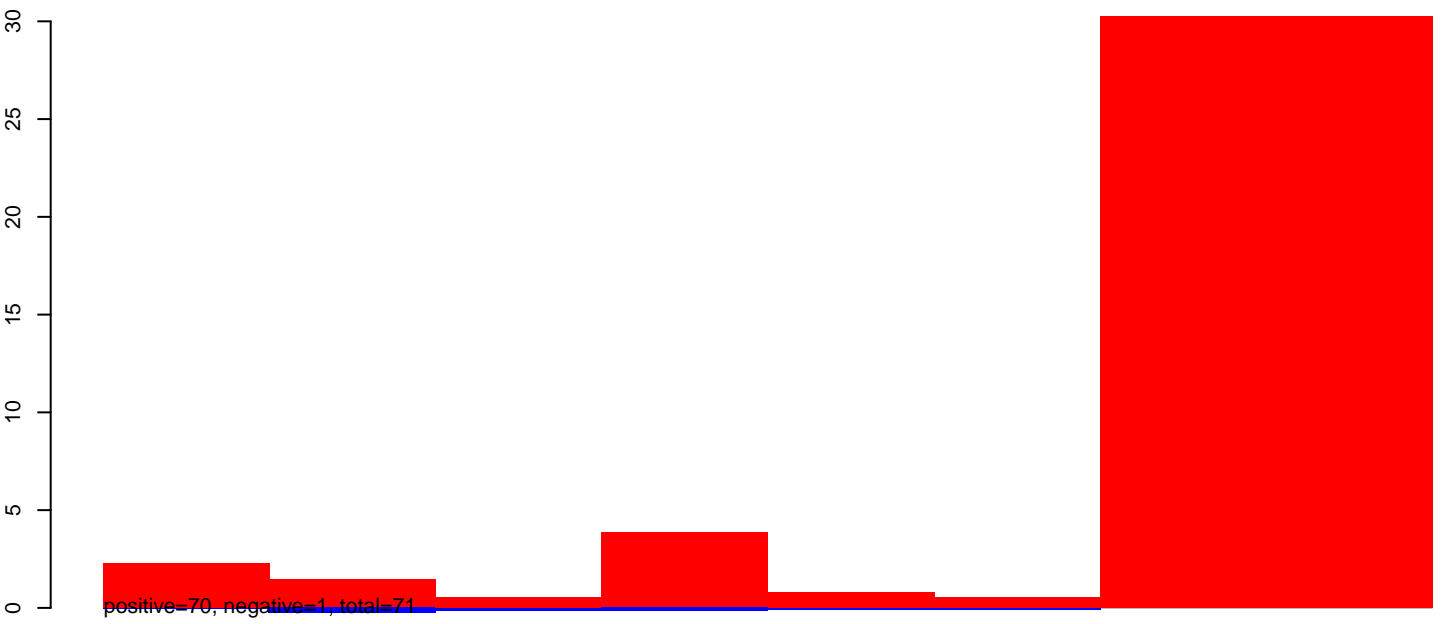
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AeAeg_Aag2.TC.Dmem.rep



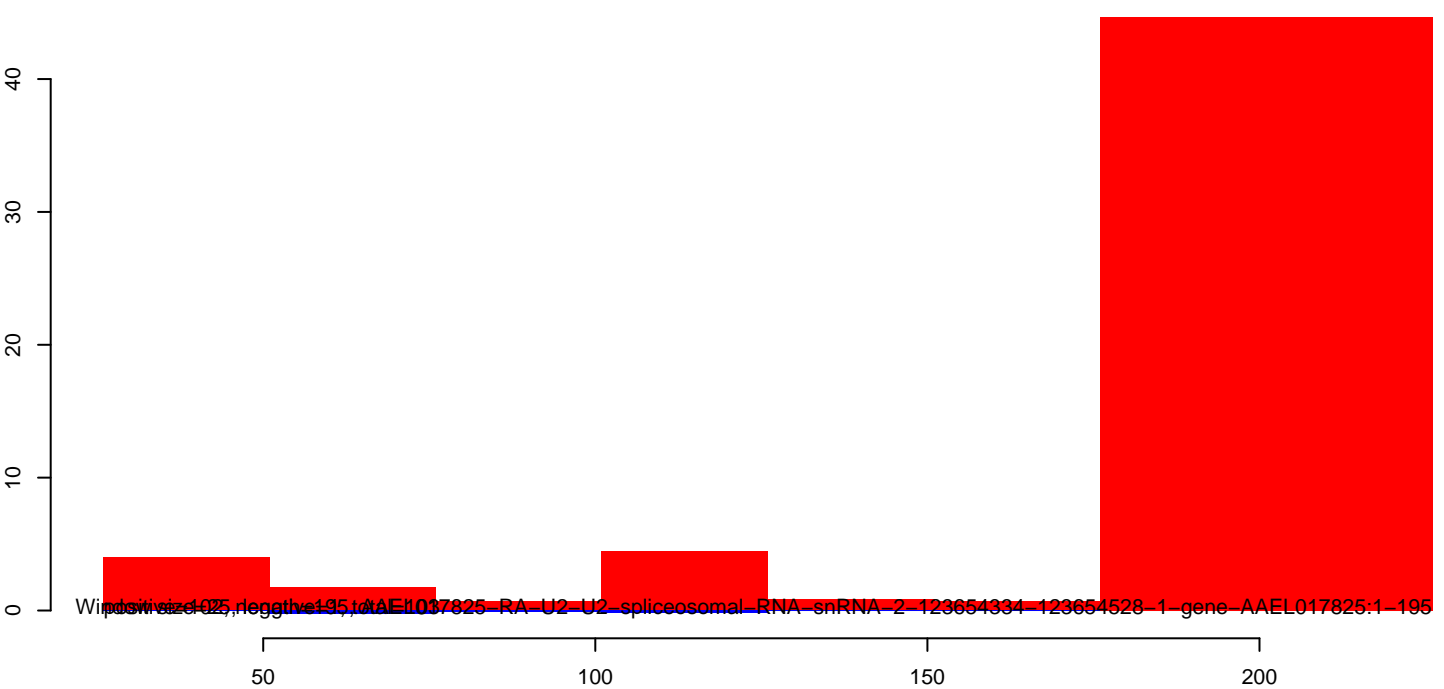
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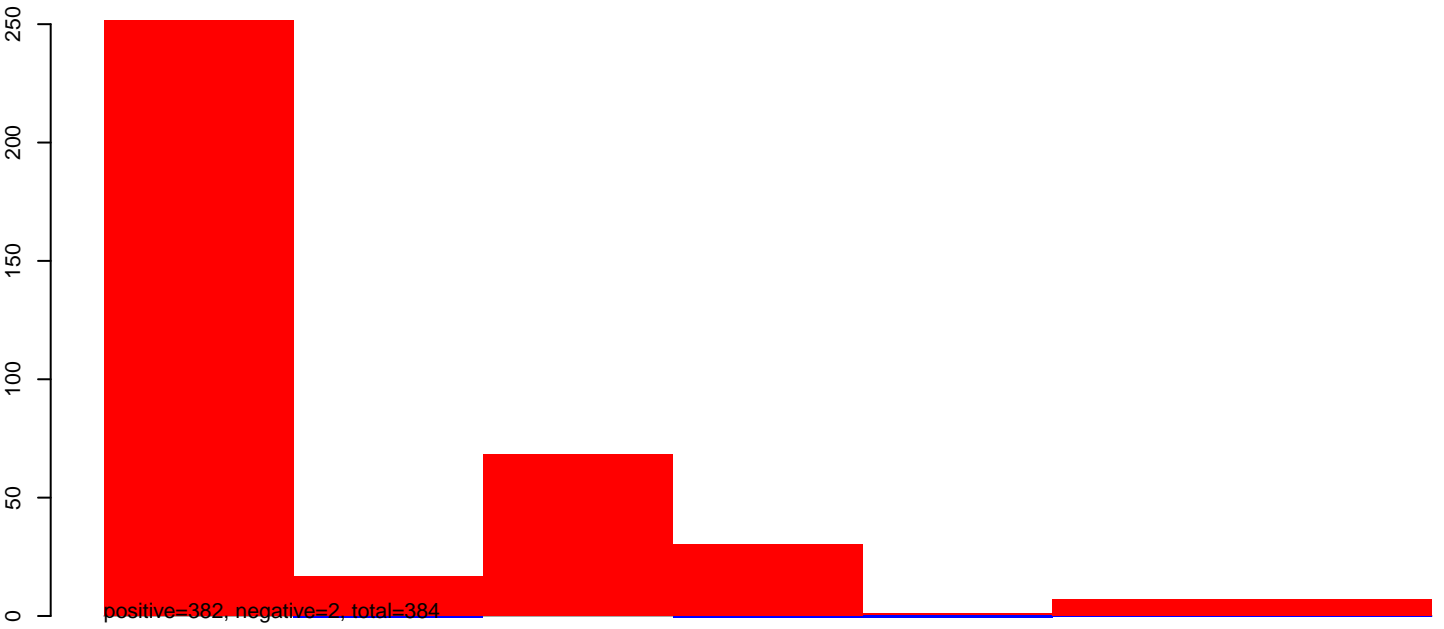
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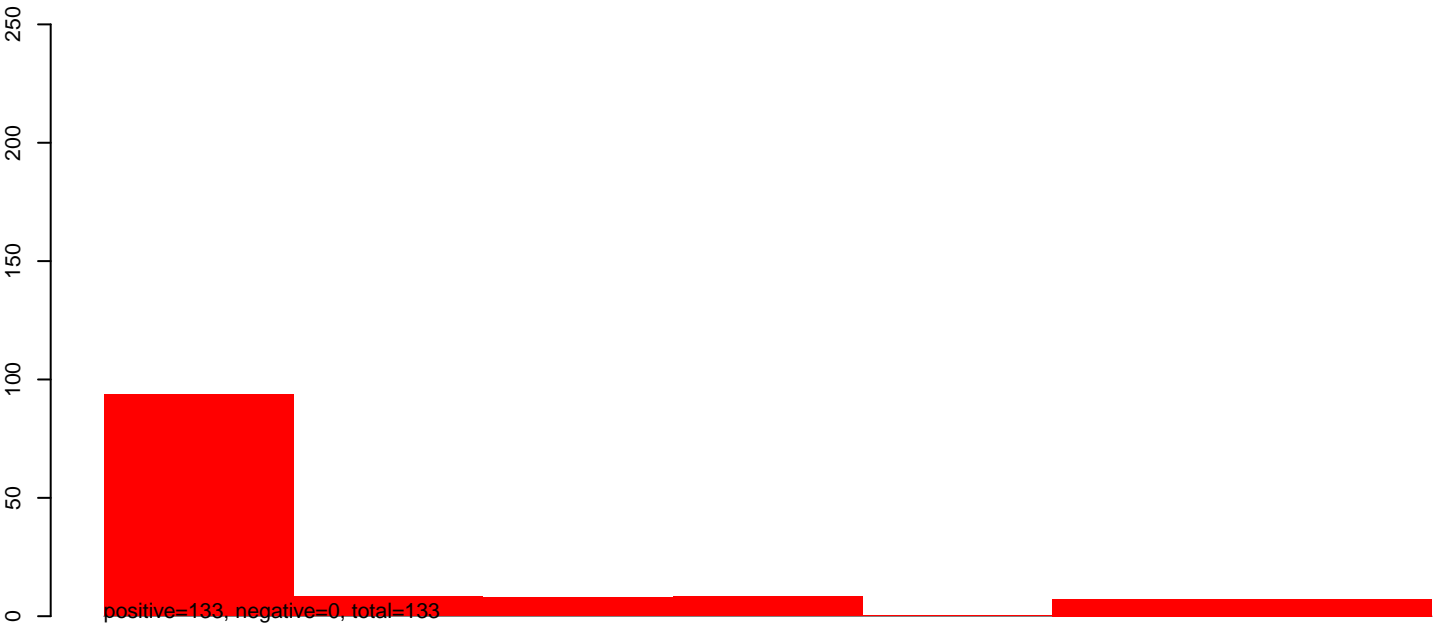
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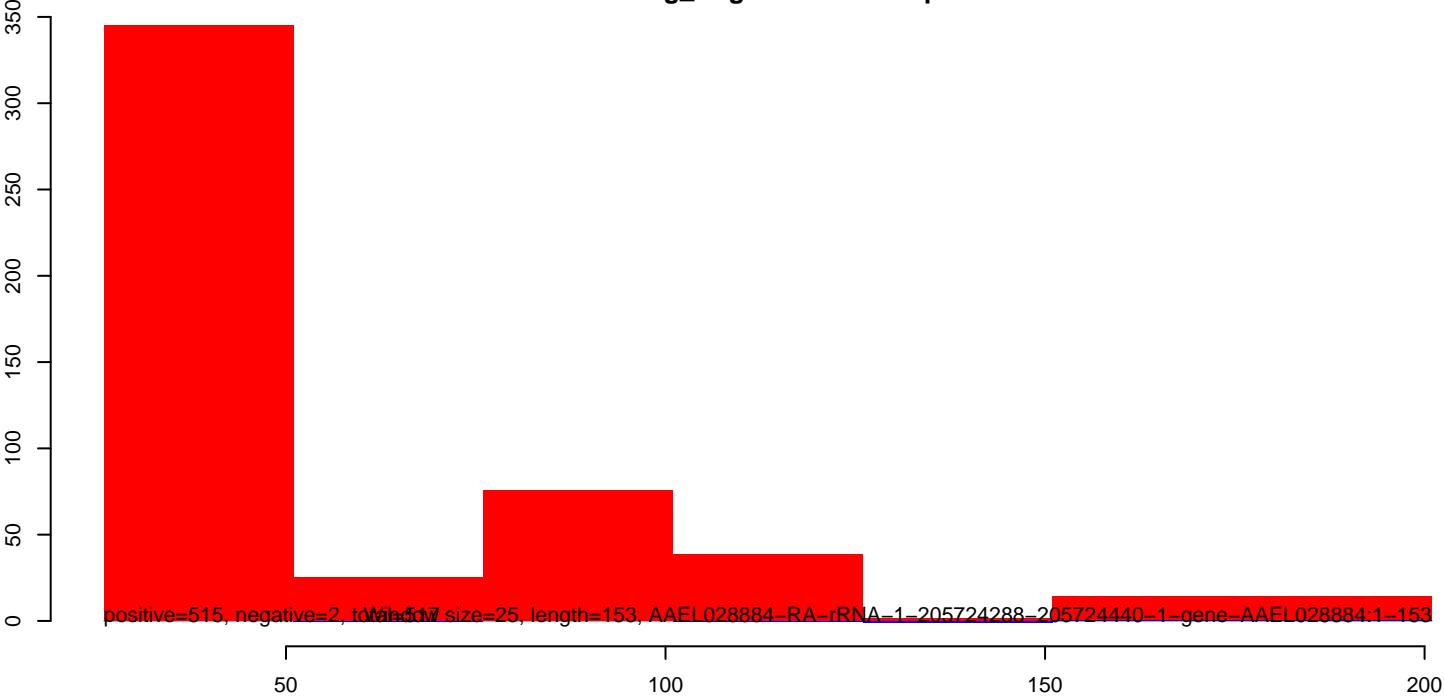
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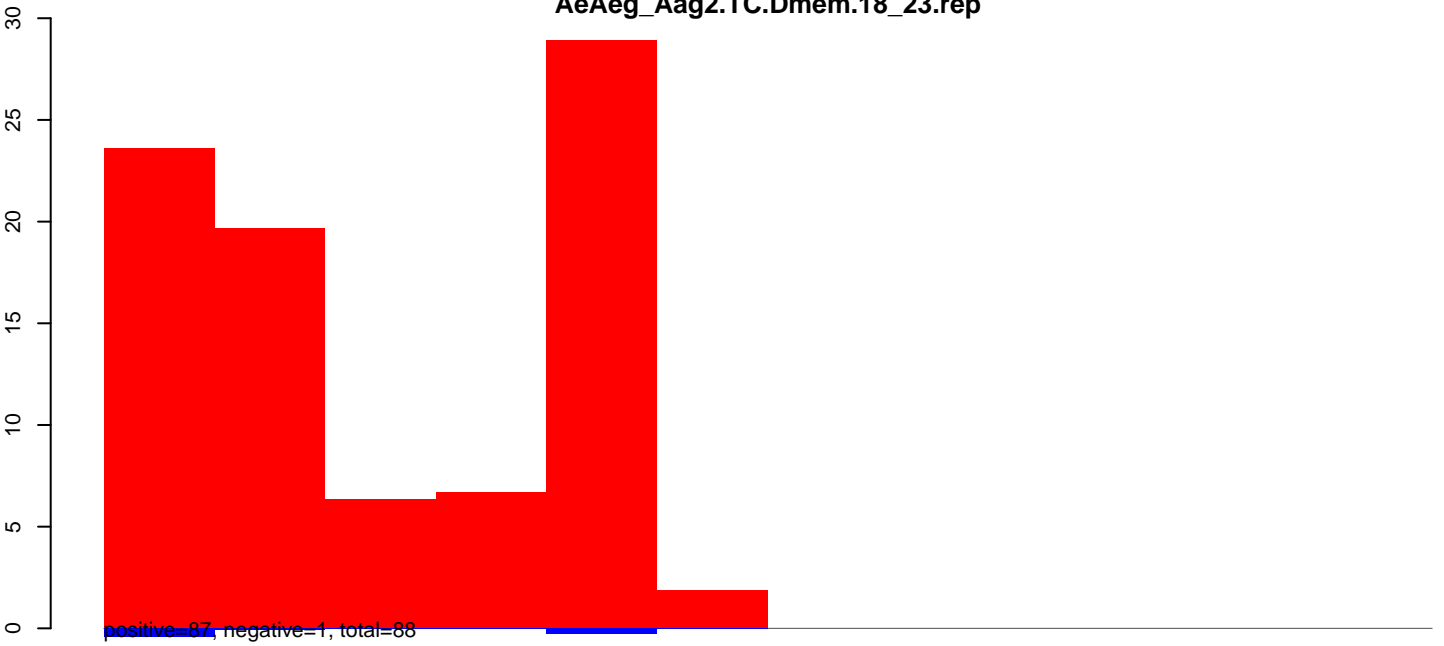
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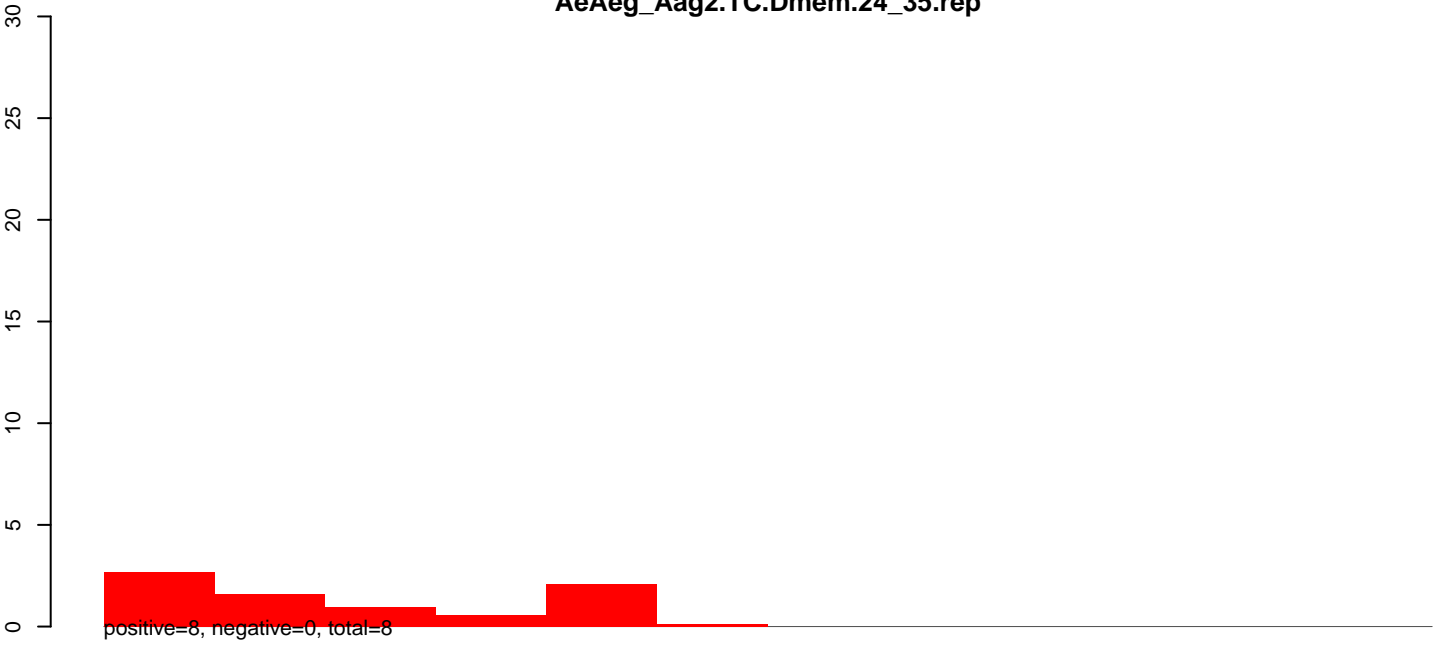
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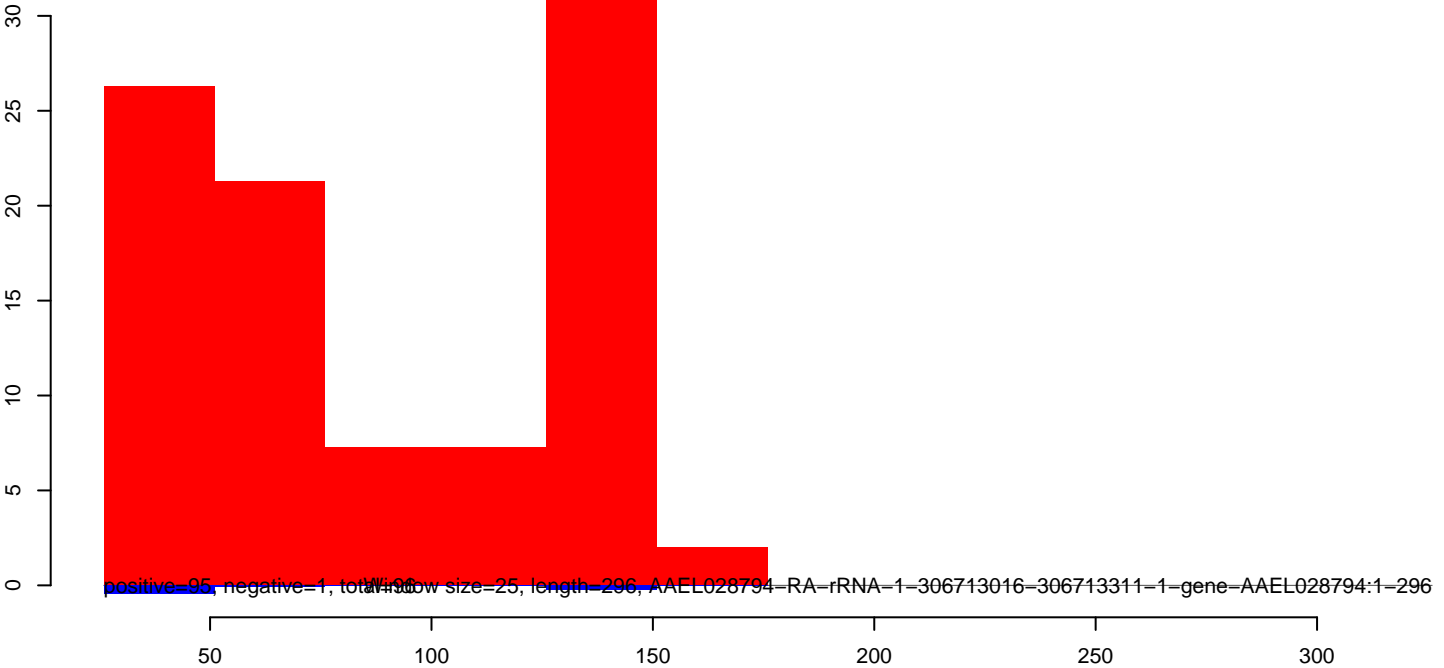
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AeAeg_Aag2.TC.Dmem.24_35.rep

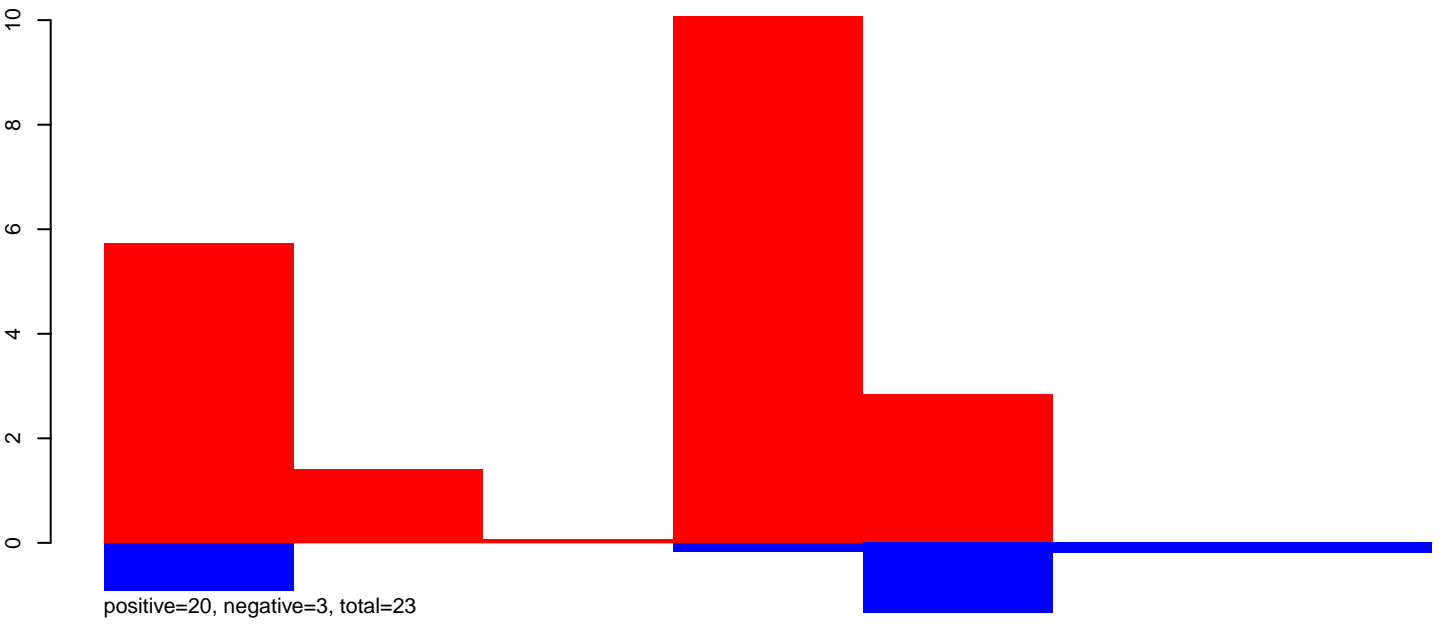


AeAeg_Aag2.TC.Dmem.rep

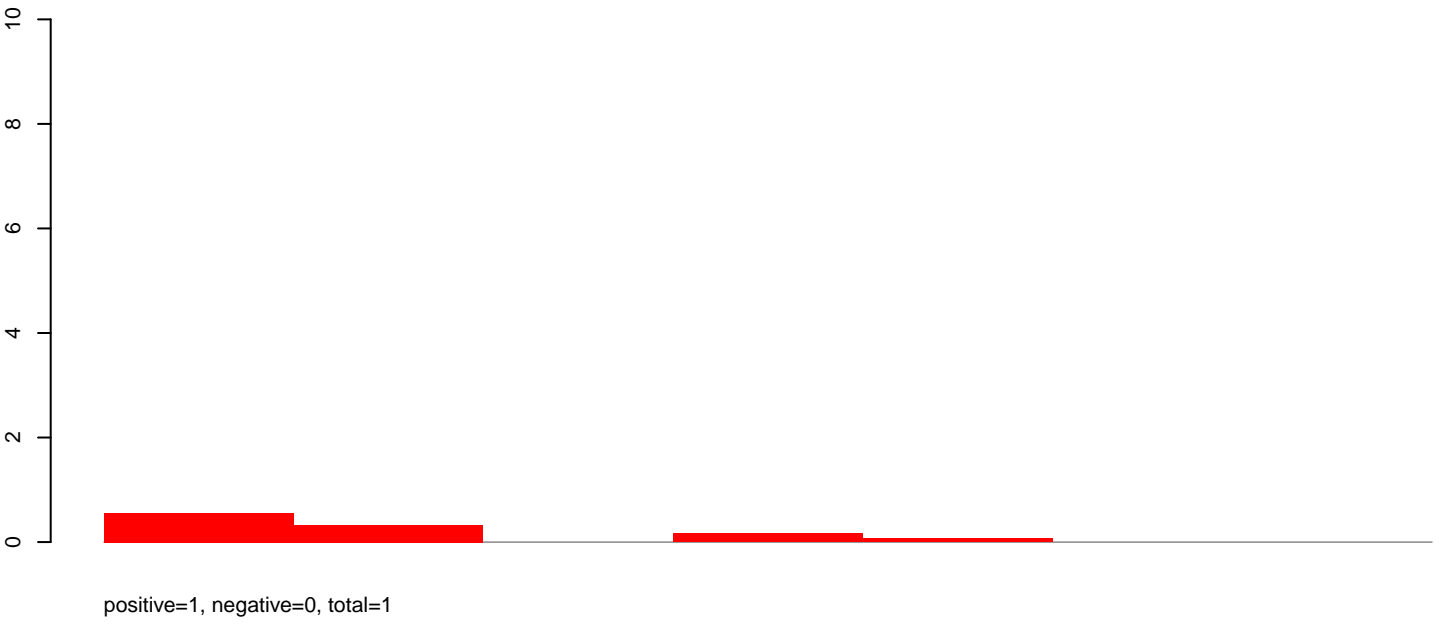


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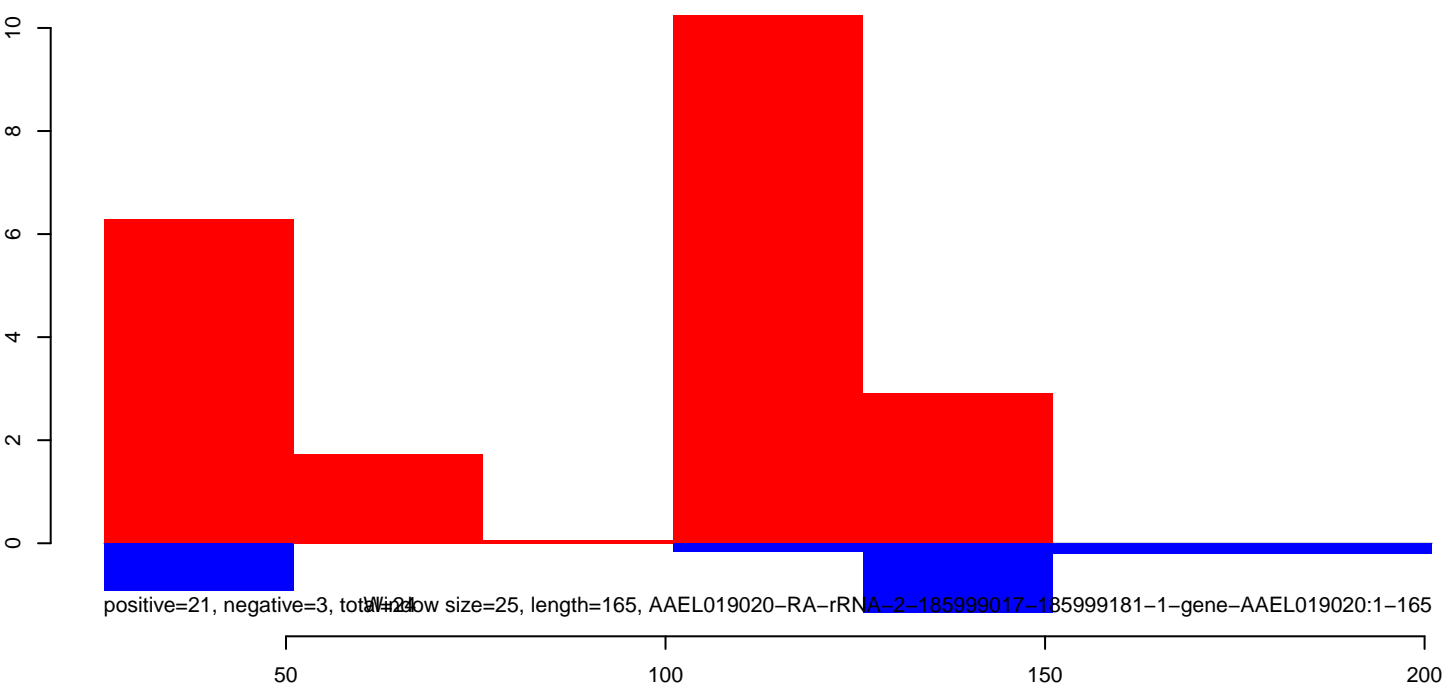
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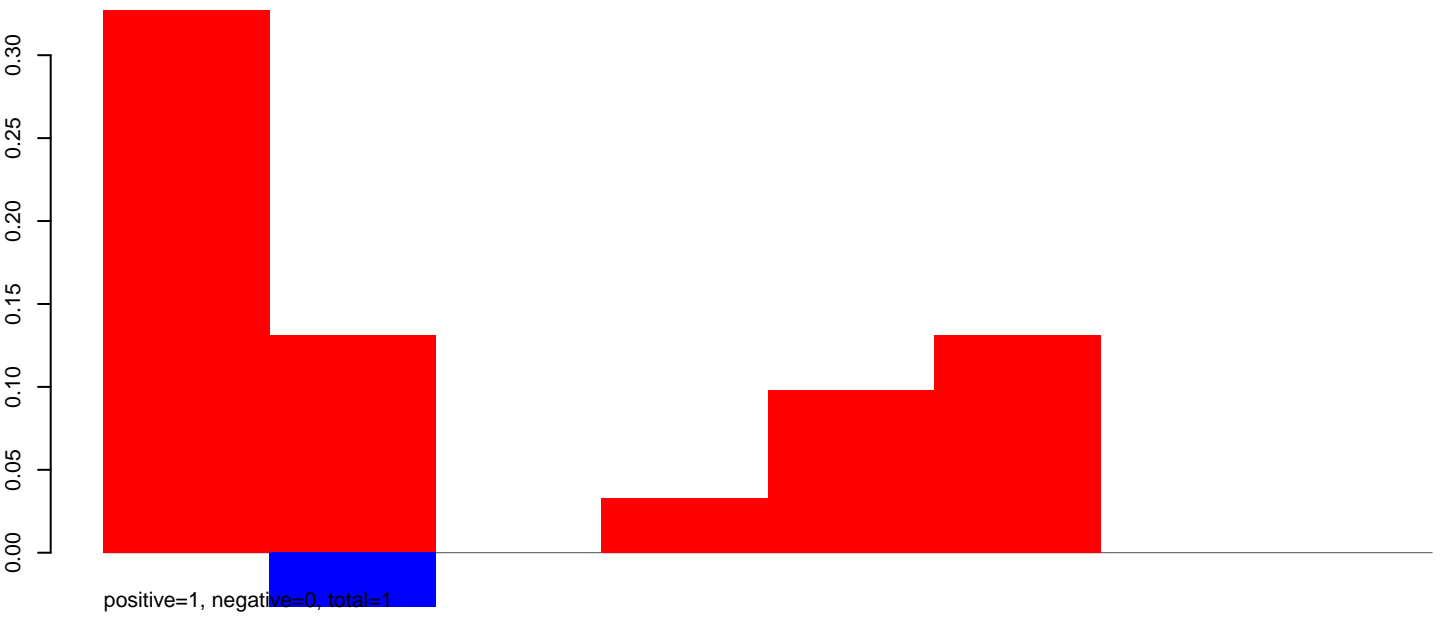
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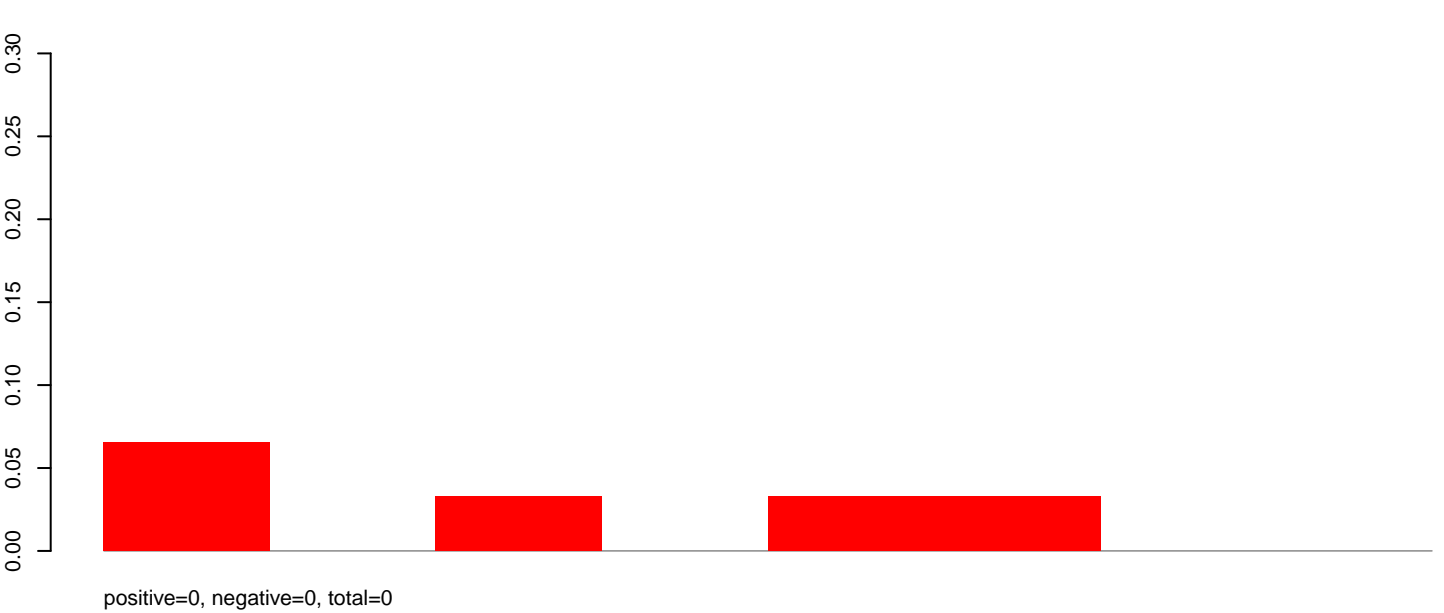
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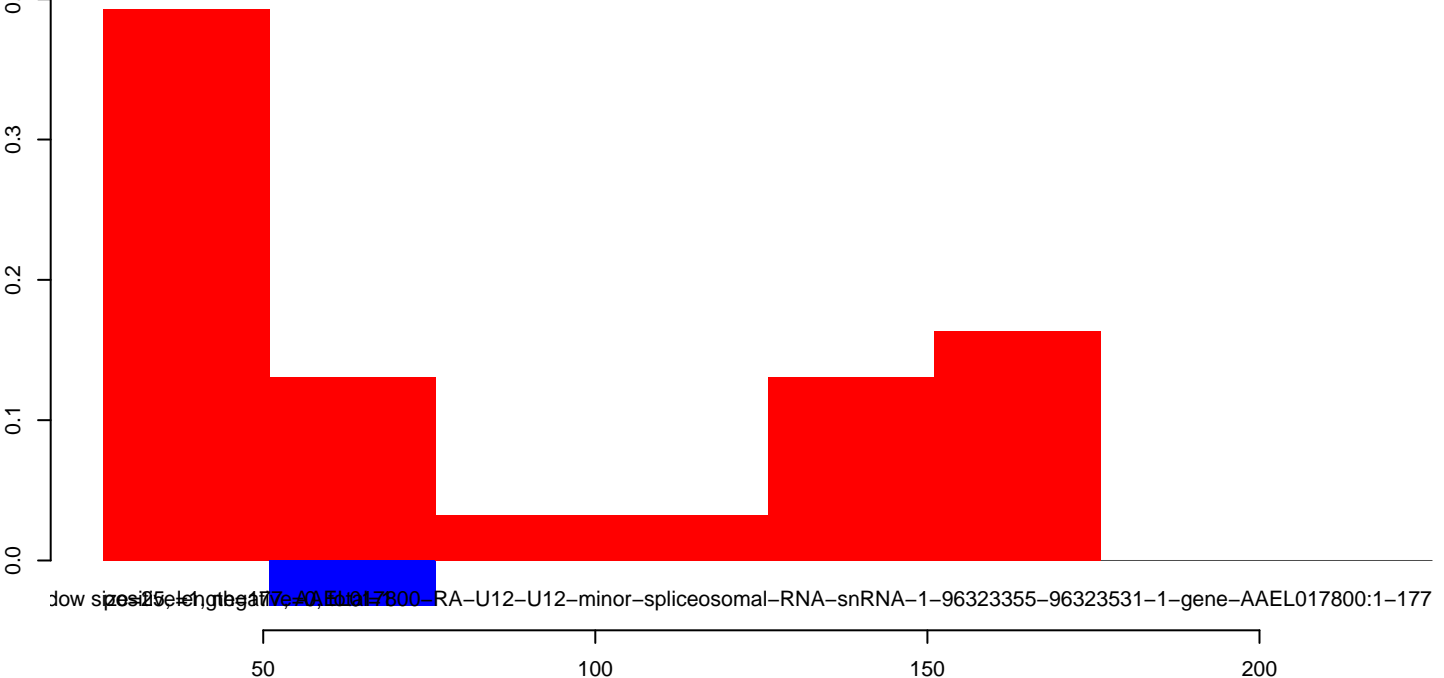
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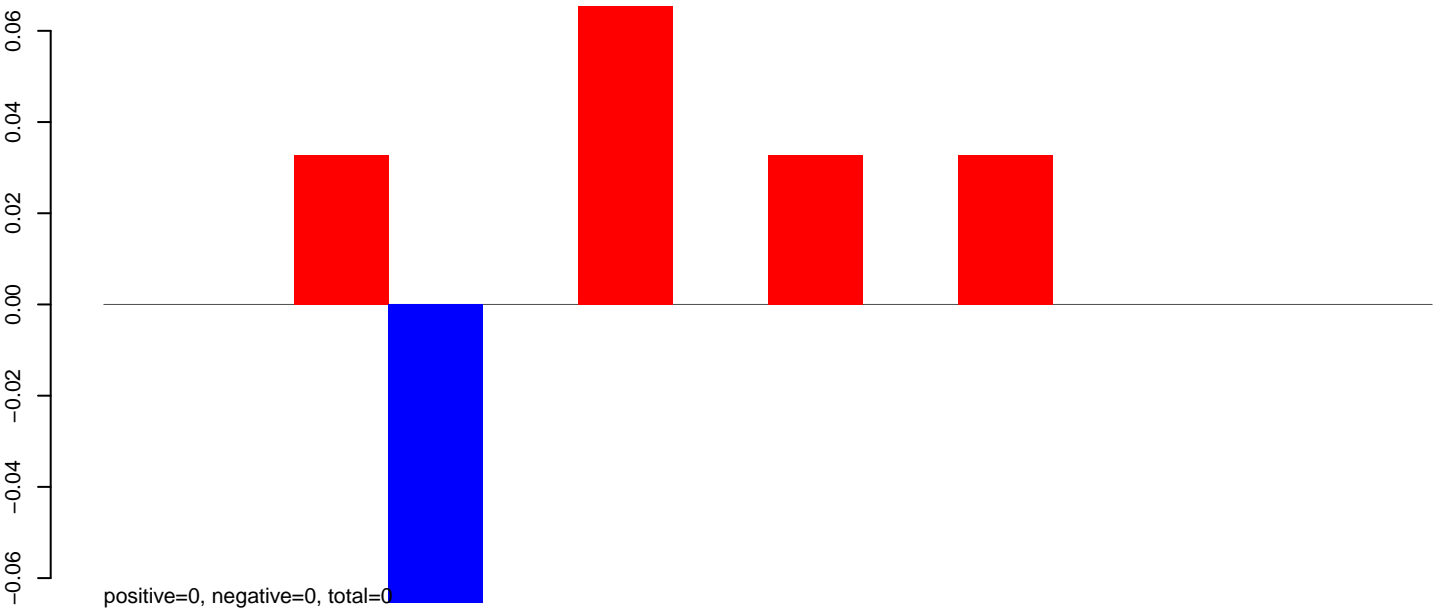
AeAeg_Aag2.TC.Dmem.24_35.rep



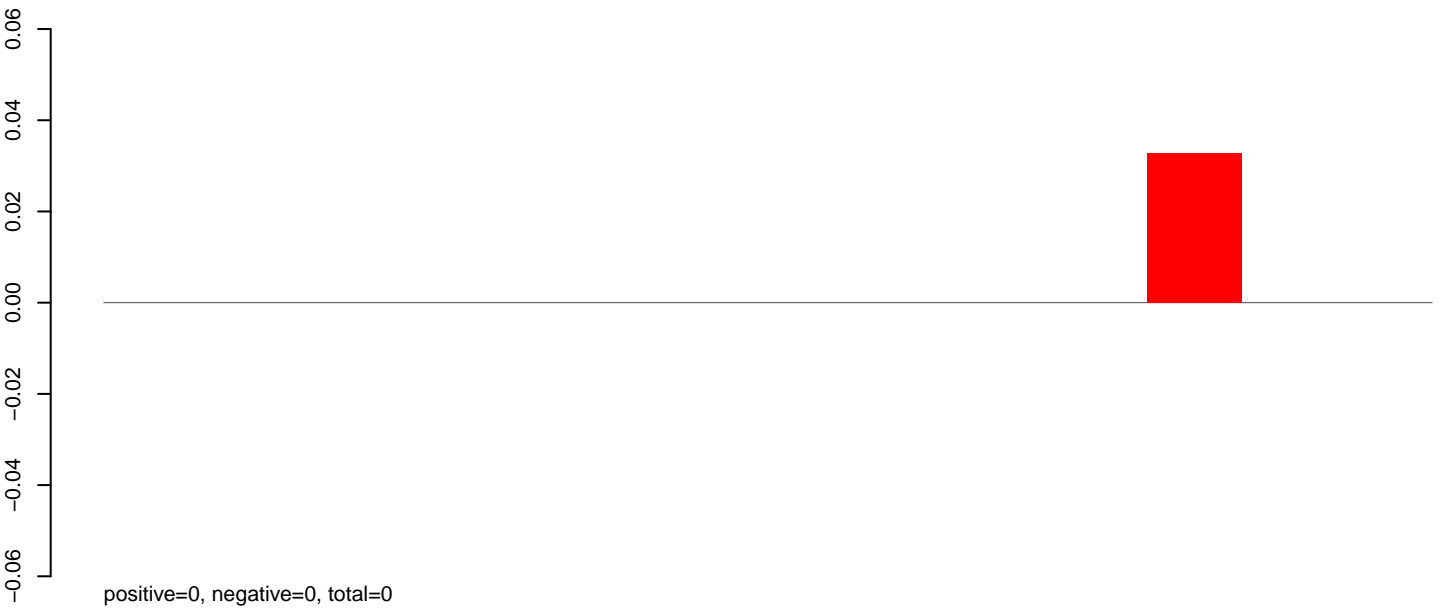
AeAeg_Aag2.TC.Dmem.rep



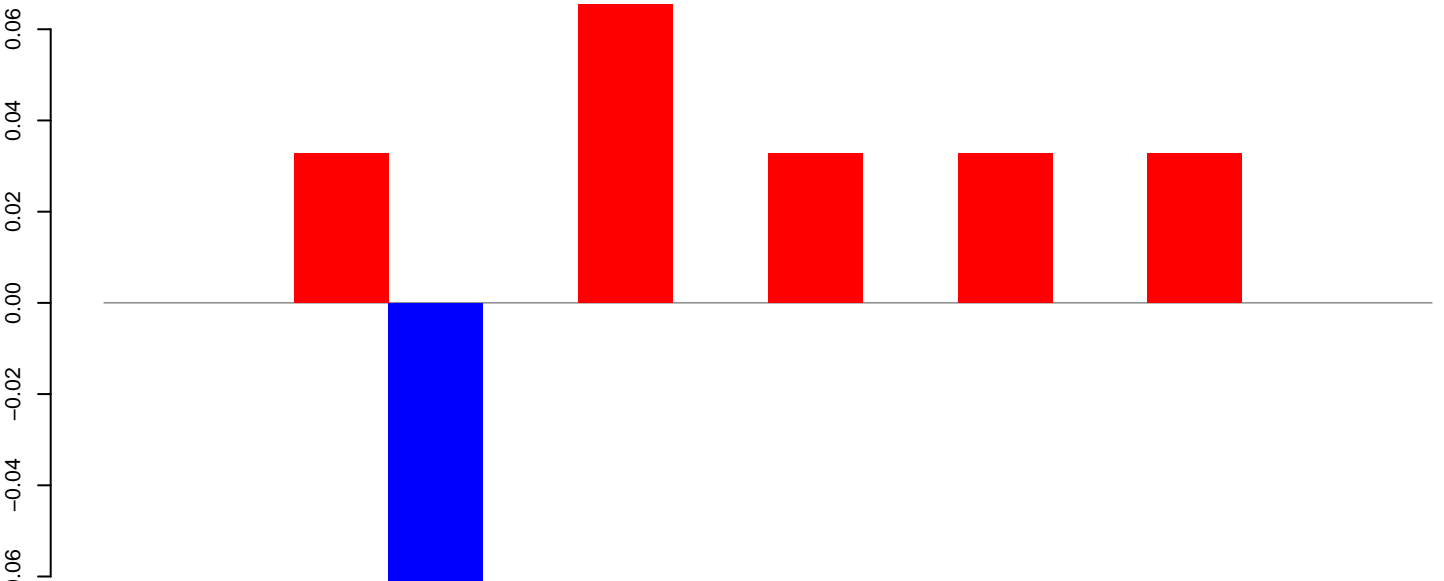
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AeAeg_Aag2.TC.Dmem.24_35.rep



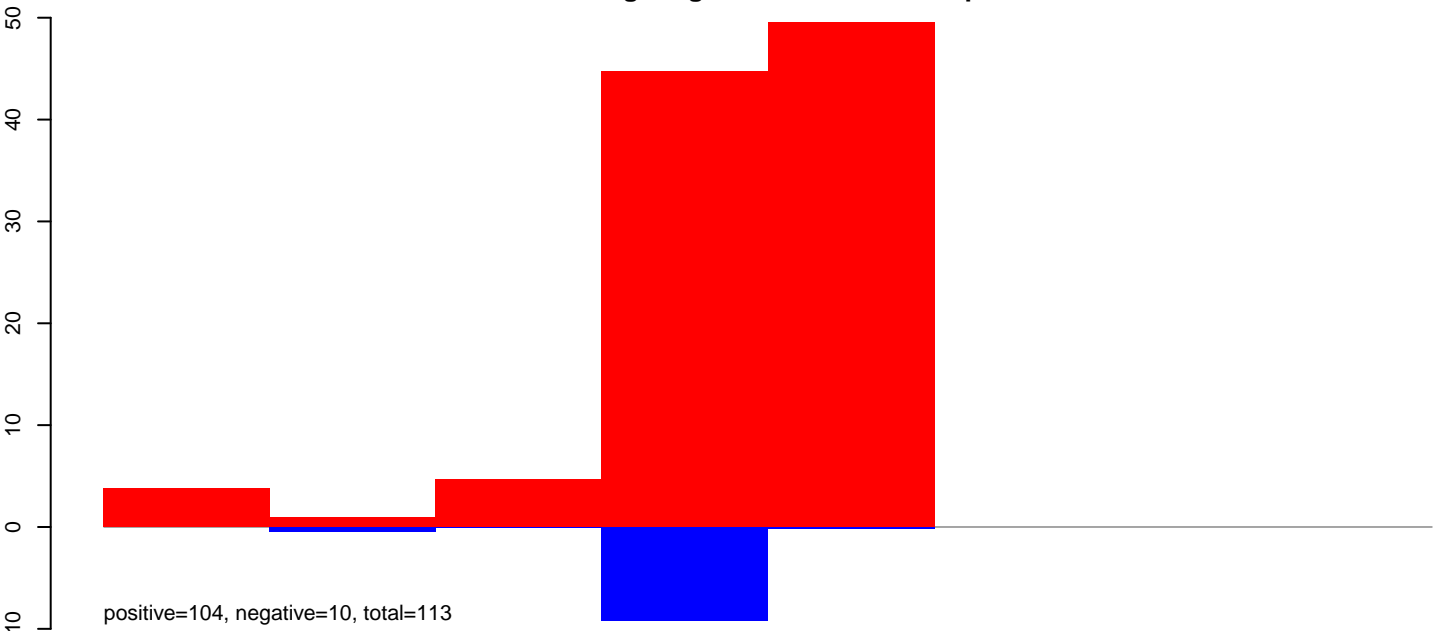
AeAeg_Aag2.TC.Dmem.rep



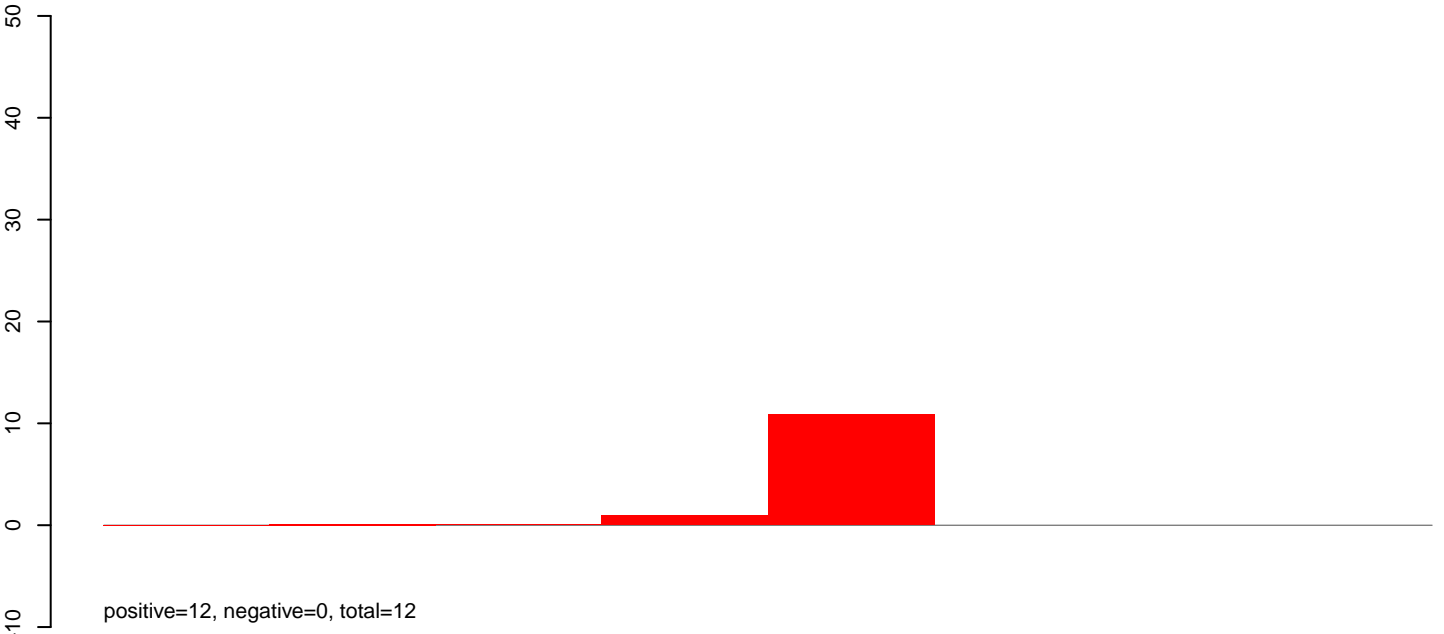
3=25, positive=30, negative=17, total=47
AeAeg_Aag2.TC.Dmem.rep - RNaseP_nuc-Nuclear-RNase-P-RNase_P_RNA-2-285037560-285037897-1-gene-AAEL017685:1-338

50 100 150 200 250 300 350

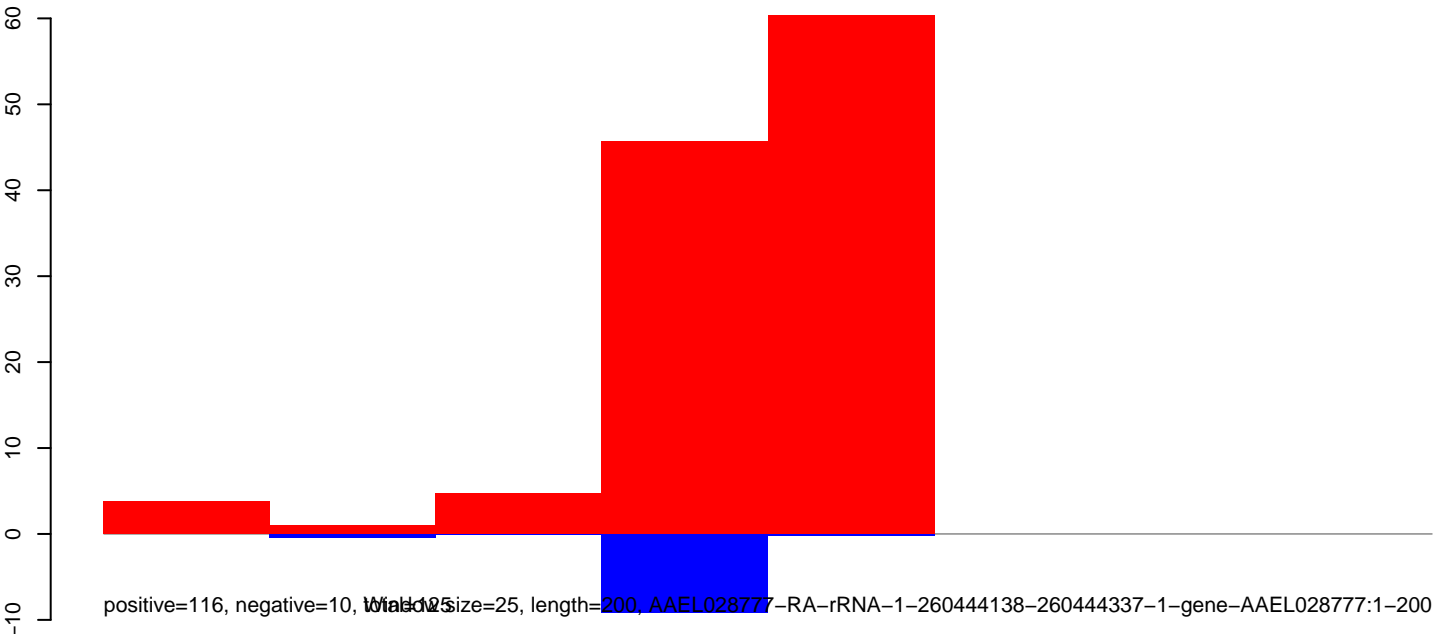
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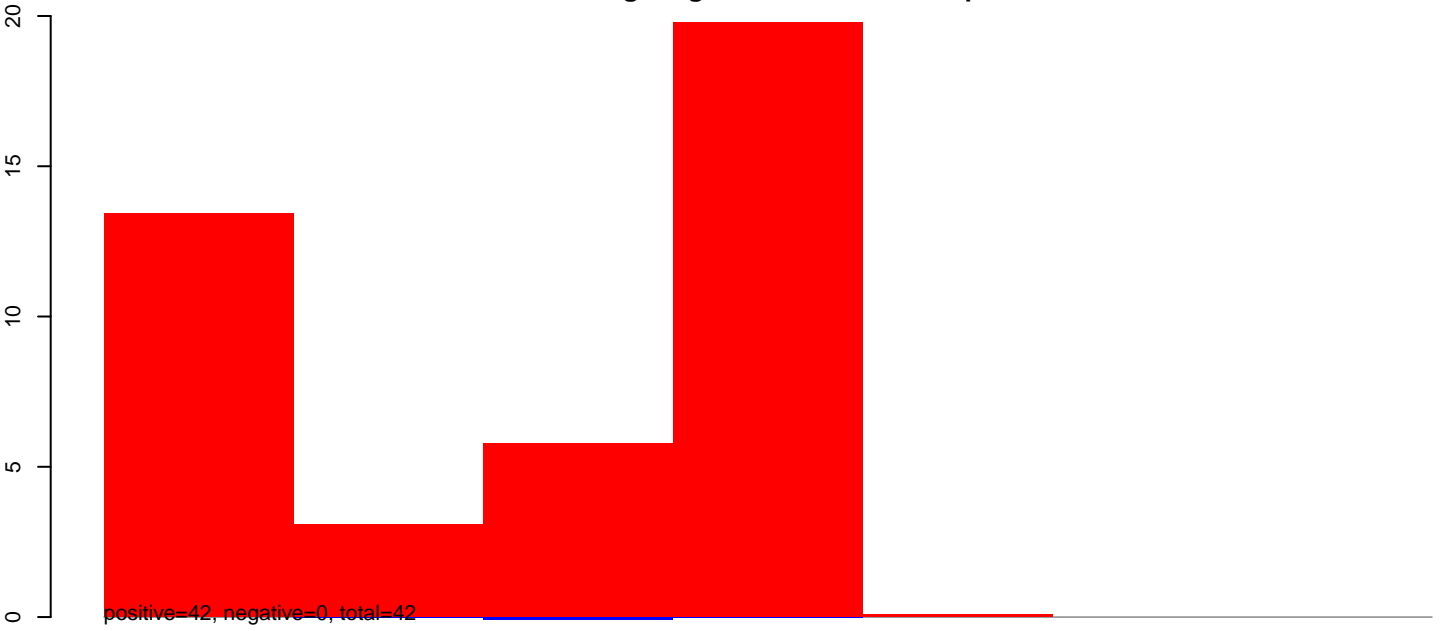
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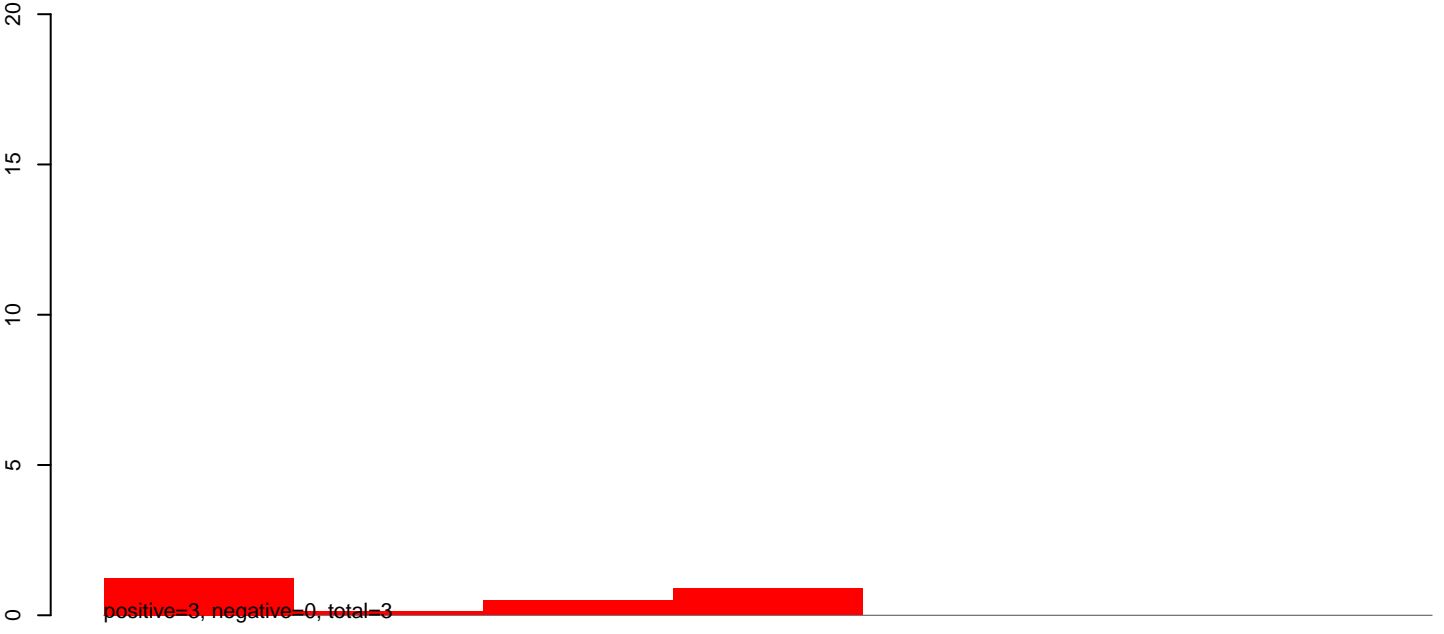
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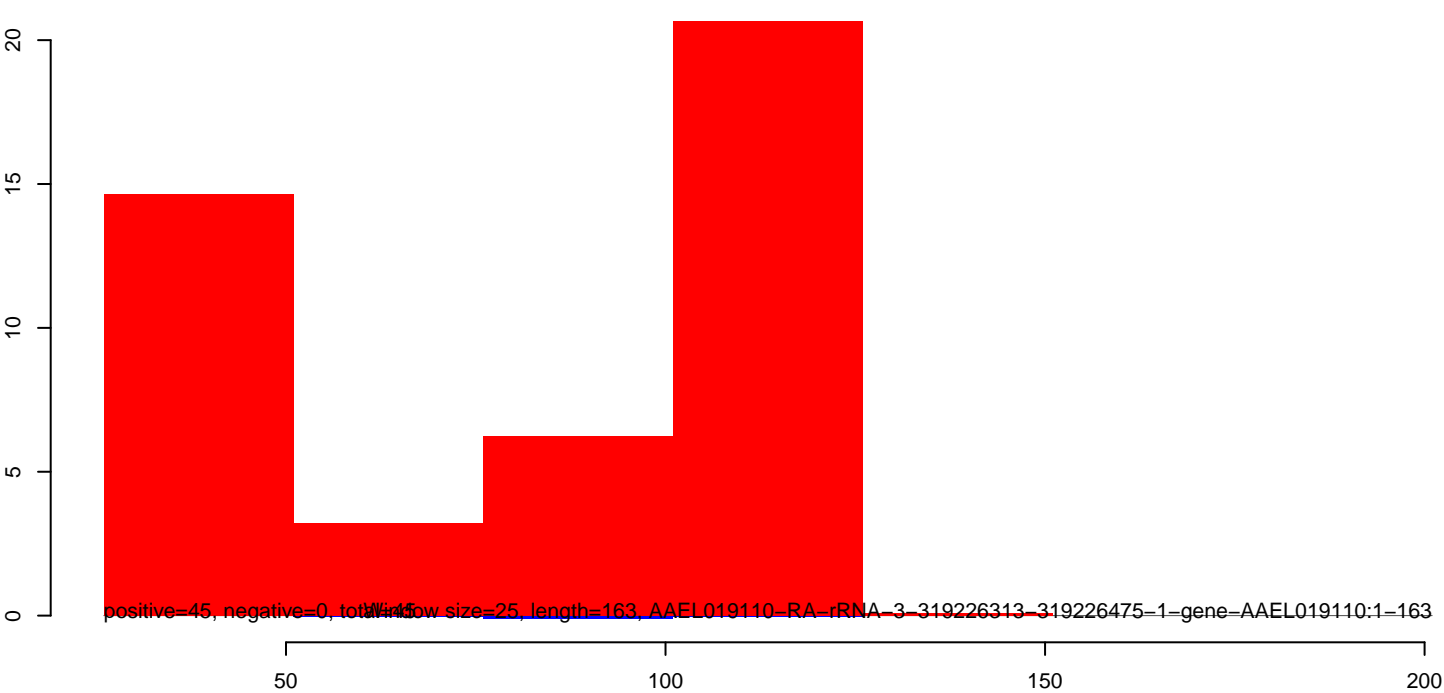
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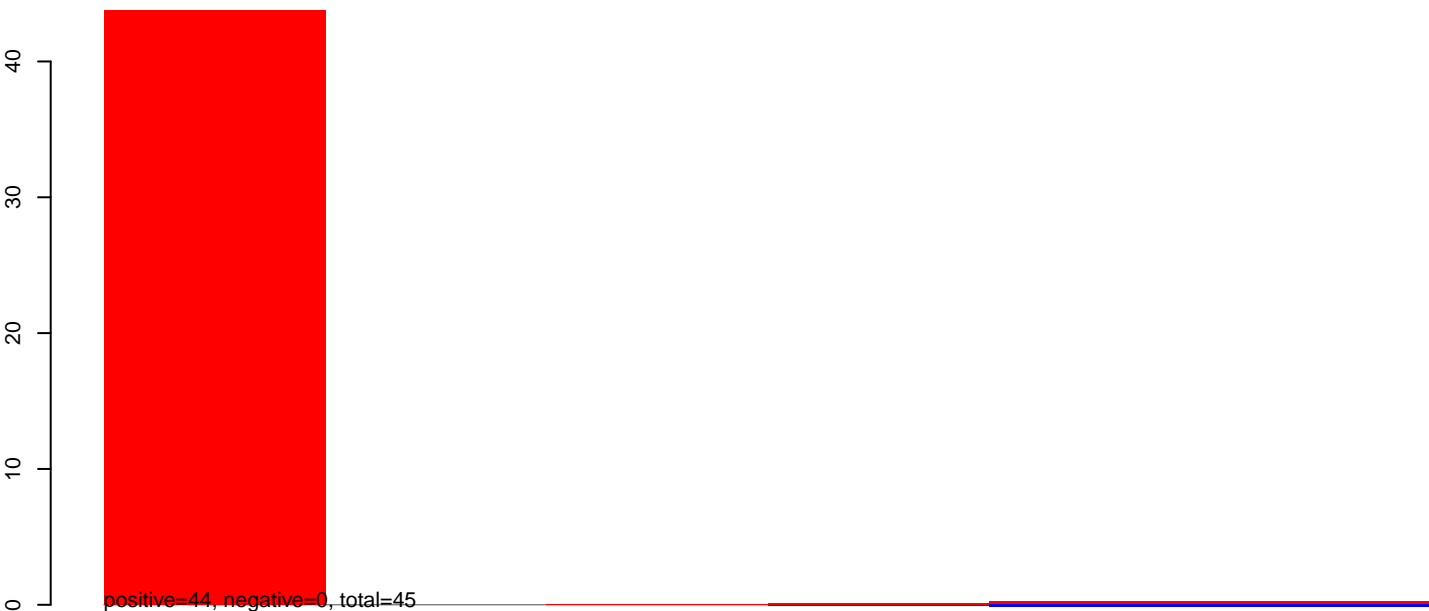
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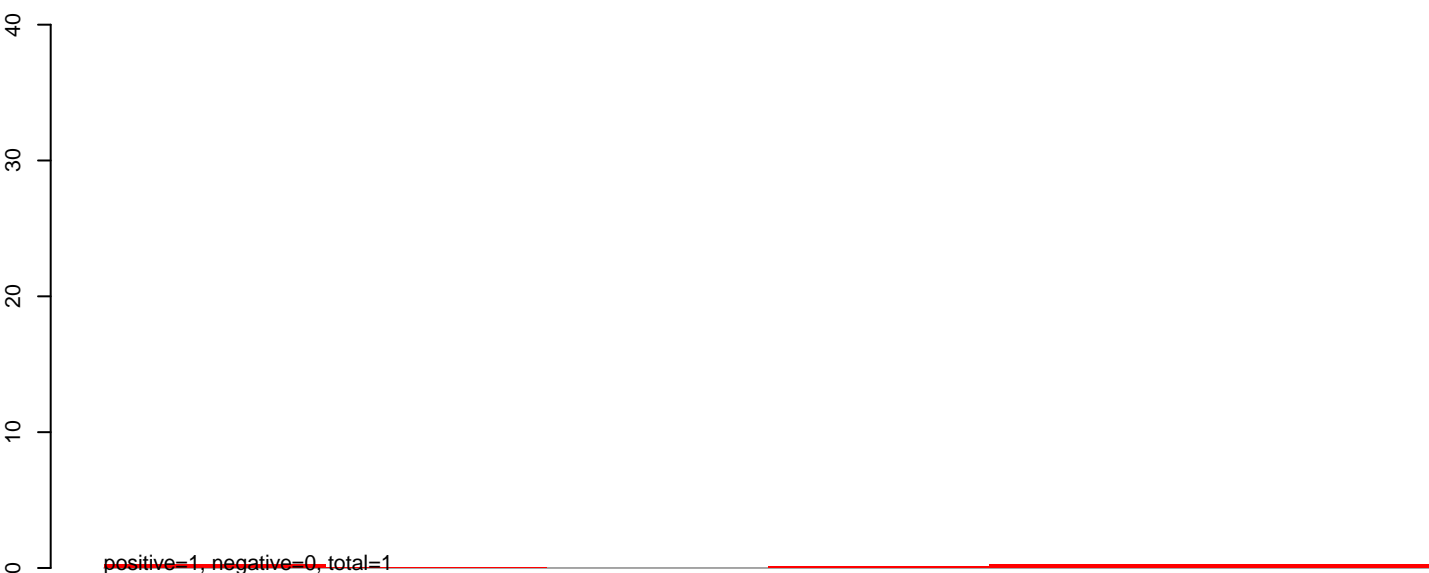
AeAeg_Aag2.TC.Dmem.rep



AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep



AeAeg_Aag2.TC.Dmem.rep



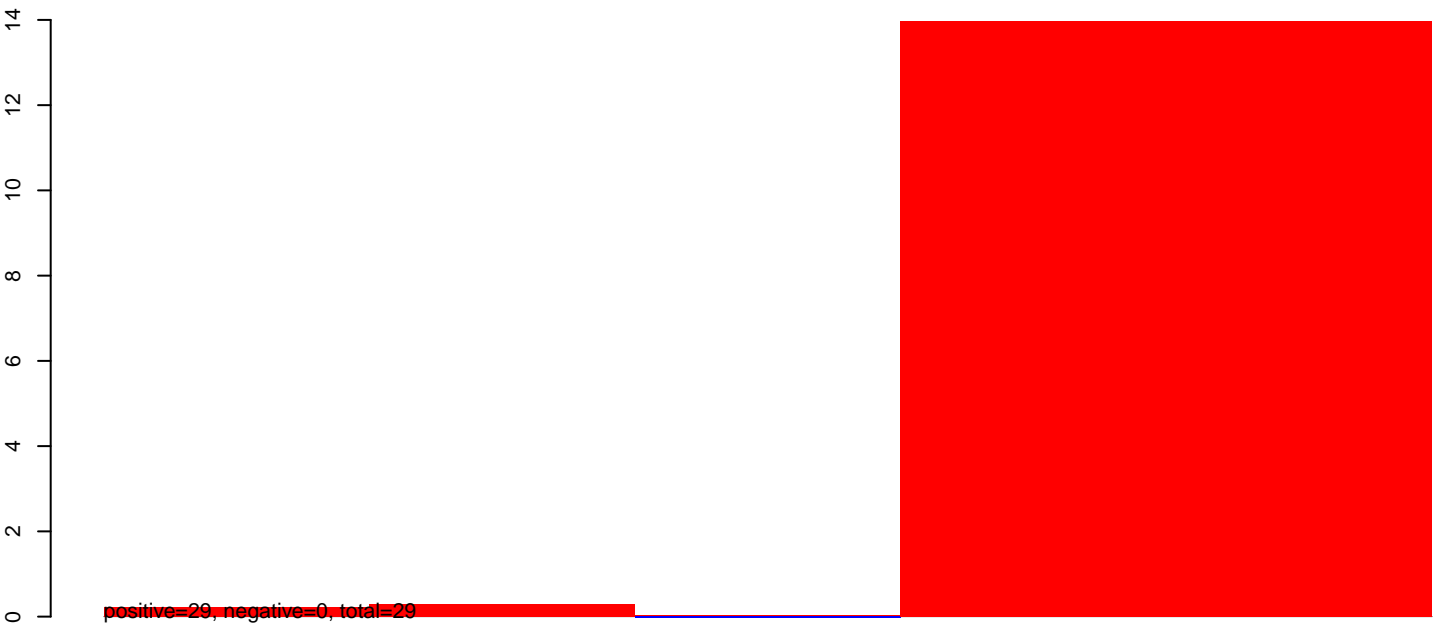
50

100

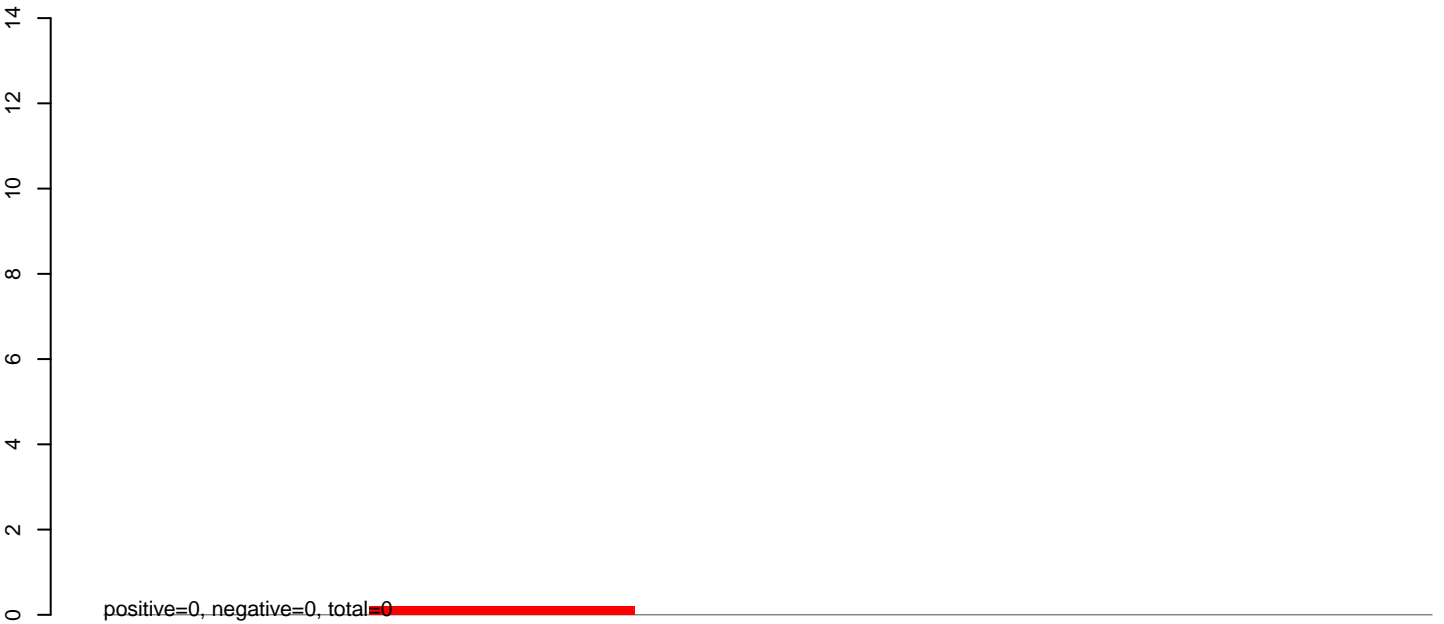
150

Window size=25, length=133, AAEL018844-RA-tRNA-1-208504156-208504288-1 gene-AAEL018844.1-133

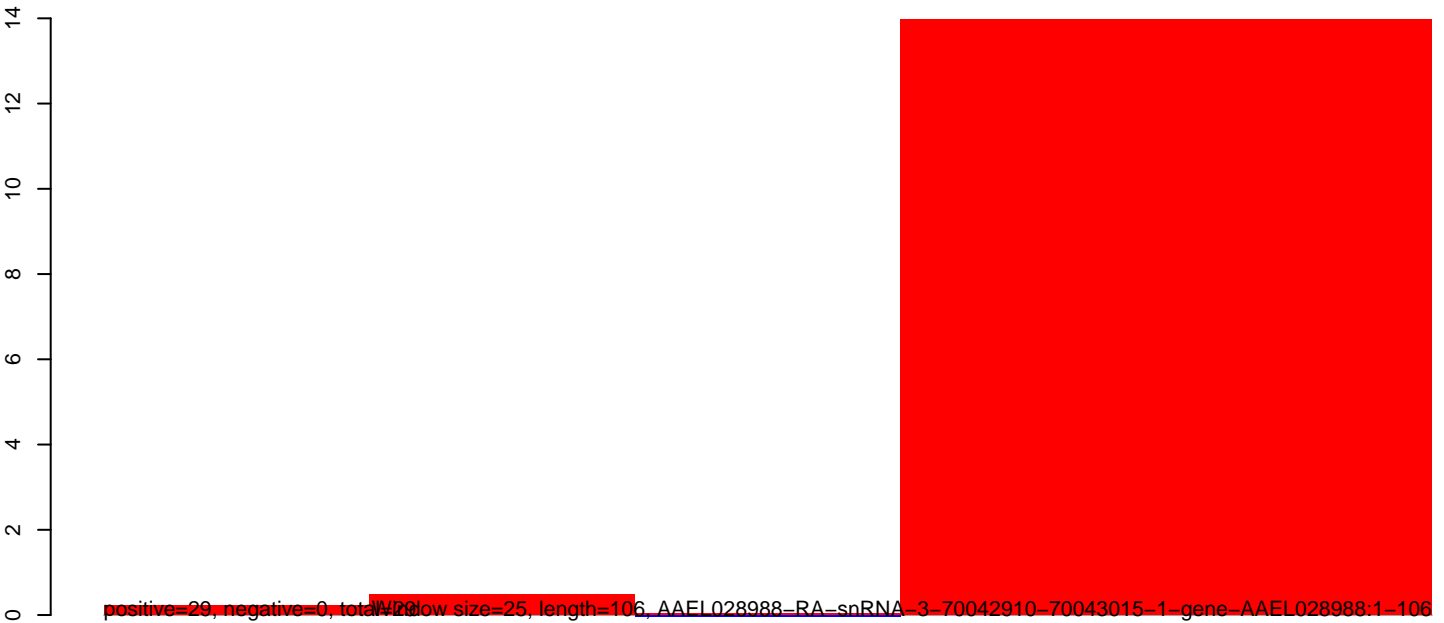
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AeAeg_Aag2.TC.Dmem.24_35.rep

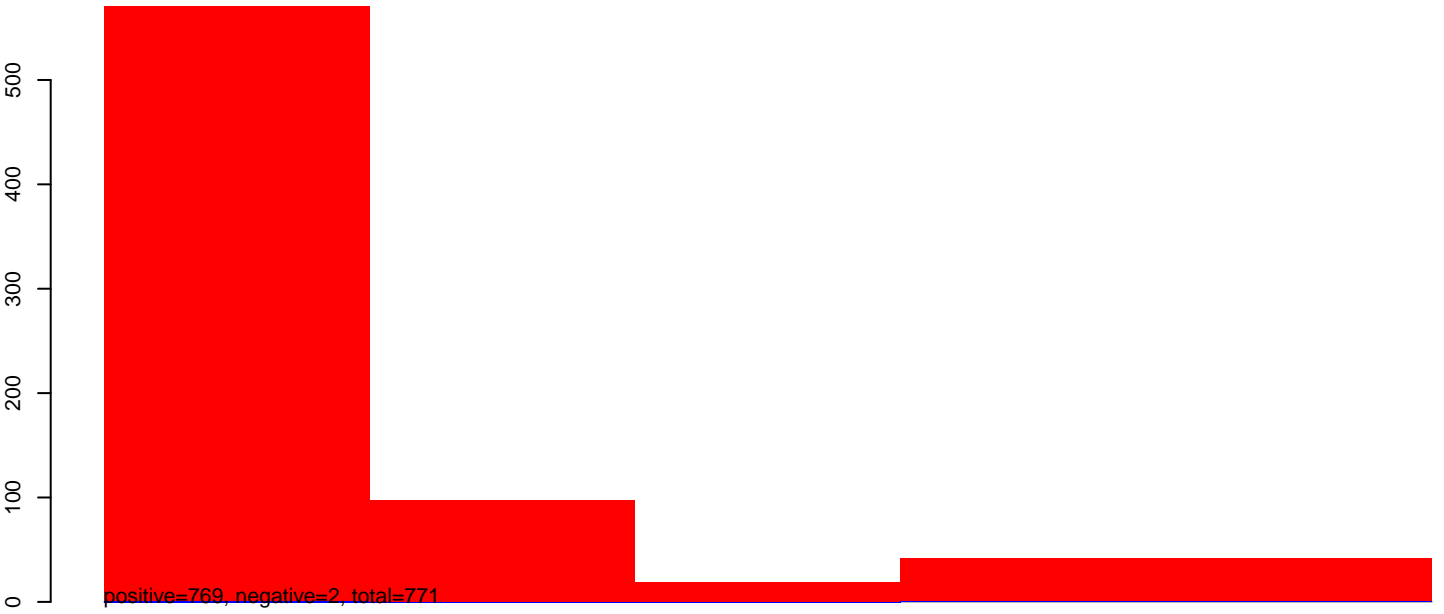


AeAeg_Aag2.TC.Dmem.rep



40 60 80 100 120 140

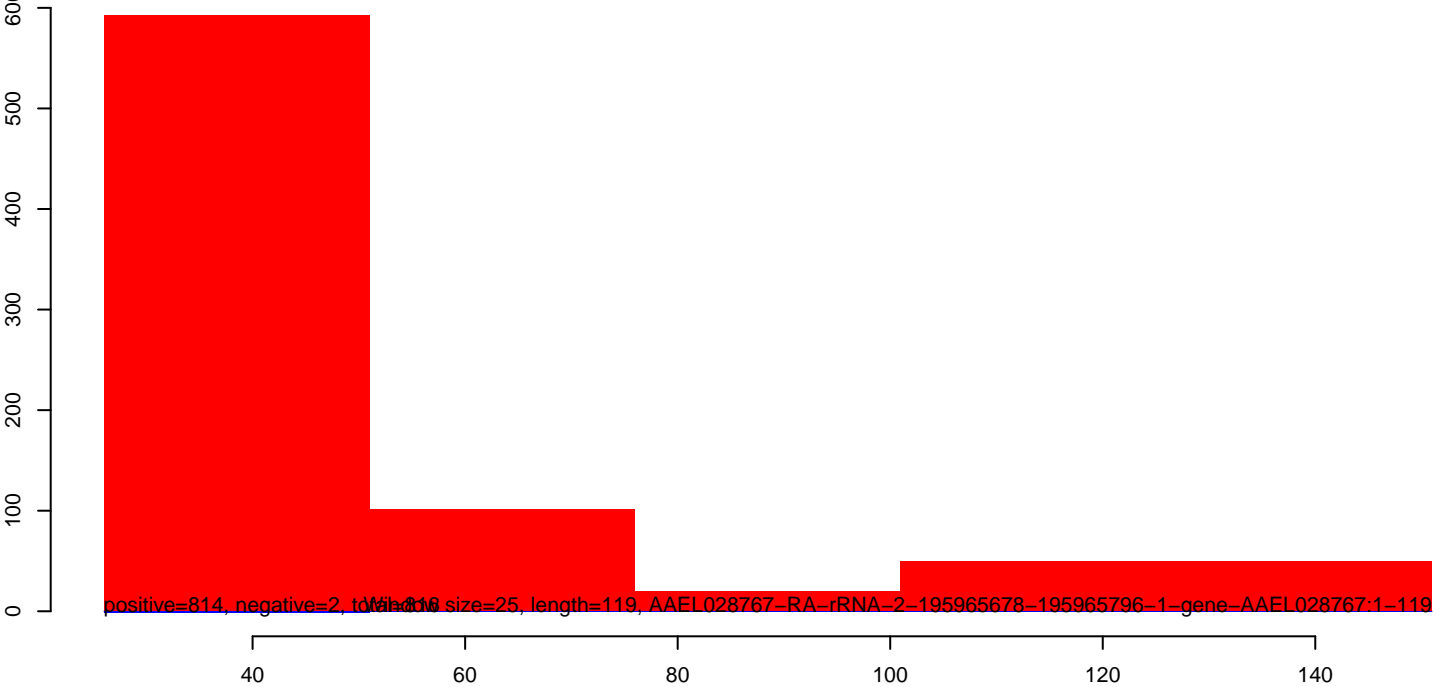
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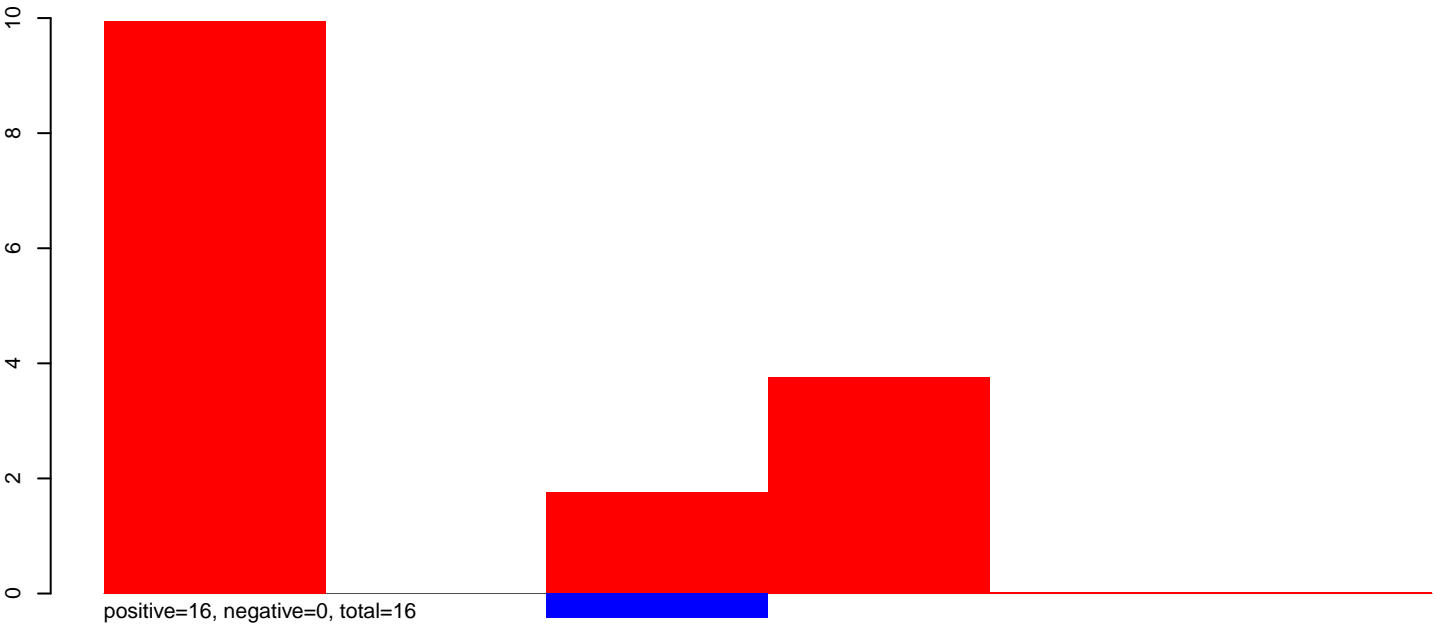
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AeAeg_Aag2.TC.Dmem.rep



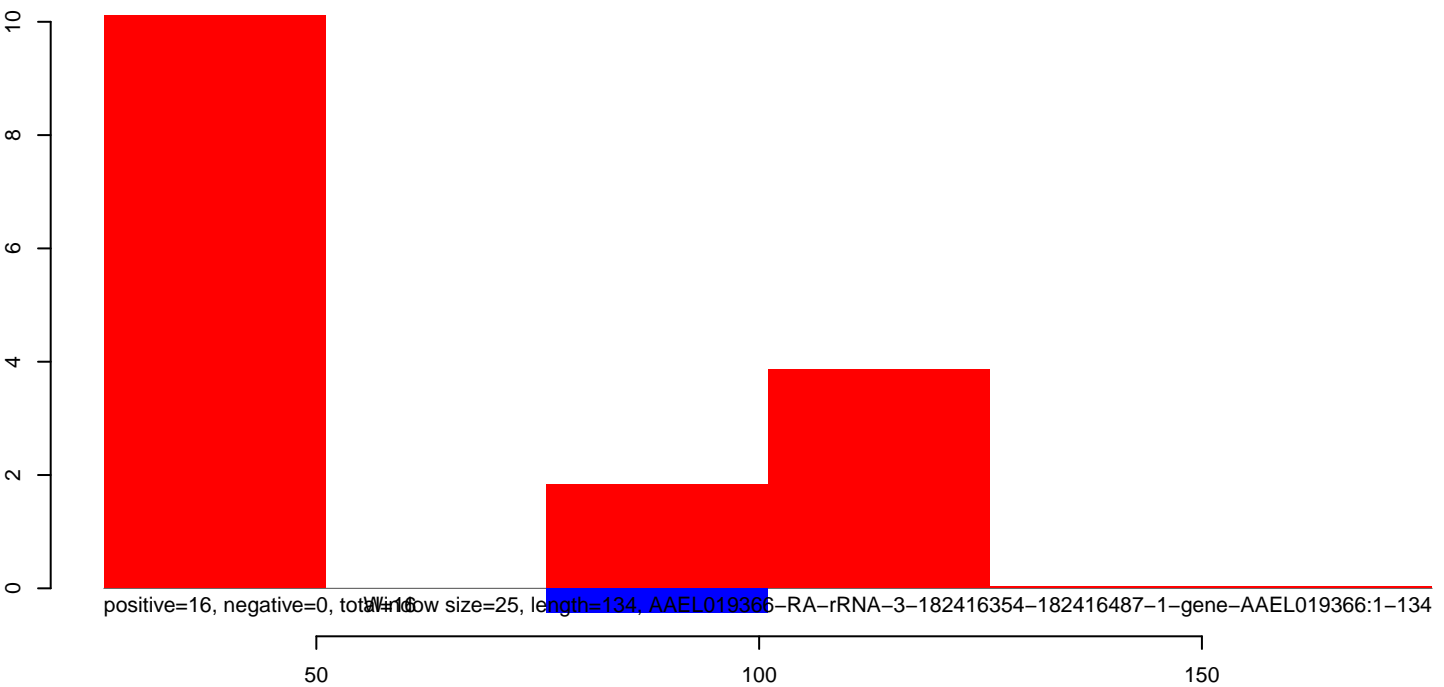
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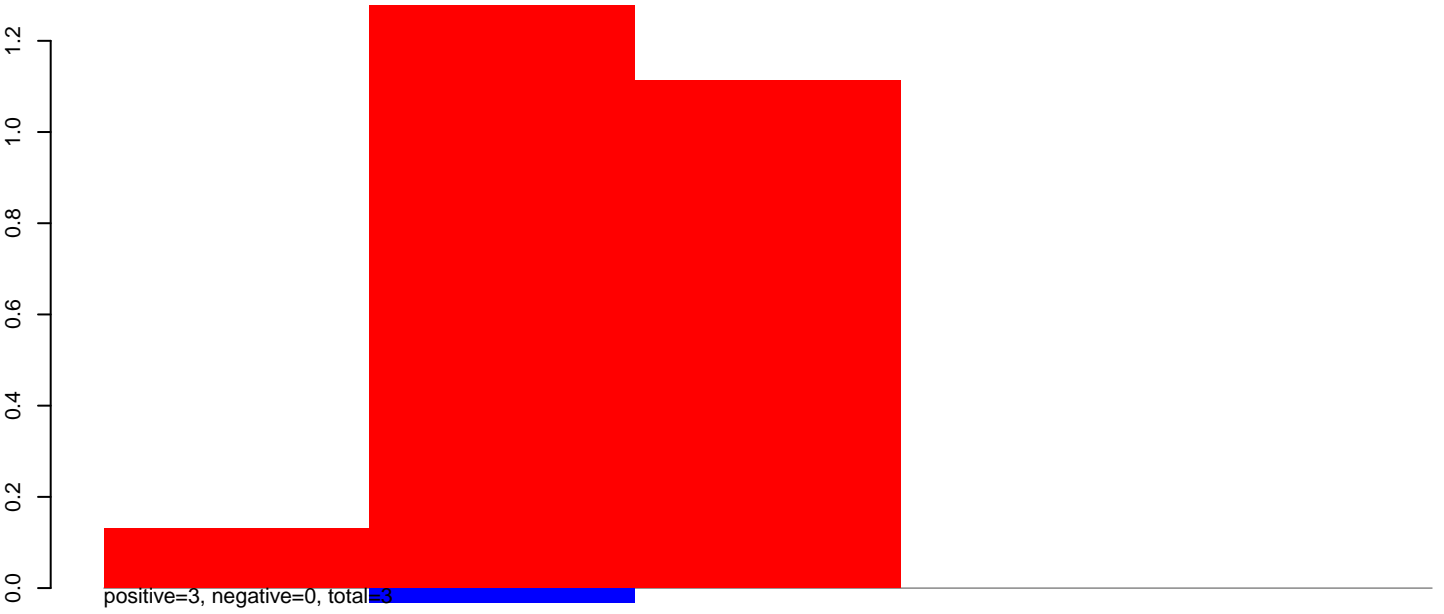
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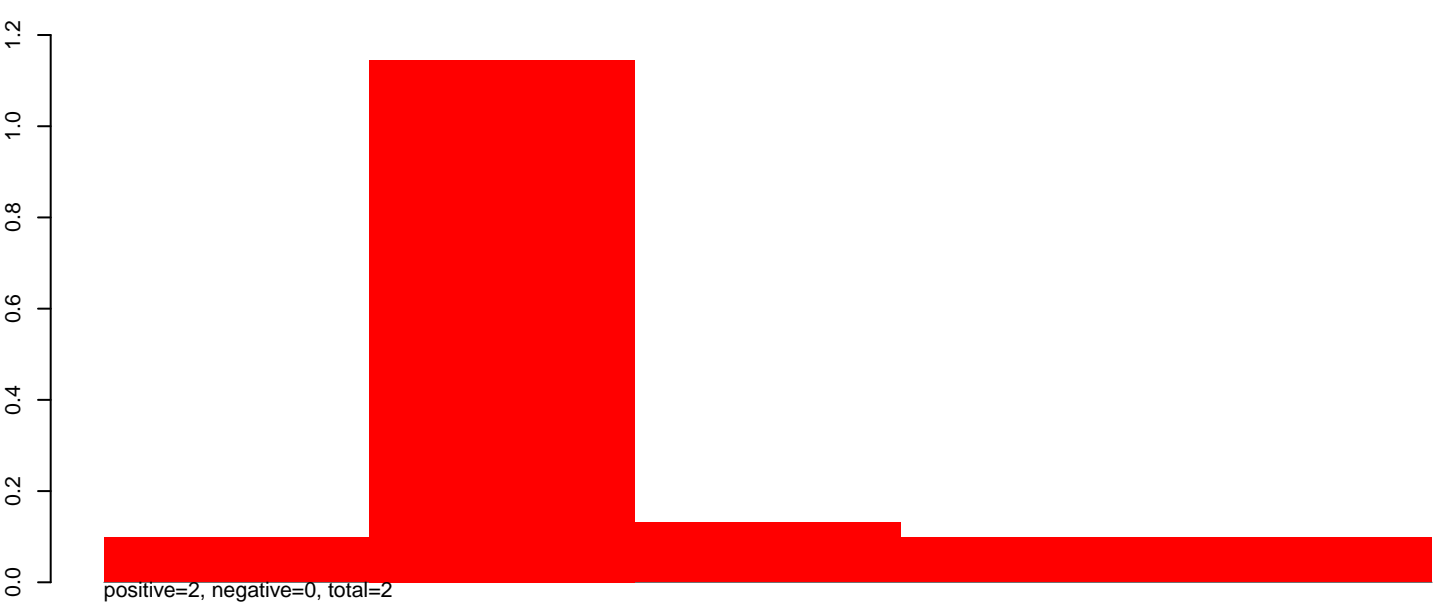
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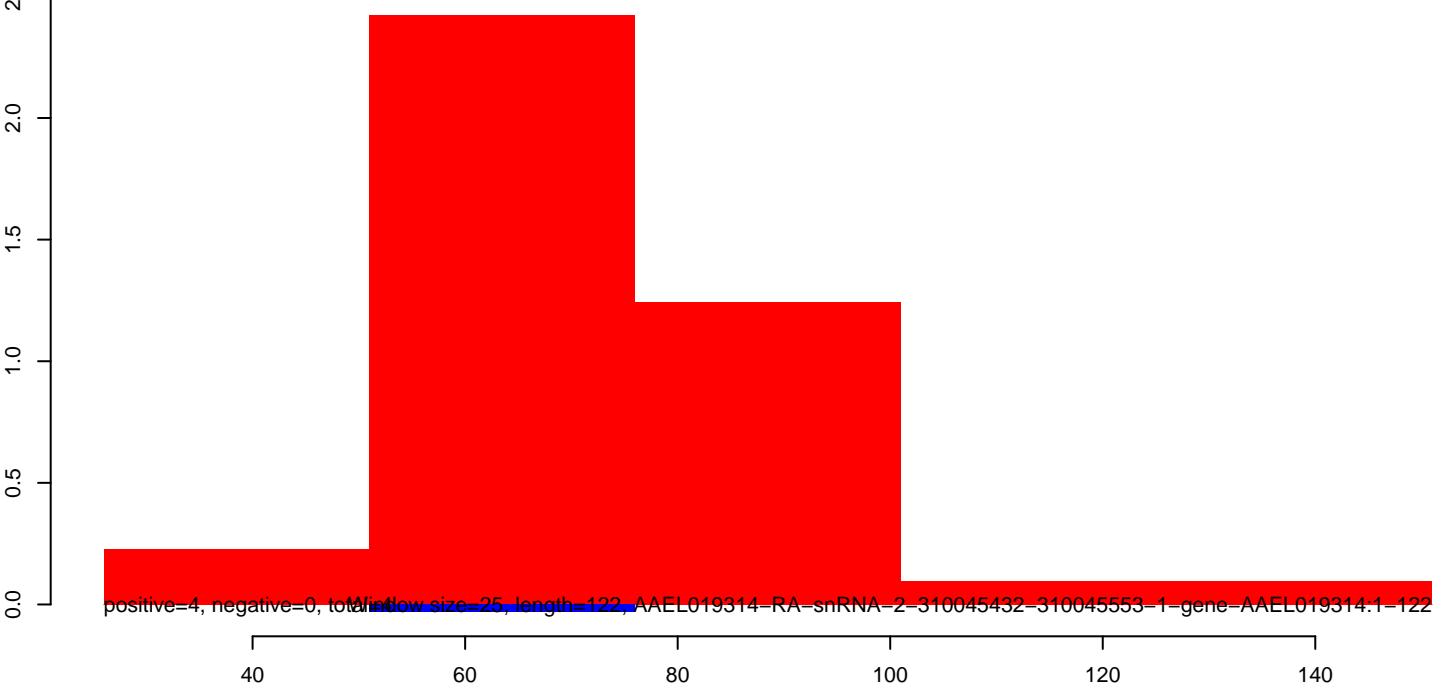
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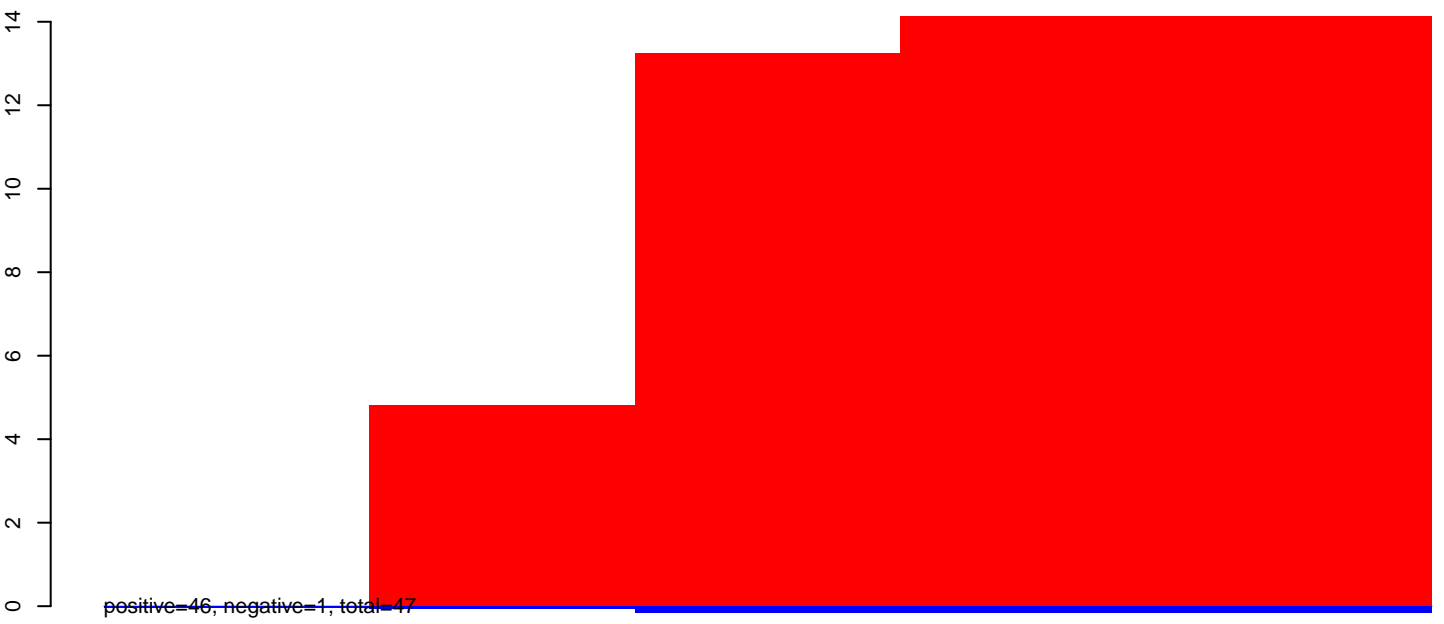
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AeAeg_Aag2.TC.Dmem.rep



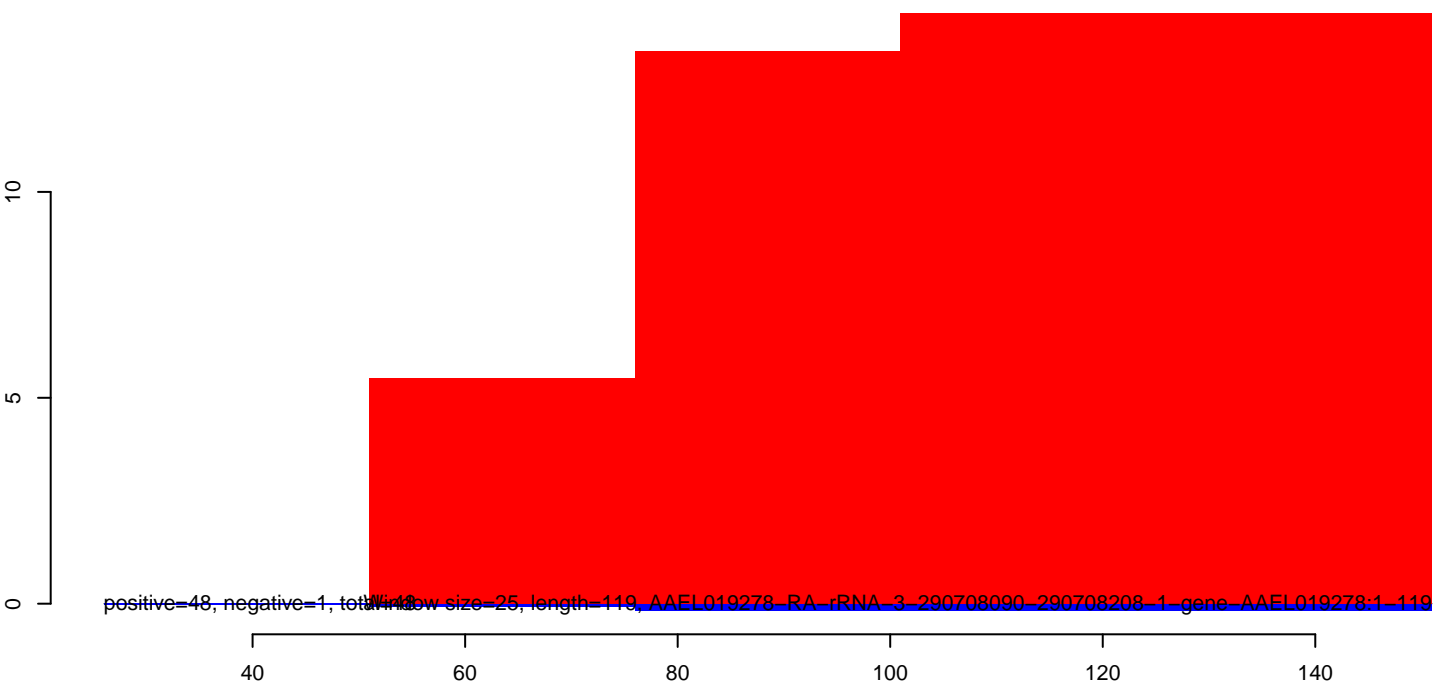
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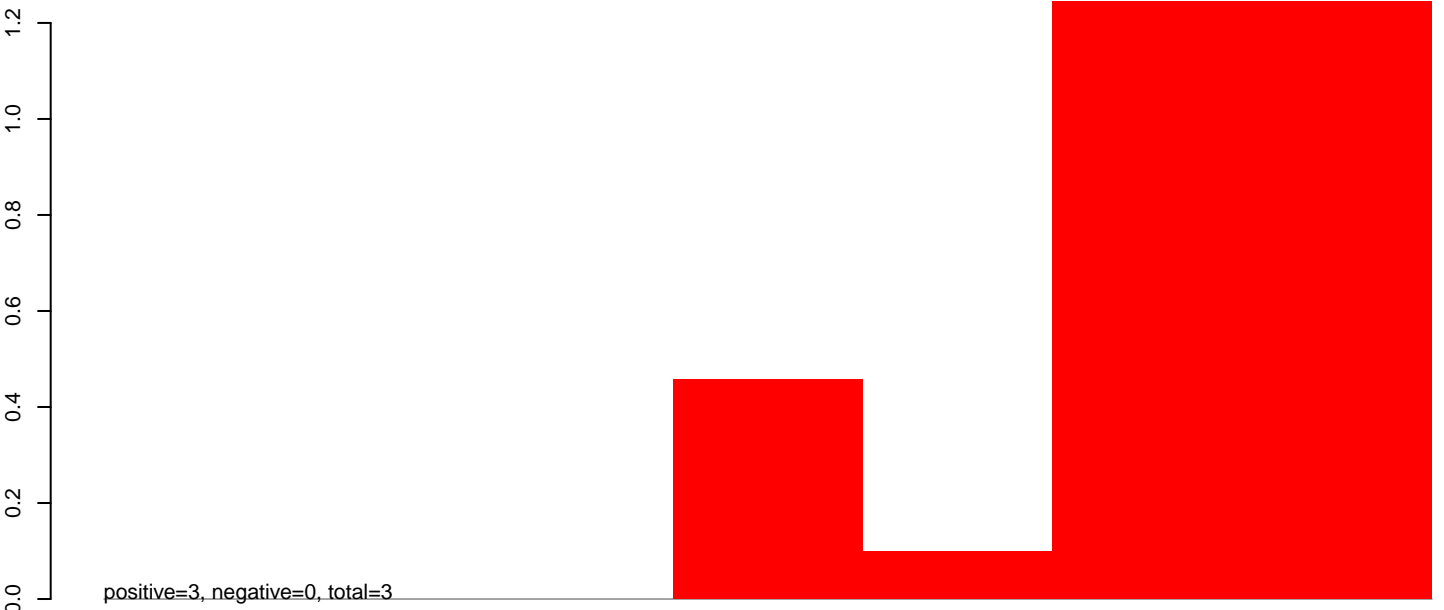
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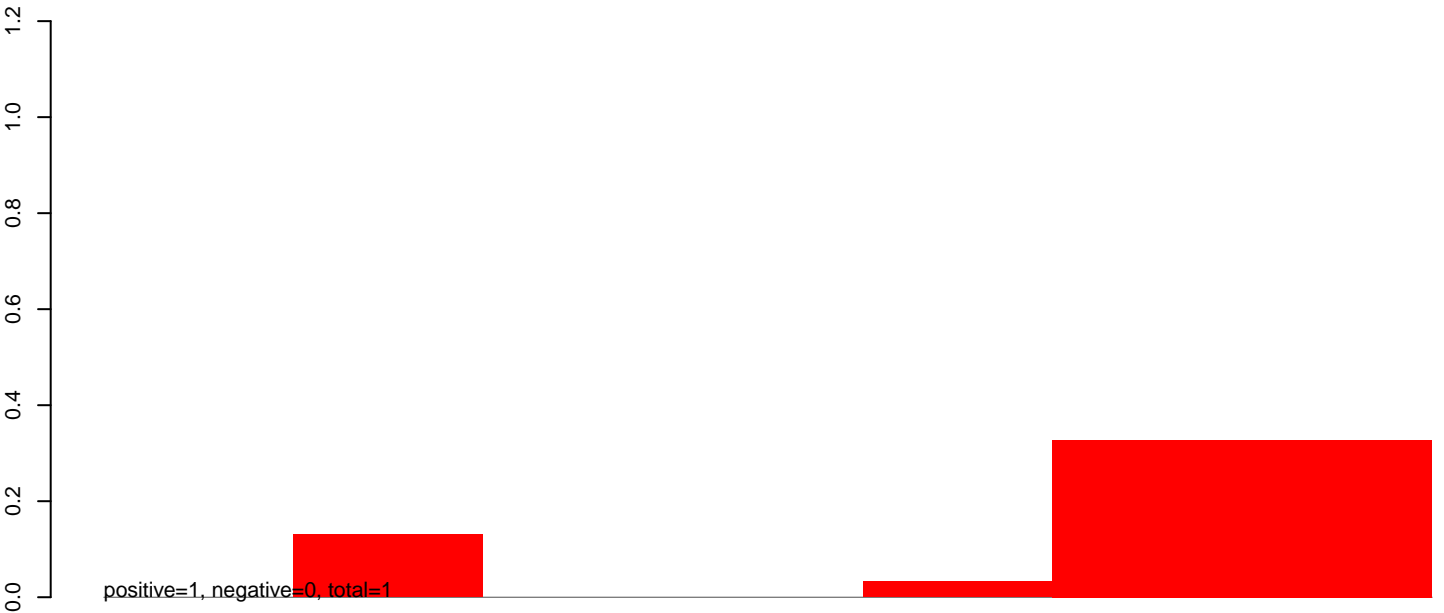
AeAeg_Aag2.TC.Dmem.rep



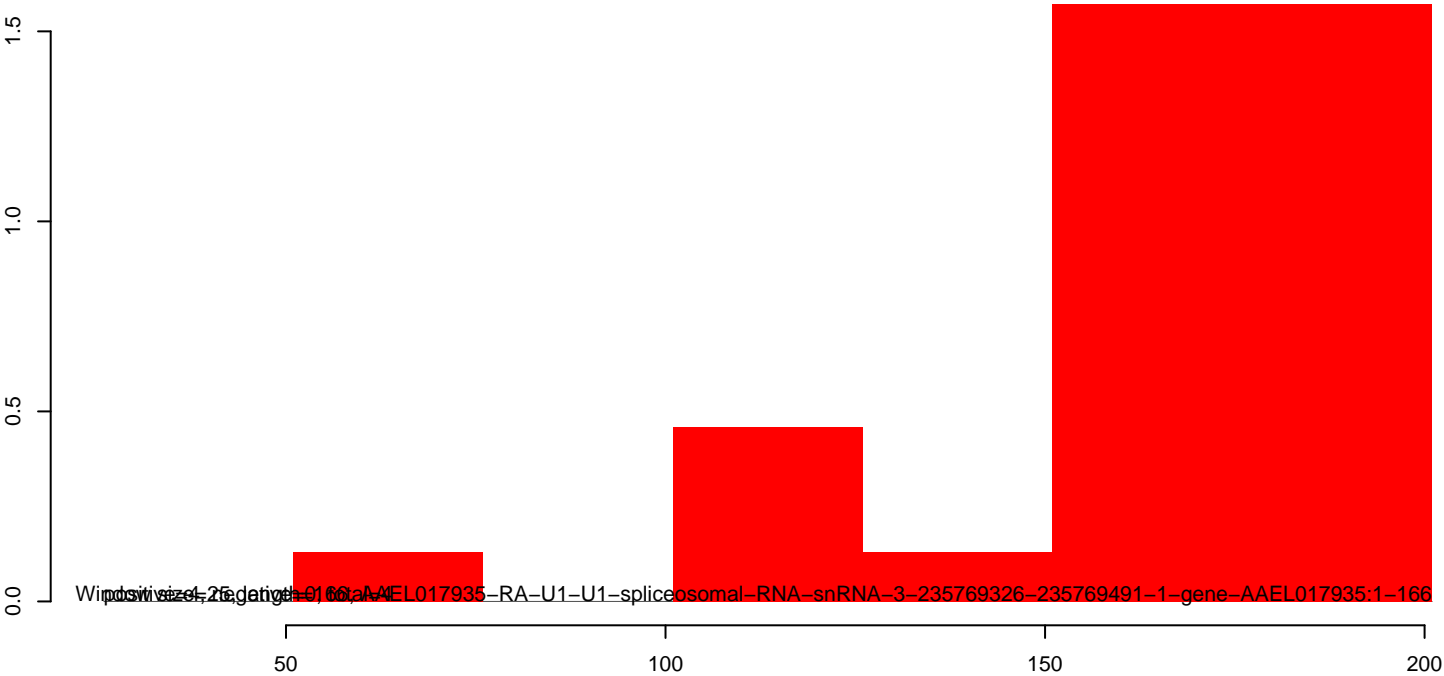
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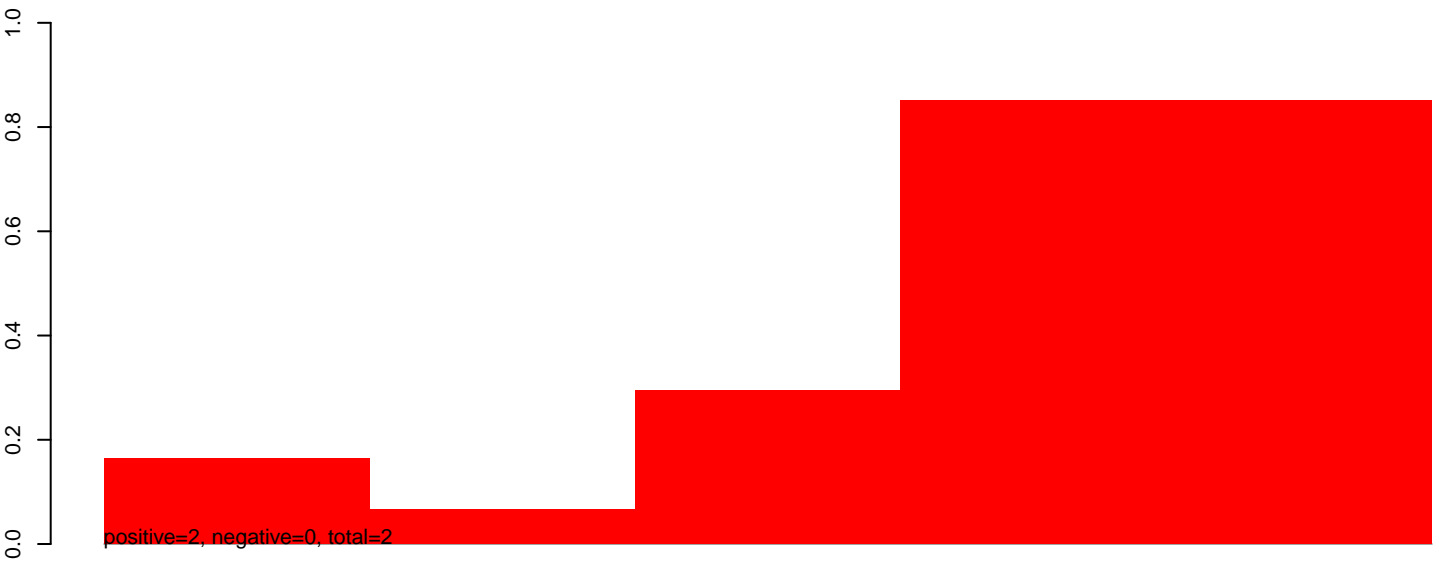
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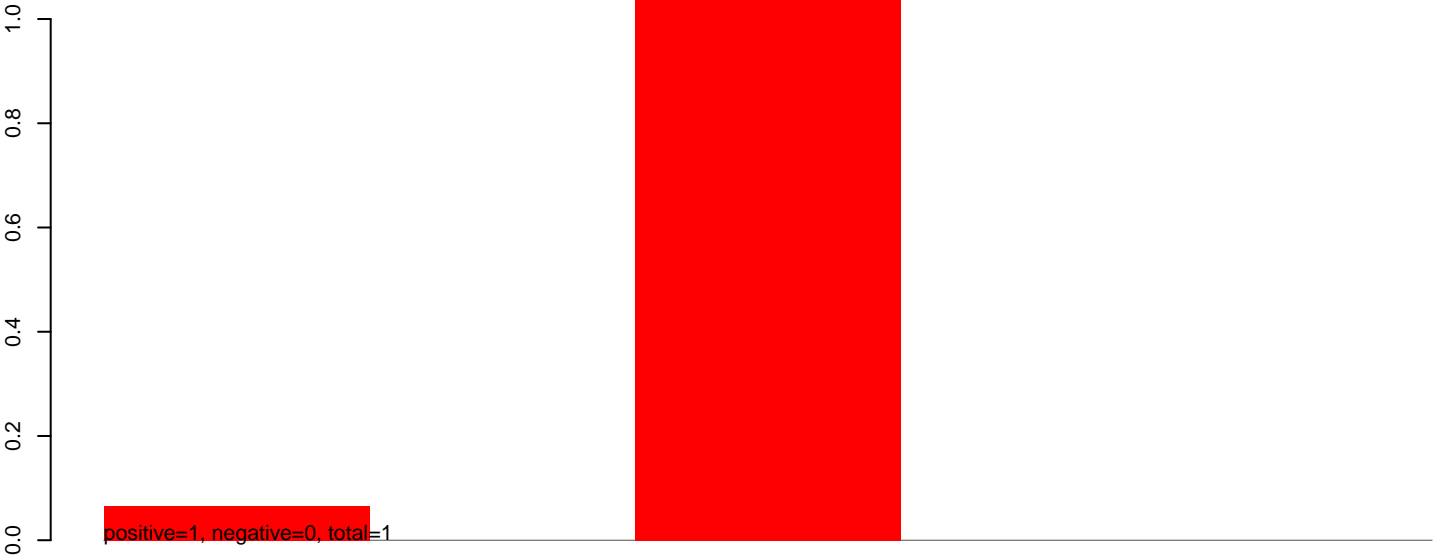
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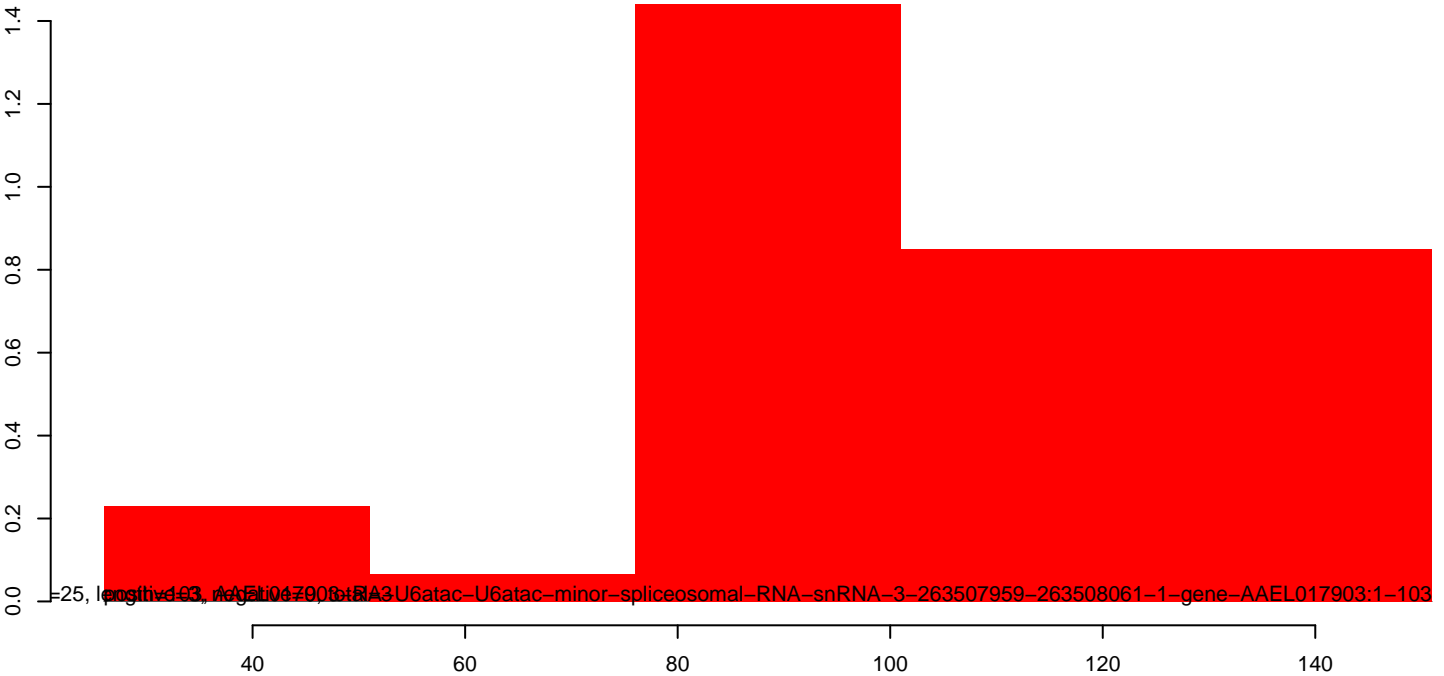
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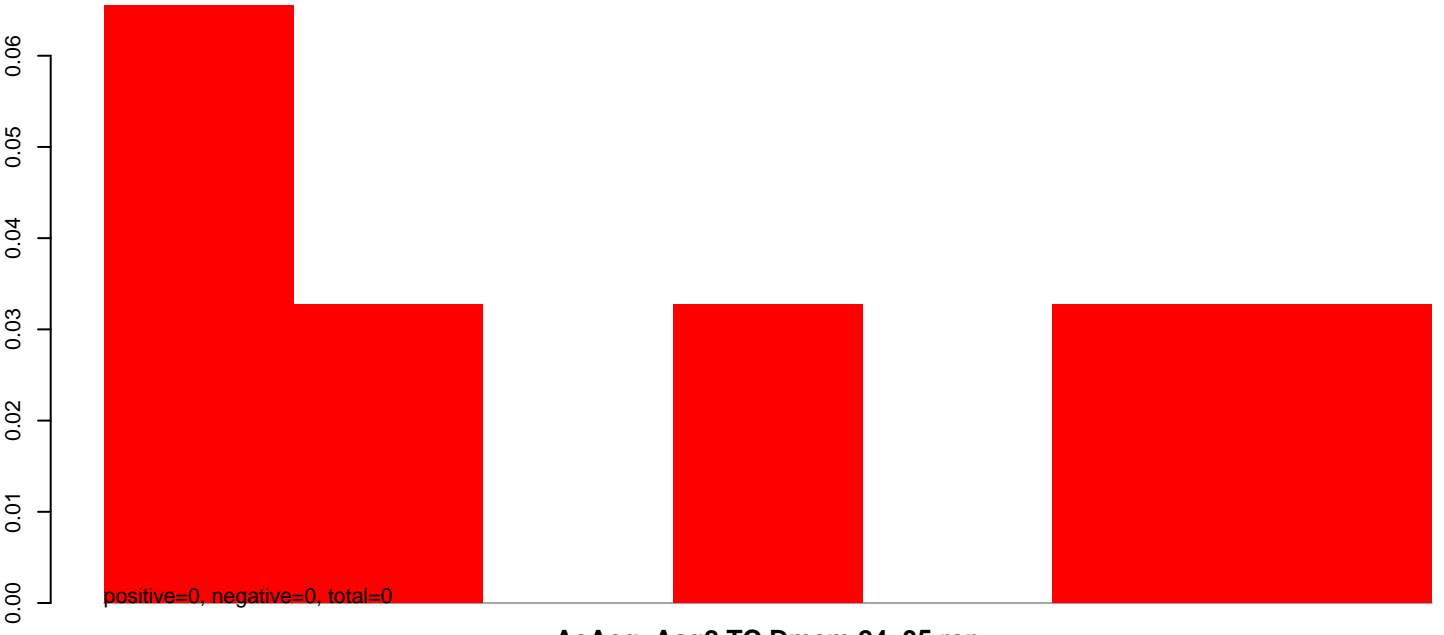
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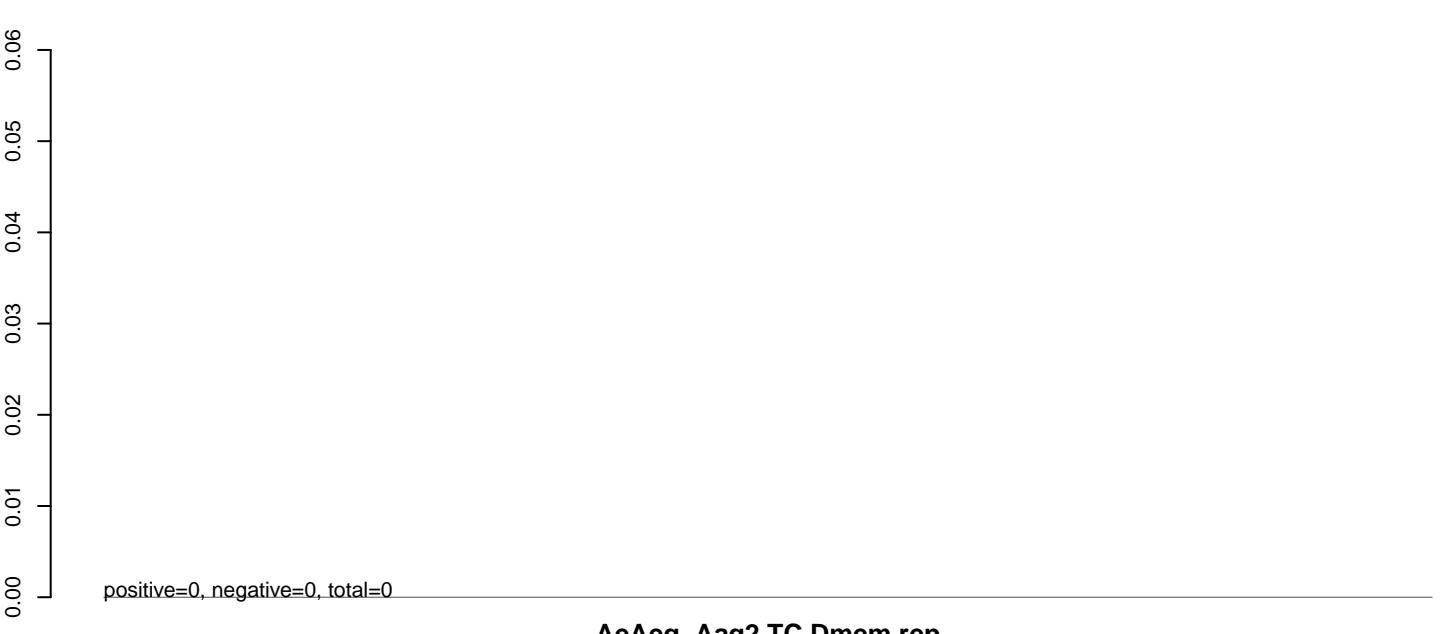
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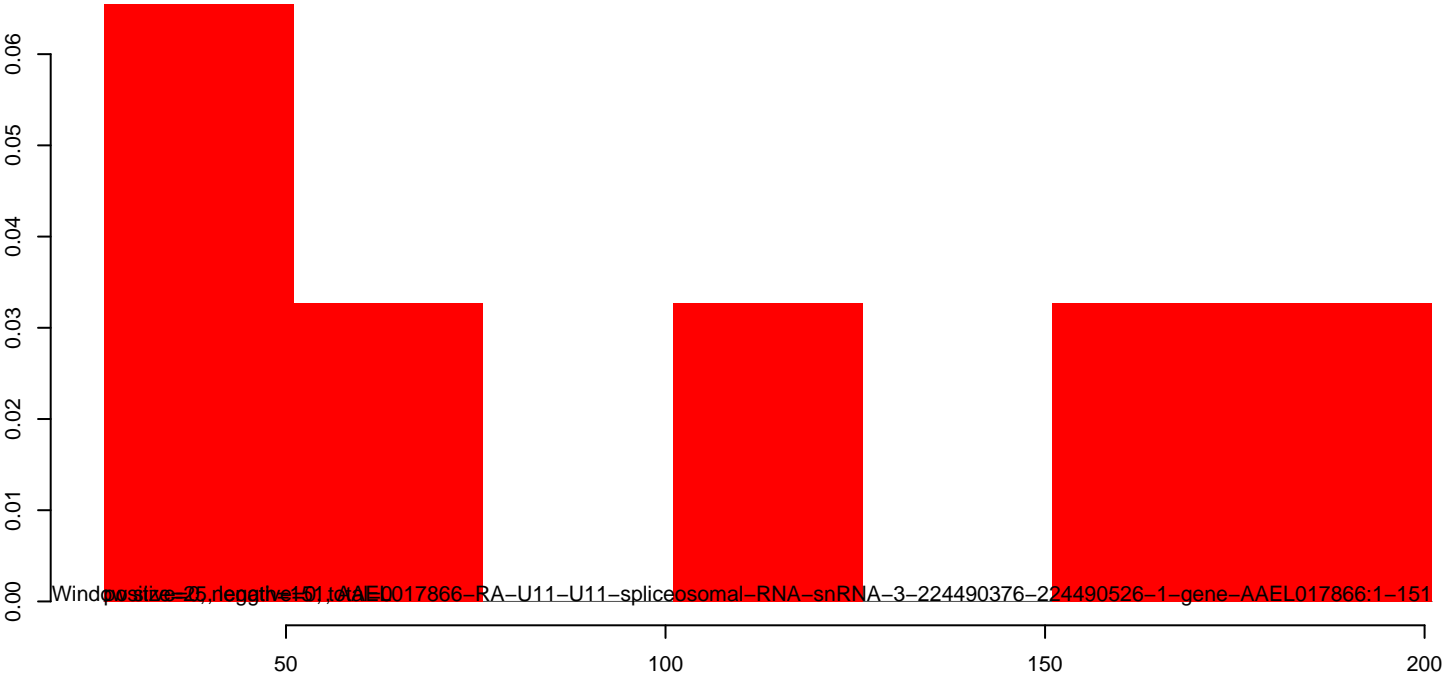
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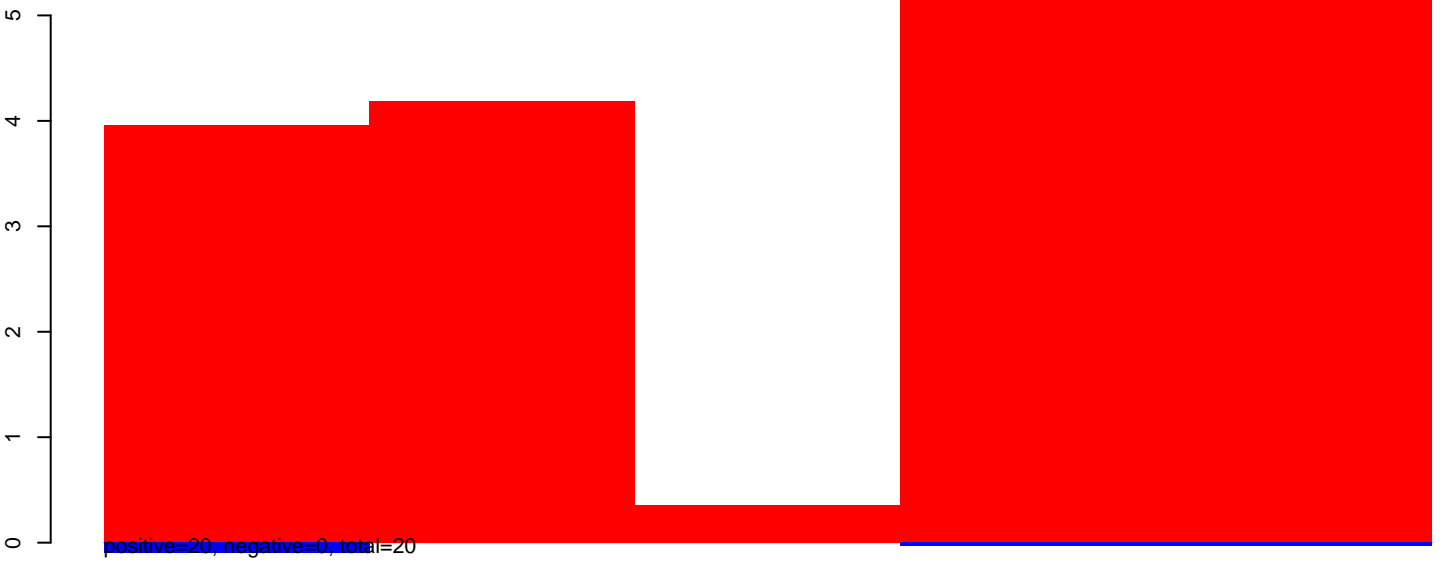
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AeAeg_Aag2.TC.Dmem.rep



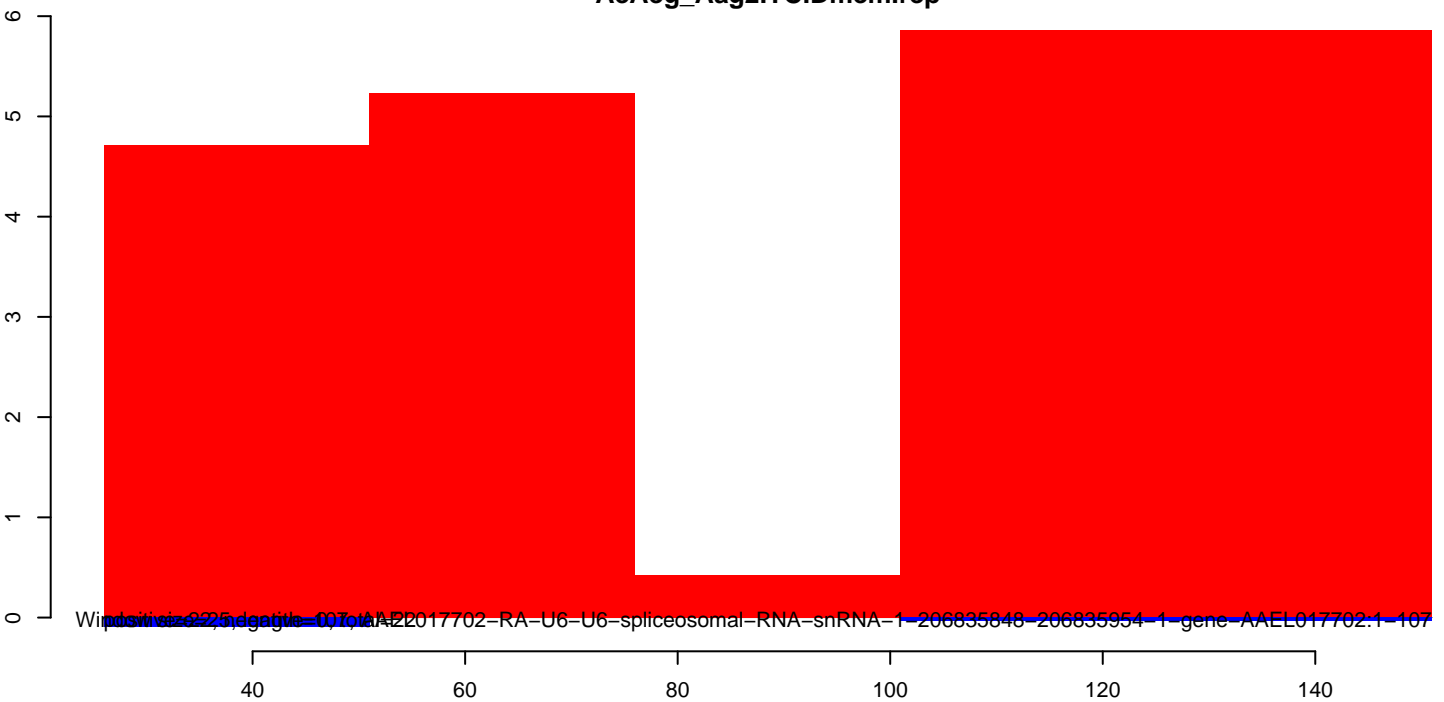
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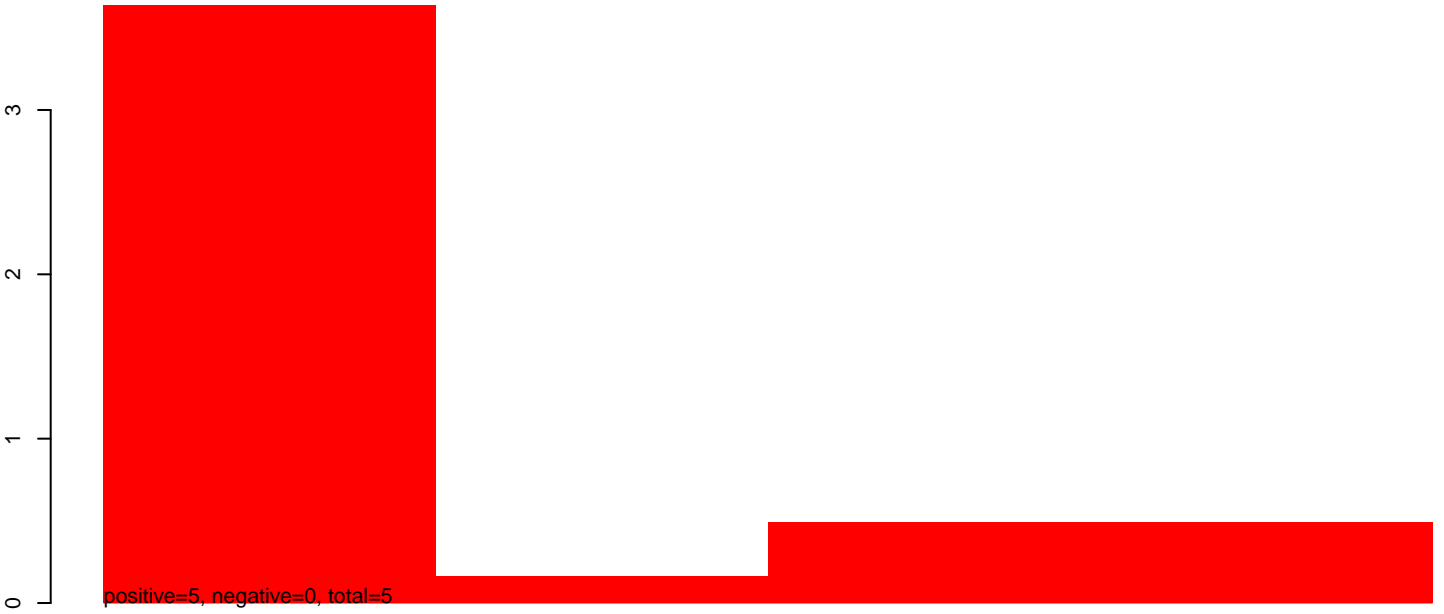
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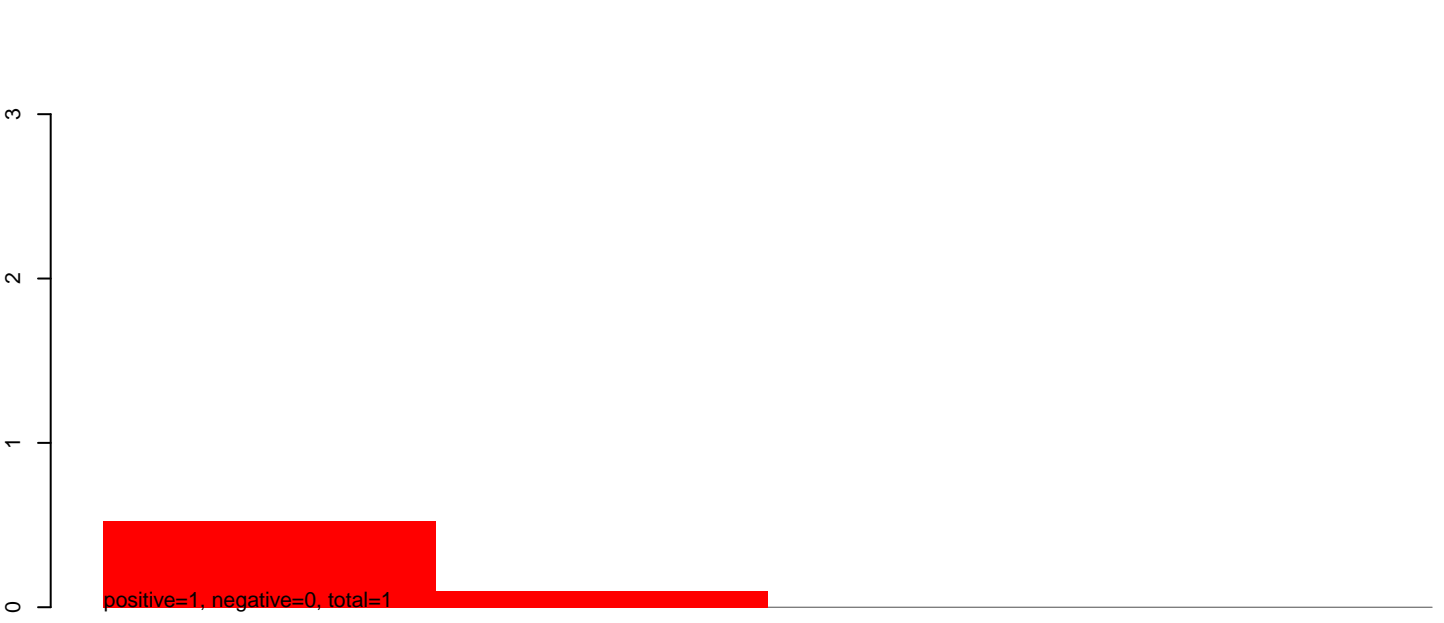
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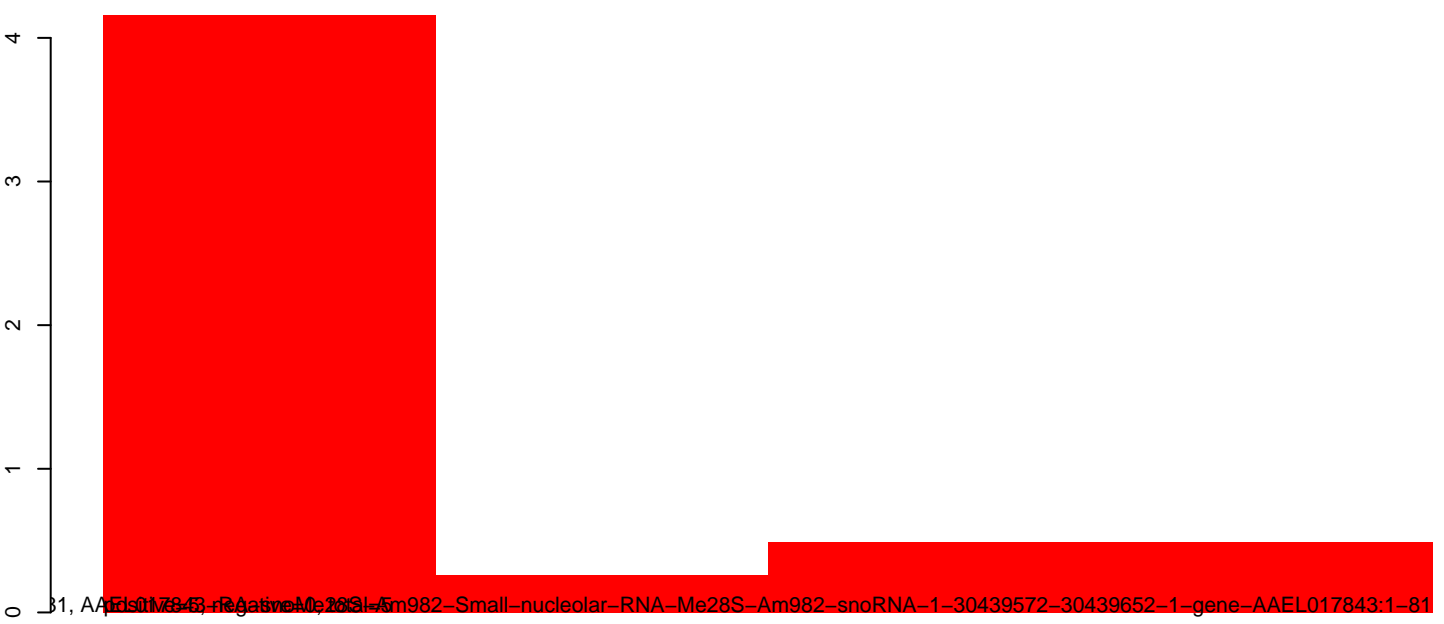
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AeAeg_Aag2.TC.Dmem.24_35.rep

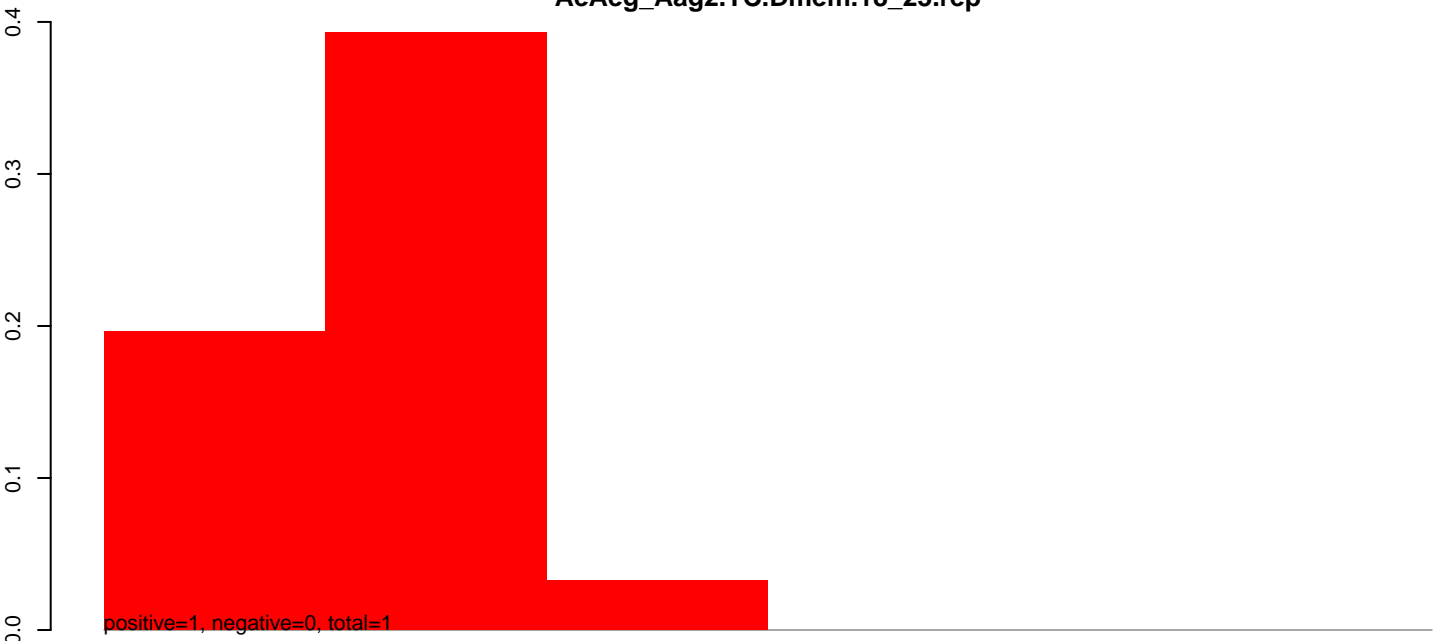


AeAeg_Aag2.TC.Dmem.rep



31, AAF017843, rRNA, 28S, 1-5, 30439572-30439652-1-gene-AAEL017843:1-81

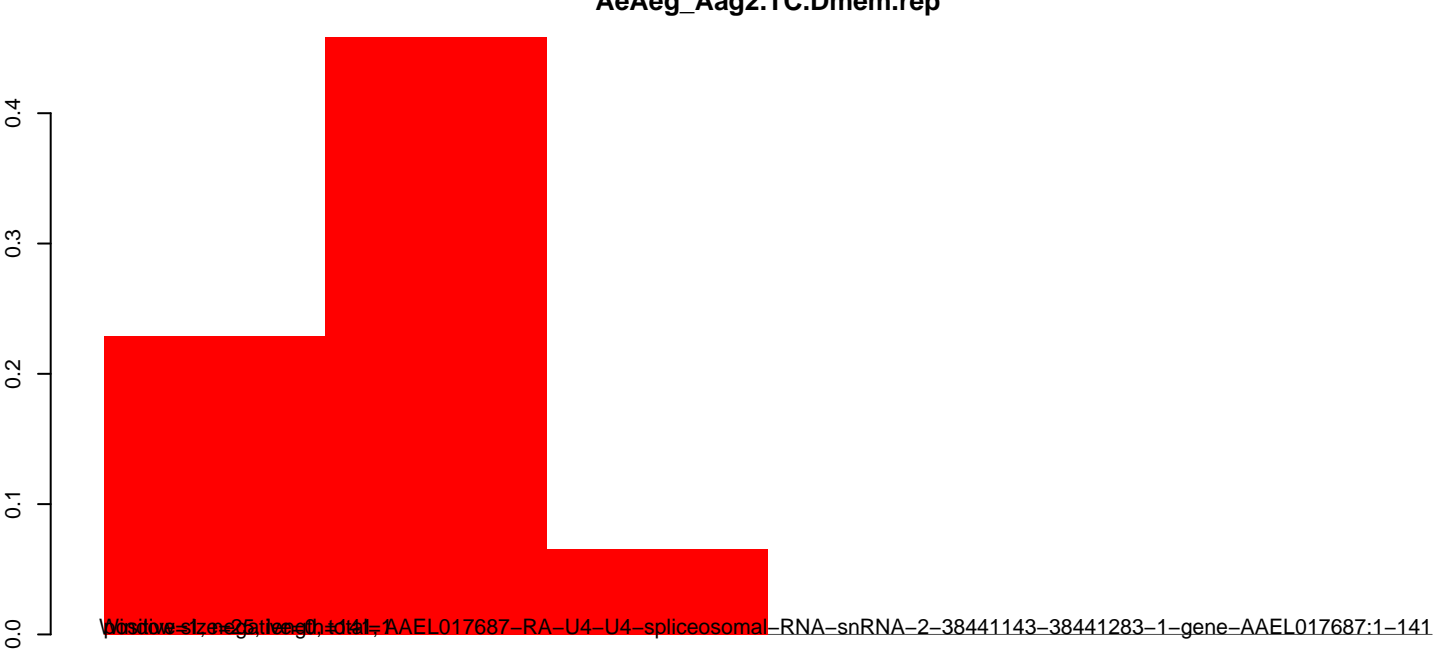
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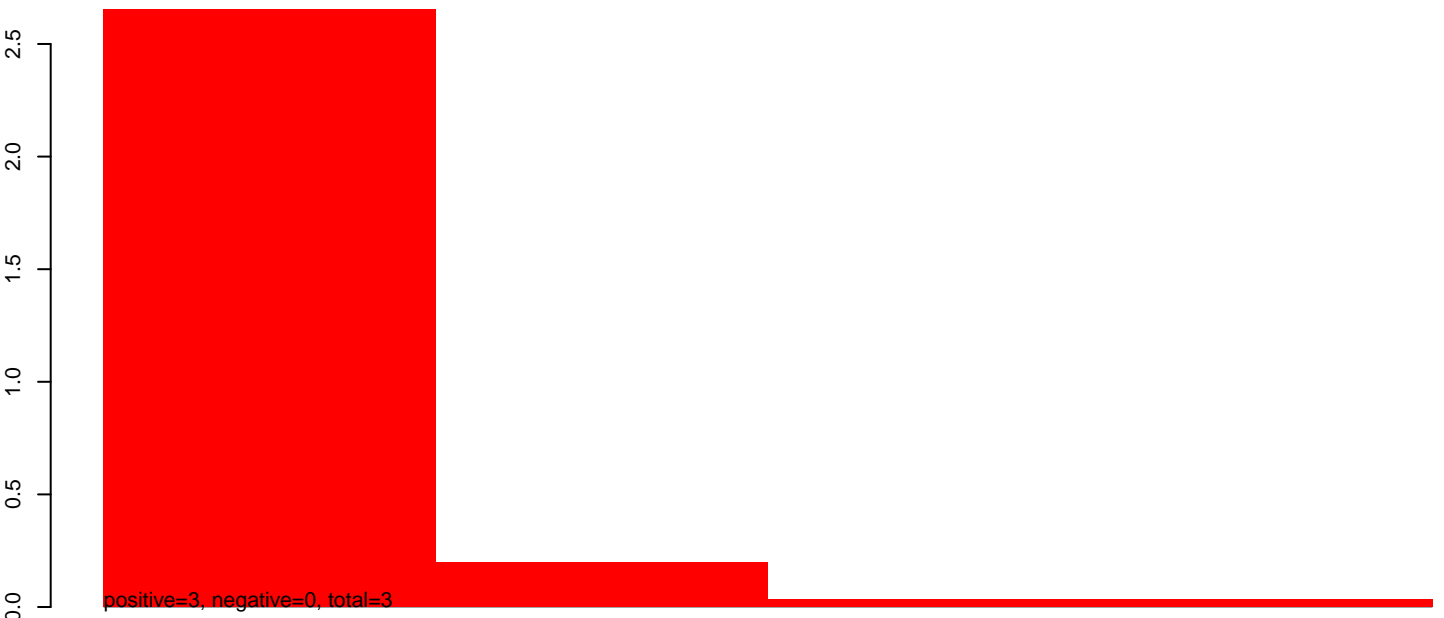
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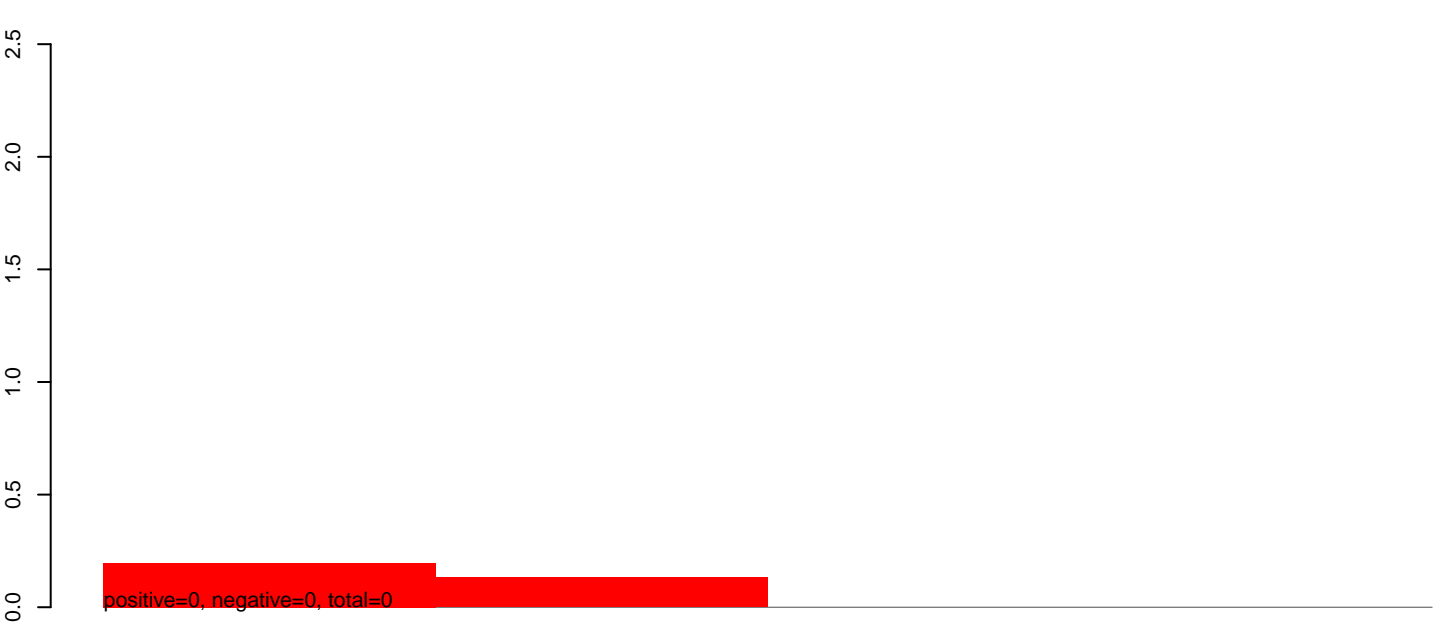
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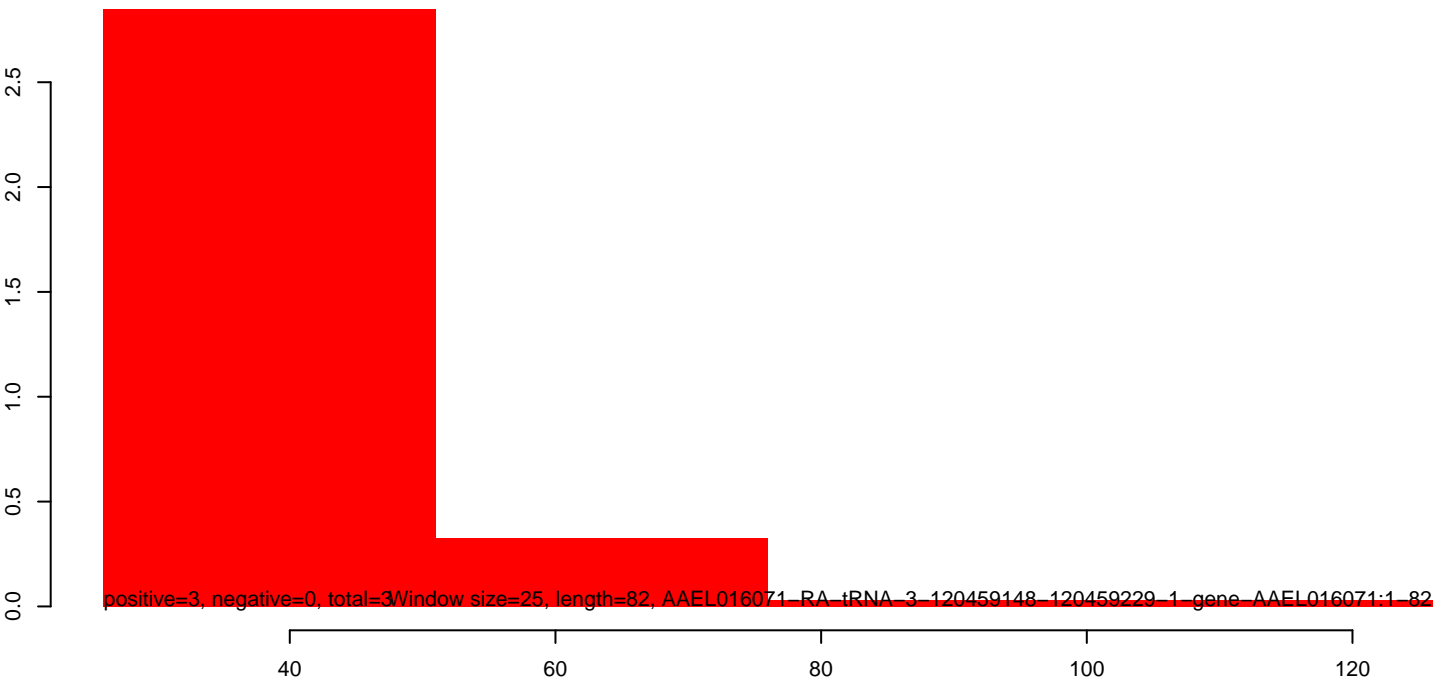
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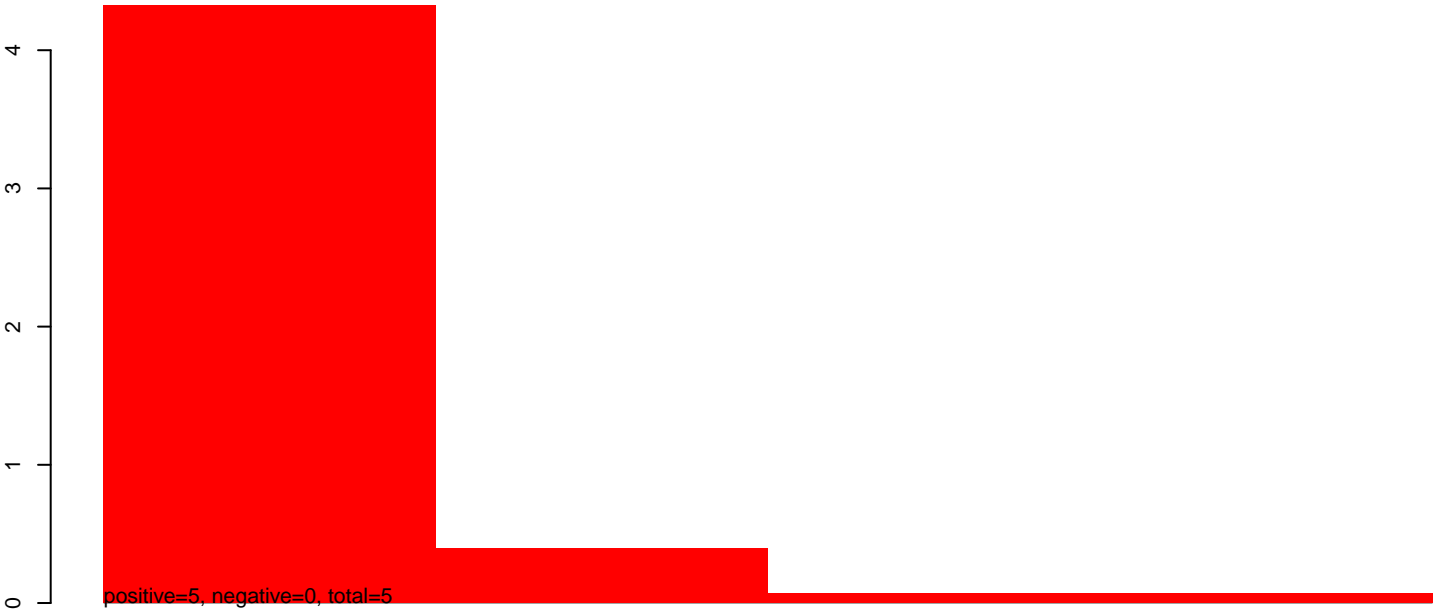
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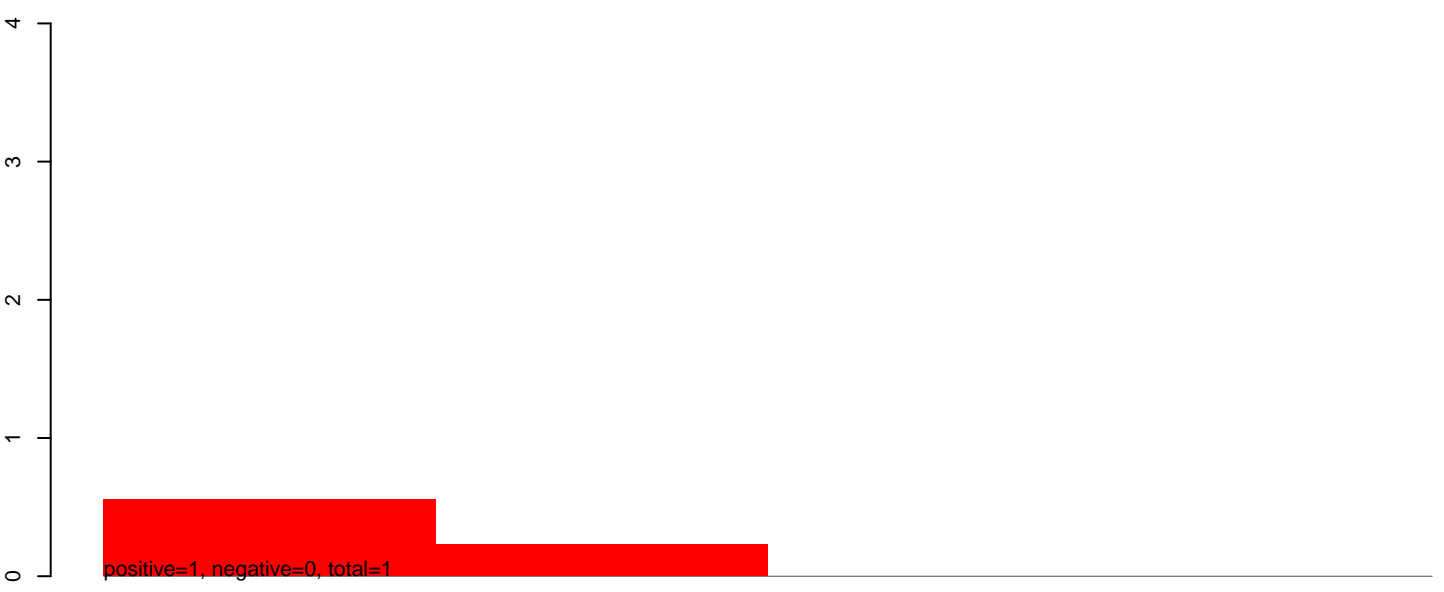
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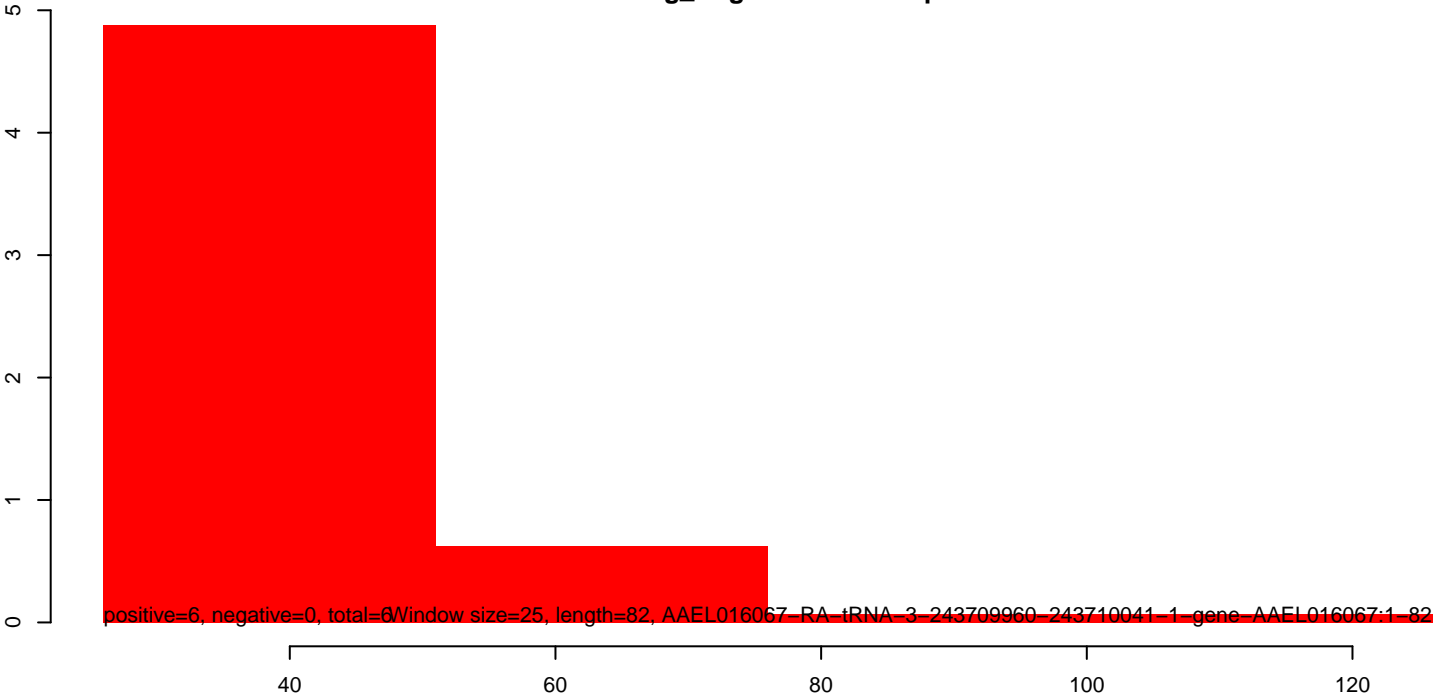
AeAeg_Aag2.TC.Dmem.18_23.rep



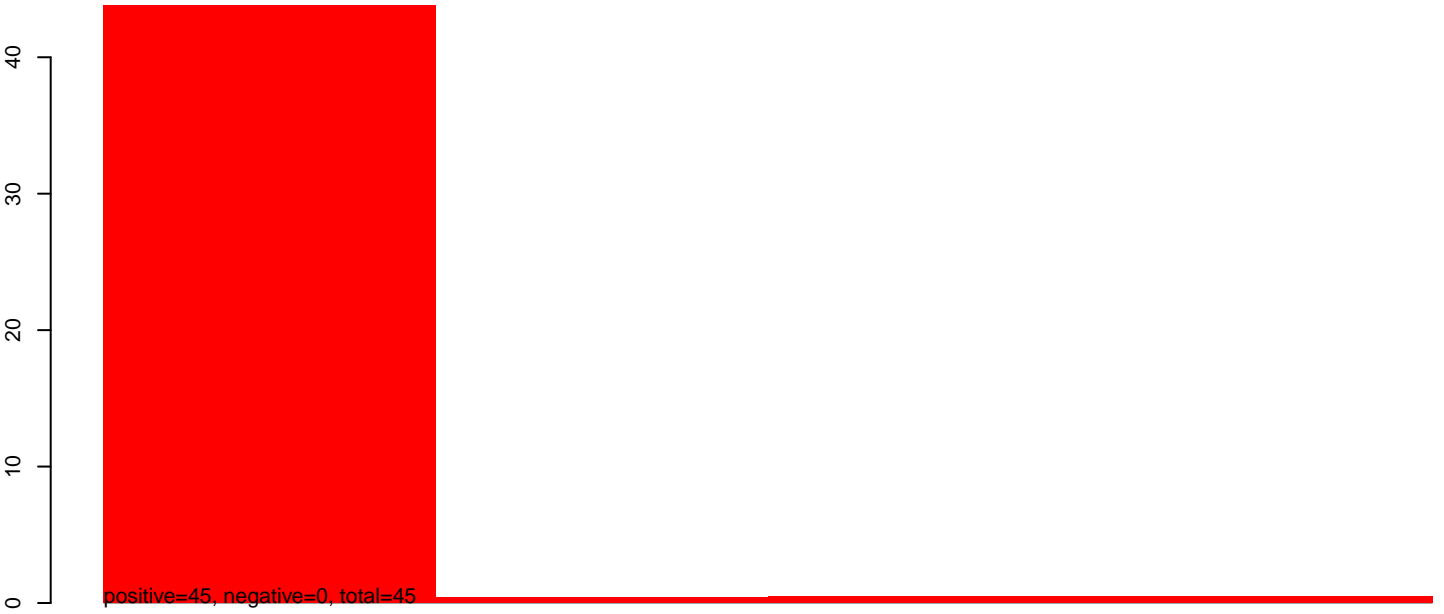
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AeAeg_Aag2.TC.Dmem.rep



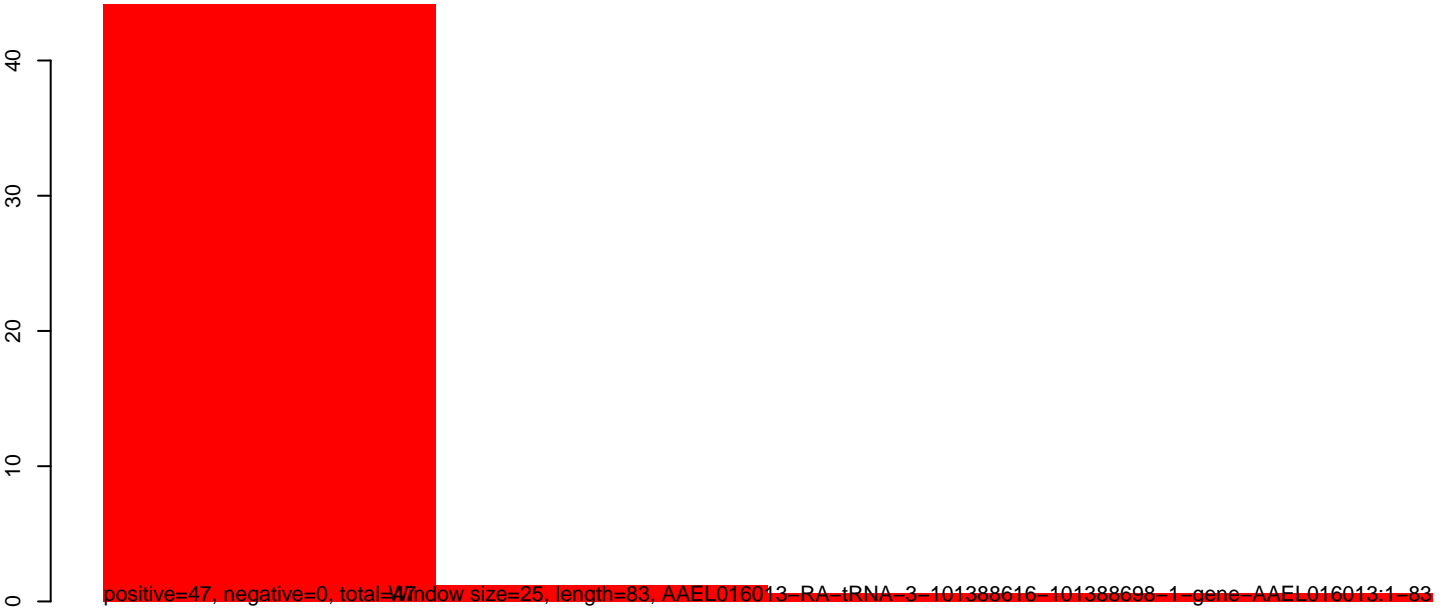
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AeAeg_Aag2.TC.Dmem.24_35.rep

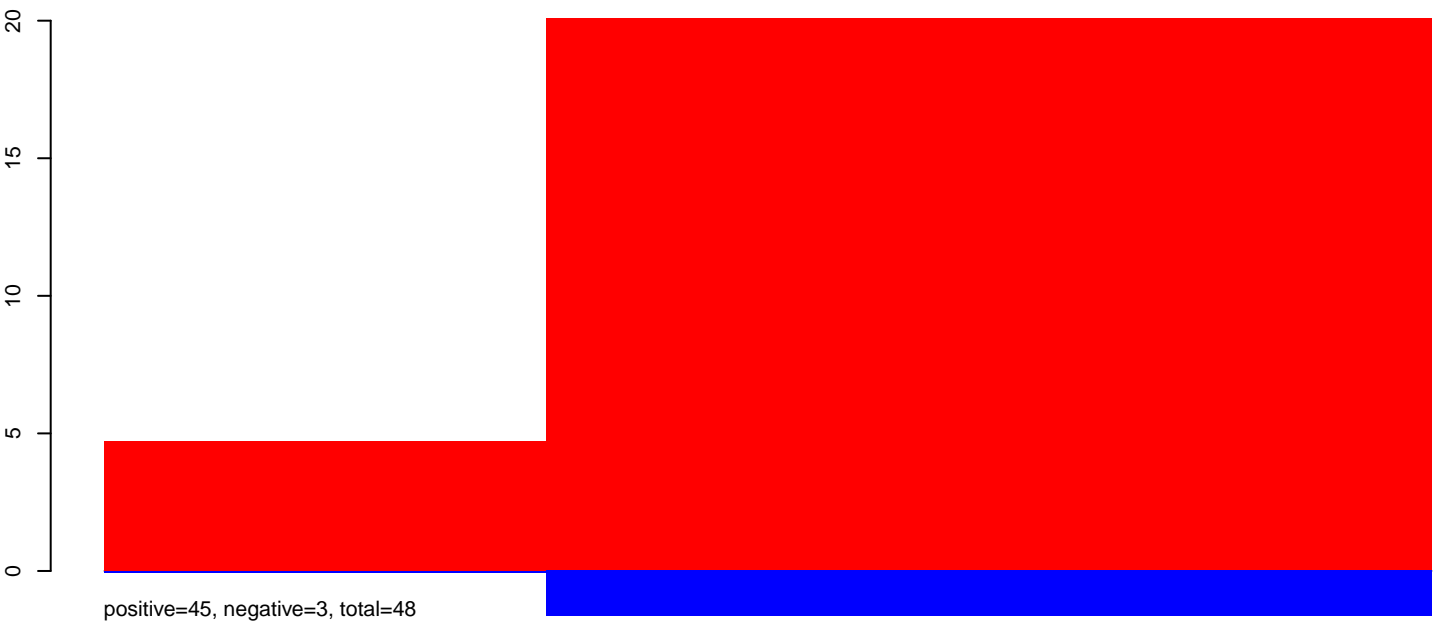


AeAeg_Aag2.TC.Dmem.rep

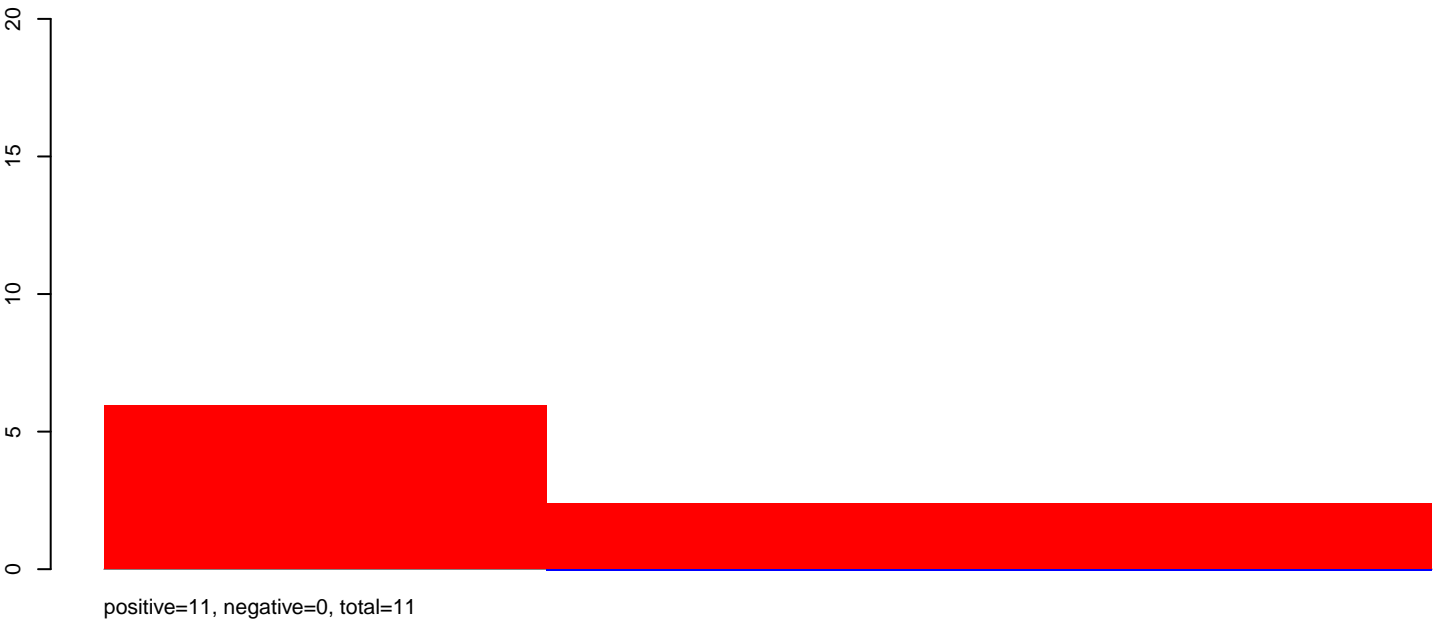


Window size=25, length=83, AAEL016013-RA-101388616-101388698-1-gene-AAEL016013:1-83

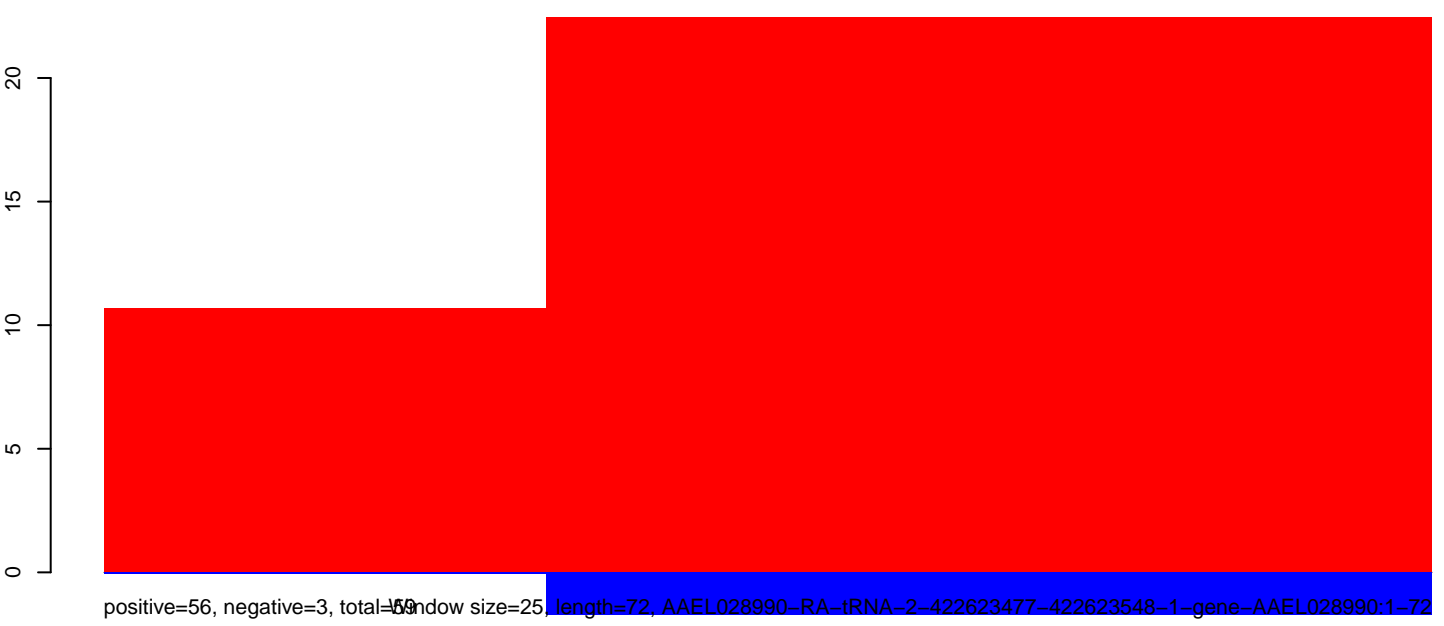
AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep

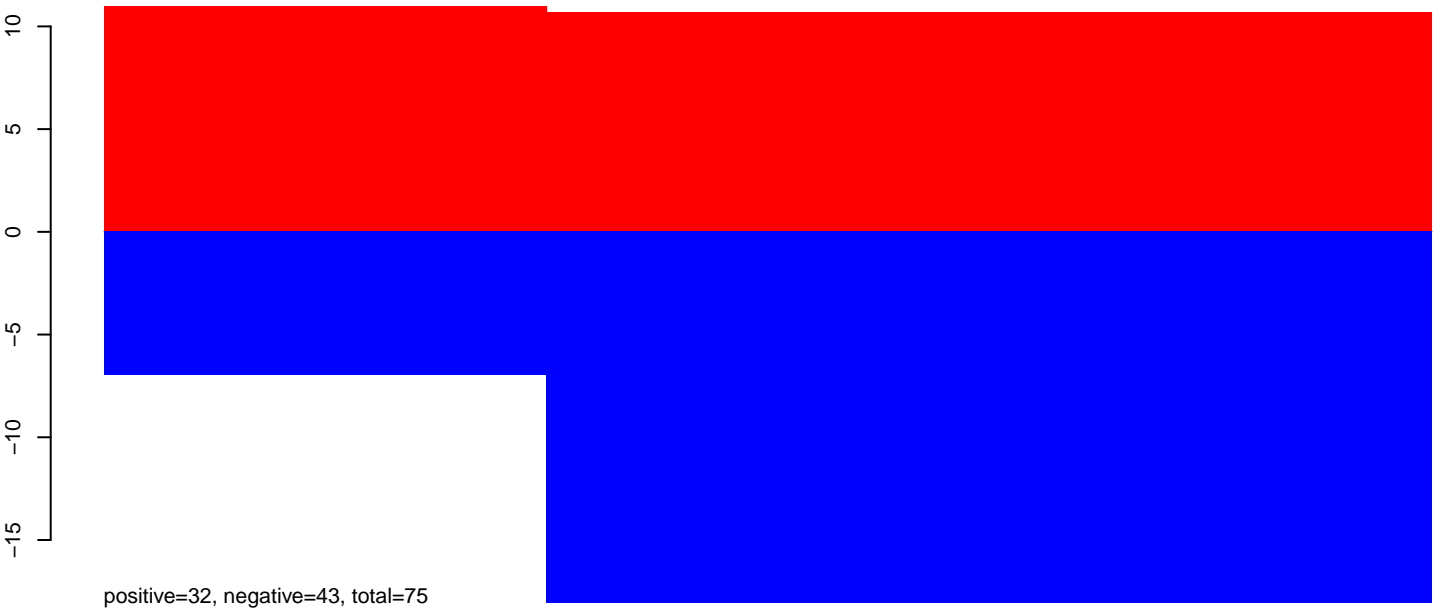


AeAeg_Aag2.TC.Dmem.rep

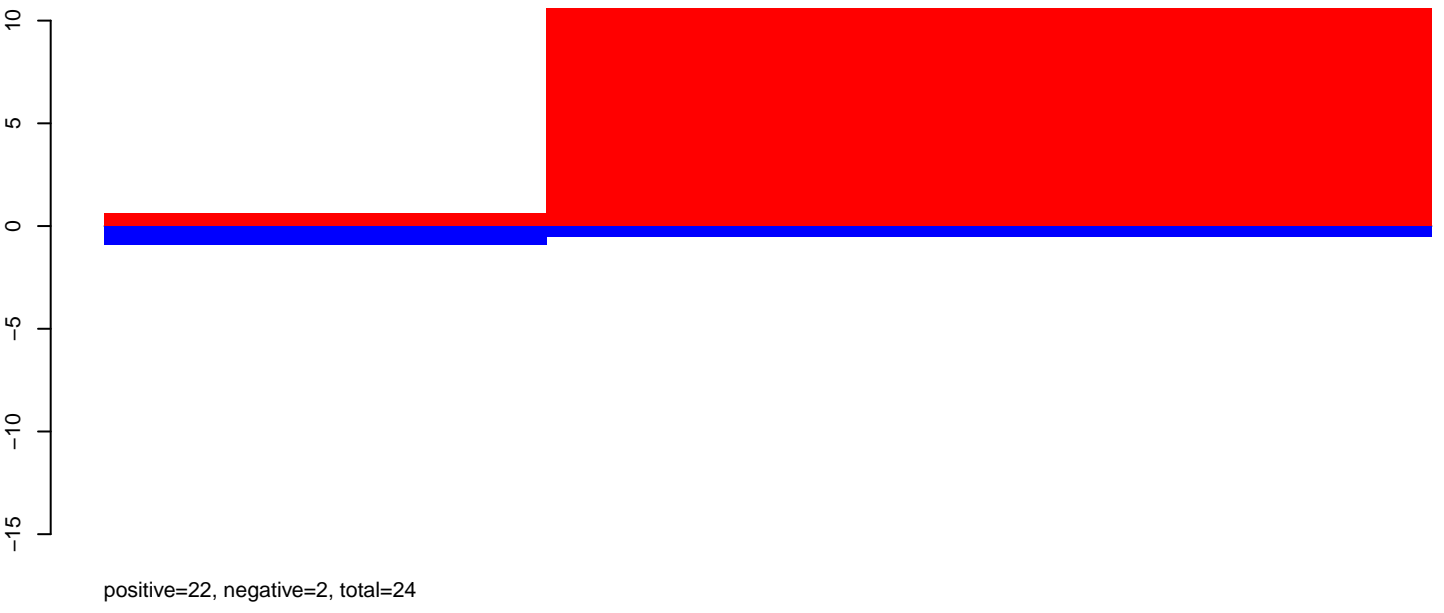


40 60 80 100
window size=25 length=72, AAEL028990-RA-tRNA-2-422623477-422623548-1-gene-AAEL028990-1-72

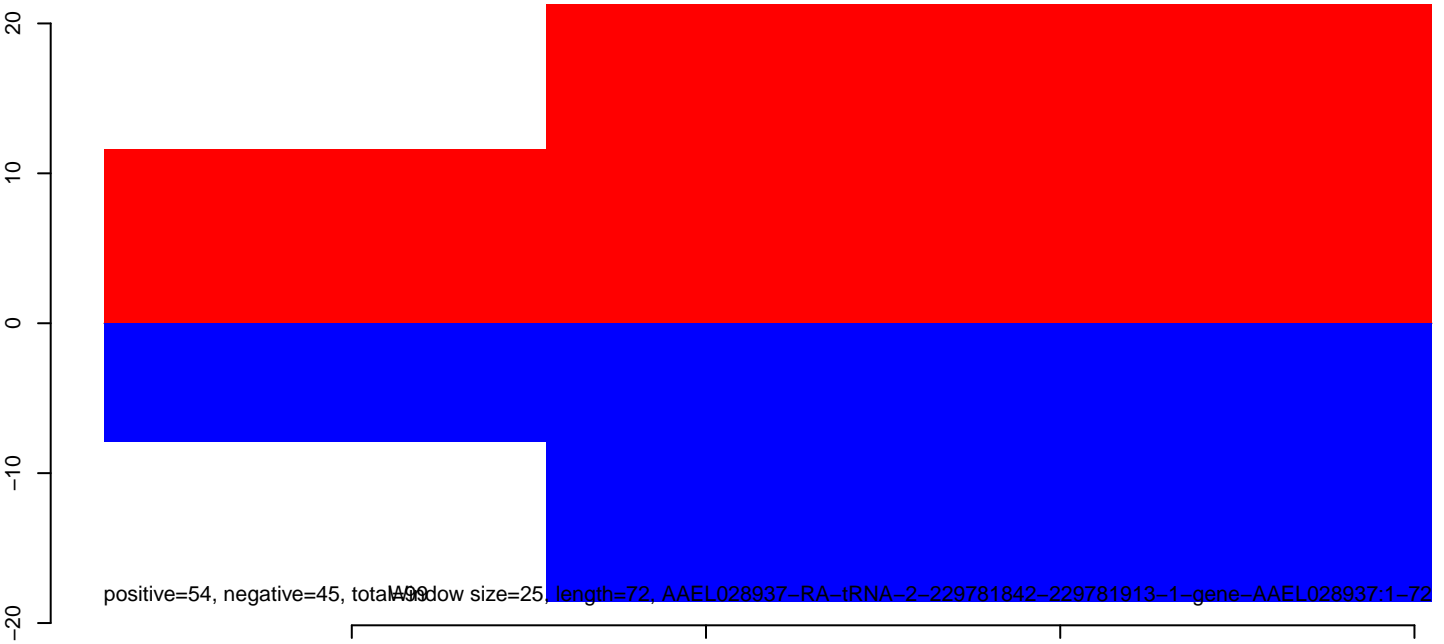
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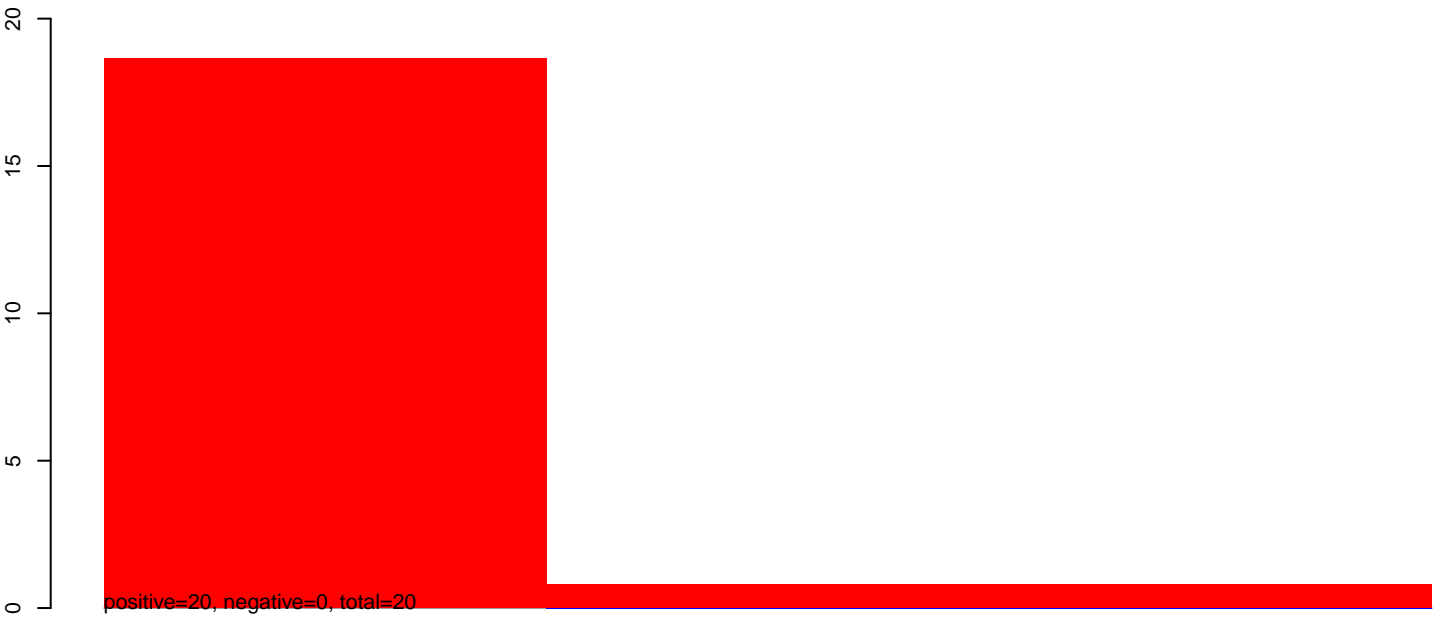
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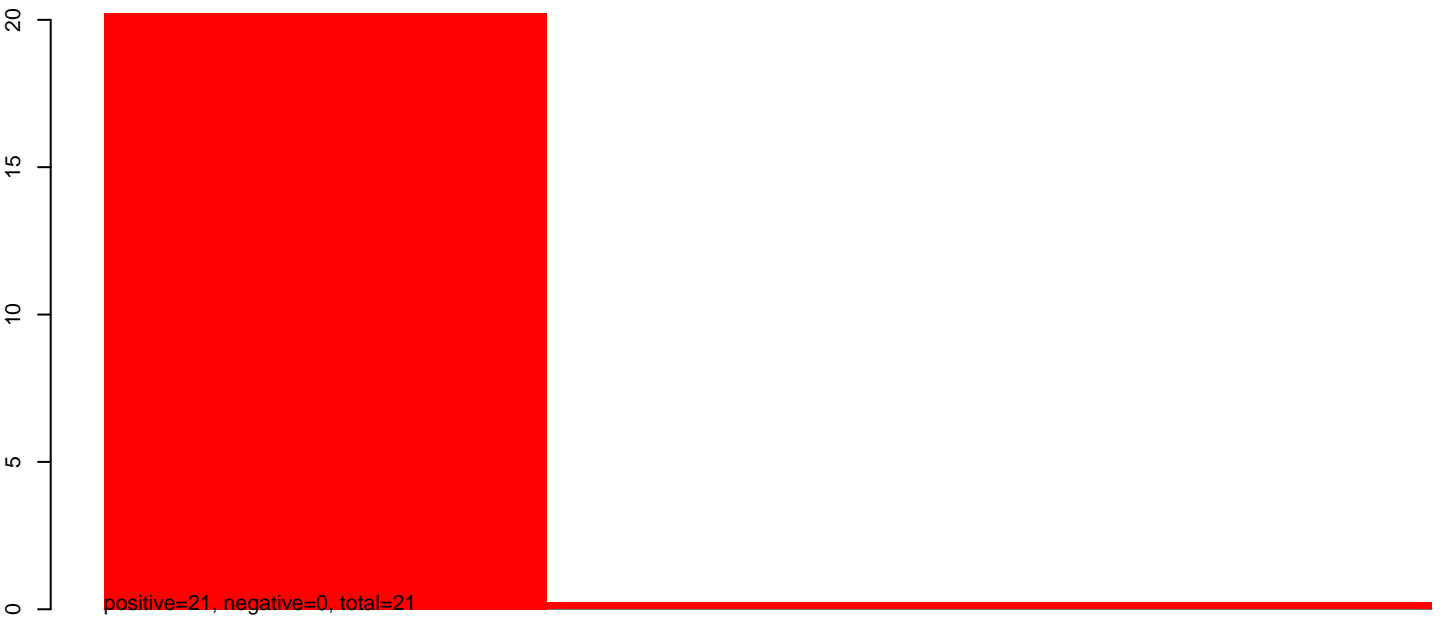
AeAeg_Aag2.TC.Dmem.rep



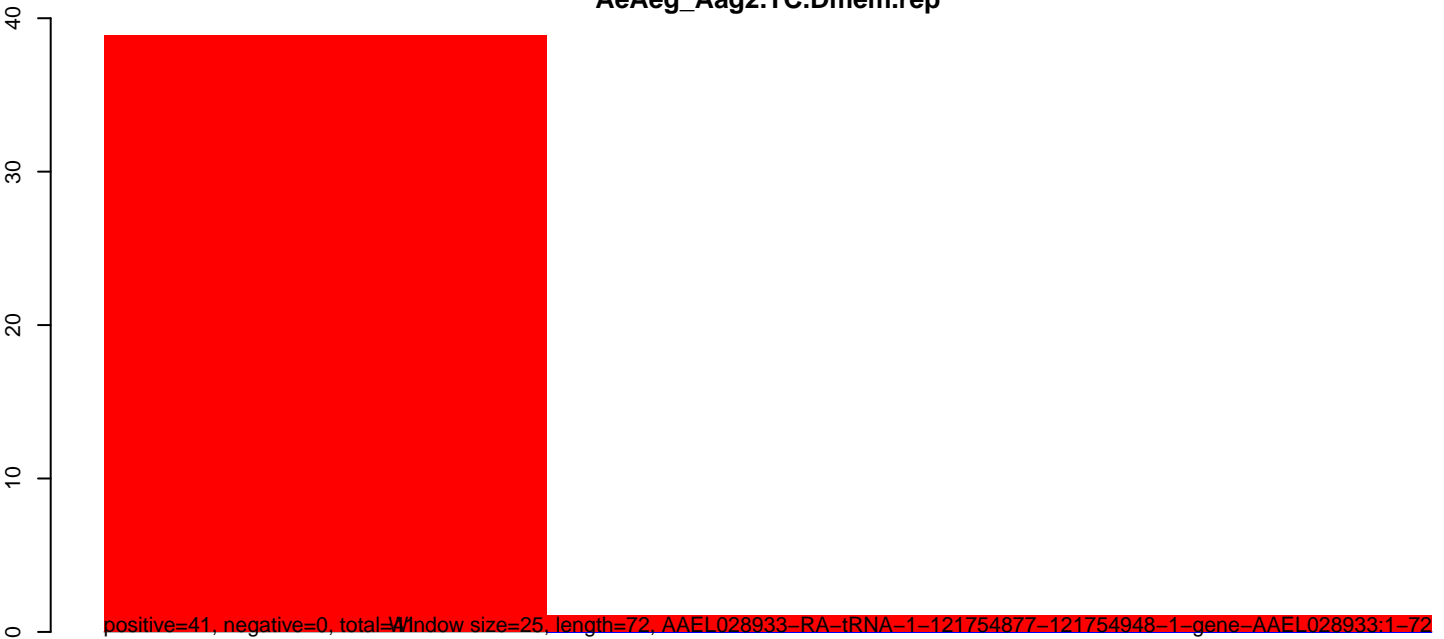
AeAeg_Aag2.TC.Dmem.18_23.rep



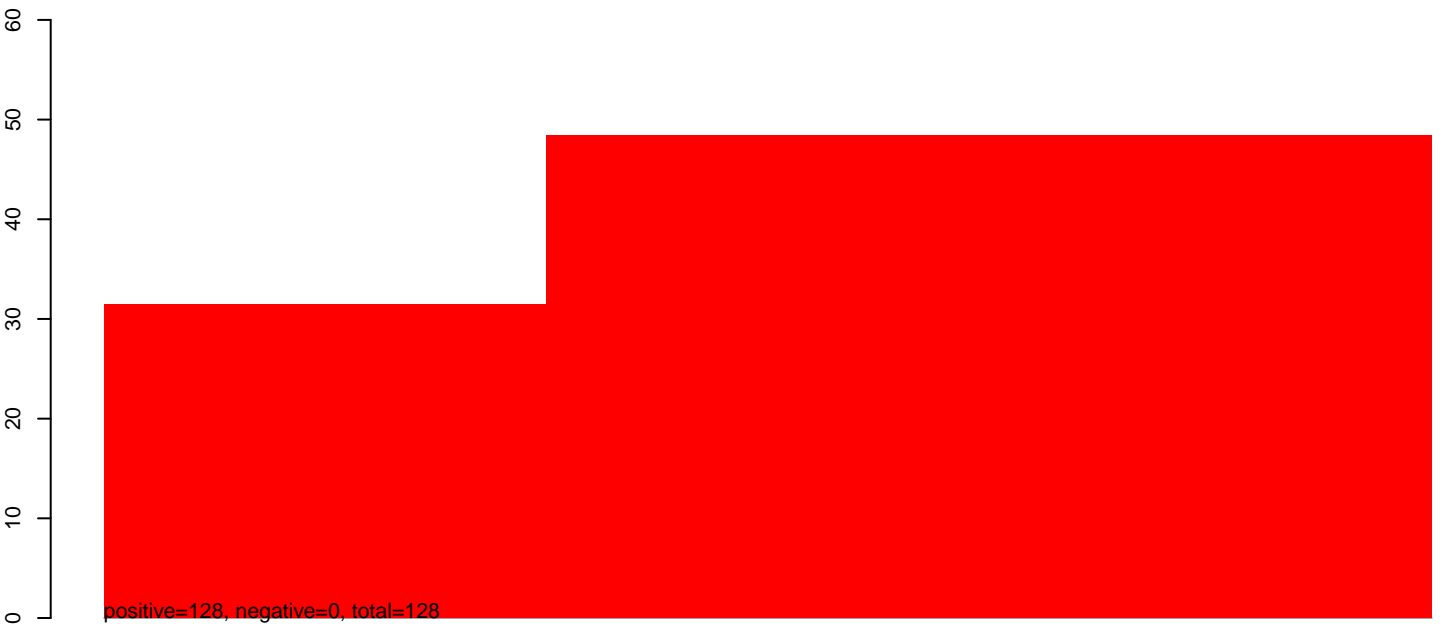
AeAeg_Aag2.TC.Dmem.24_35.rep



AeAeg_Aag2.TC.Dmem.rep



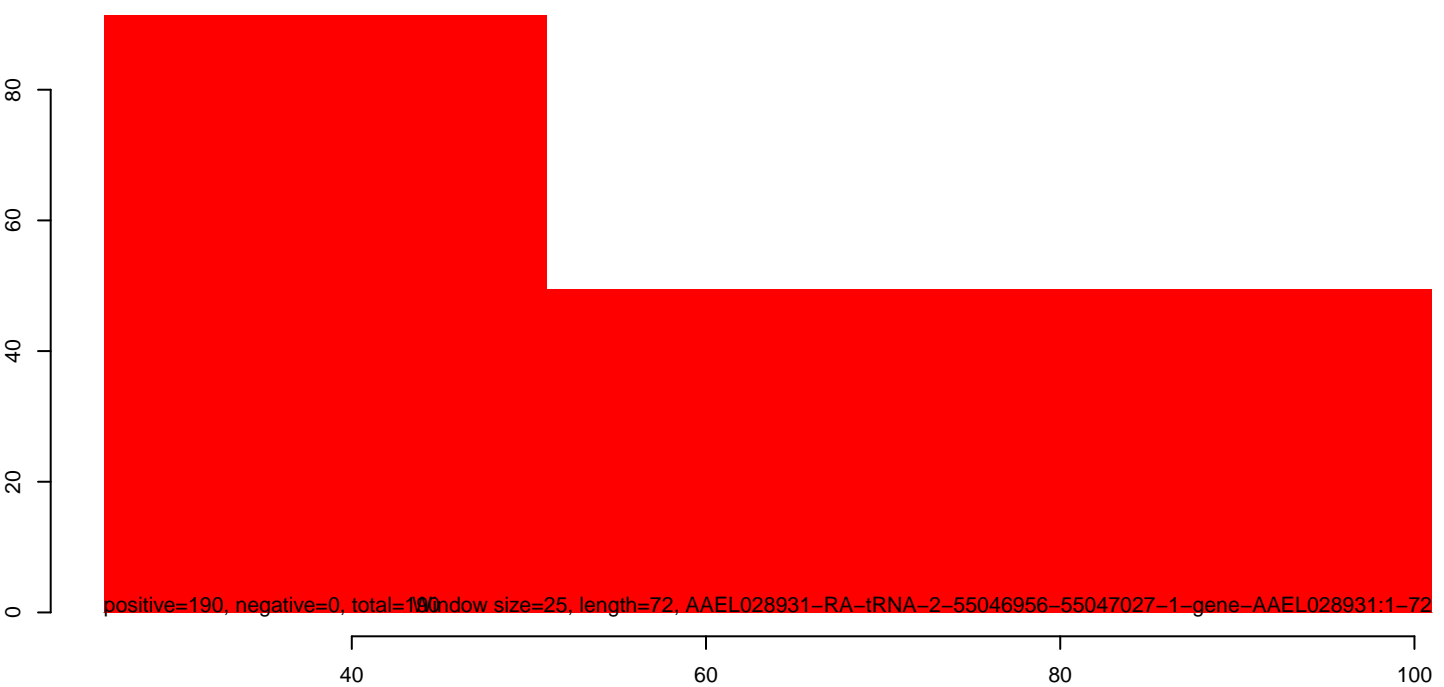
AeAeg_Aag2.TC.Dmem.18_23.rep



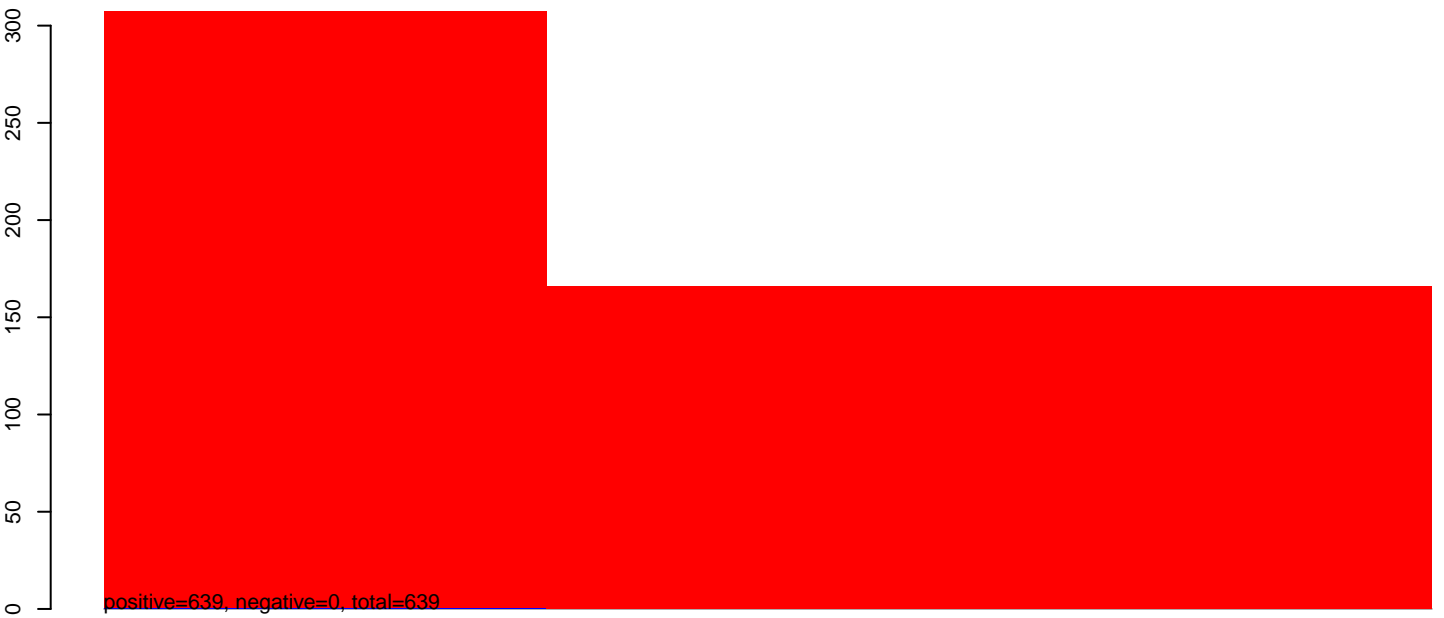
AeAeg_Aag2.TC.Dmem.24_35.rep



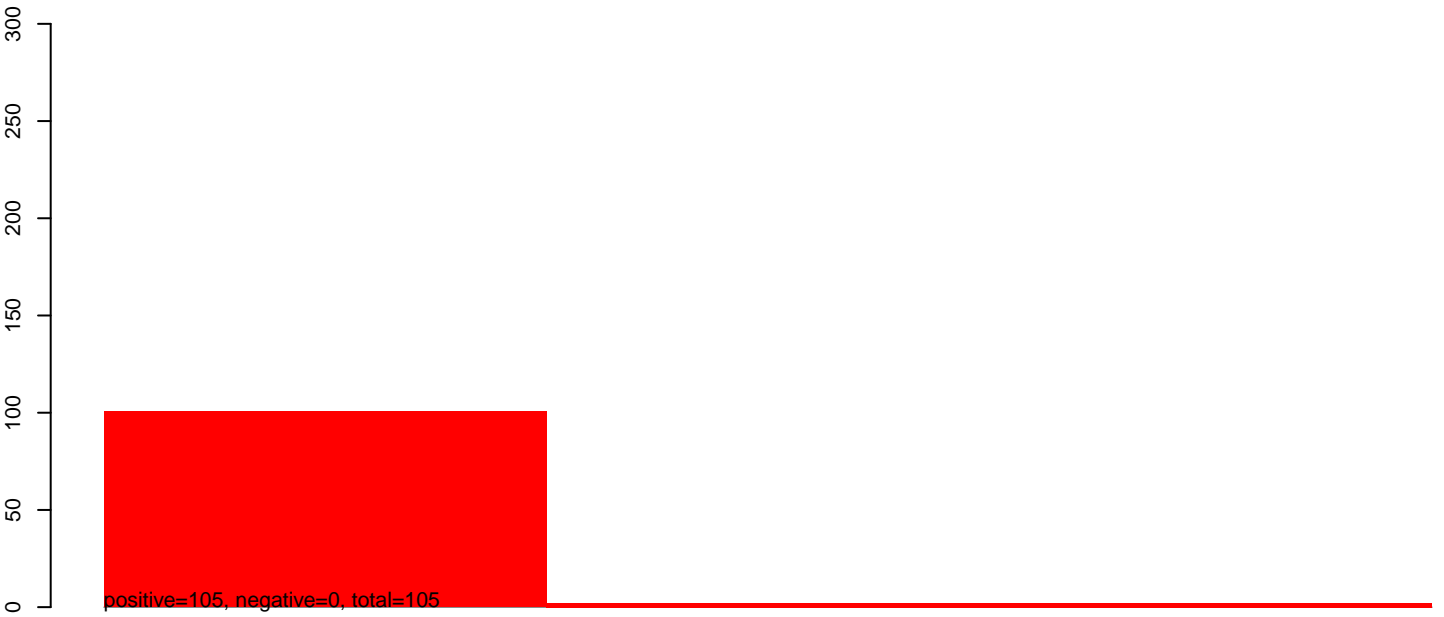
AeAeg_Aag2.TC.Dmem.rep



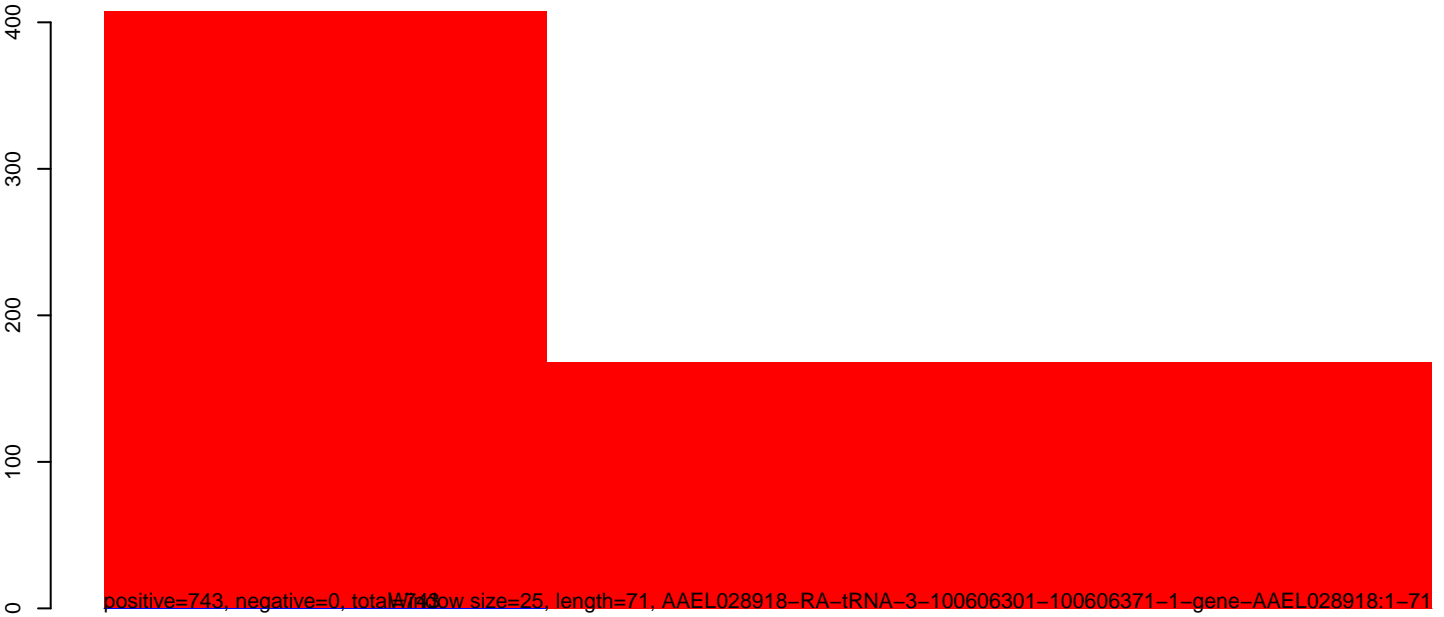
AeAeg_Aag2.TC.Dmem.18_23.rep



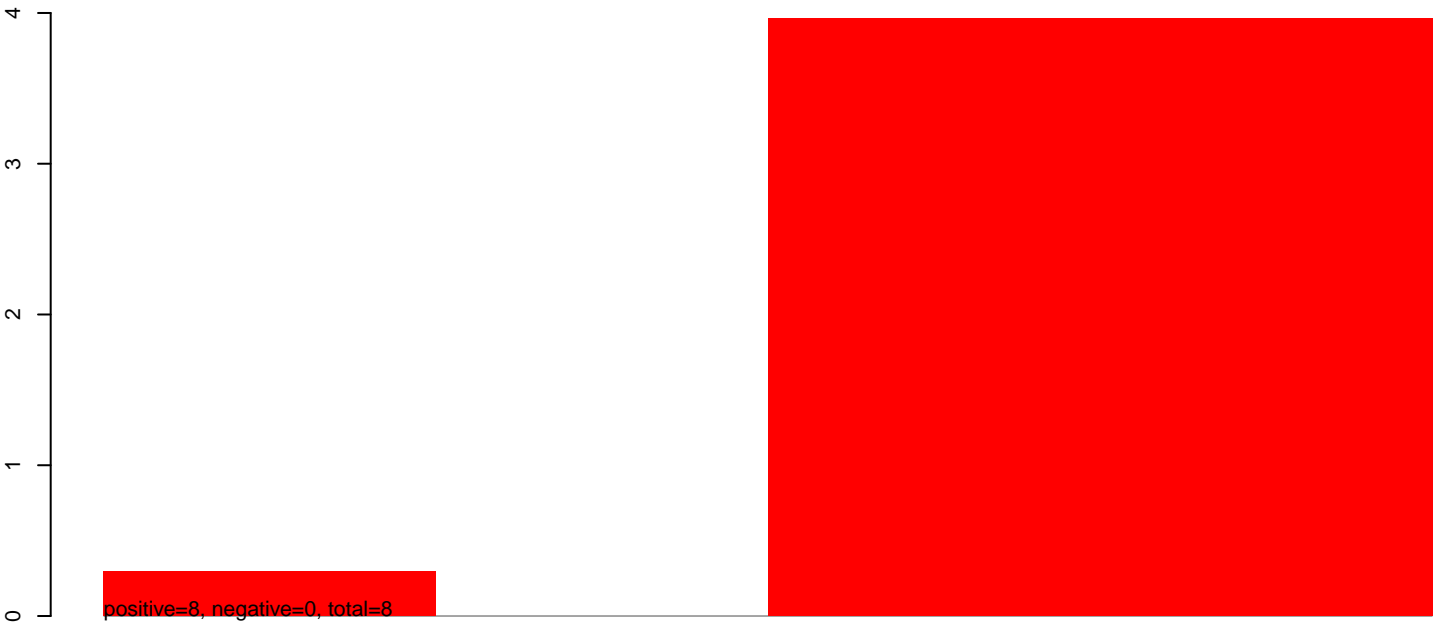
AeAeg_Aag2.TC.Dmem.24_35.rep



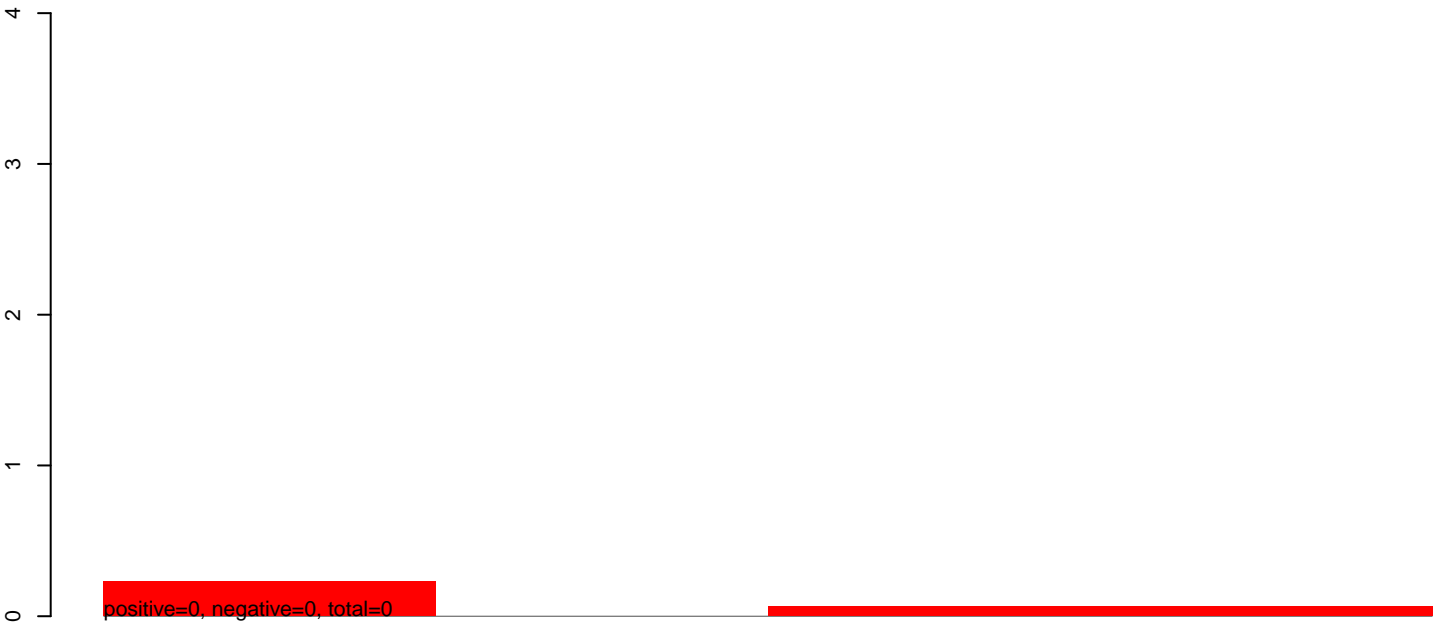
AeAeg_Aag2.TC.Dmem.rep



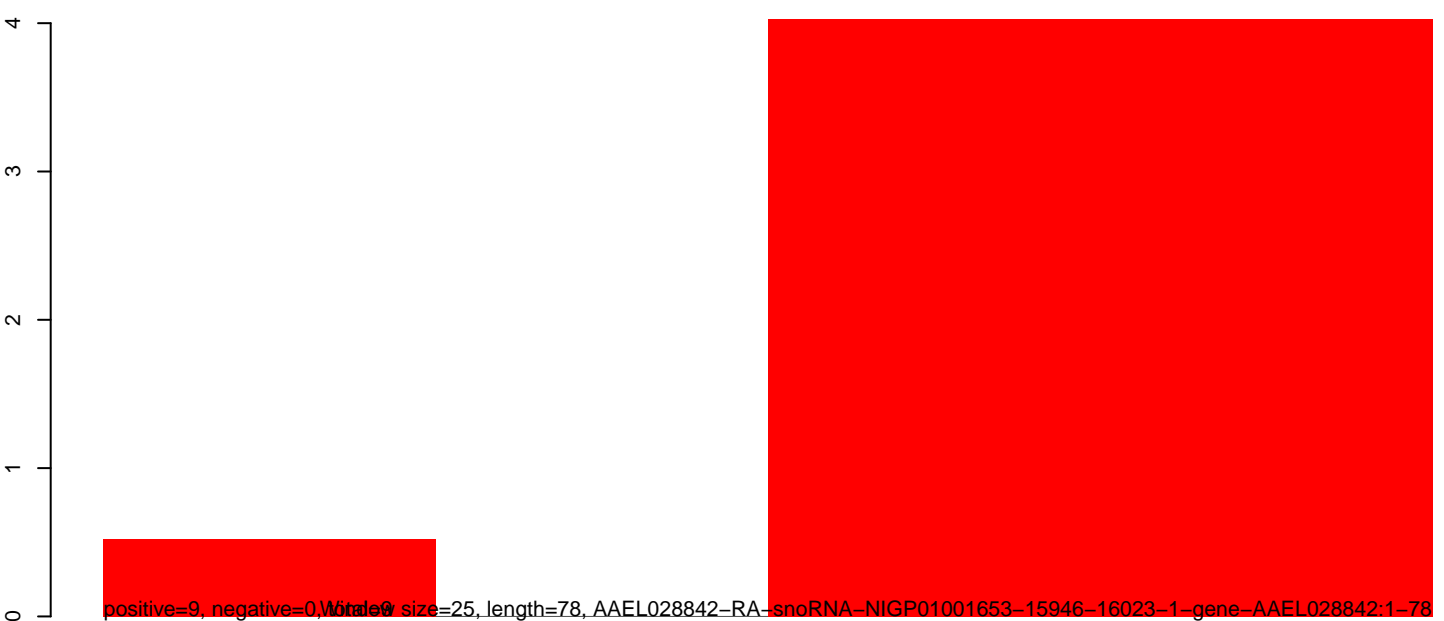
AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep



AeAeg_Aag2.TC.Dmem.rep



40 60 80 100 120

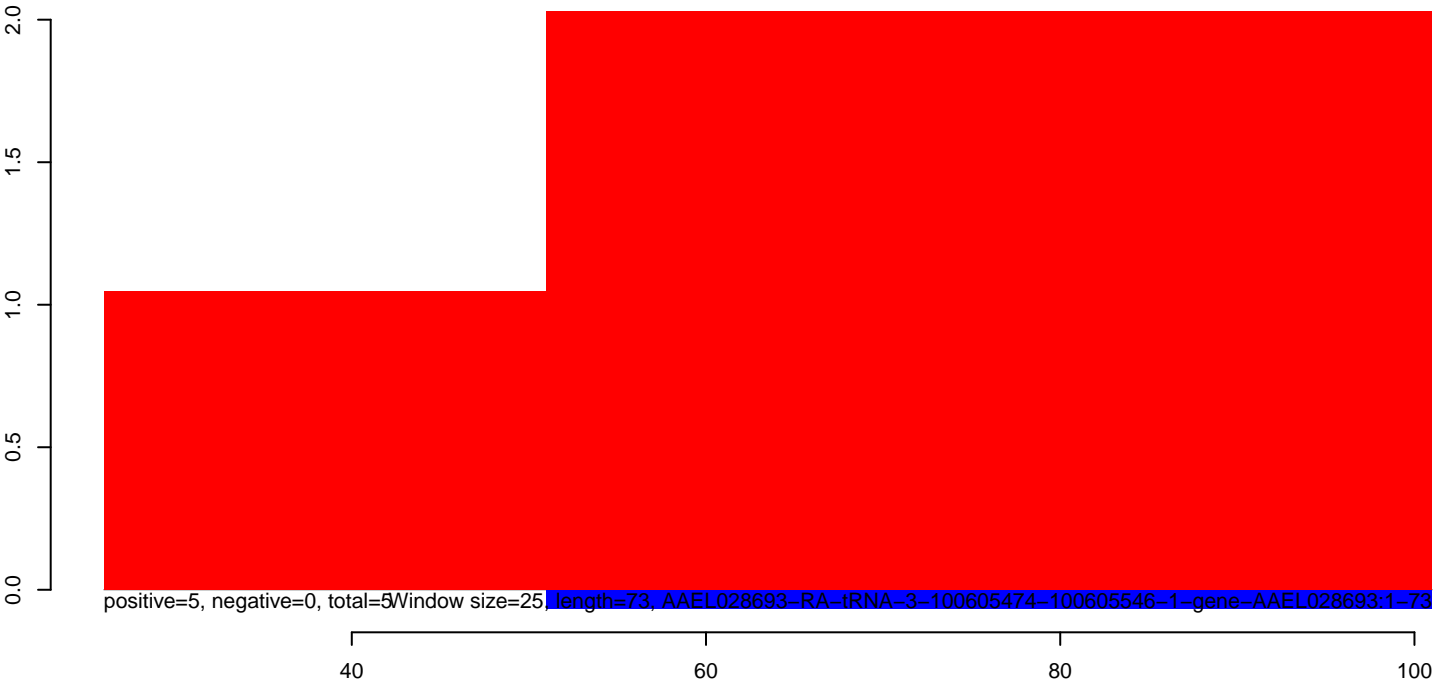
AeAeg_Aag2.TC.Dmem.18_23.rep



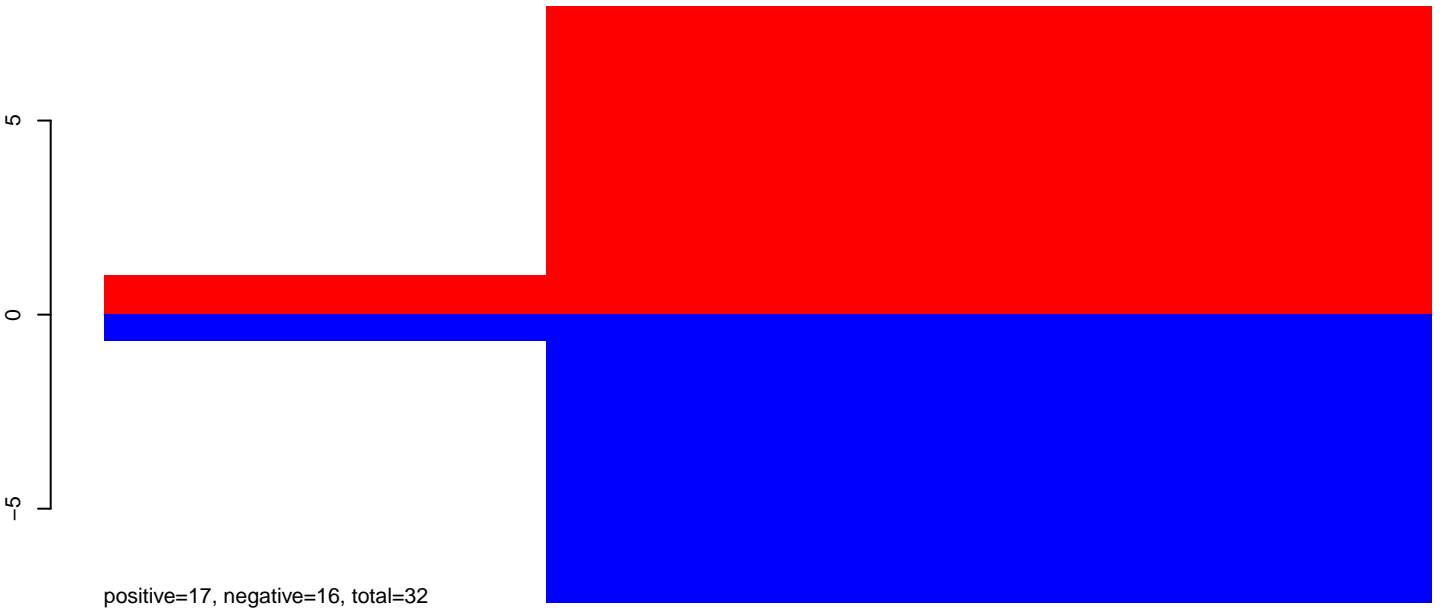
AeAeg_Aag2.TC.Dmem.24_35.rep



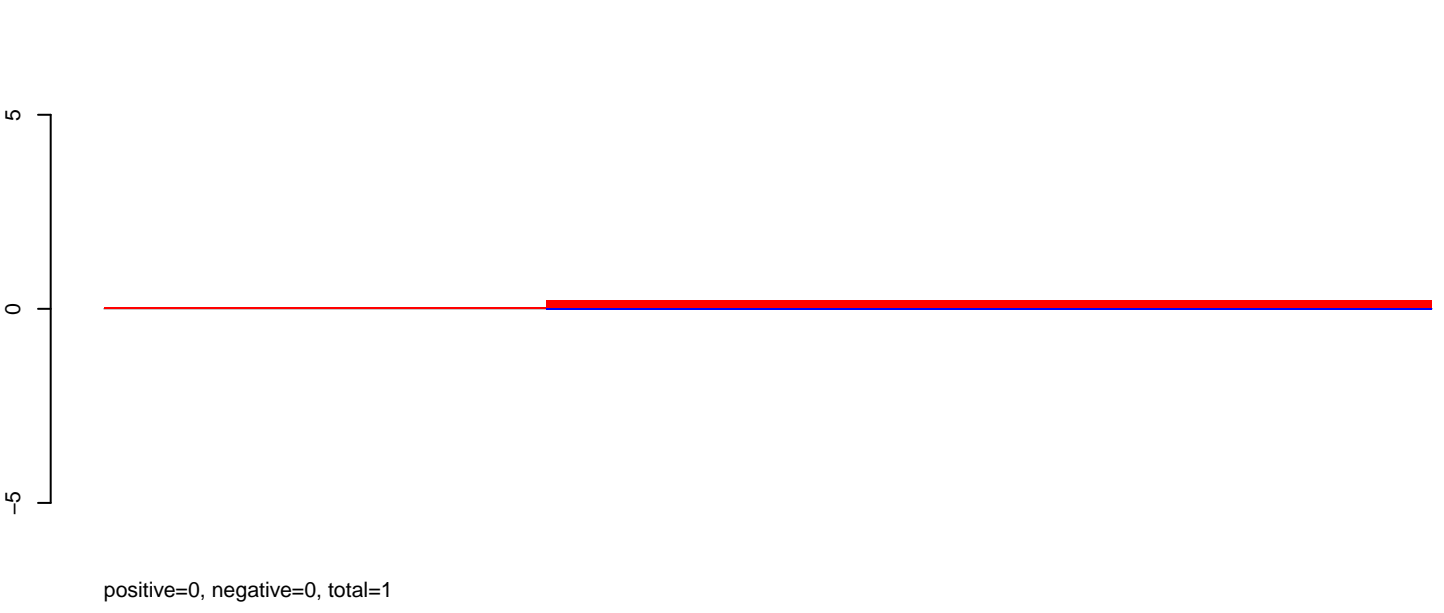
AeAeg_Aag2.TC.Dmem.rep



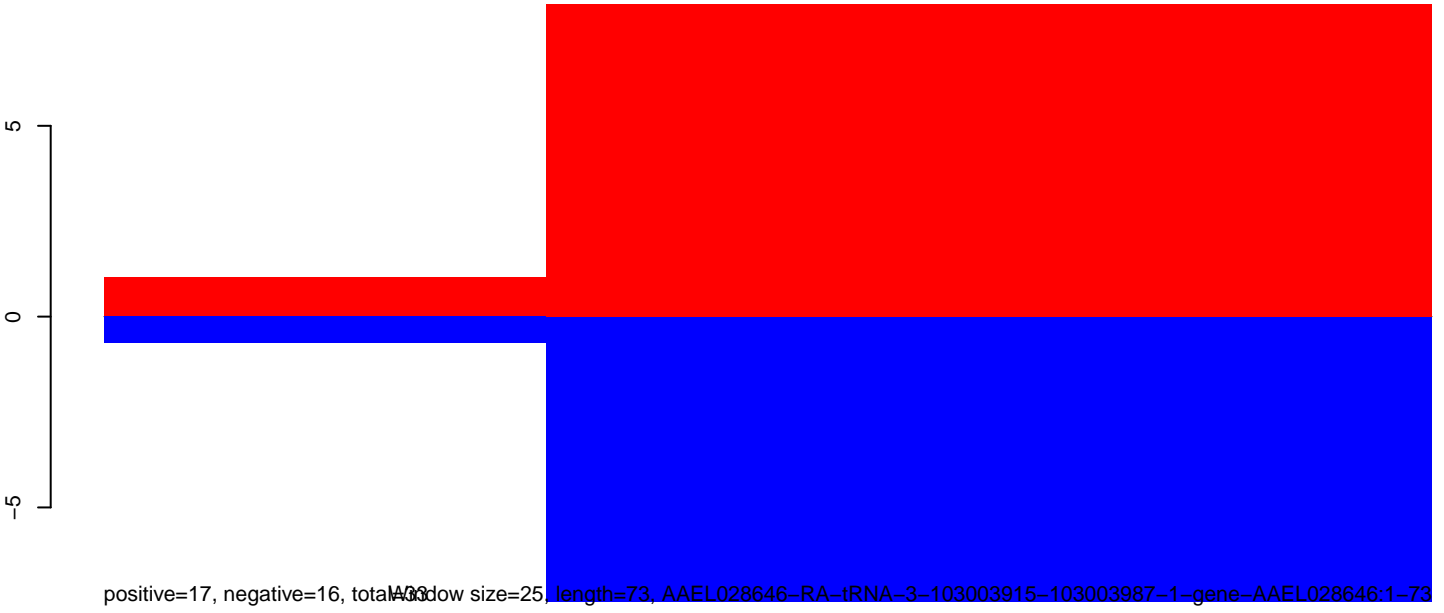
AeAeg_Aag2.TC.Dmem.18_23.rep



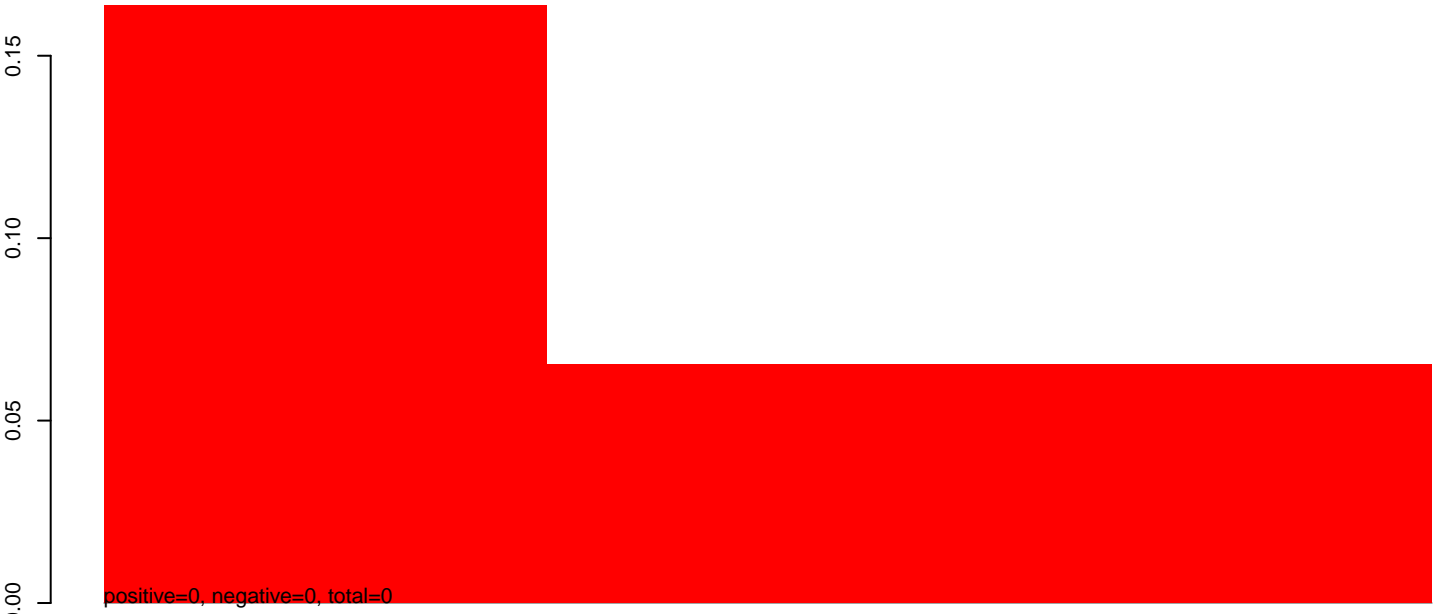
AeAeg_Aag2.TC.Dmem.24_35.rep



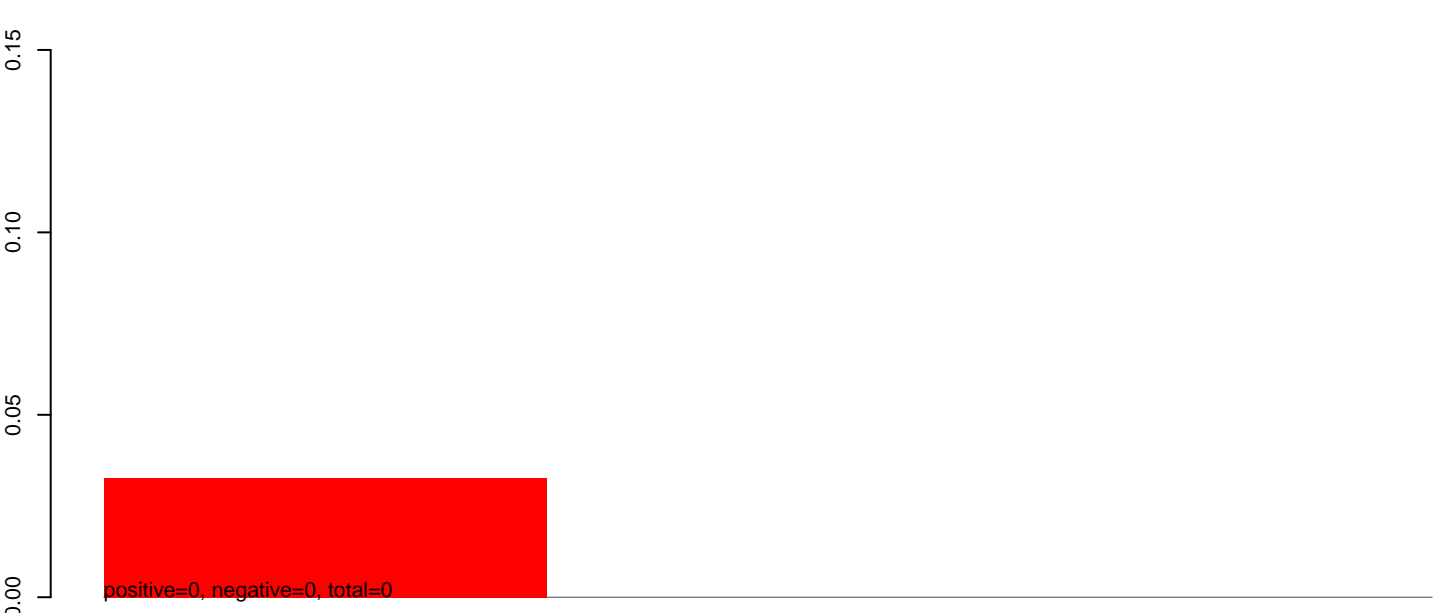
AeAeg_Aag2.TC.Dmem.rep



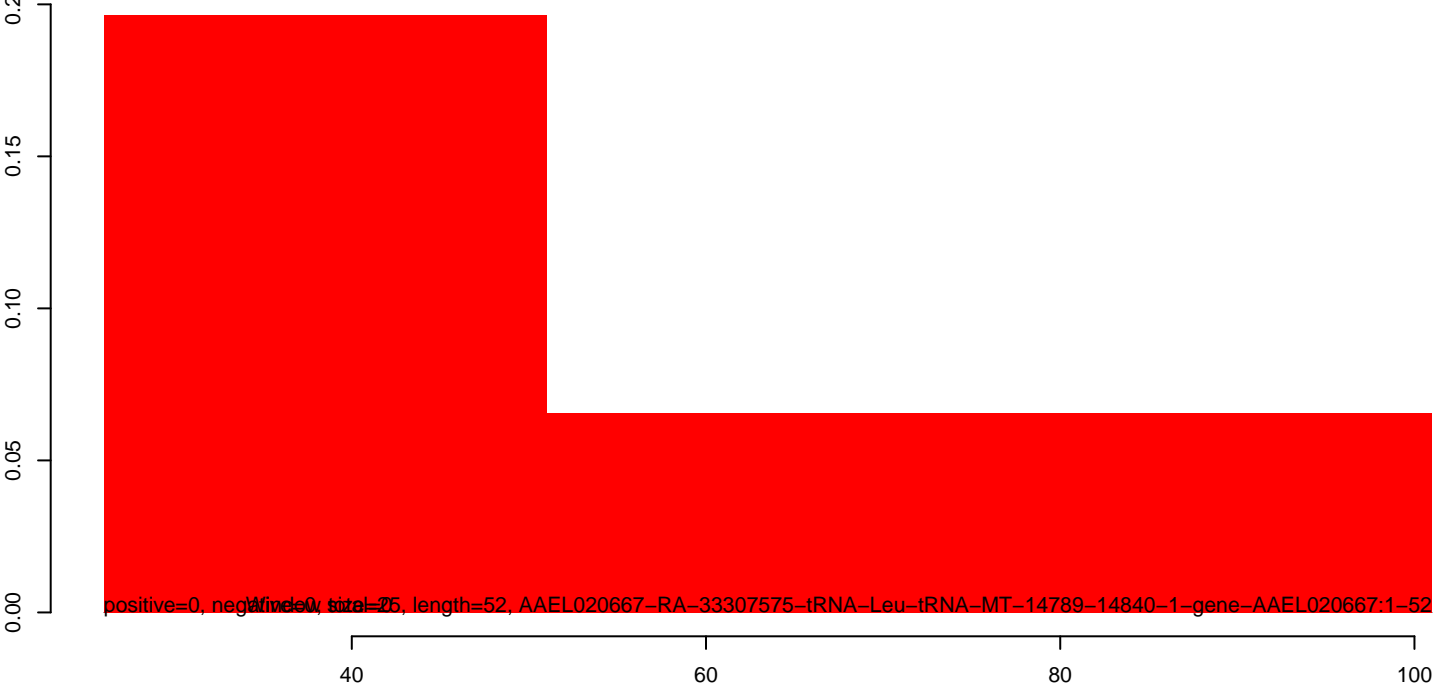
AeAeg_Aag2.TC.Dmem.18_23.rep



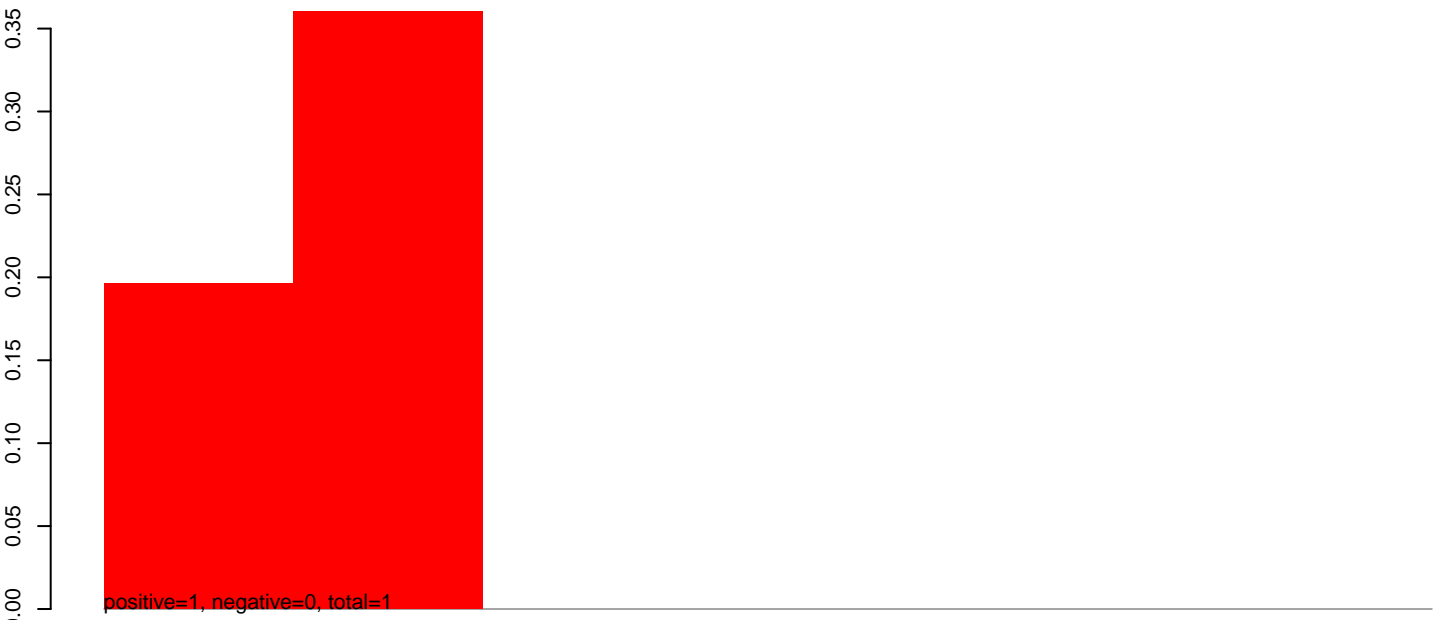
AeAeg_Aag2.TC.Dmem.24_35.rep



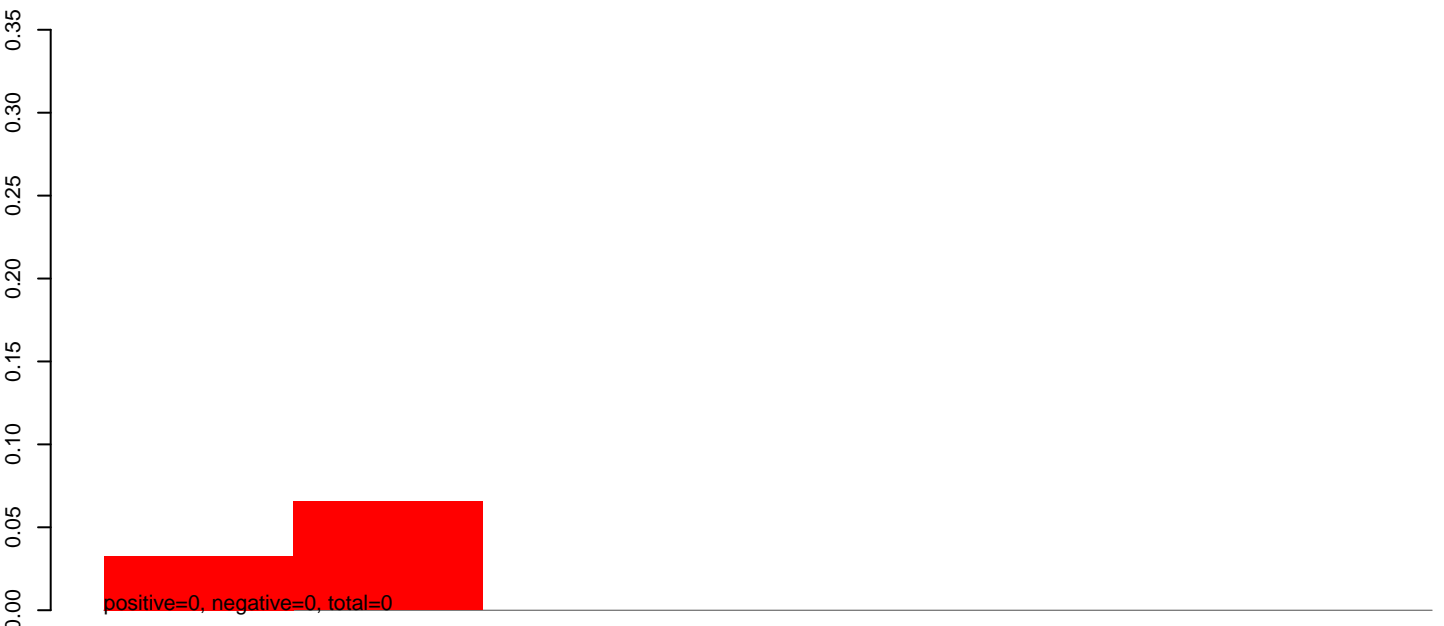
AeAeg_Aag2.TC.Dmem.rep



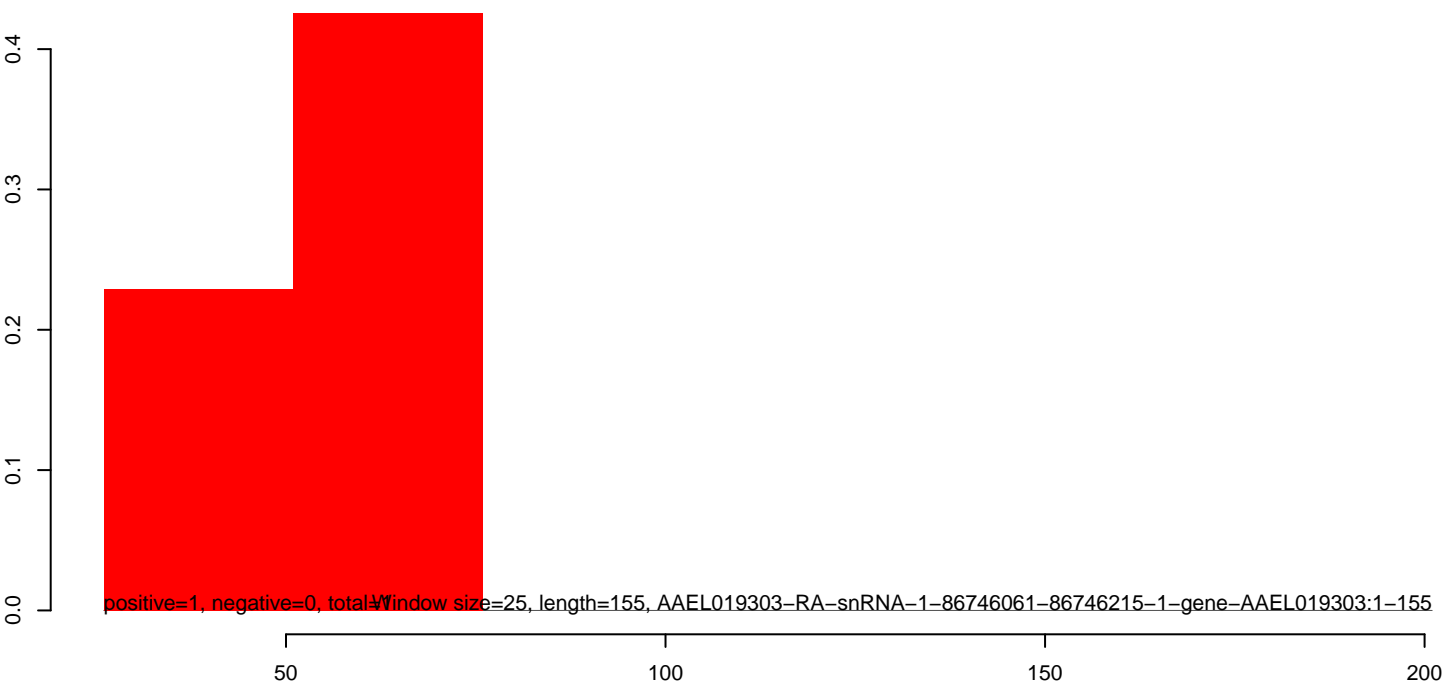
AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep

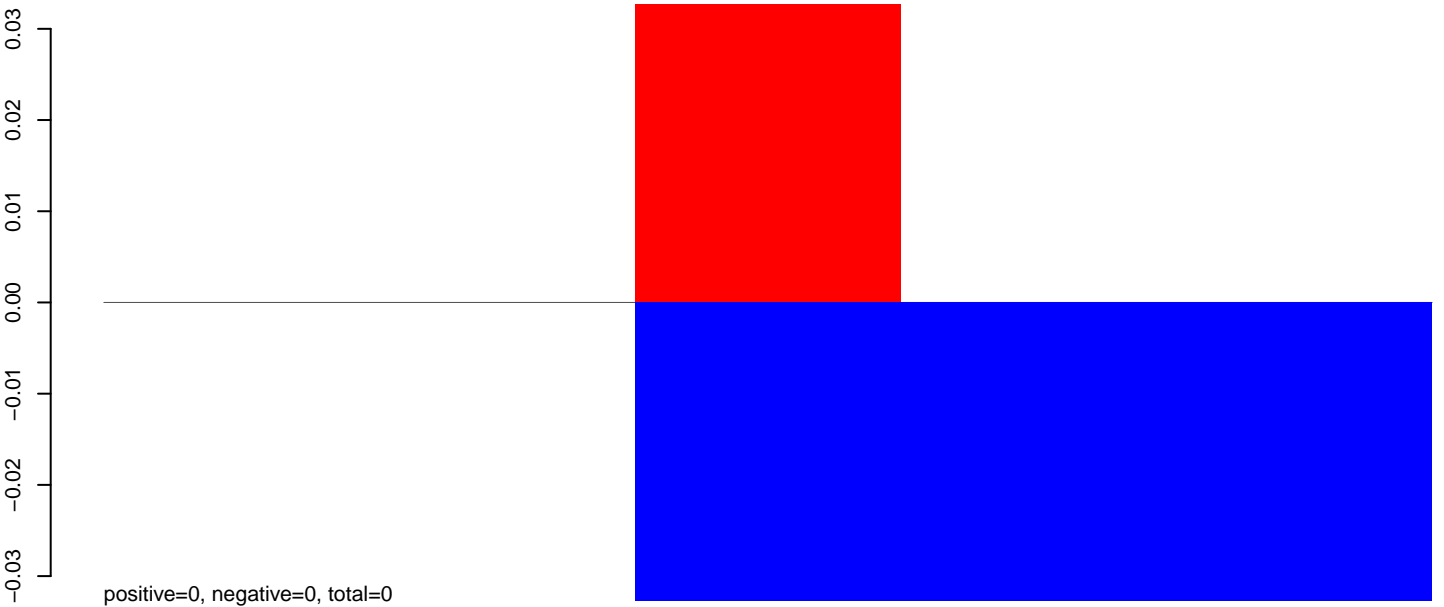


AeAeg_Aag2.TC.Dmem.rep

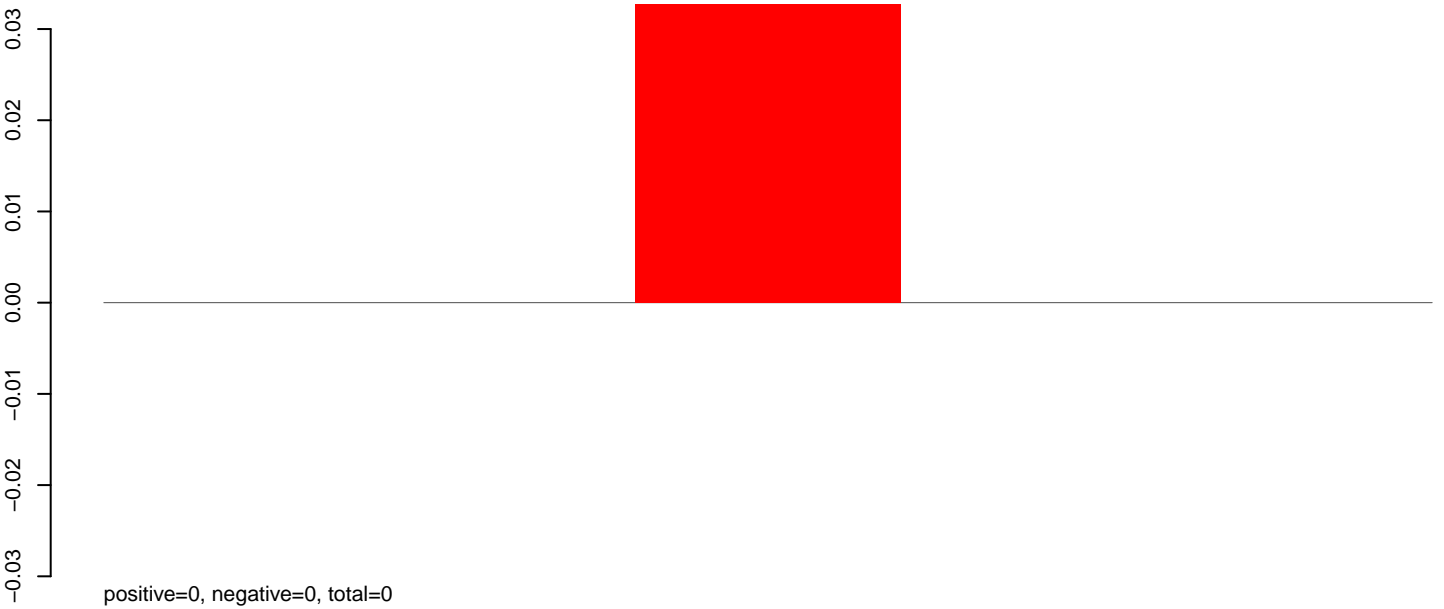


Window size=25, length=155, AAEL019303-RA-snrRNA-1-86746061-86746215-1-gene-AAEL019303:1-155

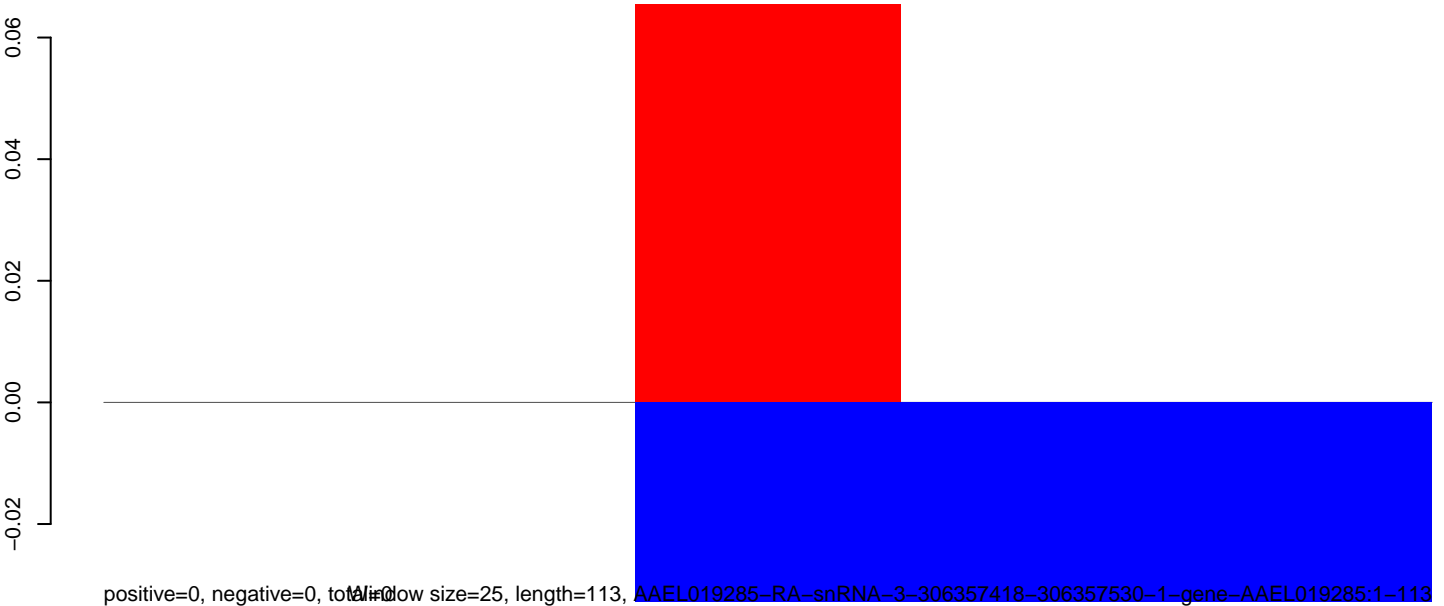
AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep



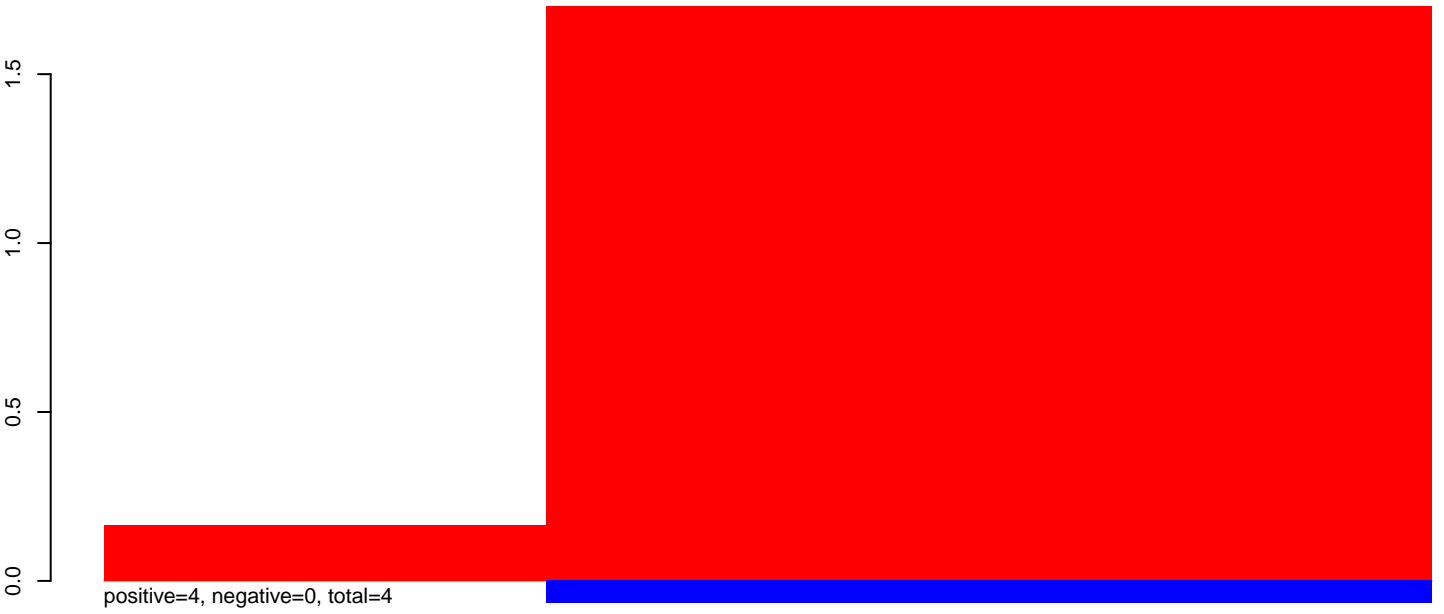
AeAeg_Aag2.TC.Dmem.rep



40 60 80 100 120 140

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

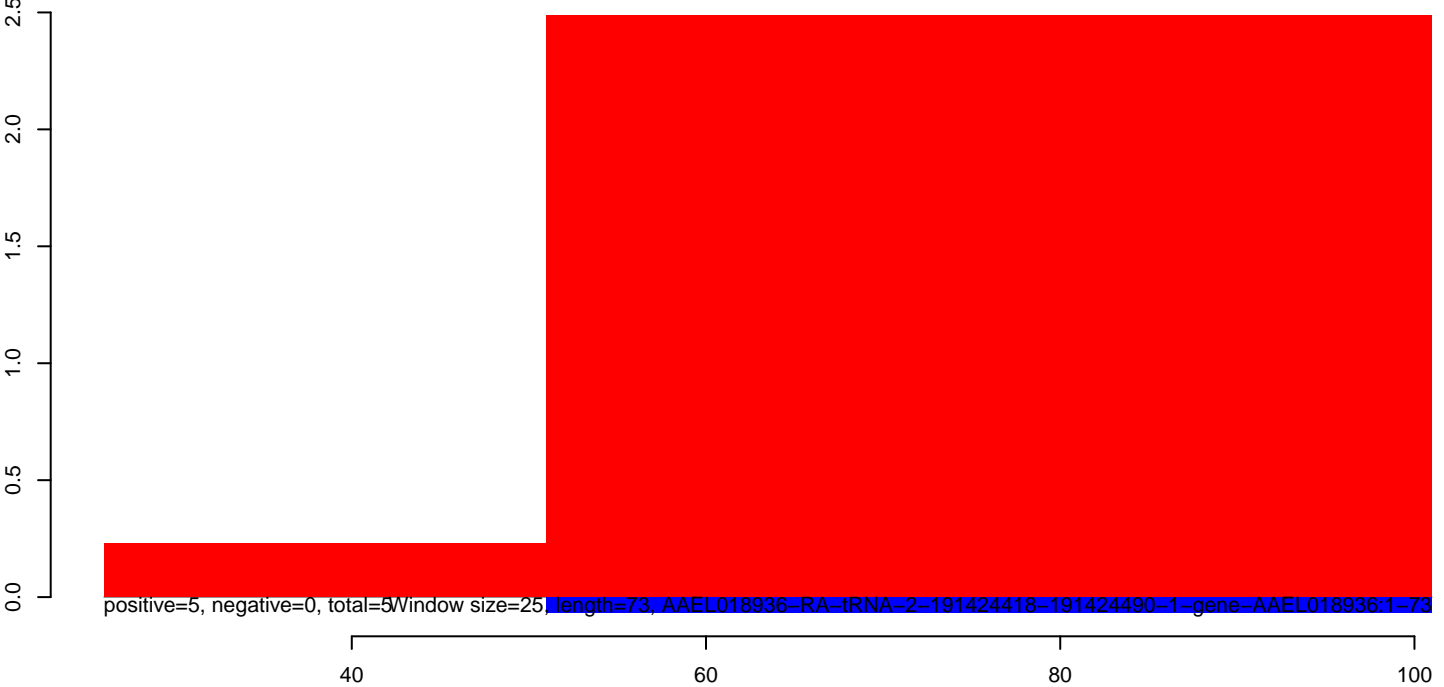
AeAeg_Aag2.TC.Dmem.18_23.rep



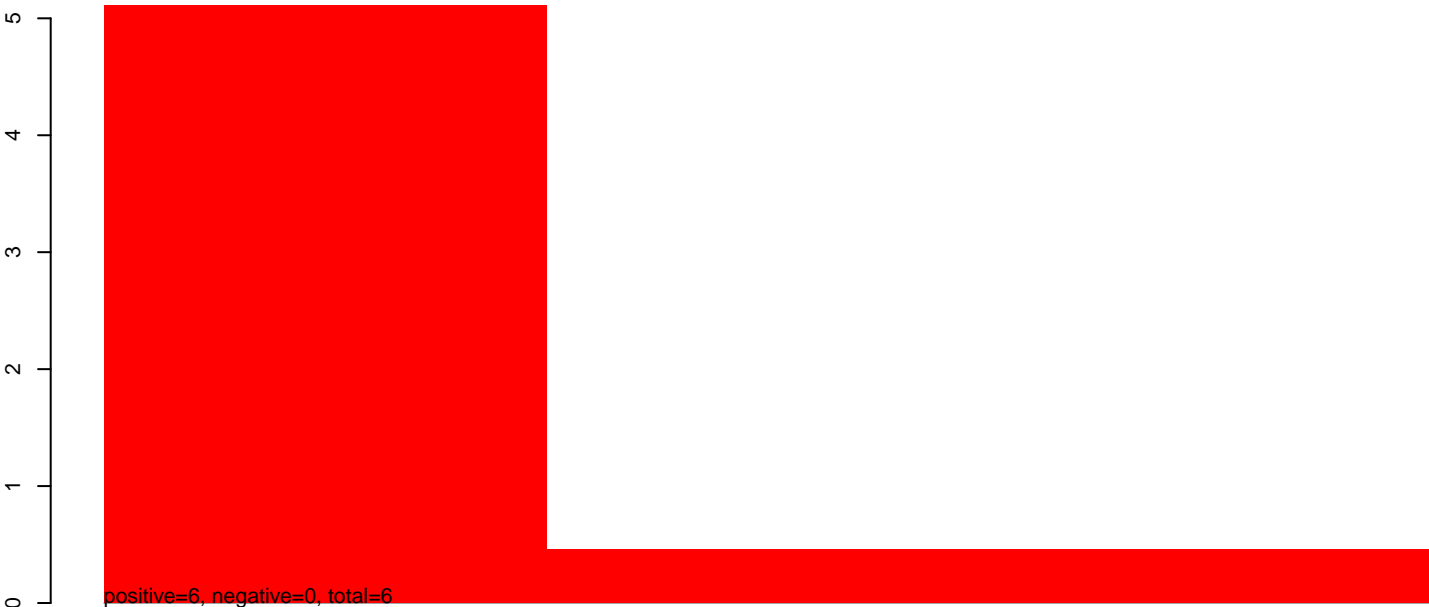
AeAeg_Aag2.TC.Dmem.24_35.rep



AeAeg_Aag2.TC.Dmem.rep



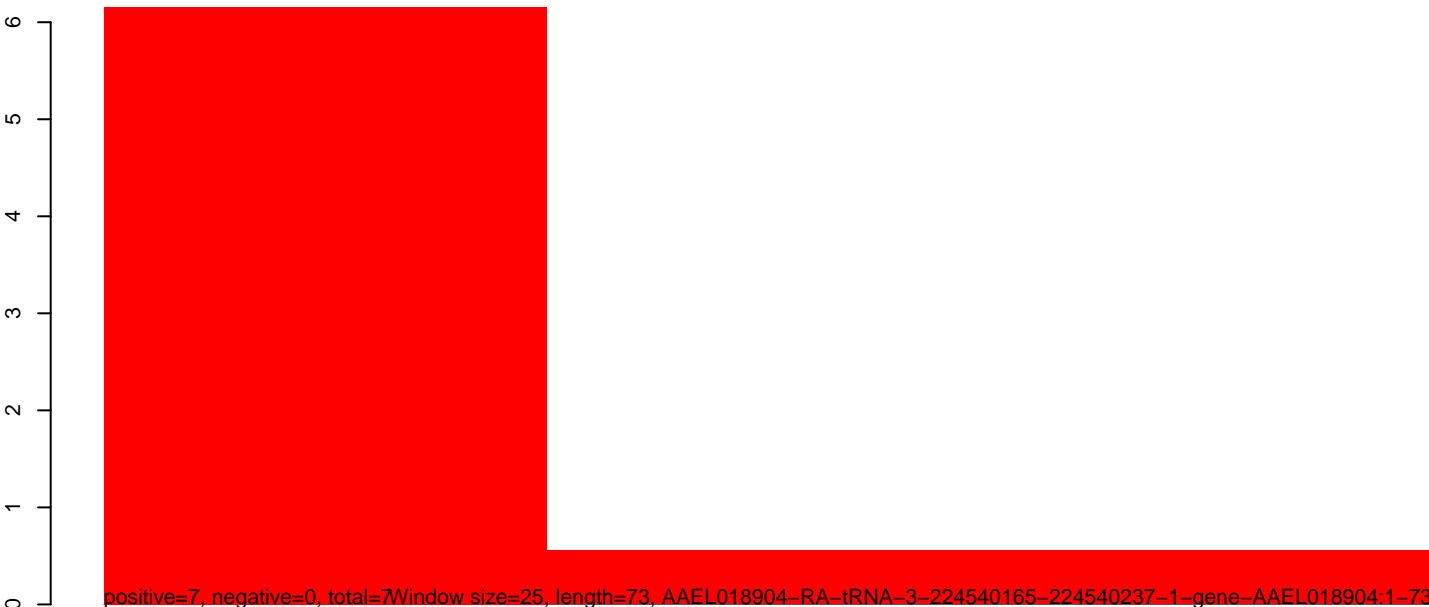
AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep



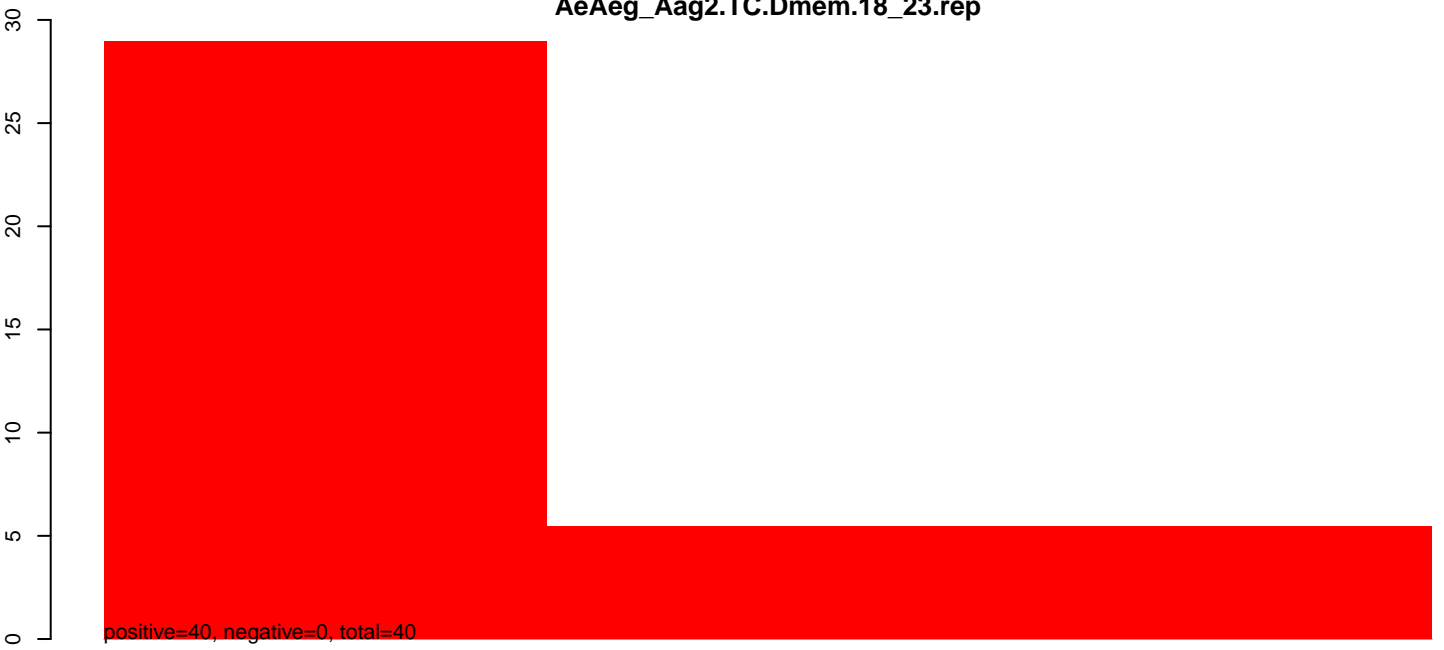
AeAeg_Aag2.TC.Dmem.rep



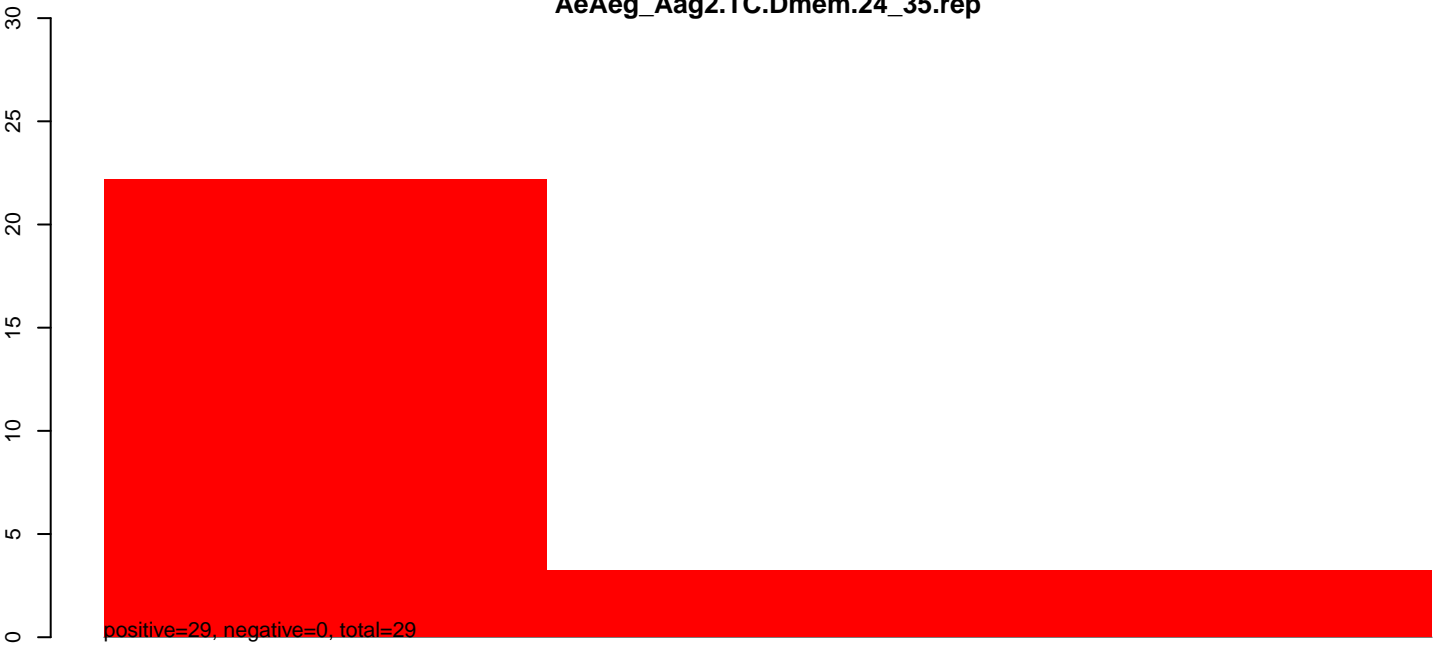
Window size=25, length=73, AAEL018904-RA-tRNA-3-224540165-224540237-1-gene-AAEL018904:1-73

40 60 80 100

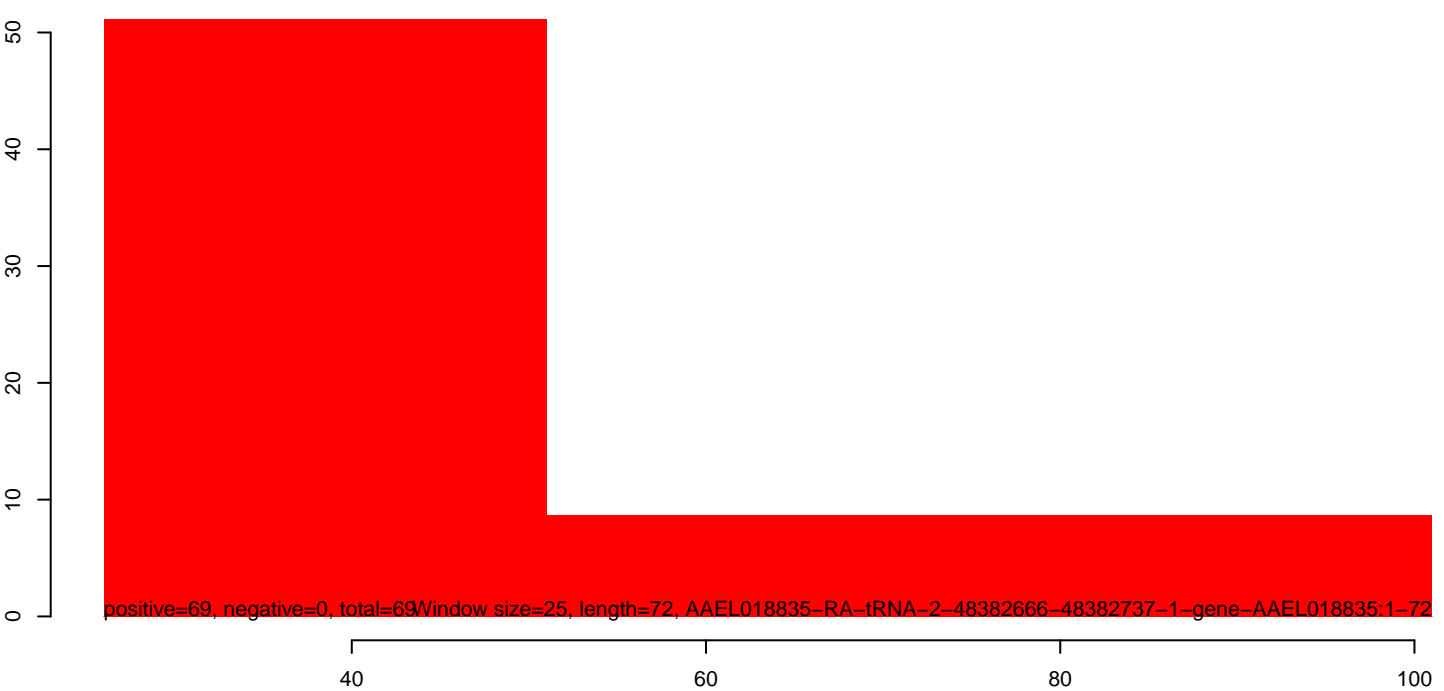
AeAeg_Aag2.TC.Dmem.18_23.rep



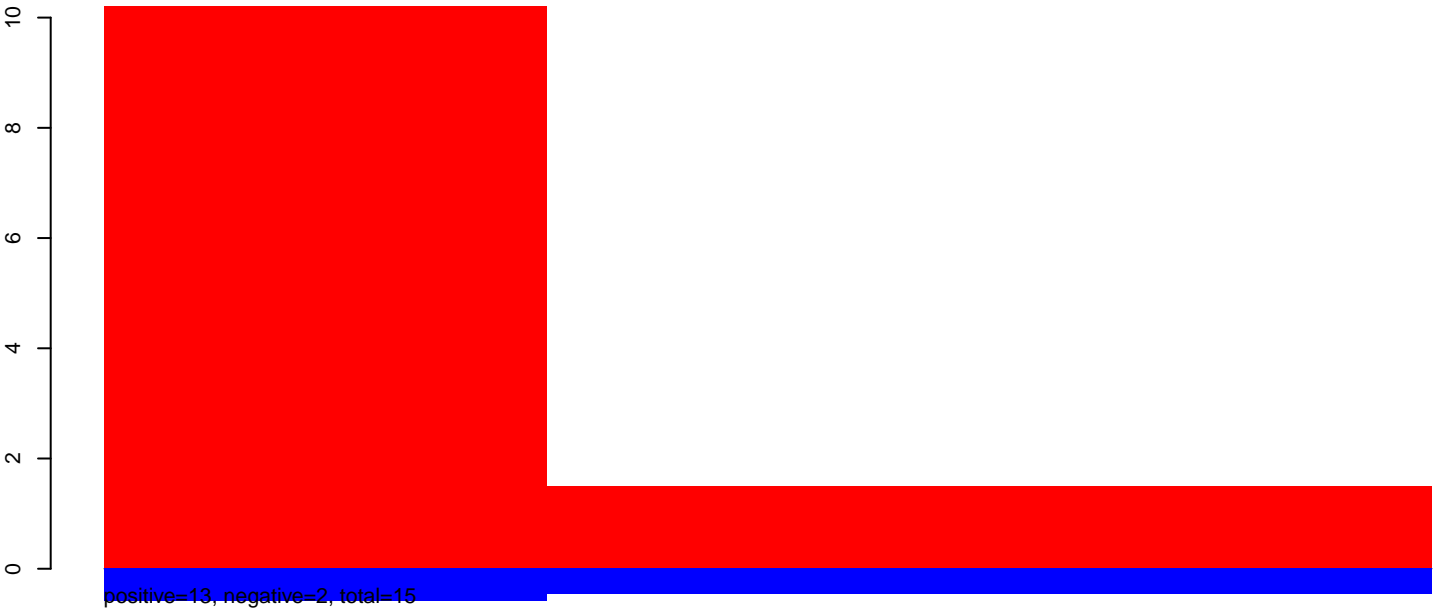
AeAeg_Aag2.TC.Dmem.24_35.rep



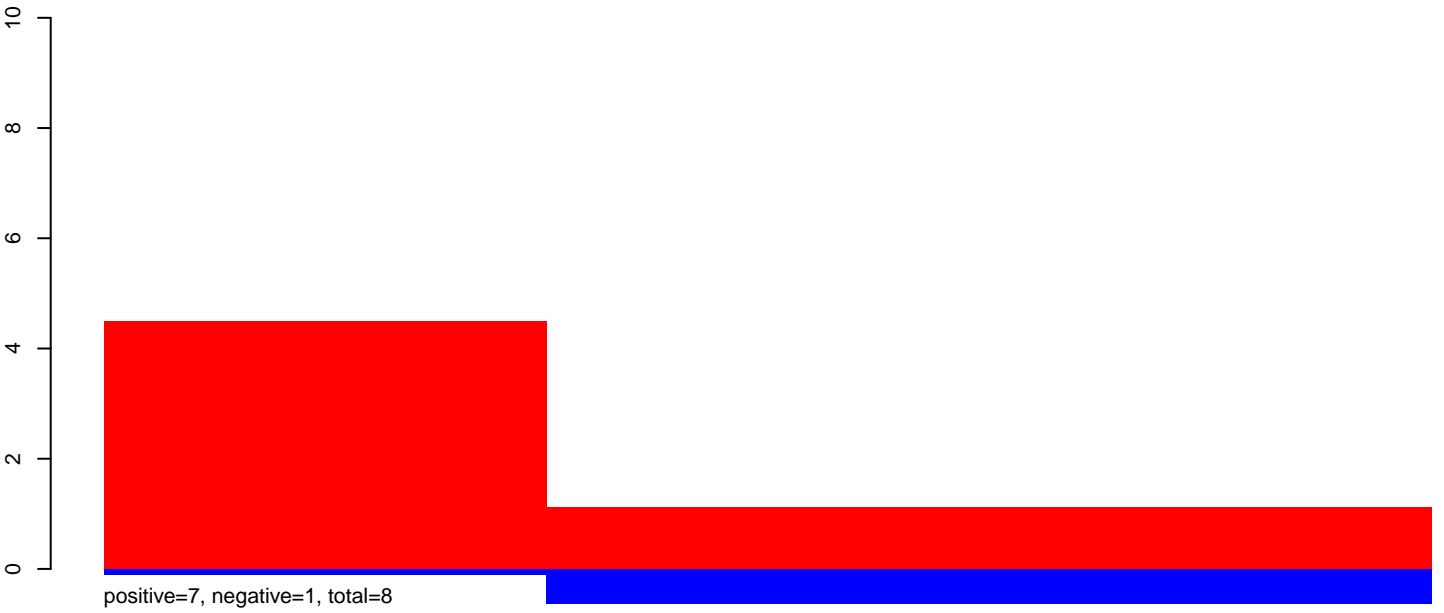
AeAeg_Aag2.TC.Dmem.rep



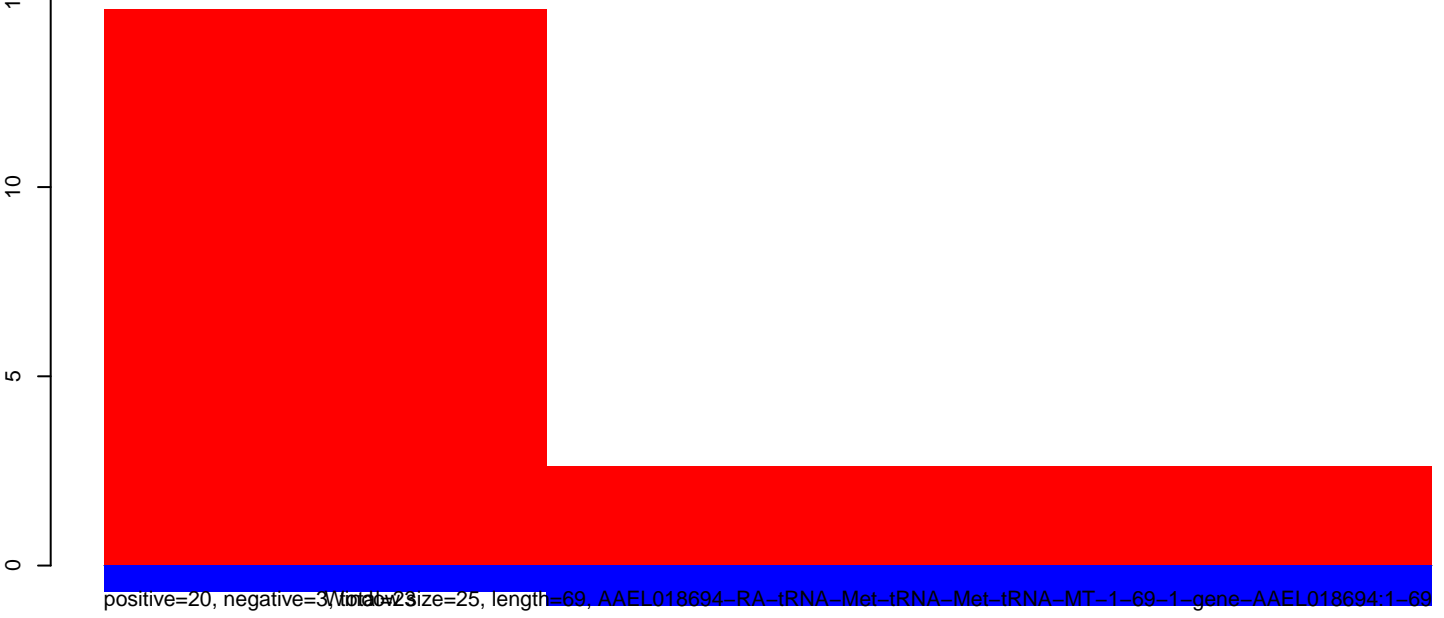
AeAeg_Aag2.TC.Dmem.18_23.rep



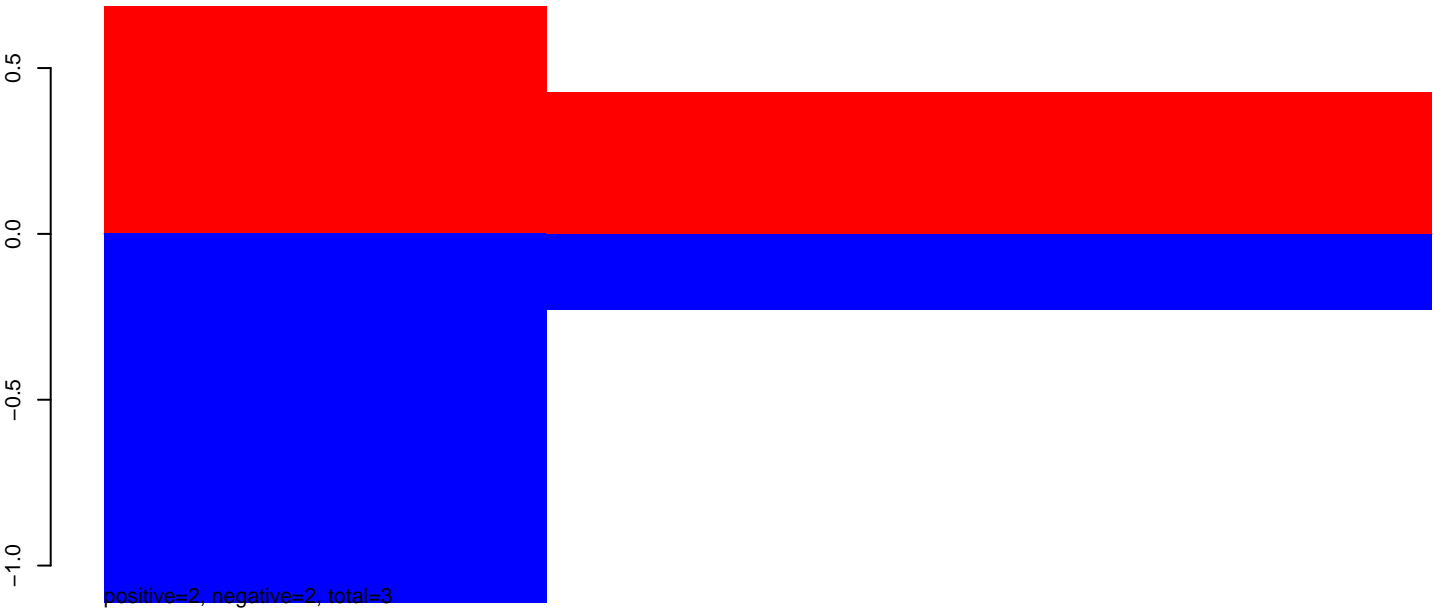
AeAeg_Aag2.TC.Dmem.24_35.rep



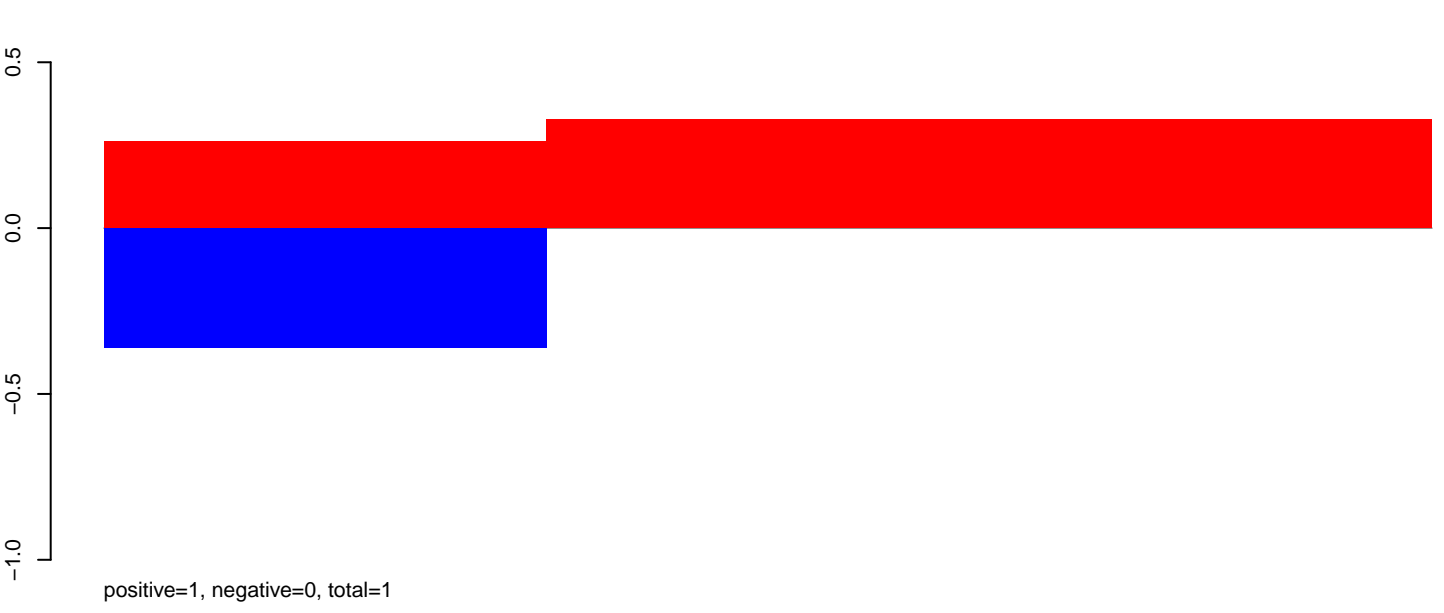
AeAeg_Aag2.TC.Dmem.rep



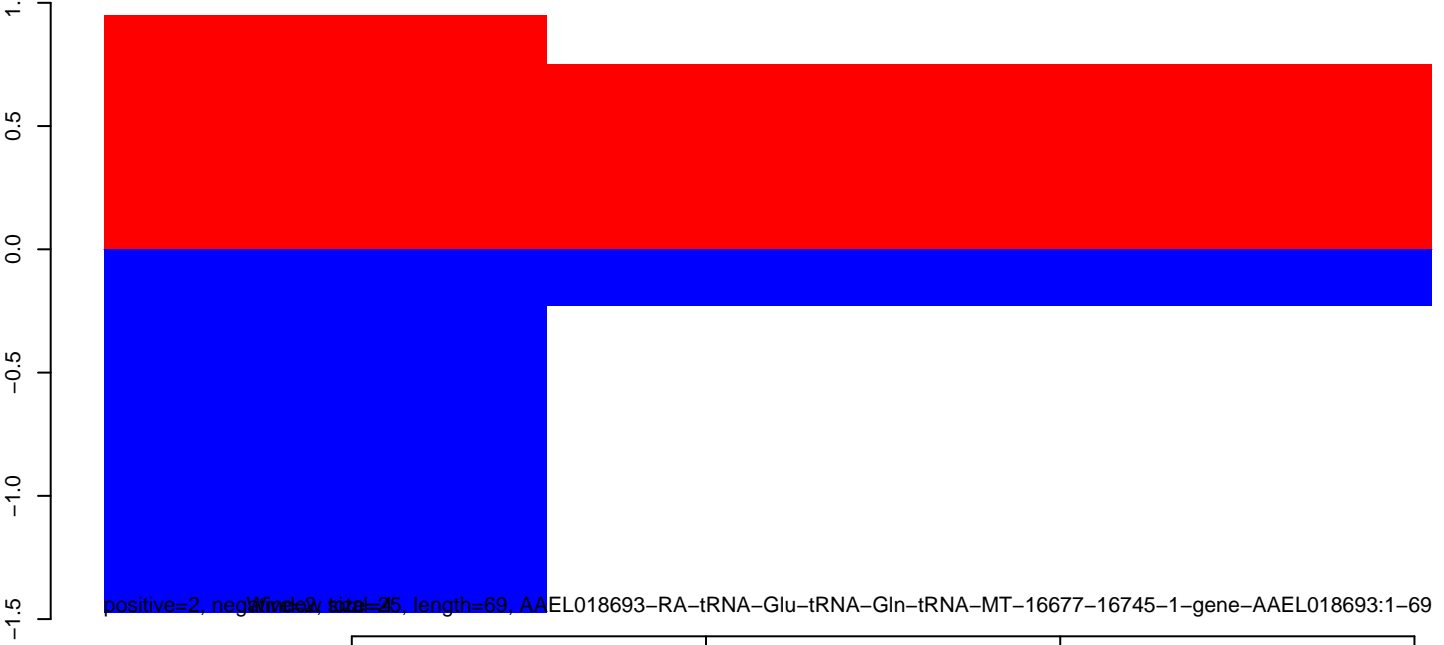
AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep



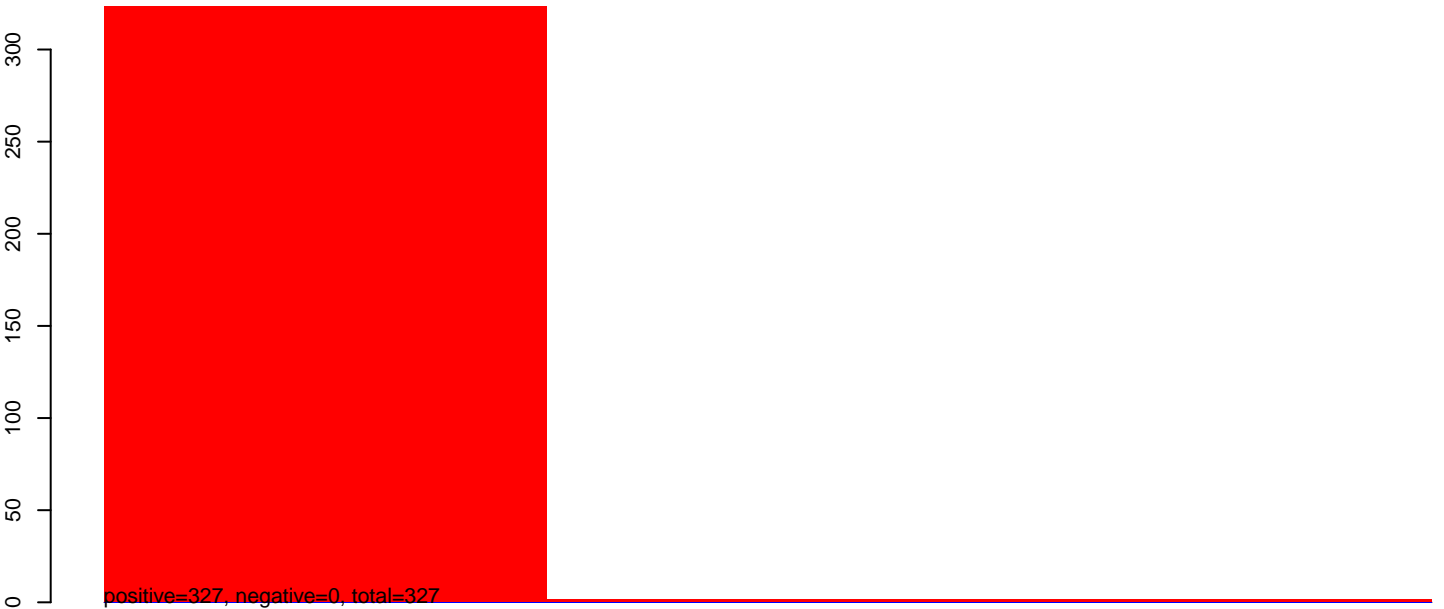
AeAeg_Aag2.TC.Dmem.rep



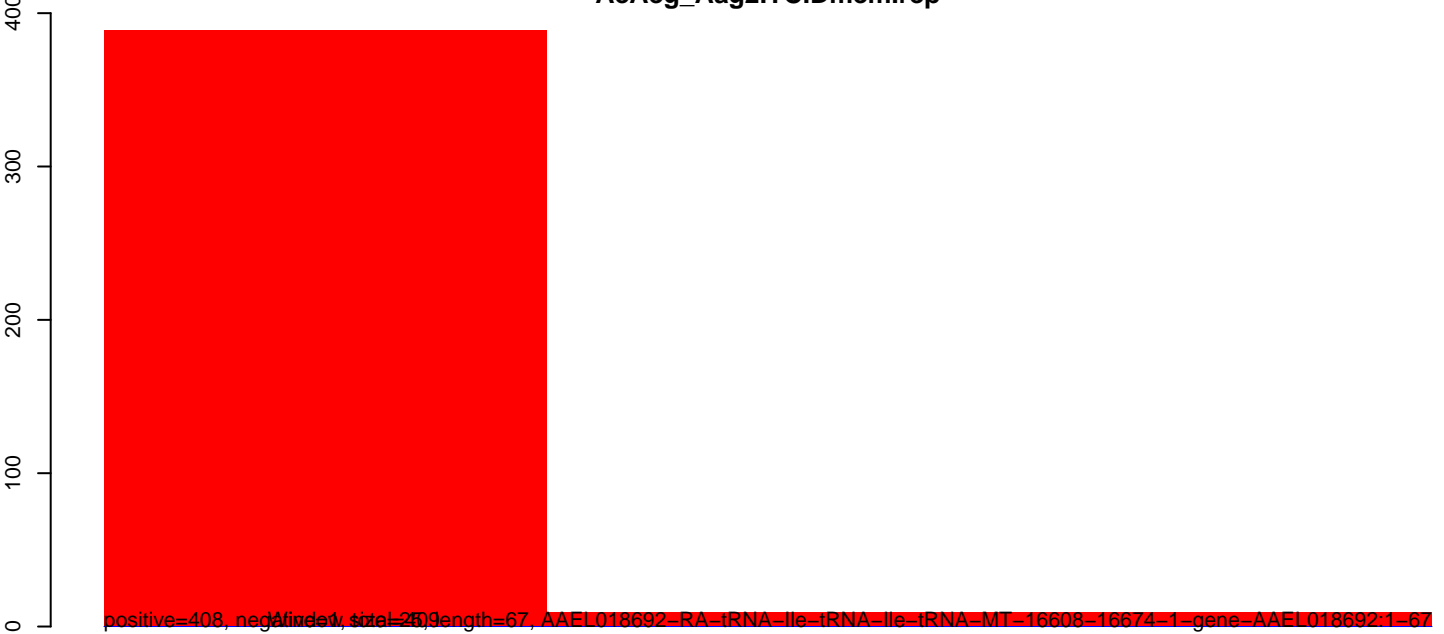
AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep



AeAeg_Aag2.TC.Dmem.rep



Window, total length=67, AAEL018692-RA-tRNA-Ile-tRNA-Ile-tRNA-MT-16608-16674-1-gene-AAEL018692:1-67

40 60 80 100

AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep



AeAeg_Aag2.TC.Dmem.rep



Window Size: 25, length=72, AAEL018690-RA-tRNA-Val-tRNA-Val-tRNA-MT-13901-13972-1-gene-AAEL018690:1-72

40 60 80 100

AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep

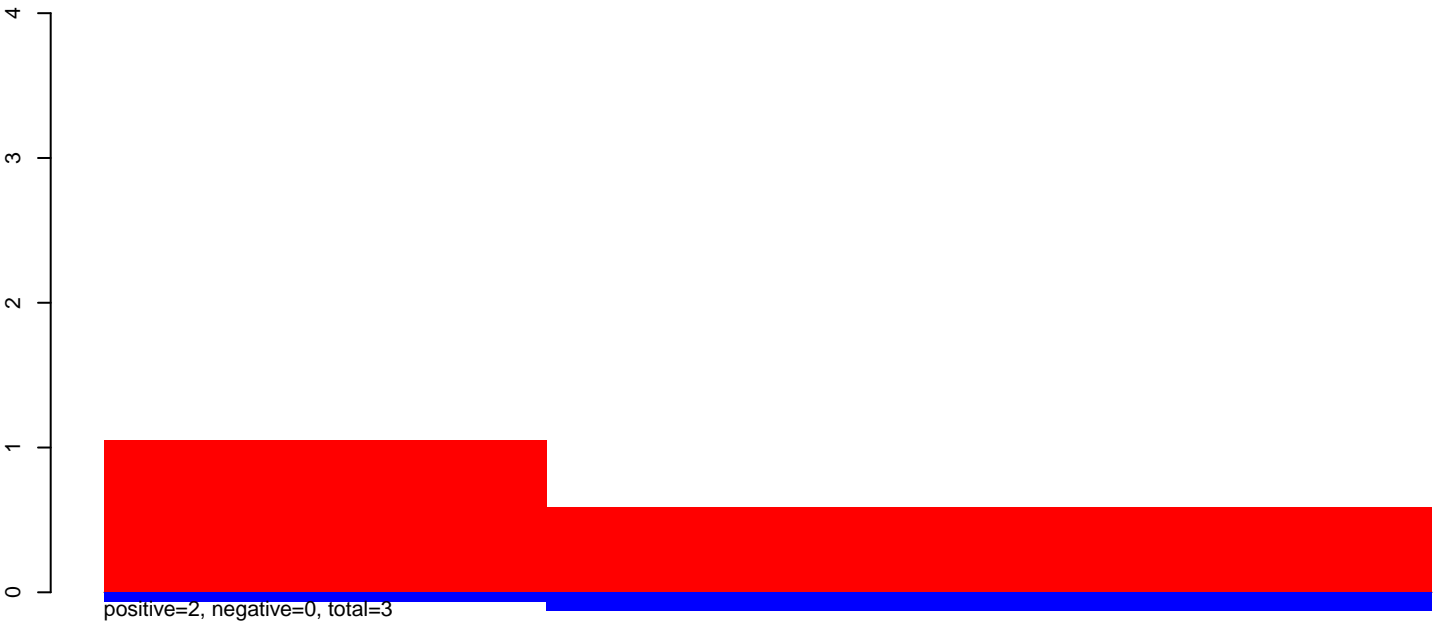


AeAeg_Aag2.TC.Dmem.rep

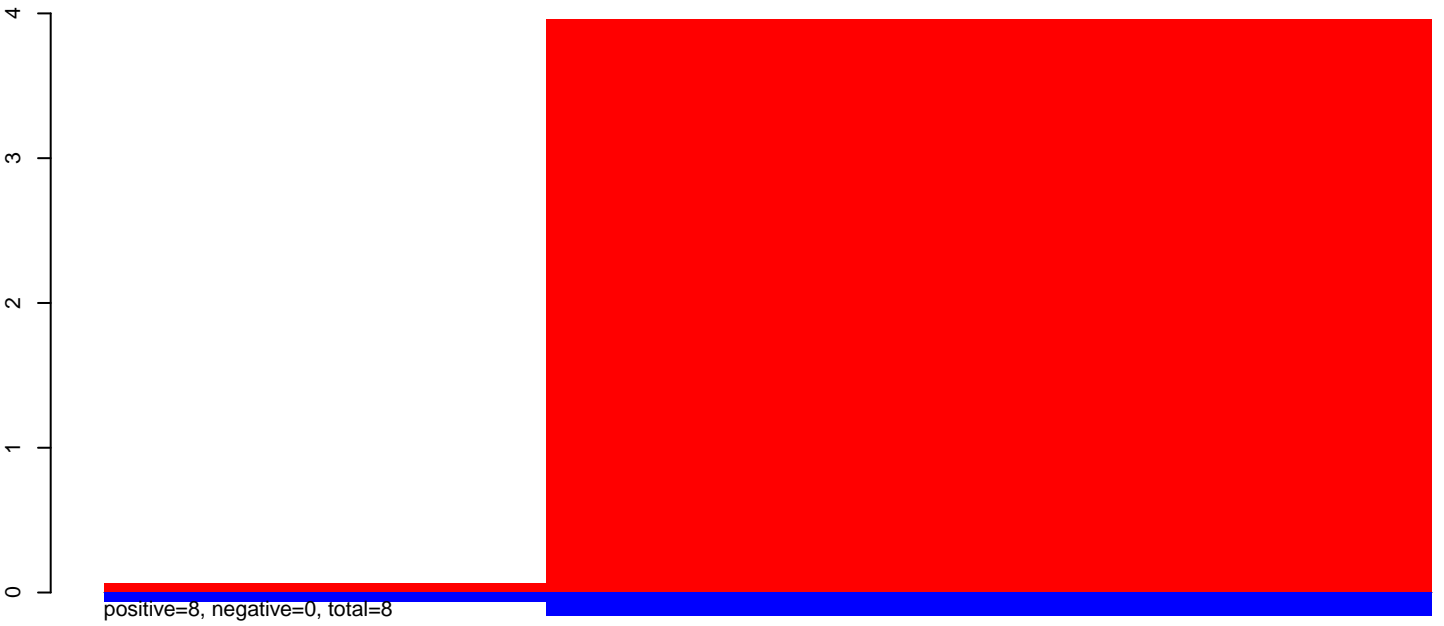


40 60 80 100

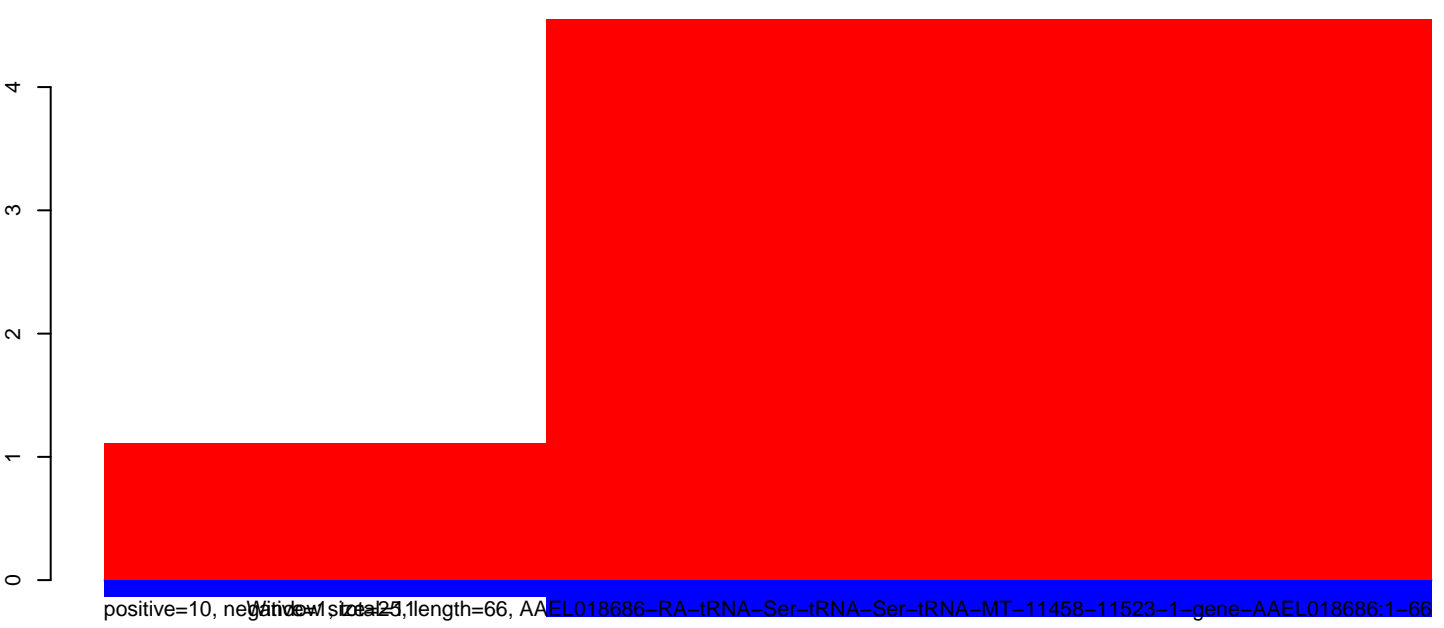
AeAeg_Aag2.TC.Dmem.18_23.rep

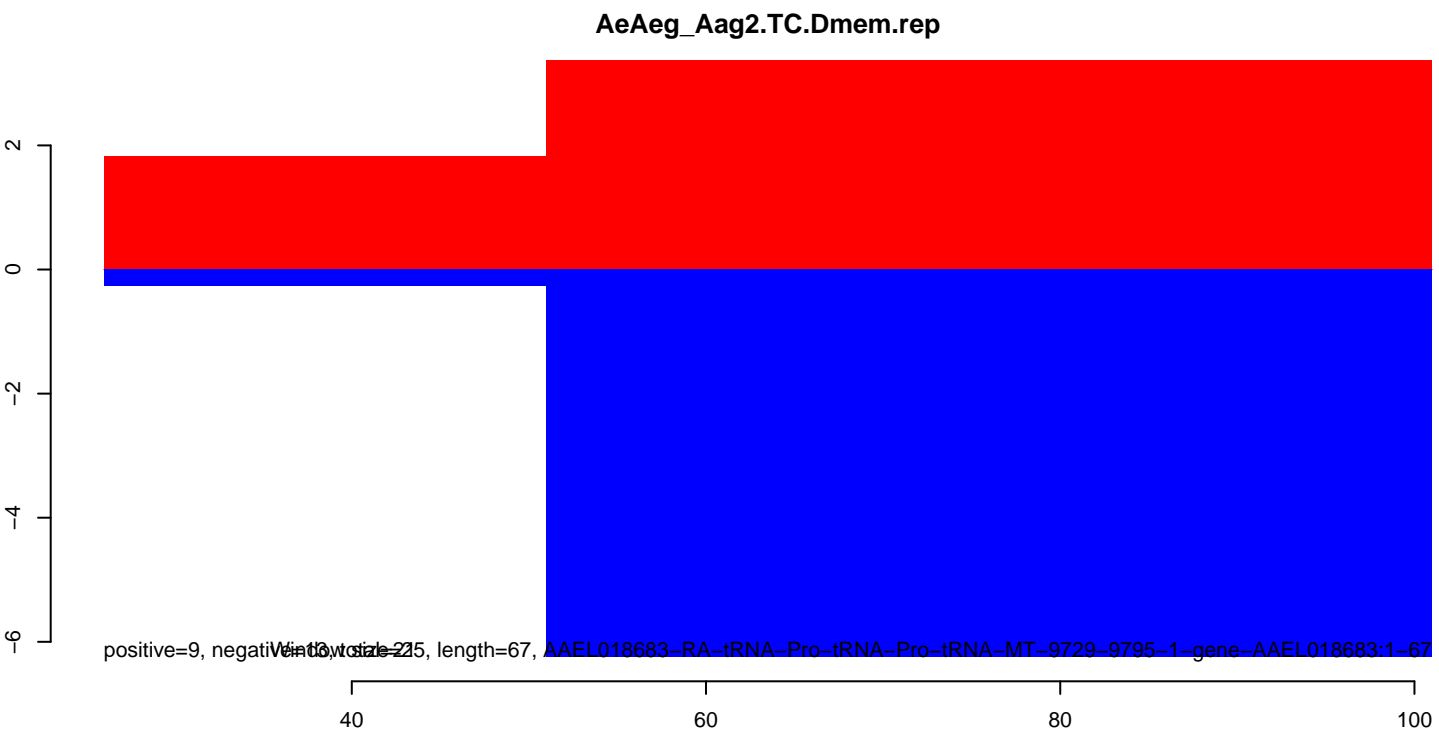
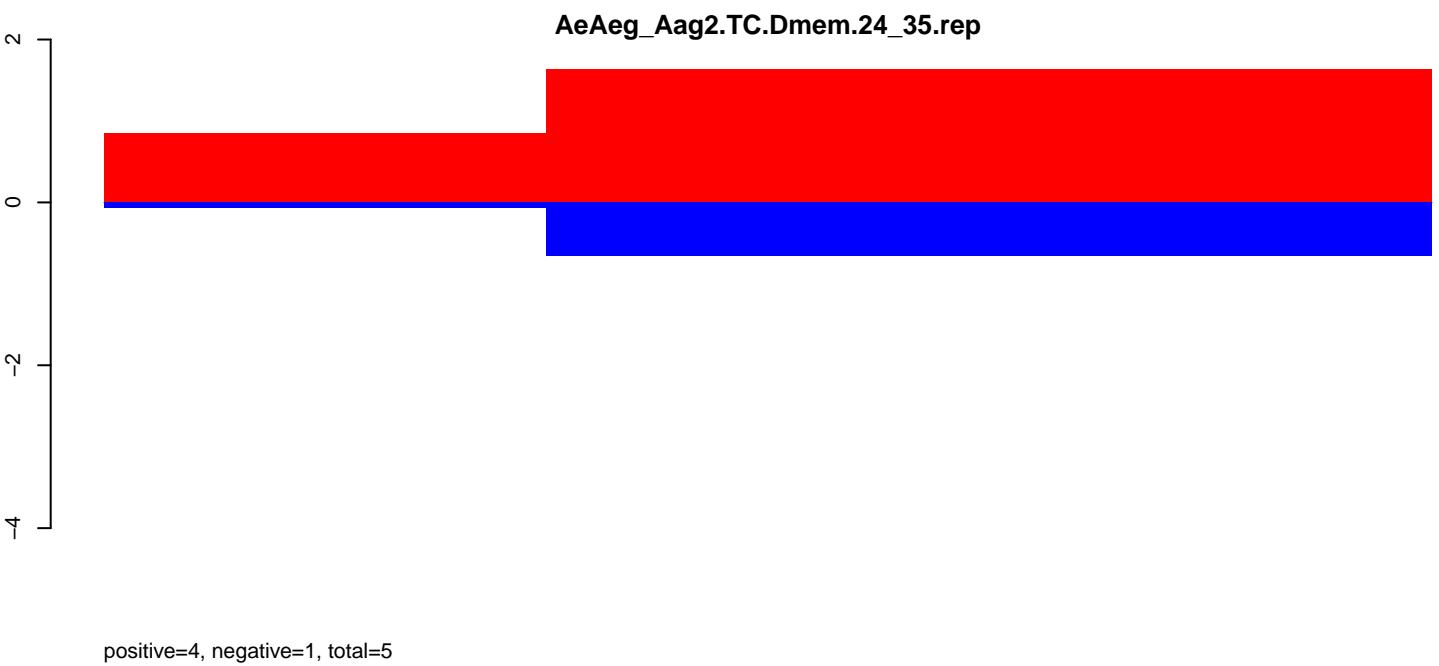
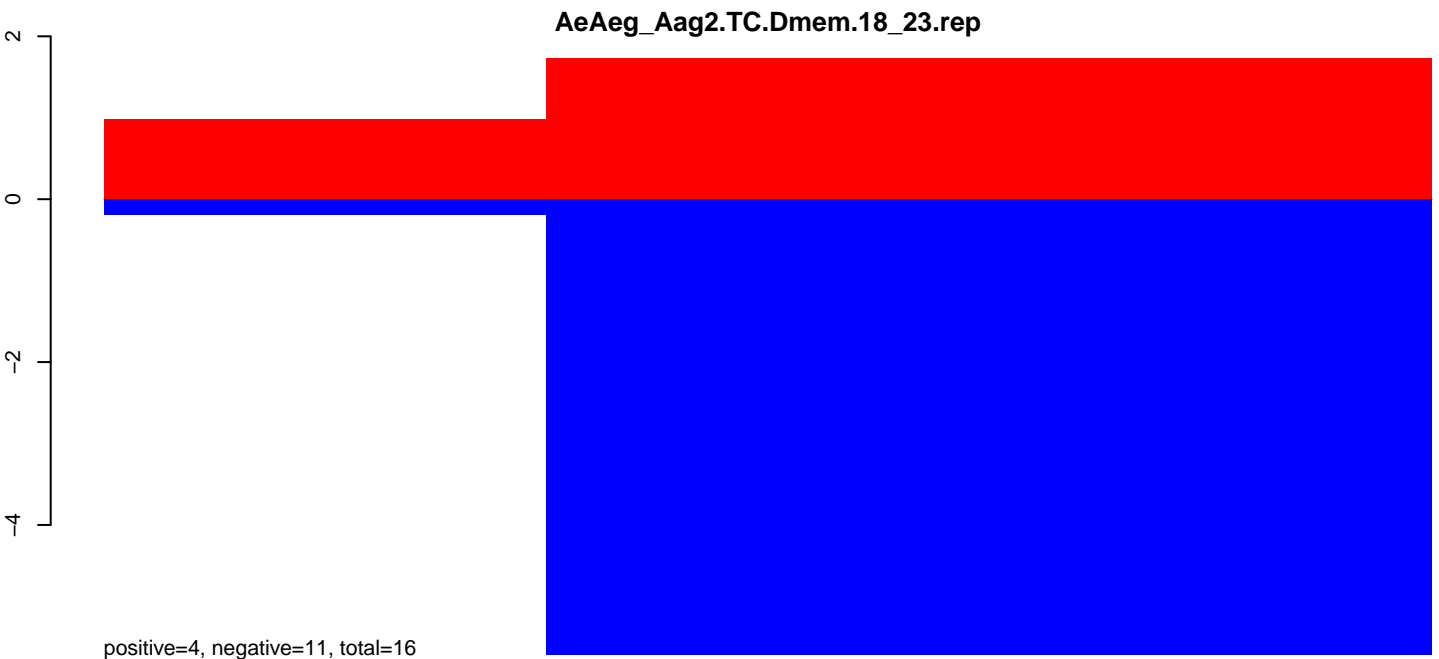


AeAeg_Aag2.TC.Dmem.24_35.rep

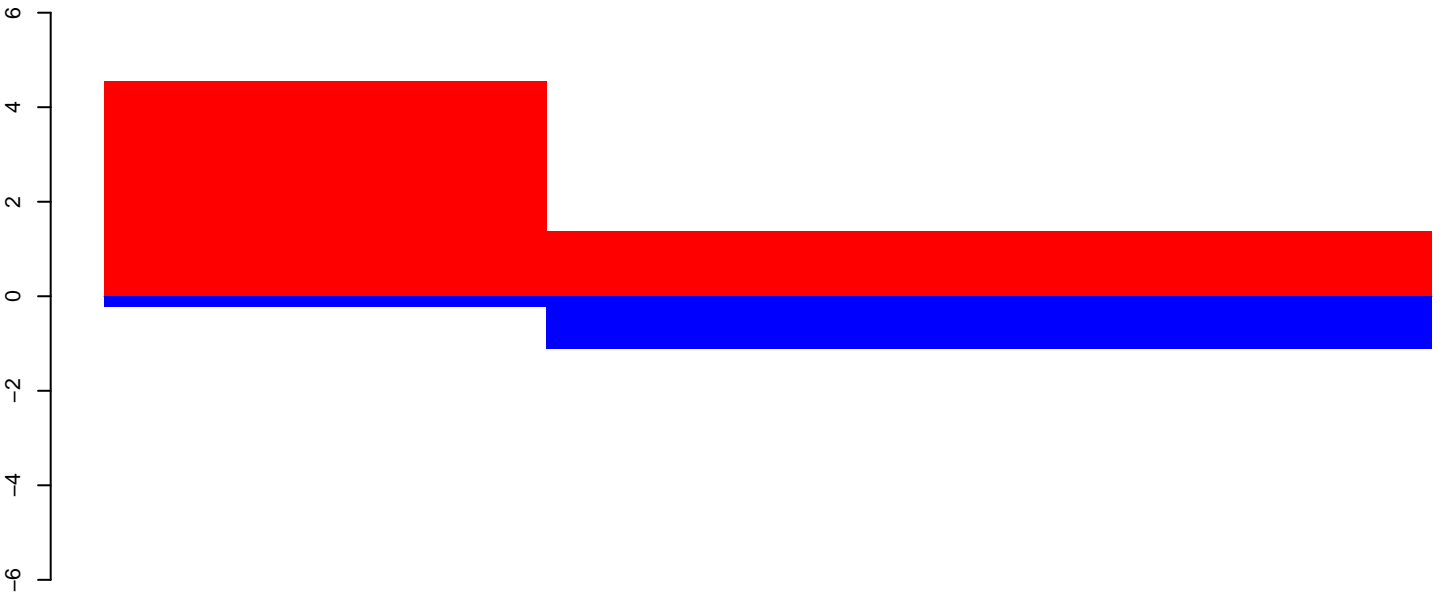


AeAeg_Aag2.TC.Dmem.rep



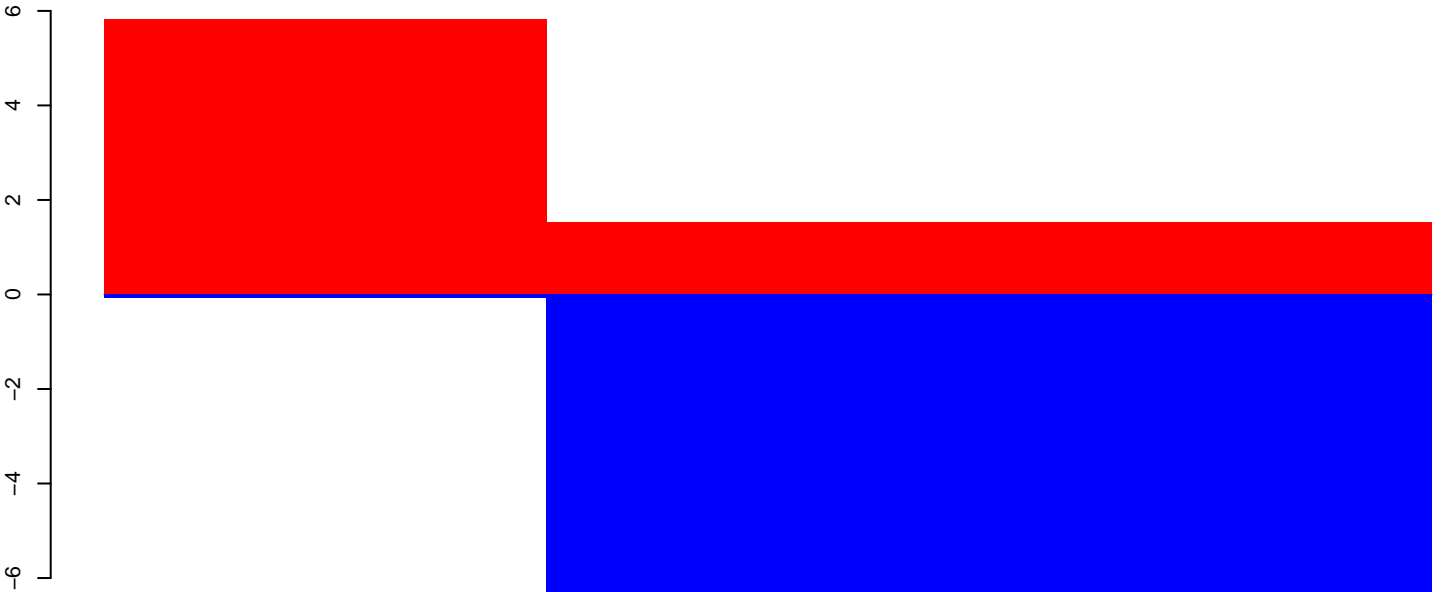


AeAeg_Aag2.TC.Dmem.18_23.rep



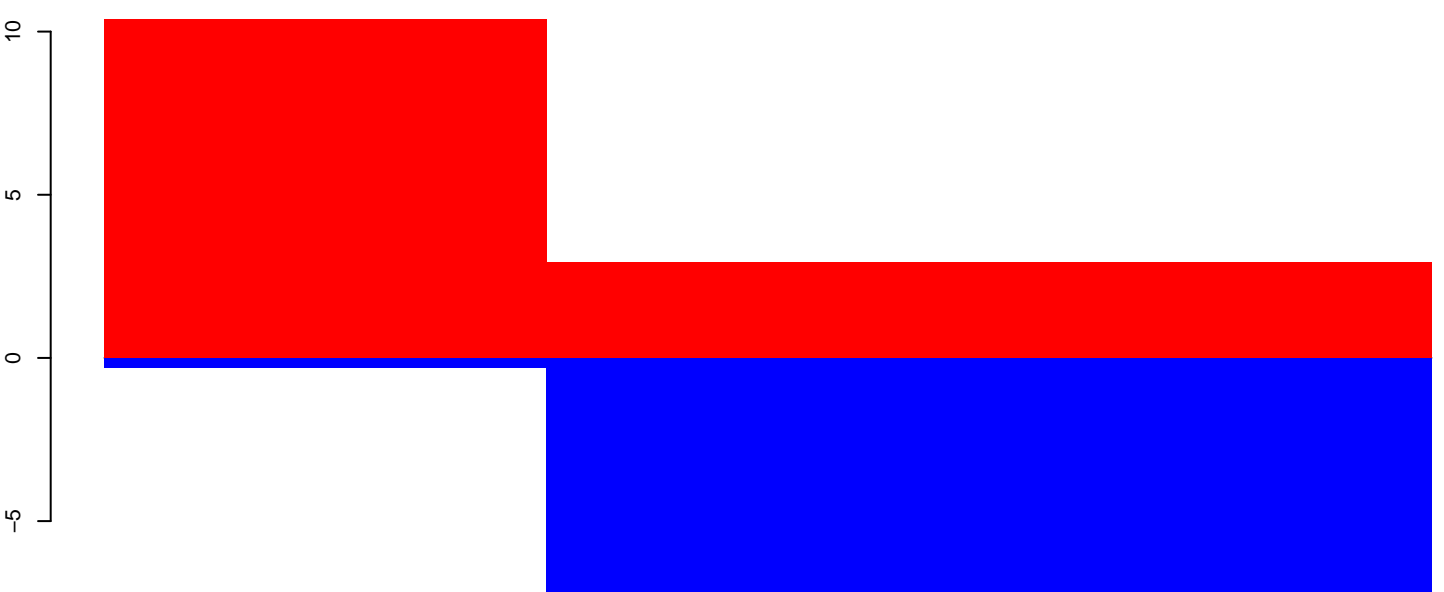
positive=7, negative=2, total=10

AeAeg_Aag2.TC.Dmem.24_35.rep



positive=9, negative=14, total=23

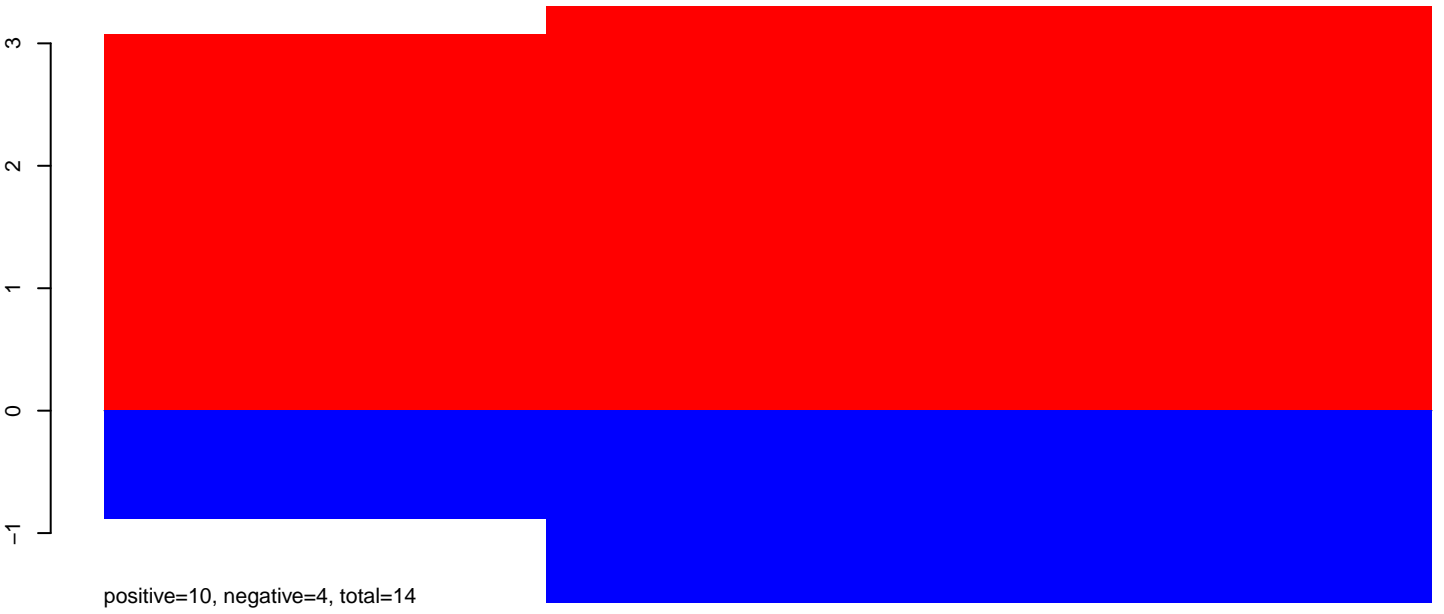
AeAeg_Aag2.TC.Dmem.rep



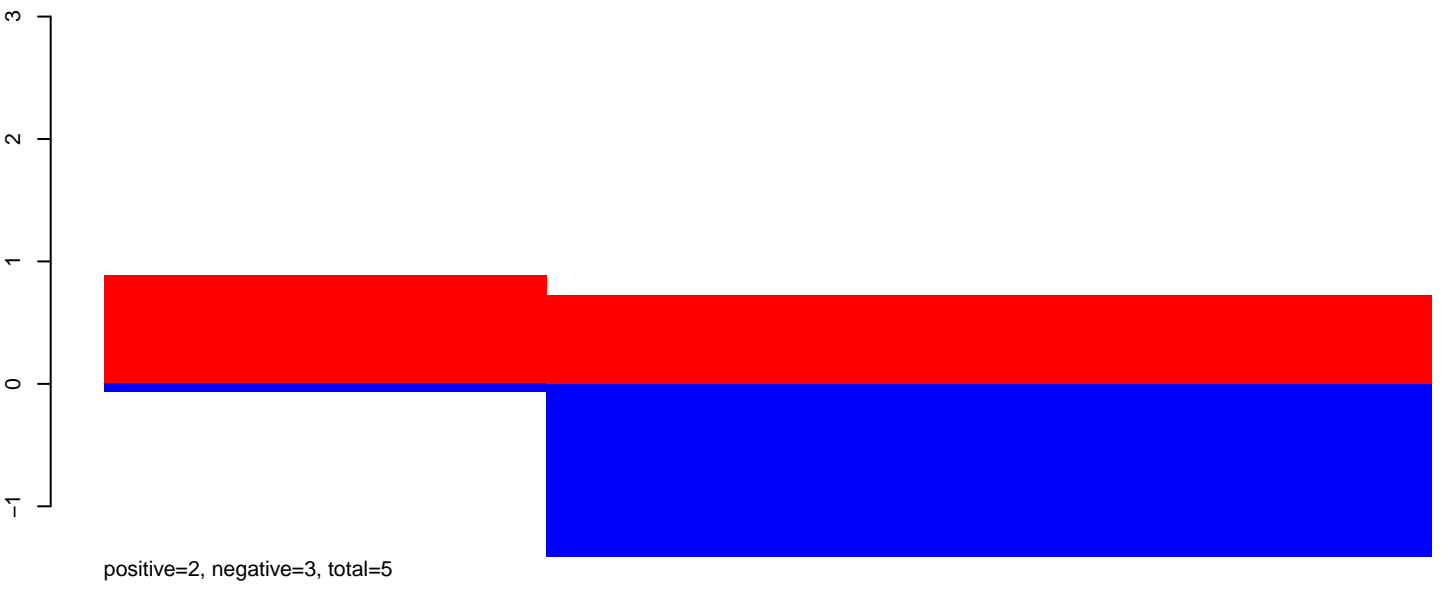
positive=16, negative=16, total=32, length=67, AAEL018677-RA-tRNA-Phe-tRNA-Phe-tRNA-MT-6139-6205-1-gene-AAEL018677-1-67

40 60 80 100

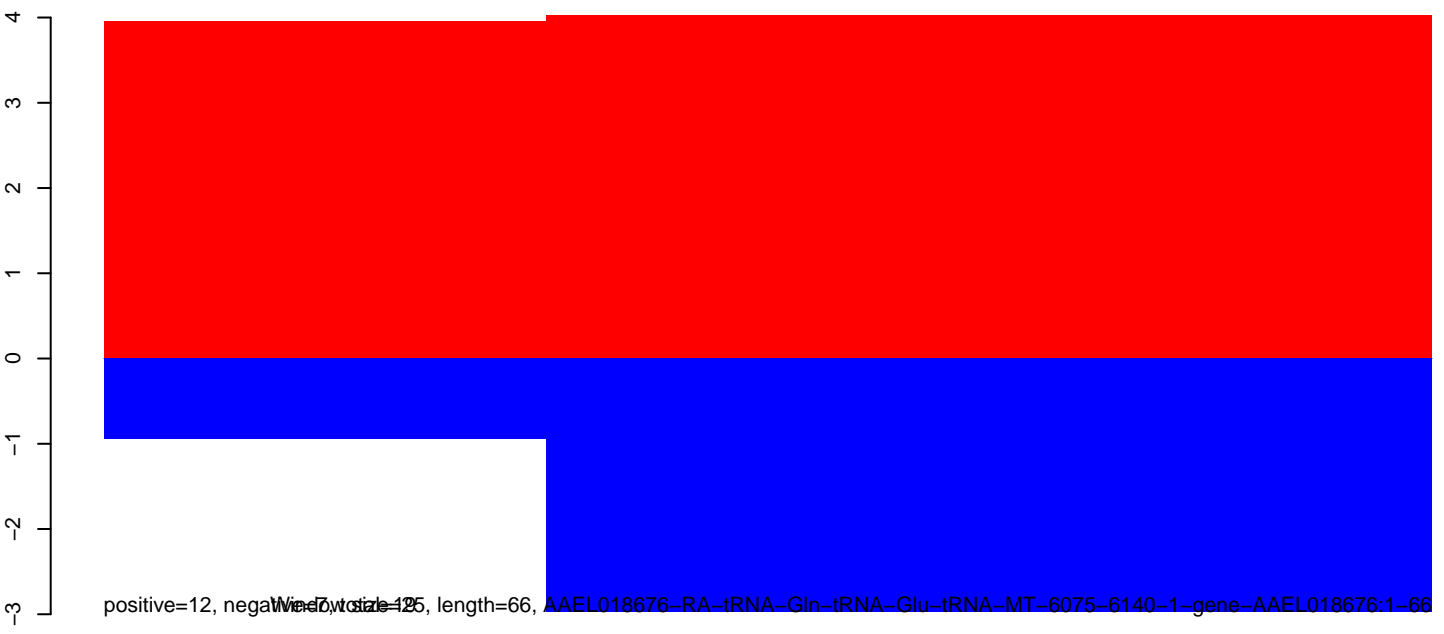
AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep

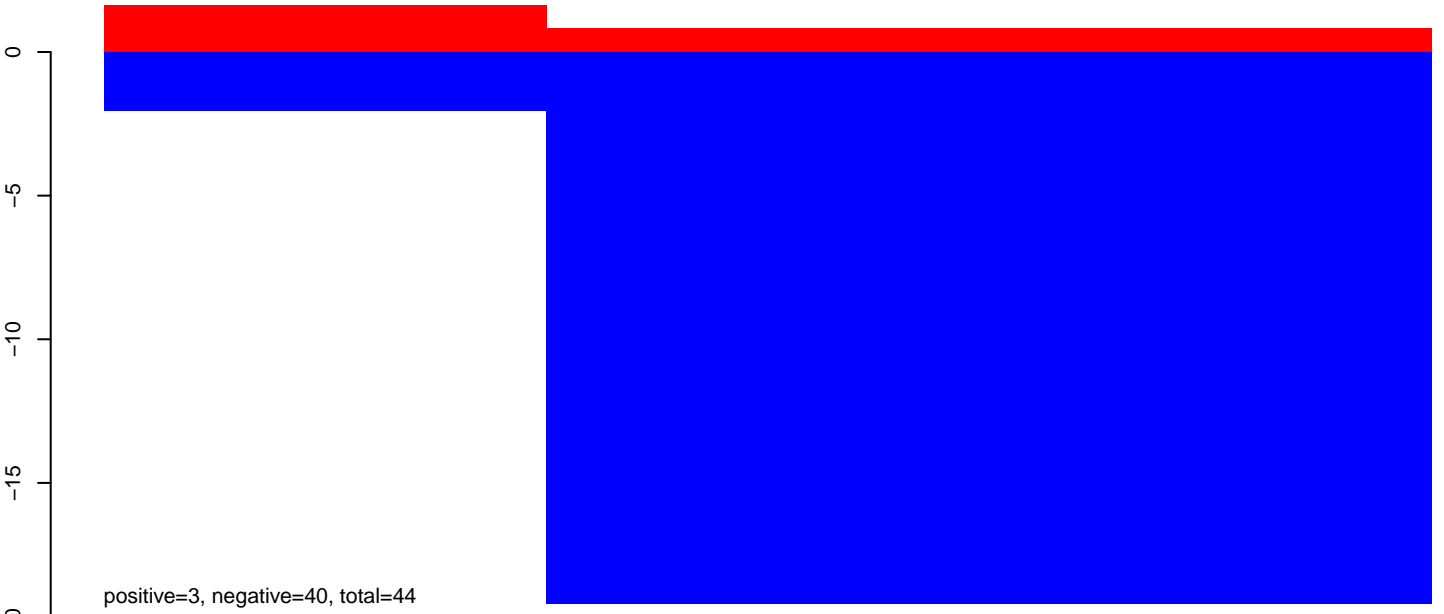


AeAeg_Aag2.TC.Dmem.rep



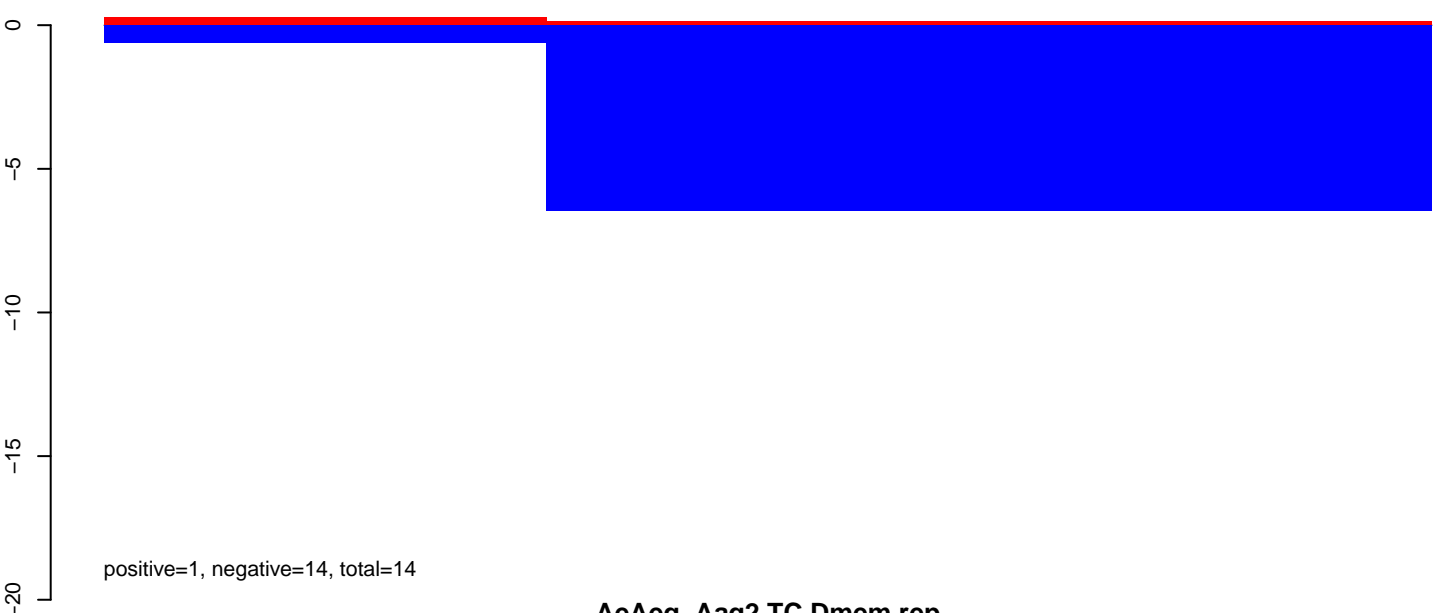
40 60 80 100

AeAeg_Aag2.TC.Dmem.18_23.rep



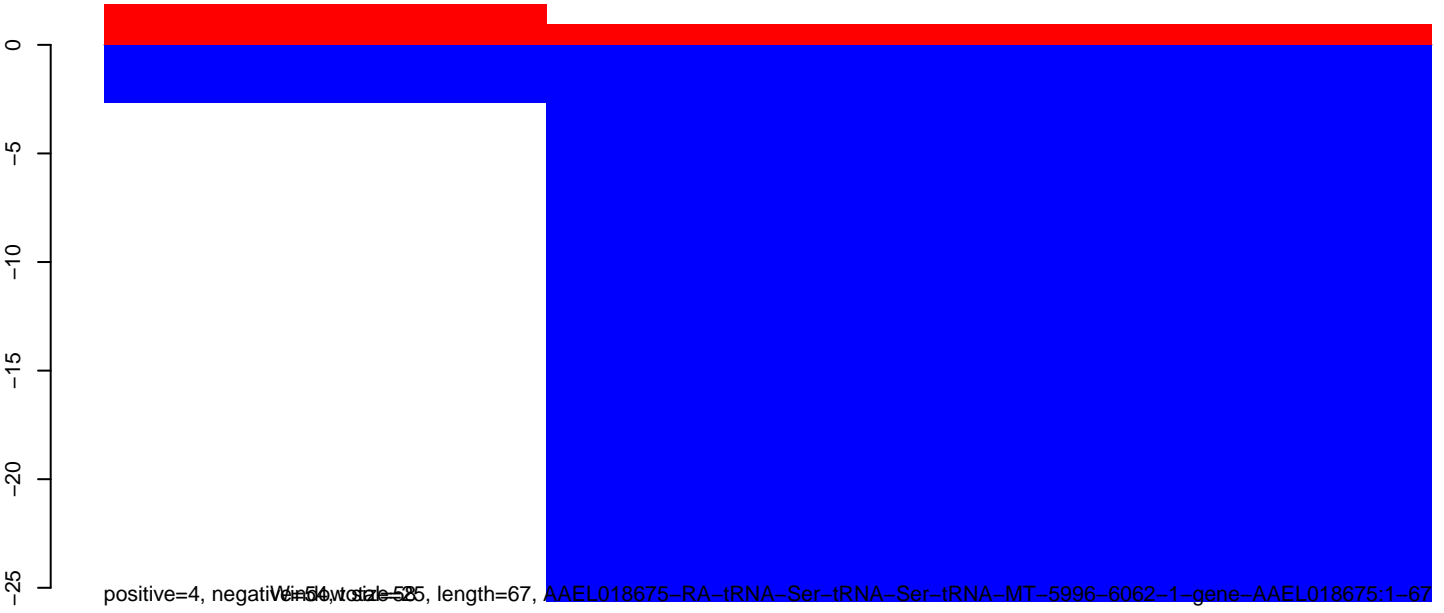
positive=3, negative=40, total=44

AeAeg_Aag2.TC.Dmem.24_35.rep



positive=1, negative=14, total=14

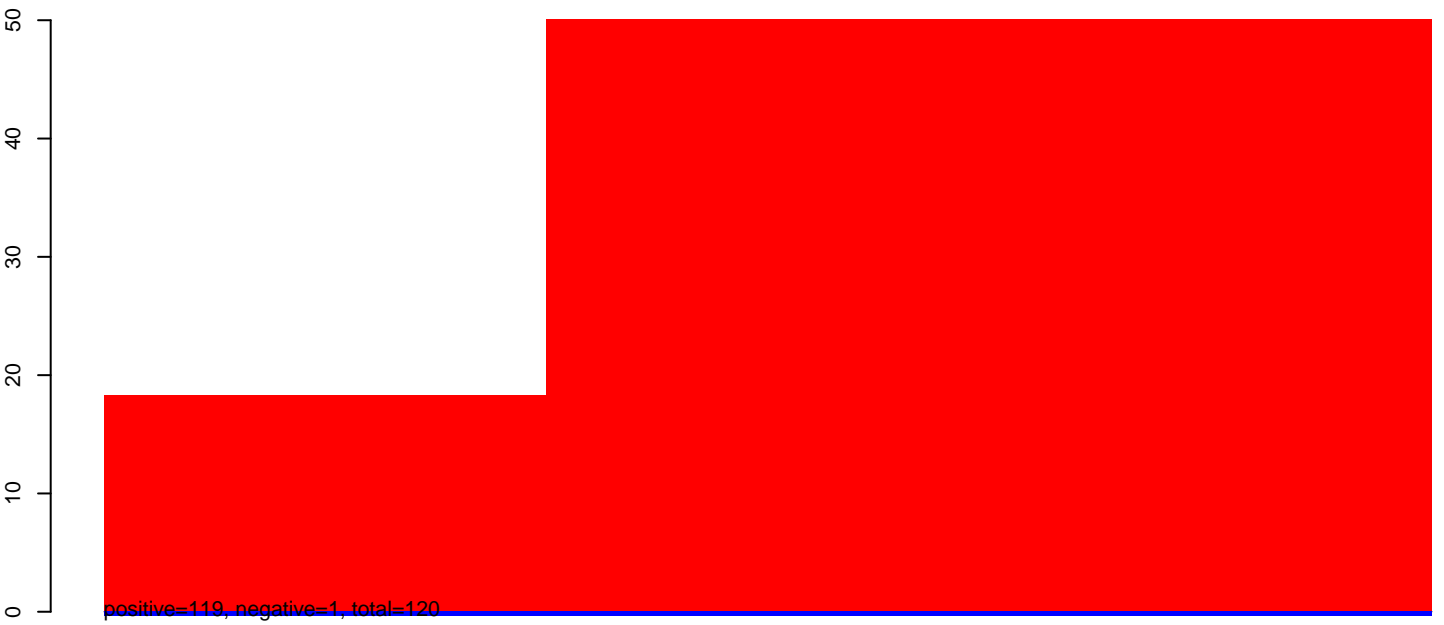
AeAeg_Aag2.TC.Dmem.rep



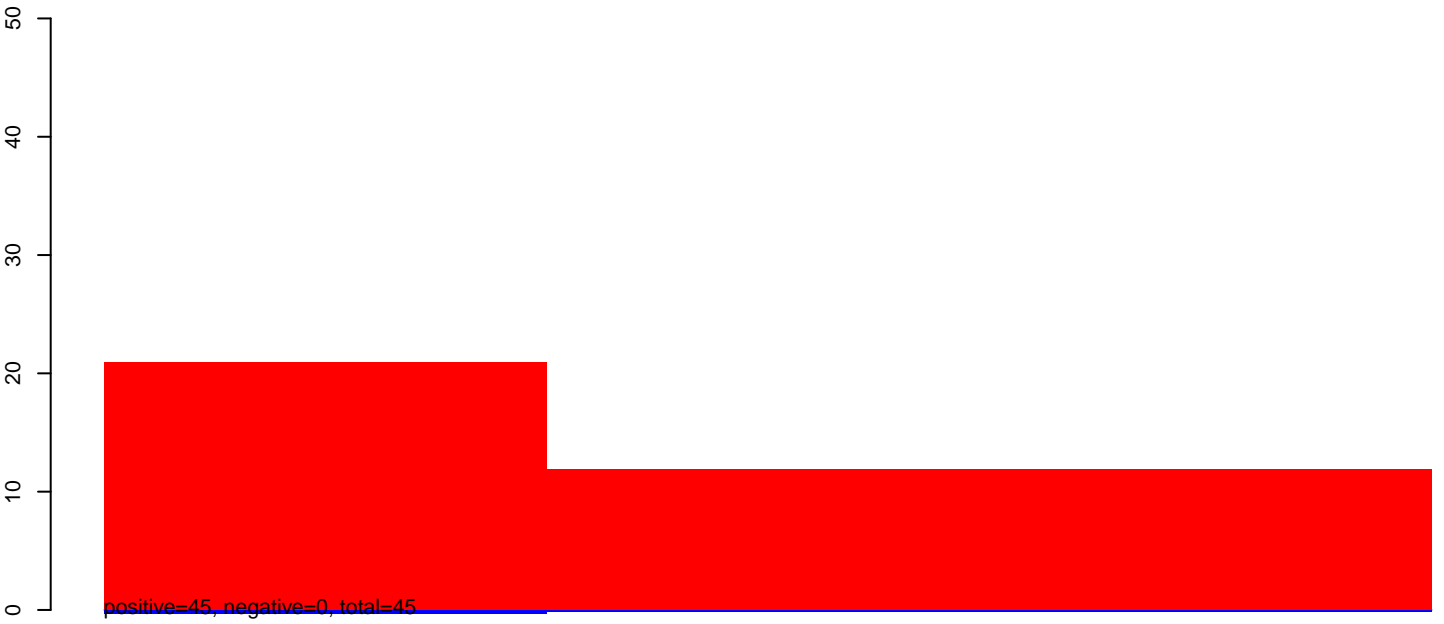
positive=4, negative=48, total=52, length=67, AAEL018675-RA-tRNA-Ser-tRNA-Ser-tRNA-MT-5996-6062-1-gene-AAEL018675:1-67

40 60 80 100

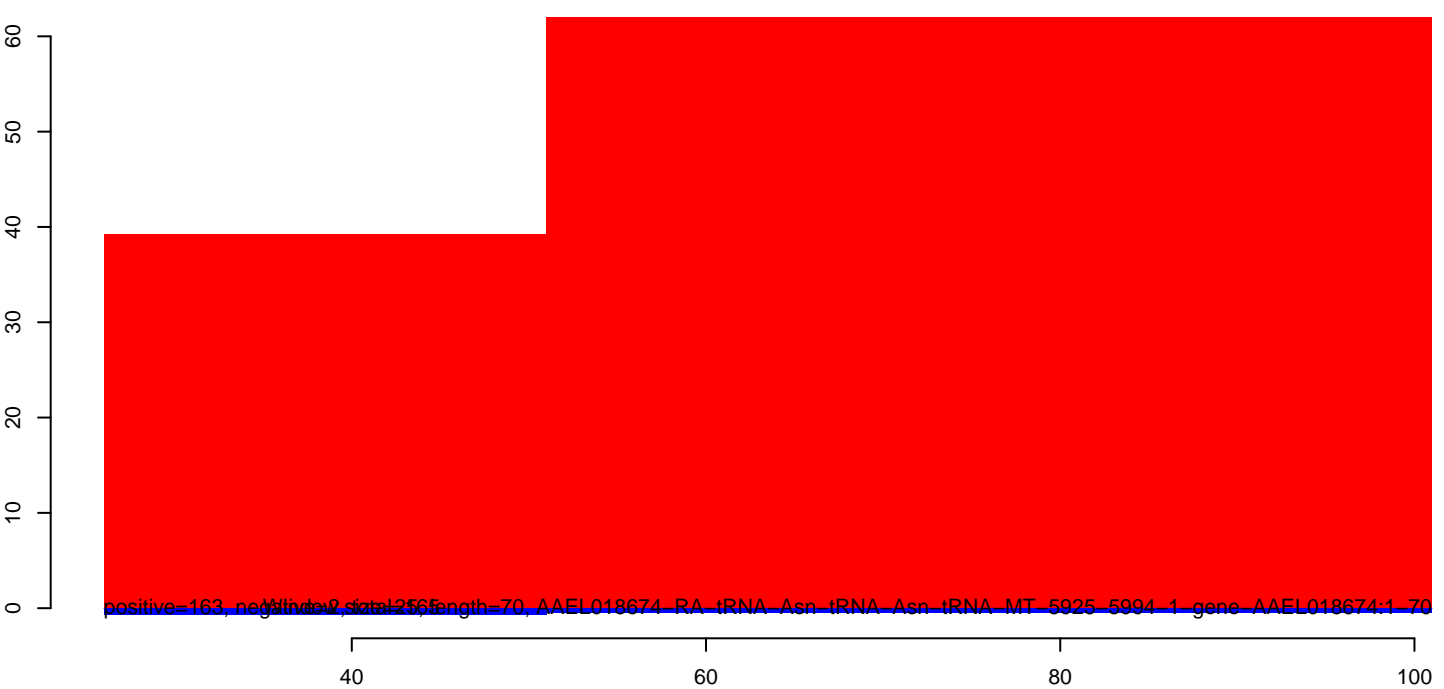
AeAeg_Aag2.TC.Dmem.18_23.rep



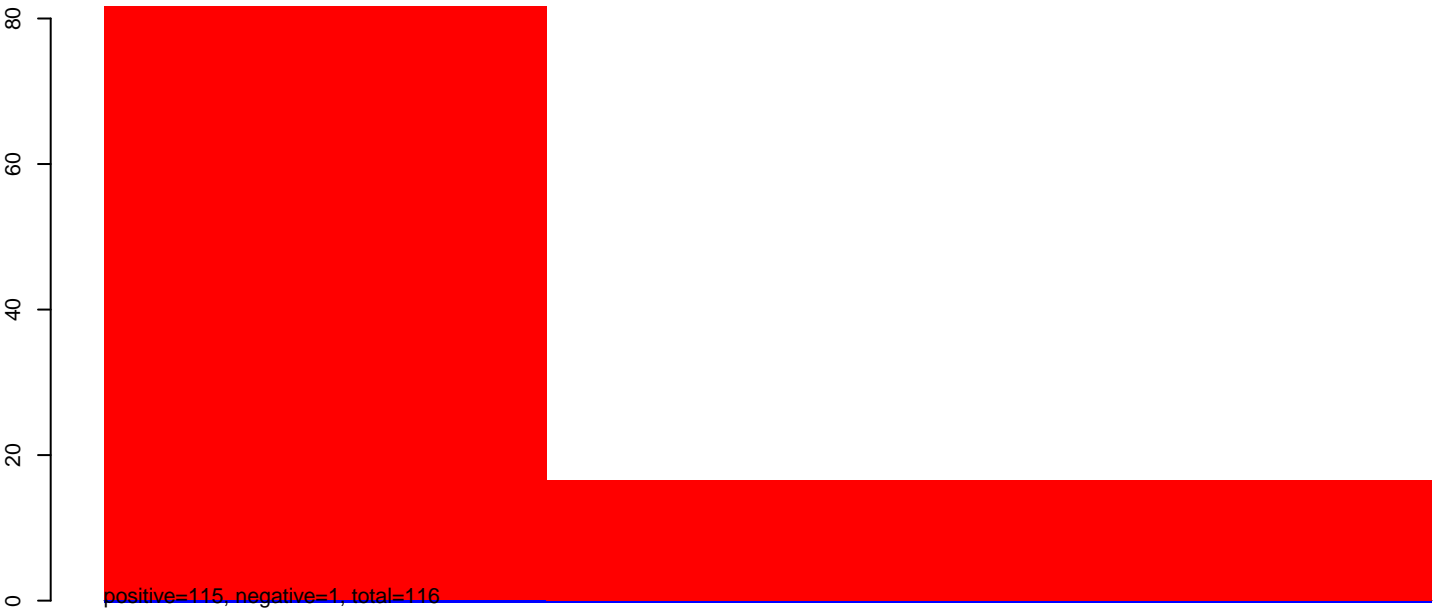
AeAeg_Aag2.TC.Dmem.24_35.rep



AeAeg_Aag2.TC.Dmem.rep



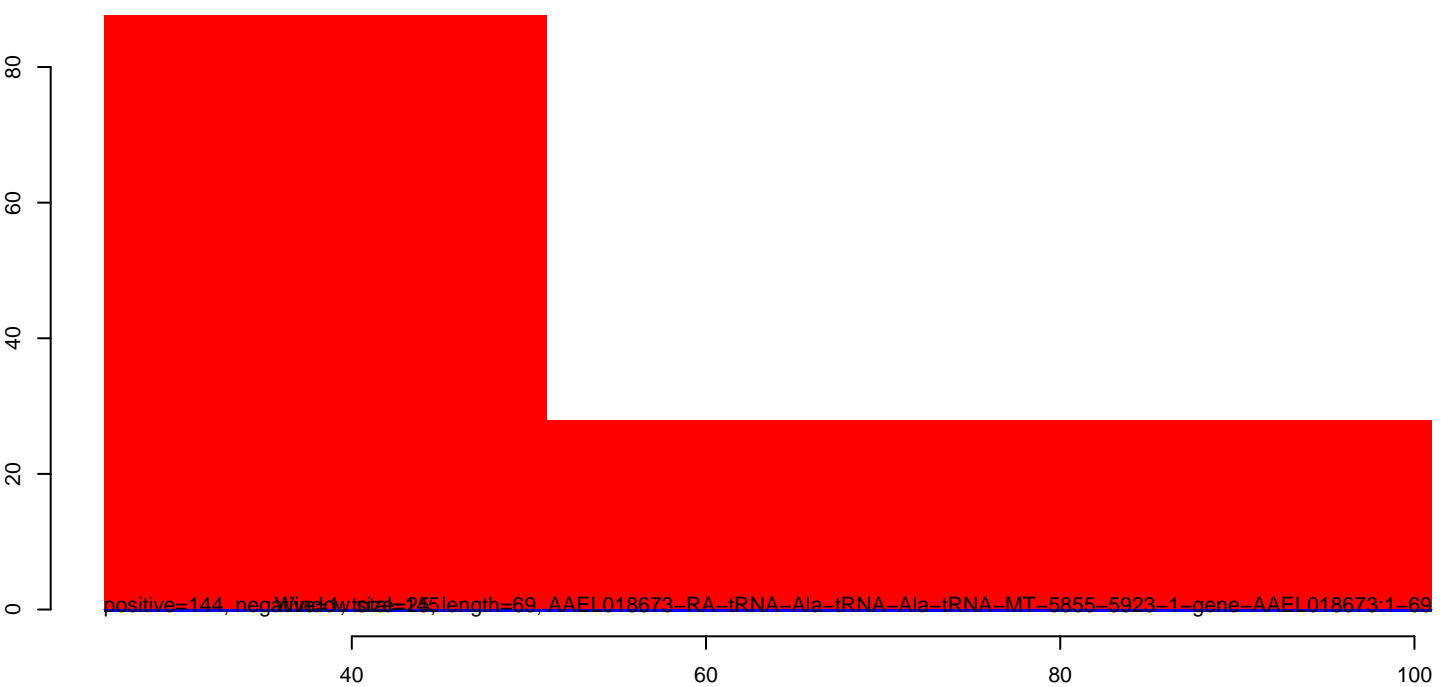
AeAeg_Aag2.TC.Dmem.18_23.rep



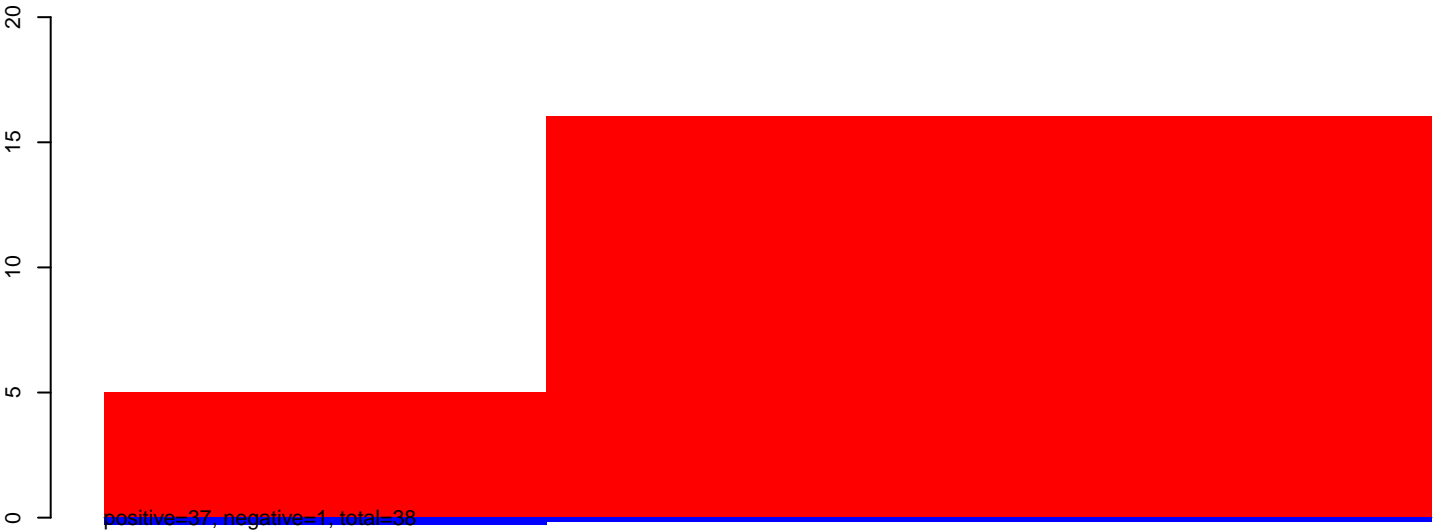
AeAeg_Aag2.TC.Dmem.24_35.rep



AeAeg_Aag2.TC.Dmem.rep



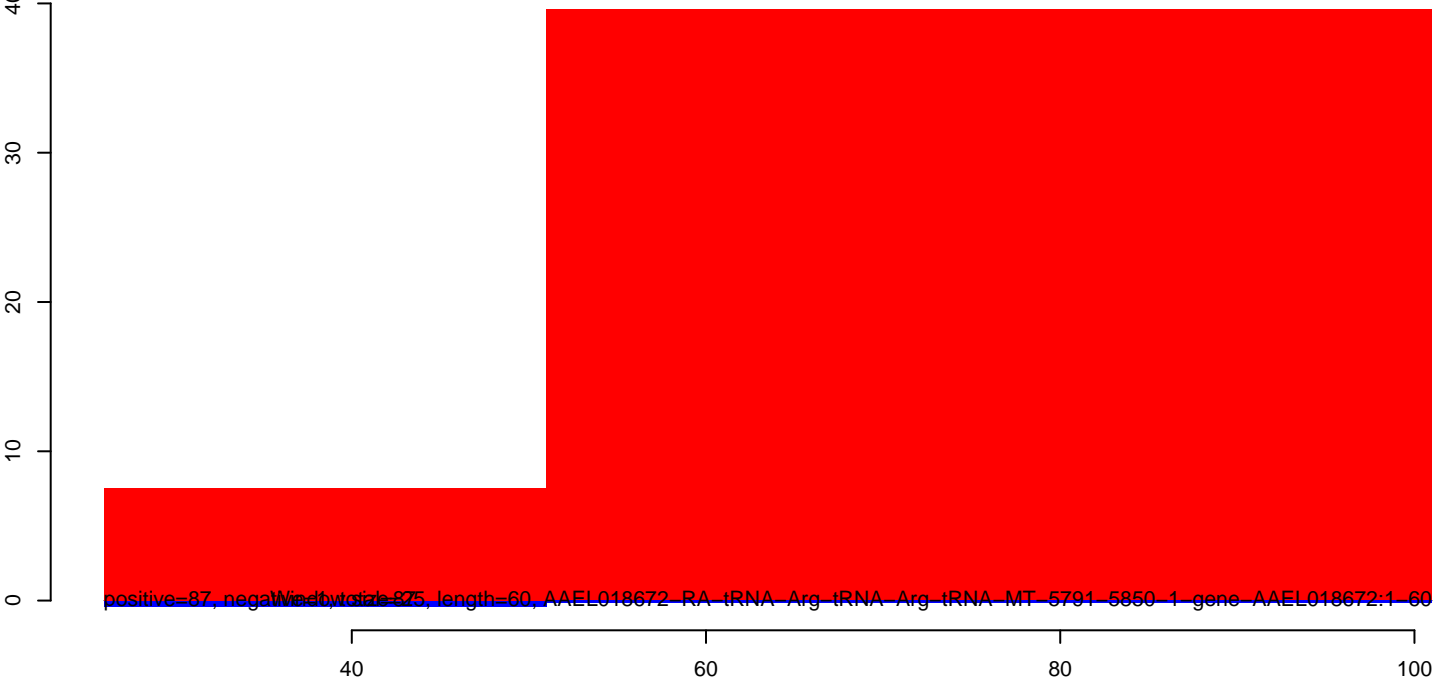
AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep



AeAeg_Aag2.TC.Dmem.rep

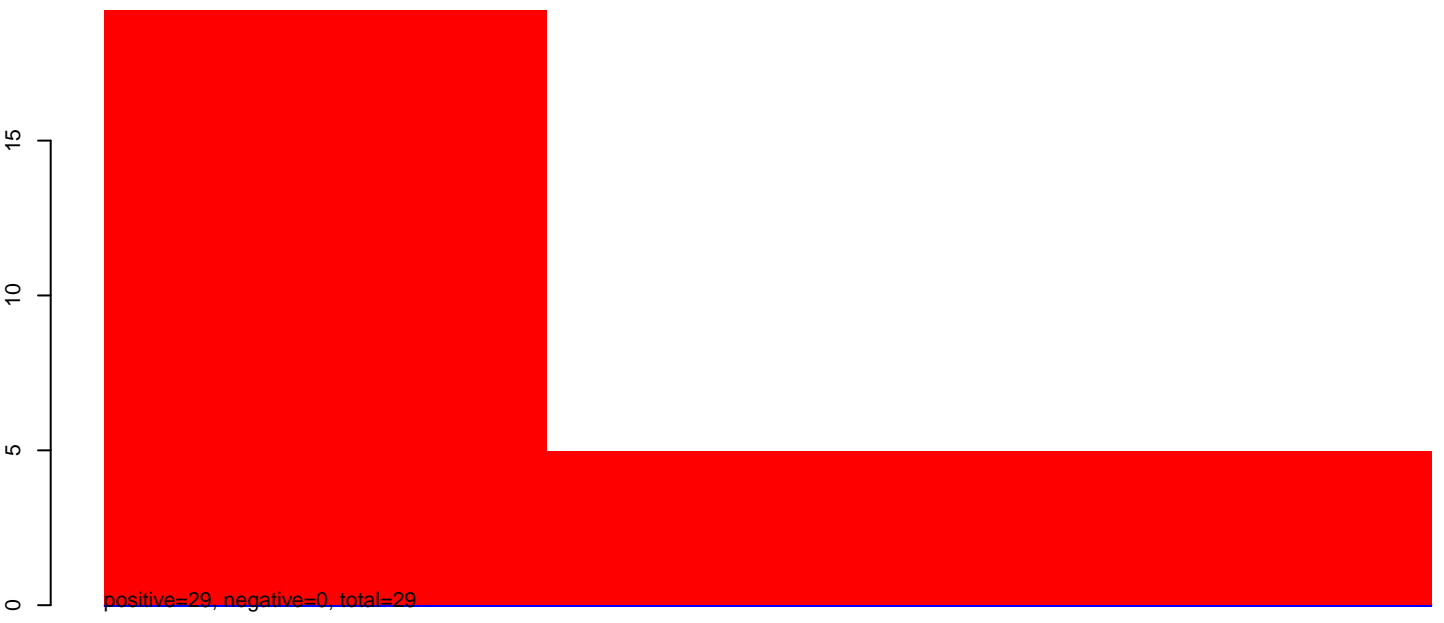


width=875, length=60, AAEL018672-RA tRNA-Arg tRNA-Arg tRNA-MT-5791-5850-1 gene-AAEL018672:1-60

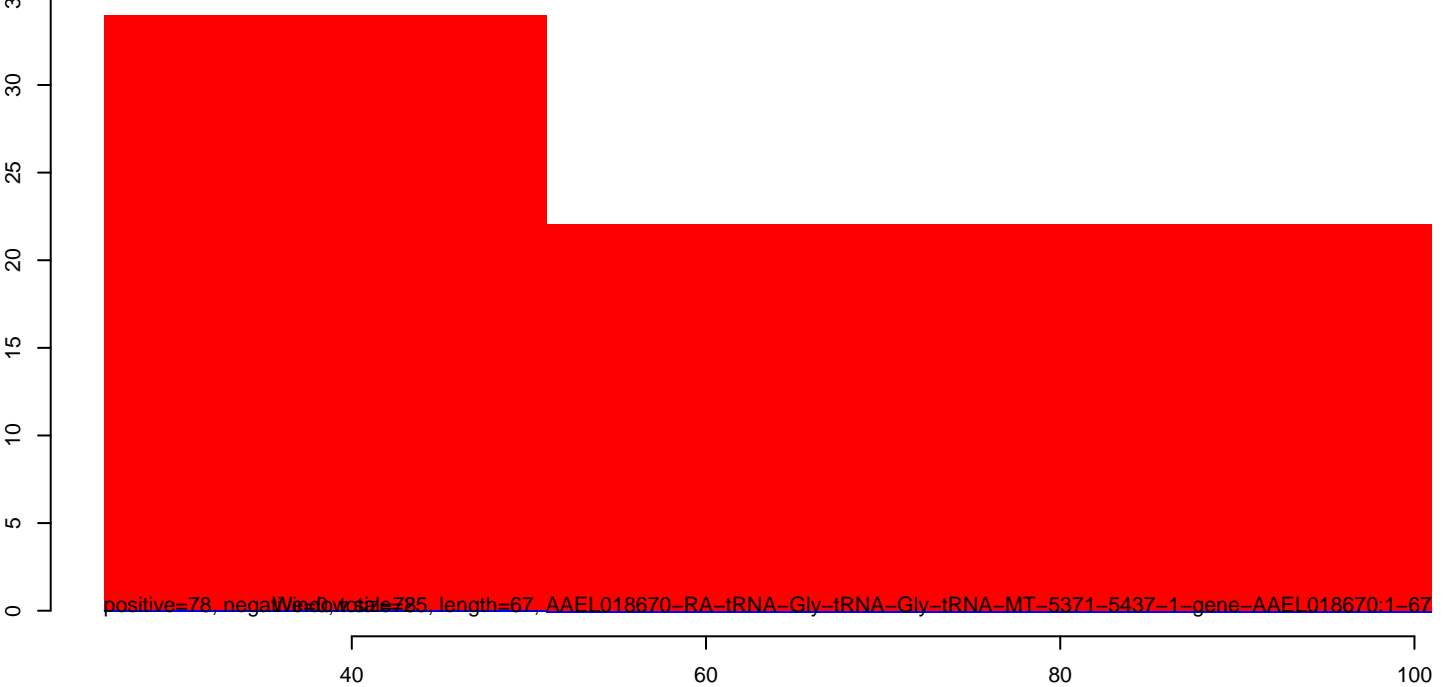
AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep



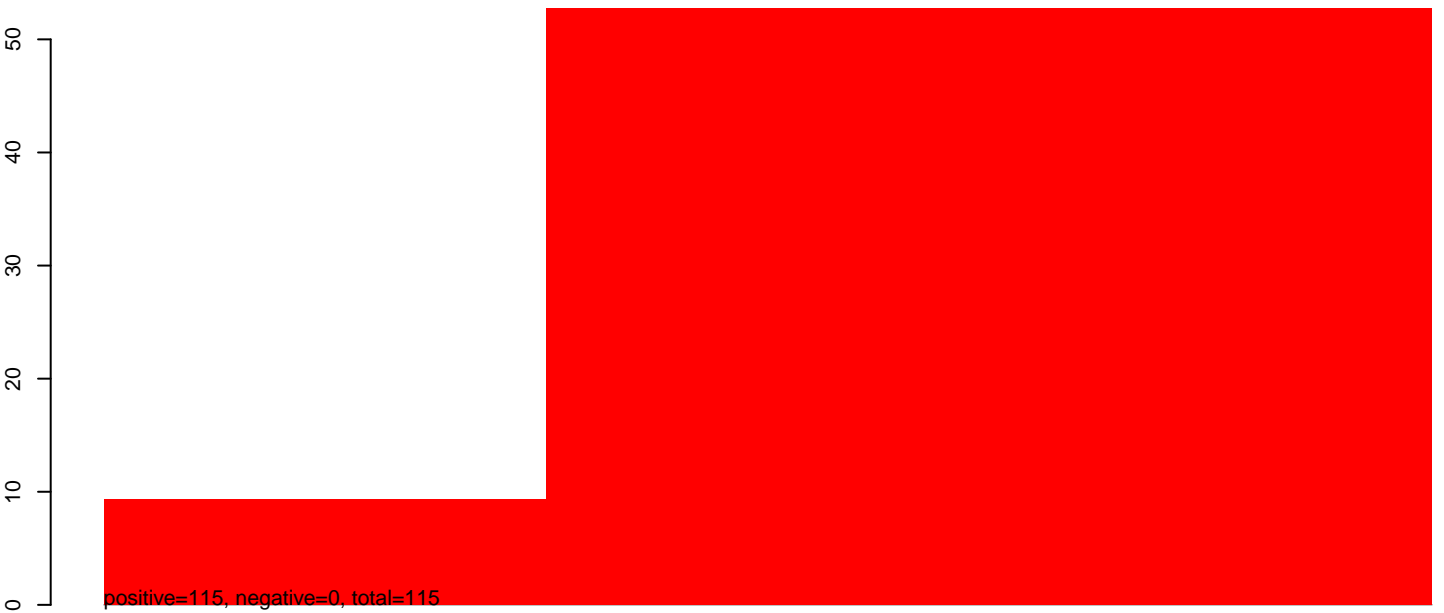
AeAeg_Aag2.TC.Dmem.rep



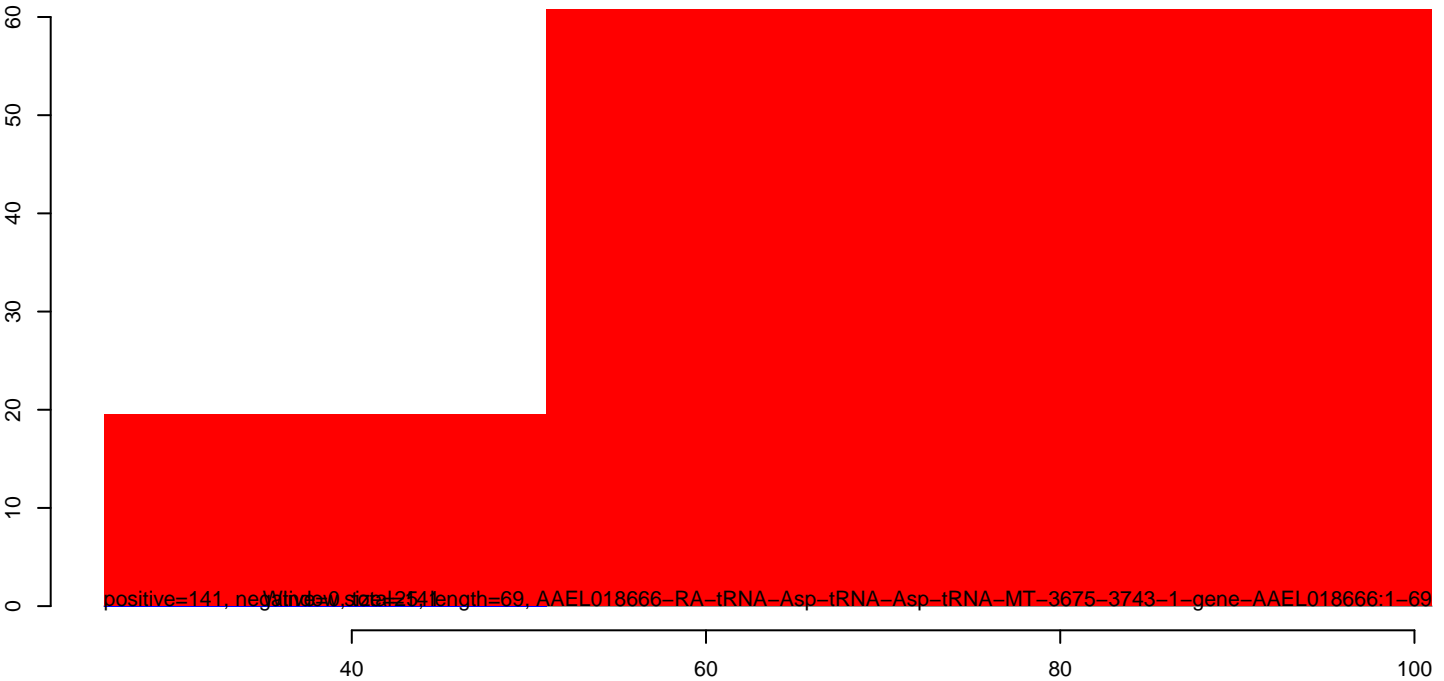
AeAeg_Aag2.TC.Dmem.18_23.rep



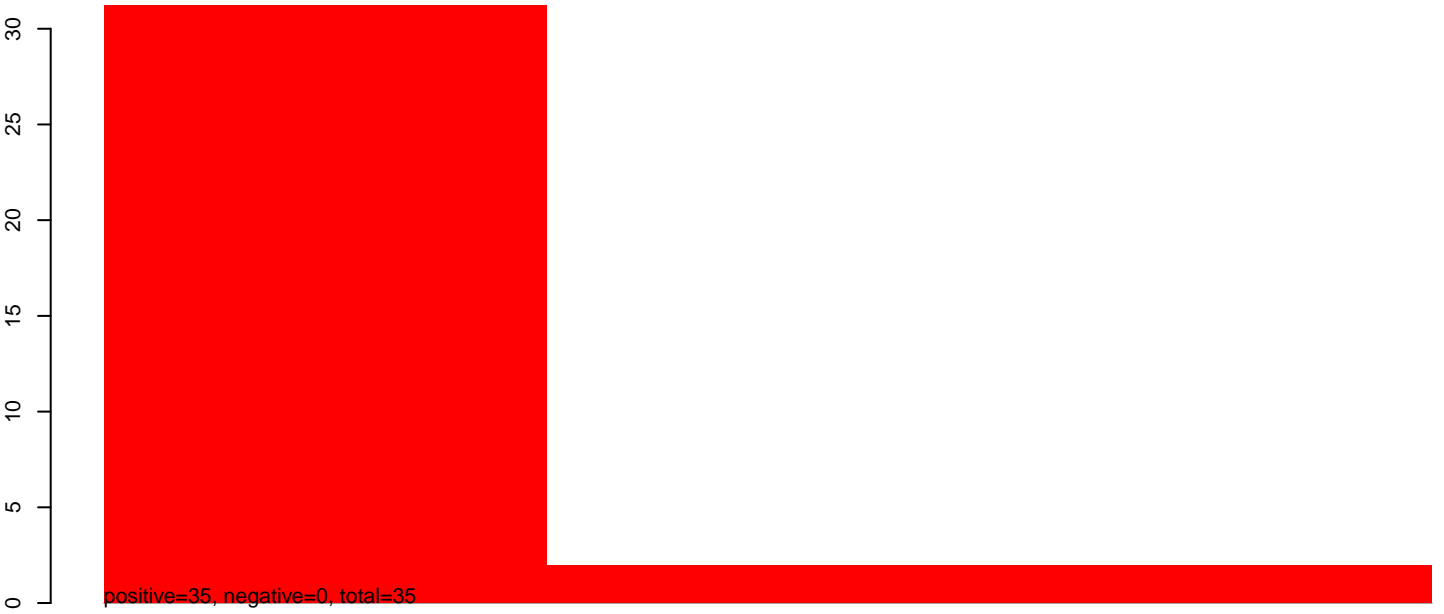
AeAeg_Aag2.TC.Dmem.24_35.rep



AeAeg_Aag2.TC.Dmem.rep



AeAeg_Aag2.TC.Dmem.18_23.rep



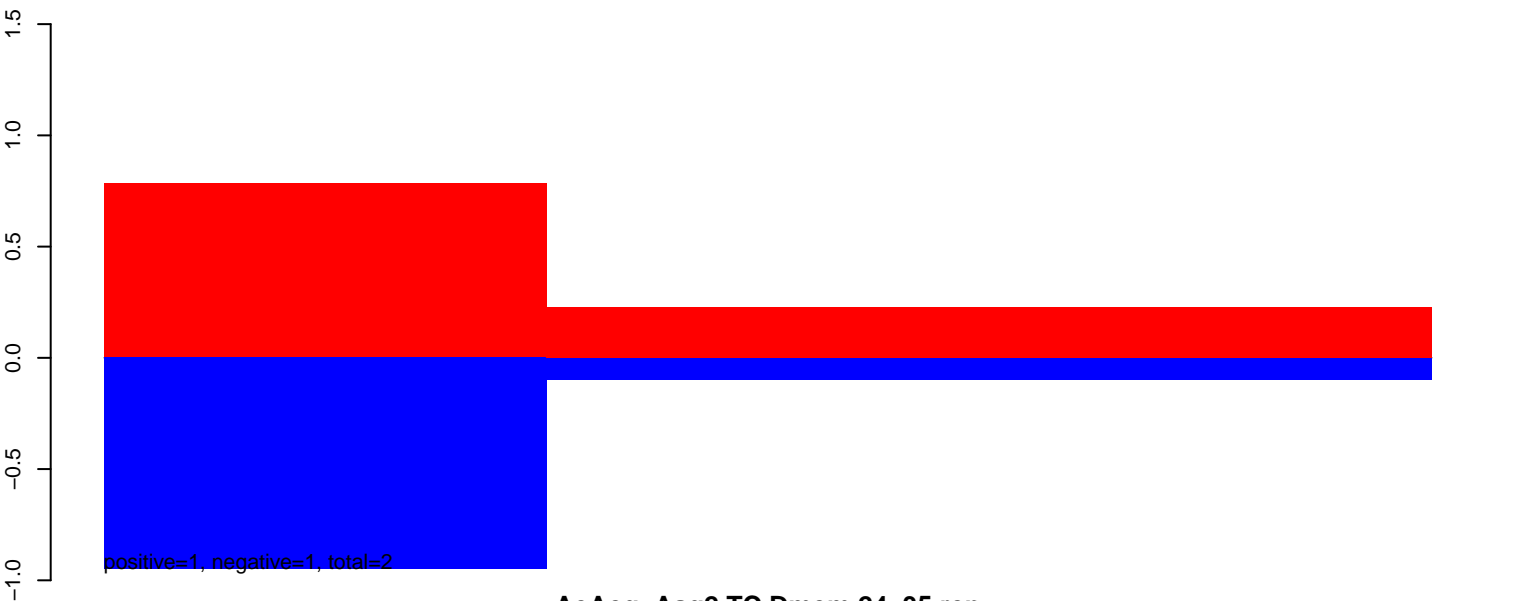
AeAeg_Aag2.TC.Dmem.24_35.rep



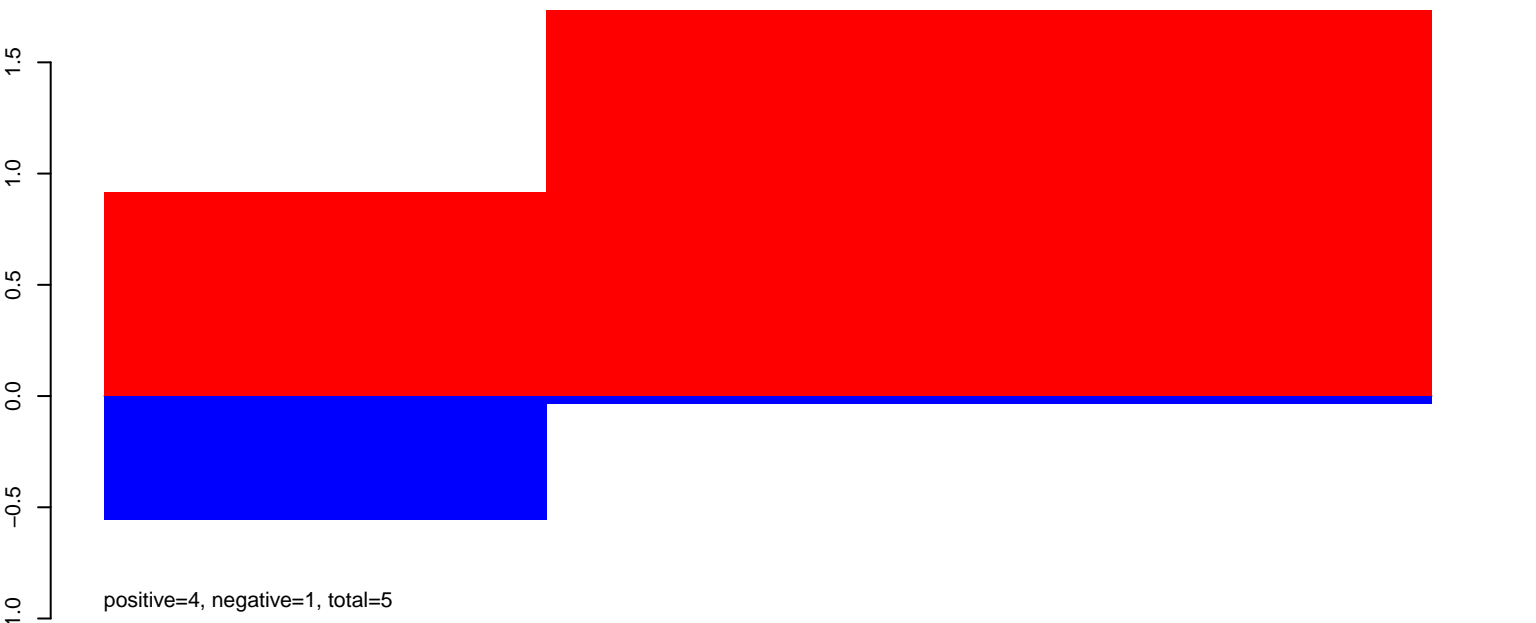
AeAeg_Aag2.TC.Dmem.rep



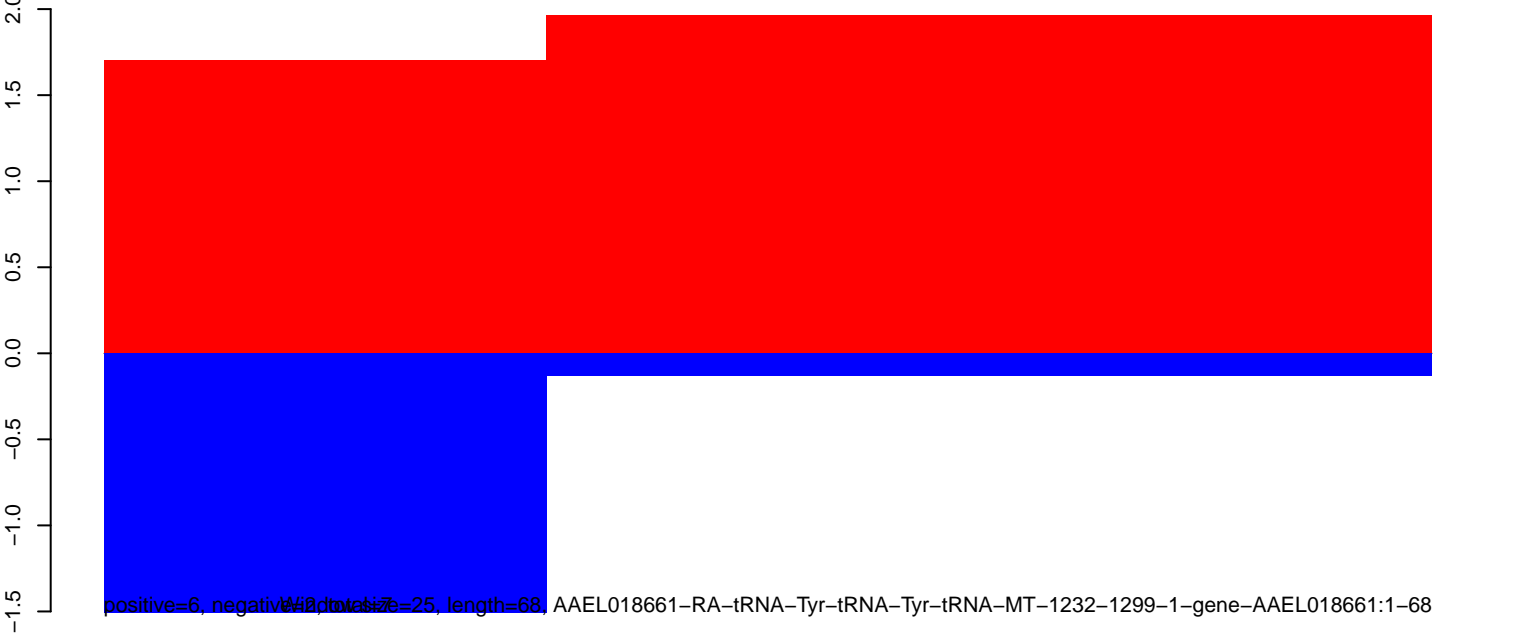
AeAeg_Aag2.TC.Dmem.18_23.rep



AeAeg_Aag2.TC.Dmem.24_35.rep



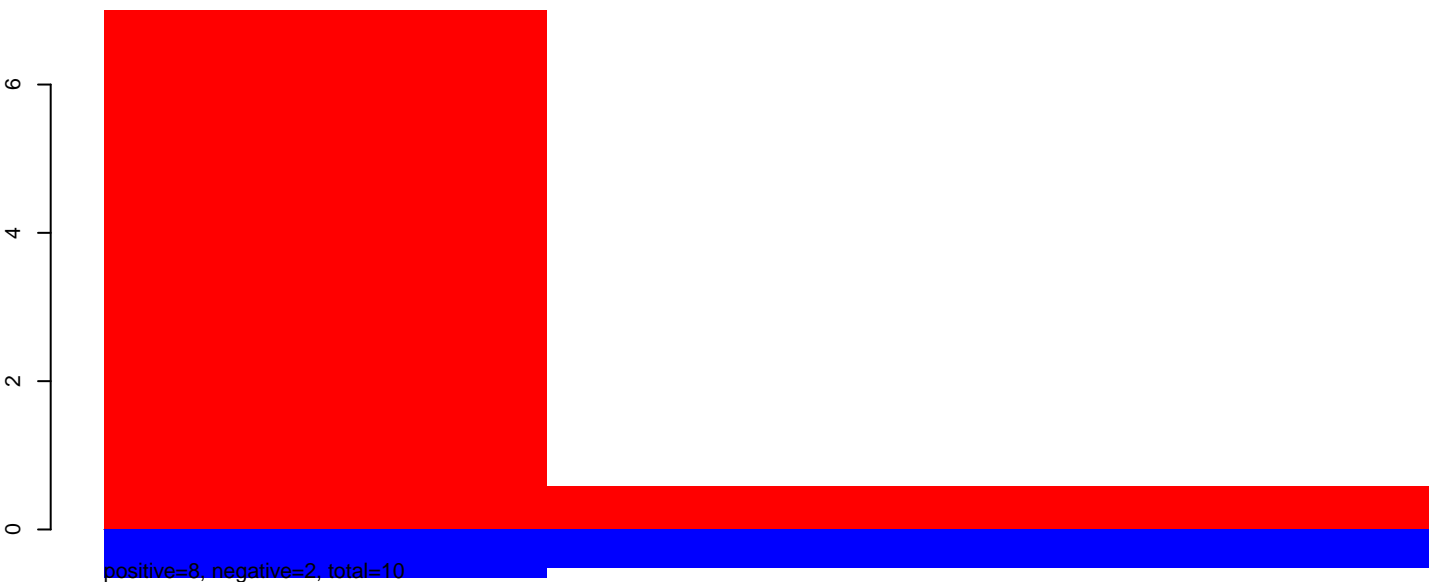
AeAeg_Aag2.TC.Dmem.rep



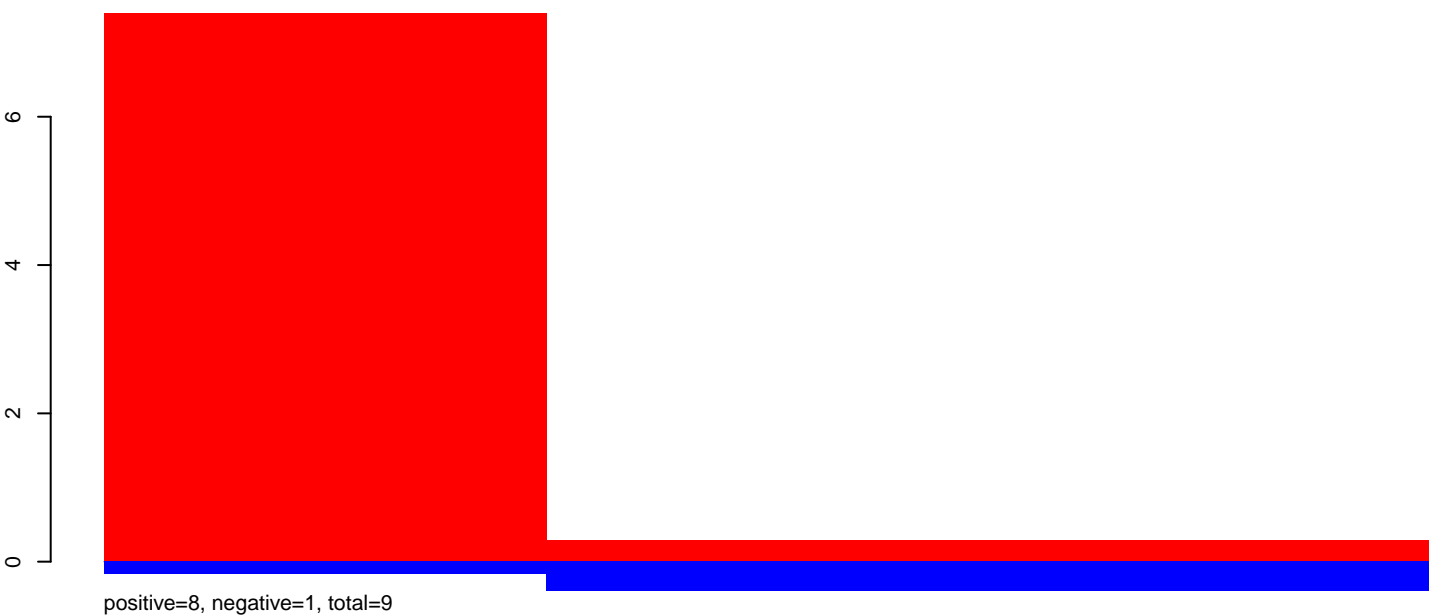
AAEL018661-RA-tRNA-Tyr-tRNA-Tyr-tRNA-MT-1232-1299-1-gene-AAEL018661:1-68

40 60 80 100

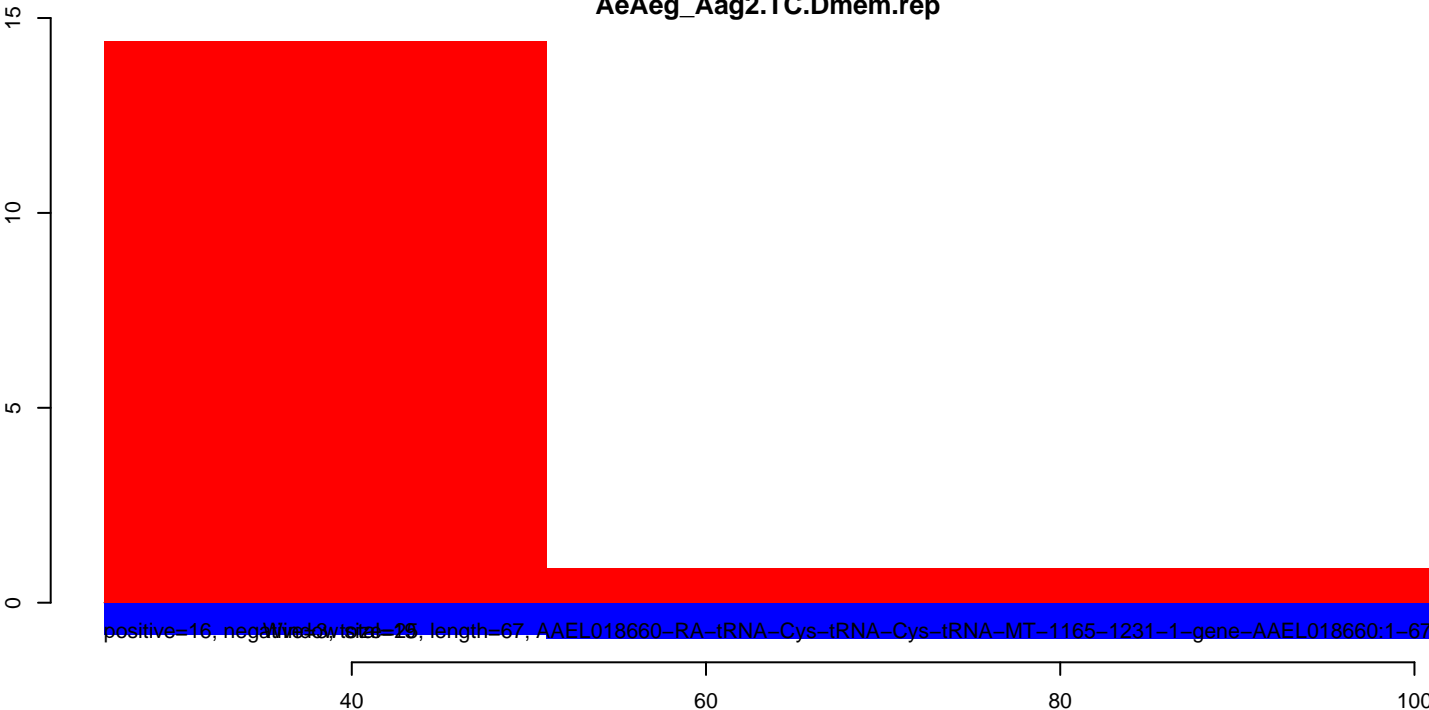
AeAeg_Aag2.TC.Dmem.18_23.rep



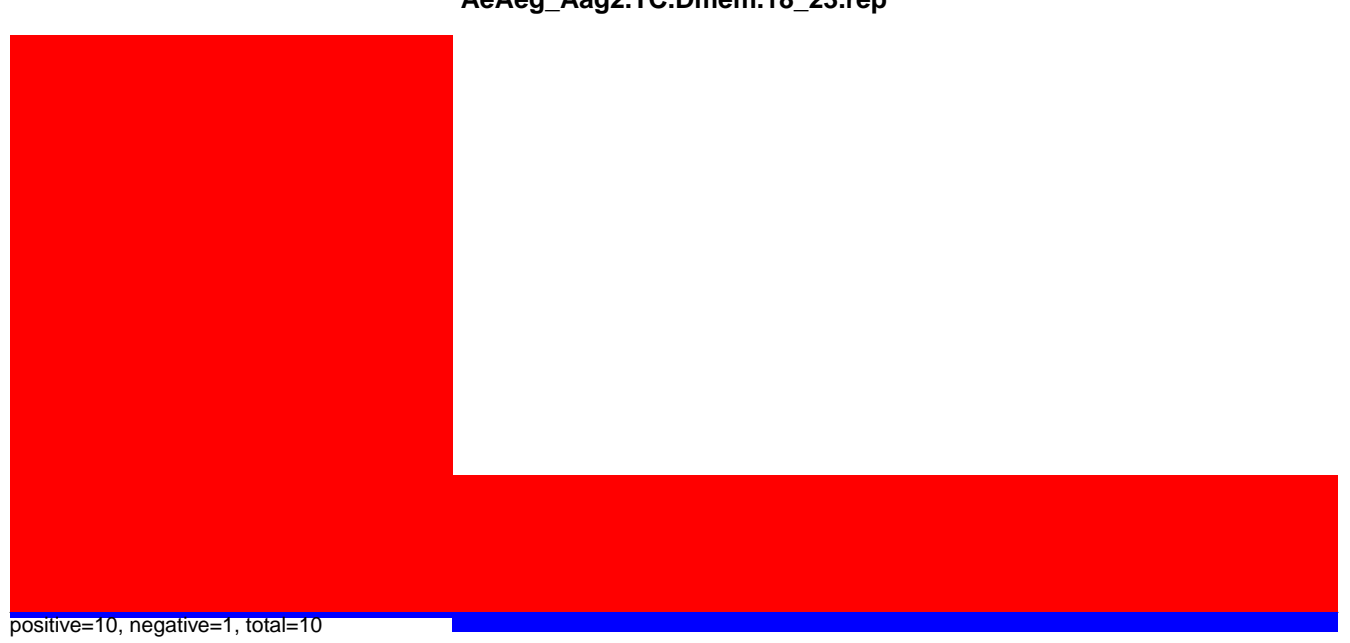
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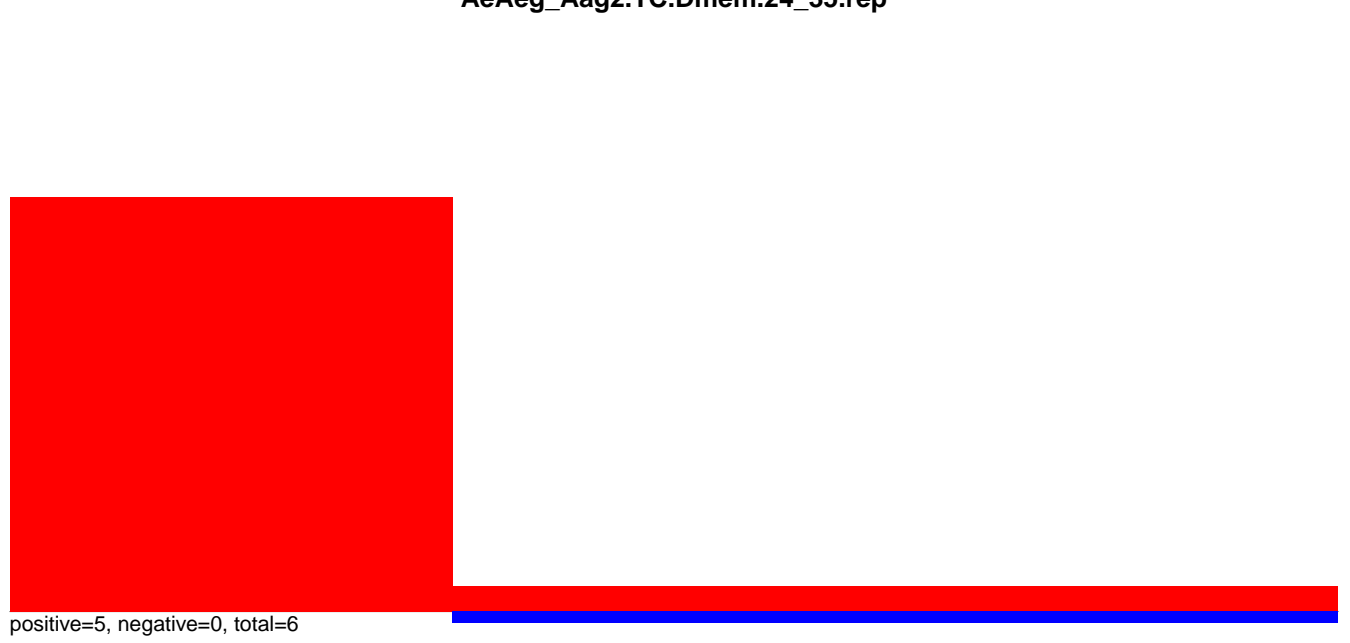
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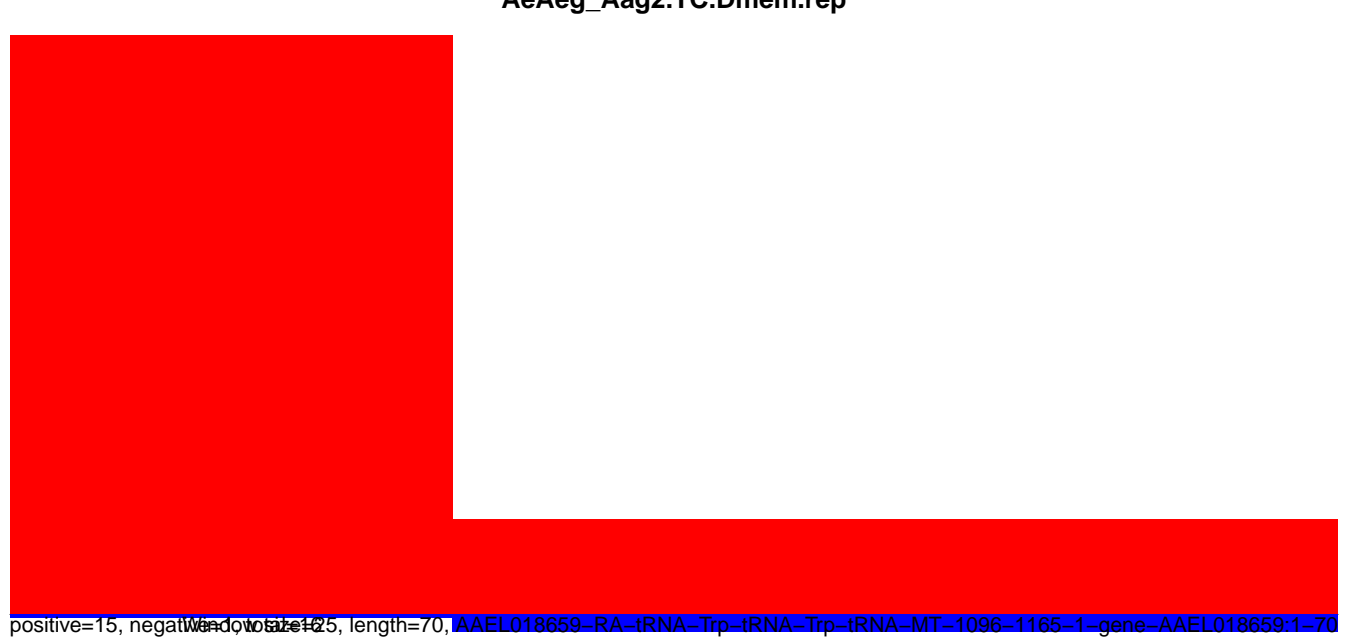
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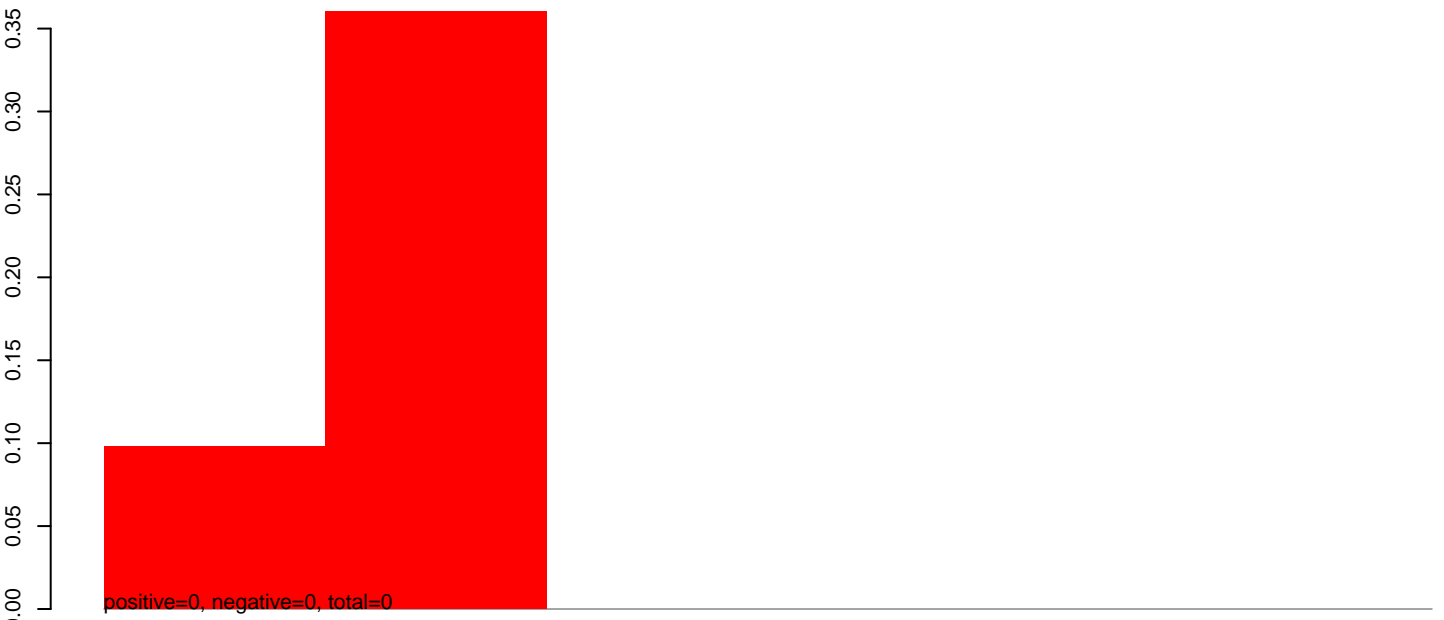


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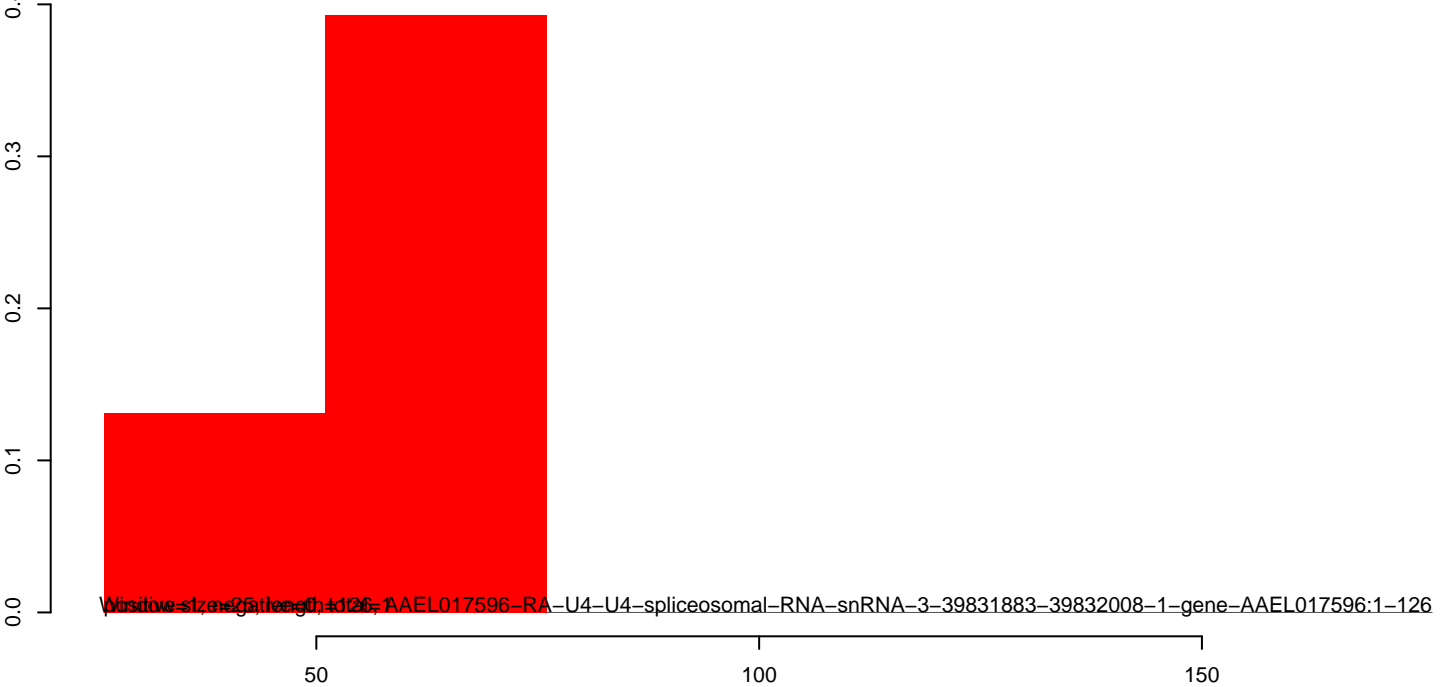
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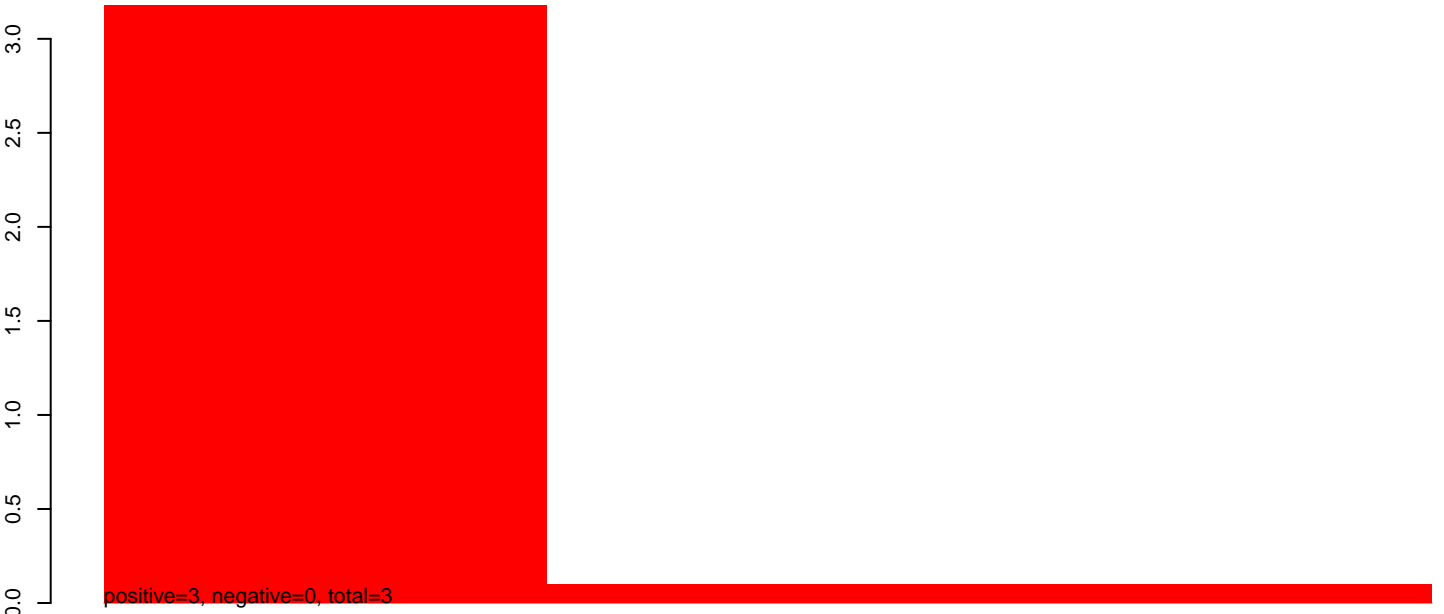
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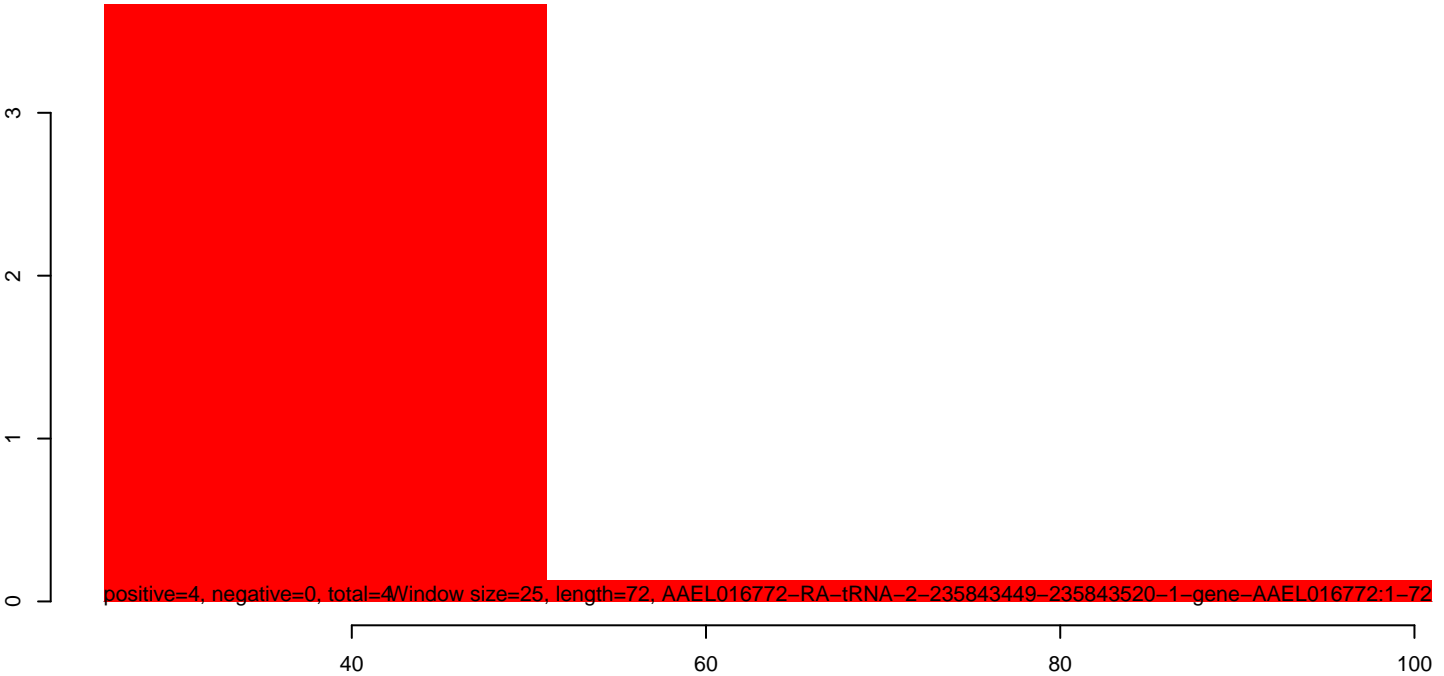
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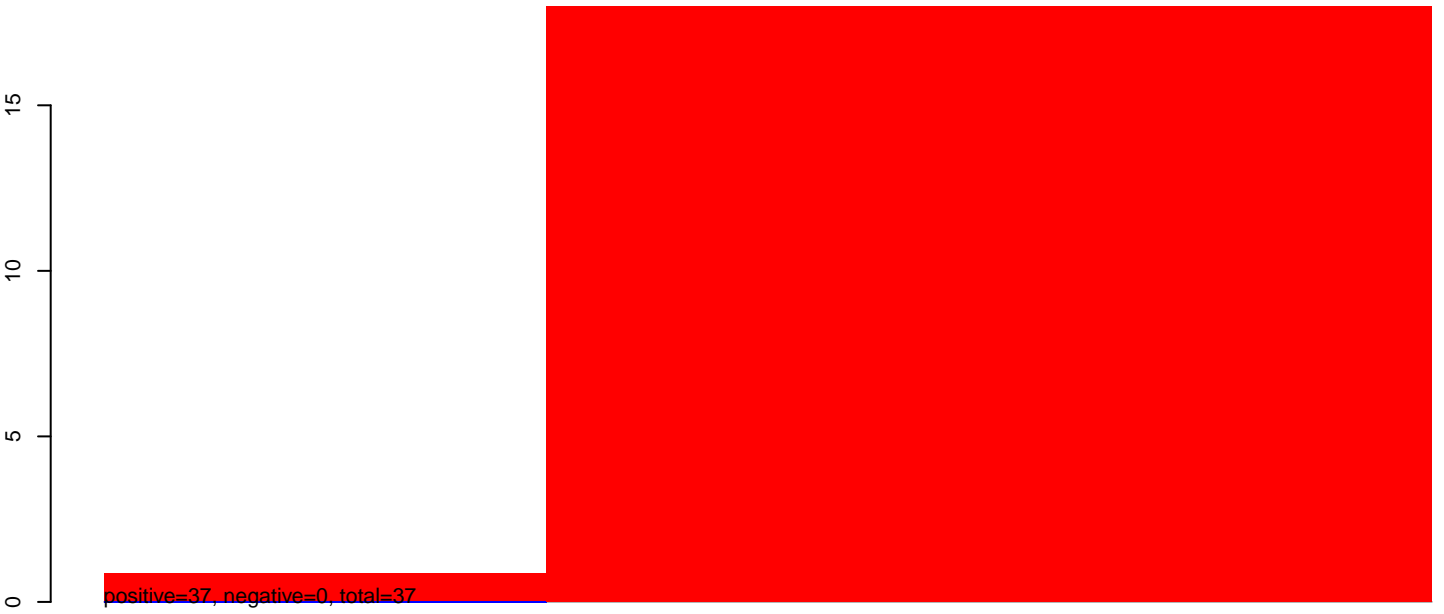
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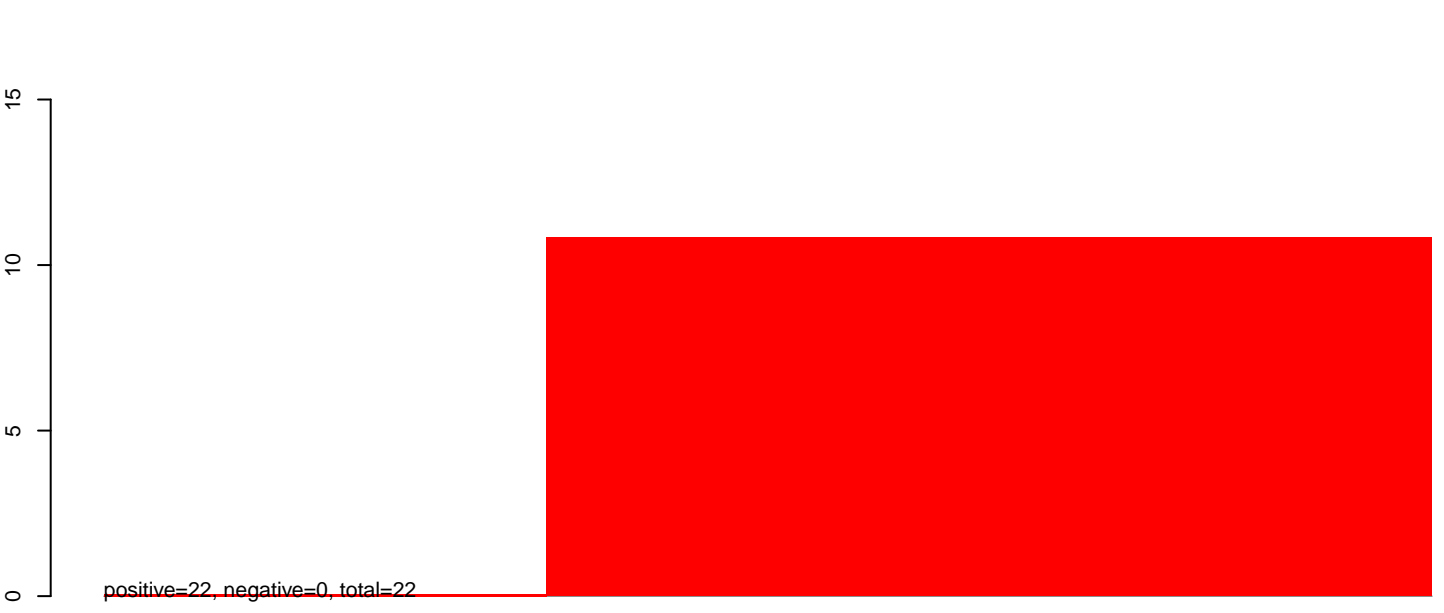
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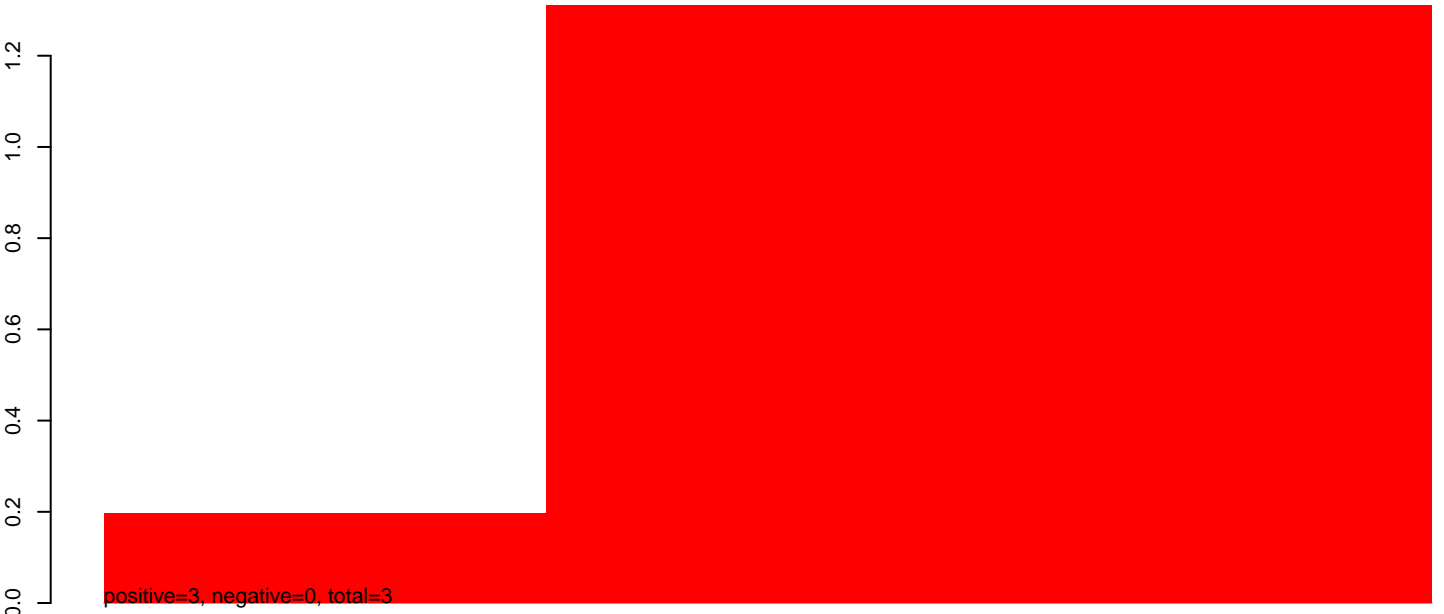


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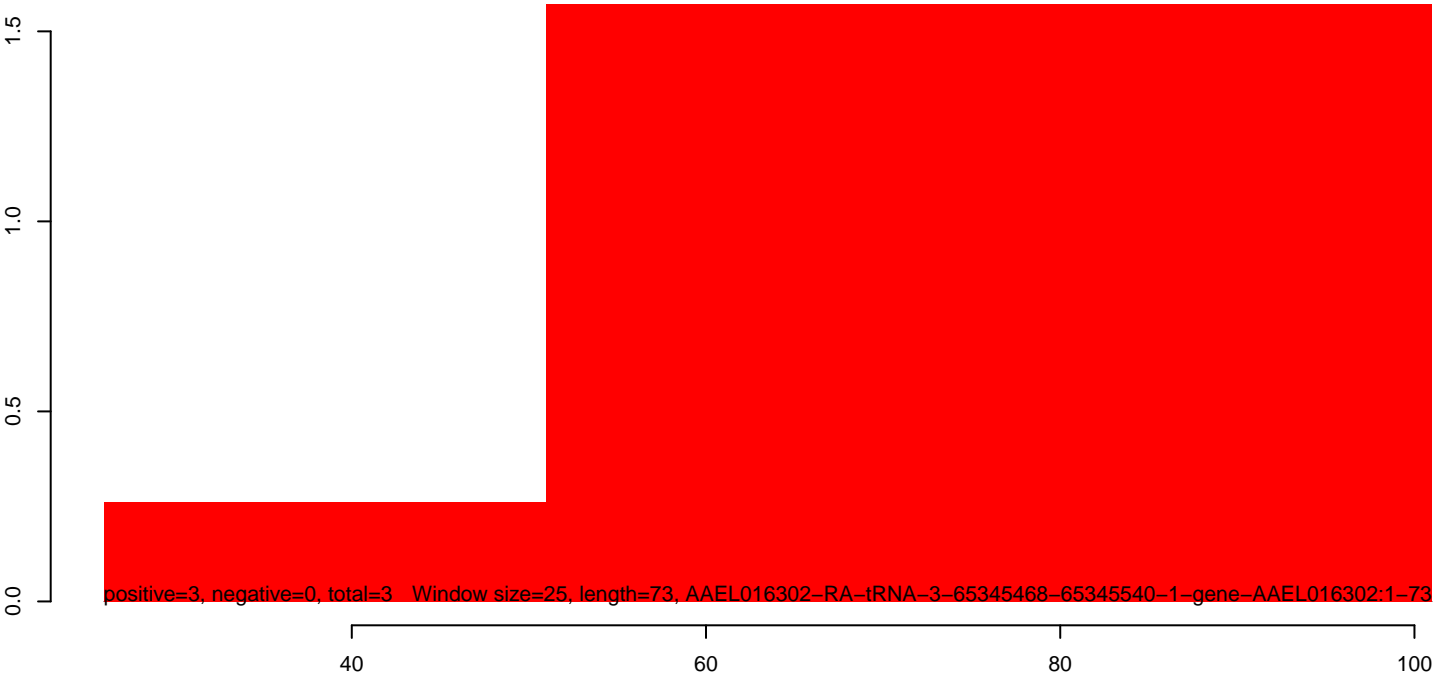
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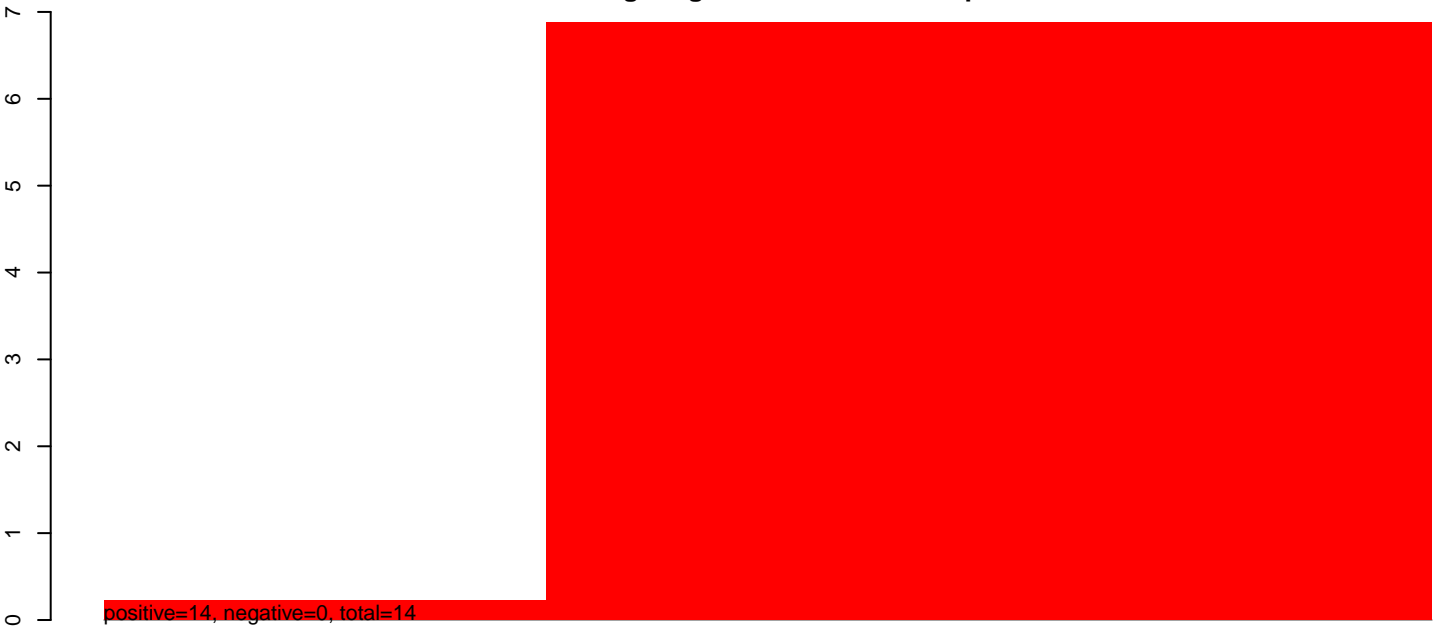
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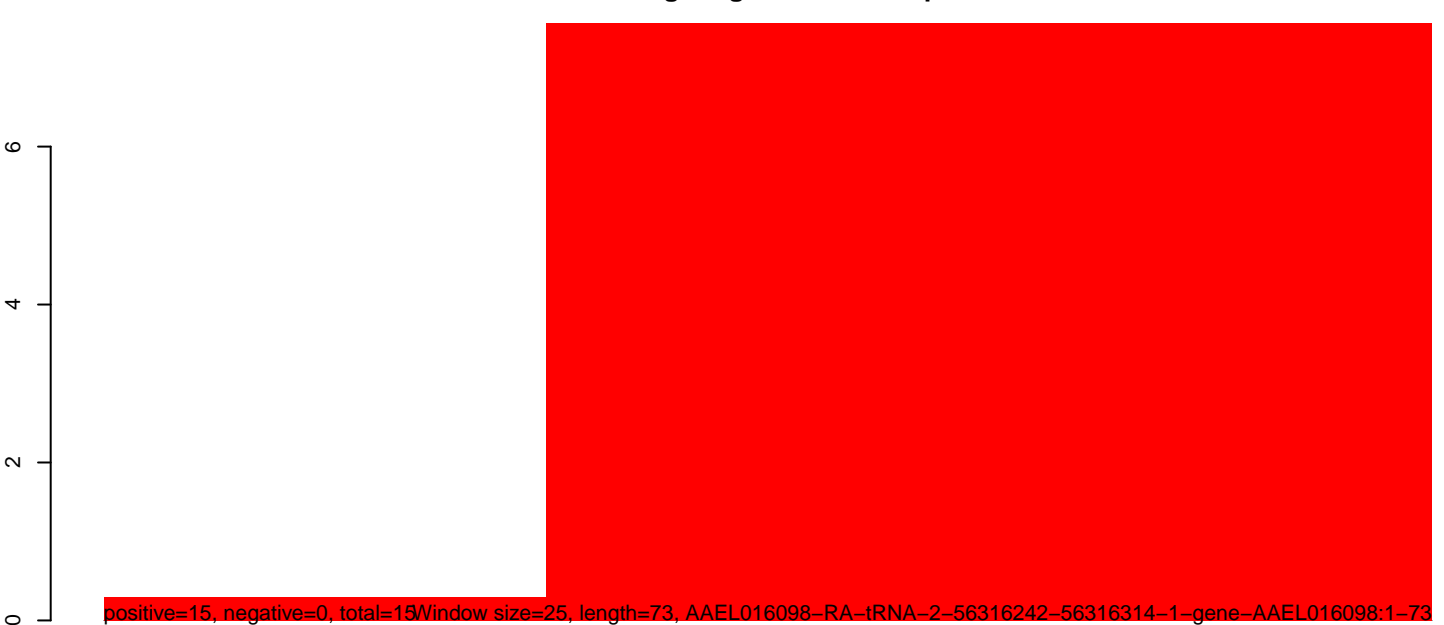
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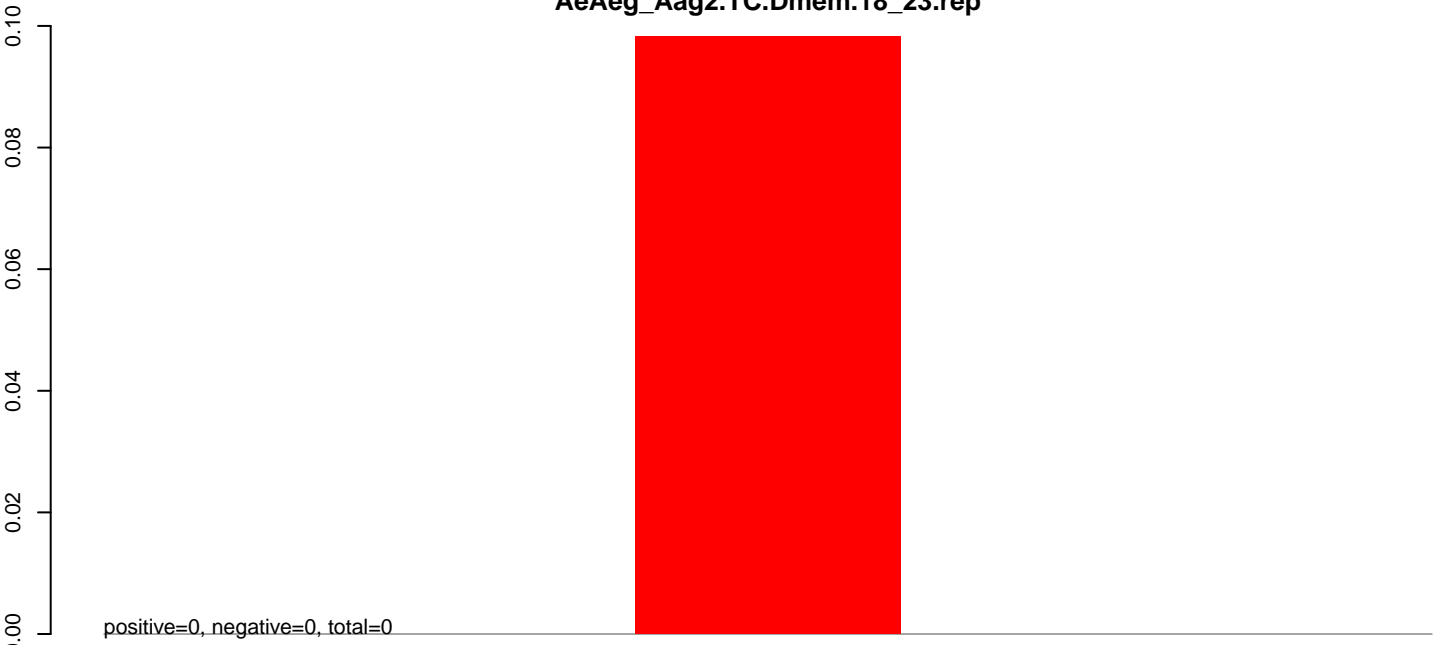


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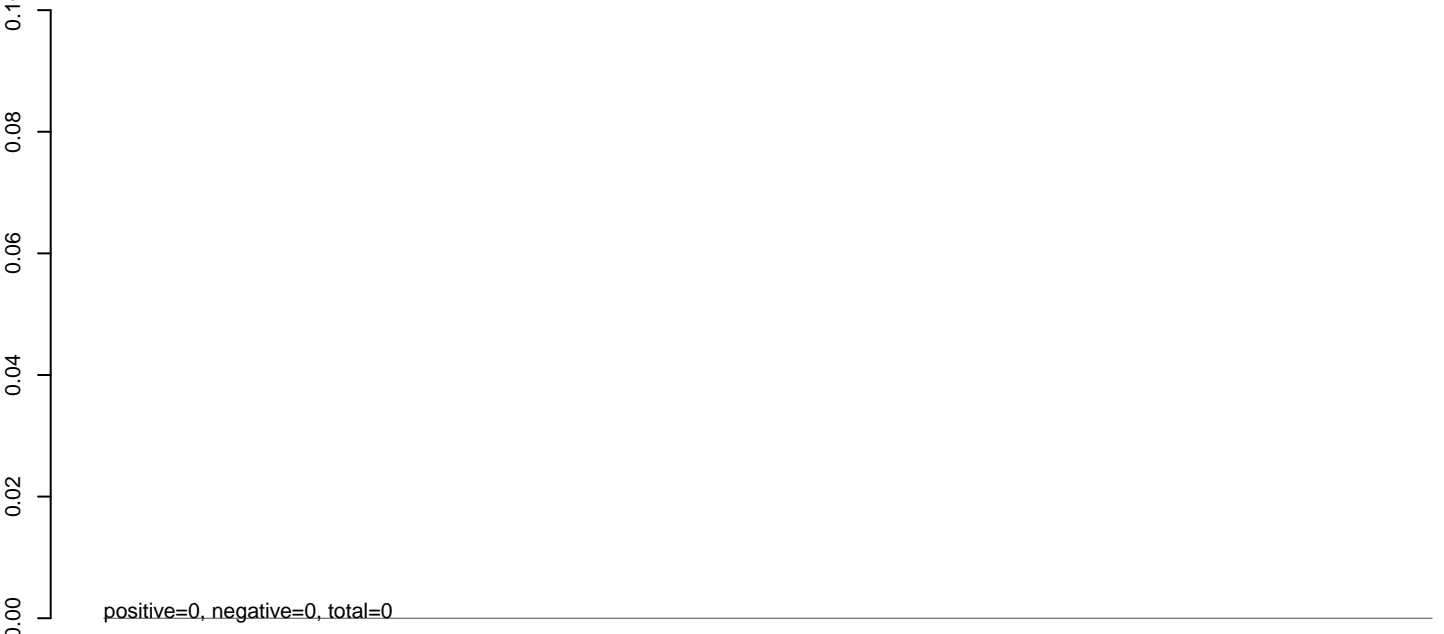


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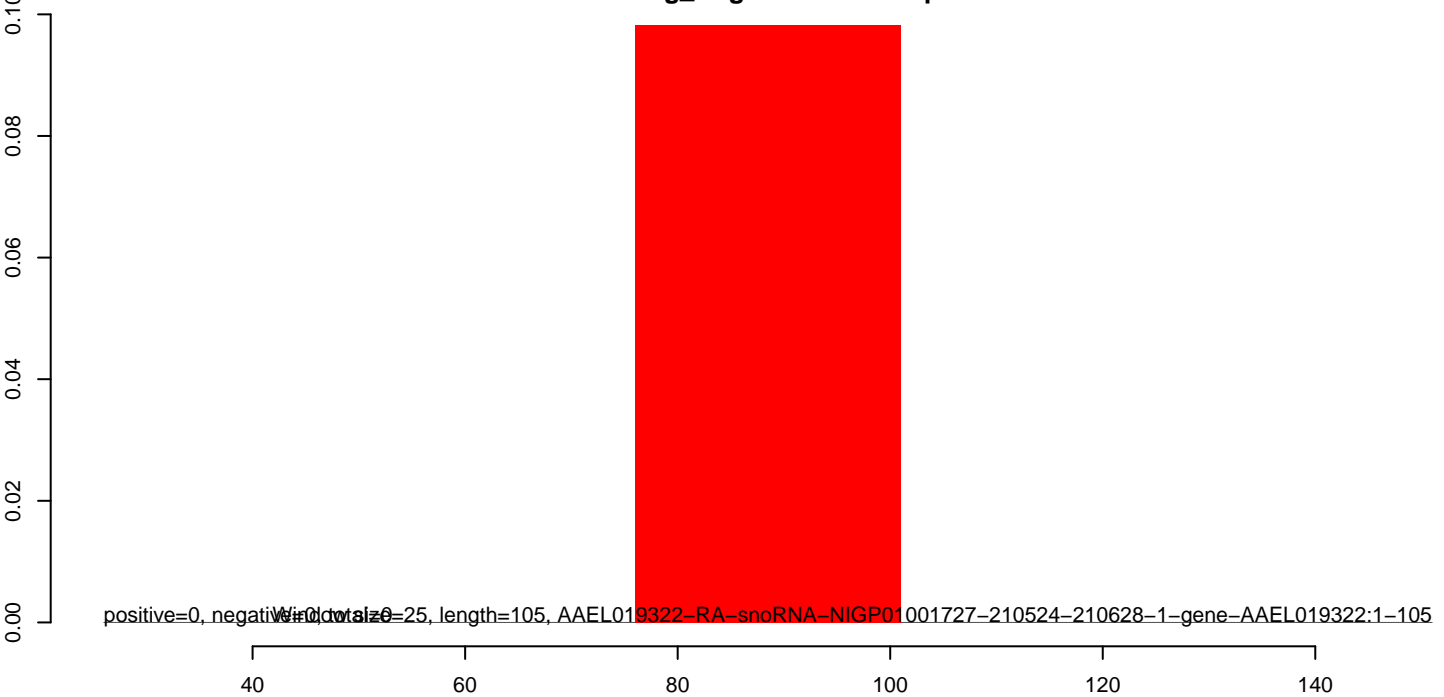
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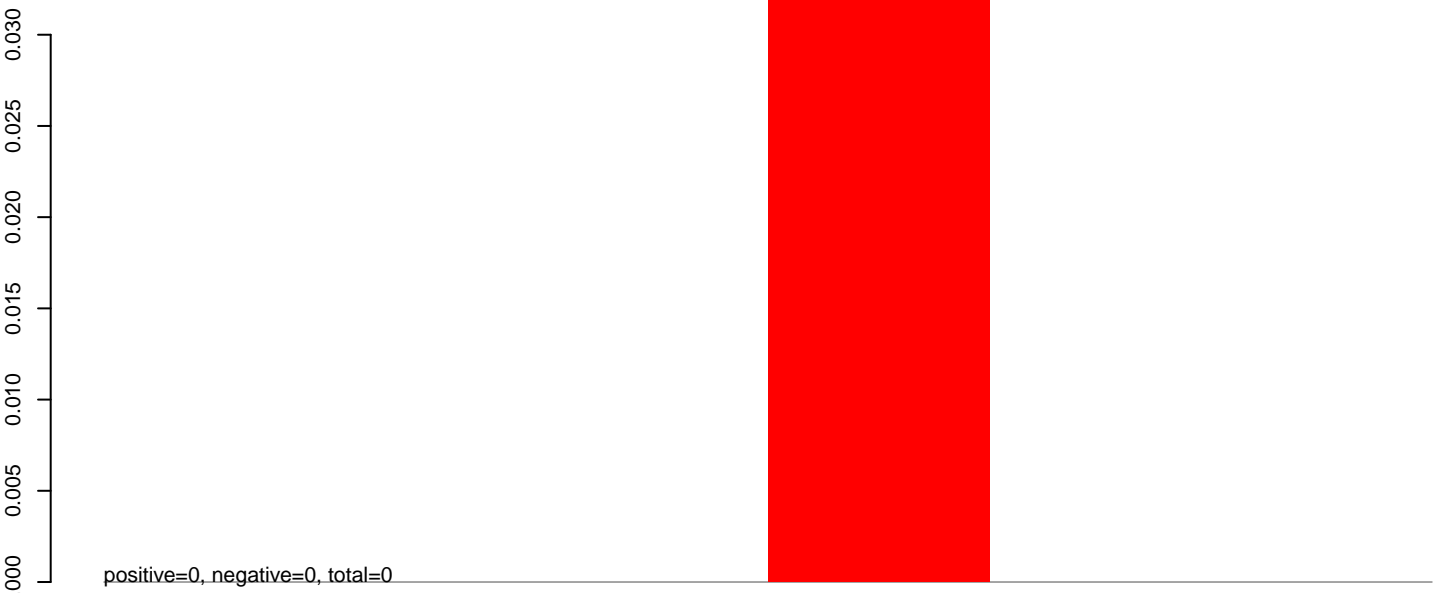
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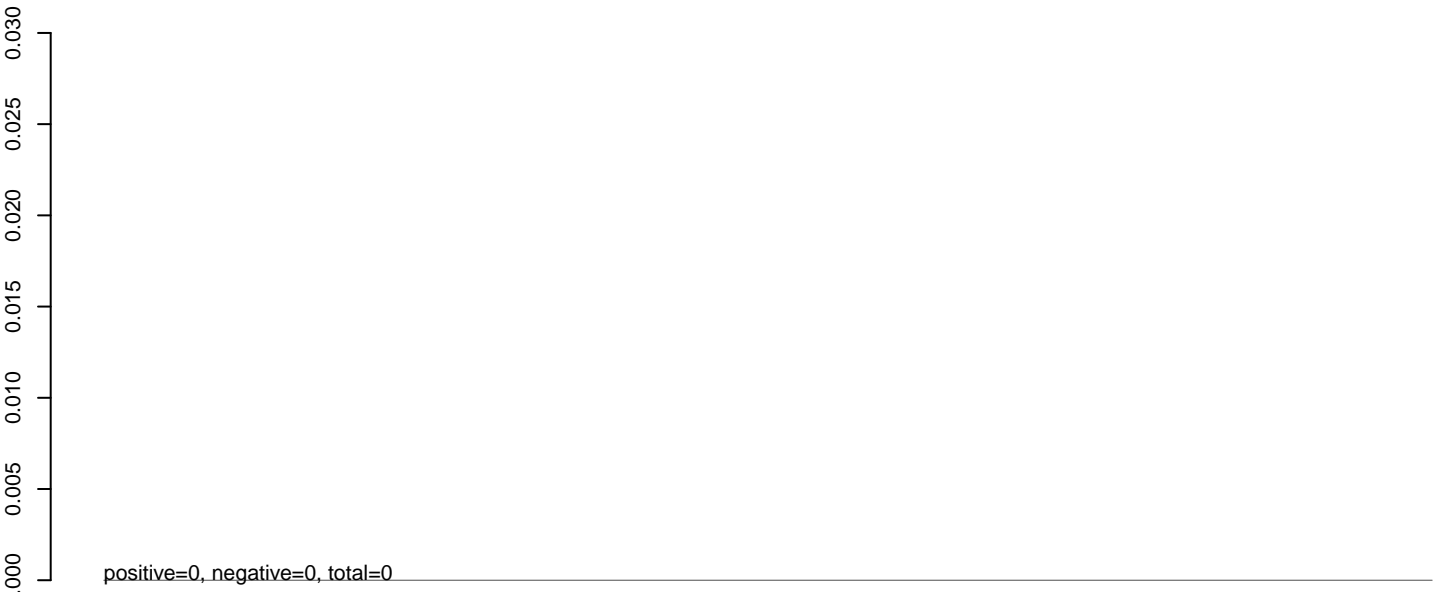
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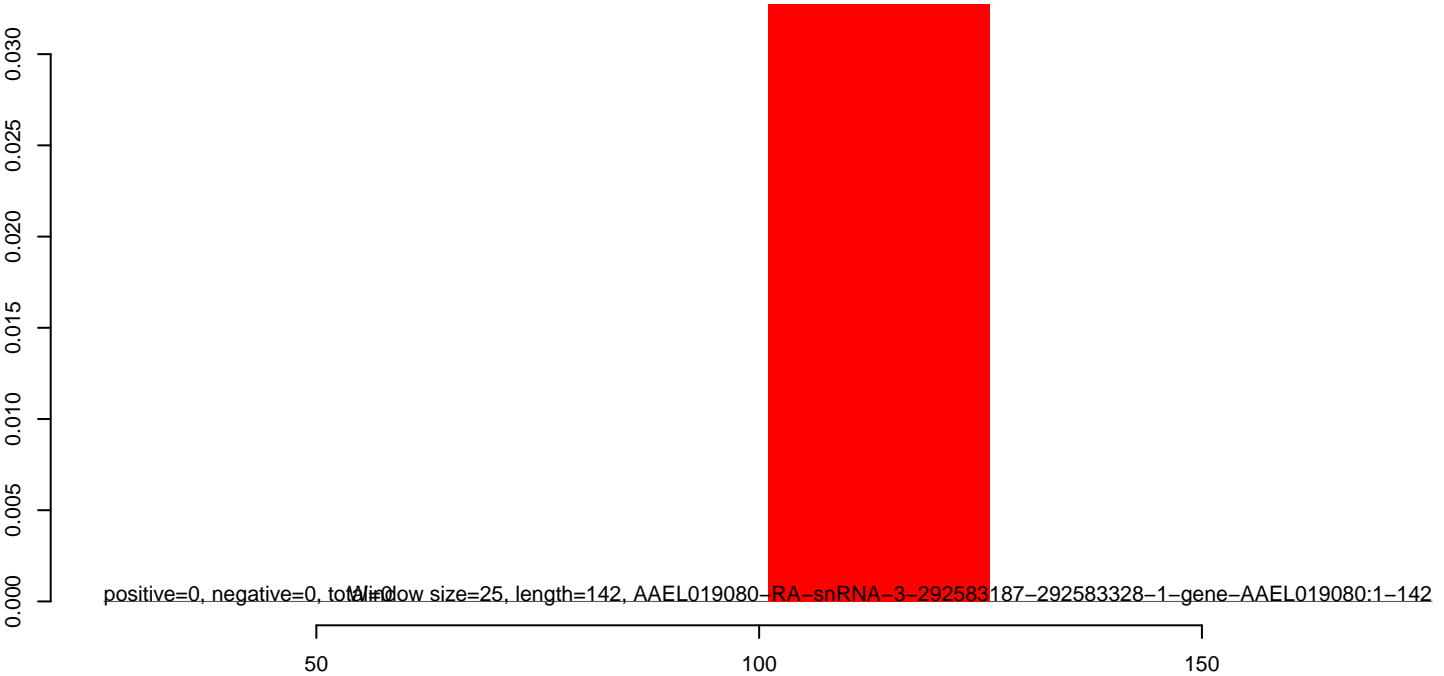
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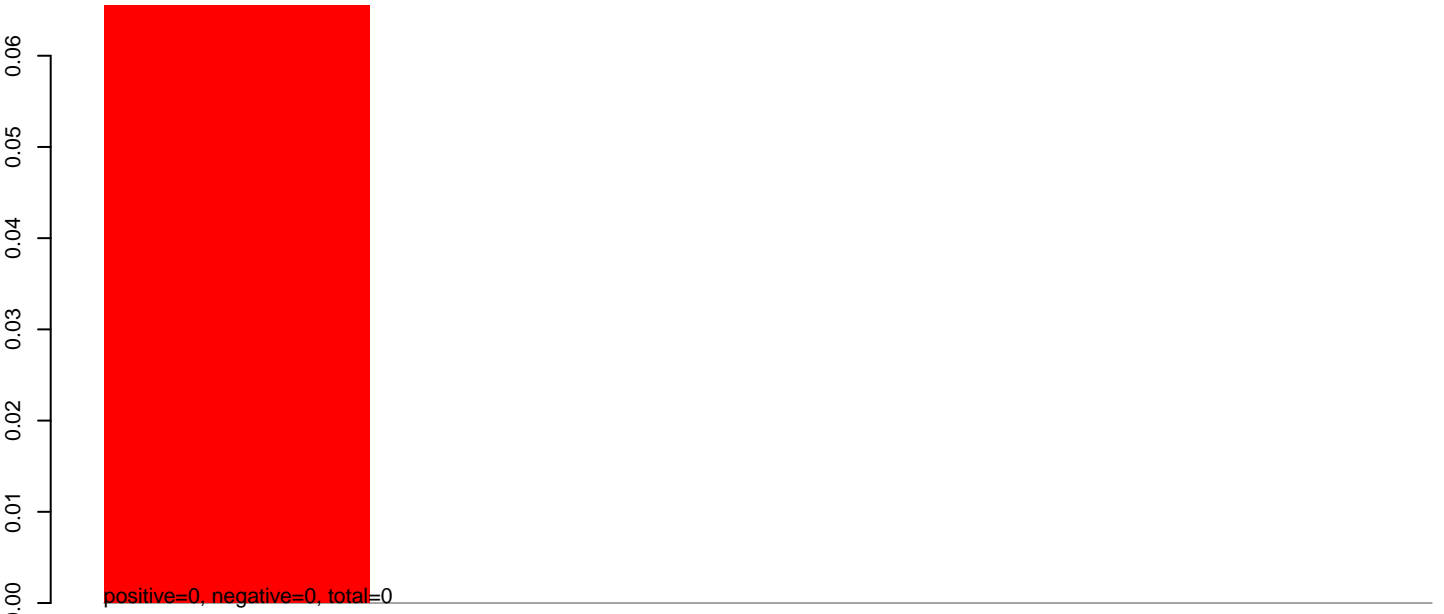
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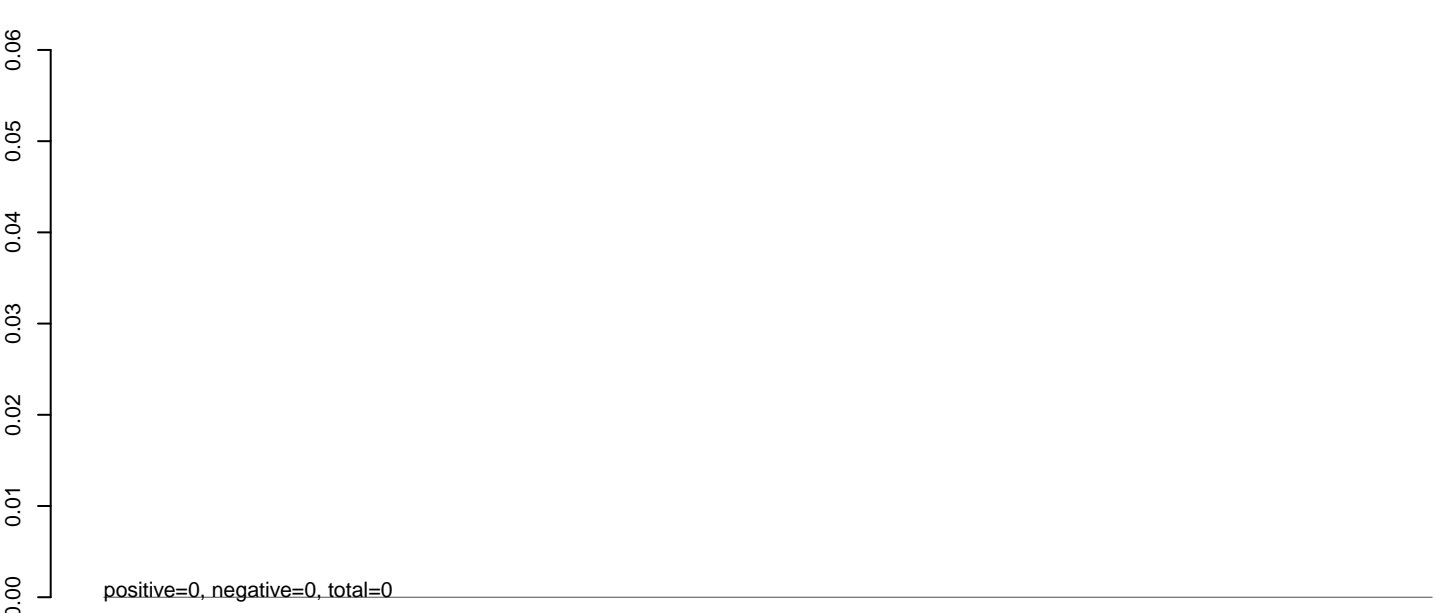
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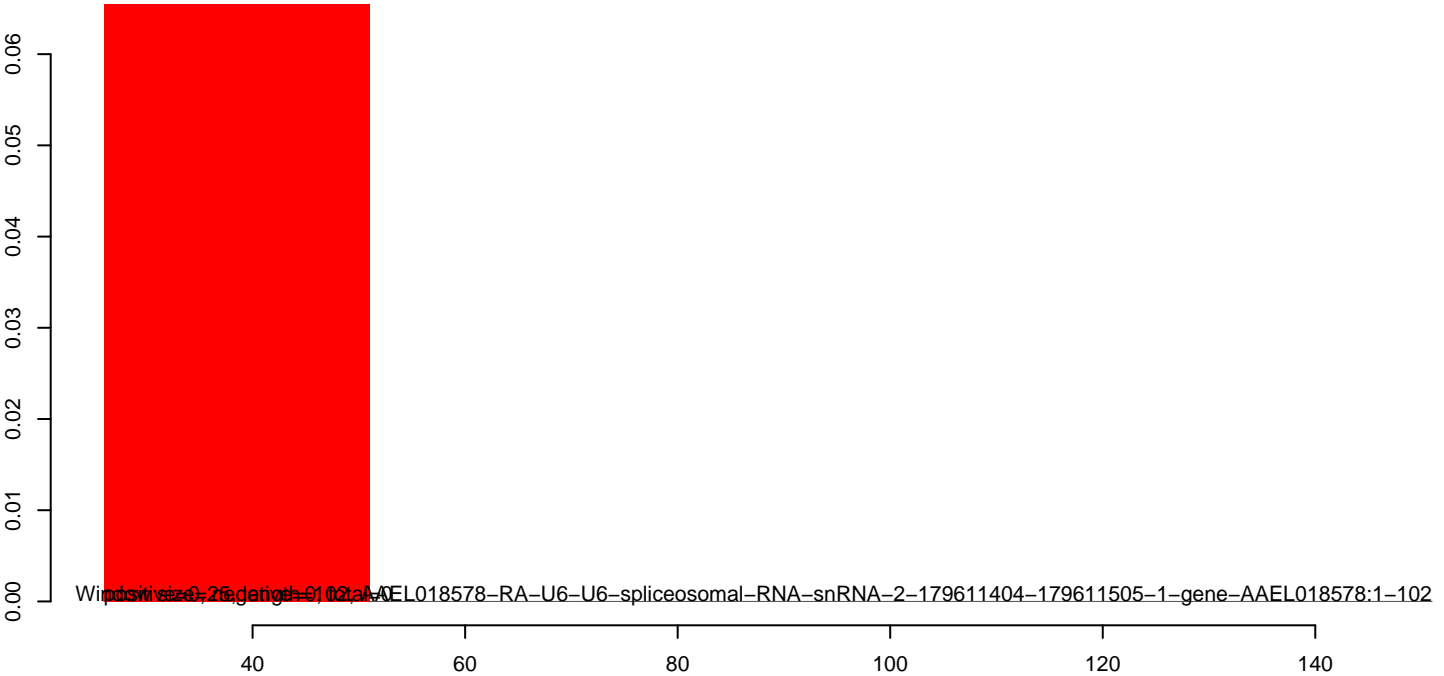
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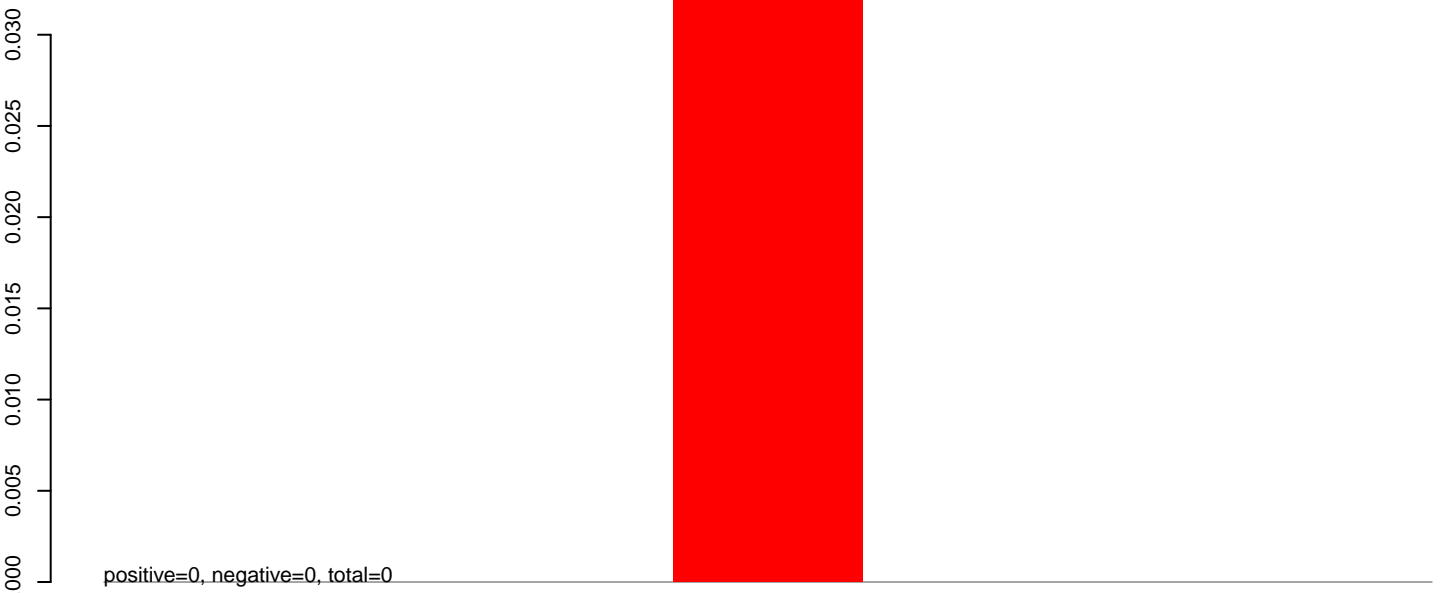
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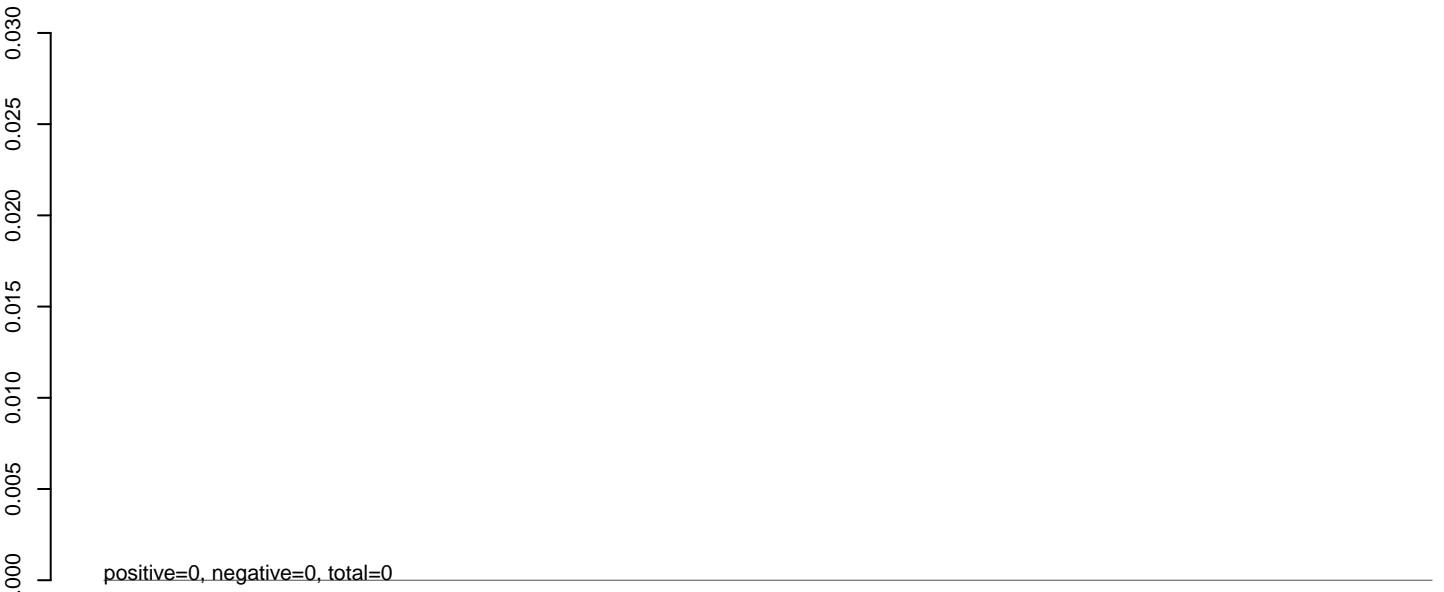
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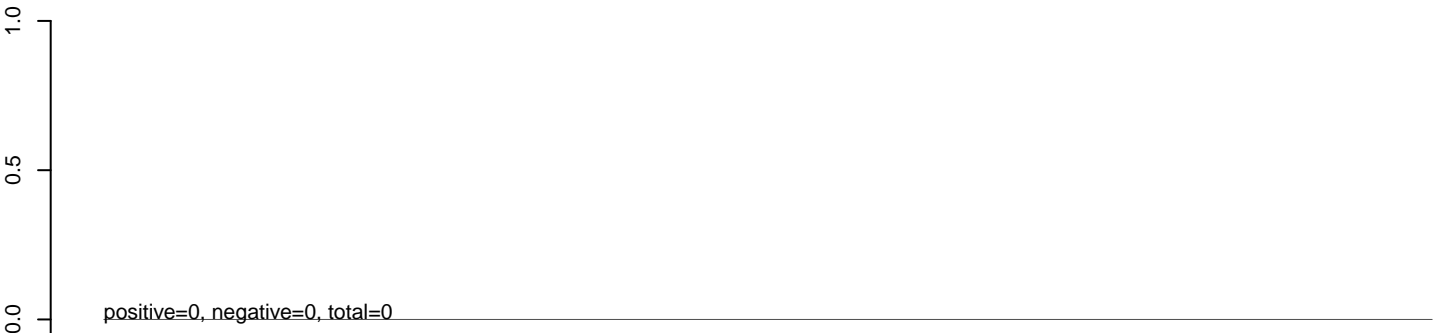
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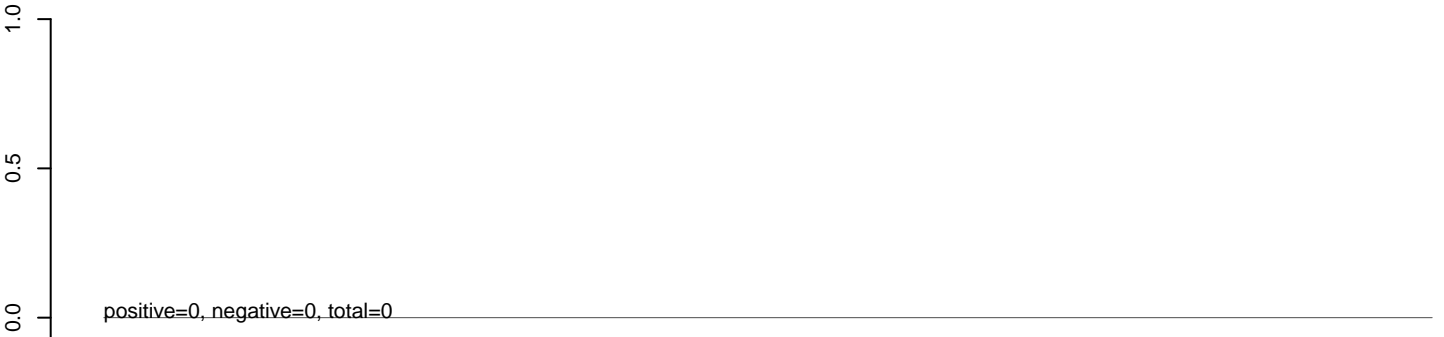
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