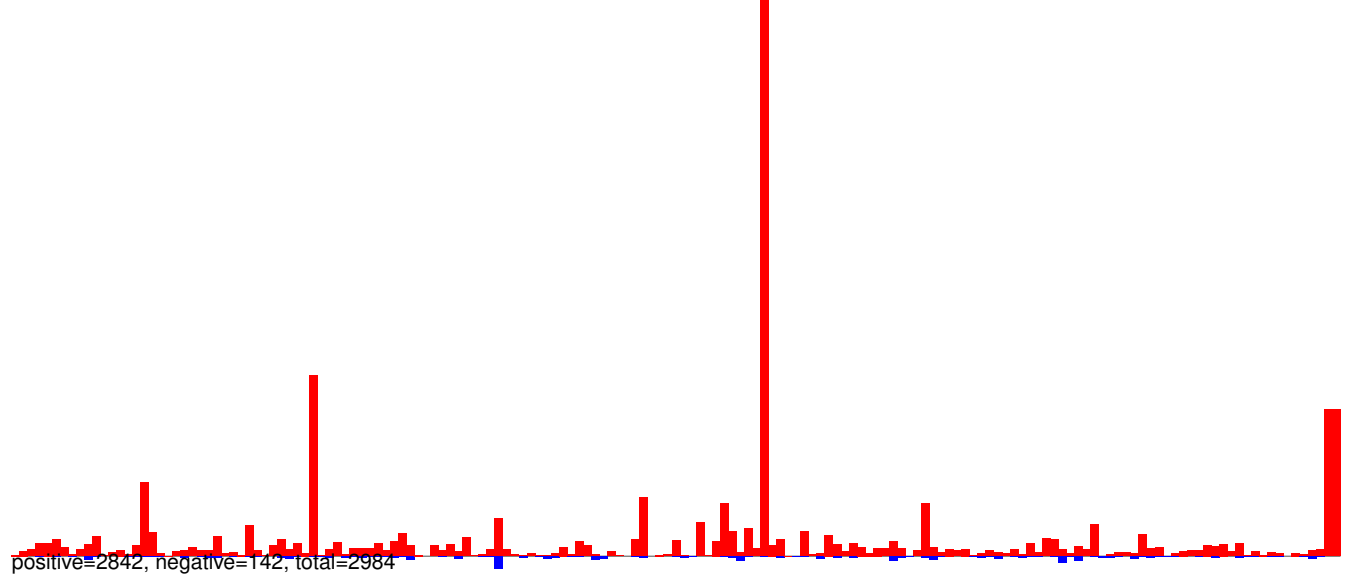
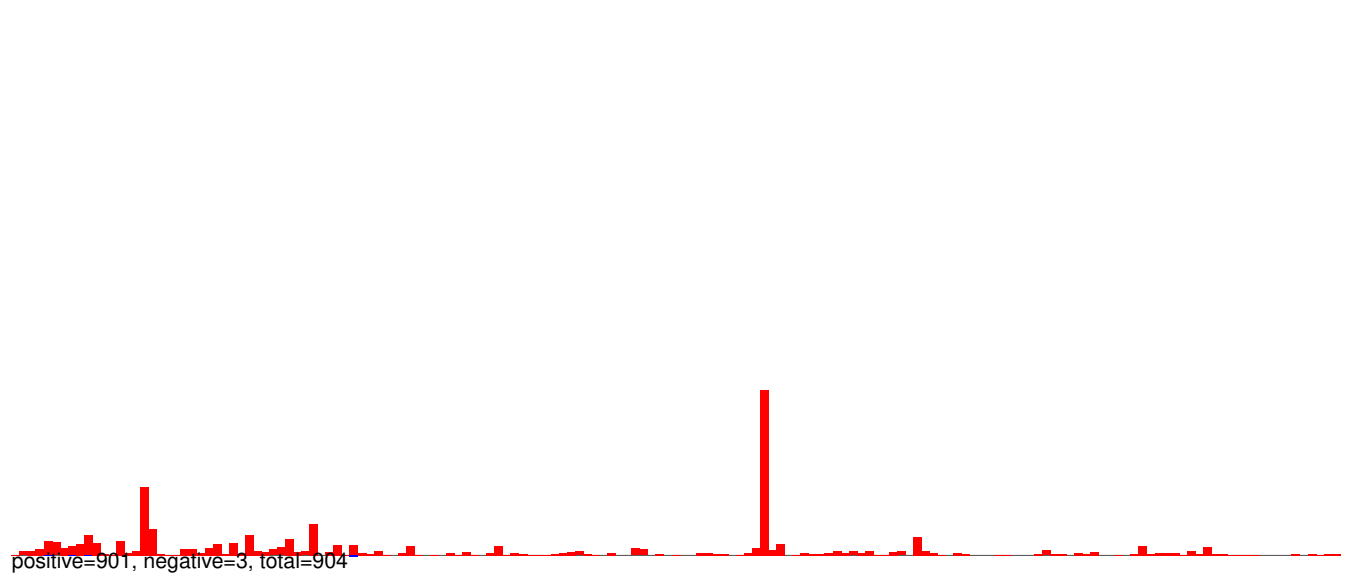


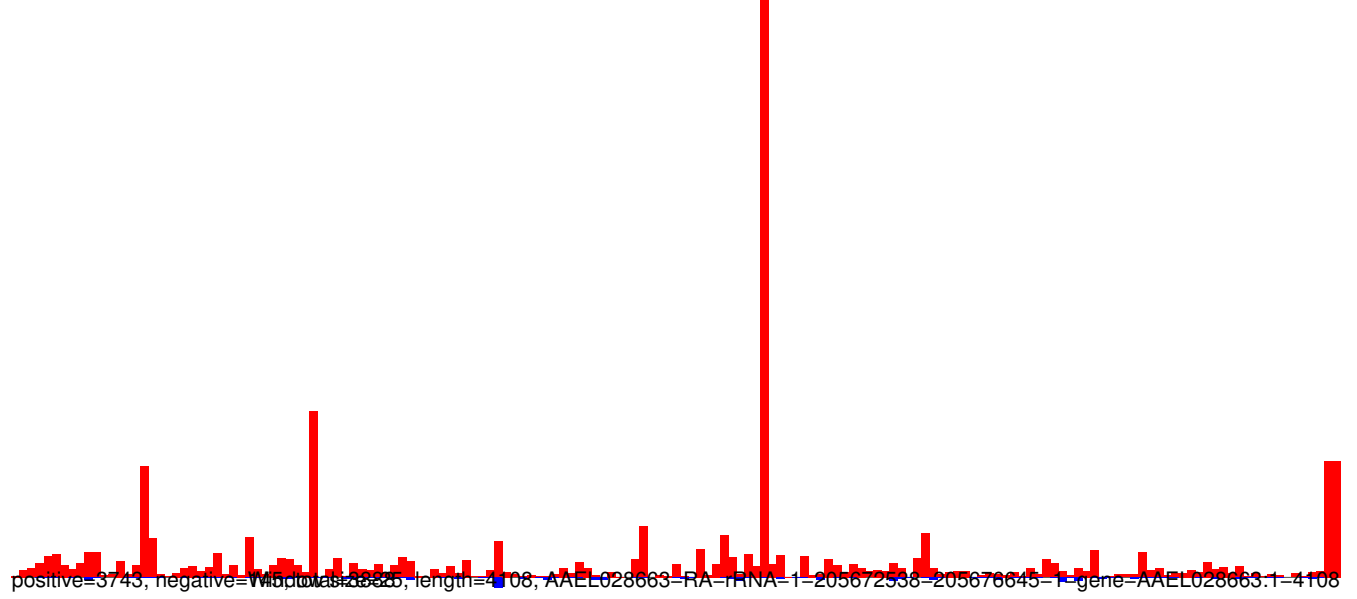
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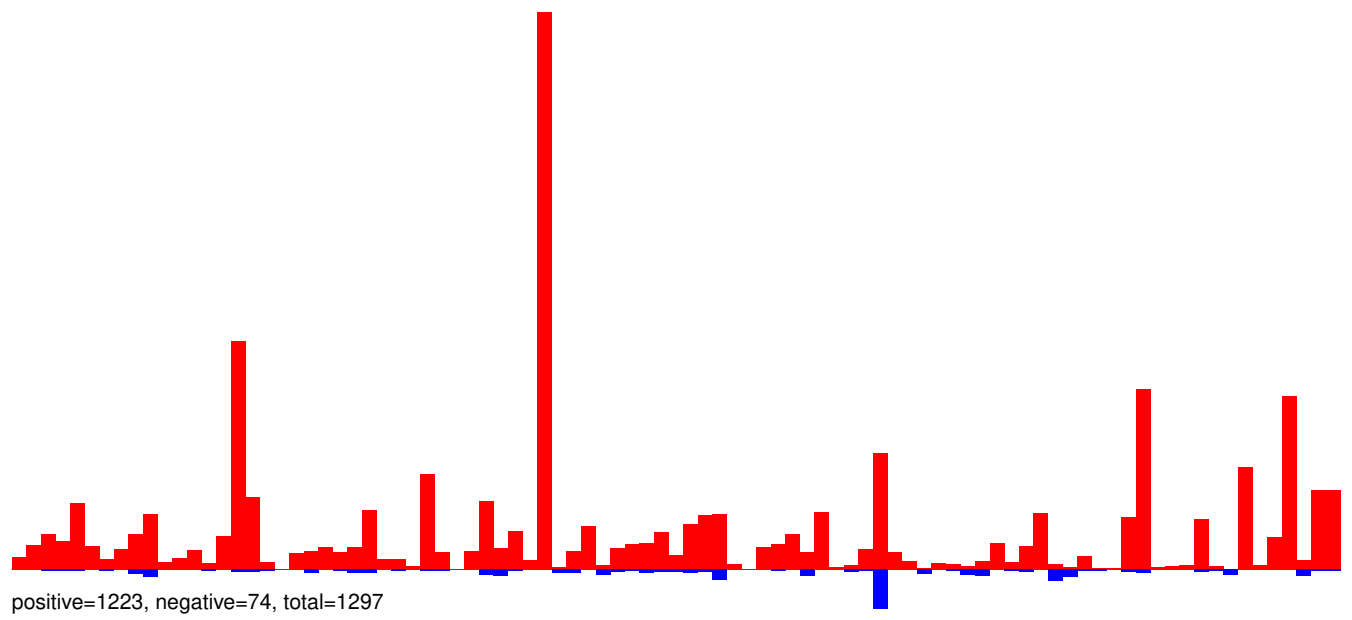
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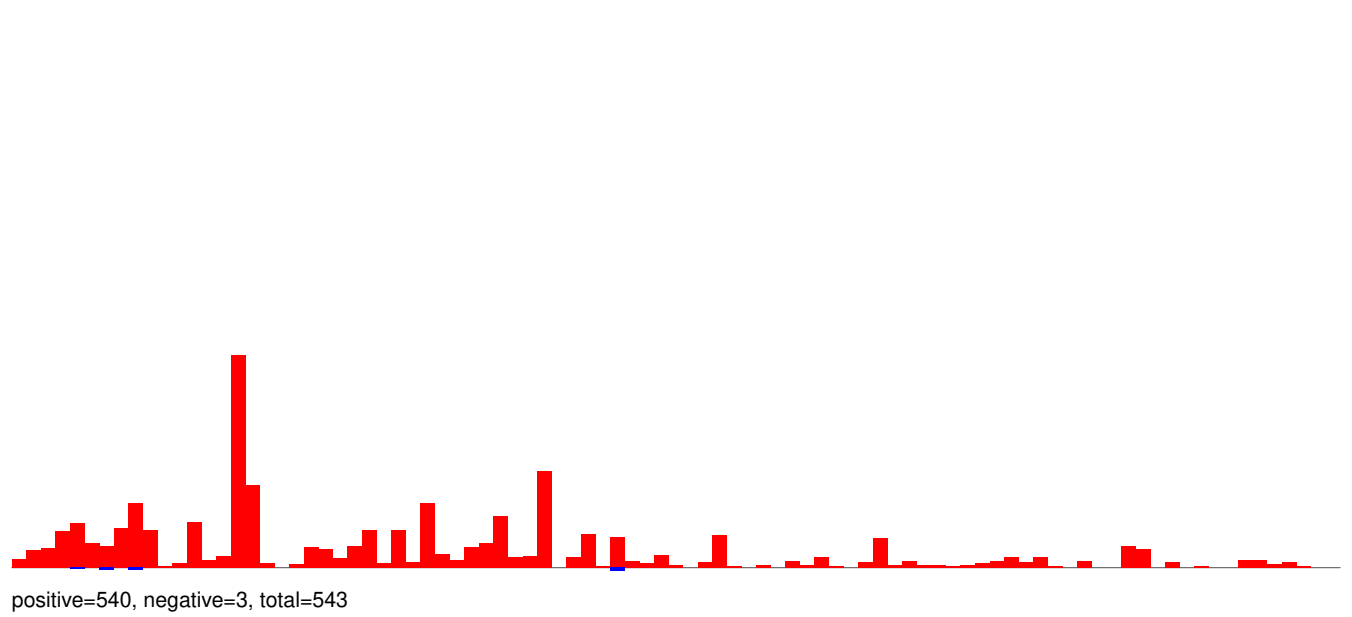
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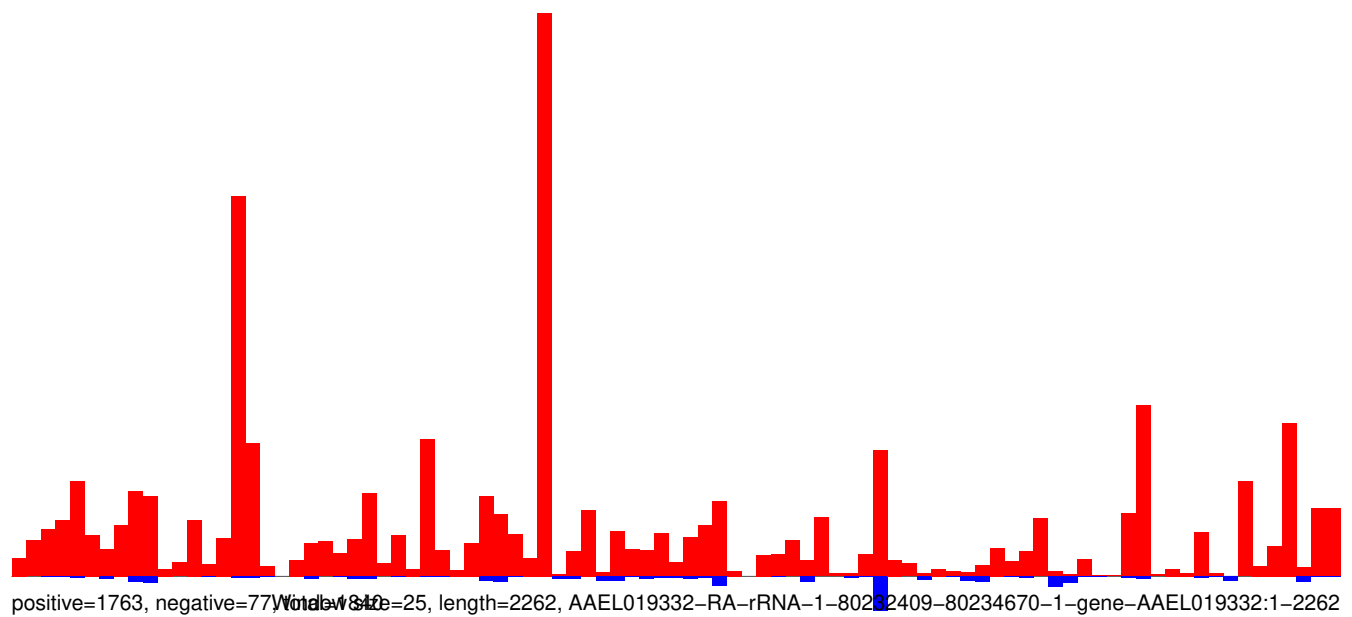
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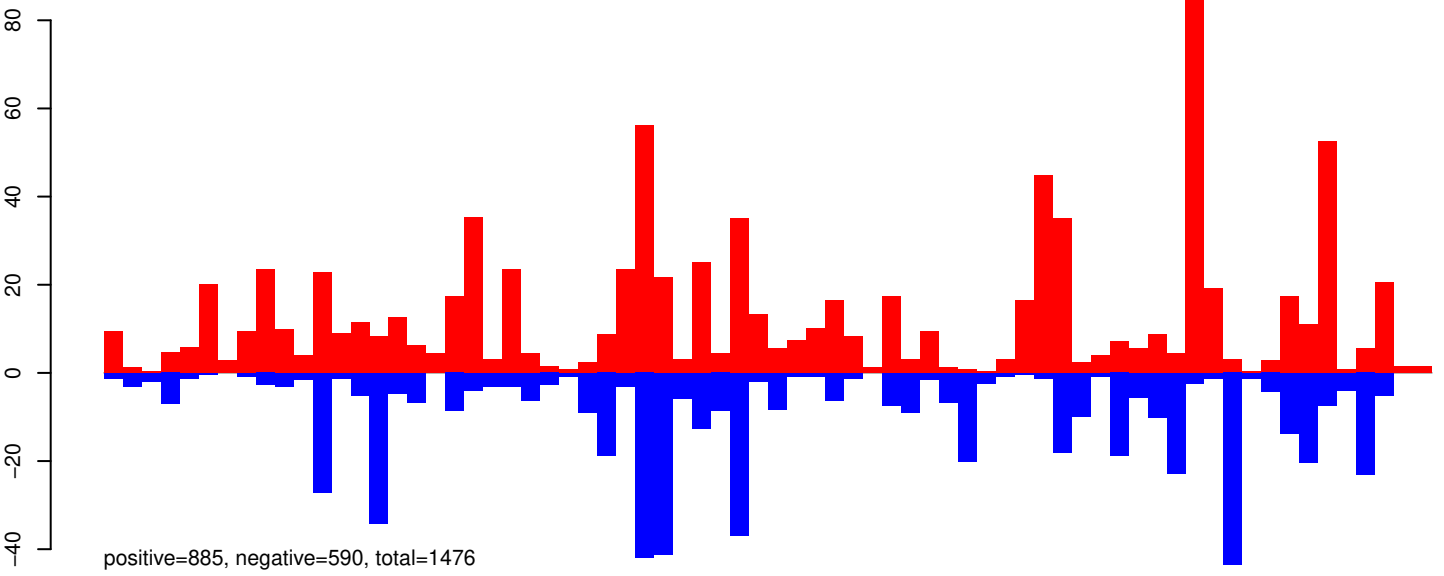
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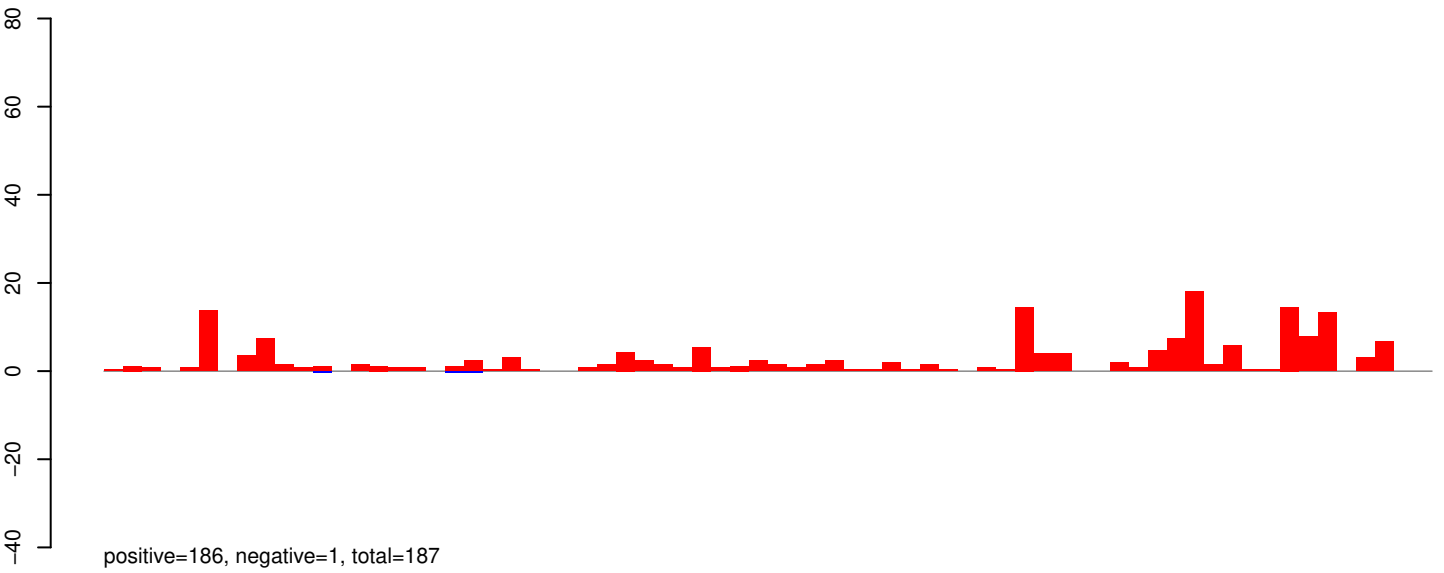
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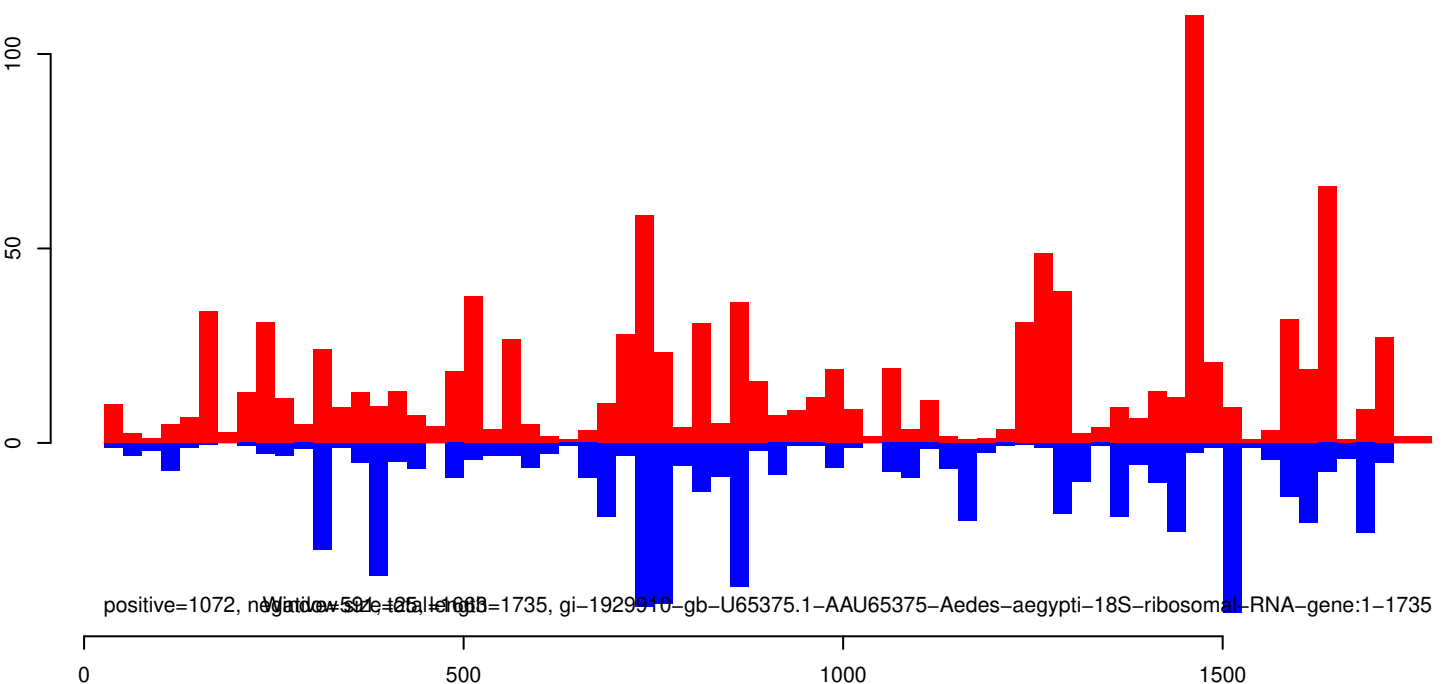
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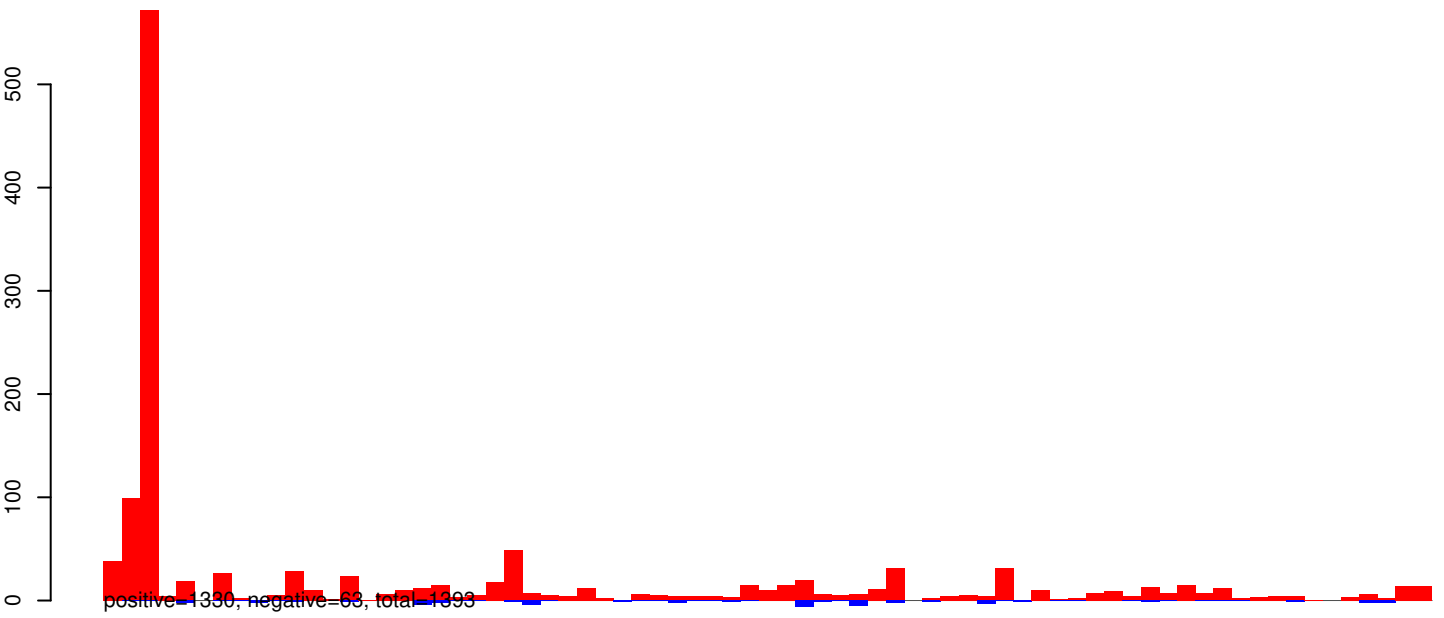
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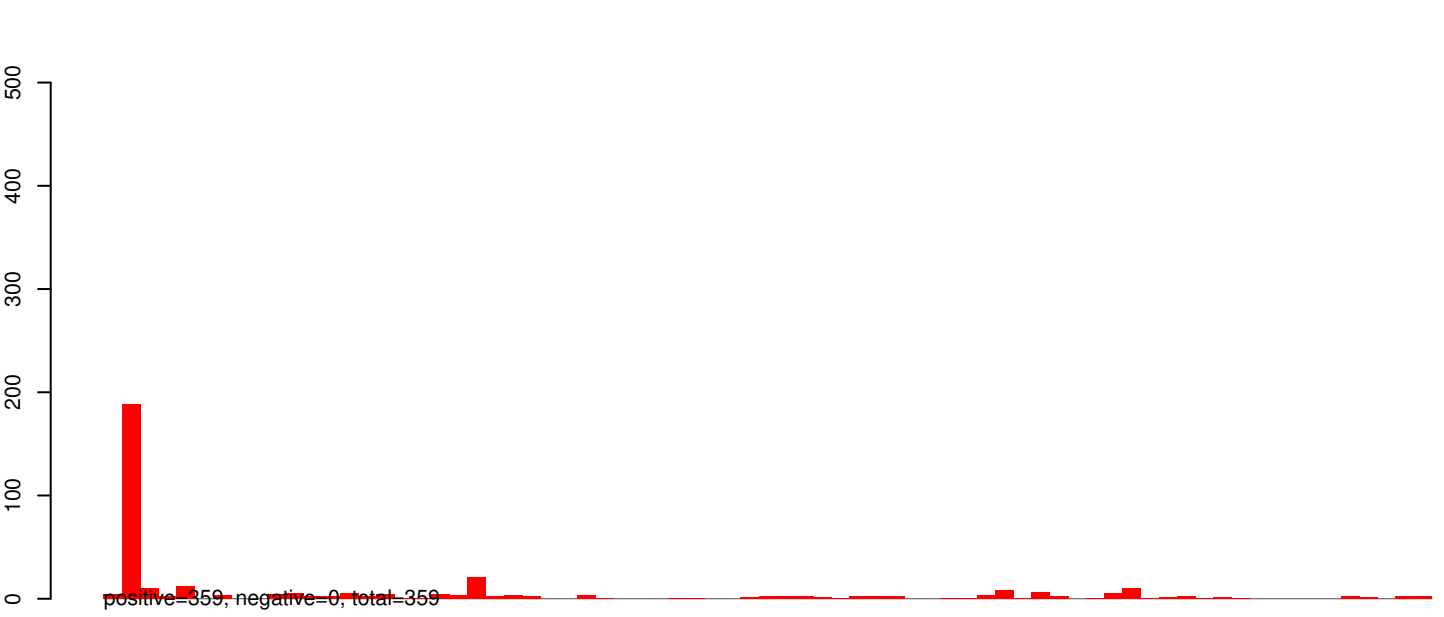
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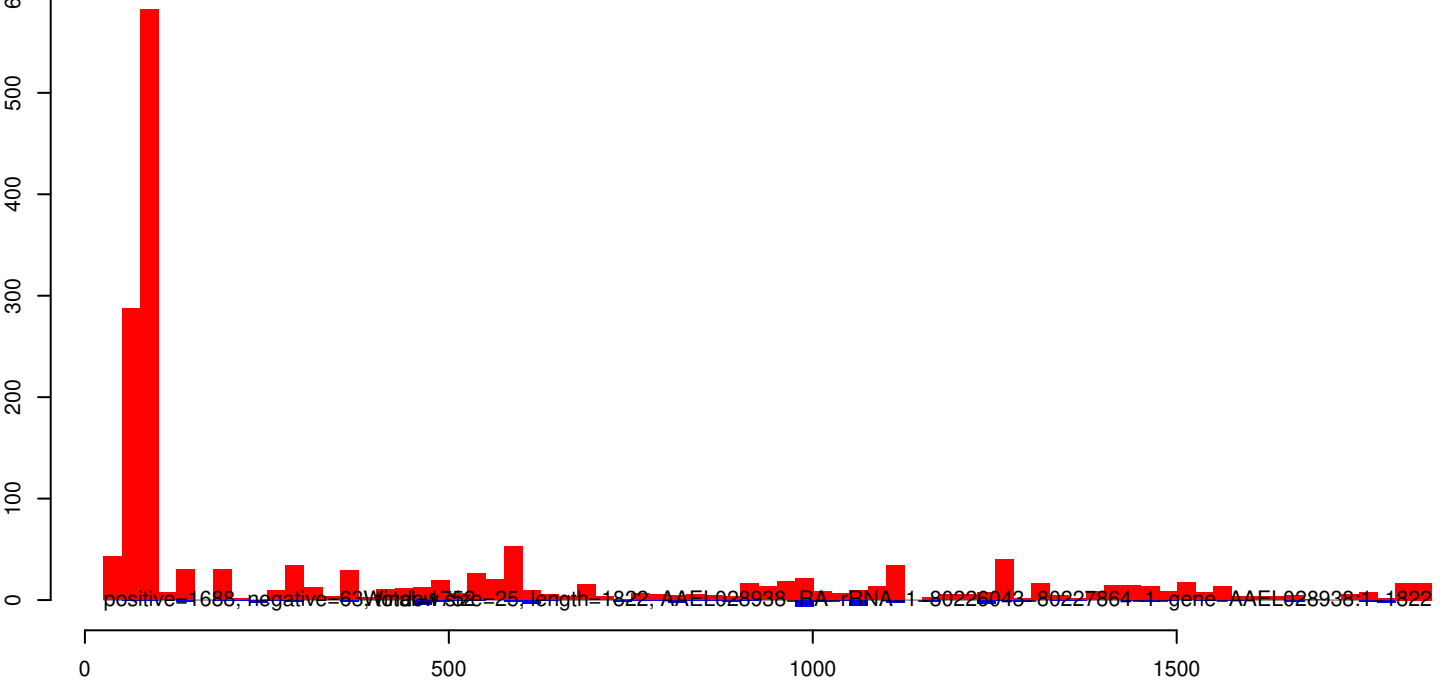
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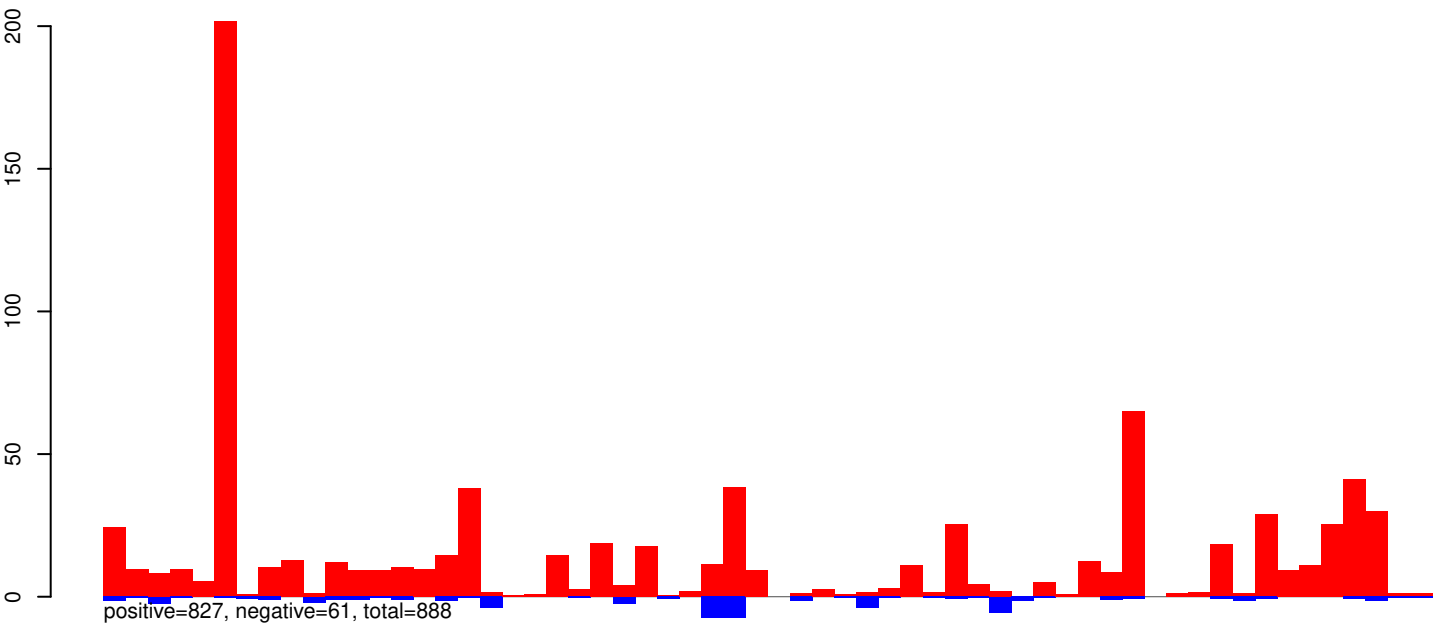
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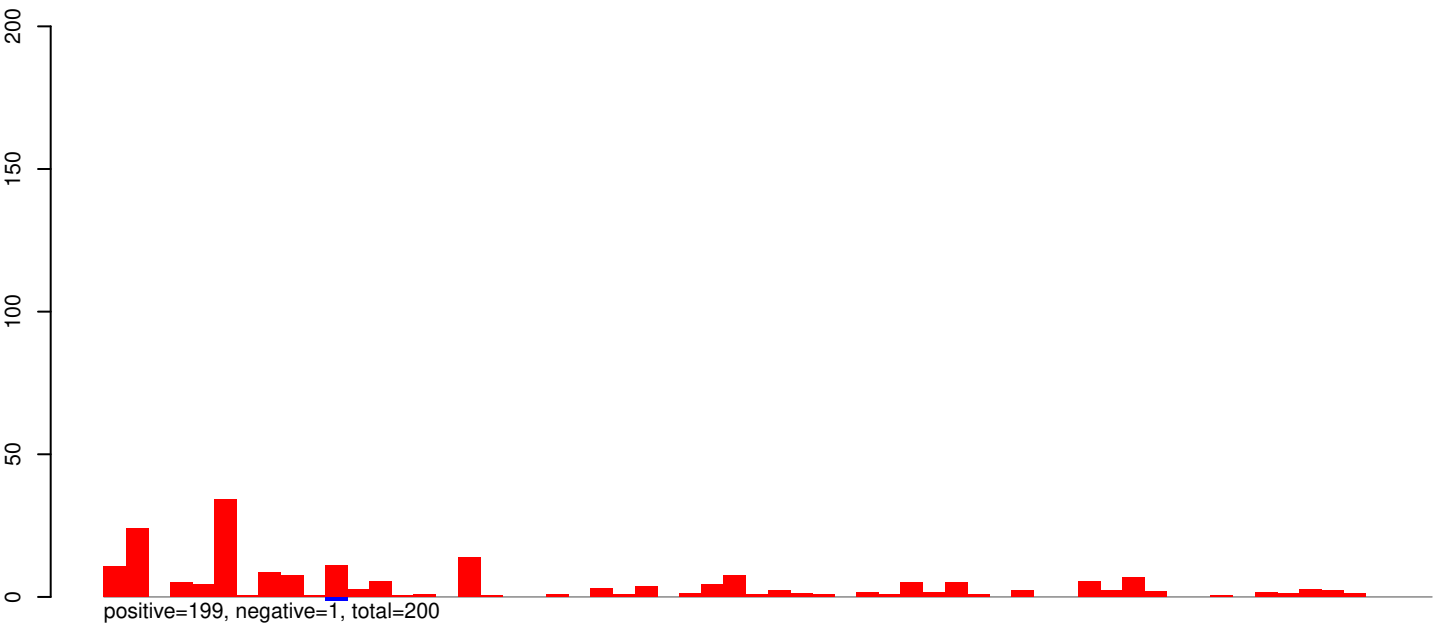
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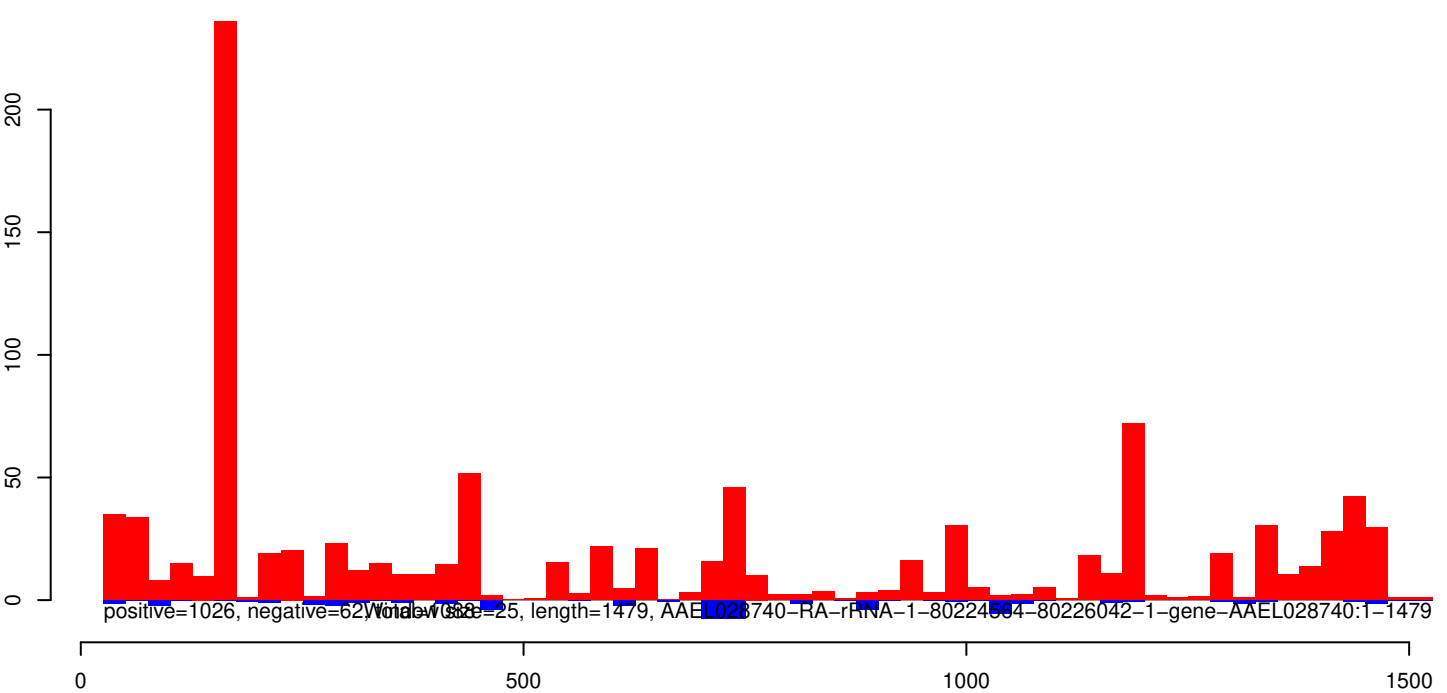
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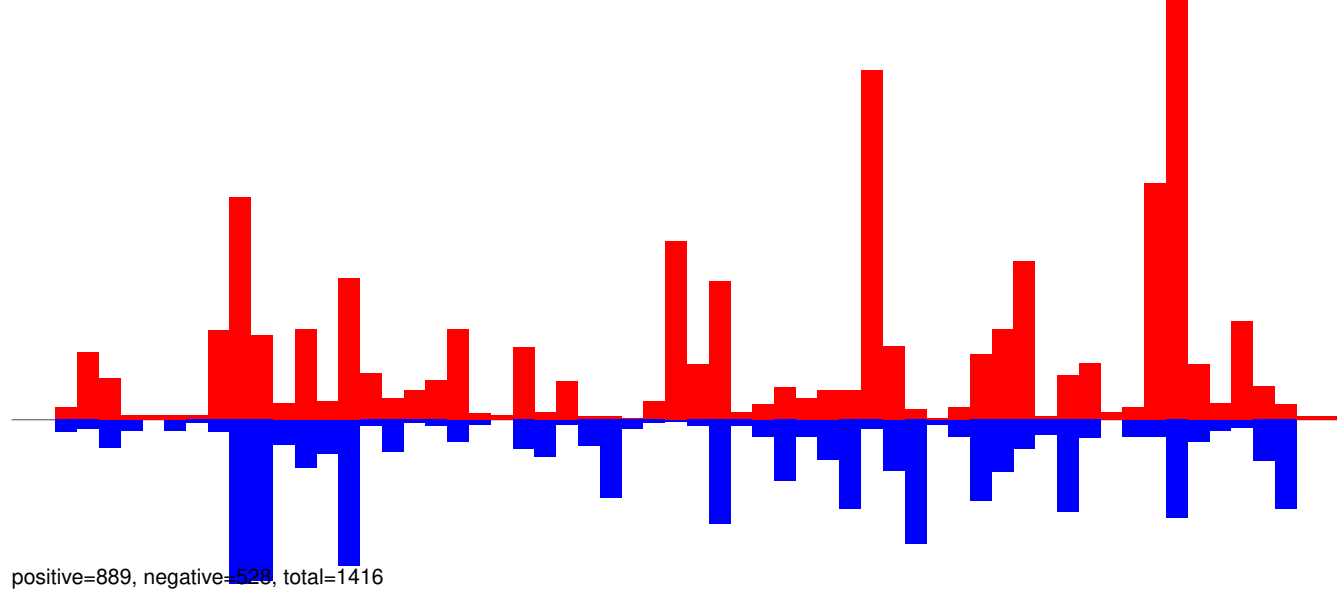
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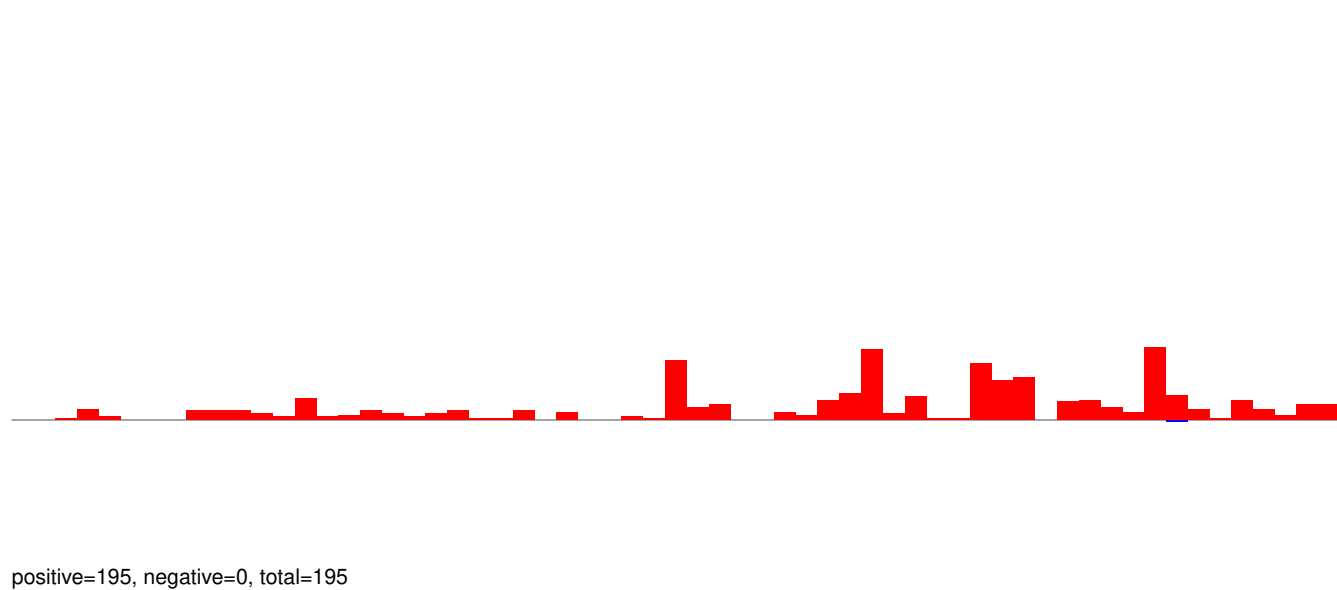
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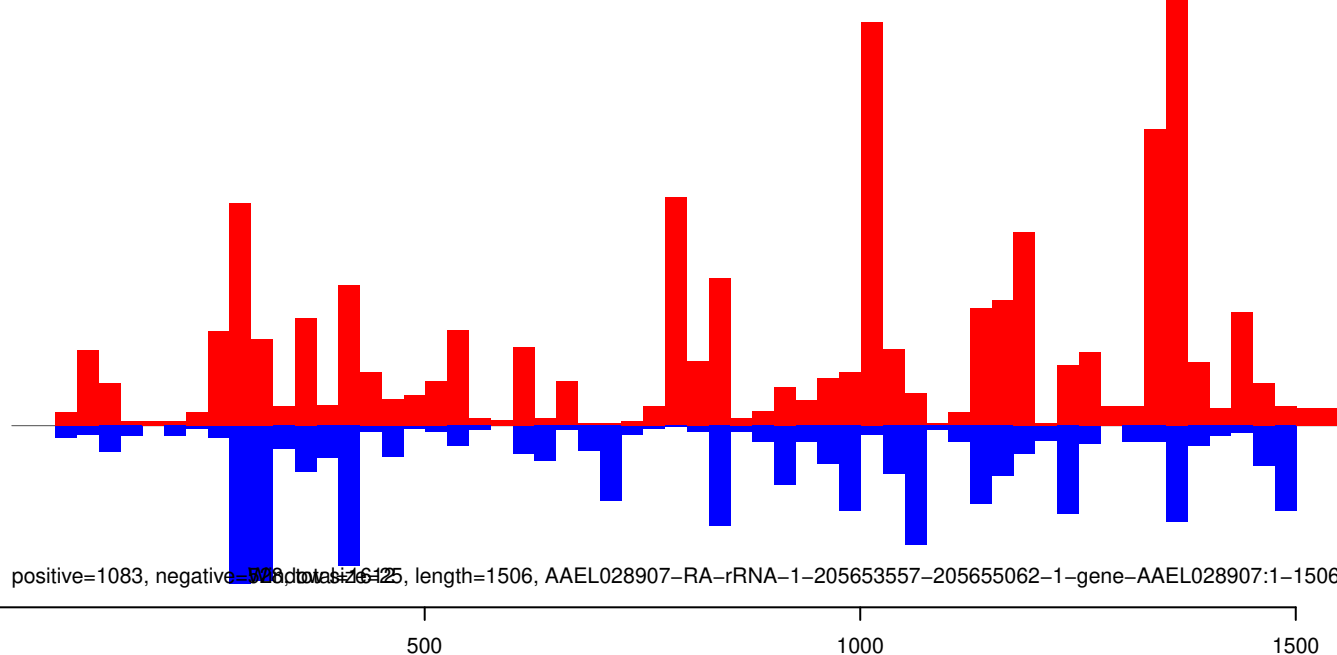
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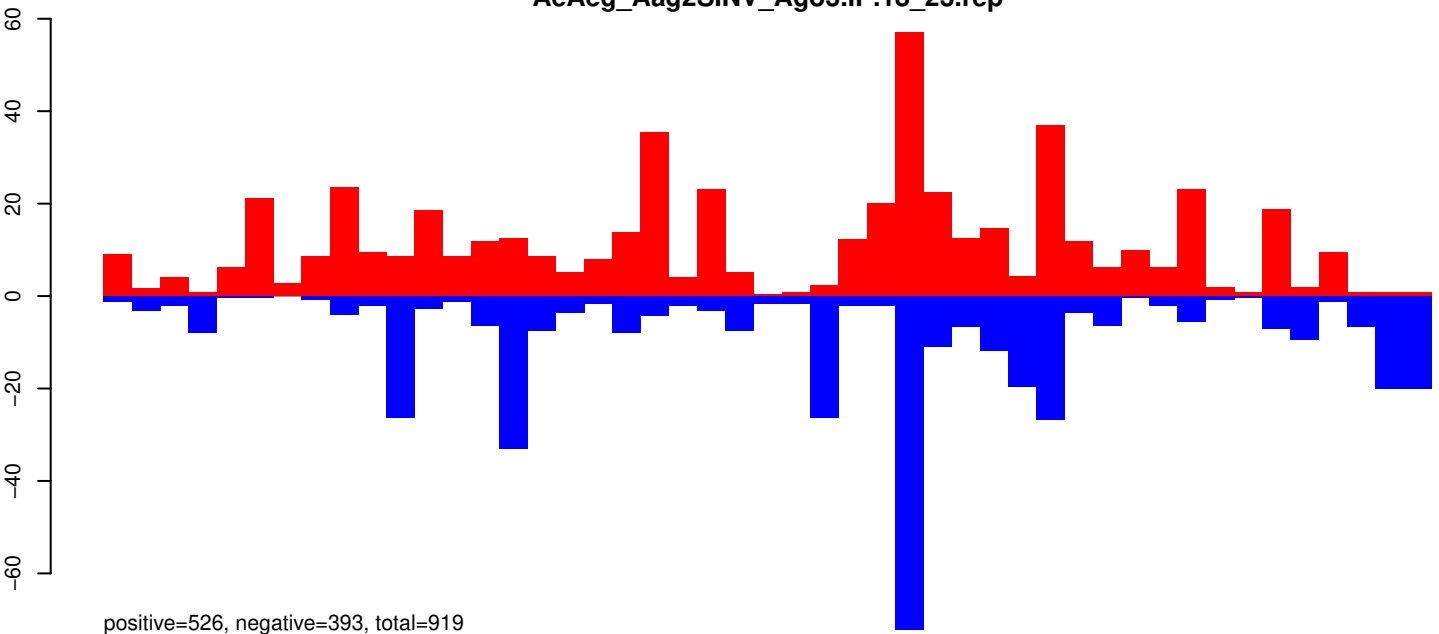
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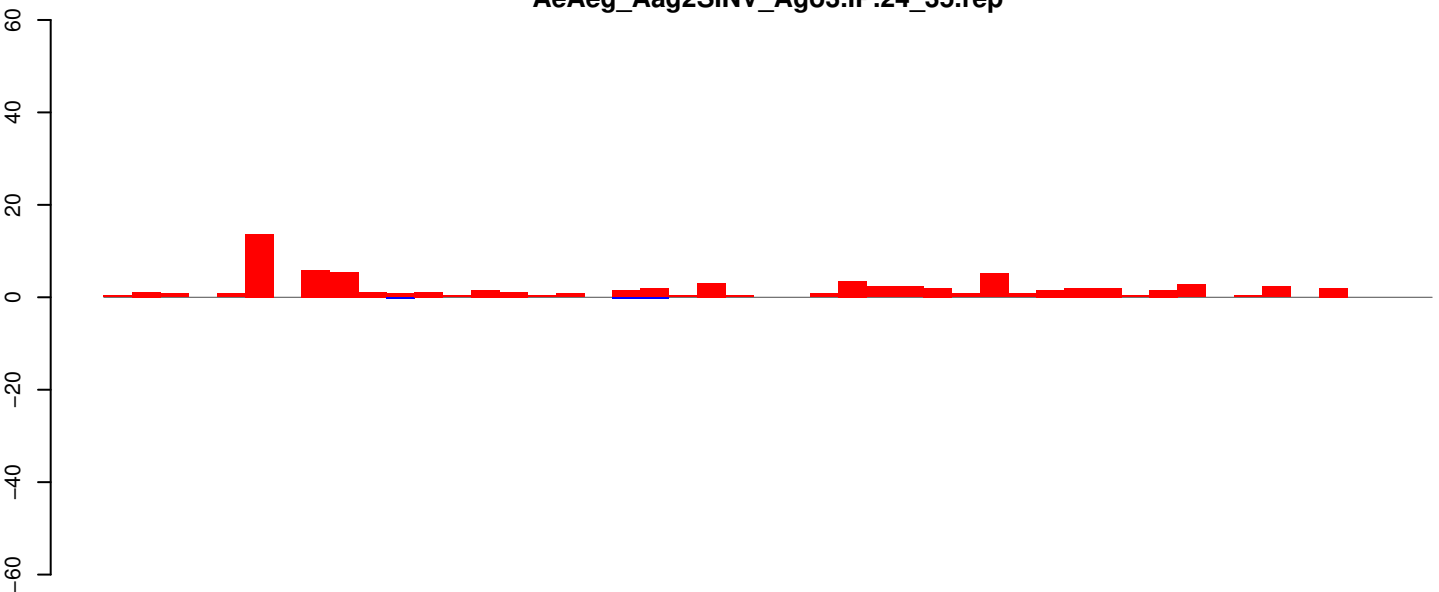


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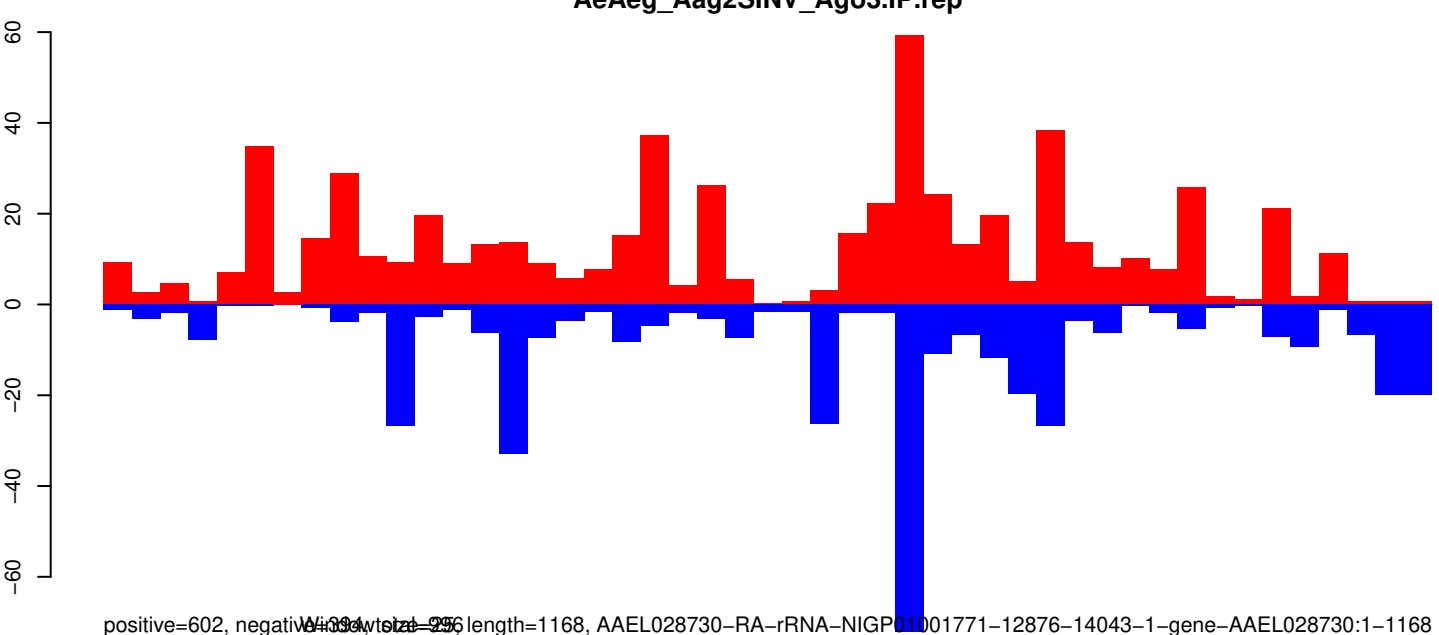
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AeAeg_Aag2SINV_Ago3.IP.24_35.rep



positive=76, negative=1, total=77

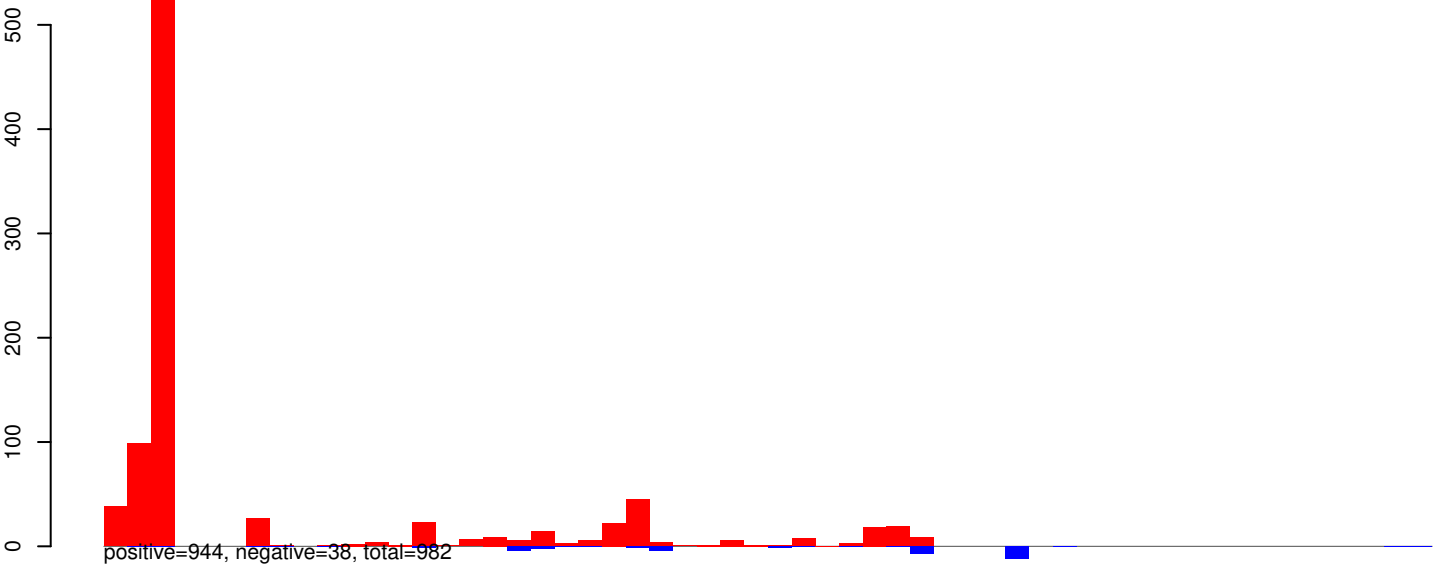
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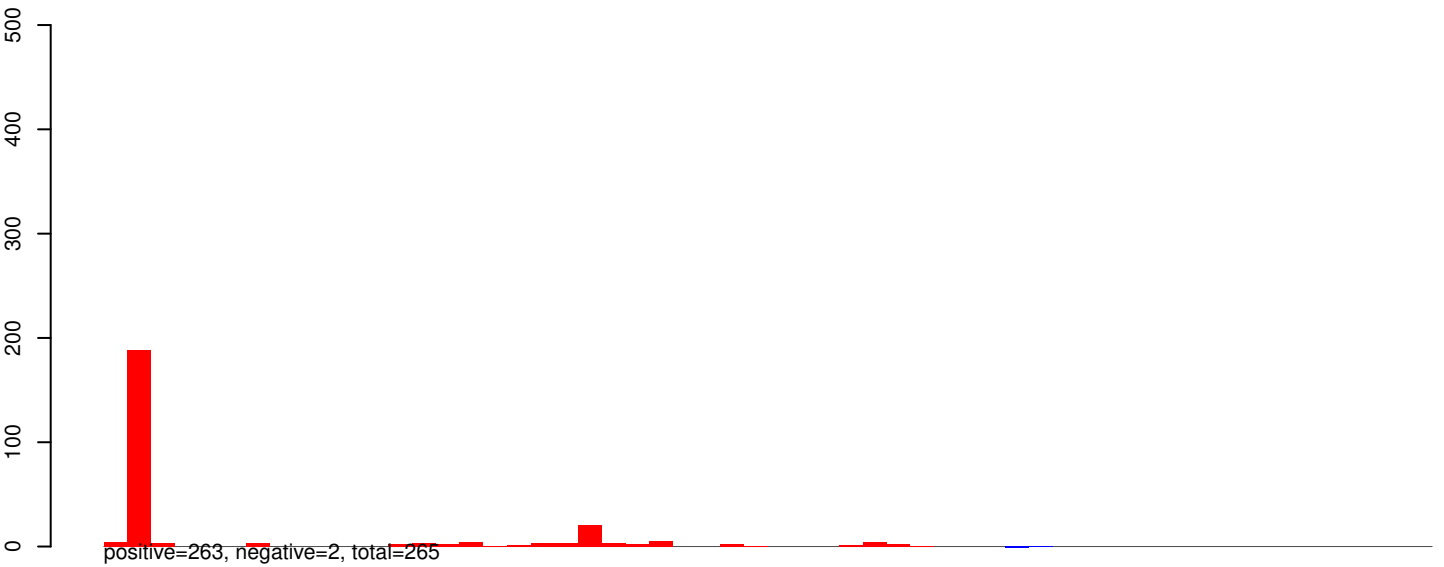
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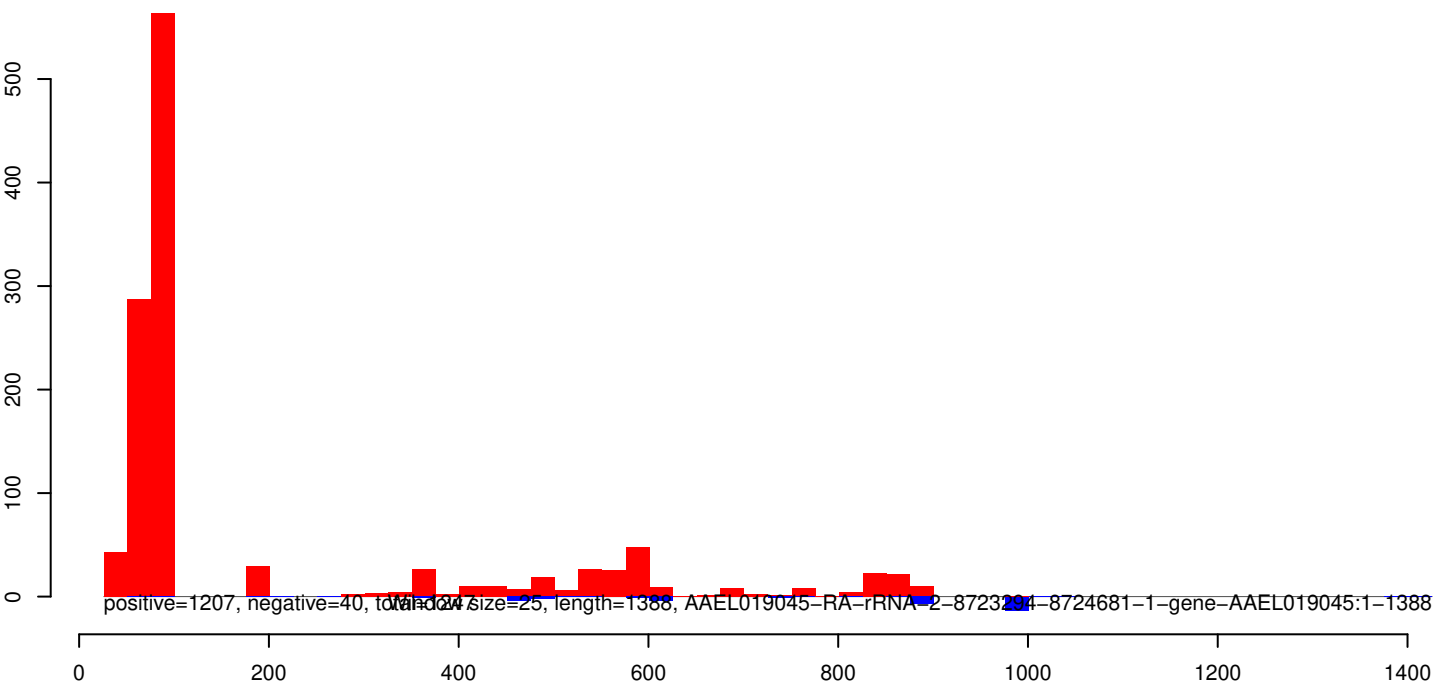
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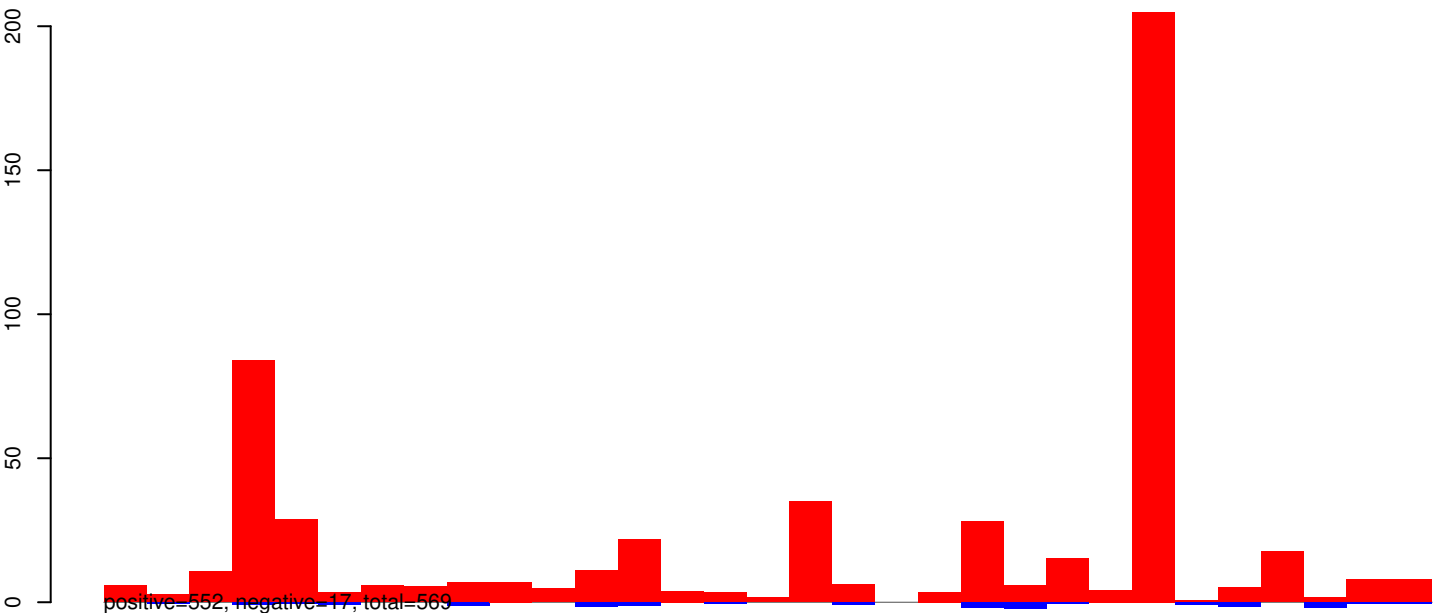
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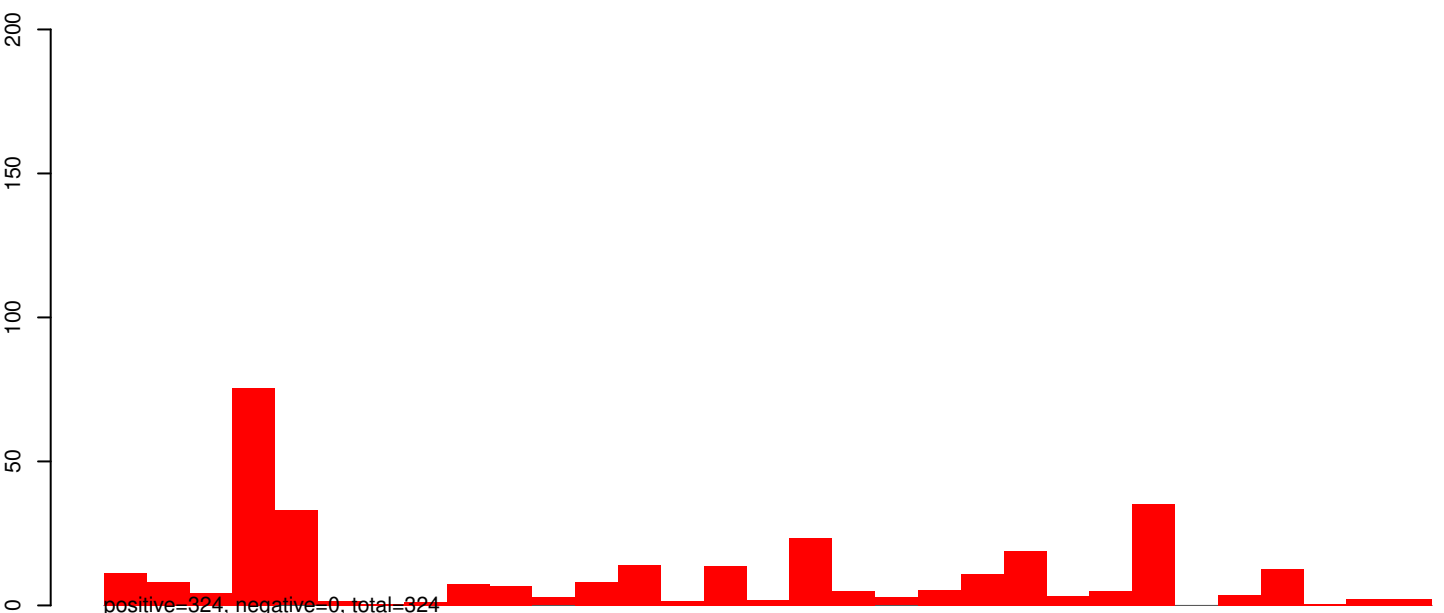
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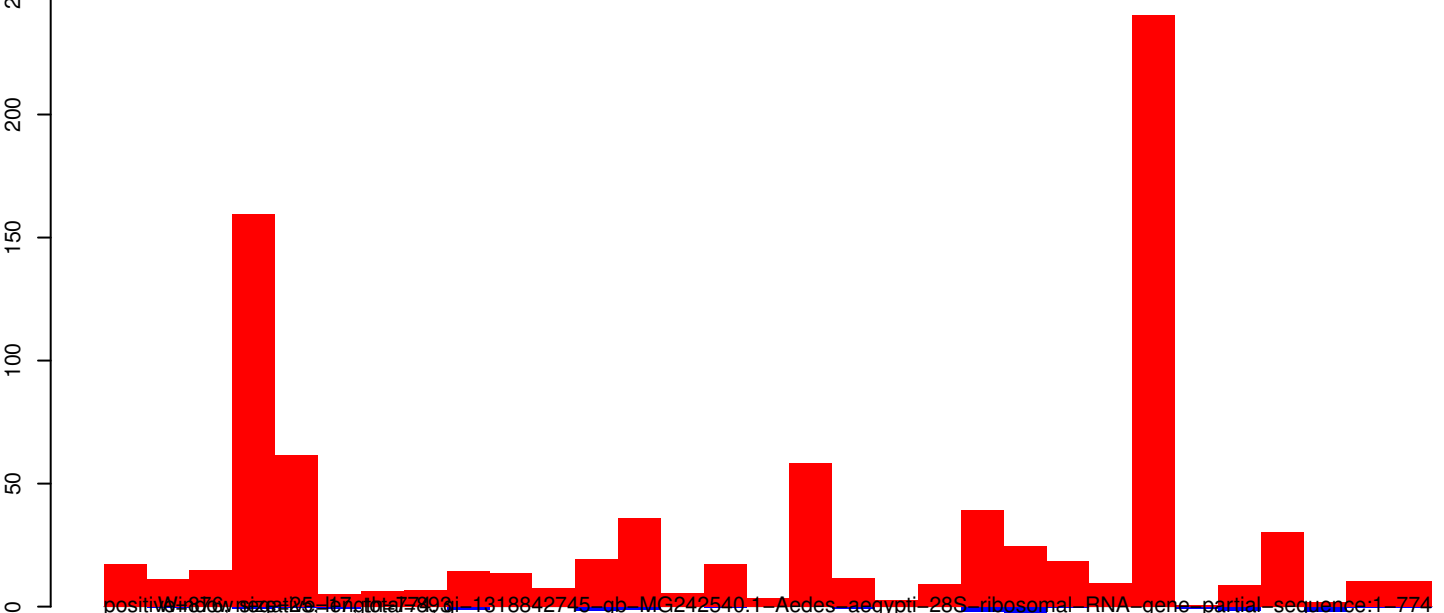
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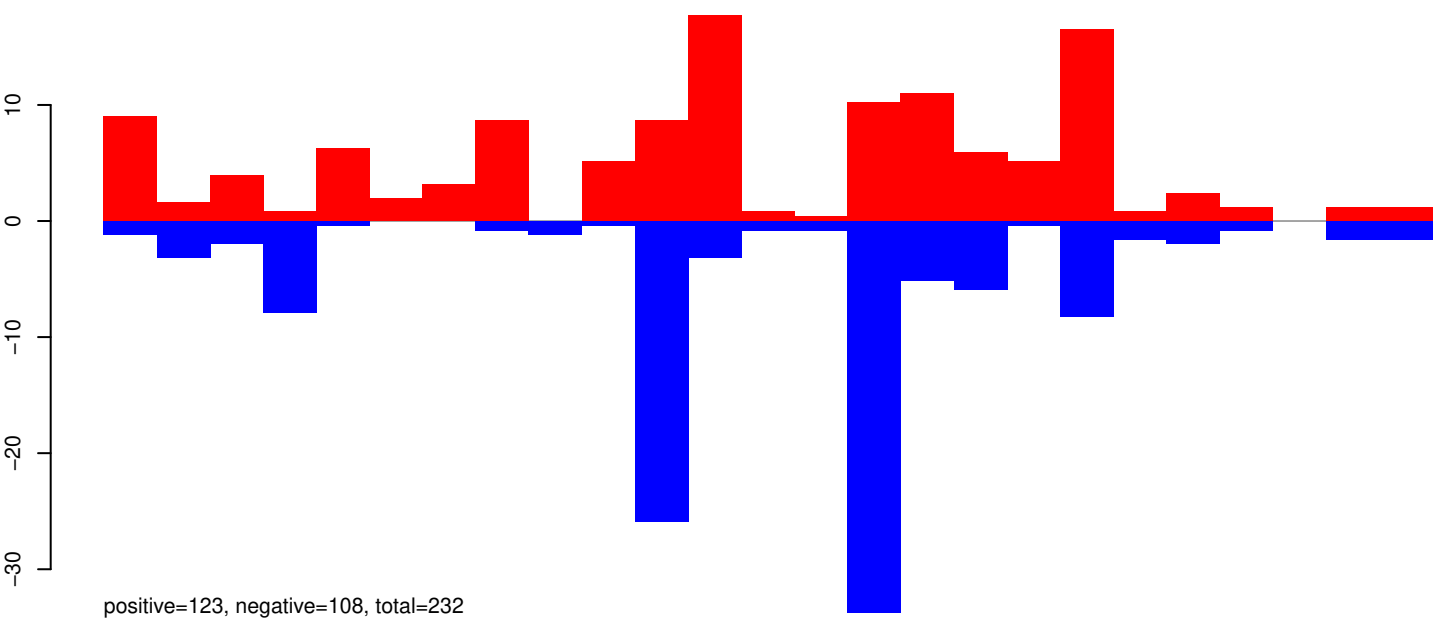
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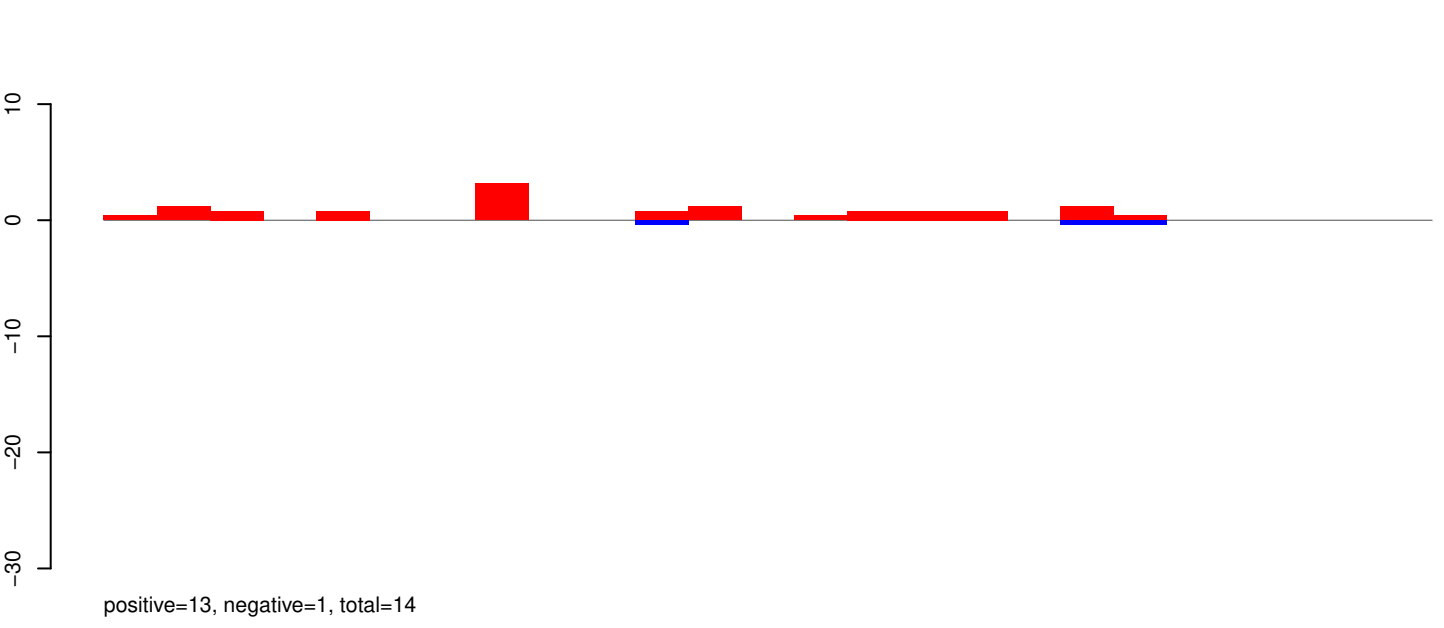
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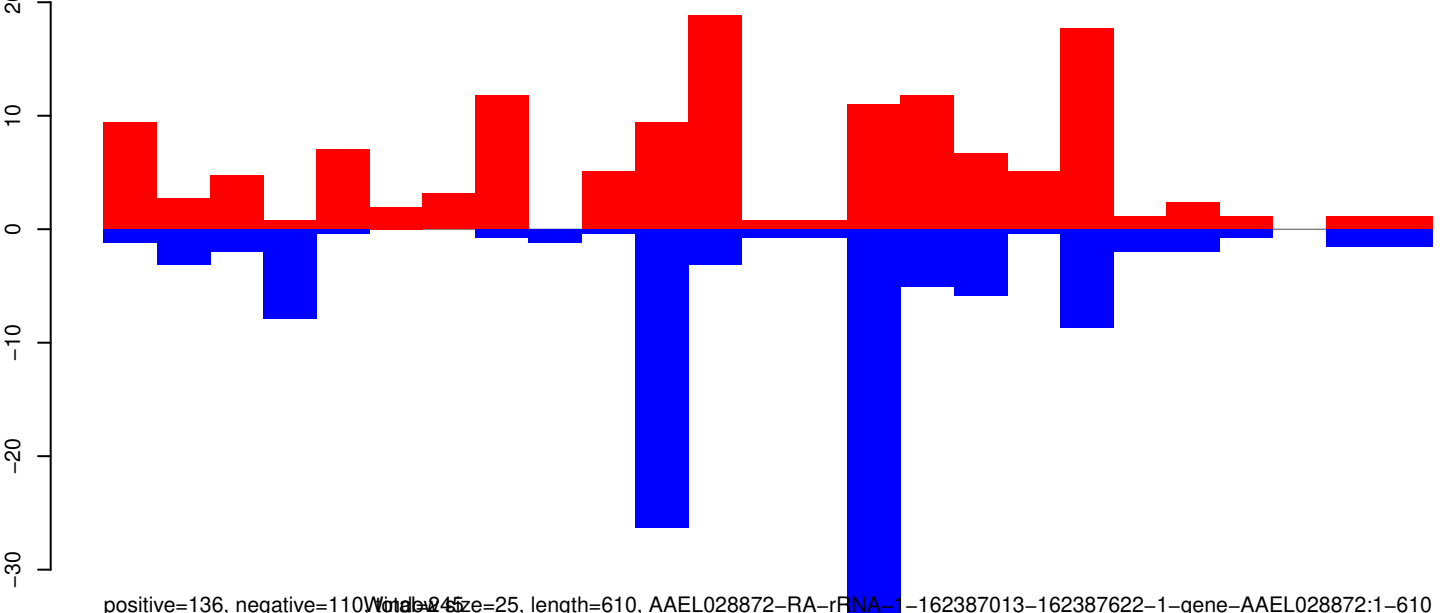
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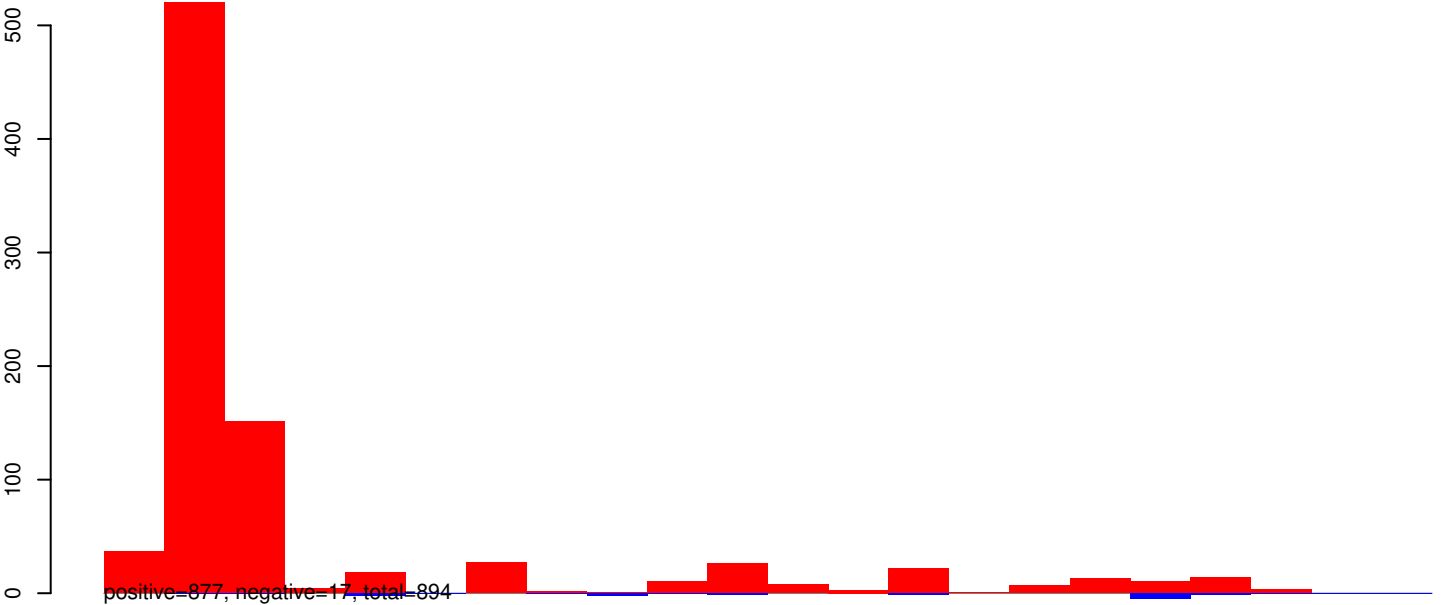
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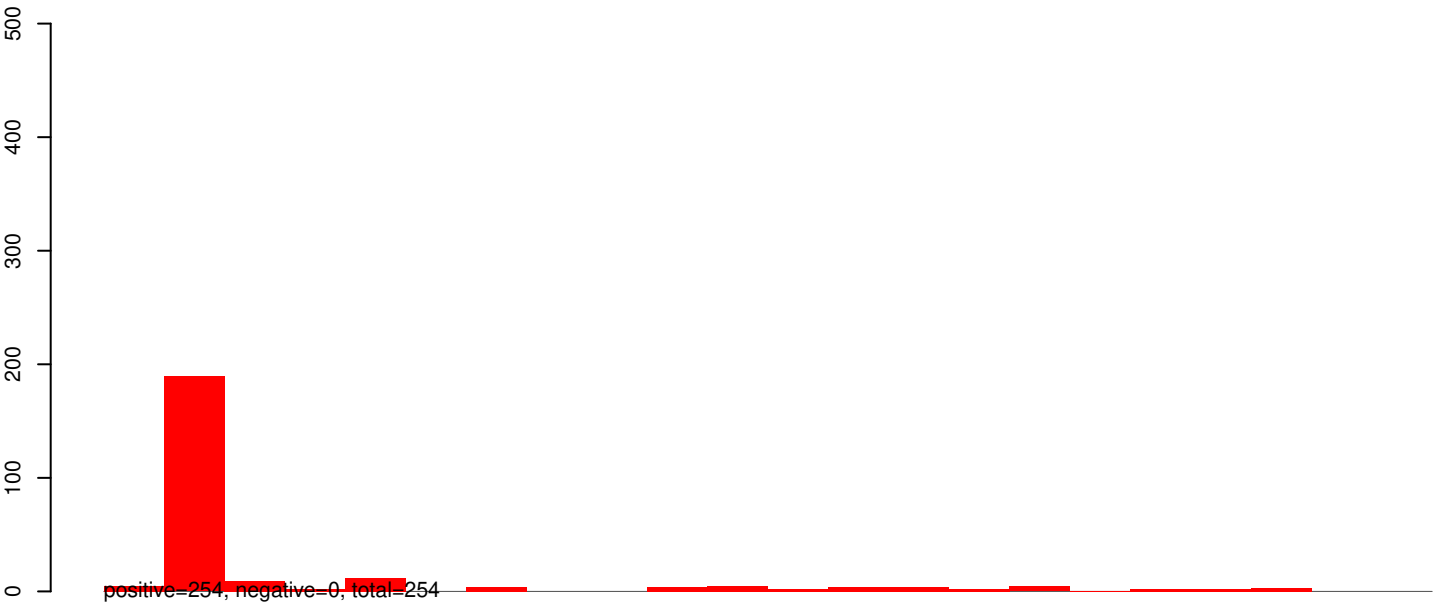
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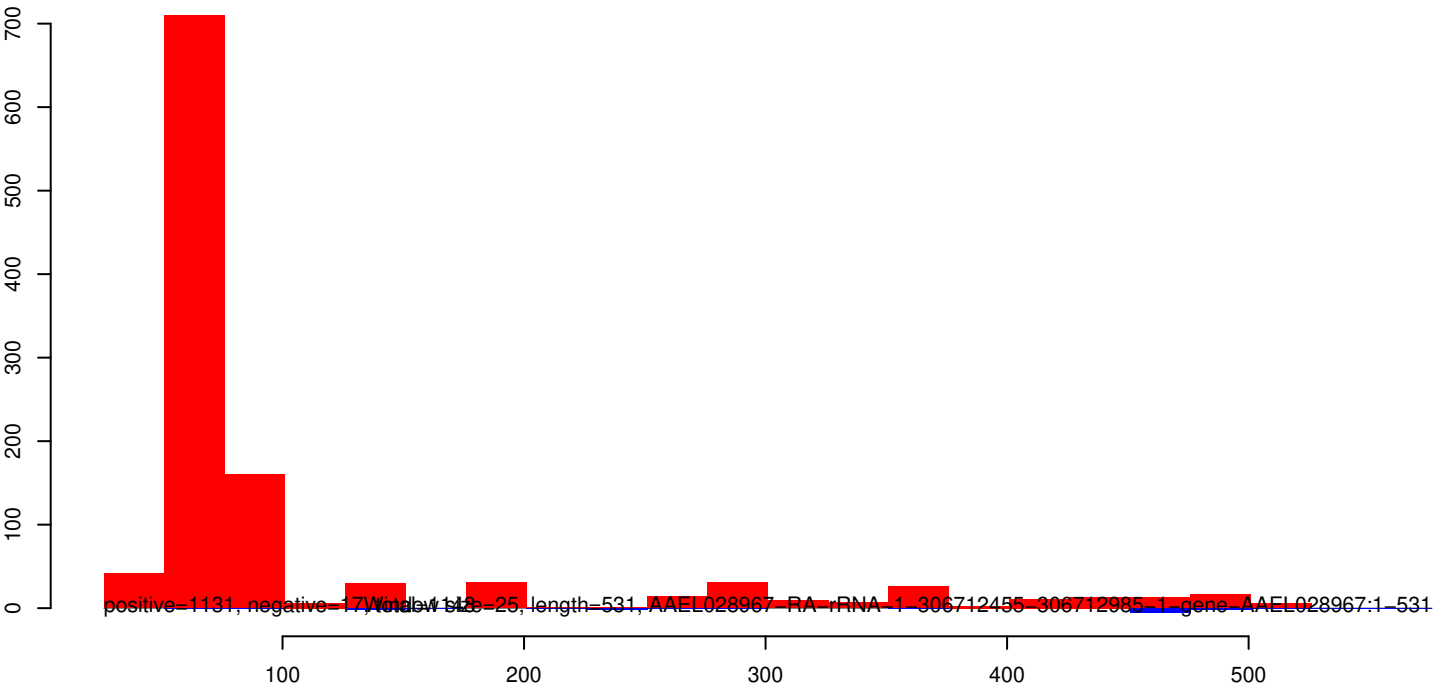
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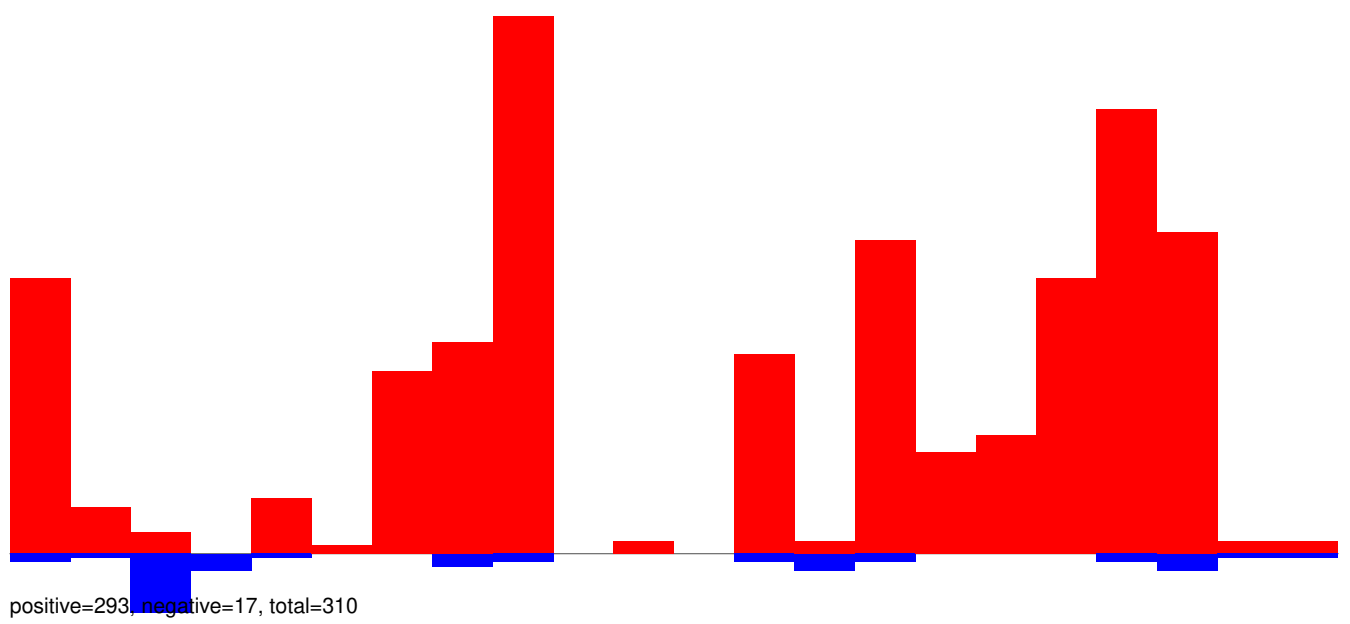
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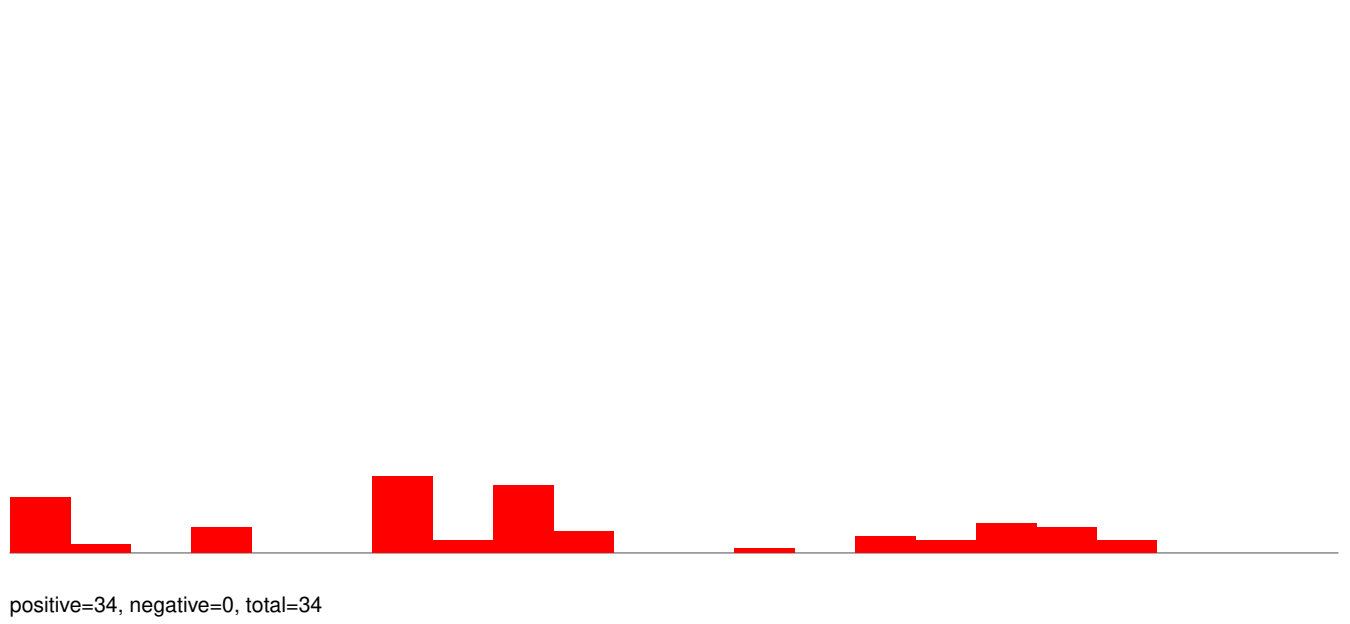
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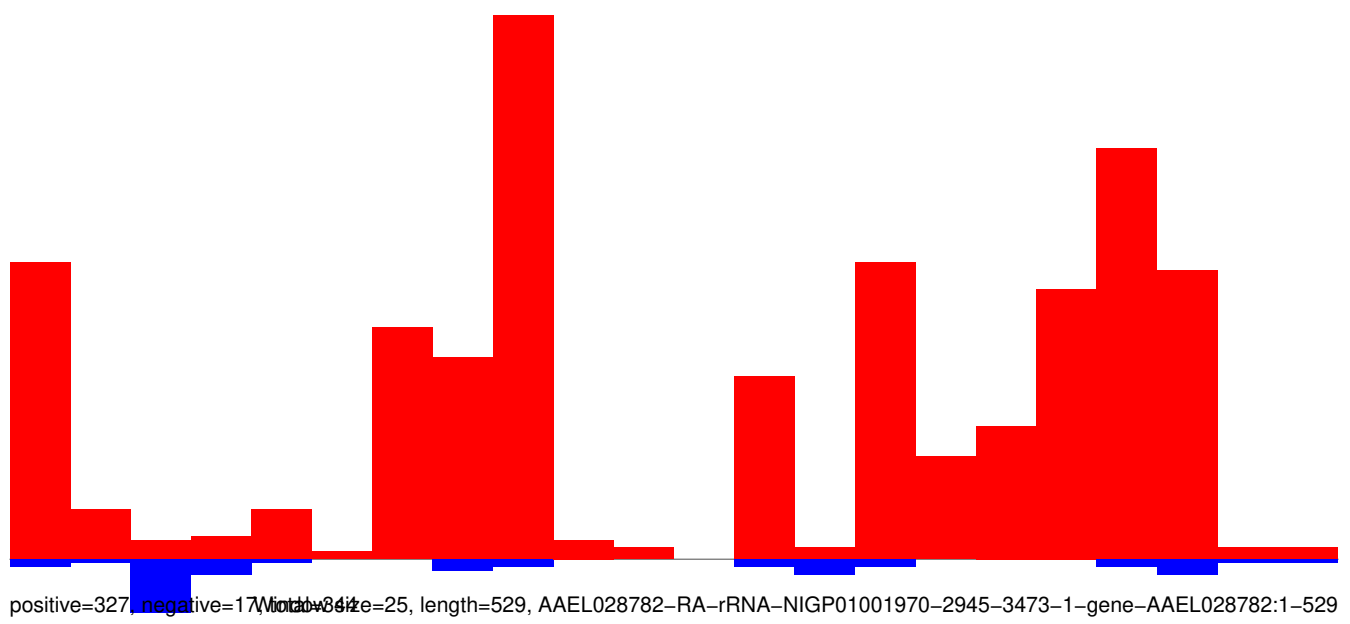
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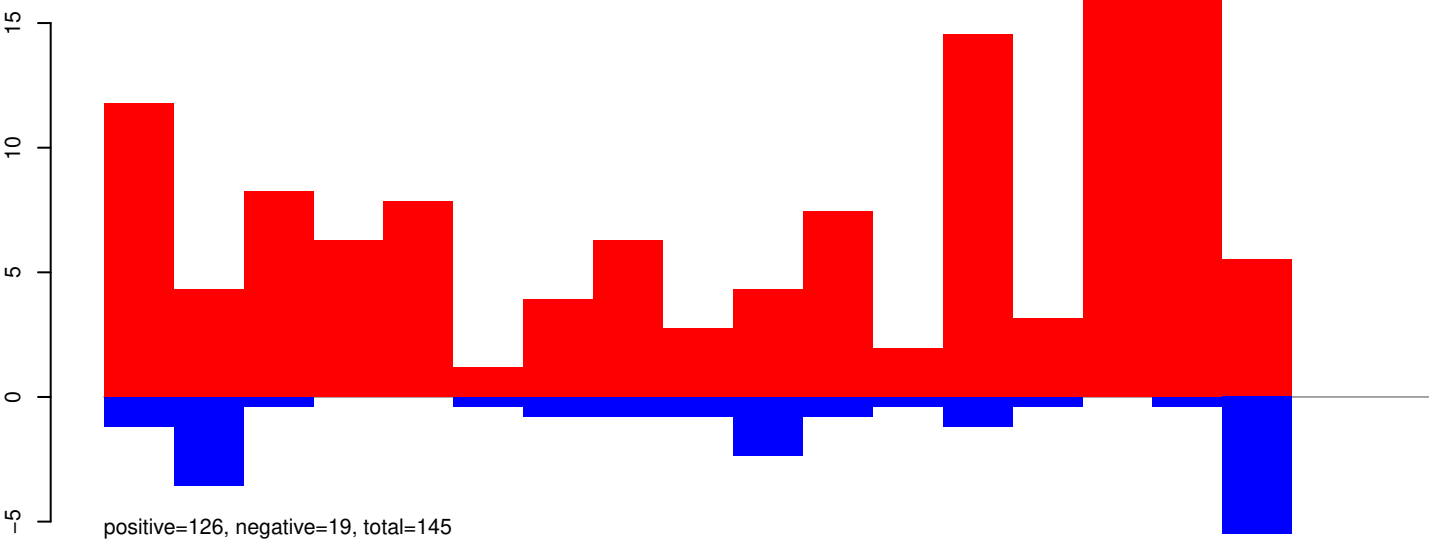
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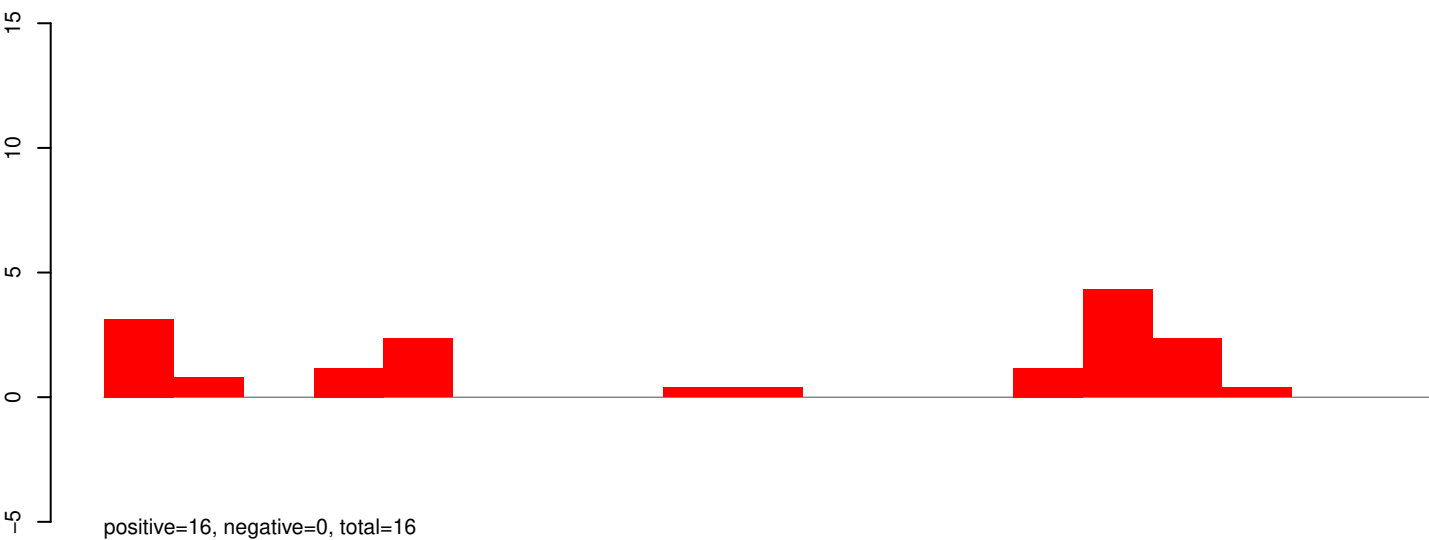
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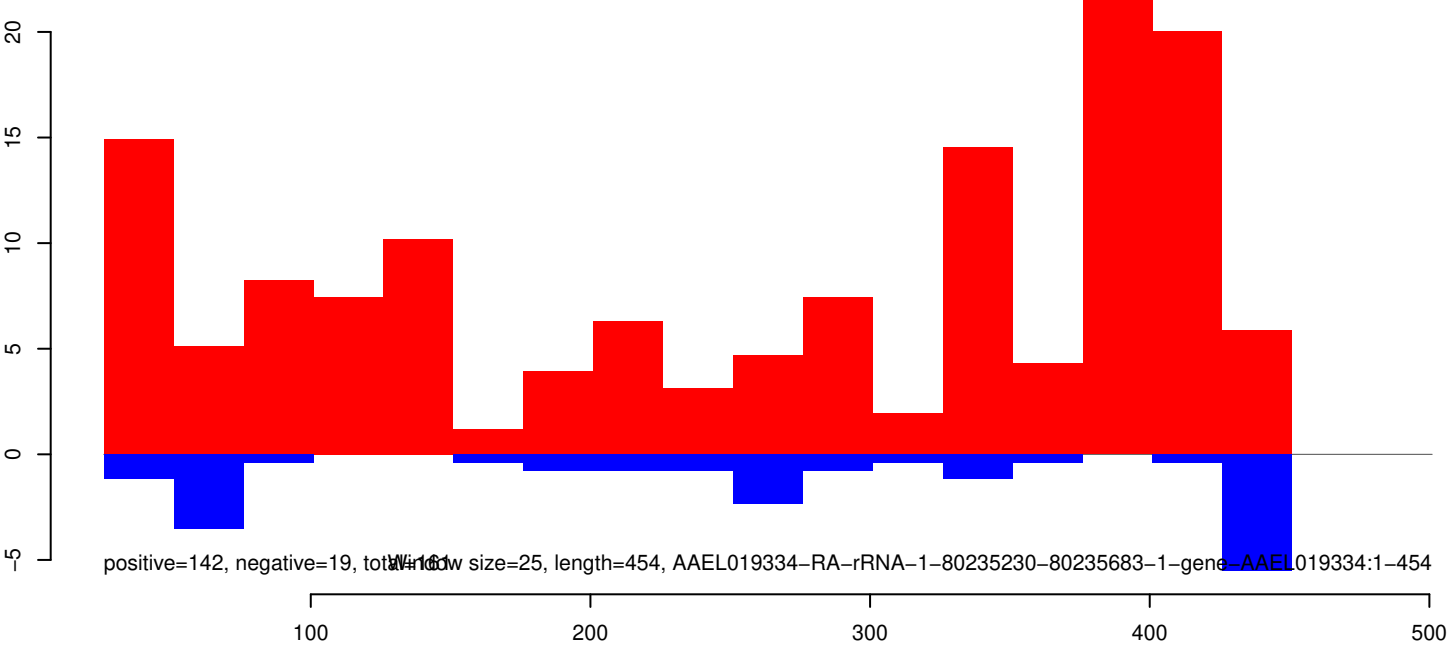
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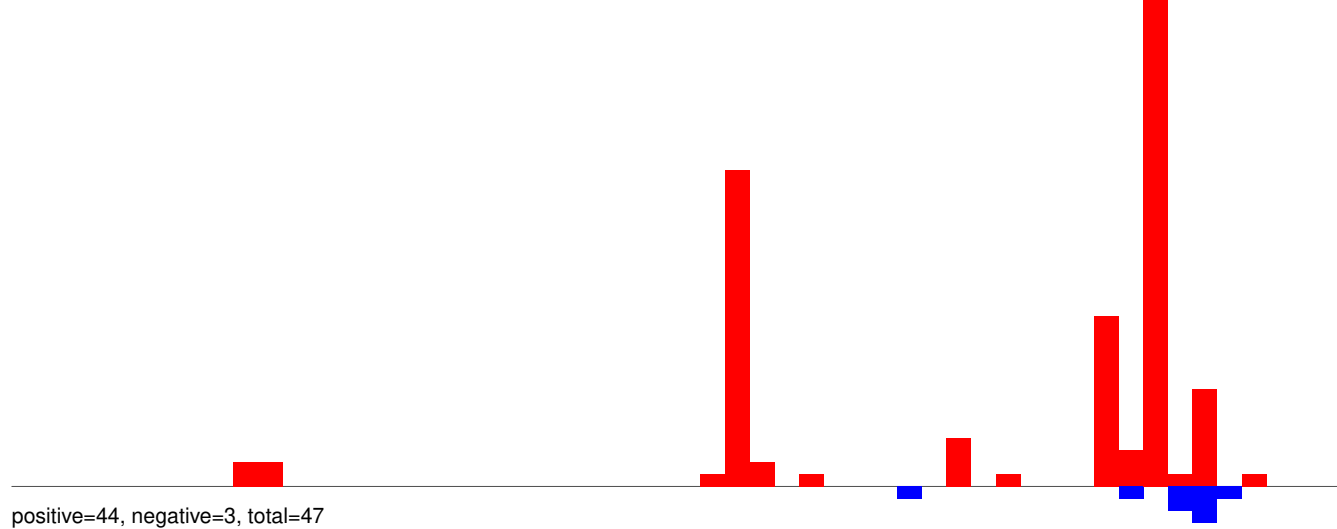
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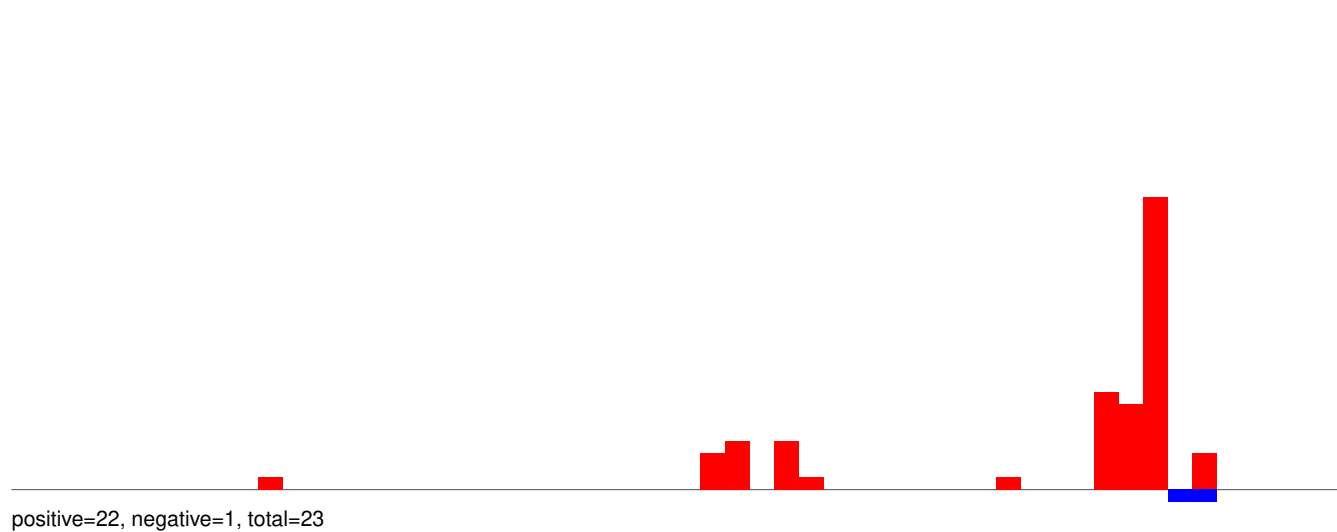
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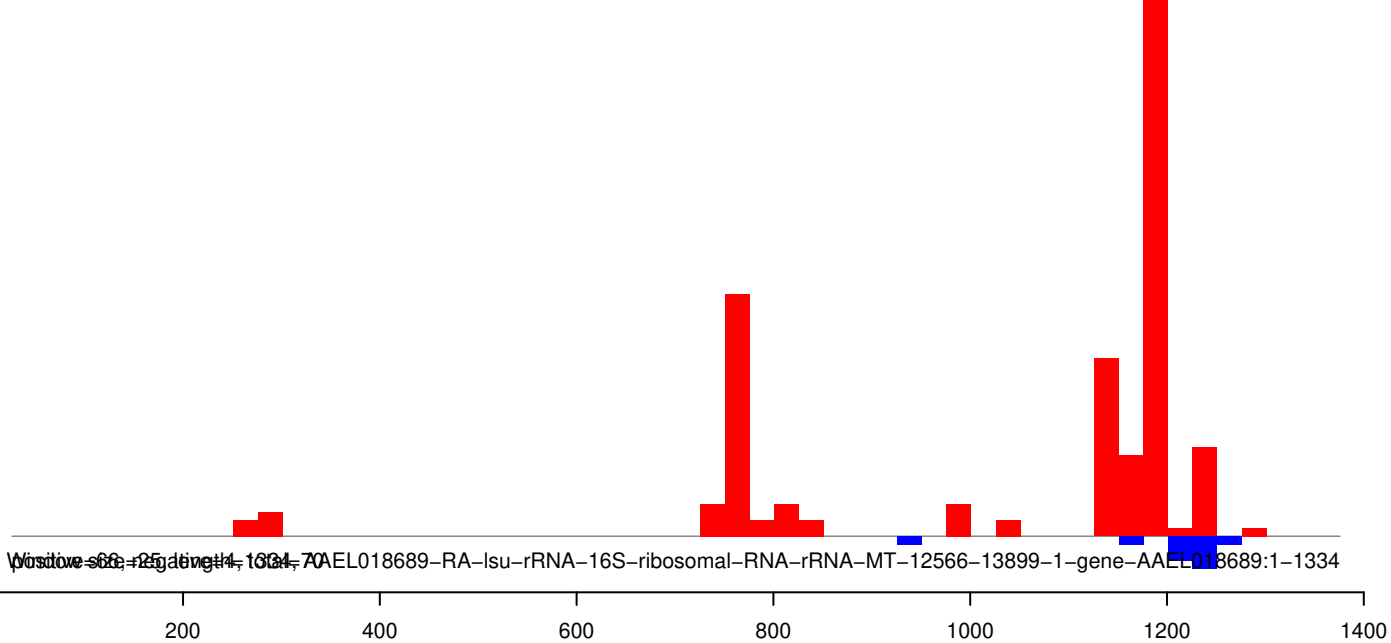
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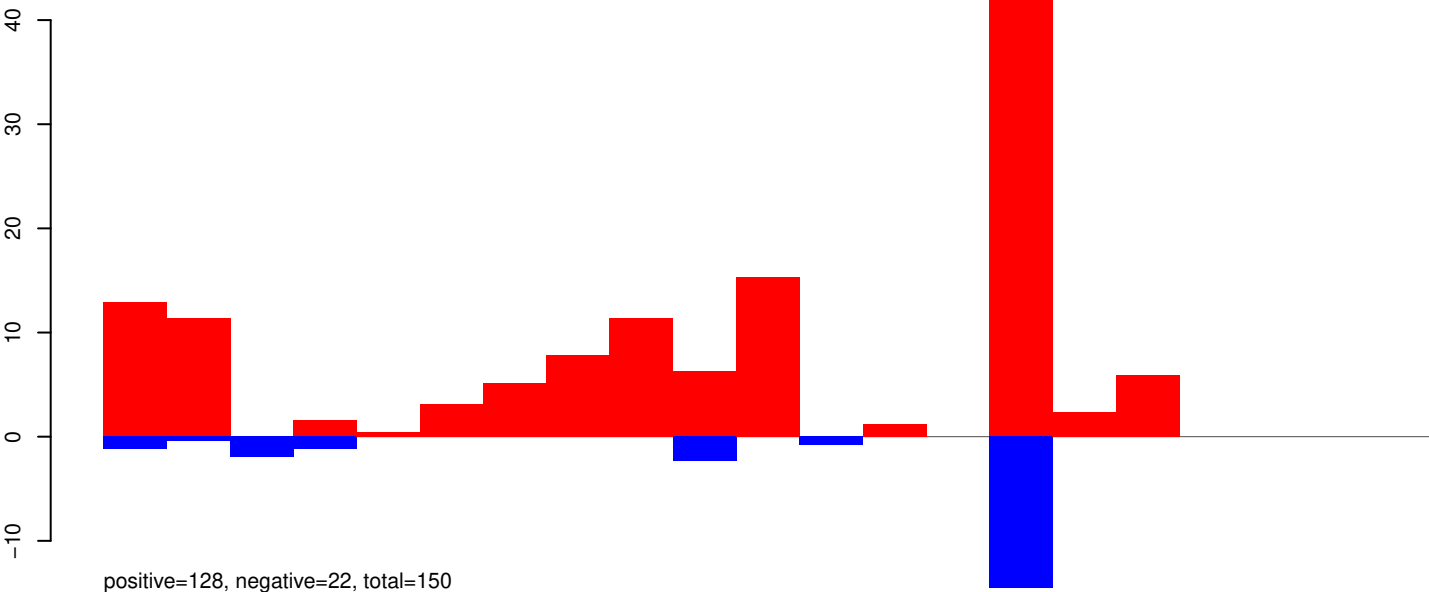
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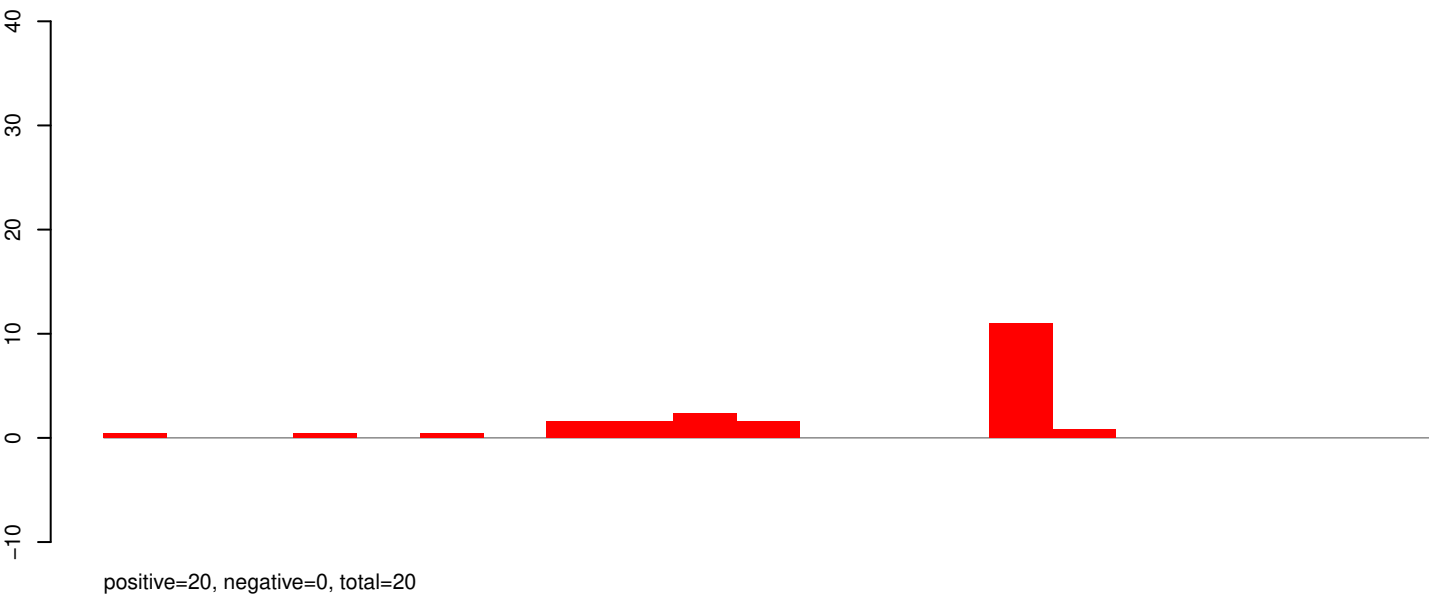
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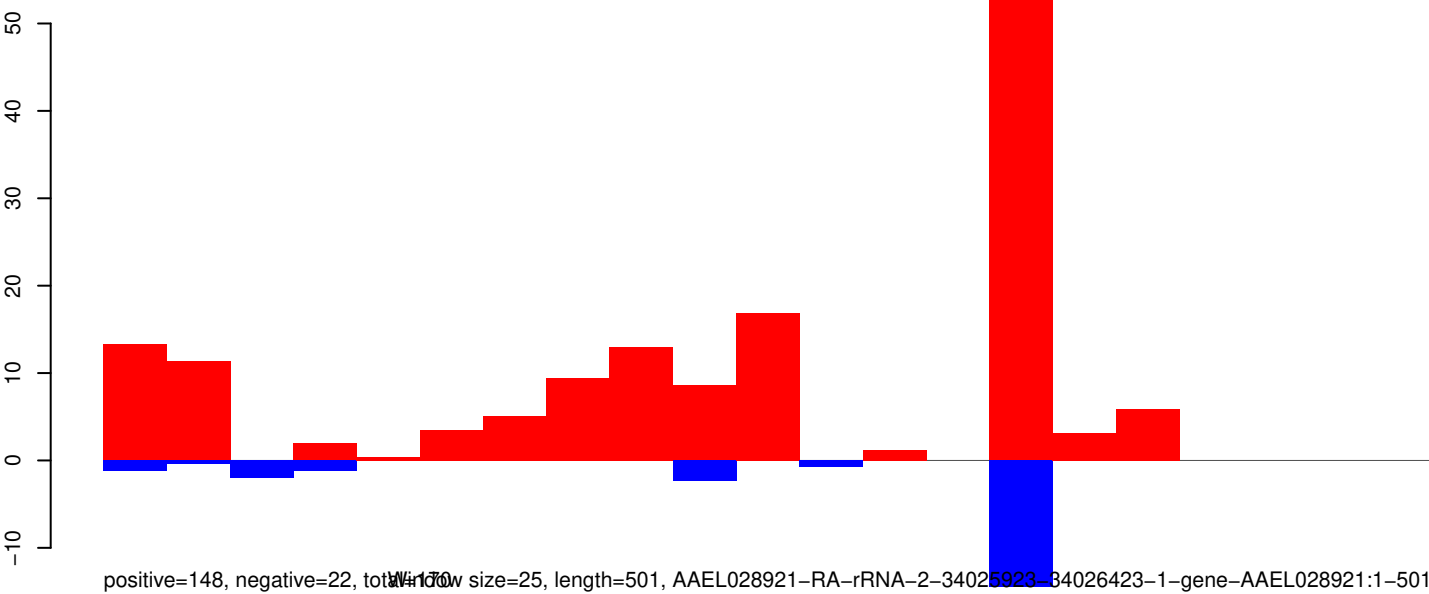
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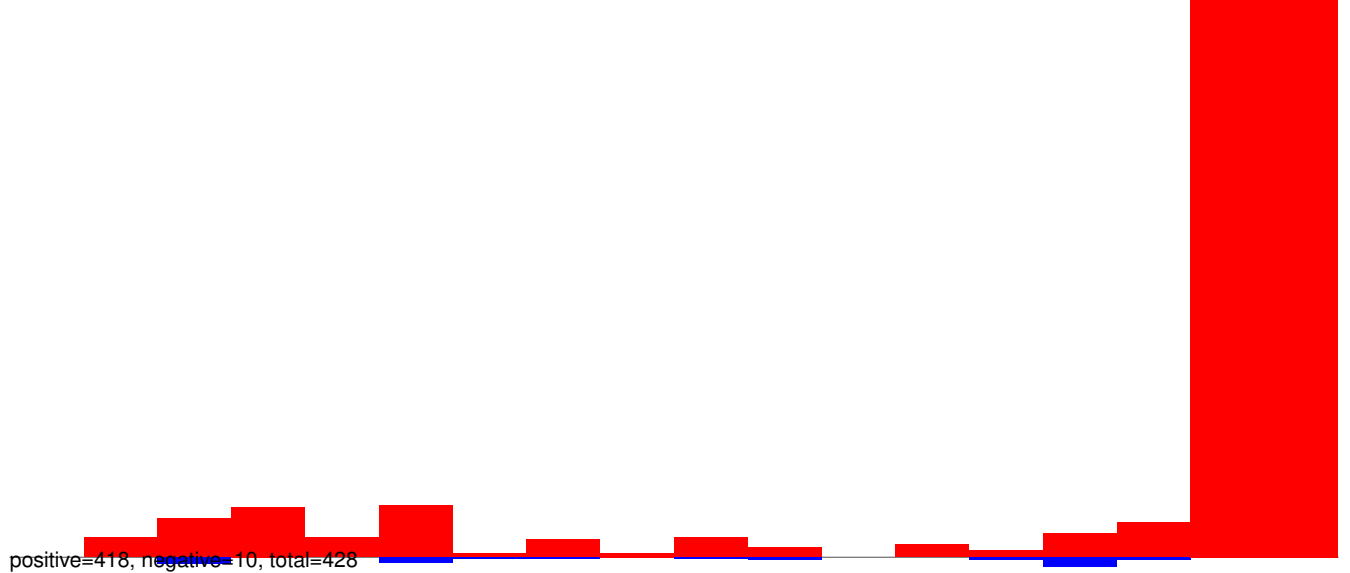


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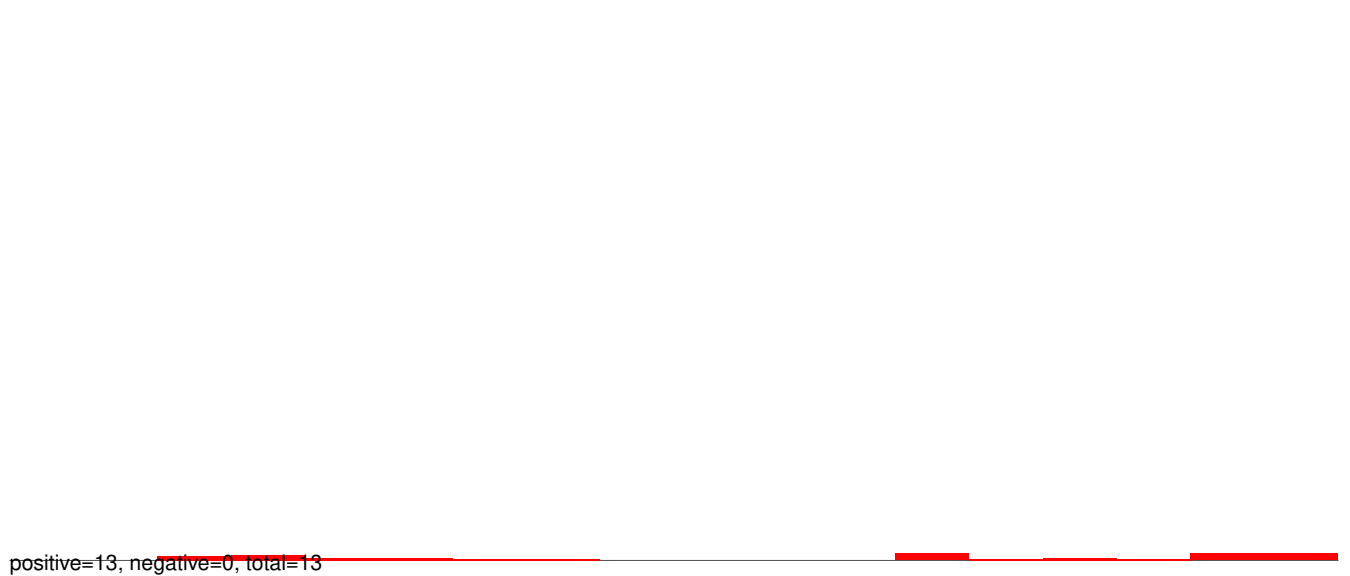


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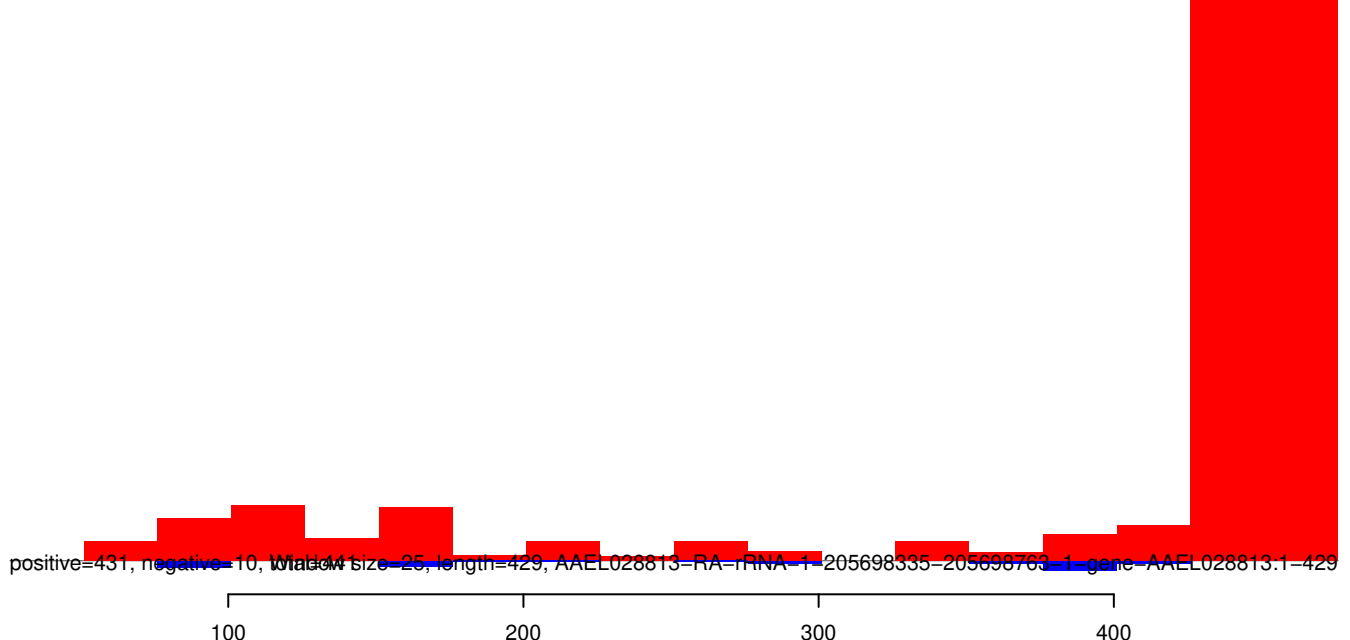
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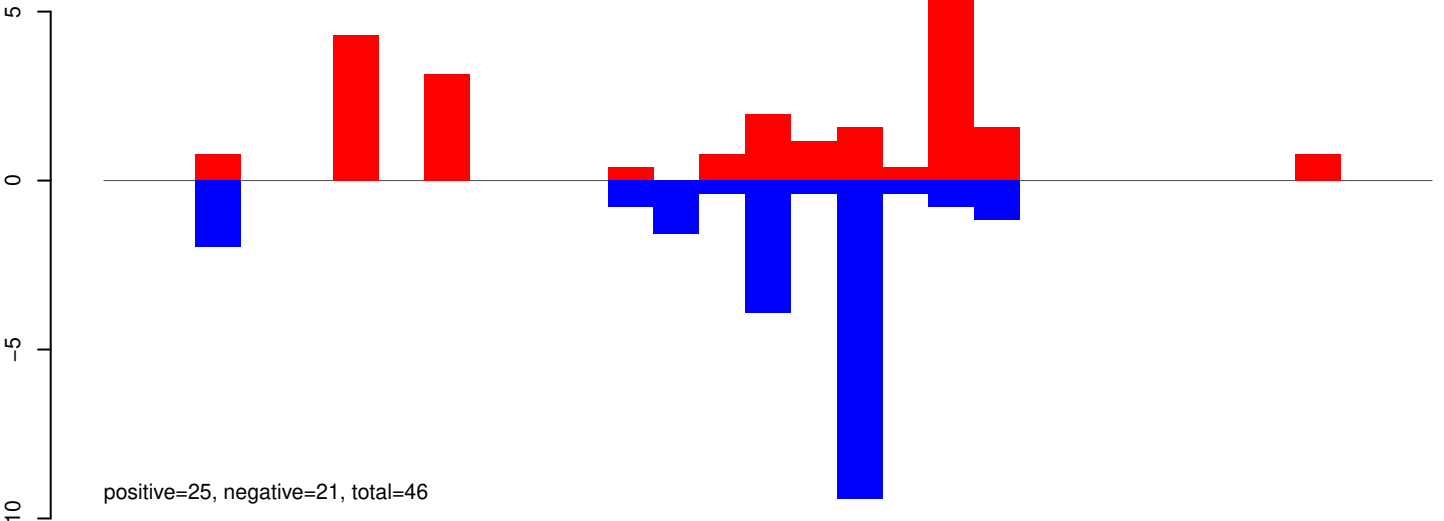
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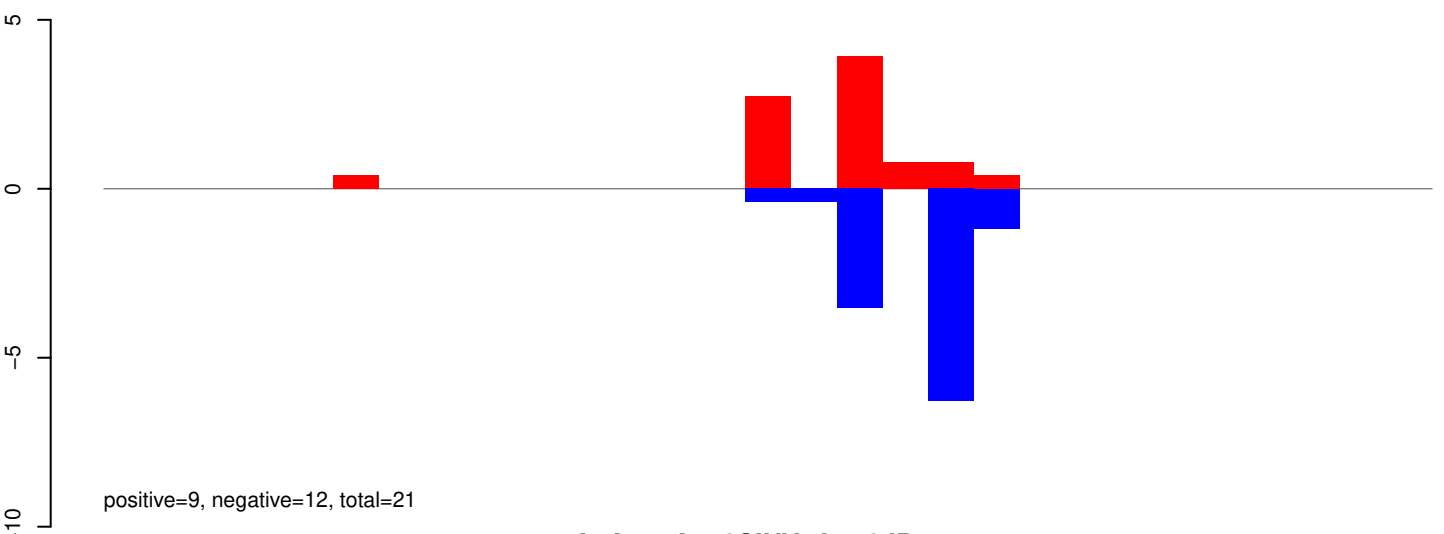
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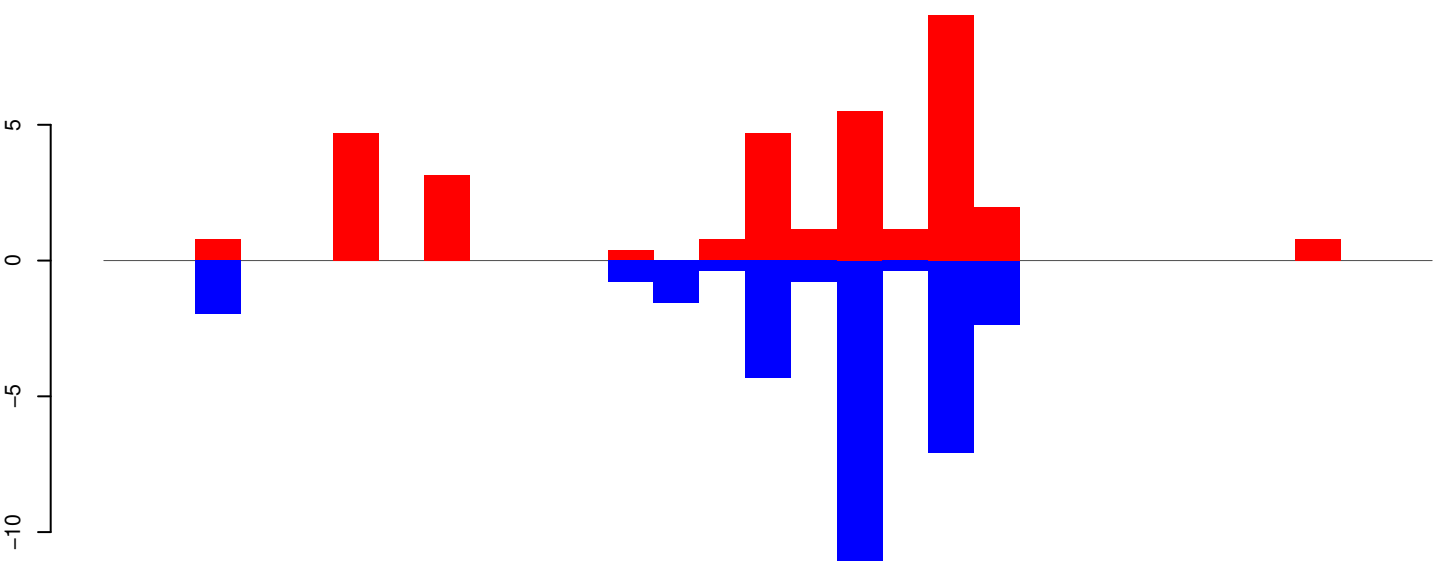
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AeAeg_Aag2SINV_Ago3.IP.24_35.rep

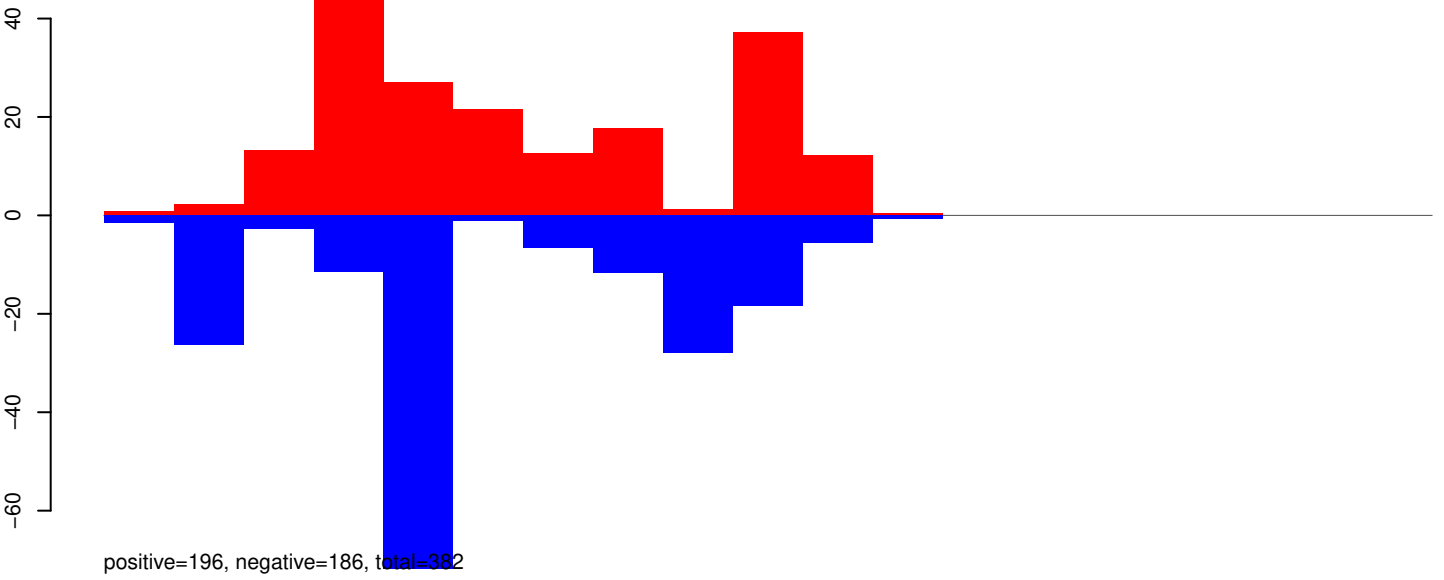


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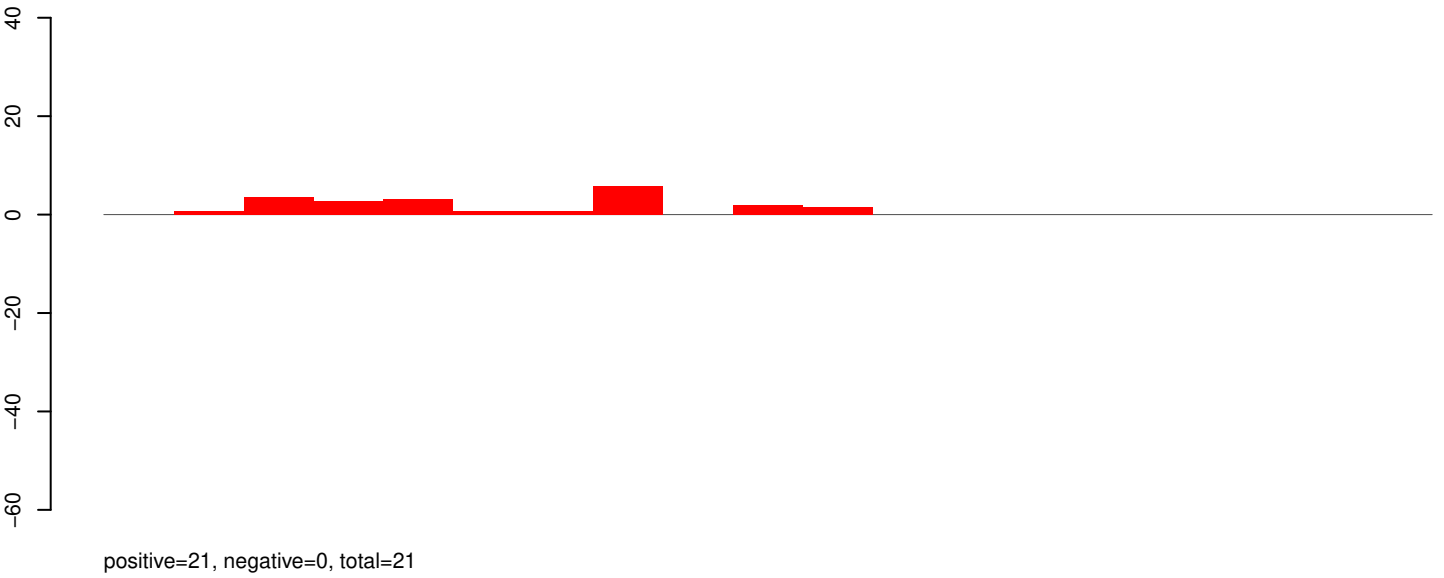


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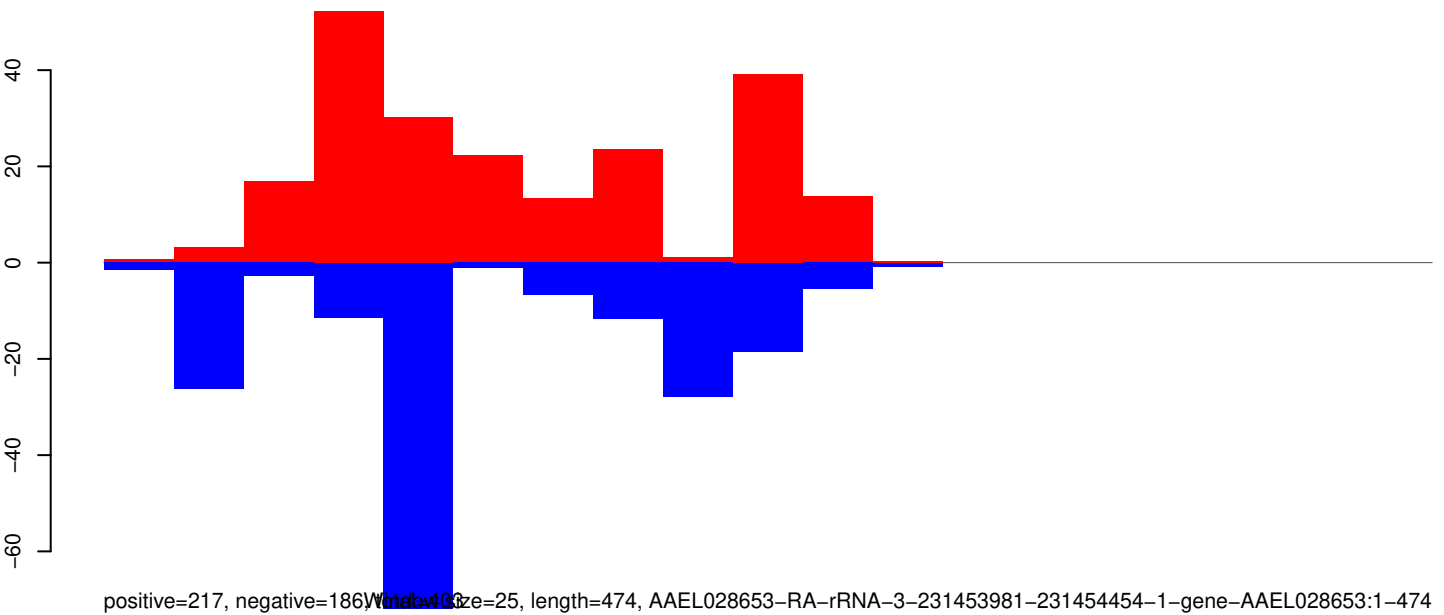
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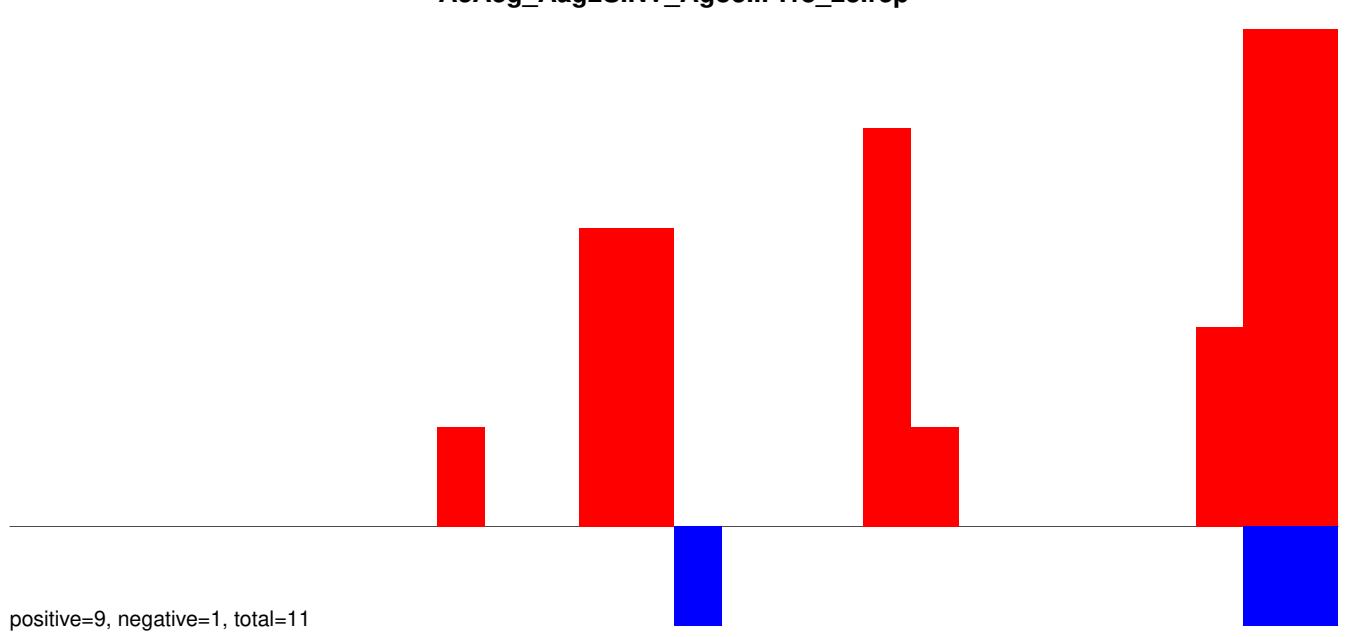
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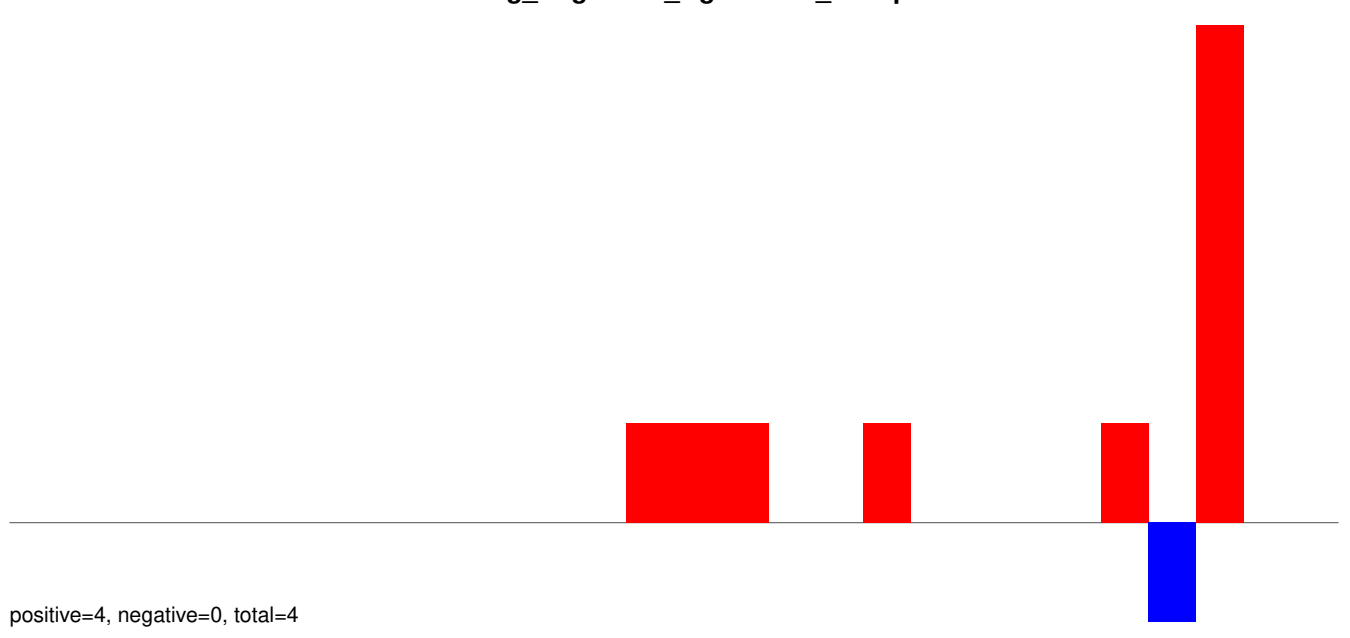
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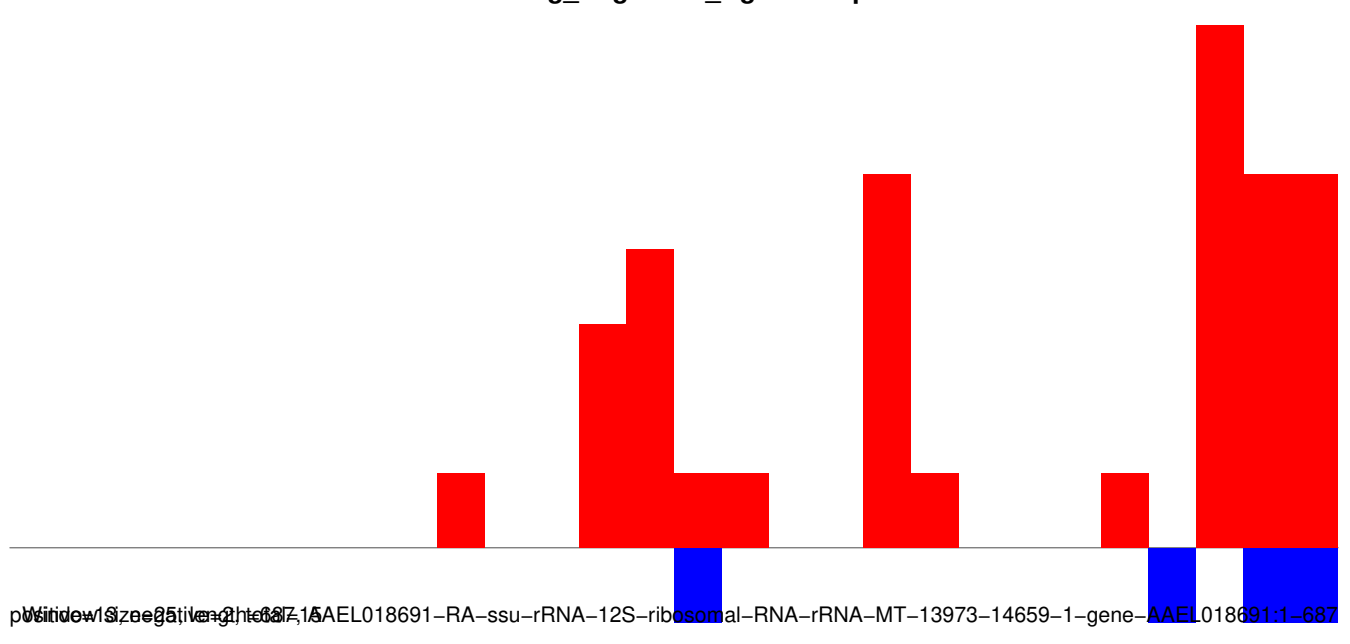
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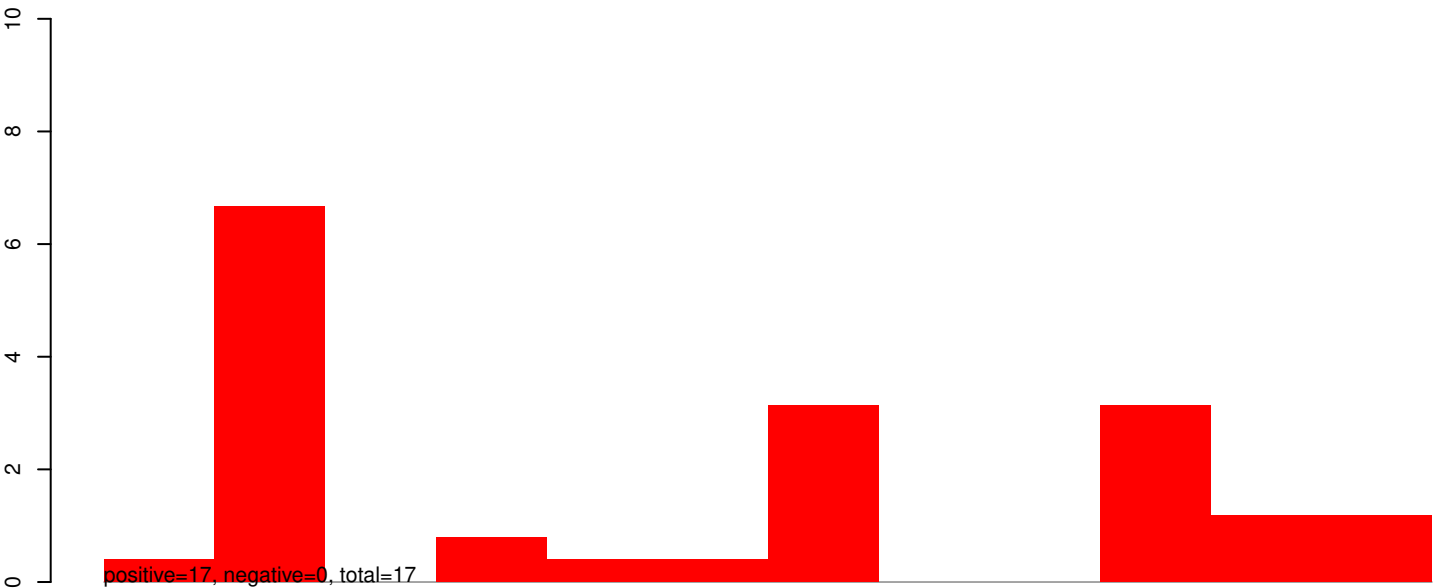


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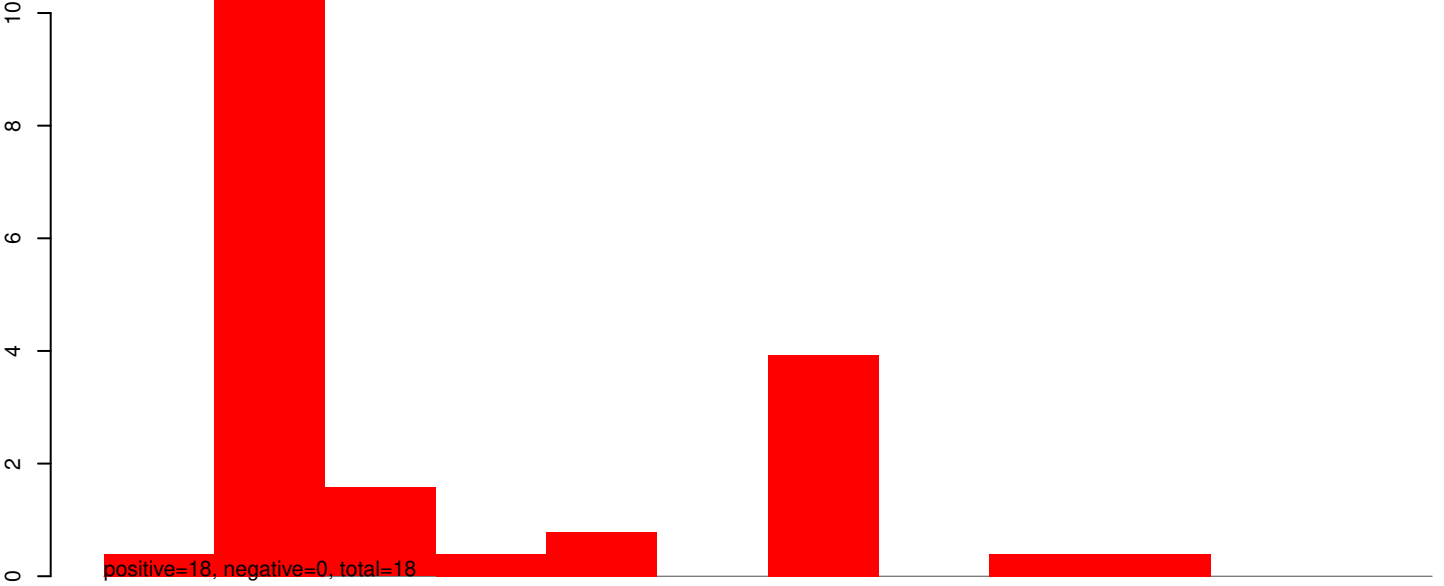
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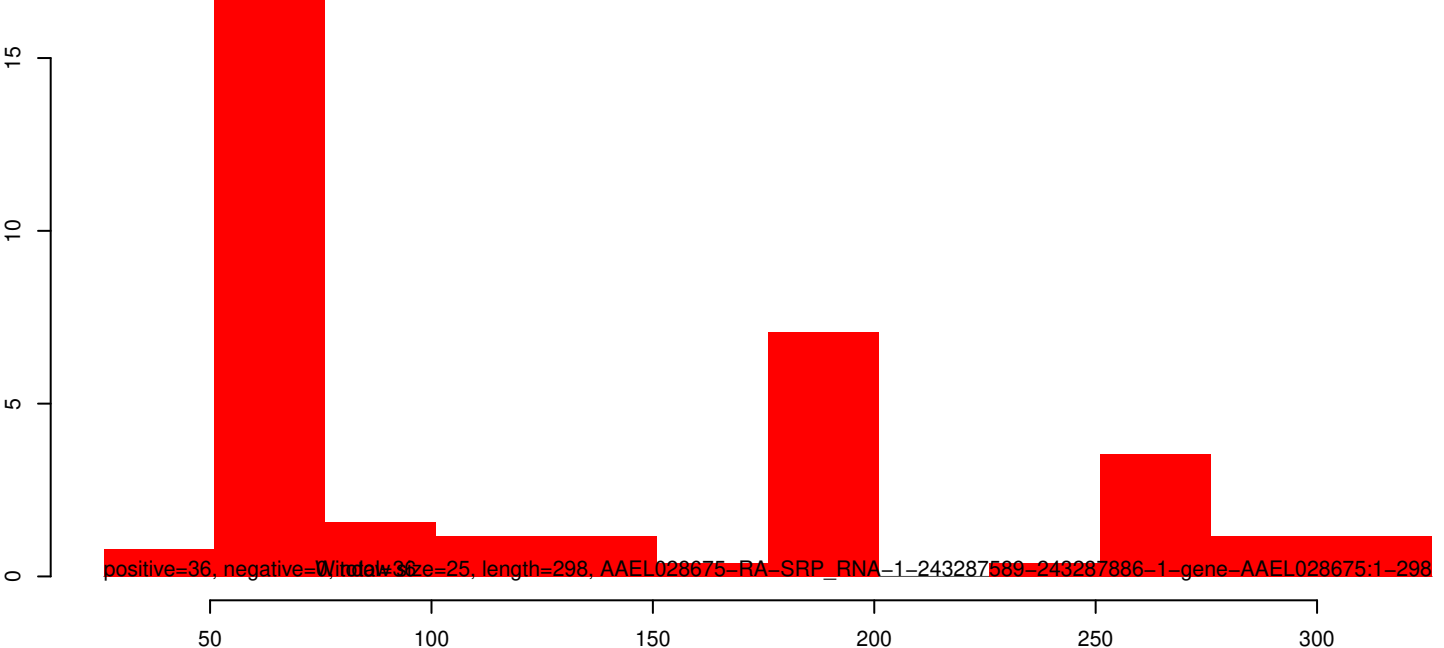
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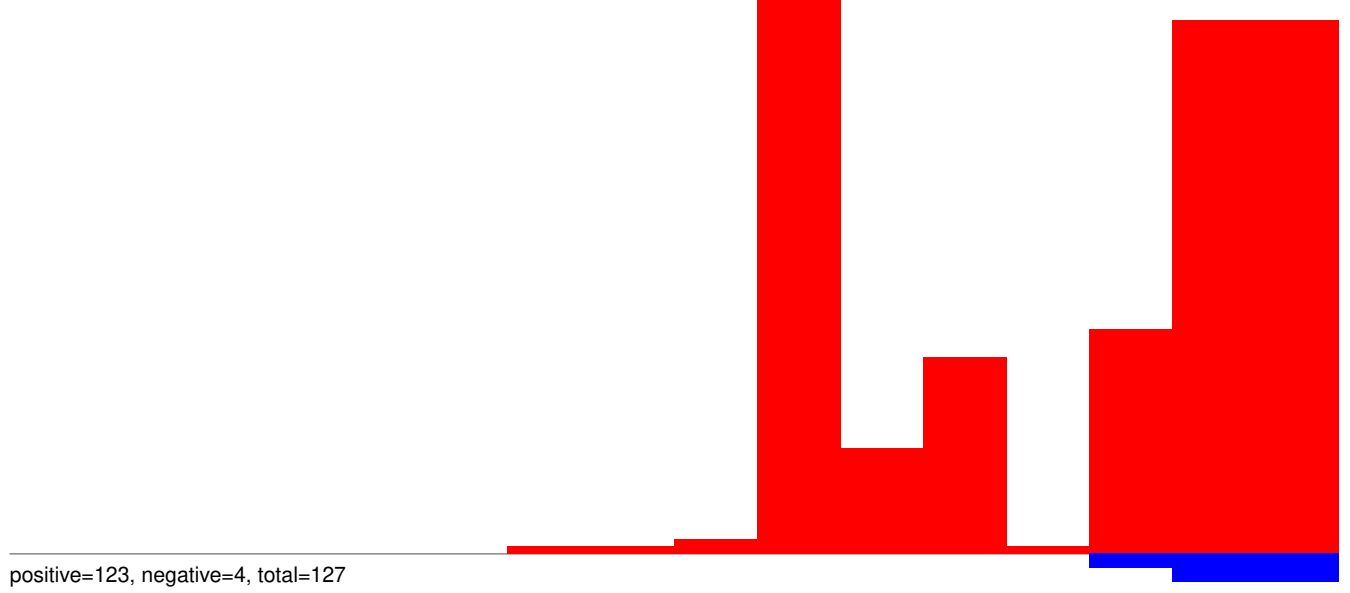
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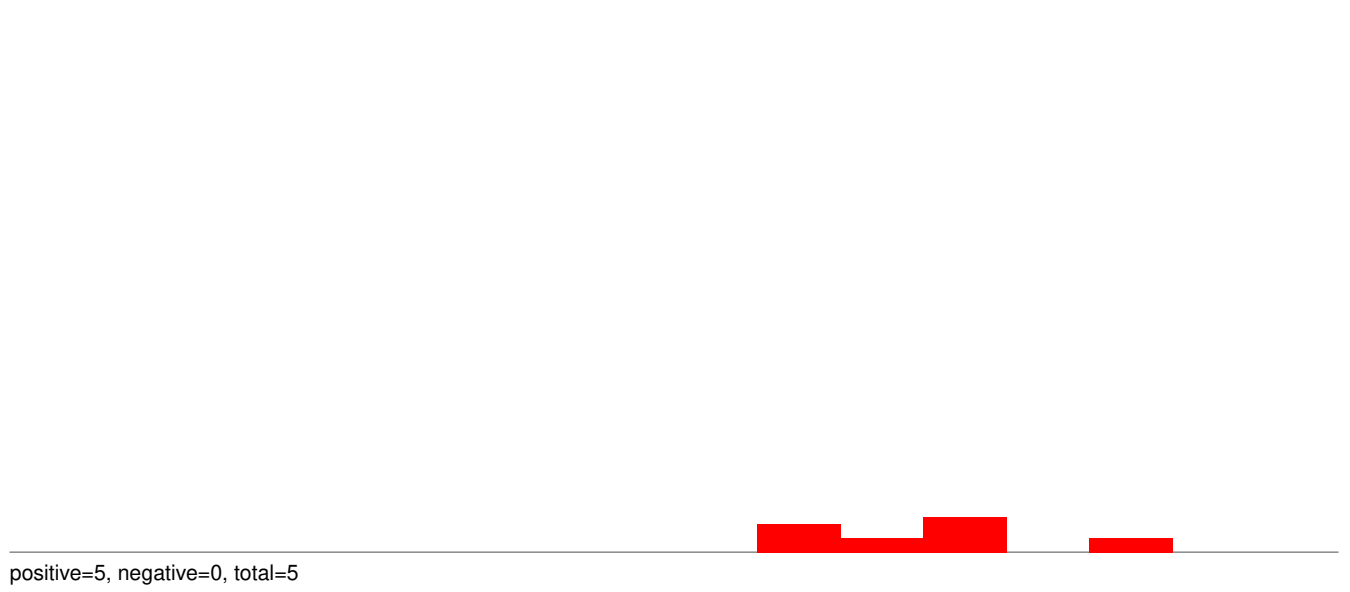
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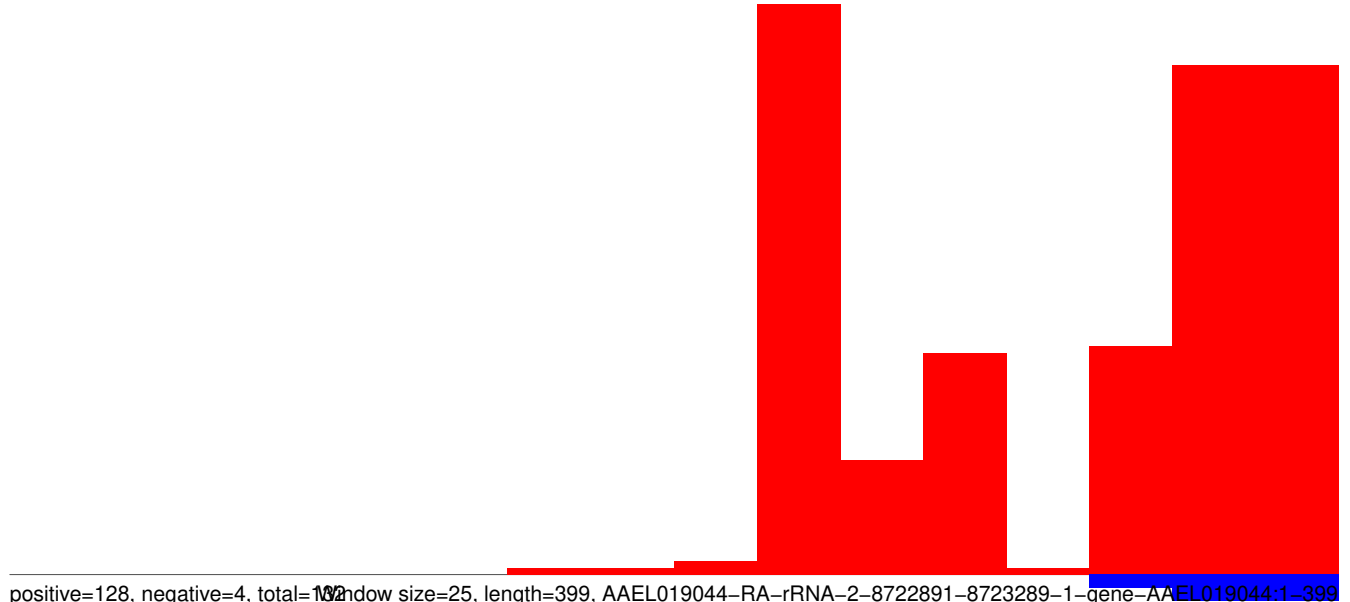
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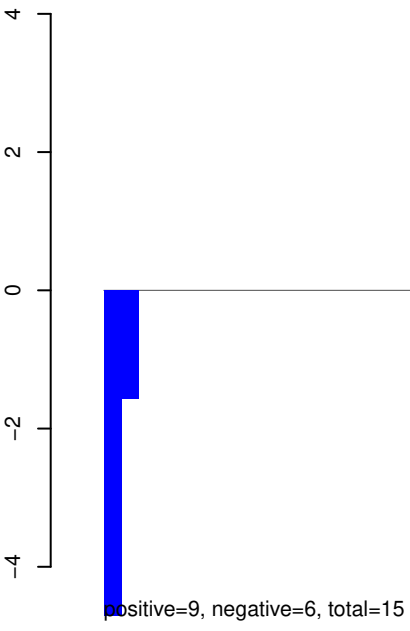
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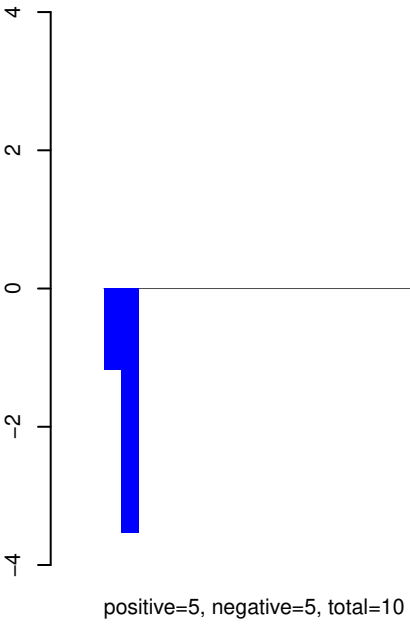
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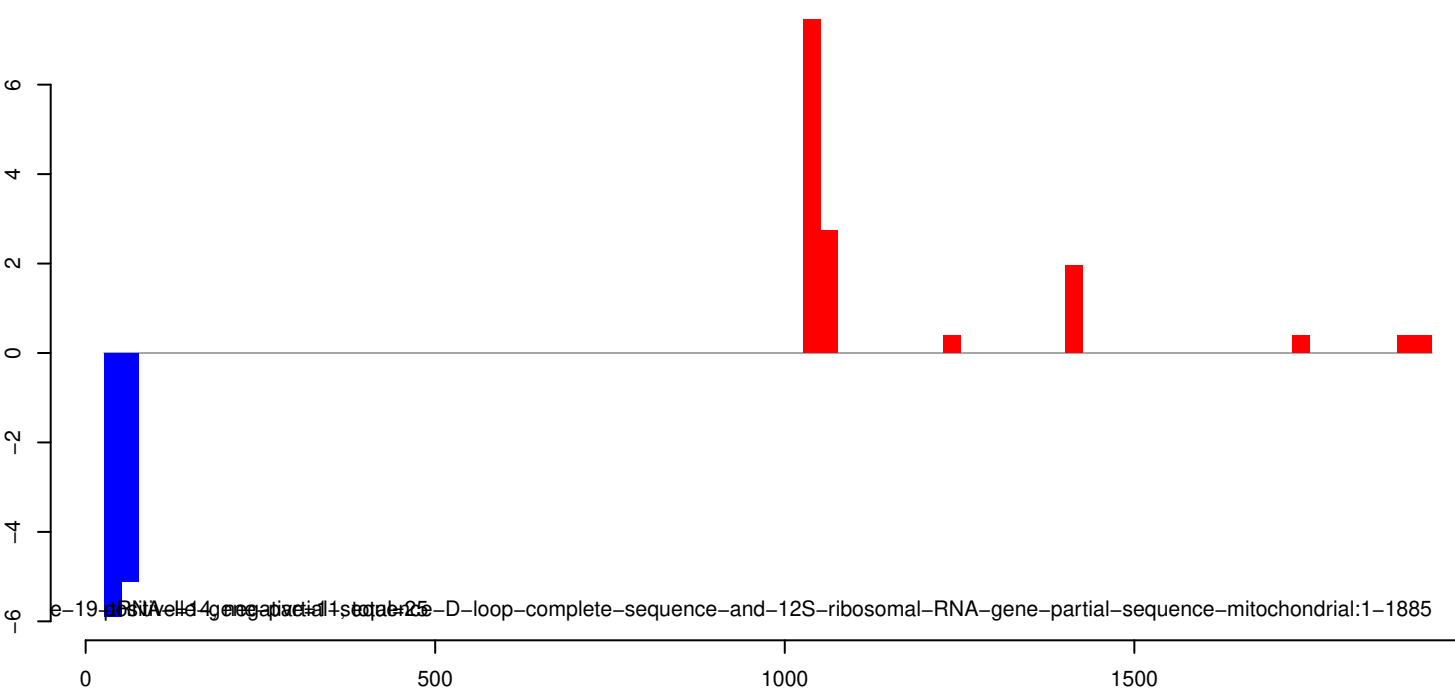
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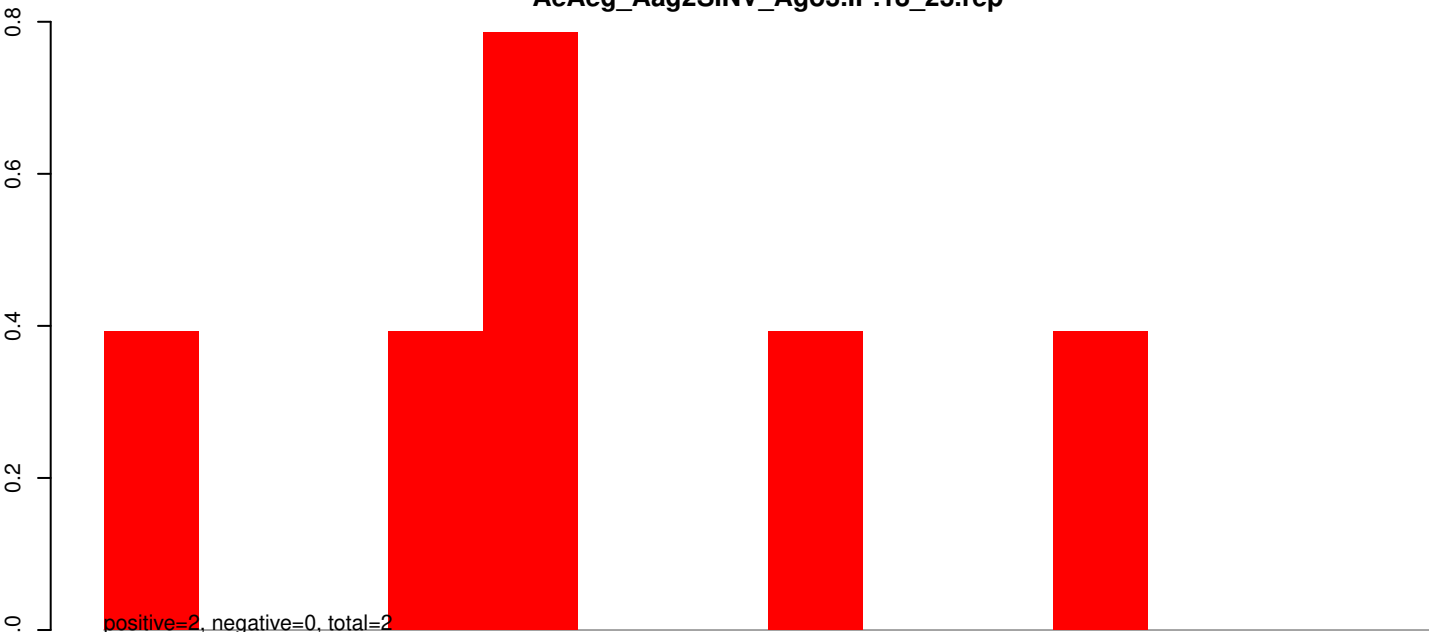


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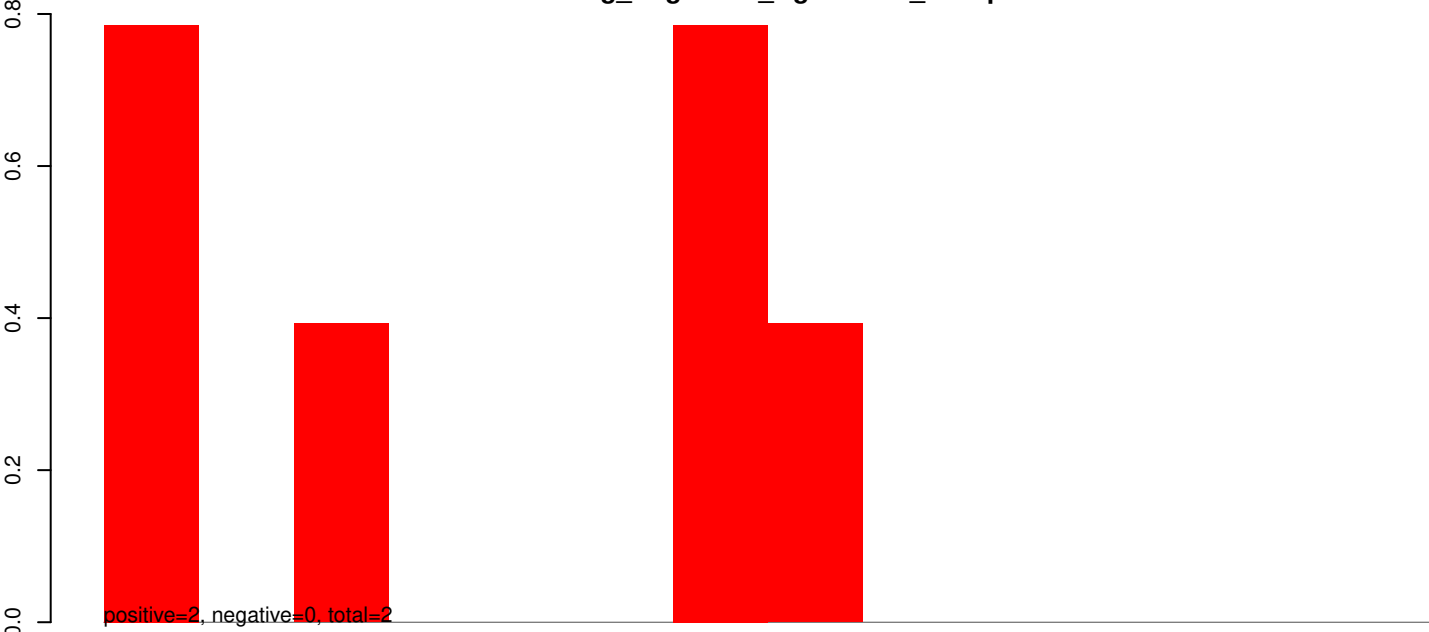


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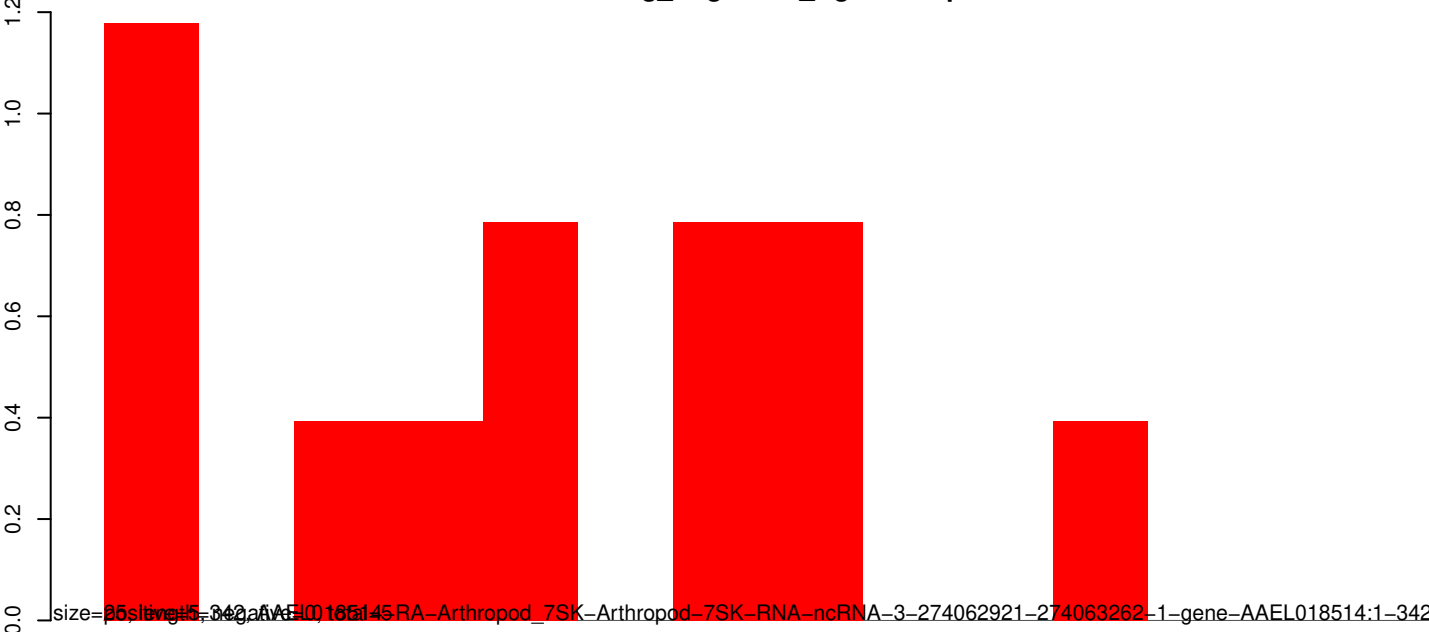
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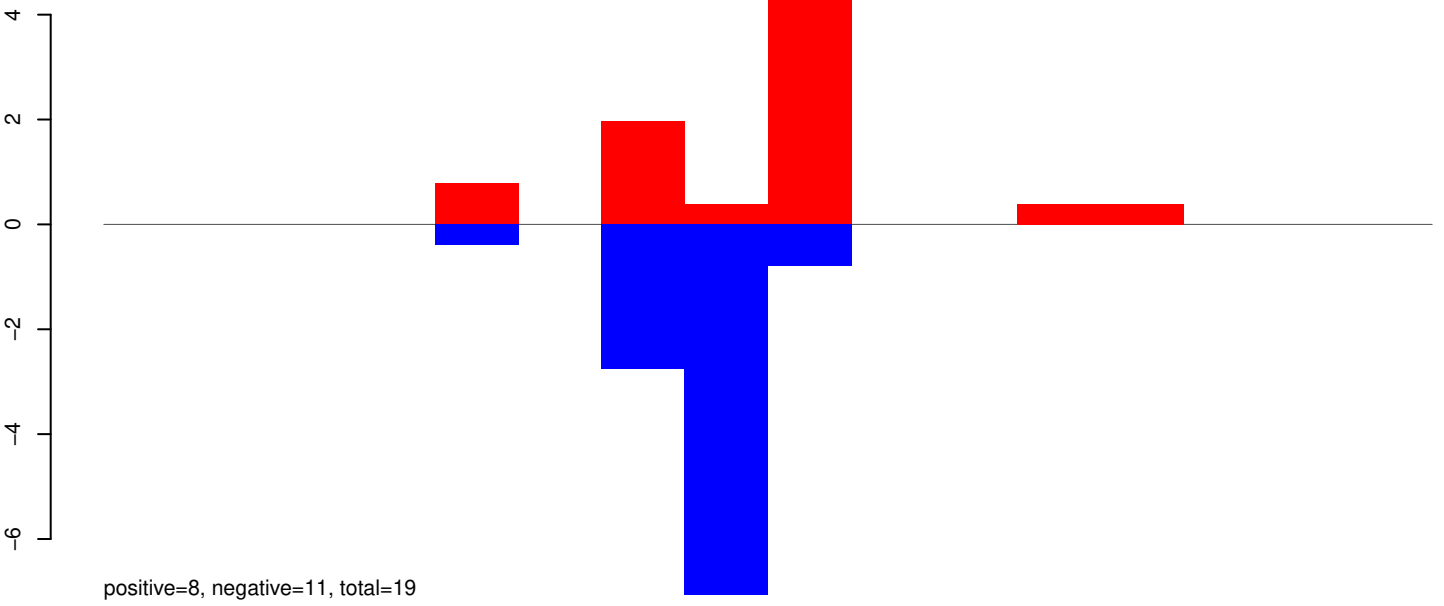
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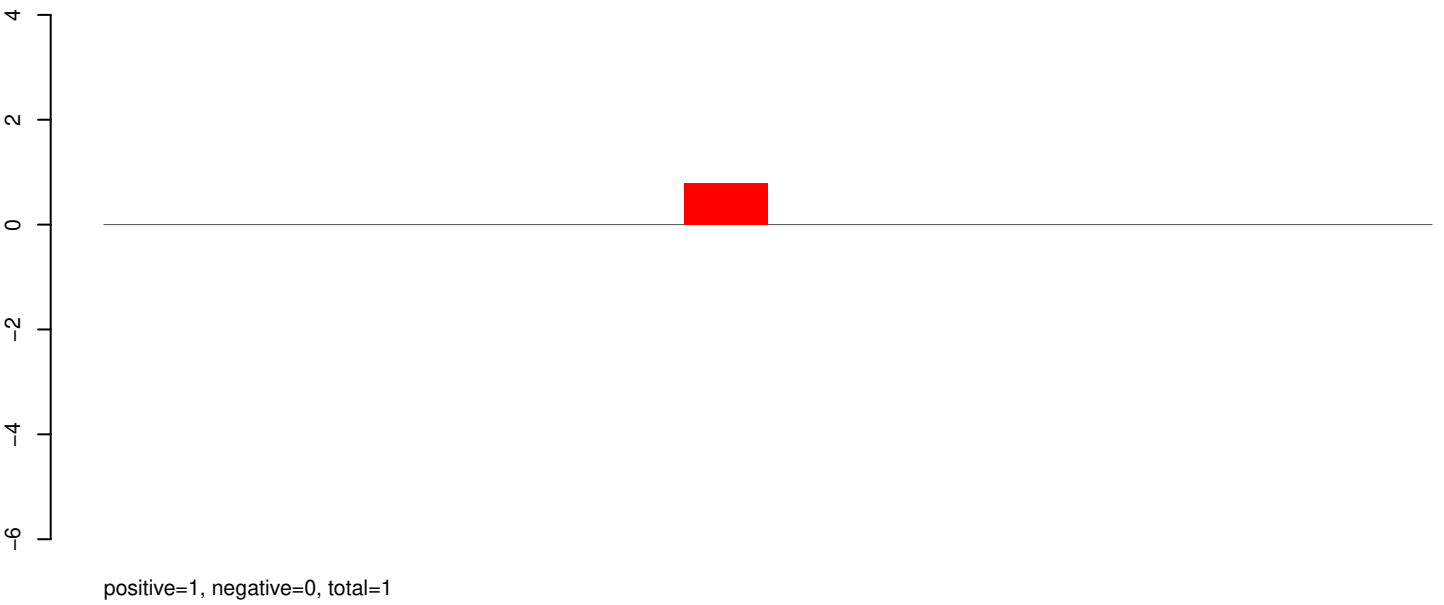
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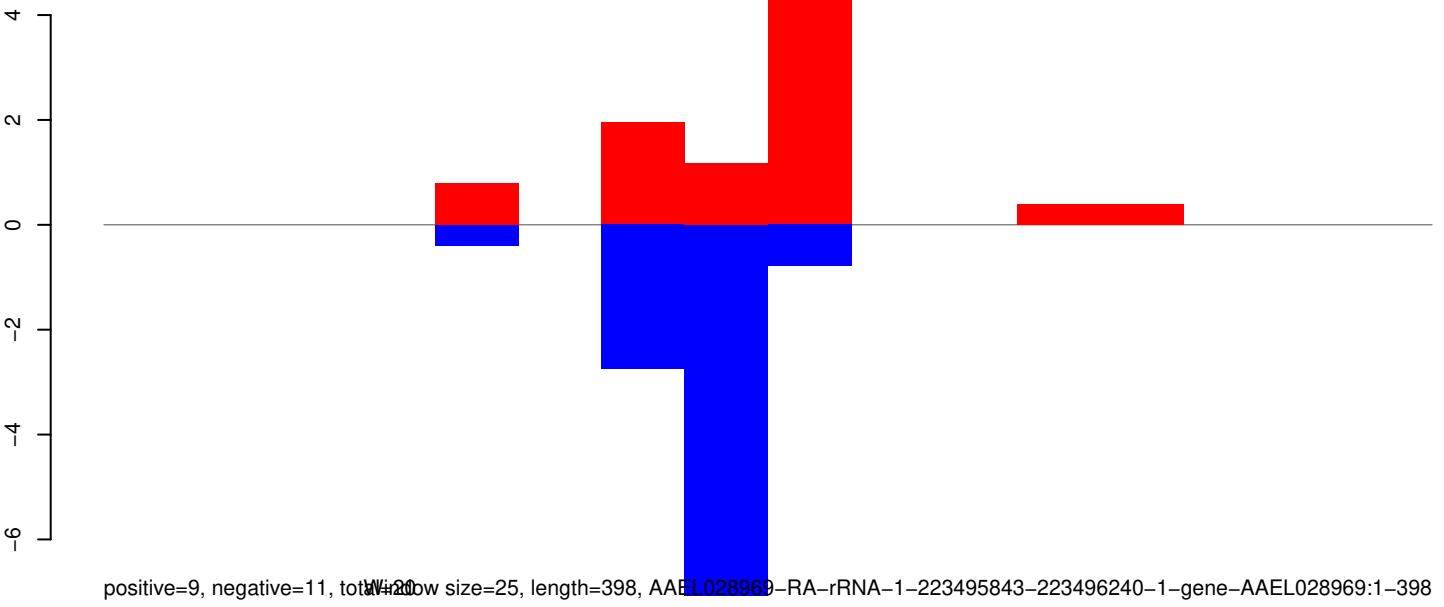
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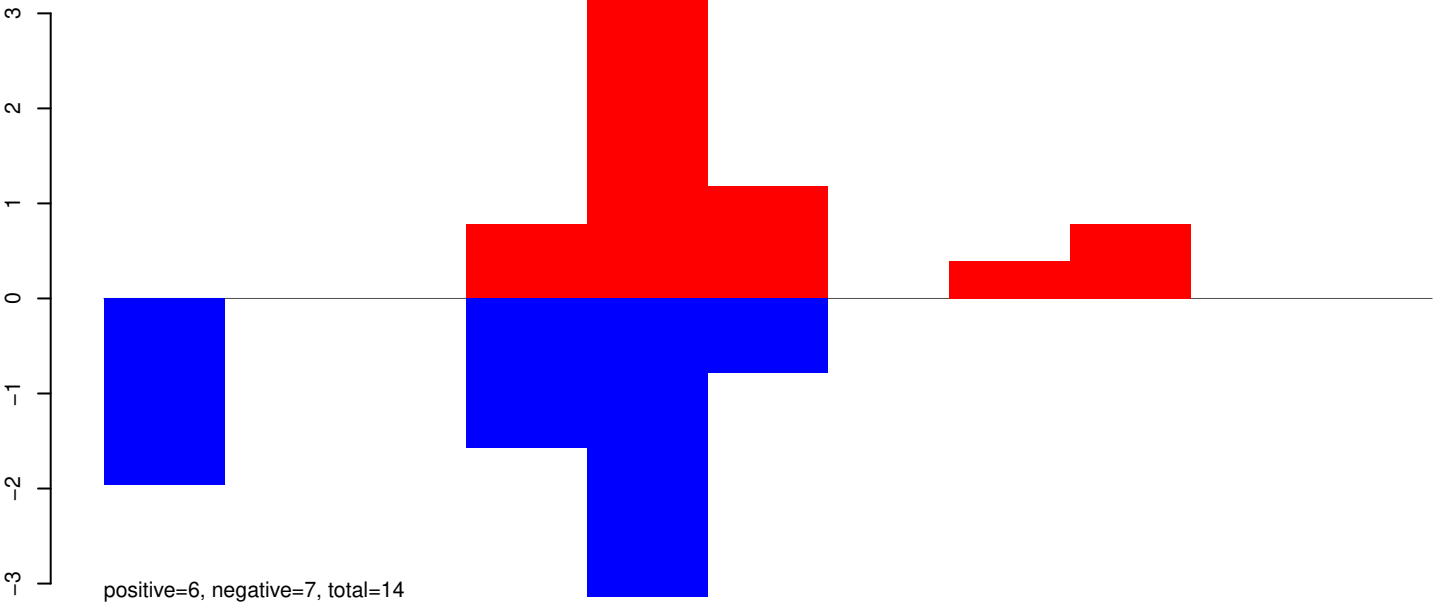


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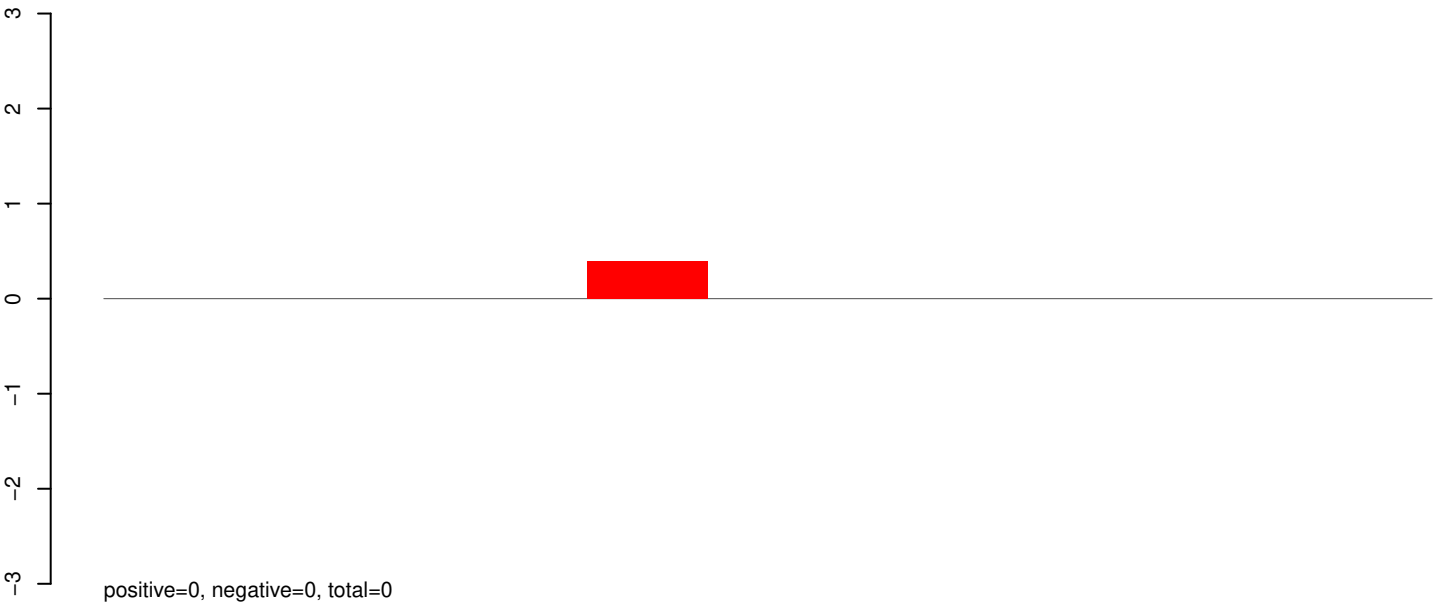


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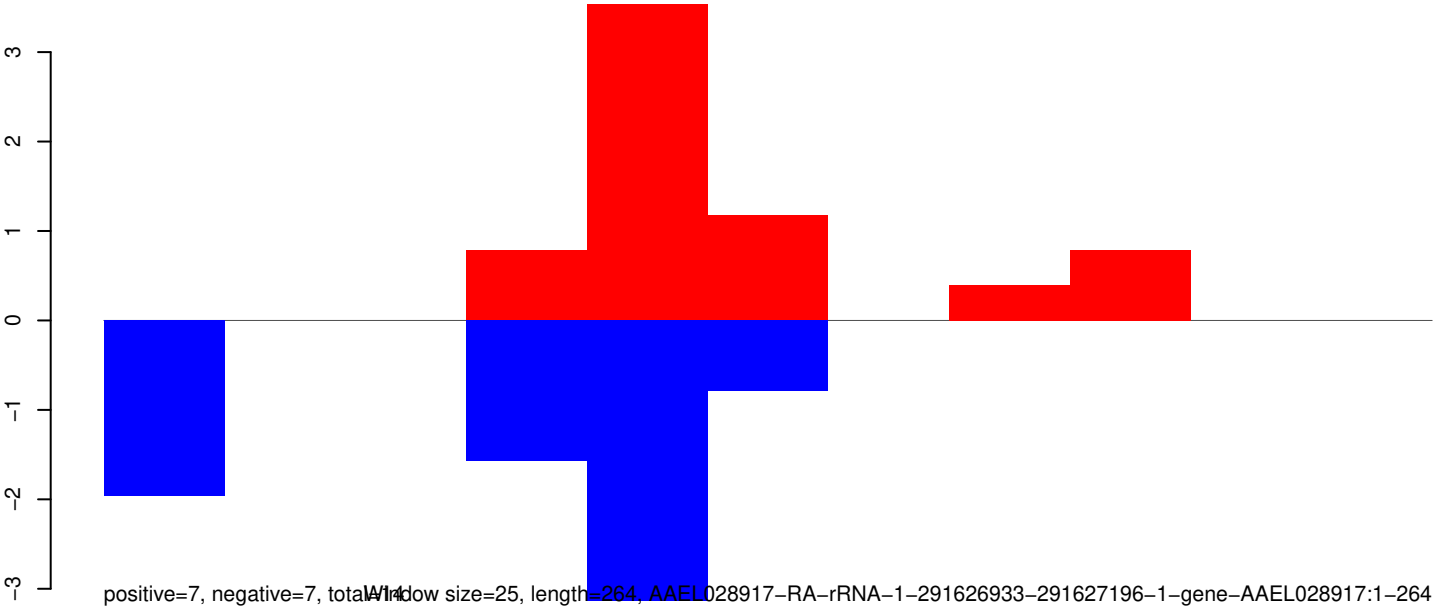
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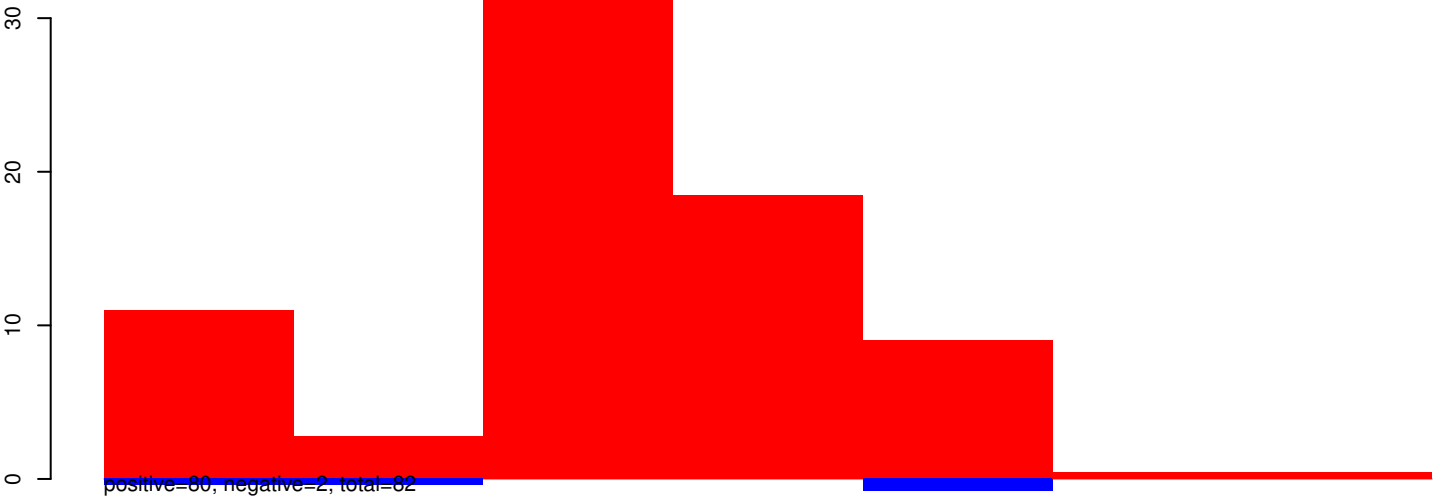


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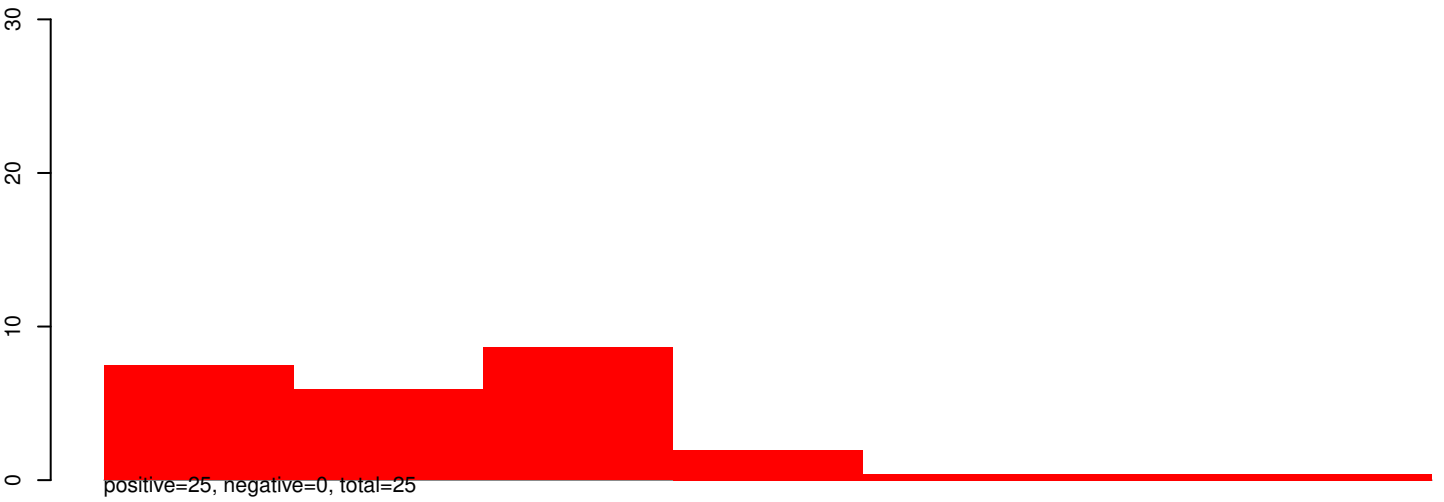


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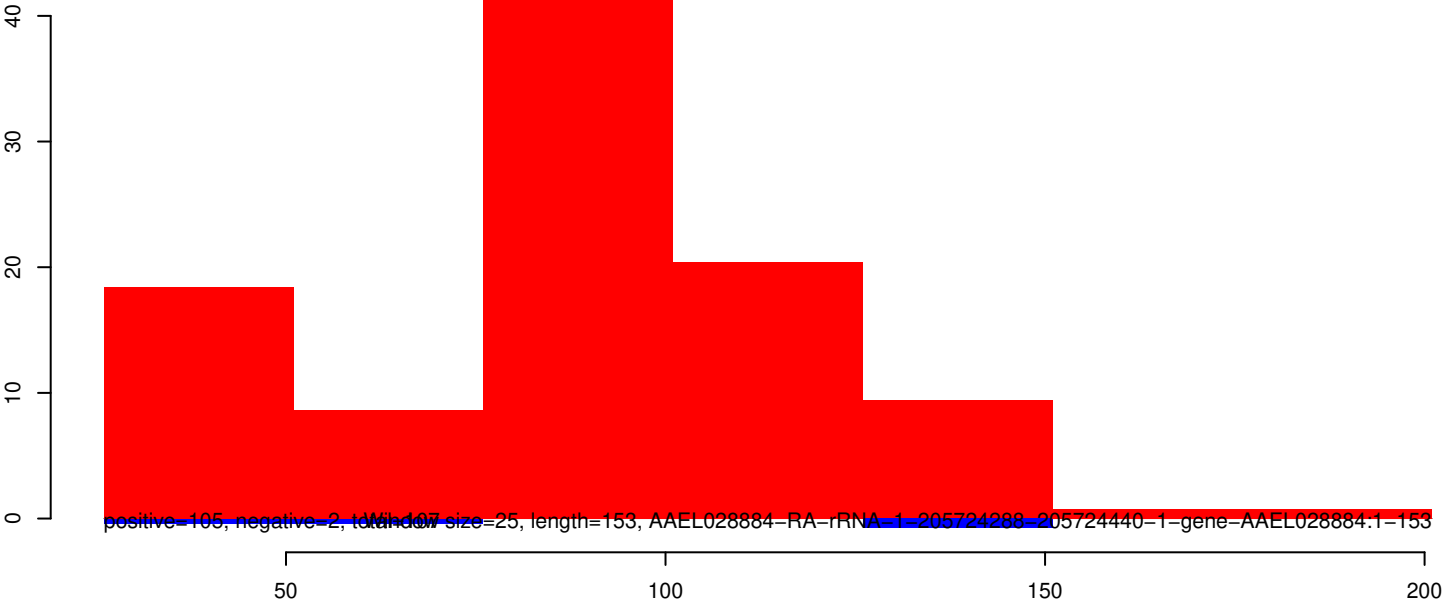
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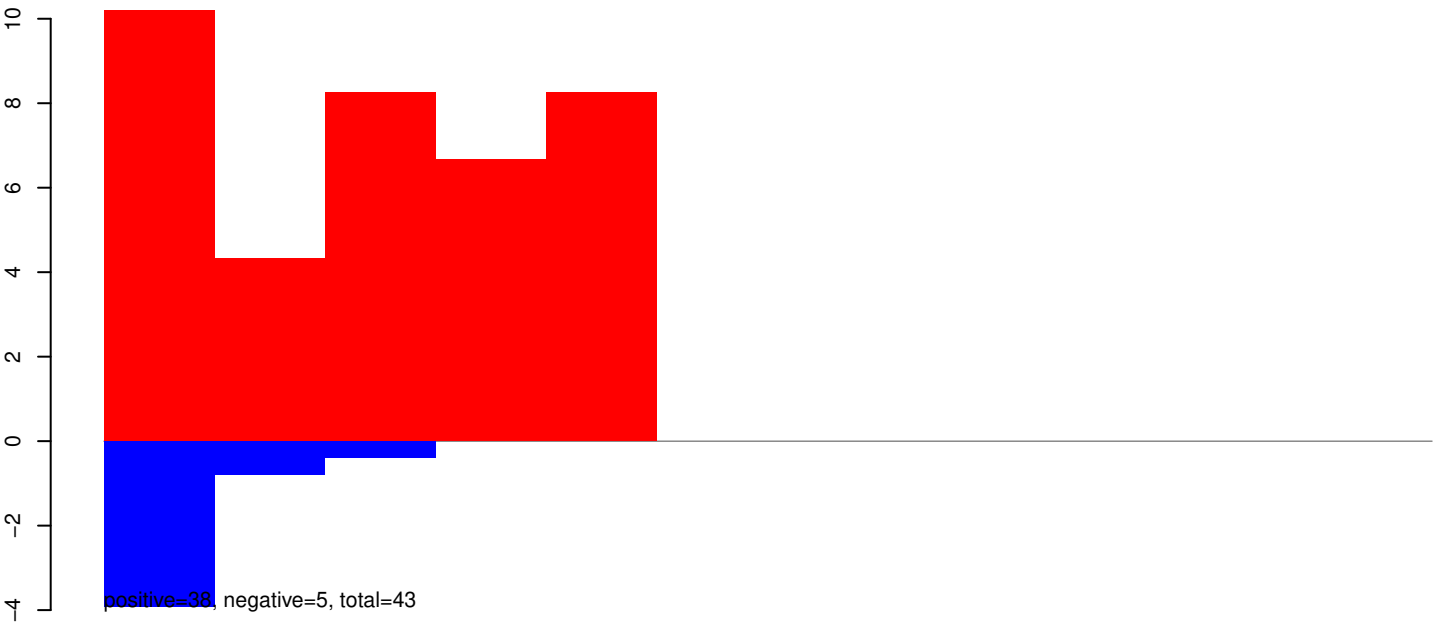
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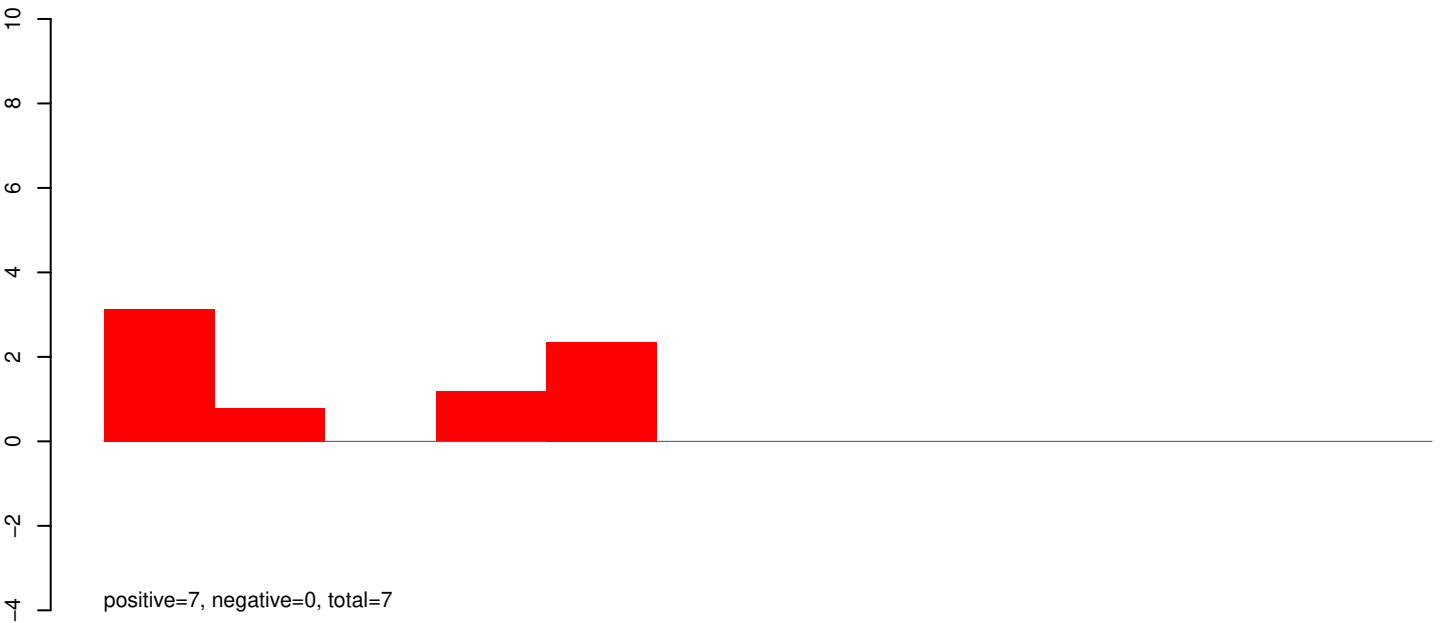
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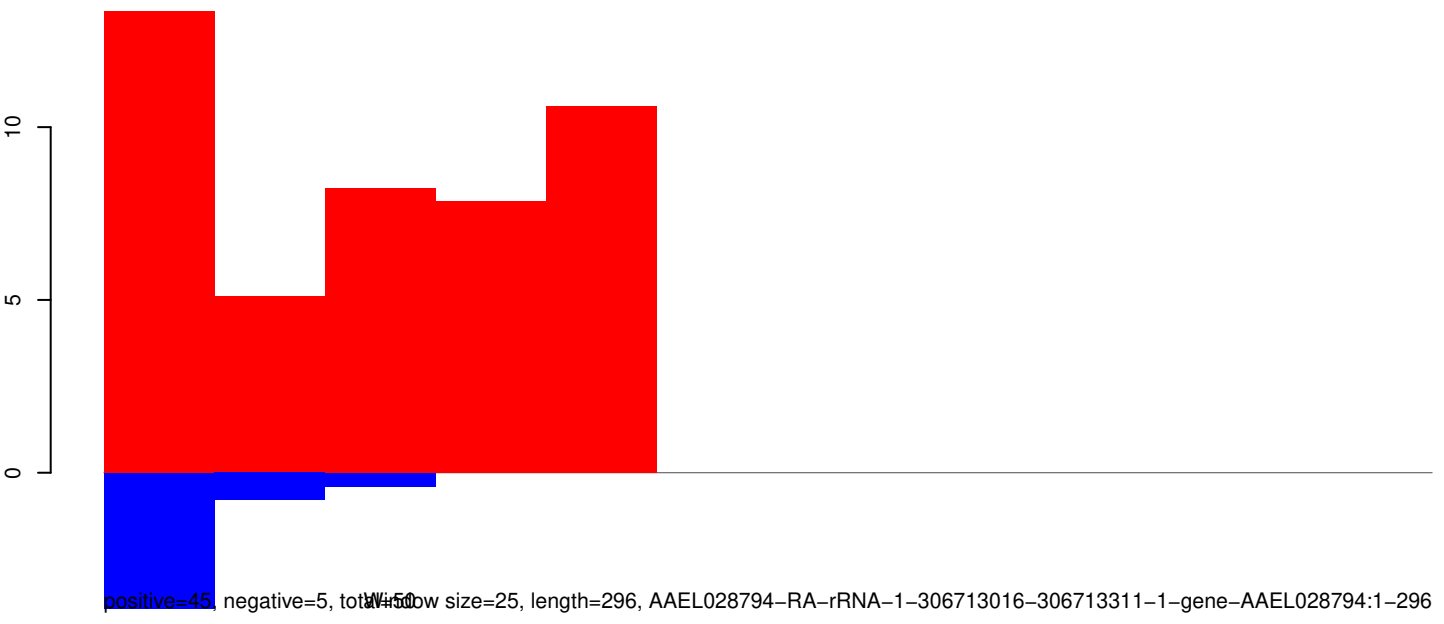
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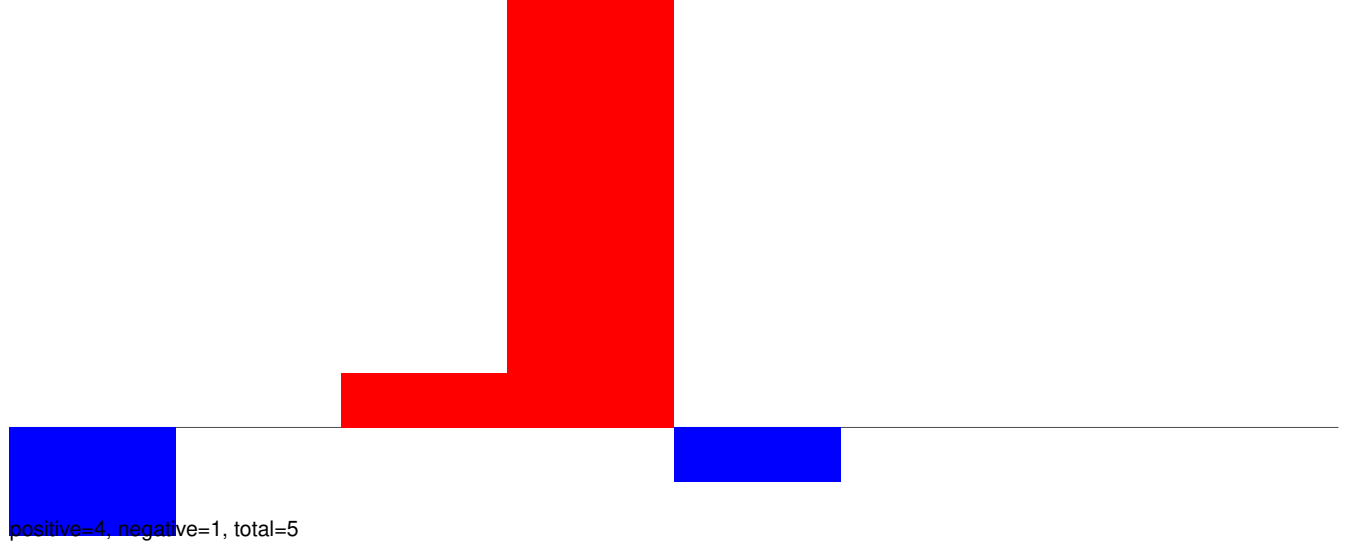
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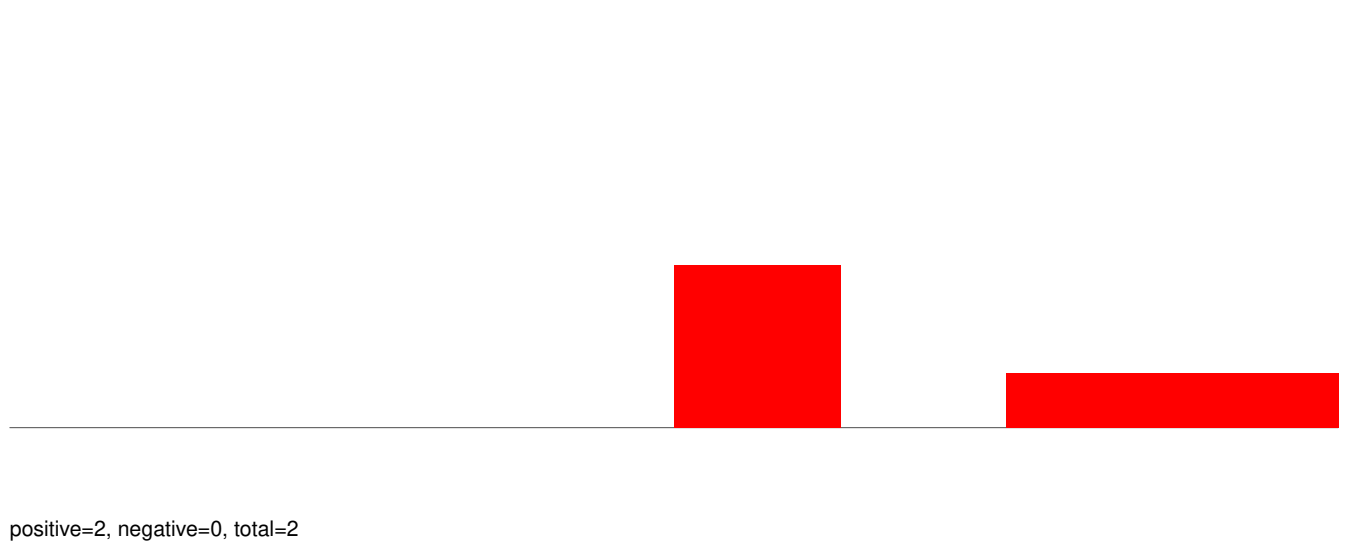
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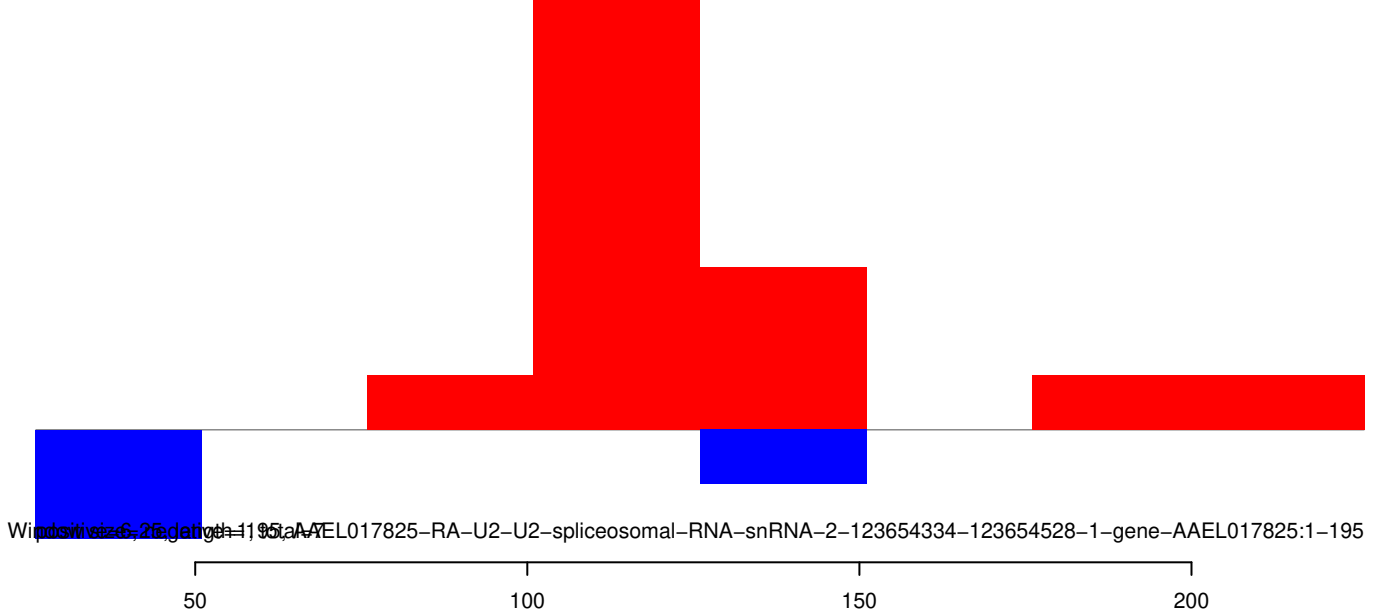
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AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep

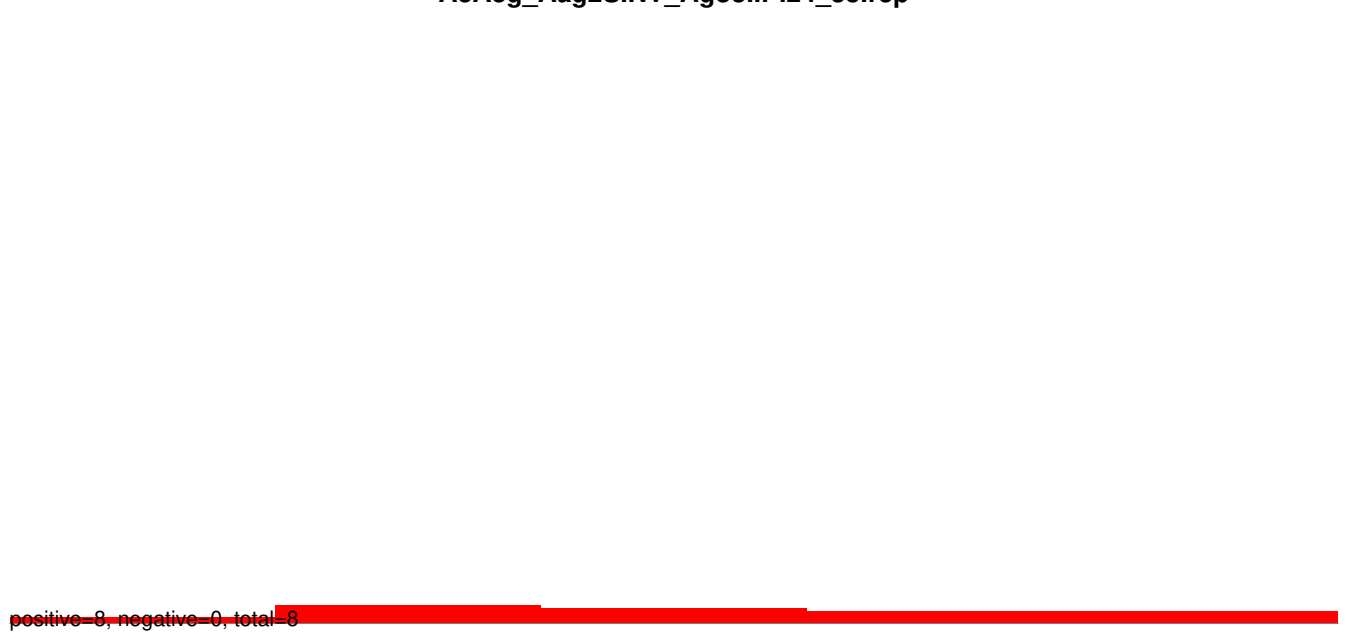


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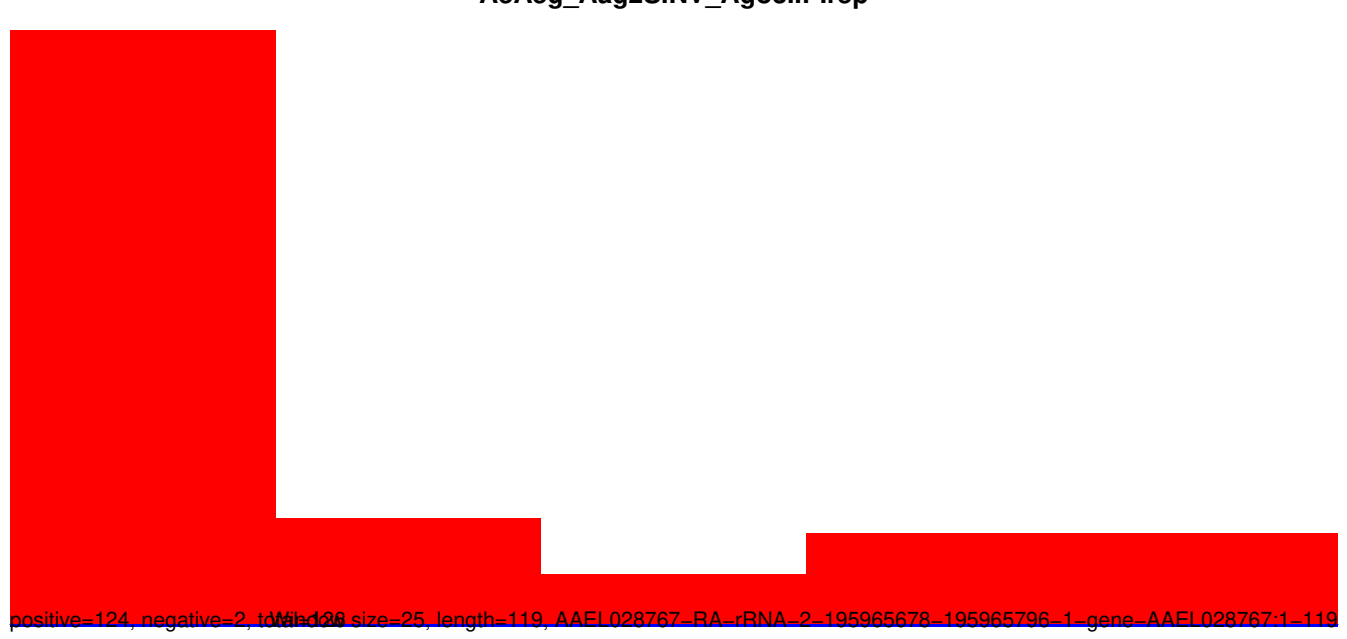
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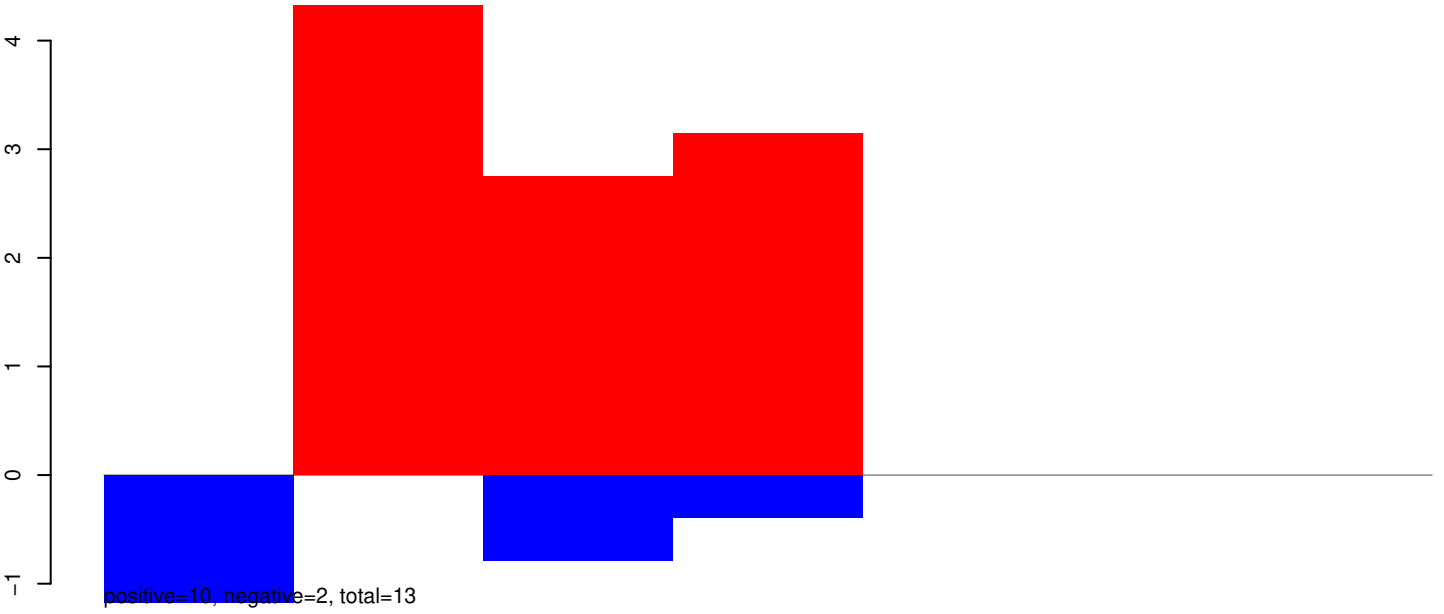
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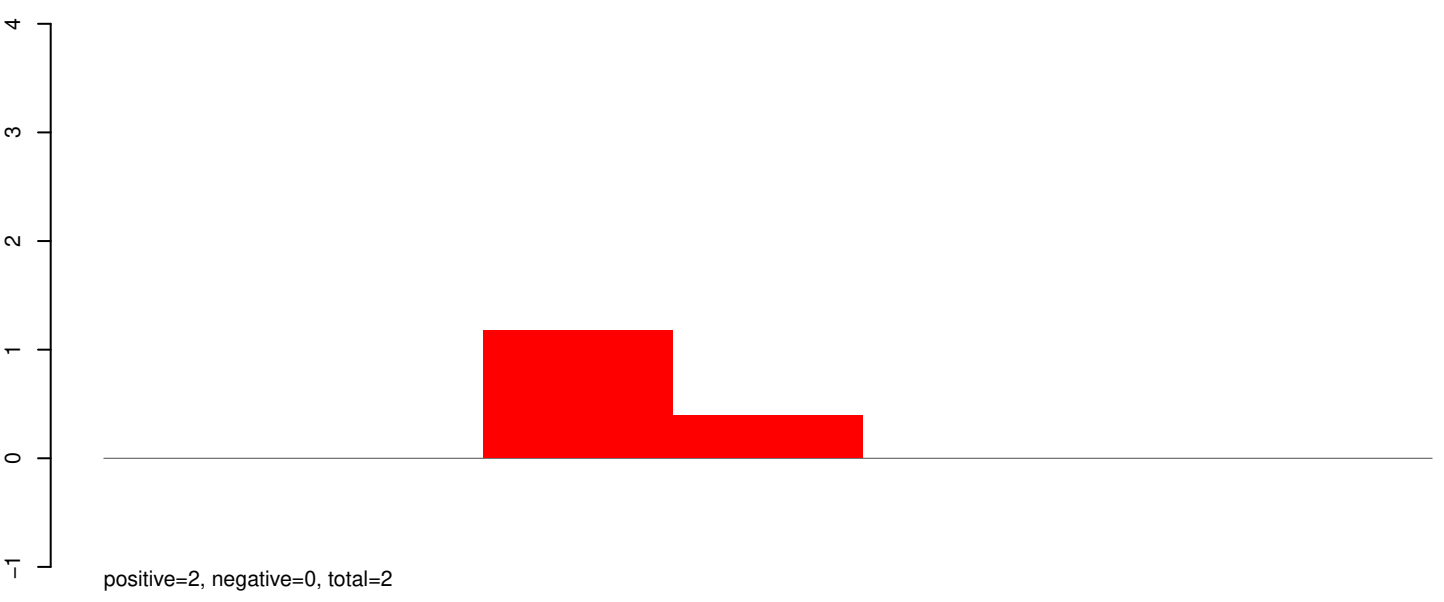
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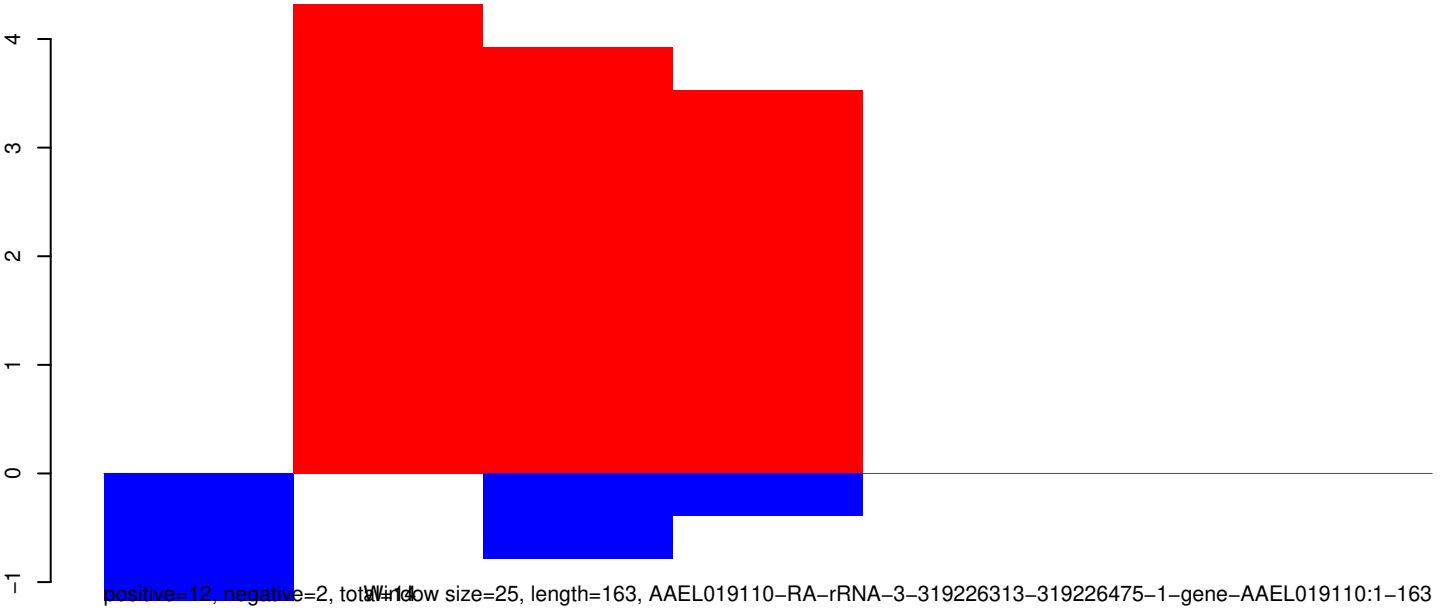
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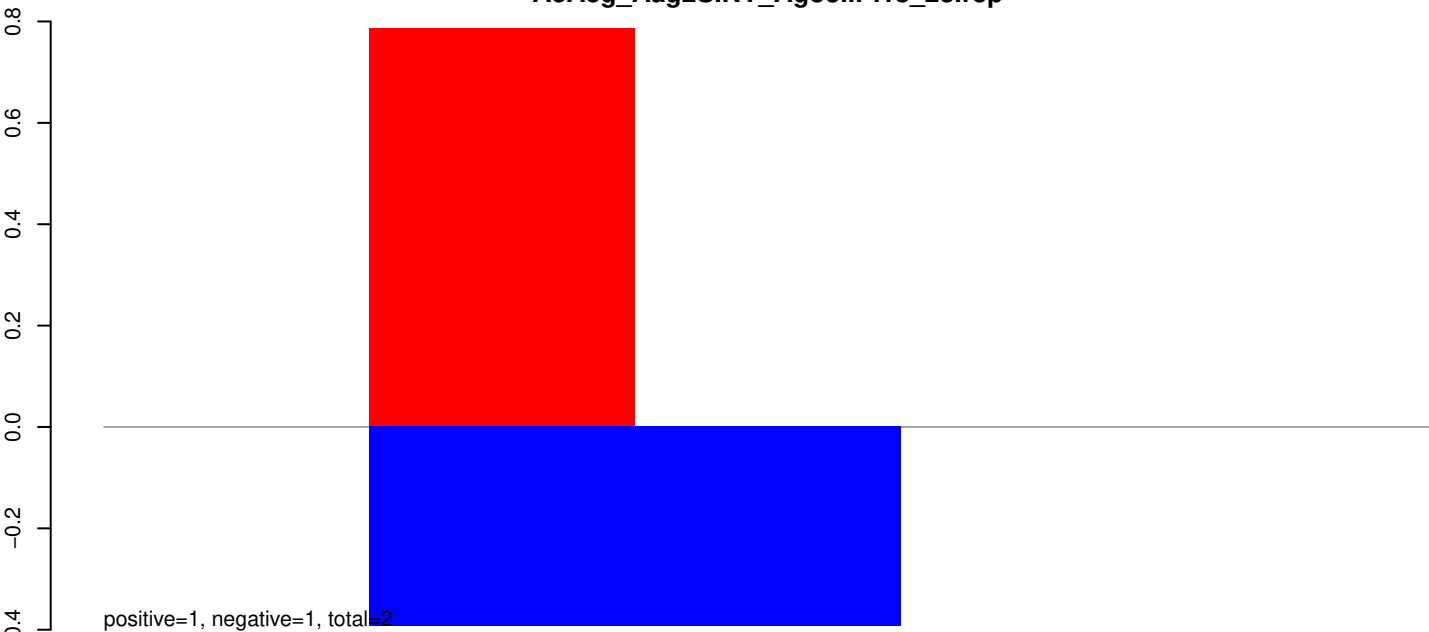


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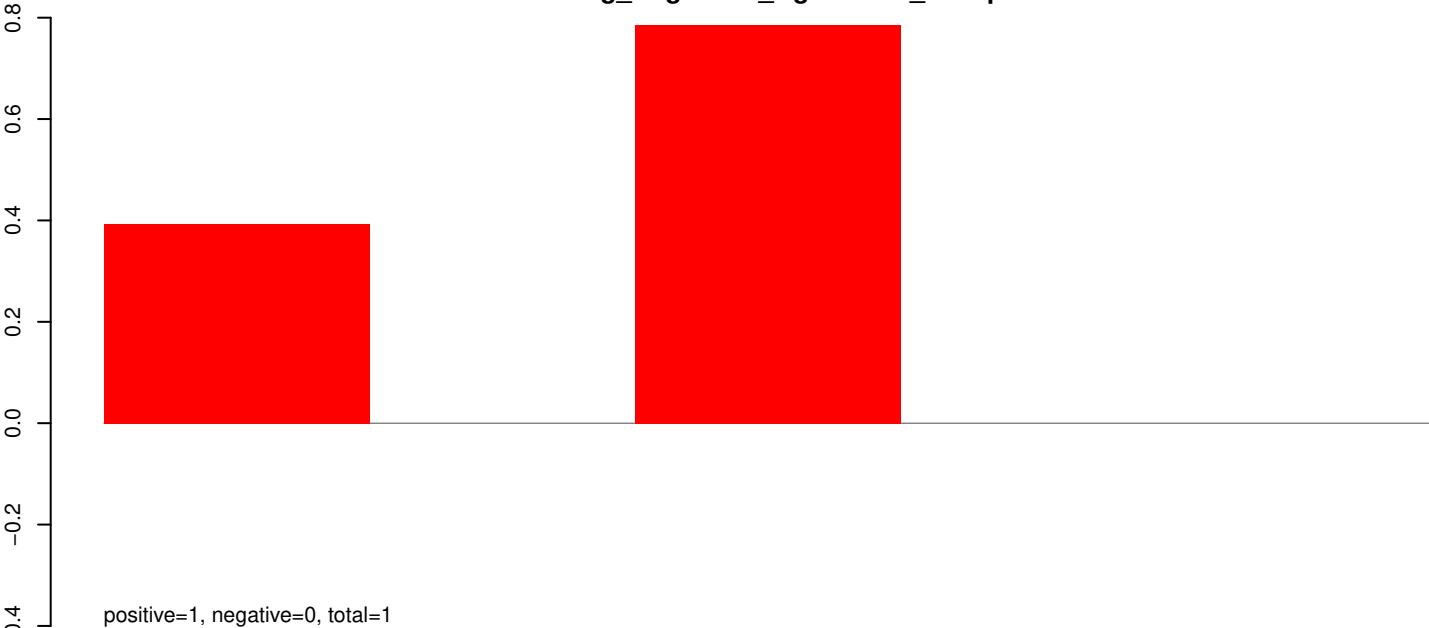


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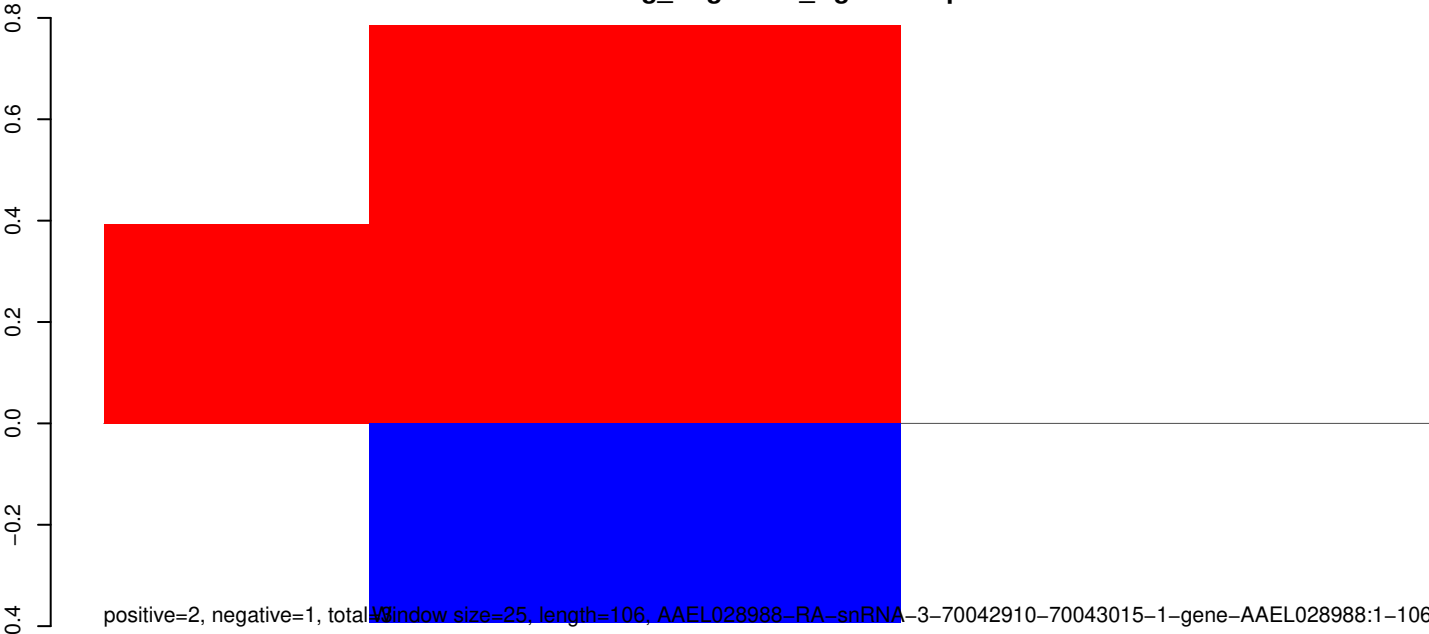
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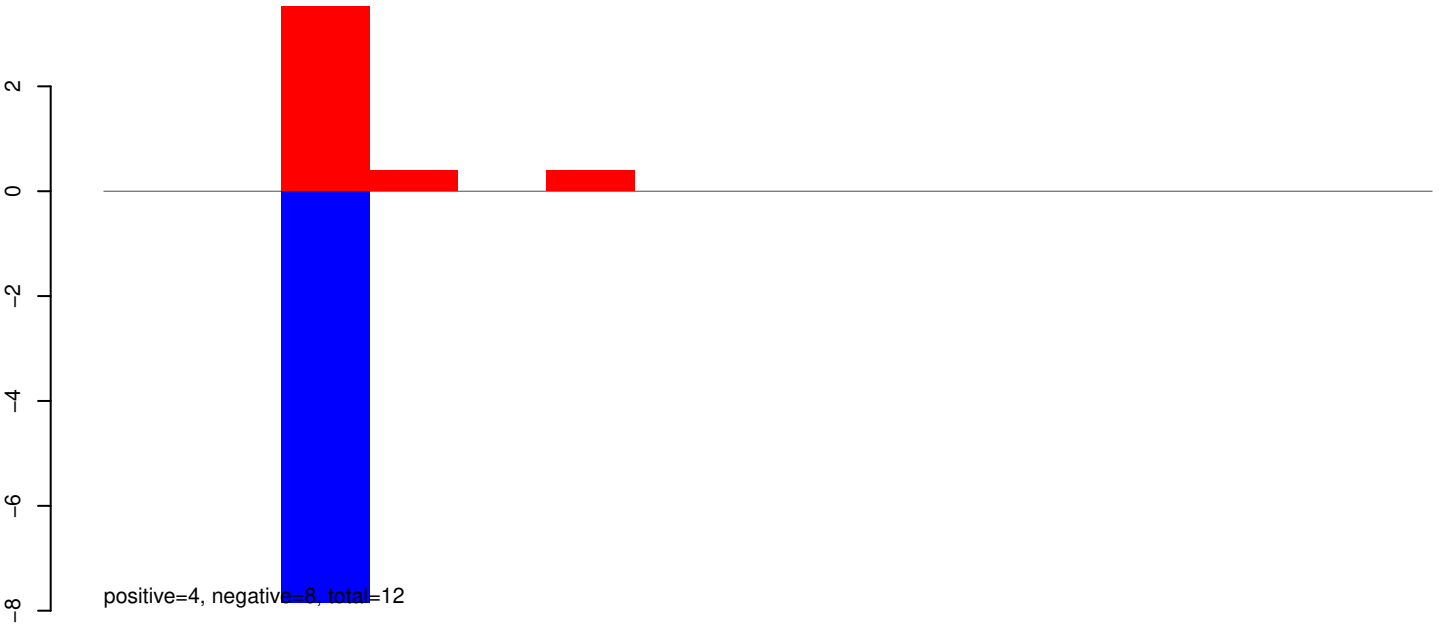
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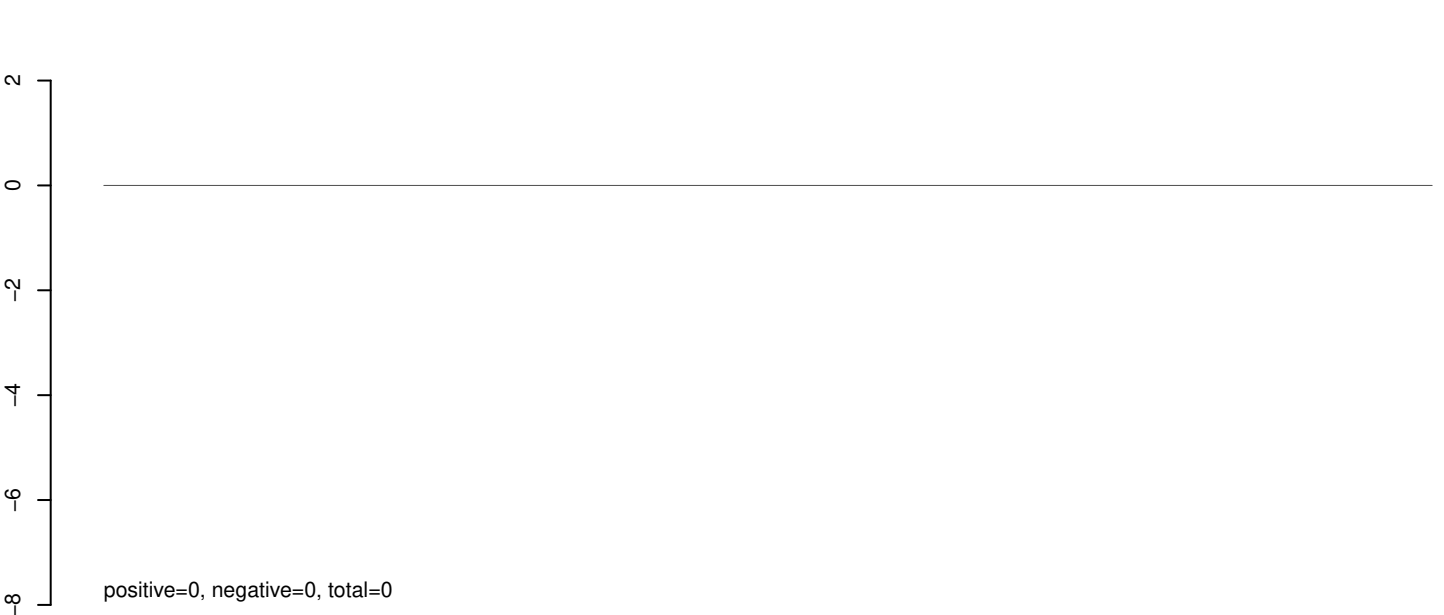
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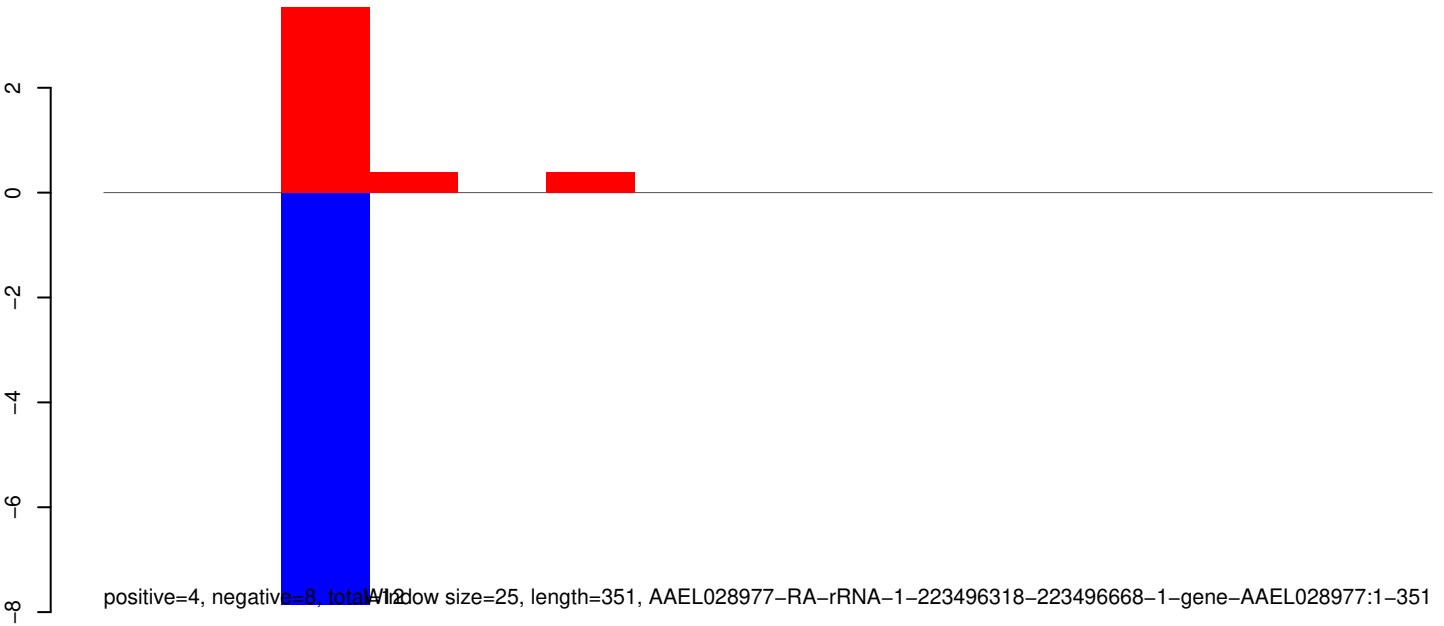
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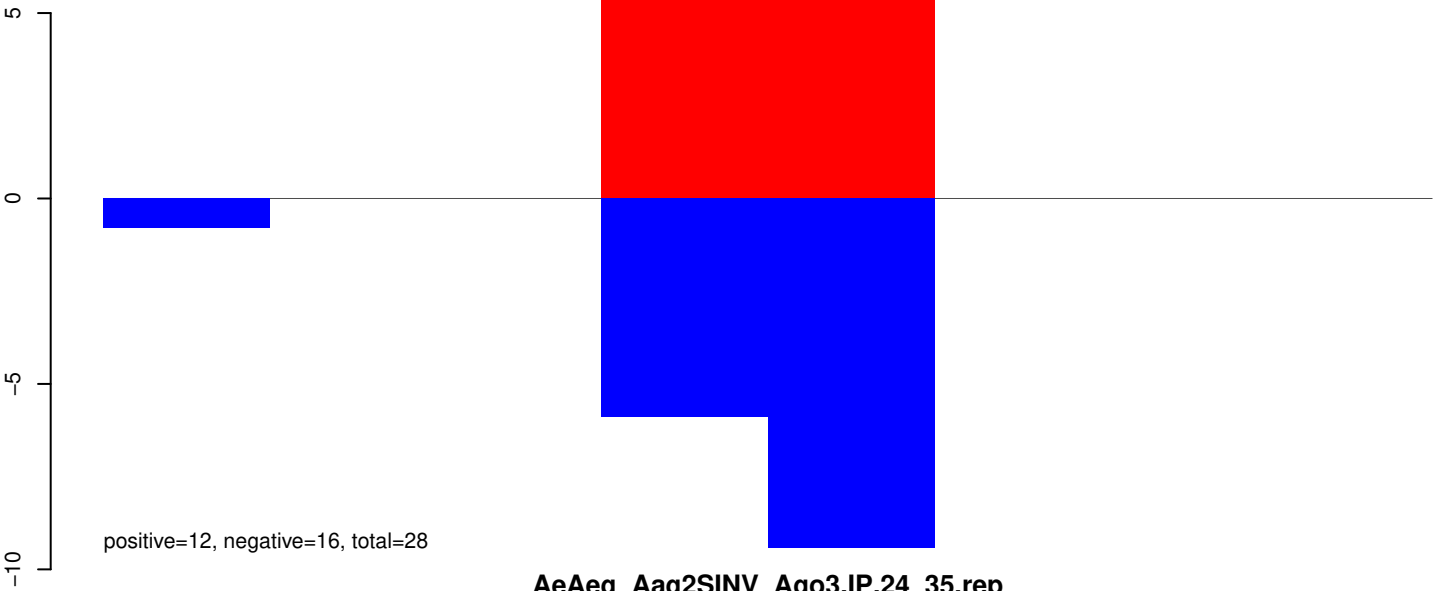
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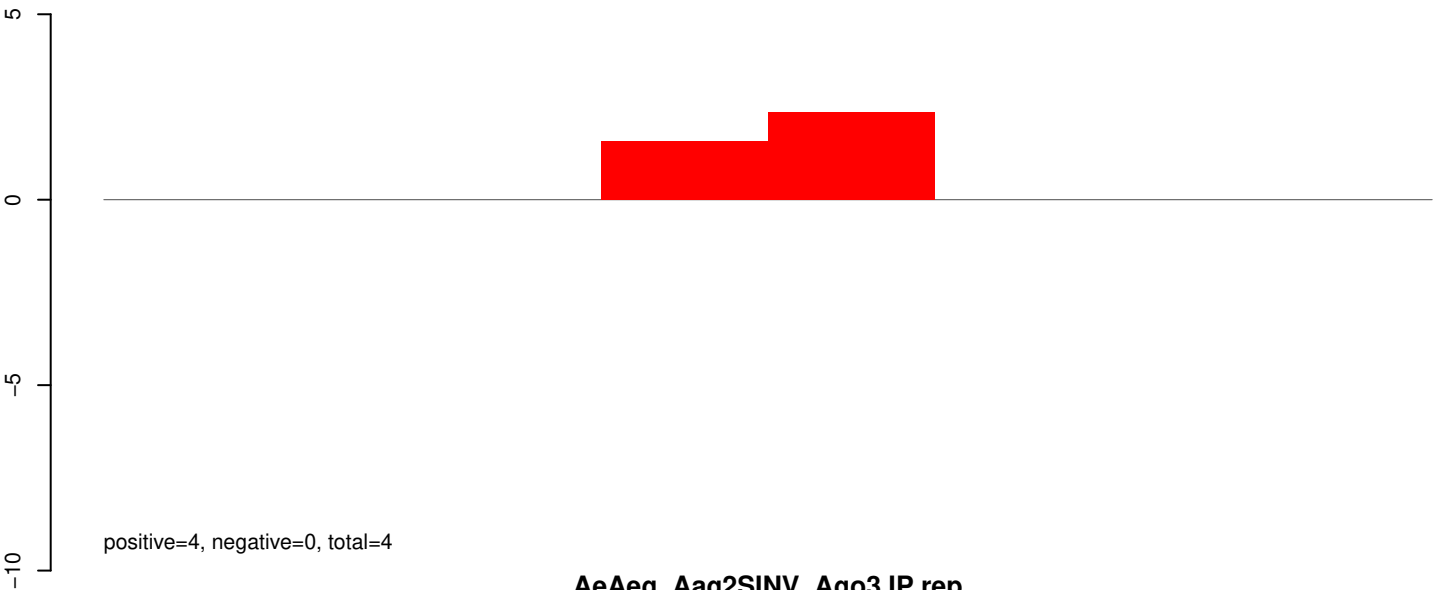
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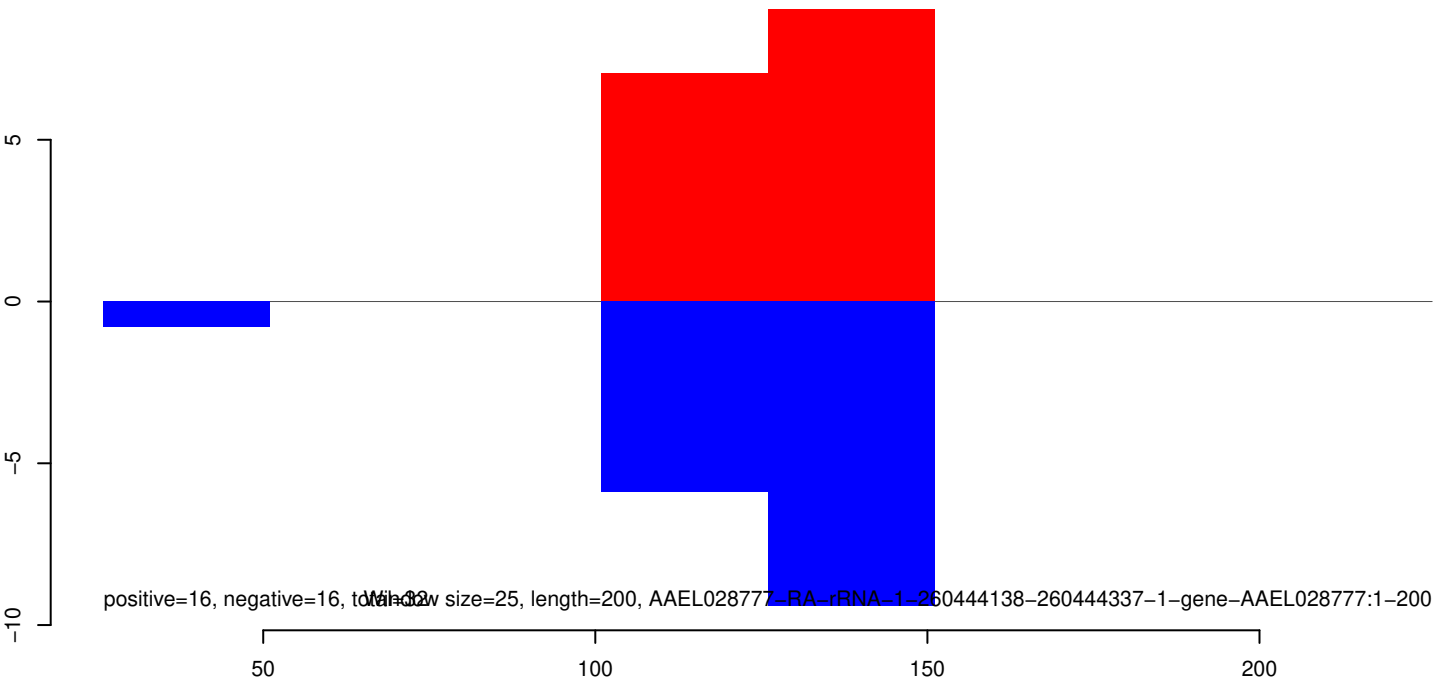
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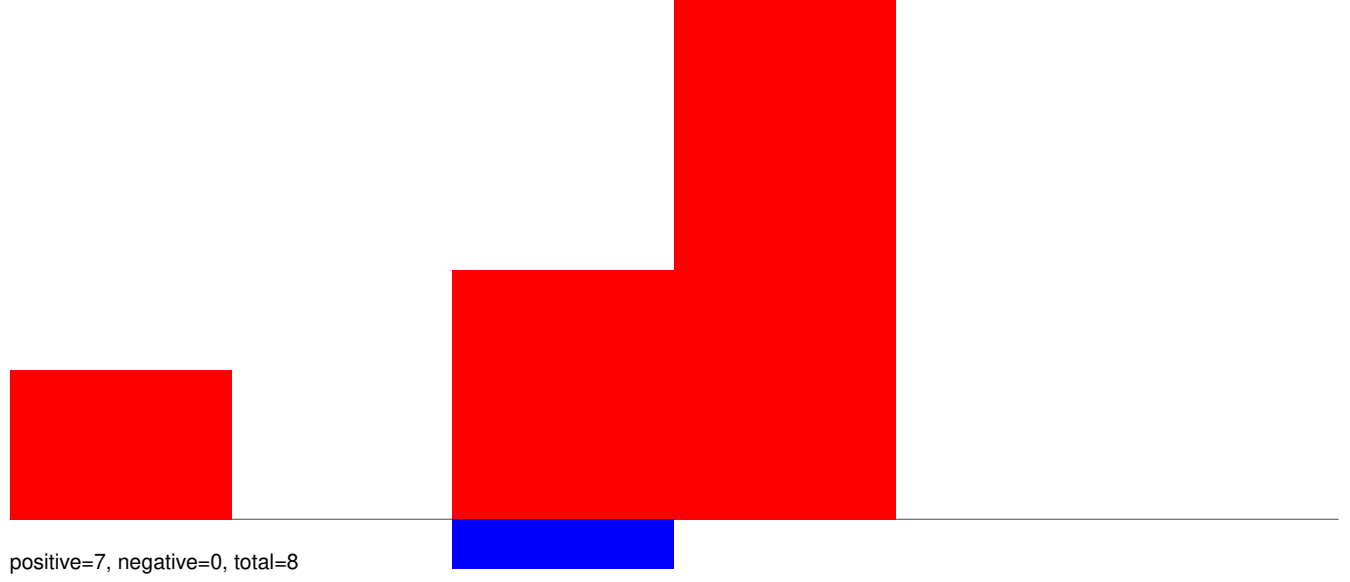
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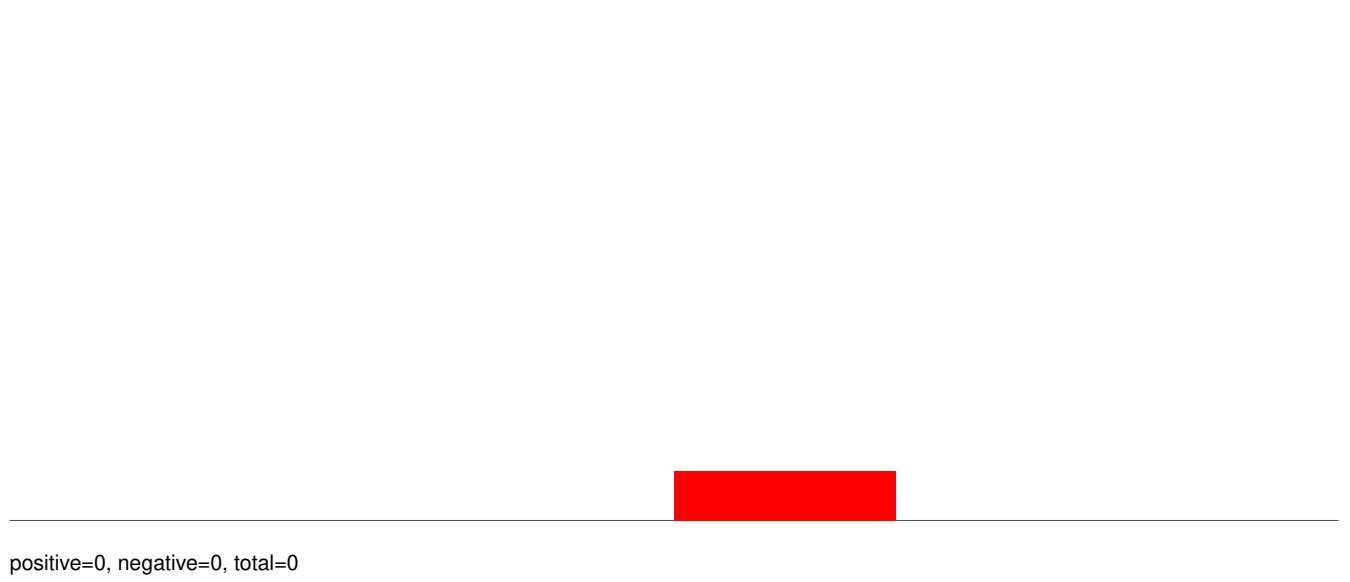
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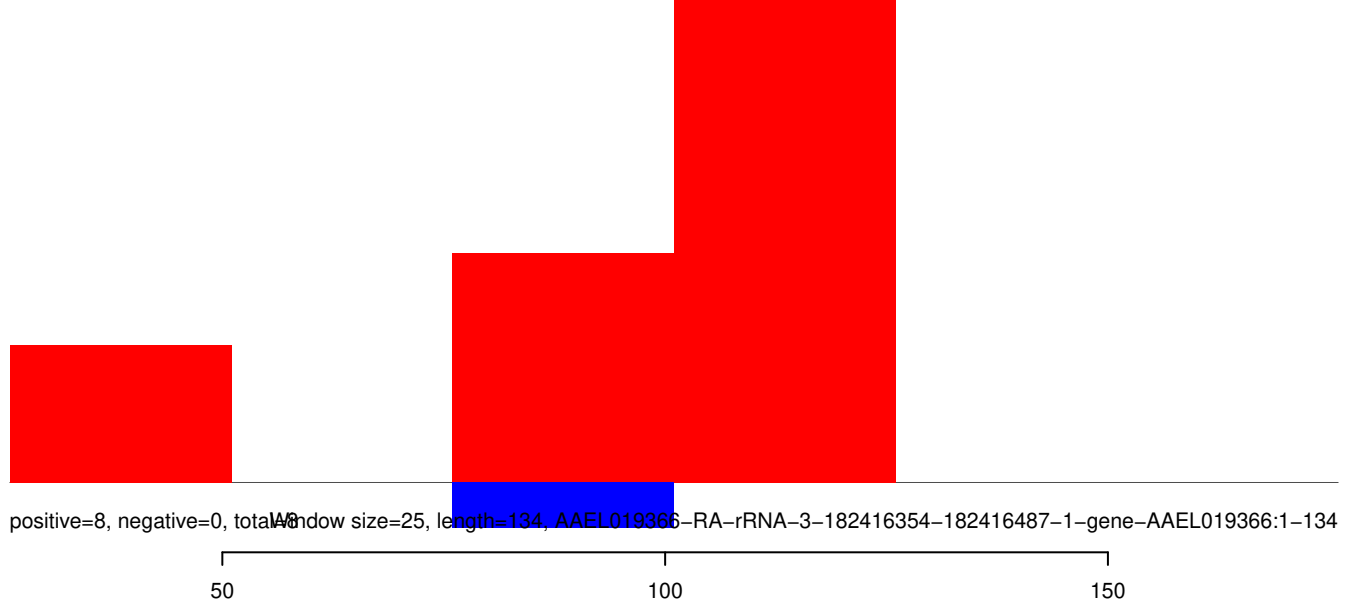
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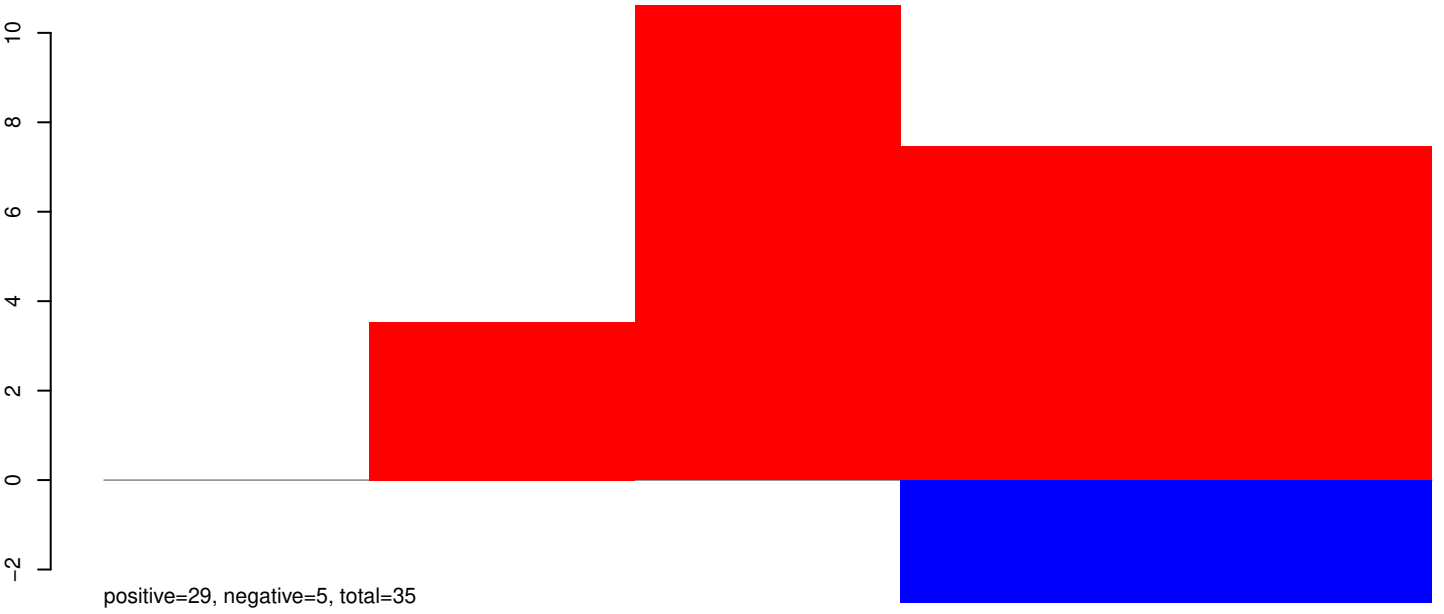
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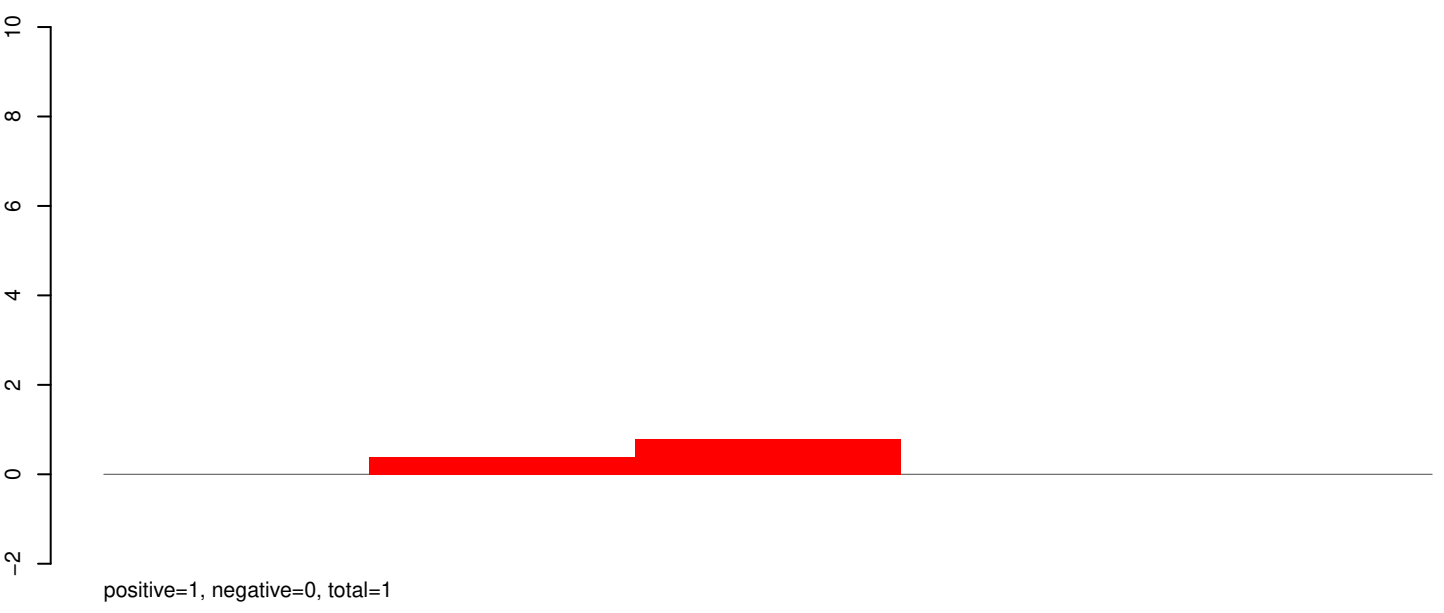
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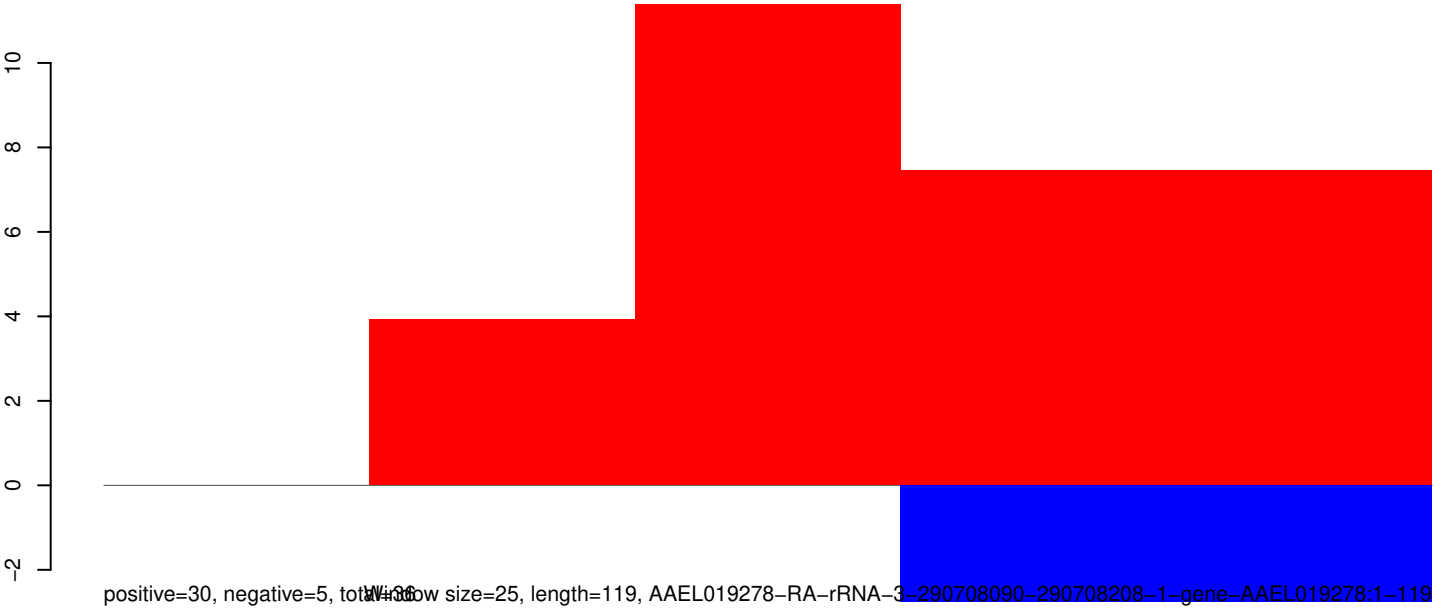
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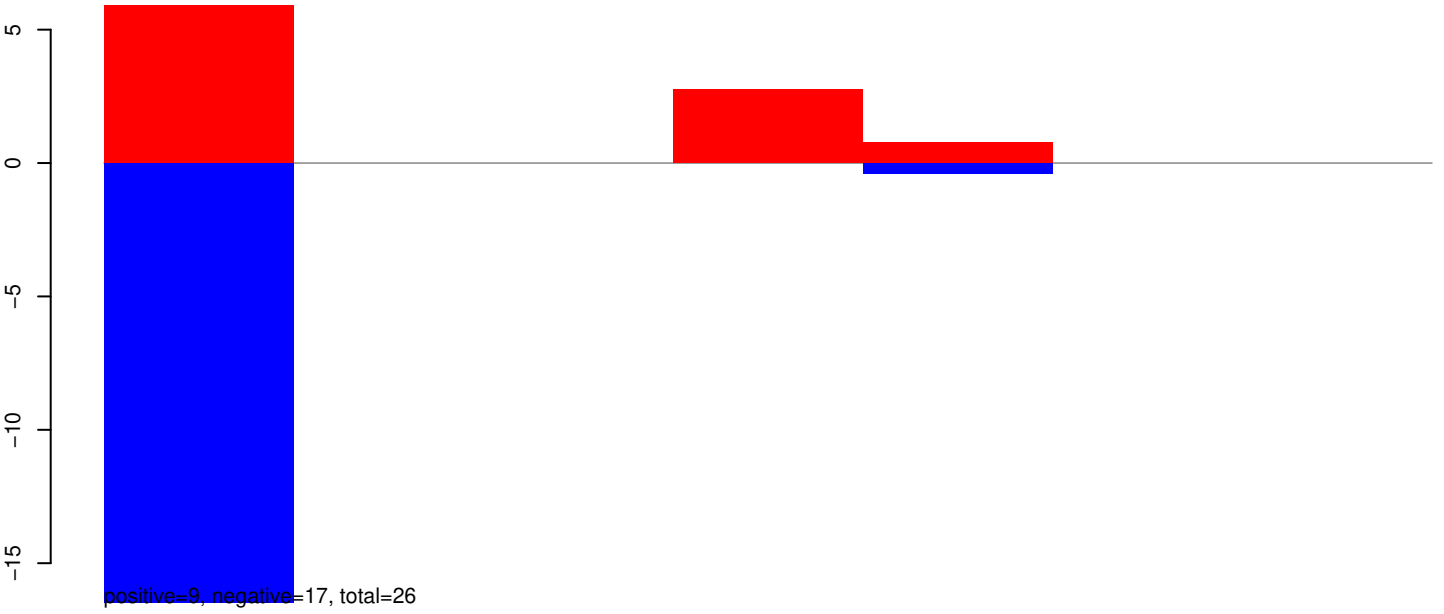
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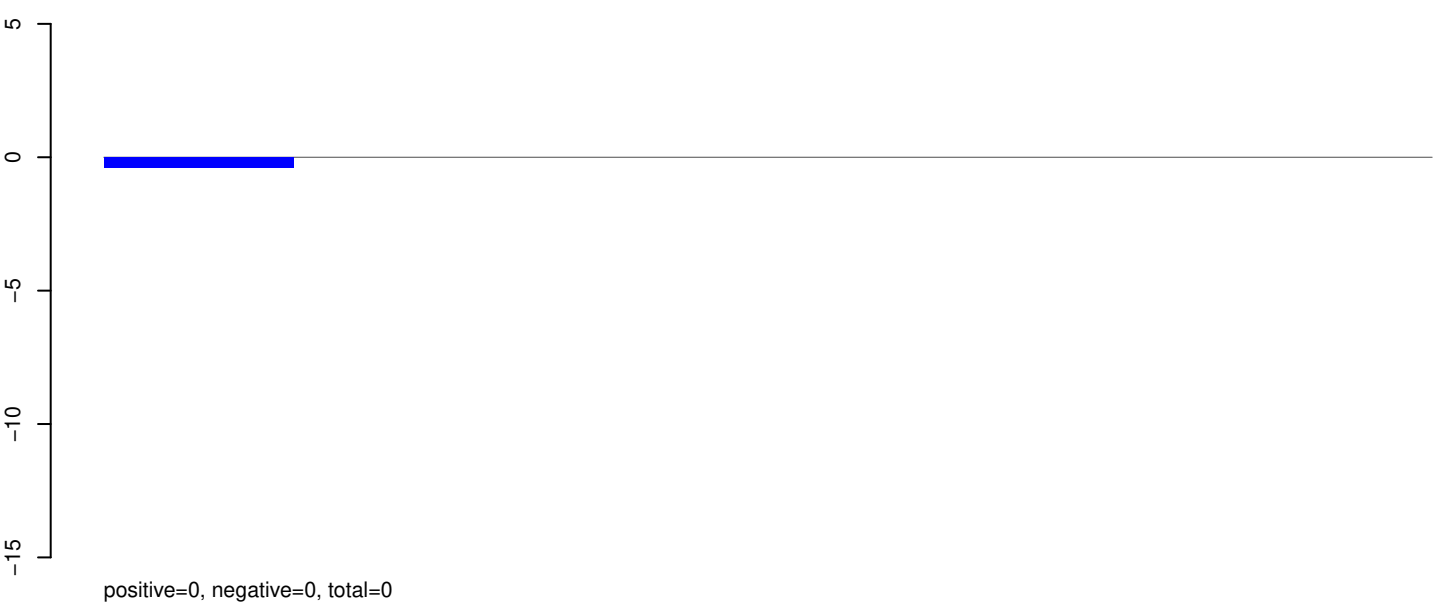
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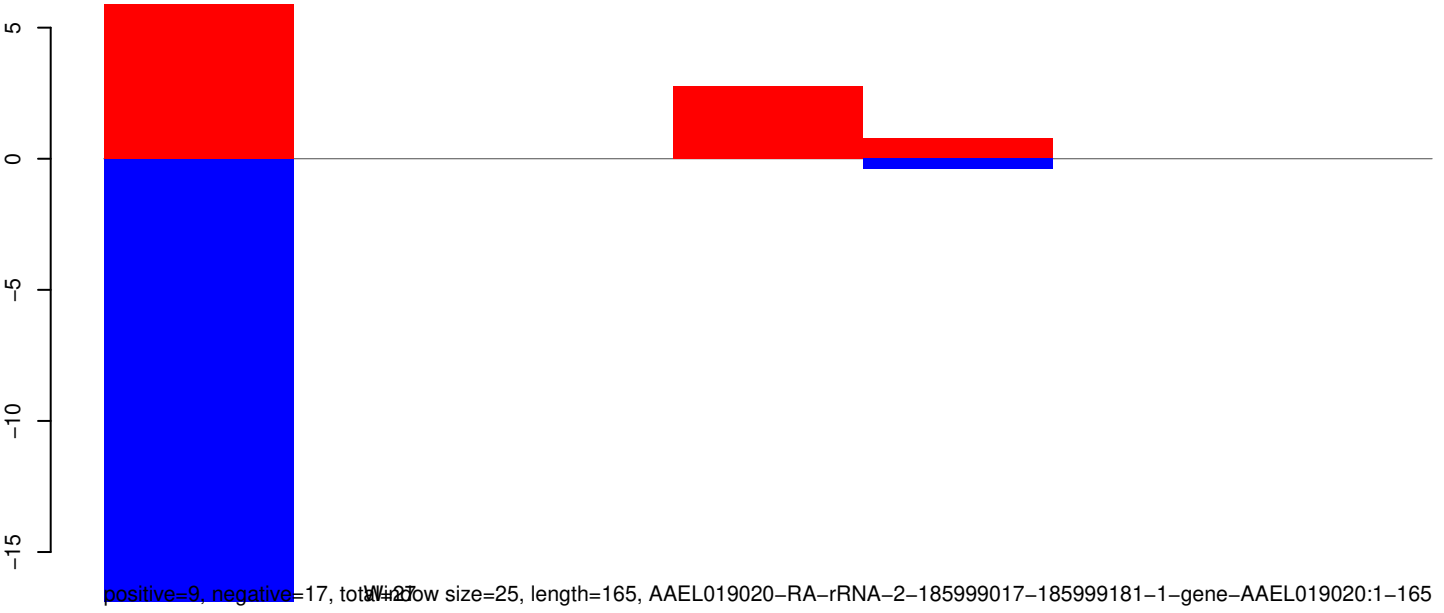
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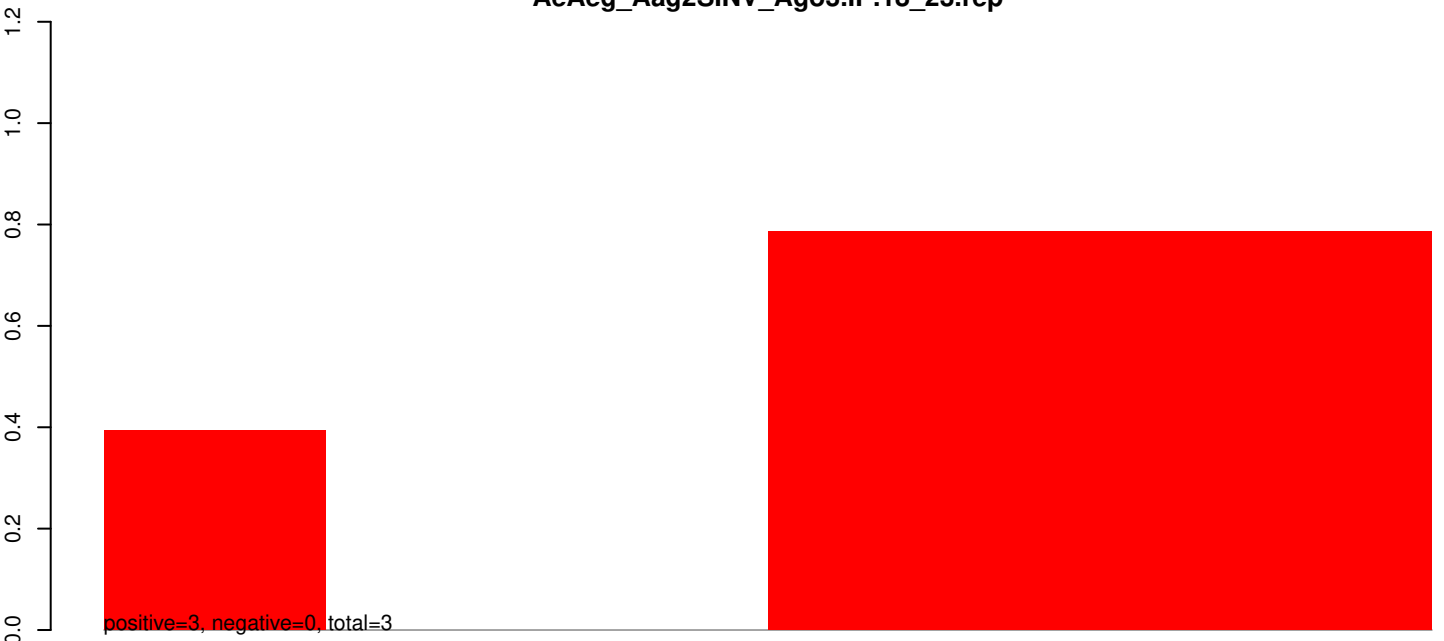


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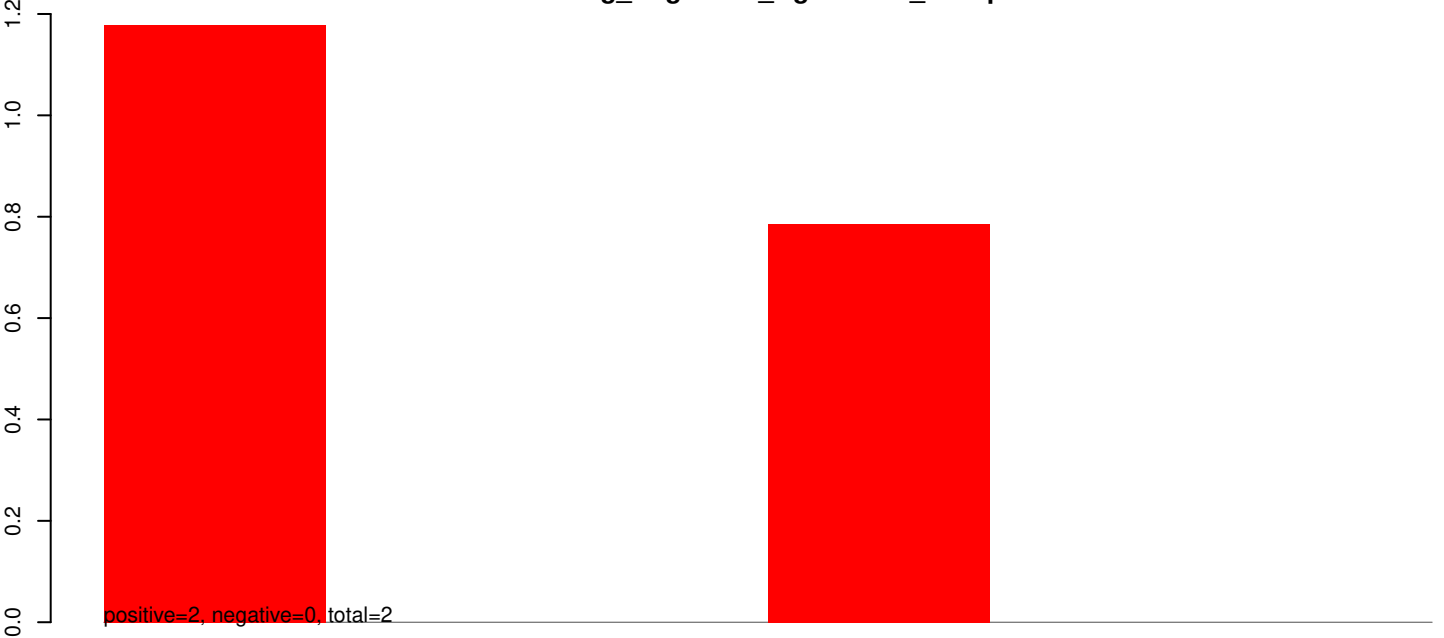


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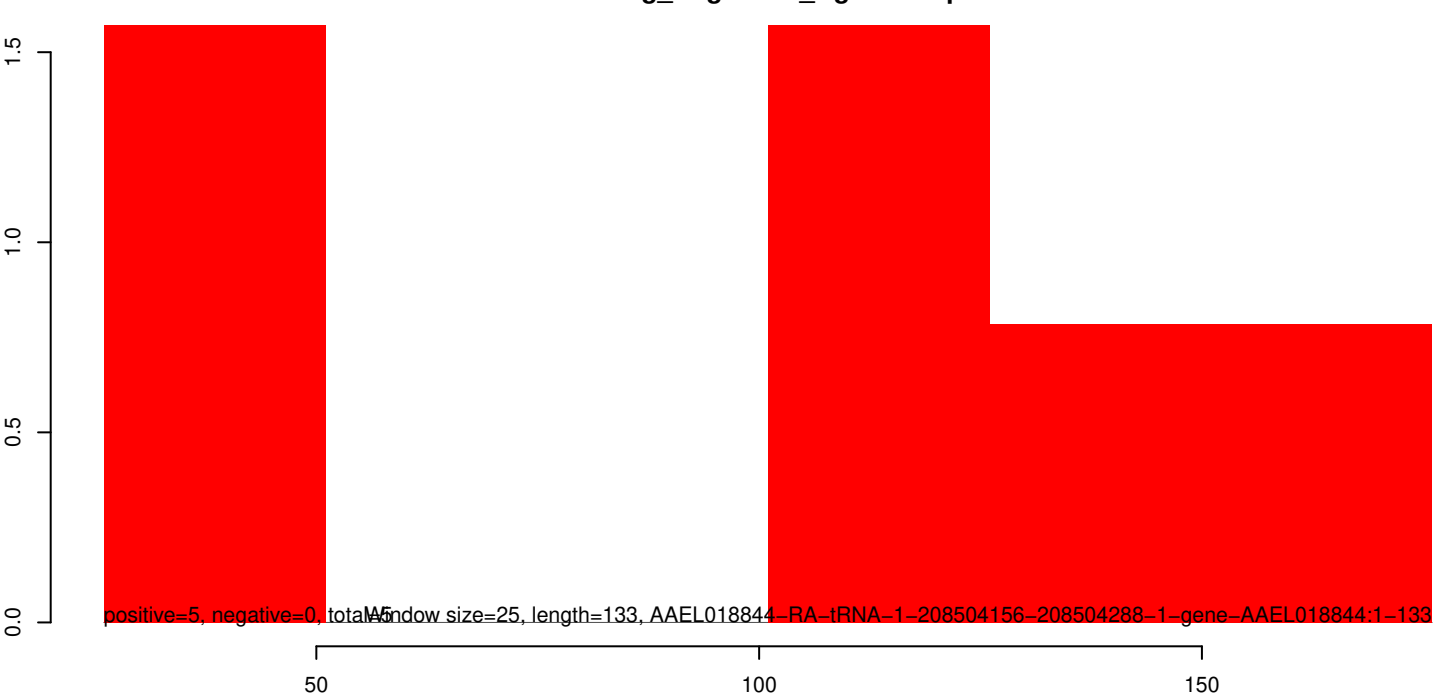
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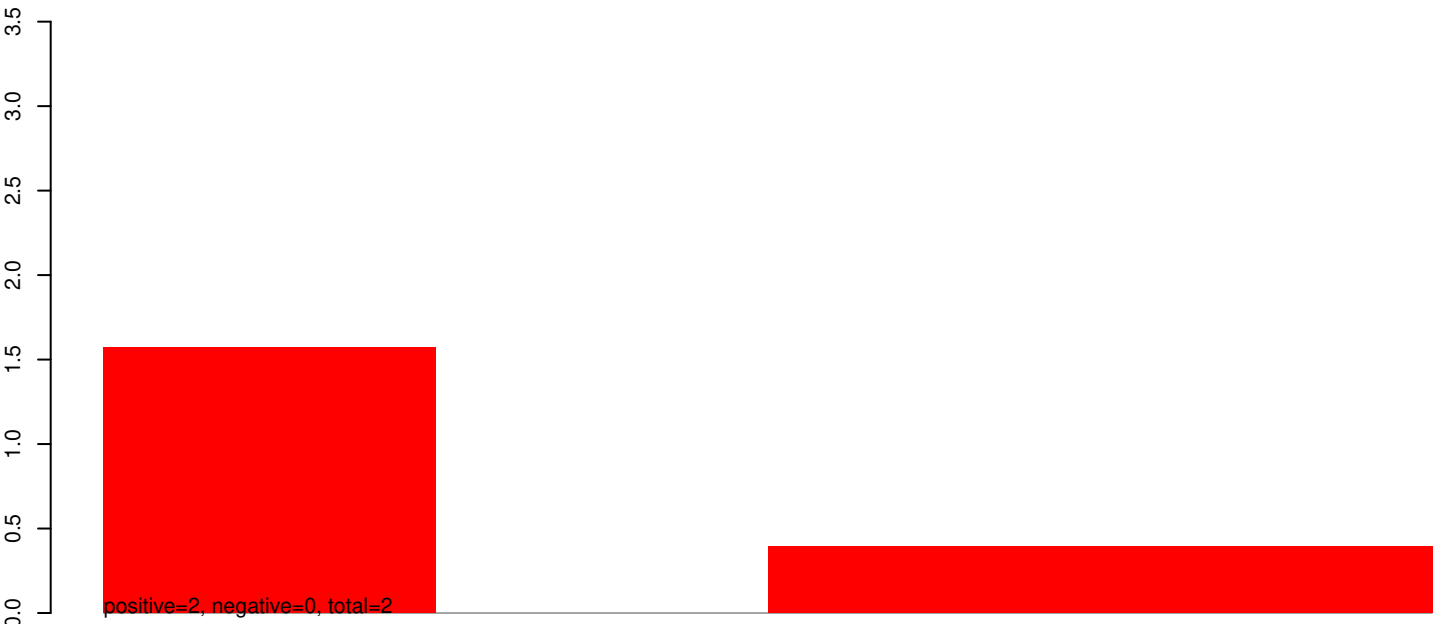
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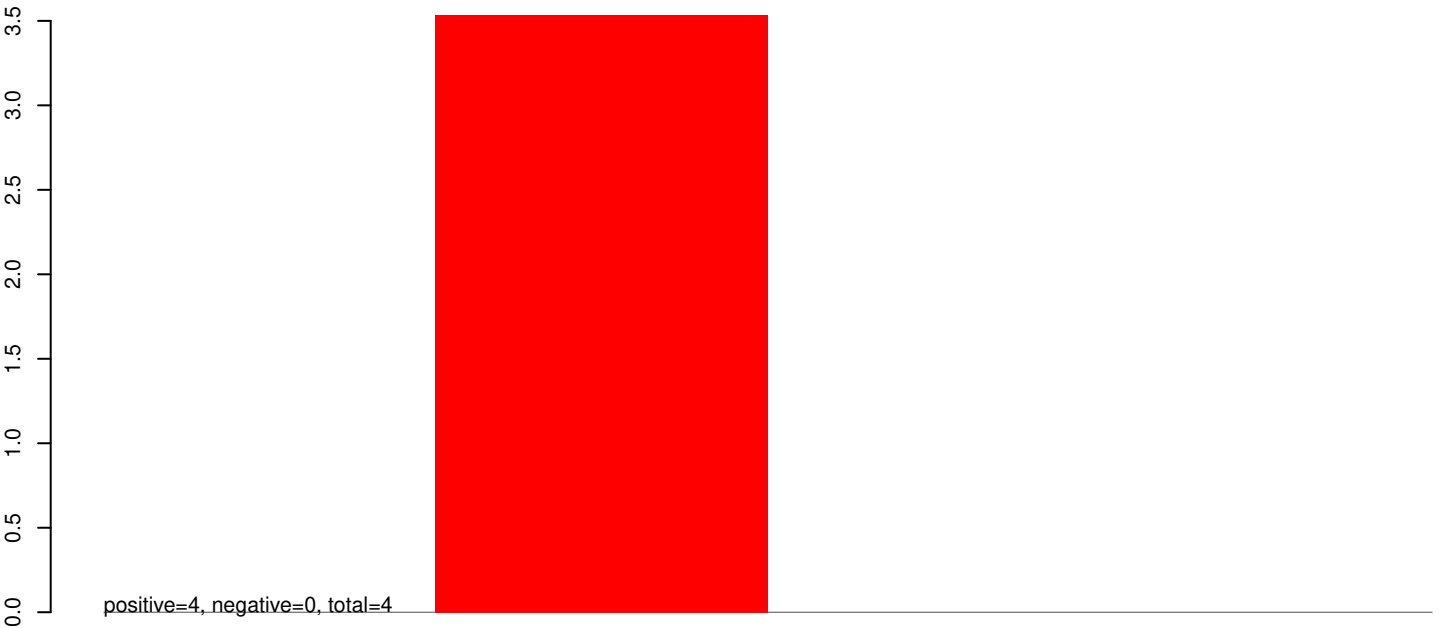
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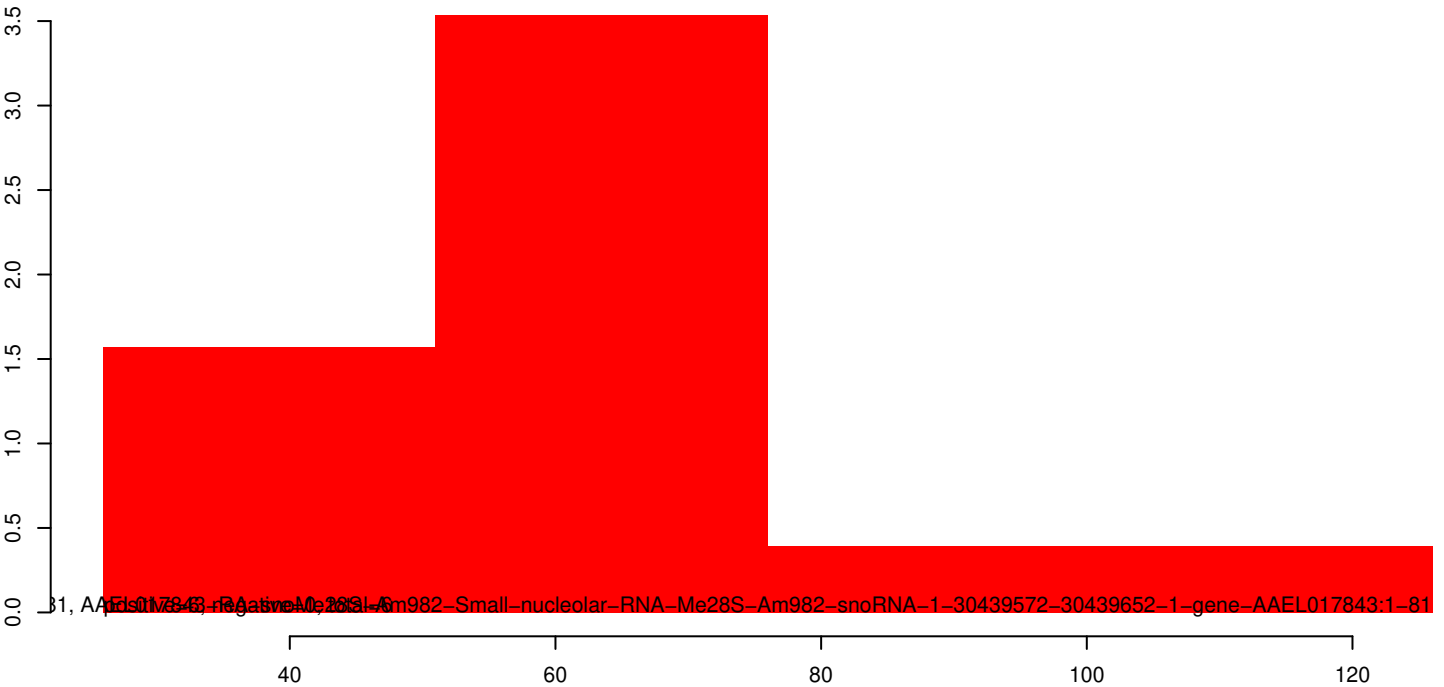
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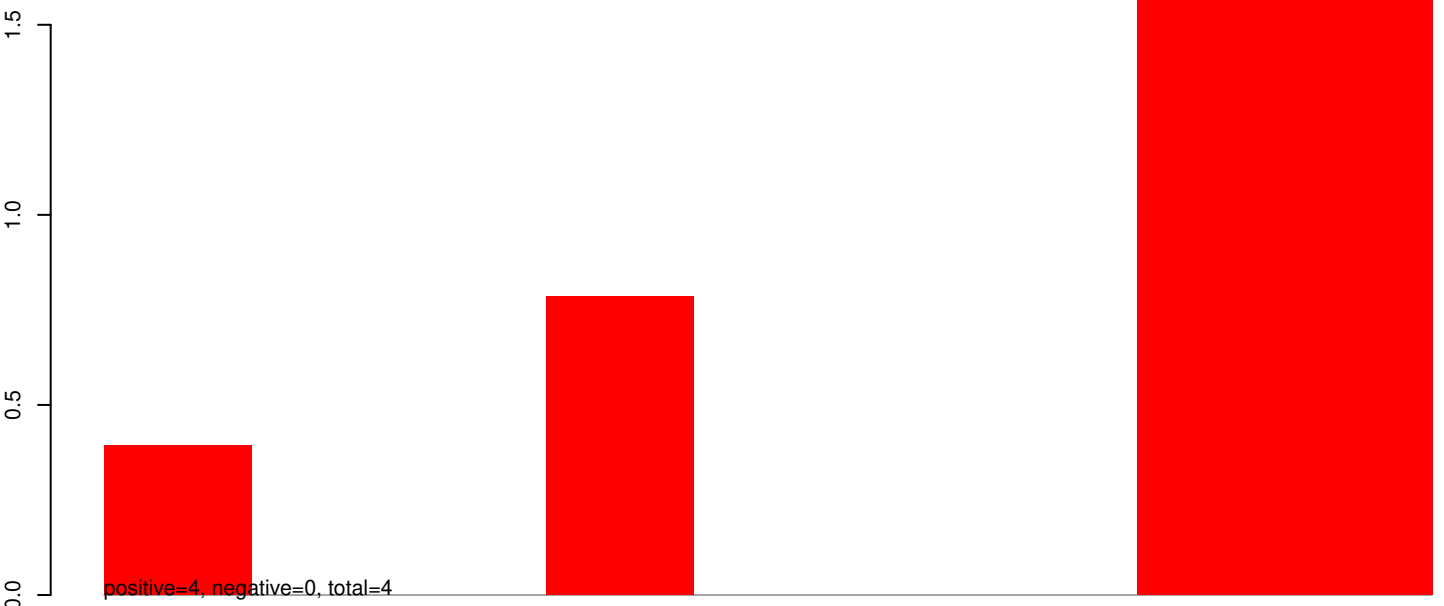
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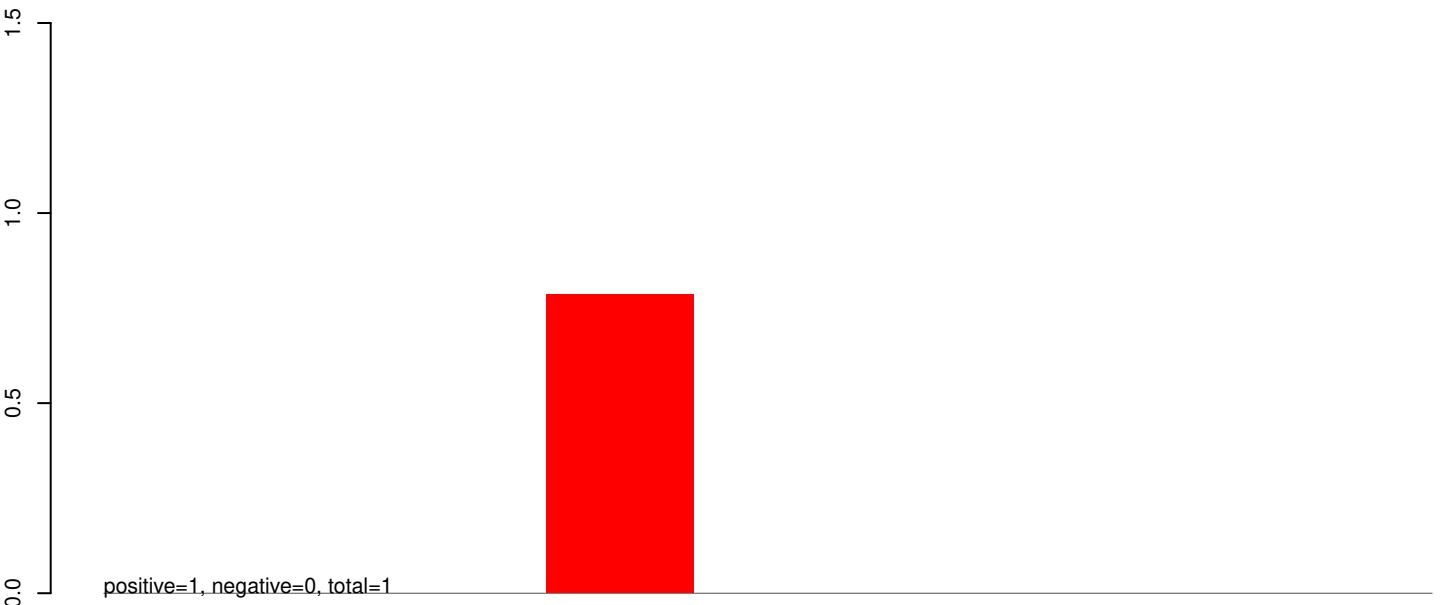
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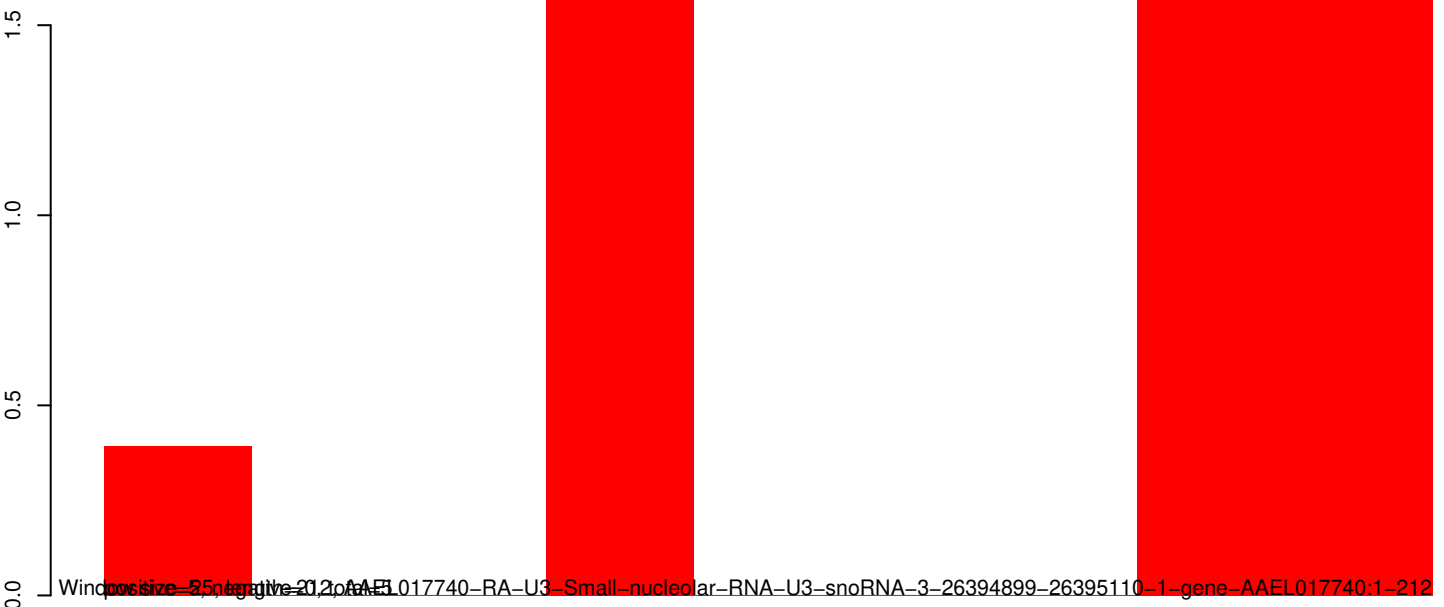
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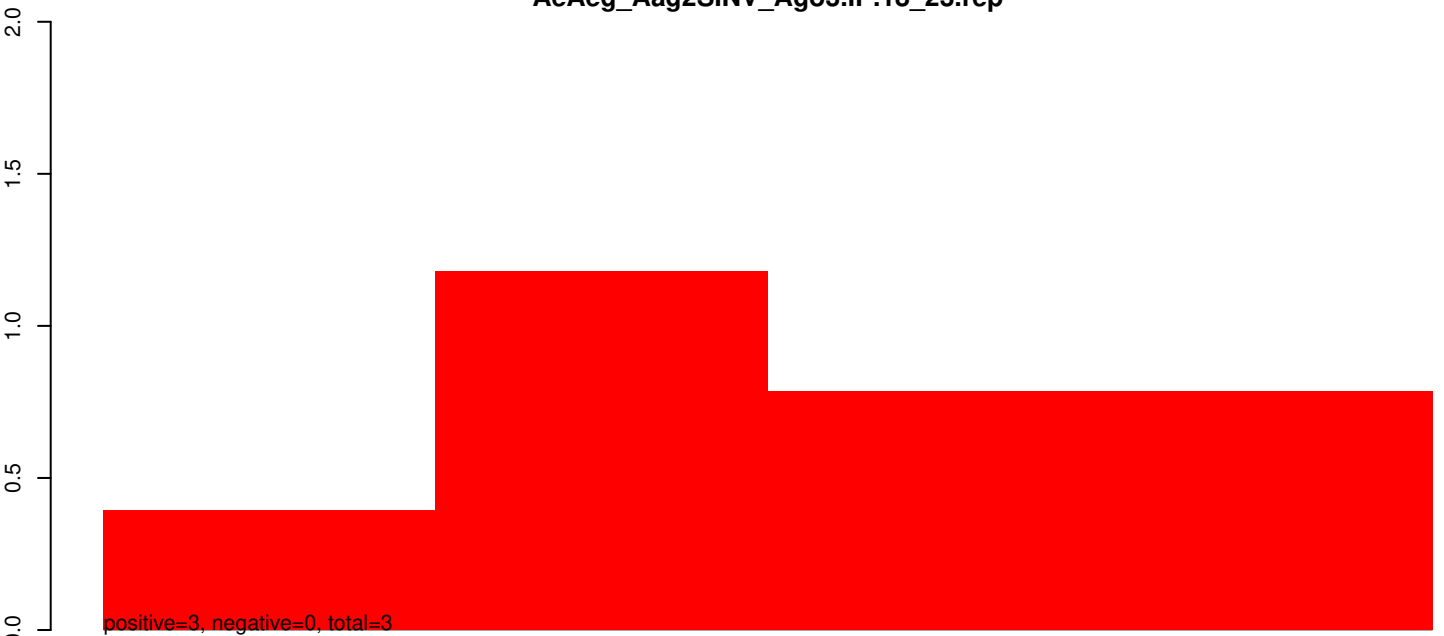
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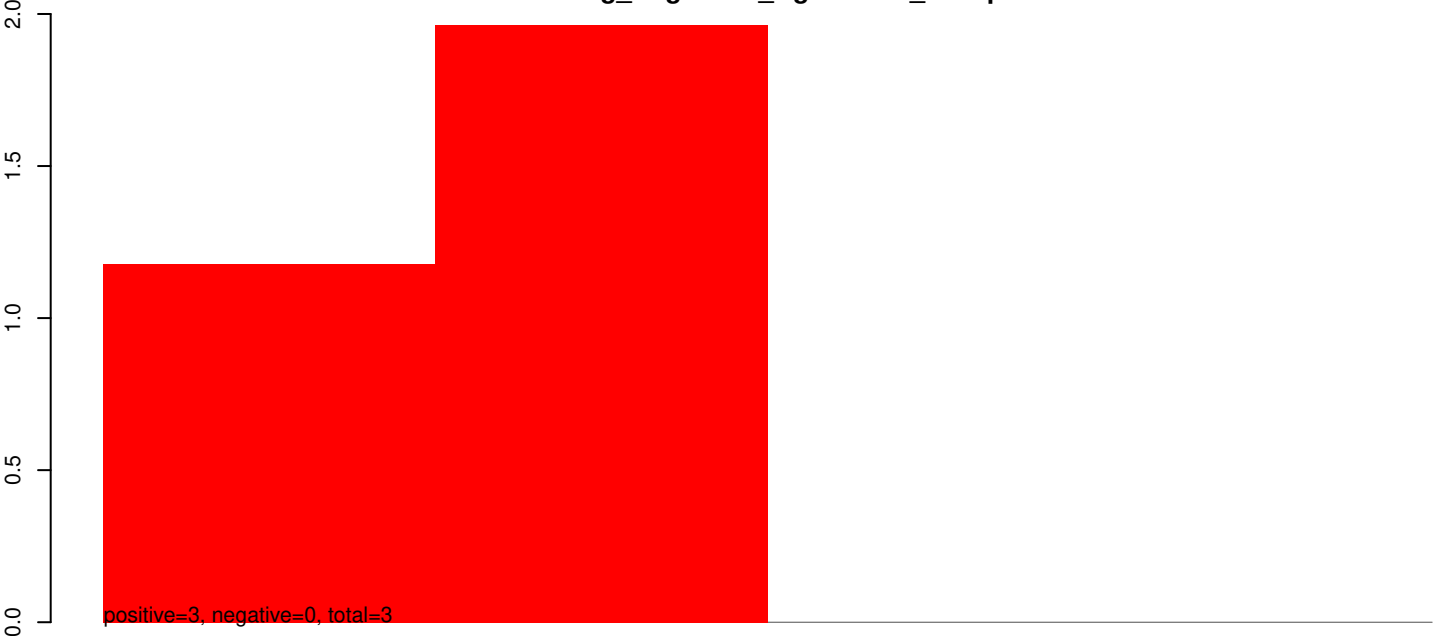
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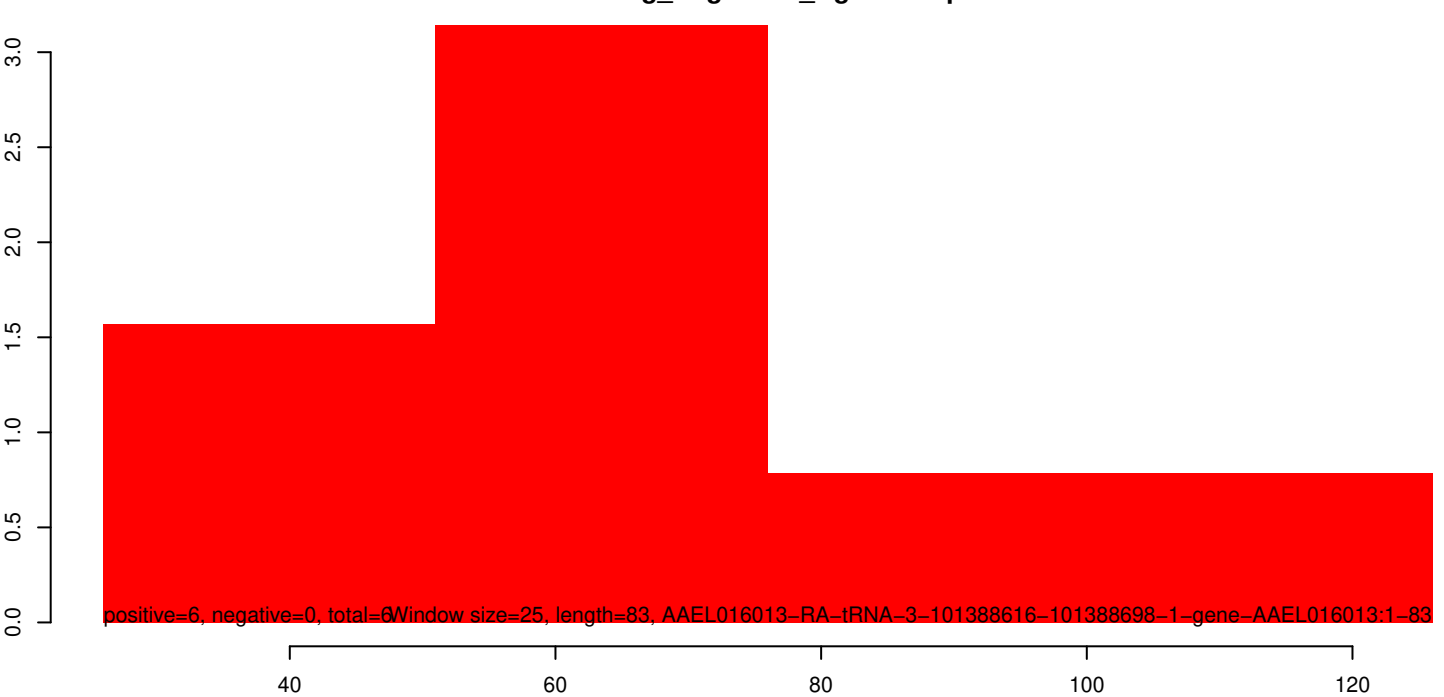
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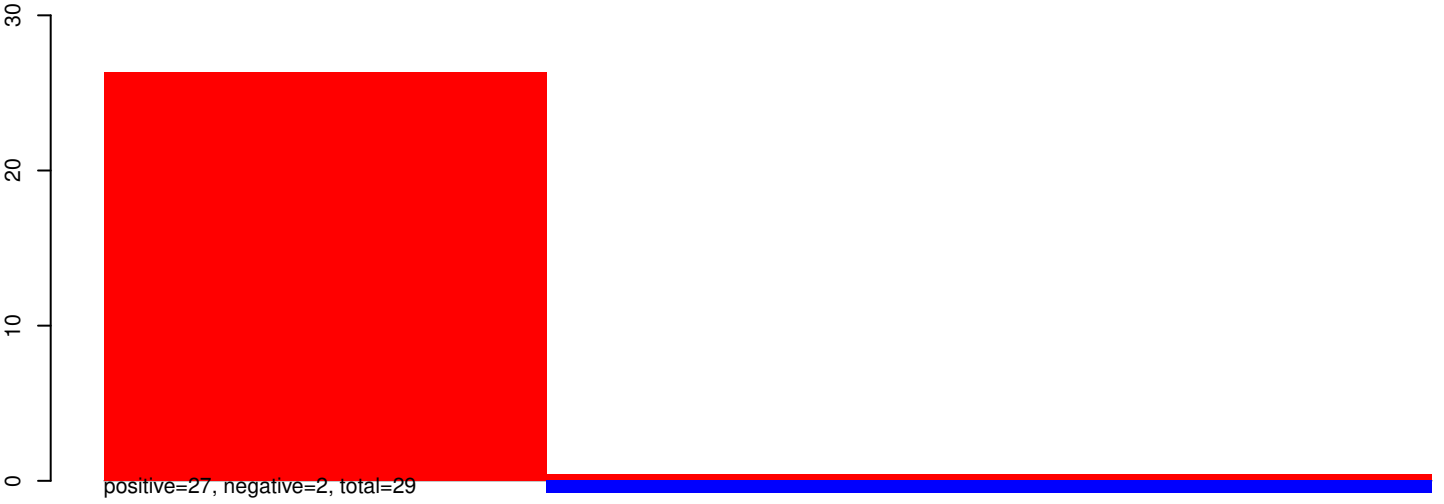
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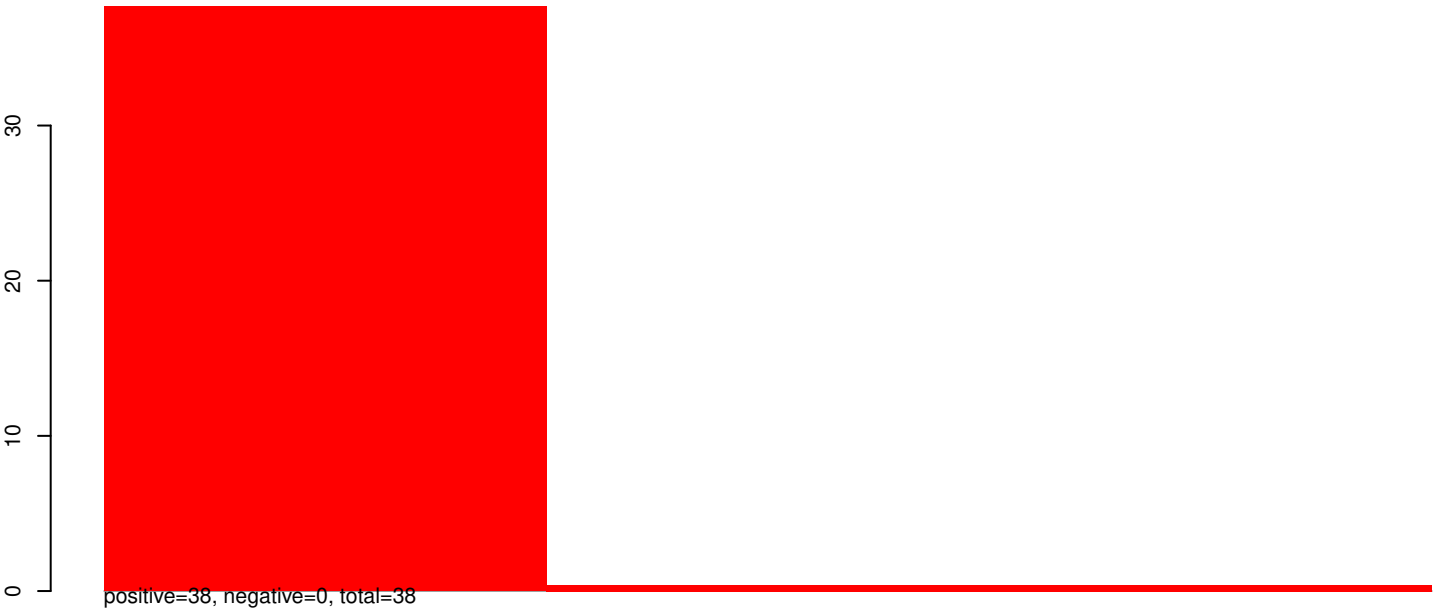
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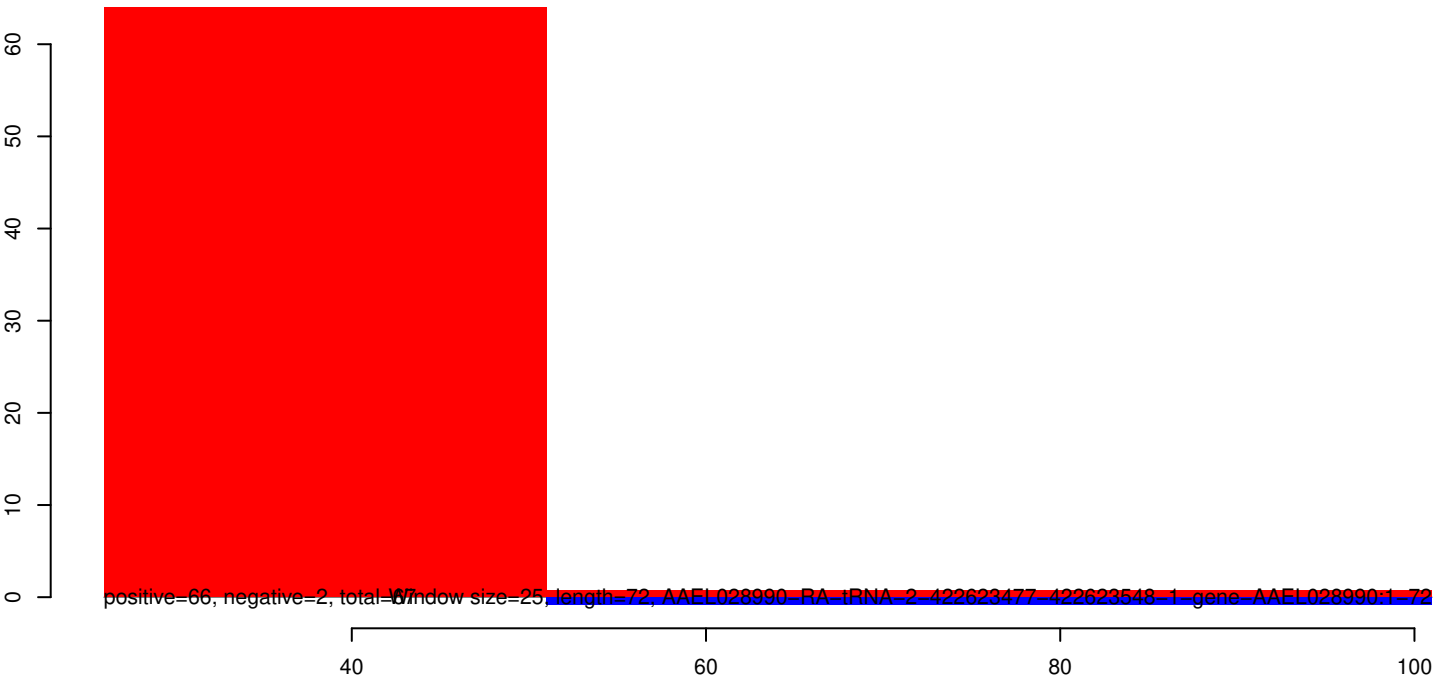
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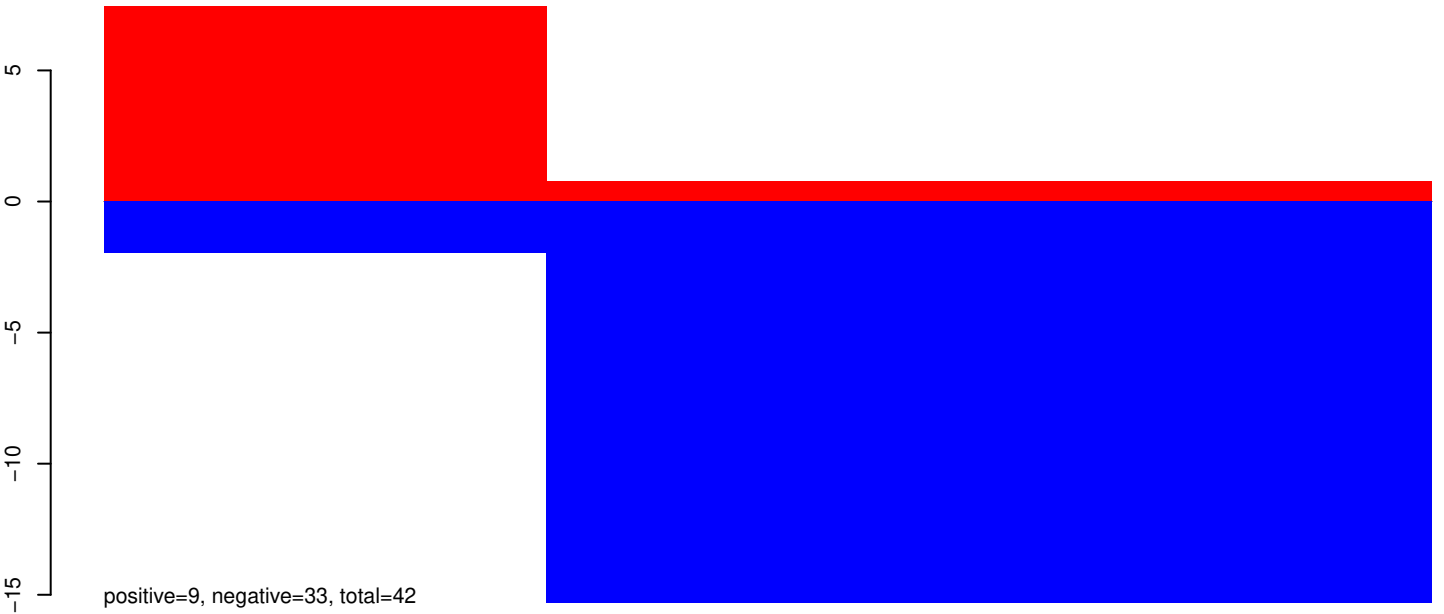
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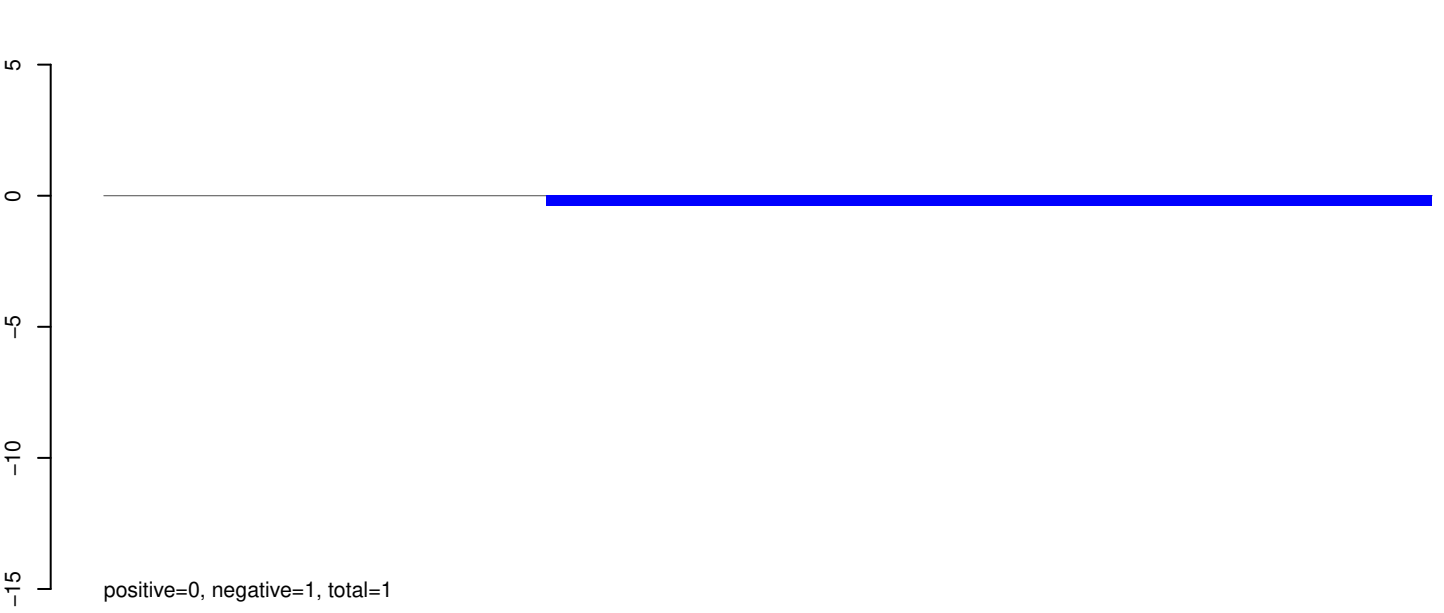
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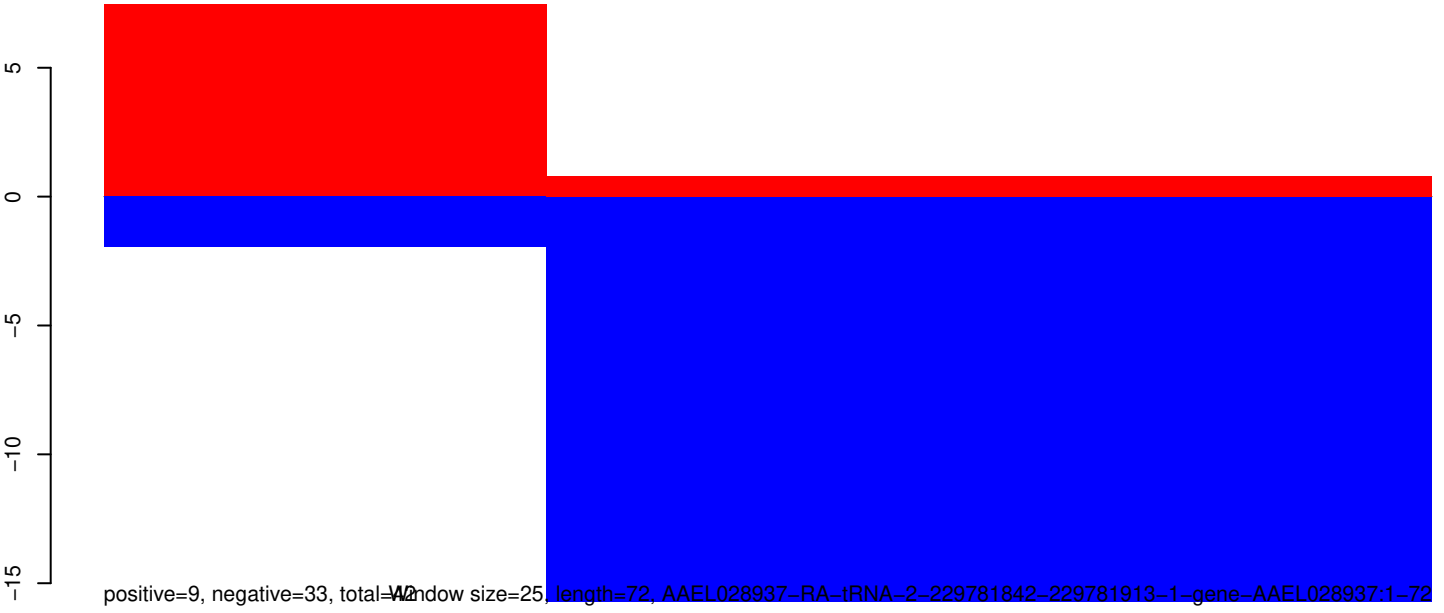
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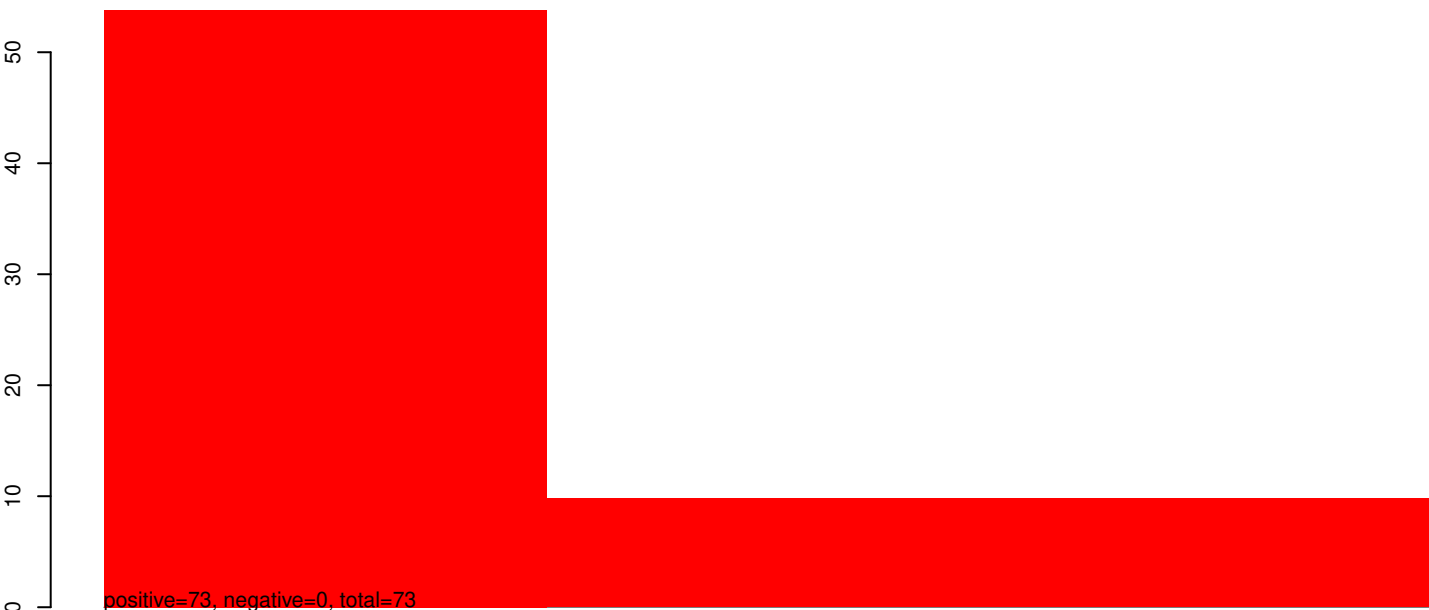


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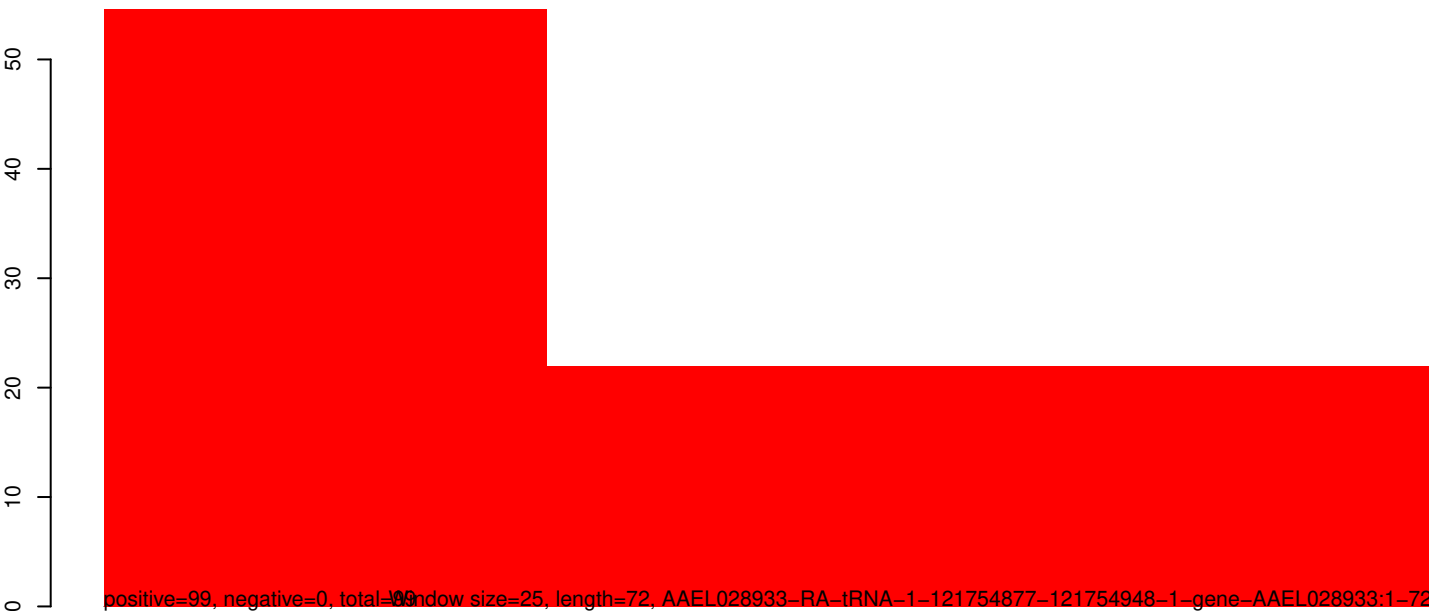
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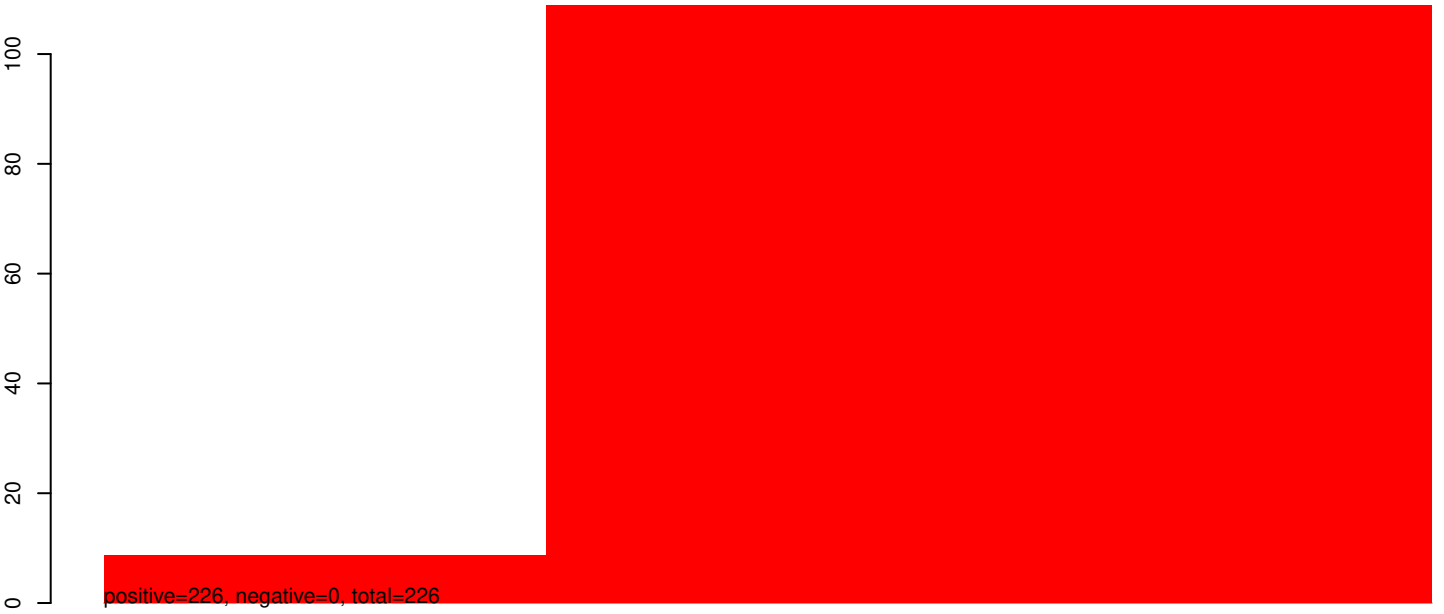
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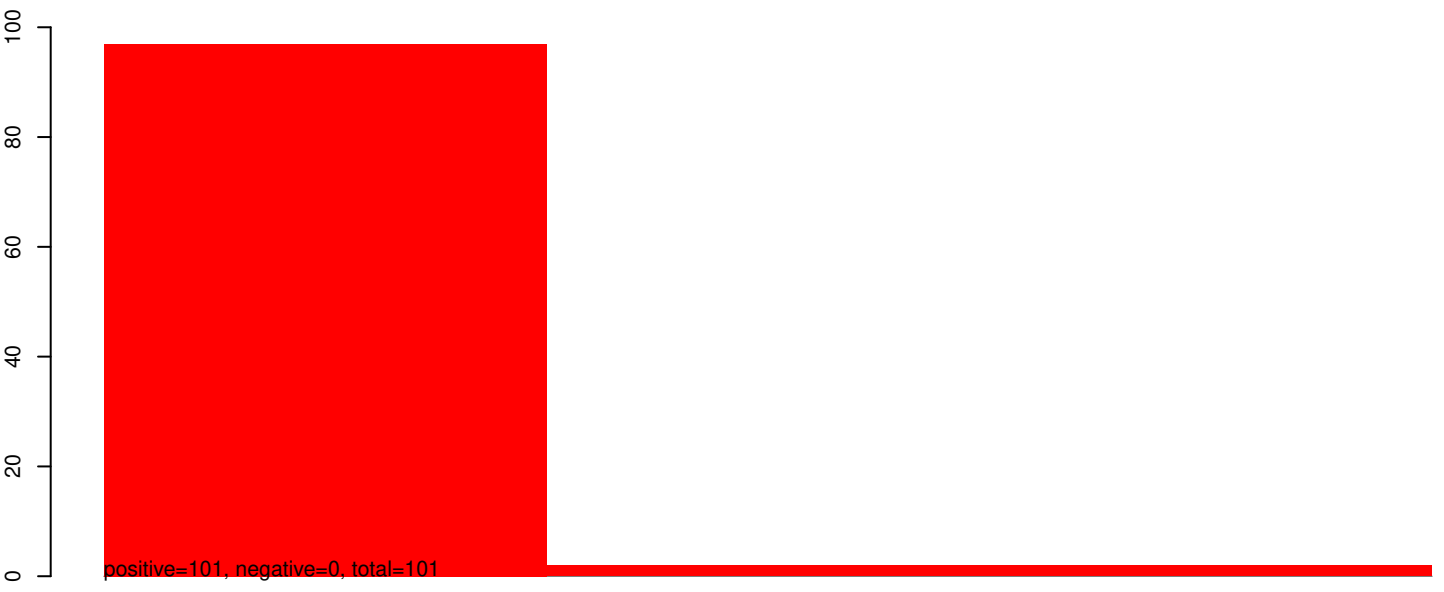
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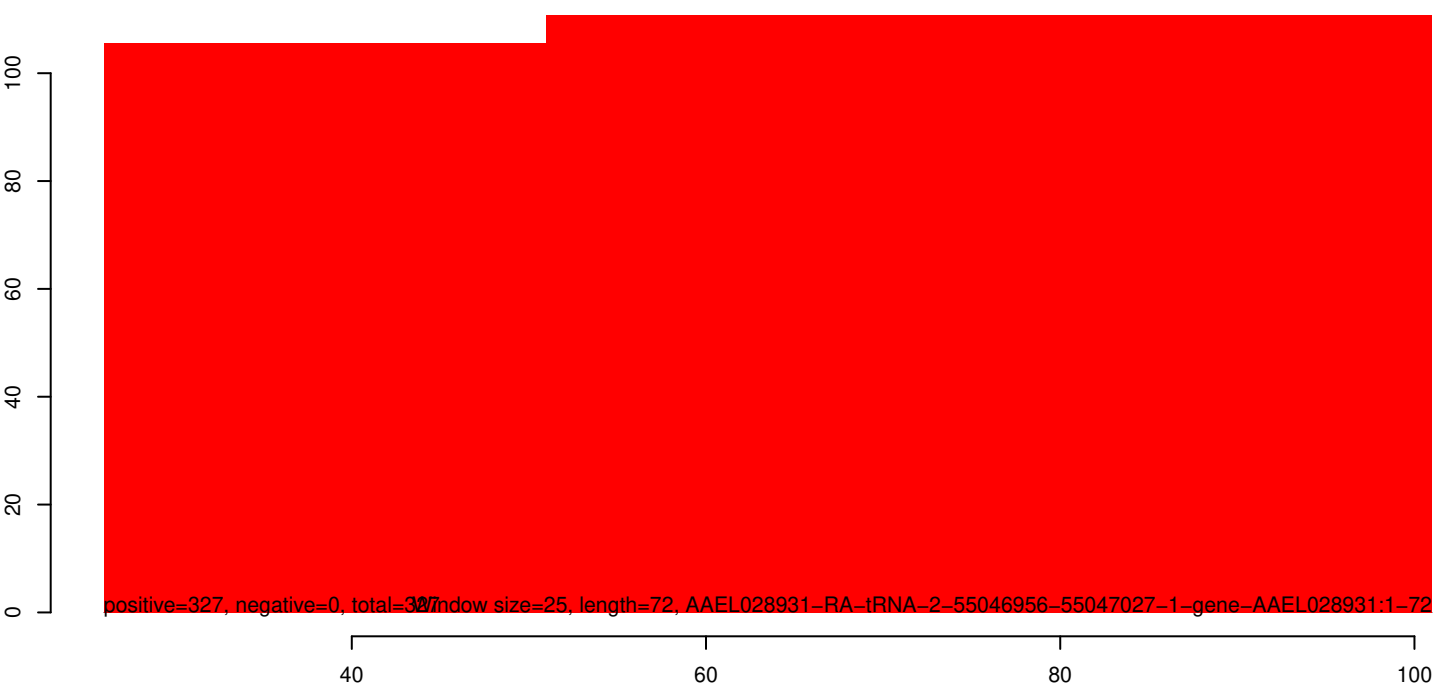
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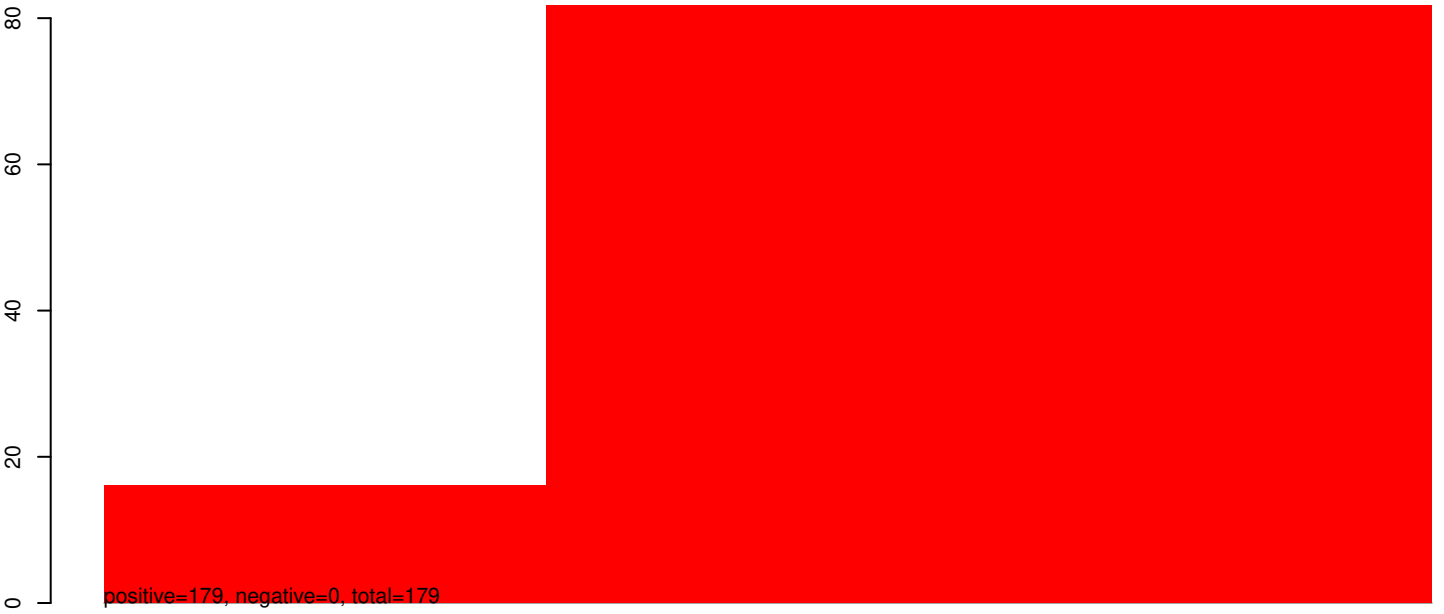
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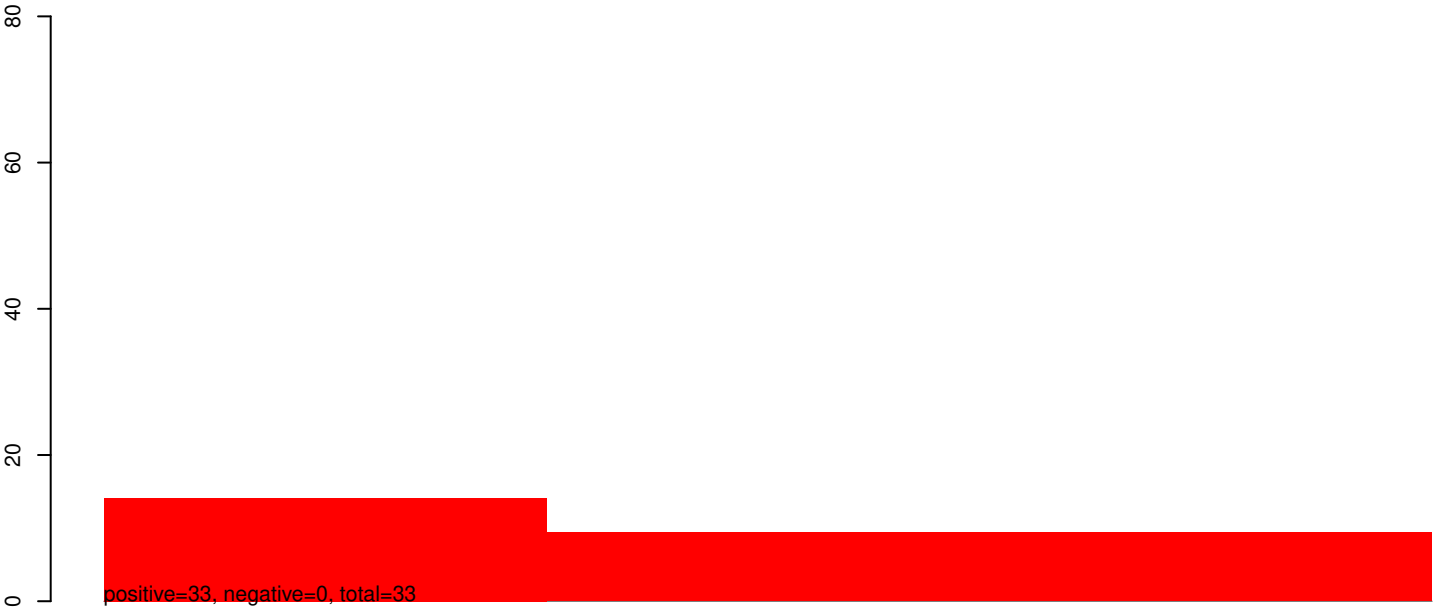
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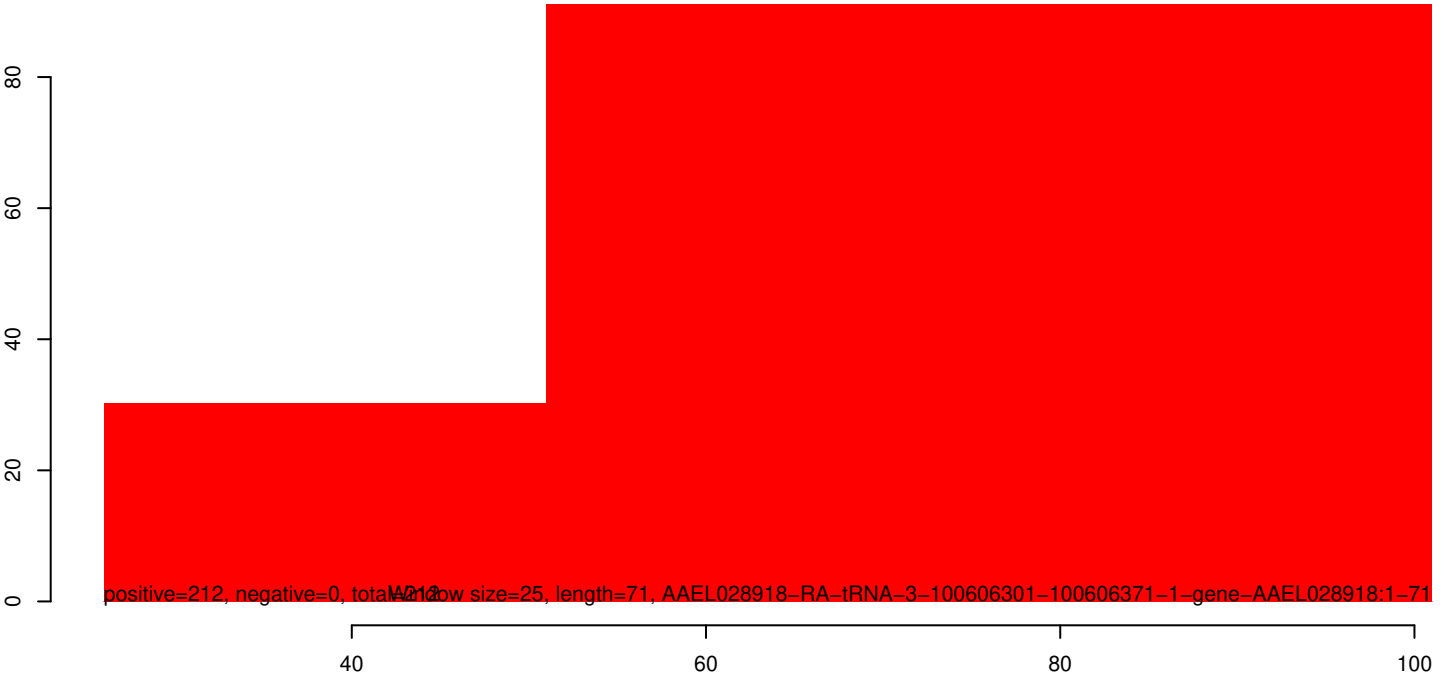
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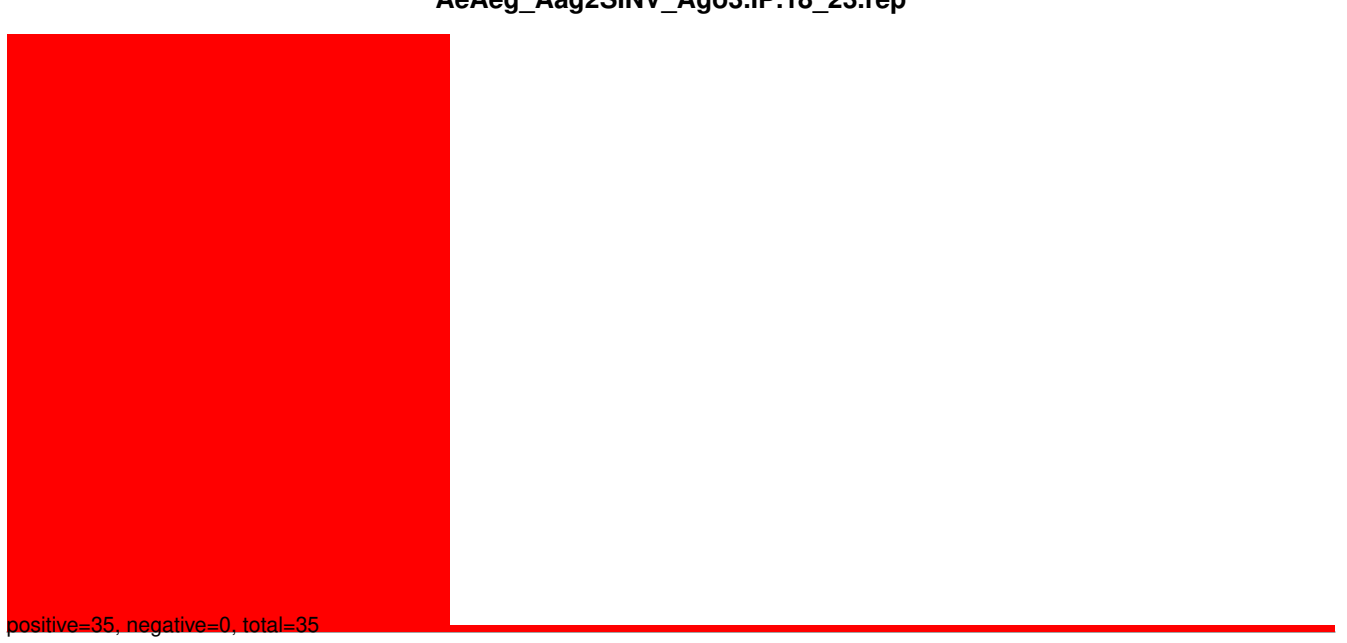
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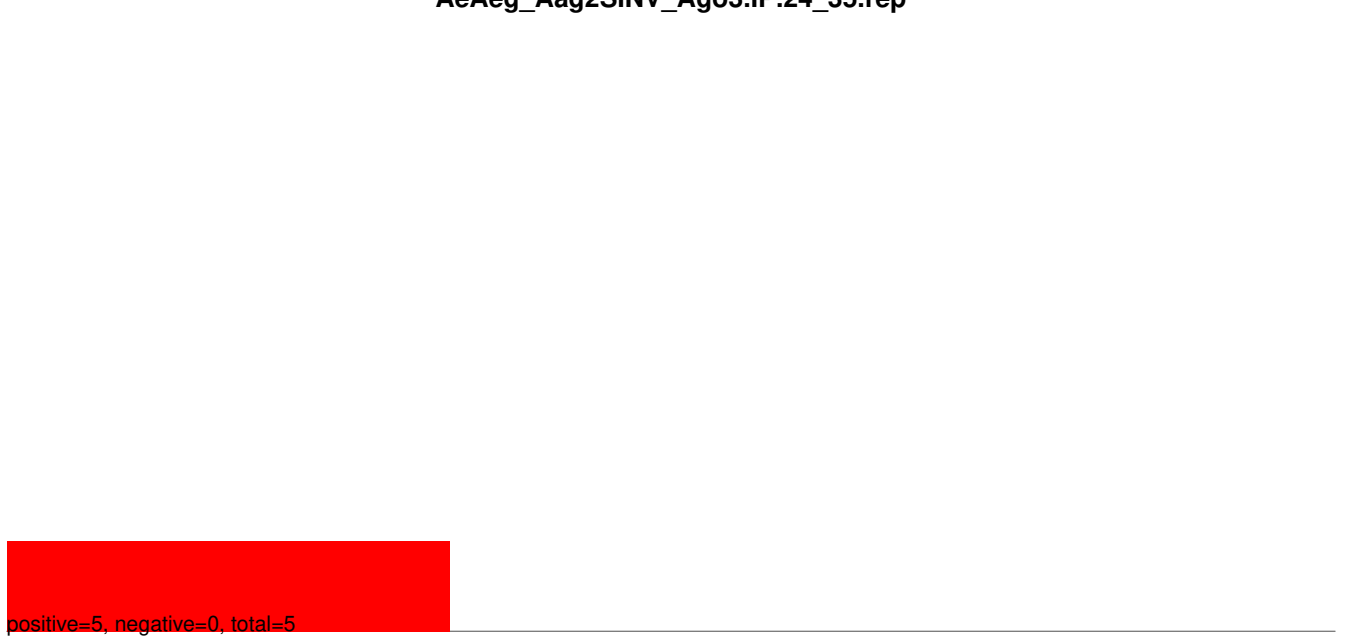
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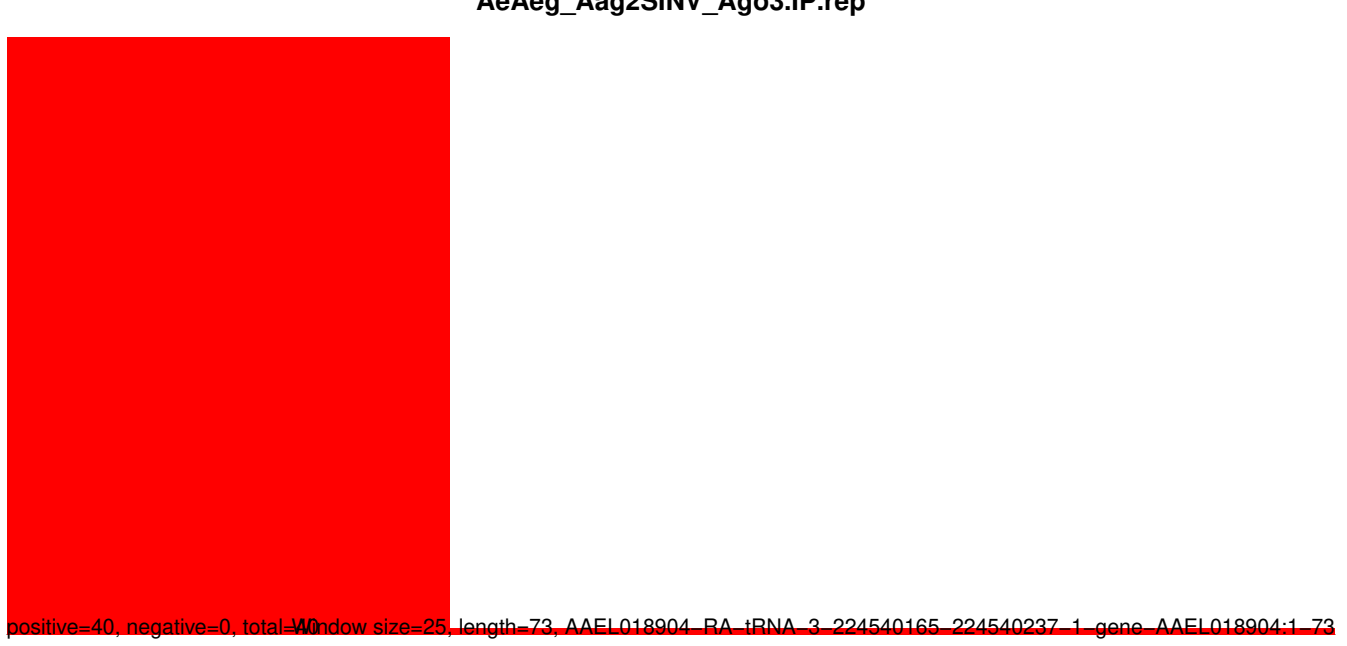
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AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep



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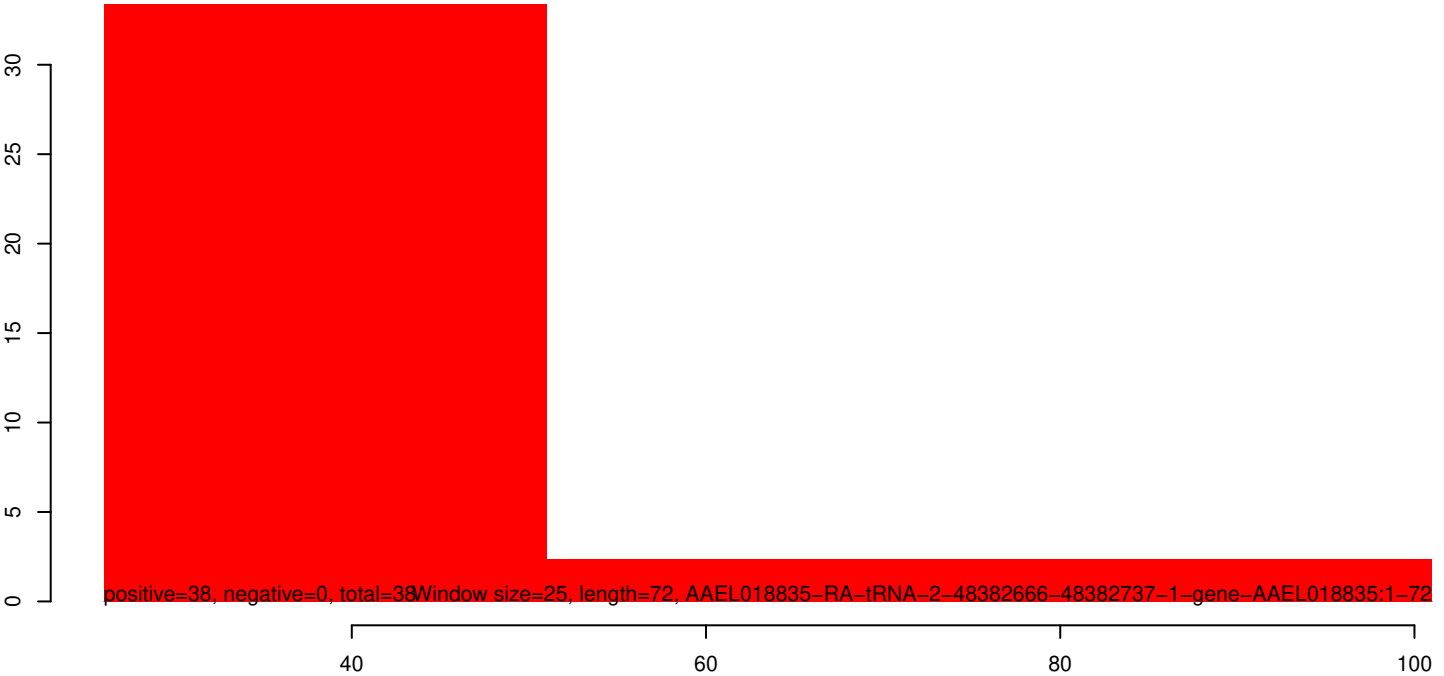
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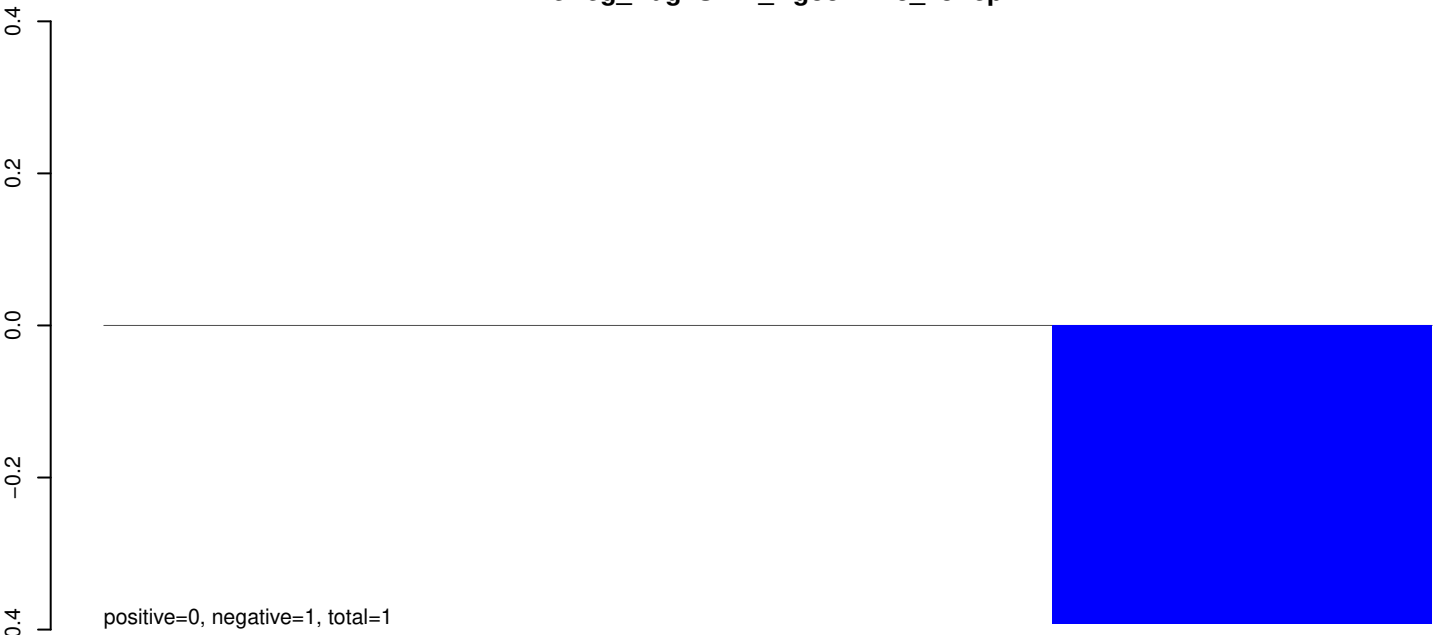
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AeAeg_Aag2SINV_Ago3.IP.rep



AeAeg_Aag2SINV_Ago3.IP.18_23.rep



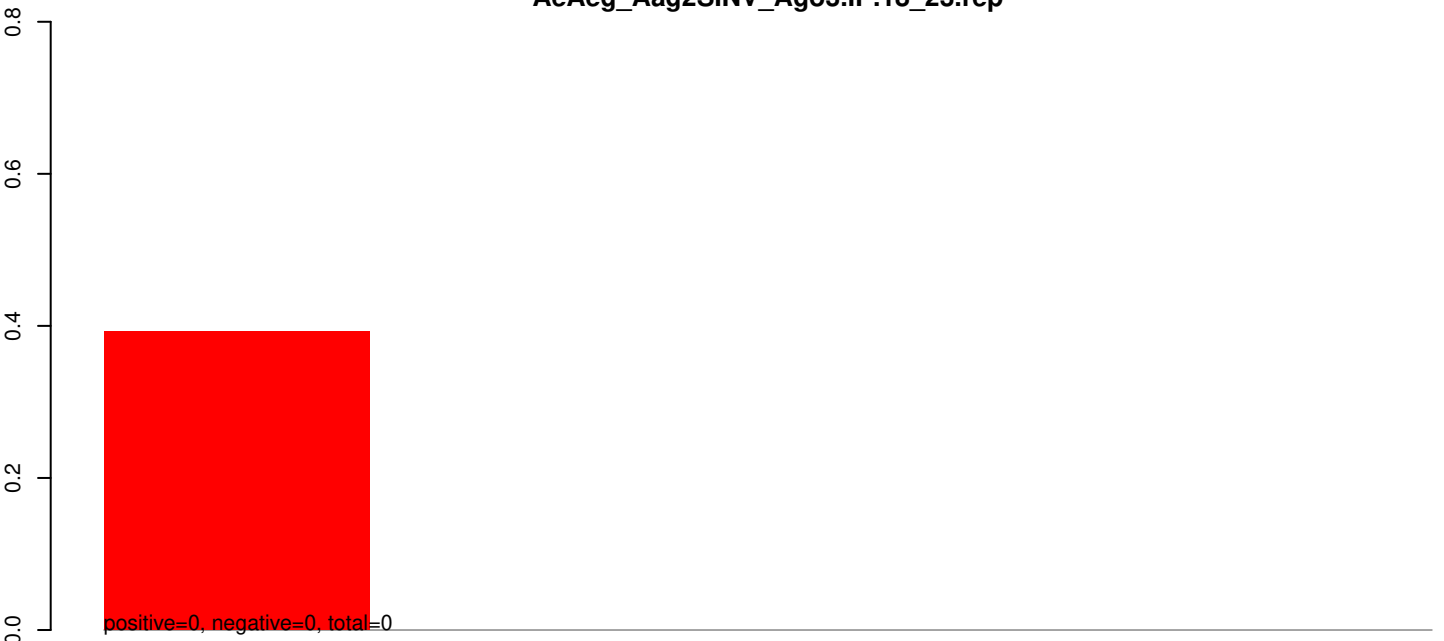
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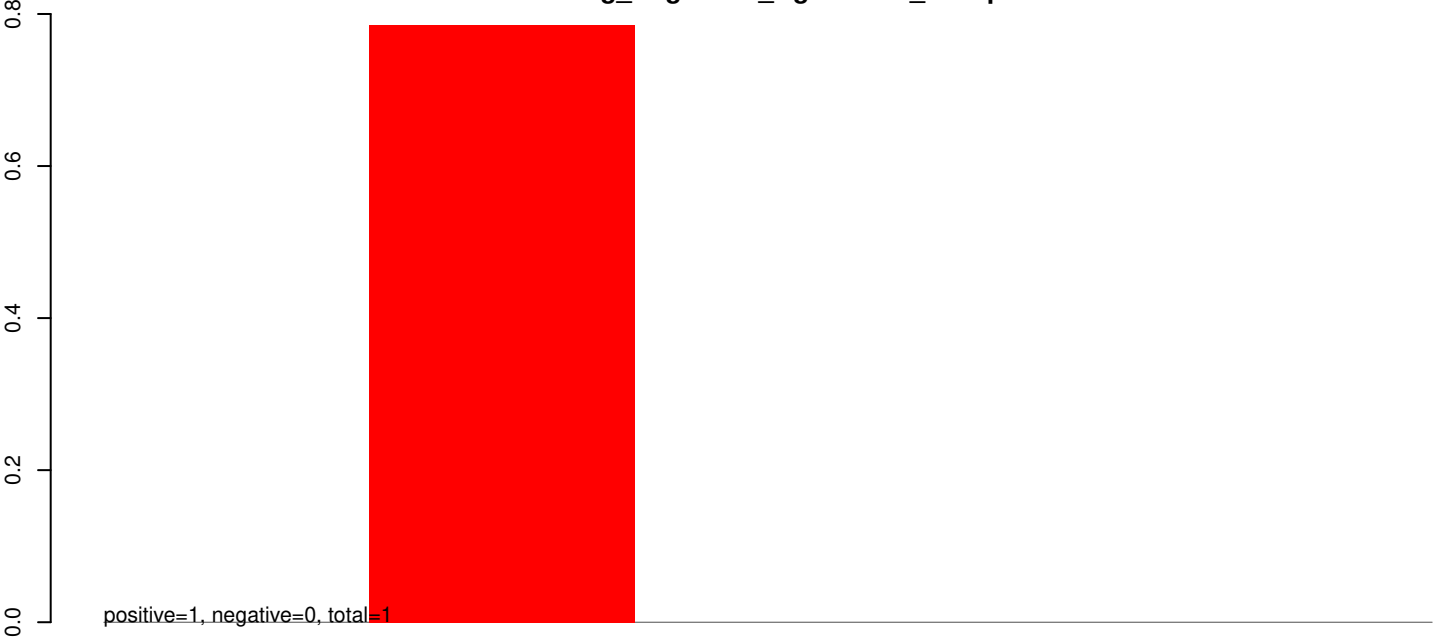
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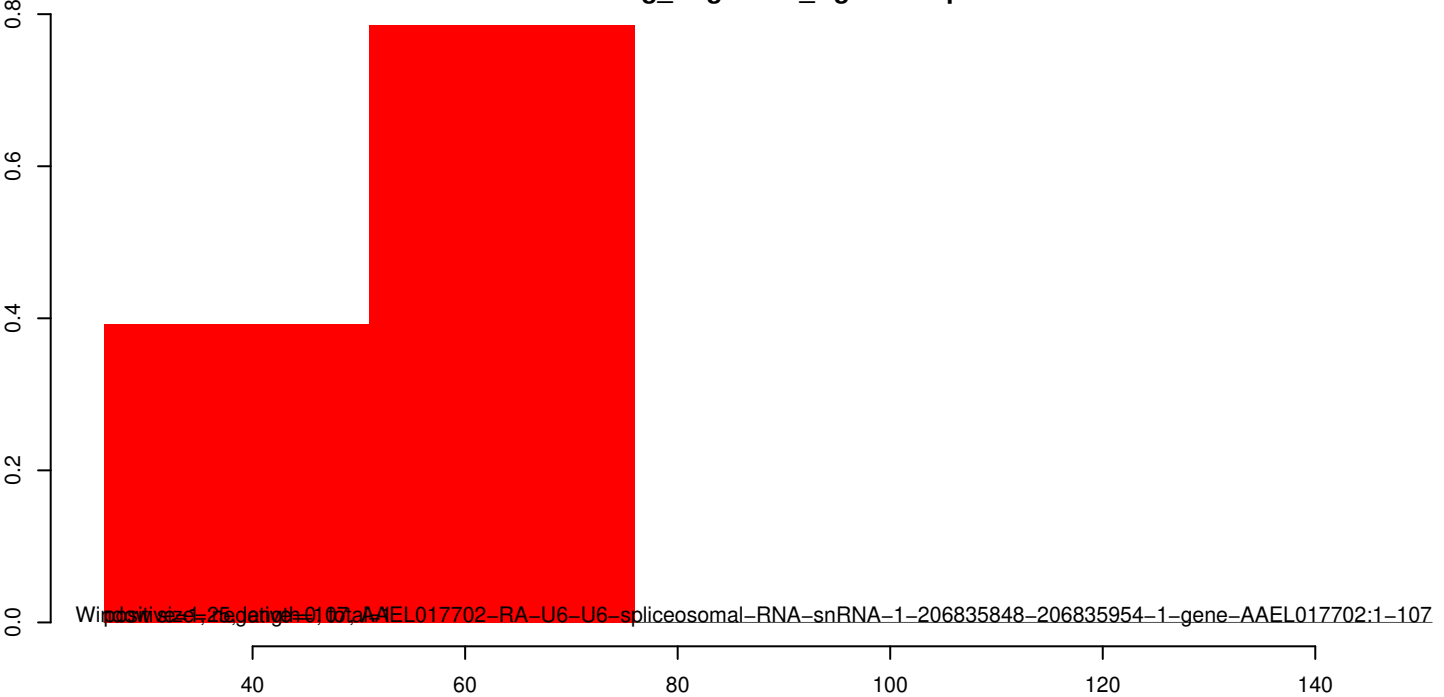
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep

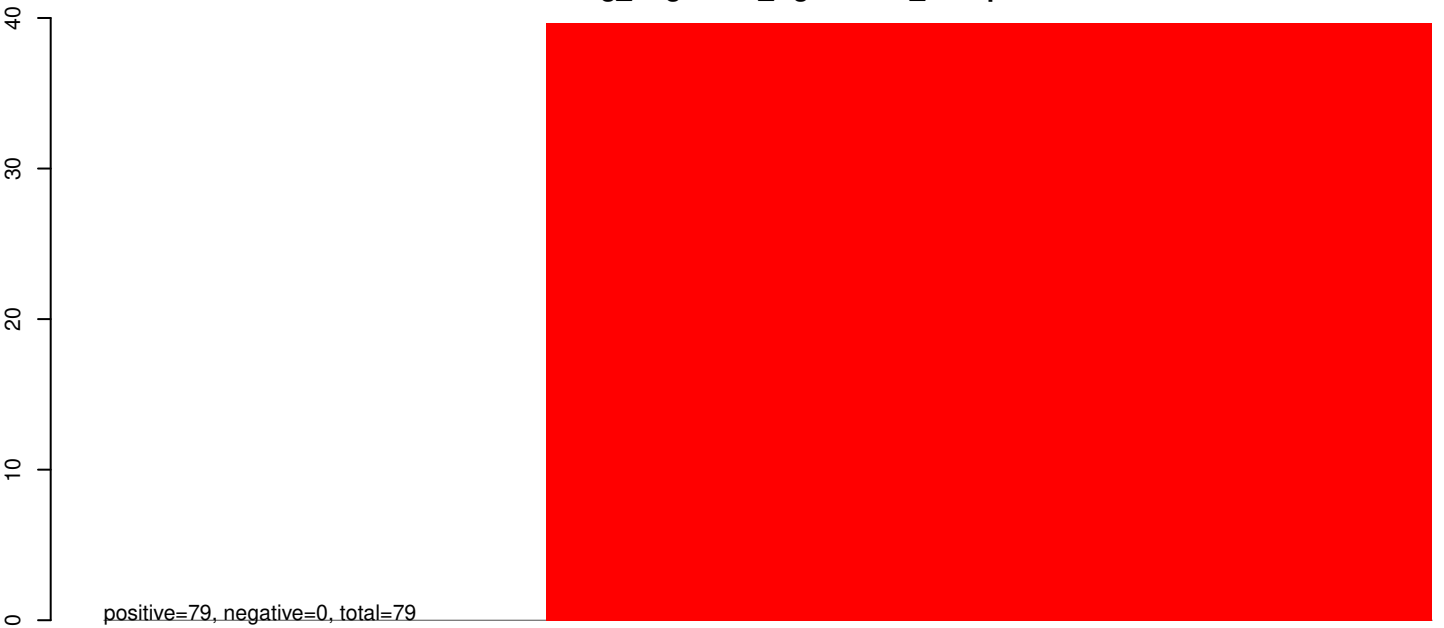


Window size = 25, gap = 0.07, AAEL017702-RA-U6-U6-spliceosomal-RNA-snRNA-1-206835848-206835954-1-gene-AAEL017702:1-107

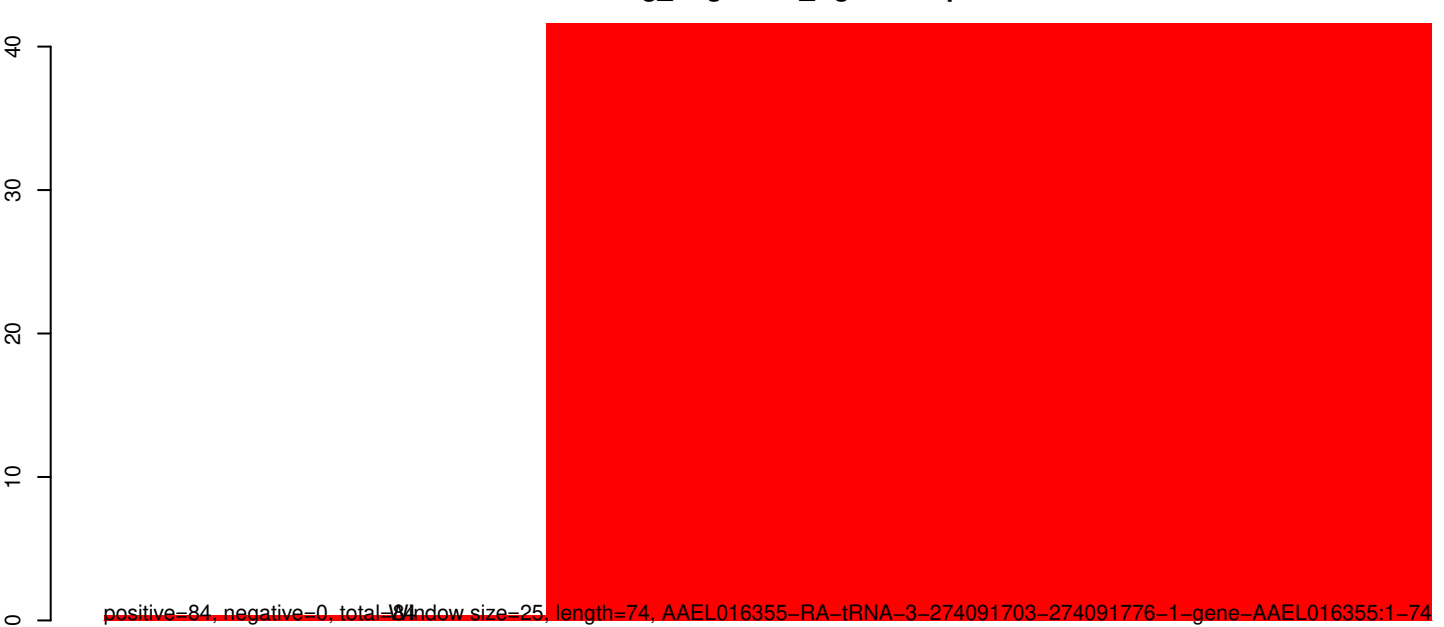
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep



Window size=25, length=74, AAEL016355-RA-tRNA-3-274091703-274091776-1-gene-AAEL016355:1-74

40 60 80 100

AeAeg_Aag2SINV_Ago3.IP.18_23.rep

positive=2, negative=0, total=2

AeAeg_Aag2SINV_Ago3.IP.24_35.rep

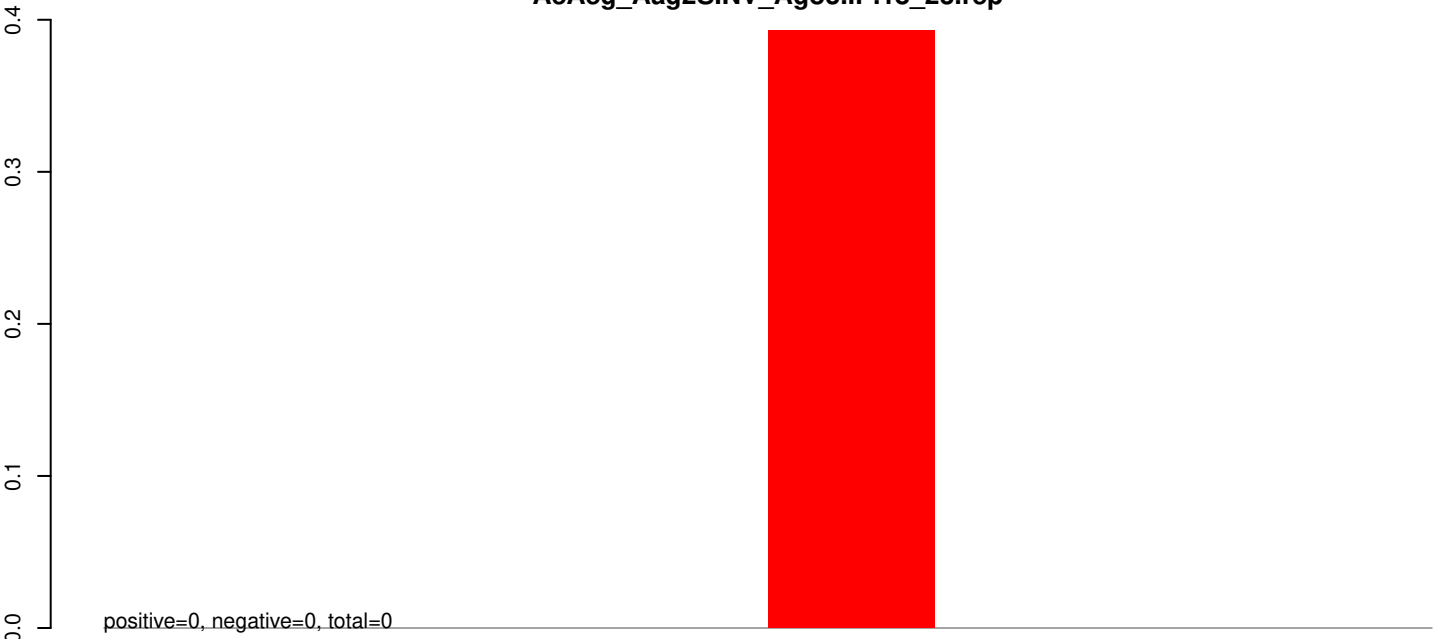
positive=2, negative=0, total=2

AeAeg_Aag2SINV_Ago3.IP.rep

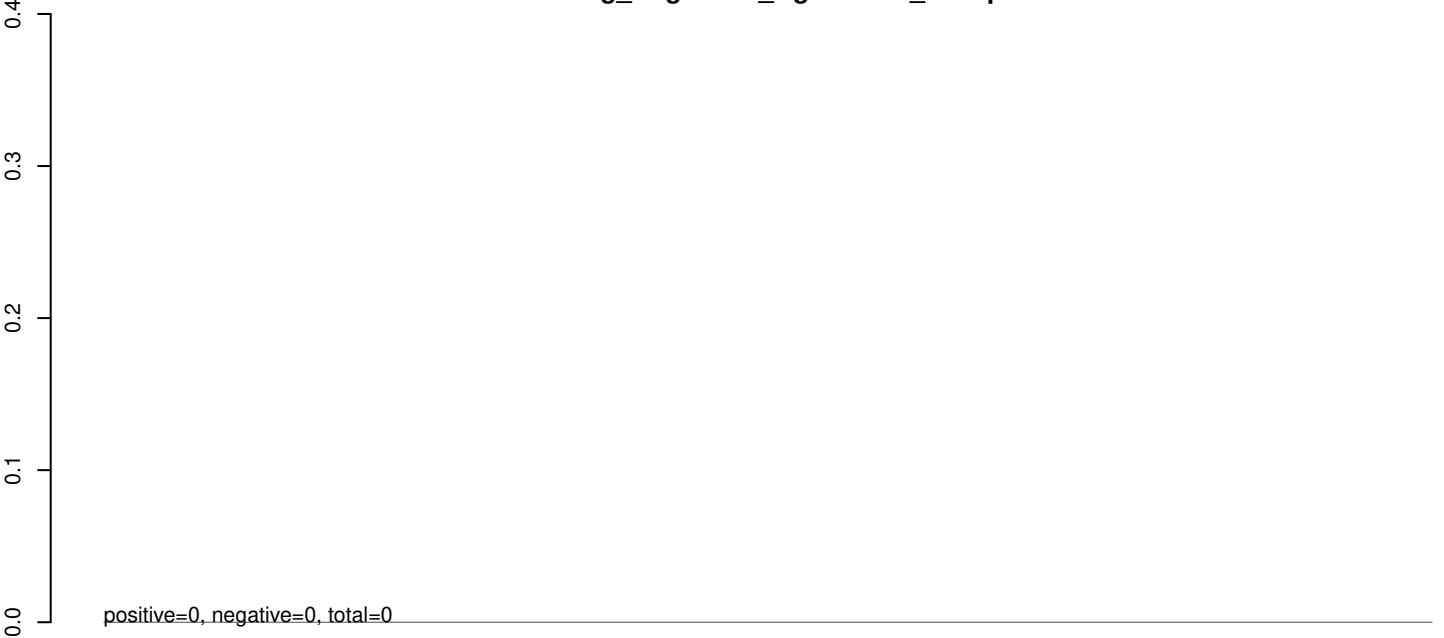
positive=3, negative=0, total=3 Window size=25, length=73, AAEL016302-RA-tRNA-3-65345468-65345540-1-gene-AAEL016302:1-73

40 60 80 100

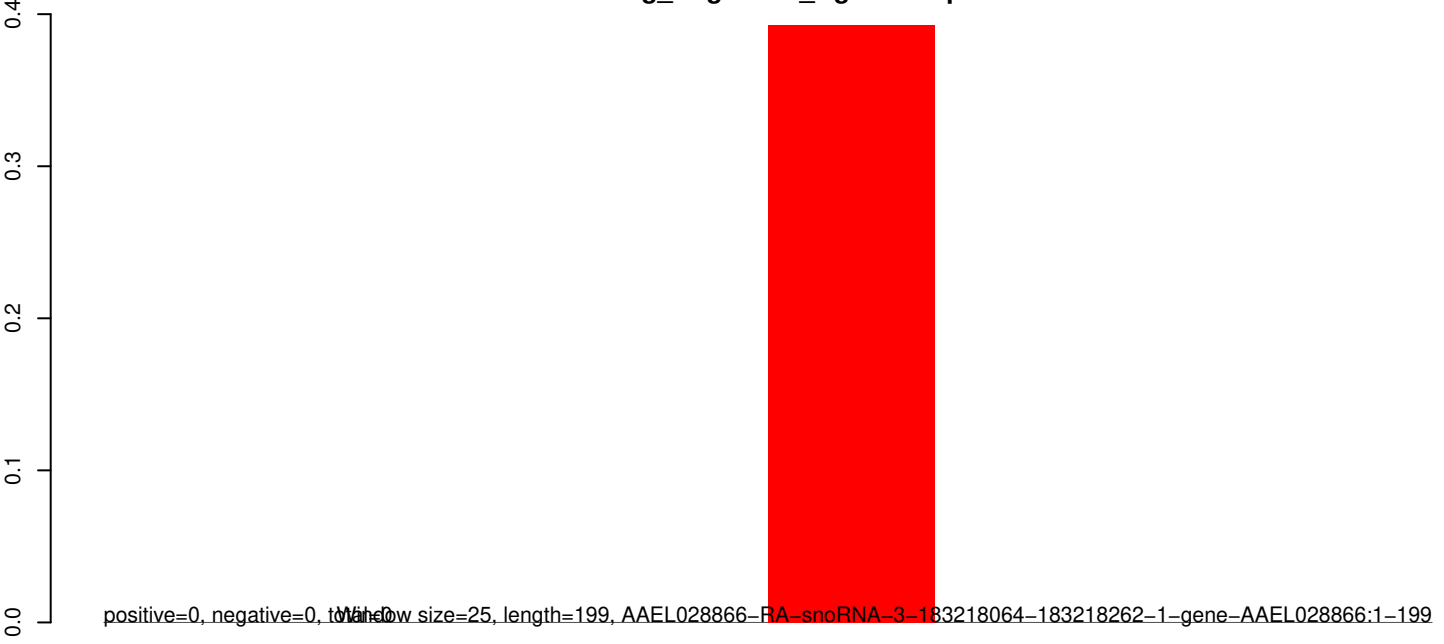
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep



Window size=25, length=199, AAEL028866-RA-snoRNA-3-183218064-183218262-1-gene-AAEL028866:1-199

AeAeg_Aag2SINV_Ago3.IP.18_23.rep

positive=1, negative=0, total=1

AeAeg_Aag2SINV_Ago3.IP.24_35.rep

positive=2, negative=0, total=2

AeAeg_Aag2SINV_Ago3.IP.rep

positive=2, negative=0, total=2 Window size=25, length=73, AAEL028693-RA-tRNA-3-100605474-100605546-1-gene-AAEL028693:1-73

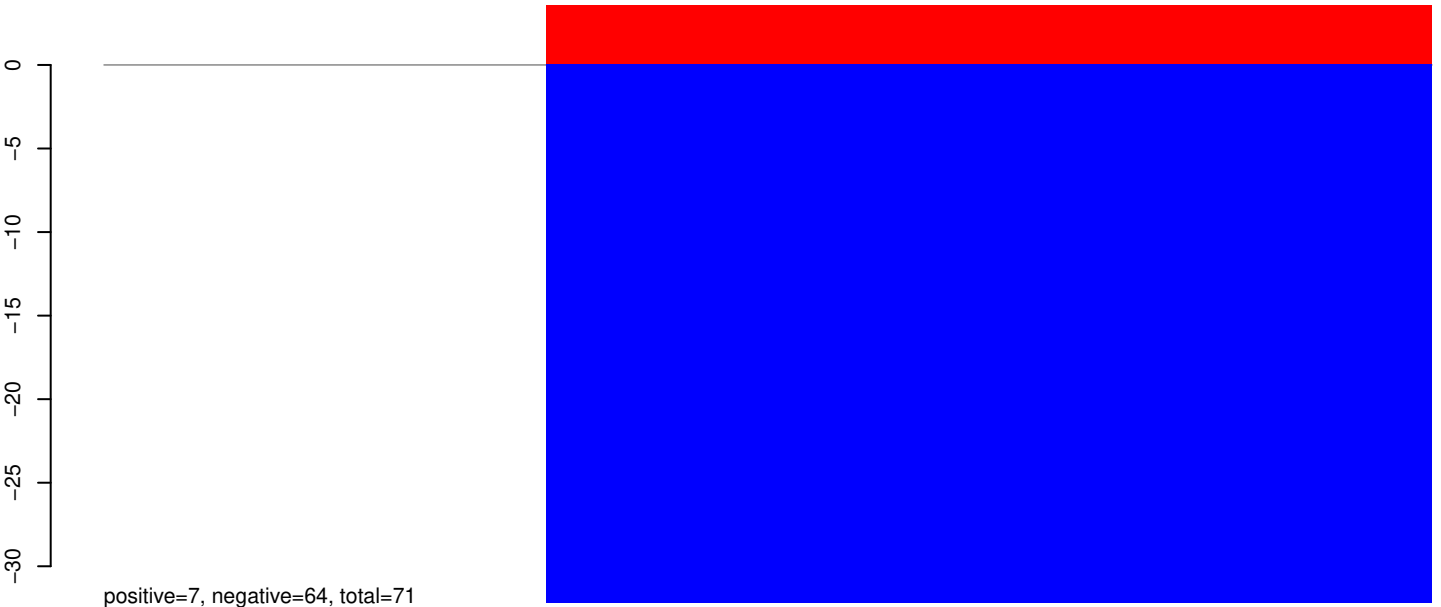
40

60

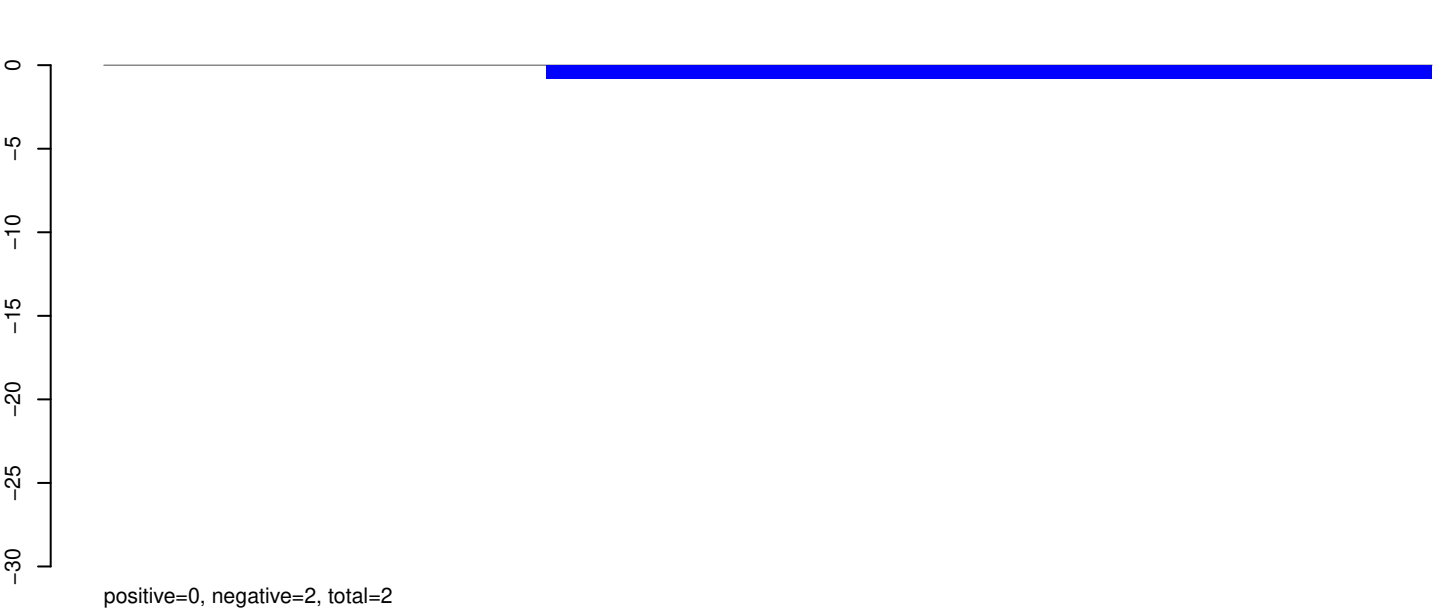
80

100

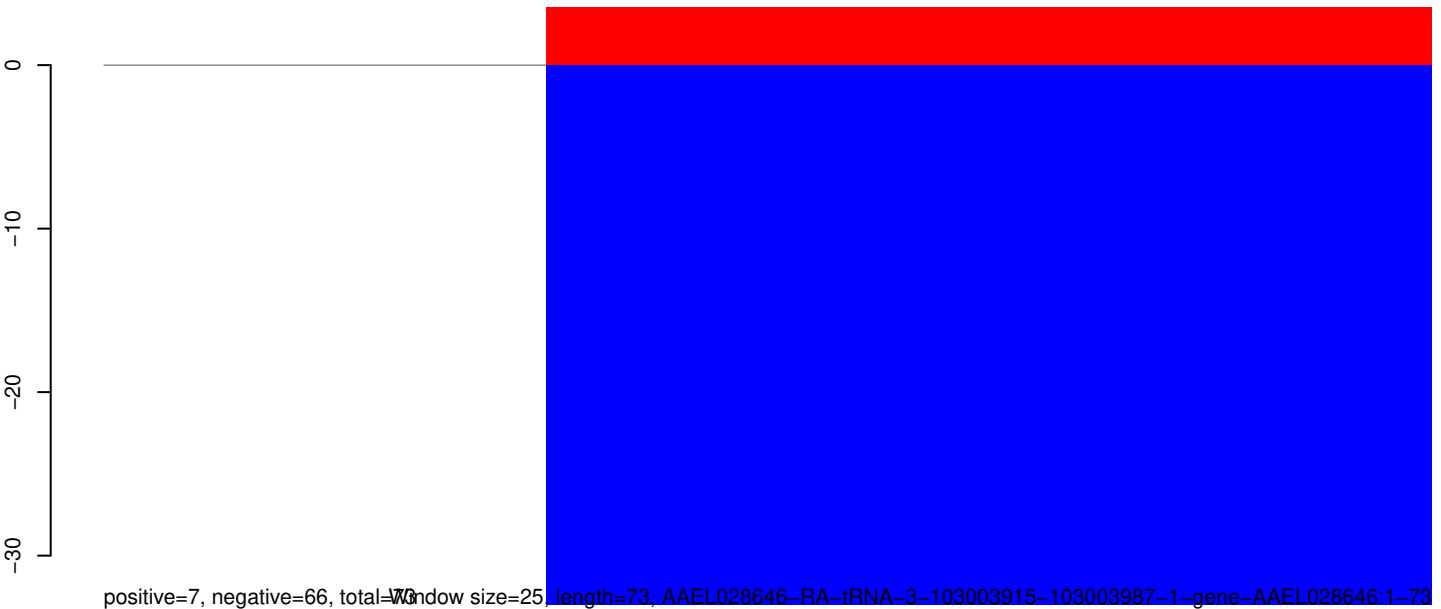
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



AeAeg_Aag2SINV_Ago3.IP.24_35.rep



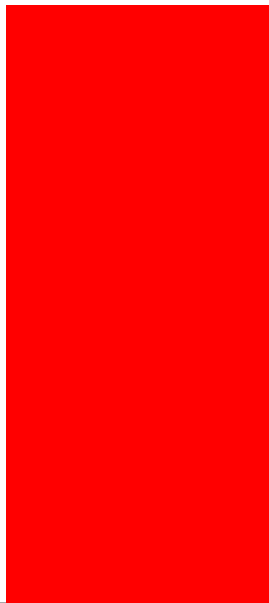
AeAeg_Aag2SINV_Ago3.IP.rep



40 60 80 100

AeAeg_Aag2SINV_Ago3.IP.18_23.rep

positive=1, negative=0, total=1



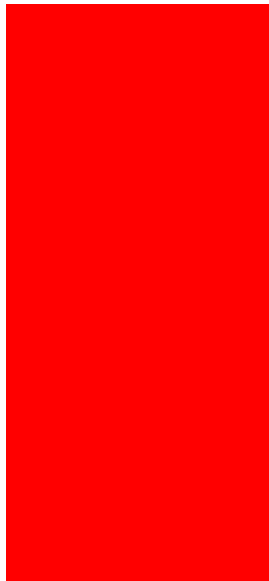
AeAeg_Aag2SINV_Ago3.IP.24_35.rep

positive=0, negative=0, total=0



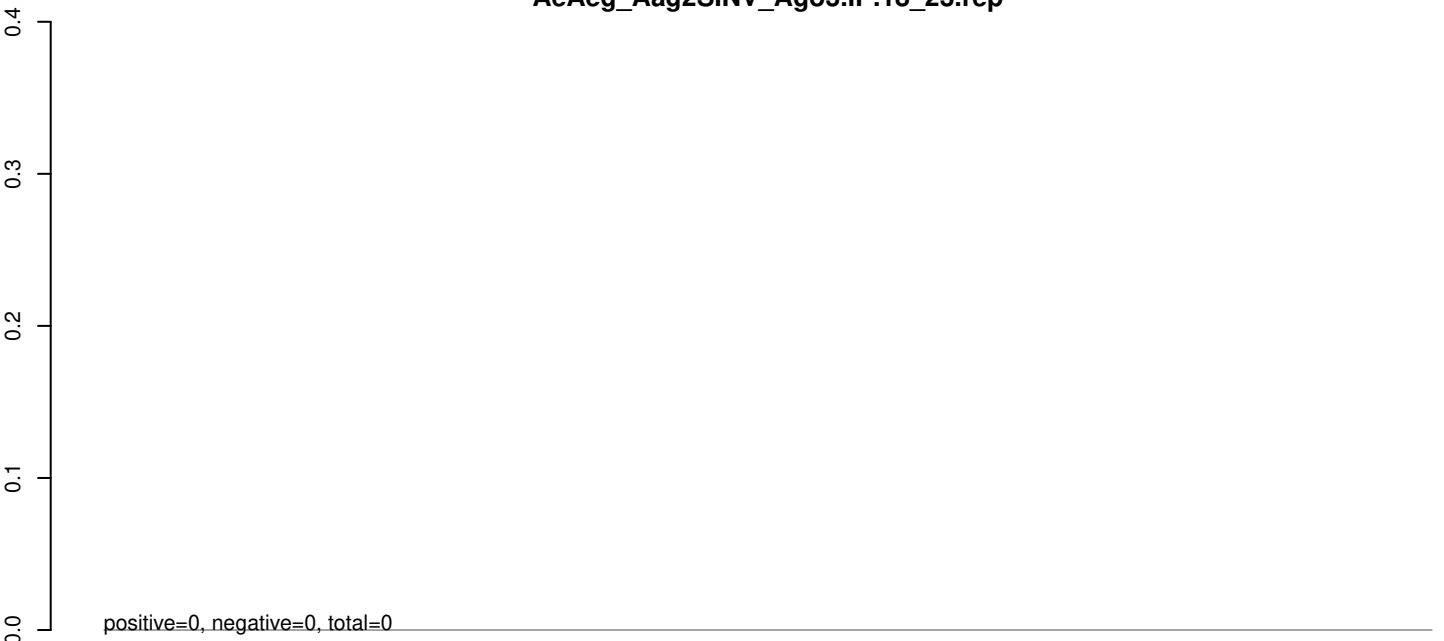
AeAeg_Aag2SINV_Ago3.IP.rep

positive=2, negative=0, total=2
Window size=25, length=122, AAEL019314-RA-siRNA-2-310045432-310045553-1-gene-AAEL019314:1-122

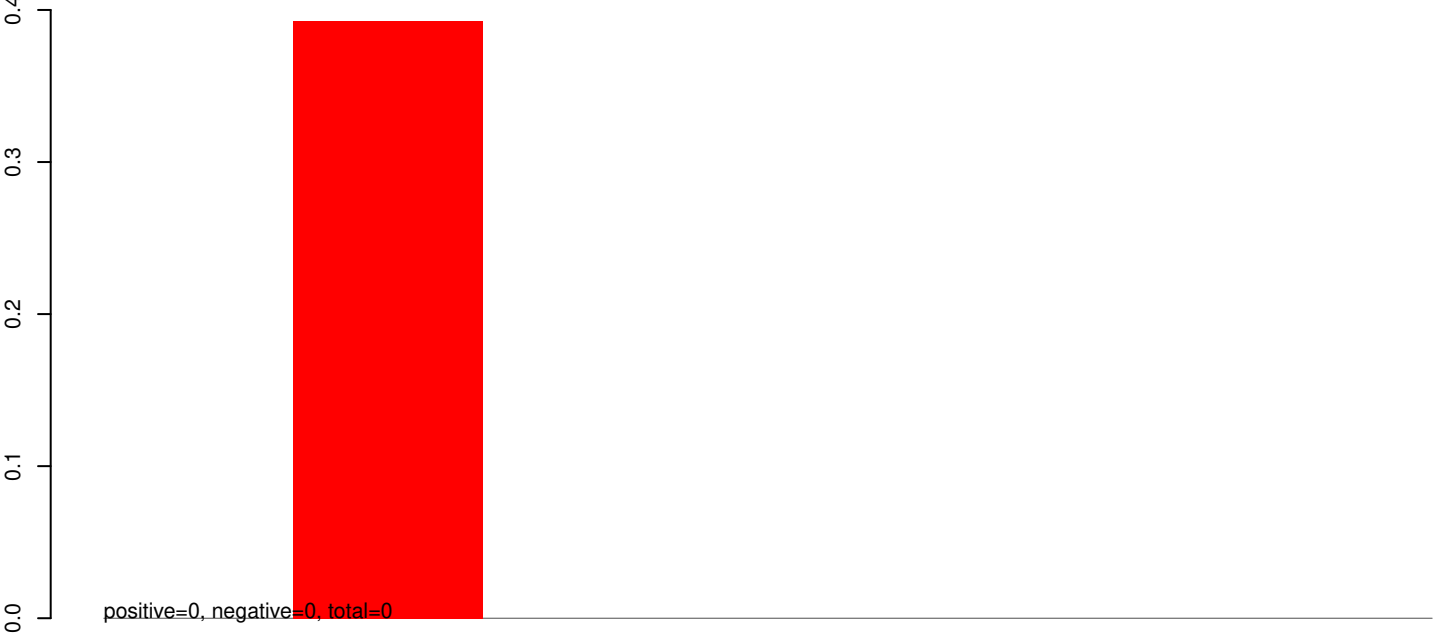


40 60 80 100 120 140

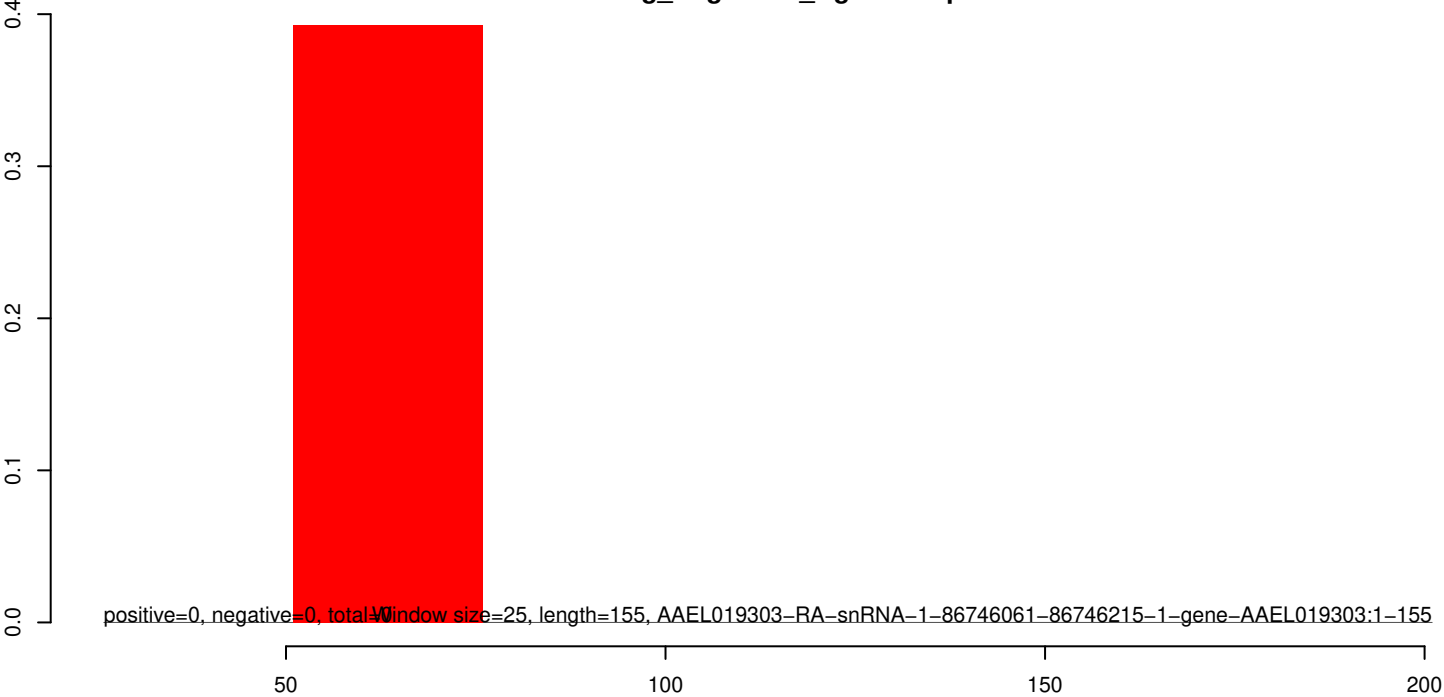
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



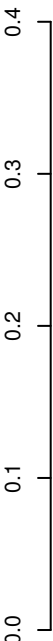
AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep

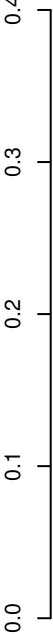


AeAeg_Aag2SINV_Ago3.IP.18_23.rep



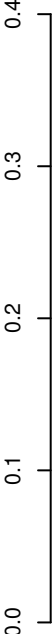
positive=0, negative=0, total=0

AeAeg_Aag2SINV_Ago3.IP.24_35.rep

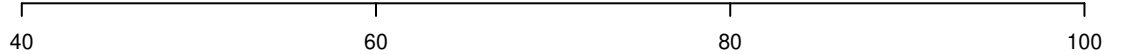


positive=1, negative=0, total=1

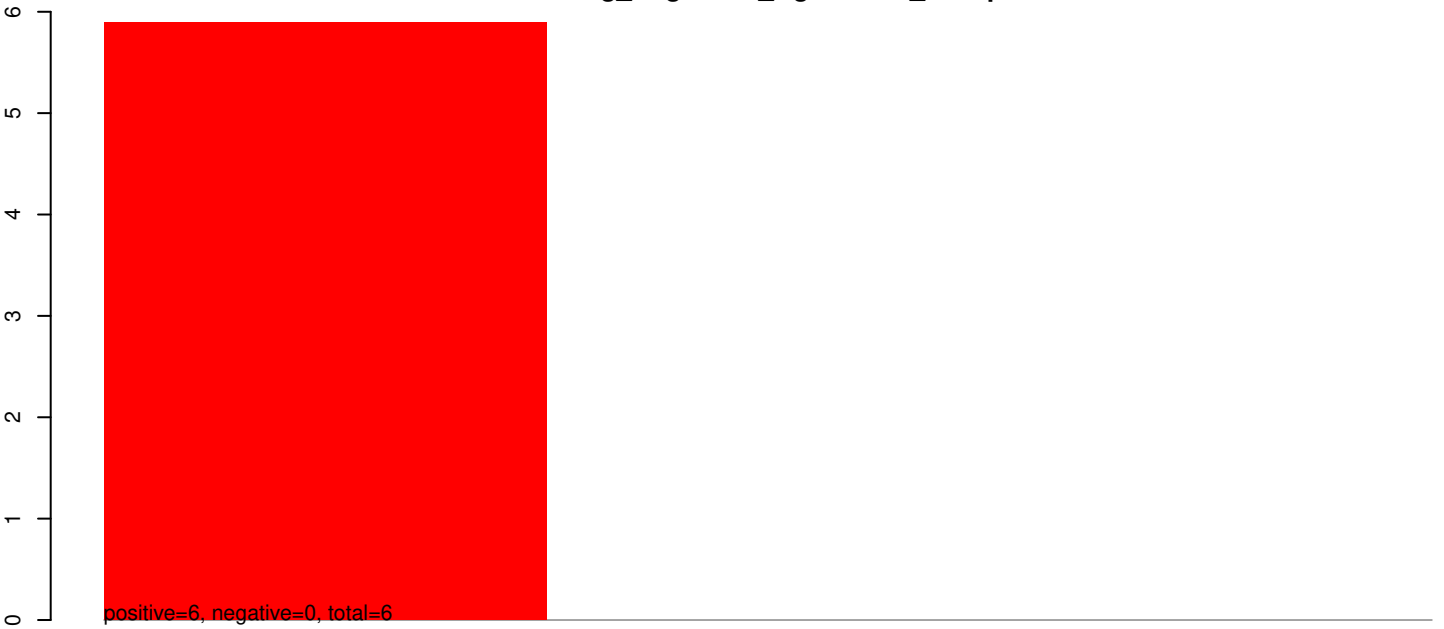
AeAeg_Aag2SINV_Ago3.IP.rep



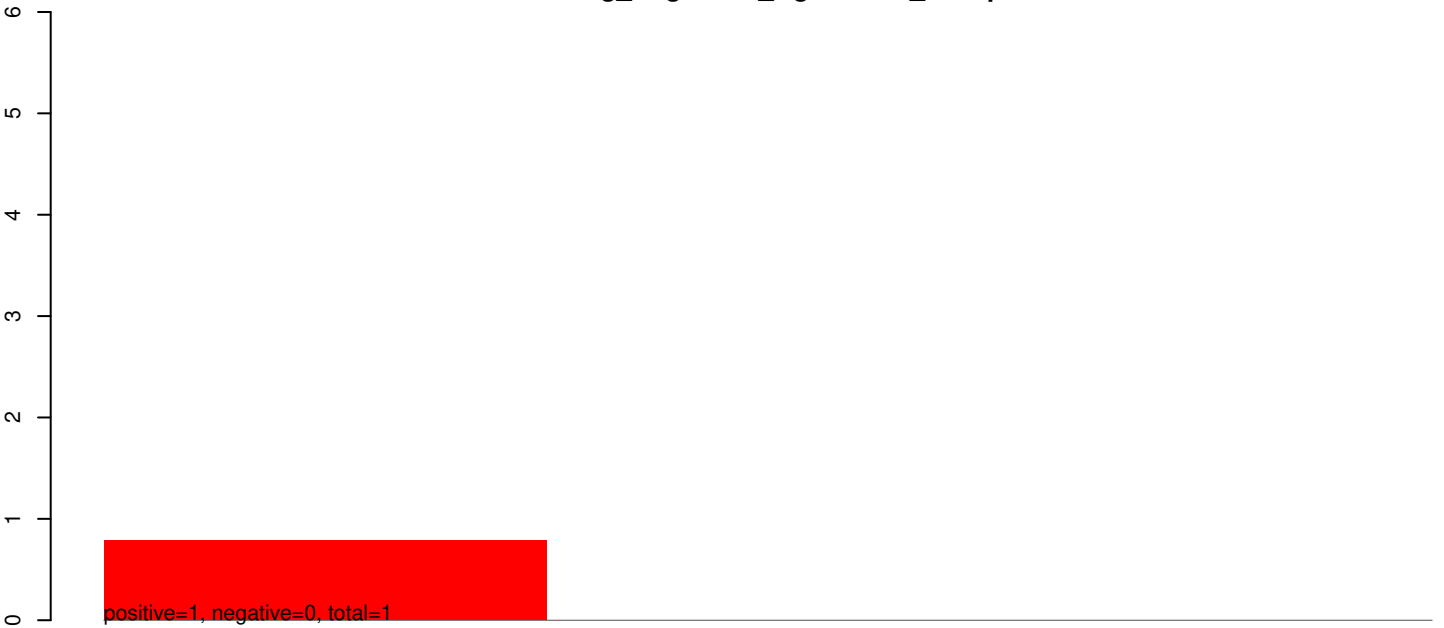
positive=1, negative=0, total=1Window size=25, length=73, AAEL018936-RA-tRNA-2-191424418-191424490-1-gene-AAEL018936:1-73



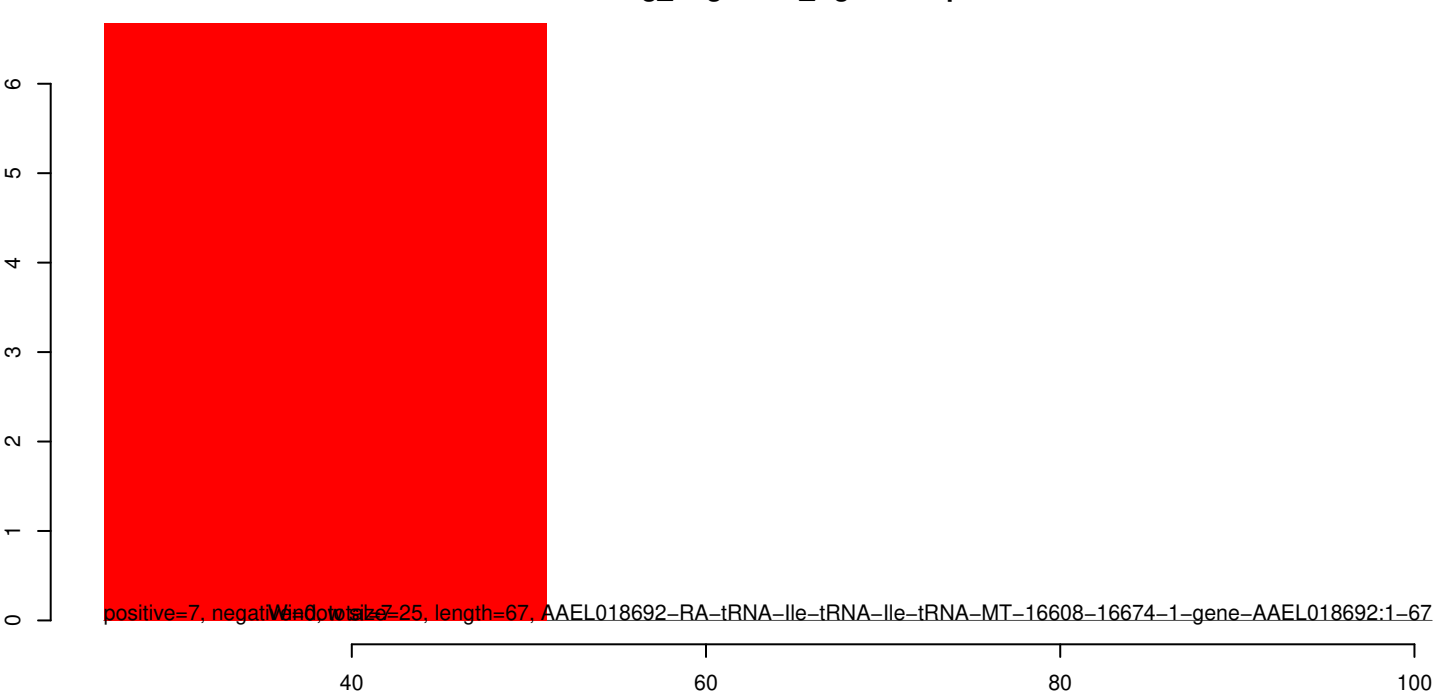
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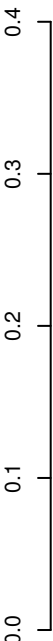
AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep

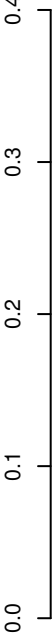


AeAeg_Aag2SINV_Ago3.IP.18_23.rep



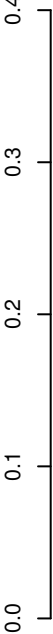
positive=0, negative=0, total=0

AeAeg_Aag2SINV_Ago3.IP.24_35.rep



positive=1, negative=0, total=1

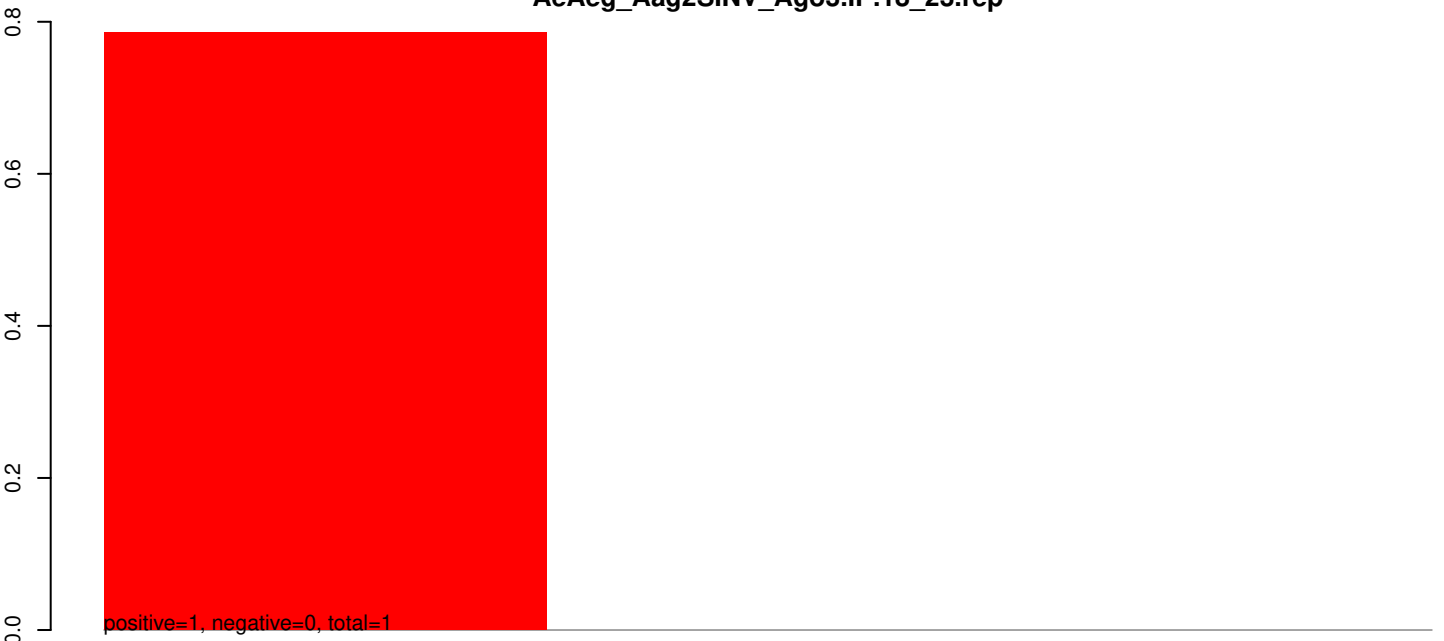
AeAeg_Aag2SINV_Ago3.IP.rep



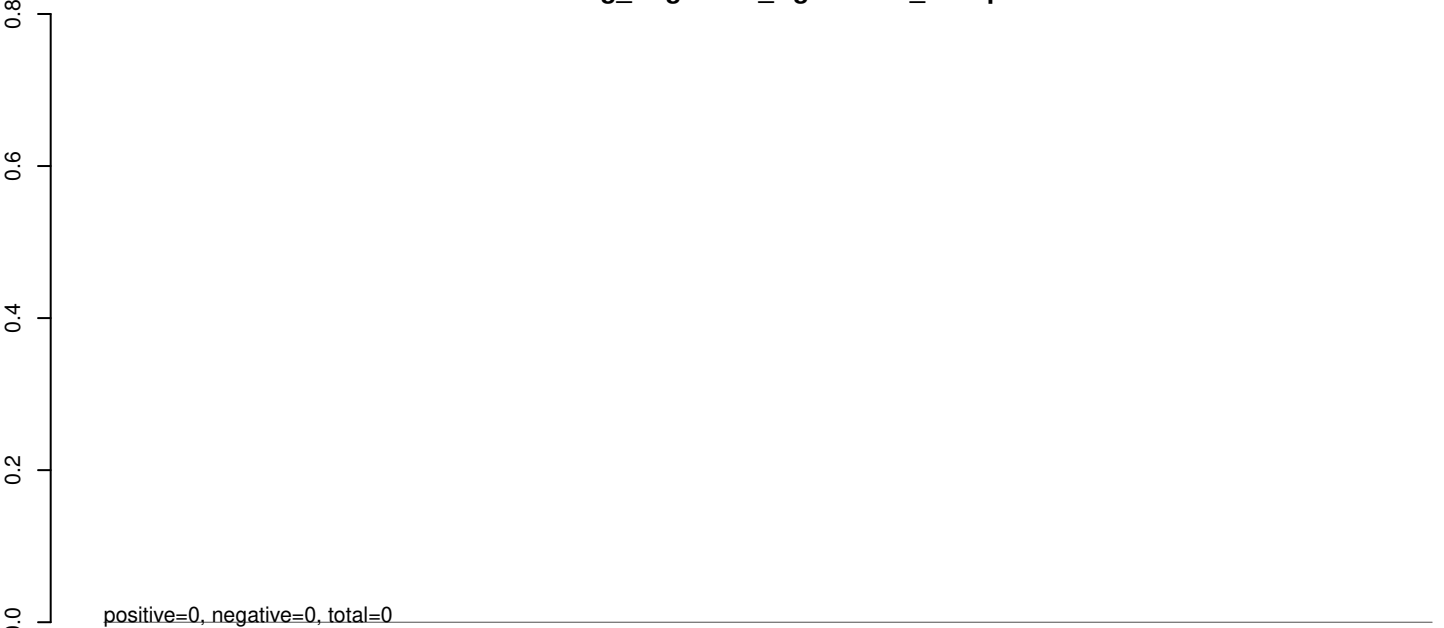
positive=1, negative=0, total=25, length=72, AAEL018690-RA-tRNA-Val-tRNA-Val-tRNA-MT-13901-13972-1-gene-AAEL018690:1-72



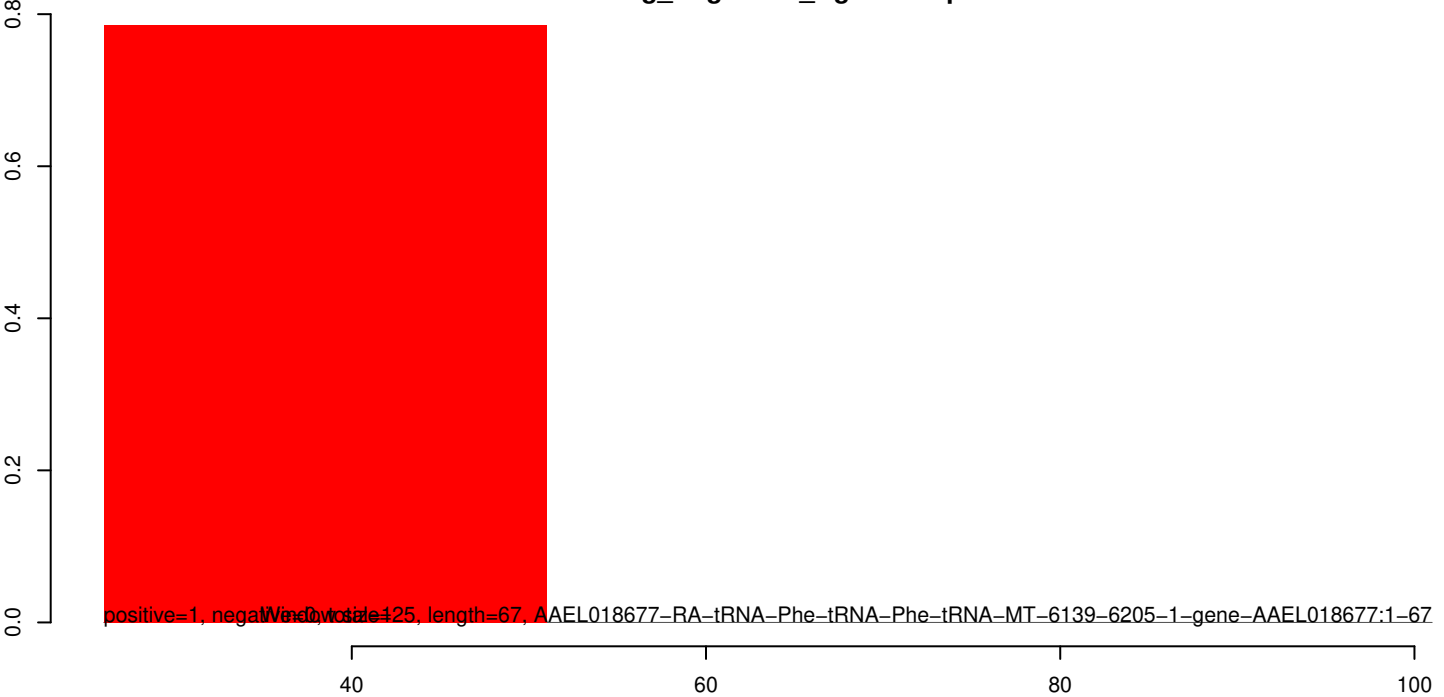
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



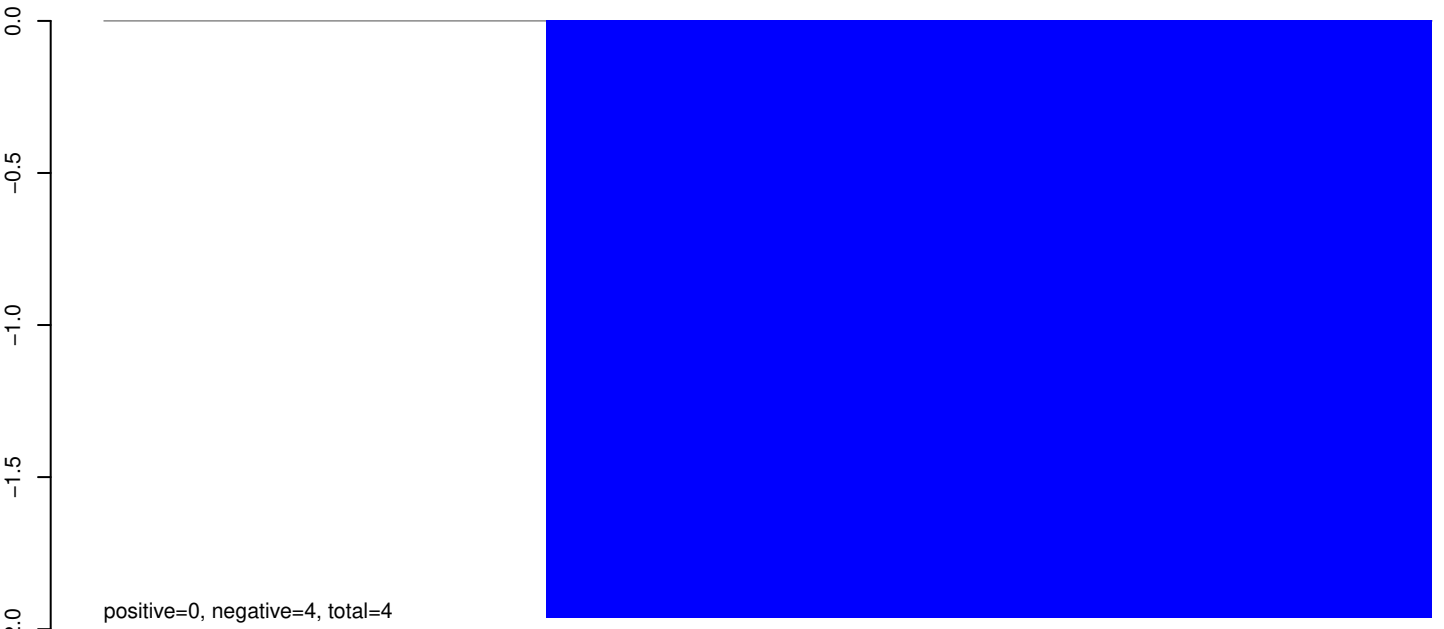
AeAeg_Aag2SINV_Ago3.IP.24_35.rep



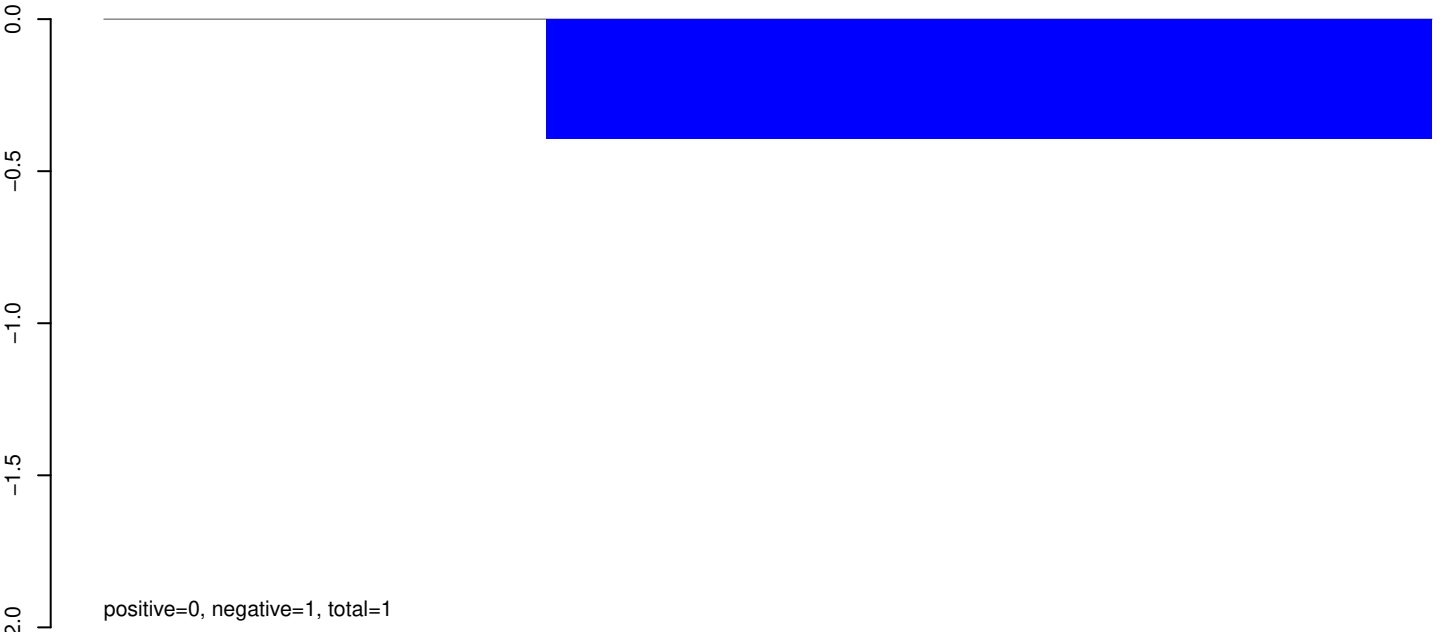
AeAeg_Aag2SINV_Ago3.IP.rep



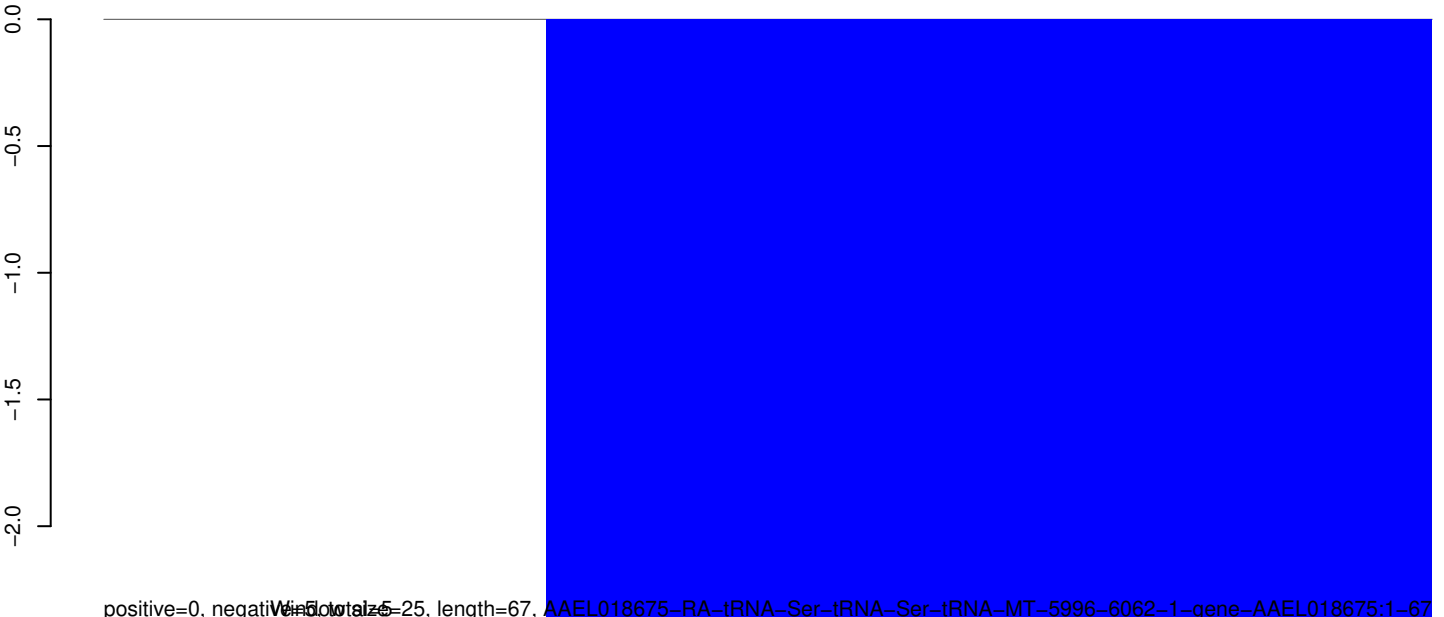
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep



Window size=25, length=67, AAEL018675-RA-tRNA-Ser-tRNA-Ser-tRNA-MT-5996-6062-1-gene-AAEL018675:1-67

40 60 80 100

AeAeg_Aag2SINV_Ago3.IP.18_23.rep

positive=2, negative=0, total=2

AeAeg_Aag2SINV_Ago3.IP.24_35.rep

positive=2, negative=0, total=2

AeAeg_Aag2SINV_Ago3.IP.rep

positive=4, negative=0, window size=25, length=67, AAEL018670-RA-tRNA-Gly-tRNA-Gly-tRNA-MT-5371-5437-1-gene-AAEL018670:1-67

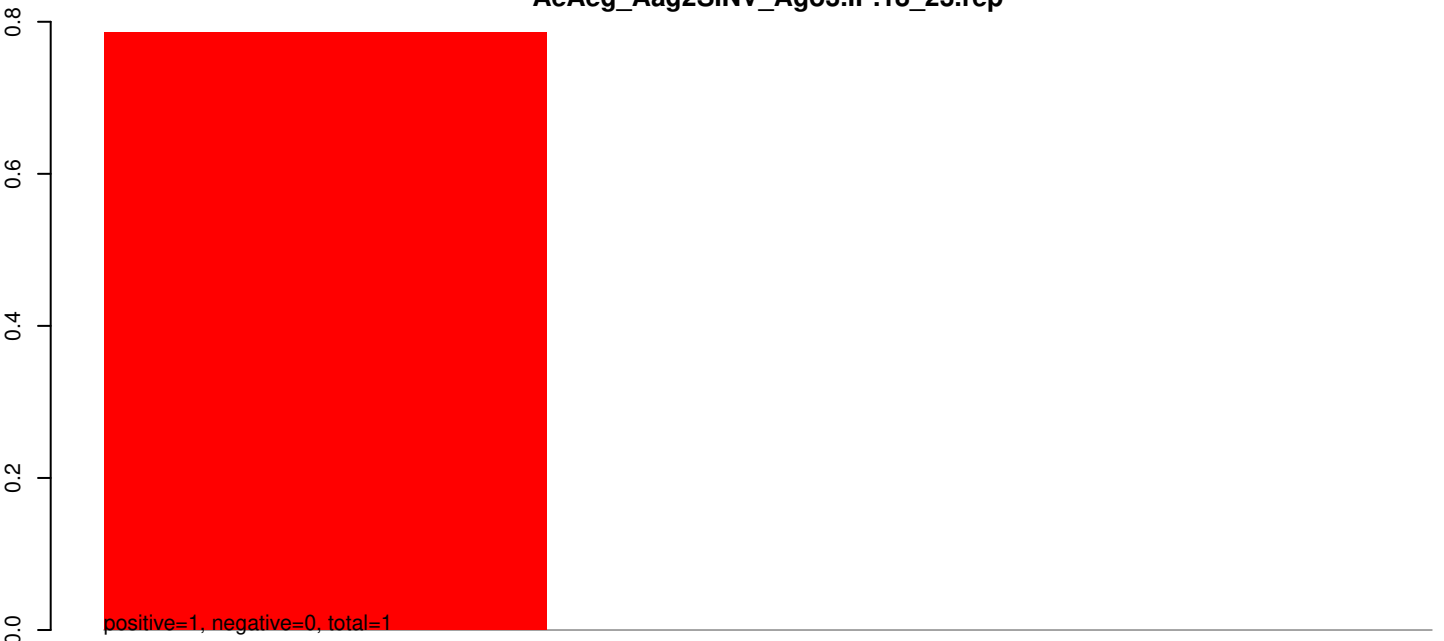
40

60

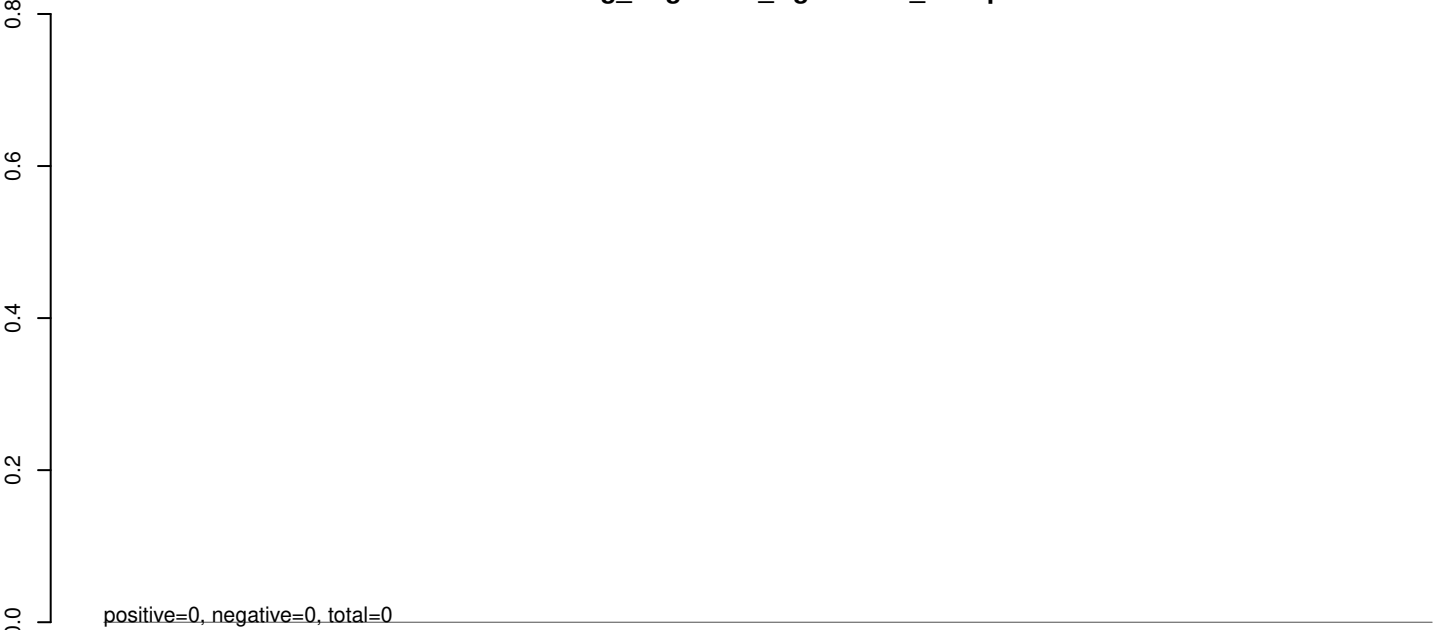
80

100

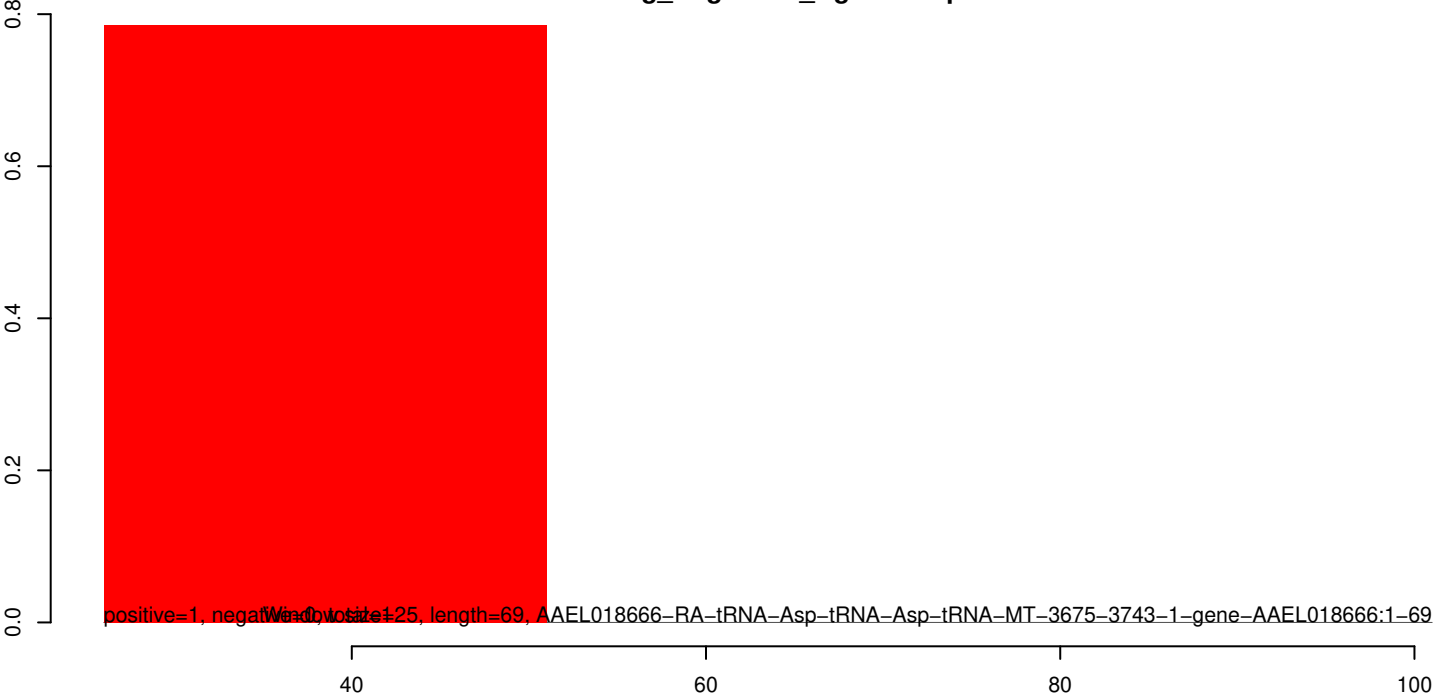
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep



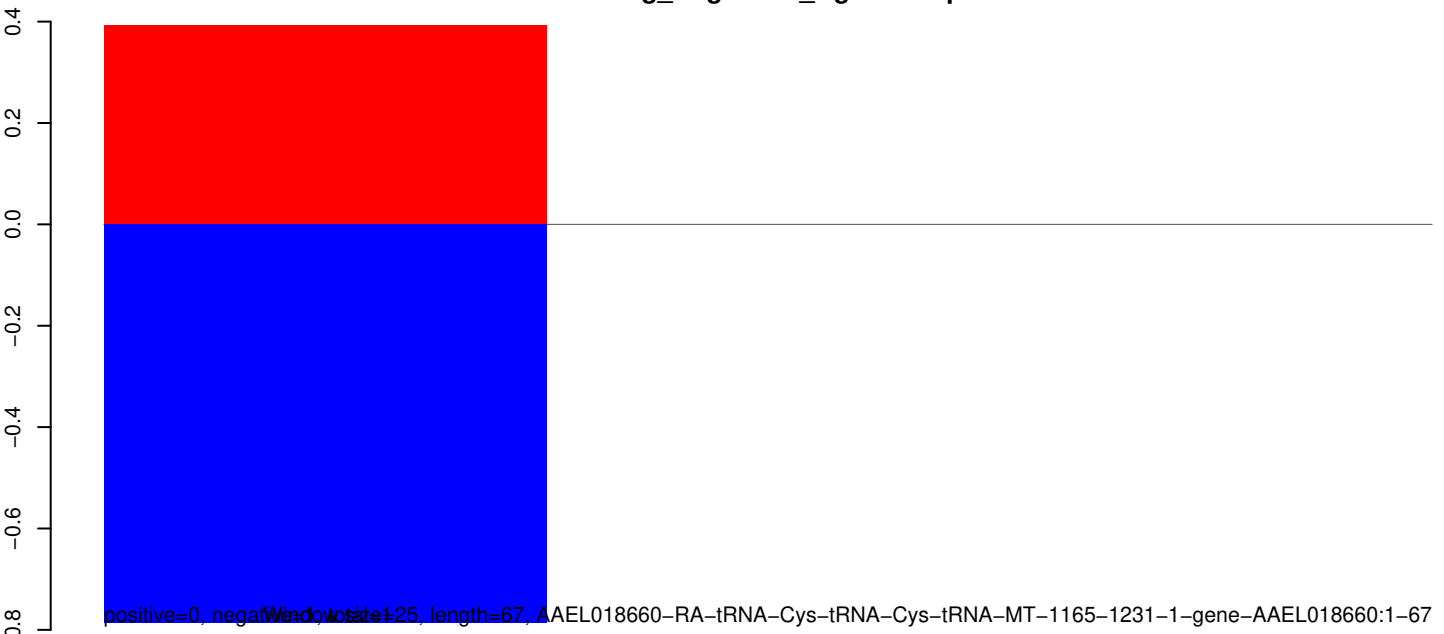
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep



Window size=25, length=67, AAEL018660-RA-tRNA-Cys-tRNA-Cys-tRNA-MT-1165-1231-1-gene-AAEL018660:1-67

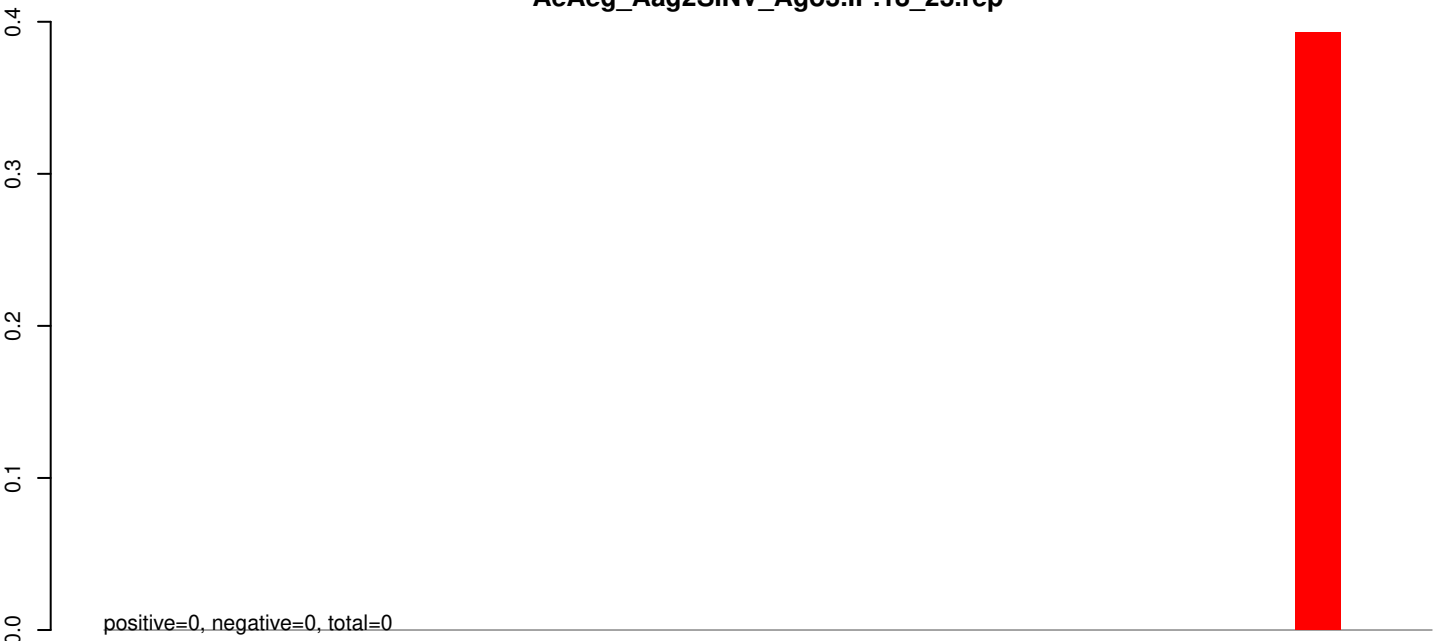
40

60

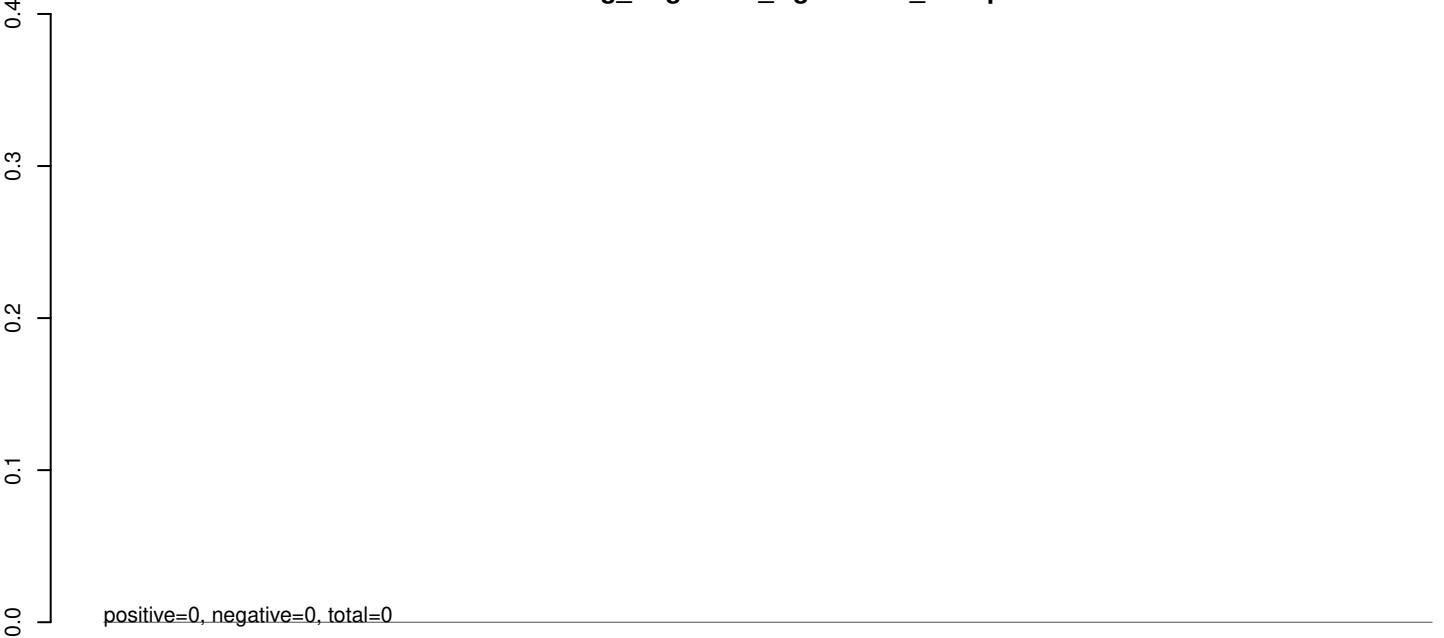
80

100

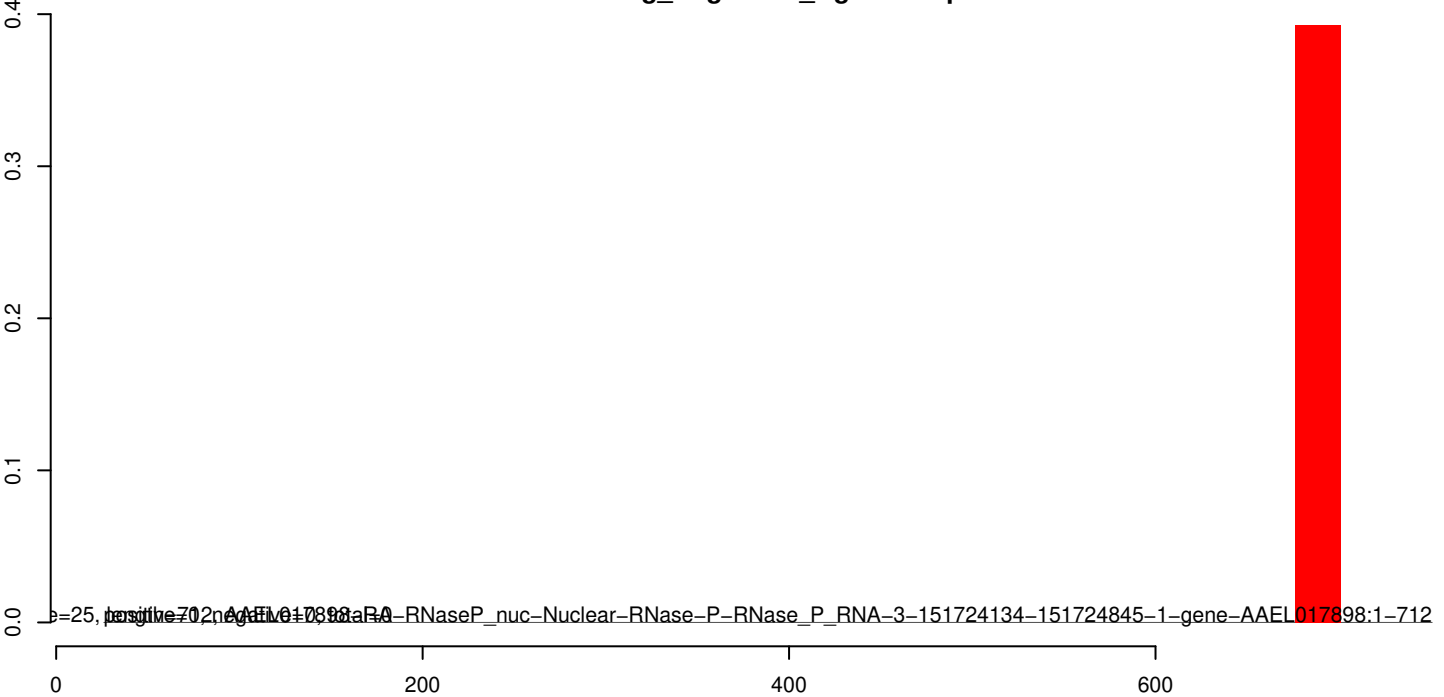
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



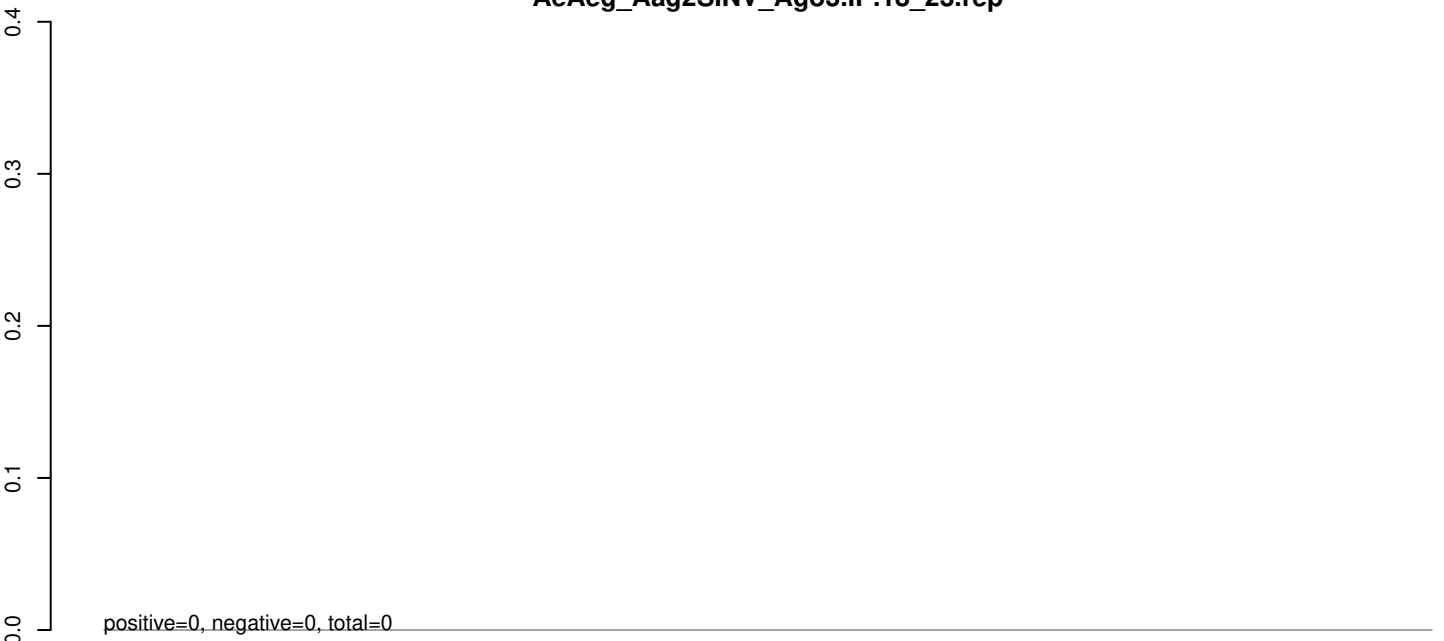
AeAeg_Aag2SINV_Ago3.IP.24_35.rep



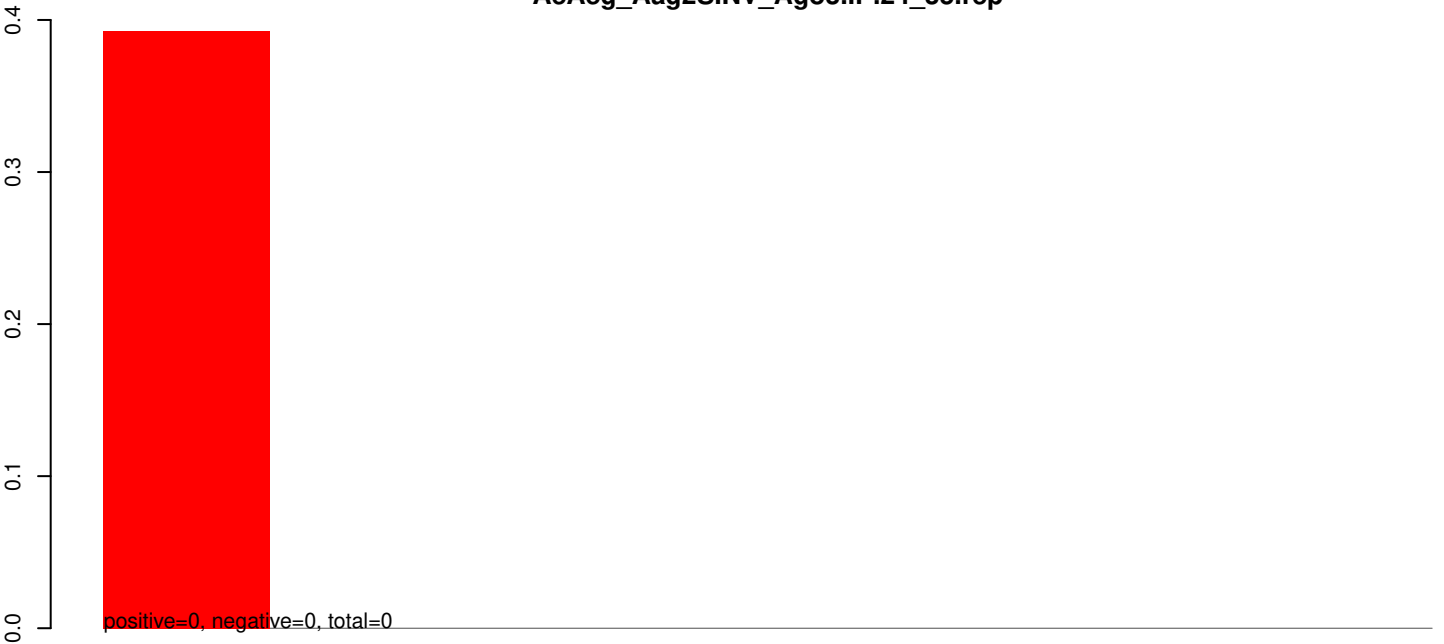
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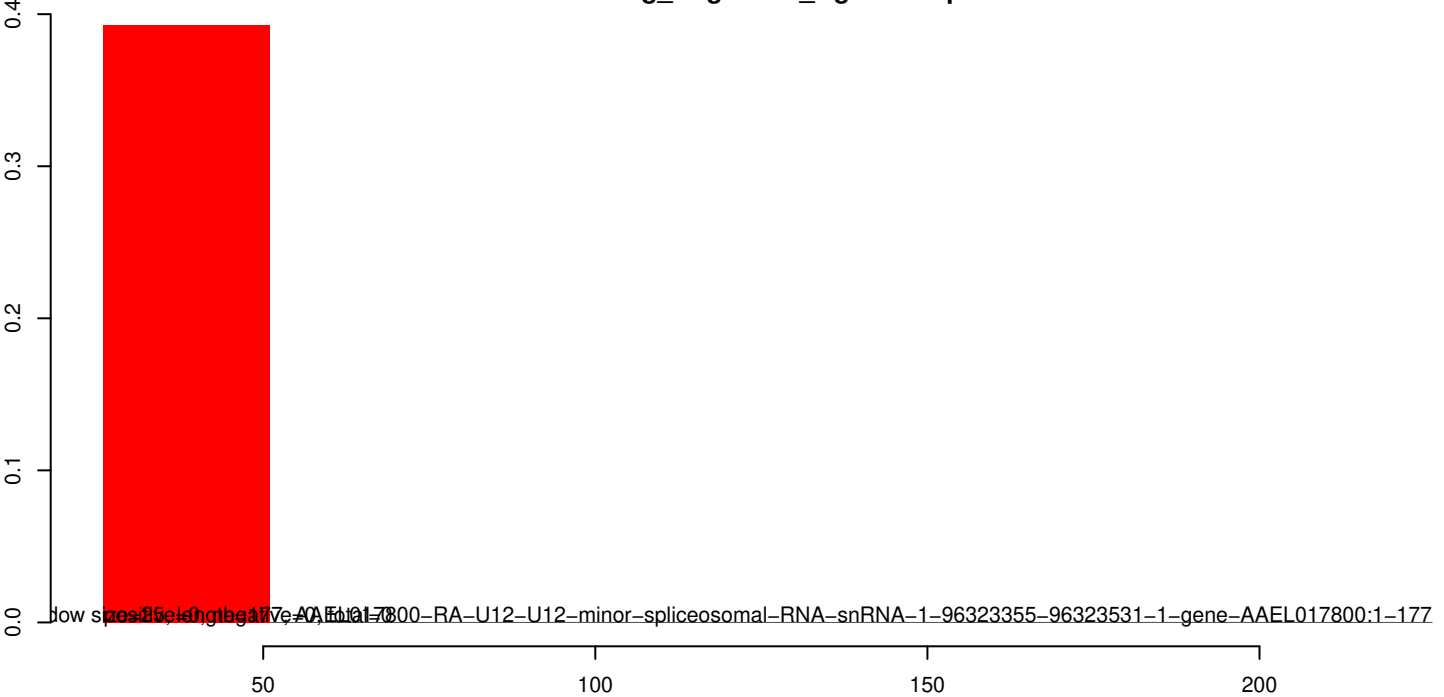
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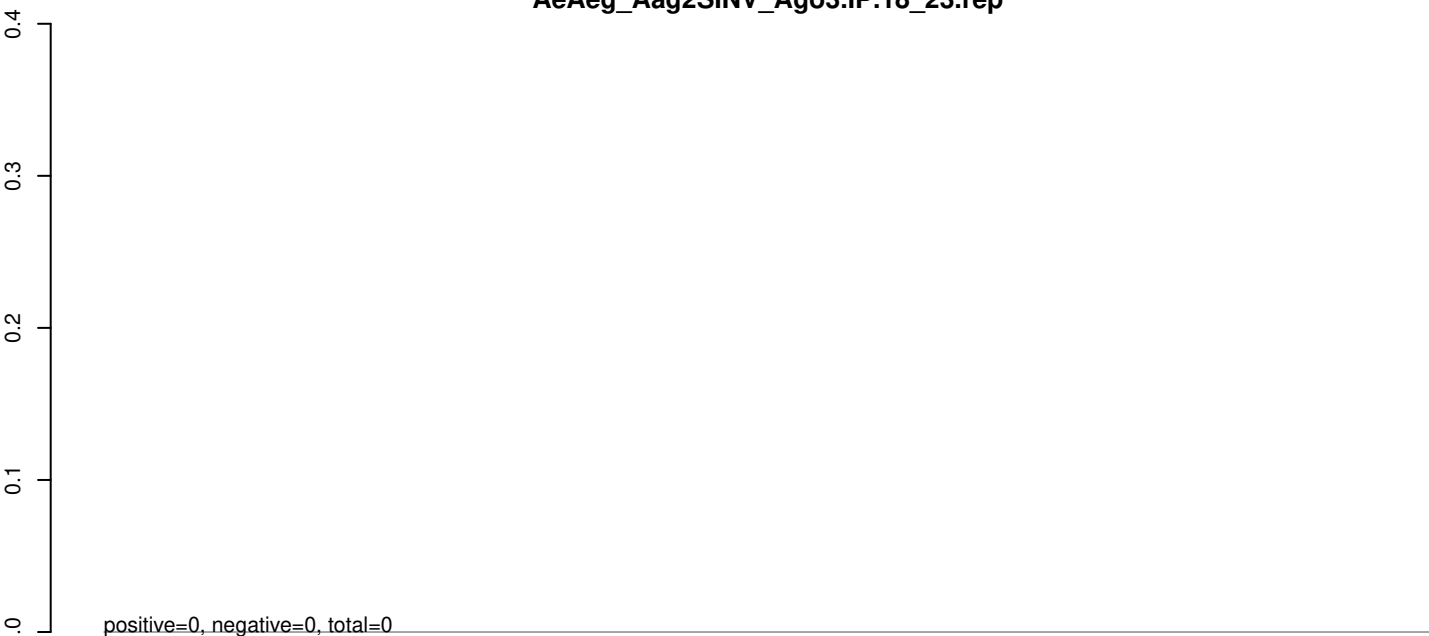
AeAeg_Aag2SINV_Ago3.IP.24_35.rep



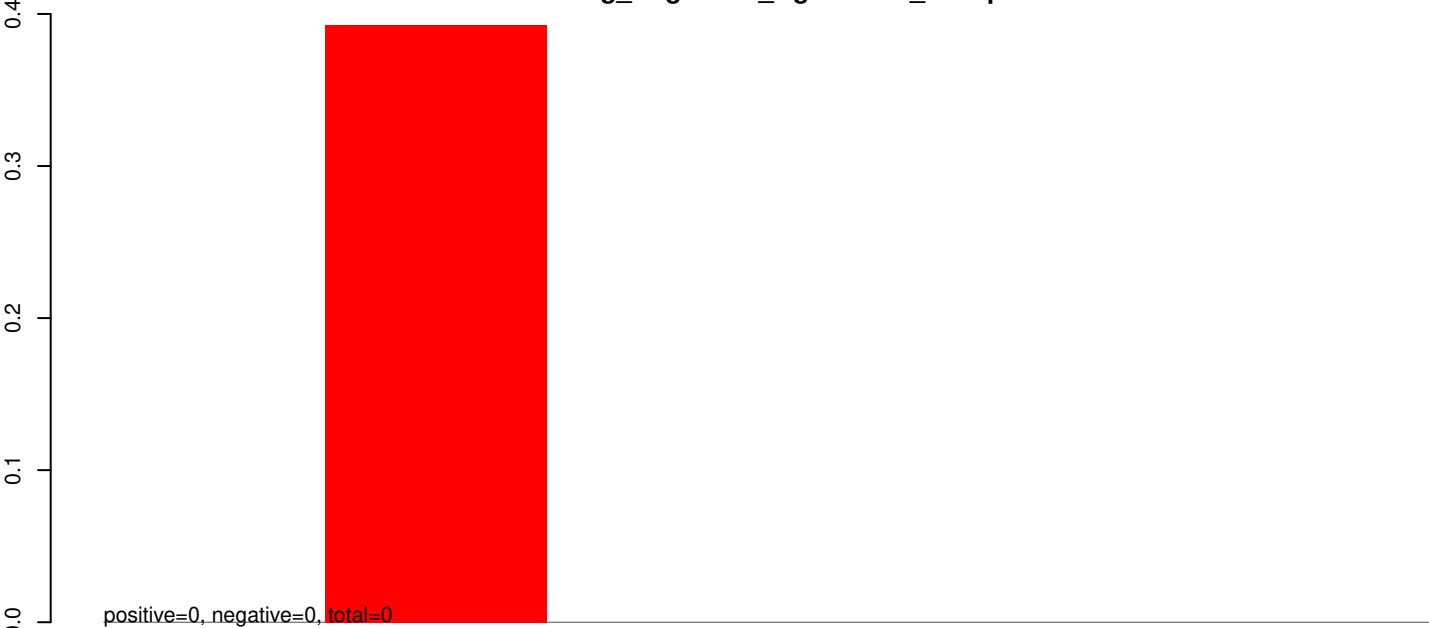
AeAeg_Aag2SINV_Ago3.IP.rep



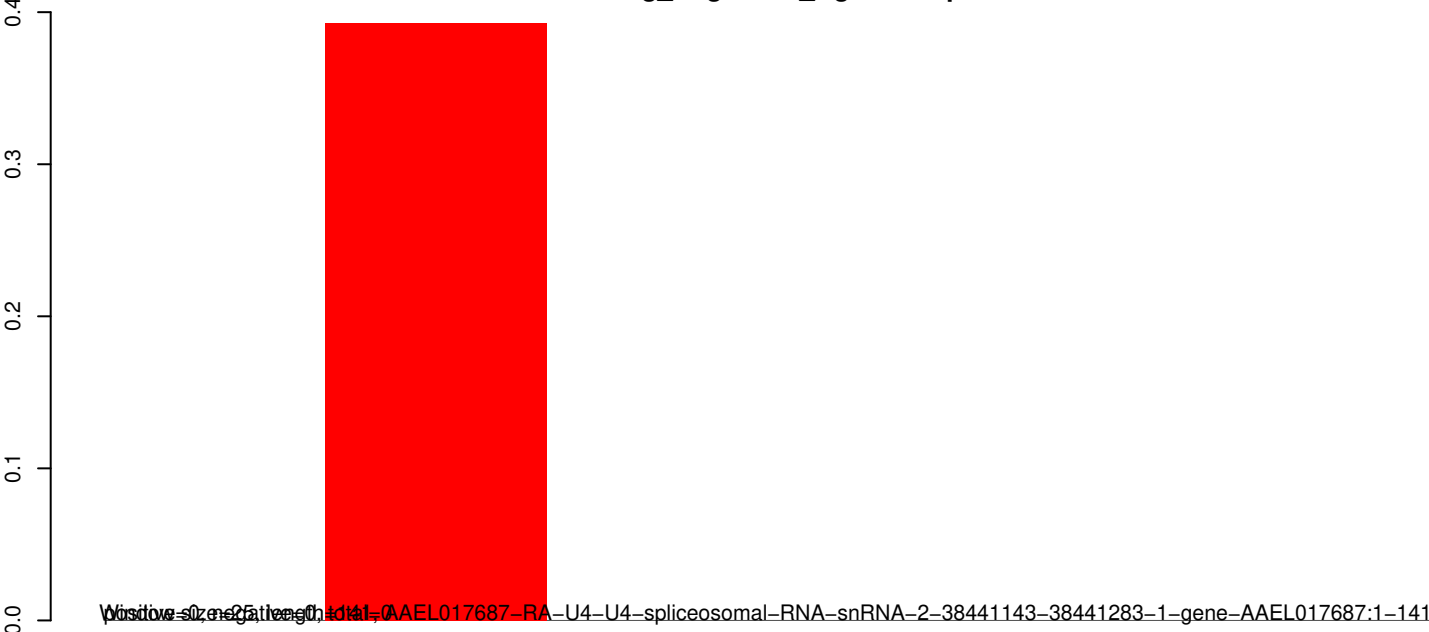
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep



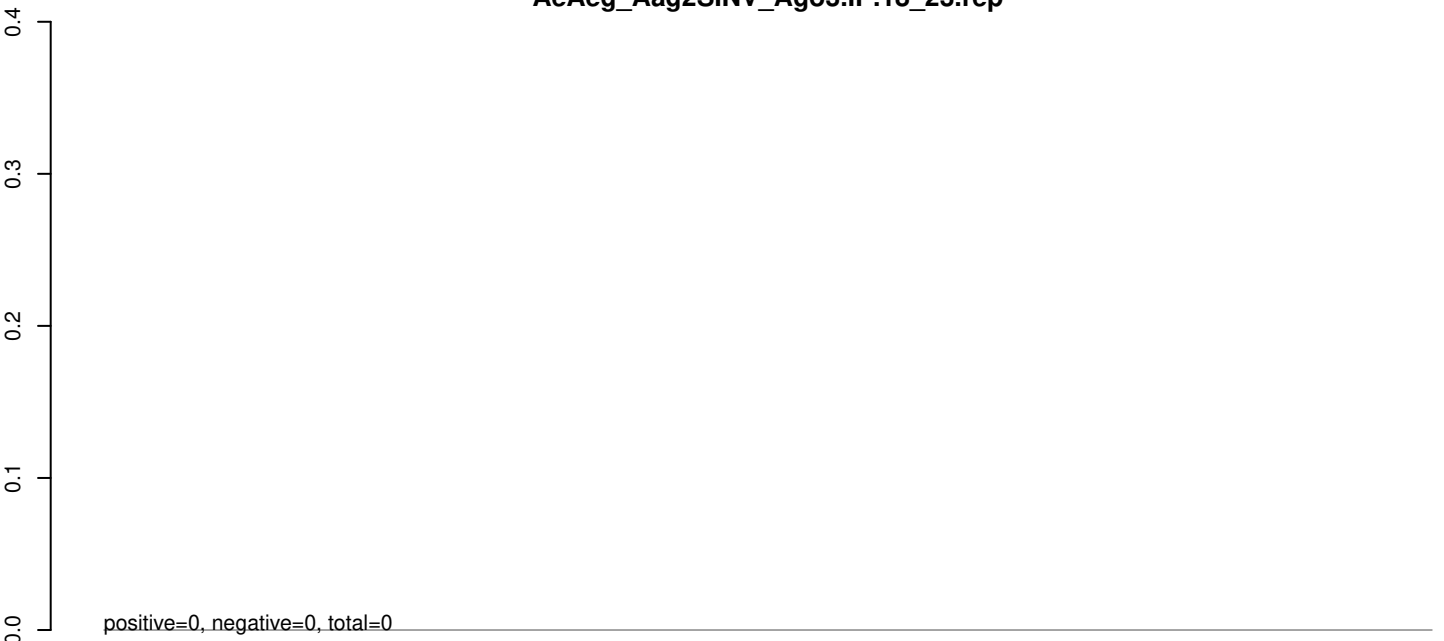
Window size: 25, step: 1, AAEL017687-RA-U4-U4-spliceosomal-RNA-snrRNA-2-38441143-38441283-1-gene-AAEL017687:1-141

50

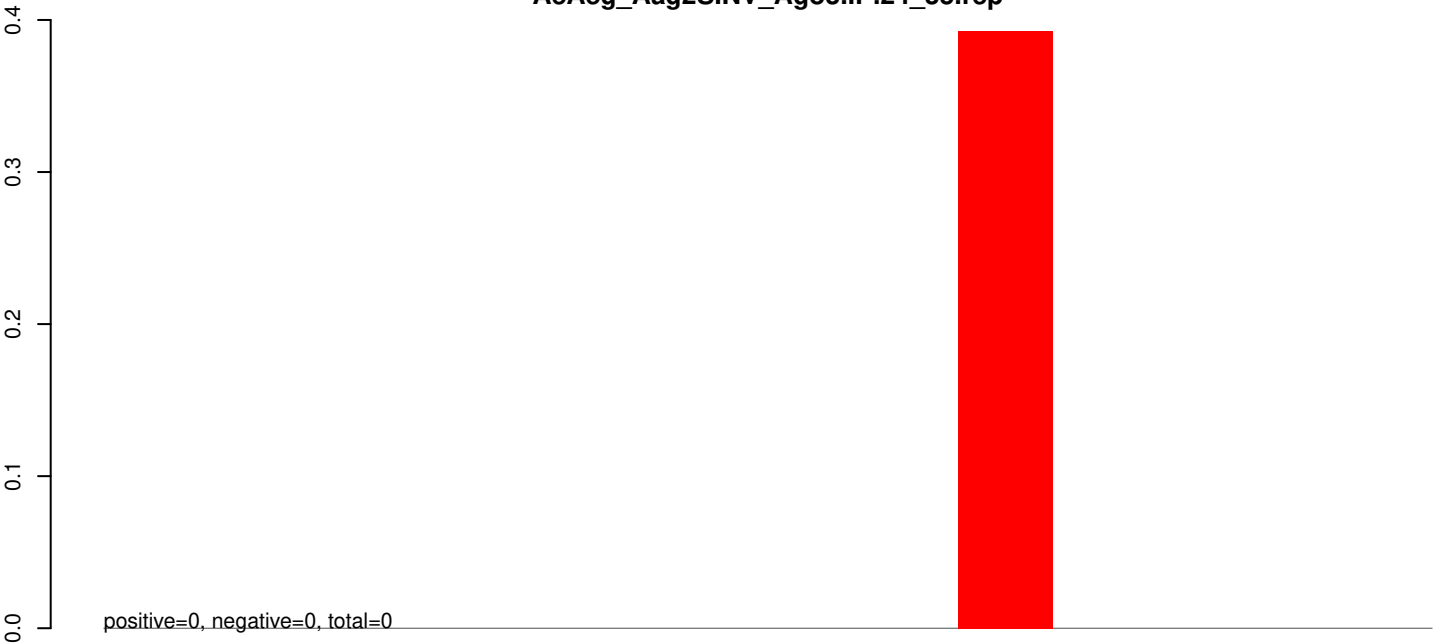
100

150

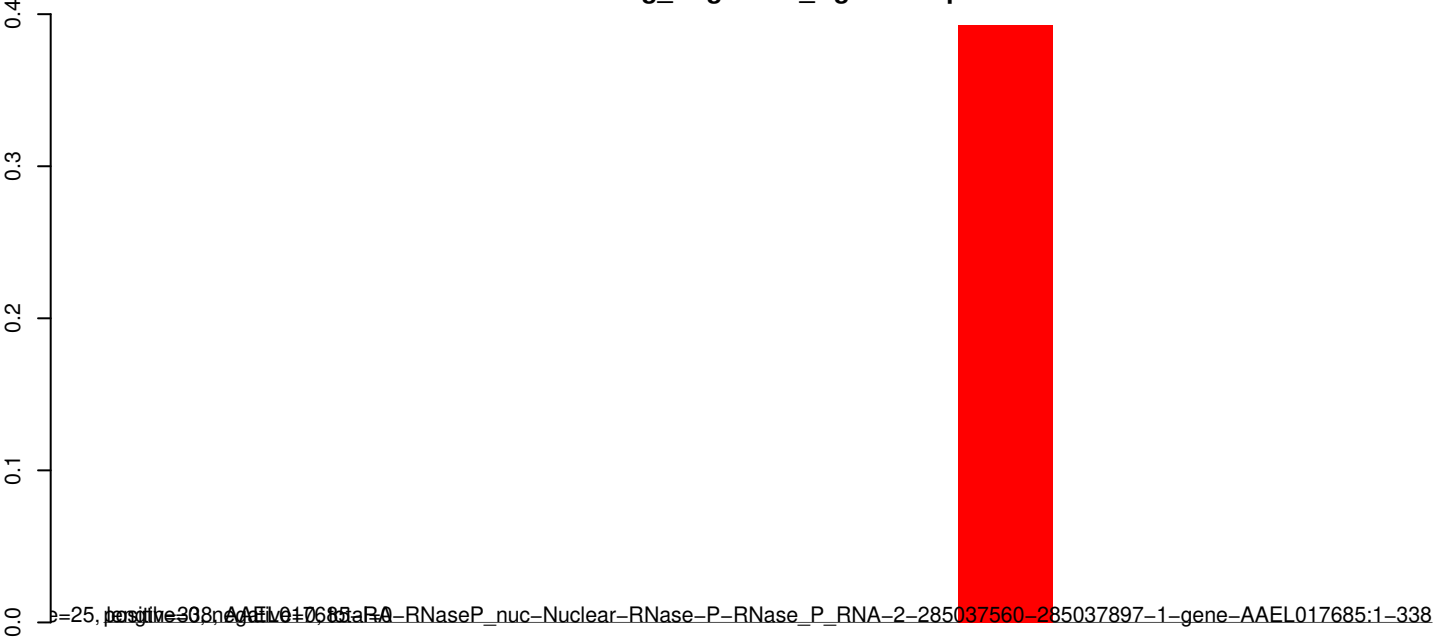
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



AeAeg_Aag2SINV_Ago3.IP.24_35.rep

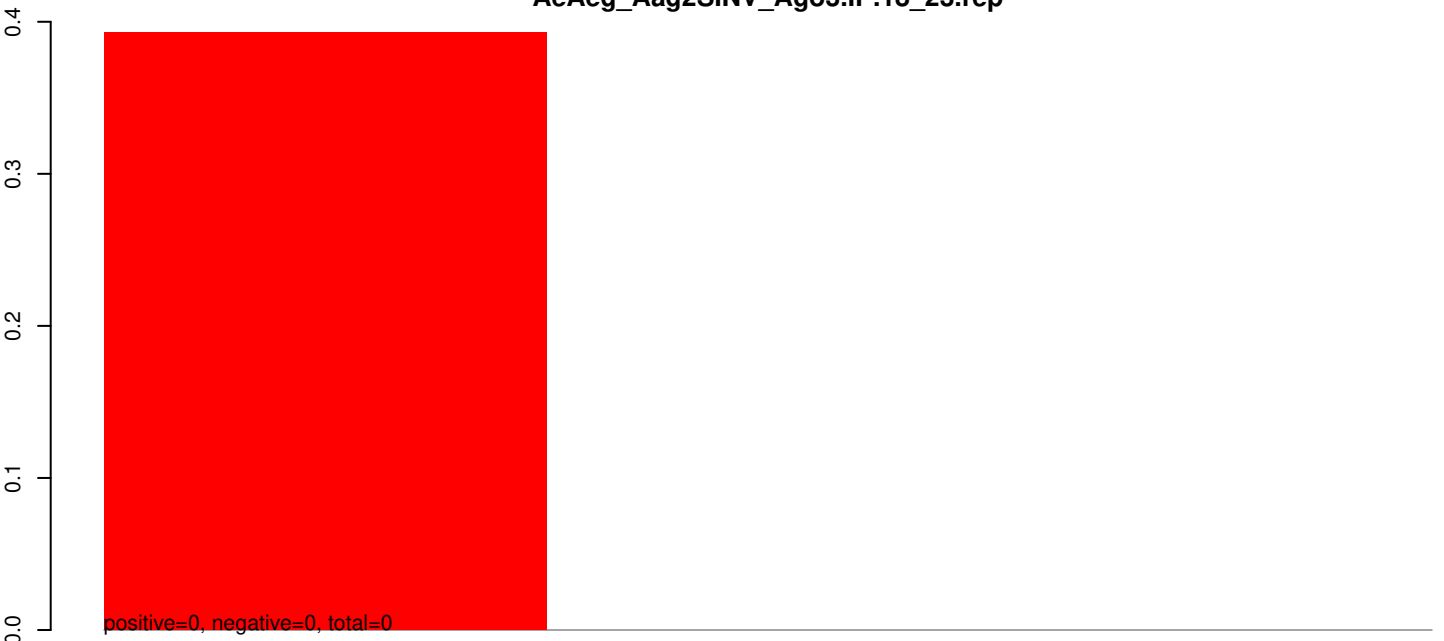


AeAeg_Aag2SINV_Ago3.IP.rep

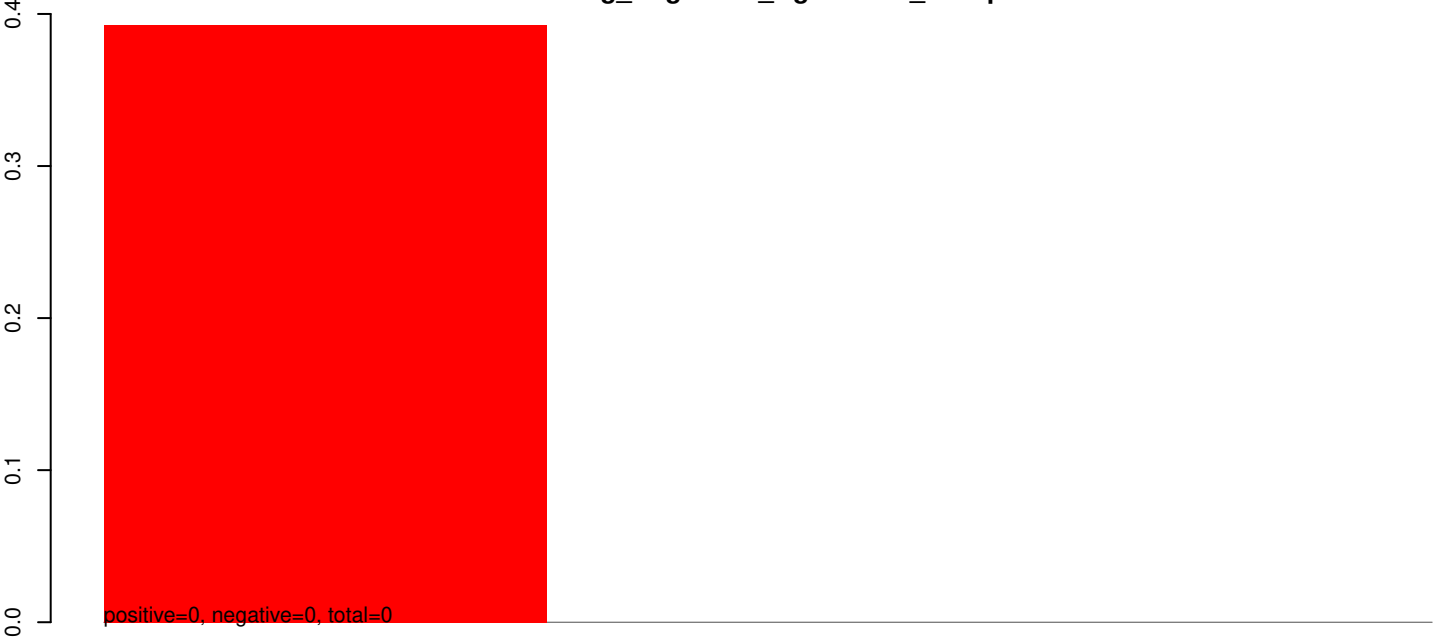


50 100 150 200 250 300 350

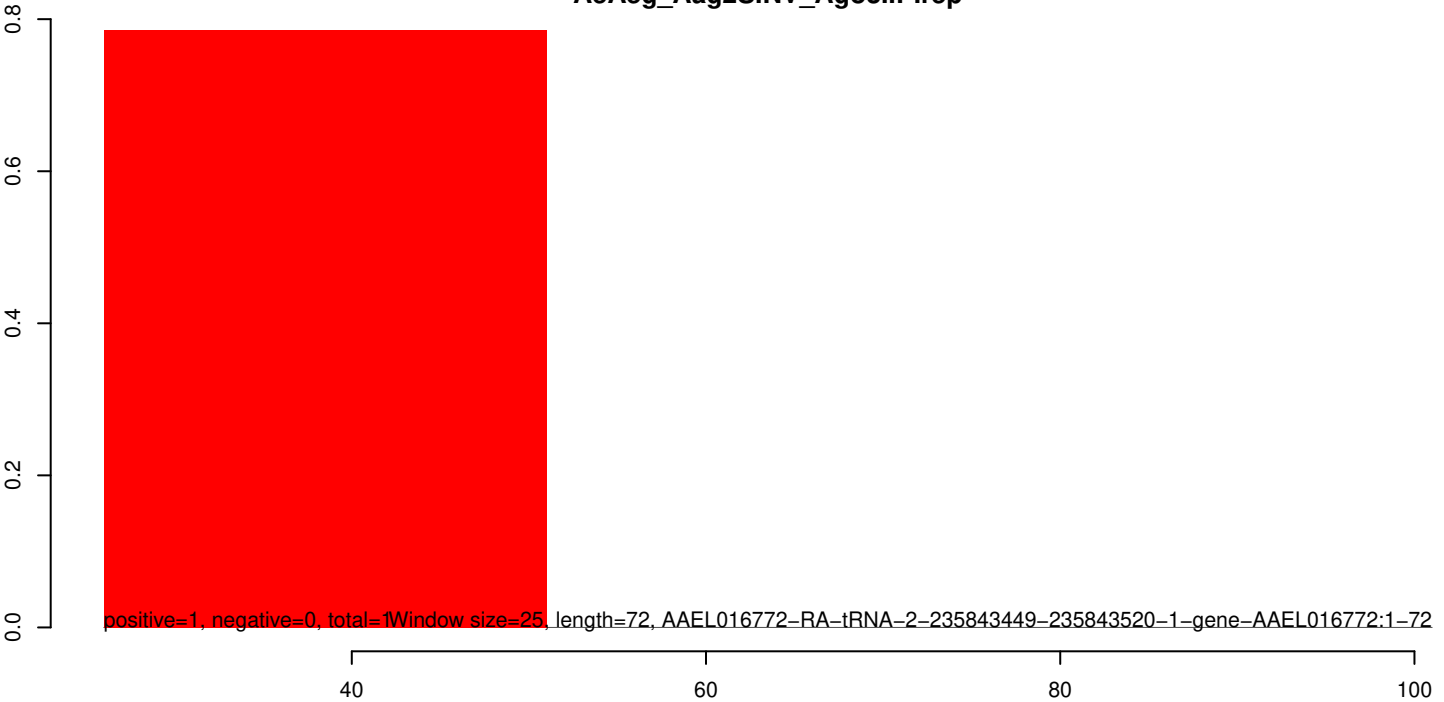
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep



AeAeg_Aag2SINV_Ago3.IP.18_23.rep

positive=2, negative=0, total=2

AeAeg_Aag2SINV_Ago3.IP.24_35.rep

positive=2, negative=0, total=2

AeAeg_Aag2SINV_Ago3.IP.rep

positive=3, negative=0, total=3 Window size=25, length=73, AAEL016098-RA-tRNA-2-56316242-56316314-1-gene-AAEL016098:1-73

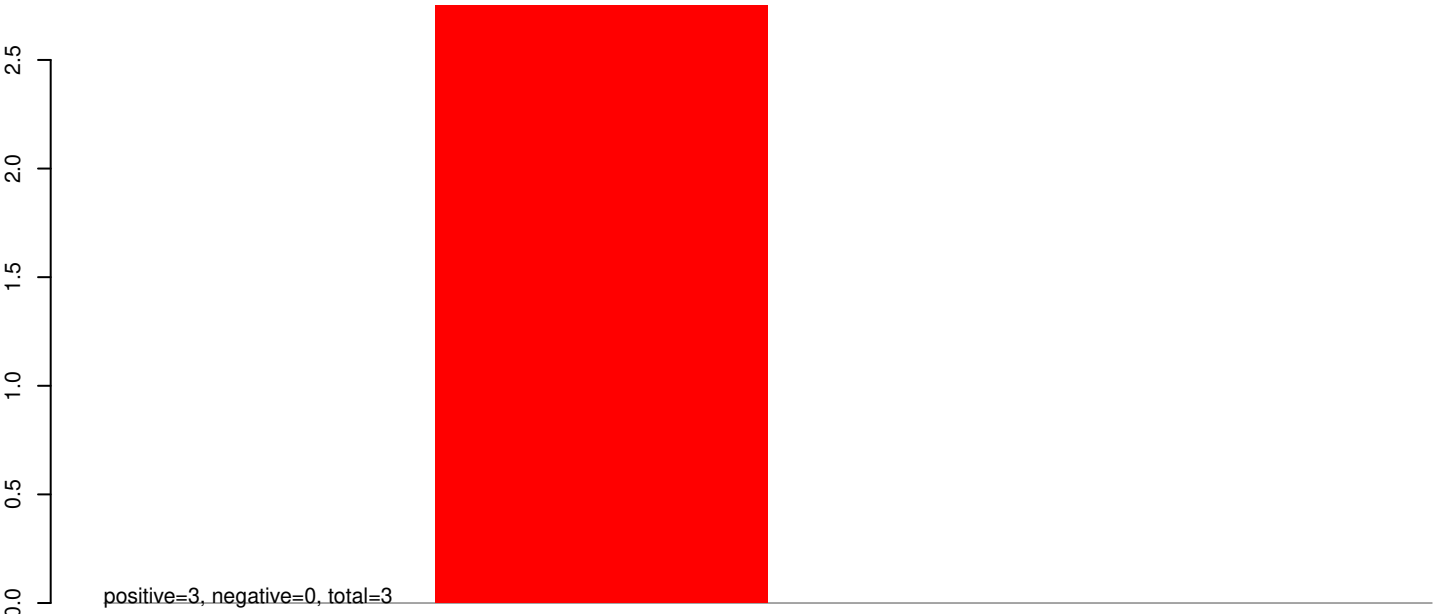
40

60

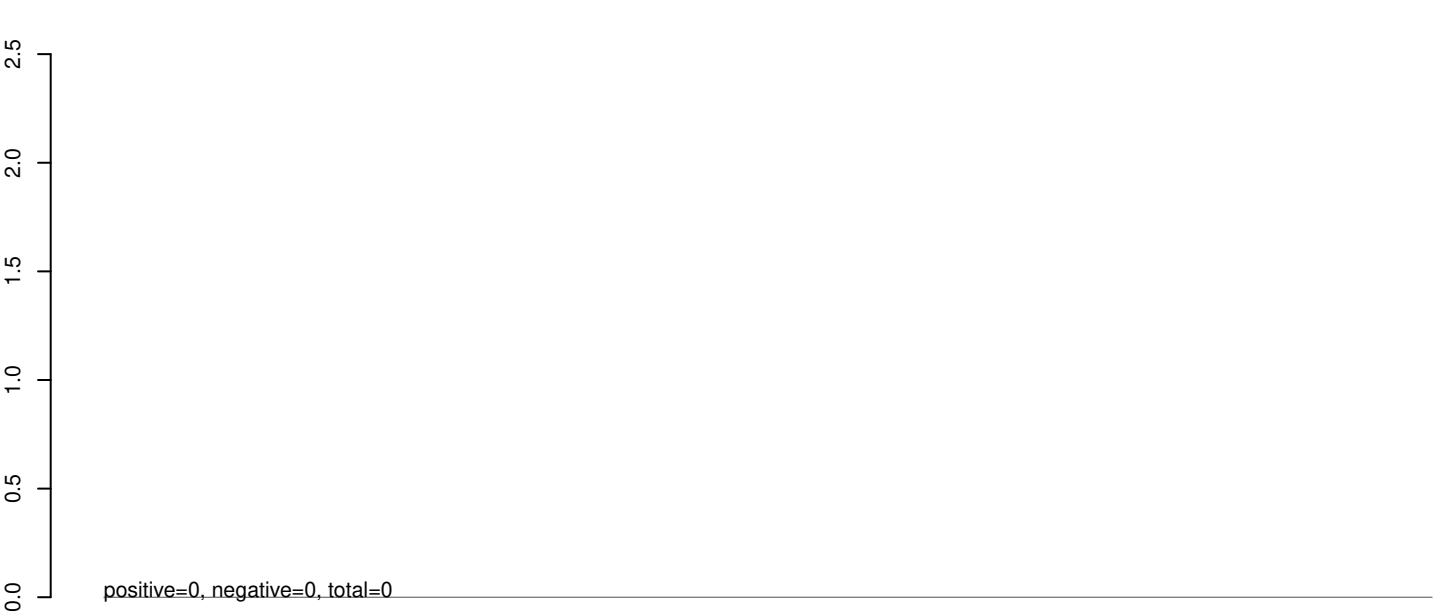
80

100

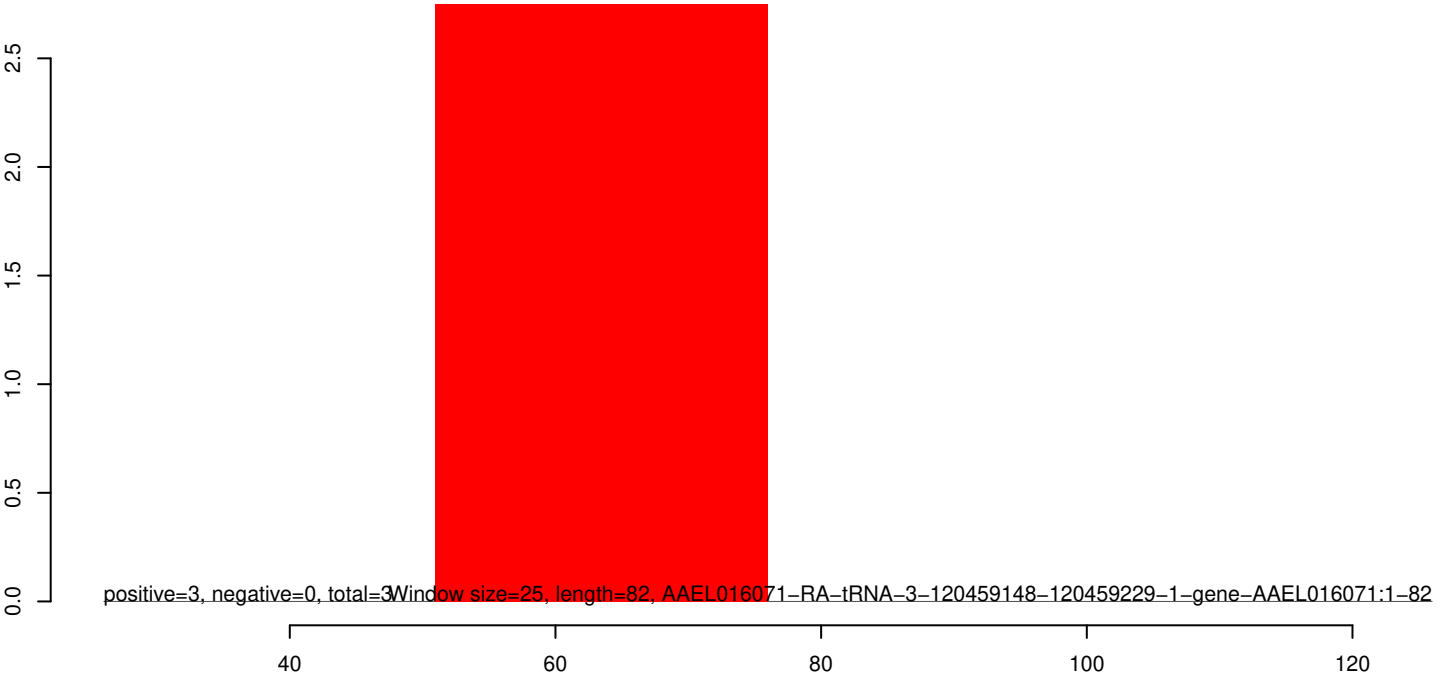
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



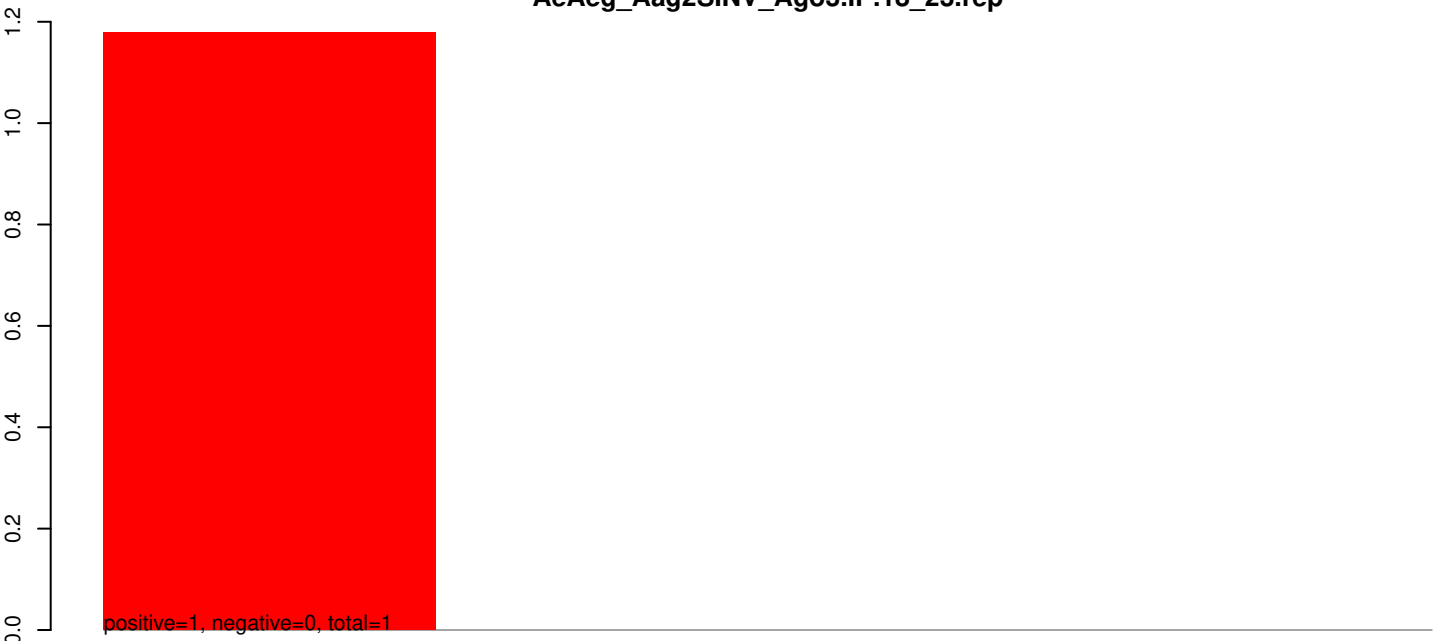
AeAeg_Aag2SINV_Ago3.IP.24_35.rep



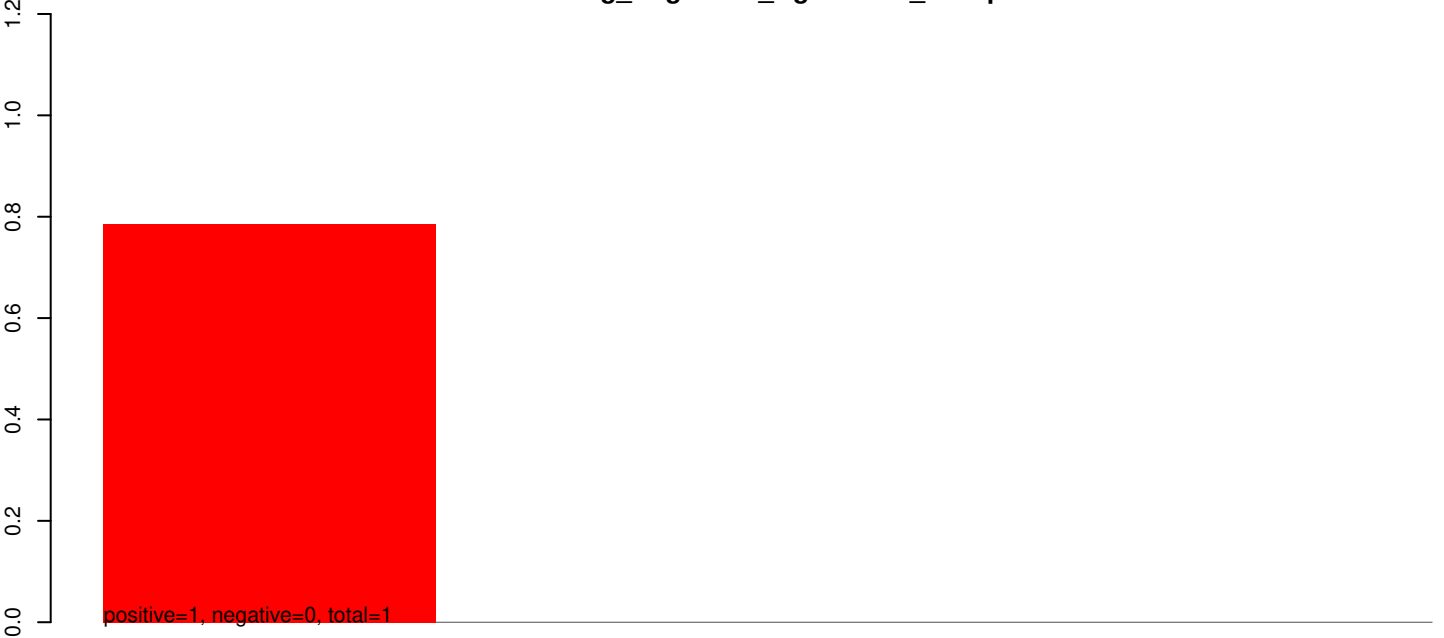
AeAeg_Aag2SINV_Ago3.IP.rep



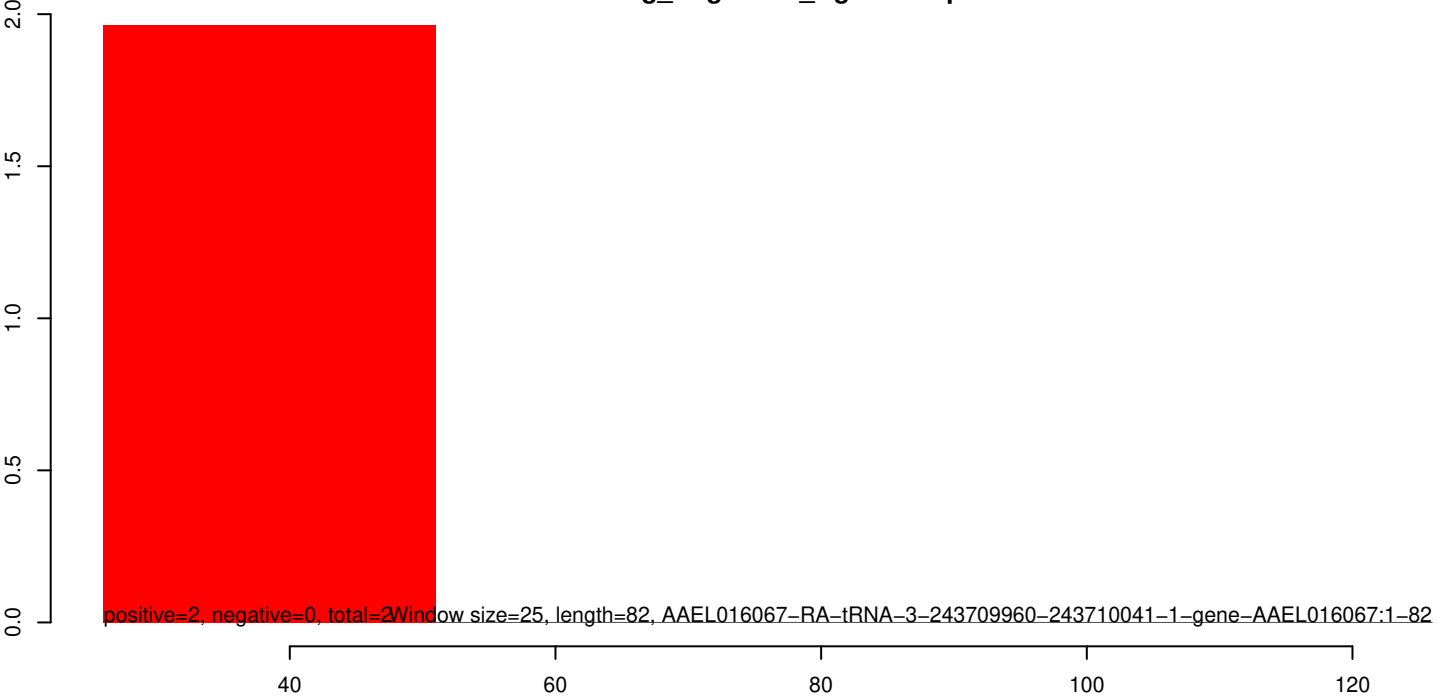
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



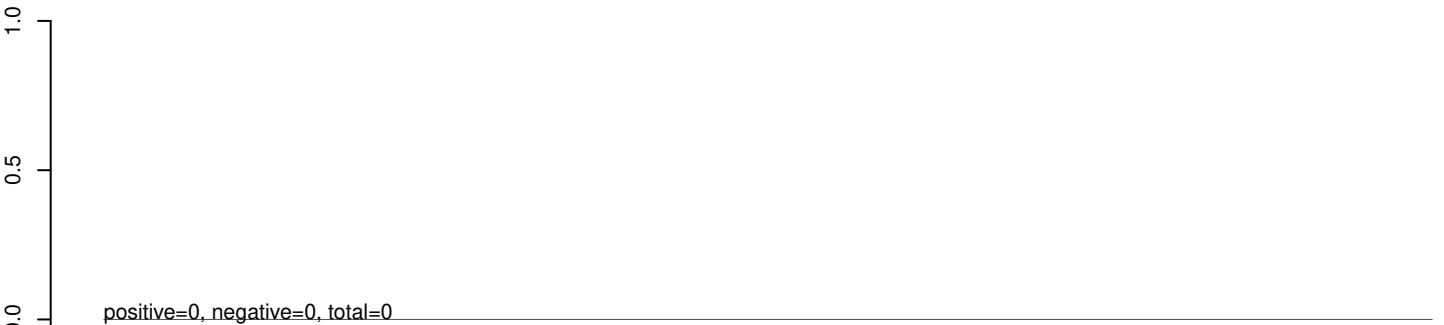
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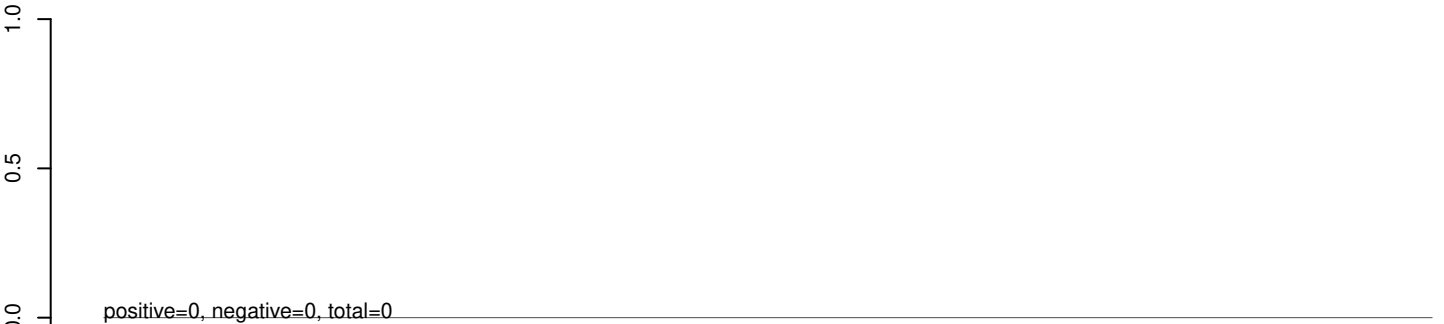
AeAeg_Aag2SINV_Ago3.IP.rep



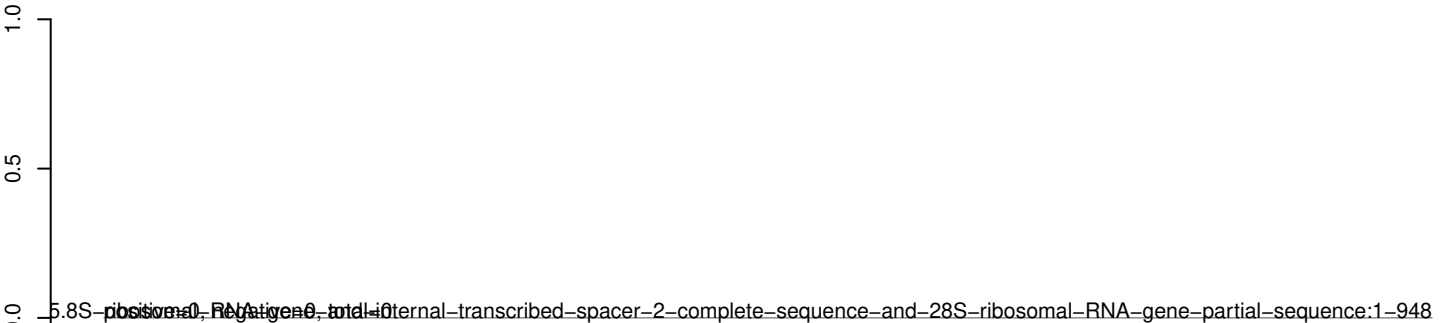
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep



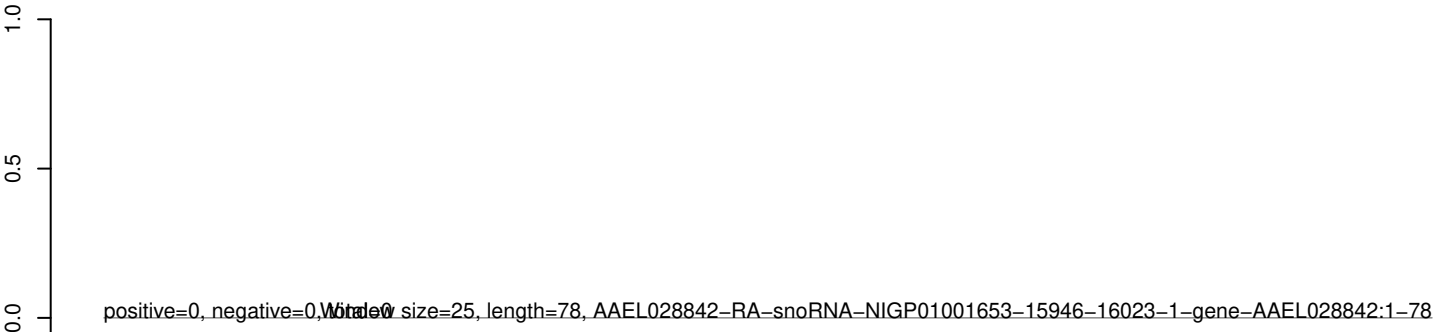
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep



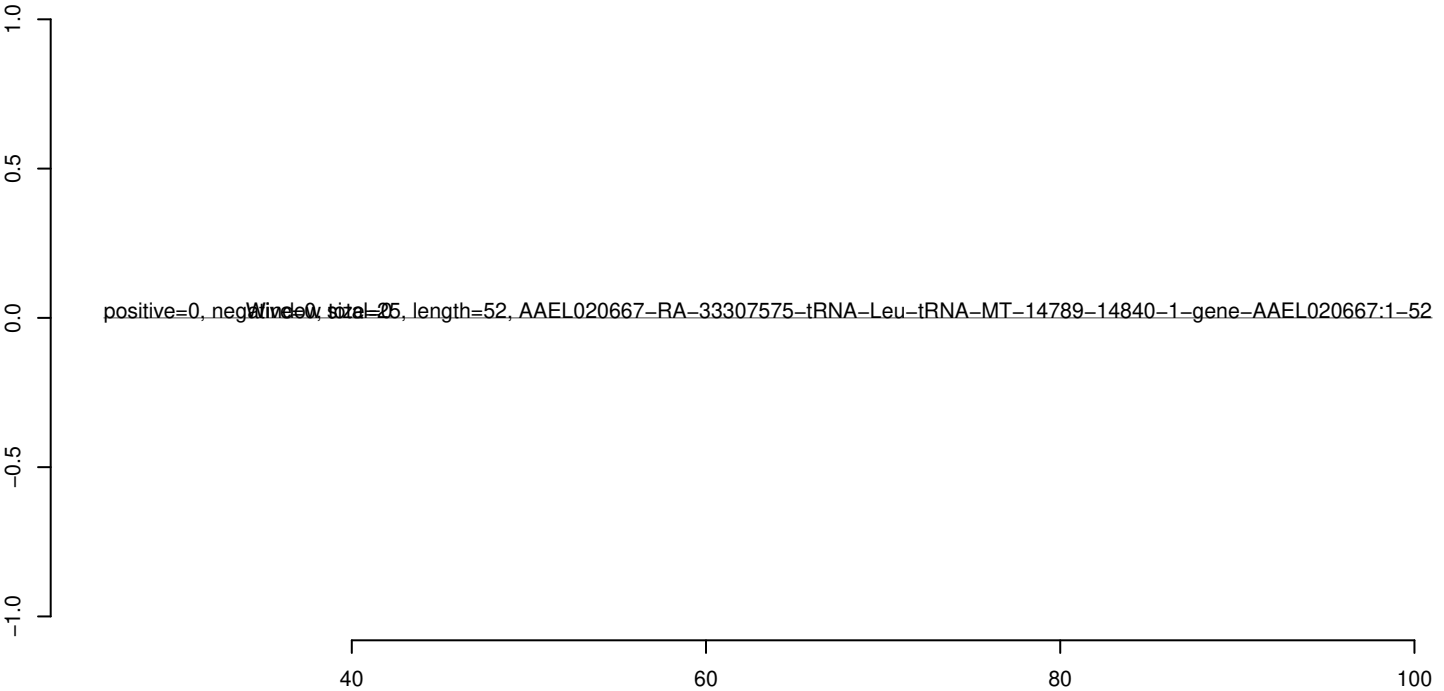
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



AeAeg_Aag2SINV_Ago3.IP.24_35.rep



AeAeg_Aag2SINV_Ago3.IP.rep



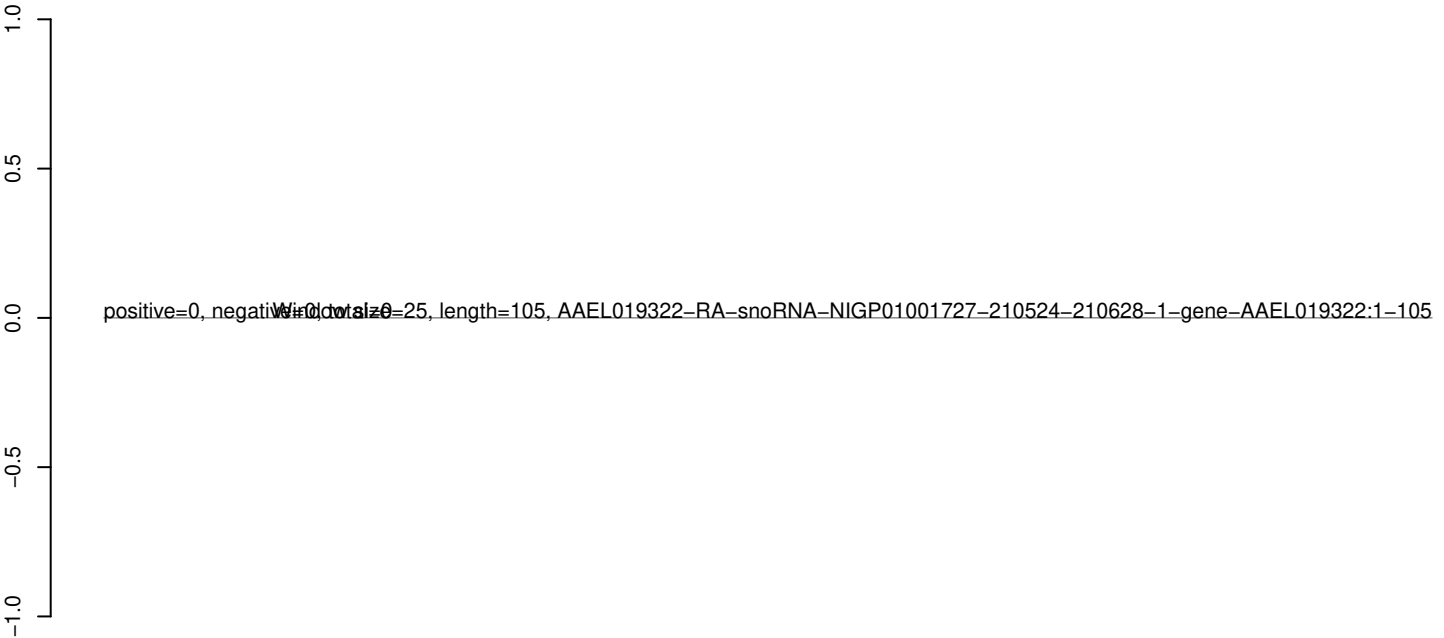
AeAeg_Aag2SINV_Ago3.IP.18_23.rep



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AeAeg_Aag2SINV_Ago3.IP.rep



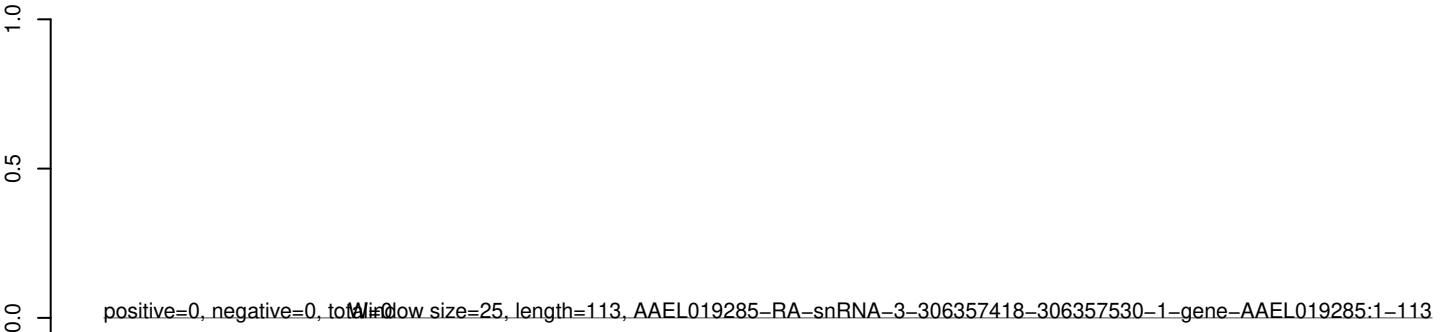
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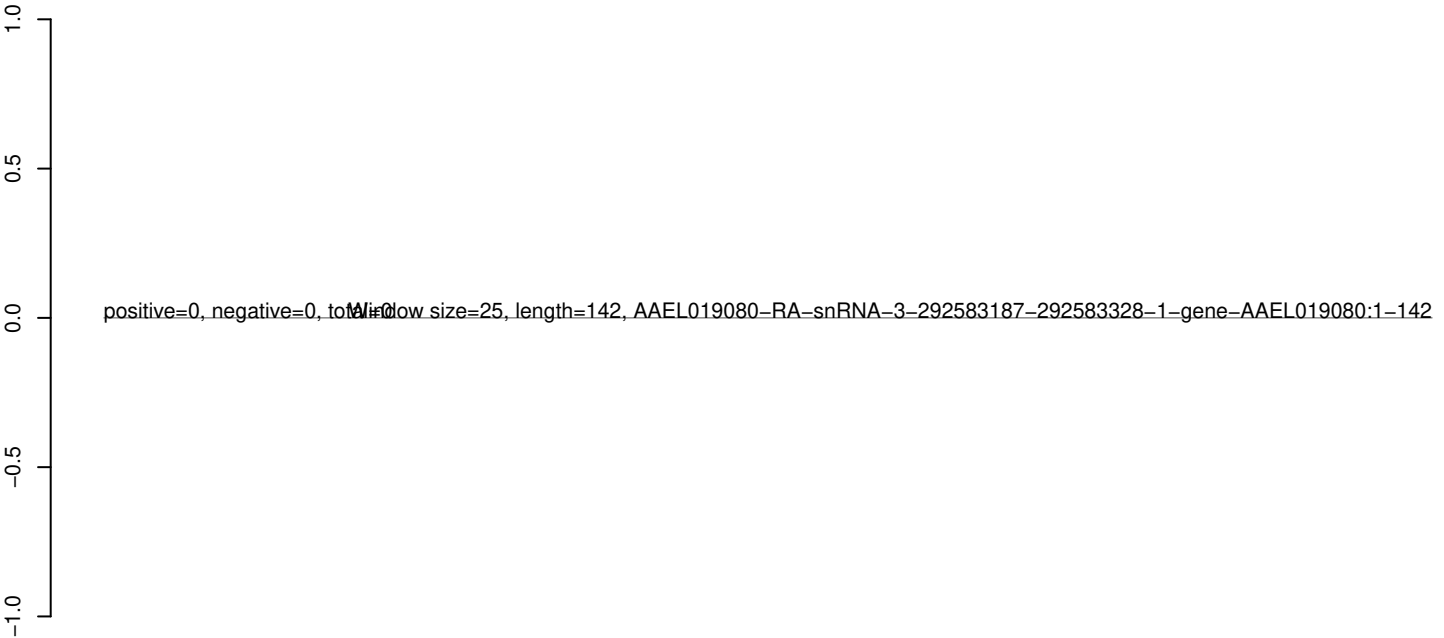
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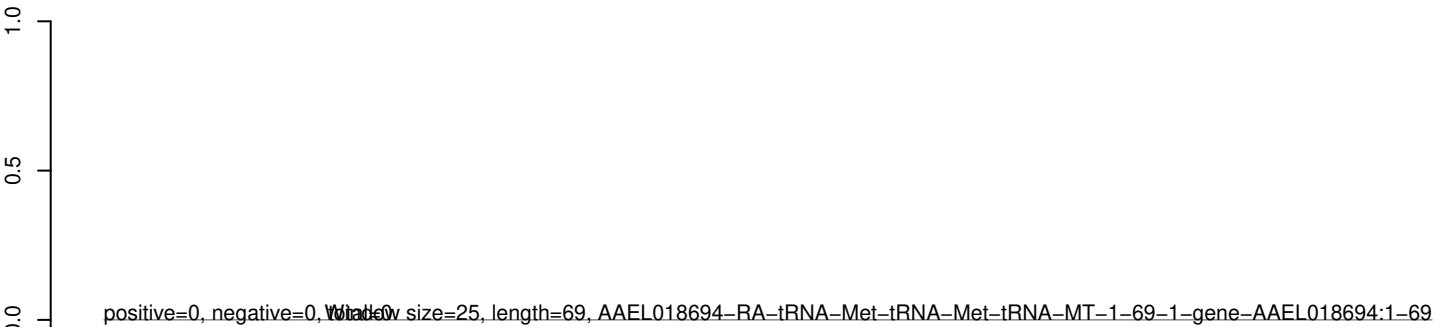
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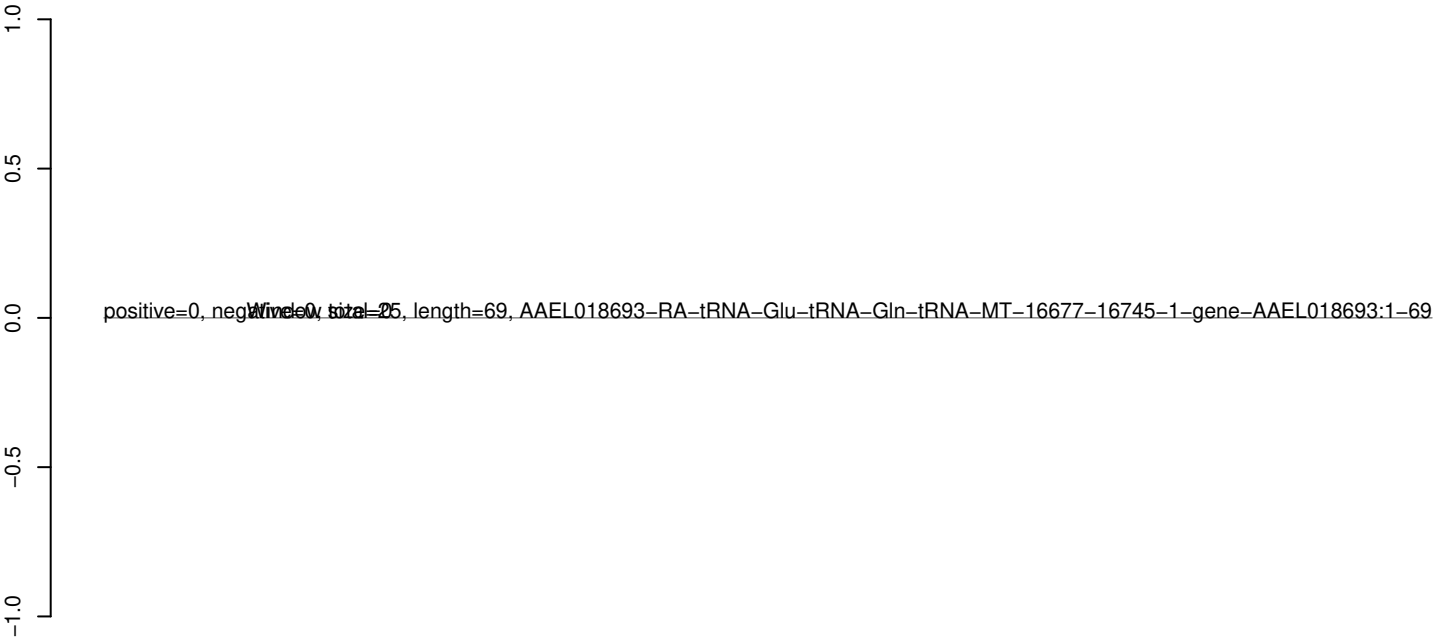
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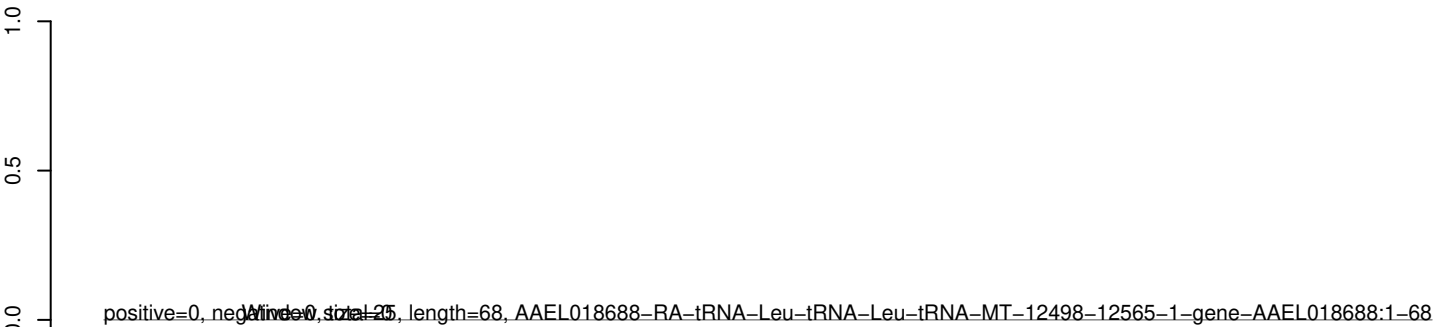
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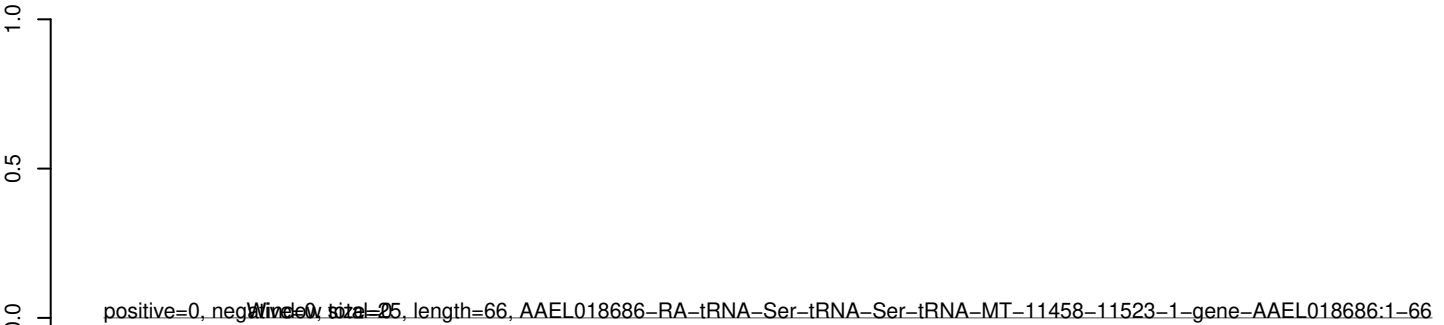
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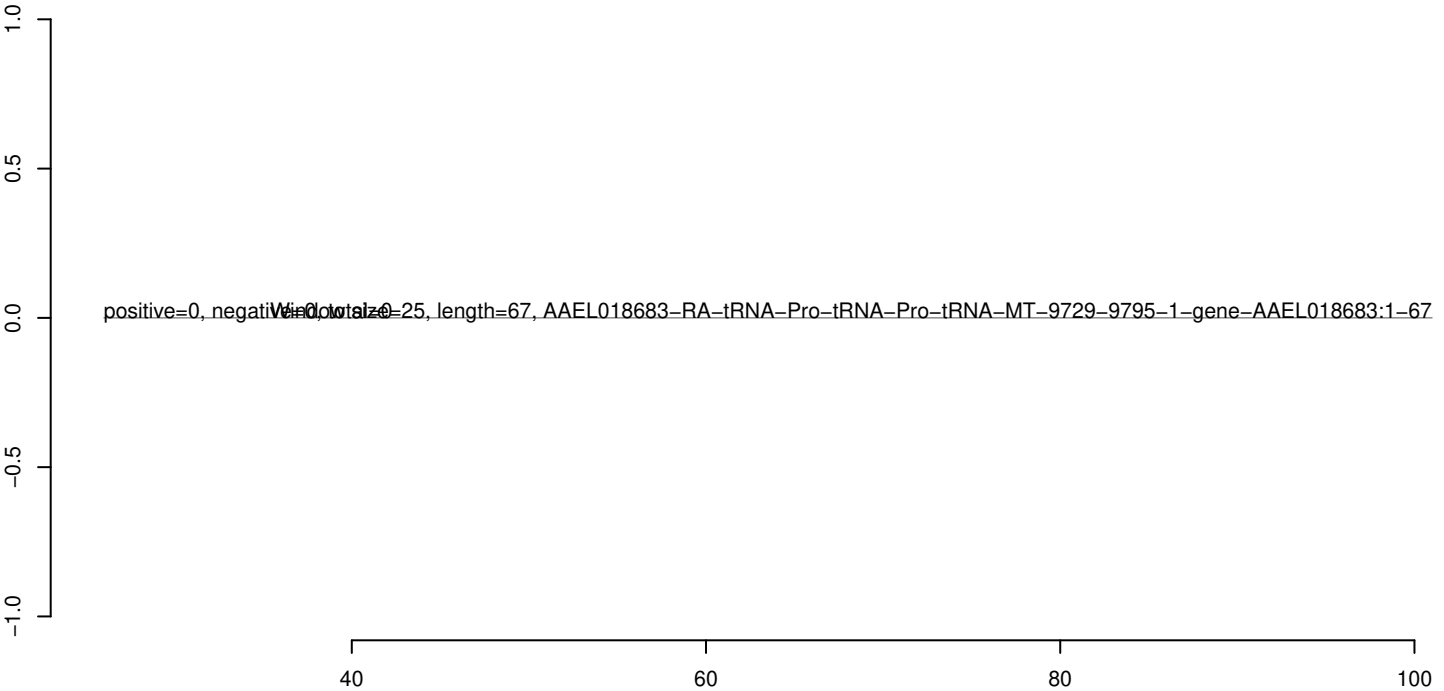
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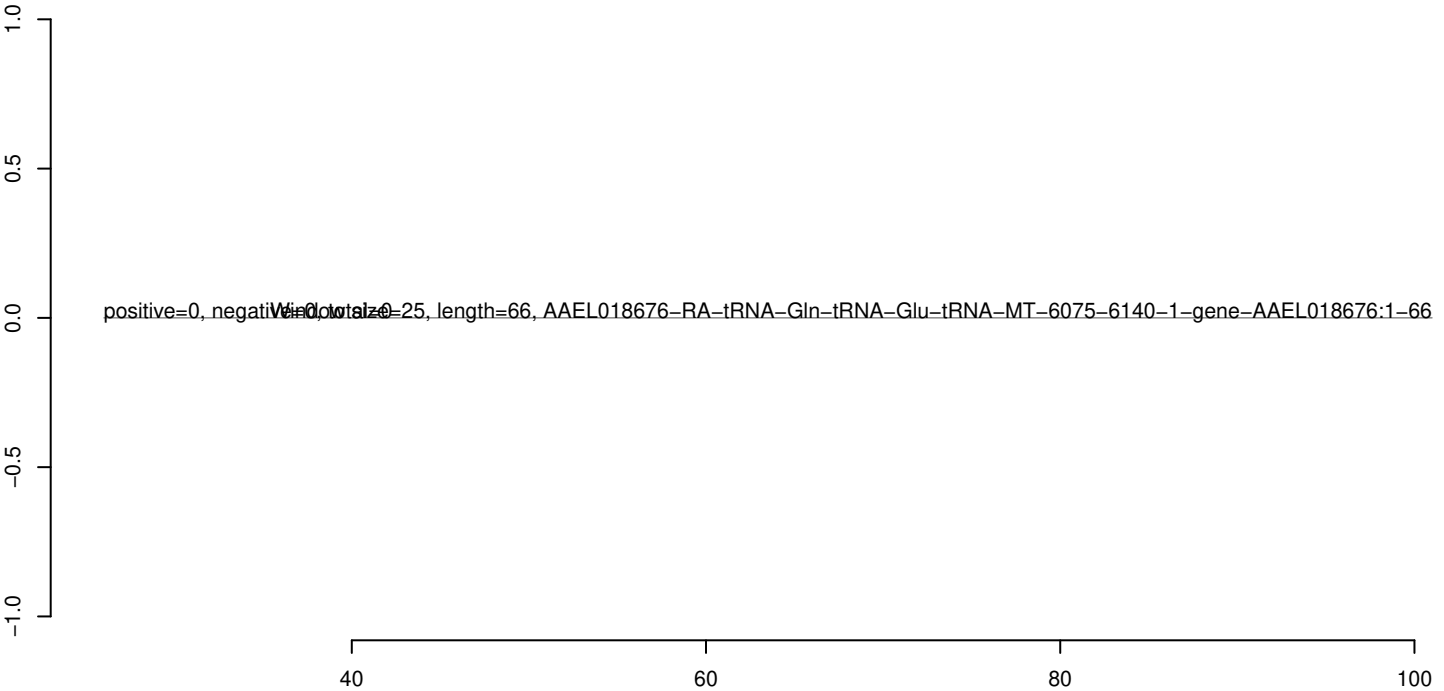
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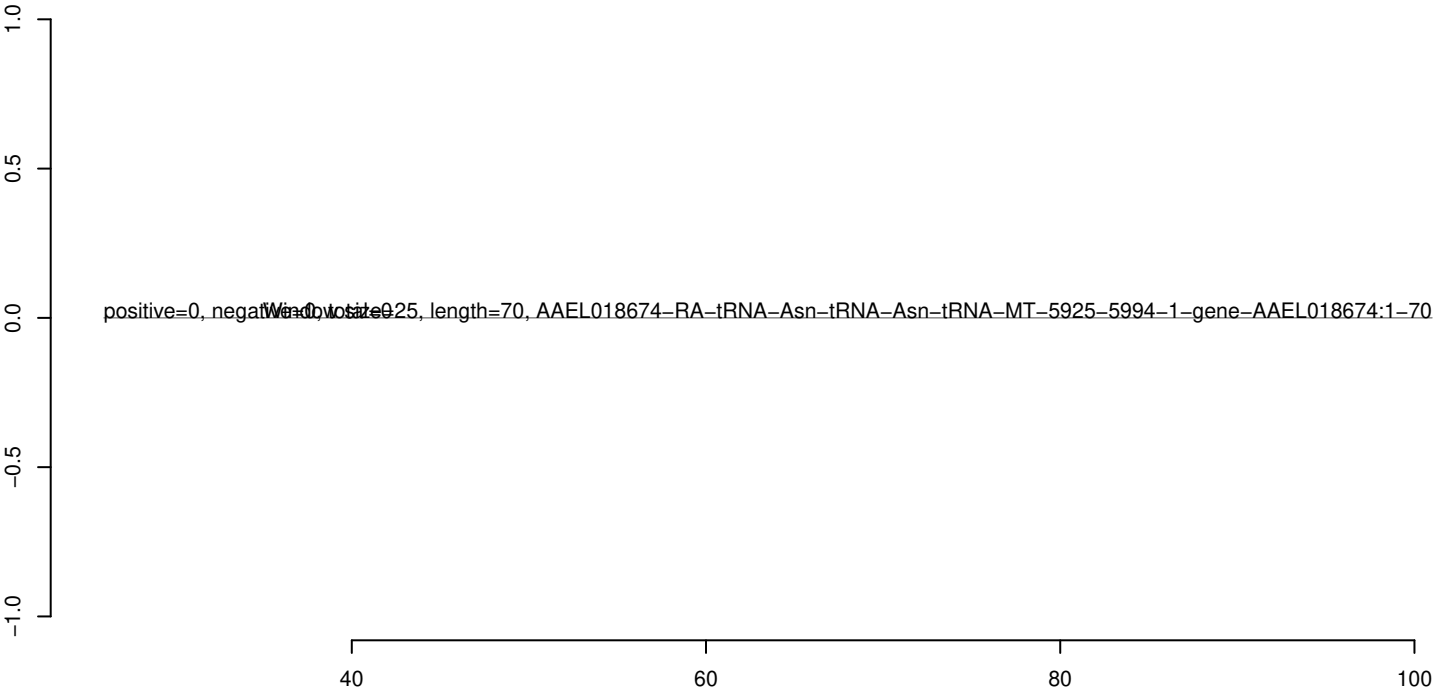
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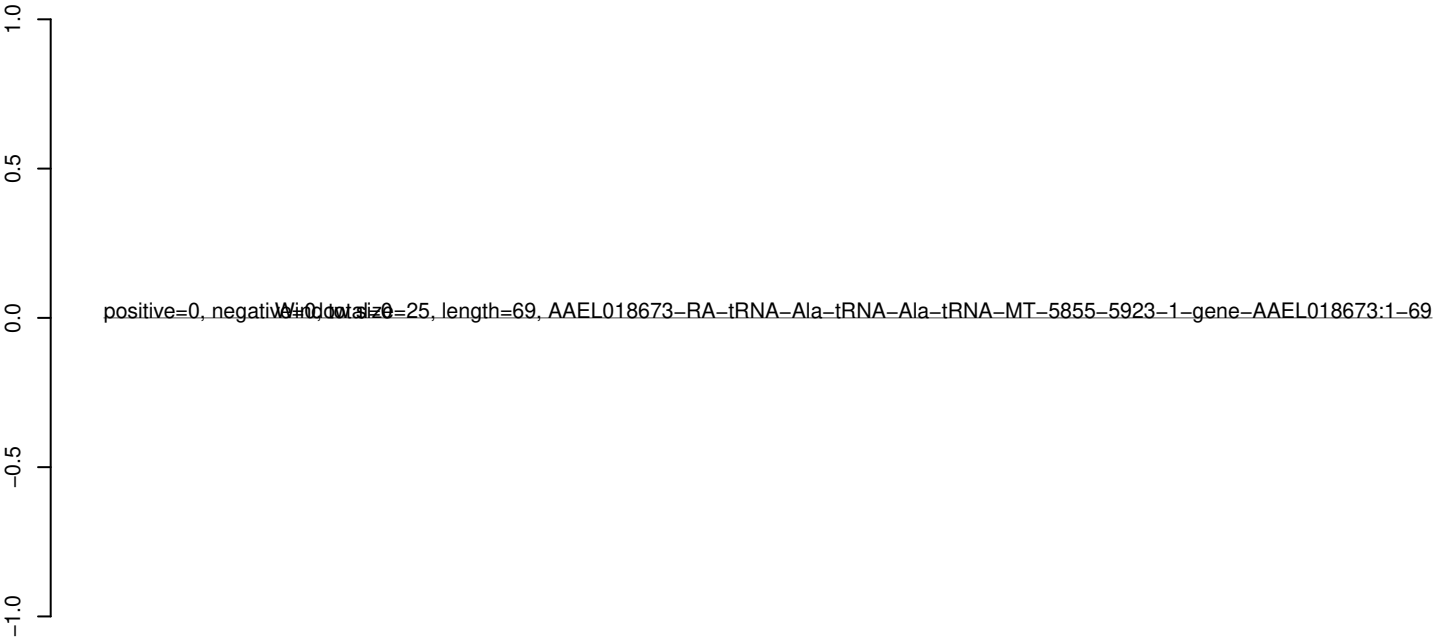
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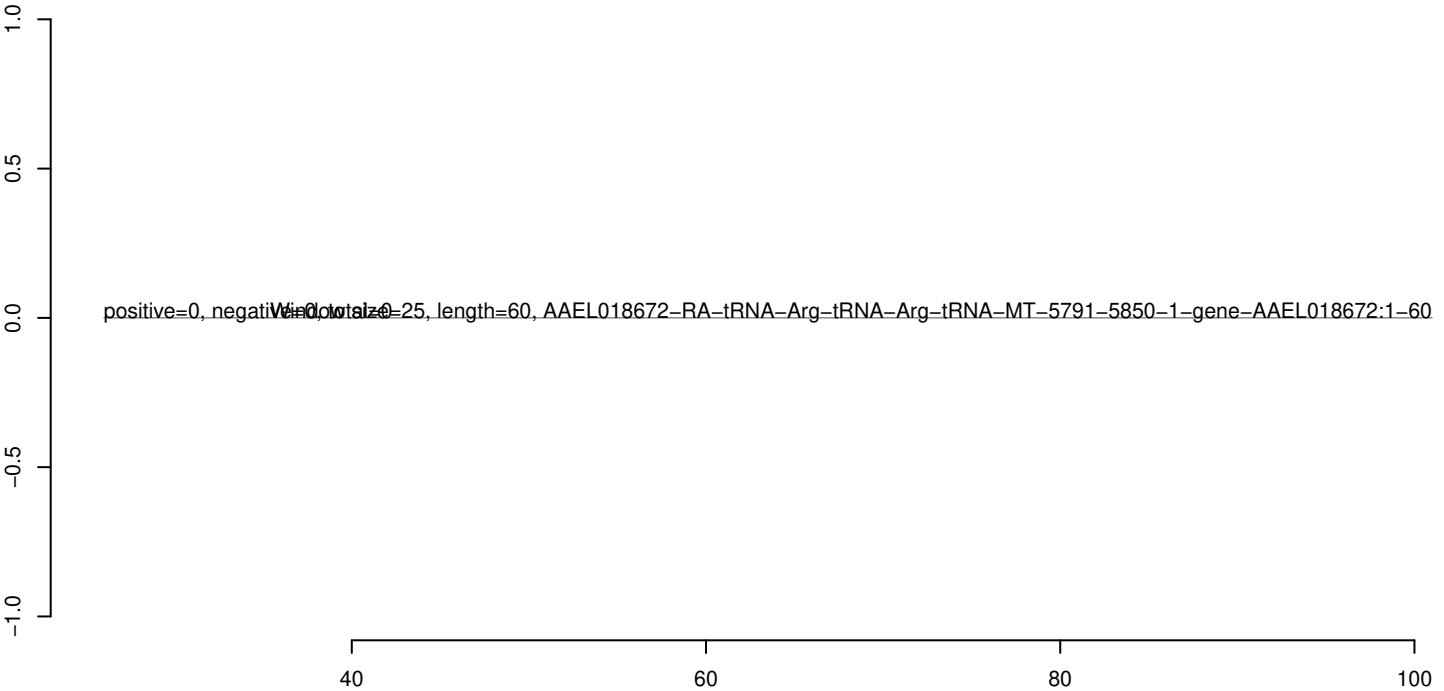
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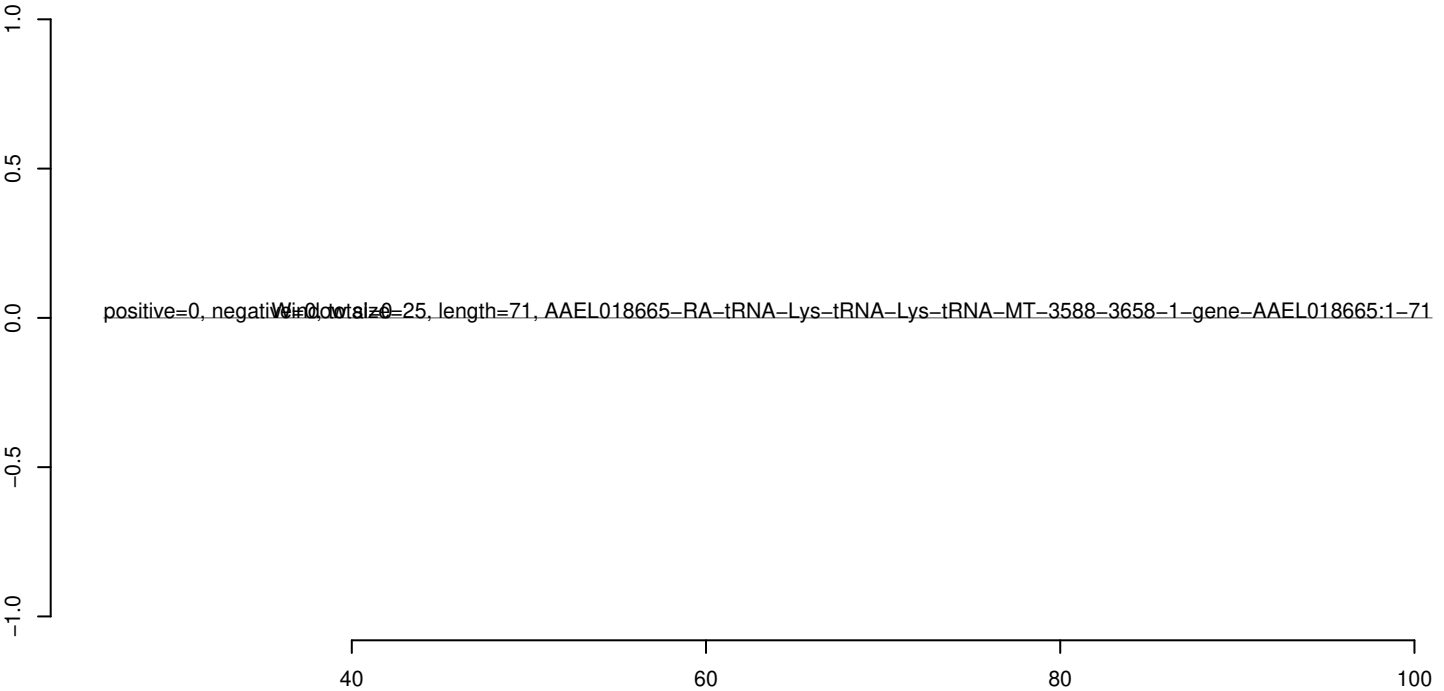
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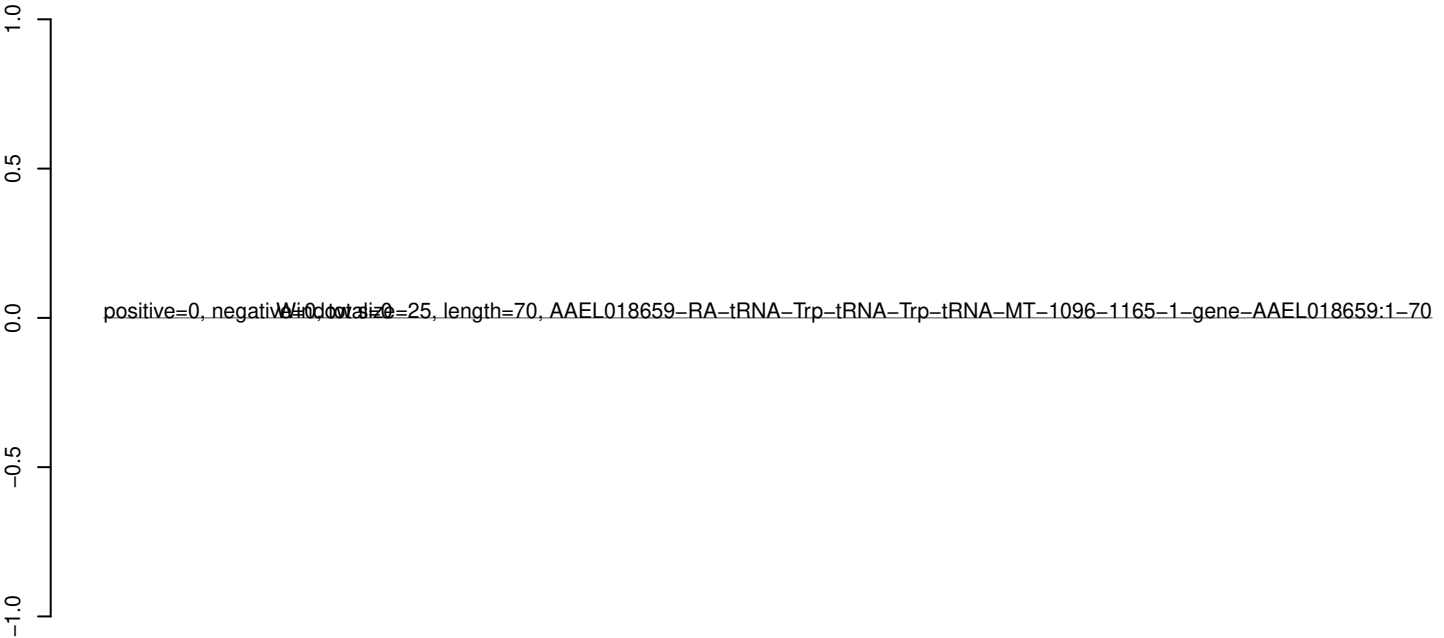
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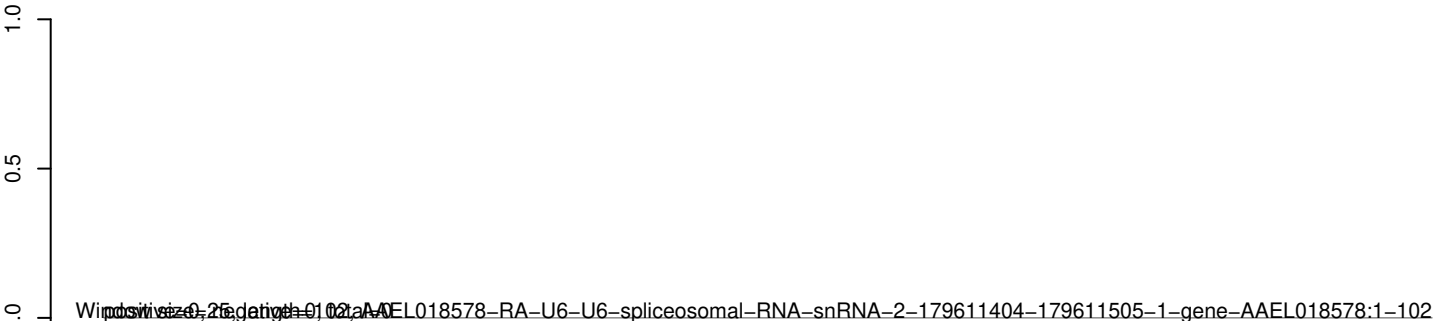
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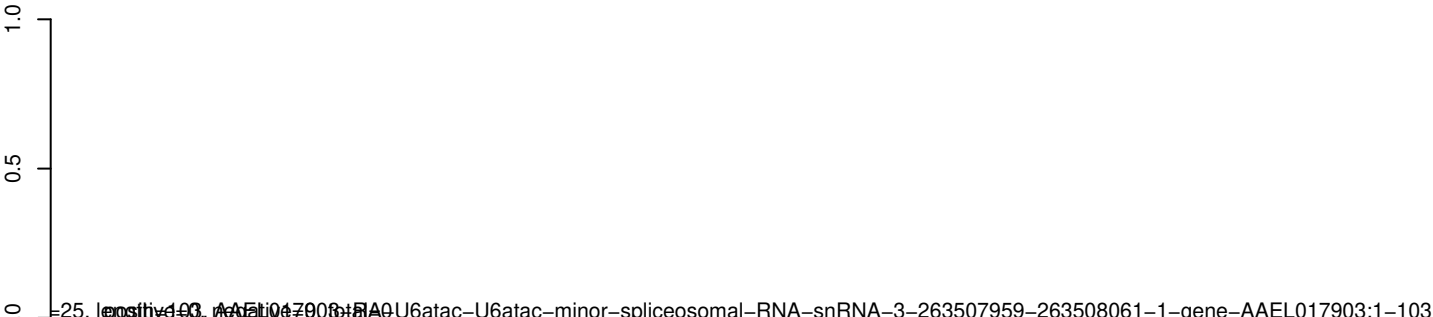
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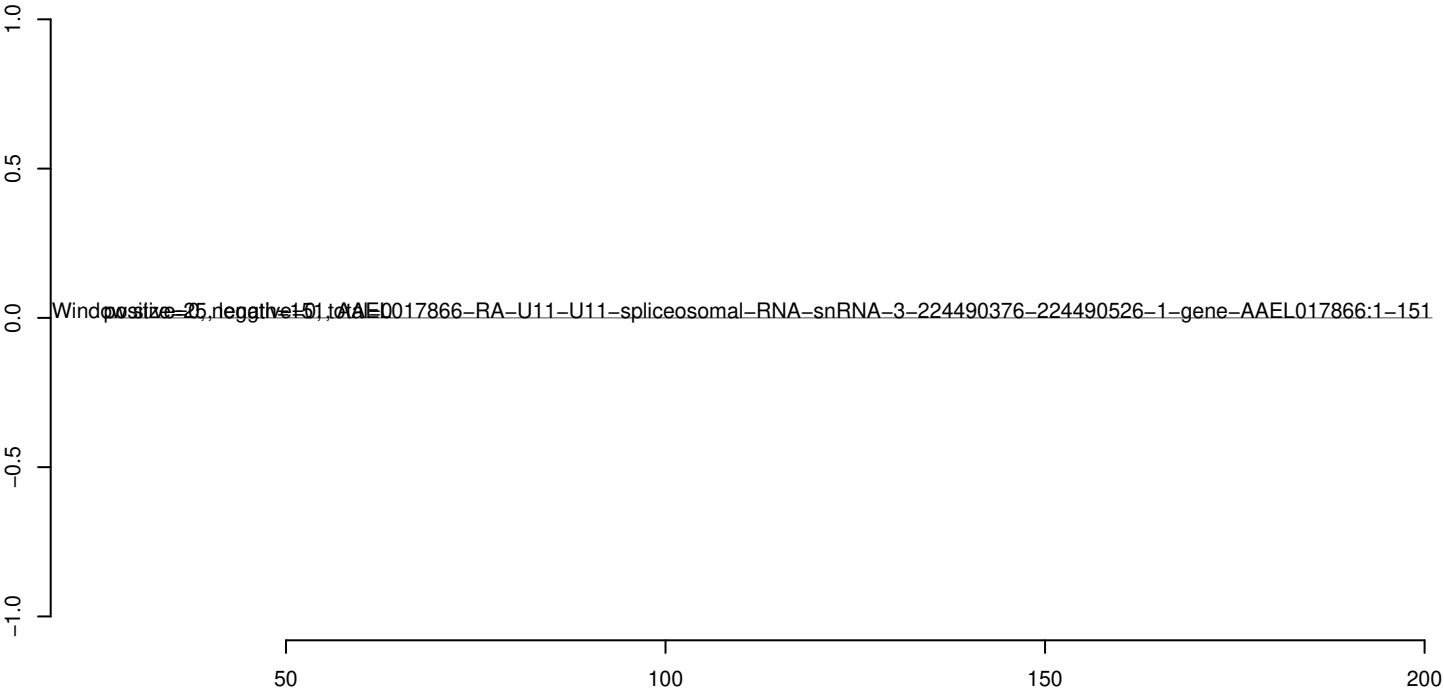
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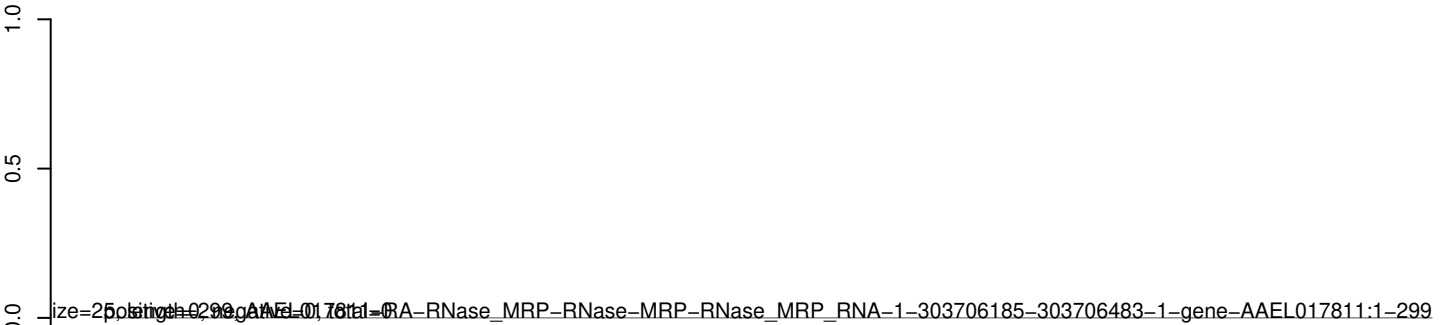
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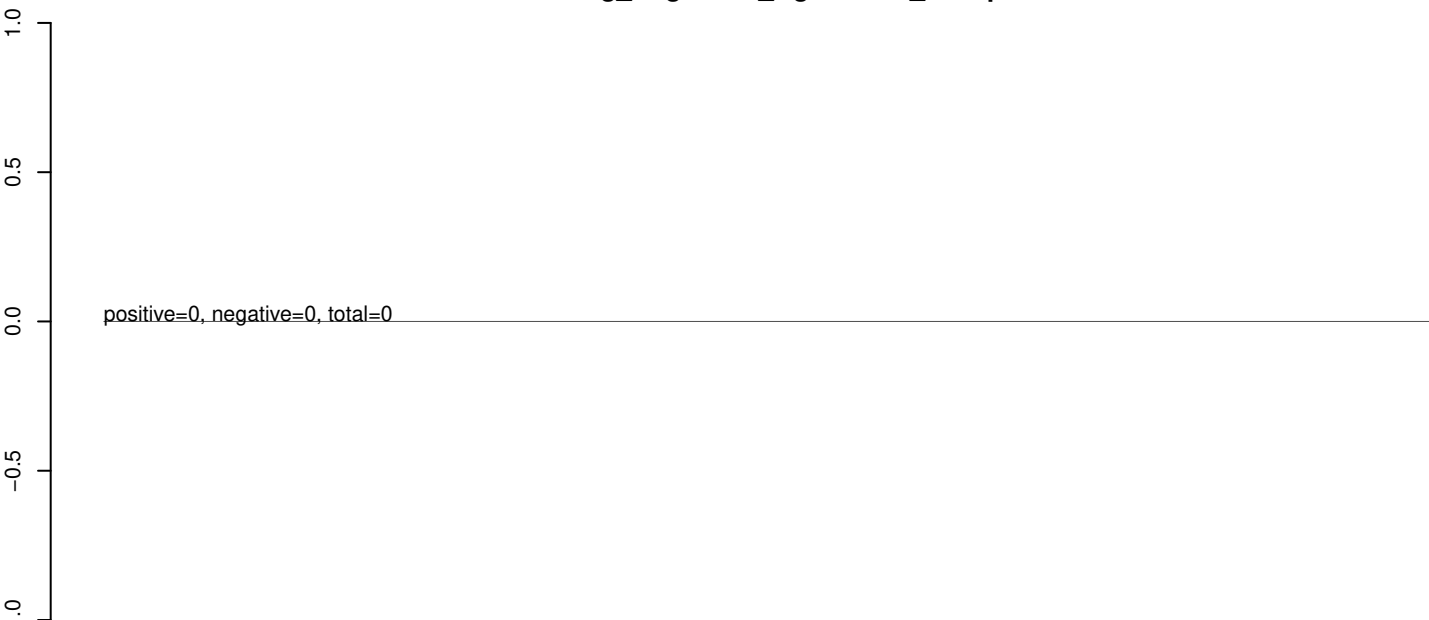
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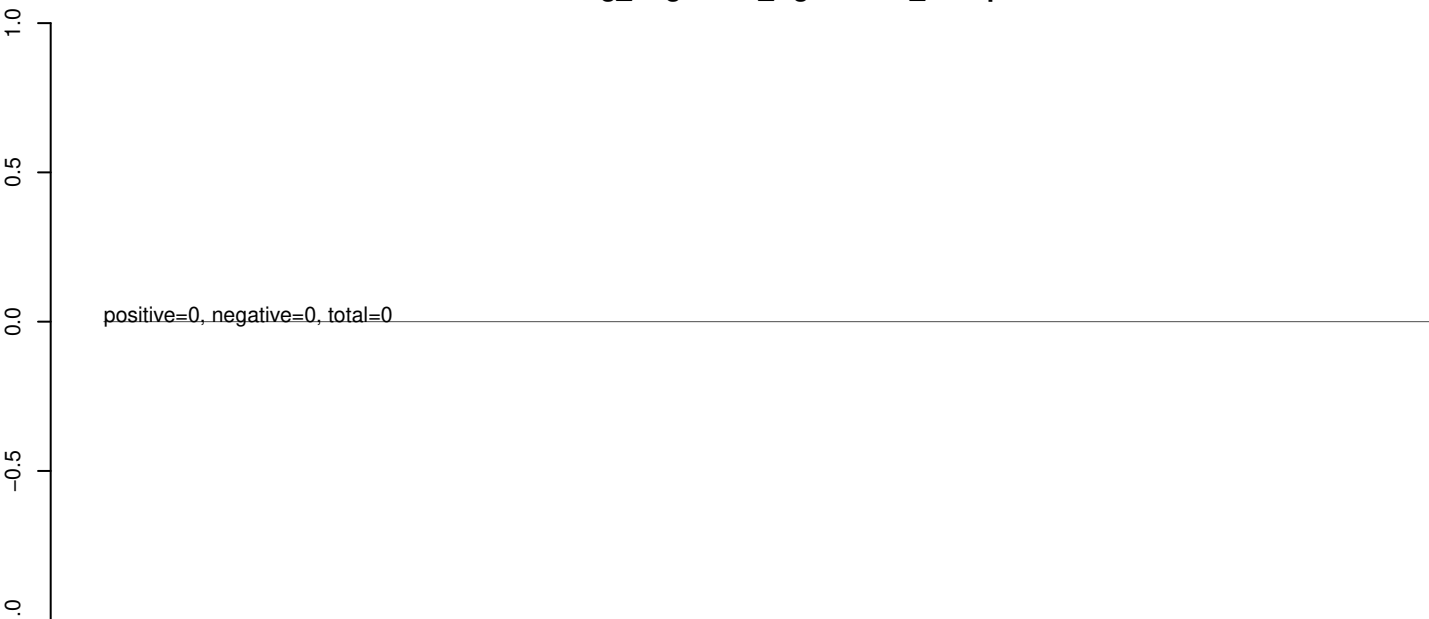
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