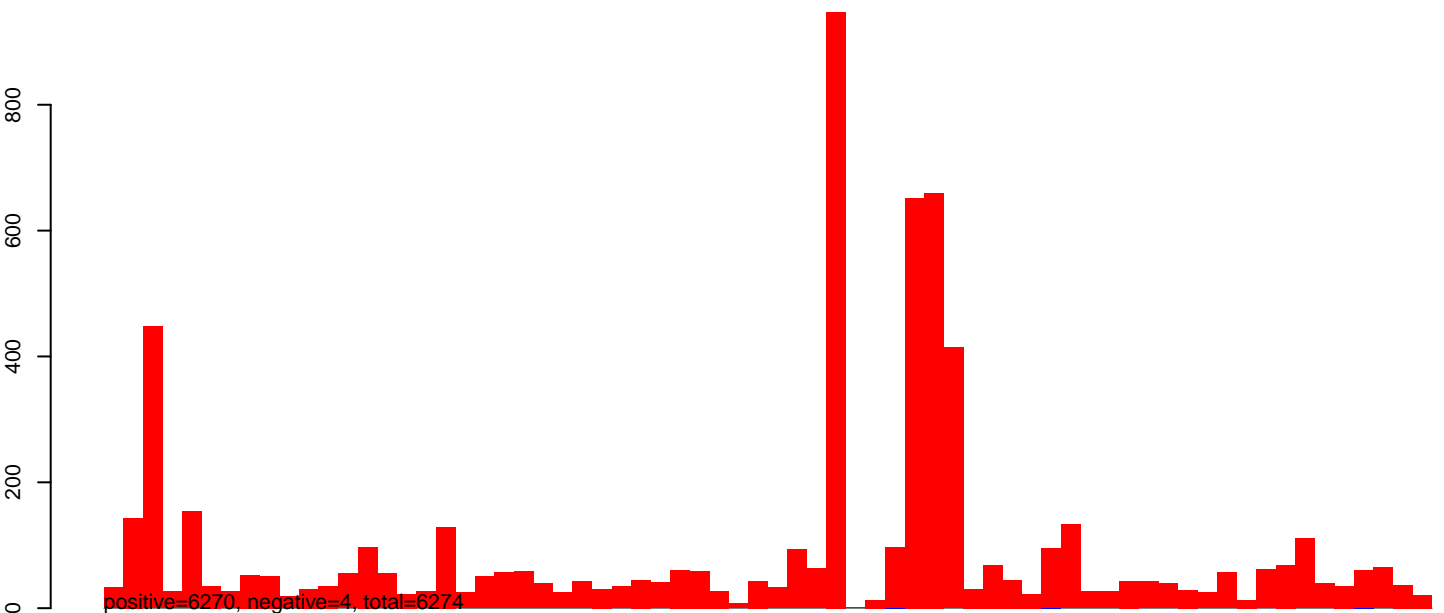
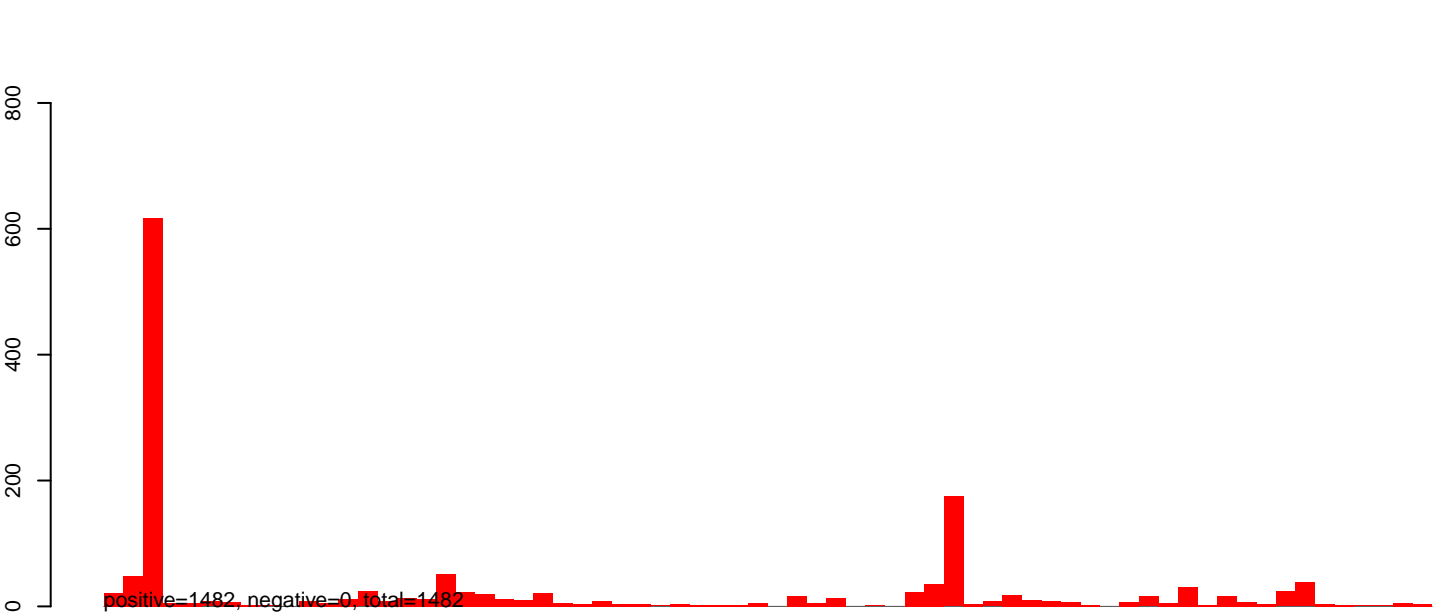


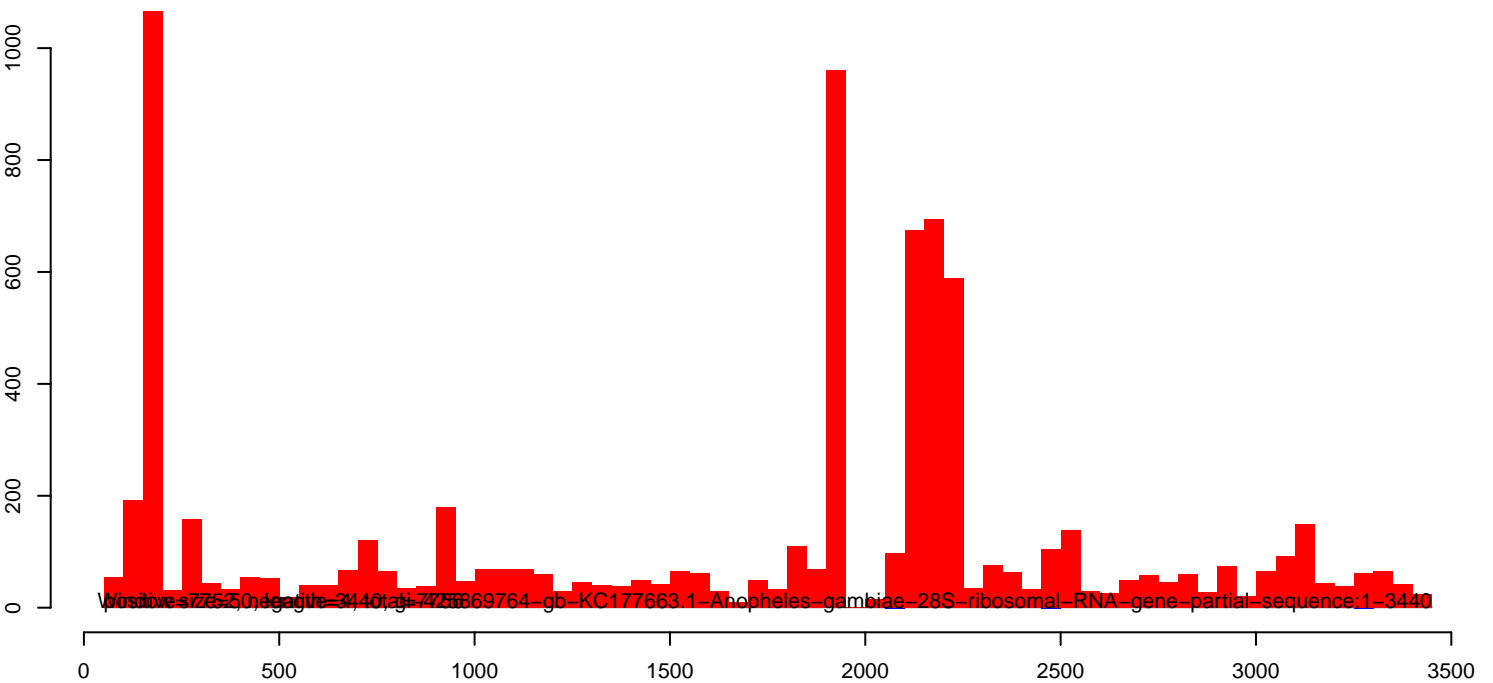
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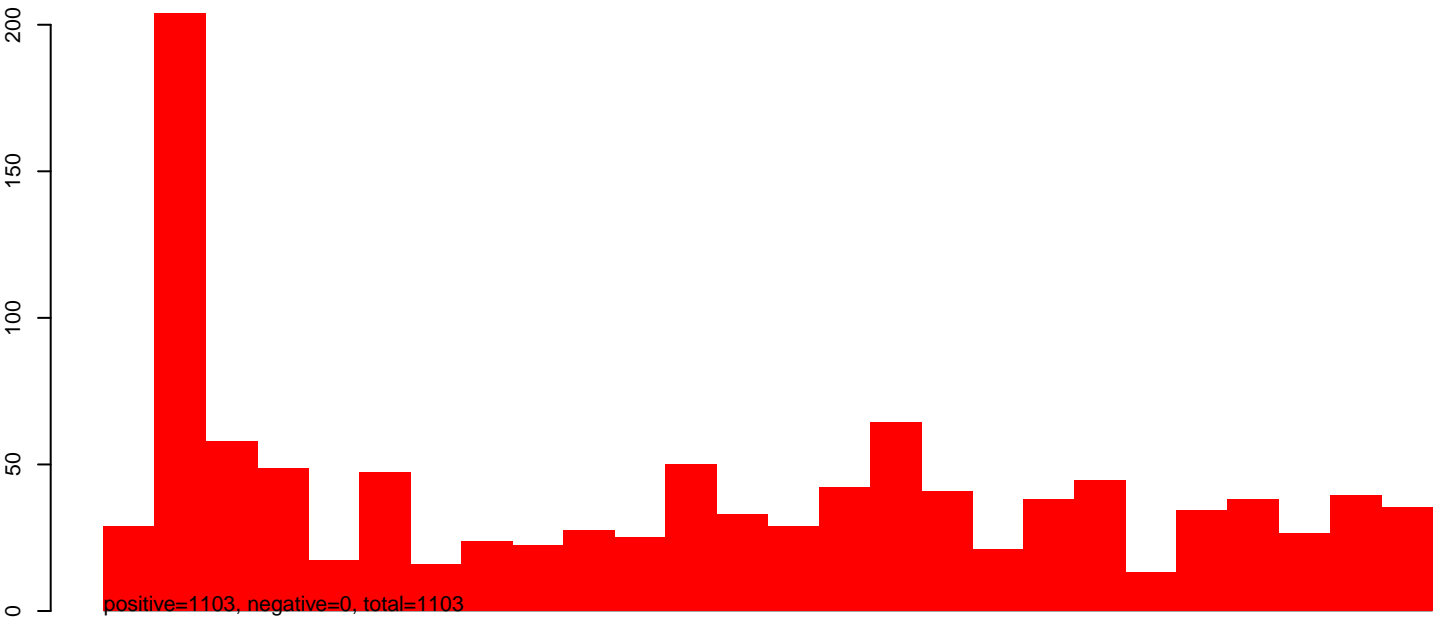


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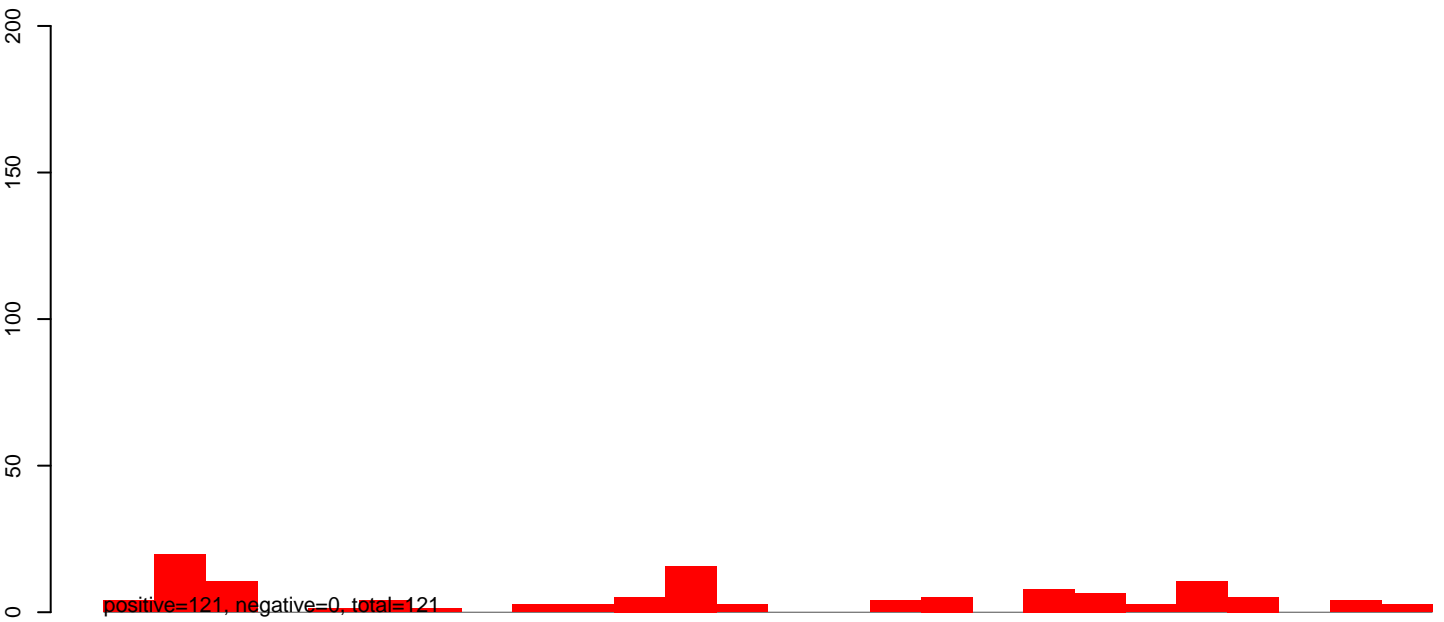




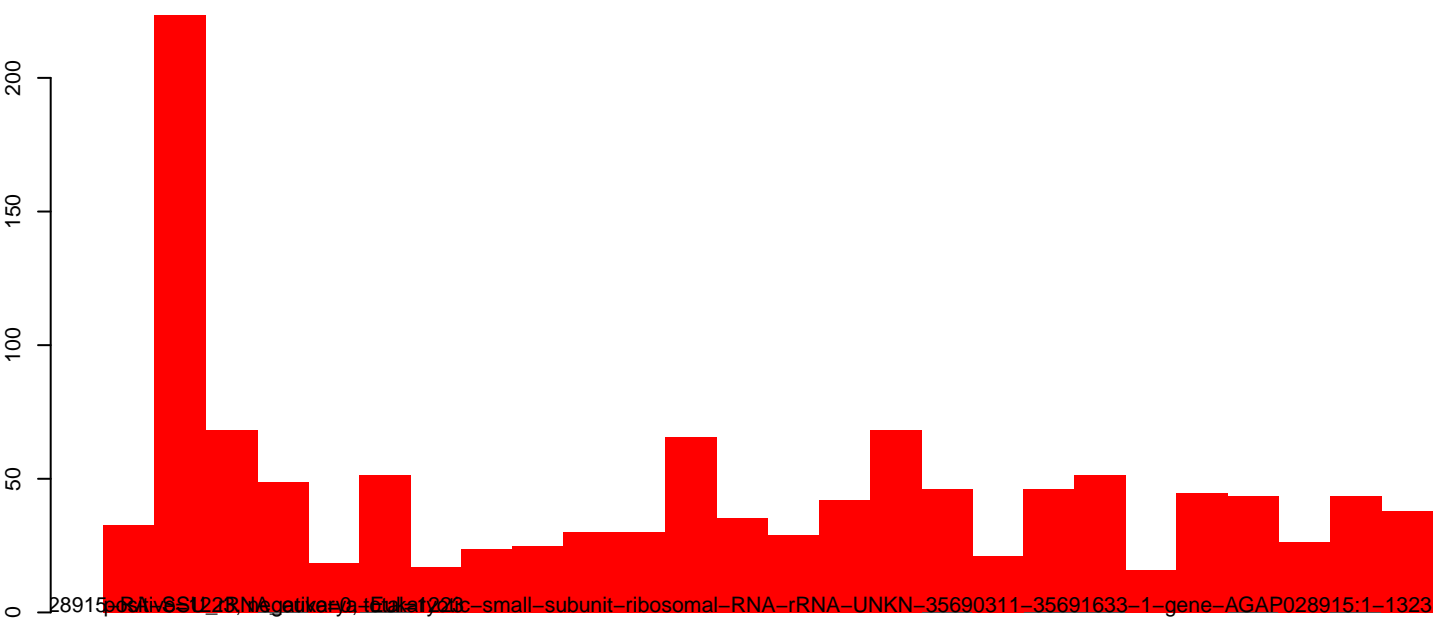
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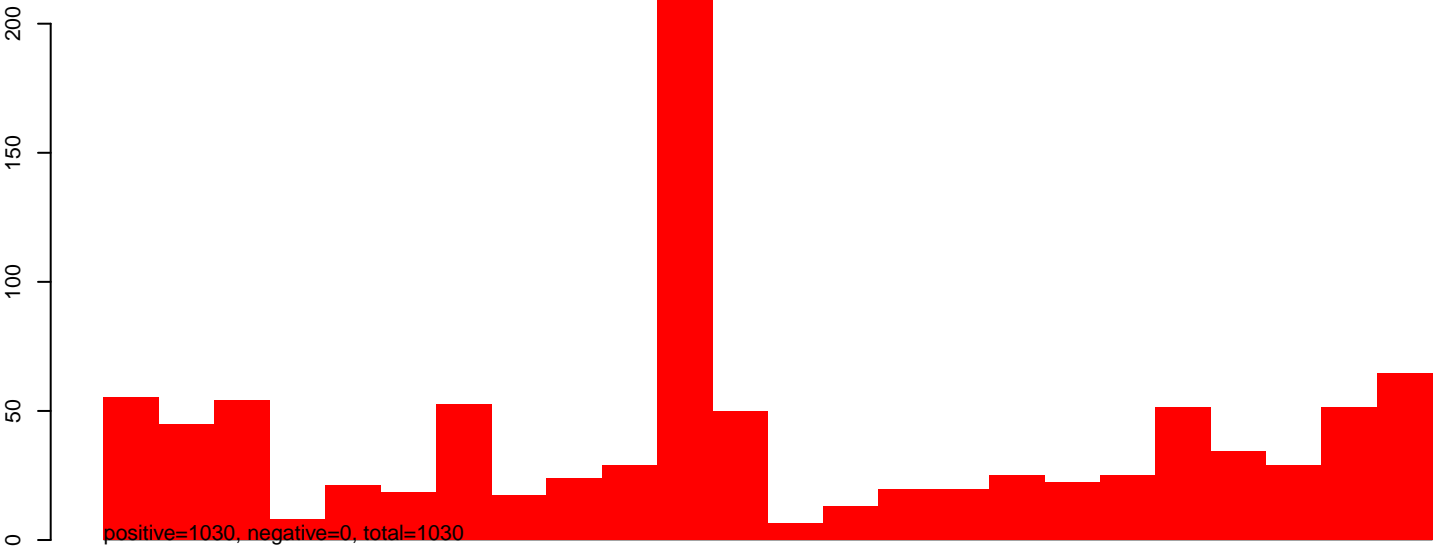


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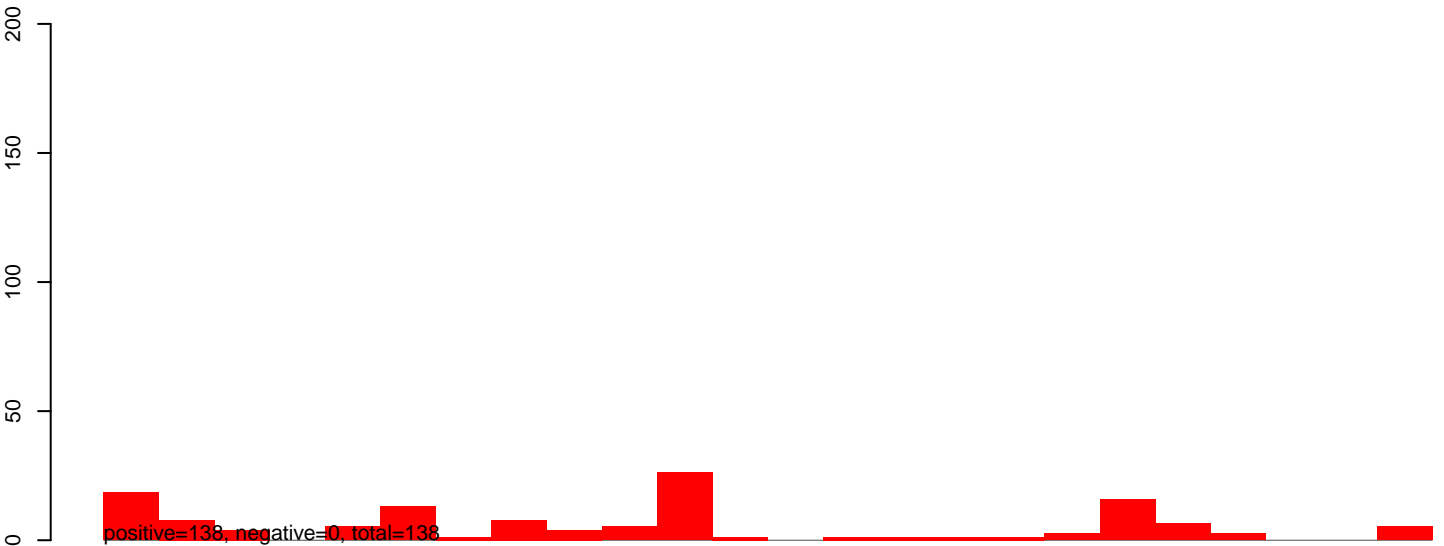


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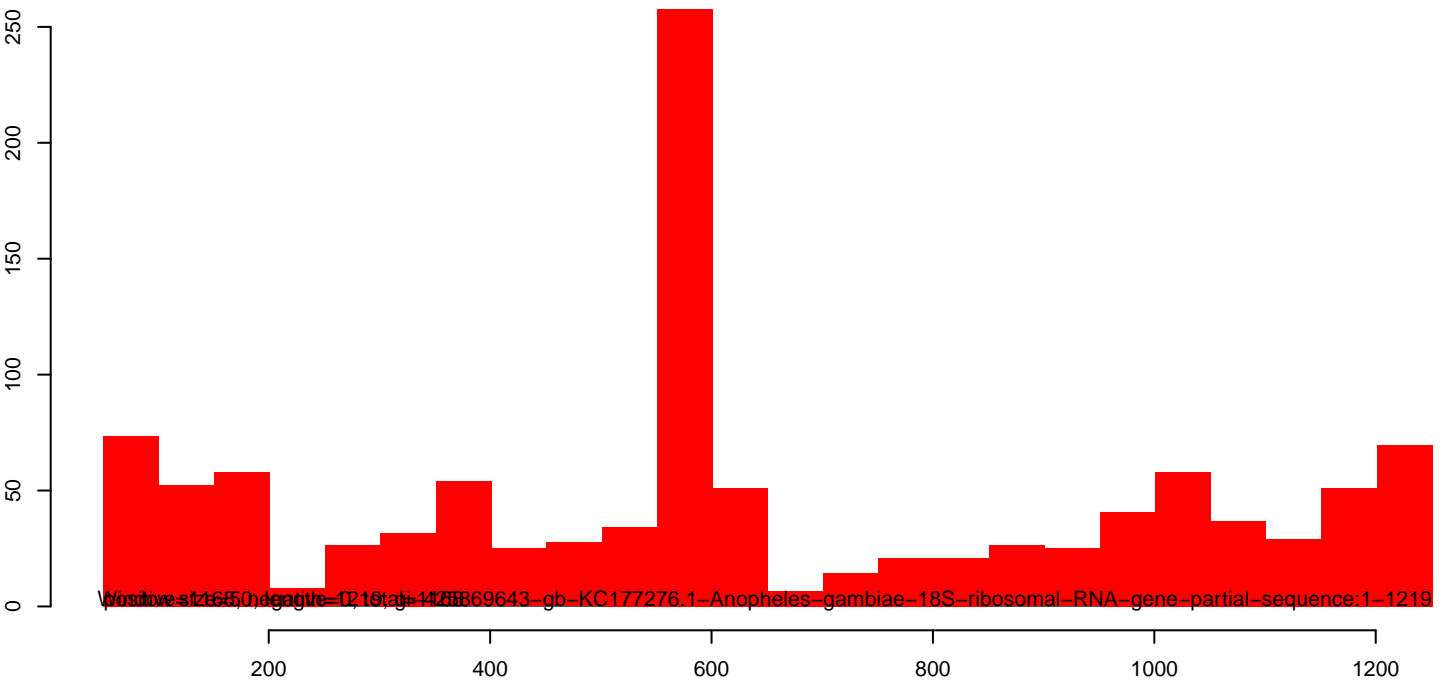
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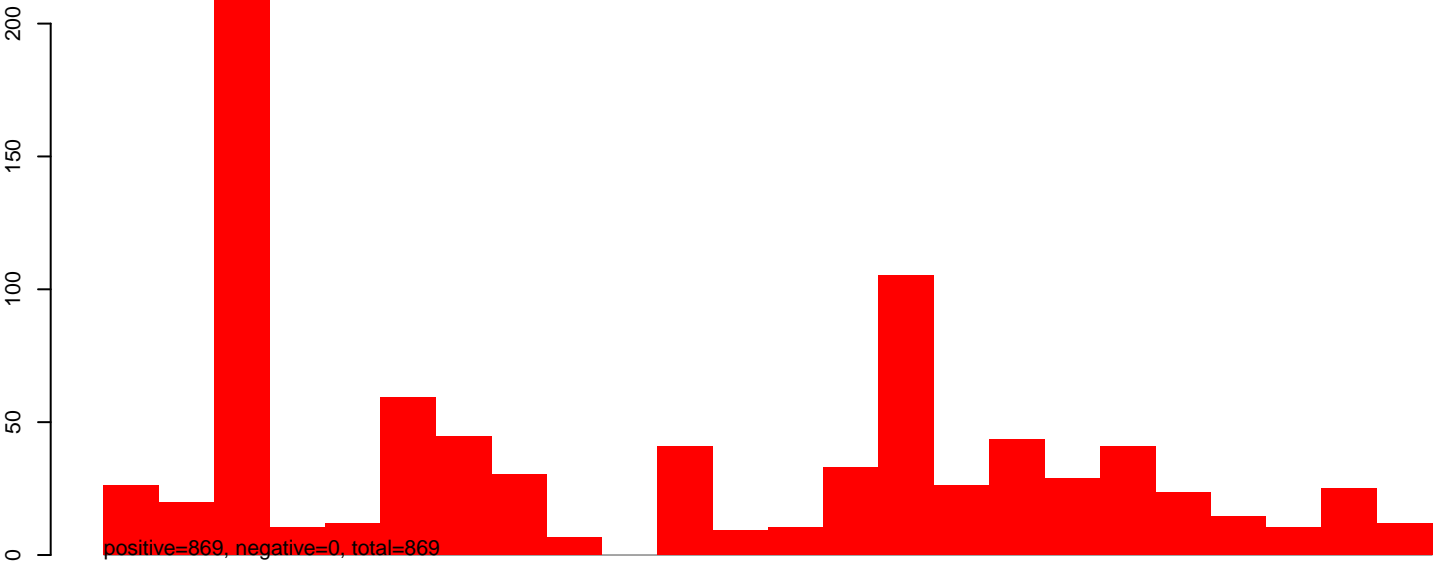


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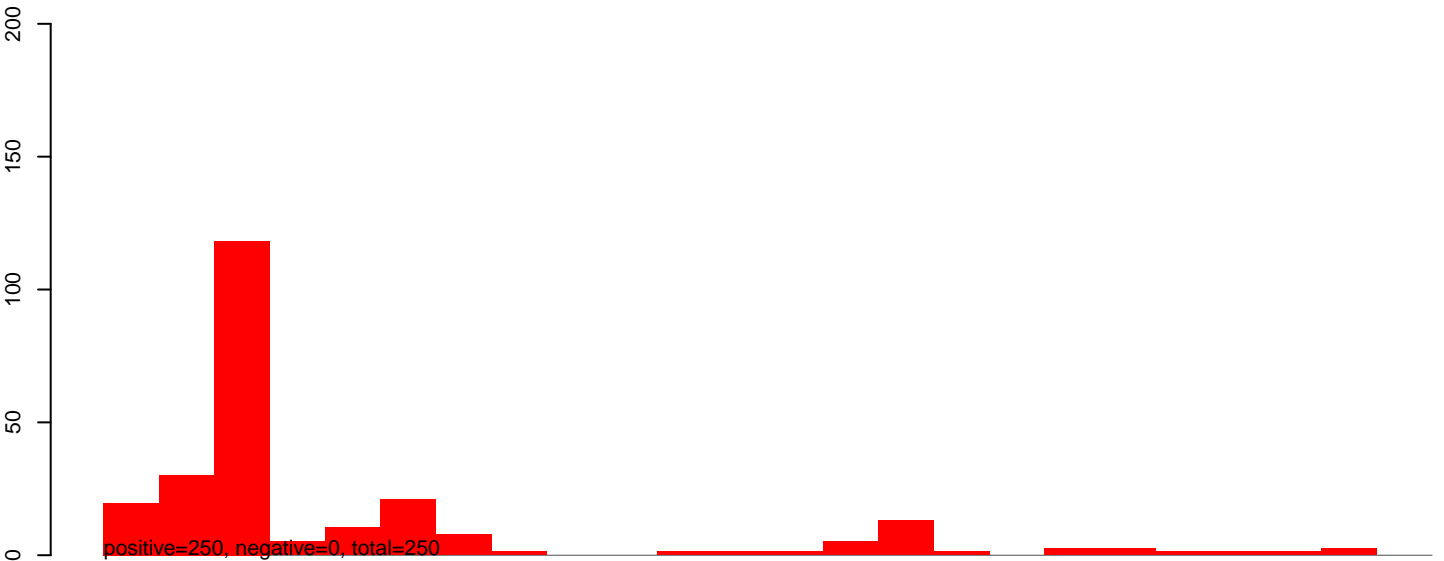




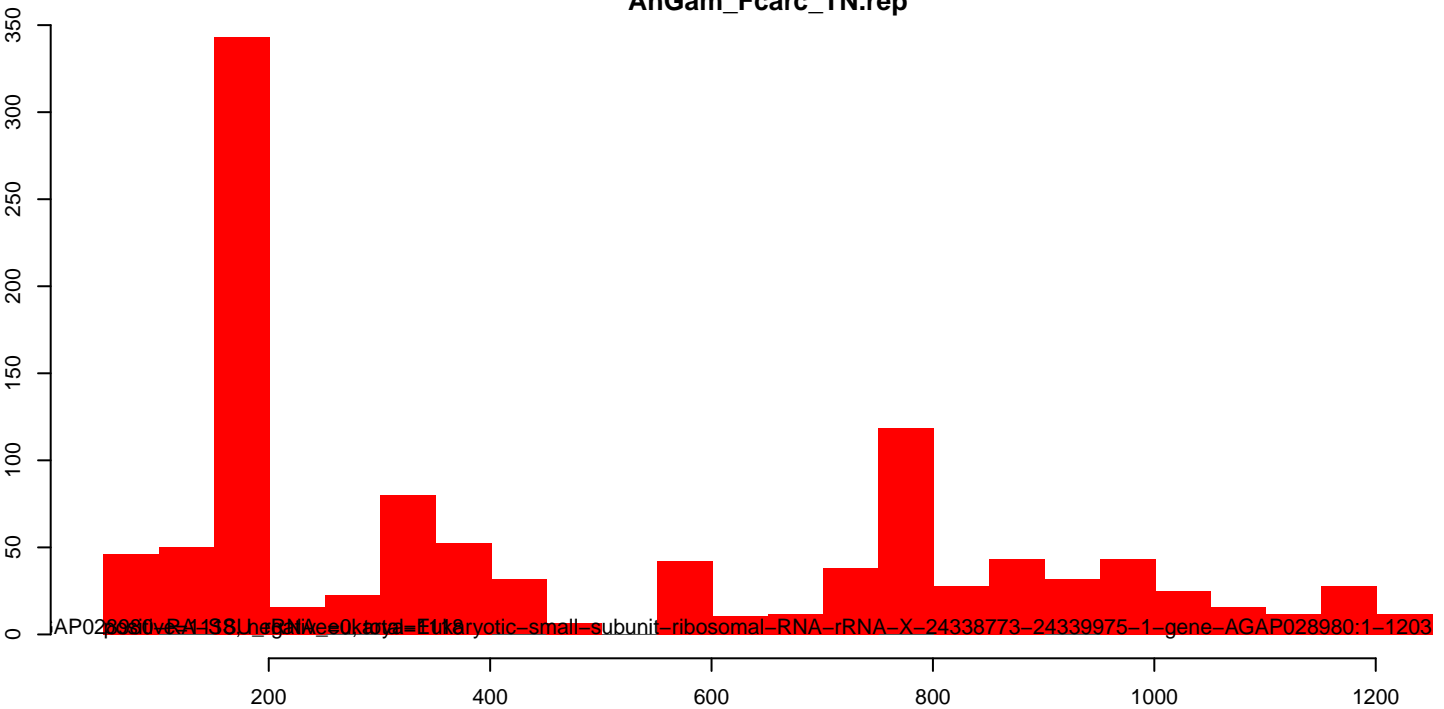
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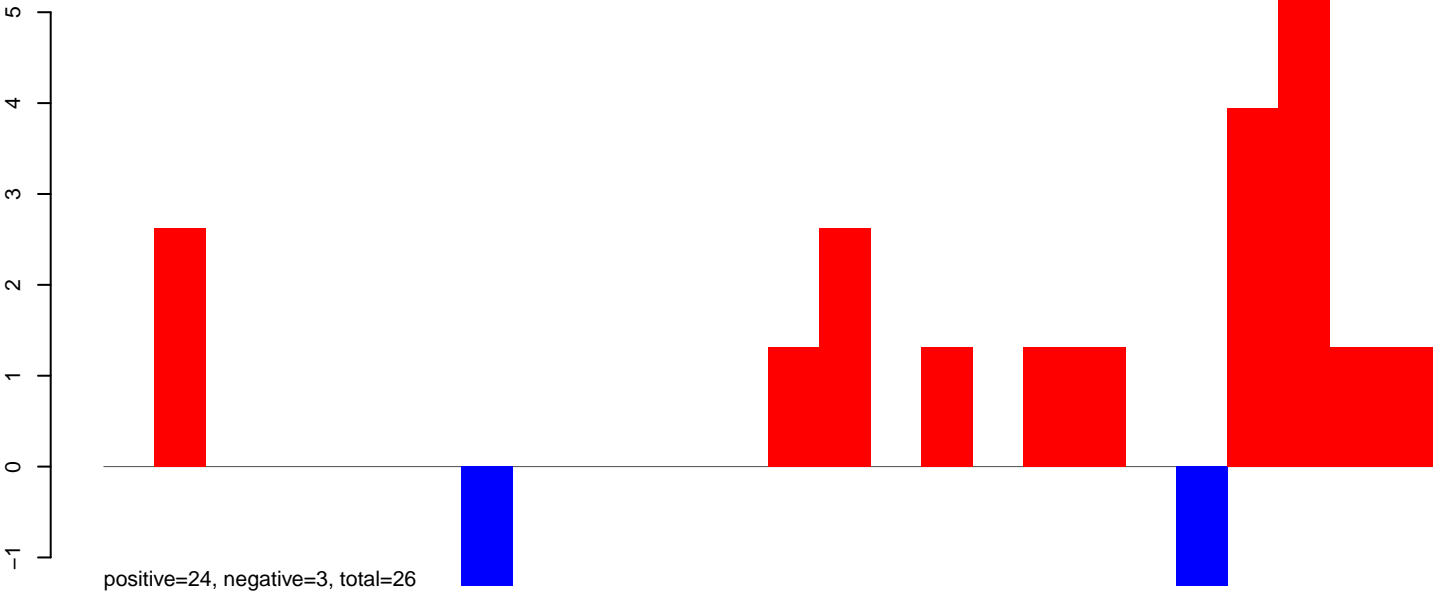
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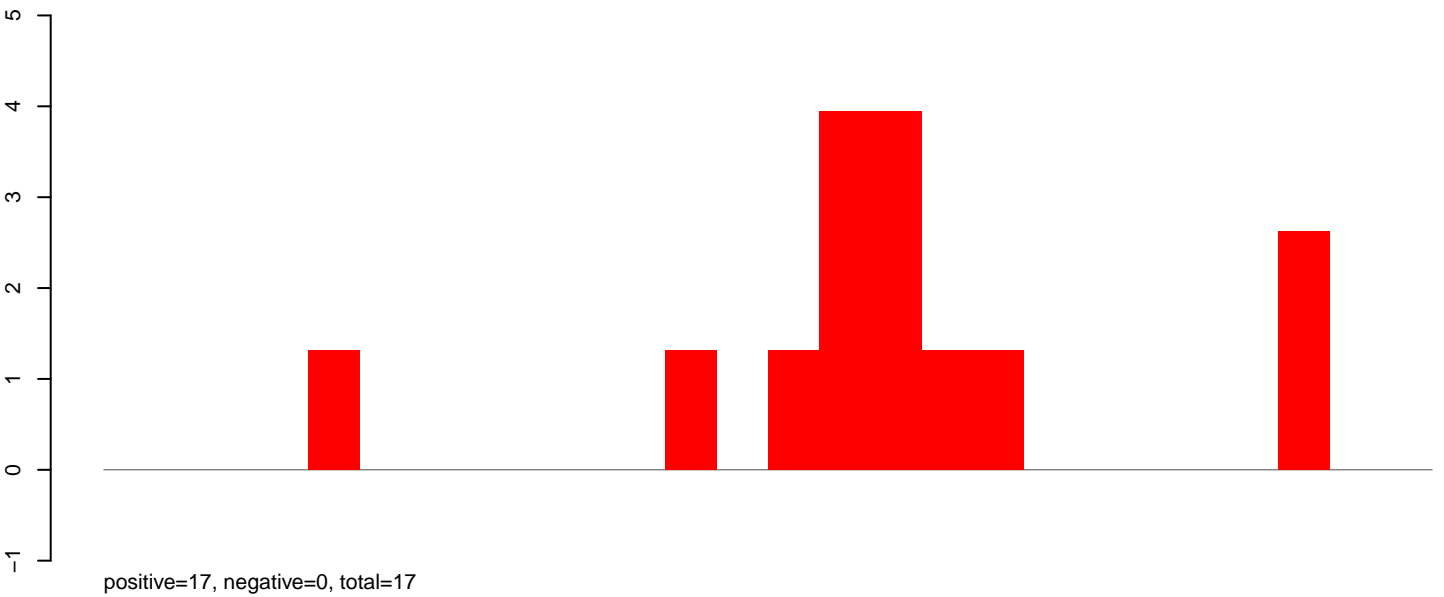
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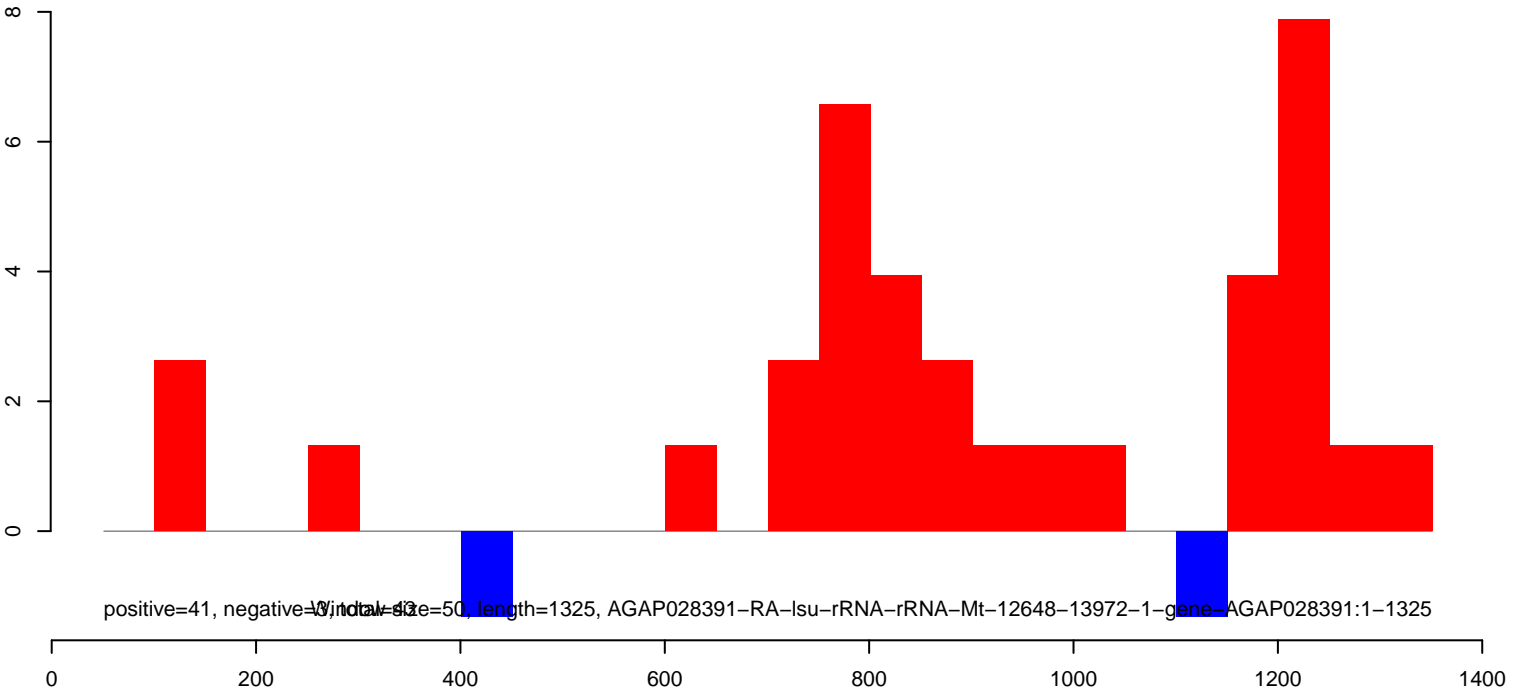
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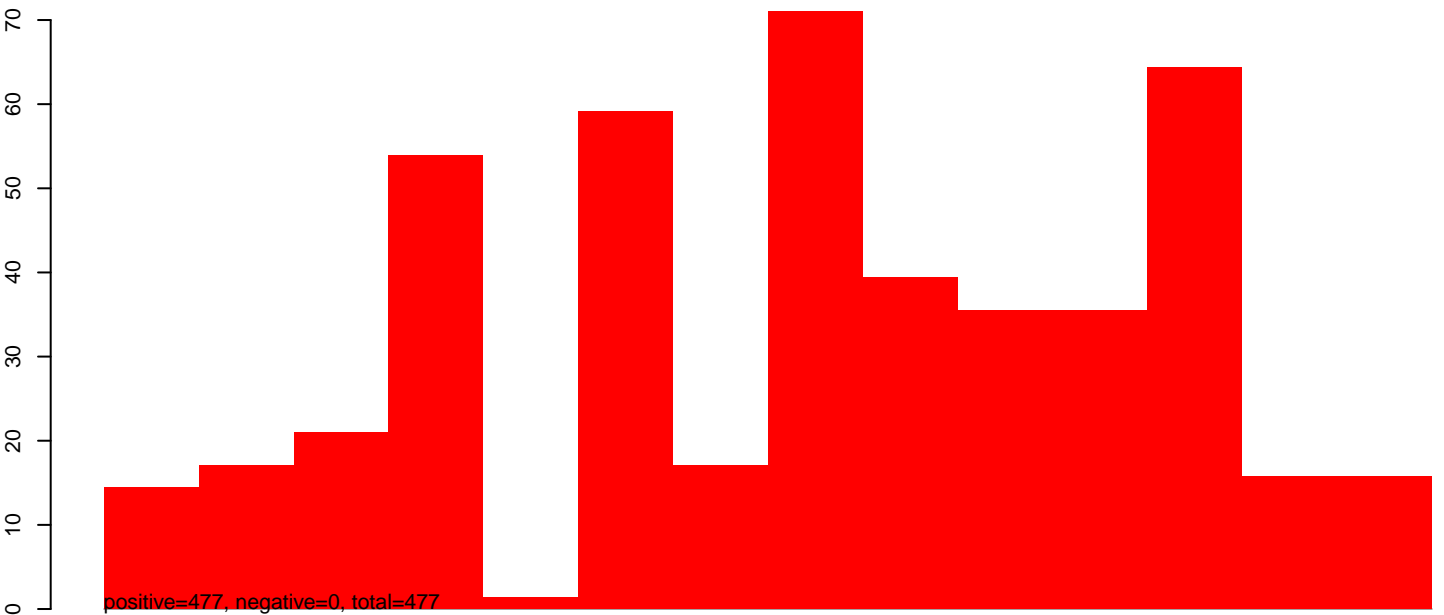
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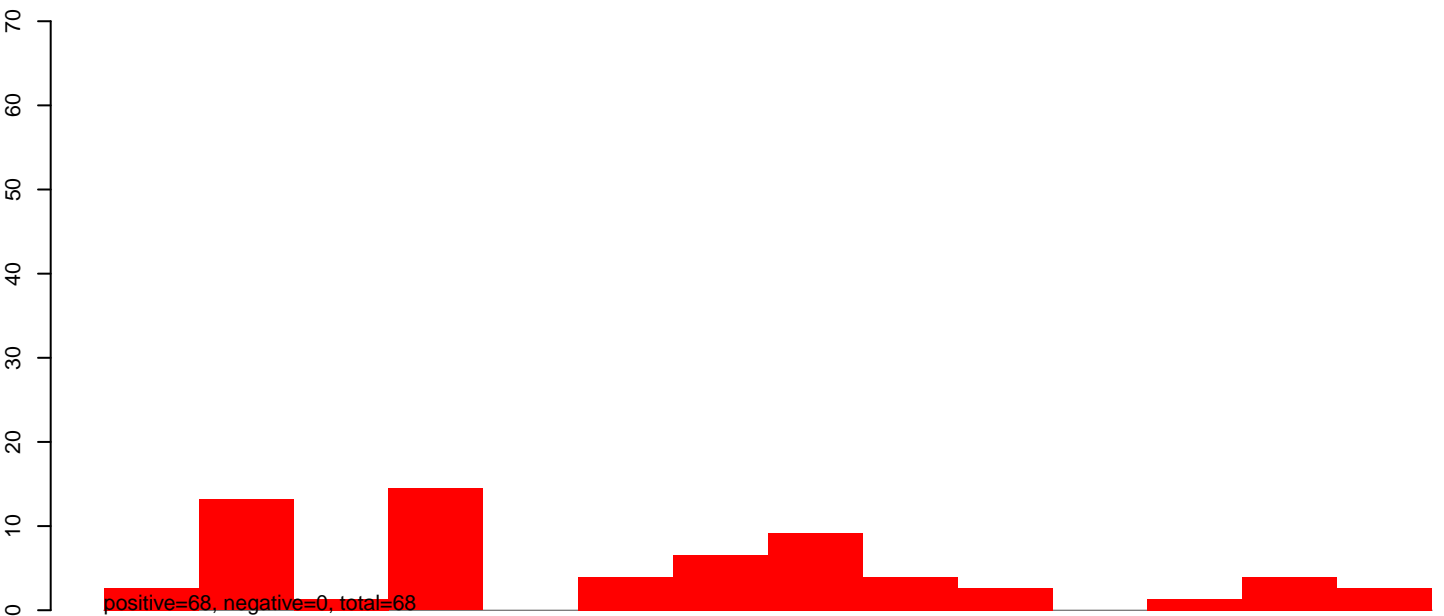




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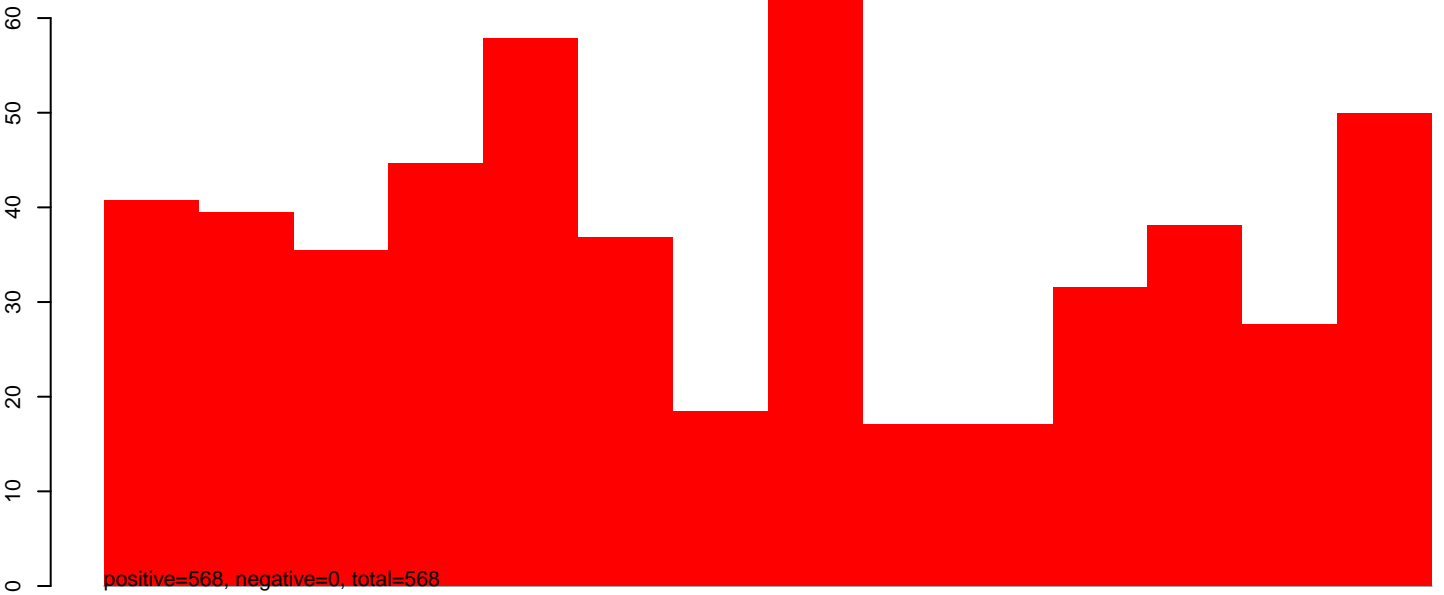
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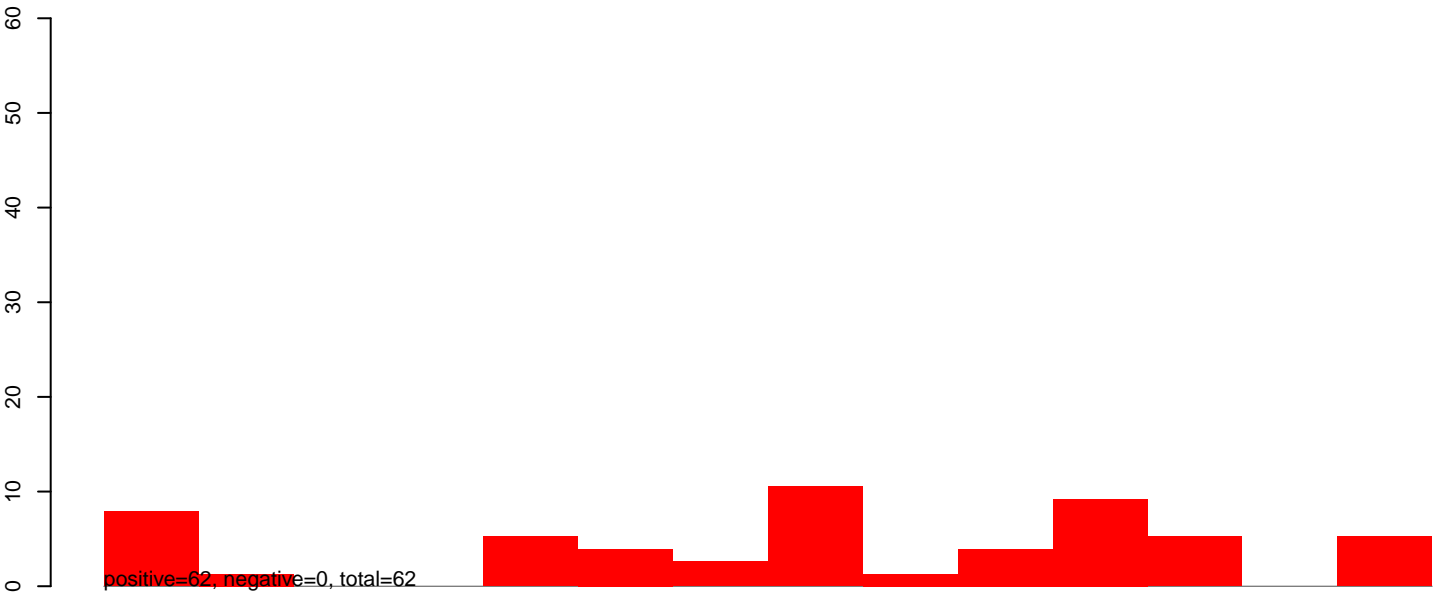
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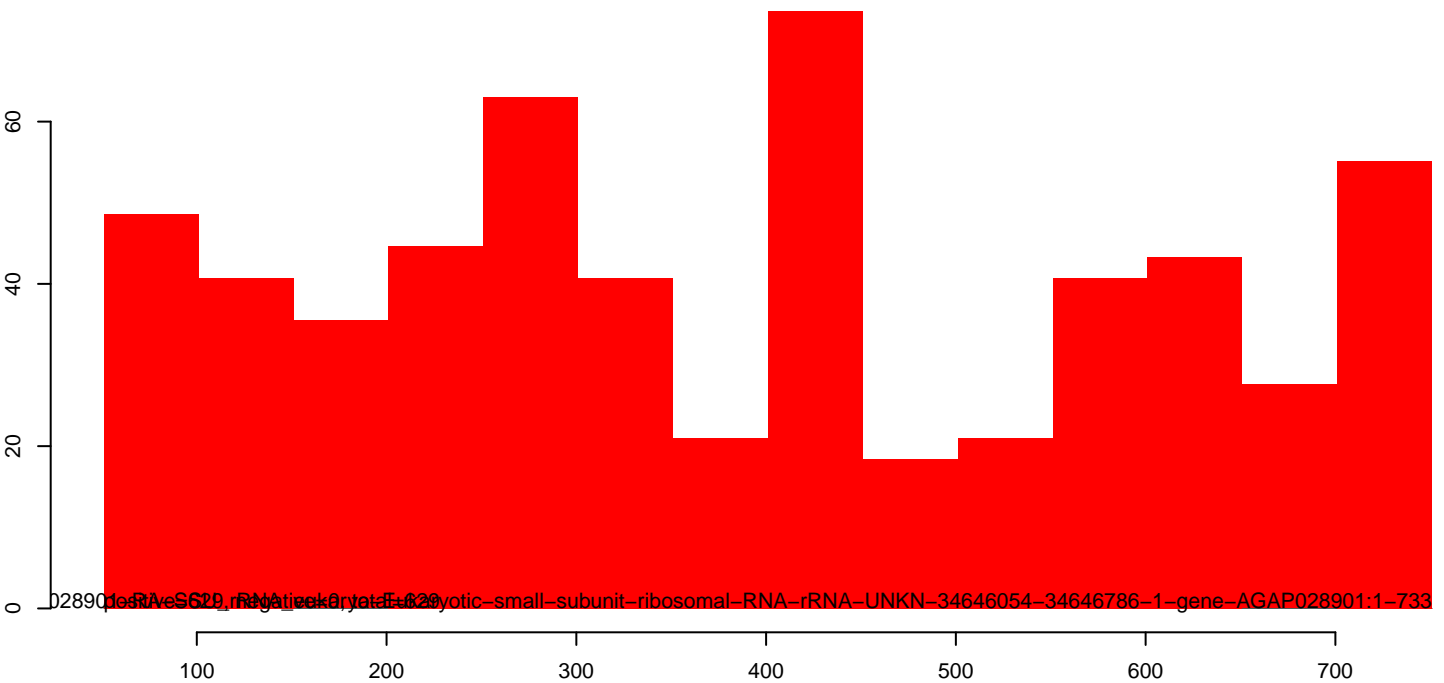
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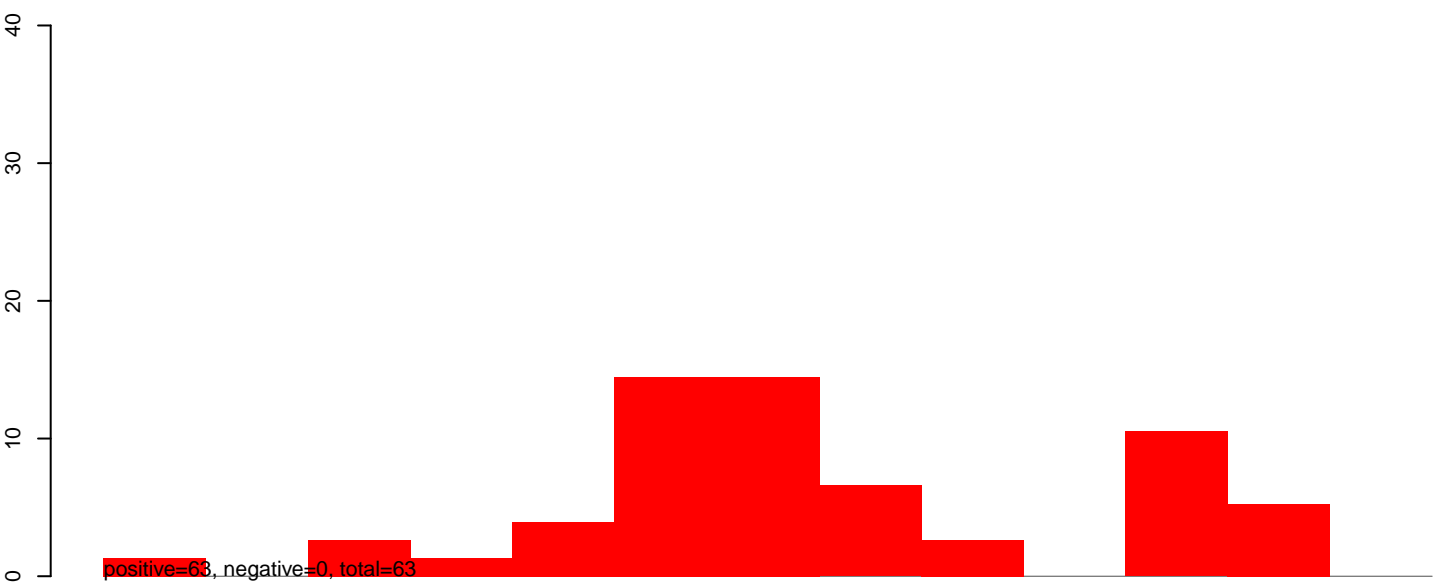
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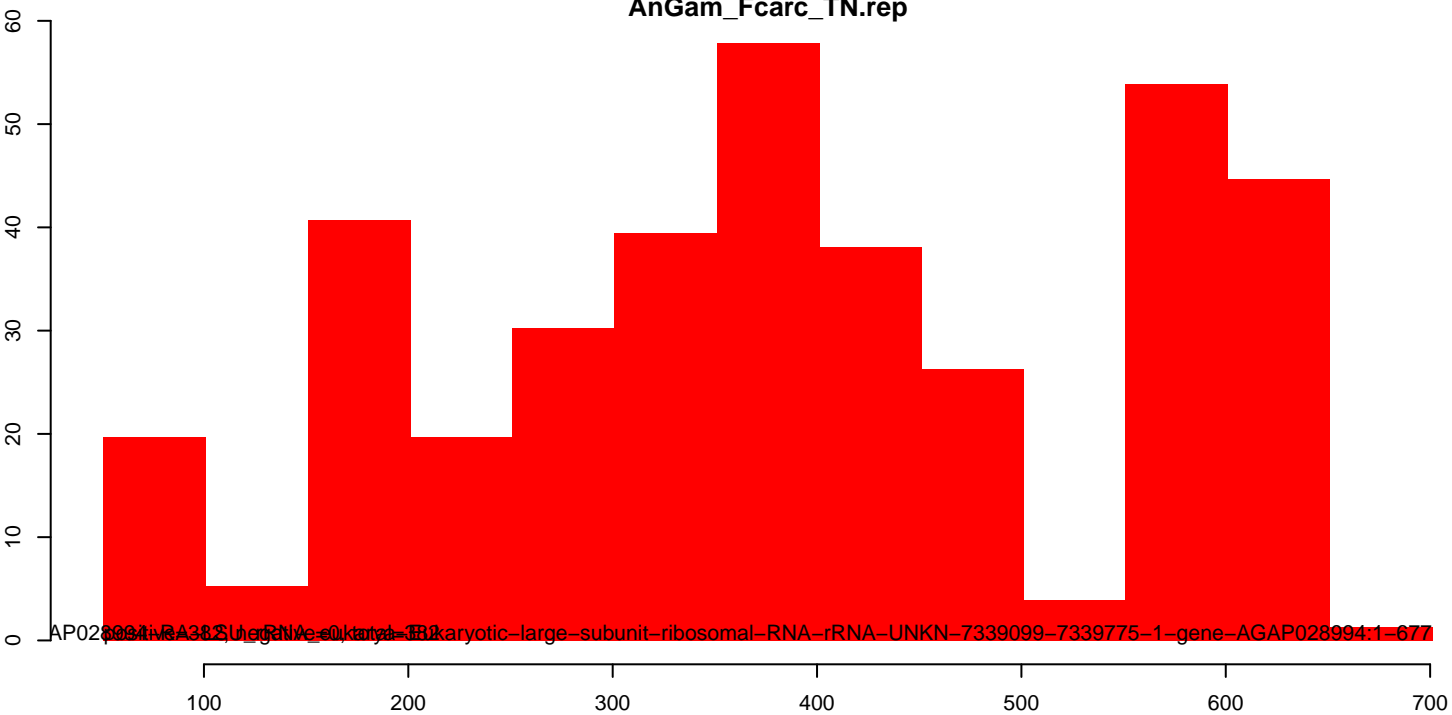
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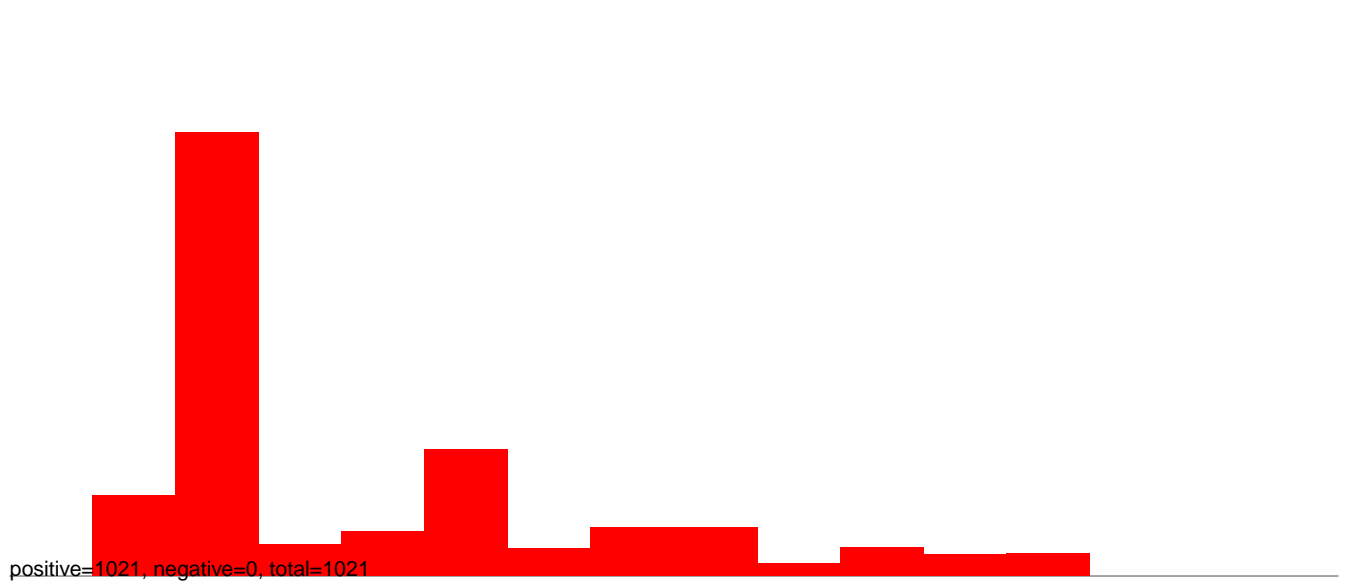
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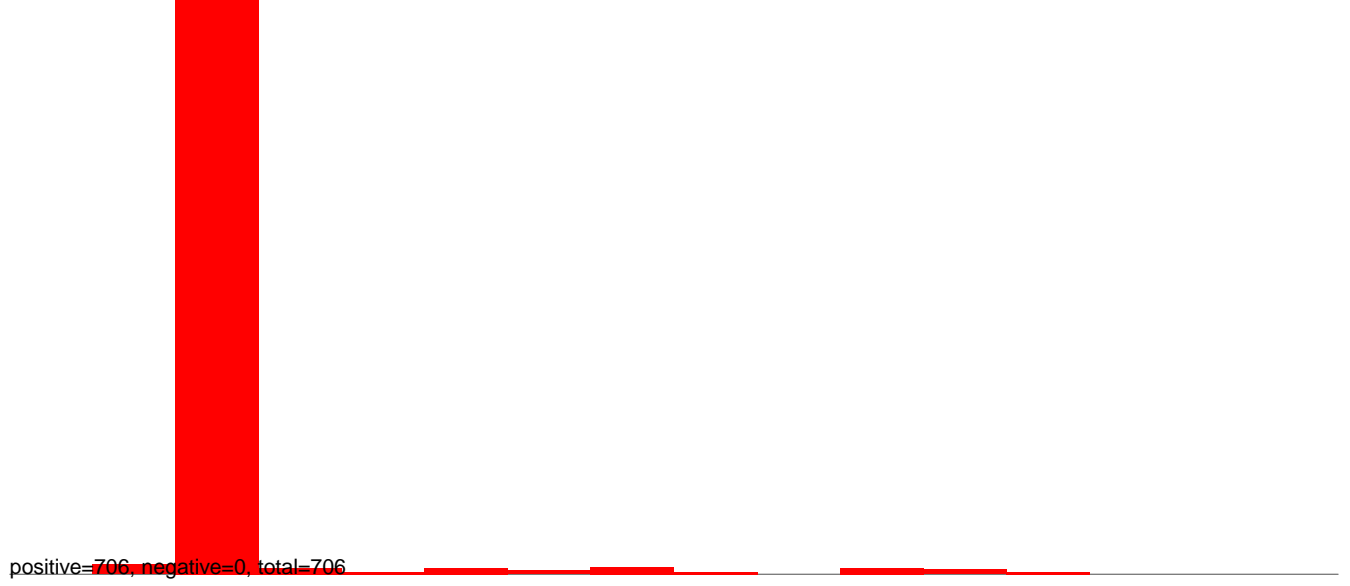
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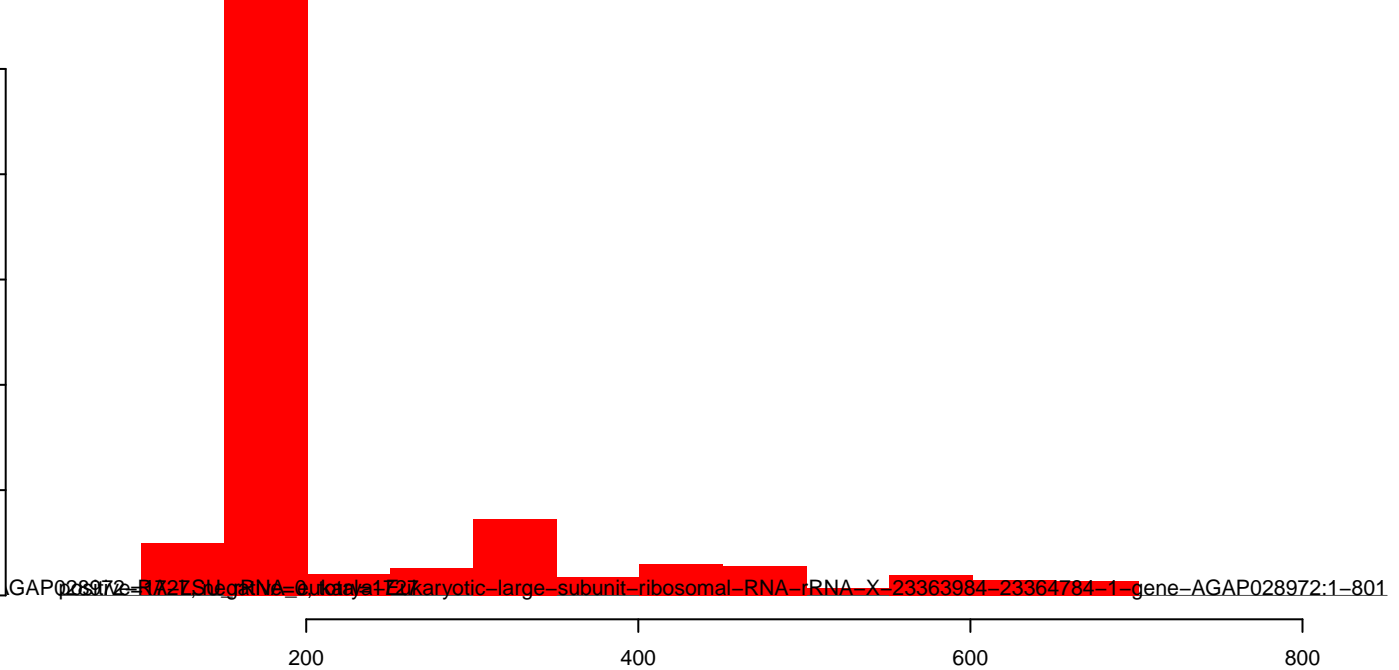
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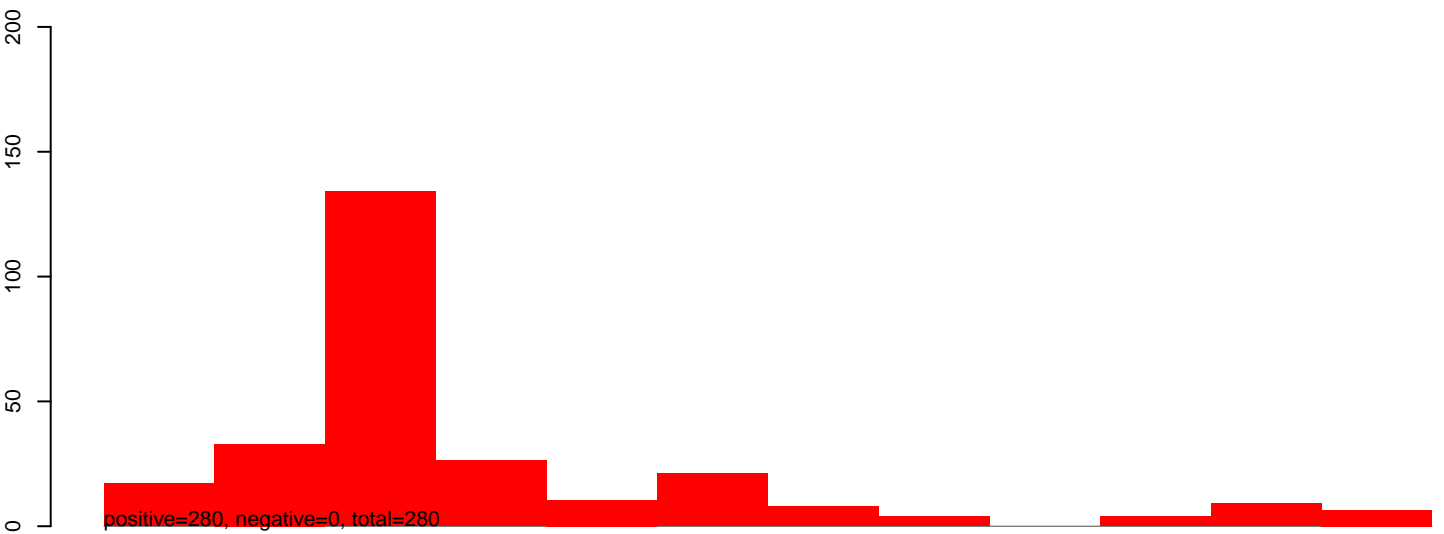


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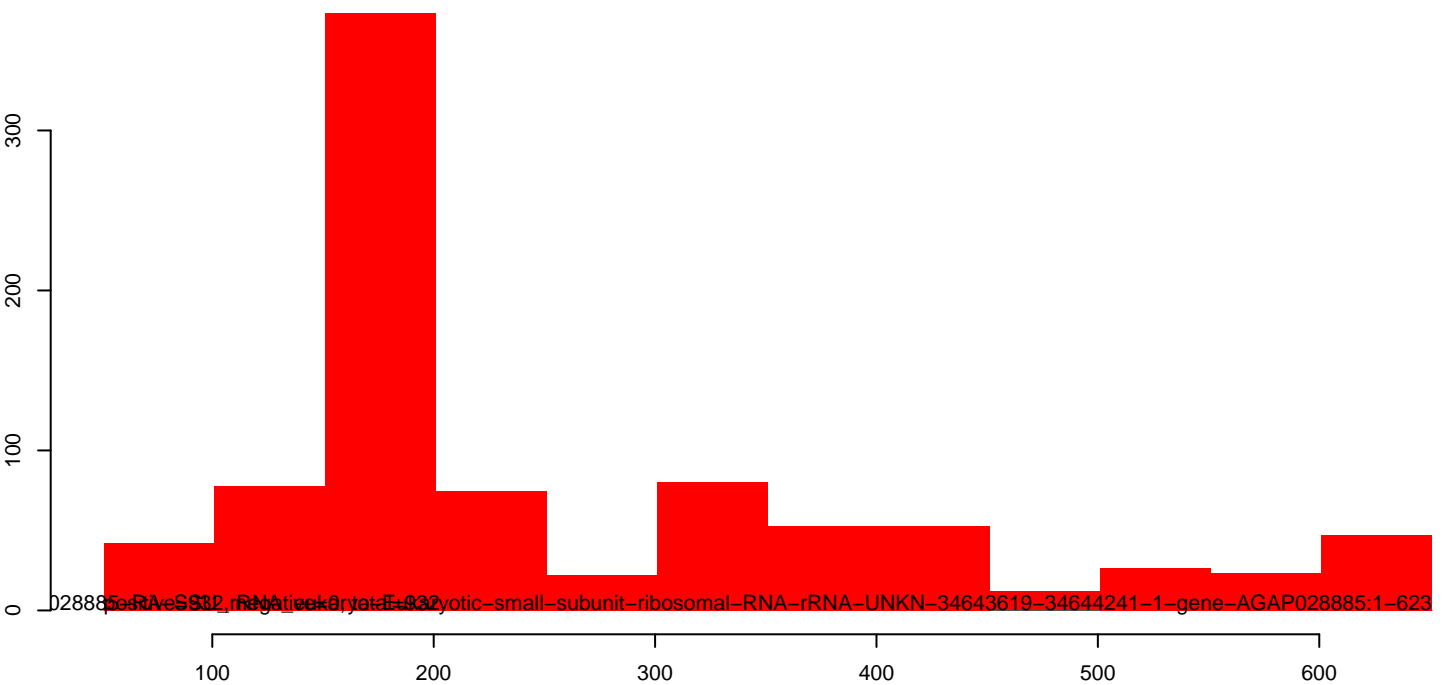
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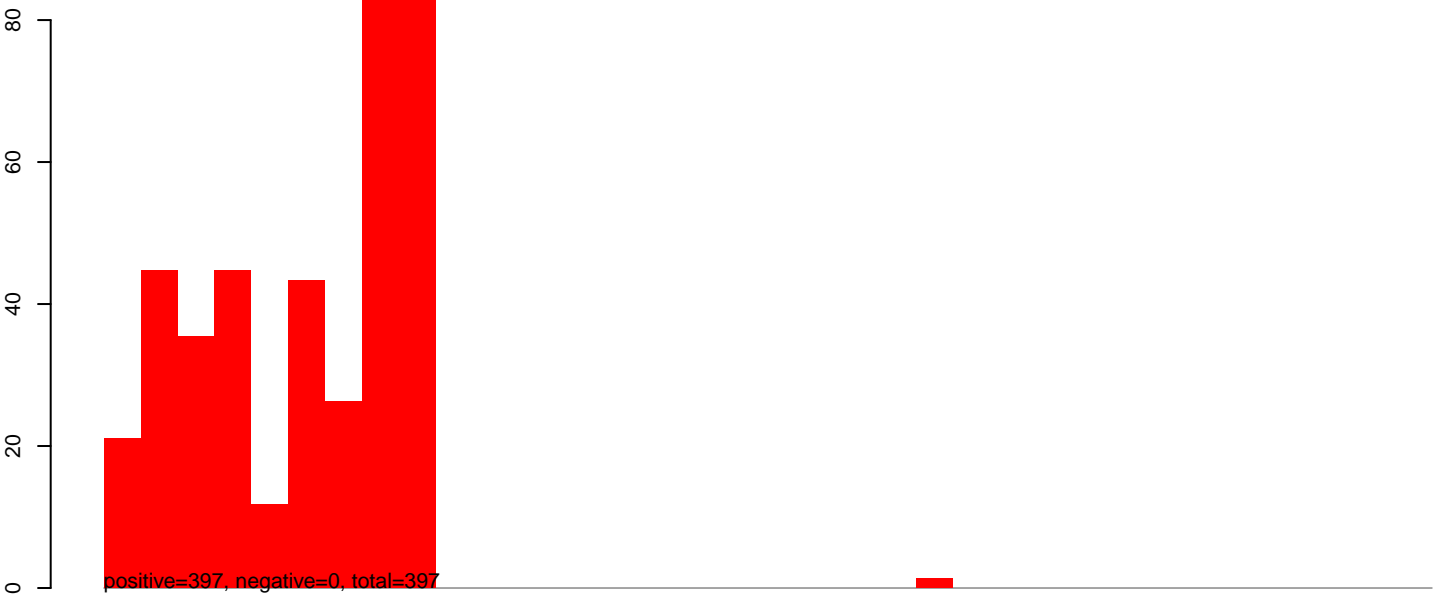
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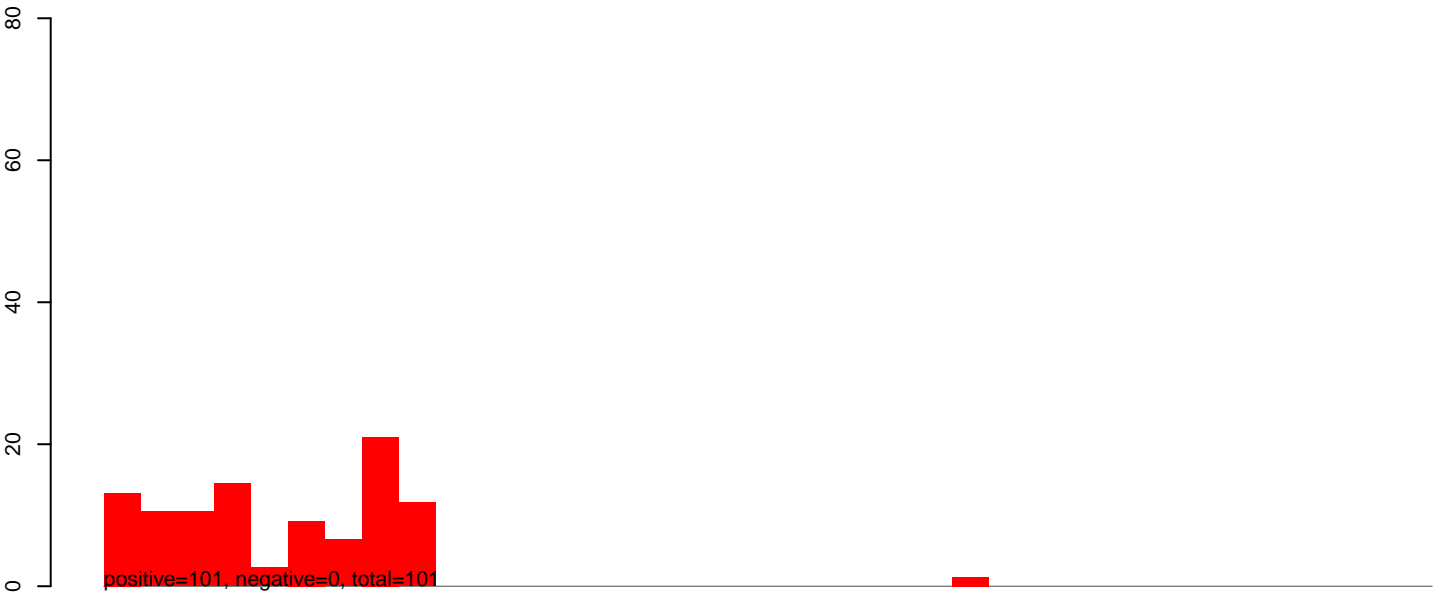
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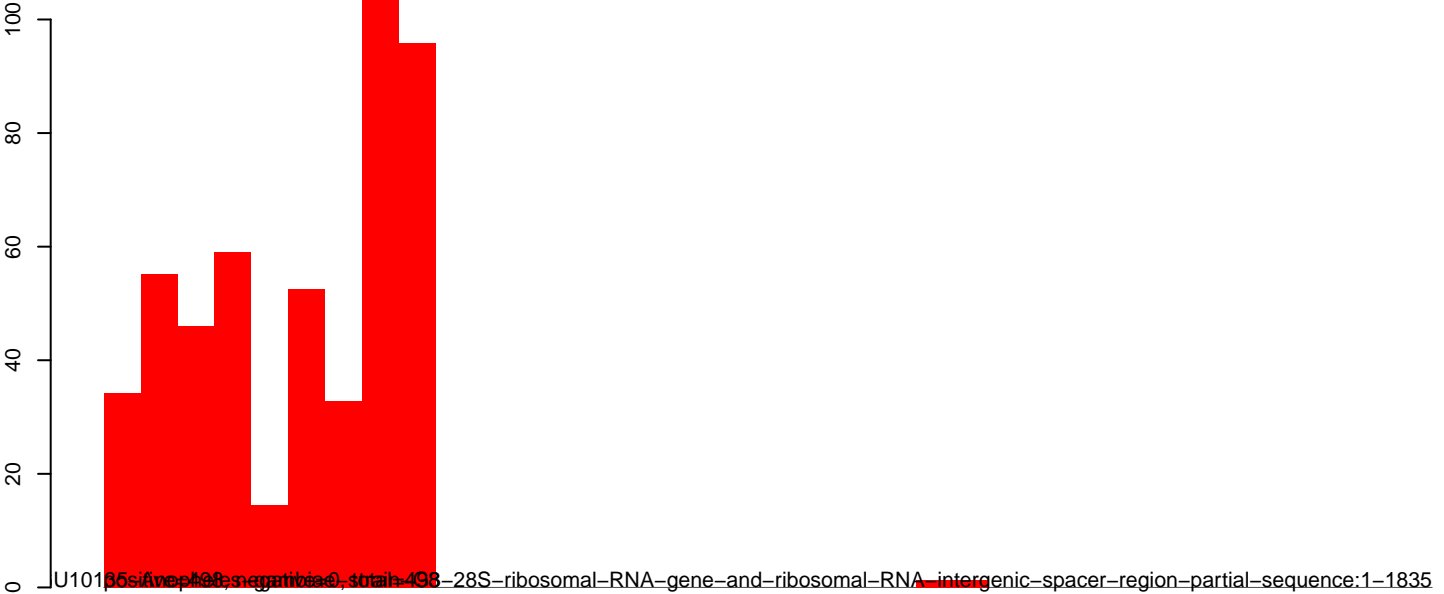
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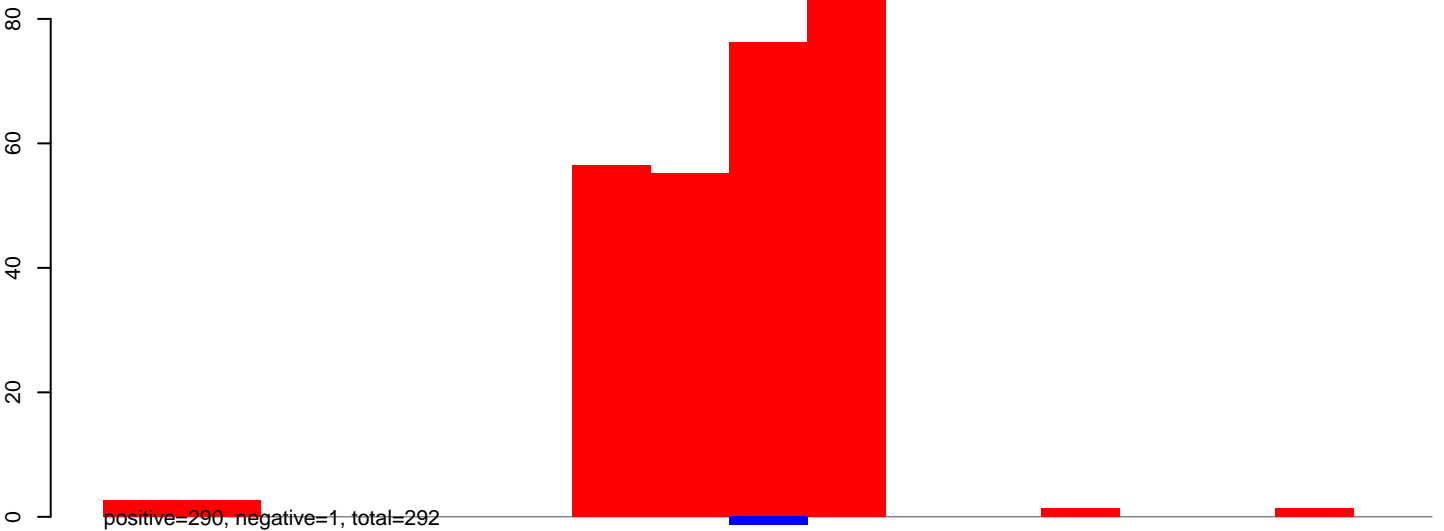
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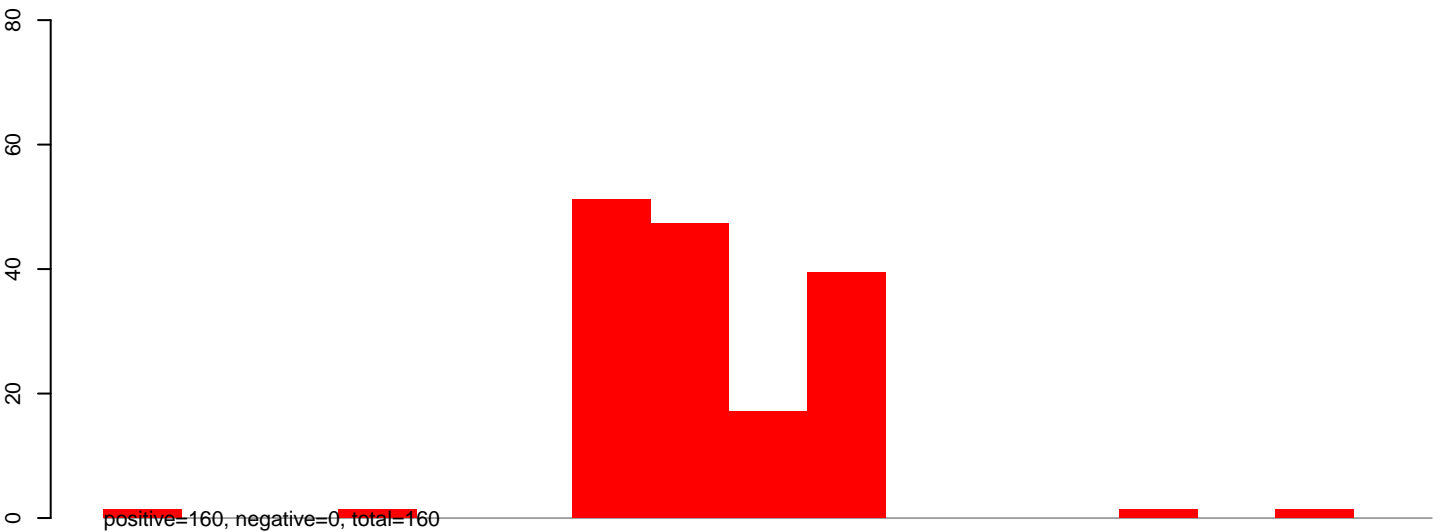
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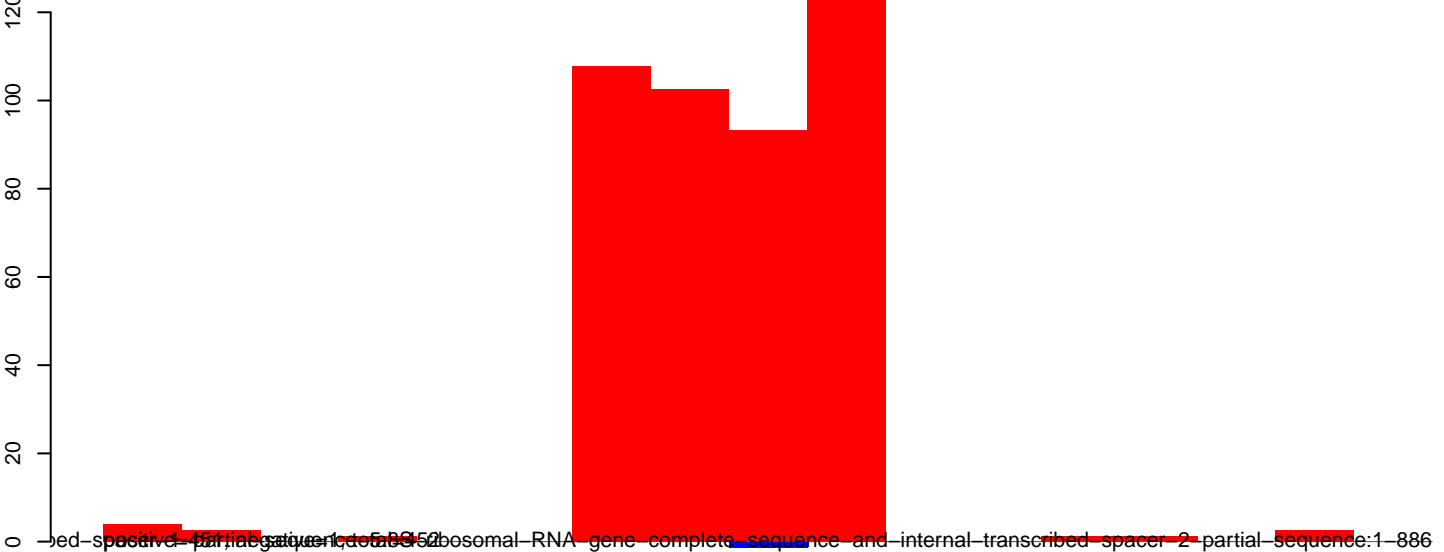
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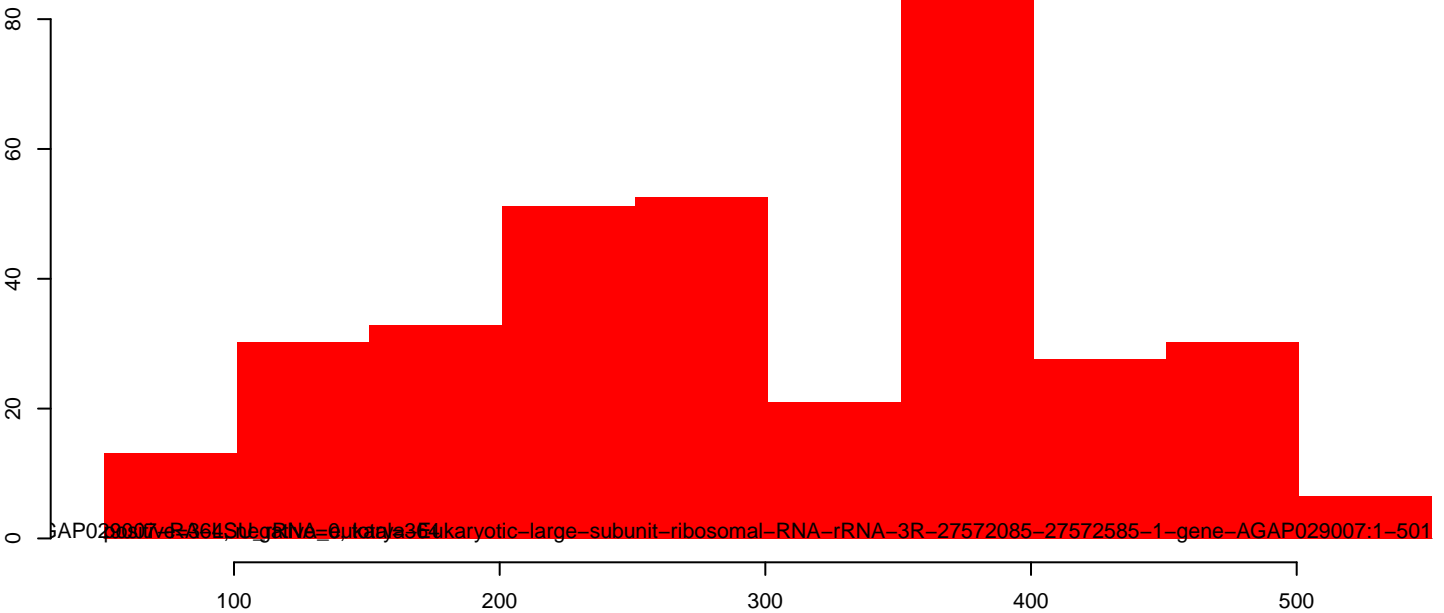
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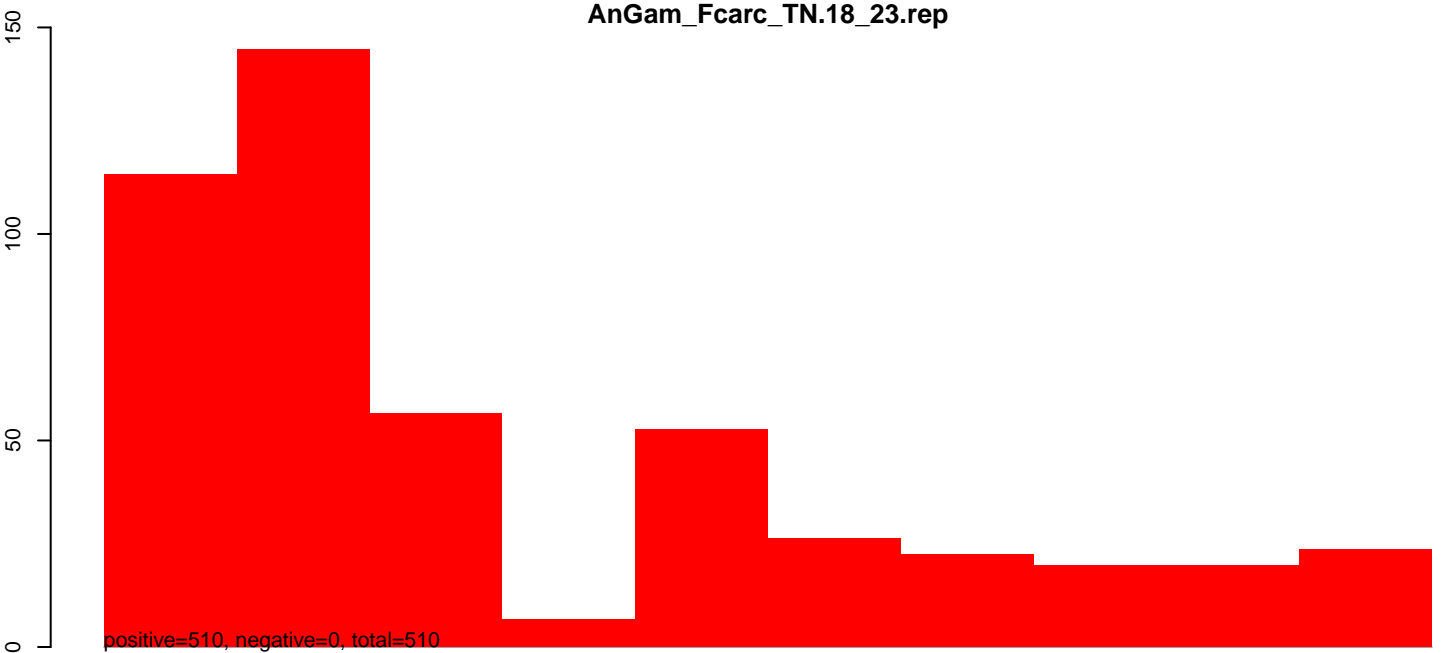


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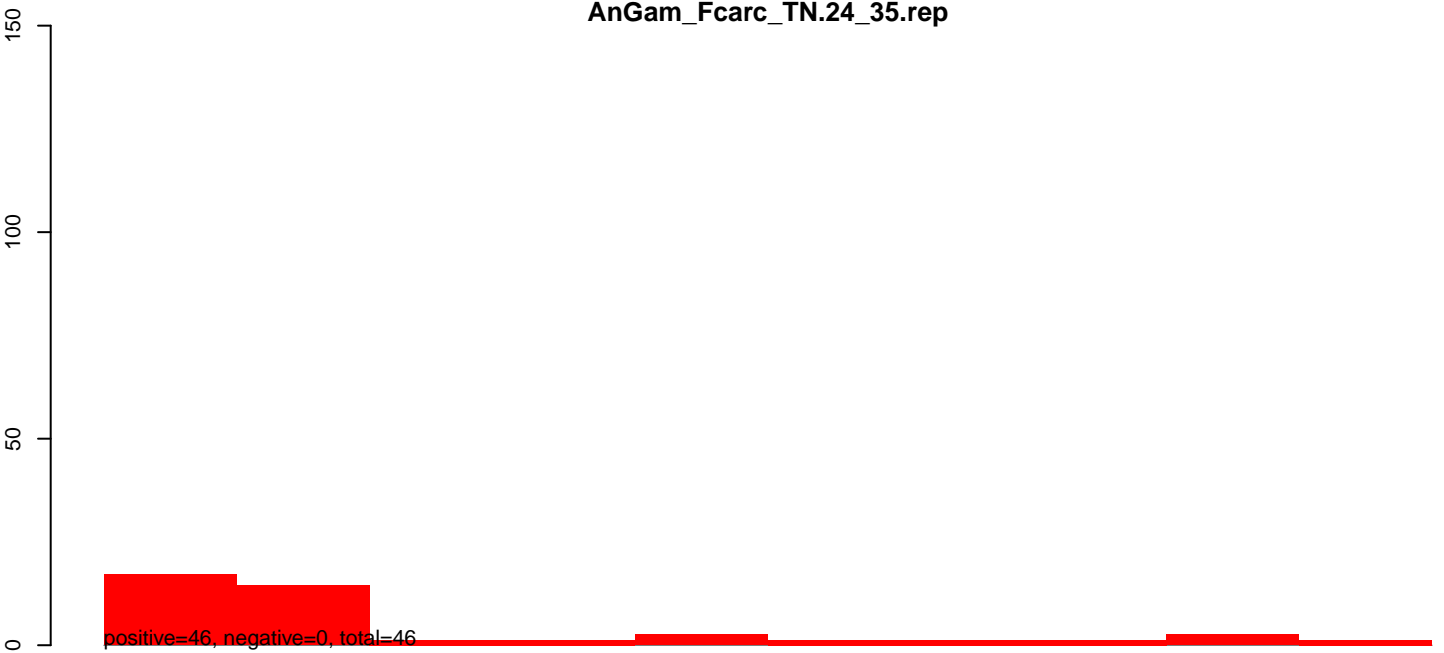




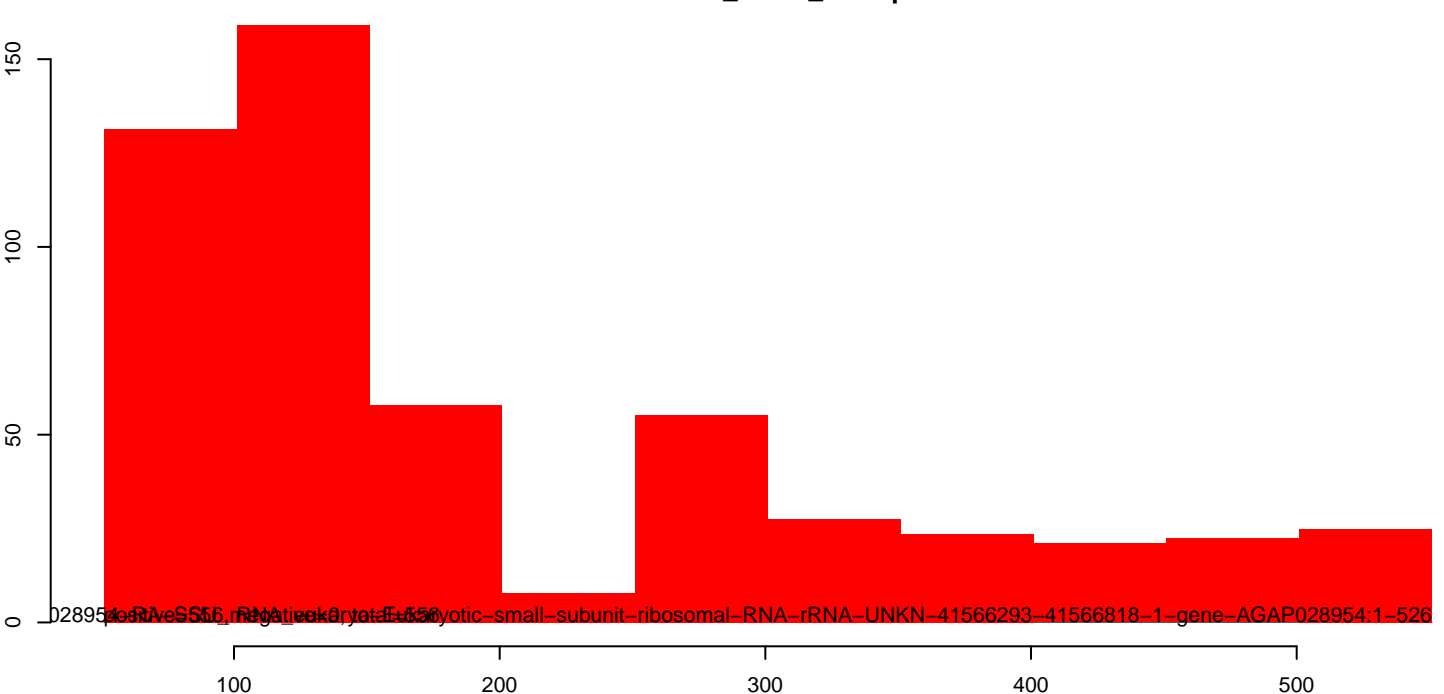
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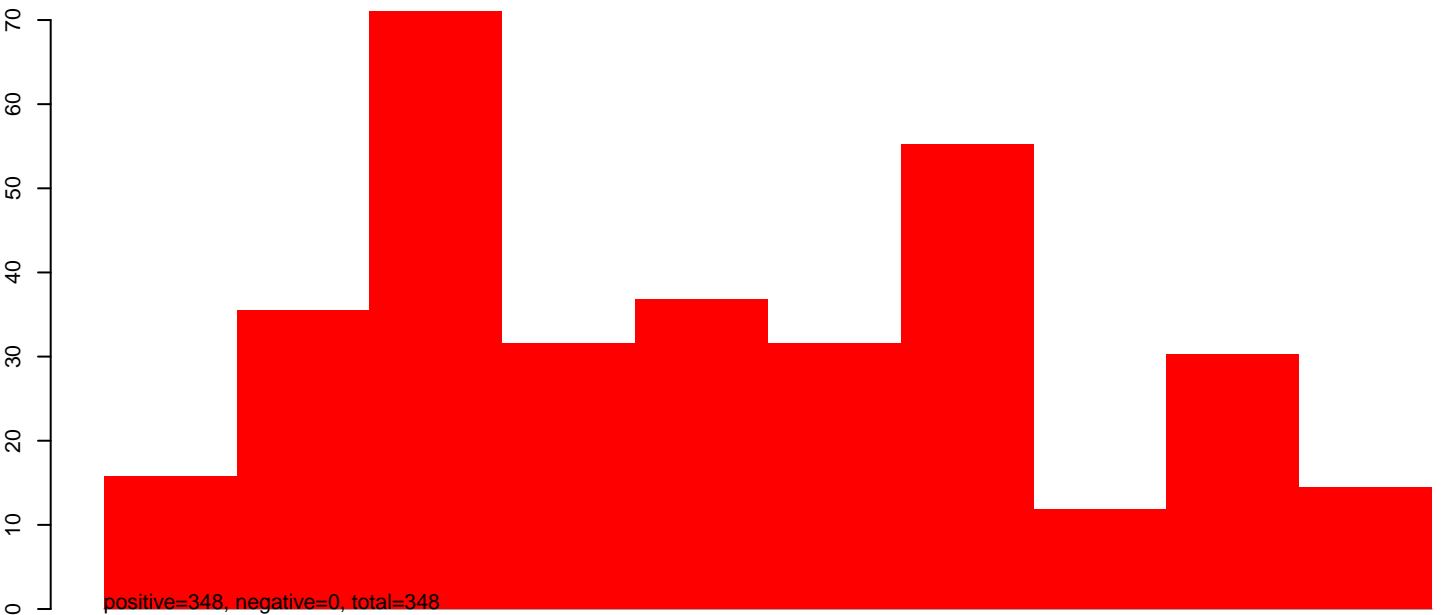
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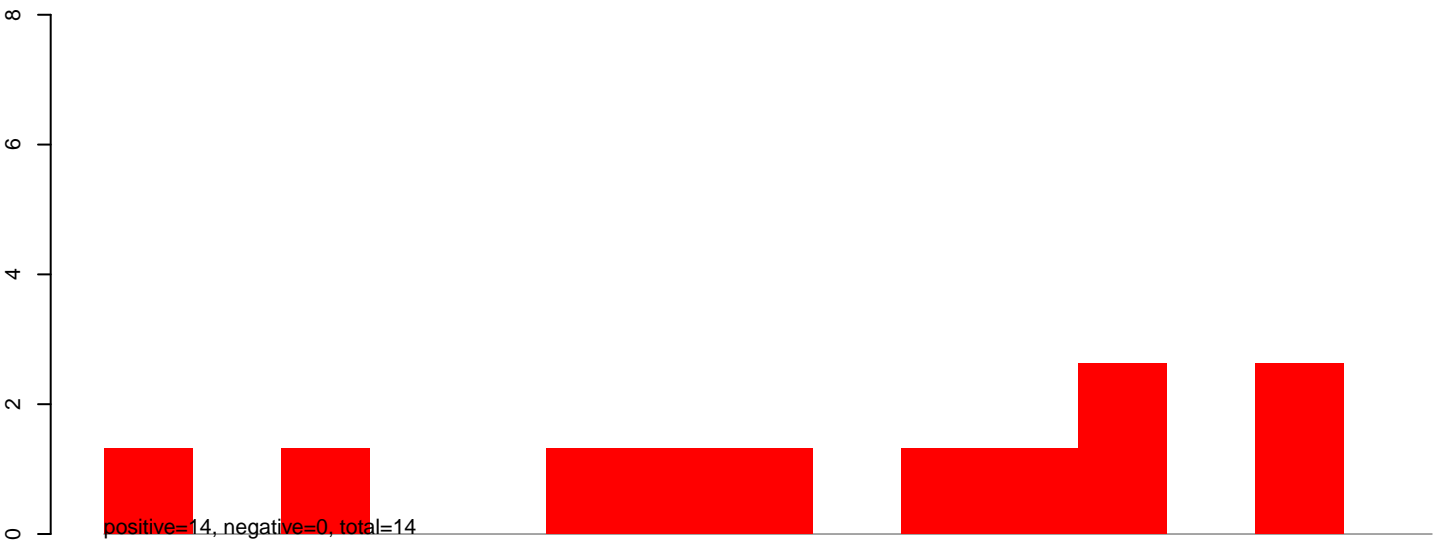
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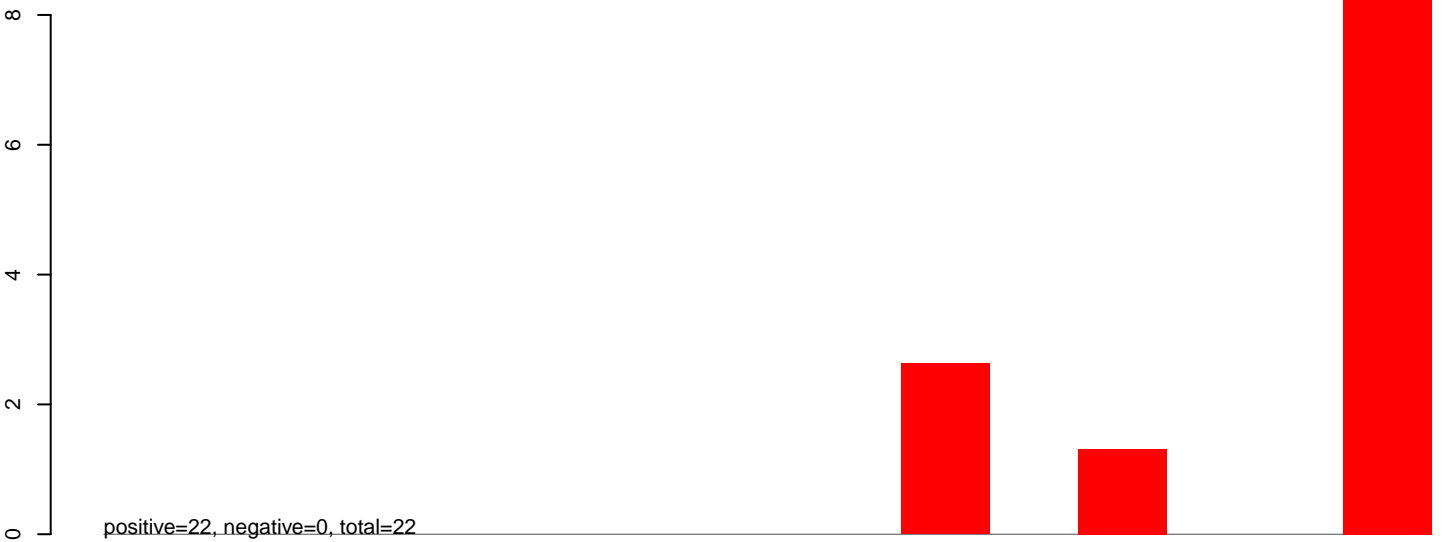
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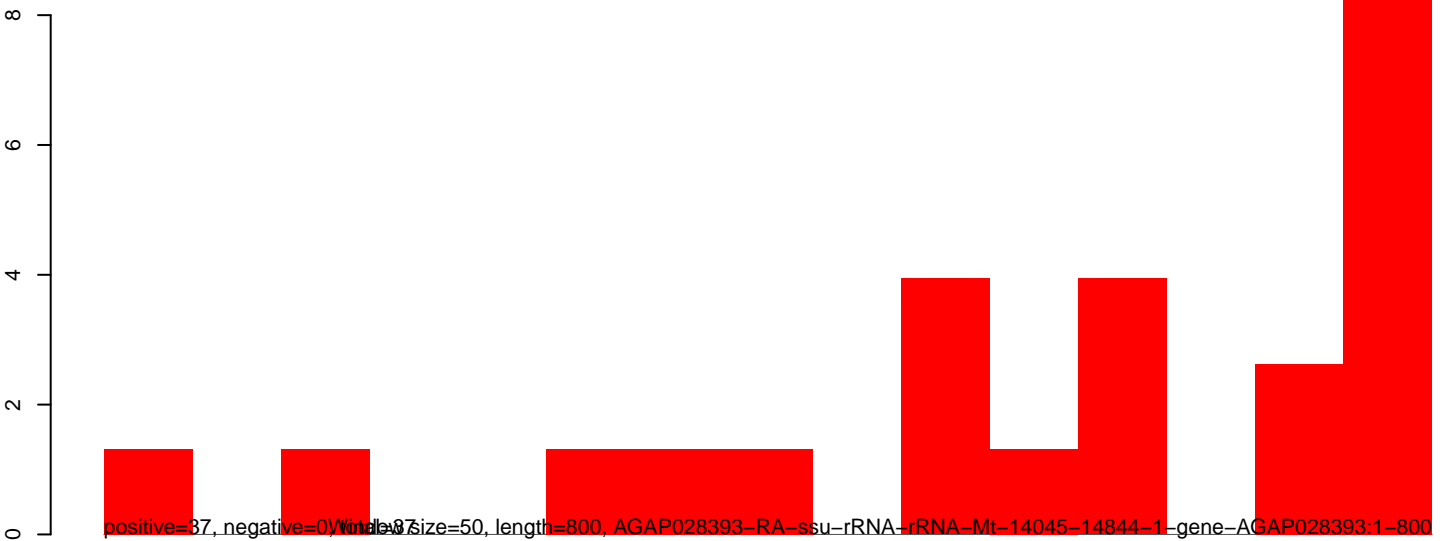
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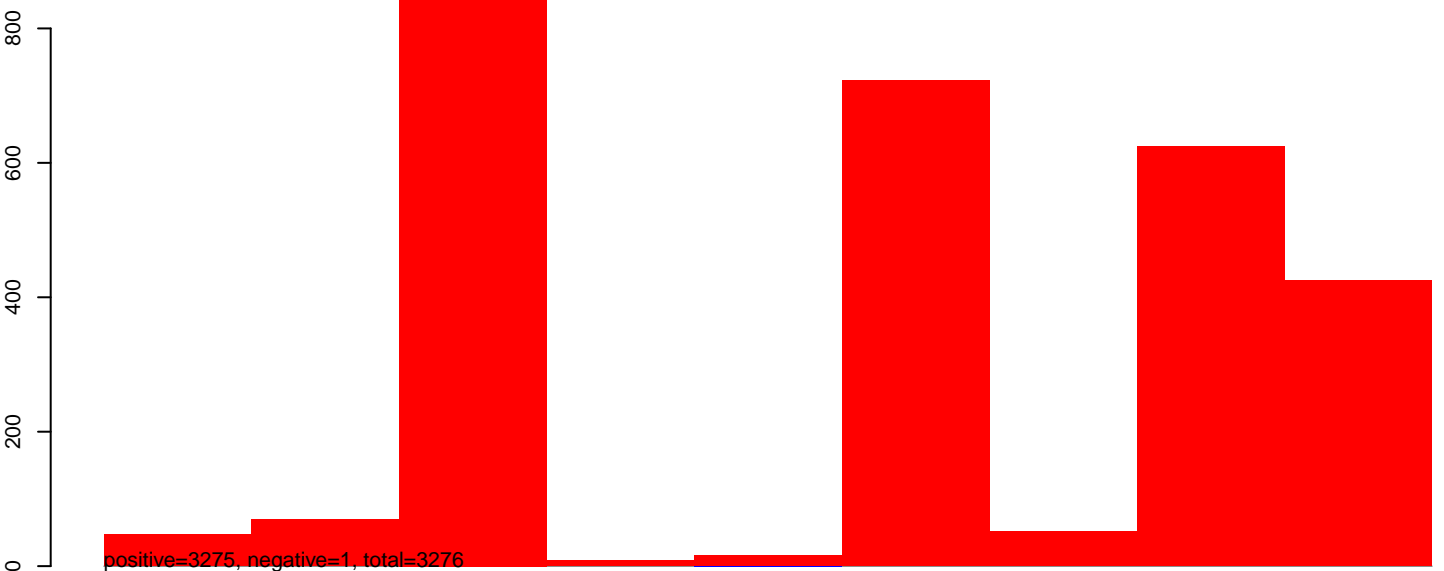
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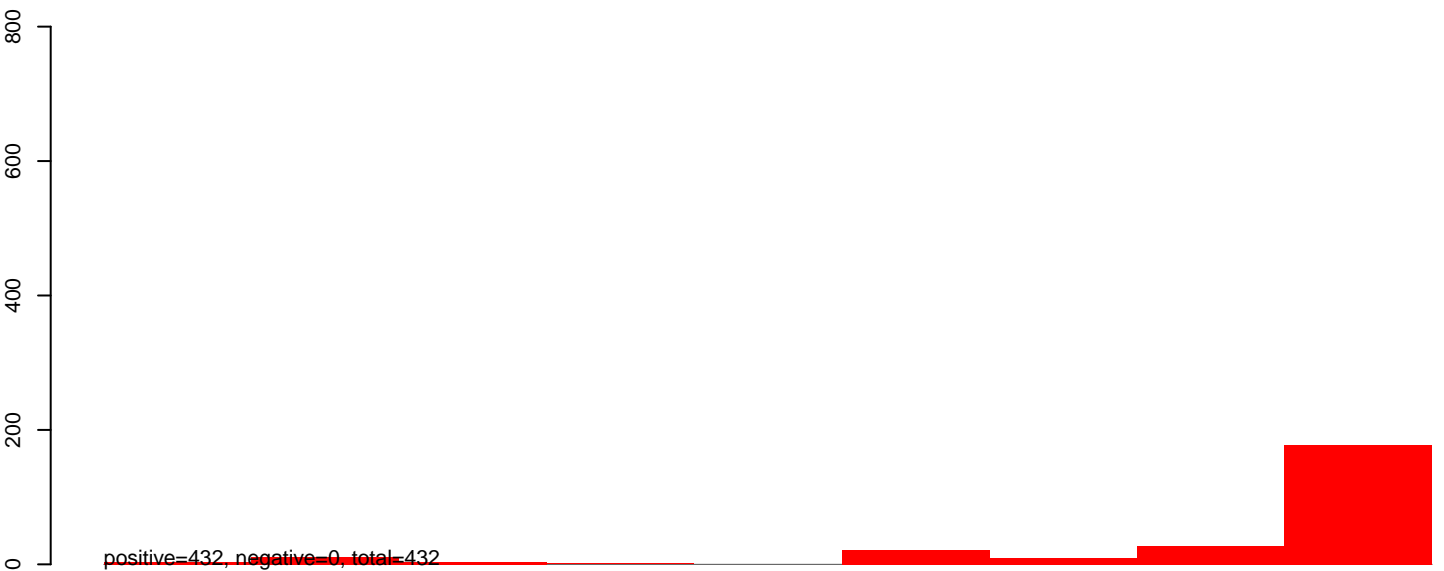
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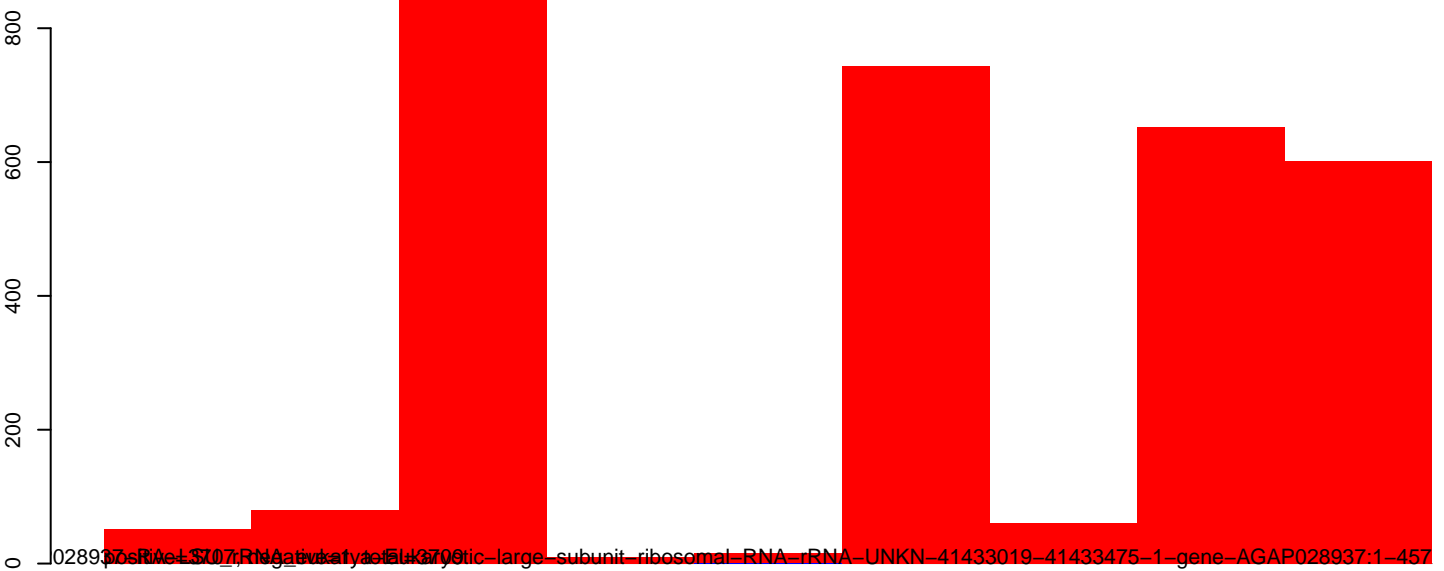
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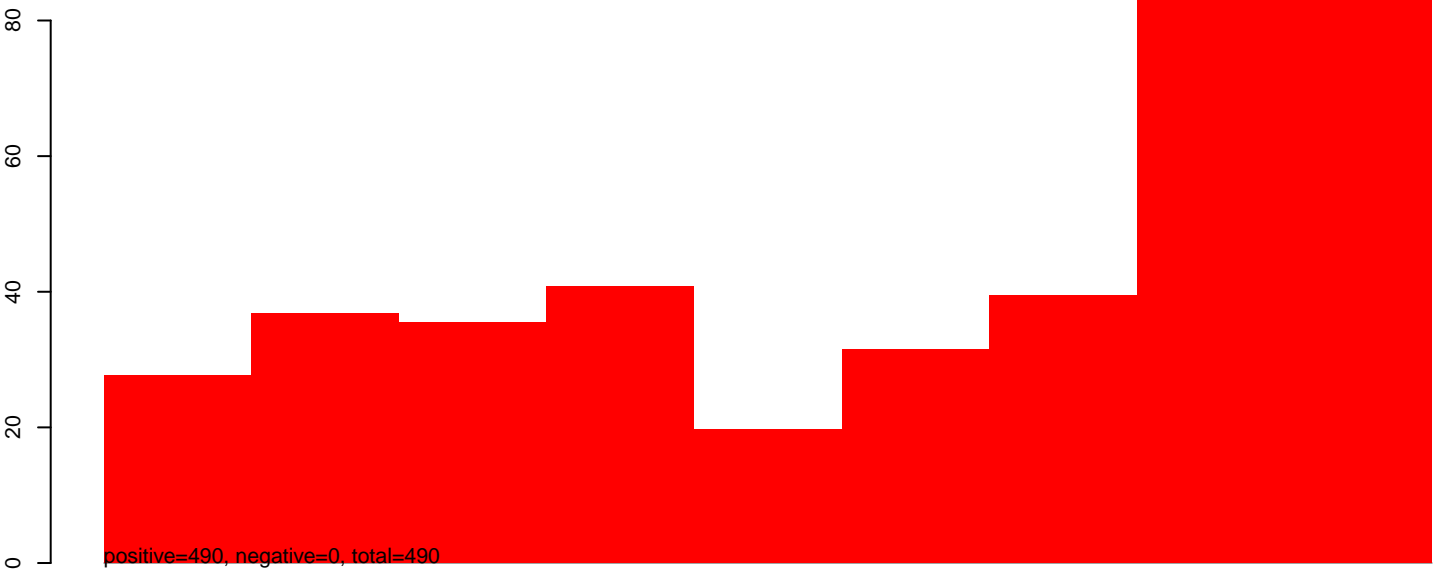
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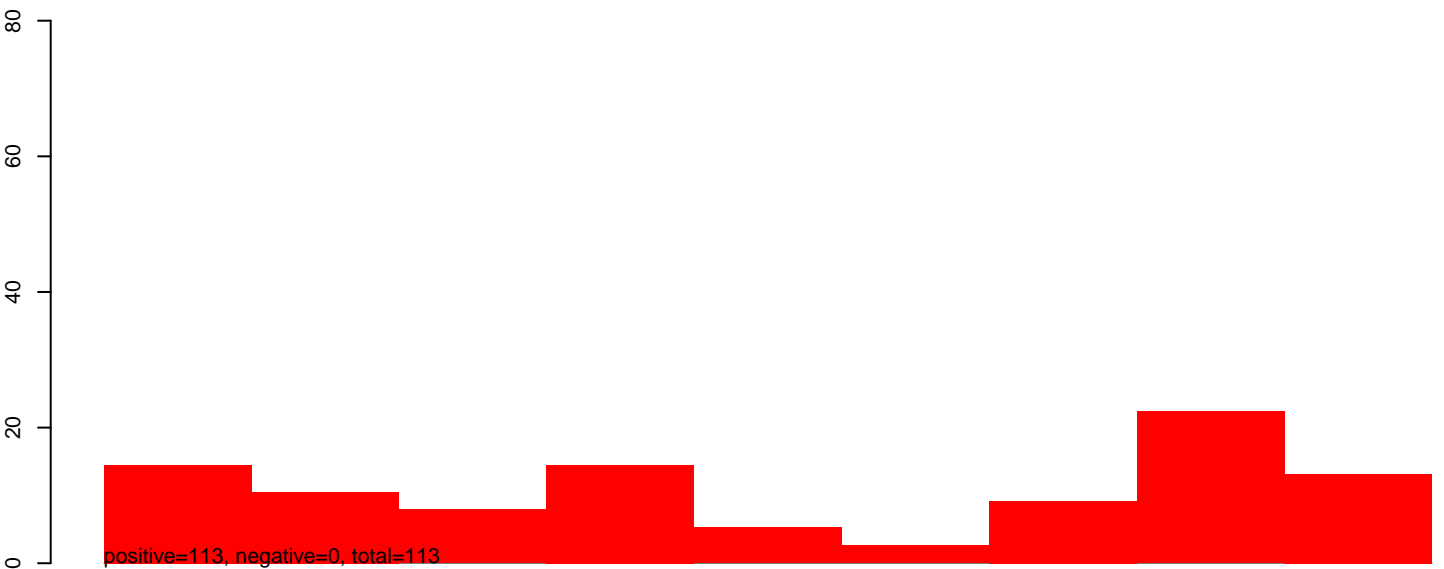
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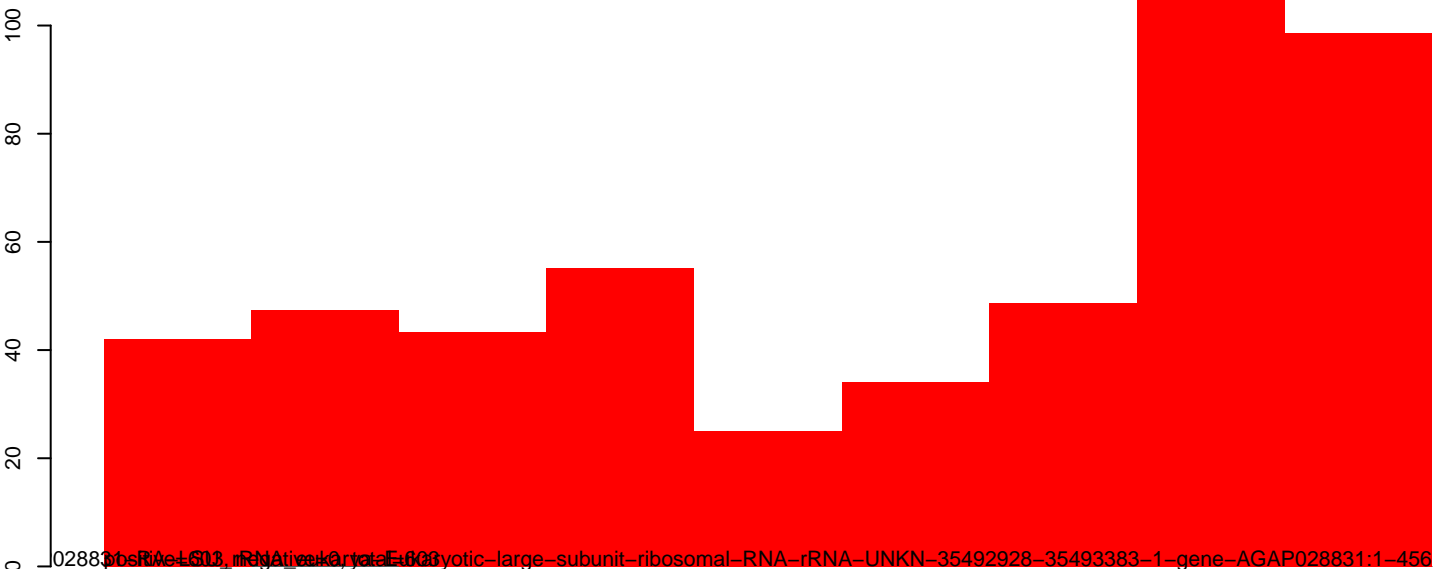
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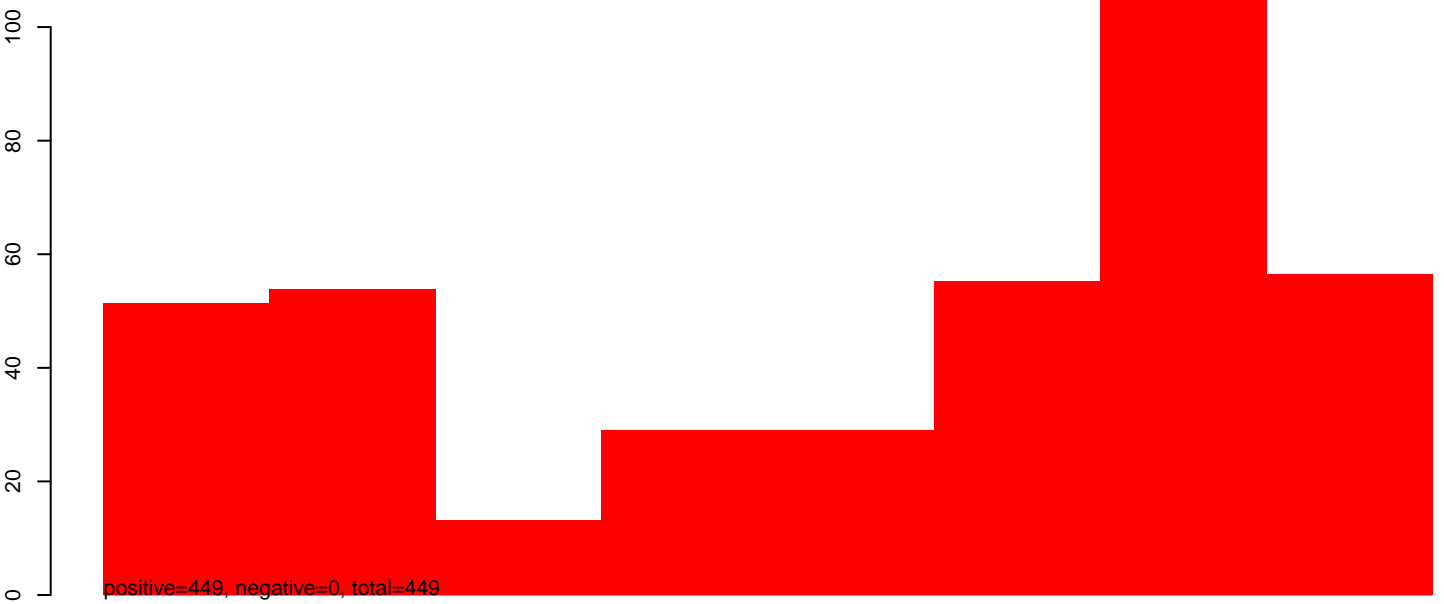
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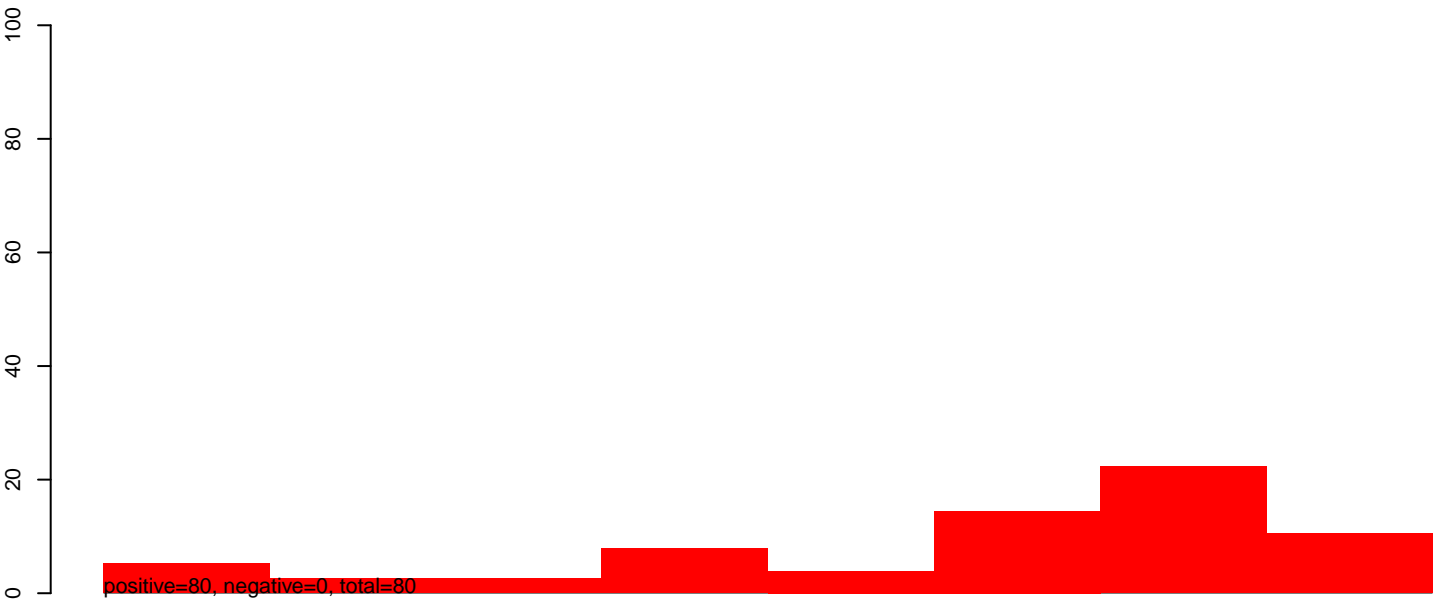
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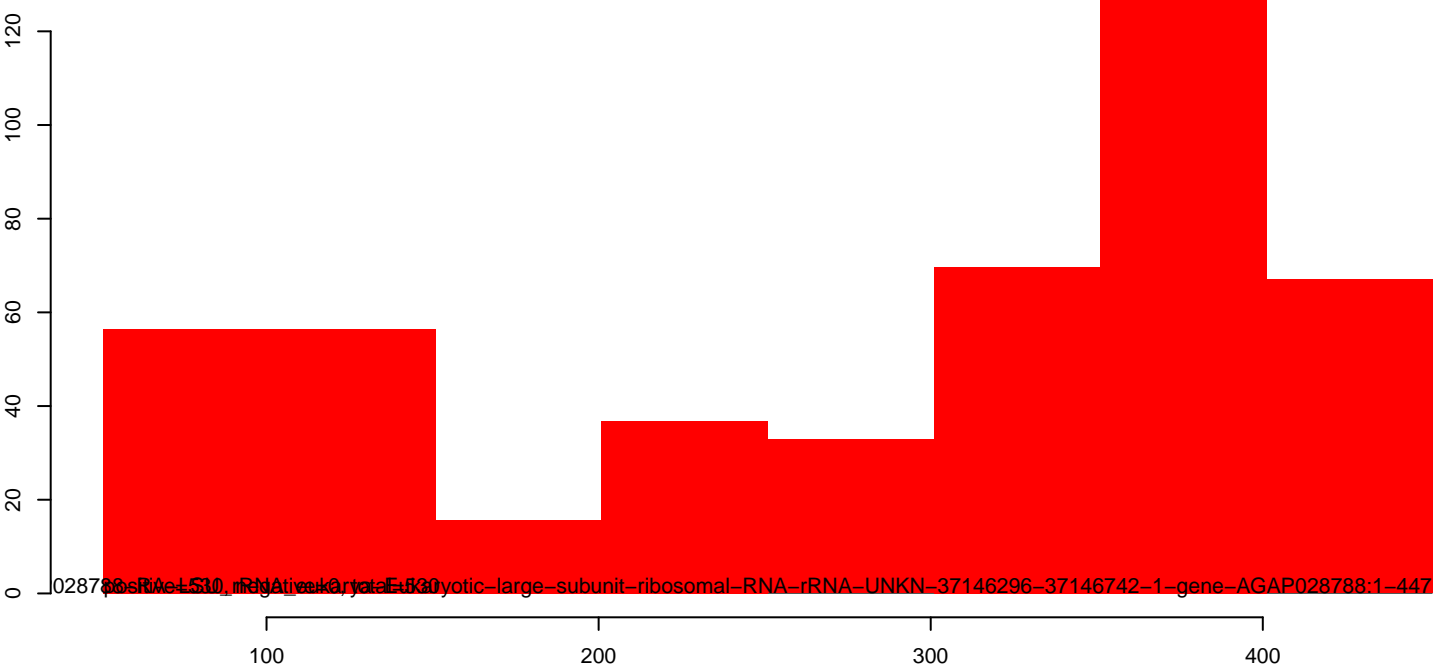
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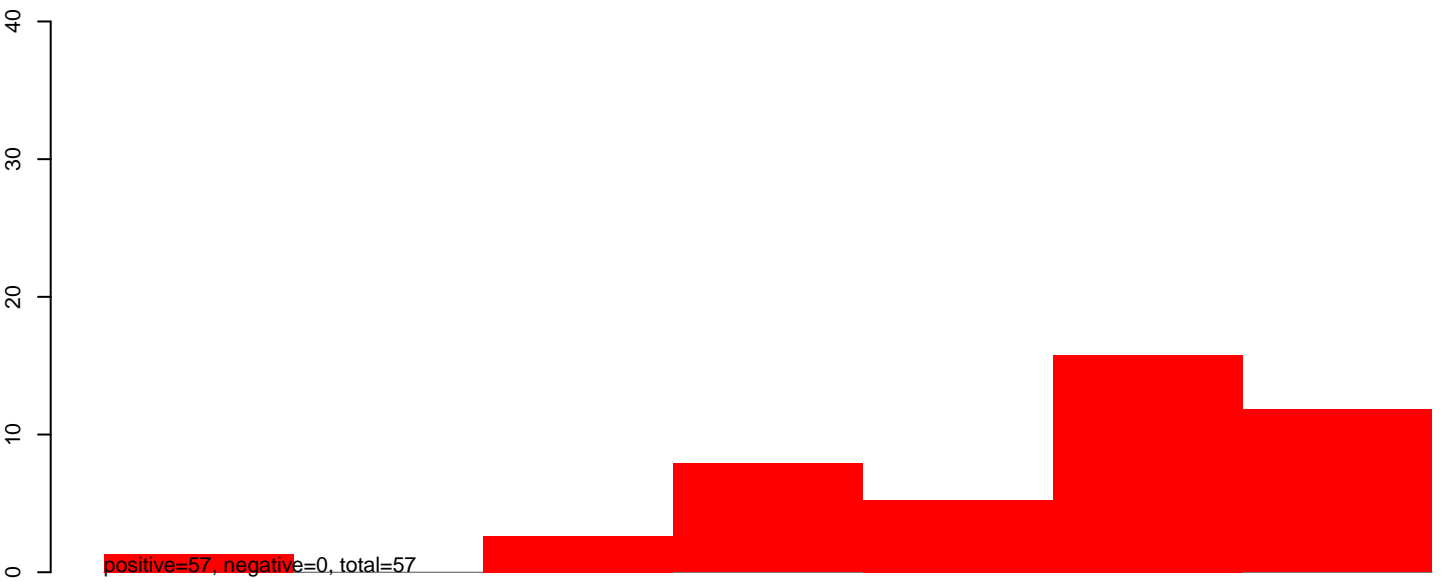
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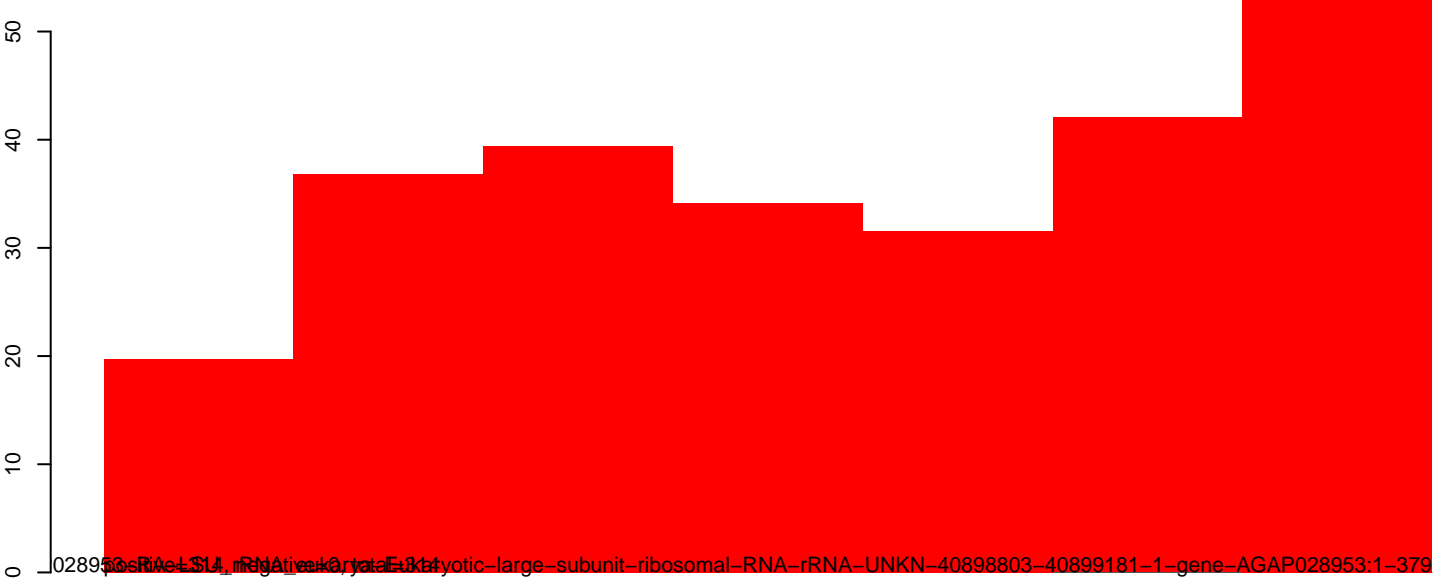
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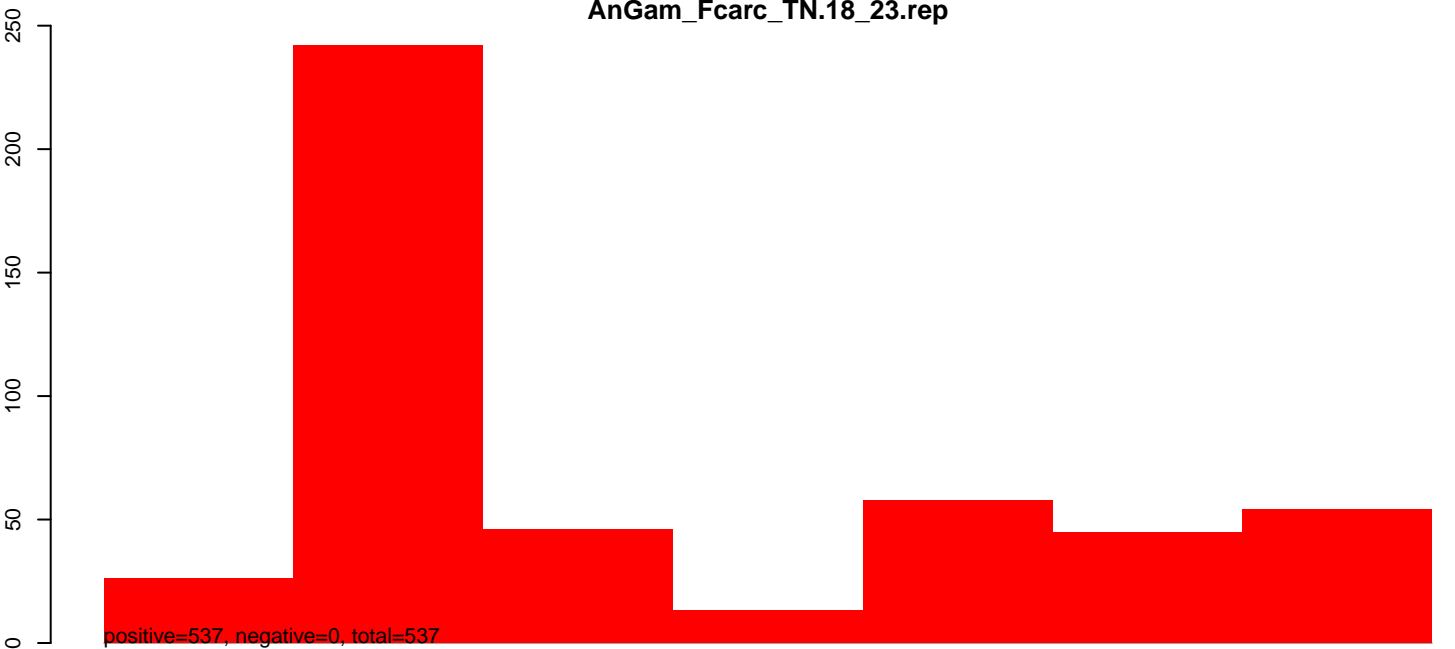
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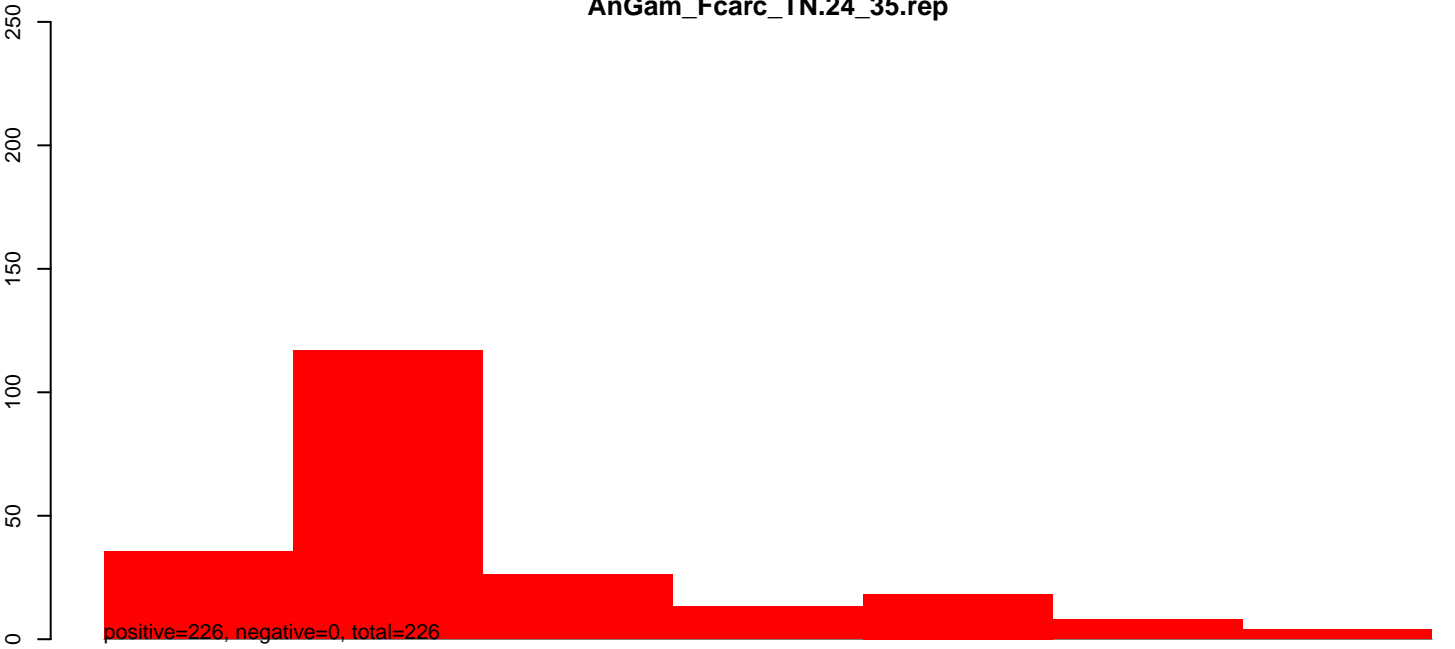
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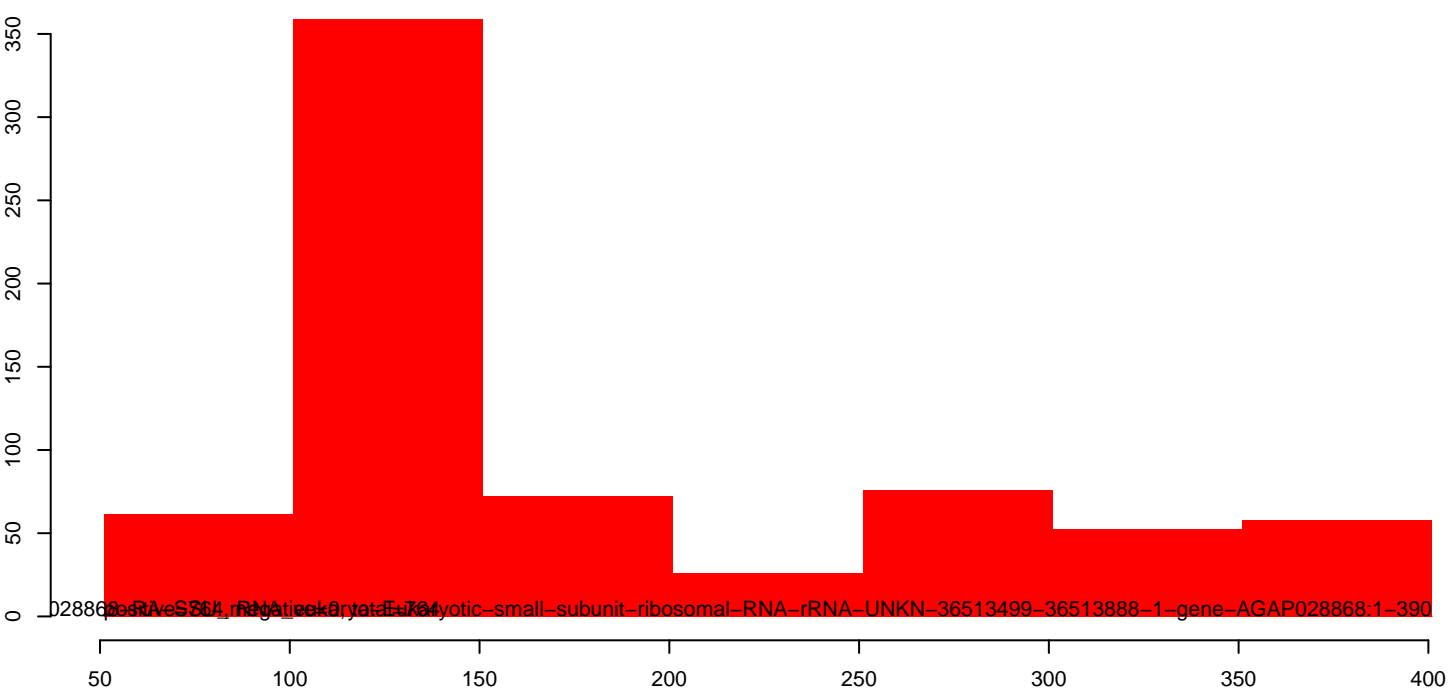
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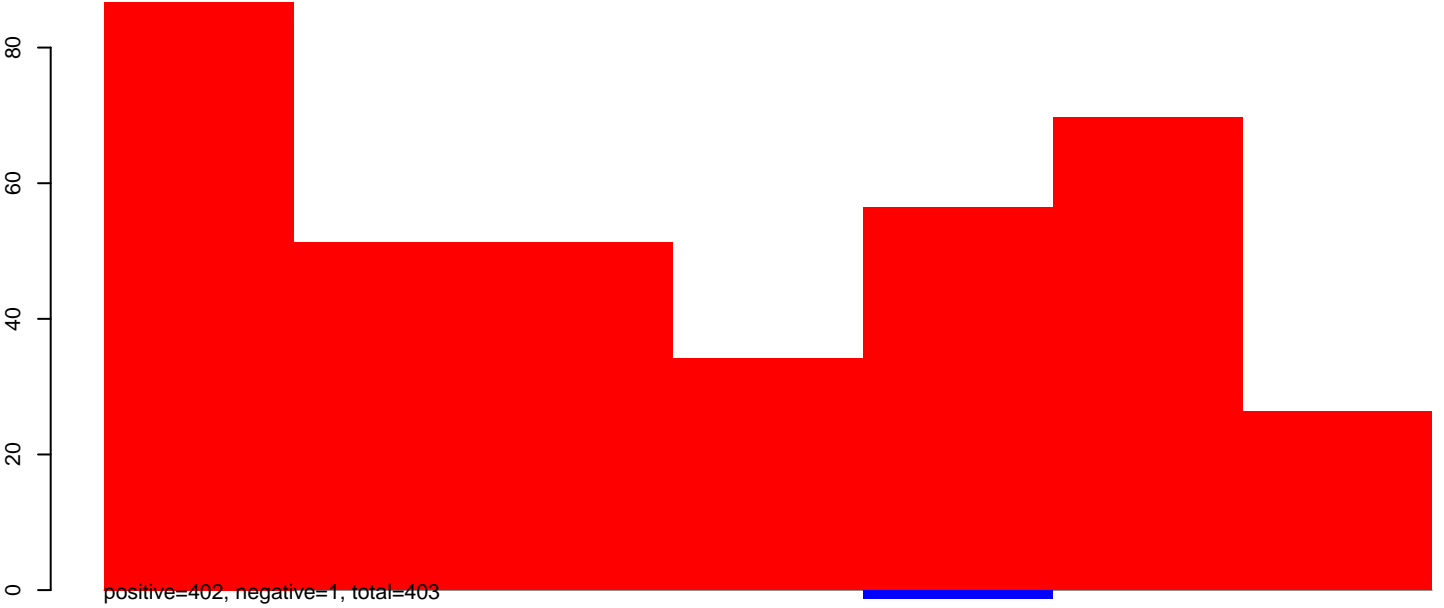


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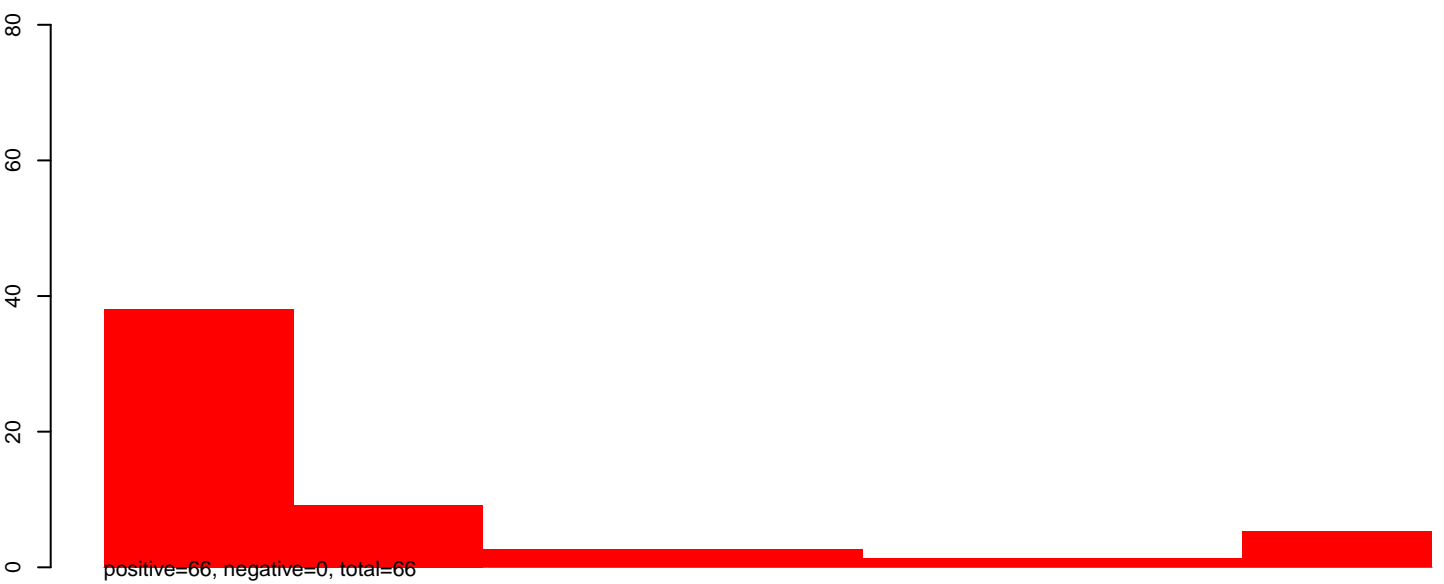




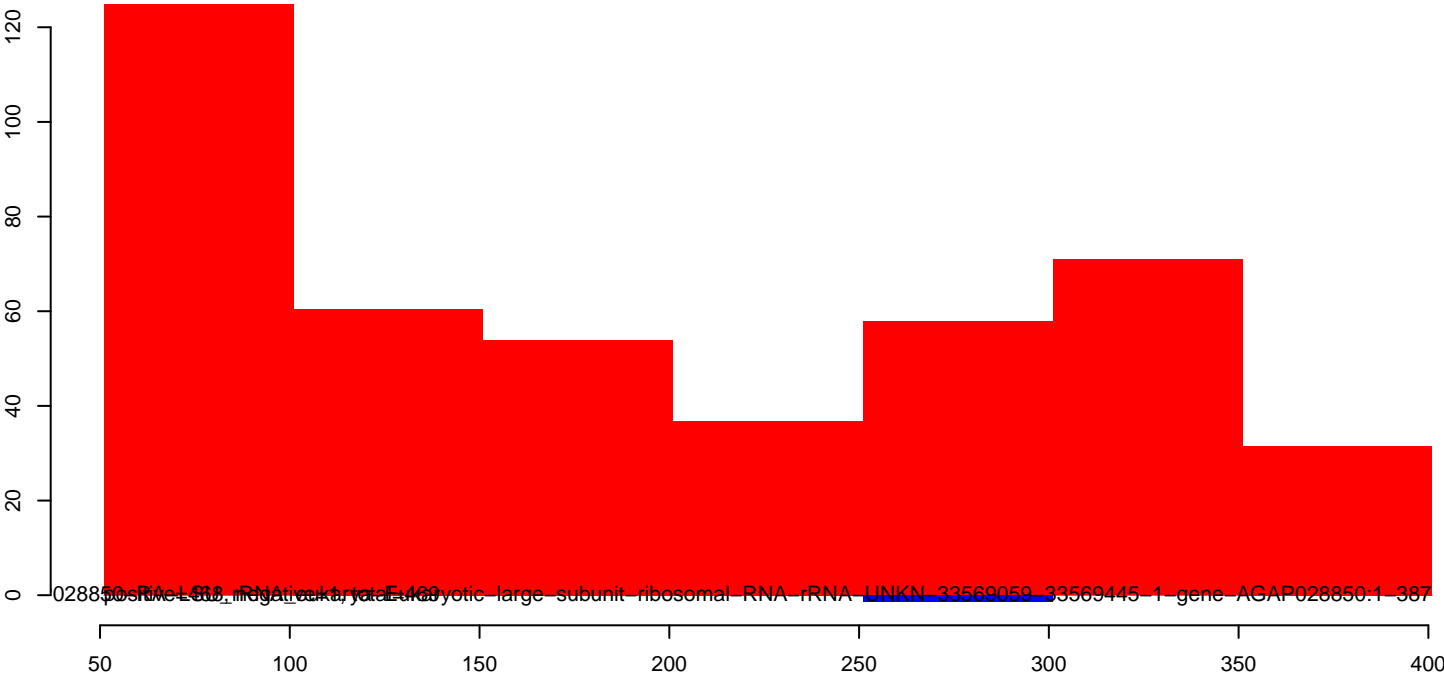
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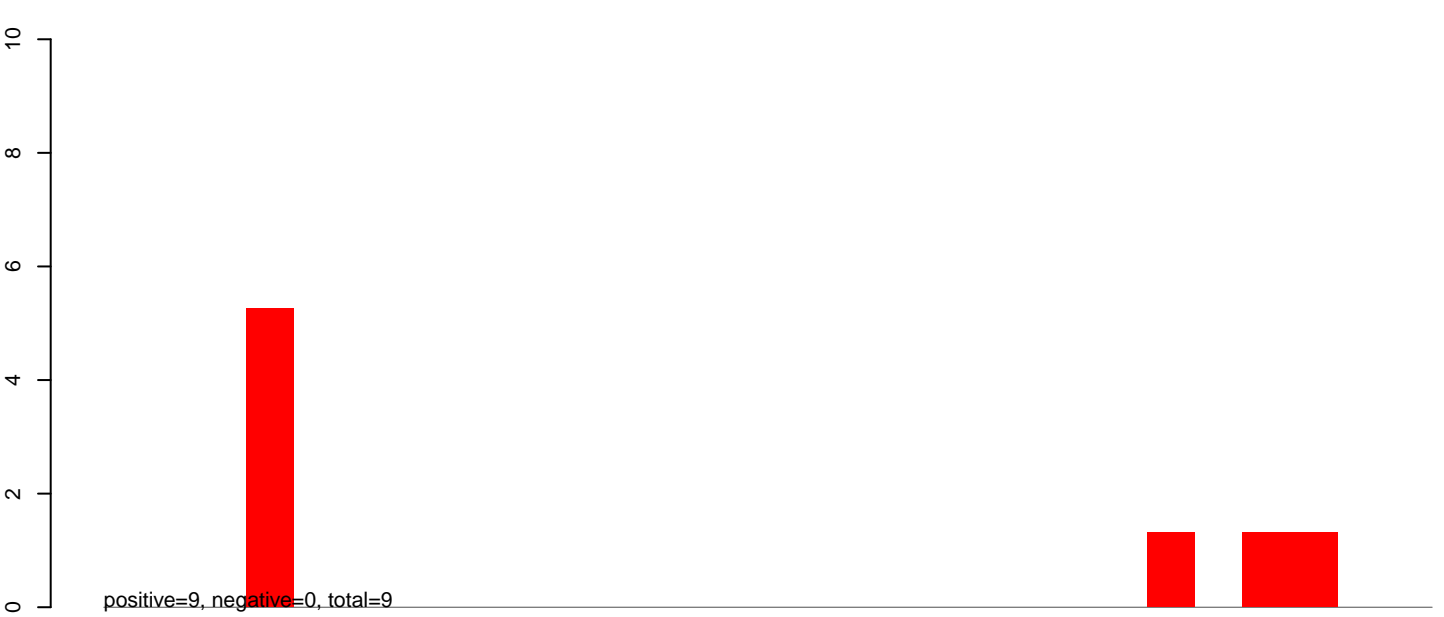
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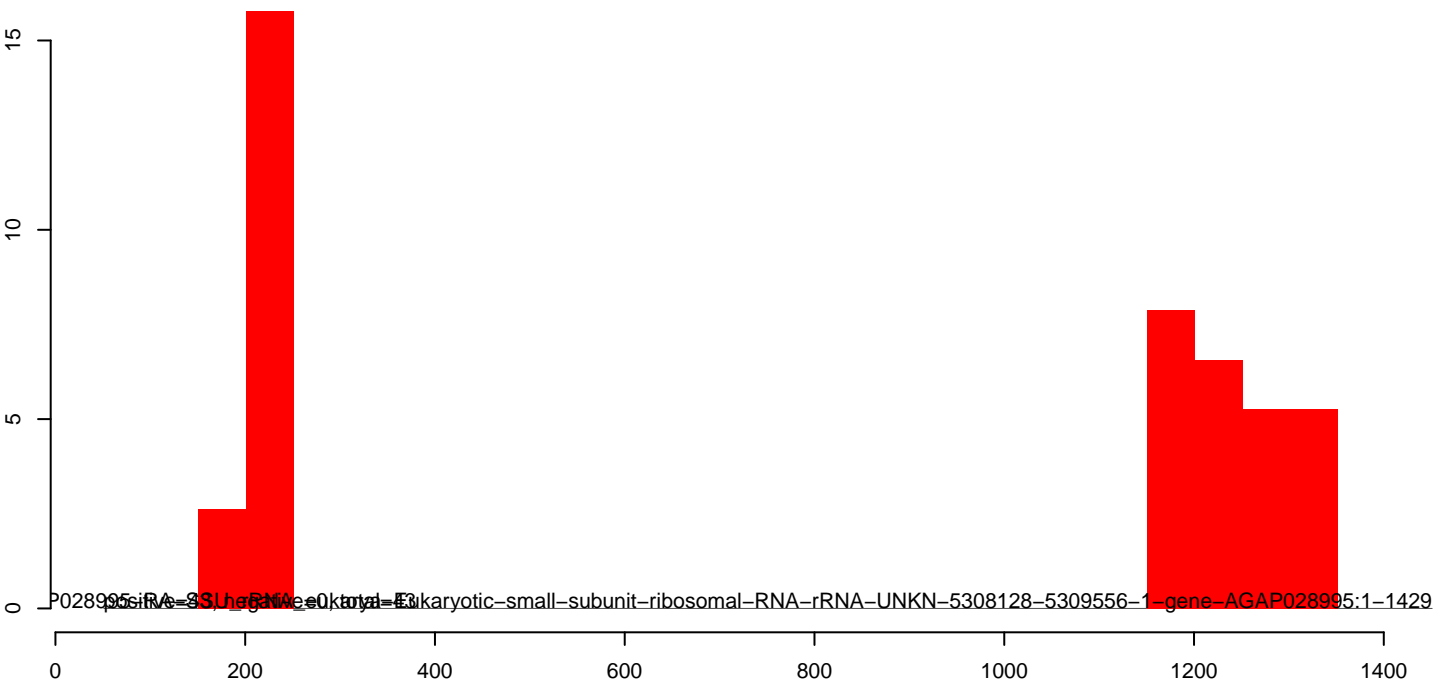
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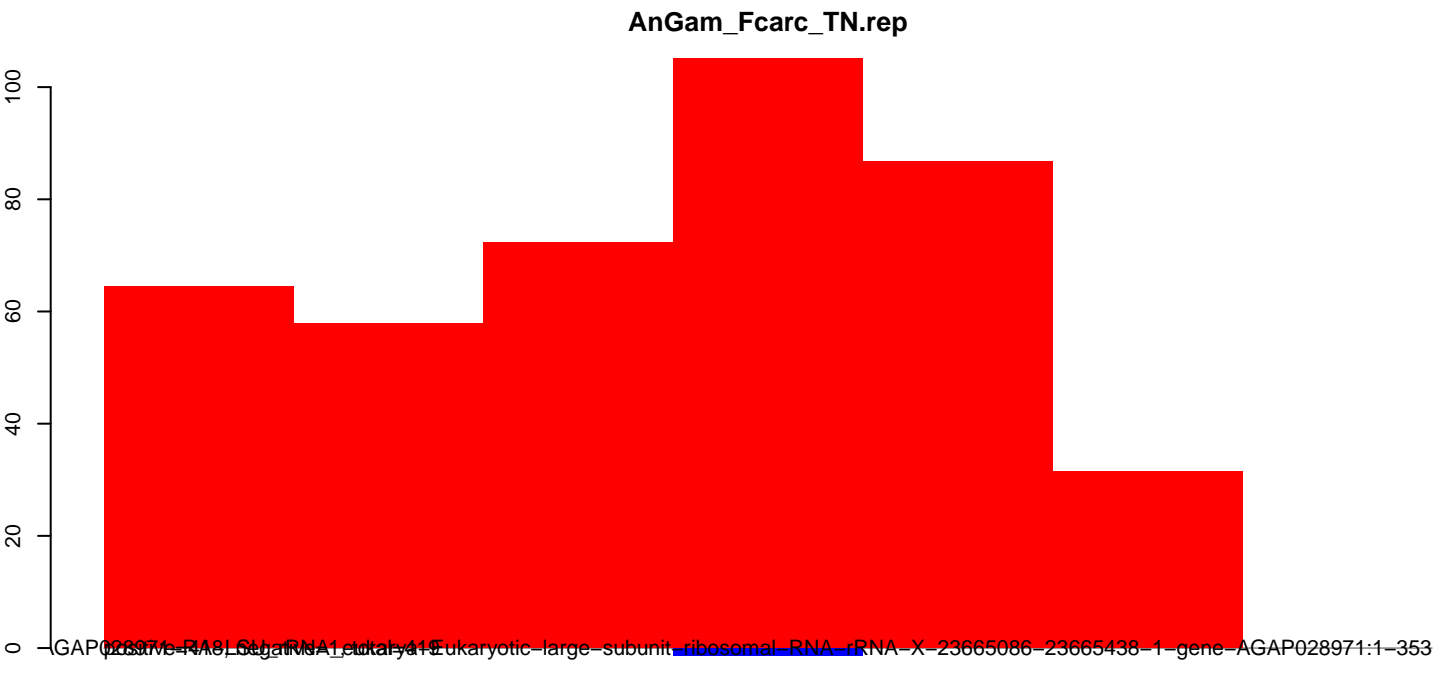
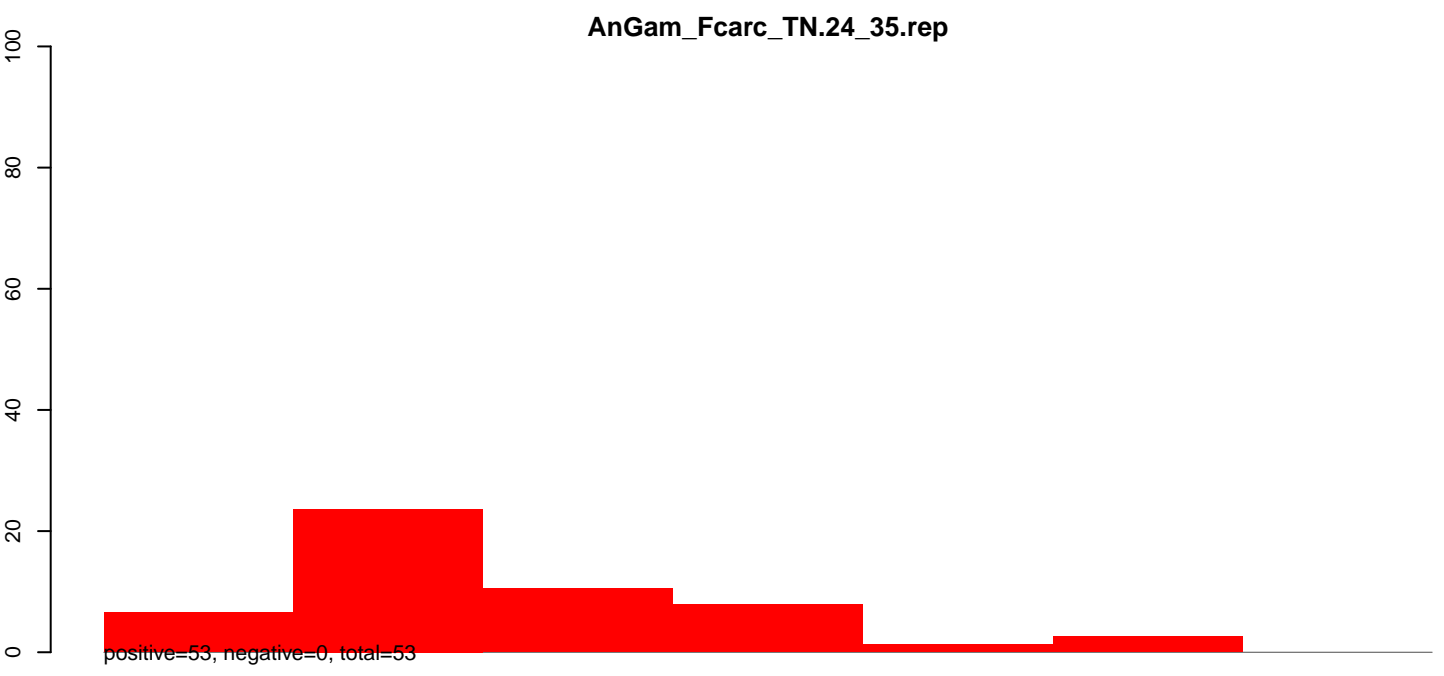
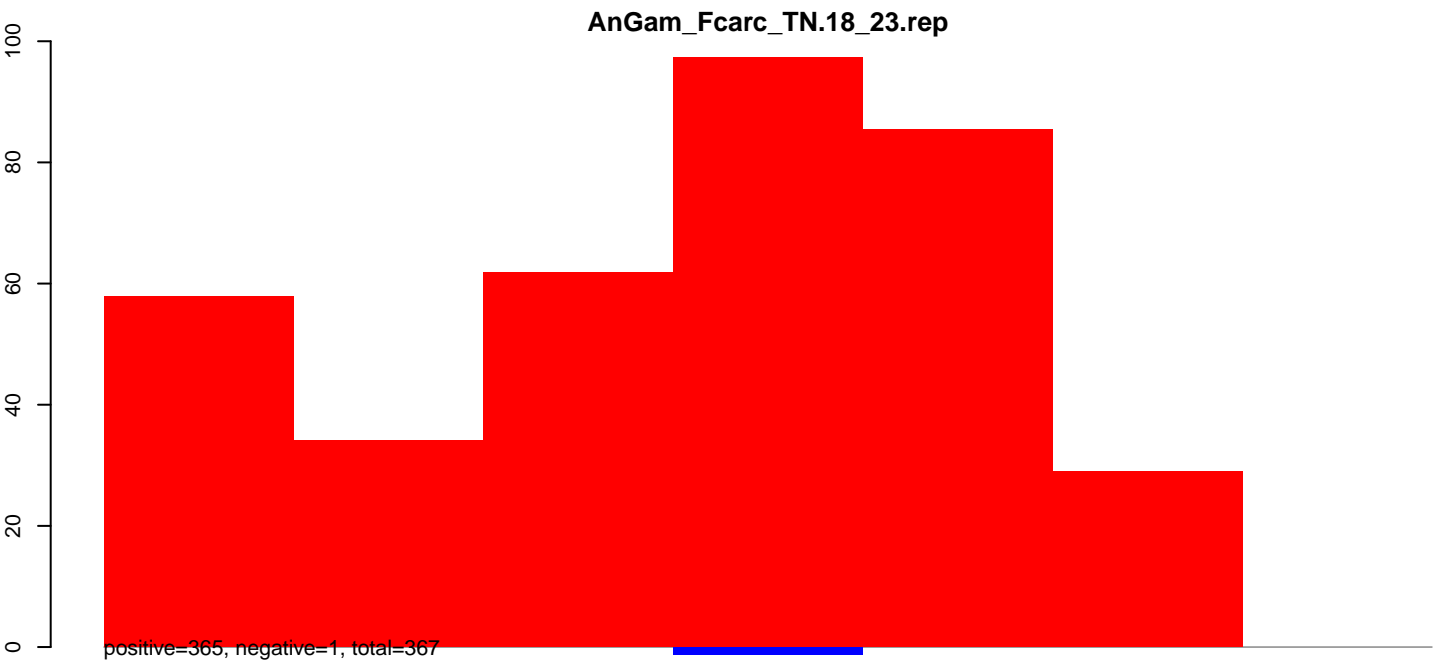


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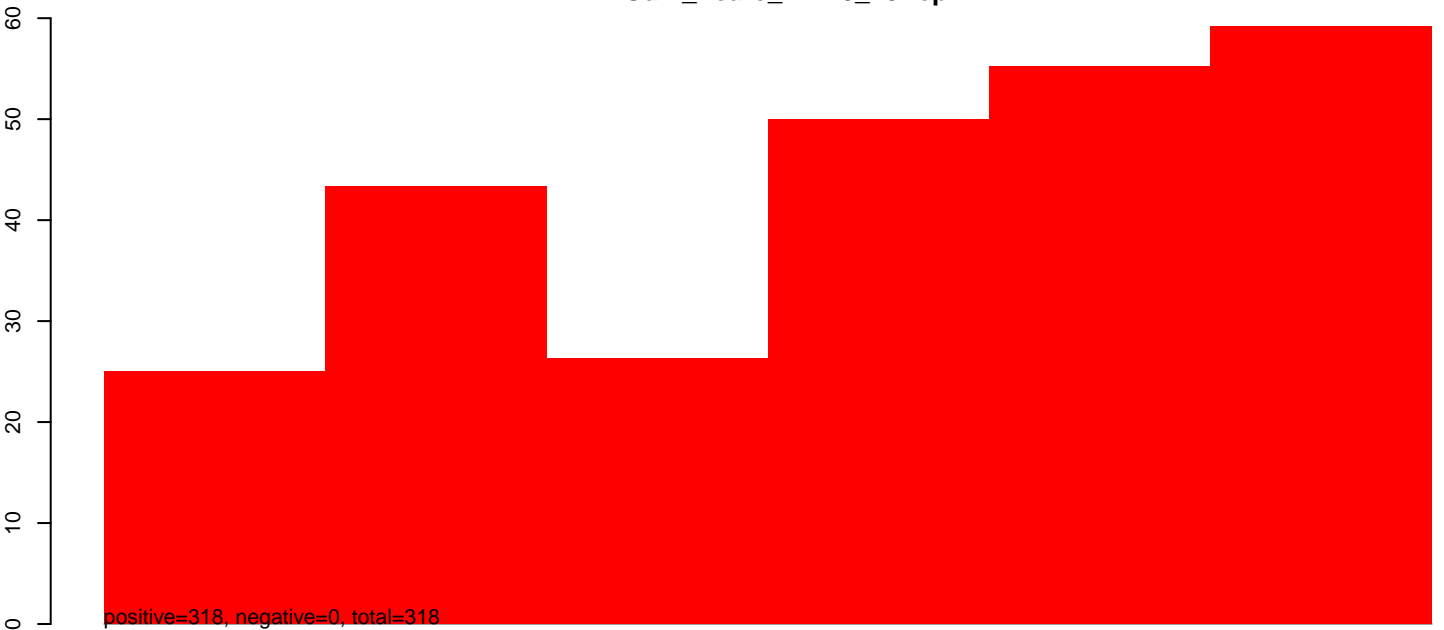


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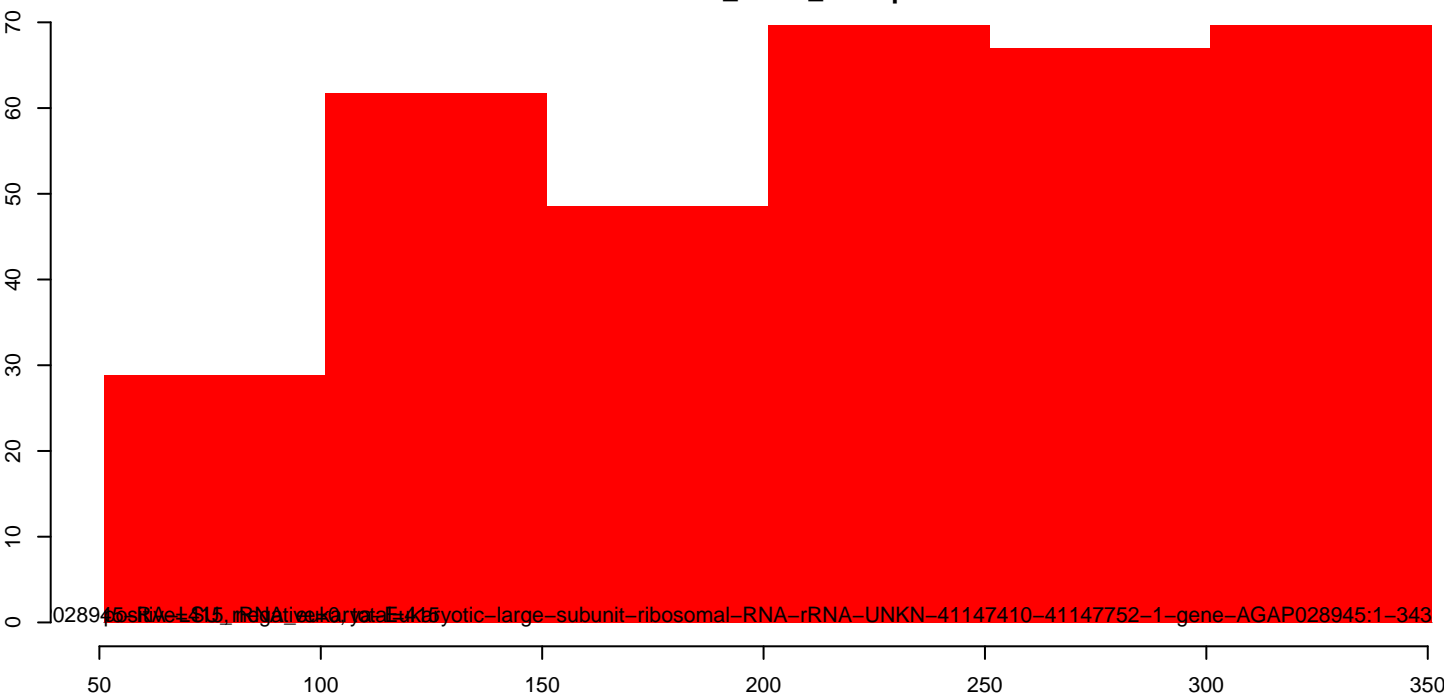
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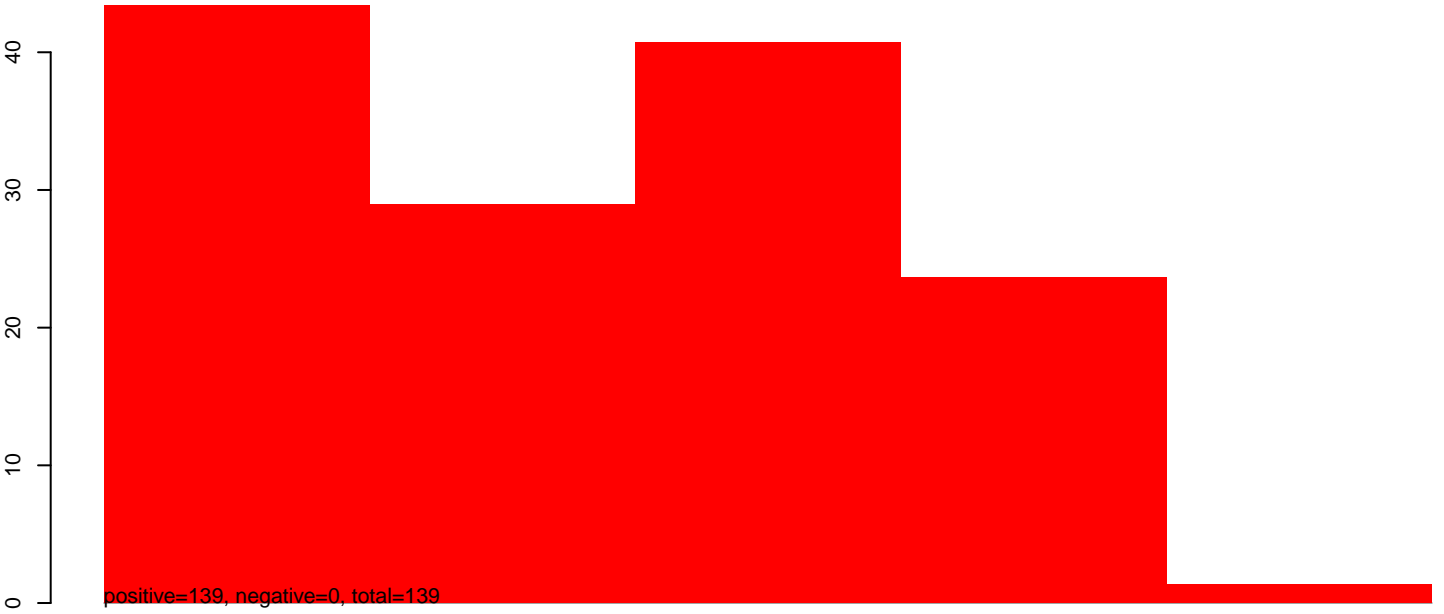
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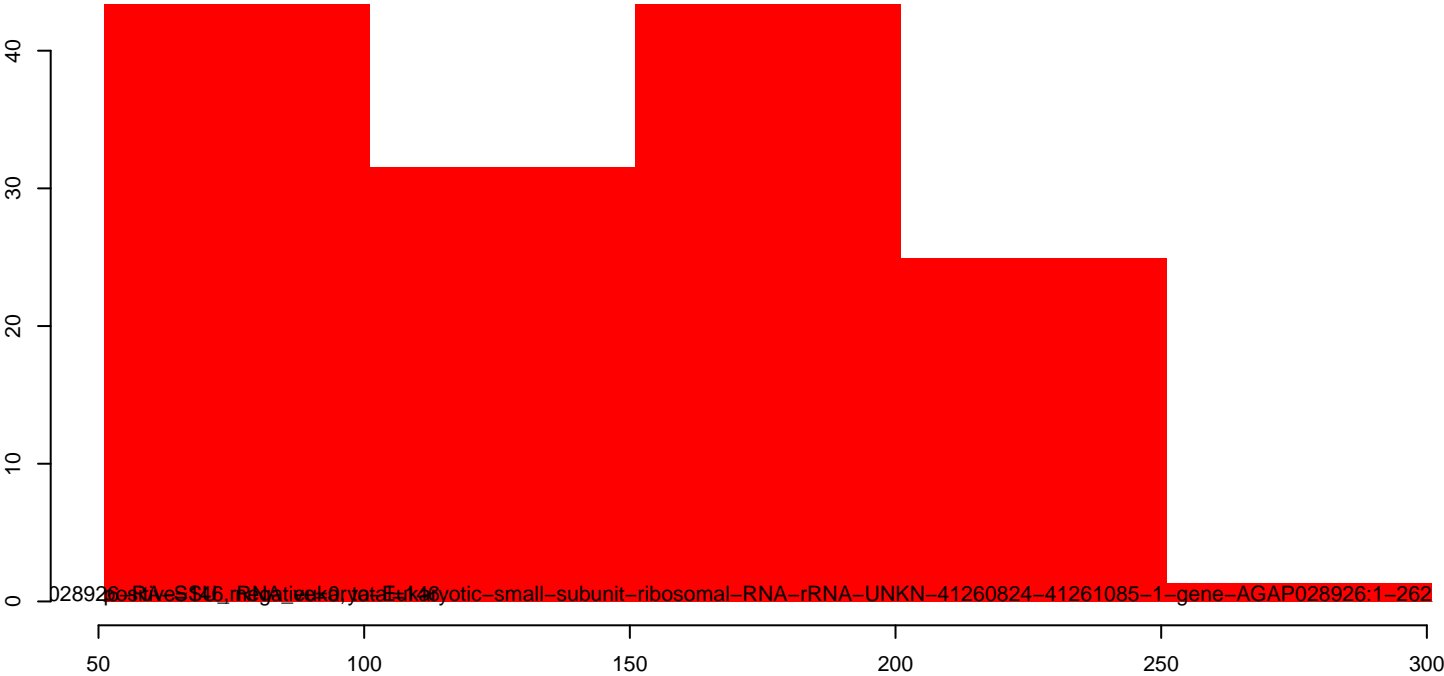
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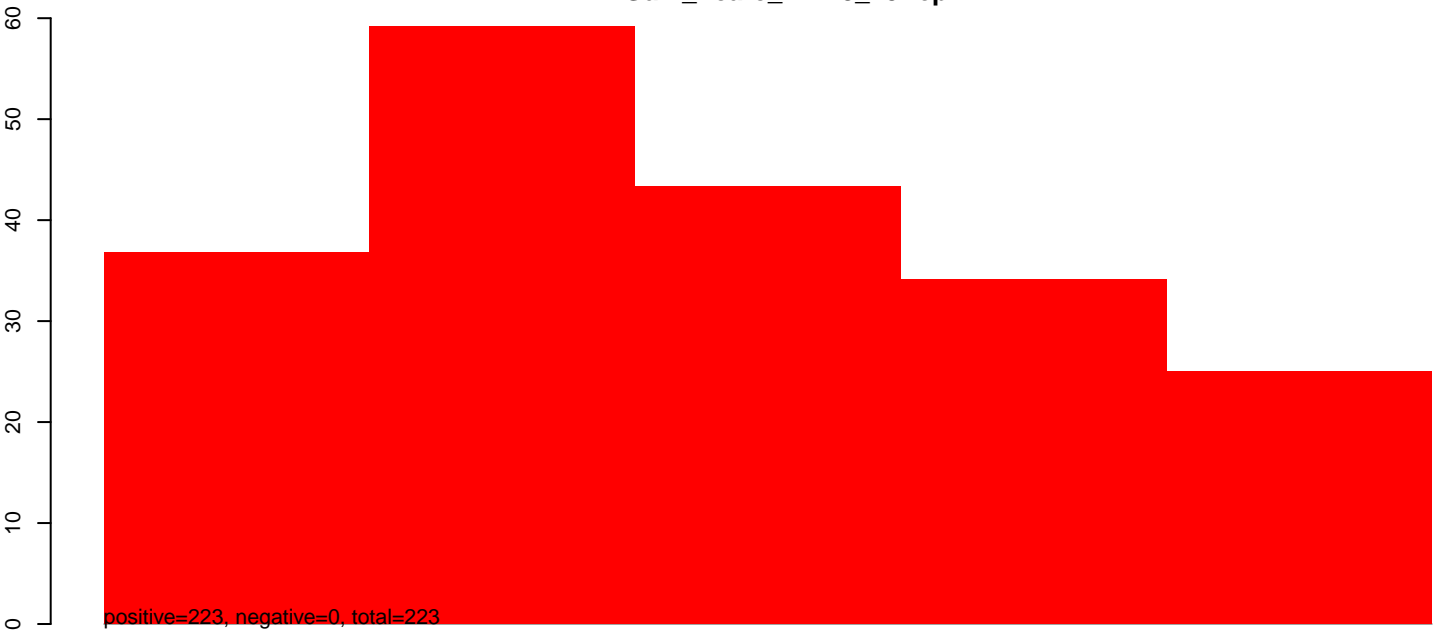
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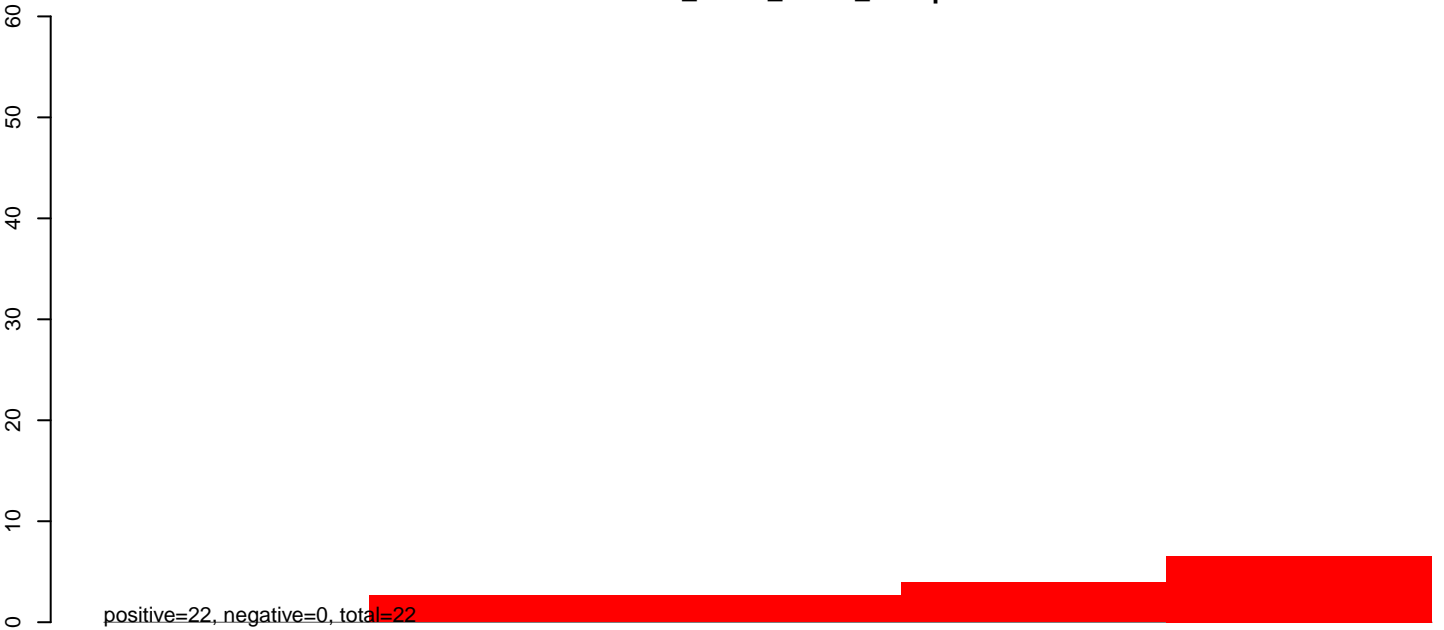
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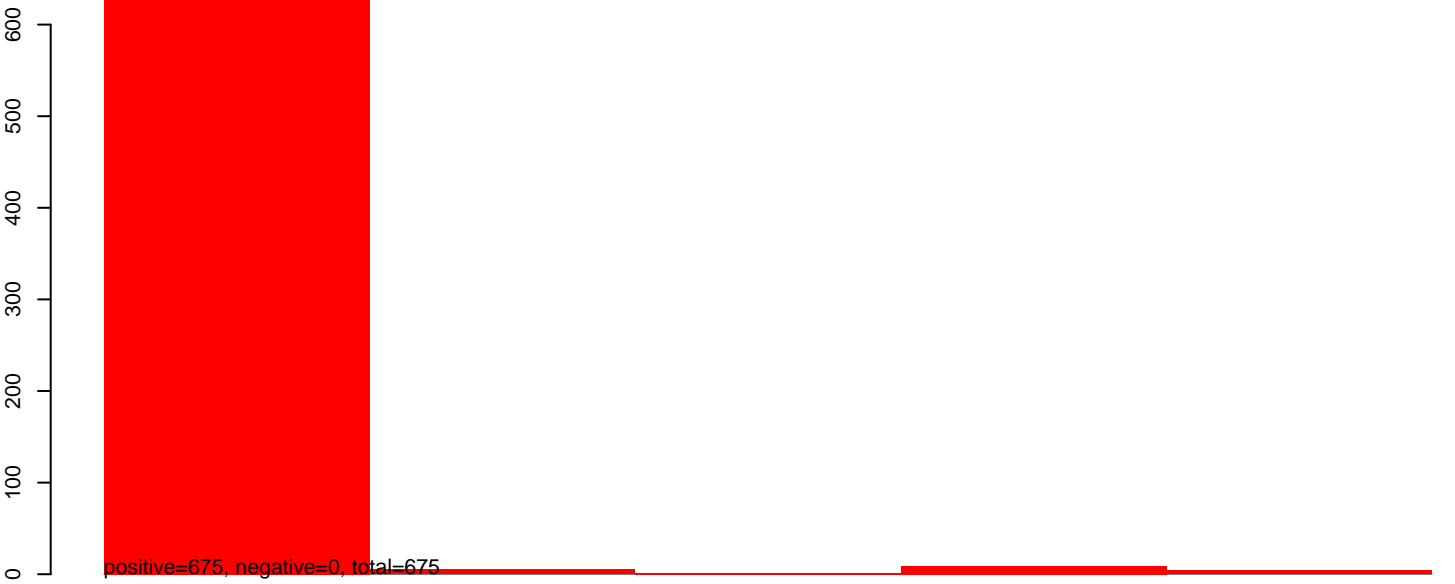
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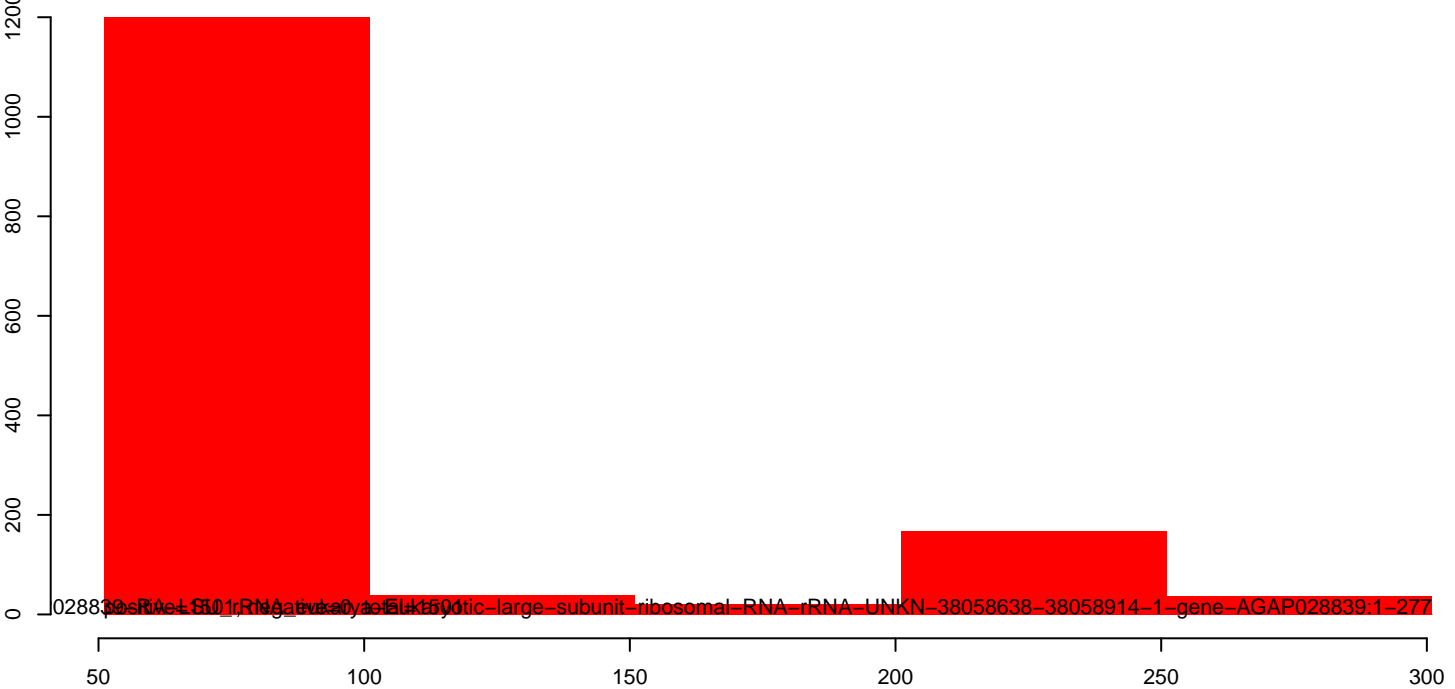
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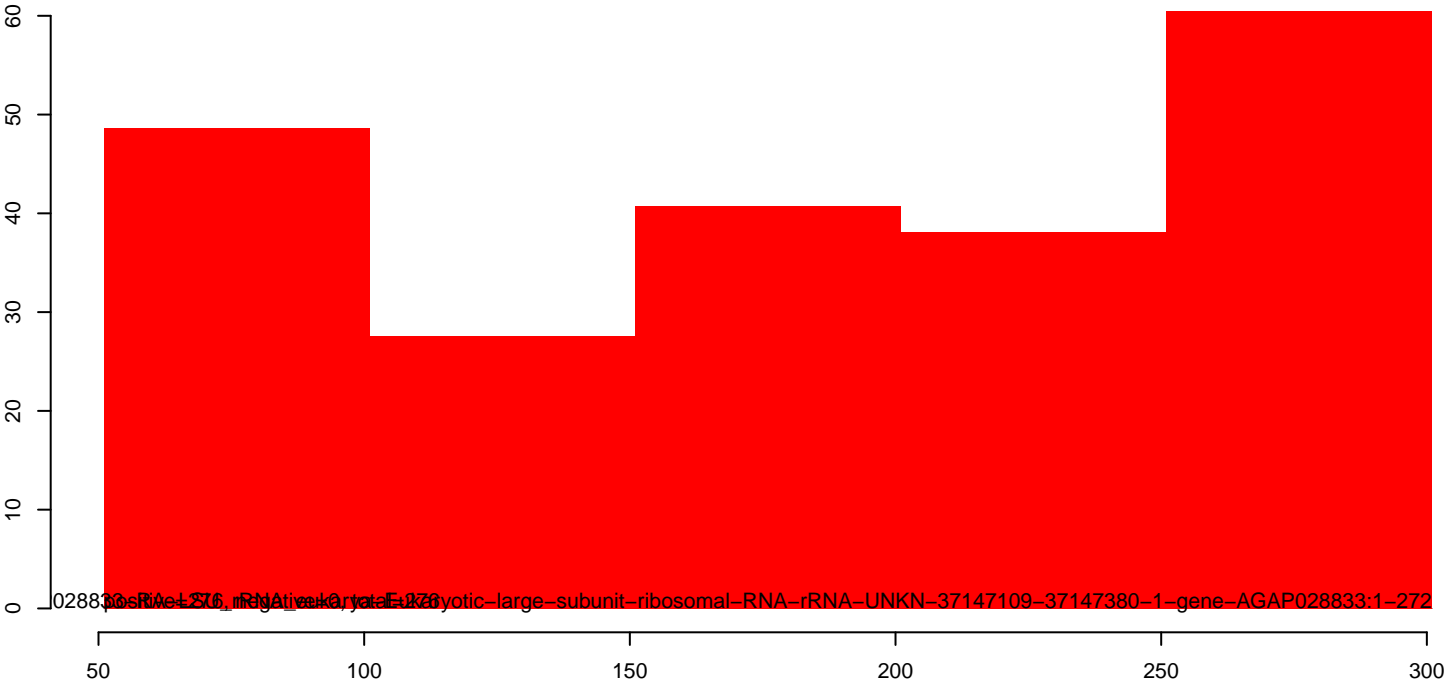
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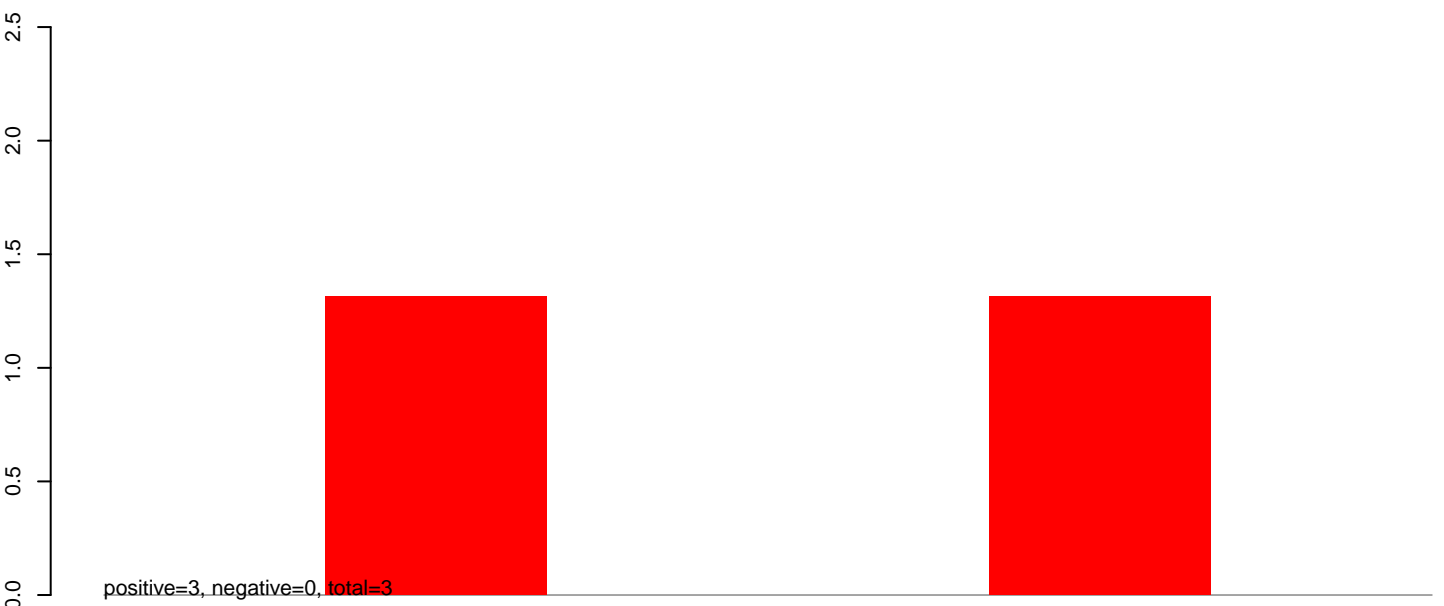


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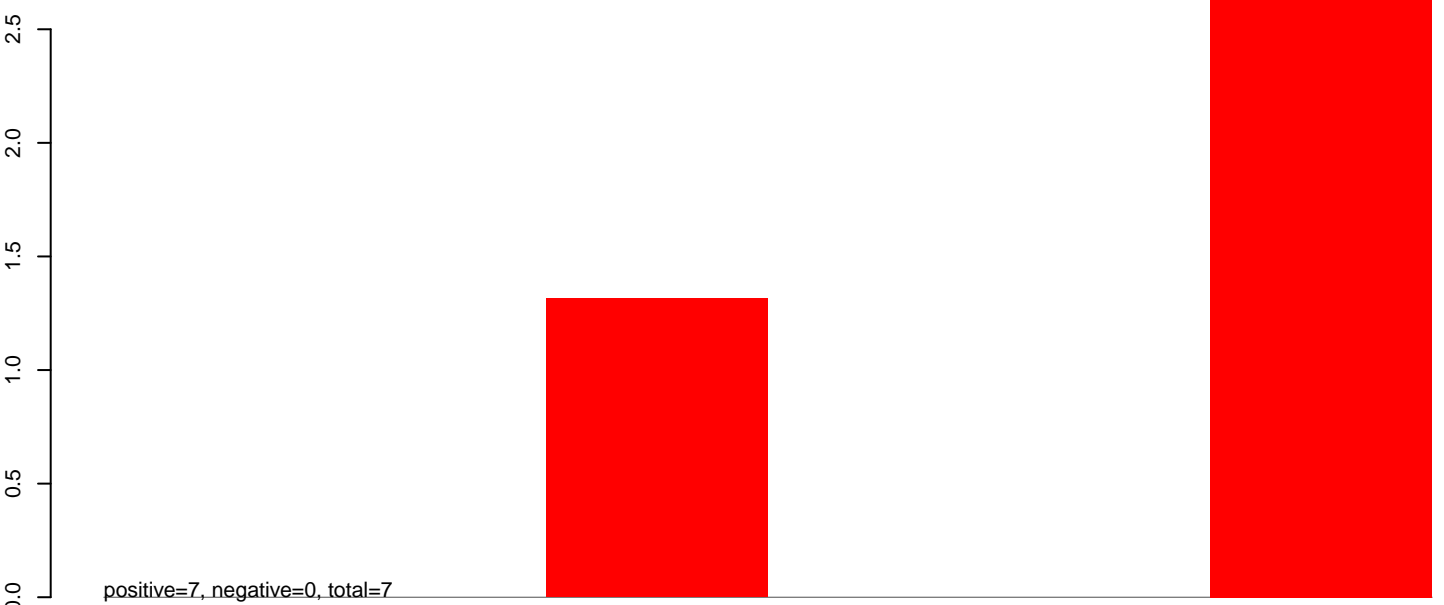




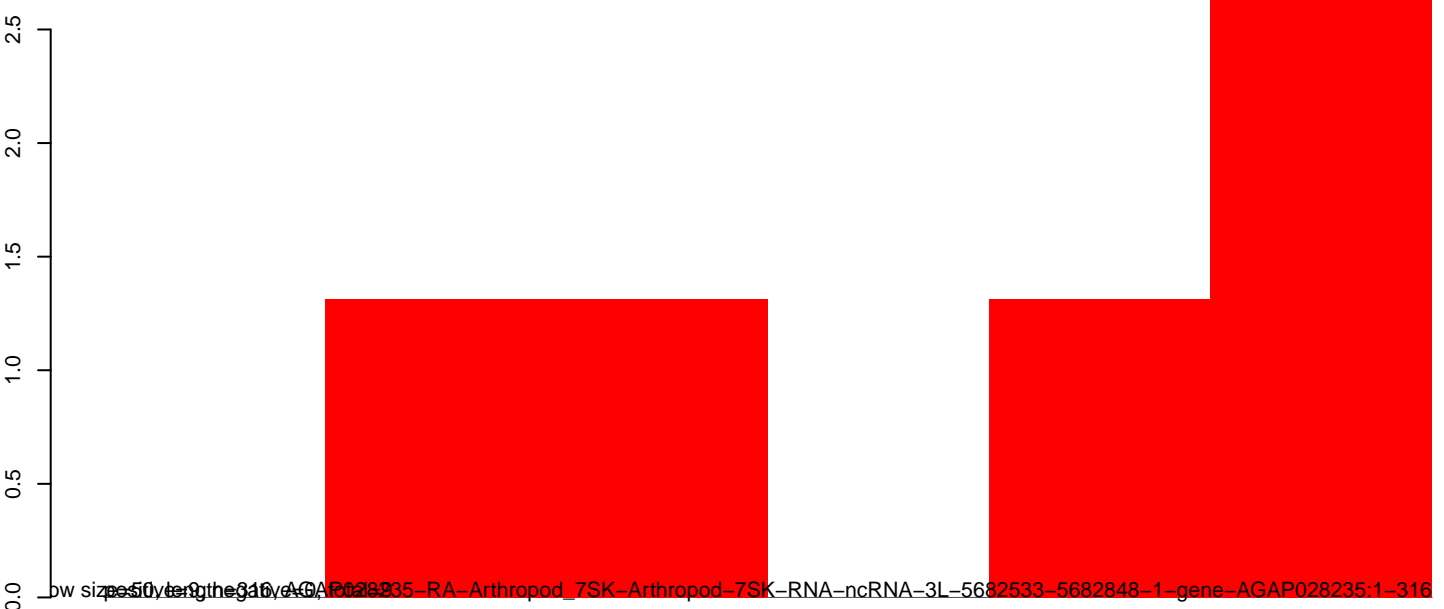
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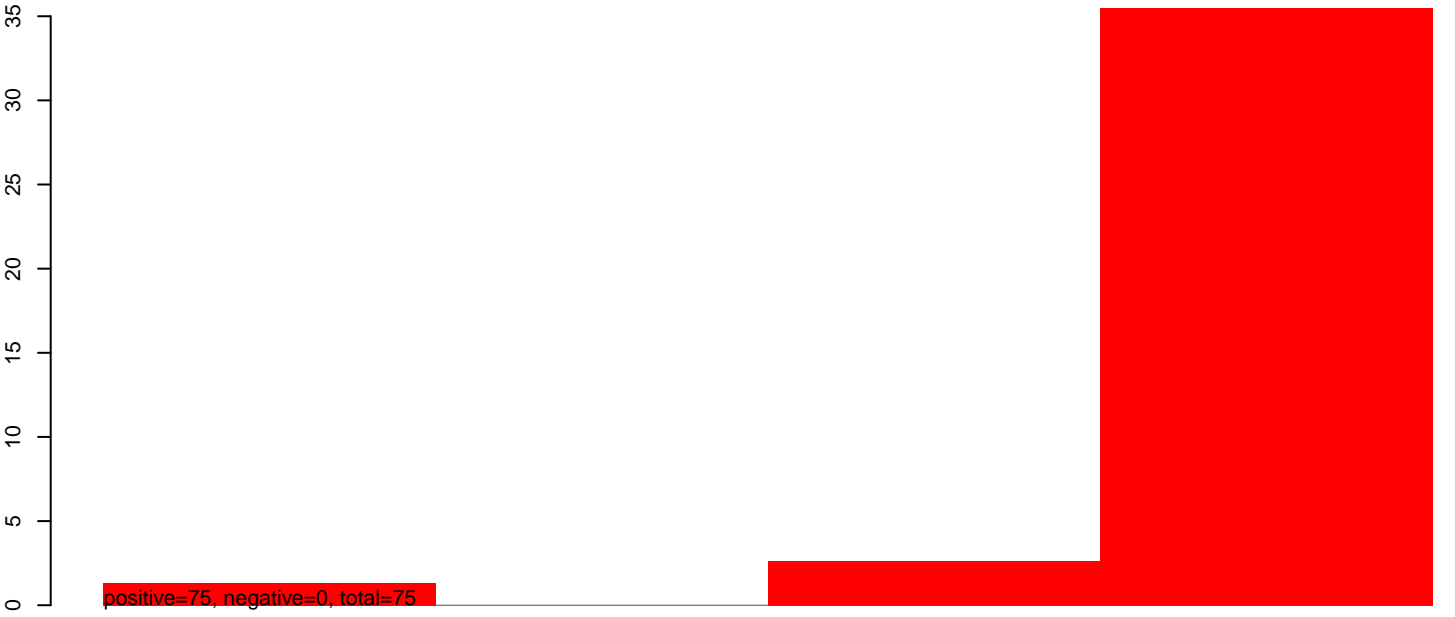
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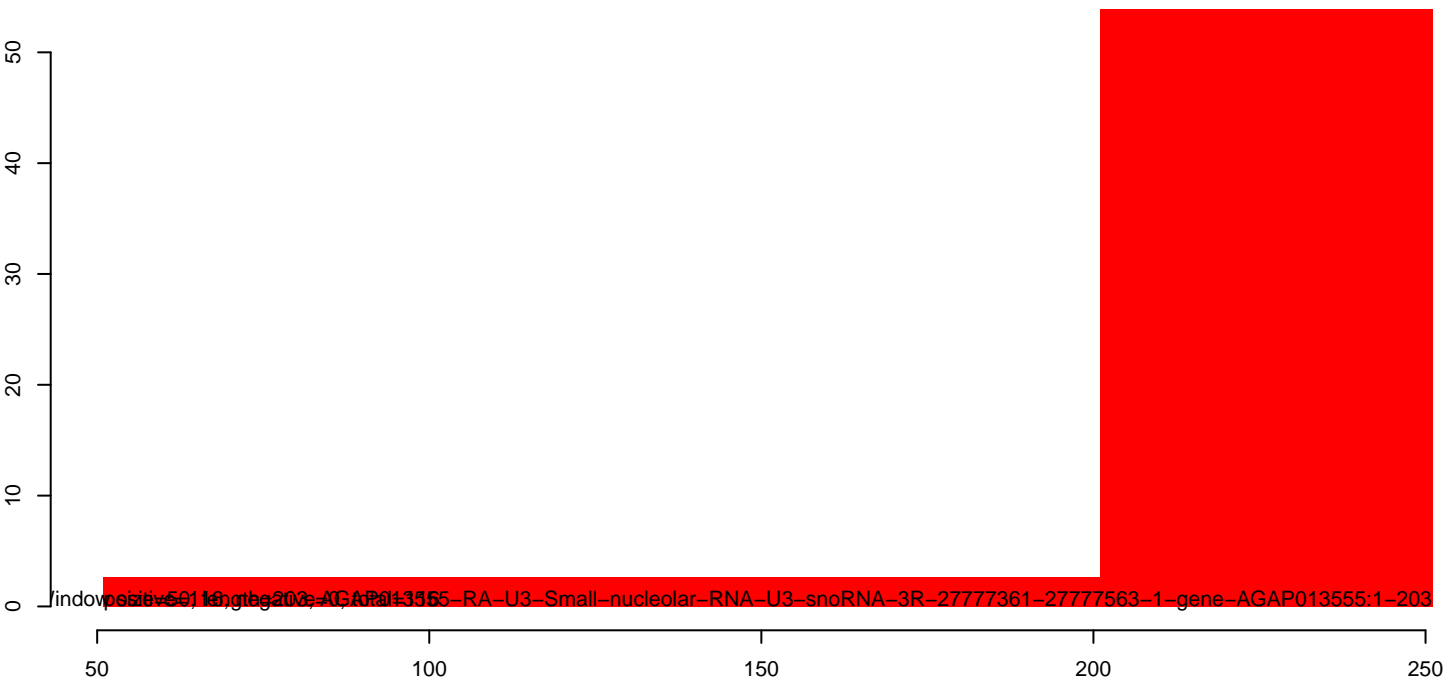
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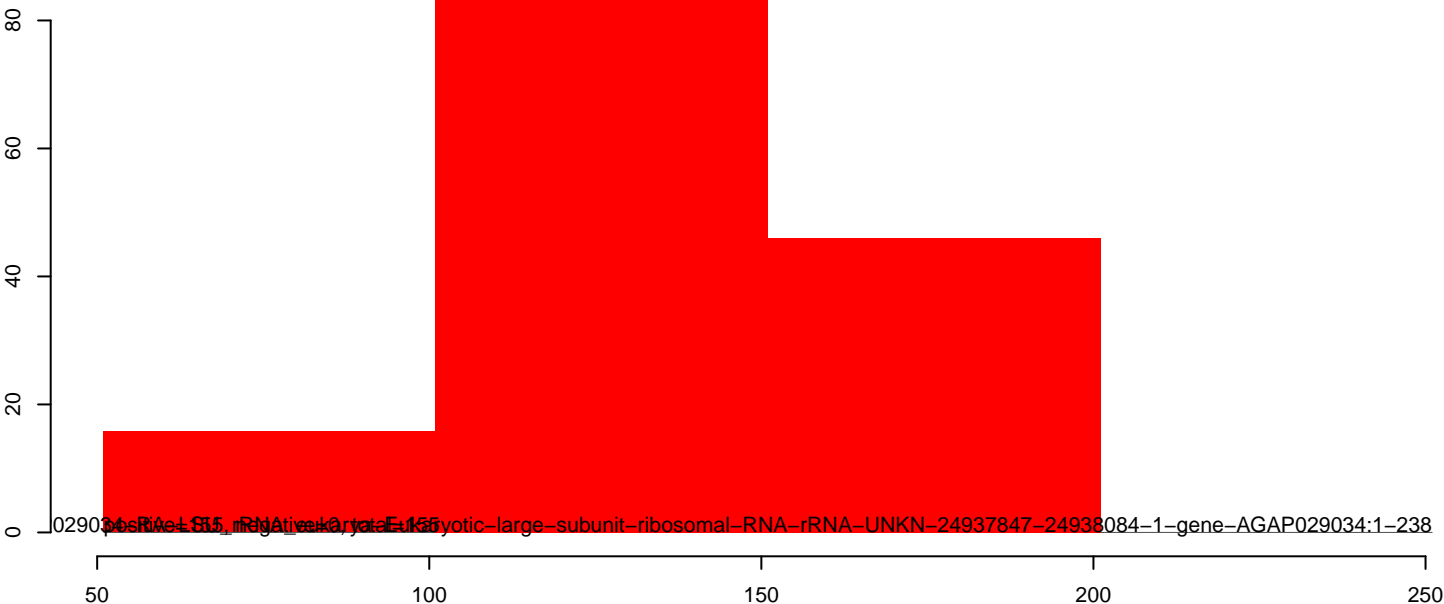
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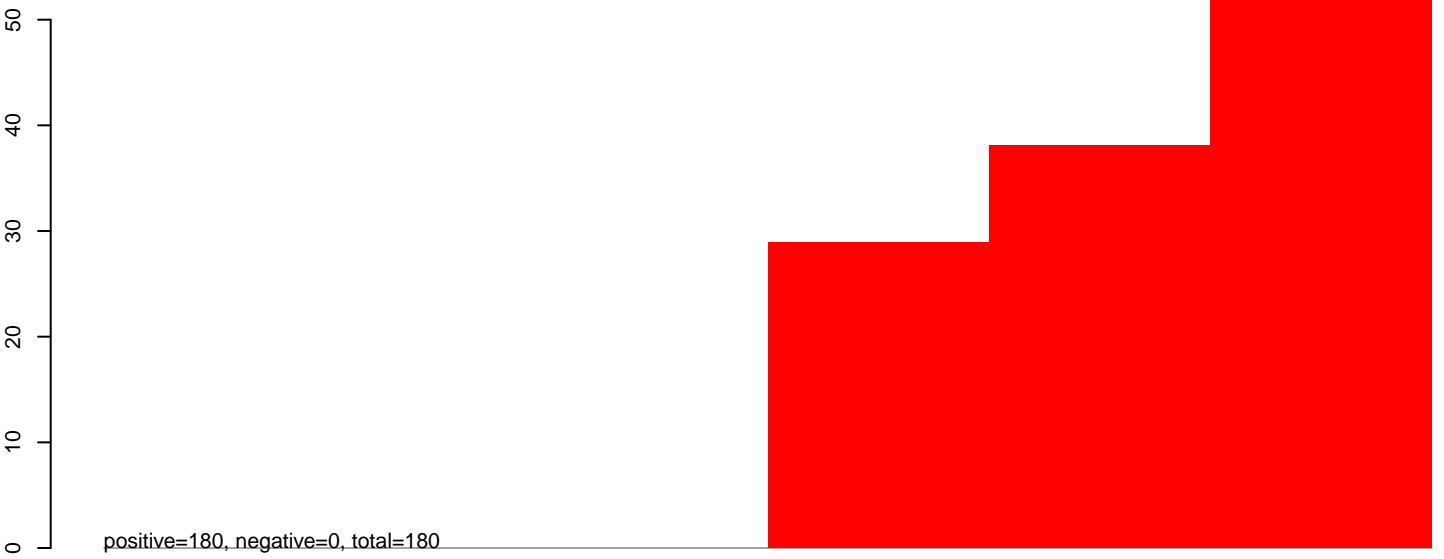


AnGam\_Fcarc\_TN.rep

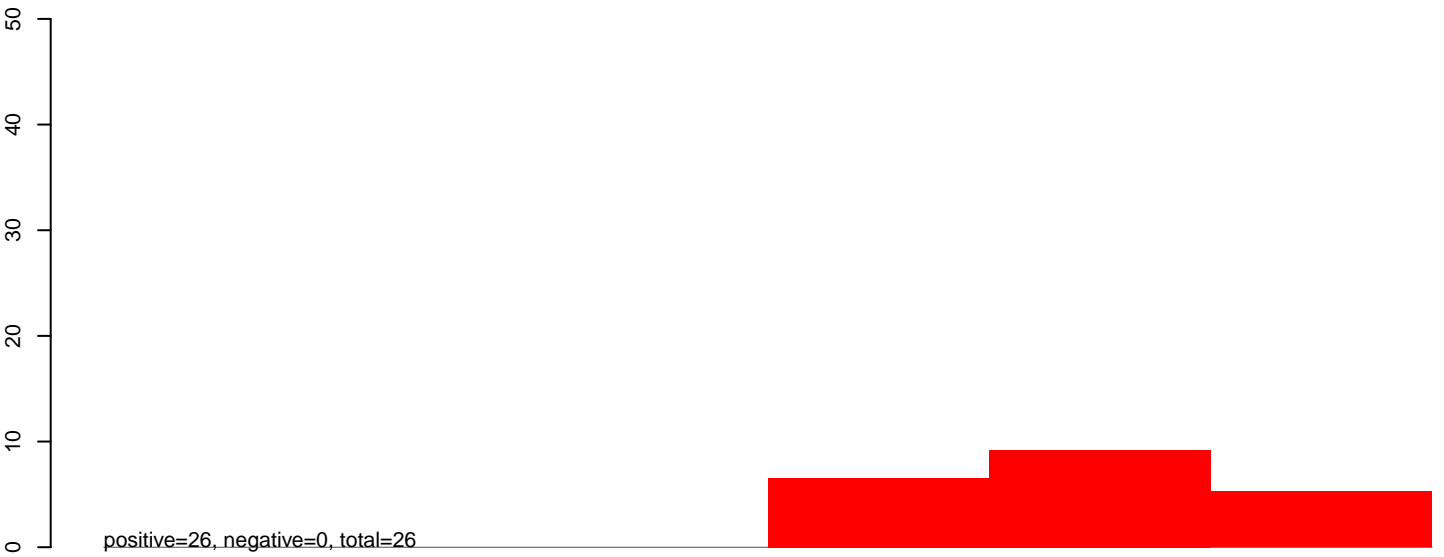


029034:1-238, rRNA-UNKN-24937847-24938084-1-gene-AGAP029034:1-238

AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



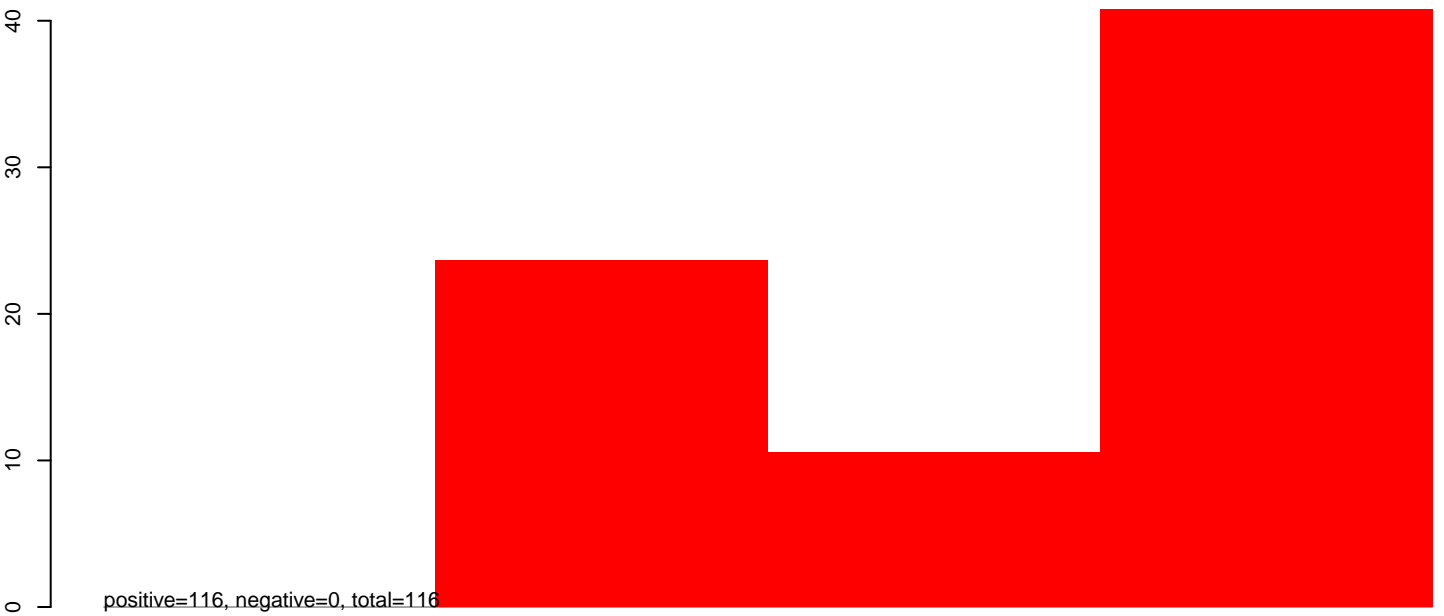
AnGam\_Fcarc\_TN.rep



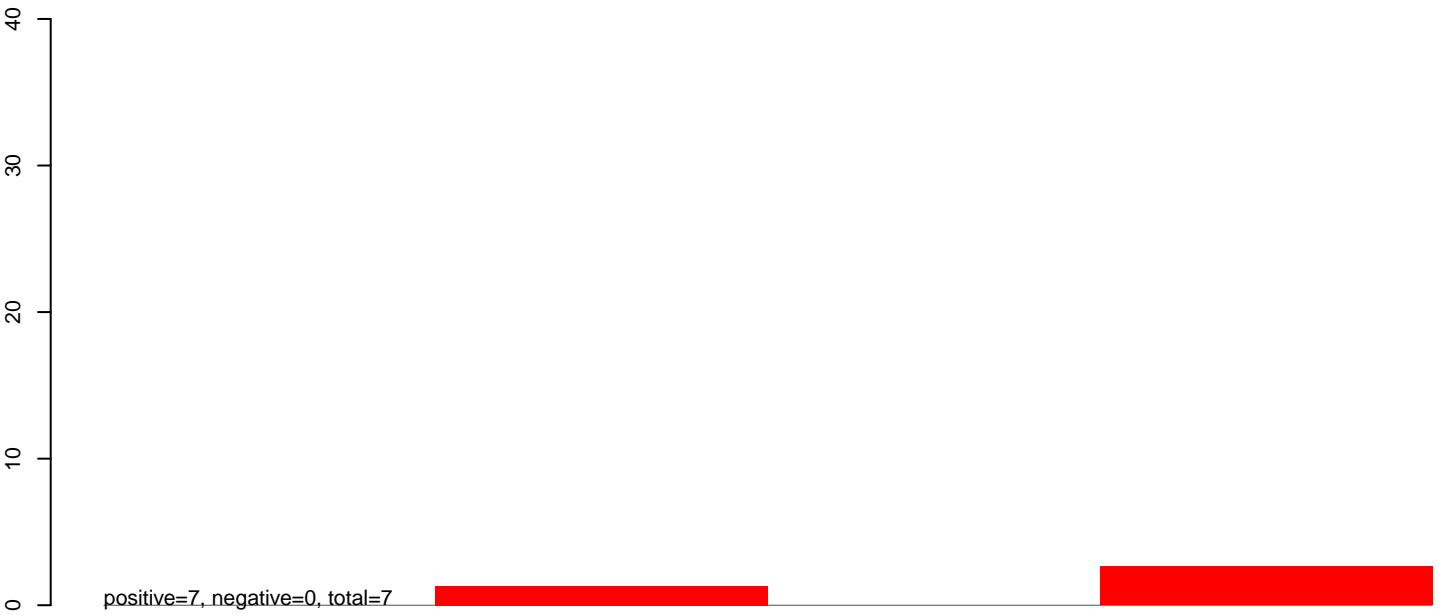
AP029006.1:205-206:R1A=0,karyo=2E6,karyotic-small-subunit-ribosomal-rRNA-rRNA-3R-21742293-21742609-1-gene-AGAP029006:1-317

50 100 150 200 250 300 350

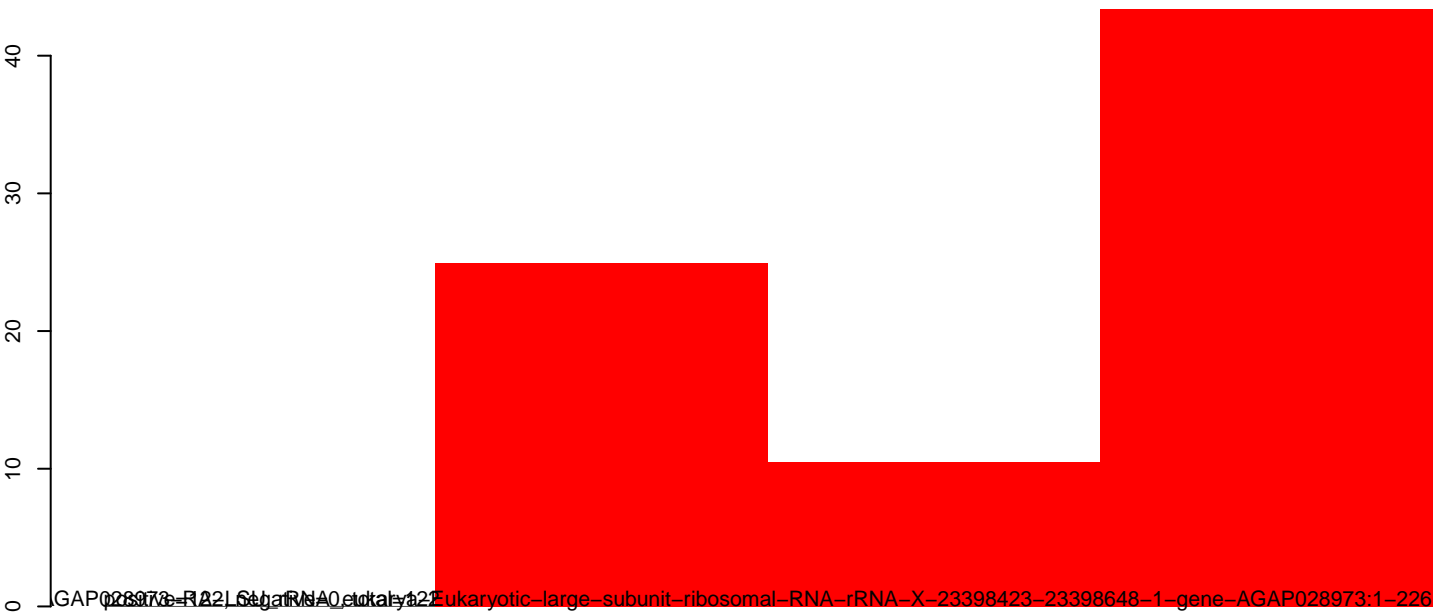
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



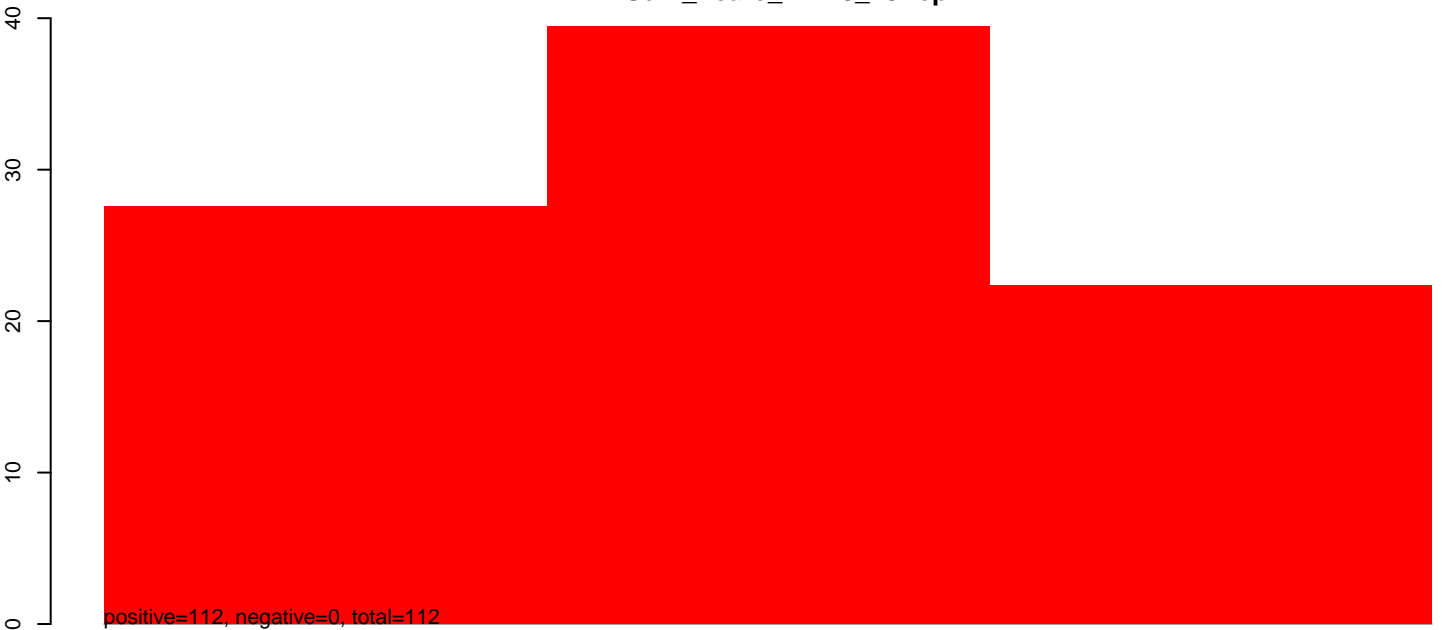
AnGam\_Fcarc\_TN.rep



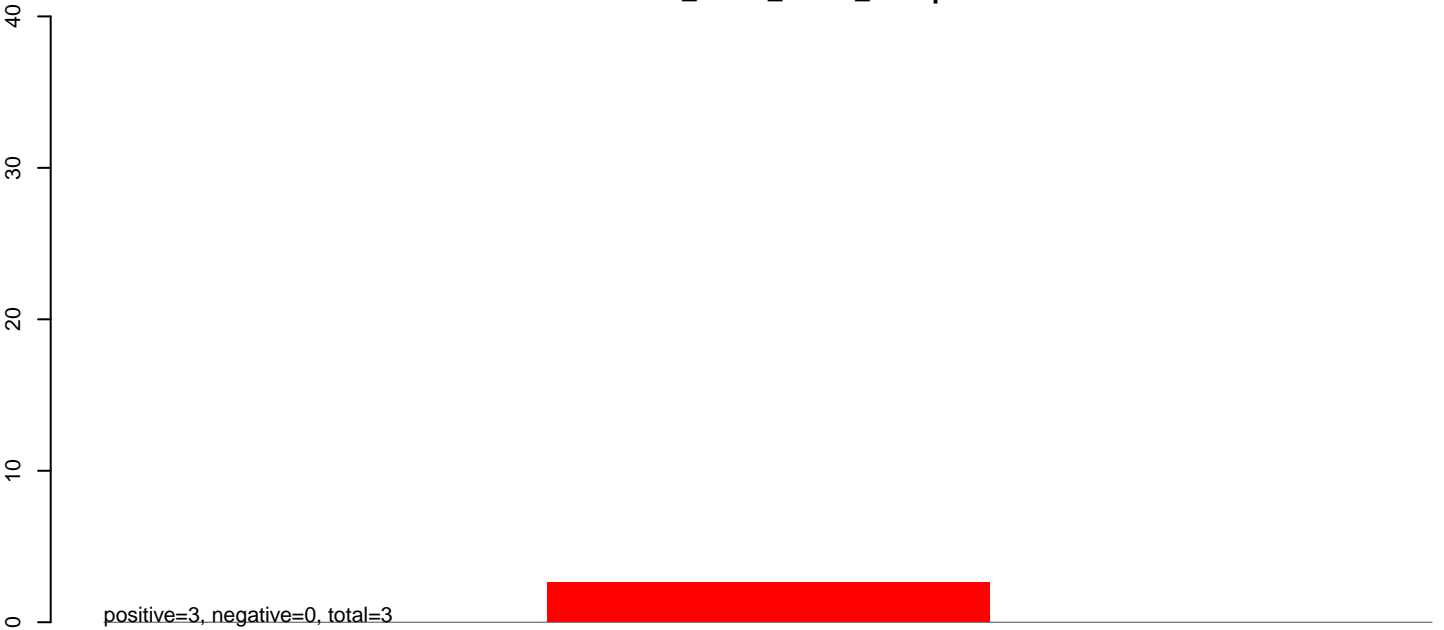
GAP028973-R12.6.gene.RNA\_eukaryotic-large-subunit-ribosomal-RNA-rRNA-X-23398423-23398648-1-gene-AGAP028973:1-226

50 100 150 200 250

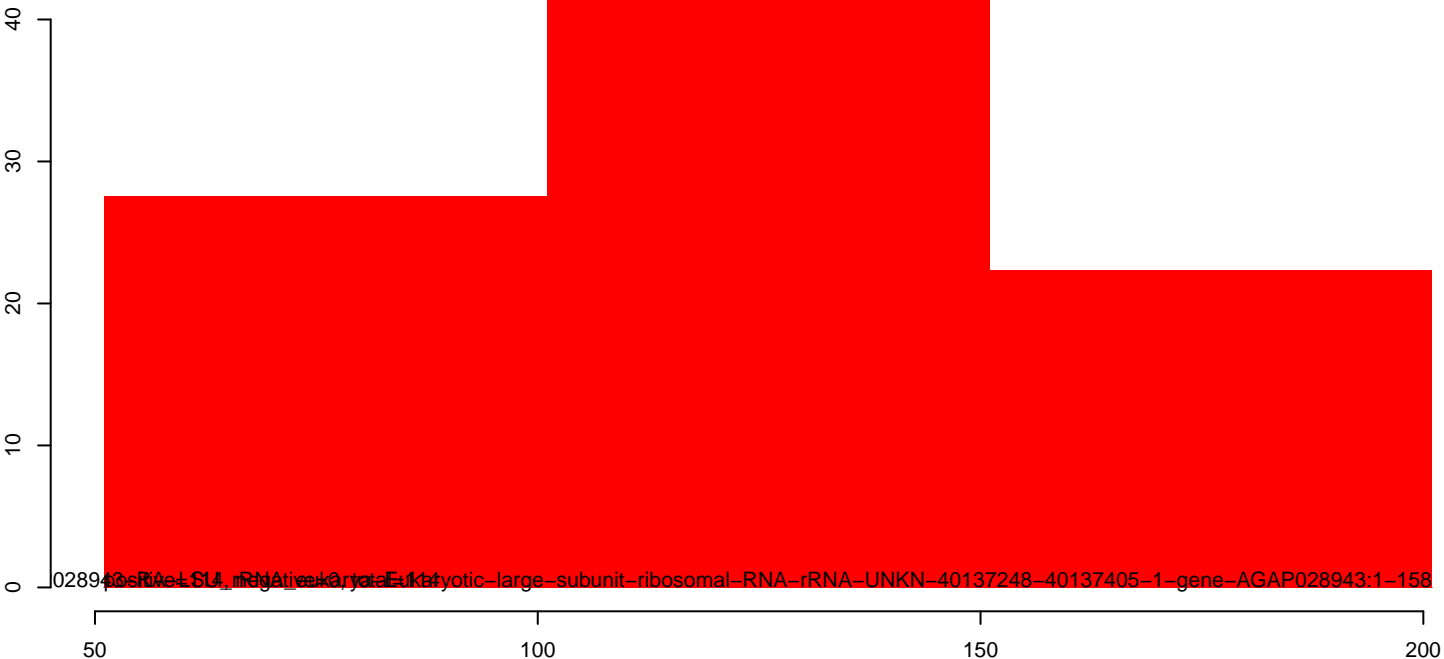
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep

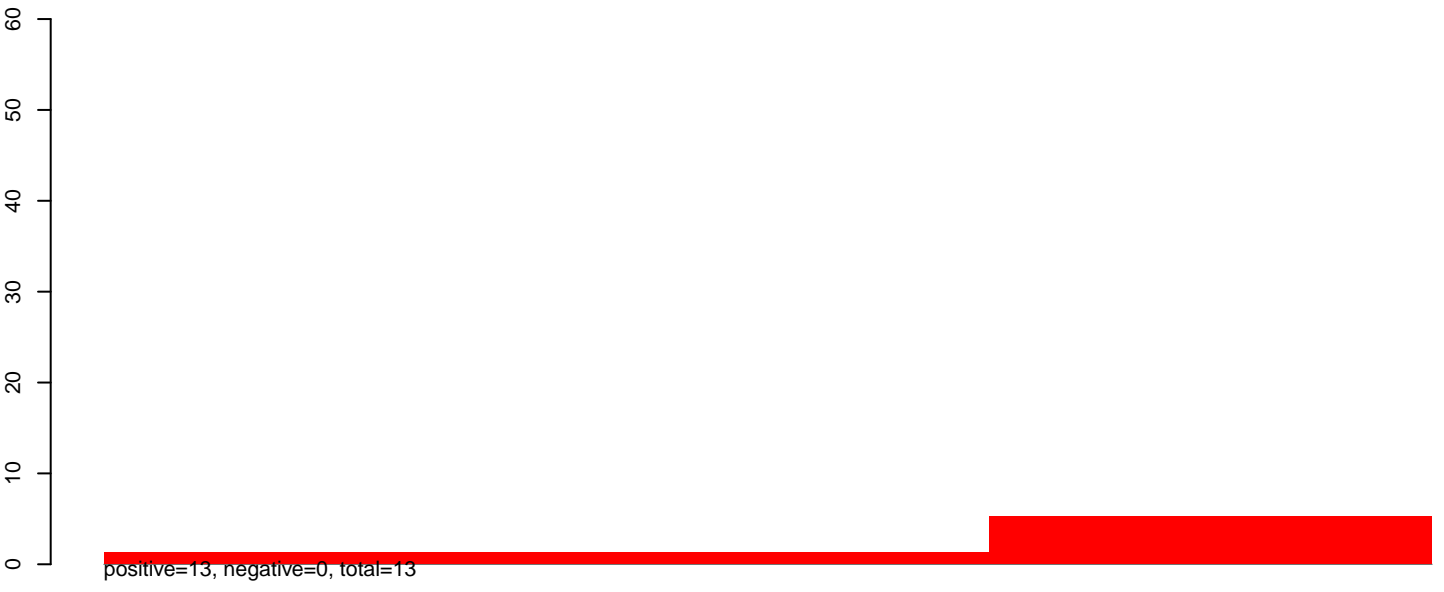




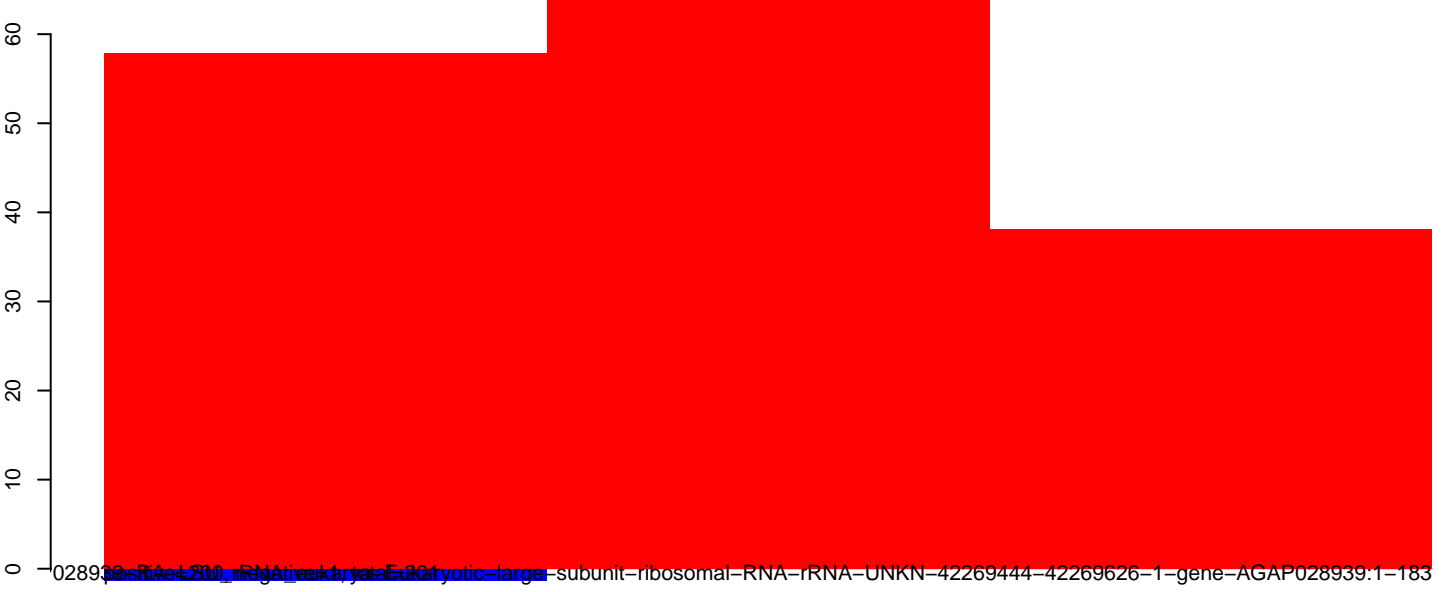
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



028939:Rice.800:Rijalva.kryta.F01:cytic-large-subunit-ribosomal-RNA-rRNA-UNKN-42269444-42269626-1-gene-AGAP028939:1-183

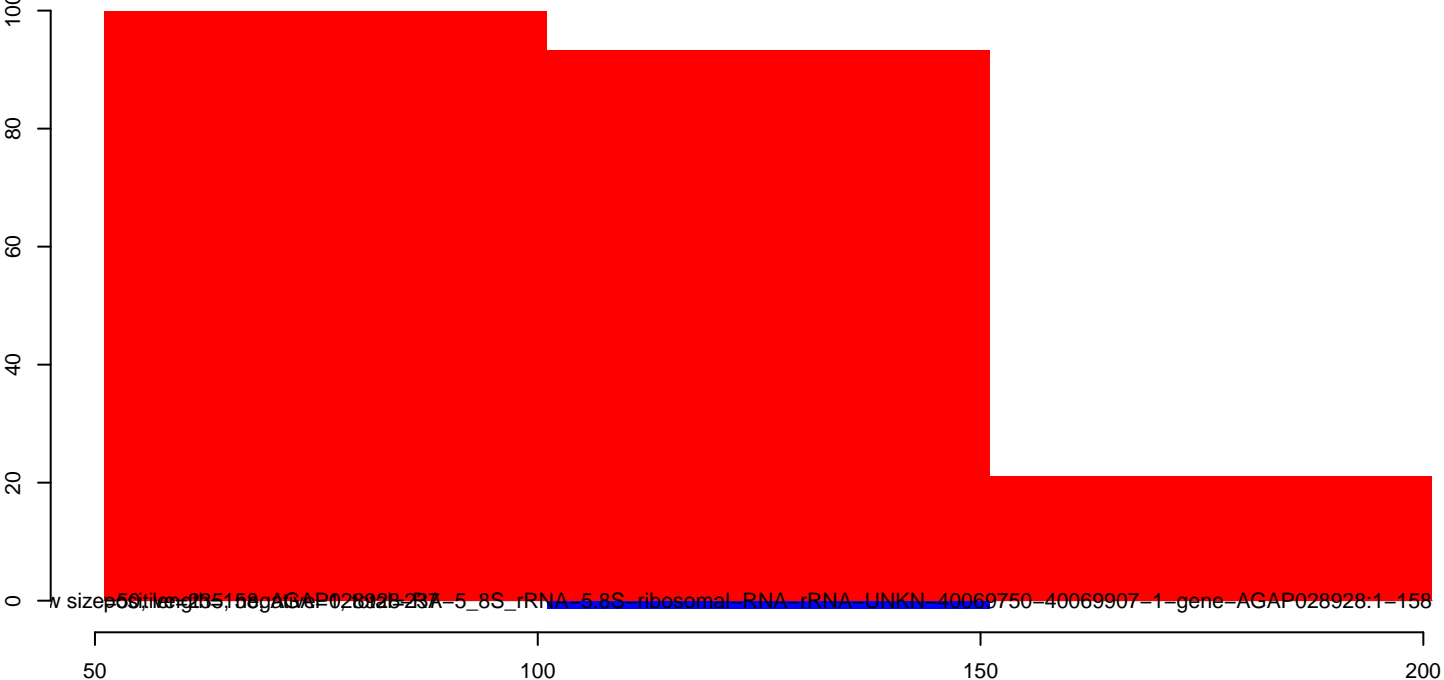
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



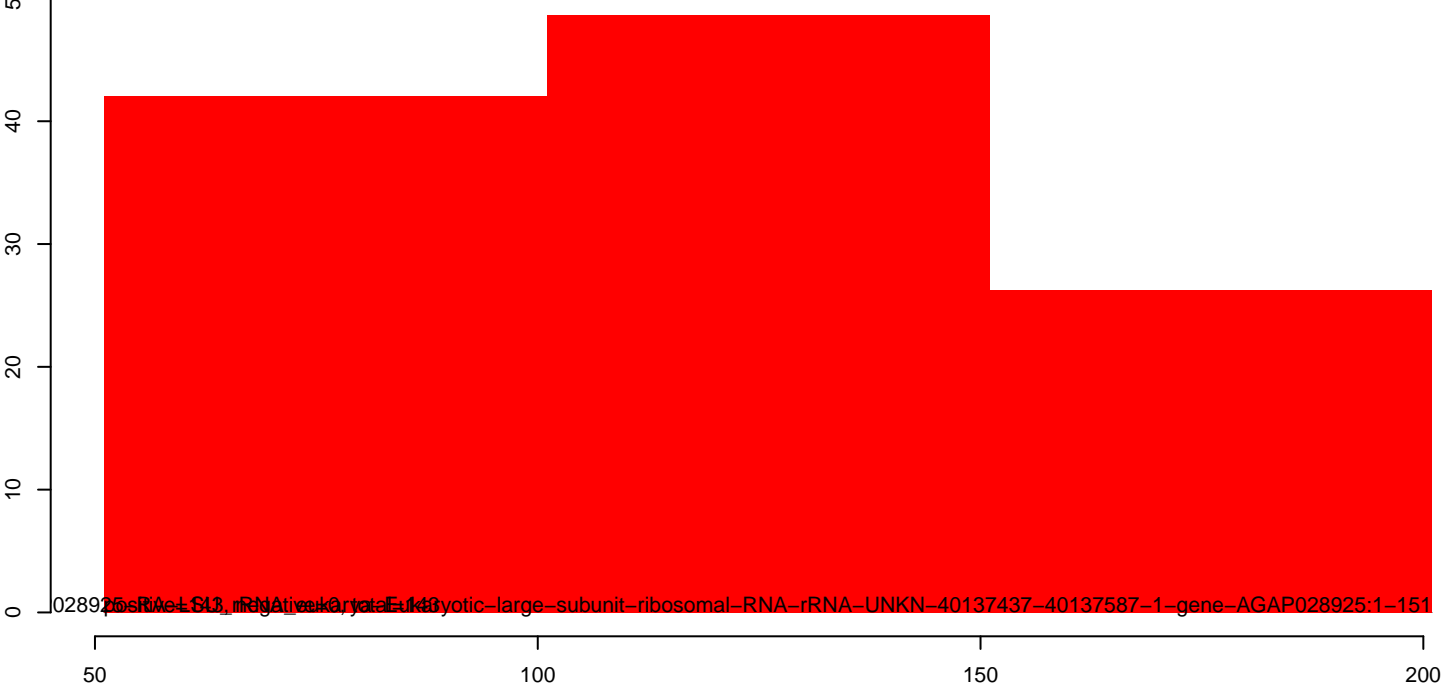
AnGam\_Fcarc\_TN.18\_23.rep



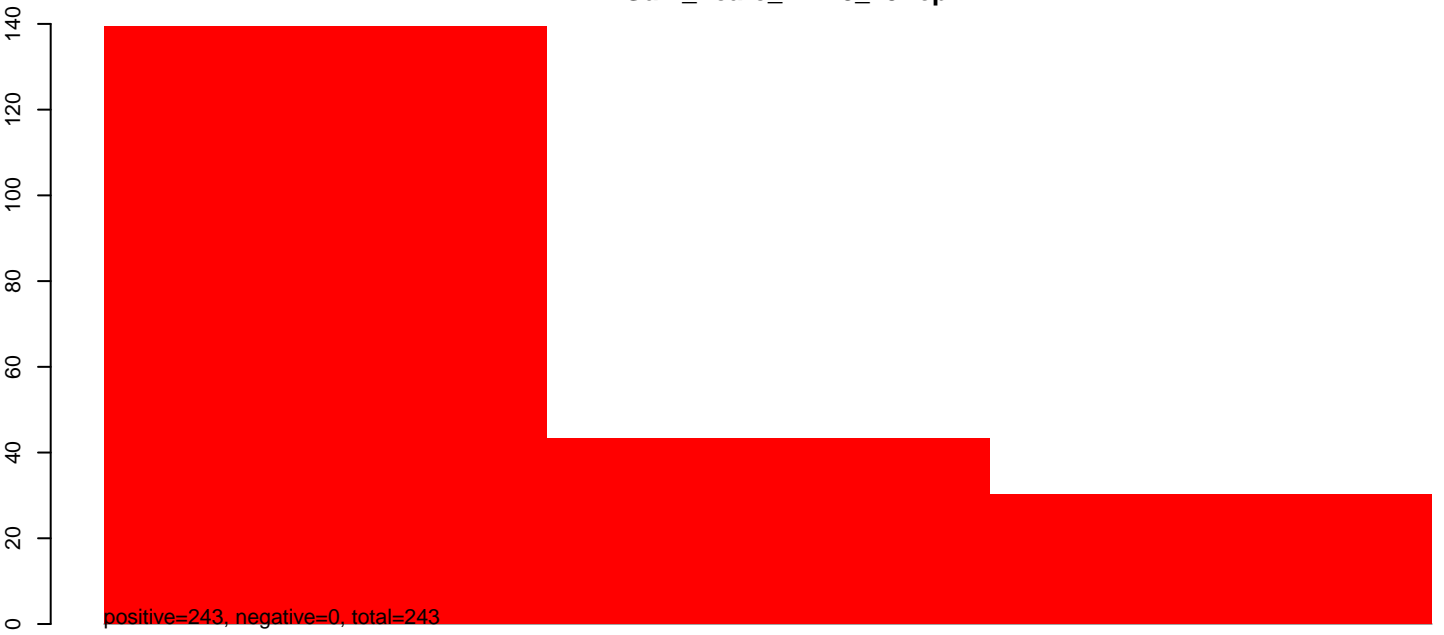
AnGam\_Fcarc\_TN.24\_35.rep



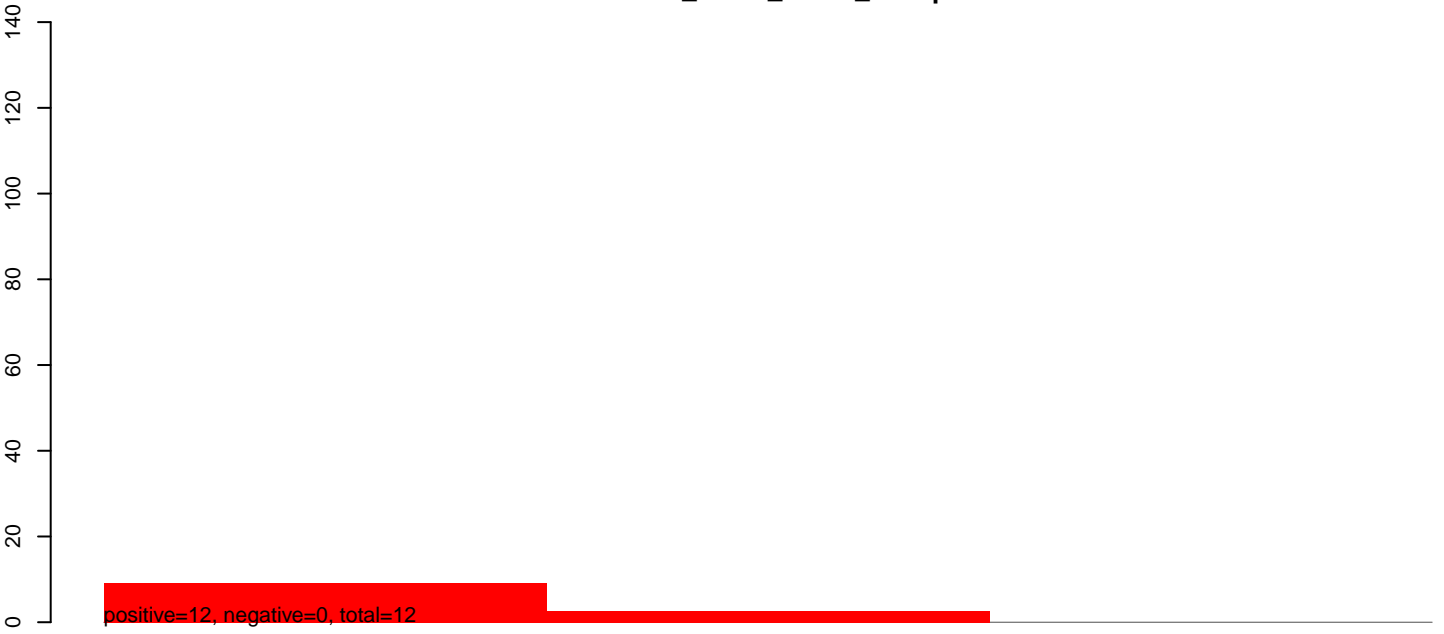
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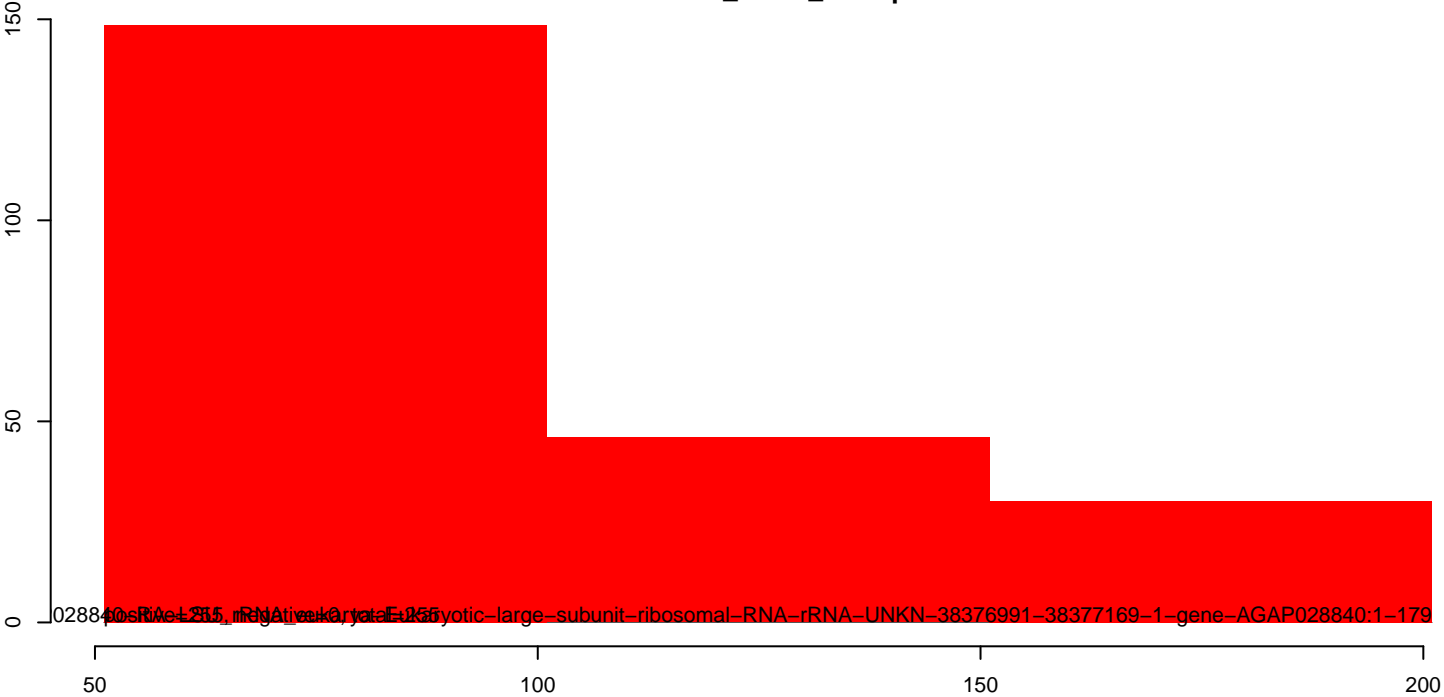
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AnGam\_Fcarc\_TN.24\_35.rep



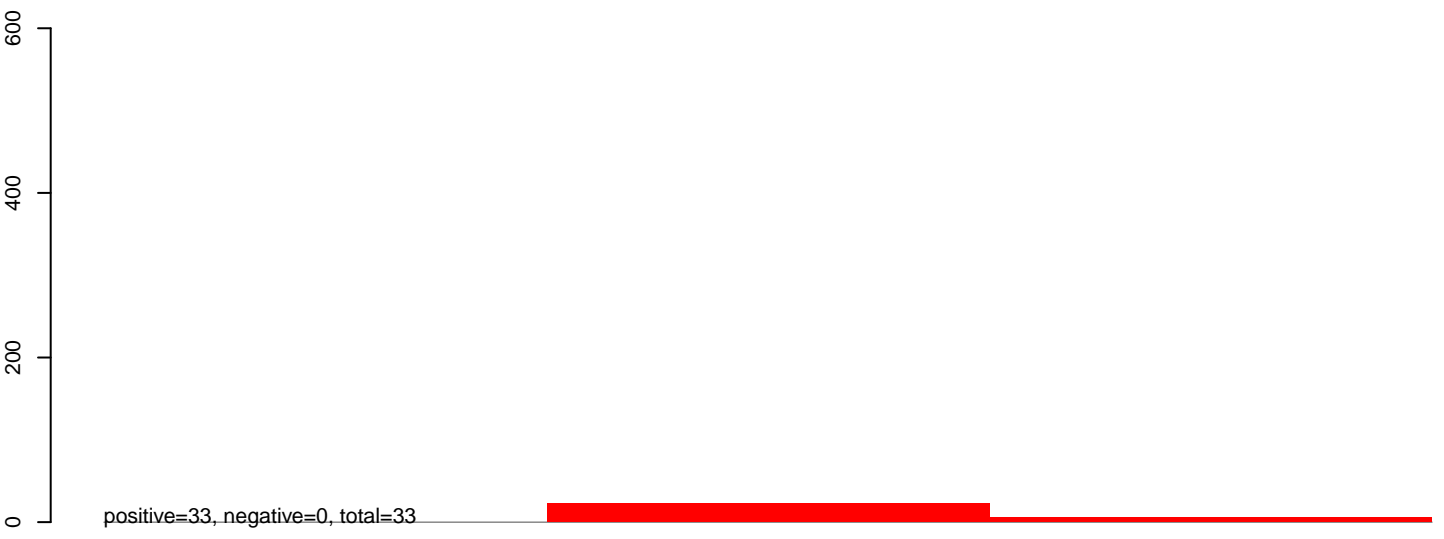
AnGam\_Fcarc\_TN.rep



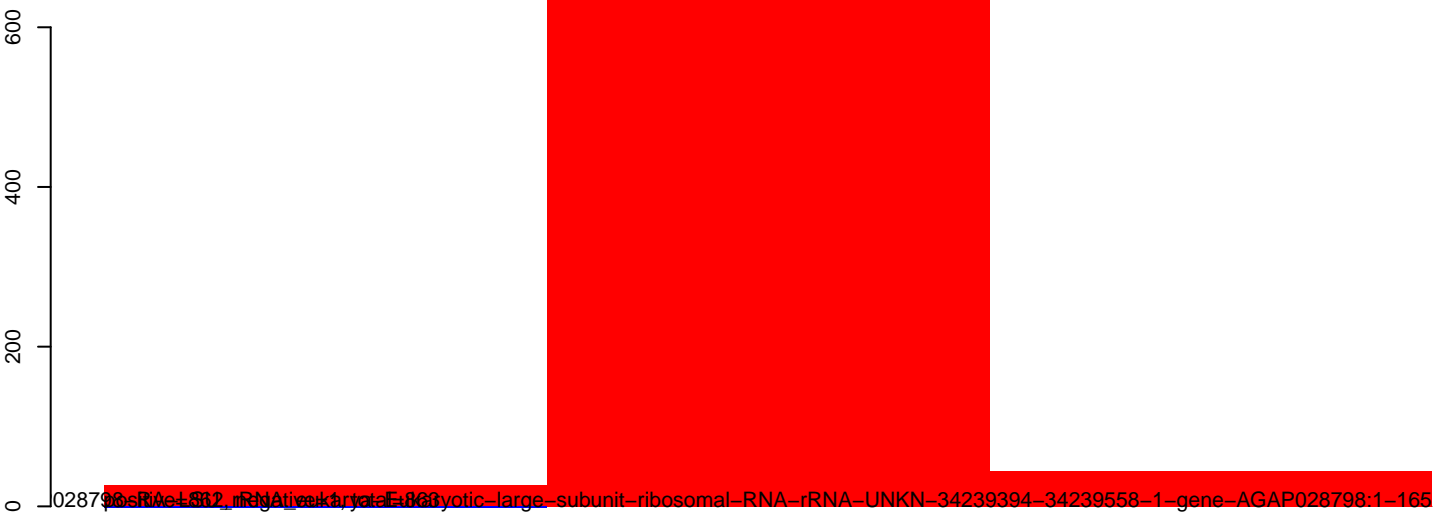
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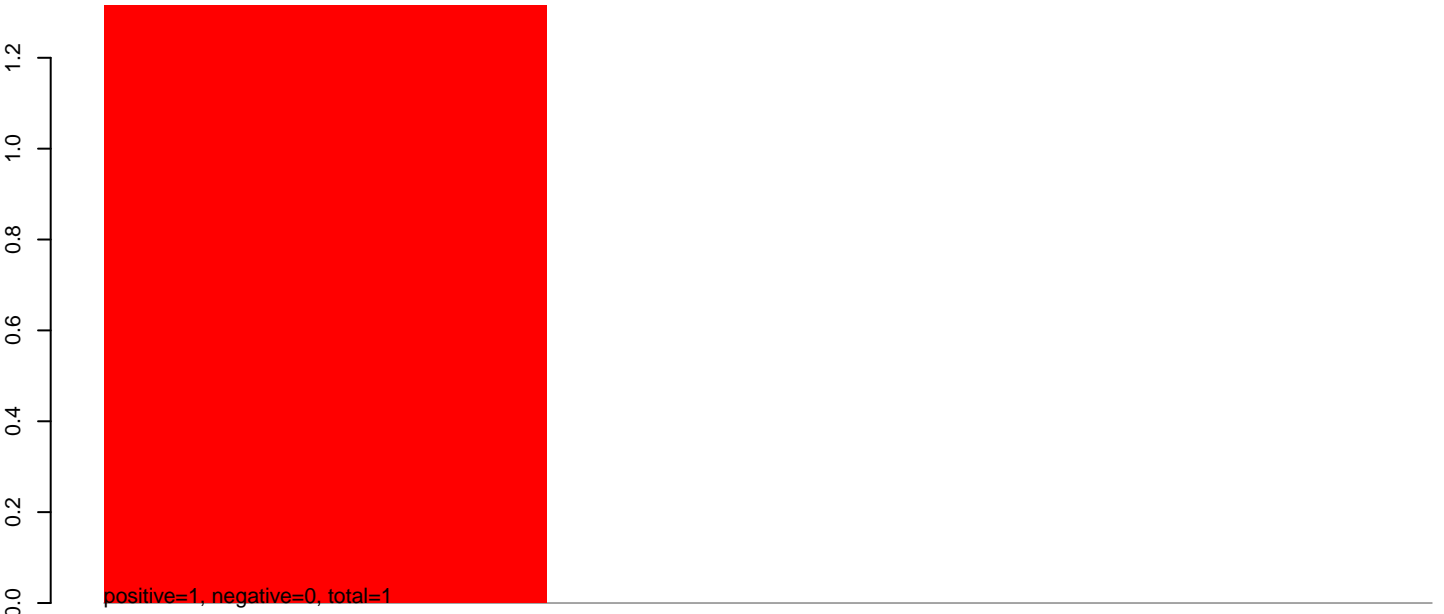
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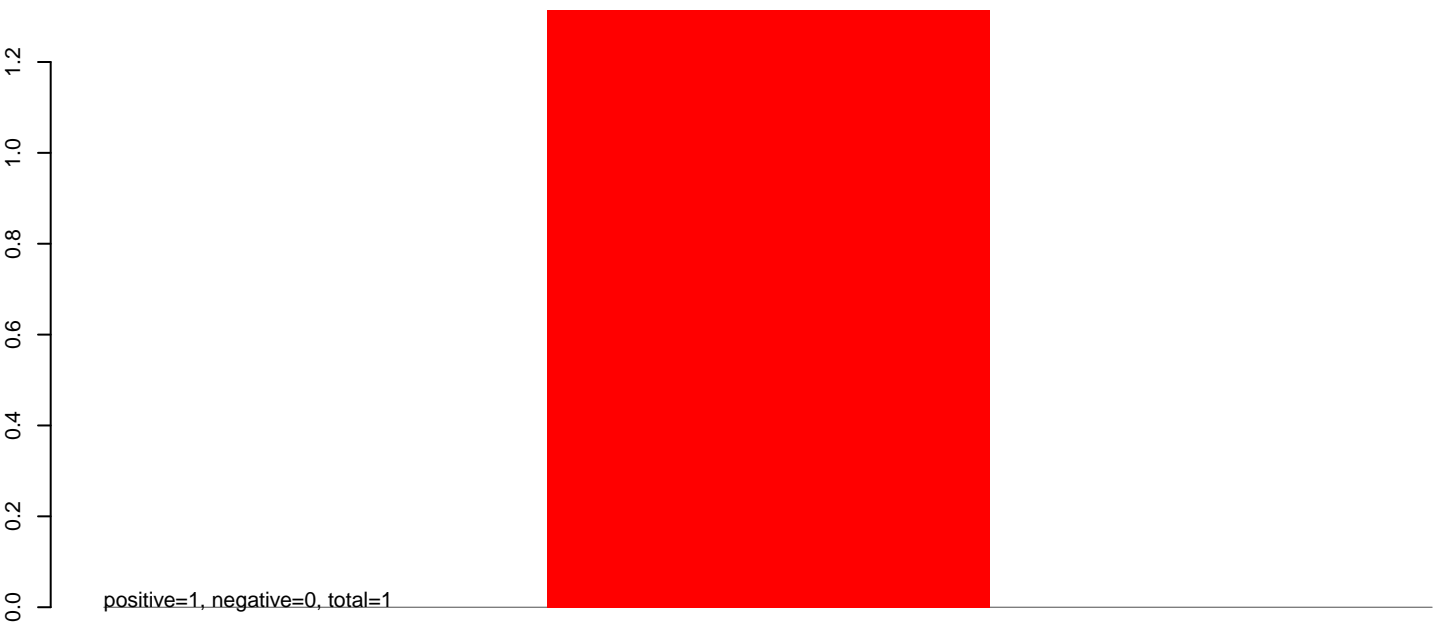
AnGam\_Fcarc\_TN.rep



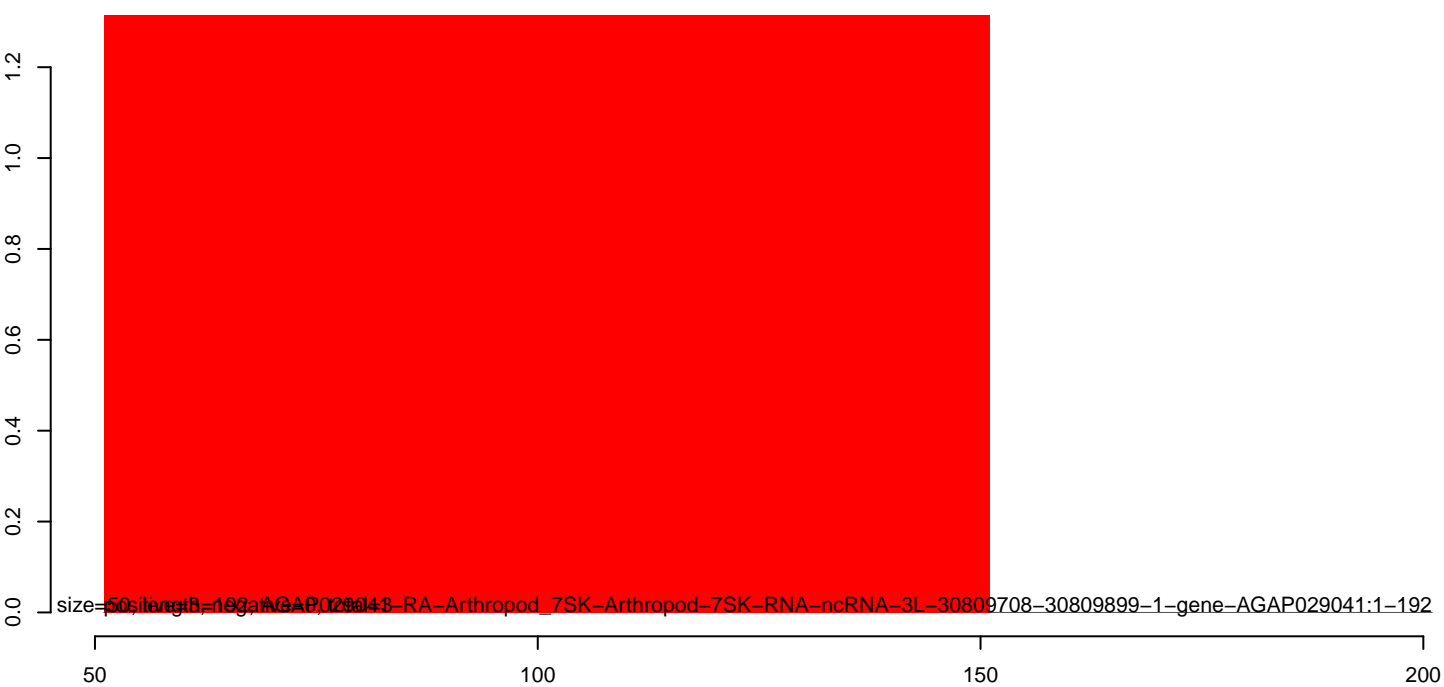
AnGam\_Fcarc\_TN.18\_23.rep



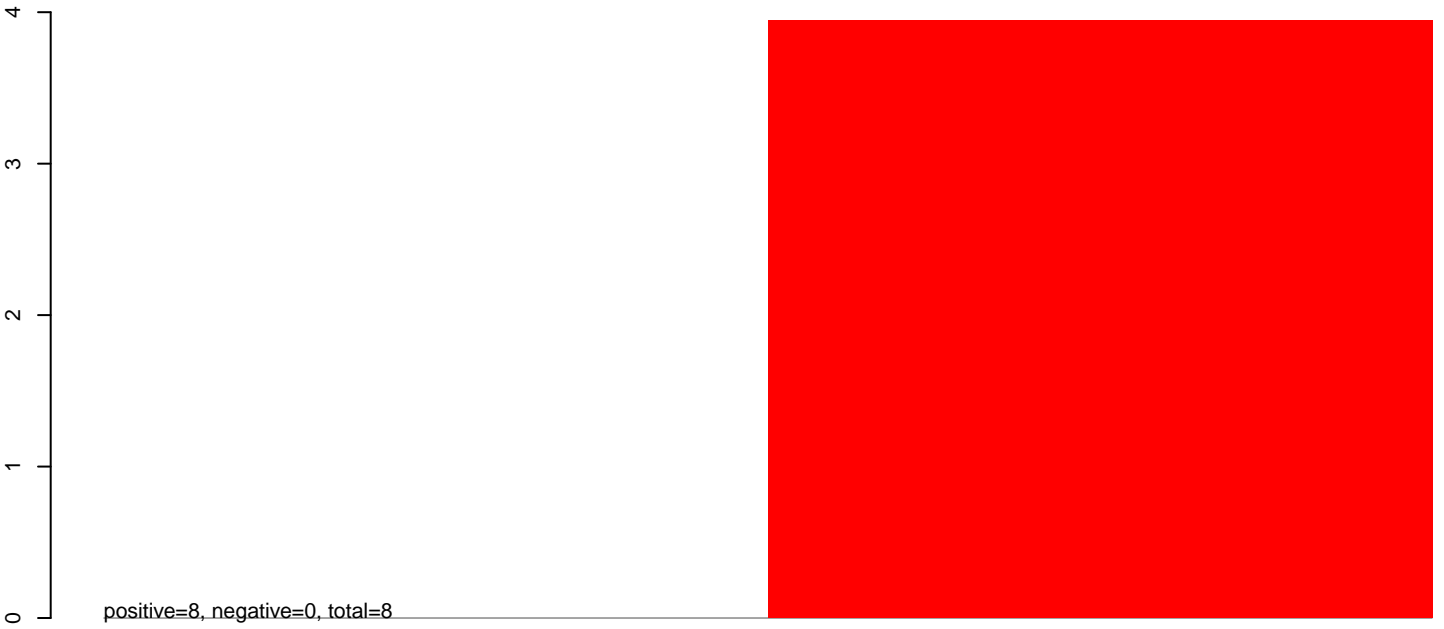
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AnGam\_Fcarc\_TN.rep



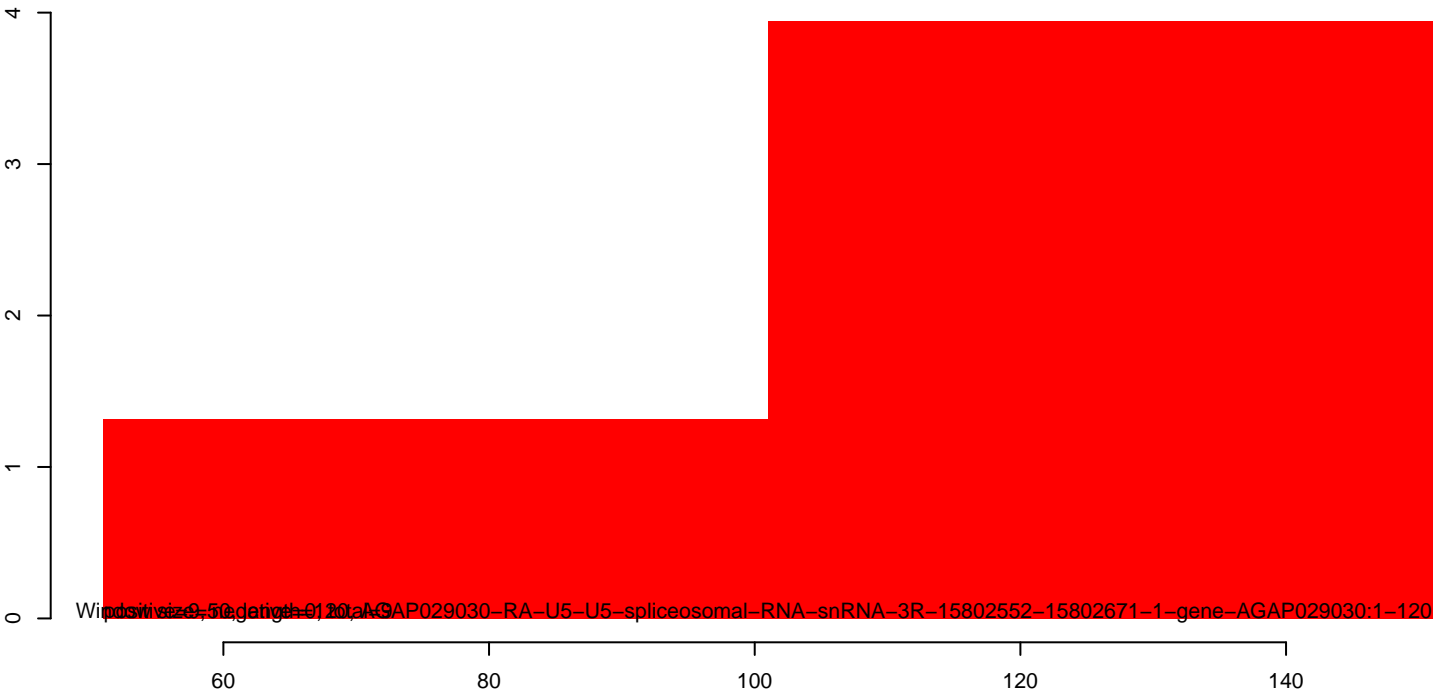
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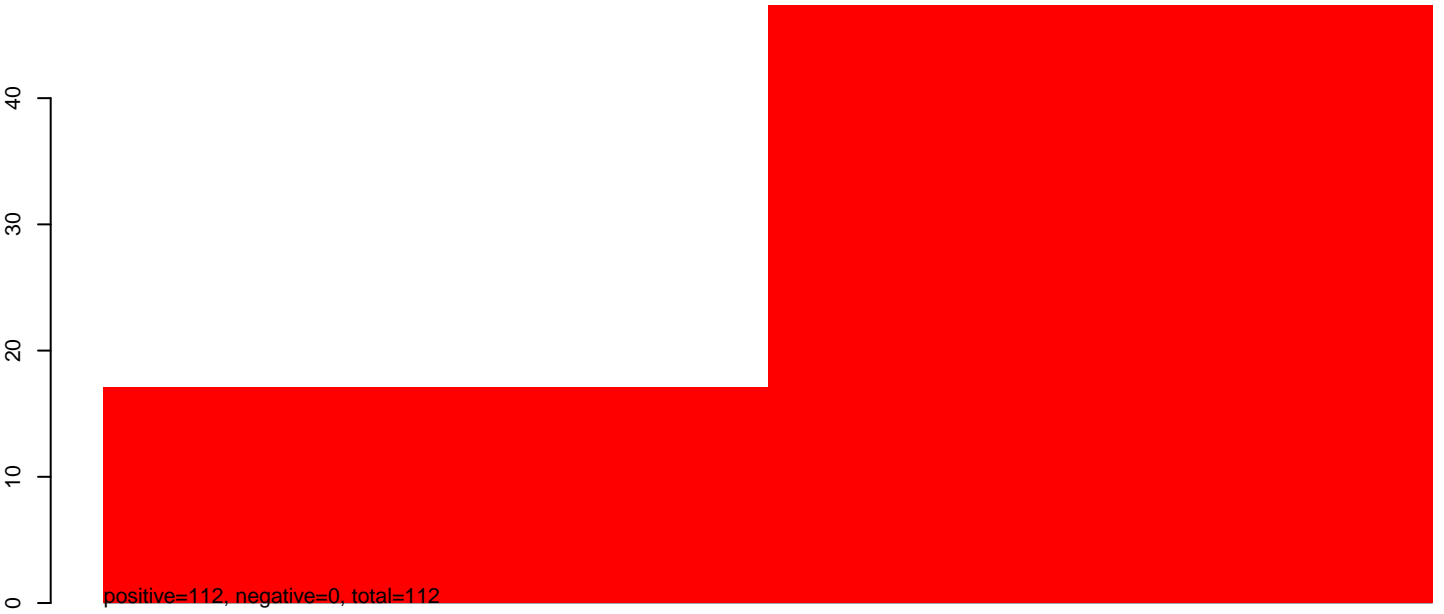
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AnGam\_Fcarc\_TN.rep



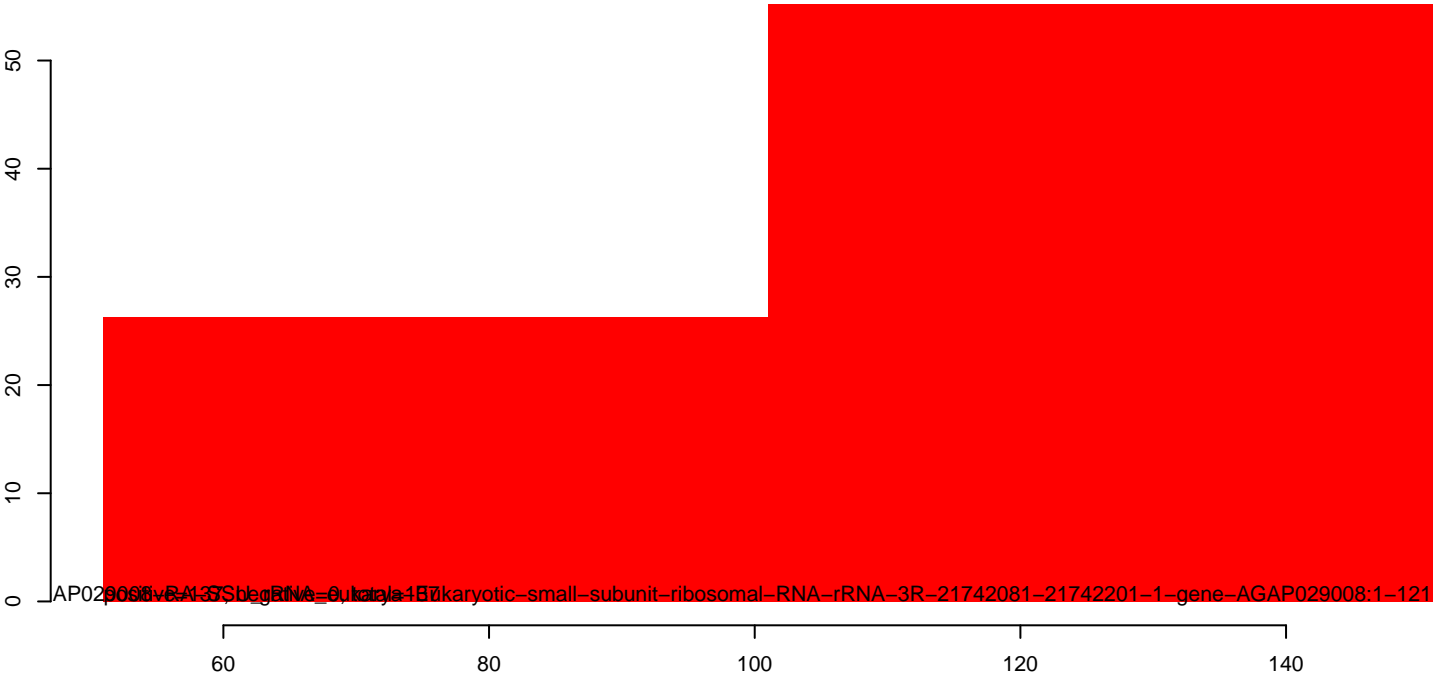
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AnGam\_Fcarc\_TN.24\_35.rep

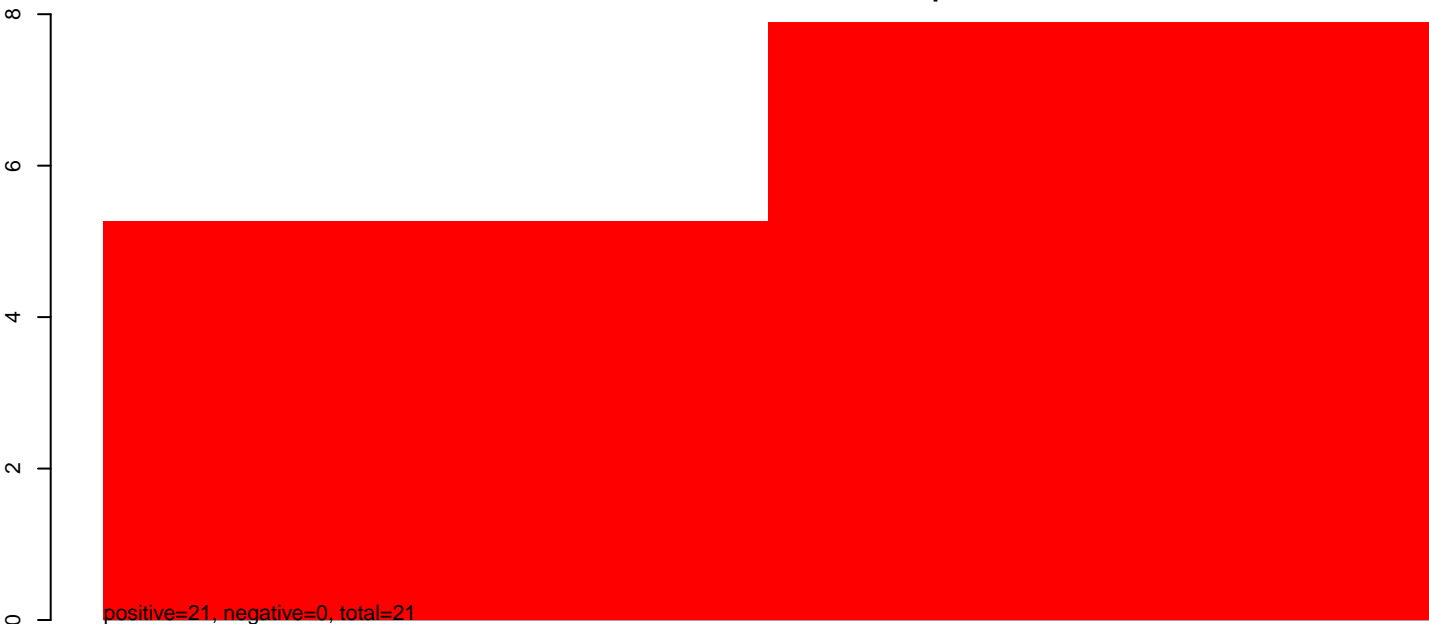


AnGam\_Fcarc\_TN.rep

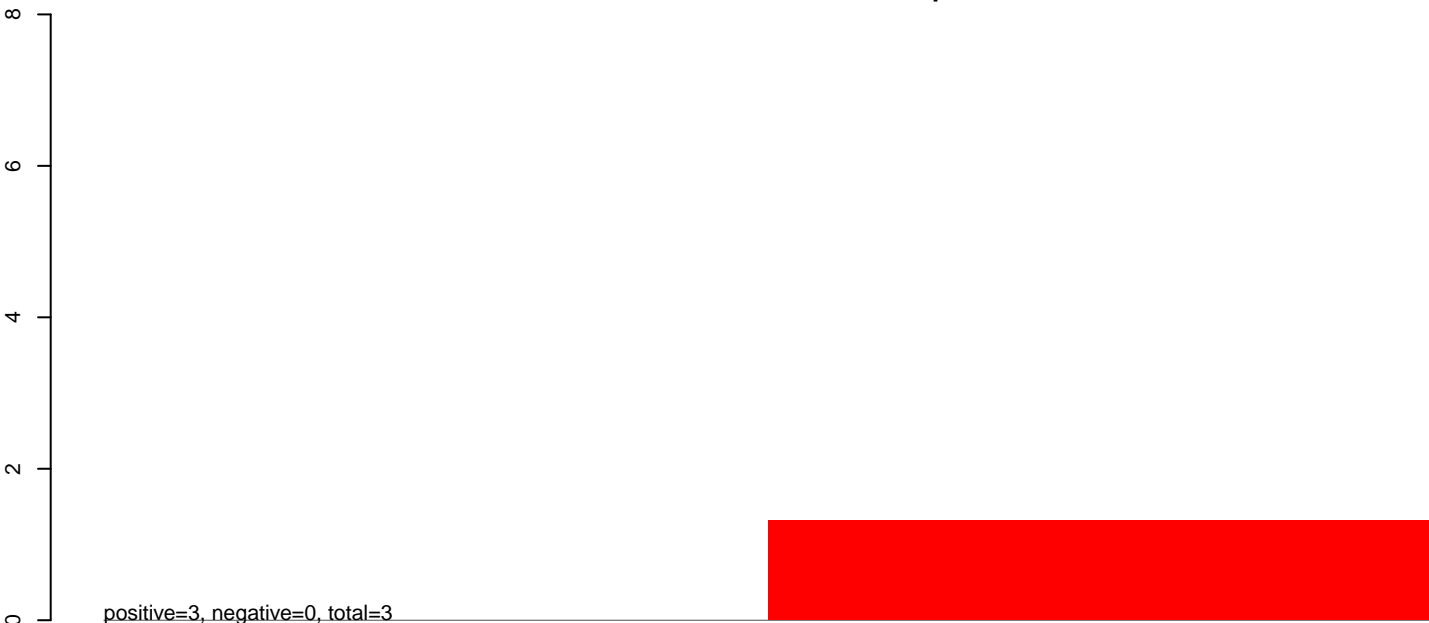




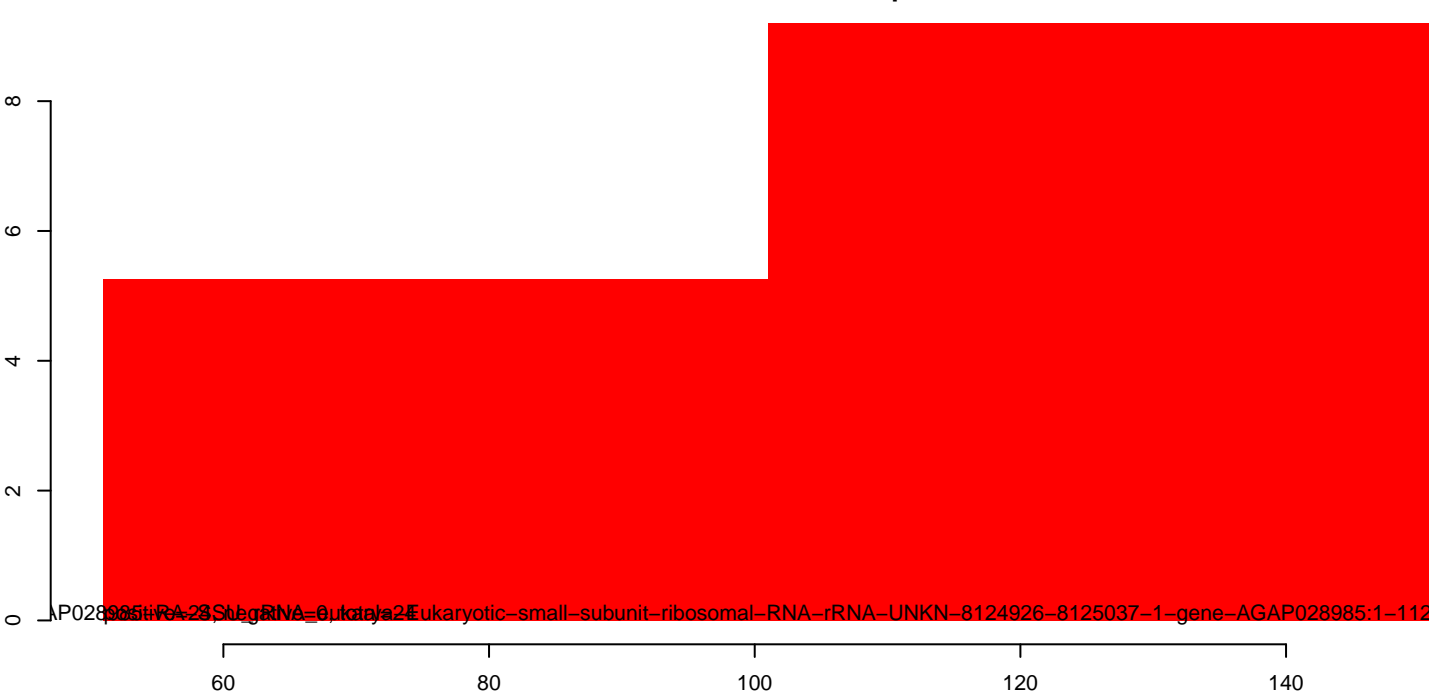
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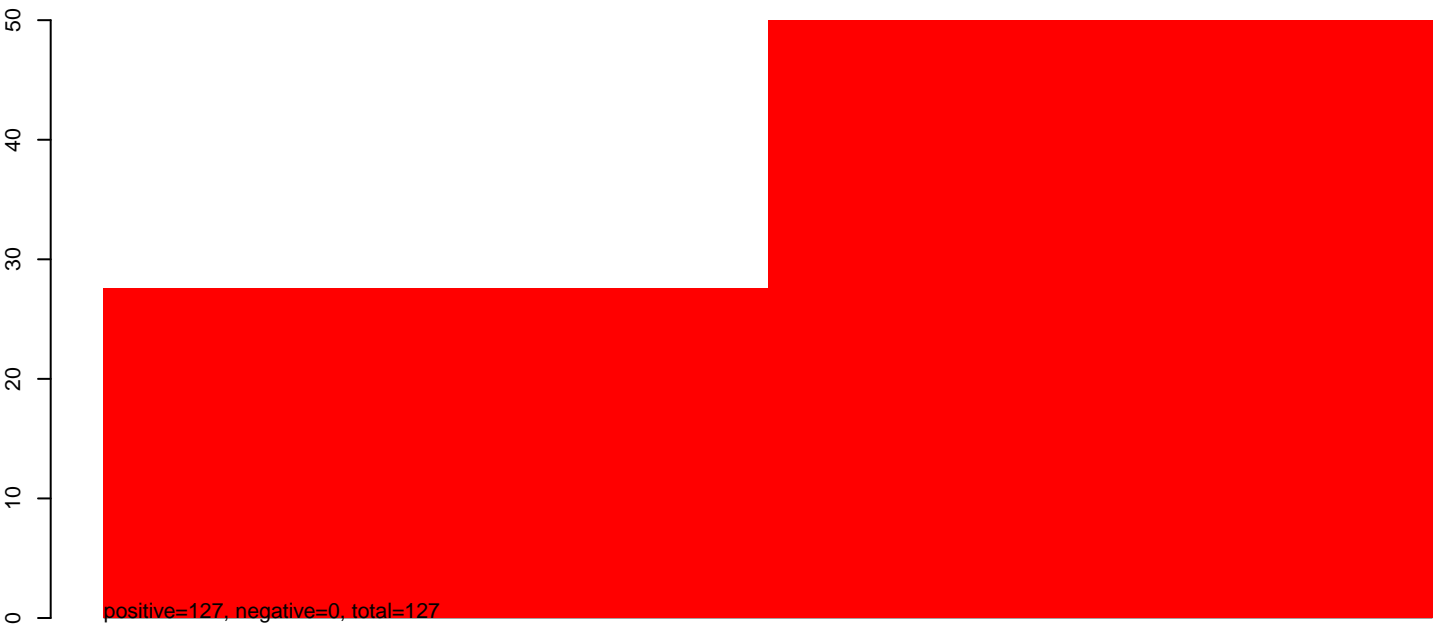
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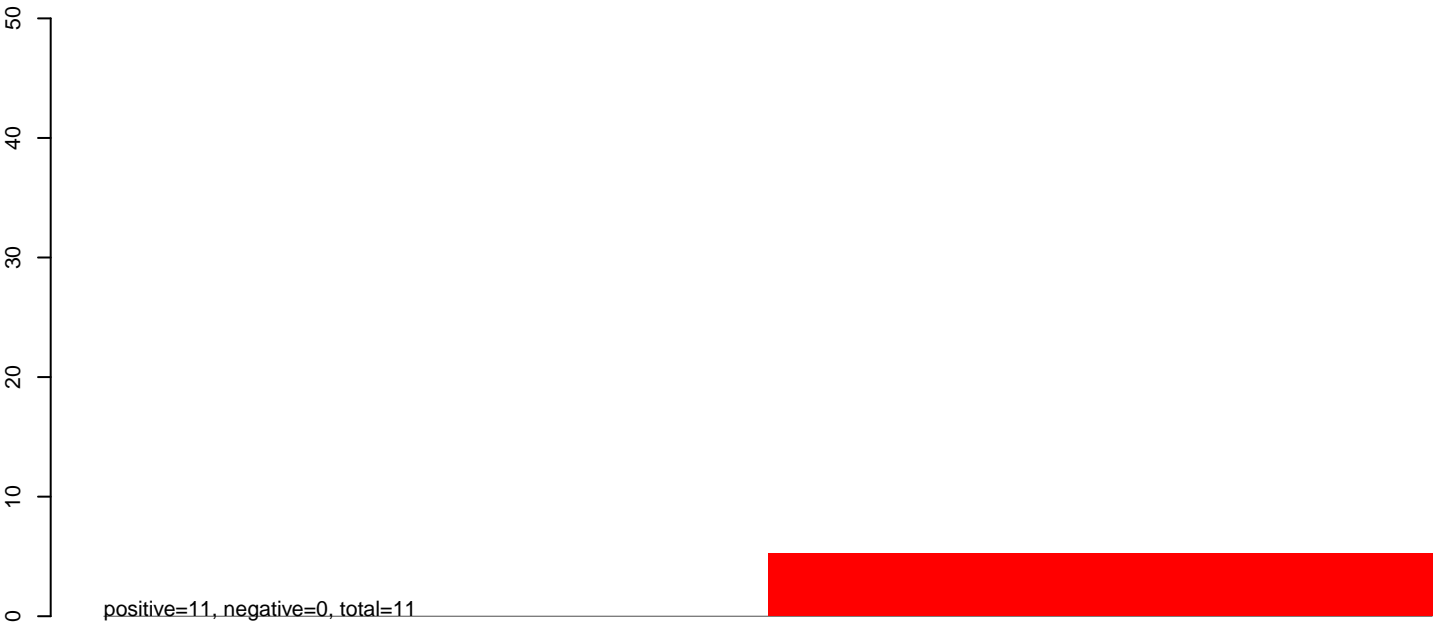
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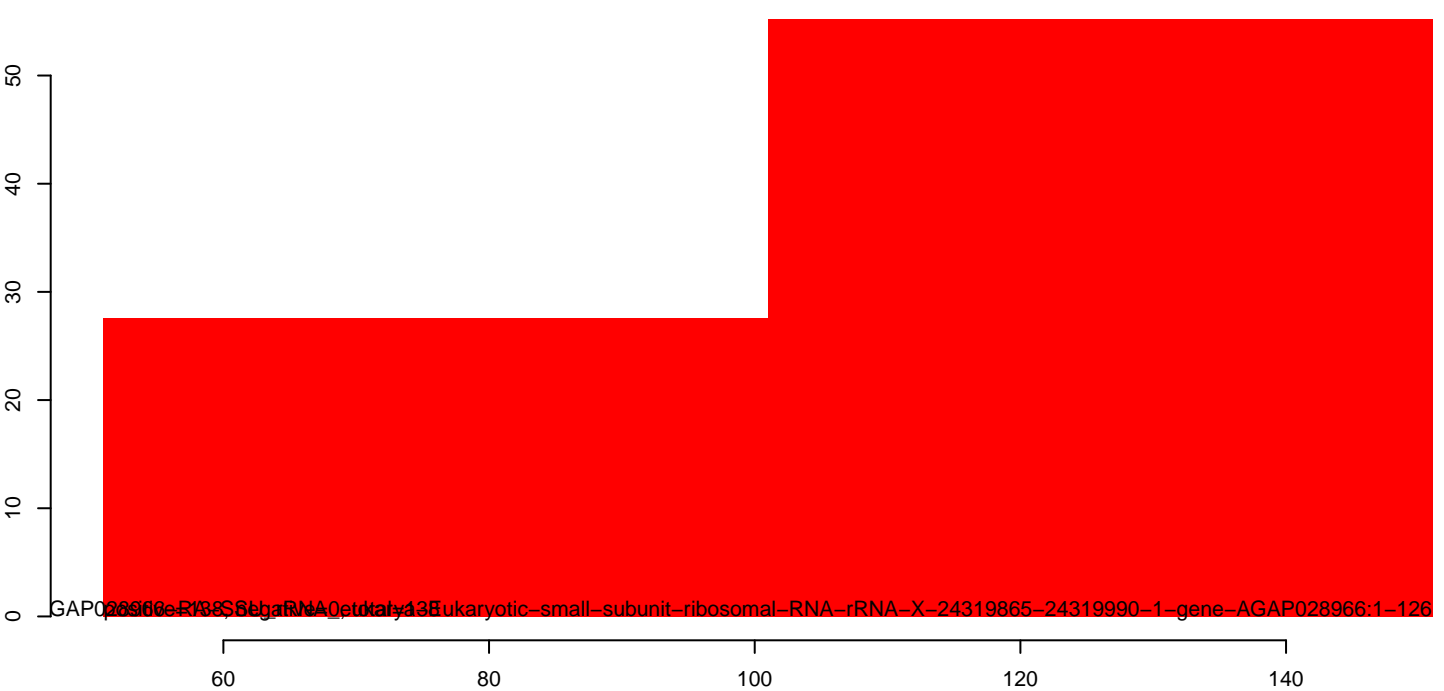
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



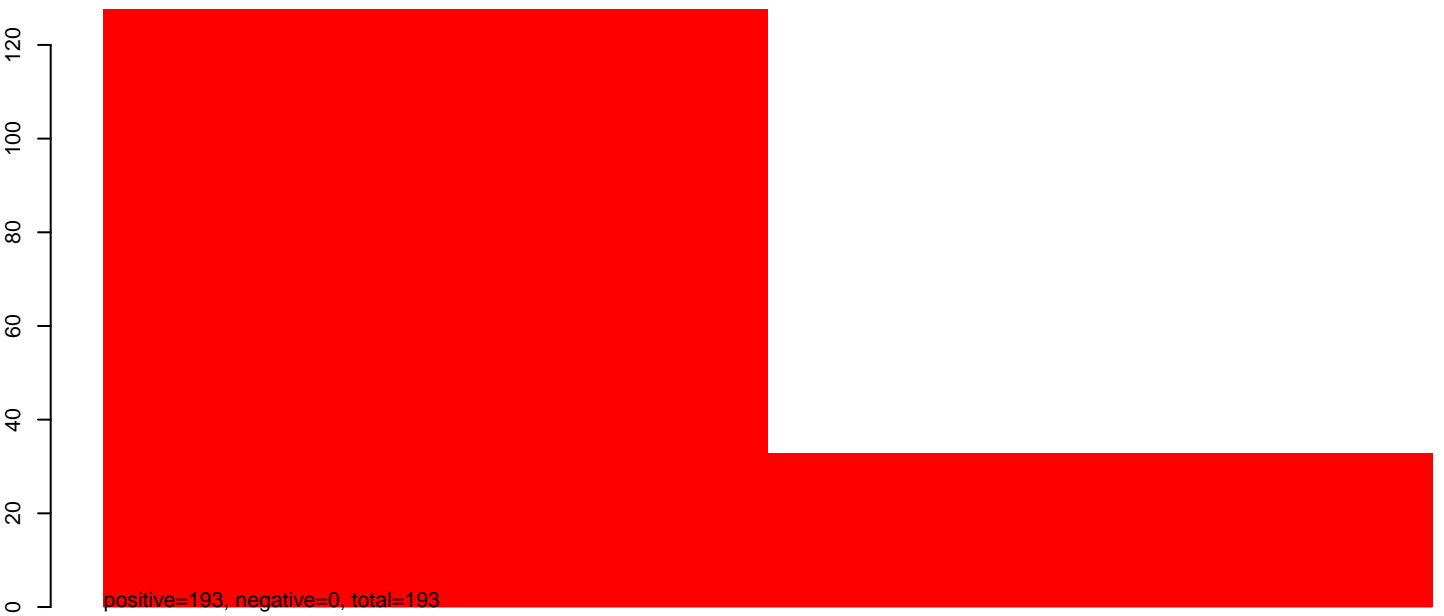
AnGam\_Fcarc\_TN.rep



028929:SSU\_rRNA=0, total=70 karyotic-small-subunit-ribosomal-RNA-rRNA-UNKN-40628342-40628476-1-gene-AGAP028929:1-135

60 80 100 120 140

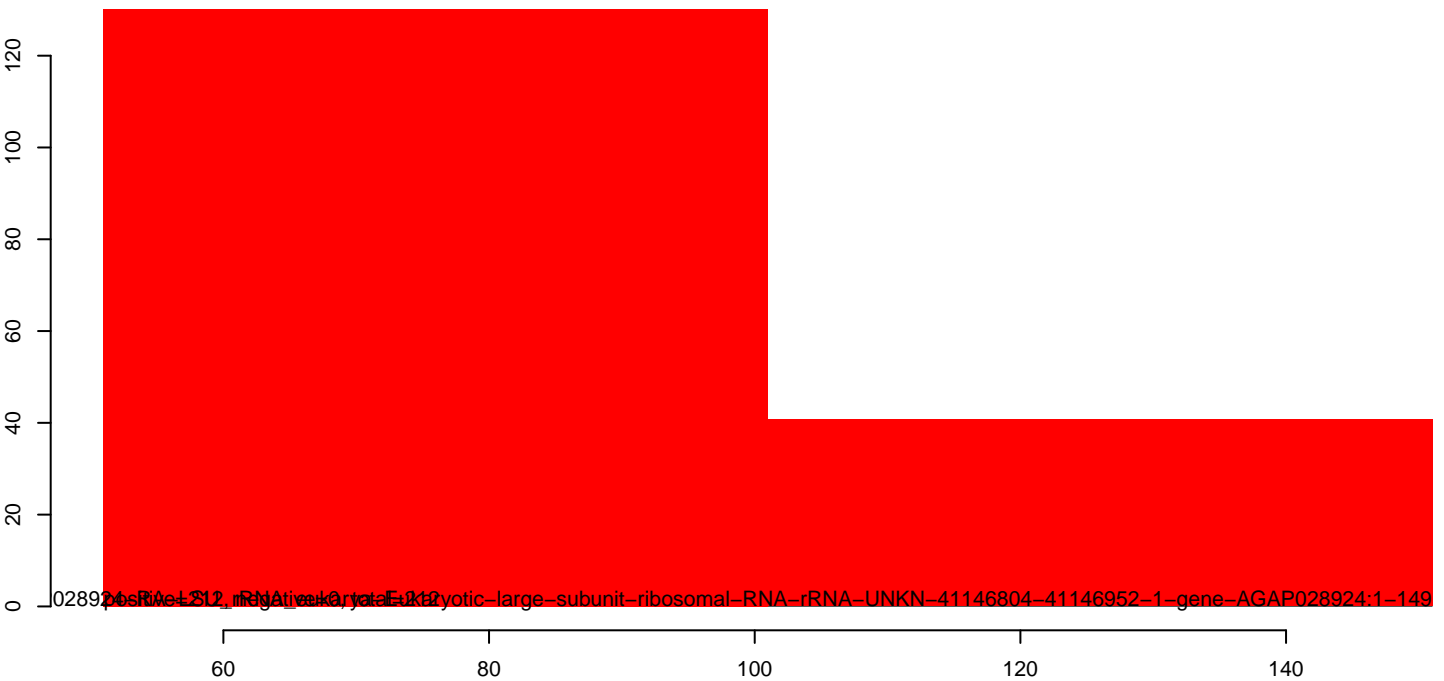
AnGam\_Fcirc\_TN.18\_23.rep



AnGam\_Fcirc\_TN.24\_35.rep



AnGam\_Fcirc\_TN.rep



028924:6sRiceLS12, negative=0, total=212, cyotic-large-subunit-ribosomal-RNA-rRNA-UNKN-41146804-41146952-1-gene-AGAP028924:1-149

60

80

100

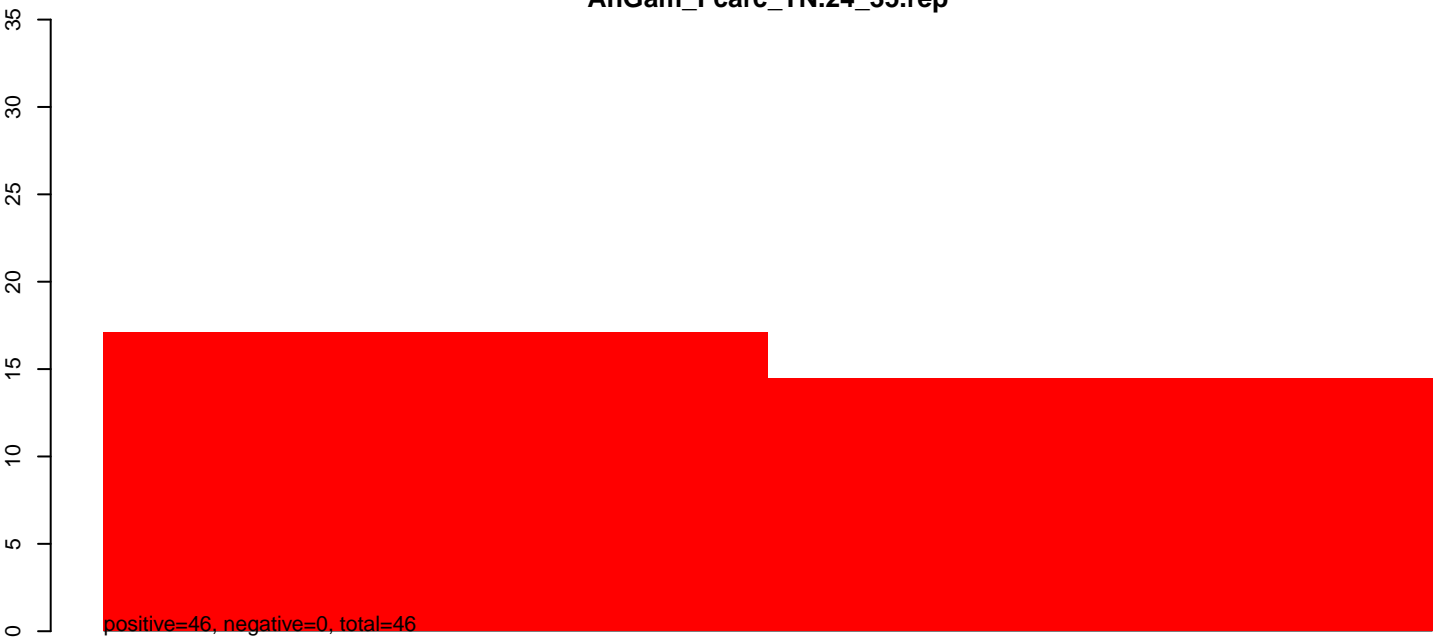
120

140

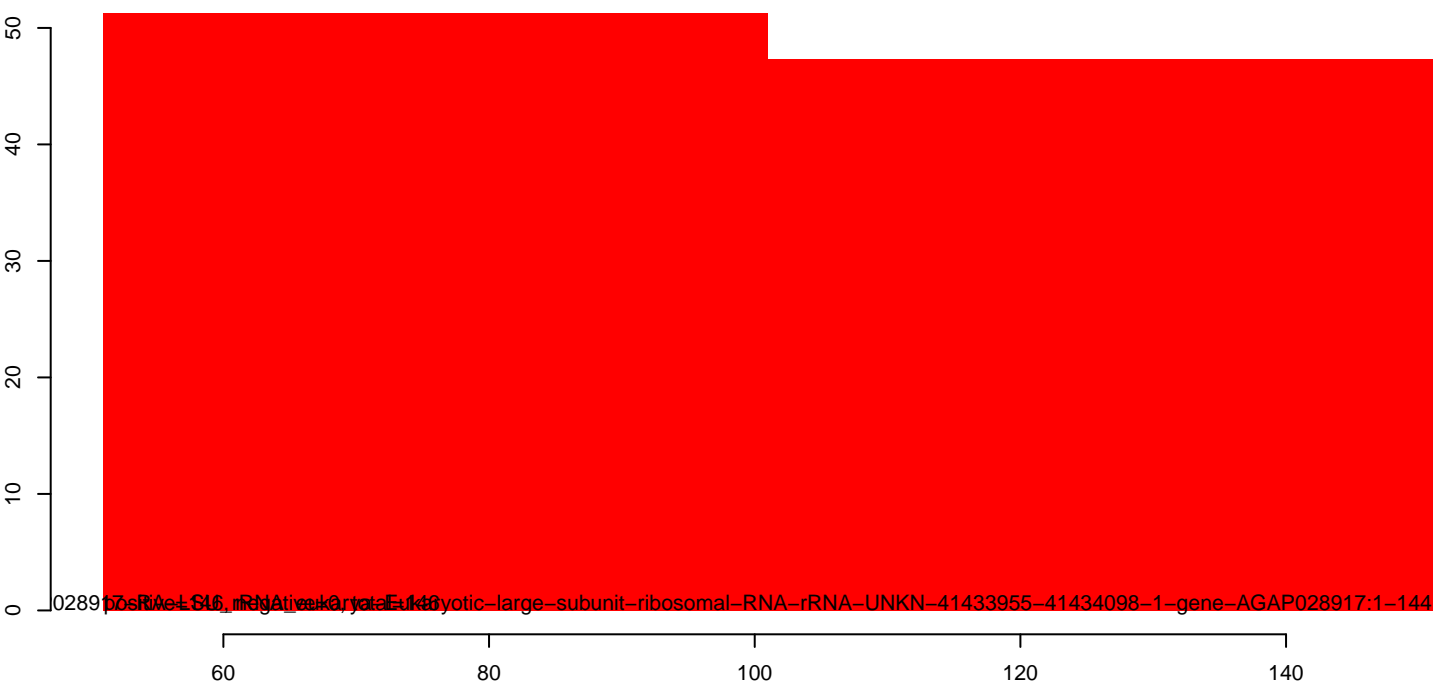
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AnGam\_Fcarc\_TN.24\_35.rep



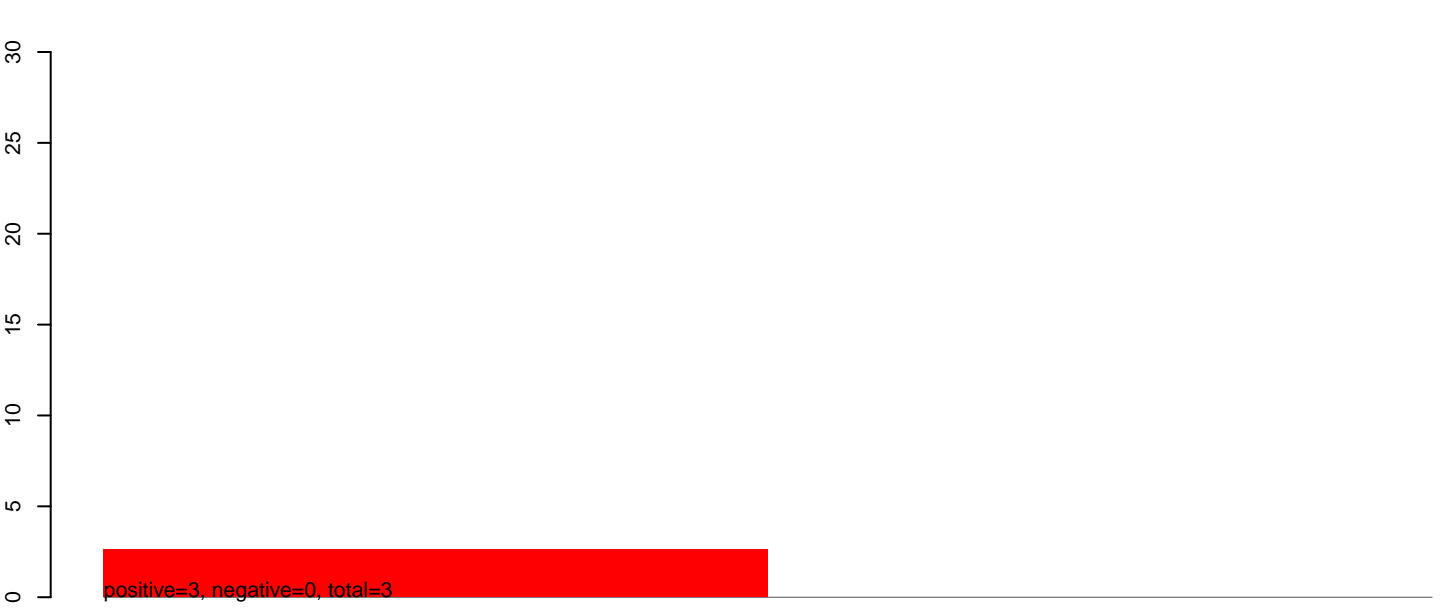
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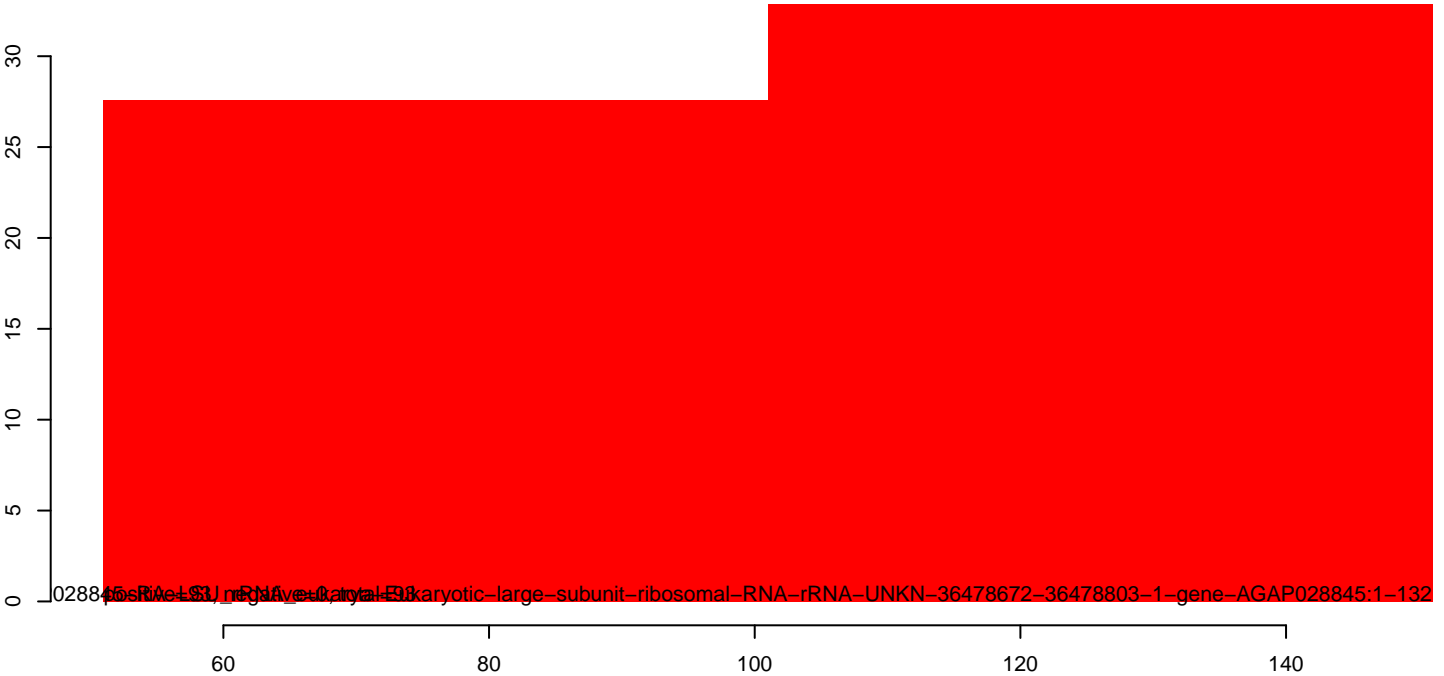
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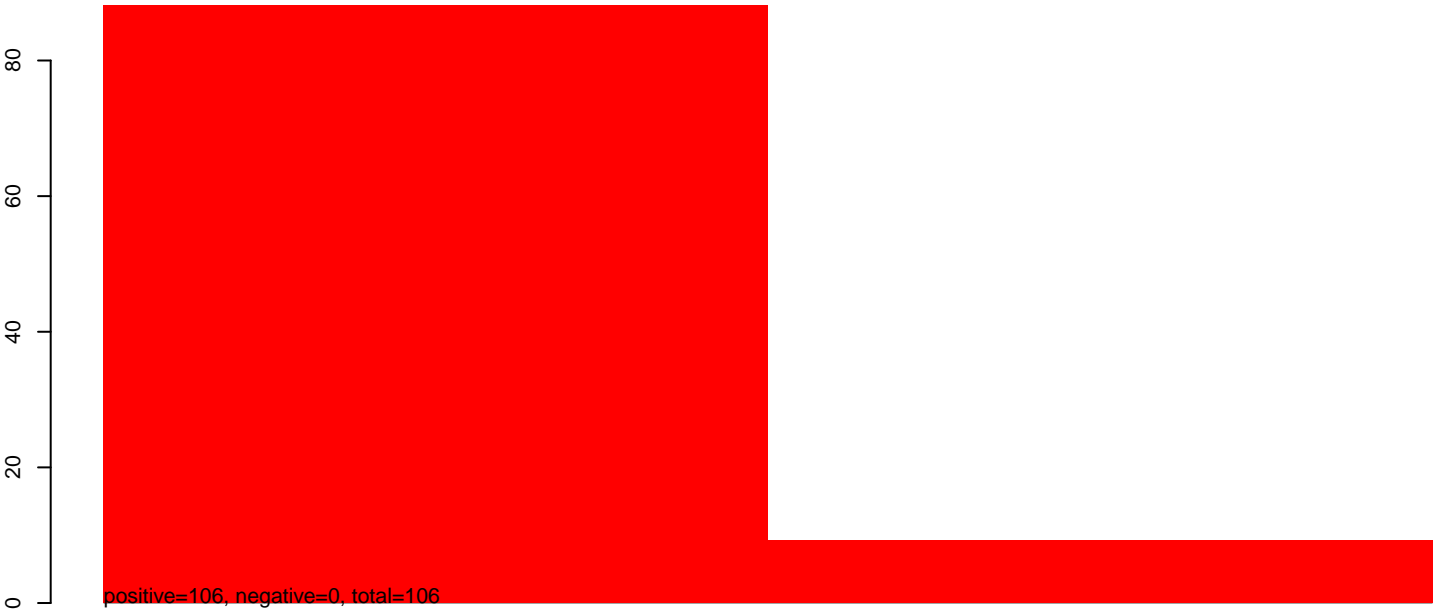
AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



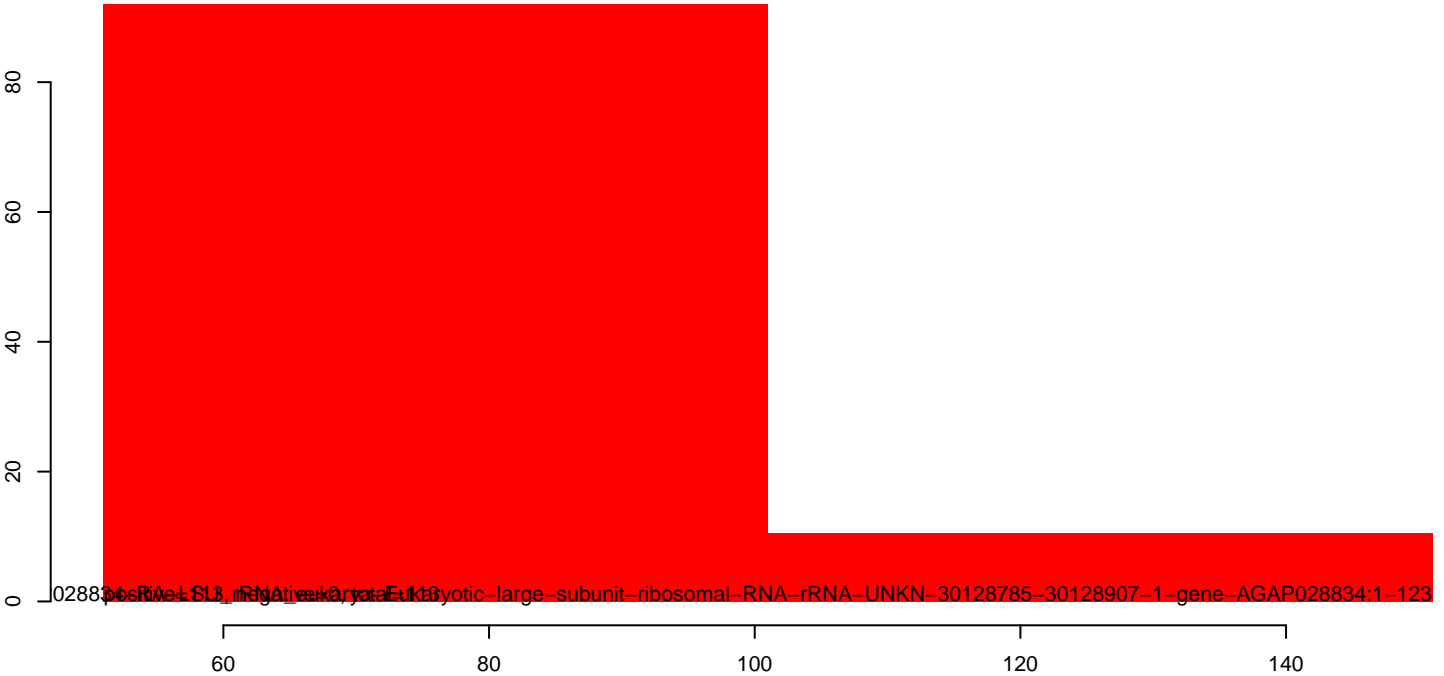
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AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



028809:SSU\_rRNA-31006077-31006203-1-gene-AGAP028809:1-127

60

80

100

120

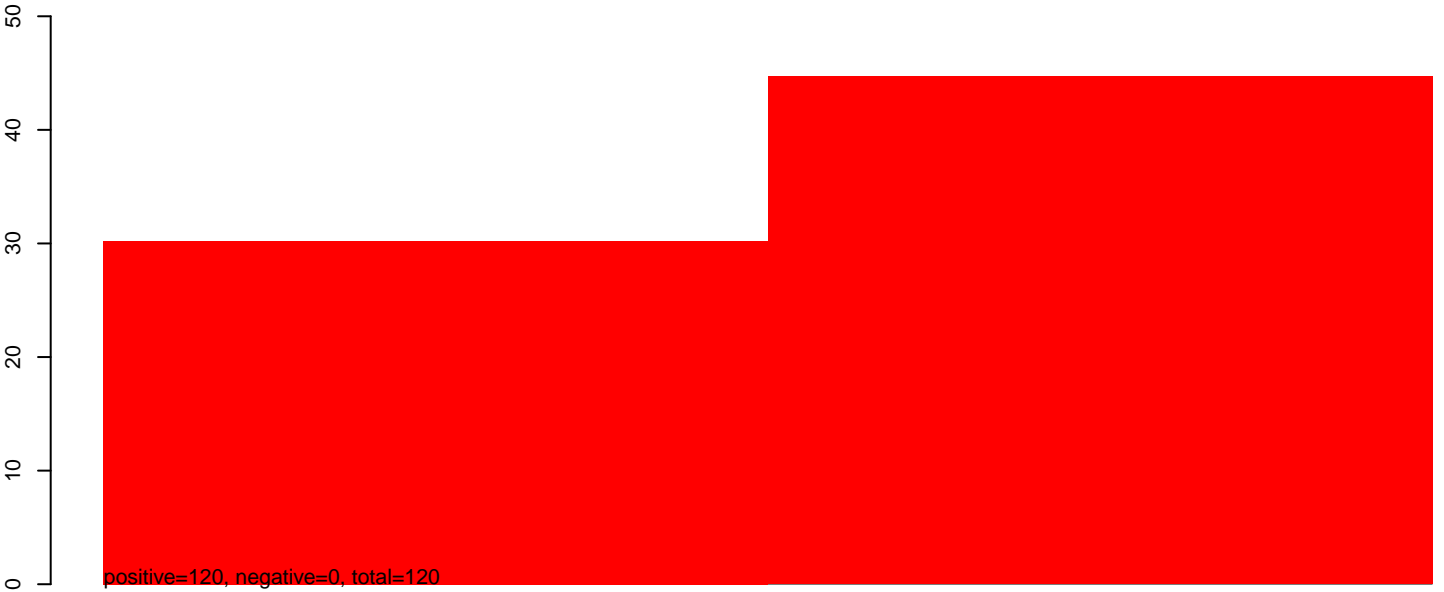
140



AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



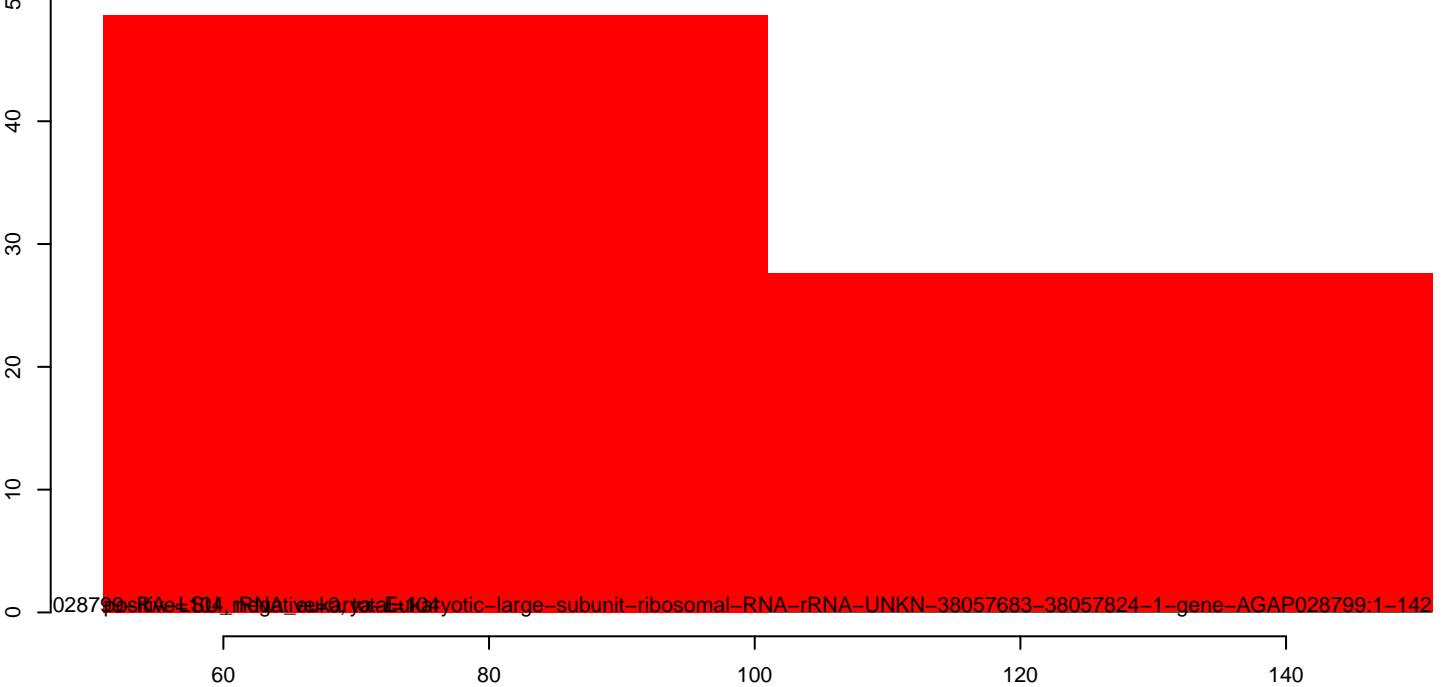
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AnGam\_Fcirc\_TN.24\_35.rep



AnGam\_Fcirc\_TN.rep



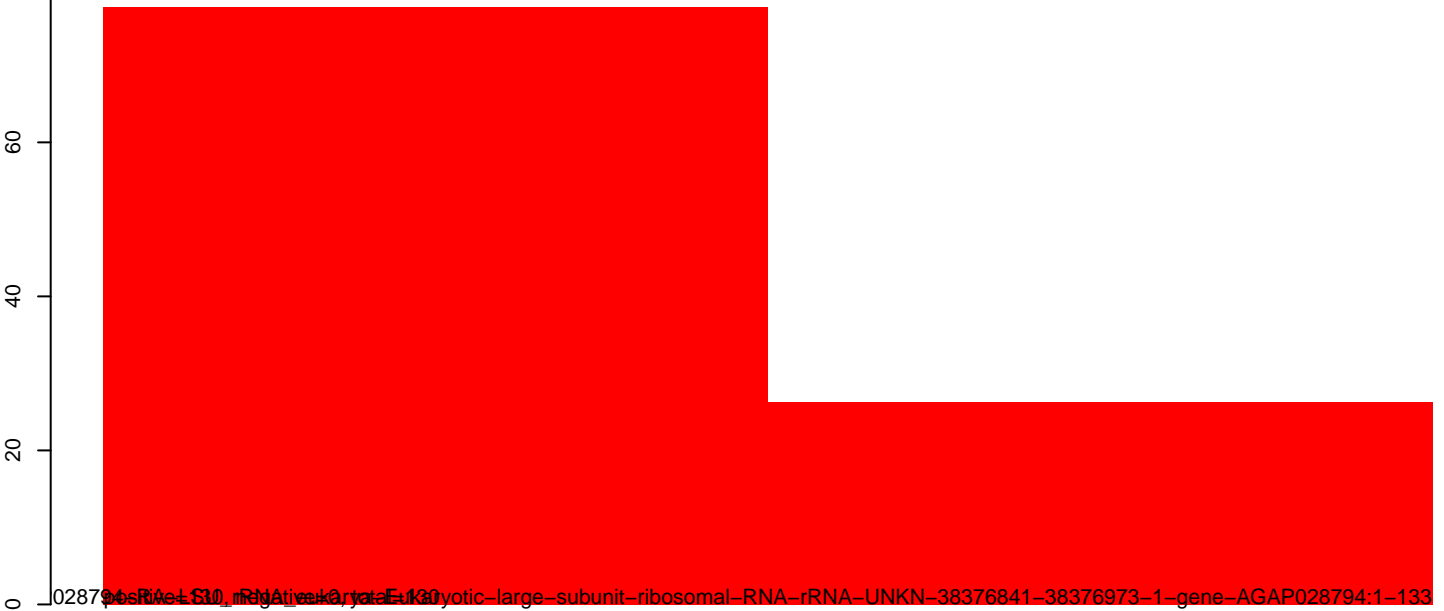
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AnGam\_Fcirc\_TN.24\_35.rep



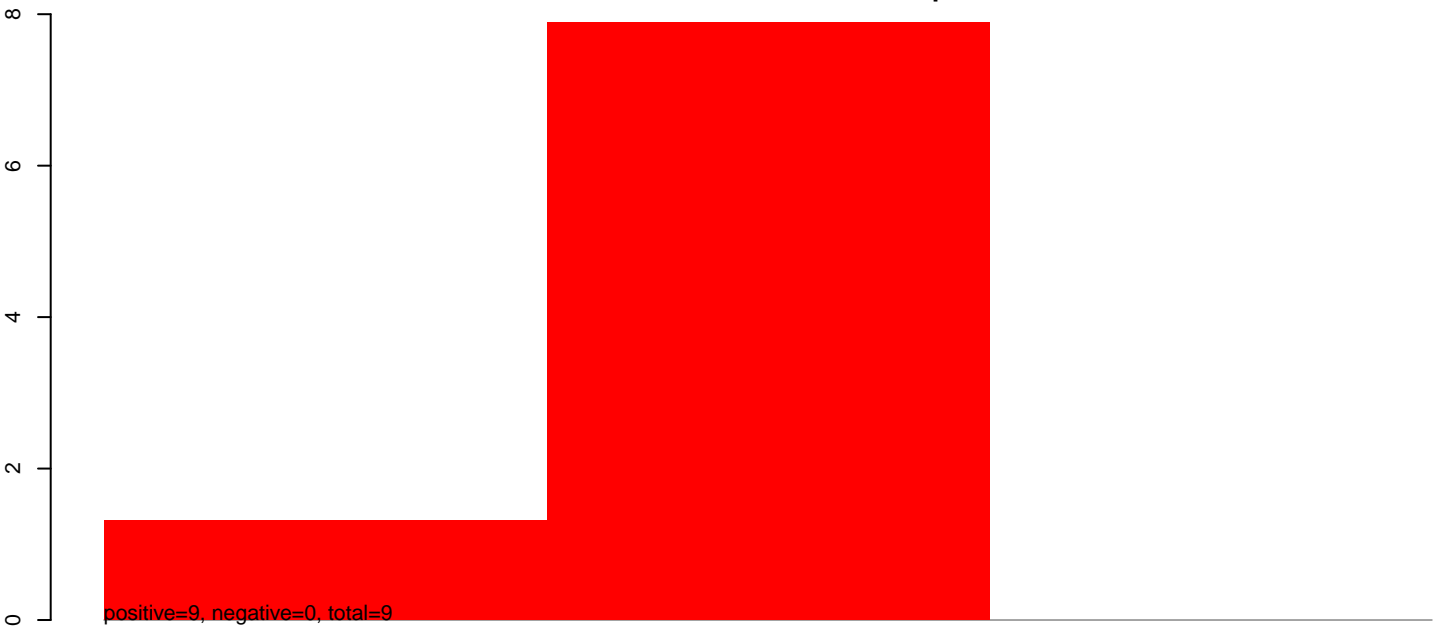
AnGam\_Fcirc\_TN.rep



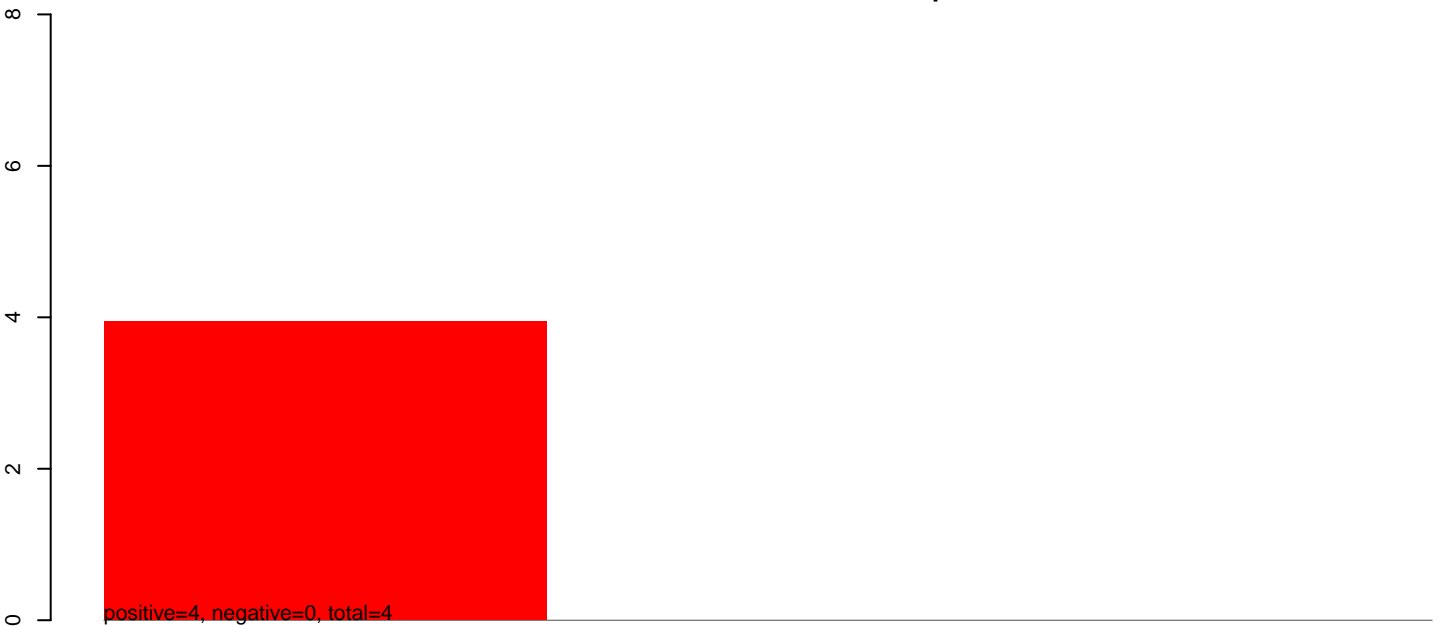
028794:1-133, positive=130, negative=0, total=130

60 80 100 120 140

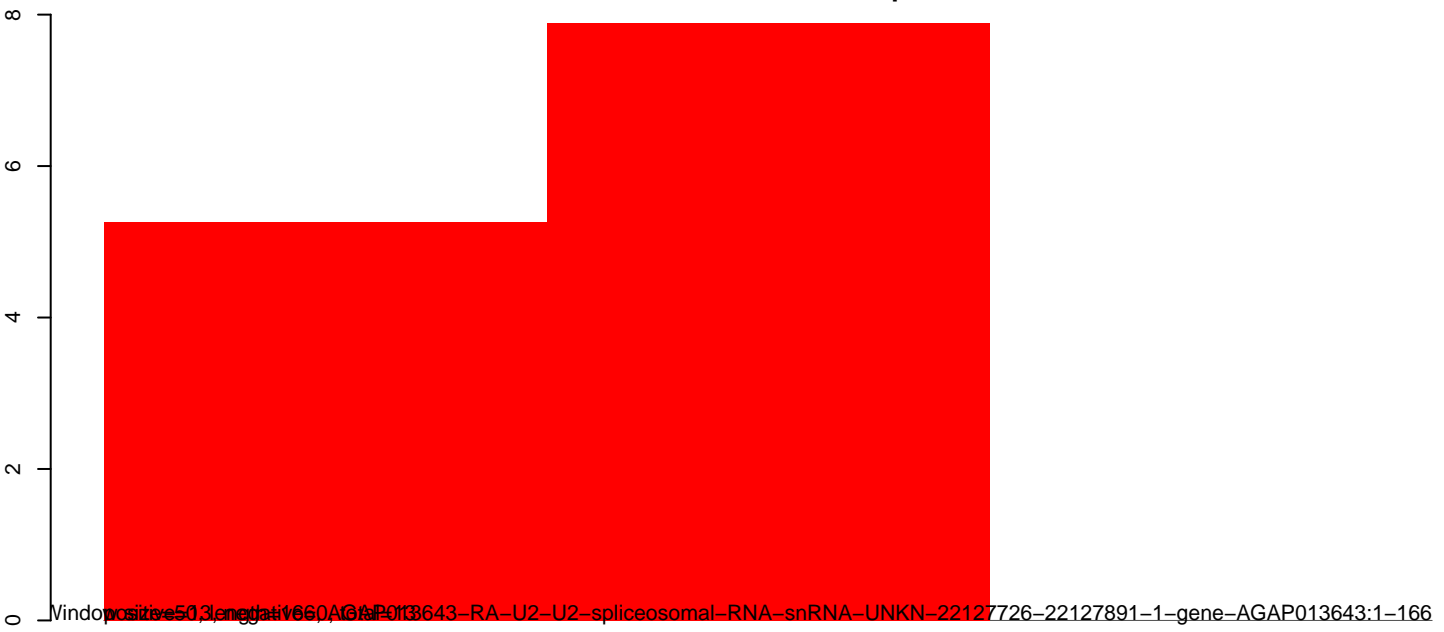
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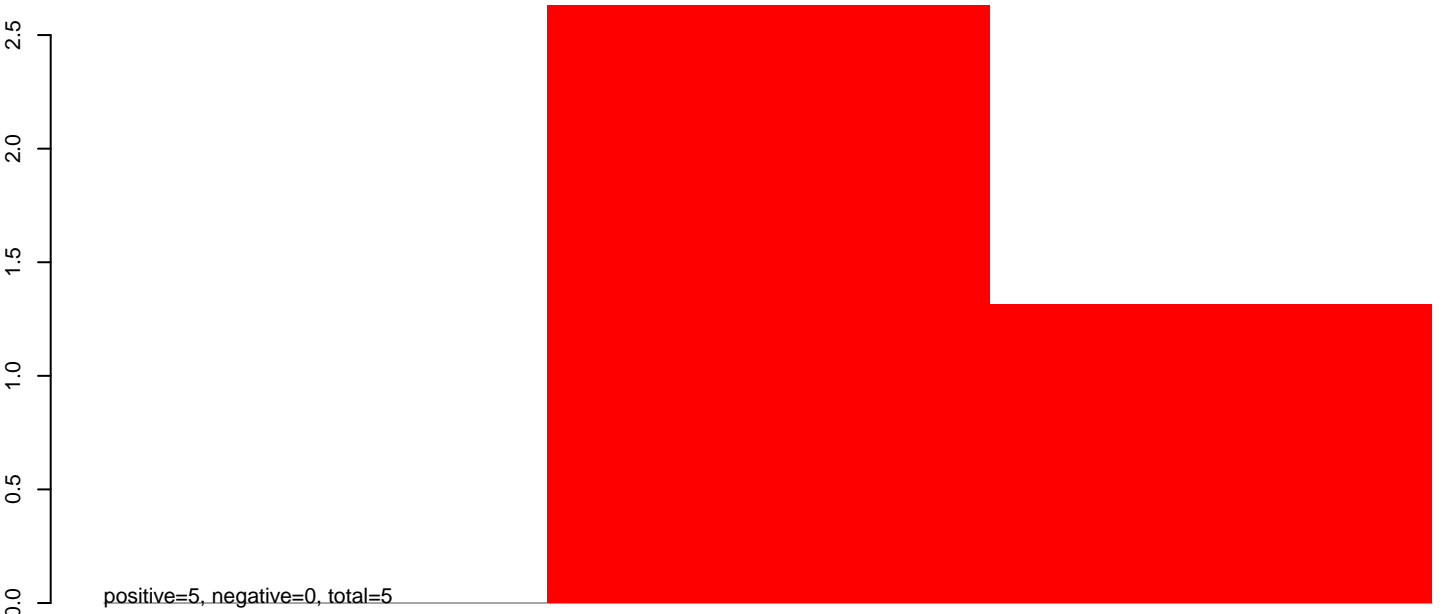
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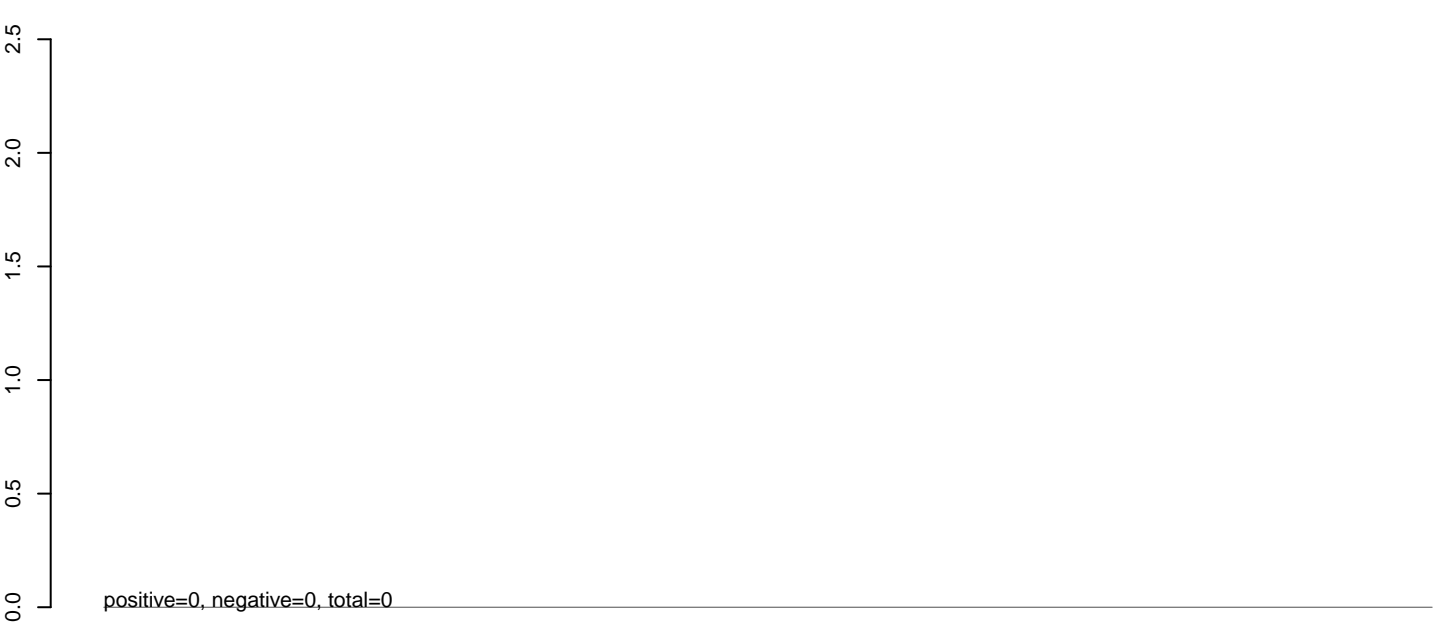
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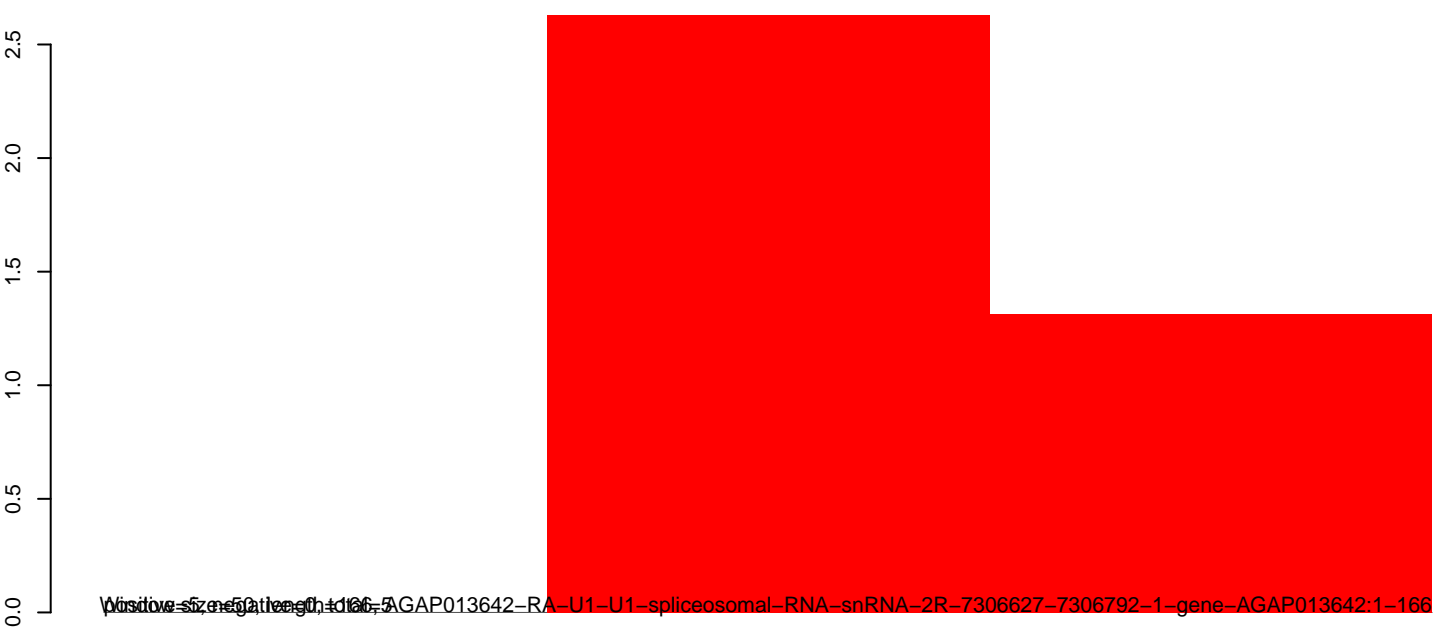
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AnGam\_Fcarc\_TN.24\_35.rep



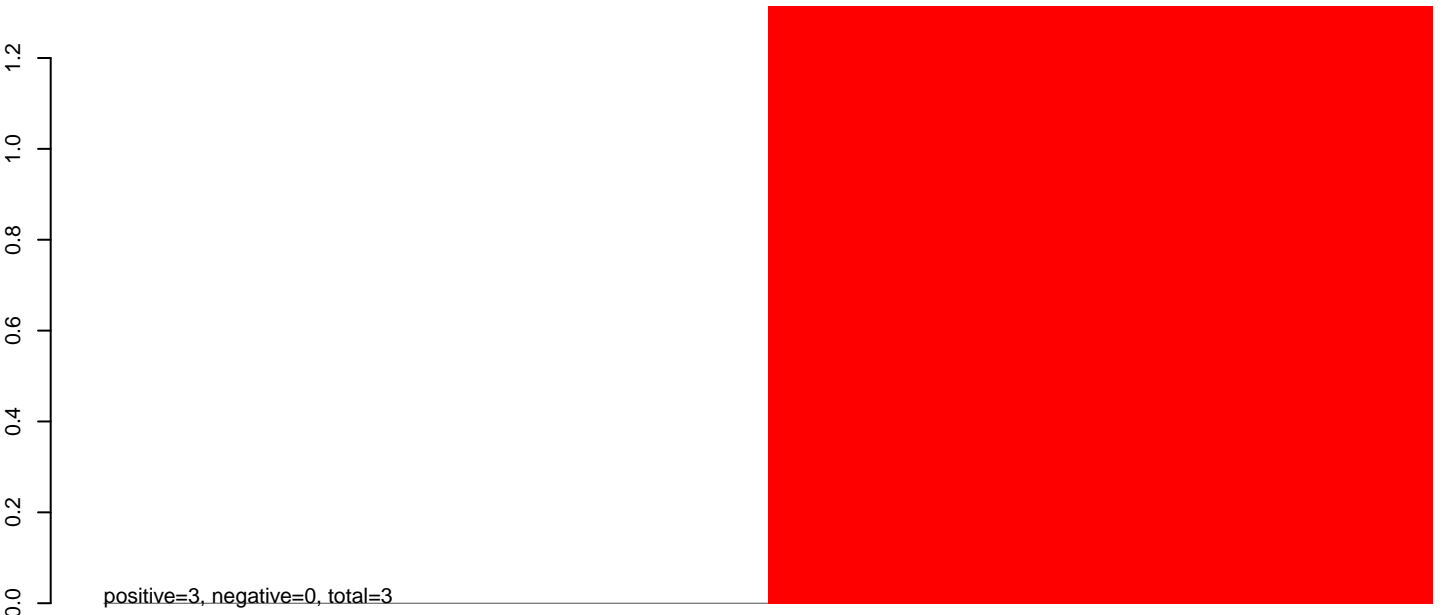
AnGam\_Fcarc\_TN.rep



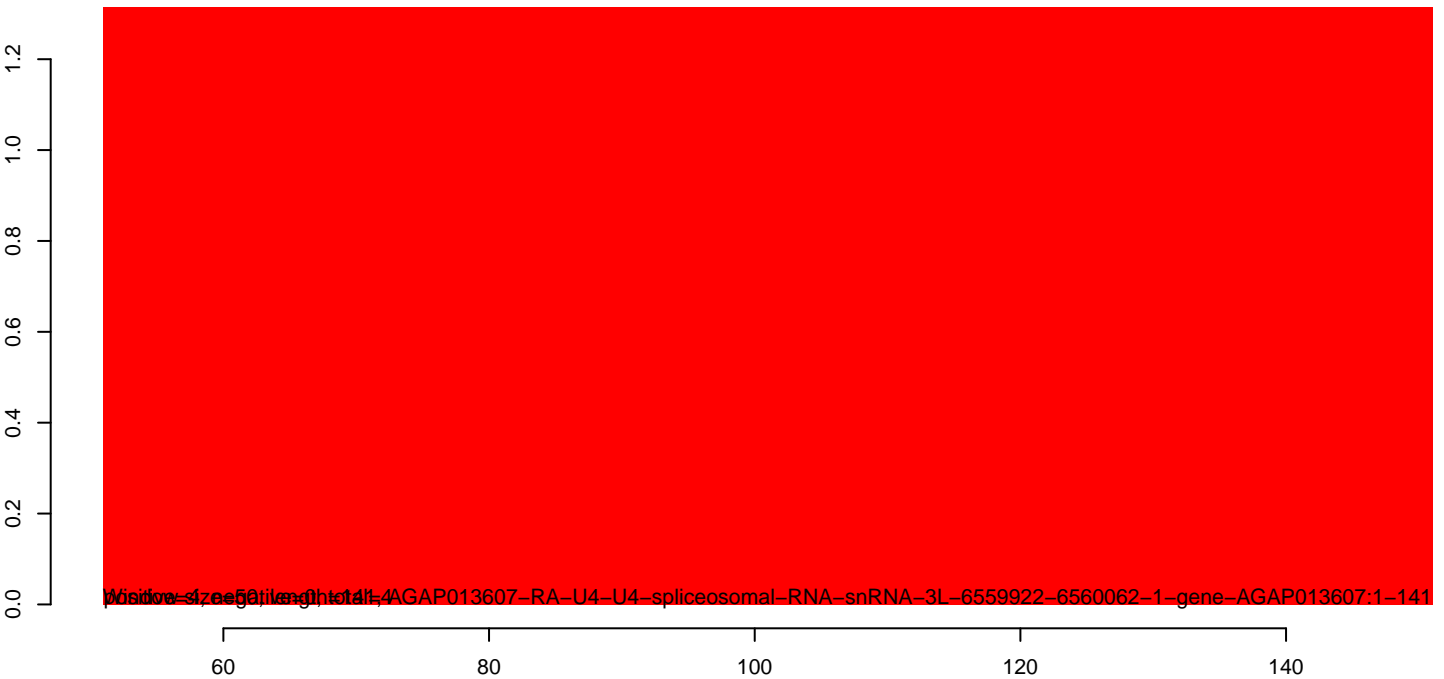
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AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



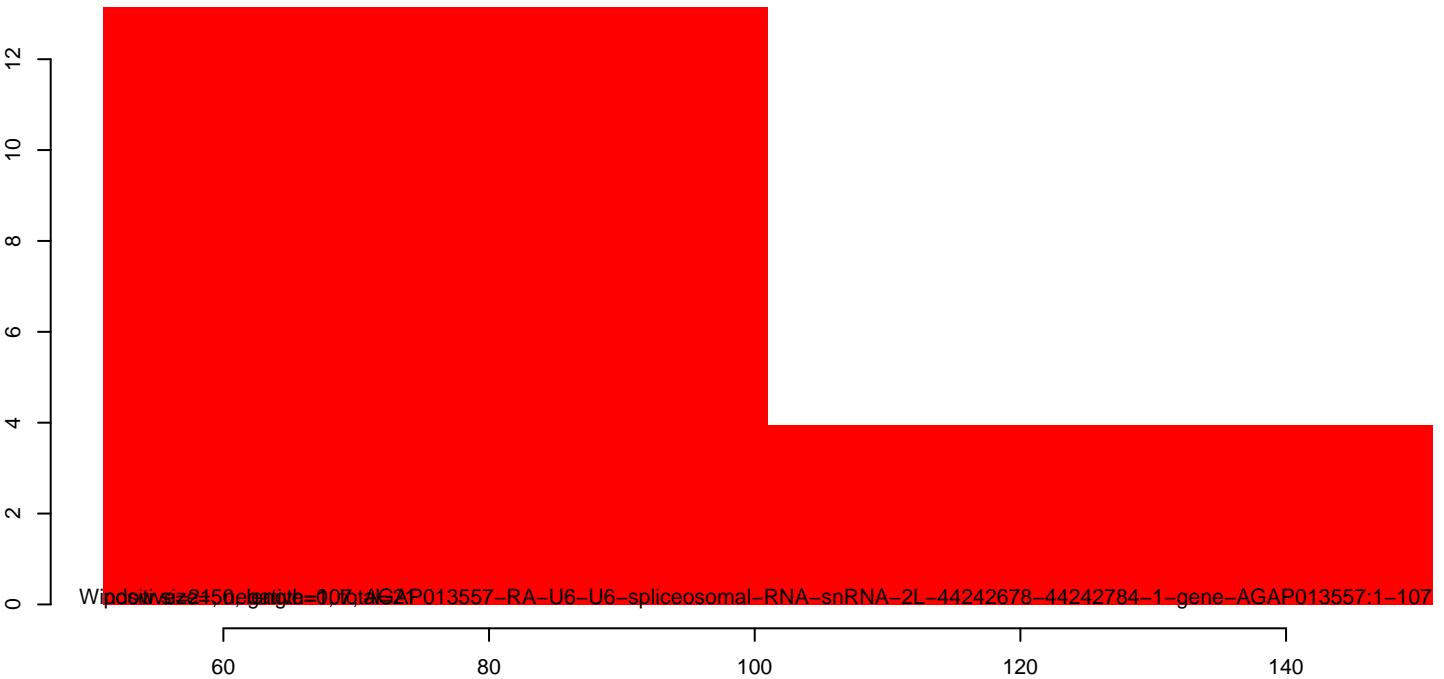
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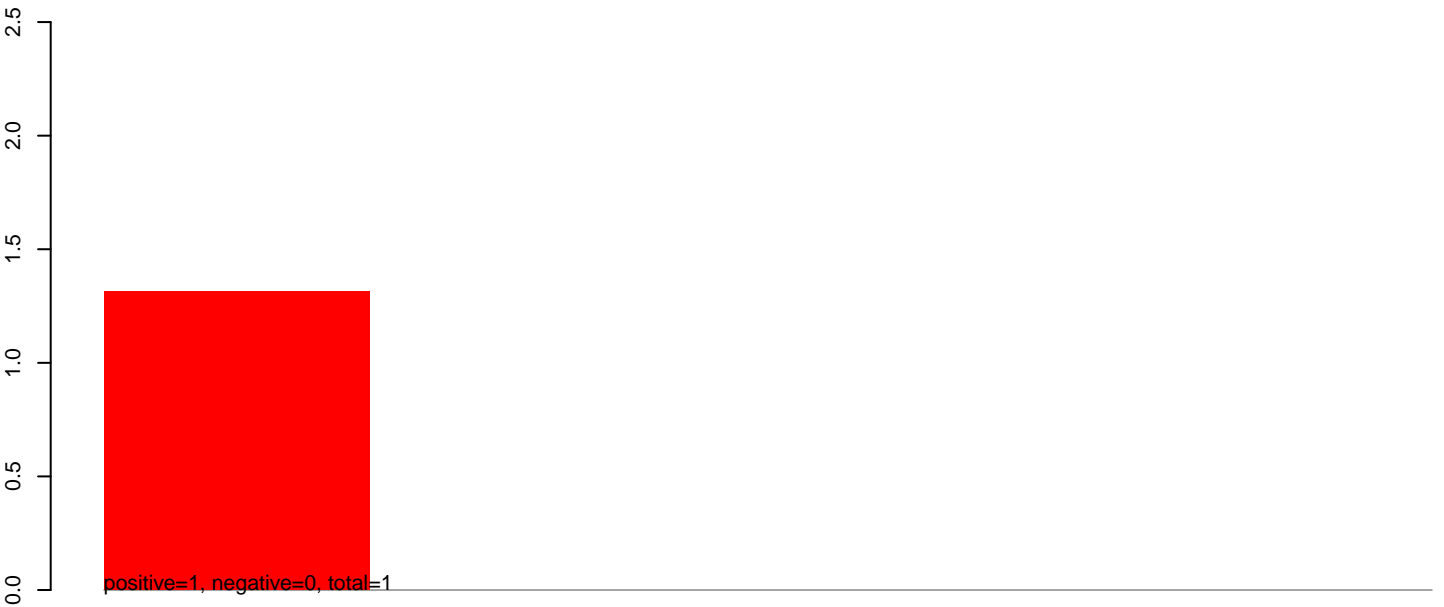
AnGam\_Fcarc\_TN.24\_35.rep



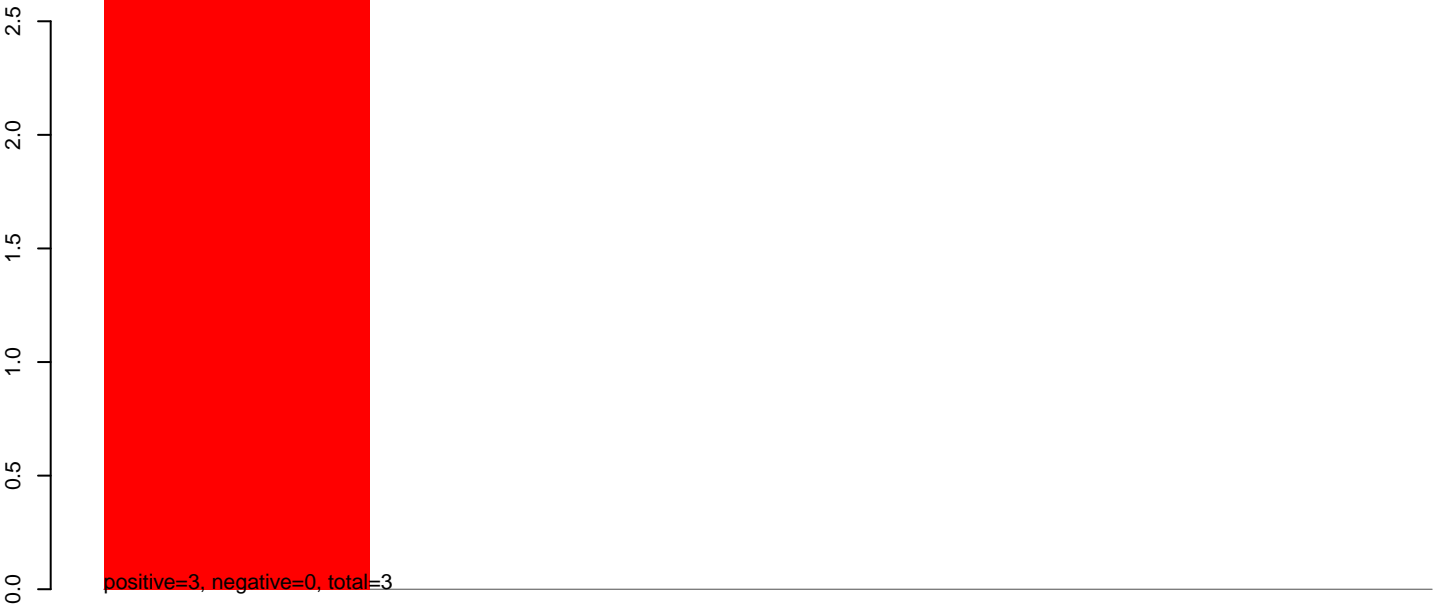
AnGam\_Fcarc\_TN.rep



### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep

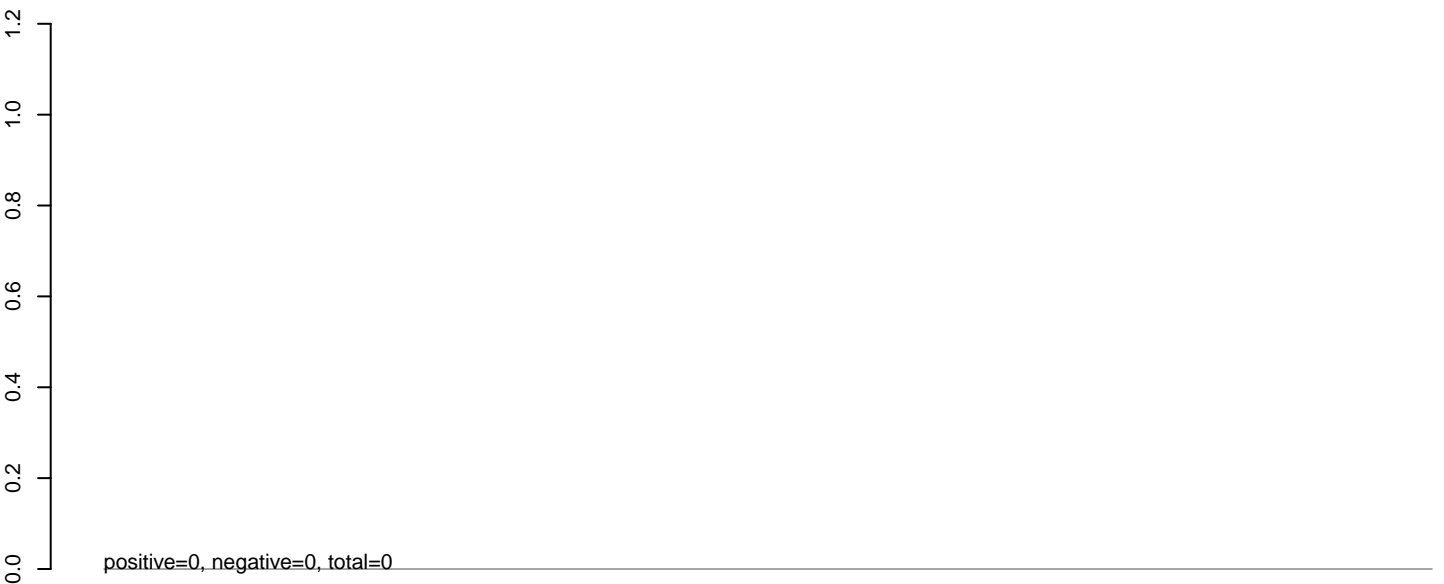


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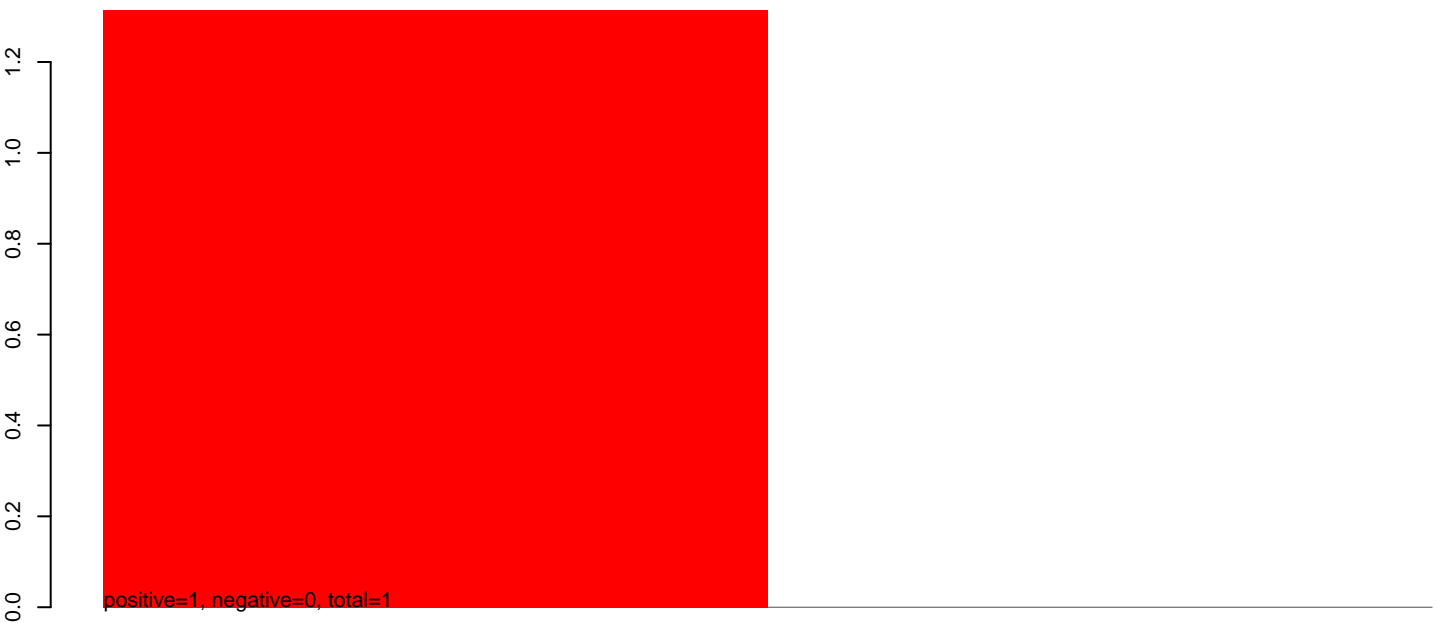




AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



Window size=50, length=100, AGAP028981-RA-U6-U6-spliceosomal-RNA-snrRNA-2L-21351634-21351782-1-gene-AGAP028981:1-149

60 80 100 120 140



### AnGam\_Fcarc\_TN.18\_23.rep

15  
10  
5  
0

positive=18, negative=0, total=18

### AnGam\_Fcarc\_TN.24\_35.rep

15  
10  
5  
0

positive=29, negative=0, total=29

### AnGam\_Fcarc\_TN.rep

20  
15  
10  
5  
0

positive=47, negative=0, total=47  
Window size=50, length=72, AGAP028392-RA-tRNA-Val-tRNA-Mt-13973-14044-1-gene-AGAP028392:1-72

50 60 70 80 90 100

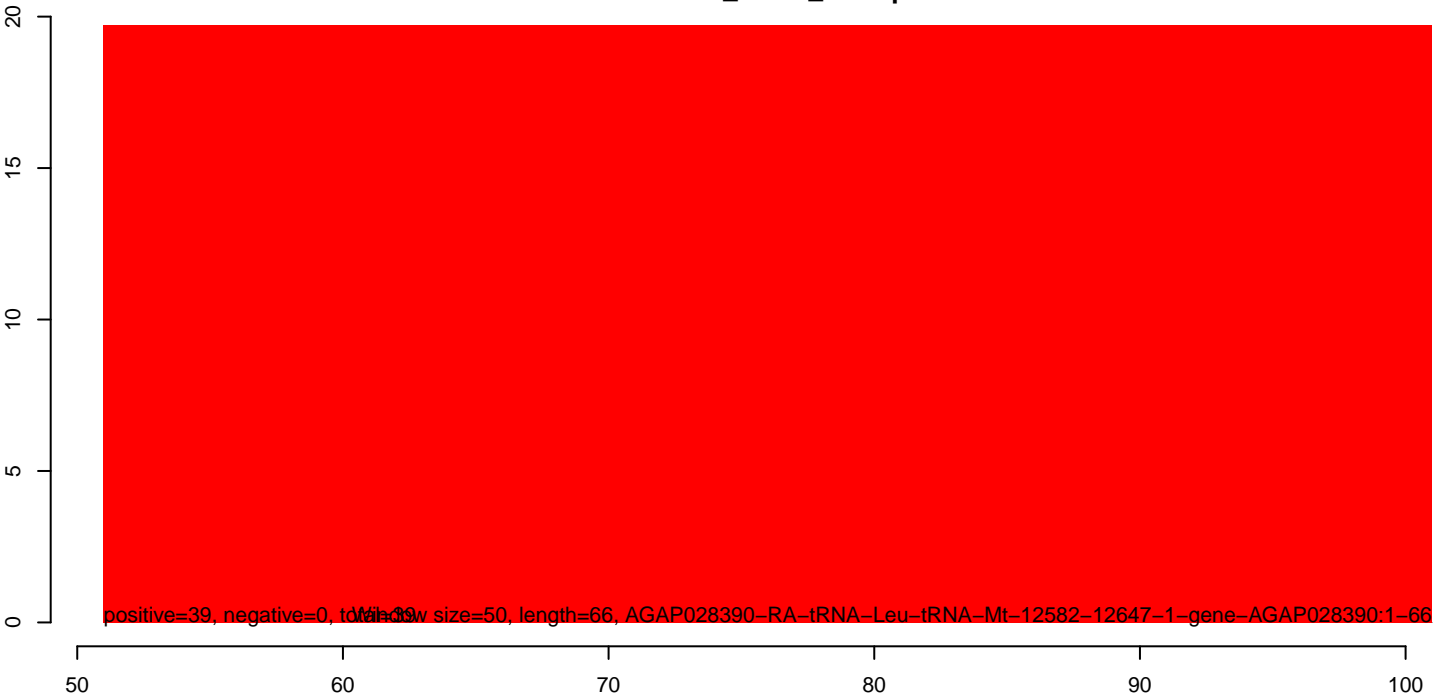
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AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



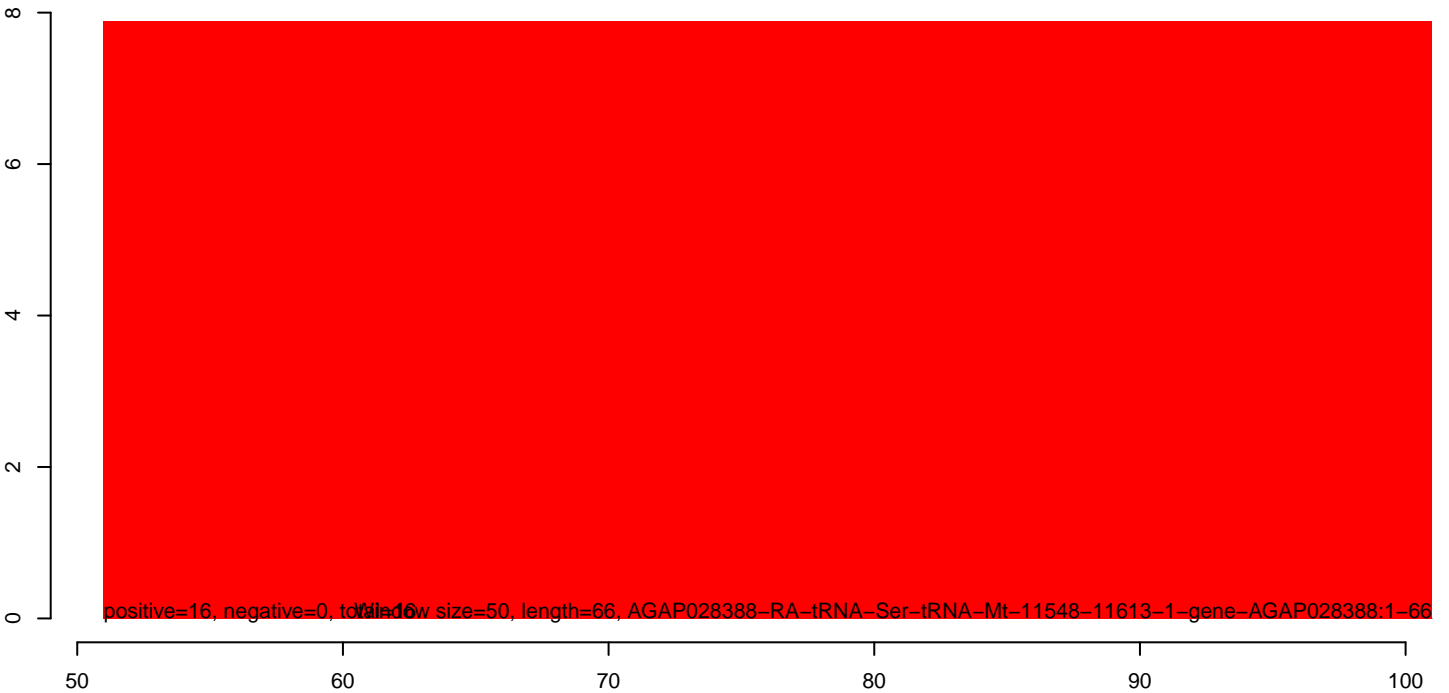
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### AnGam\_Fcarc\_TN.24\_35.rep



### AnGam\_Fcarc\_TN.rep



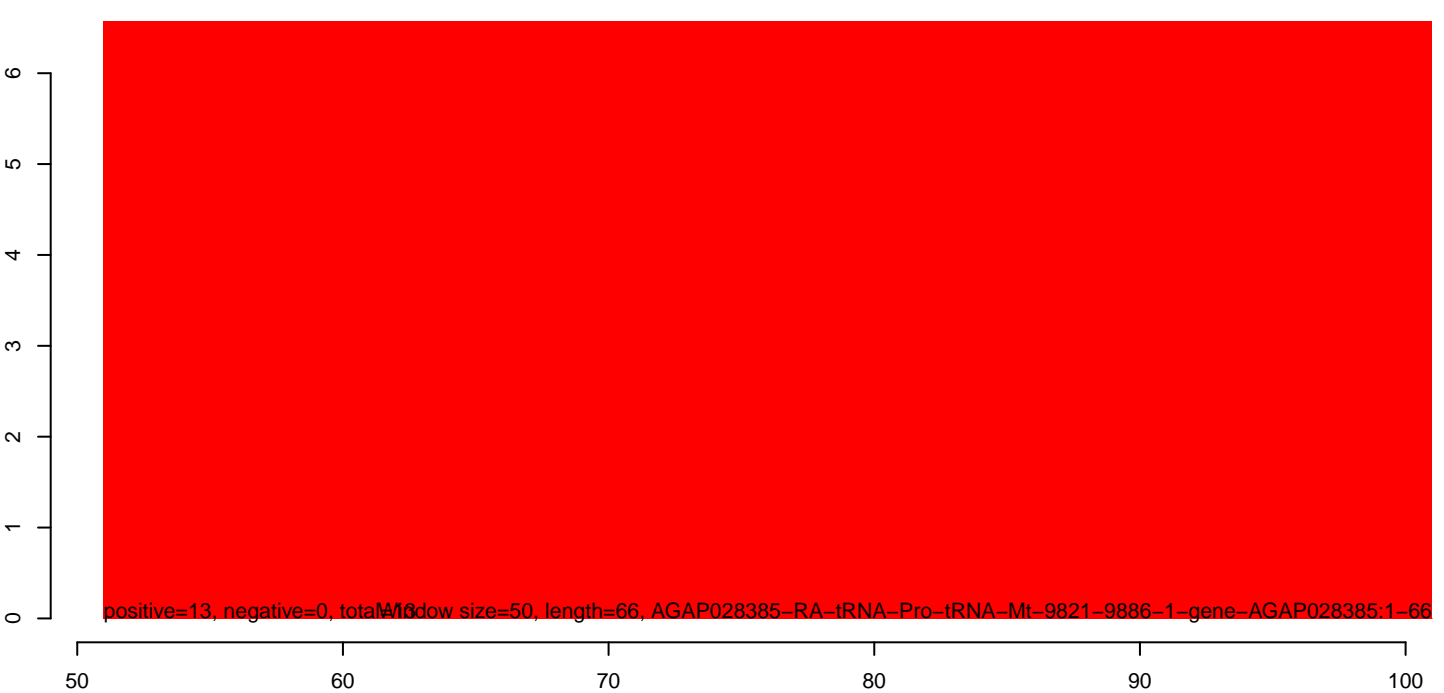
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AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



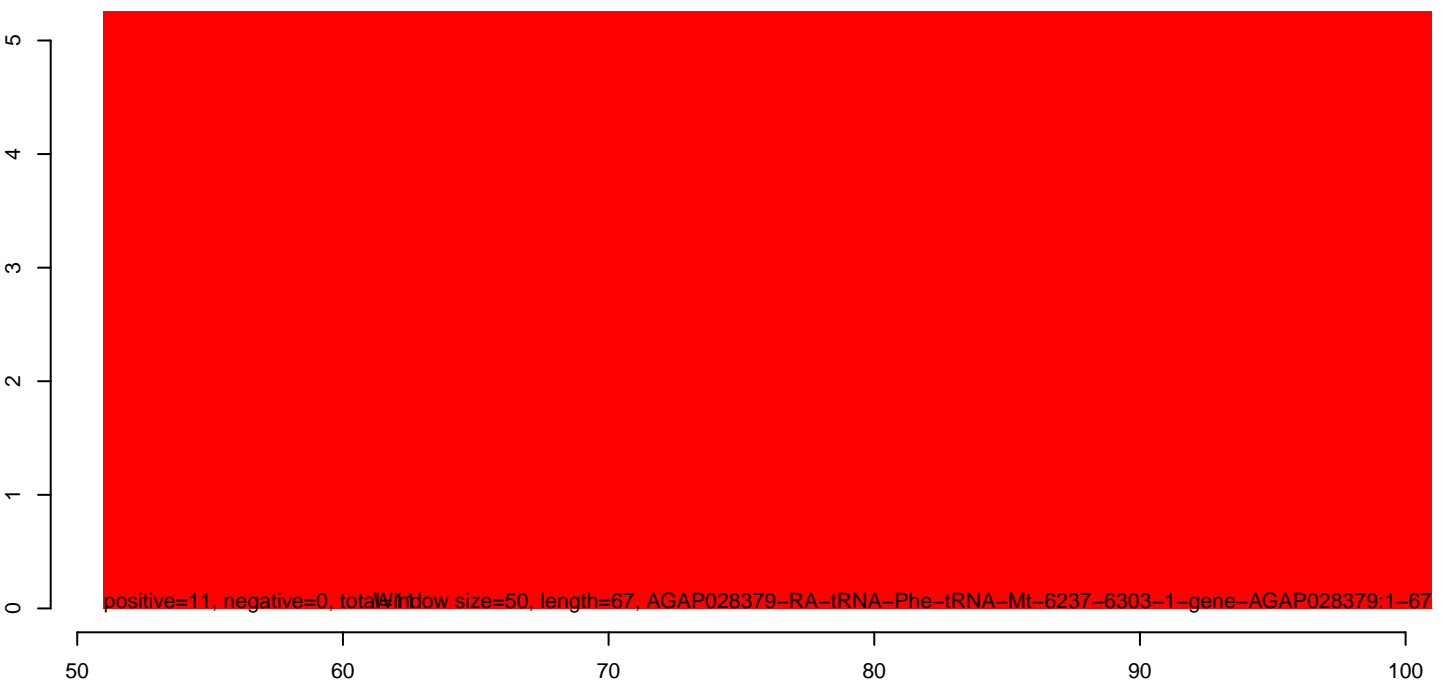
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AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep

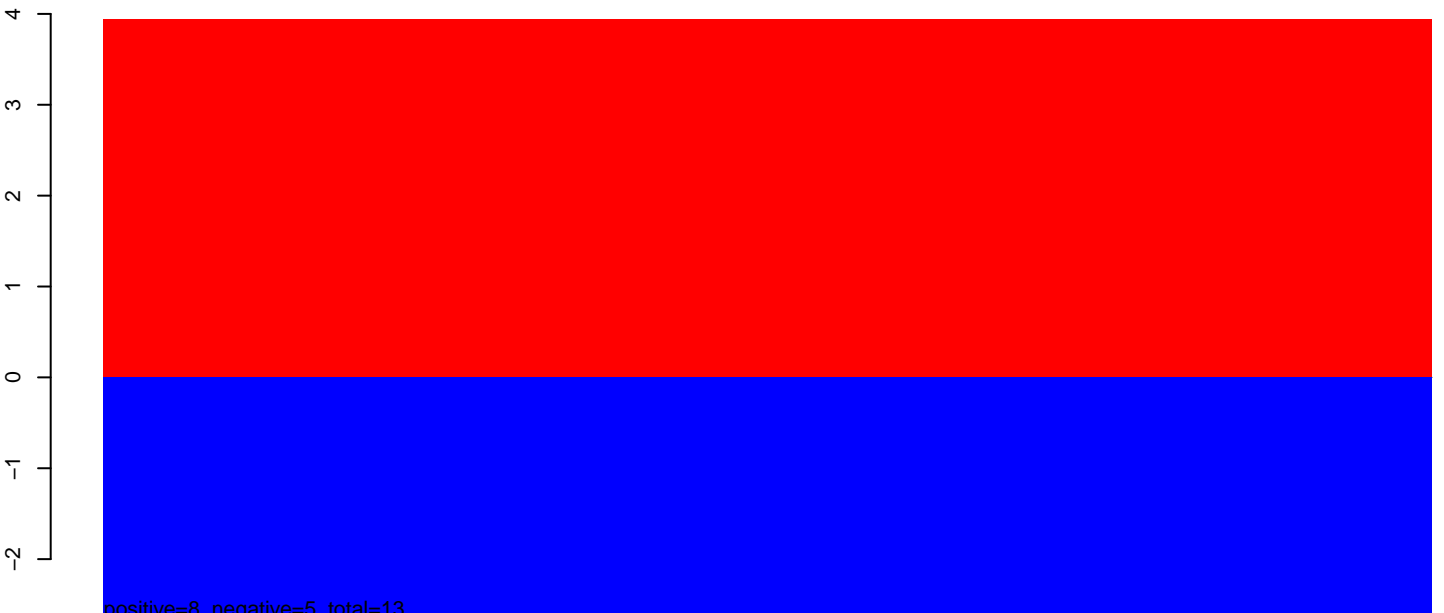


AnGam\_Fcarc\_TN.18\_23.rep



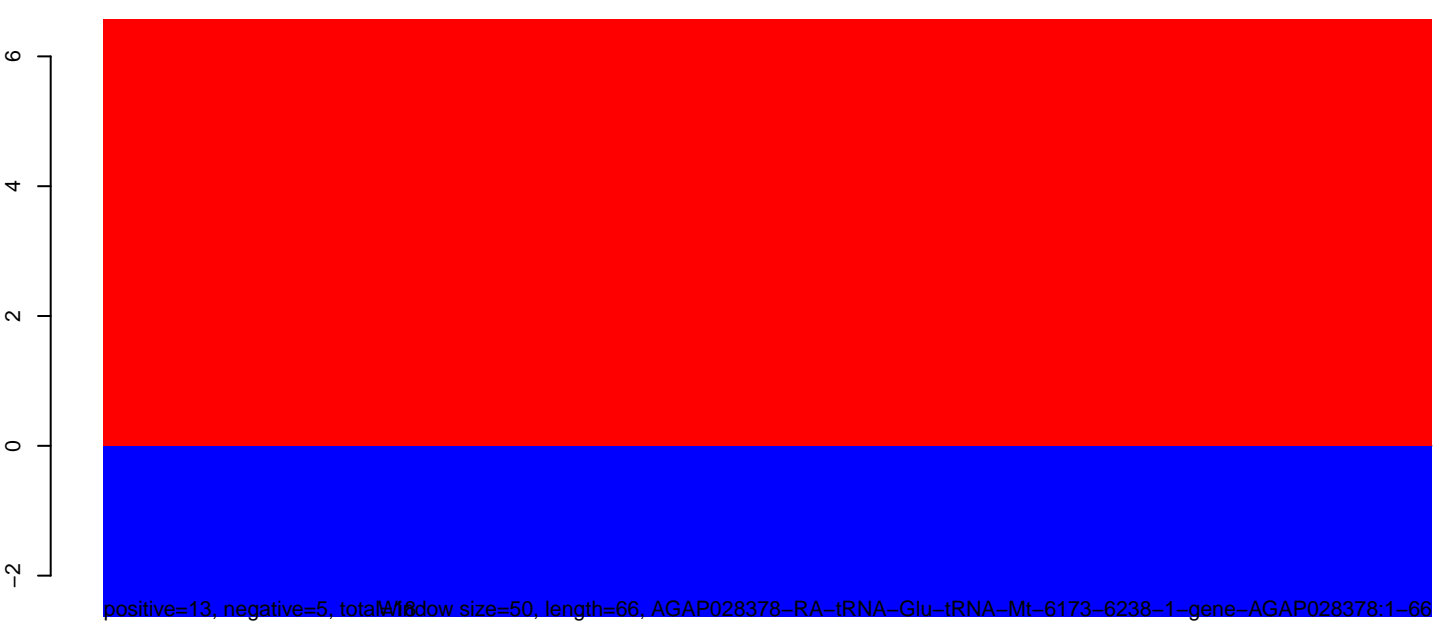
positive=5, negative=0, total=5

AnGam\_Fcarc\_TN.24\_35.rep



positive=8, negative=5, total=13

AnGam\_Fcarc\_TN.rep



positive=13, negative=5, total=18 window size=50, length=66, AGAP028378-RA-tRNA-Glu-tRNA-Mt-6173-6238-1-gene-AGAP028378:1-66

50 60 70 80 90 100



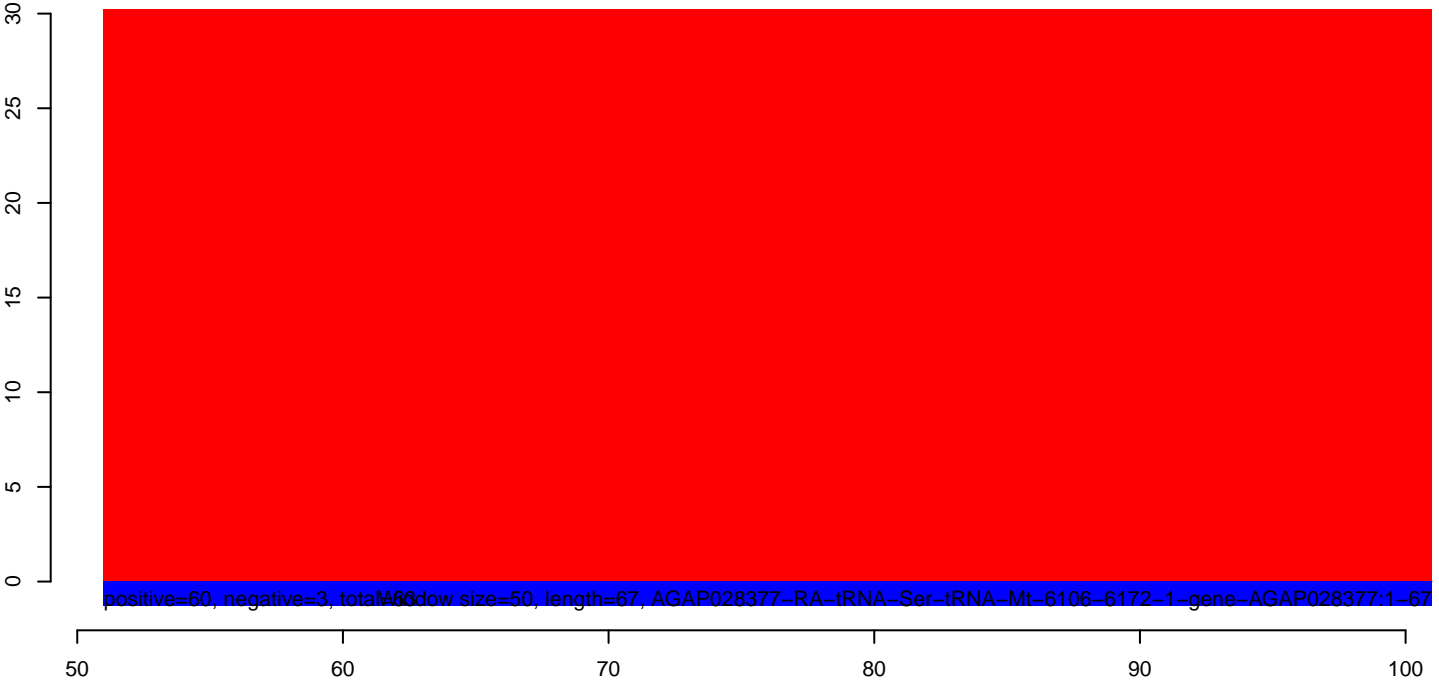
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AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



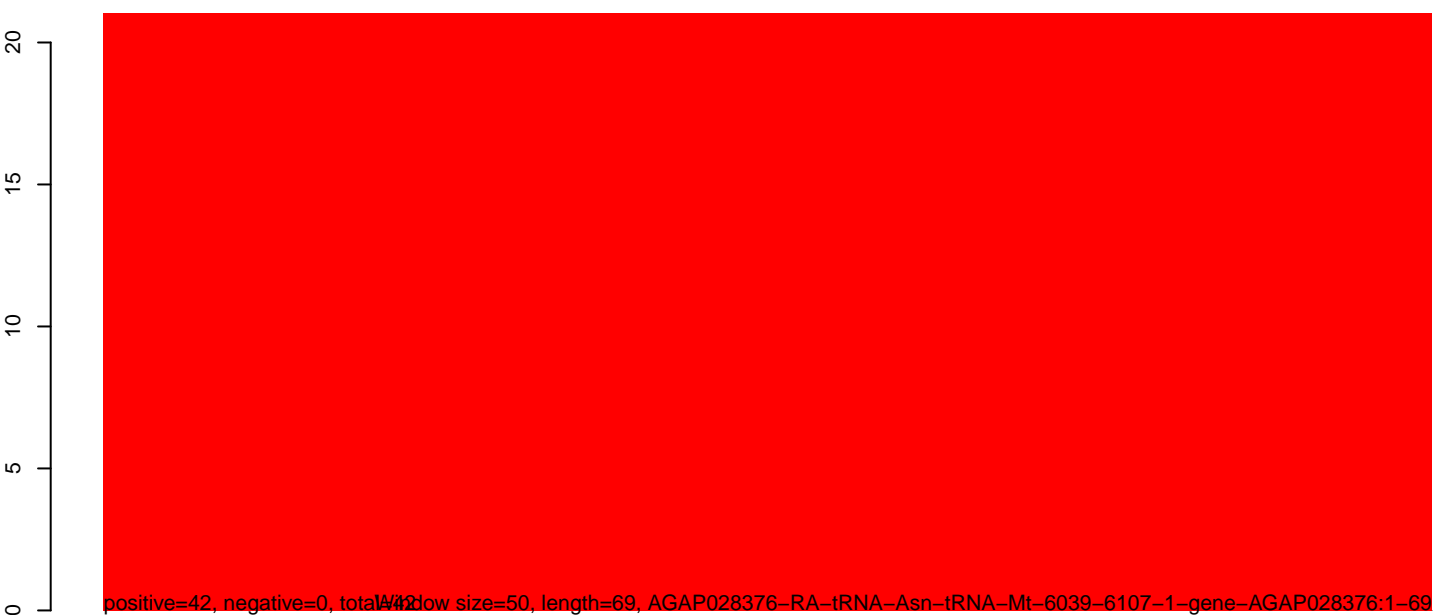
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### AnGam\_Fcarc\_TN.24\_35.rep



### AnGam\_Fcarc\_TN.rep



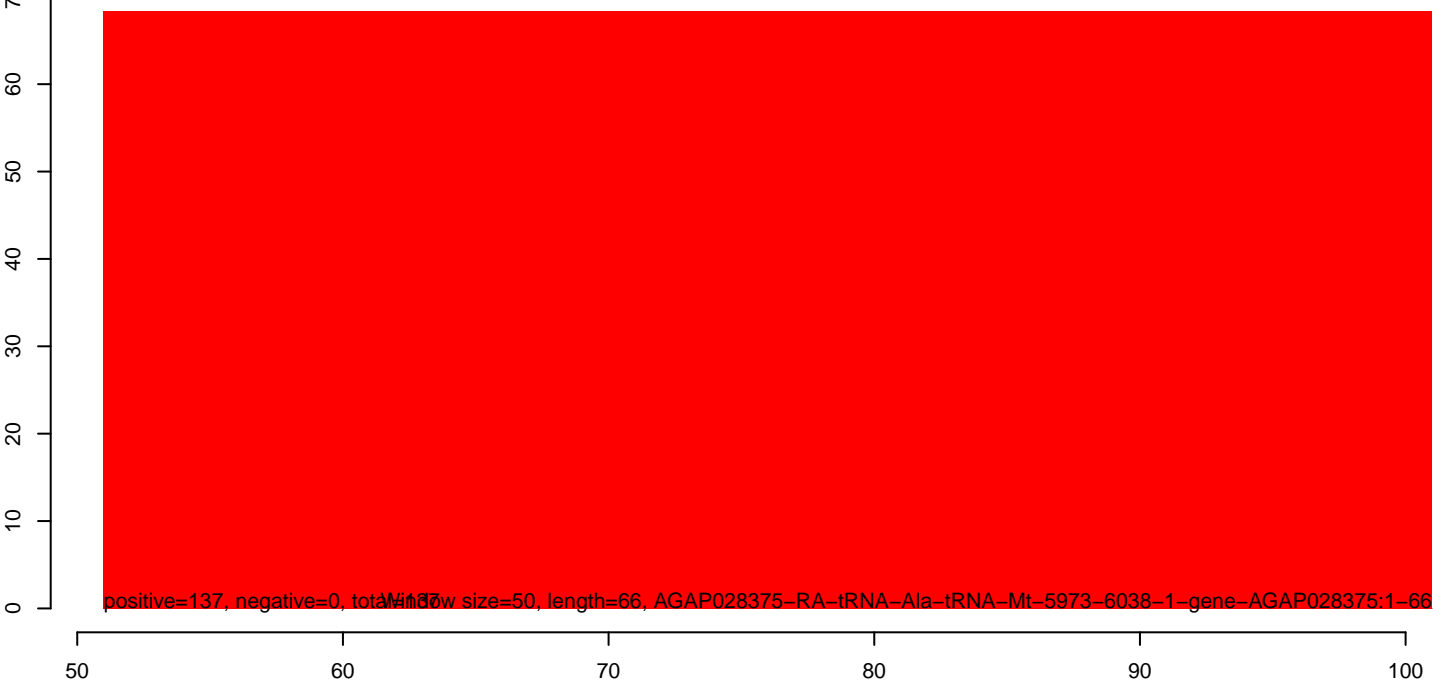
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AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



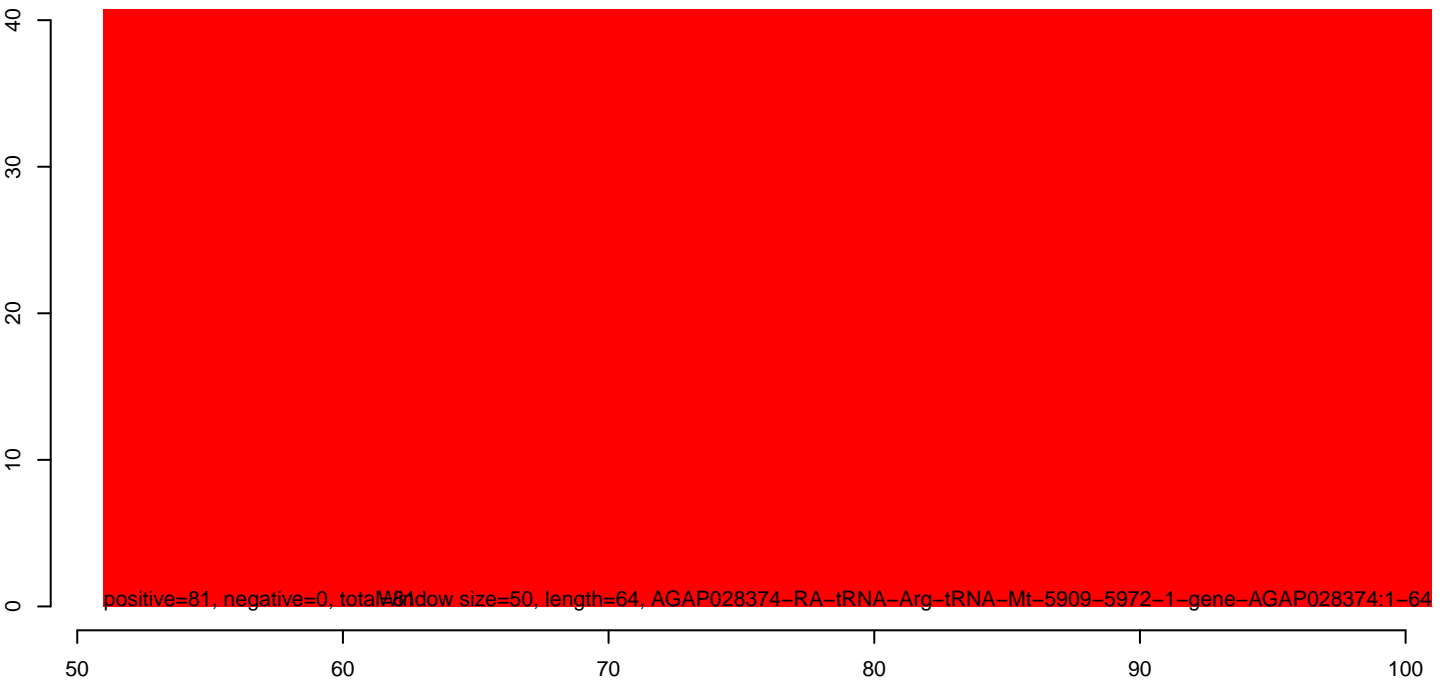
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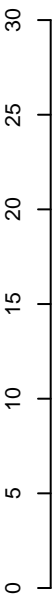
### AnGam\_Fcarc\_TN.24\_35.rep



### AnGam\_Fcarc\_TN.rep

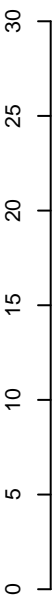


### AnGam\_Fcarc\_TN.18\_23.rep



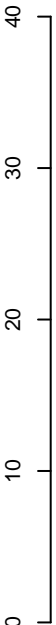
positive=16, negative=0, total=16

### AnGam\_Fcarc\_TN.24\_35.rep

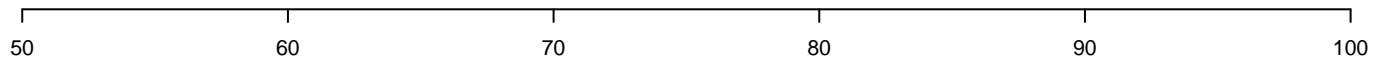


positive=63, negative=0, total=63

### AnGam\_Fcarc\_TN.rep



positive=79, negative=0, total=79 window size=50, length=67, AGAP028372-RA-tRNA-Gly-tRNA-Mt-5490-5556-1-gene-AGAP028372:1-67



### AnGam\_Fcarc\_TN.18\_23.rep

100  
80  
60  
40  
20  
0

positive=50, negative=0, total=50

### AnGam\_Fcarc\_TN.24\_35.rep

100  
80  
60  
40  
20  
0

positive=197, negative=0, total=197

### AnGam\_Fcarc\_TN.rep

120  
100  
80  
60  
40  
20  
0

positive=247, negative=0, total=247  
Window size=50, length=69, AGAP028368-RA-tRNA-Asp-tRNA-Mt-3799-3867-1-gene-AGAP028368:1-69

50 60 70 80 90 100

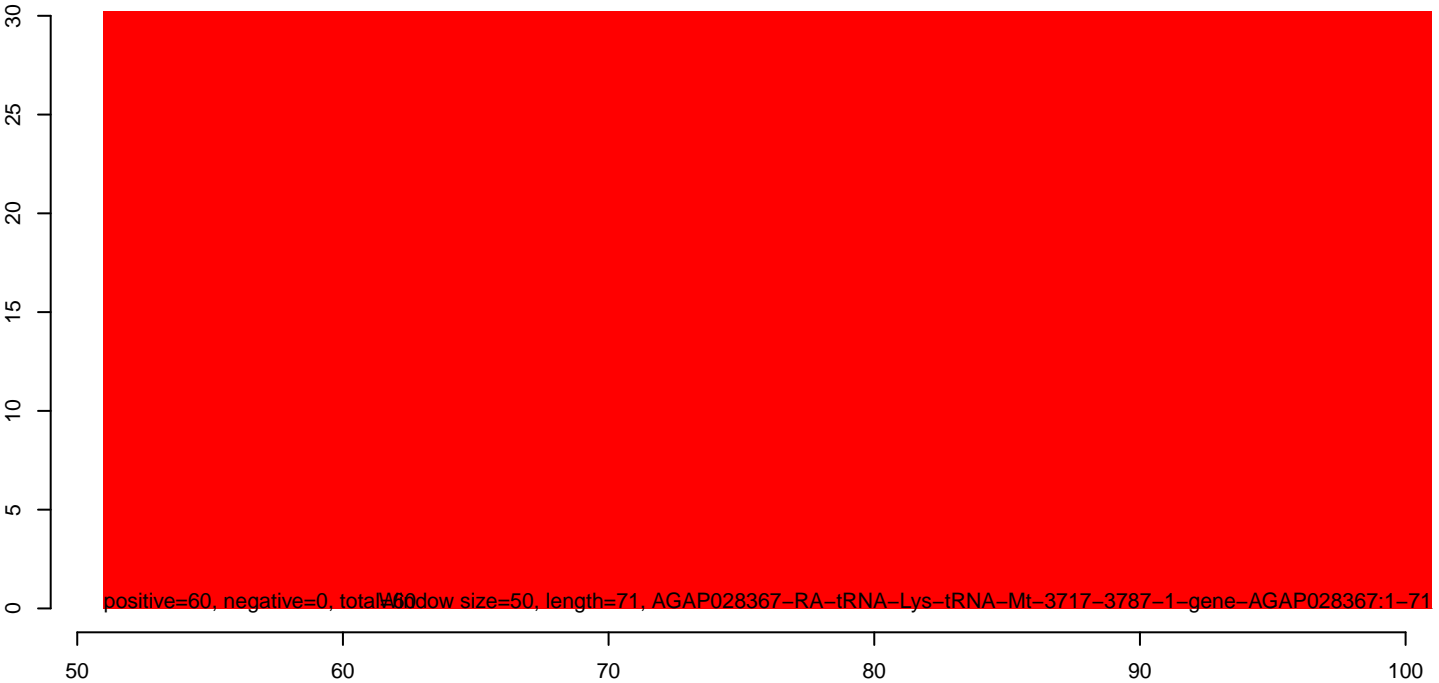
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



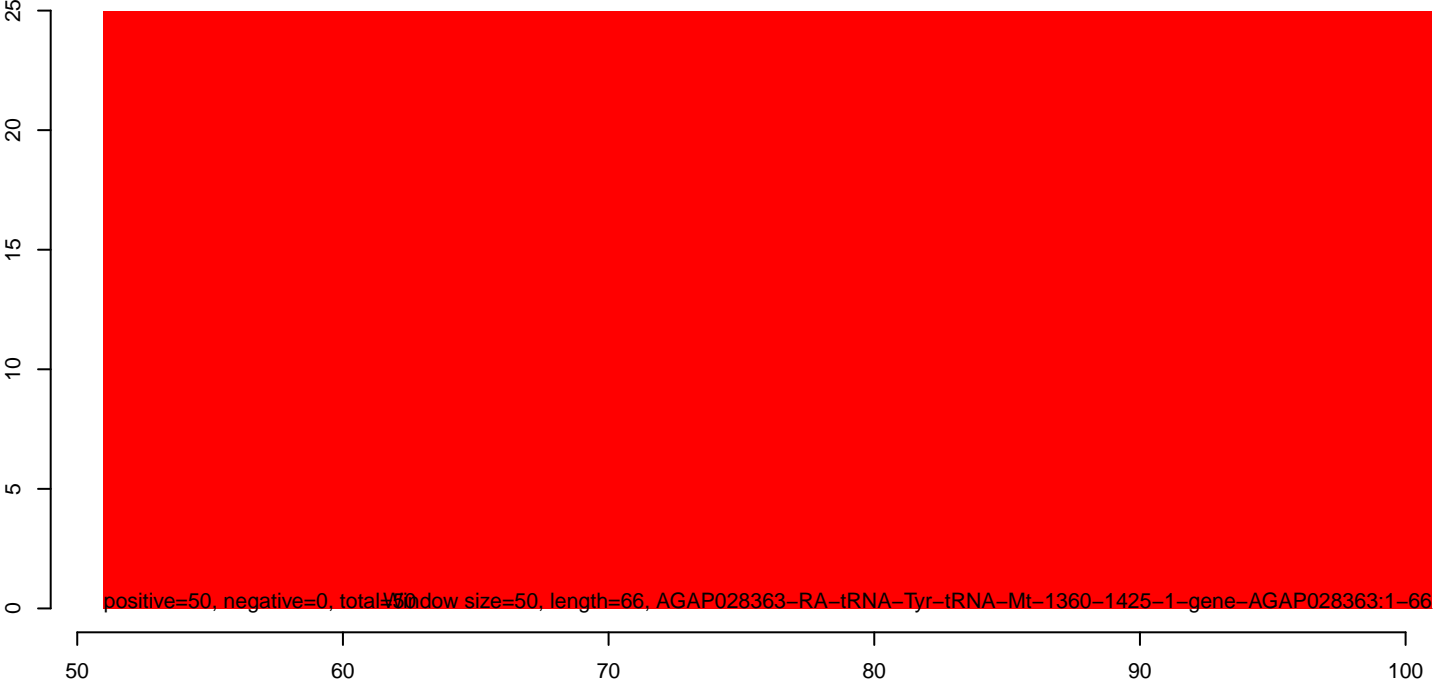
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep

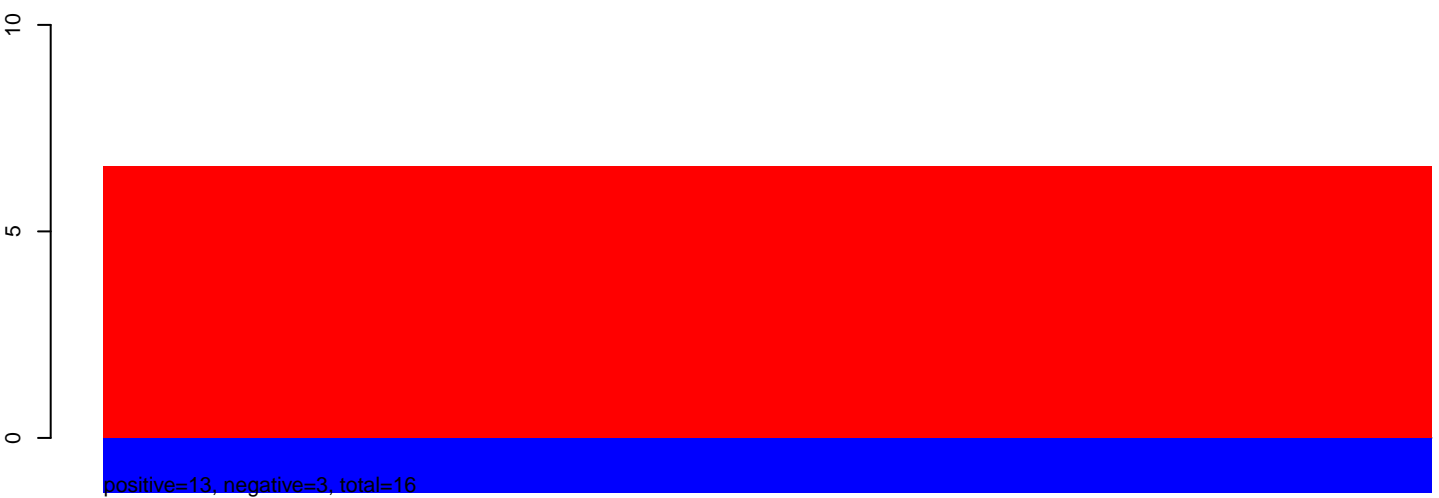




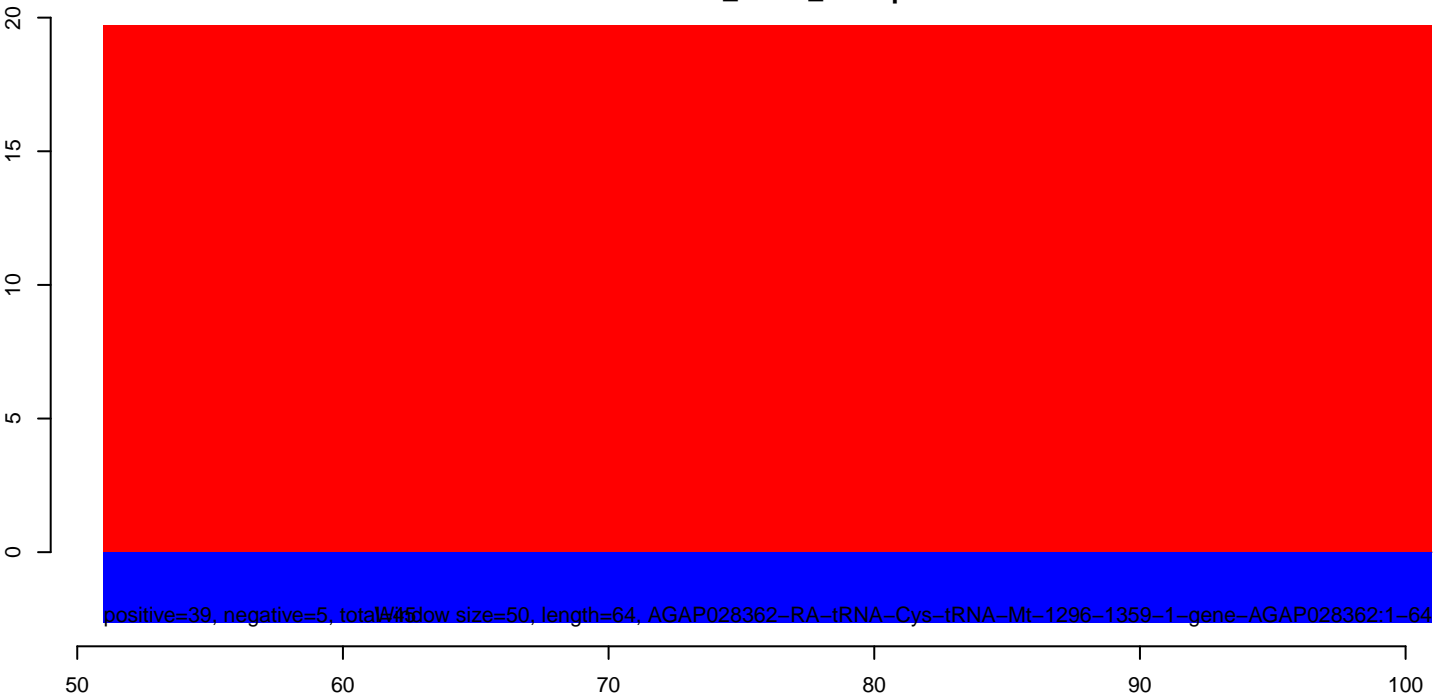
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



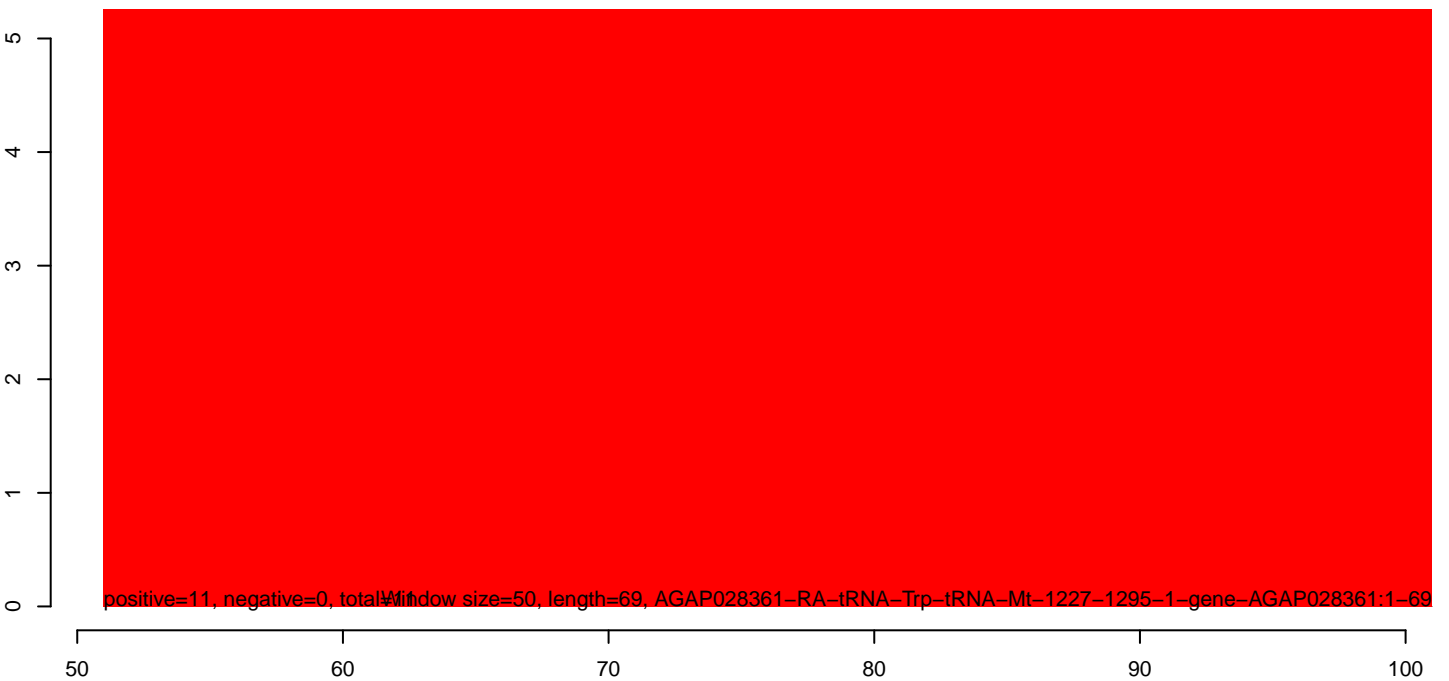
AnGam\_Fcarc\_TN.18\_23.rep



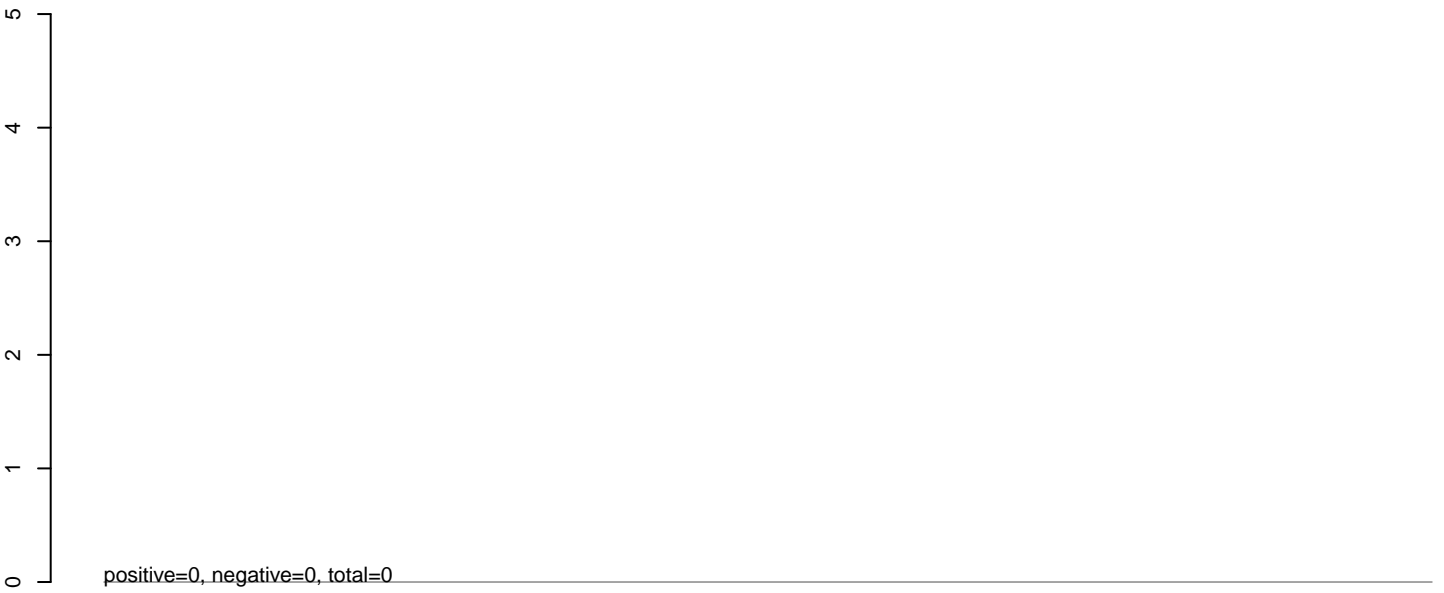
AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep

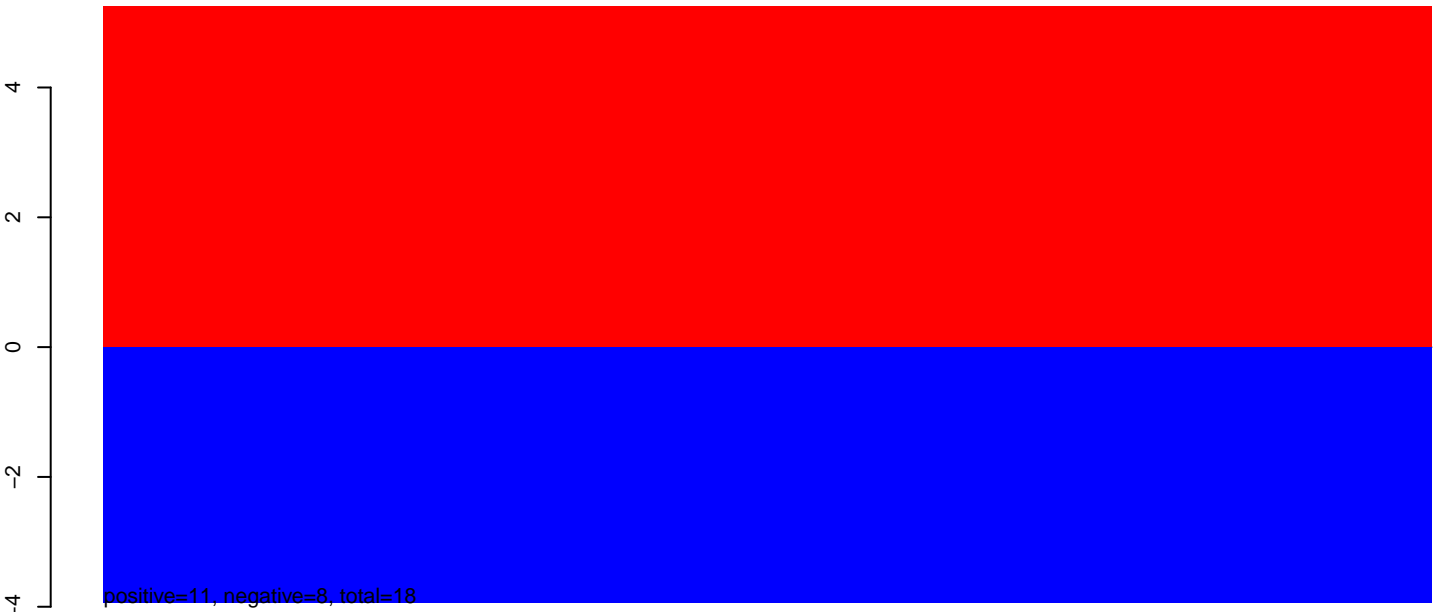


### AnGam\_Fcarc\_TN.rep



50 60 70 80 90 100

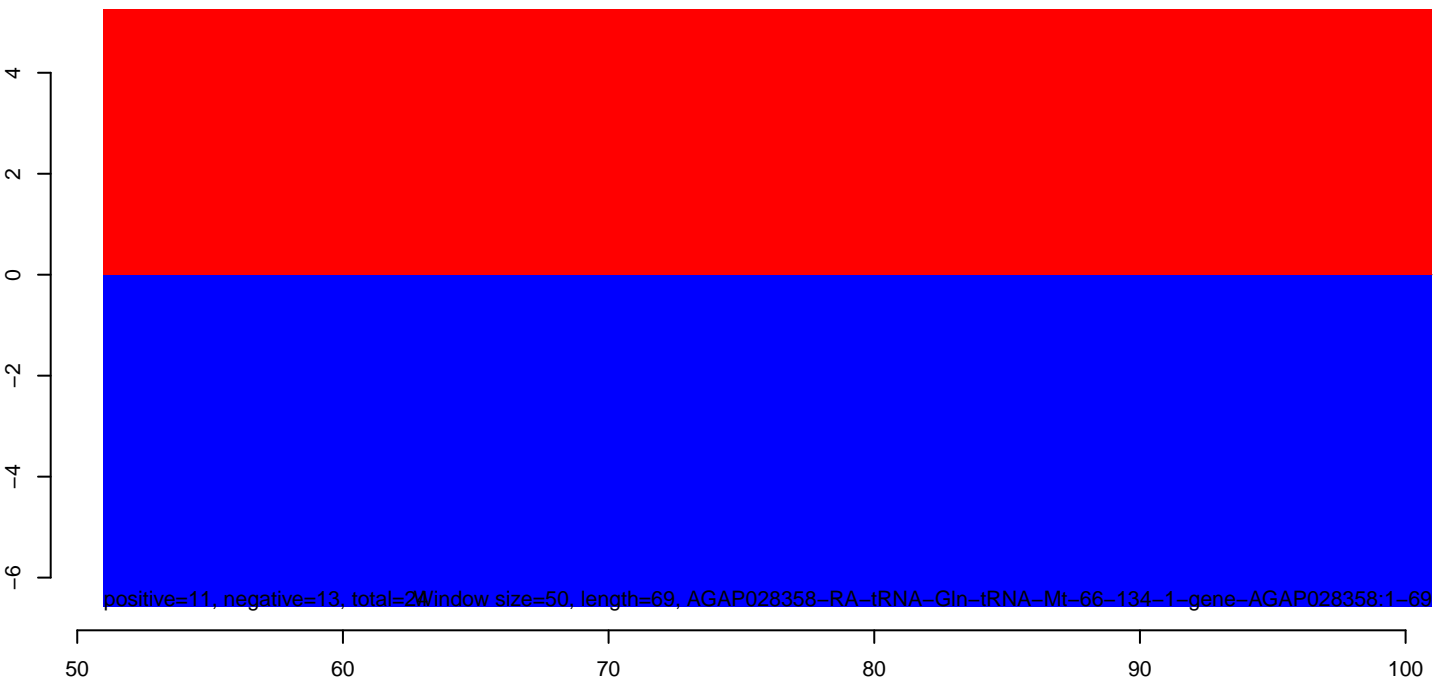
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



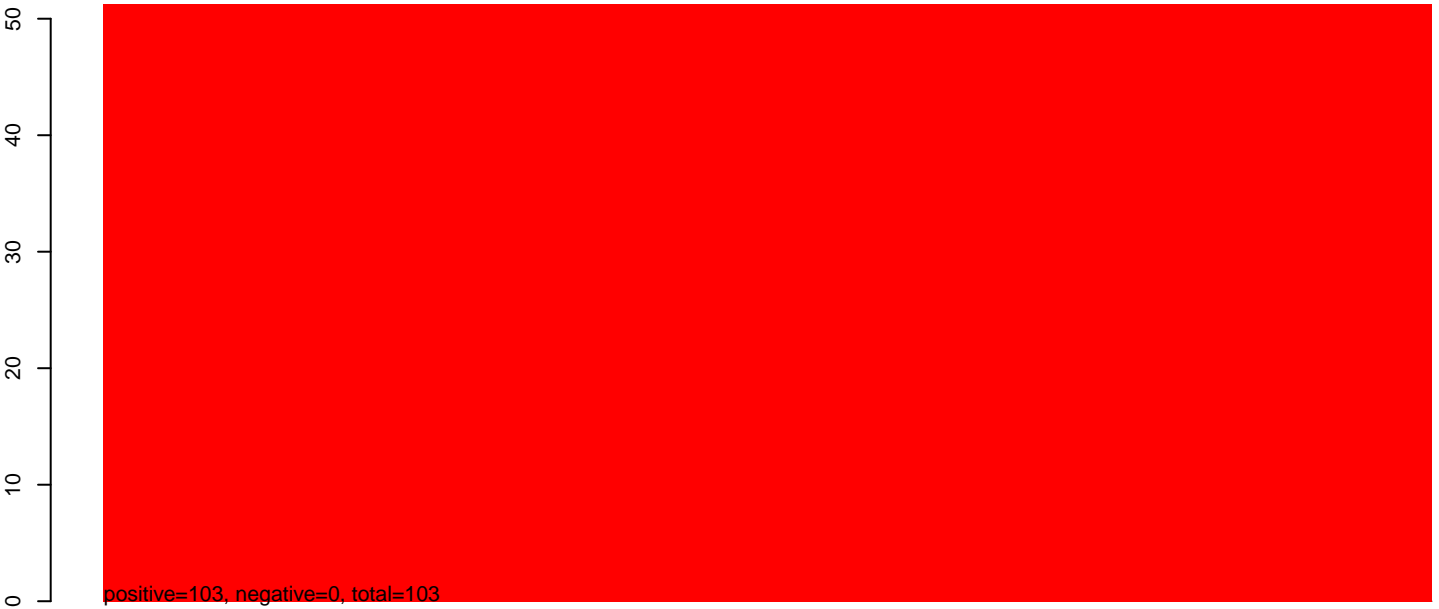
AnGam\_Fcarc\_TN.rep



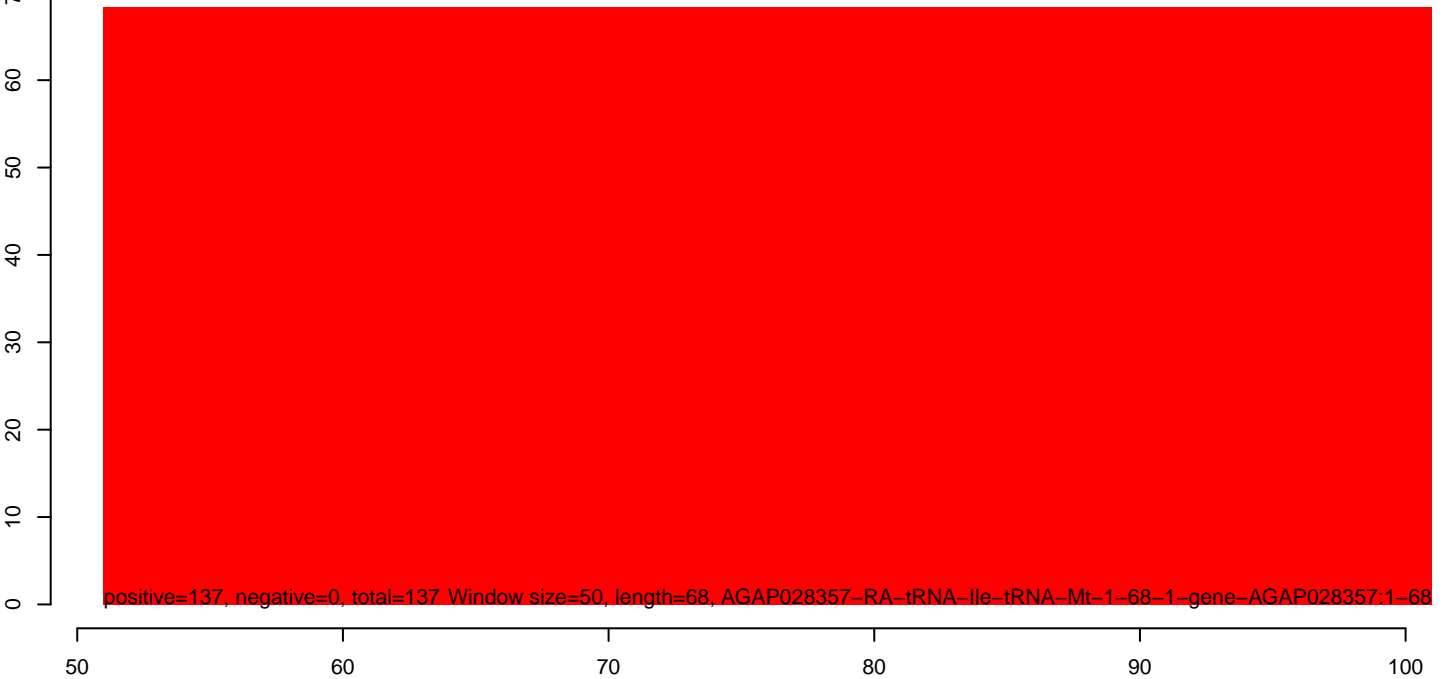
### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep



### AnGam\_Fcarc\_TN.rep



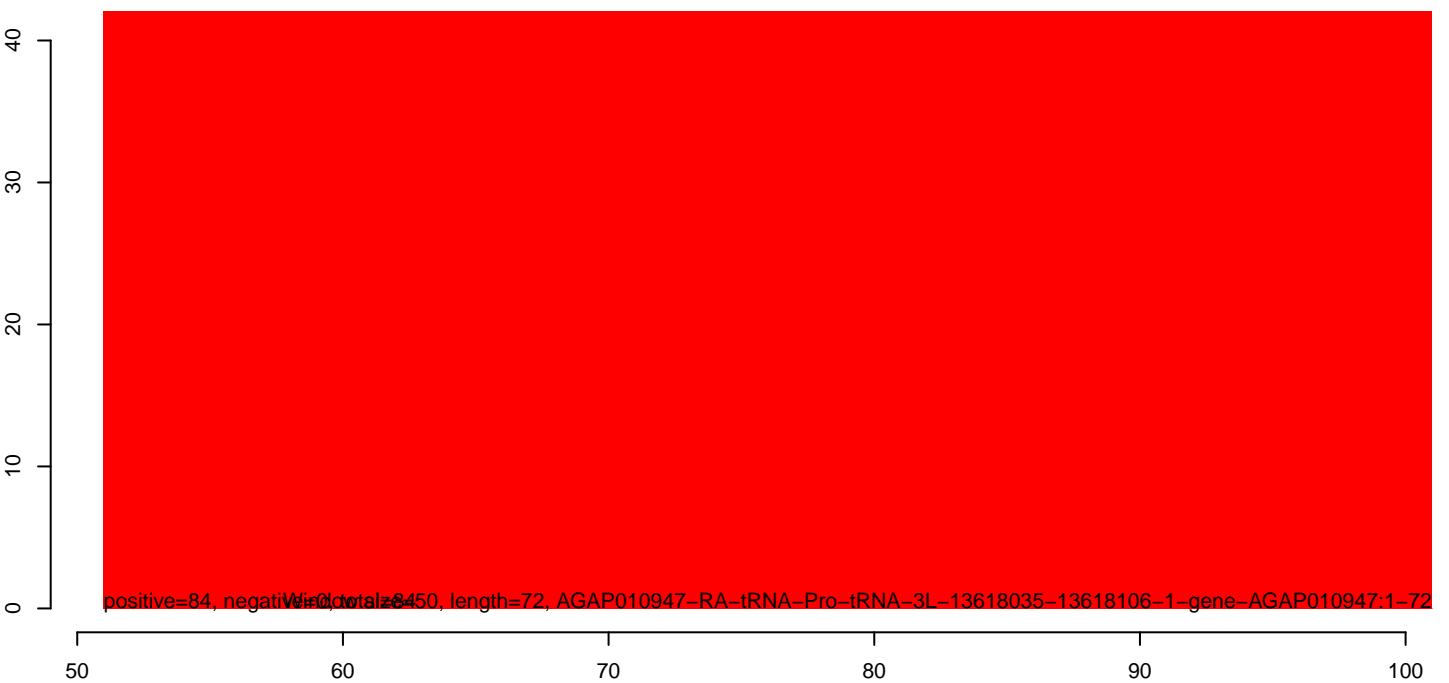
### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep



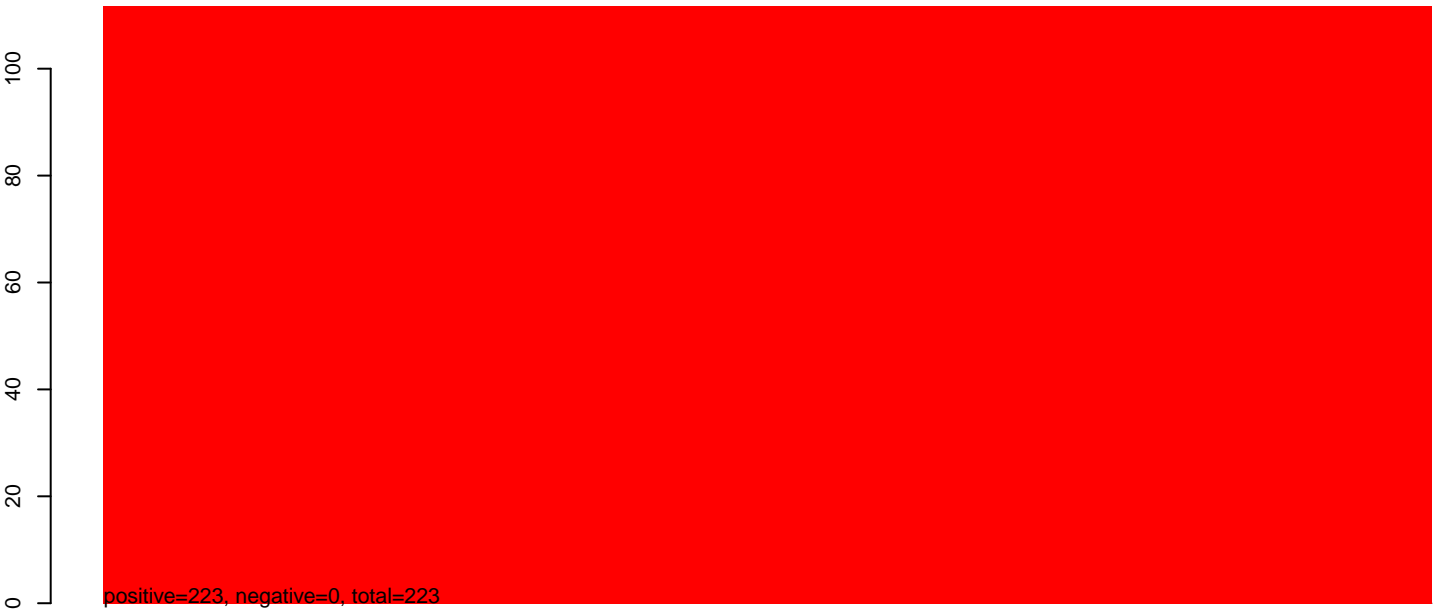
### AnGam\_Fcarc\_TN.rep



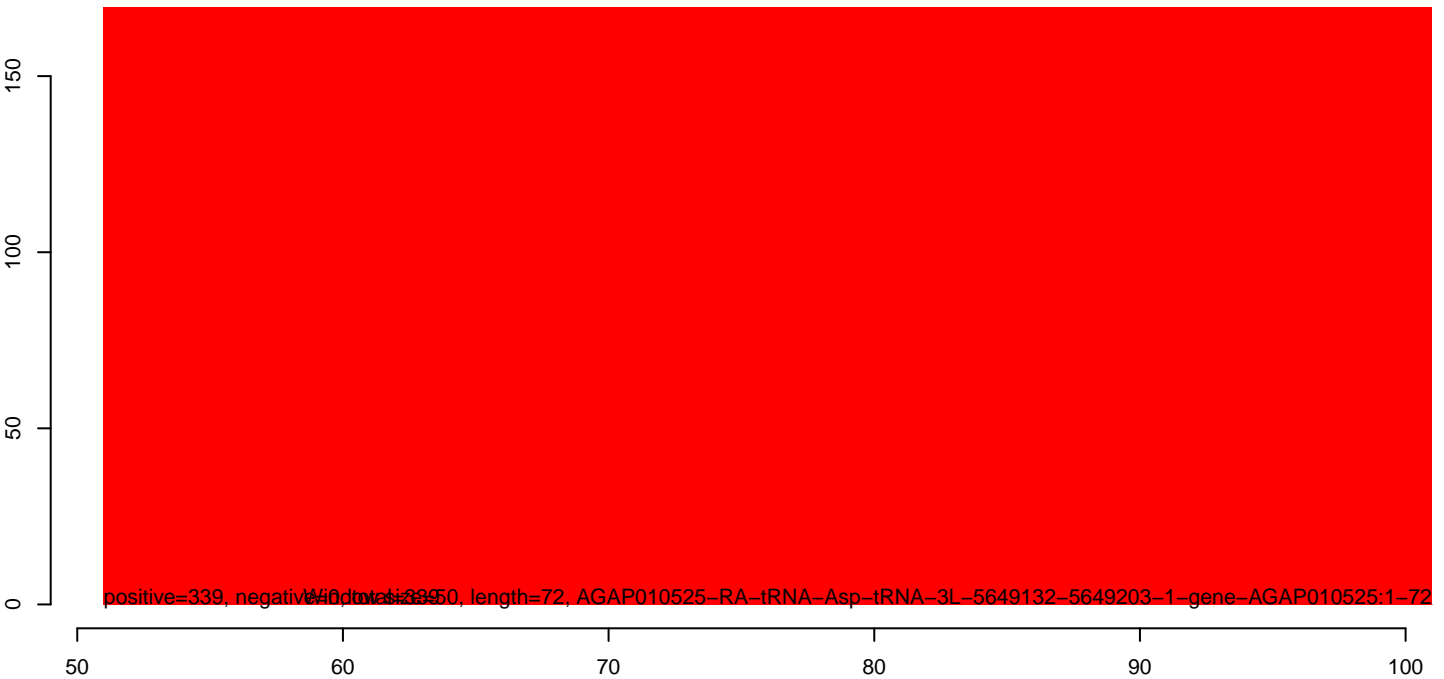
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



50 60 70 80 90 100



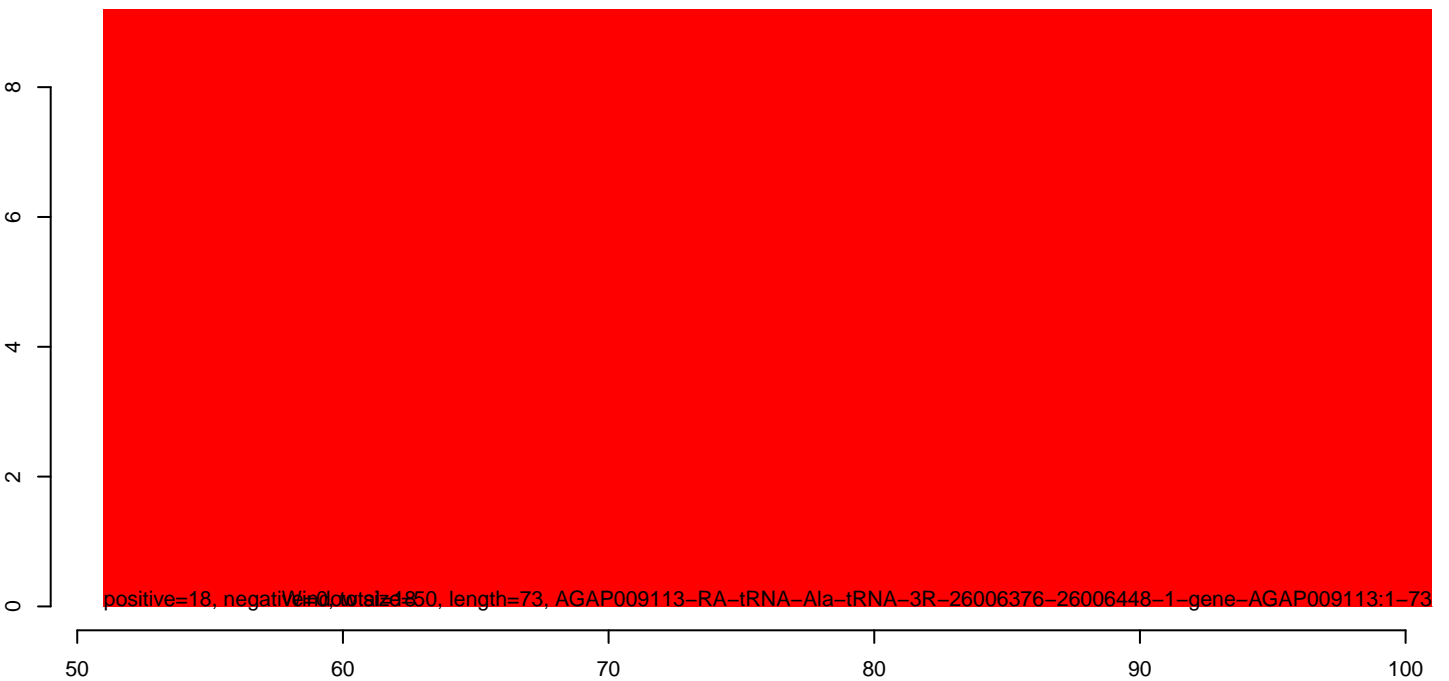
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



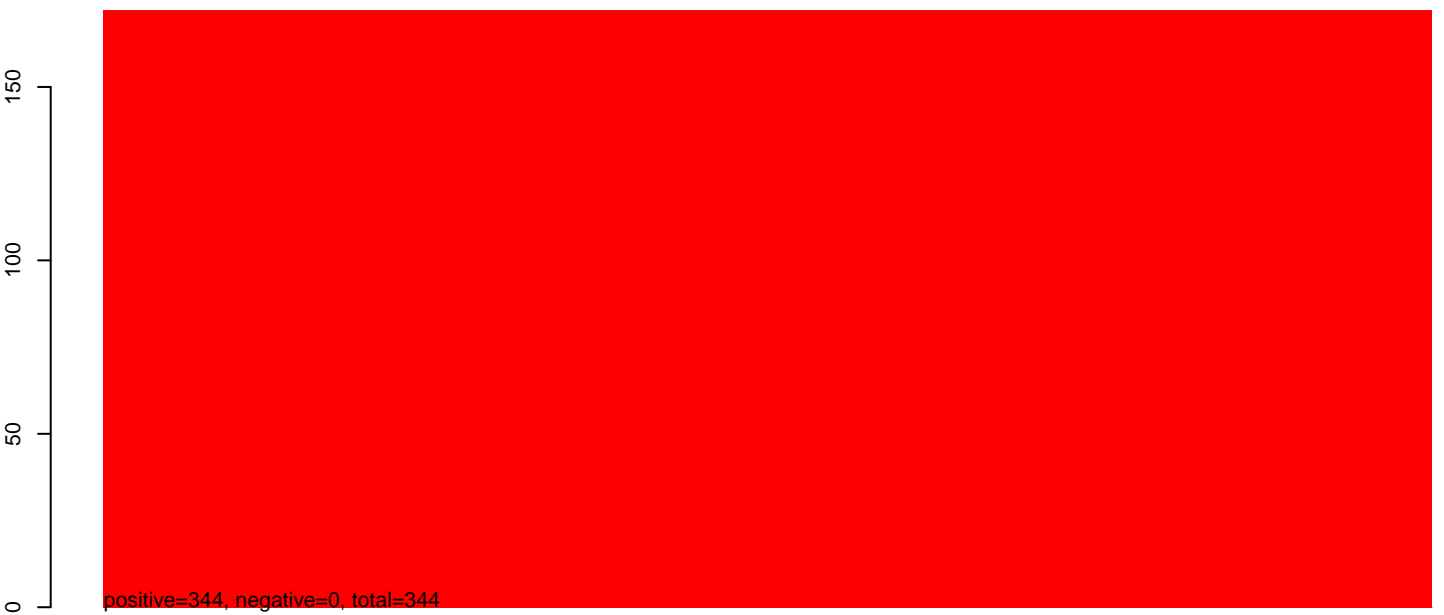
AnGam\_Fcarc\_TN.rep



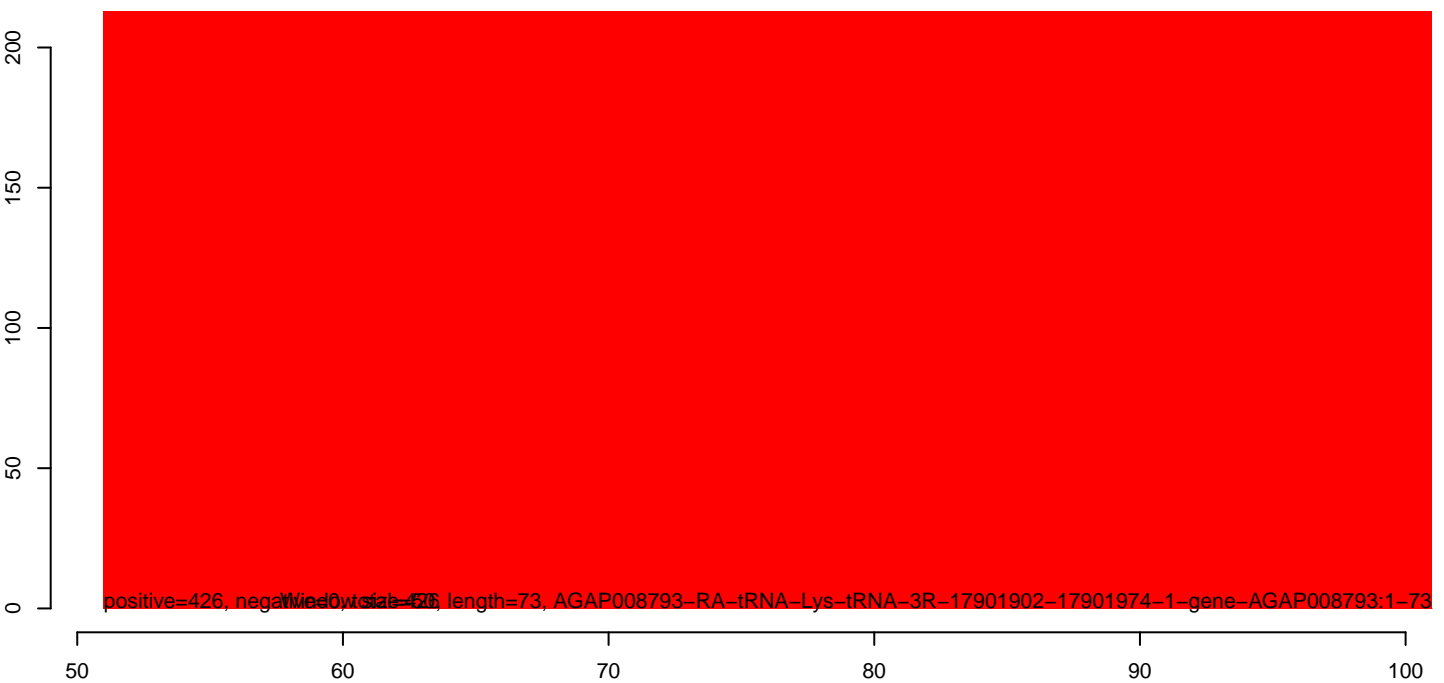
### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep



### AnGam\_Fcarc\_TN.rep



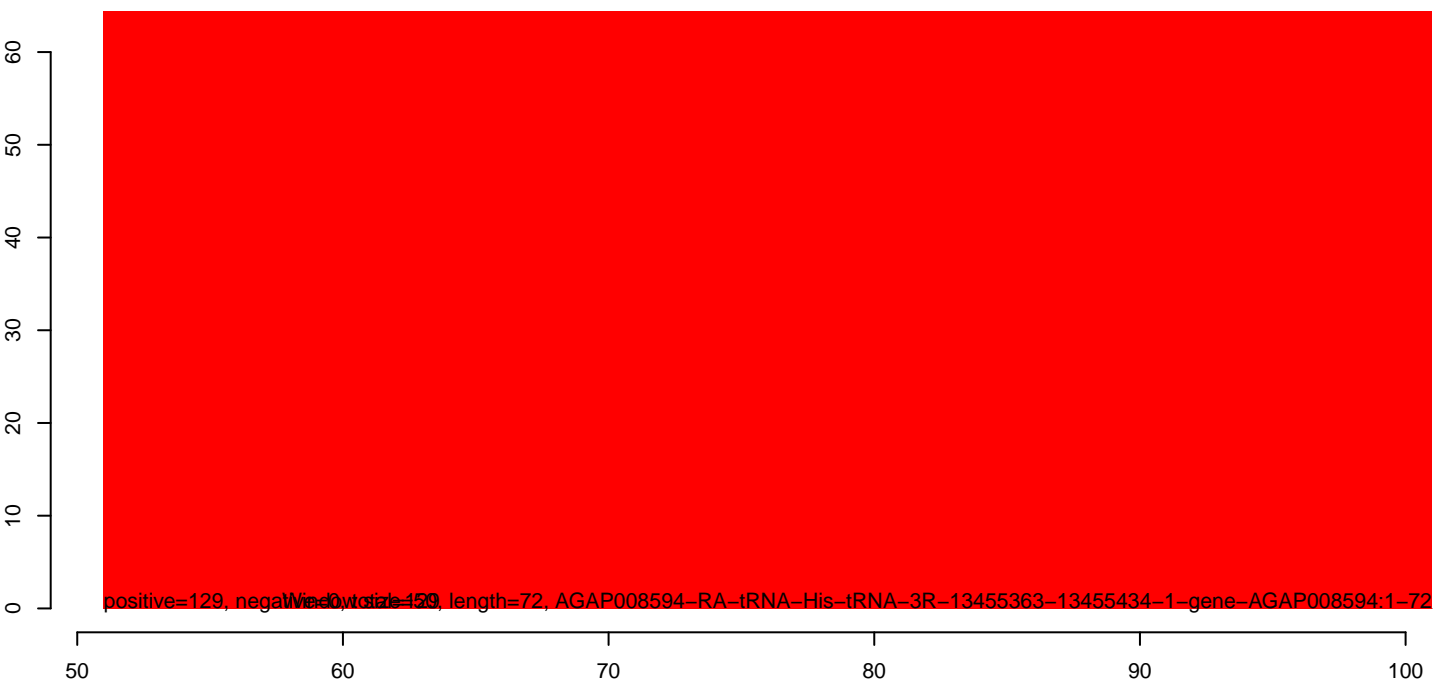
### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep



### AnGam\_Fcarc\_TN.rep

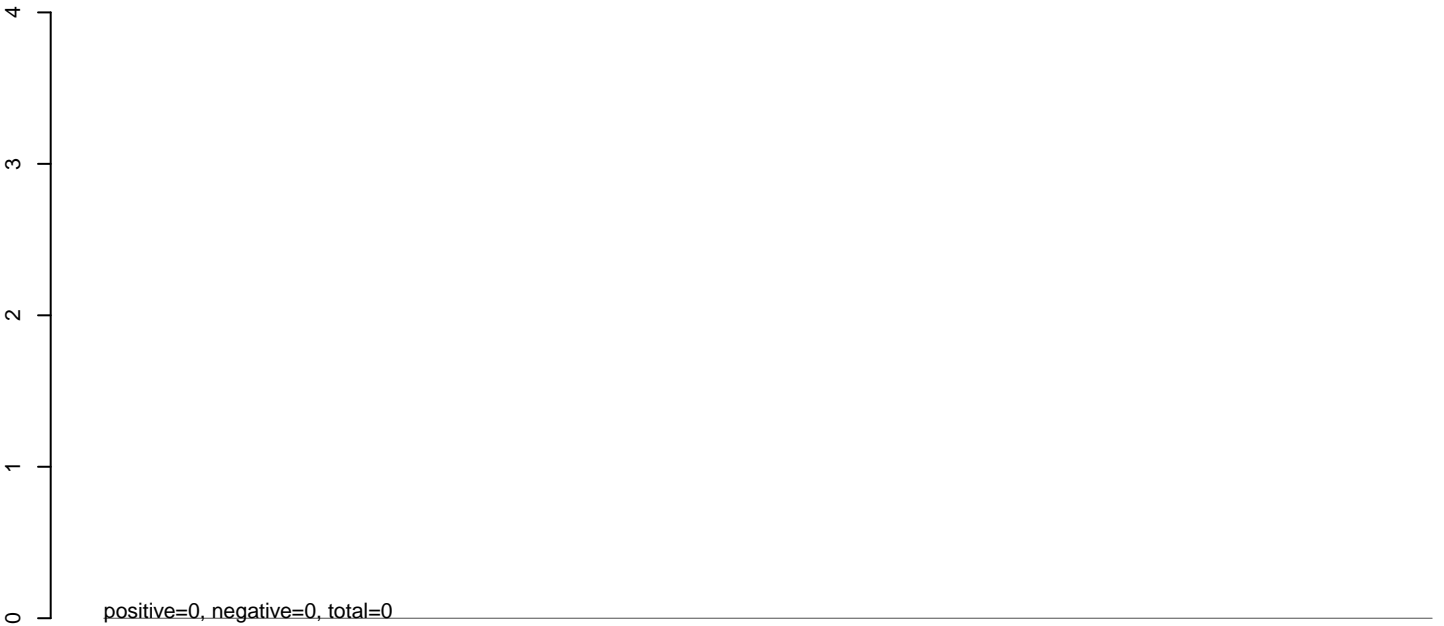


Window size=150, length=72, AGAP008594-RA-tRNA-His-tRNA-3R-13455363-13455434-1-gene-AGAP008594:1-72

AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



Window size=50, length=74, AGAP008069-RA-tRNA-Thr-tRNA-3R-5039303-5039376-1-gene-AGAP008069:1-74

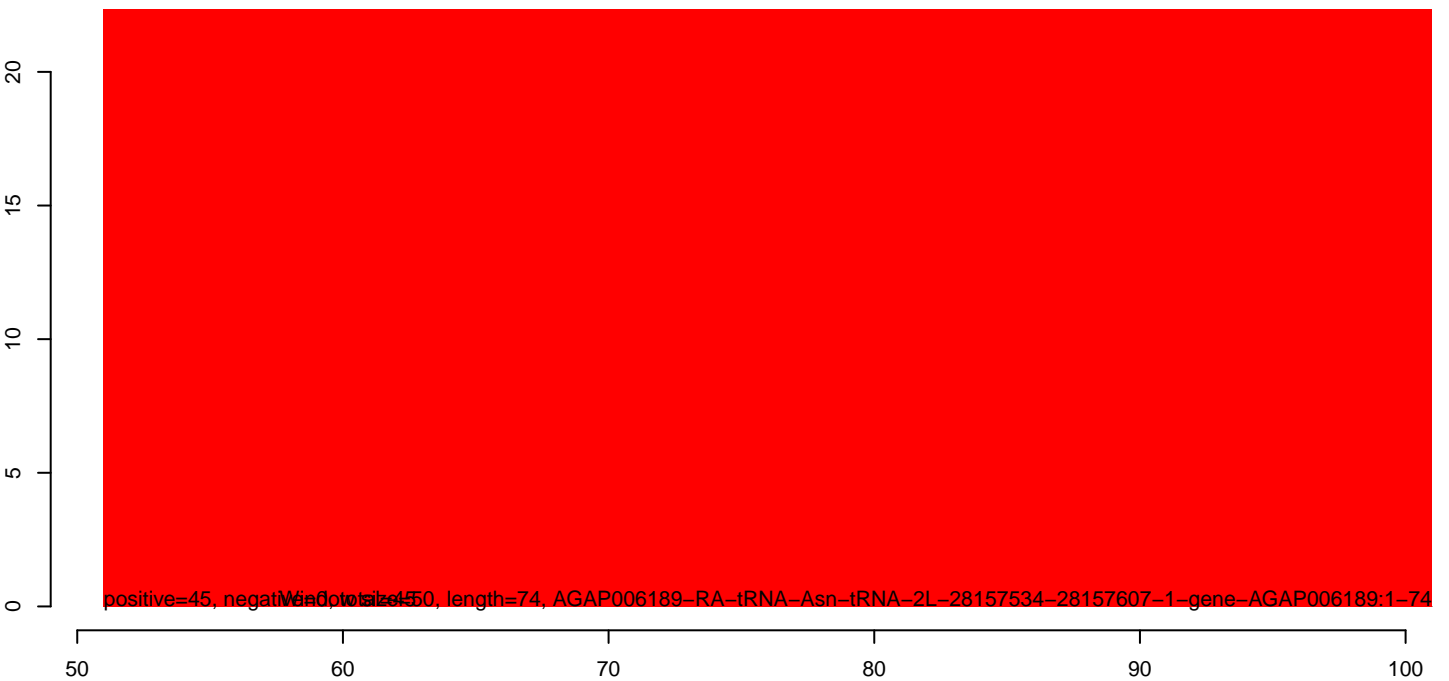
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



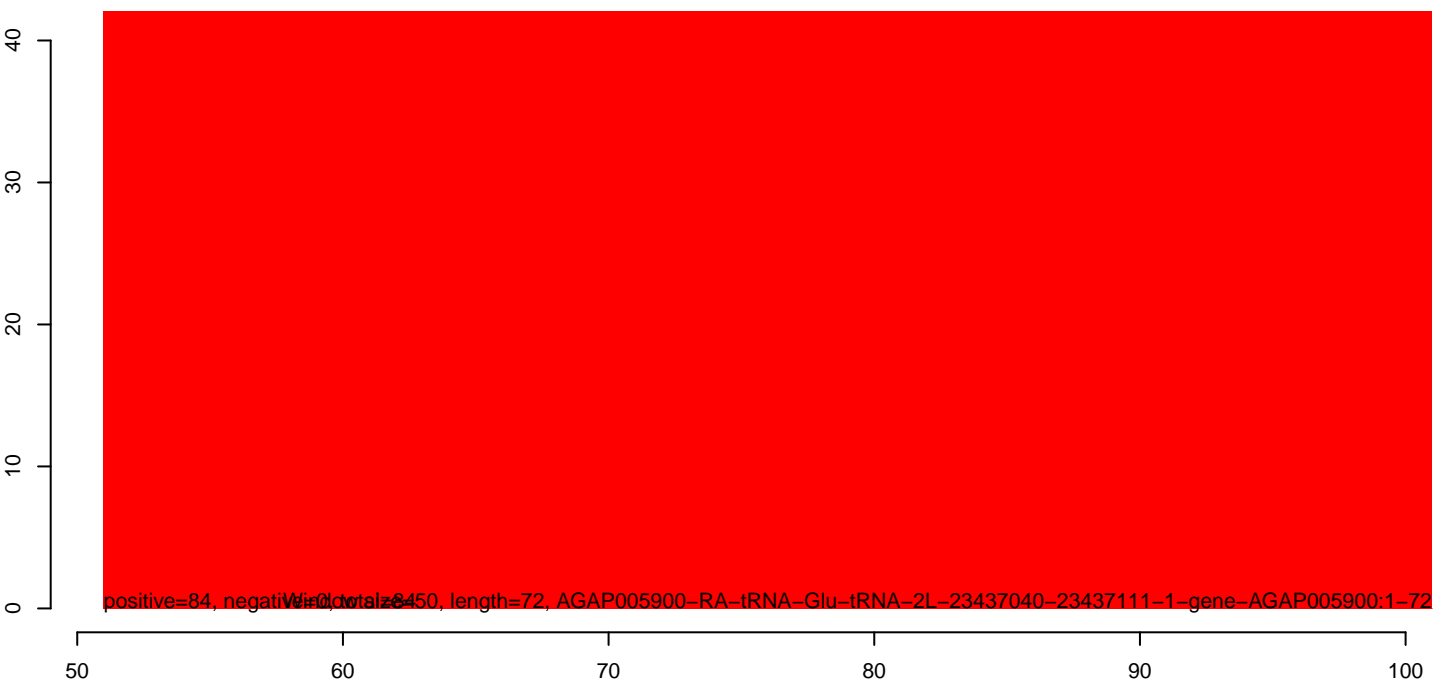
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



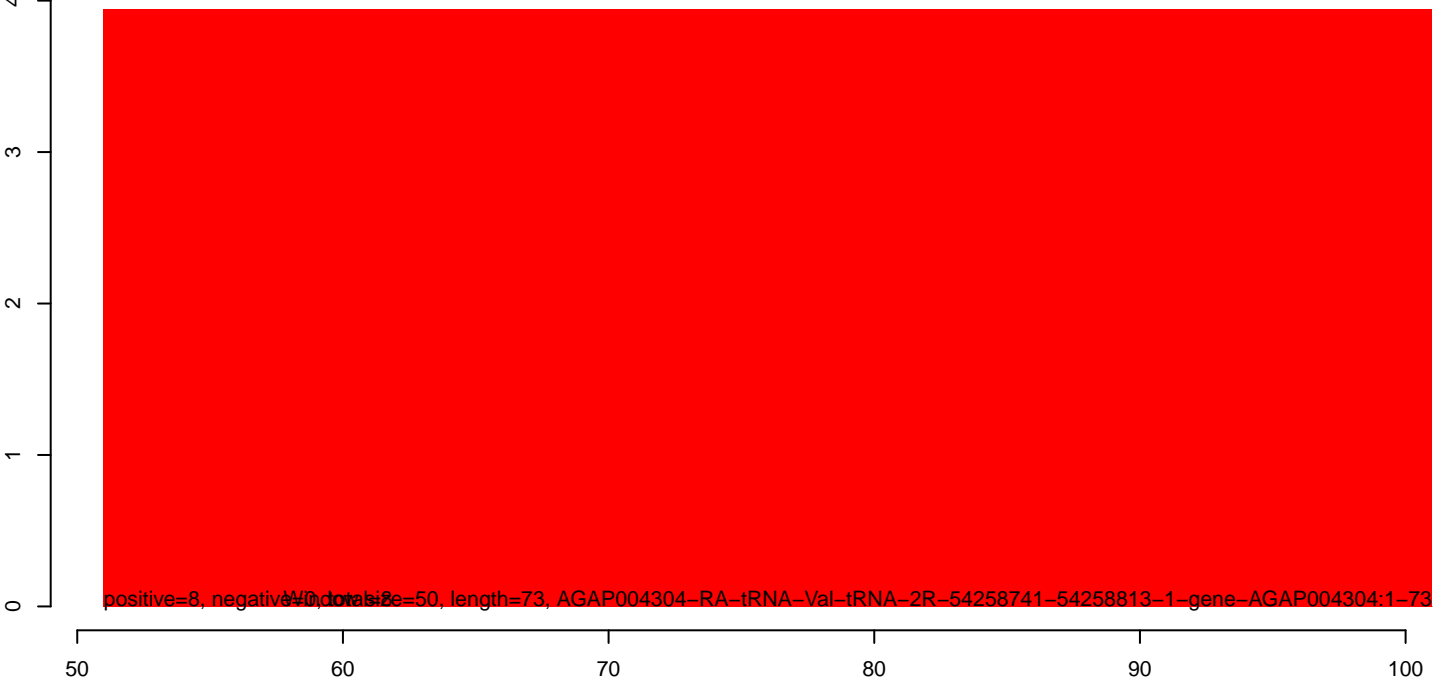
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



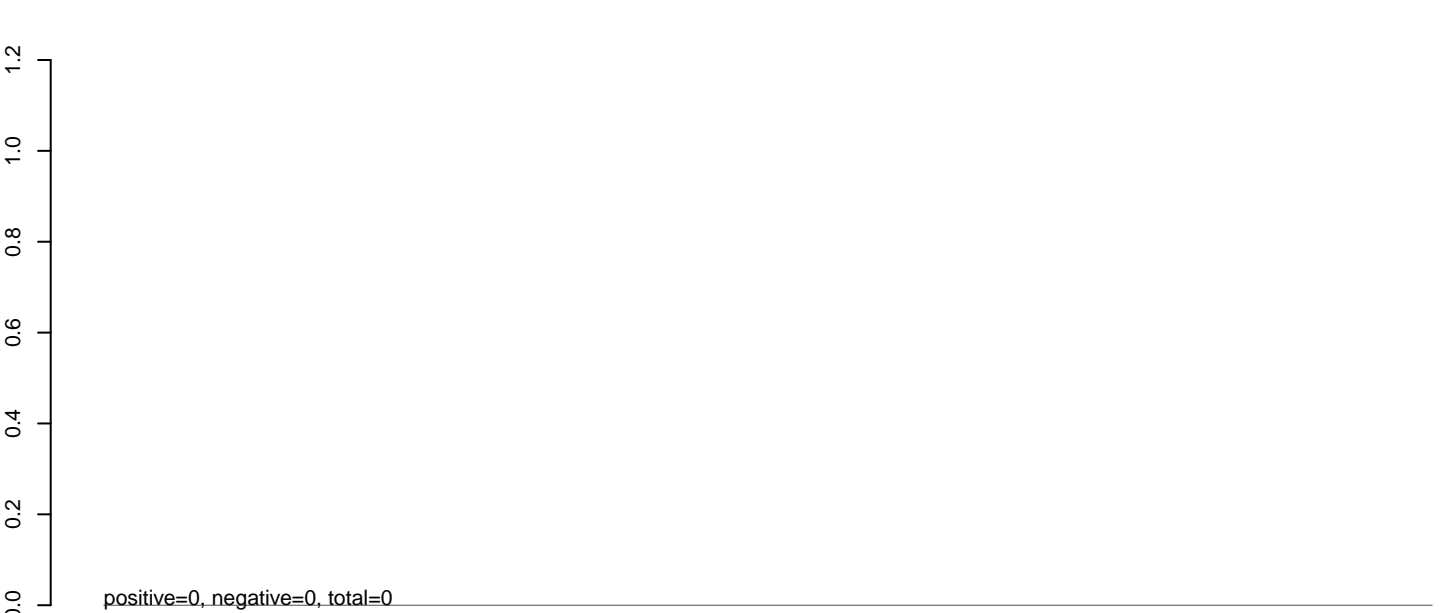
AnGam\_Fcarc\_TN.rep



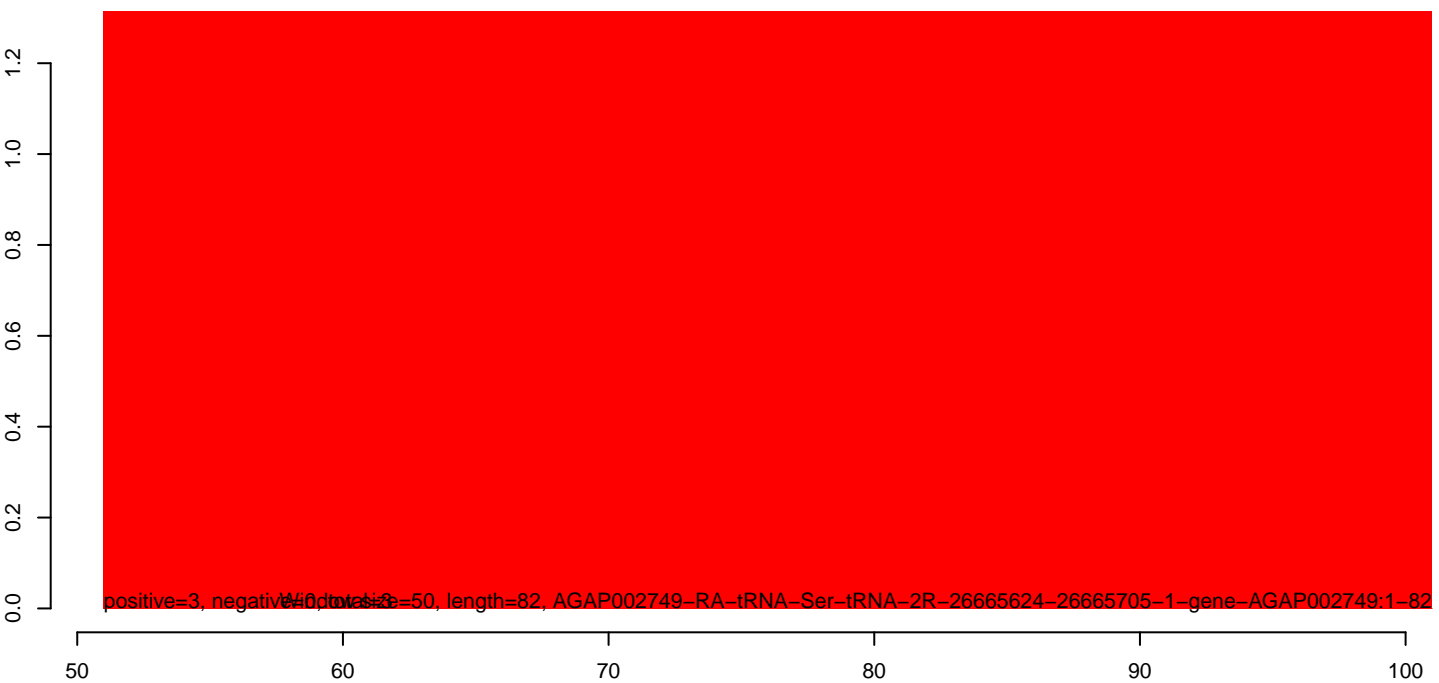
### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep

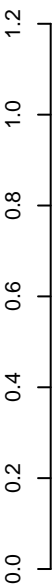


### AnGam\_Fcarc\_TN.rep



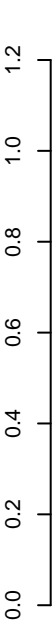


# AnGam\_Fcarc\_TN.18\_23.rep



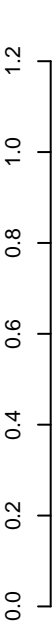
positive=0, negative=0, total=0

# AnGam\_Fcarc\_TN.24\_35.rep

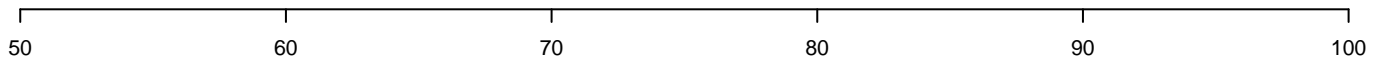


positive=3, negative=0, total=3

# AnGam\_Fcarc\_TN.rep



positive=3, negative=0, total=3, width=50, length=83, AGAP002700-RA-tRNA-Leu-tRNA-2R-25723627-25723709-1-gene-AGAP002700:1-83



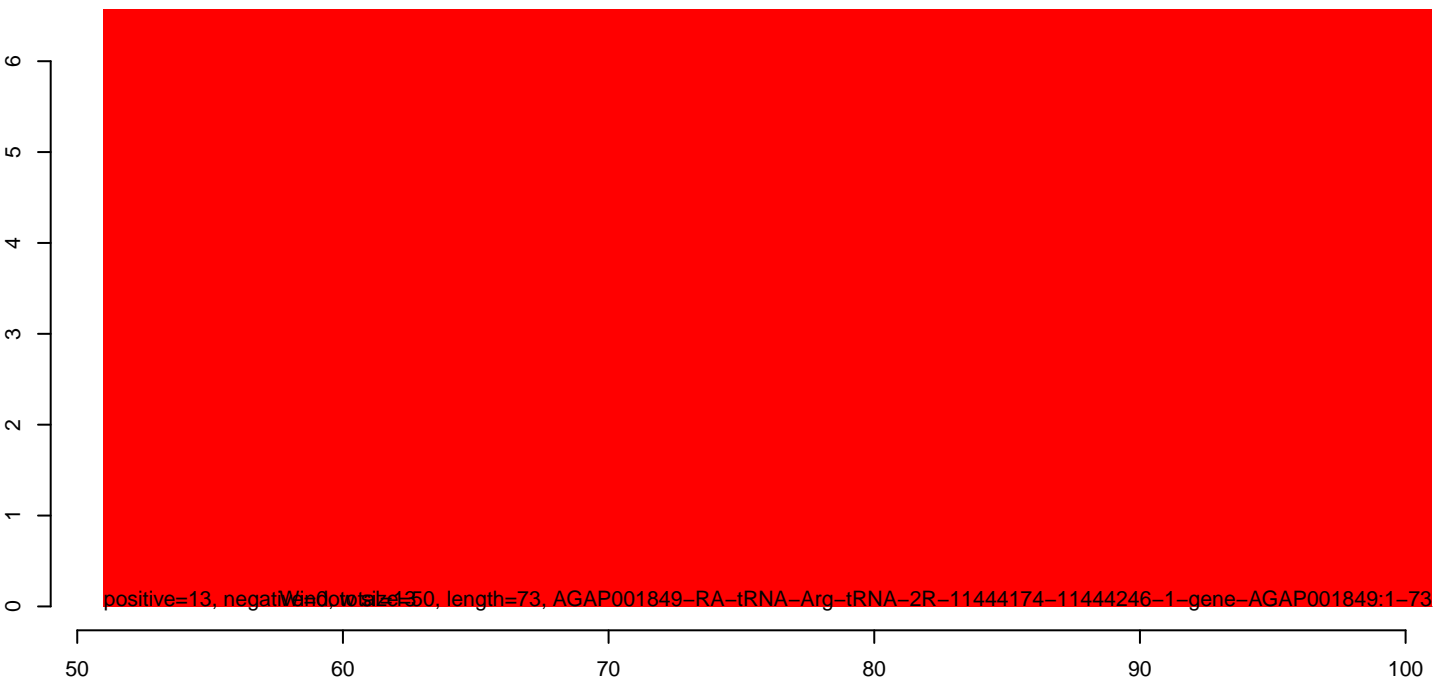
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



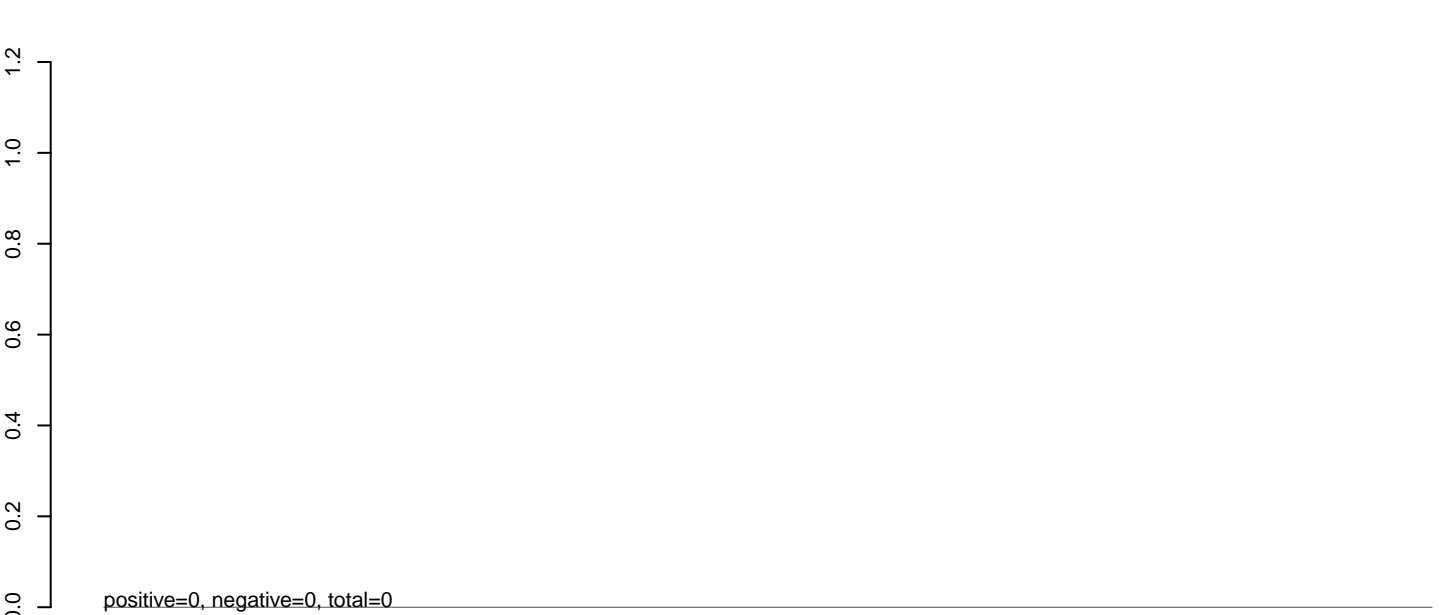
AnGam\_Fcarc\_TN.rep



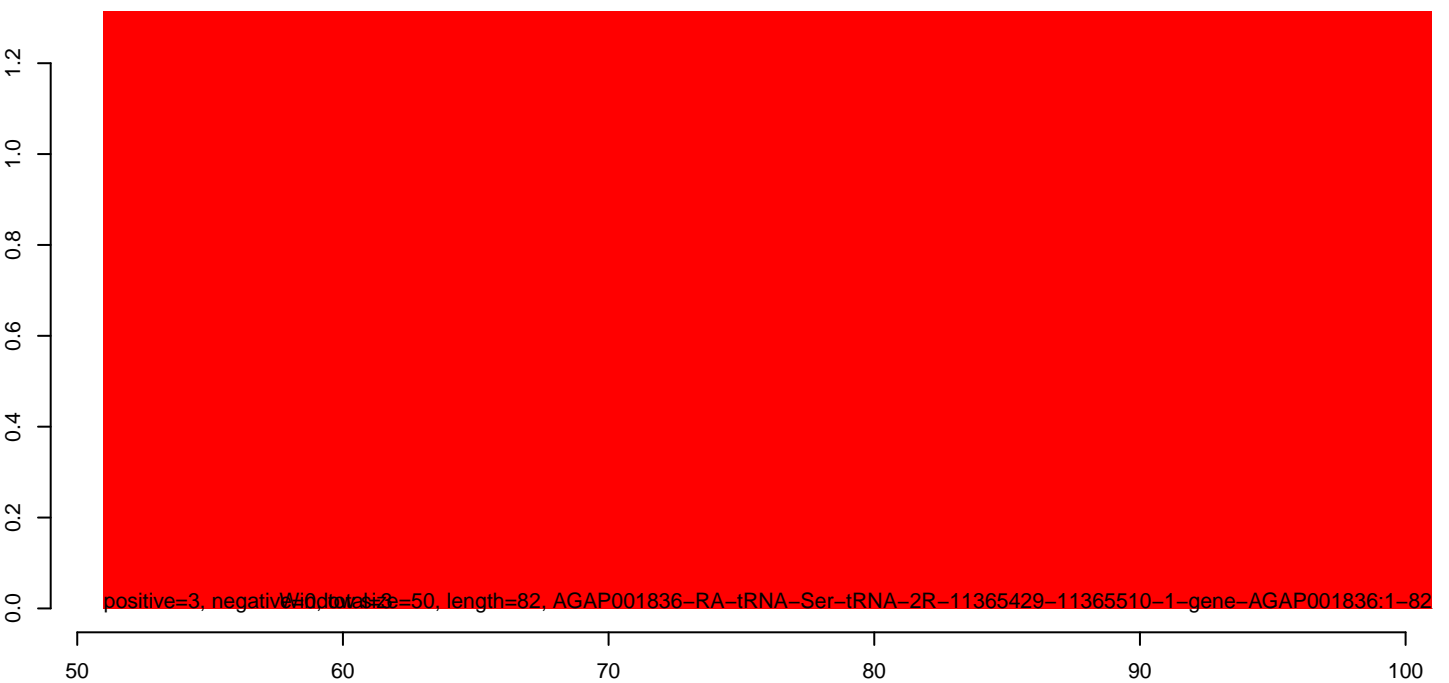
### AnGam\_Fcarc\_TN.18\_23.rep



### AnGam\_Fcarc\_TN.24\_35.rep



### AnGam\_Fcarc\_TN.rep



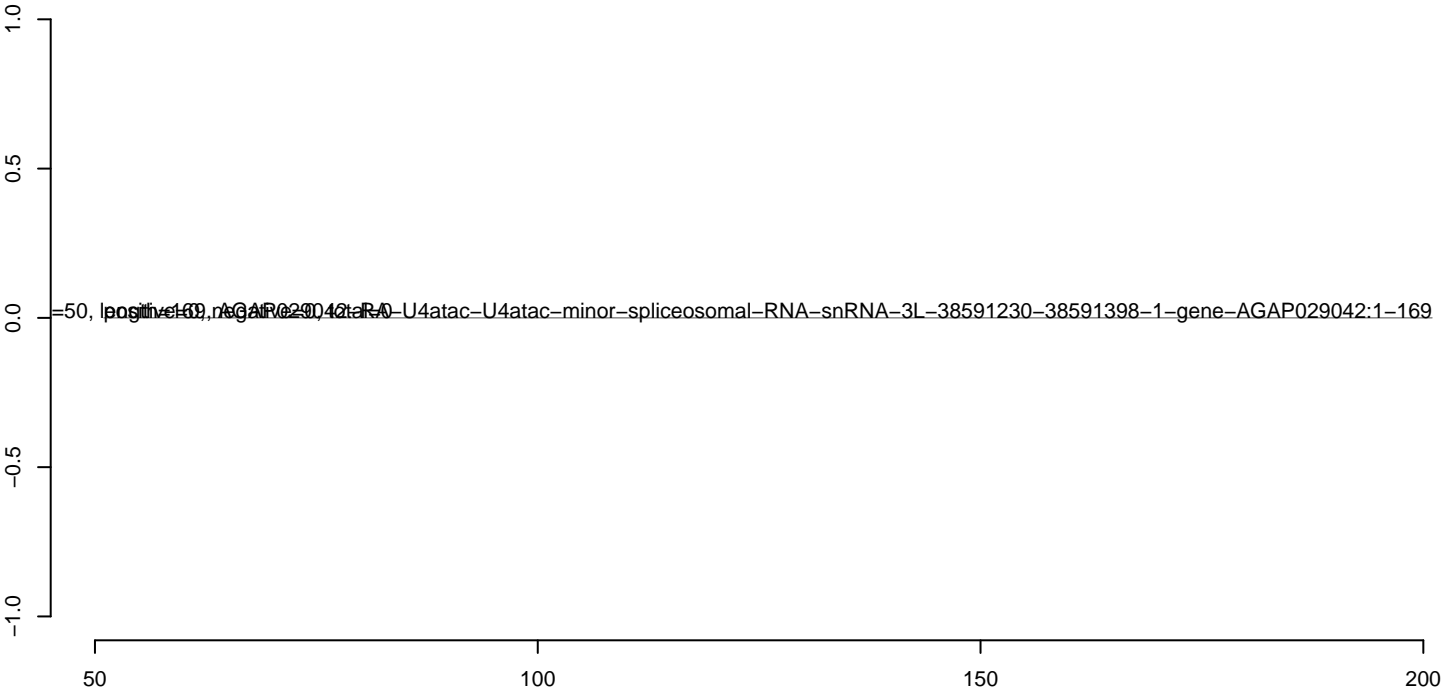
AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep



AnGam\_Fcarc\_TN.18\_23.rep



AnGam\_Fcarc\_TN.24\_35.rep



AnGam\_Fcarc\_TN.rep

