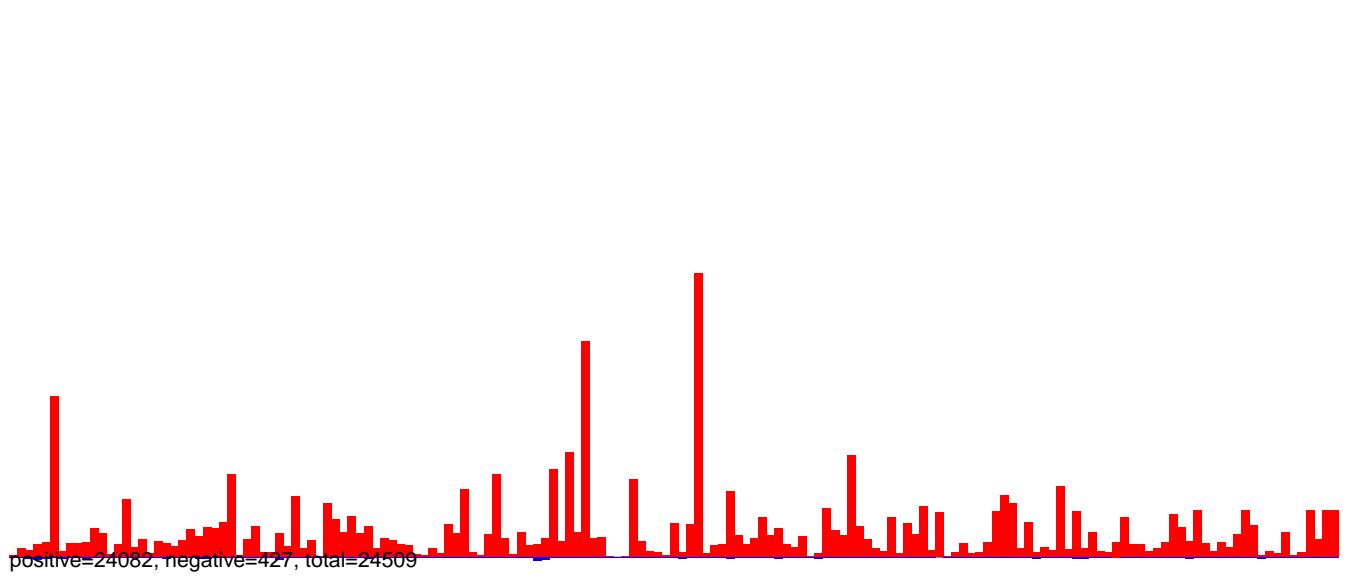
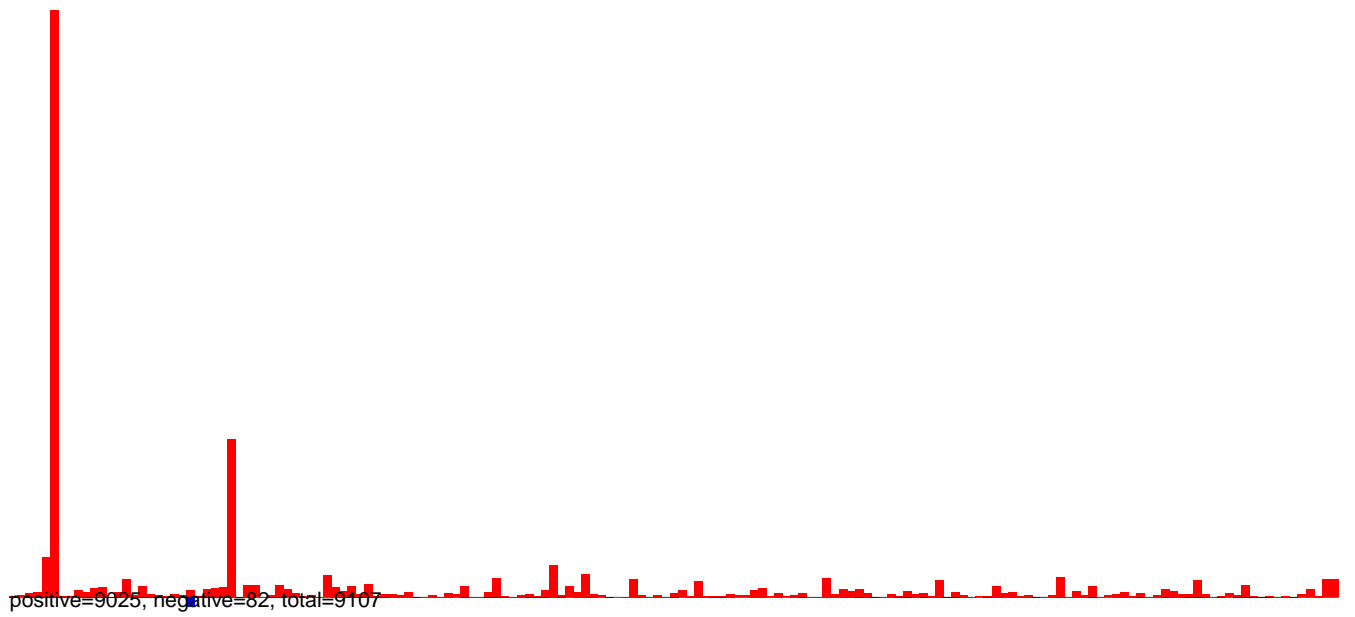


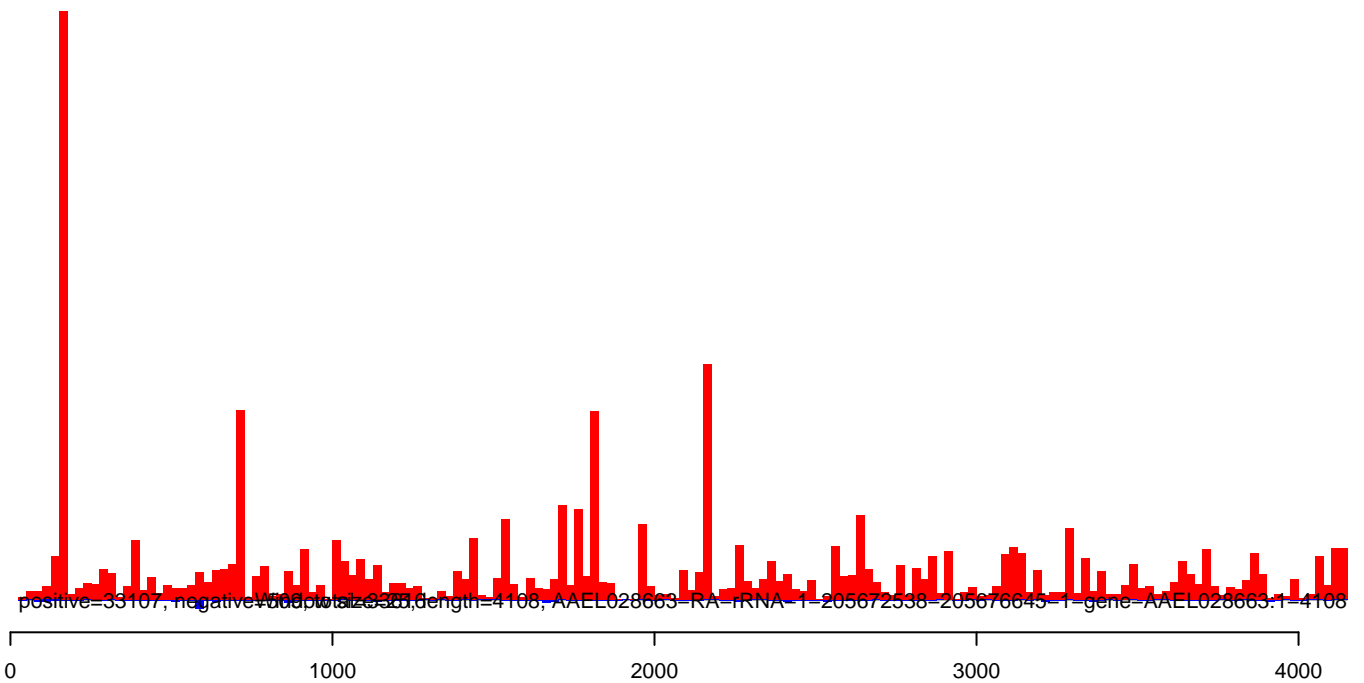
AeAeg_Testes.18_23.rep



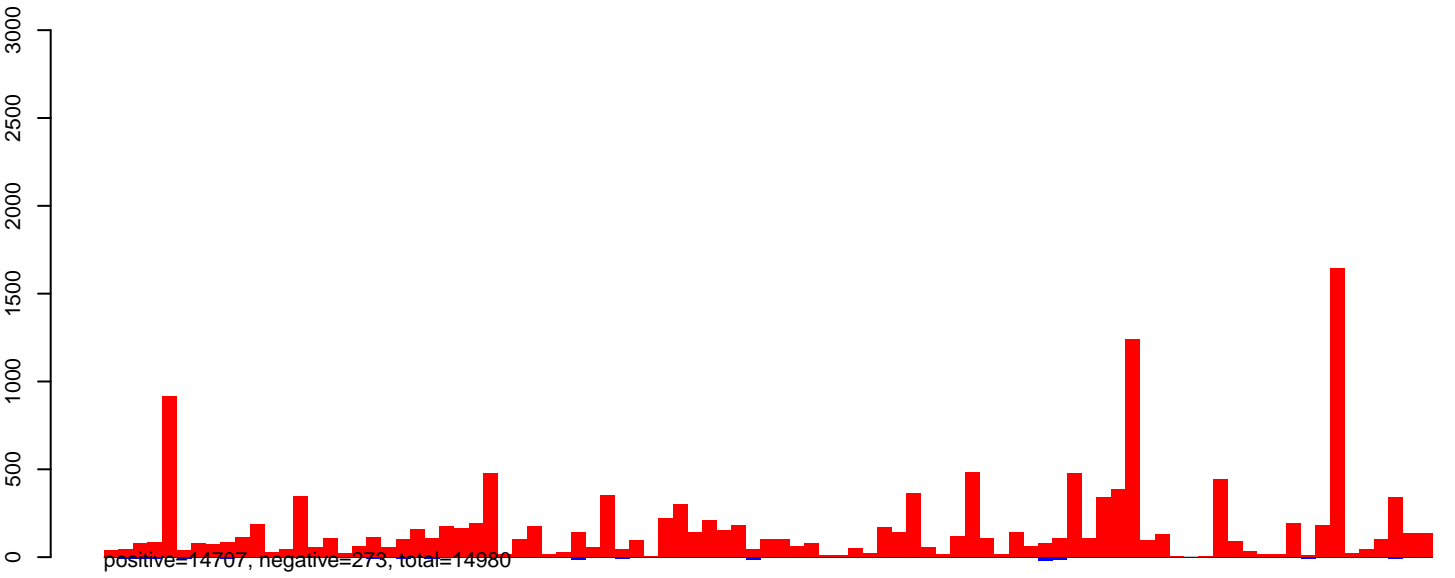
AeAeg_Testes.24_35.rep



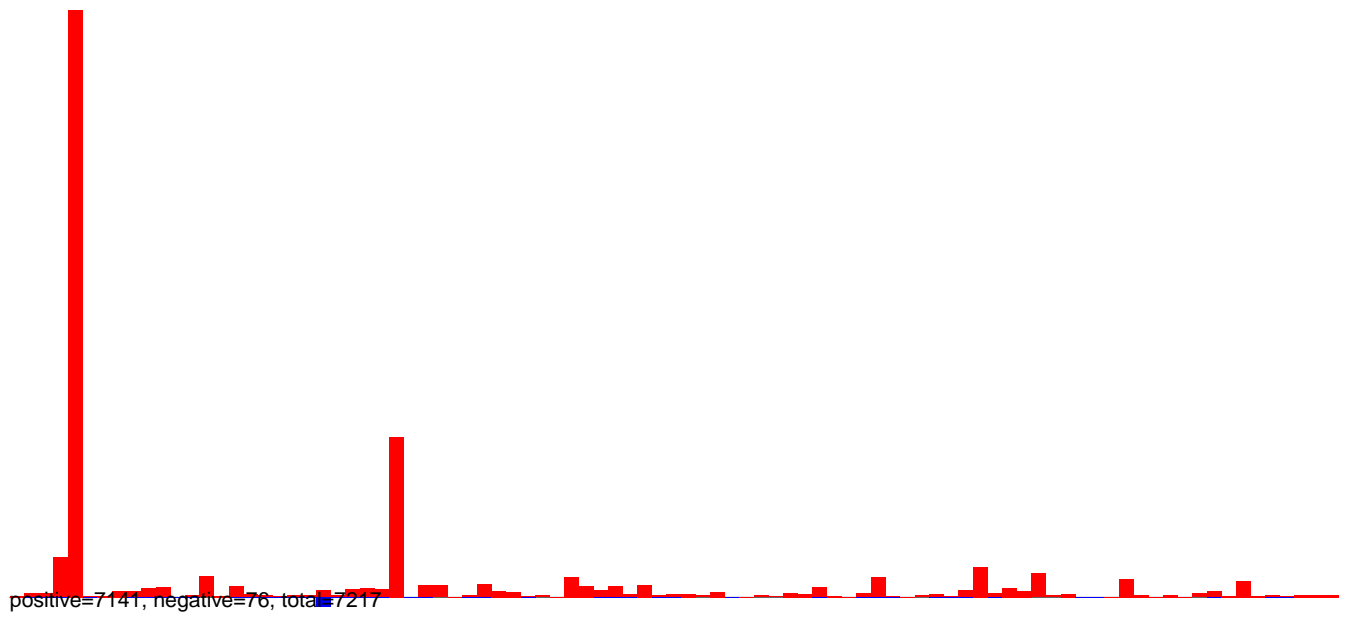
AeAeg_Testes.rep



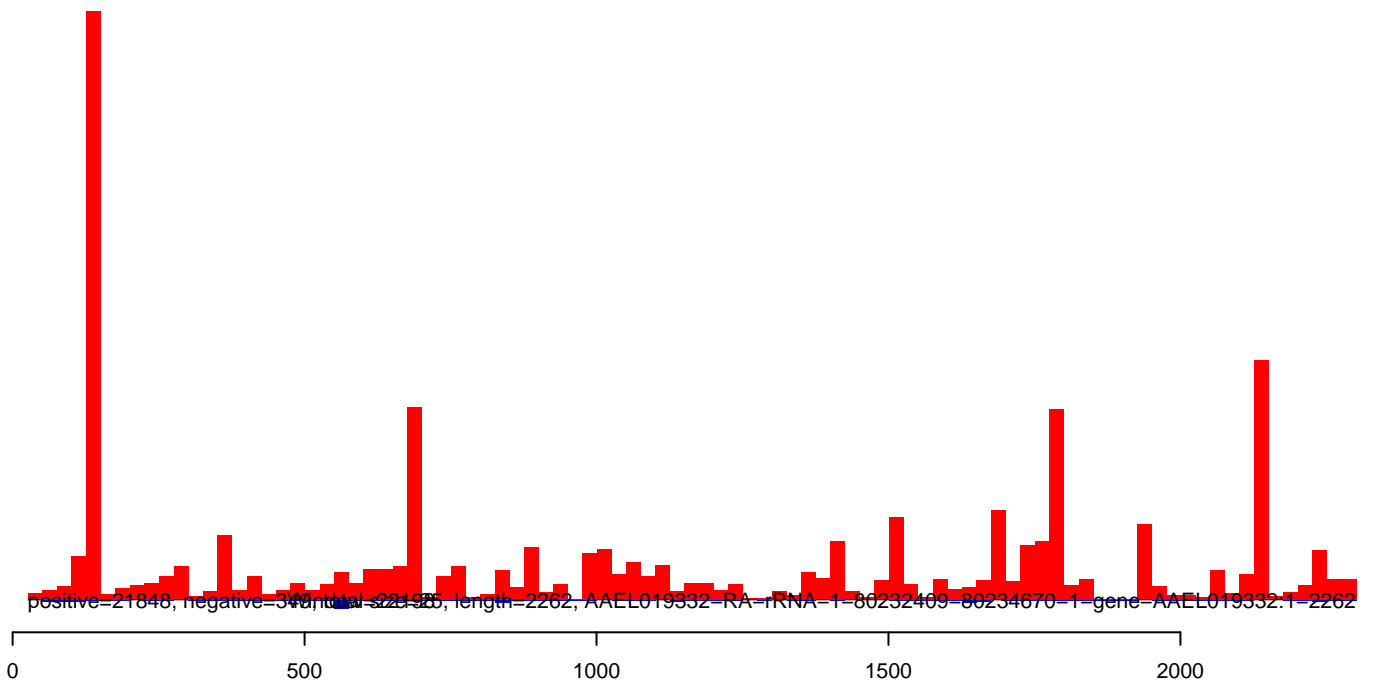
AeAeg_Testes.18_23.rep



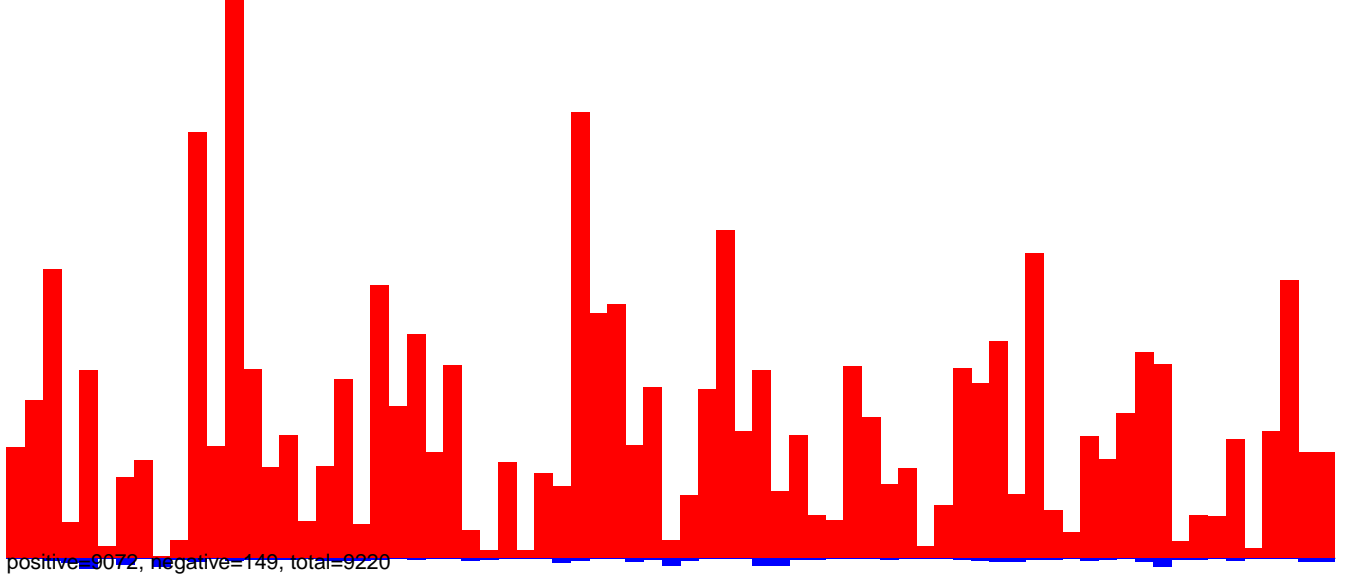
AeAeg_Testes.24_35.rep



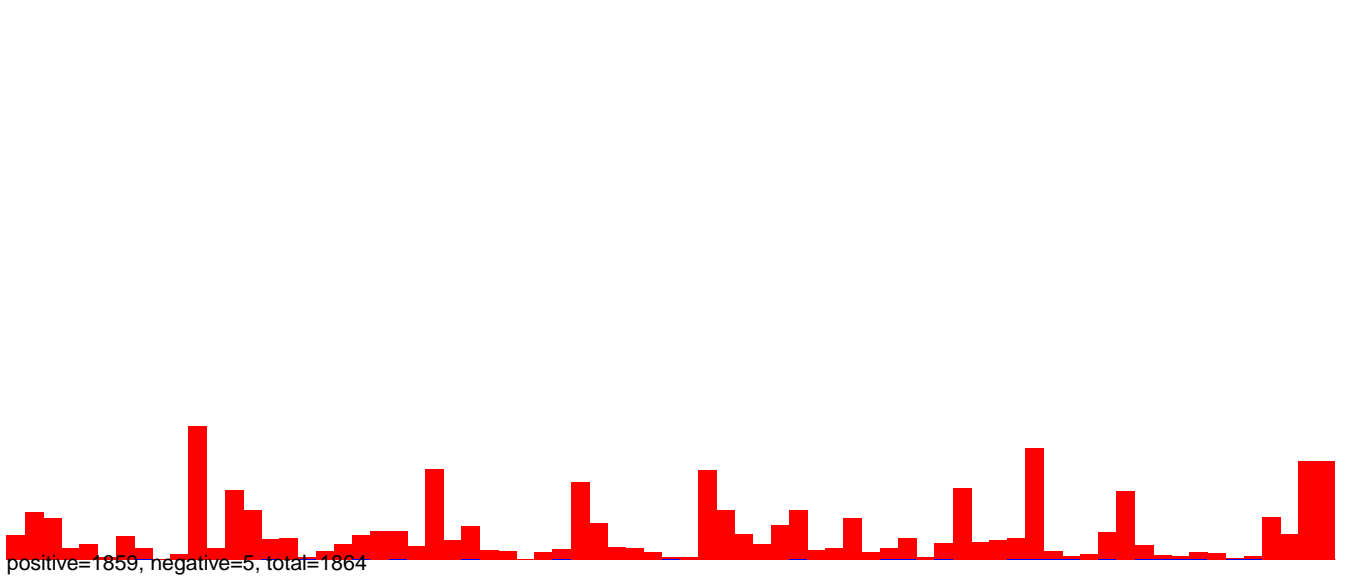
AeAeg_Testes.rep



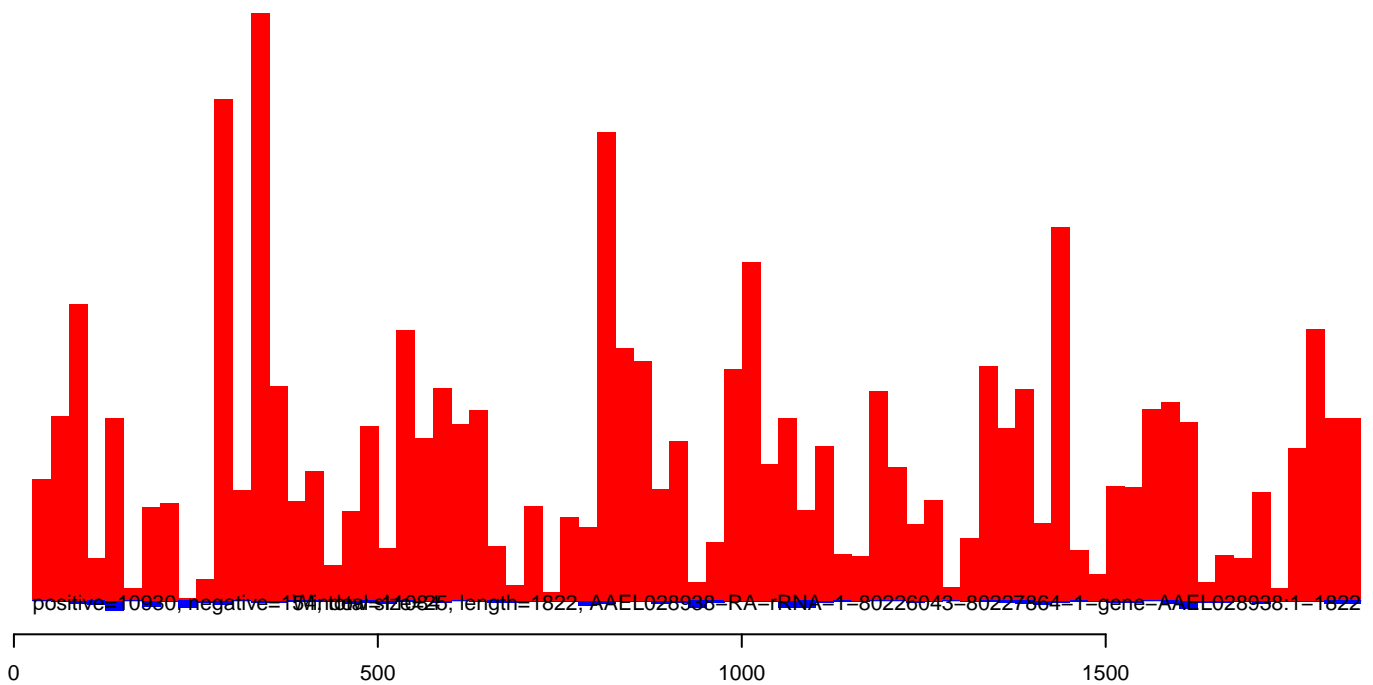
AeAeg_Testes.18_23.rep



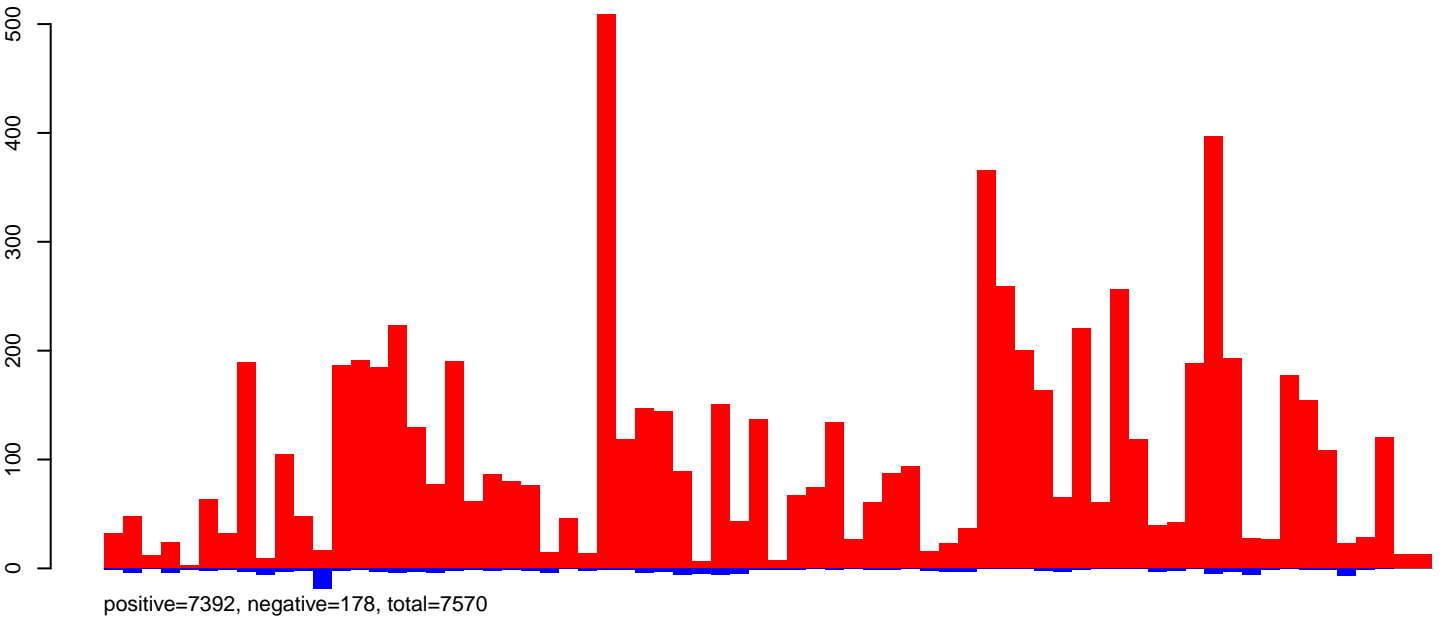
AeAeg_Testes.24_35.rep



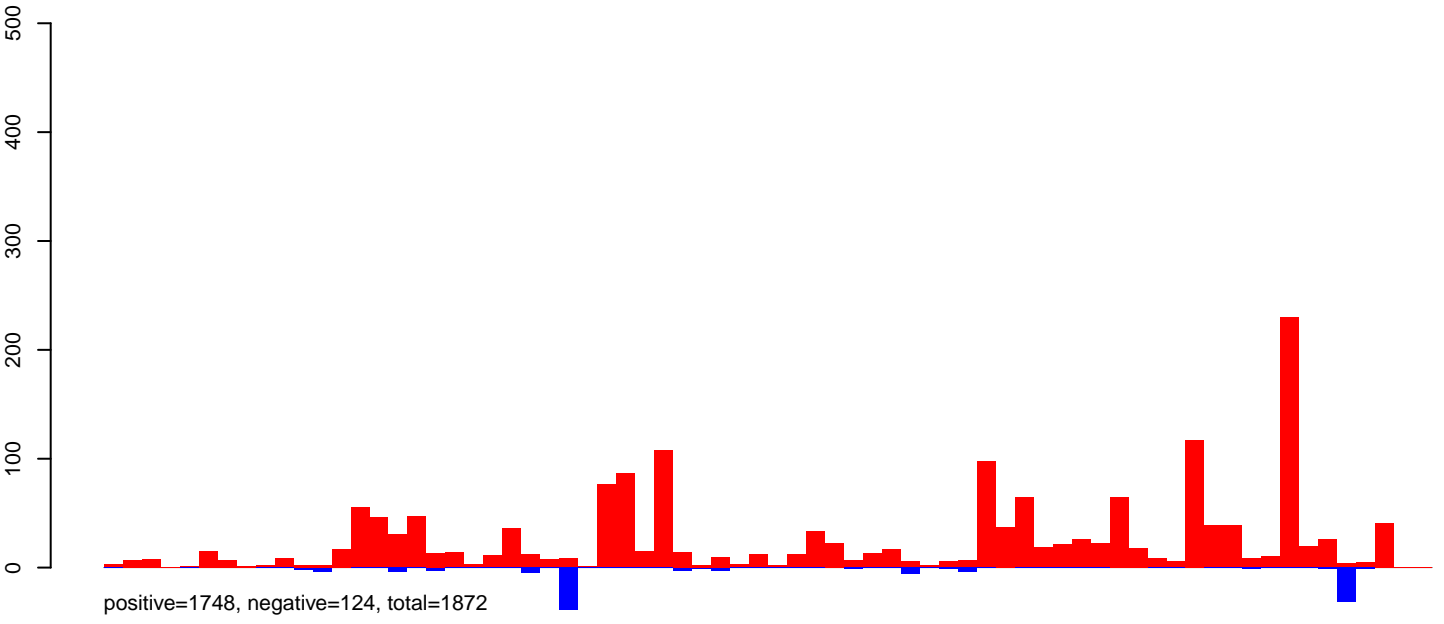
AeAeg_Testes.rep



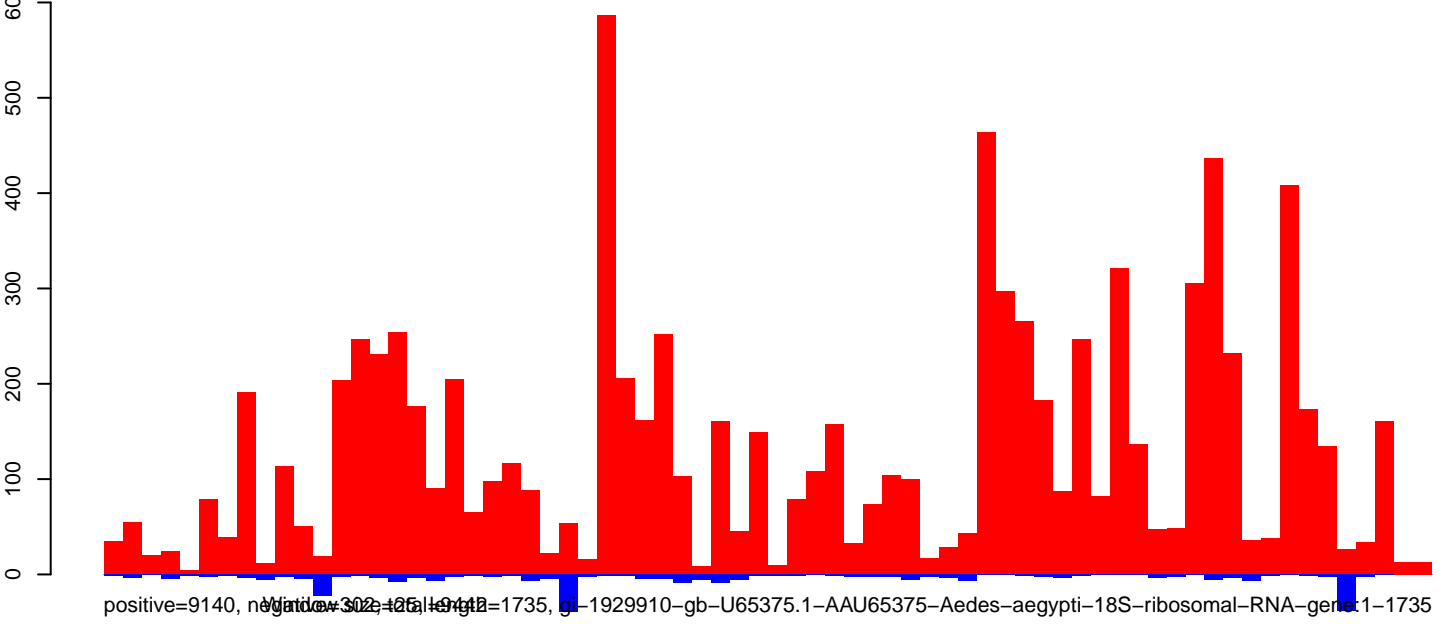
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



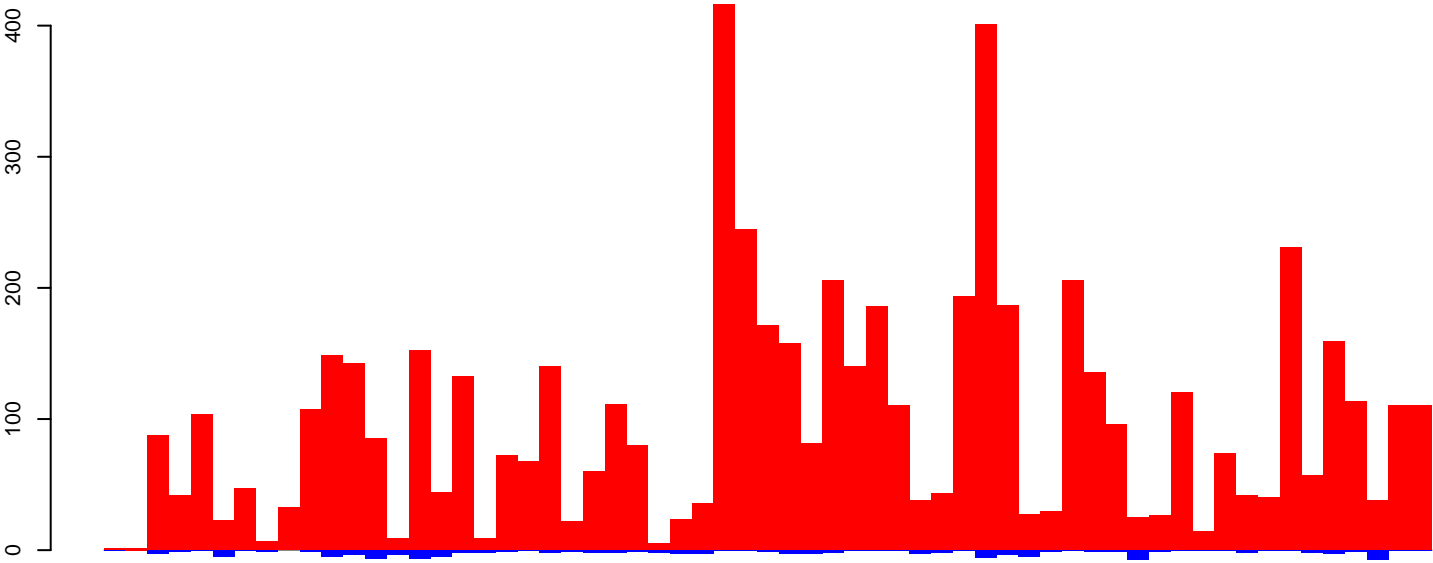
AeAeg_Testes.rep



gi|1929910-gb-U65375.1-AAU65375-Aedes-aegypti-18S-ribosomal-RNA-gene:1-1735

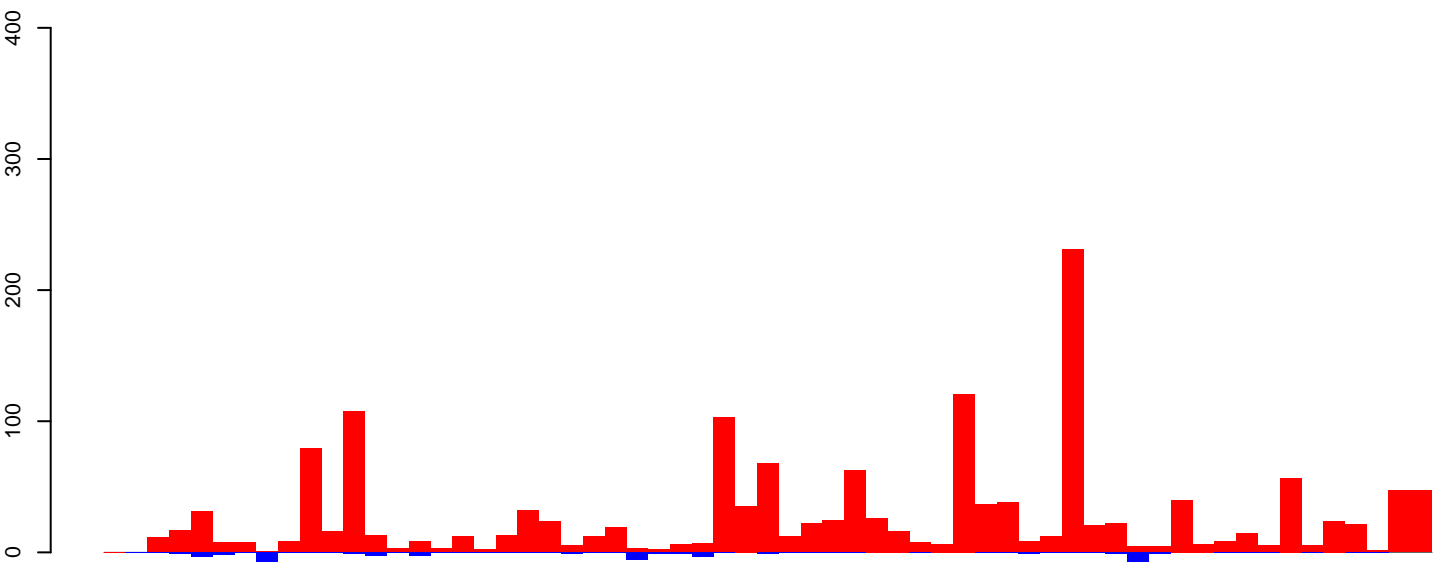
0 500 1000 1500

AeAeg_Testes.18_23.rep



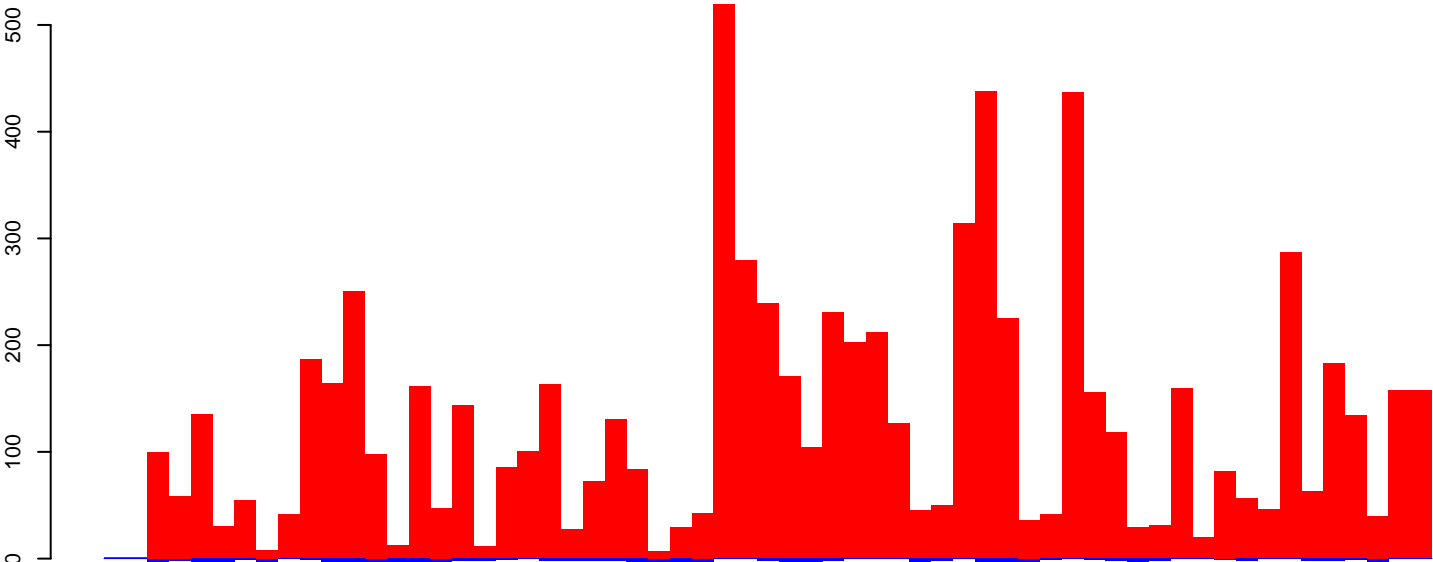
positive=6026, negative=121, total=6147

AeAeg_Testes.24_35.rep



positive=1625, negative=111, total=1735

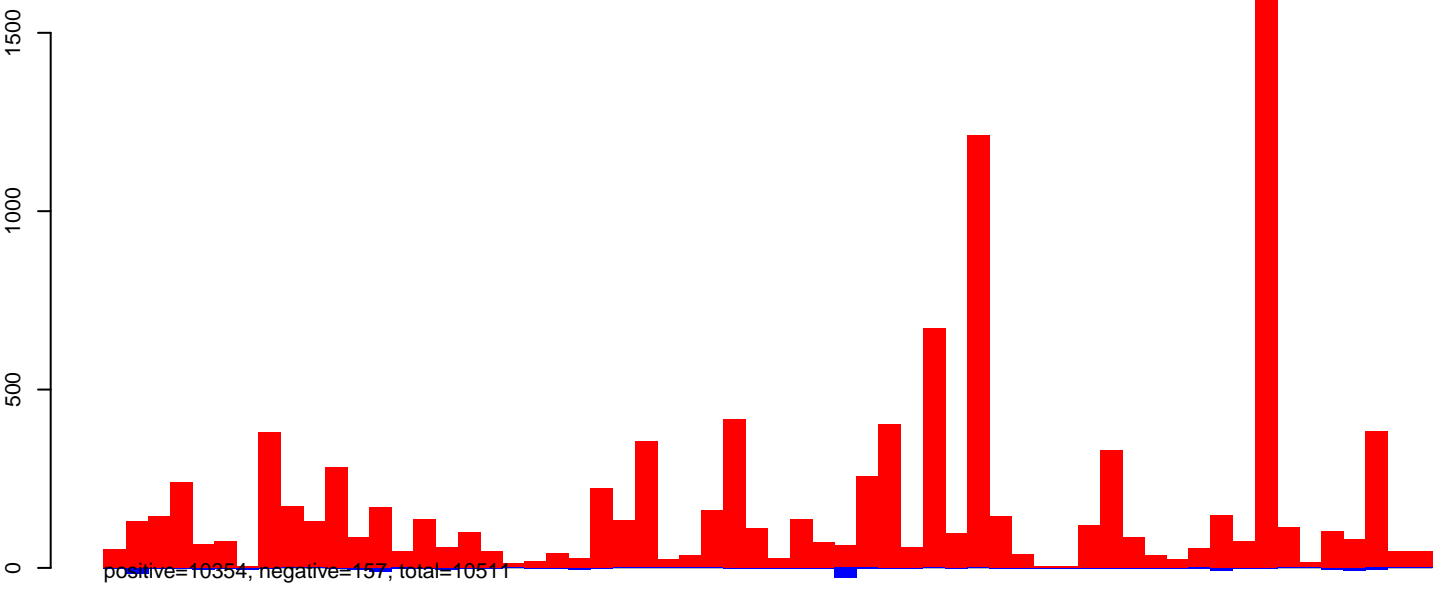
AeAeg_Testes.rep



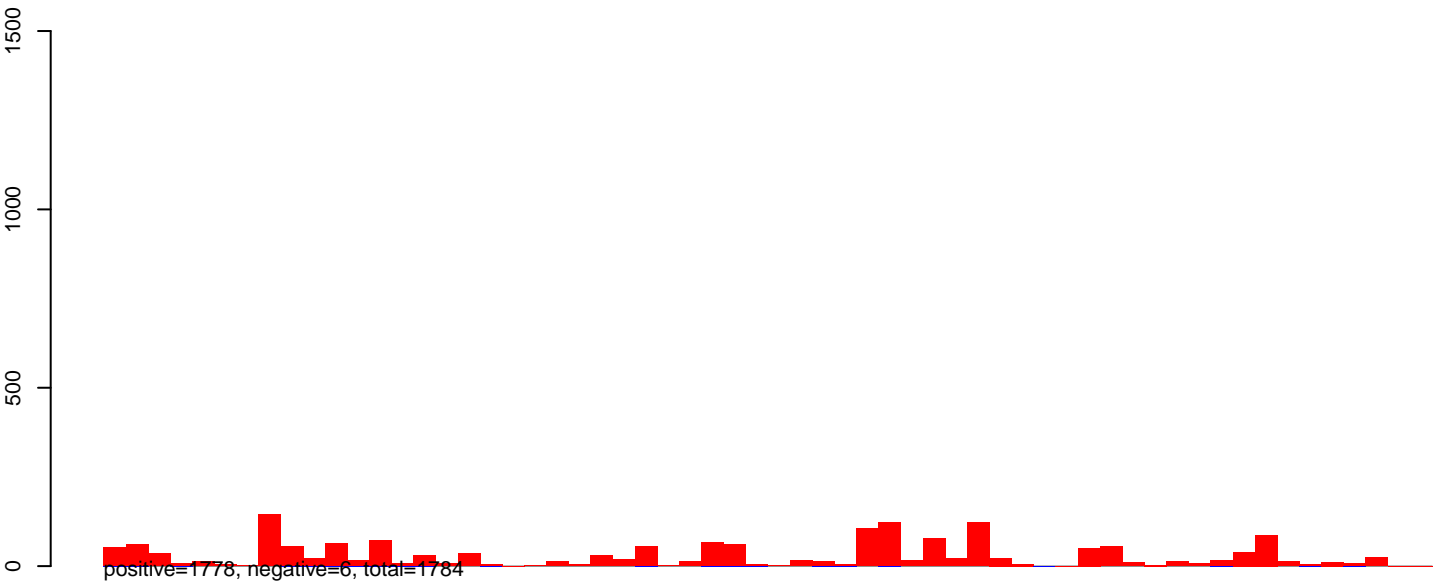
positive=7651, negative=201, total=7852, length=1506, AAEL028907-RA-rRNA-1-205653557-205655062-1-gene-AAEL028907:1-1506

0 500 1000 1500

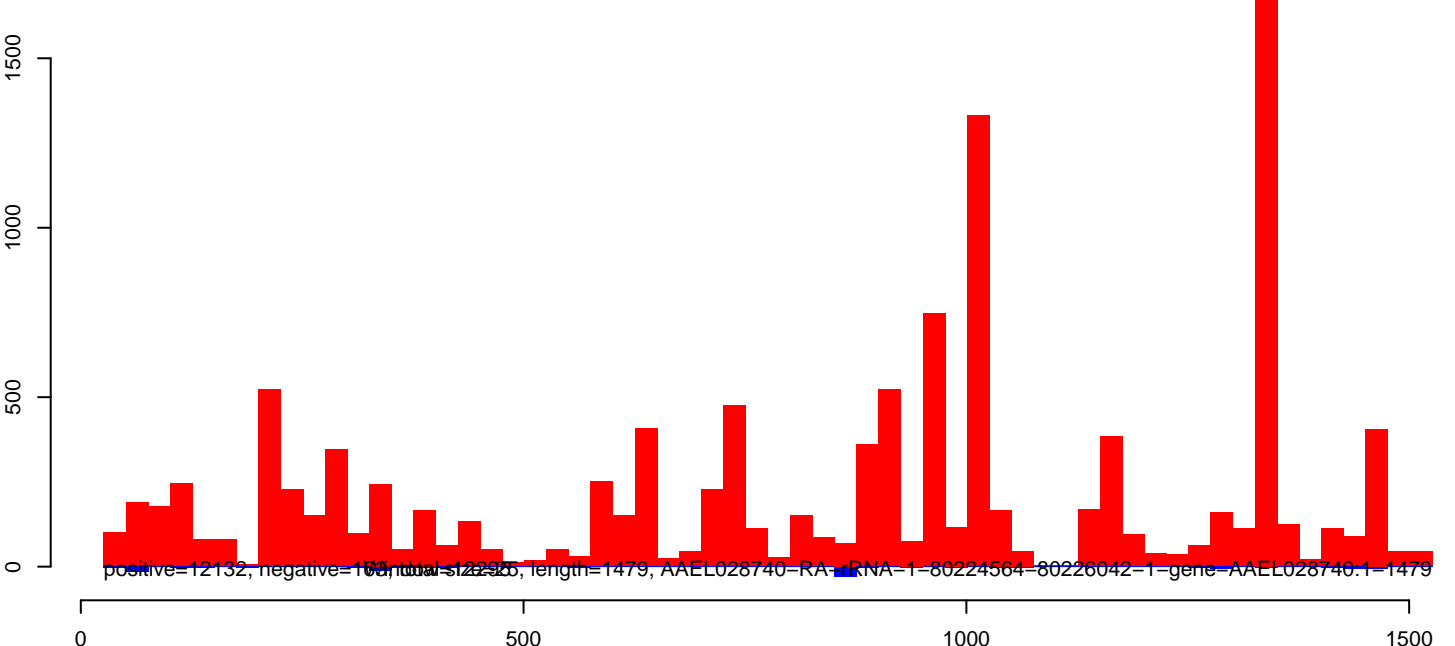
AeAeg_Testes.18_23.rep



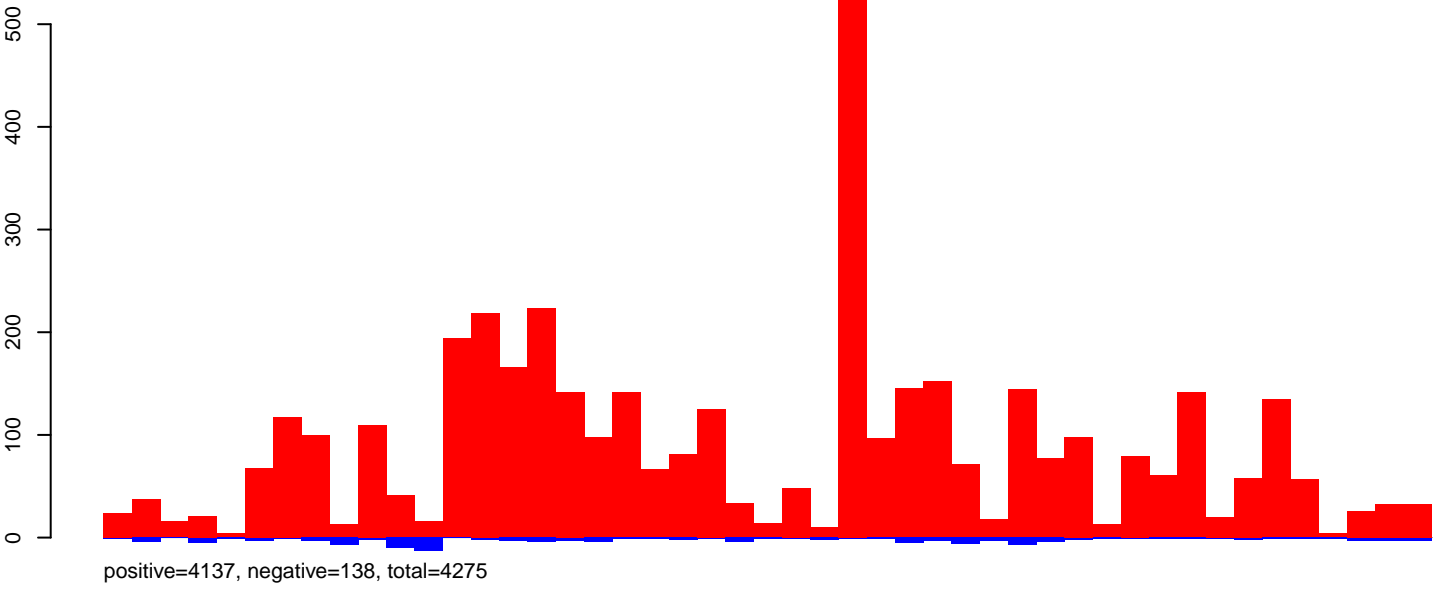
AeAeg_Testes.24_35.rep



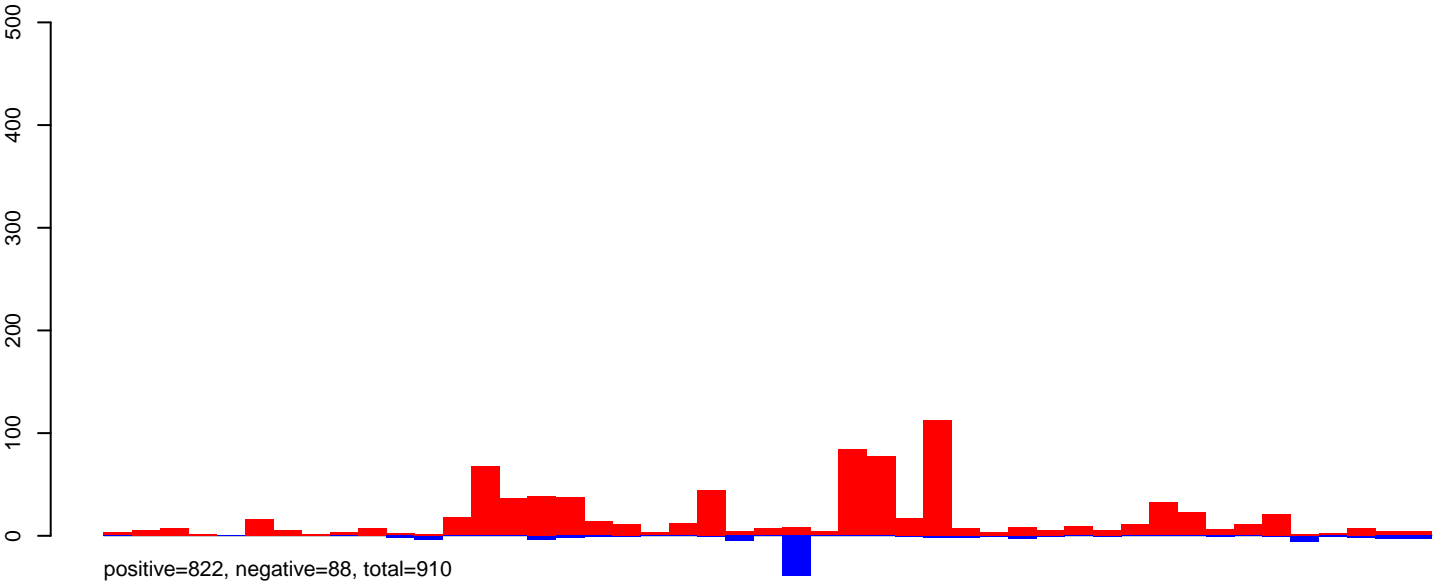
AeAeg_Testes.rep



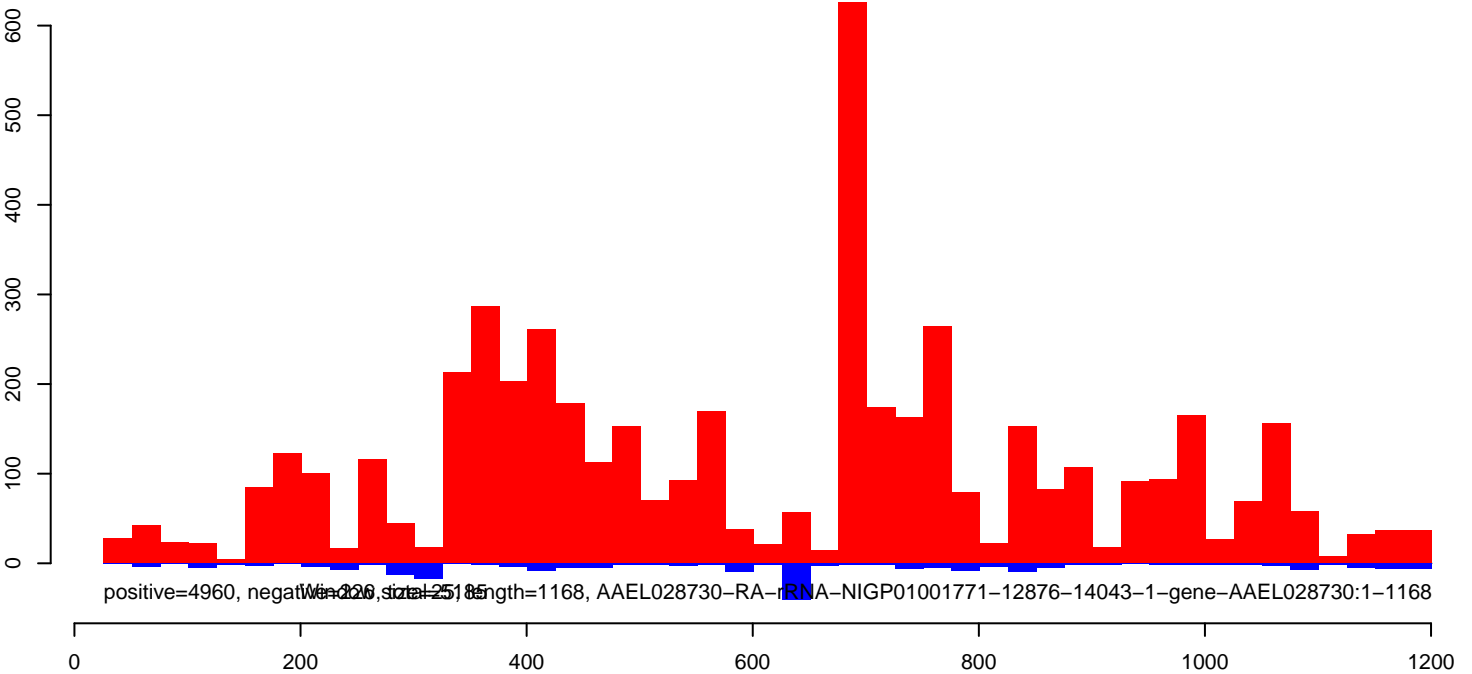
AeAeg_Testes.18_23.rep



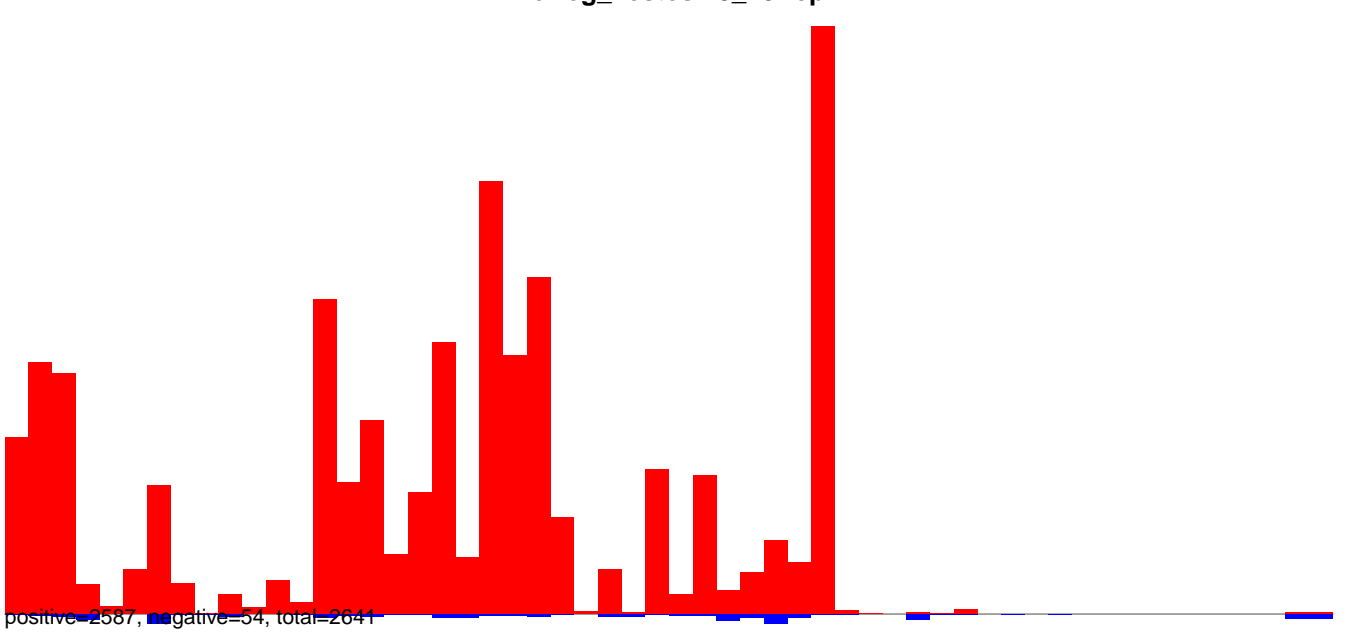
AeAeg_Testes.24_35.rep



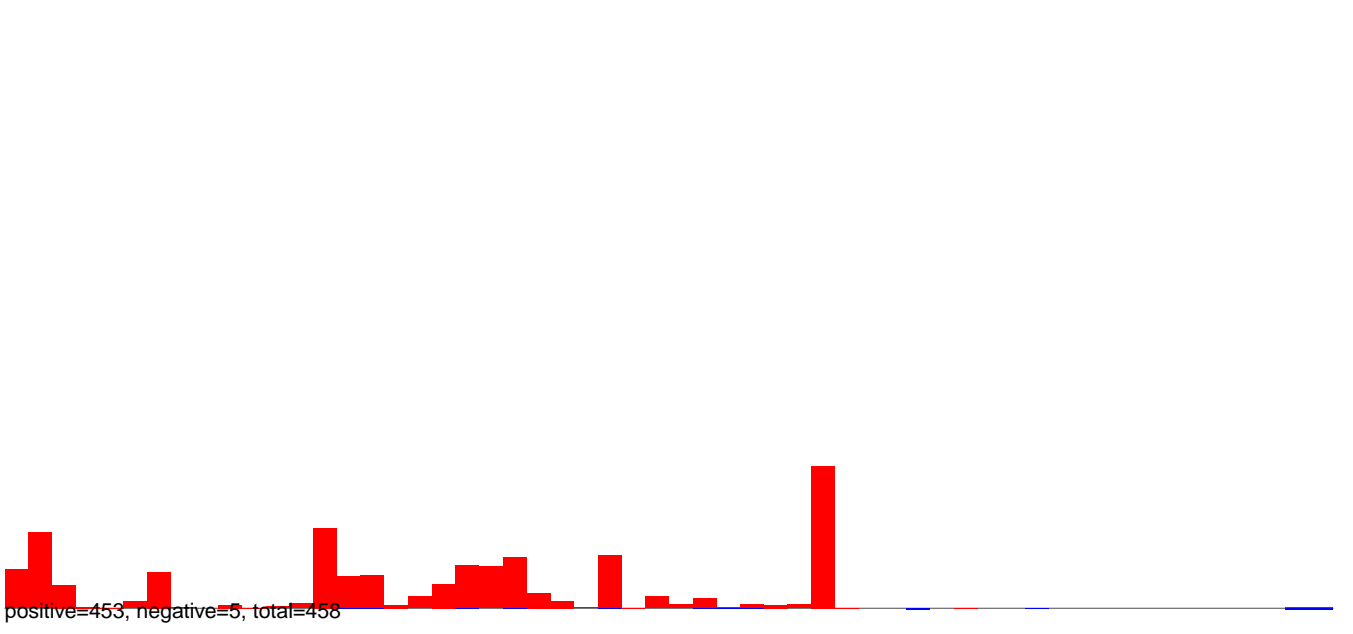
AeAeg_Testes.rep



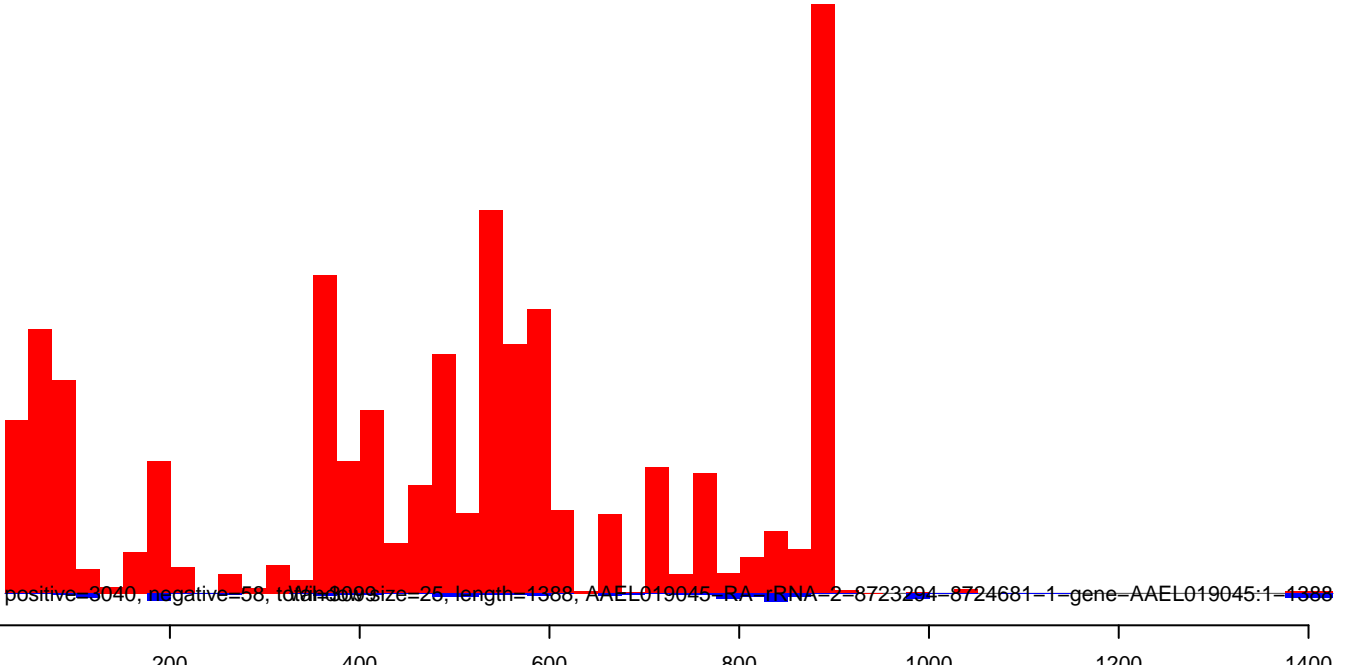
AeAeg_Testes.18_23.rep



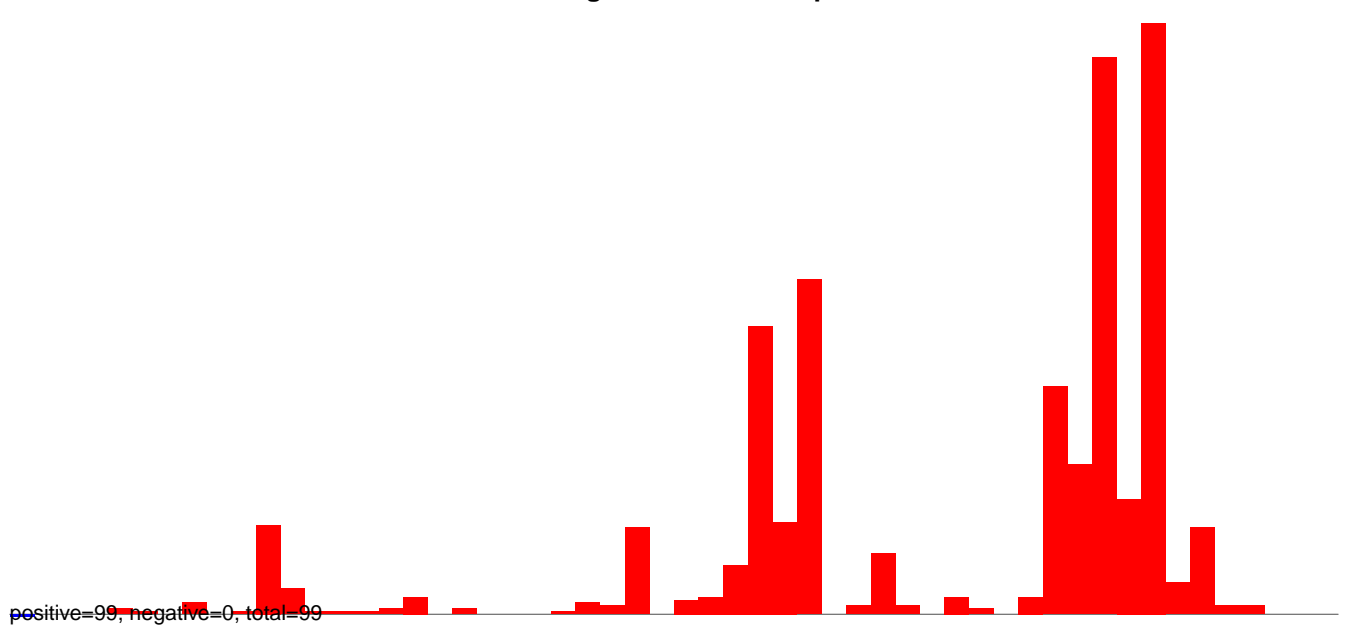
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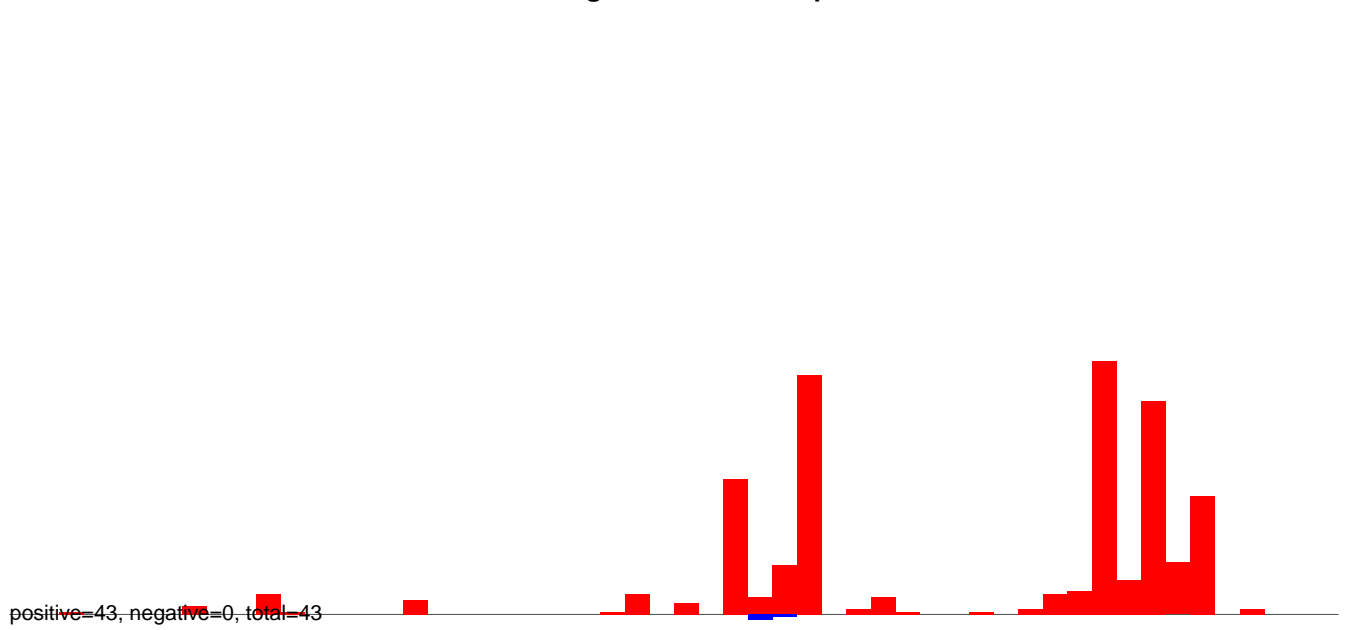
AeAeg_Testes.rep



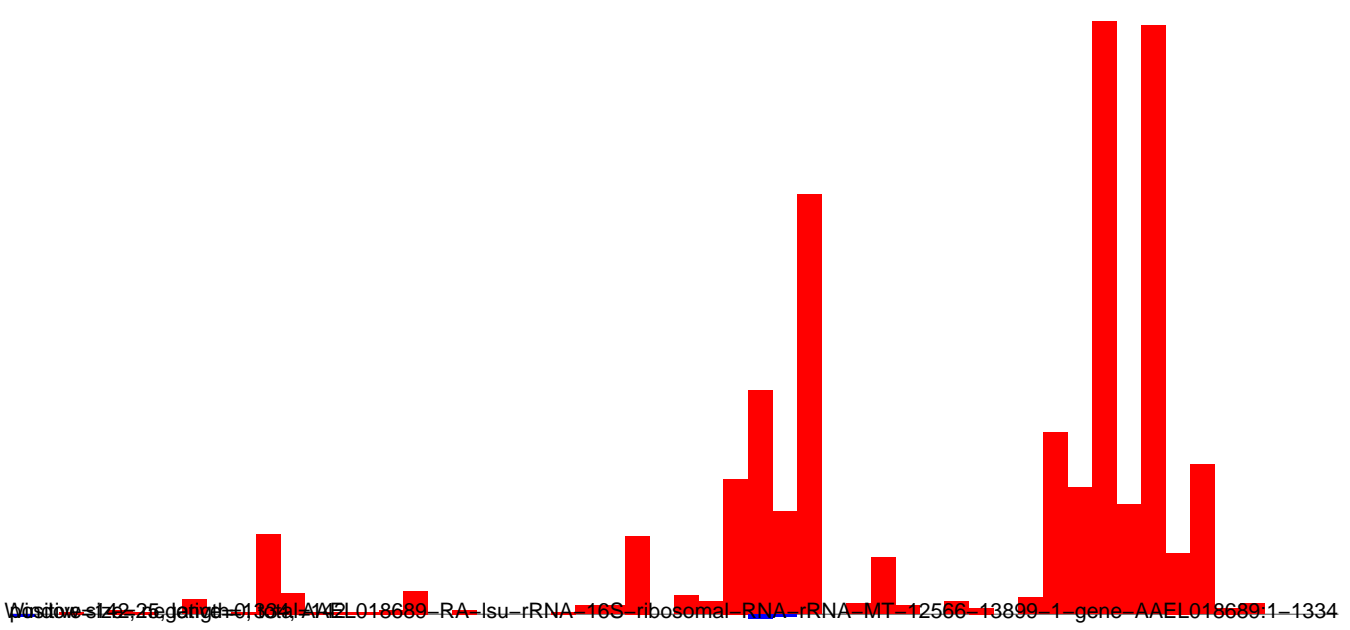
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep

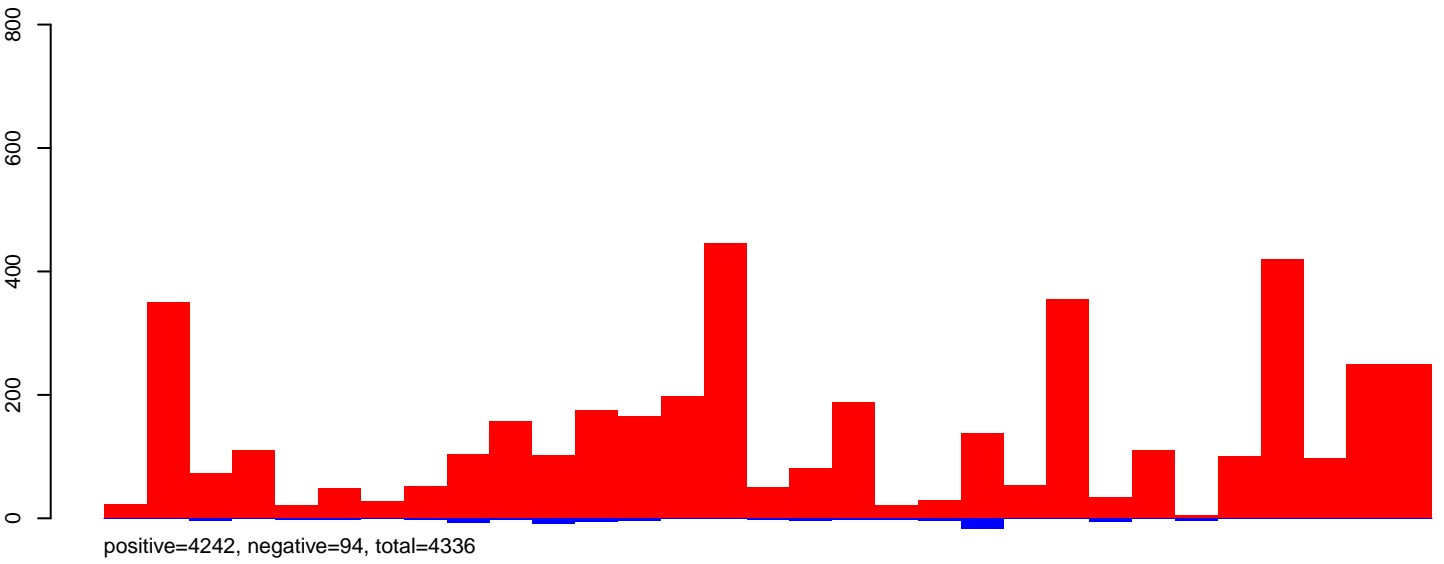


AeAeg_Testes.rep

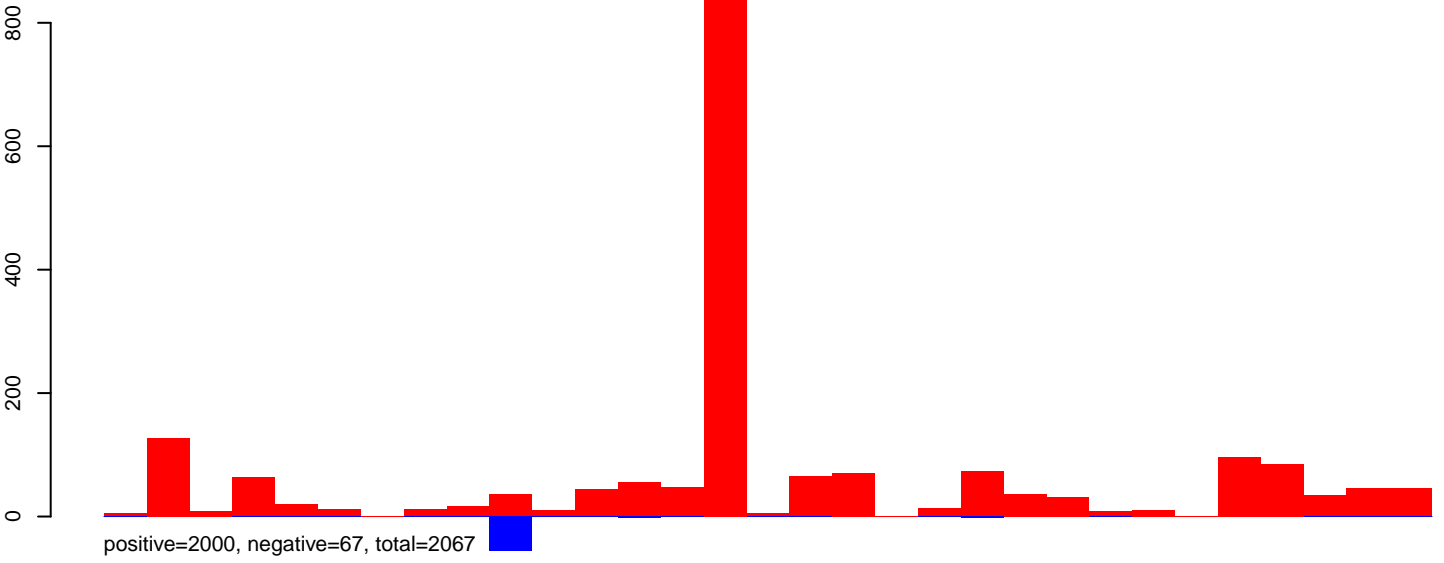


Window size: 25, gap: 0, total=142, AAEL018689-RA-Isu-rRNA-16S-ribosomal-RNA-rRNA-MT-12566-13899-1-gene-AAEL018689.1-1334

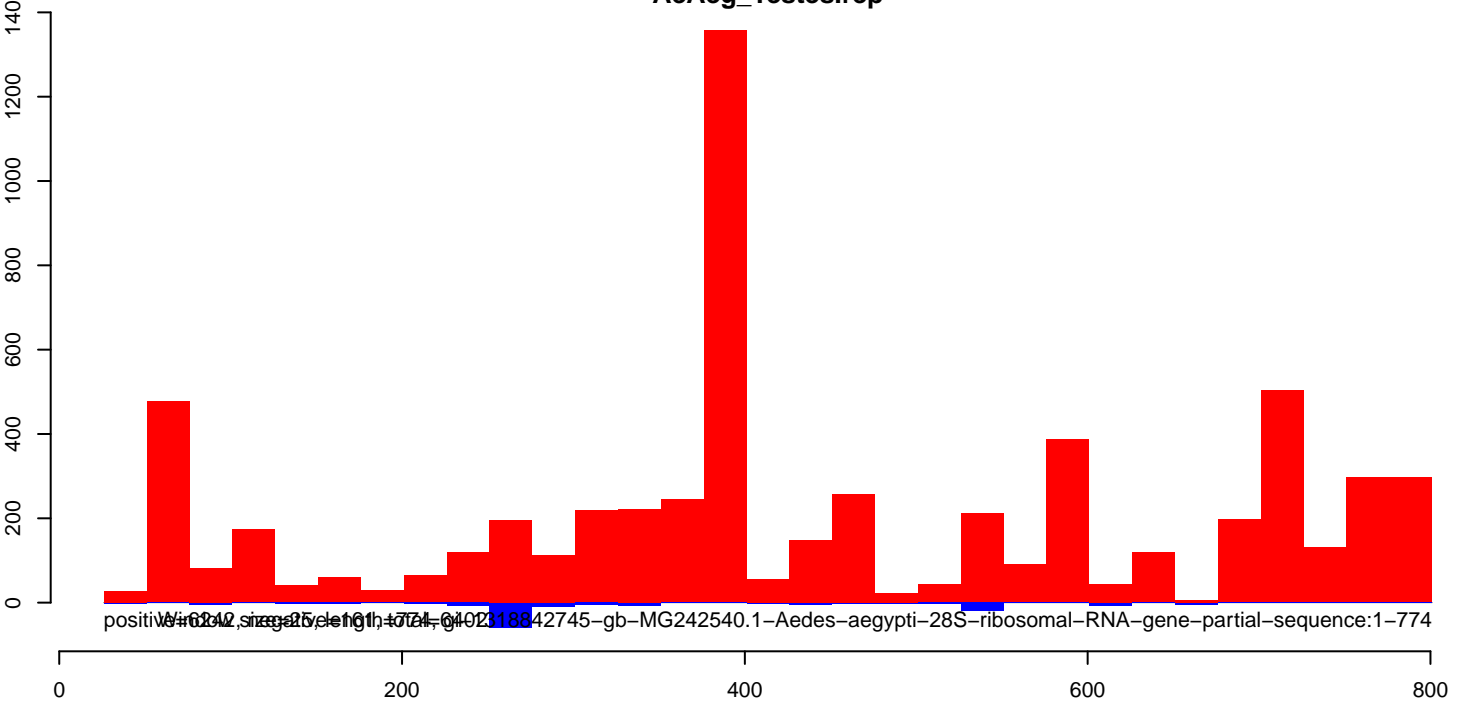
AeAeg_Testes.18_23.rep



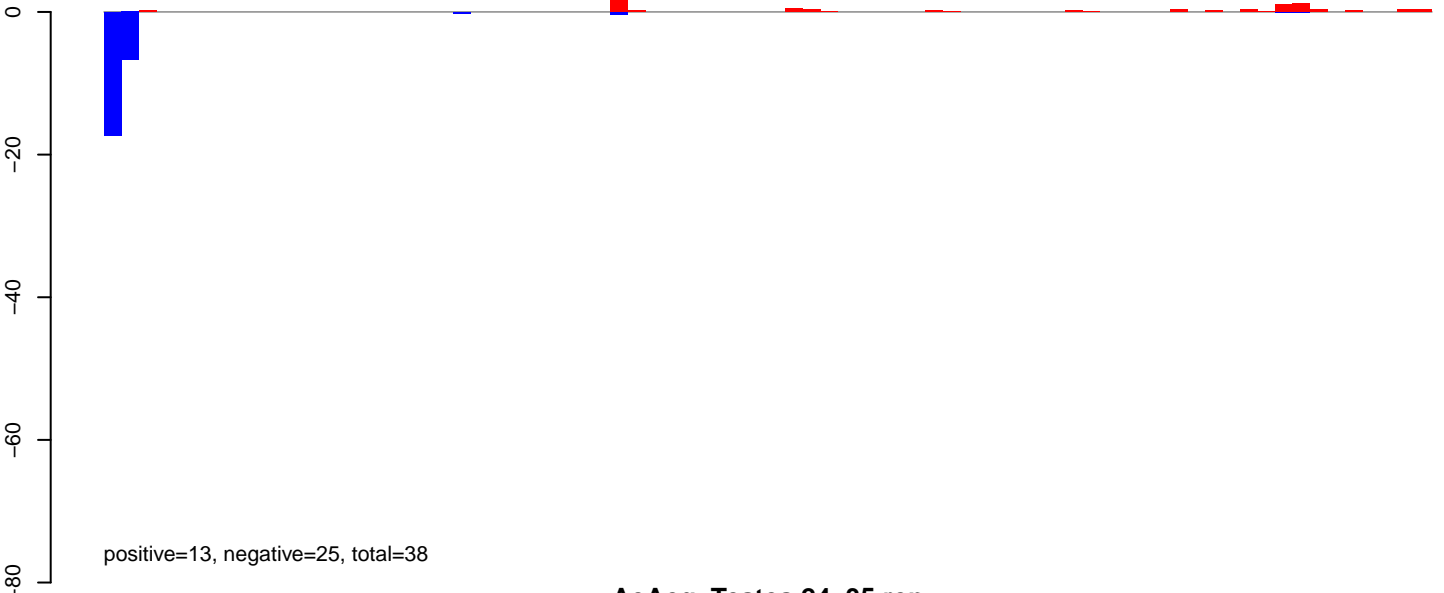
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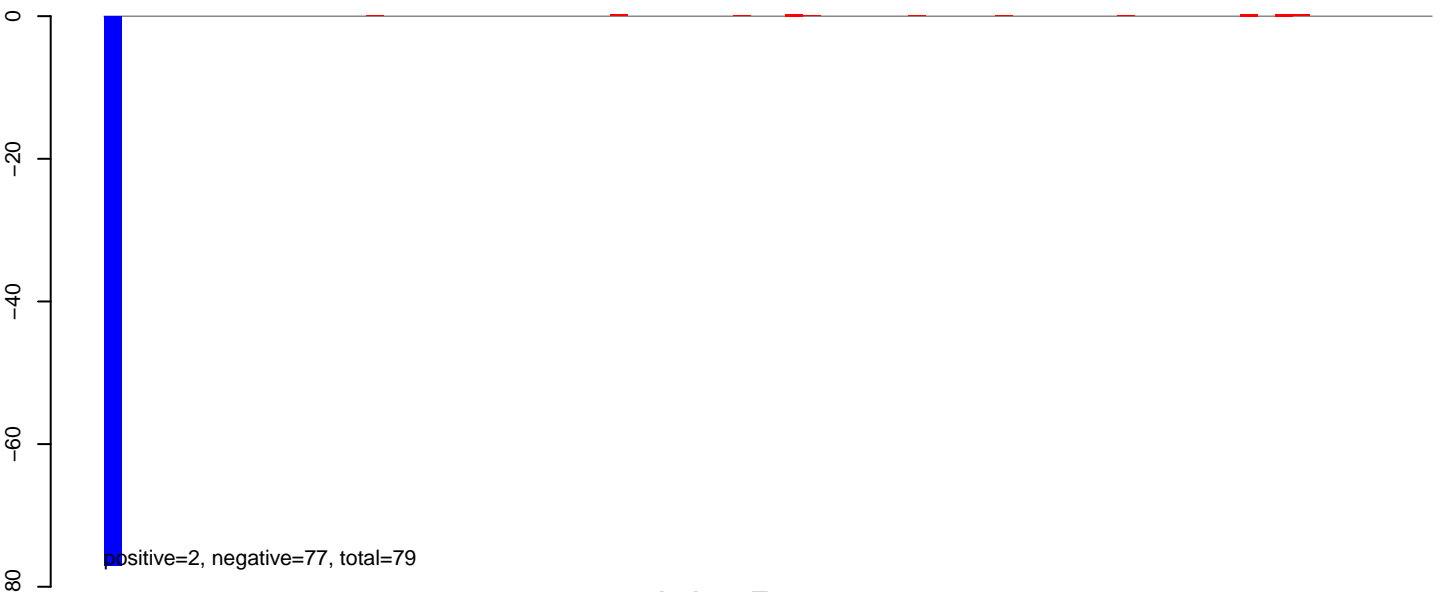
AeAeg_Testes.rep



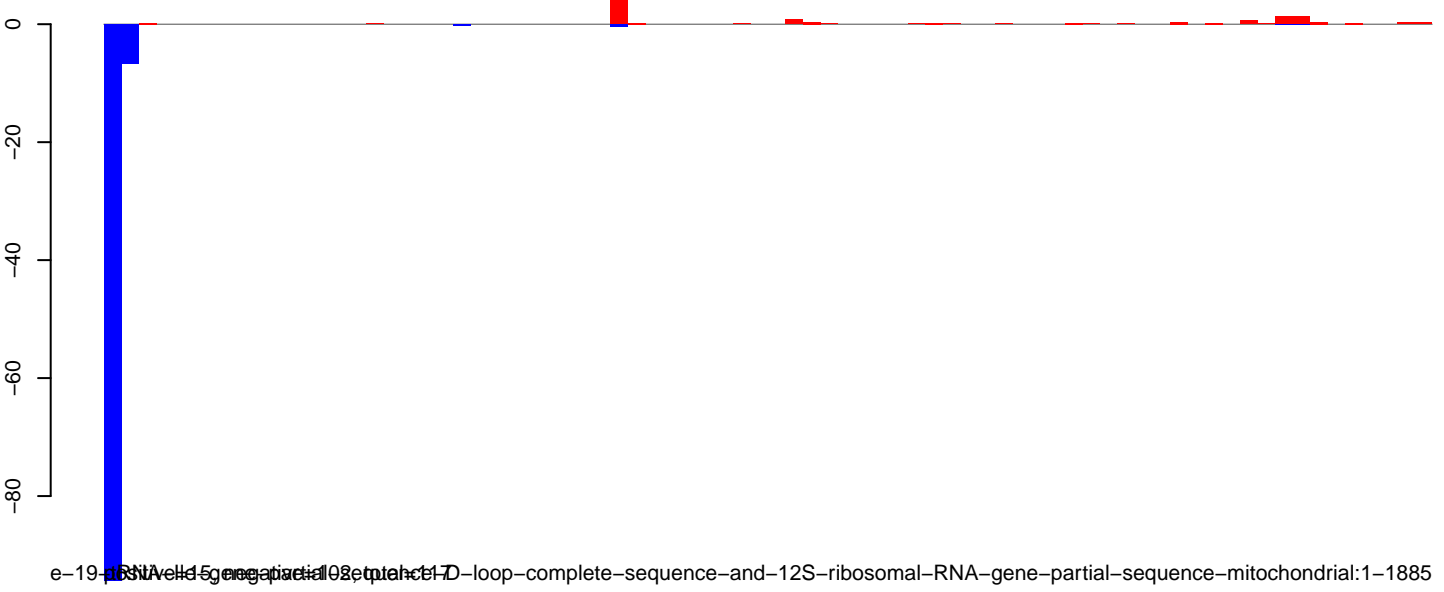
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AeAeg_Testes.24_35.rep

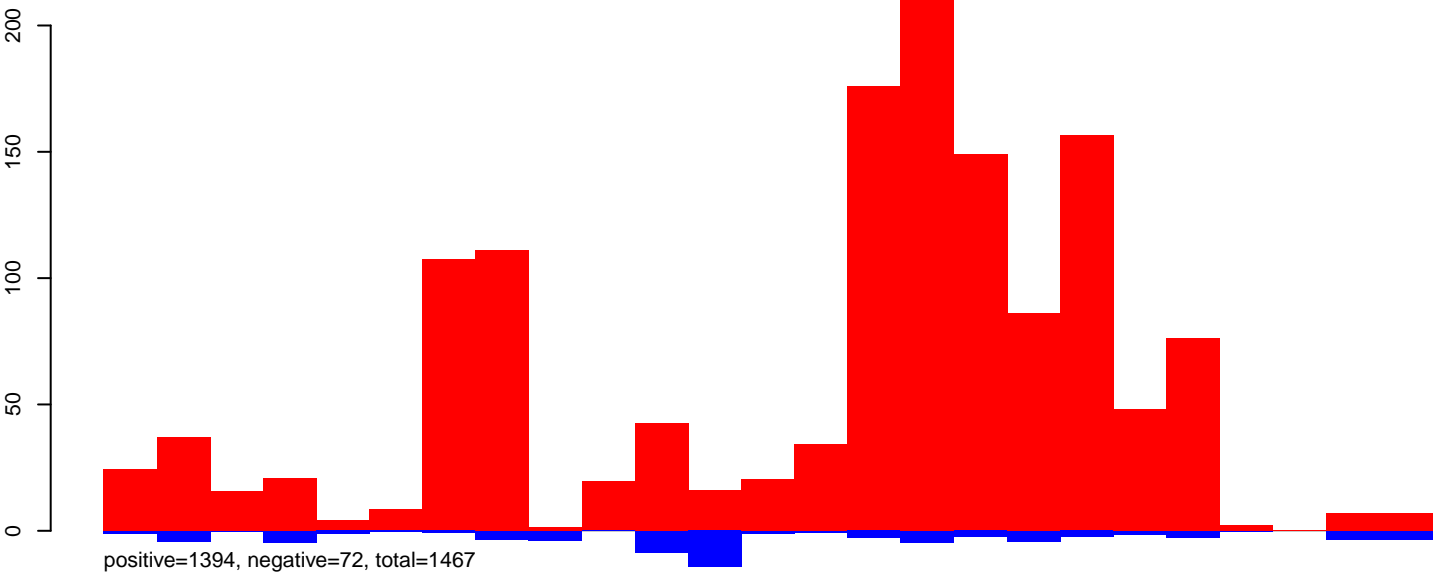


AeAeg_Testes.rep

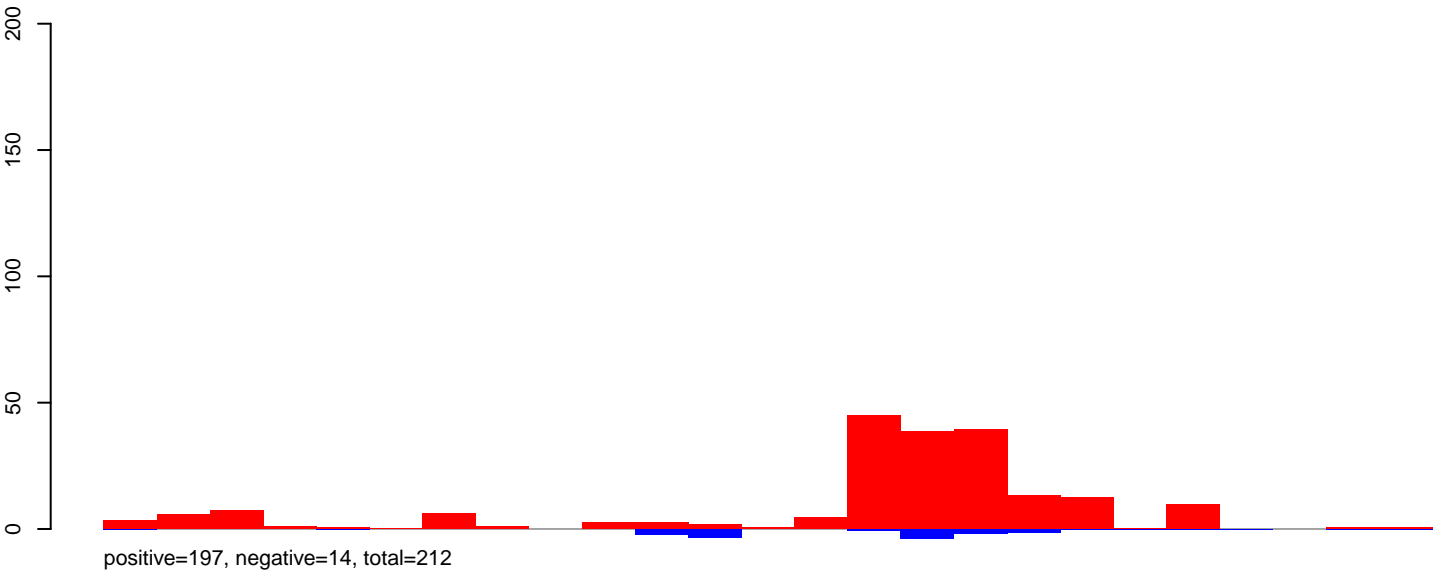


e-19-531745 gene partial 02 p1nc17 D-loop-complete-sequence-and-12S-ribosomal-RNA-gene-partial-sequence-mitochondrial:1-1885

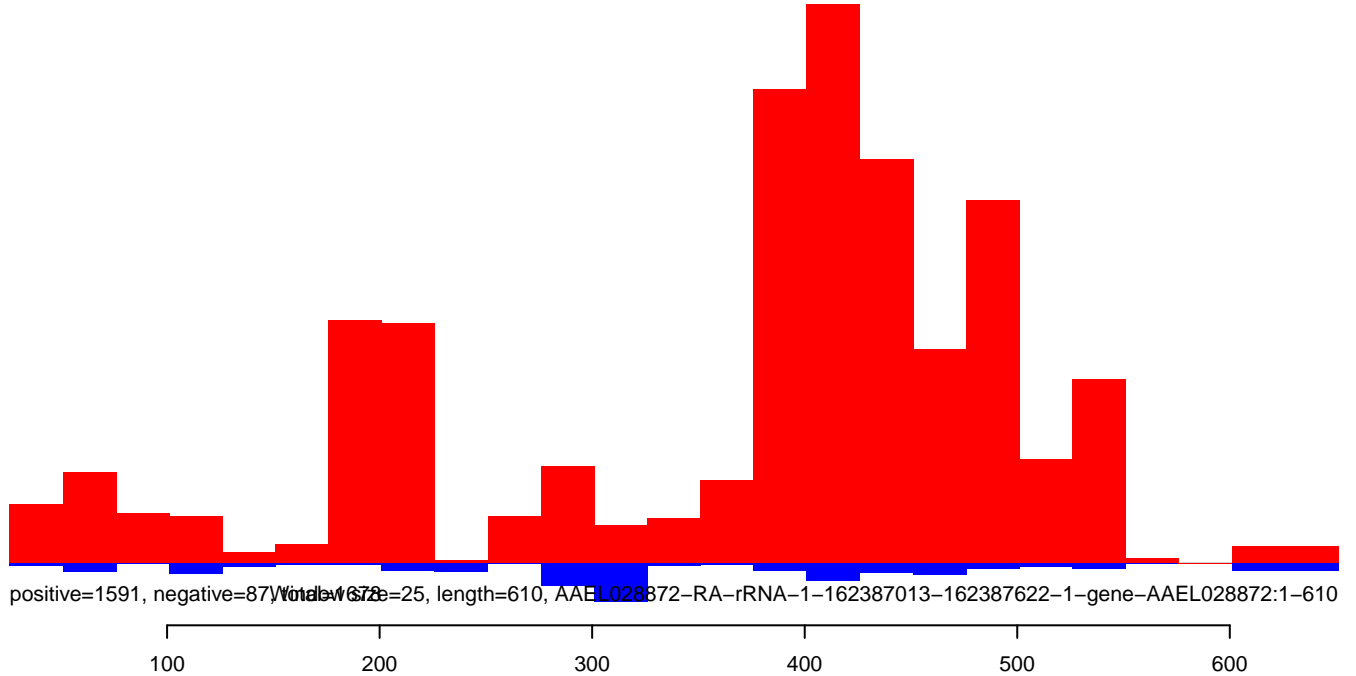
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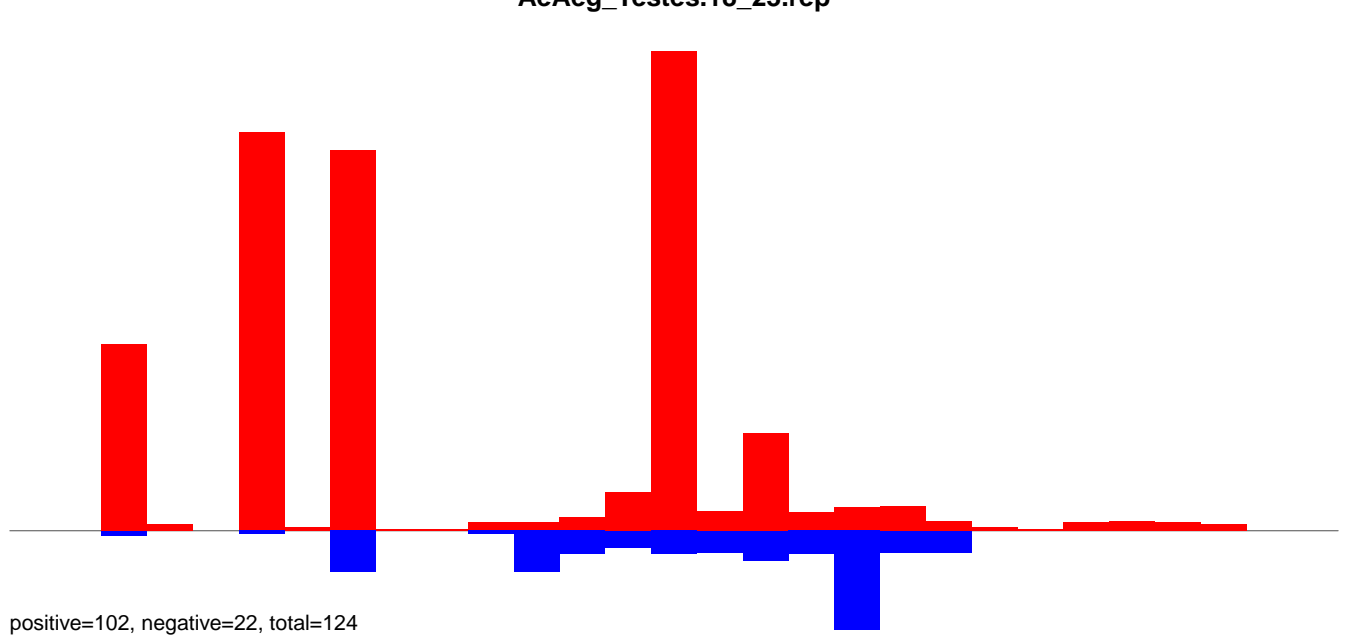
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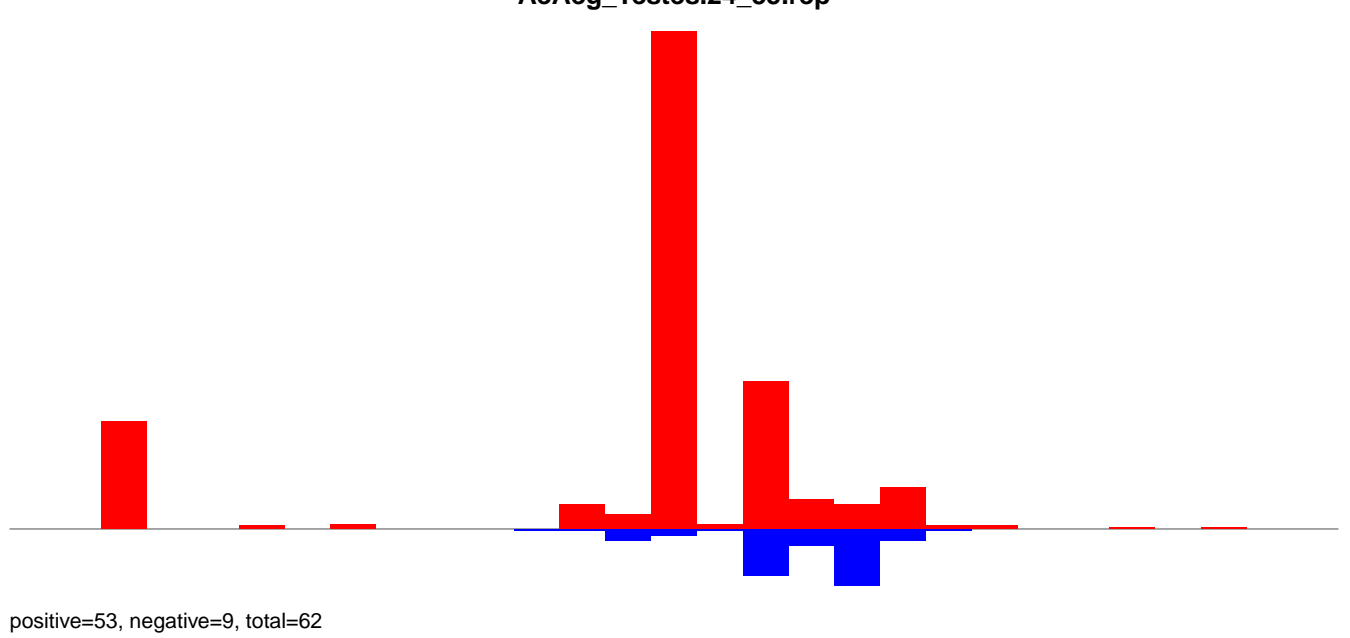
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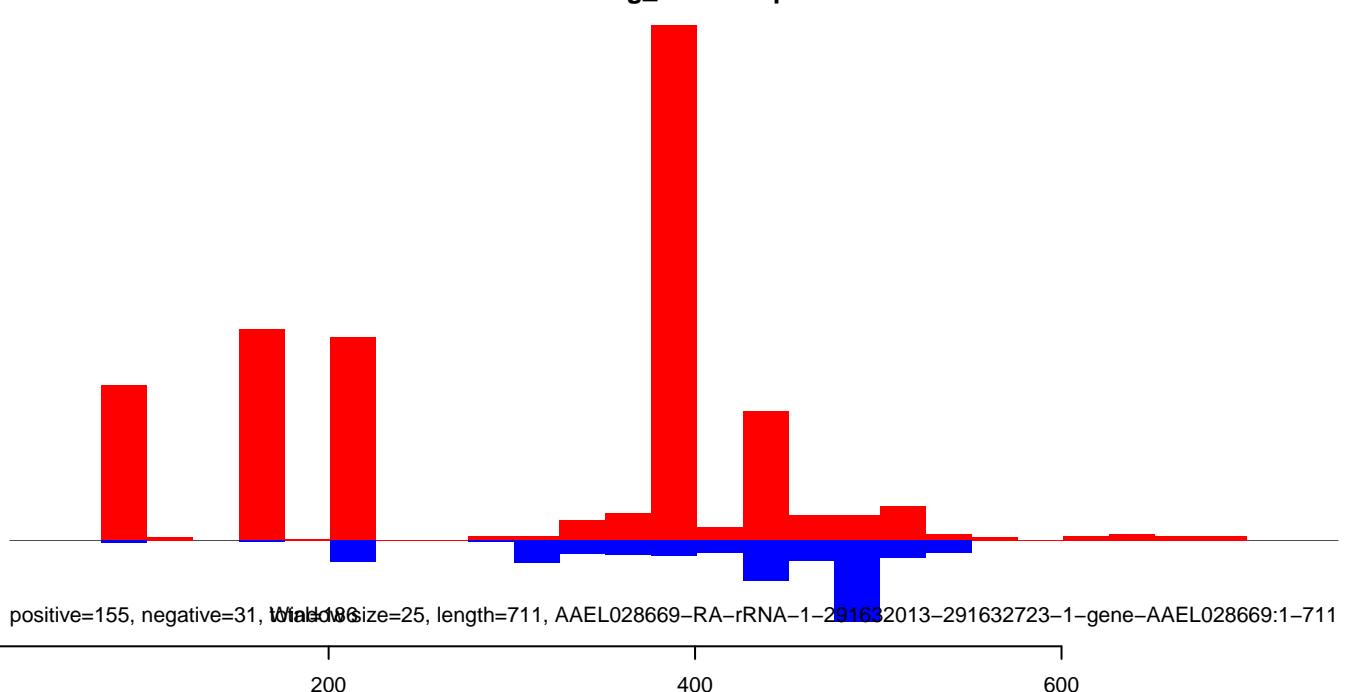
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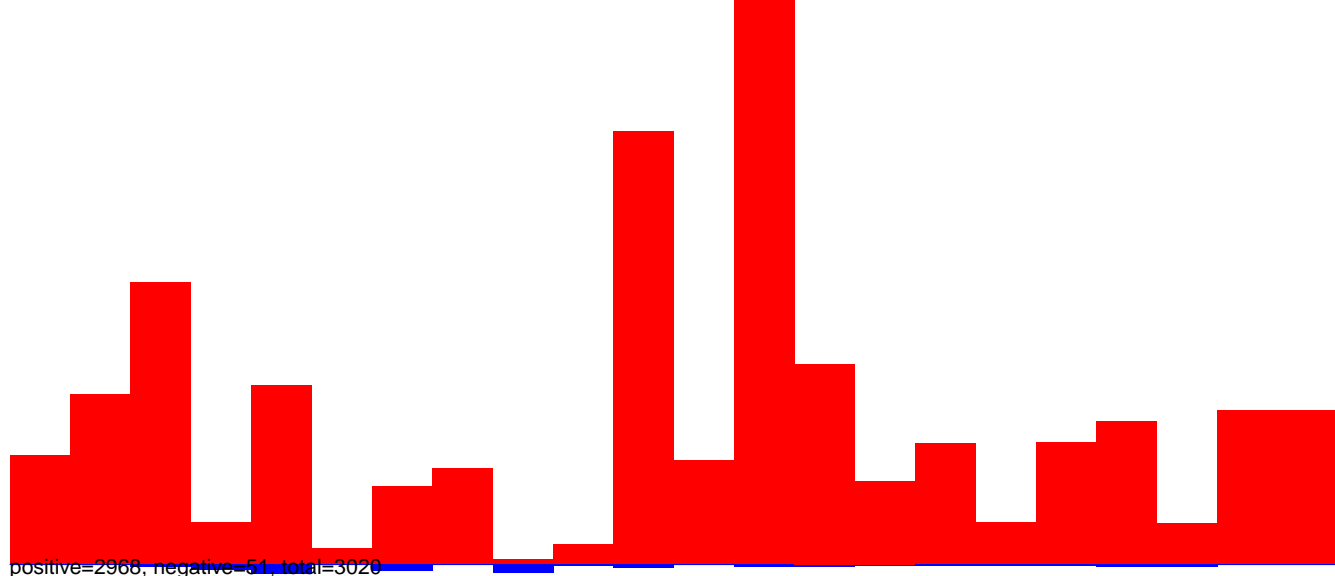
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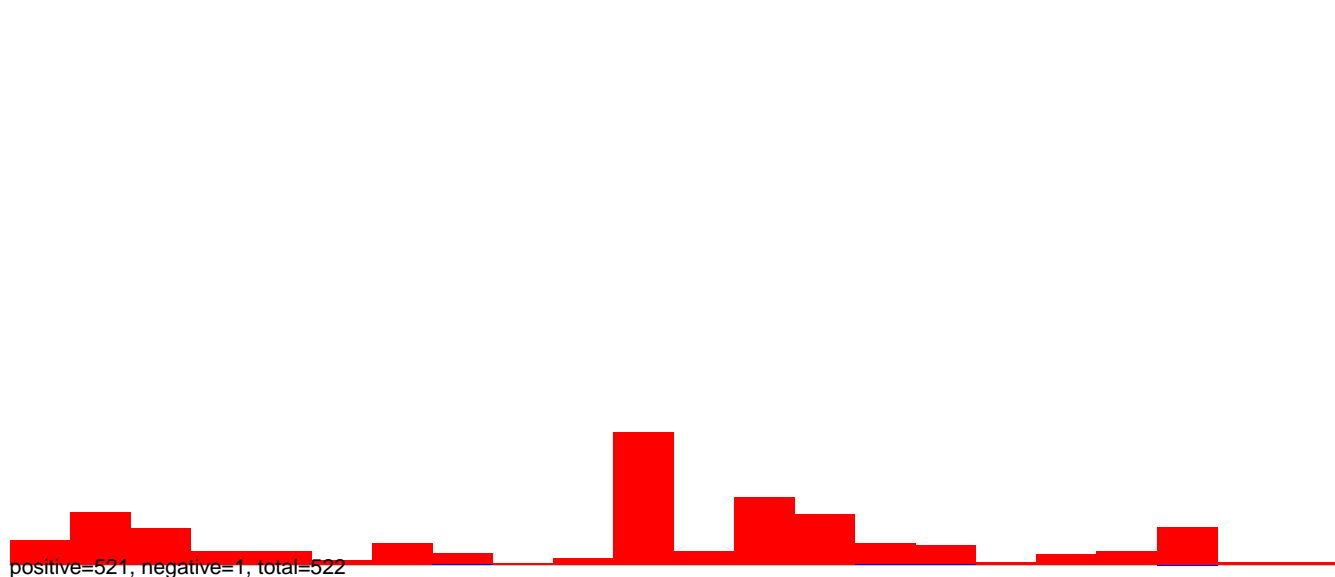
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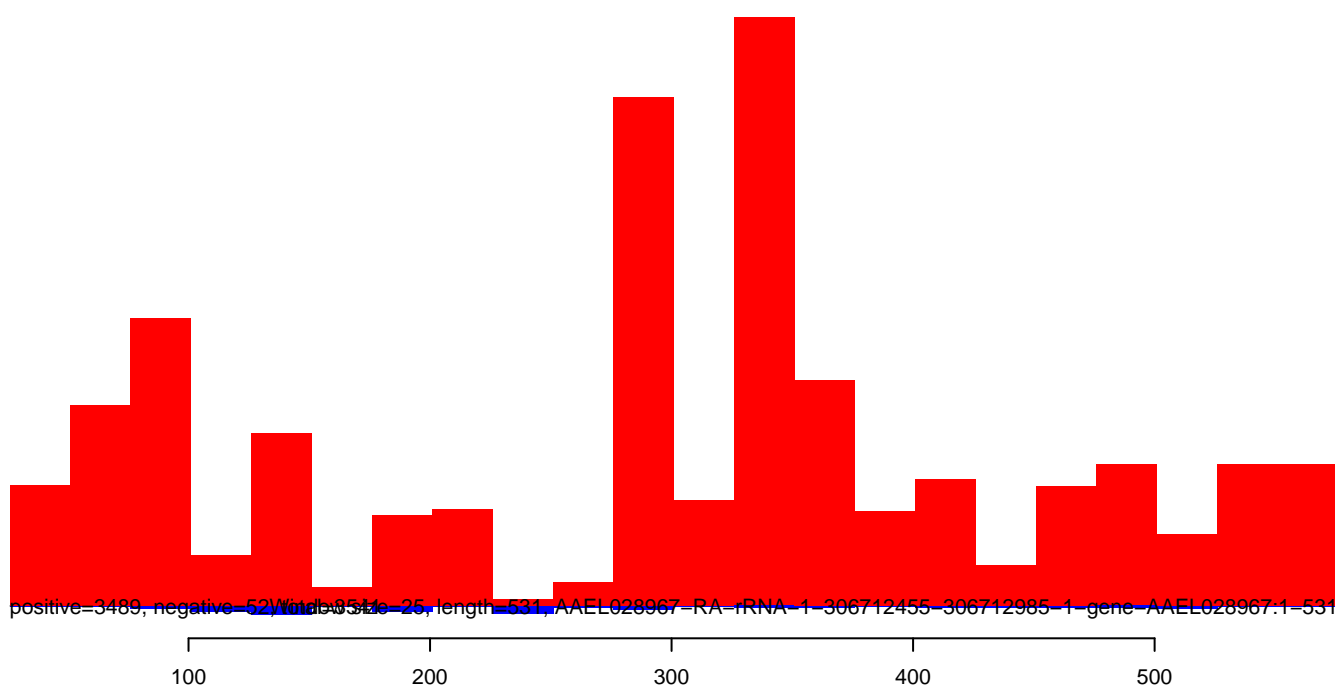
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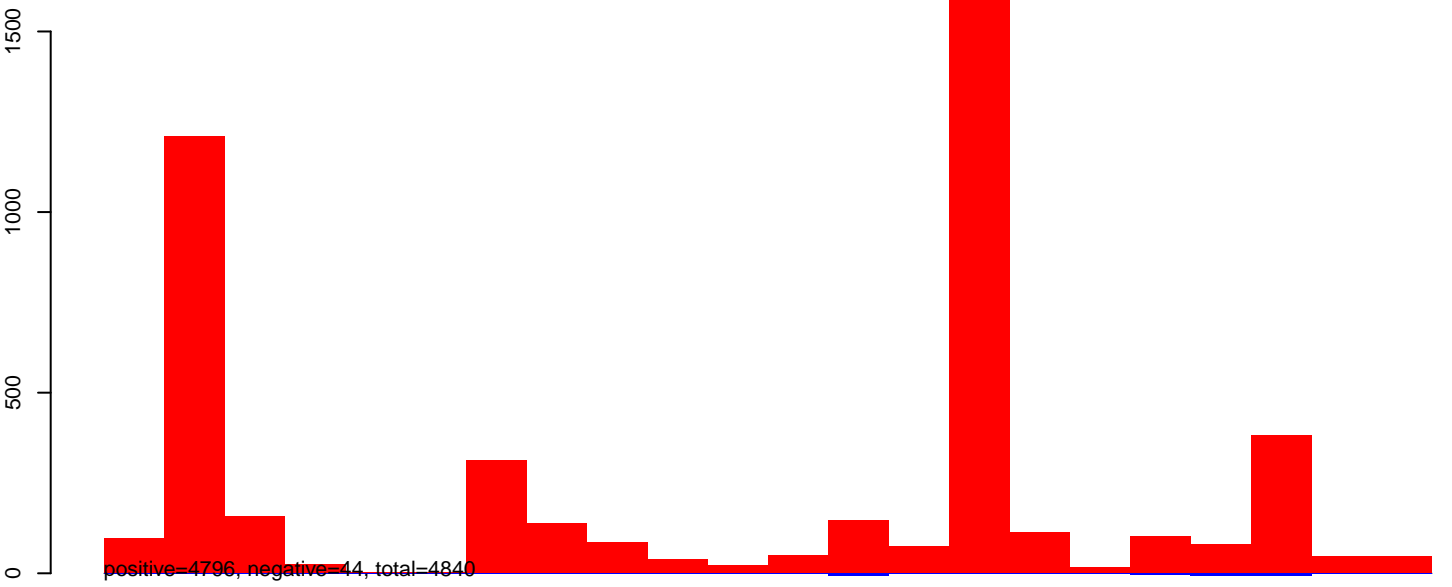
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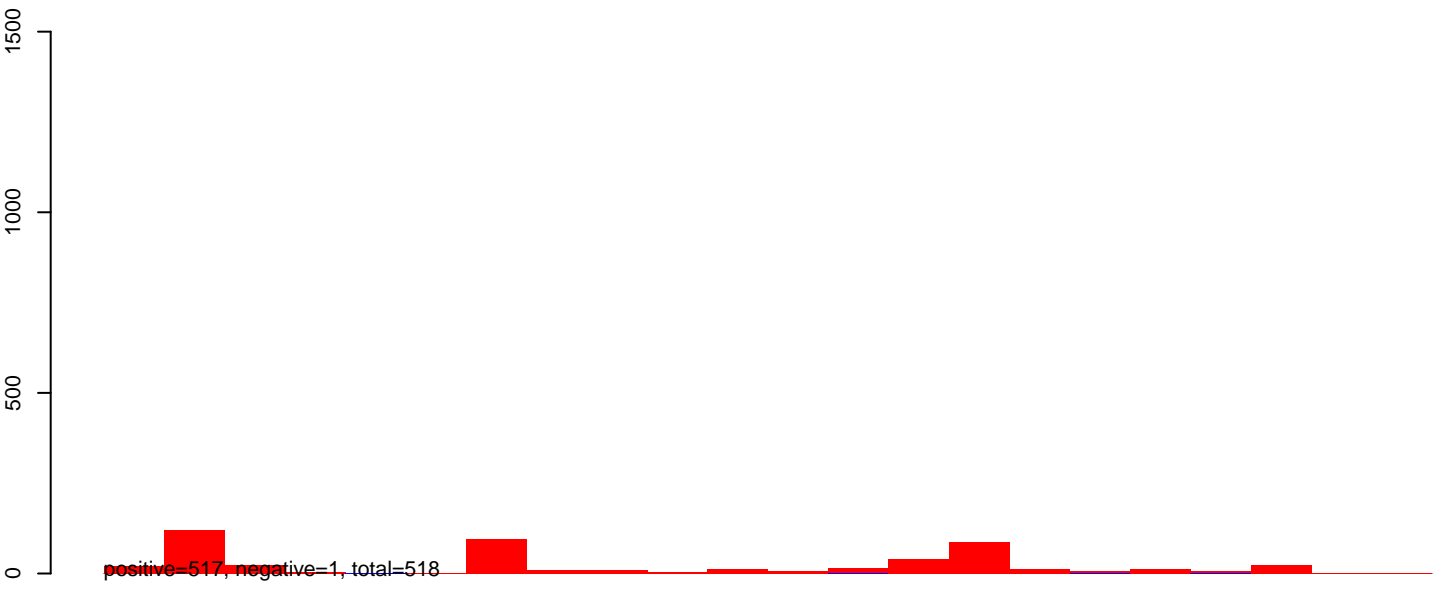
AeAeg_Testes.rep



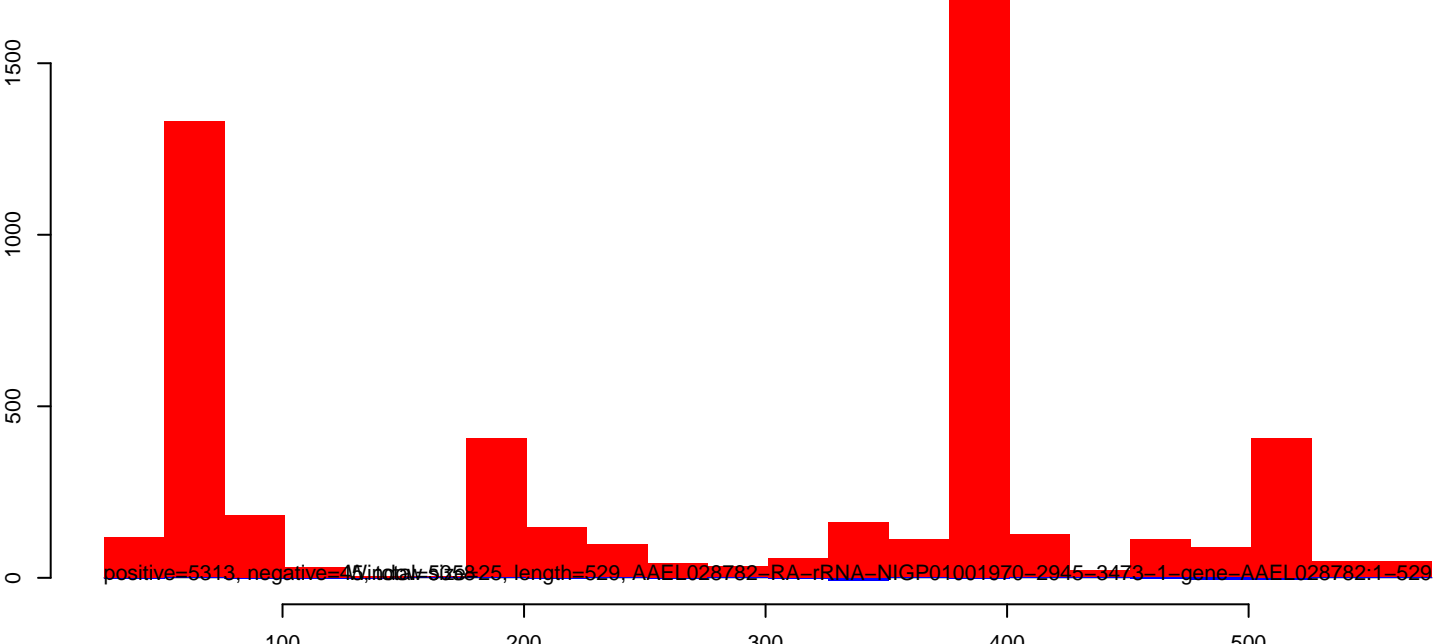
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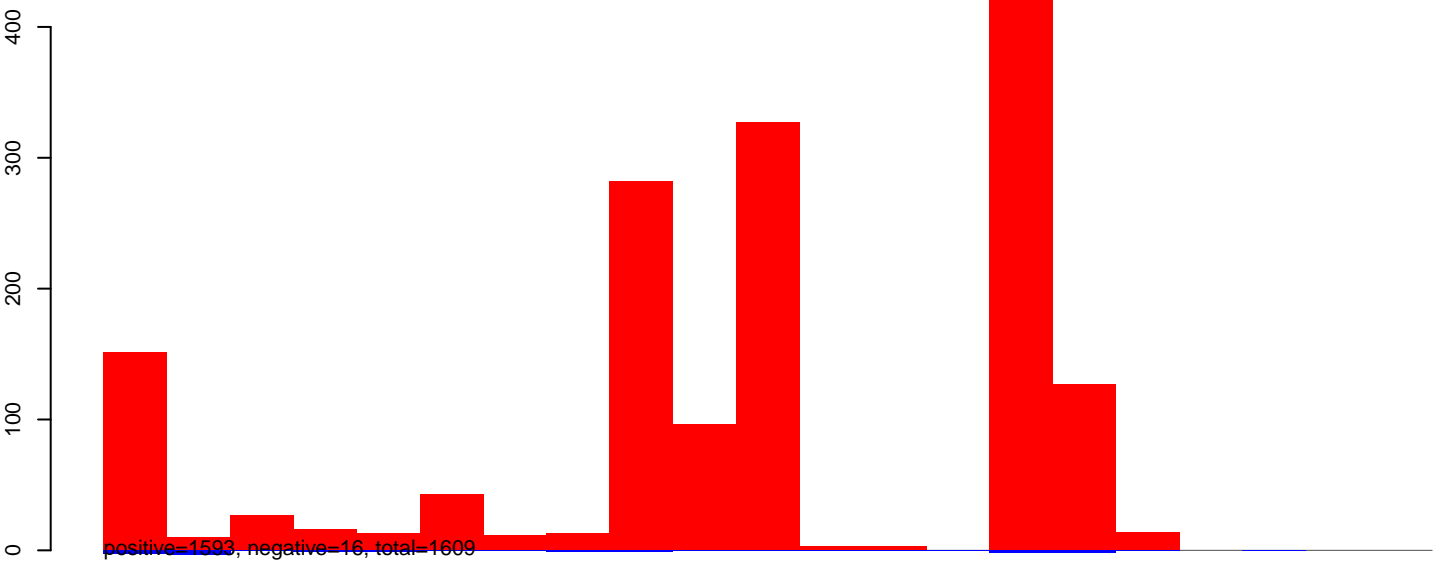
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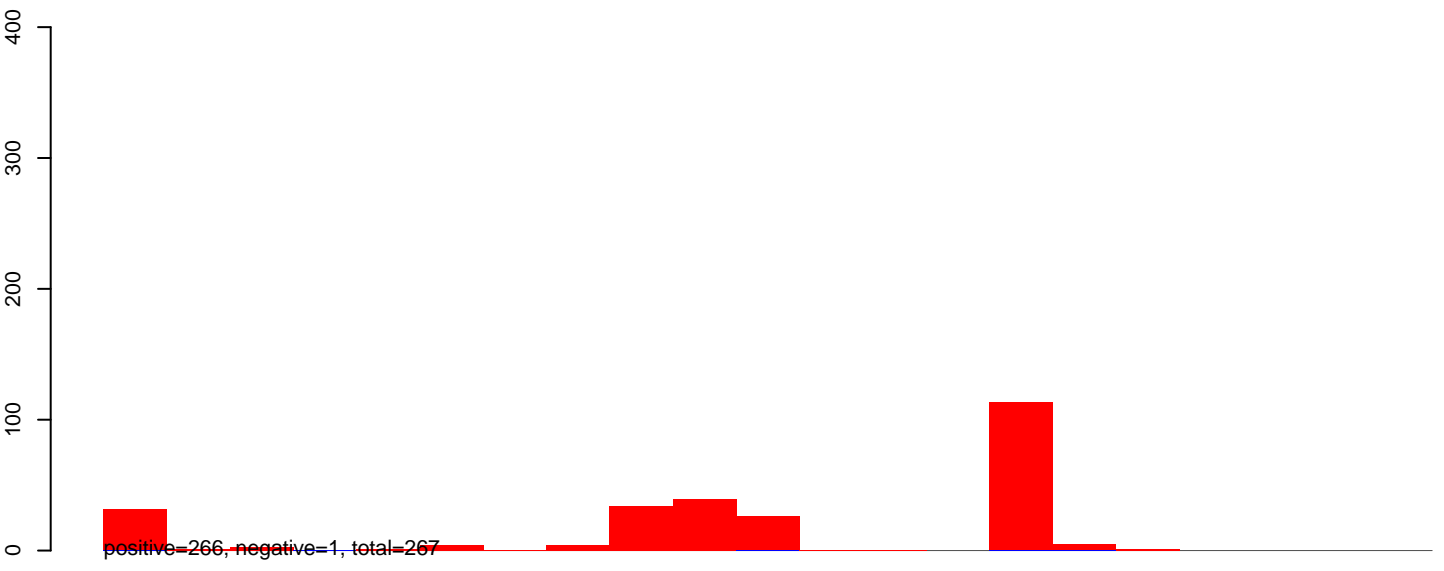
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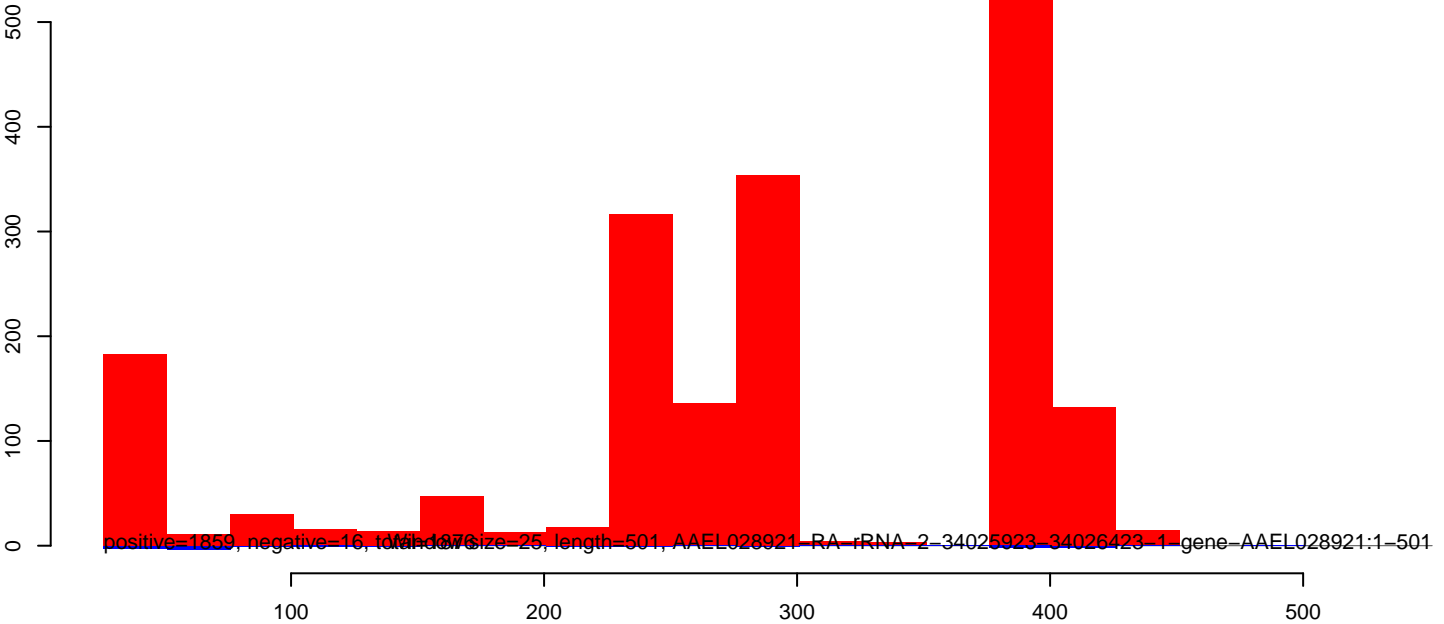
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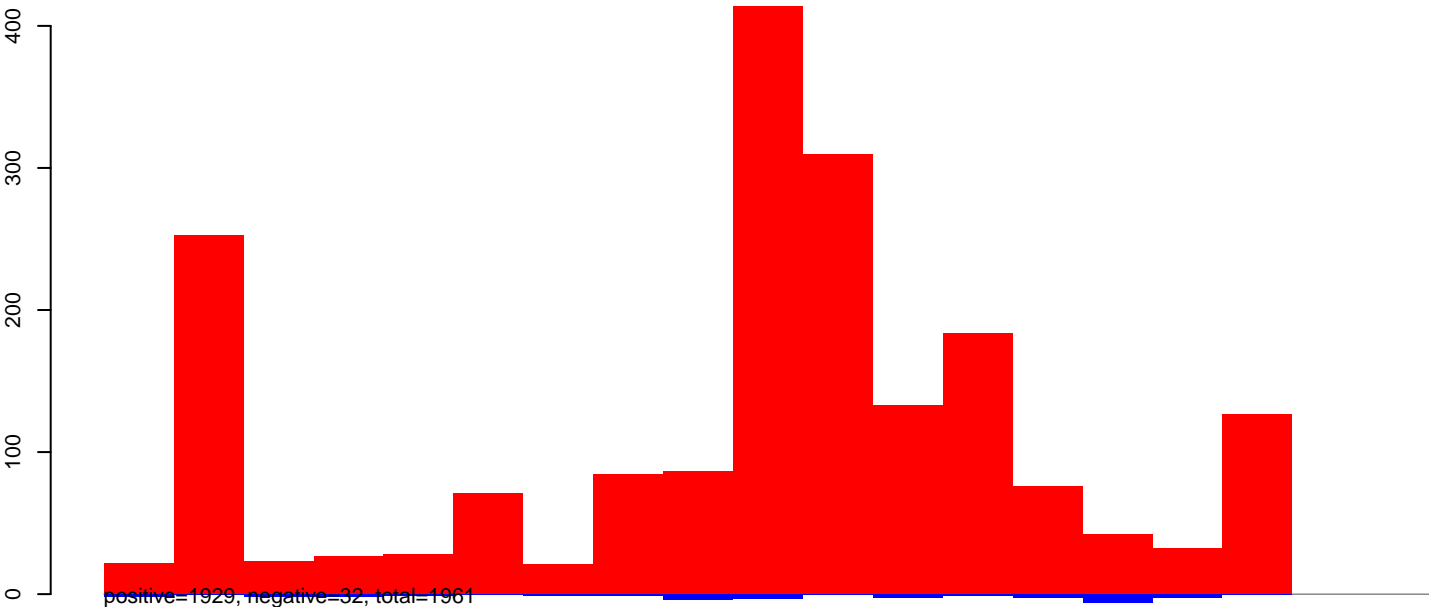
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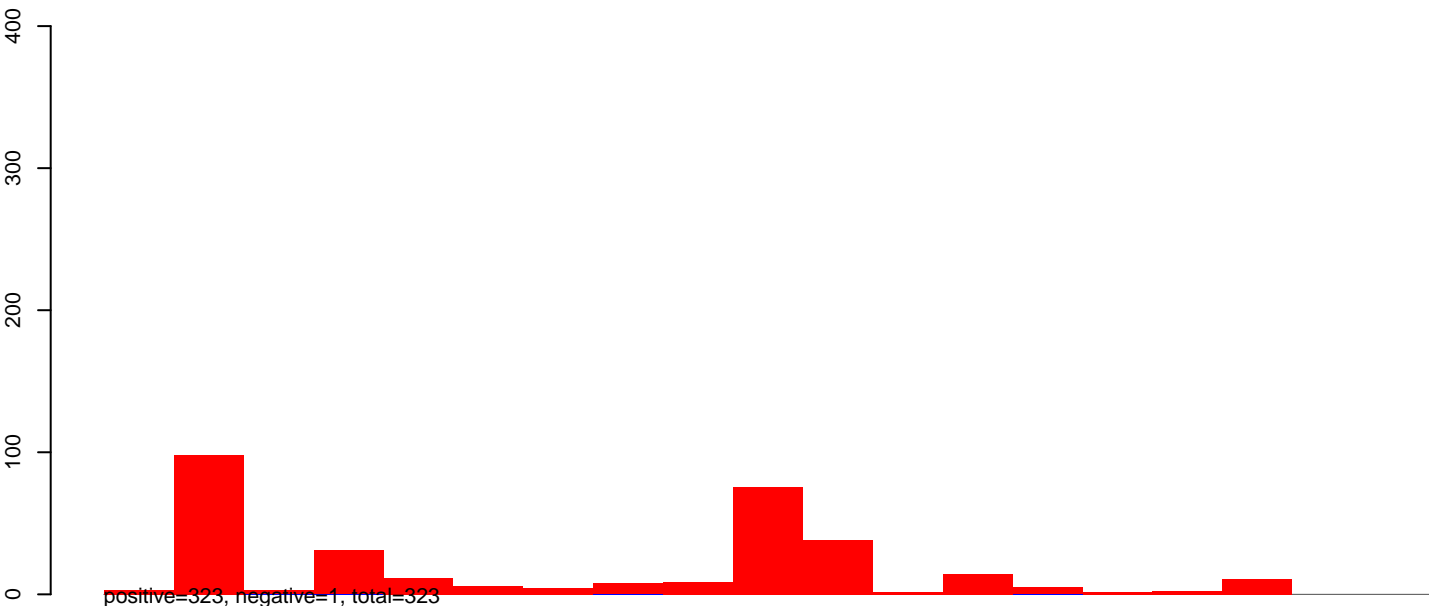
AeAeg_Testes.rep



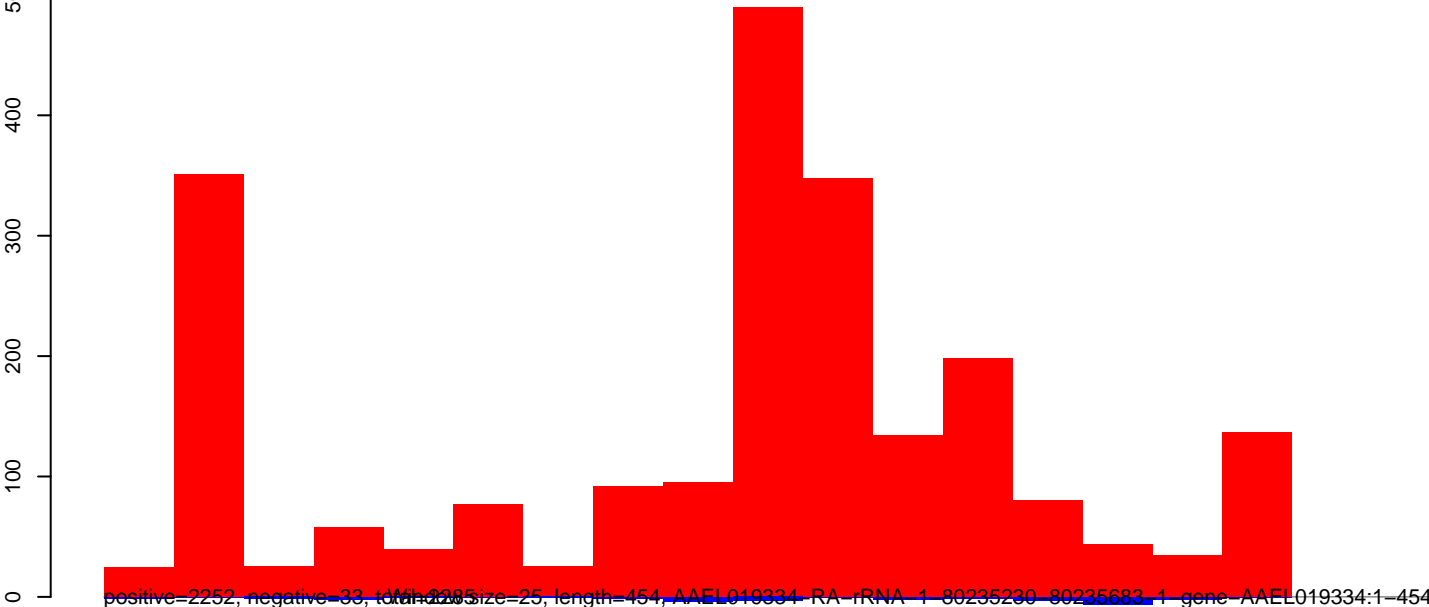
AeAeg_Testes.18_23.rep



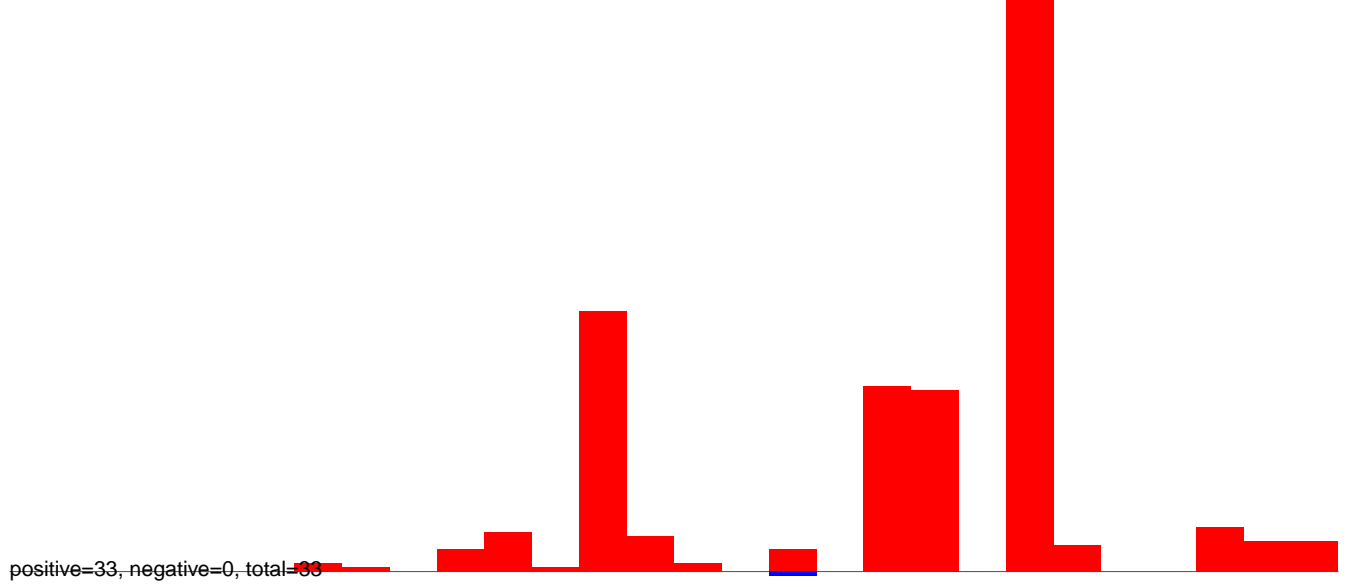
AeAeg_Testes.24_35.rep



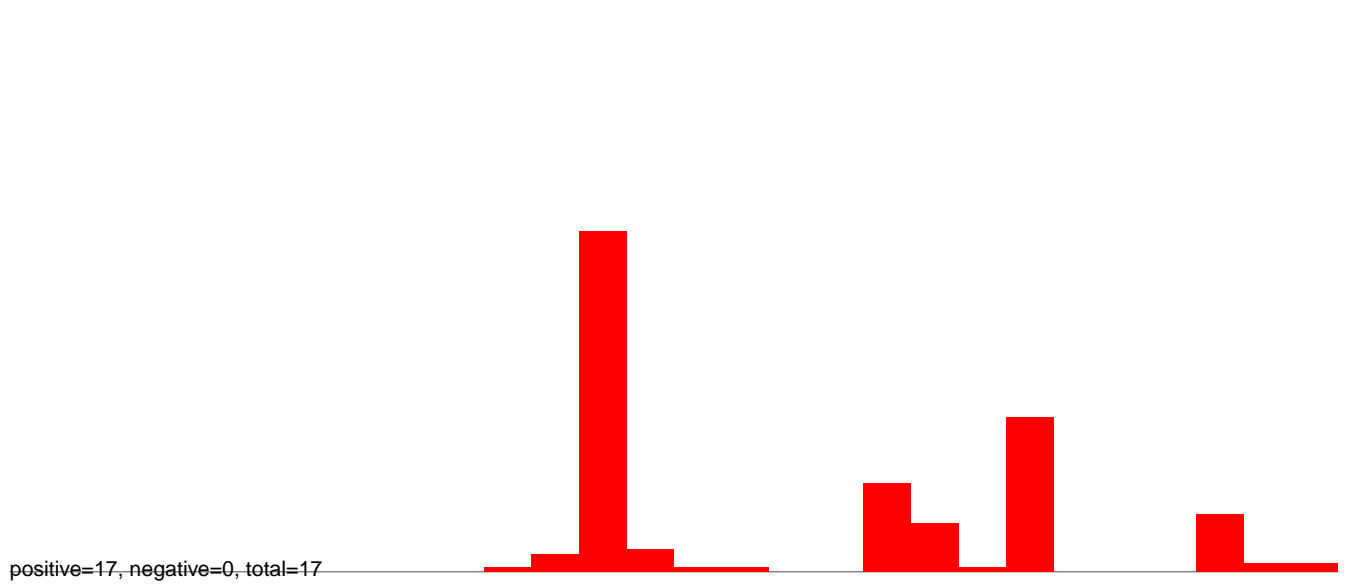
AeAeg_Testes.rep



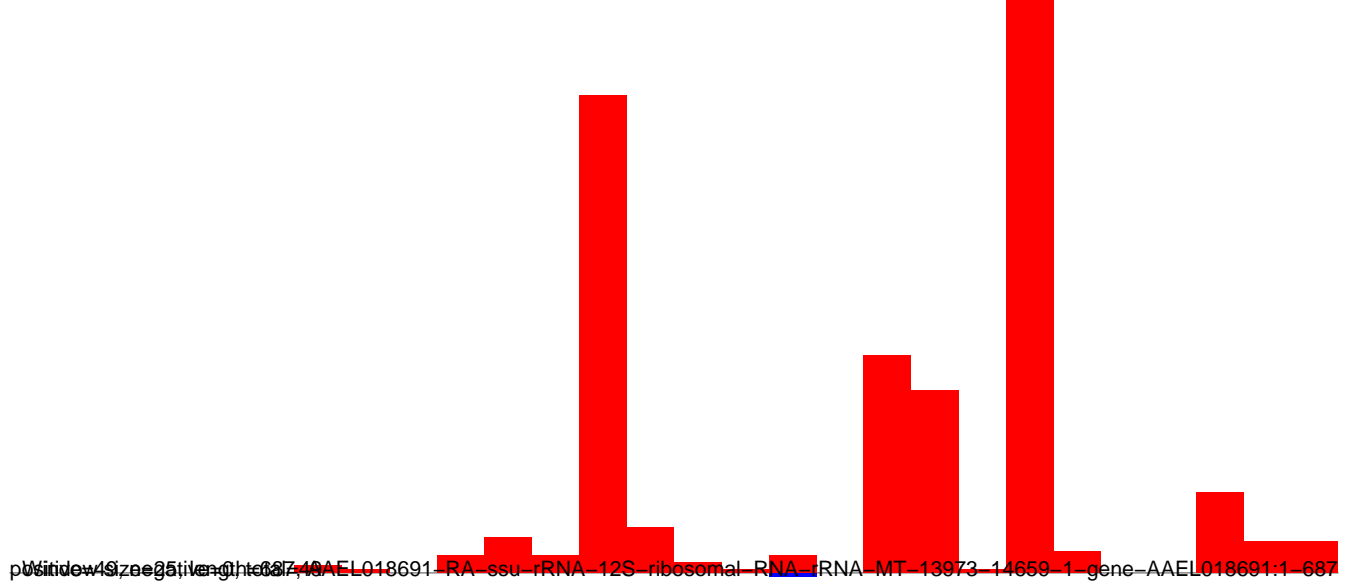
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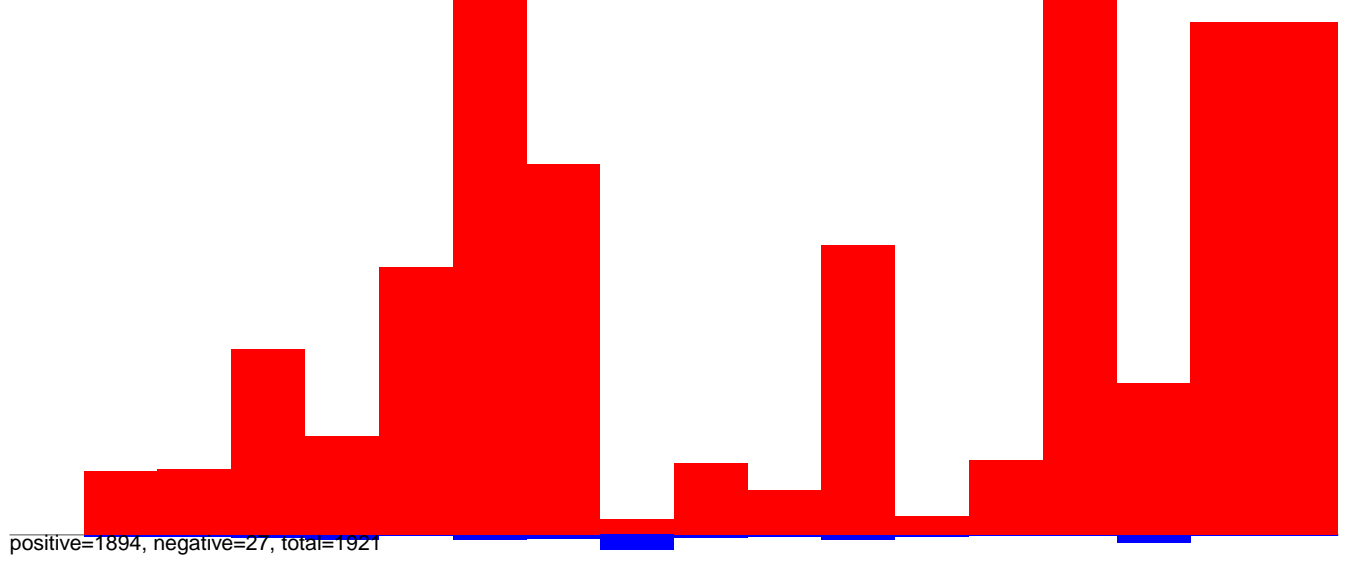
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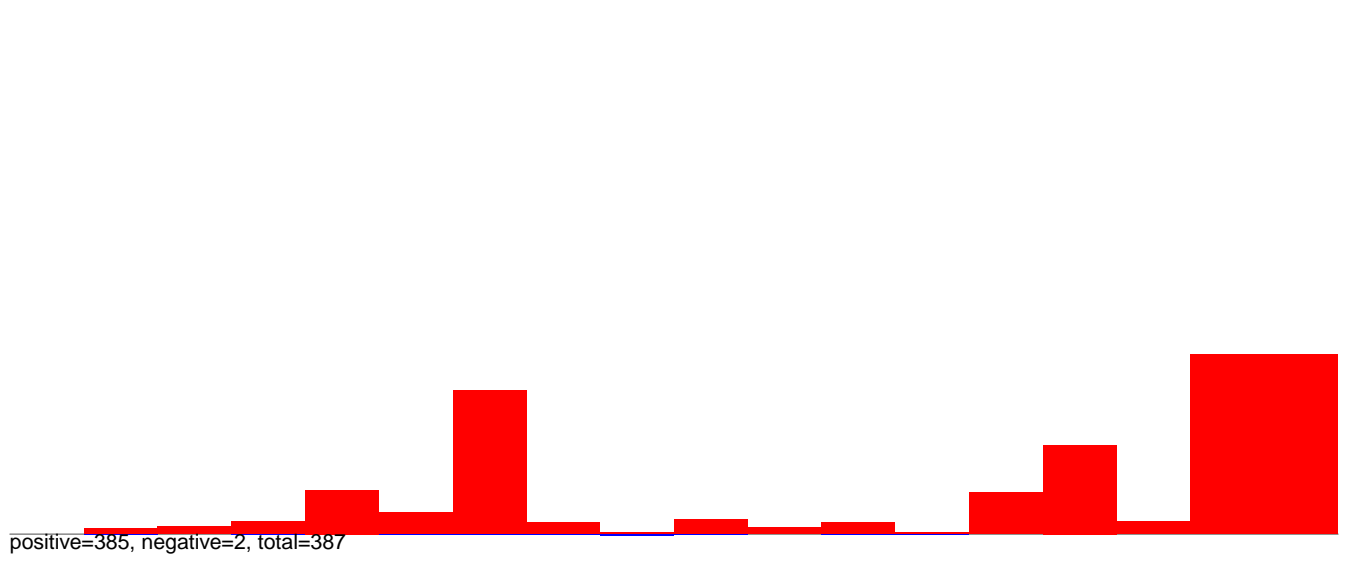
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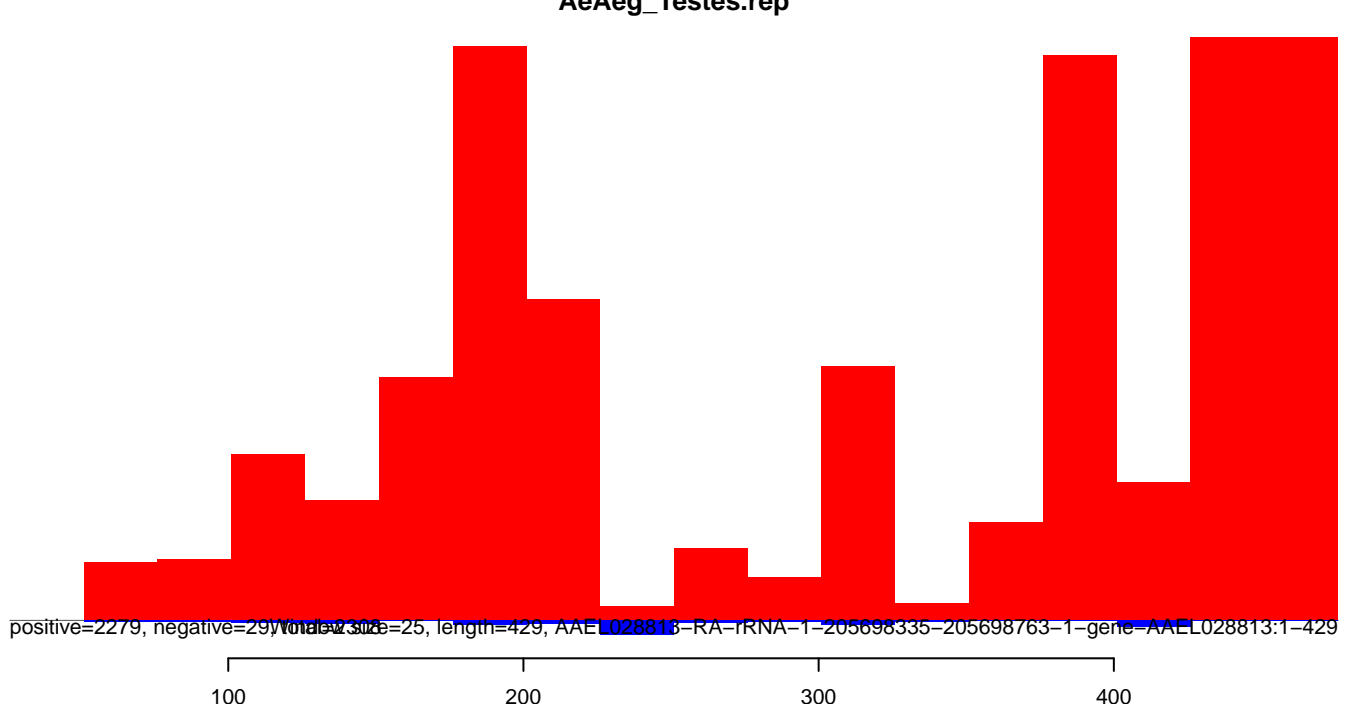
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AeAeg_Testes.24_35.rep

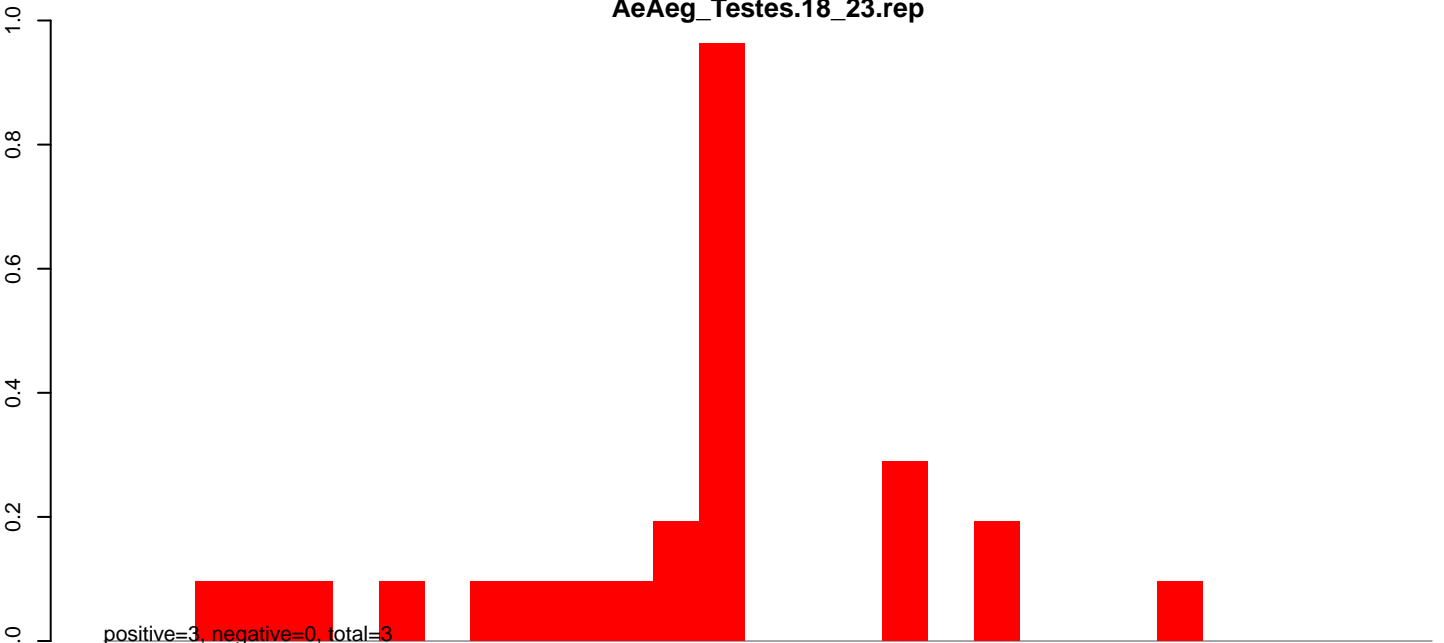


AeAeg_Testes.rep



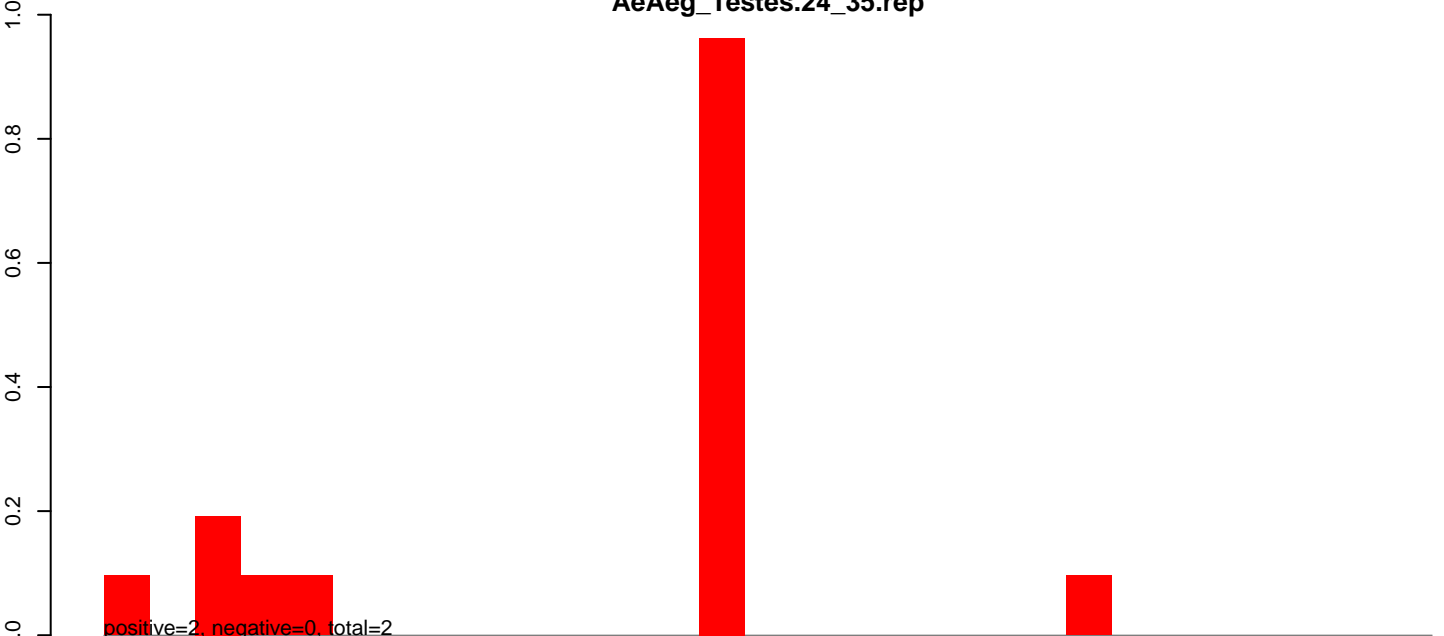
AeAeg_Testes.18_23.rep

positive=3, negative=0, total=3



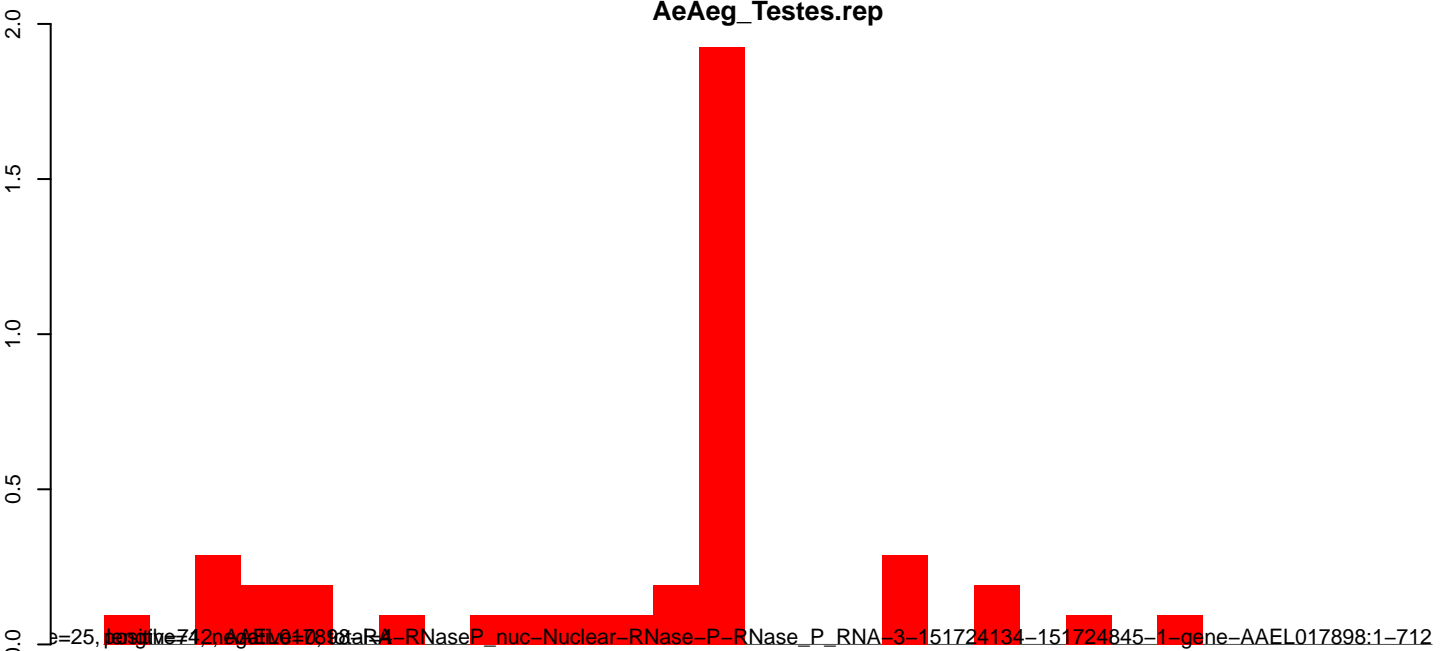
AeAeg_Testes.24_35.rep

positive=2, negative=0, total=2

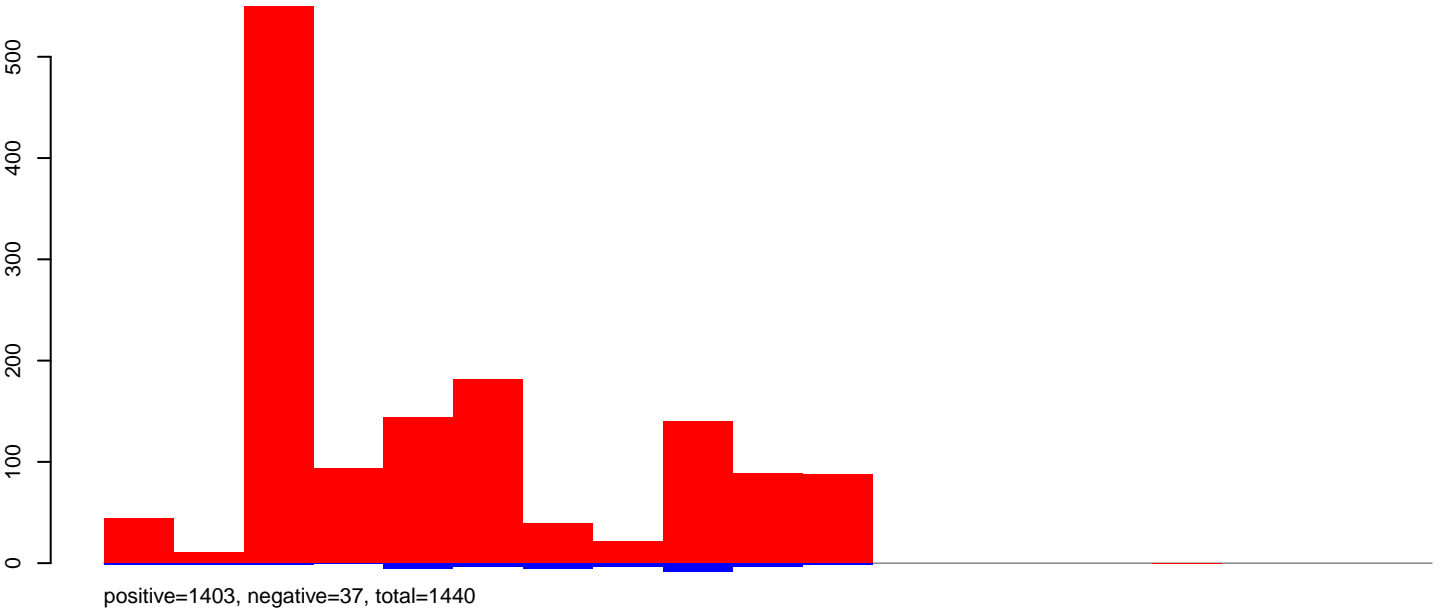


AeAeg_Testes.rep

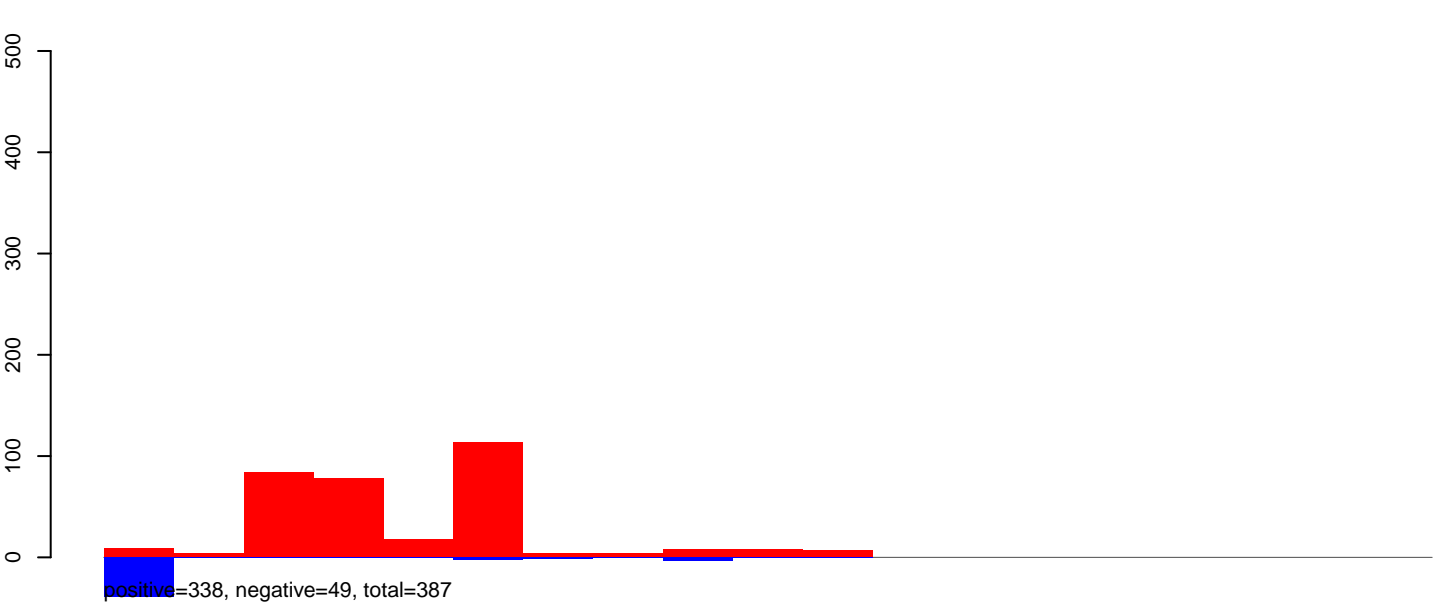
b=25, positive=74, negative=0, total=74
AeAeg_Testes.18_23.rep - RNaseP_nuc - Nuclear - RNase_P - RNase_P_RNA - 3-151724134-151724845-1-gene-AAEL017898:1-712



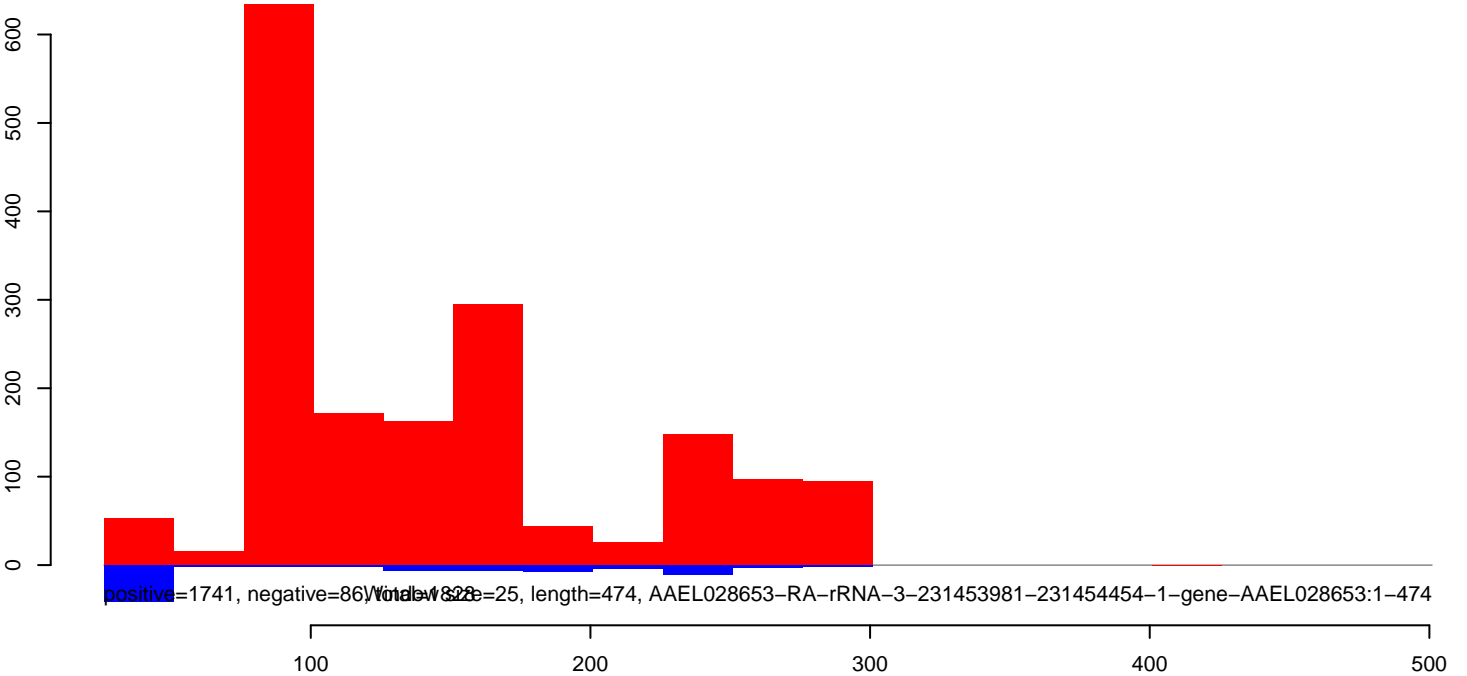
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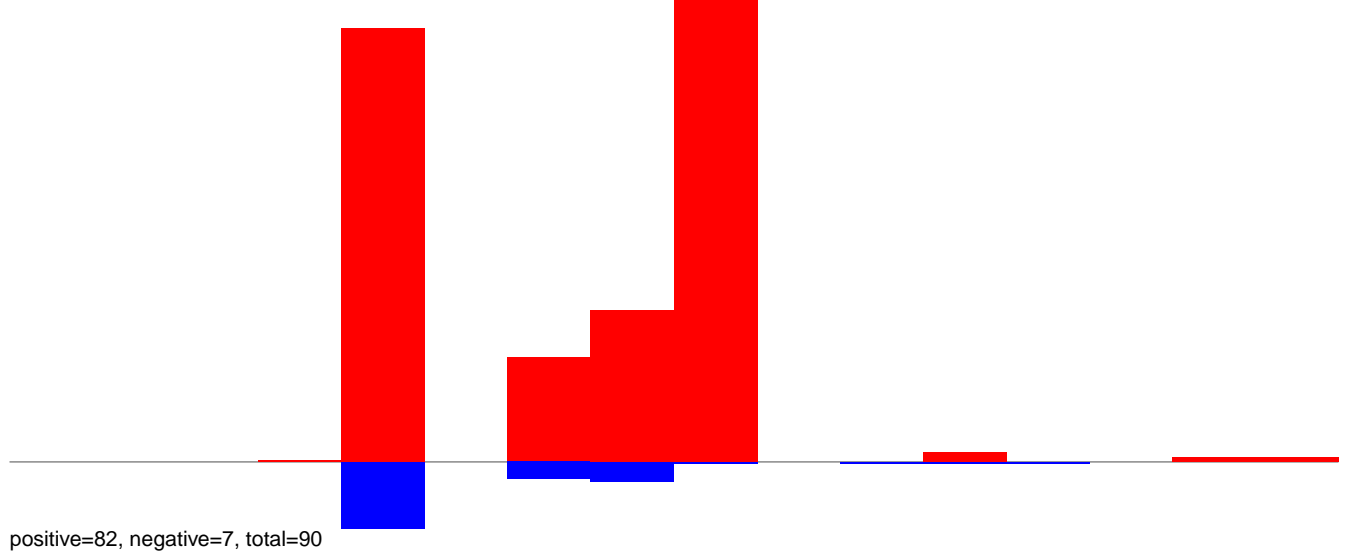
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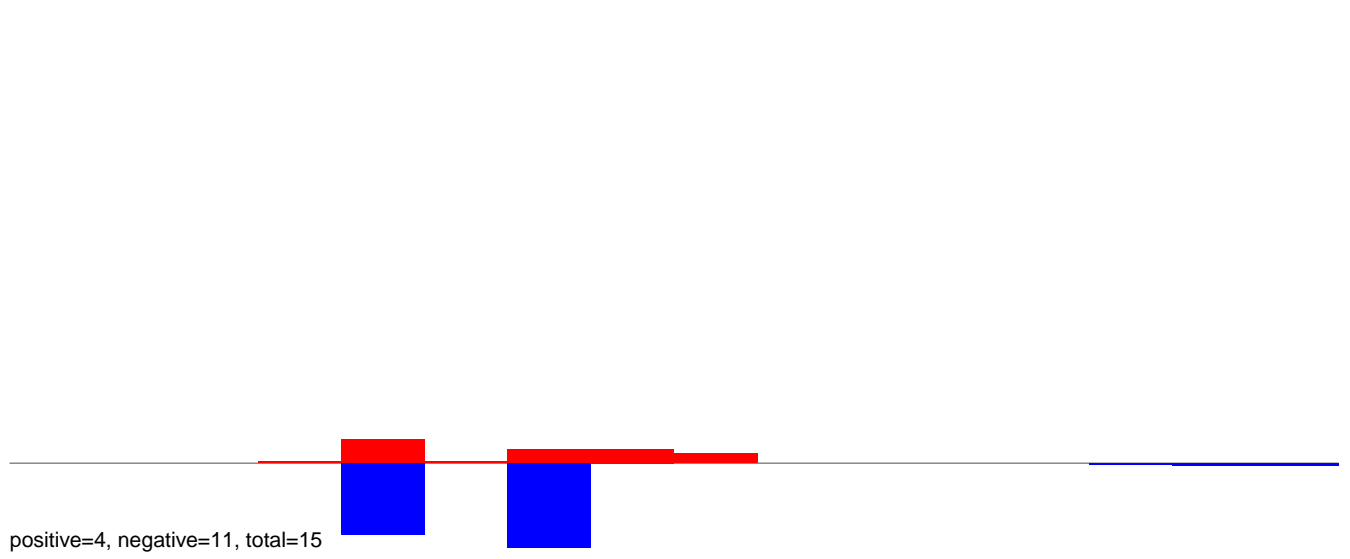
AeAeg_Testes.rep



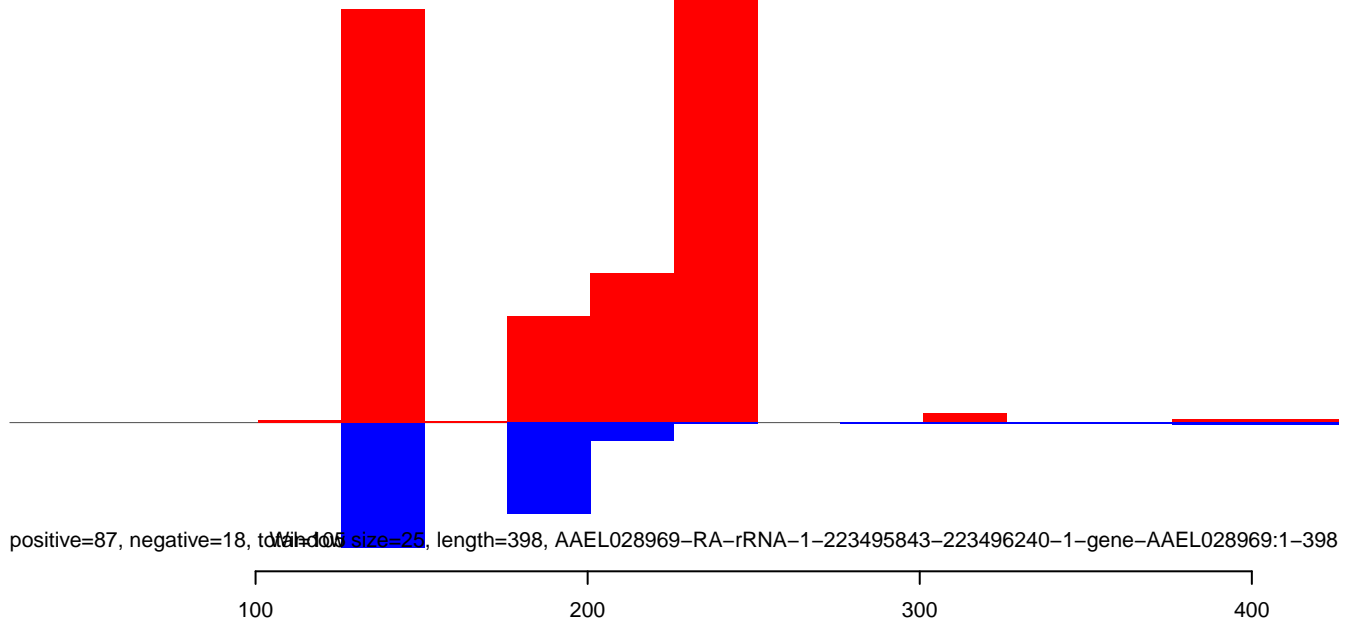
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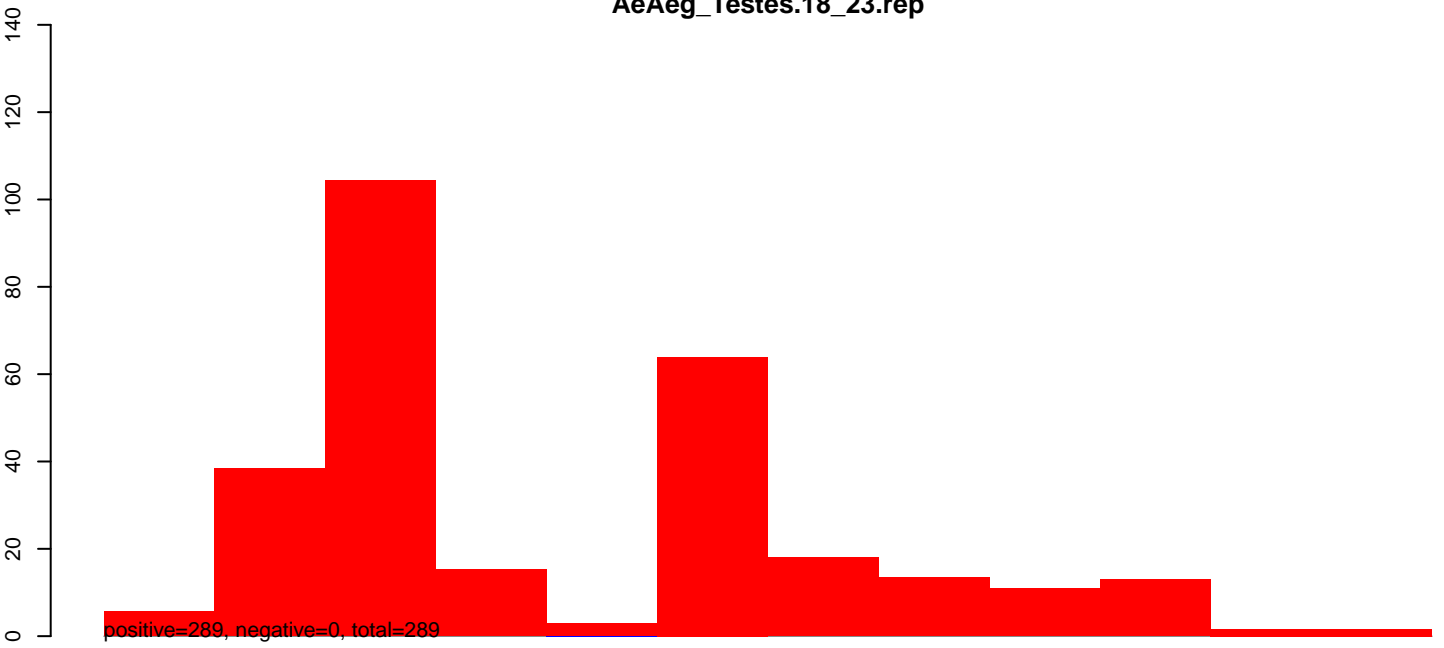
AeAeg_Testes.24_35.rep



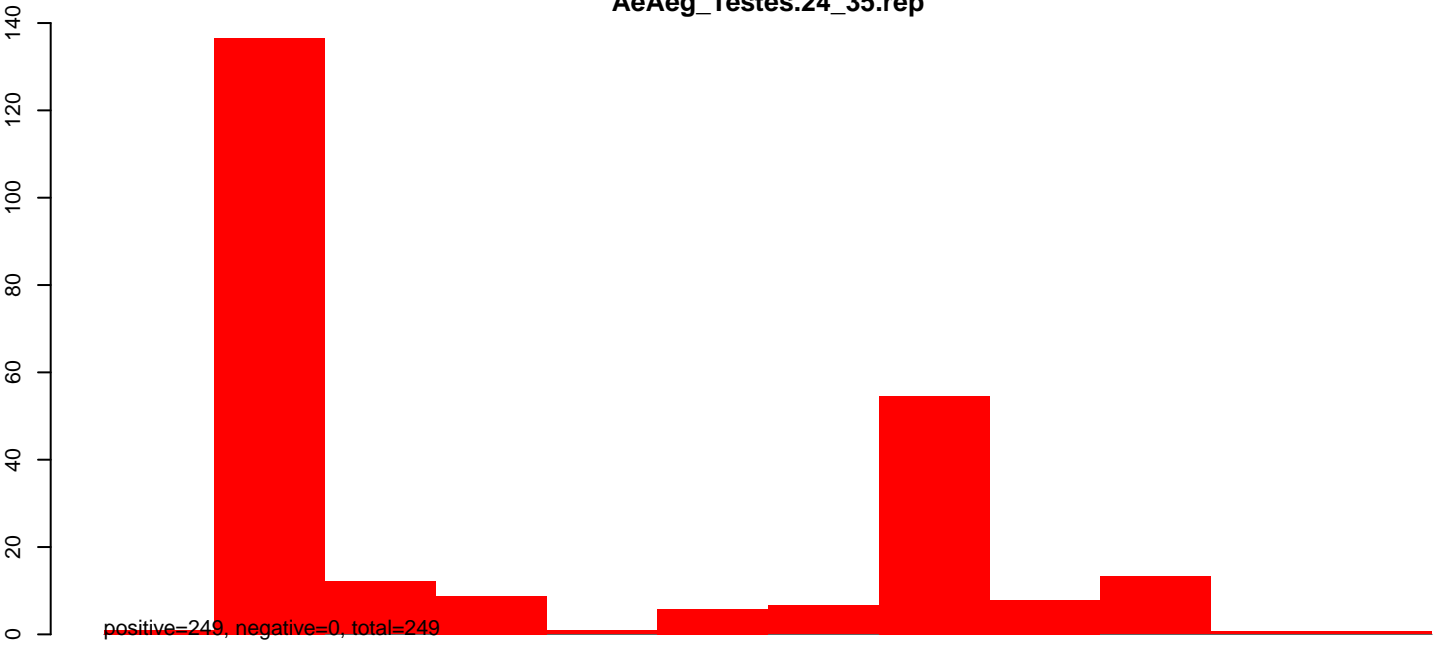
AeAeg_Testes.rep



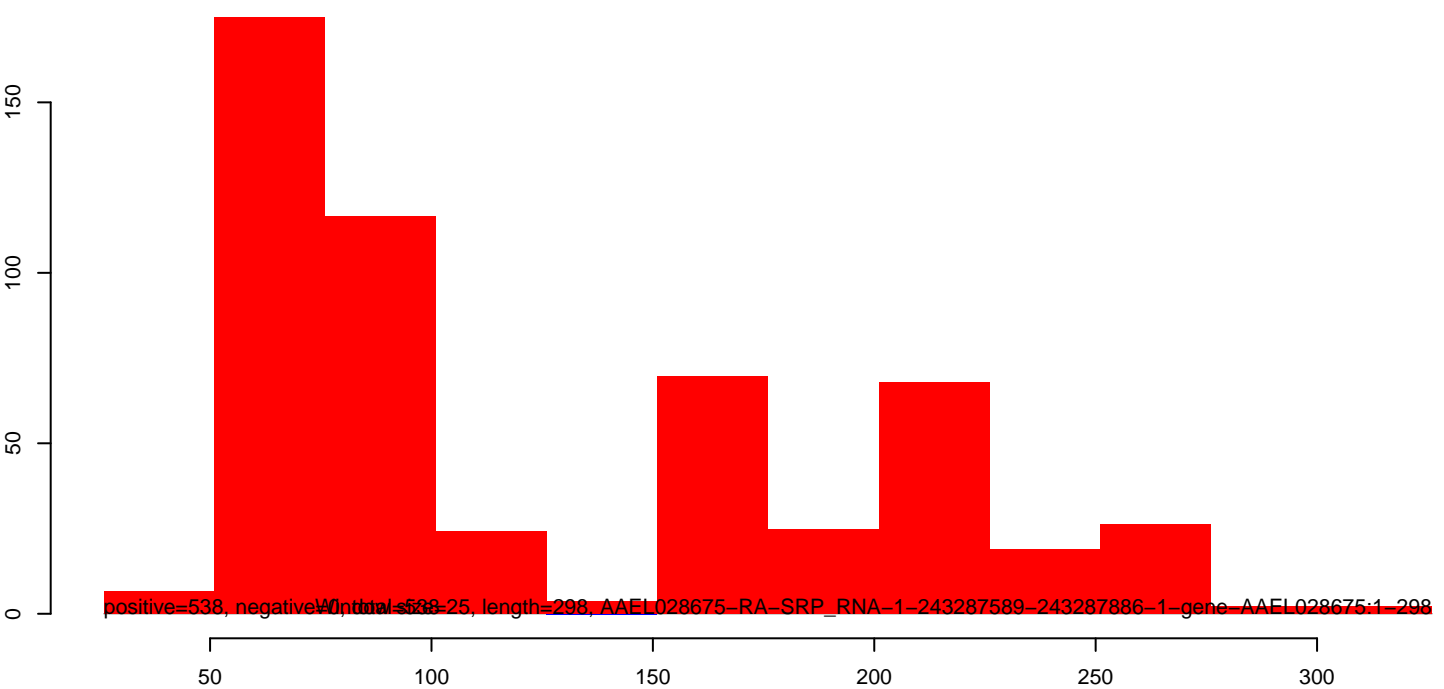
AeAeg_Testes.18_23.rep



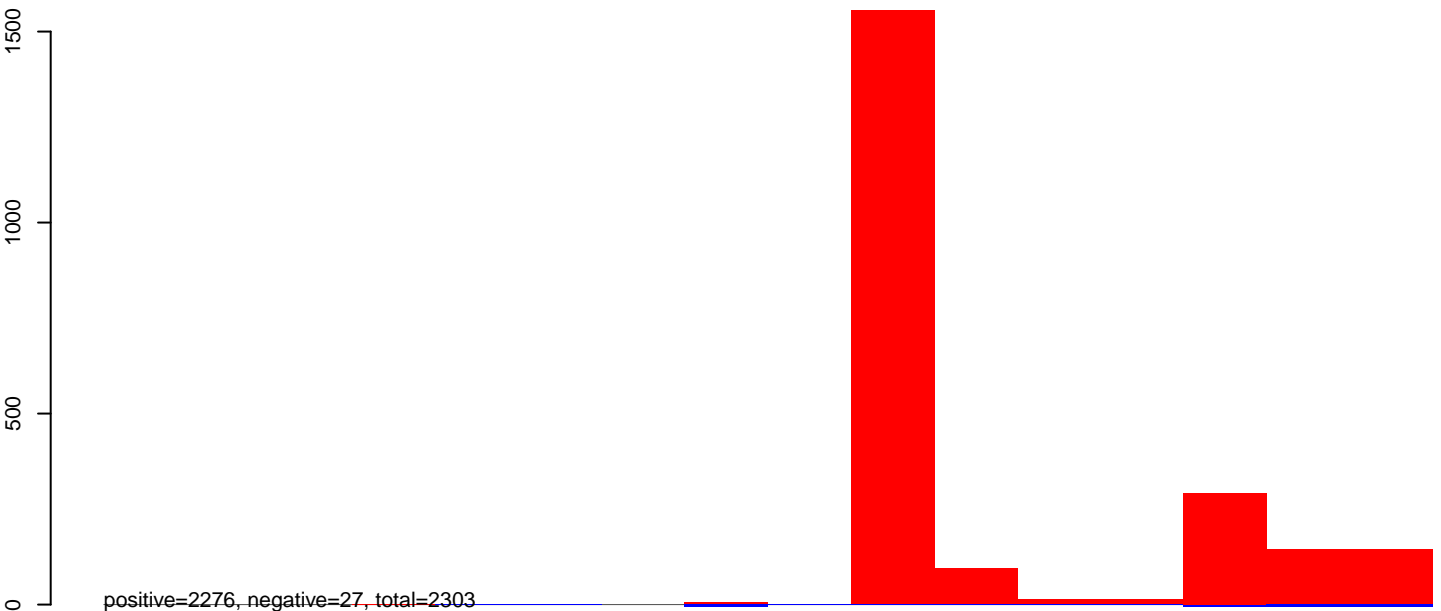
AeAeg_Testes.24_35.rep



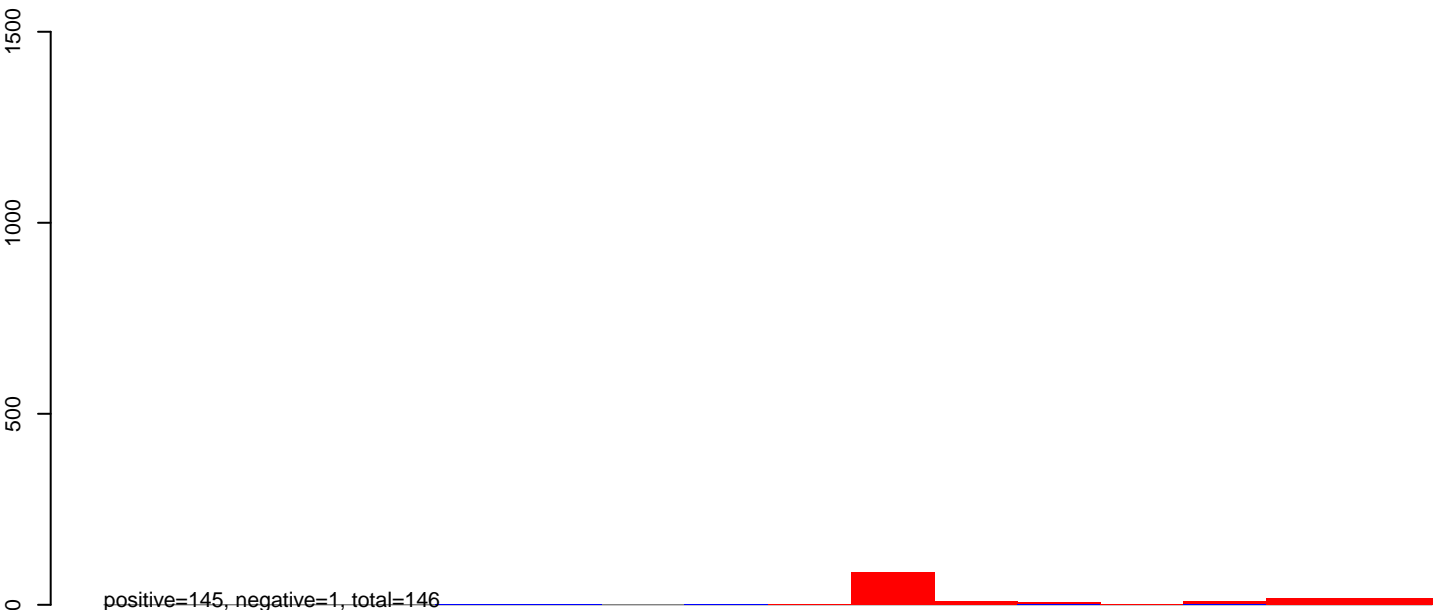
AeAeg_Testes.rep



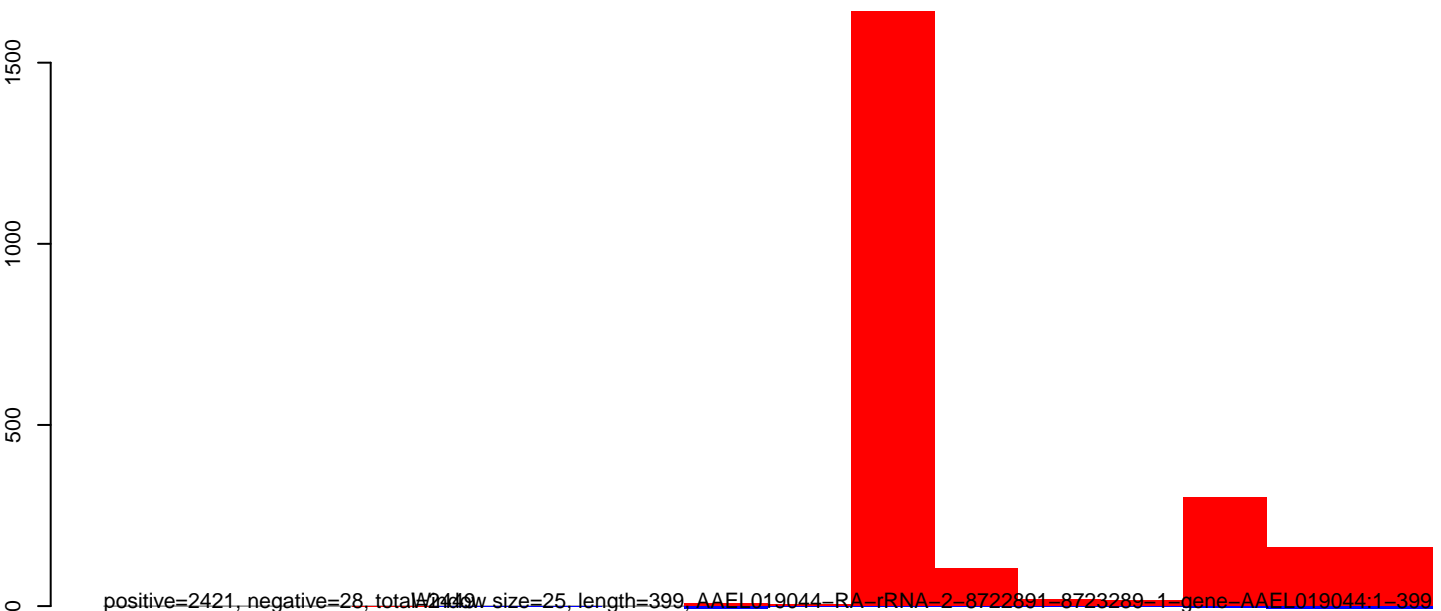
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep

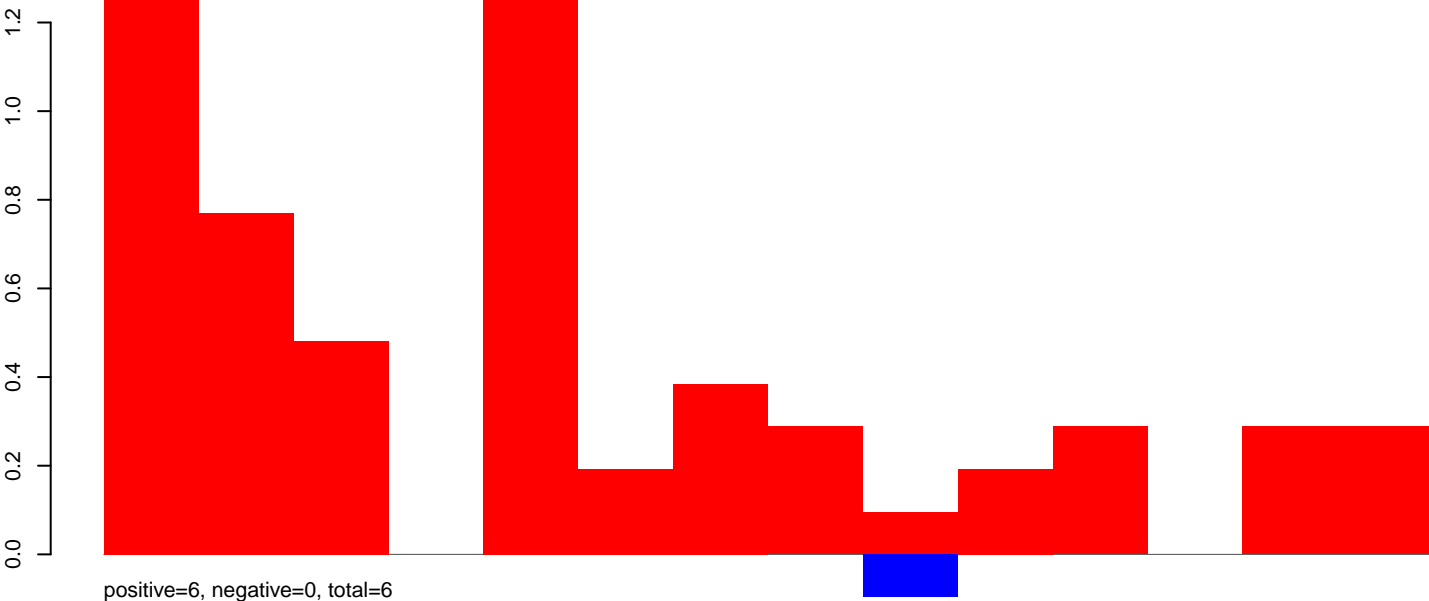


AeAeg_Testes.rep

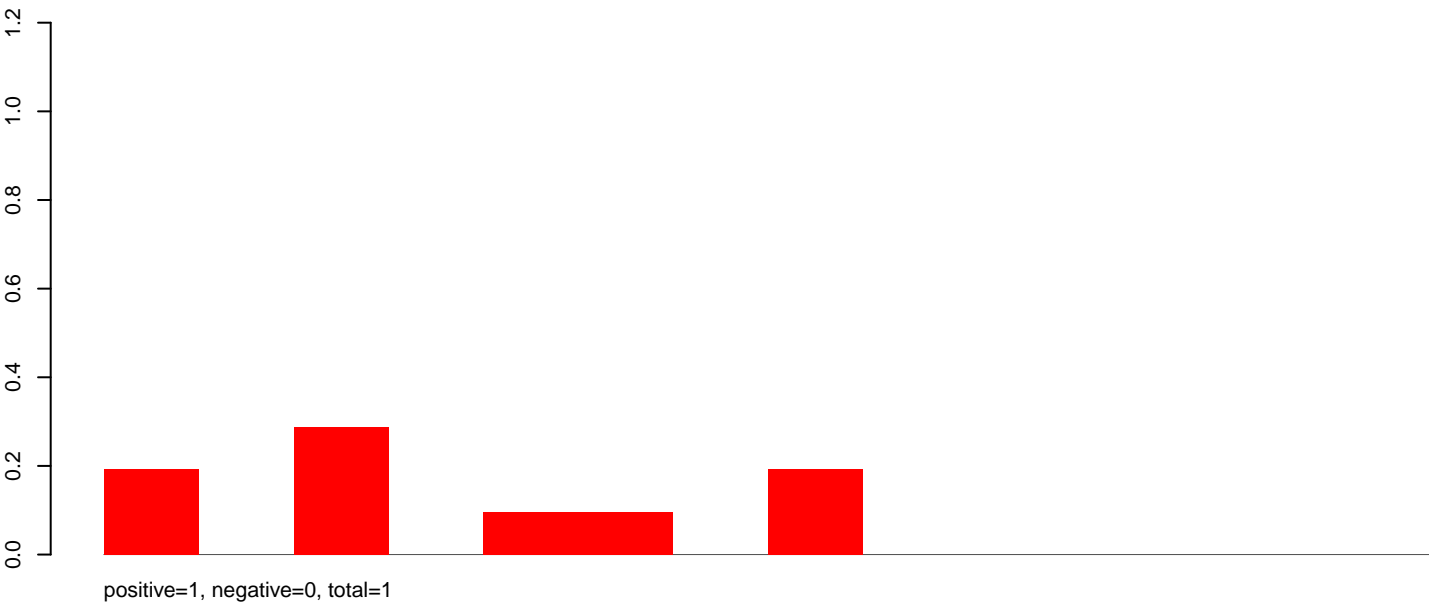


100 200 300 400

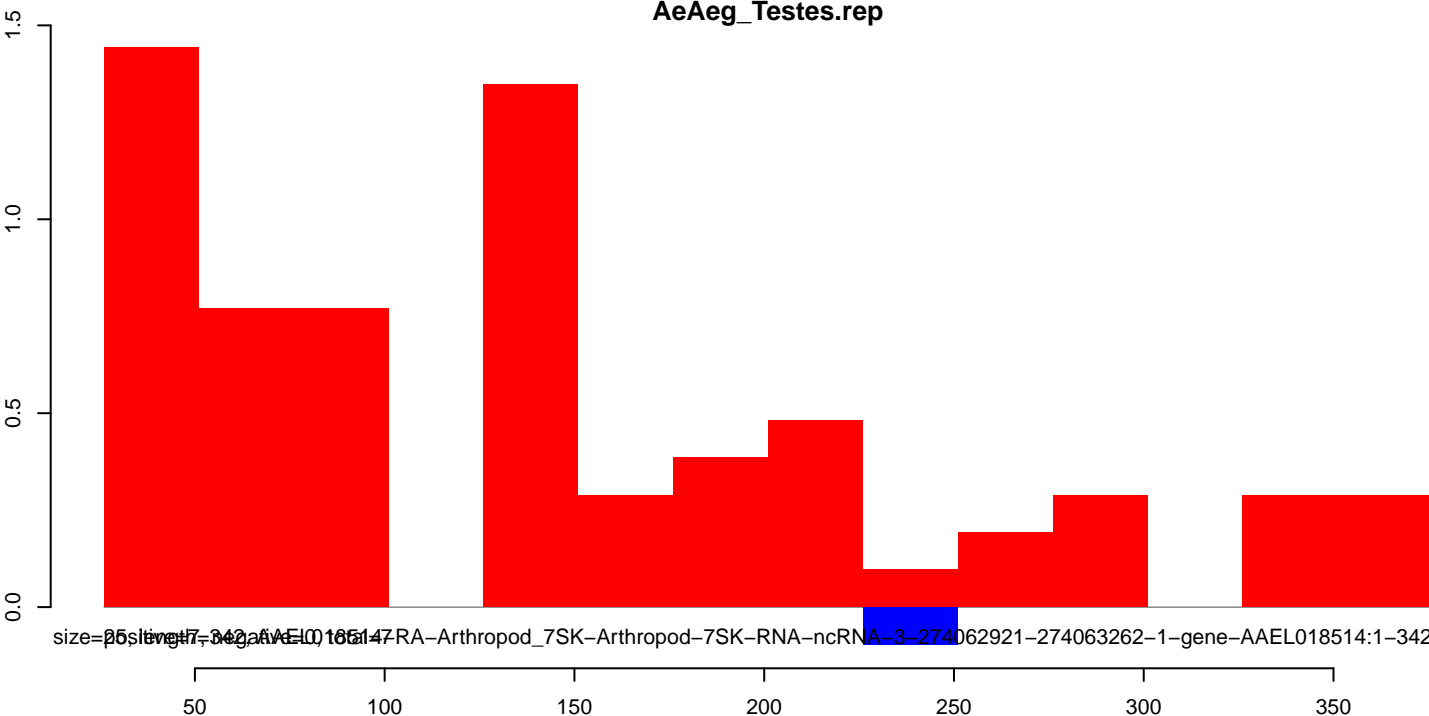
AeAeg_Testes.18_23.rep



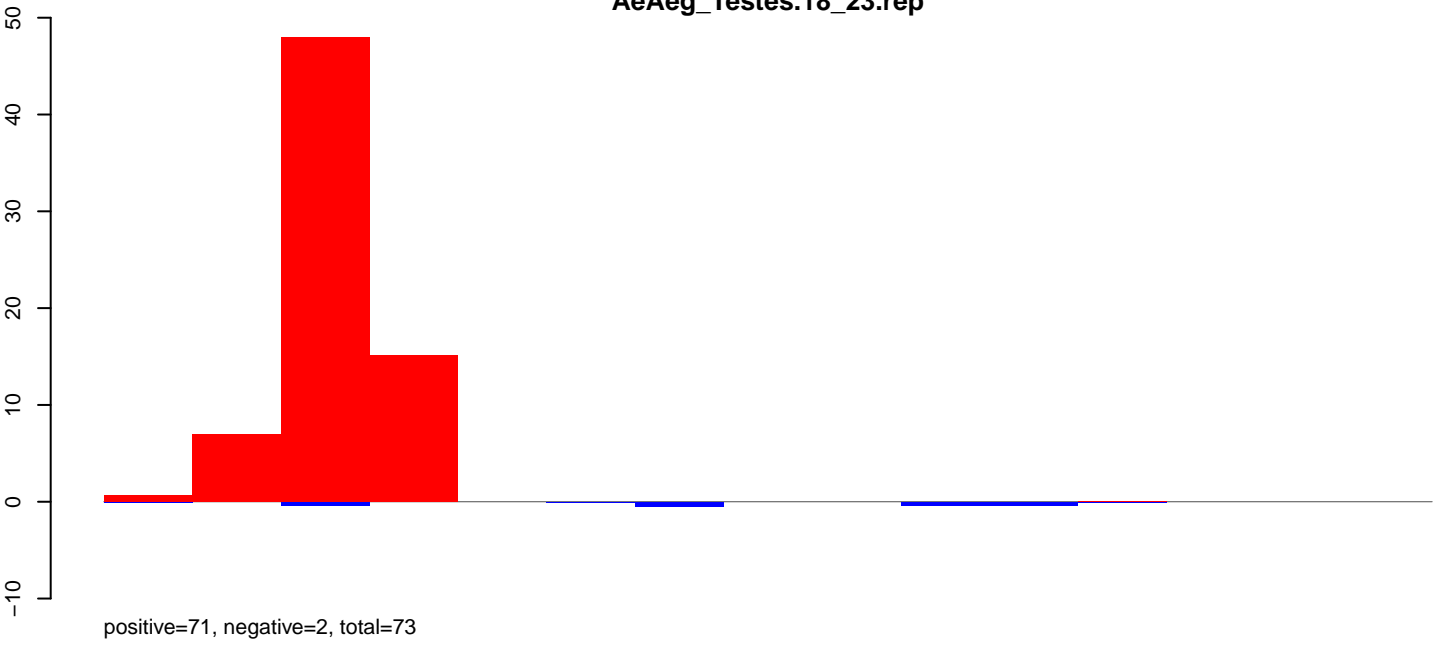
AeAeg_Testes.24_35.rep



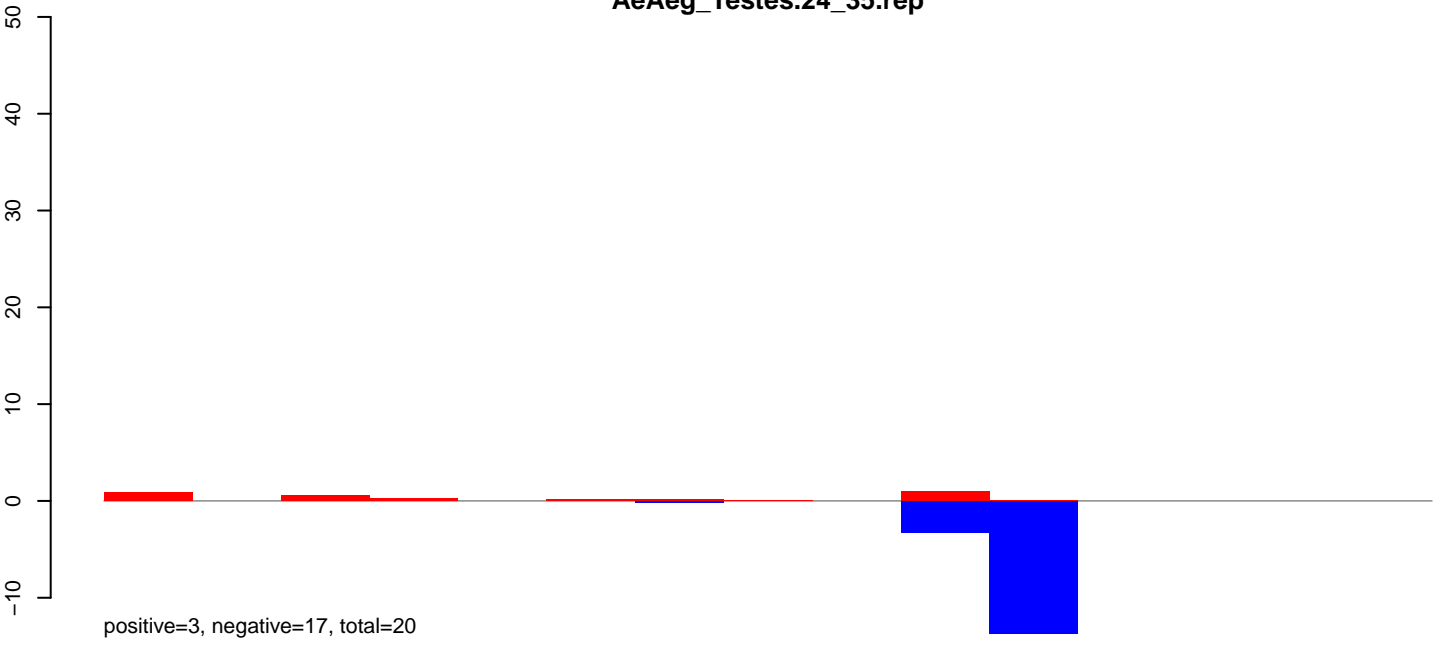
AeAeg_Testes.rep



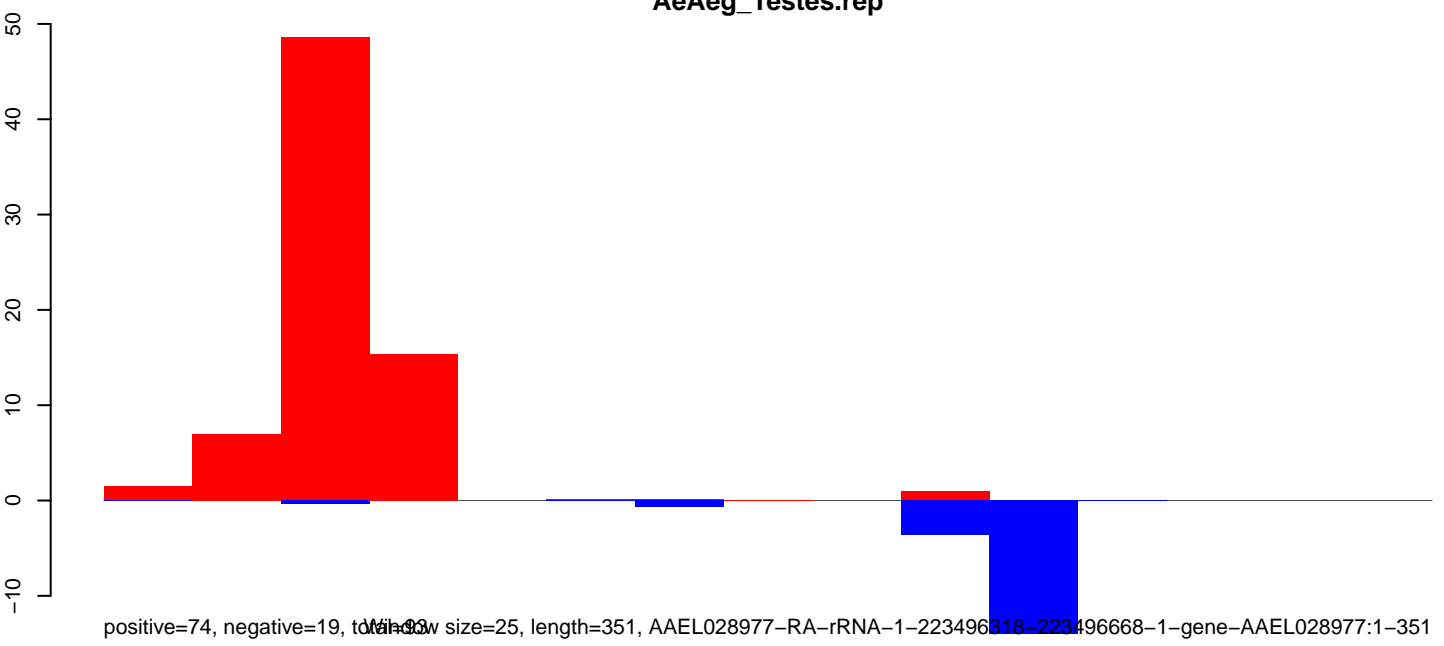
AeAeg_Testes.18_23.rep



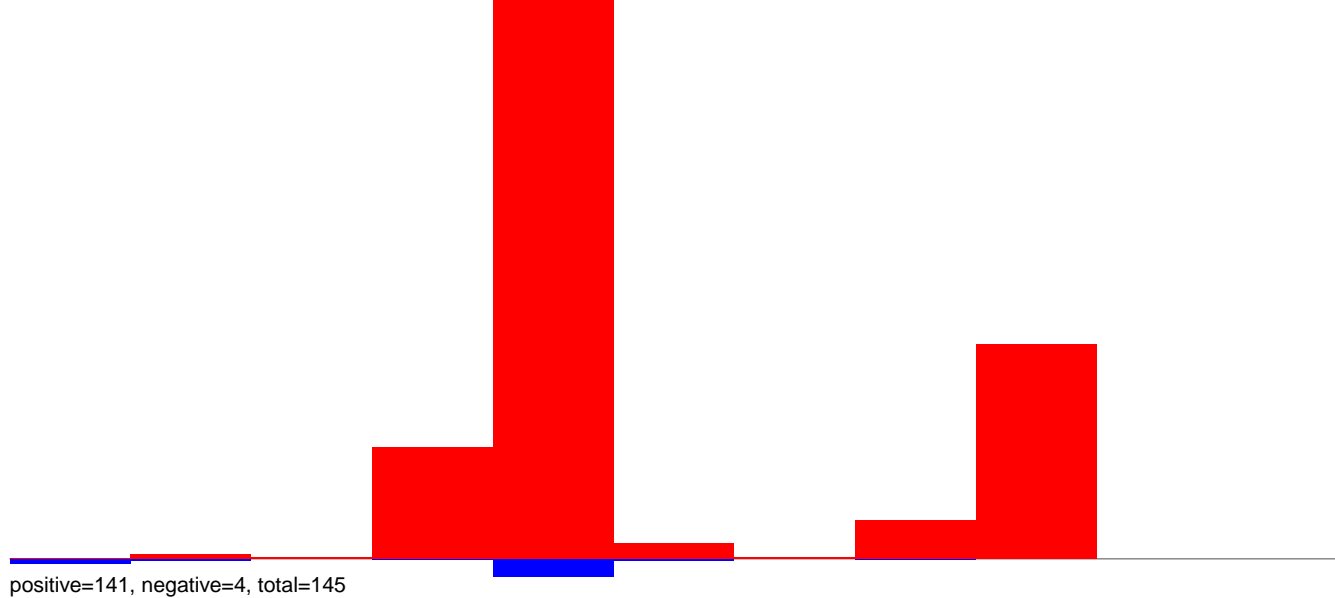
AeAeg_Testes.24_35.rep



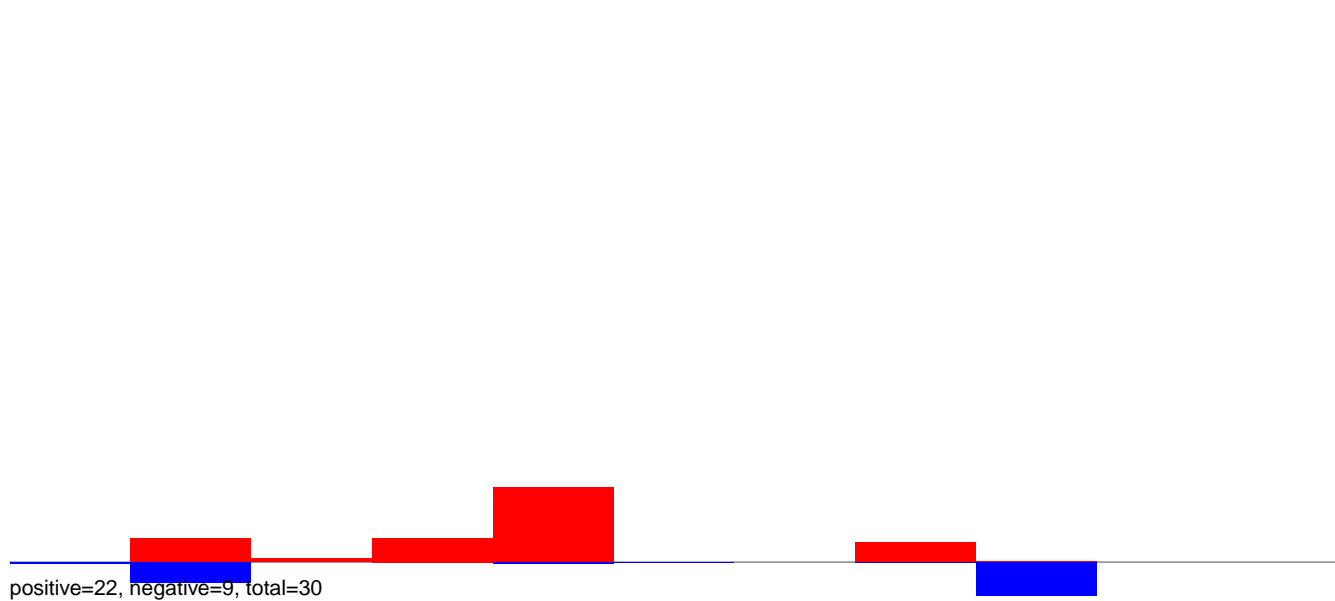
AeAeg_Testes.rep



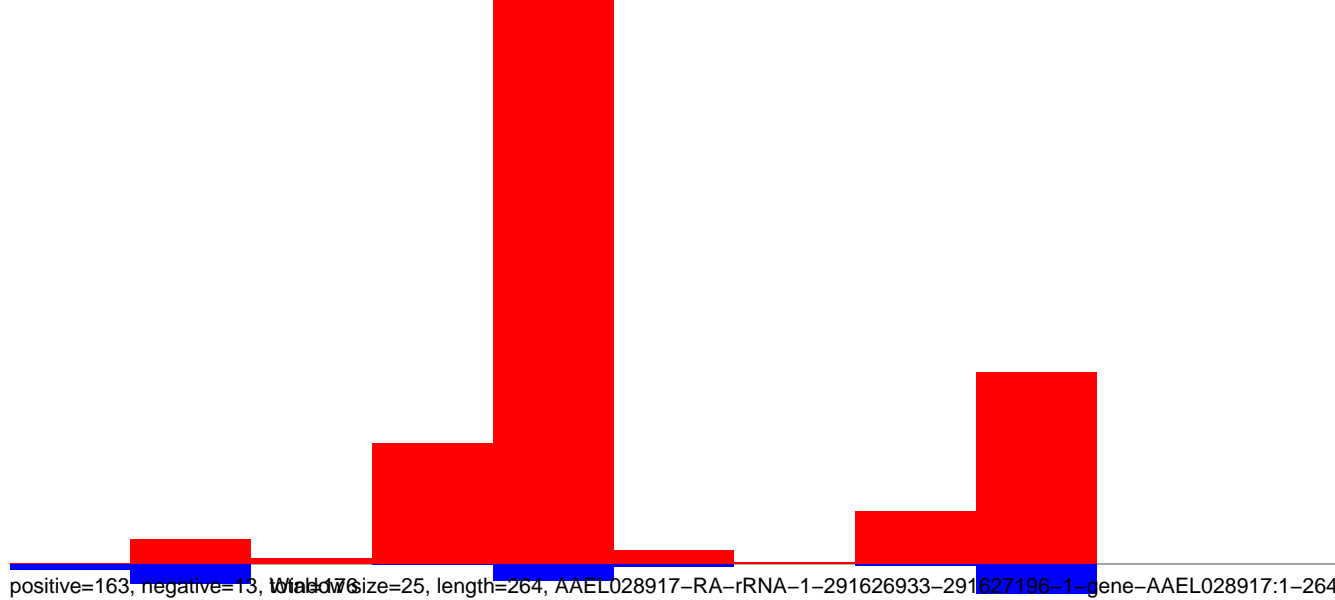
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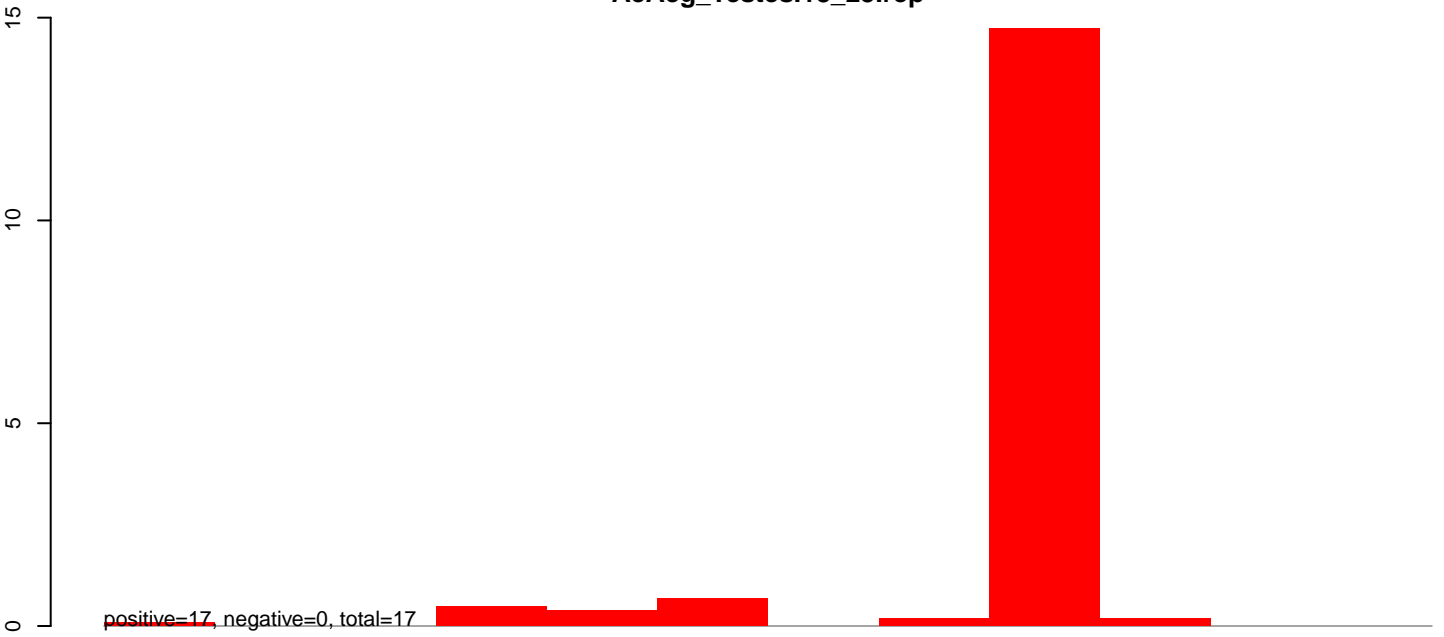
AeAeg_Testes.24_35.rep



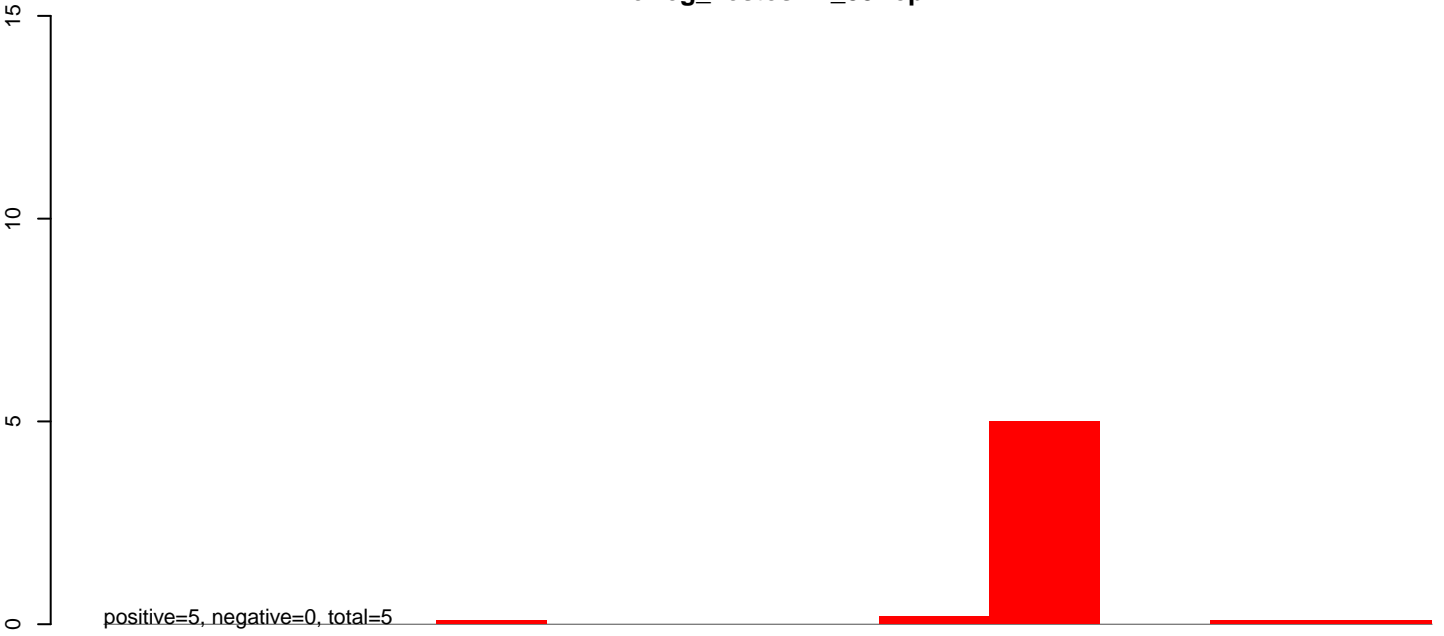
AeAeg_Testes.rep



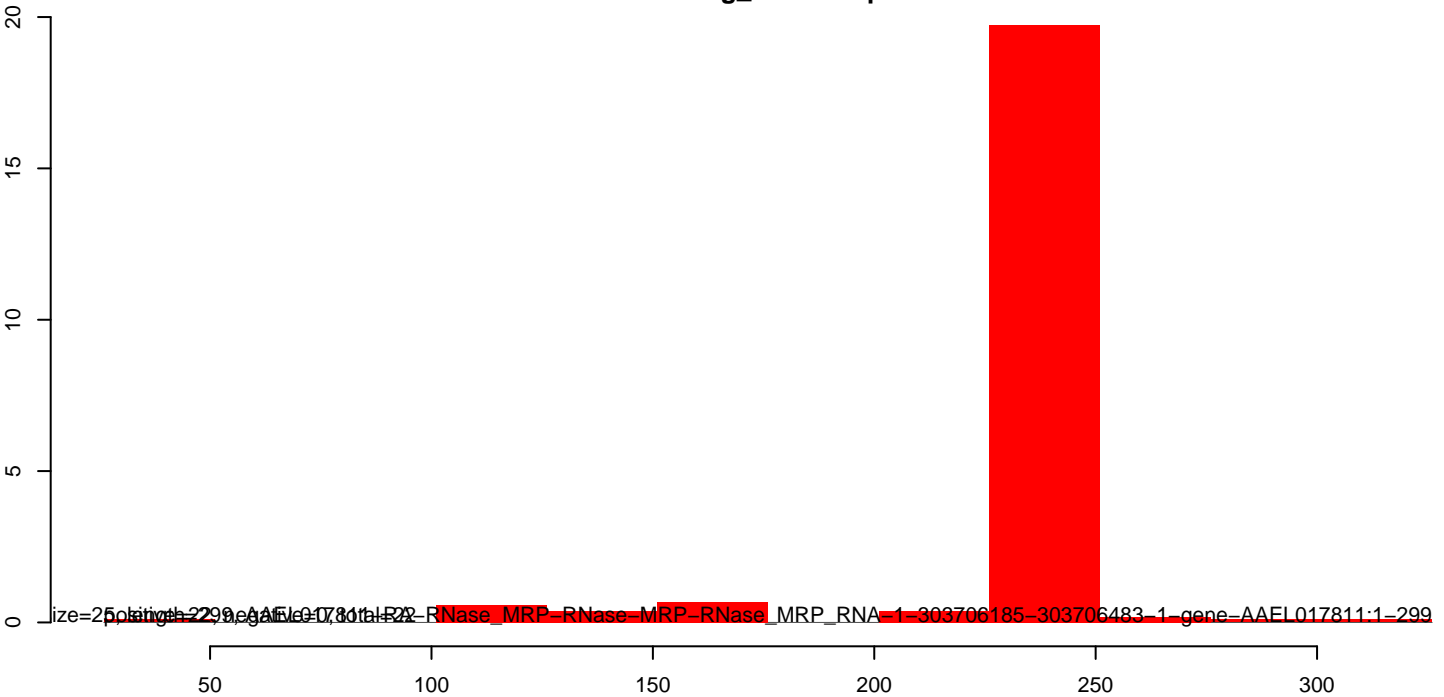
AeAeg_Testes.18_23.rep



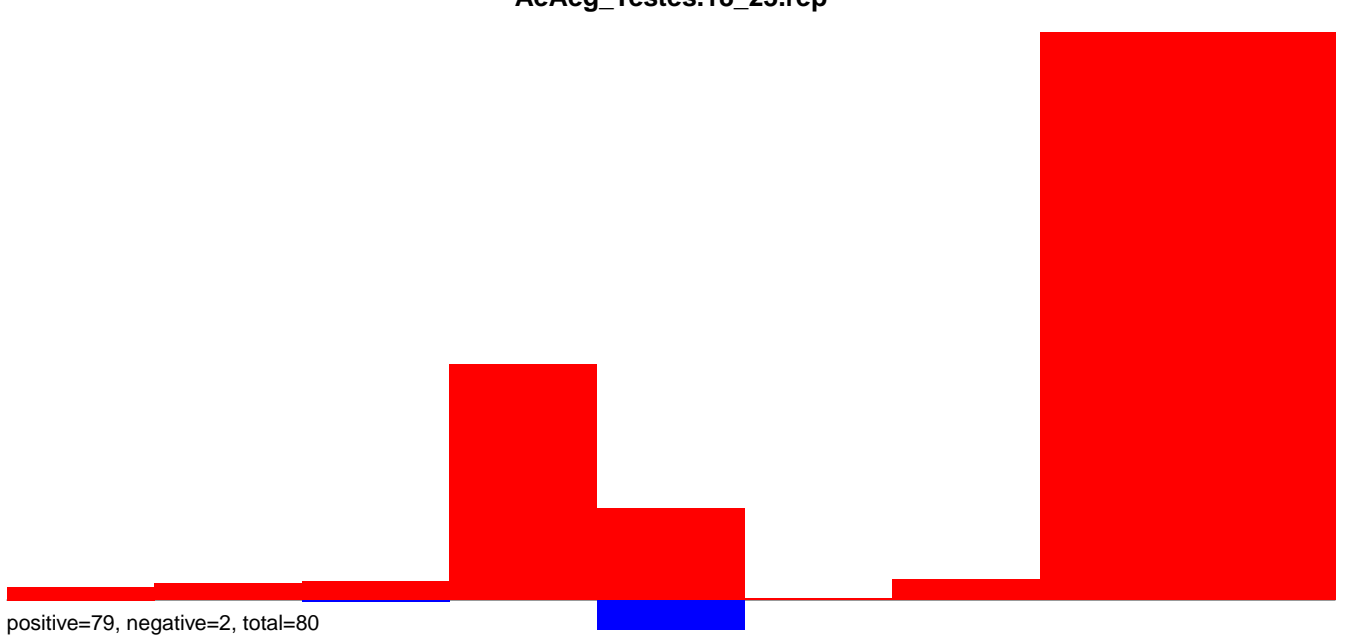
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



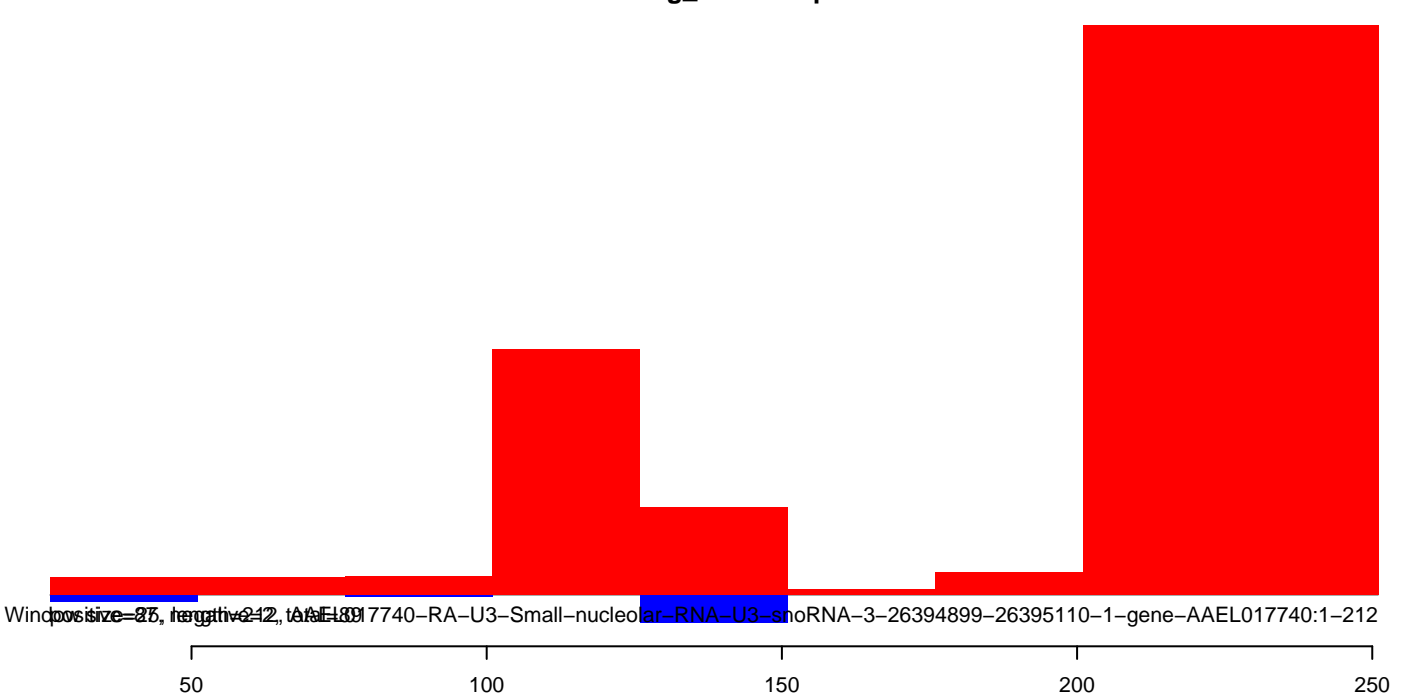
AeAeg_Testes.18_23.rep



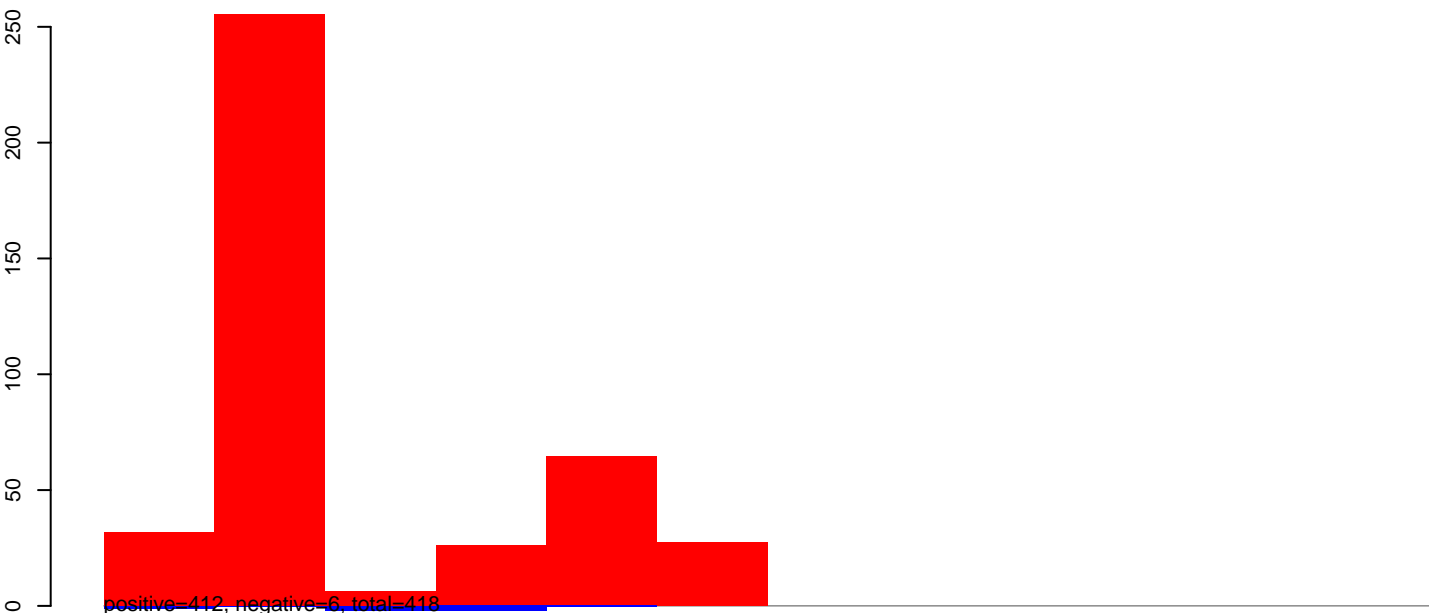
AeAeg_Testes.24_35.rep



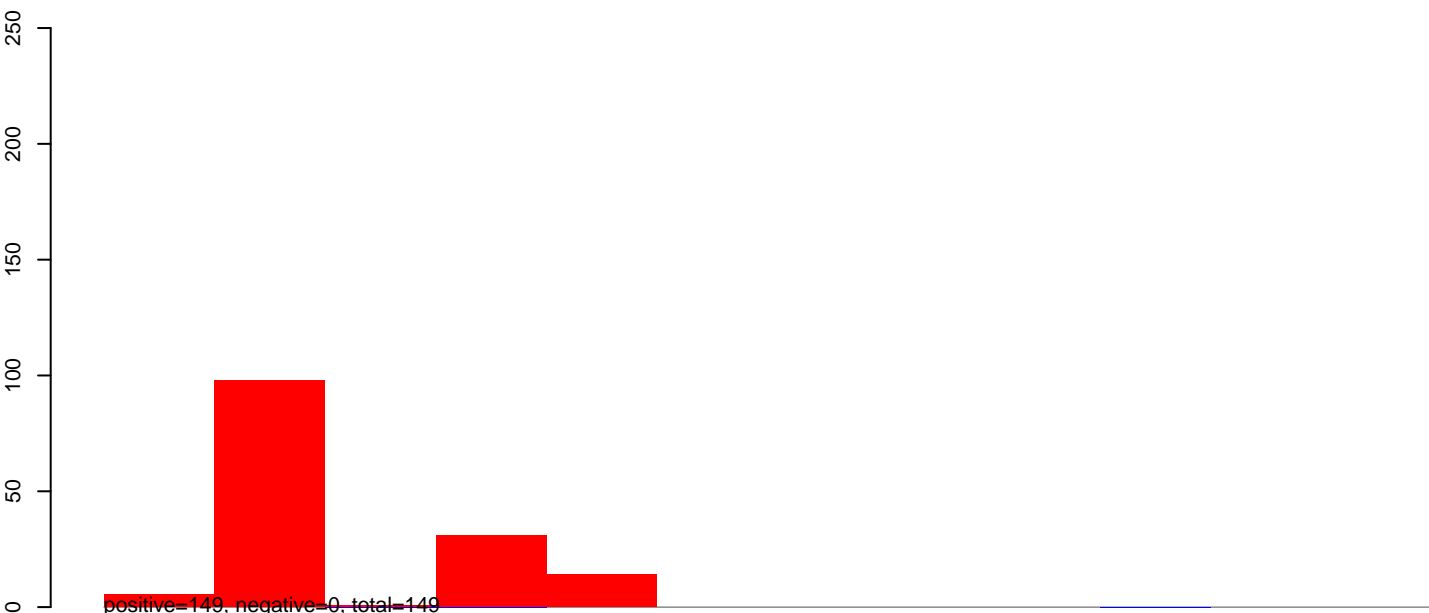
AeAeg_Testes.rep



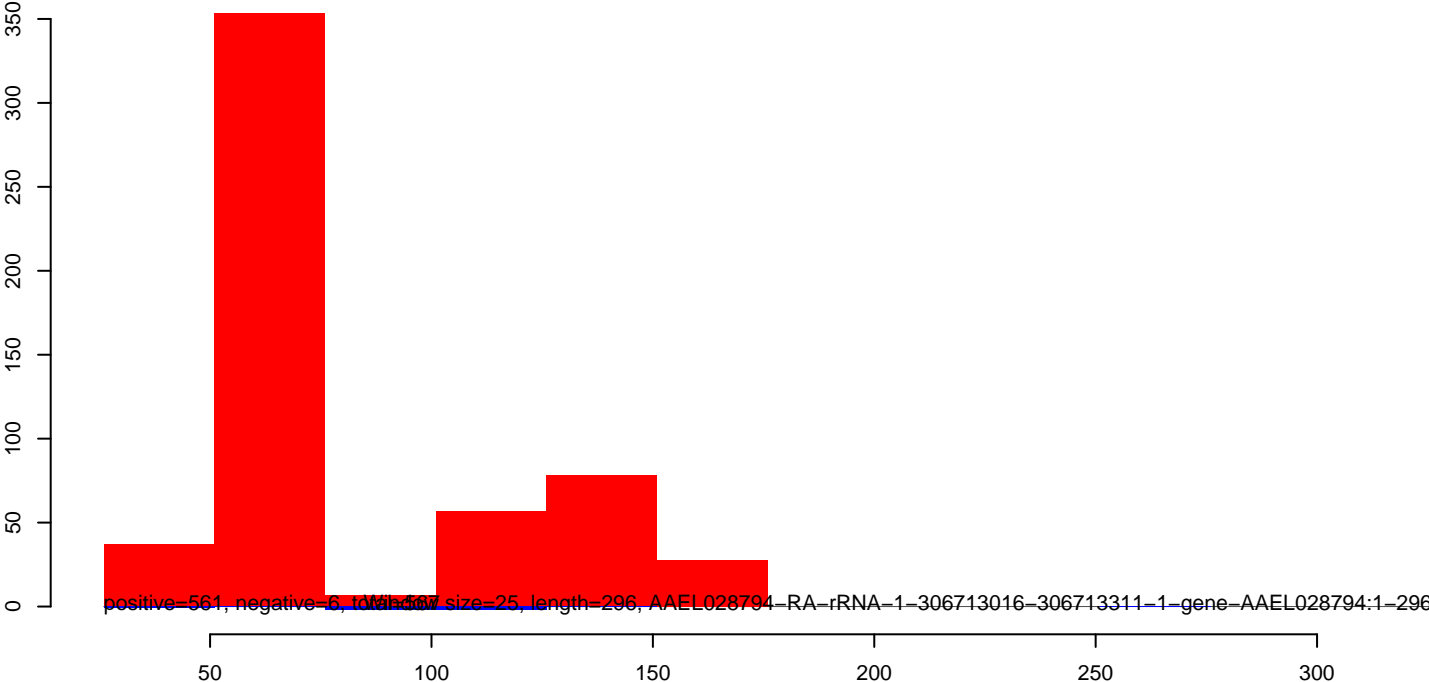
AeAeg_Testes.18_23.rep



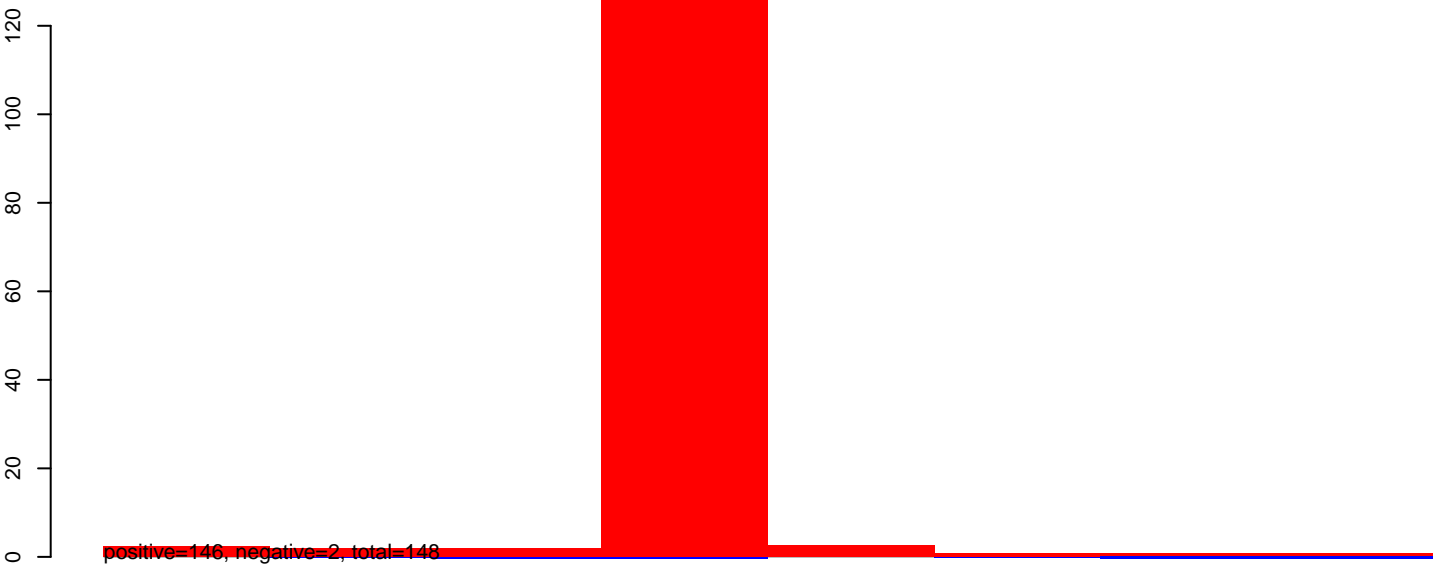
AeAeg_Testes.24_35.rep



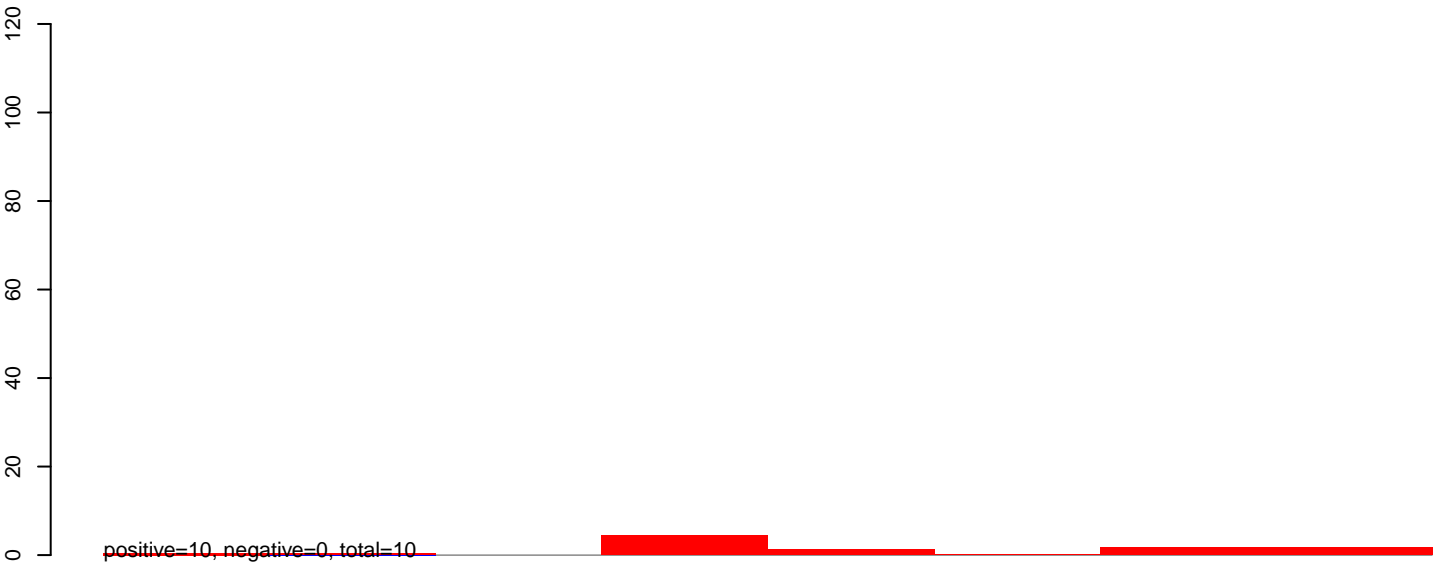
AeAeg_Testes.rep



AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



Windows: 50, negative=0, total=10, AAEL017825-RA-U2-U2-spliceosomal-RNA-snRNA-2-123654334-123654528-1-gene-AAEL017825-1-195

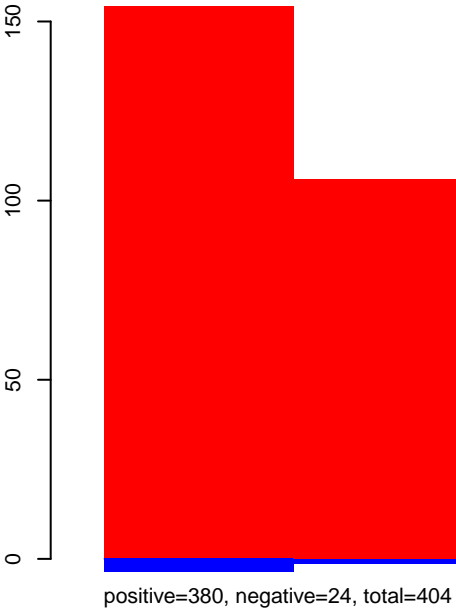
50

100

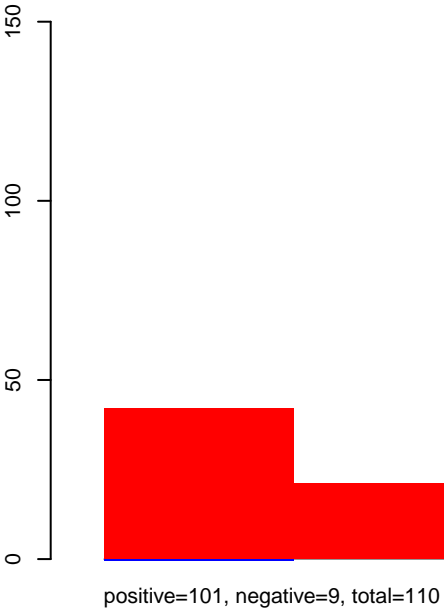
150

200

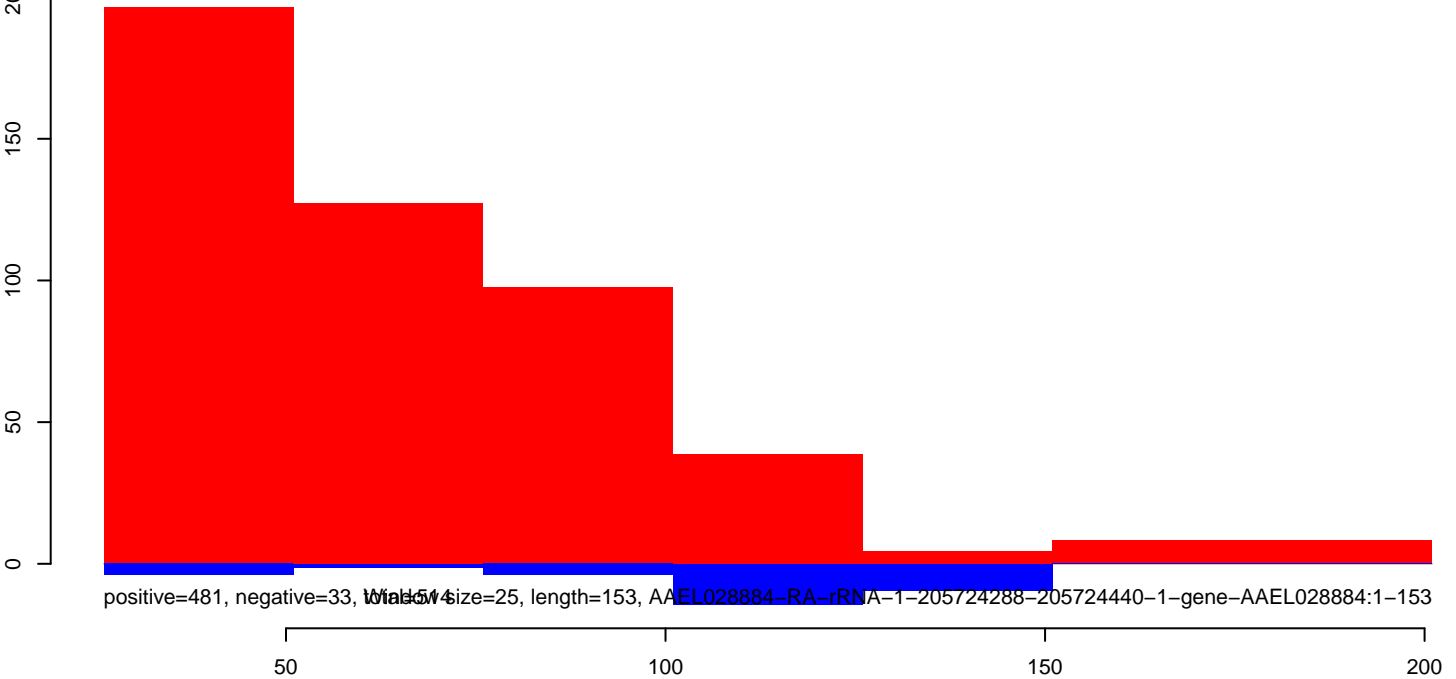
AeAeg_Testes.18_23.rep



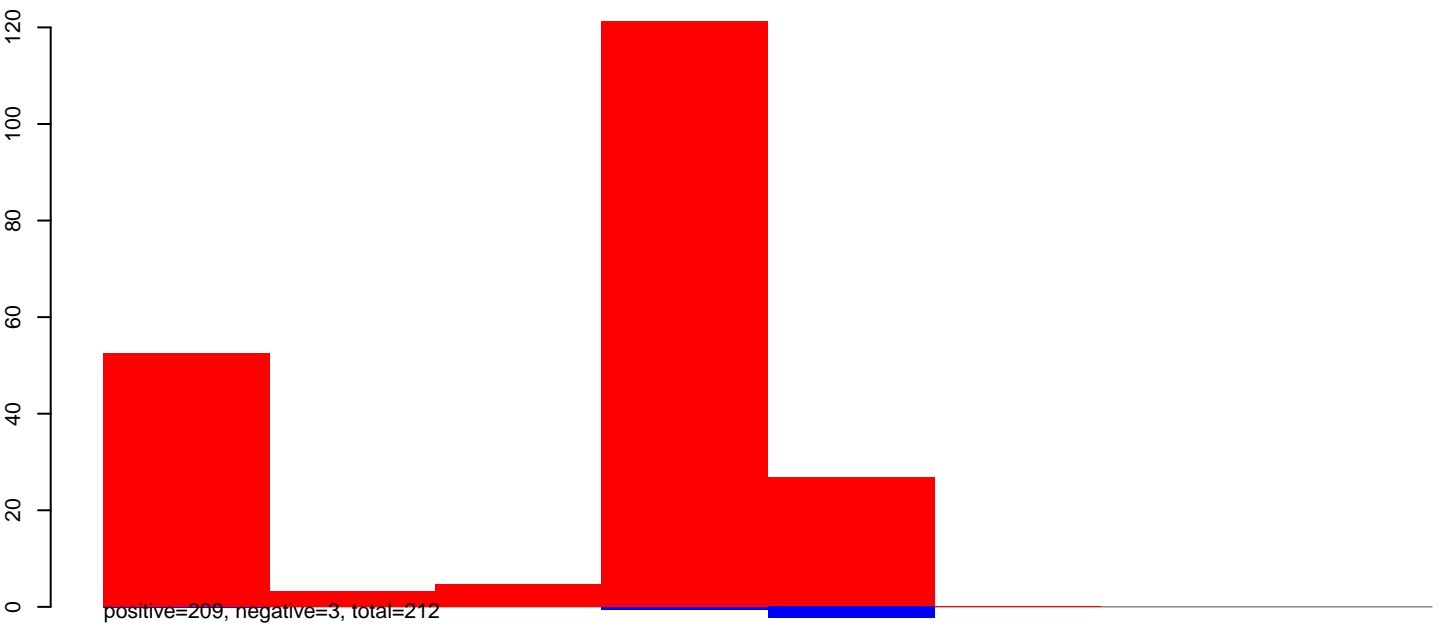
AeAeg_Testes.24_35.rep



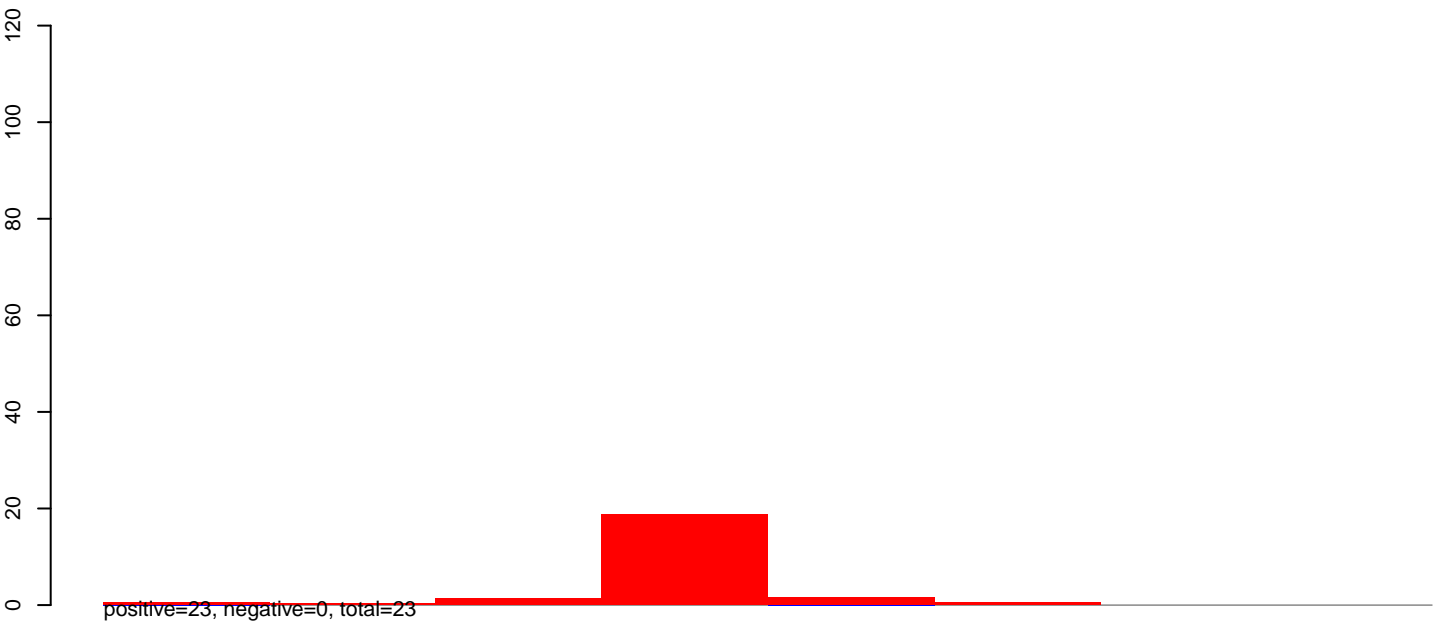
AeAeg_Testes.rep



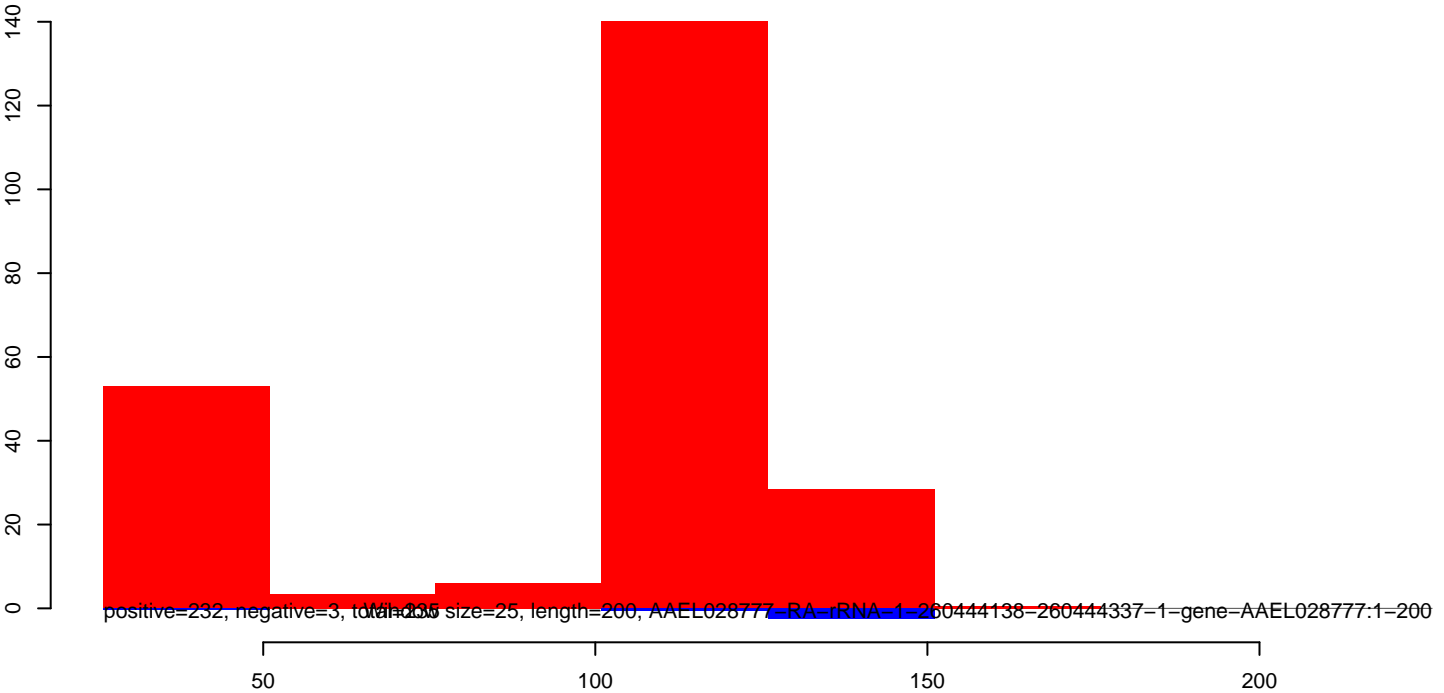
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



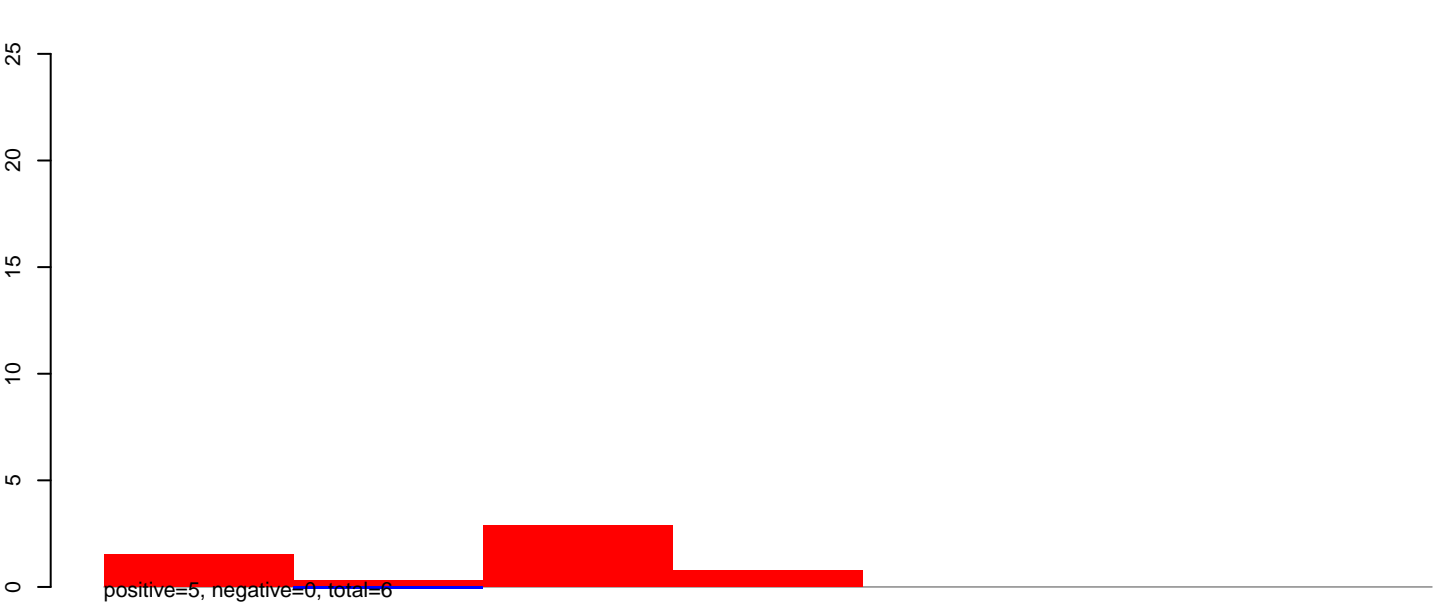
AeAeg_Testes.rep



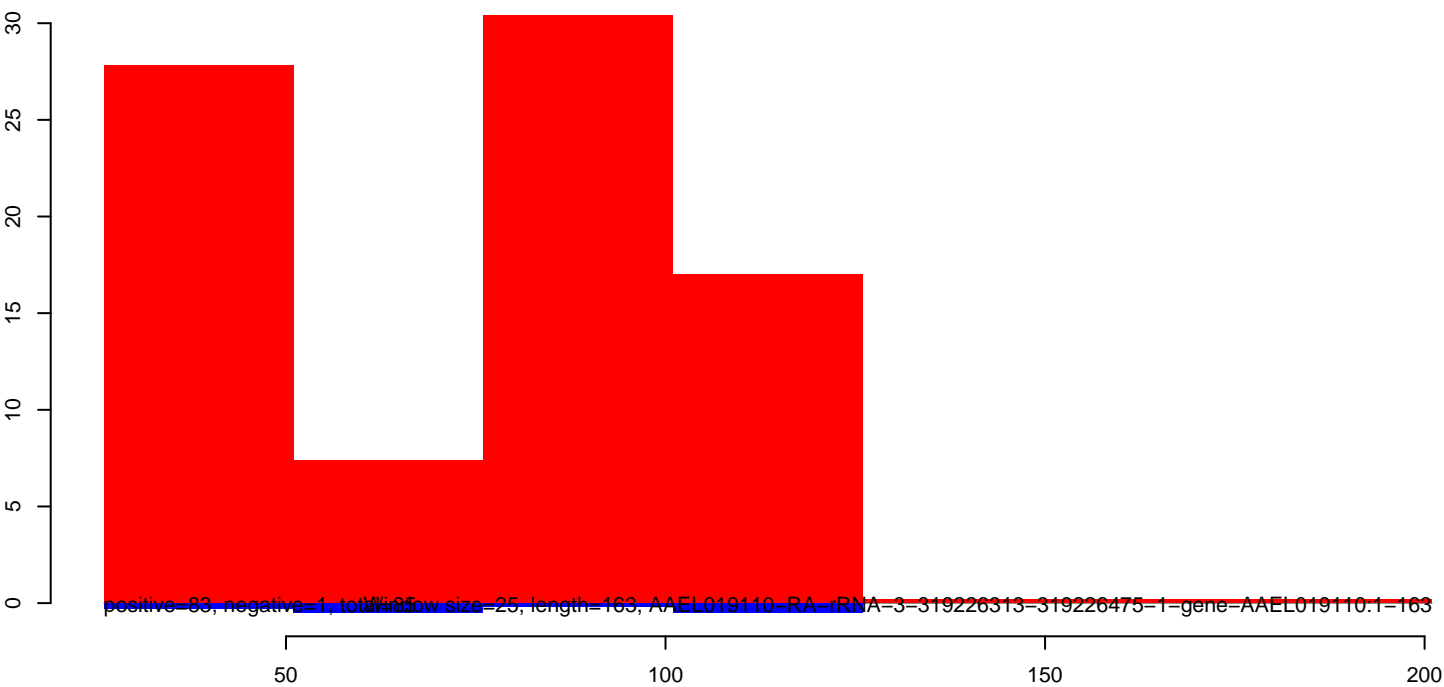
AeAeg_Testes.18_23.rep



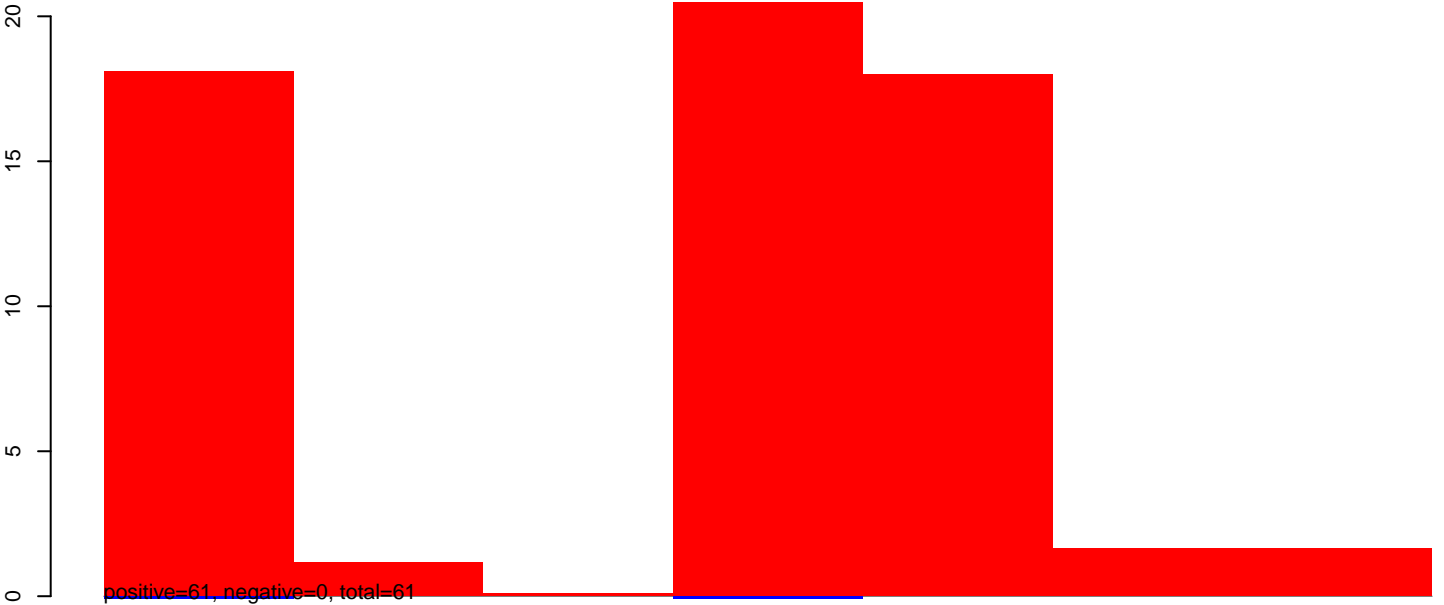
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



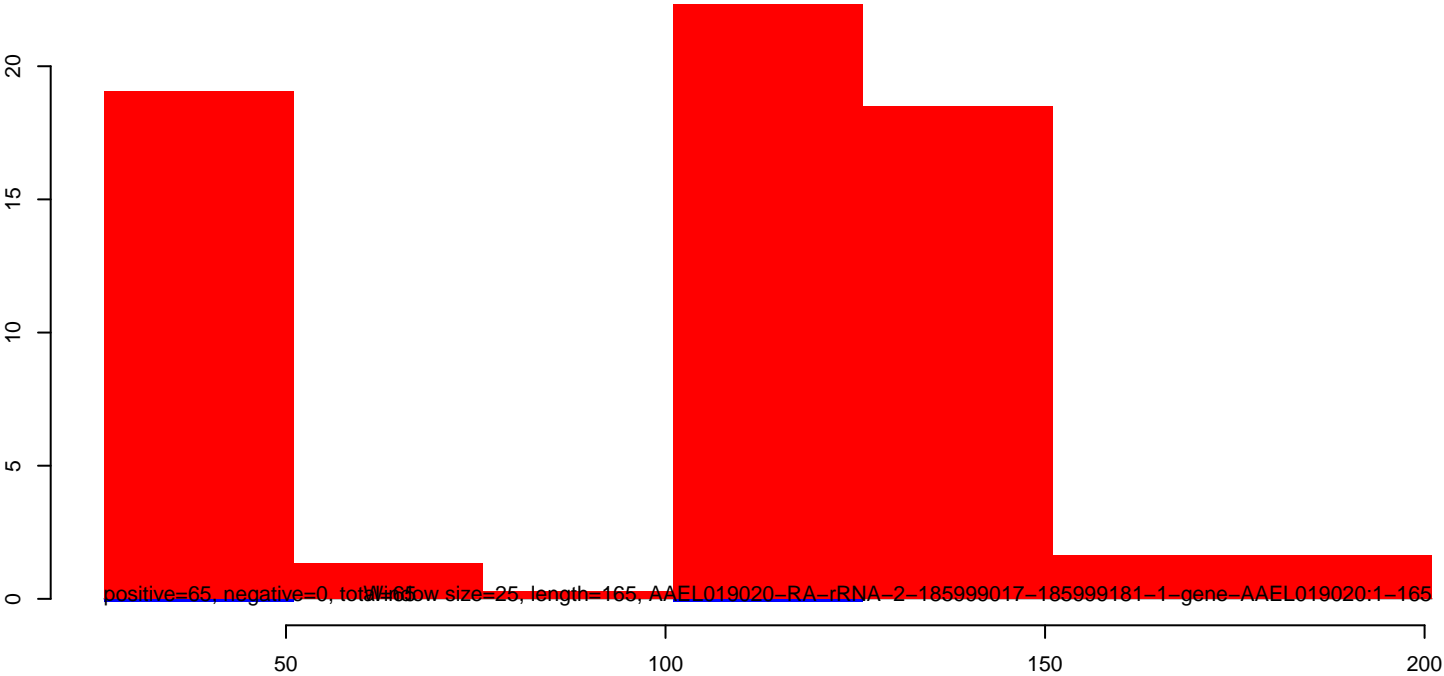
AeAeg_Testes.18_23.rep



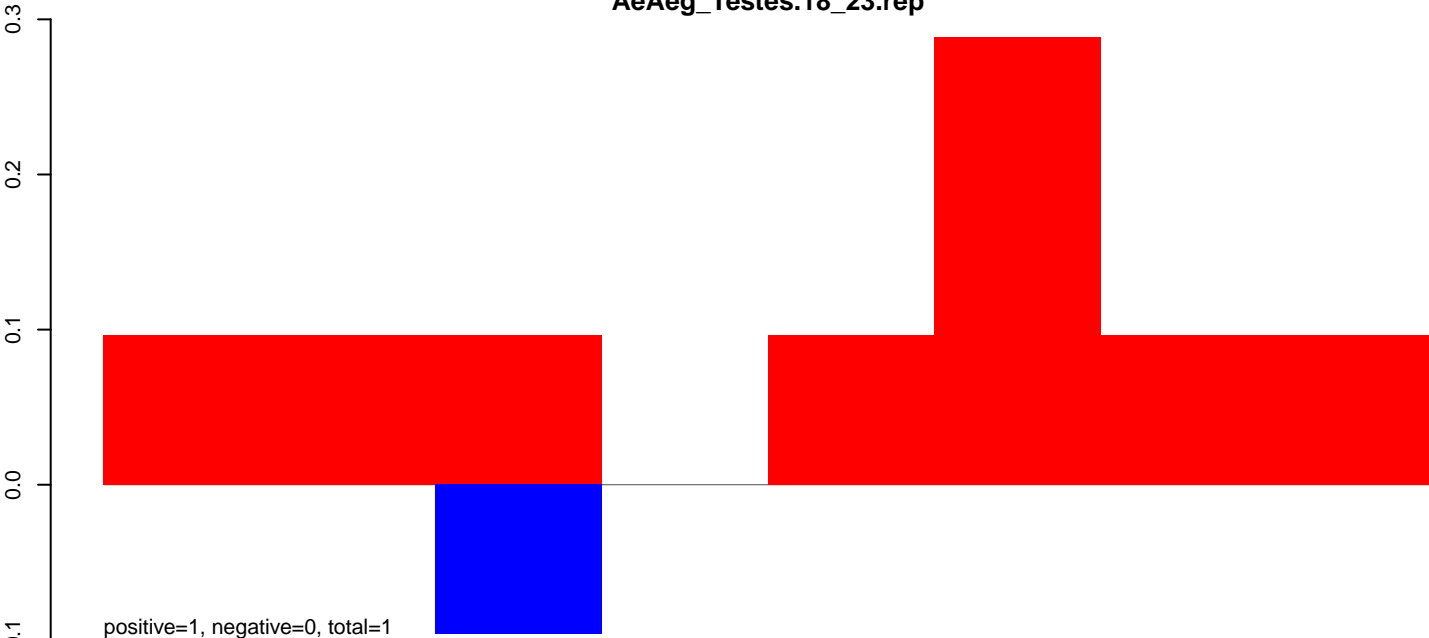
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



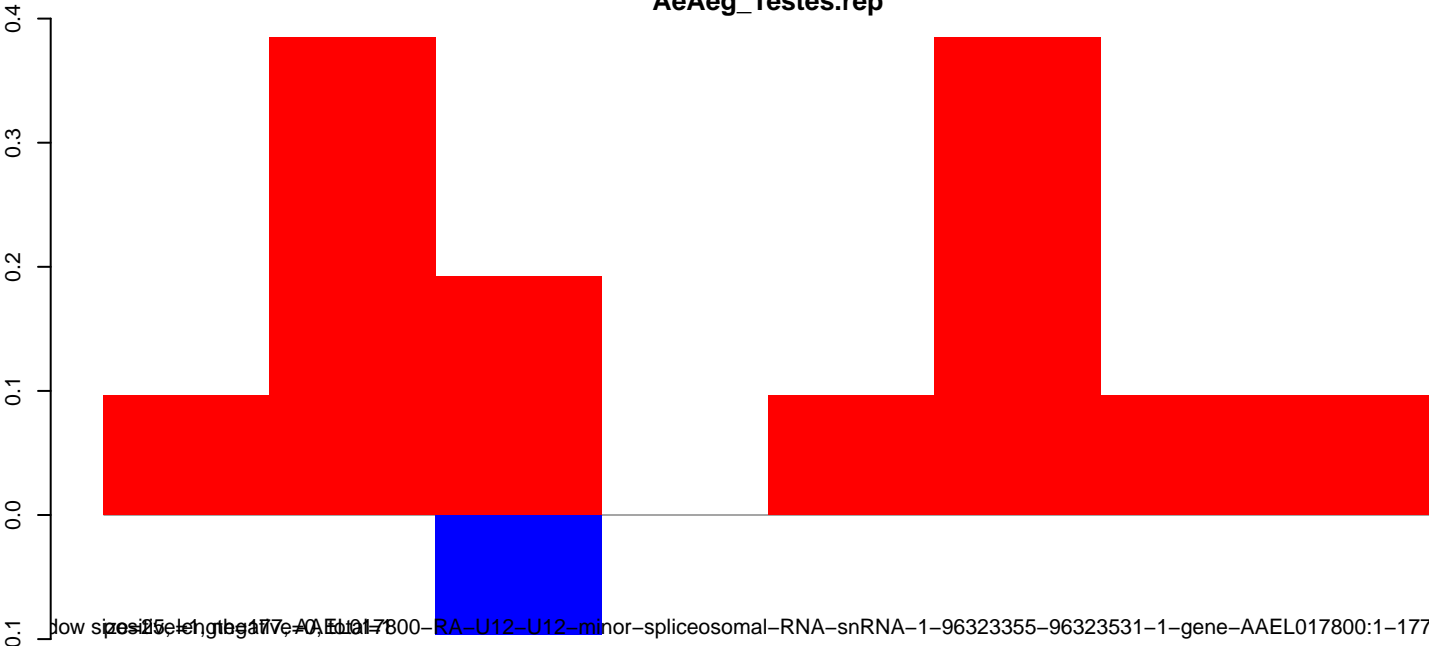
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



low size change 170, AAEL017800-RA-U12-U12-minor-spliceosomal-RNA-snrRNA-1-96323355-96323531-1-gene-AAEL017800:1-177

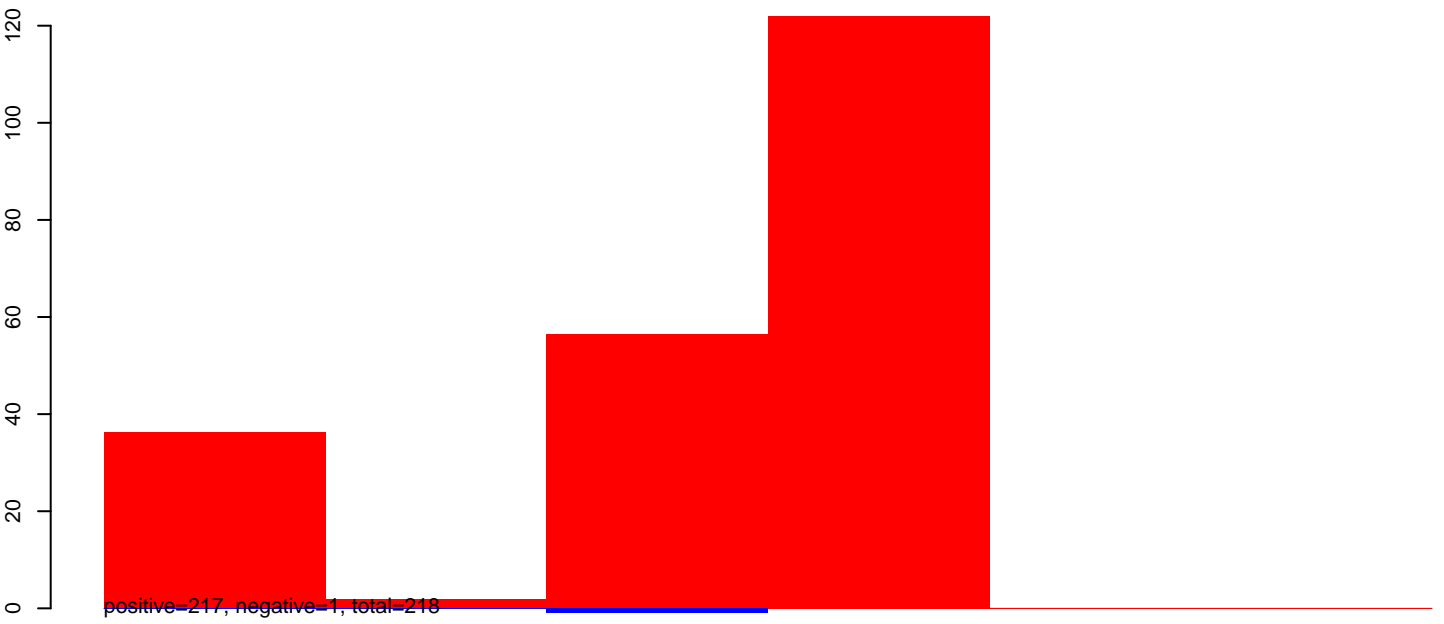
50

100

150

200

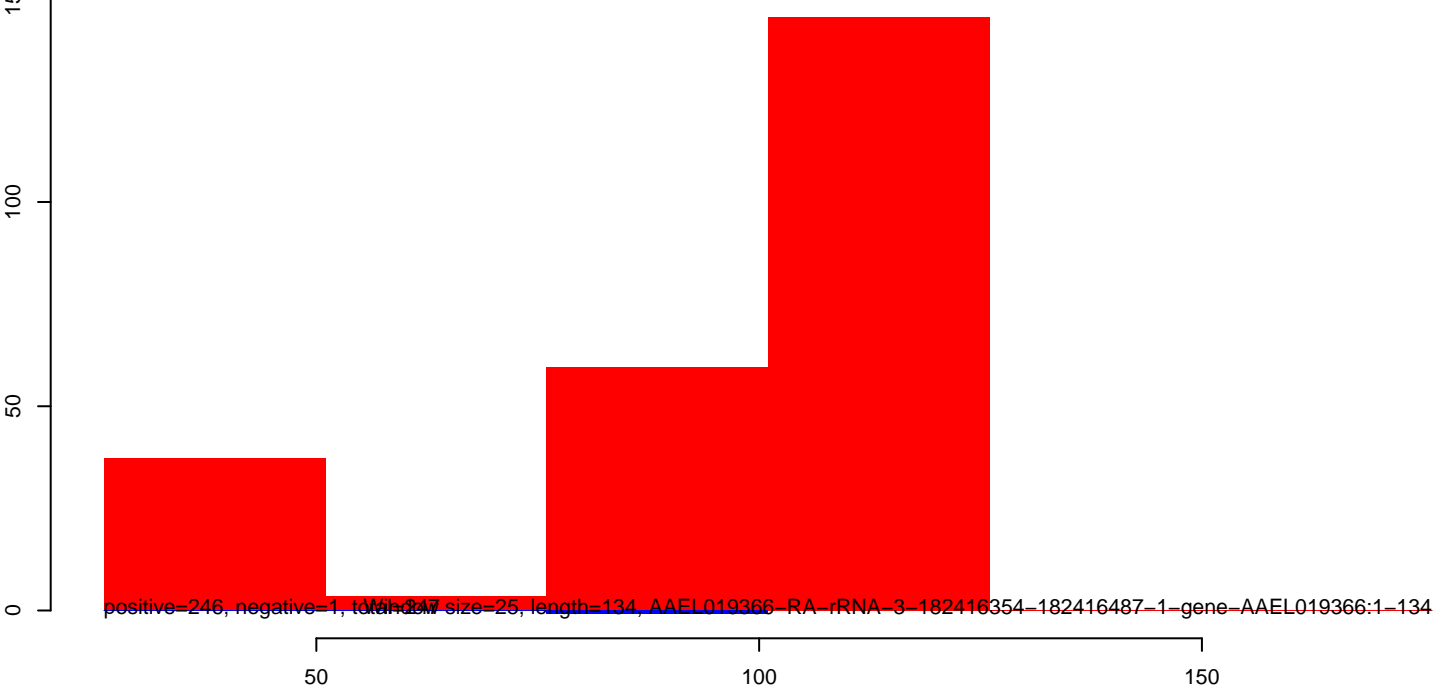
AeAeg_Testes.18_23.rep



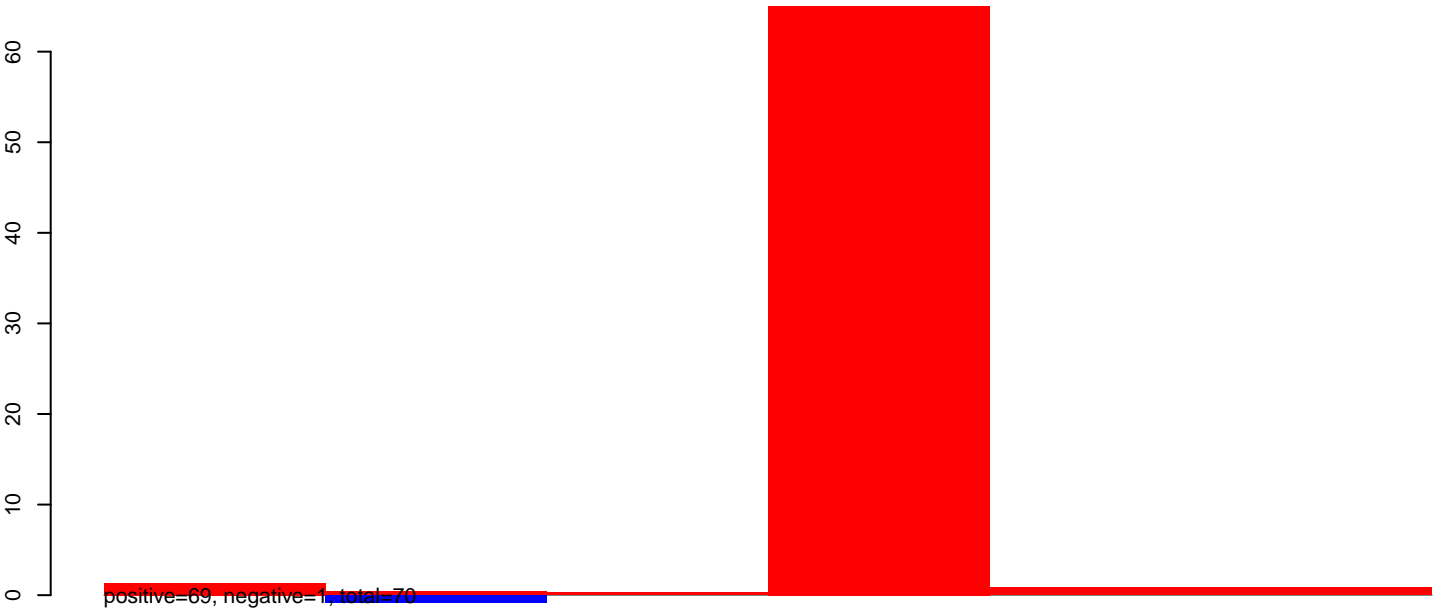
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



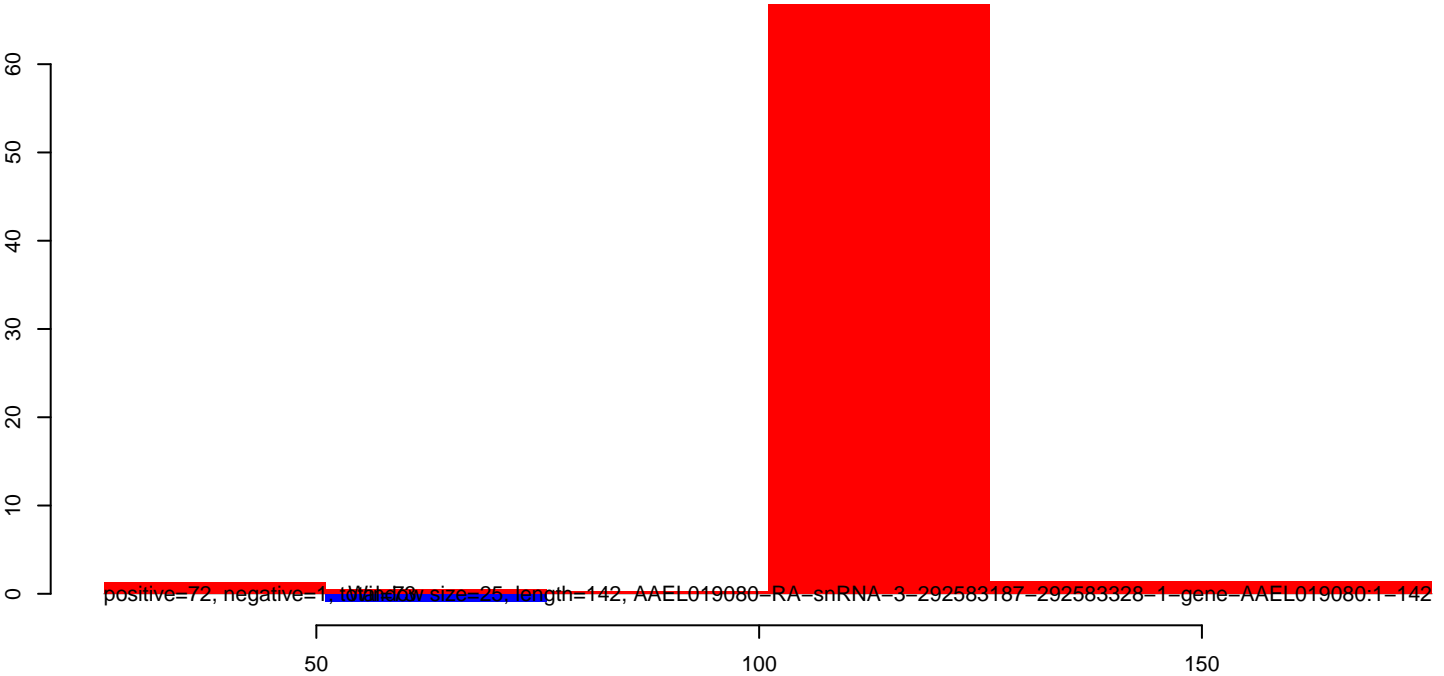
AeAeg_Testes.18_23.rep



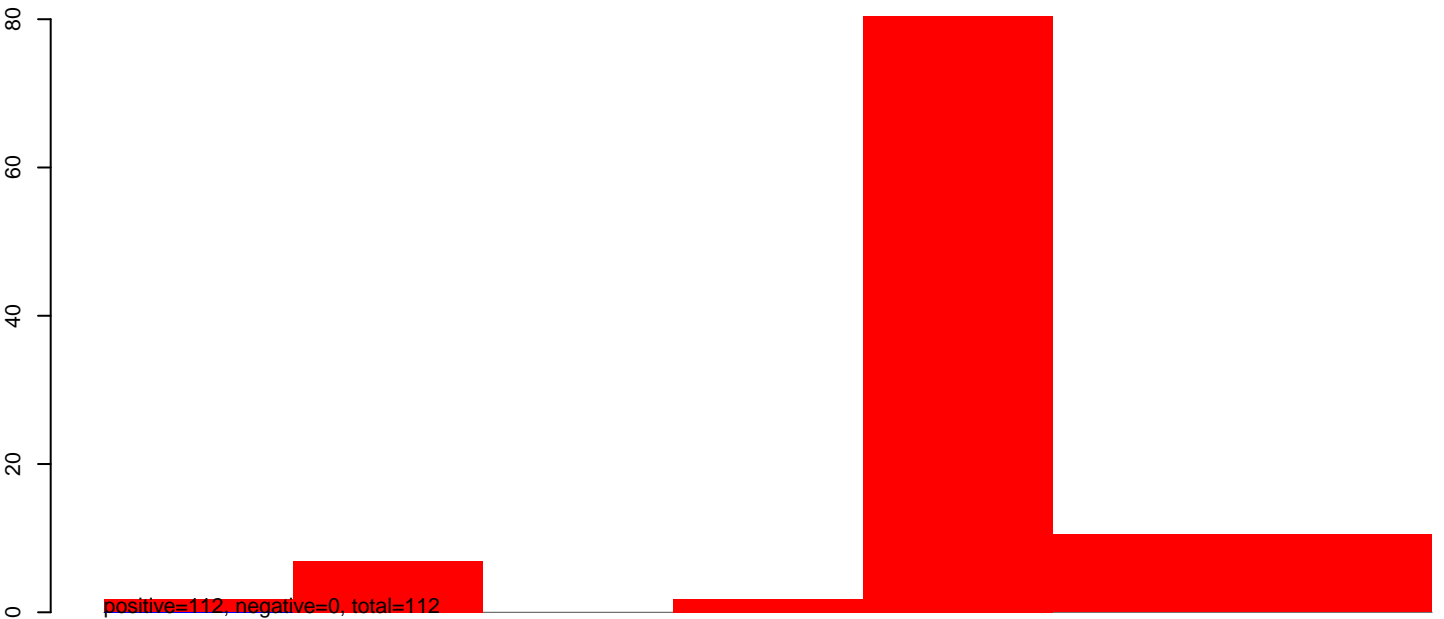
AeAeg_Testes.24_35.rep



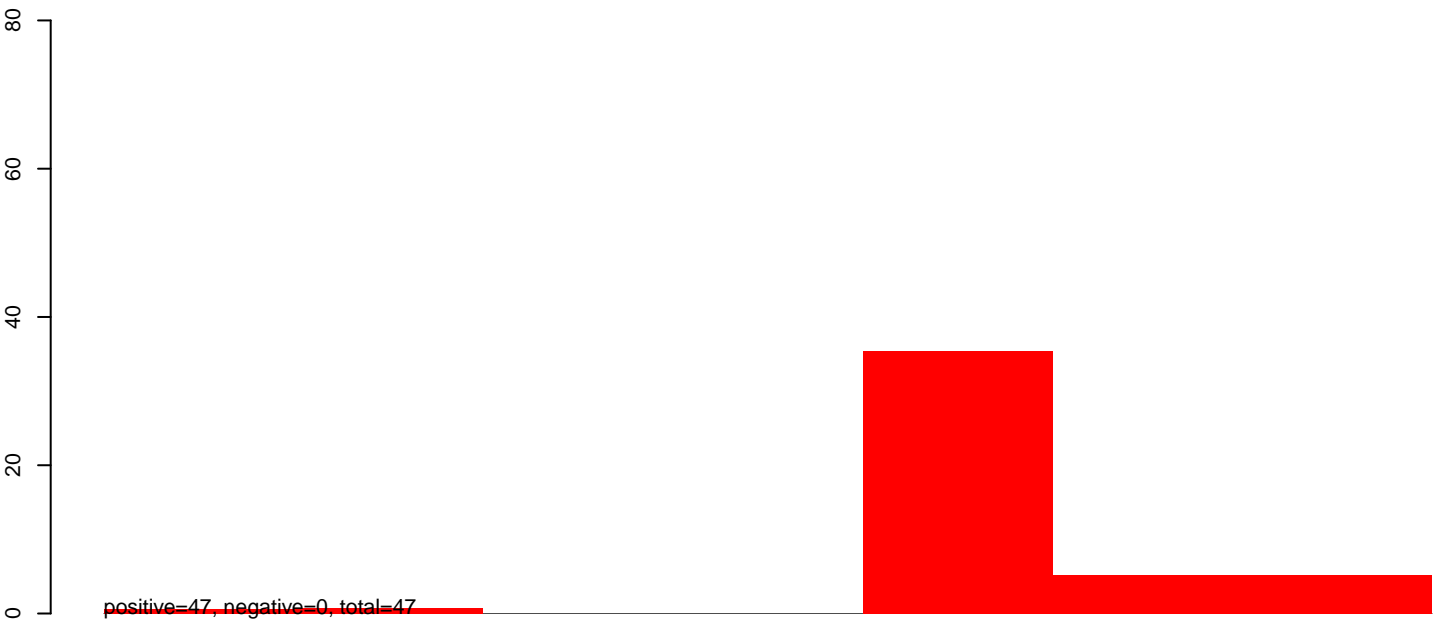
AeAeg_Testes.rep



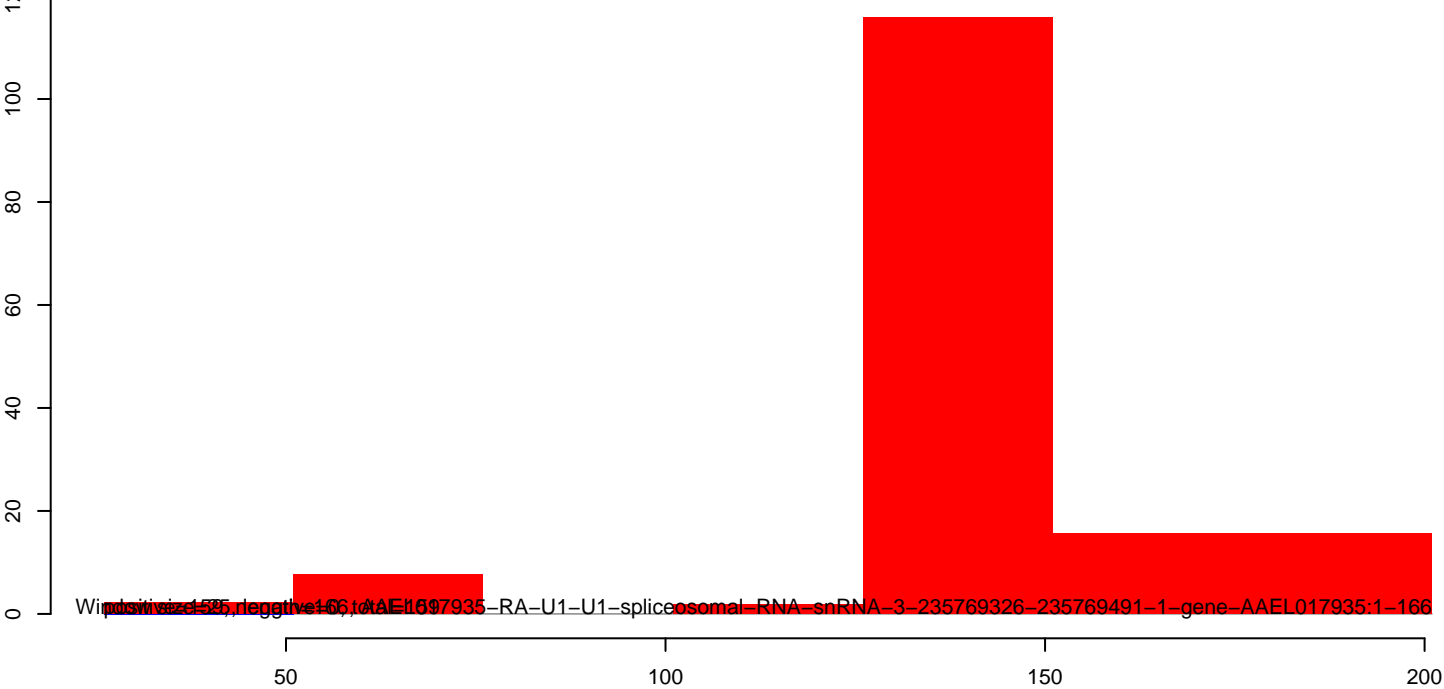
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



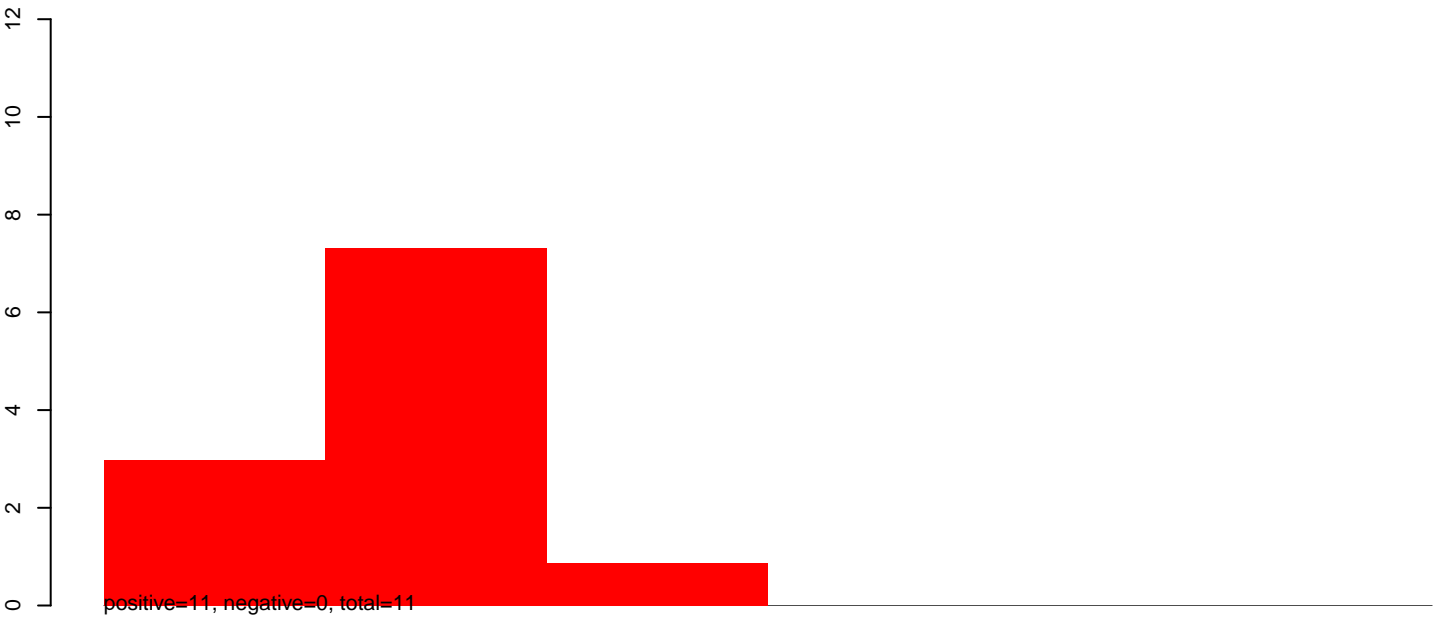
AeAeg_Testes.rep



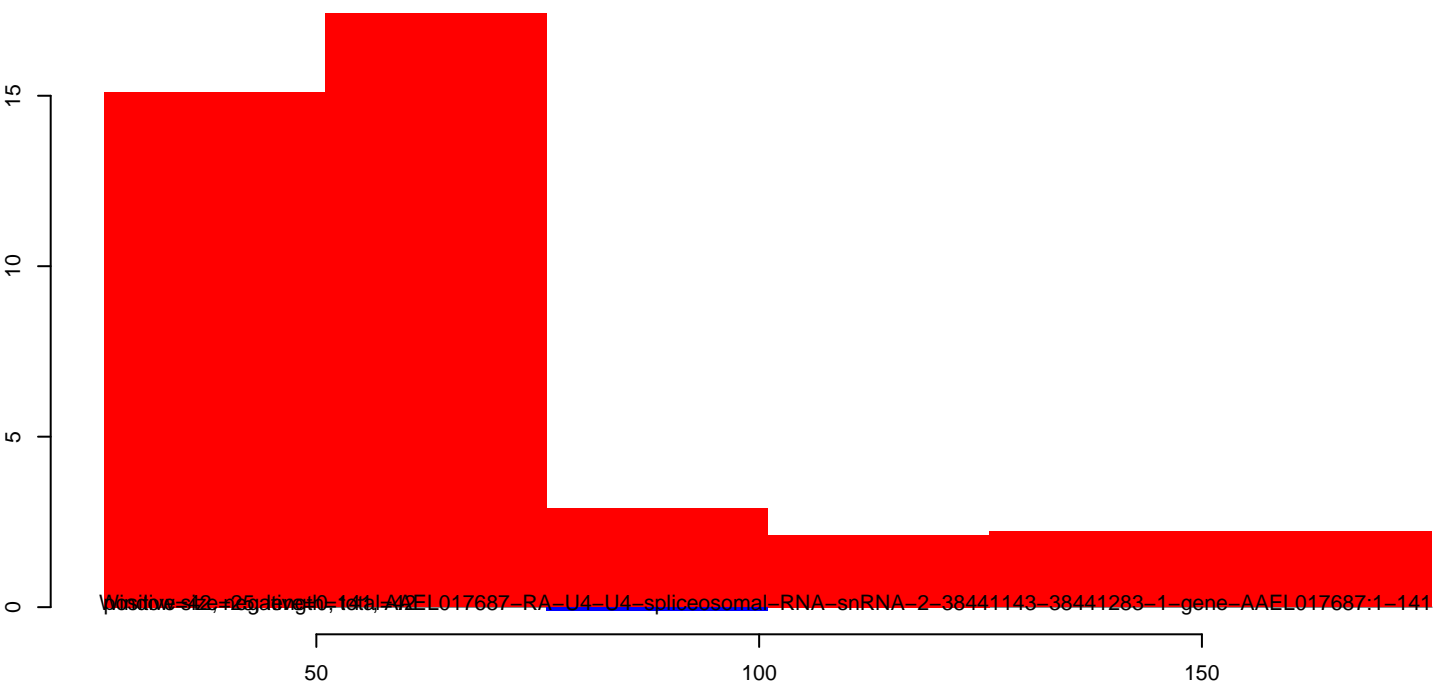
AeAeg_Testes.18_23.rep

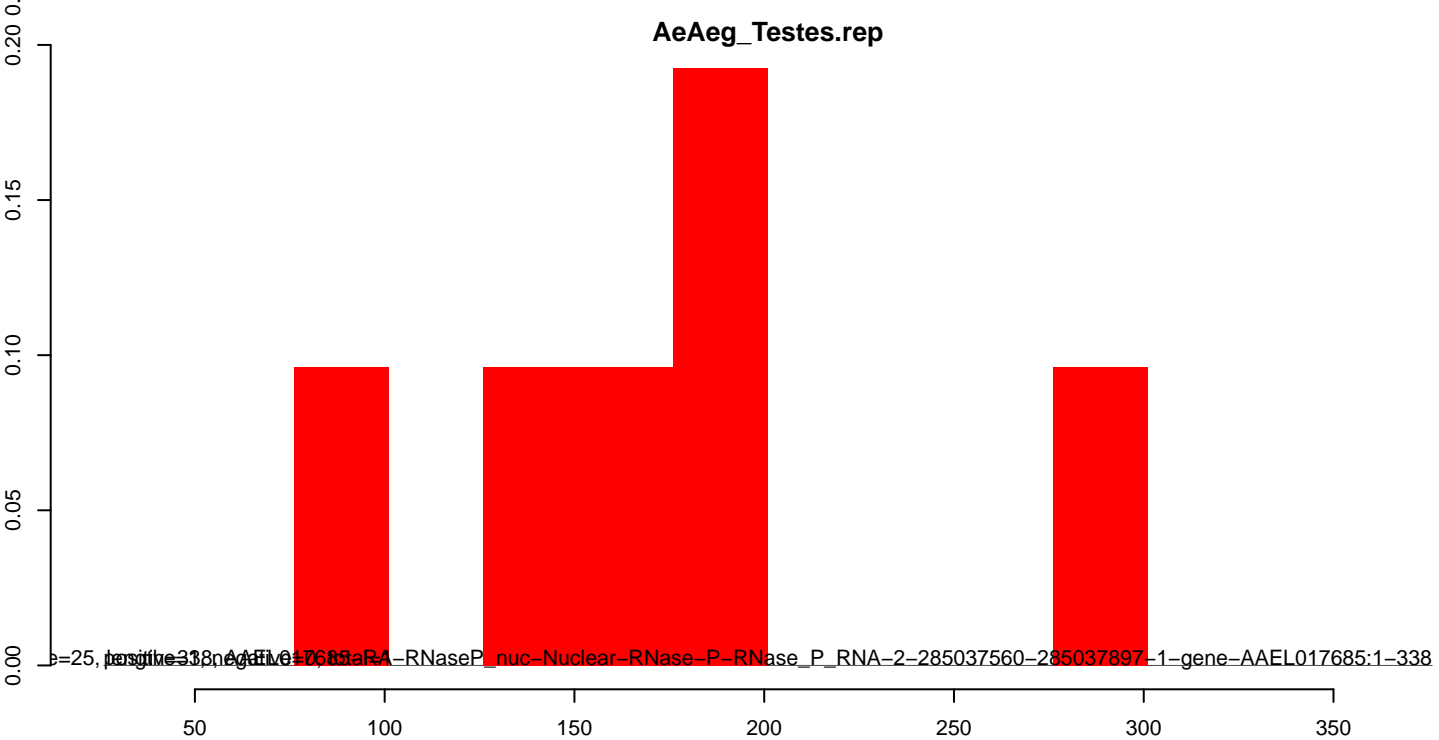
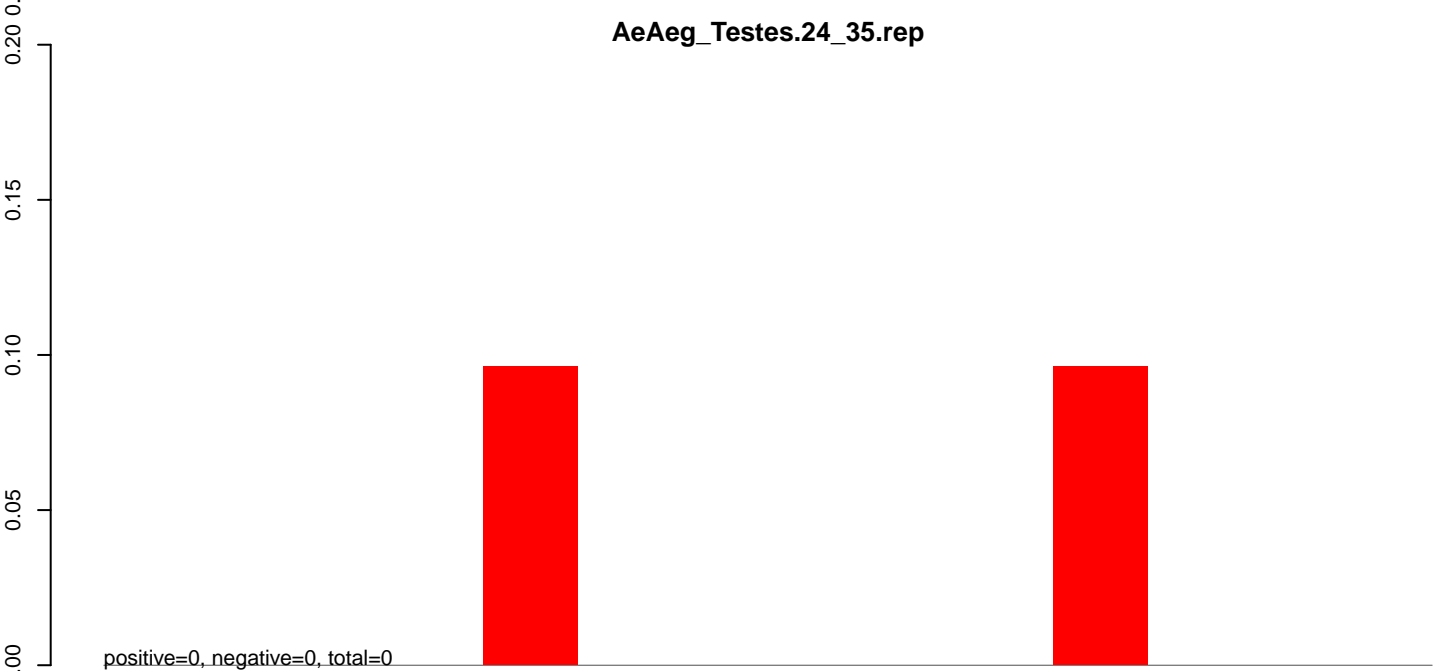
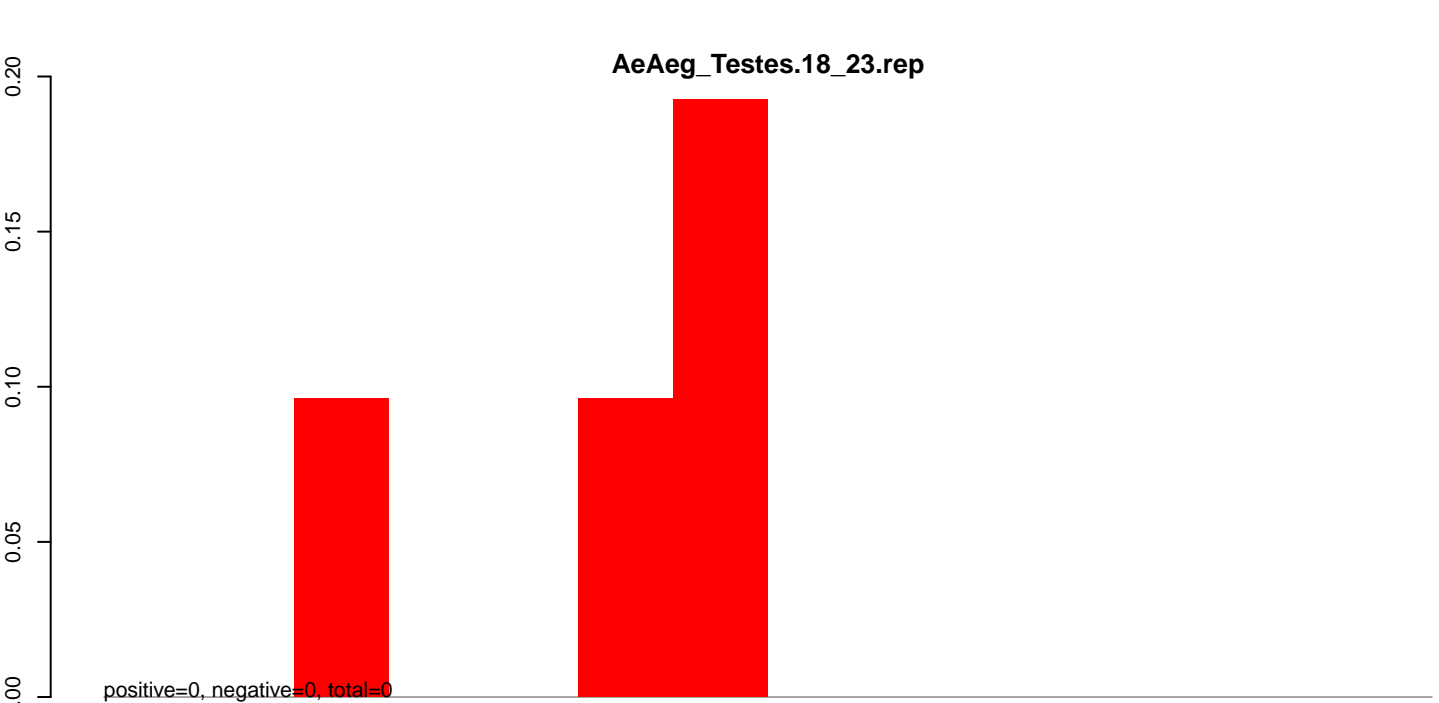


AeAeg_Testes.24_35.rep

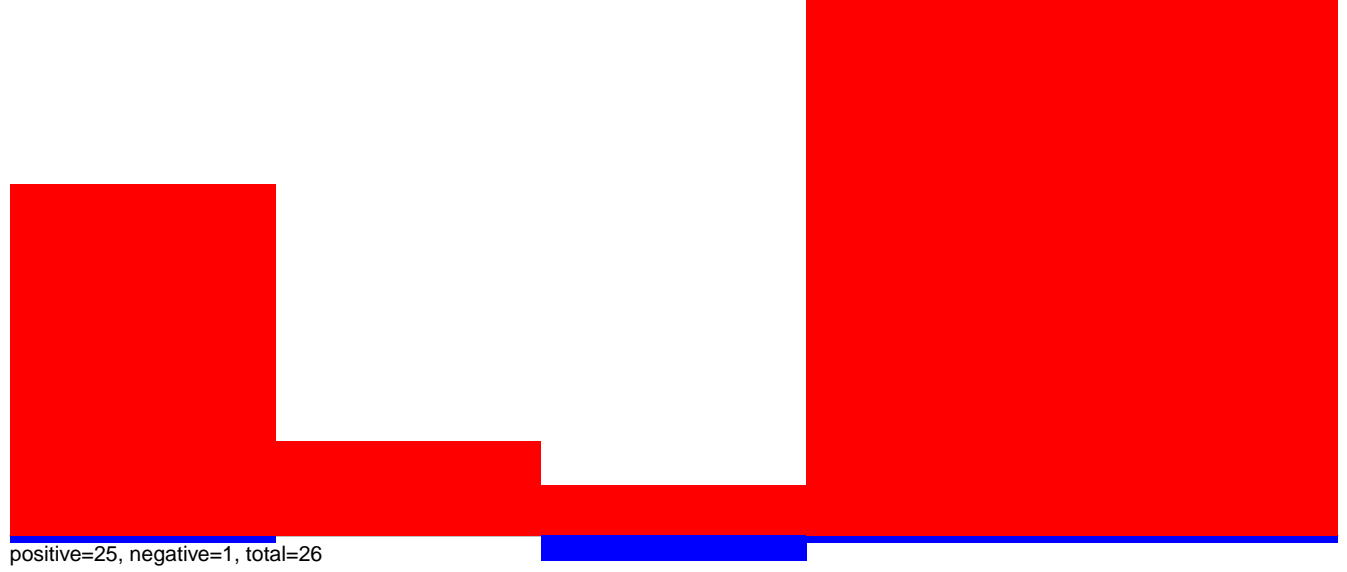


AeAeg_Testes.rep

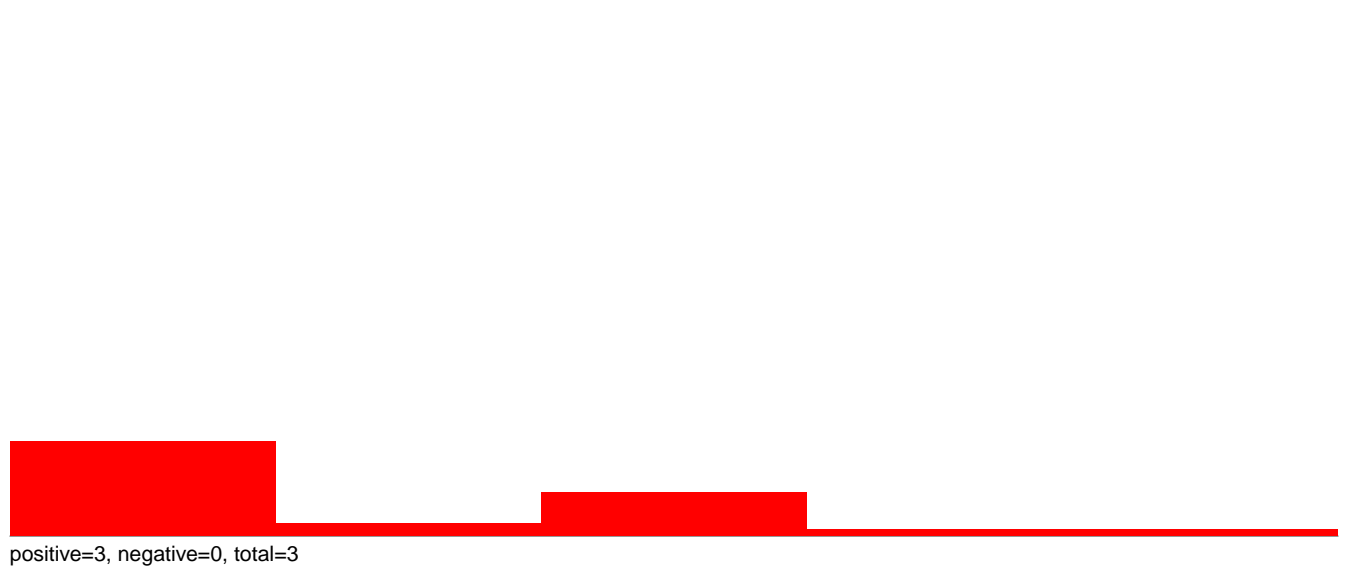




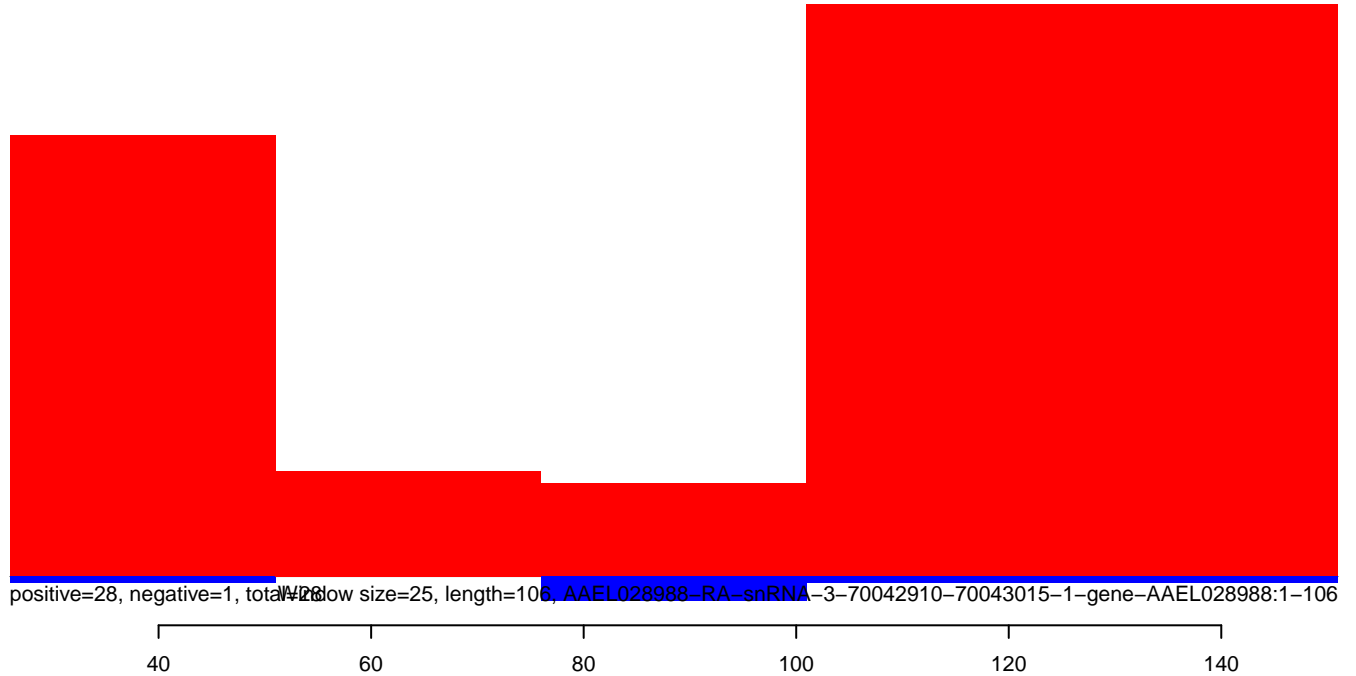
AeAeg_Testes.18_23.rep

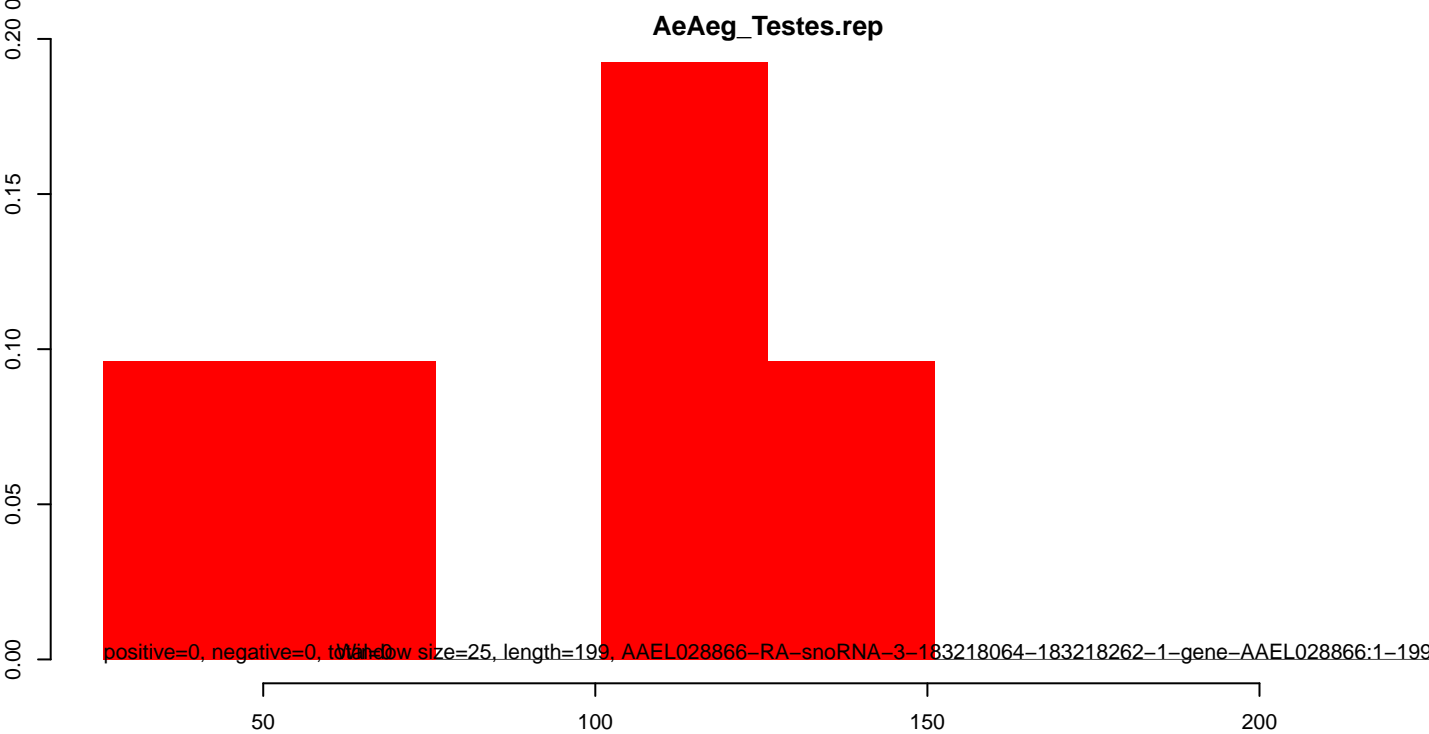
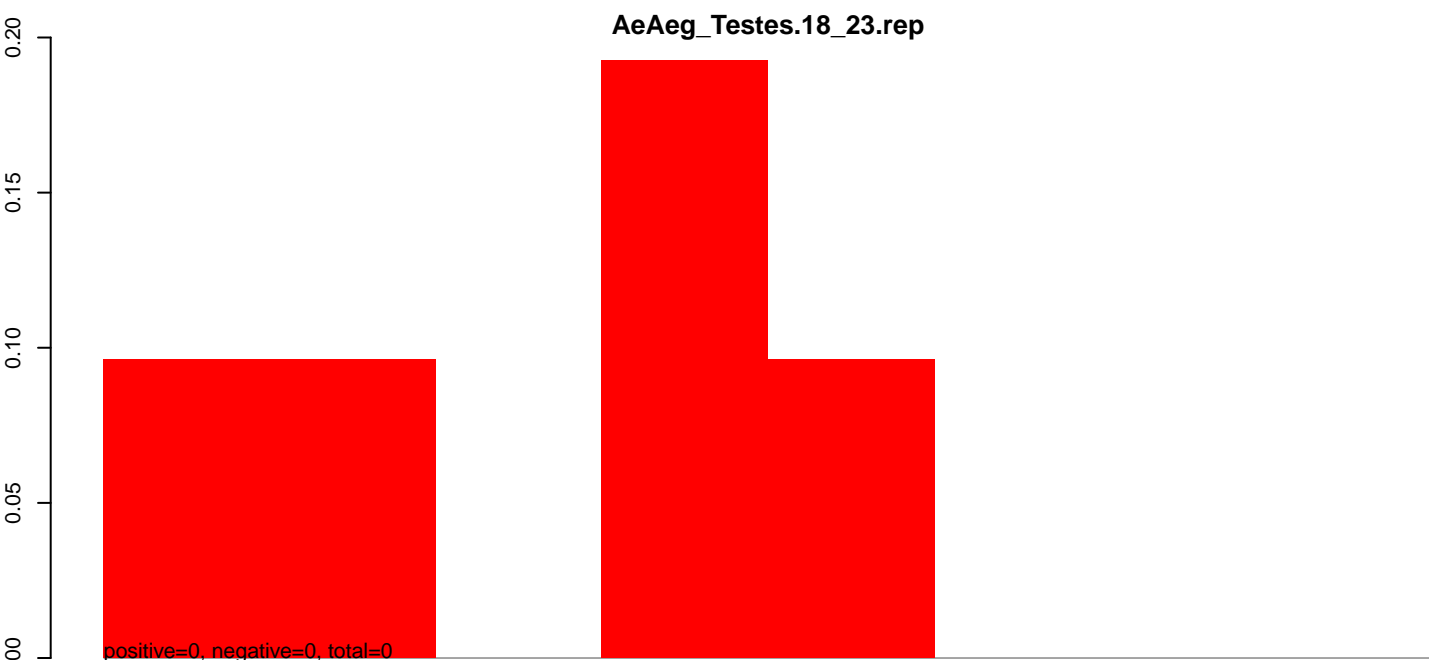


AeAeg_Testes.24_35.rep

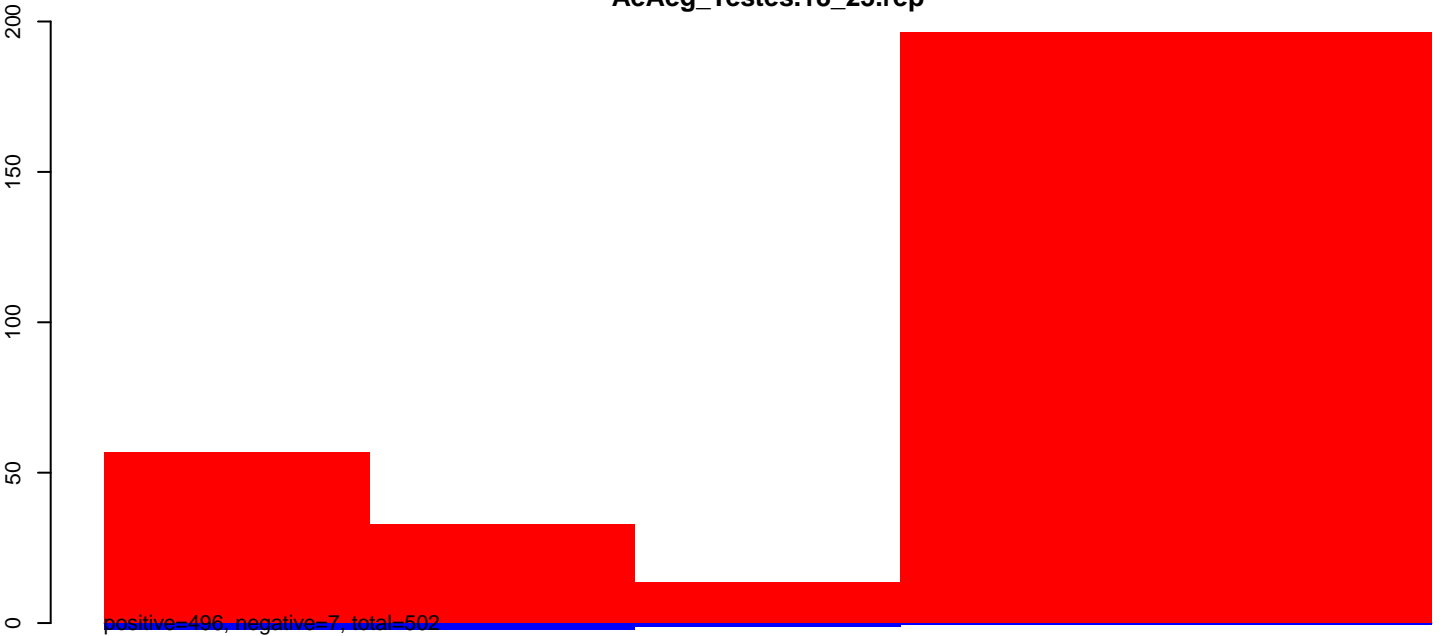


AeAeg_Testes.rep

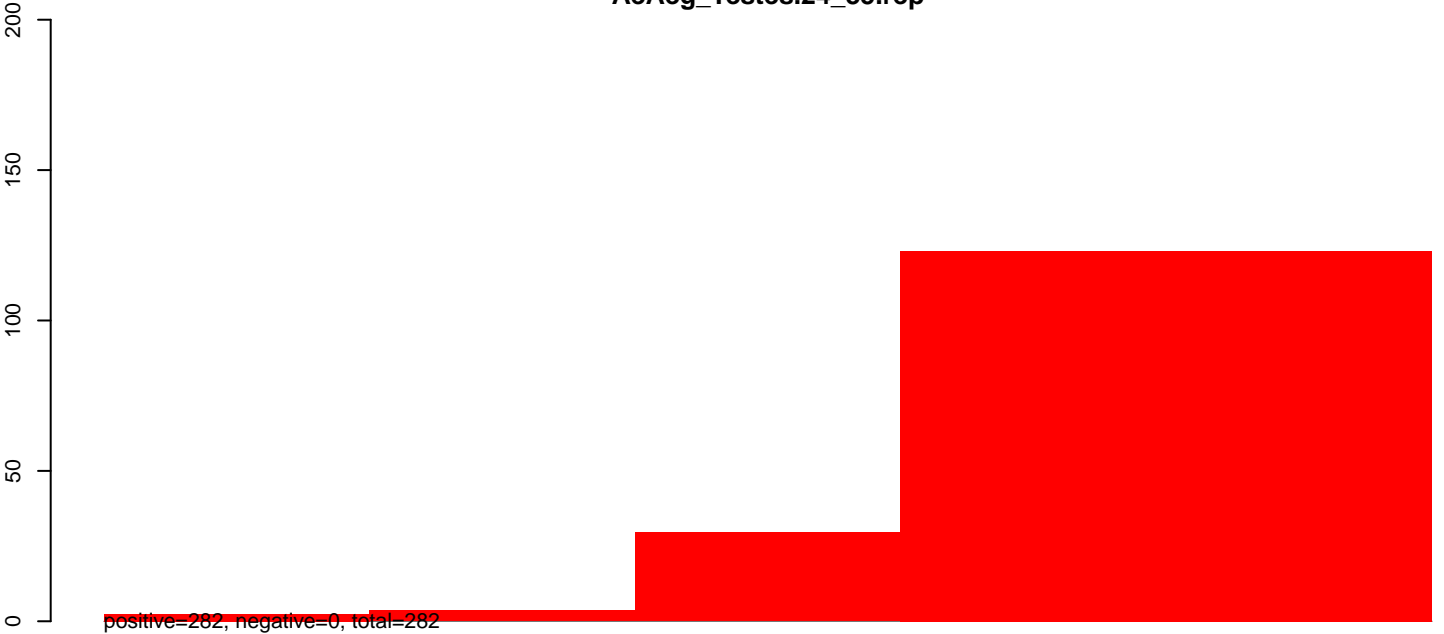




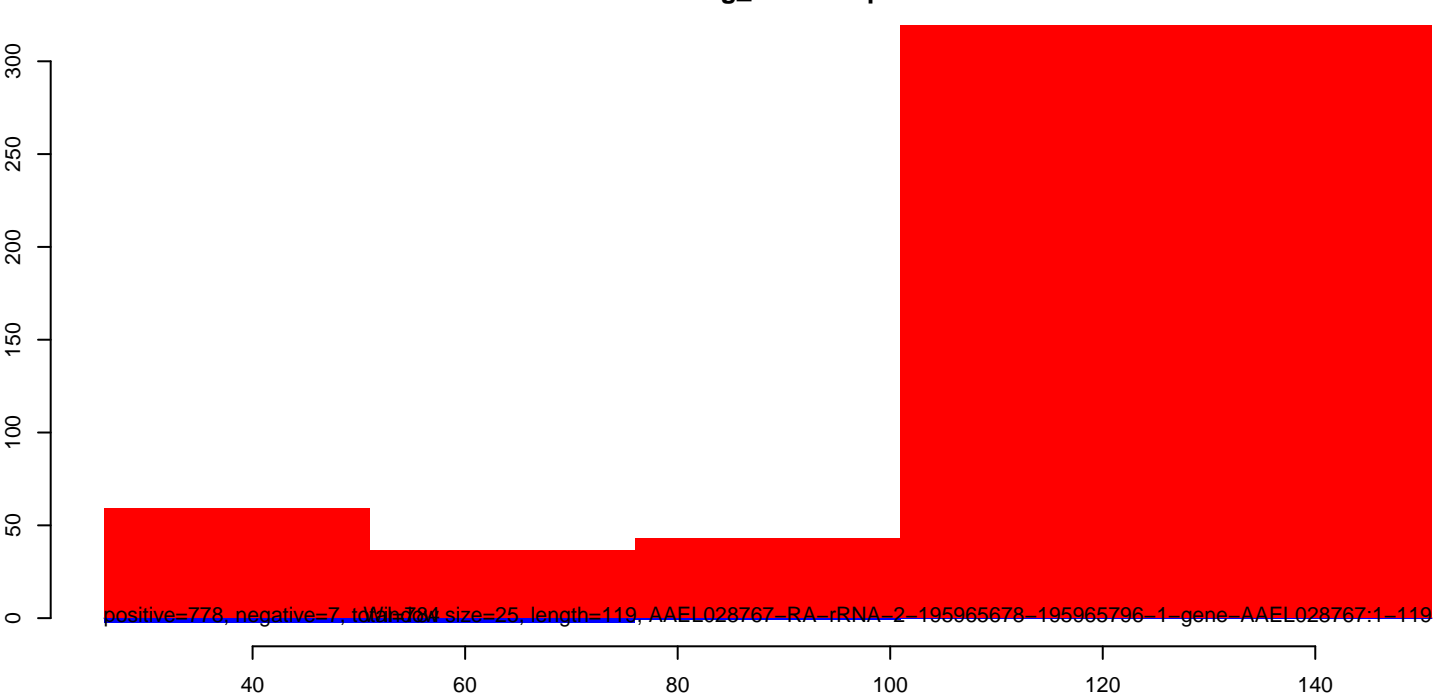
AeAeg_Testes.18_23.rep



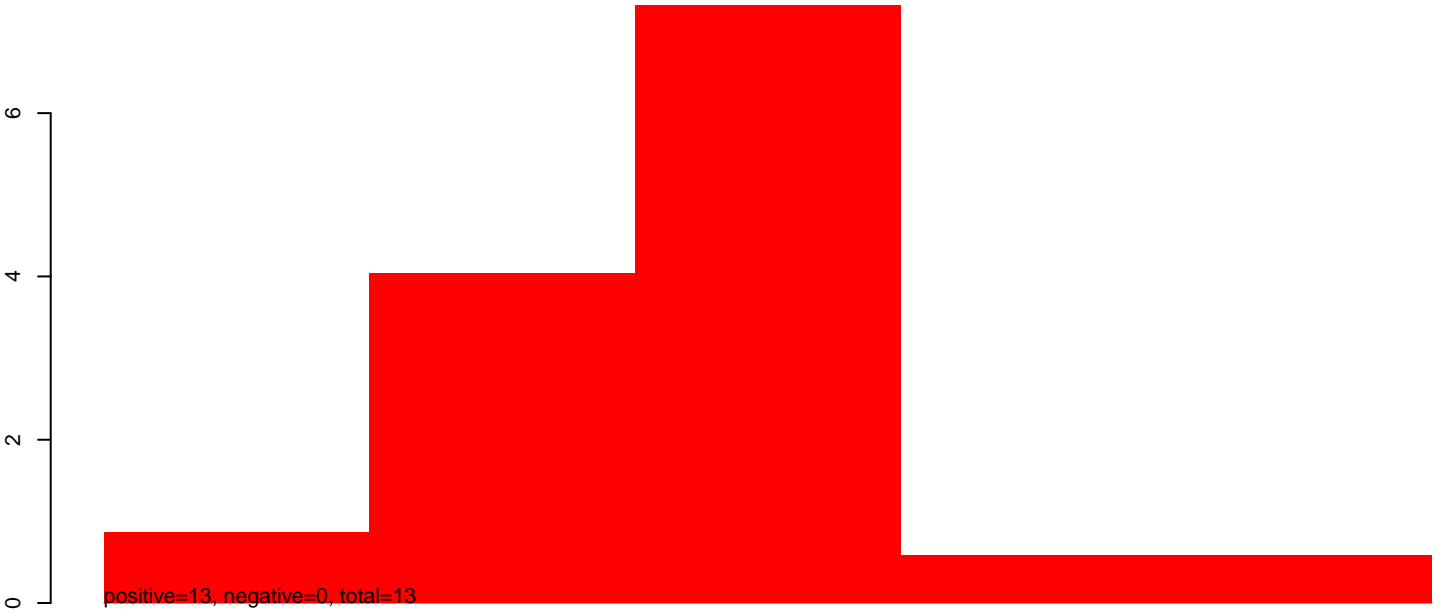
AeAeg_Testes.24_35.rep



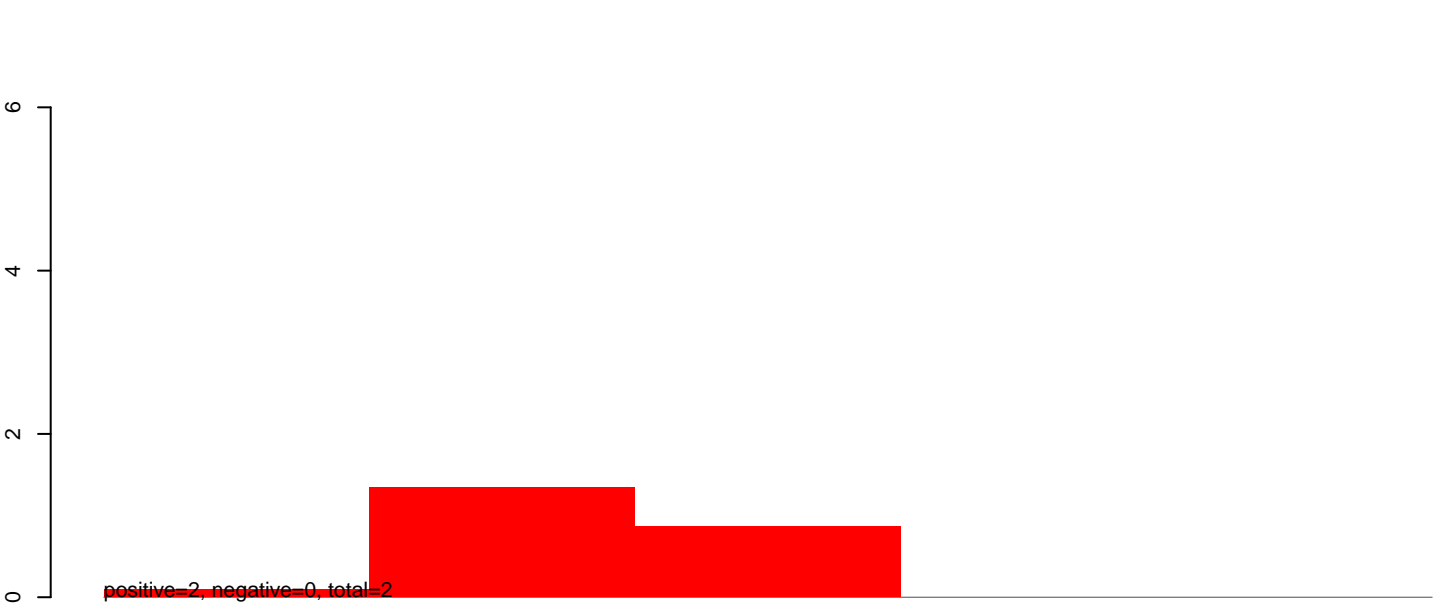
AeAeg_Testes.rep



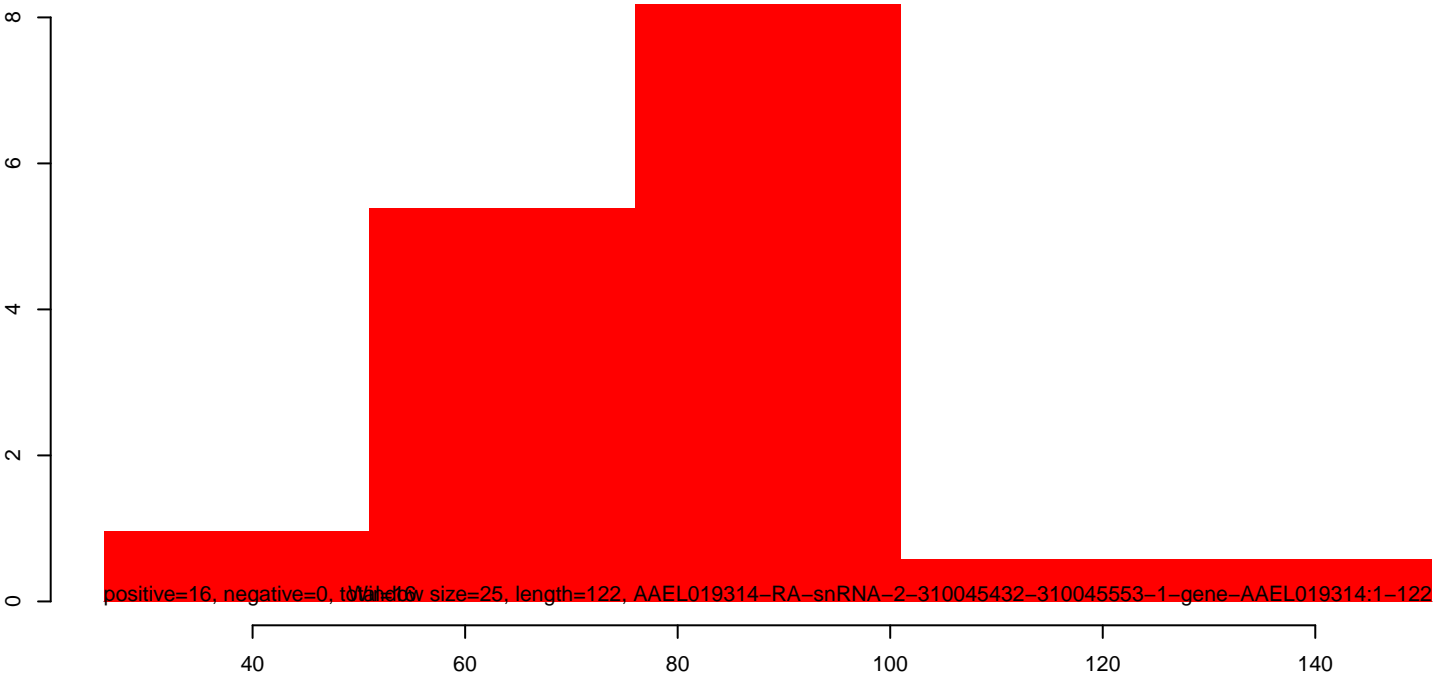
AeAeg_Testes.18_23.rep



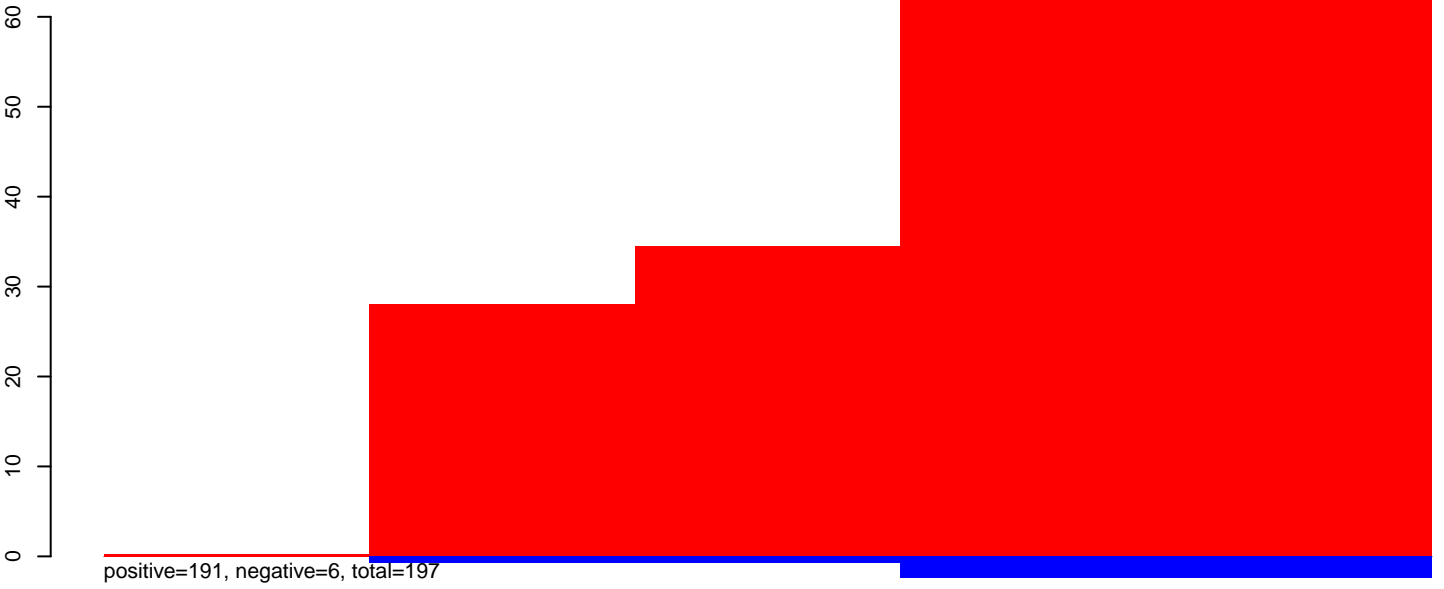
AeAeg_Testes.24_35.rep



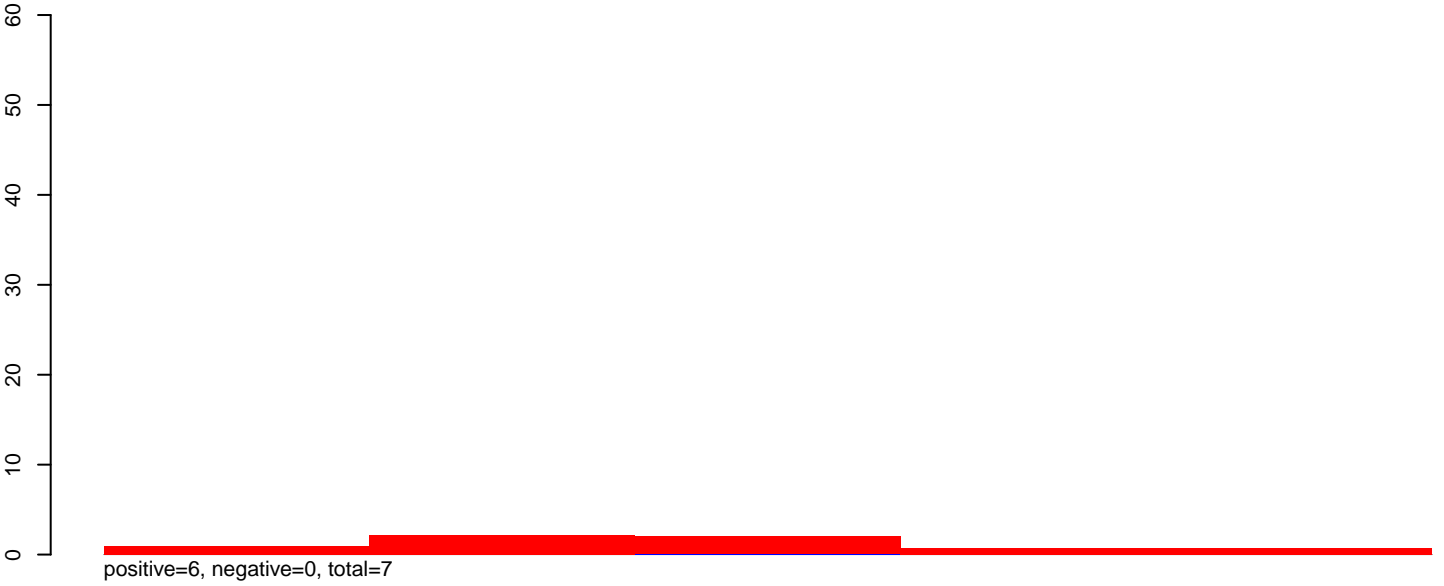
AeAeg_Testes.rep



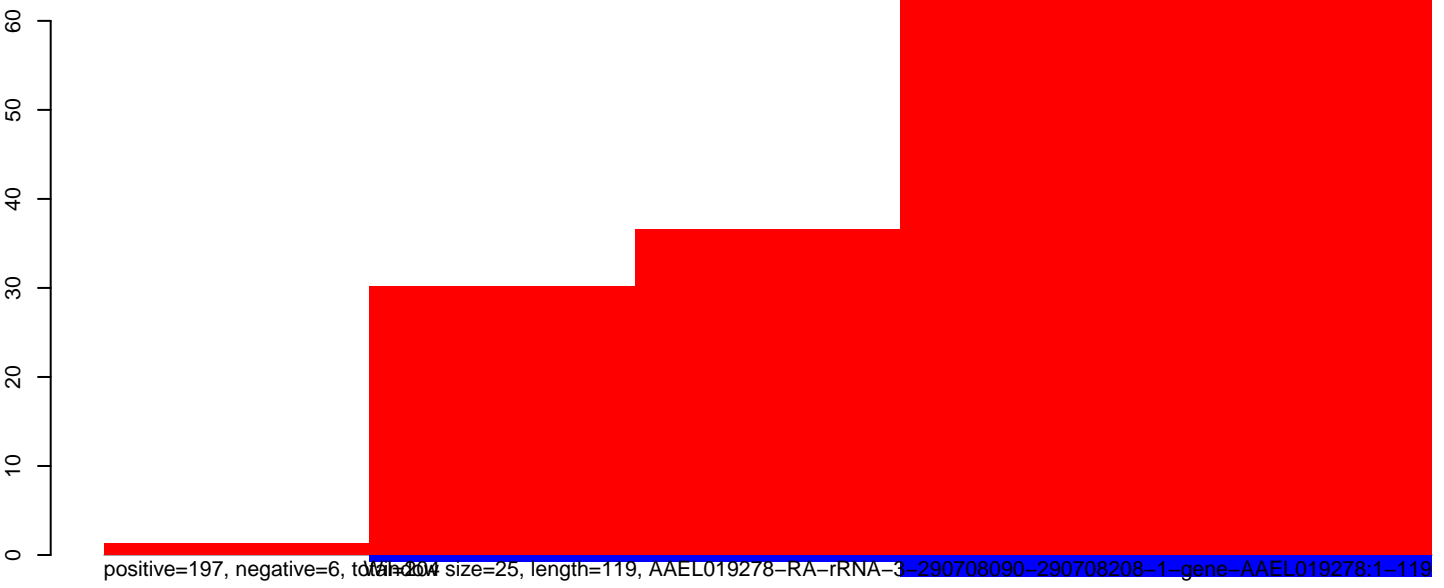
AeAeg_Testes.18_23.rep



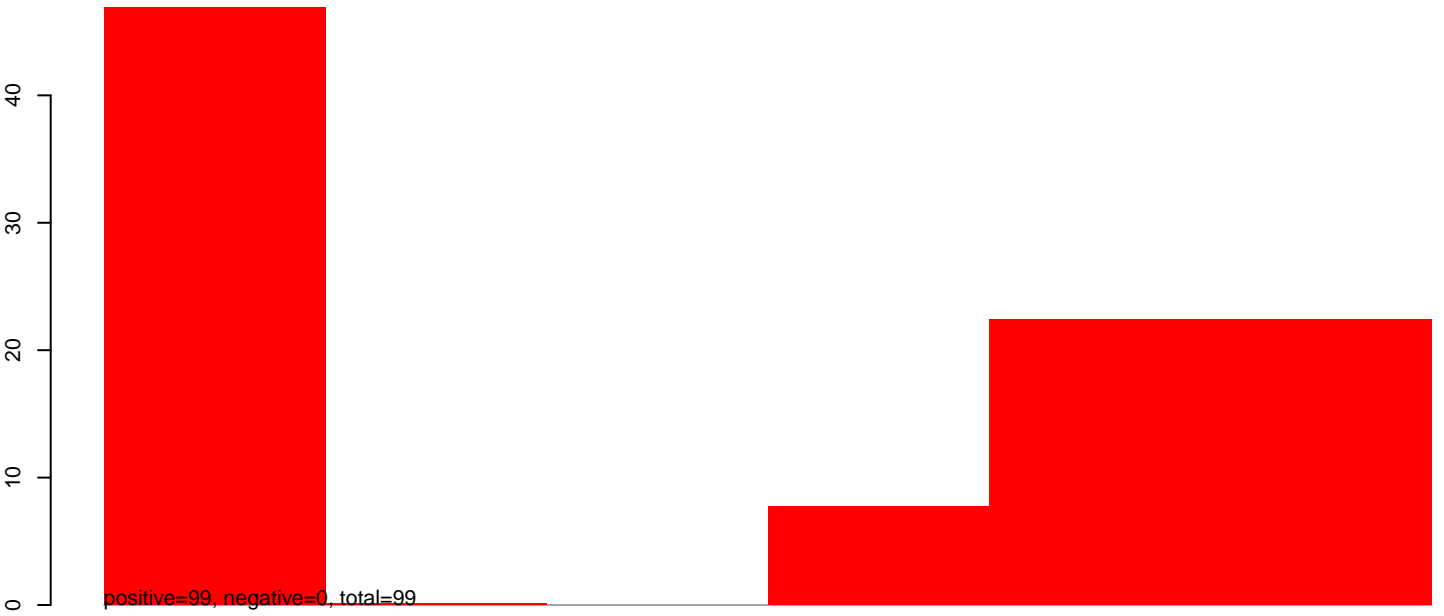
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



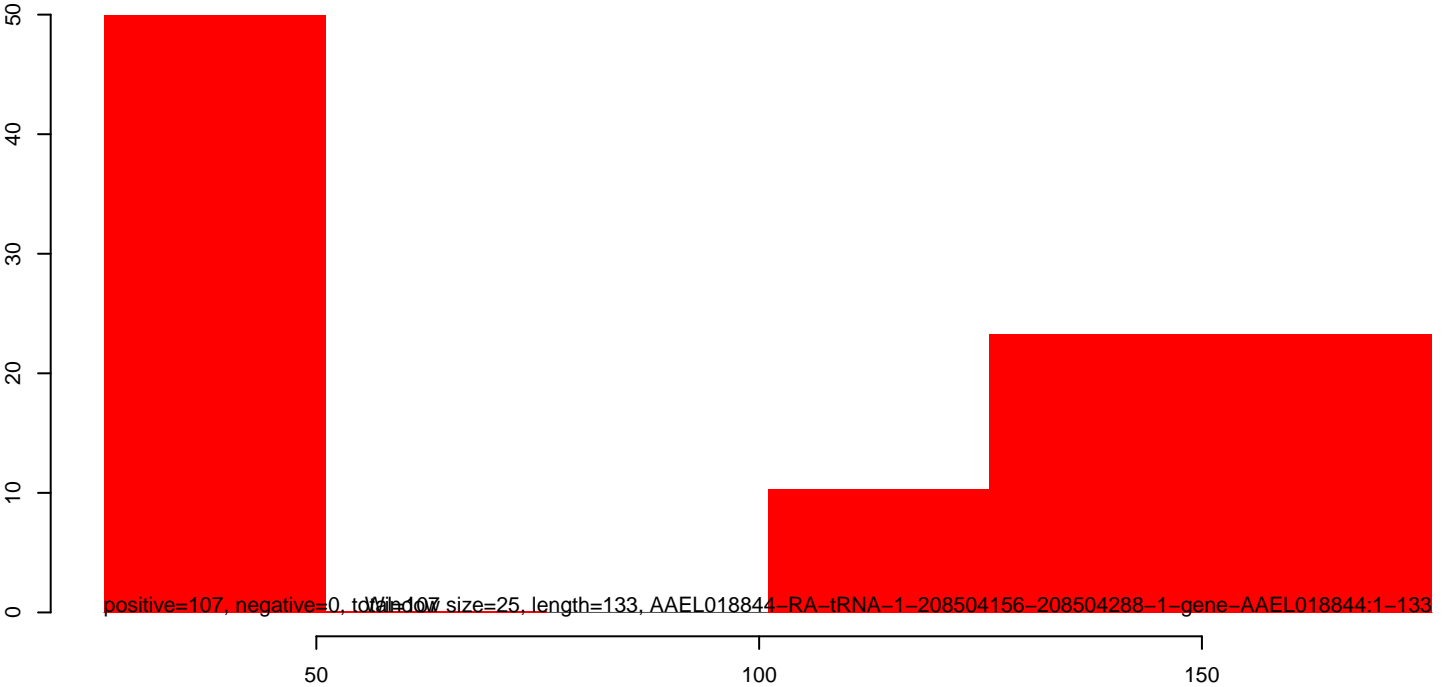
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep

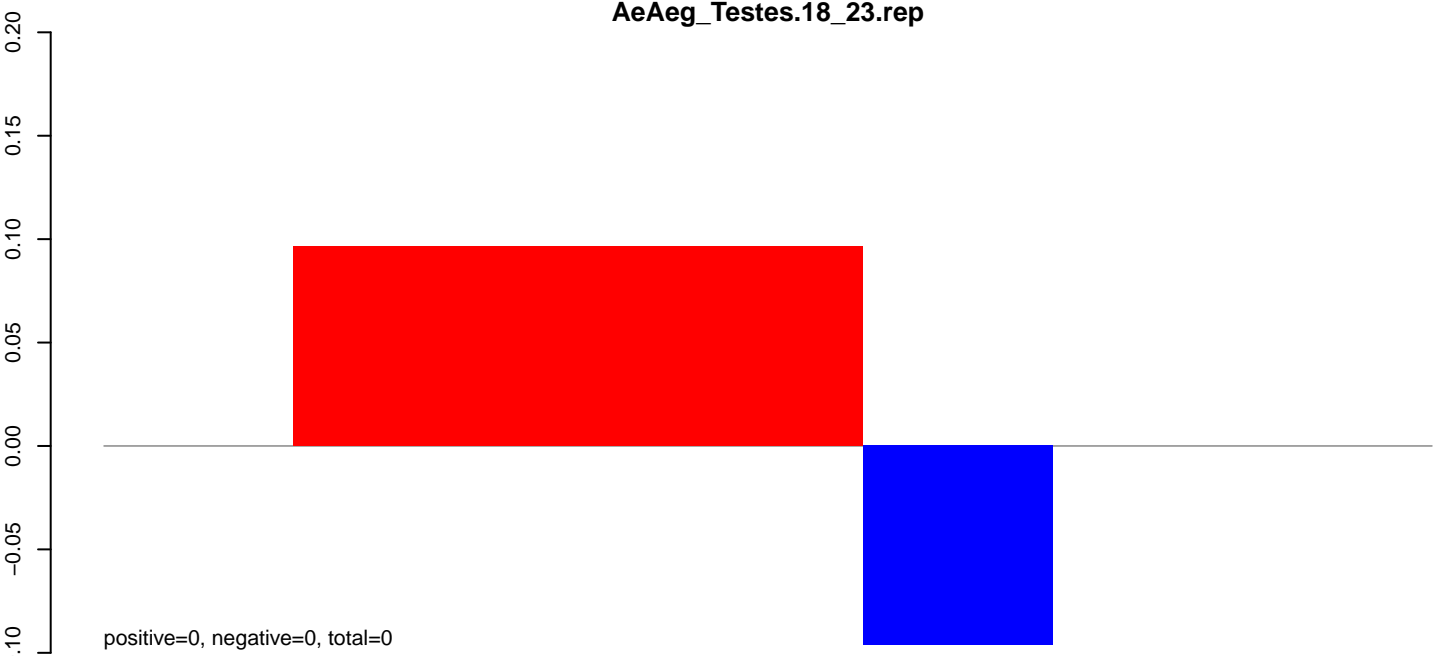


AeAeg_Testes.rep

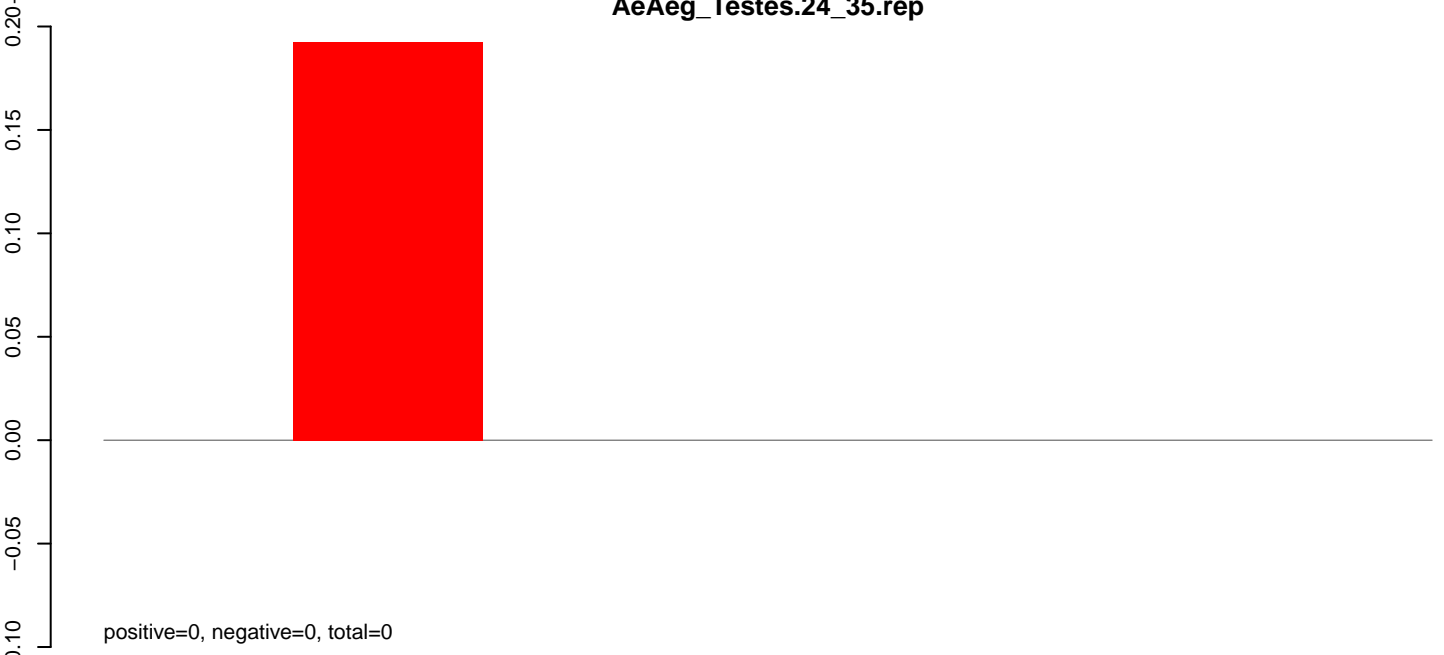


Window size=25, length=133, AAEL018844-RA-tRNA-1-208504156-208504288-1-gene-AAEL018844:1-133

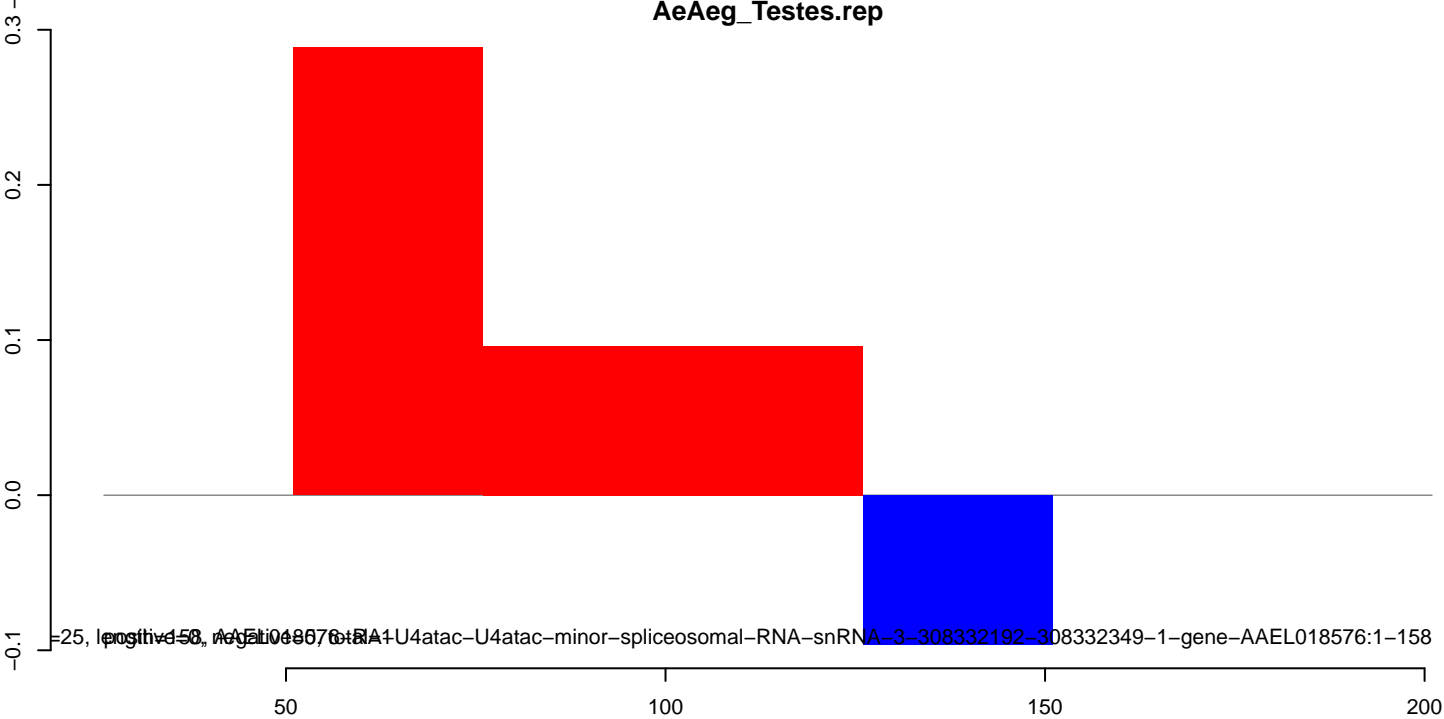
AeAeg_Testes.18_23.rep



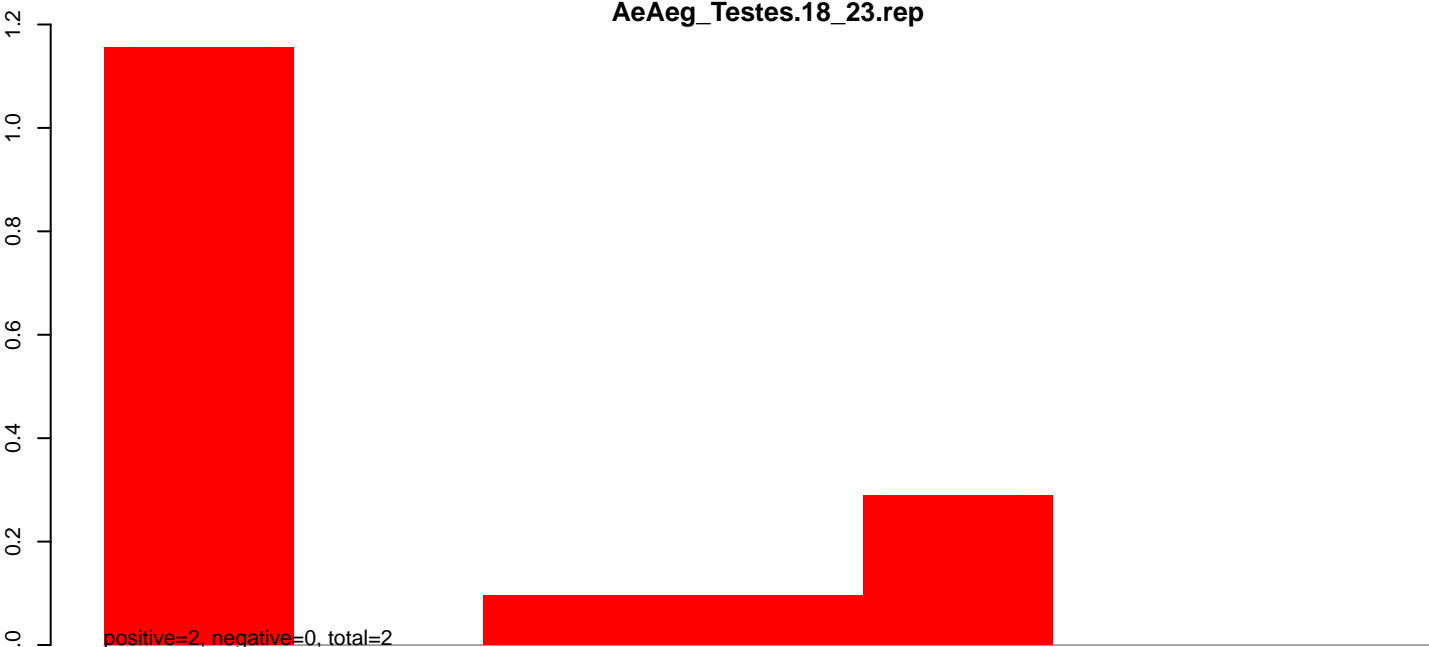
AeAeg_Testes.24_35.rep



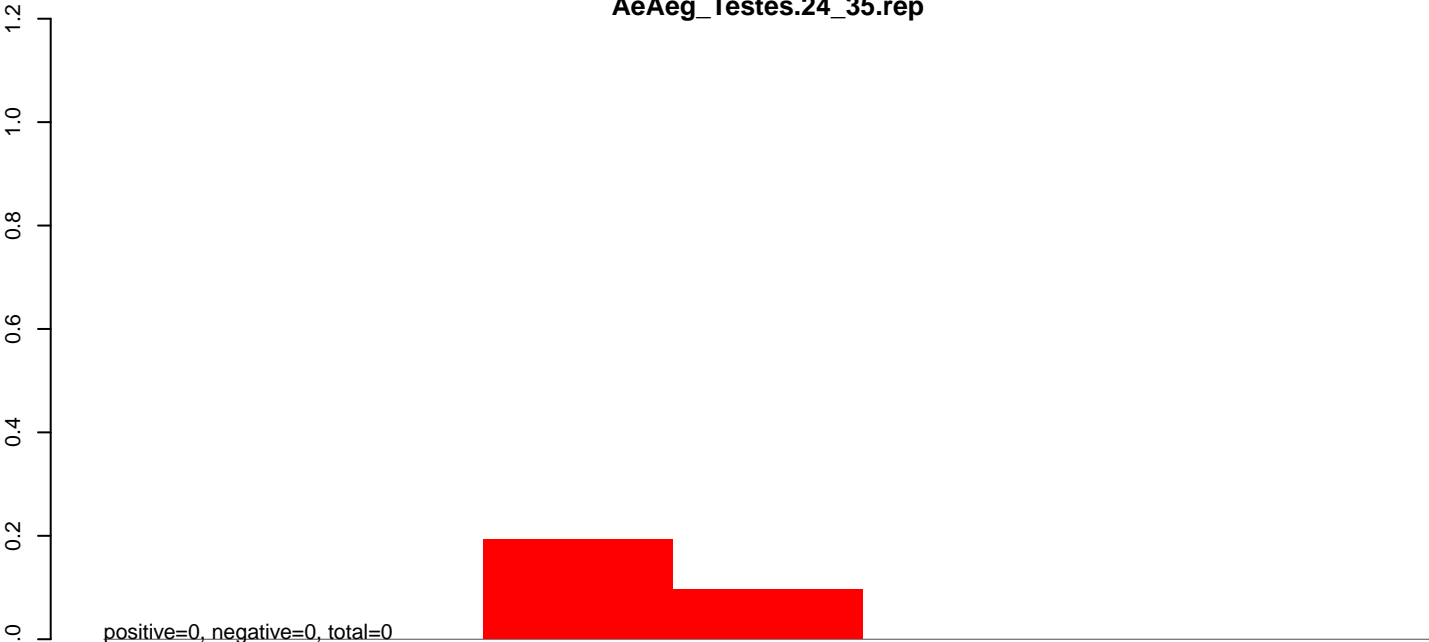
AeAeg_Testes.rep



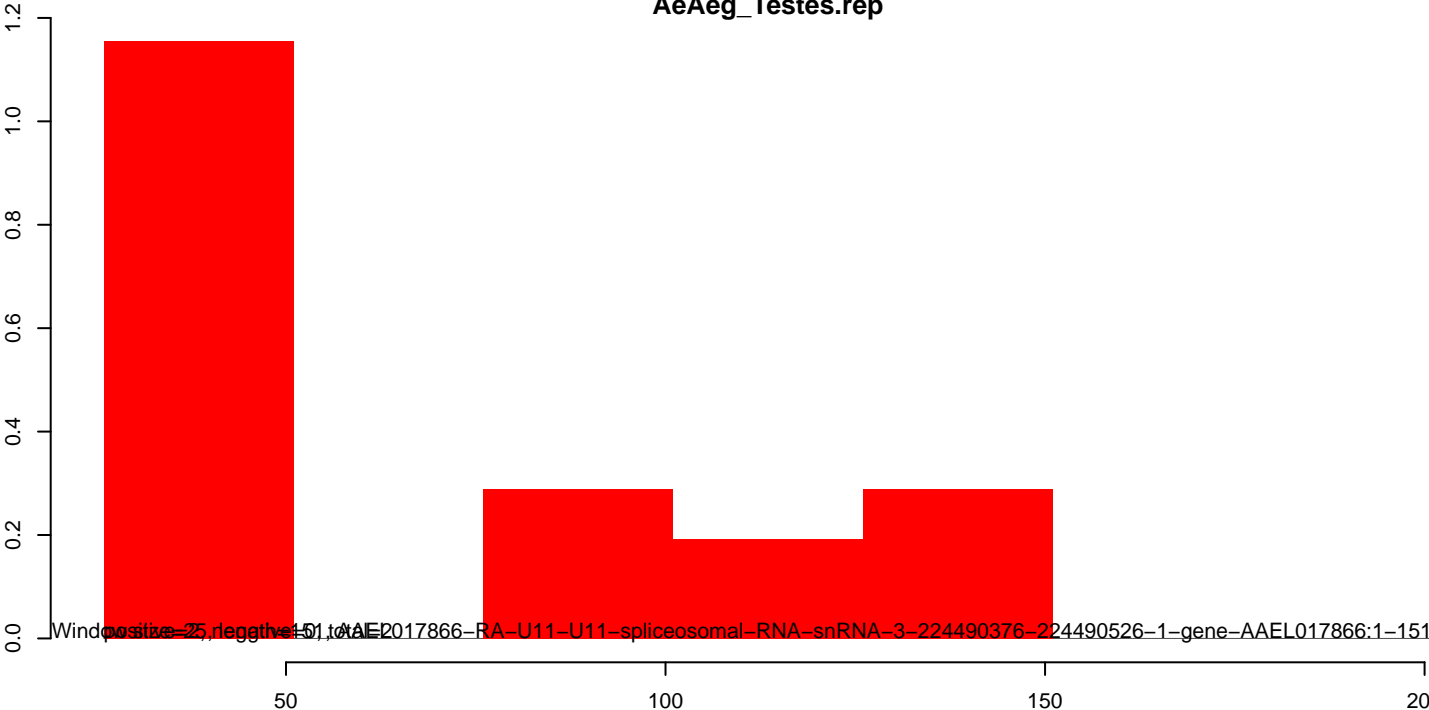
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



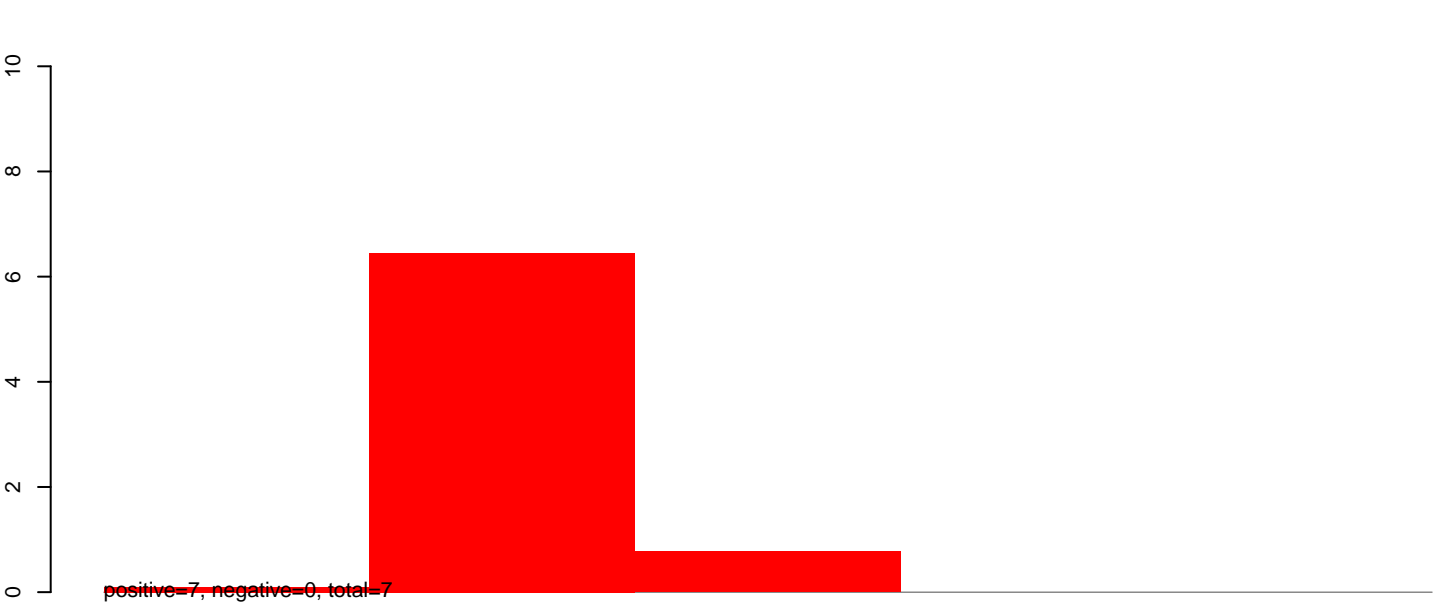
AeAeg_Testes.rep



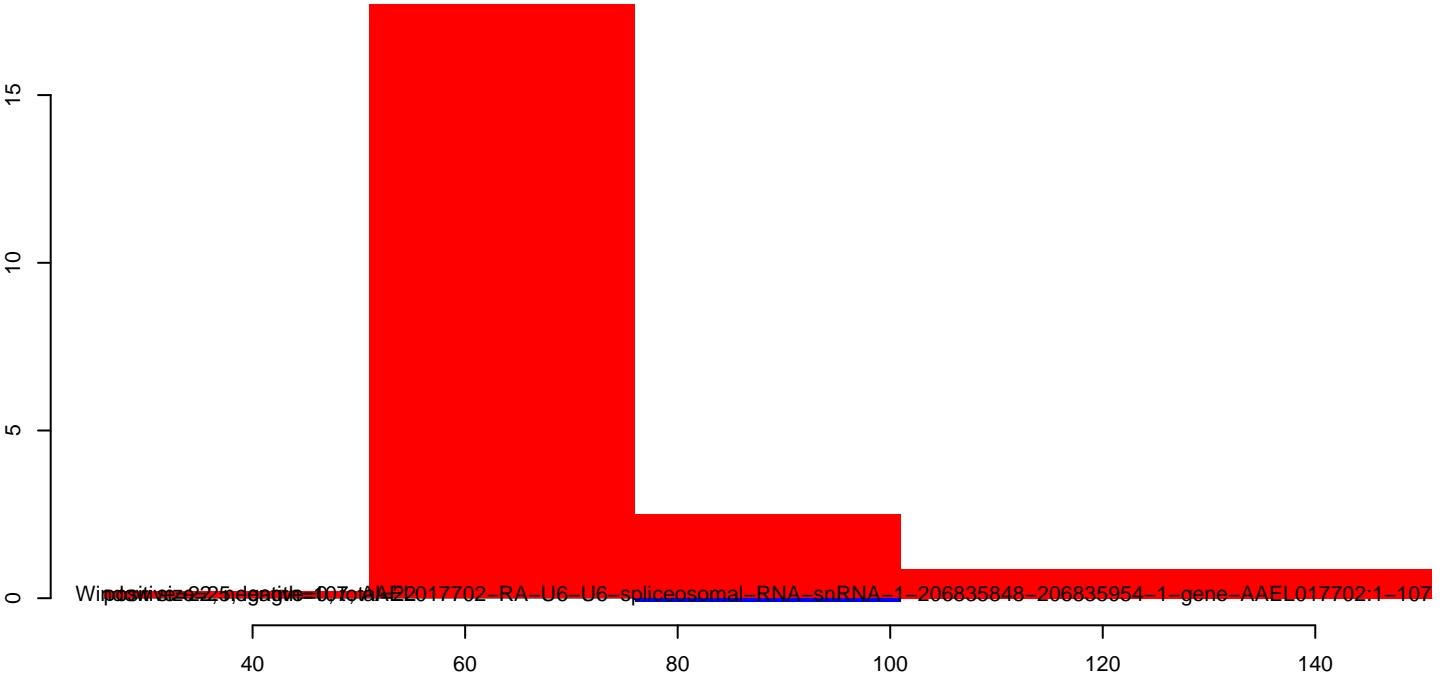
AeAeg_Testes.18_23.rep



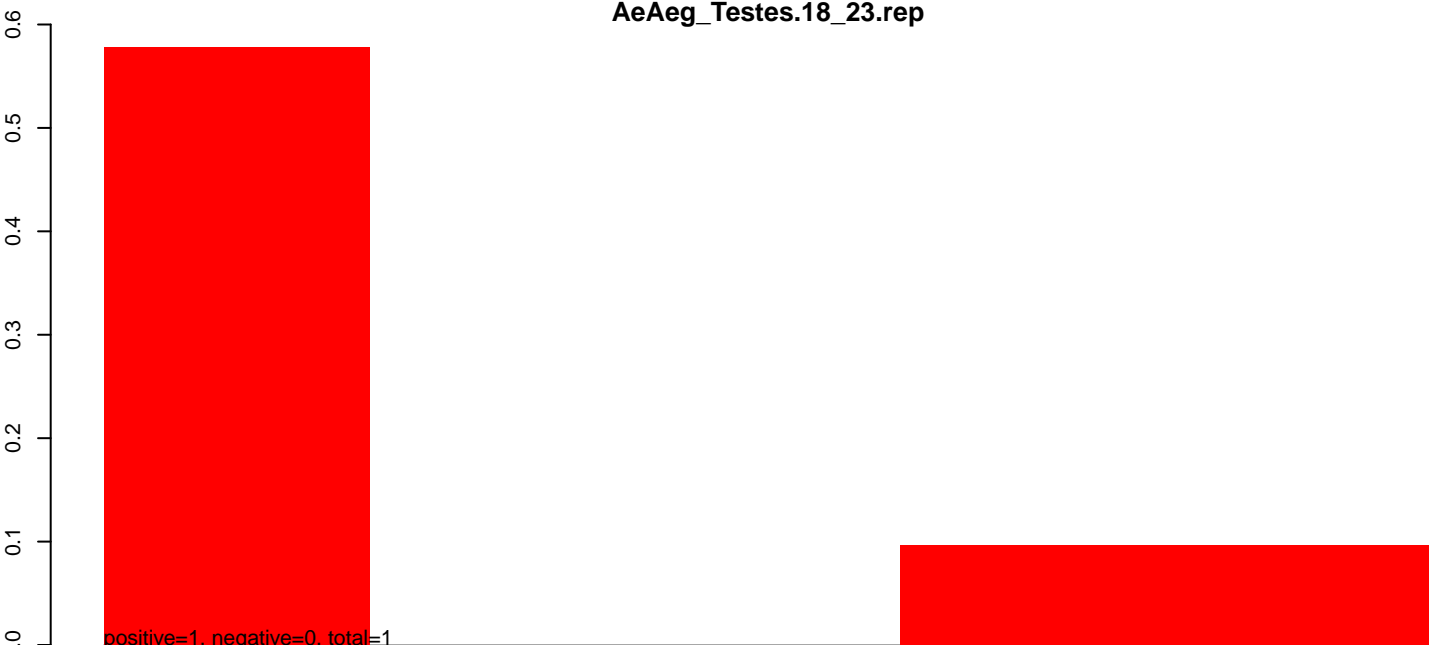
AeAeg_Testes.24_35.rep



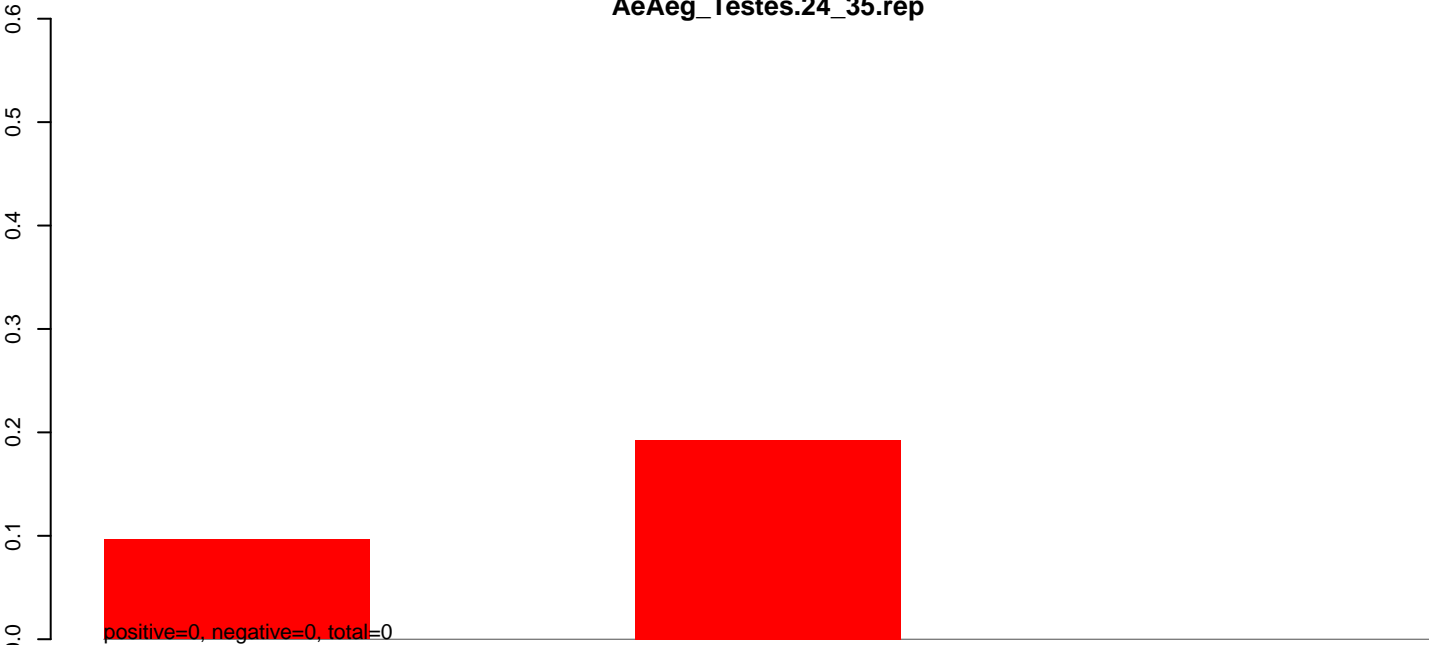
AeAeg_Testes.rep



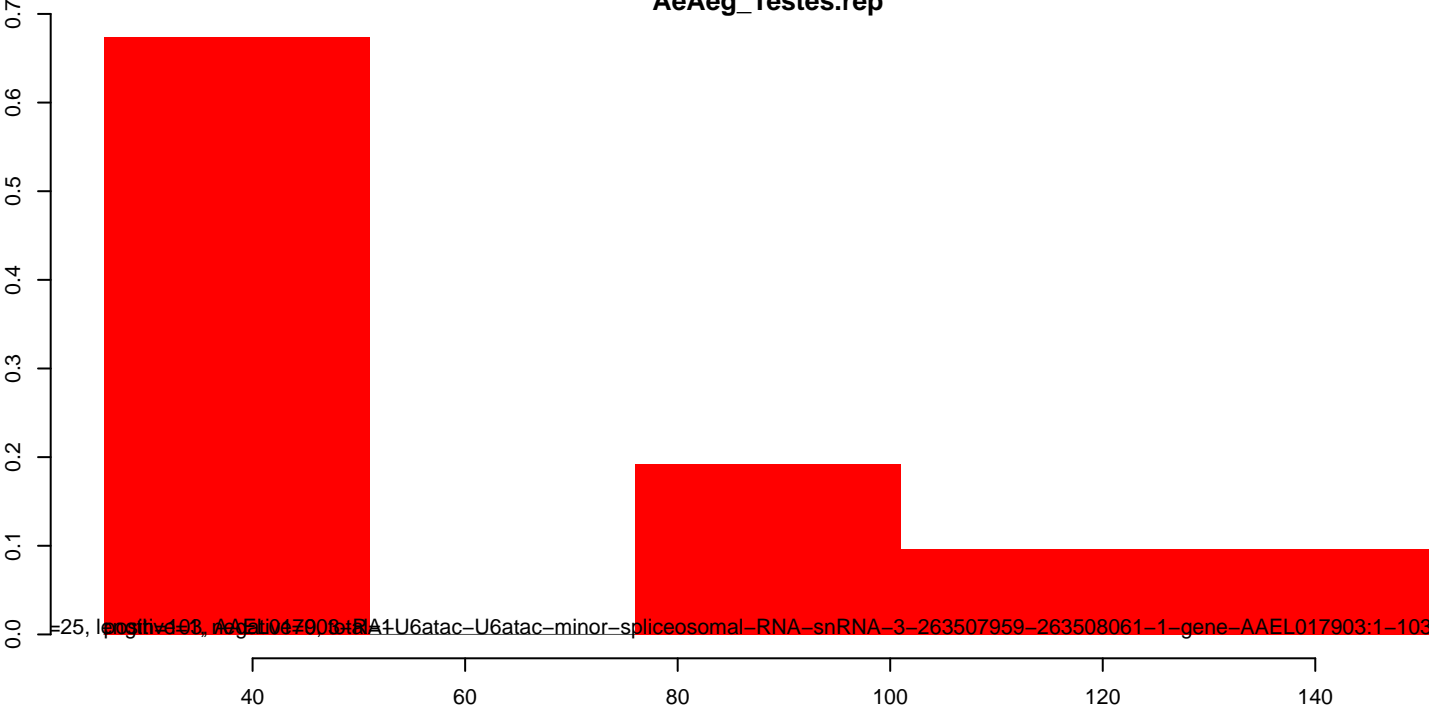
AeAeg_Testes.18_23.rep



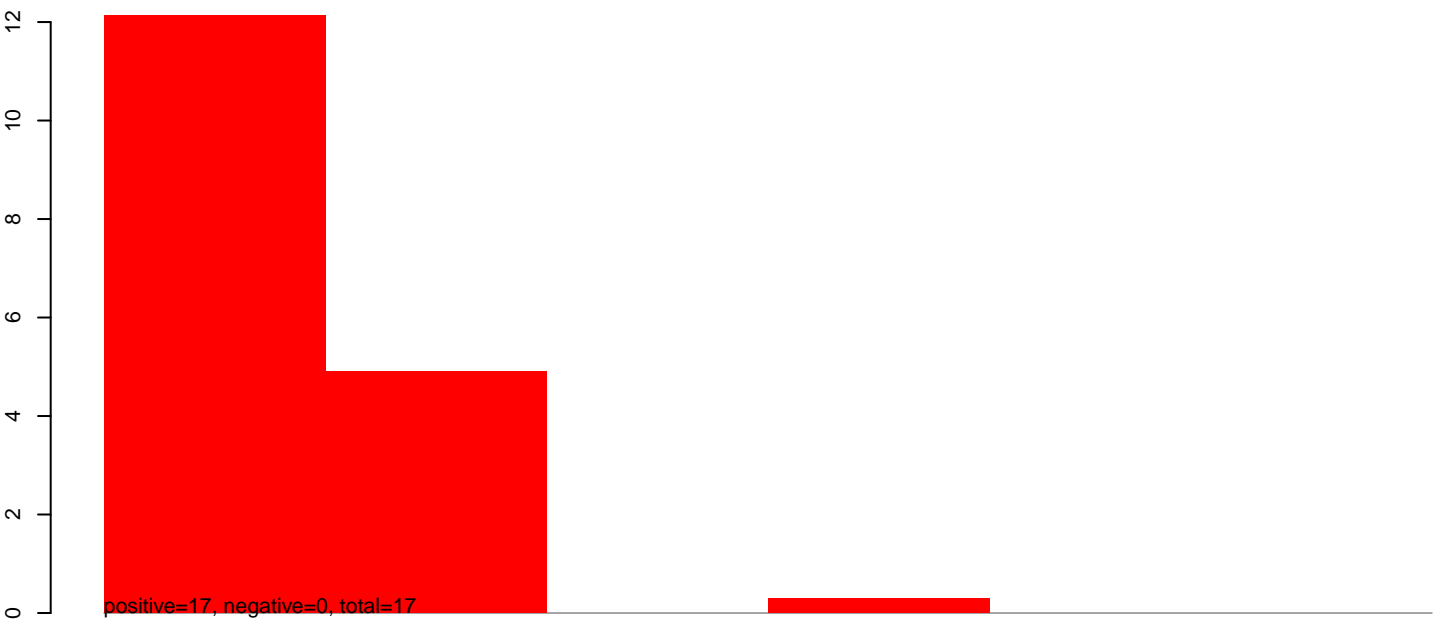
AeAeg_Testes.24_35.rep



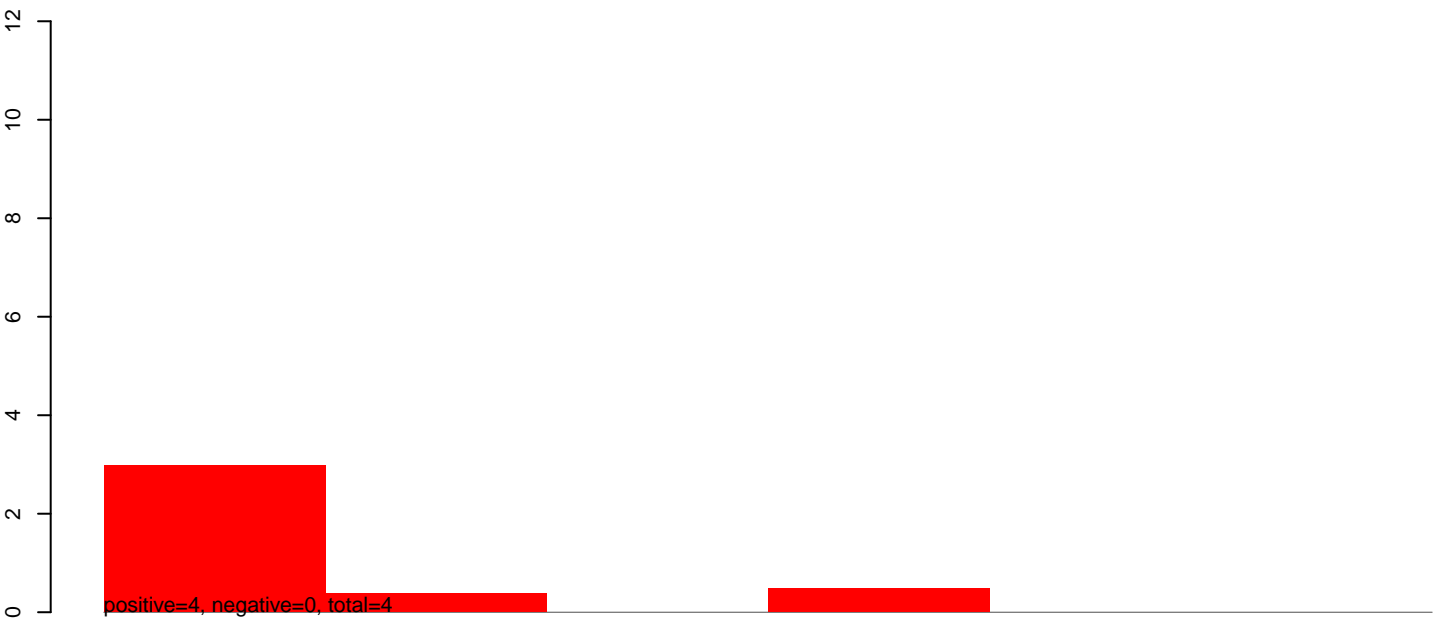
AeAeg_Testes.rep



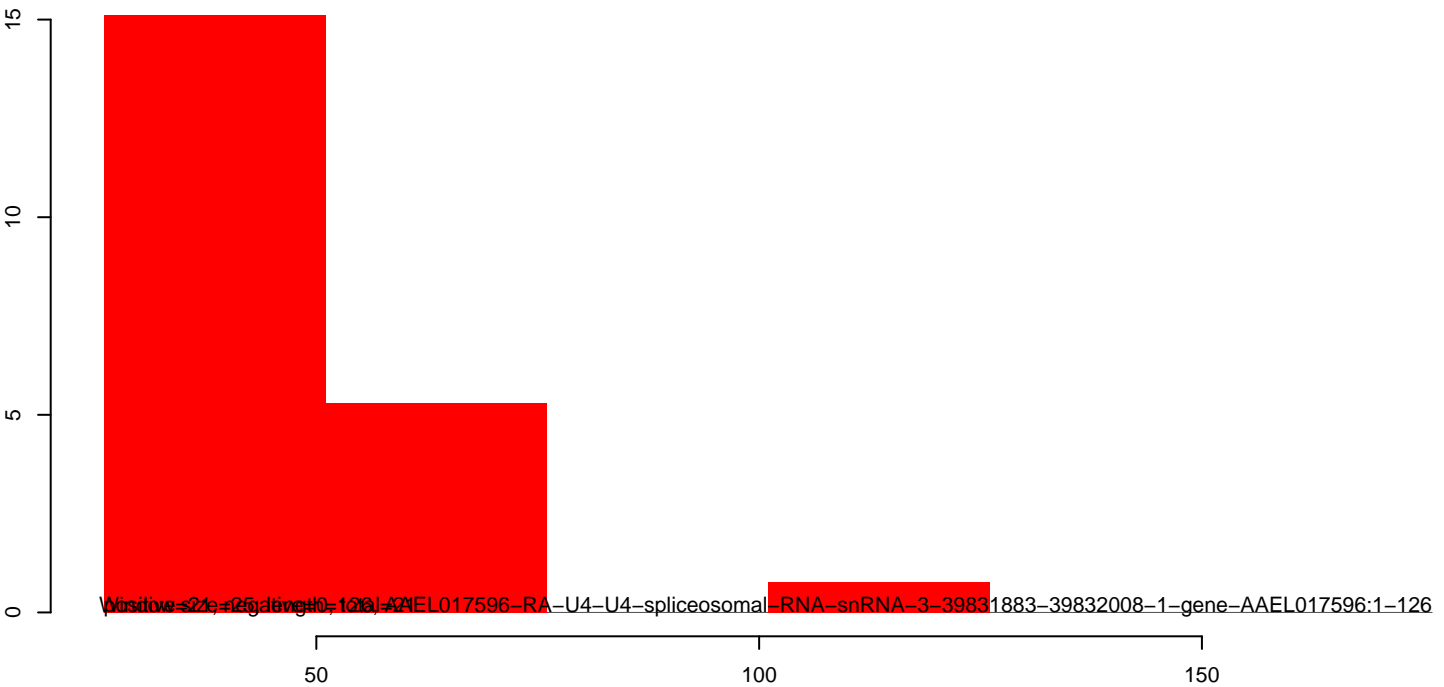
AeAeg_Testes.18_23.rep



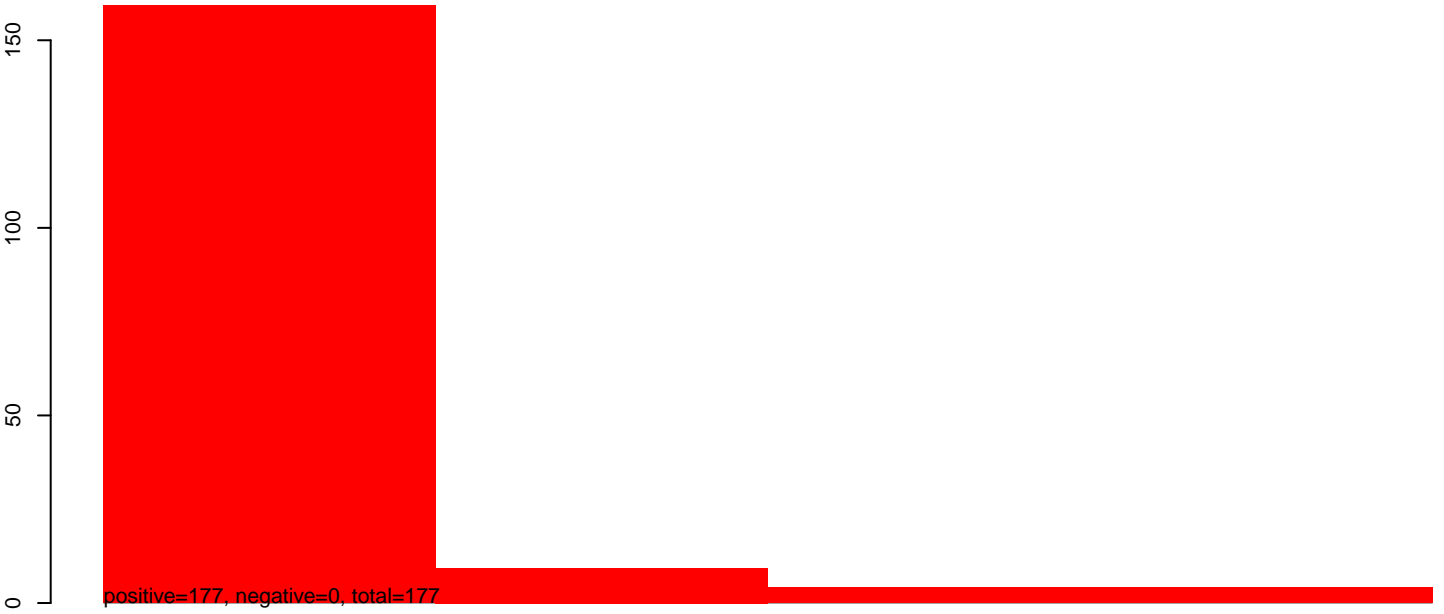
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



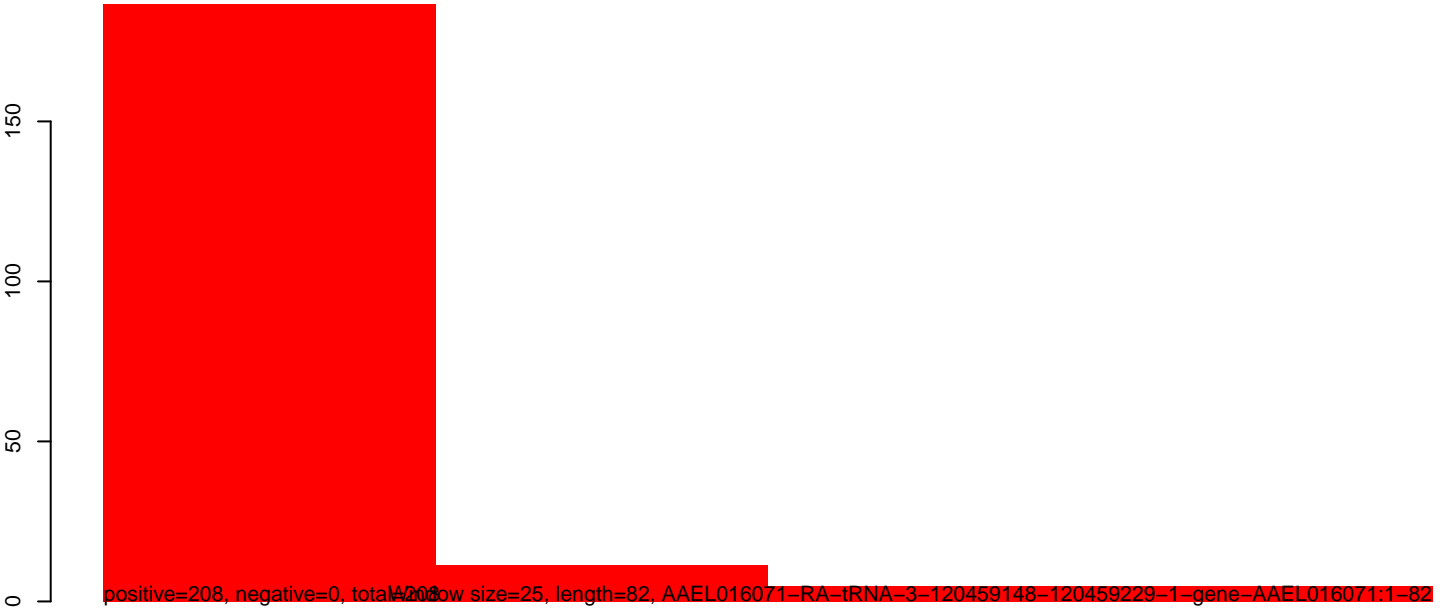
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep

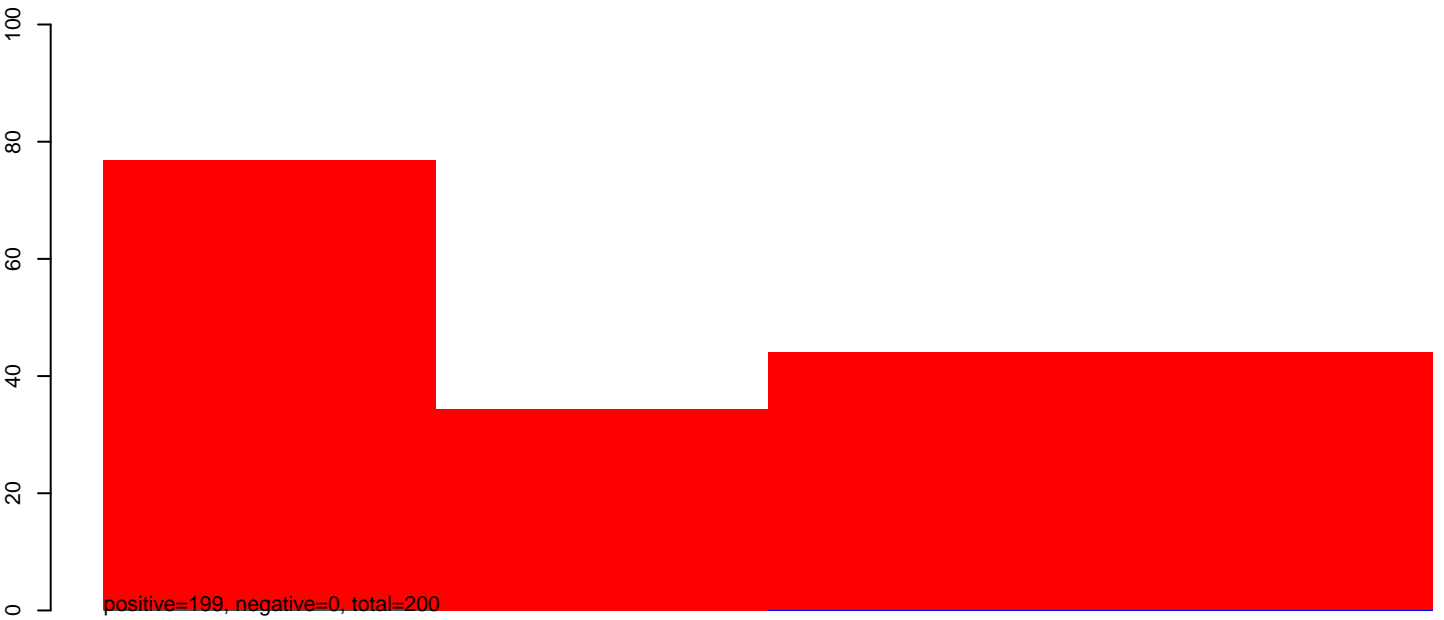


AeAeg_Testes.rep

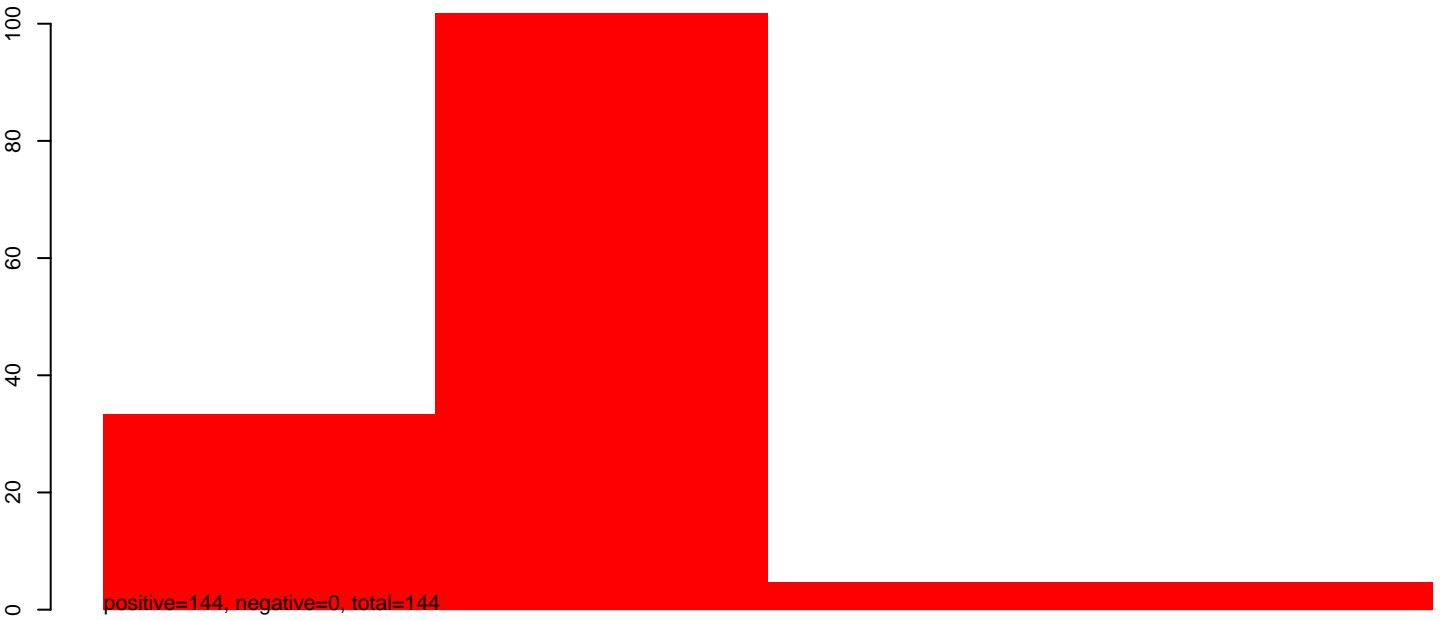


40 60 80 100 120

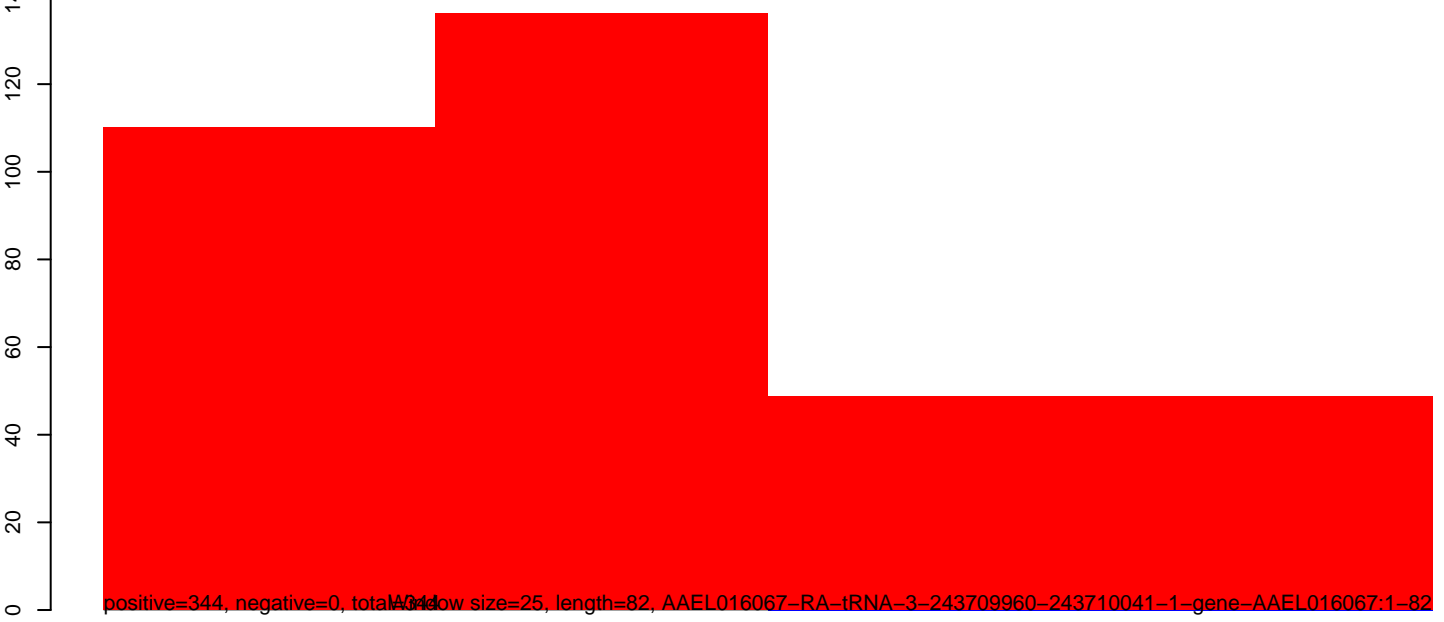
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep

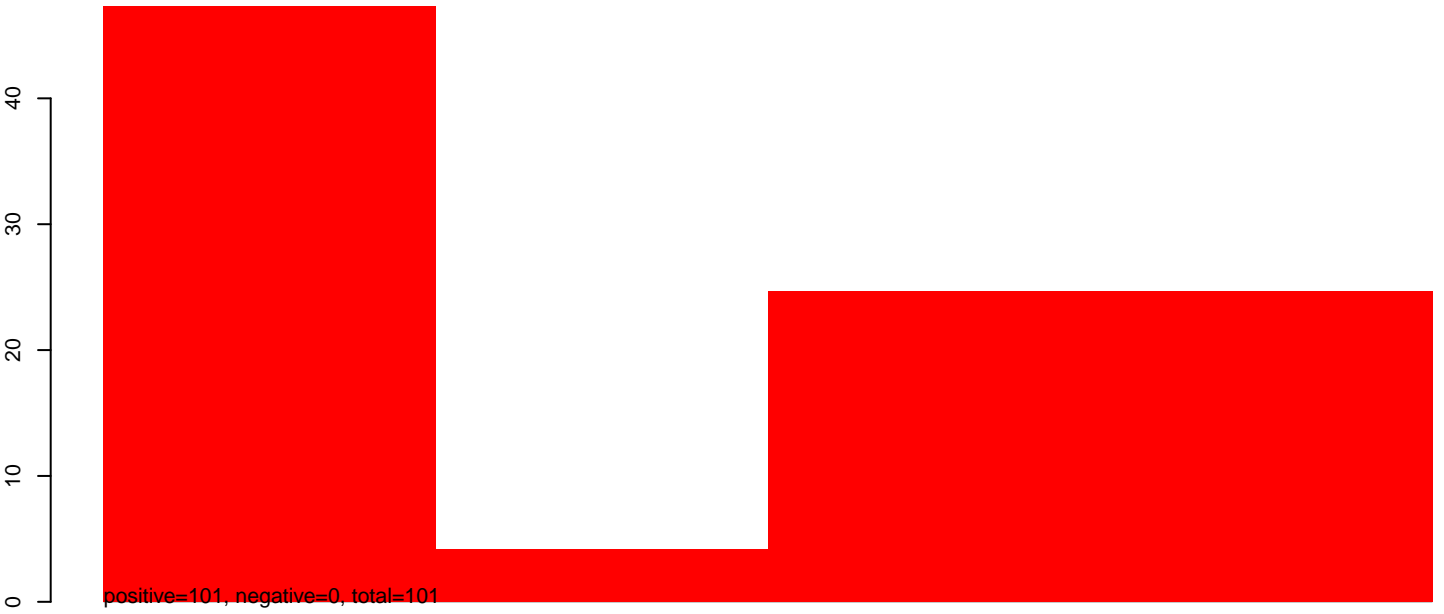


AeAeg_Testes.rep



Window size=25, length=82, AAEL016067-RA-tRNA-3-243709960-243710041-1-gene-AAEL016067:1-82

AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep

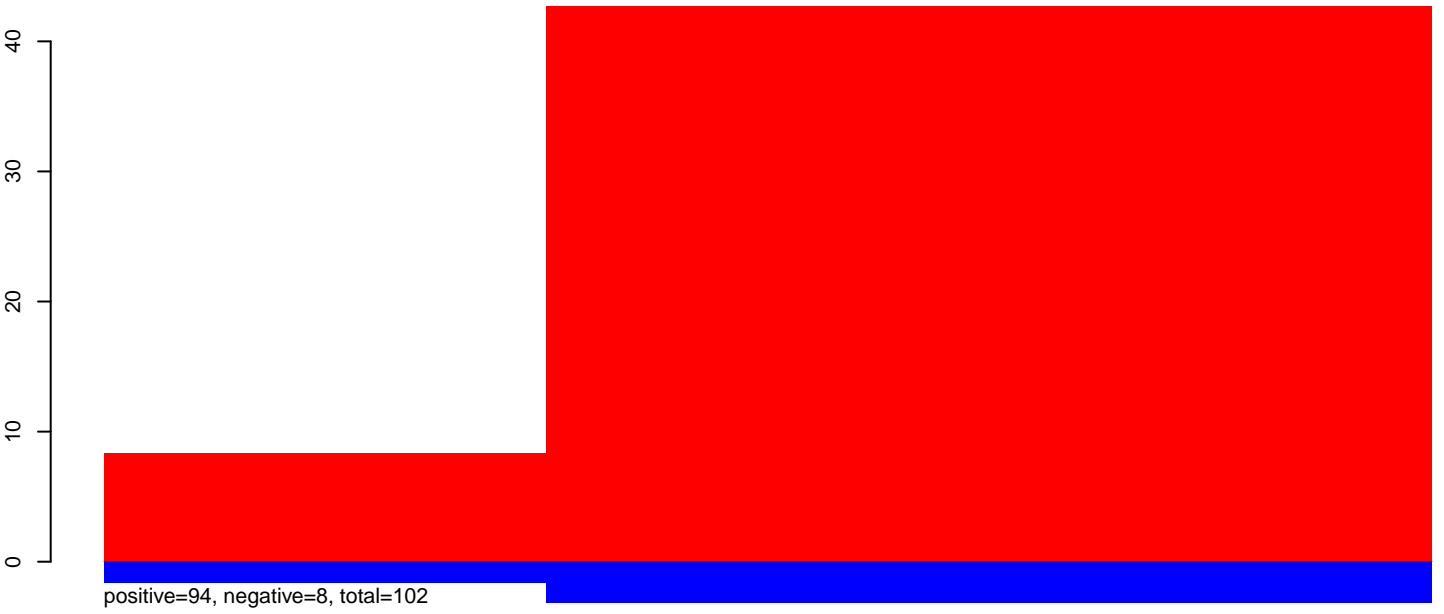


AeAeg_Testes.rep



40 60 80 100 120

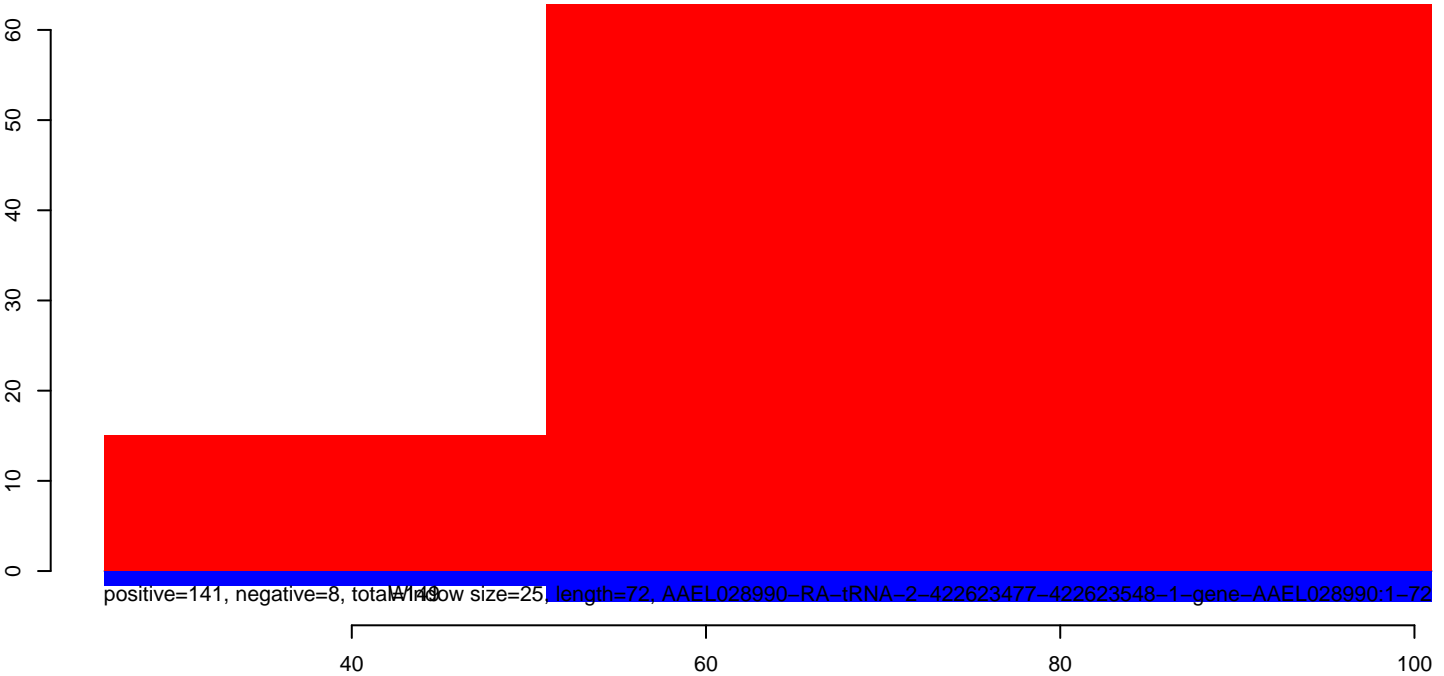
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



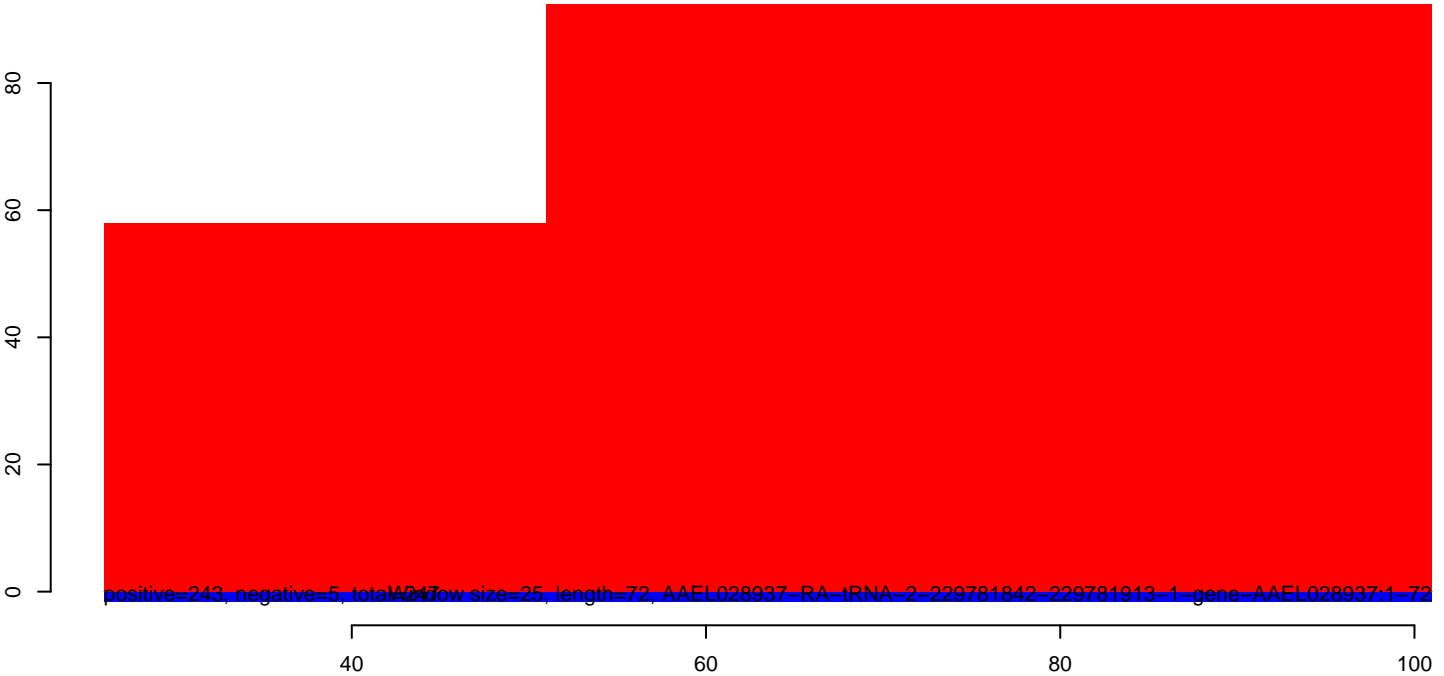
AeAeg_Testes.18_23.rep



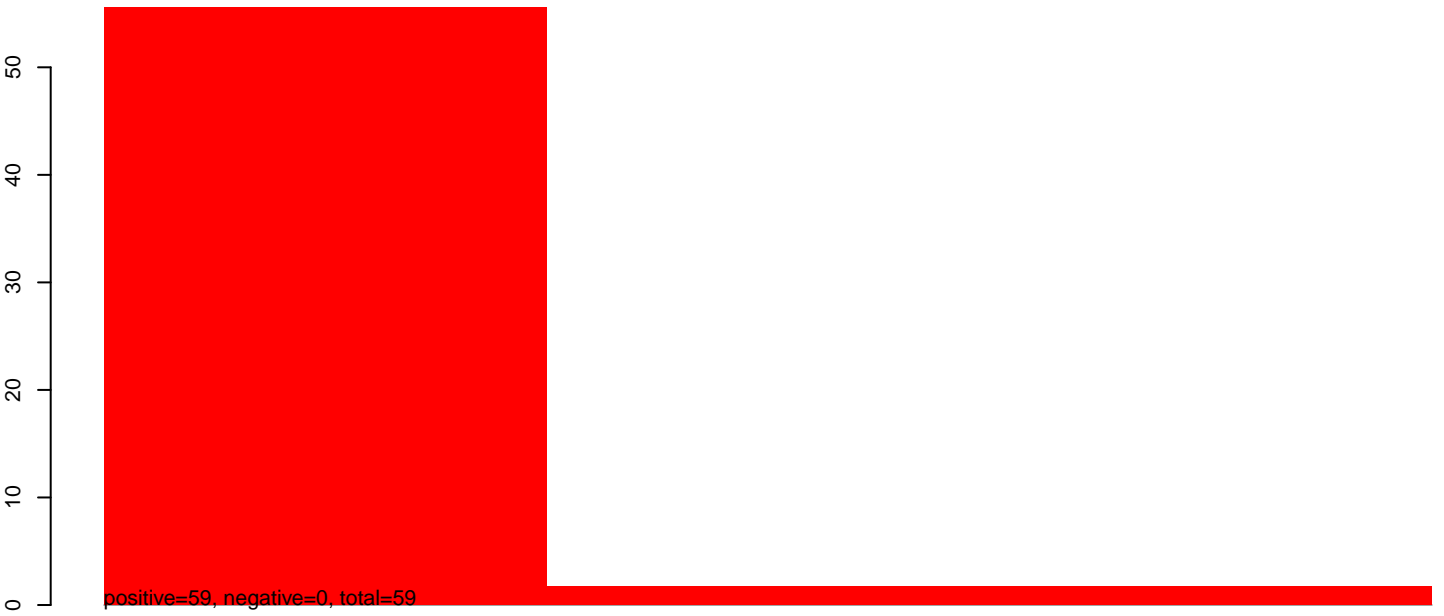
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



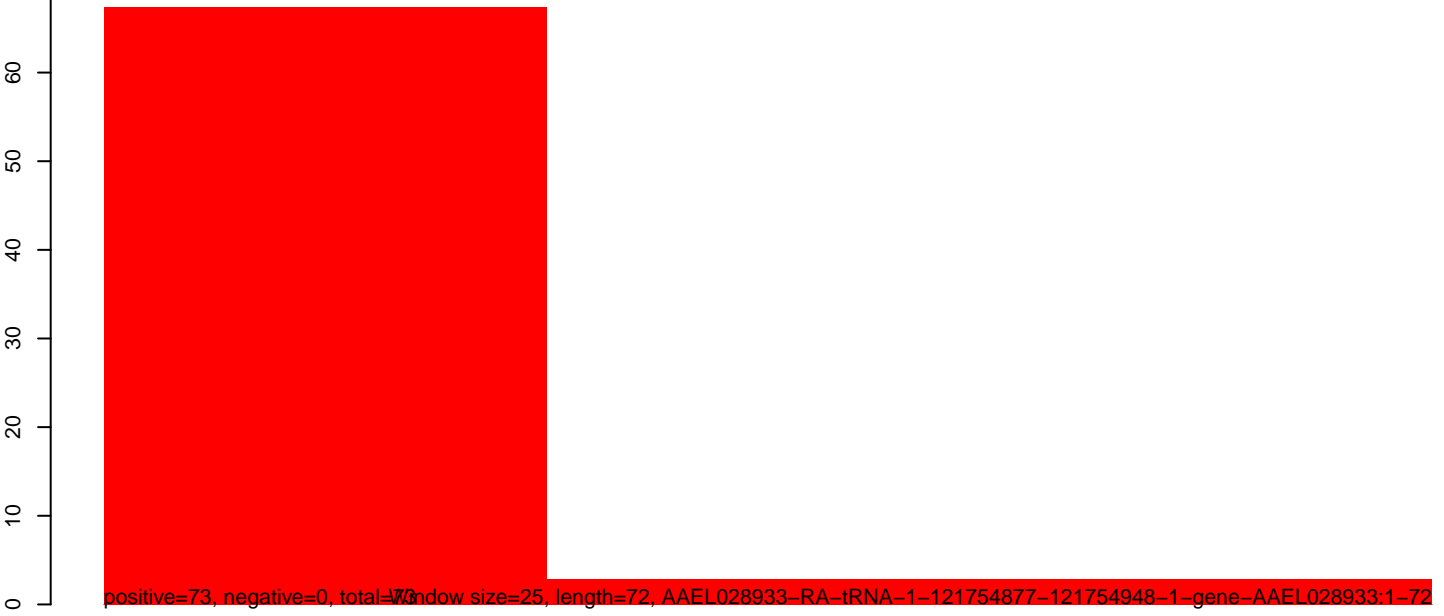
AeAeg_Testes.18_23.rep



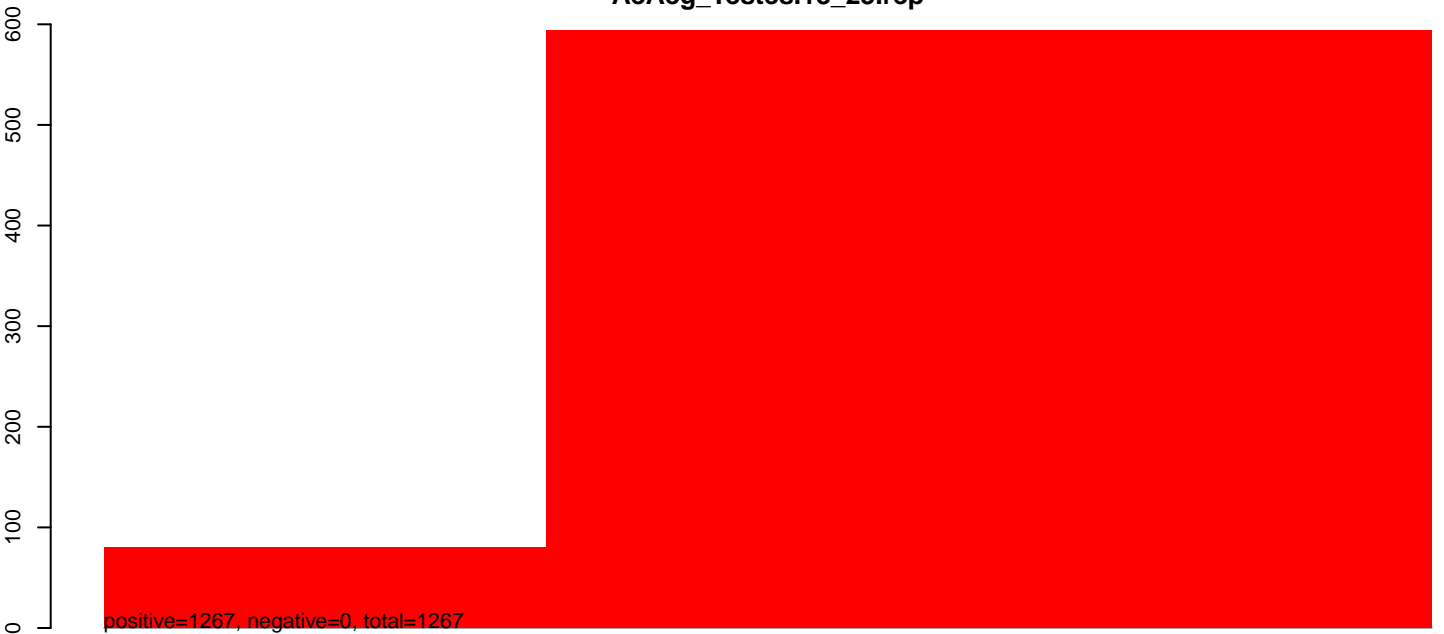
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep

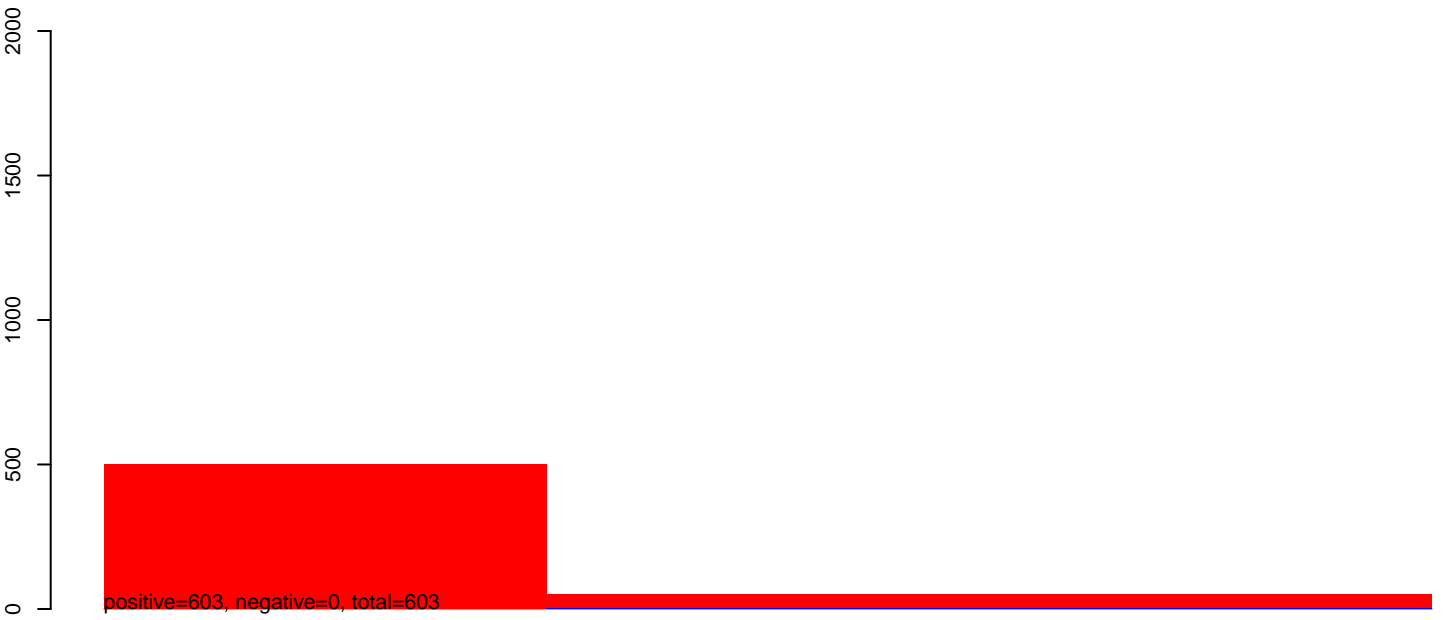


AeAeg_Testes.rep

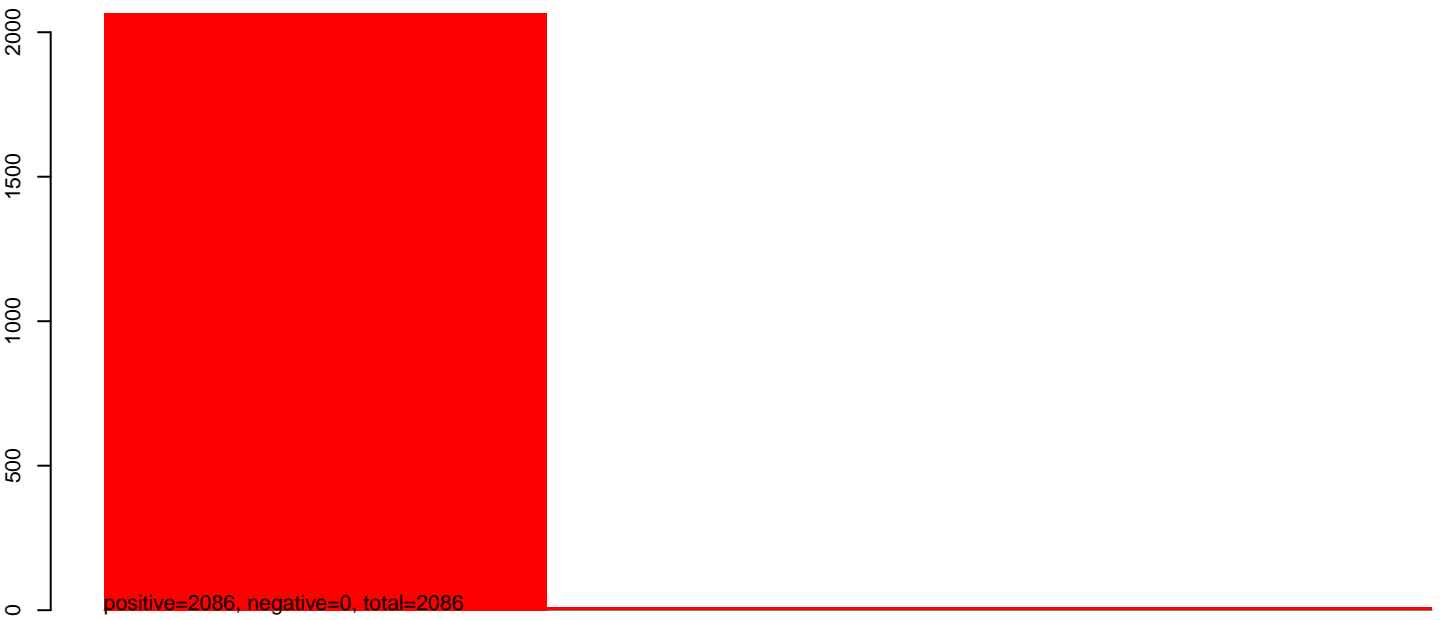


40 60 80 100
Window size=25, length=72, AAEL028931-RA-tRNA-2-55046956-55047027-1-gene-AAEL028931:1-72

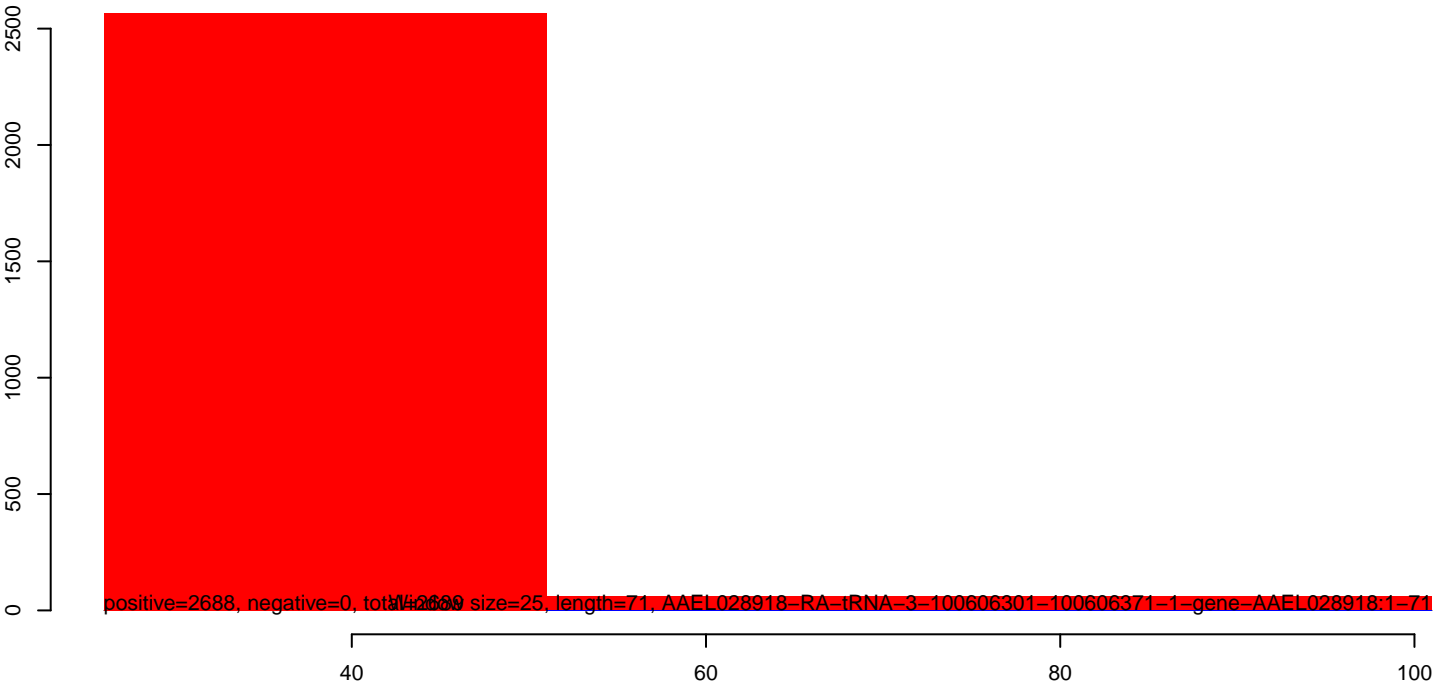
AeAeg_Testes.18_23.rep



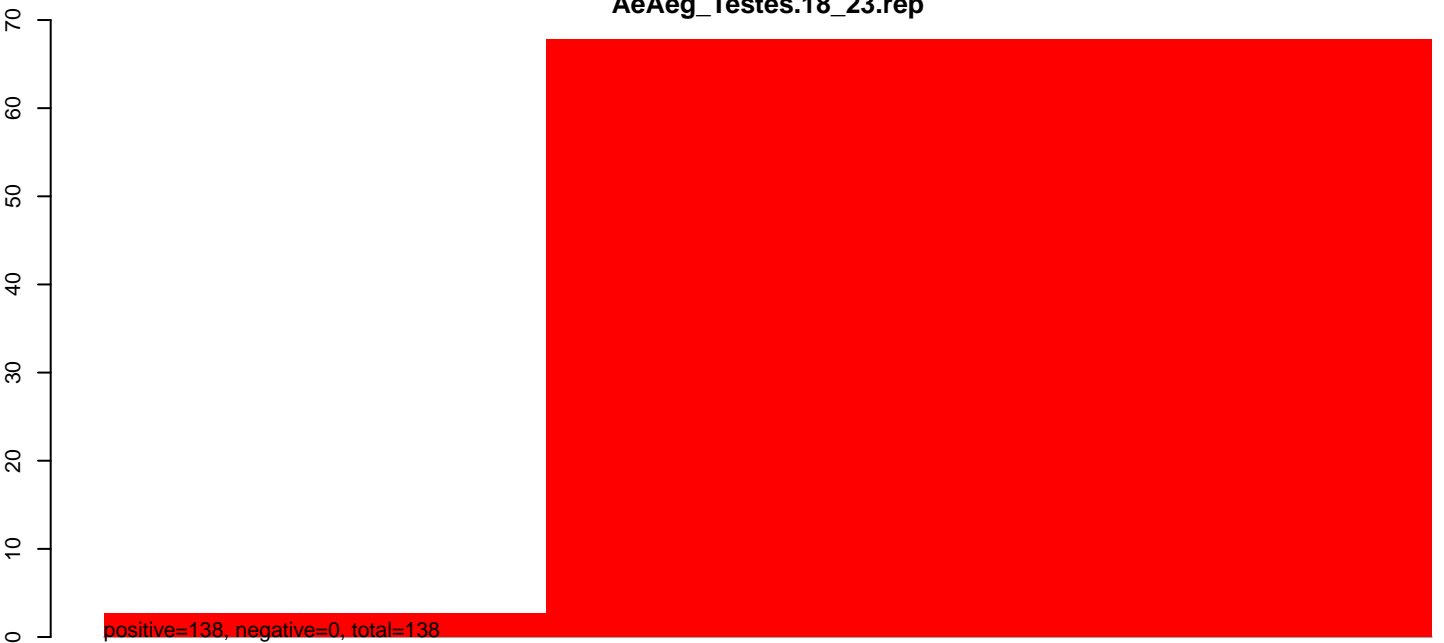
AeAeg_Testes.24_35.rep



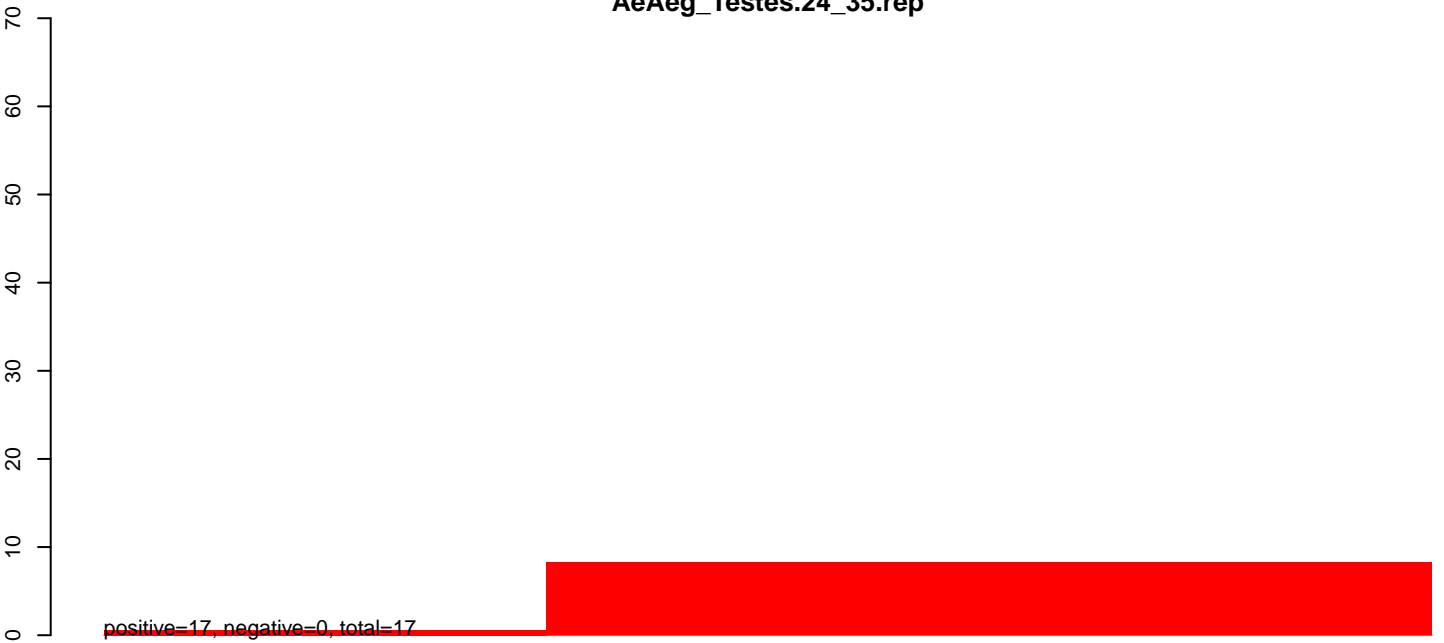
AeAeg_Testes.rep



AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



AeAeg_Testes.rep

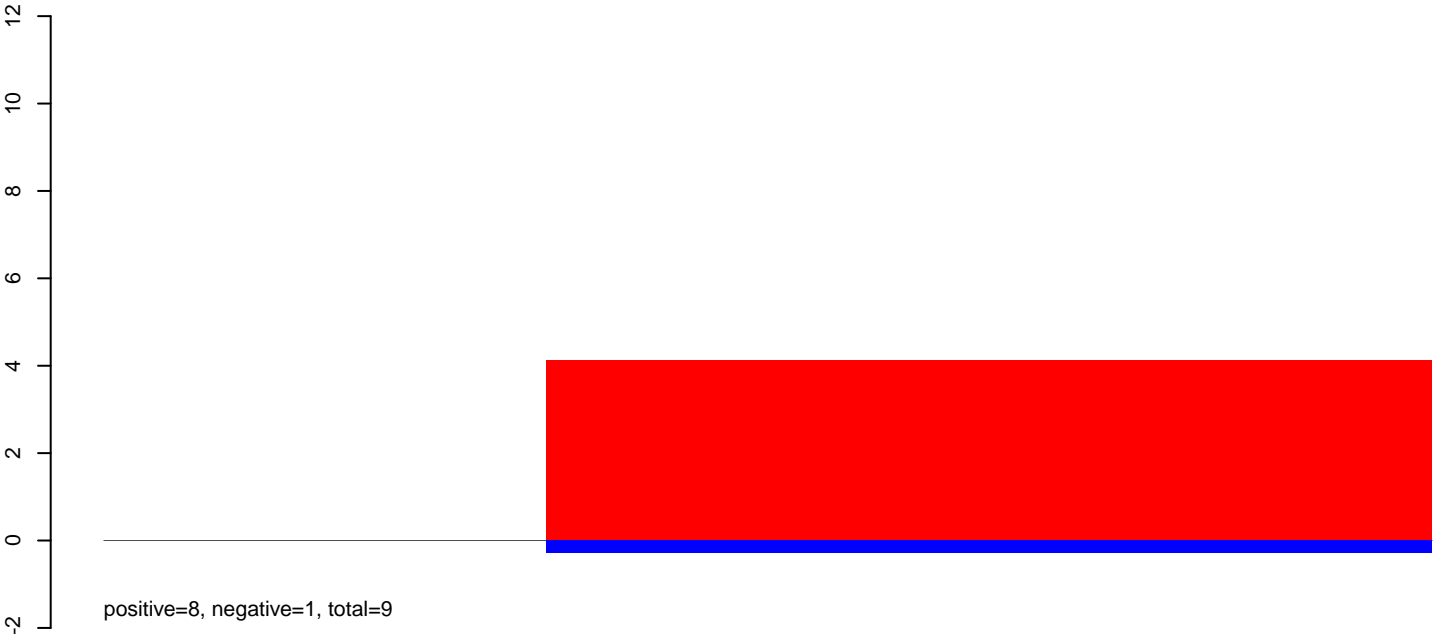


40 60 80 100
AeAeg_Testes.rep: low size=25, length=73, AAEL028693-RA-tRNA-3-100605474-100605546-1-gene-AAEL028693:1-73

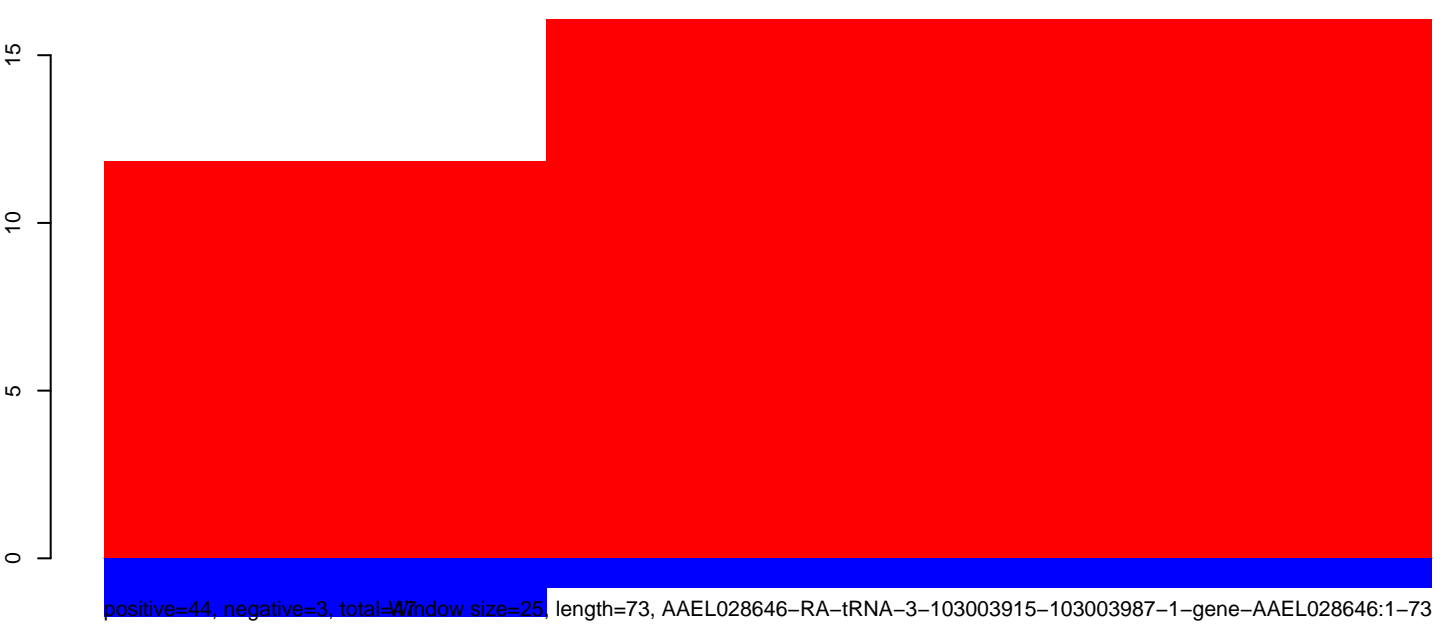
AeAeg_Testes.18_23.rep



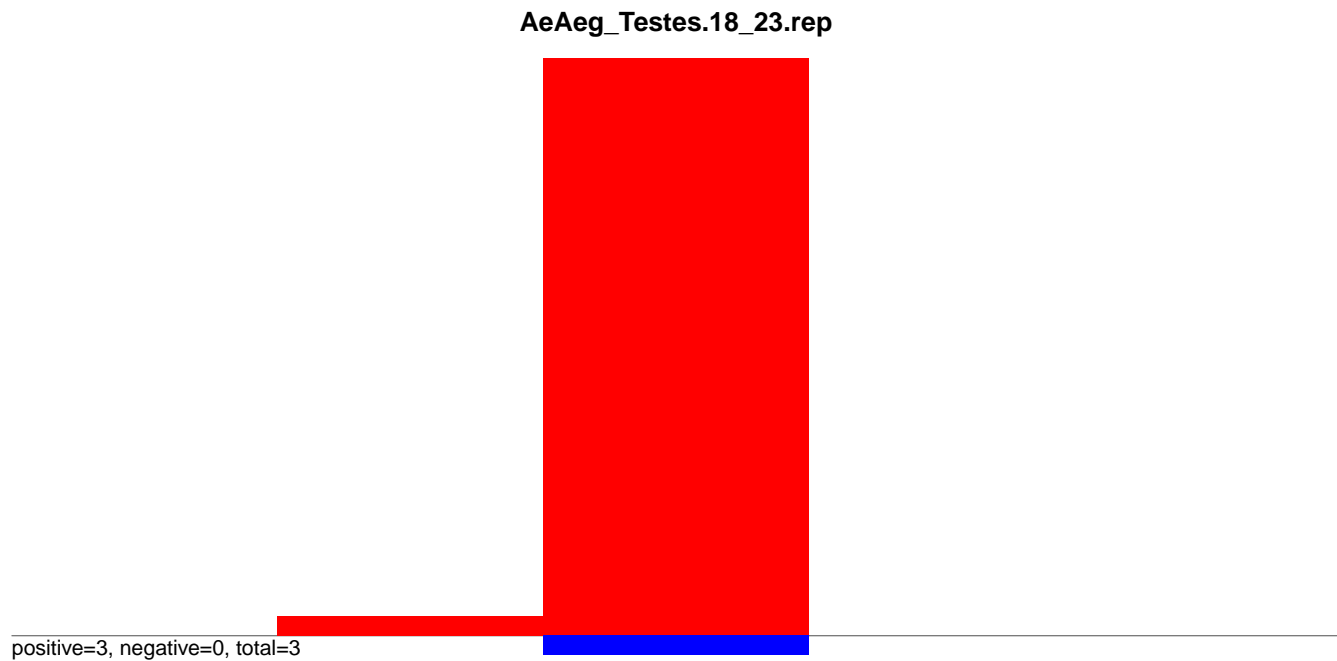
AeAeg_Testes.24_35.rep



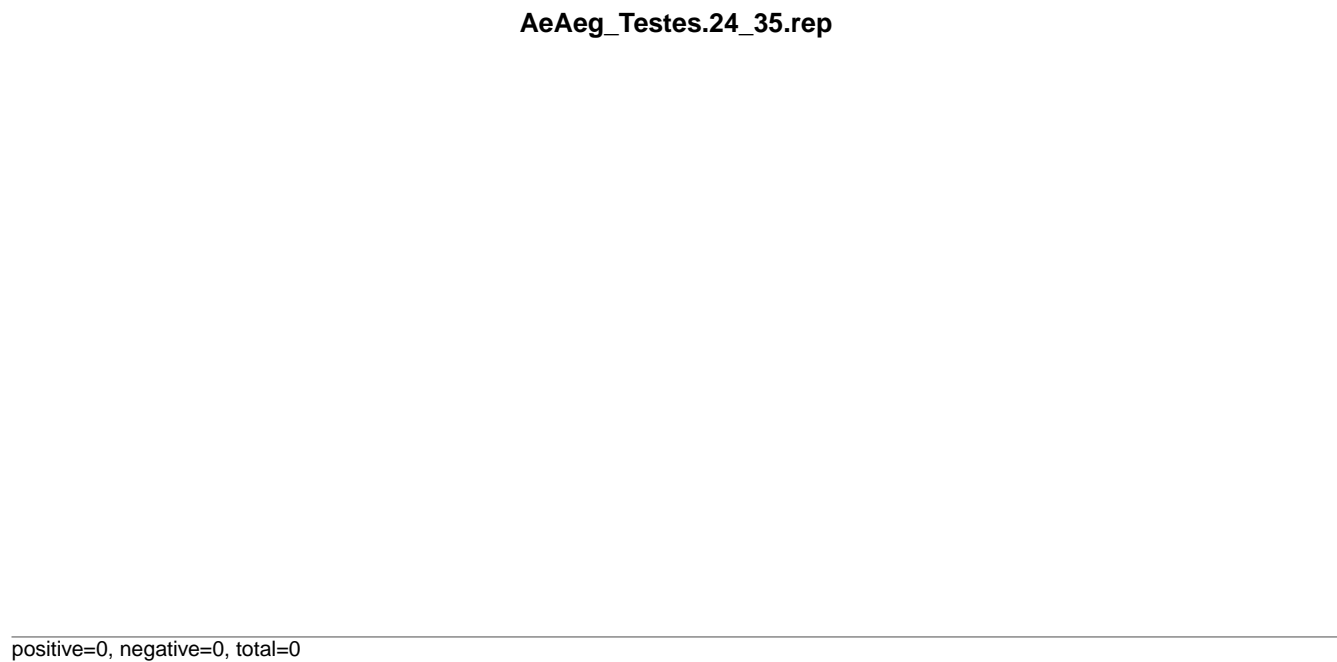
AeAeg_Testes.rep



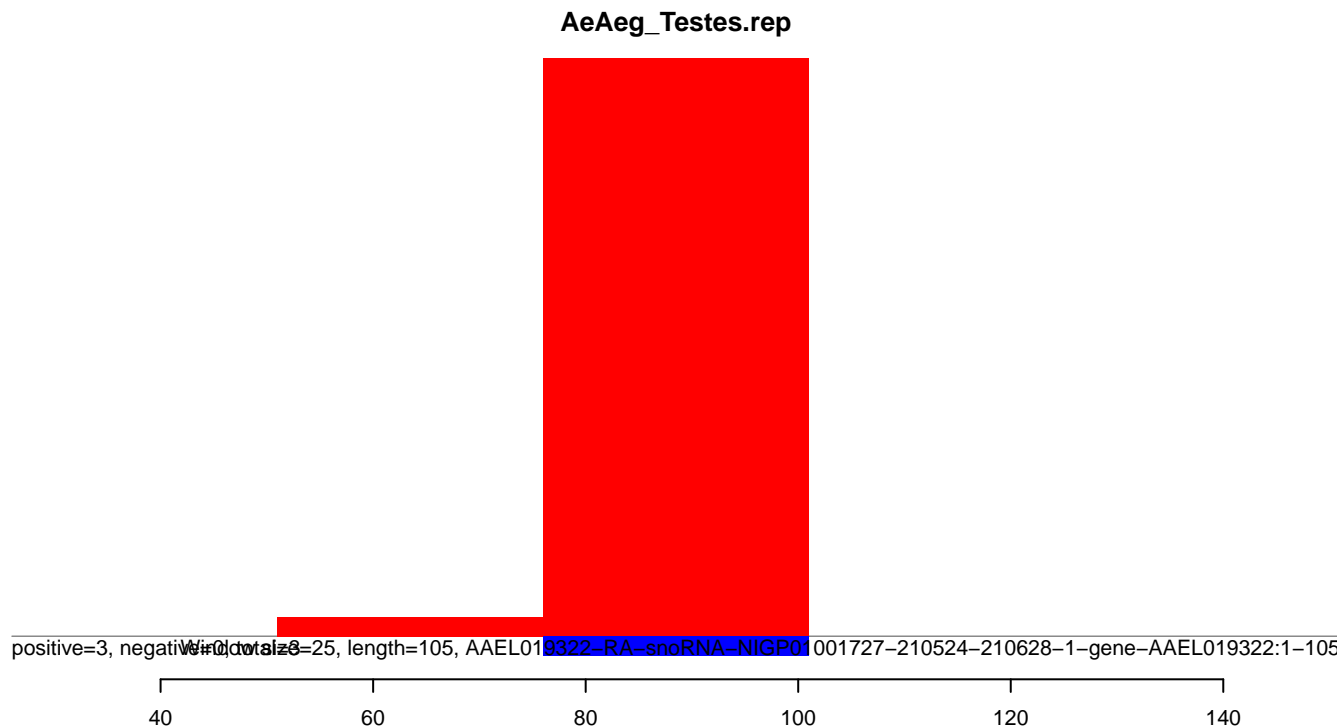
AeAeg_Testes.18_23.rep



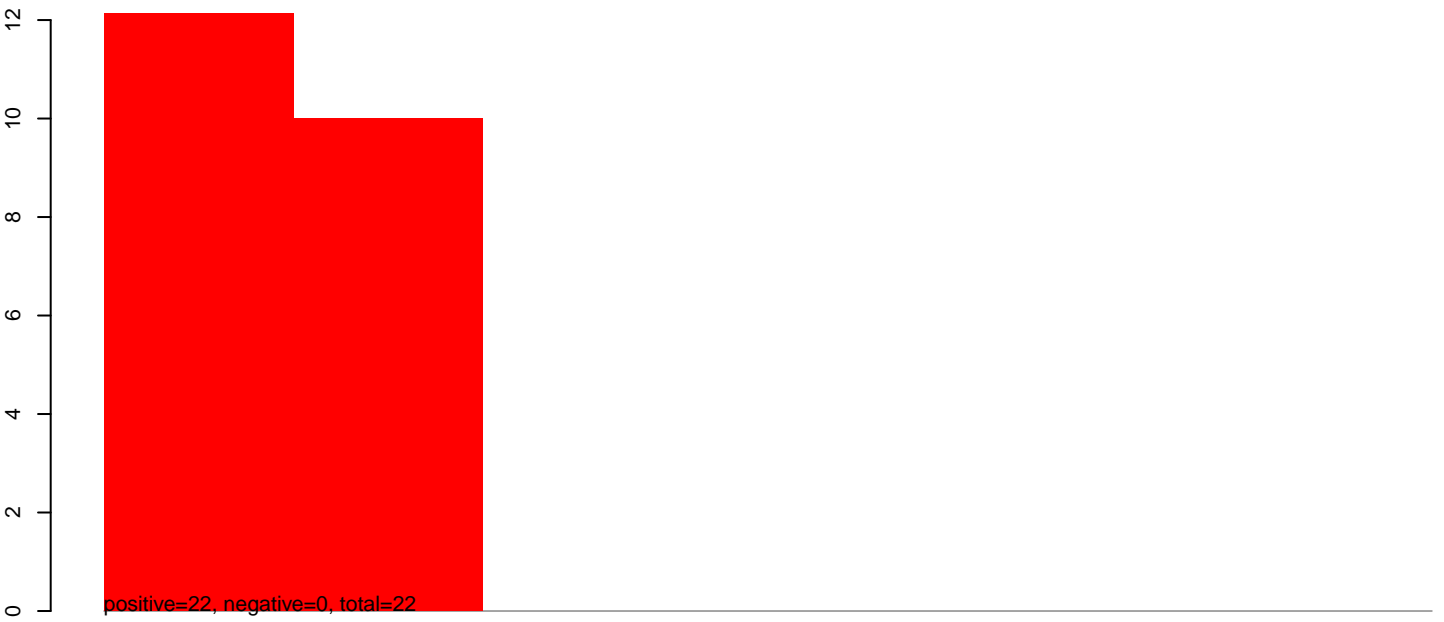
AeAeg_Testes.24_35.rep



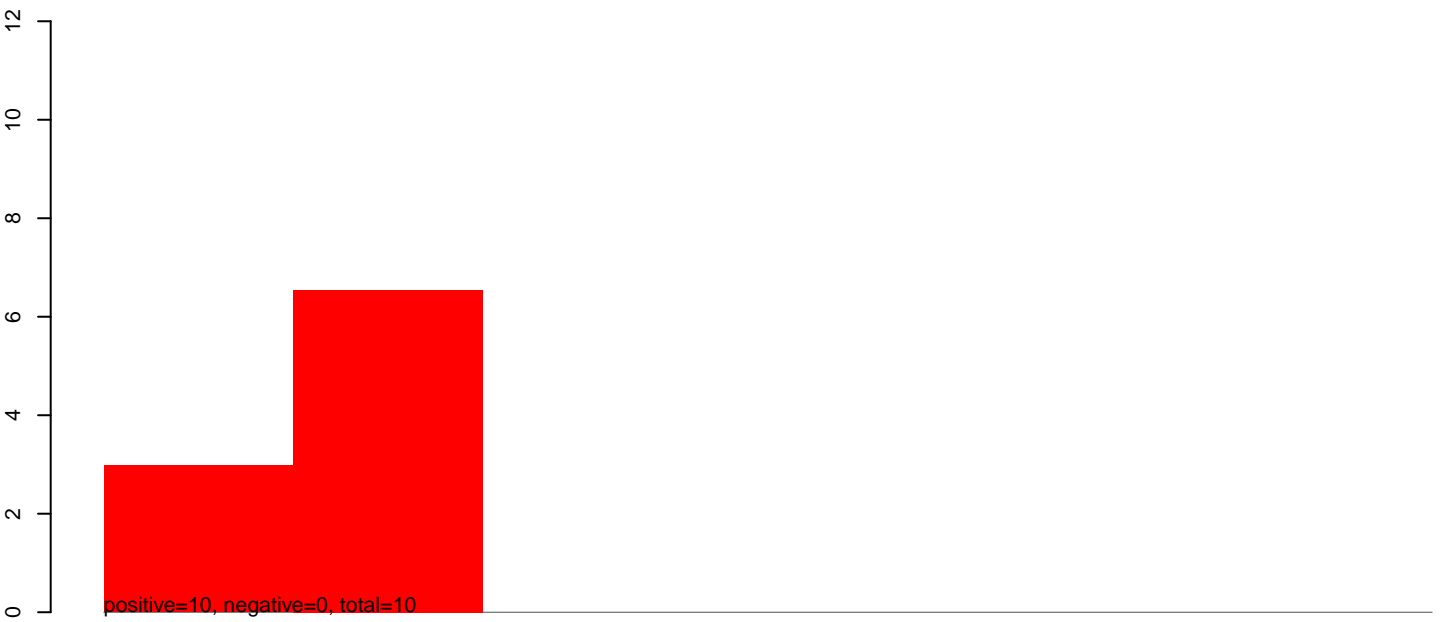
AeAeg_Testes.rep



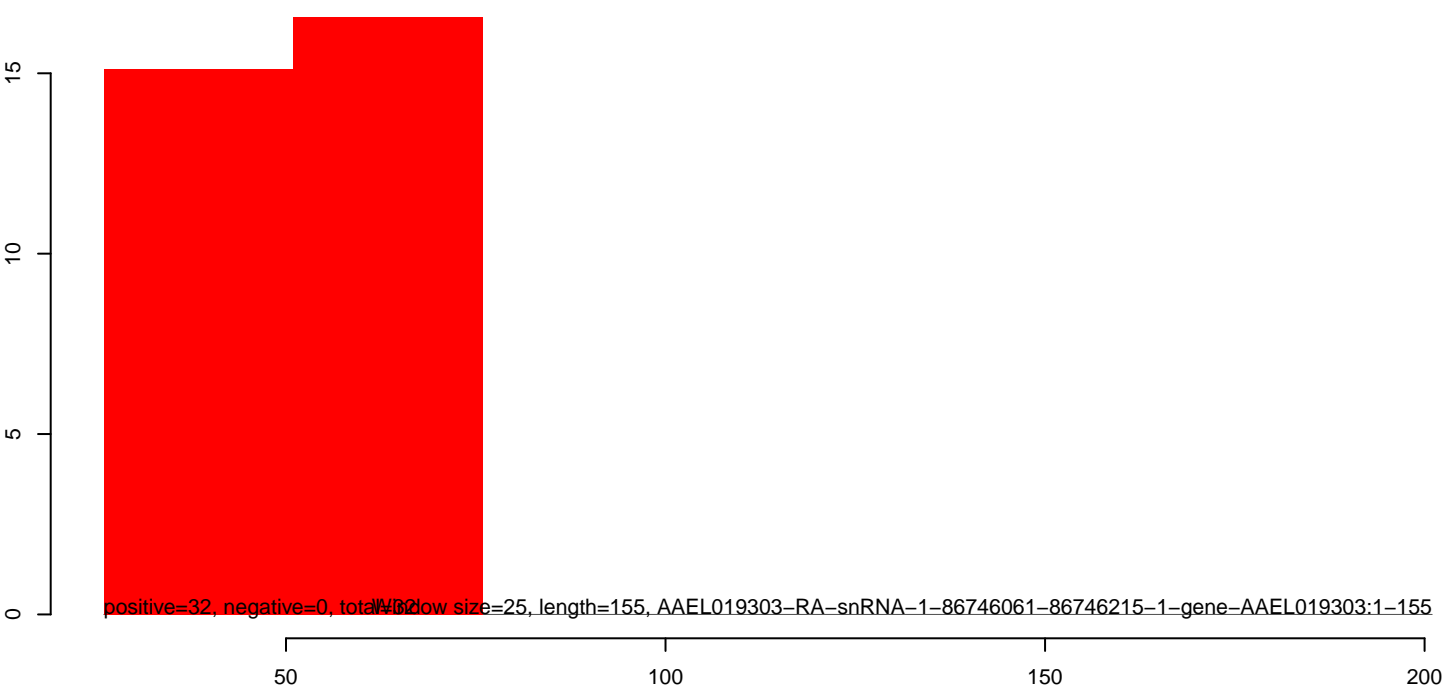
AeAeg_Testes.18_23.rep



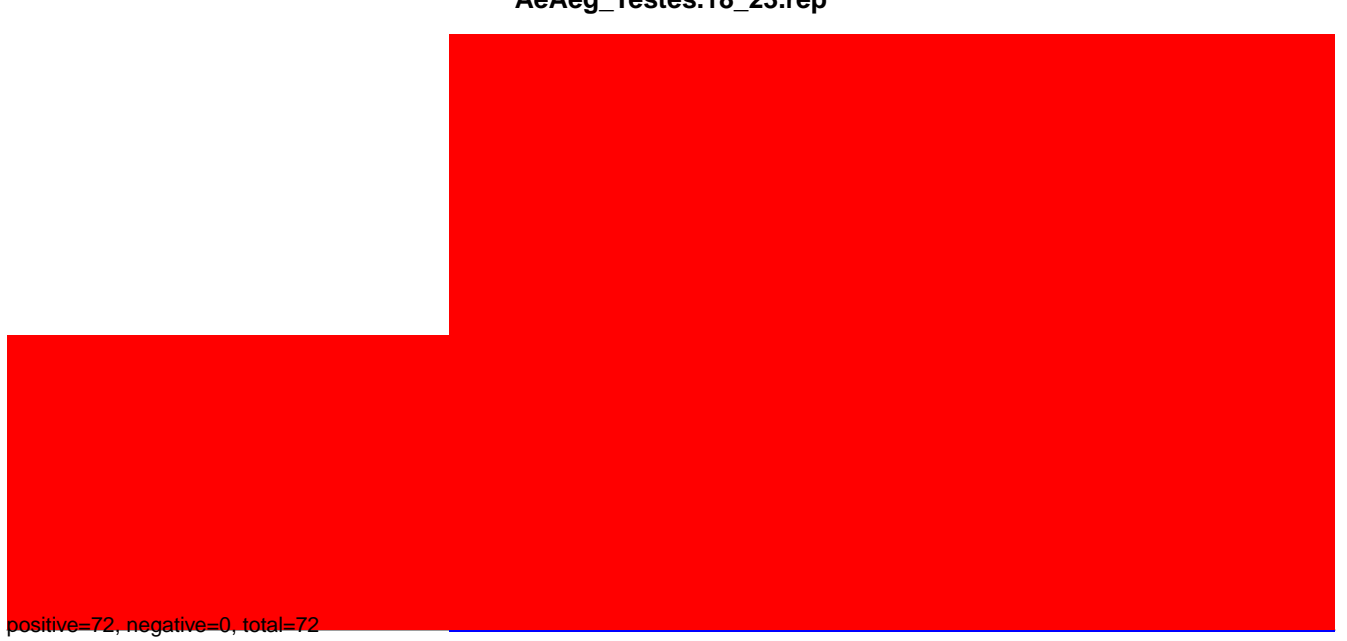
AeAeg_Testes.24_35.rep



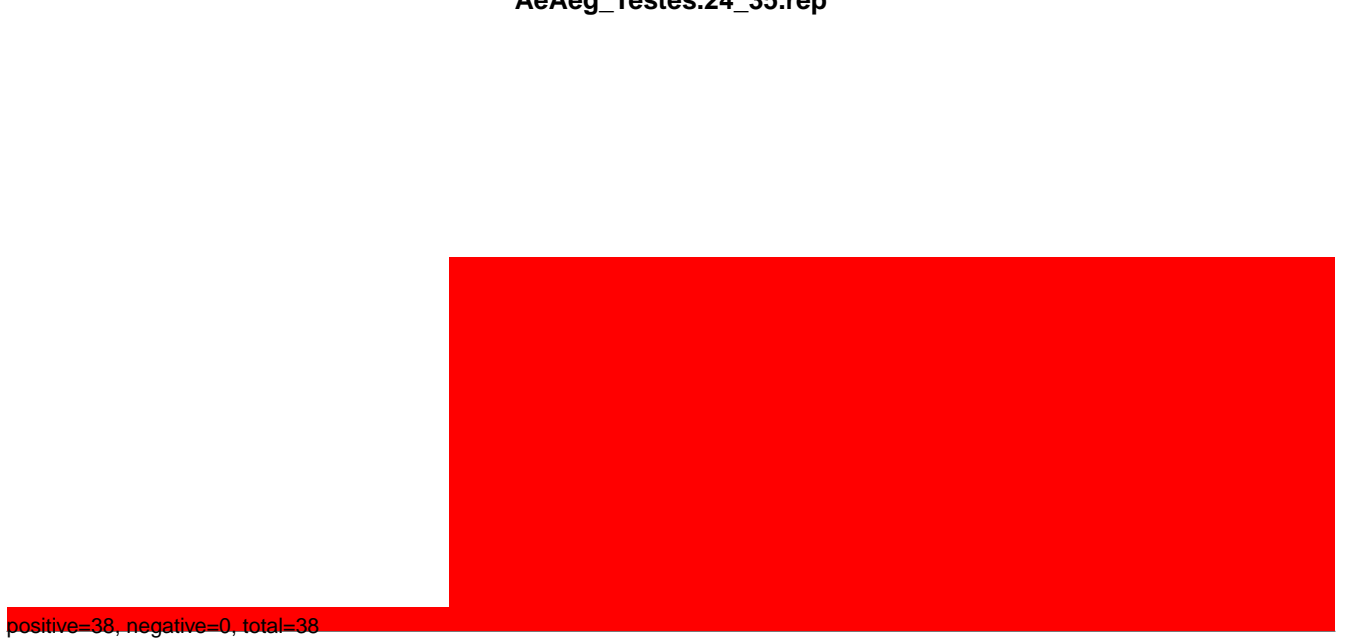
AeAeg_Testes.rep



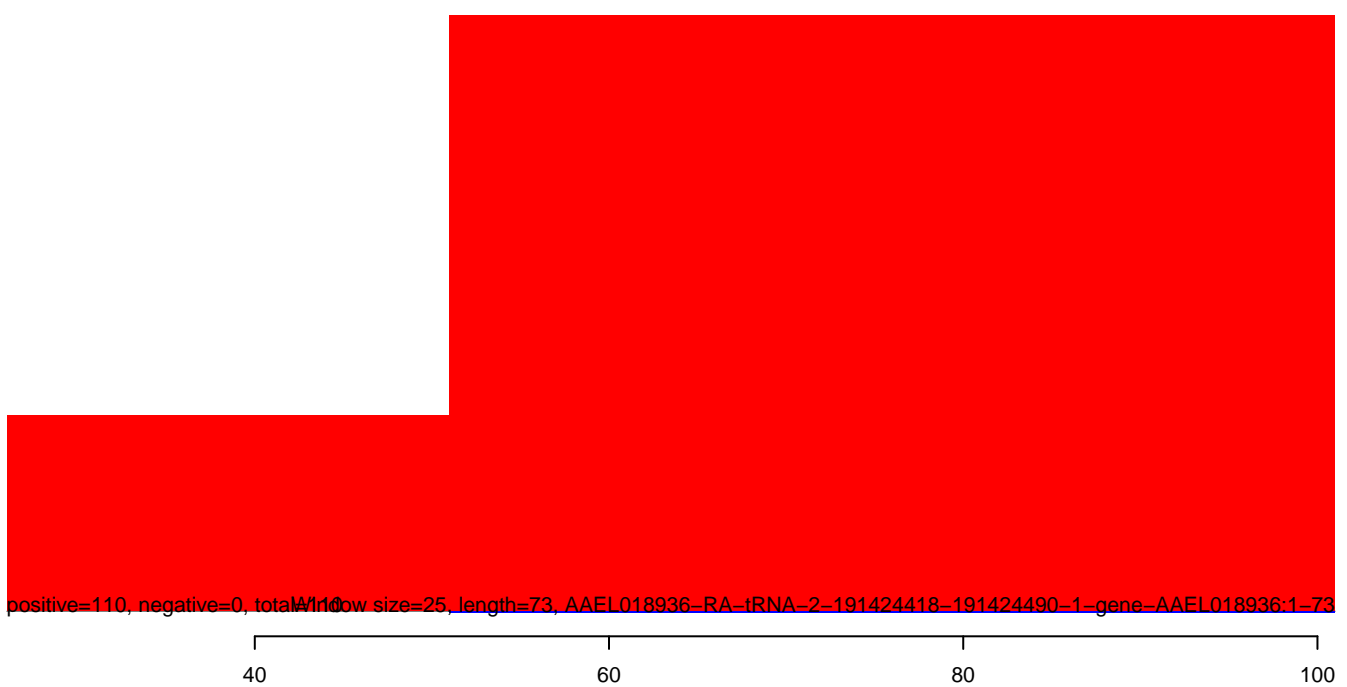
AeAeg_Testes.18_23.rep



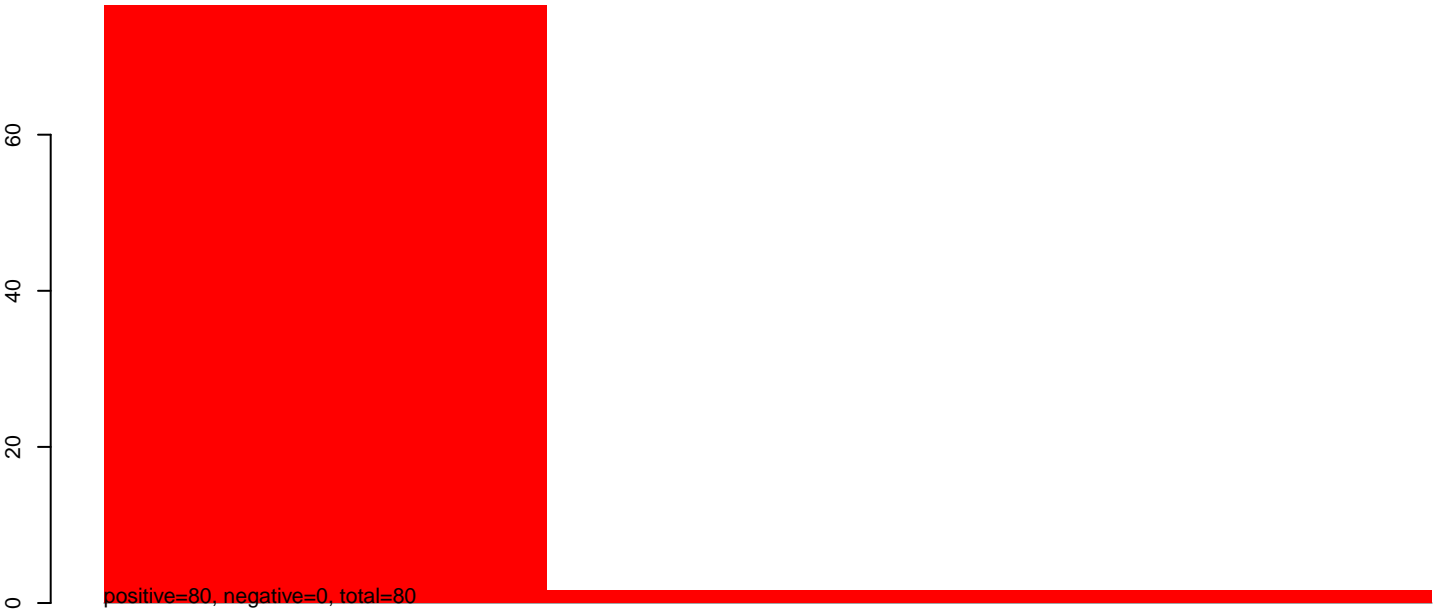
AeAeg_Testes.24_35.rep



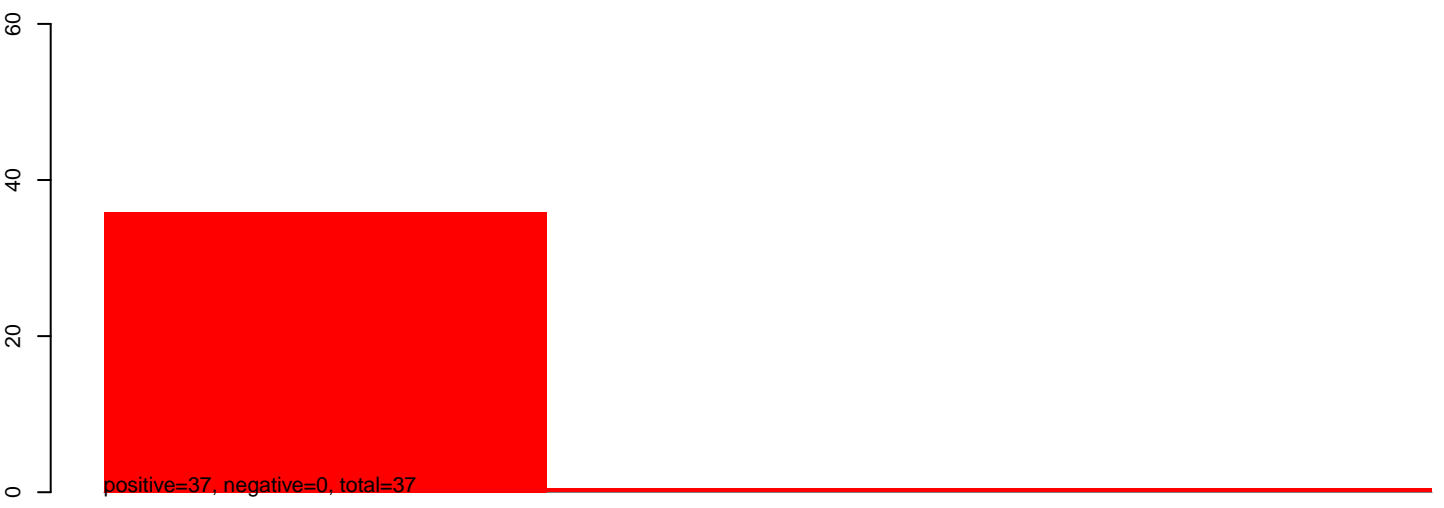
AeAeg_Testes.rep



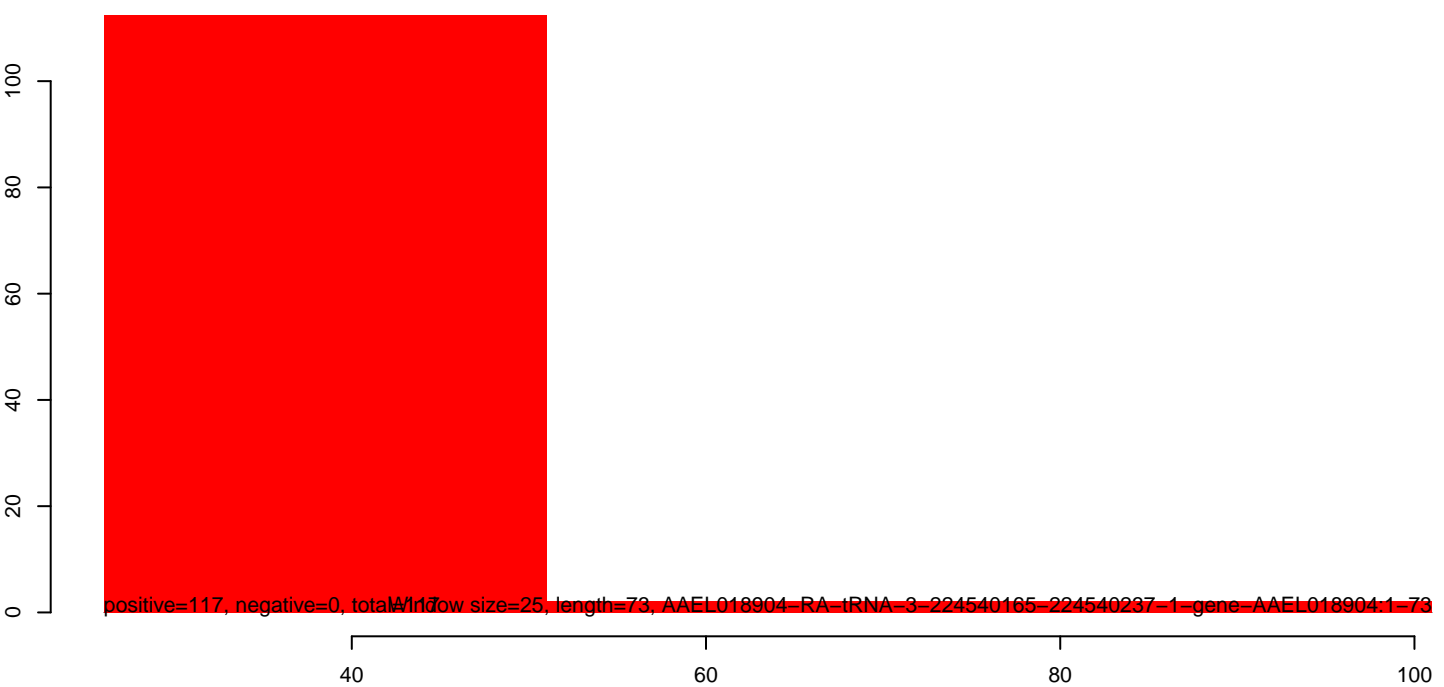
AeAeg_Testes.18_23.rep



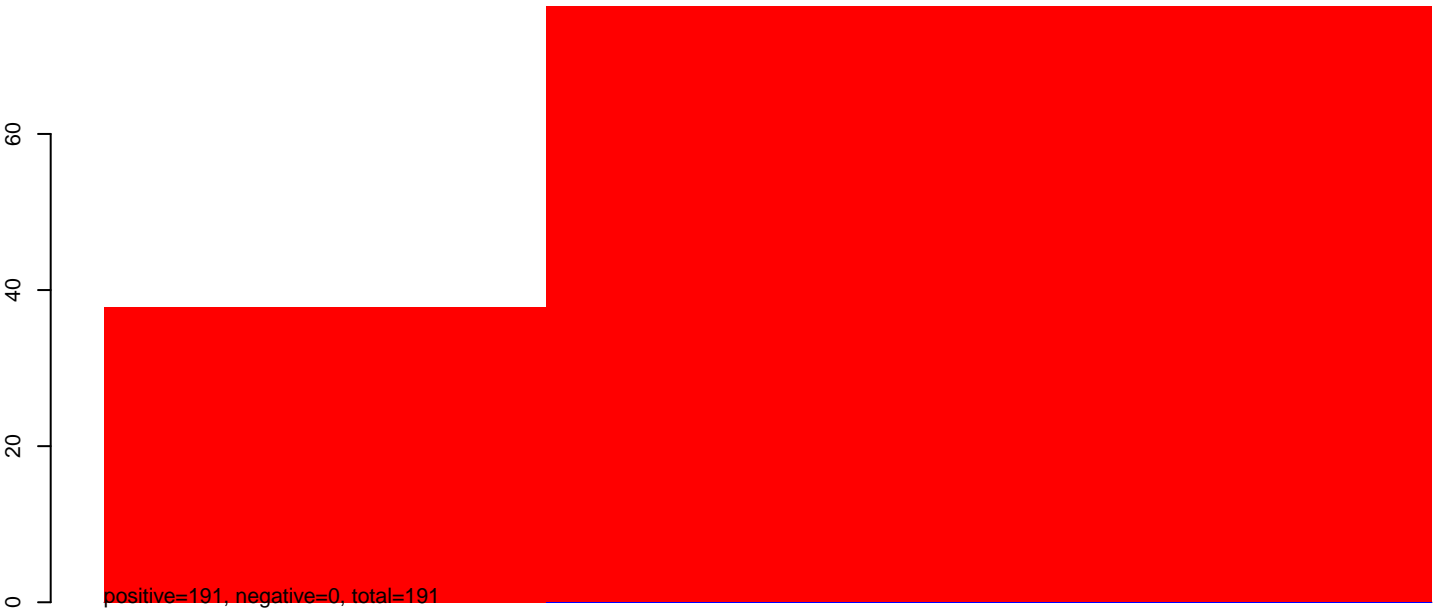
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



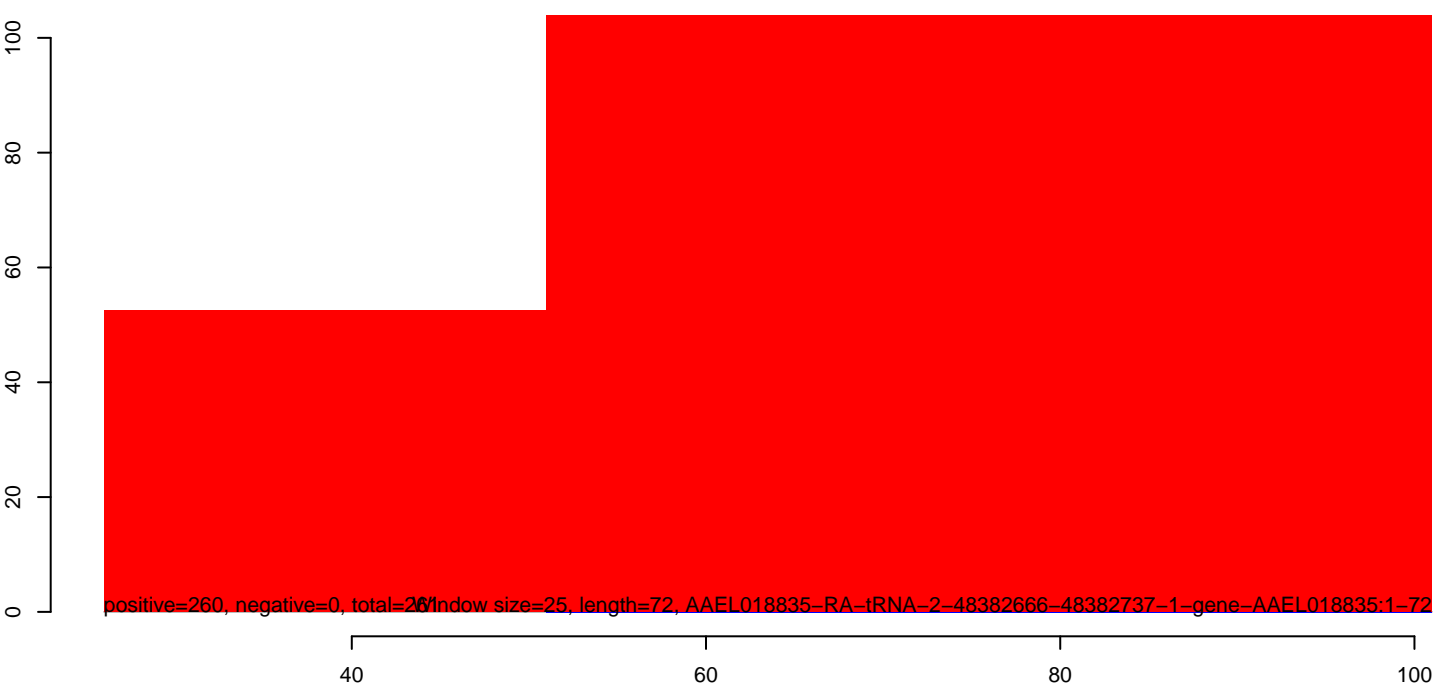
AeAeg_Testes.18_23.rep



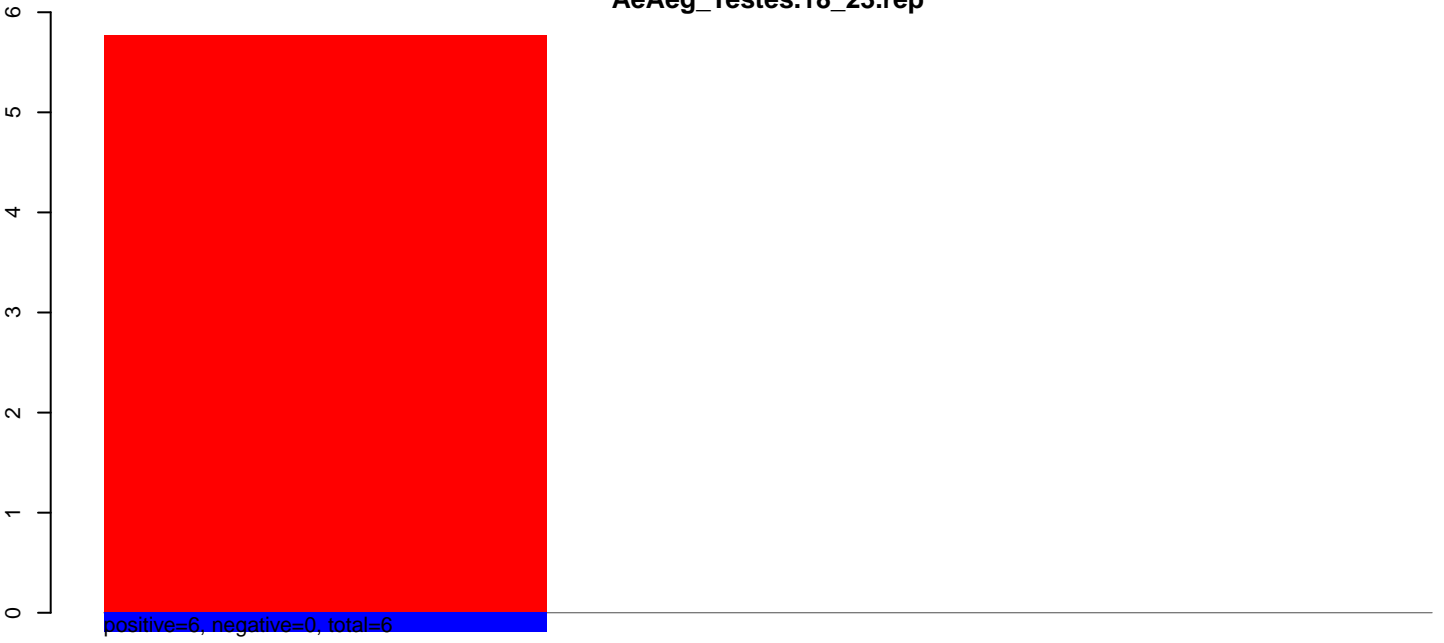
AeAeg_Testes.24_35.rep



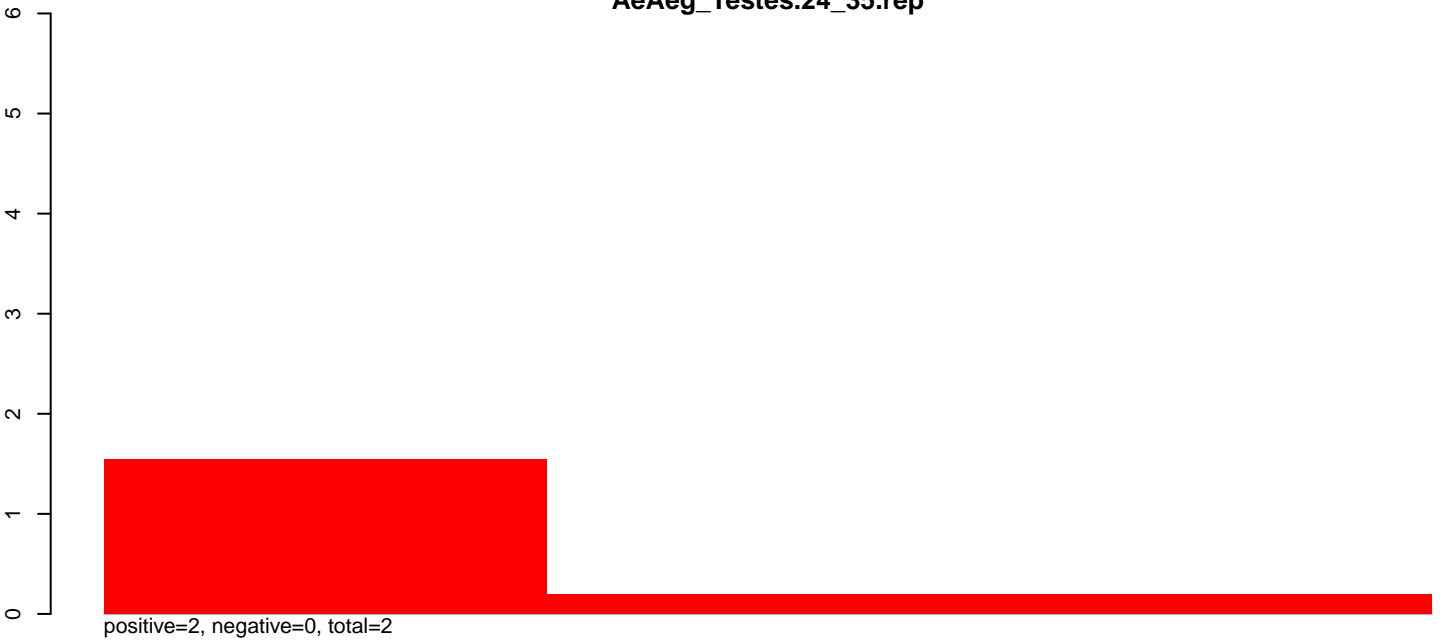
AeAeg_Testes.rep



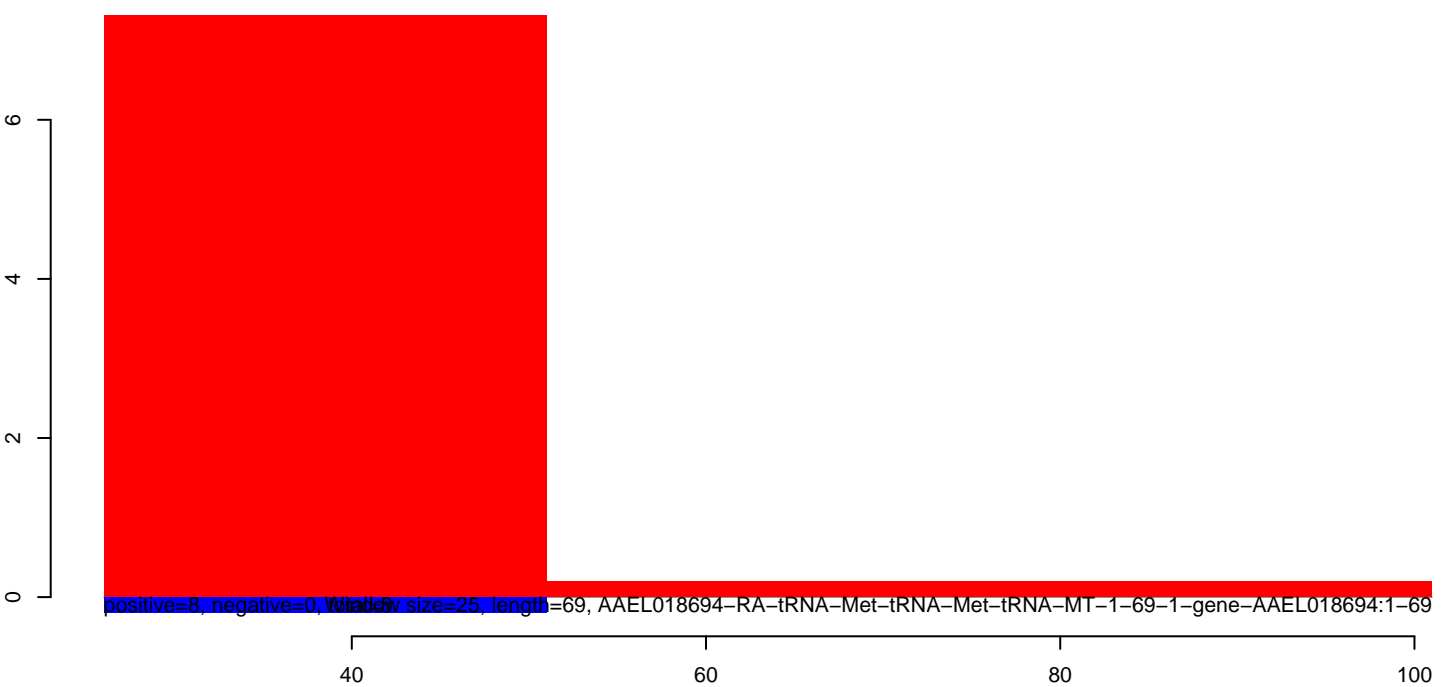
AeAeg_Testes.18_23.rep



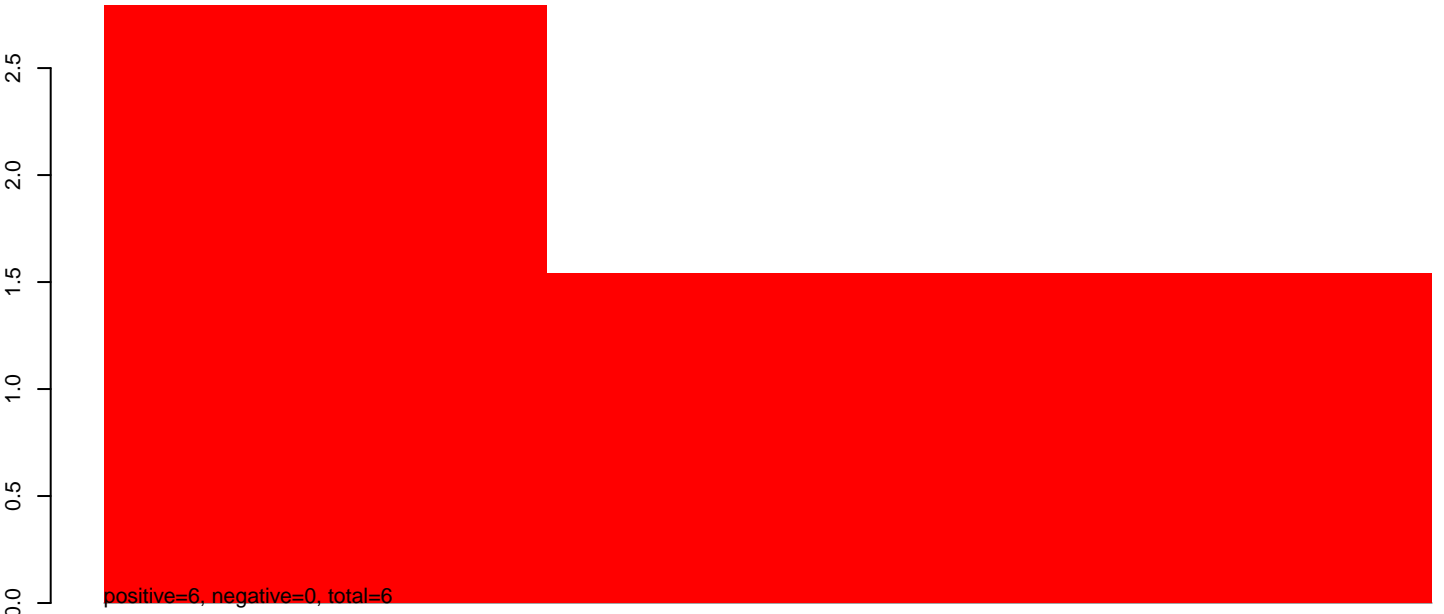
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



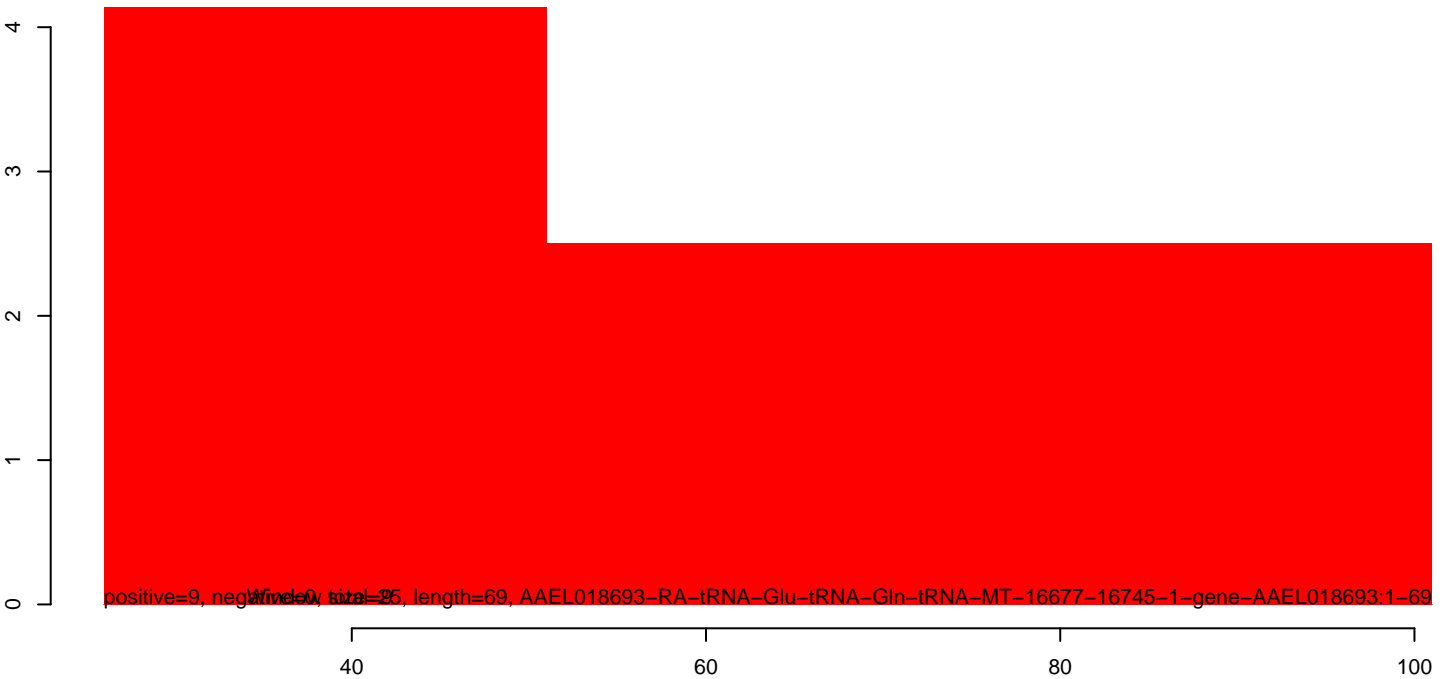
AeAeg_Testes.18_23.rep



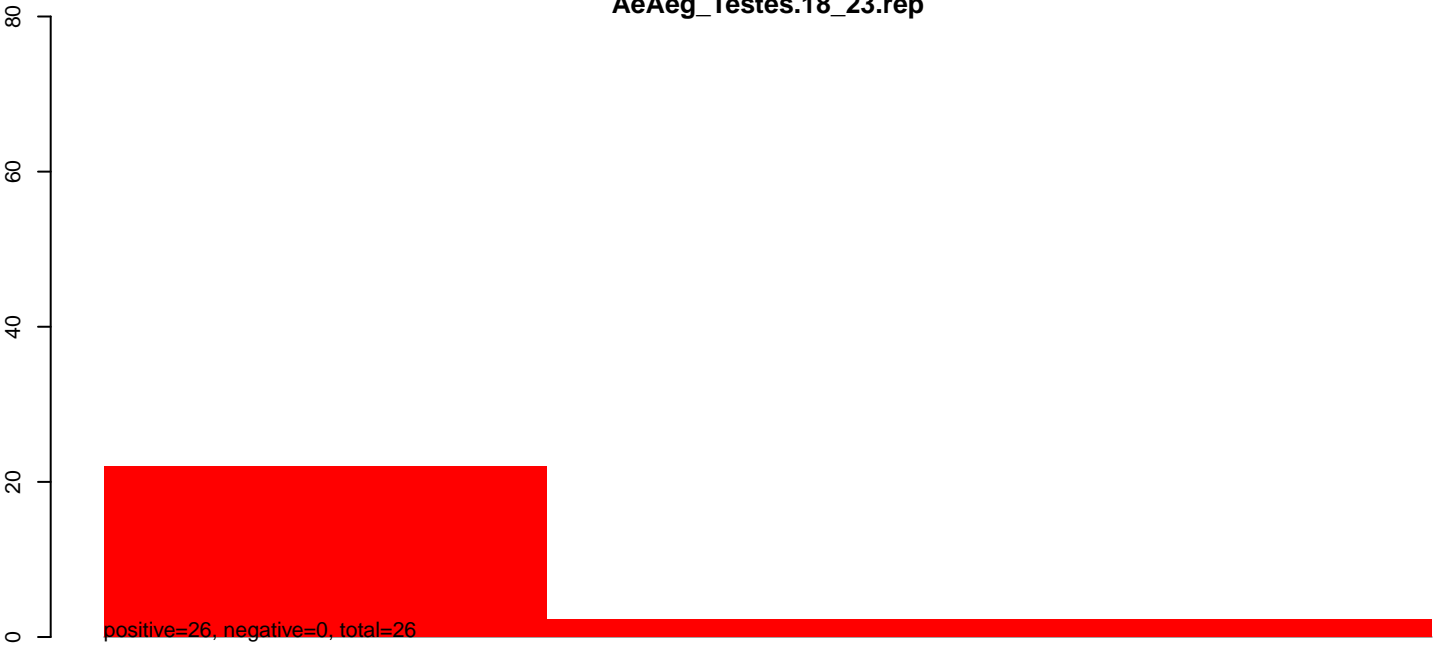
AeAeg_Testes.24_35.rep



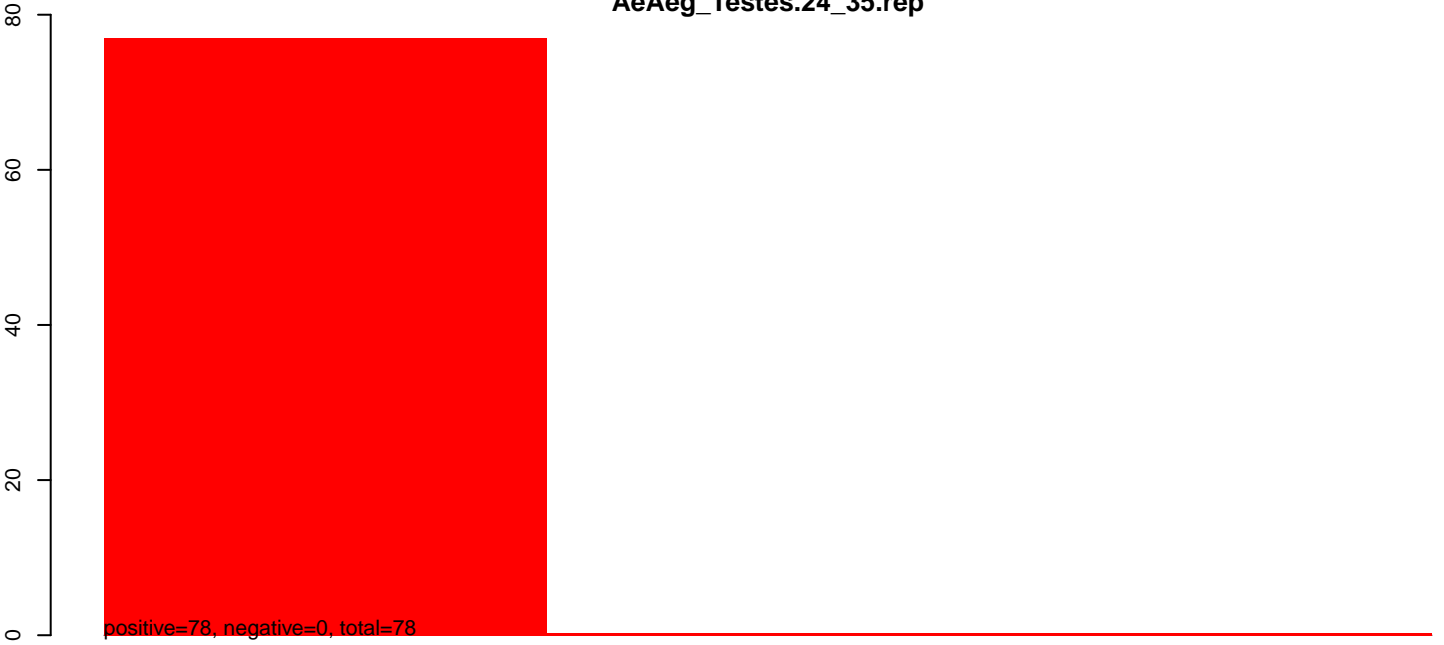
AeAeg_Testes.rep



AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep

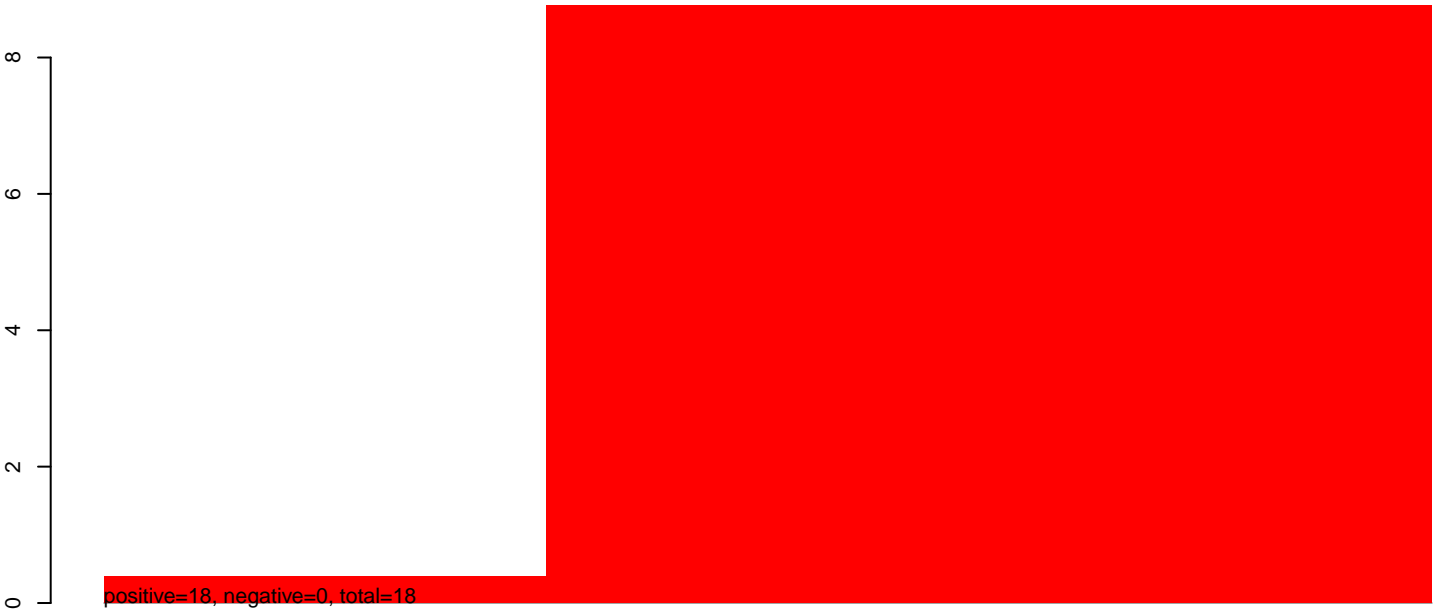


AeAeg_Testes.rep



40 60 80 100
AAEL018692-RA-tRNA-Ile-tRNA-Ile-tRNA-MT-16608-16674-1-gene-AAEL018692:1-67

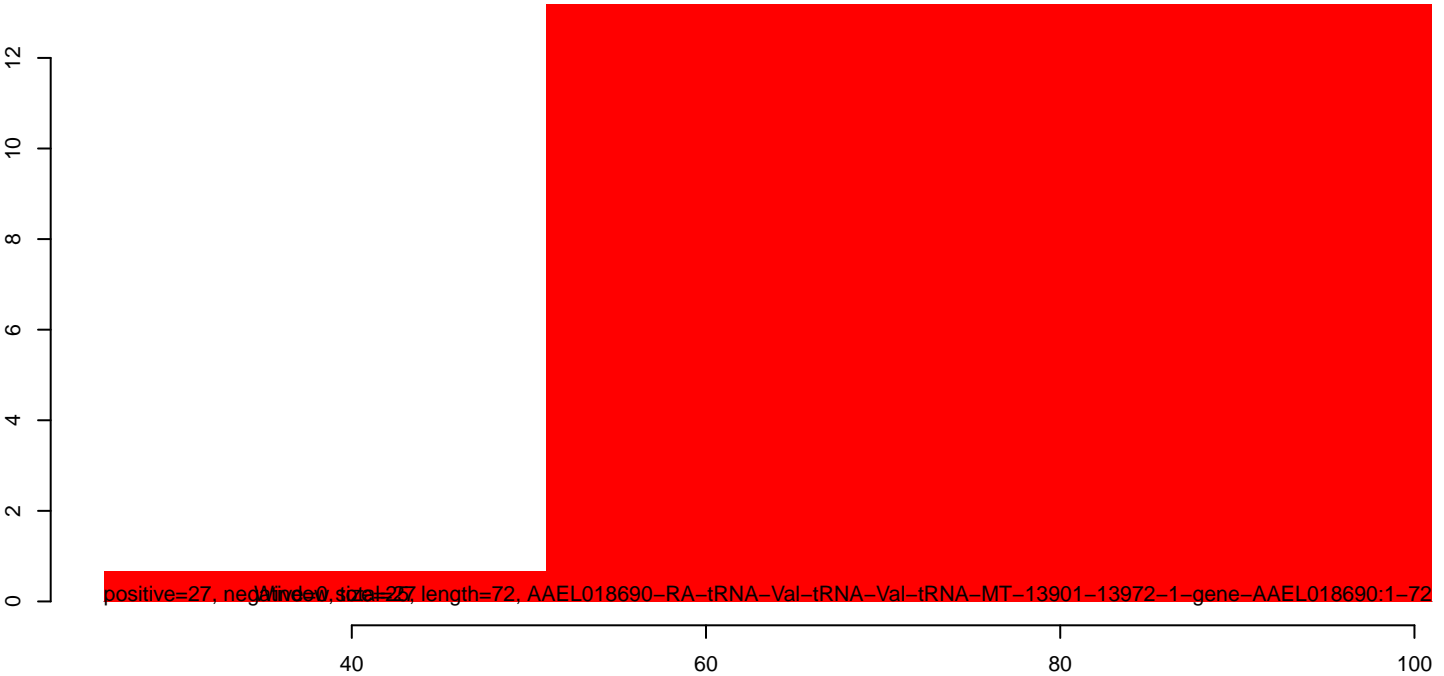
AeAeg_Testes.18_23.rep



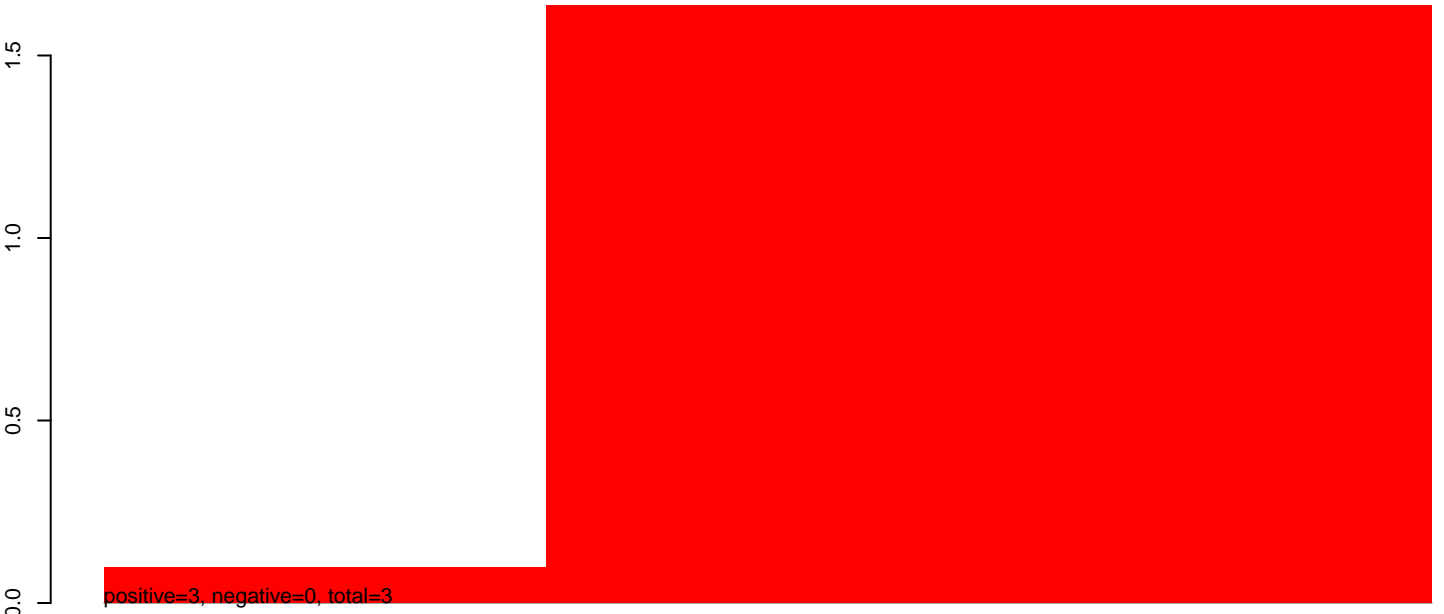
AeAeg_Testes.24_35.rep



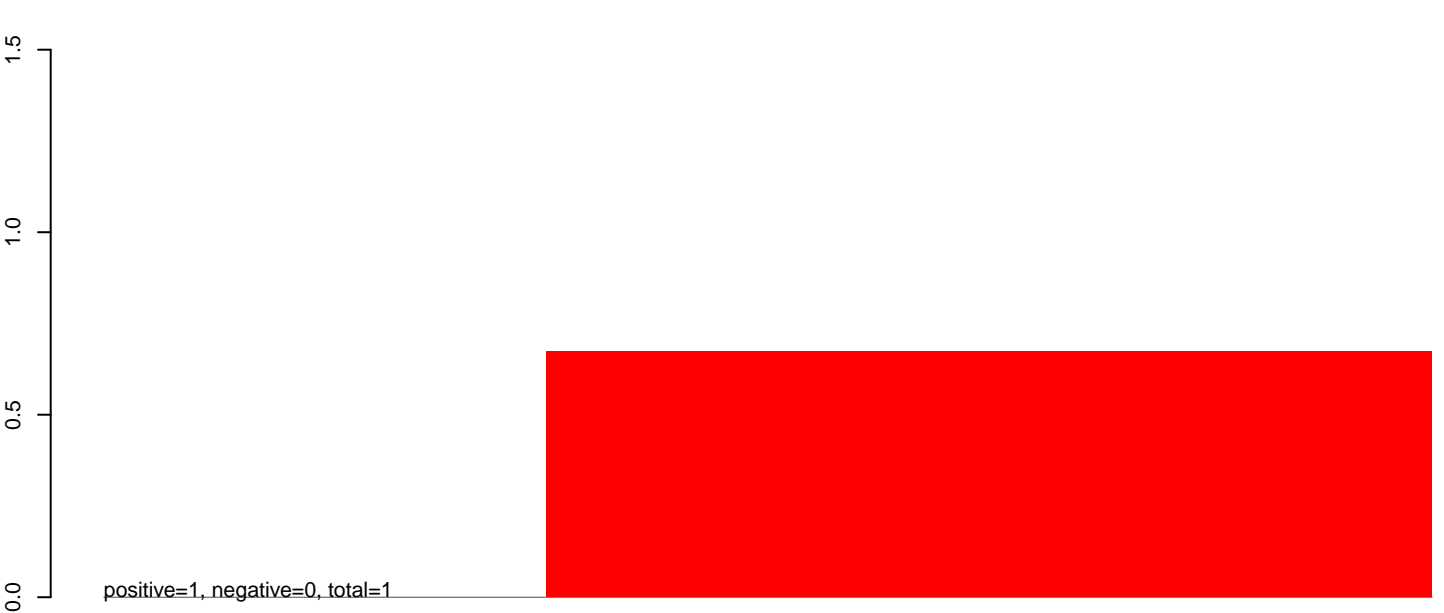
AeAeg_Testes.rep



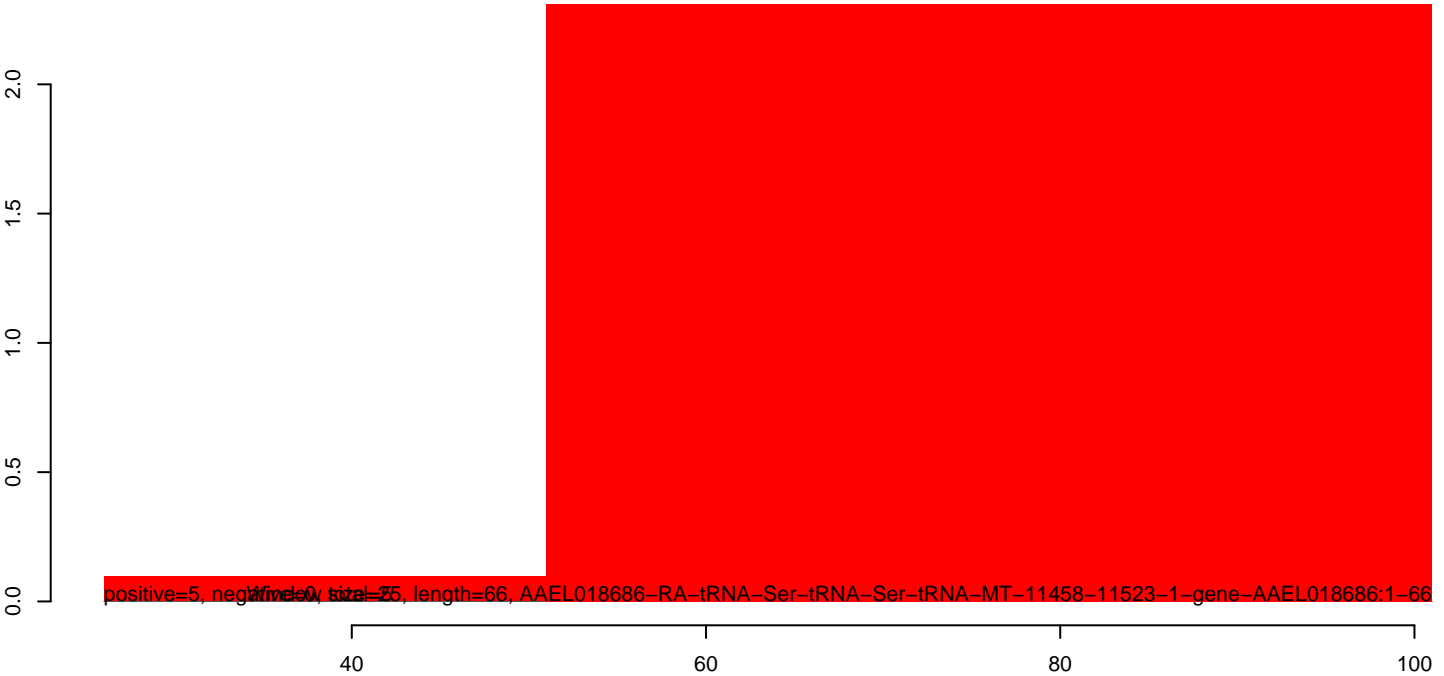
AeAeg_Testes.18_23.rep



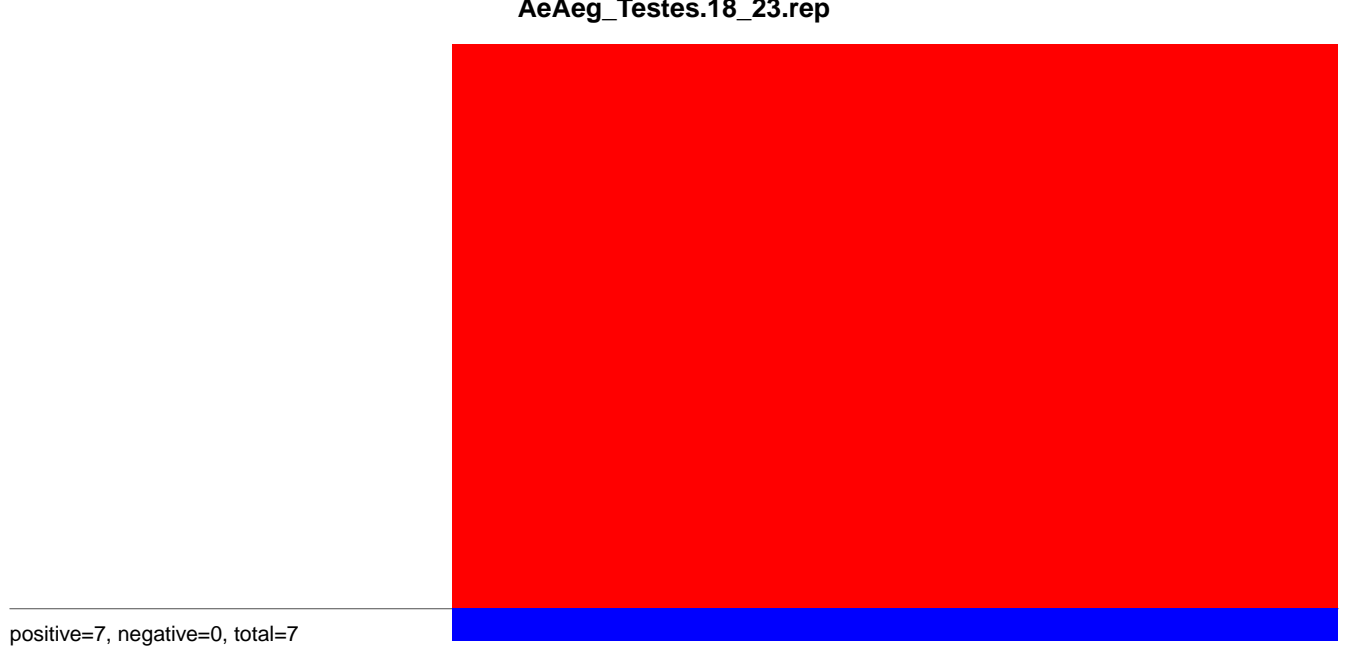
AeAeg_Testes.24_35.rep



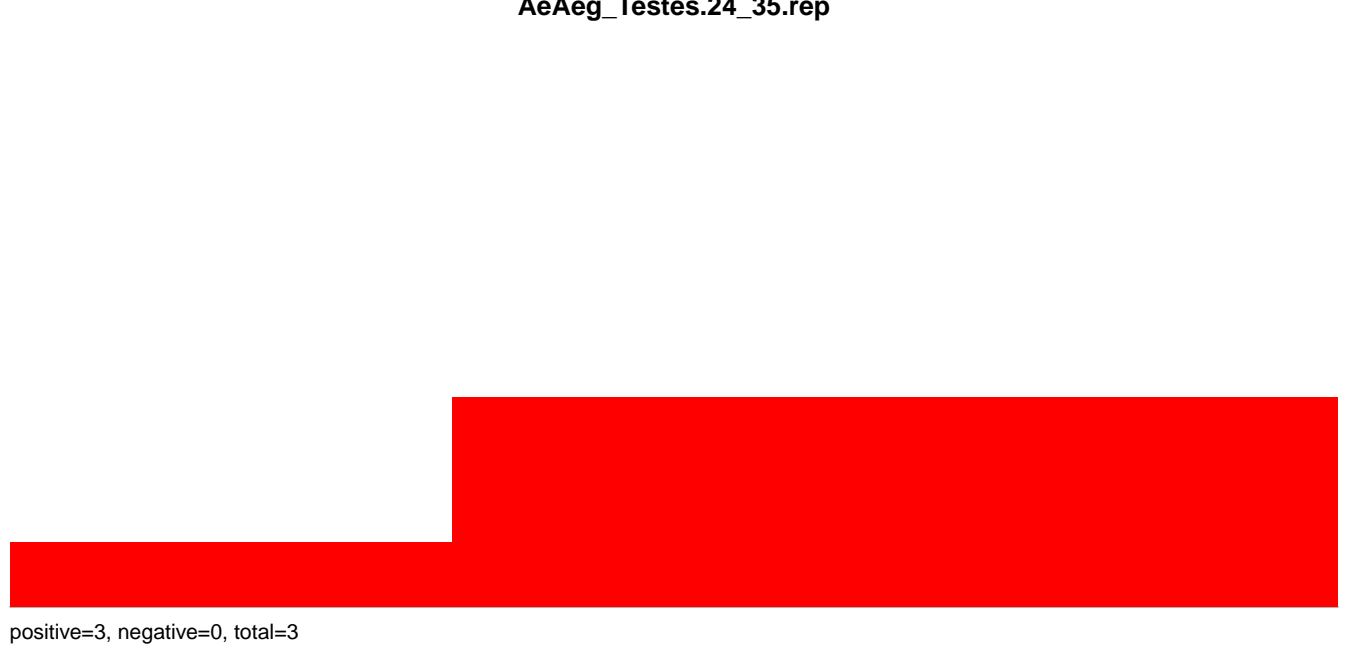
AeAeg_Testes.rep



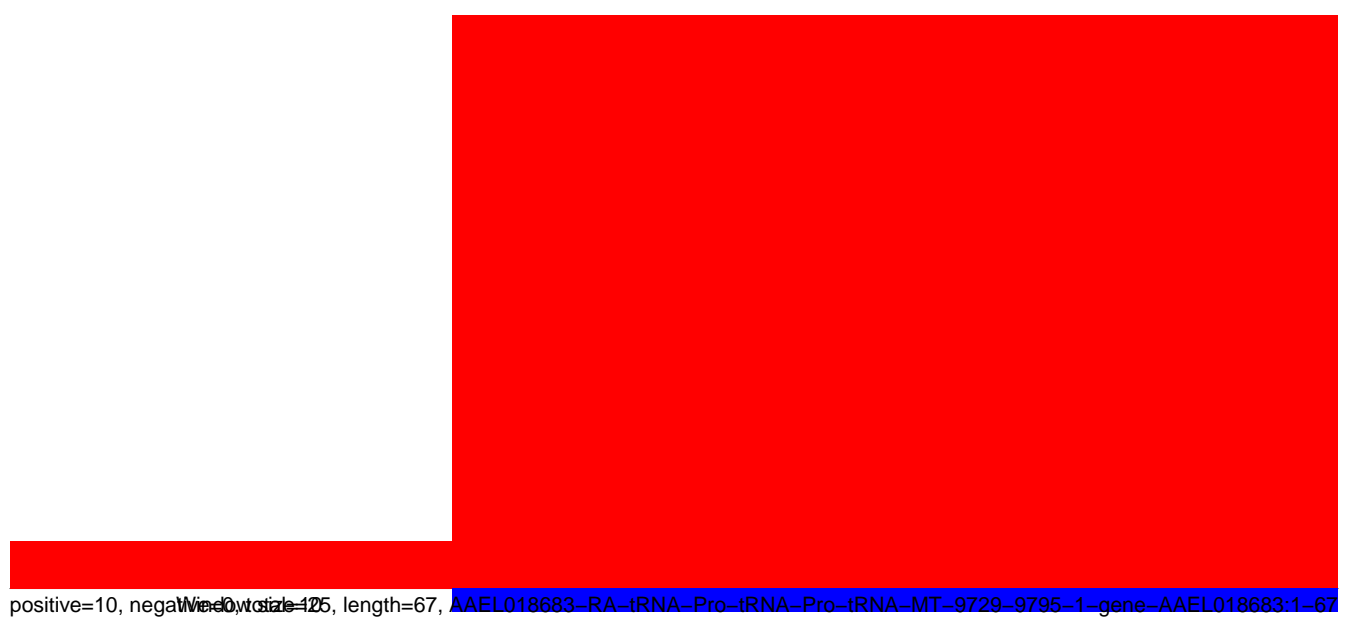
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



AeAeg_Testes.rep

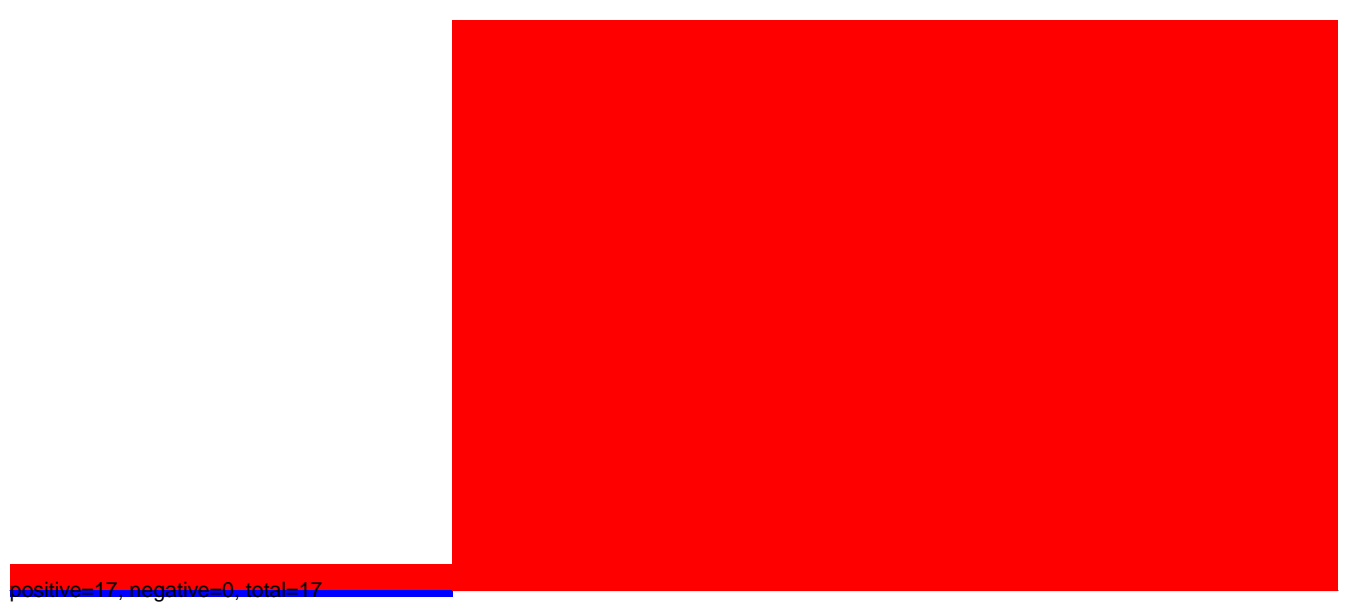


40 60 80 100

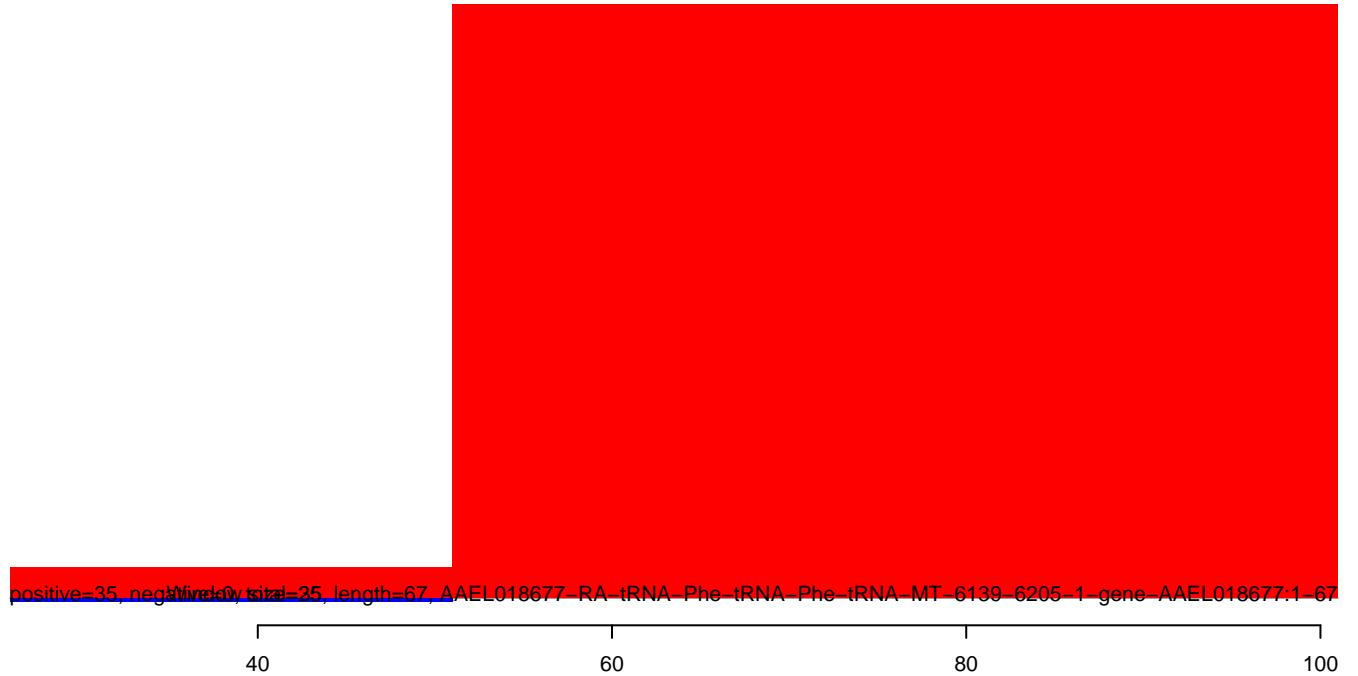
AeAeg_Testes.18_23.rep



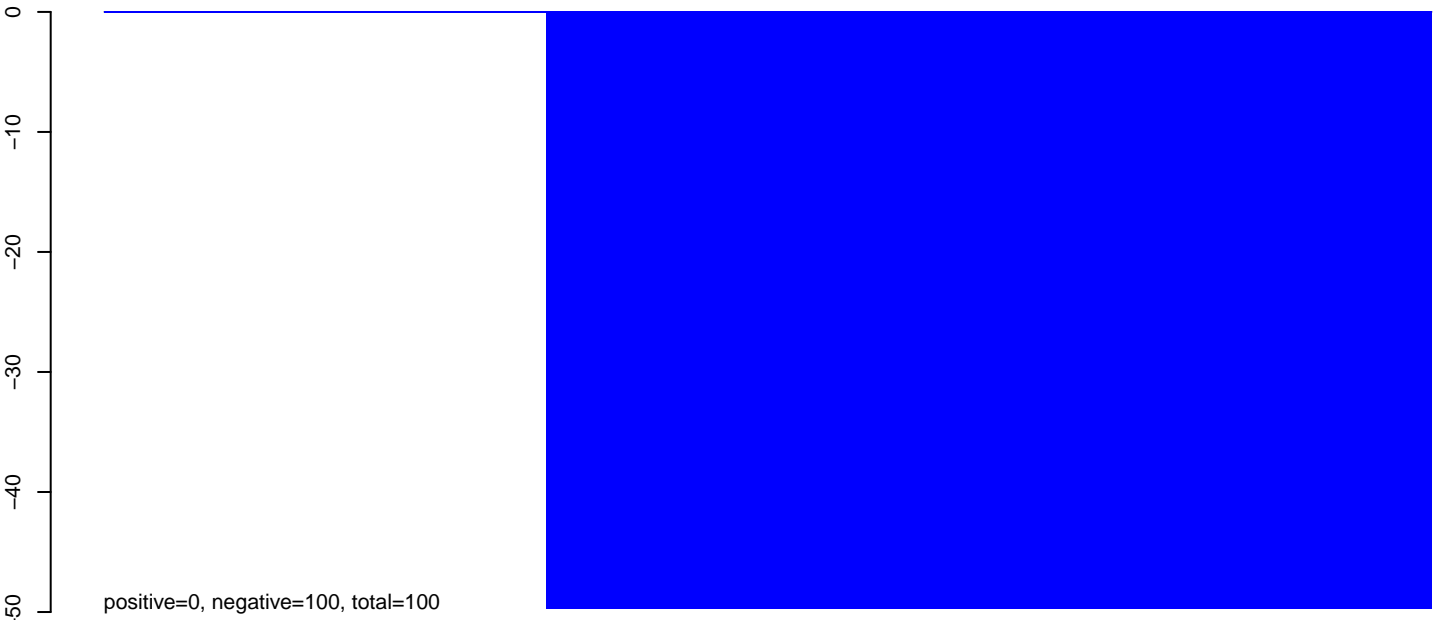
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



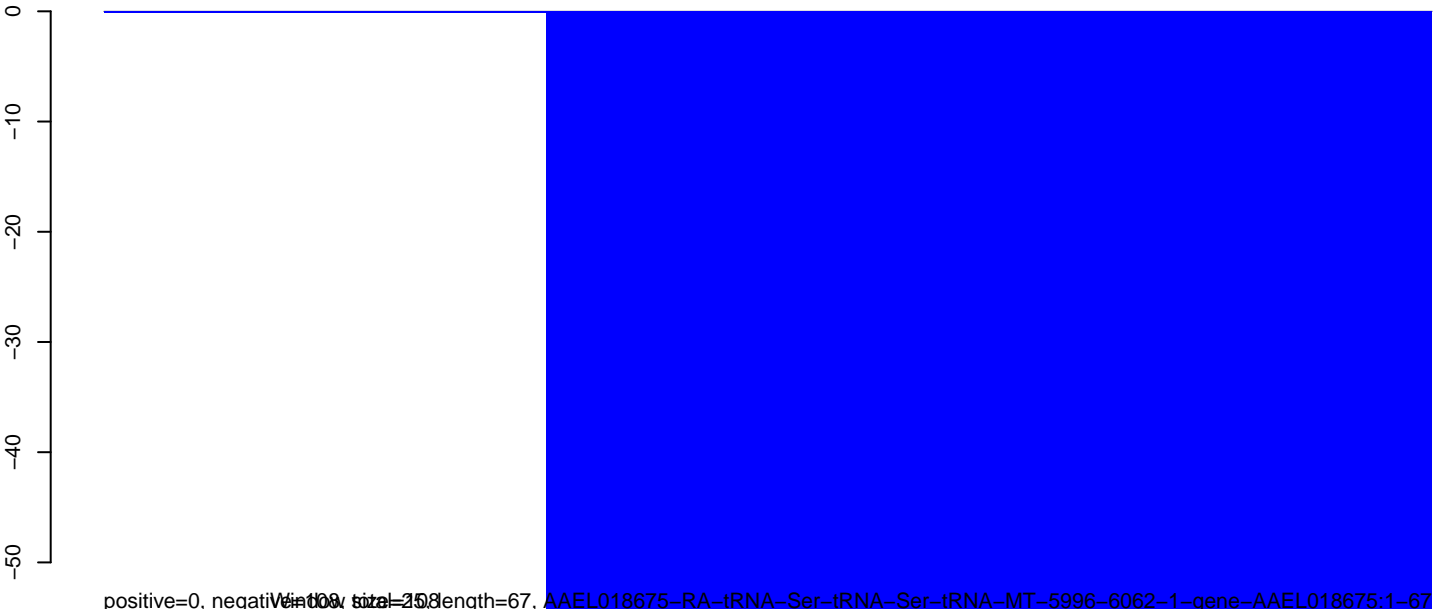
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



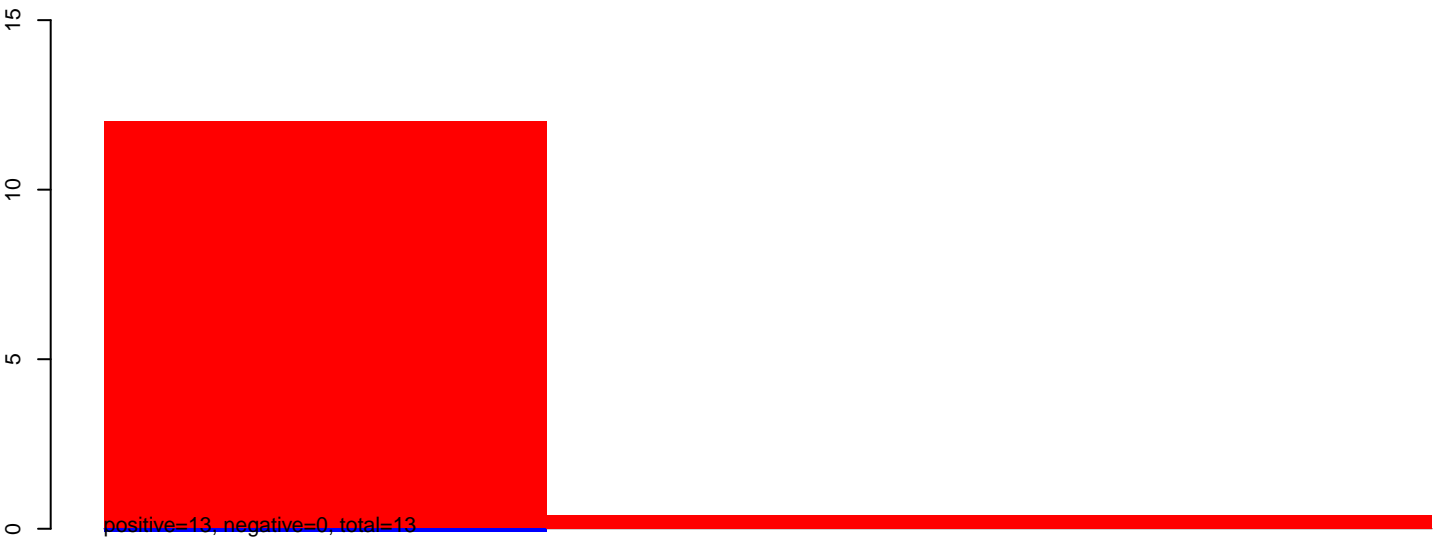
AeAeg_Testes.rep



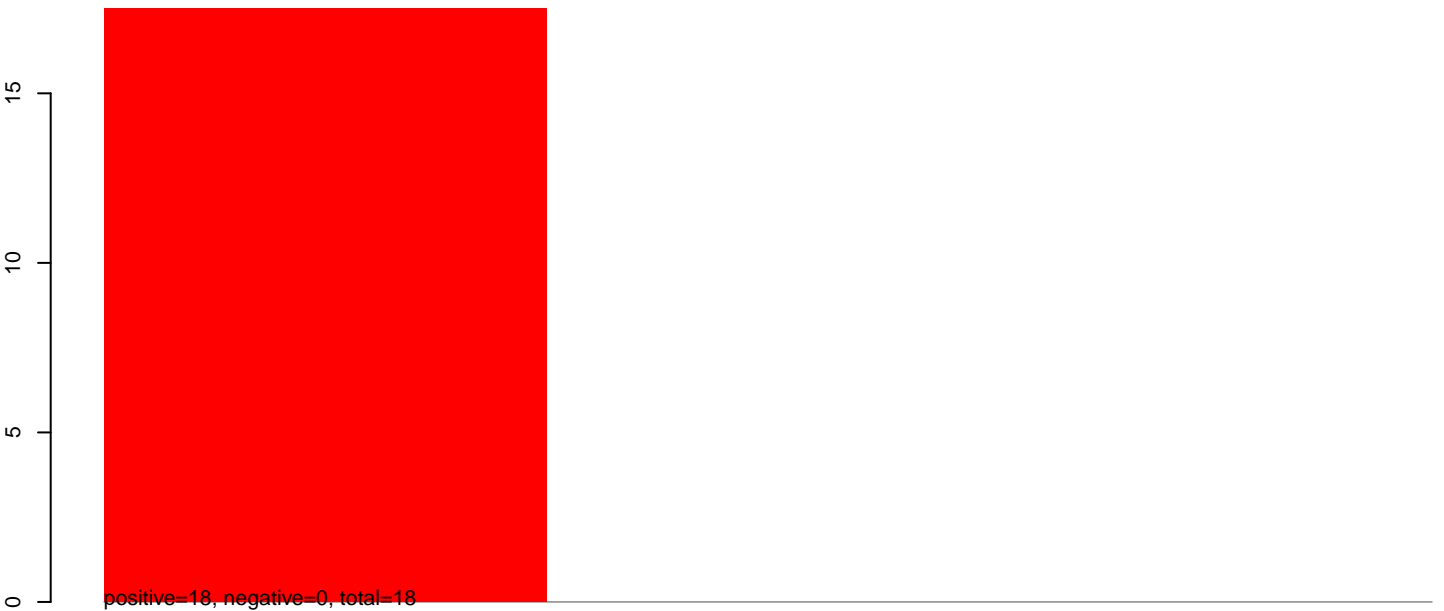
positive=0, negative=100, total=100, length=67, AAEL018675-RA-tRNA-Ser-tRNA-Ser-tRNA-MT-5996-6062-1-gene-AAEL018675:1-67

40 60 80 100

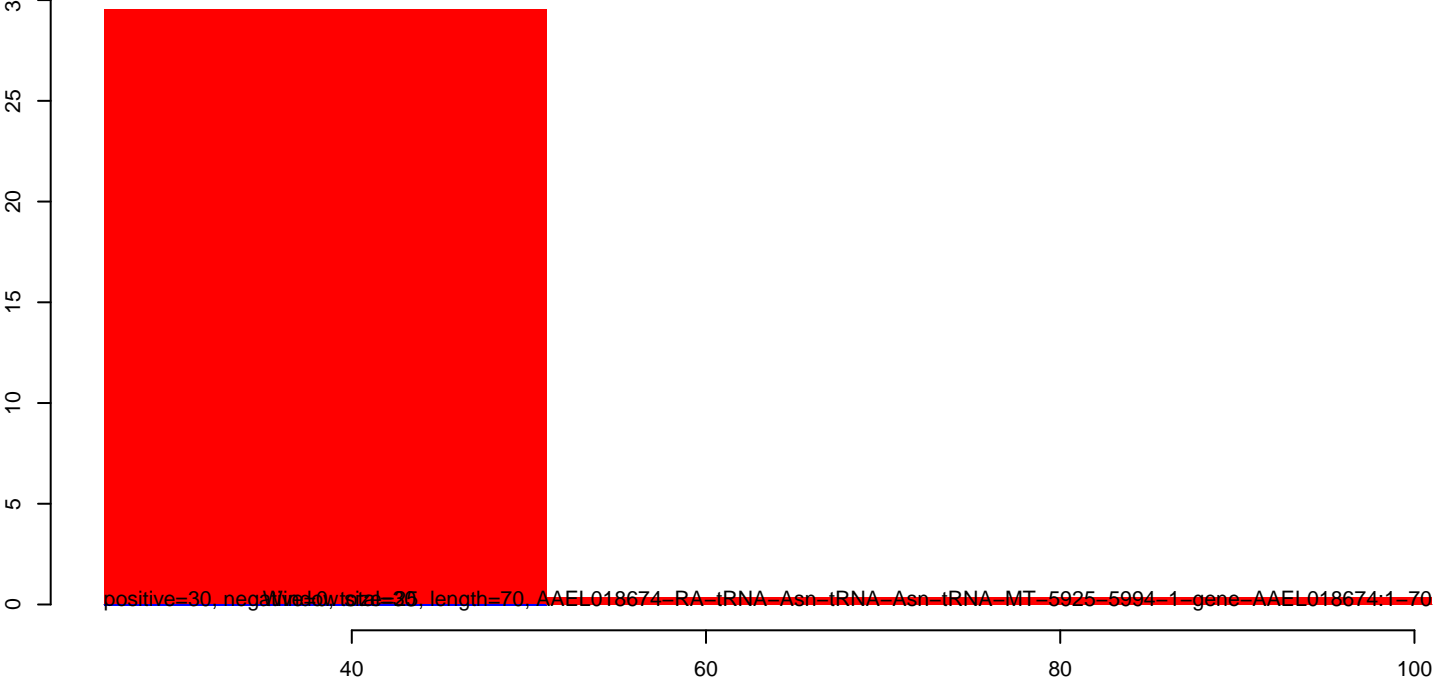
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



AeAeg_Testes.rep

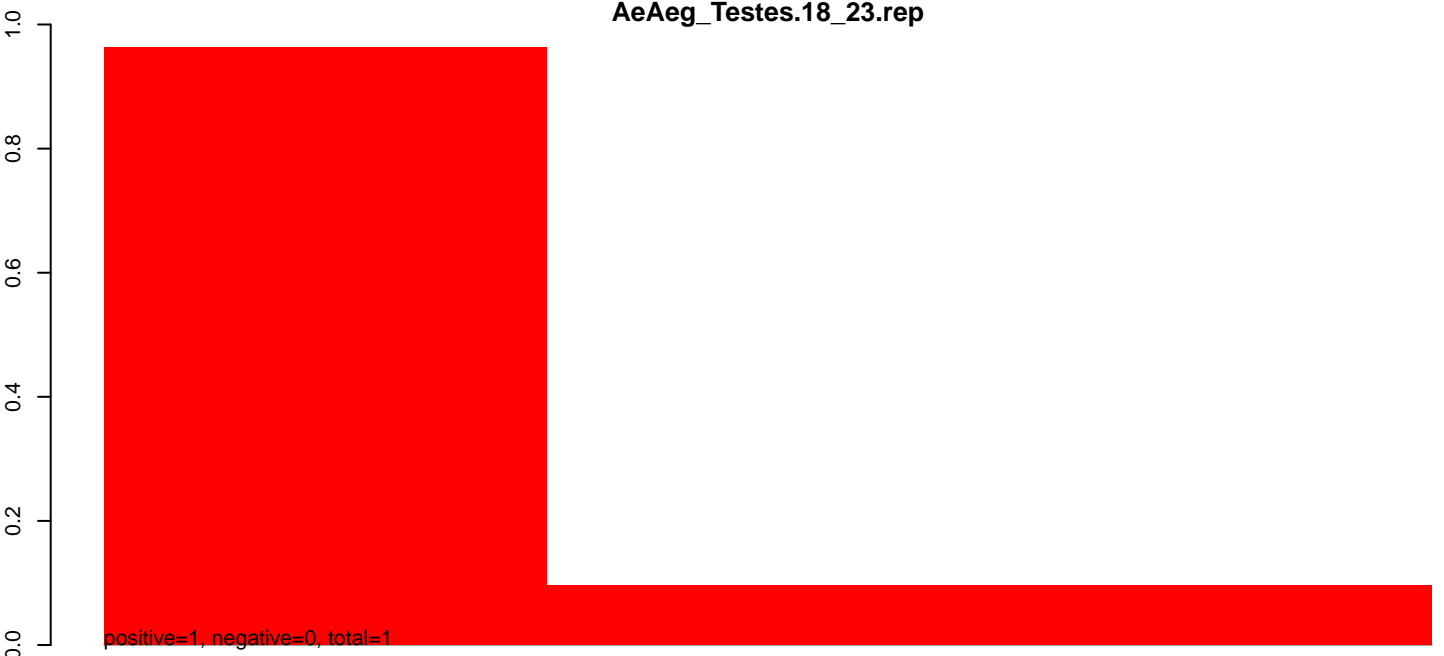


AAEL018674-RA-IRNA-Asn-IRNA-Asn-IRNA-MT-5925-5994-1-gene-AAEL018674:1-70

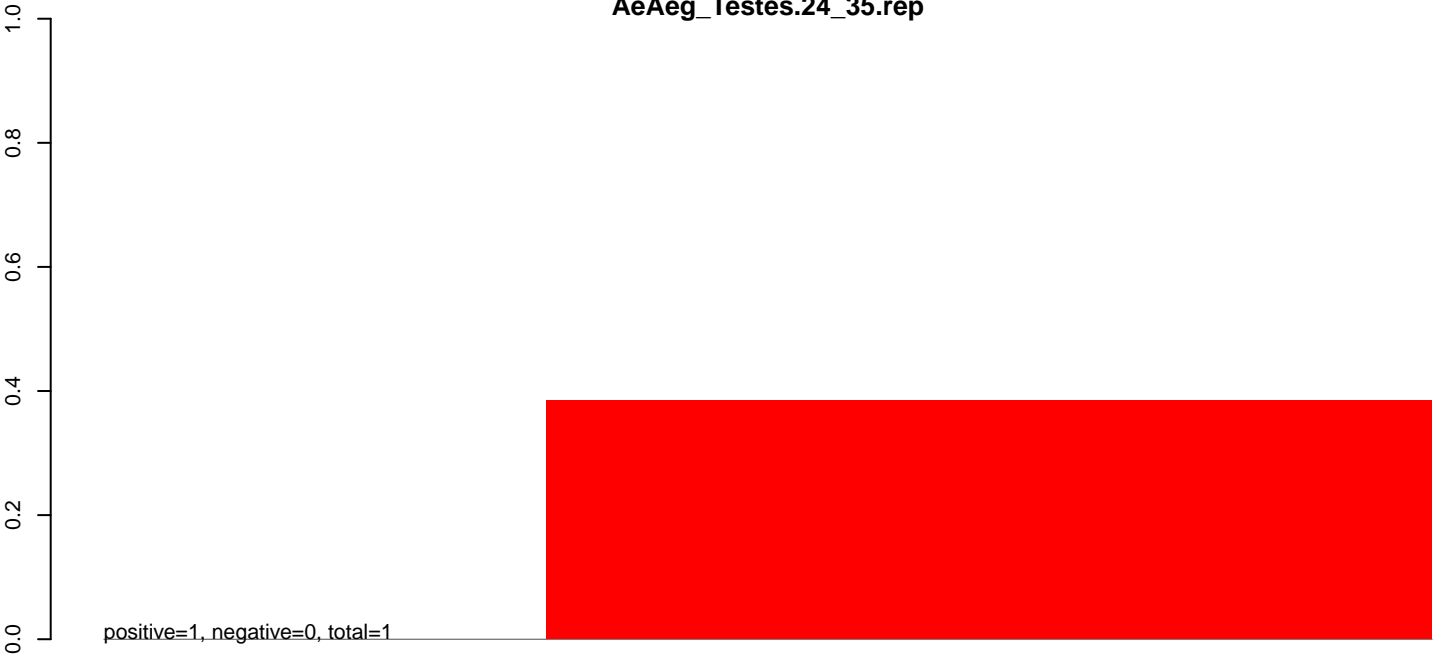
Window size=25, length=70

40 60 80 100

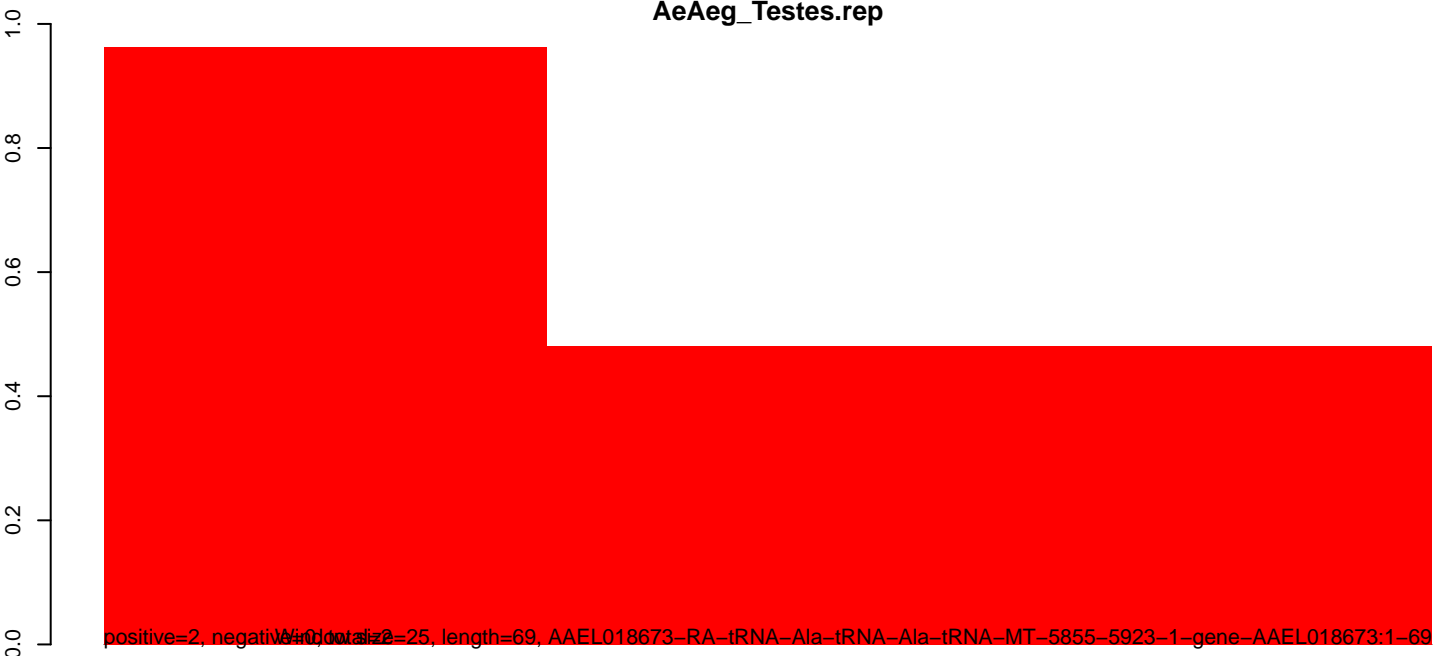
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep

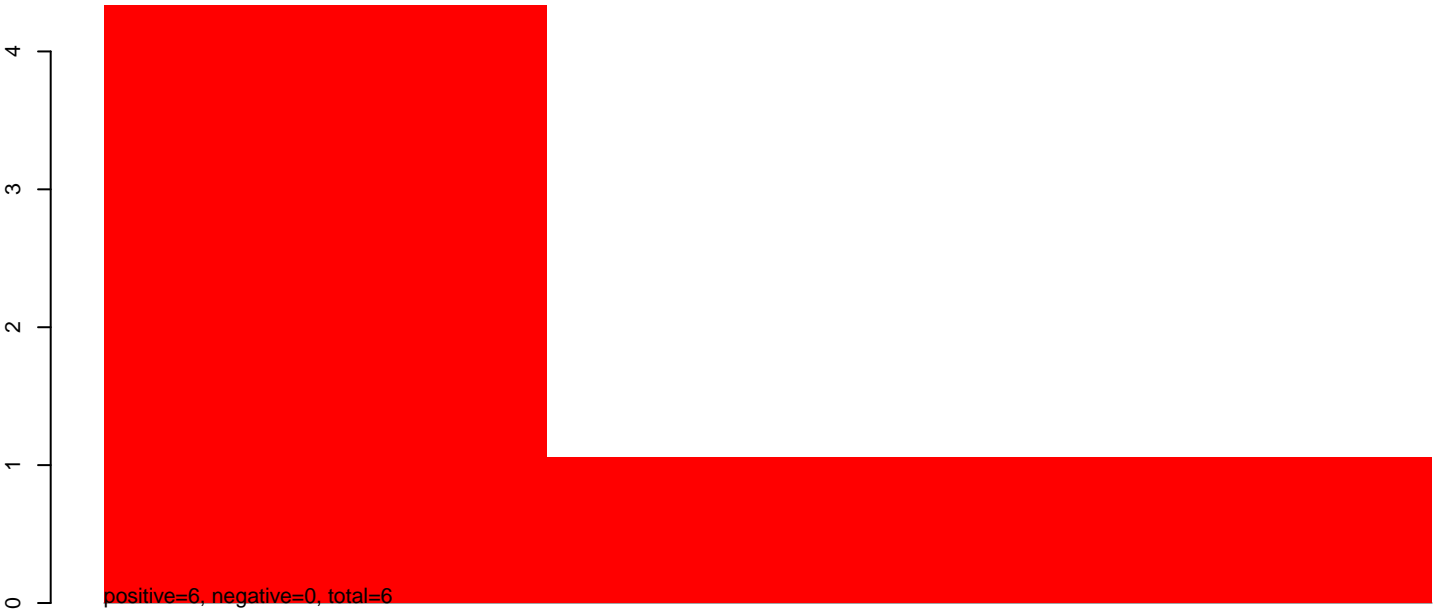


AeAeg_Testes.rep



40 60 80 100

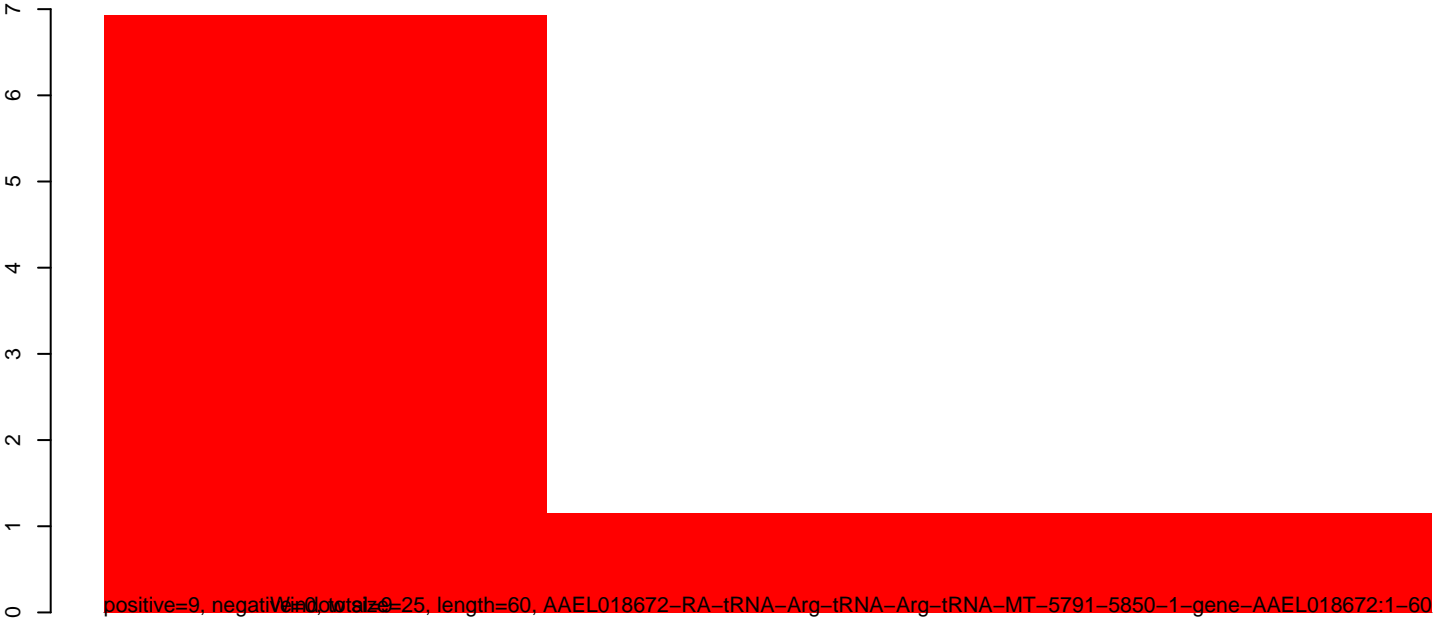
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep

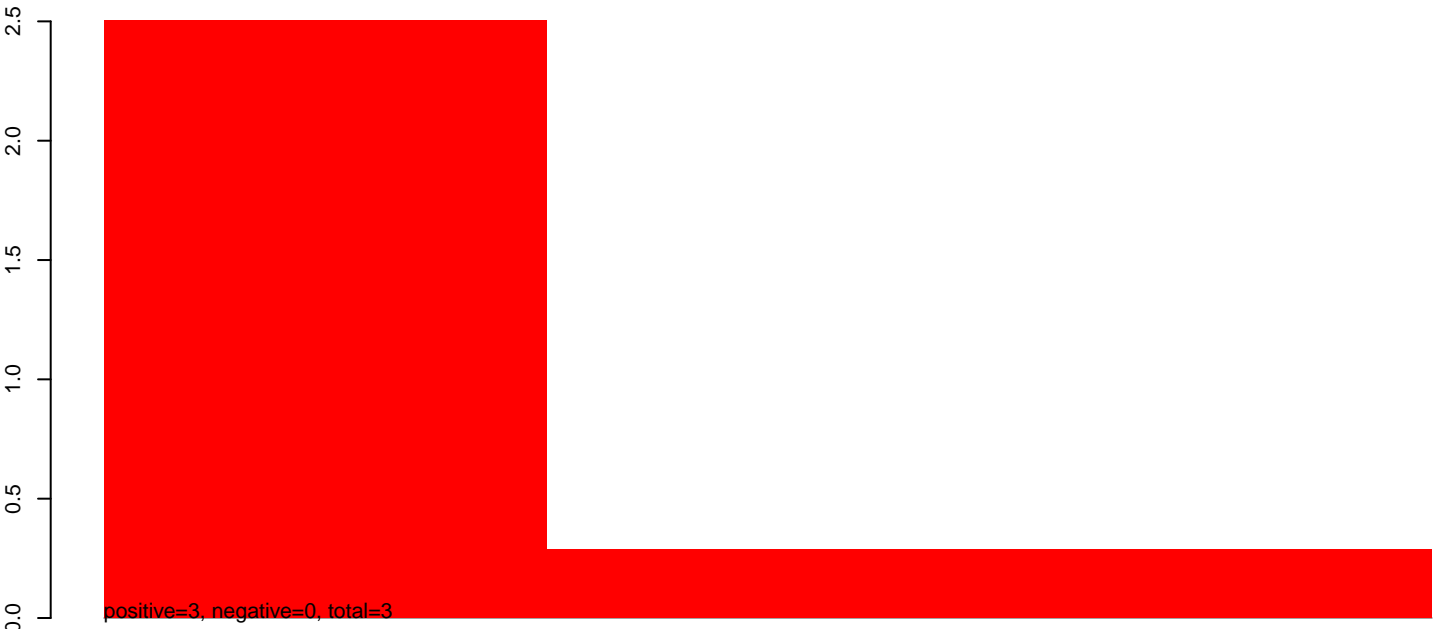


AeAeg_Testes.rep

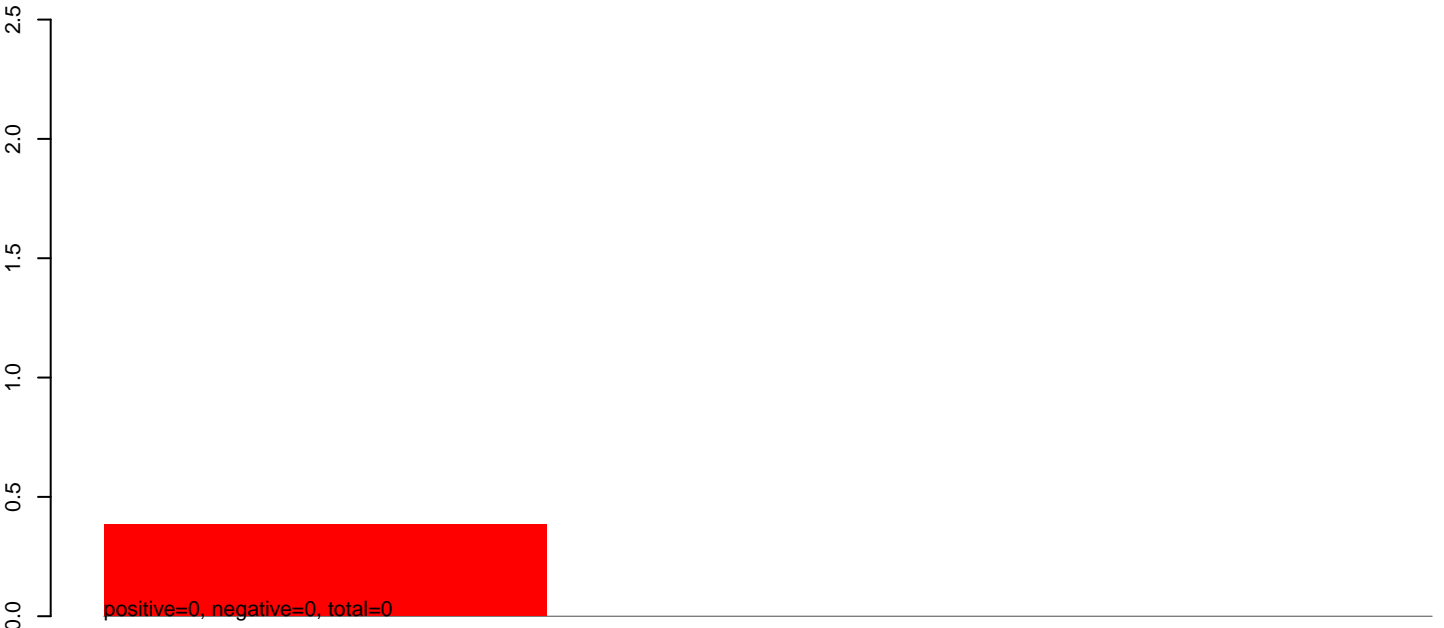


40 60 80 100

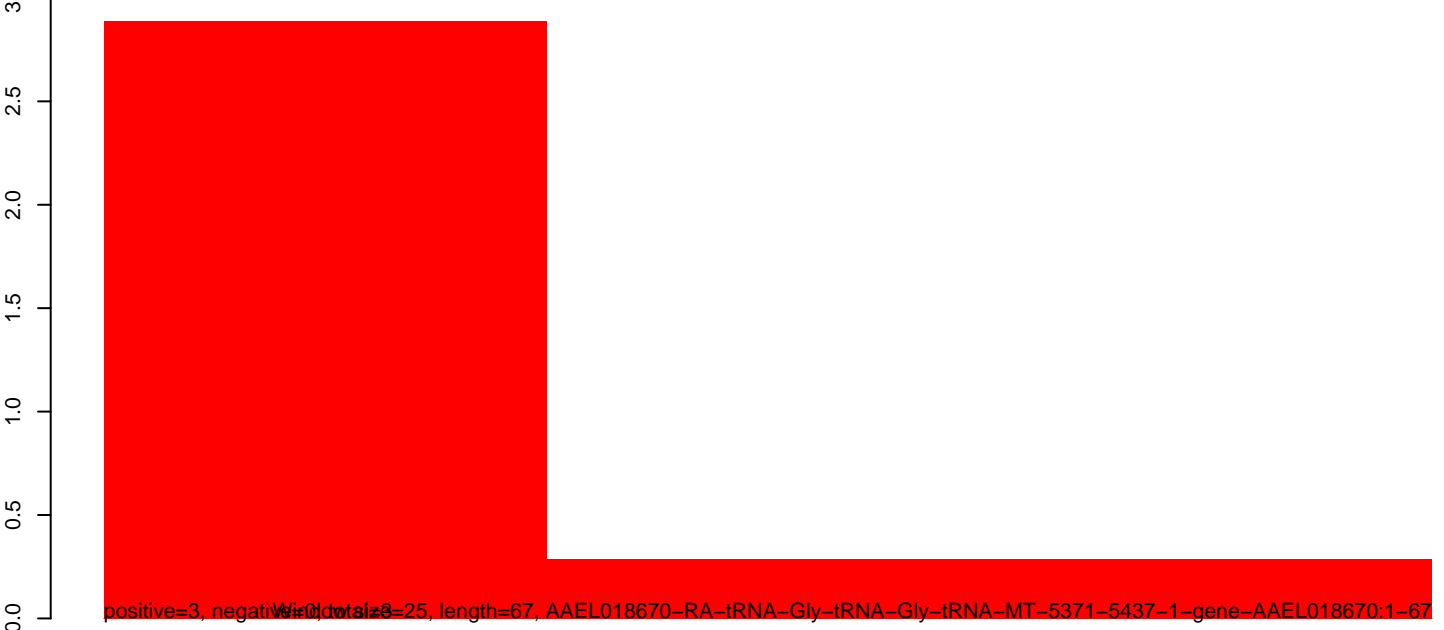
AeAeg_Testes.18_23.rep



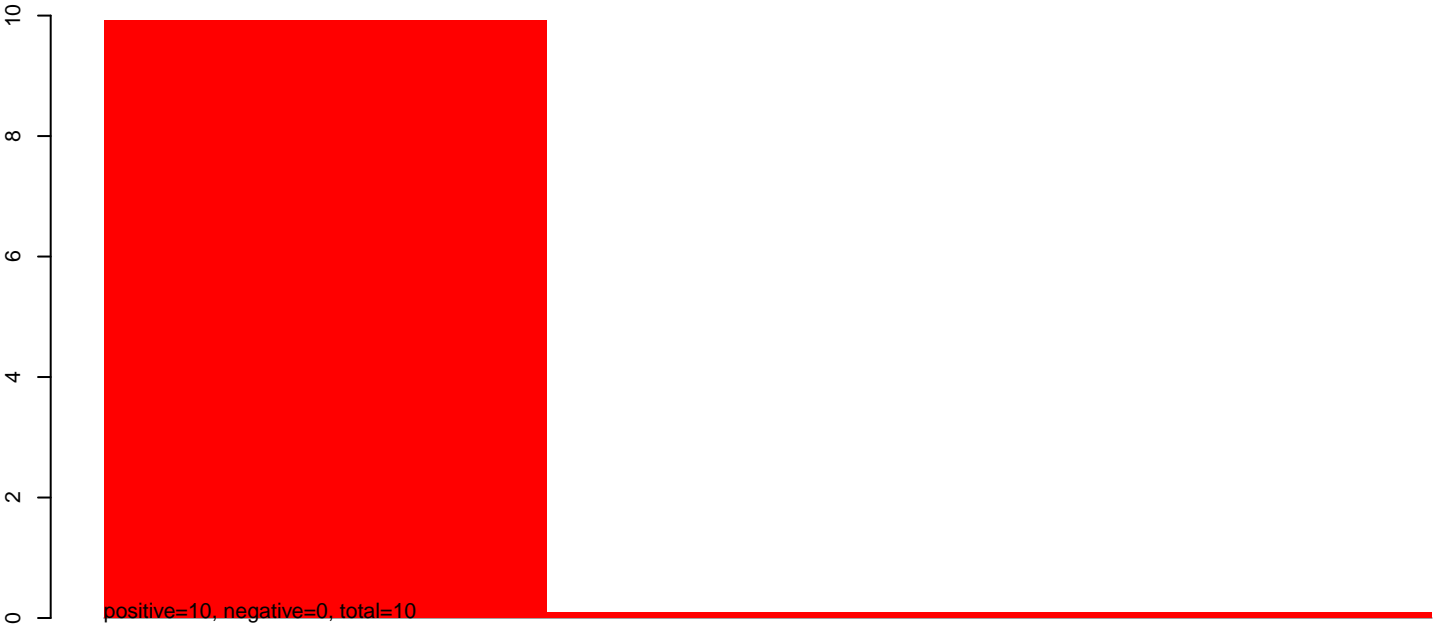
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



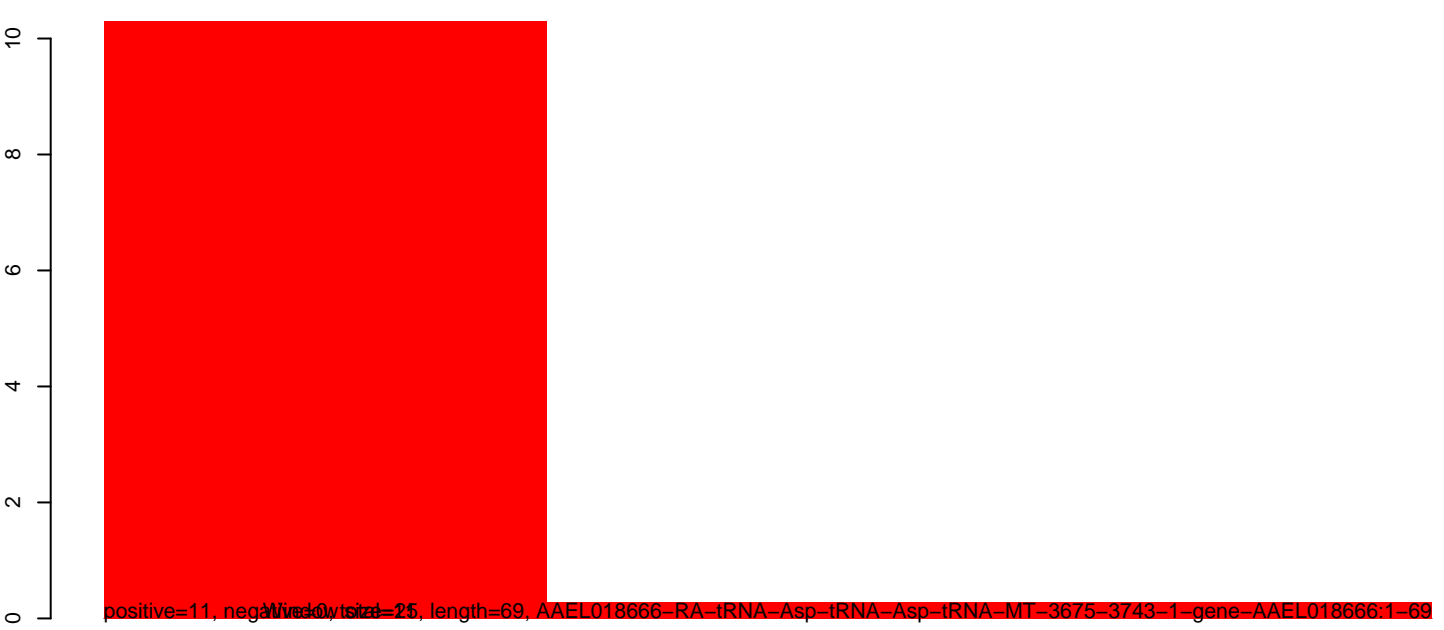
AeAeg_Testes.18_23.rep



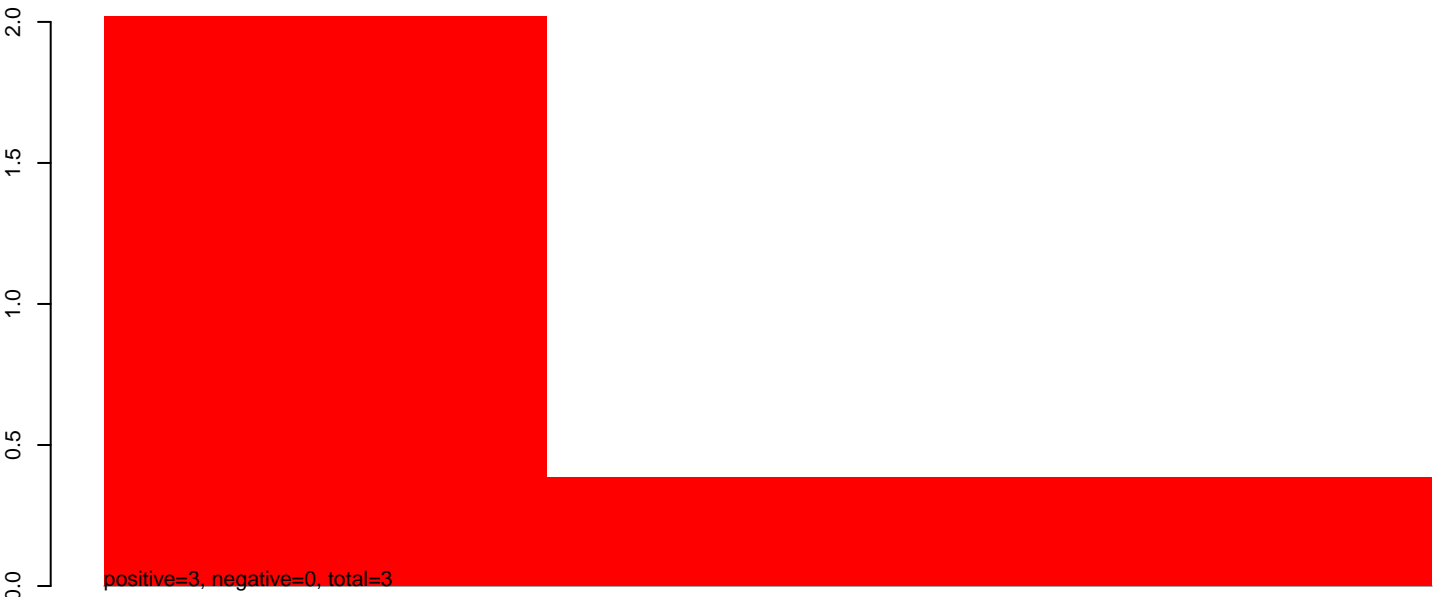
AeAeg_Testes.24_35.rep



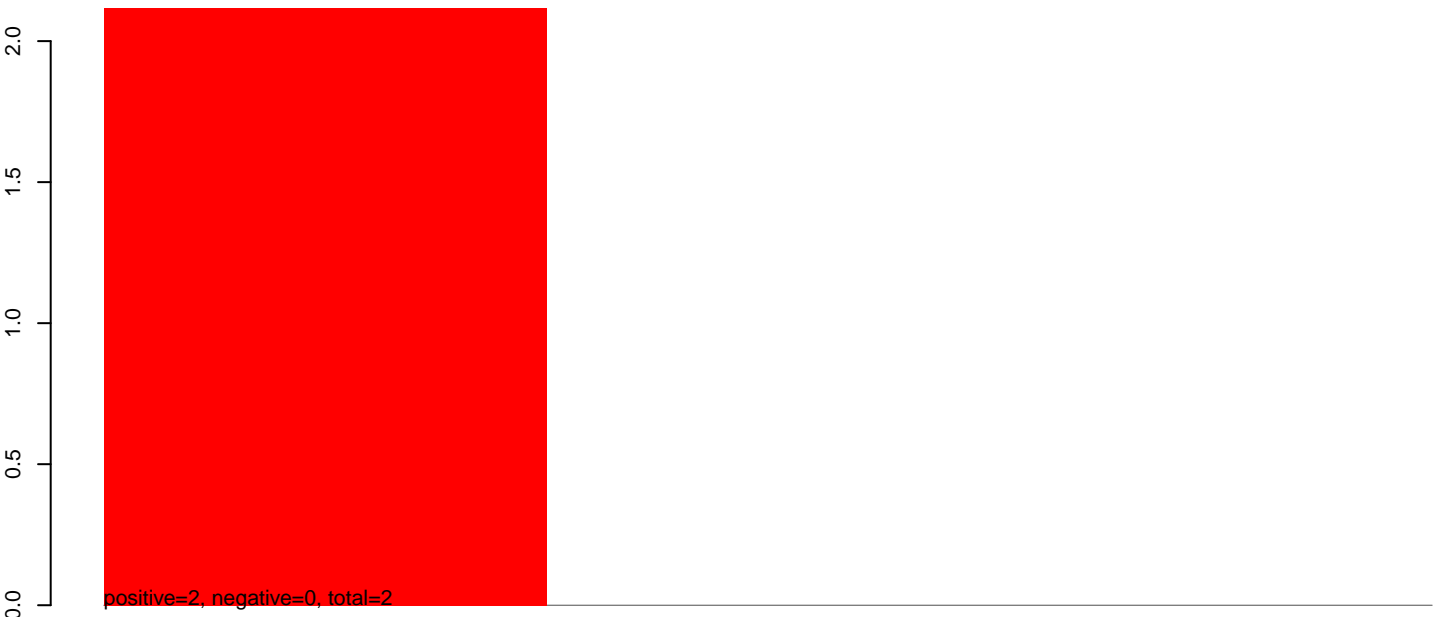
AeAeg_Testes.rep



AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep

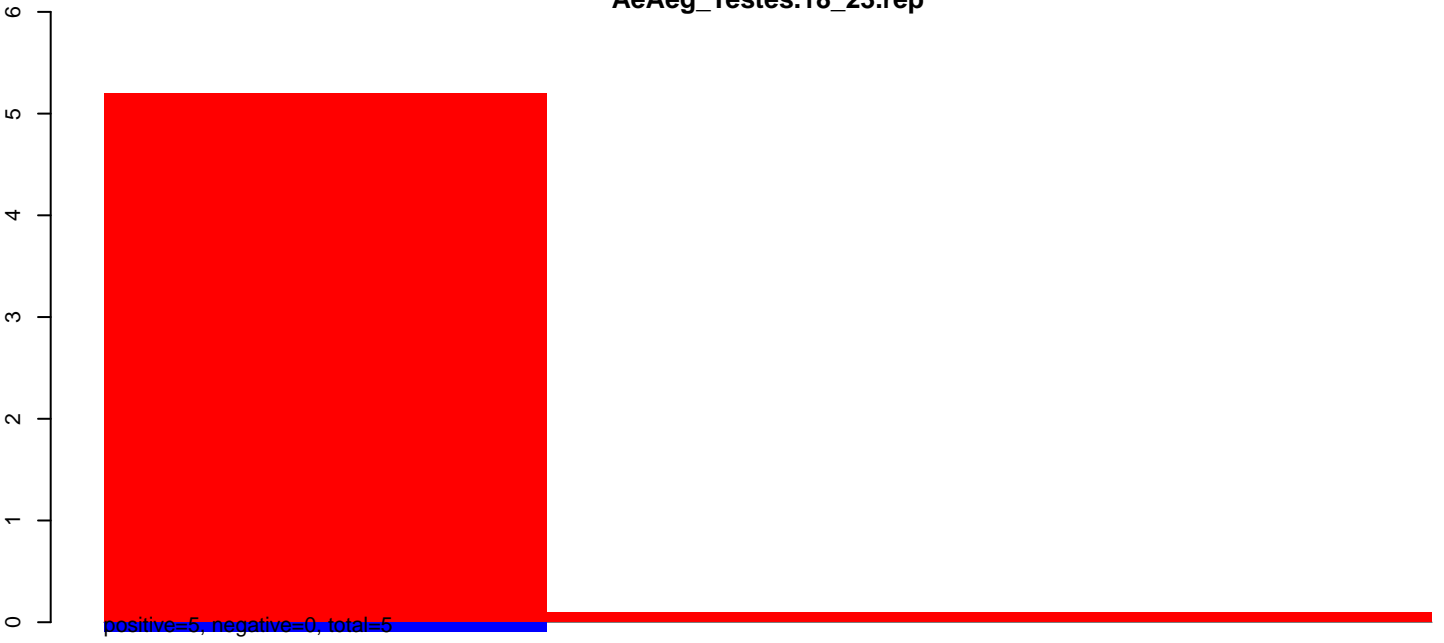


AeAeg_Testes.rep

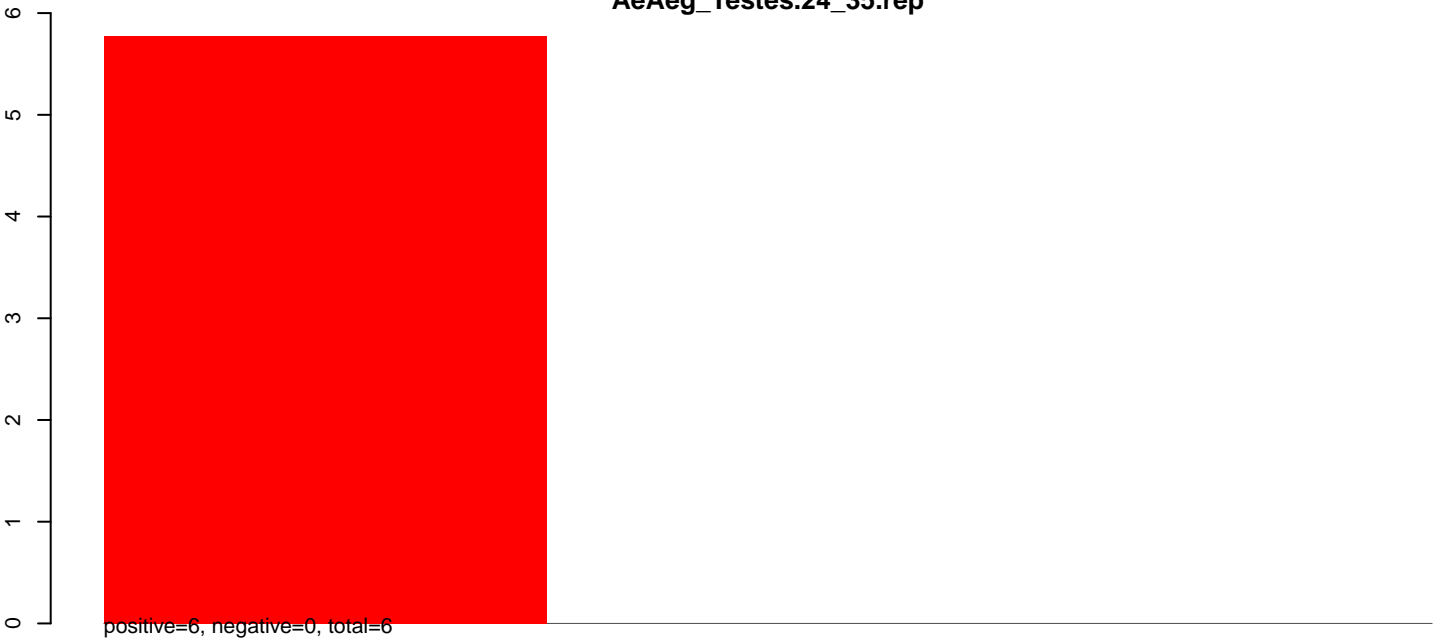


40 60 80 100

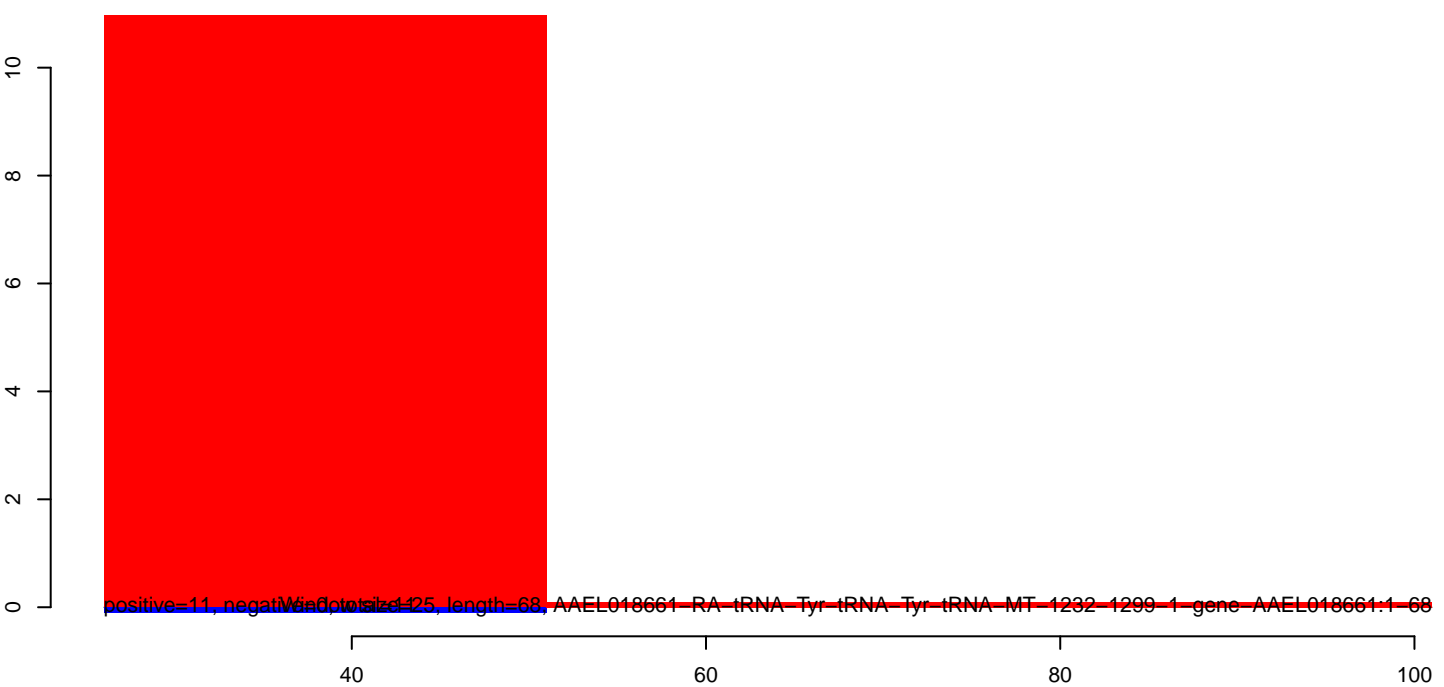
AeAeg_Testes.18_23.rep



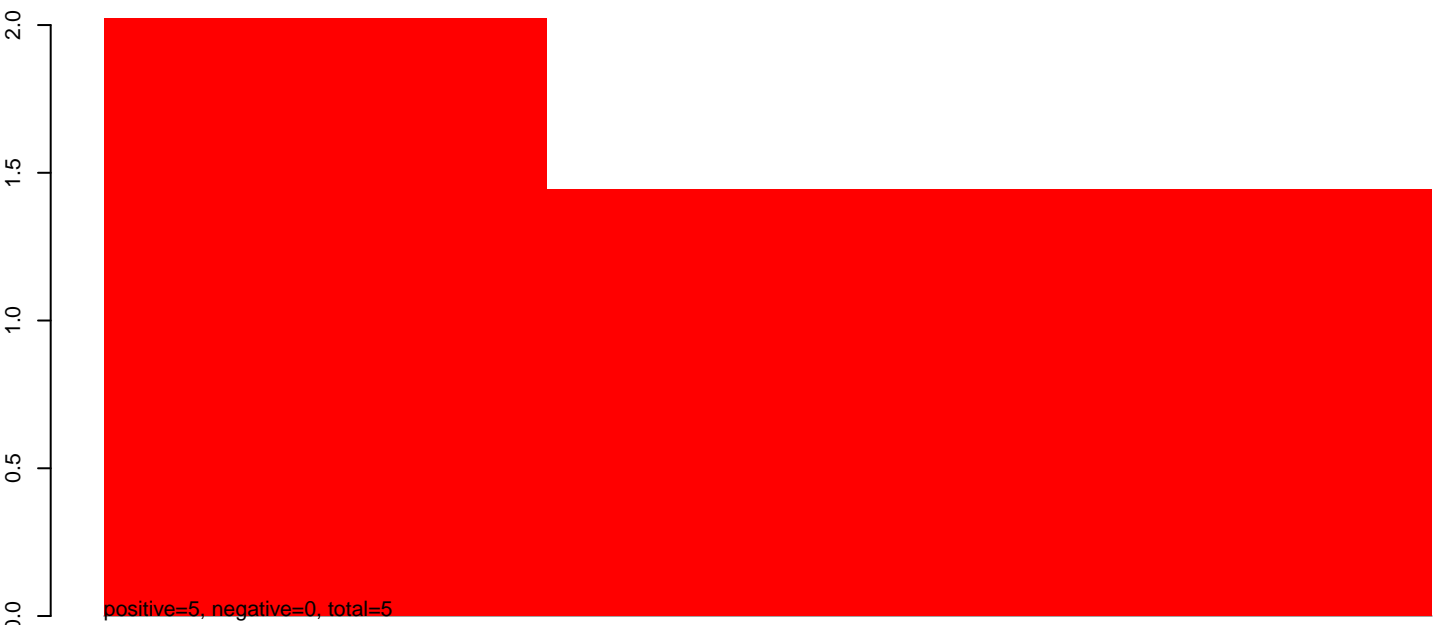
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



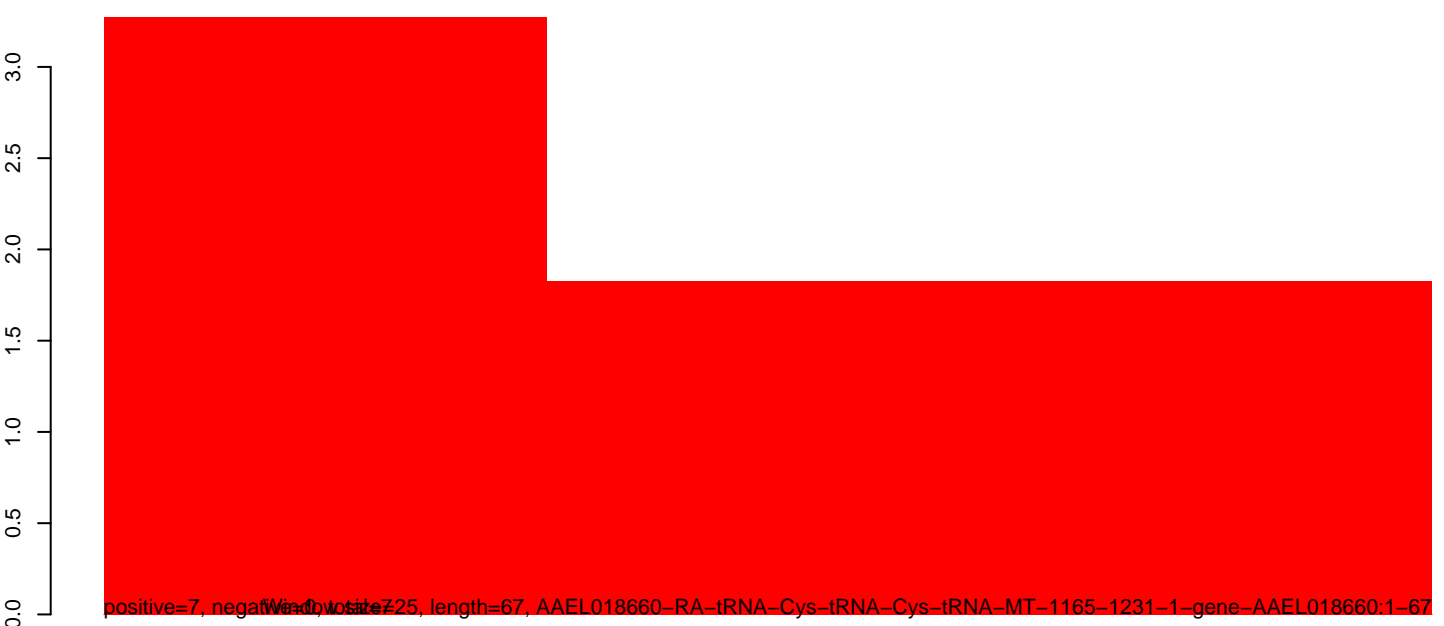
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep

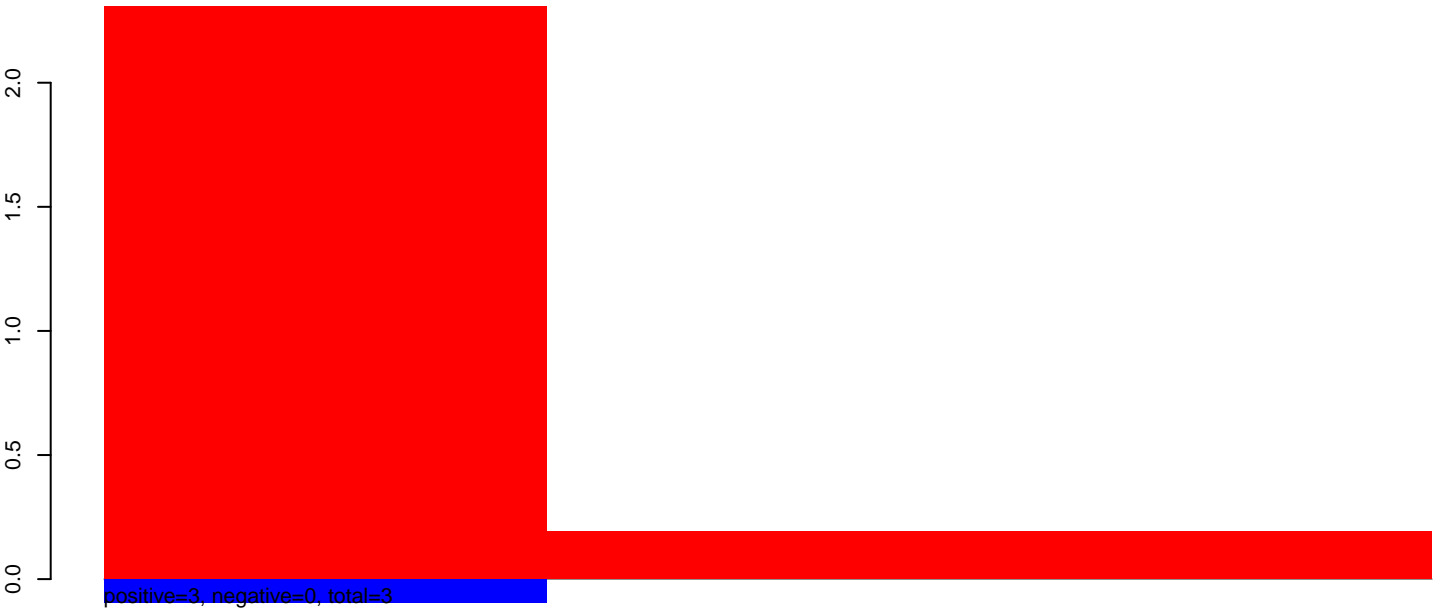


AeAeg_Testes.rep



40 60 80 100

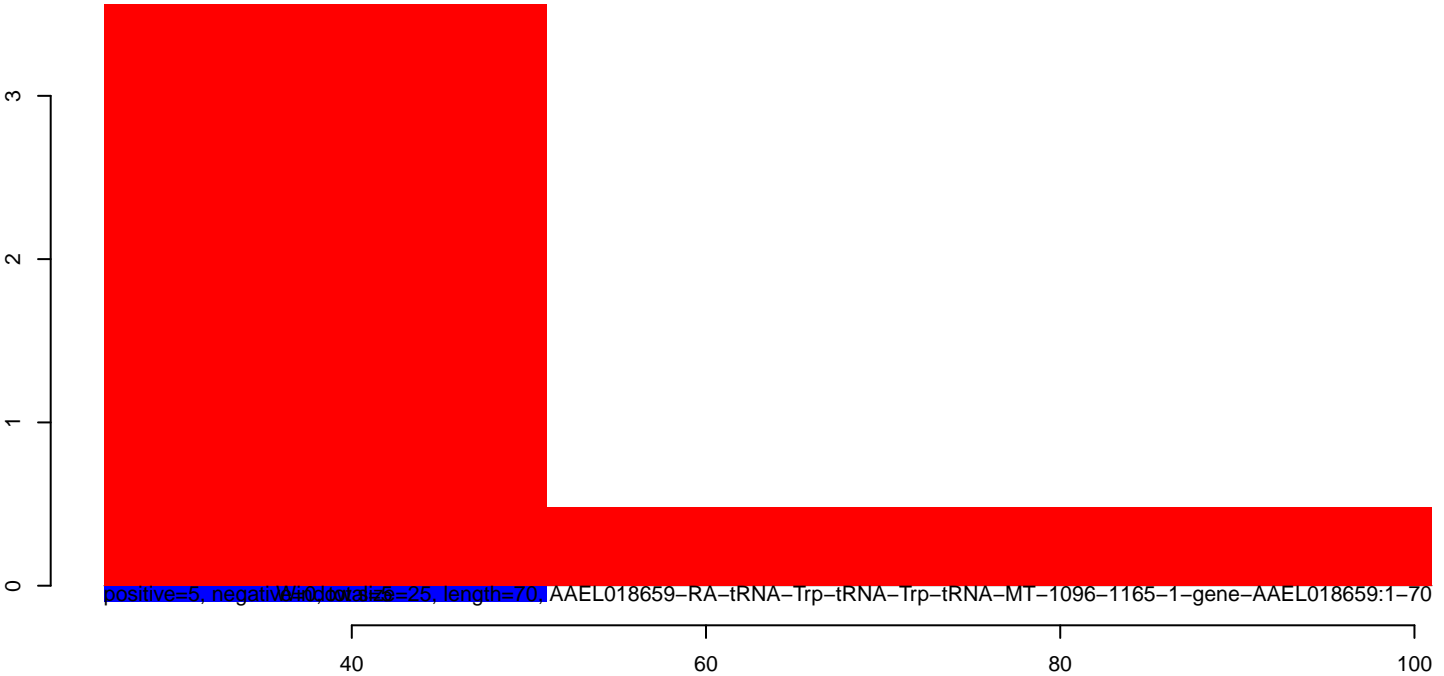
AeAeg_Testes.18_23.rep



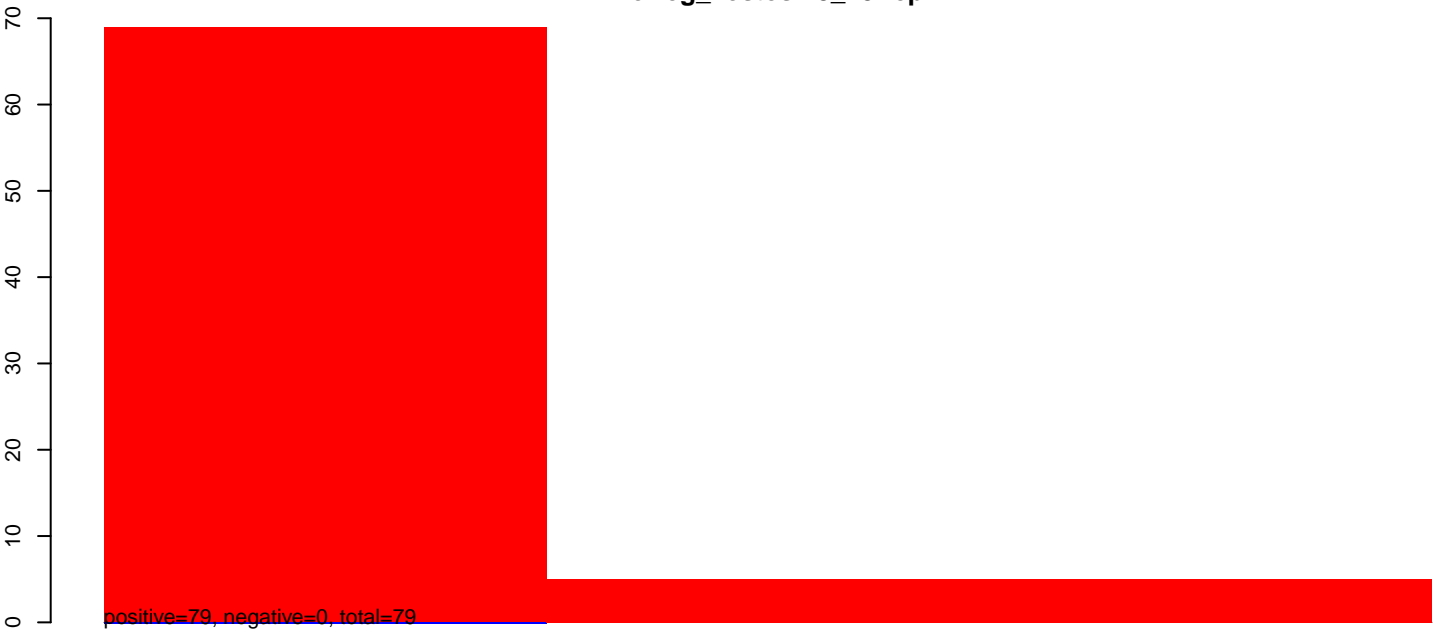
AeAeg_Testes.24_35.rep



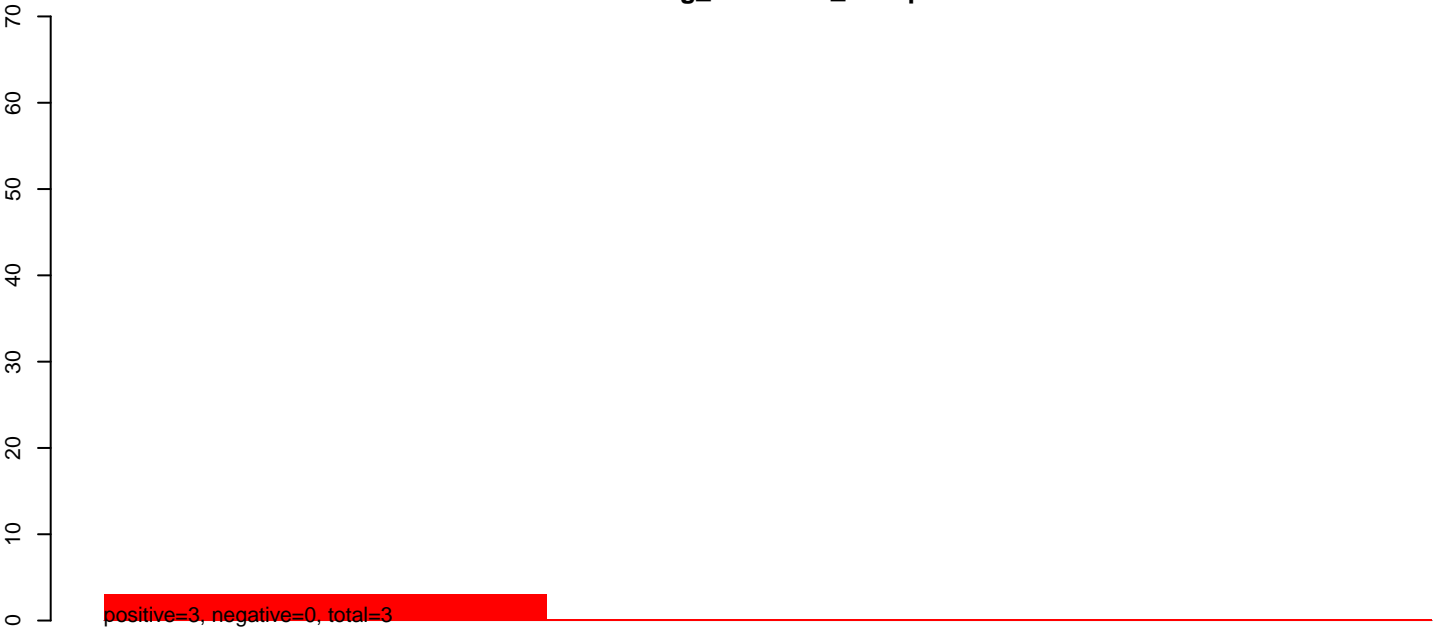
AeAeg_Testes.rep



AeAeg_Testes.18_23.rep



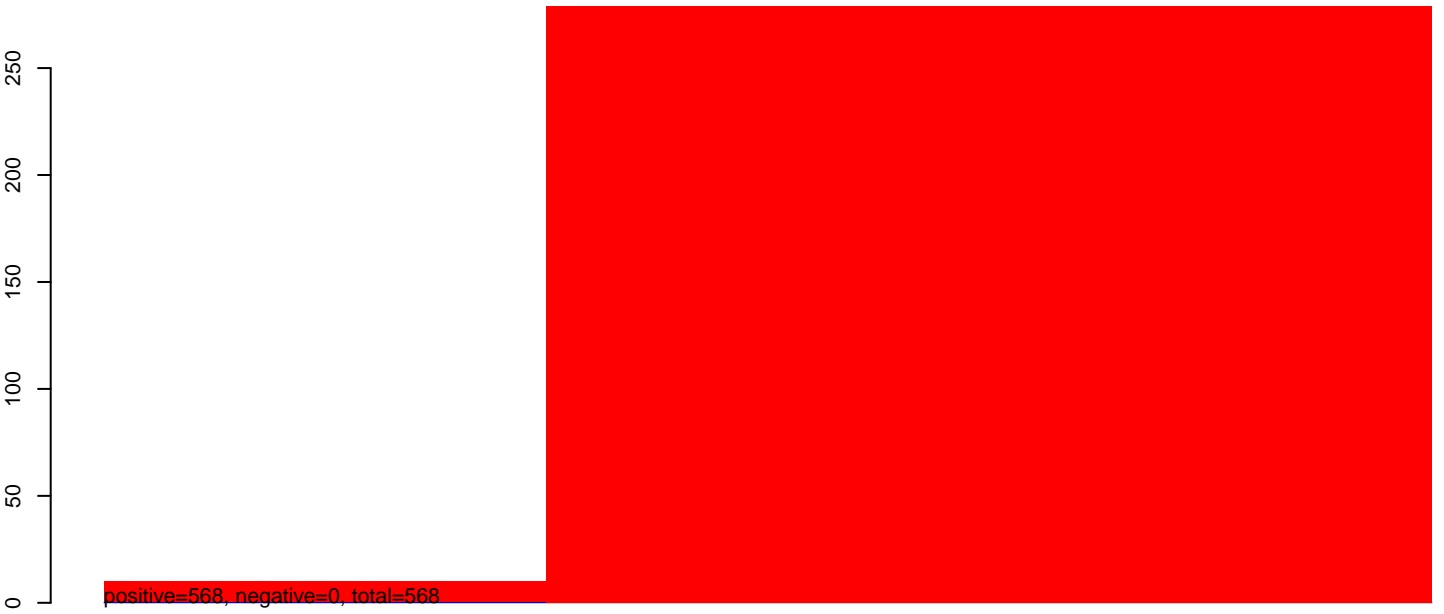
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



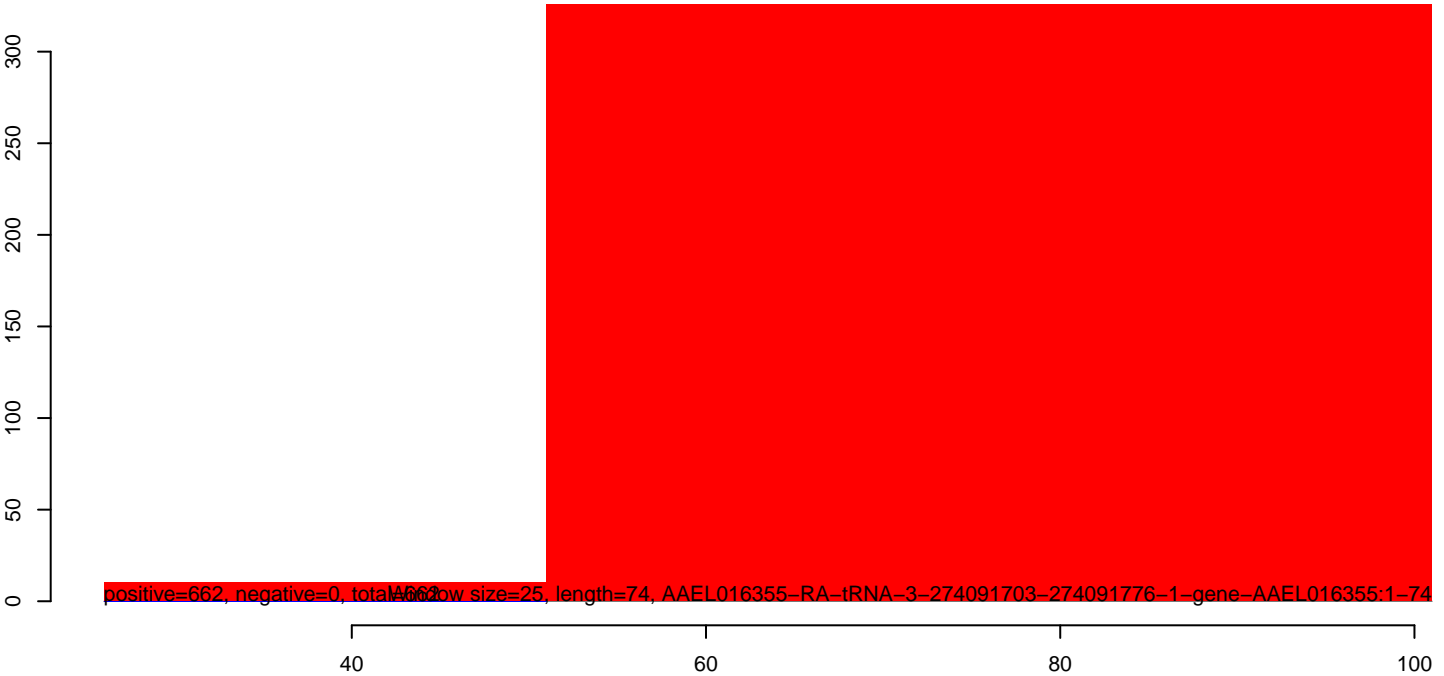
AeAeg_Testes.18_23.rep



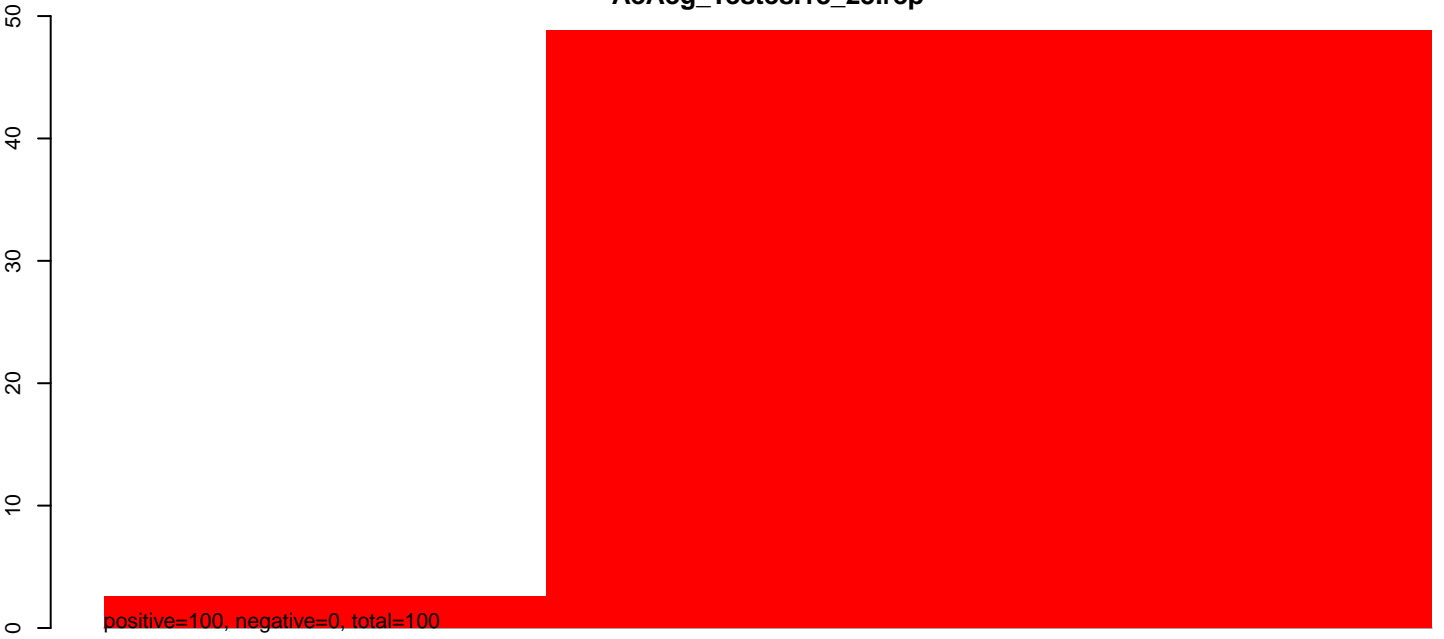
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep

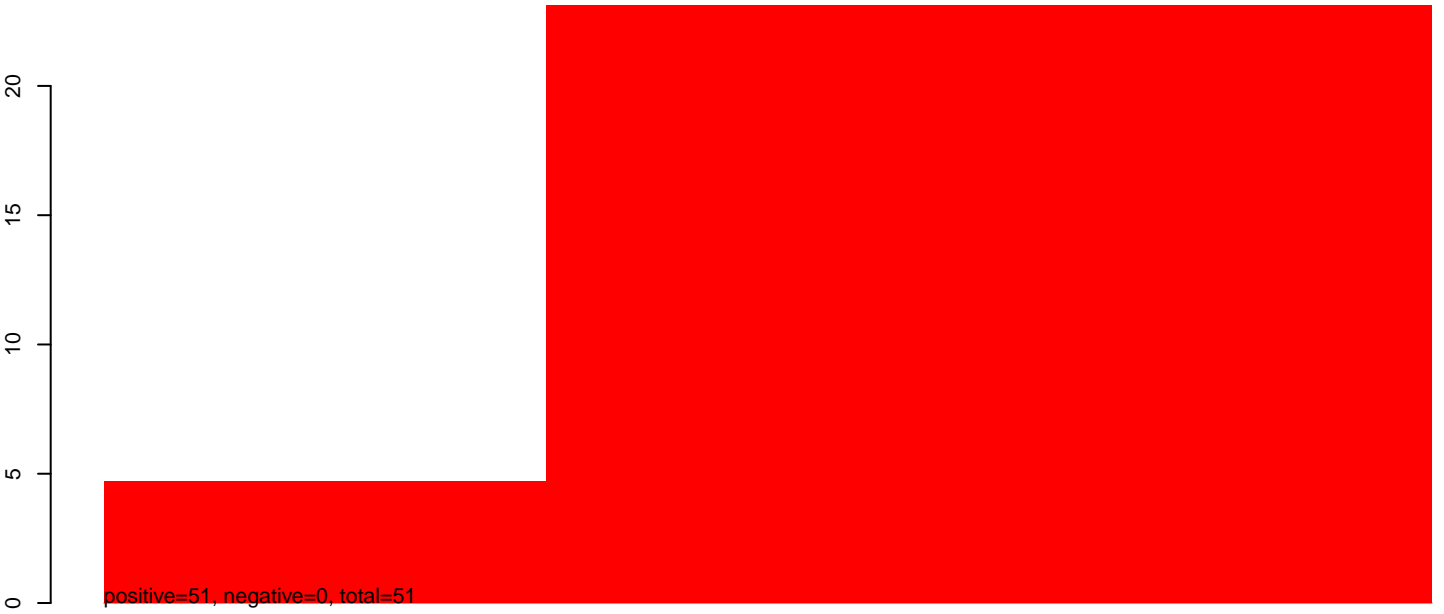


AeAeg_Testes.rep



40 60 80 100

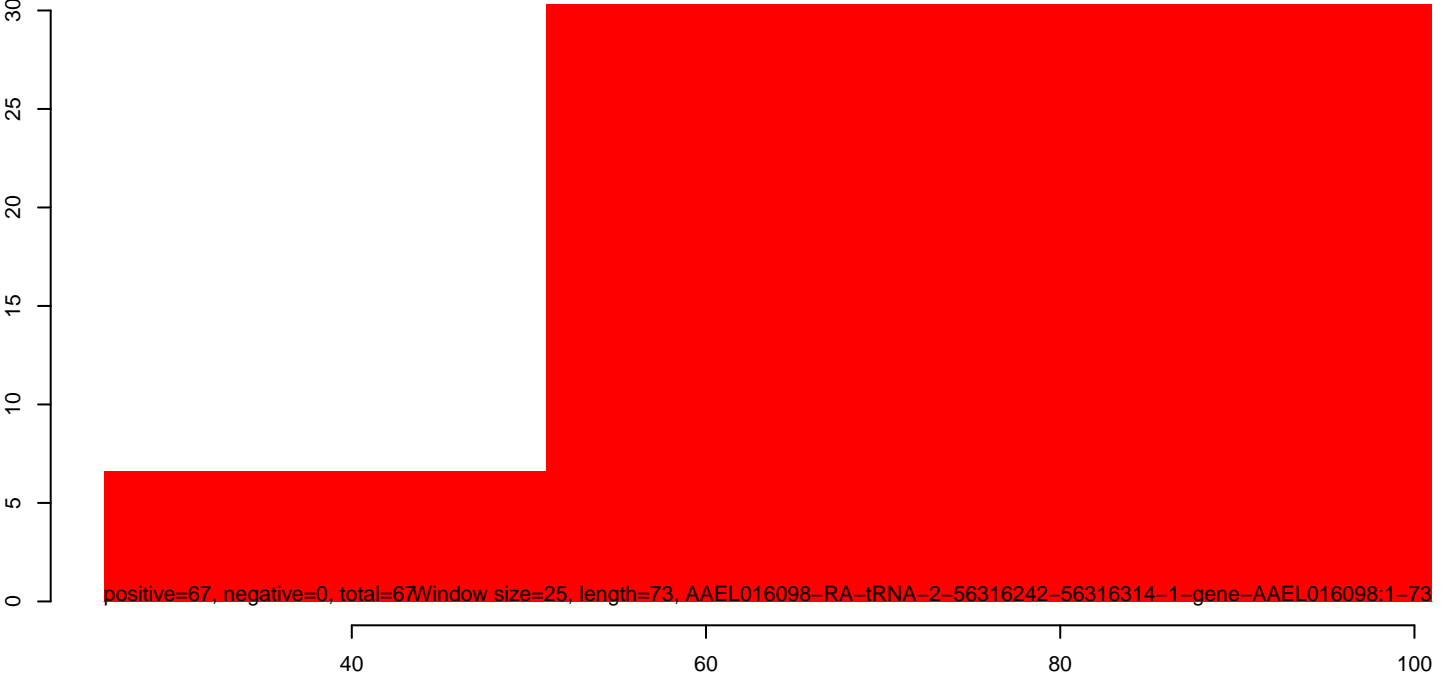
AeAeg_Testes.18_23.rep



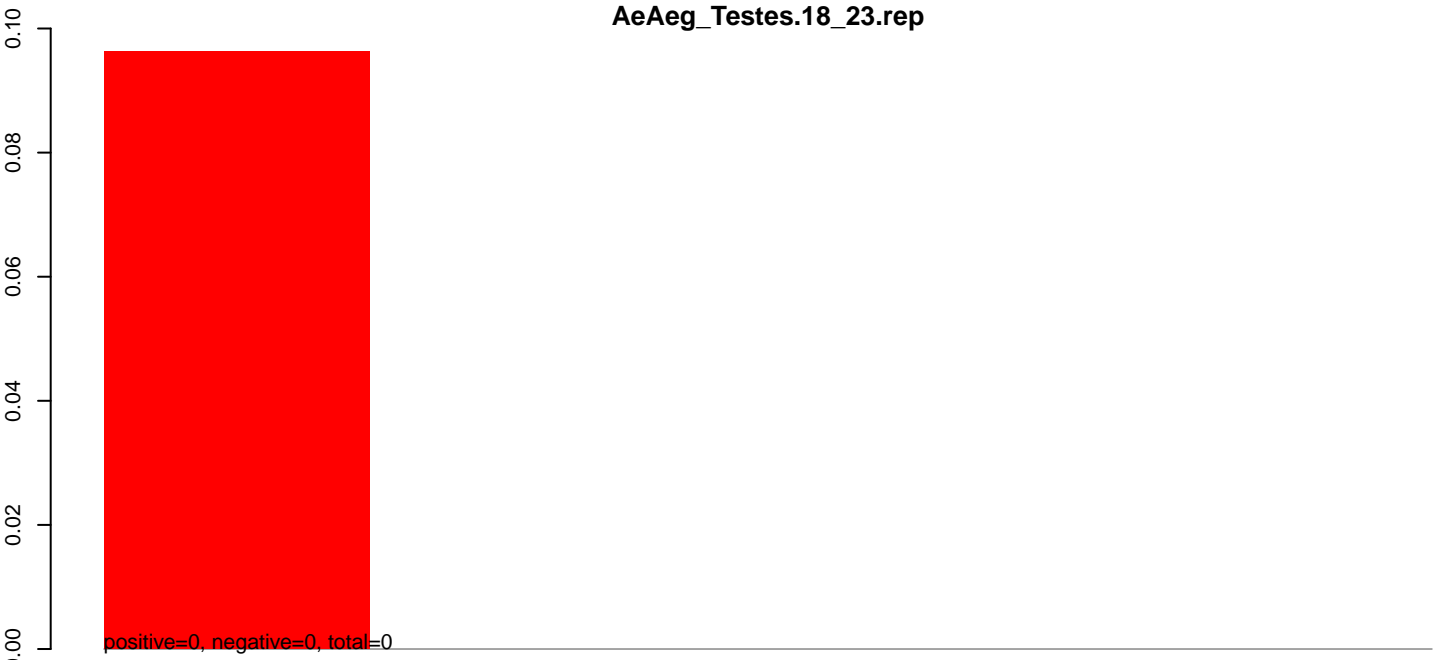
AeAeg_Testes.24_35.rep



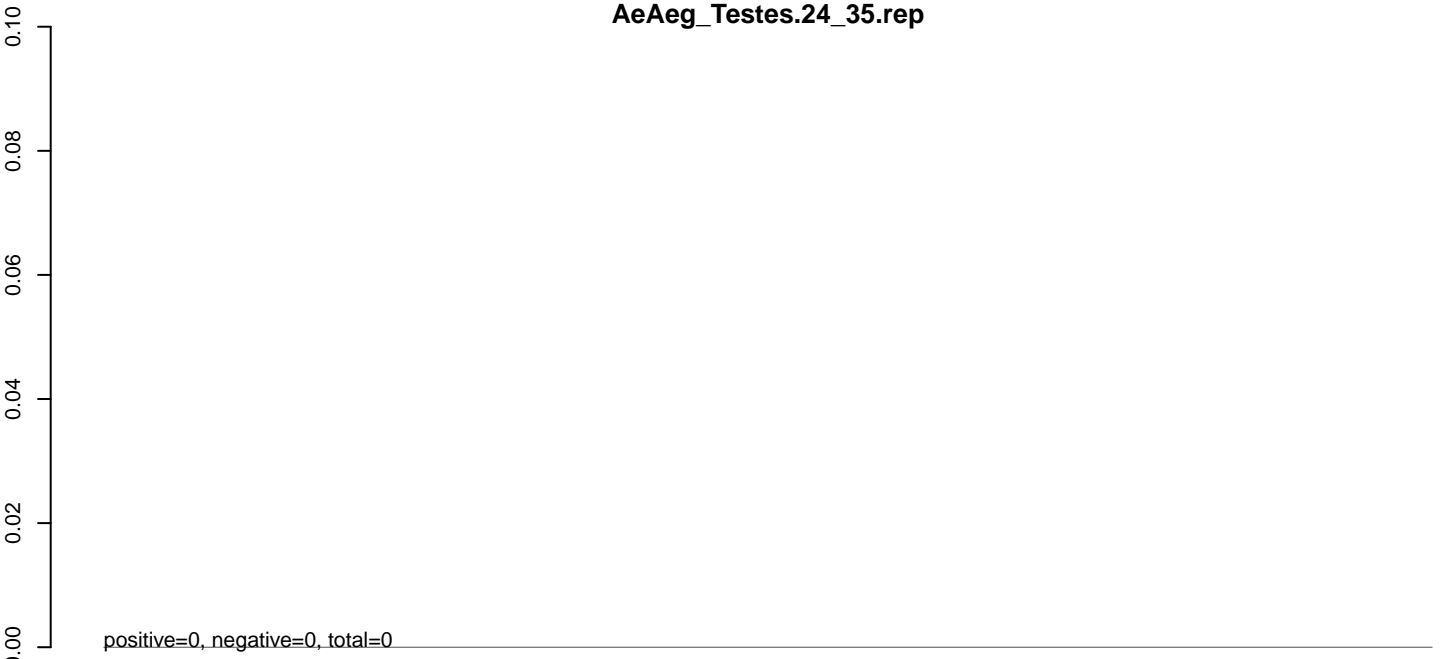
AeAeg_Testes.rep



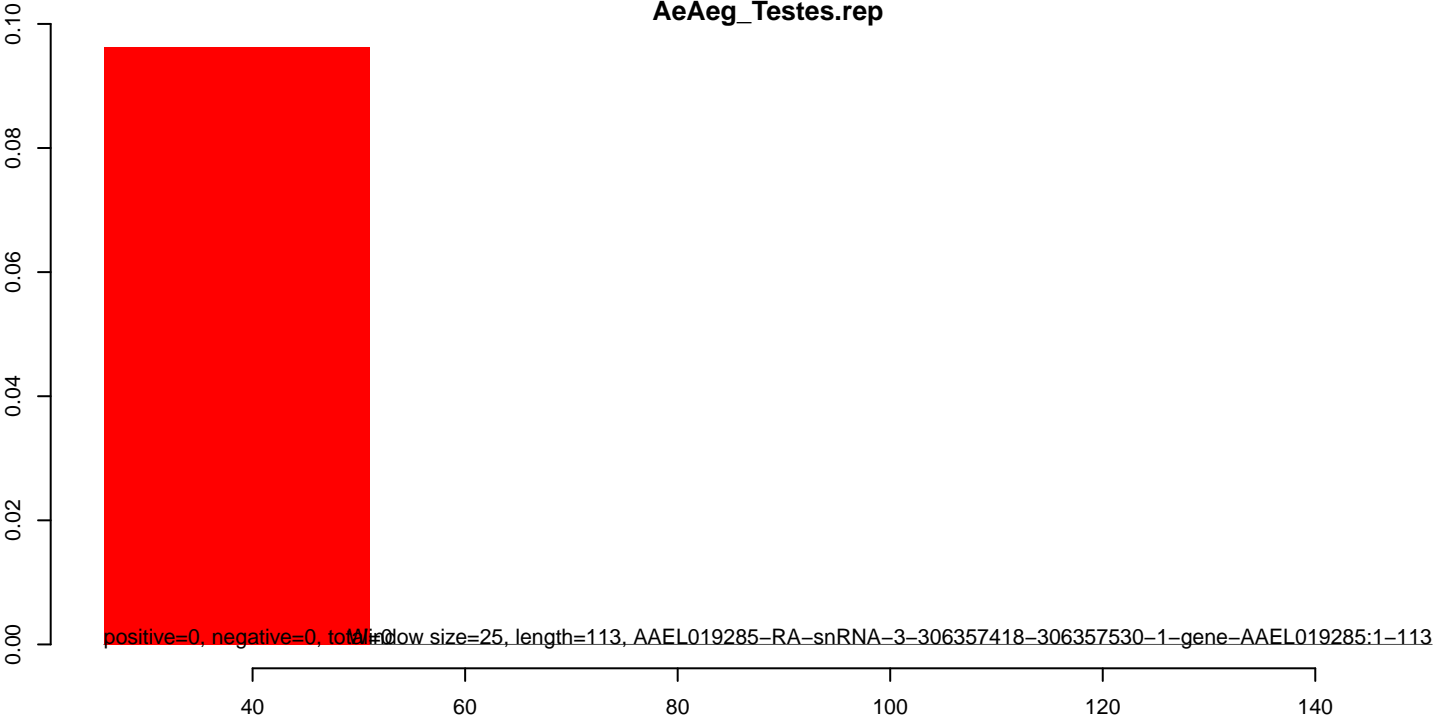
AeAeg_Testes.18_23.rep



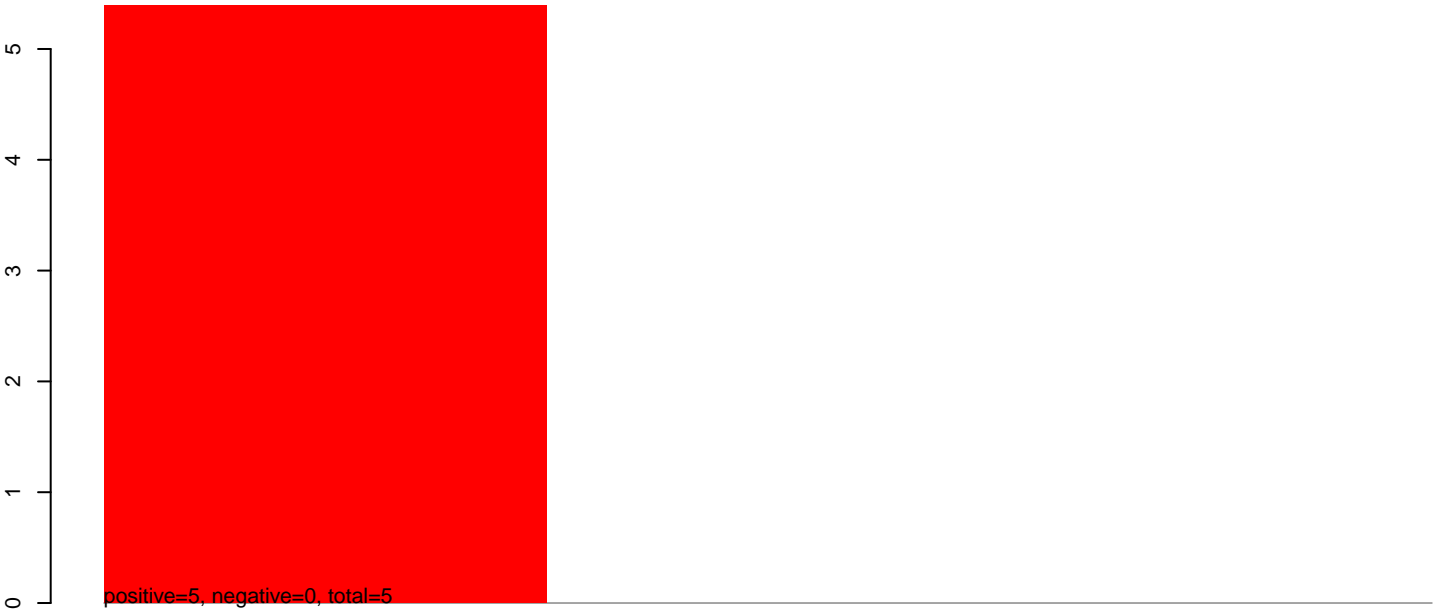
AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



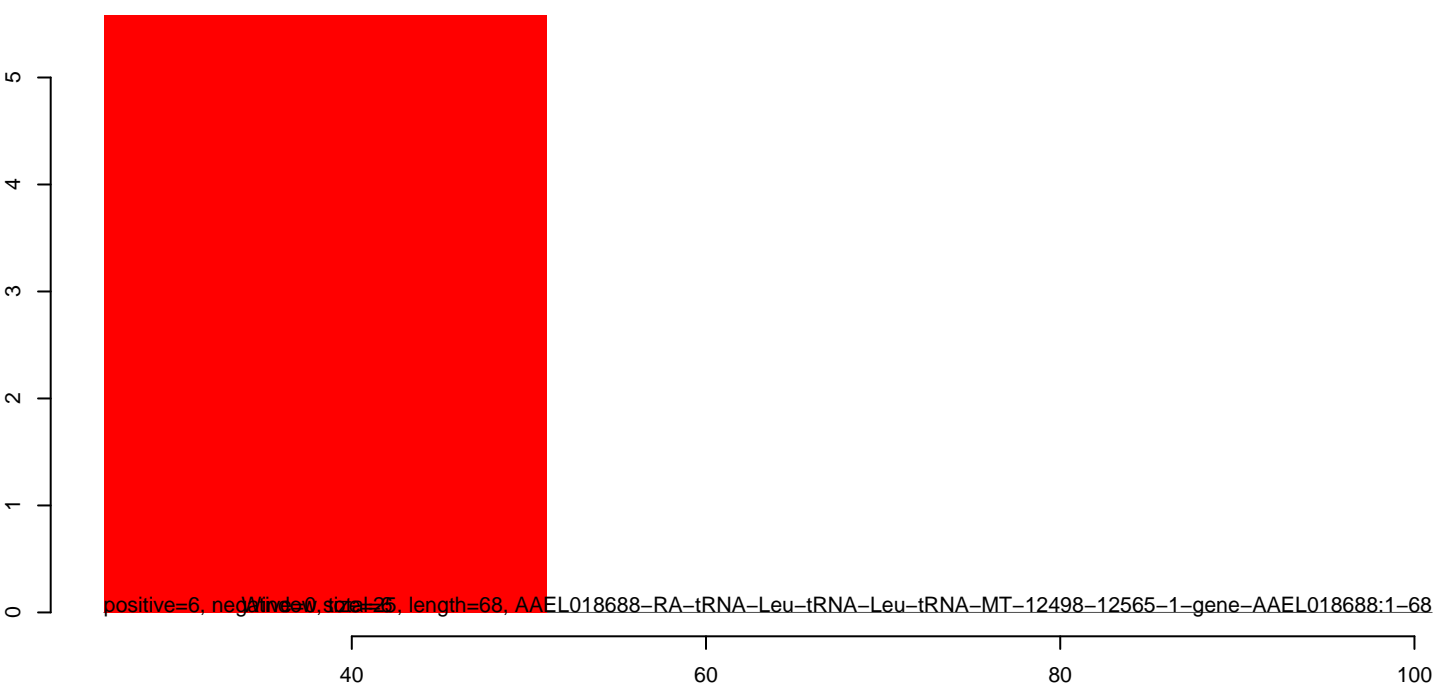
AeAeg_Testes.18_23.rep



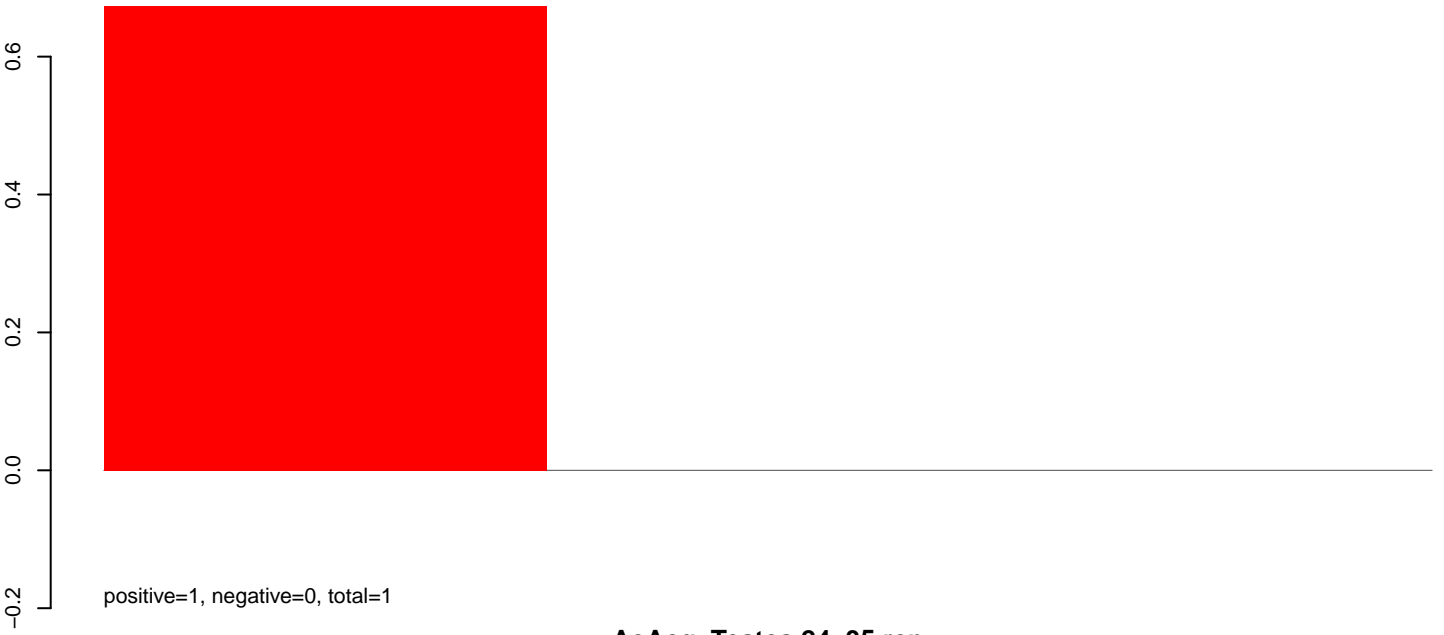
AeAeg_Testes.24_35.rep



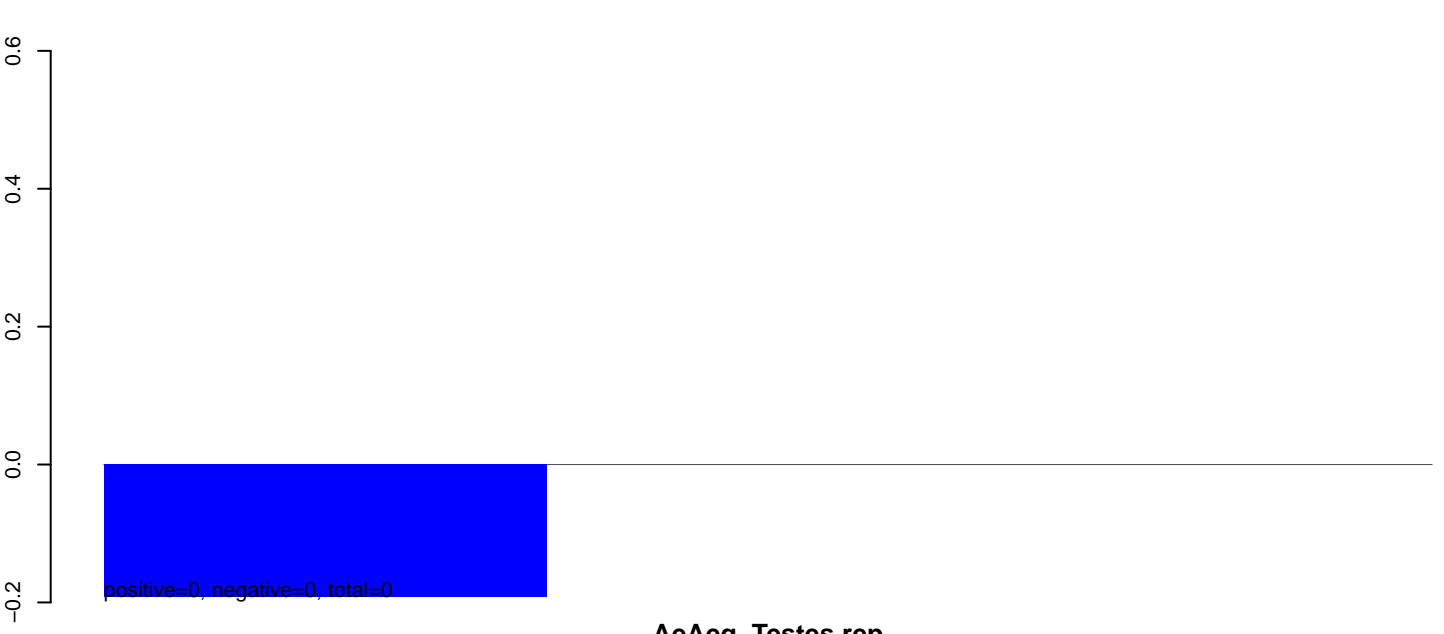
AeAeg_Testes.rep



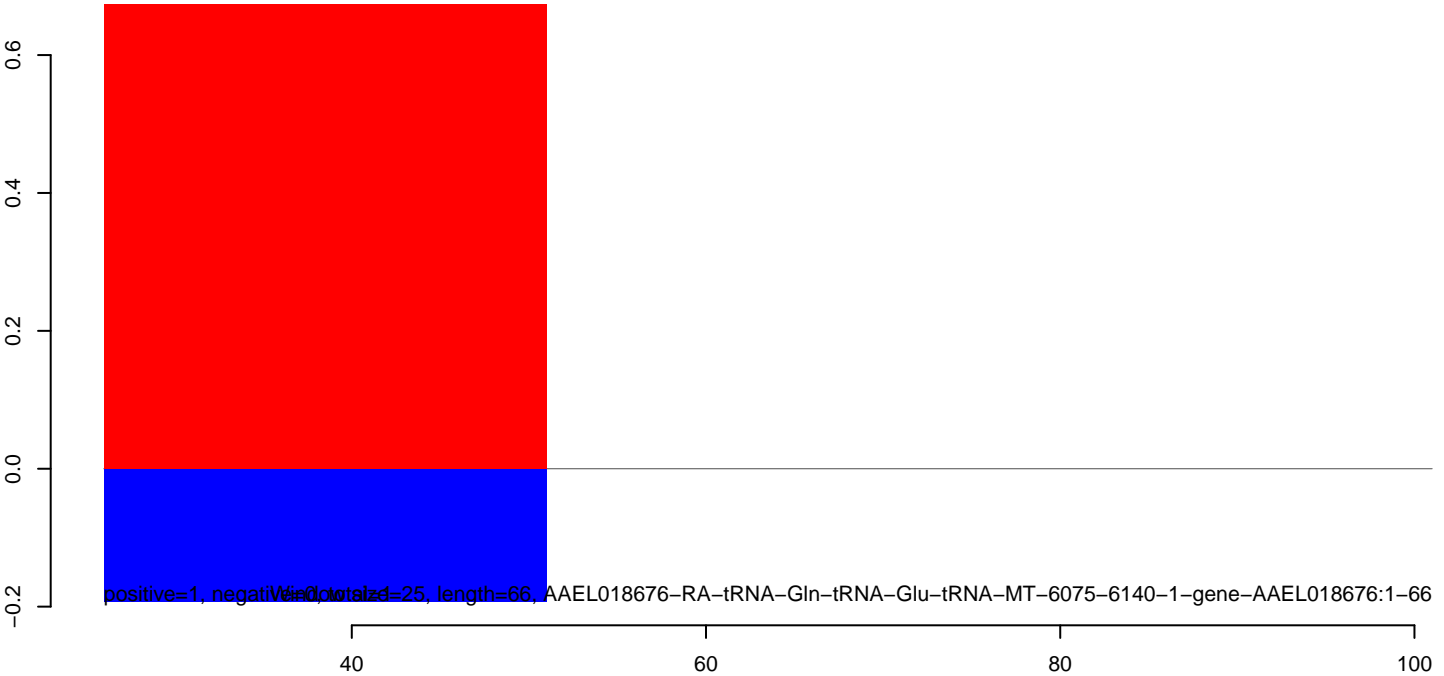
AeAeg_Testes.18_23.rep



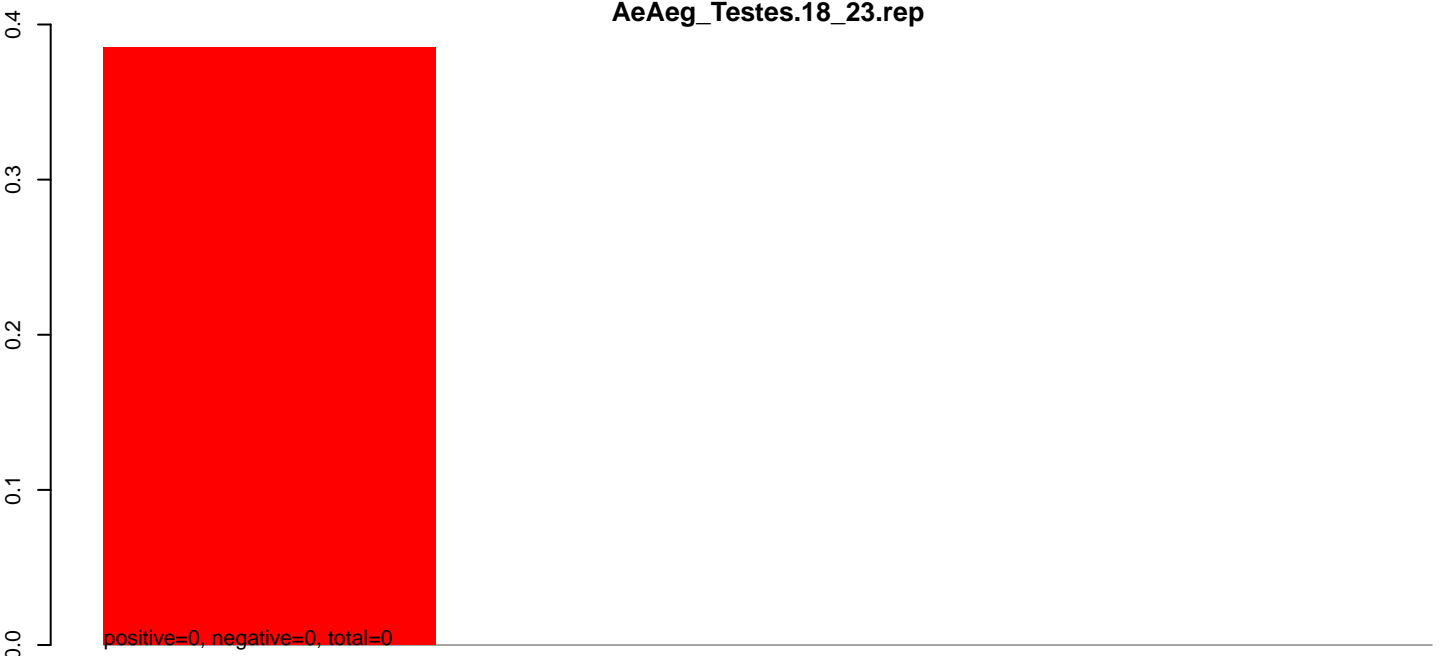
AeAeg_Testes.24_35.rep



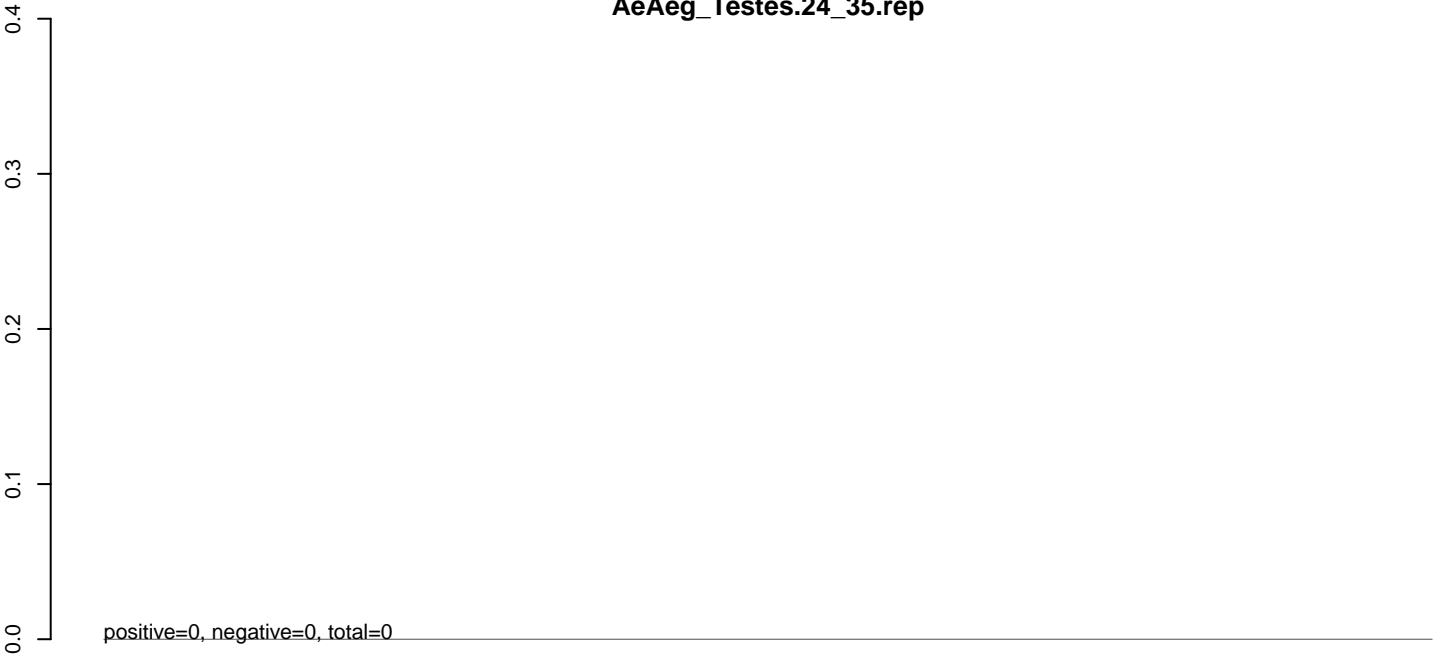
AeAeg_Testes.rep



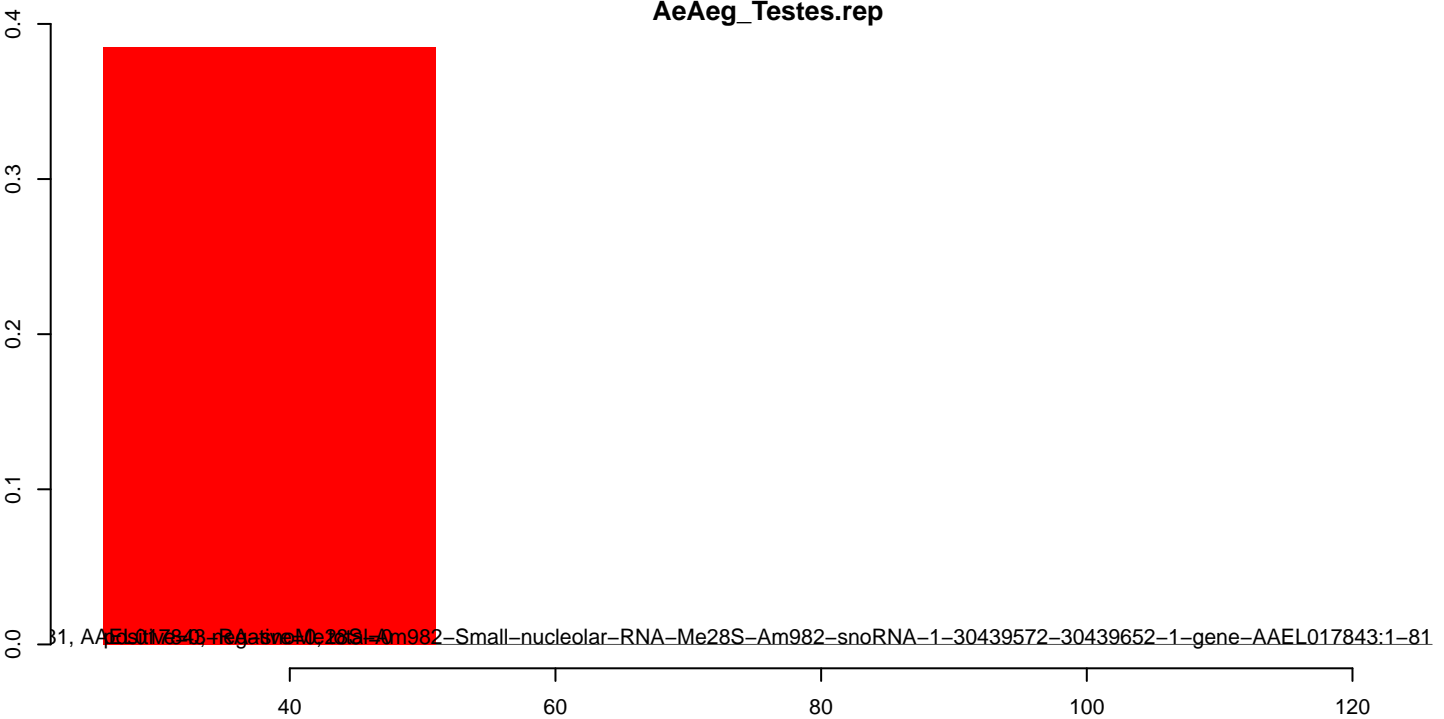
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



AeAeg_Testes.rep

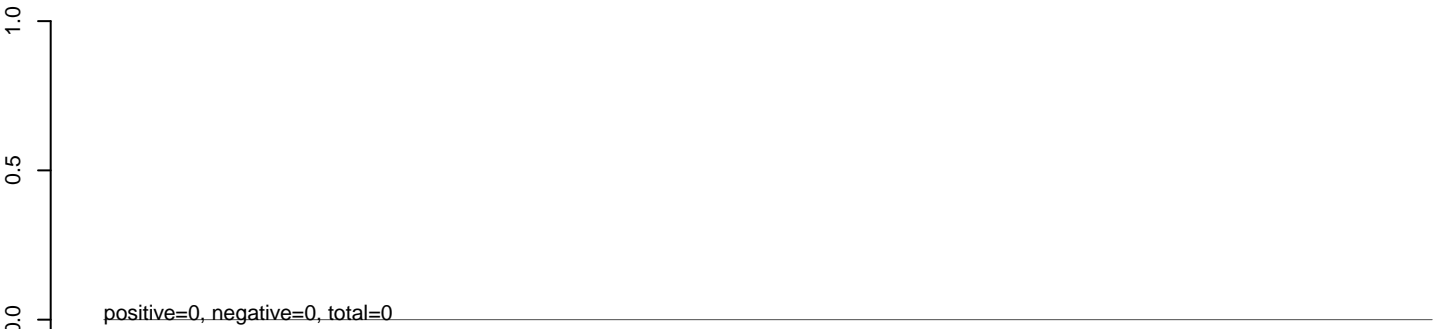


31, AAF017843, rRNA-40, 28S-40, 982-Small-nucleolar-RNA-Me28S-Am982-snoRNA-1-30439572-30439652-1-gene-AAEL017843:1-81

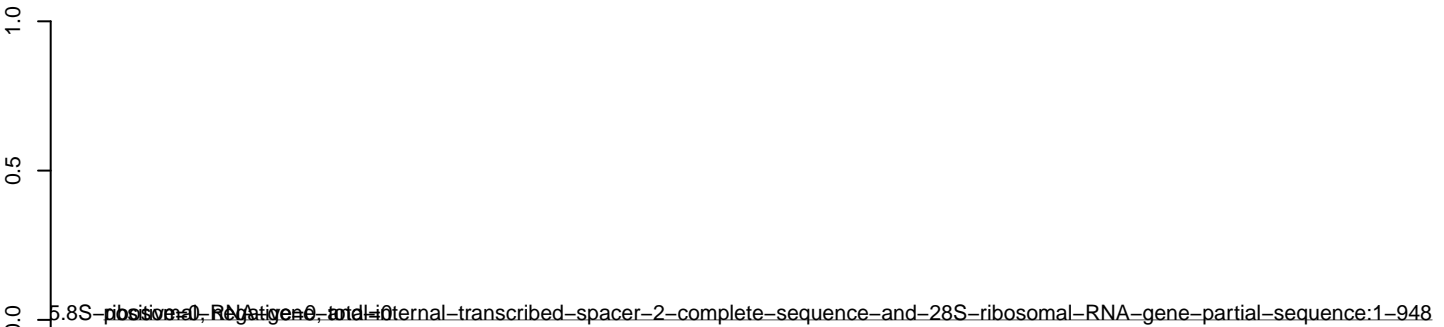
AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



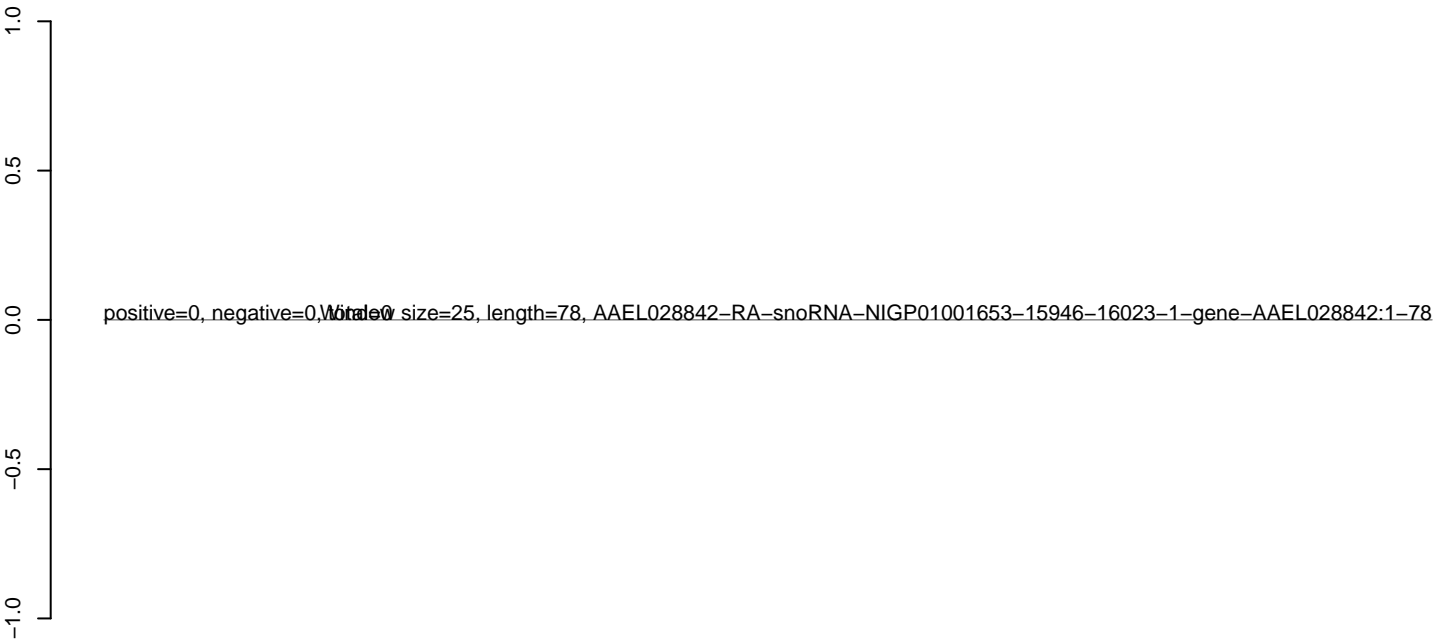
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AeAeg_Testes.rep



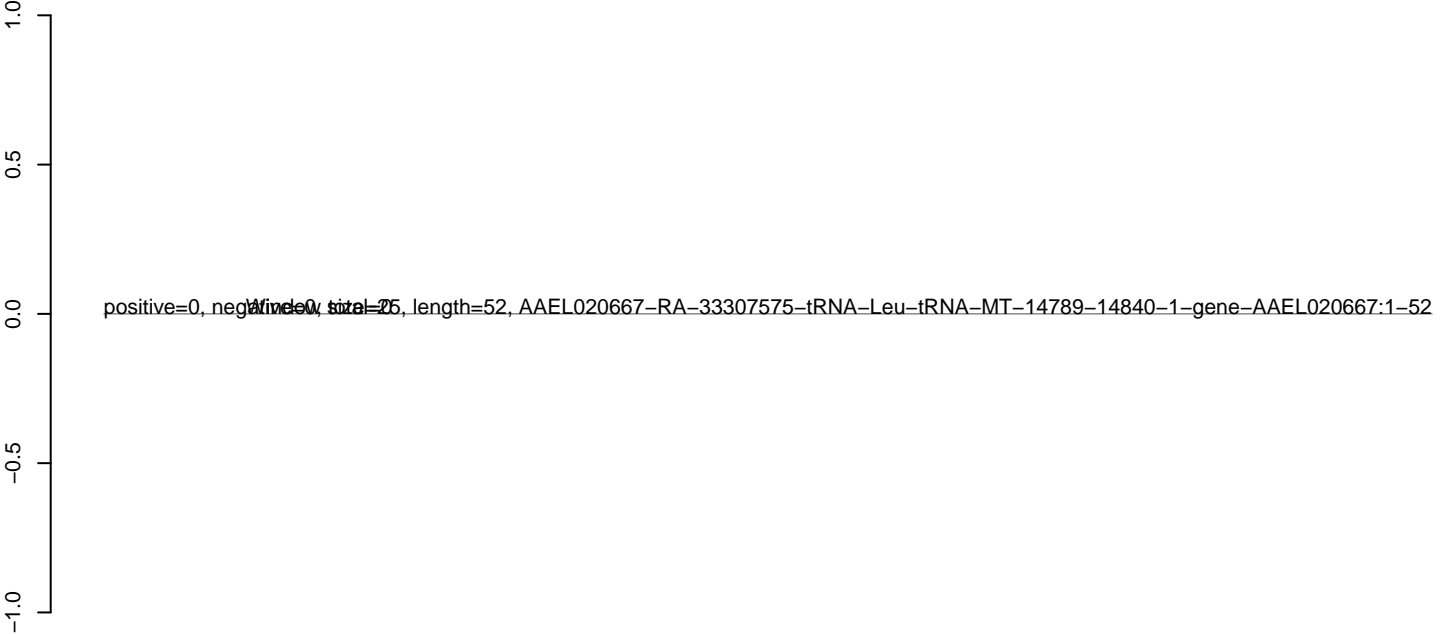
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AeAeg_Testes.24_35.rep



AeAeg_Testes.rep



AeAeg_Testes.18_23.rep



AeAeg_Testes.24_35.rep



AeAeg_Testes.rep

