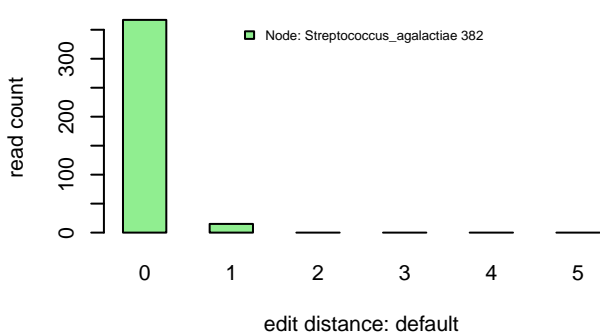
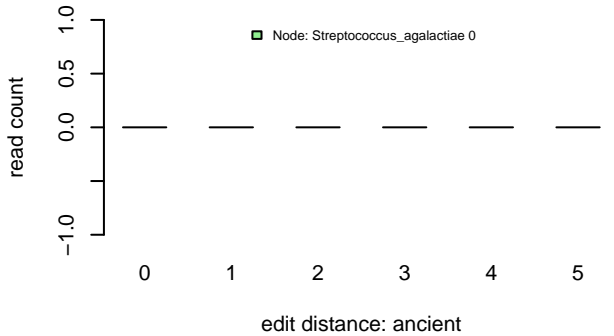


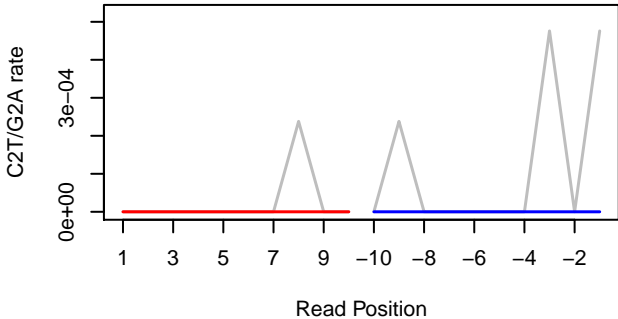
AT+Mock-1:50-3.rma6



AT+Mock-1:50-3.rma6



Damage plot for Streptococcus_agalactiae node



Node	Streptococcus_agalactiae
Top Reference	Streptococcus_agalactiae_2603V/R
assigned reads	508
all reads	508
nonDup	508
readDis	0.763
nonStacked	382
destacking?	On
C>T_1	0
G>A_-1	0
mean length (sd)	76 (1.636)

Streptococcus_agalactiae_2603V/R;_AE009948.1 100%
Streptococcus_agalactiae;_CP021867.1 092%
Streptococcus_agalactiae;_CP011329.1 091%
Streptococcus_agalactiae;_CP019979.1 089%
Streptococcus_agalactiae;_CP021868.1 075%
Streptococcus_agalactiae;_CP019978.1 075%
Streptococcus_agalactiae;_CP021862.1 074%
Streptococcus_agalactiae_A909;_CP000114.1 074%
Streptococcus_agalactiae_GD201008-001;_CP003810.1 074%
Streptococcus_agalactiae;_CP016501.1 074%