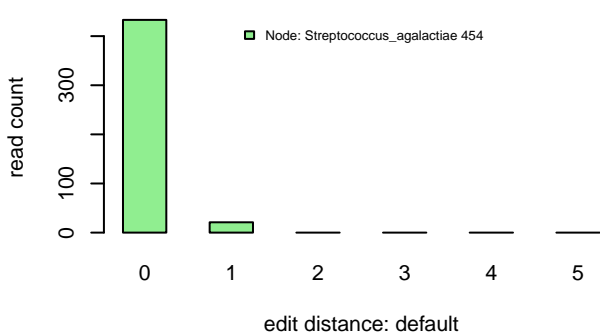
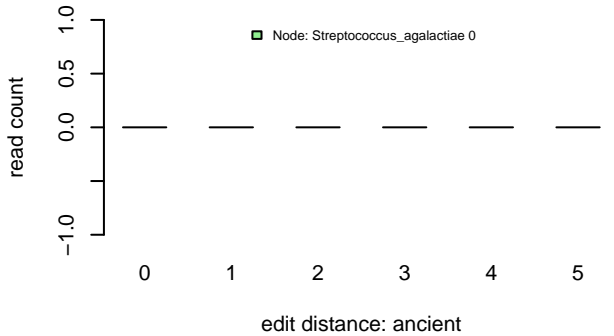


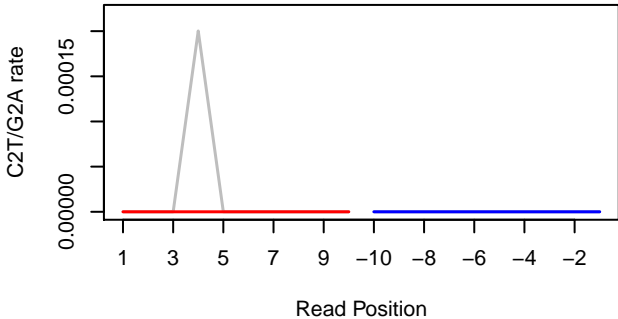
AT+Mock-1:50.rma6



AT+Mock-1:50.rma6



Damage plot for Streptococcus_agalactiae node



Node	Streptococcus_agalactiae
Top Reference	Streptococcus_agalactiae_2603V/R
assigned reads	661
all reads	658
nonDup	658
readDis	0.7
nonStacked	453
destacking?	On
C>T_1	0
G>A_-1	0
mean length (sd)	75 (2.502)

Streptococcus_agalactiae_2603V/R;_AE009948.1 100%
Streptococcus_agalactiae;_CP021867.1 092%
Streptococcus_agalactiae;_CP011329.1 090%
Streptococcus_agalactiae;_CP019979.1 089%
Streptococcus_agalactiae;_CP012503.1 075%
Streptococcus_agalactiae;_CP021868.1 073%
Streptococcus_agalactiae;_CP013202.1 072%
Streptococcus_agalactiae;_CP013908.1 071%
Streptococcus_agalactiae;_CP010867.1 071%
Streptococcus_agalactiae;_CP021862.1 071%