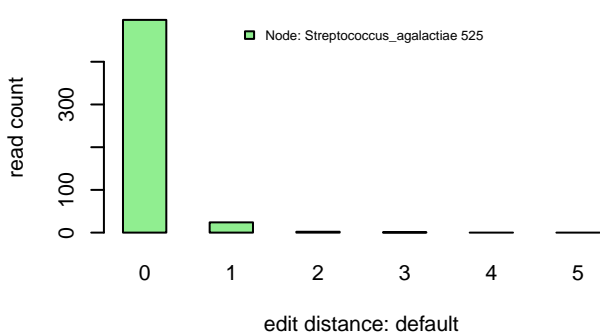
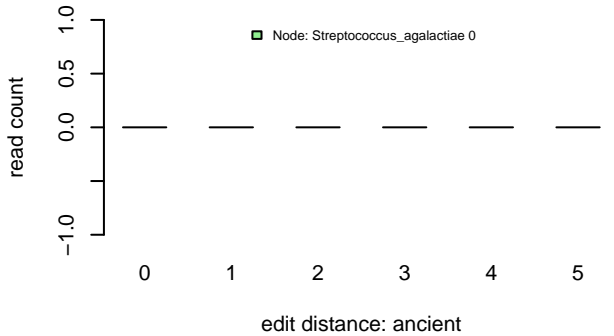


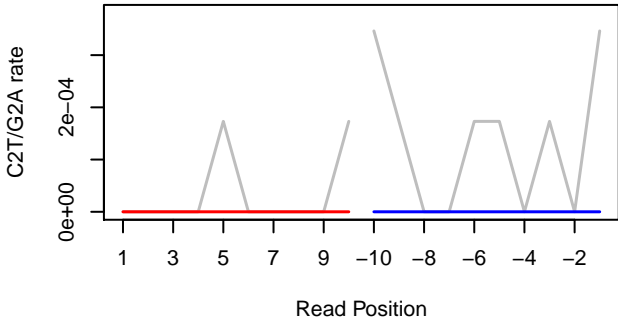
AT+Mock-1:50-1.rma6



AT+Mock-1:50-1.rma6



Damage plot for Streptococcus\_agalactiae node



Node	Streptococcus_agalactiae
Top Reference	Streptococcus_agalactiae_2603V/R
assigned reads	695
all reads	694
nonDup	694
readDis	0.773
nonStacked	524
destacking?	On
C>T_1	0
G>A_-1	0
mean length (sd)	76 (1.37)

Streptococcus_agalactiae_2603V/R;_AE009948.1 100%
Streptococcus_agalactiae;_CP021867.1 093%
Streptococcus_agalactiae;_CP011329.1 091%
Streptococcus_agalactiae;_CP019979.1 090%
Streptococcus_agalactiae;_CP021868.1 075%
Streptococcus_agalactiae;_CP012503.1 074%
Streptococcus_agalactiae;_CP013202.1 073%
Streptococcus_agalactiae_A909;_CP000114.1 072%
Streptococcus_agalactiae_09mas018883;_HF952104.1 072%
Streptococcus_agalactiae;_CP010867.1 072%