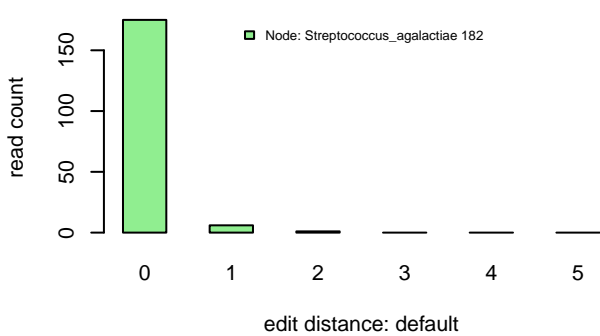
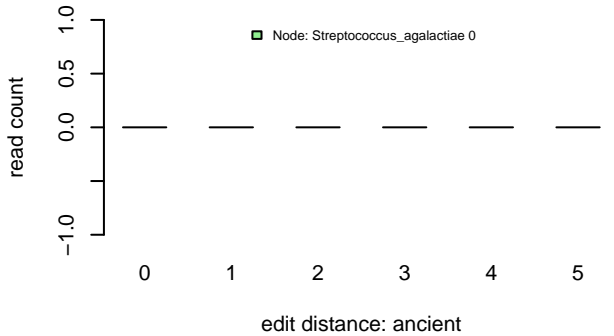


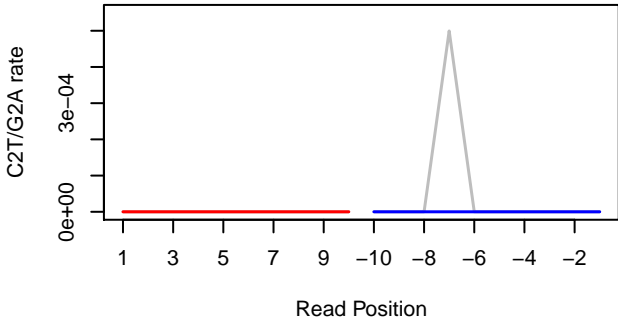
AT+Mock-1:100.rma6



AT+Mock-1:100.rma6



Damage plot for Streptococcus_agalactiae node



Node	Streptococcus_agalactiae
Top Reference	Streptococcus_agalactiae_2603V/R
assigned reads	253
all reads	252
nonDup	252
readDis	0.726
nonStacked	181
destacking?	On
C>T_1	0
G>A_-1	0
mean length (sd)	75 (1.84)

Streptococcus_agalactiae_2603V/R;_AE009948.1 100%
Streptococcus_agalactiae;_CP021867.1 091%
Streptococcus_agalactiae;_CP019979.1 088%
Streptococcus_agalactiae;_CP011329.1 087%
Streptococcus_agalactiae;_CP012503.1 076%
Streptococcus_agalactiae;_CP021868.1 075%
Streptococcus_agalactiae;_CP013908.1 074%
Streptococcus_agalactiae_NEM316;_AL732656.1 072%
Streptococcus_agalactiae;_CP021870.1 072%
Streptococcus_agalactiae;_CP019978.1 072%