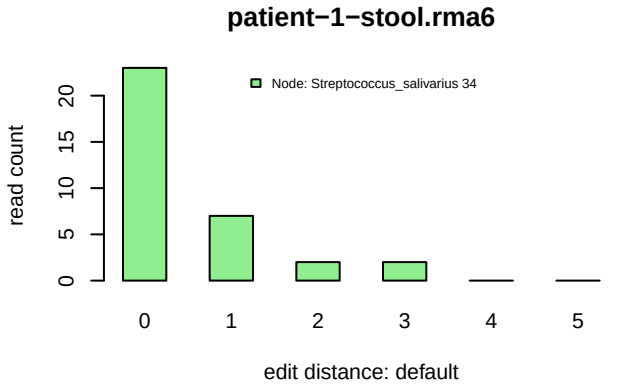
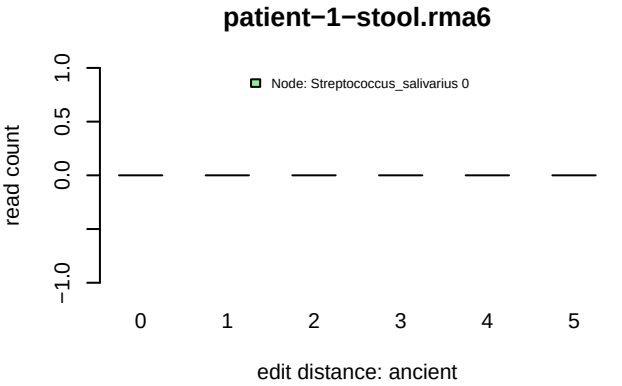
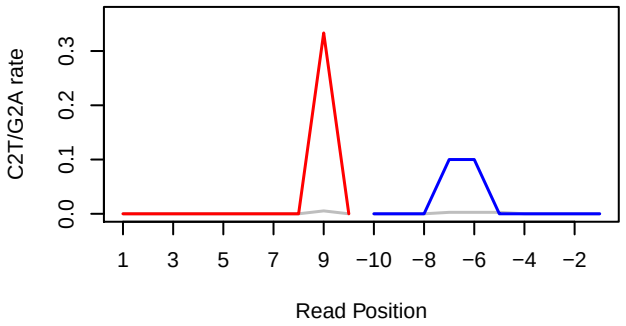




Damage plot for Streptococcus_salivarius node



Streptococcus_salivarius;_CP020451.1 100%
Streptococcus_salivarius_CCHSS3;_FR873481.1 059%
Streptococcus_salivarius_57.I;_CP002888.1 045%
Streptococcus_salivarius_JIM8777;_FR873482.1 040%
Streptococcus_equinus;_CP020438.1 013%
Streptococcus_sp._FDAARGOS_192;_CP020431.1 013%
Streptococcus_sp._HSISS1;_CM002132.1 013%
Streptococcus_equinus;_CP020439.1 004%
Streptococcus_sp._HSISS3;_CM002130.1 004%
NA%

Node	Streptococcus_salivarius
Top Reference	Streptococcus_salivarius
assigned reads	48
all reads	32
nonDup	32
readDis	0.695
nonStacked	22
destacking?	On
C>T_1	0
G>A_−1	0
mean length (sd)	76 (0.741)