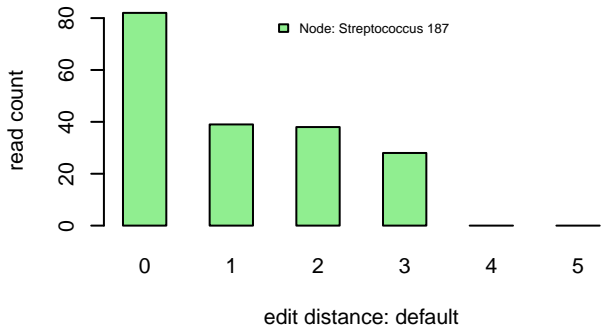
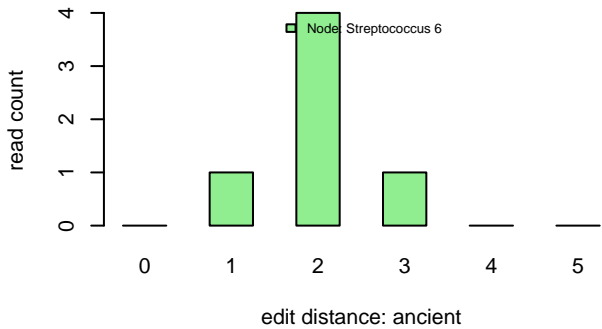


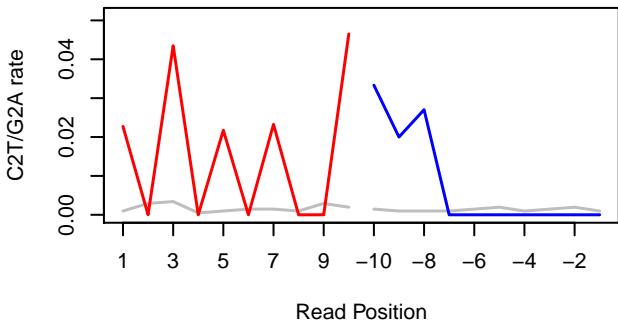
patient-1-stool.rma6



patient-1-stool.rma6



Damage plot for Streptococcus node



Streptococcus_anginosus;_CP007573.1	100%
Streptococcus_sp._HSISM1;_CM002128.1	100%
Streptococcus_parasanguinis_ATCC_15912;_CP002843.1	100%
Streptococcus_parasanguinis_FW213;_CP003122.1	100%
Streptococcus_agalactiae;_CP007570.1	084%
Streptococcus_salivarius;_CP013216.1	081%
Streptococcus_salivarius;_CP020451.1	081%
Streptococcus_sp._HSISS1;_CM002132.1	073%
Streptococcus_salivarius_JIM8777;_FR873482.1	068%
Streptococcus_equinus;_CP020438.1	060%

Node	Streptococcus
Top Reference	Streptococcus_anginosus
assigned reads	563
all reads	251
nonDup	251
readDis	0.264
nonStacked	38
destacking?	On
C>T_1	0.0227
G>A_-1	0
mean length (sd)	75 (5.018)