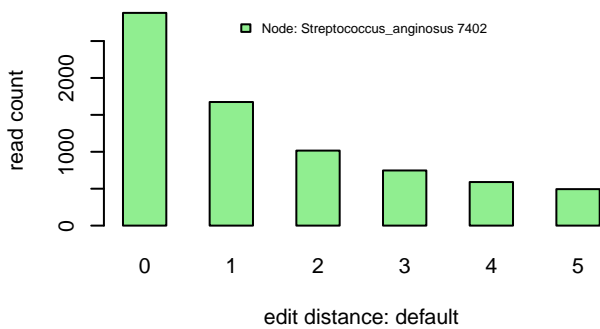
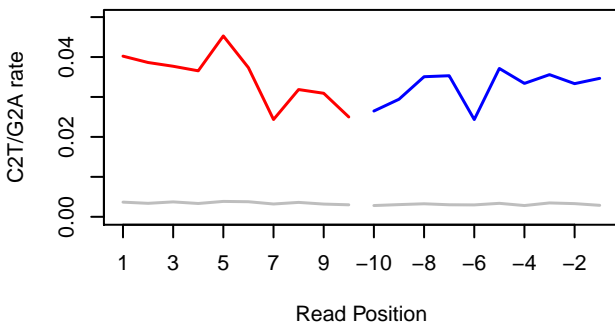


S0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractuS0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractu



### Damage plot for Streptococcus\_angerinosus node



Node	Streptococcus_angerinosus
Top Reference	Streptococcus_angerinosus
assigned reads	21214
all reads	4236
nonDup	4145
readDis	0.446
nonStacked	1363
destacking?	Off
C>T_1	0.0402
G>A_1	0.0347
mean length (sd)	107 (39.867)

Streptococcus_angerinosus;_CP012719.1 100%
Streptococcus_angerinosus_C1051;_CP003860.1 100%
Streptococcus_angerinosus_subsp._whileyi_MAS624;_AP013072.1 100%
Streptococcus_angerinosus_C238;_CP003861.1 100%
Streptococcus_angerinosus_SK52=_DSM_20563;_AB643814.1 001%
Streptococcus_sanguinis_SK36;_CP000387.1 001%
Streptococcus_intermedius_C270;_CP003858.1 000%
Streptococcus_angerinosus_SK52=_DSM_20563;_KY315441.1 000%
Streptococcus_oralis_subsp._tigurinus;_AP018338.1 000%
uncultured_bacterium;_KU606697.1 000%