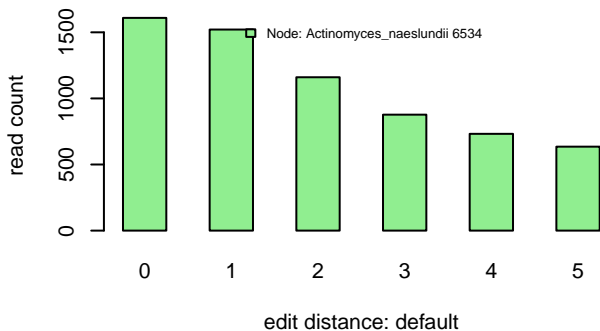
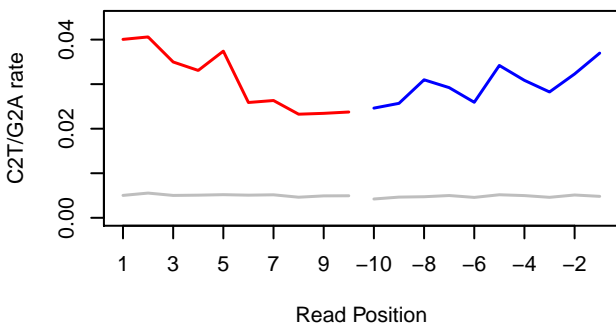


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Actinomyces_naeslundii node



Node	Actinomyces_naeslundii
Top Reference	Actinomyces_naeslundii
assigned reads	553219
all reads	9958
nonDup	9890
readDis	0.542
nonStacked	3496
destacking?	Off
C>T_1	0.0401
G>A_-1	0.037
mean length (sd)	92 (27.742)

Actinomyces_naeslundii;_AP017894.1	100%
Actinomyces_oris;_CP014232.1	0.000%
Actinomyces_sp._oral_taxon_414;_CP012590.1	0.000%
Serinicoccus_sp._JLT9;_CP014989.1	0.000%
Actinomyces_oris;_EU805700.1	0.000%
Actinomyces_oris;_EU805672.1	0.000%
Actinomyces_oris;_EU620964.1	0.000%
Actinomyces_oris;_EU805690.1	0.000%
Actinomyces_oris;_EU805653.1	0.000%
Actinomyces_oris;_EU621000.1	0.000%