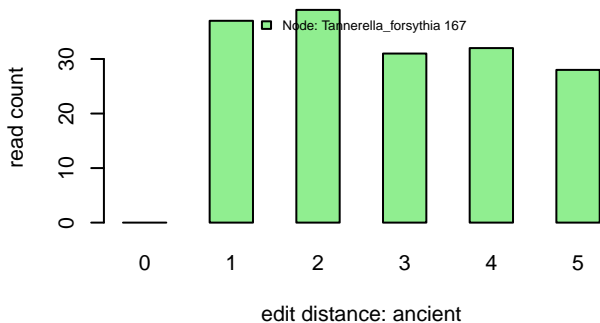
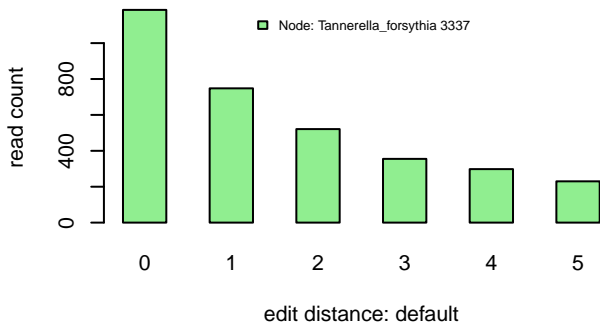
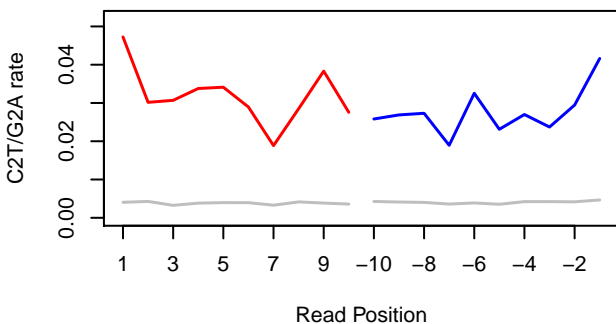


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Tannerella_forsythia node



Node	Tannerella_forsythia
Top Reference	Tannerella_forsythia_92A2
assigned reads	4418
all reads	2912
nonDup	2877
readDis	0.352
nonStacked	807
destacking?	Off
C>T_1	0.0473
G>A_-1	0.0416
mean length (sd)	82 (22.711)

Tannerella_forsythia_92A2;_CP003191.1 100%
Tannerella_forsythia_KS16;_AP013045.1 100%
Tannerella_forsythia_3313;_AP013044.1 100%
Tannerella_forsythia;_NR_076943.1 004%
Tannerella_sp._oral_taxon_HOT-286;_CP017038.1 001%
Tannerella_forsythia;_AB587729.1 001%
Tannerella_forsythia;_AJ006516.1 000%
uncultured_bacterium;_JQ999790.1 000%
Tannerella_forsythia;_AY423857.1 000%
Tannerella_forsythia;_AF054892.1 000%