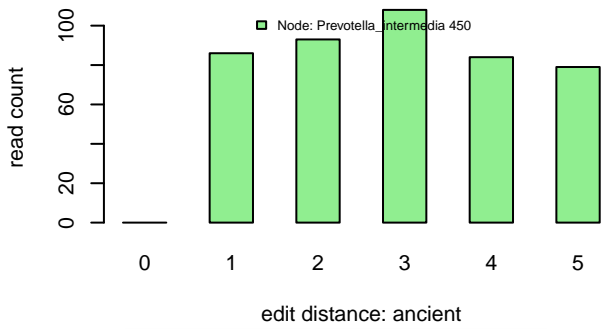
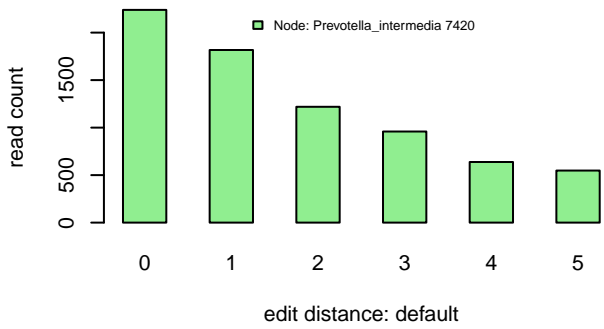
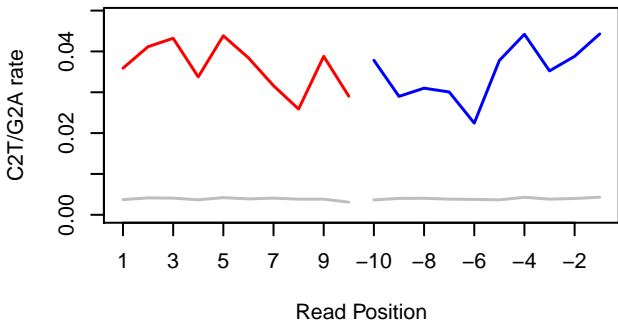


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Prevotella_intermedia node



Node	Prevotella_intermedia
Top Reference	Prevotella_intermedia
assigned reads	60567
all reads	5407
nonDup	5353
readDis	0.293
nonStacked	1365
destacking?	Off
C>T_1	0.0359
G>A_-1	0.0443
mean length (sd)	88 (28.257)

Prevotella_intermedia;_AP014597.1	100%
Prevotella_intermedia_17;_CP003503.1	100%
Prevotella_intermedia_ATCC_25611;_DSM_20706;_CP019300.1	100%
Prevotella_intermedia_17;_CP003502.1	042%
Prevotella_intermedia_ATCC_25611;_DSM_20706;_CP019301.1	035%
Porphyromonas_gingivalis;_CP011995.1	003%
Prevotella_dentalis_DSM_3688;_CP003369.1	002%
Tannerella_forsythia_KS16;_AP013045.1	002%
Bacteroides_fragilis_638R;_FQ312004.1	001%
Prevotella_fusca_JCM_17724;_CP012074.1	000%