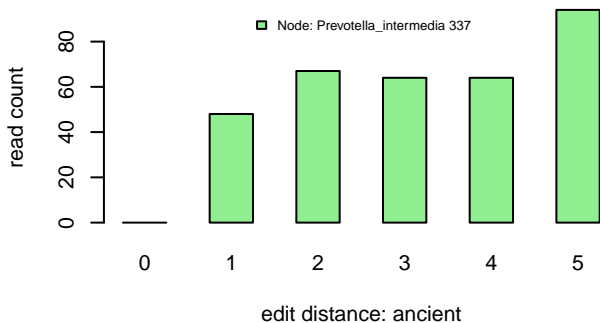
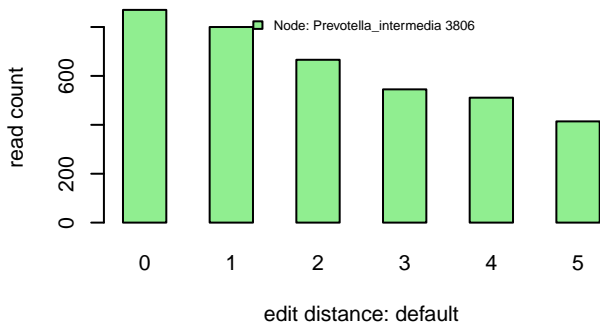
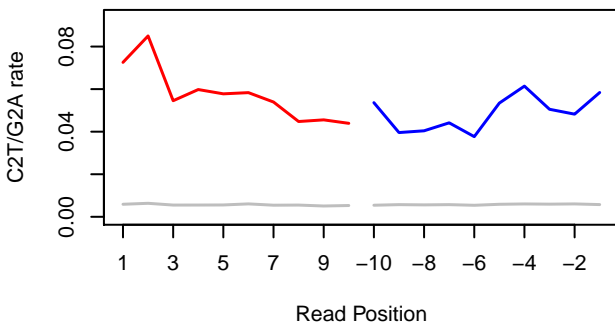


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Prevotella_intermedia node



Node	Prevotella_intermedia
Top Reference	Prevotella_intermedia
assigned reads	6031
all reads	2883
nonDup	2842
readDis	0.184
nonStacked	409
destacking?	Off
C>T_1	0.0726
G>A_-1	0.0584
mean length (sd)	83 (22.615)

Prevotella_intermedia;_AP014597.1 100%
Prevotella_intermedia_17;_CP003503.1 100%
Prevotella_intermedia_ATCC_25611_= _DSM_20706;_CP019300.1 100%
Prevotella_intermedia_ATCC_25611_= _DSM_20706;_CP019301.1 100%
Prevotella_intermedia_17;_CP003502.1 066%
Porphyromonas_gingivalis;_CP011995.1 032%
Tannerella_forsythia_KS16;_AP013045.1 004%
Bacteroides_fragilis_638R;_FQ312004.1 004%
Prevotella_fusca_JCM_17724;_CP012074.1 004%
Prevotella_dentalis_DSM_3688;_CP003369.1 003%