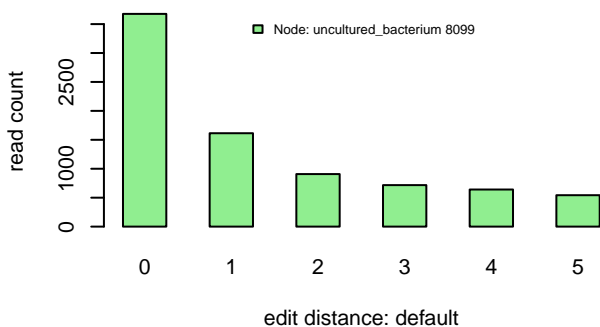
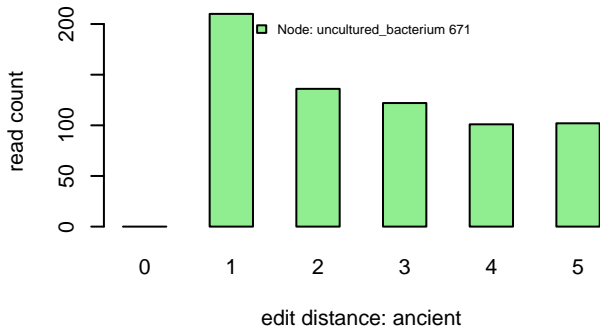


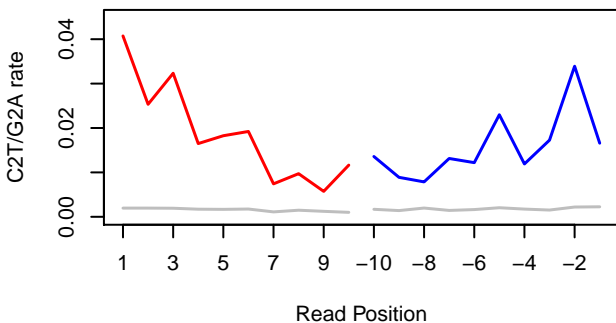
101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Damage plot for uncultured_bacterium node



uncultured_bacterium;_KX244620.1	100%
Comamonas_sp._01xTSA12B_G05;_HM113648.1	100%
Comamonas_sp._SJM;_HM140707.1	100%
Comamonas_sp._BM-9_6;_AY635871.1	100%
uncultured_anaerobic_bacterium;_AY953165.1	100%
bacterium_SRMC-52-8;_DQ104970.1	100%
Comamonas_sp._3R3-8;_GU195185.1	100%
uncultured_anaerobic_bacterium;_AY953184.1	100%
uncultured_anaerobic_bacterium;_AY953177.1	100%
Comamonas_sp._SS-1;_FJ426595.1	100%

edit distance: ancient	
Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	10863
all reads	563
nonDup	546
readDis	0.095
nonStacked	15
destacking?	Off
C>T_1	0.0407
G>A_-1	0.0166
mean length (sd)	67 (24.53)