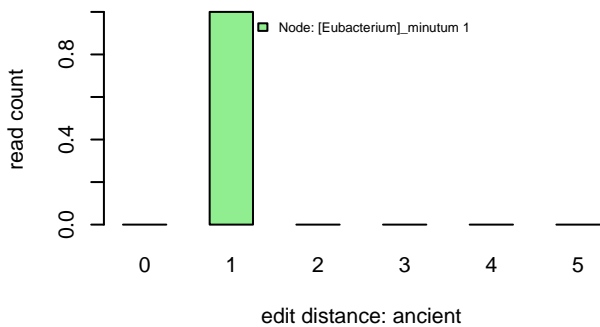
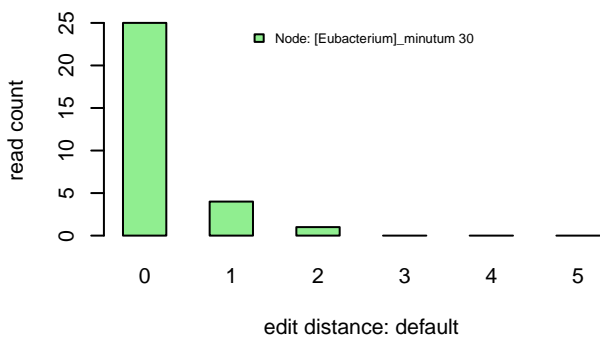
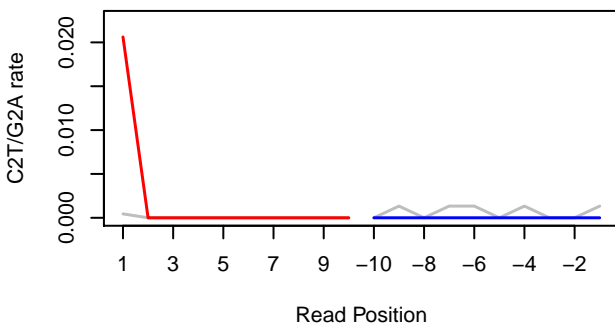


S0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractuS0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractu



### Damage plot for [Eubacterium]\_minutum node



Node	[Eubacterium]_minutum
Top Reference	[Eubacterium]_minutum
assigned reads	31
all reads	31
nonDup	30
readDis	0.039
nonStacked	1
destacking?	Off
C>T_1	NA
G>A_-1	NA
mean length (sd)	106 (50.666)

[Eubacterium]_minutum;_NR_036949.1 100%
Eubacterium_minutum_ATCC_700079;_NR_112052.1 100%
uncultured_Eubacterium_sp.;_AM420183.1 100%
Eubacterium_minutum_ATCC_700079;_CP016202.1 100%
Eubacterium_minutum_ATCC_700079;_NR_114689.1 100%
uncultured_bacterium;_KF506073.1 100%
uncultured_Clostridium_sp.;_EF704000.1 100%
Eubacterium_sp._oral_taxon_G75;_GU432190.1 100%
uncultured_Eubacterium_sp.;_AM419990.1 100%
NA%