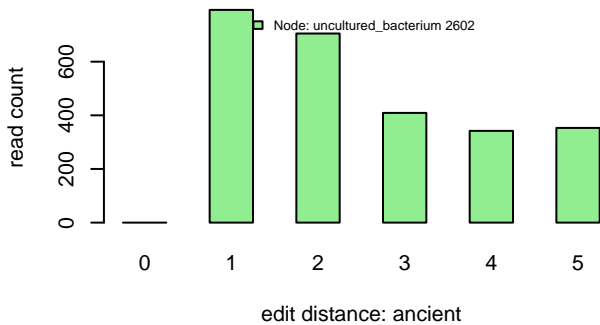
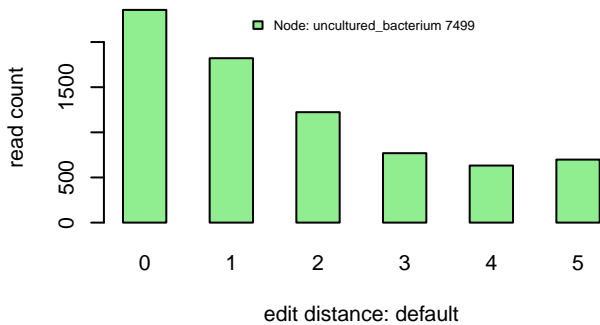
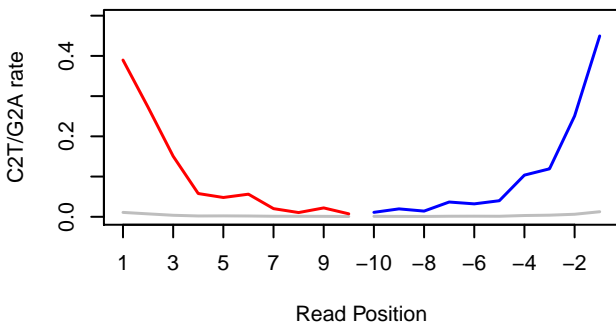


n2_S0_L000_R1_000.fastq.merged.prefixed.hg19unmn2_S0_L000_R1_000.fastq.merged.prefixed.hg19unmn



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	52832
all reads	915
nonDup	622
readDis	0
nonStacked	0
destacking?	Off
C>T_1	0.3897
G>A_-1	0.4496
mean length (sd)	63 (20.447)

uncultured_bacterium;_JQ696889.1 100%

Actinomyces_naeslundii;_JF496733.1 100%

[Eubacterium]_yurii_subsp._yurii;_NR_104843.1 100%

uncultured_Eubacterium_sp.;_AM420046.1 100%

[Eubacterium]_yurii_subsp._yurii;_L34629.2 100%

Peptostreptococcaceae_bacterium_OBRC9;_HQ616354.1 100%

Eubacterium_sp._oral_clone_OH3A;_AY947497.1 100%

TM7_phylum_sp._oral_clone_EW055;_AF385500.1 100%

[Eubacterium]_yurii_subsp._schtitka;_NR_115260.1 100%

Eubacterium_sp._oral_strain_A03MT;_AF287765.1 100%