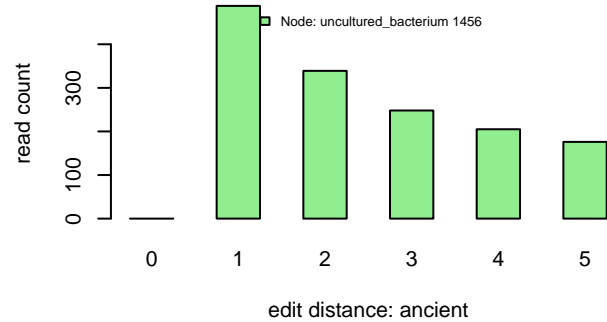
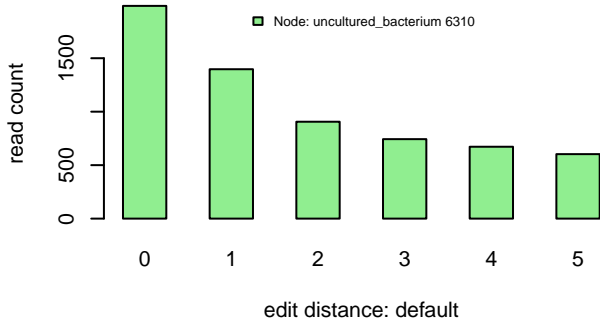
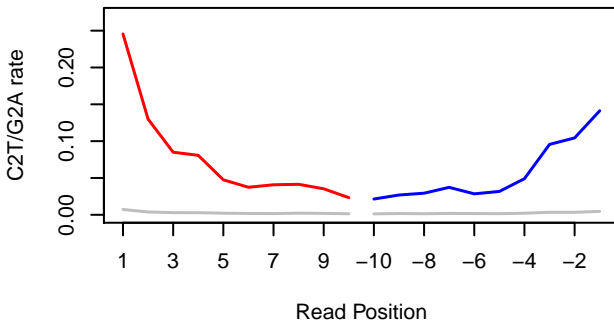


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	9229
all reads	486
nonDup	459
readDis	0.012
nonStacked	3
destacking?	Off
C>T_1	0.2456
G>A_-1	0.1412
mean length (sd)	66 (25.371)

uncultured_bacterium;_KF312324.1 100%
Abiotrophia_defectiva;_AY879307.1 100%
Abiotrophia_defectiva;_AY879306.1 100%
Abiotrophia_defectiva;_AY879308.1 100%
uncultured_Abiotrophia_sp.;_AM420170.1 100%
uncultured_Chlorobi_bacterium;_AJ272093.1 100%
uncultured_Abiotrophia_sp.;_FJ976282.1 100%
bacterium_str._86355;_AF227858.1 100%
Actinomyces_sp._oral_clone_GU009;_AY349361.1 100%
Actinomyces_odontolyticus;_GQ131411.1 100%