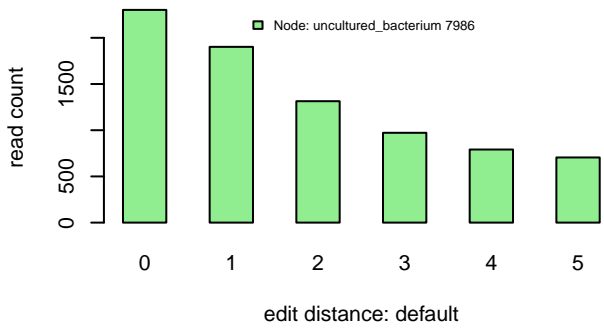
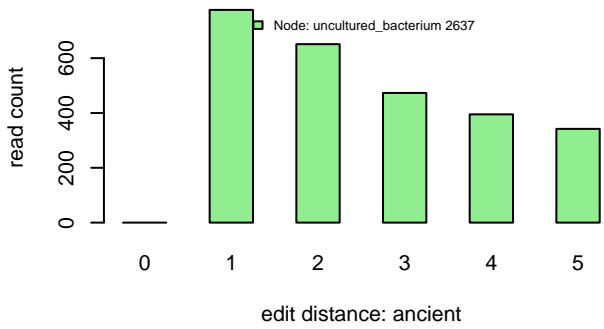


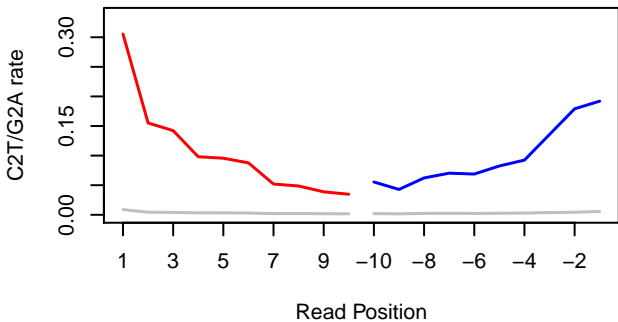
S0_L001_R1_001.fastq.combined.fq.prefixed.extractu



S0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for uncultured_bacterium node



uncultured_bacterium;_JQ696889.1	100%
Actinomyces_sp._oral_clone_GU009;_AY349361.1	100%
Actinomyces_odontolyticus;_AJ234054.1	100%
Actinomyces_sp.;_AJ234058.1	100%
Actinomyces_odontolyticus;_GQ131411.1	100%
Actinomyces_sp._oral_strain_C29KA;_AF287751.1	100%
Actinomyces_odontolyticus;_AJ234047.1	100%
uncultured_Actinomyces_sp.;_FJ976408.1	100%
uncultured_Candidatus_Saccharibacteria_bacterium;_FJ976257.1	100%
Actinomyces_sp._oral_clone_CT068;_AF385505.1	100%

Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	17909
all reads	432
nonDup	404
readDis	0.001
nonStacked	0
destacking?	Off
C>T_1	0.3056
G>A_-1	0.1921
mean length (sd)	56 (21.202)