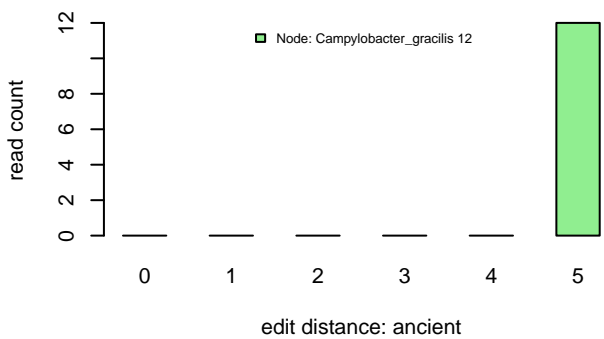
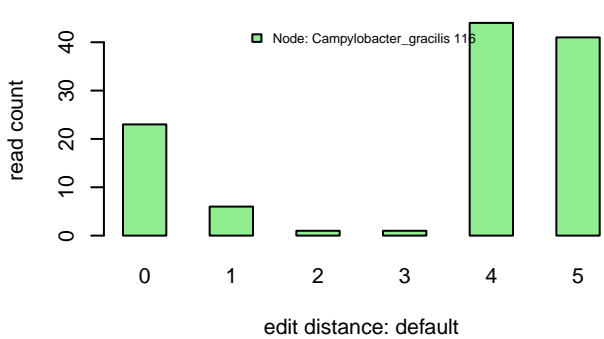
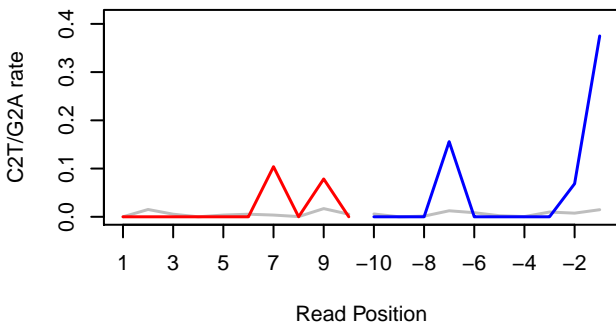


50_L001_R1_001.fastq.combined.fq.prefixed.extractu50_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for Campylobacter_gracilis node



Campylobacter_gracilis;_CP012196.1 100%
NA%
NA%
NA%
NA%
NA%
NA%
NA%
NA%
NA%
NA%

Node	Campylobacter_gracilis
Top Reference	Campylobacter_gracilis
assigned reads	224
all reads	224
nonDup	44
readDis	0.201
nonStacked	11
destacking?	Off
C>T_1	0
G>A_-1	0.375
mean length (sd)	83 (21.271)