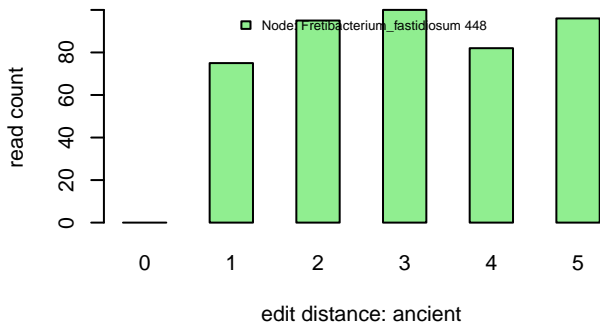
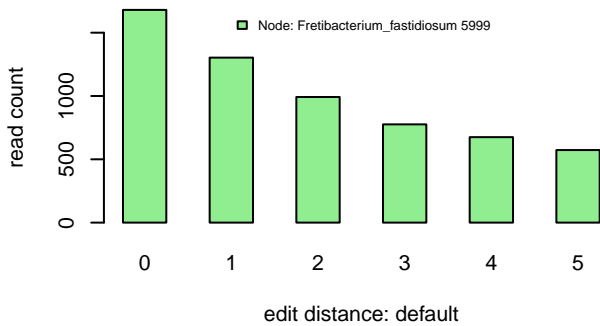
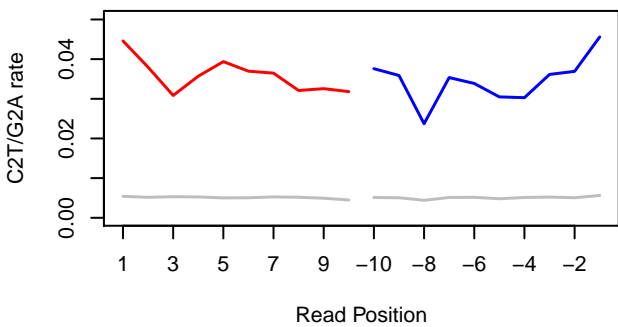


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for Fretibacterium_fastidiosum node



Fretibacterium_fastidiosum;_FP929056.1 100%
NA%
NA%
NA%
NA%
NA%
NA%
NA%
NA%
NA%

Node	Fretibacterium_fastidiosum
Top Reference	Fretibacterium_fastidiosum
assigned reads	101196
all reads	10000
nonDup	9849
readDis	0.535
nonStacked	3433
destacking?	Off
C>T_1	0.0446
G>A_-1	0.0456
mean length (sd)	106 (41.176)