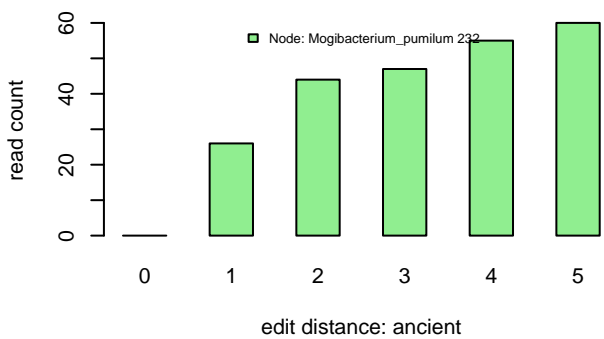
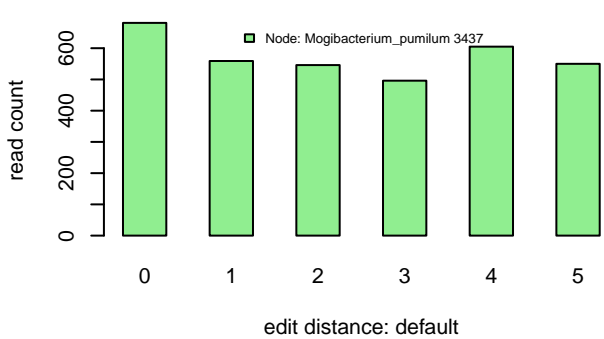
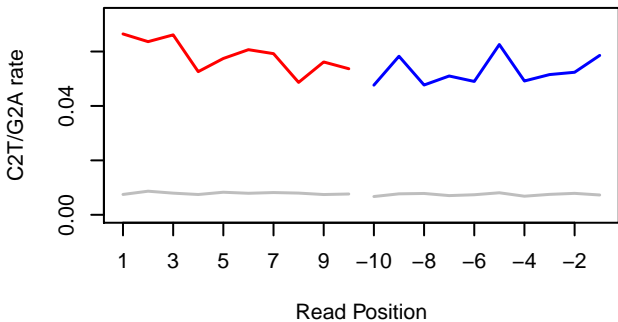


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for Mogibacterium_pumilum node



Mogibacterium_pumilum;_CP016199.1 100%

NA%

NA%

NA%

NA%

NA%

NA%

NA%

NA%

NA%

Node	Mogibacterium_pumilum
Top Reference	Mogibacterium_pumilum
assigned reads	45037
all reads	10000
nonDup	9856
readDis	0.087
nonStacked	502
destacking?	Off
C>T_1	0.0665
G>A_1	0.0586
mean length (sd)	108 (39.44)