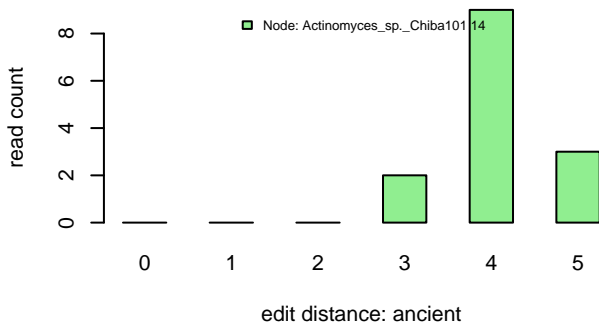
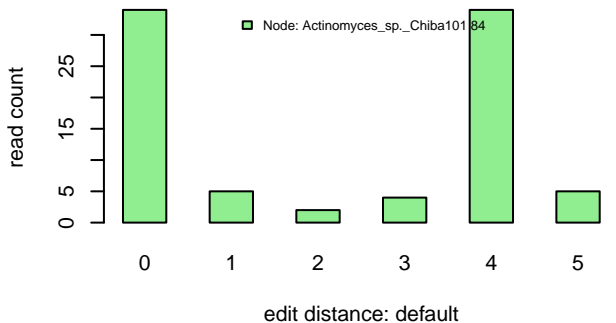
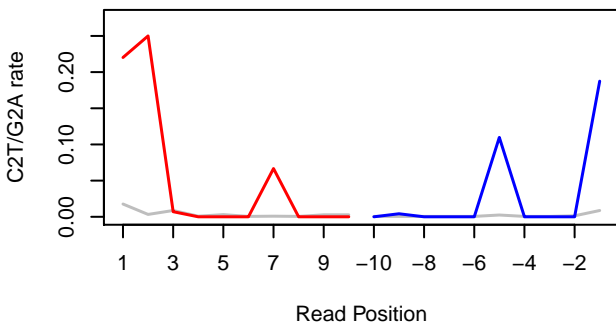


w\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19unmaw\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19unma



### Damage plot for Actinomyces\_sp.\_Chiba101 node



Actinomyces\_sp.\_Chiba101;\_AP017896.1 100%

NA%

NA%

NA%

NA%

NA%

NA%

NA%

NA%

NA%

Node	Actinomyces_sp._Chiba101
Top Reference	Actinomyces_sp._Chiba101
assigned reads	335
all reads	335
nonDup	93
readDis	0.086
nonStacked	16
destacking?	Off
C>T_1	0.2202
G>A_-1	0.1875
mean length (sd)	120 (47.354)