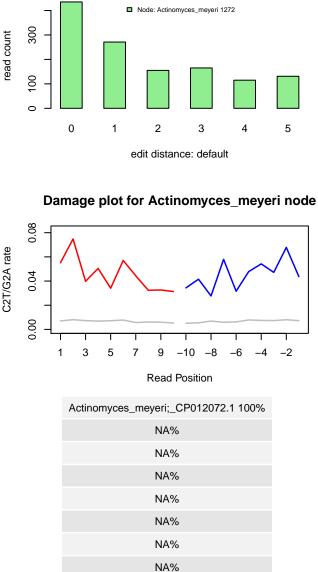
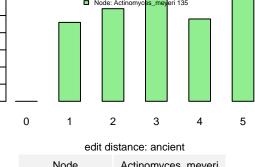
## 0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur

read count



NA% NA%



edit distance: ancient	
Node	Actinomyces_meyeri
Top Reference	Actinomyces_meyeri
assigned reads	1801
all reads	1799
nonDup	1714
readDis	0.662
nonStacked	1010
destacking?	Off
C>T_1	0.0552
G>A1	0.0437
mean length (sd)	55 (16.334)