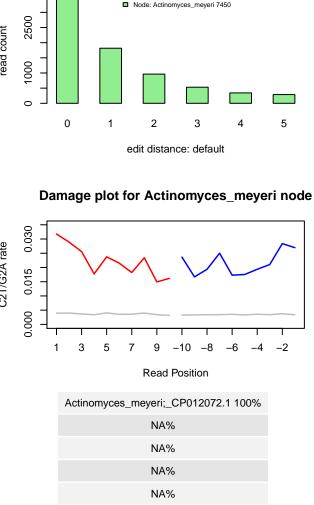
S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu

read count

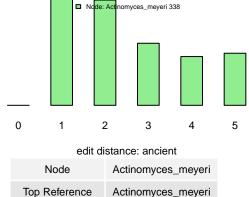
80

40 60

20



NA% NA% NA% NA%



edit distance: ancient	
Node	Actinomyces_meyeri
Top Reference	Actinomyces_meyeri
assigned reads	31824
all reads	10000
nonDup	9853
readDis	0.516
nonStacked	2979
destacking?	Off
C>T_1	0.0318
G>A1	0.027
mean length (sd)	107 (40.768)