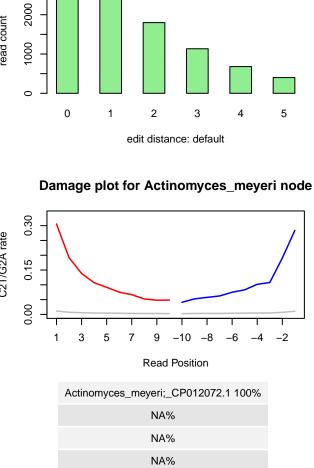
0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur

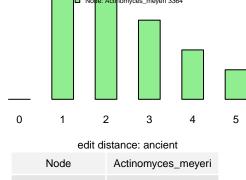
read count

900

200



NA%
NA%
NA%
NA%
NA%



edit distance: ancient	
Node	Actinomyces_meyeri
Top Reference	Actinomyces_meyeri
assigned reads	25573
all reads	10000
nonDup	9884
readDis	0.733
nonStacked	5562
destacking?	Off
C>T_1	0.3057
G>A1	0.2838
mean length (sd)	56 (21.838)