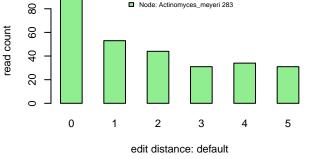
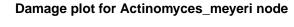
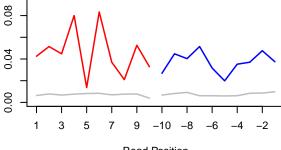
0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur

read count







C2T/G2A rate

Read Position
Actinomyces_meyeri;_CP012072.1 100%
NA%

	0	30			
0	1	2	3	4	5
	e	dit distan	ce: ancier	nt	

edit distance: ancient				
Node	Actinomyces_meyeri			
Top Reference	Actinomyces_meyeri			
assigned reads	451			
all reads	450			
nonDup	432			
readDis	0.894			
nonStacked	362			
destacking?	Off			
C>T_1	0.0426			
G>A1	0.0375			
mean length (sd)	59 (18.983)			