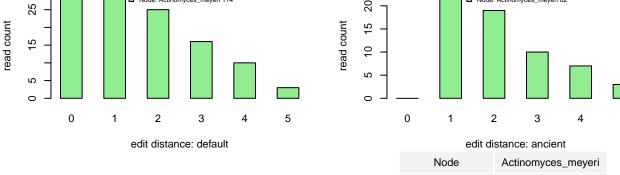
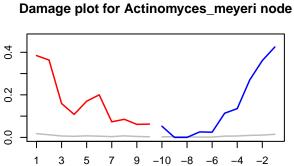
0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur





C2T/G2A rate

,	1	3	5	7	9	-10	-8	-6	-4		
	Read Position										
Actinomyces_meyeri;_CP012072.1 1000											
	NA% NA% NA%										
	NA% NA% NA%										
					NA	%					
				NA%							

0	1	2		3	4			
edit distance: ancient								
	Node		Actinomyces_meyeri					
Top Reference			Actinomyces_meyeri					
assigned reads			125					
all reads			125					
nonDup			120					
	readDis		0.961					
nonStacked			116					
de	estacking?		Off					
C>T_1			0.3846					
(G>A1		0.425					
mea	n length (s	d)	42 (12.137)					