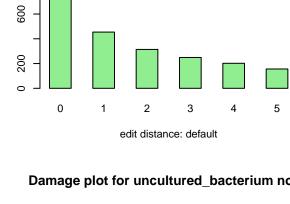
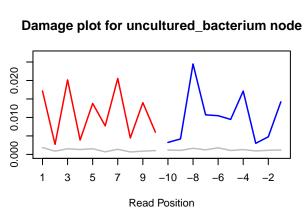
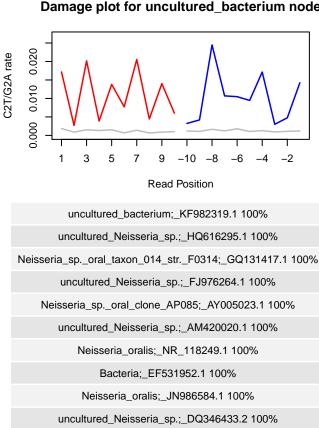
4 ■ Node: uncultured_bacterium 2234 Node: uncultured_bacterium 147 30 read count ead count 20

10

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







0	1	:	2	3	4
edit distance: ancient					
Node			uncultured_bacterium		
Top Reference			uncultured_bacterium		
assigned reads			2843		
all reads			128		
nonDup			116		
readDis			0.072		
nonStacked			7		
destacking?			Off		
C>T_1			0.0172		
G>A1			0.0142		
mean length (sd)			68 (24.619)		