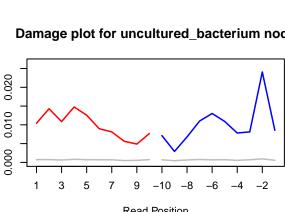
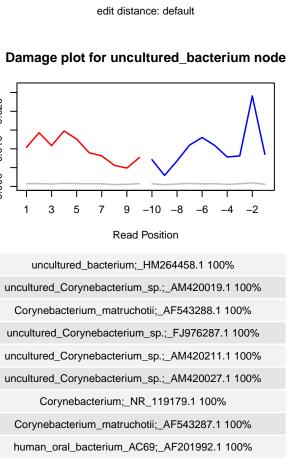
## 4000 ■ Node: uncultured\_bacterium 8055 Node: uncultured\_bacterium 473 ead count ead count 100 2000

0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un



0

C2T/G2A rate



Corynebacterium\_sp.\_oral\_clone\_AK153;\_AF287755.1 100%

0	1	:	2		3		4	
edit distance: ancient								
Node			uncultured_bacterium					m
Top Reference			uncultured_bacterium					m
assigned reads			68615					
all reads			535					
nonDup			529					
readDis			0.002					
nonStacked			1					
destacking?			Off					
C>T_1			0.0104					
G>A1			0.0085					
mean length (sd)			92 (28.444)					