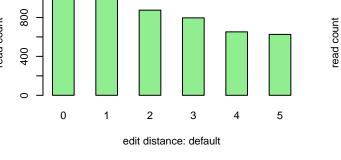
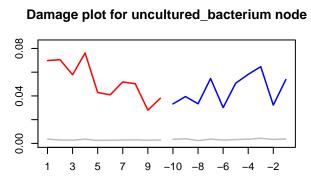
Node: uncultured_bacterium 5088 Node: uncultured_bacterium 628

100

0102_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0102_S0_L000_R1_000.fastq.merged.prefixed.hg19ur





C2T/G2A rate



Read Position

Bacteroidetes_bacterium_Oral_Taxon_274;_FJ577256.2 100%
Bacteroidetes_bacterium_oral_taxon_274;_GU409195.1 100%

Actinomyces_genomosp._C1;_AY278610.1 100% uncultured_Porphyromonadaceae_bacterium;_AM420225.1 100%

uncultured_Actinomyces_sp.;_FJ976431.1 100%

Actinomyces_sp._oral_taxon_414;_CP012590.1 100%

	0	1		2	3		4		5
	edit distance: ancient								
	Node			uncultured_bacterium					
	Top Re	uncultured_bacterium							
assigned reads				8132					
	all ı	486							
	noi	476							
	rea	0							
	nonS	0							

C>T_1 0.0698

G>A_-1 0.0538

mean length (sd) 80 (24.219)

Off

destacking?