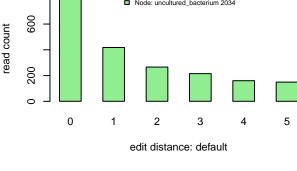
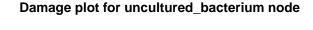
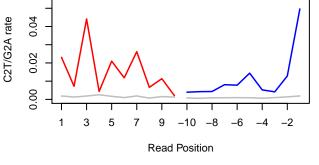
50 ■ Node: uncultured_bacterium 2034 Node: uncultured_bacterium 184

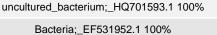
ead count 30

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









uncultured_Neisseria_sp.;_DQ346433.2 100%

Neisseria_sicca;_AJ239294.1 100%

Neisseria_sp._oral_clone_AP085;_AY005023.1 100%

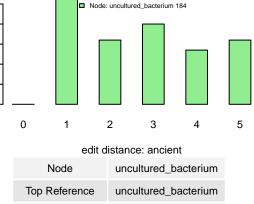
Neisseria_sp._oral_taxon_014_str._F0314;_GQ131417.1 100%

uncultured_Neisseria_sp.;_HQ616295.1 100%

uncultured_Neisseria_sp.;_FJ976264.1 100%

uncultured_Kingella_sp.;_GU227227.1 100%

Neisseria_sp._oral_clone_AP015;_AY005025.1 100%



0	1	2	3	4	
edit distance: ancient					
Node		uncultured_bacterium			
Top Reference		uncultured_bacterium			
assigned reads		2565			
all reads		75			
nonDup		72			
ı	readDis		0.023		
no	nonStacked		0		
de	destacking?		Off		
	C>T_1		0.0231		
(G>A1		0.0496		
mear	mean length (sd)		63 (23.747)		