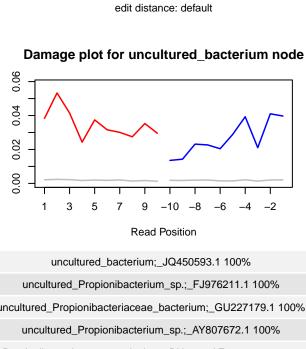


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



3

0

C2T/G2A rate uncultured_Propionibacteriaceae_bacterium;_GU227179.1 100% Propionibacterium_sp._oral_clone_BN085;_AF287757.1 100% Actinomyces_massiliensis_4401292;_NR_113059.1 100% Actinomyces_massiliensis_4401292;_NR_044288.1 100% uncultured_Propionibacterium_sp.;_AY807690.1 100% Bacteria;_FJ976437.1 100%

uncultured_human_oral_bacterium_A4;_AF201995.1 100%

0 2 3 edit distance: ancient uncultured_bacterium Node Top Reference uncultured_bacterium assigned reads 5199 all reads 149 nonDup 148 readDis 0 nonStacked 0 Off destacking? C>T_1 0.0383 G>A_-1 0.0397 mean length (sd) 78 (24.881)