0118_S0_L000_R1_000.fastq.merged.prefixed.hg19un)118_S0_L000_R1_000.fastq.merged.prefixed.hg19un 2.0 ■ Node: uncultured_bacterium 3 ■ Node: uncultured_bacterium 33 12 S read count read count 0.1 ω 0.5 0.0 0 3 0 1 2 3 5 edit distance: default edit distance: ancient Node uncultured_bacterium Top Reference uncultured_bacterium Damage plot for uncultured_bacterium node assigned reads 0.12 38 all reads 4 0.08 nonDup 1 0.04 readDis 1 nonStacked 1 0.00 destacking? Off 1 3 5 -10 C>T_1 0 G>A_-1 Read Position 0.1053 65 (27.421) mean length (sd) uncultured_bacterium;_KM657597.1 100% uncultured_actinobacterium;_AB288658.1 100% NA% NA% NA% NA% NA% NA% NA% NA%