S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu ■ Node: Anaerolineaceae_bacterium_oral_taxon_439 9211 lode: Anaerolineaceae_bacterium_oral_taxon_439 1787 9 2500 read count read count 400 1000 200 0 3 0 2 3 edit distance: default edit distance: ancient Node Anaerolineaceae_bacterium_oral_taxon_439 Top Reference Anaerolineaceae_bacterium_oral_taxon_439 age plot for Anaerolineaceae_bacterium_oral_taxon_4 assigned reads 99631 0.20 all reads 10000 C2T/G2A rate nonDup 9865 0.10 readDis 0.781 nonStacked 6151 0.00 destacking? Off 1 3 5 C>T_1 0.1789 G>A_-1 0.1809 Read Position mean length (sd) 60 (24.008) Anaerolineaceae_bacterium_oral_taxon_439;_CP017039.1 100% NA% NA% NA% NA% NA% NA% NA% NA%

NA%