S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu 100 ■ Node: Anaerolineaceae_bacterium_oral_taxon_439 312 ■ Node: Anaerolineaceae_bacterium_oral_taxon_439 88 read count read count 9 20 Ω 0 2 3 0 1 2 3 5 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 376 all reads 376 0.20 C2T/G2A rate nonDup 93 0.10 readDis 0.349 nonStacked 33 0.00 destacking? Off 1 3 5 7 -10 -8 -6 C>T_1 0.1148 0.1429 Read Position G>A_-1 mean length (sd) 77 (22.602) Anaerolineaceae_bacterium_oral_taxon_439;_CP017039.1 100% NA% NA% NA% NA% NA% NA% NA% NA%

NA%