## S0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractuS0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractu 150 ■ Node: Anaerolineaceae\_bacterium\_oral\_taxon\_439 9097 naerolineaceae\_bacterium\_oral\_taxon\_439 473 3000 read count read count 100 1000 50 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 95773 0.030 all reads 10000 C2T/G2A rate nonDup 9858 0.015 readDis 0.677 nonStacked 4696 destacking? Off 1 3 5 -2 C>T\_1 0.0327 G>A\_-1 0.03 Read Position 100 (39.625) mean length (sd) Anaerolineaceae\_bacterium\_oral\_taxon\_439;\_CP017039.1 100% NA% NA% NA% NA% NA% NA% NA% NA%

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