S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu 80 ■ Node: Adtinomyces_sp._oral_taxon_414 303 ■ Node: Actinomyces_sp._oral_taxon_414 7609 2000 9 read count 40 1000 20 0 0 2 3 0 1 2 3 edit distance: default edit distance: ancient Node Actinomyces_sp._oral_taxon_414 Top Reference Actinomyces_sp._oral_taxon_414 Damage plot for Actinomyces_sp._oral_taxon_414 no assigned reads 404946 all reads 10000 C2T/G2A rate nonDup 9899 0.015 readDis 0.729 nonStacked 5430 destacking? Off 1 3 5 -2 C>T_1 0.0255 G>A_-1 Read Position 0.021 93 (38.967) mean length (sd) Actinomyces_sp._oral_taxon_414;_CP012590.1 100% NA% NA% NA% NA% NA%

NA% NA% NA%