S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu ■ Node: Actinomyces_hongkor gensis 309 1500 ■ Node: Actinomyces_hongkongensis 5827 9 read count 40 500 20 0 2 3 0 2 3 5 edit distance: default edit distance: ancient Node Actinomyces_hongkongensis Top Reference Actinomyces_hongkongensis Damage plot for Actinomyces_hongkongensis nod assigned reads 10351 0.04 all reads 10000 C2T/G2A rate nonDup 9785 0.02 readDis 0.39 nonStacked 2434 0.00 destacking? Off 1 3 5 -2 C>T_1 0.034 0.0333 Read Position G>A_-1 106 (41.73) mean length (sd) Actinomyces_hongkongensis;_CP017298.1 100% NA% NA% NA% NA% NA% NA% NA% NA%

NA%