S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu 100 ■ Node: Actinomyc Node: Actinomyces_meyeri 4037 read count 9 read count 9 200 20 0 2 3 5 0 2 3 5 edit distance: default edit distance: ancient Node Actinomyces_meyeri Top Reference Actinomyces_meyeri Damage plot for Actinomyces_meyeri node assigned reads 33055 all reads 10000 C2T/G2A rate nonDup 9818 0.04 readDis 0.367 nonStacked 2227 0.00 Off destacking? -2 1 3 5 7 C>T_1 0.0685 G>A_-1 0.041 Read Position mean length (sd) 112 (40.996) Actinomyces_meyeri;_CP012072.1 100% NA% NA% NA% NA% NA% NA% NA% NA% NA%