0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un ■ Node: Actinomyces_oris 259 ■ Node: Actinomyces_oris 2704 9 009 read count ead count 40 20 200 0 2 3 0 2 3 5 edit distance: default edit distance: ancient Node Actinomyces_oris Top Reference Actinomyces_oris Damage plot for Actinomyces_oris node assigned reads 3060 all reads 3034 0.08 nonDup 2085 0.04 readDis 0.723 nonStacked 1478 0.00 destacking? Off 1 3 5 -2 C>T_1 0.0807 Read Position G>A_-1 0.095 60 (25.185) mean length (sd) Actinomyces_oris;_CP014232.1 100% Actinomyces_naeslundii;_AP017894.1 000% Actinomyces_oris_K20;_AB573870.1 000% Actinomyces_naeslundii;_DQ658412.1 000% Actinomyces_naeslundii;_AF107019.1 000% Actinomyces_viscosus;_M32067.1 000% Actinomyces_viscosus;_EU621125.1 000% Actinomyces_odontolyticus;_EU647590.1 000%

NA%