0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un Node: Parvimonas_micra 272 ■ Node: Parvimonas_micra 8594 3000 read count ead count 1000 20 0 3 0 2 3 edit distance: default edit distance: ancient Node Parvimonas_micra Top Reference Parvimonas_micra Damage plot for Parvimonas_micra node assigned reads 107883 0.04 all reads 9991 C2T/G2A rate nonDup 9893 0.02 readDis 0.454 nonStacked 2611 0.00 destacking? Off 1 3 5 C>T_1 0.021 G>A_-1 0.0323 Read Position mean length (sd) 86 (27.338) Parvimonas_micra;_CP009761.1 100% uncultured_Peptostreptococcus_sp.;_FJ976210.1 000% uncultured_Peptostreptococcus_sp.;_FJ268912.1 000% uncultured_Clostridium_sp.;_EF704838.1 000% uncultured_Clostridium_sp.;_EF704901.1 000% uncultured_Clostridium_sp.;_EF704816.1 000%

uncultured_Clostridium_sp.;_EF704826.1 000% uncultured_Clostridium_sp.;_EF704824.1 000% uncultured_Clostridium_sp.;_EF703937.1 000% uncultured_Clostridium_sp.;_EF698680.1 000% 5