0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un 009 Node: uncultured_bacterium 1919 ■ Node: uncultured_bacterium 6455 1500 read count read count 400 200 500 0 2 3 0 1 2 3 edit distance: default edit distance: ancient uncultured_bacterium Node Top Reference uncultured_bacterium Damage plot for uncultured_bacterium node assigned reads 10531 9.4 all reads 242 C2T/G2A rate nonDup 180 0.2 readDis 0.001 nonStacked 0 destacking? Off -2 1 3 5 7 C>T_1 0.3874 G>A_-1 0.3682 Read Position mean length (sd) 71 (25.903) uncultured_bacterium;_JQ696889.1 100%

unoundiou_buotonum;_o @ooooo.1 10070
Ottowia_sporal_taxon_894;_CP012073.1 100%
Bacteria;_FJ976437.1 100%
uncultured_Comamonadaceae_bacterium;_JQ406530.1 100%
uncultured_Lautropia_sp.;_AM419986.1 100%
Lautropia_sporal_clone_FX006;_AY134902.1 100%
Atopobium_parvulum_DSM_20469;_CP001721.1 100%
Abiotrophia_defectiva;_AY879307.1 100%
Abiotrophia_defectiva;_AY879306.1 100%
uncultured_Gemella_sp.;_EU071466.1 100%