S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu 3000 100 ■ Node: uncultured_bacterium 6614 ■ Node: uncultured_bacterium 360 2000 read count ead count 9 1000 20 0 0 2 3 edit distance: default edit distance: ancient Node uncultured_bacterium Top Reference uncultured_bacterium Damage plot for uncultured_bacterium node assigned reads 44537 0.020 all reads 527 C2T/G2A rate nonDup 515 0.010 readDis 0 nonStacked 0 destacking? Off 1 3 5 -2 C>T_1 0.0102 G>A_-1 0.0073 Read Position mean length (sd) 121 (41.336) uncultured_bacterium;_HM288087.1 100% unidentified_proteobacterium;_X73223.1 100% uncultured_Lautropia_sp.;_DQ016725.2 100% Lautropia_sp._TeTO;_FJ823153.1 100% Bacteria;_GQ073157.1 100% uncultured_Lautropia_sp.;_AM420184.1 100% Bacteria;_FJ976437.1 100% Lautropia_mirabilis;_GU397891.1 100% uncultured_Lautropia_sp.;_AY807005.1 100%

Lautropia_mirabilis;_GU397890.1 100%

5