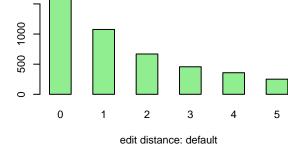
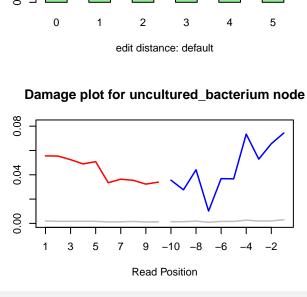
Node: uncultured_bacterium 4576 120 1000

50

0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur







C2T/G2A rate

uncultured_Comamonadaceae_bacterium;_JQ406530.1 100%

Lautropia_sp._oral_clone_FX006;_AY134902.1 100%

Ottowia_sp._oral_taxon_894;_CP012073.1 100%

uncultured_Lautropia_sp.;_AM419986.1 100%

bacterium;_MF801248.1 100%
uncultured_Brachymonas_sp.;_KR852686.1 100%

Pseudopropionibacterium_propionicum_F0230a;_CP002734.1 100%

uncultured_Lautropia_sp.;_AY806314.1 100%

