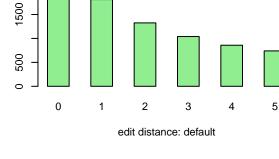
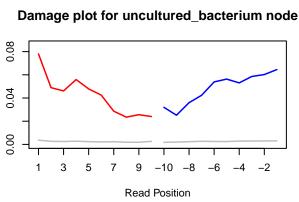


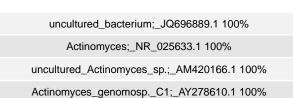
100

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





C2T/G2A rate

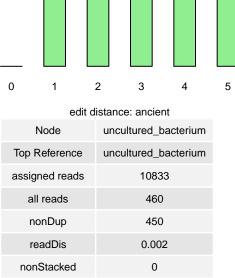




nomyces_genomospP1_oral_clone_MB6_C03;_DQ003632.1 100%
uncultured_Actinomyces_sp.;_FJ976446.1 100%

uncultured_Clostridiales_bacterium;_DQ144096.1 100%
uncultured_Bacteroidales_bacterium;_FJ976400.1 100%

uncultured_Clostridiales_bacterium;_EF012246.1 100%



Off

0.078

0.0645 55 (19.072)

destacking?

C>T_1

G>A_-1

mean length (sd)