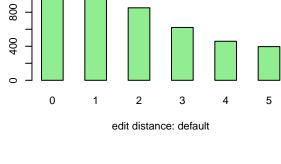
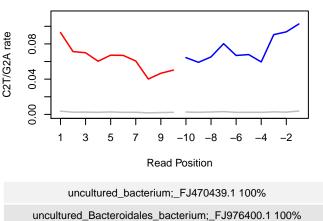


50

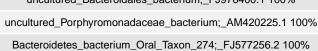
0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur





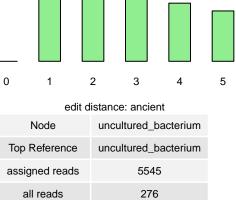


Damage plot for uncultured_bacterium node



Bacteroidetes_bacterium_oral_taxon_274;_GU409196.1 100%

Bacteroidetes_bacterium_oral_taxon_274;_GU409198.1 100% Bacteroidetes_bacterium_oral_taxon_274;_GU409193.1 100% Bacteroidetes_bacterium_oral_taxon_274;_GU409197.1 100% Bacteroides-like_sp._oral_clone_AU126;_AY005072.1 100% uncultured_Porphyromonas_sp.;_AM942563.1 100%



all reads	276
nonDup	268
readDis	0.005
nonStacked	0
destacking?	Off
C>T_1	0.093
G>A1	0.1026
mean length (sd)	52 (17.764)