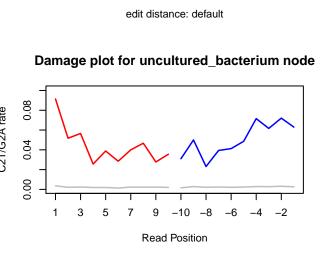


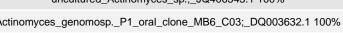
0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



3

0

uncultured\_bacterium;\_KM506769.1 100% Actinomyces;\_NR\_025633.1 100% uncultured\_Actinomyces\_sp.;\_AM420166.1 100% uncultured\_Actinomyces\_sp.;\_JQ406543.1 100%



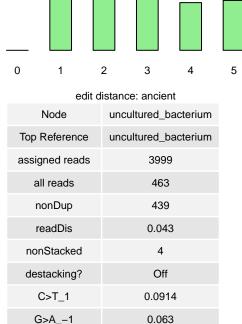
Actinomyces\_genomosp.\_C1;\_AY278610.1 100%

uncultured\_Bacteroidales\_bacterium;\_FJ976400.1 100%

Bacteroidetes\_bacterium\_Oral\_Taxon\_274;\_FJ577256.2 100%

uncultured\_Porphyromonadaceae\_bacterium;\_AM420225.1 100%

Pseudopropionibacterium\_propionicum\_F0230a;\_CP002734.1 100%



52 (17.705)

mean length (sd)