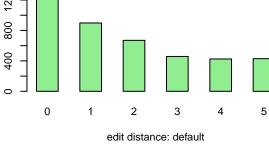
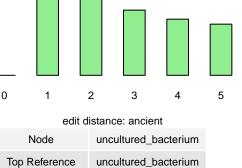
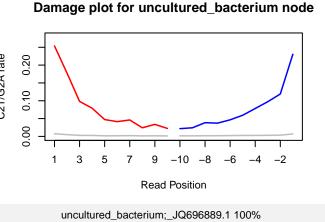
1200 ■ Node: uncultured_bacterium 4278 Node: uncultured_bacterium 1189 read count ead count 200

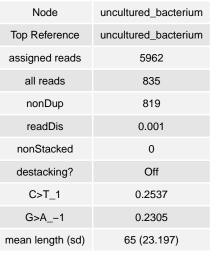
100

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









Clostridiales_bacterium_oral_taxon_G74;_GU432185.1 100% actinomyces_genomosp._P1_oral_clone_MB6_C03;_DQ003632.1 100%

uncultured_Actinomyces_sp.;_FJ976431.1 100%

uncultured_Actinomyces_sp.;_JQ406543.1 100%

ostreptococcaceae_bacterium_canine_oral_taxon_030;_JN713194.1 100%

uncultured_Actinomyces_sp.;_AM420166.1 100%

Actinomyces;_NR_025633.1 100%

tostreptococcaceae_bacterium_feline_oral_taxon_028;_KM461989.1 100%

Actinomyces_genomosp._C1;_AY278610.1 100%