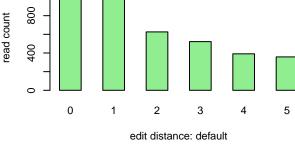
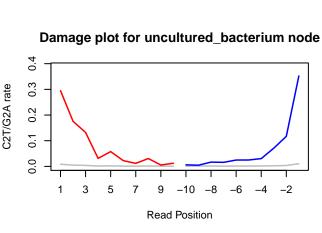
Node: uncultured_bacterium 1115 Node: uncultured_bacterium 4112 300 read count

100

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





uncultured_bacterium;_EU059897.1 100%

ncultured_Candidatus_Saccharibacteria_bacterium;_AM420132.1 100%	
ncultured_Candidatus_Saccharibacteria_bacterium;_FJ976255.1 100%	
ncultured_Candidatus_Saccharibacteria_bacterium;_AY144358.1 100%	

ncultured_Candidatus_Saccharibacteria_bacterium;_HQ616339.1 100% actinomyces_genomosp._P1_oral_clone_MB6_C03;_DQ003632.1 100% uncultured_Actinomyces_sp.;_JQ406543.1 100% ncultured_Candidatus_Saccharibacteria_bacterium;_GU227159.1 100% ncultured_Candidatus_Saccharibacteria_bacterium;_HM215448.1 100% uncultured_Actinomyces_sp.;_FJ976431.1 100%

