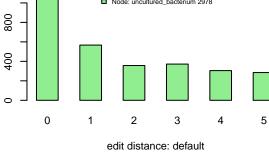
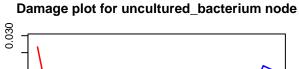
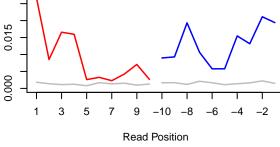
Node: uncultured_bacterium 2978 Node: uncultured_bacterium 224

read count 30

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

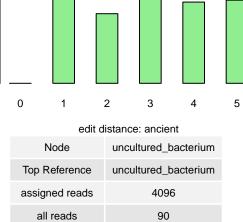






C2T/G2A rate

Read Position
uncultured_bacterium;_JF229185.1 100%
Acidithiobacillus_thiooxidans;_AF359941.1 100%
unidentified_proteobacterium;_X73223.1 100%
uncultured_Lautropia_sp.;_HQ616276.1 100%
Haemophilus_simiae;_DQ851143.1 100%
uncultured_Haemophilus_sp.;_AM420165.1 100%
Eikenella_corrodens;_NR_104685.1 100%
Lautropia_mirabilis;_GU397891.1 100%
Neisseria_canis;_AY426973.1 100%
Lautropia_mirabilis;_GU397890.1 100%



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	4096
all reads	90
nonDup	83
readDis	0.014
nonStacked	1
destacking?	Off
C>T_1	0.0267
G>A1	0.0194
mean length (sd)	66 (20.514)