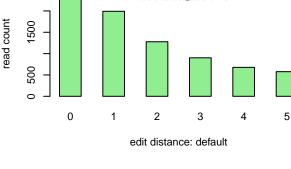
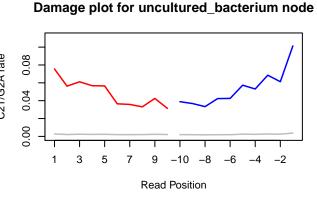
## ■ Node: uncultured\_bacterium 7917 Node: uncultured\_bacterium 1305 ead count

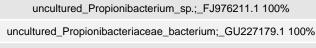
200

100

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







uncultured\_bacterium;\_JQ450593.1 100%

uncultured\_Propionibacterium\_sp.;\_AY807672.1 100% Leptotrichia\_goodfellowii;\_FJ577259.2 100%

Propionibacterium\_sp.\_oral\_clone\_BN085;\_AF287757.1 100% Leptotrichia\_goodfellowii;\_NR\_113160.1 100%

uncultured\_Propionibacterium\_sp.;\_AY807690.1 100% uncultured\_Leptotrichia\_sp.;\_FJ976432.1 100%

Leptotrichia\_goodfellowii\_DSM\_19756;\_NR\_025649.1 100%

0 2 3 edit distance: ancient

Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	10796
all reads	231
nonDup	228
readDis	0.01
nonStacked	1
destacking?	Off
C>T_1	0.0755
G>A1	0.1013

mean length (sd)

73 (27.025)