## 3000 Node: uncultured\_bacterium 798 ■ Node: uncultured\_bacterium 7475

ead count

20

50

0

Node

Top Reference

assigned reads

all reads

nonDup

readDis

nonStacked

2

edit distance: ancient

3

uncultured\_bacterium

uncultured\_bacterium

56252

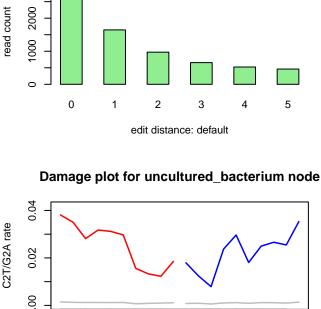
190

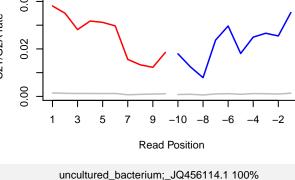
188

0.006

0

S0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractuS0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractu







Actinomyces;\_NR\_025633.1 100% Pseudopropionibacterium\_propionicum\_F0230a;\_CP002734.1 100%

uncultured\_Actinomyces\_sp.;\_AM420166.1 100%

Pseudopropionibacterium\_propionicum;\_NR\_114803.1 100% unidentified\_proteobacterium;\_X73223.1 100%

uncultured\_Propionibacterium\_sp.;\_FJ976211.1 100%

Pseudopropionibacterium\_propionicum;\_AF285117.1 100%

uncultured\_Lautropia\_sp.;\_DQ016725.2 100%

Off destacking? C>T\_1 0.0381 G>A\_-1 0.0353 mean length (sd) 108 (41.235)