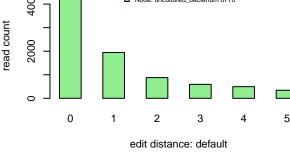
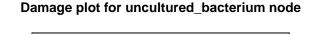
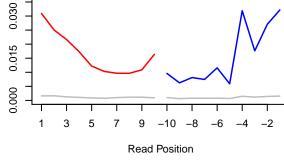
## 250 4000 ■ Node: uncultured\_bacterium 8770 Node: uncultured\_bacterium 666 ead count 150

50

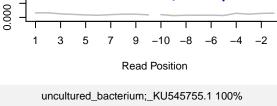
0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un)101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19un







C2T/G2A rate



| a  |    |
|--|----|
| Jeotgalicoccus_psychrophilus;_NR_025644.1 10 | 0% |
| uncultured_rumen_bacterium;_EF686620.1 100   | %  |

Jeotgalicoccus\_psychrophilus;\_EF463102.1 100%

Jeotgalicoccus\_sp.\_YIM\_KMY9-1;\_DQ358647.1 100% uncultured\_Jeotgalicoccus\_sp.;\_JX841330.1 100%

uncultured\_Jeotgalibacillus\_sp.;\_AM087528.1 100%

Jeotgalicoccus\_coquinae;\_NR\_117490.1 100% Jeotgalicoccus\_psychrophilus;\_EU194336.1 100%

Planococcaceae\_bacterium\_Storch\_2/2-2;\_KX957962.1 100%

0 2 3 edit distance: ancient Node uncultured\_bacterium Top Reference uncultured\_bacterium assigned reads 11247 all reads 857 nonDup 843 readDis 0.001 nonStacked 0 destacking? Off C>T\_1 0.0309

> 0.0322 63 (24.206)

G>A\_-1

mean length (sd)