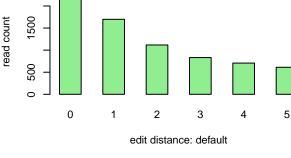
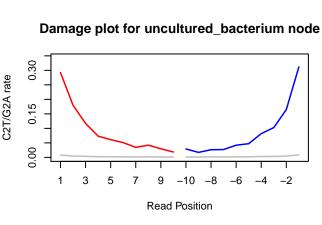
## ■ Node: uncultured\_bacterium 7374 Node: uncultured\_bacterium 2230 ead count 400

200

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





uncultured\_bacterium;\_JQ696889.1 100% [Eubacterium]\_yurii\_subsp.\_yurii;\_NR\_104843.1 100%

bacterium]\_yurii\_subsp.\_margaretiae\_ATCC\_43715;\_NR\_115259.1 100%

Eubacterium\_sp.\_oral\_strain\_A03MT;\_AF287765.1 100%

[Eubacterium]\_yurii\_subsp.\_schtitka;\_NR\_115260.1 100%

Peptostreptococcaceae\_bacterium\_AS15;\_HQ616364.1 100%

Eubacterium\_sp.\_oral\_clone\_OH3A;\_AY947497.1 100%

[Eubacterium]\_yurii\_subsp.\_yurii;\_L34629.2 100%

Pseudopropionibacterium\_propionicum\_F0230a;\_CP002734.1 100%

Peptostreptococcaceae\_bacterium\_OBRC9;\_HQ616354.1 100%

