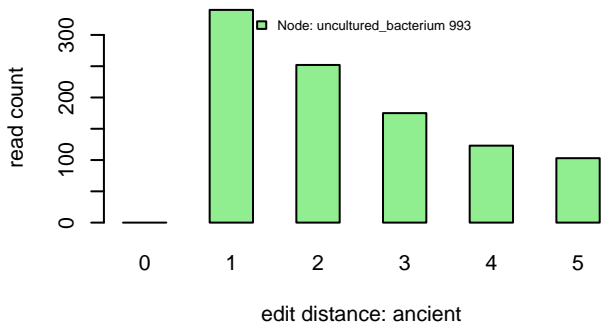
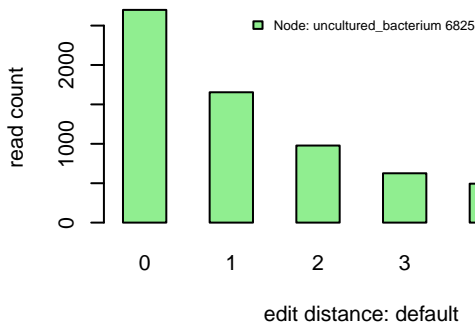
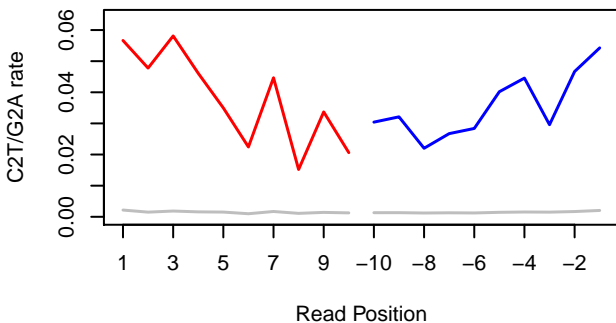


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for uncultured_bacterium node



| Node | uncultured_bacterium |
|------------------|----------------------|
| Top Reference | uncultured_bacterium |
| assigned reads | 8082 |
| all reads | 252 |
| nonDup | 246 |
| readDis | 0.029 |
| nonStacked | 1 |
| destacking? | Off |
| C>T_1 | 0.0567 |
| G>A_-1 | 0.0543 |
| mean length (sd) | 69 (25.809) |

| |
|--|
| uncultured_bacterium;_KJ910975.1 100% |
| Pseudopropionibacterium_propionicum_F0230a;_CP002734.1 100% |
| actinomyces_genomosp._P1_oral_clone_MB6_C03;_DQ003632.1 100% |
| uncultured_Actinomyces_sp.;_FJ976431.1 100% |
| Actinomyces;_NR_025633.1 100% |
| uncultured_Actinomyces_sp.;_AM420166.1 100% |
| uncultured_Actinomyces_sp.;_JQ406543.1 100% |
| Pseudopropionibacterium_propionicum;_NR_114803.1 100% |
| Bacteria;_FJ976437.1 100% |
| uncultured_Comamonadaceae_bacterium;_JQ406530.1 100% |