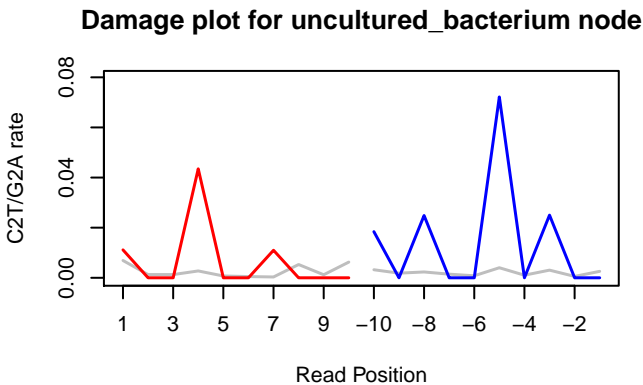
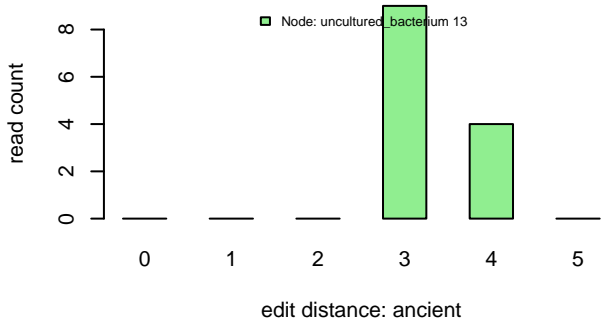
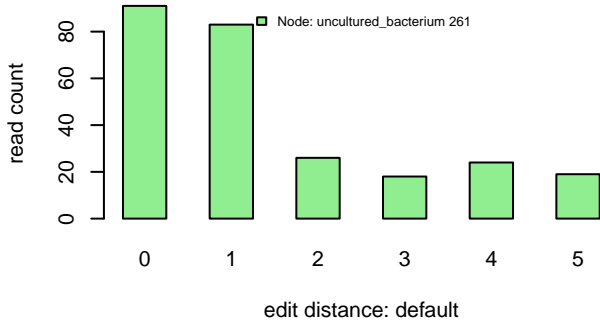


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



| Node | uncultured_bacterium |
|------------------|----------------------|
| Top Reference | uncultured_bacterium |
| assigned reads | 369 |
| all reads | 15 |
| nonDup | 1 |
| readDis | 1 |
| nonStacked | 1 |
| destacking? | Off |
| C>T_1 | 0.0111 |
| G>A_-1 | 0 |
| mean length (sd) | 60 (21.391) |

| |
|---|
| uncultured_bacterium;_KF312320.1 100% |
| uncultured_beta_proteobacterium_CBNPD1_BAC_clone_578;_EF157669.1 100% |
| uncultured_Spirochaetes_bacterium;_EU857752.1 100% |
| unassignedName;_AY344370.1 100% |
| uncultured_beta_proteobacterium;_EF702035.1 100% |
| uncultured_beta_proteobacterium;_AM706727.1 100% |
| uncultured_beta_proteobacterium;_EF562556.1 100% |
| uncultured_soil_bacterium;_AY037565.1 100% |
| uncultured_proteobacterium;_AY917252.1 100% |
| uncultured_proteobacterium;_EF019960.1 100% |