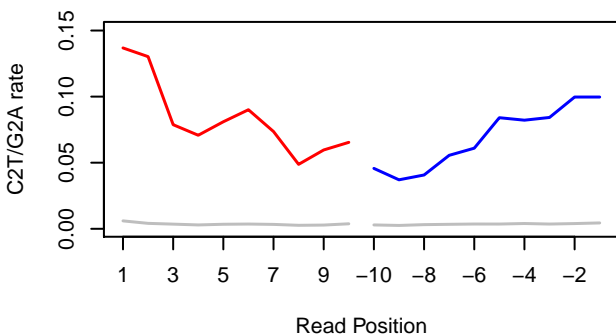


Damage plot for uncultured_bacterium node



| Node | uncultured_bacterium |
|------------------|----------------------|
| Top Reference | uncultured_bacterium |
| assigned reads | 7557 |
| all reads | 162 |
| nonDup | 161 |
| readDis | 0.03 |
| nonStacked | 0 |
| destacking? | Off |
| C>T_1 | 0.1368 |
| G>A_-1 | 0.0997 |
| mean length (sd) | 80 (25.514) |

uncultured_bacterium;_JQ972699.1 100%

uncultured_Bacteroidales_bacterium;_FJ976400.1 100%

Bacteroidetes_bacterium_Oral_Taxon_274;_FJ577256.2 100%

Bacteroidetes_bacterium_oral_taxon_274;_GU409195.1 100%

uncultured_Porphyromonadaceae_bacterium;_AM420225.1 100%

Actinomyces;_NR_025633.1 100%

uncultured_Actinomyces_sp.;_AM420166.1 100%

Actinomyces_genomosp._C1;_AY278610.1 100%

Bacteroides-like_sp._oral_clone_AU126;_AY005072.1 100%

Actinomyces_sp._oral_taxon_414;_CP012590.1 100%