

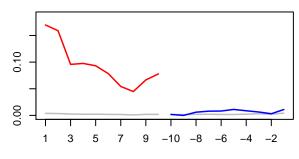
edit distance: default

0

C2T/G2A rate

2 3

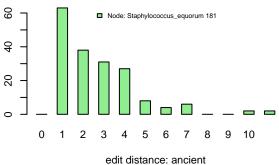
Damage plot for Staphylococcus_equorum node



Read Position

| Staphylococcus_equorum;_NZ_CP068576.1 100% | |
|--|--|
| NA;_NZ_CP068061.1 000% | |
| NA;_NZ_CP068479.1 000% | |
| NA% | |

Pool-3-031-AZP-216_S35.unmapped.rma6



read count

10

| cut distance, ancient | |
|-----------------------|------------------------|
| Node | Staphylococcus_equorum |
| Top Reference | Staphylococcus_equorum |
| all reads | 817 |
| nonDup | 638 |
| readDis | 0.913 |
| nonStacked | 541 |
| destacking? | Off |
| C>T_1 | 0.1697 |
| G>A1 | 0.011 |
| mean length (sd) | 65 (18.155) |
| | |