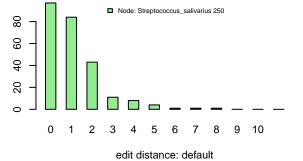
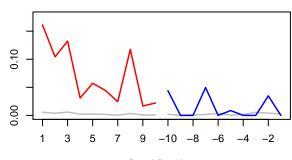
Pool-1-036-AZP-115_S36.unmapped.rma6



read count

C2T/G2A rate

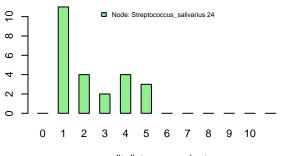
Damage plot for Streptococcus_salivarius node



Read Position

| Streptococcus_salivarius;_NZ_CP066093.1 100% | | |
|--|--|--|
| Streptococcus_salivarius_CCHSS3;_NC_015760.1 014% | | |
| Streptococcus_salivarius_57.I;_NC_017594.1 014% | | |
| Streptococcus_salivarius_JIM8777;_NC_017595.1 001% | | |
| NA;_NZ_CP065424.1 001% | | |
| NA% | | |

Pool-1-036-AZP-115_S36.unmapped.rma6



read count

edit distance: ancient

| Node | Streptococcus_salivarius |
|------------------|--------------------------|
| Top Reference | Streptococcus_salivarius |
| all reads | 125 |
| nonDup | 105 |
| readDis | 0.931 |
| nonStacked | 95 |
| destacking? | Off |
| C>T_1 | 0.161 |
| G>A1 | 0 |
| mean length (sd) | 63 (17.725) |
| | |