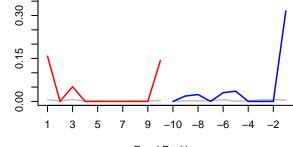


read count

C2T/G2A rate

Damage plot for Streptococcus_salivarius node



Read Position

| Streptococcus_salivarius;_NZ_LR134274.1 100% |
|---|
| Streptococcus;_NZ_CAKAKM010000001.1 077% |
| Streptococcus;_NZ_JACLQM010000007.1 066% |
| Streptococcus;_NZ_JACLQN010000005.1 066% |
| Streptococcus;_NZ_JACLQO010000003.1 066% |
| Streptococcus_salivarius_CCHSS3;_NC_015760.1 044% |
| Streptococcus_salivarius_57.I;_NC_017594.1 044% |
| Streptococcus;_NZ_JACLQJ010000017.1 033% |
| Streptococcus;_NZ_CAKAKM010000002.1 022% |

Streptococcus;_NZ_JACLQJ010000009.1 022%

Node: Streptococcus_salivarius 7

Pool-3-041-AZP-226_S45.unmapped.rma6

10

| edit | distance: | ancient |
|------|-----------|---------|
| | | |

read count

0

| euit distance, ancient | | |
|------------------------|--------------------------|--|
| Node | Streptococcus_salivarius | |
| Top Reference | Streptococcus_salivarius | |
| all reads | 15 | |
| nonDup | 12 | |
| readDis | 0.757 | |
| nonStacked | 9 | |
| destacking? | Off | |
| C>T_1 | 0.1579 | |
| G>A1 | 0.3158 | |
| mean length (sd) | 61 (16.915) | |
| | | |