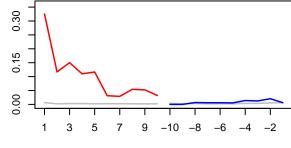


read count

C2T/G2A rate

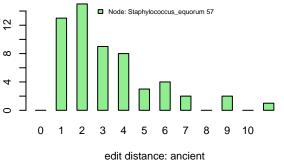
${\bf Damage\ plot\ for\ Staphylococcus_equorum\ node}$



Read Position

| Staphylococcus_equorum;_NZ_CP068576.1 100% | | |
|---|--|-------------------------------|
| NA;_NZ_JAJNIG010000028.1 010% | | |
| NA;_NZ_JAJNIH010000031.1 010% | | |
| Staphylococcus;_NZ_JAESMX010000012.1 005% | | |
| Staphylococcus;_NZ_JAGEVL010000015.1 003% Staphylococcus;_NZ_JAFENQ010000028.1 003% | | |
| | | NA;_NZ_JAJNHZ010000046.1 003% |
| NA;_NZ_JAJNHY010000042.1 003% | | |
| NA;_NZ_JAJNIV010000033.1 001% | | |
| Staphylococcus;_NZ_JAFEPM010000038.1 001% | | |

Pool-4-042-AZP-279_S43.unmapped.rma6



read count

| | Node | Staphylococcus_equorum |
|--|------------------|------------------------|
| | Top Reference | Staphylococcus_equorum |
| | all reads | 233 |
| | nonDup | 200 |
| | readDis | 0.892 |
| | nonStacked | 160 |
| | destacking? | Off |
| | C>T_1 | 0.3253 |
| | G>A1 | 0.006 |
| | mean length (sd) | 62 (17.149) |
| | | |