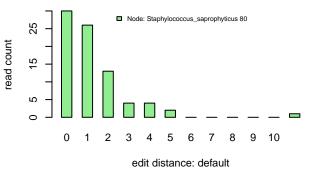
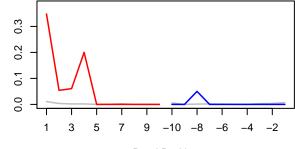
Pool-1-039-AZP-118_S39.unmapped.rma6



Damage plot for Staphylococcus_saprophyticus no

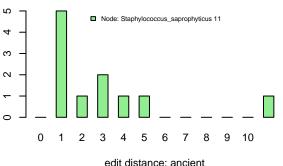


C2T/G2A rate

Read Position

| Staphylococcus_saprophyticus;_NZ_CP054440.1 100% |
|--|
| NA% |

Pool-1-039-AZP-118_S39.unmapped.rma6



read count

| 6 | cuit distance, ancient |
|------------------|------------------------------|
| Node | Staphylococcus_saprophyticus |
| Top Reference | Staphylococcus_saprophyticus |
| all reads | 79 |
| nonDup | 70 |
| readDis | 0.998 |
| nonStacked | 66 |
| destacking? | Off |
| C>T_1 | 0.3478 |
| G>A1 | 0 |
| mean length (sd) | 65 (18.037) |
| | |