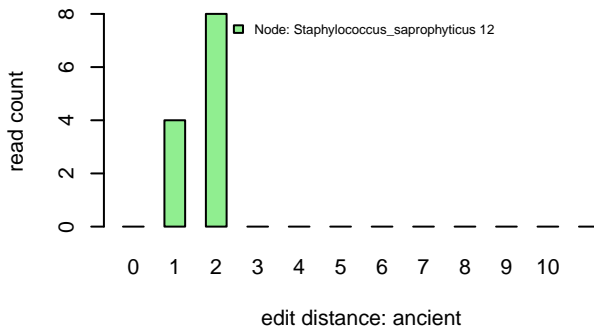


**Pool-4-040-AZP-277\_S41.unmapped.rma6**



The graph displays the C2T/G2A rate for two methods across different read positions. The red line, representing the top method, starts with a high rate of approximately 0.4 at position 1, fluctuates with peaks around 0.2 at positions 3 and 5, and then drops to near zero by position 10. The blue line, representing the bottom method, remains consistently near zero across all positions, with a very slight increase around position -2.

| Read Position | Top Method (Red Line) C2T/G2A rate | Bottom Method (Blue Line) C2T/G2A rate |
|---------------|------------------------------------|--|
| 1             | 0.40                               | 0.00                                   |
| 2             | 0.15                               | 0.00                                   |
| 3             | 0.20                               | 0.00                                   |
| 4             | 0.05                               | 0.00                                   |
| 5             | 0.20                               | 0.00                                   |
| 6             | 0.05                               | 0.00                                   |
| 7             | 0.05                               | 0.00                                   |
| 8             | 0.00                               | 0.00                                   |
| 9             | 0.05                               | 0.00                                   |
| 10            | 0.00                               | 0.00                                   |
| -10           | 0.00                               | 0.00                                   |
| -8            | 0.00                               | 0.00                                   |
| -6            | 0.00                               | 0.00                                   |
| -4            | 0.00                               | 0.00                                   |
| -2            | 0.00                               | 0.05                                   |
| 0             | 0.00                               | 0.00                                   |

| Node             | Staphylococcus_saprophyticus |
|------------------|------------------------------|
| Top Reference    | Staphylococcus_saprophyticus |
| all reads        | 83                           |
| nonDup           | 72                           |
| readDis          | 1                            |
| nonStacked       | 72                           |
| destacking?      | Off                          |
| C>T_1            | 0.4211                       |
| G>A_1            | 0                            |
| mean length (sd) | 53 (16.315)                  |

[illegible]