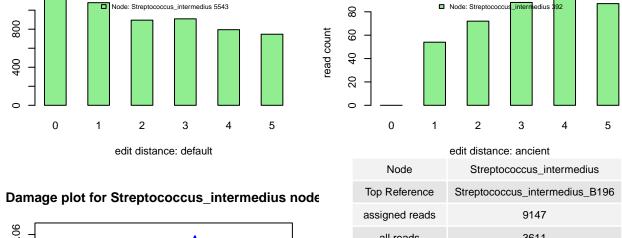
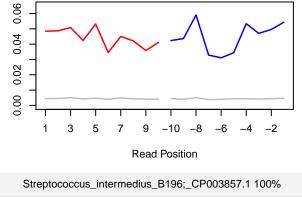
Node: Streptococcus_intermedius 5543 ■ Node: Streptococcus_intermedius

S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu





| 3 _ | _ | | | | | _ | | | | | |
|---|---|---|---|---|---|-----|----|----|----|----|---|
| | | | | | | | | | | | _ |
| | 1 | 3 | 5 | 7 | 9 | -10 | -8 | -6 | -4 | -2 | |
| Read Position | | | | | | | | | | | |
| Streptococcus_intermedius_B196;_CP003857.1 100% | | | | | | | | | | | |
| | | | | | | | | | | | |

Streptococcus_intermedius_C270;_CP003858.1 100%

Streptococcus_intermedius;_AP014880.1 100% Streptococcus_intermedius;_CP020433.1 100%

Streptococcus_intermedius;_CP012718.1 100%

Streptococcus_sanguinis_SK36;_CP000387.1 050% Streptococcus_intermedius;_AB608790.1 021%

Streptococcus_intermedius_JTH08;_AP010969.1 100%

Streptococcus_gordonii_str._Challis_substr._CH1;_CP000725.1 006%

Streptococcus_intermedius;_AB045140.2 006%

| 40 | ? - | | | | | | | | |
|----|------------------------|--------------------------------|------|---|---|---|--|--|--|
| 20 | 3 - | | | | | | | | |
| C | ,] | | | | | | | | |
| | 0 | 1 | 2 | 3 | 4 | 5 | | | |
| | edit distance: ancient | | | | | | | | |
| | Node | Streptococcus_intermedius | | | | | | | |
| | Top Referer | Streptococcus_intermedius_B196 | | | | | | | |
| | assigned rea | ads | 9147 | | | | | | |
| | all reads | | 3611 | | | | | | |
| | nonDup | | 3524 | | | | | | |
| | readDis | 0.141 | | | | | | | |
| | | | | | | | | | |

nonStacked 373 destacking? Off C>T_1 0.0485 G>A_-1 0.0543 113 (41.006) mean length (sd)