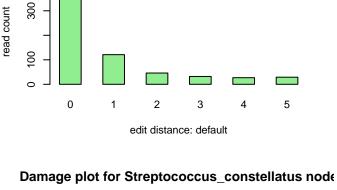
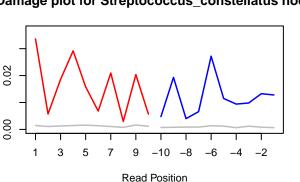
## Node: Streptococcus\_constellatus 24 ■ Node: Streptococcus\_constellatus 689 ead count

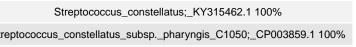
9

S0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractuS0\_L001\_R1\_001.fastq.combined.fq.prefixed.extractu





C2T/G2A rate



treptococcus\_constellatus\_subsp.\_pharyngis\_C818;\_CP003840.1 100% treptococcus\_constellatus\_subsp.\_pharyngis\_C232;\_CP003800.1 100%

Streptococcus\_constellatus\_subsp.\_constellatus;\_KY315463.1 100% Streptococcus\_constellatus\_subsp.\_constellatus;\_KY315464.1 100%

Streptococcus\_intermedius;\_KY315447.1 100%

Streptococcus\_intermedius;\_HM241929.1 100%

Streptococcus\_constellatus\_subsp.\_pharyngis;\_KY315452.1 100%

Streptococcus\_constellatus\_subsp.\_pharyngis;\_KY315450.1 100%

0 2 edit distance: ancient Node Streptococcus\_constellatus Top Reference Streptococcus\_constellatus assigned reads 918 all reads 453 nonDup 441 readDis 0.184 nonStacked 43 Off destacking? C>T\_1 0.0336 G>A\_-1 0.0128

mean length (sd)

109 (38.519)