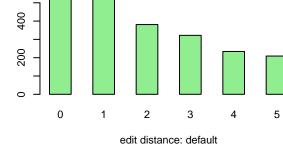
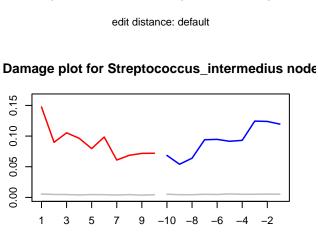
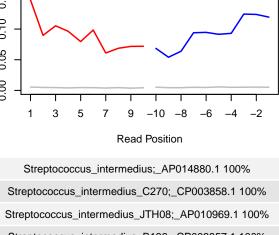
## Node: Streptococcus\_intermedius 2255 80 Node: Streptococcus\_intermedius 372 9 read count

40

0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur





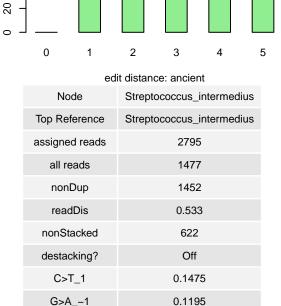


Streptococcus\_intermedius\_B196;\_CP003857.1 100%

Streptococcus\_sanguinis\_SK36;\_CP000387.1 008% Streptococcus\_cristatus\_AS\_1.3089;\_CP004409.1 004%

Streptococcus\_anginosus\_C1051;\_CP003860.1 002% Streptococcus\_sp.\_NPS\_308;\_AP01

Streptococcus_spNPS_308;_AP017652.1 001%	
Streptococcus_anginosus_C238;_CP003861.1 001%	



66 (23.692)

mean length (sd)

Streptococcus\_gordonii\_str.\_Challis\_substr.\_CH1;\_CP000725.1 001%