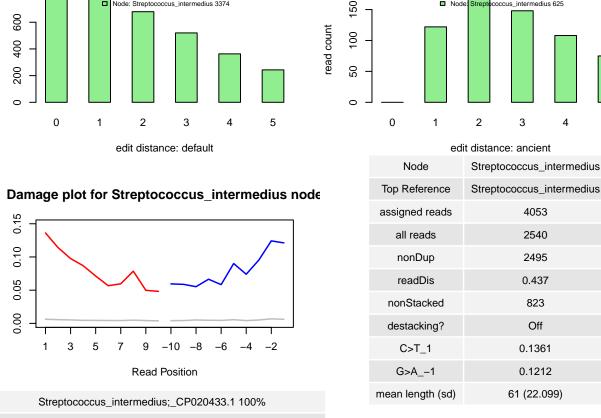
## ■ Node: Streptococcus\_intermedius 625 Node: Streptococcus\_intermedius 3374 150

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



assigned reads	4053
all reads	2540
nonDup	2495
readDis	0.437
nonStacked	823
destacking?	Off
C>T_1	0.1361
G>A1	0.1212
mean length (sd)	61 (22.099)

Streptococcus\_intermedius\_B196;\_CP003857.1 100% Streptococcus\_intermedius\_JTH08;\_AP010969.1 100%

C2T/G2A rate

Streptococcus\_sanguinis\_SK36;\_CP000387.1 030%

Streptococcus\_intermedius\_C270;\_CP003858.1 100%

Streptococcus\_cristatus\_AS\_1.3089;\_CP004409.1 005%

Streptococcus\_gordonii\_str.\_Challis\_substr.\_CH1;\_CP000725.1 004%

Streptococcus\_gordonii;\_CP017295.1 003% Streptococcus\_gordonii;\_CP020450.1 002%

Streptococcus\_gordonii;\_CP012648.1 002%