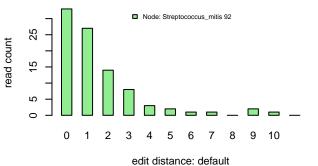
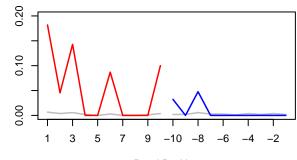
Pool-1-007-AZP-86_S7.unmapped.rma6

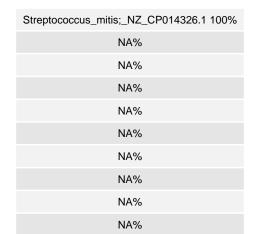


Damage plot for Streptococcus_mitis node

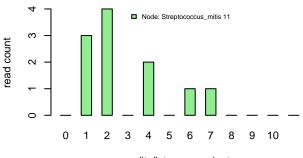


C2T/G2A rate

Read Position



Pool-1-007-AZP-86_S7.unmapped.rma6



edit distance: ancient

Node	Streptococcus_mitis
Top Reference	Streptococcus_mitis
all reads	37
nonDup	31
readDis	0.982
nonStacked	29
destacking?	Off
C>T_1	0.1818
G>A1	0
mean length (sd)	53 (17.495)