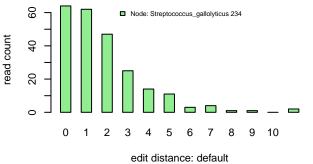
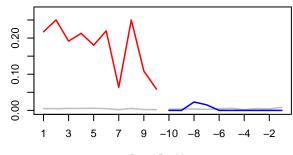
Pool-4-012-AZP-249_S13.unmapped.rma6



Damage plot for Streptococcus_gallolyticus node

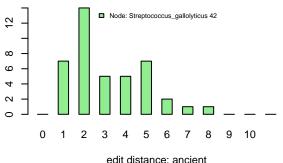


C2T/G2A rate

Read Position

Streptococcus_gallolyticus;_NZ_LS483409.1 100%
Streptococcus;_NZ_CP031733.1 000%
NA%

Pool-4-012-AZP-249_S13.unmapped.rma6



read count

edit distance: ancien

cuit distance. ancient	
Node	Streptococcus_gallolyticus
Top Reference	Streptococcus_gallolyticus
all reads	177
nonDup	151
readDis	0.931
nonStacked	140
destacking?	Off
C>T_1	0.2174
G>A1	0
mean length (sd)	60 (18.296)