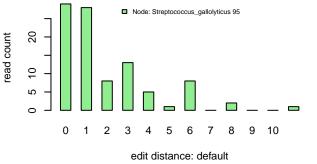
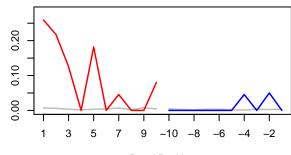
Pool-1-039-AZP-118_S39.unmapped.rma6



Damage plot for Streptococcus_gallolyticus node

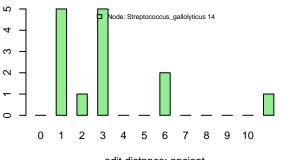


C2T/G2A rate

Read Position

Streptococcus_gallolyticus;_NZ_LS483409.1 100%	
NA%	

Pool-1-039-AZP-118_S39.unmapped.rma6



read count

edit distance: ancient

Node	Streptococcus_gallolyticus
Top Reference	Streptococcus_gallolyticus
all reads	75
nonDup	64
readDis	0.947
nonStacked	58
destacking?	Off
C>T_1	0.2593
G>A1	0
mean length (sd)	64 (20.346)