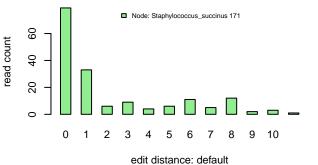
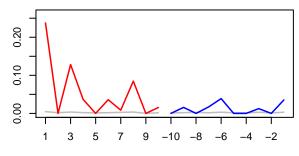
## Pool-4-031-AZP-268\_S32.unmapped.rma6



## Damage plot for Staphylococcus\_succinus node

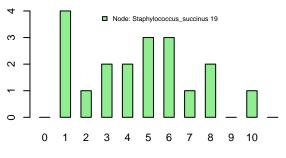


C2T/G2A rate

Read Position

NA;_NZ_CP075505.1 100%		
Staphylococcus_succinus;_NZ_CP018199.1 100%		
NA%		

## Pool-4-031-AZP-268\_S32.unmapped.rma6



read count

edit distance: ancient

Node	Staphylococcus_succinus
Top Reference	NA
all reads	98
nonDup	84
readDis	0.634
nonStacked	48
destacking?	Off
C>T_1	0.2375
G>A1	0.0353
mean length (sd)	64 (16.196)