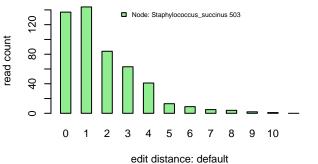
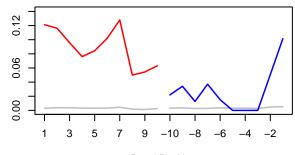
Pool-3-041-AZP-226_S45.unmapped.rma6



Damage plot for Staphylococcus_succinus node

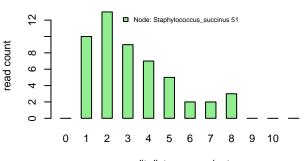


C2T/G2A rate

Read Position

Staphylococcus_succinus;_NZ_CP018199.1 100%		
NA;_NZ_CP022047.2 000%		
NA%		

Pool-3-041-AZP-226_S45.unmapped.rma6



edit distance: ancient

Node	Staphylococcus_succinus
Top Reference	Staphylococcus_succinus
all reads	503
nonDup	417
readDis	0.974
nonStacked	381
destacking?	Off
C>T_1	0.1212
G>A1	0.1014
mean length (sd)	64 (15.538)