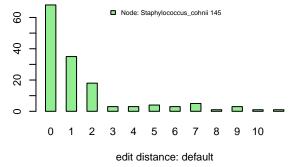
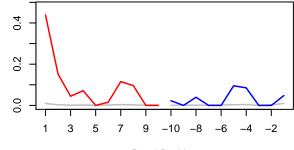
Pool-3-045-AZP-230_S49.unmapped.rma6



read count

C2T/G2A rate

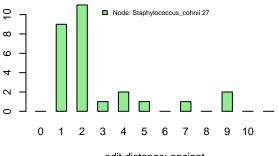
Damage plot for Staphylococcus_cohnii node



Read Position



Pool-3-045-AZP-230_S49.unmapped.rma6



ead count

edit distance: ancient

Node	Staphylococcus_cohnii
Top Reference	NA
all reads	86
nonDup	69
readDis	0.904
nonStacked	59
destacking?	Off
C>T_1	0.4388
G>A1	0.0476
mean length (sd)	55 (14.996)