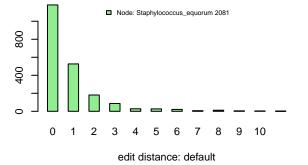
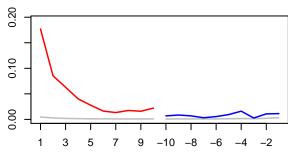
## Pool-3-027-AZP-212\_S31.unmapped.rma6



read count

C2T/G2A rate

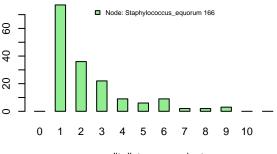
## Damage plot for Staphylococcus\_equorum node



Read Position

Staphylococcus_equorum;_NZ_CP068576.1 100%
NA;_NZ_CP065795.1 000%
NA;_NZ_CP059679.1 000%
NA;_NZ_LT906462.1 000%
NA;_NZ_CP065961.1 000%
NA;_NZ_CP080324.1 000%
NA%
NA%
NA%
NA%

## Pool-3-027-AZP-212\_S31.unmapped.rma6



read count

edit distance: ancient

euit distance, ancient	
Node	Staphylococcus_equorum
Top Reference	Staphylococcus_equorum
all reads	1407
nonDup	1168
readDis	0.883
nonStacked	917
destacking?	Off
C>T_1	0.177
G>A1	0.0113
mean length (sd)	66 (17.223)