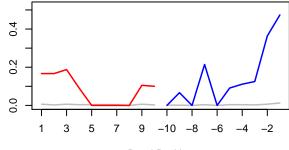


read count

C2T/G2A rate

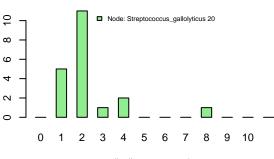
Damage plot for Streptococcus_gallolyticus node



Read Position

Streptococcus_gallolyticus;_NZ_LS483409.1 100%	
NA%	

EASI236.unmapped.rma6



edit distance: ancient

cuit distance, ancient	
Node	Streptococcus_gallolyticus
Top Reference	Streptococcus_gallolyticus
all reads	51
nonDup	51
readDis	1
nonStacked	51
destacking?	Off
C>T_1	0.1667
G>A1	0.4737
mean length (sd)	45 (12.241)