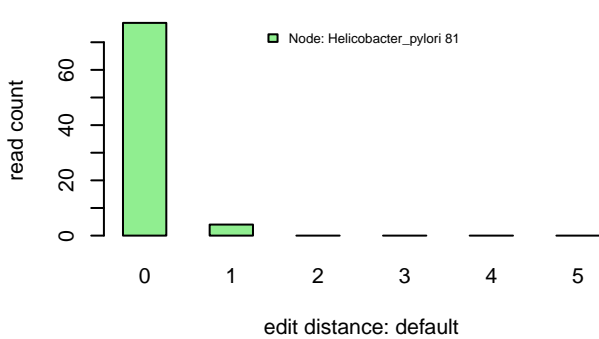
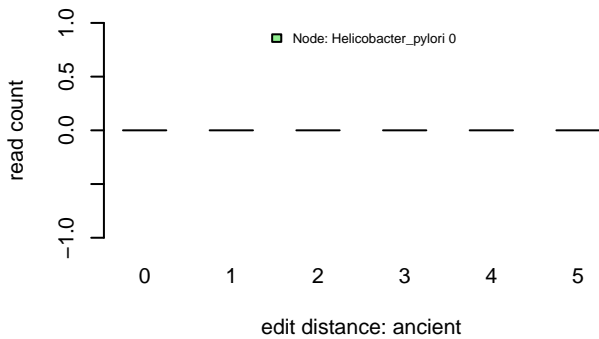


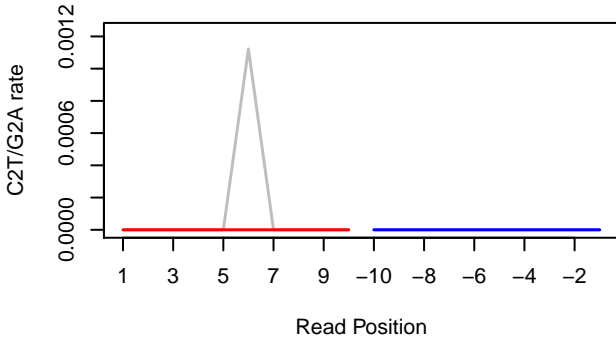
AT+Mock-1:100.rma6



AT+Mock-1:100.rma6



Damage plot for Helicobacter_pylori node



Node	Helicobacter_pylori
Top Reference	Helicobacter_pylori_Rif1
assigned reads	105
all reads	104
nonDup	104
readDis	0.78
nonStacked	80
destacking?	On
C>T_1	0
G>A_-1	0
mean length (sd)	75 (3.231)

Helicobacter_pylori_Rif1;_CP003905.1	100%
Helicobacter_pylori_Rif2;_CP003906.1	100%
Helicobacter_pylori_26695;_CP003904.1	100%
Helicobacter_pylori_26695;_AE000511.1	100%
Helicobacter_pylori;_CP010436.1	100%
Helicobacter_pylori_26695-1CH;_AP013355.1	100%
Helicobacter_pylori_26695-1CL;_AP013356.1	100%
Helicobacter_pylori_26695-1;_AP013354.1	100%
Helicobacter_pylori_26695-1;_CP010435.1	100%
Helicobacter_pylori;_CP012907.1	016%