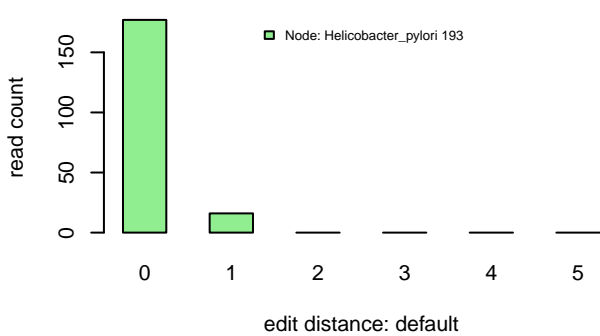
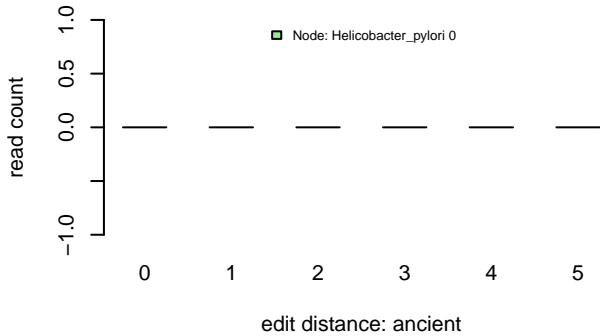


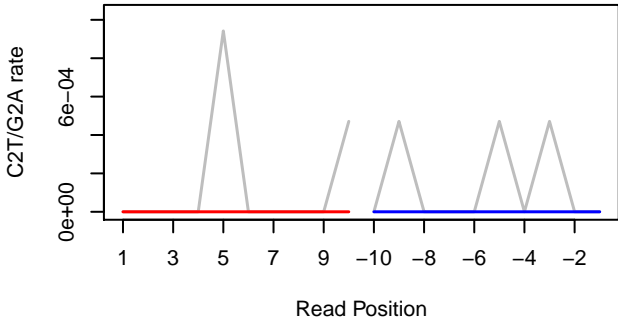
AT+Mock-1:50.rma6



AT+Mock-1:50.rma6



Damage plot for Helicobacter\_pylori node



Node	Helicobacter_pylori
Top Reference	Helicobacter_pylori_Rif1
assigned reads	264
all reads	261
nonDup	261
readDis	0.746
nonStacked	190
destacking?	On
C>T_1	0
G>A_-1	0
mean length (sd)	75 (2.145)

Helicobacter_pylori_Rif1;_CP003905.1	100%
Helicobacter_pylori_Rif2;_CP003906.1	100%
Helicobacter_pylori_26695;_AE000511.1	100%
Helicobacter_pylori;_CP010436.1	100%
Helicobacter_pylori_26695-1CH;_AP013355.1	100%
Helicobacter_pylori_26695;_CP003904.1	099%
Helicobacter_pylori_26695-1CL;_AP013356.1	099%
Helicobacter_pylori_26695-1;_AP013354.1	099%
Helicobacter_pylori_26695-1;_CP010435.1	099%
Helicobacter_pylori;_CP007605.1	012%