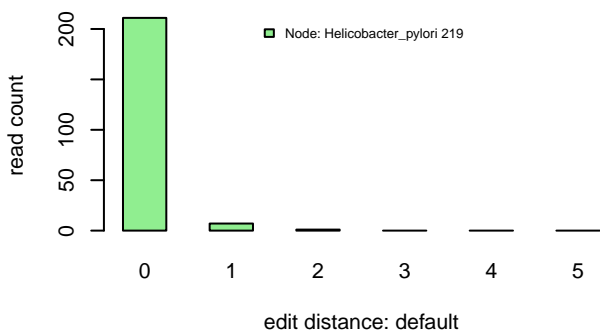
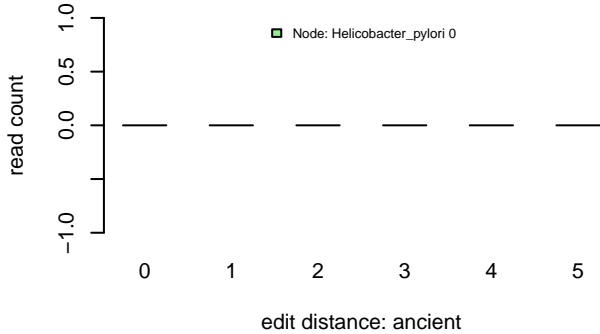


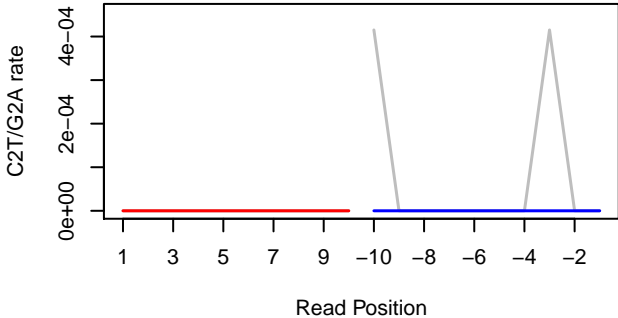
AT+Mock-1:50-1.rma6



AT+Mock-1:50-1.rma6



Damage plot for Helicobacter_pylori node



Node	Helicobacter_pylori
Top Reference	Helicobacter_pylori_26695
assigned reads	264
all reads	264
nonDup	264
readDis	0.842
nonStacked	219
destacking?	On
C>T_1	0
G>A_-1	0
mean length (sd)	75 (1.775)

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;_CP010435.1 100%
Helicobacter_pylori_Rif1;_CP003905.1 099%
Helicobacter_pylori_Rif2;_CP003906.1 099%
Helicobacter_pylori_26695-1;_AP013354.1 099%
Helicobacter_pylori_HPAG1;_CP000241.1 013%
Helicobacter_pylori_BM012S;_CP006889.1 012%