

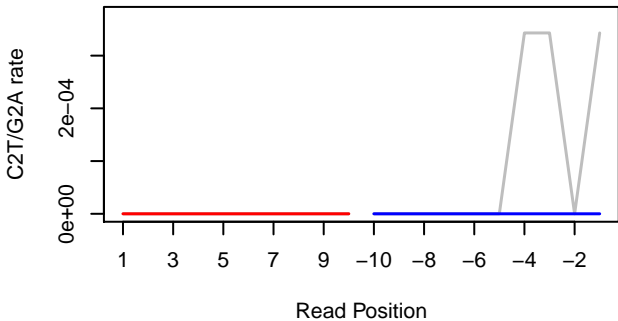
AT+Mock-1:50-4.rma6



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Damage plot for Helicobacter\_pylori node



Node	Helicobacter_pylori
Top Reference	Helicobacter_pylori
assigned reads	344
all reads	342
nonDup	342
readDis	0.779
nonStacked	263
destacking?	On
C>T_1	0
G>A_-1	0
mean length (sd)	75 (1.712)

Helicobacter_pylori;_CP010436.1 100%
Helicobacter_pylori_26695-1;_CP010435.1 099%
Helicobacter_pylori_26695;_AE000511.1 099%
Helicobacter_pylori_26695-1CH;_AP013355.1 099%
Helicobacter_pylori_26695-1CL;_AP013356.1 099%
Helicobacter_pylori_26695-1;_AP013354.1 099%
Helicobacter_pylori_Rif1;_CP003905.1 098%
Helicobacter_pylori_Rif2;_CP003906.1 098%
Helicobacter_pylori_26695;_CP003904.1 098%
Helicobacter_pylori_B8;_FN598874.1 015%