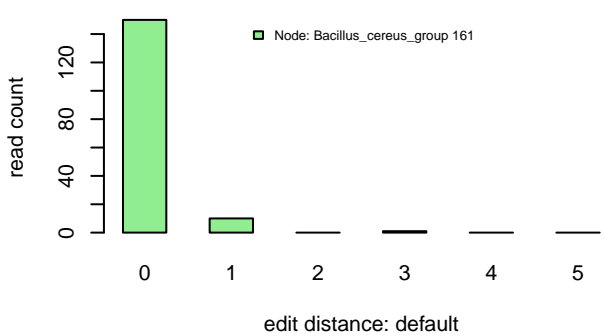
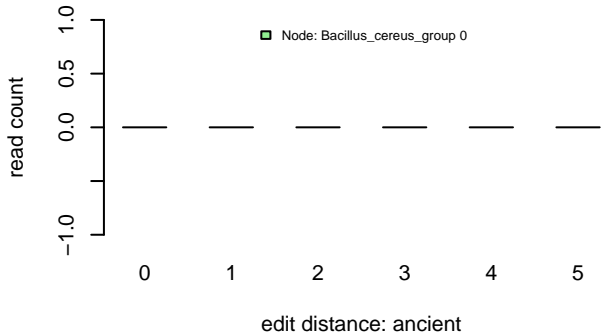


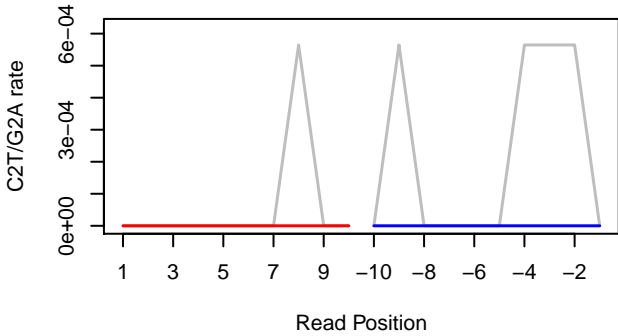
AT+Mock-1:50-3.rma6



AT+Mock-1:50-3.rma6



Damage plot for Bacillus_cereus_group node



Node	Bacillus_cereus_group
Top Reference	Bacillus_cereus_ATCC_10987
assigned reads	199
all reads	196
nonDup	196
readDis	0.806
nonStacked	158
destacking?	On
C>T_1	0
G>A_-1	0
mean length (sd)	75 (3.048)

Bacillus_cereus_ATCC_10987;_AE017194.1 100%
Bacillus_cereus_F;_CM001787.1 076%
Bacillus_cereus_FRI-35;_CP003747.1 072%
Bacillus_cereus;_CP016316.1 063%
Bacillus_thuringiensis_serovar_finitimus_YBT-020;_CP002508.1 058%
Bacillus_cereus_AH187;_CP001177.1 050%
Bacillus_cereus_NC7401;_AP007209.1 050%
Bacillus_cereus_BDRD-ST26;_CM000724.1 049%
Bacillus_cereus_Q1;_CP000227.1 048%
Bacillus_sp._ABP14;_CP017016.1 048%