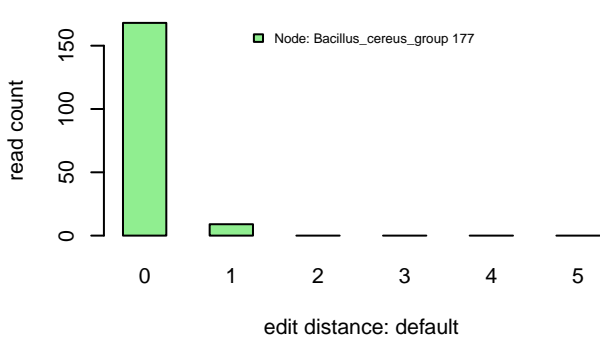
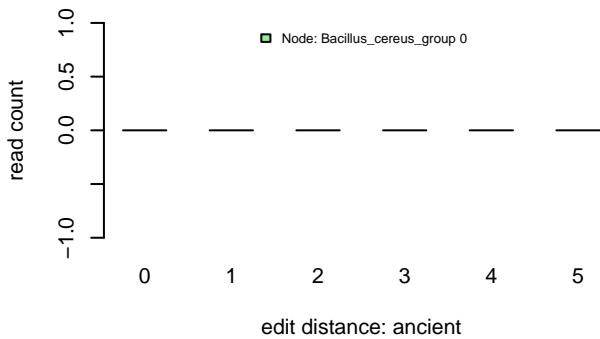


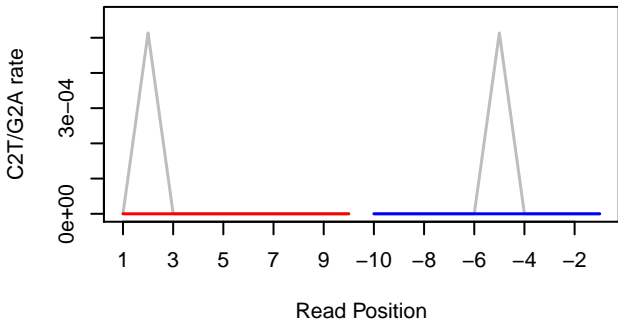
AT+Mock-1:50.rma6



AT+Mock-1:50.rma6



Damage plot for Bacillus\_cereus\_group node



Node	Bacillus_cereus_group
Top Reference	Bacillus_cereus_ATCC_10987
assigned reads	252
all reads	246
nonDup	246
readDis	0.733
nonStacked	174
destacking?	On
C>T_1	0
G>A_-1	0
mean length (sd)	75 (2.329)

Bacillus_cereus_ATCC_10987;_AE017194.1 100%
Bacillus_cereus_FRI-35;_CP003747.1 082%
Bacillus_cereus_F;_CM001787.1 076%
Bacillus_cereus;_CP016316.1 063%
Bacillus_thuringiensis_serovar_finitimus_YBT-020;_CP002508.1 062%
Bacillus_cereus_AH187;_CP001177.1 053%
Bacillus_cereus_NC7401;_AP007209.1 052%
Bacillus_cereus_BDRD-ST26;_CM000724.1 052%
Bacillus_anthraxis;_CP007618.1 051%
Bacillus_anthraxis;_CP009325.1 051%