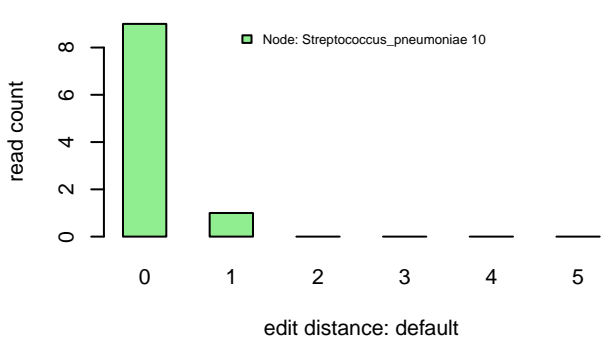
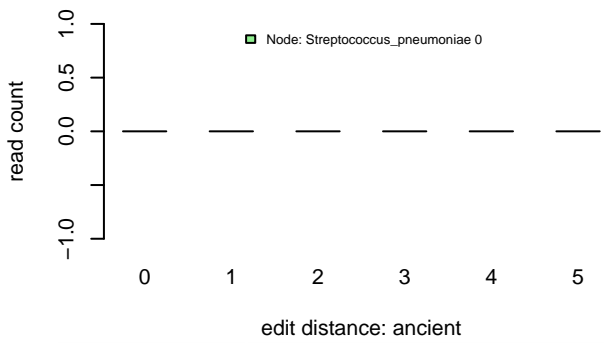


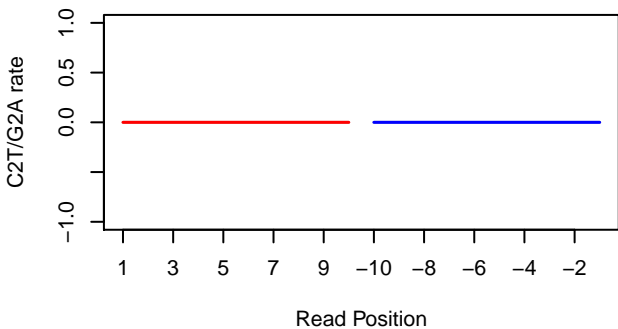
AT+Mock-1:50-4.rma6



AT+Mock-1:50-4.rma6



Damage plot for Streptococcus_pneumoniae node



Streptococcus_pneumoniae_TIGR4;_AKVY01000001.1	100%
Streptococcus_pneumoniae_JJA;_CP000919.1	077%
Streptococcus_pneumoniae_PCS8235;_CM001835.1	077%
Streptococcus_pneumoniae_70585;_CP000918.1	066%
Streptococcus_pneumoniae_AP200;_CP002121.1	055%
Streptococcus_pneumoniae_SPN033038;_FQ312042.1	055%
Streptococcus_pneumoniae_OXC141;_FQ312027.1	055%
Streptococcus_pneumoniae_INV104;_FQ312030.1	055%
Streptococcus_pneumoniae_ATCC_700669;_FM211187.1	055%
Streptococcus_pneumoniae_SPN034183;_FQ312043.1	055%

Node	Streptococcus_pneumoniae
Top Reference	Streptococcus_pneumoniae_TIGR4
assigned reads	14
all reads	13
nonDup	13
readDis	0.686
nonStacked	9
destacking?	On
C>T_1	0
G>A_-1	0
mean length (sd)	74 (3.957)