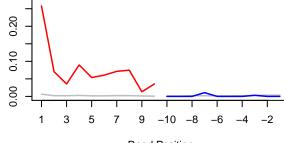


Damage plot for Streptococcus_pneumoniae node



Read Position

Streptococcus_pneumoniae_70585;_NC_012468.1 100%

Streptococcus_pneumoniae_TIGR4;_NZ_AKVY01000001.1 100% Streptococcus_pneumoniae_SPNA45;_NC_018594.1 100%

Streptococcus_pneumoniae;_NZ_LR216017.1 100%

Streptococcus_pneumoniae_TIGR4;_NC_003028.3 100%

Streptococcus_pneumoniae_SPN032672;_NC_021003.1 100%

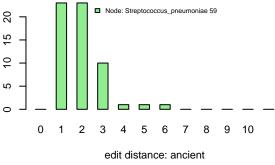
Streptococcus_pneumoniae_PCS8235;_NZ_CM001835.1 100%

Streptococcus_pneumoniae_ATCC_700669;_NC_011900.1 100% Streptococcus_pneumoniae_670-6B;_NC_014498.1 100%

Streptococcus_pneumoniae_Hungary19A-6;_NC_010380.1 100%

■ Node: Streptococcus_pneumoniae 59

Pool-1-009-AZP-88_S9.unmapped.rma6



ead count

edit distance: ancient	
Node	Streptococcus_pneumoniae
Top Reference	Streptococcus_pneumoniae
all reads	378
nonDup	311
readDis	0.968
nonStacked	295
destacking?	Off
C>T_1	0.2579
G>A1	0
mean length (sd)	55 (16.245)