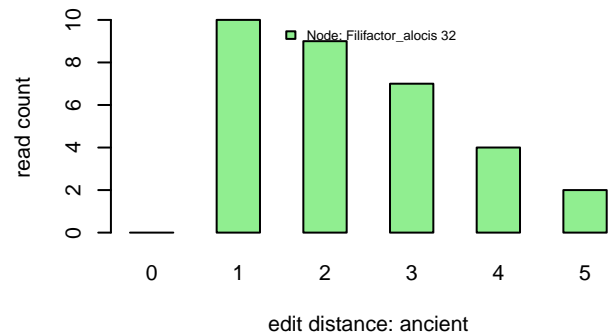
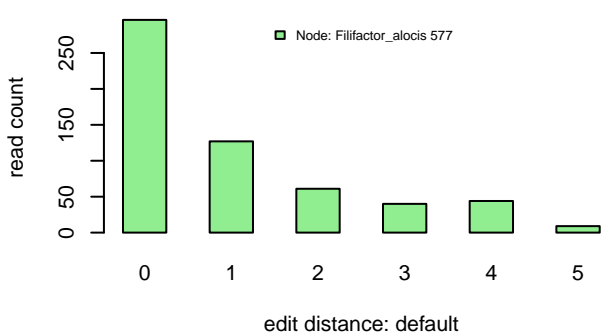
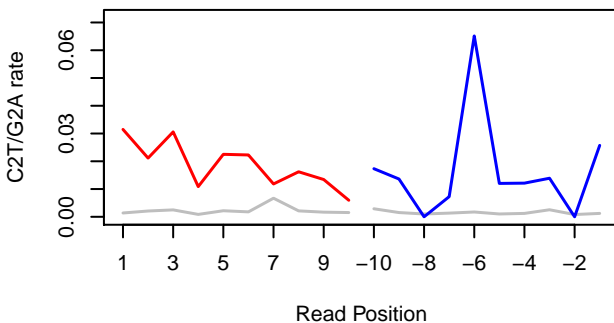


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Damage plot for Filifactor_alocis node



Node	Filifactor_alocis
Top Reference	Filifactor_alocis
assigned reads	660
all reads	647
nonDup	638
readDis	0
nonStacked	0
destacking?	Off
C>T_1	0.0315
G>A_-1	0.0257
mean length (sd)	86 (22.606)

Filifactor_alocis;_NR_076817.1 100%
Filifactor_alocis_ATCC_35896;_CP002390.1 100%
uncultured_Filifactor_sp.;_AM419956.1 100%
uncultured_Filifactor_sp.;_GU902747.1 100%
environmental_samples_<Filifactor>;_AB121851.1 100%
uncultured_bacterium;_FJ470466.1 100%
uncultured_bacterium;_HQ894888.1 100%
uncultured_bacterium;_HQ894871.1 100%
Filifactor_sp._oral_taxon_D70;_GU431025.1 100%
uncultured_bacterium;_HQ894872.1 100%