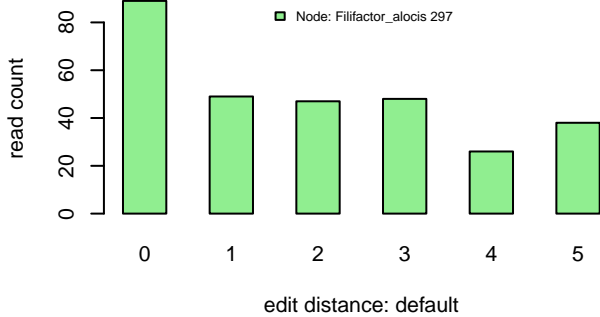
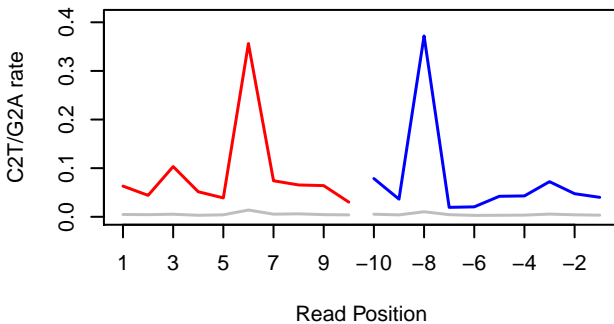


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for Filifactor_alocis node



Node	Filifactor_alocis
Top Reference	Filifactor_alocis
assigned reads	518
all reads	511
nonDup	503
readDis	0.001
nonStacked	0
destacking?	Off
C>T_1	0.0631
G>A_1	0.04
mean length (sd)	111 (42.132)

Filifactor_alocis;_NR_076817.1 100%
Filifactor_alocis_ATCC_35896;_CP002390.1 100%
unidentified_oral_bacterium_AP60-5;_AB028365.1 100%
uncultured_Filifactor_sp.;_AM419956.1 100%
uncultured_Filifactor_sp.;_GU902747.1 100%
uncultured_Filifactor_sp.;_EU071542.1 100%
environmental_samples_<Filifactor>;_AB121826.1 100%
environmental_samples_<Filifactor>;_AB121851.1 100%
uncultured_bacterium;_FJ470466.1 100%
Filifactor_sp._oral_clone_BP1-81;_AB121865.1 100%