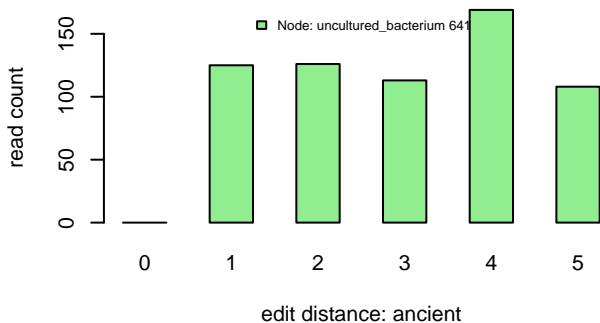
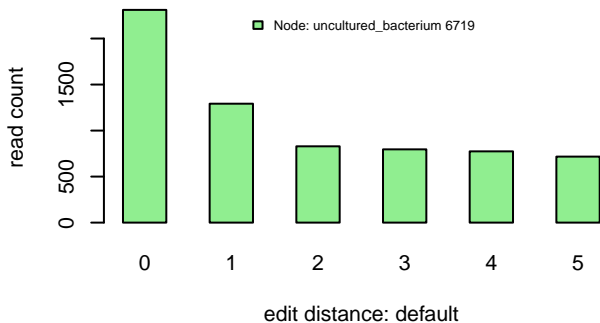
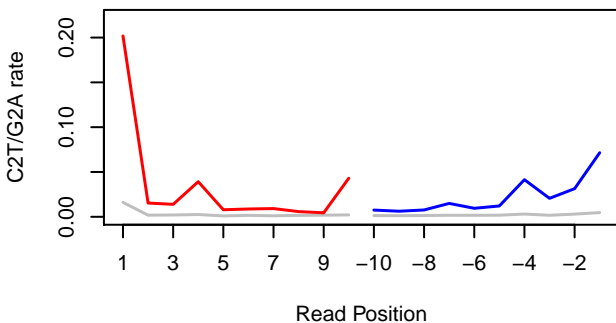


d_S0_L000_R1_000.fastq.merged.prefixed.hg19unmap_S0_L000_R1_000.fastq.merged.prefixed.hg19unmap



Damage plot for uncultured_bacterium node



Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	10059
all reads	115
nonDup	67
readDis	0.011
nonStacked	0
destacking?	Off
C>T_1	0.2018
G>A_-1	0.0715
mean length (sd)	69 (22.394)

uncultured_bacterium;_JN541288.1 100%
Brugia_timori;_LK910992.1 100%
Sphinx1.76-related_DNA;_LK931492.1 100%
uncultured_Bacteroidetes_bacterium_SBI2-18_P41A3;_DQ272742.1 100%
Sphinx1.76-related_DNA;_LK931491.1 100%
Clostridium_phoceensis;_LN846907.1 100%
Myroides_odoratus;_M62807.1 100%
Clostridium_sp._enrichment_culture_clone_7-14;_FJ625861.1 100%
Intestinimonas_butyrificiproducens;_CP011307.1 100%
uncultured_rumen_bacterium;_AY315346.1 100%