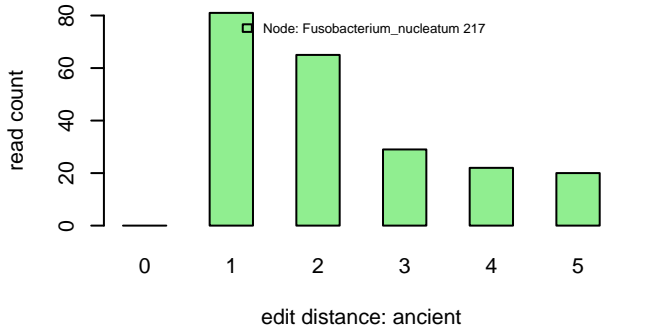
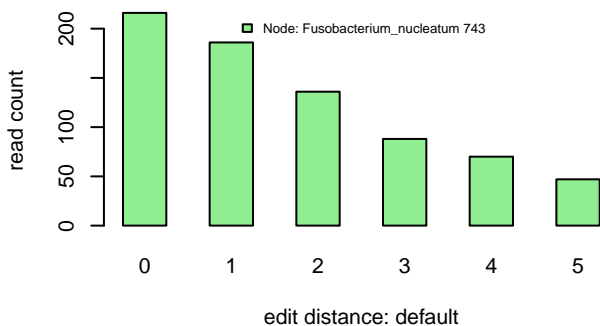
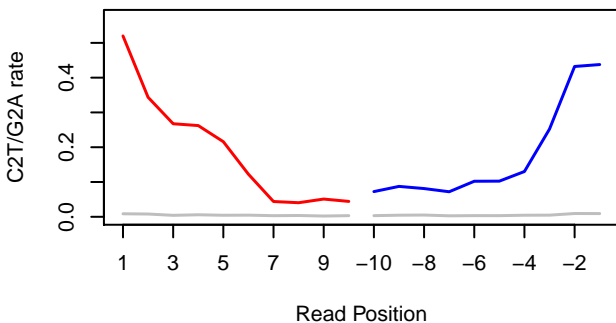


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un



Damage plot for Fusobacterium_nucleatum node



Node	Fusobacterium_nucleatum
Top Reference	Fusobacterium_nucleatum_subsp._polymorphum
assigned reads	879
all reads	238
nonDup	197
readDis	0.696
nonStacked	138
destacking?	Off
C>T_1	0.5199
G>A_-1	0.4376
length (sd)	48 (14.25)

Fusobacterium_nucleatum_subsp._polymorphum;_CP013328.1	100%
bacterium_nucleatum_subsp._nucleatum_ChDC_F316;_CP012716.1	100%
Fusobacterium_nucleatum_subsp._nucleatum;_CP012717.1	100%
Fusobacterium_nucleatum_subsp._nucleatum;_CP022122.1	100%
bacterium_nucleatum_subsp._nucleatum_ATCC_25586;_AE009951.2	100%
Fusobacterium_nucleatum_subsp._animalis;_CP012713.1	100%
Fusobacterium_nucleatum_subsp._animalis;_CP012715.1	100%
Fusobacterium_nucleatum_subsp._animalis_4_8;_CP003723.1	100%
obacterium_nucleatum_subsp._vincentii_3_1_36A2;_CP003700.1	094%
Fusobacterium_nucleatum_subsp._animalis_7_1;_CP007062.1	088%