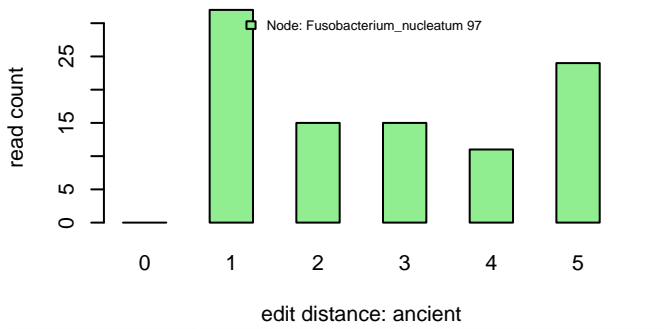
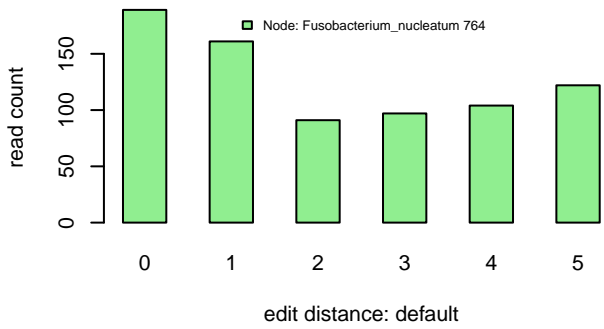
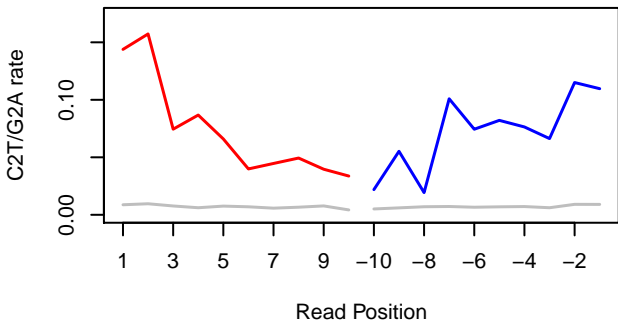


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for *Fusobacterium_nucleatum* node



Node	Fusobacterium_nucleatum
Top Reference	Fusobacterium_nucleatum_subsp._nucleatum
assigned reads	1270
all reads	466
nonDup	455
readDis	0.326
nonStacked	165
destacking?	Off
C>T_1	0.1438
G>A_-1	0.1097
n length (sd)	56 (18.288)

Fusobacterium_nucleatum_subsp._nucleatum;_CP012717.1	100%
bacterium_nucleatum_subsp._nucleatum_ATCC_25586;_AE009951.2	100%
bacterium_nucleatum_subsp._nucleatum_ChDC_F316;_CP012716.1	100%
Fusobacterium_nucleatum_subsp._animalis_4_8;_CP003723.1	100%
usobacterium_nucleatum_subsp._vincentii_ChDC_F8;_CP012714.1	100%
usobacterium_nucleatum_subsp._vincentii_3_1_36A2;_CP003700.1	100%
Fusobacterium_nucleatum_subsp._animalis;_CP012713.1	100%
Fusobacterium_nucleatum_subsp._animalis;_CP012715.1	100%
Fusobacterium_nucleatum_subsp._animalis;_CP022124.1	100%
usobacterium_nucleatum_subsp._vincentii_3_1_27;_CP007064.1	100%