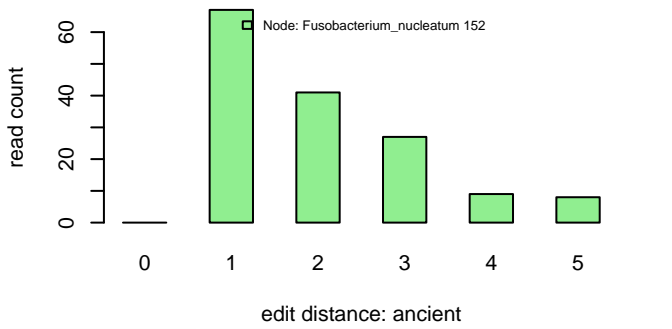
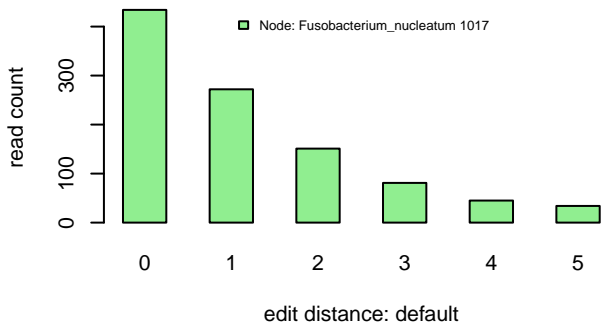
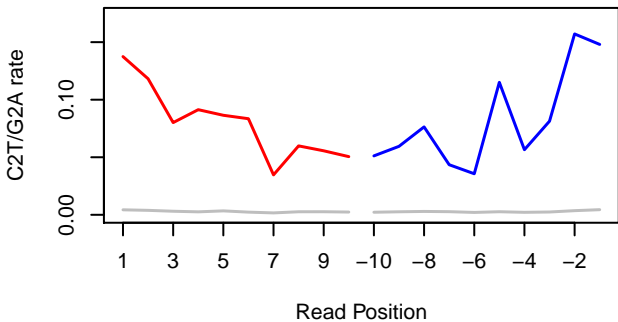


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Fusobacterium_nucleatum node



Node	Fusobacterium_nucleatum
Top Reference	Fusobacterium_nucleatum_subsp._nucleatum
assigned reads	1113
all reads	588
nonDup	573
readDis	0.94
nonStacked	525
destacking?	Off
C>T_1	0.1375
G>A_-1	0.1481
n length (sd)	55 (18.547)

Fusobacterium_nucleatum_subsp._nucleatum;_CP012717.1 100%
bacterium_nucleatum_subsp._nucleatum_ChDC_F316;_CP012716.1 100%
bacterium_nucleatum_subsp._nucleatum_ATCC_25586;_AE009951.2 100%
sobacterium_nucleatum_subsp._vincentii_3_1_36A2;_CP003700.1 091%
sobacterium_nucleatum_subsp._vincentii_ChDC_F8;_CP012714.1 090%
Fusobacterium_nucleatum_subsp._animalis;_CP012715.1 079%
Fusobacterium_nucleatum_subsp._animalis_7_1;_CP007062.1 078%
Fusobacterium_nucleatum_subsp._animalis_4_8;_CP003723.1 078%
sobacterium_nucleatum_subsp._vincentii_3_1_27;_CP007064.1 075%
Fusobacterium_nucleatum_subsp._animalis;_CP022124.1 074%