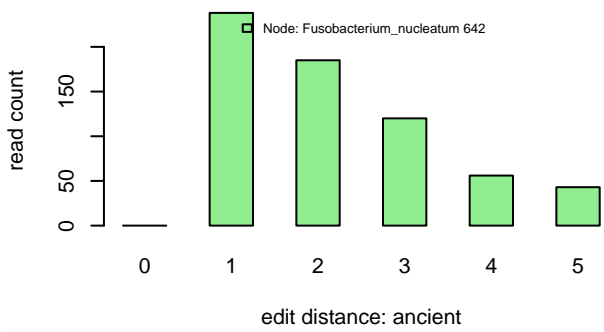
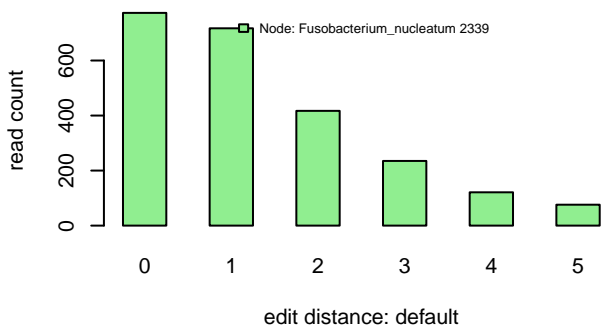
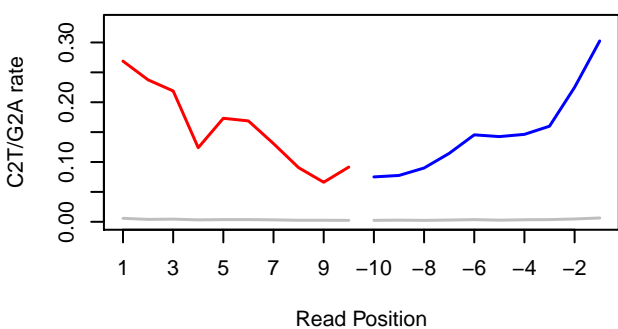


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for Fusobacterium_nucleatum node



Node	Fusobacterium_nucleatum
Top Reference	Fusobacterium_nucleatum_subsp._nucleatum
assigned reads	2455
all reads	1414
nonDup	1365
readDis	0.856
nonStacked	1089
destacking?	Off
C>T_1	0.2688
G>A_-1	0.3025
n length (sd)	48 (14.635)

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%
sobacterium_nucleatum_subsp._vincentii_3_1_36A2;_CP003700.1	100%
sobacterium_nucleatum_subsp._vincentii_ChDC_F8;_CP012714.1	100%
Fusobacterium_sp._oral_taxon_203;_CP016200.1	099%
usobacterium_nucleatum_subsp._vincentii_3_1_27;_CP007064.1	089%
Fusobacterium_nucleatum_subsp._animalis_7_1;_CP007062.1	084%
Fusobacterium_nucleatum_subsp._animalis;_CP022124.1	082%
Fusobacterium_nucleatum_subsp._animalis_4_8;_CP003723.1	081%