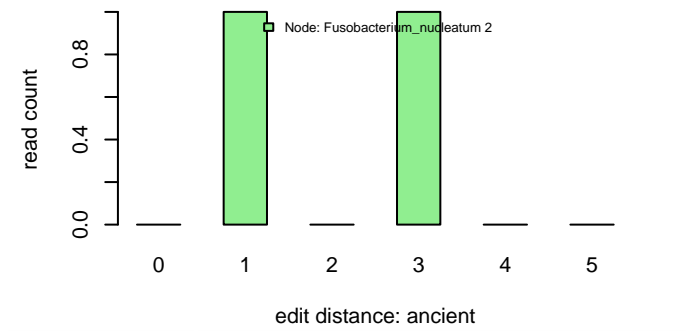
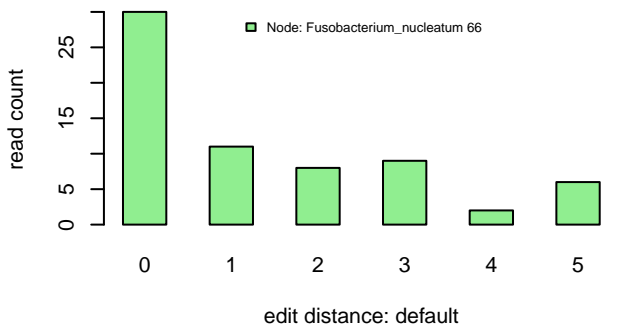
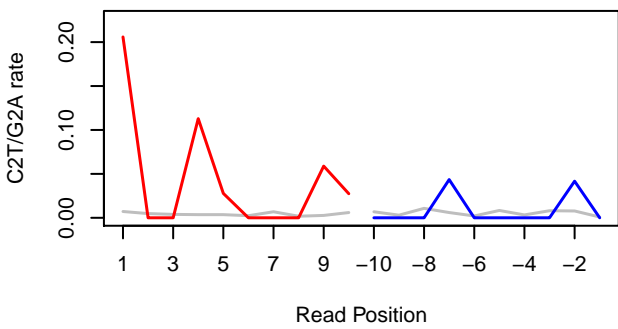


S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu



Damage plot for Fusobacterium_nucleatum node



Node	Fusobacterium_nucleatum
Top Reference	Fusobacterium_nucleatum_subsp._polymorphum
assigned reads	90
all reads	43
nonDup	43
readDis	0.966
nonStacked	41
destacking?	Off
C>T_1	0.2059
G>A_-1	0
length (sd)	68 (24.077)

Fusobacterium_nucleatum_subsp._polymorphum;_LN831027.1	100%
Fusobacterium_nucleatum_subsp._animalis_7_1;_CP007062.1	046%
Fusobacterium_nucleatum_subsp._animalis;_CP022124.1	043%
Fusobacterium_nucleatum_subsp._animalis;_CP012713.1	034%
bacterium_nucleatum_subsp._nucleatum_ChDC_F316;_CP012716.1	031%
Fusobacterium_nucleatum_subsp._nucleatum;_CP022122.1	031%
osobacterium_nucleatum_subsp._vincentii_ChDC_F8;_CP012714.1	024%
bacterium_nucleatum_subsp._nucleatum_ATCC_25586;_AE009951.2	021%
Fusobacterium_nucleatum_subsp._animalis;_CP012715.1	021%
Fusobacterium_nucleatum_subsp._animalis_4_8;_CP003723.1	019%