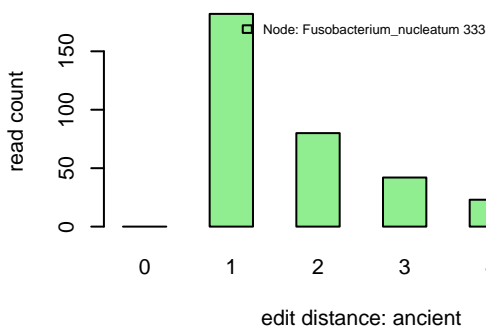
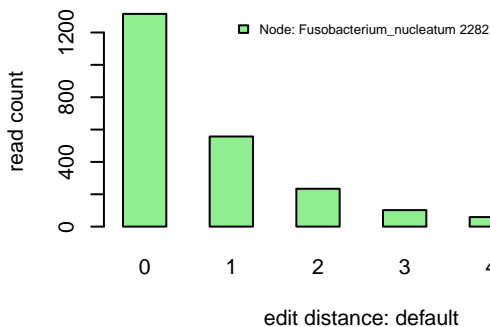
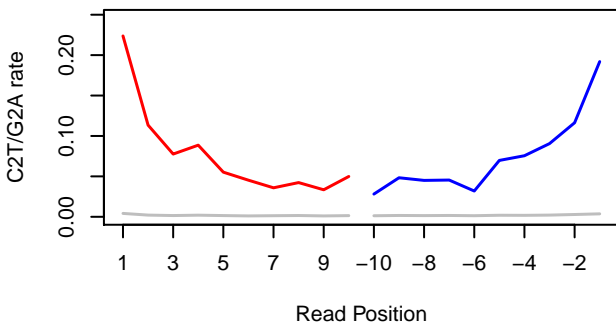


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur0101_S0_L000_R1_000.fastq.merged.prefixed.hg19ur



Damage plot for Fusobacterium_nucleatum node



Node	Fusobacterium_nucleatum
Top Reference	Fusobacterium_nucleatum_subsp._nucleatum
assigned reads	2327
all reads	1440
nonDup	1403
readDis	0.883
nonStacked	1155
destacking?	Off
C>T_1	0.2237
G>A_-1	0.192
n length (sd)	46 (13.952)

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	090%
sobacterium_nucleatum_subsp._vincentii_ChDC_F8;_CP012714.1	082%
Fusobacterium_sp._oral_taxon_203;_CP016200.1	081%
usobacterium_nucleatum_subsp._vincentii_3_1_27;_CP007064.1	076%
Fusobacterium_nucleatum_subsp._animalis_7_1;_CP007062.1	066%
Fusobacterium_nucleatum_subsp._animalis;_CP022124.1	064%
Fusobacterium_nucleatum_subsp._animalis;_CP012715.1	063%