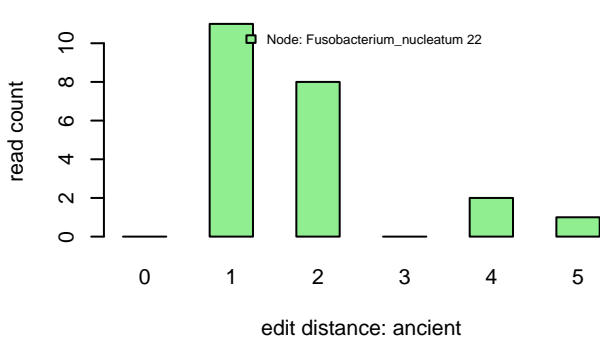
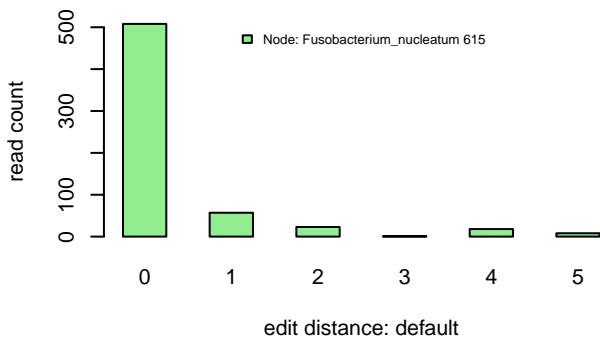
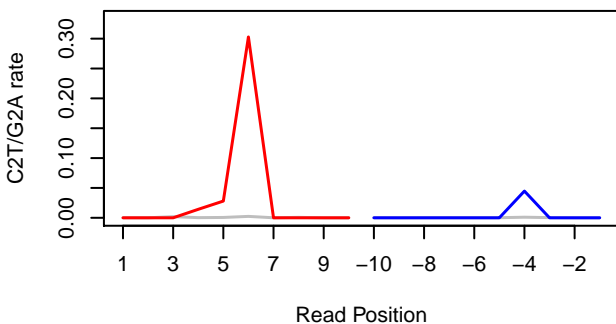


50\_L001\_R1\_001.fastq.combined.fq.prefixed.extractu50\_L001\_R1\_001.fastq.combined.fq.prefixed.extractu



### Damage plot for Fusobacterium\_nucleatum node



Node	Fusobacterium_nucleatum
Top Reference	Fusobacterium_nucleatum_subsp._nucleatum
assigned reads	627
all reads	464
nonDup	53
readDis	0.54
nonStacked	34
destacking?	Off
C>T_1	0
G>A_-1	0
n length (sd)	62 (18.303)

Fusobacterium_nucleatum_subsp._nucleatum;_CP012717.1 100%
bacterium_nucleatum_subsp._nucleatum_ATCC_25586;_AE009951.2 100%
usobacterium_nucleatum_subsp._vincentii_ChDC_F8;_CP012714.1 100%
bacterium_nucleatum_subsp._nucleatum_ChDC_F316;_CP012716.1 100%
sobacterium_nucleatum_subsp._vincentii_3_1_36A2;_CP003700.1 100%
usobacterium_nucleatum_subsp._vincentii_3_1_27;_CP007064.1 100%
Fusobacterium_nucleatum_subsp._animalis_4_8;_CP003723.1 100%
Fusobacterium_sp._oral_taxon_203;_CP016200.1 100%
Fusobacterium_nucleatum_subsp._animalis;_CP022124.1 100%
Fusobacterium_nucleatum_subsp._animalis_7_1;_CP007062.1 100%