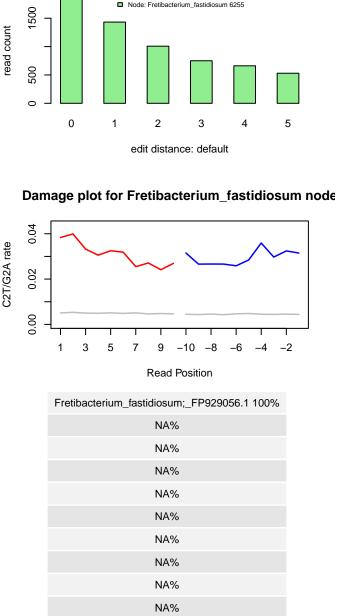
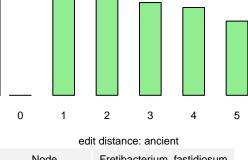
S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu Node Fretibacterium_fastidiosum 370

read count





euit distance, ancient	
Node	Fretibacterium_fastidiosum
Top Reference	Fretibacterium_fastidiosum
assigned reads	278009
all reads	10000
nonDup	9886
readDis	0.541
nonStacked	3425
destacking?	Off
C>T_1	0.0384
G>A1	0.0315
mean length (sd)	107 (41.271)