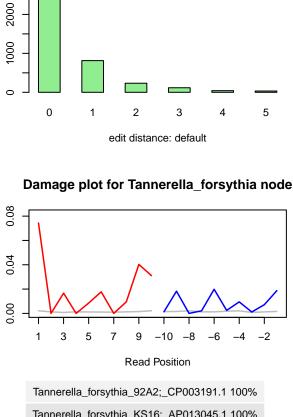
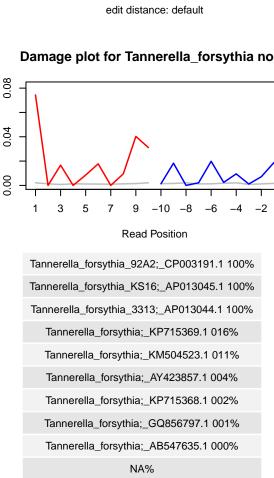
60_L001_R1_001.fastq.combined.fq.prefixed.extractu60_L001_R1_001.fastq.combined.fq.prefixed.extractu ■ Node: Tannerella_forsythia 3970 ■ Node: Tannerella_forsythia 98

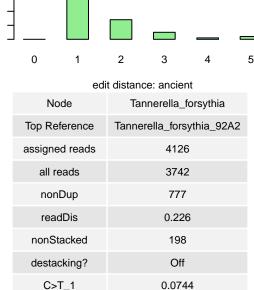
read count 4

20



C2T/G2A rate





0.0187

76 (24.979)

G>A_-1

mean length (sd)