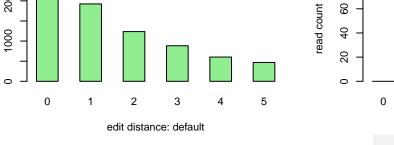
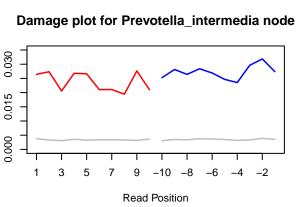


0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un0101_S0_L000_R1_000.fastq.merged.prefixed.hg19un





C2T/G2A rate

Prevotella_intermedia;_CP019302.1 100% Prevotella_intermedia_17;_CP003503.1 100%

revotella_intermedia_ATCC_25611_=_DSM_20706;_CP019300.1 100%

Prevotella_intermedia_17;_CP003502.1 048%

revotella_intermedia_ATCC_25611_=_DSM_20706;_CP019301.1 034%

Porphyromonas_gingivalis;_CP011995.1 001%

Prevotella_fusca_JCM_17724;_CP012074.1 001%

Prevotella_dentalis_DSM_3688;_CP003369.1 001%

Prevotella_denticola_F0289;_CP002589.1 000%

Prevotella_fusca_JCM_17724;_CP012075.1 000%

0 1		2	3	4	
edit distance: ancient					
Node		Prevotella_intermedia			
Top Reference		Prevotella_intermedia			
assigned reads		28062			
all reads		4960			
nonDup		4880			
readDis		0.496			
nonStacked		2054			
destacking?		Off			
C>T_1		0.0264			
G>A1		0.0274			
mean length (sd)		8	89 (28.388)		