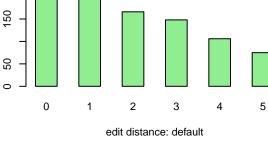
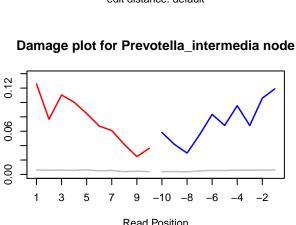


30

10

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





C2T/G2A rate



Prevotella_intermedia;_AP014925.1 100%

Prevotella_intermedia;_CP019302.1 100%

Prevotella_intermedia;_AP014597.1 100%

Prevotella_dentalis_DSM_3688;_CP003369.1 001%

revotella_intermedia_ATCC_25611_=_DSM_20706;_CP019300.1 083%

Prevotella_intermedia;_AP014598.1 051% revotella_intermedia_ATCC_25611_=_DSM_20706;_CP019301.1 036%

Prevotella_intermedia;_CP019303.1 022%

Prevotella_intermedia;_AP014926.1 021%

0 2 3 edit distance: ancient Node Prevotella_intermedia Prevotella_intermedia_17 Top Reference assigned reads 1179 all reads 532 nonDup 525 readDis 0.719 nonStacked 328 destacking? Off C>T_1 0.1257

G>A_-1

mean length (sd)

0.1189

60 (21.399)