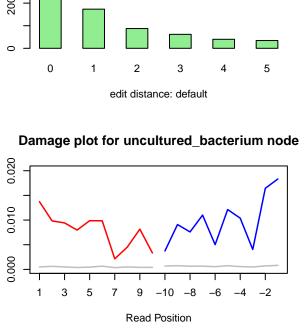
Node: uncultured_bacterium 413 ■ Node: uncultured_bacterium 8678 read count 2000

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



C2T/G2A rate

edit distance: default
Damage plot for uncultured_bacterium node
0000 0000 0000 0000 0000 0000 0000 0000 0000
Read Position
uncultured_bacterium;_KJ910972.1 100%
Neisseria_elongata;_AJ239303.1 100%
Abiotrophia_defectiva;_AY879307.1 100%
Abiotrophia_defectiva;_AY879306.1 100%
uncultured_Neisseriaceae_bacterium;_FJ976340.1 100%
Neisseria_elongata;_AJ239297.1 100%
uncultured_Abiotrophia_sp.;_AM420170.1 100%
Neisseria_elongata_subspglycolytica;_AY167422.1 100%
Abiotrophia_defectiva;_AY879308.1 100%

Neisseria_elongata_subsp._elongata;_NR_025893.1 100%

0	1	2		3		4	
edit distance: ancient							
Node			uncultured_bacterium				
Top Reference			uncultured_bacterium				
assigned reads			60240				
all reads			515				
nonDup			508				
readDis			0.005				
nonStacked			1				
destacking?			Off				
C>T_1			0.0137				
G>A1			0.0183				
mean length (sd)			93 (27.671)				