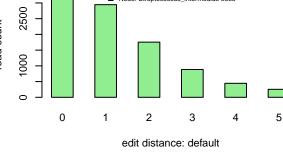
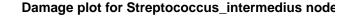
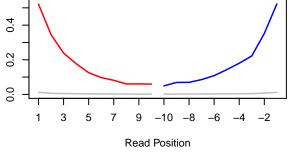
1500 ■ Node: Streptococcus_intermedius 9663 Node: Streptococcus_intermedius 4012 1000 ead count

500

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







C2T/G2A rate

Streptococcus_intermedius;_AP014880.1 100%

Streptococcus_intermedius_B196;_CP003857.1 100%

Streptococcus_intermedius_C270;_CP003858.1 100% Streptococcus_intermedius_JTH08;_AP010969.1 100%

Streptococcus_sanguinis_SK36;_CP000387.1 001%

Streptococcus_cristatus_AS_1.3089;_CP004409.1 000%

Streptococcus_anginosus_subsp._whileyi_

Streptococcus_anginosus_C238;_CP003861.1 000%

MAS624;	_AP013072.1

000%

Streptococcus_anginosus	;_KY315478.1 000%

Chrophododda_anginoddd;_ittrofoffoffoff	
Streptococcus_anginosus;_CP012719.1 000%	

