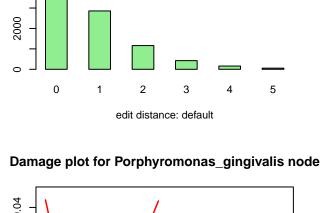
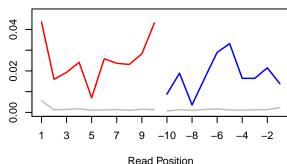


0519_S0_L000_R1_000.fastq.merged.prefixed.hg19un0519_S0_L000_R1_000.fastq.merged.prefixed.hg19un





C2T/G2A rate

0.00 0.02 0.04		<u> </u>	\bigvee			/ /	\bigvee		_	<u> </u>	
0	1	3	1 5	1 7	9	–10	-8	і –6	1 –4	1 –2	
Read Position											
Porphyromonas_gingivalis;_CP011995.1 100%											
Porphyromonas_gingivalis_W83;_AE015924.1 100%											
Po	rphyr	omor	nas_g	ingiva	alis_/	AJW4;	_CP0	01199	96.1 1	00%	
Por	rphyr	omon	as_gi	ngiva	ılis_T	DC60	;_AP	0122	03.1 1	100%	
Porp	hyro	mona	s_gin	givali	s_A7	7A1–2	8;_CI	P013	131.1	100%	
Р	orphy	/romc	nas_	gingi	valis_	_381;_	CP0	12889	9.1 10	0%	
Porph	yrom	onas_	_gingi	valis_	_ATC	C_332	277;_	AP00	9380	.1 100%	
Porphyromonas_gingivalis_ATCC_33277;_AB015879.1 033%											

Porphyromonas_gingivalis_TDC60;_AB757136.1 014% Tannerella_forsythia_92A2;_CP003191.1 012%

0	1	2	3	4				
edit distance: ancient								
N	lode	Porphyromonas_gingivalis						
Top R	eference	Porphyromonas_gingivalis						
assign	ned reads	132169						
all	reads	7630						
no	nDup	2614						
rea	adDis	0.284						
nons	Stacked	794						
dest	acking?	Off						
C	>T_1	0.0436						
G>	-A1	0.0138						
mean l	ength (sd)	54 (17.996)						