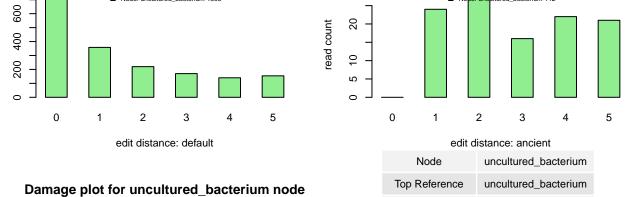
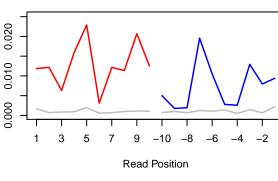
## ■ Node: uncultured\_bacterium 112 ■ Node: uncultured\_bacterium 1805 read count

0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur





C2T/G2A rate

1	3	5	7	9	-10	-8	-6	-4	-
			ı	Read	l Posit	ion			
u	ncult	ured_	bacte	rium	;_KJ9	1097	5.1 10	00%	
unio	dentifi	ed_pı	oteol	oacte	erium;_	_X732	223.1	1009	%
und	culture	ed_La	utrop	ia_s	p.;_AY	'8070	05.1	100%	, o
	Lautr	opia_	mirab	ilis;_	GU39	7891	.1 100	0%	
unc	culture	ed_La	utrop	ia_s <sub> </sub>	p.;_HC	26162	276.1	100%	6
und	culture	ed_La	utrop	ia_s	p.;_AY	′8074	08.1	100%	, o
		Bacte	eria;_	FJ97	6403.	1 100	)%		
unc	culture	ed_La	utrop	ia_s <sub> </sub>	p.;_DC	20167	727.1	100%	6
un	cultur	ed_La	autrop	oia_s	p.;_AJ	14917	77.1	100%	, O
	Lautr	opia_i	mirab	ilis;_	GU39	7890	.1 100	0%	

0	1	2	2	3	4			
	e	edit di	stance	: ancie	nt			
	Node		uncultured_bacterium					
Тор	Referen	се	uncultured_bacterium					
assi	gned rea	ds	2303					
a	all reads		84					
1	nonDup		83					
ı	readDis		0.147					
no	nStacked	t	4					
destacking?			Off					
C>T_1			0.0119					
G>A1			0.0094					
mear	n length (	sd)		67 (21.	547)			