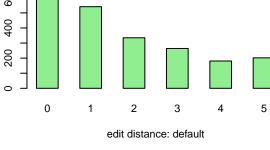
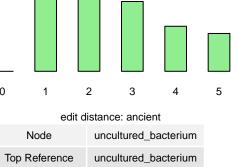
Node: uncultured_bacterium 2228 OO 1 Node: uncultured_bacterium 537

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





D	ama	ige p	OIOT	tor t	ıncı	uitur	ea_i	oact	eriu	m no	ae
0.0 0.1 0.2 0.3				_/	_	_		^			
0		ı	ı	ı	ı		ı	ı			_
	1	3	5	7	9	-10	-8	-6	-4	-2	
Read Position											
uncultured_bacterium;_JQ075394.1 100%											

C2T/G2A rate

0	1	2	3	4	Ę					
edit distance: ancient										
	Node	un	uncultured_bacterium							
Тор	Reference	un	uncultured_bacterium							
assi	gned reads		2706							
а	ıll reads		235							
r	nonDup		225							
r	eadDis		0.001							
no	nStacked		0							
de	stacking?		Off							
	C>T_1		0.33	12						
C	G>A1		0.263	33						
mear	length (sd)		56 (22.	705)						

ncultured_Candidatus_Saccharibacteria_bacterium;_AM420132.1 100% ncultured_Candidatus_Saccharibacteria_bacterium;_AY144358.1 100% ncultured_Candidatus_Saccharibacteria_bacterium;_HQ616339.1 100%

ncultured_Candidatus_Saccharibacteria_bacterium;_HM215448.1 100%

incultured_Candidatus_Saccharibacteria_bacterium;_FJ976255.1 100%

ncultured_Candidatus_Saccharibacteria_bacterium;_GU227159.1 100%
ncultured_Candidatus_Saccharibacteria_bacterium;_HM215444.1 100%

ncultured_Candidatus_Saccharibacteria_bacterium;_HM215444.1 100% uncultured_Actinomyces_sp.;_FJ976431.1 100%

uncultured_bacterium_AH040;_AF125204.1 100%