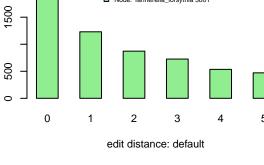
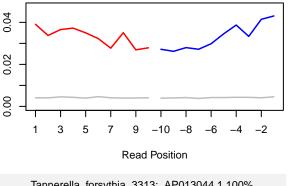
Node: Tannerella_forsythia 342 ■ Node: Tannerella_forsythia 5801

read count

S0_L001_R1_001.fastq.combined.fq.prefixed.extractuS0_L001_R1_001.fastq.combined.fq.prefixed.extractu







Tarinerella_lorsythla_5515,_AP015044.1 100%	
Tannerella_forsythia_KS16;_AP013045.1 100%	

Tannerella_forsythia_92A2;_CP003191.1 100%

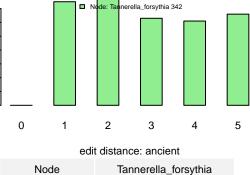
Tannerella_forsythia;_NR_076943.1 023% Tannerella_forsythia;_AB587729.1 015%

Tannerella_sp._oral_taxon_HOT-286;_CP017038.1 003% Tannerella_forsythia;_AJ006516.1 001%

uncultured_bacterium;_JQ999790.1 000%

Tannerella_forsythia;_AF054892.1 000%

uncultured_bacterium,_bQ393730.1 00	JU 76
uncultured_bacterium;_GU926679.1 0	00%



eui	edit distance: ancient		
Node	Tannerella_forsythia		
Top Reference	Tannerella_forsythia_3313		
assigned reads	7975		
all reads	5921		
nonDup	5766		
readDis	0.052		
nonStacked	233		
destacking?	Off		
C>T_1	0.0391		
G>A1	0.043		
mean length (sd)	96 (38.303)		