

Sequences producing significant alignments:

Select: [All](#) [None](#) Selected:0

Alignments Download GenBank Graphics Distance tree of results							
Description	Max score	Total score	Query cover	E value	Ident	Accession	
<input type="checkbox"/> Tatumella sp. UAPS07002 16S ribosomal RNA gene, partial sequence	774	774	99%	0.0	99%	EF012364.1	
<input type="checkbox"/> Pantoea punctata strain DSM 13700 16S ribosomal RNA gene, partial sequence	758	758	99%	0.0	99%	FJ756351.1	
<input type="checkbox"/> Tatumella punctata strain SHS 2003 16S ribosomal RNA, complete sequence >gb JN175342.1 Pantoea puncta	754	754	99%	0.0	99%	NR_104937.1	
<input type="checkbox"/> Citrobacter sp. F3-1 16S ribosomal RNA gene, partial sequence	752	752	99%	0.0	98%	EF491829.1	
<input type="checkbox"/> Pantoea terrea strain DSM 13701 16S ribosomal RNA gene, partial sequence	750	750	99%	0.0	98%	FJ756353.1	
<input type="checkbox"/> Tatumella pyseos strain LMG7888 16S ribosomal RNA gene, complete sequence	749	749	95%	0.0	99%	EU344770.1	
<input type="checkbox"/> Citrobacter sp. BinzhouCLT 16S ribosomal RNA gene, partial sequence	749	749	100%	0.0	98%	AF530068.2	
<input type="checkbox"/> Citrobacter freundii strain Y9 16S ribosomal RNA gene, partial sequence	747	747	99%	0.0	98%	KF781347.1	
<input type="checkbox"/> Citrobacter freundii strain Sn4 16S ribosomal RNA gene, partial sequence	747	747	99%	0.0	98%	KC683722.1	
<input type="checkbox"/> Citrobacter murlinae strain FFL15 16S ribosomal RNA gene, partial sequence	747	747	99%	0.0	98%	JN092600.1	
<input type="checkbox"/> Citrobacter sp. F3-4 16S ribosomal RNA gene, partial sequence	747	747	99%	0.0	98%	EF491832.1	
<input type="checkbox"/> Bacterium EMB233-1 partial 16S rRNA gene, strain EMB233-1	743	743	99%	0.0	98%	HG792123.1	
<input type="checkbox"/> Citrobacter freundii strain E087 16S ribosomal RNA gene, partial sequence	743	743	99%	0.0	98%	JX267103.1	
<input type="checkbox"/> Citrobacter sp. F3-7 16S ribosomal RNA gene, partial sequence	743	743	99%	0.0	98%	EF491835.1	
<input type="checkbox"/> Citrobacter sp. F3-6 16S ribosomal RNA gene, partial sequence	743	743	99%	0.0	98%	EF491834.1	
<input type="checkbox"/> Citrobacter sp. F3-3 16S ribosomal RNA gene, partial sequence	743	743	99%	0.0	98%	EF491831.1	
<input type="checkbox"/> Citrobacter sp. F1-3 16S ribosomal RNA gene, partial sequence	743	743	99%	0.0	98%	EF491827.1	
<input type="checkbox"/> Citrobacter sp. F1-1 16S ribosomal RNA gene, partial sequence	743	743	99%	0.0	98%	EF491825.1	
<input type="checkbox"/> Citrobacter sp. LAR-1 16S ribosomal RNA gene, partial sequence	741	741	99%	0.0	98%	KC211014.1	
<input type="checkbox"/> Citrobacter braakii strain UAAD6 16S ribosomal RNA gene, partial sequence	741	741	99%	0.0	98%	JX518488.1	
<input type="checkbox"/> Bacterium NTL113 16S ribosomal RNA gene, partial sequence	741	741	99%	0.0	98%	JQ779035.1	
<input type="checkbox"/> Enterobacter sp. C6-8 16S ribosomal RNA gene, complete sequence	741	741	100%	0.0	98%	JF939051.1	
<input type="checkbox"/> Citrobacter braakii strain A8 16S ribosomal RNA gene, partial sequence	741	741	99%	0.0	98%	HQ288930.1	
<input type="checkbox"/> Citrobacter sp. A7(2011) strain A7 16S ribosomal RNA gene, partial sequence	741	741	99%	0.0	98%	HQ288923.1	
<input type="checkbox"/> Citrobacter sp. 40 16S ribosomal RNA gene, partial sequence	741	741	99%	0.0	98%	HQ399664.1	
<input type="checkbox"/> Citrobacter freundii strain H1 16S ribosomal RNA-like gene, partial sequence	741	741	99%	0.0	98%	FJ768455.1	
<input type="checkbox"/> Citrobacter werkmanii strain IITRL7 16S ribosomal RNA gene, partial sequence	741	741	99%	0.0	98%	FJ581027.1	