

Descriptions | Graphic Summary | Alignments | Taxonomy

Sequences producing significant alignments

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GenBank | Graphics | Distance tree of results | **New** MSA Viewer

	Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Per. Ident	Acc. Len	Accession
<input type="checkbox"/>	Clostridium thiosulfatireducens strain LUP 21 16S ribosomal RNA, partial sequence	Clostridium thiosulfatireducens	2538	2538	99%	0.0	99.93%	1448	NR_042718.1
<input type="checkbox"/>	Clostridium subterminale strain JCM 1417 16S ribosomal RNA, partial sequence	Clostridium subterminale	2527	2527	99%	0.0	99.78%	1470	NR_113027.1
<input type="checkbox"/>	Clostridium thiosulfatireducens strain LUP 21 16S ribosomal RNA, partial sequence	Clostridium thiosulfatireducens	2521	2521	99%	0.0	99.71%	1447	NR_114590.1
<input type="checkbox"/>	Clostridium subterminale strain DSM 6970 16S ribosomal RNA, partial sequence	Clostridium subterminale	2521	2521	99%	0.0	99.71%	1504	NR_041795.1
<input type="checkbox"/>	Clostridium subterminale strain DSM 6970 16S ribosomal RNA, partial sequence	Clostridium subterminale	2519	2519	99%	0.0	99.64%	1452	NR_112653.1
<input type="checkbox"/>	Clostridium sulfidigenes strain SGB2 16S ribosomal RNA, partial sequence	Clostridium sulfidigenes	2519	2519	99%	0.0	99.71%	1504	NR_044161.1
<input type="checkbox"/>	Clostridium thiosulfatireducens strain DSM 13105 16S ribosomal RNA, partial sequence	Clostridium thiosulfatireducens	2422	2422	95%	0.0	99.77%	1357	NR_112656.1
<input type="checkbox"/>	Clostridium huakuii strain LAM1030 16S ribosomal RNA, partial sequence	Clostridium huakuii	2335	2335	99%	0.0	97.25%	1394	NR_134006.1
<input type="checkbox"/>	Clostridium tunisiense strain TJ 16S ribosomal RNA, partial sequence	Clostridium tunisiense	2324	2324	99%	0.0	97.10%	1475	NR_115161.1
<input type="checkbox"/>	Clostridium punense strain BLPYG-8 16S ribosomal RNA, partial sequence	Clostridium punense	2283	2283	99%	0.0	96.59%	1497	NR_145903.1
<input type="checkbox"/>	Clostridium vulturis strain YMB-57 16S ribosomal RNA, partial sequence	Clostridium vulturis	2281	2281	99%	0.0	96.53%	1412	NR_148266.1