

BLAST Results

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Job title: 16S_rRNA::tig000000001:541494-543044(+)

RID [BFKZ7J7G013](#) (Expires on 06-04 02:37 am)
Query ID lcl|Query_484981
Description 16S_rRNA::tig000000001:541494-543044(+)
Molecule type dna
Query Length 1550

Database Name nt
Description Nucleotide collection (nt)
Program BLASTN 2.12.0+ [▶ Citation](#)

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Sequences producing significant alignments:

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

<div>⌵ Alignments <div>Download</div> GenBank Graphics Distance tree of results</div>						
	Description	Max Score	Total Score	Query Cover	E value	Per. Ident
<input type="checkbox"/>	Bacillus sp. RZ2MS9 chromosome, complete genome	2863	39921	100%	0.0	100.00% CP049978.1
<input type="checkbox"/>	Bacillus cereus strain K8, complete sequence	2863	39943	100%	0.0	100.00% CP016595.1
<input type="checkbox"/>	Bacillus thuringiensis strain BM-BT15426, complete genome	2863	39842	100%	0.0	100.00% CP020723.1
<input type="checkbox"/>	Bacillus cereus strain A1, complete genome	2863	39955	100%	0.0	100.00% CP015727.1
<input type="checkbox"/>	Bacillus thuringiensis strain HER1410 chromosome, complete genome	2857	39966	100%	0.0	99.94% CP050183.1
<input type="checkbox"/>	Bacillus cereus strain FDAARGOS_798 chromosome, complete genome	2857	39921	100%	0.0	99.94% CP053954.1
<input type="checkbox"/>	Bacillus cereus strain FDAARGOS_799 chromosome, complete genome	2857	39921	100%	0.0	
<input type="checkbox"/>	Bacillus cereus strain WDC614/2 chromosome, complete genome	2857	39966	100%	0.0	