

Job Title S-05002_oECA306_A03.seq

RID [DFKMH75701R](#) Search expires on 06-05 00:23 am [Download All](#) ▼

Results for ① 37:lc|Query_44458 S-05554_oECA306_D03.seq(1024bp) ▼

Program BLASTN [?](#) [Citation](#) ▼

Database nt [See details](#) ▼

Query ID lc|Query_44458

Description ② S-05554_oECA306_D03.seq

Molecule type dna

Query Length 1024

Other reports [Distance tree of results](#) [?](#)

Filter Results

Organism only top 20 will appear ☐ exclude

Type common name, binomial, taxid or group name

[+ Add organism](#)

Percent Identity

 to

E value

 to

Query Coverage

 to

Filter

Reset

Descriptions

Graphic Summary

Alignments

Taxonomy

Sequences producing significant alignments

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☒ select all 100 sequences selected

[GenBank](#)

[Graphics](#)

[Distance tree of results](#)

	Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession
③							
☒	Caenorhabditis briggsae 18S ribosomal RNA gene, internal transcribed spacer 1, 5.8S ribosomal RNA gene, internal transcribed spacer 2	1288	1288	68%	0.0	99.43%	JN636061.1
☒	Caenorhabditis briggsae strain AF16 18S ribosomal RNA gene, internal transcribed spacer 1, 5.8S ribosomal RNA gene, internal transcribed	1284	1284	68%	0.0	99.43%	MN519141.1