

Séquence 1

Descriptions

Graphic Summary

Alignments

Taxonomy

Sequences producing significant alignments

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[Graphics](#)

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	Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession
☒	Temora stylifera isolate H04 TstCOL*42 04 R cytochrome c oxidase subunit 1 (COI) gene, partial cds; mitochondrial	1232	1232	100%	0.0	100.00%	MH159162.1
☒	Temora stylifera isolate G04 TstCOL*42 03 R cytochrome c oxidase subunit 1 (COI) gene, partial cds; mitochondrial	1232	1232	100%	0.0	100.00%	MH159161.1
☒	Temora stylifera isolate E04 TstCOL*42 01 R cytochrome c oxidase subunit 1 (COI) gene, partial cds; mitochondrial	1232	1232	100%	0.0	100.00%	MH159159.1
☒	Temora stylifera isolate F04 TstCOL*42 02 R cytochrome c oxidase subunit 1 (COI) gene, partial cds; mitochondrial	1221	1221	100%	0.0	99.70%	MH159160.1
☒	Temora stylifera voucher BIOUG<CAN>:MxTem002 cytochrome oxidase subunit 1 (COI) gene, partial cds; mitochondrial	1164	1164	95%	0.0	99.69%	DQ889132.1
☒	Temora stylifera cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial	1136	1136	98%	0.0	97.73%	MF098548.1
☒	Temora stylifera voucher P0016 cytochrome oxidase subunit I (COI) gene, partial cds; mitochondrial	1134	1134	98%	0.0	97.72%	KP861423.1
☒	Parvocalanus crassirostris isolate ICAR-CMFRI_Pc1 cytochrome c oxidase subunit I (COI) gene, partial cds; mitochondrial	556	556	100%	4e-154	81.82%	MK387708.1
☒	Parvocalanus crassirostris isolate ICAR-CMFRI_Pc1 cytochrome c oxidase subunit I (COI) gene, partial cds; mitochondrial	556	556	100%	4e-154	81.82%	MK387708.1



Temora stylifera

Crustacé, copépode

Séquence 2

BLAST[®] » blastn suite » results for RID-NPDNZ9NX014

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Job Title Nucleotide Sequence

RID [NPDNZ9NX014](#) Search expires on 09-12 16:35 pm [Download All](#) ▼

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Database nt [See details](#) ▼

Query ID lcl|Query_37261

Description None

Molecule type dna

Query Length 240

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Percent Identity to

E value to

Query Coverage to

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⚠ No significant similarity found. For reasons why, [click here](#)

Séquence trop courte ?

Trop fréquente donc non significative?

Séquence 3

Descriptions

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	Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession
☒	Posidonia oceanica mitogen-activated protein kinase kinase kinase gene, partial cds	830	830	100%	0.0	100.00%	JF811736.1

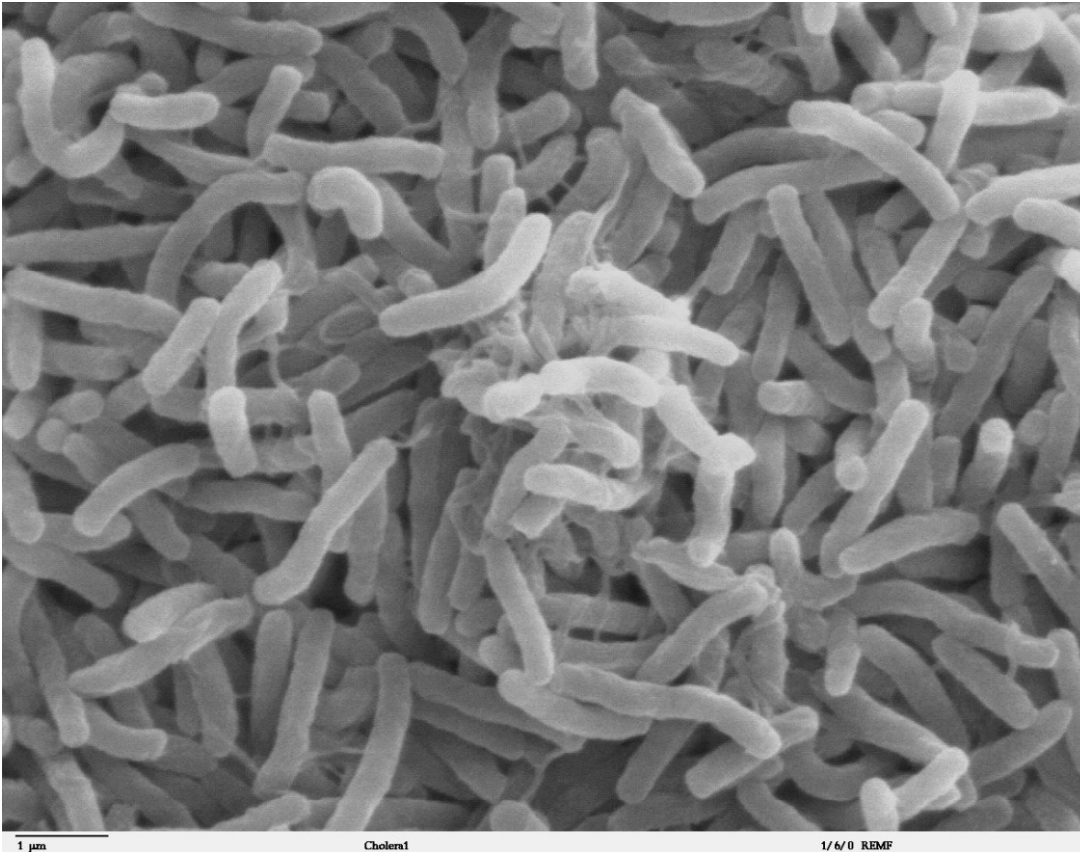


posidonia

Plante à fleur sous marine

Séquence 4

	Description	Max Score	Total Score
✓	Vibrio cholerae C6706 chromosome 1, complete sequence	1441	28
✓	Vibrio cholerae O1 biovar El Tor strain C6709 chromosome 1, complete sequence	1441	28
✓	Vibrio cholerae O1 biovar El Tor strain P27459 chromosome 1, complete sequence	1441	43
✓	Vibrio cholerae O1 biovar El Tor strain DRC-193A chromosome 1, complete sequence	1441	28
✓	Vibrio cholerae O1 biovar El Tor strain E7946 chromosome 1, complete sequence	1441	71
✓	Vibrio cholerae O1 strain AAS91 chromosome 1, complete sequence	1441	43
✓	Vibrio cholerae isolate CTMA_1441 chromosome 1, complete sequence	1441	28
✓	Vibrio cholerae strain F9993 chromosome 1, complete sequence	1441	57
✓	Vibrio cholerae C6706 chromosome 1, complete sequence	1441	28
✓	Vibrio cholerae strain 2010EL-1786 chromosome 1	1441	28
✓	Vibrio cholerae O1 biovar El Tor str. N16961 genome assembly, chromosome: 1	1441	28
✓	Vibrio cholerae strain E4 chromosome 1, complete sequence	1441	43
✓	Vibrio cholerae strain C1 CTX phage region genomic sequence	1441	72
✓	Vibrio cholerae strain No341 chromosome 1 phage sequence	1441	28



Vibrio cholerae

Bacille du choléra = bactérie

Présente sur les microplastiques
d'origine humaine