

Complete sequence of mitochondrial DNA of *Gracilaria edulis* (Rhodophyta)

<b>Publons ID</b>	41476327
<b>Wos ID</b>	WOS:000519283400001
<b>Doi</b>	10.1080/23802359.2017.1422413
<b>Title</b>	Complete sequence of mitochondrial DNA of <i>Gracilaria edulis</i> (Rhodophyta)
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<b>Publish Date</b>	APR 2 2020
<b>Journal Name</b>	MITOCHONDRIAL DNA PART B-RESOURCES
<b>Citation</b>	2
<b>Abstract</b>	Herein, the complete <i>Gracilaria edulis</i> mitogenome was determined and analysed. The complete <i>G. edulis</i> mitogenome length was 25,708 bp. The mitogenome contains 50 genes, including 24 protein-coding, two rRNA, and 23 tRNA genes and one unidentified open reading frame (ORF). Of the 24 protein-coding genes, 23 (95.83%) ended with the TAA stop codon, and one (4.17%) with TAG (rpl20). All protein-coding genes in <i>G. edulis</i> were concluded to use the start codon ATG. Mitogenome phylogenetic analysis revealed that <i>G. edulis</i> clustered together with <i>G. changii</i> and <i>G. salicornia</i> . The complete mitogenome sequence provided herein would help understand <i>Gracilaria</i> evolution.
<b>Publish Type</b>	Journal
<b>Publish Year</b>	2020
<b>Page Begin</b>	1128
<b>Page End</b>	1129
<b>Issn</b>	
<b>Eissn</b>	2380-2359
<b>Url</b>	<a href="https://www.webofscience.com/wos/woscc/full-record/WOS:000519283400001">https://www.webofscience.com/wos/woscc/full-record/WOS:000519283400001</a>
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