





http://dx.doi.org/10.11646/phytotaxa.222.3.3

## A New Tropical Species of Gracilariaceae (Rhodophyta, Gracilariales): *Gracilaria silviae sp. nov*.

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## Abstract

A new flat species of *Gracilaria*, named *Gracilaria silviae sp. nov.*, from the western Atlantic Ocean is described based on comparative morphological and molecular analyses of three molecular markers: the gene encoding cytochrome oxidase I (*cox*1), the Universal Plastid Amplicon (UPA) and the gene encoding the large subunit of ribulose-1, 5-bisphosphate carboxylase oxygenase (*rbcL*). *Gracilaria silviae* has a wide geographic distribution from the Indian River, Florida, USA, to Santa Catarina, southern Brazil, and a wide range of habit phenotypes. Across this geographic range *Gracilaria silviae* occupies a broad range of habitats, from warm temperate protected estuarine sublittoral areas to exposed tropical intertidal rocky shores and coastal reefs. The most morphologically similar species to *G. silviae* are not only the phylogenetically closest taxa to this species but also sympatric along its geographic range, i.e. *G. tikvahiae* and *G. isabellana* (formerly known as *G. lacinulata* in the northwestern Atlantic and *G. tepocensis* in southern Brazil).

Keywords: Barcoding, Red algae, Systematics, Taxonomy

## Introduction

The benthic marine red macroalgal genus *Gracilaria* Greville (1830:121) is the largest genus in the family Gracilariaceae (Gracilariales) with more than 120 currently accepted species worldwide (Guiry & Guiry 2015). Recent molecularly assisted taxonomic revisions of the order and family revealed the existence of six distinct genera, including the redefinition of *Gracilaria* to include taxa traditionally placed in the former genus *Hydropuntia* Montagne (1842:7) (Lyra *et al.* 2015). The species richness of *Gracilaria* is concentrated in tropical and sub-tropical zones, and is characterized by species bearing henriquesiana- and textorii-types of spermatangial conceptacles (*sensu* Yamamoto 1975, 1978, 1984; see also Bird 1995). The genus can also be superficially classified into flat and cylindrical species, where the former habit morphology holds the largest number of species, and has been recognized as the most taxonomically challenging group (Yamamoto 1984).

Taxonomic and species identification in *Gracilaria* based solely on morphological characters is particularly challenging due to their morphological simplicity, high levels of phenotypic plasticity, and large number of species (Gurgel & Fredericq 2004). Only with the advent and widespread use of molecular analyses based on DNA sequence data have the taxonomy, diversity, biogeography and the phylogenetics of the genus becoming better known (e.g. Gurgel *et al.* 2004a, Muangmai *et al.* 2014a,b, Lyra *et al.* 2015). The genus is also recognized for its economical importance not only as the current main source of agar around the world (Zemke-White & Ohno 1999, Armisen *et al.* 2009), but also due to other medical research applications such as antiviral, anti-inflammatory and anti-hypertensive