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New record of the non-native seaweed Gracilaria parvispora in Baja California – A note on Vergara-Rodarte et al. (2016)

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Abstract – The delimitation of species in the Gracilariales is often difficult due to the lack of diagnostic morphological characters. As a result, non-native species are often misidentified without the use of molecular tools. Recently, studies have investigated the agar quality of the dominant gracilarioid species in the Laguna San Ignacio in Baja Califronia Sur, Mexico, including Vergara-Rodarte *et al.*, (2016) published in the May edition of *Cryptogamie*, *Algologie*. The species has been reported as *Gracilaria pacifica*, *G. vermiculophylla* and *Gracilariopsis* sp. Using a combination of three mitochondrial and plastid markers, we identified this species as *G. parvispora*, extending the known distribution of this non-native species in Baja California. Due to the potential of *G. parvispora* as a source of agar, more detailed studies of on the invasion history and surveys are necessary in the eastern Pacific to determine its current distribution and impacts on the native biodiversity.

Baja California / Biological invasión / cox1 / cox2-3 spacer / Gracilaria / rbcL / Species delimitation

The delimitation of gracilarioid algae is notoriously fraught with problems due to the lack of diagnostic morphological characters (Bird, 1995) and, as a result, can often lead to species misidentification (e.g., Cohen *et al.*, 2004). Non-native species in the family Gracilariaceae often go unnoticed until molecular tools are used to determine taxonomic identity. For example, in coastal bays and estuaries in Virginia, cylindrical specimens were identified as either *Gracilaria verrucosa* (Hudson) Papenfuss (now considered a taxonomic synonym of *Gracilariopsis*

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longissima (S.G. Gmelin) M. Steentoft, L.M. Irvine & W.F. Farnham) or *G. tikvahiae* McLachlan (native to the eastern coast of the United States). However, ecological studies at the Long Term Ecological Monitoring site in Hog Island Bay were in fact assessing the impacts of the invasive *G. vermiculophylla* (Ohmi) Papenfuss (Thomsen *et al.*, 2006), a species widely introduced in the Northern Hemisphere from the northwest Pacific (Kim *et al.*, 2010a; Krueger-Hadfield *et al.*, 2016a, 2016b).

Accurate species identification is critical in order to understand the impacts of non-native species on native biodiversity. Species of economic interest, such as *Undaria pinnatifida* (Harvey) Suringar (Grulois *et al.*, 2011), *Kappaphycus alvarezii* (M.S. Doty) M.S. Doty *ex* P.C. Silva (Sellers *et al.*, 2015) or *Gracilaria salicornia* (C. Agardh) Dawson (Smith *et al.*, 2004), have been intentionally introduced around the world for aquaculture industries and have subsequently "escaped the farm". Similarly, there are algae used in aquaria by professionals and hobbyists, such as the genus *Caulerpa*, that have since found their way into nearby ecosystems (Stam *et al.*, 2006).

Recent studies have reported the presence of two invasive gracilarioid species, Gracilaria vermiculophylla and G. parvispora (C. Agardh) Dawson in Baja California, Mexico (Bellorin et al., 2004; Miller et al., 2012; García-Rodríguez et al., 2013). Kim et al., (2010a) and Krueger-Hadfield et al., (2016b) have documented the spread of G. vermiculophylla throughout the Northern Hemisphere. However, the extent of the native range of G. parvispora and its status as native or non-native has been debated. Abbott (1999) had suggested that G. parvispora was an endemic to Hawai'i. Nelson et al., (2009), by contrast, hypothesized that that G. parvispora Abbott (type locality: Kaneohe Bay, Oahu, Hawai'i) is a native to Asia, perhaps Japan, and was possibly either introduced to Hawaii or was a narrow endemic until it was intentionally introduced around the islands for aquaculture. Nelson et al., (2009) suggested a Japanese origin of G. parvispora in Hawai'i by citing evidence including the earliest Bishop Museum record is from 1908 (though this has not been confirmed to our knowledge with molecular methods), the lack of a Hawaiian name, the availability in Hawaiian fish markets since around 1945 and its disjunct distribution around the Hawaiian Islands. However, unlike Kappaphycus or G. salicornia (see Smith et al., 2004 for a discussion of these species), there is no actual evidence of human introduction of this taxon to Hawai'ian waters. Moreover, Kim et al., (2008) asserted that Gracilaria bursa-pastoris (S. Gmelin) Silva (type locality: Mediterranean Sea) reported from Korea and Japan was a misidentification of G. parvispora, basing their conclusion on rbcL sequences from Hawai'i, South Korea, Japan and Italy in which the sequences from South Korea, Japan and Hawai'i fell in a different clade from that of the Italian specimen. As a result, the distribution of G. parvispora should include central and western Pacific waters.

In northwestern Mexico, despite being large and conspicuous, *Gracilaria vermiculophylla* and *G. parvispora* were either misidentified as congeneric native species (e.g., Bellorin *et al.*, 2004) or were recorded at single sites without molecular confirmation (e.g., Dreckman, 1999). *Gracilaria vermiculophylla* has been present in Baja California since at least 1979 (Bellorin *et al.*, 2004), whereas, *G. parvispora* was not reported until 1999 (Dreckman, 1999). However, first records in taxonomically difficult genera may reflect the first time a specimen was collected, rather than the date at which the introduction occurred. Several introduced gracilarioid seaweeds in North America have been present, but unrecognized, decades longer than published first records (S.A. Krueger-Hadfield and K.A. Miller, *unpublished data*). For both *G. vermiculophylla* and *G. parvispora*, detailed study of archival herbarium collections and contemporary surveys are necessary in order to determine the

distribution of these species outside the northwest Pacific, particularly with regard to estimating the timing of the invasions.

Both Gracilaria vermiculophylla and G. parvispora have or have had some economic importance. The former was an important species in the agar industry in Japan (Okazaki, 1971) and the latter is an important alga in Hawaiian cuisine (Glenn et al., 1996, Abbott, 1999). Though it is uncertain whether aquarium shops sell G. parvispora, or some mixture of different gracilarioid seaweeds, many E-commerce sites purport to sell this species (see for example, http://marineplantbook.com). Because both species are often found together in northwestern Mexico, it is possible that they have been intentionally introduced for use in marine aquaria or for cultivation (García-Rodríguez *et al.*, 2013).

A gracilarioid seaweed, reported both as G. vermiculophylla and Gracilariopsis species, has used in studies of agar quality and biomass in the Laguna San Ignacio in Baja California Sur, Mexico (Vergara-Rodarte et al., 2010, 2016). This species demonstrated high yield (17%) and gel strength (1,132 g cm⁻²) after shorter alkaline treatments, indicating that it is suitable for agar production (Vergara-Rodarte et al., 2010). Vergara-Rodarte et al., (2016) assessed the population structure and standing crop (1,004 wet tons) of what they identified as a *Gracilariopsis* species in the same lagoon. However, the morphology of the spermatangial conceptacle (see Figure 2 from Vergara-Rodarte et al., 2016) indicated that the species in Laguna San Ignacio belongs to the genus *Gracilaria* rather than *Gracilariopsis* (Kim et al., 2008, R. Terada, pers. obs.).

In order to resolve the identity of this potentially valuable gracilariod seaweed, we used the mitochondrial barcode markers cox1 and cox2-3 spacer and the plastid rbcL marker to genotype a representative specimen sampled from El Cordón in Laguna San Ignacio (26°48'46"N 113°12'23"W) in January 2016. We extracted DNA using the DNeasy Blood and Tissue kit (Qiagen, Valencia, CA, USA) following the protocol adapted by Lindstrom et al., (2011). The cox1 and cox2-3 spacer were amplified using the primer pairs 622F (Yang et al., 2008) and 1549R (Geraldino et al., 2006) and cox2-for and cox3-rev (Zuccarello et al., 1999), respectively. The rbcL was amplified using the primer pair F57_rbcL and rbcLrevNEW (Saunders & Moore, 2013). PCR programs were described in Yang et al., (2008) for cox1, in Zuccarello et al., (1999) for cox2-3 and Saunders and Moore, (2013) and the amplification followed protocol described in Krueger-Hadfield et al., (2016b) with commercial sequencing by Eurofins Genomics (Louisville, KY, USA). Sequences were edited using 4PEAKS (Nucleobytes, The Netherlands). The El Cordón specimen is currently stored in silica gel at the University of Alabama at Birmingham (USA).

We aligned all the mitochondrial and plastid sequences using MUSCLE (Edgar 2004) in SEAVIEW ver. 4.6 (Galtier et al., 1996, Gouy et al., 2010) with default parameters. The cox1 (KY056219, this study) and rbcL sequences (KY056221, this study) matched sequences from Mexico, Hawai'i, South Korea and Japan (cox1: KC113591-94, García-Rodríguez et al., 2013; EF434921-22, Yang et al., 2008; rbcL: KC113597-99, García-Rodríguez et al., 2013; EF434942,45, Kim et al., 2008; HQ886637-38, Muangmai, unpubl. data). The cox1 alignment, when trimmed, was 785 bp and the El Cordón specimen was 100% similar to the two sequences in GenBank from Mexico (KC113591-92) and 99% similar to the Hawai'ian and Northwestern Pacific sequences (KC113593-94, EF434921-22). The *rbcL* alignment. when trimmed, was 936 base pairs and the El Cordón sequence was 100% similar to two of the three Mexican specimens (KC113597-98) and the Hawai'ian specimen (EF434945), whereas with the other Mexican sample, there was 1 base pair difference (KC113599). Both Mexican and Hawai'ian sequences were 99% similar to the other Northwest Pacific sequences (EF434942, HQ886637-38).

In contrast, the cox2-3 spacer sequence (KY056220, this study) matched sequences identified as *Gracilaria gigas* Harvey (type locality: Shimoda, Shizuoka Prefecture) from two sites in Japan and reported in Terada & Shimada (2005; AB193451-52). G. gigas is an accepted species reported from the Northwest Pacific and southwestern and southeastern Asia (Guiry & Guiry, 2016). No studies, to our knowledge, have investigated the type specimens of G. parvispora and G. gigas with molecular methods to obtain authentic sequence data or compared contemporary specimens of each species. G. parvispora was first described from a 1970 specimen from Hawai'i (Abbot, 1985), whereas G. gigas was first described in 1860 from a Japanese specimen (Masuda et al., 1995). If these two species are conspecific, the correct name for the species currently known as G. parvispora should be G. gigas by the nomenclatural law of priority. However, as the cox1 and rbcL sequences match GenBank specimens identified as G. parvispora throughout its natural and introduced range (see above) and because there is little current information about G. gigas, it is likely that the El Cordón specimen is the same as others currently identified as G. parvispora. The results of this study suggest that the specimens cited by Terada & Shimada (2005) as G. gigas were misidentifications.

Though we only sampled a single thallus, the species previously studied in Vergara-Rodarte et al., (2010, 2016) is G. parvispora. Zertuche-Gonzalez (1993) mentioned a bed of Gracilaria pacifica located in Laguna de San Ignacio, which was exploited since 1993 for the agar production. The gracilarioid bed was approximately 60 hectares and could produce a total of 900 dry tons per year, from which 20 tons were exported Japan. Further studies found that G. pacifica was the most abundant species and was present along all year at Laguna de San Ignacio (Nuñez-Lopez et al., 1998, Nuñez-Lopez & Casas-Valdez, 1998). Riosmena-Rodríguez et al., (2009) and Vergara-Rodarte et al., (2010) reported the most abundant species in Laguna de San Ignacio was G. vermiculophylla and not G. pacifica based on morphological characteristics. Moreover, the most abundant gracilarioid seaweed, thought to be G. vermiculophylla, formed monospecific bed with a maximum biomass in spring (1,004 wet tons, Vergara-Rodarte et al., 2010, 2016). Though we cannot dismiss the co-occurrence of other taxa, especially G. vermiculophylla (see García-Rodríguez et al., 2013), the dominant species previously studied in Laguna de San Ignacio and genotyped in this study is G. parvispora.

This new record adds another location along the Baja California peninsula where this species is abundant (Fig. 1). In Laguna San Ignacio, tetrasporic plants were the most common, followed by vegetative plants, though it is not known if these are sterile (Vergara-Rodarte et al., 2016). García-Rodríguez et al., (2013) sampled both tetrasporophytes and female gametophytes at the sites in northwest Mexico where they first recorded G. parvispora. Recently, Krueger-Hadfield et al., (2016a) found similar patterns in the congeneric and often associated introduced and invasive species, G. vermiculophylla. In free-floating populations, G. vermiculophylla plants were overwhelmingly tetrasporophytic (Krueger-Hadfield et al., 2016a). However, there were some sites in the introduced range with abundant hard substratum in which both tetrasporophytic and gametophytic plants were sampled (Krueger-Hadfield et al., 2016a). Due to the potential of G. parvispora as a source of agar, more detailed studies of the invasion history and extensive surveys to determine its current distribution and impacts on the recipient ecosystems.

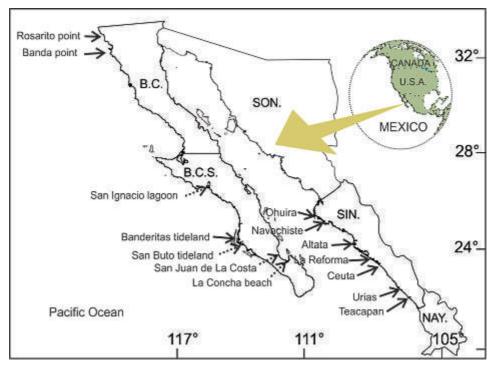


Fig. 1. The known distribution of Gracilaria parvispora (dashed arrow) and Gracilaria vermicullophylla (solid arrows) along the Mexican coast based on previous studies (see review in García-Rodríguez et al., 2013). Laguna de San Ignacio is a new record for G. parvispora.

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