



**Genetic and Morphological Diversity in the Green Algal Genus *Caulerpa*
J.V.Lamouroux (Bryopsidales, Chlorophyta) in Thailand**

Kattika Pattarach

**A Thesis Submitted in Partial Fulfillment of the Requirements for the
Degree of Master of Science in Biology (International Program)**

Prince of Songkla University

2019

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I hereby certify that this work has not been accepted in substance for any degree, and is not being currently submitted in candidature for any degree.

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ชื่อวิทยานิพนธ์	ความหลากหลายทางพันธุกรรมและสัณฐานวิทยาของสาหร่ายสีเขียวสกุล <i>Caulerpa</i> J.V.Lamouroux (Bryopsidales, Chlorophyta) ในประเทศไทย
ผู้เขียน	นางสาวกัตติกา พัฒราษ
สาขาวิชา	ชีววิทยา (นานาชาติ)
ปีการศึกษา	2561

บทคัดย่อ

สาหร่ายสีเขียวสกุล *Caulerpa* มีการกระจายในบริเวณเขตน้ำขึ้นน้ำลงและน้ำท่วมถึงของทะเลเขตร้อนและกึ่งเขตร้อน สาหร่ายบางชนิดมีความแปรผันทางลักษณะสัณฐานวิทยา ซึ่งความสามารถในการเปลี่ยนแปลงรูปร่างทางสัณฐานวิทยา และความซ้อนทับกันของลักษณะทางสัณฐานวิทยาเป็นอุปสรรคต่อการศึกษาทางด้านอนุกรมวิธานของสาหร่ายสกุลนี้ ซึ่งการศึกษาอนุกรมวิธานของสาหร่ายสกุลนี้ในประเทศไทยใช้เพียงข้อมูลทางสัณฐานวิทยาเท่านั้น ดังนั้นข้อมูลทางด้านชีวโมเลกุลจึงมีความจำเป็นในการจัดจำแนกสาหร่ายสกุลนี้ ในการศึกษาครั้งนี้เพื่อศึกษาความหลากหลายของสาหร่ายสกุล *Caulerpa* ในประเทศไทย โดยใช้ข้อมูลทางด้านสัณฐานวิทยา และชีวโมเลกุล (ยีน *tufA* และ ITS rDNA sequences) จากตัวอย่างแห้งจากพิพิธภัณฑ์และตัวอย่างสด จากการศึกษาพบ สาหร่าย *Caulerpa* ทั้งหมด 8 ชนิด ในน่านน้ำไทย คือ *C. chemnitzia* *C. lentillifera* *C. macrodisca* *C. racemosa* *C. serrulata* *C. sertularioides* *C. taxifolia* และ *C. verticillata* และพบสาหร่ายชนิด *C. corynephora* ที่เคยรายงานนั้น เป็นเพียงความแปรผันทางลักษณะสัณฐานวิทยาของสาหร่ายชนิด *C. macrodisca* ดังนั้น *C. macrodisca* จึงมีสามลักษณะทางสัณฐานวิทยา คือ *C. macrodisca* ecad *ashmeadii* *C. macrodisca* ecad *corynephora* และ *C. macrodisca* ecad *macrodisca* และสาหร่าย *C. racemosa* จะมีสองลักษณะทางสัณฐานวิทยา คือ *C. racemosa* ecad *chemnitzia* และ *C. racemosa* ecad *racemosa* ดังนั้นการศึกษาในครั้งนี้สามารถสรุปได้ว่าลักษณะทางสัณฐานวิทยาเพียงอย่างเดียวไม่เพียงพอต่อการจัดจำแนกชนิดของสาหร่ายสกุล *Caulerpa* ได้ โดยยีน *tufA* สามารถใช้ในการระบุชนิดของสาหร่ายสกุลนี้ได้ดี และ ITS rDNA sequence สามารถใช้ประกอบการยืนยันการจัดจำแนกได้ ดังนั้น ควรใช้ทั้งข้อมูลทางสัณฐานวิทยาร่วมกับข้อมูลทางพันธุกรรมในการศึกษาความ

หลากหลายของสาหร่ายสกุล *Caulerpa* นอกจากนี้ในงานวิจัยครั้งนี้ไม่พบสาหร่ายสกุล *Caulerpa* บางชนิดที่เคยรายงานไว้ ในการสำรวจภาคสนาม และไม่สามารถจัดจำแนกชนิดได้เนื่องจากไม่สามารถสกัดสารพันธุกรรมจากตัวอย่างแห้งได้

Thesis Title	Genetic and morphological diversity in the green algal genus <i>Caulerpa</i> J.V.Lamouroux (Bryopsidales, Chlorophyta) in Thailand
Author	Kattika Pattarach
Major Program	Biology (International Program)
Academic Year	2018

ABSTRACT

Caulerpa is a coenocytic green algal genus that is widely distributed in the intertidal and subtidal zones of tropical and subtropical seas. Some species show a complex external morphology. A high morphological plasticity and overlapping morphologies hamper the taxonomy of the genus tremendously. All previous taxonomic studies of *Caulerpa* in Thailand were based on morphological observations. Since then the taxonomic status of several *Caulerpa* species has been challenged by molecular studies. The present study aims to reassess the *Caulerpa* diversity in Thailand using morphological and molecular data (*tufA* gene and ITS rDNA sequences) from herbarium and fresh specimens. The result confirmed the occurrence of eight *Caulerpa* species in Thai waters, *i.e.*, *C. chemnitzia*, *C. lentillifera*, *C. macrodisca*, *C. racemosa*, *C. serrulata*, *C. sertularioides*, *C. taxifolia* and *C. verticillata*. Specimens previously identified as *C. corynephora* in Thailand actually are a morphological variety of *C. macrodisca*. Then, there were three morphological groups of *C. macrodisca*; *C. macrodisca* ecad *ashmeadii*, *C. macrodisca* ecad *corynephora* and *C. macrodisca* ecad *macrodisca*. In addition, *C. racemosa* is represented by two distinct morphologies; *C. racemosa* ecad *chemnitzia* and *C. racemosa* ecad *racemosa*. It is concluded that *Caulerpa* species identification cannot rely on morphology alone. The *tufA* gene was useful for specific identification of *Caulerpa*, and ITS rDNA sequence was supported data. So, this study suggested that morphological and molecular data were good for *Caulerpa* diversity. In addition, several morphological *Caulerpa* species previously recorded for Thailand were not found during our field survey and their identity could not be confirmed with DNA sequence data obtained from herbarium specimens.

ACKNOWLEDGEMENTS

I would like to express my gratitude to the contributed persons in this work.

I sincerely appreciate to Prof. Khanjanapaj Lewmanomont (Kasetsart University, Thailand) for her invaluable help of KUMF herbarium dried specimens.

I appreciate to Assoc. Prof. Dr Shao-Lun Liu (Tunghai University, Taiwan) for molecular research methodologies.

I deeply appreciate to Dr Jaruwat Mayakun (my advisor) and Dr Stefano G.A. Draisma (my co-advisors) for their opportunity to do this study. I most appreciated any supporting and comments, which advice throughout this work.

I also thank to Mr. Watana Chaithongkaew, Mr. Jatdilok Titioatchasai, Mr. Nutdanai Putthisawongand, Miss Surangkana Phandee and the Seaweed and Seagrass Research Unit (SSRU) team for their help in any field collection.

Finally, thanks to the Science Achievement Scholarship of Thailand (SAST) for my financial support, Faculty of Science, Prince of Songkla University (PSU) (Contact No. SCI610575S) and Graduate School, PSU for our grant.

Kattika Pattarach

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INTRODUCTION

Background and rationale

Caulerpa is one of the most common and widely distributed green algal genera in subtidal tropical and subtropical marine waters (de Senerpont Domis *et al.*, 2003; Belleza and Liao, 2007; Lee, 2008). The thallus of *Caulerpa* consists of a green creeping rhizome (stolon) with colorless downward growing rhizoids and upward growing photosynthetic fronds (assimilators). These assimilators usually bear many determinate branchlets termed ramuli. These ramuli come in a variety of shapes such as flattened, cylindrical, globular, or disk-shaped. The shape of the assimilators is the major key for species identification in this genus.

At present, there are 382 species (and infraspecific) names in the database, of which 97 species and 110 varieties and forms have been indicated as currently taxonomically accepted name (Guiry and Guiry, 2019). *Caulerpa* species show infraspecific morphological variation hampering delimiting species boundaries. This morphological plasticity may be induced by environmental factors, such as light intensity (Calvert, 1976) and temperature that affect the morphology of assimilators on *Caulerpa chemnitzia* (Esper) J.V.Lamouroux (as *Caulerpa racemosa* var. *peltata* (J.V.Lamouroux) Eubank) (Ohba *et al.*, 1992; Komatsu *et al.*, 1997). Other environmental factors; salinity, wave action and seasonal variations also influence the morphology of *Caulerpa* spp. (Calvert, 1976; Carruthers *et al.*, 1993; Meinesz *et al.*, 1995; Robledo and Freile-Pelegrín, 2005). Morphological plasticity has resulted in phenotypic variation within species of this genus and has been a long-standing source of uncertain taxonomy and causes the taxonomic havoc in *Caulerpa* (Famà *et al.*, 2002; Belleza and Liao, 2007; Sauvage *et al.*, 2013; Belton *et al.*, 2014) and the morphological characters only are not enough for the identification and classification of several species in this genus.

Recently, researchers applied molecular tools to identify some morphologically plastic species (Verbruggen *et al.*, 2009; Mahakham, 2011; de Clerck *et al.*, 2013). Molecular studies used the chloroplast-encoded *tufA* gene and the Internal Transcribed Spacers (ITS) of the nuclear ribosomal cistron to infer phylogenetic relationships and a subgeneric classification in the genus *Caulerpa* in various regions (Olsen *et al.*, 1998; Famà *et al.*, 2002; Stam *et al.*, 2006; Kazi *et al.*, 2013; Sauvage *et al.*, 2013; Belton *et al.*, 2014; Draisma *et al.*, 2014). *Caulerpa* species from the Mediterranean region, California, India and Australia have been exhaustively investigated and determined (Verlaque, 2000; Zaleski and Murray, 2006; Kazi *et al.*, 2013; Belton *et al.*, 2014).

Phang *et al.* (2016) listed 27 *Caulerpa* species and 60 taxa including varieties and formae in the South China Sea, including eight countries; Brunei, Cambodia, Indonesia, Malaysia, Philippines, Singapore, Vietnam and Thailand. In Thailand, only Lewmanomont (2008) published on the taxonomy of *Caulerpa* using morphological data and reported 16 *Caulerpa* species in Thai waters (the Andaman Sea and the Gulf of Thailand coast). Morphological variation was recorded in many herbarium specimens of *Caulerpa racemosa* (Forsskål) J.Agardh collected from various areas, such as Phuket, Surat Thani, Trang and Satun provinces, but intraspecific names were not reported (Personal Observation). However, there is no data on genetic diversity and distribution patterns of *Caulerpa* in Thailand. The present study aims to investigate the genetic diversity and distribution patterns of *Caulerpa* in Thailand using the *tufA* gene and ITS rDNA sequence data and morphological characters. This study can answer these questions; 1) how many species of *Caulerpa* are there along the coasts of Thailand? and 2) what is the extent of infraspecific morphological and genetic variation in *Caulerpa* species from Thailand? Then, the results were presented into 2 parts; 1. Diversity and distribution of the genus *Caulerpa* in Thailand and 2. An enigmatic *Caulerpa macrodisca* Decaisne (Chlorophyta) from the mangrove channels on the Andaman Sea coast of Thailand.

Review of literature

Lamouroux (1809) erected the genus *Caulerpa* and reported eight species representing five currently accepted species; *Caulerpa prolifera* (Forsskål) J.V.Lamouroux (including *Caulerpa ocellata* J.V. Lamouroux), the type of the genus that was designated in Trevisan (1849), *C. chemnitzia* (including *Caulerpa peltata* J.V.Lamouroux), *Caulerpa flexilis* J.V.Lamouroux ex C.Agardh (as *Caulerpa hypnoides* C.Agardh), *C. racemosa* (as *Caulerpa obtusa* J.V.Lamouroux, uncertain synonymy), *Caulerpa sertularioides* (S.G. Gmelin) M. Howe (as *Caulerpa myriophylla*) and *Caulerpa taxifolia* (M.Vahl) C.Agardh (as *Caulerpa pennata* J.V.Lamouroux). Kützing (1843) classified this genus into the family Caulerpaceae. *Caulerpa* consists of a creeping green stolon with colorless rhizoids and frond-like erect shoots. The upright shoots present considerable variation in morphology of the ramuli, such as sickle-shaped, disc-shaped, mushroom-shaped and globular. (Lee, 1999; Draisma *et al.*, 2014). A characteristic feature of this family is the presence of trabeculae, which are internal β -1,3 linked xylan cell wall outgrowths that give structural support the thallus (Lamouroux, 1809; Bold and Wynne, 1985; Famà *et al.*, 2002; Lee, 2008; Draisma *et al.*, 2014).

In 1899–1900, the first record of *Caulerpa* was from the vicinity of Ko Chang, the Gulf of Thailand during the Danish expedition to Siam (Rathbun, 1919). Reinbold (1901) found 11 species; *C. chemnitzia* (as *C. peltata*), *Caulerpa fastigiata* f. *minor* Weber-van Bosse, *Caulerpa lentillifera* J.Agardh (as *Caulerpa lentillifera* var. *longistipitata* Weber-van Bosse), *Caulerpa plumaris* var. *longipes* Weber-van Bosse, *C. racemosa* var. *uvifera* Weber-van Bosse, *Caulerpa scalpelliformis* (R.Brown ex Turner) C.Agardh, *Caulerpa sedoides* C.Agardh, *Caulerpa serrulata* var. *pectinata* (Weber-van Bosse) W.R.Taylor (as *Caulerpa freycinetii* var. *pectinata* Weber-van Bosse), *Caulerpa tongaensis* Papenfuss (as *Caulerpa filiformis* J.Agardh), *Caulerpa urvilleana* f. *tristicha* (J.Agardh) Weber-van Bosse and *Caulerpa verticillata* J.Agardh that grow on rocks in shallow water at Ko Chick, Ko Samae San and Ko Kahdat. In addition, Dawson (1954) found *C. lentillifera* in the northern part of Saen Soek, the Gulf of Thailand. Egerod (1974) published a second report including *Caulerpa mexicana* Sonder ex Kützing, *Caulerpa microphysa* (Weber-van Bosse) Feldmann, *Caulerpa racemosa* var. *macrophysa* (Sonder ex Kützing)

W.R.Taylor and *C. chemnitzia* (as *Caulerpa racemosa* var. *peltata* (J.V.Lamouroux) Eubank) from the Andaman Sea. The Fifth Thai Danish Expedition (1966), the third report (Seidenfaden *et al.*, 1968) showed five species of *Caulerpa*; *C. ambigua*, *Caulerpa fastigiata* Montagne, *C. mexicana*, *C. racemosa* and *C. sertularioides* at Phuket (Egerod, 1971, 1974, 1975). Lewmanomont (1978) studied some algae that are used as food in Thailand. She reported that *C. racemosa* var. *macrophysa* and *Caulerpa corynephora* Montagne (as *Caulerpa racemosa* var. *corynephora* (Montagne) Weber-van Bosse) are commonly used as food for people in the South. The Algae Checklist of Thailand (Lewmanomont *et al.*, 1995) lists 19 *Caulerpa* species, seven varieties and two formae. The most recent taxonomic morphology based treatment of the genus *Caulerpa* in Thailand lists 16 species including *Caulerpa ashmeadii* Harvey, *Caulerpa cupressoides* (Vahl) C.Agardh, *Caulerpa fastigiata* Montagne, *C. lentillifera*, *C. racemosa* var. *macrophysa* (as *Caulerpa macrophysa* (Sonder ex Kützing) G.Murray), *Caulerpa manorensis* Nizamuddin, *Caulerpa mexicana* Sonder ex Kützing, *C. microphysa*, *C. chemnitzia* (as *C. peltata*), *Caulerpa macrodisca* (Forsskål) J. Agardh (as *C. peltata* var. *macrodisca*), *C. corynephora* (as *C. racemosa* var. *corynephora*), *C. serrulata*, *C. sertularioides*, *C. taxifolia*, *C. verticillata* and *C. ambigua* currently recognized species (Lewmanomont, 2008).

A revision of the *Caulerpa* herbarium specimens in the Princess Maha Chakri Sirindhorn Natural History Museum (NHM) in Hat Yai revealed phenotypic variations of *Caulerpa* specimens (Pattarach, 2016; Personal Observation) and identification at species and subspecies level remains problematic. Seaweed expert Eric Coppejans also found *Caulerpa* with different shapes and arrangements of ramuli from the same study site in Krabi province, Thailand (personal communication). The *Caulerpa racemosa-peltata* complex along the Thai coast showed a variety of morphologies.

The branching of photosynthetic fronds exhibits a diverse range of ramuli morphology. Light intensity and temperature affect the morphology of assimilators on *C. chemnitzia* (as *C. racemosa* var. *peltata*) (Ohba *et al.*, 1992). Other environmental factors; salinity, wave action and seasonal variations also influence the morphology of *Caulerpa* spp. (Calvert, 1976; Carruthers *et al.*, 1993; Meinesz *et al.*, 1995; Robledo and Freile-Pelegrín, 2005). Morphological plasticity has resulted in phenotypic

variation within species of this genus and has been a long-standing source of uncertain taxonomy (Famà *et al.*, 2002; Belleza and Liao, 2007; Sauvage *et al.*, 2013; Belton *et al.*, 2014).

DNA barcoding is recently developed molecular tool to assist the identification of *Caulerpa* species. This method analyzes a nucleotide sequence of a standard DNA region of an unknown specimen and compares it with the nucleotide sequences of identified specimens. This can give an accurate and reliable identification in a short time (Mahakham, 2011). This technique was used to clearly identify cryptic species in the bryopsidalean genus *Codium* that showed morphological plasticity (Lee and Kim, 2015).

The *tufA* gene is a conserved sequence in green algae, located in the chloroplast genome. It also is a common DNA markers used in phylogenetic studies because it can present high variation at the species level (Famà *et al.*, 2002; Pečnikar and Buzan, 2014). Currently 1,115 *Caulerpa tufA* sequences in the GenBank nucleotide database representing 70 *Caulerpa* species (78 % of the 90 currently recognized species) (see Draisma *et al.*, 2014, suppl. table S2). In addition, the Internal Transcribed Spacers (ITS1 and ITS2) of the nuclear ribosomal cistron have also been used frequently in *Caulerpa* taxonomy (Kazi *et al.*, 2013; Yeh and Chen, 2014; Stam *et al.*, 2006). There are many reasons to use ITS rDNA as a barcoding marker. First, the nucleotide sequence of this gene is highly variable at the inter- and infraspecific level in green algae (Bakker *et al.*, 1995; Olsen *et al.* 1994, 1998). Second, the variation in ITS nucleotide sequences is about 3–4 times higher than that in chloroplast genes (Mahakham, 2011). Third, there are many ITS rDNA sequences available in GenBank, than can be used for the development of PCR primers.

Many molecular studies of *Caulerpa* have been done in many countries. For example, Olsen *et al.* (1998) used ITS rDNA sequences for the identification of *C. taxifolia* and *C. mexicana* in the Mediterranean. These two taxa have similar morphologies, but maximum parsimony and maximum likelihood phylogenies clearly separate the two taxa. The two species are thus not conspecific.

Famà *et al.* (2002) presented a molecular phylogeny of 23 *Caulerpa* taxa from tropical and subtropical zones of the world using the *tufA* gene. Their phylogenetic tree showed that the traditional morphology based infrageneric classification into 12 sections does not represent evolutionary history as most sections are polyphyletic. Sixteen clades could be discerned in their maximum likelihood phylogeny. *C. mexicana*, *C. sertularioides*, *C. webbiana*, *C. prolifera*, *C. taxifolia* and *C. racemosa* were sampled multiple times and each formed a monophyletic clade, except the latter two species. *C. distichophylla* was nested inside *C. taxifolia* and *C. racemosa* was polyphyletic divided into four clades.

Stam *et al.* (2006) did a phylogenetic survey of *Caulerpa* in the Caribbean using ITS rDNA and *tufA*. Their phylogeny showed that *C. brachypus* was sister to *C. scalpelliformis*. In addition, *C. cupressoides* and *C. serrulata* could not unambiguously be distinguished from each other and *C. racemosa* and *C. scalpelliformis* were both polyphyletic. Moreover, *C. taxifolia* and *C. racemosa* were considered as alien species record for the first time in Florida, USA.

Sauvage *et al.* (2013) studied the molecular diversity of the *C. racemosa-peltata* complex in New Caledonia and Australia using the *tufA* gene. The result showed that five lineages were identified in New Caledonia, and the main diversity may be restricted to the Indo-Pacific region. In addition, *Caulerpa cylindracea* Sonder (as *C. racemosa* var. *cylindracea*) was found as new record at Lizard Island on the Great Barrier Reef, Australia.

Kazi *et al.* (2013) studied molecular phylogeny and DNA barcoding of *Caulerpa* in India using the *tufA*, *rbcL*, 18S rDNA and ITS rDNA. The result showed that the classification of *Caulerpa* using multiple markers is more efficient than using just one gene. The molecular phylogeny result was consistent with the morphological study.

Belton *et al.* (2014) studied the character of phenotypic plasticity and identification of the *C. racemosa-peltata* complex in Australia using *tufA* and *rbcL* gene sequences. The result revealed 11 *Caulerpa* lineages within the complex, and five

of them, such as *Caulerpa lamourouxii* (Turner) C.Agardh and *Caulerpa oligophylla* Montagne showed a high degree of phenotypic plasticity and some overlap with others.

Draisma *et al.* (2014) revised the subgeneric classification of *Caulerpa* based on a time-calibrated molecular phylogeny. The phylogenetic tree based on *rbcL* and *tufA* gene sequences revealed six main clades that were each given subgenus status. Three of the six subgenera are currently monotypic, albeit *Caulerpella ambigua* (Okamura) Prud'homme van Reine & Lokhorst showed high diversity. It was concluded that *Caulerpella ambigua* represents multiple cryptic species nested inside the genus *Caulerpa*. So, *Caulerpella ambigua* was transferred back to the genus *Caulerpa* called that *Caulerpa ambigua* Okumura.

In conclusion, the phylogenetic analysis using some specific nucleotide sequences is able to identify morphological plastic species in the genus *Caulerpa*. This has been demonstrated in several regions, e.g. the Caribbean, the Mediterranean, India, New Caledonia and Australia. However, a comprehensive molecular study of *Caulerpa* in Thailand is still lacking. The present research aims to study the genetic diversity and distribution patterns of *Caulerpa* in Thai waters.

Questions of this research

1. How many species of *Caulerpa* are there along the coasts of Thailand?
2. What is the extent of infraspecific morphological and genetic variation in *Caulerpa* species from Thailand?

Objectives

1. To identify the species of *Caulerpa* using morphological and molecular evidence.
2. To assess the genetic diversity of *Caulerpa* using *tufA* and ITS rDNA sequences.
3. To confirm the identification of *Caulerpa*.

For answering the question, I begin with 1. Diversity and distribution of the genus *Caulerpa* in Thailand, followed by 2. An enigmatic *Caulerpa macrodisca* Decaisne (Chlorophyta) from the mangrove channels on the Andaman Sea coast of Thailand.

RESEARCH METHODOLOGY

Collection sites

The study sites were chosen along the Thai coast in 15 provinces following Lewmanomont (2008) (Fig. 1)

The Gulf of Thailand

- Losin, Pattani
- Kao Seng and Ko Kham,
Songkhla
- Sichon, Nakhon Si Thammarat
- Ko Samui and Ko Tan,
Surat Thani
- Mu Ko Chumphon, Chumphon
- Ao Manao,
Prachuap Khiri Khan
- Ko Kram and Ko Samae San,
Chonburi
- Ko Chik, Chanthaburi
- Saoson Beach, Rayong
- Ko Chang, Ko Kradat,
Ao Cho and Leam Sok, Trat

The Andaman Sea

- Tang Khen Bay, Phuket
- Ko Phra Thong and
Ko Similan, Phang Nga
- Ko Siboya, Khlong Yang,
Ko Lanta, Krabi
- Ko Libong, Leam Yong Lam
and Khao Bae Na, Trang
- Ko Lidee, Ko Lipe, Thung Wa
and Che Bilang, Satun

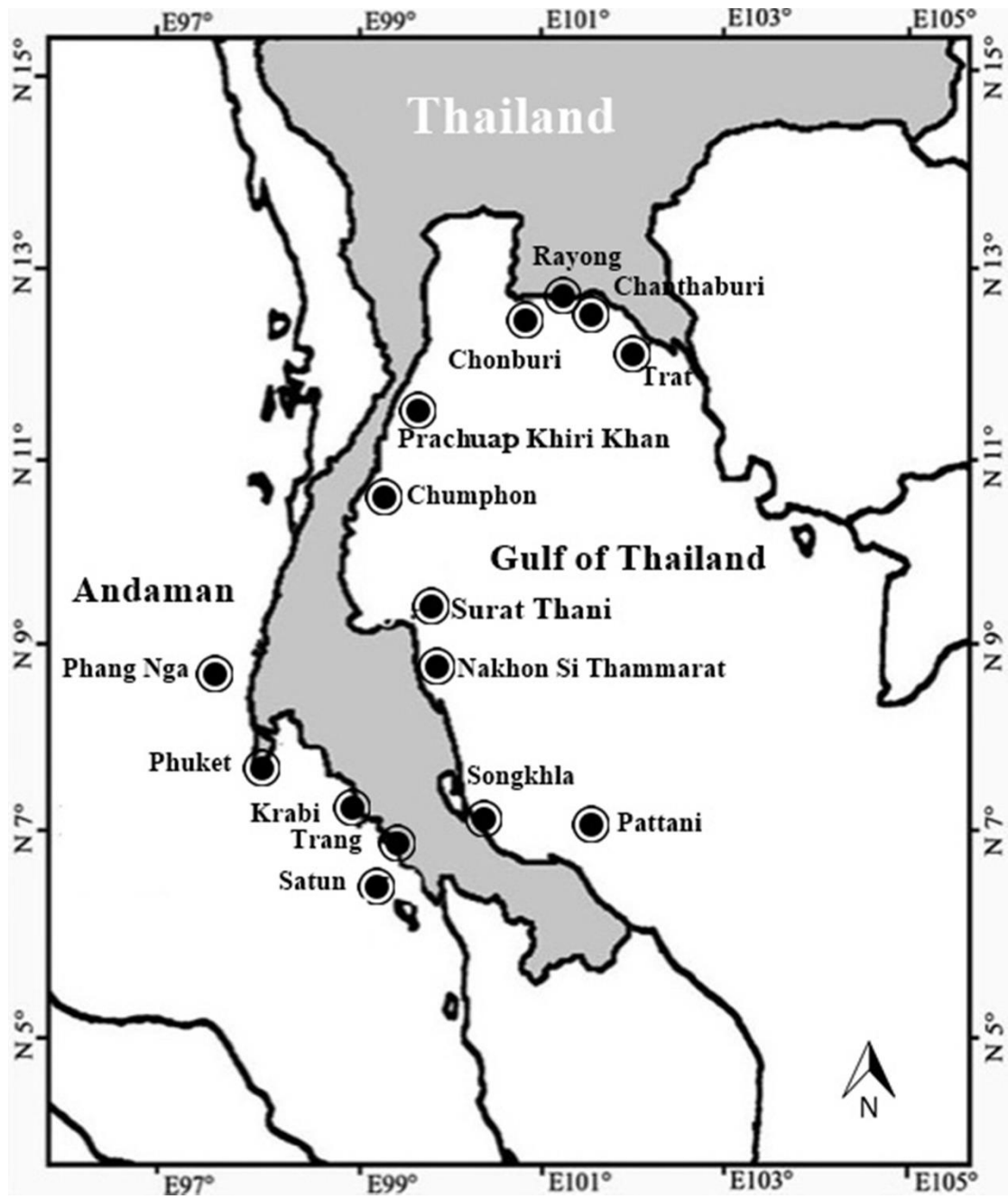


Figure 1 Map of study sites at 15 provinces in Thailand.

Environmental factors

Four environmental factors were recorded in this study, *i.e.*, habitat type, water temperature, light intensity, and salinity (Table 1). Water temperature was estimated in some sites using either a laboratory thermometer or the Onset-Hobo data logger (Onset Computer Corporation, Contoocook, NH, USA). Light intensity was also measured using the Onset-Hobo data logger. More than 800 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ was defined as high light intensity and less than 800 $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ as low intensity. Salinity was determined in some sites using a salinity reflectometer (ATC, 0–100 ppt, XHO RHS-10ATC, ATACO, China).

Table 1 Summary of environmental factors used in this study. n.m., not measured.

Study sites	GPS coordinates	Environmental factors			
		Habitat types	Water temperature (°C, mean±SE)	Light intensity	Salinity (‰)
Pattani					
Losin	7°19'N, 101°51'E	Dead coral in coral reef of subtidal zone	n.m.	n.m.	35
Songkhla					
Kao Seng	7°10'N, 100°37'E	Rocky shore in intertidal zone	34±0.12	High light	33
Ko Kham	6°58'N, 100°51'E	Rocky shore in intertidal zone	33±0.47	High light	33
Nakhon Si Thammarat					
Sichon	-	Sandy bottom and rock in intertidal zone	n.m.	n.m.	n.m.
Surat Thani					
Ko Samui	9°34'N, 100°03'E	Dead coral, sandy bottom and rock in intertidal zone	32±0.32	High light	33
Ko Tan	9°23'N, 99°57'E	Dead coral, sandy bottom and rock in intertidal and subtidal zone	32±0.45	High light	33

Study sites	GPS coordinates	Environmental factors			
		Habitat types	Water temperature (°C, mean±SE)	Light intensity	Salinity (‰)
Chumphon					
Mu Ko Chumphon	10°19'N, 99°18'E	Dead coral, sandy bottom and rock	30	Low light	33
	10°19'N, 99°16'E	in intertidal and subtidal zone		(monsoon)	
	10°17'N, 99°14'E				
Prachuap Khiri Khan					
Ao Manao	11°45'N, 99°47'E	Sandy bottom and rock in intertidal zone	34	High light	33
Chonburi					
Ko Kram	-	Dead coral, sandy bottom and rock in intertidal and subtidal zone	n.m.	n.m.	n.m.
Ko Samae San	-	Dead coral, sandy bottom and rock in intertidal and subtidal zone	n.m.	n.m.	n.m.
Chanthaburi					
Ko Chik	12°17'N, 102°14'E	Dead coral and rock in intertidal and subtidal zone	35	High light	33

Study sites	GPS coordinates	Environmental factors			
		Habitat types	Water temperature (°C, mean±SE)	Light intensity	Salinity (‰)
Rayong					
Saoson Beach	12°37'N, 101°28'E	Sandy bottom in intertidal zone	34	High light	33
Trat					
Ko Chang	12°04'N, 102°16'E	Dead coral and rock in intertidal zone	35	High light	34
Ko Kradat	-	Dead coral and rock in intertidal zone	n.m.	n.m.	n.m.
Ao Cho	12°33'N, 101°27'E	Sandy bottom and rock in intertidal zone	n.m.	n.m.	33
Leam Sok	12°02'N, 102°35'E	Rocky shore in intertidal zone	35	High light	33
Phuket					
Tang Khen Bay	-	Dead coral in intertidal zone	n.m.	n.m.	33

Study sites	GPS coordinates	Environmental factors			
		Habitat types	Water temperature (°C, mean±SE)	Light intensity	Salinity (‰)
Phang Nga					
Ko Phra Thong	9°07'N, 98°15'E	Dead coral and rock in intertidal zone	n.m.	n.m.	33
Ko Similan	8°38'N, 97°39'E	Dead coral and rock in intertidal zone	n.m.	n.m.	n.m.
Krabi					
Ko Siboya	7°53'N, 98°58'E	Sandy bottom and rock in intertidal zone	n.m.	n.m.	33
Khlong Yang	7°48'N, 99°07'E	Muddy bottom and fish cages of mangrove channels	27±0.45	Low light	~28 (27–30)
Ko Lanta	7°28'N, 99°05'E	Tide pool of rocky shore in intertidal zone	29±1.25	High light	33
Trang					
Ko Libong	7°15'N, 99°26'E	Sandy bottom of subtidal zone	30±0.19	High light	33
Leam Yong Lam	7°23'N, 99°20'E	Sandy bottom of subtidal zone	32±0.47	High light	33
Khao Bae Na	7°24'N, 99°20'E	Sandy bottom of subtidal zone	32±0.37	High light	33

Study sites	GPS coordinates	Environmental factors			
		Habitat types	Water temperature (°C, mean±SE)	Light intensity	Salinity (‰)
Satun					
Ko Lidee	6°47'N, 99°45'E	Sandy bottom and rock in intertidal zone	33±0.04	High light	~33 (31–34)
Ko Lipe	6°29'N, 99°18'E	Sandy bottom and rock in intertidal zone	35	High light	35
Thung Wa	6°37'N 100°01'E	Muddy bottom and fish cages of mangrove channels	n.m.	Low light	~30 (27–30)
Che Bilang	-	Muddy bottom and fish cages of mangrove channels	n.m.	Low light	~30 (27–30)

Specimen collection

At least three specimens (complete thalli; rhizoid, stolon and frond of each morphological species from each study site) without epiphyte were collected using scuba diving and snorkeling from the intertidal to around 5–10 m depth. The samples were preserved on herbarium sheets for morphological study. Specimens were pressed and labels with collection date, locality, collector name, and habitat type were added to the herbarium following Coppejans *et al.* (2010). All specimens were photographed using the camera Canon 60D with Canon EF-S 18–55mm f/3.5–5.6 IS. The fragments (1–2 cm) of each sample were quickly dried and preserved in silica gel for molecular studies. In addition, herbarium-dried specimens from the Princess Maha Chakri Sirindhorn Natural History Museum (NHM) in Prince of Songkla University (PSU) and Kasetsart University Museum of Fisheries (KUMF) in Kasetsart University (KU) were also sampled for nucleotide analysis.

The morphological study

Twenty-seven morphological characters (Table 2, Fig. 2) such as assimilator length, stolon and ramuli diameter (adapted from de Senerpont Domis *et al.*, 2003) were measured on each fresh specimen using a caliper, before it is processed into a herbarium specimen. Once herbarium dried specimens were measured using NIH ImageJ software (Rasband, 1997). Each specimen was investigated using field guides of seaweed (Trono, 1997; de Clerck *et al.*, 2005; Phang *et al.*, 2008; Coppejans *et al.*, 2010; 2017).

Table 2 Summary of characters to be used in morphometric analysis (adapted from de Senerpont Domis *et al.*, 2003)

No.	Morphological structures	Unit	Description
1	Number of stolon branches per cm stolon	no./cm	taken by counting the number of stolon branching and dividing it by the stolon length
2	Diameter of Stolon	cm	measured as thickest part of the stolon near the growing tip
3	Number of rhizoid clusters per cm stolon	no./cm	taken by counting the number of rhizoid clusters and dividing it by the stolon length
4	Spacing of rhizoid clusters	cm	measured as distance between rhizoid clusters
5	Length of rhizoid cluster	cm	measured as the base to the apical end of rhizoid of the thallus
6	Number of assimilators per cm stolon	no./cm	taken by counting the number of assimilator and dividing it by the stolon length
7	Spacing of assimilators	cm	measured as distance between assimilators
8	Length of assimilator	cm	measured from the base to the apical end of the upright of the thallus
9	Width of assimilator	cm	measured from the right side to the left side at the widest assimilator of the thallus
10	Assimilator shape	none	flat, feather-like (distichous), irregular and thread-like (filiform)
11	Assimilator margin	none	entire, serrate and spiny
12	Rachis length	cm	measured from the base to the first ramulus
13	Diameter of rachis	cm	measured as the thickest part of the main axis of the assimilator near the apex
14	Diameter of a constricted of rachis	cm	measured a narrow part of rachis
15	Number of laterals per cm assimilator	no./cm	taken by counting the number of assimilator branching and dividing it by the assimilator length
16	Spacing of ramuli	cm	measured as distance between ramuli on the assimilator

No.	Morphological structures	Unit	Description
17	Arrangement of ramuli	none	opposite, alternate, whorled and irregularly
18	Length of ramulus	cm	measured from the base to the apical end of ramulus
19	Diameter of ramulus	cm	measured as the broadest part of ramulus
20	Shape of ramulus	none	turbinate shape, subspherical shape, clavate shape, serrate shape, falcate shape, cylindrical shape, peltate shape, mushroom-like and filiform shape
21	Shape of ramulus tip	none	acute, acuminate, rounded, oval, mucronate, teeth and three spines
22	Shape of ramulus base	none	cuneate, obtuse, cordate and oblique
23	Diameter of a constricted of ramulus	cm	measured a narrow part of ramulus
24	Length of stalk	cm	measured from the base of ramuli to the axes of assimilator
25	Diameter of stalk	cm	measured as the broadest part of the stalk
26	Diameter of a constricted of stalk	cm	measured a narrow part of stalk
27	Shape of tip	none	acute, acuminate, rounded, mucronate, teeth and disc

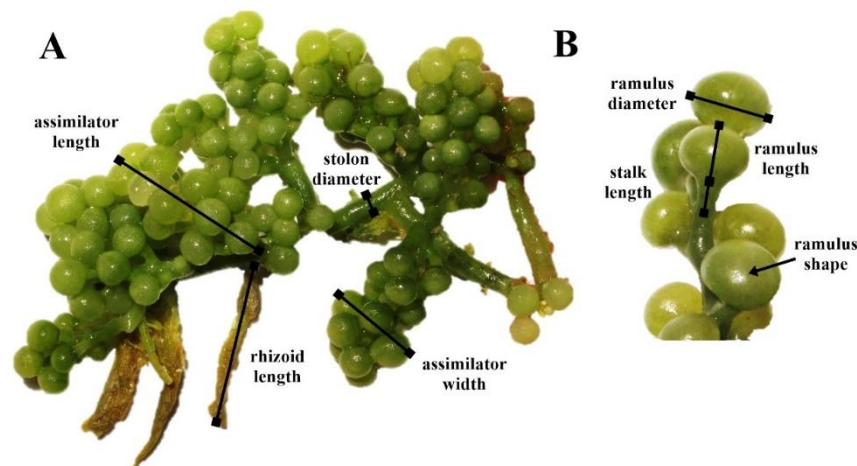


Figure 2 Morphological characters were used for morphometric measurement (A., thallus and B., ramuli of *C. racemosa* (KP40A) from Ko Lipe, Satun)

Molecular study

DNA extraction, amplification and sequencing

DNA was extracted from Silica gel dried specimens using ZR plant/Seed DNA MiniPrep™ Kit (Zymo Research Corporation, New York, USA). A small sample was gridded and added 750 µl Lysis Solution into the ZR BashingBead™ Lysis tube. The tube was processed with vortex meter at maximum speed for 40 minutes and centrifuged at 10,000 x g for 3 minutes. The supernatant (400 µl) was transferred to a Zymo-Spin™ IV Spin Filter tube and centrifuge at 7,000 x g for 3 minutes. The solution was mixed with Plant/Seed DNA Binding Buffer (1,200 µl). The mixture was filtrated by Zymo-Spin™ IIC Column tube and centrifuged at 10,000 x g for 3 minutes. DNA Pre-Wash Buffer (200 µl) was added to the Zymo-Spin™ IIC Column tube and centrifuged at 10,000 x g for 3 minutes. Plant/Seed DNA Wash Buffer (500 µl) was added to the Zymo-Spin™ IIC Column tube and centrifuged at 10,000 x g for 3 minutes (2 time). DNA Elution Buffer (80 µl) was added into the column matrix and centrifuged at 10,000 x g for 3 minutes. The eluted DNA was filtrated by Zymo-Spin™ IV-HRC Spin Filter tube and centrifuge at exactly 8,000 x g for 3 minutes. The DNA product was finally extracted and stored at -20 °C.

PCR amplification

Primers for *tufA* gene (Famà *et al.*, 2002)

Forward primer: *tufA* (5'-TGAAACAGAAMAWCGTCATTATGC-3')

Reverse primer: *tufAR* (5'-CCTTCNCGAATMGCRAAWCGC-3')

Primers for ITS sequence (Kooistra *et al.*, 2002 and White *et al.*, 1990)

Forward primer: H1F (5'-CTCTGAACCTTCGCACGTAGA-3')

Reverse primer: ITS4R (5'-TCCTCCGCTTATTGATAGATGC-3')

PCR amplification of *tufA* gene was performed in 20 μL of master mix contained 0.4 μL of each 0.2 μM primer, 0.4 μL of 0.2 mM dNTPs, 0.12 μL of 0.1X Taq DNA polymerase and 2 μL of 10X buffer (Clontech Laboratories Inc., a Takara Bio company, CA, USA) and 1 μL DNA template. Double-stranded DNA amplifications were performed in a S1000™ thermal cycler (Bio-Rad Laboratories, California, USA). An initial denaturation at 96°C for 4 minutes followed by 40 cycles of 94°C for 30 second, 52°C for 30 s and 72°C for 1 min, a final extension at 72°C for 6 min. For an annealing of ITS rDNA sequence, 48°C for 30 s (Fig. 3).

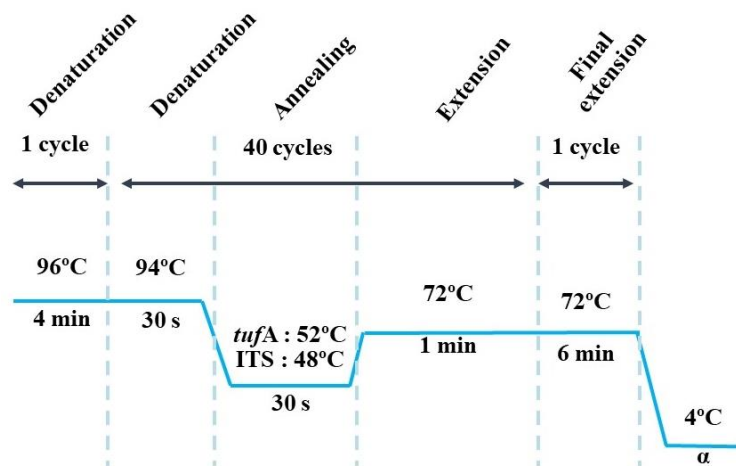


Figure 3 This is PCR amplification condition of this study following Taq DNA polymerase protocol (Clontech Laboratories Inc., a Takara Bio company, CA, USA).

Nucleotide sequencing

PCR products were purified and sequenced by Genedragon Inc. (Taiwan) and Macrogen Inc. (Seoul, Korea).

Nucleotide sequence alignment analysis

A total of 270 *Caulerpa tufA* sequences from GenBank together with 45 newly *tufA* *Caulerpa* sequences (Table 3) were used with *Dichotomosiphon tuberosus* A.Braun ex Kützing) A.Ernst (GenBank MH591082) as outgroup. For ITS sequences, 59 GenBank sequences with 56 newly generated ITS *Caulerpa* sequences (Table 3) were analyzed together with outgroup, *Rhipiliopsis reticulata* (C.Hoek) Farghaly & Denizot (GenBank AF416386). All sequences were aligned using the MUSCLE software (Edgar, 2004). Maximum Likelihood (ML) phylogenetic tree was inferred in MEGA v7 (Kumar *et al.*, 2016) using GTR+G+I model. Branch support was assessed with bootstrap (1,000 replications). Maximum Likelihood bootstrap percentages (BP) values were considered as weak (below 60%), moderate (70%–79%) and strong (80%–100%). Bayesian Inference (BI) was performed in MrBayes (Huelsenbeck and Ronquist, 2001) using Markov Chain Monte Carlo chains (MCMC) of 40,000,000 generations, sampled every 4,000 generations with 10% burnin period (Draisma *et al.*, 2014). Bayesian Inference posterior probabilities (PP) were considered as no supported (below 0.90), weak (0.90–0.94) and strong (0.95–1.00).

Table 3 Collection data, GenBank accession numbers (**bold**, indicates the sequence was determined in the present study, and ✓ as not submitted to GenBank), and references for specimens used in the present study. n.a., not available.

<i>tufA</i> NCBI accession	ITS NCBI accession	GenBank name	Location	Updated name	References	Voucher specimens	Note
-	AJ007817	<i>C. mexicana</i>	San Blas Island, Panamá	-	Olsen <i>et al.</i> (1998)	RH JUNI 98 01821	
-	AJ007818	<i>C. mexicana</i>	Florida, USA	-	Olsen <i>et al.</i> (1998)	RH JUNI 98 01943	
-	AJ007819	<i>C. taxifolia</i>	Sicily, Italy	-	Olsen <i>et al.</i> (1998)	RH JUNI 98 01941	
-	AJ007823	<i>C. taxifolia</i>	Townsville, Australia	-	Olsen <i>et al.</i> (1998)	RH JUNI 98 01946	
-	AY205288	<i>C. racemosa</i> var. <i>peltata</i>	Green Island, Taiwan	<i>C. chemnitzia</i>	Yeh and Chen (2004)	n.a.	
-	AY205289	<i>C. racemosa</i> var. <i>peltata</i>	Dali, Taiwan	<i>C. chemnitzia</i>	Yeh and Chen (2004)	n.a.	
-	AY205290	<i>C. racemosa</i> var. <i>laetevirens</i>	Fengchueisha, Taiwan	<i>C. chemnitzia</i>	Yeh and Chen (2004)	n.a.	
-	AY205291	<i>C. racemosa</i> var. <i>laetevirens</i>	Kenting, Taiwan	<i>C. chemnitzia</i>	Yeh and Chen (2004)	n.a.	
-	AY206420	<i>C. racemosa</i> var. <i>macrophysa</i>	Green Island, Taiwan	<i>C. racemosa</i>	Yeh and Chen (2004)	n.a.	
-	AY206421	<i>C. racemosa</i> var. <i>macrophysa</i>	Nanwan, Taiwan	<i>C. racemosa</i>	Yeh and Chen (2004)	n.a.	
-	AY206422	<i>C. microphysa</i>	Green Island, Taiwan	-	Yeh and Chen (2004)	n.a.	
-	AY206423	<i>C. serrulata</i>	Green Island, Taiwan	-	Yeh and Chen (2004)	n.a.	
-	AY206426	<i>C. webbiana</i>	Green Island, Taiwan	-	Yeh and Chen (2004)	n.a.	
-	AF416386	<i>Rhipiliopsis reticulata</i>	Isla Galeta, Panamá	-	Kooistra <i>et al.</i> (2002)	n.a.	
-	KF840135	<i>C. corynephora</i>	Tamil Nadu, India	-	Kazi <i>et al.</i> (Unpublished)	BHAV CSIR-CSMCRI 20130329003	
-	KR478535	<i>C. racemosa</i> var. <i>macrophysa</i>	India	<i>C. racemosa</i>	Karthick <i>et al.</i> (2015)	n.a.	
AJ417928	-	<i>C. cupressoides</i> var. <i>lycopodium</i>	Uken, Japan	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417929	-	<i>C. cupressoides</i>	St. Barthélemy, Lesser Antilles	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417930	-	<i>C. cupressoides</i> var. <i>flabellata</i>	Bocas del Toro, Panamá	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417931	-	<i>C. serrulata</i>	Dahab, Egypt	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417932	-	<i>C. serrulata</i>	Pangasinan, Philippines	-	Famà <i>et al.</i> (2002)	n.a.	

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AJ417933	-	<i>C. serrulata</i>	Colón, Panamá	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417934	-	<i>C. brachypus</i>	Pangasinan, Philippines	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417936	-	<i>C. taxifolia</i>	Moreton Bay, Australia	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417937	-	<i>C. taxifolia</i>	Safaga, Egypt	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417938	-	<i>C. taxifolia</i>	Puerto-Rico, USA	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417939	-	<i>C. taxifolia</i>	N. Kwa-Zulu Natal, South Africa	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417941	-	<i>C. ashmeadii</i>	Florida, USA	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417942	-	<i>C. prolifera</i>	Bali, Indonesia	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417944	-	<i>C. sertularioides</i>	Martinique, Lesser Antilles	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417945	-	<i>C. sertularioides</i>	Colón, Panamá	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417946	-	<i>C. sertularioides</i>	Melones Island, Panamá	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417947	-	<i>C. racemosa</i> var. <i>macrophysa</i>	Galeta, Panamá	<i>C. racemosa</i>	Famà <i>et al.</i> (2002)	n.a.	
AJ417948	-	<i>C. racemosa</i> var. <i>peltata</i>	Panamá	<i>C. chemnitzia</i>	Famà <i>et al.</i> (2002)	n.a.	
AJ417949	-	<i>C. racemosa</i> var. <i>peltata</i>	Isla Naos, Panamá	<i>C. chemnitzia</i>	Famà <i>et al.</i> (2002)	n.a.	
AJ417950	-	<i>C. racemosa</i>	Galeta, Panamá	<i>C. chemnitzia</i>	Famà <i>et al.</i> (2002)	n.a.	
AJ417951	-	<i>C. mexicana</i>	Cuba	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417953	-	<i>C. mexicana</i>	Dahab, Red Sea	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417954	-	<i>C. racemosa</i> var. <i>lamourouxii</i>	Uken, Japan	<i>C. oligophylla</i>	Famà <i>et al.</i> (2002)	n.a.	
AJ417955	-	<i>C. racemosa</i> var. <i>occidentalis</i>	Livorno, Italy	<i>C. cylindracea</i>	Famà <i>et al.</i> (2002)	n.a.	
AJ417956	-	<i>C. racemosa</i> var. <i>macrophysa</i>	Florida, USA	<i>C. racemosa</i>	Famà <i>et al.</i> (2002)	n.a.	
AJ417957	-	<i>C. racemosa</i> var. <i>turbinata</i>	Dahab, Egypt	<i>C. lamourouxii</i>	Famà <i>et al.</i> (2002)	n.a.	
AJ417958	-	<i>C. webbiana</i>	Dahab, Egypt	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417959	-	<i>C. lanuginosa</i>	Florida, USA	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417960	-	<i>C. geminata</i>	Cape Bank, Australia	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417961	-	<i>C. microphysa</i>	Texas Flower Gardens, USA	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417963	-	<i>Caulerpella ambigua</i>	Texas Flower Gardens, USA	<i>C. ambigua</i>	Famà <i>et al.</i> (2002)	n.a.	
AJ417964	-	<i>C. filiformis</i>	Bronte Beach, Australia	-	Famà <i>et al.</i> (2002)	n.a.	

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AJ417965	-	<i>C. paspaloides</i>	Florida, USA	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417967	-	<i>C. verticillata</i>	Florida, USA	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417968	-	<i>C. geminata</i>	Coffs Harbour, Australia	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417969	-	<i>C. cactoides</i>	Jervis Bay, Australia	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417970	-	<i>C. flexilis</i>	Jervis Bay, Australia	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417971	-	<i>C. scalpelliformis</i>	Cape Banks, Australia	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417972	-	<i>C. scalpelliformis</i>	Damour, Lebanon	-	Famà <i>et al.</i> (2002)	n.a.	
AJ417973	-	<i>C. selago</i>	Abu Dhiab, Egypt	-	Famà <i>et al.</i> (2002)	n.a.	
AJ512413	-	<i>C. racemosa</i> var. <i>racemosa</i>	Bolinao reef, Philippines	<i>C. oligophylla</i>	de Senerpont Domis <i>et al.</i> (2003)	n.a.	
AJ512415	-	<i>C. racemosa</i> var. <i>laetevirens</i>	Bolinao reef, Philippines	<i>C. chemnitzia</i>	de Senerpont Domis <i>et al.</i> (2003)	n.a.	
AJ512417	-	<i>C. racemosa</i> var. <i>mucronata</i>	Bolinao reef, Philippines	<i>C. racemosa</i>	de Senerpont Domis <i>et al.</i> (2003)	n.a.	
AJ512418	-	<i>C. racemosa</i> var. <i>mucronata</i>	Bolinao reef, Philippines	<i>C. racemosa</i>	de Senerpont Domis <i>et al.</i> (2003)	n.a.	
AJ512426	-	<i>C. flexilis</i>	Jerbis Bay, Australia	-	de Senerpont Domis <i>et al.</i> (2003)	n.a.	
DQ652330	-	<i>C. cupressoides</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL018	
DQ652332	DQ652237	<i>C. cupressoides</i>	US Virgin Islands2, USA	-	Stam <i>et al.</i> (2006)	FL046	
DQ652345	-	<i>C. cupressoides</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL246	
DQ652346	DQ652302	<i>C. serrulata</i>	California, USA	-	Stam <i>et al.</i> (2006)	CA033	
DQ652348	DQ652304	<i>C. serrulata</i>	California, USA	-	Stam <i>et al.</i> (2006)	CA042	
DQ652349	DQ652301	<i>C. serrulata</i>	California, USA	-	Stam <i>et al.</i> (2006)	CA068	
DQ652353	-	<i>C. brachypus</i>	California, USA	-	Stam <i>et al.</i> (2006)	CA018	
DQ652354	-	<i>C. brachypus</i>	Indiana, USA	-	Stam <i>et al.</i> (2006)	FL145	
DQ652355	-	<i>C. brachypus</i>	eBay, Union, OH	-	Stam <i>et al.</i> (2006)	FL101	
DQ652360	DQ652297	<i>C. taxifolia</i>	California, USA	-	Stam <i>et al.</i> (2006)	CA067	
DQ652361	DQ652298	<i>C. taxifolia</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL131	
DQ652362	DQ652537	<i>C. ashmeadii</i>	California, USA	-	Stam <i>et al.</i> (2006)	CA007	
DQ652363	DQ652533	<i>C. ashmeadii</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL064	
DQ652372	DQ652249	<i>C. prolifera</i>	California, USA	-	Stam <i>et al.</i> (2006)	CA008	

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DQ652373	DQ652250	<i>C. prolifera</i>	California, USA	-	Stam <i>et al.</i> (2006)	CA031	
DQ652376	DQ652247	<i>C. prolifera</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL085	
DQ652393	DQ652290	<i>C. sertularioides</i>	California, USA	-	Stam <i>et al.</i> (2006)	CA037	
DQ652400	DQ652271	<i>C. sertularioides</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL033	
DQ652409	DQ652283	<i>C. sertularioides</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL136	
DQ652420	DQ652265	<i>C. racemosa</i>	Aquarium trade, California, USA	-	Stam <i>et al.</i> (2006)	CA014	
DQ652422	DQ652263	<i>C. racemosa</i>	Aquarium trade, California, USA	-	Stam <i>et al.</i> (2006)	CA060	
DQ652424	-	<i>C. racemosa</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL030	
DQ652425	-	<i>C. racemosa</i>	US Virgin Islands2, USA	-	Stam <i>et al.</i> (2006)	FL045	
DQ652426	-	<i>C. racemosa</i>	Florida, USA	<i>C. nummularia</i>	Stam <i>et al.</i> (2006)	FL108	
DQ652428	-	<i>C. racemosa</i>	Aquarium shop (Florida, USA)	<i>C. macrodisca</i>	Stam <i>et al.</i> (2006)	FL158	
DQ652429	-	<i>C. racemosa</i>	Aquarium shop (Florida, USA)	<i>C. macrodisca</i>	Stam <i>et al.</i> (2006)	FL159	
-	✓	<i>C. racemosa</i>	Aquarium trade, California, USA	<i>C. elongata</i>	Stam <i>et al.</i> (2006), This study	CA049	for ITS: clone 1
DQ652433	DQ652238	<i>C. mexicana</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL003	
DQ652437	DQ652239	<i>C. mexicana</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL017	
DQ652446	DQ652241	<i>C. mexicana</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL077	
DQ652457	DQ652244	<i>C. mexicana</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL199	
DQ652467	DQ652321	<i>C. racemosa</i>	California, USA	<i>C. chemnitzia</i>	Stam <i>et al.</i> (2006)	CA053	
DQ652468	DQ652323	<i>C. racemosa</i>	California, USA	<i>C. chemnitzia</i>	Stam <i>et al.</i> (2006)	CA056	
DQ652470	-	<i>C. racemosa</i>	Florida, USA	<i>C. chemnitzia</i>	Stam <i>et al.</i> (2006)	FL011	
DQ652474	DQ652311	<i>C. racemosa</i>	Florida, USA	<i>C. chemnitzia</i>	Stam <i>et al.</i> (2006)	FL055	
DQ652494	DQ652259	<i>C. racemosa</i>	California, USA	<i>C. oligophylla</i>	Stam <i>et al.</i> (2006)	CA040	
DQ652495	DQ652251	<i>C. racemosa</i>	California, USA	<i>C. cylindracea</i>	Stam <i>et al.</i> (2006)	CA034	
DQ652496	-	<i>C. lanuginosa</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL166	
DQ652499	-	<i>C. paspaloides</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL013	
DQ652513	DQ652326	<i>C. microphysa</i>	California, USA	-	Stam <i>et al.</i> (2006)	CA041	
DQ652514	DQ652327	<i>C. microphysa</i>	California, USA	-	Stam <i>et al.</i> (2006)	CA045	
DQ652525	-	<i>C. verticillata</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL006	
DQ652530	-	<i>C. verticillata</i>	Florida, USA	-	Stam <i>et al.</i> (2006)	FL175	
FJ432655	-	<i>Caulerpella ambigua</i>	Hawaii, USA	<i>C. ambigua</i>	Draisma <i>et al.</i> (2014)	TS0024	

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FJ810424	-	<i>C. freycinetii</i> var. <i>integerrima</i>	Eilat, Israel	<i>C. integerrima</i>	Wynne <i>et al.</i> (2009)	n.a.	
FJ810425	-	<i>C. freycinetii</i> var. <i>integerrima</i>	Eilat, Israel	<i>C. integerrima</i>	Wynne <i>et al.</i> (2009)	n.a.	
FJ810426	-	<i>C. bartoniae</i>	South Africa	-	Wynne <i>et al.</i> (2009)	n.a.	
FM956011	-	<i>C. biserrulata</i>	Papua, Indonesia	-	Draisma and Prud'homme van Reine (Unpublished)	L SGAD0712179	
FM956012	-	<i>C. brachypus</i>	Thousand Islands, Indonesia	-	Draisma and Prud'homme van Reine (Unpublished)	L SGAD0509301	
FM956013	-	<i>C. brachypus</i>	Papua, Indonesia	-	Draisma and Prud'homme van Reine (Unpublished)	L SGAD0712020	
FM956014	-	<i>C. cupressoides</i>	Thousand Islands, Indonesia	-	Draisma and Prud'homme van Reine (Unpublished)	L SGAD0509213	
FM956015	-	<i>C. serrulata</i> cf. var. <i>pectinata</i>	Thousand Islands, Indonesia	<i>C. serrulata</i> var. <i>pectinata</i>	Draisma and Prud'homme van Reine (Unpublished)	L SGAD0509189	
FM956016	-	<i>C. lessonii</i>	Thousand Islands, Indonesia	-	Draisma and Prud'homme van Reine (Unpublished)	L SGAD0509589	
FM956017	-	<i>C. serrulata</i> f. <i>spiralis</i>	Thousand Islands, Indonesia	-	Draisma and Prud'homme van Reine (Unpublished)	L SGAD0509248	
FM956018	-	<i>C. serrulata</i> f. <i>spiralis</i>	Thousand Islands, Indonesia	-	Draisma and Prud'homme van Reine (Unpublished)	L SGAD0509249	
FM956020	✓	<i>C. serrulata</i> f. <i>spiralis</i>	Red Sea, Egypt	-	Draisma and Prud'homme van Reine (Unpublished), This study	L SGAD1033	
FM956021	✓	<i>C. lentillifera</i>	Java, Indonesia	-	Draisma <i>et al.</i> (2014), This study	L SGAD 0509343	for ITS: clone 1
-	✓	<i>C. lentillifera</i>	Papua, Indonesia	-	Draisma and Prud'homme van Reine (Unpublished), This study	L SGAD 0712132	for ITS: clone 1
FM956024	-	<i>C. lentillifera</i>	Papua, Indonesia	-	Draisma <i>et al.</i> (2014)	L SGAD 0712757	
FM956025	✓	<i>C. microphysa</i>	Tanzania	-	Draisma <i>et al.</i> (2014), This study	GENT TZ0270	for ITS: clone 1
FM956026	✓	<i>C. opposita</i>	Borneo, Indonesia	-	Draisma <i>et al.</i> (2014), This study	L 03-038	for ITS: clone 1
FM956027	-	<i>C. microphysa</i>	Sri Lanka	-	Draisma <i>et al.</i> (2014)	GENT HEC11838	
FM956029	-	<i>C. parvifolia</i>	Papua, Indonesia	-	Draisma and Prud'homme van Reine (Unpublished)	L SGAD0712794	
FM956031	-	<i>C. fastigiata</i>	Borneo, Indonesia	-	Draisma <i>et al.</i> (2014)	L 03-36	
FM956034	-	<i>C. filicoides</i>	Borneo, Indonesia	-	Draisma <i>et al.</i> (2014)	L 03-039	
FM956035	-	<i>C. filiformis</i>	South Africa	-	Draisma <i>et al.</i> (2014)	L 2005-31	
FM956036	-	<i>C. flexilis</i>	New Zealand	-	Draisma <i>et al.</i> (2014)	L 1998	
FM956037	-	<i>C. brownii</i>	New Zealand	-	Draisma <i>et al.</i> (2014)	L 81 (5)	
FM956038	-	<i>C. trifaria</i>	Tasmania, Australia	-	Draisma <i>et al.</i> (2014)	L L14	

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FM956039	-	<i>C. sedoides</i> f. <i>sedoides</i>	New Zealand	<i>C. sedoides</i>	Draisma <i>et al.</i> (2014)	L 6	
FM956040	-	<i>C. longifolia</i>	Tasmania, Australia	-	Draisma <i>et al.</i> (2014)	L L15	
FM956041	✓	<i>C. lanuginosa</i> var. <i>delicatula</i>	Tanzania	<i>C. antoensis</i>	Draisma and Prud'homme van Reine (Unpublished), This study	GENT TZ0503	for ITS: clone 1
FM956042	-	<i>C. prolifera</i>	Ischia, Italy	-	Draisma and Prud'homme van Reine (Unpublished)	L SGAD 0706105	
FM956043	✓	<i>C. racemosa</i> f. <i>requieni</i>	Thousand Islands, Indonesia	<i>C. oligophylla</i>	Sauvage <i>et al.</i> (2013), This study	SGAD 0509242	for ITS: clone 1
FM956044	✓	<i>C. racemosa</i> f. <i>requieni</i>	Berau delta, Indonesia	<i>C. oligophylla</i>	Draisma and Prud'homme van Reine (Unpublished), This study	L 03-040	for ITS: clone 1
FM956045	MK481938	<i>C. racemosa</i> f. <i>clavifera</i>	Papua, Indonesia	<i>C. racemosa</i>	Sauvage <i>et al.</i> (2013), This study	SGAD 0712247	for ITS: Clone 1
FM956046	✓	<i>C. racemosa</i> var. <i>macra</i>	Papua, Indonesia	<i>C. macra</i>	Sauvage <i>et al.</i> (2013), This study	SGAD 0509331	for ITS: Clone 1
FM956048	✓	<i>C. racemosa</i> f. <i>cylindracea</i>	Gulf of Naples, Italy	<i>C. cylindracea</i>	Sauvage <i>et al.</i> (2013), This study	SGAD 0706109	for ITS: Clone 1
FM956051	MK481937, MK481943	<i>C. racemosa</i> var. <i>occidentalis</i>	Thousand Islands, Indonesia	<i>C. racemosa</i>	Sauvage <i>et al.</i> (2013), This study	SGAD 0509638	for ITS: Clone 1, 2
FM956052	-	<i>C. racemosa</i> f. <i>macrophysa</i>	Berau delta, Indonesia	<i>C. racemosa</i>	Sauvage <i>et al.</i> (2013)	L 03–342	
FM956053*	-	<i>C. racemosa</i> var. <i>chemnitzia</i>	Thousand Islands, Indonesia	<i>C. macrodisca</i>	Sauvage <i>et al.</i> (2013)	SGAD 0509359	
FM956054	-	<i>C. racemosa</i> var. <i>chemnitzia</i>	Berau delta, Indonesia	<i>C. macrodisca</i>	Sauvage <i>et al.</i> (2013)	L 03–341	
FM956055	MK481944- MK481949	<i>C. peltata</i> var. <i>peltata</i>	Papua, Indonesia	<i>C. macrodisca</i>	Sauvage <i>et al.</i> (2013), This study	SGAD 0712635	for ITS: Clone 1-6
FM956056	-	<i>C. nummularia</i>	Thousand Islands, Indonesia	<i>C. chemnitzia</i>	Sauvage <i>et al.</i> (2013)	SGAD 0509640	
FM956057	✓	<i>C. nummularia</i>	Berau delta, Indonesia	<i>C. chemnitzia</i>	Sauvage <i>et al.</i> (2013), This study	L 03–227	
FM956058	-	<i>C. nummularia</i>	Ko Kham, Songkhla, Thailand	<i>C. chemnitzia</i>	Draisma <i>et al.</i> (2014)	GENT <i>Caulerpa peltata</i> 11-12-2006	
FM956059	✓	<i>C. nummularia</i>	Papua, Indonesia	<i>C. chemnitzia</i>	Draisma and Prud'homme van Reine (Unpublished), This study	L SGAD 0712145	for ITS: Clone 1
FM956060	-	<i>C. peltata</i> f. <i>peltata</i>	Philippines	<i>C. chemnitzia</i>	Draisma and Prud'homme van Reine (Unpublished)	GENT FL1176	

<i>tufA</i> NCBI accession	ITS NCBI accession	GenBank name	Location	Updated name	References	Voucher specimens	Note
FM956061	✓	<i>C. selago</i> cf. var. <i>selago</i>	Berau delta, Indonesia	<i>C. chemnitzia</i>	Draisma and Prud'homme van Reine (Unpublished), This study	L 03-128	for ITS: Clone 1
FM956062	-	<i>C. scalpelliformis</i>	Tasmania, Australia	-	Draisma <i>et al.</i> (2014)	L L13	
FM956063	-	<i>C. scalpelliformis</i>	South Africa	-	Draisma <i>et al.</i> (2014)	L 2005-12	
FM956064	-	<i>C. sertularioides</i> f. <i>brevipes</i>	Thousand Islands, Indonesia	-	Draisma and Prud'homme van Reine (Unpublished)	L SGAD 0509214	
FM956065	-	<i>C. taxifolia</i>	Thousand Islands, Indonesia	-	Draisma and Prud'homme van Reine (Unpublished)	L SGAD 0509244	
FM956066	-	<i>C. taxifolia</i>	Philippines	-	Draisma and Prud'homme van Reine (Unpublished)	GENT FL1126	
FM956067	-	<i>C. urvilleana</i>	Papua, Indonesia	-	Draisma and Prud'homme van Reine (Unpublished)	L SGAD 0712203	
FM956070	✓	<i>C. verticillata</i>	Berau delta, Indonesia	-	Draisma <i>et al.</i> (2014), This study	L 03-130	
FM956072	-	<i>C. verticillata</i>	Java, Indonesia	-	Draisma <i>et al.</i> (2014)	L 0509287	
FM956073	-	<i>C. webbiana</i> f. <i>disticha</i>	Azores, Portugal	-	Draisma <i>et al.</i> (2014)	n.a.	
FM956074	-	<i>C. webbiana</i>	Tanzania	-	Draisma <i>et al.</i> (2014)	GENT TZ0486	
FM956075	-	<i>C. racemosa</i> cf. var. <i>corynephora</i>	Khlong Yang, Krabi, Thailand	<i>C. macrodisca</i>	Sauvage <i>et al.</i> (2013)	HEC161156	
FN667648	-	<i>C. webbiana</i> f. <i>disticha</i>	Velasco, Palau	-	Draisma <i>et al.</i> (2014)	L V2.25.05	
FN667649	-	<i>C. manorensis</i>	Hall Island, Palau	<i>Caulerpa</i> sp.	Draisma <i>et al.</i> (2014)	L H6-25-04	
FN667650	-	<i>C. okamurae</i>	Japan	-	Draisma <i>et al.</i> (2014)	L SGAD 0909013, L:H6-25-04	
FR848331	-	<i>C. corynephora</i>	Western Australia	-	Draisma <i>et al.</i> (2014)	PERTH 6.10.9.2.6	
FR848332	-	<i>C. elongata</i>	Northern Reef, Palau	-	Draisma <i>et al.</i> (2014)	L V2.25.04	
FR848333	-	<i>C. simpliciuscula</i>	Victoria, Australia	-	Draisma <i>et al.</i> (2014)	L 09.10.048	
FR848334	-	<i>C. remotifolia</i>	Victoria, Australia	-	Draisma <i>et al.</i> (2014)	L 09.10.074	
FR848335	-	<i>C. obscura</i>	Victoria, Australia	-	Draisma <i>et al.</i> (2014)	L 09.10.052	
FR848337	-	<i>C. antoensis</i>	Northern Reef, Palau	-	Draisma and Prud'homme van Reine (Unpublished)	L NR03-25-08	
FR848338	-	<i>C. longifolia</i>	Victoria, Australia	-	Draisma <i>et al.</i> (2014)	L 09.10.042	
FR848339	-	<i>C. parvifolia</i>	Rottneest Island, Australia	-	Draisma <i>et al.</i> (Unpublished)	n.a.	
FR848340	-	<i>C. cliftonii</i>	Victoria, Australia	-	Draisma <i>et al.</i> (2014)	L 09.10.051	
FR848341	-	<i>C. brownii</i>	Victoria, Australia	-	Draisma <i>et al.</i> (2014)	L 09.10.057	

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FR848342	-	<i>C. racemosa</i> cf. var. <i>macra</i>	Palau	<i>C. macra</i>	Draisma <i>et al.</i> (Unpublished)	L C18-05-23	
FR848343	-	<i>C. sedoides</i> f. <i>geminata</i>	Western Australia	<i>C. sedoides</i>	Draisma <i>et al.</i> (2014)	PERTH -	
FR848344	-	<i>C. filicoides</i> var. <i>andamanensis</i>	Chuuk, Federated States of Micronesia	<i>C. andamanensis</i>	Draisma <i>et al.</i> (2014)	L C7-05-09	
GU592621	-	Ulvophyceae	Lizard Island, Australia	<i>C. nummularia</i>	Händeler <i>et al.</i> (2010)	n.a.	
GU592622	-	Ulvophyceae	Lizard Island, Australia	<i>C. nummularia</i>	Händeler <i>et al.</i> (2010)	n.a.	
JN645149	-	<i>C. racemosa</i>	Grande Terre, New Caledonia	<i>C. megadisca</i>	Sauvage <i>et al.</i> (2013)	IRD5639	
JN645150	-	<i>C. racemosa</i>	Porquerolles, France	<i>C. cylindracea</i>	Sauvage <i>et al.</i> (2013)	TS0133	
JN645152	-	<i>C. racemosa</i>	Fiji	<i>C. macra</i>	Sauvage <i>et al.</i> (2013)	IRD1878	
JN645153	-	<i>C. taxifolia</i>	Goeland Island, New Caledonia	-	Sauvage <i>et al.</i> (2013)	IRD5637	
JN645154	-	<i>C. racemosa</i>	Ile des Pins, New Caledonia	<i>C. megadisca</i>	Sauvage <i>et al.</i> (2013)	IRD5636	
JN645160	-	<i>C. racemosa</i>	Ile des Pins, New Caledonia	<i>C. chemnitzia</i>	Sauvage <i>et al.</i> (2013)	IRD5627	
JN645162	-	<i>C. racemosa</i>	Grande Terre, New Caledonia	<i>C. chemnitzia</i>	Sauvage <i>et al.</i> (2013)	IRD5638	
JN645163	-	<i>C. racemosa</i>	Grande Terre, New Caledonia	<i>C. cylindracea</i>	Sauvage <i>et al.</i> (2013)	IRD5629	
JN645164	-	<i>C. racemosa</i>	Ile des Pins, New Caledonia	<i>C. macra</i>	Sauvage <i>et al.</i> (2013)	IRD5624	
JN645165	-	<i>C. racemosa</i>	Grande Terre, New Caledonia	<i>C. macrodisca</i>	Sauvage <i>et al.</i> (2013)	IRD5635	
JN645167	-	<i>C. racemosa</i>	Grande Terre, New Caledonia	<i>C. macra</i>	Sauvage <i>et al.</i> (2013)	IRD5640	
JN645168	-	<i>C. racemosa</i>	Ile des Pins, New Caledonia	<i>C. oligophylla</i>	Sauvage <i>et al.</i> (2013)	IRD5633	
JN645169	-	<i>C. racemosa</i>	Grande Terre, New Caledonia	-	Sauvage <i>et al.</i> (2013)	IRD5641	
JN645170	-	<i>C. racemosa</i>	Grande Terre, New Caledonia	<i>C. oligophylla</i>	Sauvage <i>et al.</i> (2013)	IRD5642	
JN645173	-	<i>C. racemosa</i>	Minnie Waters, NSW, Australia	<i>C. nummularia</i>	Sauvage <i>et al.</i> (2013)	HV2132	
JN645174	-	<i>C. racemosa</i>	Hopetoun Beach, SWA, Australia	<i>C. cylindracea</i>	Sauvage <i>et al.</i> (2013)	HV2537	
JN645175	-	<i>C. racemosa</i>	Leander Reef, Australia	<i>C. cylindracea</i>	Sauvage <i>et al.</i> (2013)	HV2634	
JN817646	-	<i>C. obscura</i>	Australia	-	Belton <i>et al.</i> (2014)	AD A90909	
JN817647	-	<i>C. opposita</i>	Lizard Island, Australia	-	Draisma <i>et al.</i> (2014)	AD A92544	

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JN817653	-	<i>C. corynephora</i>	Australia	-	Belton <i>et al.</i> (2014)	PERTH 08292590	
JN817655	-	<i>C. fergusonii</i>	Western Australia	-	Draisma <i>et al.</i> (2014)	AD A91682	
JN817656	-	<i>Caulerpa</i> sp.	Lizard Island, Australia	<i>C. megadisca</i>	Belton <i>et al.</i> (2014)	AD-A92609	
JN817657*	-	<i>Caulerpa</i> sp.	Lizard Island, Australia	<i>C. megadisca</i>	Belton <i>et al.</i> (2014)	AD-A90107 (holotype)	
JN817659	-	<i>C. racemosa</i>	Long Reef, Kimberley, Australia	-	Belton <i>et al.</i> (2014)	PERTH 08292736	
JN817661	-	<i>C. racemosa</i>	Lizard Island, Australia	-	Belton <i>et al.</i> (2014)	AD-A92441	
JN817665*	-	<i>C. racemosa</i>	Montgomery Reef, Kimberley, Australia	-	Belton <i>et al.</i> (2014)	PERTH 08292728	
JN817666	-	<i>C. macrodisca</i>	Lizard Island, Australia	-	Belton <i>et al.</i> (2014)	AD-A88056	
JN817667	-	<i>C. peltata</i>	Montgomery Reef, Australia	<i>C. chemnitzia</i>	Belton <i>et al.</i> (2014)	PERTH 08292671	
JN817668	-	<i>C. peltata</i>	Montgomery Reef, Australia	<i>C. chemnitzia</i>	Belton <i>et al.</i> (2014)	PERTH 08292698	
JN817669	-	<i>C. chemnitzia</i>	Cassini Island, Australia	-	Belton <i>et al.</i> (2014)	PERTH 08292701	
JN817673	-	<i>C. cylindracea</i>	Ningaloo Reef, Australia	-	Belton <i>et al.</i> (2014)	AD-A88236	
JN817675	-	<i>C. cylindracea</i>	Ningaloo Reef, Australia	-	Sauvage <i>et al.</i> (2013)	AD-A91736	
JN817681	-	<i>C. racemosa</i>	Long Reef, Australia	<i>C. lamourouxii</i>	Belton <i>et al.</i> (2014)	PERTH 08292647	
JN817683	-	<i>C. racemosa</i>	Ningaloo Reef, Australia	<i>C. lamourouxii</i>	Belton <i>et al.</i> (2014)	AD-A90154	
JN817685*	-	<i>C. nummularia</i>	Heron I., Australia	-	Belton <i>et al.</i> (2014)	AD-A91369	
JN851135	-	<i>C. cactoides</i>	Australia	-	Belton <i>et al.</i> (2014)	GWS024470	
JN851137	-	<i>C. fergusonii</i>	Australia	-	Belton <i>et al.</i> (2014)	GWS025259	
JN851138	-	<i>C. nummularia</i>	Lord Howe Island, Australia	-	Belton <i>et al.</i> (2014)	GWS023933	
JN851139	-	<i>C. nummularia</i>	Lord Howe Island, Australia	-	Belton <i>et al.</i> (2014)	GWS023932	
JN851140	-	<i>C. nummularia</i>	Lord Howe Island, Australia	-	Belton <i>et al.</i> (2014)	GWS023180	
JN851143*	-	<i>C. cylindracea</i>	Point Peron, WA, Australia	-	Sauvage <i>et al.</i> (2013)	GWS025471	
JN851146	-	<i>C. flexilis</i>	Australia	-	Belton <i>et al.</i> (2014)	GWS015249	
KC153492	JF932264	<i>C. scalpelliformis</i>	Veraval, Gujarat, India	-	Kazi <i>et al.</i> (2013)	19102009001	
KC153493	JF932265	<i>C. verticillata</i>	Okha, Gujarat, India	-	Kazi <i>et al.</i> (2013)	17102009001	

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KC153495	JF932267	<i>C. racemosa</i>	Veraval, Gujarat, India	-	Kazi <i>et al.</i> (2013)	4122009001	
KC153497	JF932269	<i>C. microphysa</i>	Okha, Gujarat, India	-	Kazi <i>et al.</i> (2013)	2122009001	
KC153499	JF932271	<i>C. sertularioides</i> f. <i>brevipes</i>	Porbandar, Gujarat, India	-	Kazi <i>et al.</i> (2013)	3122009003	
KC153503	JF932275	<i>C. scalpelliformis</i> var. <i>denticulata</i>	Okha, Gujarat, India	<i>C. denticulata</i>	Kazi <i>et al.</i> (2013)	2122010002	
KC153505	JN034414	<i>C. lentillifera</i>	Poshitra rocks, Gujarat, India	-	Kazi <i>et al.</i> (2013)	20110419014	
KC153508	JQ745710	<i>C. taxifolia</i>	Poshitra rocks, Gujarat, India	-	Kazi <i>et al.</i> (2013)	20110420001	
KC153509	JQ745711	<i>C. serrulata</i>	Sadamunian, Tamilnadu, India	-	Kazi <i>et al.</i> (2013)	20101230001	
KC153510	JQ745712	<i>C. peltata</i>	Manapadu, Tamilnadu, India	<i>C. chemnitzia</i>	Kazi <i>et al.</i> (2013)	20110103001	
KC153512	JQ745714	<i>C. scalpelliformis</i>	Mandapam, Tamilnadu, India	-	Kazi <i>et al.</i> (2013)	20110104001	
KC153513	JQ745715	<i>C. sertularioides</i> f. <i>longipes</i>	Poshitra rocks, Gujarat, India	-	Kazi <i>et al.</i> (2013)	20110419019	
KC153514	JQ745716	<i>C. veravalensis</i>	Okha, Gujarat, India	-	Kazi <i>et al.</i> (2013)	20101202001	
KC153515	JQ745717	<i>C. taxifolia</i>	Kunkeshwar, Maharashtra, India	-	Kazi <i>et al.</i> (2013)	20110223001	
KC153517	JQ745719	<i>C. verticillata</i>	Krusadai Island, Tamilnadu, India	-	Kazi <i>et al.</i> (2013)	20110110001	
KC153519	JQ745721	<i>C. peltata</i>	Ratnagiri, Maharashtra, India	<i>C. chemnitzia</i>	Kazi <i>et al.</i> (2013)	20110225001	
KF256081*	-	<i>C. racemosa</i> var. <i>lamourouxii</i>	Marsa Alam, Red Sea, Egypt	<i>C. lamourouxii</i>	Belton <i>et al.</i> (2014)	HV03477	
KF256084	-	<i>C. racemosa</i> var. <i>lamourouxii</i>	Cassini Island, Australia	<i>C. lamourouxii</i>	Belton <i>et al.</i> (2014)	PERTH 08428220	
KF256085*	-	<i>C. oligophylla</i>	Heron Island, Australia	-	Belton <i>et al.</i> (2014)	AD-A95457	
KF256086	-	<i>C. oligophylla</i>	Thousand Islands, Indonesia	-	Belton <i>et al.</i> (2014)	SGAD0509292	
KF256088	-	<i>C. racemosa</i> var. <i>macra</i>	Northern Reefs, Palau	<i>C. macra</i>	Belton <i>et al.</i> (2014)	LNR13a0109	
KF256089*	-	<i>C. racemosa</i> var. <i>macra</i>	Berau delta, Indonesia	<i>C. macra</i>	Belton <i>et al.</i> (2014)	L 03-453	
KF256090	-	<i>C. macrodisca</i>	Long Reef, Australia	-	Belton <i>et al.</i> (2014)	PERTH 08292663	
KF256091	-	<i>Caulerpa</i> sp.	Yaukuvelailai Island, Fiji	<i>C. megadisca</i>	Belton <i>et al.</i> (2014)	DML40342	
KF256092	-	<i>C. macrodisca</i>	Thousand Islands, Indonesia	-	Belton <i>et al.</i> (2014)	SGAD0509415	
KF256093	-	<i>C. macrodisca</i>	Java Sea, Indonesia	-	Belton <i>et al.</i> (2014)	SGAD0509390	

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KF256094	-	<i>C. macrodisca</i>	Java Sea, Indonesia	-	Belton <i>et al.</i> (2014)	SGAD0509510	
KF256095	-	<i>C. macrodisca</i>	Papua, Indonesia	-	Belton <i>et al.</i> (2014)	SGAD0712405	
KF256097	-	<i>C. racemosa</i>	Papua, Indonesia	-	Belton <i>et al.</i> (2014)	L0789171	
KF256098	-	<i>C. nummularia</i>	Dravuni, Fiji	-	Belton <i>et al.</i> (2014)	DML40015	
KF256099	-	<i>C. chemnitzia</i>	St George's, Bermuda	-	Belton <i>et al.</i> (2014)	CWS008555	
KF256100	-	<i>C. chemnitzia</i>	Blue cut channel, Bermuda	-	Belton <i>et al.</i> (2014)	CWS008366	
KF256101*	-	<i>C. chemnitzia</i>	Matara, Sri Lanka	-	Belton <i>et al.</i> (2014)	HEC15952	
KF256105	-	<i>C. chemnitzia</i>	Lizard Island, Australia	-	Belton <i>et al.</i> (2014)	AD-A92551	
KF256106	-	<i>C. chemnitzia</i>	Exumas, Bahamas	-	Belton <i>et al.</i> (2014)	DML70471	
KF314138	-	<i>C. corynephora</i>	Australia	-	Belton <i>et al.</i> (2014)	PERTH 08195803	
KF314139	-	<i>C. agardhii</i>	Australia	-	Draisma <i>et al.</i> (2014)	PERTH 08399697	
KF314142	-	<i>C. cupressoides</i>	Australia	-	Draisma <i>et al.</i> (2014)	AD A91732	
KF314149	-	<i>C. constricta</i>	Australia	-	Belton <i>et al.</i> (2014)	AD A94521	
KF314158	-	<i>C. brachypus</i>	Australia	-	Draisma <i>et al.</i> (2014)	AD A91664	
KF649854	-	<i>C. cliffonii</i>	Australia	-	Belton <i>et al.</i> (2014)	AD A79076	
KF649856	-	<i>C. brownii</i>	Australia	-	Belton <i>et al.</i> (2014)	AD A92244	
KF649860	-	<i>C. trifaria</i>	Australia	-	Belton <i>et al.</i> (2014)	AD A92257	
KF649862	-	<i>C. alternans</i>	Australia	-	Belton <i>et al.</i> (2014)	AD A88966	
KF649865	-	<i>C. flexilis</i>	Australia	-	Draisma <i>et al.</i> (2014)	AD A92238	
KF649869	-	<i>C. hedleyi</i>	Australia	-	Draisma <i>et al.</i> (2014)	AD A89143	
KF649870	-	<i>C. fergusonii</i>	Australia	-	Belton <i>et al.</i> (2014)	AD A93587	
KF649873	-	<i>C. simpliciuscula</i>	Australia	-	Draisma <i>et al.</i> (2014)	AD A89133	
KF649874	-	<i>C. simpliciuscula</i>	Australia	-	Belton <i>et al.</i> (2014)	GWS015874	
KF649875	-	<i>C. simpliciuscula</i>	Australia	-	Belton <i>et al.</i> (2014)	AD A92243	
KF649876*	-	<i>Caulerpa</i> sp.	Australia	<i>C. lucasii</i>	Draisma <i>et al.</i> (2014)	AD A88572	
KF649878	-	<i>C. vesiculifera</i>	Kangaroo Island, Australia	-	Draisma <i>et al.</i> (2014)	AD A92234	
KF649879	-	<i>C. papillosa</i>	Kangaroo Island, Australia	-	Draisma <i>et al.</i> (2014)	AD A92242	
KF649883	-	<i>C. hodgkinsoniae</i>	Australia	-	Belton <i>et al.</i> (2019)	GWS032688	

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KF649891	-	<i>C. remotifolia</i>	Australia	-	Belton <i>et al.</i> (2019)	AD A92285	
KF649893	-	<i>C. remotifolia</i>	Australia	-	Belton <i>et al.</i> (2019)	AD A93751	
KF649900	-	<i>C. taxifolia</i>	Australia	-	Draisma <i>et al.</i> (2014)	AD A89134	
KF649902	-	<i>C. filiformis</i>	Australia	-	Belton <i>et al.</i> (2014)	AD A95186	
KF649914	-	<i>C. longifolia</i>	Australia	-	Belton <i>et al.</i> (2014)	AD A92232	
KF921072	-	<i>C. floridana</i>	Florida	-	Sauvage <i>et al.</i> (2014)	WRT345	
KM016916	-	<i>C. sedoides</i>	Australia	-	Draisma <i>et al.</i> (2014)	AD A89135	
KM186523	-	<i>Caulerpella ambigua</i>	Hawaii, USA	<i>C. ambigua</i>	Draisma <i>et al.</i> (2014)	TS0252	
KM186527	-	<i>Caulerpella ambigua</i>	Kerama Island, Japan	<i>C. ambigua</i>	Draisma <i>et al.</i> (2014)	TS1180	
KM186528	-	<i>C. okamurae</i>	Japan	-	Draisma <i>et al.</i> (2014)	GENT HV1888	
KM186529	-	<i>C. filicoides</i> var. <i>andamanensis</i>	Hawaii, USA	<i>C. andamanensis</i>	Draisma <i>et al.</i> (2014)	ARS1611 /HS2004-140	
MK497059	-	<i>C. macrodisca</i>	East Kalimantan, Indonesia	-	This study	L_03_344	
MK497060	-	<i>C. macrodisca</i>	Java Sea, Indonesia	-	This study	SGAD0509332	
MK497061	-	<i>C. macrodisca</i>	Java Sea, Indonesia	-	This study	SGAD0509539	
MK497062	-	<i>C. macrodisca</i>	West Papua, Indonesia	-	This study	SGAD0712194	
MK497063	-	<i>C. macrodisca</i>	West Papua, Indonesia	-	This study	SGAD0712198	
MK497064	-	<i>C. macrodisca</i>	West Papua, Indonesia	-	This study	SGAD0712199	
MH591082	-	<i>Dichotomosiphon tuberosus</i>	n.a.	-	Cremen <i>et al.</i> (2019)	Shimada 1401	
MK497053	MK481940	<i>C. macrodisca</i>	Market, Krabi	-	This study	KP4D	
✓	✓	<i>C. verticillata</i>	Ko Lidee, Satun	-	This study	KP5A	
✓	✓	<i>C. racemosa</i>	Ko Tan, Surat Thani	-	This study	KP10A	
✓	✓	<i>C. serrulata</i>	Ko Tan, Surat Thani	-	This study	KP11B	
✓	✓	<i>C. verticillata</i>	Ko Tan, Surat Thani	-	This study	KP13A	
✓	✓	<i>C. serrulata</i>	Big Buddha, Ko Samui, Surat Thani	-	This study	KP14C	
✓	✓	<i>C. sertularioides</i>	Big Buddha, Ko Samui, Surat Thani	-	This study	KP15C	
✓	✓	<i>C. lentillifera</i>	Ko Siboya, Krabi	-	This study	KP21A	
✓	✓	<i>C. racemosa</i>	Ko Siboya, Krabi	-	This study	KP22A	

<i>tufA</i> NCBI accession	ITS NCBI accession	GenBank name	Location	Updated name	References	Voucher specimens	Note
✓	✓	<i>C. taxifolia</i>	Ko Siboya, Krabi	-	This study	KP23A	
✓	✓	<i>C. racemosa</i>	Ko Kham, Songkhla	-	This study	KP28B	
✓	✓	<i>C. chemnitzia</i>	Ko Kham, Songkhla	-	This study	KP29A	
✓	✓	<i>C. racemosa</i>	Ko Kham, Songkhla	-	This study	KP30A	
✓	✓	<i>C. chemnitzia</i>	Ko Kham, Songkhla	-	This study	KP31A	
✓	✓	<i>C. chemnitzia</i>	Ko Kham, Songkhla	-	This study	KP32A	
✓	✓	<i>C. sertularioides</i>	Ko Lidee, Satun	-	This study	KP35A	
✓	✓	<i>C. racemosa</i>	Ko Lipe, Satun	-	This study	KP39C	
✓	✓	<i>C. racemosa</i>	Ko Lipe, Satun	-	This study	KP40A	
✓	✓	<i>C. serrulata</i>	Ko Lipe, Satun	-	This study	KP41A	
✓	-	<i>C. verticillata</i>	Ko Lipe, Satun	-	This study	KP42A	
✓	✓	<i>C. taxifolia</i>	Ko Libong, Trang	-	This study	KP43A	
✓	✓	<i>C. serrulata</i>	Taling Ngam, Ko Samui, Surat Thani	-	This study	KP45A	
✓	✓	<i>C. taxifolia</i>	Ko Matsum, Surat Thani	-	This study	KP46B	
✓	✓	<i>C. serrulata</i>	Ko Matsum, Surat Thani	-	This study	KP47B	
✓	✓	<i>C. racemosa</i>	Losin, Pattani	-	This study	KP49A	
✓	✓	<i>C. chemnitzia</i>	Losin, Pattani	-	This study	KP50A	
✓	✓	<i>C. racemosa</i>	Losin, Pattani	-	This study	KP51A	
✓	✓	<i>C. taxifolia</i>	Leam Yong Lam, Trang	-	This study	KP53A	
✓	✓	<i>C. taxifolia</i>	Leam Yong Lam, Trang	-	This study	KP55A	
✓	✓	<i>C. taxifolia</i>	Khao Bae Na, Trang	-	This study	KP57A	
MK497054	MK481941	<i>C. macrodisca</i>	Fish cage, Kilim river, Malaysia	-	This study	KP65A	
✓	✓	<i>C. racemosa</i>	Bakan Teang, Ko Lanta, Krabi	-	This study	KP68A	
✓	✓	<i>C. racemosa</i>	Bakan Teang, Ko Lanta, Krabi	-	This study	KP70A	
✓	✓	<i>C. serrulata</i>	Tangkhen Bay, Phuket	-	This study	KP72A	
MK497055	MK481942	<i>C. macrodisca</i>	Fish cage, Khlong Yang, Krabi	-	This study	KP73A	
MK497056	-	<i>C. macrodisca</i>	Fish cage, Khlong Yang, Krabi	-	This study	KP74A	
-	MK481939	<i>C. macrodisca</i>	Ao Cho, Trat	-	This study	KUMF04404	

<i>tufA</i> NCBI accession	ITS NCBI accession	GenBank name	Location	Updated name	References	Voucher specimens	Note
MK497057	-	<i>C. macrodisca</i>	Tung Wa, Satun, Thailand	-	This study	KUMF06872	
MK497058	-	<i>C. macrodisca</i>	Che Bilang, Satun, Thailand	-	This study	KUMF06874	
✓	✓	<i>C. serrulata</i>	Ko Rok, Krabi	-	This study	SP284	
✓	✓	<i>C. serrulata</i>	Ko Rok, Krabi	-	This study	SP301	
-	✓	<i>C. taxifolia</i>	Ko Rawi, Satun	-	This study	SP332	
✓	✓	<i>C. taxifolia</i>	Ko Rawi, Satun	-	This study	SP343	
✓	✓	<i>C. serrulata</i>	Ko Rok, Krabi	-	This study	SP358	
-	✓	<i>C. serrulata</i>	Ko Rok, Krabi	-	This study	SP368	
✓	-	<i>C. serrulata</i>	Ko Kradan, Trang	-	This study	SP372	
✓	✓	<i>C. taxifolia</i>	Ko Rawi, Satun	-	This study	SP474	
-	✓	<i>C. taxifolia</i>	Ko Rawi, Satun	-	This study	SP234	

* indicated as *tufA* barcode sequences (Belton *et al.*, 2014)

RESULTS AND DISCUSSION

The results and discussion are divided into two parts:

Part 1. Diversity and distribution of the genus *Caulerpa* in Thailand

Part 2. An enigmatic *Caulerpa macrodisca* Decaisne (Chlorophyta)
from the mangrove channels on the Andaman Sea coast of
Thailand.

Part 1

Diversity and distribution of the genus *Caulerpa* in Thailand

Diversity and distribution of the genus *Caulerpa* in Thailand

1. *Caulerpa* diversity in Thai waters

a. *Caulerpa* genetic diversity in Thailand

An alignment composed of 271 partial *tufA* sequences (767 bp in length) and 60 partial ITS rDNA sequences (1,114 bp in length) from GenBank with 108 newly generated *Caulerpa* sequences (45 *tufA* sequences and 63 ITS sequences) was analyzed using Maximum Likelihood (not shown) and Bayesian Inference (*tufA* in Fig. 4 and ITS sequence in Fig. 5). Both topologies confirmed a congruent result of eight *Caulerpa* species in Thai waters including *C. chemnitzia*, *C. lentillifera*, *C. macrodisca*, *C. racemosa*, *C. serrulata*, *C. sertularioides*, *C. taxifolia* and *C. verticillata* (Fig. 6). Each species clade had strong branch support (BS>80, PP>0.95).

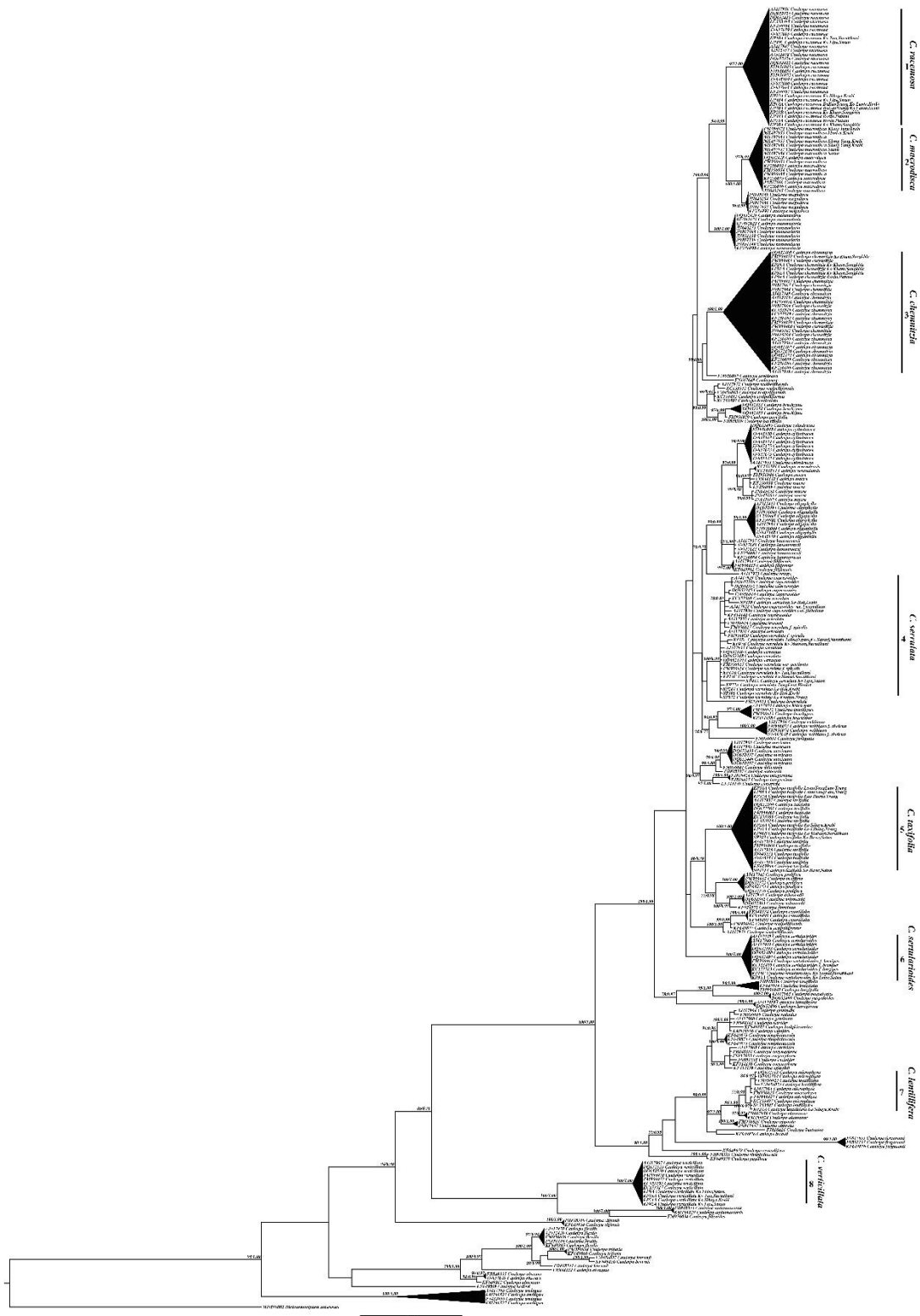


Figure 4 Bayesian phylogenetic tree constructed from a total of 316 partial *tufA* DNA sequences (alignment = 767 bp in length). Node Numbers indicate as BP and PP values. BP < 70% and PP < 0.7 are not shown. The scale is 0.04 expected changes per site.

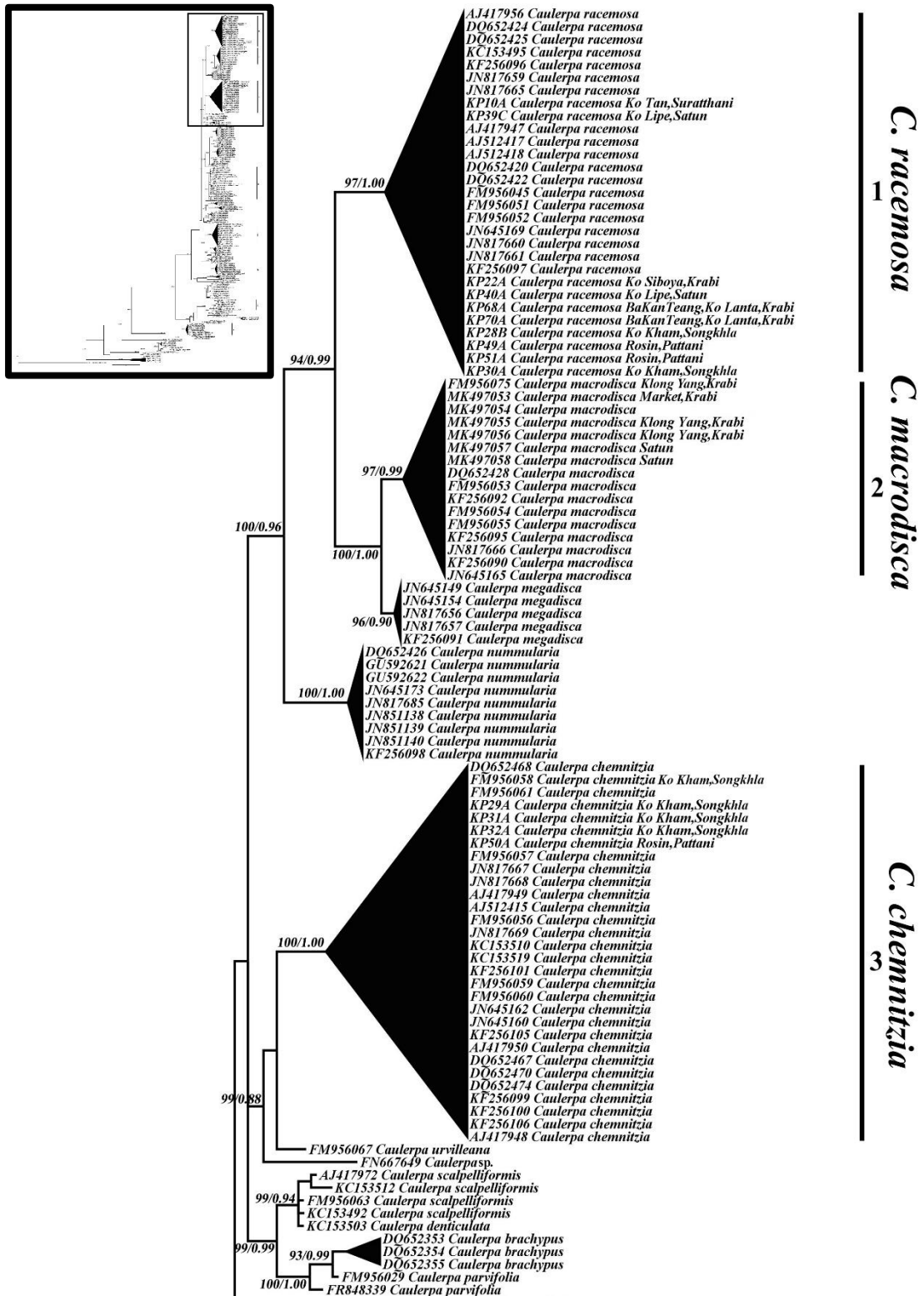


Figure 4 A., Bayesian *tufA* phylogenetic tree showed *Caulerpa* diversity from *C. racemosa* clade to *C. parvifolia*, of which there were 3 Thai *Caulerpa* clades, i.e., *C. racemosa*, *C. macrodisca* and *C. chemnitzia*. (continuous)

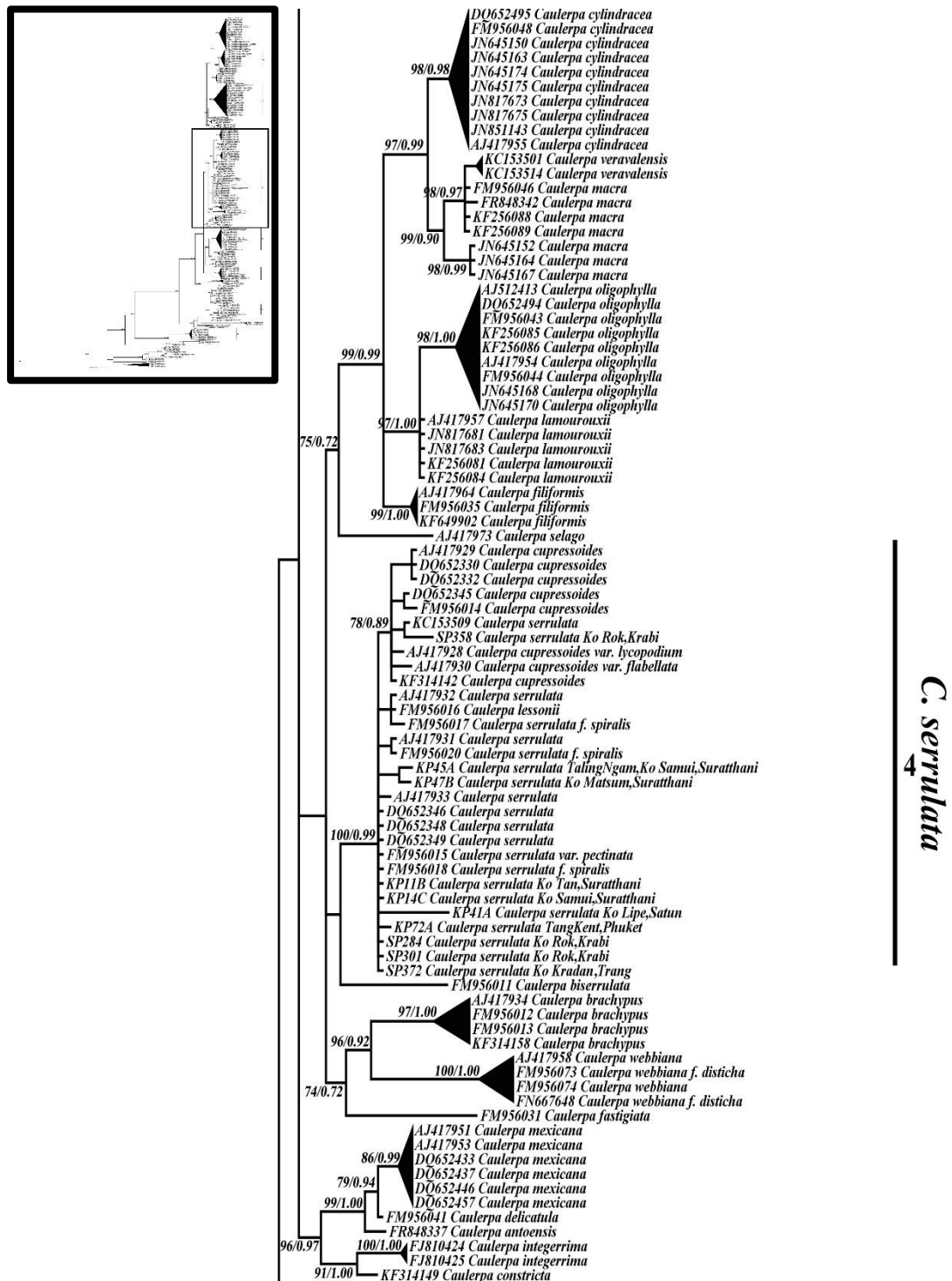


Figure 4 B., Bayesian *tufA* phylogenetic tree showed *Caulerpa* diversity from *C. cylindracea* clade to *C. constricta*, of which there was a Thai *Caulerpa* clades, i.e., *C. serrulata*. (continuous)

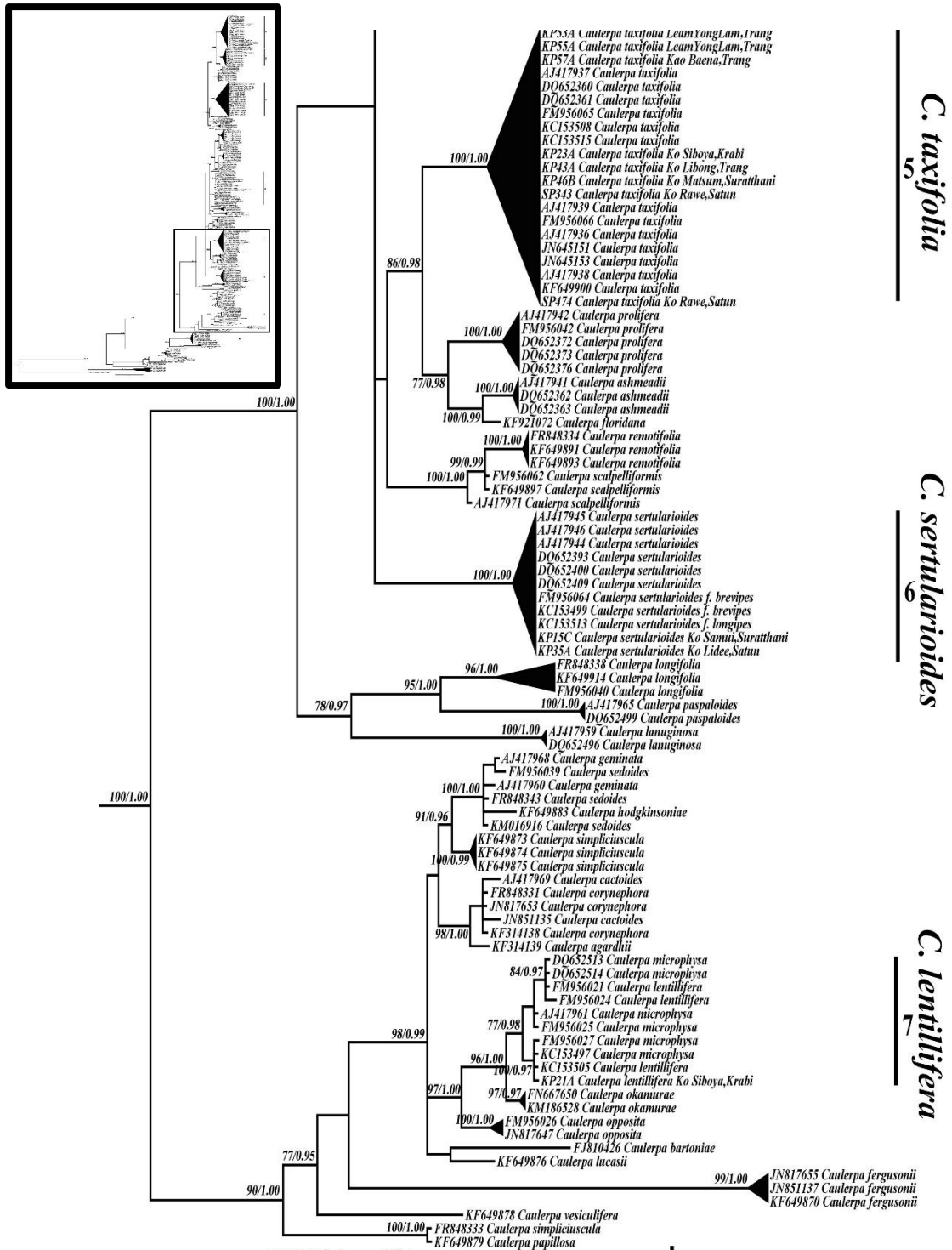


Figure 4 *C.*, Bayesian *tufA* phylogenetic tree showed *Caulerpa* diversity from *C. taxifolia* clade to *C. papillosa*, of which there were 3 Thai *Caulerpa* clades, i.e., *C. taxifolia*, *C. sertularioides* and *C. chemnitzia*. (continous)

C. verticillata

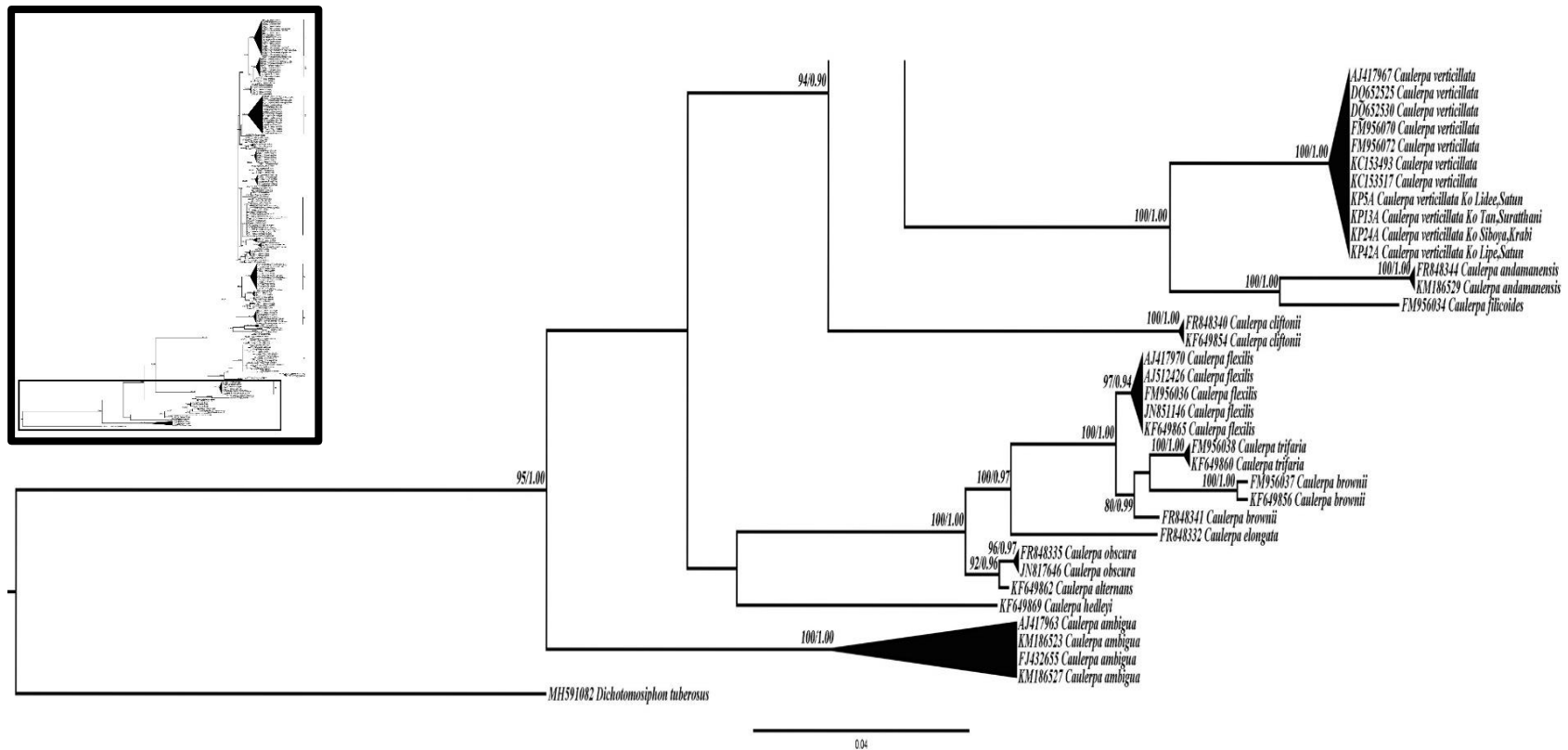


Figure 4 D., Bayesian *tufA* phylogenetic tree showed *Caulerpa* diversity from *C. verticillata* clade to *C. ambigua* clade with the outgroup (*D. tuberosus*), of which there was a Thai *Caulerpa* clades, i.e., *C. verticillata*. (continuous)

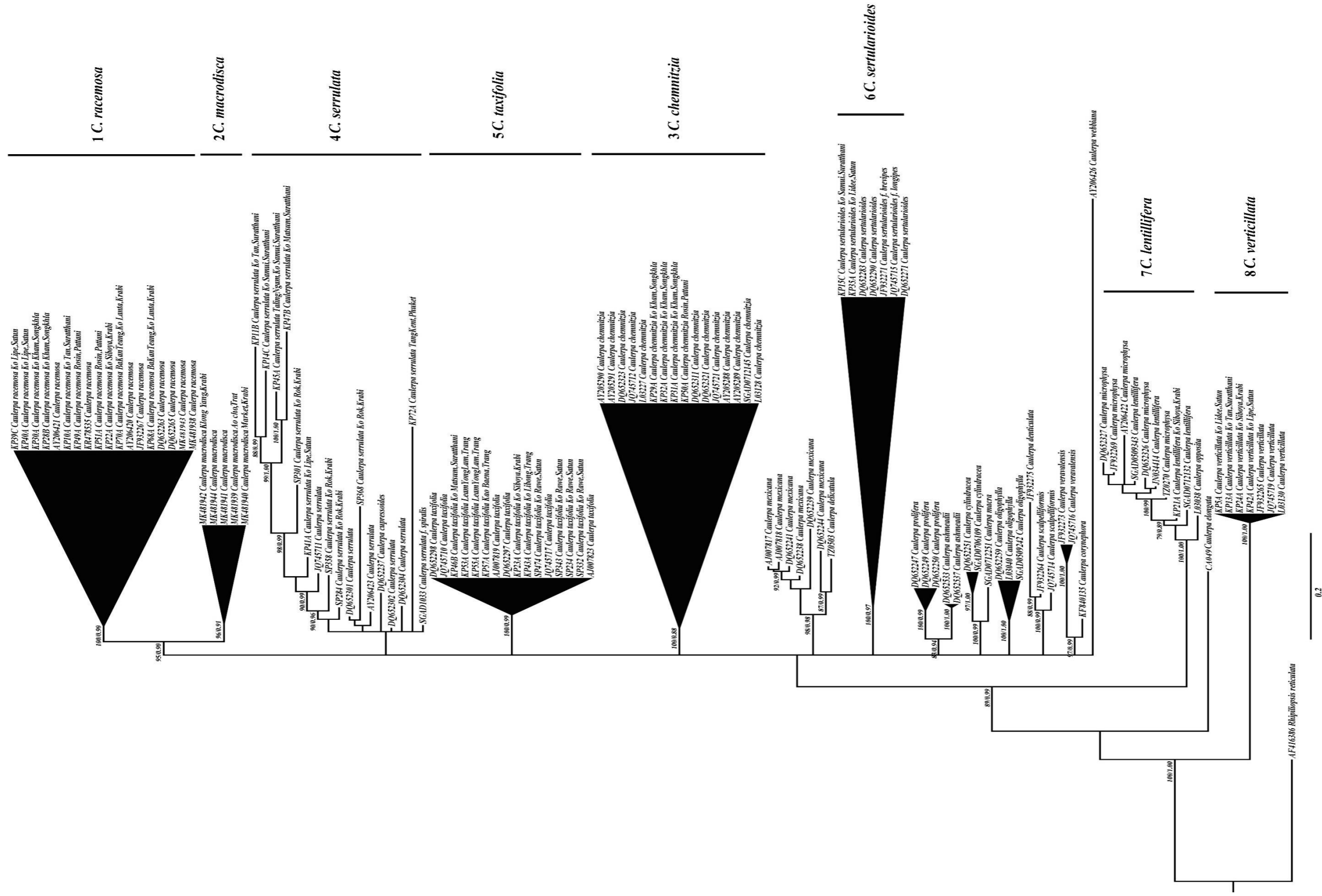


Figure 5 Bayesian phylogenetic tree constructed from a total of 123 complete ITS1, 2 sequences (alignment = 1,114 bp in length). Numbers at nodes indicate BP and PP values. BP < 70% and PP < 0.7 are not shown. The scale is 0.2 expected changes

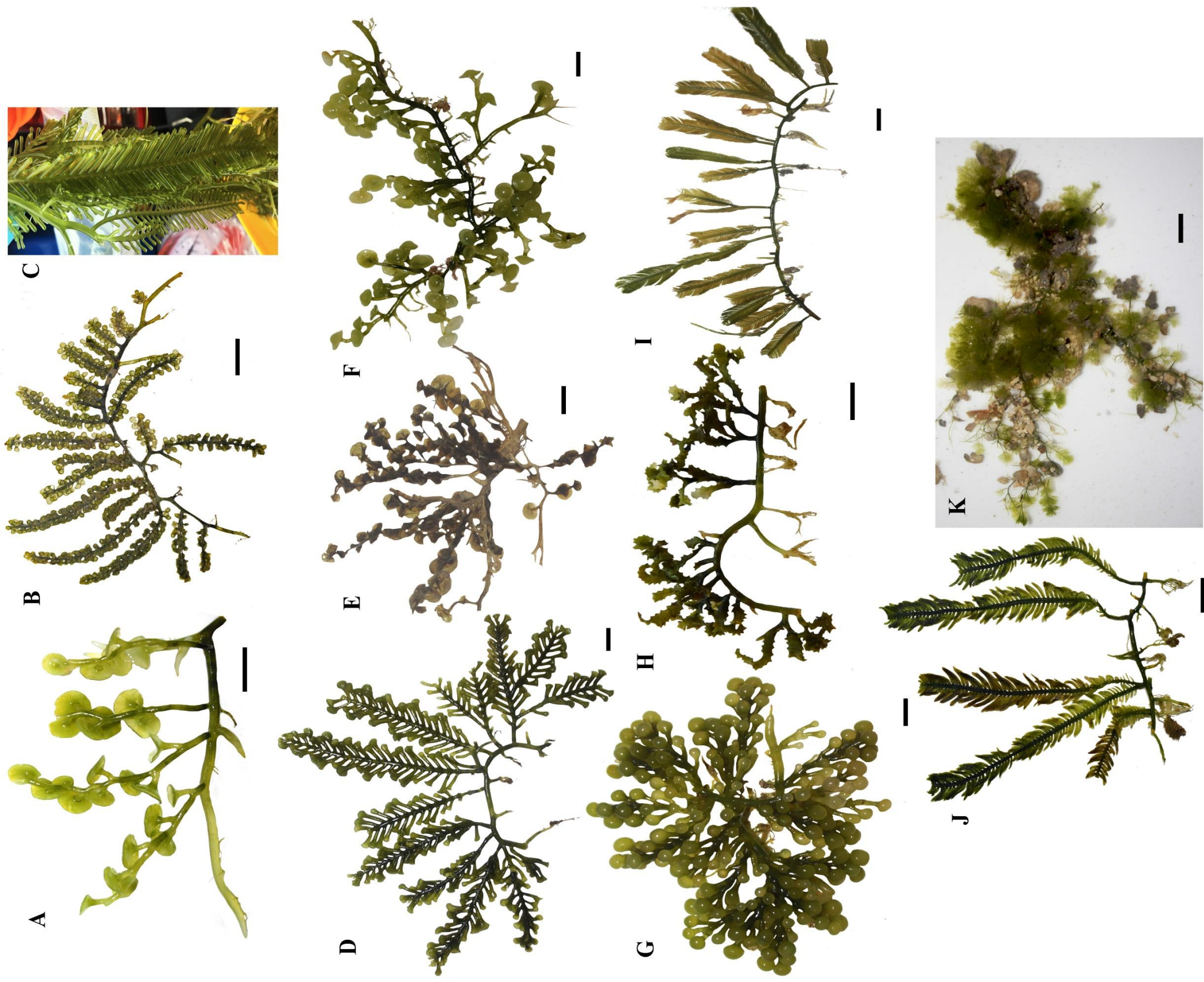


Figure 6 Eight *Caulerpa* species found in Thailand (Scale bar = 1 cm) A., *C. chemnitzia*, B., *C. lentilifera*, C-E., *C. macrodisca* (C., *C. macrodisca* ecad *ashmeadii*, no scale bar, D., *C. macrodisca* ecad *corynephora* and E., *C. macrodisca* ecad *macrodisca*), F-G., *C. racemosa* (F., *C. racemosa* ecad *chemnitzia* and G., *C. racemosa* ecad *racemosa*), H., *C. serrulata*, I., *C. sertularioides*, J., *C. taxifolia* and K., *C. verticillata*.

Table 4 Summary of morphological characteristics in the genus *Caulerpa* from Thailand (Min-Max \pm SE, cm), Measurements from dried specimens in square brackets “[]” n.m., not measured.

Features	<i>C. chemnitzia</i>	<i>C. lentillifera</i>	<i>C. macrodisca</i>				<i>C. racemosa</i>			<i>C. serrulata</i>	<i>C. sertularioides</i>	<i>C. taxifolia</i>	<i>C. verticillata</i>
			ecad <i>macrodisca</i>	ecad <i>corynephora</i>	ecad <i>ashmeadii</i>	ecad <i>racemosa</i>	ecad <i>chemnitzia</i>						
Diameter of stolon	0.17–0.34 \pm 0.009	0.13–0.18 \pm 0.006	0.15–0.18 \pm 0.003	0.23–0.34 \pm 0.006	n.m.	0.13–0.39 \pm 0.004	0.15–0.25 \pm 0.003	0.11–0.36 \pm 0.005	0.07–0.29 \pm 0.006	0.11–0.32 \pm 0.007	1.8–3.9 x 10 ⁻² \pm 0.001		
Rachis shape	Terete	Terete	Terete	Terete	Terete	Terete	Terete	Terete	Terete	Terete	Terete		
Assimilator shape	Irregularly	Irregularly	Irregularly	Distichous	Distichous	Irregularly	Irregularly	Flat, dichotomous branching	Distichous	Distichous	Filiform, dichotomous branching		
Assimilator margin	Entire	Entire	Entire	Entire	Entire	Entire	Entire	Serrate	Entire	Entire	Entire		
Assimilator length	0.87–6.75 \pm 0.280	1.8–5.7 \pm 0.441	[4.13–6.05 \pm 0.285]	2.89–10.48 \pm 0.369	n.m.	0.44–5.98 \pm 0.059	0.84–4.8 \pm 0.134	0.59–5.36 \pm 0.108	0.56–7.89 \pm 0.263	1.8–11.42 \pm 0.354	0.2–1.1 \pm 0.461		
Assimilator width	0.47–1.2 \pm 0.035	0.46–0.7 \pm 0.013	[0.54–1.23 \pm 0.112]	0.65–0.94 \pm 0.027	n.m.	0.29–1.88 \pm 0.019	0.33–1.39 \pm 0.031	0.22–4.13 \pm 0.066	0.21–1.56 \pm 0.051	0.13–1.46 \pm 0.033	0.09–0.53 \pm 0.187		
Arrangement of ramuli	Irregularly	Opposite decussate	Alternate, Irregularly	Opposite	Opposite, or alternate at base	Alternate, Irregularly	Irregularly	-	Opposite	Opposite	Whorled		
Length of ramuli	0.01–0.43 \pm 0.013	0.19–0.23 \pm 0.005	[\approx 0.01]	0.51–1.22 \pm 0.034	n.m.	0.2–0.67 \pm 0.008	0.01–0.47 \pm 0.009	0.36–4.67 \pm 0.097	0.17–0.127 \pm 0.035	0.26–0.77 \pm 0.016	0.12–0.34 x 10 ⁻² \pm 0.092		
Diameter of ramuli	0.11–0.78 \pm 0.016	0.21–0.23 \pm 0.002	[0.64–0.9 \pm 0.038]	0.14–0.5 \pm 0.021	n.m.	0.22–0.71 \pm 0.008	0.11–1.01 \pm 0.013	0.21–0.59 \pm 0.007	0.02–0.05 \pm 0.001	0.07–0.13 \pm 0.002	3–7 x 10 ⁻³ \pm 0.002		
Shape of ramuli	Mushroom-like, peltate and turbinate shape	Spherical shape	Large peltate shape	Clavate, turbinate, or rarely peltate shape	Cylindrical shape	Subspherical to spherical shape	Mushroom-like, peltate and turbinate shape	-	Cylindrical shape	Upright falcate shape	Filiform		
Shape of ramulus tip	-	-	-	-	Obtuse	Rounded	-	-	Acuminate	Acuminate	Three spines		
Shape of ramulus base	-	-	-	-	-	Cuneate	-	-	-	-	-		
Diameter of a constricted of ramuli	-	-	-	-	-	-	-	-	-	0.05–0.12 \pm 0.002	-		
Length of stalks	0.03–0.6 \pm 0.01	0.05–0.09 \pm 0.004	[0.11–0.25 \pm 0.023]	-	n.m.	0.06–0.47 \pm 0.006	0.03–0.43 \pm 0.006	-	-	-	-		
Diameter of stalks	0.05–0.23 \pm 0.005	0.06–0.07 \pm 0.001	[0.05–0.07 \pm 0.004]	-	n.m.	0.11–0.31 \pm 0.003	0.05–0.25 \pm 0.007	-	-	-	-		
Diameter of a constricted of stalks	-	0.04–0.05 \pm 0.001	-	-	n.m.	-	-	-	-	-	-		
Shape of tip	Acute	Rounded	Disc	Acute	Acute	Rounded	Acute	Teeth	Mucronate	Mucronate	Mucronate		

b. *Caulerpa* morphological diversity in Thailand

For morphological study, five species including *C. lentillifera*, *C. serrulata*, *C. sertularioides*, *C. taxifolia*, and *C. verticillata* showed clear morphological criteria for specific identification based on ramuli arrangement and shapes. Other three species including *C. racemosa*, *C. macrodisca* and *C. chemnitzia* were in the *C. racemosa - peltata* complex presenting considerable levels of morphological plasticity. Thus, another tool, *i.e.*, DNA marker, was important key for their identifications. In this study, there were two morphotypes of *C. racemosa* named here *C. racemosa* ecad *racemosa* and *C. racemosa* ecad *chemnitzia*, while *C. macrodisca* was divided into three morphologies named *C. macrodisca* ecad *macrodisca*, *C. macrodisca* ecad *corynephora* and *C. macrodisca* ecad *ashmeadii*. Of which, *C. racemosa* ecad *chemnitzia* and *C. chemnitzia* were indistinguishable because of their overlapping morphology. So, molecular data was a necessary tool for their specific identification and classification. The summary of morphological characteristics is described in Table 4.

Lineage 1 (Figs 4A, 5) represents *Caulerpa racemosa* (Forsskål) J. Agardh and does not only include specimens with spherical ramuli identified as *C. racemosa* (KP10 from Surat Thani, the Gulf of Thailand, KP39, KP40 from Satun, KP22, KP68 and KP70 from Krabi, the Andaman Sea), but also peltate specimens from the lower Gulf of Thailand morphologically attributed to *C. chemnitzia* (KP28A, KP30A from Koh Kham, Songkhla and KP49A, KP51A from Rosin Pattani). There was obviously morphological difference. Peltate specimens (as *C. chemnitzia* morphology) normally showed considerable morphological varieties of ramuli from mostly turbinate to peltate or sometimes mushroom-like, while spherical specimens (as *C. racemosa* morphology) presented erect assimilators bearing densely globose to subspherical ramuli. The spherical specimens previously morphologically identified as *C. racemosa* var. *macrophysa* (see Lewmanomont, 2008 fig. 6). However, there were extremely morphological variable of *C. racemosa* (Coppejans and Beeckman, 1989) led to consider *C. racemosa* var. *macrophysa* as belonging in this species (Coppejans, 1992). For molecular data, our results confirmed a monophyletic group of *C. racemosa*. Thus, these spherical specimens were currently regarded as *C. racemosa* ecad *racemosa*, while peltate specimens were referred as *C. racemosa* ecad *chemnitzia*.

Lineage 2 (Figs 4A, 5) represents *Caulerpa macrodisca* Decaisne containing previous specimens morphologically identified as *C. peltata* var. *macrodisca* (KUMF04404, from intertidal shore of Trat, the Gulf of Thailand), *C. corynephora* (as *C. racemosa* var. *corynephora*, *sensu* Lewmanomont, 2008, figs 11, 12) (KP4 from fresh market in Krabi, KP73, KP74 from Khlong Yang, Krabi and KUMF06874 from mangrove channel in Satun, the Andaman Sea coast) and *C. ashmeadii* (KUMF06872, *sensu* Lewmanomont, 2008, from mangrove channel in Satun, the Andaman Sea) together. The congruent *tufA* and ITS phylogenies strongly suggested that the three morphological entities in Thai waters represented a single species and will be onwards referred to as *C. macrodisca* ecad *ashmeadii*, *C. macrodisca* ecad *corynephora* and *C. macrodisca* ecad *macrodisca*. This clade will be discussed in more detail in the second part “An enigmatic *Caulerpa macrodisca* Decaisne (Chlorophyta) from the mangrove channels on the Andaman Sea coast of Thailand”.

Lineage 3 (Figs 4A, 5) represents *Caulerpa chemnitzia* (Esper) J.V.Lamouroux and includes specimens previously identified as *C. peltata* (*sensu* Lewmanomont, 2008, fig. 10). *Caulerpa chemnitzia* specimens (KP29, KP31, KP32 from Ko Kham, Songkhla and KP50 from Losin, Pattani in the lower Gulf of Thailand) showing the morphology of peltate form. Erect assimilators bearing considerable morphological varieties of ramuli from mostly turbinate to peltate shape, that disc diameter was overlapping with *C. racemosa* ecad *chemnitzia*, but smaller than *C. macrodisca* ecad *macrodisca* (Belton *et al.*, 2014; Price, 2011). Both phylogenies illustrated monophyletic clade of *C. chemnitzia*, *C. macrodisca* and *C. racemosa* respectively with strong supported values.

Lineage 4 (Figs 4B, 5) represents *Caulerpa serrulata* (Forsskål) J.Agardh from both coastlines (*sensu* Lewmanomont, 2008, fig. 14). *Caulerpa serrulata* specimens (KP11, KP14, KP45, KP47 from Surat Thani, the Gulf of Thailand, KP41 from Satun, KP72 from Phuket, SP372 from Trang and SP358, SP284 and SP301 from Krabi, the Andaman Sea) presented morphology of flattened twisted branches with serrate margin and dichotomous branching. *TufA* and ITS topologies showed a mixed of *C. serrulata* and *C. cupressoides* together as previous studies (Famà *et al.*, 2002; Stam *et al.*, 2006; Draisma *et al.*, 2014). However, *C. cupressoides* was separated from *C. serrulata* by the *rbcL* gene (Kazi *et al.*, 2013), and presented different morphology of flattened

serrate branches with upcurved pointed tip. Thus, an apparent morphological distinct was able to separate into two species (de Senerpont Domis *et al.*, 2003).

Lineage 5 (Figs 4C, 5) represents *Caulerpa taxifolia* (M.Vahl) C.Agardh growing on sandy bottom substrate from both seashores. *Caulerpa taxifolia* specimens (KP46 from Surat Thani, the Gulf of Thailand, SP234, SP332, SP343 and SP474 from Satun, KP23 from Krabi and KP43, KP53, KP55, and KP57 from Trang, the Andaman Sea coasts) (sensu Lewmanomont, 2008, fig. 16) showed a feather-like assimilator with flattened upright sickle-shaped (falcate) ramuli and opposite arrangement. Its morphological characters and molecular phylogeny result presented congruent results of a species *C. taxifolia*.

Lineage 6 (Figs 4C, 5) represents *Caulerpa sertularioides* (S.G.Gmelin) M.Howe from both seacoasts. *Caulerpa sertularioides* specimens (KP15 from Surat Thani, the Gulf of Thailand, KP35 from Satun, the Andaman Sea coasts) (sensu Lewmanomont, 2008, fig. 15) showed a feather-like assimilator bearing slender cylindrical pinnules oppositely arranged. This morphological distinct data was supported their molecular phylogenies suggested that referred as *C. sertularioides*.

Lineage 7 (Figs 4C, 5) represents *Caulerpa lentillifera* J.Agardh (KP21A) from Ko Siboya, Krabi (sensu Lewmanomont, 2008, fig.5). Four rows of spherical translucent ramuli constricted at the base appeared on each erect assimilator. For *tufA* and ITS topology, this lineage was mixed with *C. microphysa* similar to a previous result of Kazi *et al.* (2013). Moreover, both species were not difference in *rbcL* gene and 18S rDNA sequence (Kazi *et al.*, 2013). So, this study might be considered that both taxa was belong to a species.

Lineage 8 (Figs 4D, 5) represents *Caulerpa verticillata* J.Agardh from both Thai coasts. Specimens (KP13 from Surat Thani, the Gulf of Thailand, KP24 from Krabi, KP5 and KP42 from Satun, the Andaman Sea coasts) (sensu Lewmanomont, 2008, fig. 17) presented filiform thallus with dichotomous branching of whorled branched. A strongly supported of a monophyletic clade of *C. verticillata* also indicated as a single species, *C. verticillata*.

2. *Caulerpa* distribution in Thai waters

Eight species were found on the Andaman coast, *i.e.*, *C. chemnitzia*, *C. lentillifera*, *C. macrodisca* ecad *ashmeadii*, *C. macrodisca* ecad *corynephora*, *C. racemosa* ecad *racemosa*, *C. serrulata*, *C. sertularioides*, *C. taxifolia* and *C. verticillata*. Seven species were found in the Gulf of Thailand, *i.e.*, *C. chemnitzia*, *C. macrodisca* ecad *macrodisca*, *C. racemosa* ecad *chemnitzia*, *C. racemosa* ecad *racemosa*, *C. serrulata*, *C. sertularioides*, *C. taxifolia* and *C. verticillata* (Table 5).

Table 5 List of *Caulerpa* species were found from both Thai waters.

<i>Caulerpa</i> species	The Andaman Sea	The Gulf of Thailand
1. <i>C. chemnitzia</i>	✓	✓
2. <i>C. lentillifera</i>	✓	
3. <i>C. macrodisca</i>		
<i>C. macrodisca</i> ecad <i>ashmeadii</i>	✓	
<i>C. macrodisca</i> ecad <i>corynephora</i>	✓	
<i>C. macrodisca</i> ecad <i>macrodisca</i>		✓
4. <i>C. racemosa</i>		
<i>C. racemosa</i> ecad <i>chemnitzia</i>		✓
<i>C. racemosa</i> ecad <i>racemosa</i>	✓	✓
5. <i>C. serrulata</i>	✓	✓
6. <i>C. sertularioides</i>	✓	✓
7. <i>C. taxifolia</i>	✓	✓
8. <i>C. verticillata</i>	✓	✓

During the field surveys, *C. macrodisca* ecad *corynephora* and *C. macrodisca* ecad *ashmeadii* only were found in the Andaman mangrove channels. *Caulerpa lentillifera* was only found in Krabi province (the Andaman Sea coast), but there was previously reported of *C. lentillifera* from the upper Gulf of Thailand by Lewmanomont (2008). On the other hand, *C. macrodisca* ecad *macrodisca* was found in the upper Gulf of Thailand and *C. racemosa* ecad *chemnitzia* was found in the lower Gulf of Thailand (Ko Kham, Songkhla and Losin, Pattani) growing on the rock together with *C. chemnitzia*.

Sixteen *Caulerpa* species have been recorded from Thai waters (Lewmanomont, 2008; Coppejans *et al.*, 2017), but only 8 species were confirmed by molecular data in the present study. A summary of *Caulerpa* distribution in Thai waters is shown in Figure 7. This map shows the highest *Caulerpa* diversity in Krabi, Satun and Trang provinces in the upper Strait of Malacca (the Andaman Sea coast) followed by Trat and Surat Thani provinces (the upper and middle Gulf of Thailand). Seven previously recorded morphological species *C. ambigua*, *C. ashmeadii*, *C. cupressoides*, *C. fastigiata*, *C. manorensis*, *C. mexicana* and *C. microphysa* sensu Lewmanomont (2008) and Coppejans *et al.* (2017) were not found in this present study. Those recorded species were not amplified both sequences from their extracted herbarium specimens to confirm their identities. However, they have clearly their own morphological criteria leading to accept those identified species except *C. ashmeadii*, which will be discussed more in part 2 of this thesis.

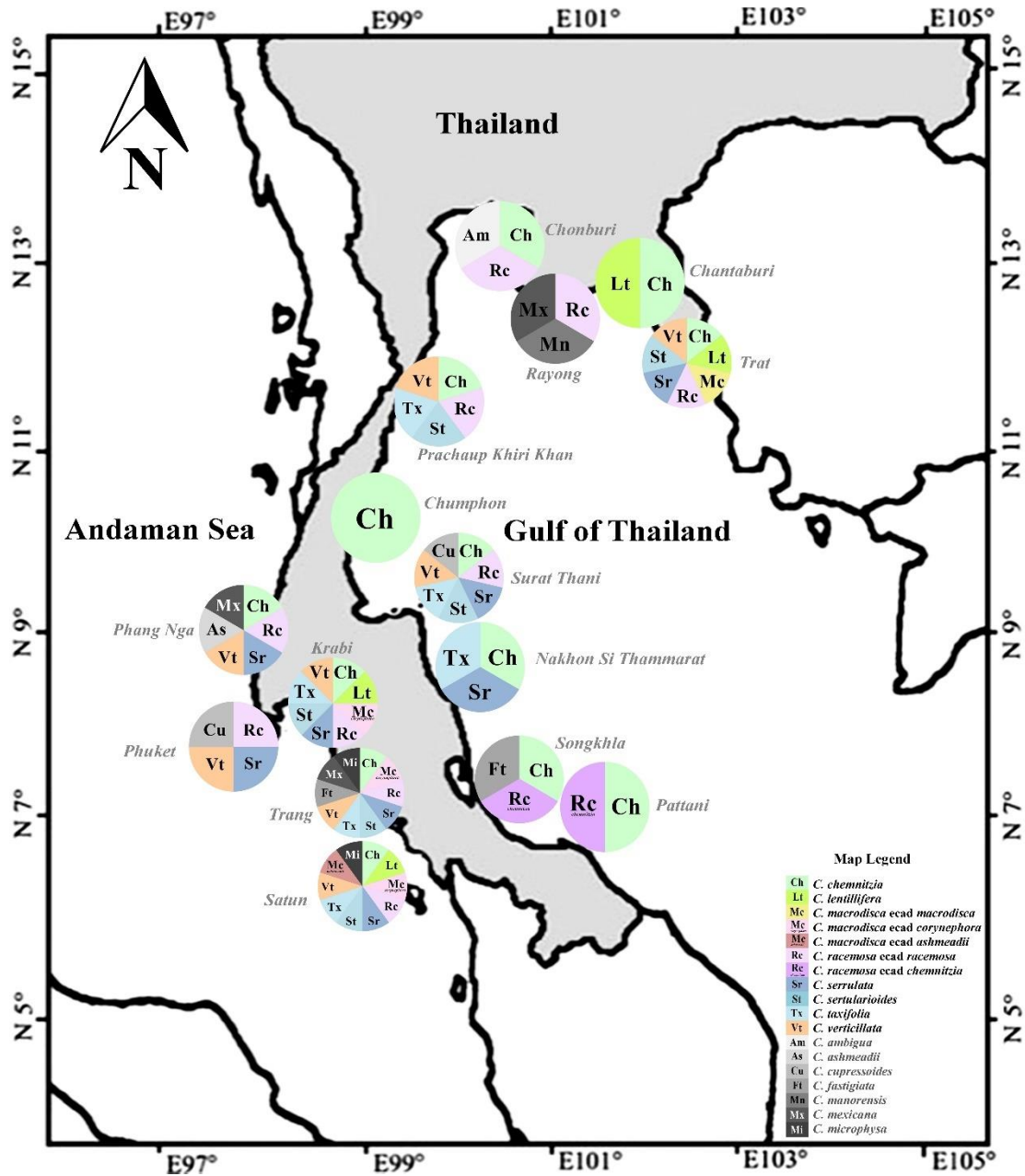


Figure 7 Map showing the distribution of *Caulerpa* species in Thailand Data from herbarium specimens in NHM, Coppejans *et al.* (2017), Lewmanomont (2008), and the present. Color legend: Thai *Caulerpa* species collected and reported in the present study, Dark-gray legend: species not found in the present study but found in the NHM herbarium or reported in the literature

Discussion

1. *Caulerpa* diversity in Thai waters

This study showed a taxonomic revision of the genus *Caulerpa* in Thailand at the species level. The congruent molecular identifications using *tufA* gene and ITS sequence in combination with morphological data of recent collections in Thailand reported the occurrence of eight *Caulerpa* species in Thai waters; *C. chemnitzia*, *C. lentillifera*, *C. macrodisca* (ecad *ashmeadii*, ecad *corynephora* and ecad *macrodisca*), *C. racemosa* (ecad *chemnitzia* and ecad *racemosa*), *C. serrulata*, *C. sertularioides*, *C. taxifolia* and *C. verticillata*. Other previously reported species, i.e., *C. ambigua*, *C. cupressoides*, *C. fastigiata*, *C. manorensis*, *C. mexicana* and *C. microphysa* sensu Lewmanomont (2008) were not found during this survey and unconfirmed their identities using molecular evidence. Thus, this study only found half of the *Caulerpa* diversity in Thailand reported in Lewmanomont (2008). That might be because of insufficient sampling, monitoring time and seasonal variations, which highly influenced on the diversity and distribution of any marine macroalgae (Prathep *et al.*, 2005; Thongroy *et al.*, 2007). Lewmanomont (2008) has been worked on her specimen collections for a decade spending more times to revisit all study sites in both monsoons, while this study ran from 2017–2019. Thus, it might be better for diversity research to extend more monitoring times and data collections.

In this study, there are eight *Caulerpa* species in Thailand. Of which, five taxa; *C. lentillifera*, *C. serrulata*, *C. sertularioides*, *C. taxifolia* and *C. verticillata* were clearly identified based on morphological criterion following Lewmanomont (2008) and Coppejans *et al.* (2010, 2017) supporting by both molecular data (Famà *et al.*, 2002; Belton *et al.*, 2014; Draisma *et al.*, 2014). However, others; *C. chemnitzia*, *C. racemosa* and *C. macrodisca* were later defined to the correct species based on molecular results.

Thai *Caulerpa* specimens previously identified as *C. corynephora* (as *C. racemosa* var. *corynephora*) sensu Lewmanomont *et al.* (2008) and Coppejans *et al.* (2017) are a variety of *C. macrodisca*. It illustrates a terete rachis with diverse ramuli from mainly clavate, turbinate to peltate formed without a constriction at the basis that not similar to

the type description by Montagne (1845) and the lectotype illustration of *C. corynephora* in Price (2011). Price (2011) reported that the true *C. corynephora* was limited in Australia presenting an annulated rachis forming transverse articulations with mostly opposite clavate ramuli arrangement. Those obvious dissimilar morphologies were supported by a phylogeny of this study indicating in a monophyletic clade of *C. macrodisca* (section *Caulerpa* crown clade) with strong support values (Fig. 4). Thus, all previous *C. corynephora* specimens in Thai were consequently renamed as *C. macrodisca* ecad *corynephora*. In addition, a confused species, *C. macrodisca* ecad *ashmeadii* (morphologically identified as *C. ashmeadii* sensu Lewmanomont (2008)) showed terete rachis consisted with cylindrical ramuli (slightly upcurved at base) with swollen tip in opposite arrangement (sometimes alternate arrangement at base), which not matched with the type illustration of *C. ashmeadii* by Harvey (1858). The type morphology of *C. ashmeadii* presented erect opposite straight cylindrical ramuli arrangement with obtuse tip. Moreover, *C. ashmeadii* is a tropical native species distributed in the Atlantic Ocean (Zaleski and Murray, 2006). Thus, the newly collected specimens previously identified as *C. ashmeadii* demonstrated as *C. macrodisca* ecad *ashmeadii* using *tufA* phylogeny in combination with morphological data. Other informations is explained in the second part.

Sympatric growing specimens (in Songkhla and Pattani) previously identified as *C. chemnitzia* based on original peltate ramuli morphology, represented 2 different species; *C. chemnitzia* and *C. racemosa*. In this study, the comparison demonstrates the biggest degrees of morphological plasticity in the complex particularly on assimilator length, ramuli length and diameter. Their qualitative characters are similar, while quantitative features are overlapping. Accordingly, PCA and ANOVA results (not shown) showed no significant difference on any character of their morphometric analysis. This corroborates the unreliability of morphological identification in the former *C. racemosa-peltata* complex (Belton *et al.*, 2014). Another study by de Senerpont Domis *et al.* (2003) showed congruous result that it was not morphological distinct exhibited on *C. racemosa* complex because its morphological plasticity responding environmental factors. Thus, the *Caulerpa racemosa-peltata* complex cannot be discriminated from a difference in morphology evidently.

Previously, there were many physiological studies about morphological plasticity of *Caulerpa*. The interrelated contribution of temperature and light intensity have important effect on external morphological constructs (Calvert, 1976; Ohba *et al.*, 1992; Komatsu *et al.*, 1997). Peterson (1972) found that *C. racemosa* (as *Caulerpa racemosa* var. *uvifera* (C.Agardh) J.Agardh) and *C. lamourouxii* (as *Caulerpa racemosa* var. *lamourouxii* (Turner) Weber-van Bosse) had diverse morphological growth under several illuminate situations that might help balancing their photosynthesis and respiration processes. In addition, Ohba and Enomoto (1987) found a high degree of morphological variation of *C. chemnitzia* (as *Caulerpa racemosa* var. *laetevirens*) that formed laetevirens-type (cylindrical ramuli) in the low temperatures-high light intensities conditions, and formed peltata-type (discoid shape ramuli) in various temperatures-low light intensities condition and intermediated type (trumpet shaped ramuli) in others. Hence, environmental variation can affect the external morphology that was causes of taxonomy problem in this genus.

Nevertheless, Yeh and Chen (2004) who studied various varieties of *C. racemosa* confirmed that nuclear rDNA and ITS sequences are helpful resolution of the inter- and infraspecific identification. The result is consistent with the study of Ohba and Enomoto (1987) showing that *C. racemosa* var. *laetevirens*, var. *peltata* and var. *turbinata* are varieties of a single species. Therefore, molecular tool is an important key to confirm specific identification of the complex. Hence, our coherent phylogenetic results apparently indicate sufficient authority for specific delimitation of the green algal genus *Caulerpa* using *tufA* gene combination with ITS region. This present study suggested that both DNA markers are appropriate makers for molecular identifications of the genus, but the reference database for *tufA* sequences is much larger. So, the *tufA* gene also provided obviously interspecific identification, while ITS rDNA sequences were still needed for subspecies level as a supporting data (Stam *et al.*, 2006; Kazi *et al.*, 2013; Draisma *et al.*, 2014).

2. *Caulerpa* distribution in Thai waters

This study showed that species compositions among sites were different. This might be a result of physical factors such as habitat types, environment factors, an indirect effect of sea surface currents (SSCs).

The biology of *Caulerpa* showed that a strong creeping stolon with numerous long rhizoids usefully attaches on various substrate in different habitat types, such as dead coral, rocky shore, muddy bottom and sandy bottom (Lee, 2008). For example, *C. taxifolia* and *C. sertularioides* normally grow on the sandy bottom in the inter- to subtidal zone. Four *Caulerpa* species; *C. racemosa*, *C. lentillifera*, *C. serrulata* and *C. chemnitzia* were generally found on hard substrates, *i.e.*, dead coral, coral rubble or rock in tide pool of rocky shore or coral reef habitat. *Caulerpa verticillata* can grow on both hard substrate and sand in the intertidal zone. Only *C. macrodisca* was found in the mangrove channels growing on floating fish cages or muddy bottom, and some specimens could attach on rock in the intertidal zone (Coppejans *et al.*, 2017; Lewmanomont, 2008). In Thailand, environment factors of both Thai coasts were different in each location (see in Table 1). Each organism requires a niche such as habitat type and physical environment for its suitable growth and reproduction (Begon *et al.*, 2006), that might mainly influence on species composition in each study site.

Pongparadon *et al.* (2015) also found higher diversity of the green algal genus *Halimeda* from the Andaman Sea coast than the Gulf of Thailand similar to another studies, *i.e.*, Wichachucherd *et al.* (2014) and Pongparadon *et al.* (2017). The Gulf of Thailand was influenced by the mix of the Philippines Sea current and the East China Sea current in the Northeast monsoon providing the clockwise current direction in the Thai Gulf. The circulated currents directly affected connectivity reduction with open water that possible limited the species distribution within the Gulf of Thailand. For the Andaman Sea coast, not only the current from Indian Ocean water masses flow through the Thai Andaman Sea coast in both monsoons, but also a current from the South China in the Northeast monsoon possibly provide more opportunity of diverse species for settlement. Thus, the water mass might be driving forces of various marine organism appearances (Chen, 1999; Huang, 1999; Hu *et al.*, 2011). Thus, the differences in SSCs

could powerfully indirectly affect genetic diversity and distribution patterns of *Caulerpa* along the Thai coastline.

In the Southeast Asia, eight species, *C. chemnitzia*, *C. lentillifera*, *C. macrodisca*, *C. racemosa*, *C. serrulata*, *C. sertularioides*, *C. taxifolia* and *C. verticillata* were generally distributed as native species in the Coral Triangle (Prud'homme van Reine *et al.*, 1996). Then, they were common dispersal along both coasts of Thailand connected to other countries in Southeast Asia region and Australia (Zaleski and Murray, 2006; Guiry and Guiry, 2019) except *C. lentillifera*, which was only found on the Andaman coast.

The present study is the first report of *C. lentillifera* from natural habitat in Krabi (the Andaman Sea coast). However, Lewmanomont (2008) recorded *C. lentillifera* in the Eastern Gulf of Thailand (Ko Chik, Chantaburi and Ko Kradat, Trat) for ten year ago. Ratana-arporn and Chirapart (2006) also reported the cultivate *C. lentillifera* from coastal shrimp aquacultures in Phetchaburi province (the Gulf of Thailand), and from Chachoengsao province by Chirapart *et al.* (2011). This species is becoming as a commercial algal species, which generally cultivated *C. lentillifera* from the Gulf of Thailand in each fishery agriculture along both sides of Thailand. *Caulerpa* was rapid growth under the culture conditions without any attentive maintenance. Thus, that might be some fragment of culture *C. lentillifera* was unexpectedly released to the Andaman Sea, and grow in the new locations here.

Although, *C. macrodisca* ecad *corynephora* and *C. macrodisca* ecad *ashmeadii* was limited in brackish water (27–30 ppt) of the Andaman Sea, while a larger peltate ramuli; *C. macrodisca* ecad *macrodisca* was restricted in the Gulf of Thailand. More information about this species has been discussed in part 2 confirmed by molecular and morphological data.

In addition, several *Caulerpa* species previously recorded in Thailand; *C. cupressoides*, *C. manorensis* and *C. mexicana* were not found during our field survey. It might be because of their rareness. *Caulerpa ambigua* and *C. fastigiata* were not found either and may have been overlooked due to their small size. Moreover, their identities could not be confirmed with DNA sequence data obtained from herbarium

specimens. However, their morphological evidences seem obvious enough to confidently predict specific identification.

Part 2

**An enigmatic *Caulerpa macrodisca* Decaisne (Chlorophyta)
from the mangrove channels on the Andaman Sea coast of
Thailand**

An enigmatic *Caulerpa macrodisca* Decaisne (Chlorophyta) from the mangrove channels on the Andaman Sea coast of Thailand

(A submitted manuscript of Journal of Fisheries and Environment, JFE)

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Introduction

The common tropically to subtropically distributed genus *Caulerpa* J.V.Lamouroux is a coenocytic siphonous green alga. Plants consist of horizontally growing stolons with downward growing rhizoids and upright branched or unbranched fronds (assimilators). Lateral branchlets (ramuli) on the assimilators can be of various shapes e.g., cylindrical (terete), trumpet-shaped (turbinate), club-shaped (clavate), sickle-shaped (falcate), disc-shaped (peltate) or globular (vesiculate). The shape and arrangement of ramuli are important keys in species identification in this genus (Coppejans and Beeckman, 1989). However, morphological plasticity is known in the genus, induced by environmental factors like light intensity (Calvert, 1976) and temperature (Ohba *et al.*, 1992). Moreover, some species have overlapping morphologies (Draisma *et al.*, 2014; Belton *et al.*, 2014, 2019). Correct species identification is therefore a challenge. The morphological plasticity within this genus has resulted in a long-standing source of uncertain and unstable taxonomy (Famà *et al.*, 2002; Sauvage *et al.*, 2013; Belton *et al.*, 2014).

It was long debated whether several species with vesiculate and peltate ramuli represent different species or varieties of a single or a few species. These taxa have been referred to as the “*Caulerpa racemosa-peltata* complex”. Belton *et al.* (2014) recognized 11 distinct species in the complex based on chloroplast-encoded *tufA* gene and RUBISCO large subunit (*rbcL*) gene sequence data. They provided names, DNA barcodes (*i.e.*, reference *tufA* sequences), and morphological descriptions for nine species, but stated that reliable morphological characterization remains not possible for several species due to high levels of phenotypic plasticity and morphological overlap.

They strongly suggested the use of molecular-based identifications and to refrain from recognizing any infra-specific ranks (*i.e.*, varieties and forms) within these species, but instead the use of morphological entities without formal taxonomic status (*e.g.*, ecad, ecotype) for highly plastic species.

Belton *et al.* (2014) recognized *Caulerpa macrodisca* Decaisne (homotypic synonyms: *Caulerpa racemosa* var. *macrodisca* (Decaisne) Weber Bosse 1898 and *Caulerpa peltata* var. *macrodisca* (Decaisne) Weber Bosse 1898) as sister-species of a newly described *C. megadisca* Belton & Gurgel. They described *C. macrodisca* as having large peltate ramuli arranged around an upright axis, and noted morphological variations of ramuli from disc-like to slightly mushroom-like. The specimens molecularly identified as *C. macrodisca* in Belton *et al.* (2014) came from Indonesia (7 specimens), Thailand (1), Australia (2), New Caledonia (1), and the aquarium trade (1), of which some were previously published (Stam *et al.*, 2006; Sauvage *et al.*, 2013). Three specimens from the Thousand Islands in the Java Sea were collected nearest to the type location of *C. macrodisca* (Anambas Islands, Indonesia) and from one of these three the *tufA* sequence was selected as reference sequence for the species. The DNA sequence of the Thai specimen was originally published in Sauvage *et al.* (2013), but its morphology was not discussed and neither in Belton *et al.* (2014). However, it was submitted to Genbank as *C. racemosa* var. *corynephora* (Montagne) Weber-van Bosse, the currently accepted name of which is *C. corynephora* Montagne (Guiry and Guiry, 2019) and its morphology deviates from the description of *C. macrodisca sensu* Belton *et al.* (2014). Moreover, molecular studies showed that the two species are not closely related, *C. corynephora* belonging to the *Caulerpa* section *Sedoideae* J.Agardh ex De Toni and *C. macrodisca* to the section *Caulerpa* (Belton *et al.*, 2015, 2019).

In Thailand, *C. corynephora* has only been reported from the Andaman Sea coast growing in mangrove channels, often on floating fish cages (Lewmanomont 1978, 2008; Coppejans *et al.*, 2017). In Thailand, *Caulerpa macrodisca* with characteristic peltate ramuli has only been reported from the Gulf of Thailand (Lewmanomont, 2008, as *C. peltata* var. *macrodisca*). Its identity was not confirmed with DNA sequence data. The aims of the present study are to confirm the identity of *C. macrodisca* in the Gulf of Thailand and the morphological *Caulerpa* entity previously reported as *C. corynephora*

(or *C. racemosa* var. *corynephora*) from mangrove channels on the Andaman Sea coast using DNA sequence data and to describe their morphological variation.

Materials and methodology

Specimen collection and morphological characterization

The intertidal and subtidal Andaman Sea coast and the Gulf of Thailand were explored by snorkeling and SCUBA diving from 2016–2018. Six *Caulerpa* specimens used in the present study were newly collected from mangrove channels on the Thai Andaman Sea coast and Langkawi, Malaysia, and from a market in Krabi province, Thailand (Table 3). Freshly collected specimens were stripped of epiphytes, photographed using a CANON 60D camera, and various morphological characters such as stolon diameter, assimilator length and ramulus diameter were measured using a caliper. Once dried, specimens were measured using NIH ImageJ software (Rasband, 1997). A small piece (1–2 cm) was preserved in silica gel for later DNA extraction. Specimens were herbarium pressed and labeled following Coppejans *et al.* (2010). Subsequently, specimens were identified following key references (Lewmanomont, 2008; Coppejans *et al.*, 2017). Attempts to make new collections of *C. macrodisca* with peltate ramuli in Thailand were unsuccessful. Therefore, we attempted to determine DNA sequence data from a herbarium specimen (KUMF04404) with peltate ramuli from the Gulf of Thailand.

Molecular study

DNA was extracted using the ZR Plant/Seed DNA MiniPrep™ Kit (Zymo Research Corporation, New York, USA) following manufacturer's instructions. Two DNA markers were targeted, *i.e.*, the chloroplast-encoded *tufA* gene and the nuclear internal transcribed spacers (ITS1 and ITS2) of the ribosomal cistron. *TufA* and ITS amplifications were done in a final reaction volume of 20 µl containing 0.2 mM dNTPs, 0.2 µM of each primer, 0.1× Titanium® Taq DNA polymerase and 10× buffer (Clontech Laboratories Inc., Takara Bio company, CA, USA), and 1 µl DNA template (3–20 ng/µl). Forward and reverse primers for *tufA* amplifications were, respectively, *tufA* (5'-TGAAACAGAAMAWCGTCATTATGC-3') and *tufAR* (5'-CCTTCNCGAATMGCRAAWCGC-3') (Famà *et al.*, 2002) or *tufAR1* (5'-CCATAGGAATTGGACTATCA-3') (Stam *et al.* 2006), annealing at, respectively, nucleotide (nt) positions 210, 1184, and 1096 in a complete *tufA* gene (1230 nt) of *C. chemnitzia* (Esper) J.V.Lamouroux (Genbank NC032042, Lam and Lopez-Bautista, 2016, as *C. racemosa*). For ITS amplifications, the primers H1F (5'-CTCTGAACCTTCGCACGTAGA-3') (Kooistra *et al.*, 2002) and ITS4 (5'-TCCTCCGCTTATTGATATGC-3') (White *et al.*, 1990) were used. Double-stranded DNA amplifications were performed in a S1000™ thermal cycler (Bio-Rad Laboratories, California, USA). PCR amplification was started at 96 °C for 4 minutes as a denaturation step, followed by 40 cycles of 30 s at 94 °C, 30 s at 52 °C (for *tufA*) or 48 °C (for ITS), and 60 s at 72 °C for denaturing, annealing and extension steps, and a final extension step at 72 °C for 6 minutes. PCR purification and sequencing were done by Macrogen Inc. (Seoul, Korea) using the amplification primers.

The DNA sequence data set was complemented with sequences from Genbank. In addition, previously unpublished (S.G.A. Draisma) *tufA* and/or ITS sequences of *C. macrodisca* and *C. racemosa* specimens from Indonesia were available for this study (Table 3). These *tufA* sequences were determined as described in Draisma *et al.* (2014). ITS was amplified as described in Stam *et al.* (2006) and subsequently sequenced from cloned amplicons as described in Draisma *et al.* (2012). The herbarium vouchers of these Indonesian specimens are housed in the Naturalis Biodiversity Center in Leiden, The Netherlands, which at the time of our study did not loan any specimens. Sequences were aligned using the MUSCLE software (Edgar, 2004). Molecular species

identification was done by Maximum Likelihood (ML) and Bayesian Inference (BI) phylogenetic inference. ML was performed in MEGA v7 (Kumar *et al.*, 2016) using the General Time Reversible model with Gamma distribution and invariable sites. Clade support was assessed by bootstrap analysis (Felsenstein, 1985) performed with 1,000 pseudoreplicates. BI was performed in MrBayes (Huelsenbeck and Ronquist, 2001) using Markov Chain Monte Carlo chains (MCMC) for 40,000,000 generations, sampled every 4,000th generation with 10% burnin period. The *tufA* dataset included *C. macrodisca* and its sister-species *C. megadisca*. *C. racemosa* was used as an outgroup, because it was shown to be the sister-clade of these two species in Belton *et al.* (2014). A specimen was identified as *C. macrodisca* if in the ML and BI *tufA* trees it was a member of a supported (ML bootstrap percentage (BP) $\geq 80\%$, BI posterior probability (PP) ≥ 0.95) clade that also included the *C. macrodisca* DNA barcode sequence (FM956053) and was sister to a clade including the *C. megadisca* DNA barcode sequence (JN817657). No previously published ITS sequences were available for *C. macrodisca* and *C. megadisca*. Newly generated ITS sequences were analyzed with previously published *C. racemosa* ITS sequences as an outgroup.

Results

Phylogenetic analyses and species confirmation

The *tufA* alignment (40 taxa) was 843 nt positions (234–1076) in length after trimming. The ITS alignment (17 sequences from 12 taxa) was 652 nt positions long including 66 gapped positions. Figure 1 shows the BI *tufA* phylogeny of the *Caulerpa* species under study. The ML tree (not shown) was congruent with the BI tree, revealing the same clades. ML BP and BI PP values are shown in Figure 1. A strongly supported *C. macrodisca* clade (ML BP = 80%, BI PP = 0.98) was sister to a strongly supported *C. megadisca* clade (ML BP = 94%, BI PP = 0.97). The sister-relationship had high to maximum support (ML BP = 96%, BI PP = 1.00). Within the *C. macrodisca* clade four subclades could be discerned, *i.e.*, an Australasian clade (Australia and New Caledonia), an Andaman Sea clade (Thailand and Malaysia), a Java Sea clade, and a Coral Triangle clade (East Kalimantan and West Papua) (Fig. 1). Specimens from the aquarium trade grouped with the Java Sea clade. Relationships among the four subclades remained unresolved. Only the Australasian subclade showed sequence variation (the New Caledonian specimen differed from the Australian specimens). All *Caulerpa* specimens from the Malaysian and Thai mangrove channels on the Andaman Sea coast grouped together in the same *C. macrodisca* subclade.

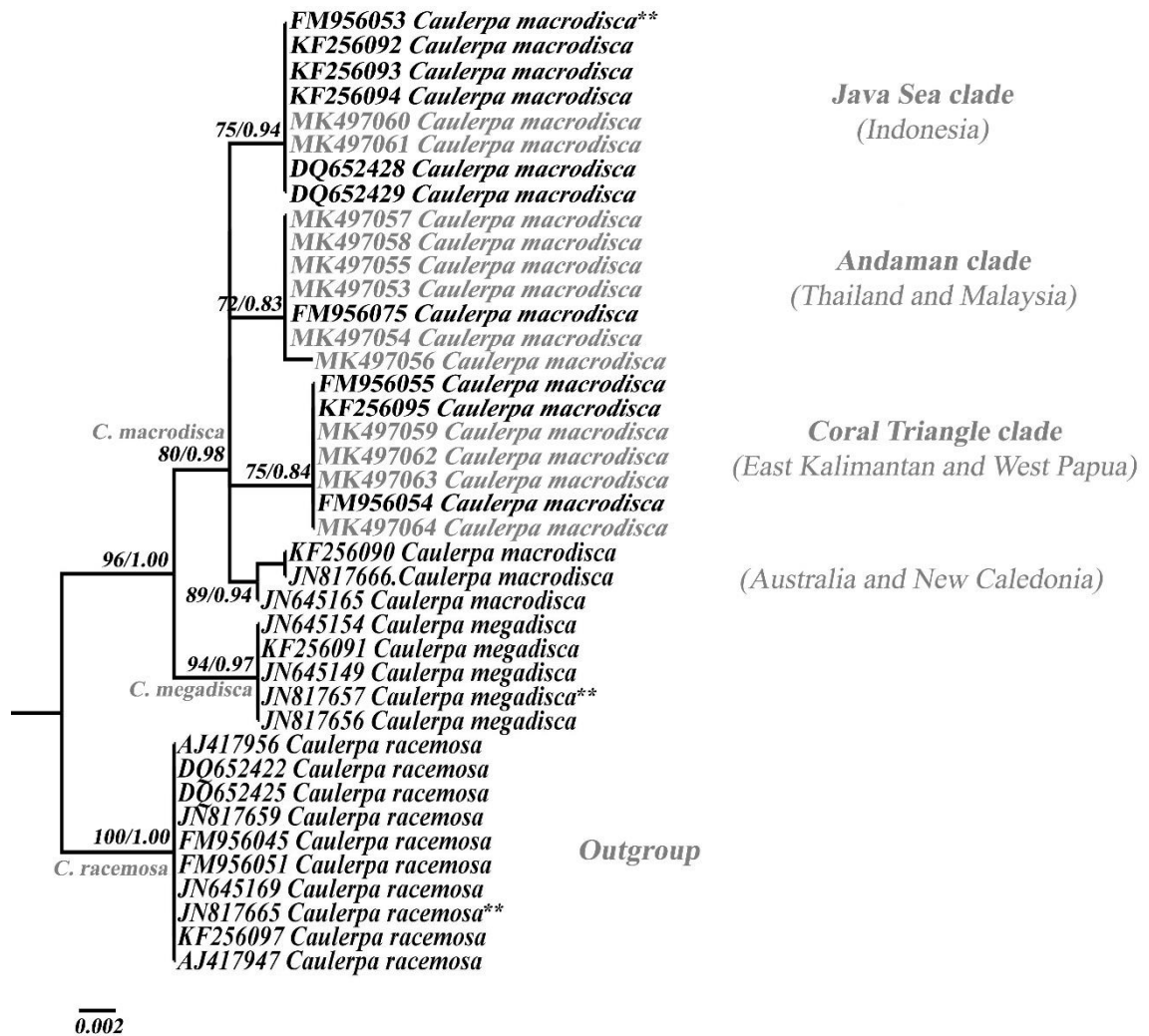


Figure 8 Bayesian phylogenetic tree constructed from 40 partial *tufA* DNA sequences of three *Caulerpa* species; *C. racemosa* (outgroup, n=10), *C. megadisca* (n=5), and *C. macrodisca* (n=25). (alignment = 843 bp in length). Taxon labels include GenBank accession numbers. Numbers at branch nodes correspond to Maximum Likelihood bootstrap percentages (BP) and Bayesian Inference posterior probabilities (PP). BP < 70% and PP < 0.7 are not shown. The scale is 0.002 expected changes per site. Taxon labels in gray are newly generated sequences.

** indicate proposed DNA barcode sequences for each species (Belton *et al.*, 2014).

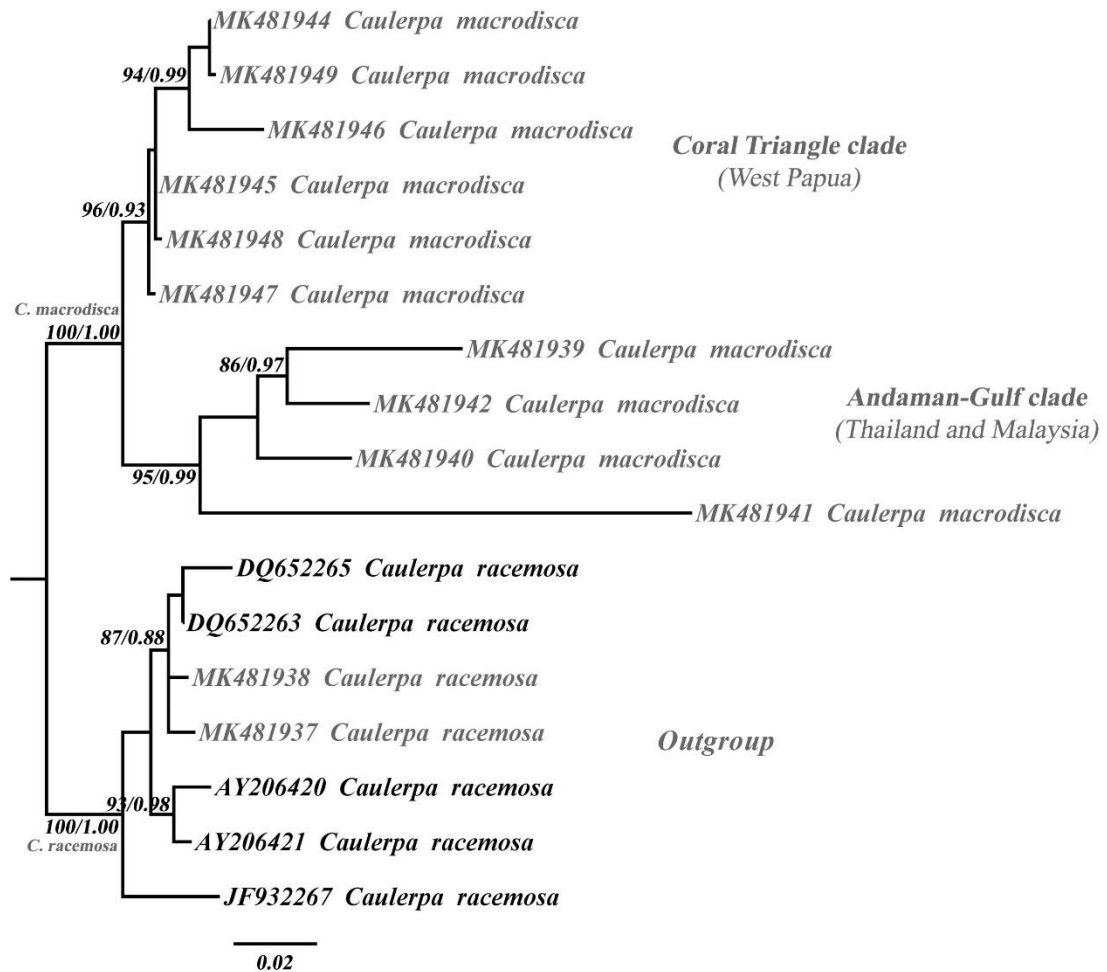


Figure 9 Bayesian phylogenetic tree constructed from 10 *C. macrodisca* and 7 *C. racemosa* (outgroup) ITS rDNA sequences (alignment = 652 bp in length). Taxon labels include GenBank accession numbers. Numbers at branch nodes correspond to Maximum Likelihood bootstrap percentages (BP) and Bayesian Inference posterior probabilities (PP). BP < 70% and PP < 0.7 are not shown. The scale is 0.02 expected changes per site. Taxon labels in gray are newly generated sequences. The six sequences in the Coral Triangle clade were generated from a single individual.

We only succeeded in amplifying the ITS sequence from the peltate *C. macrodisca* herbarium specimen (KUMF04404) from the Gulf of Thailand. It was nested inside the *C. macrodisca* clade in the BI ITS tree (Fig. 9). The ML ITS tree (not shown) was congruent with the BI tree and ML BP and BI PP are shown in Figure 2. *C. macrodisca* from the Gulf of Thailand was nested inside a strongly supported (ML BP = 95%, BI PP = 0.99) clade comprised of *Caulerpa* from the Andaman Sea. This clade was sister to *C. macrodisca* from West Papua (represented by six ITS sequences from a single individual) with maximum support.

Morphological characterization

Caulerpa specimens from the Andaman Sea resembled *C. (racemosa* var.) *corynephora* *sensu* Lewmanomont (2008, figs 12, 13) and *sensu* Coppejans *et al.* (2017, fig. 32) with the exception of KUMF06872, which resembled *Caulerpa ashmeadii* Harvey *sensu* Lewmanomont (2008, figs 1, 2). However, the DNA sequence analysis described above clearly identified them as *C. macrodisca*. We will onwards refer to the three morphological *C. macrodisca* entities in Thailand as ecad: *C. macrodisca* ecad *macrodisca* (typical form bearing peltate ramuli, Figs 10A, B), ecad *corynephora* (with clavate and/or turbinate ramuli, Figs 10C–H, J–M), and ecad *ashmeadii* (with cylindrical ramuli, Figs 10I, N). Table 2 summarizes the morphological features of *C. macrodisca sensu* Belton *et al.* (2014) (*i.e.*, ecad *macrodisca*) and the three ecads in Thailand. The measurements for *C. macrodisca* ecad *corynephora* specimens (KP73 and KP74) were from both fresh and herbarium specimens (indicated in Table 2). The recently collected specimens KUMF06872 and KUMF06874 were not available for measurement..

Table 6 Features of specimens assigned to *C. macrodisca* in the present study and the *C. macrodisca* description provided in Belton *et al.* (2014). Measurements from fresh specimens in square brackets “[]”. n.a., not available. n.m., not measured.

Characters	<i>C. macrodisca</i> Decaisne	<i>C. macrodisca</i> ecad <i>macrodisca</i>	<i>C. macrodisca</i> ecad <i>corynephora</i>	<i>C. macrodisca</i> ecad <i>ashmeadii</i>
	<i>sensu</i> Belton <i>et al.</i> (2014)	from the Gulf of Thailand	from the Andaman Sea	from the Andaman Sea
Specimens examined	see Belton <i>et al.</i> (2014)	KUMF04404	PSU KP73A, PSU KP74A, GENT HEC16156, KUMF06874	KUMF06872
Figures	fig. 9C in Belton <i>et al.</i> (2014)	Fig. 10A	Figs 10C–D, F–H, Figs 11A–B	Fig. 10I
Habitat	n.a.	Intertidal	Mangrove channel, subtidal	Mangrove channel, subtidal
Stolon diameter (mm)	(1.0–)1.5–3.0	1.5–1.8	1.4–2.3 [2.3–3.4]	n.m.
Assimilator height (cm)	1.0–5.0	4.1–6.0	2.7–9.2 [2.9–10.5]	n.m.
Ramuli				
Arrangement	Semi-crowded, radially arranged around an axis	Alternate or irregular	In opposite pairs along the axis, but sometimes alternately opposite (distichous) near the base. KUMF06874 distichous throughout	Mostly in opposite pairs along the axis
Shape	Distinctly peltate	Distinctly peltate	Clavate to mostly turbinate (rarely peltate)	Cylindrical with a swollen tip
Length (mm)	5.0–8.0	<i>ca.</i> 1.0	4.0–10.9 [5.1–12.2]	n.m.
Diameter (mm)	5.0–10.0	6.4–9.0	1.2–4.0 [1.4–5.0]	n.m.



Figure 10 *Caulepa macrodisca* from Thailand and Malaysia (A, C–G, scale bar = 1 cm; B, J–L, scale bar = 0.5 cm; H–I, M–N, no scale bar): (A, B) *C. macrodisca* ecad *macrodisca* (KUMF04404) from Ao Cho, Trat; (A) Habit (herbarium); (B) Drawings of peltate ramuli in apical (left) and lateral (right) views; (C, D) *C. macrodisca* ecad *corynephora* (KP73) from Klong Yang, Krabi; (C) Fresh specimen; (D) Herbarium; (E) *C. macrodisca* ecad *corynephora* (KP65) from Kilim river, Langkawi, Malaysia (fresh); (F, G) *C. macrodisca* ecad *corynephora* (KP74) from Klong Yang, Krabi; (F) Fresh specimen; (G) Herbarium; (H) *C. macrodisca* ecad *corynephora* (KUMF06874) from Che Bilang, Satun; (I) *C. macrodisca* ecad *ashmeadii* (KUMF06872) from Tung Wa, Satun; (J–L) Drawings of ramuli of *C. macrodisca* ecad *corynephora*; (J, K) club-shaped, clavate and (L, M) trumpet-shaped, turbinate; (N) Drawings of cylindrical ramuli of *C. macrodisca* ecad *ashmeadii* (KUMF06872).

Discussions

The *tufA* and ITS phylogenies (Figs 8, 9) showed that *Caulerpa* taxa from Thailand that were previously assigned to *Caulerpa* (*racemosa* var.) *corynephora* and *Caulerpa ashmeadii* (Lewmanomont, 2008; Phang *et al.*, 2008) actually belong to *Caulerpa macrodisca sensu* Belton *et al.* (2014) in the *Caulerpa* section *Caulerpa*. The ITS phylogeny supported that *Caulerpa macrodisca* ecad *macrodisca* from the Gulf of Thailand is conspecific with the other two ecads from the Andaman Sea coast.

The general habit of the *C. macrodisca* ecad *macrodisca* specimen (KUMF04404) (Fig. 10) matched the lectotype illustration of *C. macrodisca* by Decaisne (1846 (1846–1864), pl. 1, fig. 1), but Decaisne's illustration has no scale and dimensions are therefore unknown. However, Decaisne (1842) reported that the peltate disc was 1 cm in his original description. The habit and dimensions of KUMF04404 were in agreement with those of *C. macrodisca* given in Belton *et al.* (2014, table 1, fig. 9C) (Table 6). However, KUMF04404 (Fig. 10A) showed a branched assimilator, which was not reported by Belton *et al.* (2014). The assimilator length of the dried specimen KUMF04404 was 4–6 cm and its ramulus discs 6–9 mm in diameter (Fig. 10B). Lewmanomont (2008, fig. 11) reported an (unbranched) assimilator length of 2.5–7 cm and a disc diameter of 4.5–17.5 mm in intertidal *C. macrodisca* (as *C. peltata* var. *macrodisca*) from Trat province (no voucher numbers given). Specimens from the mangrove channels on the Andaman Sea coast molecularly identified as *C. macrodisca* (Fig. 10) also showed branched assimilators (Figs 10C–N), but deviated in ramuli shape and arrangement from ecad *macrodisca*. The ramuli in *C. macrodisca* ecad *corynephora* (KP73 and KP74) can be up to 12 mm long and are gradually widening from 1.4 mm towards a 5 mm wide rounded (club-shaped, clavate, Figs 10J–K) or blunt (trumpet-shaped, turbinate, Figs 10L–M) terminus. The ramuli in *C. macrodisca* ecad *ashmeadii* (KUMF06872) are cylindrical throughout except for a swollen terminus (Fig. 10N). Ramuli in *C. macrodisca* ecad *corynephora* and ecad *ashmeadii* were arranged in opposite pairs or distichously (alternating) along the rachis.

Earlier molecular phylogenies unambiguously demonstrated that true *C. corynephora* (type location Torres Strait, Australia) belongs to the *Caulerpa* section *Sedoideae* which is characterized by species with pyrenoid-associated chloroplasts and ramuli on

a constricted pedicel (Draisma *et al.*, 2014; Belton *et al.*, 2015, 2019). Several species in the section *Sedoideae*, including *C. corynephora*, have an annulated rachis (Draisma *et al.*, 2014), a character not found outside the section *Sedoideae*. Pyrenoids and an annulated rachis are present in *Caulerpa corynephora* (Price, 2011; Belton *et al.*, 2015, 2019) and this species has only been confirmed from northern and western Australia (Belton *et al.*, 2019). Lewmanomont's (1978, 2008) and Phang *et al.*'s (2008, no description given) identification of *C. racemosa* var. *corynephora* from mangrove channels on the Andaman Sea coast of Thailand and Malaysia may have been based on illustrations in Weber-van Bosse (1898, pl. xxxiii, figs 10–14, 1913, figs 27, 28). However, these illustrations are not an accurate representation of either the holotype specimen or Montagne's (1845) own illustration of *C. corynephora* (Price, 2011). Pyrenoids, constricted pedicels, and annulations were not observed in *C. macrodisca* ead *corynephora* specimens from Thailand and Malaysia, although Coppejans *et al.* (2017, p. 41) stated in the description of *C. corynephora* from Thailand “*The plasts possess pyrenoids, separating C. corynephora from the C. racemosa-peltata complex; this is confirmed by molecular data (Prud’homme van Reine in litteris 2014)*”. However, this statement was based on a miscommunication between Coppejans and Prud’homme van Reine (E. Coppejans and W.F. Prud’homme van Reine, pers. comm. to S.G.A. Draisma 2018). Coppejans *et al.* (2017, fig. 32C) also showed a specimen with cylindrical ramuli, yet they assigned it to *C. corynephora*, not to *C. ashmeadii*. Collection no. HEC16156 A–D (in GENT, as *C. racemosa* var. *corynephora*) consists of multiple sheets, *i.e.*, HEC16156-A (Fig. 11A) and HEC16156-A' (Fig. 11B). The latter contains four separate stolon fragments with assimilators with various ramuli shapes including cylindrical with swollen tips.

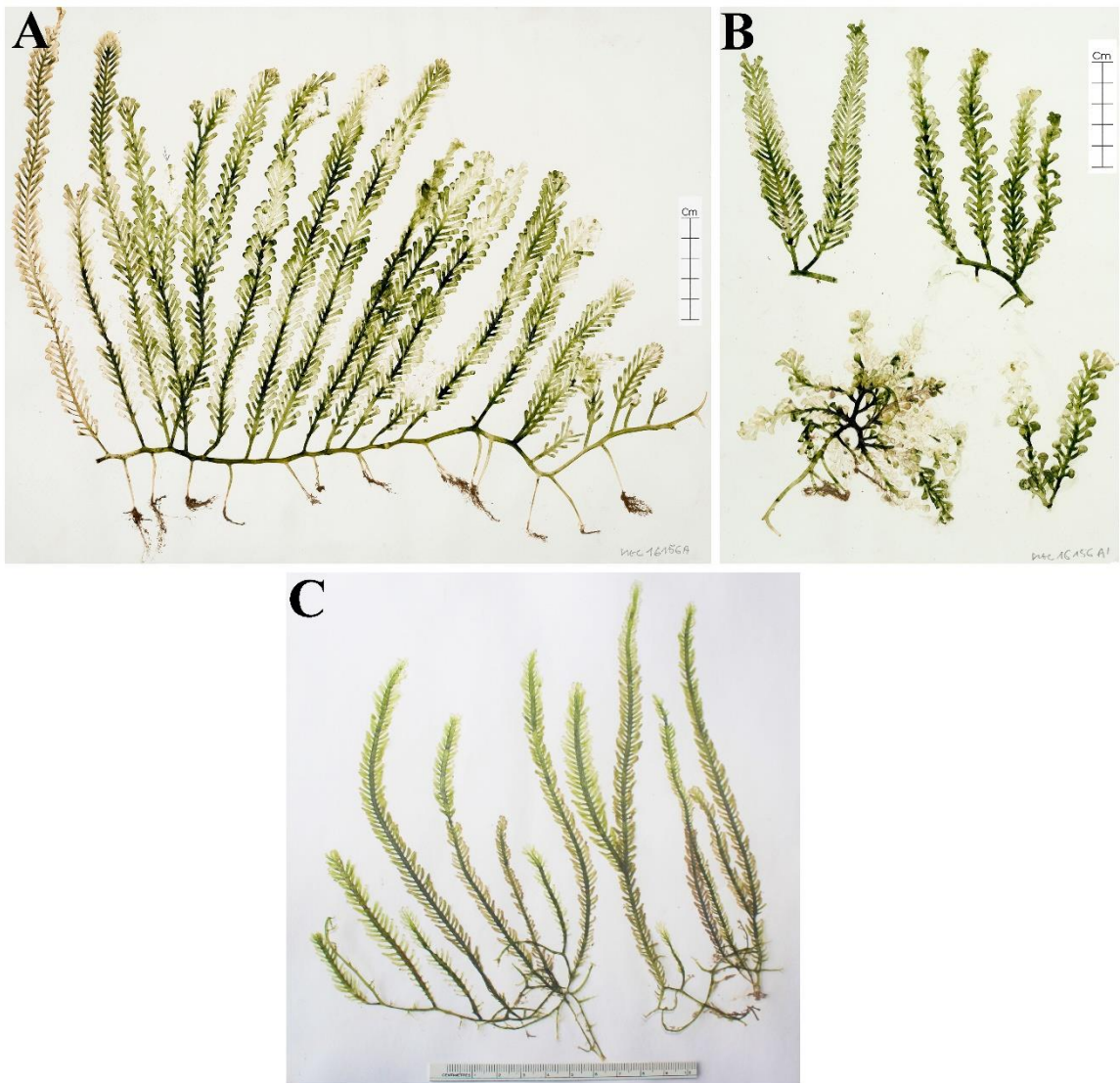


Figure 11 Herbarium specimens of A., *C. macrodisca* ecad *corynephora* HEC16156A-D in GENT, as *C. racemosa* var. *corynephora* (Herbarium sheet HEC16156A), B., *C. macrodisca* ecad *corynephora* HEC16156A-D in GENT, as *C. racemosa* var. *corynephora* (Herbarium sheet HEC16156A) and C., *C. ashmeadii* from Phang Nga province (KL8301 in KUMF)

KUMF06872 (Fig. 10I) from Satun province, originally identified as *C. ashmeadii*, also represents an ecad of *C. macrodisca*. PCR amplification was unsuccessful for KL8301 (in KUMF, Fig. 11C) from Phang Nga province (also on the Andaman Sea coast), which was also morphologically identified as *C. ashmeadii*. It appears not to be the same specimen as the *C. ashmeadii* from Phang Nga depicted in Lewmanomont (2008, figs 1, 2, no voucher number given). However, specimens identified as *C. ashmeadii* from Phang Nga differ from the type specimen of *C. ashmeadii* (type locality Florida) illustrated in Harvey (1858, pl. XXXVIII.A), which shows straight cylindrical ramuli with obtuse tips in an opposite arrangement, whereas ramuli in KL8301 (Fig. 11C) and the specimen depicted in Lewmanomont (2008, figs 1, 2) are slightly curved up. *Caulerpa ashmeadii* has only been confirmed with molecular data from the Caribbean and the Gulf of Mexico and is thought to be confined to the Atlantic (Famà *et al.*, 2002; Stam *et al.*, 2006; Sauvage *et al.*, 2014). However, *C. ashmeadii* has been reported from the Indo-Pacific. Besides the above mentioned Thai records, it has been reported from India (Umamaheswara Rao 1969) and Vietnam (Nguyen *et al.*, 1993). Subsequent Indo-Pacific *C. ashmeadii* reports (Silva *et al.*, 1996; Sahoo *et al.*, 2001; Nguyen, 2007; Nguyen *et al.*, 2013; Phang *et al.*, 2016) can all be traced back to the original reports which could not be accessed by the present authors. *Caulerpa ashmeadii* was not found by Kazi *et al.* (2013) who molecularly identified Indian *Caulerpa* species. Kazi *et al.* (2013) found and molecularly identified *Caulerpa veravalensis* Thivy & V.D.Chauhan from India (Thivy and Chauhan, 1963), a species morphologically similar to *C. ashmeadii*, but with compressed ramuli. *Caulerpa veravalensis*, *C. ashmeadii*, and *C. macrodisca* all belong to the *Caulerpa* section *Caulerpa*, but are not closely related to each other. Each has another sister-species. Herbarium specimens from Thailand identified as *C. ashmeadii* (voucher KL8301 in KUMF (Fig. 11C) and figs 1–2 in Lewmanomont, 2008) are morphologically similar to *C. veravalensis* (see fig. S9 in Kazi *et al.*, 2013). However, whether ramuli originally were compressed or cylindrical cannot be discerned from herbarium specimens. In her description of *C. ashmeadii* from Phang Nga mangroves, Lewmanomont (2008) mentioned that ramuli are cylindrical, whereas ramuli in Indian intertidal *C. veravalensis* are compressed (Kazi *et al.*, 2013).

Another brackish water *Caulerpa* species with terete ramuli, reminiscent of a lanky *C. macrodisca* ecad *ashmeadii*, was described from Swan River, Western Australia,

i.e. Caulerpa lagara Carruthers, Walker & Huisman (Carruthers *et al.*, 1993). It has not been reported since its original description and was therefore not included in Belton *et al.* (2019), a re-assessment of southern Australian *Caulerpa* using DNA sequence data. Conspecificity of *C. lagara* and *C. macrodisca* is not supported by Carruthers *et al.*'s study (1993). They cultured estuarine *C. lagara* in 20, 30, and 40 ‰ salinity, where it did not develop peltate ramuli. In Draisma *et al.* (2014, table S2) it was suggested that *C. lagara* might be a synonym of *C. pinnata* C.Agardh (type location Sri Lanka), but DNA sequence data is lacking for both taxa. Carruthers *et al.* (1993) and Silva *et al.* (1996) mention a resemblance of *C. racemosa* var. *corynephora* (Montagne) Weber-van Bosse to, respectively, *C. lagara* and *C. pinnata*.

Caulerpa macrodisca ecad *corynephora* and ecad *ashmeadii* are morphologically distinct from *C. macrodisca* ecad *macrodisca* and form a distinct clade within *C. macrodisca* in the *tufA* phylogeny (Fig. 8) and therefore could merit the official taxonomic status variety or form. However, this is not supported by the ITS phylogeny (Fig. 9) where *C. macrodisca* ecad *macrodisca* from the Gulf of Thailand (not included in the *tufA* tree) is nested inside the Andaman Sea clade. The phylogenetic pattern may rather be a reflection of a biogeographic pattern. Any infra-specific classification is currently not warranted without *tufA* sequence data of *C. macrodisca* ecad *macrodisca* from the Gulf of Thailand and a study of the morphology of the members of the other *C. macrodisca* subclades (Fig. 8). The differences in morphology are more likely environmentally induced. *C. macrodisca* var. *macrodisca* grew intertidally in the Gulf of Thailand and *C. macrodisca* on the west coast of the Thai-Malay peninsula always grew subtidal in mangrove channels hundreds of meters from the sea. Morphological variation in *Caulerpa* can be caused by several environmental factors such as light intensity, temperature and salinity (Peterson, 1972; Calvert, 1976; Ohba *et al.*, 1992). Ohba and Enomoto (1987) reported that *Caulerpa racemosa* var. *laetevirens* (Montagne) Weber Bosse (probably *C. chemnitzia*) exhibited various ramulus shapes (cylindrical, turbinate and peltate) at different temperatures and light intensities. The estuarine environment of the Andaman mangrove channels may induce the *C. macrodisca* ecad *corynephora* and ecad *ashmeadii* morphologies. Culture experiments should provide more insight into the morphological plasticity of *C. macrodisca*.

CONCLUSION

In conclusion, this study was found eight *Caulerpa* species, i.e., *C. chemnitzia*, *C. lentillifera*, *C. macrodisca*, *C. racemosa*, *C. serrulata*, *C. sertularioides*, *C. taxifolia* and *C. verticillata*. There were three distinct morphotypes of *C. macrodisca* in Thailand. *C. macrodisca* ecad *macrodisca* was only found in the Gulf of Thailand. *C. macrodisca* ecad *corynephora* and *C. macrodisca* ecad *ashmeadii* were only found in mangrove channels on the Andaman Sea coast. In addition, there were two distinct morphotypes of *C. racemosa* in Thai water including *C. racemosa* ecad *chemnitzia* and *C. racemosa* ecad *racemosa*.

The present study highlights the importance of DNA sequence data for reliable species identification in the *Caulerpa racemosa-peltata* complex and to determine the true extent of morphological variation within each species. It is recommended to apply the use of morphological entities like ‘ecads’ which do not have formal taxonomic status for highly plastic species such as *C. macrodisca* and *C. racemosa*. Moreover, morphological variations especially on thallus size and ramuli shape of *Caulerpa* in Thailand might be affected by environmental factors because of its morphological plasticity.

For seven previous species, this study did not find any specimens during this field work, and their identities were not confirmed by molecular study. Then, this result led to a decrease in macroalgal diversity in Thailand. Thus, further studies may focus on a modeling approach to estimate any environmental conditions predicting the future situation of any marine macroalgae in Thai water.

Taxonomic conclusions

1. *Caulerpa chemnitzia* (Esper) J.V.Lamouroux (Fig. 12A)

Synonyms : *Fucus chemnitzia* Esper, *Ahnfeldtia chemnitzia* (Esper) Trevisan, *Chauvinia chemnitzia* (Esper) Kützing and *Caulerpa racemosa* var. *chemnitzia* (Esper) Weber-Van Bosse

Description : Green thallus; rhizoid 0.23–1.56 cm long; large widely horizontal stolon 0.17–0.34 cm in diameter, abundant growth on rock at intertidal zone; erect assimilator, 0.47–1.2 cm in width, 0.87–6.75 cm in length; terete axis 0.1–0.22 cm in diameter; bearing various ramulus shapes from mostly turbinate to peltate into mushroom-like, 0.11–0.78 cm in diameter, supported by short stalks 0.03–0.6 cm long; acute tips.

Vouchers, collection month and location : KP29, KP31 (Ko Kham, Songkhla in October, 2016), KP32 and KP50 (Losin, Pattani in August, 2017)

2. *Caulerpa lentillifera* J.Agardh (Figs 12B–C)

Synonyms : *Ahnfeldtia lentillifera* (J.Agardh) Trevisan and *Chauvinia lentillifera* (J.Agardh) Kützing

Description : Light green thallus; rhizoid 0.11–0.78 cm long; horizontal stolon 0.13–0.18 cm in diameter, abundant growth on rock and dead coral at intertidal zone; Erect branch with dense and short ramuli on axis; terete axis, 0.1–0.15 cm in diameter; spherical translucent ramulus shape, 0.21–0.23 cm in diameter, supported by short stalks 0.05–0.09 cm long, prominently constricted at the base, imbricately arranged or in rows of four; rounded tips; chloroplast with pyrenoid.

Vouchers, collection month and location : KP6 and KP21 (Ko Siboya, Krabi in February and April, 2016)

3. *Caulerpa macrodisca* Decaisne (Figs 12D–G)

Synonyms : *Ahnfeldtia macrodisca* (Decaisne) Trevisan, *Chauvinia macrodisca* (Decaisne) Kützing, *Caulerpa peltata* var. *macrodisca* (Decaisne) Weber-van Bosse and *Caulerpa racemosa* var. *macrodisca* (Decaisne) Weber-van Bosse

Caulerpa macrodisca ead *macrodisca* (Figs 12D–E)

Description : Green thallus; rhizoid 0.61–2.35 cm long; terete stolon, 0.15–0.18 cm diameter, abundant growth on sandy or muddy bottoms of turbid water; erect assimilators, 0.54–1.23 cm in width, 4.13–6.05 cm in length; terete axis 0.06–0.1 cm diameter, bearing large thin disc, 0.64–0.9 cm diameter; with short stalks, sometimes, a single peltate branch (generally 1 cm. or more diameter) was consisted along terete stolon; disk tips.

Vouchers, collection month and location : KUMF04404 (Ao Cho, Trat in January, 1991)

Caulerpa macrodisca ead *corynephora* (Figs 12F–G)

Description : Light green thallus; rhizoid 0.84–5.05 cm long; horizontal stolon, 0.23–0.34 mm in diameter, abundant growth on fish's cage in the brackish water of the Andaman mangrove channels; erect assimilators, 0.65–0.94 cm in width, 2.89–10.48 cm in length; terete axis, 0.14–0.21 cm in diameter, constriction at the basis; bearing generally 2 to 4 longitudinal rows with opposite various shapes of ramulus from mostly clavate (1 mm in diameter) to turbinate into rarely peltate (disc 5 mm in diameter, nearly the top), 0.14–0.5 cm in diameter; without constriction at the base; acute tips; chloroplasts without associated pyrenoids.

Vouchers, collection month and location : KP65 (Kilim river, Malaysia in December, 2017), KP73, KP74 (Khleng Yang, Krabi in July, 2018) and KUMF06874 (mangrove channel, Satun in September, 2018)

Caulerpa macrodisca ecad *ashmeadii*

Description : light green thallus, growth in the Andaman mangrove channel; erect assimilators bearing generally opposite ramuli arrangement (sometime alternate at base); cylindrical ramulus shape with swollen tips

Vouchers, collection month and location : KUMF06872 (mangrove channel, Satun in September, 2018)

4. *Caulerpa racemosa* (Forsskål) J.Agardh (Figs 12H–L)

Synonyms : *Fucus racemosus* Forsskål

Caulerpa racemosa ecad *racemosa* (Figs 12H–I)

Description : Dark green thallus; rhizoid 0.28–4.64 cm long; large widely horizontal stolon 0.13–0.39 cm in diameter, abundant growth on rock and dead coral at intertidal zone; strong short erect branch, 0.29–1.88 cm in width, 0.44–5.98 cm in length; with dense ramuli; terete axis 0.09–0.47 cm in diameter, with constriction at the basis; subspherical to spherical (vesiculate) ramulus shape with rounded tip, 0.22–0.71 cm in diameter; supported by short stalk 0.06–0.47 mm long, 0.11–0.31 cm in diameter; with constriction at the base, 0.07–0.26 cm in diameter; rounded tip.

Vouchers, collection month and location : KP10 (Ko Tan, Surat Thani in February, 2016), KP7, KP22 (Ko Siboya, Krabi in February and April, 2016), KP39, KP40 (Ko Lipe, Satun in January, 2017), KP68, KP69 and KP70 (Ko Lanta, Krabi in March, 2018)

Caulerpa racemosa ead *chemnitzia* (Figs 12J–L)

Description : Green thallus; rhizoid 0.12–2.76 cm long; horizontal stolon 0.15–0.25 cm in diameter, abundant growth on rock at intertidal zone; erect assimilators, 0.33–1.39 cm in width, 0.84–4.81 cm in length; terete axis 0.1–0.19 cm in diameter, with constriction at the basis; bearing various ramulus shapes from mostly turbinate to peltate or sometimes mushroom-like, 0.11–1.01 cm in diameter, supported by short stalks 0.03–0.43 cm long; acute tips.

Vouchers, collection month and location : KP28, KP30 (Ko Kham, Songkhla in October, 2016), KP49 and KP51 (Losin, Pattani in August, 2017)

5. *Caulerpa serrulata* (Forsskål) J.Agardh (Fig. 12M)

Synonyms : *Fucus serrulatus* Forsskål and *Caulerpa freycinetii* f. *serrulata* (Forsskål) Weber-van Bosse

Description : Heavy green thallus with yellow tip; descending branches having branched rhizoid at the ends 0.1–5.74 cm long; large widely horizontal stolon 0.11–0.36 cm in diameter, widespread and dense colonies on rock and dead coral at intertidal zone; dichotomous branching, twist once or twice; upright flattened branches with serrate margin; branch 0.21–0.59 cm in width, 0.36–4.67 cm in length; supported by cylindrical stalks 4 mm long or less; teeth tips.

Vouchers, collection month and location : KP11 (Ko Tan, Surat Thani in February, 2016), KP14, KP45 (Ko Samui, Surat Thani in February, 2016 and August, 2017), KP41 (Ko Lipe, Surat Thani in January, 2017), KP47 (Ko Matsum, Surat Thani in August, 2017), KP72 (Tang Khen Bay, Phuket in March, 2018), SP358, SP284, SP301, SP358, SP368 (Ko Rok, Krabi in February, 2011) and SP372 (Ko Kradan, Trang in February, 2011)

6. *Caulerpa sertularioides* (S.G.Gmelin) M.Howe (Figs 12N–O)

Synonyms : *Fucus sertularioides* S.G.Gmelin

Description : Green widespread thallus, feather like erect branches; rhizoid 0.06–3.78 cm long; large strong stolon 0.07–0.29 cm in diameter, abundant growth on sand, rock and dead coral at intertidal zone; distichously arranged assimilators along axis, slender, cylindrical, oppositely arranged, pinnules ramulus; branch, 0.21–1.56 cm in width, 0.56–7.89 cm in length, rarely dichotomous, branching 1–2 times; yellow mucronate tips.

Vouchers, collection month and location : KP15 (Ko Samui, Surat Thani in February, 2016), KP34, KP35 and KP61 (Ko Lidee, Satun in November-December, 2016 and December, 2017)

7. *Caulerpa taxifolia* (M.Vahl) C.Agardh (Figs 12P–R)

Synonyms : *Fucus taxifolius* M.Vahl

Description : Green thallus; rhizoid 0.58–6.6 cm long; horizontal stolon 0.11–0.32 cm in diameter, abundant growth on rock at intertidal zone; upright branch with opposite ramuli arrangement, usually branched; branches, 0.133–1.46 cm in width, 1.8–11.42 cm in length; clearly flattened and upright falcate ramulus shape 0.07–0.13 cm wide, 0.26–0.77 cm long, with obviously constricted at base; mucronate tips.

Vouchers, collection month and location : KP8, KP23 (Ko Siboya, Krabi in February and April, 2016), KP43 (Ko Libong, Trang in May, 2017), KP46 (Ko Matsum, Surat Thani in August, 2017), KP53, KP54, KP55, KP56 (Leam Yong Lam, Trang in September, 2017), KP57 (Khao Bae Na, Trang in September, 2017), SP332, SP343, SP234 (Ko Rawi, Satun in February, 2011) and SP474 (Ko Adang-Rawi, Satun in December, 2011)

8. *Caulerpa verticillata* J.Agardh (Figs 12S–U)

Synonyms : *Stephanocoelium verticillatum* (J.Agardh) Kützing

Description : Green filiform thallus; rhizoid 0.2–2.81 mm long; stolon 0.18–0.39 mm diameter, abundant tuft on rock and dead coral; 4–7 times dichotomously branched, whorled branching 0.94–5.28 mm wide, 2–10.97 mm high, dense overlap layers (more than 5 layers); branchlets cylindrical 0.03–0.07 mm width, 0.19–3.36 mm long; tree spine tips.

Vouchers, collection month and location : KP1, KP5, KP13, KP62 (Ko Lidee, Satun in January, 2015–2016 and December, 2017), KP9, KP24 (Ko Siboya, Krabi in February and April, 2016) and KP42 (Ko Lipe, Satun in January, 2017)

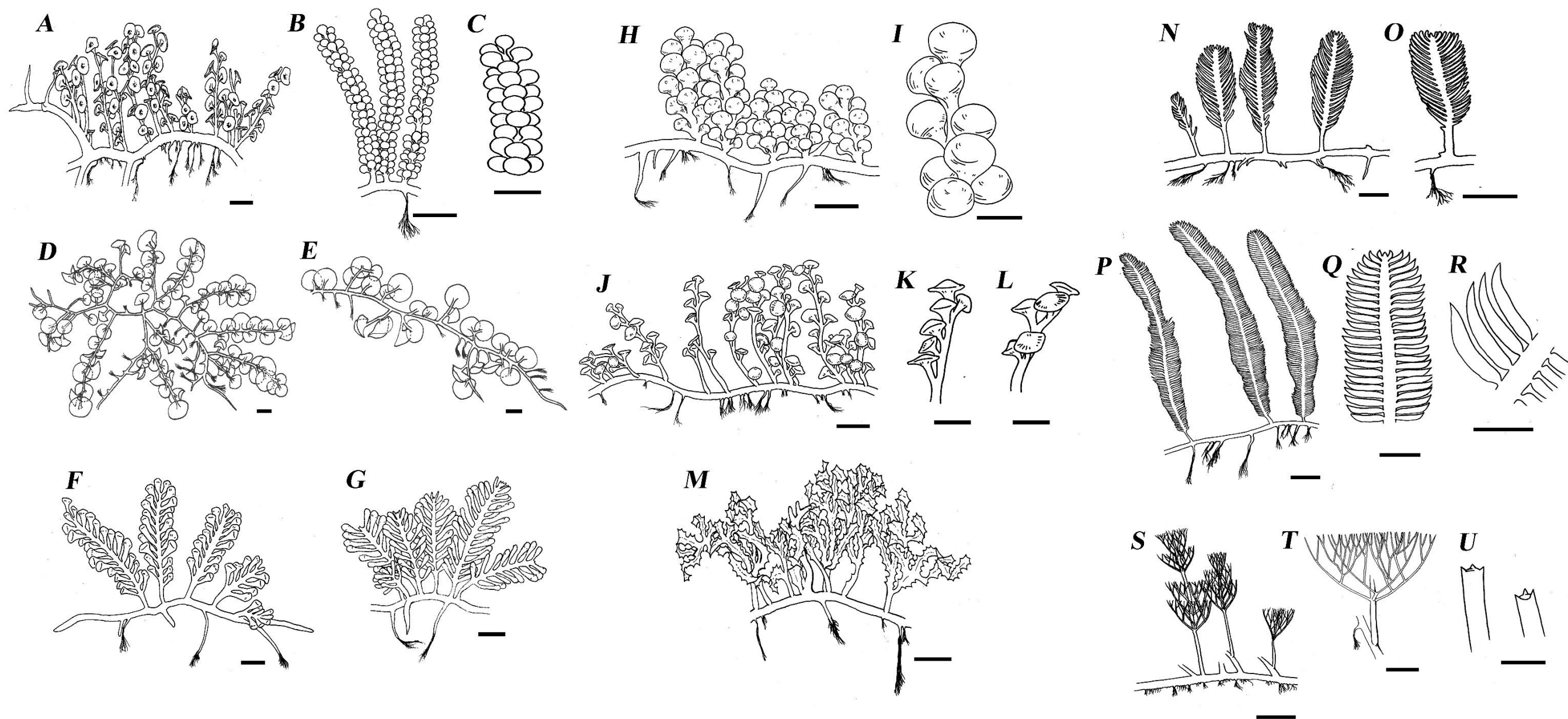


Figure 12 Eight *Caulerpa* species were found in Thailand (A–G., H., J., M., P., scale bar = 1 cm; I., K–L., N–O., Q–R., T., scale bar = 0.5 cm; S., scale bar = 1 mm; U., scale bar = 50 μ m) including A., *C. chemnitzia*, B–C., *C. lentillifera*, D–G., *C. macrodisca* (D–E., *C. macrodisca* ecaud *macrodisca* and F–G., *C. macrodisca* ecaud *corynephora*), H–L., *C. racemosa* (H–I., *C. racemosa* ecaud *racemosa* and J–L., *C. racemosa* ecaud *chemnitzia*), M., *Caulerpa serrulata*, N–O., *C. sertularioides*, P–R., *C. taxifolia* and S–U., *C. verticillata*.

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Pattarach, K., Phetcharat, S. and Mayakun, J. 2019. Some species of *Gracilaria* (Gracilariaceae, Rhodophyta) from Southern Thailand. In: *Taxonomy of Southeast Asian Seaweeds*, Institute of Ocean and Earth Sciences, University of Malaya, Monograph Series 17 (eds S. M. Phang, S. L. Song and P. E. Lim). UM Press, Kuala Lumpur. 47–57.