IRISH CLAIRE LITERATUS

MASTER OF SCIENCE IN PLANT PATHOLOGY

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> GRADUATE SCHOOL CHIANG MAI UNIVERSITY MAY 2023

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A THESIS SUBMITTED TO CHIANG MAI UNIVERSITY IN PARTIAL FULFILMENT OF THE REQUIREMENTS FOR THE DEGREE OF MASTER OF SCIENCE IN PLANT PATHOLOGY

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GRADUATE SCHOOL, CHIANG MAI UNIVERSITY MAY 2023

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ACKNOWLEDGEMENTS

I dedicate this thesis, first and foremost, to our omnipotent, omniscient, and omnipresent loving God for the provision in every aspect of my life throughout my graduate study journey.

To my supportive adviser, Assoc. Prof. Dr. Ratchadawan Cheewangkoon, the commencement of this study is the result of your amazing leadership, supervision, and encouragement. I will always be grateful to you. Your brilliance, kindness, and generosity will be remembered. To Dr. Milan C. Samarakoon and Asst. Prof. Dr. Sararat Monkhung, thank you for imparting your knowledge and expertise during the thesis examination and revisions.

To my seniors in the laboratory, Miss Sukanya Haituk, Miss Patchareeya Withee, Miss Thiyagaraja Vinodhini, Mister Anuruddha Karunarathna, and Miss Dulanjalee Harishchandra, I couldn't thank you enough for every effort, time, and knowledge that you have given in the pursuit of completing this work.

I also dedicate this work to my late father, Jovanie Literatus, whose zeal for his work had inspired me to pursue a degree in Agriculture, my beloved mother, Marilou Literatus, who has been my greatest supporter, and my best friend, Lane Shaikoski, whose life has been positively influencing mine.

To my dear friends CJ, Gemma, Lai, Lissa, and Myla, thank you for your prayers and unwavering support. Lastly, to these people, my church family- Living Praise of Zion-Valencia Bukidnon, Philippines, my pastors Anacorita Querubin, Joey and Raya Gregorio, my cell group leader, Deesri Jenn de Isidro-Lerit, and my cell groupmates-Navigators, I am blessed with your lives. Thank you for your unceasing prayers.

Lastly, thank you to Chiang Mai University Graduate School for offering the CMU Presidential Scholarship. I am positive that more graduate students will be able to pursue further studies and establish connections through this scholarship program.

Irish Claire Literatus

หัวข้อวิทยานิพนธ์	อนุกรมวิธานและวงศ์วานวิวัฒนาการของราสาเหตุใบจุดนูนคำบน โพธิ์ <i>Ficus religiosa</i> ในจังหวัดเชียงใหม่	
ผู้เขียน	นางสาว ไอริช แคลร์ ลิตเทอราตัส	
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บทคัดย่อ

ใบจดนนดำเป็นโรคที่มีสาเหตุมาจากราวงศ์ Phyllachoraceae สามารถพบได้บ่อยในเขตร้อน และชิ้น ส่วนใหญ่พบในประเทศแถบเอเชีย เช่น ประเทศไทย โดยอาการมักเกิดที่ขึ้นใบ ลำต้น และผล ในปัจจุบันมีการศึกษาที่สำคัญเกี่ยวกับราชนิดนี้เพียงไม่กี่ชิ้นเท่านั้น เนื่องจากราสาเหตุโรคดังกล่าวไม่ ้สามารถเจริญเติบโตในอาหารเลี้ยงเชื้อได้ ส่งผลให้ข้อมูลการกระจายตัวทางภูมิศาสตร์ และข้อมูลทาง ซึ่งการศึกษาส่วนใหญ่รายงานว่าโรคนี้เกิดจากสกุล อณูชีวิวทยาจึงเป็นเรื่องยากสำหรับการศึกษา Phyllachora อย่างไรก็ตาม มีรายงานโรคใบจุดนูนคำบางส่วนเกิดจากสกุลอื่น และยังพบการแพร่ ระบาคของโรกใบจุดนูนดำหลายจุดซึ่งยังไม่สามารถระบุสาเหตุได้ ในประเทศไทยต้นโพธิ์มีบทบาท สำคัญในการอนุรักษ์วัฒนธรรมและประวัติศาสตร์ อย่างไรก็ตาม ต้นโพธิ์กำลังเผชิญกับภาวะที่กลืน ไม่เข้าคายไม่ออกเนื่องจากการโจมตีของเชื้อโรค โดยเฉพาะบนใบ ซึ่งทำหน้าที่เป็นจุคศูนย์กลางของ ดังนั้นการศึกษาครั้งนี้จึงมีจุดประสงค์เพื่อสำรวจและเก็บรวบรวมโรคใบจุดนูนคำใบโพธิ์ ความงาม ในพื้นที่จังหวัดเชียงใหม่ และทำการศึกษาความสัมพันธ์ทางพันธุกรรรมของรา โดยใช้ข้อมูลทาง สัณฐานวิทยา ร่วมกับข้อมูลทางอณูชีววิทยา โคยสามารถรวบรวมตัวอย่างของใบโพธิ์ที่แสดงอาการ ใบจุดนูนคำได้ทั้งหมด 12 ตัวอย่าง จาก 7 อำเภอ ได้แก่ แม่แจ่ม แม่วาง แม่แตง แม่ริม สันทราย หางคง โดยใบจุดนูนดำจะแสดงอาการแผลจุดเดี่ยวสีดำ หรือรวมกันเป็นจุดใหญ่เรียก สันป่าตอง ແລະ Pseudostromata มีลักษณะยาว ไม่สม่ำเสมอ ไม่ต่อเนื่อง เบาบาง รวมตัวกัน เกลี้ยง เป็นมันเงา ใน ้ผิวหนังถึงชั้นใต้ผิวหนัง epiphyllous จากนั้นทำ การวิเคราะห์ลำดับนิวคลิโทไทด์ของรา ในตำแหน่ง ITS และ LSU พบว่า จัดจำแนกราได้จำนวน 1 สกุล คือ Neophyllachora ในการศึกษาครั้งนี้ได้พบ Neophyllachora fici ซึ่งเป็นการรายงานชนิดใหม่จากประเทศไทย ซึ่งได้อธิบายลักษณะสัณฐานวิทยา และวงศ์วานวิวัฒนาการไว้ในการศึกษาครั้งนี้ด้วย



Thesis Title	Taxonomy and Phylogeny of Fungi Causing Tar Spo	
	Ficus religiosa in Chiang Mai Province	
Author	Miss Irish Claire Literatus	
Degree	Master of Science (Plant Pathology)	
Advisory Committee	Assoc. Prof. Dr. Ratchadawan Cheewangkoon Dr. Milan C. Samarakoon	Advisor Co-advisor

ABSTRACT

กมยนติ

Tar spot is a common fungal disease often found in tropical and damp areas. They mostly exist in Asian countries like Thailand, and the symptoms usually occur on leaves, stems, and fruits. Only a few significant studies about this fungus are available because of their biotrophic nature, which makes them unable to grow in culture, therefore, sequence data from fresh collections are difficult to obtain. On one hand, most studies recorded this disease to be caused by the genus Phyllachora. On the other hand, some Tar spot symptoms are known to be caused by other genera, and there are still several Tar spot infestations in which the causal agent is yet to identify. In Thailand, the Bodhi tree plays an important role in culture and historic preservation. However, the Bodhi tree has been facing a dilemma due to attacks of pathogens, particularly on the leaves, which serve as the center of its beauty. This study aimed to identify the causal agent of the Tar spot on Bodhi tree leaves taken from different locations around Chiang Mai province, northern Thailand, through morpho-molecular evaluation. We isolated eight leaf samples of *Ficus* religiosa taken from Chiang Mai, Thailand. Symptoms on the host appear as black, solitary to gregarious, mainly on the upper surface. Pseudostromata is elongated, irregular, discrete, sparse, coalescent, glabrous, shiny, intraepidermal to subepidermal, epiphyllous, multilocular, occasionally amphigenous, rarely covering the leaf surface. Morphologically the new species was characterized by pseudostromatic ascomata with ostiole, septate paraphyses, cylindrical to fusiform asci, and globose to elliptical ascospores with sometimes 1-2 guttules, central concave depression present mostly in the globose form, a mucilaginous sheath that is irregularly thickened and widely thickened in the lateral part. The asexual morph produces ellipsoidal and hyaline conidia. The newly

obtained sequences were positioned within *Neophyllachora* and formed a distinct clade but close to *Neophyllachora fici* with high bootstrap support in the phylogenetic analyses. In addition, both species are reported in the same host genus (*Ficus*) but with different locality (Thailand Vs. Taiwan). Further the synopsis table for *Neophyllachora* and the identification key to the genus are provided.



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CHAPTER 1

INTRODUCTION

1.1 Introduction

Tar spot is a common fungal leaf disease characterized by the formation of black stromata, which is a slightly raised, semi-circular, dark brown to black and lustrous structure (Cannon 1991). The leaf spots caused by tar spot pathogens on leaves decrease the rate of photosynthesis and alter the physiological and biochemical aspects resulting in a reduction in crop production (Ballesta et al. 2010, Karami et al. 2014). The fungal disease has a wide host range, and the size of the spots can range up to several cm across (Hsiang et al. 2008). Tar spot also affects some economically and culturally important crops such as wheatgrass (Conners 1967), job's tears (Titatarn et al. 1984), sorghum (Long et al. 1985), bodhi tree (Hsieh et al. 2003), and corn (Ruhl et al. 2016).

Members of Phyllachoraceae are mostly obligate parasites which form "tar spots" on leaves and occasionally on stems and fruits with most are host-specific (Cannon 1997). The taxa are known as minor pathogens that rarely kill the host tissues but provide pathway for the secondary infection from severe pathogens by retaining for a long time in the host tissue (Cannon 1991, Hock et al. 1992, Parbery 1996). Phyllachoraceae was established by Theissen & Sydow (1915) with *Phyllachora* as the type genus. The family accommodates more than thousand species worldwide and are included in 54 genera (Tennakoon et al. 2020, Wijayawardene et al. 2022). However, molecular data are available only for a few members because of the difficulties in obtaining cultures from fresh samples (Hyde et al. 2020).

The taxa are characterized by ascohymenial development with paraphyses, thinwalled asci, which may have an apical ring that does not stain blue in iodine (J-) and often hyaline and 1-celled ascospores. The asexual morph is coelomycetes, spermatial or disseminative (Hawksworth et al. 1983). The family was placed in different orders including Dothideales (Horst 1990), Sphaeriales (Nannfeldt 1932, Miller 1949, Müller and von Arx 1962, Wehmeyer 1975), Xylariales (Luttrell 1951, Barr1990), Glomerellales (Chadefaud 1960, Locquin 1984), Phyllachorales (Barr 1976a, b, 1983), Polystigmatales (Eriksson 1982, Hawksworth et al. 1983), and Diaporthales (Cannon 1988). Maharachchikumbura et al. (2016) placed Phyllachoraceae in Phyllachorales with Phaeochoraceae and this classification has been accepted by further studies (Hongsanan et al. 2017, Hyde et al. 2020). Consequently, Mardones et al. (2017) and Guterres et al. (2019) added two new families namely Telimenaceae and Phaeochorellaceae to this order.

In Thailand, several studies have reported *Phyllachora* species, including *Phyllachora bambusae*, *P. chloridis*, *P. coicis*, *P. cynodontis*, P. cynodonticola, *P. digitariae*, *P. graminis*, *P. pterocarpi*, *P. repens*, *P. thysanolaenae* and *P. vetiveriana* (Giatgong 1980, Nuangpai et al. 1984, Lenne 1990, Athipunyakom & Likhitekaraj 2009, Dayarathne et al. 2017, Tamakaew et al. 2017). This study provides the first report of the genus *Neophyllachora* from Thailand which was also reported from the leaves of *Ficus religiosa* for the first time. We also introduce a new species to *Neophyllachora* based on the evidence from both morphology and phylogeny. The synopsis table for *Neophyllachora* with the key to the species are provided.

1.2 Research Objectives

1.2.1 To provide morpho-molecular identification of the fungi causing Tar spots on *Ficus religiosa*.
1.2.2 To assess the biogeography of the fungi causing Tar spots on *Ficus religiosa* in Chiang Mai Province.

1.3 Usefulness of the Research (Theoretical and/or applied)

1.3.1 This study aims to increase the knowledge and attention about the flourishing Tar spot infestation in *Ficus religiosa* leaves in Chiang Mai Province. The results of taxonomy and phylogeny will be published for the scientific community for the use of pathologists, taxonomists, and quarantine systems. The sequences and phylogenetic data will be submitted to databases for future studies.

CHAPTER 2

LITERATURE REVIEW

2.1 An overview of Tar spot

Tar spot is a fungal leaf disease that infects several plants. The symptom starts with small yellow spots on growing leaves, then expands into large black blotches. In addition, the pathogen produces a circular pattern of black fruiting bodies within these spots (Zacaroni et al. 2013). As the fungus grows, the spreading yellow spot slowly turns from a yellow-green to a deep, tarry black, referred to as a tar spot (https://www.hamilton.ca/home-property-and-development/property-gardens-trees/tar-

spot-disease). In the winter season, the disease can cause intense defoliation in susceptible plants (Zacaroni et al. 2013). However, tar spots can be easily confused with the black saprophytic organisms that grow on dead leaf tissue. To elude this, take notice that saprophytes usually have a dusty appearance and can be rubbed off the leaf tissue while tar spot stromata cannot be rubbed off, and it is usually surrounded by a narrow tan halo (Kleczewski et al. 2019). According to Hudler et al. (1998), the disease was first reported in Ohio in the 1940s, and their study unveiled that the fungus *Rhytisma acerinum*, was responsible for the tar spot occurring on a variety of maples in Europe, where various studies about tar spot diseases were conducted (Hsiang et al. 2008). The disease has diverse host plants, including some economically important crops and trees. The causal agents are commonly from *Rhytismales* and *Phyllachorales*. The former is frequently found in forest areas or foothills of high ranges occupying a special niche mostly on forest plants, while the former prefers a cropland environment or open plain (Thaung, 2008).

2.2 Tar spot disease caused by Phyllachorales

Phyllachorales are mostly linked to angiosperms, with the following families preferentially parasitized: *Arecaceae, Fabaceae, Lauraceae, Melastomataceae,*

Moraceae, Myrtaceae, and Poaceae (Cannon 1997). Additional host families studied include Asclepiadaceae (Pearce et al. 2001), Erythroxylaceae (Cannon 1988), Proteaceae (Pearce et al. 2001), and Rosaceae (Cannon 1988).

Currently, there are at least 3 families belonging to *Phyllachorales* - *Phaeochoraceae*, *Phyllachoraceae* (Maharachchikumbura et al. 2016; Hongsanan et al. 2017), and *Telimenaceae* (Mardones et al. 2017). According to Maharachchikumbura et al. (2016), members of the *Phyllachoraceae* is characterized by forming leaf spots on the host that are abundant but scattered, raised, mostly rounded to oblong or elongated, sometimes parallel with leaf venation, surrounded by a light-brown necrotic region; lacking periphyses; having numerous paraphyses, branched or unbranched; 8-spored asci, persistent, cylindrical to fusiform, often present with an apical ring; ascospores fusiform to narrowly oval, hyaline, often with a mucilaginous sheath. *Phyllachoraceae* is similar *to Phaeochoraceae*, but the former species are characterized by 8-spored asci, an often-present apical ring, usually hyaline ascospores, rarely pale brown, thin and smoothwalled, while the latter is characterized by 6-8-spored asci, usually without apical structure, yellow to olivaceous ascospores or in various shades of brown, thick-walled (Maharachchikumbura et al. 2016; Mardones et al. 2017).

Phyllachora is the largest genus of *Phyllachoraceae*, and they are morphologically characterized by clypeate pseudostroma in leaf tissues, generalized infection of the entire section of the mesophyll forming leaf spots on the host, mostly rounded to oblong or elongated, surrounded by,light-brown necrotic region; perithecium globose; numerous paraphyses, branched, slightly longer than asci; asci 8-spored, persistent, cylindrical to fusiform, a short pedicellate, an apical ring often present; and ascospores 1–3 seriate, fusiform to narrowly oval, hyaline, sometimes with a gelatinous sheath (Dos Santos et al, 2016; Maharachchikumbura et al. 2016; Yang et al. 2019). *Phyllachora* species are commonly found with *Poaceae* but have been reported to infect more than 1000 plant species including *Cyperaceae*, *Fabaceae*, *Lauraceae*, *Moraceae*, *Myrtaceae*, *Poaceae*, *Proteaceae*, and *Rosaceae* (Li et al. 2022).

In a study conducted by Li et al. (2022), they found that the *Phyllachora* genus is paraphyletic since the host of *Phyllachora pomigena* remains unknown, the species formed a single clade. They further explain that in the phylogenetic analysis, the new species described are included within the *Phyllachora* genus and separated from other taxa with a single subclade. Their hosts are *Cenchrus flaccidus* and *Chloris virgata*, both belonging to *Poaceae* (graminicolous).

2.3 Reports on Tar spot disease caused by *Phyllachora* worldwide

In 1991, black spots (caused by *Phyllachora repens*) were seen on the leaves of Bodhi trees at the National Chung Hsing University Campus in Taiwan. The diseased trees resulted in severe defoliation out of season. The obligate parasite produced a teleomorph state and spermogonia however, no anamorph was found (Hsieh et al. 2003). Although this incident is interesting, no recent studies are available after the report.

The tar spot of corn (*Phyllachora maydis*) was first confirmed in the United States in 2015. In 2018, a yield-reducing epidemic of tar spots occurred in northern Indiana and surrounding states. Following this epidemic, tar spot was detected in 172 counties across six states in the Midwest. Fields in the most severely affected regions reached 100% disease incidence and over 50% severity on the ear leaf (Kleczewski et al. 2019).

2.4 Reports on Tar spot disease caused by Phyllachora in Thailand

In a disease survey conducted in November 1983, tar spots (*Phyllachora coicis*) were identified from the upper and lower leaf surfaces of Job's tears (*Coix lachryma-jobi*) in the vicinity of Dong Ma Da village of Mae Suay district of Chiang Rai province. The symptoms were characterized by roughly circular raised black bodies (fungal stromata) about 1.5 mm in diameter and were estimated to have a more than 70 percent level of infection (Nuangphai 1984).

A different study by Boon-Long et al. (1987) reported the occurrence of tar spot disease in sorghum and the pathogen was identified as *Phyllachora sorghi*. Hitherto, no further studies are available after the report.

2.5 An overview of the Bodhi Tree

The Bodhi tree (*Ficus religiosa*) is a perennial tropical tree that belongs to *Moraceae*. It has grayish bark and heart-shaped leaves and can grow up to 9 feet in diameter. This large broadleaf evergreen tree is native to Southeast Asia and India and recorded from the mid-19th century as the 'tree of knowledge (https://www.encyclopedia.com/plants-and-animals/plants/plants/bo-tree). The Bodhi

tree has other common names like the Bo tree, Peepul tree, and Sacred fig. It is culturally, spiritually, and historically significant to Buddhism. Medically, based on some historical facts, the fig tree was used to treat ailments and disorders including asthma, diabetes, diarrhea, epilepsy, gastric problems, and inflammatory, infectious, and sexual disorders (http://selectree.calpoly.edu/tree-detail/612). According to the record, the oldest and largest Bodhi tree in Thailand is located in the eastern part of the country in Wat Ton Pho Si Maha Pho, Tambon Khok Pip, Prachin Buri (https://www.tourismthailand.org/Attraction/ton-pho-si-maha-pho-the-great-badhitree). Most Thai people if not all according to the Redhitree hely and an according to the second the Redhitree hely and an according to the record.

bodhitree). Most Thai people, if not all, considered the Bodhi tree holy and an essential part of their culture and identity.

2.6 Reports on diseases of Bodhi Tree caused by various pathogens

According to a report by Schrader (2020), the common causal agents of Bodhi tree diseases are ascomycetes, and these pathogens can cause many small black spots on the leaf surface with different shapes of the bulge. As the disease advances, the Bodhi tree will eventually die of withered leaves. As of this time, the only way to manage the disease is by cutting off and burning the diseased leaves as soon as symptoms appear.

Like any other plant, the Bodhi tree is not exempt from different thriving diseases. Abeygunawardhane (1969) reported the leaf spot disease in the Bodhi tree, which is caused by *Glomerella cingulata*, and the brown root disease caused by *Phellinus noxius*, which was responsible for the mysterious death of individual Bodhi trees. Leaf blight disease was reported in Bodhi trees from Jaipur, India, and the causal agent was identified as *Phyllosticta* sp. (Sharma et al. 2011). Another leaf spot disease was reported from Lahore, Pakistan, caused by *Curvularia aeria* (Nayab & Akhtar 2016).

In 2007, a news report from Hindustan Times (2007), surfaced on the internet raising concern about the unknown disease of the sacred Bodhi tree in Bodh Gaya, northeastern India. The symptom includes hundreds of fresh leaves falling off daily which is quite unusual. It was reported that the cause of the incident was undernutrition and treatments were done but then no further details about the diagnosis were revealed and no further reports were given.

A study conducted by Li et al. (2022) in Zhanjiang, Guangdong, China reported the leaf spot disease on the Bodhi tree which was identified as caused by *Diaporthe tulliensis*. They describe the symptoms as circular to oval-shaped spots with pale white centers and brown-black edges surrounded by a chlorotic halo. Proper control management is needed to address this potential loss as repeated annual defoliation may weaken the tree and decrease its aesthetic value in the landscape.

2.7 Reports on diseases of Ficus spp. caused by Phyllachora spp.

Based on the recent data, there are 69 species of *Phyllachora* that are pathogenic to *Ficus* spp., but only 3 species are pathogenic to *Ficus* religiosa (USDA Fungal Databases; retrieved from https://nt.ars-grin.gov/fungaldatabases).

2.8 The genus Neophyllachora

In 2017, Dayarathne et al. introduced a new genus, *Neophyllachora* to accommodate the following: *Neophyllachora cerradensis*, *N. myrciae*, *N. myrciariae*, *N. subcircinans* and *N. trucantispora*, which are related to *Phyllachora* species but constitutes an independent strongly supported monophyletic clade within Phyllachoraceae. Some species of the aforesaid genus can badly infect some economically important crops which could lead to yield loss (Li et al. 2022).

2.9 Reports on diseases of Ficus spp. caused by Neophyllachora spp.

A study conducted by Tennakoon et. al (2021) is the first report on the occurrence of the Tar spot in *Ficus septica*, which is a promising topical herbal medicine to cure small cutaneous ulcers (Deli et al. 2022). The causal agent is named *Neophyllachora fici*. This study will also report one disease of *Ficus religiosa* caused by the genus *Neophyllachora*. Nonetheless, only a few significant studies about the aforementioned genera are available because of their biotrophic nature, which makes them unable to grow in culture, therefore, sequence data from fresh collections are difficult to obtain (Tamakaew et al. 2017).

CHAPTER 3

MATERIALS AND METHODS

3.1 Sample collection and disease symptoms

A survey of *Ficus religiosa* (Bodhi trees) with tar spot symptoms was carried out from September to December 2022. Twelve samples were taken from seven different locations in Chiang Mai Province, Thailand (Figure 1). Leaf samples were kept in plastic bags, labeled properly, and carried to the laboratory within 24 h of collection.



Figure 1. Chiang Mai Province map showing the location of the collected samples

3.2 Morphological studies

After being transferred to the laboratory, leaf samples were examined using a stereo microscope (Zeiss Stemi 305). Observations, photographs, measurements, and descriptions were made from squash mounts of fresh fruiting bodies and sections of the ascomata mounted in water and 95% lactic acid. Melzer's reagent was used to check the apical apparatus in the ascus. Photographs were made using a Nikon SMZ745T stereo microscope (×4.5) and ZEISS Scope A1 microscope (×40-100) and measurements were carried out using the Tarosoft (R) Image Frame Work program (Tarosoft, Bangkok, Thailand. Photoplates were prepared in Adobe Photoshop 2020 version 21.0.2 software.

3.3 Spore germination test

For the spore germination test, a small amount of the sticky-like fruiting bodies was taken from the fresh samples using a sterile scalpel and spread in a zigzag on Potato Dextrose Agar (PDA) plate. The PDA plate was then incubated at room temperature and was observed under the stereo microscope after 24 h to check for spore germination. Newly germinated spores were picked up from the incubated PDA plate using a sterile scalpel and transferred to new PDA and water agar (WA) plates. The plates were incubated at room temperature and were checked 3 times (after 24 h, 48 h, and 72 h) under the stereo microscope to see the fungal growth.

3.4 DNA extraction, PCR amplification, and sequencing

DNA was extracted from eight samples directly from ascomata using the DNA extraction kit (FAVORGEN, Ping-Tung, Taiwan), following the protocols in the manufacturer's instructions. Partial small subunit nuclear rDNA (SSU), large subunit (LSU), and internal transcribed spacer (ITS) genes were amplified using the primer pairs NS1 and NS4 (White et al. 1990), LROR and LR5 (Vilgalys & Hester 1990) and ITS4 and ITS5 (White et al. 1990) respectively. Amplifications were performed in 25 μ l of PCR mixtures containing 12.5 μ l PCR Master Mix, 9.5 μ l deionized water, 1 μ l of DNA template, and 1 μ l of each primer. The PCR thermal cycle program was performed with an initial step of 3 mins at 94°C, followed by 35 cycles of 30 sec at 94°C, 58 sec at 30°C, and 1 min at 72°C, with a final extension of 10 mins at 72°C. The amplifications were visualized by gel electrophoresis at 100V for 30 min. The PCR products were sequenced

and automatically determined in a genetic analyzer at the 1st Base Company (Kembangan, Malaysia) using the aforementioned PCR primers.

3.5 Taxon sampling

DNA sequences generated in this study were subjected to BLAST searches (NCBI) (https://www.ncbi.nlm.nih.gov) to select taxa for phylogenetic analyses. The sequences for the phylogenetic analyses were retrieved from GenBank following Tennakoon et al. (2021). The final combined alignment comprised 53 sequences including the new sequence.

3.6 Phylogenetic analyses and species identification

Multiple alignments were automatically made with MAFFT v. 7 (http://mafft.cbrc.jp/alignment/server), using default settings (Katoh & Standley 2013) and further edited manually by BioEdit v. 7.0.5.2. The ML phylogenetic tree was generated using the RAxML-HPC2 on XSEDE (8.2.8) (Stamatakis 2014) in the CIPRES Science Gateway platform (Miller et al. 2010) with 1000 separate runs. MrBayes v. 3.1.2 was used to perform Bayesian analysis (Huelsenbeck & Ronqvist 2001). GTR+I+G model was selected as the best-fit model for each gene using MrModeltest v. 2.3 (Nylander 2004) under the Akaike information criterion (AIC). Markov Chain Monte Carlo sampling (MCMC) was run for 5,000,000 generations and trees were sampled every 100th generation. The first 10% of trees that represented the burn-in phase were discarded and only the remaining 90% of trees were used for calculating posterior probabilities (PP) for the majority rule consensus tree. The resulting trees were drawn in FigTree v1.4.0 (Rambaut 2012), then edited in Microsoft PowerPoint (2013) and Adobe Photoshop CS6 version 10.0.

CHAPTER 4

RESULTS

4.1 Sample collection and disease symptoms

In this study, the leaf samples were collected from October to December 2022. In our survey of Chiang Mai Province, we found small and large black spots. We collected 12 samples (7 small black spots and 5 large spots) from 7 different locations (Figure 1.)

4.2 Morphological studies

Based on our observation, those leaf samples with large spots were found from strangler trees (Figure 2). Strangling is one of the characteristics of the tropical figs of the genus *Ficus* in the family *Moraceae*. This growth pattern upon host trees, which often results in the host's death, is common in tropical forests worldwide. Species of trees that possess the ability to strangle are called strangler figs or just stranglers (Britannica, 2019). Contrariwise, leaf samples with small spots were collected from non-strangler trees.

The symptom morphology of our collections (Figure 3) was found to fit with the generic concept of the Tar spot symptom in having the formation of black stromata, which is a slightly raised, semi-circular to irregular, dark brown to black lustrous structure as described by Hyde & Cannon (1999). However, the large spots are found to have craggy bulging structures and mostly have irregular sizes. Our samples are parasitic on leaves of *Ficus religiosa (Moraceae)*. Symptoms on the host appear as black, solitary to gregarious, mainly on the upper surface (Figure 4). The measurements and descriptions of the sexual morph characters and the asexual spores (conidia and spermatia) are shown in Tables 1 and 2, respectively. Photographs of both sexual morph and asexual morph characters are shown in Figures 5 and 6, respectively. In addition, all samples were stained with a drop of Melzer's reagent, and the asci were confirmed to be J- as they all did not stain blue (Figure 7).

sexual morph characters	Measurements	descriptions
Ascomata	$83-172 \times 127-349 \ \mu m$	perithecial, globose to subglobose, solitary or aggregates, ostiolate (conspicuous)
Peridium	21–30 thick	dark brown to black, Lateral part wider than basal part, compactly arranged strongly melanized cells
paraphyses	1–2 μm wide	filiform, numerous, persistent, septate, unbranched, longer than asci
Asci	$59-140 \times 14-25 \ \mu m$ ($\bar{x} = 88 \times 20 \ \mu m$, n = 30)	8-spored, unitunicate, persistent, cylindrical to fusiform, short pedicellate, walls uniform in thickness but not specially thickened at the apex and without visible apical structures
ascospores	Copyright [©] by Chiang Mai A I I r ights res $8-12 \times 5-9 \ \mu m$ $(\bar{x} = 13 \times 7 \ \mu m, n = 30)$	uniseriate to biseriate, overlapping, unicellular, hyaline, globose to elliptical, 1-2 guttulates, with a central concave depression, covered by a mucilaginous sheath, irregularly thickened sheath, 1–4.5 µm thickness

 Table 1. Sexual morph characters measurements and descriptions

12

asexual morph characters	Measurement	descriptions
conidiomata	Str.	visible conidiogenous cells, spermatial or conidial wide conidial locules
conidiogenous cells		holoblastic, covering the base of the conidiomata
conidia	$2.0-4.0 \times 3.5-6.0 \ \mu m$ ($\bar{x} = 3.33 \times 2.28 \ \mu m, n = 50$)	ellipsoidal, hyaline, rarely septate
spermatia a a a	$0.90-1.40 \times 7.0-16.0 \ \mu m$ ($\bar{x} = 1.03 \times 11.48 \ \mu m, n = 50$)	botuliform or falciform, narrowly rounded at both ends, curved, aseptate, hyaline, smooth-walled
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 Table 2. Asexual morph characters measurements descriptions



Figure 2. *Ficus religiosa*. a. strangler tree. b. non-stangler tree and Mai University



Figure 3. Symptom morphology of the 12 collected leaf samples from *Ficus religiosa*. a. large spots- samples 6 and 9-12. b. small spots- samples 1-5 and 7-8.



Figure 4. Ficus religiosa leaves. a. large spots. b. small spots



Figure 5. Sexual morph of *Neophyllachora* **sp.** (large and small spots). a. Tar spot on living leaves, b. Upper and lower leaves of with large spot, c. Upper and lower leaves of with small spot. D. Pseudostroma, e. Horizontal section through pseudostroma, f. Vertical section through pseudostroma, g. Vertical section through ascomata, h. Vertical section through peridium, i, j Asci and paraphyses. k. Paraphyses, l–s. Asci, t1–t7. Ascospores, u. Conidia, w. Germinated spores. Scale bars: $f = 1000 \,\mu\text{m}$, $g = 100 \,\mu\text{m}$, $h = 100 \,\mu\text{m}$, $k = 5 \,\mu\text{m}$, I, j, i–s= 50 μm . t1–t7, u, w= 10 μm .



Figure 6. Asexual morph of *Neophyllachora sp.* a. spermogonia. b. spermatia. c. conidiomata. d. conidiogenous cells.



Figure 7: Asci stained with Melzer reagent. a. small spots. b. large spots

4.3 Spore germination test

Spores germinated (Figure 8) after the PDA plates were incubated for 24 h at room temperature. Then, we attempted to isolate the fungus by transferring the newly germinated spores to PDA and WA plates which were then incubated at room temperature. However, there was no sign of any fungal growth after 24h, 48h, and 72 h of incubation.





Figure 8: Germinated spores. a-b. small spots. c-d. large spots. Copyright[©] by Chiang Mai University All rights reserved

4.4 Phylogenetic analysis

The PCR amplification for SSU and LSU gene regions failed following the condition referred to by Dayararthne et al. (2017). Thus, the phylogenetic analyses were conducted only using the ITS gene region. All *Neophyllachora* species lack the LSU gene and the SSU is available only for several species including *N. cerradensis*, *N. subcircinans* and *N. truncatispora* and the phylogenetic placement of our new sequences concurred with the combined multigene analysis. The newly generated sequences were cladded together with *Neophyllachora fici* with high bootstrap support (Figure 7). The base pair comparison between *Neophyllachora fici* and the newly obtained sequences shows more than 2% differences in the ITS region. In addition, the blast hits of all the new sequences showed closest to *Neophyllachora*. The final dataset comprised 65 strains including 7 new sequences with 525 aligned characters including gaps (Table 3.)

All genera in Phyllachoraceae are well-resolved except the generic type *Phyllachora* which has recovered as polyphyletic. The best scoring RAxML tree was selected to represent the relationships among the taxa, with the final ML optimization likelihood value of -8093.468208 (Figure 9). The parameters for the GTR+I+G model of ITS were as follows: estimated base frequencies; A-0.235449, C-0.269319, G-0.261529, T-0.233702, substitution rates AC-1.107447, AG-2.360770, AT-1.392100, CG-0.585365, CT-3.555061 and GT = 1.000000. The ML and Bayesian analyses both resulted in trees with similar topologies. Bayesian posterior probabilities from MCMC were evaluated with a final average standard deviation of split frequencies of 0.002338.

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Table 3. Taxa table

No.	Taxon	Strain	Genbank Accession (ITS)
1	Camarotella costaricensis	MM_149	KX451913
2	Camarotella costaricensis	MM_21	KX451900
3	Camarotella sp.	MM_27	KX451901
4	Coccodiella miconiae	ppMP1342	MF460365
5	Coccodiella miconiicola	CBMAP_H290A	MF460368
6	Coccodiella sp.	MM_165	KX451917
7	Neophyllachora cerradensis	UB15626	KC683454
8	Neophyllachora cerradensis	UB16014	KC683455
9	Neophyllachora cerradensis	UB21823	KC683470
10	Neophyllachora cerradensis	UB21908	KC683471
11	Neophyllachora fici	MFLU 19_2702	MW114384
12	Neophyllachora fici	NCYU 19_0326	MW114386
13	Neophyllachora myrciae	UB21292	KC683463
14	Neophyllachora myrciae	UB22192	KC683476
15	Neophyllachora myrciariae	UB21781	KC683469
16	Neophyllachora religiosa	CRC-H190	XXX
17	Neophyllachora religiosa	CRC-H191	XXX
18	Neophyllachora religiosa	CRC-H192	xxx
19	Neophyllachora religiosa	CRC-H193	xxx xxx
20	Neophyllachora religiosa	CRC-H194	a d xxx
21	Neophyllachora religiosa	CRC-H195	XXX
22	Neophyllachora religiosa	CRC-H196	XXX
23	Neophyllachora subcircinans	UB21747	KC683467
24	Neophyllachora subcircinans	UB09748	KC683441
25	Neophyllachora subcircinans	UB21238	KC683461
26	Neophyllachora subcircinans	UB21347	KC683466

Table 3. Taxa table (continued	Table 3.	Taxa	table ((continued)
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No.	Taxon	Strain	Genbank Accession (ITS)
27	Neophyllachora truncatispora	UB14083	KC683448
28	Phyllachora arthraxonis	MHYAU_072	MG269749
29	Phyllachora arundinellae	MHYAU_108	MG269761
30	Phyllachora capillipediicola	MHYAU_20089	KY498084
31	Phyllachora capillipediicola	MHYAU_20090	KY498115
32	Phyllachora chloridis	MFLU 15_0173	KY594026
33	Phyllachora chloridis-virgatae	MHYAU_20058	KY498102
34	Phyllachora chloridis-virgatae	MHYAU_20137	KY498092
35	Phyllachora cynodonticola	MFLU 16_2978	KY594025
36	Phyllachora cynodonticola	MFLU 16_2977	KY594024
37	Phyllachora cynodontis	MHYAU_20043	KY471329
38	Phyllachora cynodontis	MHYAU_20042	KY471328
39	Phyllachora flaccidudis	IFRD9445	ON075524
40	Phyllachora graminis	101486	AF257111
41	Phyllachora graminis	DAOM_2409	HQ317550
42	Phyllachora graminis	MM-166_P	KX451920
43	Phyllachora heterocladae	MFLU 18_1221	MK305902
44	Phyllachora imperatae	MHYAU_014	MG269746
45	Phyllachora indosasae	MHYAU_125	MG195637
46	Phyllachora isachnicola	MHYAU_179	MH018561
47	Phyllachora isachnicola	MHYAU_180_P	MH018562
48	Phyllachora jiaensis	IFRD9448	ON075527
49	Phyllachora keralensis	MHYAU_20082	KY498106
50	Phyllachora miscanthi	MHYAU_167	MG195644
51	Phyllachora miscanthi	MHYAU_157	MG195643
52	Phyllachora panicicola	MFLU16_2979	KY594028

No.	Taxon	Strain	Genbank Accession (ITS)	
53	Phyllachora pogonatheri	MHYAU_071	MG269748	
54	Phyllachora pogonatheri	MHYAU_070	MG269747	
55	Phyllachora sandiensis	IFRD9446	ON075525	
56	Phyllachora sinobambusae	MHYAU_085	MG195630	
57	Phyllachora sp.	MHYAU_123	MG195631	
58	Phyllachora sp.	MHYAU_158	MG195633	
59	Phyllachora sphaerocaryi	MHYAU_178	MH018560	
60	Phyllachora sphaerocaryi	MHYAU_217	MK614100	
61	Phyllachora virgataes	IFRD9447	ON075526	
62	Polystigma pusillum voucher	MM_113	KX451907	
63	Telimena bicincta	MM_108	KX451910	
64	Telimena canafistulae	MM_13	KX451906	
65	Telimena leeae	TH549	KX451934	

Table 3. Taxa table (continued)



ลิ<mark>ปสิทธิ์มหาวิทยาลัยเชียงใหม่</mark> Copyright[©] by Chiang Mai University All rights reserved



Figure 9. RAxML tree based on analyses of ITS sequence data. Bootstrap support values for ML equal or greater than 70%, and Bayesian posterior probabilities (BP) equal or greater than 0.95 are given as ML/BP above the nodes. The tree is rooted to *T. bicincta* (MM-108), *T. canafistulae* (MM-13) and *T. leeae* (TH549).

4.5 Taxonomy

Phyllachorales M.E. Barr

MycoBank number: MB 90495; Facesoffungi number: FoF 06410; Index Fungorum number: IF 90495

Notes: *Phyllachorales* was established by Barr (1983), and currently has three families, namely *Phaeochoraceae*, *Phyllachoraceae*, and *Telimenaceae* (Dayarathne et al. 2017; Mardones et al. 2017; Yang et al. 2019: Hyde et al. 2020). According to Parbery (1967) and Cannon (1991), *Phyllachorales* species are leaf- or steam-inhabiting microfungi with shiny black stromata and are morphologically characterized by deep black stromata of various shapes. This characteristic, however, is not seen in species of *Polystigma* which have brightly colored stromata. In addition, the perithecia of *Phyllachorales* are usually strongly melanized that may be superficial, erumpent, or immersed in the host tissue, have thin-walled paraphyses that frequently deliquesce, unitunicate asci of cylindrical to clavate shape, with an ascus crown and an inconspicuous apical ring not staining blue in iodine; and globose to filiform ascospores, which in most species are hyaline and 1-celled, with only a few genera including species with brown or septate ascospores.

Phyllachoraceae Theiss. & P. Syd.

MycoBank number: MB 81156; Facesoffungi number: FoF 01329; Index Fungorum number: IF 81156

Notes: *Phyllachoraceae* species have ascohymenial development with paraphyses and thin-walled asci. Some species may have an apical ring, that does not stain blue in iodine (J-) and ascospores that are often hyaline and 1-celled (Cannon 1991; Maharachchikumbura et al. 2015, 2016; Hyde et al. 2020). Members of this family are mostly reputed to be highly host-specific, and the majority of them lack sequence data (Dayarathne et al. 2017, Hyde et al. 2020). In 1915, Theissen and Sydow introduced *Phyllachoraceae* to accommodate *Phyllachora*.

Phyllachora Nitschke ex Fuckel

MycoBank number: MB 4049; Facesoffungi number: FoF 02126; Index Fungorum number: IF 4049

Type species – *Phyllachora graminis* (Pers.) Fuckel

Notes – *Phyllachora* species are named based on their host association (Cannon 1988). Their

clypeate pseudostroma in the leaf appears in various sizes from a subcuticular or intraepidermal to a generalized infection of the entire section of the mesophyll, inducing characteristic black shiny superficial symptoms (Dos Santos et al. 2016). According to Cannon (1991), the depth of ascomata is not a valid character to distinguish genera as it can be influenced by the consistency of the host.

Neophyllachora Dayar. & K.D. Hyde

MycoBank number: MB 553633; Facesoffungi number: FoF 13499; Index Fungorum number: IF 553633

Type species – Neophyllachora myrciae Dayar. & K.D. Hyde

Notes: *Neophyllachora myrciae*, which was previously known as *Dothidea myrciae* (Léveillé 1846) is the type species of *Neophyllachora*. Dayarathne & Hyde (2017) introduced *Neophyllachora* to accommodate this novel species. They described the members of this genus as having sub-epidermal, intra-epidermal stromata without a deeper invasion of mesophyll, and clavate asci. At present, there are 6 *Neophyllachora* species listed in Species Fungorum (accession date: 07.02.2023). Five of them were listed in 2017, namely *N. cerradensis* (Dayarathne & Hyde), *N. myrciae* (Dayarathne & Hyde), *N. myrciariae* (Dayarathne et al.), *N. subcircinans* (Dayarathne et al.), *N. truncatispora* (Dayarathne et al.), and *N. fici* (Tennakoon et al.) was added in 2021.



Figure 10. *Neophyllachora* **sp. (large and small spots**). a. Tar spot on living leaves, b. Upper and lower leaves of with large spot, c. Upper and lower leaves of with small spot. D. Pseudostroma, e. Horizontal section through pseudostroma, f. Vertical section through pseudostroma, g. Vertical section through ascomata, h. Vertical section through peridium, i, j Asci and paraphyses. k. Paraphyses, 1–s. Asci, t1–t7. Ascospores, u. Conidia, w. Germinated spores. Scale bars: $f = 1000 \mu m$, $g = 100 \mu m$, $h = 100 \mu m$, $k = 5 \mu m$, I, j, i–s= 50 μm . t1–t7, u, w= 10 μm .

Material examined: Thailand, Chiang Mai Province, Mae Wang District, Wat Ampharam, Ban Kad, on living leaves of *Ficus religiosa (Moraceae)*, 19 November 2022.



Figure 11. *Neophyllachora* sp. (sample 1). a. Host (*Ficus religiosa*), b. Upper and lower leaves with small spots, c. Pseudostroma, d. Horizontal section through pseudostroma, e. Vertical section through pseudostroma, f. Vertical section through peridium, g. Asci and paraphyses. h. Paraphyses, i–j. Asci, k-l. Ascospores. Scale bars: $e-f = 100 \mu m$, g, $i-j = 50 \mu m$, $h = 5 \mu m$, $k-l = 10 \mu m$.

Material examined: Thailand, Chiang Mai Province, Hang Dong District on living leaves of *Ficus religiosa (Moraceae*), 19 November 2022.



Figure 12. *Neophyllachora* **sp. (sample 2)**. a. Host (*Ficus religiosa*), b. Upper and lower leaves with small spots, c. Pseudostroma, d. Horizontal section through pseudostroma, e. Vertical section through pseudostroma, f. Vertical section through peridium, g. Asci and paraphyses. h. Paraphyses, i–j. Asci, k-l. Ascospores. Scale bars: $e-f = 100 \mu m$, g, $i-j = 50 \mu m$, $h = 5 \mu m$, $k-l = 10 \mu m$.

Material examined: Thailand, Chiang Mai Province, Hang Dong District on living leaves of *Ficus religiosa (Moraceae)*, 19 November 2022.



Figure 13. *Neophyllachora* **sp. (sample 3)**. a. Host (*Ficus religiosa*), b. Upper and lower leaves with small spots, c. Pseudostroma, d. Horizontal section through pseudostroma, e. Vertical section through pseudostroma, f. Vertical section through peridium, g. Asci and paraphyses. h. Paraphyses, i–j. Asci, k-l. Ascospores. Scale bars: $e-f = 100 \mu m$, g-h, $j = 50 \mu m$, $i = 5 \mu m$, $k-l = 10 \mu m$.

Material examined: Thailand, Chiang Mai Province, San Patong District on living leaves of *Ficus religiosa (Moraceae*), 19 November 2022.



Figure 14. *Neophyllachora* **sp. (sample 4)**. a. Host (*Ficus religiosa*), b. Upper and lower leaves with small spots, c. Pseudostroma, d. Horizontal section through pseudostroma, e. Vertical section through pseudostroma, f. Vertical section through peridium, g. Asci and paraphyses. h. Paraphyses, i–j. Asci, k-l. Ascospores. Scale bars: $e-f = 100 \mu m$, g-h, $j = 50 \mu m$, $i = 5 \mu m$, $k-l = 10 \mu m$.

Material examined: Thailand, Chiang Mai Province, Mae Wang District, Wat Ampharam, Ban Kad, on living leaves of *Ficus religiosa (Moraceae)*, 19 November 2022.



Figure 15. *Neophyllachora* sp. (sample 5). a. Host (*Ficus religiosa*), b. Upper and lower leaves with small spots, c. Pseudostroma, d. Horizontal section through pseudostroma, e. Vertical section through pseudostroma, f. Vertical section through peridium, g. Asci and paraphyses. h. Paraphyses, i–j. Asci, k-l. Ascospores. Scale bars: $e-f = 100 \mu m$, $g = 5 \mu m$, g, $h-j = 50 \mu m k-l = 10 \mu m$.

Material examined: Thailand, Chiang Mai Province, Mae Wang District, Wat Ampharam, Ban Kad, on living leaves of *Ficus religiosa (Moraceae)*, 19 November 2022.



Figure 16. *Neophyllachora* sp. (sample 6). a. Host (*Ficus religiosa*), b. Upper and lower leaves with small spots, c. Pseudostroma, d. Horizontal section through pseudostroma, e. Vertical section through pseudostroma, f. Vertical section through peridium, g. Asci and paraphyses. h. Paraphyses, i–j. Asci, k-l. Ascospores. Scale bars: $e-f = 100 \mu m$, $g = 5 \mu m$, g, $h-k = 50 \mu m$, $l-m = 10 \mu m$.

Material examined: Thailand, Chiang Mai Province, Mae Wang District, Wat Ampharam, Ban Kad, on living leaves of *Ficus religiosa (Moraceae)*, 19 November 2022.



Figure 17. *Neophyllachora* sp. (sample 7). a. Host (*Ficus religiosa*), b. Upper and lower leaves with small spots, c. Pseudostroma, d. Horizontal section through pseudostroma, e. Vertical section through pseudostroma, f. Vertical section through peridium, g. Asci and paraphyses. h. Paraphyses, i–j. Asci, k-l. Ascospores. Scale bars: $e-f = 100 \mu m$, $g = 5 \mu m$, g, $h-k = 50 \mu m$, $l-m = 10 \mu m$.

Material examined: Thailand, Chiang Mai Province, San Patong District on living leaves of *Ficus religiosa (Moraceae)*, 19 November 2022.



Figure 18. *Neophyllachora* **sp. (sample 8)**. a. Host (*Ficus religiosa*), b. Upper and lower leaves with small spots, c. Pseudostroma, d. Horizontal section through pseudostroma, e. Vertical section through pseudostroma, f. Vertical section through peridium, g. Asci and paraphyses. h. Paraphyses, i–j. Asci, k-l. Ascospores. Scale bars: $e-f = 100 \mu m$, $g = 5 \mu m$, g, $h-k = 50 \mu m$, $1-m = 10 \mu m$.

Material examined: Thailand, Chiang Mai Province, Hang Dong District on living leaves of *Ficus religiosa (Moraceae)*, 19 November 2022.



Figure 19. *Neophyllachora* sp. (sample 9). a. Host (*Ficus religiosa*), b. Upper and lower leaves with small spots, c. Pseudostroma, d. Horizontal section through pseudostroma, e. Vertical section through pseudostroma, f. Vertical section through peridium, g. Asci and paraphyses. h. Paraphyses, i–j. Asci, k-l. Ascospores. Scale bars: e-f = $100 \mu m$, g = $5 \mu m$, h-k = $50 \mu m$, l-m = $10 \mu m$.

Material examined: Thailand, Chiang Mai Province, San Sai District on living leaves of *Ficus religiosa (Moraceae)*, 19 November 2022.



Figure 20. *Neophyllachora* **sp. (sample 10)**. a. Host (*Ficus religiosa*), b. Upper and lower leaves with small spots, c. Pseudostroma, d. Horizontal section through pseudostroma, e. Vertical section through pseudostroma, f. Vertical section through peridium, g. Asci and paraphyses. h. Paraphyses, i–j. Asci, k-l. Ascospores. Scale bars: $e-f = 100 \mu m$, $g = 5 \mu m$, $h-k = 50 \mu m$, $l-m = 10 \mu m$.

Material examined: Thailand, Chiang Mai Province, San Sai District on living leaves of *Ficus religiosa (Moraceae)*, 19 November 2022.



Figure 21. *Neophyllachora* **sp.** (sample 11). a. Host (*Ficus religiosa*), b. Upper and lower leaves with small spots, c. Pseudostroma, d. Horizontal section through pseudostroma, e. Vertical section through pseudostroma, f. Vertical section through peridium, g. Asci and paraphyses. h. Paraphyses, i–j. Asci, k-l. Ascospores. Scale bars: $e-f = 100 \mu m$, $g = 5 \mu m$, $h-k = 50 \mu m$, $l-m = 10 \mu m$.

Material examined: Thailand, Chiang Mai Province, Mae Rim District on living leaves of *Ficus religiosa (Moraceae*), 19 November 2022.



Figure 22. *Neophyllachora* **sp.** (sample 12). a. Host (*Ficus religiosa*), b. Upper and lower leaves with small spots, c. Pseudostroma, d. Horizontal section through pseudostroma, e. Vertical section through pseudostroma, f. Vertical section through peridium, g. Asci and paraphyses. h. Paraphyses, i–j. Asci, k-l. Ascospores. Scale bars: $e-f = 100 \mu m$, $h = 5 \mu m$, $h-k = 50 \mu m$, $l-m = 10 \mu m$.

Material examined: Thailand, Chiang Mai Province, Mae Rim District on living leaves of *Ficus religiosa (Moraceae*), 19 November 2022.

CHAPTER 5

DISCUSSION AND CONCLUSION

In our survey around Chiang Mai Province, we found small and large black spots on the leaves of Bodhi tree. We collected 12 samples (7 small black spots and 5 large black spots) from seven different locations (Figure 1). Based on our observation, those leaf samples with large spots were found from strangler trees. Strangling is one of the characteristics of the tropical figs of the genus *Ficus* in the family *Moraceae*. This growth pattern upon host trees, which often results in the host's death, is common in tropical forests worldwide (Britannica, 2019). Contrariwise, leaf samples with small spots were collected from non-strangler trees. However, the large spots are found to have craggy bulging structures and mostly have irregular sizes.

In this study, we attempted to isolate the fungus by transferring the newly germinated spores to PDA and WA plates which were then incubated at room temperature. However, there was no sign of any fungal growth after 24 h, 48 h, and 72 h of incubation. We selected 8 samples for morpho-phylogenetic studies; 4 with small black spots and 4 with large black spots. The former showed the same morphological characteristics of the black spot in having black, abundant, scattered, raised, mostly rounded to oblong or elongated, epiphyllous and shiny, superficial pseudostromata, which is sometimes parallel with the leaf venation. The latter almost have the same morphological characteristics of the black spot as the former, except that they are bulky, bristly, and bigger in size. Spermatia was produced in spermogonia and was grown with ascomata in the same pseudostromata. Spermatia has also has been reported in several species of *Phyllachoraceae* including *Parberya arxii* (P.F. Cannon) C.A. Pearce & K.D. Hyde. (Pearce and Hyde 2001), Phyllachora maydis Maubl. (Monteiro et al. 2013) and Phyllachora heterocladae C.L. Yang, X.L. Xu & K.D. Hyde. (Yang et al. 2019). Pearce and Hyde (2001) reported filiform spemartia produced in Spermatogonia in Parberya arxii that was originally placed in Sphaerodothis (Sacc. & P. Syd.) Shear and referred these structures as andromorph. Previously, Hyde and Cannon (1999) described an anamorphic state of Sphaerodothis arengae (Racib.) Shear ex Theiss. & Syd., which included ellipsoidal α -conidia and filiform β -conidia. However, the filiform spermatia found in P. arxii differed from S. arengae but are similar to those found in many Phyllachora species, particularly those taxa on grasses. In Australian Phyllachora collections spermatogonia developed either within the stromatic development of the teleomorph, in small irregular shaped locules at the side or above the developing ascomata mainly on moraceous hosts (Pearce and Hyde 2006). Evidently, our species also reported from a moraceous host and the spermatial locule lies aside the developing ascomata. Neophyllachora myrciae also shows sexual, spermacial and conidial structures and the species was originally introduced as Dothidea myrciae Lév. However, the author mentioned both spermacial and conidial structures as asexual states (dos Santos et al. 2017). Nonetheless, the relationship between spermatial state and telemorphic state remains unresolved (Cannon 1991). Moreover, we are also unable to make a conclusion about the lifecycle of the new species as the sampling was done only for a particular period in both young and mature trees. Thus, additional taxon sampling with detailed study of each stage of disease symptoms are needed for further investigation.

All samples were stained with a drop of Melzer's reagent, and the asci were confirmed to be J- as they all did not stain blue. The new species shows close phylogenetic association to N. fici and both species are reported in the same host genus (Ficus) but with different locality (Thailand Vs. Taiwan). Morphologically, our new species differs from N. fici in the size of asci $(55-185 \times 11-26 \text{ vs. } 90-100 \times 15-19 \text{ } \mu\text{m})$ and the thickness of the peridium (15-40 vs. 20-25 µm), and mainly differs in the ascospore characteristics. Our new species possesses ascospores which covered with thick gelatinous sheath $(1-4.5 \mu m)$ whereas N. fici lacks sheath in the ascospores. Our new species further differs from the type N. myrciae in the smaller pseudostromata (2-3)vs. 3–6 µm), thinner paraphyses (1.5–2.5 vs. 2.5–4.5 µm), shape of asci (fusoid vs. cylindrical to fusiform), shape of ascospores (lunate vs. globose to elliptical) and the presence of sheath in the ascospores, arrangement of ascospores (1-2 seriate vs. biseriate to multiseriate), color of ascospores (hyaline to olivaceous vs. hyaline), host (Ficus religiosa vs. Myrcia sp.) and the distribution (Thailand vs. Brazil) but shares similar characteristics in the size of ascomata, asci and coelomycetes asexual state. N. thailandica shares similar ascospores characteristics to N. cerradensis in elliptic-oblong shape ascospores with gelatinous sheath. However, our new species differs in the size of ascospores $(8-15 \times 5-12 \text{ vs. } 15-22 \times 6-9 \mu\text{m})$, thickened gelatinous sheath with hyaline to olivaceous color ascospores while *N. cerradensis* has thin walled and hyaline ascospores (Table 4). In summary, our new species mainly differs from the extant species in the ascospores characteristics which has thick gelatinous sheath that are irregularly thickened, irregularly guttulate at the immature stage and a large gattulate present in the maturity, and the shape ranges from globose to elliptical and a central concave depression present in the globose shaped ascospores.



Neophyllachora species							
Morphological							
characters	N. cerradensis	N. fici	N. myrciae	N. myrciariae	N. subcircinans	N. truncatispora	N. religiosa
Pseudostromata (diam)							
(mm)	2–4	2–3	3–6	0.5-1.5	1-4	2–5	2–3
	Ampulliform	Globose to	Globose to	ile ?	20/1		Globose to
Shape of Ascomata	to Globose	Subglobose	Ampulliform	Ampulliform	Ampulliform	Globose to Ovoid	Subglobose
					Immersed in the	T 1.	
	Occasionally	D (1 1)	Occasionally	Immersed in the	pseudostromatic	Immersed in	F · 1 · 11
Position of Ascomata	coalescing	Epiphyllous	coalescent	pseudostromata	tissue	pseudostroma	Epiphyllous
Peridium thickness	17.01	20. 25	A	10.24	15 01		15 40
(µm)	1/-21	20-25	-	19-24	15-21	-	15-40
Size of Ascomata	260-362 ×	$150-300 \times 200-$	205–485 × 148–	$161-432 \times 126-$	$329-417 \times 193-$	1 (0, 200, 100, 250	150–300 × 200–
(µm)	168-264	400	207	267	275	$160-200 \times 100-350$	400
Size of Asci (µm)	69–92 × 17–28	90–100 × 15–19	89–117 × 13–19	63–90 × 11–15	73–100 × 12–19	69–117 × 14–25	$55-185 \times 11-26$
		Cylindrical to	$\langle A \rangle = \left\{ A \right\}$	ESSEL A	. //		Cylindrical to
Shape of Asci	Fusoid	Fusiform	Fusoid	Clavate-fusoid	Cylindrical	-	Fusiform
Width of Paraphyses			MA	RS'			
(µm)	1.6–3	1.5–2.5	2.5-4.5	2-3	2–3.5 µm	2.5–5	1.5-2.5
Septate of Paraphyses	Septate	Aseptate	Septate	Septate	Septate	Septate	Septate
Size of Ascospores		8	6	Y. d.	?!		
(µm)	$15-22 \times 6-9$	$12-13 \times 10-11$	$14-18 \times 5-7$	$14-19 \times 5-8$	$11-16 \times 7-9$	$18-26 \times 7-8$	$8-15 \times 5-12$
	Elliptic-	Globose to					Globose to
Shape of Ascospores	Oblong	Subglobose	Lunate	Elliptical	Oblong to ellipsoid	Sublunate to fusoid	elliptical
			a tom have		hyaline to light		Hyaline to
Color of Ascospores	Hyaline	Hyaline	Hyaline	Hyaline	olivaceous	Hyaline	Olivaceous
							Mostly
					Mostly uniseriate,		uniseriate,
Ascospores			Biseriate to		sometimes with		sometimes with
arrangement	Biseriate	1–2 seriate	multi-seriate	Obliquely biseriate	biseriate	-	biseriate

Table 4. Synopsis of Neophyllachora species
Table 4.	(continued)
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Neophyllachora species							
Morphological characters	N corradonsis	N fici	N myrciae	N myrciariae	N subcircinans	N truncatispora	N religiosa
	Covered by a thin gelatinous	Absont	Thin welled	Covered by a thin	Thin wall surrounded by a	Wall thickenings at	Covered by a thick gelatinous
Gattules	microgattulate cytoplasm	-	-	Irregularly guttulate	Centrally gattulate	-	Irregularly gattulate
Asexual morph	Coelomycete	Unknown	Coelomycete	Unknown	Unknown	Coelomycete	Coelomycete
Host	Leaves of <i>Myrcia torta</i>	Leaves of <i>Ficus</i> septica	Myrcia sp.	Leaves of Myrciaria delicatula	<i>Psidium</i> sp.	Leaves of Myrcia camapuanensis	Leaves of Ficus religiosa
Distribution	Brazil	Taiwan	Brazil	Brazil	Brazil	Brazil	Thailand
References	Dos Santos <i>et</i> <i>al.</i> 2016	Tennakoon <i>et al.</i> 2021	Dos Santos <i>et</i> <i>al</i> . 2016	Dos Santos <i>et al.</i> 2016	Dos Santos <i>et al.</i> 2016	Dos Santos <i>et al</i> . 2016	This study
MAI UNIVERSIT							



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Key to species of Neophyllachora

1. Parasitic on Ficus, Myrcia, and Myrciaria species	2
1'. Parasitic on Psidium species, ascospores thin-walled, short-ellipsoidal covered with thin-walled gelatinous	
sheath	cinans
2. Parasitic on <i>Ficus</i> , <i>Myrcia</i> species	3
2'. Parasitic on Myrciaria species, clavate-fusoid asci	ciariae
3. Parasitic on <i>Myrcia</i> species, fusoid asci, ascospores covered with gelatinous sheath	4
3'. Parasitic on Myrcia species, lunate-reniform to half-moon shape ascospores with thick	
walled	ispora
4. Elliptic-Oblong ascospores with microguttulate cytoplasm	ıdensis
4'. Lunate ascospores with thin-walled	iyrciae
5. Parasitic on Ficus, globose to ellipsoidal ascospores with gelatinous sheath	
Neophyllach	ora sp.
5'. Parasitic on <i>Ficus</i> , globose to subglobose ascospores without gelatinous sheath	
~	.N. fici

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