



***IDENTIFICATION OF A MYCORRHIZA ISOLATE FROM PAPHIOPEDILUM
BARBATUM (LINDL.) PFITZER FROM EX SITU LOCATION AND
DETERMINATION OF ITS GROWTH-ENHANCING CAPABILITIES***

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By

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IDENTIFICATION OF A MYCORRHIZA ISOLATE FROM *PAPHIOPEDILUM BARBATUM* (LINDL.) PFITZER FROM *EX SITU* LOCATION AND DETERMINATION OF ITS GROWTH-ENHANCING CAPABILITIES

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Orchids and mycorrhiza are known to have a symbiotic relationship that function particularly in seed germination and also in promoting overall plant growth. It is believed that a certain orchid mycorrhiza that functions to enhance plant growth also lives symbiotically in the roots of *Paphiopedilum barbatum*. Thus, the objectives of this research are to identify the mycorrhizal fungi that form a symbiosis with the terrestrial tropical orchid, *P. barbatum* and evaluate its capabilities to enhance growth in certain *in vitro* *Paphiopedilum* seedlings. Mycorrhizal fungi were isolated from the roots of *P. barbatum* collected from five different locations in Peninsular Malaysia. Isolation of fungi was carried out according to the Currah method for initial morphological identification and then through molecular identification using the internal transcribed spacer (ITS) region of nuclear ribosomal DNA of the isolates. The identified

mycorrhiza was then inoculated on *in vitro* selected *Paphiopedilum* seedlings on a double slanting media to observe mycorrhization and the occurrence of enhancement of plant growth. Only one mycorrhiza was isolated and identified; namely *Tulasnella calospora* (anamorph: *Rhizoctonia repens*) which is a ubiquitous orchid mycorrhiza. Phylogenetic analyses with sequences originating from different hosts from different countries indicated a close genetic relatedness of the sequences studied. *In vitro* inoculation of *T. calospora* on seedlings from the *Paphiopedilum* genus after incubation for 24 weeks showed a significant 1.1% increase in fresh weight in *P. rothschildianum* with 68% root mycorrhization. However, no significant growth was observed in the seedlings of *P. sanderianum*, *P. gigantifolium* x *P. rothschildianum* and *P. esquirolei* x *P. rothschildianum* when inoculated with *T. calospora* after incubation for 24 weeks even though each had 32%, 24% and 13% root mycorrhization. This preliminary finding present an insight into the orchid mycorrhiza of the *Paphiopedilum* but a more extensive study is needed to further explore this symbiotic relationship and furthermore contribute in developing a beneficial application whether commercially or for the conservation of this genus.

Abstrak thesis yang dikemukakan kepada Senat Universiti Putra Malaysia sebagai memenuhi keperluan untuk ijazah Master Sains

**PENGENALPASTIAN SEJENIS MIKORIZA TERPENCIL DARI
PAPHIOPEDILUM BARBATUM (LINDL.) PFITZER DARI LOKASI *EX SITU*
DAN PENENTUAN KEBOLEHAN UNTUK MENINGKATKAN
PERTUMBUHAN**

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Disember 2010

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Orkid dan mikoriza diketahui mempunyai hubungan simbiotik yang berfungsi terutamanya dalam percambahan biji benih dan juga merangsang keseluruhan pertumbuhan pokok. Di percayai bahawa terdapat mikoriza orkid tertentu membantu dalam pertumbuhan pokok yang hidup secara simbiotik di dalam akar *Paphiopedilum barbatum*. Maka, objektif kajian ini adalah untuk mengenalpasti mikoriza yang boleh membina hubungan simbiotik dengan orkid terrestrial *P. barbatum* dan menguji kebolehannya merangsangkan pertumbuhan dalam anak pokok tertentu dari genus *Paphiopedilum* secara *in vitro*. Kulat mikoriza telah dipencilkan dari akar *P. barbatum* yang dikumpulkan dari lima lokasi yang berbeza di Semenanjung Malaysia. Pemencilan telah dilakukan menggunakan kaedah Currah untuk mengenalpastian awal

secara morfologi dan kemudian menggunakan kaedah molekul dengan menggunakan bahagian Internal Transcribed Spacer (ITS) dari DNA ribosom nukleus kulat tersebut. Mikoriza yang telah dikenalpasti itu kemudiannya telah diinokulasi kepada anak pokok *Paphiopedilum* yang terpilih secara *in vitro* di atas media dua sendeng untuk memerhatikan mikorisasi dan kesannya ke atas pertumbuhan. Hanya satu mikoriza iaitu *Tulasnella calospora* (anamorf: *Rhizoctonia repens*) iaitu mikoriza orkid yang telah dipencilkan dan dikenalpasti. Analisis filogenetik bersama jujukan DNA yang diperolehi daripada perumah yang berbeza dan negara yang berbeza menunjukkan perkaitan genetic yang rapat antara jujukan yang telah diuji. Inokulasi secara *in vitro*, *T. calospora* ke atas anak pokok dari genus *Paphiopedilum* selepas inkubasi selama 24 minggu menunjukkan peningkatan signifikan dari segi berat basah sebanyak 1.1% bagi *P. rothschildianum* dengan 68% mikorisasi akar. Namun begitu, tiada peningkatan pertumbuhan yang signifikan dapat dilihat bagi anak pokok *P. sanderianum*, *P. gigantifolium* x *P. rothschildianum* dan *P. esquirolei* x *P. Rothschildianum* yang diinokulasi dengan *T. calospora* selepas diinkubasi selama 24 minggu walaupun masing-masing menunjukkan 32%, 24% dan 13% mikorisasi akar. Pertemuan awal ini memberi maklumat awal tentang mikoriza orkid dalam *Paphiopedilum* tetapi kajian yang lebih menyeluruh diperlukan untuk mendalami hubungan simbiotik ini dan seterusnya menyumbang dalam pembangunan aplikasi yang berguna sama ada secara komersial atau bagi tujuan konservasi genus ini.

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LIST OF ABBREVIATIONS

AG	Anastomosis Group
dDH ₂ O	Double distilled water
DNA	Deoxyribonucleic Acid
dNTP	Deoxynucleotide triphosphate
DSE	Dark Septate Endophytes
ECM	Ectomycorrhizas
EDTA	Ethylenediaminetetraacetic acid
HCl	Hydrochloric acid
ITS	Internal Transcribed Spacers
KCl	Potassium chloride
KOH	Potassium Hydroxide
MgCl ₂	Magnesium chloride
NaCl	Sodium chloride
OM	Orchid Mycorrhiza
PCR	Polymerase Chain Reaction
PDA	Potato Dextrose Agar
RAPD	Random Amplified Polymorphic DNA
rDNA	Ribosomal Deoxyribonucleic Acid
<i>s.l.</i>	<i>sensu lato</i>
TBE	Tris Borate EDTA
TE	Tris-EDTA
VAM	Vesicular Arbuscular Mycorrhizas

CHAPTER 1

INTRODUCTION

The roots of a plant is by far one of the most important structures for plant performance as the roots have direct contact with a substrate whether soil, water or other organic mediums as a source of nutrients. In view of that, roots are affected by a complex array of microorganisms that inhabit a variety of niches with their own function and purpose in the substrate. Generally, these functions include nitrogen fixation, phosphate solubilization, plant growth promotion, and biological control of plant pathogens.

This association between microorganisms and roots can be beneficial (water uptake, soil stabilization, growth promotion, N₂ fixation, biocontrol, antibiosis, symbiosis), harmful (infection, phytotoxicity) or neutral (nutrient flux, free enzyme release, attachment, allelopathy, competition); depending on soil conditions (Rao, 1993). However, the most beneficial microorganisms that associate with plants are soil inhabiting fungi that form mutualistic associations with the root of plants, referred to as mycorrhizas (Smith and Read, 1997).

According to Brundrett (2004), mycorrhiza is defined as a symbiotic association essential for one or both partners, between a fungus (specialized for life in soils and plants) and a root (or other substrate-contacting organ) of a living plant, that is primarily responsible for nutrient transfer. Mycorrhizas occur in a specialized plant organ such as the roots where intimate contact results from a synchronized plant-fungus development.

This obligate relationship is particularly important in orchids as most orchids whether terrestrial or epiphytic, are known to establish a mycorrhizal association in at least one stage in their life cycle (Harvais and Hadley, 1967; Dearnaley, 2007; Stewart and Kane, 2007) or in the entire life cycle of chlorophyll-deficient species (Rasmussen, 1995).

The distinctiveness of mycorrhiza symbiosis in orchids can be seen in the occurrence of fungal interaction with embryo cells of germinating orchid seeds as well as with the roots of seedlings and mature plants (Peterson *et al.*, 2004). Orchid seeds are fairly minute and lack the nutritional reserves available in other types of seeds. Therefore, in nature, most orchids require the availability of mycorrhizal fungi as a carbon source to initiate seed germination, facilitate protocorm development and seedling growth (Rasmussen, 1995; Rasmussen and Whigham, 2002; Dong *et al.*, 2008).

These orchid fungal symbionts are also known to produce cellulases and polyphenol oxidases, enzymes that enable the fungi to digest soil organic matter to simple sugars that may be utilized by both partners in the symbiosis (Peterson *et al.*, 2004). Because of this prevalent and imperative association, the term orchid mycorrhiza (OM) is used exclusively for the mycorrhiza that occurs in orchids.

Most orchid mycorrhizal fungi are Basidiomycetes and belong to the form genus *Rhizoctonia*, a diverse polyphyletic group of pathogens, endophytes, saprophytes and mycorrhizal fungi (Warcup, 1981; Rasmussen, 1995; Currah *et al.*, 1997). Even though these fungi are known to be pathogenic to a variety of plants, they may also form

beneficial symbiotic relationship with orchids, as can be seen by *R. solani*, an active crop plant pathogen. Identification of these fungi is difficult due to the common absence of sexual reproductive structures and inconsistent vegetative characters that are traditionally used in morphological identification.

As most of orchid mycobionts are discovered to be from the *Rhizoctonia s.l.*, certain mycelial characters including the size and shape of moniloid cells, the formation of sclerotia, enzyme activity and particularly the ultrastructures of the septal pore are exclusively used to distinguish the sterile mycelia but only to the generic level (Zelmer and Currah, 1995). Due to this, fungal systematics is rather incoherent with different naming in taxa for the anamorph (fungal form with vegetative structures) and teleomorph (fungal form which produces sexual reproductive structures) from that of the same species (Currah *et al.*, 1997).

The advance of molecular techniques has greatly facilitated the progress of identification of mycorrhizal fungi (McCormick *et al.*, 2004). Research in this field relies mainly on PCR amplification of ribosomal DNA (rDNA) sequences with nuclear encoded small subunit (17S/18S rDNA) being the most common but nuclear encoded rDNA internal transcribed sequences (ITS), mitochondrial small subunit rDNA (16S) and nuclear encoded large subunit (25S rDNA) are also utilized significantly for research purposes (Gardes and Bruns, 1993; Kristiansen *et al.*, 2001; 2004; Otero *et al.*, 2005; Dearnaley, 2007; Giuseppe and Bellusci, 2009; Wright *et al.*, 2010). Examples of use of molecular techniques for mycorrhizal research are the study by Taylor and

McCormick (2008) to characterize basidiomycetous orchid mycorrhizas using internal transcribed spacer primer (ITS), molecular identification of orchid mycorrhizal fungi using mitochondrial ribosomal large subunit DNA (Kristiansen *et al.*, 2004) and a more recent study by Shimura *et al.* (2009) using ITS for phylogenetic analyses. On the other hand, random amplified polymorphic DNAs (RAPDs) and microsatellites have been far less popular as molecular markers for mycorrhizal fungi (Barker and Larkan, 2002) as ITS can provide a higher degree of variation even between closely related species.

The understanding of orchid mycorrhiza remains fragmented despite increase in research over the past four to five decades and vast increase in information (Brundrett, 2002; McCormick *et al.*, 2004). Tropical orchid mycorrhiza in particular has long been neglected by researchers (Dixon *et al.*, 2003) as studies on orchid mycorrhizal fungi have long focused on orchids from temperate regions even though the majority of orchids are tropical (Rasmussen, 1995). Even less studies on orchid mycorrhizal fungi from Southeast Asia have been done with only known studies done by Ma *et al.* (2003), Kristiansen *et al.* (2004) and Athipunyakom *et al.* (2004).

With the worldwide threat of extinction faced by the flora and fauna of the planet, orchids are acknowledged to be among the most severely threatened. Orchid conservation efforts are furthermore complicated due to a complex combination of external and internal factors. The external factors are the commonly referred reasons believed to cause extinction, such as habitat destruction and commercial exploitation

while symbiotic relationships that are essential in the life of an orchid is held to be a more difficult internal factor.

The unique association between orchids and their mycorrhiza is thought to present the foremost limitations for orchid conservation. Orchid conservation practices currently incorporate *ex situ* efforts that focus on isolation of mycorrhizal fungi used for artificial germination of orchid seeds and later to supplement existing populations or introduced to new areas of suitable habitat other than *in situ* efforts such as managing existing orchid populations and assisted migration of orchid species to new sites (Swartz and Dixon, 2009). However, in view of this, orchid conservation efforts in Malaysia still lack the inclusion of a mycorrhizal approach as relatively few studies have ever been done to investigate the mycorrhizas of Malaysian orchids (Jutta, 2007).

Orchid species from the genus *Paphiopedilum* (Orchidaceae) are among the most widely cultivated and hybridized of the 912 orchid genera. Commonly referred as Slipper Orchids, the *Paphiopedilum* is native to South China, India, Southeast Asia and the Pacific Islands. They are highly popular in horticulture and a prized collection for orchid lovers because of the flower's distinctively atypical form and rarity in nature. However, mycorrhizal studies on the *Paphiopedilum* are surprisingly scarce even though the significance of such knowledge is beneficial for conservation and even commercial purposes of the genus. *Paphiopedilum barbatum* (Lindl.) Pfitzer is a relatively more common species in Malaysia but increasingly threatened. A preliminary study by Jutta (2007) discovered the presence of *Rhizoctonia* spp. which is a known

OM when isolating endophytes from the roots of *P. barbatum*. This leads to the hypothesis that certain orchid mycorrhizal fungi that are beneficial for plant growth live symbiotically in the roots of *P. barbatum*. Thus, this study aims to isolate and identify the mycorrhizal fungi associated with *P. barbatum*, a relatively widespread Malaysian species in the *Paphiopedilum* genus and furthermore assess its growth enhancing capabilities with *Paphiopedilum* plant materials *in vitro*.

1.1 Objectives:

The objectives of this study are:

1. To isolate and identify (via molecular techniques) mycorrhizal fungi in *P. barbatum*.
2. To evaluate the growth performance of *in vitro* *Paphiopedilum* plant materials inoculated with the isolated mycorrhiza.

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