Assessing the invasiveness of alien aroids using modelling techniques and ecological assessments

by

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Westville campus



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As the candidate's supervisor I ha	ave approved this thesis for	submission.
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Abstract

Biological invasions represent one of the main drivers of the present decline in biodiversity worldwide and are difficult and costly to control. Consequently, identifying which factors allow a small proportion of species to successfully invade is a key area of research in invasion biology and is essential for effective management. In this thesis, I studied invasion patterns of the Araceae family, explored some of their ecological drivers, and unravelled mechanistic relationships that caused species to become successful. There are several emerging generalizations in invasion biology, but often the factors determining invasiveness are group-specific. Therefore the primary aim of this thesis was to establish whether general patterns of invasion biology also applied to Araceae. At a global scale, I found that, similar to other plant families, species with large native ranges and those that have been widely introduced were more likely to become invasive. What is unique to the family is the great diversity of growth forms, some of which are more likely to become invasive than others. I identified nine lineages in the family that have a greater tendency to invasiveness and recommended a precautionary approach be taken for these clades. At a regional scale, I used Epipremnum aureum as my case study species, because of the detection of the species in the country, as well as knowledge on its invasive cogener. In the KwaZulu-Natal province of South Africa, I found 78 naturalized E. aureum populations and 321 cultivated populations, of which the naturalized populations covered nearly 3 hectares in total. Disturbance played a major role in facilitating invasions and species distribution models indicated that E. aureum has a high probability of expanding its current range. Due to the invasion threat of this species, I recommended that all plants outside cultivation be removed. Lastly, I assessed a unique case where a widely planted species, Monstera deliciosa, has not yet become a global invader. I explored whether introduction history drives invasiveness in the Monsteroideae subfamily. I found that long residence times and high propagule pressure facilitated invasiveness in this subfamily. This was followed by as a local scale approach to identify factors influencing invasion success. The naturalization of *Monstera deliciosa* was largely driven by anthropogenic effects in Limpopo, South Africa, despite the plants' occurence in suitable habitat. Therefore, I concluded that M. deliciosa poses a low

invasion risk to South Africa. Overall, this thesis demonstrated the importance of using a taxonomic group to identify the contribution of multiple factors in the success of invasive species, but that species-specific assessments will still be required for effective

management.

Preface

The work described in this thesis was carried out in the School of Sciences, University of KwaZulu-Natal, Westville campus from June 2013 to December 2016, under the supervision of Professor Şerban Procheş and co-supervision of Professor John R. Wilson.

I hereby declare that the entirety of the work contained in this thesis is my own, original work, that I am the sole author thereof (unless to the extent explicitly otherwise stated) and that I have not previously in its entirety or in part submitted it for obtaining any degree or diploma to any tertiary institution.

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Declaration 1 - Plagiarism

l,	declare that:
1.	The research reported in this thesis, except where otherwise indicated, is my original research.
2.	This thesis has not been submitted for any degree or examination at any other university.
3.	This thesis does not contain other persons' data, pictures, graphs or other information, unless specifically acknowledged as being sourced from other persons.
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Declaration 2 - Publications

Publication 1: Moodley, D., Procheş, Ş., Wilson, J.R.U., 2016. A global assessment of a large monocot family highlights the need for group-specific analyses of invasiveness. AoB PLANTS 8, 1–14. (Editor's choice)

Contribution of each author:

DM, SP & JRUW: Planning and discussion of the study.

DM: Database compilation, statistical analyses, led the writing.

SP & JRUW: Provided comments on the manuscript.

Publication 2: Moodley, D., Procheş, Ş., Wilson, J.R.U. (2017). Assessing and managing the threat posed by *Epipremnum aureum* in South Africa. South African Journal of Botany 109, 178–188

Contribution of each author:

DM, \$P & JRUW: Planning and discussion of the study.

DM: Data collection, statistical analyses, led the writing.

\$P & JRUW: Provided comments on the manuscript.

Poem - Entropy

Sunlight raced towards Earth's embrace,
and in that twilit moment,
love seemed to fill the air.
peace prevailed as sunlight unveiled
the magnificence of mother nature.
life received a borrowed grandeur
sunlight was the enlightening lender.
have you ever seen such a beauty?

as the moment passed, entropy!
the gradual decline into disorder,
with humans circumventing the natural borders
causing unnatural movements of fauna and flora
to regions near and far, opening up pandora.
the sunlight bore witness to such calamities!
silent cries of dying trees filled the air,
leaving mother nature in much despair!

the hands of time sing a song with a constant rhyme
each hand takes away a moment, and never gives it back again,
soon only echoes and silent cries of dying trees will be heard.
the bees will dance no more, the birds will never sing,
the flowers will cease to bloom, the forests will be doomed
wild and scenic rivers will become a remnant,
only then will man be repentant
and as sunlight becomes disasters spotlight,
life will never again receive a borrowed light!

now is the time to step up to the plate, to conserve Earth's natural habitats, before it's too late!

Acknowledgements

Time flies (especially when you are doing a PhD), but memories last forever. Now, at the end of this journey, it is probably the best time to thank all the people, who made the completion of my PhD possible by providing significant guidance, advice, and encouragement which culminated in a wonderful experience.

First and foremost, I am filled with gratitude to the indivisible invisible frequency (God) that always resonated words of confidence and faith; I offer my profound thankfulness unto He, the Guru perceptor, whom has always guided me; and my sincere thanks and appreciation to my beloved grandmother, mom and brother for their continuous support, patience, and confidence in me.

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For further details see the acknowledgement sections in chapters two and three.

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Chapter 1: Introduction

For millions of years the distribution of the global flora and fauna was restricted by natural barriers such as mountains, deserts, rivers, and oceans. Anthropogenic activities have circumvented these barriers and provided a conduit for some species to disperse much farther than they could naturally and subsequently increased the rate and spread of alien species (Hulme, 2009; Meyerson and Mooney, 2007b; Wilson *et al.*, 2009b). As such, biological invasions can broadly be defined by the expansion of a species' geographic range into novel regions. The study of invasion biology has allowed us to progress in our understanding of invasions and our ability to manage them.

Following species introductions, many of the alien species remain innocuous and some eventually die out. For plants, only a small proportion are able to produce self-sustaining populations, and an even smaller proportion are able to spread great distances from their point of introduction (Blackburn *et al.*, 2011a; Williamson, 1996). This small percentage of species has caused significant environmental, economic and social impacts (Levine and D'Antonio, 2003; Mack *et al.*, 2000; Pimentel, 2011). Accordingly, this phenomena has directed some of the key research questions in invasion biology, such as: 1) why do some species become invasive once introduced, while others either fail or remain innocuous; 2) why are some habitats more vulnerable to invasions than other habitats; and 3) how can the knowledge developed in response to these two questions be used to predict invasion patterns, as well as develop effective policies and prioritize the management of biological invasions.

Many hypotheses have been proposed to explain and predict biological invasions. For example, Baker (1965; 1974) discussed the ability of plants to reproduce both sexually and asexually, as well as adapt to environmental stress (i.e. phenotypic plasticity), while Goodwin *et al.* (1999) showed that invasion success was related to plant life form. In addition, higher specific leaf area (Gallagher *et al.*, 2015; Grotkopp and Rejmánek, 2007) and the capability to climb (Paul and Yavitt, 2011) also predispose plant species to invasiveness.

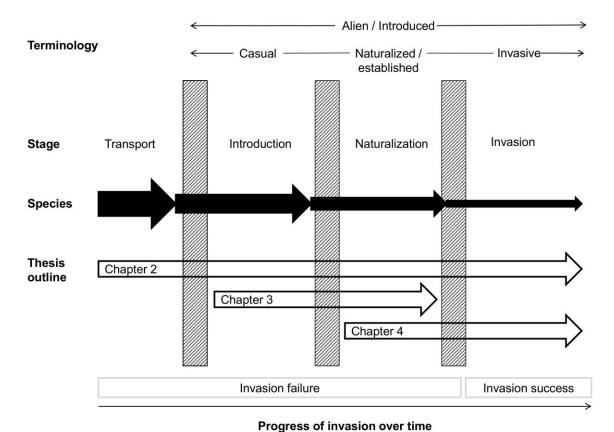
In the same way, not all habitats are equally vulnerable to invasion, and indeed habitats vary in their plant invasion level. Often, invaded habitats are commonly associated with anthropogenic disturbances, however, many invasive alien plants also thrive in natural habitats (Chytrý et al., 2008; Pyšek et al., 2010a; Pyšek et al., 2010b). Disturbance increases invasibility because it eliminates resident species, disrupts species interactions, and increases the amount of available resources such as space, light, water and nutrients, consequently reducing the effects of competition and creating opportunities which an alien species may be able to exploit (Davis et al., 2000; Shea and Chesson, 2002). In addition, favourable climatic factors and the availability of resources such as light, water, and fertile soil also predispose a habitat to invasion (Parepa et al., 2013; Richardson and Pyšek, 2006a; Thuiller et al., 2007).

Some species are, however, not immediately successful at invading. They remain at low numbers for many years (i.e. lag phase) before the population suddenly explodes, turning a quiescent alien species into an invader (Kowarik, 1995; Zenni and Nuñez, 2013). Lag phases can be associated with several factors, such as delayed introduction of a required mutualist (Richardson *et al.*, 2000b), the absence of disturbance events such as fire for fire-dependent species (Geerts *et al.*, 2013), genetic constraints (Ellstrand and Schierenbeck, 2000), climate change (Salo, 2004), and the build up to achieve significant propagule pressure. It is also possible that lag phases are an artefact of sampling effort over time (Cousens and Mortimer, 1995). Consequently, lag phases can vary greatly across species and are difficult to predict.

Numerous studies have shown that aspects of the species' introduction history are also important drivers of plant invasions and therefore should complement studies that are looking at invasiveness. In particular, a greater influx of propagules (i.e. propagule pressure) and a longer time since a species was first introduced into a region (i.e. residence time) often correlates well with successful invasions (Colautti *et al.*, 2006; Lockwood *et al.*, 2005b; Pyšek *et al.*, 2009b; Simberloff, 2009; Wilson *et al.*, 2007). Given these complex drivers of biological invasions, it is evident that understanding these context-dependent effects across species, habitats and spatial scales will provide important insights to develop successful management strategies and ultimately transform invasion biology into a predictive science.

Finally, in order to advance our understanding of invasion patterns and processes we need to consider two other important components for invasion biology research comprising taxonomy and the stages of invasion. Although a combination of traits (some of which have been cited above) has been shown to have high predictive power in identifying drivers of invasiveness, it has not yet been possible to generalize these factors across all but fairly narrow taxonomic groups. This highlights the context-dependent nature of invasions and suggests the need to study invasions using the appropriate taxonomic level. Model groups studied to date have provided important insights, and even though they share similar drivers of invasiveness (e.g. native range size), they also differ in their mechanism associated with invasion success. For instance, the level of susceptibility to the root rot fungus (Phytophthora) significantly influenced naturalization success in Proteaceae (Moodley et al., 2013); specific growth forms facilitate vegetative dispersal in Cactaceae which correlates with invasiveness (Novoa et al., 2015); long-distance seed dispersal drives invasiveness in Pinus (Richardson, 2006); and the ability of Australian acacias to form effective mutualisms with nitrogen-fixing bacteria and their long-lived seed banks are important for their invasion success (Gibson et al., 2011; Le Maitre *et al.*, 2011).

In addition, since the relative importance of factors influencing invasiveness vary at different stages along the introduction-naturalization-invasion continuum (Figure 1; Gravuer *et al.*, 2008; Lloret *et al.*, 2005a; Pyšek *et al.*, 2009b; Richardson and Pyšek, 2006a; Williamson, 2006), it is important to take into account how different traits play a role at different invasion stages, as failure to do so can obscure the results. Moreover, this integrative line of research (i.e. species traits, habitat characteristics, cognizance of lag phases, introduction history, and invasion stages) using taxonomic groups will provide a greater understanding of the context dependencies of biological invasions.



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Fig. 1.1. A simplified illustration of the invasion process as a series of distinct stages (i.e. introduction-naturalization-invasion) which a species must transition through in order to successfully invade (adapted from Blackburn et al., 2011a). Species statuses are defined according to the stage they occupy. Alien species refers to those that are intentionally introduced by humans only; casual are those alien species that do not form self-replacing populations thereby depending on repeated introductions; naturalized or established are those alien species that can form selfsustaining populations for at least 10 years for plants without direct human intervention, or despite human intervention; and invasive species are a subset of naturalized species that reproduce in large numbers, spread great distances from the initial point of introduction (i.e. > 100 m; < 50 years for taxa spreading by seeds and 6 m/3 years for taxa spreading vegetatively), and cause impacts to the economy, environment or health (Richardson et al., 2011b). Geographic barriers and survival rates are important for overcoming the introduction stage, whereas overcoming various barriers to reproduction is important for naturalization, and dispersal ability is an important barrier for invasion (Richardson et al., 2000a). These progressive stages are recognized as the introduction-naturalization-invasion (INI)

continnum. Intuitively, a species fails to become invasive if it fails to progress through the barriers at any stage of the invasion process. The stages investigated throughout the thesis are also outlined. The hatched bars represent the various barriers and the black arrows show that the proportion of species that progress from one stage to the next is less than the previous one.

Pinus L. (Richardson, 2006), Australian acacias Mill. (sensu lato, Castro-Díez *et al.*, 2011; Hui *et al.*, 2011; Richardson *et al.*, 2011a), and Proteaceae (Moodley *et al.*, 2013) are model groups of woody plants that have been well studied in invasion biology and generated useful insights on traits that are important for invasiveness. However, much less work has been done on herbaceous plants. For this thesis I chose to explore the Araceae Juss., or the aroid family, as I will argue that this family which contains many well-known species of horticultural importance provides an excellent study group for identifying determinants of species invasiveness and habitat invasibility in herbaceous plants.

THE ARACEAE

The Araceae is recognized as a diverse family of monocotyledonous herbs, the third largest monocotyledon family after orchids and grasses, and the seventh largest of all flowering plants (Mayo *et al.*, 1997; Nauheimer *et al.*, 2012). The uniqueness of this ancient family is evident by its diversity in life forms, morphology and anatomy (Bogner, 1987; Cabrera *et al.*, 2008; Croat, 1990; Cusimano *et al.*, 2011; Grayum, 1990). Species range from gigantic to tiny, from arborescent to floating plants, and they occupy terrestrial, epiphytic and aquatic habitats (although most species in the family are epiphytes and climbers) (Boyce and Croat, 2011 onwards).

The Araceae occur naturally on every continent except for Antarctica, however their distribution is predominantly tropical with two main centers of diversity, tropical Asia and tropical America (Grayum, 1990; Nauheimer *et al.*, 2012). Although the paleotropics comprise more genera, the neotropics comprise most of the total species resulting in regions such as America being extremely species rich (Mayo *et al.*, 1997). Ecologically, the Araceae is an important herbaceous family because of the species' dominance of the understory and inter-canopy herb layer (Croat, 1990; Nadkami *et al.*, 2001), which subsequently also serve as indicators of forest

condition. Conversely, the ability to dominate on the ground and in the sub- or intercanopy may also pre-adapt these species to become invasive. Furthermore, aroids have been used for many decades as a food source and for medicinal purposes, and they are amongst the most horticulturally important families (Acebey *et al.*, 2010; Bienz, 1980; Boyce and Croat, 2011 onwards; Croat, 1994; Kubitzki, 1998). Consequently, comparable to the above-mentioned model groups, many species have had a long history of introduction to regions outside their native ranges.

Because of the increasing interest in this family in horticulture, introduction pathways are increasing. Certain introduction pathways enhance the likelihood of invasive success by ensuring high propagule pressure (Wilson et al., 2009b). Since many Araceae species are popular in horticulture (e.g. ornamentals in gardens and inside homes, decoration in public spaces, and grown in aquariums) which is as an important pathway for invasive alien plants in general (Dehnen-Schmutz et al., 2007b; Reichard and White, 2001; Richardson and Rejmánek, 2011), this group may exhibit different invasion patterns compared to Pinus and Australian acacias. Pinus and Australian acacia species were mainly planted for forestry purposes worldwide. Although species introduced through horticulture and forestry are categorized under the same introduction pathway (i.e. the importation of a commodity), they differ in several aspects such as their cultivation, trade, propagule pressure, residence time, and probability of escape. Hence, these different human effects may result in different invasion patterns. Additionally, compared to *Pinus* and Australian acacias, only a few Araceae species are currently known to be invasive globally and some others are naturalized. Given these dynamics, important insights can be gleaned from seeking patterns and correlations from a group with large numbers of introduced species over large geographical areas. Consequently, the long history of widespread transfers and planting of Araceae in many parts of the world has created a natural global-scale experiment with many opportunities to explore different aspects of plant invasion biology. This makes the Araceae an excellent taxonomic group for uncovering invasion patterns and processes in herbaceous plants, and this thesis will therefore also supplement studies on model groups.

HYPOTHESES PROPOSED TO EXPLAIN THE INVASION SUCCESS IN ARACEAE

The global problem of biological invasions is multifaceted and therefore, as outlined above, there are several ecological attributes and hypotheses proposed to explain the success of invasive species (Catford *et al.*, 2009). Furthermore, a single theory cannot account for invasion success among all environments since this varies spatially and temporally. Therefore, in order to develop appropriate management plans for invasive species, it is necessary to develop a synoptic view of the dynamic processes involved in the invasion process. In the past, studies mostly focused on species invasiveness and habitat invasibility (Alpert *et al.*, 2000; Rejmánek *et al.*, 2005b; Richardson and Pyšek, 2006a). More recently, an increasing number of studies recognized the importance of quantifying the likelihood of invasion by also focusing on pathways (Essl *et al.*, 2015a; Kumschick *et al.*, 2015a; McGeoch *et al.*, 2016; Puth and Post, 2005). This complementary approach of targeting all three aspects (species, sites and pathways) is necessary to facilitate reactive and proactive management, and develop effective invasion policies.

Species

Given the introduction history of Araceae, as well as a very invasive genus in the family (Lemnaceae), I expect that the tens rule, traits of an ideal weed, propagule pressure, and the residence time hypotheses will play a role in driving species invasiveness at broad and small spatial scales. The tens rule was proposed as the quantitative estimate of the proportion of introduced species becoming invasive (Williamson and Fitter, 1996). As such, I expect to find that 10% of Araceae will be introduced, of which 1 in 10 introduced species will become naturalized and that 1 in 10 of those naturalized species becomes invasive. Several studies have attempted to profile successful invaders by identifying which traits and attributes facilitate their invasion (Baker, 1965; Elton, 1958; Goodwin *et al.*, 1999; Grotkopp *et al.*, 2002; Pyšek *et al.*, 2003; Rejmánek *et al.*, 2005a; van Kleunen *et al.*, 2016). I will attempt to identify traits driving invasiveness within Araceae based on evidence of other successful invaders, as well as data availability.

Propagule pressure, both in space (by widespread distribution and abundant plantings) and/or temporally (by a long history of cultivation) increases the chance of

successful invasions (Lonsdale, 1999; Rouget and Richardson, 2003). Propagules include seeds, seedlings, adult plants and reproductive vegetative fragments. Due to the challenges of measuring propagule pressure, this frequently cited determinant is studied in many different forms in the invasion biology literature. These associated proxy variables encompass propagule abundance, propagule richness and propagule frequency to better understand the processes involved (Ricciardi et al., 2010). In the chapters that ensue I will use the number of introduced regions (i.e. propagule frequency) and the number of stems (i.e. propagule abundance) as proxies for propagule pressure. As a result, I predict that introduced Araceae with widespread dissemination and/or species planted in large numbers will have a higher probability of invasion. One of the most robust generalizations in invasion biology is that the probability of invasion increases with the time since introduction (i.e. residence time or time lag; Hulme, 2003; Kowarik, 1995; Pyšek and Jarošík, 2005). In addition, the importance of residence time is also associated with propagule pressure, because species that were introduced a long time ago are likely to have been introduced many times since their first introduction. In the case of Araceae, I expect that species present in their new ranges for a longer period of time will be naturalized and/or invasive since they had the opportunity to fulfil more life cycles and spread further. On the other hand, given the context dependency of invasions, I also expect to find that propagule pressure and residence time are dependent on optimal site conditions at a local scale.

Sites

The capacity of species to tolerate the abiotic conditions of the site it was introduced in is another predictor coinciding with successful establishment. Hypotheses attributing invasion success to environmental factors are often based on fluctuation in resource availability and anthropogenic or natural disturbances (Funk and Vitousek, 2007; Levine and D'Antonio, 1999; Seabloom *et al.*, 2003). Resource availability can fluctuate following an increase in resource supply (e.g. water, light and nutrients) which will facilitate invasion success in the new site as long as the alien species can outcompete the resident species (Blumenthal, 2006; Davis *et al.*, 2000). In addition, natural and anthropogenic disturbances increase resource levels which give alien species a better chance of success at survival and establishment (Mack *et al.*, 2000). Therefore, I predict that successful Araceae will comprise of

species that are in the right place (i.e. suitable environmental conditions) at the right time (i.e. have access to available resources and also occupy disturbed areas which increase resource levels and decrease competition).

Pathways

Recent studies have highlighted the significance between pathways of introduction and invasion success (Essl *et al.*, 2015a; Faulkner *et al.*, 2016; Perrings *et al.*, 2005; Pyšek *et al.*, 2011). Prioritizing pathways uses information on vectors and routes of introduced species, thereby ensuring pre-border management and regulation of high-risk invasion pathways, as well as post-border management and legislation (Carlton and Ruiz, 2005; Hulme *et al.*, 2008). Hulme *et al.* (2008) outlined six principal pathways which was later refined into categories by Essl *et al.* (2015a). These six principal pathway categories comprise intentional release, escape from containment, transport as contaminant, transport as a stowaway, spread through corridors, and spread through unaided natural dispersal. Because Araceae are mainly introduced for horticultural purposes, I postulate that intentional releases and escape from containment are the main pathways since these include ornamental species.

THESIS STRUCTURE

This thesis comprises five chapters, three of which have been written in manuscriptstyle and are intended for publication in peer-reviewed journals. The chapters are structured as follows:

Chapter 1 provides a general introduction to biological invasions and includes a background, some of the research conducted, gaps in our knowledge, and objectives of this thesis.

Studies have shown that the determinants of naturalization and invasion success vary across taxonomic groups, invasion stages, and spatial scales. In chapter 2, I explore this variation for the Araceae family, a group remarkable through its life form diversity and horticultural importance globally. To do this, I created a species inventory of the family, described the invasion status of all species, identified factors influencing introduction, naturalization and invasion success and tested whether this

varied for different life forms, and then predicted which species will become invasive in future.

Chapter 3 explores the invasion risk posed by *Epipremnum aureum* in South Africa, a species that was recently recognized as a potential invader in the country, but a well-known invader in tropical regions. Here, I demarcate the species' current distribution in South Africa, describe factors driving naturalization, identify potential areas suitable for spread that are currently unoccupied by the species, and finally provide management options.

In chapter 4, I examine a paradoxical case in invasion biology; a popular horticultural species (*Monstera deliciosa*) that has been grown in most of the warm countries of the world for many decades, however it is not yet a major invader. Hence, this species does not conform to the established invasion theories. To unravel the mechanisms and factors behind the dynamics of this poorly studied species, I first assess whether attributes related to introduction history influences the invasion status of the Monsteroideae subfamily globally, subsequently I quantify the invasion risk of Monsteroideae in South Africa, and finally, at a local scale (i.e. using the only known invasive population in the Limpopo province) I describe the potential conditions under which *M. deliciosa* can become naturalized.

Chapter 5 provides a synthesis of the results of the three research studies and highlights the knowledge added to invasion biology by assessing the practicality of using a taxonomic group to explain invasion successes and failures. I emphasize the effectiveness of this approach in revealing the mechanisms of both successful and failed invasions, and encourage future comparisons within taxonomic groups.

THESIS OBJECTIVES

While research to date has described a multitude of factors influencing biological invasions (Hui and Richardson, 2017; Kolar and Lodge, 2001; Pyšek and Richardson, 2007; Richardson and Pyšek, 2012; van Kleunen *et al.*, 2015; Wilson *et al.*, 2017), the conclusions that can be drawn are limited by the lack of taxonomic group studies that identify drivers affecting invasion successes and failures. Using Araceae as a test case, this thesis examines the role of drivers of invasiveness (i.e.

the inherent ability of a species to invade) and invasibility (i.e. the susceptibility of the habitat to being invaded) in relation to the different invasion stages, spatial scales and introduction pathways. This approach will also account for the taxonomic bias in Araceae regarding general invasion principles in an herbaceous plant group since the invasion biology aspect is currently understudied. The overall goal of my research is to conduct a global assessment of the invasion processes driving Araceae invasions and to examine these processes at multiple spatial scales (Fig. 1.2).

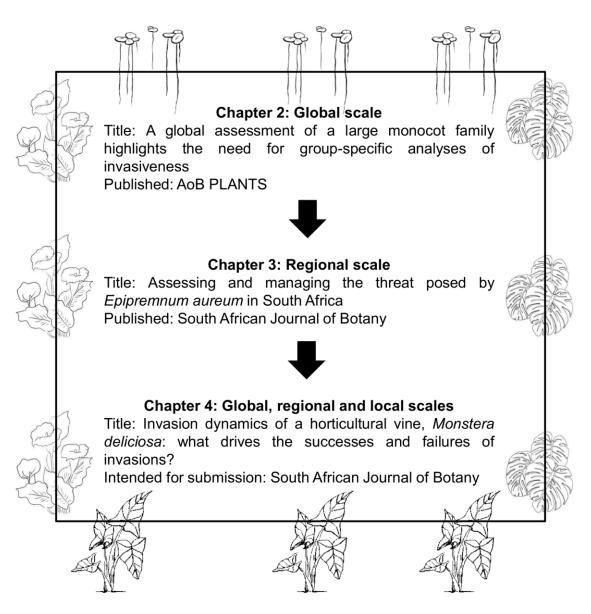


Fig. 1.2. The titles of each chapter, the associated spatial scales studied and the journal where it was published or submitted. The box encapsulating the chapters

symbolizes the unique taxonomic group used, which ranges from tiny floating aquatic plants to forest climbers and tuberous plants.

My specific research questions were:

- 1. What are the suites of factors underlying invasion success in Araceae?
- 2. Does *Epipremnum aureum* pose an invasion threat in South Africa?
- 3. Why is the world's most common house plant (*Monstera deliciosa*) not yet a major invader?

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Chapter 2: A global assessment of a large monocot family highlights the need for group-specific analyses of invasiveness

ABSTRACT

Significant progress has been made in understanding biological invasions recently and one of the key findings is that the determinants of naturalization and invasion success vary from group to group. Here we explore this variation for one of the largest plant families in the world, the Araceae. This group provides an excellent opportunity for identifying determinants of invasiveness in herbaceous plants, since it is one of the families most popular with horticulturalists, with species occupying various habitats and comprising many different life forms. We first developed a checklist of 3,494 species of Araceae using online databases and literature sources. We aimed to determine whether invasiveness across the introduction-naturalizationinvasion continuum is associated to particular traits within the family, and whether analyses focused on specific life-forms can reveal any mechanistic correlates. Boosted regression tree models were based on species invasion statuses as the response variables and traits associated with human use, biological characteristics, and distribution as the explanatory variables. The models indicate that biological traits such as plant life form and pollinator type are consistently strong correlates of invasiveness. Additionally, large scale drivers such as the number of native floristic regions and numbers of introduced regions are also influential at particular stages in the invasion continuum. We used these traits to build a phenogram showing groups defined by the similarity of characters. We identified nine groups that have a greater tendency to invasiveness (including Alocasia, the Lemnoideae and Epipremnum). From this we propose a list of species that are not currently invasive for which we would recommend a precautionary approach be taken. The successful management of plant invasions will depend on understanding such context-dependent effects across taxonomic groups, and across the different stages of the invasion process.

KEYWORDS: Araceae, biological invasions, boosted regression trees, invasiveness, predictions, stages of invasion, traits.

INTRODUCTION

Trade and transport of goods by humans have connected regions across the globe (Hulme, 2009; Pyšek *et al.*, 2010c). These pathways break down geographic barriers which results in thousands of species being introduced outside their native ranges (Pyšek *et al.*, 2011; Wilson *et al.*, 2009a). Of the introduced species, some are able to reproduce and form self-replacing populations to become naturalized but only a small subset progress to become invasive (Blackburn *et al.*, 2011b; Lockwood *et al.*, 2005a; Richardson *et al.*, 2000c; Williamson and Fitter, 1996). Identifying why some species become invasive in the introduced range while others do not, is one of the most important but challenging questions in invasion ecology. By improving our understanding of the drivers linked to biological invasions we can also develop better management practices and predict potential invasions.

The conceptualized invasion process comprises a series of barriers which a species must overcome to become naturalized and invasive in the introduced range (Blackburn et al., 2011b; Richardson et al., 2000c). A general understanding over the last several decades is that invasive species possess particular traits which allow them to overcome the invasion barriers in the introduced range. In the literature, species traits such as rapid growth rates and high reproductive output (Grotkopp and Rejmánek, 2007; Pyšek and Richardson, 2007; van Kleunen et al., 2010), as well as, their introduction history, such as high propagule pressure and a long residence time (Pyšek et al., 2009b; Simberloff, 2009) have been shown to be important determinants of invasiveness, but their relative importance varies across studies. The likelihood of invasiveness has also been predicted by attributes of the native range, such as large range sizes, and environmental similarity with the introduced range (Guisan and Thuiller, 2005; Hui et al., 2011). In addition, different traits become important at different stages of the invasion process (Richardson and Pyšek, 2012). For example, a large proportion of the alien plants have been introduced by humans over many years via the horticultural pathway, and this facilitates invasions through high propagule pressure and long residence times (Dehnen-Schmutz and Touza, 2008; Lambdon et al., 2008; Pyšek et al., 2009b).

Although there are several hypotheses explaining traits driving invasiveness, identifying a general suite of traits has proved difficult (Jeschke et al., 2012;

Richardson and Pyšek, 2006b). To date, empirical evidence shows that different sets of traits become important in different situations and the determinants of invasiveness are context-dependent (e.g. Funk, 2013; Moodley *et al.*, 2013; Prinzing *et al.*, 2002; Pyšek *et al.*, 2009a; Rejmánek, 1996; Thompson *et al.*, 1995; van Kleunen *et al.*, 2010). Furthermore, while some species perform better with the predicted invasive traits, it is not a feature shared by all invasive species (Alpert *et al.*, 2000; Lloret *et al.*, 2005b; Richardson and Pyšek, 2006b; Tecco *et al.*, 2010). One line of reasoning is that invasive species are associated with invasion syndromes. For example, invasion success may be specific to particular taxonomic groups, habitats or species life history traits (Kueffer *et al.*, 2013; Perkins and Nowak, 2013; Pyšek *et al.*, 2012). Therefore, instead of trying to identify general trends between invasive and non-invasive species across a wide range of taxa, it would be ideal to conduct in-depth case studies within taxonomic groups.

Araceae, also known as the arum or aroid family, is one of the oldest and the third largest monocotyledonous family in the world, after orchids and grasses (Mayo *et al.*, 1997; Nauheimer *et al.*, 2012). A unique feature of all species in this family is that their inflorescences consist of a spadix and a spathe (Chartier *et al.*, 2014). Aroids mostly occur in the tropics where they are concentrated in Southeast Asia, tropical America and the Malay Archipelago, and they comprise diverse life forms which occupy a wide range of habitats such as aquatic, terrestrial and ephiphytic (Cabrera *et al.*, 2008; Grayum, 1990; Mayo *et al.*, 1997). In addition, aroids have been used for decades as a food source, for medicinal purposes and in horticulture (Croat, 1994; Kubitzki, 1998; Mayo *et al.*, 1997). Given their large diversity and distribution, as well as their long history of introduction, Araceae serves as an excellent taxonomic group for identifying determinants of invasiveness in herbaceous plants.

In this study, we focused on introduction dynamics, characteristics of species' native ranges and biological traits to identify correlates of invasiveness within the Araceae family. Given that there are a variety of life-forms in Araceae, we hypothesized that when all species were analysed together, the only factors that would be significantly correlated to invasiveness would be factors seen to have a consistent influence across previously studied groups (e.g. native range size). However, repeating the analyses separately for different life forms would reveal specific mechanistic

correlates of invasiveness. Our objectives were therefore to: (1) create a species inventory using databases and literature sources, (2) describe the invasion status of all species, (3) identify which factors (native range characteristics, introduction dynamics and biological traits), influence introduction, naturalization and invasion success and whether this varied for different life forms, and (4) predict which species will become invasive in future.

METHODS

Global aroid database

Currently, there are no global databases listing all species belonging to Araceae. However, recent publications by Boyce and Croat (2011 onwards) provide the number of published and estimated species for each genus. This key resource gave us an initial idea of the aroid taxonomy. In order to create a comprehensive species inventory which includes data on accepted genera, species and synonyms, we surveyed a wide range of online databases (eMonocot, International aroid society, The Plant List, USDA Germplasm Resources Information Network, and World Checklist of Selected Plant Families). Given the large number of estimated and undescribed species in this family, it is likely there are aroid species that we did not include in our list.

Species status

The status of introduced, naturalized and invasive species is described in a wide variety of sources (e.g. on the internet, in published and unpublished literature). Since the criteria for defining naturalized and invasive species differ across studies, it is important to use reliable sources (Falk-Petersen *et al.*, 2006). We used multiple sources which contain a broad range of taxa, habitats and ecosystem types. This included 1) online databases (Atlas of living Australia, Calflora, Center for invasive species and ecosystem health, DAISIE, eMonocot, FloraBase, GBIF, GCW, GISD, HEAR, Invasive species of Japan, Randall (2007), and The PLANTS database), 2) published literature (New Zealand naturalized plant checklist), and 3) expert opinion (Haigh, A., Köster, N., Li, R., Seznec, G., Boyce, P., pers.comm.).

Determinants of invasiveness

Explanatory variables related to biological traits, biogeographical factors and human usage were selected to predict invasiveness (Table 2.1). We used these traits and factors because they were shown to be important drivers of invasiveness in other taxonomic groups such as Australian acacias (Castro-Díez *et al.*, 2011; Gibson *et al.*, 2011), Cactaceae (Novoa *et al.*, 2015), pines (Zenni and Simberloff, 2013), and Proteaceae (Moodley *et al.*, 2013). Binary response variables were categorized into three groups: non-introduced vs introduced (but not naturalized) species; introduced (but not naturalized) vs naturalized (but not invasive) species; and naturalized (but not invasive) vs invasive species. These groupings describe the stages that species need to successfully transition through to become invasive (Blackburn *et al.*, 2011b).

Table 2.1. Summary of traits used as explanatory variables in the analyses for identifying potential drivers of invasiveness in Araceae. The number of species is indicative of available data in terms of the listed traits (out of a total of 3,494 species). The range and median values for integer variables are shown in parentheses.

Trait	Levels	Number of species	Type of variable
Introduction dynamics	Use (food source; medicine; fibre production; horticulture; agroforestry; phytoremediation)	546	Categorical
	Total number of uses	546	Integer (1-5;1)
	Number of introduced regions (proxy for propagule pressure)	514	Integer (1-50;1)
Native range	34 floristic native regions classified according to Ronald Good (1974)	3490	Categorical, binary
	Total number of native regions (proxy for range size)	3490	Integer (1-31;1)
	Habitat (desert & xeric shrubland; mediterranean forests, woodland and scrub; temperate mixed forest; tropical dry forest; tropical moist forest)	3494	Categorical
Biological traits	Pollinator type (bees; beetles; flies; combination)	3250	Categorical

Flower sexuality (bisexual; unisexual)	3470	Categorical, binary
Regeneration mechanism (seed; vegetative; both)	444	Categorical
Life form chamaephyte; epiphyte; geophyte; helophyte; hemicryptophyte; hemiepiphyte; hyrdophyte; lithophyte; phanerophyte)	3426	Categorical

Statistical analyses

All analyses were performed in the R software version 2.15.1 (R Development Core Team, 2012). We used boosted regression trees (BRTs) to assess the relationship of the explanatory variables with the three transition stages, first using all species belonging to Araceae, followed by models developed for particular life forms. The BRT models were fitted using the 'gbm.step' function from the gbm package version 1.6-3.2 (Ridgeway, 2012).

Boosted regression trees are an advanced machine learning technique that applies an iterative method which sequentially builds multiple simple models, using the residuals from each subset of data during model fitting, to produce one ensemble model (Elith *et al.*, 2008; Friedman, 2001). This technique improves the models' predictive performance (Elith *et al.*, 2006). Among some of the advantages of this technique are that it can be fitted to a variety of response types (e.g. Gaussian, Poisson and binomial), it handles complex interactions between variables more efficiently than traditional methods (i.e. generalized linear models), it identifies important predictor variables, and it addresses issues like missing data and outliers (Elith *et al.*, 2008; Friedman, 2002).

Elith *et al.* (2008) provide details on selecting optimal settings for model fitting. These settings include the learning rate (shrinkage parameter that determines the contribution of each tree to the growing model) and tree complexity (specifies the number of nodes on each tree which controls whether interactions are fitted) which must be adjusted to produce a model comprising at least 1000 trees. BRT results include a measure of the comparative strength of association between the response variable and predictor variables (i.e. percentage deviance explained), and a cross-

validation coefficient (CV) which indicates the degree to which the model fits withheld data.

For this study, we first built preliminary models for each stage of the invasion continuum using all the predictor variables listed in Table 2.1 so that we could identify those with the greatest predictive contributions and reduce the overall number of variables in our analyses. The models were built with the default 10-fold cross-validation. The relative influence of predictor variables are determined by how often a variable was selected for splitting, weighted by the improvement of the models results (Elith *et al.*, 2008). From these results, we only kept predictors that contributed at least 5% to the models. From those, we performed a correlation test using Kendall's rank correlation to remove correlated variables ($r^2 > 0.65$); however all variables conformed to the correlation criterion. The models that were developed for particular life forms were only run for the introduction stage because of small datasets.

Boosted regression tree model calibration is prone to overfitting, and there are several ways to reduce this behaviour. A key approach of the model building process is to use validation processes which require a proportion of the dataset to be withheld. Here, cross-validation was performed using 75% of the data for training the model and the remaining 25% for testing. We used the caret package, version 6.0-24 (Kuhn, 2014), which creates random training and test sets while stratifying by the y variable. To evaluate model performance we used the average percentage deviance explained and the average cross-validation area under the receiver operating characteristic (AUC). Hosmer and Lemeshow (2000) state that an AUC value between 0.7-0.8 can be regarded as an acceptable model performance, 0.8-0.9 is excellent and higher than 0.9 is considered outstanding. A value of 0.5 or lower indicates predictions are worse than random. Due to the relatively low number of invasive Araceae, we could not fit training and testing datasets for the invasion model. Therefore, we only used 10-fold CV for model development and the crossvalidation AUC-value for evaluation (Elith et al., 2008). CV provides a means for testing the model on withheld portions of data, while still using the full data set at some stage to fit the model. The optimal parameter settings that were used in the final models are presented in Table 2.2.

Lastly, using predictors that met the BRT criteria (i.e. predictors that contributed at least 5% to the model) we either built generalized linear models with binomial errors, or used independent t-tests. This step provided insight into the individual explanation potential of each variable.

Table 2.2. Optimal parameter settings used in calibrating the boosted regression trees that produced the best performing introduction-naturalization-invasion models. To reduce overfitting, we used cross-validation which was performed by splitting 75% of the data for training the model and 25% for testing. We tested various learning rates (0.1–0.0005), bag 10 fractions (0.1–0.8) and levels of tree complexity (1–5). By trial-and-error we determined the most effective algorithm parameters for our dataset which is depicted below.

	Introduction Model	Naturalization Model	Invasion Model
Sample size (n):			
Full dataset	3,494	514	46
Training Data	2,621	386	_
Test Data	873	128	_
Parameters:			
Learning rate	0.001	0.001	0.001
Tree complexity	3	3	3
Bag Fraction	0.5	0.5	0.75

Predicting potentially invasive species

Using published literature, the first step was to examine the family tree and only select monophyletic groups. This selection controlled for phylogenetic effects as best as possible. Given that very few genera have published species level phylogenies, and most genera contain only non-introduced species, we decided to only include genera with known invasive species records. However, most of the invasive genera also lacked complete species level phylogenies. Selecting groups with invasive genera was important as it allowed inferring potentially invasive species in a more insightful manner (i.e. the selected groups comprised traits that are already known to confer invasiveness). In addition, it is assumed that species that have the potential to become invasive will be ones that 1) have relatives that are invasive, and 2) have

similar traits as invasive species. These two assumptions were used to formulate criteria to shortlist genera that have a known history of invasiveness.

Second, using the results from the BRT analyses, we scored species on traits that have already been shown to facilitate naturalization and invasion success in Araceae (Appendix 1). Following species scoring, we removed uninformative character states from the matrix. Finally, we constructed the phenograms using Jaccard's index and the unweighted pair group method with arithmetic mean (UPGMA) implemented in the Freetree software version 0.9.1.50 (Pavlicek *et al.*, 1999) which ranked species based on their overall similarity of characters. The phenograms clustered species based on the statistical similarity of their traits and also reflect evolutionary relatedness since only monophyletic groups were selected (see above). This allowed us to match species clusters with their associated invasion status. We used this approach as a tool to predict species that are not yet invasive but potentially pose an invasion risk.

RESULTS

Global aroid list

The Araceae database comprises 115 genera with 3,494 species worldwide (see Appendix 2), predominantly tropical in their distribution. Relatively few species 468 (13%) have been introduced (not yet naturalized or invasive) outside their native ranges, of which 27 (5% of the introduced species pool) species are classified as naturalized (not yet invasive) and 19 (4% of the introduced species pool) as invasive (Fig. 2.1A). Chamaephytes (Fig. 2.1B) and geophytes (Fig. 2.1C) contain the largest numbers of species, as well as, large proportions of introduced (not naturalized or invasive) species (11.98% and 17.34%, respectively), but they have low numbers of invasive species. Helophytes have the greatest proportion of introduced (not naturalized or invasive) species (23.86%) and also a relatively high proportion of naturalized (not invasive) and invasive species (Fig. 2.1D). Hemicryptophytes had 17.50% introduced (not naturalized or invasive) species, no naturalized (not invasive) species and 1% of the species are invasive (Fig. 2.1E). Hydrophytes seem to be the most successful with 13.15% introduced (not naturalized or invasive) species, 10.52% naturalized (not invasive) species and 13.15% invasive species (Fig. 2.1F). In contrast, phanerophytes have a large proportion of introduced (not

naturalized or invasive) species and naturalized (not invasive) species (16.67%) but no invasive species (Fig. 2.1g).

	a) All species	b) Chamae- phyte	c) Geophyte	d) Helophyte	e) Hemicryp- tophyte	f) Hydro- phyte	g) Phanero- phyte
Species Pool	3494	893	980	88	257	38	12
Introduced (not yet naturalized or invasive)	468	107	170	21	45	5	2
Naturalized (not yet invasive)	27 ↓ ↓ 19	6	11	1 1	0	4	2 0
	19	1	4	2	3	5	0

Fig. 2.1. Numbers of Araceae species at different stages along the introduction-naturalization-invasion continuum. The selected plant life forms that are depicted here tend to be introduced more often.

Model performance

The predictive performance of the models varied from acceptable (for the introduction and invasion model) to outstanding (for the naturalization model). The final BRT introduction model explained 13% of the mean total deviance (1 - mean residual deviance/mean total deviance). The test data AUC score was 0.72 and the full dataset cross-validation coefficient AUC score was 0.70 \pm 0.011 (mean \pm standard error). The naturalization model accounted for 59% of the total deviance and the test data AUC score was 0.98 while the cross-validation coefficient AUC score was 0.93 \pm 0.021. The invasion model accounted for 36% of the total deviance and the cross-validation coefficient AUC score was 0.74 \pm 0.072.

Factors associated with species' native range, introduction dynamics and biological traits in explaining INI success

The number of native floristic regions, which we used as a proxy for range size, was an important predictor for introduction (Table 2.3; Appendix 3). Species that occur over more floristic regions in their native range tend to be introduced more often (Fig. 2.2A; $F_{3.3490} = 46.7$, p < 0.001).

Table 2.3. Variables shown in the boosted regression tree analyses to have the greatest influence on the prediction of introduction, naturalization and invasion. The percentage contribution of a variable is based on the number of times the variable is selected for splitting, weighted by the squared improvement to the model as a result of each split, and averaged over all trees. For each model, the contribution of the variables is scaled to add up to 100%, with higher numbers indicating stronger influence on the response.

Model	Variable	Percentage contribution
Introduction	Number of native regions	30.00
	Life form	26.00
	Pollinator type	17.70
	Species native to Polynesia	9.90
	Flower sexuality	8.20
	Habitat	8.20
Naturalization	Number of introduced regions	65.90
	Life form	16.00
	Habitat	9.80
	Number of uses	8.30
Invasion	Life form	48.90
	Number of introduced regions	35.30
	Pollinator type	15.90

The number of introduced regions, which we used as a proxy for propagule pressure, was an important predictor of naturalization and invasion (Table 2.3). This suggests that species that are introduced to more regions in their new range tend to overcome the naturalization and invasion barriers (Fig. 2.2B; $F_{2.511} = 266$, p < 0.001).

Flower sexuality was significant for species overcoming the introduction barrier (Table 2.3). Relative to non-introduced species there are significantly more unisexual flowers among introduced species, but there are no significant differences across the naturalization and invasion stages (Fig. 2.2C; $F_{3,3466} = 11.29$, p < 0.001). Tropical climbers largely comprise species with unisexual flowers which explains why species with this flower type is likely to be introduced.

Data on the purpose of introduction were limited, as only 12% (n = 409) of the species had information on human usage. Nevertheless, we found number of uses to be an important predictor of naturalization (Table 2.3). Introduced species that had failed to naturalize tended to have fewer uses than naturalized and invasive species (Fig. 2.2D; $F_{2,406} = 53.55$, p < 0.001).

In comparison to other plant life forms, chamaephytes (z = -19.165; p < 0.001), geophytes (z = 3.587; p < 0.001), helophytes (z = 3.626; p < 0.001), hemicryptophytes (z = 2.386; p = 0.0170), hydrophytes (z = 3.940; p < 0.001), and phanerophytes (z = 1.980; p = 0.0477) have been introduced more frequently outside their native ranges. After introduction, hydrophytes (z = 4.870; p < 0.001) are the most successful in overcoming the naturalization barriers (Fig. 2.2E). These successful species are mainly used as ornamentals (including plants used in gardens, landscaping, cut flowers, aquariums and ponds) or as a food source. This demonstrates that horticulture provides a major pathway for plant invasions in Araceae. Even though life form was the most important factor across all stages (Table 2.3), we did not find a significant difference between the different life forms for the invasion stage. This can be attributed to the large number of naturalized species across the range of life forms that were able to become invasive.

The method of pollination was an important correlate for species introduction and invasion (Table 2.3). Species pollinated by bees (z = -7.930; p < 0.001) and flies (z = 3.149; p = 0.00164) were introduced more often. Although not significant, the combination of pollinators (z = 0.007; p > 0.05) and fly-pollinated (z = 0.007; p > 0.05) species are more invasive (Fig. 2.2F). Pollination by flies is typical of plants in the Araceae family. Fly-pollinated species being able to overcome the introduction barrier is probably an artefact of human use, since fly-pollinated species comprise

popular ornamental plants that are used for their unique inflorescences (e.g. *Amorphophallus*, *Anthurium*, *Arisaema* and *Zantedeschia*), decorative foliage (e.g. *Philodendron* and *Schismatoglottis*), or as aquarium plants (e.g. *Cryptocoryne*). Nevertheless, these pollinators highlight a specialized pollination syndrome in Araceae.

The type of habitat a species occupies in its native range was an important correlate of introduction and naturalization (Table 2.3). Although most of the species originating in desert and xeric shrublands are introduced (z = -2.587; p = 0.00969), they have not yet been recorded to naturalize or invade (Fig. 2.2G). Species native to humid regions, mediterranean forests (z = -3.569; p = 0.00289) and temperate mixed forests (z = -3.922; p < 0.001) in particular, tend to overcome the introduction and naturalization barriers.

From the thirty-four native floristic regions that Araceae occupy, species native to the Polynesian province were introduced more often (Table 2.3). While larger native floristic regions such as Malaysia and Euro-Siberia were more important in terms of the total number of invasive species originating there, Polynesia had the largest proportion of introduced species (64%), with 24% classified as naturalized and 12% as invasive.

Lastly, after incorporating particular life forms into the analyses, we did not find specific correlates of invasiveness that differed from the original model, therefore we rejected the second hypothesis. We found the number of introduced regions and reproductive characteristics to be important for chamaephytes; the number of native floristic regions, pollinator type, species native to West African rainforests and human use were important for epiphytes; and the number of introduced regions and reproductive characteristics were important for geophytes. In addition, we did not find a strong clustering in life forms across the family (see Appendix 4).

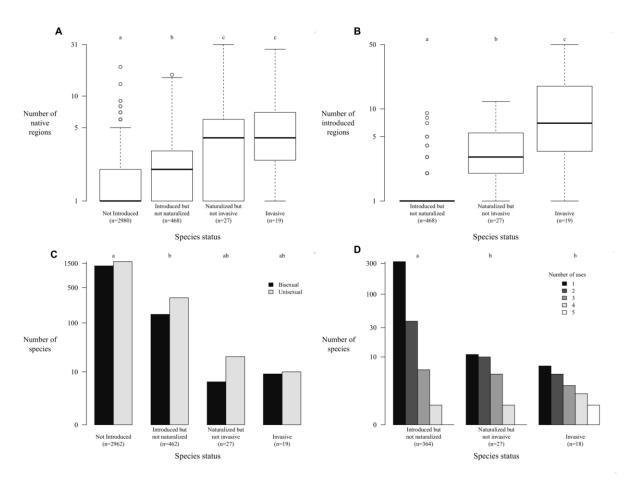


Fig. 2.2. The relationship between the introduction status of Araceae species and the parameters found to have a significant effect using boosted regression trees. A) Invasive taxa have larger native range sizes. Native range size is measured here in terms of the number of floristic regions based on Good's (1974) classification. Araceae naturally occur in 34 of the 37 floristic regions. B) Invasive species tend to have been introduced to more regions than naturalized species, and almost 90% of species which have been introduced to only one region have not yet naturalized. C) Species with unisexual flowers tend to have overcome more of the barriers to invasion than species with bisexual flowers. **D)** Species with a broad range of uses have naturalized and become invasive more often. Five different categories of human usage were considered: food source, medicine, fibre production, horticulture, agroforestry, and phytoremediation. E) Different life forms varied in their importance at different stages of the invasion. Introduced hydrophytes have naturalized far more than any other life form. F) Species that were fly-pollinated or had a combination of pollinator types became invasive compared to bee or beetle pollinated species. G) Species native to mediterranean and temperate mixed forests tend to naturalize more often. There were few data on the human uses of species that had not been

introduced outside their native range and so this category was excluded. In panels A and B, the box is the interquartile range, and the bold centre line is the median. Different letters denote different values using Tukey's multiple comparisons of means test. In panels E-G, tests were done using the original data, though the panels actually show plots of the fitted functions produced by boosted regression trees which indicate the effect on species presence/absence across the INI stages (y-axes) by each predictor variable (x-axes). For the relative contribution of each variable to the total deviance explained see Table 2.3. Grey panels indicate factors with low importance in the INI continuum, and therefore exclusion from the model.

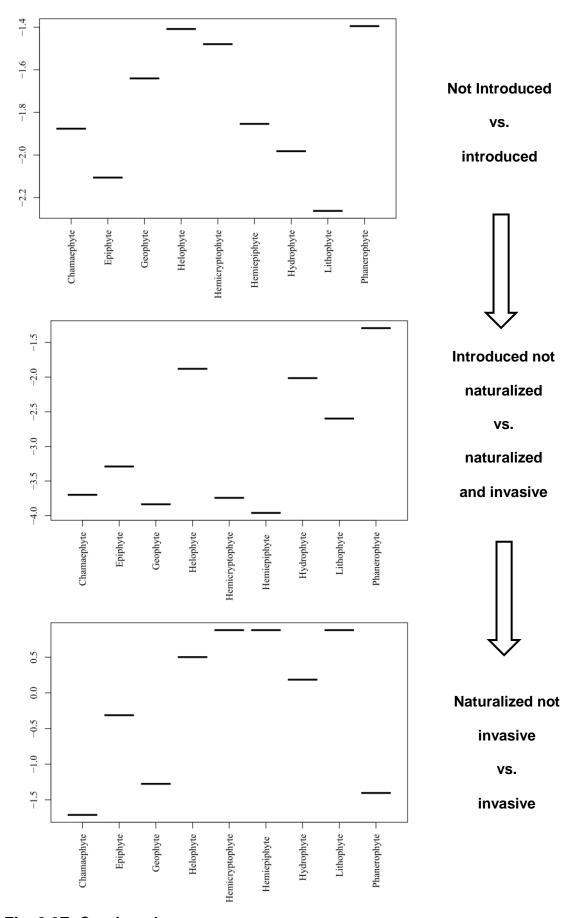


Fig. 2.2E. Continued

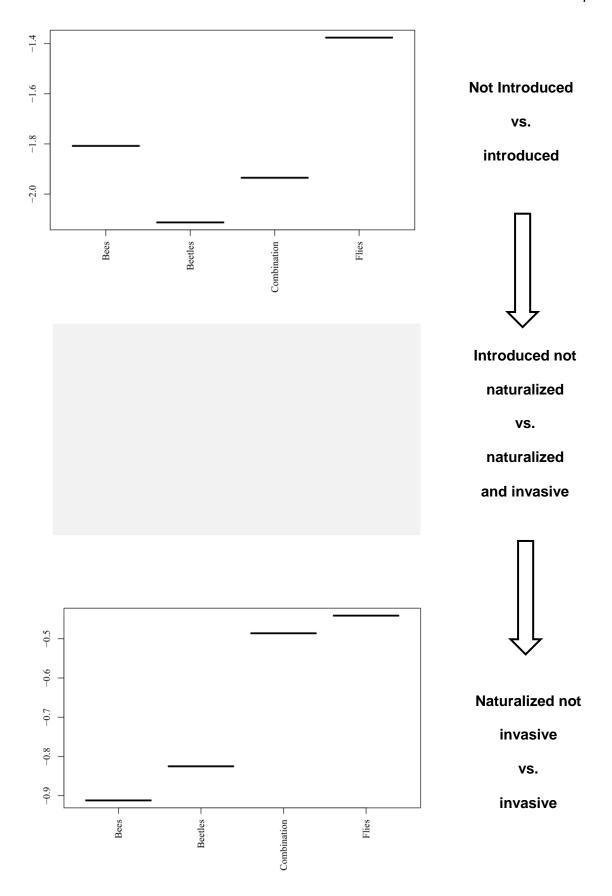


Fig. 2.2F. Continued

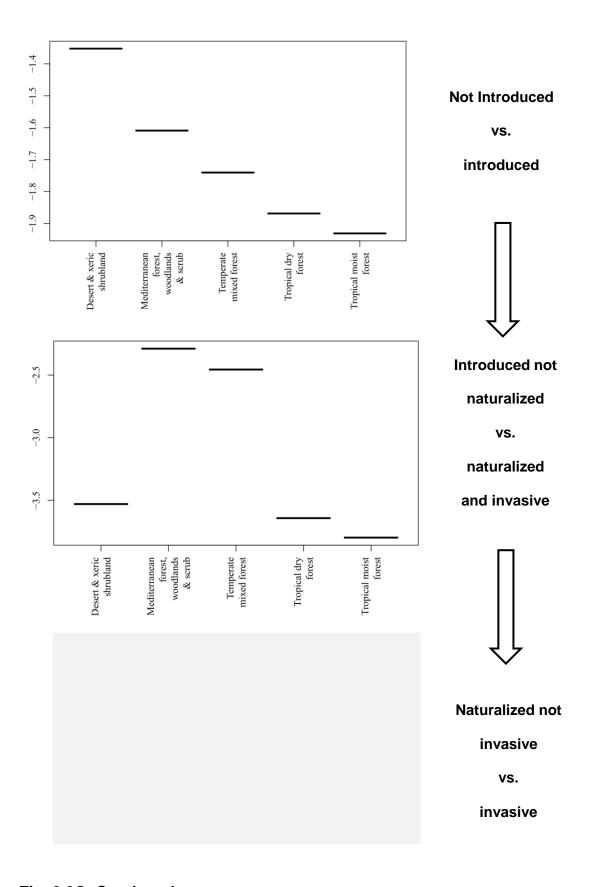


Fig. 2.2G. Continued

Predicting potentially invasive species

From the BRT models, we identified eight characteristics which facilitate species to overcome the INI barriers (see Appendix 1). Of the fifteen invasive genera in Araceae, we constructed phenograms inclusive of fourteen genera. The arguments used to identify potentially invasive species from the phenogram were based on 1) overall similarity in the character states of species, 2) whether species group with naturalized or invasive species, and 3) whether species cluster with naturalized or invasive sister groups. From the nine monophyletic groups, species with a high risk of becoming invasive are listed in Table 2.4 and their respective phenograms are illustrated in Appendix 5.

Table 2.4. A list of potentially invasive Araceae species constructed from model-based statistical inferences (i.e. UPGMA phenograms). These species are placed into groupings that are based on evolutionary relatedness (i.e. monophyletic groups) and similar ecological traits. Phenograms are illustrated in Appendix 5.

Monophyletic group	No. of species evaluated	No. of potentially high risk species	Potentially invasive species list [see Appendix 5]	Comments
Alocasia	77	5	Alocasia longiloba, Alocasia odora, Alocasia acuminata, Alocasia brisbanensis, Alocasia hypnosa	High likelihood for the listed non- introduced and introduced species to become invasive.
Amydrium, Anadendrum, Epipremnum, Monstera, Rhaphidophora, Scindapsus	82	38	Monstera adansonii var. adansonii, Monstera deliciosa, Anadendrum microstachyum, Anadendrum latifolium	Most species in this group are not yet introduced, however since this group already contains two invasive species, all species that are not listed requires further evaluation.

Ariopsis, Colocasia, Remusatia, Steudnera	20	11	Remusatia hookeriana, Remusatia pumila, Colocasia affinis	One cluster contains the invasive Colocasia esculenta, therefore, species in this group requires more attention.
Arophyton, Carlephyton, Colletogyne, Peltandra, Typhonodorum	2	1	Peltandra sagittifolia	Peltandra virginica is invasive and sister species Peltandra sagittifolia has been introduced outside its native range.
Arum, Biarum, Dracunculus, Eminium, Helicodiceros, Sauromatum, Theriophonum, Typhonium	55	23	Arum maculatum, Dracunculus vulgaris, Typhonium blumei, Typhonium roxburghii, Sauromatum venosum, Sauromatum horsfieldii, Typhonium trilobatum	Many species require further evaluation. Risk assessments must be conducted prior to species introductions.
Caladium, Chlorospatha, Filarum, Hapaline, Jasarum, Scaphispatha, Syngonium, Ulearum, Xanthosoma, Zomicarpa and Zomicarpella	169	~107	See clusters marked with asterisks in Appendix 5	Large group with five naturalized, but not invasive species and three invasive species scattered in the phenogram. All groups containing high risk species need to be evaluated further.

Cryptocoryne, Lagenandra	86	65	All species that clusters with invasive species	Phenogram shows very little structure (i.e. many species nested within groups) because fewer informative traits were used. Nevertheless, a single cluster contains the naturalized and invasive species. Therefore all species within this group pose an invasion risk.
Gymnostachys, Lysichiton, Orontium, Symplocarpus	8	6	Lysichiton camtschatcensis, Symplocarpus egorovii, Symplocarpus foetidus, Symplocarpus nabekuraensis, Symplocarpus nipponicus, Symplocarpus renifolius	High likelihood for non-introduced and introduced species to become invasive.
Lemna, Spirodela, Wolffia and Wolffiella	31	8	Lemna aequinoctialis, Lemna minor, Lemna perpusilla, Spirodela oligorrhiza, Wolffia arrhiza, Wolffia brasiliensis, Wolffiella lingulata, Wolffiella welwitschii	Many invasive species in this group. The listed non-invasive species have a high invasion risk because they cluster with the invasive species.

DISCUSSION

Identifying characteristics of successful invaders has been a major goal in invasion biology (Pyšek and Richardson, 2007; Rejmánek, 1996; Rejmánek and Richardson, 1996; Rejmánek *et al.*, 2005b; Richardson *et al.*, 2011a; Richardson and Pyšek,

2012). Although there are quite a few studies that have looked at traits in this family, these are mainly restricted to the most invasive group, the Duckweeds, and do not necessarily examine drivers of invasiveness. Some examples of studies conducted to date include; life traits and nutrient uptake of *Lemna minuta* and *Landoltia punctata* (Gérard and Triest, 2014); duckweeds as a valuable meal for domestic animals and fish due to their high protein content (Leng *et al.*, 1995); nutrient responses of *Lemna minuta* and *Lemna minor* to different nutrient availabilities (Paolaccia *et al.*, 2016); dispersal of the invasive *Lemna minuta* mediated by mallard ducks (Coughlan *et al.*, 2015); traits driving invasiveness of the first invasive *Colocasia esculenta* population in the Iberian Peninsula (García-de-Lomas *et al.*, 2012); and the phytochemical, pharmacological, medicinal, bioremediation potential, allelopathy, utilization and management of *Pistia stratiotes* (Khan *et al.*, 2014). Evidently, literature on the traits and mechanisms facilitating invasions in the Araceae family is resource poor, making this one of the most comprehensive studies to improve our understanding of invasion patterns in this family.

Our study supports the understanding that although some invasive traits are shared between invasive species, this is not consistent among all taxa and they are context-specific (Alpert *et al.*, 2000; Moodley *et al.*, 2013; Novoa *et al.*, 2015; Potgieter *et al.*, 2014; Richardson and Pyšek, 2006b; Theoharides and Dukes, 2007). Our main observations were that species that have large native floristic ranges are more likely to be introduced, and introduced species that are introduced to more regions are more likely to naturalize and invade; life form is consistently a major predictor; pollinator type might also be an important correlate and this is arguably specific to Araceae. Additionally, we found that particular traits or a combination of traits become important at different stages of the invasion continuum.

The importance of native range size (measured here in terms of the number of native floristic regions) is consistent with other studies (Hui *et al.*, 2011; Moodley *et al.*, 2013; Procheş *et al.*, 2012; Pyšek *et al.*, 2009a; Rejmánek, 1996), which also showed that species with larger native ranges are more likely to be introduced and become naturalized. A large native distribution is often correlated with invasiveness because there is a higher probability that wide ranging species will be picked up and intentionally or accidentally introduced (Blackburn and Jeschke, 2009). It is also

reflective of species being tolerant to a wide range of environmental conditions which pre-adapts them to survive and become established in the new region (Goodwin *et al.*, 1999; Pyšek *et al.*, 2009a).

High introduction efforts across novel ranges translate to a high propagule pressure. This finding is also in agreement with other studies (Colautti *et al.*, 2006; Moodley *et al.*, 2013; Pauchard and Shea, 2006; Zenni and Simberloff, 2013), where higher propagule pressure facilitates naturalization and invasion. This concept is based on the principle that species which are introduced across a wide area of the new region have a better chance of landing in localities that are suitable for establishment (Lockwood *et al.*, 2005a).

A large proportion of plant invasions result from horticultural introductions (Dehnen-Schmutz *et al.*, 2007b; Keller *et al.*, 2011; Reichard and White, 2001). Araceae are often used in horticulture, with hundreds of species and cultivars. Araceae that are used by humans for more purposes have a higher probability of being introduced and becoming naturalized. In addition, the invasion stage included species with the most number of uses. Other studies also found that species used by humans have a greater chance of becoming established in the introduced region because of a higher probability of being transported, and higher propagule pressure (Pyšek *et al.*, 2003; Thuiller *et al.*, 2006; van Kleunen *et al.*, 2007).

Plant life form is a common predictor of invasiveness for Araceae species since this trait is shared across the INI stages. This includes species (a) that are classified as hydrophytes; and (b) used for ornamental purposes. Araceae species that conform to these categories often reproduce vegetatively and this regeneration strategy is frequently linked to invasiveness (Kolar and Lodge, 2001). Although vegetative reproduction is not associated with long-distance spread, it can play an important role in the establishment of invasive species under suitable conditions in their new range (Daehler, 1998; Lloret et al., 2005b). Given that hydrophytes are more likely to overcome the introduction and naturalization barriers, species belonging to this life form pose a greater invasion risk. Furthermore, once species overcome the introduction and naturalization barriers, species of any life form have the potential to become invasive.

Ornamental species topped the list of invasive Araceae. It is well known that species deliberately introduced for ornamental purposes are associated with successful invasion because high market availability allows for high propagule pressure (Dehnen-Schmutz, 2011; Dehnen-Schmutz et al., 2007a). Species comprising invasive life forms with a potential for ornamental use should be carefully evaluated prior to introduction and management plans specific for these plants should be put in place. In addition, any species that is likely to be introduced with high propagule pressure poses a high risk and therefore efforts to reduce propagule pressure may successfully prevent a proportion of invasions. Propagules include seeds, seedlings, adult plants and reproductive vegetative fragments, and a high propagule pressure refers to propagules introduced or planted in large numbers, disseminated across a wide area, and/or with several introduction events. Efforts to reduce such pressures include regulating high-risk species that have not yet been introduced (i.e. pre-border management), as well as, regulating high invasion risk species that have already been introduced accompanied by management efforts that aim to eradicate, contain and control (i.e. post-border management). Additionally, control of propagules through effective management of the transport pathways and vectors (i.e. the importation of a commodity is the mechanism through which Araceae is introduced) will serve as a superior tool for the management and control of human-mediated biological introductions.

The main centres of origin and diversity of aroids are tropical regions such as tropical Asia and tropical America (Croat, 1998). However we found that species native to the Polynesian province were more successful in overcoming the introduction barriers. Forests in these Paleotropical regions are classified as one of the most wide-ranging and species-rich terrestrial habitats in the world (Whitmore, 1984) across taxa, and the Araceae are no exception. A higher introduction effort of wide-ranging species could be attributable to a higher abundance and tolerance to diverse conditions in any new area and so a relatively higher ease of cultivation (Dehnen-Schmutz *et al.*, 2007b; Forcella and Wood, 1984; Goodwin *et al.*, 1999; Prinzing *et al.*, 2002). In addition, since Polynesia is made up of islands, introduction effort from these islands is a key driver for Araceae dispersal. It is also possible that some of

these "native" Polynesian species were introduced by humans (and so pre-selected for an ability to be introduced), though this remains to be determined.

Some model groups demonstrate strong mechanistic correlation to invasion, such as Phytophthora susceptibility in Proteaceae (Moodley et al., 2013) and the many growth forms in Cactaceae which contain detachable vegetative propagules (Novoa et al., 2015). In Araceae we found that most correlates are universal. However, specialized pollinator types (e.g. flies and beetles) were important for introduction and invasion and this factor might be specific to Araceae. Most Araceae species are dependent on specialized pollinators (n = 900 beetles, n = 653 flies), and this may be limiting species that cannot spread vegetatively from becoming invasive. Species that require specialized pollination can encounter barriers to invasion when there is a lack of suitable pollinators or pollinator functional groups in their new range (Geerts and Pauw, 2012). The prevention and management of potentially high-risk species is required to help reduce the threats posed by invasive alien species. On one hand, there should be management plans put in place for species that are already introduced or species with a few naturalized populations, but which pose an invasion risk (e.g. prohibit further dissemination of potentially invasive species, remove highrisk species or issue permits for the possession of high-risk species, and consider attempting eradication or containment).

On the other hand, prevention is the best line of defence and can be applied to species that are not yet introduced but have similar traits to naturalized and invasive species. For instance, groups that so far lack invasive species may contain potentially invasive species which haven't been given an opportunity to invade. Therefore, phenograms should also be used for non-invasive groups that comprise species with the same suite of characteristics as the invasive groups. Screening high-risk species using a simple method based on evolutionary history and trait similarity is a conceptual step forward that provides a general framework in trying to predict invasiveness, however, this has ample room for improvement. In practice this will contribute towards the battle against invasive species, since risk assessment has its greatest impact when integrated into early invasive species management planning (Hulme, 2012; Wilson *et al.*, 2013).

CONCLUSION

Araceae conforms to some, but not all, of the emerging generalizations in invasion biology. As in other studies with other taxa, Araceae species that have been widely introduced (i.e. high propagule pressure) and which have large native range sizes are more likely to be invasive. However, unlike many other groups, there was little evidence of a link between invasiveness and regeneration mechanism (i.e. by seed, vegetative or both). Instead, there was a significant effect of plant life form and pollinator syndrome. Moreover, the importance of factors varied across the INI continuum.

Since the mechanisms associated with invasiveness differ between taxa and across the INI continuum, group and stage-specific analyses are required. As more complete phylogenies and better knowledge of traits become available, these analyses are likely to become increasingly sophisticated and able to produce valuable insights into risk assessments.

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Chapter 3: Assessing and managing the threat posed by *Epipremnum aureum* in South Africa

ABSTRACT

The predictive success of risk assessments is still largely a function of invasiveness elsewhere. Therefore, species that are invasive elsewhere should be prioritised for management, and where possible eradicated. We set out to investigate the threat posed by the alien climber Epipremnum aureum (Araceae) and assess techniques for controlling the spread of the species in South Africa. Epipremnum aureum is highly invasive in Hawaii and Sri Lanka, and has recently been considered as a potential invader in South Africa. However, no study has examined the invasion dynamics of the species. We mapped the species' current distribution in South Africa, modelled its potential distribution globally, and explored control methods. We only recorded the species in the KwaZulu-Natal province of South Africa, which comprised 78 naturalized populations and 321 cultivated populations. Delimitation surveys of the naturalized populations revealed ~187,000 plants over ~3 hectares. Several of these populations comprised plants as tall as the trees they were growing on, and were often found flourishing in dump sites, along roadsides or as a result of escaping cultivation. Species distribution models showed that *E. aureum* has a high probability of expanding its current range primarily along the coastal regions of South Africa and into neighbouring countries on Africa's eastern seaboard. Due to the invasion threat of the species, we recommend that all plants outside cultivation be removed. To achieve this, we found that applying herbicides to freshly cut stems significantly reduced plant growth. Given the species' limited dispersal ability and effective chemical control methods, we propose that *E. aureum* should be listed as category 3 under South Africa's National Environmental Management: Biodiversity Act (10/2004) Alien and Invasive Species regulations, i.e. naturalized populations need to be managed, it cannot be propagated or sold in future, but current garden plantings may remain.

Keywords: Biological invasions, climatic suitability, management, post-border risk assessment, South Africa

INTRODUCTION

Biological invasions are a global threat to agriculture, natural ecosystems, human and animal health, biodiversity and the economy (Davis, 2009; Drake *et al.*, 1989; Mazza *et al.*, 2014; McNeely, 2001a; Pimentel, 2011; Vitousek *et al.*, 1997). The mechanisms that underlie plant invasions are multifaceted, and for this reason no single predictor exists. Factors that facilitate the success of invasive alien plants include a combination of species invasiveness (i.e. intrinsic properties of a species), habitat invasibility (i.e. properties of a community that make it vulnerable or resistant), and the history of introduction (i.e. propagule pressure and residence time) (Lockwood *et al.*, 2005a; Nentwig, 2007; Richardson and Pyšek, 2006b; Wilson *et al.*, 2007). Understanding the conditions that facilitate biological invasions is a critical step in the prevention and management of invasions.

Understanding pathways of introductions is also important if species invasions are to be effectively regulated (Essl *et al.*, 2015b). Humans have both intentionally (e.g. import for horticulture) and unintentionally (e.g. as contaminants or stowaways) introduced species into new environments (Dehnen-Schmutz *et al.*, 2007b; Lambdon *et al.*, 2008; Mack and Lonsdale, 2001; Pyšek *et al.*, 2011; Reichard and White, 2001). With increased globalization, the number of introduced species has increased exponentially, and the number of pathways by which species may spread has also increased (Hulme, 2009; Hulme *et al.*, 2008; Pyšek *et al.*, 2011). The impact of biological invasions will inevitably continue increasing if management efforts against those species and pathways that pose the greatest threat are not prioritised (Rouget *et al.*, 2016).

Consequently, the development of country-level risk assessments, spanning the preborder, border and post-border stages, has become a high priority for managing invasive species. Weed risk assessments (WRA) represent a tool that is used to support the exclusion of potentially invasive alien species (IAS) from being introduced (i.e. pre-border screening), as well as assessing the potential impact of already introduced species that occupy different stages along the introductionnaturalization-invasion continuum (i.e. post-border prioritization and management) (Conser et al., 2015; Groves et al., 2001; Hulme, 2012; Pheloung et al., 1999a; Reichard and Hamilton, 1997; Weber and Gut, 2004). WRA tools are developed through classification of consistent patterns in traits of species that have previously become invasive. The Australian WRA (Pheloung *et al.*, 1999a) is widely recognised as one of the best systems to identify plant species, particularly terrestrial plant species, that are likely to become invasive and cause negative impacts. Several peer-reviewed papers have supported the accuracy of the Australian WRA system and recommend its wider application (Gordon *et al.*, 2008a; Hulme, 2012).

Once a species has been introduced and becomes established at a site (i.e. postborder), preventing its spread through containment or eradication becomes a priority. However, to fully understand the risks of IAS and to develop effective invasion policies and management, post-border weed risk assessments also need to take into account the context of the invasion. For example, introduction pathways (Hulme, 2009), species traits and their associated impacts (Blackburn *et al.*, 2014; Pyšek and Richardson, 2010), and the sites' susceptibility to invasion (Catford *et al.*, 2011), depict the plethora of conditions that can facilitate the success of IAS. As a result, the context for prioritization can vary widely across taxa, habitats and countries. Furthermore, delimiting the current geographic extent of the species and estimating their abundance are crucial steps towards understanding the likelihood of successfully implementing control and eradication plans (Panetta and Lawes, 2005; Wilson *et al.*, 2014). Consequently, standardized approaches for prioritizing pathways, sites and species impacts have recently been proposed (Dawson *et al.*, 2015; Essl *et al.*, 2015b; Kumschick *et al.*, 2015a; McGeoch *et al.*, 2016).

Araceae, one of the largest plant families, has been introduced globally through the horticultural pathway and currently contains 19 invasive species (Moodley *et al.*, 2016b). Of these, *Epipremnum aureum* has recently been detected in KwaZulu-Natal (KZN), South Africa (Sithole and Nzama, 2012). The global invasive status of *E. aureum* incited an evaluation of the species invasion risk and management in South Africa by the South African National Biodiversity Institute's Invasive Species Programme, which is responsible for detecting new invasions, conducting post-border risk assessments, and coordinating the eradication of high risk species that have a limited distribution (Wilson *et al.*, 2013).

Epipremnum aureum (Linden & André) G.S. Bunting, commonly referred to as devil's ivy, silver vine and golden pothos, is an evergreen epiphyte and a widely planted invasive species in subtropical and tropical climates around the world (Moodley *et al.*, 2016b; PIER). This species, a creeper and climber, is widely cultivated for ornamental use (i.e. as garden and indoor plants) because of its popular variegated foliage, reputation as a low maintenance plant and its efficiency in removing indoor pollutants such as formaldehyde and benzene (Dela Cruz *et al.*, 2014; Xu *et al.*, 2011). *E. aureum* reproduces easily from cuttings and detached pieces of stem, and is mainly dispersed vegetatively. The species rarely flowers in the wild, and in South Africa the species has not been seen in flower (pers.obs.).

The botanical classification of *E. aureum* has been controversial throughout its history. The species is sometimes considered a variety of *Epipremnum pinnatum* (L.) A. Engler, however Boyce (2004) identified *E. aureum* as a completely different species. There are quite a few distinctions between *E. pinnatum* and *E. aureum*; in *E. pinnatum* mature leaves are strongly pinnatifid, resembling the foliage of *Monstera deliciosa* Liebm. more than that of *E. aureum*; *E. pinnatum* has lanceolate to elliptic adult leaves while *E. aureum* has ovate to ovate-lanceolate leaves; and *E. pinnatum* flowers abundantly in the wild and in cultivation whereas *E. aureum* seldom flowers in the wild and flowering in cultivation is extremely rare (Boyce, 2004). *E. pinnatum* has not been recorded in South African herbaria, and based on morphology, the species was also not found during our field surveys.

While *E. pinnatum* is known to have a broad native range (i.e. temperate and tropical Asia, Australia, and the Pacific Islands; GISD), until 2004, there was uncertainty surrounding the origin of *E. aureum*. Finally, aroid botanist Peter Boyce established that the species was originally collected as *Epipremnum mooreense* Nadeaud (a synonym of *E. aureum*) from natural forest on the island of Moorea in French Polynesia (Boyce, 2004). Following the release of imported plants, *E. aureum* is now common in many countries including several Pacific Islands, the Caribbean, China, Hawaii, Central and South America, Southeast Asia, India, Pakistan and Singapore (eMonocot; HEAR; PIER).

One criterion commonly used to infer the status of a species as an invasion risk, is evidence that the species has a history of being invasive elsewhere, especially under similar climatic conditions (Faulkner *et al.*, 2014; Kolar and Lodge, 2001; Reichard and Hamilton, 1997; Thuiller *et al.*, 2005). *Epipremnum aureum* has a pan-tropical distribution and is regarded as a species of considerable concern because it has invaded several tropical and sub-tropical forests in Asia and the new world (Center for invasive species and ecosystem health; FLEPPC; PIER). Additionally, in Sri Lanka the species densely covers the forest floor and the trunks of trees, causing severe ecological disruption (Nyanatusita and Dissanayake, 2013). In South Africa, *E. aureum* was first detected in the Southbroom and Durban areas in the KZN province were it was observed to be invading forest margins (Sithole and Nzama, 2012). The identity of the species has been confirmed as *E. aureum* and specimens are lodged at the KZN herbarium (Appendix 6).

The purpose of this study is to determine the risk posed by *E. aureum* in South Africa. Specifically, we aim to: 1) delineate the current distribution of the species in South Africa; 2) describe factors that have contributed to its successful invasion; 3) identify the threat *E. aureum* poses in South Africa and globally using bioclimatic models, the Australian WRA, and notes on its behaviour in South Africa; 4) investigate best management practices to control the species; and 5) provide a recommendation as to whether regulation should be considered. This study is the first focussing on the invasion dynamics of *E. aureum*.

MATERIALS AND METHODS

Identifying study sites

Detection is important to ensure that new invasive species are promptly identified, reported, and contained or eradicated. A systematic search for the invasive species is the first component when managing a newly detected incursion (Wilson *et al.*, 2017). Initial localities of *E. aureum* were identified using the Southern African Plant Invaders Atlas (SAPIA) database in which there were 6 records (Henderson, 2007). To identify additional localities in the country, we contacted eight herbaria which cover five South African provinces (i.e. Bews, Bolus, Charles E. Moss, Compton, H.G.W.J. Schweickerdt, KwaZulu-Natal, Larry Leach and Selmar Schonland herbaria). As far as the species on record are concerned, *E. aureum* was only

recorded in KZN. For that reason, to determine if there are other populations in the province we distributed information flyers in KZN (Appendix 7). The flyers were targeted at spotter networks (i.e. groups of volunteers located in different towns or provinces in South Africa who identify and report invasive species) in the south coast area, however, many of the spotters often travelled throughout KZN. Spotters comprised members from the Pondoland custodians of rare and endangered wildflowers (CREW) group, members affiliated with the south coast conservancies, and residents.

Delimiting the extent of *E. aureum* populations in South Africa

Delimiting the spatial extent of the detected populations is necessary to determine the area to which the species has already spread (i.e. size of populations). Such delimitation surveys are also important for selecting appropriate control actions (e.g. eradication, containment, or no action) and assessing management feasibility (Hauser *et al.*, 2016; Moore *et al.*, 2011; Panetta and Lawes, 2005; Tobin *et al.*, 2013). Once a locality was confirmed through our detection efforts, we scanned the entire town using road surveys at a driving speed limited to 20km/hr. In addition, during our ad hoc drives through the nearby towns we detected populations that were previously not recorded. When we spotted *E. aureum*, we marked the locality with a handheld GPS (Garmin GPSMAP 64s). We also recorded whether the species was planted in a garden or in the wild (i.e. unmanaged area), and for populations growing in the wild, we collected information about the plants and the site (e.g. area, abundance, and disturbance).

Field assessments of wild populations

The extent of occurrence of invasive species is often used as proxy of success, and over time can be used as a measure of spread (Wilson *et al.*, 2014). In addition, understanding the occupied area inhabited by IAS will assist in guiding management strategies. We determined the extent of occurrence for each population by connecting the outlying GPS waypoints for each population and then calculating the enclosed area (m²), using the mapping software ExpertGPS 5.71.

Species abundance is another important indicator used to characterize the risk of IAS because it is associated with the impact of biological invasions (Kumschick et

al., 2015b). In addition, the number of stems is important for the expansion of E. aureum populations because the main mode of reproduction occurs vegetatively and seldom via seed. Consequently, since most populations comprised extremely dense plants and it was difficult to count all individuals, we measured abundance in terms of the number of stems in a population (i.e. each stem was considered as a single plant). To count the number of stems we laid three 1×1 m² transects, oriented perpendicular to the plant and its climbing structure, on different plant cover types (i.e. a representation from dense to sparse cover) and counted every stem within the transect. Plant abundance was then estimated using the average numbers of stems counted in a 1×1 m² perpendicular transect and multiplied by the populations extent of occurrence (m²).

A number of studies have shown that plant height is a consistent trait facilitating invasion success (Pyšek and Richardson, 2007). One explanation is that height is a major determinant of a plants' ability to compete for limiting resources, such as light (Moles et al., 2009; Westoby et al., 2002). For vines, the maximum height will, however, be limited by the height of the supporting structures on which they grow. Therefore, we estimated the height of the tallest *E. aureum* plant and the height of its associated support structure in order to demonstrate the competitive ability of the species. Height was estimated by one person holding a 2 m stick against the supporting structure and the other person standing 10 m away used their index finger and thumb to count the number of sticks from the ground to the top of the structure and to the top of the tallest *E. aureum* plants. If a vine can overtop a tree or shrub, then they can potentially have much greater impacts. Where the support structures were trees, the trees were grouped into size classes and the height of the tallest E. aureum plants relative to the height of the tree species they were growing on (one sample per population) were plotted on a boxplot and analysed using a generalized linear model with a Poisson distribution.

Lastly, we looked at land-use types where the study species occurred and their associated disturbance levels. Disturbance is commonly linked to increasing habitat invasibility (Alpert *et al.*, 2000; Pyšek *et al.*, 2010b; Rejmánek *et al.*, 2005b). Disturbance creates a window of opportunity during which IAS might benefit if they can reproduce and thrive better than the native species under such conditions (Davis

et al., 2000). For example, there is a strong correlation between plant invasions and disturbed habitats such as roadsides (Gelbard and Belnap, 2003; Meunier and Lavoie, 2012; Parendes and Jones, 2000). The land-use types at each site were initially categorized in the same way as the data collected for the SAPIA database but this was modified during field work. The following categories were used: canal, garden escapee (i.e. most of the population is still thriving in a garden but the species is in the early stages of spreading into the adjacent land), natural vegetation, roadside, transformed vegetation (i.e. sites dominated with alien plants), vacant land (i.e. land set aside for development which also includes vacant or abandoned properties), or wasteland (i.e. dump sites). We plotted the frequency of the number of populations and their land-use types, and analyzed the data using Pearson's Chisquared Test for Count Data. All statistical analyses were performed using R Studio software (R Development Core Team, 2012).

Predicting potentially suitable climatic areas for *E. aureum*

The predictive success of invasive weed risk assessments is still largely a function of invasiveness elsewhere, as well as, climatic suitability, therefore an understanding of these criteria provides significant value for management (Rouget *et al.*, 2004; Thuiller *et al.*, 2005).

There are no geo-referenced records of E. aureum in its native range. Moreover, although a number of sources report on the species invasiveness in several countries, very few global geo-referenced records exist (18 in total). Therefore, our presence-only data included the global records and the South African records obtained from field surveys conducted in this study (N = 171 with duplicate records removed).

To develop the niche model, current environmental data (1950–2000) were downloaded from the WorldClim database at a resolution of 30 arc-seconds (www.worldclim.org, accessed June 2012). The entire dataset of the 19 raster predictor variables was reduced through pairwise evaluation (Kendall rank correlation coefficient < 0.65) to reduce multi-collinearity among the predictors (Elith et al., 2010). We chose mean temperature of the coldest quarter as a primary predictor variable because *E. aureum* is limited by cold tolerance (http://aggie-

horticulture.tamu.edu/syllabi/308/Lists/Fourth%20Edition/Epipremnumaureum.pdf, accessed June 2012; Floridata plant encyclopedia, http://floridata.com/Plants/Araceae/Epipremnum%20aureum/1210, accessed June 2012). Subsequent variable selection was based on predictors with the lowest pairwise correlations. The resulting variables included four predictors; one rainfall (i.e. annual precipitation) and three temperature (i.e. minimum temperature of the coldest month, mean temperature of the wettest quarter and mean temperature of the coldest quarter) variables.

We used maximum entropy modelling to quantify habitats at potential risk of invasion and map the potential global geographic distribution of *E. aureum* (MaxEnt version 3.3.3 k; Phillips *et al.*, 2006). The selection of MaxEnt was based on the following reasons: (1) MaxEnt provides an appropriate strategy for working with presence-only data such as our *E. aureum* data set, (2) MaxEnt consistently outperformed other species distribution model implementations across taxa and geographic regions (Elith *et al.*, 2006), and (3) models are not strongly influenced by small sample sizes or irregularly sampled data and hence prediction is relatively robust (Pearson *et al.*, 2007, Elith *et al.*, 2011).

We generally opted for default MaxEnt settings: 10,000 random background points (i.e. pseudo-absences), from which the algorithm will select random points that are assumed as pseudo-absences, create response curves to evaluate *E. aureum* response to individual variables, logistic output to produce a continuous map, and jackknife procedure to measure variable importance. Appropriate selection of background points is essential for presence-only species distribution modelling because these points are generated from a random sample of non-occurrences in the region of interest. We ensured that the background points were geographically (i.e. selected from the spatial extent of presence records) and environmentally (i.e. selected from intersecting climate classes using the Köppen-Geiger climate classification) stratified. Furthermore, occurrence data is often spatially biased which generally results in environmental bias. However, spatial aggregation is reduced when using background data with the same bias as occurrence data but this does not correct for the lack of data due to low sampling effort. This selection method, together with 10,000 pseudo-absence points, is recommended when using a

machine learning technique in order to improve the models predictive accuracy (Barbet-Massin *et al.* 2012; Phillips and Dudik, 2008; VanDerWal *et al.* 2009).

In addition, we changed the following settings for model parameterization: (1) selected hinge features for smoother response curves, (2) regularization parameter = 1 to control over-fitting and clamping, (3) selected a random seed, (4) set random test percentage at 25 to evaluate model performance and reduce bias (75% of the data trained the model), (5) set replicates at 10 to ensure variability, (6) replicated run-type was set as subsample, and (7) set maximum iterations to 5,000 allowing the model adequate time for convergence.

The percent contribution of each variable and jackknife procedures were used to investigate the relative importance of the bioclimatic predictors while the 'area under the curve' (AUC) of the receiver operating characteristic was used to evaluate model performance. The AUC is a threshold-independent measure of model performance that ranges from 0 to 1. From the 10 replicates that were run we used the average AUC values for training and test datasets. Values > 0.9 indicate high accuracy, 0.7–0.9 indicates moderate accuracy, 0.5–0.7 indicates poor performance and values below 0.5 indicate that predictions are worse than random (Townsend Peterson *et al.*, 2011). Finally, using ArcMap version 10.2.2, the ASCII file containing the average model results were converted to binary maps in raster format. The colour distribution from light to dark represents increasing habitat suitability for *E. aureum*.

Weed risk assessment

We used the Australian Weed Risk Assessment protocol (A-WRA) to categorize the risk of invasiveness of *E. aureum* in South Africa based on its biology and ecology, climatic requirements, history, and biogeography (Pheloung *et al.*, 1999a). In the absence of any direct measure of impact, we used the A-WRA as a tool to predict potential weed impacts.

The A-WRA is a question-based scoring system, subdivided into sections on biology and ecology, climatic requirements, history, and biogeography. The assessment involves answering up to 49 questions and each question is awarded between -3 and 5 points. The final WRA score is the sum of points for all answered questions.

The answers generate a numerical score relating to the plants' invasive potential and the score is then used to determine one of three outcomes: the species is accepted for introduction (score < 1); rejected (score > 6); or rejected pending further evaluation of invasive potential (score 1–6). A minimum of 10 answers are needed for a species to be evaluated. The A-WRA system can therefore be used to identify useful non-problematic plants, as well as to predict potentially invasive plants of the agricultural and/or environmental sectors. In our study, questions related to geography and climate, were modified to reflect the conditions of South Africa (i.e. question 2.01).

Identifying best methods for control

We selected one research site in Umtentwini, KZN (-30.7224 °S, 30.464 °E), because this site contained many E. aureum plants growing on many trees over a large area (4,262 m²). This accounted for independent treatments and replication. Additionally, this selection intentionally factored in soil properties and climatic conditions. The uniformity of these variables across the trial site is essential because changes in these factors can influence the effects of the herbicide. The objectives of this trial were to evaluate the performance of three methods for *E. aureum* control: 1) cut treatment: cutting plants at 1.5 m above ground level with no herbicide application (i.e. the stem was severed completely); 2) glyphosate treatment: cutting plants at 1.5 m and tying plastic packets filled with 100 ml of herbicide mixture (2% Clearout 360 diluted with actipron, water and a marker dye) to the fresh cut stem on the aerial part of the plant, as well as, spraying the mixture on the fresh cut stem on the rooted plant; and 3) triclopyr and picloram treatment: cutting plants at 1.5 m and applying the herbicide gel (Kaput 100 Gel) with a brush to the fresh cut aerial and rooted parts of the plant. The control group, located at the same study site, received no treatment.

Measurements were done on the width of the stems using callipers 100 mm above the cut-stem and 100 mm below cut-stem. For the control we measured the width of the stem once at 1.5 m, i.e. height at which the plants were cut for the treatments. Stem width was used as a measure for effective control because previous trials that attempted to control this species revealed that the deterioration of the stems is more symptomatic of plant death compared to wilting leaves (Skene, pers. comm.). In

addition, measurements were taken above and below the cut-stem because the plants above the cut may still survive as they will be reliant on stored root reserves, while plants below the cut will primarily rely on nutrients obtained from the soil (Skene, pers. comm.). The reason for cutting the plants at 1.5 m was to ensure that the dead rooted part of the plant could be pulled out of the ground.

Glyphosate is the active ingredient in the 2% Clearout 360 herbicide, and triclopyr and picloram are the active ingredients in the Kaput 100 Gel. These two systemic herbicides were selected because literature searches suggest that these herbicides are commonly used to treat other Araceae species (e.g. *Colocasia esculenta, Philodendron* spp., *Pistia stratiotes* and *Syngonium podophyllum*). The experimental and control groups were set up on 27 August 2015 using 20 climbing *E. aureum* plants per treatment. The treatments and control group were not pseudo-replicated (n = 80). Thereafter, evaluations of the groups were done once a week for 6 weeks (3, 10, 17, 24 September; 1, 8 October). Statistical analyses were performed using the Kruskal-Wallis rank sum test.

RESULTS

Current distribution

Epipremnum aureum is present in several coastal towns in the KZN province, South Africa (Table 3.1; Appendix 8). From our information flyers, spotter networks and enquiries with staff from other herbaria across South Africa, we did not discover additional localities in other parts of the country. However, it must be noted that during this study our field survey effort was focussed in KZN because we only found records of naturalized *E. aureum* populations in this province (i.e. there may be populations in other provinces). Through road surveys, we found a total of 399 populations in the KZN province, of which 321 occur in gardens and 78 populations were growing in unmanaged sites which increases the species' potential to spread. Furthermore, these populations were distributed across 35 towns in KZN and populations in 9 of these towns were reported by spotters. This implies that although passive surveillance (e.g. through spotters) is valuable and less costly, active surveillance provides more accurate and timely information which is fundamental for post-border risk assessments.

Table 3.1. The distribution of *Epipremnum aureum* across KwaZulu-Natal, South Africa, as determined by spotters and road-side surveys. For wild populations, we collected information about the plants and the site. However, for populations that were cultivated in gardens, we only recorded their locality. Some towns are grouped together because they are located in close proximity.

Town	No. of garden populations	No. of unmanaged populations
Anerley	7	
Bazley Beach	1	_
Durban North	12	_
Eshowe	9	_
Glenwood	8	_
Hibberdene	1	_
Margate + Uvongo	15	9
Marina Beach	_	1
Melville	15	_
Mtwalume	3	_
Munster	7	_
Oslo Beach	8	_
Park Rynie	7	1
Pendale + Pennington +		
Kelso	37	5
Port Edward	_	8
Port Shepstone	_	5
Ramsgate	1	23
Scottburgh	15	1
Seapark	24	_
Sezela	2	_
Shelly Beach	7	3
Southbroom	4	4
St. Michael's on Sea	7	_
Sunwich Port	8	3
Trafalgar	3	_
Umgeni Park	7	_
Umkomaas	5	_
Umtentwini	16	9
Virginia	1	1
Westville	86	3
Winkelspruit + Warner Beach	5	2

Status of wild populations

The majority of the surveyed wild populations were observed as naturalized given that the plants were self-sustaining (i.e. reproducing), healthy and in large numbers.

This is a problem of great concern because these populations are wild (i.e. they are currently not managed) and therefore require active management. We found a total of 76 invasion foci requiring management across KZN with their extent of occurrence ranging from 1 m² to 3,383 m² within a population to a total occupancy area of 25,660.20 m² across all populations. Invasion foci were defined by populations that are clearly distinct containing the parent and/or initial site of introduction. For example, a garden population and the adjacent land in which the plants escaped reflected one invasion foci.

Abundance (i.e. the number of stems) varied widely across the populations, with sites comprising 3–34,011 plants. The total abundance across all sites was estimated at 186,667 individual stems (this can be interpreted as the estimated number of plants). Given the extent and abundance of the species, we classify *E. aureum* as category E under the Blackburn scheme (Blackburn *et al.*, 2011b). This category classifies *E. aureum* as a fully invasive species since there are several invasion foci comprising self-sustaining individuals that are distributed across a wide area.

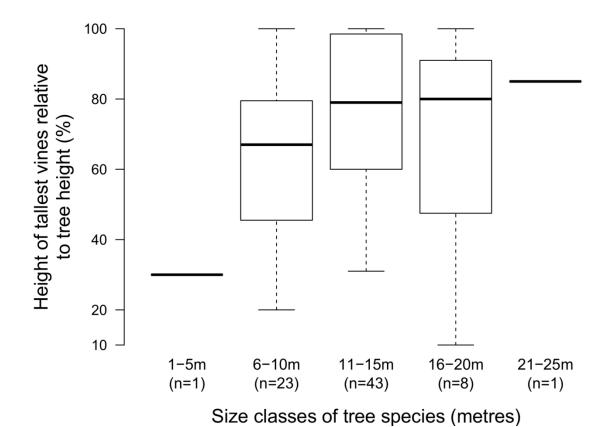


Fig. 3.1. Boxplot showing the relationship between the heights of the tallest vines relative to the heights of the supporting trees across 76 of the 77 populations surveyed. One population was supported by a boundary wall and therefore not included. Tree height was categorized according to size classes with population sample sizes shown in parentheses: 1-5 m (n = 1), 6-10 m (n = 23), 11-15 m (n = 43), 16-20 m (n = 8), 21-25 m (n = 1). The box is the interquartile range, and the bold centre line is the median.

The height of *E. aureum* also varied across populations (Fig. 3.1). Most of the populations comprised plants that were growing on trees ranging from 6 to 10 m (30%), 11–15 m (57%), and 16–20 m (11%). Of concern is that many populations consist of vines that have already reached the canopy. Furthermore, we observed that *E. aureum* does not have any preference for the support structures they climb. The species climbs up anything in its path, native or alien plant species, telephone poles, light poles and walls (Fig. 3.2A–C). When no structures were available to climb, *E. aureum* rapidly grows on the ground with a dense cover of its variegated foliage.



Fig. 3.2. Examples depicting the behaviour of *E. aureum* populations outside cultivation. **A)** Plants escaping from a garden and spreading in the neighbouring transformed vegetation, **B)** plants climbing up electricity poles, **C)** plants covering a telephone box, **D)** dense stands of *E. aureum* as a result of dumping garden refuse, and **E)** an additional example of illegal dump sites as a major source of *E. aureum* populations.

In South Africa, naturalized populations of *E. aureum* were primarily found in landuse types associated with some level of disturbance (Fig. 3.3). A chi-square goodness of fit test revealed a significant difference across the seven disturbance types and the number of *E. aureum* populations ($\chi^2 = 98.63$, df = 6, P < 0.001). This also demonstrates that the species can succeed across a range of land-use types. Furthermore, when comparing all land use types, significantly more populations were present in wastelands and this is a result of the disposal of garden cuttings (Fig. 3.2D–E). Thus, wastelands are a major source of *E. aureum* populations and characterize habitats in which the species flourishes. Garden escapees and roadsides also contributed significantly to the naturalization of *E. aureum*

populations. Of concern is the one population growing in natural vegetation and this population already covers a large area. Since we only found one population in natural vegetation this may represent an anomaly, however if left unmanaged, this could portray the future (i.e. many naturalized populations overgrowing natural vegetation).

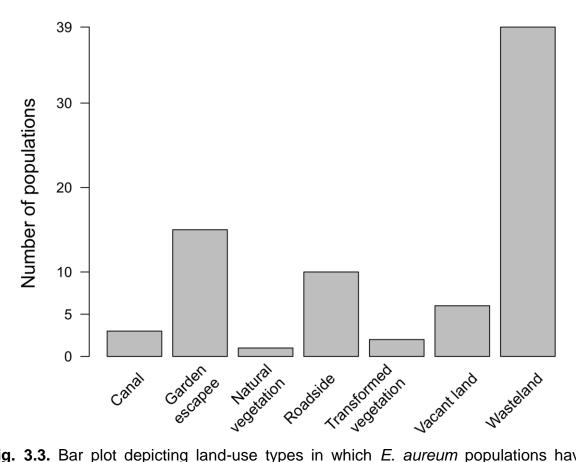


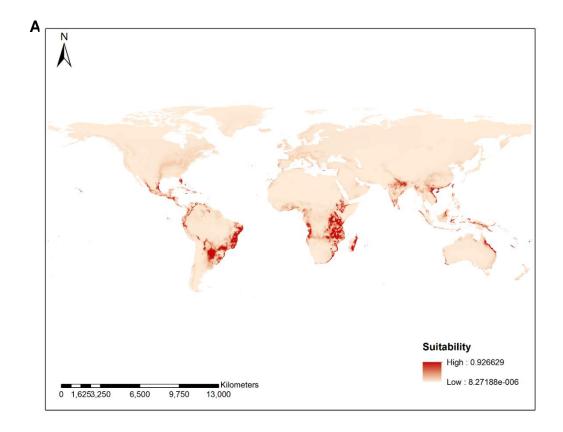
Fig. 3.3. Bar plot depicting land-use types in which *E. aureum* populations have naturalized.

Potentially suitable climatic areas for *E. aureum*

The average testing AUC value across the 10 iterations of the MaxEnt model was 0.967 ± 0.0156 (\pm standard deviation) which indicates good model performance for predicting suitable climatic conditions for *E. aureum*. The major contributors to invasion risk in descending order included minimum temperature of the coldest month (39.0730%) annual precipitation (33.546%), mean temperature of the coldest quarter (23.606%), and mean temperature of the wettest quarter (3.775%).

Based on model projections, many coastal tropical and sub-tropical regions provide suitable habitats for the species (Fig. 3.4A), including large parts of Mexico, Brazil, Angola, South Africa through to Mozambique and up to Kenya extending inland, India, Vietnam, and Australia (Queensland). The most climatically suitable islands for *E. aureum* include the Bahamas, Hawaii, Madagascar, Réunion Island, New Caledonia and several islands in Southeast Asia.

In South Africa, potential climatic suitability for *E. aureum* was largely restricted to the Indian Ocean coast (Fig. 3.4B). Projected habitat suitability occurs from Cape St. Francis in the Eastern Cape to KZN and its two neighbouring countries (Swaziland and Mozambique). Suitability also extends inland into the Mpumalanga and Limpopo provinces. On the southern Cape and west coast, Knysna and the Cape Peninsula were projected to be climatically suitable. Although we did not expect this region to be suitable, we must include it as a high risk region because both areas are home to National Parks. In addition, the entire predicted range includes many coastal towns within South Africa that are not yet known to be colonized. As such, there is a potential for further spread within South Africa, as well as, across the borders into Swaziland and Mozambique. This further highlights high priority locations for early detection and invasion control.



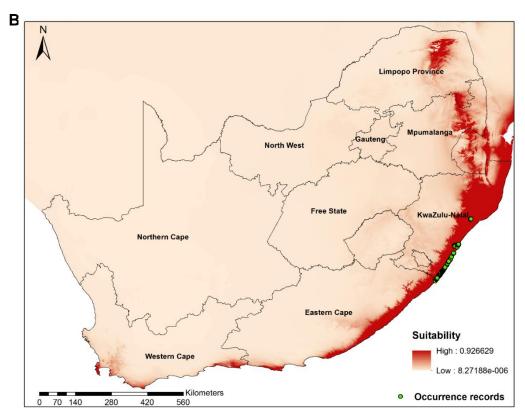


Fig. 3.4. Average suitability map showing the potential distribution of *E. aureum*, **A)** globally and **B)** zoomed into South Africa, based on existing occurrence data and climate variables modelled using MaxEnt. The maps represent an average of 10

replicates created using the subsampling method. The scale is a logistic probability with values between 0 (low probability; light shading) and 1 (high probability; darker shading).

Weed risk assessment

We were able to complete the A-WRA for *E. aureum*, answering 39 of the 49 questions (Table 3.2). Main gaps of knowledge correspond to reproductive characteristics (5 unanswered questions). This scheme predicted *E. aureum* to have a high probability of invasion in South Africa with a resulting score of 9. Consequently, pre-border screening should deny species entry into South Africa. Moreover, the environmental sector has a higher risk of impacts by *E. aureum* than the agricultural sector. The domestication of the species, climatic suitability, invasion history, weedy growth habit, high species densities, and vegetative propagation, preadapts *E. aureum* to becoming invasive. In addition, these characteristics suggest that *E. aureum* is likely to have profound negative impacts similar to other invasive vines (Blaustein, 2001; Pavlovic and Leicht-Young, 2011; *et al.*Yurkonis and Meiners, 2004; Zhang *et al.*, 2004).

Table 3.2. Australian weed risk assessment for *Epipremnum aureum*.

Question	Answer	Score	Possible scores
1.01 Is the species highly domesticated?	y ^a	-3	0 or -3
1.02 Has the species become naturalized where grown?	y ^b	1	-1 or 1
1.03 Does the species have weedy races?	n°	-1	-1 or 1
2.01 Species suited to South African climates	1 ^d	1	0, 1 or 2
2.02 Quality of climate match data (0-low; 1-intermediate; 2-high)	2 ^d	2	0, 1 or 2
2.03 Broad climate suitability (environmental versatility)	n ^d	0	0, 1 or 2
2.04 Native or naturalized in regions with tropical or subtropical climates	n ^e	0	0 or 1
2.05 Does the species have a history of repeated introductions outside its natural range?	y ^f	1	Refer to "lookup" table (Appendix 2) from Pheloung <i>et al.</i> , 1999
3.01 Naturalized beyond native range	У ^d	1	Refer to "lookup" table (Appendix 2) from

Pheloung et al., 1999

3.02 Garden/amenity/disturbance weed	у ^g	1	Refer to "lookup" table (Appendix 2) from Pheloung <i>et al.</i> , 1999
3.03 Weed of agriculture/horticulture/forestry	y ^h	2	Refer to "lookup" table (Appendix 2) from Pheloung <i>et al.</i> , 1999
3.04 Environmental weed	у ^с	2	Refer to "lookup" table (Appendix 2) from Pheloung <i>et al.</i> , 1999
3.05 Congeneric weed	λ _a	1	Refer to "lookup" table (Appendix 2) from Pheloung <i>et al.</i> , 1999
4.01 Produces spines, thorns or burrs	n ^d	0	0 or 1
4.02 Allelopathic	n°	0	0 or 1
4.03 Parasitic	n°	0	0 or 1
4.04 Unpalatable to grazing animals	f		-1 or 1
4.05 Toxic to animals	y ^a	1	0 or 1
4.06 Host for recognised pests and pathogens	y ⁱ	1	0 or 1
4.07 Causes allergies or is otherwise toxic to humans	y ^j	1	0 or 1
4.08 Creates a fire hazard in natural ecosystems	n°	0	0 or 1
4.09 Is a shade tolerant plant at some stage of its life cycle	y ^k	1	0 or 1
4.10 Grows on infertile soils	y ^k	1	0 or 1
4.11 Climbing or smothering growth habit	y ^l	1	0 or 1
4.12 Forms dense thickets	y ^m	1	0 or 1
5.01 Aquatic	n ^h	0	0 or 5
5.02 Grass	n ^h	0	0 or 1
5.03 Nitrogen fixing woody plant	n ^h	0	0 or 1
5.04 Geophyte	n ^h	0	0 or 1
6.01 Evidence of substantial reproductive failure in native habitat	n°	0	0 or 1
6.02 Produces viable seed	f		-1 or 1
6.03 Hybridises naturally	f		-1 or 1
6.04 Self-fertilisation	f		-1 or 1

6.05 Requires specialist pollinators	f		0 or -1
6.06 Reproduction by vegetative propagation	y ⁿ	1	-1 or 1
6.07 Minimum generative time (years)	f		-1, 0, or 1
7.01 Propagules likely to be dispersed unintentionally	y ^d	1	-1 or 1
7.02 Propagules dispersed intentionally by people	y ^d	1	-1 or 1
7.03 Propagules likely to disperse as a produce contaminant	n°	-1	-1 or 1
7.04 Propagules adapted to wind dispersal	n ⁿ	-1	-1 or 1
7.05 Propagules buoyant	n°	-1	-1 or 1
7.06 Propagules bird dispersed	f		-1 or 1
7.07 Propagules dispersed by other animals (externally)	f		-1 or 1
7.08 Propagules dispersed by other animals (internally)	f		-1 or 1
8.01 Prolific seed production	n ^f	-1	-1 or 1
8.02 Evidence that a persistent propagule bank is formed (>1 yr)	n ^f	-1	-1 or 1
8.03 Well controlled by herbicides	y ^d	-1	-1 or 1
8.04 Tolerates or benefits from mutilation, cultivation or fire	y ^a	1	-1 or 1
8.05 Effective natural enemies present in South Africa	f		-1 or 1

^aMeshram, A. and Srivastava, N. 2014. Molecular and physiological role of Epipremnum aureum. International Journal of Green Pharmacy. 8: 73-76; bPIER. Pacific Island **Ecosystems** at Risk, http://www.hear.org/pier/species/epipremnum_pinnatum_cv_aureum.htm, accessed ^dThis cNo February 2016; evidence; paper; ehttp://mobile.floridata.com/Plants/Araceae/Epipremnum%20aureum/1210; fUnknown; ⁹Global compendium of weeds, http://www.hear.org/gcw/species, accessed February 2016; ^hBoyce, P., 2004. A review of Epipremnum (araceae) in cultivation. Aroideana 27, 199-205; Wick, R.L. and Dicklow, M.B. 2002. Epipremnum, a new host for *Phytophthora capsici*. Plant Disease 86 (9): 1050; ^jSpoerke, D.G. and Smolinske, S.C. 1990. Toxicity of houseplants.CRC Press, Florida; https://edis.ifas.ufl.edu/fp194, accessed March 2016; https://plantdatabase.kpu.ca/plant/plantDetail/54, accessed March 2016; "Nyanatusita, B., Dissanayake, R., 2013. Udawattakele: a sanctuary destroyed from within. Journal of the Wildlife and Nature Protection Society of Sri Lanka 26; "http://aggie-horticulture.tamu.edu/syllabi/308/Lists/Fourth%20Edition/Epipremnumaureum.pdf, accessed March 2016.

Best management practices

The impacts of the control group (i.e. no treatment) on stem widths did not differ significantly to plants above the cut stem treatment (Fig. 3.5A). In both cases 35% of the plants had no change in their stem width while some of the plants continued to flourish (indicated by the 20% increase in stem width of the control group and 7% in cut treatment). However, although not significant, there was also some decrease in stem width (control group: 19% and the cut treatment: 11%). Interestingly, seeing as plants cut off from their roots were still able to grow and persist without any indication of new roots being produced, we can deduce that *E. aureum* does not only absorb nutrients from the soil but also from the air and/or stored reserves. The herbicide treatments were significantly superior since stem widths deteriorated by 47% after Clearout 360 (i.e. glyphosate) and 41% after the Kaput treatment (i.e. triclopyr and picloram). Furthermore, the herbicide treatments resulted in mortality of three plants during the trial (i.e. whole plants dropped to the ground with no chance of recovery).

The impacts of the treatments below the fresh cut stem showed a similar trend to the aerial part of the plant (Fig. 3.5B). However, Clearout 360 performed significantly better (56% decrease in stem width) on the roots of *E. aureum* than Kaput (31% decrease in stem width). Furthermore, triclopyr and picloram did not appear to be translocated into the roots because three weeks into the trial we observed growth of new shoots in plants treated with Kaput, but this was not observed in the glyphosate treatment.

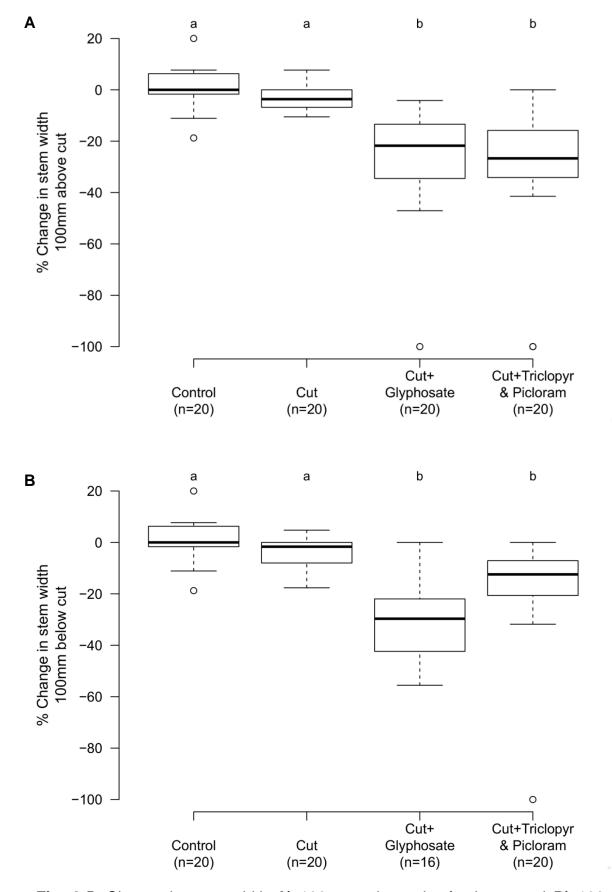


Fig. 3.5. Change in stem width A) 100 mm above the fresh cut and B) 100 mm below the fresh cut *E. aureum* plants. Stem width was measured across four

treatment groups during a six week trial. Boxplots display the median with a solid line, 25th and 75th percentiles in the lower and upper boxes, respectively, and the data range is indicated by the whiskers. Open circles indicate outliers (values > 1.5 times interquartile distance below 25th percentile). We used the non-parametric Kruskal-Wallis test to compare means ranks between treatment groups. Letters denote significance of the mean ranks (P < 0.05).

DISCUSSION

The most cost-effective strategy against IAS following post-border introductions is early detection and eradication when populations are small and localized. Following quantification of the spatial extent, abundance, competitive ability, potential for further spread into suitable habitats, and the high invasion risk of *E. aureum* it can be concluded that this evergreen climber poses a significant but currently manageable threat to biodiversity in the KZN province of South Africa. Our results suggest that it is possible to control naturalized populations of *E. aureum* because, although the species is spread across the coastal areas of the KZN province, the total condensed extent of occurrence (~3 hectares) and vine abundance (i.e. number of plant stems, ~187,000) are still low. For these reasons it is necessary to take action and manage the species in order to curb potential impacts on the ecosystem. We believe that containment is a viable strategy for managing *E. aureum* in South Africa.

During our field surveys (i.e. over a 2 year period) no flowers were seen, therefore reproduction and spread via seeds might be negligible. As a result, the more stems in a population, the higher the propagule pressure for establishment and spread. Propagule pressure is regarded as a major driver of invasive success of alien species (Colautti *et al.*, 2006; Lockwood *et al.*, 2005a; Simberloff, 2009), and our results on plant abundance also supports this hypothesis. This is based on the idea that increasing the number of individuals should increase the success of establishment. *Epipremnum aureum* grows rapidly and sustains its populations via vegetative reproduction and stem fragments. Therefore maintaining its populations with high plant numbers was expected.

Despite the thick stems (e.g. largest stem measured was 48 mm in diameter) and large leaves, this vine can still grow high into the canopies (e.g. tallest vine was 18

m). The maximum height that a species can reach is an indicator of its light capture strategy and competitive ability. Moreover, tall plants are associated with invasion success (Bucharova and Van Kleunen, 2009; Pyšek and Richardson, 2007). Climbing capacity enables climbers to ascend up to a height to compete for light and space and this facilitates their success as an invader (Paul and Yavitt, 2011). Epipremnum aureum thrives under high light conditions and therefore creeps, trails and clings around anything in its path. In doing so, the species shades out its competitors and monopolizes the light environment for photosynthesis (cf. Hejda et al., 2009).

Epipremnum aureum is commonly grown as an ornamental plant in gardens, worldwide and in South Africa (for example, we found 321 cultivated populations in KZN, South Africa). Humans also contribute significantly to the spread of *E. aureum*, by dumping their garden waste in unmanaged spaces, illegal dumping sites, parks and natural areas, as well as allowing their garden plants to skip the fence. Seeing as invasive species often establish more frequently in disturbed rather than pristine habitats (Didham et al., 2005), it is clear that anthropogenic disturbances exacerbate the threat of *E. aureum*. In the field we also observed a striking correlation between dump sites and numerous dense E. aureum populations. Being a vegetativelypropagated plant, E. aureum is able to rapidly spread and become established in dense monocultures on the ground, up the trees and telephone poles, and across walls and fences. Repeated patterns of human-mediated disturbances give alien plants a chance to establish and spread. Populations growing along roadsides were also common. Roads are particularly good corridors as they alter conditions, stress indigenous species, and allow easier access of humans as vectors of plant dispersal. They also have higher light conditions and bare soil, which favour alien plant establishment (Gelbard and Belnap, 2003; Mortensen et al., 2009; Pauchard and Alaback, 2004). Therefore, management measures to reduce human disturbance need to be implemented. For example, increase awareness and knowledge of invasive plants within the community and encourage the disposal of garden waste through the local municipalities.

From a management point of view, it is imperative to identify areas that are not yet invaded but where early warning, detection, and control programmes can be

implemented. *Epipremnum aureum* has a wide potential distribution across the coastal regions of South Africa including numerous unoccupied regions by the species. In view of the fact that *E. aureum* has only been reported in KZN thus far, there is a high possibility of control and eradication. If not contained, there is a high invasion risk in the Eastern Cape, southern Cape and parts of the Mpumalanga and Limpopo regions which provides suitable habitats. Among many management activities, monitoring and mapping the occurrence of invasive species is important for control action. The model presented in this study can be used for informing management plans and guiding monitoring efforts in preventing further spread of *E. aureum* in South Africa.

Since *E. aureum* is already a problem in the KZN province, and given its high potential habitat suitability in regions that it is currently not known to occupy, it has a high potential to expand its range. Therefore control efforts of populations present in KZN must be prioritised. Mechanical control, by itself, of *E. aureum* is not feasible given the vines' abundance in a single population, thick stems and great heights reached on trees. In addition, although cutting weakened the plant, the vegetative nature and rapid growth of *E. aureum* allows the species to persist and quickly reoccupy the space. Hence, the problem is alleviated only temporarily. Fortunately, chemical control on cut stems proved successful in destroying plants.

Between the two chemicals tested, field observations and experiments indicate that glyphosate is more successful in affecting *E. aureum*. This treatment works from the canopy down to the ground seeing as both leaf chlorosis and the loss of stem turgor were observed in that order. The triclopyr and picloram treatment primarily affected the stems causing them to lose turgor and change colour. Therefore, chemical control is preferable to mechanical control and indicates promising potential for the control of *E. aureum*. Cost–benefit analyses of biological control programs usually show an overpowering economic justification for the use of biological control agents (De Clercq *et al.*, 2011; Olckers, 2004). However, for *E. aureum* biological control is probably not ideal since a) there is no tested agent available, b) effective chemical control methods are available, and c) most populations detected to date are accessible with low levels of spread.

CONCLUSION

Epipremnum aureum is a highly invasive species in Hawaii and many parts of Asia. In South Africa, given the large number of naturalized populations and the lack of records from the other provinces, the KZN province appears to be an invasion hotspot. The species has a high potential to become invasive under favourable conditions and therefore has a high risk of becoming invasive along the east and south coasts of South Africa. Although many of the populations comprise a large number of plants, herbicidal control appears to be effective and feasible.

To minimise the threat posed, we recommend that new plantings be prohibited and all populations outside cultivation (i.e. 78 naturalized populations) be controlled and ideally extirpated. However, given the fact that the species appears to be relatively limited in its dispersal ability, has no seed banks, and no chance of maintaining populations vegetatively if treated with herbicides, it might pose a manageable threat if allowed to remain on a property provided that the species is contained within the property. However, we recommend that the species should not be further propagated or sold (i.e. category 3 under South Africa's National Environmental Management: Biodiversity Act (10/2004) A&IS regulations).

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Chapter 4: Invasion dynamics of a horticultural vine, Monstera deliciosa: what drives the successes and failures of invasions?

ABSTRACT

To understand the success of invading species, knowledge on factors driving invasiveness is needed. Here, we look at Monstera deliciosa (Araceae) and its relatives in a South African and global context. First, we explore the subfamily Monsteroideae to assess whether attributes related to introduction history influences the invasion status of species. Next, we classify the invasion risk of *M. deliciosa* in South Africa using a weed risk assessment and species distribution models. Lastly, using a local scale study, we identify potential conditions under which M. deliciosa can become naturalized. Overall, Monsteroideae species that have been introduced for a longer period of time, and those that have been introduced to several new regions have a significantly higher likelihood of becoming naturalized and invasive. Monstera deliciosa was ranked as having a high impact risk in South Africa and also modelled to have a wide potential global distribution. On a local scale, naturalization success was driven by warm temperatures and sufficient water. However, despite the risk, we conclude that *M. deliciosa* will not become a rampant invader based on the fact that the species already had a sufficiently long minimum residence time and high propagule pressure in South Africa and in several currently occupied regions globally, yet there are no records of invasiveness or impacts. Even if the species becomes invasive under ideal conditions, it can be controlled manually. As such, M. deliciosa can be recommended to the horticultural trade, landscaping sector and gardeners.

Keywords

Biological invasion, green list, horticulture, low risk, Monsteroideae, time lag

INTRODUCTION

The growing magnitude of worldwide interconnectedness through globalization has intensified the flow of trade, transport, and travel. Consequently, this increased mobility of people and their goods removes natural barriers between and across ecosystems. This facilitates the introduction of new species to ecosystems, as well as, intensifies the spread of invasive alien species (IAS) through intentional or accidental introductions (Hulme, 2009; McNeely, 2001b; Meyerson and Mooney, 2007a; Perrings *et al.*, 2010). In addition, IAS are now widely cited as the second greatest global threat to biodiversity. Therefore, understanding the factors that facilitate or limit invasions is an important step towards predicting and managing invasive species, and ultimately conserving biodiversity.

For a species to become invasive, it must be: introduced deliberately or unintentionally through human activities to an area where it is not native; resistant enough to survive in the new area; establish self-sustaining populations; disperse; and spread (Blackburn *et al.*, 2011a; Richardson *et al.*, 2000a). However, not all species become invasive when introduced to novel systems. Only a proportion of species introduced to novel systems become naturalized, and of those, a small subset spread and become invasive (Richardson and Pyšek, 2006a; Williamson and Fitter, 1996). Although only a small percentage of species become invasive, their negative impacts (i.e. ecological, economic, and/or social impacts) can be extensive, and over time causes substantial damage which requires high control costs (Mack *et al.*, 2000; Pimentel, 2011; Pimentel *et al.*, 2005; Vilá *et al.*, 2010). For that reason urgent action is required.

Furthermore, although many introduced species fail to become invasive (e.g. Bufford and Daehler, 2014; Copp *et al.*, 2007; Miller *et al.*, 2007; Reichard and Hamilton, 1997; Rejmánek and Richardson, 1996; Young, 2015), there are many studies looking at species that were able to persist and comparatively fewer studies that have addressed the issue of failed invasions. Instances of failed invasions can highlight factors that limit the expansion of an invasive species (Lonsdale, 1999; Zenni and Nuñez, 2013). This type of knowledge will further our understanding of biological invasions and also identify species with a low invasion risk that can be recommended for introduction.

Identifying and prioritizing pathways of introduction is major component of managing biological invasions (McGeoch *et al.*, 2016). Species introduced for horticulture, ornamental use in particular, are dominating invasion pathways worldwide (Dehnen-Schmutz and Touza, 2008; Ööpik *et al.*, 2013; Reichard and White, 2001; Richardson and Rejmánek, 2011). The invasion success of plants introduced via this pathway is a result of importers selecting plants with specific characteristics that inadvertently also match characteristics associated with invasiveness (Bucharova and Van Kleunen, 2009; Dehnen-Schmutz *et al.*, 2007b). Furthermore, ornamentals that are marketed at a good price also influence invasion success. For example, Dehnen-Schmutz *et al.* (2007b) showed that species marketed at low prices and species that were regularly available in the 19th and 20th centuries were strong predictors of the current range of alien plants in Britain. Lastly, the high number of introduction events and the nurturing of plants by gardeners also enable species to overcome the various barriers to invasion (Kowarik, 2003).

Preventing biological invasions is the most cost-effective and efficient approach, however, if the IAS has already been introduced, then early detection and management become important to prevent establishment. As a general rule, it is understood that under certain conditions any widely planted species will have a greater likelihood of becoming naturalized and invasive, even more so, when species have been introduced long ago. Introduction history has been identified as an important driver of invasiveness because it usually influences a high propagule pressure (Cassey *et al.*, 2004; Colautti *et al.*, 2006; Lockwood *et al.*, 2005b; Simberloff, 2009; Von Holle and Simberloff, 2005) and sufficiently long residence time (Dehnen-Schmutz *et al.*, 2007a; Phillips *et al.*, 2010; Pyšek and Jarošík, 2005; Wilson *et al.*, 2007) which are significant drivers of invasion. In addition, introduction history as a correlate of invasiveness (i.e. residence time in particular) also facilitates invasiveness in alien vines (Harris *et al.*, 2007).

Another challenge in invasion biology, which led to the premise of this study, is to understand the patterns and conditional nature of invasions. Across the spectrum of species introduced to novel systems there are; species that clearly become a problem; species that will not become a problem; and a unique category of species that fall somewhere in between – widely planted but not yet a major invader. This

unique pattern may be explained by a poorly studied species, *Monstera deliciosa* Liebm., which is widely planted yet it has seemingly minimal records of being invasive.

Although *M. deliciosa* is our focal case study species, we first selected the subfamily Monsteroideae to assess whether attributes related to introduction history influences the invasion status of species within this clade. Residence time is one of the most important correlates of vine invasiveness (Harris *et al.*, 2007). The reasoning behind this is that species that are present for a longer time have a higher probability of spreading more propagules and forming new populations (Wilson *et al.*, 2007). Propagule pressure has also been identified as one of the most consistent predictors of invasiveness (Colautti *et al.*, 2006; Simberloff, 2009). Species that are introduced across a wider area in a new environment have a better chance of landing in localities that are suitable for establishment (Lockwood *et al.*, 2005b). Therefore, to identify whether species that are introduced into more regions have a greater chance of becoming invasive, we used the number of introduced regions as a proxy for propagule pressure. Additionally, a species' continent of origin is another important predictor associated with introduction history (Zenni, 2014). This attribute takes into account the chance of dispersal by humans from particular source areas.

We then explore these trait findings to unravel possible explanations of the global status of *M. deliciosa*. Next, at a regional scale, we aim to classify the invasion risk of *M. deliciosa* in South Africa using a weed risk assessment and species distribution models. Finally, on a local scale, we describe potential conditions under which *M. deliciosa* can become invasive.

MATERIALS AND METHODS

Study species

Monstera deliciosa, commonly known as the Swiss cheese plant, is an evergreen climber native to tropical moist forests in Mexico, Costa Rica, Guatemala, Honduras, Nicaragua, and Panama (eMonocot, http://emonocot.org/, accessed August 2013). This climber has been recorded to reach 20 m or more in height, often depending on the height of the support structure, and can also form a dense mat on the ground when unsupported. Monstera deliciosa's attractive large leaves and tolerance of

shade and humidity make it one of the most widely cultivated ornamental plants (Madison, 1977). *Monstera deliciosa* is now grown in most of the warm countries of the world (temperate and tropical regions) as a potted indoor plant or as a garden plant. However, it has only been recorded as naturalized in a few regions, and is surprisingly not a major invader (Table 4.1).

Table 4.1. List of introduced regions where *M. deliciosa* is grown and its associated invasion status. This list relies on published literature and online databases. Searches used for identifying the relevant literature and databases were undertaken online using the following combination of terms in Google: *M. deliciosa* was used in conjunction with "grown", "cultivated", "alien", "exotic", "introduced", "naturalized", "invasive", "impact".

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Region	Year of record	Status	Reference
Ascension	2005	Introduced	Varnham, K. 2005. Non-native species in UK Overseas Territories: a review. JNCC Report 372. Peterborough, United Kingdom
Australia	1991	Introduced	Global Compendium of Weeds (GCW). http://www.hear.org/gcw/species/monstera_deliciosa/, accessed September 2016
Azores	_	Naturalized	Delivering Alien Invasive Species In Europe (DAISIE). http://www.europe- aliens.org/speciesFactsheet.do?speciesId =717#, accessed September 2016
Bermuda	2005	Introduced	Varnham, K. 2005. Non-native species in UK Overseas Territories: a review. JNCC Report 372. Peterborough, United Kingdom

California	2002	Introduced	Jepsen, E.P.B., and Murdock, A.G. 2002. Inventory of native and non-native vegetation on John Muir national historic site, Eugene O'Neill national historic site, and Port Chicago national monument, http://www.sfnps.org/download_product/2 606/0, accessed September 2016
England	1752	Introduced	http://floridata.com/Plants/Araceae/Monste ra+deliciosa/794, accessed November 2016; http://edis.ifas.ufl.edu/hs311, accessed November 2016
Florida	1976	Introduced	Morton, J.F. 1976. Pestiferous spread of many ornamental and fruit species in south Florida. Proceedings of the Florida State Horticultural Society, 89: 348-353
Galapagos	2004	Naturalized	Global Compendium of Weeds (GCW). http://www.hear.org/gcw/species/monstera_deliciosa/, accessed September 2016
Hawaii	2010	Introduced	Benitez, D.M., R. Loh, T. Tunison, N.G. Zimmer, J. Makaike, R. Mattos and M. Casali. 2012. The distribution of invasive plant species of concern in the Kīlauea and Mauna Loa strip areas of Hawaiʻi Volcanoes National Park, 2000-2010. Technical Report No. 179. The Hawaiʻi-Pacific Islands Cooperative Ecosystem Studies Unit & Pacific Cooperative Studies Unit, University of Hawaiʻi, Honolulu, Hawaiʻi. 120 pp.
India	1878	Introduced	http://edis.ifas.ufl.edu/hs311, accessed November 2016

London	_	Introduced	http://www.clifton.co.uk/cg- monstera_deliciosa.htm, accessed November 2016
Madeira	2002	Naturalized	Delivering Alien Invasive Species In Europe (DAISIE). http://www.europe- aliens.org/speciesFactsheet.do?speciesId =717#, accessed September 2016
Mediterranean	2012	Introduced	Heywood, V.H. 2012. The role of New World biodiversity in the transformation of mediterranean landscapes and culture. Bocconea 24: 69-93
Motuora Island, Hauraki Gulf	2006	Introduced	Heiss-Dunlop, S., and Fillery, J. 2006. Vascular flora of Motuora Island, Hauraki Gulf. Auckland Botanical Society Journal, 61(2): 113-120
Pitcairn Islands	2009	Introduced	Varnham, K. 2009. Non-native species in UK Overseas Territories: an updated review. JNCC Report 372. Peterborough, United Kingdom
Puerto Rico	2000	Introduced	Global Compendium of Weeds (GCW). http://www.hear.org/gcw/species/monstera _deliciosa/, accessed September 2016
Saint Helena	2005	Introduced	Varnham, K. 2005. Non-native species in UK Overseas Territories: a review. JNCC Report 372. Peterborough, United Kingdom
Singapore	1877	Introduced	http://edis.ifas.ufl.edu/hs311, accessed November 2016
South Africa	1989	Naturalized	Henderson, L. 1989. Invasive alien woody plants of Natal and the north-eastern Orange Free State. Bothalia 19(2): 237-261

Spain – Introduced Delivering Alien Invasive Species In

Europe (DAISIE). http://www.europe-aliens.org/speciesFactsheet.do?speciesId

=717#, accessed September 2016

New Zealand – Naturalized Howell, C. 2008. Consolidated list of

environmental weeds in New Zealand. DOC Research & Development Series 292. Department of Conservation,

Wellington. 42 pp.; Global Compendium of

Weeds (GCW).

http://www.hear.org/gcw/species/monstera_deliciosa/, accessed September 2016

Global introduction history of Monsteroideae

To assess whether introduction history drives invasiveness in this subfamily we only looked at species that are introduced but not yet naturalized versus naturalized and invasive species. This included a total of 42 species belonging to 9 genera (Moodley et al., 2016a). Since we usually do not know exactly when a species was introduced, we used minimum residence time (MRT) based on the year in which the first herbarium specimen was collected. We estimated MRT using the formula 2016-x, with x being the earliest introduced record of the species and 2016 being the current year. Herbarium records were obtained from the Global Biodiversity Information Facility (http://www.gbif.org/, accessed September 2016) and Tropicos (http://www.tropicos.org, accessed September 2016). We identified the species' continent of origin by first obtaining their countries of origin using eMonocot and then assigning them to Africa, Asia, Australia/Oceania, Europe, North America, and South America. The number of introduced regions were extracted from a recently complied Araceae database (Moodley et al., 2016a).

Statistical Analysis

To explore whether minimum residence time, country of origin, and the number of introduced regions (i.e. explanatory variables) influences the invasion status (i.e.

response variable) of introduced Monsteroideae, a one-way analysis of variance (ANOVA) or the non-parametric alternative (Kruskal-Wallis rank sum test) were used. The ANOVA was used if the data satisfied the following assumptions: observations are independent; have a normal distribution; and scores in different groups have homogeneous variances (i.e. minimum residence time and country of origin). We used the Kruskal-Wallis rank sum test when the assumptions were violated (i.e. number of introduced regions). All statistical analyses were performed in R (version 3.3.1, R Development Core Team, 2016).

Weed risk assessment of *M. deliciosa* in South Africa

One of the most popular screening tools to date is the Australian Weed Risk Assessment (A-WRA), which consists of 49 questions that categorizes the risk of species becoming invasive (Pheloung et al., 1999b). The questions focus on biogeography, biology/ecology, and undesirable plant attributes, and the answers are scored from least likely (-3) to most likely (5). Several studies demonstrated the consistency and accuracy of the A-WRA in predicting invaders (Daehler et al., 2004; Gassó et al., 2010; Gordon et al., 2008b; Gordon et al., 2008c; Křivánek and Pyšek, 2006). Hence, we selected this screening protocol. In our study, questions related to geography and climate, were modified to reflect the conditions of South Africa. Suitability of species to Australian climate was changed to suitability to South African climate (question 2.01) and native or naturalized in regions with extended dry periods was changed to native or naturalized in regions with tropical or subtropical climates (question 2.04). Species are identified as potentially high risk and rejected for import if the score is > 6, potentially low risk and accepted for import if the score is < 1, and species with intermediate scores (1-6) are recommended for further evaluation.

Potential global distribution of *M. deliciosa* based on climate

We used the widely accepted maximum entropy model (MaxEnt version 3.3.3k) to predict the probability of *M. deliciosa* occurrence and map its potential geographic distribution (Phillips *et al.*, 2006). MaxEnt estimates the probability distribution of a species based on the maximum entropy principle by comparing environmental conditions and associated presence records to background points (Elith *et al.*, 2011;

Phillips and Dudík, 2008). This software was implemented in R (version 3.3.1, R Development Core Team, 2016).

Occurrence records, background data and environmental layers

We downloaded georeferenced occurrence records from both the native and introduced ranges because this distribution encompassed the most realistic estimation of *M. deliciosa's* climate niche. We also downloaded georeferenced records for two closely related introduced *Monstera* species (i.e. *Monstera obliqua* Miq. and *Monstera punctulata* (Schott) Schott ex Engl.) to use as background data (i.e. pseudo-absence). These two species were native to the same regions as *M. deliciosa*. This method is referred to as target group sampling and is recommended over randomly generated background points because it accounts for sampling bias (Phillips *et al.*, 2009). All records were downloaded from GBIF (http://www.gbif.org/) using the package rgbif (Chamberlain *et al.*, 2016). We used the "getData" function from the raster package (Hijmans, 2016) to download bioclimatic data from the WorldClim database at a spatial resolution of 10 arc-minutes. This database provides climatic data averaged over the years 1950–2000.

The 19 climatic variables were reduced through pair-wise correlation to avoid multicollinearity among these variables (Kendall rank correlation coefficient < 0.65), as suggested by Elith et al. (2010). We chose annual mean temperature as our primary predictor variable because M. deliciosa grows best between 20–30°C and is killed by frost (Lim, 2011). Subsequent variable selection was based on predictors with the lowest pair-wise correlations. The resulting variables included four predictors; two temperature (annual mean temperature, temperature annual range (i.e. maximum temperature of the warmest month - minimum temperature of the coldest month)) and two rainfall (annual precipitation, precipitation seasonality) variables. Lastly, the biogeo package (Robertson, 2016) was used for data cleaning and this involved removing fossil records, duplicates within 10' grid cells, points in the sea, points that do not have environmental data, as well as pseudo-absences with the same coordinates as presence points. In addition, M. deliciosa records present in areas that experience frost, such as in Europe and the United States of America, were also removed to ensure data reliability. This resulted in 210 occurrences and 390 pseudo-absences.

Model building and evaluation

We used the dismo package (Hijmans *et al.*, 2016) to build the model using the occurrence records mentioned above and 10,000 pseudo-absence records. Pseudo-absence data are used in lieu of observed absence data to define environmental conditions in which the species has not been recorded. This means that biased sampling can influence the accuracy of the model (Phillips *et al.*, 2009; VanDerWal *et al.*, 2009). Therefore, MaxEnt assumes that species occurrence data are unbiased (Phillips *et al.*, 2006). As such, we limited the spatial extent from which MaxEnt could select pseudo-absence points to locations within 2 degree (i.e. ~240 kilometres) of occurrence records (Merow *et al.*, 2013). Bias in occurrence records was further addressed by producing a bias grid derived from a Gaussian kernel density map of the occurrence locations (Elith *et al.*, 2010).

MaxEnt was run using only linear, quadratic and hinge features. This reduces the chances of model overfitting and results in smoother response curves, as well as, models that are focussed on the strongest trends in the data (Elith et al., 2010; Merow et al., 2013). Models were trained using a randomly selected 70% of the data (both occurrences and pseudo-absences) and model accuracy was then tested on the remaining 30%. This process was repeated 100 times in order to account for variability in model performance arising from the selection of training and testing data. From this, we received training and testing area under the receiver operating characteristic curve (ROC) values, also known as AUC. The AUC value is a widely used measure of model performance and is interpreted as the probability that a randomly selected pair of occupied and unoccupied sites are correctly predicted (Elith et al., 2006). Values range from 0 to 1, and values > 0.9 indicate high accuracy, values of 0.7-0.9 indicate good accuracy, and values < 0.7 indicate low accuracy (Swets, 1988). Lastly, since the regularization coefficient in Maxent (β) can impact on model predictions, we set the model to explore different values for β and select a model based on the best AUC value (Merow et al., 2013).

Model projection

We used the raw output from MaxEnt to predict the potential distribution of *M. deliciosa* based on climatic suitability. Using the raw output is preferable to the logistic transformation output since it is based on the probability of climatic suitability

(Merow *et al.*, 2013). The final MaxEnt output provides a calculation of the percent contribution each variable has on the model building process and also reveals a surface with a continuous climatic suitability gradient with values ranging from 0 (least suitable) to 1 (most suitable).

A qualitative assessment of an invasive population in the Limpopo Province, South Africa

Only two entries of *M. deliciosa* populations in South Africa are recorded in the Southern African Plant Invader Atlas (Henderson, 2007). This database lists one population in the KwaZulu-Natal Province, with no additional information provided, and the other in the Limpopo province. Hence, we could only follow up one the Limpopo record. This record was lodged in 2012 and describes the population as a garden escapee that is invading the moist high-rainfall southern slopes of the Soutpansberg Mountain. The garden, surrounding the main house and originally created by the first landowner, is situated within the lajuma research centre high up in the Soutpansberg mountain range. Lajuma has been declared a natural heritage site and comprises a variety of veld types, including montane forests and grasslands, and a remarkable diversity of plants and animals.

Field work in 2016 for this study involved mapping the extent of this garden population and evaluating reasons for its spread. The extent was determined by calculating the area of a convex hull drawn around the most outlying points in ArcGIS 10.4. Spread was measured by identifying the initial plantings by the landowners and then mapping the plants expansion. Given the ruggered terrain, surveys were conducted by foot along the roads, trails and fence lines in the reserve in order to track every plant in the garden, as well as those which may have escaped into the surrounding reserve. Because this was a qualitative assessment of the first detected naturalized population, we were more interested in the land use type, whether the area was managed, modes of spread, and the total area of spread.

RESULTS

Global introduction history of Monsteroideae

The Monsteroideae subfamily comprises 362 species belonging to 9 genera. Out of these, 39 species (11%) have been introduced but are not yet naturalized and 3

species (0.82%) have become naturalized or invasive (Fig. 4.1A, Appendix 9). Species that have been introduced over a longer period of time were more likely to naturalize and invade in their new region (F(1, 31) = 6.41, P = 0.017) (Fig. 4.1B). The most common native continent for introduced Monsteroideae was the American continent (n = 21), followed by Asia (n = 8) and Oceania (n = 7) (Fig. 4.1C). However, continent of origin was not significantly related to invasion status (F(4, 37) = 0.81, P = 0.52). Species that were introduced into more regions had a higher incidence of naturalizing and invading than those that were introduced into fewer regions ($X^2 = 10.98$, df = 1, P = 0.00091) (Fig. 4.1D).

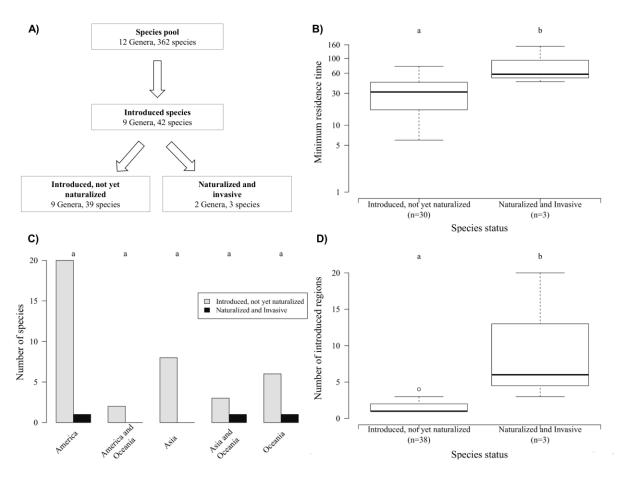


Fig. 4.1. Factors associated with invasion success in Monsteroideae. **A)** a narrative on the introduction history of the species pool; **B)** the minimum residence time; **C)** country of origin; and **D)** number of introduced regions. Different lower-case letters denote significant differences (P < 0.05) amongst invasion categories.

Weed risk assessment of *M. deliciosa* in South Africa

Out of the 49 questions in the weed risk assessment, 39 were answered based on published literature and online databases (Appendix 10). *Monstera deliciosa* scored a total of 7 points, with the biology/ecology of the species contributing the most to the total score. This score portrays a high probability of invasion and would have resulted in the species being rejected in a pre-border evaluation. In addition, we also attempted to perform an impact-based risk assessment following the generic impact scoring system (Nentwig *et al.*, 2016). However, this scoring system relies on published evidence of environmental and socio-economic impact and after searching the literature we could not find any reports of impact.

Potential global distribution of M. deliciosa based on climate

The MaxEnt model exhibited a high degree of predictive accuracy (AUC = 0.90 ± 0.0041 95% CI). M. deliciosa occurrence is predicted across much of the tropical countries and a few temperate countries. In particular, Mexico (the species' native range), the west coast of the United States, Hawaii, Chile, parts of Argentina and Brazil, England, Ireland, Spain, France, Italy, southern Africa, Madagascar, Iran, Nepal, Bhutan, west coast of India, parts of Southeast Asia, large parts of China, south Korea, Japan, Taiwan, Australia, and New Zealand (Fig. 4.2). Analysis of variable contribution indicates that among the four climatic parameters, annual mean temperature was the most influential predictor of *M. deliciosa* occurrence (84.8%), followed by temperature annual range (9.4%), precipitation seasonality (3.8%), and annual precipitation (2%). In addition, the relationship between the most influential predictor (annual mean temperature) and probability of occurrence is bell shaped (Appendix 11). This indicates reduced suitability as the annual mean temperatures shift from the optimums of 10-20 °C. This is particularly observable in the lowest and highest temperature ranges. Additionally, suitable climate ranges for the remaining variables comprised regions experiencing 12-28 °C as their temperature annual range; an increasing trend was observed for precipitation seasonality with a spike in regions receiving more than 110 mm of rain seasonally; and the likelihood of suitability increased in regions experiencing 1,000-4,000 mm of rain annually but suitability decreased in regions experiencing more than 4,000 mm of rain which could be indicative of winter rainfall regions.

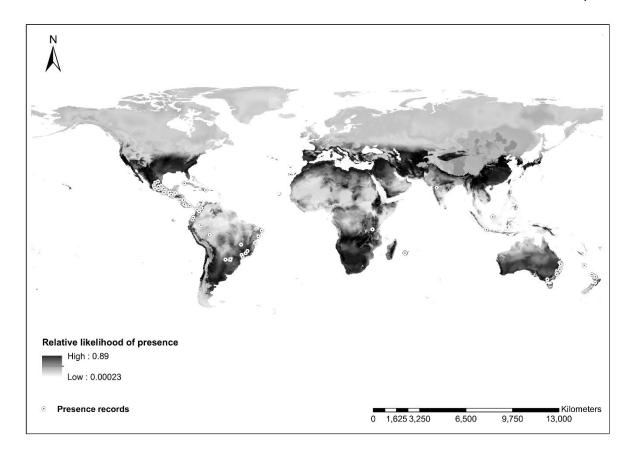


Fig. 4.2. Predicted global distribution of *M. deliciosa* based on occurrences in both its native and introduced ranges. Occurrences are indicated by white dots and projected presences are shown on a gradient from dark shading (high suitability) to white shading (low suitability).

A qualitative assessment of an invasive population in the Limpopo Province, South Africa

A densely scattered population was found growing and spreading at the site, covering approximately 1.4 ha (Fig. 4.3). The garden was established in 1948 at this site and *M. deliciosa* is believed to be grown here since the creation of the garden (68 years ago, I. Gaigher, pers. comm.). This species is now the dominant plant in the garden (Fig. 4.4). The plants have moved outside the garden through dispersal by vegetative means which is occurring at a very slow rate since they are currently not too far from the garden area. The garden has a lot of trees and is therefore suitable to support the growth and spread of *M. deliciosa* within the garden, and the dense clusters of trees in the Bushveld surrounding the garden also provide sufficient host structures. This site also receives a lot of shade since most of the plants were under 40–80% shade.

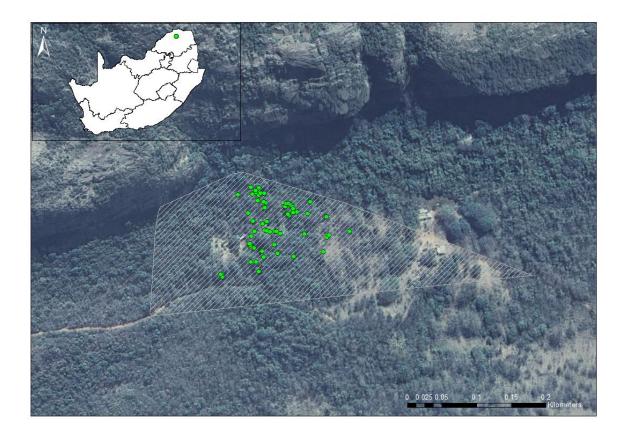


Fig. 4.3. The naturalized *M. deliciosa* population on the Soutpansberg Mountains in the Limpopo province, South Africa. The striped area represents the total area surveyed (~59331.50 m²) and the green dots depicts where we found the plants.

On the northern slopes of the garden (i.e. dry soils) we observed that *M. deliciosa's* distribution clearly follows an irrigation system occurring along the garden just outside the garden fence. On the southern slopes (i.e. wetter soils), the plants are much more abundant and dense and they are slowly moving downhill vegetatively, but the plants are only concentrated along the waterway. Over the years the landowner has tried to restrict the plants to the garden and commented that manual removal of the plants is effective. *M. deliciosa* can be cleared by chopping off the base stem and removing the rooted base, as well as, removing any aerial roots hanging down. The rest of the plant will then slowly die in the trees and/or on the ground.

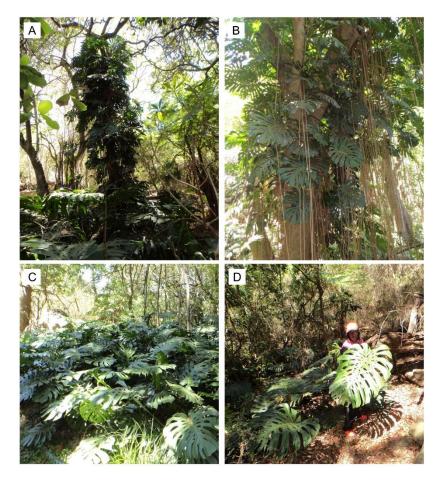


Fig. 4.4. *Monstera deliciosa* in Limpopo, South Africa. **A)** Climbing 10 m high, **B)** aerial roots growing downward out of the thick stems and taking root where they touch the ground, **C)** creeping densely on the ground, and **D)** showing off its massive leathery leaves with deep splits and oblong holes.

DISCUSSION

Many different mechanisms promoting invasions have been proposed, however recent studies have shown that invasions are context dependent (González-Moreno et al., 2014; Kueffer et al., 2013; Perkins and Nowak, 2013; Pyšek et al., 2012). These context dependent effects include interactions between climate, species traits, introduction history, conditions of the local habitat, and human activity. We support the argument that many cases are different, but strongly advocate that future research which aims to identify factors facilitating plant invasions should focus on species in particular functional groups (Moodley et al., 2013; Moodley et al., 2016a; Novoa et al., 2015).

In general, species within the Monsteroideae subfamily are likely to naturalize and become successful invaders if they have been introduced for a longer period of time. The likelihood of alien species becoming naturalized increases with residence time because the longer a species is present in a specific region, more propagules can be produced and dispersed (Pyšek and Jarošík, 2005; Pyšek *et al.*, 2011). Additionally, Monsteroideae species that are introduced to several new regions are more likely to overcome the naturalization and invasion barriers. This can be attributed to species having a better chance of landing in localities that are suitable for establishment if they are introduced across a wider area (Lockwood *et al.*, 2005b). Similar patterns have been reported for other alien flora (Dawson *et al.*, 2009; Moodley *et al.*, 2013; Trueman *et al.*, 2010). Even though *M. deliciosa* possesses these introduction history traits, it is not considered a major invader.

Monstera deliciosa has a high impact risk in South Africa. This is largely driven by the species' intrinsic traits and undesirable attributes in particular (e.g. toxicity, climbing growth habit, tolerates infertile soils and shade). This suggests that intrinsic traits which have been shown to drive invasiveness are not sufficient, and *M. deliciosa* requires additional interactions to overcome the invasion barriers.

Globally, the species is generally present in climatic regions that are moderately to highly suitable (Fig. 4.2). In addition, *M. deliciosa* has a wide potential geographical range. Therefore, it is possible that the species may become an invader with impacts if introduced to regions that have highly suitable climatic conditions but are currently unoccupied by the species (i.e. such as Korea or Southwest China).

On a local scale we found that water availability was the most important limiting factor since the plants were confined around an irrigation system or a stream (i.e. human intervention). Additionally, though the population occurs in a moderately suitable region (50–70% relative likelihood of presence), human intervention was necessary to sustain the population. Furthermore, while the population is naturalized and spreading within the modified landscape, it is not yet widespread. Taking into consideration the naturalized status as well as the restricted spread, the population can be classified as D1 under the Blackburn scheme (Blackburn *et al.*, 2011a). This

status denotes that the population is self-sustaining with individuals surviving a significant distance from the original point of introduction.

Following an introduction event, many species are not immediately successful at establishing and it often takes a considerable amount of time for an invasive species to transition to an exponential growth phase in the new geographical area. Herbaceous invasive plants can show a delay in invasion in the order of decades (Pyśek and Prach, 1993). Initially, growth rate is seemingly slow, but after a trigger there may be rapid population expansion. This is either due to the slow initial phase of exponential growth from small populations or a true lag phase (Aikio *et al.*, 2010). Understanding this dynamic is critical for management success (Essl *et al.*, 2011; Wilson *et al.*, 2017).

The determinants for the naturalization success of *M. deliciosa* at the Limpopo site comprise a suite of characters. This includes the undesirable attributes, climatically suitable regions, and a continuous source of water, which was facilitated by human-mediated opportunities in the Limpopo population. In the absence of these characteristics, populations remain in a lag phase. Similar aspects inducing a lag phase were identified for other taxa (Larkins, 2012).

Another example of a species that overcame a lag phase, even though it possessed all the traits of an invader, is *Banksia ericifolia*. This species required a trigger, such as disturbance, to influence the species establishment, growth and spread. A large population (n = 100 plants) in the Western Cape province, South Africa, had a sufficiently long residence time (35 years) but remained under the invasion radar until multiple fire events caused the population to spread rapidly and form dense monotypic stands (Geerts *et al.*, 2013). Consequently, this highlights that the lag phase is the best time to eradicate an invasive species population.

Using several examples, the role of failed invasions is also clearly demonstrated by a single species that can be highly invasive at one site and either fail or have minor impact at another (Zenni and Nuñez, 2013). However, much is still unknown on this crucial topic and such instances are often difficult to detect. Nevertheless, more detailed studies on invasion failures (e.g. why invasions are unsuccessful and what

changes in the natural landscape contribute to these failures) are required to advance our understanding of invasions.

Even though we identified several reasons why *M. deliciosa* should be invasive such as; widespread planting (globally and locally); non-reliance on pollinators and dispersal vectors in the introduced region due to vegetative reproduction; and a high likelihood of invasion risk, there is still a question of why *M. deliciosa* is in a lag phase and what will trigger an exponential phase. Species may establish in an area because they are better competitors than the resident species, but in order to become dominant, their competitive advantage must be persistent. We postulate that this is where *M. deliciosa* is failing. The species can become established and densified within a restricted area over a period of time, but its growth rate may be limiting its ability to spread fast enough. It is probable that given sufficient time and if left unmanaged the species may become an invader, however this should be explored in future studies.

CONCLUSION

Our analyses highlights that there are a number of characters, both of *M. deliciosa* (e.g. the species contains characteristics that pose a high invasion risk) and of the introduced habitats (e.g. warm temperatures and sufficient water) that increases the probability of the species naturalizing. However, even though these attributes favoured successful naturalization, the species is not a predominant invader. Moreover, when populations have naturalized it appears that manual control is effective and cost-efficient. Therefore, we recommend that *M. deliciosa* should not be regulated under South Africa's National Environmental Management: Biodiversity Act (10/2004) Alien and Invasive Species regulations. Given the sufficiently long minimum residence time, high propagule pressure, and no records of invasiveness or impacts elsewhere in the globe, we would also suggest that in many situations the species should be considered as "safe" and added to green lists of species that, despite lots of opportunities to invade, have never caused significant negative impacts (Dehnen-Schmutz, 2011).

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Chapter 5: Synthesis

THEORY

This thesis uncovered patterns, processes and invasion risks of a very popular horticultural plant group, the Araceae, which is not well represented in the invasion biology literature. Comparable to other model groups (e.g. Australian acacias, Cactaceae, Eucalyptus, *Pinus* and Proteaceae), the global distribution of the Araceae has been radically changed by humans in the last few centuries. However, in contrast to these groups, very little is known about the invasion status of such a widely introduced family, as well as the determinants underlying species' introduction, naturalization and invasion. The findings presented in this thesis address these knowledge gaps and assist in developing management approaches. Each study evaluated determinants of invasions at different stages along the introduction-naturalization-invasion (INI) continuum. The following predictions were tested:

- (i) A taxonomic group approach which explores invasiveness across the INI continuum to reveal specific mechanistic correlates within the family, as well as correlates that similarly influence other model groups. To identify these correlates I created the first comprehensive species inventory of Araceae using databases and literature sources; described the invasion status of all species; identified whether factors associated with the native range characteristics, introduction dynamics and biological traits influenced INI success and whether this varied for different life forms; and predicted which species have the potential to become invasive in future (Chapter 2).
- (ii) Invasion history elsewhere and/or impact elsewhere are reliable predictors of potential invasiveness and invasibility (Blackburn *et al.*, 2014; Williamson, 1996; Wilson *et al.*, 2011). For that reason, the global invasive status of *Epipremnum aureum* incited an evaluation of the species' invasion risk and management in South Africa, and was also the first invasion study of this species. To determine the potential risk posed by *E. aureum*, I delineated the current distribution of the species in South Africa; described factors that have contributed to its successful invasion;

identified the threat *E. aureum* poses globally and in South Africa; tested best management practices to control the species; and provided a recommendation on regulating the species (Chapter 3).

(iii) Invasion failure can occur at any stage of the INI continuum (Blackburn et al., 2011a; Zenni and Nuñez, 2013). Consequently, understanding why some alien plant species fail to become invasive provides insights on the relative importance of drivers of invasiveness and invasibility, and also help inform suitable management strategies. To unravel why *Monstera deliciosa* has failed to invade in much of its novel ranges, I assessed whether the popular hypotheses related to introduction history influences the invasion status of species within the Monsteroideae clade; and at a regional scale, I classified the invasion risk of *M. deliciosa* in South Africa; and on a local scale, I described potential conditions under which *M. deliciosa* can become naturalized (Chapter 4).

SUMMARY OF MAJOR FINDINGS

The Araceae conform to some, but not all, of the emerging generalizations in the invasion biology literature (Chapter 2; Table 5.1). My findings closely parallel those in many other studies; Araceae species that have been widely introduced (i.e. high propagule pressure) and which have large native range sizes were more likely to be invasive. However, unlike many other groups, there was little evidence of a link between invasiveness and regeneration mechanism (i.e. by seed, vegetative or both). Instead, there was a significant effect of plant life form and pollinator syndrome. Moreover, the importance of these factors varied across the INI continuum. I found that species classified as hydrophytes were more likely to overcome the introduction and naturalization barriers, and these species were mainly used as ornamentals. In addition, species requiring specialized pollinator types (e.g. flies and beetles) were able to overcome the invasion barriers, and this factor might be specific to Araceae. The type of habitat a species occupied in its native range was also an important correlate of introduction and naturalization success. Species native to humid regions, such as mediterranean forests and temperate mixed forests, were more likely to be introduced and become naturalized. Furthermore, I identified nine monophyletic groups that comprise species that are not currently invasive but have a high risk of becoming invasive in future, and I recommend a precautionary

approach be taken for these species. This includes species belonging to the following clades: *Alocasia*, *Amydrium*, *Ariopsis*, *Arum*, *Caladium*, *Cryptocoryne*, *Gymnostachys*, Lemnoideae, and *Peltandra*.

The global invasion patterns of Epipremnum aureum are mirrored in South Africa (Chapter 3; Table 5.1). In particular, the KwaZulu-Natal province of South Africa appears to be an invasion hotspot for *E. aureum*, with 78 naturalized populations and 321 cultivated populations. The naturalized populations consist of ~187,000 plants over ~3 hectares. Many of these extremely successful populations comprised plants as tall as the trees they were growing on, and were often found flourishing in dump sites, along roadsides or as a result of escaping cultivation. In addition, E. aureum has a high probability of expanding its current range along the coastal regions of South Africa, as well as, into neighbouring countries on Africa's eastern seaboard. Due to the invasion threat of the species, I recommend that all plants outside cultivation be removed, but current garden plantings could remain (though are not replaced). To control populations outside of cultivation, I found that applying herbicides to freshly cut stems effectively reduced plant growth. In view of that, I proposed that E. aureum should be listed as category 3 under South Africa's National Environmental Management: Biodiversity Act (10/2004) Alien and Invasive Species regulations, i.e. naturalized populations need to be managed, existing cultivated indivudals may remain, but the plant may not be propagated or sold in future.

It is well acknowledged that not all species become invasive when introduced to novel systems and the illustrious *Monstera deliciosa* distinctly falls into this category (Chapter 4; Table 5.1). Following an introduction event it often takes a considerable amount of time for an invasive species to transition to an exponential growth phase in the new geographical area. Globally, *M. deliciosa* has been subjected to a long period of time in the introduced regions, high propagule pressure, and suitable climatic conditions, while possessing undesirable species attributes, yet the species has not been documented as invasive, neither has there been any impacts. A qualitative local scale investigation of the only recorded naturalized population in South Africa, highlights that, invasions are in essence human driven. At the site in Limpopo the climate is suitable, the propagules are plentiful but the species only

spreads where a continual water supply is available, and this pattern follows the irrigation systems. Furthermore, the species can be removed manually which is a very efficient and cost effective means of control. Taking all aspects into account, the sufficiently long minimum residence time, high propagule pressure, and no records of invasiveness or impacts elsewhere, I recommended that *M. deliciosa* should not be regulated under South Africa's National Environmental Management: Biodiversity Act (10/2004) Alien and Invasive Species regulations, but should rather be considered as "safe" and added to green lists of species that, despite lots of opportunities to invade, have never caused significant negative impacts.

Table 5.1. Hypotheses evaluated in playing a role in Araceae invasions for each chapter and their associated traits tested. It is important to note that these findings vary across the INI continuum, as well as spatially and temporally. Empty cells indicate hypotheses not included in a chapter.

	Chapter 2	Chapter 3	Chapter 4
Species			
1. Tens rule	15% of the species pool have been introduced, of which 10% have naturalized, and 70% of the naturalized species are invasive		
2. Traits of an ideal weed	The number of native floristic regions (i.e.proxy for native range size), life form (e.g. hydrophytes), pollinator type, native floristic	Growth habit (climbing and creeping), height, and vegetative reproduction	Continent of origin

		regions (e.g. Polynesia), flower sexuality, habitat type (e.g. Mediterranean forests and temperate mixed forests), and the number of uses		
3.	Propagule pressure	The number of introduced regions (i.e. proxy for propagule pressure)	The number of stems in a population (i.e. proxy for propagule pressure)	The number of introduced regions (i.e. proxy for propagule pressure)
4.	Residence time		_	Minimum residence time (i.e. the year in which the first herbarium specimen was collected)
Sit	es			
1.	Resource availability	-	Habitat suitability (i.e. temperature and precipitation)	Habitat suitability (i.e. temperature and precipitation) and water supply
2.	Disturbances	_	Land-use types (i.e. canal, garden escapee, natural vegetation, open area, roadside,	_
			transformed	

vegetation,

vacant land or wasteland)

Pathways

1. Intentional	Phytoremediation	_	_
release			
2. Escape from	Agroforestry, food	Cultivation	Cultivation
containment	source, fibre		
	production,		
	horticulture, and		
	medicine		

IMPLICATIONS FOR THE THEORY OF INVASION BIOLOGY

In general, the analytical, experimental and observational aspects of this thesis supported the invasion biology literature which shows that invasions are complex processes synergistically driven by factors associated with introduction history, intrinsic species traits, characteristics of the recipient environment, and anthropogenic influences. What is more, the role of these factors differed at different stages of the INI continuum, and this facet is lacking in the literature.

There are a few universal attributes associated with plant invasions, the most common being propagule pressure, native range size and residence time (Hui and Richardson, 2017; Pyšek *et al.*, 2009a; Pyšek *et al.*, 2015; Richardson and Pyšek, 2012). These findings are also supported by Araceae. However, progress towards understanding the invasion process has largely been made by examining these invasion correlates, in isolation, rather than linking their degree of success to particular stages of the continuum. As a result, robust conclusions about which species will make the shift from introduced to naturalized and invasive, as well as the mechanisms driving these transitions, remain somewhat elusive. These comparisons provide critical insights into the transitions along the INI continuum and this thesis helps to bridge this gap. This approach helped to identify processes relevant for understanding invasions and their underlying mechanisms.

In addition to the invasion stages, the patterns of plant invasions and the mechanisms which generate these patterns, vary across spatial scales (Hamilton *et al.*, 2005; Pauchard and Shea, 2006; Pyšek and Hulme, 2005). Considering spatial scales in invasion studies offered insights into appropriate management strategies. For instance, if I only looked at *Monstera deliciosa* on a local scale in South Africa, I would have initially deemed the species as invasive. However, taking into account the global invasion dynamics of the species, fittingly changed the status to being a low invasion risk, and compelled me to identify the fine scale drivers of invasions.

Lastly, analyzing a single taxonomic or functional group provides a relatively accurate understanding of the mechanisms facilitating invasions. For example, specialized pollination syndromes were identified as a driver of invasiveness in Araceae, and this is unique to the family. Vegetative reproduction has been shown to be a common predictor of invasiveness in other groups (Kolar and Lodge, 2001; Lloret *et al.*, 2005a), but it was not significant in Araceae (chapter 2). In addition, a comprehensive look at two case study species with different invasion patterns revealed important information on species invasion, such as why some species become invasive (chapter 3 – primarily driven by human-mediated disturbances) while others fail (chapter 4 – lag phase). As a result, future studies need to continue looking at different groups in order to develop robust generalizations in invasion biology.

IMPLICATIONS FOR METHODOLOGY

This project successfully revealed the suitability of various approaches when studying processes that promote invasions. I demonstrated that collating a comprehensive species list, compiling trait databases, mapping species distributions, conducting population surveys and performing field experiments are useful for understanding invasion dynamics. In addition, risk assessments and species distribution models are also valuable tools for, among other things, informing management.

IMPLICATIONS FOR INVASIVE ALIEN SPECIES MANAGEMENT

Whilst research in invasion biology has progressed rapidly over the last few decades, a recurring criticism is linking existing knowledge into policies and management strategies (Hulme, 2006; Lodge, 1993). The results of this thesis have practical implications for invasion biologists, conservation managers and legislation. In chapter 2, I predicted which species have the potential to become invasive in the future and therefore recommended a precautionary approach be taken for these species. In chapter 3, I recommended that *E. aureum* be regulated, and in chapter 4, I recommended that *M. deliciosa* is reasonably innocuous and can continue being introduced and planted.

FUTURE RESEARCH

A stage-based approach to improve our understanding of the factors important throughout the invasion process should consider the themes proposed below (in combination):

- Identification of species traits combined with characteristics of the recipient habitats. In the global study, I established that hydrophytes pose a greater invasion risk than most other life forms. One of the limitations of this study was the exclusion of dispersal vectors due to the lack of available information. The inclusion of this trait will identify which trait is more important in driving hydrophyte invasiveness, regeneration mechanism or dispersal vectors;
- Research focused on particular taxonomic or functional groups;
- Studies should also consider traits of non-introduced species. This was a
 major limitation in this thesis. This information will be crucial for refining
 methods of predicting which species are likely to become invasive which will
 facilitate prevention methods, as well as better management of invasions;
- Ignoring failed invasions may hinder our understanding of the invasion process and these types of studies are still rare. For that reason, we should focus more effort on compiling and analyzing data on failed invasions. In the local case study of a failed invasion one of the major limitations was the lack of additional *M. deliciosa* localities for comaprisons. Other localities would have significantly strengthened the study.

CONCLUSION

The study of invasion biology contributes to a better understanding of processes that facilitate or limit invasions. This thesis illustrated that the mechanisms associated with invasiveness differ between taxa and across the INI continuum, therefore group and stage-specific analyses are essential. Despite the advances made through this work, much remains to be done to improve our knowledge and control of invasive alien species. To achieve the goal of predicting successful invasions, the recommendations mentioned above (i.e. spatial scale, invasion stage, species traits and characteristics of the recipient habitat) must be jointly accounted for in order to increase our predictive power. I strongly recommend that future studies use a similar framework for other plant groups.

These types of studies will provide a better understanding of why some introduced species become invasive while others fail, and will ultimately assist in managing biological invasions. In addition, similar studies of other plant groups are necessary to confirm the general applicability of these results. As more complete phylogenies and better knowledge of traits become available, these analyses are likely to become increasingly sophisticated and able to produce valuable insights into risk assessments. The work presented in this thesis contributes to understanding the causes and mechanisms of plant invasions and addresses questions of species invasiveness and community invasibility. Consequently, this research improves our ability to predict and manage the invasion risk of existing, emerging and potential invasive Araceae species, which is the ultimate research mission in invasion biology.

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Appendices

Appendix 1. Eight characteristics used to construct phenograms for invasive genera.

Characteristics	Reason for selection	Binary code
Number of native regions	Species occurring over a larger range have a greater chance of becoming introduced and naturalized (proxy for range size).	Native regions ≥ 3 = 1; Native regions < 3 = 0
Plant life form	Chamaephytes, geophytes, helophytes, hemicryptophytes or hydrophytes were found to be the most important life forms for Araceae to overcome the introduction and naturalization barriers.	Chamaephytes, geophytes, helophytes, hemicryptophytes or hydrophytes = 1; Other life forms = 0
Pollinator type	Fly pollinated species have a higher likelihood of being introduced by humans.	Fly pollinated = 1; Other pollinators = 0
Native floristic regions	Species native to the Polynesian province had a higher likelihood of being introduced.	Polynesian = 1; Non- Polynesian = 0
Flower sexuality	Unisexual flowers were favoured over bisexual flowers.	Unisexual = 1; Bisexual = 0
Habitat	Araceae genera are concentrated in the tropics. Species from tropical moist forests have a greater chance of being introduced.	Tropical moist forests = 1; Other habitats = 0
Number of introduced regions	Species introduced into more regions have a greater chance of becoming naturalized and invasive (proxy for propagule pressure).	Introduced regions ≥ 3 = 1; Introduced regions < 3 = 0
Number of uses	Species with more uses overcame the naturalization barriers.	Number of uses $\geq 3 =$ 1; Number of uses $< 3 =$ 0

Appendix 2. A

comprehensive species checklist developed for Araceae in 2013– 2014.

Scientific name

Aglaodorum griffithii Aglaonema brevispathum Aglaonema chermsiriwattanae Aglaonema cochinchinense Aglaonema commutatum var.commutatum Aglaonema commutatum var. elegans Aglaonema commutatum var. maculatum Aglaonema commutatum var. warburgii Aglaonema cordifolium Aglaonema costatum Aglaonema crispum Aglaonema densinervium Aglaonema flemingianum Aglaonema hookerianum Aglaonema marantifolium Aglaonema modestum Aglaonema nebulosum Aglaonema nitidum Aglaonema ovatum Aglaonema philippinense var. philippinense Aglaonema philippinense var. stenophyllum Aglaonema pictum Aglaonema pumilum Aglaonema rotundum Aglaonema simplex Aglaonema tricolor Aglaonema vittatum Alloschemone inopinata Alloschemone occidentalis Alocasia acuminata Alocasia aequiloba Alocasia alba Alocasia arifolia Alocasia atropurpurea Alocasia augustiana Alocasia baginda Alocasia balgooyi Alocasia beccarii Alocasia boa Alocasia boyceana

Alocasia brancifolia

Alocasia brisbanensis

Scientific name Alocasia cadieri Alocasia celebica Alocasia chaii Alocasia clypeolata Alocasia cucullata Alocasia culionensis Alocasia cuprea Alocasia decipiens Alocasia decumbens Alocasia devansayana Alocasia fallax Alocasia flabellifera Alocasia flemingiana Alocasia fornicata Alocasia gageana Alocasia grata Alocasia hainanica Alocasia heterophylla Alocasia hollrungii Alocasia hypnosa Alocasia hypoleuca Alocasia infernalis Alocasia inornata Alocasia jiewhoei Alocasia kerinciensis Alocasia lancifolia Alocasia lauterbachiana Alocasia lecomtei Alocasia longiloba Alocasia macrorrhizos Alocasia maquilingensis Alocasia megawatiae Alocasia melo Alocasia micholitziana Alocasia miniuscula Alocasia monticola Alocasia navicularis Alocasia nebula Alocasia nicolsonii Alocasia nycteris Alocasia odora Alocasia pangeran Alocasia peltata Alocasia perakensis Alocasia portei Alocasia princeps Alocasia principiculus Alocasia puber Alocasia puteri Alocasia pyrospatha Alocasia ramosii Alocasia reginae

Alocasia reginula

Alocasia reversa

Alocasia robusta

Alocasia sanderi

Alocasia ridleyi

Scientific name Alocasia sarawakensis Alocasia scabriuscula Alocasia scalprum Alocasia simonsiana Alocasia sinuata Alocasia suhirmaniana Alocasia venusta Alocasia wentii Alocasia wongii Alocasia zebrina Ambrosina bassii Amorphophallus aberrans Amorphophallus abyssinicus subsp. abyssinicus Amorphophallus . abyssinicus subsp. akeassii Amorphophallus abyssinicus subsp. unyikae Amorphophallus albispathus Amorphophallus albus Amorphophallus amygdaloides Amorphophallus andranogidroensis Amorphophallus angolensis subsp. angolensis Amorphophallus angolensis subsp. maculatus Amorphophallus angulatus Amorphophallus angustispathus Amorphophallus ankarana Amorphophallus annulifer Amorphophallus antsingyensis Amorphophallus aphyllus Amorphophallus asper Amorphophallus asterostigmatus Amorphophallus atrorubens Amorphophallus atroviridis Amorphophallus bangkokensis Amorphophallus barthlottii Amorphophallus baumannii Amorphophallus beccarii Amorphophallus bequaertii Amorphophallus bonaccordensis Amorphophallus borneensis Amorphophallus boyceanus Amorphophallus brachyphyllus Amorphophallus brevispathus Amorphophallus bufo Amorphophallus bulbifer Amorphophallus calabaricus subsp. calabaricus

Scientific name

Amorphophallus cirrifer Amorphophallus coaetaneus Amorphophallus commutatus Amorphophallus consimilis Amorphophallus corrugatus Amorphophallus costatus Amorphophallus coudercii Amorphophallus croatii Amorphophallus curvistylis Amorphophallus dactylifer Amorphophallus declinatus Amorphophallus decus-silvae Amorphophallus discophorus Amorphophallus dracontioides Amorphophallus dunnii Amorphophallus dzui Amorphophallus eburneus Amorphophallus echinatus Amorphophallus eichleri Amorphophallus elatus Amorphophallus elegans Amorphophallus elliottii Amorphophallus erythrororrhachis Amorphophallus excentricus Amorphophallus flotoi Amorphophallus forbesii Amorphophallus fuscus Amorphophallus galbra Amorphophallus gallaensis Amorphophallus gallowayi Amorphophallus gigas Amorphophallus glaucophyllus Amorphophallus gliruroides Amorphophallus glossophyllus Amorphophallus goetzei Amorphophallus gomboczianus Amorphophallus gracilior Amorphophallus gracilis Amorphophallus haematospadix Amorphophallus harmandii Amorphophallus hayi Amorphophallus henryi Amorphophallus hetterscheidii Amorphophallus hewittii

Amorphophallus hildebrandtii

Amorphophallus hohenackeri

Amorphophallus impressus

Amorphophallus incurvatus

Amorphophallus interruptus

Amorphophallus josefbogneri

Amorphophallus johnsonii

Amorphophallus hirsutus

Amorphophallus hirtus

Amorphophallus hottae

Amorphophallus

infundibuliformis

Amorphophallus calabaricus

Amorphophallus canaliculatus

Amorphophallus chlorospathus

Amorphophallus carneus

Amorphophallus cicatricifer

subsp. mayoi

Scientific name

Amorphophallus julaihii Amorphophallus juliae Amorphophallus kachinensis Amorphophallus kiusianus Amorphophallus konjac Amorphophallus konkanensis Amorphophallus koratensis Amorphophallus krausei Amorphophallus lacourii Amorphophallus lambii Amorphophallus lanuginosus Amorphophallus laoticus Amorphophallus lewallei Amorphophallus linearis Amorphophallus linguiformis Amorphophallus longicomus Amorphophallus longiconnectivus Amorphophallus Iongispathaceus Amorphophallus longistylus Amorphophallus longituberosus Amorphophallus lunatus Amorphophallus luzoniensis Amorphophallus lyratus Amorphophallus macrorhizus Amorphophallus mangelsdorffii Amorphophallus manta Amorphophallus margaritifer Amorphophallus margretae Amorphophallus maximus subsp. maximus Amorphophallus maximus subsp. fischeri Amorphophallus maxwellii Amorphophallus mekongensis Amorphophallus merrillii Amorphophallus mildbraedii Amorphophallus minor Amorphophallus mossambicensis Amorphophallus muelleri Amorphophallus mullendersii Amorphophallus myosuroides Amorphophallus mysorensis Amorphophallus napalensis Amorphophallus napiger Amorphophallus natolii Amorphophallus niahensis Amorphophallus nicolsonianus Amorphophallus obovoideus Amorphophallus obscurus Amorphophallus ochroleucus Amorphophallus ongsakulii Amorphophallus operculatus Amorphophallus opertus Amorphophallus verticillatus Amorphophallus paeoniifolius Amorphophallus vogelianus Amorphophallus palawanensis Amorphophallus xiei Amorphophallus paucisectus

Scientific name Amorphophallus pendulus Amorphophallus perakensis Amorphophallus pilosus Amorphophallus plicatus Amorphophallus polyanthus Amorphophallus prainii Amorphophallus preussii Amorphophallus prolificus Amorphophallus purpurascens Amydrium sinense Amorphophallus pusillus Amorphophallus putii Amorphophallus pygmaeus Amorphophallus ranchanensis Amorphophallus reflexus Amorphophallus rhizomatosus Amorphophallus richardsiae Amorphophallus rostratus Amorphophallus rugosus Amorphophallus sagittarius Amorphophallus salmoneus Amorphophallus saraburensis Amorphophallus saururus Amorphophallus scaber Amorphophallus schmidtiae Amorphophallus scutatus Amorphophallus serrulatus Amorphophallus sinuatus Amorphophallus sizemoreae Amorphophallus smithsonianus Amorphophallus sparsiflorus Amorphophallus spectabilis Amorphophallus staudtii Amorphophallus stuhlmannii subsp. stuhlmannii Amorphophallus stuhlmannii subsp. congoensis Amorphophallus subcymbiformis Amorphophallus sumawongii Amorphophallus suwidiianus Amorphophallus sylvaticus Amorphophallus symonianus Amorphophallus synandrifer Amorphophallus taurostigma Amorphophallus tenuispadix Amorphophallus tenuistylis Amorphophallus teuszii Amorphophallus thaiensis Amorphophallus tinekeae Amorphophallus titanum Amorphophallus tonkinensis Amorphophallus tuberculatus Amorphophallus variabilis Amorphophallus venustus

Scientific name

Amorphophallus yuloensis Amorphophallus yunnanensis Amorphophallus zenkeri subsp. zenkeri Amorphophallus zenkeri subsp. mannii Amydrium hainanense Amydrium humile Amydrium medium Amydrium zippelianum Anadendrum affine var. affine Anadendrum affine var. semivestitum Anadendrum angustifolium Anadendrum badium Anadendrum cordatum Anadendrum ellipticum Anadendrum griseum Anadendrum latifolium Anadendrum marcesovaginatum Anadendrum marginatum Anadendrum microstachyum Anadendrum montanum Anadendrum superans Anaphyllopsis americana Anaphyllopsis cururuana Anaphyllopsis pinnata Anaphyllum beddomei Anaphyllum wightii Anchomanes abbreviatus Anchomanes boehmii Anchomanes dalzielii Anchomanes difformis Anchomanes giganteus Anchomanes nigritianus Anthurium abelaezii Anthurium acanthospadix Anthurium acaule Anthurium acebeyae Anthurium achupallense Anthurium acutangulum Anthurium acutibacca Anthurium acutifolium var. acutifolium Anthurium acutifolium var. herrerae Anthurium acutissimum Anthurium acutum Anthurium aduncum Anthurium affine Anthurium agnatum Anthurium alatipedunculatum Anthurium alatum Anthurium albertiae Anthurium albidum Anthurium albispatha

Scientific name

Anthurium albovirescens Anthurium alcatrazense Anthurium alegriasense Anthurium algentryi Anthurium alluriquinense Anthurium alstonii Anthurium alticola Anthurium amargalense Anthurium ameliae Anthurium amnicola Anthurium amoenum var amoenum Anthurium amoenum var. humile Anthurium anceps Anthurium anchicayense Anthurium ancuashii Anthurium andicola Anthurium andinum Anthurium andraeanum Anthurium andreslovinense Anthurium angelopolinense Anthurium angosturense Anthurium angustatum Anthurium angustilaminatum subsp. angustilaminatum Anthurium angustilaminatum subsp. cibuserpentis Anthurium angustilobum Anthurium angustisectum Anthurium angustispadix Anthurium anorianum Anthurium antioquiense Anthurium antonioanum Anthurium antrophyoides Anthurium apanui Anthurium apaporanum Anthurium apiaense Anthurium arenasense Anthurium argyrostachyum Anthurium aripoense Anthurium arisaemoides Anthurium aristatum Anthurium armeniense Anthurium aroense Anthurium arusiense Anthurium asplundii Anthurium atamainii Anthurium atramentarium **Anthurium** atropurpureum var. atropurpur eum Anthurium atropurpureum var. arenicola Anthurium atropurpureum var. thomasii Anthurium atroviride Anthurium augustinum Anthurium aureum

Anthurium auritum

Anthurium albobueyense

Scientific name Anthurium austin-smithii Anthurium aylwardianum Anthurium baguense Anthurium bakeri Anthurium balaoanum Anthurium balslevii Anthurium barbacoasense Anthurium barclayanum Anthurium barreranum Anthurium barrieri Anthurium barrvi Anthurium basirotundum Anthurium bayae Anthurium beckii Anthurium bellum Anthurium beltianum Anthurium benktsparrei Anthurium bernardii Anthurium berriozabalense Anthurium berryi Anthurium besseae Anthurium betanianum Anthurium bicordoense Anthurium bimarginatum Anthurium binotii Anthurium birdseyanum Anthurium bittneri Anthurium bocainense Anthurium boekei Anthurium bogneri Anthurium bogotense Anthurium bonplandii subsp. bonplandii Anthurium bonplandii subsp. guayanum Anthurium boosianum Anthurium boudetii Anthurium brachvpodum Anthurium bradeanum Anthurium bragae Anthurium bredemeyeri Anthurium brenesii Anthurium brent-berlinii Anthurium brevipedunculatum Anthurium brevipes Anthurium breviscapum Anthurium brevispadix Anthurium brittonianum Anthurium bromelicola subsp. bromelicola Anthurium bromelicola subsp. bahiense Anthurium brownii Anthurium bucayanum Anthurium buchtienii Anthurium buganum

Anthurium bullianum

Anthurium bullosum

Anthurium burgeri

Scientific name Anthurium bushii Anthurium cabrerense Anthurium cabuyalense Anthurium cachabianum Anthurium cainarachense Anthurium caldodsonii Anthurium calimense Anthurium callejasii Anthurium caloveboranum Anthurium campii Anthurium camposii Anthurium canaliculatum Anthurium candolleanum Anthurium caperatum Anthurium caraboboense Anthurium caramantae Anthurium carchiense Anthurium cardenasii Anthurium carinatum Anthurium caripense Anthurium carneospadix Anthurium carnosum Anthurium carpishense Anthurium cartiense Anthurium cartilagineum Anthurium cataniapoense Anthurium caucanum Anthurium caucavallense Anthurium caulorrhizum Anthurium ceratiinum Anthurium ceronii Anthurium cerrateae Anthurium cerrobaulense Anthurium cerrocampanense Anthurium cerropelonense Anthurium cerropirrense Anthurium chacoense Anthurium chamberlainii Anthurium chamulense subsp. chamulense Anthurium chamulense subsp. oaxacanum Anthurium chiapasense subsp. chiapasense Anthurium chiapasense subsp. tlaxiacense Anthurium chimborazense Anthurium chinchipense Anthurium chinimense Anthurium chiriquense Anthurium chocoense Anthurium chorense Anthurium chorranum Anthurium chrysolithos Anthurium churutense Anthurium cinereopetiolatum Anthurium cipoense Anthurium circinatum

Scientific name Anthurium citrifolium Anthurium clarinervium Anthurium clarkei Anthurium clathratum Anthurium clavatum Anthurium clavigerum Anthurium cleistanthum Anthurium clidemioides subsp. clidemioides Anthurium clidemioides subsp. pacificum Anthurium coclense Anthurium cocornaense Anthurium coerulescens Anthurium cogolloanum Anthurium coleomischum Anthurium coleorrhiza Anthurium collettianum Anthurium collinsii Anthurium colonchense Anthurium colonense Anthurium colonicum Anthurium coloradense Anthurium comtum Anthurium concinnatum Anthurium concolor Anthurium coniunctum Anthurium consimile Anthurium consobrinum Anthurium conspicuum Anthurium constrictum Anthurium conterminum Anthurium corallinum Anthurium cordatotriangulum Anthurium cordatum Anthurium cordiforme Anthurium cordobense Anthurium cordulatum Anthurium coriaceum Anthurium coripatense Anthurium correae Anthurium corrugatum Anthurium cotejense Anthurium cotobrusii Anthurium cowanii Anthurium crassifolium Anthurium crassilaminum Anthurium crassinervium var. crassinervium Anthurium crassinervium var. caatingae Anthurium crassiradix var. crassiradix Anthurium crassiradix var. purpureospadix Anthurium crassitepalum Anthurium crassivenium Anthurium cremersii Anthurium crenatum Anthurium croatii

Scientific name Anthurium crystallinum Anthurium cuasicanum Anthurium cubense Anthurium cucullispathum Anthurium cultrifolium Anthurium cupreonitens Anthurium cupreum Anthurium cupulispathum Anthurium curicuriariense Anthurium curtispadix Anthurium curvilaminum Anthurium curvispadix Anthurium cuspidatum Anthurium cuspidiferum Anthurium cutucuense Anthurium cuyabenoense Anthurium cylindratum Anthurium cymbiforme Anthurium cymbispatha Anthurium daguense Anthurium darcyi Anthurium davidsei Anthurium davidsoniae Anthurium debilipeltatum Anthurium debilis Anthurium decurrens Anthurium deflexum Anthurium delannayi Anthurium dendrobates Anthurium denudatum Anthurium diazii Anthurium digitatum Anthurium diversicaudex Anthurium dolichocnemum Anthurium dolichophyllum Anthurium dolichostachyum Anthurium dombeyanum Anthurium dominicense Anthurium dorbayae Anthurium draconopterum Anthurium dressleri Anthurium dukei Anthurium durandii Anthurium dussii Anthurium dwyeri Anthurium dylanii Anthurium ecuadorense Anthurium effusilobum subsp. effusilobum Anthurium effusilobum subsp.pallidispadix Anthurium effusispathum Anthurium eggersii Anthurium eichleri Anthurium emarginatum Anthurium eminens subsp. eminens Anthurium eminens subsp. longispadix

Anthurium cirinoi

Scientific name Scientific name Scientific name Scientific name Anthurium ensifolium Anthurium gladiifolium Anthurium icanense Anthurium laucheanum Anthurium ericae Anthurium glanduligerum Anthurium idmense Anthurium lautum Anthurium ernestii Anthurium glaucophyllum Anthurium illepidum Anthurium lechlerianum var. ernestii Anthurium glaucospadix Anthurium iltisii Anthurium lehmannii **Anthurium** Anthurium lennartii Anthurium globosum Anthurium imperiale ernestii var. oellgaardii Anthurium gomesianum Anthurium impolitum Anthurium lentii Anthurium erskinei Anthurium gonzalezii Anthurium incomptum Anthurium leonianum Anthurium erythrostachyum Anthurium gracile Anthurium inconspicuum Anthurium leonii Anthurium esmeraldense Anthurium gracililaminum Anthurium espinae Anthurium incurvatum Anthurium leptocaule Anthurium gracilipedunculatum Anthurium incurvum Anthurium leuconeurum Anthurium eximium Anthurium gracilispadix Anthurium infectorium Anthurium leveaui Anthurium expansum Anthurium intermedium Anthurium lezamae Anthurium grande Anthurium exstipulatum Anthurium fasciale Anthurium grandicataphyllum Anthurium interruptum Anthurium Ihotzkyanum Anthurium fatoense Anthurium grandifolium Anthurium inzanum Anthurium libanoense Anthurium grex-avium Anthurium ionanthum Anthurium licium Anthurium faustomirandae Anthurium iramireziae Anthurium lievenii Anthurium gualeanum Anthurium fendleri Anthurium guanacense Anthurium isidroense Anthurium ligulare Anthurium fernandezii Anthurium filiforme Anthurium guanchezii Anthurium jaramilloi Anthurium lilacinum Anthurium quatemalense Anthurium jefense Anthurium limonense Anthurium flavescens Anthurium guayaquilense Anthurium jenmanii Anthurium lindenianum Anthurium flavidum Anthurium gustavii Anthurium jesusii Anthurium lindmanianum Anthurium flavolineatum Anthurium lineolatum Anthurium flavoviride Anthurium gymnopus Anthurium iilekii Anthurium flexile Anthurium hacumense Anthurium jimenae Anthurium linganii subsp. flexile Anthurium hagsaterianum Anthurium joaquinense Anthurium lingua **Anthurium** Anthurium halmoorei Anthurium johnmackii Anthurium linguifolium flexile subsp. muelleri Anthurium hamiltonii Anthurium llanense Anthurium johnsoniae Anthurium folsomianum Anthurium hammelii Anthurium iosei Anthurium Ilewellvnii Anthurium fontellanum Anthurium harrisii Anthurium julianii Anthurium Iloense Anthurium fontoides Anthurium hastifolium Anthurium julospadix Anthurium loefgrenii Anthurium foreroanum Anthurium hatschbachii Anthurium jureianum Anthurium lojtnantii Anthurium forgetii Anthurium hebetatilaminum Anthurium kajekai Anthurium longicaudatum Anthurium formosum Anthurium kallunkiae Anthurium longicuspidatum Anthurium fornicifolium Anthurium hebetatum Anthurium herthae Anthurium kamemotoanum Anthurium longifolium Anthurium fosteri Anthurium hieronymi Anthurium karstenianum Anthurium longigeniculatum Anthurium fragae Anthurium hinoideum Anthurium kastelskii Anthurium longipeltatum Anthurium fragrans Anthurium hodgei Anthurium kayapii Anthurium longipes Anthurium fragrantissimum Anthurium hoehnei Anthurium knappiae Anthurium longispadiceum Anthurium fraseri Anthurium hoffmannii Anthurium krukovii Anthurium longissimilobum Anthurium friedrichsthalii Anthurium longissimum Anthurium holm-nielsenii Anthurium kugkumasii Anthurium funiferum subsp. longissimum Anthurium holquinianum Anthurium kunayalense Anthurium furcatum Anthurium longissimum Anthurium kunthii Anthurium hookeri Anthurium fuscopunctatum subsp. nirquense var. kunthii Anthurium hornitense Anthurium fusiforme Anthurium longistamineum Anthurium kunthii Anthurium huacamayoense var. cylindricum Anthurium longistipitatum Anthurium gaffurii Anthurium huallagense Anthurium kusuense Anthurium longiusculum Anthurium galactospadix Anthurium huampamiense Anthurium lacerdae Anthurium loretense Anthurium galeanoae Anthurium huanucense Anthurium laciniosum Anthurium louisii Anthurium galeottii Anthurium huashikatii Anthurium lactifructum Anthurium lucens Anthurium galileanum Anthurium huautlense Anthurium lucidum Anthurium lancea Anthurium gaskinii Anthurium huberi Anthurium lancetillense Anthurium lucioi Anthurium gaudichaudianum Anthurium lancifolium Anthurium huixtlense Anthurium gehrigeri Anthurium lucorum var. lancifolium Anthurium humboldtianum Anthurium luschnathianum Anthurium geitnerianum Anthurium lancifolium subsp. humboldtianum Anthurium lutescens Anthurium genferryae Anthurium humboldtianum var. albifructum subsp. viridispadix Anthurium lutevnii Anthurium geniculatum Anthurium langendoenii Anthurium gentryi Anthurium humoense Anthurium langsdorffii Anthurium luxurians Anthurium giganteum Anthurium hutchisonii Anthurium lanjouwii Anthurium lygrum Anthurium hygrophilum Anthurium lynniae Anthurium ginesii Anthurium latemarginatum Anthurium ianthinopodum Anthurium maasii Anthurium giraldoi Anthurium latissimum

Scientific name

Anthurium macarenense Anthurium macbridei Anthurium macdanielii Anthurium machetioides Anthurium macleanii Anthurium macphersonii Anthurium macrocephalum Anthurium macrolonchium Anthurium macrophyllum Anthurium macropodum Anthurium macrospadix Anthurium macrourum Anthurium maculosum Anthurium madisonianum Anthurium magdae Anthurium magnificum Anthurium magnifolium Anthurium maguirei Anthurium malagaense Anthurium malianum Anthurium manabianum Anthurium mancuniense Anthurium manuanum Anthurium marense Anthurium margaricarpum Anthurium marginellum Anthurium mariae Anthurium maricense Anthurium marinoanum Anthurium marleenianum Anthurium marmoratum Anthurium martianum Anthurium masfense Anthurium maximiliani Anthurium maximum Anthurium megapetiolatum Anthurium melastomatis Anthurium membranaceum Anthurium mendietae Anthurium merlei Anthurium metallicum Anthurium miaziense Anthurium michelii Anthurium microphyllum Anthurium microspadix Anthurium minarum Anthurium mindense Anthurium miniatum Anthurium modicum Anthurium molaui Anthurium molle Anthurium montanum Anthurium monteverdense Anthurium monticola var. monticola Anthurium monticola var. attenuatum

Anthurium monzonense

Scientific name Anthurium moonenii Anthurium morae Anthurium morii Anthurium moronense Anthurium mostaceroi Anthurium mourae Anthurium multinervium Anthurium multisulcatum Anthurium munchiquense Anthurium myosuroides Anthurium myosurus Anthurium nakamurae Anthurium nangaritense Anthurium nanum Anthurium napaeum Anthurium narinoense Anthurium navasii Anthurium naviculare Anthurium nelsonii Anthurium nemorale Anthurium nemoricola Anthurium nervatum Anthurium nicolasianum Anthurium nigrescens Anthurium nigropunctatum Anthurium niqueanum Anthurium nitens Anthurium nitidulum Anthurium nitidum Anthurium nizandense Anthurium novitaense Anthurium nubicola Anthurium nutibarense Anthurium nymphaeifolium Anthurium obliquatum Anthurium oblongocordatum Anthurium obpyriforme Anthurium obscurinervium Anthurium obtusatum Anthurium obtusifolium Anthurium obtusilobum Anthurium obtusum subsp. obtusum Anthurium obtusum subsp. puntarenense Anthurium occidentale Anthurium ochranthum Anthurium ochreatum Anthurium ocotepecense Anthurium oerstedianum Anthurium oreodoxa Anthurium oreophilum Anthurium organense Anthurium orientale Anthurium orteganum

Anthurium ottobuchtienii

Anthurium ottonis

Anthurium ovatifolium

Scientific name

Anthurium oxyanthum Anthurium pichindense Anthurium oxybelium Anthurium pilonense Anthurium oxycarpum Anthurium oxyphyllum Anthurium oxystachyum Anthurium pachylaminum Anthurium pachyspathum Anthurium pageanum Anthurium palacioanum Anthurium palenquense Anthurium pallatangense Anthurium pallens Anthurium pallidicaudex Anthurium pallidiflorum Anthurium palmarense Anthurium palmatum Anthurium paludosum Anthurium panamense Anthurium panduriforme Anthurium papillilaminum Anthurium paradisicum Anthurium paraguasense Anthurium paraguayense var. paraguayense Anthurium paraguayense var. coroicoanum Anthurium parambae Anthurium parasiticum Anthurium pariense Anthurium parile Anthurium parvispathum Anthurium parvum Anthurium pastasanum Anthurium patens Anthurium pauciflorum Anthurium paucinerve Anthurium payaminoense Anthurium pedatoradiatum subsp. pedatoradiatum Anthurium pedatoradiatum subsp. helleborifolium Anthurium pedatum Anthurium pedunculare Anthurium pellucidopunctatum Anthurium peltatum Anthurium peltigerum Anthurium pulidoae Anthurium penae Anthurium pulverulentum Anthurium pendens var. pulverulentum Anthurium pendulifolium Anthurium penningtonii Anthurium pentaphyllum var. pentaphyllum Anthurium pentaphyllum var. bombacifolium Anthurium perijanum Anthurium perviride Anthurium pescadilloense

Scientific name

Anthurium pinkleyi Anthurium pirottae Anthurium pirrense Anthurium pittieri var. pittieri Anthurium pittieri var. fogdeniorum Anthurium pittieri var. morii Anthurium piurensis Anthurium plantagineum Anthurium platyglossum Anthurium platyrhizum Anthurium plowmanii Anthurium plurisulcatum Anthurium pluviaticum Anthurium podophyllum Anthurium pohlianum Anthurium polydactylum Anthurium polynervium Anthurium polyneuron Anthurium polyphlebium Anthurium polyschistum Anthurium polystictum Anthurium porcesitoense Anthurium potarense Anthurium pradoense Anthurium praealtum Anthurium pranceanum Anthurium prolatum Anthurium prominens Anthurium promininerve Anthurium protensum subsp. protensum Anthurium protensum subsp. arcuatum Anthurium pseudospectabile Anthurium psilostachyum Anthurium ptarianum Anthurium puberulinervium Anthurium puberulum Anthurium pucayacuense Anthurium pulcachense Anthurium pulchellum

Anthurium pulverulentum var. adsimile Anthurium punctatum Anthurium punkuyocense Anthurium purdieanum Anthurium purpureospathum Anthurium purpureum Anthurium queirozianum Anthurium quinindense Anthurium quinquenervium Anthurium quinquesulcatum

Anthurium petrophilum

Anthurium phyllobaris

Anthurium pichinchae

Scientific name
Anthurium quipuscoae
Anthurium radiatum
Anthurium radicans
Anthurium raimundii
Anthurium ramoncaracasii
Anthurium ramonense
Anthurium ramosense
Anthurium ramosii
Anthurium ranchoanum
Anthurium ravenii
Anthurium recavum
Anthurium redolens
Anthurium reflexinervium

Anthurium renexinervium
Anthurium regale
Anthurium remotigeniculatum
Anthurium remotum
Anthurium resectum
Anthurium resectum
Anthurium reticulatum
Anthurium retiferum
Anthurium rhizophorum
Anthurium rhodorhizum
Anthurium ribeiroi
Anthurium rigidifolium
Anthurium rimbachii
Anthurium riodocense
Anthurium riofrioi

Anthurium riograndicola
Anthurium riojaense
Anthurium rionegrense
Anthurium riparium
Anthurium rivulare
Anthurium rodrigueziae
Anthurium roezlii
Anthurium rojasiae
Anthurium roraimense
Anthurium roseospadix
Anthurium rotundatum

Anthurium rotundistigmatum
Anthurium rubrifructum
Anthurium rubrivellus
Anthurium rugulosum
Anthurium rupestre
Anthurium rupicola
Anthurium rzedowskii
Anthurium saccardoi
Anthurium sagawae
Anthurium sagittale
Anthurium sagittaria

Anthurium rotundilobum

Anthurium sagittatum
Anthurium sagittellum
Anthurium salgarense
Anthurium salvadorense
Anthurium salvinii
Anthurium samamaense
Anthurium sanctifidense

Scientific name Anthurium sanguineum Anthurium santaritensis Anthurium santiagoense Anthurium sapense Anthurium sarmentosum Anthurium sarukhanianum Anthurium scaberulum Anthurium scandens subsp. scandens Anthurium scandens subsp. pusillum Anthurium scherzerianum Anthurium schlechtendalii subsp. schlechtendalii Anthurium schlechtendalii subsp. jimenezii Anthurium schottianum Anthurium schunkei Anthurium sebastianense Anthurium seibertii Anthurium seleri Anthurium sellowianum Anthurium septuplinervium Anthurium shinumas Anthurium siccisilvarum Anthurium sidneyi Anthurium sierpense Anthurium signatum Anthurium silanchense Anthurium silverstonei Anthurium silvicola Anthurium silvigaudens

Anthurium silverstonei
Anthurium silvicola
Anthurium silvigaudens
Anthurium simonii
Anthurium simpsonii
Anthurium sinuatum
Anthurium siqueirae
Anthurium smaragdinum
Anthurium smithii
Anthurium sodiroanum

Anthurium soejartoi

subsp. soejartoi
Anthurium soejartoi
subsp. ascendens
Anthurium solitarium
Anthurium solomonii
Anthurium soukupii
Anthurium sparreorum
Anthurium spathiphyllum
Anthurium spathulifolium
Anthurium spectabile
Anthurium splendidum

Anthurium stephanii
Anthurium stipitatum
Anthurium straminopetiolum
Anthurium striatipes
Anthurium striatum
Anthurium striolatum
Anthurium stuebelii

Anthurium subaequans

Anthurium standleyi

Scientific name Anthurium subcarinatum Anthurium subcaudatum Anthurium subcoerulescens Anthurium subcordatum subsp. subcordatum Anthurium subcordatum subsp. chlorocardium Anthurium subhastatum Anthurium subovatum Anthurium subrotundum Anthurium subsagittatum Anthurium subscriptum Anthurium subsignatum Anthurium subtriangulare Anthurium subtrilobum Anthurium subtruncatum Anthurium subulatum Anthurium sucrii Anthurium sulcatum Anthurium superbum subsp. superbum Anthurium superbum subsp. brentberlinii Anthurium supianum Anthurium supraglandulum Anthurium suramaense Anthurium sylvestre Anthurium sytsmae Anthurium tacarcunense Anthurium tachiranum Anthurium talmonii Anthurium tamaense Anthurium tarapotense Anthurium tatei Anthurium teimosoanum Anthurium tenaense Anthurium tenerum Anthurium tenuicaule Anthurium tenuifolium Anthurium tenuispica Anthurium teribense Anthurium ternifolium Anthurium terryae Anthurium testaceum

Anthurium thompsoniae

Anthurium thrinax

Anthurium tikunorum

Anthurium tilaranense

Anthurium titanium

Anthurium tolimense

Anthurium tomasiae

Anthurium tonduzii

Anthurium tonianum

Anthurium torraense

Anthurium treleasei

Anthurium tremulum

Anthurium tricarinatum

Anthurium trianae

Anthurium timplowmanii

Scientific name Anthurium triciafrankiae Anthurium trifidum Anthurium trilobum Anthurium trinervium Anthurium triphyllum Anthurium trisectum Anthurium truncatulum Anthurium truncatum Anthurium truncicola Anthurium tsamajainii Anthurium tunquii Anthurium tutense Anthurium tysonii Anthurium uasadiensis Anthurium uleanum var. uleanum Anthurium uleanum var. nanayense Anthurium umbraculum Anthurium umbricola Anthurium umbrosum Anthurium unense Anthurium upalaense Anthurium urbanii Anthurium urvilleanum Anthurium utleyorum Anthurium validifolium Anthurium validinervium Anthurium vallense Anthurium vanderknaapii Anthurium variegatum Anthurium variilobum Anthurium vaupesianum Anthurium veitchii Anthurium velutinum Anthurium venadoense Anthurium venosum Anthurium ventanasense Anthurium verapazense Anthurium verrucosum Anthurium versicolor var. versicolor Anthurium versicolor var. azuayense Anthurium vestitum Anthurium victorii Anthurium vientense Anthurium vinillense Anthurium viridescens Anthurium viridispathum Anthurium vittariifolium Anthurium vomeriforme Anthurium wagenerianum Anthurium wallisii Anthurium walujewii Anthurium warintsense

Anthurium warocqueanum

Anthurium watermaliense

Anthurium wattii

Scientific name Anthurium weberbaueri Anthurium wedelianum subsp. wedelianum Anthurium wedelianum subsp. viridispadix Anthurium wendlingeri Anthurium werffii Anthurium werneri Anthurium whitmorei Anthurium willdenowii Anthurium willifordii Anthurium wintersii Anthurium wurdackii Anthurium xanthoneurum Anthurium xanthophylloides Anthurium yamayakatense Anthurium yarumalense Anthurium yatacuense Anthurium yetlense Anthurium yungasense Anthurium yurimaguense Anthurium yutajense Anthurium zappiae Anthurium zeneidae Anthurium zuloagae Anubias afzelii Anubias barteri var. barteri Anubias barteri var. angustifolia Anubias barteri var. caladiifolia Anubias barteri var. alabra Anubias barteri var. nana Anubias gigantea Anubias gilletii Anubias gracilis Anubias hastifolia Anubias heterophylla Anubias pynaertii Apoballis acuminatissima Apoballis belophylla Apoballis brevipes Apoballis grandiflora Apoballis hastifolia Apoballis javanica Apoballis longicaulis Apoballis mutata Apoballis okadae Apoballis ovata Apoballis rupestris Apoballis sagittifolia Aridarum borneense Aridarum burttii Aridarum caulescens Aridarum crassum

Aridarum incavatum

Aridarum minimum

Scientific name Aridarum montanum Aridarum nicolsonii Aridarum purseglovei Aridarum rostratum Ariopsis peltata Ariopsis protanthera Arisaema abei Arisaema aequinoctiale Arisaema agasthyanum Arisaema album Arisaema amurense Arisaema anomalum Arisaema aprile Arisaema aridum Arisaema asperatum Arisaema auriculatum Arisaema austroyunnanense Arisaema averyanovii Arisaema balansae Arisaema bannaense Arisaema barbatum Arisaema barnesii Arisaema bockii Arisaema bonatianum Arisaema bottae Arisaema brucei Arisaema calcareum Arisaema candidissimum Arisaema caudatum Arisaema chuanxiense Arisaema chumponense Arisaema ciliatum Arisaema clavatum Arisaema concinnum Arisaema condaoense Arisaema consanguineum subsp. consanguineum Arisaema consanguineum subsp. kelung-insulare Arisaema constrictum Arisaema cordatum Arisaema costatum Arisaema cucullatum Arisaema dahaiense Arisaema decipiens Arisaema dracontium Arisaema echinatum Arisaema echinoides Arisaema ehimense Arisaema elephas Arisaema enneaphyllum Arisaema erubescens Arisaema exappendiculatum Arisaema fargesii Arisaema filiforme Arisaema fimbriatum subsp. fimbriatum

Scientific name Arisaema flavum subsp. flavum Arisaema flavum subsp. tibeticum Arisaema formosanum Arisaema franchetianum Arisaema fraternum Arisaema galeatum Arisaema garrettii Arisaema ghaticum Arisaema grapsospadix Arisaema griffithii Arisaema hainanense Arisaema handelii Arisaema heterocephalum subsp. heterocephalum Arisaema heterocephalum subsp. okinawaense Arisaema heterophyllum Arisaema hunanense Arisaema ilanense Arisaema inclusum Arisaema intermedium Arisaema ishizuchiense subsp. ishizuchiense Arisaema ishizuchiense subsp. brevicollum Arisaema iyoanum Arisaema jacquemontii Arisaema jethompsonii Arisaema jingdongense Arisaema kawashimae Arisaema kerrii Arisaema kishidae Arisaema kiushianum Arisaema kuratae Arisaema lackneri Arisaema laminatum Arisaema leschenaultii Arisaema lichiangense Arisaema lidaense Arisaema lihengianum Arisaema limbatum Arisaema linearifolium Arisaema lingyunense Arisaema lobatum Arisaema longipedunculatum Arisaema macrospathum Arisaema maekawae Arisaema mairei Arisaema maximowiczii subsp. maximowiczii Arisaema maximowiczii subsp. tashiroi Arisaema maxwellii Arisaema meleagris Arisaema menglaense Arisaema microspadix Arisaema mildbraedii Arisaema minamitanii

Scientific name Arisaema monophyllum var. monophyllum Arisaema monophyllum var. atrolinguum Arisaema mooneyanum Arisaema muratae Arisaema muricaudatum Arisaema murrayi var. murrayi Arisaema murrayi var. sahyadricum Arisaema murrayi var. sonubeniae Arisaema nagiense Arisaema nambae Arisaema negishii Arisaema nepenthoides Arisaema nikoense var. nikoense Arisaema nikoense var. australe Arisaema nikoense var. kaimontanum Arisaema nilamburense Arisaema odoratum Arisaema ogatae Arisaema omkoiense Arisaema ornatum Arisaema ovale var. ovale Arisaema ovale var. inaense Arisaema ovale var. sadoense Arisaema pachystachyum Arisaema pallidum Arisaema parisifolia Arisaema parvum Arisaema pattaniense Arisaema penicillatum Arisaema petelotii Arisaema petiolulatum Arisaema pianmaense Arisaema pingbianense Arisaema polyphyllum Arisaema prazeri Arisaema propinguum Arisaema psittacus Arisaema quinatum Arisaema quinquelobatum Arisaema ramulosum Arisaema ringens Arisaema rostratum Arisaema roxburghii Arisaema rubrirhizomatum Arisaema ruwenzoricum Arisaema sachalinense Arisaema saddlepeakense Arisaema sarracenioides Arisaema saxatile Arisaema sazensoo Arisaema schimperianum

Arisaema minus

Arisaema fimbriatum

subsp. bakerianum

Scientific name

Arisaema scortechinii Arisaema seppikoense Arisaema serratum var. serratum Arisaema serratum var. izuense Arisaema serratum var. mavebarae Arisaema serratum var. suwoense Arisaema setosum

Arisaema siamicum Arisaema siangense Arisaema sikokianum Arisaema silvestrii

Arisaema sizemoreae Arisaema smitinandii

Arisaema sinii

Arisaema somalense Arisaema souliei Arisaema speciosum var. speciosum Arisaema speciosum var. mirabile Arisaema speciosum var. ziroense

Arisaema sukotaiense Arisaema taiwanense var. taiwanense Arisaema taiwanense var. brevipedunculatum

Arisaema tengtsungense Arisaema ternatipartitum Arisaema thunbergii subsp. thunbergii Arisaema thunbergii subsp. autumnale Arisaema thunbergii subsp. geomundoense Arisaema thunbergii subsp. urashima Arisaema tortuosum subsp. tortuosum Arisaema tortuosum var. neglectum Arisaema tortuosum subsp. sivadasanii Arisaema tosaense Arisaema translucens

Arisaema tsangpoense Arisaema tuberculatum Arisaema ulugurense Arisaema umbrinum Arisaema undulatifolium subsp. undulatifolium Arisaema undulatifolium subsp. uwajimense

Arisaema triphyllum

Arisaema utile Arisaema vexillatum Arisaema victoriae Arisaema wangmoense Arisaema wardii Arisaema wattii Arisaema wilsonii

Scientific name

Arisaema wrayi Arisaema xuanweiense Arisaema yamatense subsp. yamatense Arisaema yamatense subsp. sugimotoi Arisaema yanxianum Arisaema yunnanense Arisaema zhui

Arisarum proboscideum Arisarum simorrhinum Arisarum vulgare subsp. vulgare Arisarum vulgare subsp. clusii Arisarum vulgare subsp. hastatum Arophyton buchetii Arophyton crassifolium

Arophyton humbertii Arophyton pedatum

Arophyton rhizomatosum Arophyton simplex Arophyton tripartitum var. tripartitum Arophyton tripartitum var. masoalaense

Arum alpinariae Arum apulum Arum balansanum Arum besserianum Arum byzantinum Arum concinnatum

Arum creticum Arum cylindraceum subsp. cylindraceum Arum cylindraceum subsp. pitsyllianum

Arum cyrenaicum Arum dioscoridis var. dioscoridis Arum dioscoridis var. cyprium Arum dioscoridis var. philistaeum Arum dioscoridis var. syriacum Arum elongatum Arum euxinum Arum gratum

Arum hainesii Arum hygrophilum Arum idaeum Arum italicum subsp. italicum Arum italicum subsp. albispathum Arum italicum subsp. canariense Arum italicum subsp. neglectum Arum jacquemontii

Arum korolkowii

Arum maculatum

Arum lucanum

Arum megobrebi Arum nigrum Arum orientale subsp. orientale Arum orientale subsp. longispathum Arum palaestinum Arum pictum

Scientific name

Arum purpureospathum Arum rupicola var. rupicola Arum rupicola var. virescens Arum sintenisii Asterostigma cryptostylum

Asterostigma cubense Asterostigma lividum Asterostigma lombardii Asterostigma luschnathianum Asterostigma reticulatum Asterostigma riedelianum

Asterostigma tweedieanum Bakoa brevipedunculata Bakoa lucens

Bakoa nakamotoi

Biarum aleppicum Biarum angustatum Biarum auraniticum Biarum bovei Biarum carduchorum Biarum carratracense Biarum crispulum Biarum davisii Biarum dispar Biarum ditschianum

Biarum eximium Biarum fraasianum Biarum kotschvi Biarum marmarisense Biarum mendax Biarum olivieri Biarum pyramid var. pyrami

Biarum pyrami

var. serotinum Biarum rhopalospadix Biarum straussii Biarum syriacum Biarum tenuifolium subsp. tenuifolium

Biarum tenuifolium subsp. abbreviatum Biarum tenuifolium subsp. arundanum Biarum tenuifolium subsp. galianii Biarum tenuifolium subsp. idomenaeum Biarum tenuifolium subsp. zelebori

Bognera recondita Bucephalandra gigantea Scientific name

Bucephalandra motleyana Caladium andreanum Caladium bicolor Caladium clavatum Caladium coerulescens Caladium humboldtii Caladium lindenii Caladium macrotites Caladium picturatum Caladium praetermissum Caladium schomburgkii Caladium smaragdinum Caladium steyermarkii

Caladium ternatum Caladium tuberosum Calla palustris Callopsis volkensii Carlephyton diegoense Carlephyton glaucophyllum Carlephyton madagascariense

Cercestis afzelii

Cercestis camerunensis Cercestis congoensis Cercestis dinklagei Cercestis hepperi Cercestis ivorensis Cercestis kamerunianus Cercestis mirahilis Cercestis sagittatus Cercestis taiensis Chlorospatha amalfiensis

Chlorospatha antioquiensis Chlorospatha atropurpurea Chlorospatha besseae Chlorospatha betancurii Chlorospatha bogneri Chlorospatha callejasii Chlorospatha castula Chlorospatha cogolloi Chlorospatha corrugata Chlorospatha croatiana subsp. croatiana

Chlorospatha croatiana subsp. enneaphylla Chlorospatha cutucuensis Chlorospatha dodsonii Chlorospatha feuersteiniae Chlorospatha gentryi Chlorospatha hammeliana Chlorospatha hastifolia Chlorospatha ilensis Chlorospatha kolbii Chlorospatha kressii Chlorospatha lehmannii

Chlorospatha luteynii Chlorospatha macphersonii Chlorospatha mirabilis

Chlorospatha longipoda

Scientific name Scientific name Scientific name Scientific name Chlorospatha nicolsonii Cryptocoryne keei Culcasia seretii Dieffenbachia obscurinervia Chlorospatha planadensis Cryptocoryne lingua Culcasia simiarum Dieffenbachia oerstedii Chlorospatha ricaurtensis Cryptocoryne loeiensis Culcasia striolata Dieffenbachia olbia Colletogyne perrieri Cryptocoryne longicauda Culcasia tenuifolia Dieffenbachia paludicola Culcasia yangambiensis Colocasia affinis Cryptocoryne mekongensis Dieffenbachia panamensis Colocasia antiquorum Cryptocoryne minima Cyrtosperma beccarianum Dieffenbachia parlatorei Colocasia esculenta Cryptocoryne moehlmannii Cyrtosperma bougainvillense Dieffenbachia parvifolia Colocasia fallax Cryptocoryne nevillii Cyrtosperma brassii Dieffenbachia pittieri Colocasia gigantea Cryptocoryne noritoi Cyrtosperma carrii Dieffenbachia seguine Colocasia mannii Cryptocoryne nurii Cyrtosperma cuspidispathum Dieffenbachia shuttleworthiana Cryptocoryne pallidinervia Colocasia menglaensis Dieffenbachia standleyi Cyrtosperma giganteum Colocasia oresbia Cryptocoryne parva Cyrtosperma gressittiorum Dieffenbachia tonduzii Croatiella integrifolia Cryptocoryne pontederiifolia Cyrtosperma hambalii Dieffenbachia weberbaueri Cryptocoryne affinis Cryptocoryne pygmaea Cyrtosperma johnstonii Dieffenbachia weirii Cryptocoryne alba Cryptocoryne retrospiralis Cyrtosperma kokodense Dieffenbachia wendlandii Dieffenbachia williamsii Cryptocoryne albida Cryptocoryne schulzei Cyrtosperma macrotum Cryptocoryne annamica Cryptocoryne scurrilis Cyrtosperma merkusii Dieffenbachia wurdackii Cryptocoryne sivadasanii Dieffenbachia aglaonematifolia Dracontioides desciscens Cryptocoryne aponogetifolia Cryptocoryne spiralis Cryptocoryne auriculata Dieffenbachia antioquensis Dracontioides salvianii var. spiralis Cryptocoryne bangkaensis Dieffenbachia aurantiaca Dracontium amazonense Cryptocoryne spiralis Cryptocoryne beckettii Dieffenbachia beachiana Dracontium angustispathum var. cognatoides Cryptocoryne bogneri Dieffenbachia bechiana Dracontium asperispathum Cryptocoryne striolata Cryptocoryne tambraparaniana Dieffenbachia bowmannii Cryptocoryne bullosa Dracontium asperum Cryptocoryne ciliata Cryptocoryne thwaitesii Dieffenbachia brittonii Dracontium bogneri Cryptocoryne cognata Dieffenbachia burgeri Dracontium croatii Cryptocoryne uenoi Cryptocoryne consobrina Dieffenbachia cannifolia Dracontium dubium Cryptocoryne undulata Cryptocoryne cordata Dieffenbachia concinna Dracontium gigas Cryptocoryne usteriana var. cordata Cryptocoryne versteegii Dieffenbachia copensis Dracontium grandispathum Cryptocoryne cordata var. diderici Cryptocoryne vietnamensis Dieffenbachia cordata Dracontium grayumianum Cryptocoryne cordata Dieffenbachia costata Dracontium guianense Cryptocoryne villosa var. evae Dieffenbachia crebripistillata Dracontium iquitense Cryptocoryne walkeri Cryptocoryne cordata Dieffenbachia daguensis var. grabowskii Cryptocoryne wendtii Dracontium longipes Cryptocoryne cordata Cryptocoryne yujii Dieffenbachia davidsei Dracontium margaretae var. zonata Dieffenbachia duidae Dracontium nivosum Cryptocoryne zaidiana Cryptocoryne coronata Dieffenbachia elegans Dracontium peruvianum Cryptocoryne zukalii Cryptocoryne crispatula Culcasia angolensis Dieffenbachia enderi Dracontium pittieri var. crispatula Cryptocoryne crispatula Dieffenbachia fortunensis Culcasia annetii Dracontium plowmanii var. balansae Dieffenbachia fosteri Dracontium polyphyllum Culcasia bosii Cryptocoryne crispatula Dieffenbachia fournieri Culcasia brevipetiolata Dracontium prancei var. decus-mekongensis Cryptocoryne crispatula Dieffenbachia galdamesiae Dracontium purdieanum Culcasia caudata var. flaccidifolia Dieffenbachia gracilis Dracontium soconuscum Culcasia dinklagei Cryptocoryne crispatula Dieffenbachia grayumiana Dracontium spruceanum Culcasia ekongoloi var. planifolia Culcasia falcifolia Dieffenbachia hammelii Dracontium ulei Cryptocoryne crispatula var. tonkinensis Dieffenbachia herthae Dracunculus canariensis Culcasia glandulosa Cryptocoryne crispatula Culcasia insulana Dieffenbachia horichii Dracunculus vulgaris var. yunnanensis Dieffenbachia humilis Eminium albertii Culcasia lanceolata Cryptocoryne cruddasiana Dieffenbachia imperialis Eminium heterophyllum Culcasia liberica Cryptocoryne decus-silvae Dieffenbachia isthmia Eminium intortum Culcasia linearifolia Cryptocoryne dewitii Dieffenbachia killipii Eminium jaegeri Culcasia loukandensis Cryptocoryne edithiae Dieffenbachia lancifolia Eminium koenenianum Culcasia mannii Cryptocoryne elliptica Eminium lehmannii Dieffenbachia leopoldii Culcasia obliquifolia Cryptocoryne ferruginea Eminium rauwolffii Dieffenbachia longispatha Culcasia orientalis Cryptocoryne fusca var. rauwolffii Dieffenbachia lutheri Culcasia panduriformis Eminium rauwolffii Cryptocoryne griffithii Dieffenbachia macrophylla Culcasia parviflora var. kotschyi Cryptocoryne hudoroi Culcasia rotundifolia Dieffenbachia meleagris Eminium regelii Cryptocoryne ideii Culcasia sanagensis Dieffenbachia nitidipetiolata Eminium spiculatum Cryptocoryne jacobsenii Dieffenbachia obliqua Culcasia scandens Epipremnum amplissimum

Scientific name Epipremnum aureum Epipremnum carolinense Epipremnum ceramense Epipremnum dahlii Epipremnum falcifolium Epipremnum giganteum Epipremnum meeboldii Epipremnum moluccanum Epipremnum moszkowskii Epipremnum nobile Epipremnum obtusum Epipremnum papuanum Epipremnum pinnatum Epipremnum silvaticum Filarum manserichense Furtadoa mixta Furtadoa sumatrensis Gearum brasiliense Gonatopus angustus Gonatopus boivinii Gonatopus clavatus Gonatopus marattioides Gonatopus petiolulatus Gorgonidium beckianum Gorgonidium bulbostylum Gorgonidium cardenasianum Gorgonidium intermedium Gorgonidium mirabile Gorgonidium striatum Gorgonidium vargasii Gorgonidium vermicidum Gymnostachys anceps Hapaline appendiculata Hapaline benthamiana Hapaline brownii Hapaline celatrix Hapaline colaniae Hapaline ellipticifolia Hapaline kerrii Hapaline locii Helicodiceros muscivorus Hestia longifolia Heteropsis boliviana Heteropsis croatii Heteropsis duckeana Heteropsis ecuadorensis Heteropsis flexuosa var. flexuosa Heteropsis flexuosa var. maguirei Heteropsis linearis Heteropsis longispathacea Heteropsis macrophylla Heteropsis melinonii Heteropsis oblongifolia Heteropsis peruviana

Heteropsis rigidifolia

Heteropsis robusta

Scientific name Heteropsis salicifolia Heteropsis spruceana Heteropsis steyermarkii Heteropsis tenuispadix Holochlamys beccarii Homalomena adiensis Homalomena aeneifolia Homalomena agens Homalomena ardua Homalomena argentea Homalomena aromatica Homalomena asmae Homalomena asperifolia Homalomena atroviridis Homalomena atrox Homalomena batoeensis Homalomena bellula Homalomena burkilliana Homalomena clandestina Homalomena cochinchinensis Homalomena confusa Homalomena consobrina Homalomena cordata Homalomena corneri Homalomena crinipes Homalomena cristata Homalomena curtisii Homalomena curvata Homalomena davidiana Homalomena debilicrista Homalomena distans Homalomena doctersii Homalomena elegans Homalomena elegantula Homalomena erythropus subsp. erythropus Homalomena erythropus subsp. allenii Homalomena expedita Homalomena gadutensis Homalomena gaudichaudii Homalomena giamensis Homalomena gillii Homalomena griffithii Homalomena hainanensis Homalomena hammelii Homalomena hanneae Homalomena hastata Homalomena havilandii Homalomena hendersonii Homalomena hooglandii Homalomena humilis Homalomena impudica Homalomena insignis Homalomena jacobsiana Homalomena josefii Homalomena kalkmanii

Scientific name Homalomena kiahii Homalomena korthalsii Homalomena kualakohensis Homalomena kvistii Homalomena lancea Homalomena lancifolia Homalomena latifrons Homalomena lauterbachii Homalomena lindenii Homalomena longipes Homalomena magna Homalomena major Homalomena matangae Homalomena megalophylla Homalomena melanesica Homalomena metallica Homalomena minor Homalomena minutissima Homalomena moffleriana Homalomena monandra Homalomena montana Homalomena nigrescens Homalomena nutans Homalomena obovata Homalomena obscurifolia Homalomena occulta Homalomena ovalifolia Homalomena ovata Homalomena padangensis Homalomena palawanensis Homalomena peekelii Homalomena peltata Homalomena pendula Homalomena philippinensis Homalomena picturata Homalomena pineodora Homalomena pontederifolia Homalomena producta Homalomena pseudogeniculata Homalomena pulleana Homalomena punctulata Homalomena pyrospatha Homalomena robusta Homalomena roezelii Homalomena rostrata Homalomena rubescens Homalomena rusdii Homalomena sarawakensis Homalomena saxorum Homalomena schlechteri Homalomena scortechinii Homalomena sengkenyang Homalomena silvatica Homalomena singaporensis Homalomena soniae Homalomena speariae

Scientific name Homalomena steenisiana Homalomena stollei Homalomena striatieopetiolata Homalomena subcordata Homalomena symplocarpifolia Homalomena tenuispadix Homalomena terajaensis Homalomena treubii Homalomena truncata Homalomena vagans Homalomena vietnamensis Homalomena vittifolia Homalomena vivens Homalomena wallichii Homalomena wallisii Homalomena wendlandii Homalomena wongii Homalomena zollingeri Incarum pavonii Jasarum steyermarkii Lagenandra bogneri Lagenandra dewitii Lagenandra erosa Lagenandra gomezii Lagenandra jacobsenii Lagenandra keralensis Lagenandra koenigii Lagenandra lancifolia Lagenandra meeboldii Lagenandra nairii Lagenandra ovata Lagenandra praetermissa Lagenandra thwaitesii Lagenandra toxicaria Lagenandra undulata Lasia concinna Lasia spinosa Lasimorpha senegalensis Lemna aequinoctialis Lemna disperma Lemna gibba Lemna japonica I emna minor I emna minuta Lemna obscura Lemna perpusilla Lemna tenera Lemna trisulca Lemna turionifera Lemna valdiviana Lemna yungensis Lorenzia umbrosa Lysichiton americanus Lysichiton camtschatcensis Mangonia tweedieana Mangonia uruguaya Monstera acacoyaguensis

Homalomena kelungensis

Scientific name

Monstera acuminata Monstera adansonii var. adansonii Monstera adansonii var. klotzschiana Monstera adansonii var. laniata Monstera amargalensis Monstera aureopinnata Monstera barrieri Monstera busevi Monstera cenepensis Monstera costaricensis Monstera deliciosa Monstera dissecta Monstera dubia Monstera epipremnoides Monstera filamentosa Monstera florescanoana Monstera glaucescens Monstera gracilis Monstera kessleri Monstera lechleriana Monstera lentii Monstera luteynii Monstera maderaverde Monstera membranacea Monstera minima Monstera molinae Monstera obliqua Monstera oreophila Monstera pinnatipartita Monstera pittieri Monstera planadensis Monstera praetermissa Monstera punctulata Monstera siltepecana Monstera spruceana Monstera standleyana Monstera subpinnata Monstera tenuis Monstera tuberculata var. tuberculata Monstera tuberculata var. brevinoda Monstera vasquezii Monstera xanthospatha Montrichardia arborescens Montrichardia linifera Nephthytis afzelii var. afzelii Nephthytis afzelii var. graboensis Nephthytis bintuluensis Nephthytis hallaei Nephthytis mayombensis Nephthytis poissonii var. poissonii Nephthytis poissonii

var. constricta

Nephthytis swainei

Scientific name

Ooia grabowskii Ooia kinabaluensis Orontium aquaticum Pedicellarum paiei Peltandra sagittifolia Peltandra virginica Philodendron acreanum Philodendron acuminatissimum Philodendron acutifolium Philodendron adamantinum Philodendron adhatodifolium Philodendron advena Philodendron aemulum Philodendron alatum Philodendron albisuccus Philodendron alliodorum Philodendron alternans Philodendron alticola Philodendron altomacaense Philodendron amargalense Philodendron ampamii Philodendron amplisinum Philodendron ampullaceum Philodendron anaadu Philodendron ancuashii Philodendron angustialatum Philodendron angustilobum Philodendron angustisectum Philodendron anisotomum Philodendron annulatum Philodendron antonioanum Philodendron appendiculatum Philodendron applanatum Philodendron appunii Philodendron aristeguietae Philodendron aromaticum Philodendron asplundii Philodendron atabapoense Philodendron aurantiifolium subsp. aurantiifolium Philodendron aurantiifolium subsp. calderense Philodendron aurantispadix Philodendron aureimarginatum Philodendron clarkei Philodendron auriculatum Philodendron auritum Philodendron auyantepuiense Philodendron avenium Philodendron azulitense Philodendron bahiense Philodendron bakeri Philodendron balaoanum Philodendron barbourii Philodendron barrosoanum

Scientific name

Philodendron beniteziae Philodendron bernardopazii Philodendron billietiae Philodendron bipennifolium Philodendron bipinnatifidum Philodendron biribiriense Philodendron blanchetianum Philodendron bogotense Philodendron borgesii Philodendron brandii Philodendron brandtianum Philodendron brasiliense Philodendron breedlovei Philodendron brenesii Philodendron brent-berlinii Philodendron brevispathum Philodendron brewsterense Philodendron brunneicaule Philodendron buchtienii Philodendron buntingianum Philodendron burgeri Philodendron burle-marxii Philodendron calatheifolium Philodendron callosum subsp. callosum Philodendron callosum subsp. ptarianum Philodendron campii Philodendron camposportoanum Philodendron canaimae Philodendron canicaule Philodendron cardonii Philodendron cardosoi Philodendron carinatum Philodendron cataniapoense Philodendron caudatum Philodendron chimantae Philodendron chimboanum Philodendron chiriquense Philodendron chirripoense Philodendron chrysocarpum Philodendron cipoense Philodendron clewellii Philodendron colombianum Philodendron coloradense Philodendron condorcanquense Philodendron conforme Philodendron consanguineum Philodendron consobrinum Philodendron copense Philodendron corcovadense Philodendron cordatum Philodendron coriaceum Philodendron correae

Scientific name Philodendron cotobrusense Philodendron cotonense Philodendron craspedodromum Philodendron crassinervium Philodendron crassispathum Philodendron crassum Philodendron cremersii Philodendron cretosum Philodendron croatii Philodendron cruentospathum Philodendron cruentum Philodendron cuneatum Philodendron curvilobum Philodendron daniellii Philodendron danteanum Philodendron dardanianum Philodendron davidsei Philodendron davidsonii subsp. davidsonii Philodendron davidsonii subsp. bocatoranum Philodendron deflexum Philodendron delascioi Philodendron delinksii Philodendron deltoideum Philodendron densivenium Philodendron devansayanum Philodendron devianum Philodendron dioscoreoides Philodendron discretivenium Philodendron distantilohum Philodendron divaricatum Philodendron dodsonii Philodendron dolichophyllum Philodendron dominicalense Philodendron dressleri Philodendron dryanderae Philodendron duckei Philodendron chinchamayense Philodendron dunstervilleorum Philodendron dussii Philodendron dwyeri Philodendron dyscarpium var. dyscarpium Philodendron dyscarpium var. ventuarianum Philodendron eburneum Philodendron ecordatum Philodendron edenudatum Philodendron edmundoi Philodendron effusilobum Philodendron elaphoglossoides Philodendron elegans Philodendron elegantulum Philodendron englerianum subsp. englerianum Philodendron englerianum subsp. duidae Philodendron ensifolium

subsp. ensifolium

Philodendron cotapatense

Philodendron basii

Philodendron basivaginatum

Philodendron baudoense

Scientific name Philodendron ensifolium subsp. campanense Philodendron ensifolium subsp. colonense Philodendron ernestii Philodendron erubescens Philodendron escuintlense Philodendron exile Philodendron eximium Philodendron fendleri Philodendron ferrugineum Philodendron fibrillosum Philodendron fibrosum Philodendron findens Philodendron flumineum Philodendron folli Philodendron folsomii Philodendron fortunense Philodendron fragile Philodendron fragrantissimum Philodendron fraternum Philodendron furcatum Philodendron giganteum Philodendron gigas Philodendron glanduliferum subsp. glanduliferum subsp. camiloanum

Philodendron glanduliferum Philodendron glaziovii Philodendron gloriosum Philodendron goeldii Philodendron gonzalezii Philodendron grandifolium Philodendron grandipes Philodendron granulare Philodendron graveolens

Philodendron grayumii Philodendron grazielae Philodendron grenandii Philodendron guaiquinimae Philodendron gualeanum Philodendron guianense

Philodendron guttiferum Philodendron hammelii Philodendron hastatum

Philodendron hebetatum Philodendron hederaceum var. hederaceum

var. ovatum

var. linnaei

Philodendron lindenianum

Philodendron linguifolium

Philodendron lingulatum

Philodendron lindenii

Philodendron linnaei

Philodendron linnaei

Philodendron llanense

Philodendron loefgrenii

Philodendron longilaminatum

Philodendron longilobatum

var. rionegrense

Philodendron hatschbachii

Philodendron hederaceum var. kirkbridei Philodendron hederaceum var. oxycardium Philodendron heleniae subsp. heleniae

Philodendron heleniae subsp. amazonense Philodendron henry-pittieri

Philodendron herbaceum Philodendron herthae

Philodendron heterocraspedon

Scientific name Philodendron heterophyllum Philodendron heteropleurum Philodendron holstii Philodendron hooveri Philodendron hopkinsianum Philodendron houlletianum Philodendron huanucense Philodendron huashikatii Philodendron huaynacapacense Philodendron humile Philodendron hylaeae Philodendron ichthyoderma Philodendron immixtum Philodendron inaequilaterum Philodendron inconcinnum Philodendron inops Philodendron insigne Philodendron jacquinii Philodendron jefense Philodendron jodavisianum Philodendron jonkerorum Philodendron juninense Philodendron kautskyi Philodendron killipii Philodendron knappiae Philodendron krauseanum Philodendron kroemeri Philodendron krugii Philodendron lacerum Philodendron laticiferum Philodendron latifolium Philodendron lazorii Philodendron leal-costae Philodendron lechlerianum Philodendron lehmannii Philodendron lemae Philodendron lentii Philodendron leucanthum Philodendron leyvae Philodendron liesneri Philodendron ligulatum var. ligulatum Philodendron ligulatum var. heraclioanum Philodendron ligulatum

Scientific name Philodendron longipedunculatum Philodendron longipes Philodendron longirrhizum Philodendron longistilum Philodendron lundii Philodendron lupinum Philodendron macroglossum Philodendron macropodum Philodendron maculatum Philodendron madronense Philodendron maguirei Philodendron malesevichiae Philodendron mamei Philodendron marahuacae Philodendron maroae Philodendron martianum Philodendron martini Philodendron mathewsii Philodendron mawarinumae Philodendron maximum Philodendron mayoi Philodendron mcphersonii Philodendron megalophyllum Philodendron melanochrysum Philodendron melinonii Philodendron mellobarretoanum Philodendron membranaceum Philodendron merenbergense Philodendron meridense Philodendron mesae Philodendron mexicanum Philodendron micranthum Philodendron microstictum Philodendron millerianum Philodendron minarum Philodendron misahualliense Philodendron missionum Philodendron modestum Philodendron monsalveae Philodendron montanum Philodendron moonenii

Philodendron multispadiceum Philodendron muricatum Philodendron musifolium Philodendron myrmecophilum Philodendron nadruzianum Philodendron nanegalense Philodendron narinoense Philodendron nebulense Philodendron ninoanum

Philodendron morii

Philodendron multinervum

Philodendron nullinervium Philodendron oblanceolatum Philodendron obliquifolium

Philodendron niqueanum

Scientific name

Philodendron oblongum Philodendron obtusilobum Philodendron ochrostemon Philodendron oligospermum Philodendron opacum Philodendron orionis Philodendron ornatum Philodendron pachycaule Philodendron pachyphyllum Philodendron palaciosii Philodendron paludicola Philodendron panamense Philodendron panduriforme var. panduriforme Philodendron panduriforme var. reichenbachianum Philodendron parvilobum Philodendron pastazanum Philodendron patriciae Philodendron paucinervium Philodendron paxianum Philodendron pedatum Philodendron pedunculum Philodendron peraiense Philodendron perplexum

Philodendron peperomioides Philodendron phlebodes var. phlebodes Philodendron phlebodes var. kermesinum Philodendron pimichinese Philodendron pinnatifidum Philodendron pinnatilobum

Philodendron pipolyi Philodendron pirrense Philodendron placidum Philodendron planadense Philodendron platypetiolatum Philodendron platypodum Philodendron pogonocaule Philodendron polliciforme Philodendron popenoei Philodendron populneum Philodendron prominulinervium

Philodendron propinguum Philodendron pseudauriculatum Philodendron pseudoundulatum Philodendron pteropus Philodendron pterotum Philodendron puhuangii Philodendron pulchellum Philodendron pulchrum Philodendron purpureoviride Philodendron purulhense Philodendron pusillum Philodendron quinquelobum Philodendron quinquenervium

Philodendron quitense

Scientific name Philodendron radiatum var. radiatum Philodendron radiatum var. pseudoradiatum Philodendron rayanum Philodendron recurvifolium Philodendron remifolium subsp. remifolium Philodendron remifolium subsp. sabulosum Philodendron renauxii Philodendron reticulatum Philodendron rhizomatosum Philodendron rhodoaxis subsp. rhodoaxis Philodendron rhodoaxis subsp. lewisii Philodendron Philodendron rhodospathiphyllum Philodendron ricardoi Philodendron rigidifolium subsp. rigidifolium Philodendron rigidifolium subsp. sanctae-ritae Philodendron rimachii Philodendron riparium Philodendron robustum Philodendron rodrigueziae Philodendron roezlii Philodendron rojasianum Philodendron romeroi Philodendron roraimae var. roraimae Philodendron roraimae subsp. aracamuniense Philodendron roseocataphyllum Philodendron roseopetiolatum Philodendron roseospathum var. roseospathum Philodendron roseospathum var. angustilaminatum Philodendron rothschuhianum Philodendron rubrocinctum Philodendron rubromaculatum Philodendron rudgeanum Philodendron rugosum Philodendron ruizii Philodendron ruthianum Philodendron sagittifolium Philodendron samayense Philodendron santodominguense Philodendron saxicola Philodendron scalarinerve Philodendron scherberichii Philodendron schottianum Philodendron schottii subsp. schottii Philodendron schottii subsp. talamancae Philodendron scitulum Philodendron scottmorianum

Philodendron seguine

subsp. seguine

Scientific name Philodendron seguine subsp. lingua-bovis Philodendron senatocarpium Philodendron serpens Philodendron silverstonei Philodendron simmondsii Philodendron simonianum Philodendron simsii Philodendron simulans Philodendron smithii Philodendron solimoesense Philodendron sonderianum Philodendron sousae Philodendron sparreorum Philodendron speciosum Philodendron sphalerum Philodendron spiritus-sancti Philodendron splitgerberi Philodendron spruceanum Philodendron squamicaule Philodendron squamiferum Philodendron squamipetiolatum Philodendron standleyi Philodendron stenolobum Philodendron stenophyllum Philodendron stevermarkii Philodendron straminicaule Philodendron striatum Philodendron strictum Philodendron suberosum Philodendron subhastatum Philodendron subincisum Philodendron sucrense Philodendron sulcatum Philodendron sulcicaule Philodendron surinamense Philodendron swartiae Philodendron tachirense Philodendron tarmense Philodendron tatei subsp. tatei Philodendron tatei subsp. melanochlorum Philodendron tenue Philodendron tenuipes Philodendron tenuispadix Philodendron teretipes Philodendron thalassicum Philodendron thaliifolium Philodendron tortum Philodendron toshibae Philodendron traunii Philodendron triangulare Philodendron tricostatum Philodendron tripartitum Philodendron triplum Philodendron trojitense

Scientific name Philodendron tuerckheimii Philodendron tweedieanum Philodendron tysonii Philodendron ubigantupense Philodendron uleanum Philodendron uliginosum Philodendron undulatum Philodendron urraoense Philodendron ushanum Philodendron utleyanum Philodendron validinervium Philodendron vargealtense Philodendron variifolium Philodendron venezuelense Philodendron venosum Philodendron ventricosum Philodendron venulosum Philodendron venustifoliatum Philodendron venustum Philodendron verapazense Philodendron verrucapetiolum Philodendron verrucosum Philodendron victoriae Philodendron vinaceum Philodendron viride Philodendron wadedavisii Philodendron wallisii Philodendron warszewiczii Philodendron weberbaueri Philodendron wendlandii Philodendron werkhoveniae Philodendron wilburii var. wilburii Philodendron wilburii var. longipedunculatum Philodendron williamsii Philodendron wittianum Philodendron woronowii Philodendron wullschlaegelii Philodendron wurdackii Philodendron xanadu Philodendron yavitense Philodendron yutajense Philodendron zhuanum Philonotion americanum Philonotion bolivaranum Philonotion spruceanum Phymatarum borneense Pichinia disticha Pinellia cordata Pinellia fujianensis Pinellia integrifolia Pinellia pedatisecta Pinellia peltata Pinellia polyphylla Pinellia ternata Pinellia tripartita Pinellia yaoluopingensis

Scientific name Piptospatha burbidgei Piptospatha elongata Piptospatha impolita Piptospatha insignis Piptospatha manduensis Piptospatha marginata Piptospatha perakensis Piptospatha remiformis Piptospatha repens Piptospatha ridleyi Piptospatha truncata Piptospatha viridistigma Pistia stratiotes Podolasia stipitata Pothoidium lobbianum Pothos armatus Pothos atropurpurascens Pothos barberianus Pothos beccarianus Pothos brassii Pothos brevistylus Pothos brevivaginatus Pothos chinensis Pothos clavatus Pothos crassipedunculatus Pothos curtisii Pothos cuspidatus Pothos cylindricus Pothos dolichophyllus Pothos dzui Pothos englerianus Pothos falcifolius Pothos gigantipes Pothos gracillimus Pothos grandis Pothos hellwigii Pothos hookeri Pothos inaequilaterus Pothos insignis Pothos junghuhnii Pothos keralensis Pothos kerrii Pothos kingii Pothos lancifolius Pothos laurifolius Pothos leptostachyus Pothos longipes Pothos longivaginatus Pothos luzonensis Pothos macrocephalus Pothos mirabilis Pothos motleyanus Pothos oliganthus Pothos ovatifolius Pothos oxyphyllus Pothos papuanus Pothos parvispadix

Philodendron trujilloi

Scientific name

Pothos philippinensis Pothos pilulifer Pothos polystachyus Pothos remotiflorus Pothos repens Pothos roxburghii Pothos salicifolius Pothos scandens Pothos tener Pothos thomsonianus Pothos touranensis Pothos versteegii Pothos volans Pothos zippelii Protarum sechellarum Pseudohydrosme buettneri Pseudohydrosme gabunensis Pycnospatha arietina Pycnospatha palmata Remusatia hookeriana Remusatia pumila Remusatia vivipara Remusatia yunnanensis Rhaphidophora acuminata Rhaphidophora africana Rhaphidophora angustata Rhaphidophora araea Rhaphidophora australasica Rhaphidophora balgooyi Rhaphidophora banosensis Rhaphidophora beccarii Rhaphidophora bonii Rhaphidophora brevispathacea Rhaphidophora burkilliana Rhaphidophora calophylla Rhaphidophora chevalieri Rhaphidophora conica Rhaphidophora conocephala Rhaphidophora corneri Rhaphidophora crassicaulis Rhaphidophora crassifolia Rhaphidophora cravenschoddeana Rhaphidophora cretosa Rhaphidophora cryptantha Rhaphidophora cylindrosperma Rhaphidophora dahlii Rhaphidophora decursiva Rhaphidophora discolor Rhaphidophora dulongensis Rhaphidophora elliptica Rhaphidophora elliptifolia Rhaphidophora elmeri Rhaphidophora falcata Rhaphidophora floresensis Rhaphidophora foraminifera

Scientific name

Rhaphidophora formosana Rhaphidophora fortis Rhaphidophora geniculata Rhaphidophora glauca Rhaphidophora gorokensis Rhaphidophora guamensis Rhaphidophora hayi Rhaphidophora hongkongensis Rhaphidophora hookeri Rhaphidophora intonsa Rhaphidophora intrusa Rhaphidophora jubata Rhaphidophora kokodensis Rhaphidophora koordersii Rhaphidophora korthalsii Rhaphidophora laichauensis Rhaphidophora lancifolia Rhaphidophora latevaginata Rhaphidophora liukiuensis Rhaphidophora lobbii Rhaphidophora luchunensis Rhaphidophora maingayi Rhaphidophora megaphylla Rhaphidophora megasperma Rhaphidophora megastigma Rhaphidophora microspadix Rhaphidophora mima Rhaphidophora minor Rhaphidophora moluccensis Rhaphidophora montana Rhaphidophora monticola Rhaphidophora neoguineensis Rhaphidophora nicolsonii Rhaphidophora okapensis Rhaphidophora oligosperma Rhaphidophora ovoidea Rhaphidophora pachyphylla Rhaphidophora parvifolia Rhaphidophora peepla Rhaphidophora peeploides Rhaphidophora perkinsiae Rhaphidophora pertusa Rhaphidophora petrieana Rhaphidophora philippinensis Rhaphidophora pilosa Rhaphidophora puberula Rhaphidophora sabit Rhaphidophora sarasinorum Rhaphidophora schlechteri Rhaphidophora schottii Rhaphidophora spathacea Rhaphidophora spuria Rhaphidophora stenophylla Rhaphidophora stolleana

Scientific name

Rhaphidophora talamauana Rhaphidophora tenuis Rhaphidophora ternatensis Rhaphidophora tetrasperma Rhaphidophora teysmanniana Rhaphidophora todayensis Rhaphidophora tonkinensis Rhaphidophora typha Rhaphidophora ustulata Rhaphidophora versteegii Rhaphidophora waria Rhodospatha acosta-solisii Rhodospatha arborescens Rhodospatha badilloi Rhodospatha bolivarana Rhodospatha boliviensis Rhodospatha brachypoda Rhodospatha brent-berlinii Rhodospatha cardonae Rhodospatha densinervia Rhodospatha dissidens Rhodospatha falconensis Rhodospatha forgetii Rhodospatha guasareensis Rhodospatha herrerae Rhodospatha katipas Rhodospatha kraenzlinii Rhodospatha latifolia Rhodospatha monsalveae Rhodospatha moritziana Rhodospatha mukuntakia Rhodospatha oblongata Rhodospatha pellucida Rhodospatha perezii Rhodospatha piushaduka Rhodospatha robusta Rhodospatha statutii Rhodospatha stevermarkii Rhodospatha venosa Rhodospatha wendlandii Sauromatum brevipes Sauromatum brevipilosum Sauromatum diversifolium Sauromatum gaoligongense Sauromatum giganteum Sauromatum hirsutum Sauromatum horsfieldii Sauromatum tentaculatum Sauromatum venosum Scaphispatha gracilis Scaphispatha robusta Schismatoglottis acutifolia Schismatoglottis adoceta Schismatoglottis ahmadii Schismatoglottis ardenii

Scientific name Schismatoglottis bauensis Schismatoglottis bifasciata Schismatoglottis bogneri Schismatoglottis brevicuspis Schismatoglottis calyptrata Schismatoglottis canaliculata Schismatoglottis ciliata Schismatoglottis clarae Schismatoglottis clausula Schismatoglottis clemensiorum Schismatoglottis confinis Schismatoglottis conoidea Schismatoglottis convolvula Schismatoglottis corneri Schismatoglottis crinitissima Schismatoglottis cyria Schismatoglottis decipiens Schismatoglottis dilecta Schismatoglottis dulosa Schismatoglottis ecaudata Schismatoglottis edanoi Schismatoglottis elegans Schismatoglottis erecta Schismatoglottis eximia Schismatoglottis eymae Schismatoglottis ferruginea Schismatoglottis gamoandra Schismatoglottis gillianiae Schismatoglottis glauca Schismatoglottis grabowskii Schismatoglottis hainanensis Schismatoglottis harmandii Schismatoglottis hayana Schismatoglottis hayi Schismatoglottis hottae Schismatoglottis ifugaoensis Schismatoglottis inculta Schismatoglottis jelandii Schismatoglottis jepomii Schismatoglottis jitinae Schismatoglottis josefii Schismatoglottis kurzii Schismatoglottis lancifolia Schismatoglottis latevaginata Schismatoglottis linae Schismatoglottis lingua Schismatoglottis longispatha Schismatoglottis luzonensis Schismatoglottis maelii Schismatoglottis matangensis Schismatoglottis mayoana Schismatoglottis merrillii Schismatoglottis mindanaoana Schismatoglottis mira Schismatoglottis modesta Schismatoglottis monoplacenta

Schismatoglottis asperata

Schismatoglottis barbata

Rhaphidophora sulcata

Rhaphidophora sylvestris

Scientific name

Schismatoglottis moodii Schismatoglottis motleyana Schismatoglottis multiflora Schismatoglottis multinervia Schismatoglottis nervosa Schismatoglottis niahensis Schismatoglottis nicolsonii Schismatoglottis patentinervia Schismatoglottis pectinervia Schismatoglottis penangensis Schismatoglottis petri Schismatoglottis platystigma Schismatoglottis plurivenia Schismatoglottis puberulipes Schismatoglottis pudenda Schismatoglottis pumila Schismatoglottis pusilla Schismatoglottis pyrrhias Schismatoglottis retinervia Schismatoglottis roseospatha Schismatoglottis samarensis Schismatoglottis sarikeensis Schismatoglottis schottii Schismatoglottis scortechinii Schismatoglottis sejuncta Schismatoglottis silamensis Schismatoglottis simonii Schismatoglottis subundulata Schismatoglottis tahubangensis Schismatoglottis tecturata Schismatoglottis tessellata Schismatoglottis thelephora Schismatoglottis trifasciata Schismatoglottis trivittata Schismatoglottis trusmadiensis Schismatoglottis turbata Schismatoglottis ulusarikeiensis Schismatoglottis unifolia Schismatoglottis venusta Schismatoglottis viridissima Schismatoglottis wahaiana Schismatoglottis wallichii Schismatoglottis warburgiana Schismatoglottis wongii Schismatoglottis zonata Schottariella mirifica Scindapsus alpinus Scindapsus altissimus Scindapsus beccarii Scindapsus carolinensis Scindapsus coriaceus Scindapsus crassipes Scindapsus curranii Scindapsus cuscuaria Scindapsus cuscuarioides

Scindapsus falcifolius

Scientific name Scindapsus geniculatus Scindapsus glaucescens Scindapsus grandifolius Scindapsus hederaceus Scindapsus javanicus Scindapsus latifolius Scindapsus longipes Scindapsus longistipitatus Scindapsus lucens Scindapsus maclurei Scindapsus mamilliferus Scindapsus marantifolius Scindapsus officinalis Scindapsus perakensis Scindapsus pictus Scindapsus roseus Scindapsus rupestris Scindapsus salomoniensis Scindapsus schlechteri Scindapsus scortechinii Scindapsus splendidus Scindapsus subcordatus Scindapsus suffruticosus Scindapsus sumatranus Scindapsus treubii Spathantheum fallax Spathantheum orbignyanum Spathicarpa gardneri Spathicarpa hastifolia Spathicarpa lanceolata Spathiphyllum atrovirens Spathiphyllum barbourii Spathiphyllum bariense Spathiphyllum blandum Spathiphyllum brent-berlinii Spathiphyllum brevirostre Spathiphyllum buntingianum Spathiphyllum cannifolium Spathiphyllum cochlearispathum Spathiphyllum commutatum Spathiphyllum cuspidatum Spathiphyllum diazii Spathiphyllum dressleri Spathiphyllum floribundum Spathiphyllum friedrichsthalii Spathiphyllum fulvovirens Spathiphyllum gardneri Spathiphyllum gracile Spathiphyllum grandifolium Spathiphyllum grazielae Spathiphyllum humboldtii Spathiphyllum jejunum Spathiphyllum juninense

Spathiphyllum kalbreyeri

Spathiphyllum kochii

Spathiphyllum laeve

Scientific name

Spathiphyllum lanceifolium Spathiphyllum lechlerianum Spathiphyllum maguirei Spathiphyllum matudae Spathiphyllum mawarinumae Spathiphyllum minor Spathiphyllum monachinoi var. monachinoi Spathiphyllum monachinoi var. perangustum Spathiphyllum montanum Spathiphyllum neblinae Spathiphyllum ortgiesii Spathiphyllum patinii Spathiphyllum patulinervum Spathiphyllum perezii Spathiphyllum phryniifolium Spathiphyllum pygmaeum Spathiphyllum quindiuense Spathiphyllum schlechteri Spathiphyllum schomburgkii Spathiphyllum silvicola Spathiphyllum solomonense Spathiphyllum tenerum Spathiphyllum uspanapaensis Spathiphyllum wallisii Spathiphyllum wendlandii Spirodela oligorrhiza Spirodela polyrhiza Spirodela punctata Spirodela sichuanensis Stenospermation adsimile Stenospermation ammiticum Stenospermation amomifolium Stenospermation ancuashii Stenospermation andreanum Stenospermation angosturense Stenospermation angustifolium Stenospermation arborescens Stenospermation archeri Stenospermation benavidesae Stenospermation brachypodum Stenospermation crassifolium Stenospermation densiovulatum Stenospermation dictyoneurum Stenospermation ellipticum Stenospermation escobariae Stenospermation flavescens Stenospermation flavum Stenospermation gentryi Stenospermation glaucophyllum

Scientific name

Stenospermation latifolium Stenospermation longifolium Stenospermation *longipetiolatum* Stenospermation longispadix Stenospermation maguirei Stenospermation majus Stenospermation marantifolium Stenospermation mathewsii var. mathewsii Stenospermation mathewsii var. stipitatum Stenospermation monsalvae Stenospermation 5 multiovulatum Stenospermation nebulense Stenospermation olgae Stenospermation parvum Stenospermation peripense Stenospermation pittieri Stenospermation popayanense Stenospermation pteropus Stenospermation robustum Stenospermation rusbyi Stenospermation sessile Stenospermation spruceanum Stenospermation subellipticum Stenospermation ulei Stenospermation velutinum Stenospermation wallisii Stenospermation zeacarpium Steudnera assamica Steudnera capitellata Steudnera colocasiifolia Steudnera colocasioides Steudnera discolor Steudnera gagei Steudnera griffithii Steudnera henryana Steudnera kerrii Stylochaeton angolense Stylochaeton bogneri Stylochaeton borumense Stylochaeton crassispathum Stylochaeton cuculliferum Stylochaeton euryphyllum Stylochaeton grande Stylochaeton hypogeum Stylochaeton kornasii Stylochaeton lancifolium Stylochaeton malaissei Stylochaeton milneanum Stylochaeton natalense subsp. natalense Stylochaeton natalense subsp. maximum Stylochaeton natalense subsp. obliquinerve

Stylochaeton oligocarpum

Stenospermation gracile

Stenospermation hilligii

Stenospermation laevis

Stenospermation interruptum

Scientific name

Stylochaeton pilosum Stylochaeton puberulum Stylochaeton salaamicum Stylochaeton shabaense Stylochaeton tortispathum Stylochaeton zenkeri Symplocarpus egorovii Symplocarpus foetidus Symplocarpus nabekuraensis Symplocarpus nipponicus Symplocarpus renifolius Synandrospadix vermitoxicus Syngonium angustatum Syngonium armigerum Syngonium atrovirens Syngonium auritum Syngonium castroi Syngonium chiapense Syngonium chocoanum Syngonium crassifolium Syngonium dodsonianum Syngonium erythrophyllum Syngonium foreroanum Syngonium gentryanum Syngonium harlingianum Syngonium hastiferum Svngonium hastifolium Syngonium hoffmannii Syngonium laterinervium Syngonium Ilanoense Syngonium macrophyllum Syngonium mauroanum Syngonium meridense Syngonium neglectum Syngonium oduberi Syngonium podophyllum var. podophyllum Syngonium podophyllum var. peliocladum Syngonium rayi Syngonium sagittatum Syngonium salvadorense Syngonium schottianum Syngonium sparreorum Syngonium standleyanum Syngonium steyermarkii Syngonium triphyllum Syngonium wendlandii Syngonium yurimaguense Taccarum caudatum Taccarum crassispathum Taccarum peregrinum Taccarum ulei Taccarum warmingii Taccarum weddellianum Theriophonum dalzellii Theriophonum danielii

Theriophonum fischeri

Scientific name Theriophonum infaustum Theriophonum manickamii Theriophonum minutum Theriophonum sivaganganum Typhonium acetosella Typhonium adnatum Typhonium albidinervium Typhonium albispathum Typhonium alismifolium Typhonium angustilobum Typhonium bachmaense Typhonium baoshanense Typhonium blumei Typhonium bognerianum Typhonium brownii Typhonium bulbiferum Typhonium circinnatum Typhonium cochleare Typhonium conchiforme Typhonium cordifolium Typhonium digitatum Typhonium echinulatum Typhonium eliosurum Typhonium filiforme Typhonium flagelliforme Typhonium fultum Typhonium gagnepainii Typhonium gallowayi Typhonium glaucum Typhonium griseum Typhonium hayatae Typhonium huense Typhonium hunanense Typhonium inopinatum Typhonium jinpingense Typhonium johnsonianum Typhonium jonesii Typhonium laoticum Typhonium liliifolium Typhonium lineare Typhonium listeri Typhonium medusae Typhonium mirabile Typhonium neogracile Typhonium nudibaccatum Typhonium orbifolium Typhonium pedatisectum Typhonium pedunculatum Typhonium peltandroides Typhonium penicillatum Typhonium pottingeri Typhonium praecox Typhonium praetermissum Typhonium pusillum

Typhonium reflexum

Typhonium roxburghii

Typhonium russell-smithii

Scientific name

Typhonium sagittariifolium Typhonium saraburiensis Typhonium sinhabaedyae Typhonium smitinandii Typhonium stigmatilobatum Typhonium subglobosum Typhonium taylorii Typhonium trifoliatum Typhonium trilobatum Typhonium tubispathum Typhonium varians Typhonium vermiforme Typhonium violifolium Typhonium watanabei Typhonium weipanum Typhonium wilbertii Typhonodorum lindleyanum Ulearum donburnsii Ulearum sagittatum var. sagittatum Ulearum sagittatum var. viridispadix Urospatha angustiloba Urospatha antisylleptica Urospatha caudata Urospatha edwallii Urospatha friedrichsthalii Urospatha loefgreniana Urospatha meyeri Urospatha riedeliana Urospatha sagittifolia Urospatha somnolenta Urospatha wurdackii Wolffia angusta Wolffia arrhiza Wolffia australiana Wolffia borealis Wolffia brasiliensis Wolffia columbiana Wolffia cylindracea Wolffia elongata Wolffia globosa Wolffia microscopica Wolffia neglecta Wolffiella caudata Wolffiella denticulata Wolffiella gladiata Wolffiella hyalina Wolffiella lingulata Wolffiella neotropica Wolffiella oblonga Wolffiella repanda Wolffiella rotunda Wolffiella welwitschii Xanthosoma acutum Xanthosoma akkermansii Xanthosoma aristeguietae

Scientific name

Xanthosoma baguense Xanthosoma bayo Xanthosoma belophyllum Xanthosoma bilineatum Xanthosoma bolivaranum Xanthosoma brasiliense Xanthosoma brevispathaceum Xanthosoma caladioides Xanthosoma caracu Xanthosoma caulotuberculatum Xanthosoma conspurcatum Xanthosoma contractum Xanthosoma cordatum Xanthosoma cordifolium Xanthosoma cubense Xanthosoma daguense var. daguense Xanthosoma daguense var. amargalense Xanthosoma dealbatum Xanthosoma eggersii Xanthosoma exiguum Xanthosoma flavomaculatum Xanthosoma fractum Xanthosoma granvillei Xanthosoma guttatum Xanthosoma hebetatum Xanthosoma helleborifolium Xanthosoma herrerae Xanthosoma hylaeae Xanthosoma latestigmatum Xanthosoma longilobum Xanthosoma lucens Xanthosoma mafaffoides Xanthosoma mariae Xanthosoma maroae Xanthosoma maximiliani Xanthosoma mendozae Xanthosoma mexicanum Xanthosoma narinoense Xanthosoma nitidum Xanthosoma obtusilobum Xanthosoma orinocense Xanthosoma paradoxum Xanthosoma pariense Xanthosoma peltatum Xanthosoma pentaphyllum Xanthosoma platylobum Xanthosoma plowmanii Xanthosoma poeppigii Xanthosoma pottii Xanthosoma puberulum Xanthosoma pubescens Xanthosoma pulchrum Xanthosoma riedelianum Xanthosoma riparium Xanthosoma robustum

Xanthosoma auriculatum

Appendices

Scientific name

Xanthosoma sagittifolium Xanthosoma saguasense Xanthosoma seideliae Xanthosoma stenospathum Xanthosoma striatipes Xanthosoma striolatum Xanthosoma syngoniifolium Xanthosoma taioba Xanthosoma tarapotense

Scientific name

Xanthosoma trichophyllum Xanthosoma trilobum Xanthosoma ulei Xanthosoma undipes Xanthosoma viviparum Xanthosoma weeksii Xanthosoma wendlandii Xanthosoma yucatanense Zamioculcas zamiifolia

Scientific name

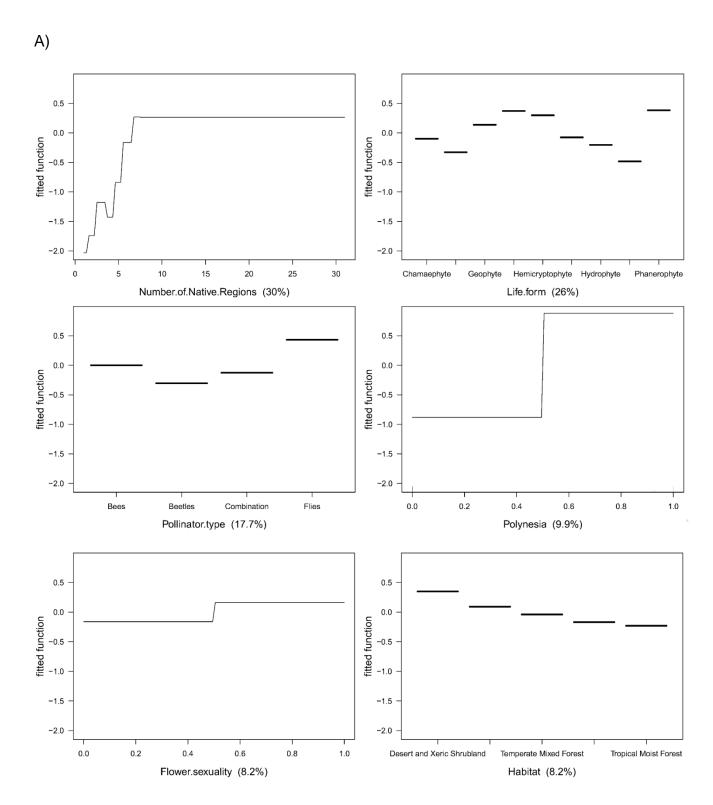
Zantedeschia aethiopica Zantedeschia albomaculata subsp. albomaculata Zantedeschia albomaculata subsp. macrocarpa Zantedeschia elliottiana Zantedeschia jucunda Zantedeschia odorata Zantedeschia pentlandii Zantedeschia rehmannii

Scientific name

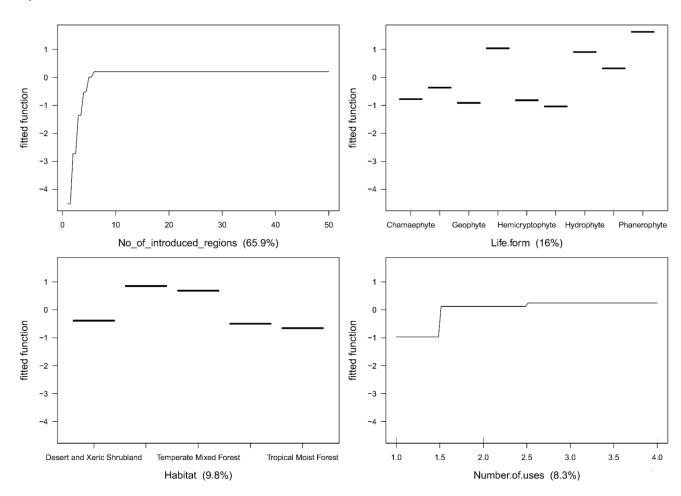
Zantedeschia valida Zomicarpa pythonium Zomicarpa steigeriana Zomicarpella amazonica

Zomicarpella maculata

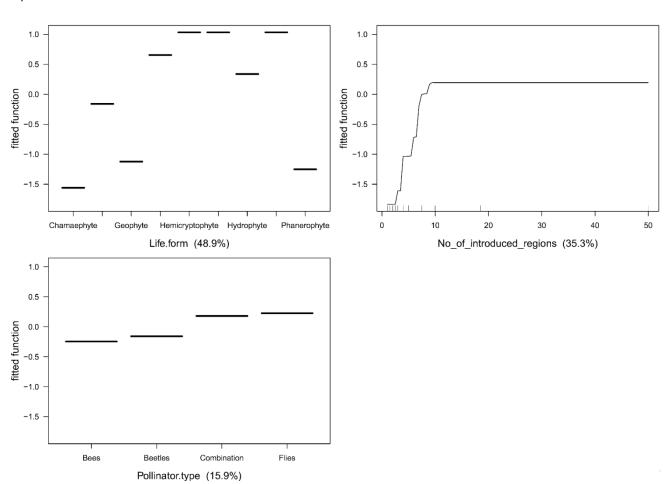
Appendix 3. Fitted function plots produced from the boosted regression tree models for species categorised in the (A) introduction, (B) naturalization and (C) invasion stages.



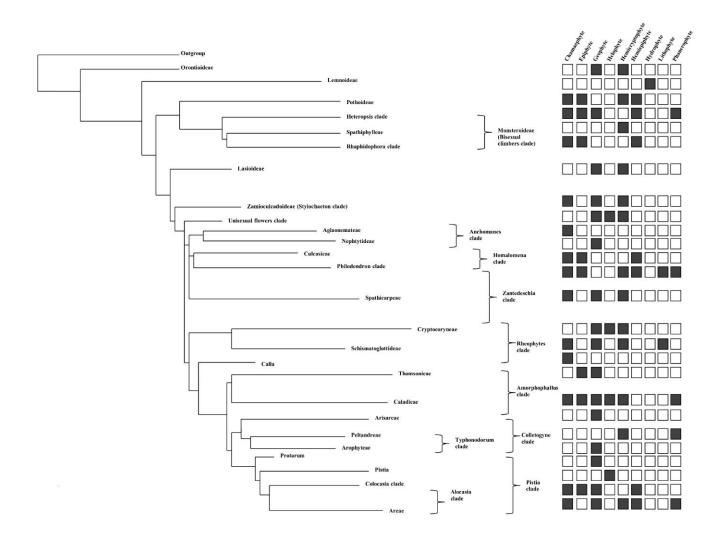






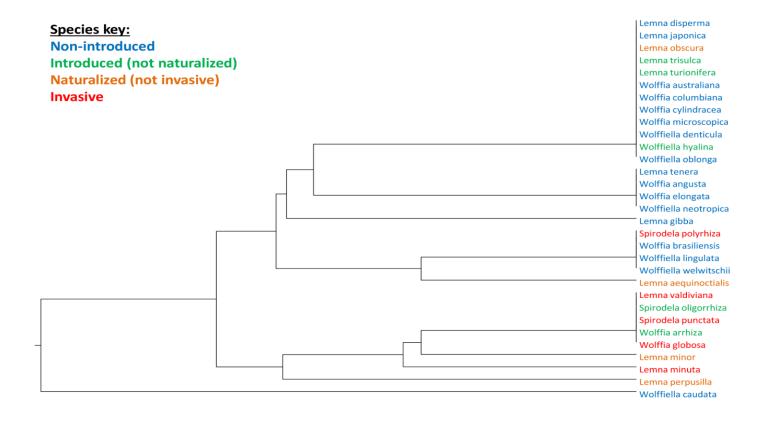


Appendix 4. Stick phylogeny of Araceae lineages. Black squares correspond to each clade and their associated life form(s). For further details on the phylogeny see (Cusimano *et al.*, 2011). The tree reveals that life-forms are spread across the phylogeny.

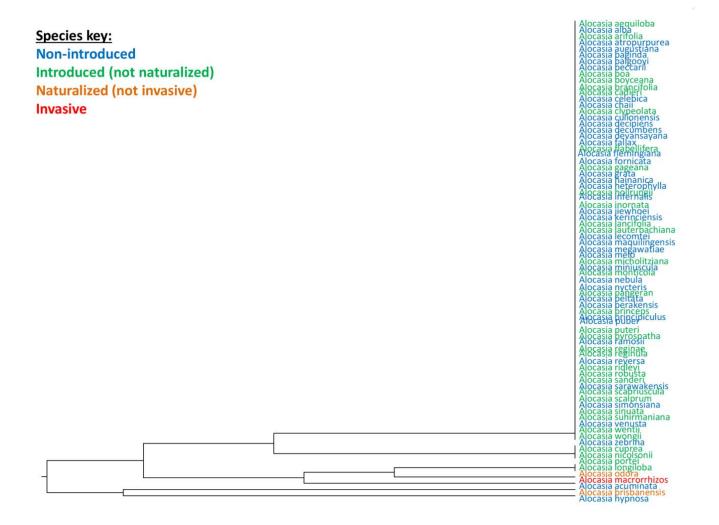


Appendix 5. Phenograms illustrating species that have a potential to become invasive based on shared traits within the following monophyletic groups: (A) Lemnoideae, (B) *Alocasia*, (C) *Amydrium*, (D) *Ariopsis*, (E) *Arum*, (F) *Caladium*, (G) *Cryptocoryne*, and (H) *Gymnostachys*.

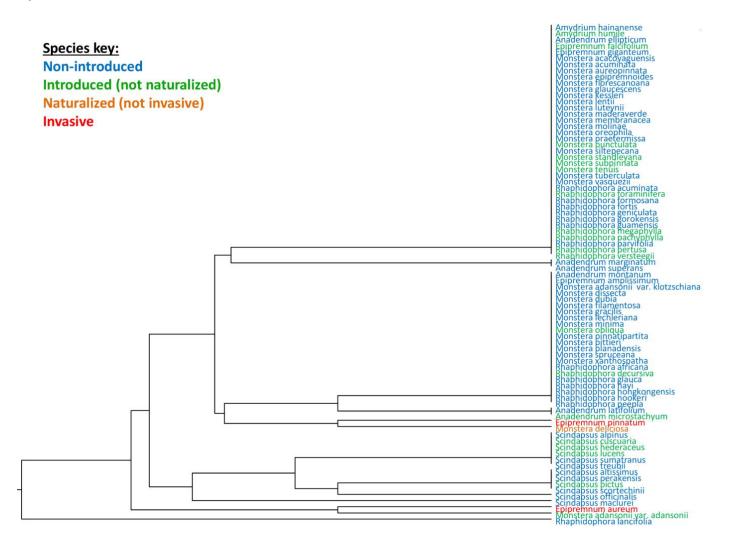
A)

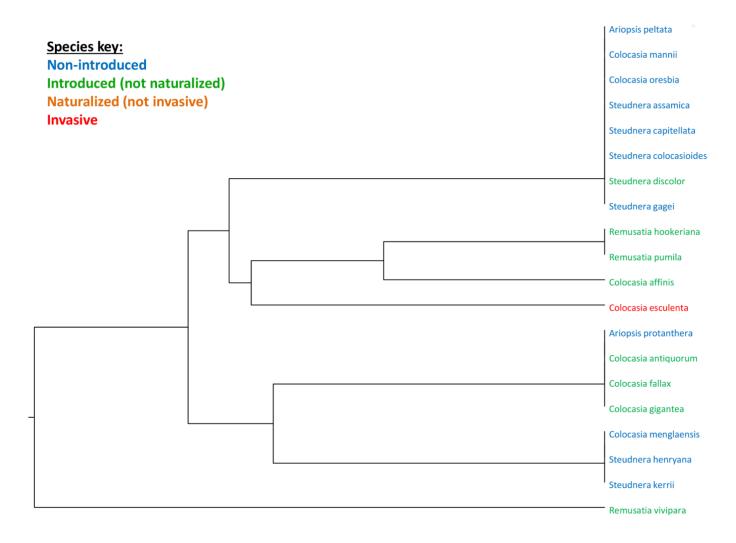


B)

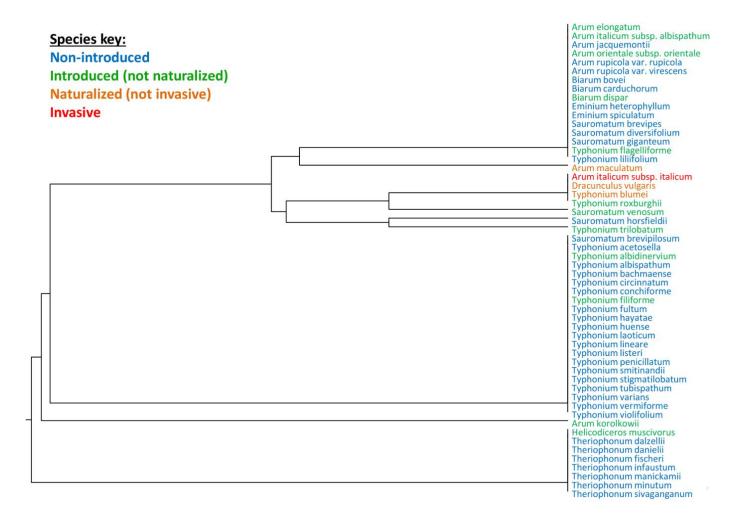


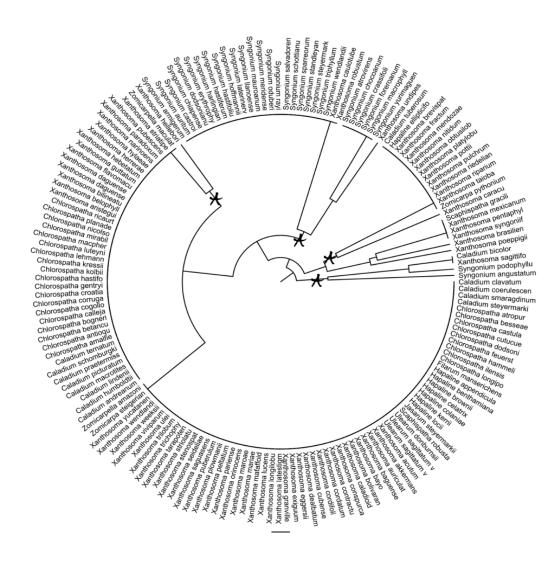
C)



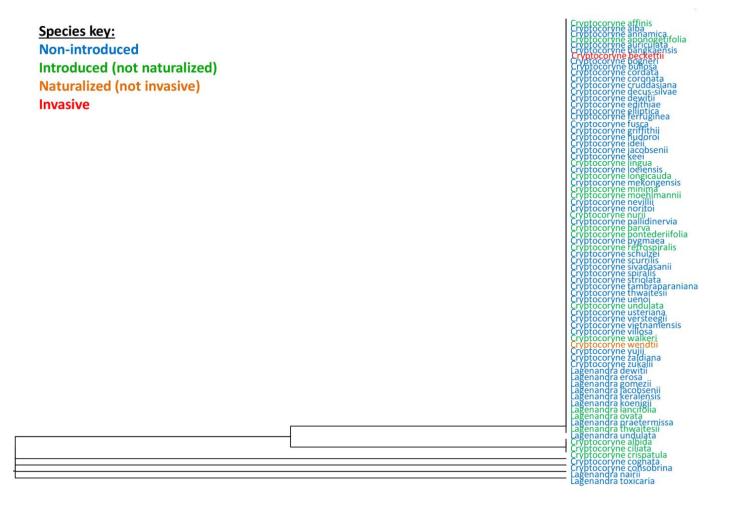


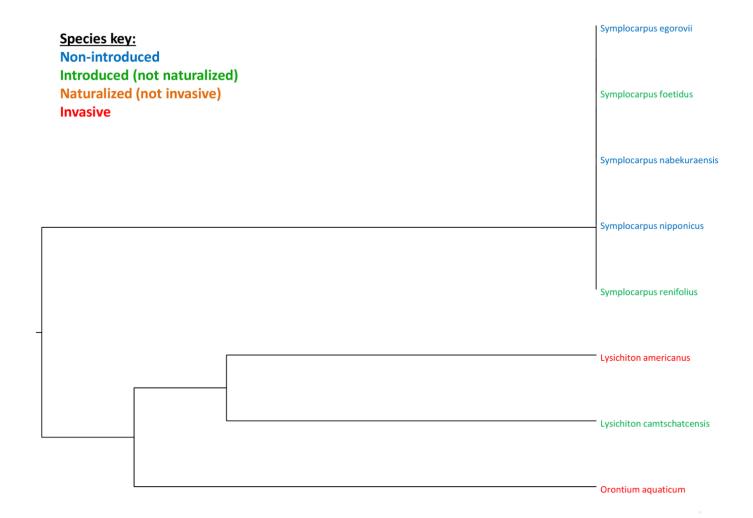
E)





G)



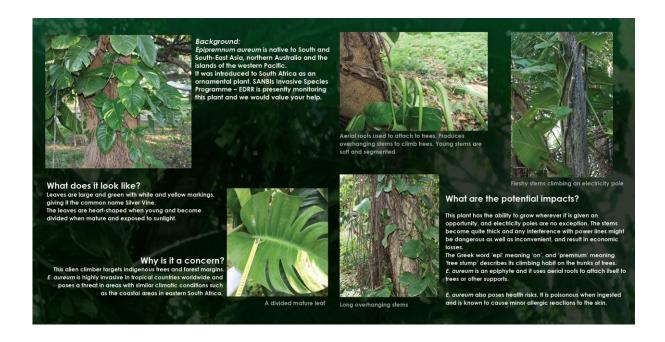


Appendix 6. Specimens of *Epipremnum aureum* in the KwaZulu-Natal herbarium.

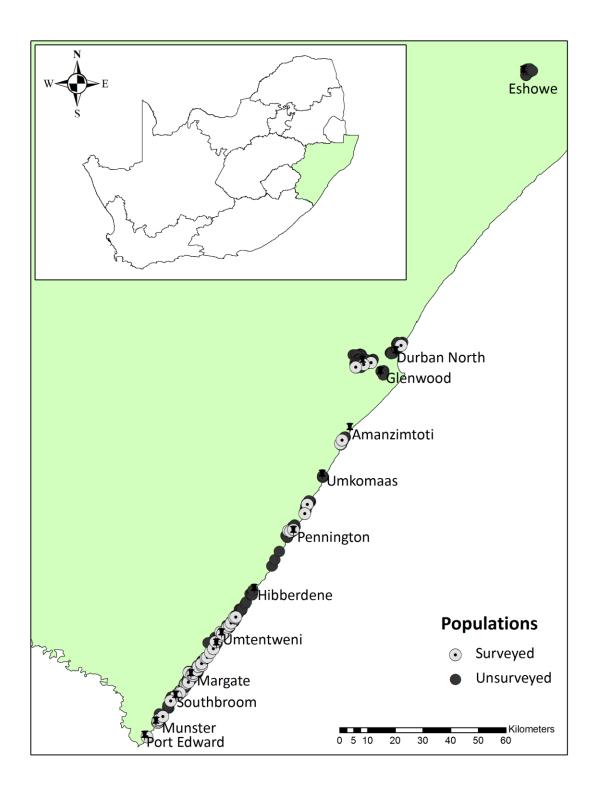
Record number	Accession number	Locality	GPS coordinates
1322	1230365	Kwambonambi, Marche Hotel garden	28.59972° S 32.08361° E
120	1250948	Pinetwon, Japanese Garden	29.83556° S 30.87028° E

Appendix 7. *Epipremnum aureum* pamphlets distributed for additional sightings.





Appendix 8. *Epipremnum aureum* in the KwaZulu-Natal province, South Africa. The geographical distribution contains 321 unsurveyed (i.e. cultivated) and 78 surveyed (i.e. naturalized) populations.



Appendix 9. Raw data of all introduced Monsteroideae and their associated traits that were analysed.

Scientific Name	Introduced	Naturalized	Invasive	Status	Minimum residence time (2016-x)	Minimum residence time_References	Continent of origin	Number of Introduced Regions	Number of Introduced Regions_References
Amydrium humile Schott	1			0	30	Tropicos.org. 2016. Missouri Botanical Garden, St. Louis, MO, USA. http://www.tropicos.org, Accessed 26/09/2016	Asia	1	(Randall, 2007)
Amydrium zippelianum (Schott) Nicolson	1			0	NA		Asia & Oceania	1	(Randall, 2007)
Anadendrum microstachyum (de Vriese & Miq.) Backer & Alderw.	1			0	41	Tropicos.org. 2016. Missouri Botanical Garden, St. Louis, MO, USA. http://www.tropicos.org, Accessed 26/09/2016	Asia	1	(Randall, 2007)
Epipremnum amplissimum (Schott) Engl.	1			0	49	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	Oceania	1	Atlas of Living Australia

Epipremnum aureum (Linden & André) G.S.Bunting	1		1	1	58	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	Oceania	20	eMonocot
Epipremnum falcifolium Engl.	1			0	33	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	Asia	1	GBIF
Epipremnum pinnatum (L.) Engl.	1		1	1	152	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	Asia & Oceania	3	eMonocot; USDA, 2011; HEAR
<i>Monstera acuminata</i> K.Koch	1			0	NA		North America	1	GBIF
Monstera adansonii Schott var. adansonii	1			0	35	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	Oceania	2	eMonocot, 2010
Monstera deliciosa Liebm.	1	1		1	45	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	North America	6	eMonocot, 2010; USDA, 2011; GBIF; Randall, 2007

Appendices

<i>Monstera obliqua</i> Miq.	1	0	NA		North & South America & Oceania	1	Randall, 2007
Monstera punctulata (Schott) Schott ex Engl.	1	0	30	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	North America	1	Randall, 2007
Monstera standleyana G.S.Bunting	1	0	37	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	North America	1	Randall, 2007
Monstera subpinnata (Schott) Engl.	1	0	NA		South America	1	Randall, 2007
Monstera tenuis K.Koch	1	0	22	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	North America	1	Randall, 2007
Rhaphidophora australasica F.M.Bailey	1	0	35	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	Oceania	1	GBIF
Rhaphidophora conica Engl.	1	0	63	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	Oceania	1	GBIF

Rhaphidophora decursiva (Roxb.) Schott	1	0	44	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	Asia	3	GBIF
Rhaphidophora korthalsii Schott	1	0	NA		Asia & Oceania	1	Randall, 2007
Rhaphidophora megaphylla H.Li	1	0	11	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	Asia	1	GBIF
Rhaphidophora pachyphylla K.Krause	1	0	44	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	Oceania	1	Atlas of Living Australia
Rhaphidophora pertusa (Roxb.) Schott	1	0	47	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	Asia	2	GBIF
Rhaphidophora schottii Hook.f.	1	0	21	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	Asia	1	GBIF
<i>Rhaphidophora</i> <i>versteegii</i> Engl. & K.Krause	1	0	NA		Oceania	1	Randall, 2007

Rhodospatha moritziana Schott	1	0	15	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	North & South America		
Rhodospatha venosa Gleason	1	0	11	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	South America	1	GBIF
Scindapsus pictus Hassk.	1	0	58	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	Asia	3	GBIF
Spathiphyllum blandum Schott	1	0	26	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	North America	1	Randall, 2007
Spathiphyllum cannifolium (Dryand. ex Sims) Schott	1	0	8	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	South America & Oceania	3	GBIF
Spathiphyllum cochlearispathum (Liebm.) Engl.	1	0	27	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	North America	2	GBIF

Spathiphyllum commutatum Schott	1	0	NA		Asia & Oceania	1	Randall, 2007
Spathiphyllum floribundum (Linden & André) N.E.Br.	1	0	12	Tropicos.org. 2016. Missouri Botanical Garden, St. Louis, MO, USA. http://www.tropicos.org, Accessed 26/09/2016	South America	2	GBIF
Spathiphyllum friedrichsthalii Schott	1	0	76	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	North & South America	1	GBIF
Spathiphyllum kochii Engl. & K.Krause	1	0	54	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	South America	2	GBIF
Spathiphyllum lanceifolium (Jacq.) Schott	1	0	NA		South America	1	GBIF
Spathiphyllum ortgiesii Regel	1	0	17	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	North America	1	GBIF
Spathiphyllum patinii (R.Hogg) N.E.Br.	1	0	44	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	South America	2	Randall, 2007; GBIF

Appendices

Spathiphyllum phryniifolium Schott	1	0	11	Tropicos.org. 2016. Missouri Botanical Garden, St. Louis, MO, USA. http://www.tropicos.org, Accessed 26/09/2016	North & South America	1	Randall, 2007
<i>Spathiphyllum wallisii</i> Regel	1	0	40	GBIF (2016). Global Biodiversity Information Facility, http://www.gbif.org/, Accessed 26/09/2016	South America	4	GBIF; eMonocot, 2010; HEAR
Stenospermation amomifolium (Poepp.) Schott	1	0	6	Tropicos.org. 2016. Missouri Botanical Garden, St. Louis, MO, USA. http://www.tropicos.org, Accessed 26/09/2016	South America	1	GBIF
Stenospermation multiovulatum (Engl.) N.E.Br.	1	0	20	Tropicos.org. 2016. Missouri Botanical Garden, St. Louis, MO, USA. http://www.tropicos.org, Accessed 26/09/2016	North & South America	1	Randall, 2007
Stenospermation popayanense Schott	1	0	NA		South America	1	Randall, 2007

Appendix 10. The outcome of the weed risk assessment of *M. deliciosa* in South Africa.

Family:	Araceae	Date assessed:	15/09/2016
Taxon:	Monstera deliciosa Liebm.	Assessor:	D. Moodley
Common name:	Swiss Cheese Plant	AWRA score:	7
Synonyms:	Monstera borsigiana K.Koch Monstera deliciosa var. borsigiana Engl. Monstera deliciosa var. sierrana G.S.Bunting Monstera lennea K.Koch Monstera tacanaensis Matuda Philodendron anatomicum Kunth	Recommendation:	Reject

Refer to:

Pheloung *et al.* (1999) A weed risk assessment model for use as a biosecurity tool evaluating plant introductions. *Journal of Environmental Management* 57, pp 239-251.

Gordon *et al.* (2010) Guidance for addressing the Australian Weed Risk Assessment questions. *Plant Protection Quarterly* 25, pp 56-74.

Question	Answer	Score	Possible scores
1.01 Is the species highly domesticated?	У	-3	0 or -3
1.02 Has the species become naturalized where grown?	у	1	-1 or 1
1.03 Does the species have weedy races?	n	-1	-1 or 1
2.01 Species suited to South African climates	2	2	0, 1 or 2
2.02 Quality of climate match data (0-low; 1-intermediate; 2-high)	2	2	0, 1 or 2
2.03 Broad climate suitability (environmental versatility)	n	0	0, 1 or 2
2.04 Native or naturalized in regions with tropical or subtropical climates	У	1	0 or 1
2.05 Does the species have a history of repeated introductions outside its natural range?	у	2	Refer to "lookup" table (Appendix 2) from Pheloung et al., 1999
3.01 Naturalized beyond native range	У	2	Refer to "lookup" table (Appendix 2) from Pheloung et al., 1999
3.02 Garden/amenity/disturbance weed	У	2	Refer to "lookup" table (Appendix 2) from Pheloung et al., 1999
3.03 Weed of agriculture/horticulture/forestry	n	0	Refer to "lookup" table (Appendix 2) from Pheloung et al., 1999
3.04 Environmental weed	n	0	Refer to "lookup" table (Appendix 2) from Pheloung et al., 1999
3.05 Congeneric weed	n	0	Refer to "lookup" table (Appendix 2)

			from Pheloung et al., 1999
4.01 Produces spines, thorns or burrs	n	0	0 or 1
4.02 Allelopathic	n	0	0 or 1
4.03 Parasitic	n	0	0 or 1
4.04 Unpalatable to grazing animals			-1 or 1
4.05 Toxic to animals	у	1	0 or 1
4.06 Host for recognised pests and pathogens	у	1	0 or 1
4.07 Causes allergies or is otherwise toxic to humans	у	1	0 or 1
4.08 Creates a fire hazard in natural ecosystems	n	0	0 or 1
4.09 Is a shade tolerant plant at some stage of its life cycle	у	1	0 or 1
4.10 Grows on infertile soils	у	1	0 or 1
4.11 Climbing or smothering growth habit	у	1	0 or 1
4.12 Forms dense thickets	n	0	0 or 1
5.01 Aquatic	n	0	0 or 5
5.02 Grass	n	0	0 or 1
5.03 Nitrogen fixing woody plant	n	0	0 or 1
5.04 Geophyte	n	0	0 or 1
6.01 Evidence of substantial reproductive failure in native habitat	n	0	0 or 1
6.02 Produces viable seed	у	1	-1 or 1
6.03 Hybridises naturally			-1 or 1
6.04 Self-fertilisation	n	-1	-1 or 1
6.05 Requires specialist pollinators	n	0	0 or -1
6.06 Reproduction by vegetative propagation	у	1	-1 or 1
6.07 Minimum generative time (years)	3.0	0	-1, 0, or 1
7.01 Propagules likely to be dispersed unintentionally	у	1	-1 or 1
7.02 Propagules dispersed intentionally by people	у	1	-1 or 1
7.03 Propagules likely to disperse as a produce contaminant	n	-1	-1 or 1
7.04 Propagules adapted to wind dispersal	n	-1	-1 or 1
7.05 Propagules buoyant			-1 or 1

7.06 Propagules bird dispersed			-1 or 1
7.07 Propagules dispersed by other animals (externally)	n	-1	-1 or 1
7.08 Propagules dispersed by other animals (internally)			-1 or 1
8.01 Prolific seed production	n	-1	-1 or 1
8.02 Evidence that a persistent propagule bank is formed (>1 yr)	n	-1	-1 or 1
8.03 Well controlled by herbicides			-1 or 1
8.04 Tolerates or benefits from mutilation, cultivation or fire	у	1	-1 or 1
8.05 Effective natural enemies present in South Africa			-1 or 1

Notes and References:

1.01	"a popular foliage	http://floridata.com/Plants/Araceae/Monstera+deliciosa/794; Mayo, S. 1985. Araceae.
	houseplant"; "common	In: R. Polhill (ed.). Flora of Tropical East Africa. 71 pp.
	ornamental climber	
	throughout the world"	
1.02	Florida; New Zealand and	http://regionalconservation.org/ircs/database/plants/PlantPage.asp?TXCODE=Monsdeli;
	Galapagos	http://www.hear.org/gcw/species/monstera_deliciosa/
1.03	"Although its been around for	http://www.ourhouseplants.com/plants/monstera-deliciosa;
	ages, there are still very few	http://keyserver.lucidcentral.org/weeds/data/media/Html/monstera_deliciosa.htm
	cultivars you can buy". There	
	is no evidence that cultivars	
	are weedy	
2.01	Bioclimatic model	Current study
2.02	Bioclimatic model	Current study
0.00	0.76.146.666.166.1	1.11
2.03	Suited to tropical and warm	http://www.gbif.org/species/2868241
	subtropical climates	
2.04	"Occasionally naturalised in	http://keyserver.lucidcentral.org/weeds/data/media/Html/monstera_deliciosa.htm

	the warmer temperate, sub- tropical and tropical regions of eastern Australia"	
2.05	"It was introduced into cultivation in England in 1752; reached Singapore in 1877 and India in 1878. Specimens of the fruit were exhibited by the Massachusetts Horticultural Society in 1874 and 1881. It has become familiar as an ornamental in most of the warm countries of the world and is widely used in warm and temperate regions"	Morton, J. 1987. Ceriman. p. 15–17. In: Fruits of warm climates. Julia F. Morton, Miami, FL.
3.01		http://www.hear.org/gcw/species/monstera_deliciosa/
3.02	"This very common garden plant has become a weed of riparian areas and urban bushland, particularly in the warmer parts of eastern Australia. It is relatively common in coastal areas, usually growing where garden refuse has been dumped"	http://keyserver.lucidcentral.org/weeds/data/media/Html/monstera_deliciosa.htm
3.03	No evidence	

3.04	Not known to be invasive	
3.05	No evidence	
4.01	Species does not have these traits	
4.02	No evidence	
4.03	No evidence	
4.04	Unknown	
4.05	"The oxalic acid, and possibly other unidentified principles, in the unripe fruit, the floral remnants of the ripe fruit, and all parts of the plant, cause oral and skin irritation"	Morton, J. 1987. Ceriman. p. 15–17. In: Fruits of warm climates. Julia F. Morton, Miami, FL.
4.06	"The following diseases have been recorded in Florida: leaf spot caused by Leptosphaeria sp., Macrophoma philodendri, Phytophthora sp., and Pseudomonas cichorri; anthracnose from Glomerella cingulata; bacterial soft rot from infection by Erwinia carotovora; and root rot caused by Pythium splendens and Rhizoctonia	Morton, J. 1987. Ceriman. p. 15–17. In: Fruits of warm climates. Julia F. Morton, Miami, FL.

	solani"	
4.07	"The oxalic acid, and possibly other unidentified principles, in the unripe fruit, the floral remnants of the ripe fruit, and all parts of the plant, cause oral and skin irritation"	Morton, J. 1987. Ceriman. p. 15–17. In: Fruits of warm climates. Julia F. Morton, Miami, FL.
4.08	Lush evergreen foliage	
4.09	"Plant grows in the shade"	http://hort.ufl.edu/database/documents/pdf/shrub_fact_sheets/mondela.pdf
4.10	"Soil tolerances: slightly alkaline; clay; sand; acidic; loam"	http://hort.ufl.edu/database/documents/pdf/shrub_fact_sheets/mondela.pdf
4.11	"Vigorous vine which quickly climbs up any shaded tree trunk"	http://hort.ufl.edu/database/documents/pdf/shrub_fact_sheets/mondela.pdf
4.12	No evidence	
5.01	Epiphytic	
5.02	Vine	
5.03	Araceae family	
5.04	Climber	
6.01	Reproduction is usually by vegetative spread	http://www.terrain.net.nz/friends-of-te-henui-group/weeds/fruit-salad-plant-monstera- deliciosa.html
6.02	"Due to recalcitrant nature of	http://www.tropilab.com/swisscheeseplant.html

	the seeds, they have a short viable life"	
6.03	Unknown	
6.04	"The stigma is receptive before the ripening of the stamens, thus avoiding the self-fertilization"	Madison, M. 1977. A revision of Monstera (Araceae). Gray Herbarium of Harvard University. 100pp.
6.05	Bees and Insects	http://tropical.theferns.info/viewtropical.php?id=Monstera+deliciosa
6.06	"From leaf cuttings, herbaceous stem cuttings, and softwood cuttings"	http://davesgarden.com/guides/pf/go/1204/#b
6.07	"Suckers will fruit in 2 to 4 years, cuttings in 4 to 6 years"	
7.01	"It is mainly found growing where garden waste has been dumped along roadsides and waterways"	http://keyserver.lucidcentral.org/weeds/data/media/Html/monstera_deliciosa.htm
7.02	"Fruit salad plant has a long history as a popular houseplant throughout the world"	Martin, T.J. 2002. A Mexican migrant the naturalisation of <i>Monstera deliciosa</i> (fruit salad plant) in New Zealand. Auckland Botanical Society Journal, 57: 151-154
7.03	No evidence	
7.04	No evidence	
7.05	Unknown	
7.06	Unknown, but likely, since berries are edible	
7.07	No evidence, but fruits have no means for attachment	
7.08	Unknown	

8.01	"Generally there are no seeds, but sometimes, palegreen, hard seeds the size of large peas, may occur in a dozen or so of the segments"; "Seeds are occasionally produced but seedlings take a long time to	Morton, J. 1987. Ceriman. p. 15–17. In: Fruits of warm climates. Julia F. Morton, Miami, FL.; http://www.dpi.nsw.gov.au/data/assets/pdf_file/0003/119775/monsterosa-deliciosa.pdf
9.02	develop fully"	Mustofin A. M. (1092) Manatara propagation by good. Pyulloton' Cocyderaty appaga
8.02	Seeds must be sown immediately	Mustafin, A. M. (1983) Monstera propagation by seed. Byulleten' Gosudarstvennogo Nikitskogo Botanicheskogo Sada, 1983, No.51, pp.29-33, 8
8.03	Unknown. In New Zealand they recommend cut and treat stump using Triclopyr	-
8.04	"I sometimes chop the stem into 3 inch pieces, let them harden off for a day or two, then lay with nodes down on top of the soil. You don't even need leaves to do this. These things will practically propagate themselves"	http://forums.gardenweb.com/discussions/1689993/propagating-monstera-deliciosa
8.05	Unknown	

Appendix 11. Response curves demonstrating the relationship between the environmental variables and the probability of *M. deliciosa* presence. The environmental variables comprise Bio 1 (annual mean temperature), Bio 7 (temperature annual range (i.e. maximum temperature of the warmest month – minimum temperature of the coldest month)), Bio 12 (annual precipitation), and Bio 15 (precipitation seasonality). Temperature variables are measured in degrees Celsius and precipitation variables in mm. Rug lines on the lower x-axes represents values of *M. deliciosa* occurrences, and pseudo-absence values are shown on the upper x-axes.

