# KAJIAN SISTEMATIK KE ATAS GENUS *PLAGIOSTACHYS* RIDL. (ALPINIOIDEAE: ZINGIBERACEAE) DI BORNEO

**AVELINAH JULIUS** 

# **INSTITUT BIOLOGI TROPIKA**

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## DECLARATION

The materials in this thesis are original except for quotations, excerpts, summaries and references, which have been duly acknowledged.

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> AVELINAH JULIUS ITBC, UMS

#### ABSTRACT

The genus Plagiostachys of Borneo was revised. Morphological characteristics in living plants were observed to find new significant characters for the genus classification. This study showed that several characters, such as the shape and the presence or absence of anther crest, the branching pattern, and the bracteole shape were consistent within a species and taxonomically important to differentiate between the closely related species. Seven new species and one new variety were recognized, described and illustrated, namely P. brevicucullata A. Julius & A. Takano, P. longicaudata A. Julius & A. Takano, P. megacarpa A. Julius & A. Takano, P. pustulosa A. Julius & A. Takano, P. roseiflora A. Julius & A. Takano, P. subulata A. Julius & A. Takano, P. viridisepala A. Julius & A. Takano and P. breviramosa J. Cowley var. ecristata A. Julius & A. Takano. A key to species of Bornean Plagiostachys was constructed based on morphological characteristics. Molecular phylogenetic analysis using nuclear DNA sequence data of internal transcribed spacer region (ITS1, ITS2) and 5.8S gene was conducted to examine the phylogenetic status of the genus. The strict consensus tree (Length = 1216; CI = 0.4120; RI = 0.7395; RC = 0.3047) obtained from parsimony analysis of the ITS sequence data showed that the species of Plagiostachys made a strongly supported clade with some Alpinia species which belong to section Alpinia (BS = 92 %). On the other hand, species of *Plagiostachys* comprise three subclades and each subclade is moderately to strongly support with relatively high bootstrap value: Plagiostachys subclade A (BS = 70%), Plagiostachys subclade B (BS = 81%) and Plagiostachys subclade C (BS = 99%). Alpinia is not included into the subclades. It suggests that the Plagiostachys is a different entity from Alpinia but more data is needed to verify this. Therefore, Plagiostachys remains in its current status until further analysis is carried out. The ITS sequences also provided new data for inferring relationships within *Plagiostachys* and allow new interpretations of capsule character (pubescence or glabrous capsule) that maybe of value in future classification.



#### ABSTRAK

# KAJIAN SISTEMATIK KE ATAS GENUS PLAGIOSTACHYS RIDL. (ALPINIOIDEAE: ZINGIBERACEAE) DI BORNEO

Kajian telah dibuat ke atas genus Plagiostachys di Borneo. Ciri-ciri morfologi ke atas tumbuhan hidup telah dikaji untuk mencari ciri-ciri penting bagi pengkelasan genus tersebut. Hasil kajian ini menunjukkan bahawa beberapa ciri seperti bentuk dan ada atau tidak ada 'anther crest', corak percabangan dan bentuk 'bracteole' adalah konsisten di dalam spesies dan penting secara taksonomi untuk membezakan di antara spesies-spesies terdekat. Tujuh spesies baru dan satu variasi baru telah dikenalpasti, dihurai dan diilustrasi, iaitu P. brevicucullata A. Julius & A. Takano, P. longicaudata A. Julius & A. Takano, P. megacarpa A. Julius & A. Takano, P. pustulosa A. Julius & A. Takano, P. roseiflora A. Julius & A. Takano, P. subulata A. Julius & A. Takano, P. viridisepala A. Julius & A. Takano dan P. breviramosa J. Cowley var. ecristata A. Julius & A. Takano. Kekunci spesies Plagiostachys di Borneo telah dibuat berdasarkan ciri-ciri morfologi. Analisis molekul filogenetik menggunakan data jujukan nuklear DNA, 'internal transcribed spacer region' (ITS1, ITS2) dan '5.85' gen telah dibuat untuk mengkaji status filogenetik genus tersebut. Pokok 'strict consensus' (Panjang = 1216; indeks kestabilan = 0.4120; indeks penahanan = 0.7395; indeks terskala = 0.3047) vang diperolehi daripada analisis parsimoni data jujukan ITS menunjukkan bahawa gabungan spesies Plagiostachys dengan beberapa spesies Alpinia adalah disokong kuat (BS = 92 %). Sebaliknya, spesies Plagiostachys vang terdiri daripada tiga 'subclades' dan setiap 'subclade' disokong secara sederhana dan kuat mempunyai nilai 'bootstrap' yang tinggi secara relatif: Plagiostachys subclade A (BS = 70%), Plagiostachys subclade B (BS = 81%) dan Plagiostachys subclade C (BS = 99%). Spesies Alpinia tidak terdapat di dalam setiap 'subclade'. Ini menunjukkan bahawa Plagiostachys mungkin entiti lain daripada Alpinia tetapi data lanjut adalah perlu untuk mengesahkannya. Dengan itu, status semasa Plagiostachys dikekalkan sehingga kajian lanjut dilakukan. Penjujukan ITS juga memberikan informasi baru mengenai hubungan di dalam Plagiostachys seperti kapsul berbulu atau licin yang mana ciri tersebut adalah penting untuk pengkelasan di masa akan datang.



# **KEYWORDS**

Borneo, ITS, *Plagiostachys*, Systematic, Zingiberaceae



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# LIST OF ABBREVIATIONS

BO	:Herbarium Bogor
BORH	:BORNEENSIS herbarium
BS	:Bootstrap value
CI	:Consistency Index
CRP	:Crocker Range Park
DVCA	:Danum Valley Conservation Area
E	:Herbarium Royal Botanic Gardens, Edinburgh
ETS	:External Transcribed Spacer
Fl	:Herbarium Universitatis Florentinae
G	:Herbarium Conservatoire et Jardin botaniques de la Ville de Geneva
HI	:Homoplasy Index
HYO	:Herbarium of Museum of Human Nature & Activities, Hyogo
IBSC	:Herbarium South China Botanical Garden
ITS	:Internal Transcribed Spacer
JSPS	:Japan Society for Promotion of Science (BioTropical resources)
К	:Herbarium Royal Botanic Gardens, Kew
MBCA	:Maliau Basin Conservation Area
ML	:Maximum Likelihood
MP	:Maximum Parsimony
MULTREES	:Multiple trees
MULTREES NHM	:Multiple trees :Herbarium University of Nottingham
NHM	:Herbarium University of Nottingham
NHM NNI	:Herbarium University of Nottingham :Nearest-Neighbour Interchange
NHM NNI PCR	:Herbarium University of Nottingham :Nearest-Neighbour Interchange :Polymerase Chain Reaction
NHM NNI PCR PR	:Herbarium University of Nottingham :Nearest-Neighbour Interchange :Polymerase Chain Reaction :Herbarium National Museum in Prague
NHM NNI PCR PR RC	:Herbarium University of Nottingham :Nearest-Neighbour Interchange :Polymerase Chain Reaction :Herbarium National Museum in Prague :Rescaled Consistency Index
NHM NNI PCR PR RC RDC	:Herbarium University of Nottingham :Nearest-Neighbour Interchange :Polymerase Chain Reaction :Herbarium National Museum in Prague :Rescaled Consistency Index :Rainforest Discovery Centre
NHM NNI PCR PR RC RDC RI	:Herbarium University of Nottingham :Nearest-Neighbour Interchange :Polymerase Chain Reaction :Herbarium National Museum in Prague :Rescaled Consistency Index :Rainforest Discovery Centre :Retention Index
NHM NNI PCR PR RC RDC RI SAN	<ul> <li>:Herbarium University of Nottingham</li> <li>:Nearest-Neighbour Interchange</li> <li>:Polymerase Chain Reaction</li> <li>:Herbarium National Museum in Prague</li> <li>:Rescaled Consistency Index</li> <li>:Rainforest Discovery Centre</li> <li>:Retention Index</li> <li>:Herbarium Sandakan</li> </ul>
NHM NNI PCR PR RC RDC RI SAN SAR	<ul> <li>:Herbarium University of Nottingham</li> <li>:Nearest-Neighbour Interchange</li> <li>:Polymerase Chain Reaction</li> <li>:Herbarium National Museum in Prague</li> <li>:Rescaled Consistency Index</li> <li>:Rainforest Discovery Centre</li> <li>:Retention Index</li> <li>:Herbarium Sandakan</li> <li>:Herbarium Sarawak</li> </ul>
NHM NNI PCR PR RC RDC RI SAN SAR SING	<ul> <li>:Herbarium University of Nottingham</li> <li>:Nearest-Neighbour Interchange</li> <li>:Polymerase Chain Reaction</li> <li>:Herbarium National Museum in Prague</li> <li>:Rescaled Consistency Index</li> <li>:Rainforest Discovery Centre</li> <li>:Retention Index</li> <li>:Herbarium Sandakan</li> <li>:Herbarium Sarawak</li> <li>:Herbarium Royal Botanic Garden, Singapore</li> </ul>
NHM NNI PCR PR RC RDC RI SAN SAR SING SNP	<ul> <li>:Herbarium University of Nottingham</li> <li>:Nearest-Neighbour Interchange</li> <li>:Polymerase Chain Reaction</li> <li>:Herbarium National Museum in Prague</li> <li>:Rescaled Consistency Index</li> <li>:Rainforest Discovery Centre</li> <li>:Retention Index</li> <li>:Herbarium Sandakan</li> <li>:Herbarium Sarawak</li> <li>:Herbarium Royal Botanic Garden, Singapore</li> <li>:Sabah Parks herbarium</li> </ul>
NHM NNI PCR PR RC RDC RI SAN SAR SING SNP TBR	<ul> <li>:Herbarium University of Nottingham</li> <li>:Nearest-Neighbour Interchange</li> <li>:Polymerase Chain Reaction</li> <li>:Herbarium National Museum in Prague</li> <li>:Rescaled Consistency Index</li> <li>:Rainforest Discovery Centre</li> <li>:Retention Index</li> <li>:Herbarium Sandakan</li> <li>:Herbarium Sarawak</li> <li>:Herbarium Royal Botanic Garden, Singapore</li> <li>:Sabah Parks herbarium</li> <li>:Tree Bisection-Reconnections</li> </ul>
NHM NNI PCR PR RC RDC RI SAN SAR SING SNP TBR <i>matK</i> rDNA rRNA	<ul> <li>:Herbarium University of Nottingham</li> <li>:Nearest-Neighbour Interchange</li> <li>:Polymerase Chain Reaction</li> <li>:Herbarium National Museum in Prague</li> <li>:Rescaled Consistency Index</li> <li>:Rainforest Discovery Centre</li> <li>:Retention Index</li> <li>:Herbarium Sandakan</li> <li>:Herbarium Sarawak</li> <li>:Herbarium Royal Botanic Garden, Singapore</li> <li>:Sabah Parks herbarium</li> <li>:Tree Bisection-Reconnections</li> <li>:maturase-K</li> </ul>
NHM NNI PCR PR RC RDC RI SAN SAR SING SNP TBR <i>matK</i> rDNA	<ul> <li>:Herbarium University of Nottingham</li> <li>:Nearest-Neighbour Interchange</li> <li>:Polymerase Chain Reaction</li> <li>:Herbarium National Museum in Prague</li> <li>:Rescaled Consistency Index</li> <li>:Rescaled Consistency Index</li> <li>:Retention Index</li> <li>:Herbarium Sandakan</li> <li>:Herbarium Sarawak</li> <li>:Herbarium Royal Botanic Garden, Singapore</li> <li>:Sabah Parks herbarium</li> <li>:Tree Bisection-Reconnections</li> <li>:maturase-K</li> <li>:ribosomal Deoxyribonucleic acid</li> </ul>

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#### **CHAPTER 1**

#### INTRODUCTION

Zingiberaceae, commonly known as the ginger family, is the largest of the eight families comprising the monophyletic tropical order Zingiberales. The family form a monophyletic group together with Costaceae, Marantaceae, and Cannaceae, and Costaceae is the sister family to Zingiberaceae (Clark et al., 1993; Kress, 1990). It is distributed pantropically with the center of diversity in South and South East Asia. 53 genera and over 1300 species are known in this family (Kress et al., 2002). Of these, 24 genera and about 600 species occur in the Malesian region, throughout Malaysia, Indonesia, Brunei, Singapore, the Philippines, and Papua New Guinea (Larsen et al., 1999). While for Borneo, there are 18 genera and over 200 species have been reported (Poulsen, 2006). Among these, the genera such as Alpinia, Amomum, Boesenbergia, Etlingera, Hornstedtia and Plagiostachys are commonly found in Borneo. Within the family Zingiberaceae, four tribes are traditionally recognized, namely Alpinieae Meisn., Globbeae Meisn., Hedychieae Petersen and Zingibereae Burtt & Olatunji (Burtt & Smith, 1972; Smith, 1982). However, the current classification based on DNA sequence data presented by Kress et al. (2002) has recognized four subfamilies; the Siphonochiloideae



W. J. Kress (the genus *Siphonochilus* only), the Tamijioideae W. J. Kress (*Tamijia*), the Zingiberoideae Haask. (including the former tribes Hedychieae and Globbeae), and the Alpinioideae Link (comprises most of the former Alpinieae). The subfamily Alpinioideae, which is characterized by the plane of distichy of leaves perpendicular to rhizome and the reduction or absence of the two lateral staminodes, are further subdivided into two tribes, i.e. Alpinieae A. Rich. and Riedelieae W. J. Kress. The Alpinieae, defined by having fleshy or indehiscent fruit and lacking extrafloral nectarines, includes all remaining former Alpinieae, e.g. *Alpinia, Amomum, Etlingera, Hornstedtia* and *Plaglostachys*.

*Plaglostachys* Ridl., is a complex genus and comprising about 24 species. It is distributed in Indochina to Malesia with the center of diversity in Borneo (Smith, 1985). The genus is generally the constituent of the undergrowth in the tropical forests. The plants inhabiting damp places in the primary forest, but sometimes they are found in old secondary forest and disturbed sites. It has been distinguished from the other genera in Zinglberaceae by the apparently lateral, tightly congested inflorescence, which is, in fact, terminal on the short stem of the leafy shoots, and breaks through the leaf sheaths usually just above ground level or sometimes in the middle (Smith, 1990). The flower is subtended by a usually tubular bracteole, and the labellum, although small and rather fleshy, is petaloid with divergent venation (for some species). The



inflorescence is, in certain species, very mucilaginous, where the bracteole and calyx are fleshy, and tend to decay soon.

In many instances, the taxonomic study of *Plagiostachys* have been devoted to Borneo (Gobilik et al., 2005; Sakai & Nagamasu, 2003) compared to the other areas such as the Philippines, Indonesia and Peninsular Malaysia, as very few literature is available. The genus is still poorly known among gingers and needs critical revision and more research work (Larsen et al., 1999; Smith, 1982). The problem is mainly because of the mucilaginous nature of the inflorescences. This hinders close examination of inflorescences and flowers in dried herbarium specimens. It is not only because of the flower structure will be lost but also the fleshy and decaying calyx and bracteole, virtually unusable when dealing with dry specimen. Another problem is, most of the herbarium specimens are incomplete and seem to have been collected in fruits rather than in flower (Cowley, 1999). Thus, studying on many living plants and doing observations in the field are the most important steps in revising the genus. These will enable one to gather many data on variation of important characters in the species of Plagiostachys that are useful for its classification.

The existing classification of Bornean *Plagiostachys* has been based on Smith (1985). She classified the genus into two informal groups: the Group I, defined by having mucilaginous inflorescences, the bracteoles distinctly tubular, early decaying



and only the very basal part remaining, the calyx is fleshy and decaying in the upper part at least, the style sometimes adnate to the wall of the corolla tube above the epigynous glands, the capsule is oblong-pyriform, angled and rarely globose, the lique is bilobed, and the flowers are yellow and white with some pink; whereas the Group II is characterized by having non-mucilaginous inflorescences, the bracteoles tubular at base or open, and partially decaying or persistent, the calyx is not fleshy and not decaying, the style usually free from the wall of the corolla tube at the base, the capsule is globose, the ligule is truncate or emarginate and rarely bilobed, and the flowers is pinkish-red with a yellow labellum. The classification is, however, disputed by Cowley (1999) and Sakal & Nagamasu (1998) as the characters used for the grouping did not fit with any species described recently. This indicates Smith's grouping can no longer be used. Further investigation of the usefulness of the grouping in the present day is needed.

The study by Kress *et al.* (2005) is the most thorough paper to date addressing the relationships among genera in the subfamily Alpinioideae. The result of their molecular assessment shows that *Amomum* and *Alpinia* are polyphyletic groups and *Plaglostachys* embedded within *Alpinia*. *Alpinia* is the largest and most widespread genus in Zingiberaceae with more than 200 species (Larsen, 1998), occurring from India and Sri Lanka to China, Japan, Southeast Asia, New Guinea, Australia, Fiji and



Samoa (Smith, 1990). The genus, defined by having terminal inflorescences on the leafy shoots and the lateral staminodes are absent or represented by small tooth-like structures at the base of the labellum. This polyphyletic genus was splitted into six clades of species each with varying statistical support. The analysis of their combined data sets (ITS and matk) have demonstrated that Clades I, II, III and IV are each strongly supported as monophyletic (BS > 90%), whereas Clades V (BS = 61%) and Clade VI (BS < 50%) are weakly to poorly supported groups. On the other hand, Plagiostachys is moderately to strongly supported (BS > 70%) as a monophyletic group, but they only included two species in their analyses. Therefore, the monophyletic status of Plagiostachys requires further confirmation by adding more species of the genus. It is also expected that by increasing the taxon sampling in the phylogenetic analysis will illustrate the relationships within Plagiostachys in order to see how congruent are Smith's (1985) informal grouping of the genus. Additionally, any attempt to reclassify Plaglostachys are should make reference to its closely related genus Alpinia. This is not only because Alpinia is the sister genus of Plagiostachys (Kress et al., 2005; Pedersen, 2004), but the laterally produced inflorescence in the latter is also shared with A. hansenii R. M. Sm. and A. havilandii K. Schumn. (Smith, 1990). Both of these species are also apparently having inflorescence breaking through the leaf sheaths near the top and are, as far as is known, never mucilaginous.



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