

Research Article

# New Genus *Lobidiscus* and New Combinations of Genera of *Moutan*, *Liquidambar* and *Hamamelis* (Hamamelidales)

Da-Li Fu<sup>1, 2,\*</sup> 

<sup>1</sup>Research Institute of Non-Timber Forestry, Chinese Academy of Forestry, Zhengzhou, China

<sup>2</sup>Key Laboratory of Non-Timber Forest Germplasm Enhancement & Utilization of National Forestry and Grassland Administration, Zhengzhou, China

## Abstract

Hamamelidales Griseb. (1854) order serves as a foundational group and plays a significant role in the research of evolutionary systematics of the class Rosopsida Batsch (1802) of Fructophyta D. L. Fu & H. Fu (2018). To address the ambiguities surrounding the circumscription of certain genera within this order, such as *Paeonia* L. (1753), *Liquidambar* L. (1753), and *Hamamelis* L. (1753), relevant chloroplast complete genome sequences from the NCBI database were retrieved, and evolutionary analyses were conducted on these sequences in this study. Notably, *Lobidiscus* D. L. Fu, gen. nov., a new genus endemic to America characterized by its distinctly lobed floral discs, has been separated from *Paeonia* L. belonging to Paeoniaceae (Bercht. & J. Presl) Rudolphi, nom. cons. (1830), and *Lobidiscus californicus* (Nutt.) D. L. Fu is designated as its type species. This new genus derives from *Moutan* Rchb. (1827), rather than *Paeonia* L., given that a maximum PHS value of 0.837 were observed between *Lobidiscus brownii* (Hook.) D. L. Fu and *Moutan delavayi* (Franch.) D. L. Fu based on CPCG evolutionary analyses. Meanwhile, the application of the minimum criterion PHS  $\leq$  0.928 (intergeneric, CPCG) for genus classification of Fructophyta D. L. Fu & H. Fu, has led to confirmation of two synonyms for *Liquidambar* L.: *Altingia* Noronha and *Semiliquidambar* H. T. Chang; additionally, three current synonyms for *Hamamelis* L., including *Distylium* Siebold & Zucc., *Parrotia* C. A. Mey., and *Sycopsis* Oliv., have been scientifically identified. In total, 13 novel family names like Liquidambaraceae D. L. Fu, Dianthaceae D. L. Fu, Diospyraceae D. L. Fu and Ilicaceae D. L. Fu have been established, along with two new specific epithets, *Hamamelis hubeiensis* D. L. Fu and *Hamamelis grandifolia* D. L. Fu, and 39 newly valid combinations involving *Hamamelis* L., *Liquidambar* L., *Lobidiscus* D. L. Fu, and *Moutan* Rchb. also been published, such as *Hamamelis annamica* (Gagnep.) D. L. Fu, *Liquidambar cambodiana* (Lecomte) D. L. Fu, *Lobidiscus brownii* (Hook.) D. L. Fu, and *Moutan suffruticosus* (Andrews) D. L. Fu. These contributions will effectively clarify taxonomic nomenclature confusions in a scientific manner while establishing a robust foundation for further research into the evolutionary systems within the order Hamamelidales Griseb.

## Keywords

*Lobidiscus*, *Moutan*, *Liquidambar*, *Hamamelis*, New Genus, New Combination, CPCG (Chloroplast Complete Genome), Genus Minimum Criterion, Typical Algorithm

\*Corresponding author: fu\_dali@163.com (Da-Li Fu)

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## 1. Introduction

The order Hamamelidales Griseb. (1854) is foundational and plays a significant role in the research of evolutionary systematics of the class Rosopsida Batsch (1802) of Fructophyta D. L. Fu & H. Fu (2018). This order encompasses families such as Hamamelidaceae R. Br., nom. cons. (1818), Vitaceae Juss., nom. cons. (1789), Penthoraceae Rydb. ex Britt., nom. cons. (1901) and Paeoniaceae (Bercht. & J. Presl) Rudolphi, nom. cons. (1830) [1, 2]. However, traditional taxonomy and modern phylogenetic approaches reveal persistent taxonomic challenges [3-9], particularly regarding ambiguities in the circumscription of certain genera including *Paeonia* L. (1753), *Liquidambar* L. (1753), and *Hamamelis* L. (1753) within this order.

In 1804, the species of Mudan, renowned as the "King of Flowers" in China, was officially published and named *Paeonia suffruticosa* Andrews (Woody Peony). This was also the first species of *Moutan* taxa to be released, in accordance with the relevant articles of the International Code of Botanical Nomenclature, the name holds absolute precedence in the names of *Moutan* taxa. Consequently, the assertion that "many scholars have confirmed that *P. suffruticosa* is a hybrid formed by repeated hybridization of several species of subsect. *Vaginatae*, based on morphological and molecular biological evidence" [10] is wholly incorrect. In other words, as long as the type of *P. suffruticosa* is authentic, regardless of whether it is collected from cultivated or wild plants, the species must be acknowledged. It implies that all other subsequent species of *Moutan* taxa can be regarded as the hybrids (alternative forms of varieties) of *P. suffruticosa*, yet *P. suffruticosa* cannot be considered as a hybrid of these later species. In 1824, the *Paeonia* sect. *Moutan* DC. was published [11], and in 1827, the genus *Moutan* Rchb. was established. This was a typical course of development in plant taxonomy; however, it failed to draw the attention of taxonomists, despite the *Moutan* taxa having very distinct characteristics, such as the woody stems and the envelope discs [11-19]. Subsequently, species within the *Moutan* taxa such as *Paeonia delavayi* Franch (1887), *Paeonia potaninii* Komarov (1921), and *Paeonia decomposita* Hand-Mazz. (1939), have continued to utilize the genus name *Paeonia* L. Over the past three decades, Chinese *Moutan* taxonomist Hong T. [12-14] and world-renowned *Paeonia* taxonomist Hong D. Y. [15-19] have also maintained the use of the genus name in their publications concerning *Moutan* taxa. However, contemporary phylogenetic studies [20-21] have demonstrated that the genus *Paeonia* L. can be clearly divided into three distinct taxa, now recognized as 3 subgenera or 3 sections. Further investigations are requisite to ascertain whether these two additional taxa may warrant classification as separate genera based on their unique taxonomic characteristics [12-19, 22].

In another instance, phylogenetic analyses have revealed that *Altingia* Noronha and *Semiliquidambar* H. T. Chang are

nested within *Liquidambar* L., leading to the formal transfer of all *Altingia* and *Semiliquidambar* species to *Liquidambar* L. [23-25]. The reasons for this nesting typically involve the synonymy of *Altingia* and *Semiliquidambar* with *Liquidambar* L., or the characterization of *Liquidambar* L. as a diverse genus, with exceptions arising from sampling and analysis errors. Consequently, further analyses are warranted to ascertain whether the other two genera should be considered synonyms of *Liquidambar* L.

Similarly, phylogenetic analyses have indicated a close relationship between the genus *Hamamelis* L. and *Distylium* Siebold & Zucc., *Parrotia* C. A. Mey., and *Sycopsis* Oliv. [26, 27], some of which are nested. Therefore, further analyses are required to determine whether the other three genera should be considered synonyms of *Hamamelis* L. or if *Distylium* Sieb. & Zucc. represents a diverse genus.

Both traditional taxonomy and modern phylogeny encounter limitations in their objectivity and impartiality when it comes to determine whether a genus is the synonym or a diverse genus. The new science evolutionomy has been developed with the publications of the evolutionary continuity principle, the evolutionary particularity principle, the theoretical monograph as *the Theory and Practice of Evolutionomy*, and so on [1-9]. The establishment, publication, and implementation of the minimum criterion  $\text{PHS} \leq 0.928$  (intergeneric, CPCG) for the classification of genus of Fructophyta D. L. Fu & H. Fu has scientifically identified 107 current genus synonyms within the class Scutellopsida D. L. Fu, and the taxonomic confusions of the class has also been scientifically resolved to a certain extent. [4-9].

To scientifically identify the circumscription of certain genera and resolve the synonyms or diverse genera leading to the taxonomic confusions within the three genera of Hamamelidales Griseb., some relevant CPCG sequences from the NCBI (National Center for Biotechnology Information, USA) database have been downloaded and the evolutionary analyses on these sequences have been conducted, and the results are as follows.

## 2. Materials and Methods

### 2.1. CPCG of Hamamelidales

Total 28 CPCG of representative species of three families of Hamamelidales Griseb. were selected from the NCBI database. Their current names, scientific names and CPCG numbers of NCBI are listed in Table 1 to Table 6.

### 2.2. Evolutionary Analyses of CPCG

The evolutionary analyses of CPCG mainly use the typical algorithm [3-9] to determine the relative evolutionary relationships between different taxa by comparing the phyloge-















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