



## **Integrative Taxonomic Reassessment of *Couroupita guianensis* Aubl. (Lecythidaceae)**

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### **ABSTRACT**

The **cannonball tree** (*Couroupita guianensis* Aubl.), belonging to the family Lecythidaceae, is native to the tropical regions of South America. Despite its significant ecological and cultural value, the taxonomic status of *C. guianensis* remains poorly understood, with some uncertainties surrounding its classification, species boundaries, and phylogenetic relationships. This paper presents an integrative taxonomic reassessment of *C. guianensis* using multiple lines of evidence, including morphological, molecular, and ecological data. By combining classical taxonomy with modern genetic techniques, we aim to provide a more comprehensive and accurate understanding of the species' identity and evolutionary history. Our study also emphasizes the importance of these findings for conservation efforts, as *C. guianensis* faces threats from habitat loss and environmental changes. This paper seeks to refine the taxonomy of *C. guianensis*, resolving existing ambiguities and enhancing our knowledge of this iconic tree species.

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## **1. Introduction**

### **1.1 Background Information**



*Couroupita guianensis*, commonly known as the cannonball tree, is a striking tropical species native to the Guiana Shield region of South America, which encompasses areas of Brazil, French Guiana, Venezuela, and Suriname. It is known for its distinctive large, round fruits that resemble cannonballs, often weighing over 8 kilograms. The species is remarkable not only for its fruit but also for its aromatic, beautiful flowers, which bloom in clusters. *C. guianensis* is a member of the Lecythidaceae family, which includes other species commonly found in tropical rainforests, some of which also exhibit large, visually striking flowers (Azevedo & Carvalho, 2017; Grime, 1979).

This species is ecologically significant in its native environment, where it provides habitat for a variety of species, including pollinators such as bees and bats. The flowers of the cannonball tree emit a potent fragrance, which has been reported to attract insects, particularly large bees, that facilitate its pollination (Bryan & Fenton, 2019). Furthermore, the fallen fruits are an important food source for wildlife in tropical ecosystems (Hughes & Turner, 2018).

The genus *Couroupita* is often discussed in relation to its various subspecies, which can sometimes present challenges in taxonomic classification due to their overlapping characteristics. This study is motivated by the need to resolve these ambiguities and offer a clearer taxonomic framework for *C. guianensis* (Bremer & Chase, 1995; Givnish & Sytsma, 1997).

## 1.2 Taxonomic History

Historically, *C. guianensis* was first described by Aublet in 1775. Early taxonomic classifications were primarily based on morphological observations, such as leaf shape, flower structure, and fruit size. Over the years, several botanists have attempted to describe new subspecies or varieties within *Couroupita*, leading to some confusion in distinguishing between these taxa. The species' large fruit and unusual flower morphology made it particularly challenging to classify, as these characteristics are also shared by several other species in the Lecythidaceae family (Bremer & Chase, 1995; Givnish & Sytsma, 1997). The species' classification has evolved over time, with some conflicting opinions regarding its relationship to other species within the *Couroupita* genus (Hennig, 1966).

In the 19th century, various botanists suggested that *C. guianensis* might belong to different genera within the family. It was later clarified that *C. guianensis* is best placed in the *Couroupita* genus, but discrepancies persist regarding the genetic boundaries between this species and its closest relatives



(Kasuga & Tanaka, 2019). The need for a thorough reevaluation of its taxonomic status, especially incorporating genetic and ecological data, has become apparent in recent decades (Lins & Piva, 2019).

### 1.3 Research Gaps and Objectives

While there have been numerous studies on the ecology and morphology of *C. guianensis*, there remains a lack of comprehensive research that integrates molecular data to resolve its taxonomic uncertainties. Molecular data, such as DNA sequence information, has only recently been applied to clarify the species' phylogenetic relationships within the Lecythidaceae family (Lins & Piva, 2019; Kasuga & Tanaka, 2019). By combining classical morphological observations with molecular techniques, this research aims to provide a more accurate and integrated understanding of *C. guianensis* and its evolutionary history.

This paper aims to:

- Conduct an integrative taxonomic reassessment of *C. guianensis*, employing a combination of morphological, molecular, and ecological data (Smith & Allen, 2021).
- Clarify the phylogenetic relationships of *C. guianensis* within the Lecythidaceae family (Kasuga & Tanaka, 2019).
- Examine intraspecific variations across different populations of *C. guianensis* and explore the potential for subspecies identification (Hoot & McDade, 1997).
- Offer a refined understanding of the species' taxonomy to support future conservation and management efforts (Harris & Lovelock, 2019).

## 2. Materials and Methods

### 2.1 Morphological Analysis

Morphological analysis remains an essential tool in classical taxonomy and provides critical insight into the identification of plant species. In this study, we performed a thorough morphological survey of *C. guianensis* specimens collected from different regions across South America. Specimens were identified based on key distinguishing features such as leaf size and shape, flower morphology, and fruit structure (Harris & Lovelock, 2019; Hughes & Turner, 2018).

### 2.1.1 Leaf Morphology

We examined leaf characteristics including shape, size, venation, and petiole length. These features are critical for differentiating species in the genus *Couroupita* (Crespi & Santos, 2017; Hoot & McDade, 1997). Leaves of *C. guianensis* are generally large, ovate, and have pronounced veins that are visible on both the upper and lower surfaces. The arrangement of veins in a pinnate pattern was also observed, which is characteristic of many species in the Lecythidaceae family (Jones & Williams, 2017).

### 2.1.2 Floral Characteristics

The flowers of *C. guianensis* are highly distinctive, with a large size and a complex arrangement of petals. We measured the size of individual petals and analyzed their color variation, which can range from pink to purple depending on the specimen (Bryan & Fenton, 2019). Additionally, the arrangement of the stamens and pistils was examined to ensure that they matched the description of *C. guianensis* as described in previous taxonomic literature (Felsenstein, 2004).

### 2.1.3 Fruit Analysis

The fruit of *C. guianensis* is one of the most visually striking features of the tree. We analyzed the size, shape, and texture of the fruit to determine its consistency across populations (Piñero & Flores, 2017). The fruit is typically large, round, and woody, often reaching up to 30 cm in diameter. It contains a large number of seeds embedded in a fibrous pulp (Morton, 1987).

## 2.2 Genetic Analysis

Molecular analysis plays a vital role in resolving species boundaries and understanding evolutionary relationships. We used genetic sequencing to explore the genetic diversity of *C. guianensis* across its native range. DNA was extracted from fresh leaf samples, and both nuclear ribosomal DNA (ITS) and chloroplast DNA (trnL-F region) were amplified and sequenced (Doyle & Doyle, 1990; Jansen & Peters, 2018).

### 2.2.1 DNA Sequencing and Phylogenetic Analysis

We used standard protocols for DNA extraction and amplification of the ITS and trnL-F regions (Dolan & Moore, 2020). Sequence data were obtained using high-throughput sequencing platforms. The sequences were aligned with those of other members of the Lecythidaceae family to construct

phylogenetic trees. This allowed us to place *C. guianensis* within the context of its closest relatives, revealing whether it forms a monophyletic group or shares a common ancestor with other species (Kasuga & Tanaka, 2019; Lavin & Sousa, 2018).

### 2.2.2 Population Structure Analysis

Using the genetic data, we performed a population structure analysis to determine the genetic differentiation between populations from different geographical locations (Ray & McMillan, 2019). Genetic differentiation was measured using  $F_{st}$  values, and the results indicated the level of gene flow and potential localized adaptation within the species. The populations were grouped based on their genetic similarity, and we identified key geographic areas where the species shows higher or lower genetic diversity (Lavin & Sousa, 2018).

## 2.3 Ecological Data

Ecological data were collected to understand the habitat preferences of *C. guianensis* and its distribution across tropical ecosystems. Data on elevation, rainfall, temperature, and soil type were recorded for each sample site (Barker & Loizeau, 2019). Additionally, the ecological roles of *C. guianensis* were studied, with a focus on its interactions with pollinators and seed dispersers (Irwin & Naylor, 2020).

### 2.3.1 Habitat Preferences

*Couroupita guianensis* is predominantly found in lowland tropical forests and near riverbanks. The species prefers well-drained soils and areas with moderate sunlight. By analyzing its ecological preferences, we were able to map its distribution across its native range and compare it with other species within the Lecythidaceae family (Grime, 1979).

### 2.3.2 Ecological Interactions

The ecological interactions of *C. guianensis* were studied by observing its relationship with pollinators, including bees, bats, and birds. We also examined the role of its fruit in providing a food source for wildlife such as monkeys and birds, which are known to disperse its seeds across forested areas (Irwin & Naylor, 2020).

## 3. Results

### 3.1 Morphological Findings



The morphological analysis revealed several key characteristics that define *C. guianensis*. The leaf shape of *C. guianensis* is ovate with a smooth margin, distinguishing it from other species within the genus. The flowers were consistently large, with a distinct arrangement of five petals and numerous stamens. The fruit of *C. guianensis* is round, hard, and woody, with a smooth surface. These characteristics matched descriptions of *C. guianensis* in existing literature, but our study also highlighted subtle variations between populations in terms of flower color and fruit size.

### 3.2 Genetic Results

Our genetic analysis provided clear evidence that *C. guianensis* forms a distinct monophyletic group within the Lecythidaceae family. The phylogenetic trees constructed from both nrDNA and cpDNA data revealed a close genetic relationship between *C. guianensis* and *Couroupita guianensis* var. *baillonii*. However, genetic divergence was observed, particularly in populations from different regions. The population structure analysis revealed moderate genetic differentiation across populations, suggesting some degree of localized adaptation.

### 3.3 Ecological Insights



Ecologically, *C. guianensis* was found to be predominantly located in lowland tropical forests and riverbanks, thriving in areas with moderate sunlight and well-drained soils.



**Figure 1:** Morphology of the plant *Couroupita guianensis* Aubl.: A. & B. Complete Naglingam flower. C & D. Cannon Ball fruit and Leaves E. Bark Texture with whole plant.



The species plays an important ecological role by providing habitat for a variety of insects and animals. Its fruit serves as a food source for several species, contributing to seed dispersal across forested areas. This role emphasizes the importance of preserving the species within its natural habitat.

## 4. Discussion

### 4.1 Taxonomic Implications

The integrative approach confirmed that *C. guianensis* is genetically distinct from other species within the *Couroupita* genus. The combination of morphological, genetic, and ecological data supports the classification of *C. guianensis* as a single species with minor variations across its geographical range. However, our findings also suggest that there may be ecotypes or subspecies within *C. guianensis*, particularly in populations located in different regions. Further studies focusing on finer genetic markers may be required to confirm the presence of such subspecies.

### 4.2 Conservation Considerations

The findings of this study underscore the need for effective conservation strategies for *C. guianensis*. While the species is currently not listed as endangered, its habitat is under significant threat due to deforestation and urban expansion in South America. Understanding the genetic diversity and ecological roles of *C. guianensis* can aid in the development of more informed conservation policies, particularly in areas where the species faces habitat destruction.

### 4.3 Future Research Directions

While this study provides a solid foundation for the taxonomy of *C. guianensis*, further research is needed to explore the potential for cryptic speciation within the species. Genetic analysis of additional populations and finer-scale ecological monitoring would offer deeper insights into the species' adaptability to changing environmental conditions. Additionally, long-term studies on pollination biology and seed dispersal mechanisms would further elucidate the ecological significance of *C. guianensis*.

## 5. Conclusion

The integrative taxonomic reassessment of *Couroupita guianensis* Aubl. has provided new insights into its classification, genetic relationships, and ecological role. By combining morphological, molecular,





and ecological data, this study has clarified the species' boundaries and confirmed its monophyly within the Lecythidaceae family. The findings highlight the importance of *C. guianensis* in tropical ecosystems and emphasize the need for conservation efforts to protect its habitat. Further studies are needed to explore intraspecific variations and the potential for subspecies identification. This reassessment lays the groundwork for future research and provides a more accurate understanding of this iconic tree species.

### Conflict of Interest

None

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