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Evolution of species diversity in the genus *Chamaecostus* (Costaceae): molecular phylogenetics and morphometric approaches

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Abstract

While most species within the genus *Chamaecostus* (Costaceae) are well defined, the broad geographic range and long list of synonyms associated with *Chamaecostus subsessilis* led us to believe there may be some cryptic species within the complex. We thus investigate the phylogenetic relationships of species in the *Chamaecostus* lineage and specifically test the monophyly and diversity of the *Chamaecostus subsessilis* species complex from a population perspective by analyzing molecular sequence data and leaf morphometrics. We interpret evolutionary trends across the entire genus based on a molecular character-based phylogenetic hypothesis that includes all currently described species of *Chamaecostus*. Our results show that while *Chamaecostus* is strongly monophyletic, *C. cuspidatus* is found to be sister to a clade of some but not all samples of *C. subsessilis*, making it necessary to acknowledge more than one species in the *C. subsessilis* complex. Herbarium specimens of the *C. subsessilis* complex could be assigned based on geographic proximity to one of the major three clades recovered in the phylogenetic analysis. Leaf morphometric measurements were performed on each of these lineages and traits were tested to detect differences among phylogenetic lineages. We conclude by proposing the recognition of a new combination, *Chamaecostus acaulis*, which we describe.

Keywords: species complex, morphometrics, tropical plant taxonomy, spiral gingers, South America

Introduction

Most species are identified on the basis of incongruent patterns and discontinuity of trait variation across individual specimens. Individuals comprising a species manifest their variability in rather continuous variation, and their integrity can only be understood by sampling through the range of this variation (Frost & Kluge 1994), which can then provide the basis for ascertaining trends across the species unit. These observed trends frequently support the recognition of taxa. However, morphological discontinuity does not always co-occur with lineage splitting: isolated subpopulations bearing significant genetic structure can lack phenotypic differences between them, making species delimitations challenging (DeSalle *et al.* 2005, Padial & De La Riva 2009, Padial *et al.* 2010, Florio *et al.* 2012). Such cryptic speciation is characterized by two or more morphologically indistinguishable groups of organisms that are found to belong to distinct evolutionary lineages (Sáez & Lozano 2005). Perceived cryptic speciation can also derive from our inability to distinguish important, and not always prominent, morphological differences (Shaffer & Thomson 2007). Using a phylogenetic lineage approach, species-level phylogenies and networks are able to provide a consistent and predictive evolutionary understanding of species limits by helping to identify unique evolutionary entities among population-level sampling (Funk & Omland 2003, Goldstein & DeSalle 2000). The phylogenetic approach to species delimitation is particularly promising because distinct species are interpreted as being on separate evolutionary trajectories (Hey & Pinho 2012), which, in some cases, are expected to continue to diverge even in the absence of reproductive barriers (Rieseberg *et al.* 2004).

Species of *Chamaecostus* Specht & Stevenson (2006: 157) are fairly distinguishable from one another, with the sole exception being the *C. subsessilis* (Nees & Mart.) Specht & Stevenson (2006: 158) species complex. Taxonomic

Since there is substantial overlap in characters between the two (Figure 3, Table 1), we strongly suggest that location should be taken into account when identifying both species, in particular because of the strong geographic structure resolved between the populations analyzed here. *Chamaecostus acaulis* occurs West of the Araguaia River valley, through Peruvian, Bolivian and Brazilian South Amazonia, and in Western and Southern portions of the Central Brazilian Shield, while *Chamaecostus subsessilis* occurs East from the Araguaia River valley, and within most of the Central Brazilian Shield and in transition forests between Cerrado and Central Atlantic Rain Forest (Figure 2B). However, potentially sympatric populations may occur within the Araguaia River valley, and further detailed analyses of the populations in this transition zone are necessary.

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References

- Darriba, D., Taboada, G.L., Doallo, R. & Posada, D. (2012) jModelTest 2: more models, new heuristics and parallel computing. *Nature Methods* 9: 772.
<http://dx.doi.org/10.1038/nmeth.2109>
- de Queiroz, K. (1998) The general lineage concept of species, species criteria, and the process of speciation: A conceptual unification and terminological recommendations. In: Howard, D.J. & Berlocher, S.H. (Eds.) *Endless forms: Species and speciation* Oxford University Press, New York, pp. 57–75.
- de Queiroz, K. (2007) Species concepts and species delimitation. *Systematic Biology* 56: 879–886.
<http://dx.doi.org/10.1080/10635150701701083>
- DeSalle, R., Egan, M., Sidall, M. (2005) The unholy trinity: taxonomy, species delimitation and DNA barcoding. *Philosophical Transactions of the Royal Society B* 360: 1905–1916.
<http://dx.doi.org/10.1098/rstb.2005.1722>
- Doyle, J.J. & Doyle, J.L. (1990) A rapid total DNA preparation procedure for fresh plant tissue. *Focus* 12:13–15.
- Edgar, R.C. (2004) MUSCLE: multiple sequence alignment with high accuracy and high throughput. *Nucleic Acids Research* 32:1792–1797.
<http://dx.doi.org/10.1093/nar/gkh340>
- Engler, H.G.A. (Ed.) (1904) *Pflanzenreich IV*. 46 (Heft 20).
- Faria, A.P.G., Wendt, T., Brown, G.K. (2010) A revision of *Aechmea* subgenus *Macrochordion* (Bromeliaceae) based on phenetic analyses of morphological variation. *Botanical Journal of the Linnean Society* 162: 1–27.
<http://dx.doi.org/10.1111/j.1095-8339.2009.01019.x>
- Florio, A.M., Ingram, C.M., Rakotondravony, H.A., Louis, Jr. E.E. & Raxworthy, C.J. (2012) Detecting cryptic speciation in the widespread and morphologically conservative carpet chameleon (*Furcifer lateralis*) of Madagascar. *Journal of Evolutionary Biology* 25: 1399–1414.
<http://dx.doi.org/10.1111/j.1420-9101.2012.02528.x>
- Frost, D.R. & Kluge, A.G. (1994) A consideration of epistemology in systematic biology, with special reference to species. *Cladistics* 10: 259–294.
<http://dx.doi.org/10.1111/j.1096-0031.1994.tb00178.x>
- Funk, D.J. & Omland, K.E. (2003) Species-level paraphyly and polyphyly: frequency, causes, and consequences, with insights from animal mitochondrial DNA. *Annual Review of Ecology, Evolution and Systematics* 34: 397–423.
<http://dx.doi.org/10.1146/annurev.ecolsys.34.011802.132421>
- Gagnepain, F. (1902) *Bulletin de la Société Botanique de France, Series 4* 2: 102–103.
- Goldstein, P.Z. & DeSalle, R. (2000) Phylogenetic species, nested hierarchies, and character fixation. *Cladistics* 16: 364–384.
<http://dx.doi.org/10.1111/j.1096-0031.2000.tb00356.x>
- Goldstein, P.Z., DeSalle, R., Amato, G. & Vogler, A.P. (2000) Conservation genetics and the species boundary. *Conservation Biology* 14: 120–131.
<http://dx.doi.org/10.1046/j.1523-1739.2000.98122.x>
- Guindon, S. & Gascuel, O. (2003) A simple, fast, and accurate algorithm to estimate large phylogenies by maximum likelihood. *Systematic*

- Biology* 52: 696–704.
<http://dx.doi.org/10.1080/10635150390235520>
- Hey, J. & Pinho, C. (2012) Population genetics and objectivity in species diagnosis. *Evolution* 66: 1413–1429.
<http://dx.doi.org/10.1111/j.1558-5646.2011.01542.x>
- Huelsenbeck, J.P. & Ronquist, F. (2001) MRBAYES: Bayesian inference of phylogenetic trees. *Bioinformatics* 17: 754–755.
<http://dx.doi.org/10.1093/bioinformatics/17.8.754>
- Hwang, L.H., Hwang, S.Y. & Lin, T.P. (2000) Low chloroplast DNA variation and population differentiation of *Chamaecyparis formosensis* and *Chamaecyparis taiwanensis*. *Taiwan Journal of Forest Science* 15: 229–236.
- Irwin, D.E. (2002) Phylogeographic breaks without geographic barriers to gene flow. *Evolution* 56: 2383–2394.
[http://dx.doi.org/10.1554/0014-3820\(2002\)056\[2383:PBWGBT\]2.0.CO;2](http://dx.doi.org/10.1554/0014-3820(2002)056[2383:PBWGBT]2.0.CO;2)
- Kay, K.M., Reeves, P.A., Olmstead, R.G. & Schemske, D.W. (2005) Rapid speciation and the evolution of hummingbird pollination in neotropical *Costus* subgenus *Costus* (Costaceae): evidence from nrDNA ITS and ETS sequences. *American Journal of Botany* 92: 1899–1910.
<http://dx.doi.org/10.3732/ajb.92.11.1899>
- Kuntze, C.E.O. (1891) *Revisio Generum Plantarum* 2: 687.
- Linnaeus, C. (1753) *Species Plantarum* 1: 1–560.
- Loesener, L.E.T. (1929) *Notizblatt des Botanischen Gartens und Museums zu Berlin-Dahlem* 10: 714.
<http://dx.doi.org/10.2307/3994705>
- Maas, P.J.M. (1972) Costoideae (Zingiberaceae). In: *Flora Neotropica. Monograph* 8. Hafner, New York, New York, USA.
- Maas, P.J.M. (1976) Notes on New World Zingiberaceae. *Acta Botanica Neerlandica* 24: 469.
- Maas, P.J.M. (1977) *Renealmia* (Zingiberaceae-Zingiberoideae) Costoideae (Additions) (Zingiberaceae). *Flora Neotropica. Monograph* 18. Hafner, New York, New York, USA.
- Martius, C.F.P. (Ed.) (1890) *Flora Brasiliensis*. Vol III. Pars III, pp. 1–128.
- Moore, S. (1895) *Transactions of the Linnean Society Series 2* 4: 480.
- Nees, C.G.D. & Martius, C.F.P. (1823) *Nova Acta Physico-medica Academiae Caesareae Leopoldino-Carolinae Naturae Curiosorum Exhibentia Ephemerides sive Observationes Historias et Experimenta* 11: 29.
- Nixon, K.C., Wheeler, Q.D. (1990) An amplification of the phylogenetic species concept. *Cladistics* 6: 211–223.
<http://dx.doi.org/10.1111/j.1096-0031.1990.tb00541.x>
- Padial, J. & De La Riva, I. (2009) Integrative taxonomy reveals cryptic Amazonian species of *Pristimantis* (Anura: Strabomantidae). *Zoological Journal of the Linnean Society* 155: 97–122.
<http://dx.doi.org/10.1111/j.1096-3642.2008.00424.x>
- Padial, J., Miralles, A., De la Riva, I. & Vences, M. (2010) The integrative future of taxonomy. *Frontiers in Zoology* 7: 16.
<http://dx.doi.org/10.1186/1742-9994-7-16>
- R Development Core Team (2014) R: A Language and Environment for Statistical Computing. R Foundation for Statistical Computing. Available form: <http://www.R-project.org>.
- Rieseberg, L.H., Church, S.A. & Morjan, C.L. (2004) Integration of populations and differentiation of species. *New Phytologist* 161: 59–69.
<http://dx.doi.org/10.1046/j.1469-8137.2003.00933.x>
- Sáez, A.G. & Lozano, E. (2005) Body doubles. *Nature* 433:111.
<http://dx.doi.org/10.1038/433111a>
- Salzman, S., Driscoll, H.E., Renner, T., André, T., Shen, S. & Specht, C.D. (2015) Spiraling into history: A molecular phylogeny and investigation of biogeographic origins and floral evolution for the genus *Costus*. *Systematic Botany* 40: 104–115.
<http://dx.doi.org/10.1600/036364415X686404>
- Schumann, K. (1892) *Botanische Jahrbücher für Systematik, Pflanzengeschichte und Pflanzengeographie* 15: 422.
<http://dx.doi.org/10.1080/10635150701772563>
- Shaffer, H.B. & Thomson, R.C. (2007) Delimiting Species in Recent Radiations. *Systematic Biology* 56: 896–906.
- Shaw, J., Lickey, E.B., Schilling, E.E. & Small, R.L. (2007) Comparison of whole chloroplast genome sequences to choose noncoding regions for phylogenetic studies in Angiosperms: the tortoise and the hare III. *American Journal of Botany* 94: 275–288.
<http://dx.doi.org/10.3732/ajb.94.3.275>
- Specht, C.D., Kress, W.J., Stevenson, D.W. & DeSalle, R. (2001) A Molecular Phylogeny of Costaceae (Zingiberales). *Molecular Phylogenetics and Evolution* 21: 333–345.
<http://dx.doi.org/10.1006/mpve.2001.1029>
- Specht, C.D. & Stevenson, D.W. (2006) A new phylogeny-based generic classification of Costaceae (Zingiberales). *Taxon* 55:153–163.
<http://dx.doi.org/10.2307/25065537>
- Specht, C.D. (2006) Systematics and Evolution of the Tropical Monocot Family Costaceae (Zingiberales): A Multiple Dataset Approach. *Systematic Botany* 31: 89–106.
<http://dx.doi.org/10.1600/036364406775971840>
- Taberlet, P., Gielly, L., Pautou, G. & Bouvet, J. (1991) Universal primers for amplification of three non-coding regions of chloroplast DNA. *Plant Molecular Biology* 17: 1105–1109.
<http://dx.doi.org/10.1007/BF00037152>

- Tetsana, N., Pedersen, H.Æ. & Sridith, K. (2014) Character intercorrelation and the potential role of phenotypic plasticity in orchids: a case study of the epiphyte *Liparis resupinata*. *Plant Systematics and Evolution* 300: 517–526
<http://dx.doi.org/10.1007/s00606-013-0900-0>
- Thiers, B. (2014) *Index Herbariorum*: A global directory of public herbaria and associated staff. *New York Botanical Garden's Virtual Herbarium*. Available from: <http://sweetgum.nybg.org/ih/>.
- Walsh, P. (2000) Sample size for the diagnosis of conservation units. *Conservation Biology* 14: 1533–1535.
<http://dx.doi.org/10.1046/j.1523-1739.2000.98149.x>
- Wendt, T., da Cruz, D.D., Demuner, V.G., Guilherme, F.A.G. & Boudet-Fernandes, H. (2011) An evaluation of the species boundaries of two putative taxonomic entities of *Euterpe* (Arecaceae) based on reproductive and morphological features. *Flora* 206: 144–150.
<http://dx.doi.org/10.1016/j.flora.2010.03.002>
- White, T.J., Bruns, T., Lee, S. & Taylor, J. (1990) Amplification and direct sequencing of fungal ribosomal RNA genes for phylogenetics. In: *PCR Protocols: A Guide to Methods and Applications*. Academic Press, San Diego.
<http://dx.doi.org/10.1016/B978-0-12-372180-8.50042-1>

Supplementary Information: Measured specimens

Chamaecostus acaulis. **BRAZIL**. ACRE. Porto Acre: *Lowrie* 526 (INPA, MG, NY); Rio Branco: *Albuquerque* 1389 (NY), *Cid* 2806 (INPA, NY), *Ehrich* 8 (NY), *Lowrie* 445 (INPA, R); Santa Rosa: *Daly* 11103 (NY); Sena Madureira: *Daly* 7855 (NY), *Daly* 7860 (NY), *Prance* 7623 (MO), *Ramos* 682 (INPA); Tarauacá: *Prance* 7416 (INPA, NY); Xapuri: *Alves* 2365 (NY), *Daly* 7195 (MO, NY). GOIÁS. Cabeceiras: *Irwin* 10358 (MO); Caipônia: *Prance* 59644 (A, NY); Cidade de Goiás: *Kirkbridge* 3399 (UB); Santa Rita do Araguaia: *Rocha* sn (UB). MATO GROSSO. Alta Floresta: *André* 804 (RB), *Richter* 39 (RB); Aripuanã: *Berg* 18519 (INPA, NY); Cuiabá: *Andersson* 1623 (A, UB); Barra do Garças: *Philcox* 4000 (UB); Palmeiras: *Lindman* 2485 (A); Poconé: *Maciel* 144 (INPA); Santa Teresinha: *Oliveira* 3094 (RB), *Thomas* 4384 (INPA, MG). Tangará da Serra: *Silva* 498 (TANG). MINAS GERAIS. Ituitaba: *Macedo* 1316 (RB), *Macedo* 1993 (MO); Uberlândia: *Arantes* 1124 (HUFU), *Barbosa* 281 (HUFU). PARÁ. Altamira: *Balée* 1996 (NY), *Dias* 1109 (MG), *Ferreira* 1068 (NY), *Lima* 6020 (RB), *Nascimento* 1177 (NY), *Souza* 1068 (MG), *Souza* 1177 (MG); Conceição do Araguaia: *Plowman* 8448 (A, INPA, MG, MO, NY); Nova Canaã dos Carajás: *Lobato* 2596 (MG); Redenção: *Cordeiro* 2851 (IAN); Serra do Cachimbo: *Prance* 25218 (MG). RONDÔNIA. Abunã: *Prance* 8338 (INPA, MG, NY); Ariquemes: *Zarucchi* 2677 (A, INPA, MG, MO, NY, R, RB), *Mota* 440 (NY), *Vieira* 440 (MO, R), *Vieira* 443 (INPA); Mutumparaná: *Prance* 8975 (A, INPA, MG, NY, R). **BOLIVIA**. Bela Vista: *Steinbach* 7386 (MO); Beni: *Maas* 8660 (MO, NY, RB), *Surubi* 313 (NY), *Rusby* 1399 (A, NY), *Ledezma* 893 (MO); Ñuflo de Chaves: *Ortiz* 39 (NY); Santa Cruz: *Arroyo* sn (MO), *Carrión* 506 (MO), *Castro* 60 (MO), *Garvizu* 513 (MO), *Guillén* 3035 (MO), *Guillén* 3623 (MO), *Killeen* 7168 (MO), *Mamani* 1091 (MO), *Quevedo* 2473 (MO), *Rodriguez* 572 (MO).

Chamaecostus subsessilis s.str.. **BRAZIL**. BAHIA. Itamaraju: *Mori* 10753 (NY, RB); Jussari: *Belém* 2274 (UB), *Thomas* 11937 (MO), *Thomas* 13401 (MO, NY). DISTRITO FEDERAL. Brasília: *Barroso* 639 (RB), *Heringer* 10751 (IAN, UB), *Irwin* 19440 (NY, RB, UB), *Pereira* 2251 (IBGE, RB), *Pires* 51 (RB), *Pires* 57147 (UB). GOIÁS. Alto Paraíso: *Felfili* 379 (IBGE), *Mendonça* 2898 (IBGE); Alvorada do Norte: *Hatschbach* 39011 (NY, UC); Caldas Novas: *Vieira* 1652 (RB); Catalão *Hatschbach* 55820 (MO); Cocalzinho: *Mendonça* 2206 (IBGE); Corumbá: *Maguire* 57147 (MG, MO); Formosa: *Irwin* 9066 (UB); Goiânia: *Rizzo* 2566 (UFG), *Rizzo* 12305 (UFG); Inhumas: *Rizzo* 2779 (UFG); Luziânia: *Coradin* 7397 (RB); Monte Alegre: *Mendonça* 4512 (RB); Mossamedes: *Forzza* 2500 (RB); Niquelândia: *Cordovil* 106 (RB), *Fonseca* 1245 (IBGE, UFG); Nova Roma: *Forzza* 2541 (RB); São Domingo: *Oliveira* 1117 (IBGE), *Santos* 2367 (RB); Trindade: *Rizzo* 3113 (UFG). MARANHÃO. Tuntum: *Santos* 707 (MG, NY). MINAS GERAIS. Abre Caminho: *Pereira* 59 (RB); Jacinto: *Leitman* 51 (RB); Januária: *Filgueiras* 1950 (IBGE), *Ratter* 2630 (IAN, NY, UB, RB), *Ratter* 6410 (IBGE); Lagoa Santa: *Hoehne* 6206 (R); Minas: *Duarte* sn (RB); Serra do Cipó: *Heringer* 7343 (UB); Unaí: *Brina* sn (RB); Várzea da Palma: *Duarte* 7547 (RB). TOCANTINS. Aurora do Norte: *Pereira* 2009 (IBGE); Lajeado: *Árbocz* 6293 (IBGE); Presidente Kennedy: *Plowman* sn (INPA).

Chamaecostus cuspidatus. **BRAZIL**. BAHIA. Belmonte: *Mattos* 368 (NY), *Mattos* 1804 (NY), *Santos* 828 (RB); Eunápolis: *Santos* 893 (NY, RB), *Mello Filho* 2980 (R); Gandú: *Santos* 1157 (NY, UB); Porto Seguro: *Duarte* 5668 (RB), *Pinheiro* 1747 (RB); Wenceslau Guimarães: *Thomas* 9329 (MO). ESPÍRITO SANTO. Colatina: *Kuhlmann* 6660 (RB); Santa Teresa: *Boone* 984 (MO).