Check for updates



Research Article

A new classification of Cyperaceae (Poales) supported by phylogenomic data

Isabel Larridon^{1,2*} , Alexandre R. Zuntini¹, Étienne Léveillé-Bourret³, Russell L. Barrett⁴, Julian R. Starr⁵, A. Muthama Muasya^{1,6}, Tamara Villaverde⁷, Kenneth Bauters⁸, Grace E. Brewer¹, Jeremy J. Bruhl⁹, Suzana M. Costa¹⁰, Tammy L. Elliott¹¹, Niroshini Epitawalage¹, Marcial Escudero¹², Isabel Fairlie^{1,13}, Paul Goetghebeur², Andrew L. Hipp^{14,15}, Pedro Jiménez-Mejías^{16,17}, Izai A.B. Sabino Kikuchi^{1,18}, Modesto Luceño¹⁹, José Ignacio Márquez-Corro¹⁹, Santiago Martín-Bravo¹⁹, Olivier Maurin¹, Lisa Pokorny^{1,20}, Eric H. Roalson²¹, Ilias Semmouri²², David A. Simpson^{1,23}, Daniel Spalink²⁴, W. Wayt Thomas²⁵, Karen L. Wilson⁴, Martin Xanthos¹, Félix Forest¹, and William J. Baker¹

Received 30 September 2020; Accepted 3 May 2021; Article first published online 8 May 2021

Abstract Cyperaceae (sedges) are the third largest monocot family and are of considerable economic and ecological importance. Sedges represent an ideal model family to study evolutionary biology due to their species richness, global distribution, large discrepancies in lineage diversity, broad range of ecological preferences, and adaptations including multiple origins of C_4 photosynthesis and holocentric chromosomes. Goetghebeur's seminal work on Cyperaceae published in 1998 provided the most recent complete classification at tribal and generic level, based on a morphological study of Cyperaceae inflorescence, spikelet, flower, and embryo characters, plus anatomical and other information. Since then, several family-level molecular phylogenetic studies using Sanger sequence data have been published. Here, more

This is an open access article under the terms of the Creative Commons Attribution License, which permits use, distribution and reproduction in any medium, provided the original work is properly cited.

¹Royal Botanic Gardens, Kew, Richmond, Surrey TW9 3AE, UK

²Systematic and Evolutionary Botany Lab, Department of Biology, Ghent University, K.L. Ledeganckstraat 35, 9000 Ghent, Belgium

³Institut de recherche en biologie végétale (IRBV), Université de Montréal, Montréal, QC H1X 2B2, Canada

⁴National Herbarium of New South Wales, Royal Botanic Gardens, Mrs Macquaries Road, Sydney, Sydney, NSW 2000, Australia

⁵Department of Biology, University of Ottawa, Ottawa, ON K1N 6N5, Canada

⁶Department of Biological Sciences, University of Cape Town, Bolus Herbarium, Private Bag X3, Rondebosch, 7701 Cape Town, South Africa

⁷Universidad de Almería, Carretera Sacramento s/n, La Cañada de San Urbano, Almería 04120, Spain

⁸Botanic Garden Meise, Nieuwelaan 38, Meise 1860, Belgium

⁹Botany, School of Environmental and Rural Science, University of New England, Armidale, NSW 2351, Australia

¹⁰Departamento de Biologia, Universidade Federal de Lavras, Caixa Postal 3037, Lavras, Minas Gerais CEP 37200-000, Brazil

¹¹Department of Botany and Zoology, Faculty of Science, Masaryk University, Kotlarska 2, Brno 611 37, Czech Republic

¹²Departamento de Biología Vegetal y Ecología, Universidad de Sevilla, Carretera Profesor García González s/n, Seville 41012, Spain

¹³Department of Animal and Plant Sciences, Univeristy of Sheffield, Alfred Denny Building, Western Bank, Sheffield S10 2TN, UK

¹⁴The Morton Arboretum, 4100 Illinois Route 53, Lisle, IL 60532-1293, USA

¹⁵The Field Museum, 1400S. Lake Shore Dr., Chicago, IL 60605, USA

¹⁶Departamento de Biología (Botánica), Facultad de Ciencias Biológicas, Universidad Autónoma de Madrid, C/Darwin, 2, 28049 Madrid, Spain

¹⁷Centro de Investigación en Biodiversidad y Cambio Global (CIBC-UAM), Universidad Autónoma de Madrid, 28049 Madrid, Spain

¹⁸Hortus Botanicus Leiden, Universiteit Leiden, PO Box 9500, Leiden 2300 RA, The Netherlands

¹⁹Department of Molecular Biology and Biochemical Engineering, Botany area, Universidad Pablo de Olavide, Carretera de Utrera km 1, 41013 Seville, Spain

²⁰Centre for Plant Biotechnology and Genomics (CBGP, UPM-INIA), Pozuelo de Alarcón, Madrid 28223, Spain

²¹School of Biological Sciences, Washington State University, Pullman, WA 99164-4236, USA

²²Laboratory of Environmental Toxicology and Aquatic Ecology, Faculty of Bioscience Engineering, Ghent University, 9000 Ghent, Belgium

²³Department of Botany, School of Natural Sciences, Trinity College Dublin, Dublin 2, Ireland

²⁴Department of Ecology and Conservation Biology, Texas A&M University, College Station, TX 77843, USA

²⁵The New York Botanical Garden, 2900 Southern Blvd, Bronx, NY 10458, USA

^{*}Author for Correspondence. E-mail: i.larridon@kew.org

than 20 years after the last comprehensive classification of the family, we present the first family-wide phylogenomic study of Cyperaceae based on targeted sequencing using the Angiosperms353 probe kit sampling 311 accessions. In addition, 62 accessions available from GenBank were mined for overlapping reads and included in the phylogenomic analyses. Informed by this backbone phylogeny, a new classification for the family at the tribal, subtribal, and generic levels is proposed. The majority of previously recognized suprageneric groups are supported, and for the first time, we establish support for tribe Cryptangieae as a clade including the genus Koyamaea. We provide a taxonomic treatment including identification keys and diagnoses for the 2 subfamilies, 24 tribes, and 10 subtribes, and basic information on the 95 genera. The classification includes five new subtribes in tribe Schoeneae: Anthelepidinae, Caustiinae, Gymnoschoeninae, Lepidospermatinae, and Oreobolinae.

Key words: Angiosperms353, classification, Cyperaceae, phylogenomics, Poales, systematics, targeted sequencing, taxonomy.

1 Introduction

Cyperaceae (sedges) are the third largest monocot family (>5600 spp.; Govaerts et al., 2020) and are of considerable economic and ecological importance (Simpson & Inglis, 2001; Spalink et al., 2016a, 2018). Cyperaceae are an ideal model family to study evolutionary biology due to their species richness, global distribution, large discrepancies in lineage diversity (Escudero & Hipp, 2013), broad range of ecological preferences and diverse phenotypes (Naczi & Ford, 2008), multiple origins of C₄ photosynthesis (Besnard et al., 2009), and the presence of holocentric chromosomes (Márquez-Corro et al., 2019, 2021). The family is species-rich in the tropics where it exhibits high generic diversity and a remarkable species richness in the genus Cyperus L. with >960 spp. (Govaerts et al., 2020). High diversity in temperate regions is mostly due to the megadiverse genus Carex L. with >2000 spp. (Govaerts et al., 2020). The history of the family goes back to the early Cenozoic, as supported by a reliable fossil record dating back to the Paleocene (Smith et al., 2009; Spalink et al., 2016b) and evidence of large genera already established by the end of the Eocene (Jiménez-Mejías et al., 2016a), with a probable origin in South America (Spalink et al., 2016b).

Phenotypic diversity in Cyperaceae is represented among others by a wide range of growth forms, from tiny ephemerals less than 3 cm high such as Isolepis inconspicua (Levyns) J.Raynal to climbing herbs that may measure to more than 12 m long such as Scleria boivinii Steud. (Fig. 1A), and to the long-lived dwarf-tree-like Microdracoides squamosa Hua (Fig. 1B) and the similar looking species of the New Caledonian endemic genus Chamaedendron (Kük.) Larridon (Larridon et al., 2018a). It is also ecologically diverse and occurs in a wide variety of habitats ranging from truly aquatic plants in rivers and lakes to annuals and perennials in fireprone grasslands, forests and high-elevation vegetations (Simpson et al., 2011; e.g., Carex moorcroftii Falc. ex Boott has been found up to elevations of 5700 m; Dai et al., 2010), deserts, and coastal sands (Väre & Kukkonen, 2005; Reznicek et al., 2021). There are even species with a tendency for epiphytism (i.e., Coleochloa domensis Muasya & D.A.Simpson; Muasya et al., 2010). In some ecosystems, such as wetlands and inselbergs, Cyperaceae are true ecosystem builders providing habitats for many other species (e.g., mats of Afrotrilepis pilosa (Boeckeler) J.Raynal; Fig. 1C). In temperate Australia, the genera Ammothryon R.L.Barrett, K.L.Wilson & J.J.Bruhl, Caustis R.Br., Gahnia J.R.Forst. & G.Forst., Lepidosperma Labill., Netrostylis R.L.Barrett, J.J.Bruhl & K.L.Wilson, Schoenus L., and Tricostularia Nees ex Lehm. can be dominant components of the understorey in numerous dryland vegetation communities, with high ecological significance (Barrett, 2013; Barrett et al., 2021a, 2021b). Furthermore, the family exhibits high diversity in reproductive and seed dispersal structures, and use of these morphological characters to define taxon limits has resulted in much of the observed taxonomic complexity of, for example, Cyperus (Fig. 1D; Larridon et al., 2011a, 2013) and Carex (Global Carex Group, 2015; Jiménez-Mejías et al. 2016b), and the wide range of dispersal vectors known for the family (Goetghebeur, 1998; Larridon et al., 2011b, 2021a), for example, birds in species of Gahnia (Fig. 1E), and ants in selected species of Carex (Handel, 1976; Vellend et al., 2000) and Lepidosperma (Barrett, 2013). Long-distance dispersal has led to unusual disjunct distributions in genera such as Morelotia Gaudich. (Barrett et al., 2021c) and Tetraria P.Beauv. (Larridon et al., 2018b). Although predominantly wind-pollinated, there are many transitions from wind to insect pollination in Cyperaceae (Fig. 1F; Wragg & Johnson, 2011; Yano et al., 2015; Costa et al., 2018a; Villa-Machío et al., 2020). The sedge family is also physiologically diverse, with species using C_3 , C_4 , or C_3-C_4 intermediate photosynthesis (Soros & Bruhl, 2000; Bruhl & Wilson, 2007; Besnard et al., 2009). At least six independent origins of C₄ photosynthesis have been recorded in tribe Abildgaardieae and in Cyperus, Eleocharis R.Br., and Rhynchospora Vahl (Bruhl & Wilson, 2007; Besnard et al., 2009; Larridon et al., 2011a, 2021b). Cyperaceae are of worldwide economic significance with about 10% of species used by humans, particularly in the tropics (Simpson & Inglis, 2001). Globally important crop species include Cyperus esculentus L. (tiger nuts) as snacks or for the production of "horchata de chufa" (tiger nut milk), and Eleocharis dulcis (Burm.f.) Trin. ex Hensch. (water chestnuts) as a popular ingredient in Asian dishes. Several species are notorious weeds (Bryson & Carter, 2008), such as Cyperus aromaticus (Ridl.) Mattf. & Kük. (Vitelli et al., 2010; Biosecurity Queensland, 2018), C. esculentus (Dodet et al., 2008; Rogers et al., 2008), and C. rotundus L. (Goetghebeur, 1998; Rogers et al., 2008; ISSG, 2019). Cyperaceae also have a remarkable and often overlooked importance in human history (e.g., totora or Schoenoplectus californicus (C.A.Mey.) Soják, papyrus or Cyperus papyrus L.; Banack et al., 2004; Bagnall, 2009) and in human evolution as a food source



Fig. 1. Morphological and ecological diversity of Cyperaceae. **A,** *Scleria boivinii* Steud. **B,** *Microdracoides squamosa* Hua. **C,** Afrotrilepis pilosa (Boeckeler) J.Raynal. **D,** *Carex lechleriana* (Steud.) J.R.Starr formerly placed in the segregate genus *Uncinia* Pers. **E,** *Gahnia tristis* Nees. **F,** *Rhynchospora alba* (L.) Vahl. Photos **A** by Javier Galán Díaz; **B** by Charlotte Couch; **C** by Xander van der Burgt; **D** by Modesto Luceño; **E** by Russell Barrett; **F** by Juan Carlos Zamora.

(Sponheimer et al., 2005, 2013; Dominy, 2012; Wynn et al., 2013; Cerling et al., 2013a, 2013b).

Cyperaceae are a well-supported monophyletic family within the monocot order Poales, sister to Juncaceae (Simpson et al., 2007; Muasya et al., 2009a; Givnish et al., 2010; Escudero & Hipp, 2013; Jung & Choi, 2013; Bouchenack-Khelladi et al., 2014; Ruhfel et al., 2014; Spalink et al., 2016b; Semmouri et al., 2019). Goetghebeur's (1998) seminal work on Cyperaceae still provides the most recent complete classification at tribal and generic level, based on a morphological study of Cyperaceae inflorescence, spikelet, flower, and embryo characters, plus anatomical and other information (Goetghebeur, 1986). Since then, a range of molecular phylogenetic studies using Sanger sequence data has been published at the family level (Simpson et al., 2007; Muasya et al., 2009a; Escudero & Hipp, 2013; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Spalink et al., 2016b; Semmouri et al., 2019; Larridon et al., 2021b). Two main clades have been recovered within Cyperaceae, corresponding to subfamilies Mapanioideae and Cyperoideae. Mapanioideae mainly includes broad-leaved tropical forest understory herbs, whereas Cyperoideae is much more diverse in terms of species richness, morphology, and ecology. Other studies have focused on particular tribes or genera, including Carex (e.g., Global Carex Group, 2015; Jiménez-Mejías et al., 2016c; Martín-Bravo et al., 2019), Cyperus (e.g., Larridon et al., 2011a, 2013; Bauters et al., 2014), Abildgaardieae (e.g., Reutemann et al., 2018; Roalson et al., 2019; Muasya et al., 2020), Schoeneae (e.g., Viljoen et al., 2013; Larridon et al., 2018a), or the Scirpo–Caricoid Clade (e.g., Léveillé-Bourret et al., 2014, 2015, 2018a, 2018b; Léveillé-Bourret & Starr, 2019), increasing our understanding of the relationships in Cyperaceae. Despite previous studies, systematic relationships in Cyperaceae at the tribal and generic level are not yet fully resolved because (i) some genera have not yet been sampled and (ii) some relationships are conflicting in different studies.

Molecular phylogenetic studies on Cyperaceae have relied heavily on relatively few loci, such as a selection of plastid markers and the nuclear markers ITS and ETS (e.g., Semmouri et al., 2019). However, phylogeny estimation is more accurate when conducted with tens to hundreds of nuclear loci, because larger numbers of informative characters help to resolve short branches, and historical processes such as deep coalescence can be taken into account (Johnson et al., 2019). Hence, reduced-representation sequencing methods have been developed to sample hundreds of nuclear, orthologous single-copy genes for plant phylogenetic studies (Kadlec et al., 2017; Couvreur et al., 2019; Johnson et al., 2019; Villaverde et al., 2018, 2020), allowing users to yield data sets of a larger scale for phylogenetics without the bioinformatic challenges and

costs associated with whole-genome sequencing. Larridon et al. (2020) provided an overview of earlier high-throughput sequencing studies on Cyperaceae, whereas more recent studies relying on genomic data already show alternative phylogenetic structure in certain sedge groups not previously recovered using Sanger sequencing (Léveillé-Bourret et al., 2018c; Larridon et al., 2020; Starr et al., 2021; Villaverde et al., 2020, 2021).

The aim of this study is to resolve the high-level relationships in Cyperaceae and to test the monophyly of the tribes and genera as currently accepted to generate a new classification from subfamily to generic level. We hypothesize that using genome-scale data and an in-depth sampling will provide significantly more phylogenetic information to resolve the topology of the Cyperaceae Tree of Life. Equally, we postulate that the high-throughput technique-targeted sequencing will enable sequencing historical herbarium specimens with poor DNA quality (Brewer et al., 2019), allowing us to place previously unplaced genera in the family phylogeny for the first time.

2 Material and Methods

2.1 Taxon sampling

A total of 361 accessions of Cyperaceae were sampled, along with 21 accessions representing other families in order Poales as outgroups (Table S1). The sampling includes nearly all currently accepted genera of Cyperaceae (Govaerts et al., 2020). Three monotypic genera were not sampled, and have never been successfully sequenced using Sanger methods: Nelmesia Van der Veken and Trichoschoenus J.Raynal, which are only known from their type collections, and Rhynchocladium T.Koyama. Costa et al. (2021a) recently changed the generic circumscription in tribe Cryptangieae, re-establishing the monotypic genus Didymiandrum Gilly, whereas Barrett et al. (2021b) recently described a new monotypic genus Ammothryon. These two genera were not sampled. Lab work for samples of three additional monotypic genera, Blysmopsis Oteng-Yeb., Capeobolus Browning, and Khaosokia D.A.Simpson, and the small genus Blysmus Panz. ex Schult. did not provide data of sufficient quality. These genera have been previously successfully placed in the Cyperaceae Tree of Life (e.g., Léveillé-Bourret et al., 2014, 2018c; Larridon et al., 2018a; Semmouri et al., 2019). In total, 311 of the 382 accessions were sequenced after enrichment with the Angiosperms353 probes. In addition, 36 accessions enriched with the Angiosperms I kit for Anchored Phylogenomics (Léveillé-Bourret et al., 2018c), including Khaosokia caricoides D.A.Simpson, were mined for reads overlapping with the data generated using the Angiosperms353 probes, as were 6 accessions enriched with Cyperaceae-specific probes (Villaverde et al., 2020), and 20 transcriptomes available on GenBank (Table S1). Angiosperms353 data for most accessions were newly generated for this study, following the protocol established by Baker et al. (2021). In addition, some data were obtained from recent studies (Larridon et al., 2020, 2021c; Starr et al., 2021; Table S1).

2.2 DNA extraction, library preparation, hybridization, and sequencing

The voucher information and treatment of each sample are provided (Table S1). Molecular work for accessions enriched with

the Angiosperms353 probes was carried out at the Sackler Phylogenomics Laboratory, within the Jodrell Laboratory at Royal Botanic Gardens, Kew (Richmond, Surrey, UK). Genomic DNA was extracted from leaf tissue obtained from herbarium specimens or silica-dried samples, using either a modified CTAB approach (Doyle & Doyle, 1987) or a CTAB protocol, based on Beck et al. (2012), modified for optimal simultaneous extraction of 96 to 192 samples (i.e., one or two plates) from degraded (i.e., herbarium) samples (see Supplementary Data Sheet 1 in Larridon et al., 2020). Lastly, 76 accessions were sourced from the Kew DNA Bank (http://dnabank.science.kew.org/) (Table S1). The samples extracted using a CTAB approach were purified using Agencourt AMPure XP Bead Clean-up (Beckman Coulter, Indianapolis, IN, USA). All DNA extracts were quantified using a Quantus™ Fluorometer (Promega Corporation, Madison, WI, USA) and then run on a 1% agarose gel to assess the average fragment size. Samples with a very low concentration (not visible on a 1% agarose gel) were assessed on an Agilent Technologies 4200 TapeStation System using Genomic DNA ScreenTape (Santa Clara, CA, USA). DNA extracts with average fragment sizes above 350 bp were sonicated using a Covaris M220 Focusedultrasonicator™ (Covaris, Woburn, MA, USA) following the manufacturer's protocol to obtain an average fragment size of 350 bp. Dual-indexed libraries for Illumina® sequencing were prepared using the DNA NEBNext® Ultra™ II Library Prep Kit and the NEBNext® Multiplex Oligos for Illumina® (Dual Index Primers Set 1 and 2) from New England BioLabs® (Ipswich, MA, USA) following the manufacturer's instructions but at half the recommended volumes. The quality of the libraries was evaluated on the TapeStation using High Sensitivity D1000 ScreenTape and the libraries were quantified using a Quantus Fluorometer. The final average library size including the adapters was c. 500 bp. Afterward, the samples were pooled and enriched with the Angiosperms353 probes (Johnson et al., 2018) following the manufacturer's instructions (myProbes® Manual v4.01, Arbor Biosciences, Ann Arbor, MI, USA). Final products were again run on the TapeStation to assess quality (i.e., average fragment size) so they could be pooled equimolarly for sequencing. After multiplexing library pools, sequencing was performed on an Illumina® MiSeq instrument (San Diego, CA, USA) with v2 (300 cycles at 2×150 bp) or v3 (600 cycles at 2×300 bp) chemistry at Royal Botanic Gardens, Kew (Richmond, Surrey, UK), or on an Illumina® HiSeq (San Diego, CA, USA) at either Macrogen (Seoul, South Korea) or GENEWIZ® (Leipzig, Germany), producing 2 × 150 bp long reads. Raw reads for all accessions are available from the NCBI GenBank Sequence Read Archive (SRA) under Bioproject numbers PRJNA553989 (http://www.ncbi.nlm.nih.gov/ bioproject/PRJNA553989), PRJNA649146 (http://www.ncbi.nlm. nih.gov/bioproject/PRJNA649146), PRJNA668802 (http://www. ncbi.nlm.nih.gov/bioproject/PRJNA668802), and PRJNA669 051 (http://www.ncbi.nlm.nih.gov/bioproject/PRJNA669051), and from the European Nucleotide Archive (ENA) under EMBL Project number PRJEB35285 (https://www.ebi.ac.uk/ena/ browser/view/PRJEB35285).

2.3 Read processing, assembly, and phylogenomic analyses Bioinformatics settings follow Larridon et al. (2021c). Raw reads were trimmed to remove adapter sequences and portions of low quality with Trimmomatic v.0.39 (Bolger et al., 2014) using the following settings: LEADING:30 TRAILING:30 SLIDINGWINDOW:4:2:30 MINLEN:36. HybPiper

v.1.3.1 (Johnson et al., 2016) was used to process the quality-checked, trimmed reads, with default settings except for minimum coverage set to 4x. Paired and unpaired reads from all accessions were mapped to targets with BLASTx (Altschul et al., 1990) using the Angiosperms353 target loci amino acid (AA) sequences (see Supplementary Data Sheet 3 in Larridon et al., 2020). Mapped reads were then assembled into contigs with SPAdes v.3.13.1 (Bankevich et al., 2012). Subsequently, exonerate v.2.2 (Slater & Birney, 2005) was used to align the assembled contigs to their associated target sequence and remove intronic regions (exons data set). HybPiper flags potential paralogs when multiple contigs are discovered mapping well to a single reference sequence. As few random paralog warnings were raised, no sequence was excluded.

Phylogenomic analyses were executed in two rounds (following Zuntini et al., 2021) to improve the inference results. In the first round, all exon sequences with at least 50 bp were recovered and then aligned with MAFFT v.7 (Katoh and Standley, 2013) with the "localpair max iterations 1000" option; sites with more than 30% missing data were removed using Phyutility (Smith & O'Meara, 2012), after which IQ-TREE v.2.1.0 (Minh et al., 2020) was run per gene, followed by TreeShrink (Mai & Mirarab, 2018) with threshold set to 0.05. After this, a quality check was performed to see how many times each accession appeared in each gene tree. Finally, ASTRAL-III v.5.5.11 (Zhang et al., 2018) was run after collapsing branches ≤10% bootstrap (BS) support using Newick Utilities (Junier & Zdobnov, 2010). This round provided the preliminary result. In the second round, again all sequences with at least 50 bp were recovered, those flagged by TreeShrink were removed, and then aligned with MAFFT, after which we generated summary stats in AMAS (Borowiec, 2016). Short alignments (<100 bp) were removed.

For the coalescent ASTRAL analysis, IQ-TREE was run per gene, after which ASTRAL was executed after collapsing branches ≤10% BS support to provide the final result. Tree images were plotted in R (R Core Team, 2020), using the packages ape (Paradis & Schliep, 2018), ggimage (Yu, 2019a), ggtree (Yu et al., 2017), treeio (Yu, 2019b), and their dependencies.

For the concatenated IQ-TREE analysis, the individual gene alignments were concatenated in AMAS, and IQ-TREE was run with mode set to "MFP + MERGE" and 10 000 replicates of ultrafast bootstrap replications (Hoang et al., 2018) to generate the final result. We also calculated two measures of genealogical concordance in our data set, the gene concordance factor (gCF) and the site concordance factor (sCF), using the options "-gcf" and "-scf" in IQ-TREE. Trees were plotted in FigTree v.1.4.4 (https://github.com/rambaut/figtree/releases).

3 Results

3.1 Capture success and data quality

The success of sequence recovery was variable, with an average of 177 genes per sample (above 25% of target size) and 41% of the total potential target (260 802 bp), as indicated in Johnson et al. (2019). The recovery of samples hybridized with other kits was significantly lower: for

samples hybridized with the Angiosperms I kit for Anchored Phylogenomics, the recovery was 65 genes, on average, above 25% of target length and 17% of total potential length, whereas samples hybridized with Cyperaceae-specific probes yielded, on average, 45 genes and 18% of potential length (Table S2, Fig. S1).

The gene alignments length varied between 123 and 2439 bp (average 623 bp), harboring between 16 and 368 samples (average 236). In total, gene alignments present 196 726 characters, with 30.6% missing data (3.4%–57.6%). The proportion of variable sites and parsiomony-informative sites was, respectively, 0.79 and 0.64 (Table S3).

3.2 Phylogenetic relationships

The tree resulting from the coalescent ASTRAL analysis is shown in Fig. 2, and the tree resulting from the concatenated IQ-TREE analysis is shown in Fig. S2. As relationships are very congruent, below we will discuss the relationships as shown in Fig. 2.

Cyperaceae are retrieved as a monophyletic family sister to Juncaceae with strong support (Fig. 2). Within Cyperaceae, 16 main clades are recovered (Fig. 2). Clade 1 represents subfamily Mapanioideae and includes two sister clades representing the tribes Chrysitricheae and Hypolytreae. Clade 2 represents tribe Trilepideae. Clade 3 represents the speciespoor tribe Cladieae. Clade 4 is the Bisboeckelereae-Sclerieae Clade, which includes two subclades representing tribe Bisboeckelereae and tribe Sclerieae, respectively. Clade 5 represents the species-poor tribe Carpheae. Clade 6 is formed of the genus Koyamaea W.W.Thomas & G.Davidse sister to a clade representing tribe Cryptangieae. Clade 7 represents the diverse and species-rich tribe Schoeneae. Tribe Schoeneae includes a range of well-supported clades; however, the nodes in the backbone of the tribe are not all well supported. Also, its position in the backbone of the family is not well supported (LPP = 0.76). Clade 8 consists of tribe Rhynchosporeae. Clade 9 or the Scirpo-Caricoid Clade (SCC Clade) includes a range of species-poor and species-rich lineages: Dulichieae, Khaosokieae, Calliscirpeae, Scirpeae, Trichophoreae, Sumatroscirpeae, and Cariceae. Each tribe is well supported as a monophyletic group as are the backbone nodes. Clade 10 is the Abildgaardieae-Eleocharideae Clade, which falls apart into two sister clades representing the tribes Abildgaardieae and Eleocharideae. Clades 11-14 are often referred to as the Fuireneae s.l. grade, with Clade 11 representing tribe Bolboschoeneae, Clade 12 tribe Fuireneae s.s., Clade 13 tribe Schoenoplecteae, and Clade 14 tribe Pseudoschoeneae. Each tribe is well supported as a monophyletic group and is placed with high support in the backbone of the family. Clades 15 and 16 represent the two main clades of tribe Cypereae, that is, the Ficinia Clade and the Cyperus Clade.

4 Discussion

4.1 Family Cyperaceae

In our results, Cyperaceae are confirmed as a monophyletic family within the monocot order Poales, sister to Juncaceae (Figs. 2, 3). The relationships inferred within Cyperaceae are mostly congruent with those of previous analyses

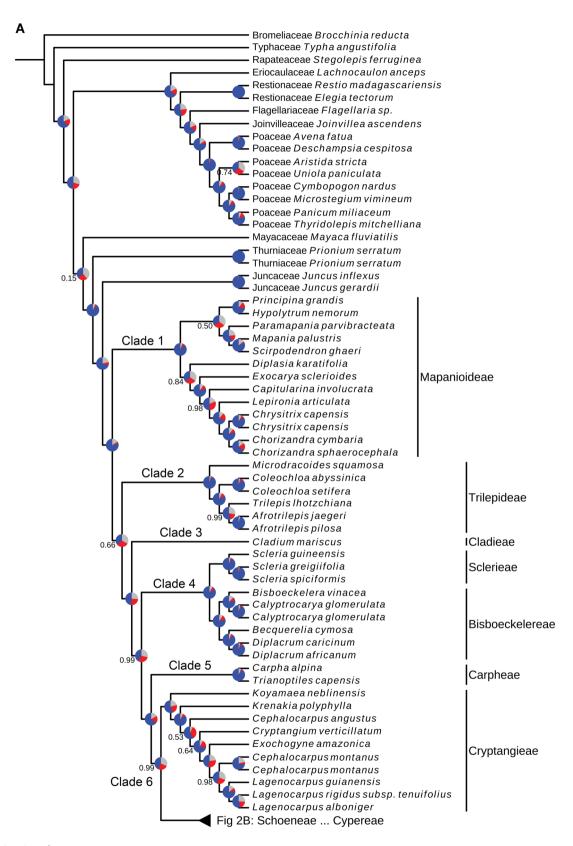


Fig. 2. Continued

(Simpson et al., 2007; Muasya et al., 2009a; Escudero & Hipp, 2013; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Spalink et al., 2016b; Semmouri et al., 2019), with some exceptions. Table S4 provides an overview of the main published classifications of the Cyperaceae and the classification proposed in this study, clearly indicating which changes occurred as more data became available. Table 1 provides an overview of the proposed classification.

Most previous molecular studies, which were largely based on chloroplast sequence data, recognized two subfamilies in Cyperaceae (Muasya et al., 2009a; Escudero & Hipp, 2013; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Spalink et al., 2016b; Semmouri et al., 2019). The targeted sequencing data (Fig. 2) confirm the established topology with subfamily Mapanioideae sister to subfamily Cyperoideae. The main morphological arguments to recognize two subfamilies in Cyperaceae relate to the differences in the morphology of the basic units of the inflorescence. In Cyperoideae, inflorescences are composed of one to many spikelets, each consisting of a rachilla bearing few to many glumes that may or may not subtend a flower (e.g., Goetghebeur, 1998). In contrast, the inflorescence units of Mapanioideae are frequently referred to as spicoids (e.g., Kukkonen, 1984; Simpson, 1992; Simpson et al., 2003; Beentje, 2016 and the preferred term here) or pseudospikelets (e.g., Eiten, 1976; Dai et al., 2010), and comprise 1–13(–100) scales. The homology of these units is still unclear. Many authors consider them to be a much-reduced spikelet (the basic inflorescence unit found in most other Cyperaceae; Dahlgren et al., 1985; Simpson, 1992; Vrijdaghs et al., 2006; Prychid & Bruhl, 2013), whereas others view them as a flower in which the regular trimerous structure of the cyperaceous flower has been disturbed (Goetghebeur, 1986, 1998).

Most previous studies retrieved tribe Trilepideae as sister to all remaining Cyperoideae (Simpson et al., 2007; Muasya et al., 2009a; Escudero & Hipp, 2013; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Spalink et al., 2016b; Semmouri et al., 2019). This relationship is here confirmed (Figs. 2, 3). Otherwise, little congruence can be found concerning the relationships between the early diverging lineages of Cyperoideae in the literature. In Simpson et al. (2007; using only the plastid marker *rbcL*), a Bisboeckelereae–Sclerieae Clade is the second branching clade in Cyperoideae before Cladium P.Browne. In Muasya et al. (2009a; using the plastid markers *rbcL* and *trnL-F*), Cladium branches off before Bisboeckelereae and Sclerieae that form a polytomy with the remainder of Cyperoideae; in Jung & Choi (2013; using the plastid markers *rbcL* and *trnL-F* plus one nuclear marker ITS),

both clades form a polytomy with the remainder of the Cyperoideae (cf. the Maximum Parsimony results of Semmouri et al., 2019; using five plastid markers and two nuclear markers ETS and ITS). However, in Hinchliff & Roalson (2013; supermatrix approach with scaffold based on two chloroplast markers ndhF and rbcL), Cladium is strongly supported as sister to Schoeneae, with the Bisboeckelereae-Sclerieae Clade retrieved as a separate lineage. In the Maximum Likelihood and Bayesian Inference results of Semmouri et al. (2019), and in the results of Spalink et al. (2016b; using the plastid markers matK, ndhF, rbcL, and trnL-F), Cladieae branches off right after Trilepidae, followed by a Bisboeckelereae-Sclerieae Clade sister to the rest of Cyperoideae. In our results (Fig. 2A), after subfamily Mapanioideae (Clade 1), tribe Trilepideae (Clade 2), tribe Cladieae (Clade 3), the Bisboeckelereae-Sclerieae Clade (Clade 4), tribe Carpheae (Clade 5), and tribe Cryptangieae (Clade 6), followed by the rest of subfamily Cyperoideae, branch off subsequently.

The topology of the family (Figs. 2, 3) raises interesting evolutionary and developmental questions in that Clades 1-6 are largely composed of tribes that are characterized by species having unisexual flowers (with the exception of Cladieae and Carpheae), in contrast to the remaining Cyperaceae tribes that are largely characterized by having bisexual flowers (with the exception of the tribes Khaosokieae and Cariceae in the Scirpo-Caricoid Clade). These clades are also characterized by having embryo types that were placed close to the ancestral Juncus-type embryo in the semophylesis (evolutionary sequence) of the embryo types according to Goetghebeur (1986; see also fig. 3 of Semmouri et al., 2019). Goetghebeur (1998) placed most of these tribes in two subfamilies: Chrysitricheae and Hypolytreae in subfamily Mapanioideae, and Trilepideae, Bisboeck-Sclerieae, and Cryptangieae in subfamily elereae, Sclerioideae. Tribes Carpheae and Cladieae were only recently recognized (Semmouri et al., 2019) and were previously treated as part of tribe Schoeneae (e.g., Goetghebeur, 1998). Simpson et al. (2007) showed that subfamily Sclerioideae was not monophyletic and suggested maintaining only two subfamilies in Cyperaceae, that is, Mapanoideae and Cyperoideae.

4.2 Subfamily Mapanioideae

On the basis of pollen data, Simpson et al. (2003) supported the recognition of the two tribes in subfamily Mapanioideae, that is, Chrysitricheae (Fig. 5D) and Hypolytreae (Fig. 4B). Most Cyperaceae, including tribe Chrysitricheae, have

Fig. 2. Phylogenetic reconstruction of the relationships in Cyperaceae based on analysis of the exons data set. Coalescent ASTRAL analysis. Values by nodes represent local posterior probabilities (LPPs) and pie charts at nodes correspond to quartet support with blue: agreeing genes, red: disagreeing genes, and gray: uninformative genes. **A,** showing Poales outgroups and Cyperaceae Clades 1–6 (Mapanioideae, Trilepideae, Cladieae, the Bisboeckelereae–Scleria Clade, Carpheae, and Cryptangieae). **B,** Clades 7 and 8 (Schoeneae and Rhynchosporeae). **C,** Clade 9 or the Scirpo-Caricoide Clade (Dulichieae, Khaosokieae, Calliscirpeae, Scirpeae, Trichophoreae, Sumatroscirpeae, and Cariceae). **D,** Clade 10 or the Abildgaardieae–Eleocharideae Clade (Abildgaardieae and Eleocharideae). **E,** Clades 11–14 representing the Fuireneae s.l. Grade (Bolboschoeneae, Fuireneae, Schoenoplecteae, and Pseudoschoeneae). **F,** Clades 15 and 16 or tribe Cypereae (subtribes Ficiniinae and Cyperinae).

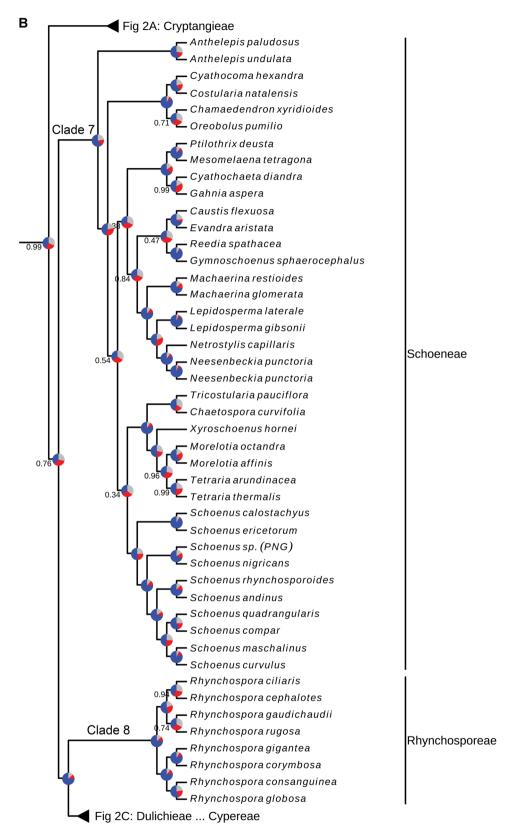


Fig. 2. Continued

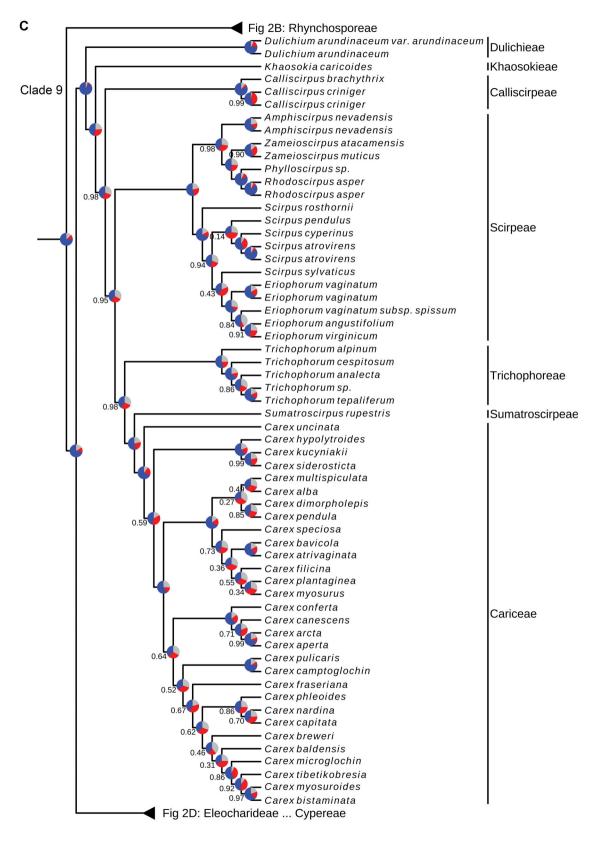


Fig. 2. Continued

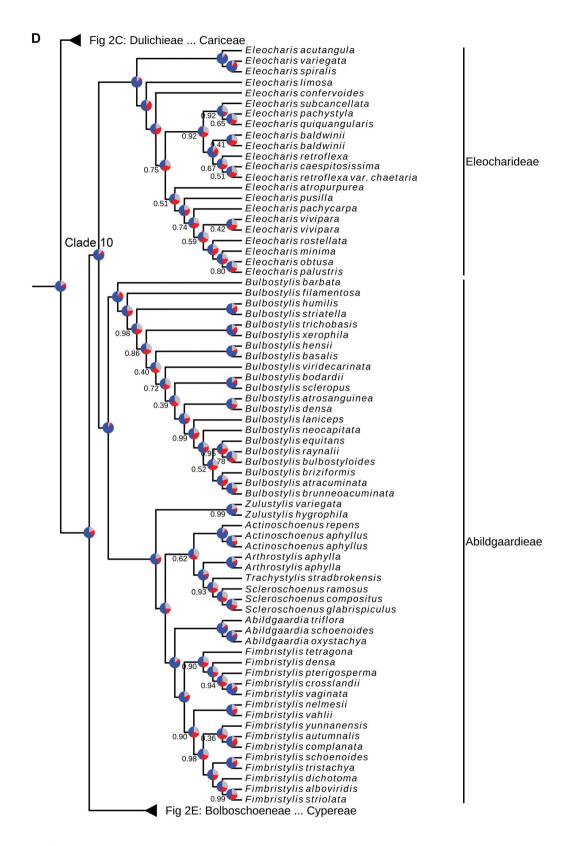


Fig. 2. Continued

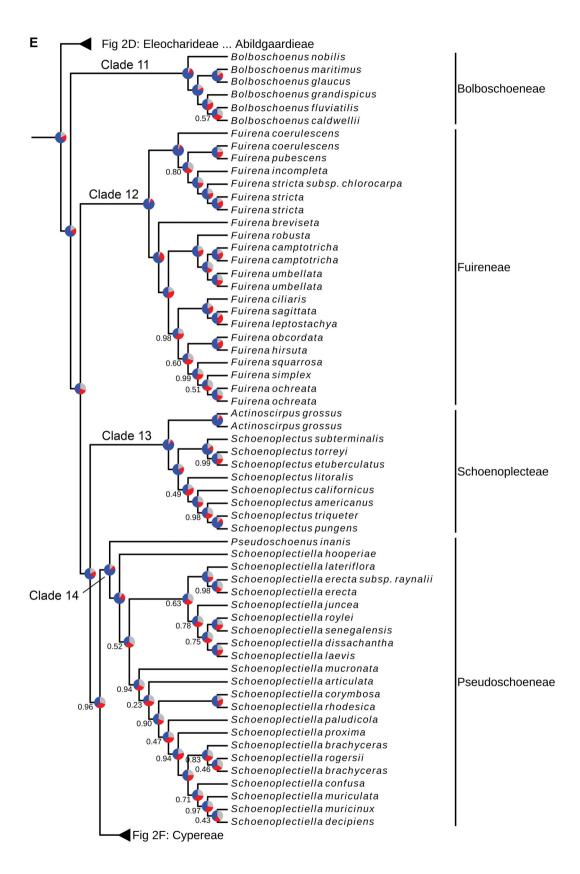


Fig. 2. Continued

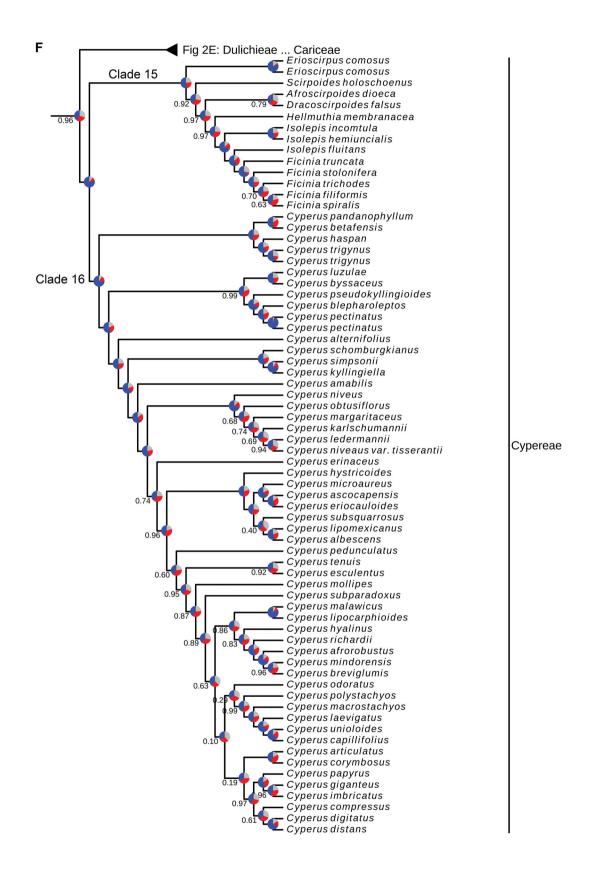


Fig. 2. Continued

thin-walled, pyriform, pseudomonad pollen, whereas Hypolytreae (forest or forest-margin dwellers, where wind pollination is less or not effective) have thick-walled, spheroidal, "Mapania-type" pollen that is coated with lipids, supporting earlier studies, especially Lorougnon (1973), which suggest that Hypolytreae use animal vectors for pollination (Simpson et al., 2003; Nagels et al., 2009). Simpson et al. (2003) indicated that younger, developmental stages of "Mapania-type" pollen were not available for their study and that pollen ontogeny could not be examined. However, Coan et al. (2010) showed that several Hypolytrum species have pseudomonads, suggesting that "Mapania-type" pollen in general is pseudomonad.

In most molecular studies, the circumscription of Chrysitricheae and Hypolytreae and the relationships between and within these tribes are not well resolved or have been conflicting (Simpson et al., 2003, 2007; Muasya et al., 2009a; Hinchliff & Roalson, 2013; Semmouri et al., 2019). A case in point is the inconsistent position of Diplasia karatifolia Rich. Simpson et al. (2003) and Muasya et al. (2009a) placed Diplasia Pers. within tribe Hypolytreae, whereas Semmouri et al. (2019) placed it in a nested position within tribe Chrysitricheae, and its relationship was unresolved in Hinchcliff & Roalson (2013). In Spalink et al. (2016b), Diplasia was positioned as a separate lineage sister to the often retrieved Hypolytreae-Chrysitricheae Clade. Our results recover the tribes Chrysitricheae and Hypolytreae as monophyletic, with Diplasia as sister to the rest of tribe Chrysitricheae with moderate support (LPP = 0.84; Fig. 2A).

A recent molecular phylogenetic study showed that the formerly recognized monotypic genus *Principina* Uittien in nested within *Hypolytrum* (A. Mesterházy et al., unpublished data). In our results, the sample of *Principina* is retrieved as sister to the single included accession of *Hypolytrum*, confirming a close relationship. In Section 5, we follow A. Mesterházy et al. (unpublished data) and relegate *Principina* to synonymy.

4.3 Subfamily Cyperoideae 4.3.1 Tribe Trilepideae

In earlier studies (e.g., Simpson et al., 2007; Muasya et al., 2009a; Hinchliff & Roalson, 2013; Semmouri et al., 2019) and in this study (Figs. 2, 3), tribe Trilepideae (Fig. 5C) is retrieved as the first branching clade sister to the remaining lineages of subfamily Cyperoideae. Trilepideae includes four genera: Afrotrilepis (Gilly) J.Raynal, Coleochloa Gilly, Microdracoides Hua, and Trilepis Nees. In our results (Fig. 2A), within Trilepideae, Microdracoides appears sister to the other three genera, with Coleochloa sister to an Afrotrilepis—Trilepis Clade. This agrees with morphological argument (see Key to the genera of Trilepideae).

4.3.2 Tribe Cladieae

Tribe Cladieae (Fig. 5B) is monogeneric including only the cosmopolitan genus *Cladium*. Uncertainty remains concerning the relationship between *Cladium* and the monotypic genus *Rhynchocladium* from the Guiana Shield in Guyana and Venezuela. Despite several attempts, *Rhynchocladium* has never been successfully included in a molecular study.

Morphological similarities point at possible relationships between Rhynchocladium and Cladium (Cladieae), Rhynchospora (Rhynchosporeae), and Machaerina (Schoeneae) (Maguire, 1972; Goetghebeur, 1986). Consequently, Rhynchocladium is the only genus here considered as unplaced in the proposed classification.

4.3.3 Bisboeckelereae-Sclerieae Clade

The circumscription of tribe Bisboeckelereae (Fig. 4A) and tribe Sclerieae (Fig. 5A) has not changed since the study of Goetghebeur (1998). Becquerelia cymosa Brong. and Diplacrum africanum (Benth.) C.B.Clarke have always been retrieved as a clade, ever since the study of Simpson et al. (2007). Muasya et al. (2009a) also included Calyptrocarya Nees in their study; however, this resulted in Bisboeckelereae not being monophyletic in their results (but sister to Scleria P.J.Bergius and Becquerelia Brongn. + Diplacrum R.Br.). With a wider species sampling, Hinchliff & Roalson (2013) found a grade of Bisboeckelereae leading to a monophyletic Sclerieae. Only recent studies (Bauters et al., 2016; Spalink et al., 2016b; Semmouri et al., 2019) retrieved a monophyletic Bisboeckelereae sister to the monogeneric Sclerieae. Our results confirm this topology with a monogeneric Sclerieae Clade sister to a Bisboeckelereae Clade (Figs. 2, 3). The latter includes two subclades: (i) Becquerelia + Diplacrum and (ii) Bisboeckelera + Calyptrocarya (Fig. 2A), which agrees with morphological arguments (see key to the genera of Bisboeckelereae in Section 5).

4.3.4 Tribe Carpheae

The position of the clade including the genera Carpha Banks & Sol. ex R.Br. and Trianoptiles Fenzl ex Endl. is variable in the literature. In some studies, Carpha is positioned within Schoeneae s.l. clade (Zhang et al., 2004, 2007; Verboom, 2006; Muasya et al., 2009a; Hinchliff & Roalson, 2013), whereas here and in other studies, Carpha (+ Trianoptiles) is placed outside Schoeneae (Simpson et al., 2007; Jung & Choi, 2013; Viljoen et al., 2013; Larridon et al., 2018a; Semmouri et al., 2019). This clade is also set apart by its unique combination of embryo morphology characters, having an embryo that is more or less rhomboid to top-shaped with a tapered scutellum, with a well-differentiated root cap in a lateral position separated from the coleoptile by a notch. This led Semmouri et al. (2019) to erect a new tribe to accommodate the genera Carpha and Trianoptiles. We retrieve tribe Carpheae (Fig. 4C) as a separate speciespoor clade (Figs. 2, 3).

4.3.5 Tribe Cryptangieae

The relationship between tribe Cryptangieae (Fig. 5G) and the other Cyperaceae tribes varied in different studies (cf. Muasya et al., 2009a; Escudero & Hipp, 2013; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Spalink et al., 2016b; Semmouri et al., 2019). In earlier studies, the diversity of the tribe was represented by only two to three species. On the basis of sequence data of an increased sampling (18 spp.), Costa et al. (2018b) recovered Cryptangieae as monophyletic, supporting its recognition as a tribe in combination with its morphological distinctiveness. However, it remained unclear

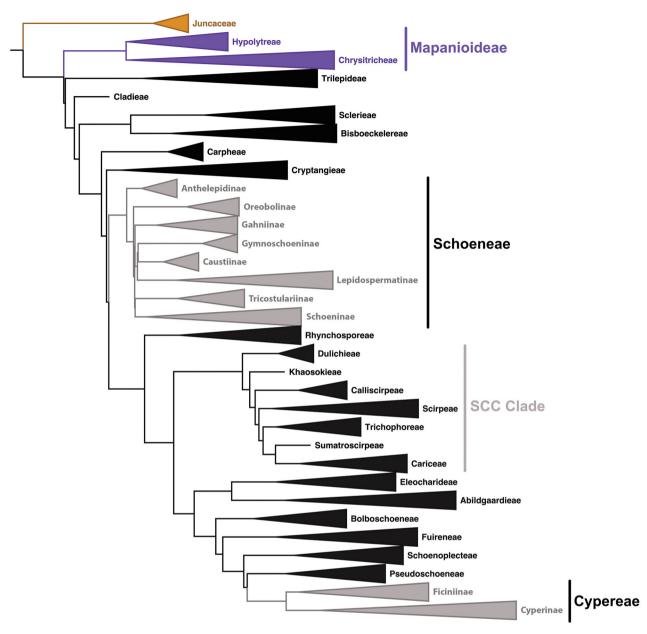


Fig. 3. Summary phylogeny of Cyperaceae at the tribal level showing the topology resulting from the coalescent ASTRAL analysis.

if the tribe is more closely related to tribe Schoeneae, tribe Carpheae, or to a Carpheae–Schoeneae Clade. Also, it did not include sequence data of the genus Koyamaea. Our targeted sequencing results show that a clade of Koyamaea plus Cryptangieae sensu Costa et al. (2018b) branches after tribe Carpheae and before tribe Schoeneae in the coalescent ASTRAL analysis (Figs. 2, 3); however, in the concatenated IQTREE analysis, the positions of Cryptangieae and Schoeneae are inverted (Fig. S2).

Koyamaea neblinensis W.W.Thomas & G.Davidse was described as a new genus and species to science by Thomas & Davidse (1989). Due to its bisexual spikelets, each bearing one pistillate flower and many staminate flowers, presence of well-developed perianth bristles in both kinds of flowers,

spirally arranged glumes, and regular nutlet without a cupule, Koyamaea was classified as a new genus of the then recognized subfamily Sclerioideae (Goetghebeur, 1998). As the authors believed that their new genus was not closely related to any other genus, they placed it in its own tribe Koyamaeae. Sanger sequence data could not be obtained for this extremely rare species to test its placement in the family and the value of erecting a separate monotypic tribe for it. High-throughput sequencing techniques are better at dealing with fragmented DNA obtained from historical herbarium specimens (e.g., Buerki & Baker, 2015; Hart et al., 2016; Bakker, 2017; Zeng et al., 2018; Brewer et al., 2019). Our targeted sequencing results are the first to place the genus Koyamaea in the Cyperaceae Tree of Life. It is here inferred as

 Table 1
 Proposed classification with number of species per genus and tribe, and number of genera per tribe

				Species/	Species/	Genera/
Subfamily	Tribe	Subtribe	Genera	genus	tribe	tribe
Mapanioideae C.B.Clarke	Hypolytreae Pers.		Hypolytrum Pers.	63		
			Mapania Aubl.	100		
			Paramapania Uittien	7		
			Scirpodendron Zipp. & Kurz	2	172	4
	Chrysitricheae Nees		Capitularina J.Kern.	-		
			Chorizandra R.Br.	9		
			Chrysitrix L.	4		
			Diplasia Pers.	-		
			Exocarya Benth.	τ-		
			Lepironia Pers.	-	41	9
Cyperoideae Beilschm.	Trilepideae Goetgh.		Afrotrilepis (Gilly) J.Raynal	2		
			Coleochloa Gilly	8		
			Microdracoides Hua	-		
			Trilepis Nees	5	16	4
	Cladieae Nees		Cladium P.Browne	Μ	Μ	-
	Sclerieae Wight & Arn.		Scleria P.J.Bergius	258	258	-
	Bisboeckelereae Pax		Becquerelia Brongn.	9		
			Bisboeckelera Kuntze	4		
			Calyptrocarya Nees	8		
			Diplacrum R.Br.	10	28	4
	Carpheae Semmouri & Larridon		Carpha Banks & Sol. ex R.Br.	15		
			Trianoptiles Fenzl ex Endl.	٣	18	2
	Cryptangieae Benth.		Cephalocarpus Nees	20		
			Cryptangium Schrad. ex Nees	-		
			Didymiandrum Gilly	-		
			Exochogyne C.B.Clarke	2		
			Koyamaea W.W.Thomas & Davidse	-		
			Krenakia S.M.Costa	10		
			Lagenocarpus Nees	15	50	7
	Schoeneae Dumort.	Anthelepidinae R.L.Barrett	Anthelepis R.L.Barrett, K.L.Wilson &	4		
			J.J.Bruni			
		Caustiinae R.L.Barrett	Caustis R.Br.	7		
			Evandra R.Br.	7		
		Gahniinae Pax	Cyathochaeta Nees	5		
			Gahnia J.R.Forst. & G.Forst.	14		
			Mesomelaena Nees	5		
			Ptilothrix K.L.Wilson	-		
						Continued

7	C	3
	a	j
	Ξ	2
	C	
:	F	3
	C	
	c	٥
ι	٥	j

Table 1 Continued						
				Species/	Species/	Genera/
Subfamily	Tribe	Subtribe	Genera	genns	tribe	tribe
		Gymnoschoeninae	Gymnoschoenus Nees	7		
		R.L.Barrett	Reedia F.Muell.	~		
		Lepidospermatinae R.L.Barrett	repiaosperna Labin.	80		
			Machaerina Vahl	55		
			Neesenbeckia Levyns	-		
			Netrostylis R.L.Barrett, J.J.Bruhl &	£		
			K.L.Wilson			
		Oreobolinae R.L.Barrett	Capeobolus Browning	-		
			Chamaedendron (Kük.) Larridon	2		
			Costularia C.B.Clarke	15		
			Cyathocoma Nees	Μ		
			Oreobolus R.Br.	17		
		Schoeninae Fenzl	Schoenus L.	149		
		Tricostulariinae R.L.Barrett	Ammothryon R.L.Barrett, K.L.Wilson	_		
			& J.J.Bruhl			
			Chaetospora R.Br.	\sim		
			Morelotia Gaudich.	9		
			Tetraria P.Beauv.	39		
			Tricostularia Nees	F		
			Xyroschoenus Larridon	_	466	25
	Rhynchosporeae Nees		Rhynchospora Vahl*	399	399	-
	Dulichieae W.Schultze-Motel		Blysmopsis Oteng-Yeb.	_		
			Blysmus Panz. ex Schult.	\sim		
			Dulichium Pers.	-	7	\sim
	Khaosokieae LévBourret & J.R.Starr		Khaosokia D.A.Simpson	-	-	-
	Calliscirpeae LévBourret & J.R.Starr		Calliscirpus C.N.Gilmour, J.R.Starr &	7	2	-
			Naczi			
	Scirpeae T.Lestib.		Amphiscirpus Oteng-Yeb.	~		
			Eriophorum L.	48		
			Phylloscirpus C.B.Clarke	\sim		
			Rhodoscirpus Léveillé-Bourret,	_		
			Donadío & J.R.Starr			
			Scirpus Tourn. ex L.	47		
			Zameioscirpus Dhooge & Goetgh.	٣	73	9
	Trichophoreae LévBourret & J.R.Starr		Trichophorum Pers.	19	19	-
	Sumatroscirpeae LévBourret & J.R.Starr		Sumatroscirpus Oteng-Yeb.	4	4	-

_	_	
ĺ	ă	
	<u> </u>	
	Ē	
:	Ē	
1	Ċ	
,	C	
ļ	_	,
	•	
_	q	Į
1	c	١
Ī	n	
ŀ	-	
		Table 1 Continued

				Species/	Species/	Genera/
Subfamily	Tribe	Subtribe	Genera	genus	tribe	tribe
	Cariceae Dumort.		Carex L.	2003	2003	-
	Eleocharideae Goetgh.		Eleocharis R.Br.*	302	302	-
	Abildgaardieae Lye		Abildgaardia Vahl*	6		
			Actinoschoenus Benth.	2		
			Arthrostylis R.Br.	2		
			Bulbostylis Vahl*	227		
			Fimbristylis Vahl*	320		
			Nelmesia Van der Veken*	-		
			Scleroschoenus K.L.Wilson, J.J.Bruhl	9		
			& R.L.Barrett			
			Trachystylis S.T.Blake	~		
			Trichoschoenus J.Raynal	-		
			Zulustylis Muasya	2	571	10
	Bolboschoeneae (Tatanov) J.R.Starr		Bolboschoenus (Asch.) Palla	15	15	-
	Fuireneae Rchb. ex Fenzl		Fuirena Rottb.	55	55	-
	Schoenoplectieae Lye		Actinoscirpus (Ohwi) R.W.Haines	~		
			& Lye			
			Schoenoplectus (Rchb.) Palla	16	17	7
	Pseudoschoeneae J.R.Starr		Pseudoschoenus (C.B.Clarke)	~		
			Oteng-Yeb.			
			Schoenoplectiella Lye	63	64	2
	Cypereae Dumort.	Ficiniinae Fenzl	Afroscirpoides García-Madr. &	~		
			Muasya			
			Dracoscirpoides Muasya	κ		
			Erioscirpus Palla	2		
			Ficinia Schrad.	87		
			Hellmuthia Steud.	~		
			Isolepis R.Br.	69		
			Scirpoides Ség.	4		
		Cyperinae Pax	Cyperus L.*	964	1131	8
	Incertae cedis		Rhynchocladium T.Koyama	-	-	-
2 subfamilies	24 tribes	10 subtribes		5687		95 genera
				species		

follows Novoselova (1994a, 1994b); (v) Trichophorum that follows Léveillé-Bourret et al. (2020); (vi) Abildgaardieae that follow Larridon et al. (2021); (vii) Ficinia and Isolepis that follow Muasya & Larridon (2021); and (ix) Cyperus where the number reflects the recent synonymization of Androtrichum (Brongn.) Brongn. (Pereira-Silva et al., 2020). Genera including species using the C₄ photosynthetic pathway are indicated with an asterisk (*). pers. comm., 2020); (iii) Rhynchospora where the number reflects the recent synonymization of Pleurostachys Brongn. (Thomas, 2020); (iv) Eriophorum where the number of species The number of species follows Govaerts et al. (2020) except for (i) Cryptangieae that follow Costa SM (pers. comm., 2020); (ii) Schoeneae that follow Barrett RL, Elliott TL, and Larridon I

sister to the genera of Cryptangieae (Fig. 2A). As there are other arguments linking *Koyamaea* to Cryptangieae, that is, morphological (flowers with spiral glumes, bearing perianth and lacking cupule, beaked fruits) and anatomical (thickened pericarp) shared features, we opt to include *Koyamaea* in Cryptangieae.

Generic delimitation in Cryptangieae has fluctuated over the years, either with the number of genera considered in the strict sense, including just one or few species, or lumped into a broader Lagenocarpus Nees. Recent molecular studies highlighted the need of an updated generic circumscription (Costa et al. 2018b, 2021a). In the new interpretation, Cephalocarpus Nees includes the species formerly placed in Everardia Ridley and now encompasses the 20 species of Cryptangieae with an elongate caudex and lateral inflorescences (Costa et al., 2021a, 2021b). In our results, the monophyly of the newly enlarged genus Cephalocarpus is supported by the concatenated IQ-TREE analysis (Fig. S2), but not by the coalescent ASTRAL analysis where Cephalocarpus angustus (N.E.Brown) S.M.Costa (syn. Everardia angusta N.E.Brown) and Cephalocarpus montanus (Ridl.) S.M.Costa (syn. Everardia montana Ridl.) are not retrieved in a single clade (Fig. 2A). It should be noted that both species were formerly placed in Everadia; we did not sequence the type species of Cephalocarpus (Cephalocarpus dracaenula Nees). Lagenocarpus (sensu Koyama 2005) species have been split in five genera (Costa et al., 2021a): three of them (Cryptangium Schrader ex Nees, Dydimiandrum Gilly, and Exochogyne C.B.Clarke) with 1-2 species and the others with 10 (Krenakia S.M.Costa; Costa et al., 2021a) and 15 species (Lagenocarpus). The genera are distinguished mostly by leaf, inflorescence, and fruit characters (Costa et al., 2021a). The results of Costa et al. (2021a) place Krenakia as sister to a clade encompassing three subclades: (i) Didymiandrum + Exochogyne; (ii) Cryptangium sister to Cephalocarpus; and (iii) Lagenocarpus s.s. Our targeted sequencing results show Koyamaea sister to a clade encompassing the Cryptangieae sensu Costa et al. (2018b, 2021a).

4.3.6 Tribe Schoeneae

Tribe Schoeneae (Figs. 5E, 5F) has been widely recognized, but its exact circumscription has long remained unclear. However, tribe Schoeneae s.s. (excluding Carpheae and Cladieae) has been shown to be monophyletic in recent studies (Viljoen et al., 2013; Spalink et al., 2016b; Larridon et al., 2018a; Semmouri et al., 2019). Goetghebeur (1998) and Simpson et al. (2007) had also placed *Pleurostachys* Brongn. and *Rhynchospora* in Schoeneae s.l., but these genera were moved to a separate tribe Rhynchosporeae by Muasya et al. (2009). Our targeted sequencing results (Fig. 2B) confirm a monophyletic tribe Schoeneae sensu Semmouri et al. (2019).

Relationships within tribe Schoeneae have not been entirely resolved; however, progress has been made in our understanding of its evolution. Morphologically, tribe Schoeneae is a highly variable group. Previous molecular analyses of the group recovered six main clades: the Caustis Clade, Gahnia Clade, Lepidosperma Clade, Oreobolus Clade, Schoenus Clade, and Tricostularia Clade (Viljoen et al., 2013; Larridon et al., 2018a). In the more deeply sampled phylogenetic study of Semmouri et al. (2019), two additional clades became visible, a clade including the genera *Reedia*

F.Muell. and Gymnoschoenus Nees and a separate lineage of Schoenus paludosus (R.Br.) Roem. & Schult. Our targeted sequencing results confirm the presence of eight main clades in Schoeneae (Fig. 2B). To facilitate the morphological characterization of the main clades in this morphologically diverse tribe, they are recognized as subtribes in Section 5.

Previous molecular studies remarked on the polyphyly of genera in Schoeneae (Zhang et al., 2004; Muasya et al., 2009a; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Viljoen et al., 2013; Larridon et al., 2018a; Barrett et al., 2019). This mostly concerned the genera Costularia C.B.Clarke, Epischoenus C.B.Clarke, Schoenus, Tetraria P.Beauv., and Tricostularia, which had species scattered across different clades in tribe Schoeneae, greatly confusing the boundaries of the subtribes recognized here. Recently, efforts have been made to realign the taxonomy of these genera to make them monophyletic.

Anthelepis Clade: In the BI and ML results of Semmouri et al. (2019), Schoenus paludosus formed a polytomy with the Gahnia Clade and the Oreobolus Clade, revealing its isolated position from other Schoenus species. In fact, Schoenus paludosus also differs morphologically from the true Schoenus species in having one or sometimes two lower male flowers and an upper bisexual flower at each spikelet, besides a non-zigzag rachilla (as opposed to the usual states for the genus of bisexual flowers and upper internodes of the rachilla elongated and prominently zigzag; Wilson, 1993). Schoenus paludosus was recently placed in a new genus Anthelepis R.L.Barrett, K.L.Wilson & J.J.Bruhl together with the species previously named Schoenus guillauminii Kük. and Tricostularia undulata (Thwaites) J.Kern (Barrett et al., 2019). The Anthelepis Clade is here strongly supported as sister to the remainder of Schoeneae (Fig. 2B).

Caustis Clade: This clade includes the genera *Caustis* R.Br. and *Evandra* R.Br. (Fig. 2B). The unexpected placement of a lineage previously included in *Tetraria*, that is, *Tetraria borneensis* J.Kern, in the Caustis Clade (Larridon et al., 2018a) is being explored further (Barrett RL & Larridon I, unpublished data). This species could not be sequenced for this study.

Gahnia Clade: The circumscription of the Gahnia Clade, including the genera *Cyathochaeta* Nees, *Gahnia* J.R.Forst. & G.Forst., *Mesomelaena* Nees, and *Ptilothrix* K.L.Wilson (Fig. 2B), remains unchanged, but notably excludes *Morelotia* Gaudich., previously included under *Gahnia*, which is placed in the Tricostularia Clade (e.g., Verboom, 2006; Muasya et al., 2009a; Jung & Choi, 2013; Hinchliff & Roalson, 2013; Viljoen et al., 2013; Larridon et al., 2018a; Semmouri et al., 2019; this study).

Gymnoschoenus—Reedia Clade: This is a small clade of just three morphologically distinctive species placed in the genera *Gymnoschoenus* and *Reedia* (Fig. 2B), each with restricted distributions in southern Australia whose affinities have been much debated.

Lepidosperma Clade: Another lineage previously included in *Tetraria*, that is, the *Tetraria capillaris* (F.Muell.) J.M.Black species complex, native to Australia and New Zealand, was found to be part of the Lepidosperma Clade (Viljoen et al., 2013; Larridon et al., 2018a; Barrett et al., 2019). A recent taxonomic revision of the *Tetraria capillaris* species complex resulted in the publication of

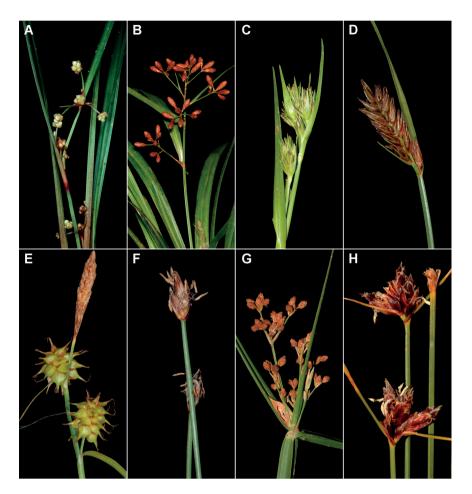


Fig. 4. Morphological diversity of the Cyperaceae tribes. A, Hypolytreae, Mapania floribunda (Nees ex Steud.) T.Koyama. B, Bisboeckelereae, Calyptrocarya poeppigiana Kunth. C, Carpheae, Carpha capitellata (Nees) Boeckeler. D, Dulichieae, Blysmus compressus (L.) Panz. ex Link. E, Cariceae, Carex lepidocarpa Tausch. F, Eleocharideae, Eleocharis quinqueflora (Hartmann) O.Schwarz. G, Schoenoplecteae, Actinoscirpus grossus (L.f.) Goetgh. & D.A.Simpson. H, Cypereae: Ficiniinae, Ficinia acuminata (Nees) Nees. Photos by Modesto Luceño.

a new genus Netrostylis (Barrett et al., 2021a). The Lepidosperma Clade appears to have originated in Australia (Viljoen et al., 2013). Previous studies (e.g., Verboom, 2006; Muasya et al., 2009a; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Viljoen et al., 2013; Larridon et al., 2018a; Semmouri et al., 2019) indicated that the clade includes (i) the species-rich genus Lepidosperma occurring from China to Australasia; (ii) Machaerina Vahl (including Baumea Gaudich.), which is widespread from Australia to the Americas; and (iii) the monotypic genus Neesenbeckia Levyns endemic from the South African Cape Floristic Region. As retrieved in, for example, Viljoen et al. (2013) and Larridon et al. (2018a), our results confirm that within the Lepidosperma Clade, the genus Machaerina is sister to a clade with two subclades: (i) Lepidosperma and (ii) Netrostylis sister to Neesenbeckia (Fig. 2B). The latter sister relationship between Netrostylis and Neesenbeckia suggests an unusual dispersal event from Australia to southern Africa. Although some taxonomic issues remain in Machaerina (Barrett RL, Wilson KL & Bruhl JJ, unpublished data), more work is required in Lepidosperma, which has c. 200 undescribed species in southern Australia (Barrett & Wilson, 2012, 2013).

Oreobolus Clade: Larridon et al. (2018a) found that Costularia s.l. was composed of four distinct evolutionary lineages with two lineages being part of the Oreobolus Clade: (i) a much-reduced genus Costularia (Larridon et al., 2019a) and (ii) a small New Caledonian endemic genus Chamaedendron. The circumscription of the other genera in this clade (Fig. 2B), that is, Capeobolus, Cyathocoma Nees, and Oreobolus R.Br., remains unchanged.

Schoenus Clade: As some species of *Tetraria* and *Epischoenus* had been shown to be nested within *Schoenus* (Viljoen et al., 2013), Elliott & Muasya (2017) transferred these species to *Schoenus*. The broader circumscription of *Schoenus* is supported by our targeted sequencing results (Fig. 2B), and only a single morphologically variable and geographically widespread genus is recognized in this clade.

Tricostularia Clade: Larridon et al. (2018a) found that the two other distinct evolutionary lineages previously placed in Costularia s.l. were part of the Tricostularia Clade: (i) a single-species lineage from the Seychelles Xyroschoenus hornei

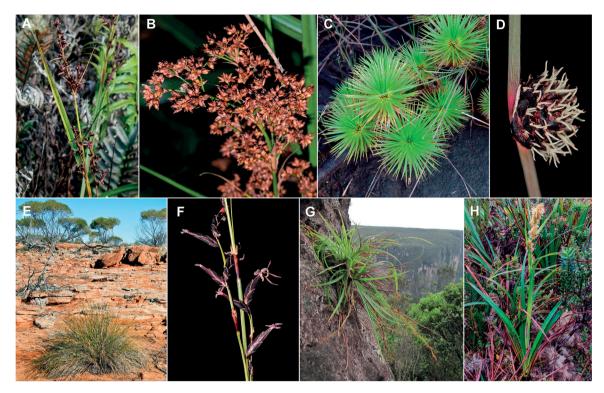


Fig. 5. Morphological diversity of the Cyperaceae tribes. **A,** Sclerieae, *Scleria gaertneri* Raddi. **B,** Cladieae, *Cladium mariscus* subsp. *intermedium* Kük. **C,** Trilepideae, *Microdracoides squamosa* Hua. **D,** Chrysitricheae, *Chorizandra enodis* Nees. **E,** Schoeneae, *Lepidosperma* Labill. sp. **F,** Schoeneae, *Schoenus melanostachys* R.Br. **G,** Cryptangieae, *Cephalocarpus* cf. *maguireanus* (T.Koyama) S.M.Costa. **H,** Rhynchosporeae, *Rhynchospora* Vahl sp. Photos **A, B,** and **D–F** by Russell Barrett; **C** by Isabel Larridon; **G** by Suzana Costa.

(C.B.Clarke) Larridon and (ii) the rest of Costularia subgenus Lophoschoenus that formed a grade including the African "reticulate-sheathed" Tetraria species, with a single Epischoenus species nested within this clade (i.e., Epischoenus cernuus Levyns). As Elliott & Muasya (2017) had already transferred the "non-reticulate sheathed" species of Tetraria, including the original type species of Tetraria, that is, T. thuarii P.Beauv. (= Schoenus compar L.), and most of Epischoenus into Schoenus, Larridon et al. (2017) proposed to conserve the genus Tetraria with a new type, T. thermalis (L.) C.B.Clarke. The same authors recircumscribed Tetraria as including c. 30 "reticulate sheathed" Tetraria species, Epischoenus cernuus, and most of Costularia subgenus Lophoschoenus sensu Kükenthal (1939) (Larridon et al., 2018b).

The Tricostularia Clade also includes the Australian species Tetraria octandra (Nees) Kük., which Larridon et al. (2018a) suggested should be accepted as T. octandra (Nees) C.B.Clarke, as it is not related to Tetraria. The taxonomic changes made to Costularia s.l. and Tetraria by Larridon et al. (2017, 2018a, b) are supported by our targeted sequencing results (Fig. 2B). More recent research has shown that three Australian species until recently placed in Tetraria, T. australiensis C.B.Clarke, T. microcarpa S.T.Blake, and T. octandra are closely related to Morelotia and Xyroschoenus (Barrett et al., 2021b). Therefore, the decision has been taken to expand the circumscription of Morelotia, by including the three Australian Tetraria species and a Pacific Island species

(originally described as *Machaerina* involuta H.St.John) (Barrett et al., 2021b).

Some Schoenus species have been recovered in the Tricostularia Clade, that is, Schoenus curvifolius (R.Br.) Roem. & Schult., S. grandiflorus (Nees) F.Muell., and S. turbinatus (R.Br.) Roem. & Schult. (Viljoen et al., 2013; Larridon et al., 2018a; Barrett et al., 2019; Semmouri et al., 2019). Further research has led to the resurrection and recircumscription of the genus Chaetospora R.Br. (Barrett et al., 2019, 2020). Schoenus grandiflorus has been recognized as a monotypic genus Ammothryon as it is not recovered as sister to any genus in the clade (Barrett et al., 2021b).

Within the Tricostularia Clade, the genus *Tricostularia* itself has been reduced in morphological circumscription with the removal of species now placed in *Anthelepis* (Barrett et al., 2019). In parallel, the number of species was enlarged with the addition of *Lepidosperma aphylla* R.Br. and *L. exsul* C.B.Clarke (Barrett & Wilson, 2012) and ongoing taxonomic revision of species boundaries in southern Western Australia (Barrett, 2012; Barrett et al., 2021b).

4.3.7 Tribe Rhynchosporeae

The topology of the remaining Cyperoideae lineages in our results largely matches previous studies (Simpson et al., 2007; Muasya et al., 2009a; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Spalink et al., 2016b; Costa et al., 2018b; Semmouri et al., 2019), with tribe Rhynchosporeae (Fig. 5H) sister to the Scirpo-Caricoid Clade (SCC Clade) plus the

Abildgaardieae—Eleocharideae—Fuireneae—Cypereae Clade (FAEC Clade) (Fig. 2). Tribe Rhynchosporeae is now accepted as monogeneric. The nested position of *Pleurostachys* Brongn. in *Rhynchospora* Vahl has been known for a long time (Thomas et al., 2009; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Semmouri et al., 2019). Moreover, their similar embryo morphology (*Carex*-type embryo; Semmouri et al., 2019) provided an additional argument for the merging of the two taxa. Thomas (2020) recently merged *Pleurostachys* into *Rhynchospora*.

4.3.8 Scirpo-Caricoid Clade

A Scirpo-Caricoid Clade (SCC Clade), referred to as the Scirpeae-Dulichieae-Cariceae Clade (SDC Clade) in some previous studies, was recognized in all recent molecular phylogenetic studies of Cyperaceae as a sister group to the Abildgaardieae–Eleocharideae–Fuireneae–Cypereae (FAEC Clade) (Muasya et al., 2009a; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Spalink et al., 2016b; Léveillé-Bourret et al., 2014, 2015, 2018b, 2018c; Semmouri et al., 2019). The Scirpo-Caricoid Clade contains 41% of all recognized Cyperaceae species (Léveillé-Bourret & Starr, 2019; Govaerts et al., 2020), comprising a curious assemblage of tribes that illustrates the full breadth of inflorescence and floral diversity of Cyperoideae. This includes bisexual, monoecious, and dioecious species with empty proximal glumes or all glumes fertile, spirally or distichously inserted flowers, sterile or fertile prophylls, as well as setiform, tepaliform, or absent perianth (Léveillé-Bourret & Starr, 2019). There are no recognized synapomorphies for this clade, whose only recognizable characteristic is its center of diversity in cold temperate regions of the Northern Hemisphere (Léveillé-Bourret & Starr, 2019; Martín-Bravo et al., 2019), contrasting with the southern temperate distribution for Schoeneae or mostly tropical diversity of other major Cyperaceae lineages.

The monogeneric tribe Cariceae contains most of the diversity of the clade, with c. 2000 species (Roalson et al., 2021; Villaverde et al., 2020, 2021), and is characterized by a highly derived inflorescence morphology formed of perianthless unisexual flowers, with female flowers strictly associated with the production of secondary branches, and pistils contained within or subtended by the first bract of secondary branches (a prophyll called a perigynium or utricle if closed forming a bottle-like structure; Jiménez-Mejías et al., 2016b). Although relationships within this tribe are not highly supported in the present study, they have been already addressed in Villaverde et al. (2020).

The other 13 genera (c. 88 species) of the Scirpo-Caricoid Clade have all been placed at one point in their history within a broadly circumscribed "tribe Scirpeae," which was essentially defined by a lack of derived characters. Unsurprisingly, most recent studies suggested paraphyly of Scirpeae when thus circumscribed (Hinchliff & Roalson, 2013; Jung & Choi, 2013; Spalink et al., 2016b). However, an ancient rapid radiation near the crown of the Scirpo-Caricoid Clade made previous phylogenetic analyses extremely difficult, with different analyses supporting different topologies with consistently low support (Hinchliff & Roalson, 2013; Jung & Choi, 2013; Spalink et al., 2016b; Semmouri et al., 2019).

A series of recent studies combining plastid and nuclear ribosomal markers (Gilmour et al., 2013; Léveillé-Bourret et al.

2014, 2015, 2018a, 2019), genomic data (Léveillé-Bourret et al., 2018c; Villaverde et al., 2020, 2021), and morphological data (Léveillé-Bourret & Starr, 2019) were able to resolve the most recalcitrant backbone branches of the Scirpo-Caricoid phylogeny. Our present results (Fig. 2C) are in agreement with these recent studies and support the taxonomic treatment of the clade as presented in Léveillé-Bourret & Starr (2019).

4.3.8.1 Tribe Dulichieae. As in previous studies, tribe Dulichieae (Fig. 4D), excluding *Sumatroscirpus* Oteng-Yeb., is strongly supported as monophyletic (e.g., Gilmour et al., 2013; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Léveillé-Bourret et al., 2014, 2015, 2018a, 2018c; Léveillé-Bourret & Starr, 2019; Semmouri et al., 2019). It is easily diagnosed by the presence of distichous spikelets and flowers, scale-like spikelet prophylls subtending flowers, and long narrow beaks on fruits. Two genera have commonly been recognized in Dulichieae, that is, *Blysmus* and *Dulichium* Pers. (Goetghebeur, 1998), but recent studies provide some support for recognizing *Blysmopsis* (Léveillé-Bourret et al., 2014; Semmouri et al., 2019), which was commonly treated as a synonym of *Blysmus* (Goetghebeur, 1998).

4.3.8.2 Tribe Khaosokieae. The monogeneric tribe Khaosokieae (Fig. 6B) represents a single recently described genus and species (Simpson et al., 2005) that has proven difficult to place in the suprageneric classification based on morphological arguments due to characters suggesting affinities with several different tribes, that is, unisexual flowers like Cariceae, seven perianth bristles and narrow elongate spikelets suggestive of Dulichium (tribe Dulichieae), but spirally inserted flowers and sterile prophylls typical of tribe Scirpeae. All phylogenetic studies have supported its isolated position within the Scirpo-Caricoid Clade, either as sister to all other lineages of the tribe (Muasya et al., 2009a; Jung & Choi, 2013; Semmouri et al., 2019) or as the next diverging lineage after tribe Dulichieae (Simpson et al., 2007; Escudero & Hipp, 2013; Hinchliff & Roalson, 2013; Léveillé-Bourret et al., 2014, 2015, 2018a, 2018c; Spalink et al., 2016b; this study, Fig. 2C).

4.3.8.3 Tribe Calliscirpeae. This monogeneric lineage was recently recognized (Léveillé-Bourret & Starr, 2019) based on species formerly placed in *Scirpus* L. or *Eriophorum* L., but that differ by having antrorsely barbed perianth bristles and a Carextype embryo (Gilmour et al., 2013). All previous studies have consistently supported the isolated position of this lineage in the Scirpo-Caricoid Clade, but its phylogenetic position as sister to a Scirpeae–Trichophoreae–Sumatroscirpeae–Cariceae Clade has never received strong support (Léveillé-Bourret et al., 2014, 2015, 2018a, 2018c; Semmouri et al., 2019). In our results, tribe Calliscirpeae (Fig. 6A) branches after Khaosokieae, sister to the remaining lineages of the Scirpo-Caricoid Clade (Fig. 2C).

4.3.8.4 Tribe Scirpeae. Tribe Scirpeae (Fig. 6C) is a lineage of the Scirpo-Caricoid Clade (e.g., Simpson et al., 2007; Muasya et al., 2009a; Hinchliff & Roalson, 2013; Léveillé-Bourret et al., 2014, 2018a, 2018c; Spalink et al., 2016b; Semmouri et al., 2019). As previously discussed (see Section 4.3.8), tribe Scirpeae sensu Goetghebeur (1998) is not mono-



Fig. 6. Morphological diversity of the Cyperaceae tribes. A, Calliscirpeae, Calliscirpus brachythrix C.N.Gilmour, J.R.Starr & Naczi. B, Khaosokieae, Khaosokia caricoides D.A.Simpson. C, Scirpeae, Scirpus sylvaticus L. D, Trichophoreae, Trichophorum alpinum (L.) Pers. E, Sumatroscirpeae, Sumatroscirpus rupestris Lév.-Bourret & J.R.Starr. Photos A and E by Julian Starr; B by Rachun Pooma; C and D by Modesto Luceño.

phyletic and consists of three separate lineages. As a result, Scirpeae was recircumscribed by Léveillé-Bourret & Starr (2019) to include only species possessing a (sub-)lateral germ pore in their embryos, corresponding to Schoenustype, Fimbristylis-type, or intermediate embryo types. Under this circumscription, Scirpeae is monophyletic. No visible macromorphological character has been found that can unambiguously diagnose this tribe, which means that identification must be done by means of exclusion. Two major subclades are found within this monophyletic Scirpeae (Fig. 2C): (i) a mostly South American group that has been dubbed "Zameioscirpus Clade" is supported in many studies (Dhooge et al., 2003; Muasya et al., 2009a; Léveillé-Bourret et al., 2015), including Amphiscirpus Oteng-Yeb., Phylloscirpus C.B.Clarke, Rhodoscirpus Léveillé-Bourret, Donadío & J.R.Starr, and Zameioscirpus Dhooge & Goetgh.; and (ii) a mostly circumboreal "Scirpus Clade," well supported in our analyses and consistently found in other studies, with the genus Eriophorum L. forming a wellsupported clade nested within Scirpus, thus making Scirpus paraphyletic (e.g., Gilmour et al., 2013; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Léveillé-Bourret et al., 2014, 2018a, 2018c).

The circumscription of Eriophorum and Scirpus is one of the last adjustments needed to achieve a classification of Cyperaceae where all genera are circumscribed as monophyletic entities. Two options for resolving this issue appear viable: (i) Eriophorum can be merged with Scirpus as proposed by Koyama (1958), or (ii) Eriophorum can be maintained by splitting Scirpus into a series of new genera. Both of these solutions have drawbacks. If Eriophorum is treated within Scirpus, the specific epithets for some wellknown species such as Eriophorum gracile W.D.J.Koch would suddenly be unfamiliar to most in the botanical community (i.e., =Scirpus ardea T.Koyama). However, maintaining Eriophorum would require splitting Scirpus into six to eight genera, each consisting of one to a dozen species. As a taxonomically well-sampled and strongly supported phylogeny for Scirpeae is still lacking, the extent of the taxonomic changes needed to split Scirpus remains unclear. Consequently, a decision on merging or splitting should wait until conclusive phylogenetic data are gathered.

4.3.8.5 Tribe Trichophoreae. The recently recognized tribe Trichophoreae (Fig. 6D) contains species that were previously

associated with Scirpeae, but lack the lateral embryo germ pore that now defines tribe Scirpeae s.s. (Léveillé-Bourret et al., 2019). A clade comprising species placed in this tribe has been retrieved within the Scirpo-Caricoid Clade in nearly all phylogenetic studies (Muasya et al., 2009a; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Léveillé-Bourret et al., 2014, 2015, 2018a; Spalink et al., 2016b; Semmouri et al., 2019), with the exception of early analyses based only on rbcL (e.g., Muasya et al., 1998; Simpson et al., 2007). However, the phylogenetic placement of Trichophoreae within the Scirpo-Caricoid Clade was settled only recently using a phylogenomic approach (Léveillé-Bourret et al., 2018c). Trichophoreae is monogeneric with the recent inclusion of the formerly recognized genera Oreobolopsis T.Koyama & Guagl. and Cypringlea M.T.Strong within Trichophorum Pers. (Léveillé-Bourret et al., 2020). Under this monophyletic circumscription (Fig. 2C), Trichophorum is highly variable in habit, distribution, and ecology, including short unispicate arctic-alpine species, as well as large paniculate subtropical species. Nonetheless, the genus can be distinguished by having only basal leaves usually with mucronate lamina and proximal glumes empty and/or with a longer awn.

4.3.8.6 Tribe Sumatroscirpeae. The genus Sumatroscirpus Oteng-Yeb. was formerly placed in Dulichieae (Goetghebeur 1998), but unlike other members of this tribe, it possesses tubular fertile prophylls similar to the perigynia of Cariceae. Léveillé-Bourret et al. (2018a) demonstrated that Sumatroscirpus is actually sister to Cariceae and that it corresponds to a morphologically transitional lineage between Cariceae and Scirpeae. This result was confirmed by Semmouri et al. (2019) and is here again confirmed with a completely independent data set (Fig. 2C). We, thus, support its recognition as a monogeneric tribe Sumatroscirpeae (Fig. 6E).

4.3.8.7 Tribe Cariceae. Tribe Cariceae (Fig. 4E) is strongly supported as monophyletic as in previous studies (e.g., Global Carex Group, 2015; Starr et al., 2015; Jiménez-Mejías et al., 2016c; Martín-Bravo et al., 2019; Semmouri et al., 2019; Villaverde et al., 2020). The genus Carex has become monophyletic by the inclusion of the formerly recognized segregate genera Cymophyllus Mack., Kobresia Willd., Schoenoxiphium Nees, and Uncinia Pers. (Global Carex Group, 2015). This taxonomic decision agrees with our results (Fig. 2C). The previously cited genera were the only ones included in the most recent treatments of the tribe (e.g., Kükenthal, 1909; Egorova, 1999; Ball & Reznicek, 2002), whose circumscription remains otherwise unaltered. Although the topology within Carex largely reflects recent studies focused on Carex (e.g., Villaverde et al., 2020), the placement of some species such as Carex ncinate L.f.is not well supported, and for a deeper systematic analysis of the genus Carex, we refer to those studies. Data of Carex species used in this study were generated using three different targeted sequencing probe kits: Angiosperms353 (Johnson et al., 2019), Cyperaceae-specific (Villaverde et al., 2020), and Angiosperms I kit for Anchored Phylogenomics (Léveillé-Bourret et al., 2018c). Lower recovery of the Angiosperms353 genes from data generated with the other probe kits may have contributed to lower resolution within Carex in this study.

4.3.9 Abildgaardieae–Eleocharideae–Fuireneae–Cypereae Clade

The Abildgaardieae–Eleocharideae–Fuireneae–Cypereae Clade (FAEC Clade) has long been retrieved in molecular studies of the Cyperaceae family (Simpson et al., 2007; Muasya et al., 1998, 2009a; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Spalink et al., 2016b; Semmouri et al., 2019). The relationships between the lineages have not always been clear; however, most studies recovered an Abildgaardieae–Eleocharideae Clade, a Fuireneae s.l. grade, and a monophyletic tribe Cypereae (e.g., Hinchliff & Roalson, 2013; Jung & Choi, 2013; Spalink et al., 2016b; Semmouri et al., 2019; this study). Within the FAEC Clade, the Fuireneae s.l. grade and tribe Cypereae are characterized by embryos with horizontal germ pores (Léveillé-Bourret & Starr, 2019).

4.3.9.1 Abildgaardieae—**Eleocharideae Clade.** The monophyly of a clade containing tribe Abildgaardieae and *Eleocharis* R.Br. is well supported in recent phylogenetic studies (Hinchliff & Roalson, 2013; Jung & Choi, 2013; Spalink et al., 2016b; Semmouri et al., 2019; Larridon et al., 2021c), and in this study (Fig. 2D).

Tribe Eleocharideae. Tribe Eleocharideae (Fig. 4F) sensu Goetghebeur (1986) included Eleocharis and three monotypic genera (Chillania Roiv., Egleria L.T.Eiten, and Websteria S.H.Wright), whereas Goetghebeur (1998) recognized Eleocharis, Egleria, and Websteria. On the basis of morphological data (e.g., an Eleocharis-type embryo) and molecular evidence, these monotypic genera have been combined into Eleocharis, resulting in a monogeneric tribe (Roalson & Friar, 2000; Hinchliff et al., 2010; Roalson et al., 2010; Hinchliff & Roalson, 2013; Semmouri et al., 2019). Our results support a monophyletic and monogeneric Eleocharideae (Fig. 2D). However, within the species-rich genus Eleocharis, there are significant problems with the infrageneric classification as most of the currently recognized sections and series (González-Elizondo & Peterson, 1997) are not monophyletic (Roalson et al., 2010).

Tribe Abildgaardieae. The monophyly of tribe Abildgaardieae (Fig. 7A) is well supported (Muasya et al., 2009a; Hinchliff & Roalson, 2013; Spalink et al., 2016b; Semmouri et al., 2019; Larridon et al., 2021c; this study, Fig. 2D), as is the monophyly of both clades within this tribe: the Bulbostylis Clade and the Fimbristylis Clade (Muasya et al., 2009a; Jung & Choi, 2013; Hinchliff & Roalson, 2013; Spalink et al., 2016b; Roalson et al., 2019; Semmouri et al., 2019; Larridon et al., 2021c; this study). Until the inclusion of Nemum Desv. in Bulbostylis Kunth (Roalson et al., 2018, 2019; Larridon et al., 2019b), Bulbostylis was paraphyletic (Simpson et al., 2007; Muasya et al., 2009a; Hinchliff & Roalson, 2013; Spalink et al., 2016b; Reutemann et al., 2018; Semmouri et al., 2019; Larridon et al., 2021c).

The isolated position of Fimbristylis hygrophila Gordon-Gray and F. variegata Gordon-Gray (Hinchliff & Roalson, 2013; Spalink et al., 2016b; Semmouri et al., 2019; Muasya et al., 2020; Larridon et al., 2021c), sister to the rest of the Fimbristylis Clade, recently led to the publication of these species as the new genus Zulustylis Muasya (Muasya et al., 2020). The Fimbristylis Clade also includes the genera Actinoschoenus Benth., Arthrostylis R.Br., and Trachystylis S.T.Blake (Muasya et al., 2009a; Hinchliff & Roalson, 2013; Spalink et al., 2016b, Semmouri et al., 2019;

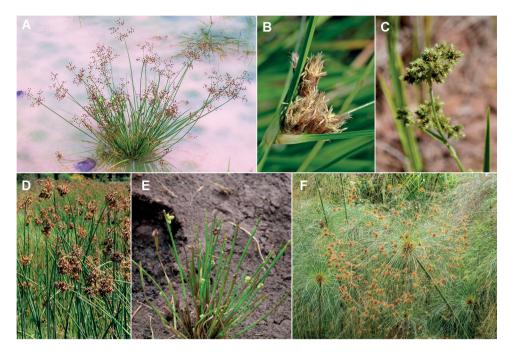


Fig. 7. Morphological diversity of the Cyperaceae tribes. **A,** Abildgaardieae, *Fimbristylis* Vahl sp. **B,** Bolboschoeneae, Bolboschoenus caldwellii (V.J.Cook) Soják. **C,** Fuireneae, *Fuirena umbellata* Rottb. **D,** Schoenoplecteae, *Schoenoplectus tabernaemontani* (C.C.Gmel.) Palla. **E,** Pseudoschoeneae, *Schoenoplectiella senegalensis* (Steud.) Lye. **F,** Cypereae: Cyperinae, Cyperus papyrus L. Photos **A, B, D,** and **F** by Russell Barrett; **C** and **E** by Isabel Larridon.

Muasya et al., 2020; Larridon et al., 2021c). These genera had been regarded as constituting the "Arthrostylideae" (Goetghebeur, 1986; Bruhl, 1995). They were later placed in Schoeneae (Goetghebeur, 1998) and transferred to tribe Abildgaardieae by Muasya et al. (2009a). Larridon et al. (2021c) showed that the "Arthrostylideae" genera form two clades: (i) a clade of Australian species encompassing Arthrostylis, Trachystylis, and a new genus Scleroschoenus K.L.Wilson, J.J.Bruhl & R.L.Barrett published to place several species recently described in Actinoschoenus (Rye et al., 2015); and (ii) a clade of Actinoschoenus s.s. These species-poor lineages within the Fimbristylis Clade are characterized by (i) the C₃ photosynthetic pathway and (ii) Carex- or Schoenus-type embryos, whereas the remaining lineages use the C₄ photosynthetic pathway and have Abildgaardia-, Fimbristylis-, and Tylocarya-type embryos (Semmouri et al., 2019; Larridon et al., 2021c). The genus Abildgaardia has been treated in a variety of ways, but Larridon et al. (2021c) circumscribe it as a separate genus sister to Fimbristylis. Embryo morphology data, in agreement with phylogenetic data, support the recognition of the genus Abildgaardia, as the latter differs from Fimbristylis by having an Abildgaardia-type embryo (Semmouri et al., 2019). Crosslandia W.Fitzg. has been clearly established as being nested within Fimbristylis (Hinchliff & Roalson, 2013; Spalink et al., 2016b; Semmouri et al., 2019; Roalson et al., 2019; Larridon et al., 2021c), and a new combination has been published to move it into Fimbristylis (Roalson et al., 2019).

4.3.9.2 Fuireneae s.l. grade. Here and in the more recent molecular studies, tribe Fuireneae s.l. (which includes tribes Bolboschoeneae, Fuireneae s.s., Schoenoplecteae, and Pseudoschoeneae sensu Starr et al., 2021) has been retrieved as a grade

toward branching off after the Cypereae, Abildgaardieae-Eleocharideae Clade (Escudero & Hipp, 2013; Hinchliff & Roalson, 2013; Spalink et al., 2016b; Glon et al., 2017; Semmouri et al., 2019; Starr et al., 2021; this study, Fig. 2E). Fuireneae s.l. is not monophyletic, but falls apart into three (Semmouri et al., 2019) or four clades (Muasya et al., 2009a; Escudero & Hipp, 2013; Hinchliff & Roalson, 2013; Glon et al., 2017). Within the Fuireneae grade, a first clade of Bolboschoenus (Asch.) Palla branches off first, followed by a clade of Fuirena Rottb., with a third clade consisting of a Schoenoplectiella Lye and Pseudoschoenus (C.B.Clarke) Oteng-Yeb., and a fourth clade of Schoenoplectus (Rchb.) Palla and Actinoscirpus (Ohwi) R.W.Haines & Lye clade being sister to Cypereae in Glon et al. (2017) and Spalink et al. (2016b). In Escudero & Hipp (2013), the third and fourth clades were inverted in position, whereas in Semmouri et al. (2019), the third and fourth clades form a clade together, sister to Cypereae. Starr et al. (2021) enriched samples of more than a third of Fuireneae s.l. using the Angiosperms353 probes. Their results strongly support the recognition of six genera and four major Fuireneae lineages that they recognize as tribes (Starr et al., 2021), that is, tribe Bolboschoeneae (Fig. 7B), tribe Fuireneae s.s. (Fig. 7C), tribe Schoenoplecteae (Figs. 4G, 7D), and tribe Pseudoschoeneae (Fig. 7E), a conclusion supported by this study. At the generic level, using both Angiosperms353 and nrDNA data sets, Starr et al. (2021) show that most sub-Saharan African perennial species previously treated in Schoenoplectus needed to be transferred to Schoenoplectiella to make both genera monophyletic.

4.3.9.3 Tribe Cypereae. On the basis of molecular data (Muasya et al., 2002, 2009a), two clades are recognized in tribe Cypereae. Most species of the Ficinia Clade (Fig. 4H) are

characterized by spikelets with spirally arranged glumes, whereas most species of the Cyperus Clade (Fig. 7F) usually have spikelets with distichously arranged glumes (Muasya et al., 2009a, 2009b; Larridon et al., 2011a). Although commonly referred to as the Ficinia Clade and Cyperus Clade, subtribal names exist for these groupings that we accept to use in Section 5 (Ficiniinae and Cyperinae; Fig. 3).

In the Ficinia Clade, Erioscirpus Palla is the first genus to diverge, before Scirpoides Ség (Fig. 2F). Erioscirpus was previously thought to be more allied to Scirpus and Eriophorum, but molecular studies (Yano et al., 2012; García-Madrid et al., 2015; Muasya & Larridon, 2021) supported its inclusion in the Ficinia Clade. The decision of Reid et al. (2017) to combine Karinia mexicana (C.B.Clarke ex Britton) Reznicek & McVaugh into Scirpoides Ség. (as Scirpoides mexicanus (Reznicek & McVaugh) Goetghebeur ex C.S. Reid and J.R. Carter), based on molecular and morphological evidence, was supported by Semmouri et al. (2019). Scirpoides mexicanus was not sampled in this study. Semmouri et al. (2019) strongly supported a sister relationship between Ficinia Schrad. and Isolepis R.Br. However, several other studies including a larger species sampling have shown that Isolepis is paraphyletic and includes Ficinia (Muasya et al., 2009b; Muasya & de Lange, 2010; Hinchliff & Roalson, 2013; Spalink et al., 2016b). Recently, Muasya & Larridon (2021) sampled 78% of the Ficinia Clade for a nuclear data set including ETS and ITS and in a chloroplast data set including the genes matK, ndhF, rbcL, and rps16, the trnL intron, and trnL-F spacer with the aim to recircumscribe Ficinia and Isolepis as monophyletic genera. On the basis of the topology obtained with their nuclear data set, Muasya & Larridon (2021) (i) broadened the circumscription of Ficinia to include the annual Isolepis species characterized by cartilaginous glumes and including all Isolepis species retrieved outside the core Isolepis clade, and (ii) narrowed the circumscription of Isolepis to encompass only those species retrieved as part of the core Isolepis clade. Two southern African genera that were recently described, Afroscirpoides García-Madr. & Muasya (García-Madrid et al., 2015) and Dracoscirpoides Muasya (Muasya et al., 2012), form a clade in this study (Fig. 2F). Species segregated into Dracoscirpoides and Hellmuthia are atypical for tribe Cypereae, all bearing perianth, and were originally described as part of Scirpus. Hellmuthia is strongly supported as sister to a clade including Ficinia and Isolepis (e.g., Simpson et al., 2007; Hinchliff & Roalson, 2013; Jung & Choi, 2013; Semmouri et al., 2019). The scale-like perianth of Hellmuthia, interpreted to be analogous to similar structures in Chrysitricheae (subfamily Mapanioideae, here as tribe Hypolytreae s.l.) by Haines & Lye (1976), is now thought to be ontogenetically similar to perianth in other Cyperaceae (Vrijdaghs et al., 2006; Muasya et al., 2009b).

Until recently, the Cyperus Clade included two genera, that is, Cyperus L. (962 species; Govaerts et al., 2020) and Androtrichum (Brongn.) Brongn. (2 species; Govaerts et al., 2020). Thirteen segregate genera recognized by Goetghebeur (1998), that is, Courtoisina Soják, Kyllingiella R.W.Haines & Lye, and Oxycaryum Nees (C₃ photosynthesis), plus Alinula J.Raynal, Ascolepis Nees ex Steud., Ascopholis C.E.C.Fisch., Kyllinga Rottb., Lipocarpha R.Br., Pycreus P.Beauv., Queenslandiella Domin, Remirea Aubl.,

Sphaerocyperus Lye, and Volkiella Merxm. & Czech (C₄ photosynthesis), had already since been synonymized with Cyperus (Larridon et al., 2011a, 2011b, 2013, 2014; Bauters et al., 2014). The small genus Androtrichum (C₂ photosynthesis) had not yet been combined into Cyperus due to a lack of data (only rbcL sequences were available for Androtrichum) and conflicting results (Muasya et al., 2009a; Hinchliff & Roalson, 2013; Jung & Choi, 2013). Recently, Pereira-Silva et al. (2020) sank Androtrichum into Cyperus, based on the results of Semmouri et al. (2019). Our results confirm the placement of the two species previously placed in Androtrichum, that is, Cyperus byssaceus Pereira-Silva and C. trigynus Spreng., among the C3 lineages of Cyperus (Fig. 2F). Previous studies based on a limited set of chloroplast and nuclear ribosomal DNA (nrDNA) markers were able to resolve relationships between species of Cyperus using C₃ photosynthesis (Larridon et al., 2011a, 2011b), but not between sections and species using C₄ photosynthesis (Bauters et al., 2014; Larridon et al., 2013, 2014). Owing to the massive size of the genus and the apparent fast radiation of the C4 Cyperus Clade (Spalink et al., 2016b), a comprehensive and robust phylogeny for Cyperus is not easily accomplished. Recently, Larridon et al. (2020) tested the ability of two enrichment probe kits to resolve low-level relationships in the C₄ Cyperus Clade, that is, the universal Angiosperms353 probes (Johnson et al., 2019) and Cyperaceae-specific probes (Villaverde et al., 2020). As in previous studies (e.g., Larridon et al., 2011a, 2013), they found that species of Cyperus section Amabiles, for example, Cyperus cuspidatus Kunth and C. amabilis Vahl, are sister to all other taxa in the C₄ Cyperus Clade. Interestingly, after Cyperus section Amabiles, a clade of white-glumed Cyperus species, that is, margaritaceus-C. niveus complex, was retrieved as sister to the rest of the C4 Cyperus species. Here, a similar topology is obtained (Fig. 2F) based on an analysis of a wider sampling of C₄ Cyperus Clade species.

5 Taxonomic Treatment

5.1 Family Cyperaceae

Cyperaceae Juss. Gen. Pl. [Jussieu] 26. 1789, nom. cons. Type Cyperus L.

Description (adapted from POWO, 2020): Perennial or annual herbs, terrestrial, rarely submerged aquatics, or scandent; perennial species rhizomatous, stoloniferous, bulbous, or sub-bulbous; annual species generally caespitose. Some genera have a caudex (e.g., Microdracoides). Culms usually trigonous to triquetrous, less often terete, flattened (e.g., Fimbristylis spp.), polygonal (e.g., Fuirena umbellata, Schoenoplectiella heptangularis), or irregular, often with prominent ribs. Leaves basal or basal and cauline, generally tristichous with a closed sheath; blade usually linear, glabrous, scabrous, or hairy, with central midrib prominent, sometimes with an expanded 3-veined blade (Hypolytrum, Mapania) or elliptic and constricted into a pseudopetiolate base (e.g., Mapania spp., Carex spp.); ligules present or absent. Inflorescence bracts usually present, leaf-like, bristlelike, or glume-like. Inflorescence simple to compound, lax to condensed, and usually highly branched, paniculate, cor-

ymbose, spicate, or capitate, comprising 1 to many ultimate inflorescence units, very rarely reduced to a single flower (e.g., Eleocharis confervoides), the inflorescence units either indeterminate (= spikelets) or, in a few genera, determinate (= spicoids). In subfamily Cyperoideae, spikelets few to many, sometimes reduced to a single spikelet (e.g., Eleocharis), terminal or lateral comprising 1-many scales, usually spirally arranged, or sometimes distichous; flowers bisexual, unisexual, or both types present. In subfamily Mapanioideae, spicoids few to many, terminal, with 2-13(-100) membranous scales on a much reduced axis, the lowest 2 scales opposite (sometimes fused), keeled, longer than the flowers within, the whole spicoid subtended and usually hidden by a larger scale-like bract; flowers unisexual and comprising a single naked stamen or pistil. Perianth absent or of 1-many bristles or scales. Stamens 1-3 (up to 6 in Cyathocoma and up to 8(-9) in Morelotia). Ovary superior, trimerous, or dimerous (extremely rarely tetramerous; e.g., Carex concinnoides), locule 1, ovule 1; style short to elongate, base sometimes thickened persistent or not; stigma usually 2–3-branched, rarely 1 (e.g., Rhynchospora spp.). Fruit a nutlet, rarely drupe-like (e.g., Cladium), sometimes accompanied by a cupule and hypogynium (e.g., Scleria).

Includes 24 tribes, 10 subtribes, 95 genera, 5796 species. **Distribution** Cosmopolitan

Key to the subfamilies of Cyperaceae

5.2 Subfamily Mapanioideae

Mapanioideae C.B.Clarke in W.H. Harvey & O.W. Sonder (ed. W.T. Thiselton-Dyer), Fl. Cap. 7: 150. (1897).

Type Mapania Aubl.

Diagnosis Characterized by a different morphology of the flowering units compared with Cyperoideae.

Includes 2 tribes, 11 genera, 185 species.

Distribution Mainly tropical and austral temperate.

Key to the tribes of subfamily Mapanioideae

5.2.1 Tribe Chrysitricheae

Chrysitricheae Nees, Linnaea 10: 144. (1835).

Type Chrysitrix L.

Diagnosis Chrysitricheae are robust, rhizomatous, or stoloniferous perennials. The leaves are linear or reduced to basal sheaths. The inflorescence bracts are leaf-like to culm-like and continuous with the culm. The inflorescences are paniculate, anthelate, capitate, or reduced to a single spike. The basic reproductive units comprise spicoids.

The spicoids each have 4–100 (or more) floral bracts and the lowest two bracts are opposite and keeled or, in *Chrysitrix*, spirally arranged. The floral bracts may or may not subtend a single stamen and each spicoid is terminated by a single pistil. The spicoids are subtended and usually hidden by glume-like spicoid bracts and these units are aggregated into spikes that are analogous to spikelets in Cyperoideae genera. The pollen is pyriform, with the exception of *Diplasia* in which it is spheroidal. The style is 2–3-fid and the fruits are 2–3-sided or terete with a hard exocarp.

Accepted genera 6; Capitularina J.Kern. (1 sp.), Chorizandra R.Br. (6 spp)., Chrysitrix L. (4 spp.), Diplasia Pers. (1 sp.), Exocarya Benth. (1 sp.), and Lepironia Pers. (1 sp.). For descriptions and notes on the genera, see Goetghebeur (1998).

Distribution Chrysitricheae mainly have a southern hemisphere (Gondwanan) distribution with the exception of *Diplasia*, which is present in Trinidad and Central and tropical South America. They are found in open swamps or forest.

Key to the genera of tribe Chrysitricheae

- 2a. Leaf blade absent; inflorescence always a single spike... 3
- 2b. Leaf blade present; inflorescence capitate, anthelate, or rarely a single spike4
- 3a. Spikes fusiform. Madagascar to Polynesia Lepironia

- 4b. Culm not septate; inflorescence anthelate.....5

5.2.2 Tribe Hypolytreae

Hypolytreae Nees ex Wight & Arn., Contr. Bot. India: 69 (1834).

Type Hypolytrum Pers.

Diagnosis Hypolytreae are rather delicate to very robust (up to 5 m tall), rhizomatous or stoloniferous perennials. The leaves are linear or sometimes with an expanded, linearoblong to broadly oblong blade and pseudopetiole between the blade and sheath, or rarely reduced to bladeless sheaths. The inflorescence bracts are leaf-like to glume-like. The inflorescences are paniculate, capitate, or reduced to a single spike, rarely anthelate. The basic reproductive units comprise spicoids. The spicoids each have 4–15 floral bracts and the lowest two bracts are opposite and keeled. The lowest two bracts usually subtend a single stamen, whereas the remaining floral bracts may or may not subtend a single stamen and each spicoid is terminated by a single pistil. The spicoids are subtended and usually hidden by glume-like spicoid bracts, and these units are aggregated into spikes that are analogous to spikelets in non-mapiniid genera. The pollen is spheroidal. The style is 2-3-fid and the

fruits are 2–3-sided or terete with a hard, succulent or occasionally berry-like exocarp.

Accepted genera 4; Hypolytrum Pers. (63 spp.; including the formerly recognized segregate genus *Principina* Uittien), *Mapania* Aubl. (100 spp.), *Paramapania* Uittien (7 spp.), and *Scirpodendron* Zipp. ex Kurz (2 spp.). For descriptions and notes on the genera, see Simpson (1992), Goetghebeur (1998), and Alves (2015).

Distribution Hypolytreae have a pantropical distribution and occur primarily in forest or forest margins, rarely in savannah.

Key to the genera of tribe Hypolytreae

- 2a. Floral bracts 2(-3) Hypolytrum
- 2b. Floral bracts 4-63

5.3 Subfamily Cyperoideae

Cyperoideae Beilschm. in Flora 16 (Bieb. 7): 52, 106. (1833). Type Cyperus L.

Diagnosis Cyperoideae flowering units lack the pair of lateral, opposing and keeled floral bracts found in Mapanioideae.

Includes 22 tribes, 85 genera, 5488 species. **Distribution** Cosmopolitan

Key to the tribes of Cyperoideae

- 2a. Pistillate flower with hairy perianth scales or bristles3
- 2b. Pistillate flower without perianth, sometimes with hypogynium4

- 4b. Glumes distichous; pistil subtended directly by a single scale-like glume (or in Bisboeckelereae enclosed in a sac-like organ or within 2 opposite scales, but then pistil seemingly terminal on axis, and fruit seated on a hypogynium); embryo Juncus- or Fimbristylis-type5

- 7a. Many prophylls containing a flower.....8
- 7b. All prophylls sterile.....9

- 9b. Pistils >2 per spikelet; glume wings not or scarcely enveloping the flower of the node below; embryo various......14

- 11a. Style 2-fid, fruit flattened, straight; style base enlarged, persistent as a tubercle on fruit Rhynchosporeae

14b. Flowers bisexual, or if unisexual then never with the above combination of characters15 15a. Style base distinct and often thickened or fimbriate, persistent on the fruit or deciduous with the style16 15b. Style base neither distinct nor thickened17 16a. Leaf blade usually present; inflorescence often corymbose or anthelate, rarely unispicate but then with one involucral bract larger than glumes; perianth absent; embryo not mushroom-shaped, Abildgaardia-, Bulbostylis-, or Fimbristylis-type Abildgaardieae (in part) 16b. Leaf reduced to a bladeless sheath, or blade sometimes evident as a tiny mucro; inflorescence unispicate, without larger involucral bract; perianth usually present; embryo mushroom-shaped, Eleocharis- or Websteriatype Eleocharideae 17a. Embryo with basal or lateral root cap, and perpendicular 17b. Embryo with a lateral root cap and parallel germ pore20 18a. Perianth bristles 6–12, antrorsely scabrous their whole length; ligule and glume ciliate; all glumes fertile 18b. Not this combination of characters19 19a. Spikelet with 1–5 sterile proximal glumes, rarely all fertile but then basal glume with longer mucro, 1.5-4 mm wide; perianth parts o-6; perianth barbs antrorse or divaricate when present; cauline leaves absent; embryo with basal root cap, Carex-type Trichophoreae 19b. Spikelet with all glumes fertile, or rarely with sterile proximal glumes but then basal glume not with longer mucro, 6–15+ mm wide; perianth bristles >10 (Eriophorum p.p.); perianth barbs retrorse when present; cauline leaves present or absent; embryo with lateral root cap, Schoenus- or Fimbristylis-type 20a. Glumes often distichous; perianth usually absent; usually with bladed leaves all basal; embryo not mushroomshaped, Cyperus-typeCypereae 20b. Glumes never distichous; perianth usually present; often bladeless, or bladed with leaves basal and cauline; mushroom-shaped, Bolboschoenus-Schoenoplectus-type21 21a. Embryo 176–305(–382 μm) long, with scutellum 39%–60% of total embryo length; bracts sheathing, leaf-like, rarely cusp-like; leaves with well-developed blades, hairy at least at the junction of blade and sheath, rarely glabrous when the blade is reduced to a mucronate sheath Fuireneae 21b. Embryo 315–1269 μm long, when <380 μm, scutellum is 28%-32% of total embryo length; bracts sheathless, leaflike or appearing to be a continuation of the stem; leaves well developed or reduced to sheaths, glabrous22 22a. Lowermost primary bract leaf-like with spikelets 10-40 mm long; embryo with three primordial leaves, notch below the root cap presentBolboschoeneae 22b. Lowermost primary bract patent to erect, but stem-like, when leaf-like, patent to reflexed with spikelets to 5 mm long; embryo with two primordial leaves, notch below root

cap absent......23

- 23b. Embryo scutellum umbonate or distinctly pileate; nutlet epidermal cells linear, (8.0–)9.2–20.2 times longer than wide, rarely isodiametric to oblong, 1.5–3.8 times longer than wide; nutlet surface smooth or transversely rugose; basal flowers sometimes present Pseudoschoeneae

5.3.1 Tribe Trilepideae

Trilepideae Goetgh. in Taxon 34: 629. (1985)

Type Trilepis Nees

Diagnosis Trilepideae are characterized by a panicle composed of many dense spikes of many tiny spikelets with few distichous glumes, unisexual flowers, a perianth usually formed by 3 fimbriate scales opposite the flat sides of the nutlet, and a *Trilepis*-type embryo (Goetghebeur, 1985; 1998; Semmouri et al., 2019).

Accepted genera 4; Afrotrilepis (Gilly) J.Raynal (2 spp.), Coleochloa Gilly (8 spp.), Microdracoides Hua (1 sp.), and Trilepis Nees (5 spp.). For descriptions and notes on the genera, see Goetghebeur (1998).

Distribution West and West Central Africa (Afrotrilepis, Microdracoides), Tropical and southern Africa and Madagascar (Coleochloa), northern South America to Brazil (Trilepis). Occurring in tropical areas mostly on inselbergs, growing on shallow soils; one species epiphytic in submontane tropical rain forest.

Key to the genera of Trilepideae (based on Goetghebeur, 1998)

- 3b. Leaves eligulate; spikes always unisexual Trilepis

5.3.2 Tribe Cladieae

Cladieae Nees, Linnaea 9: 297. 1834.

Type Cladium P.Browne

Diagnosis (Semmouri et al., 2019) Herbs perennial, rhizomatous, and stoloniferous; culms few-noded, hollow; leaves eligulate, following a 1/3 phyllotaxis; inflorescence paniculate, with many spikelets, primary bracts leaflike, sheathing; spikelets with few to many, spirally arranged persistent glumes, lower flower mostly functionally male, upper flowers bisexual with two stamens, bristles absent, nutlet ovoid, with a thick corky beak, surface smooth to wrinkled, embryo small and poorly developed, broadly obovate in outline, with a basal, poorly developed root cap and without a leaf primordium (*Juncus*-type embryo).

Accepted genus *Cladium* P.Browne (3 spp.). For a description and notes on the genus, see Goetghebeur (1998).

Distribution *Cladium* is subcosmopolitan and occurs in swamps and marshes, often in brackish or calcareous habitats.

5.3.3 Tribe Bisboeckelereae

Bisboeckelereae Pax in H.G.A. Engler & K.A.E. Prantl, Nat. Pflanzenfam. Nachtr.: 48. (1897).

Type Bisboeckelera Kuntze

Diagnosis Bisboeckelereae are recognized by the (sometimes connate) empty glumes surrounding the apparently terminal female flower, male spikelets with glumes each with a single stamen, and *Carex-*, *Schoenus-*, or *Fimbristylis-*type embryos (Goetghebeur, 1998; Semmouri et al., 2019).

Accepted genera 4; Becquerelia Brongn. (6 spp.), Bisboeckelera Kuntze (4 spp.), Calyptrocarya Nees (8 spp.), and Diplacrum R.Br. (10 spp.). For descriptions and notes on the genera, see Goetghebeur (1998).

Distribution Tropical America (Becquerelia, Calyptrocarya), southern Tropical South America (Bisboeckelera), Tropics and subtropics (Diplacrum).

Key to the genera of Bisboeckelereae (based on Goetghebeur, 1998)

- 1b. Nutlets rather tightly enclosed by either a shortly pubescent or a many-nerved utriculiform glume.......3

5.3.4 Tribe Sclerieae

Sclerieae Wight & Arn., Contr. Bot. India: 71. (1834)

Type Scleria P.J.Bergius

Diagnosis Sclerieae have a basically paniculate inflorescence, the spikelets are bisexual or unisexual, flowers unisexual, the nutlet is surrounded at the base by a hypogynium and a cupula (sometimes reduced), and a *Fimbristylis*-type embryo (Goetghebeur, 1998; Semmouri et al., 2019).

Accepted genus Scleria P.J.Bergius (258 spp.). For a description and notes on the genus, see Goetghebeur (1998). For infrageneric classification and recent taxonomic treatments, see Bauters et al. (2016, 2018, 2019), Bauters (2018), and Galán Díaz et al. (2019).

Distribution Tropics and subtropics to North America.

5.3.5 Tribe Carpheae

Carpheae Semmouri & Larridon, Bot. Rev. 85: 33 (2019).

Type Carpha Banks & Sol. ex R.Br.

Diagnosis (Semmouri et al., 2019) Tribe Carpheae is characterized by the *Carpha*-type embryo, whereas *Carex*-and *Schoenus/Helothrix*-type embryos are prevalent in tribe

Schoeneae. Anthers are typically conspicuously greenish-yellow in this tribe, whereas they are yellow to red-colored in the morphologically similar Schoeneae. Many species of tribe Schoeneae mainly occur in austral temperate dryland habitats that are only seasonally damp (e.g., woodland and heathland), whereas Carpheae occur typically in wetlands and damp areas.

Accepted genera 2; Carpha Banks & Sol. ex R.Br. (15 spp.), and Trianoptiles Fenzl ex Endl. (3 spp.). For descriptions and notes on the genera, see Goetghebeur (1998).

Distribution Whereas the annual *Trianoptiles* species are endemic to the wetlands of South Africa (SW Cape), perennial *Carpha* occurs in swamps and along stream sides in the southern and central African mountains, Madagascar, Mascarenes, New Guinea, southern Japan, southeastern Australia, New Zealand, and Chile.

Key to the genera of Carpheae (based on Goetghebeur, 1998)

5.3.6 Tribe Cryptangieae

Cryptangieae Benth. in J. Linn. Soc. London, Bot. 18: 366. (1881).

Type Cryptangium Schrad. ex Nees

Diagnosis Cryptangieae are mostly characterized by unisexual spikelets (except for *Koyamaea*, with a more basal single female flower and many male flowers above), spirally arranged glumes (distichously arranged in *Exochogyne*), fruit usually triangular or trigonous in cross-section with three fimbriate perianth scales opposite the flat sides of the nutlet (biconvex and without hypogynous scales in *Exochogyne*), and *Juncus-* or *Carex-*-type embryos, although few species have been studied (Goetghebeur, 1998; Semmouri et al., 2019). It seems that all species present a red-pinkish style and stigma, except for some populations in the "campos rupestres" of Chapada Diamantina localities (Bahia, Brazil).

Accepted genera 7; Cephalocarpus Nees (20 spp.), Cryptangium Schrad. ex Nees (1 spp.), Didymiandrum Gilly (1 spp.), Exochogyne C.B.Clarke (2 spp.), Koyamaea W.W.Thomas & Davidse (1 sp.), Krenakia S.M.Costa (10 spp.), and Lagenocarpus Nees (15 spp.). For descriptions and notes on the genera, see Costa et al. (2021a).

Distribution Tropical America, in forested (Koyamaea, Didymiandrum) and open vegetation, mostly at sandy nutrient-poor soils and/or rocky places, from seashores and sandy temporarily wet plains (such as the Amazonian "campinaranas") to high altitudes (such as the "tepuis" and "campos rupestres"). Also, in some mountains associated with the Andes, but with older and nutrient-poor soils, such as the Cordillera del Condor (Ecuador, Peru).

Key to the genera of Cryptangieae (modified from Costa et al., 2021a)

	Pierce I will also with a baselone of Course Course
1a.	Bisexual spikelets with a basalmost female flower and many male flowers above
1b.	Unisexual spikelets only
2a.	Inflorescence spike-like; glumes distichously arranged;
24.	nutlet biconvex in cross-section Exochogyne
2b.	Inflorescence raceme/panicle-like or head-like; glumes
	spirally arranged; nutlet triangular or trigonous in cross-
	section3
3a.	Inflorescence lateral; hypogynous scales with fimbriate
	margins and glabrous dorsal surface Cephalocarpus
3b.	Inflorescence terminal; hypogynous scales inconspicuous
	and/or glabrous or with sparsely short-ciliate margins or
	pubescent at margin and dorsal surface4
4a.	Male spikelets with (1–)4–5 stamens; nutlet trigonous
	(trisulcate or inconspicuously ridged) Lagenocarpus
4b.	1 (3)
	sharp ridges5
5a.	Dioecious herbs; involucral bracts (sterile or fertile)
	solitary or forming pseudowhorls with elliptic to oblong-
	elliptic blades Didymiandrum
5b.	Monoecious herbs; involucral bracts (sterile or fertile)
_	never forming pseudowhorls6
6a.	Leaf with a developed blade, rarely reduced to sheath;
	female spikelets with 1(-2) flowers and 3 glumes;
	nutlet with three conspicuous cavities at the base
6h	Cryptangium Leaf always reduced to sheath; female spikelets with 1–5
ob.	flowers and 3–7 glumes; nutlet cavities absent or
	inconspicuous
	inconspicuous Ni chakia

5.3.7 Tribe Schoeneae

Schoeneae Dumort., Fl. Belg. 145 (1827).

Type Schoenus L.

Diagnosis Schoeneae are characterized by a (mostly) restricted number of bisexual flowers per spikelet, a \pm well-developed perianth (sometimes absent), spikelets in which the flower(s) are surrounded by the wings of the next glume, and mostly *Carex*- and *Schoenus*-type embryos, although more specialized *Helothrix*-type embryos are also known (Goetghebeur, 1998; Semmouri et al., 2019).

Includes 8 subtribes, 25 genera.

Distribution The tribe has a mostly southern hemisphere distribution, in temperate and subtropical areas, with just a small number of taxa in the northern hemisphere.

Key to the subtribes of Schoeneae

,
1a. Rachilla elongate and flexuose around nutlet(s)2
1b. Rachilla straight3
2a. Leaves ligulate (usually a prominent tuft of hairs);
rhachilla zigzagSchoeninae
2b. Leaves eligulate; rhachilla once-elongate
Tricostulariinae (in part)
3a. Leaves ligulate4
3b. Leaves eligulate9
4a. Inflorescence capitate or subcapitate (lacking spreading
bracts that greatly exceed the inflorescence)
Gymnoschoeninae (in part)
4b. Inflorescence not capitate (may be fan-shaped in
Gahniinae; Mesomelaena, Ptilothrix, but then with
spreading bracts that greatly exceed the inflores-

cence)......5

5a. Glumes distichous6
5b. Glumes spirodistichous8
6a. Inflorescence compact, dense, with two rigid involucral
bracts greatly exceeding the inflorescence (or one bract
in Mesomelaena stygia)
6b. Inflorescence open to dense paniculate, sometimes
reduced, but not as above7
7a. Leaf blades bifacial, well developed (sometimes sen-
escent) Anthelepidinae
7b. Leaf blades conduplicate, unifacial, or highly reduced
Lepidospermatinae (in part)
8a. Ramets distinctly candelabriform
Lepidospermatinae (in part)
8b. Ramets all produced at a similar level11
9a. Perianth inflated, scale-like, persistent on nutlet
Lepidospermatinae (in part)
9b. Perianth absent or bristle-like, usually not persistent on
nutlet
10a. Inflorescence enclosed by two greatly enlarged invo-
lucral bracts
10b. Inflorescence not enclosed by two greatly enlarged
involucral bracts
11b. Glumes distichous (a few glumes spirodistichous in
Cyathocoma hexandra and sometimes in other species)12
12a. Glumes deciduous; plants with leaves well developed
Oreobolinae
12b. Glumes distichous, mostly persistent (deciduous in
Tricostularia, but then leaf blades usually highly reduced)
,

5.3.7.1 Subtribe Anthelepidinae Anthelepidinae R.L.Barrett, subtr. nov.

Type Anthelepis R.L.Barrett, K.L.Wilson & J.J.Bruhl

......Tricostulariinae

Diagnosis Tufted, sometimes rhizomatous, perennial or annual graminoids; leaves mostly basal; culms semi-terete; leaves well developed; ligulate; leaf margins scaberulous or glabrous, flat to channeled; inflorescence terminal, paniculate or subracemose; glumes obscurely distichous, usually deciduous; rhachilla non-flexuous, straight; flowers subtended by upper glumes; lower flower(s) functionally male, upper bisexual; upper glumes longer than lower; spikelets ranging from few to many grouped together in spikelet bundles; 3 stamens, stigma 3-fid; nutlets ranging in shape from narrow-ellipsoid to obovoid; perianth bristles (3)6, shorter or longer than the nutlet.

Accepted genus Anthelepis R.L.Barrett, K.L.Wilson & J.J.Bruhl (4 spp.). For a description and notes on the genus, see Barrett et al. (2019).

Distribution From Sri Lanka to Hainan, New Caledonia to Australia. Whereas *A. undulatus* is widespread, the other three species are localized.

5.3.7.2 Subtribe Caustiinae

Caustiinae R.L.Barrett, subtr. nov.

Type Caustis R.Br.

Diagnosis Tufted, sometimes loosely so, distinctly rhizomatous, perennial graminoids; leaves basal and cauline; culms semi-terete, sometimes ribbed or distinctly grooved, or sometimes trigonous; leaves not or well

developed; eligulate; leaf margins ciliate, scaberulous or glabrous, flat to channeled; inflorescence terminal, spike-like or paniculate; glumes spirally arranged or subdistichous, usually persistent; rhachilla non-flexuous, straight; flowers subtended by upper glumes; lower flower (when present) usually functionally male, sometimes bisexual, upper flower(s) bisexual or sometimes unisexual; upper glumes longer than lower; spikelets ranging from few to many, solitary on often long peduncles or sometimes grouped together in spikelet bundles; 3–6 or 12–20+ stamens, stigma 3–5- or c. 8-fid; nutlets ranging in shape from cylindric, ovoid, or obovoid; perianth bristles absent or 2–3(–5), c. as long as the nutlet.

Accepted genera 2; *Caustis* R.Br. (7 spp.) and *Evandra* R.Br. (2 spp.) plus the unplaced species *Tetraria borneensis* J.Kern. For descriptions and notes on the genera, see Goetghebeur (1998).

Distribution Australia (Caustis and Evandra) and Borneo (Tetraria borneensis).

Key to the genera of Caustiinae

- 1a. Glumes subdistichously arranged; stamens 6; style branches 3; perianth present. Tetraria borneensis
- 2a. Stamens 3–6 or 8; style branches 3–8; leaves reduced to a sheath or sometimes with a blade up to 15 cm long.

 Caustis
- 2b. Stamens 12–20 or more; style branches 8; leaves well developed, up to 90 cm long. Evandra

5.3.7.3 Subtribe Gahniinae

Gahniinae Pax, Bot. Jahrb. Syst. 7: 308 (1886), (as Gahninae) **Type** *Gahnia* J.R.Forst. & G.Forst.

Diagnosis Tufted, rhizomatous, sometimes only shortly so, perennial graminoids; leaves mostly basal or cauline on an erect caudex; culms terete or rarely trigonous, sometimes grooved; leaves well developed or reduced; ligulate; leaf margins scaberulous or glabrous, flat to channeled or filiform; inflorescence terminal, paniculate, or compact and obconical to fan-shaped; glumes spirally arranged or distichous, not deciduous; rhachilla non-flexuous, straight; flowers subtended by upper glumes; lower flower (when present) functionally male, upper bisexual; upper glumes usually longer than lower, but sometimes shorter; spikelets ranging from few to many grouped together in spikelet bundles; 2–6 stamens, stigma 2–5-fid; nutlets ranging in shape from narrow-ellipsoid to narrow-oblong or obovoid; perianth bristles absent or 3, 4, or 5, shorter or longer than the nutlet.

Accepted genera 4; Cyathochaeta Nees (5 spp.), Gahnia J.R.Forst. & G.Forst. (41 spp., plus several undescribed), Mesomelaena Nees (5 spp.), and Ptilothrix K.L.Wilson (1 sp.). For descriptions and notes on the genera, see Wilson (1981, 1993) and Goetghebeur (1998).

Distribution Australia (*Cyathochaeta*, *Mesomelaena*, *Ptilothrix*) and Australasia and the Pacific (*Gahnia*).

Key to the genera of Gahniinae

- 3b. Style base deciduous, shortly pubescent; nut obtusely angled or subterete, seated on a hypogynous disc (except Mesomelaena graciliceps); hypogynous scales slender at first, but at maturity, very much broadened below and enclosing the nut, glabrous below, very shortly antrorsely scabrous aboveMesomelaena

5.3.7.4 Subtribe Gymnoschoeninae

Gymnoschoeninae R.L.Barrett, subtr. nov.

Type Gymnoschoenus Nees

Diagnosis Tufted, robust, rhizomatous, perennial; leaves basal (*Gymnoschoenus*) or cauline on a thick, erect caudex (*Reedia*; and order followed for characters below); culms terete; leaves well developed; ligulate or eligulate; leaf margins pilose or pungently toothed, thickly to thinly lunate; inflorescence terminal, subglobular or elongate and mostly enclosed by two greatly enlarged inflorescence bracts; glumes distichous or spirally arranged to subdistichous, lower glumes persistent or all deciduous; rhachilla nonflexuous, straight; flowers subtended by upper glumes; lower flower functionally male, upper bisexual; upper glumes longer than lower; spikelets many, in spikelet bundles; 3 or 6 stamens, stigma 3-fid; nutlets obovoid to broad-ellipsoid; perianth bristles 0–6, longer than the nutlet.

Accepted genera 2; Gymnoschoenus Nees (2 spp.) and Reedia F.Muell. (1 sp.). For descriptions and notes on the genera, see Goetghebeur (1998).

Distribution Southern Australia.

Key to the genera of Gymnoschoeninae

5.3.7.5 Subtribe Lepidospermatinae

Lepidospermatinae R.L.Barrett, subtr. nov.

Type Lepidosperma Labill.

Diagnosis Tufted (rarely cushion-forming in *Lepidosperma*; or rarely semi-scandent in *Machaerina*), rhizomatous, perennial; leaves mostly basal, sometimes cauline (*Machaerina*), or ramets proliferous (in a few *Lepidosperma*); culms highly variable, commonly biconvex, but ranging from flat to terete or quadrangular or occasionally biconvex; leaves well developed or higly reduced (*Neesenbeckia*, *Netrostylis*; some *Lepidosperma*

and Machaerina); ligulate or eligulate (Netrostylis); leaf margins scaberulous, hispid, ciliate or glabrous, highly variable, commonly biconvex or concavo-convex, but ranging from flat to terete or \pm quadrangular or occasionally biconvex; inflorescence terminal or pseudoaxillary, paniculate, or sometimes appearing spike-like or subracemose (or subcapitate in Neesenbeckia); glumes distichous or spirodistichous, usually not deciduous; rhachilla non-flexuous, straight; flowers 1–5, subtended by upper glumes; lower flower (s) functionally male, upper 1(2) usually bisexual, sometimes functionally male; upper glumes longer than lower (or 2 middle glumes longest in Neesenbeckia); spikelets ranging from few to many grouped together in spikelet bundles; 3 stamens, stigma 3-fid or 6-fid (Neesenbeckia), style base conspicuous in Neesenbeckia and Netrostylis; nutlets ellipsoid to obovoid (sometimes with corky and angular epidermis in Machaerina); perianth (3)6 inflated scales (Lepidosperma) or absent or bristles o-6, shorter or longer than the nutlet.

Accepted genera 4; Lepidosperma Labill. (80 spp., plus many yet to be described species), Machaerina Vahl (55 spp.), Neesenbeckia Levyns (1 sp.), and Netrostylis R.L.Barrett, J.J.Bruhl & K.L.Wilson (11 spp.). For descriptions and notes on the genera, see Goetghebeur (1998), Barrett & Wilson (2012) for Lepidosperma, and Barrett et al. (2021a) concerning Netrostylis.

Distribution: Australia, Pacific, and southeast Asia (*Lepidosperma*, *Netrostylis*), Tanzania, West Indian Ocean to Pacific and tropical America (*Machaerina*), South African Cape Province (*Neesenbeckia*).

Key to the genera of Lepidospermatinae

5.3.7.6 Subtribe Oreobolinae

Oreobolinae R.L.Barrett, subtr. nov.

Type Oreobolus R.Br.

Diagnosis Tufted or cushion-forming, rhizomatous, perennial graminoids; leaves basal or basal and cauline, sometimes cauline on an erect or pseudodendroid caudex; culms terete, elliptical or partially flattened; leaves well developed; eligulate; leaf margins scaberulous or with prickle hairs, flat to thickly lunate; inflorescence terminal, fastigiate paniculate or open paniculate (sometimes a solitary spikelet

or capitate to subcapitate in *Oreobolus*); glumes distichous to spirodistichous, deciduous; rhachilla non-flexuous, straight; flowers subtended by upper glumes; lower flower(s) functionally male (rarely bisexual or absent), upper bisexual (rarely functionally male or female); upper glumes longer than lower; spikelets ranging from few to many in spikelet bundles; usually 3 stamens (6 in *Cyathocoma*), stigma 3-fid; nutlets ranging in shape from ellipsoid to ovoid or obloid; perianth bristles 6 (sometimes not all developing), shorter or longer than the nutlet.

Accepted genera 5; Capeobolus Browning (1 sp.), Chamaedendron (Kük.) Larridon (5 spp.), Costularia C.B.Clarke (15 spp.), Cyathocoma Nees (3 spp.), and Oreobolus R.Br. (17 spp.). For descriptions and notes on the genera, see Goetghebeur (1998), Browning & Gordon-Gray (1999), and Larridon et al. (2018a, 2019).

Distribution South African (*Capeobolus*, *Cyathocoma*), Africa, Madagascar, and West Indian Ocean (*Costularia*), New Caledonia (*Chamaedendron*), Malesia to Australasia and the Hawaiian Islands, and from Costa Rica to the Falkland Islands (*Oreobolus*).

Key to the genera of Oreobolinae

- 3b. Inflorescence a robust panicle of many branches; perianth bristles feathery; 3 stamens......4
- 4a. Leaf sheath only slightly wider than leaf bladeCostularia

5.3.7.7 Subtribe Schoeninae

Schoeninae Fenzl in S.F.L. Endlicher, Gen. Pl.: 114 (1836), (as Schoenoideae).

Type Schoenus L.

Diagnosis Usually tufted (rarely rhizomatous), perennial or annual graminoids; leaves basal; culms usually terete, but also angular or flattened; leaves sometimes reduced to a sheath but usually well developed; usually ligulate; leaf margins serrate, usually terete, but also flat; inflorescence terminal or sometimes pseudolateral; glumes distichous, deciduous; flowers subtended by upper glumes; lower flowers bisexual upper functionally male; upper glumes longer than lower; spikelets ranging from few (e.g., S. fliculmis T.L.Elliott & Muasya) to many grouped together in spikelet bundles; 1–6 stamens, stigma usually 3-fid; nutlets ranging in shape from obovate to rounded trigonous; perianth bristles vary in number from none to many (e.g.,

S. albovaginatus T.L.Elliott & Muasya). Perianth bristles can also be rudimentary or longer than the nutlet (e.g., S. albovaginatus).

Accepted genus Schoenus L. (149 spp.; including some species placed in the formerly recognized genus Epischoenus C.B.Clarke and some species previously placed in Tetraria P.Beauv.). For information on the recircumscription of Schoenus, see Musili et al. (2016), Elliott & Muasya (2017, 2018, 2020), Elliott et al. (2019, 2021), and Barrett et al. (2021b).

Distribution Primarily Australasia and South Africa, with a few species in Europe, the Americas, and Caribbean Islands (Kern, 1974; Viljoen et al., 2013).

5.3.7.8 Subtribe Tricostulariinae

Tricostulariinae R.L.Barrett, K.L.Wilson & J.J.Bruhl, Telopea 24: 74 (2021).

Type Tricostularia Nees ex Lehm.

Diagnosis Tufted, rhizomatous, perennial graminoids; leaves basal and commonly cauline, sometimes on an erect caudex (Ammothryon, some Tetraria and Xyroschoenus); culms trigonous or terete; leaves well developed or much reduced; eligulate; leaf margins scaberulous to denticulate, flat to channeled, sometimes V- or M-shaped in section, sometimes involute or revolute; inflorescence terminal, paniculate, sometimes appearing headlike (Chaetospora, some Tricostularia) or spike-like (some Tricostularia); cladoprophyll present at base of spikelets (except Ammothryon), glumes distichous, mostly persistent (deciduous in Tricostularia); rhachilla mostly non-flexuous, straight (elongate and curved around nutlet in Morelotia affinis, M. gahniiformis, M. octandra, and Xyroschoenus); flowers subtended by upper glumes (except upper glume reduced and infertile in Pacific Morelotia); lower flower(s) variously bisexual fertile, functionally male or bisexual sterile, upper bisexual or functionally male (Morelotia octandra); upper glumes longer than lower; spikelets ranging from few to many in spikelet bundles; mostly 3 stamens ((3)6 in Morelotia australiensis, (4, 6)8(9) in M. octandra), stigma mostly 3-fid (3- or 4-fid in some Tetraria, 3-5-fid in Morelotia octandra); nutlets ranging in shape from ovoid to broad ellipsoid or globose; perianth absent or bristles 2, 6, or 7-8, shorter or longer than the nutlet.

Accepted genera 6; Ammothryon R.L.Barrett, K.L.Wilson & J.J.Bruhl (1 sp.), Chaetospora R.Br. (3 spp.), Morelotia Gaudich. (6 spp.; including the formerly recognized genus Tetrariopsis C.B.Clarke), Tetraria P.Beauv. (39 spp.; including a species previously placed in the formerly recognized genus Epischoenus C.B.Clarke, and some species previously placed in Costularia C.B.Clarke s.l.; Larridon et al., 2017, 2018b), Tricostularia Nees ex Lehm. (11, plus 4 undescribed species), and Xyroschoenus Larridon (1 sp.). For descriptions and notes on the genera, see Goetghebeur (1998); Larridon et al. (2018a): Chamaedendron and Xyroschoenus; Barrett et al. (2020): Chaetospora; Barrett et al. (2021b): all genera in the subtribe.

Distribution Southern Australia (*Tricostularia*), Southern Australia, New Zealand, French Polynesia, Hawaiian Islands (*Morelotia*), New Caledonia to Borneo and New Guinea, South African Cape Floristic Region to tropical Africa (*Tetraria*), the Seychelles (*Xyroschoenus*).

Key to the genera of Tricostulariinae

- 3b. Leaf blades well-formed, 4–100 cm long; involucral bracts leaf-like; inflorescence open, elongate or dense panicle-, head-like, or subglobular, 0.5–1.5 or 5–70 cm long......4
- 4b. Culms with 0–5 nodes; leaf blades (0.5–)1–20 mm wide; inflorescence open, elongate, or dense panicle-like, 5–70 cm long; branchlets erect to spreading......5
- 5b. Nutlets stipitate. Tetraria

5.3.8 Tribe Rhynchosporeae

Rhynchosporeae Nees, Linnaea 9: 294 (1834).

Type Rhynchospora Vahl

Diagnosis Usually small to medium-sized perennials, rarely annuals, inflorescence very variable; glumes spirally arranged (rarely distichous); anthers (1–2) 3, inconspicuous; style 2-branched; perianth bristles usually 3–6, or absent; nutlet usually lenticular to globose, topped with a persistent style base (tubercle).

Embryo top shaped in frontal view, root cap developed in a (sub)basal position, and first leaf primordium developed in a lateral position (*Carex*-type embryo).

Accepted genus *Rhynchospora* Vahl (399 spp.; including the formerly recognized genus: *Pleurostachys* Brongn.; Thomas, 2020).

Distribution Cosmopolitan.

5.3.9 Tribe Dulichieae

Dulichieae W.Schultze-Motel in Willdenowia 2: 173. 14 (1959). **Type** *Dulichium* Pers.

Diagnosis Flowers bisexual, ligule glabrous, spikelet prophyll fertile and squamiform, spikelets distichous on rachis, glume disposition usually distichous at least on terminal spikelet of main stem, all glumes of spikelet fertile, flowers 3–7 per spikelet, perianth setiform, style base

continuous in texture with fruit, leaving a long narrow beak of variable length on fruit, embryo with basal root cap and lateral plumule (Carex-type).

Accepted genera 3; Blysmopsis Oteng-Yeb. (1 sp.), Blysmus Panz. ex Schult. (2 spp.), and Dulichium (1 sp.). For descriptions and notes on Blysmus and Dulichium, see Goetghebeur (1998); and on Blysmopsis, see http://www.efloras.org/florataxon.aspx?flora_id=1&taxon_id=104130).

Distribution Temperate northern hemisphere (*Blysmopsis*, *Blysmus*), Canada to U.S.A. (*Dulichium*).

Key to the genera of Dulichieae

- 1b. Spikelets in a single terminal spike, rarely with an additional lateral spike; perianth of 0–6 bristles2

5.3.10 Tribe Khaosokieae

Khaosokieae Lév.-Bourret & J.R.Starr, Taxon 68: 239 (2019).

Type Khaosokia D.A.Simpson

Diagnosis Dioecious, prophylls sterile, proximal glumes of spikelet sterile, flowers ≥10 per spikelet, perianth of 7 antrorsely scabrous bristles. Embryo not available.

Accepted genus *Khaosokia* D.A.Simpson (1 sp.). For further notes on the genus, see Simpson et al. (2005).

Distribution Endemic to Thailand.

5.3.11 Tribe Calliscirpeae

Calliscirpeae Lév.-Bourret & J.R.Starr, Taxon 68: 238 (2019).

Type Calliscirpus C.N.Gilmour, J.R.Starr & Naczi

Diagnosis Flower bisexual, ligule ciliate, all glumes of spikelet fertile, flowers ≥10 per spikelet, perianth of 6–7(–12) long silky antrorsely scabrous bristles forming a cottony mass at maturity, anthers white or yellowish-white, embryo with

basal root cap and lateral plumule (Carex-type).

Accepted genus *Calliscirpus* C.N.Gilmour, J.R.Starr & Naczi (2 spp.). For further notes on the genus, see Gilmour et al. (2013).

Distribution West Pacific North America (Oregon to California, U.S.A.).

5.3.12 Tribe Scirpeae

Scirpeae T.Lestib. in B.C.J. Dumortier, Fl. Belg.: 143. (1827).

Note: Validated by a reference to [unranked] Scirpeae T. Lestib., Essai Cyper. 22, 39. 29 (1819).

Type Scirpus Tourn. ex L.

Diagnosis Flower bisexual or rarely functionally unisexual with remnant of opposite sex, ligule glabrous or ciliate, prophyll sterile, all glumes of spikelet fertile or rarely 1–12 proximal glumes sterile, flowers (3–)10+ per spikelet, perianth setiform or absent, embryo with lateral

root cap and (sub-)basal plumule (Schoenus-type or Fimbristylis-type).

Accepted genera 6; Amphiscirpus Oteng-Yeb. (1 sp.), Eriophorum L. (18 spp.), Phylloscirpus C.B.Clarke (3 spp.), Rhodoscirpus Léveillé-Bourret, Donadío & J.R.Starr (1 sp.), Scirpus Tourn. ex L. (47 spp.), and Zameioscirpus Dhooge & Goetgh. (3 spp.). For descriptions and notes on the genera, see Novoselova (1994a, b), Goetghebeur (1998), Dhooge et al. (2003), and Léveillé-Bourret et al. (2015).

Distribution Tropical to subarctic northern hemisphere south to southern South America, and Australia through Malesia.

Key to the genera of Scirpeae

- 1b. Cauline leaves absent, leaves all basal with node of the distal leaf hidden in the sheath of the leaf below4

- 4a. Inflorescence, a dense head of several to many spikelets, rarely unispicate; perianth of retrorsely barbed bristles

5.3.13 Tribe Trichophoreae

Trichophoreae Lév.-Bourret & J.R.Starr, Taxon 68: 239 (2019).

Type Trichophorum Pers.

Diagnosis Flower bisexual or rarely functionally unisexual with remnant of opposite sex, ligule glabrous, prophyll sterile, basal (o–)1–9 glumes of spikelet sterile, lowest glume often with conspicuously longer awn than following glumes, flowers 1–10+ per spikelet, perianth setiform, squamiform, or absent, embryo with a basal root cap and lateral plumule (Carex-type).

Accepted genus Trichophorum Pers. (19 spp.; including the formerly recognized genera: Oreobolopsis T.Koyama & Guagl.

and Cypringlea M.T.Strong; Léveillé-Bourret et al., 2020). For notes on the genus, see Léveillé-Bourret et al. (2020).

Distribution Temperate and subtropical northern hemisphere, Andean South America to northwest Argentina.

5.3.14 Tribe Sumatroscirpeae

Sumatroscirpeae Lév.-Bourret & J.R.Starr, Mol. Phyl. Evol. 119 93–104 (2018).

Type Sumatroscirpus Oteng-Yeb.

Diagnosis Flowers bisexual, spikelet prophyll (perigynium) fertile, tubular, spikelets spirally inserted on rachis, glume disposition spiral, sometimes pseudodistichous, all glumes of spikelet fertile, glume wings sometimes partially enveloping the flower of the node below, flowers 7–10+ per spikelet, perianth setiform, style base enlarged, differentiated and persistent on fruit, embryo with basal root cap and lateral plumule (embryo type undetermined).

Accepted genus Sumatroscirpus Oteng-Yeb. (4 spp.). For a description and notes on the genus, see Léveillé-Bourret et al. (2017, 2018b).

Distribution Western China to West Sumatra.

5.3.15 Tribe Cariceae

Cariceae Dumort., Fl. Belg.: 145. (1827).

Type Carex L.

Diagnosis Cariceae is characterized by unisexual flowers, the female ones being enclosed by a prophyll called a perigynium or if closed forming a bottle-like structure, a utricle (see Jiménez-Mejías et al., 2016b), absence of perianth parts, and a *Carex*-type (very rarely a *Schoenus*-type) embryo.

Accepted genus Carex L. (2003 spp.; including the formerly recognized genera: Cymophyllus Mack., Kobresia Willd., Schoenoxiphium Nees, and Uncinia Pers.; Global Carex Group, 2015).

Distribution Cosmopolitan, although absent from Antarctica mainland it is the only Cyperaceae group present in a true Antarctic archipelago (South Georgia; Philcox, 1961; Govaerts et al., 2020; see notes in Jiménez-Mejías & Dorr, 2018).

5.3.16 Tribe Eleocharideae

Eleocharideae Goetgh. in Taxon 34: 629. (1985).

Type Eleocharis R.Br.

Diagnosis (Goetghebeur, 1998): Eleocharideae is characterized by its reduced vegetative morphology, leaves reduced to a sheath (no blade), unispiculate inflorescence, *Eleocharis*-type embryo, and a helio- and helophilous ecology. Characters shared with its sister tribe Abildgaardieae include a differentiated and thickened style base, and moniliform stigmatic hairs. Characters in common with many Fuireneae include a bristle-like perianth, and an embryo with a broadened cotyledon.

Accepted genus Eleocharis R.Br. (c. 302 spp.; including the formerly recognized genera: Chillania Roiv., Egleria L.T.Eiten, and Websteria S.H.Wright; Hinchliff et al., 2010). For a description and notes on the genus, see Goetghebeur (1998) and Hinchliff et al. (2010).

Distribution Cosmopolitan.

5.3.17 Tribe Abildgaardieae Abildgaardieae Lye in Bot. Not. 126: 328 (1973).

Type Abildgaardia Vahl

Diagnosis (Goetghebeur, 1998): Abildgaardieae is characterized by its clearly differentiated style base, which is often thickened and persistent on the nutlet, but it is deciduous in a number of species. Glumes of the spikelet are typically spirally arranged, but distichous glumes are present in some species. Moniliform stigmatic hairs present. Embryos are of the related Abildgaardia-, Bulbostylis-, Carex-, Fimbristylis-, Schoenus- and Tylocarya-type. (Semmouri et al., 2019).

Accepted genera 10; Abildgaardia Vahl (9 spp.), Actinoschoenus Benth. (2 spp.), Arthrostylis R.Br. (2 spp.), Bulbostylis Kunth (227 spp.; including the formerly recognized segregate genus: Nemum Desv.; Roalson et al., 2018, 2019; Larridon et al., 2019b), Fimbristylis Vahl (320 spp.), Nelmesia Van der Veken (1 sp.), Scleroschoenus K.L.Wilson, J.J.Bruhl & R.L.Barrett (6 spp.), Trachystylis S.T.Blake (1 sp.), Trichoschoenus J.Raynal (1 sp.), and Zulustylis Muasya (2 spp.). For descriptions and notes on the genera, see Larridon et al. (2021c).

Distribution Cosmopolitan.

Key to the genera of Abildgaardieae (based on Larridon et al., 2021c)

- 3a. Fertile flowers per spikelet >2.....4
- 4b. Not this combination of characters......5

- 7a. Style 2-fid; leaf blades well developedTrachystylis
- 7b. Style 3-fid; leaf blades very poorly developed (<1 cm long)8

- 9b. Style base deciduous with rest of style, slightly enlarged10
- 10b. Culm intercostal zones undifferentiated (in Transverse Section); Culms 3–9-angled (always some culms at least 4-angled); nutlet large (>2 mm long), long-stipitate; embryo Abildgaardia-typeScleroschoenus

5.3.18 Tribe Bolboschoeneae

Bolboschoeneae (Tatanov) J.R.Starr, J. Syst. Evol. 59: 825 (2021).

≡ Schoenoplecteae subtrib. Bolboschoeninae Tatanov, in Novosti Sist. Vyssh. Rast. 39: 33 (2007).

Type Bolboschoenus (Asch.) Palla

Diagnosis Differs from all other Cyperaceae tribes by this unique combination of characters: Perennials with long rhizomes often forming hard ovoid tubers at tips. Culms many-noded, 3-sided, thickened at base. Leaves well developed, basal and cauline, eligulate with blade often reduced in lower leaves. Inflorescence terminal (in reduced inflorescences, bract may be erect, but clearly leaf-like), a (compound) corymb-like anthela or capitate with 1 to many spikelets. Inflorescence bracts leaf-like, patent, lowermost often suberect. Spikelets with many spirally arranged, deciduous glumes, each subtending a flower. Glumes puberulent, the apex entire to emarginate or deeply 2-fid, awned or mucronate. Flowers bisexual, perianth present, formed by 3-6 parts, shorter to longer than the nutlet, bristle-like, deciduous with fruit. Stamens 3. Styles 2 or 3. Style base persistent, barely thickened, if at all. Nutlets obovate, dorsiventrally lenticular, or trigonous. Pericarp with the three highly differentiated layers, exocarp cells often enlarged and hollow, surface smooth, epidermal cells roughly isodiametric. Embryo fungiform with three primordial leaves and a notch below the root cap (Bolboschoenus-type).

Accepted genus *Bolboschoenus* (Asch.) Palla (15 spp.). **Distribution** Temperate to tropical regions worldwide.

5.3.19 Tribe Fuireneae

Fuireneae Rchb. ex Fenzl, Gen. Pl.: 116 (1836).

Type Fuirena Rottb.

Diagnosis Differs from all other Cyperaceae tribes by this unique combination of characters: Annuals or rhizomatous perennials. Culms many-noded, rarely scapose, 3–5-sided, sometimes thickened at base. Leaves usually well developed, basal and cauline, ligule tubular, membranous, with blade often reduced in lower leaves (rarely all leaf blades reduced).

Inflorescence terminal (in reduced inflorescences, bract may be erect, but clearly leaf-like), paniculate to capitate with few to many spikelets. Inflorescence bracts leaf-like, usually sheathing, lowermost bract sometimes erect. Spikelets with many spirally or rarely pentastichously arranged, deciduous glumes, each subtending a flower. Glumes often pubescent, the apex entire and mucronate to awned. Flowers bisexual, perianth present, as long or shorter than nutlet, formed by 3 parts, or when 6 in 2 whorls, the inner parts scale-like, the outer parts bristle-like, rarely all parts reduced or absent or only 1 scale developed, deciduous with the fruit. Stamens 1 to 3. Styles 3. Style base persistent, barely thickened, if at all. Nutlets obovate, triquetrous to trigonous, frequently stipitate, smooth or variously ornamented. Embryo turbinate to weakly fungiform with a horizontally broadened scutellum, first leaf primordium not strongly outgrown, the second leaf primordium either absent or poorly developed (Fuirena-type).

Accepted genus Fuirena Rottb. (55 spp.).

Distribution Tropical and warm temperate regions worldwide, especially in the Americas and Africa.

5.3.20 Tribe Schoenoplecteae

Schoenoplecteae Lye, in Blyttia 29: 147 (1971).

Type Schoenoplectus (Rchb.) Palla

Diagnosis Differs from all other Cyperaceae tribes by this unique combination of characters: Perennials with long rhizomes sometimes ending in tubers at tips. Culms nodeless, scapose, trigonous to terete, thickened at base. Leaves usually reduced to a sheath, sometimes developing a ligulate blade, but rarely well developed. Inflorescence pseudolateral, rarely clearly terminal, corymb-like anthela or capitate with (1-)few to many spikelets. Inflorescence bracts often large, erect, stem-like, rarely leaf-like, and patent to reflexed (Actinoscirpus). Spikelets with many spirally arranged, deciduous glumes, each subtending a flower. Glumes puberulent to glabrous, the margins often ciliate or laciniate distally, apex entire to emarginate or deeply 2-fid, awned or mucronate. Flowers bisexual. Perianth present, formed by (-5)6 parts, smooth to retorsely scabrid, bristle-like or sometimes plumose, longer or shorter than nutlet, deciduous with fruit. Stamens 2 or 3. Styles 2 to 3. Style base not thickened, persistent. Nutlets smooth, obovate, trigonous, or dorsiventrally lenticular, yellow to dark brown when mature. Fruit epidermal cells isodiametric to narrowly oblong. Embryo fungiform, scutellum turbinate to rhomboid in shape, root cap lateral, first (well developed) and second embryonic leaves basal (Schoenoplectus-type I).

Accepted genera 2; Actinoscirpus (Ohwi) R.W.Haines & Lye (1 sp.), and Schoenoplectus (Rchb.) Palla (17 spp.). For descriptions and notes on the genera, see Starr et al. (2021).

Distribution Tropical and subtropical Asia from India east to China and south to Northeast Australia (Actinoscirpus), predominantly temperate (Schoenoplectus).

Key to the genera of Schoenoplecteae (based on Starr et al., 2021)

5.3.21 Tribe Pseudoschoeneae

Pseudoschoeneae J.R.Starr, J. Syst. Evol. 59: 826 (2021).

Type Pseudoschoenus (C.B.Clarke) Oteng-Yeb.

Diagnosis Differs from all other Cyperaceae tribes by this unique combination of characters: Annuals or perennials, tufted or with firm, short to creeping rhizomes. Culms nodeless and scapose or 1(-3) noded above the base, trigonous, terete or rarely 7-sided. Leaves reduced to a mucronate sheath, rarely with well-developed blades, ligulate or eligulate. Inflorescence pseudolateral, rarely appearing terminal, a corymb-like anthela or capitate with one to many spikelets, rarely compound paniculate with a conspicuously sinuous main axis (Pseudoschoenus). Inflorescence bracts culm-like, erect, or patent while fruiting, rarely short, rigid, and sheathing, but then appearing as a continuation of the stem. Spikelets with many spirally arranged, deciduous or persistent glumes, each subtending a flower. Scale apex entire to apiculate. Flowers bisexual, rarely polygamodioecious. Perianth present of absent, formed by 0-10 parts, smooth or retrorsely scabrid, bristle-like, as long as or longer than the nutlet, deciduous with the fruit. Stamens 2 or 3, rarely vestigial in female flowers. Basal flowers often present in the axil of leaf sheaths. Styles 2 or 3. Style base undifferentiated, rarely distinct and somewhat thickened, persistent. Nutlets smooth or transversely rugose to distinctly ridged, obovate, trigonous to planoconvex or biconvex, dark nearing black when mature, sometimes brown. Nutlets from basal flowers (when present) are much larger and bear an elongated lignified style (amphicarpy). Fruit epidermal cells longitudinally elongated in shape. Embryo fungiform, scutellum umbonate or distinctly pileate, root cap lateral, first (well developed) and second embryonic leaves basal (Schoenoplectus-type II).

Accepted genera 2; Pseudoschoenus (C.B.Clarke) Oteng-Yeb. (1 sp.) and Schoenoplectiella Lye (62 spp.). For descriptions and notes on the genera, see Starr et al. (2021).

Distribution Southern Africa (*Pseudoschoenus*), temperate to tropical regions worldwide (*Schoenoplectiella*).

Key to the genera of Pseudoschoeneae (based on Starr et al., 2021)

- 1a. Inflorescence paniculate or racemose, with a definite main axis of well-developed internodesPseudoschoenus
- 1b. Inflorescence corymb-like or reduced to one or a cluster of sessile spikelets, without a definite main axis due to highly reduced internodesSchoenoplectiella

5.3.22 Tribe Cypereae

Cypereae Dumort., Anal. Fam. Pl.: 65. (1829).

Type Cyperus L.

Diagnosis Cypereae are characterized by a Cyperus-type or the strongly similar Ficinia-type embryo (Van der Veken, 1965; Goetghebeur, 1998; Semmouri et al, 2019).

Includes 2 subtribes, 8 genera. **Distribution** Cosmopolitan.

Key to the subtribes of Cypereae

1a. Glumes usually spirally arranged; anatomy C_3 . If distichously arranged, then 2 or more parallel veins

Subtribe Ficiniinae. Ficiniinae Fenzl in Endl., Gen. Pl.: 117. (1836).

Type Ficinia Schrad.

Diagnosis Annuals or perennials, tufted or with short to creeping rhizomes. Culms nodeless and scapose or noded above the base, trigonous or terete. Leaves reduced to a mucronate sheath or with well-developed blades, ligulate or eligulate. Inflorescence comprising a single terminal spikelet, or pseudolateral, anthelate, or capitate. Spikeletets usually terete, flowers bisexual (except in *Afroscirpoides*), perianth absent or varying from scales to cotton-like bristles, nutlet sessile or bearing a gynophore (*Ficinia*). Embryo Cyperus or Ficinia-type.

Accepted genera 7; Afroscirpoides García-Madr. & Muasya (1 sp.), Dracoscirpoides Muasya (3 spp.), Erioscirpus Palla (2 spp.), Ficinia Schrad. (87 spp.), Hellmuthia Steud. (1 sp.), Isolepis R.Br. (69 spp.), and Scirpoides Ség. (4 spp.; including the formerly recognized segregate genus: Karinia Reznicek & McVaugh). For descriptions and notes on the genera, see Muasya et al. (2012), Yano et al. (2012), García-Madrid et al. (2015), Reid et al. (2017), and Muasya & Larridon (2021).

Distribution Cosmopolitan.

Key to the genera of Ficiniinae (adapted from García-Madrid et al., 2015)

- 3b. Leaf blades and inflorescence bracts smooth, nutlets without gynophore......4
- 4b. Glumes chartaceous to hyalineIsolepis

- 6a. Anthelate inflorescence (sometimes ± capitate in *Erioscirpus microstachyus*) with few to many sessile spikelets in clusters of (1) 2–3. Long perianth bristles cotton-like Erioscirpus

Subtribe Cyperinae. Cyperinae Pax, Bot. Jahrb. Syst. 7: 306. 28 (1886).

Type Cyperus L.

Diagnosis Annuals or perennials, tufted or with short to creeping rhizomes. Culms usually nodeless and scapose, trigonous or terete. Leaves reduced to a mucronate sheath or with well-developed blades, eligulate. Inflorescence comprising one to many spikelets, capitate or anthelate. Spikelets usually with distichous glumes, flowers usually bisexual, perianth absent, lacking a gynophore (except Cyperus lipocarphioides (Kük.) Lye). Embryo Cyperus-type.

Accepted genus Cyperus L. (964 spp.; including the formerly recognized segregate genera: Alinula J.Raynal, Androtrichum (Brongn.) Brongn., Ascolepis Nees ex Steud., Ascopholis C.E.C.Fisch., Courtoisina Soják, Kyllinga Rottb., Kyllingiella R.W.Haines & Lye, Lipocarpha R.Br., Oxycaryum Nees, Pycreus P.Beauv., Queenslandiella Domin, Remirea Aubl., Sphaerocyperus Lye, and Volkiella Merxm. & Czech). For taxonomic changes to the circumscription of Cyperus, see Larridon et al. (2011b, 2014), Bauters et al. (2014), Pereira-Silva et al. (2020).

Distribution Cosmopolitan.

5.3.23 Incertae sedis

The monotypic genus Rhynchocladium T.Koyama remains unplaced. It is limited in distribution to Guyana and southern Venezuela.

Acknowledgements

The molecular lab work and initial bioinformatics were funded by the Calleva Foundation and the Sackler Trust through the Plant and Fungal Trees of Life programme (https://www.kew.org/science/our-science/projects/plant-and-fungal-trees-of-life), by funding to IL from the Royal Botanic Gardens, Kew, including a pilot study grant. This study was supported by a NSERC grant to JRS enabling a 6-week research stay at Royal Botanic Gardens, Kew, covering some of the lab work and sequencing costs. The authors are grateful to Juan Carlos Zamora for allowing them to use his photo of bee pollination in Rhynchospora alba and to Rachun Pooma for his photo of Khaosokia caricoides.

References

- Altschul SF, Gish W, Miller W, Myers EW, Lipman DJ. 1990. Basic local alignment search tool. *Journal of Molecular Biology* 215: 403–410.
- Alves M. 2015. Hypolytrum (Cyperaceae): Taxonomic and nomenclatural notes, geographical distribution and conservation status of Neotropical species. Rodriguésia 66: 379–392.
- Bagnall RS. 2009. The Oxford handbook of papyrology. Oxford: Oxford University Press.

- Baker WJ, Bailey P, Barber V, Barker A, Bellot S, Bishop D, Botigué LR, Brewer G, Carruthers T, Clarkson JJ, Cook J, Cowan RS, Dodsworth S, Epitawalage N, Françoso E, Gallego B, Johnson MG, Kim JT, Leempoel K, Maurin O, McGinnie C, Pokorny L, Roy S, Stone M, Toledo E, Wickett NJ, Zuntini AR, Eiserhardt WL, Kersey PJ, Leitch IJ, Forest F. A comprehensive phylogenomic platform for exploring the angiosperm Tree of Life. BioRxiv 2021.02.22.431589
- Banack SA, Rondón XJ, Diaz-Huamanchumo W. 2004. Indigenous cultivation and conservation of Totora (Schoenoplectus californicus, Cyperaceae) in Peru. Economic Botany 58: 11–20.
- Bankevich A, Nurk S, Antipov D, Gurevich AA, Dvorkin M, Kulikov AS, Lesin VM, Nikolenko SI, Pham S, Prjibelski AD, Pyshkin AV, Sirotkin AV, Vyahhi N, Tesler G, Alekseyev MA, Pevzner PA. 2012. SPAdes: A new genome assembly algorithm and its applications to single-cell sequencing. *Journal of Computational Biology* 19: 455–477.
- Bakker FT. 2017. Herbarium genomics: Skimming and plastomics from archival specimens. Webbia 72: 35–45.
- Barrett RL. 2013. Ecological importance of sedges: A survey of the Australasian Cyperaceae genus *Lepidosperma*. *Annals of Botany* 111: 499–529.
- Barrett RL, Wilson KL. 2012. A review of the genus *Lepidosperma* Labill. (Cyperaceae: Schoeneae). *Australian Systematic Botany* 25: 225–294.
- Barrett RL, Wilson KL. 2013. Two new species of *Lepidosperma* (Cyperaceae) occurring in the Perth area of Western Australia. Nuytsia 23: 173–187.
- Barrett RL, Wilson KL, Bruhl JJ. 2019. Anthelepis, a new genus for four mainly tropical species of Cyperaceae from Australia, New Caledonia and South-East Asia. Australian Systematic Botany 32: 269–289.
- Barrett RL, Wilson KL, Bruhl JJ. 2020. Reinstatement and revision of the genus *Chaetospora* (Cyperaceae: Schoeneae). *Telopea* 23: 95–112.
- Barrett RL, Bruhl JJ, Wilson KL. 2021a. Netrostylis, a new genus of Australasian Cyperaceae removed from *Tetraria*. *Telopea* 24: 53–60.
- Barrett RL, Bruhl JJ, Wilson KL. 2021b. Revision of generic concepts in Schoeneae subtribe Tricostulariinae (Cyperaceae) with a new Australian genus Ammothryon and new species of Tricostularia. Telopea 24: 61–169.
- Barrett RL, Taputuarai R, Meyer J-Y, Bruhl JJ, Wilson KL. 2021c. Reassessment of the taxonomic status of Cyperaceae on Rapa Iti, Austral Islands, French Polynesia, with a new combination, Morelotia involuta. Telopea 24: 171–187.
- Bauters K. 2018. Scleria revized: A molecular approach towards a new classification. Ph.D. Dissertation. Gent: Ghent University.
- Bauters K, Larridon I, Reynders M, Huygh W, Asselman P, Vrijdaghs A, Muasya AM, Goetghebeur P. 2014. A new classification for Lipocarpha and Volkiella as infrageneric taxa of Cyperus s.l. (Cypereae, Cyperoideae, Cyperaceae): Insights from species tree reconstruction supplemented with morphological and floral developmental data. Phytotaxa 166: 1–32.
- Bauters K, Asselman P, Simpson DA, Muasya AM, Goetghebeur P, Larridon I. 2016. Phylogenetics, ancestral state reconstruction, and a new infrageneric classification of Scleria (Cyperaceae) based on three DNA regions. *Taxon* 65: 444–466.
- Beck JB, Alexander PJ, Applhin L, Al-Shehbaz IA, Rushworth C, Bailey CD, Windham MD. 2012. Does hybridization drive the transition to asexuality in diploid Boechera? *Evolution* 66: 985–995.

Beentje H. 2016. The Kew plant glossary. 2nd edition. Royal Botanic Gardens, Kew, Richmond: Kew Publishing.

- Besnard G, Muasya AM, Russier F, Roalson EH, Salamin N, Christin P-A. 2009. Phylogenomics of C4 photosynthesis in sedges (Cyperaceae): Multiple appearances and genetic convergence. *Molecular Biology and Evolution* 26: 1909–1919.
- Biosecurity Queensland. 2018. *Technical s. Invasive plant and animal research* 2017–2018. State of Queensland: Queensland Government.
- Bouchenack-Khelladi Y, Muasya AM, Linder HP. 2014. A revised evolutionary history of Poales: Origins and diversification. Botanical Journal Linnean Society 175: 4–16.
- Bolger AM, Lohse M, Usadel B. 2014. Trimmomatic: A flexible trimmer for Illumina Sequence Data. *Bioinformatics* 30: 2114–2120.
- Borowiec ML. 2016. AMAS: A fast tool for alignment manipulation and computing of summary statistics. *PeerJ (Corta Madera, CA and London)* 4: e1660.
- Brewer GE, Clarkson JJ, Maurin O, Zuntini AR, Barber V, Bellot S, Cowan R, Davies NMJ, Dodsworth S, Edwards SL, Eiserhardt WL, Epitawalage N, Frisby S, Grall A, Kersey PJ, Pokorny L, Leitch IL, Forest F, Baker WJ. 2019. Factors affecting the targeted sequencing of 353 nuclear genes from herbarium specimens spanning the diversity of angiosperms. Frontiers in Plant Science 10: 1102.
- Browning J, Gordon-Gray KD. 1999. Studies in Cyperaceae in southern Africa. 33: A new monotypic genus, Capeobolus. South African Journal of Botany 65: 218–222.
- Bruhl JJ, Wilson KA. 2007. Towards a comprehensive survey of C3 and C4 photosynthetic pathways in Cyperaceae. In: Columbus JT, Friar EA, Porter JM, Prince LM, Simpson MG eds. Monocots III/ grasses IV. Aliso 23. Claremont, CA: Rancho Santa Ana Botanic Garden. 99–148.
- Bryson CT, Carter R. 2008. The significance of Cyperaceae as weeds. In: Naczi RFC, Ford BA eds. Sedges: Uses, diversity and systematics of the Cyperaceae. Monographs in Systematic Botany from the Missouri Botanical Garden. 108: 15–101. St. Louis: Missouri Botanical Garden.
- Buerki S, Baker WJ. 2015. Collections-based research in the genomic era. Biological Journal of the Linnean Society 117: 5–10.
- Cerling TE, Kyalo Manthi F, Mbua EN, Leakey LN, Leakey MG, Leakey RE, Brown FH, Grine FE, Hart JA, Kaleme P, Roche H, Uno KT, Wood BA. 2013a. Stable isotope-based diet reconstructions of Turkana Basin hominins. Proceedings of the National Academy of Sciences of the United States of America 110: 10501–10506.
- Cerling TE, Chritz KL, Jablonski NG, Leakey MG, Kyalo Manthi F. 2013b. Diet of Theropithecus from 4 to 1 Ma in Kenya. Proceedings of the National Academy of Sciences of the United States of America 110: 10507–10512.
- Coan AI, Alves MV, Scatena VL. 2010. Evidence of pseudomonad pollen formation in *Hypolytrum* (Mapanioideae, Cyperaceae). Australian Journal of Botany 58: 663–672.
- Costa ACG, Albuquerque IS, Thomas WW, Machado IC. 2018a. Influence of environmental variation on the pollination of the ambophilous sedge *Rhynchospora ciliata* (Cyperaceae). *Plant Ecology* 219: 241–250.
- Costa SM, Morokawa R, Bittrich V, Thomas WW, Amaral MCE. 2018b. What about Cryptangieae (Cyperaceae)—A molecular hypothesis about its tribal status and circumscription. *Phytotaxa* 347: 127–149.
- Costa SM, Vitta FA, Thomas WW, Muasya AM, Morokawa R, Bittrich V, Shepperd GJ, Amaral MCE. 2021a. An updated generic

- circumscription of Cryptangieae (Cyperaceae, Poales) based on a molecular phylogeny and a morphological character reconstruction. *Phytotaxa* 483: 211–228.
- Costa SM, Amaral MCE, Thomas WW. 2021b. Two new species of Cephalocarpus (Cryptangieae, Cyperaceae) from the Venezuelan Guayana Highland. Brittonia. https://doi.org/10.1007/s12228-021-09652-1
- Couvreur TLP, Helmstetter AJ, Koenen EJM, Bethune K, Brandão RD, Little SA, Sauquet H, Erkens RHJ. 2019. Phylogenomics of the major tropical plant family Annonaceae using targeted enrichment of nuclear genes. Frontiers in Plant Sciences 9: 1941.
- Dai LK, Liang SY, Zhang SR, Tang YC, Koyama T, Tucker GC. 2010. *Carex* L. In: Wu ZY, Raven PH, Hong DY eds. *Flora of China*. Beijing: Science Press; St. Louis: Missouri Botanical Garden Press. 23: 285–461.
- Dhooge S, Goetghebeur P, Muasya AM. 2003. Zameioscirpus, a new genus of Cyperaceae from South America. Plant Systematics and Evolution 243: 73–84.
- Dodet M, Petit RJ, Gasquez J. 2008. Local spread of the invasive Cyperus esculentus (Cyperaceae) inferred using molecular genetic markers. Weed Research 48: 19–27.
- Dominy NJ. 2012. Hominins living on the sedge. Proceedings of the National Academy of Sciences of the United States of America 109: 20171–20172.
- Doyle JJ, Doyle JL. 1987. A rapid DNA isolation procedure for small quantities of fresh leaf tissue. Phytochemical Bulletin, Botanical Society of America 19: 11–15.
- Eiten LT. 1976. Inflorescence units in Cyperaceae. Annals of the Missouri Botanical Garden 63: 81–112.
- Elliott TL, Muasya AM. 2017. Taxonomic realignment in the southern African *Tetraria* (Cyperaceae, tribe Schoeneae; Schoenus clade). South African Journal of Botany 112: 354–360.
- Elliott TL, Muasya AM. 2018. A taxonomic revision of Schoenus compar–Schoenus pictus and allies (Cyperaceae, tribe Schoeneae) with three new species described from South Africa. South African Journal of Botany 114: 303–315.
- Elliott TL, Barrett RL, Muasya AM. 2019. A taxonomic revision of Schoenus cuspidatus and allies (Cyperaceae, tribe Schoeneae)—Part 1. South African Journal of Botany 121: 519–536.
- Elliott TL, Muasya AM. 2020. A taxonomic revision of Schoenus cuspidatus and allies (Cyperaceae, tribe Schoeneae)—Part 2. South African Journal of Botany 130: 327–347.
- Elliott TL, van Mazijk R, Barrett RL, Bruhl JJ, Joly S, Muthaphuli N, Wilson KL, Muasya AM. 2021. Global dispersal and diversification of the genus Schoenus (Cyperaceae) from the Western Australian biodiversity hotspot. *Journal of Systematics and Evolution* 59: 791–808.
- Escudero M, Hipp AL. 2013. Shifts in diversification rates and clade ages explain species richness in higher-level sedge taxa (Cyperaceae). American Journal of Botany 100: 2403–2411.
- Galán Díaz J, Bauters K, Rabarivola L, Xanthos M, Goetghebeur P, Larridon I. 2019. A revision of Scleria (Cyperaceae) in Madagascar. Blumea 64: 195–213.
- García-Madrid AS, Muasya AM, Álvarez I, Cantó P, Molina JA. 2015. Towards resolving phylogenetic relationships in the Ficinia clade and description of the new genus Afroscirpoides (Cyperaceae: Cypereae). Taxon 64: 688–702.
- Gilmour CN, Starr JR, Naczi RFC. 2013. *Calliscirpus*, a new genus for two narrow endemics of the California Floristic Province, *C. criniger* and *C. brachythrix* sp. nov. (Cyperaceae). *Kew Bulletin* 68: 85–105.

- Givnish TJ, Ames M, McNeal JR, McKain MR, Steele PR, dePamphilis CW, Graham SW, Pires JC, Stevenson DW, Zomlefer WB, Briggs BG, Duvall MR, Moore MJ, Heaney JM, Soltis DE, Soltis PS, Thiele K, Leebens-Mack JH. 2010. Assembling the tree of the monocotyledons: Plastome sequence phylogeny and evolution of Poales. Annals of the Missouri Botanical Garden 97: 584–616.
- Global Carex Group. 2015. Making Carex monophyletic (Cyperaceae, tribe Cariceae): A new broader circumscription. Botanical Journal of the Linnean Society 179: 1–42.
- Glon HE, Shiels DR, Linton E, Starr JR, Shorkey AL, Fleming S, Lichtenwald SK, Schick ER, Pozo D, Monfils AK. 2017. A five gene phylogenetic study of Fuireneae (Cyperaceae) with a revision of Isolepis humillima. Systematic Botany 42: 26–36.
- Goetghebeur P. 1986. Genera Cyperacearum. Een bijdrage tot de kennis van de morfologie, systematiek en fylogenese van de Cyperaceae-genera. Ph.D. Dissertation. Gent: Ghent University.
- Goetghebeur P. 1998. Cyperaceae. In: Kubitzki K ed. The families and genera of vascular plants. Berlin: Springer-Verlag. 4: 141–190.
- González-Elizondo MS, Peterson PM. 1997. A classification and key to the supraspecific taxa in *Eleocharis* (Cyperaceae). *Taxon* 46: 443–449.
- Govaerts R, Jiménez-Mejías P, Koopman J, Simpson DA, Goetghebeur P, Wilson KL, Egorova T, Bruhl JJ. 2020. World checklist of Selected Plant Families. Cyperaceae. Facilitated by the Royal Botanic Gardens, Kew [online]. Available from http://www.kew.org/wcsp/monocots/ [accessed 2 June 2020].
- Handel SN. 1976. Dispersal ecology of Carex pedunculata (Cyperaceae), a new North American myrmecochore. American Journal of Botany 63: 1071–1079.
- Hart ML, Forrest LL, Nicholls JA, Kidner CA. 2016. Retrieval of hundreds of nuclear loci from herbarium specimens. *Taxon* 65: 1081–1092.
- Hinchliff CE, Lliully AE, Carey T, Roalson EH. 2010. The origins of *Eleocharis* (Cyperaceae) and the status of *Websteria*, Egleria, and *Chillania*. Taxon 59: 709–719.
- Hinchliff CE, Roalson EH. 2013. Using supermatrices for phylogenetic inquiry: An example using the sedges. Systematic Biology 62: 205–219.
- Hoang DT, Chernomor O, von Haeseler A, Minh BQ, Vinh LS. 2018. UFBoot2: improving the ultrafast bootstrap approximation. *Molecular Biology and Evolution* 35: 518–522.
- ISSG. 2019. Global Invasive Species Database. Developed and managed by the Invasive Species Specialist Group (ISSG) of the Species Survival Commission (SSC) of the International Union for Conservation of Nature (IUCN) [online]. Available from http://www.iucngisd.org/gisd/ [accessed 13 May 2019].
- Jiménez-Mejías P, Martinetto E, Momohara A, Popova S, Smith SY, Roalson EH. 2016a. Commented synopsis of the Pre-Pleistocene fossil record of *Carex* (Cyperaceae). Botanical Review 82: 258–345.
- Jiménez-Mejías P, Luceño M, Wilson KL, Waterway MJ, Roalson EH. 2016b. Clarification of the use of the terms perigynium and utricle in Carex L. (Cyperaceae). Systematic Botany 41: 519–528.
- Jiménez-Mejías P, Hahn M, Lueders K, Starr JR, Brown BH, Chouinard BN, Chung K-S, Escudero M, Ford BA, Ford KA, Gebauer S, Gehrke B, Hoffmann MH, Jin X-F, Jung J, Kim S, Luceño M, Maguilla E, Martín-Bravo S, Míguez M, Molina A, Naczi RFC, Pender JE, Reznicek AA, Villaverde T, Waterway MJ, Wilson KL, Yang J-C, Zhang S, Hipp AL, Roalson EH. 2016c. Megaphylogenetic specimen-level approaches to the *Carex* (Cyperaceae) phylogeny using ITS, ETS, and *matK* sequences: Implications for classification. *Systematic Botany* 41: 500–518.

- Johnson MG, Gardner EM, Liu Y, Medina R, Goffinet B, Shaw AW, Zerega NJC, Wickett NJ. 2016. HybPiper: Extracting coding sequence and introns for phylogenetics from high-throughput sequencing reads using target enrichment. *Applications in Plant Sciences* 4(7): 1600016.
- Johnson MG, Pokorny L, Dodsworth S, Botigué LR, Cowan RS, Devault A, Eiserhardt WL, Epitawalage N, Forest F, Kim JT, Leebens-Mack JH, Leitch IJ, Maurin O, Soltis DE, Soltis PS, Wong GK, Baker WJ, Wickett NJ. 2019. A universal probe set for targeted sequencing of 353 nuclear genes from any flowering plant designed using k-medoids clustering. Systematic Biology 68: 594–606.
- Jung J, Choi H-K. 2013. Recognition of two major clades and early diverged groups within the subfamily Cyperoideae (Cyperaceae) including Korean sedges. *Journal of Plant Research* 126: 335–349.
- Junier T, Zdobnov EM. 2010. The Newick utilities: High throughput phylogenetic tree processing in the UNIX shell. *Bioinformatics* 26: 1669–1670.
- Kadlec M, Bellstedt DU, Le Maitre NC, Pirie MD. 2017. Targeted NGS for species level phylogenomics: "made to measure" or "one size fits all"? PeerJ (Corta Madera, CA and London) 5: e3569.
- Katoh K, Standley DM. 2013. MAFFT multiple sequence alignment software version 7: improvements in performance and usability. *Molecular Biology and Evolution* 30: 772–780.
- Kern JH. 1974. Cyperaceae. In: van Steenis CGGJ ed. Flora Malesiana Series I. Groningen: Wolters-Noordhoff Publishing. 435–753.
- Koyama T. 1958. Taxonomic study of the genus Scirpus Linné. Journal of the Faculty of Science, the University of Tokyo 7: 324–325.
- Kukkonen I. 1984. On the inflorescence structure in the family Cyperaceae. Annales Botanici Fennici 21: 257–264.
- Larridon I, Reynders M, Huygh W, Bauters K, Van de putte K, Muasya AM, Boeckx P, Simpson DA, Vrijdaghs A, Goetghebeur P. 2011a. Affinities in C₃ Cyperus lineages (Cyperaceae) revealed using molecular phylogenetic data and carbon isotope analysis. Botanical Journal of the Linnean Society 167: 19–46.
- Larridon I, Reynders M, Huygh W, Bauters K, Vrijdaghs A, Leroux O, Muasya AM, Simpson DA, Goetghebeur P. 2011b. Taxonomic changes in C₃ Cyperus (Cyperaceae) supported by molecular phylogenetic data, morphology, embryology, ontogeny and anatomy. *Plant Ecology and Evolution* 144: 327–356.
- Larridon I, Bauters K, Reynders M, Huygh W, Muasya AM, Simpson DA, Goetghebeur P. 2013. Towards a new classification of the giant paraphyletic genus Cyperus (Cyperaceae): Phylogenetic relationships and generic delimitation in C4 Cyperus. Botanical Journal of the Linnean Society 172: 106–126.
- Larridon I, Bauters K, Huygh W, Reynders M, Goetghebeur P. 2014. Taxonomic changes in C₄ Cyperus (Cypereae, Cyperoideae, Cyperaceae): combining the sedge genera Ascolepis, Kyllinga and Pycreus into Cyperus s.l. Phytotaxa 166: 33–48.
- Larridon I, Verboom GA, Muasya AM. 2017. (2555) Proposal to conserve the name *Tetraria* (Cyperaceae) with a conserved type. Taxon 66: 1226–1227.
- Larridon I, Semmouri I, Bauters K, Viljoen JA, Prychid CJ, Muasya AM, Bruhl JJ, Wilson KL, Goetghebeur P. 2018a. Molecular phylogenetics of the genus Costularia (Schoeneae, Cyperaceae) reveals multiple distinct evolutionary lineages. Molecular Phylogenetics and Evolution 126: 196–209.
- Larridon I, Verboom GA, Muasya AM. 2018b. Revised delimitation of the genus *Tetraria*, nom. cons. prop. (Cyperaceae, tribe Schoeneae, Tricostularia clade). South African Journal of Botany 118: 18–22.

Larridon I, Rabarivola L, Xanthos M, Muasya AM. 2019a. Revision of the Afro-Madagascan genus Costularia (Cyperaceae): Infrageneric relationships and species delimitation. *PeerJ* (Corta Madera, CA and London) 7: e6528.

- Larridon I, Goetghebeur P, Roalson EH. 2019b. New combinations and a new name for including *Nemum* species in *Bulbostylis* (Abildgaardieae: Cyperaceae). *Phytotaxa* 418: 112–114.
- Larridon I, Villaverde T, Zuntini AR, Pokorny L, Brewer G, Epitawalage N, Fairlie I, Hahn M, Kim J, Maguilla E, Maurin O, Xanthos M, Hipp A, Forest F, Baker WJ. 2020. Tackling rapid radiations with targeted sequencing. Frontiers in Plant Science 10: 1655.
- Larridon I, Galán Díaz J, Bauters K, Escudero M. 2021a. What drives diversification in a pantropical plant lineage with extraordinary capacity for long-distance dispersal and colonisation? *Journal of Biogeography* 48: 64–77.
- Larridon I, Spalink D, Jiménez-Mejías P, Márquez-Corro JI, Martín-Bravo S, Muasya AM, Escudero M. 2021b. The evolutionary history of sedges in Madagascar. *Journal of Biogeography* 48: 917–932.
- Larridon I, Zuntini AR, Barrett RL, Wilson KL, Bruhl JJ, Goetghebeur P, Baker WJ, Brewer GE, Epitawalage N, Fairlie I, Forest F, Kikuchi I, Pokorny L, Semmouri I, Spalink D, Simpson DA, Muasya AM, Roalson EH. 2021c. Resolving the generic limits in Cyperaceae tribe Abildgaardieae using targeted sequencing. Botanical Journal of the Linnean Society 196: 163–187.
- Léveillé-Bourret É, Gilmour CN, Starr JR, Naczi RFC, Spalink D, Sytsma KJ. 2014. Searching for the sister to sedges (*Carex*): resolving relationships in the Cariceae-Dulichieae-Scirpeae clade (Cyperaceae). Botanical Journal of the Linnean Society 176: 1–21.
- Léveillé-Bourret É, Donadío S, Gilmour CN, Starr JR. 2015. Rhodoscirpus (Cyperaceae: Scirpeae), a new South American sedge genus supported by molecular, morphological, anatomical and embryological data. *Taxon* 64: 931–944.
- Léveillé-Bourret É, Starr JR, Ford BA. 2018a. Why are there so many sedges? Sumatroscirpeae, a missing piece in the evolutionary puzzle of the giant genus Carex (Cyperaceae). Molecular Phylogenetics and Evolution 119: 93–104.
- Léveillé-Bourret É, Starr JR, Ford BA. 2018b. A Revision of Sumatroscirpus (Sumatroscirpeae, Cyperaceae) with discussions on Southeast Asian biogeography, general collecting, and homologues with Carex (Cariceae, Cyperaceae). Systematic Botany 43: 510–531.
- Léveillé-Bourret É, Starr JR, Ford BA, Moriarty Lemmon E, Lemmon AR. 2018c. Resolving rapid radiations within angiosperm families using Anchored Phylogenomics. Systematic Biology 67: 94–112.
- Léveillé-Bourret É, Starr JR. 2019. Molecular and morphological data reveal three new tribes within the Scirpo-Caricoid Clade (Cyperoideae, Cyperaceae). *Taxon* 68: 218–245.
- Léveillé-Bourret É, Chen B-H, Garon-Labrecque M-È, Ford BA, Starr JR. 2020. RAD sequencing resolves the phylogeny, taxonomy and biogeography of Trichophoreae despite a recent rapid radiation (Cyperaceae). *Molecular Phylogenetics and Evolution* 145: 106727.
- Lorougnon G. 1973. Le vecteur pollinique chez les *Mapania* et les *Hypolytrum* Cypéracées de sous-bois des foréts tropicales ombrophiles. *Bulletin Jardin Botanique National de Belgique* 43: 33–36.
- Maguire B. 1972. The botany of the Guayana Highland—part IX. Memoirs of the New York Botanical Garden 23: 1–832.
- Márquez-Corro JI, Martín-Bravo S, Spalink D, Luceño M, Escudero M. 2019. Inferring hypothesis-based transitions in clade-specific

- models of chromosome number evolution in sedges (Cyperaceae). Molecular Phylogenetics and Evolution 135: 203–209.
- Márquez-Corro JI, Martín-Bravo S, Jiménez-Mejías P, Hipp AL, Spalink D, Naczi RFC, Roalson EH, Luceño M, Escudero M. 2021. Macroevolutionary insights in sedges (*Carex*: Cyperaceae): The effects of rapid chromosome number evolution on lineage diversification. *Journal of Systematics and Evolution* 59: 776–790.
- Martín-Bravo S, Jiménez-Mejías P, Villaverde T, Escudero M, Hahn M, Spalink D, Roalson EH, Hipp AL, the Global *Carex* Group: Benítez-Benítez C, Bruederle LP, Fitzek E, Ford BA, Ford KA, Garner M, Gebauer S, Hoffmann MH, Jin X-F, Larridon I, Léveillé-Bourret É, Lu Y-F, Luceño M, Maguilla E, Márquez-Corro JI, Míguez M, Naczi R, Reznicek AA, Starr JR. 2019. A tale of worldwide success: Behind the scenes of *Carex* (Cyperaceae) biogeography and diversification. *Journal of Systematics and Evolution* 57: 695–718.
- Minh BQ, Schmidt HA, Chernomor O, Schrempf D, Woodhams MD, von Haeseler A, Lanfear R. 2020. IQ-TREE 2: New models and efficient methods for phylogenetic inference in the genomic era. *Molecular Biology and Evolution* 37: 2461.
- Muasya AM, de Lange PJ. 2010. Ficinia spiralis (Cyperaceae) a new genus and combination for Desmoschoenus spiralis. New Zealand Journal of Botany 48: 31–39.
- Muasya AM, Larridon I. 2021. Delimiting the genera of the Ficinia Clade (Cypereae, Cyperaceae) based on molecular phylogenetic data. *PeerJ (Corta Madera, CA and London)* 9: e10737.
- Muasya AM, Simpson DA, Chase MW, Culham A. 1998. An assessment of suprageneric phylogeny in Cyperaceae using rbcL DNA sequences. *Plant Systematics and Evolution* 211: 257–271.
- Muasya AM, Simpson DA, Chase MW. 2002. Phylogenetic relationships in *Cyperus L. s.l.* (Cyperaceae) inferred from plastid DNA sequence data. Botanical Journal Linnean Society 138: 145–153.
- Muasya AM, Simpson DA, Verboom GA, Goetghebeur P, Naczi RFC, Chase MW, Smets E. 2009a. Phylogeny of Cyperaceae based on DNA sequence data: Current progress and future prospects. Botanical Review 75: 2–21.
- Muasya AM, Vrijdaghs A, Simpson DA, Chase MW, Goetghebeur P, Smets E. 2009b. What is a genus in Cypereae: Phylogeny, character homology assessment and generic circumscription in Cypereae. Botanical Review 75: 52–66.
- Muasya AM, Harvey Y, Cheek M, Tah K, Simpson DA. 2010. *Coleochloa domensis* (Cyperaceae), a new epiphytic species from Cameroon. *Kew Bulletin* 62: 323–325.
- Muasya AM, Goetghebeur P, Larridon I. 2020. Zulustylis (Abildgaardieae, Cyperaceae)—A new genus from sub-Saharan Africa. South African Journal of Botany 128: 326–332.
- Musili PM, Gibbs AK, Wilson KL, Bruhl JJ. 2016. Schoenus (Cyperaceae) is not monophyletic based on ITS nrDNA sequence data. Australian Systematic Botany 29: 265–283.
- Naczi RFC, Ford BA. 2008. Sedges: uses, diversity, and systematics of Cyperaceae. St Louis: Missouri Botanical Garden Press.
- Nagels A, Muasya AM, Huysmans S, Vrijdaghs A, Smets E, Vinckier S. 2009. Palynological diversity and major evolutionary trends in Cyperaceae. *Plant Systematics and Evolution* 277: 117–142.
- Novoselova MS. 1994a. The system of the genus Eriophorum (Cyperaceae). I. Subgenera Erioscirpus, Eriophoropsis, Phyllanthela. Botanicheskiy Zhurnal 79: 77–89.
- Novoselova MS. 1994b. The system of the genus *Eriophorum* (Cyperaceae). II. Subgenerus *Eriophorum*. Botanicheskiy Zhurnal 79: 66–75.

- Paradis E, Schliep K. 2018. ape 5.0: An environment for modern phylogenetics and evolutionary analyses in R. *Bioinformatics* 35: 526–528.
- Pereira-Silva L, Trevisan R, Rodrigues AC, Larridon I. 2020. Combining the small South American genus Androtrichum in Cyperus (Cyperaceae). Plant Ecology and Evolution 153: 446–454.
- Philcox D. 1961. An Uncinia from South Georgia. Kew Bulletin 15: 229.
- POWO. 2020. Plants of the World Online. Board of Trustees of the Royal Botanic Gardens, Kew [online]. Available from http://www.plantsoftheworldonline.org/ [accessed 15 January 2020].
- Prychid CJ, Bruhl JJ. 2013. Floral ontogeny and gene protein localization rules out euanthial interpretation of reproductive units in *Lepironia* (Cyperaceae, Mapanioideae, Chrysitricheae). Annals of Botany 112: 161–177.
- R Core Team. 2020. R: A language and environment for statistical computing. Vienna: R Foundation for Statistical Computing. https://www.R-project.org/
- Reutemann AG, Ardissone RE, López MG, Muchut SE, Boldrini I, Vegetti AC, Giussani LM. 2018. Phylogenetic relationships in *Bulbostylis* (Abildgaardieae: Cyperaceae) inferred from nuclear and plastid DNA sequence data. Systematics and Biodiversity 16: 441–452.
- Reznicek AA, González-Elizondo MS, Hahn M, Garner M, Hipp AL. 2021. Monograph of Carex section Schiedeanae (Cyperaceae): Unexpected taxonomic and ecological diversity in a Mexican sedge clade. Journal of Systematics and Evolution 59: 698–725.
- Roalson EH, Friar EA. 2000. Infrageneric classification of *Eleocharis* (Cyperaceae) revisited: Evidence from the internal transcribed spacer (ITS) region of nuclear ribosomal DNA. Systematic Botany 25: 323–336.
- Roalson EH, Hinchliff CE, Trevisan R, da Silva CRM. 2010. Phylogenetic relationships in *Eleocharis* (Cyperaceae): C₄ photosynthesis origins and patterns of diversification in the spikerushes. Systematic Botany 35: 257–271.
- Roalson EH, Simpson DA, Larridon I. 2018. (2618) Proposal to conserve *Bulbostylis*, nom. cons. (Cyperaceae) against an additional name, *Nemum. Taxon* 67: 642.
- Roalson EH, Prata AP, Mesterházy A, Chase MW, Simpson DA, Thomas WW, Larridon I. 2019. A broader circumscription of Bulbostylis as to include Nemum (Abildgaardieae: Cyperaceae). Phytotaxa 395: 199–208.
- Roalson EH, Jiménez-Mejías P, Hipp AL, Benítez-Benítez C, Bruederle LP, Chung K-S, Escudero M, Ford BA, Ford K, Gebauer S, Gehrke B, Hahn M, Hayat MQ, Hoffmann MH, Jin X-F, Kim S, Larridon I, Léveillé-Bourret É, Lu Y-F, Luceño M, Maguilla E, Márquez-Corro JI, Martín-Bravo S, Masaki T, Míguez M, Naczi RFC, Reznicek AA, Spalink D, Starr JR, Uzma, Villaverde T, Waterway MJ, Wilson KL, Zhang S-R. 2021. A framework infrageneric classification of *Carex* (Cyperaceae) and its organizing principles. *Journal of Systematics and Evolution* 59: 726–762.
- Rogers HH, Runion GB, Prior SA, Price AJ, Torbert HA, Gjerstad DA. 2008. Effects of elevated atmospheric CO₂ on invasive plants: comparison of purple and yellow nutsedge (*Cyperus rotundus* L. and C. esculentus L.). Journal of Environmental Quality 37: 395–400.
- Ruhfel BR, Gitzendanner MA, Soltis PS, Soltis DE, Burleigh JG. 2014. From algae to angiosperms-inferring the phylogeny of green plants (Viridiplantae) from 360 plastid genomes. BMC Evolutionary Biology 14: 23.
- Rye BL, Barrett RL, Barrett MD, Bruhl JJ, Clarke KL, Wilson KL. 2015. Five new species and a new combination in Cyperaceae from the Kimberley region of Western Australia. *Nuytsia* 26: 167–184.

- Semmouri I, Bauters K, Léveillé-Bourret É, Starr JR, Goetghebeur P, Larridon I. 2019. The phylogeny and systematics of Cyperaceae, the evolution and importance of embryo morphology. *Botanical Review* 85: 1–39.
- Simpson DA. 1992. A revision of the genus Mapania. Richmond: Royal Botanic Gardens, Kew.
- Simpson DA, Inglis CA. 2001. Cyperaceae of economic, ethnobotanical and horticultural importance: A checklist. *Kew Bulletin* 56: 257–360.
- Simpson DA, Furness CA, Hodkinson TR, Muasya AM, Chase MW. 2003. Phylogenetic relationships in Cyperaceae subfamily Mapanioideae inferred from pollen and plastid DNA sequence data. American Journal of Botany 90: 1071–1086.
- Simpson DA, Muasya AM, Chayamarit K, Parnell JAN, Suddee S, De Wilde B, Jones MB, Bruhl JJ, Pooma R. 2005. Khaosokia caricoides, a new genus and species of Cyperaceae from Thailand. Botanical Journal of the Linnean Society 149: 357–364.
- Simpson DA, Muasya AM, Alves M, Bruhl JJ, Dhooge S, Chase MW, Furness CA, Ghamkhar K, Goetghebeur P, Hodkinson TR. 2007. Phylogeny of Cyperaceae based on DNA sequence data—A new rbcL analysis. Aliso 23: 72–83.
- Simpson DA, Yesson C, Culham A, Couch CA, Muasya AM. 2011. Climate change and Cyperaceae. In: Hodkinson T, Jones M, Waldren S, Parnell J eds. Climate change, ecology and systematics. Cambridge: Cambridge University Press. 439–456
- Slater GS, Birney E. 2005. Automated generation of heuristics for biological sequence comparison. *BMC Bioinformatics* 6: 31.
- Smith SA, O'Meara BC. 2012. treePL: Divergence time estimation using penalized likelihood for large phylogenies. *Bioinformatics* 28: 2689–2690.
- Smith SY, Collinson ME, Simpson DA, Rudall PJ, Marone F, Stampononi M. 2009. Elucidating the affinities and habitat of ancient, widespread Cyperaceae: Volkeria messelensis Gen. Et Sp. Nov., a fossil mapanioid sedge from the Eocene of Europe. American Journal of Botany 96: 1506–1518.
- Soros CL, Bruhl JJ. 2000. Multiple evolutionary origins of C₄ photosynthesis in the Cyperaceae. In: Wilson KL, Morrison DA eds. Monocots: Systematics and Evolution. Melbourne: CSIRO. 629–636
- Spalink D, Drew BT, Pace MC, Zaborsky JG, Li P, Cameron KM, Givnish TJ, Sytsma KJ. 2016a. Evolution of geographical place and niche space: Patterns of diversification in the North American sedge (Cyperaceae) flora. Molecular Phylogenetics Evolution 95: 183–195.
- Spalink D, Drew BT, Pace MC, Zaborsky JG, Starr JR, Cameron KM, Givnish TJ, Sytsma KJ. 2016b. Biogeography of the cosmopolitan sedges (Cyperaceae) and the area-richness correlation in plants. *Journal of Biogeography* 43: 1893–1904.
- Spalink D, Pender J, Escudero M, Hipp AL, Roalson EH, Starr JR, Waterway MJ, Bohs L, Sytsma KJ. 2018. The spatial structure of phylogenetic and functional diversity in the United States and Canada: An example using the sedge family (Cyperaceae). *Journal of Systematics and Evolution* 56: 449–465.
- Sponheimer M, Lee-Thorp J, de Ruiter D, Codron D, Codron J, Baugh AT, Thackeray F. 2005. Hominins, sedges, and termites: new carbon isotope data from the Sterkfontein valley and Kruger National Park. *Journal of Human Evolution* 48: 301–312.
- Sponheimer M, Alemseged Z, Cerling TE, Grine FE, Kimbel WH, Leakey MG, Lee-Thorp JA, Kyalo Manthi F, Reed KE, Wood BA, Wynn JG. 2013. Isotopic evidence of early hominin diets. Proceedings of the National Academy of Sciences of the United States of America 109: 10513–10518.
- Starr JR, Jansen FH, Ford BA. 2015. Three new early diverging *Carex* (Cariceae–Cyperaceae) lineages from East and Southeast Asia

with important evolutionary and biogeographic implications. *Molecular Phylogenetics Evolution* 88: 105–120.

- Starr JR, Jiménez-Mejías P, Zuntini AR, Léveillé-Bourret É, Semmouri I, Muasya AM, Baker WJ, Brewer GE, Epitawalage N, Fairlie I, Forest F, Sabino Kikuchi IAB, Pokorny L, Larridon I. 2021. Targeted sequencing supports morphology and embryo features in resolving the classification of Cyperaceae tribe Fuireneae s.l. *Journal of Systematics and Evolution* 59: 809–832.
- Thomas WW. 2020. Two new species of Rhynchospora (Cyperaceae) from Bahia, Brazil, and new combinations in Rhynchospora section Pleurostachys. Brittonia 72: 273–281.
- Thomas WW, Araújo AC, Alves MV. 2009. A preliminary molecular phylogeny of the Rhynchosporeae (Cyperaceae). *Botanical Review* 75: 22–29.
- Thomas WW, Davidse G. 1989. Koyamaea neblinensis, a new genus and species of Cyperaceae (Sclerioideae) from Cerro de la Neblina, Venezuela and Brazil. Systematic Botany 14: 189–196.
- Van der Veken P. 1965. Contribution à l'embryographie systématique des Cyperaceae-Cyperoideae. Bulletin du Jardin botanique de l'État à Bruxelles 35: 285–354.
- Väre H, Kukkonen I. 2005. Seven new species of *Cyperus* (Cyperaceae) section *Arenarii* and one new combination and typification. *Annales Botanici Fennici* 42: 473–483.
- Vellend M, Lechowicz MJ, Waterway MJ. 2000. Environmental distribution of four *Carex* species (Cyperaceae) in an old-growth forest. *American Journal of Botany* 87: 1507–1516.
- Verboom GA. 2006. A phylogeny of the schoenoid sedges (Cyperaceae: Schoeneae) based on plastid DNA sequences, with special reference to the genera found in Africa. Molecular Phylogenetics Evolution 38: 79–89.
- Viljoen J-A, Muasya AM, Barrett RL, Bruhl JJ, Gibbs AK, Slingsby JA, Wilson KL, Verboom GA. 2013. Radiation and repeated transoceanic dispersal of Schoeneae (Cyperaceae) through the southern hemisphere. American Journal of Botany 100: 2494–2508.
- Villa-Machío I, Zamora JC, Sandoval-Sierra JV, Blanco-Pastor JL, Fernández-Mazuecos M, Jiménez-Mejías P. 2020. Insect pollination in temperate sedges? A case study in Rhynchospora alba (Cyperaceae). Plant Biosystems. https://doi.org/10.1080/11263504.2020.1845846
- Villaverde T, Pokorny L, Olsson S, Rincón-Barrado M, Johnson MG, Gardner EM, Wickett NJ, Molero J, Riina R, Sanmartín I. 2018. Bridging the micro- and macroevolutionary levels in phylogenomics: Hyb-Seq solves relationships from populations to species and above. New Phytologist 220: 636–650.
- Villaverde T, Jiménez-Mejías P, Luceño M, Roalson EH, Hipp AL, the Global Carex Group: Wilson KL, Larridon I, Gebauer S, Hoffmann MH, Simpson DA, RFC Naczi, Reznicek AA, Ford BA, Starr JR, Park J, Escudero M, Martín-Bravo S. 2020. A new classification of Carex subgenera supported by a HybSeq backbone phylogeny. Botanical Journal of the Linnean Society 194: 141–163.
- Villaverde T, Maguilla E, Luceño M, Hipp AL. 2021. Assessing the sensitivity of divergence time estimates to locus sampling, calibration points, and model priors in a RAD-seq phylogeny of Carex section Schoenoxiphium. Journal of Systematics and Evolution 59: 687–697.
- Vitelli JS, Madigan BA, van Haaren PE. 2010. Control techniques and management strategies for the problematic Navua sedge (*Cyperus aromaticus*). *Invasive Plant Science and Management* 3: 315–326.
- Vrijdaghs A, Goetghebeur P, Smets E, Muasya AM. 2006. The floral scales in Hellmuthia (Cyperaceae, Cyperoideae) and Paramapania (Cyperaceae, Mapanioideae): An ontogenetic study. Annals of Botany 98: 619–630.

- Wilson KL. 1981. Revision of the genus Mesomelaena (Cyperaceae). Telopea 2: 181–195.
- Wilson KL. 1993. Cyperaceae. In: Harden GJ ed. Flora of New South Wales. Kensington: University of New South Wales Press. 4: 293–396.
- Wragg PD, Johnson SD. 2011. Transition from wind pollination to insect pollination in sedges: experimental evidence and functional traits. New Phytology 191: 1128–1140.
- Wynn JG, Sponheimer M, Kimbel WH, Alemseged Z, Reed K, Bedaso ZK, Wilson JN. 2013. Diet of Australopithecus afarensis from the Pliocene Hadar Formation, Ethiopia. Proceedings of the National Academy of Sciences of the United States of America 109: 10495–10500.
- Yano O, Ikeda H, Watson MF, Rajbhandari KR, Jin X-F, Hoshino T, Muasya AM, Ohba H. 2012. Phylogenetic position of the Himalayan genus Erioscirpus (Cyperaceae) inferred from DNA sequence data. Botanical Journal of the Linnean Society 170: 1–11.
- Yano O, Fuse S, Fujiki T, Tamura MN, Yago M, Sueyoshi M, Yang Y-P, Ikeda H. 2015. Insect Pollination of *Carex* (Cyperaceae) from Yunnan, SW China. *Journal of Japanese Botany* 90: 407–412.
- Yu G. 2019a. ggimage: Use Image in 'ggplot2'. R package version 0.2.1 [online]. Available from https://CRAN.R-project.org/package= ggimage/ [accessed 15 October 2019].
- Yu G. 2019b. treeio: Base classes and functions for phylogenetic tree input and output. R package version 1.8.1 [online]. Available from https://guangchuangyu.github.io/software/treeio/ [accessed 15 October 2019].
- Yu G, Smith D, Zhu H, Guan T, Lam T-Y. 2017. ggtree: An R package for visualization and annotation of phylogenetic trees with their covariates and other associated data. Methods in Ecology and Evolution 8: 28–36.
- Zeng C-X, Hollingsworth PM, Yang J, He Z-S, Zhang Z-R, Li D-Z, Yang J-B. 2018. Genome skimming herbarium specimens for DNA barcoding and phylogenomics. *Plant Methods* 14: 43.
- Zhang C, Rabiee M, Sayyari E, Mirarab S. 2018. ASTRAL-III: Polynomial time species tree reconstruction from partially resolved gene trees. BMC Bioinformatics 19: 153.
- Zhang X, Marchant A, Wilson KL, Bruhl JJ. 2004. Phylogenetic relationships of *Carpha* and its relatives (Schoeneae, Cyperaceae) inferred from chloroplast *trnL* intron and *trnL*–*trnF* intergenic spacer sequences. *Molecular Phylogenetics and Evolution* 31: 647–657.
- Zhang X, Bruhl JJ, Wilson KL, Marchant A. 2007. Phylogeny of *Carpha* and related genera (Schoeneae, Cyperaceae) inferred from morphological and molecular data. *Australian Systematic Botany* 20: 93–106.
- Zuntini AR, Frankel LP, Pokorny L, Forest F, Baker WJ. 2021. A comprehensive phylogenomic study of the monocot order Commelinales, with a new classification of Commelinaceae. American Journal of Botany. https://doi.org/10.1002/ajb2.1698

Supplementary Material

The following supplementary material is available online for this article at http://onlinelibrary.wiley.com/doi/10.1111/jse.12757/suppinfo:

Table S1. Voucher information for accessions included in the targeted sequencing study.

Table S2. Recovery statistics for the genes targeted by the Angiosperms353 probes for the accessions included in this study. **Table S3.** AMAS summary statistics generated for the exons data set. Invariable columns were removed.

Table S4. Overview of the main published classifications of the family Cyperaceae and the classification proposed in this study,

clearly indicating which changes occurred as more data became available. See legend on the right of the table and the included notes.

Fig. S1. Heatmap of recovery of the Angiosperms353 probes for the accessions included in this study.

Fig. S2. Phylogenetic reconstruction of the relationships

in Cyperaceae based on analysis of the exons data set. Concatenated IQ-TREE analysis. Values above branches represent UltraFast Bootstrap support; missing values indicate maximum support. Values below the branches represent gCF/sCF values. Bars on the right indicate subfamilial and tribal classification.