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"The present authors have devoted a total of 50 years to the study of grasses, and present here the first complete reworking of the Gramineae at generic level since 1883 ..."--Preface.

This volume is the outcome of a modern phylogenetic analysis of the grass family based on multiple sources of data, in particular molecular systematic studies resulting from a concerted effort by researchers worldwide, including the author. In the classification given here grasses are subdivided into 12 subfamilies with 29 tribes and over 700 genera. The keys and descriptions for the taxa above the rank of genus

are hierarchical, i.e. they concentrate upon characters which are deemed to be synapomorphic for the lineages and may be applicable only to their early-diverging taxa. Beyond the treatment of phylogeny and formal taxonomy, the author presents a wide range of information on topics such as the structural characters of grasses, their related functional aspects and particularly corresponding findings from the field of developmental genetics with inclusion of genes and gene products instrumental in the shaping of morphological traits (in which this volume appears unique within this book series); further topics addressed include the contentious time of origin of the family, the emigration of the originally shade-loving grasses out of the forest to form vast grasslands accompanied by the switch of many members to C4 photosynthesis, the impact of herbivores on the silica cycle housed in the grass phytoliths, the reproductive biology of grasses, the domestication of major cereal crops and the affinities of grasses within the newly circumscribed order Poales. This volume provides a comprehensive overview of existing knowledge on the Poaceae (Gramineae), with major implications in terms of key scientific challenges awaiting future research. It certainly will be of interest both for the grass specialist and also the generalist seeking state-of-the-art information on the diversity of grasses, the most ecologically and economically important of the families of flowering plants.

Three subtribes of the Bambuseae are present on Sri Lanka, including six genera and 12 species, and they are described both morphologically and anatomically. In the Arundinariinae, five species of fargesioid arundinarias occur: *Arundinaria debilis*, *A. densifolia*, *A. floridunda*, *A. walkeriana*, and the new species *A. scandens*; all are shrubby plants that grow in cool mountain forests and bogs. Three genera of Bambusinae are present. *Bambusa* is represented by three introduced, cultivated species, and *Dendrocalamus* by the new species *D. cinctus*. The new genus *Pseudoxytenanthera* is described, and the single species *P. monadelphica* is unusual in the subtribe in its thin, soft culms produced in open clumps, vine-like habit, and combination of 2-3 stigmas and a branching pattern in which the large central bud remains dormant with the simultaneous production of basal branches, followed by development of the central bud into an elongate whip-like shoot. The Schizostachydinae are represented by *Ochlandra stridula* and *Davidsea attenuata*, the latter a new monotypic segregate of *Teinostachyum* and differing from that genus in its intravaginal branching pattern with three major and several subsidiary branches with the major branches elongating and becoming whip-like, and in its elongate style and plumose stigmas.

Plant molecular biology has produced an ever-increasing flood of data about genes and genomes. Evolutionary biology and systematics provides the context for synthesizing this information. This book brings together contributions from evolutionary biologists, systematists, developmental geneticists, biochemists, and others working on diverse aspects of plant biology whose work touches to varying degrees on plant molecular evolution. The book is organized in three parts, the first of which introduces broad topics in evolutionary biology and summarizes advances in plant molecular phylogenetics, with emphasis on model plant systems. The second segment presents a series of case studies of gene family evolution, while the third

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gives overviews of the evolution of important plant processes such as disease resistance, nodulation, hybridization, transposable elements and genome evolution, and polyploidy.

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