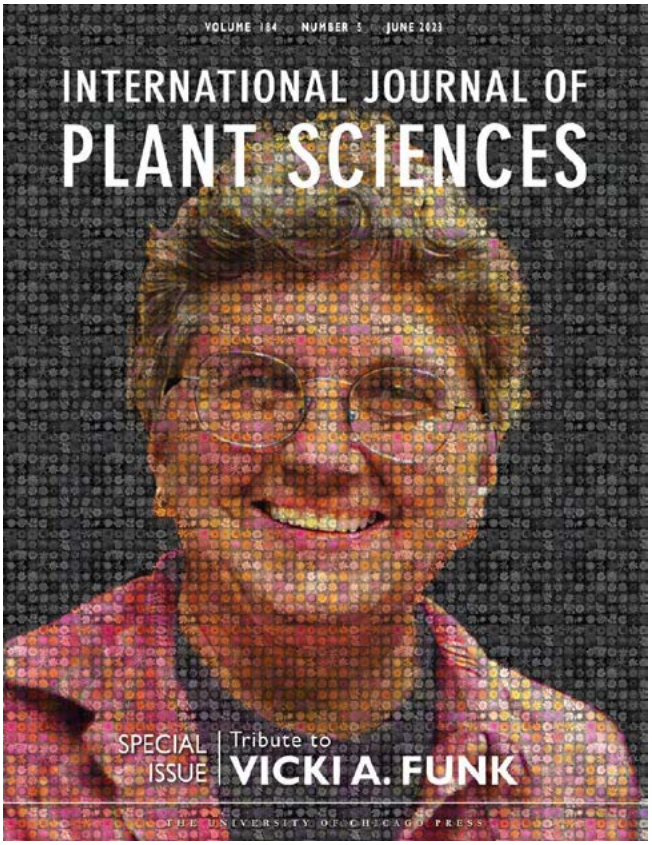
TICATIMES

selected Compositae news and updates from
THE INTERNATIONAL COMPOSITAE ALLIANCE

Edited by Jennifer R. Mandel



JULY 2023

IJPS SPECIAL ISSUE PUBLISHED HONORING VICKI A. FUNK

This July, a special issue was published in the International Journal of Plant Sciences: "A Botanist at the Extreme: Honoring the Great Contributions of Dr. Vicki A. Funk." This issue was guest edited by Jennifer R. Mandel (University of Memphis, USA), J. Mauricio Bonifacino (Universidad de la República, Uruguay), and Erika R. Moore (University of Memphis, USA). This issue was spurred by the symposium at the Botany Meeting in Alaska, 2022, that was hosted in Vicki's honor. Recognizing the achievements from a long and distinguished career by a pioneer in systematic botany, the issue brought together a diverse set of research topics made possible in Compositae through Vicki's generous and collaborative spirit: Biogeography, Collections, Genomics, Palynology, Phylogeny, and Systematics.

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AUGUST 2023

TICA TO HOST SYMPOSIUM AT IBC MADRID IN 2024

In Madrid next year, TICA will host a symposium: "Synantherology reloaded: Recent advances and the future of evolutionary studies in Compositae." Organizers of the symposium are Jennifer R. Mandel (University of Memphis, USA) and J. Mauricio Bonifacino (Universidad de la República, Uruguay). Technological advances in high-throughput sequencing, large-scale phylogenomics, and improved database infrastructure have revolutionized the fields of plant systematics and evolution. The symposium will highlight the impact these technologies have had on our understanding of the biogeography, systematics, and evolution of the world's largest flowering plant family. Speakers will present major findings from the past decade including how new data have led to major changes in classification and have improved data accessibility and infrastructure. The symposium will address the future of studies in the family and what some of the newest technologies, e.g., highly accurate long-read sequencing, CRISPR/functional genetics, and single-cell/nuclei sequencing, will have to offer for further studying the evolution of this fascinating plant group. There are still a couple of slots left in the symposium, reach out to the organizers if you are interested in speaking. We will aim to maximize diversity of speakers and topics.