Unveiling the rapid radiation of *Helianthemum* subg. *Eriocarpum* (Cistaceae) through target capture sequencing

María VALERIO DE ARANA¹, Sara MARTÍN-HERNANZ¹,²,³, Juan VIRUEL³, Polina VOLKOVA⁴, Zhihao SU⁵, Encarnación RUBIO¹, Amelia SHEPHERD-CLOWES³, Rafael G. ALBALADEJO¹, Abelardo APARICIO¹

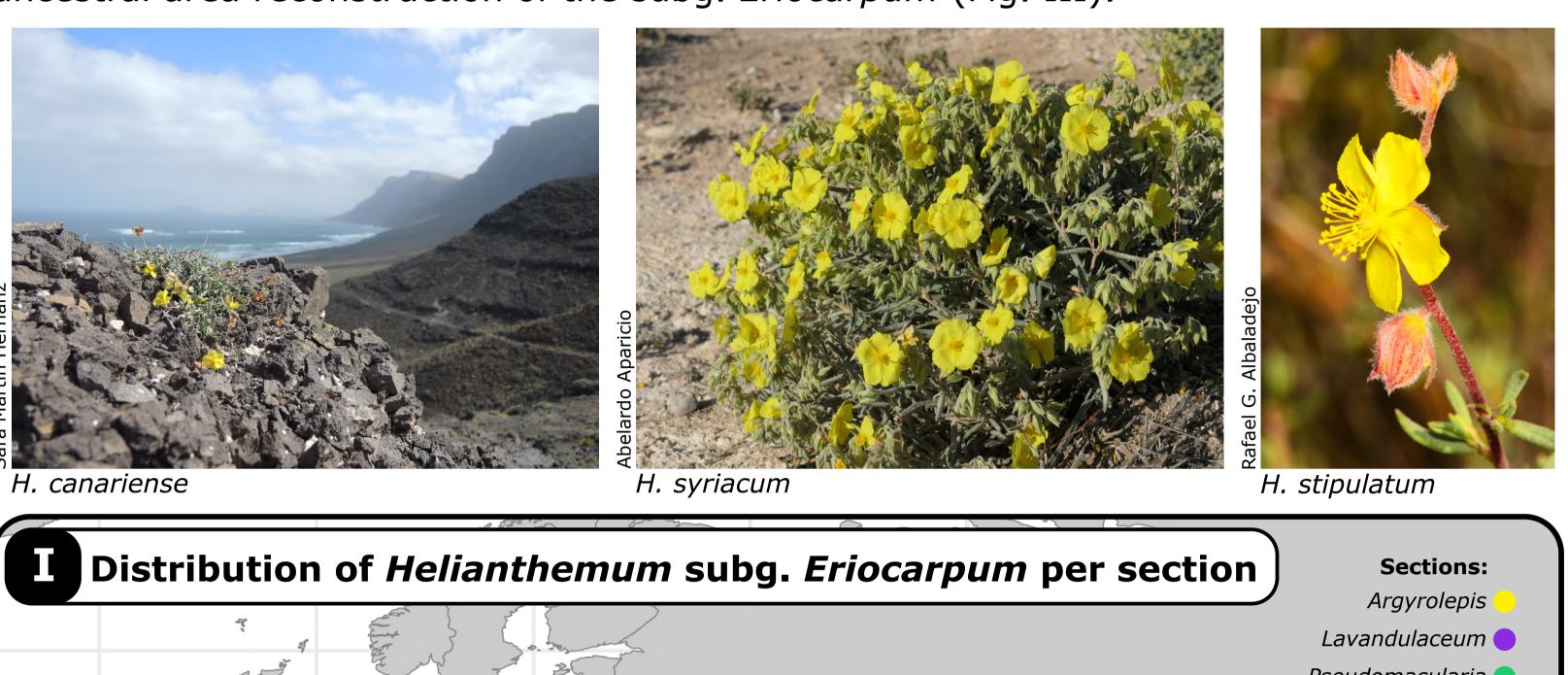
¹Universidad de Sevilla, Seville, Spain ²Universidad Complutense de Madrid, Madrid, Spain ³Royal Botanic Gardens Kew, London, United Kingdom ⁴Russian Academy of Sciences, Borok, Russia

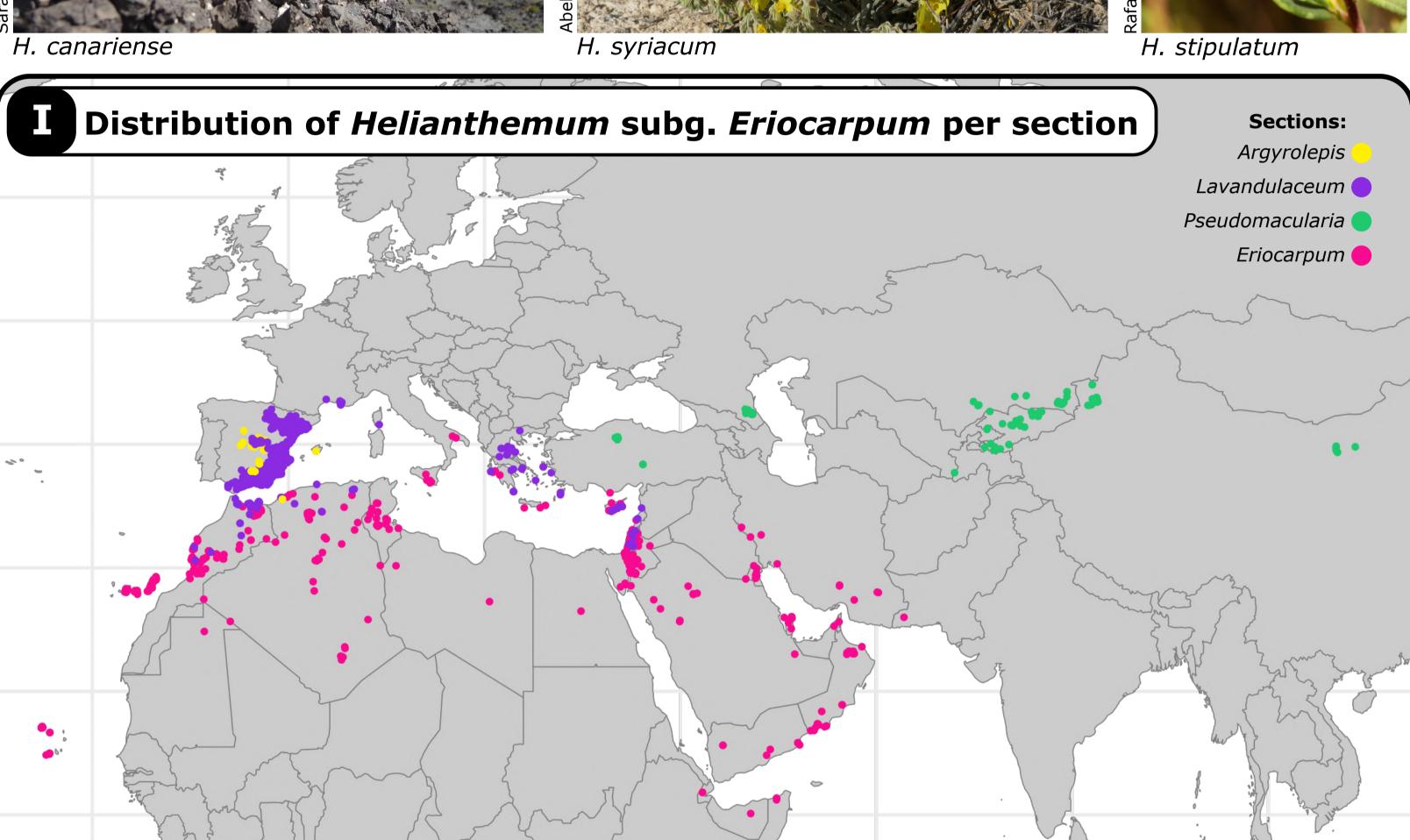
⁵Chinese Academy of Sciences, Beijing, China

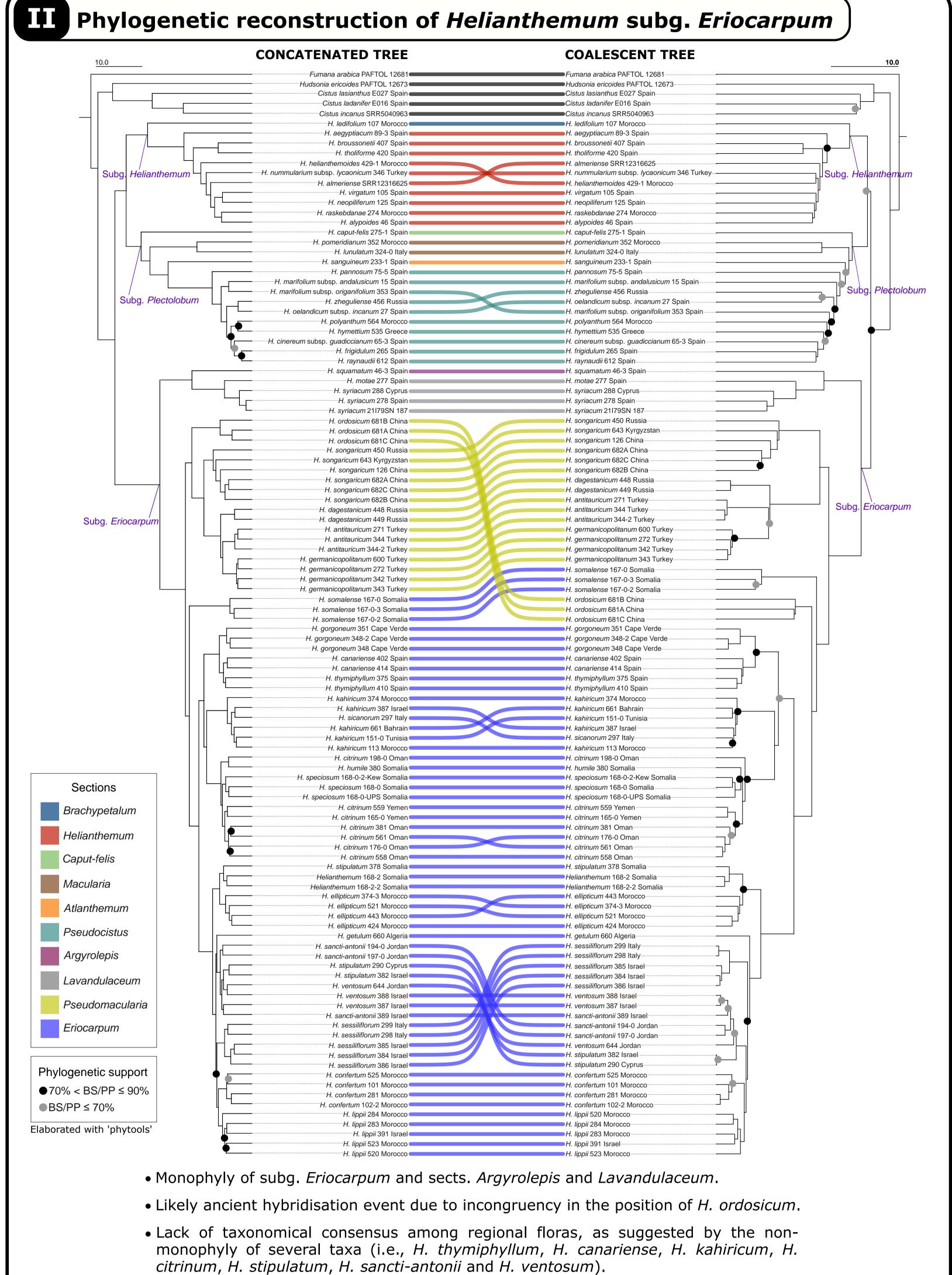
Poster ID: 613 / 2411 mariavaleriodearana@gmail.com

INTRODUCTION: The newly described subg. *Eriocarpum* (Martín-Hernanz et al., 2021) is formed by 32 subdesert specialists with the most widespread and disjunct distribution in the genus *Helianthemum*. It occurs in Macaronesia, the Mediterranean Basin, the Sahara, Arabia, Eastern and Central Asia and the Horn of Africa (Fig. I), where five out of the six species present in northern Somalia are endemic. Previous phylogenetic and biogeographic reconstructions of genus *Helianthemum* generally failed to include these taxonomically valuable endemic species (Aparicio et al., 2017; Martín-Hernanz et al., 2019). Here, we aim to uncover the phylogenetic relationships and biogeographic history of subg. *Eriocarpum* using genomic data and intensive taxon sampling.

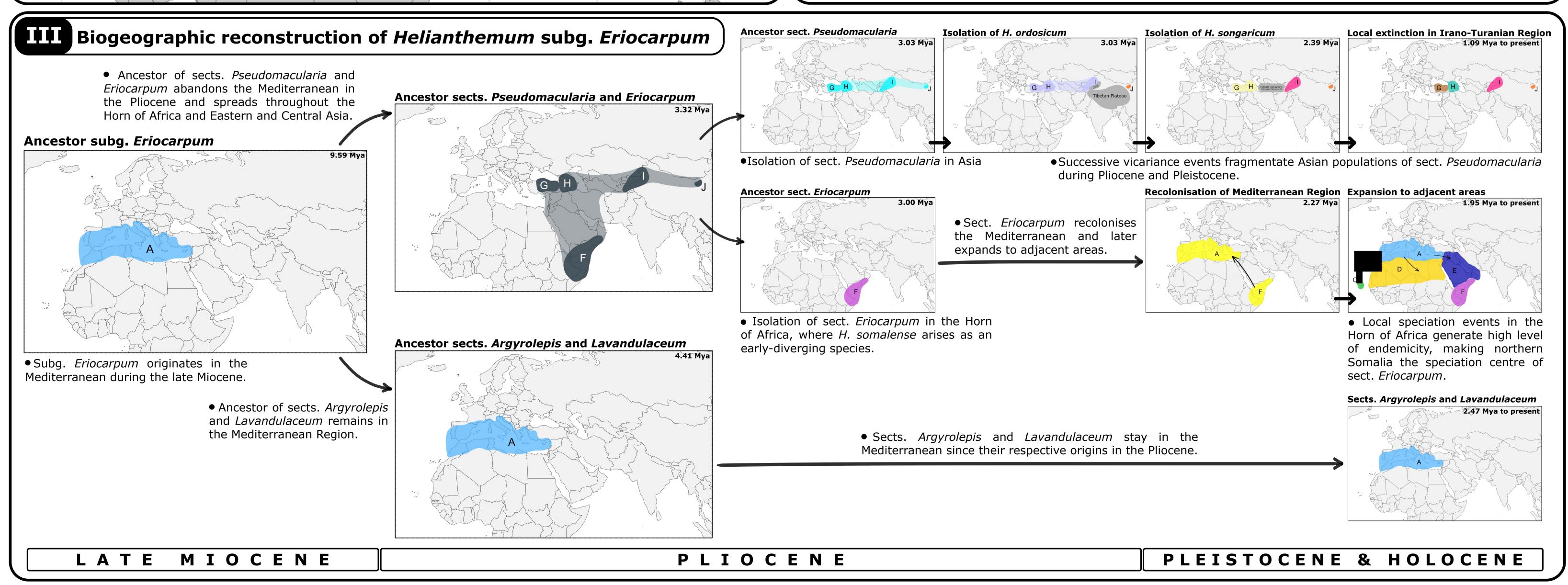
M&M: We obtained 74 samples of subg. *Eriocarpum*, representing the 78% of its taxa, including four of the Horn of Africa endemics, and sequenced them through target capture sequencing with the Angiosperms353 kit. We reconstructed phylogenetic trees using supermatrix (partitioned concatenated matrix) and supertree (coalescent) approaches (Fig. II), estimated the lineage divergence times, and elaborated an ancestral area reconstruction of the subg. *Eriocarpum* (Fig. III).







• New ascription of *H. dagestanicum* to sect. Pseudomacularia.







BOTÁNICO



UNIVERSITE7







REFERENCES:

- ◆ Aparicio A, Martín-Hernanz S, Parejo-Farnés C, Arroyo J, Yeşilyurt EB, Yeşilyurt ML, Yeşilyurt, ML, Rubio E, Albaladejo RG (2017).
 Phylogenetic reconstruction of the genus Helianthemum (Cistaceae) using plastid and nuclear DNA-sequences: Systematic and evolutionary inferences. TAXON 66, 868–885.
- Martín-Hernanz S, Aparicio A, Fernández-Mazuecos M, Rubio E, Reyes-Betancort JA, Santos-Guerra A, Olangua-Corral M, Albaladejo RG (2019). Maximize Resolution or Minimize Error? Using Genotyping-By-Sequencing to Investigate the Recent Diversification of Helianthemum (Cistaceae). Front. Plant Sci. 10, 1416.
- Martín-Hernanz S, Velayos M, Albaladejo RG, Aparicio A, (2021). Systematic implications from a robust phylogenetic reconstruction of the genus *Helianthemum* (Cistaceae) based on genotyping-by-sequencing (GBS) data. *Anales del Jardín Botánico de Madrid* 78.