POLLEN EVOLUTION IN THE GENUS *ECHINOPS* (CARDUEAE, ASTERACEAE): DECIPHERING THE ORIGIN OF GIANT POLLEN GRAINS

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Premise of research. The genus *Echinops* is unique among the Cardueae tribe of Asteraceae for presenting two distinctive features, both related to reproductive structures: a syncephalium (=secondary capitulum) and an impressively large pollen grain with a triangular section and probably the thickest of all plant cell walls. While the syncephalium constitutes a synapomorphy for the genus, recent evidence suggests that some *Echinops* species have pollen similar to that of other Cardueae. This study therefore seeks to contribute insights into the spatiotemporal frame of pollen evolution within the genus.

Methodology. Micromorphological characterization was provided for 35 specimens from 28 *Echinops* species using light and scanning electron microscopy. Pollen counts were carried out for two *Echinops* species and *Cardopatium corymbosum*. Pollen data are discussed in the context of a dated *Echinops* phylogeny. For comparison purposes, new and published pollen data of 622 Cardueae species and 303 taxa of Vernonieae, another tribe where syncephaly has evolved, were collated.

Pivotal results. The "Perennial" *Echinops* pollen type of huge size and triangular section likely derived from the "Annual" *Echinops* pollen type, more similar in shape, size, and exine structure to that of other Cardueae. Pollen type transition took place in the genus long after syncephaly evolved. Pollen size increase did not occur at the expense of pollen quantity and could respond to warmer environmental conditions and increased male competition.

Conclusions. This study of *Echinops* pollen evidenced the evolutionary exploration of novel phenotypic space in the genus, most certainly in response to the climatic context in which the species have diversified.

Keywords: Compositae, exine, light microscopy (LM), scanning electron microscopy (SEM), syncephalium.

Online enhancements: tables.

Introduction

Echinops L. is a genus of ca. 120 species from Eurasia and Africa, classified in the monotypic subtribe Echinopsinae (Cardueae, Asteraceae; Susanna et al. 2020). The genus is notable among Cardueae in that it is the only one to have evolved a syncephalium (=secondary capitulum), which represents a synapomorphy for the taxon. The spherical shape of this syncephalium, its basipetal flowering sequence, and the fact that it is constituted by uniflowered capitula (often with blue flowers, hence the common name "blue thistle"), make *Echinops* species easily recognizable (fig. 1). The genus also stands out for producing impressively large pollen grains, triangular prisms rounded

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at the poles with slightly sunken lateral sides (Mohl 1835), and probably the thickest of all plant cell walls (Gabarayeva et al. 2018). It is likely that both traits (i.e., syncephalium and giant pollen grains) have had a considerable impact on the reproductive biology of *Echinops* species, which are mainly pollinated by insects (Grant 1950), although cases of bird-pollinated species are also reported (Vogel 2015).

Of the published classifications of Asteraceae and Cardueae pollen grains, most consider that *Echinops* pollen should constitute a separate pollen type (Mohl 1835; Stix 1960; Pla Dalmau 1961; Díez 1987; Tormo and Ubera 1988, 1995; Perveen 1999; Jafari and Ghanbarian 2007; Punt and Hoen 2009). However, while *Echinops* has long been model for the study of pollen morphogenesis (e.g., Stix 1964, 1970; Blackmore 1990; Gabarayeva et al. 2018), little attention has been paid to describing and interpreting pollen diversity within the genus. To our knowledge, pollen characterization data are limited to about 20 *Echinops* species (i.e., only ca. 17% of the species; Tormo and Ubera 1988;

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Fig. 1 *Echinops ritro* subsp. *ruthenicus*. Photo of an individual in the field (*left*) and dissection of its syncephalium (*right*), showing 184 uniflowered capitula and the secondary receptacle (protocol described in Fu et al. 2023). Photograph credit: Oriane Hidalgo.

Garnatje and Martín 2007; Osman 2009; Punt and Hoen 2009; Vural et al. 2010; Kallajxhiu et al. 2014; Tahmasebi et al. 2022). Nevertheless, the few data available revealed a particularly wide range of pollen size, from polar axis (P) of 14.50 to >100 μ m (Erdtman 1952; Tormo and Ubera 1988; Garnatje and Martín 2007; Tahmasebi et al. 2022). Garnatje and Martín (2007) consider that pollen grains of the lower size range, corresponding to annual Echinops species, should constitute a different type, which they call the "Annual" Echinops pollen type. This pollen type characterized by a P <50 μ m and a circular section in polar view shows affinities with other pollen types in the tribe Cardueae (e.g., the crupina type; Wagenitz 1955). The pollen of P >50 μ m with triangular section and prominent bridge-shaped intercolpia, exclusive to Echinops and in particular its perennial species, constitutes the "Perennial" Echinops pollen type (Garnatje and Martín 2007).

Whether the arising of giant pollen grains could be related to syncephaly is unknown, although one would certainly expect the evolution of these two highly distinctive features of Echinops, both related to reproductive structures, to be linked to some extent. A temporal frame of pollen type transitions within the genus is lacking; however, it was suggested that the Annual Echinops type could have evolved in the most recent common ancestor of *Echinops* species, while the Perennial *Echinops* type could have evolved when perennial species diverged (Sánchez-Jiménez et al. 2010). If this were the case, this would mean that the pollen of huge size and triangular section evolved much after the syncephalium. Nevertheless, the possibility that the Annual Echinops type derived from the Perennial type cannot be totally excluded, and its similarities with other Cardueae pollen would then be convergences due to sharing a short growth cycle (e.g., as found in some annual/biennial, high-mountain, or invasive species; Hidalgo et al. 2008b). Regarding the possible impact of the inflorescence context on pollen evolution, the examination of pollen traits in other syncephalous Asteraceae could shed some light on the subject. The tribe Vernonieae is particularly interesting in this respect, as it includes representatives with syncephalia resembling those of *Echinops* (e.g., *Chresta* Vell. ex DC.; Siniscalchi et al. 2017).

The aims of this study were to (i) assess the extent of pollen diversity in *Echinops*, (ii) provide insights into pollen type transitions, and (iii) address the evolutionary and functional significance of pollen traits in the genus.

Material and Methods

Plants

Floral buds were obtained from herbarium specimens (table 1) and preserved in paper envelopes until processed for pollen examination by light microscope (LM) and scanning electron microscope (SEM). Sampling covers the nine Echinops sections whose circumscription was verified and redefined by Sánchez-Jiménez et al. (2010; i.e., sects. Acantholepis (Less.) Jaub. & Spach, Chamaechinops Bunge, Echinops, Hamolepis R.E.Fr., Hololeuce Rech.f., Oligolepis Bunge, Phaeochaete Bunge, Psectra Endl., and Ritropsis Greuter & Rech.f.), along with one representative of sect. Cenchrolepis Hochst. (E. hispidus Fresen) and two tropical African species, E. eryngiifolius O.Hoffm. and E. gracilis O.Hoffm., that are not attributed to any section. For pollen counting, fresh plant material was collected from cultivated plants at the Botanical Garden of Barcelona and the Royal Botanic Gardens, Kew. The five mature, undehisced anthers of a floral bud were collected and stored in an open PCR tube in a box with silica gel and left to dry for at least 1 wk.

Light Microscopy

Pollen was acetolyzed following the micromethod described by Avetissian (1950), mounted on glycerogelatine, and sealed with nail polish. Pollen plates were photographed with a Zeiss AxioCam HRm digital camera mounted on a Zeiss Axioplan microscope, and images were analyzed with AxioVision LE version 4.8.2. Depending on the specimen, 7–50 fully developed Table 1

Pollen Size, Shape, and Type of <i>Echinops</i> Accessions, with Indication of Collection Data							
Taxon	$P \pm SD (\mu m)$	E \pm SD (μ m)	P/E	Shape	Туре	п	Accession data
E. albicaulis	88.80 ± 7.3	72.99 ± 6.1	1.22	SP	EA	15	Kazakhstan; Goloskokov s.n. (GB)
E. amplexicaulis	80.21 ± 6.3	51.95 ± 4.7	1.55	Р	Pe	13	Cameroon; Jacques-Felix 9083 (YA)
E. bannaticus	72.12 ± 4.8	55.22 ± 4.0	1.31	SP	Pe	23	Montenegro; Garnatje & Sánchez-Jiménez 14 (BC)
E. cephalotes	78.77 ± 3.5	63.63 ± 4.6	1.24	SP	Pe	50	Iran; Babakanlou & Amin s.n., 13.VII.1973 (W 1973-05105)
E. cornigerus	99.62 ± 5.8	71.50 ± 5.2	1.40	Р	Pe	39	Afghanistan; Grey-Wilson & Hewer 1656 (GB)
E. davuricus	64.01 ± 3.7	46.10 ± 3.2	1.39	Р	Pe	47	China; Liu, Cao & Vallès s.n., 29.VIII.2007 (BC)
E. elbursensis					EA		Iran, Tehran; 25.VIII.1975 (W)
E. echinatus	76.45 ± 4.5	61.43 ± 3.7	1.25	SP	Pe	46	Pakistan; (W 1979-05252)
E. emiliae	61.21 ± 4.3	57.07 ± 4.5	1.07	Sph	EA	44	Turkey; Ertuğrul, Garcia-Jacas & Susanna 2262 (BC)
E. eryngiifolius	103.80 ± 5.9	64.28 ± 5.6	1.62	P	Pe	35	Rwanda; Raynal 20690 (B 10 0024097)
E. exaltatus	66.89 ± 5.4	55.75 ± 5.6	1.20	SP	Pe	50	Slovenia; Garnatje & Sánchez-Jiménez 6 (BC)
E. fastigiatus	$21.84 \pm .96$	$20.19 \pm .90$	1.08	Sph	An	31	Kazakhstan; Goloskokov s.n. (LE)
E. fontqueri	103.70 ± 3.5	74.98 ± 2.8	1.38	Р	Pe	15	Morocco; Gómiz s.n., 19.V.2002 (BC)
E. freitagii	55.14 ± 2.1	44.13 ± 3.5	1.26	SP	Pe	40	Iran, Khorasan; 1.VIII.1975 (W 1979-07328)
E. gmelinii	41.91 ± 3.7	36.36 ± 2.9	1.15	SP	An	7	China; Vallès, Yan &. Zhao s.n., 2.IX.2007 (BC)
E. gracilis	67.69 ± 6.1	50.71 ± 5.9	1.34	Р	Pe	48	Cameroon; Letouzey 8784 (YA)
E. graecus	67.01 ± 3.7	61.77 ± 3.6	1.09	Sph	Pe	27	Greece; Garnatje & Sánchez-Jiménez 35 (BC)
E. griffithianus	103.3 ± 3.6	76.70 ± 4.3	1.35	Р	Pe	42	Afghanistan; Jörgensen 569 (GB)
E. hoehnelii	80.65 ± 4.1	51.19 ± 4.1	1.58	Р	Pe	45	Kenya; Galbany & Arrabal s.n., 10.IX.2006 (BC)
E. hololeucus	58.14 ± 3.7	46.49 ± 3.2	1.25	SP	Ho	37	Afghanistan; 34°00'N, 64°45'E, 27.VII.1962 (W 1972-19421)
E. humilis	33.67 ± 1.1	37.49 ± 1.0	.90	Sph	An	47	Russia; Tipokhina & Danilyuk s.n., 10.VIII.1973 (LE)
E. hystrichoides	99.22 ± 3.2	70.24 ± 4.3	1.41	Р	Pe	29	Saudi Arabia; Collenette 1570 (E00276929)
E. hystrichoides	101.10 ± 7.3	70.84 ± 6.1	1.43	Р	Pe	21	Yemen; Hein 3806 (BC)
E. hystrichoides	135.02 ± 14.9	94.33 ± 7.0	1.43	Р	Pe	35	Yemen; Hein 3806 (B 10 0220615)
E. hystrichoides	96.95 ± 5.4	71.02 ± 7.0	1.37	Р	Pe	35	Yemen; Hein 3806 (B 10 0220613)
E. hystrichoides	122.34 ± 6.9	97.23 ± 11.4	1.26	SP	Pe	16	Yemen; Hein 3942 (B 10 0220608)
E. integrifolius	26.34 ± 1.60	24.05 ± 1.07	1.09	Sph	An	15	China; (PE 00554105)
E. longifolius	78.15 ± 4.7	48.23 ± 6.9	1.62	Р	Pe	16	Togo; Hein & Pircher 772 (B 10 0293577)
E. przewalskyi	$68.82~\pm~2.7$	51.21 ± 3.5	1.35	Р	Pe	43	China; Vallès & Zhao s.n., 5.IX.2007 (BC)
E. spinosissimus	125.14 ± 6.2	88.22 ± 6.7	1.42	Р	Pe	7	Somalia; Bally 9941 (G00160590)
E. spinosissimus	94.64 ± 6.0	66.36 ± 2.8	1.42	Р	Pe	16	Yemen; Hein 6936 (B 10 0220616)
E. spinosissimus	95.85 ± 4.9	74.00 ± 5.9	1.30	SP	Pe	32	Chad; Pappi 131 (G00160589)
E. spinosissimus subsp.	145.50 ± 4.1	106.30 ± 4.1	1.37	Р	Pe	39	Greece; Rechinger12702 (W)
spinosissimus							
E. talassicus	76.24 ± 3.7	58.64 ± 4.3	1.30	SP	Pe	32	Kazakhstan; Ivaschenko, Susanna 2180 & Vallès (BC)
E. transcaucasicus					Pe	•••	Armenia; Avetisian, Arevschatian & Pogosian s.n., 30.VIII.1979 (W)

Note.	P = polar axis;	E = equatoria	il axis. Shape: Sph	= spheroidal P/E	[0.88-1.14); SP =	= subprolate P/E	E [1.14–1.33); P	= prolate P/E
[1.33–2).	Pollen type: An =	= annual; EA =	Echinate-Annual;	Ho = Hololeucus	; $Pe = Perennial. n$	n = number of p	pollen grains me	asured by LM.

grains were measured. The parameters considered were polar and equatorial (E) diameters and shape (P/E ratio). For pollen grains with a triangular cross section, we consider that the diameter E corresponds to the length of the side.

Scanning Electron Microscopy

We followed the procedure of Halbritter (1998) with slight modifications. Pollen grains from herbarium vouchers were rehydrated by placing the anthers into an Eppendorf tube filled with pure water and further dehydrated overnight in 30 mL of 2,2-dimethoxypropane (DMP) acidified with a drop of 0.2 M HCl. Anther tissues were subsequently removed, and the tube was centrifuged before eliminating the DMP. Pollen grains were criticalpoint dried in CO₂, using ethanol as intermediate fluid, further mounted on stubs, sputter-coated with graphite, and observed with an ESEM Quanta 200 FEI XTE 325/D8395 at high-vacuum conditions at the Centres Científics i Tecnològics of the Universitat de Barcelona. In the case of *E. elbursensis* Rech.f. and *E. transcaucasicus* Iljin, pollen grains were acetolyzed as described above, gold-coated with diode sputtering, and observed under a Hitachi 52300. Pollen description followed the nomenclature recommendations of Punt et al. (2007).

Pollen Counts by Flow Cytometry

We used lycopod spore tablets as a standard for pollen counting (9666 \pm 671 spores per capsule; batch 3862; Department of Quaternary Geology, Lund University, Lund, Sweden) following the protocol of J. M. de Vos, Y. Woudstra, I. J. Leitch, and O. Hidalgo (personal communication). Samples were analyzed using a CyFlow SL3 flow cytometer (Partec, Münster, Germany) fitted with a 100-mW green solid-state laser (Cobolt Samba, Solna, Sweden). For each species, we analyzed two flowers per individual, with three technical replicates each.

Divergence Time Estimation

Sequences of nr-ITS and cp-trnLF regions were gathered from GenBank and edited with BioEdit (ver. 7.2.3; Hall 1999) to constitute a data set of 97 taxa, including 85 Echinops and 12 outgroups species. Alignment was carried out on GUIDANCE web server (ver. 1.5; Penn et al. 2010) using MAFFT multiple alignment algorithm (ver. 7; Katoh and Standley 2013), with 100 bootstrap repeats, a max iterate of 1000, and a local pair refinement strategy. Divergence times were estimated with BEAST (ver. 2.1.3; Bouckaert et al. 2014) using the combined ITS and trnLF data set. A minimum age constraint of 47.5 mya was applied based on the fossil pollen grain Mutisiapollis telleriae for calibration (Palazzesi et al. 2022b and references therein). Each gene was analyzed under separate substitution models selected using the Akaike information criterion implemented in jModelTest (ver. 2.1.4; Darriba et al. 2012)-namely, SYM+I+G for ITS and GTR+G for trnLF-while assuming the clock and tree were linked. Uncorrelated, lognormal relaxed clock model and birth-death tree prior model were specified.

Tribe-Level Surveys

New and published P and E length data were collated for 997 accessions of Cardueae (see raw data in table A1). Plant names were updated primarily using the Plants of the World Online and Euro+Med Plantbase databases (Greuter 2006; POWO 2022), leading to a total of 622 species represented. P and E values were then averaged per species. Subtribe assignment follows the circumscriptions of Herrando-Moraira et al. (2019, 2020). For comparative purposes, we also compiled data for another tribe, the Vernonieae, that includes representatives with a syncephalium of globular shape and a small number of flowers per capitulum resembling that of Echinops (e.g., Chresta; Siniscalchi et al. 2017) as well as other types of syncephalium (e.g., Eremanthus Less [Loeuille et al. 2012]; Paralychnophora MacLeish [Souza-Souza et al. 2016]). P and E length data were gathered from the literature for 303 taxa of Vernonieae (table A2), together with information on whether these taxa were syncephalous.

Data Visualization and Analysis

Data visualization, processing, and analysis were carried out in R (ver. 3.2.2; R Core Team 2019) using the package ggplot2 (Wickham 2016) and the function ace in phytools (Revell 2012) for ancestral state reconstruction.

Results

This study contributed pollen measurements (images not shown; table 1) and SEM views (figs. 2, 3) for 35 specimens from 28 *Echinops* species. Our results further extend the already remarkable pollen size diversity known in the genus, to a range of P length spanning 1 order of magnitude, up to 145.5 μ m in *E. spinosissimus* Freyn subsp. *spinosissimus* (table 1). The study provides the first pollen count data for the genera *Cardopatium* Juss. and *Echinops*, with mean values of 2669.84 ± 754.01 for *C. corymbosum* (L.) Pers., 4716.34 ± 651 for *E. pappii* Chiov., and 4142.95 ± 321.90 for *E. sphaerocephalus* L. Details of pollen count data are provided in table A3.

We collated new and previously published pollen data for about 50 *Echinops* species, which were then used to assign pollen type—either Annual or Perennial—following the description of Garnatje and Martín (2007). However, the pollen of some species did not correspond to either type, which led us to describe two new pollen types: the Echinate-Annual and the Hololeucus pollen types (description provided in table 2). It is worth noting that the affinities of Annual, Echinate-Annual, and Hololeucus pollen types, while differing in size (fig. 4) and exine ornamentation (figs. 2, 3), are much closer to each other than to the Perennial type (table 2). The latter is clearly distinguished by its peculiar shape and its exceptionally complex exine (Gabarayeva et al. 2018; table 2).

The mapping of pollen types onto the phylogeny shows an almost complete correspondence to *Echinops* sections (fig. 4). Indeed, excluding two species, each section is characterized by a single pollen type: Annual for sections *Acantholepis* and *Chamaechinops*, Echinate-Annual for sect. *Psectra*, Hololeucus for sect. *Hololeuce*, and Perennial for the remaining sections (table 2; fig. 4). The two exceptions are *E. strigosus* in sect. *Psectra*, with Annual type, and *E. albicaulis* Kar. & Kir. in sect. *Echinops*, with Echinate-Annual type. This attests to the low evolutionary lability of pollen type despite the great diversity of pollen size and shape displayed by the genus.

The ancestral state reconstruction suggests that *Echinops* Annual pollen type characterizes the backbone of the genus, while *Echinops* Perennial pollen type evolved in the common ancestor of the clade comprising sects. *Echinops*, *Hamolepis*, *Oligolepis*, *Phaeochaete*, and *Ritropsis*, 6.10 mya (node marked with a B in the chronogram of fig. 4).

A general trend in pollen evolution in *Echinops* is an increase in size, especially of the P-axis (fig. 4) that varies over a 10.03-fold range and reaches in sect. *Ritropsis* and *Oligolepis* the largest values for the Cardueae (fig. 5A, 5B). Changes in size are also accompanied by changes in shape, the smallest grains being spheroidal or subprolate and the largest ones being prolate (fig. 5A). In comparison, such size and shape changes are not observed in the tribe Vernonieae (fig. 5C, 5D). Indeed, the range of P-axis values in this tribe is only 4.30-fold, and all of the 303 taxa studied so far have spheroidal or suboblate pollen (fig. 5C).

Discussion

Evolution of Echinops within Cardueae: Climatic Context

The phylogenomics hypothesis of Mandel et al. (2019) places *Echinops* as a sister group to the remaining Cardueae, with the exception of *Carlina* L. and *Cardopatium*; these last two taxa form a clade sister to the remaining Cardueae. The tribe Cardueae may have diverged from the rest of Asteraceae about 40 mya in western Asia according to dating analyses and ancestral area reconstructions (Barres et al. 2013; Mandel et al. 2019). We inferred here that the origin of Cardueae occurred during a global warming event (i.e., Mid-Eocene Climatic Optimum [MECO]), which has been recorded throughout the world (Bijl et al. 2010). Aridification appears to have intensified during and after the MECO, particularly in places today occupied by massive deserts such as the Tibetan Plateau (Bosboom et al. 2014). The crown Cardueae—that is, the most recent common ancestor of the tribe—evolved about 36 mya, during the Late Eocene, when



Fig. 2 Scanning electron microscopy views of pollen grains for *Echinops* species. A–C, *Echinops albicaulis*. D–G, *Echinops bannaticus*. H, I, Echinops cephalotes. J, K, Echinops cornigerus. L, M, Echinops elbursensis. N, O, Echinops emiliae. P, Echinops eryngiifolius. Q, R, Echinops exaltatus. S–U, Echinops fontqueri. V, W, Echinops freitagii. X, Echinops gmelini.



Fig. 3 Scanning electron microscopy views of pollen grains for *Echinops* species. A–C, *Echinops* gracilis. D–F, *Echinops* graecus. G–I, *Echinops* griffithianus. J, K, *Echinops hoehnellii*. L–N, *Echinops hololeucus*. O, P, *Echinops humilis*. Q–S, *Echinops hystricoides*. T, *Echinops latifolius*. U, V, *Echinops longifolius*. W, X, *Echinops przewalskyi*. Y–AB, *Echinops spinosissimus* subsp. spinosissimus. AC–AE, *Echinops talassicus*. AF, AG, *Echinops transcaucasicus*.

Table	2
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Echinops pollen type	Description	Taxa included
Annual	Pollen grains small or medium, $P < 50 \ \mu$ m, spheroidal to subprolate; exine ornamentation microechinated with supratectal pointed elements from 0.10 to 0.15 μ m, which are regularly distributed; ² circular section in polar view	Sect. Acantholepis: E. acantholepis, ¹ E. gmelinii, E. nanus; ¹ sect. Chamaechinops: E. fastigiatus, E. humilis, E. integrifolius; sect. Psectra: E. strigosus ¹
Echinate-Annual	Pollen grains medium or large, $20 < P < 100 \ \mu m$, spheroidal to subprolate, resembling the Annual type but with ornamentation echinated consisting of spines with rounded tips; circular section in polar view	Sect. Echinops clade VIII: E. albicaulis; sect. ¹ Psectra: E. elbursensis, E. emiliae
Hololeucus	Pollen grains of medium size, P around 60 µm, subprolate in shape; exine ornamentation consisting of sparsely distributed spines with rounded tips; circular section in polar view	Sect. Hololeuce: E. hololeucus
Perennial	Pollen grains large or very large, $P > 50 \ \mu m$, subprolate to prolate shape, in few cases sphe- roidal (e.g., <i>E. graecus</i> , <i>E. transcaucasicus</i>); exine ornamentation microechinated/echinated- verrucated; intercolpal zone raised, producing a pronounced bridge; triangular section in polar view ²	 Sect. Cenchrolepis: E. giganteus,¹ E. hispidus;¹ sect. Echinops clade VIII: E. bannaticus, E. davuricus, E. exaltatus, E. freitagi, E. graecus, E. microcephalus,¹ E. obliquilobus,¹ E. ossicus,¹ E. przewalskyi, E. ritro,¹ E. sphaerocephalus,¹ E. talassicus; sect. Echinops clade IX: E. transcaucasicus; sect. Hamolepis: E. hoehnelii; sect. Oligolepis: E. cornigerus, E. dumanii,² E. echinatus, E. griffithianus, E. phaeocephalus,² E. polygamus;³ sect. Phaeochaete: E. amplexicaulis, E. cephalotes, E. longifolius, E. mildbraedii;¹ sect. Ritropsis: E. fontqueri,¹ E. glaberrimus,⁴ E. hystrichoides, E. machrochaetus,⁴ E. macrophyllus,⁵ E. spinosissimus, E. tenuisectus;³ unknown or doubtful sect.: E. chlorophyllus,^{6,a} E. eryngiifolius, E. galalensis,⁴ E. gracilis, E. hussonii,⁴ E. robustus,^{6,a} E. saissanicus¹

Pollen Types in the Genus Echinops

Sources. 1 = Garnatje and Martín (2007); 2 = Vural et al. (2010); 3 = Tahmasebi et al. (2022); 4 = Osman (2009); 5 = Jafari and Ghanbarian (2007); 6 = Ikuse (1962).

Note. In one case, for *E. taeckholmianus* (Osman 2009), we were unable to assign a pollen type. P = polar axis.

^a Echinops chlorophyllus Rech.f. and E. robustus Bunge have been traditionally ranged into sect. Oligolepis. However, a very similar species, E. cephalotes, was removed from this section and placed in sect. Phaeochaete (Sánchez-Jiménez et al. 2010). These species should likely be reassigned to sect. Phaeochaete; however, pending molecular phylogenetic data, we chose to let them be unassigned.

ephemeral continental-scale glaciations occurred in Antarctica (Van Breedam et al. 2022), anticipating a more widespread glaciation event at the Eocene-Oligocene transition (34.44-33.65 mya). This period of a large temperature drop across the Eocene-Oligocene transition marks the onset of the modern "icehouse" world. In this context, the family Asteraceae as well as other grasslands' components (e.g., Poaceae) started to rise in diversification linked to a global decrease in CO₂ (Palazzesi et al. 2022a). Also, Echinops may have split from its sister group during this climatic scenario, although previously published data suggested a later origin of the genus, during the Miocene (Montazerolghaem et al. 2017). The lack of fossil records assigned to Echinops prevents any substantial interpretation about the origin and diversification of the genus. In line with this, Partridge (1978) recorded some fossil pollen grains that he claimed to belong to Echinops or Mutisia L.f. from Neogene offshore sediments in western Africa. However, the overall morphology of the pollen grains-for example, the size and distribution of the spines-appears not to be related to either Echinops or Mutisia but to other Cardueae. The unquestionable Quaternary record of Echinops, however, allows us to better interpret the climatic context in which the genus grew and expanded. For example, dispersed pollen grains recorded by Jiang et al. (2014) in China and deposited during the Last Glacial Maximum, about 20,000 years ago, lead them to suggest that *Echinops*, associated with other arid-adapted shrubs and herbs (e.g., *Taraxacum* F.H.Wigg. and Chenopodiaceae), bloomed during very dry conditions at the expense of humid-demanding taxa (e.g., grasses). Further interpretations of the climatic context during the early evolution of *Echinops* (late Paleogene–early Neogene) await new fossil discoveries.

Evolution of Pollen in Echinops: Decoupled from That of the Syncephalium and from Any Polyploidization Event

Our results show that the Perennial pollen type appeared 6.10 mya at the earliest, well after the syncephalium, which evolved in the most recent common ancestor of all *Echinops* 15.11 mya (fig. 4). Previous studies suggest that one or more polyploidization events may have accompanied the evolution of these characters in *Echinops* (Sánchez-Jiménez et al. 2009). Polyploidy and subsequent genome diploidization mechanisms are widely recognized as key players promoting evolutionary diversification (e.g., Dodsworth et al. 2016; Landis et al. 2018). However, mixed evidence has been found for their effect as drivers of pollen morphological evolution (Jardine et al. 2022), including pollen size (reviewed in Sanders 2021).



Fig. 4 Distribution of mean polar axis (P) values per species and pollen types across the different sections of *Echinops* (following the taxonomic circumscription of Sánchez-Jiménez et al. 2010). Ancestral state reconstruction results of pollen type are presented as pie charts at selected nodes of the dated phylogeny. Branch support is provided for *Echinops*, with bold branches indicating a posterior probability \geq 95%. Photos (at the same scale for a better comparison) illustrate the different types of pollen in the genus: *Echinops albicaulis* for Echinate-Annual type (*a*), *E. spinosissimus* subsp. *spinosissimus* for Perennial type (*b*), *E. hololeucus* for Hololeucus type (*c*), and *E. gmelinii* for Annual type (*d*).

The most frequent chromosome numbers in *Echinops* are superior or equal to 30 and considered to correspond to diploidized polyploids (Garnatje et al. 2004). Nevertheless, some annual species have much lower chromosome numbers, the most extreme case being *E. acantholepis* Jaub. & Spach with 2n = 14. Consequently, two possible configurations were proposed (Sánchez-Jiménez et al. 2009). One situates a polyploidization event at the base of the genus with subsequent drastic dysploidy in the clade of annual species. The other involves several independent polyploidization events (at least four) that affected all but a few annual species. Unexpectedly, the most recent developments in transcriptomics and phylogenomics support another configuration, where the last polyploidization event in the evolutionary

history of *Echinops* would be the one common to all Asteraceae except Barnadesieae (Zhang et al. 2021; Palazzesi et al. 2022b).

The pollen size increase observed in *Echinops* Perennial type is then probably not related to polyploidization; however, its decrease in the annual species may well be associated with other genomic processes that impact on genome size. Indeed, genome size correlates with cell cycle duration, and annual species that are generally fast growing therefore tend to have smaller genomes than their perennial relatives (in *Echinops*; Garnatje et al. 2004; Sánchez-Jiménez et al. 2009). This downsizing is achieved through genomic restructuring, which often also results in chromosome number changes. Since genome size also correlates with cell size (Beaulieu et al. 2008), genome downsizing would then result



Fig. 5 Diversity of pollen shape and size in 622 species of Cardueae (*A*, *B*) and 303 species and subspecies of Vernonieae (*C*, *D*). P indicates polar axis; E indicates equatorial axis. Dashed lines indicate P/E ratio of 0.88, 1.14, 1.33, and 2, which delimitate the suboblate P/E [0.75–0.88), spheroidal P/E [0.88–1.14), subprolate P/E [1.14–1.33), prolate P/E [1.33–2), and perprolate P/E ≥ 2 shape in the morphospace. *B*, *D*, Boxplots with individual jitter values depicting the distribution of P size. Colors correspond to subtribes for Cardueae (all presenting a simple capitulum except the Echinopsinae and possibly the Dipterocominae, with a syncephalium) and to the inflorescence type (simple capitulum vs. syncephalium) for Vernonieae. A picture of a syncephalous representative is provided for each tribe: *Echinops* (Cardueae) and *Chresta* (Vernonieae). Photograph credits: Oriane Hidalgo (*Echinops ritro*), Carolina M. Siniscalchi (*Chresta sphaerocephala*).

in smaller cells, including pollen. *Echinops acantholepis* and *E. gmelinii* Turcz. illustrate this trend, where annual taxa show both genome downsizing—here accompanied by dysploidy—and reduced pollen size, as previously found in other Asteraceae such as in *Callicephalus* C.A.Mey., *Centaurea* L., and *Oligochaeta* K.Koch (Hidalgo et al. 2008*a*, 2008*b*) within Cardueae and in *Reichardia* Roth (Siljak-Yakovlev et al. 2017).

Increase of Pollen Size in Echinops Perennial Pollen Type Did Not Occur at the Expense of Pollen Quantity

Because of the existence of a trade-off between the number and size of pollen grains (Cruden and Miller-Ward 1981; Vonhof and Harder 1995), it is important to also consider the quantitative aspects of *Echinops* pollen evolution. We carried out pollen counts to find out whether *Echinops* species with Perennial pollen type have a particularly low amount of pollen per flower. Furthermore, reproductive systems are reflected in male and female investments through the pollen-to-ovule ratio (Cruden 1977; Cruden 2000), and the number of pollen grains in *Echinops* species could therefore provide information on where they fall on the spectrum of self-compatibility.

Echinops sphaerocephalus presented a mean of 4142.95 pollen grains per flower, suggesting that the pollen-to-ovule ratio is maintained at a level corresponding to xenogamy (Cruden 1977) despite a huge pollen size. This fits our expectations, given our unsuccessful attempts to reproduce *Echinops* species in the past. Unfortunately, there is little data to compare these results with, as the breeding system of *Echinops*—and even more generally that of Cardueae-has so far been poorly studied (East 1940). Previous reports of pollen counts in the tribe show mean values per flower of 5895 for Berardia lanuginosa (Lam.) Fiori & Paol. (facultative xenogamy; Guerrina et al. 2016), ca. 850 for Carduus baeocephalus Webb subsp. baeocephalus (facultative xenogamy), ca. 190 for Carduus baeocephalus subsp. microstigma Gaisberg & Wagenitz (facultative autogamy; Gaisberg 2002), 3650.01-6500.01 for Carthamus tinctorius L. (facultative xenogamy; Shao et al. 2012), and 107.17-138.92 and 926.58-2308.38, respectively, for cleistogamous and chasmogamous flowers of Centaurea melitensis L. (Porras and Muñoz Álvarez 2000). In this sense, the values for Echinops and Cardopatium are at the high end of the range for Cardueae. Taken together, rather than a rebalancing of male reproductive investment to favor pollen grain size over number, the pattern observed in Echinops species with Perennial pollen type more likely suggests an increase of the overall resources devoted to male function.

Functional Significance of the Evolution of the Echinops Perennial Pollen Type

Among the many factors that contribute to determining pollen size, postpollination processes, rather than pollen transport, appear to be prevalent in animal-pollinated species (Harder 1998; Sarkissian and Harder 2001). In particular, a significant relationship exists between pollen volume and style length (Torres 2000). Indeed, pollen tube length is predetermined by the nutrient reserves of the pollen and must adjust to the pistil length (Cruden 2009). In addition, pollen tube growth rate also depends on nutrient reserves, which gives the large pollen grains an advantage in the competition for fertilization. Pollen size increase in Echinops could therefore respond to an evolutionary trend toward producing larger flowers and syncephalia and/or a selection pressure for improved male fitness. Flower length varies considerably in Echinops, up to 4-4.5 cm in the gigantic Ethiopian species E. ellenbeckii O.Hoffm. and E. longisetus A.Rich. (Tadesse 2004), with enormous syncephalia up to 25 cm in diameter and pollinated by birds (Vogel 2015). Certainly, these Echinops species, for which unfortunately no palynological or phylogenetic data are available, should constitute a target for future studies on Echinops pollen evolution. Indeed, in Chresta (a syncephalous representative of the Vernonieae tribe), the largest pollen grains are found in C. curumbensis (Philipson) H.Rob. and C. speciosa Gardner, which also have the largest flowers (up to 3.3 cm) and are bird pollinated (Siniscalchi et al. 2017).

There is growing evidence that environmental factors also participate in determining pollen size and shape (Ejsmond et al. 2011, 2015). Pollen undergoes dehydration when released from the anther and rehydration once deposited on the stigma, processes that can alter pollen viability and whose intensity depends on the level of dryness of the environment. Accordingly, a trend toward larger pollen (i.e., with a lower surface-to-volume ratio and therefore a lower desiccation rate) is observed at high temperatures (Ejsmond et al. 2015). This trend, however, is interpreted as more likely resulting from increased competition between pollen on the stigma than representing an adaptation to high desiccation stress (Ejsmond et al. 2015). Larger pollen grains in *Echinops* may therefore reflect the adaptation to warmer environmental conditions.

In addition to the effect of water stress on pollen viability, dehydration and rehydration can also affect the structure of the exine and cause the pollen to break. The size and shape of pollen grains influence their harmomegathic functioning (i.e, the capacity to accommodate changes in volume through exine elasticity and harmomegathic structures), which is optimized in smaller and more elongated pollen (Wodehouse 1935). In this sense, the particular structure of *Echinops* Perennial pollen type, its exceptionally complex wall and elongated shape, could certainly have the function of allowing a considerable increase in size while maintaining sufficient harmomegathic capacity.

Although some taxa may share certain pollen characteristics with *Echinops*, such as an elongated shape in the case of Arctiinae (fig. 5A), the evolutionary trajectory of *Echinops* pollen remains quite exceptional. Actually, other strategies have been explored in Asteraceae lineages to cope with arid and hot climates, which do not involve the huge increase of pollen size and a dramatic shape change seen in *Echinops*. For example, in Centaureinae,

adaptation to such climates is achieved through the development of a harmomegathic structure, the cavea (Hidalgo et al. 2008a, 2008b and references therein). Similarly, in Vernonieae, type F pollen (sensu Keeley and Jones 1979) is described by a reduced amount of tectum, thick columellae around the lacunae and germinal pores, and a highly geometric form with patterned elevated ridges and lacunae regularly spaced. These characteristics probably confer improved tensile strength that may be related to volume changes and an increased resistance to desiccation (Bolick 1978; Keeley and Jones 1979). Type F pollen is found in very few Vernonieae species that grow in very harsh conditions (e.g., the syncephalous Chresta martii (DC.) H.Rob. over rock outcrops in the dry Brazilian caatinga and Linzia Sch.Bip. ex Walp. spp. in dry or recently burnt grasslands; Jeffrey and Beentje 2000; Siniscalchi et al. 2017) and in some syncephalous weedy species (e.g., Elephantopus mollis Kunth and Rolandra fruticosa (L.) Kuntze; Robinson 1992; Skvarla et al. 2005).

Concluding Remarks

In this work, we provide insights into the evolution of pollen gigantism in *Echinops* and its potential relationships with adaptation to warmer environmental conditions. As evidence accumulates across plant lineages of a response of pollen morphology to climate, we can anticipate that current global changes might also influence pollen phenotypic spectrum as well as pollination success. In this sense, the present study contributes to increasing the body of knowledge on pollen traits that is still too limited and yet crucial for modeling the impact of global change.

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