

Pollen Morphology of Cotoneaster (Rosaceae) Species in Türkiye and Its Systematic Significance

Serap İşik Seylan¹* D, Zübeyde Uğurlu Aydın² D and Ali A. Dönmez² D

¹Department of Mathematics and Science Education, Faculty of Education, Hacettepe University, 06800 Beytepe, Ankara, Türkiye

²Department of Biology, Molecular Plant Systematic Laboratory (MOBIS), Faculty of Science, Hacettepe University, 06800 Beytepe, Ankara, Türkiye

Received December 12, 2023; revision accepted September 6, 2024

Pollen morphology of 14 specimens from all nine Turkish native species along with one undescribed species of *Cotoneaster* was examined by light microscopy (LM) and scanning electron microscopy (SEM). The most frequent pollen types are trizonocolporate and tetracolporate, respectively and they occur together. Moreover, trisyncolporate, 3-parasyncolporate, 4-parasyncolporate, and pentacolporate pollen grains were observed in *Cotoneaster* for the first time in this study. Trizonocolporate pollen grains are radially symmetrical, isopolar, 22.5–35 × 26.25–42.5 µm in size and suboblate in shape. The amb is triangular, except *C. integerrimus*, which has a circular outline. Two endoaperture types can be distinguished. One is rectangular/square, the other is circular (endopore) and present only in *C. integerrimus*, *C. melanocarpus* and *C. tomentosus*. The exine pattern can be divided into two types, namely striate-perforate and rugulate-perforate. The quantitative pollen characteristics were analyzed using principal component analysis (PCA) and all pollen features were considered for hierarchical cluster analysis (HCA). Principal component analysis revealed that colpus width, endoaperture length, and endoaperture width are the most powerful metrical pollen traits in *Cotoneaster* identification. According to hierarchical cluster analysis, pollen traits enable division of *Cotoneaster* into two subgenera. The present study provides insights into pollen morphology of *Cotoneaster*, as well as its use for taxonomic purposes.

Keywords: Cotoneaster, hierarchical cluster analysis, pollen morphology, principal component analysis, taxonomy

INTRODUCTION

Rosaceae, comprising ca. 90 genera and 3000 species, is a relatively large family, notable for its taxonomic difficulty due to polyploidization, hybridization, apomixis, and radial evolution (The Angiosperm Phylogeny Group, 2016). As a consequence of complex breeding system, the genus *Cotoneaster* Medik. is one of the complicated taxa in Rosaceae and it includes 50 to 370 species according to different taxonomic approaches (Flinck and Hylmö, 1966; Phipps et al., 1990; Fryer and Hylmö, 2009; Dickoré and

Kasperek, 2010). The genus is a woody member of Malinae subtribe (referred to Maloideae in Morgan et al., 1994; Evans and Campbell, 2002 and Pyrinae in Campbell et al., 2007; Potter et al., 2007; Li et al., 2012; Lo and Donoghue, 2012) and chiefly distributed in the northern hemisphere (Fryer and Hylmö, 2009). Some species of *Cotoneaster* are cultivated for ornamental usage because of the flower, colored fruit and cold tolerance (Li et al., 2012).

The complex breeding system allows to increase morphologically intermediate offspring. For instance, apomictic breeding is a well

^{*} Corresponding author, serapi@hacettepe.edu.tr

known phenomenon in Cotoneaster as in other genera of Rosaceae (Li et al., 2012; Rothleutner et al., 2016). Likewise, hybridization is a frequent occurrence in cultivation and the genus is composed of more than 50% tetraploids species (Fryer and Hylmö, 2009; Dickinson, 2018), although more comprehensive studies are needed to prove natural hybridization in Cotoneaster (Dickoré and Kasperek, 2010). Relationships within Cotoneaster are not deeply resolved, owing to complex species groups and lack of diagnostic morphological characters (Li et al., 2012; Meng et al., 2021). Initially, some flower characters such as petal color (white vs. pink or red), flower number and spreading or erect petals were used to divide Cotoneaster into two subgenera/sections, Chaenopetalum and Orthopetalum (syn. Cotoneaster) (Koehne, 1893; Flinck and Hylmö, 1966; Phipps et al., 1990; Fryer and Hylmö, 2009). Recently, molecular phylogenetic studies (Li et al., 2012; Meng et al., 2021) have focused on infrageneric taxonomy, hybridization and character evolution in the genus and showed the incongruence between the nrITS and cpDNA data indicating hybridization.

Rosaceae pollen morphology regarding subfamilies, tribes and genera has been described by many researchers (Zhou et al., 1999; 2000; Song et al., 2016) and pollen morphology of Cotoneaster has been studied by several authors (Eide, 1981; Hebda and Chinnappa, 1990; Sáez and Rosselló, 2012; Perveen and Qaiser, 2014; Ghosh and Saha, 2017). Some researchers (Hsieh and Huang, 1997; Chang et al., 2011a; Chang et al., 2011b) also examined pollen morphology of four Taiwanese Cotoneaster species, while four species from China were studied by Jarvis et al. (1992) and Zhou et al. (2000). Pollen morphology of several Cotoneaster species that occur in Iran (16 species) was examined by Raei Niaki et al. (2020), five of which (C. integerrimus Medik., C. melanocarpus (Bunge) Fischer, C. nummularius Fisch. & C.A.Mey., C. morulus Pojark. and C. multiflorus Bunge.) also grow in Türkiye and were investigated in our study. It is known that pollen morphology contains unique characters with a large amount of genetic information and they can play important roles for species identification and phylogenetic studies of fruit trees (Song et al., 2017). Some researches focused on taxonomic importance of pollen characters and showed valuable palynological knowledge for taxonomy of the target taxa (Işık

and Dönmez, 2004; Oybak Dönmez, 2008; Ullah et al. 2022). However, pollen morphology of Turkish *Cotoneaster* taxa has been poorly understood from a taxonomic perspective.

The main object of the study was to investigate pollen morphology of all Turkish native *Cotoneaster* species and to evaluate pollen features in respect of taxonomy of the genus. Pollen features were examined by light and scanning electron microscopy, and multivariate analyses were performed using nine quantitative and four qualitative pollen characters. More specifically, palynological knowledge was employed to understand the taxonomic implications of pollen morphology in *Cotoneaster*.

MATERIALS AND METHODS

PLANT MATERIAL

The pollen material used in the present study was collected in Türkiye by the second and third authors (Table 1), and the voucher specimens are deposited at the Herbarium of Hacettepe University (HUB). For pollen morphological studies, a total of 13 specimens from eight known species, and one newly described species in preparation for publication were investigated. Whenever possible, more than one specimen was investigated for each species. The following specimens were examined: one from *C. integerrimus*, *C. melanocarpus*, *C. meyeri*, *C. transcaucasicus*, *C. morulus* and *Cotoneaster sp.*, two from *C. multiflorus* and *C. tomentosus*, and four from *C. nummularius*.

LIGHT AND SCANNING ELECTRON MICROSCOPIC STUDIES

For light microscopic (LM) studies, pollen placed on a microscope slide was first treated with 2–3 drops of 70% ethyl alcohol to remove oily substances (like pollenkitt). After the ethyl alcohol on the slide had evoparated on a hot plate, the pollen material was embedded in glycerine-jelly, stained with basic fuchsine, following the method of Wodehouse (1935). The aperture form was determined for a minimum of 100 grains per specimen, and the frequency occurrence of each aperture type was then calculated. Size measurements were made on the most frequent trizonocolporate pollen grains. The following param-

TABLE 1. Voucher information of the studied *Cotoneaster* specimens in this study.

Specimens	Voucher information
C. integerrimus Medik.	Sivas: 1662 m, 40°10'11.0"N, 037°52'715"E, 11.6.2014, AAD 19111
C. melanocarpus (Bunge) Fisch.	Kars: Kağızman, 1331 m, 40°03'54.3"N, 042°51'163"D, 05.05.2013, AAD 18486
C. multiflorus Bunge.	Erzurum: 2172 m, 40°40'00.7"N, 040°32'824"E, 29.06.2012, AAD 18020-M. Beilstein & J. Brock
	Sivas: Divriği, 1691 m, 39°36'991"N, 037°43'113"E, 11.6.2014, AAD 19101
C. meyeri Pojark.	Erzurum: Oltu, 1660 m, 40°23'99.9"N, 041°32'906"E, 30.05.2012, AAD 17960 - Z. Uğurlu
C. nummularius Fisch. & C.A. Mey.	Sivas: from Refahiye to Sivas, 25.05.2012, AAD 17912 - Z. Uğurlu
	Bingöl: 1145 m, 38°56'431"N, 040°38'105"E, 27.05.2012, AAD 17934 - Z. Uğurlu
	Kars: 1195 m, 40°10'815"N, 043°08'557"E, 30.05.2012 AAD 17947- Z. Uğurlu
	Çankırı: Kurşunlu, 1431 m, 40°55'78.7"N, 033°14'743"E, 30.05.2013, AAD 18530
C. transcaucasicus Pojark.	Kars: 1195 m, 40°10'815"N, 043°08'557"E, 30.05.2012, AAD 17944 - Z. Uğurlu
C. morulus Pojark.	Artvin: Yaylabaşı, 1468 m, 40°52'95.0"N, 041°20'795"E, 03.06.2013, AAD 18565
C. tomentosus (Aiton) Lindl.	Çankırı: Kurşunlu, 1431 m, 40°55'78.7"N, 033°14'743"E, 30.5.2013, AAD 18528.
	Çankırı: 30.05.2014, AAD 19045
Cotoneaster sp.	A4 Çankırı: Kurşunlu, 13.05.2014, AAD 19036

eters were measured: pollen size, given by the polar axis (P) and equatorial axis (E); equatorial diameter in polar view; distance between two ectocolpi at poles; colpus length (Clg) and colpus width (Clt); endoaperture length (Enaplg) and endoaperture width (Enaplt); exine and intine thickness. The pollen shape was determined by the ratio of mean length of polar axis to the mean length of equatorial axis (P/E). The polar area index (PAI) was calculated as the ratio of the distance between the apices of two ectocolpi to its equatorial diameter in polar view. P and E were measured for 50 pollen grains per specimen, and the other measurements were made on ten grains. IBM SPSS Statistics version 23 programme (George and Mallery, 2016) was utilized to calculate the means (M), standard deviations (SD) and ranges (V) for pollen size (P and E). Photomicrographs were taken with a Leica DFC 320 digital camera connected to a Leica DM 4000 B microscope.

For scanning electron microscopic (SEM) study, the pollen was first treated with 70% ethyl alcohol, then air-dried before being mounted on stubs, subsequently coated with gold. The photomicrographs were taken using a JSM 6490 LV electron microscope.

The palynological terminology mainly follows Punt et al. (2007) and Hesse et al. (2009). In addition, Van Bergen et al. (1995) and Grímsson et al. (2017) were used as sources for de-

scribing 3-parasyncolporate and 4-parasyncolporate pollen types.

MULTIVARIATE DATA ANALYSES

Thirteen pollen features, nine quantitative (polar axis, equatorial axis, polar area index, colpus length, colpus width, endoaperture length, endoaperture width, exine thickness, intine thickness) and four qualitative (pollen shape, amb shape, endoaperture shape, exine pattern) were analyzed for fourteen specimens of *Cotoneaster*.

For principal component analysis (PCA), arithmetic means of quantitative variables were used and the results were shown in a two-dimensional plot of the first and second PCs with eigenvalues of the characters. Vector values are provided in the results. All computations were run with the statistical software R version 4.0.5 using the FactoMineR and Factoextra packages (R Core Team, 2020). Hierarchical cluster analvsis (HCA) based the Gower coefficient (Gower, 1971) was performed to group the examined taxa based on all the obtained quantitative and qualitative pollen features presented in Tables 2-3. The multivariate data analysis was conducted using Past version 2.17c (Hammer et al., 2001).

TABLE 2. Quantitative pollen characteristics of the studied Turkish Cotoneaster taxa.

			Pollen 🤇	Pollen Type (%)				Ъ	P (µm)		a	E (µm)
Taxa-Voucher	trizc	tetrac	trisync	3-psc	4-psc	pentac	M	SD	۸	M	SD	Δ
C. integerrimus-AAD 19111	59	22	15	3	ı	1	28.85	±0.59	26.25-32.5	34.1	±0.83	28.75-37.5
C. melanocarpus-AAD 18486	65	20	12	2	П	l	27	±0.81	22.5-32.5	31.2	±0.94	26.25-37.5
C. multiflorus-AAD 18020 – M. Beilstein & J. Brock	66	1	1	I		ı	26.33	±0.67	22.5-30	31.88	±0.95	26.25–35
C. multiflorus-AAD 19101	26	40	13	17	4	I	26.25	±0.57	22.5 - 26.25	32.05	±0.81	26.25-37.5
C. meyeri-AAD 17960 – Z. Uğurlu	26	32	2	10	ı	ı	26.58	±0.61	24.3-29.16	31.03	±0.98	26.73-34.02
C. nummularius-AAD 17912 - Z. Uğurlu	94	9	I	I	ı	I	30.08	±0.7	26.73-34.02	36.94	±1.11	29.16-41.31
C. nummularius-AAD 17934 – Z. Uğurlu	49	47	2	2	ı	I	26	9.0∓	24.3-29.16	30.57	Ŧ	26.73-36.45
C. nummularius-AAD 17947 – Z. Uğurlu	66	1	ı	ı	ı	ı	27.41	€0.68	24.3-30.38	32.08	₹0.88	26.73-36.45
C. nummularius-AAD 18530	82	15	ı	1	ı	1	27.25	±0.54	25–30	32.33	∓0.85	27.5–37.5
C. transcaucasicus-AAD 17944 – Z. Uğurlu	40	29	ı	1	ı	ı	27.27	9.0∓	24.3-29.16	31.54	±0.82	26.73-36.45
C. morulus-AAD 18565	88	11	1	I	ı	ı	26.48	±0.65	22.5-31.25	32.4	±0.94	26.25–38.75
C. tomentosus-AAD 18528	65	25	10	1	ı	1	27.55	±0.47	25–30	31.68	±0.85	27.5–37.5
C. tomentosus-AAD 19045	46	53	1	1	ı	1	31.33	±0.75	27.5–35	36	±0.84	32.5-41.45
C. spAAD 19036	61	22	3	14	1	ı	27.25	99.0∓	22.5-31.25	33.65	±1.24	27.5-42.5

TABLE 2. Continued

Taxa-Voucher	PAI	Clg (µm)	Clt (µm)	Enaplg (µm) Enaplt (µm)	Enaplt (µm)	Exine thickness (µm)	Intine thickness (µm)
C. integerrimus-AAD 19111	0.18	19 (21) 23	8 (10.7) 16	7 (10.8) 13	8 (10.6) 14	1.75 (1.92) 2	0.75 (0.77) 1
C. melanocarpus-AAD 18486	0.23	18 (19.60) 22	8 (8.8) 10	6 (8.6) 11	7 (8.4) 10	1.5 (1.78) 2	0.75 (0.78) 1
C. multiflorus-AAD 18020 – M. Beilstein & J. Brock	0.21	18 (20.7) 23	9 (11.7) 14	12 (15.5) 18	11 (12) 14	1.5 (1.7) 1.75	0.75 (0.8) 1
C. multiflorus-AAD 19101	0.21	18 (19.80) 21	9 (12.7) 14	12 (15.2) 17	10 (12.6) 14	1.5 (1.67) 1.75	0.75 (0.8) 1
C. meyeri-AAD 17960 – Z. Uğurlu	0.20	19 (21.80) 24	12.5 (13.75) 17.5	11 (14.8) 19	11 (13.4) 17	1.5(1.8)2	0.75 (0.8) 1
C. nummularius-AAD 17912 – Z. Uğurlu	0.21	23 (24.20) 25	15 (18) 20	19 (20.4) 23	15 (16.3) 20	1.5 (1.83) 2	0.75 (0.78) 1
C. nummularius-AAD 17934 – Z. Uğurlu	0.22	19 (21.20) 23	12.5 (15.5) 20	12 (15.8) 18	12 (13.3) 16	1.5 (1.78) 2	0.75 (0.8) 1
C. nummularius-AAD 17947 – Z. Uğurlu	0.23	21 (23.4) 25	12.5 (15.25) 17.5	14 (15.5) 18	12 (14.2) 16	1.5(1.8)2	0.75 (0.8) 1
C. nummularius-AAD 18530	0.25	19 (21.20) 24	9 (11.6) 13	10 (12.1) 15	9 (11.5) 13	1.75 (1.8) 2	0.75 (0.8) 1
C. transcaucasicus-AAD 17944 – Z. Uğurlu	0.22	20 (21.38) 25	9 (11.6) 13	10 (12.1) 15	9 (11.5) 13	1.75 (1.8) 2	0.75 (0.8) 1
C. morulus-AAD 18565	0.26	20 (22.20) 25	10 (11.8) 15	11 (13.7) 18	10 (12.2) 16	1.5 (1.8) 2	0.75 (0.78) 1
C. tomentosus-AAD 18528	0.29	21 (23.3) 26	6 (7.6) 10	6 (7.8) 10	6 (7.3) 9	1.5 (1.83) 2	0.75 (0.78) 1
C. tomentosus-AAD 19045	0.34	22 (23.3) 26	6 (8.30) 10	5 (8.2) 10	5 (8.05) 10	1.5 (2) 2.5	0.75 (0.8) 1
C. spAAD 19036	0.29	21 (21.90) 24	9 (12.7) 17	9 (12.6) 15	8 (11.4) 15	1 (1.55) 2	0.50 (0.63) 0.75

porate, trisync - trisyncolporate, psc - parasyncolporate, pentac - pentacolporate, PAI - polar area index, Clg - colpus length, Clt - colpus width, Abbreviations: P - polar axis, E - equatorial axis, M - mean value, SD - standard deviation, V - variation, trizc - trizonocolporate, tetrac - tetracol-Enaplg - endoaperture length, Enaplt - endoaperture width.

TABLE 3. Qualitative pollen characteristics of the studied Turkish Cotoneaster taxa.

Taxa-Voucher	Pollen shape	Amb	Endoaperture shape	Exine pattern
C. integerrimus-AAD 19111	suboblate	circular	circular (endopore)	Striate-perforate
C. melanocarpus-AAD 18486	suboblate	triangular	circular (endopore)	Rugulate-perforate
C. multiflorus-AAD 18020 – M. Beilstein & J. Brock	suboblate	triangular	rectangular/square	Striate-perforate
C. multiflorus-AAD 19101	suboblate	triangular	rectangular/square	Striate-perforate
C. meyeri-AAD 17960 – Z. Uğurlu	suboblate	triangular	rectangular/square	Striate-perforate
C. nummularius-AAD 17912 – Z. Uğurlu	suboblate	triangular	rectangular/square	Striate-perforate
C. nummularius-AAD 17934 – Z. Uğurlu	suboblate	triangular	rectangular/square	Striate-perforate
C. nummularius-AAD 17947 – Z. Uğurlu	suboblate	triangular	rectangular/square	Striate-perforate
C. nummularius-AAD 18530	suboblate	triangular	rectangular/square	Striate-perforate
C. transcaucasicus-AAD 17944 – Z. Uğurlu	suboblate	triangular	rectangular/square	Striate-perforate
C. morulus-AAD 18565	suboblate	triangular	rectangular/square	Rugulate-perforate
C. tomentosus-AAD 18528	suboblate	triangular	circular (endopore)	Rugulate-perforate
C. tomentosus-AAD 19045	suboblate	triangular	circular (endopore)	Rugulate-perforate
C. spAAD 19036	suboblate	triangular	rectangular/square	Rugulate-perforate

RESULTS

GENERAL POLLEN MORPHOLOGY

A summary of pollen morphological observations of the Turkish Cotoneaster species examined in this study under LM and SEM is given Tables 2-3 and Figs. 1-2. Pollen grains occur as monads. The most frequent pollen types are trizonocolporate (26-99%) (Fig. 1a, b and Fig. 2a, b) and tetracolporate (1-59%) (Fig. 1c and Fig. 2c). However, in C. multiflorus (AAD 19101), C. transcaucasicus Pojark. and C. tomentosus (Aiton) Lindl. (AAD 19045) samples, the ratio of tetracolporate pollen type is higher than the ratio of trizonocolporate pollen. In all specimens both trizonocolporate and tetracolporate pollen grains occur together. In addition, some Cotoneaster specimens also have trisyncolporate (1-15%; C. integerrimus, C. melanocarpus, C. multiflorus-AAD 19101, C. meyeri Pojark., C. nummularius-AAD 17934, C. morulus, C. tomentosus, C. sp.) (Fig. 1d), 3-parasyncolporate (1-17%; C. integerrimus, C. melanocarpus, C. multiflorus-AAD 19101, C. meyeri, C. nummularius-AAD 17934, C. transcaucasicus, C. sp.) (Fig. 1e), 4-parasyncolporate (1-4%; C. melanocarpus, C. multiflorus-AAD 19101) (Fig. 1f) and pentacolporate (1%; C. integerrimus) (Fig. 1g) pollen grains. Trizonocolporate pollen grains are radially symmetrical, isopolar

and small to medium in size: the polar axis (P) measures 22.5–35 µm, and the equatorial axis (E) measures 26.25–42.5 μm. The pollen grains are suboblate in shape. The equatorial outline is transversely elliptic. The amb (the outline of a pollen grain or spore seen in polar view) is triangular (Fig. 1a), with the exception of C. integerrimus, which has a circular amb (Fig. 1b). The polar area index (PAI) ranges from 0.18 to 0.34. The apertures situated at the equator and the aperture structure consist of a colpus and endoaperture. Margins of colpi are straight. Colpus length (Clg) ranges from 18 to 26 µm and colpus width (Clt) ranges from 6 to 20 um. Colpus ends are acute (Fig. 1a) and colpus membrane is without any sculpturing element. Endoapertures are distinct and two endoaperture types can be distinguished. One is rectangular/square in outline (Fig. 1h), the other is circular (endopore) (Fig. 1i), and occurs only in C. integerrimus, C. melanocarpus and C. tomentosus. Rectangular endoapertures are slightly equatorially elongate (lalongate) in some pollen grains of C. multiflorus and C. morulus. Endoaperture length (Enaplg) ranges from 5 to 23 µm and endoaperture width (Enaplt) ranges from 5 to 20 µm. The exine wall is tectate-columellate and 1–2.5 µm in thickness. Exine pattern types are striate-perforate (Fig. 2d-h) and rugulate-perforate (Fig. 2i-l). The striate-perforate type consists of lirae (which vary in size, orientation and anastomosing) separated by grooves (varied in size), having perforations (which vary in size and frequency) and the rugulate-perforate type consists of elongated exine elements longer than 1 μ m, irregularly arranged (with perforations). Intine thickness is $\leq 1 \mu$ m.

EVALUTION OF MULTIVARIATE ANALYSES

The metric data of pollen grains were presented using PCA to reveal taxonomic relationship among the taxa (Fig. 3, Table 4). The first and

second axes from the PCA explained 80.9% of the accumulated variance of the analyzed data. The first axis (PC1) accounted for 36.8% of the variance, mainly based on pollen size (P and E). Cumulative variance of the second axis (PC2) totaled 44.1% and it was associated with the endoaperture size (Enaplg and Enaplt) and colpus width. Most of the target taxa were nested on the negative side of axis 1 and had the highest size of colpus width and endoaperture. The remaining taxa, i.e., *C. melanocarpus*, *C. tomentosus*, *C. integerrimus* and *Cotoneaster sp.*,

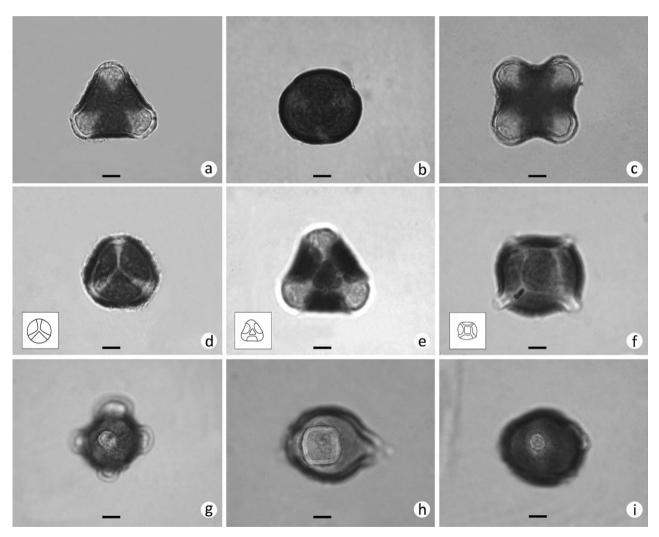


Fig. 1. Light microscopy (LM) photomicrographs of pollen grains of selected *Cotoneaster* species. **(a-b)** Trizonocolporate pollen grain in polar view **(a)** triangular in *C. morulus* (AAD 18565), **(b)** circular in *C. integerrimus* (AAD 19111). **(c)** Tetracolporate pollen grain in polar view in *C. multiflorus* (AAD 19101). **(d)** Trisyncolporate pollen grain in polar view in *C. multiflorus* (AAD 19101). **(f)** 4-parasyncolporate pollen grain in polar view in *C. multiflorus* (AAD 19101). **(g)** Pentacolporate pollen grain in polar view in *C. integerrimus* (AAD 19111). **(h)** Rectangular/square endoaperture in equatorial view in *C. meyeri* (AAD 17960 – Z. Uğurlu). **(i)** Circular endoaperture (endopore) in equatorial view in *C. melanocarpus* (AAD 18486). Scale: 10 μm.

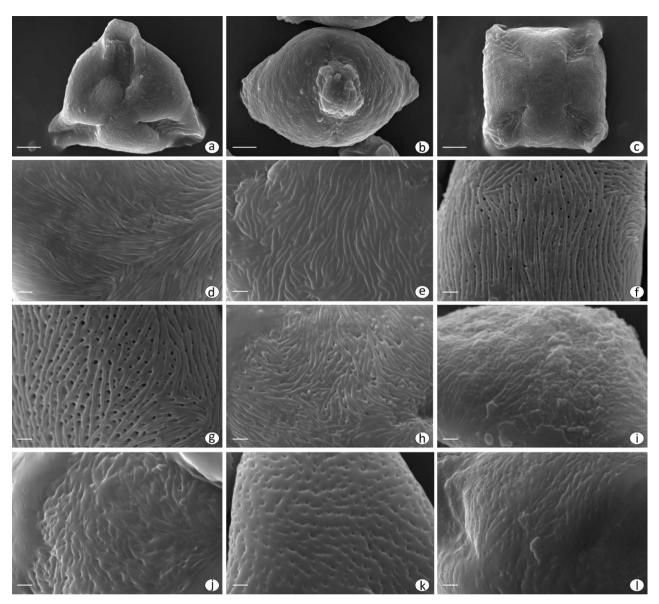


Fig. 2. Scanning electron microscopy (SEM) photomicrographs of pollen grains of *Cotoneaster* species. **(a–c)** *C. transcaucasicus* (AAD 17944 – Z. Uğurlu) **(a–b)** Tricolporate pollen grain **(a)** in polar view, **(b)** in equatorial view. **(c)** Tetracolporate pollen grain in polar view. **(d–h)** Striate-perforate exine patterns **(d)** *C. integerrimus* (AAD 19111), **(e)** *C. multiflorus* (AAD 19101), **(f)** *C. meyeri* (AAD 17960 – Z. Uğurlu), **(g)** *C. transcaucasicus* (AAD 17944 – Z. Uğurlu), **(h)** *C. nummularius* (AAD 17934 – Z. Uğurlu). **(i-1)** Rugulate-perforate exine patterns **(i)** *C. melanocarpus* (AAD 18486), **(j)** *C. morulus* (AAD 18565), **(k)** *C. tomentosus* (AAD 19045), **(l)** *C. sp.* (AAD 19036). Scale: a–c = 10 μm; d-l = 2 μm.

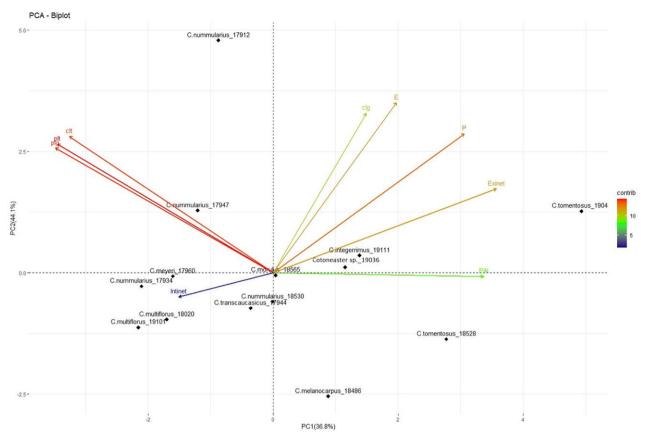


Fig. 3. Principal Component Analysis (PCA) performed with the pollen metric data of *Cotoneaster* taxa. Character codes follow Table 4. (contrib: contribution of variables to principal component).

TABLE 4. Vector values of the principal component analysis (PCA) of the nine quantitative pollen morphological characters of *Cotoneaster* taxa. The highest values are marked in bold.

Name of variables	PC1 (36.8%)	PC2 (44.1%)
Polar axis (P)	-0.042	0.592
Equatorial axis (E)	0.053	0.702
The polar area index (PAI)	-0.004	0.006
Colpus length (clg)	0.058	0.389
Colpus width (clt)	0.553	0.011
Endoaperture length (plg)	0.677	-0.039
Endoaperture width (plt)	0.476	-0.030
Exine thickness (Exinet)	-0.006	0.024
Intine thickness (Intinet)	0.040	-0.001

were positioned on the positive side of this axis. The common feature of these species was the highest value of polar and equatorial axes. Most of the *Cotoneaster* taxa were grouped around the centre of PC1 and PC2, while different specimens of *C. nummularius* and *C. to-*

mentosus were distributed at the extreme sides of PC1 and PC2.

Based on the qualitative and quantitative pollen characters, hierarchical cluster analysis (HCA) was performed to explore the relationship of *Cotoneaster* taxa. The phenogram produced three main sharp groups (Fig. 4). Group I includes *C. multiflorus*, *C. meyeri*, *C. nummularius* (AAD 17934, AAD 17947, AAD 17912) and *C. transcaucasicus*. Four species of *Cotoneaster*. *C. integerrimus*, *C. melanocarpus*, *C. nummularius* (AAD 18530) and *C. tomentosus* appear in Group II and the remaining two taxa (*C. morulus* and *C. sp.*) are placed in Group III.

Among the examined taxa, *C. morulus* and *C. sp.* show a rugulate-perforate exine pattern with a rectangular/square endoaperture shape. Likewise, *C. tomentosus* and *C. melanocarpus* have the same exine pattern (rugulate-perforate) but the endoaperture shape is circular (endopore). Only *C. integerrimus* is characterized by a striate-perforate exine pattern and a circular (endopore) endoaperture shape.

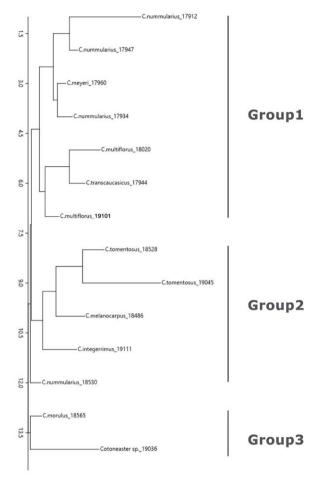


Fig. 4. Hierarchical Cluster Analysis performed with all pollen variables of *Cotoneaster* taxa.

The largest group of the phenogram including all populations of *C. nummularius*, *C. multiflorus*, *C. meyeri* and *C. transcaucasicus* is characterized by a striate-perforate exine pattern and rectangular/square endoaperture shape. It is clear that *Cotoneaster* taxa are classified based on the exine pattern features and endoaperture shape.

DISCUSSION

General outlines of the pollen grains in the Maleae tribe, which also comprises *Cotoneaster*, vary from oblate through spherical to prolate in equatorial view and the amb can be triangular through circular to three-lobed (Reitsma, 1966; Hebda et al., 1988; Perveen and Qaiser, 2014; Zhou et al., 2000; Ghosh and Saha, 2017). The main aperture type is tricolporate (Reitsma, 1966; Eide, 1981). However, some members (i.e.,

Mespilus, Amelanchier, Crataegus, Sorbus, Malus, ×Malosorbus, Erilobus) also have non-tricolporate pollen types such as tricolpate, tetracolporate, syncolpate, syncolporate, pericolp(or)ate and inaperturate (Reitsma, 1966; Hebda et al., 1988; Christensen, 1992; Işık and Dönmez, 2004; Oybak Dönmez, 2008). Pollen size in the tribe varies from 10.7 µm (Cotoneaster) to 56 µm (Crateagus, longest axis) (Christensen, 1992; Ghosh and Saha, 2017). Two basic types of sculpturing are observed: rugulate-striate and non-rugulate-striate. Rugulate to striate pollen grains are most frequent (Hebda and Chinnappa, 1990;1994; Zamani et al., 2010). Non-rugulate-striate pollen includes granulate-microscabrata (Pyrus) (Ghosh and Saha, 2017), perforate and microreticulate (Crataegus) (Oybak Dönmez, 2008), and reticulate (Mespilus, Pyrus, Pyracantha) (Byatt et al., 1977; Jarvis et al., 1992; Zhou et al., 2000).

Pollen grains of all the examined Turkish Cotoneaster species are suboblate in equatorial view (Table 3), unlike the spherical to prolate pollen grains recorded in previous Cotoneaster studies (Eide, 1981; Hsieh and Huang, 1997; Sáez and Rosselló, 2012; Perveen and Qaiser, 2014; Raei Niaki et al., 2020). The trizonocolporate pollen type is predominant in the Rosaceae family (Reitsma, 1966; Hebda and Chinnappa, 1990; Ghosh and Saha, 2017) as in Cotoneaster taxa. It was revealed that Turkish Cotoneaster species possess other pollen types along with trizonocolporate pollen within the same sack, including tetracolporate, trisyncolporate, 3-parasyncolporate, 4-parasyncolporate pentacolporate ones. In all the specimens trizonocolporate and tetracolporate pollen grains co-occur but the others are present only in some specimens (Table 2). Tetracolporate pollen grains in three Cotoneaster species from Iran were also reported by Raei Niaki et al. (2020). However, in several previous pollen morphological studies on Cotoneaster, only trizonocolporate pollen was recorded (Eide, 1981; Hebda and Chinnappa, 1990; 1994; Jarvis et al., 1992; Hsieh and Huang, 1997; Zhou et al., 2000; Chang et al., 2011a; 2011b; Perveen and Qaiser, 2014; Ghosh and Saha, 2017). Trisyncolporate, 3-parasyncolporate, 4-parasyncolporate, and pentacolporate pollen grains were observed in Cotoneaster for the first time in this study (Fig. 1d-g). Polyploidy and apomixis are important phenomena among plants (Richards, 1990) and they are common in Cotoneaster species (Sax, 1954; Campbell and

Dickinson, 1990; Campbell et al., 1991; Bartish et al., 2001; Gregor, 2013; Rothleutner et al., 2016; Dickinson, 2018). Variation in pollen aperture numbers is generally related to different levels of ploidy (Borsch and Wilde, 2000) and the occurrence of pollen polymorphism associated with polyploid taxa has been recorded in some species (Chinnappa and Warner, 1982). In the case of the Turkish *Cotoneaster* samples, polyploidy and apomixis have not been examined yet. However, pollen type diversity observed in Turkish *Cotoneaster* species may be associated with polyploidy and apomixis.

The pollen grains of Turkish *Cotoneaster* are small to medium and 22.5–35µm (P), 26.25–42.5 µm (E) in size (Table 2). The range of (P) values is almost consistent with those of *Cotoneaster* species (23.2–34 µm) from Pakistan (Perveen and Qaiser, 2014) and 19.6–37.6 µm (Eide, 1981) from north-west Europe. However, Ghosh and Saha (2017) reported the smallest *Cotoneaster* pollen size (P: 10.4–11.2 µm, E: 7.2–8.8 µm) from India. There is no significant variation in pollen size between the Turkish species and within them, but only, *C. meyeri, C. transcaucasicus* and *C. tomentosus* show a narrower range of polar axis and equatorial axis.

The aperture form is also an important feature of Rosaceae pollen. In the family, the aperture structure usually consists of a colpus and equatorial endoaperture, which varies from being well-defined to weakly defined (Hebda and Chinnappa, 1990). Although the endoaperture shape is rectangular/square in most Turkish Cotoneaster specimens, C. integerrimus, C. tomentosus and C. melanocarpus can be distinguished from other species by their circular endoaperture shape (endopore) (Table 3). In previous studies on Cotoneaster, there was no detailed information about the endoaperture shape feature. Only Jarvis et al. (1992) divided rosaceous pores into three main types and established that C. horizontalis, which grows in China, has a star-shaped pore region. Furthermore, Hebda and Chinnappa (1990) mentioned the presence of a pore flap in *Cotoneaster* pollen. However, a pore flap is known to be characteristic of Amelanchier and Crataegus in the Maleae tribe (Hebda et al., 1988).

All the Turkish *Cotoneaster* specimens have a triangular amb, as in the Iranian samples described by Raei Niaki et al. (2020), except *C. integerrimus*, which has a circular outline (Table 3).

It was also noted that the pollen grains of some *Cotoneaster* species in China and Taiwan have a circular amb (Hsieh and Huang, 1997; Zhou et al., 2000; Chang et al., 2011b).

Polar area index (PAI) and aperture size (colpus length and width, endoaperture length and width) vary slightly between species (Table 2). Polar area index is often an informative differentiating character for distinguishing between types. It depends highly on the length of the ectocolpi. Long ectocolpi result in a small apocolpium index, short ectocolpi give large indices (Punt and Hoen, 2009). On the other hand, such a relationship was not observed in the studied Turkish Cotoneaster species. However, colpus width (Clt), endoaperture length (Enaplg) and width (Enaplt) values were observed to be shorter in C. integerrimus, C. tomentosus and C. melanocarpus species with a circular endoaperture (endopore), and the PCA shows that these features are important to group the analyzed species (Tables 2-4). The exine pattern is considered to be one of the most important distinguishing characters in Rosaceae (Eide, 1981; Hebda and Chinnappa, 1990; 1994; Ghosh and Saha, 2017). In the Turkish Cotoneaster species, the most frequent exine patterns are rugulate-perforate and striate-perforate (Table 3 and Fig. 2d-l). The examined specimens can be divided into two groups, based on the exine pattern. Also, in previous studies on Cotoneaster, striate perforate (Hebda and Chinnappa, 1994; Hsieh and Huang, 1997; Zhou et al., 2000; Raei Niaki et al., 2020) and rugulate perforate (Eide, 1981; Hsieh and Huang, 1997; Chang et al., 2011a) exine patterns were reported. However, psilate (Eide, 1981; Hebda and Chinnapa, 1990; Raei Niaki et al., 2020) and foveolate (Zhou et al., 2000) exine patterns are also seen in Cotoneaster species.

Two of multivariate analysis, namely PCA and HCA, were performed in this study. Most of the studied taxa were placed around the negative side of PC1, except *C. melanocarpus*, *C. tomentosus*, *C. integerrimus* and *Cotoneaster sp.* The PCA biplot in this study shows that pollen, colpus width and endoaperture size offer important information for *Cotoneaster* identification (Fig. 3). Similarly, Li et al. (2012) investigated palynological traits in *Prunus* taxa to assess taxonomic relationships, and found colpus width with the other pollen characteristics (equatorial size, colpus length and ridge width) to be the

most powerful diagnostic characters for apricot identification.

The results of PCA based on metric variables are not strictly correlated with clustering analysis loadings for all pollen characters. All specimens are classified into two groups on the plot of PCA, which included only quantitative characters, while three sharp groups are nested on the phenogram, which is composed of both quantitative and qualitative characters (Fig. 4). Four species, C. tomentosus, C. melanocarpus, C. integerrimus and C. morulus, belonging to subgenus Cotoneaster, were grouped in the same group with C. nummularius (AAD 18530) and the unidentified Cotoneaster specimen. It is in accordance with previous studies (Li et al., 2012; Raei Niaki et al., 2020) and supports monophyly of subgenus Cotoneaster. Raei Niaki et al. (2020) consider the correlation between a particular pollinator group and pollen type to explain pollen similarity in subgenus Cotoneaster. Interestingly, one population of C. nummularius (AAD 18530) belonging to subgenus Chaenopetalum is placed in the same group with the taxa of subgenus Cotoneaster. This may be because it has smaller colpus width (Clt), endoaperture length (Enaplg) and endoaperture width (Enaplt) values than other C. nummularius populations. The remaining studied taxa belonging to subgenus Chaenopetalum are clustered into the second branch of the phenogram. From a taxonomic point of view, our results confirm that pollen features are important to divide Cotoneaster taxa into two subgenera.

CONCLUSION

The present study highlights pollen features of all nine Turkish native taxa such as pollen type, pollen size, pollen shape, polar area index, aperture size and characteristics, exine-intine thickness, exine pattern characteristics as well as taxonomic usefulness of pollen characters. Quantitative and qualitative pollen characteristics are approximately consistent between populations of the same taxa. Four new pollen types (trisyncolporate, 3-parasyncolporate, 4-parasyncolporate, and pentacolporate) for *Cotoneaster* were observed for the first time in this study, unlike in previous studies on the pollen morphology of the genus. PCA shows that pol-

len data are limited to resolve taxonomic relationships at the species level. The HCA results are more useful to split *Cotoneaster* taxa into two main subgenera based on the studied palynological features. Specifically, the exine pattern and endoaperture shape help to elucidate their taxonomic rank at the subgenus level. The unidentified *Cotoneaster* specimen was nested seperately in the phenogram and the pollen characterictics may provide additional clues to describe this taxon as a new taxonomic level. However, further knowledge is needed to clarify its systematic position.

AUTHORS' CONTRIBUTIONS

Serap Işık Seylan: pollen morphological studies and writing. Zübeyde Uğurlu Aydın: data analyses, fieldwork and writing. Ali A. Dönmez: Project administration, fieldwork, review and editing. The authors declare that there is no conflict of interest.

ACKNOWLEDGEMENTS

This study was supported by Scientific and Technical Research Council of Türkiye with project number 111 T 850. The authors thank Emel Oybak Dönmez for English revision of the manuscript.

REFERENCES

Bartish IV, Hylmö B, and NYBOM H. 2001. RAPD analysis of interspecific relationships in presumably apomictic *Cotoneaster* species. *Euphytica* 120: 273–280. https://doi.org/10.1023/A:1017585600386

Borsch T, and Wilde V. 2000. Pollen variability within species, populations, and individuals, with particular reference to *Nelumbo*. In: Harley MM, Morton CM, Blackmore S [ed.], *Pollen and Spores: Morphology and Biology*, 285–299. Royal Botanic Gardens, Kew.

Byatt JI, Ferguson IK, and Murray BG. 1977. Intergeneric hybrids between *Crataegus* L. and *Mespilus* L.: a fresh look at an old problem. *Botanical Journal of the Linnean Society* 74(4): 329–343. https://doi.org/10.1111/j.1095-8339.1977.tb01185.x

Campbell CS, and Dickinson TA. 1990. Apomixis, patterns of morphological variation, and species concepts in subfam. Maloideae (Rosaceae). *Systematic Botany* 15(1): 124–135. https://doi.org/10.2307/2419022

- CAMPBELL CS, GREENE CW, and DICKINSON TA. 1991. Reproductive biology in Subfam. Maloideae (Rosaceae). Systematic Botany 16(2): 333–349. https://doi.org/10.2307/2419284
- CAMPBELL CS, EVANS RC, MORGAN DR, DICKINSON TA, and ARSE-NAULT MP. 2007. Phylogeny of subtribe Pyrinae (formerly the Maloideae, Rosaceae): limited resolution of a complex evolutionary history. *Plant Systematics and Evolution* 266: 119–145. https://doi.org/10.1007/s00606-007-0545-y
- Chang KC, Wang CC, Deng SL, Kono Y, Lu FY, and Peng CI. 2011a. Cotoneaster rosiflorus (Rosaceae), a new species from Taiwan. Botanical Studies 52: 211–218.
- Chang KC, Wang CM, Deng SL, and Wang CC. 2011b, A new species *Cotoneaster chingshuiensis* (Rosaceae) from Taiwan. *Taiwania* 56(2): 125–131.
- CHINNAPPA CC, and WARNER BG. 1982. Pollen morphology in the genus *Coffea* (rubiaceae): ii. pollen polymorphism. *Grana* 21(1): 29–37. https://doi.org/10.1080/00173138209427677
- Christensen KI. 1992. Revision of *Crataegus* Sect. *Crataegus* and Nothosect. Crataeguineae (Rosaceae-Maloideae) in the Old World. *Systematic Botany Monographs* 35: 1–199. https://doi.org/10.2307/25027810
- Dickinson TA. 2018. Sex and Rosaceae apomictics. *Taxon* 67(6): 1093–1107. https://doi.org/10.12705/676.7
- DICKORÉ WB, and KASPEREK G. 2010. Species of *Cotoneaster* (Rosaceae, Maloideae) indigenous to, naturalising or commonly cultivated in Central Europe. *Willdenowia* 40(1): 13–45. https://doi.org/10.3372/wi.40.40102
- EIDE F. 1981. Key for Northwest European Rosaceae pollen. Grana 20(2): 101–118. https://doi.org/10.1080/00173138109427651
- EVANS RC, and CAMPBELL CS. 2002. The origin of the apple subfamily (Maloideae; Rosaceae) is clarified by DNA sequence data from duplicated GBSSI genes. *American Journal of Botany* 89(9): 1478–1484. https://doi.org/10.3732/ajb.89.9.1478
- FLINCK KE, and HYLMÖ B. 1966. A list of series and species in the genus *Cotoneaster*. Botaniska Notiser 119: 445–463.
- Fryer J, and Hylmö B. 2009. Cotoneasters: A Comprehensive Guide to Shrubs for Flowers, Fruit, and Foliage. Timber Press, Portland and London.
- George D, and Mallery P. 2016. *IBM SPSS Statistics 23 Step* by Step. A simple Guide and Reference. Routledge, New York.
- GHOSH A, and SAHA I. 2017. Pollen morphological study of some selected Indian taxa of Rosaceae. *Indian Journal Applied and Pure Biology* 32(2): 121–130.
- GREGOR T. 2013. Apomicts in the vegetation of Central Europe. Tuexenia 33: 233–257.
- Grimsson F, Grimm GW, and Zetter R. 2017. Evolution of pollen morphology in Loranthaceae. *Grana* 57(1–2): 16–116. https://doi.org/10.1080/00173134.2016.1261939
- Gower JC. 1971. A general coefficient of similarity and some of its properties. *Biometrics* 27(4): 857–871. https://doi.org/10.2307/2528823

- Hammer Ø, Harper DAT, and Ryan PD. 2001. Past: paleontological statistics software package for education and data analysis. *Palaeontologia Electronica* 4(1): 1–9.
- Hebda RJ, and Chinnappa CC. 1990. Studies on pollen morphology of Rosaceae in Canada. *Review of Palaeobotany and Palynology* 64(1-4): 103–108. https://doi.org/10.1016/0034-6667(90)90123-Z
- Hebda RJ, and Chinnappa CC. 1994. Studies on pollen morphology of Rosaceae. *Acta Botanica Gallica* 141(2): 183–193. https://doi.org/10.1080/12538078.1994.10515150
- Hebda RJ, Chinnappa CC, and Smith BM. 1988. Pollen morphology of the Rosaceae of Western Canada: I. *Agrimonia* to *Crataegus. Grana* 27(2): 95–113. https://doi.org/10.1080/00173138809432836
- Hesse M, Halbritter H, Zetter R, Weber M, Bruchner R, Frosch-Radivo A, and Ulrich S. 2009. *Pollen Terminology. An Illustrated Handbook*. Springer-Verlag, Wien.
- HSIEH TH, and HUANG TC. 1997. Notes on the Flora of Taiwan (28)-the genus *Cotoneaster* Medik (Rosaceae). *Taiwania*. 42(1): 43–52.
- Işik S, and Dönmez AA. 2004. Pollen morphology of the three Pomoid Genera × *Malosorbus* Browicz, *Mespilus* L., and *Eriolobus* (Ser.) Roemer (Rosaceae). *Hacettepe Journal* of *Biology and Chemistry* 33: 65–75.
- Jarvis DI, Leopold EB, and Liu Y. 1992. Distinguishing the pollen of deciduous oaks, evergreen oaks, and certain rosaceous species of southwestern Sichuan Province, China. Review of Palaeobotany and Palynology 75(3–4): 259–271. https://doi.org/10.1016/0034-6667(92)90019-D
- KOEHNE E. 1893. *Deutsche Dendrologie*. Verlag von Ferdinand Enke, Stuttgart.
- Li QY, Guo W, Liao WB, Macklin JA, and Li JH. 2012. Generic limits of Pyrinae: Insights from nuclear ribosomal DNA sequences. *Botanical Studies* 53(1): 151–164.
- Lo EYY, and Donoghue MJ. 2012. Expanded phylogenetic and dating analyses of the apples and their relatives (Pyreae, Rosaceae). *Molecular Phylogenetics and Evolution* 63(2): 230–243. https://doi.org/10.1016/j. ympev.2011.10.005
- Meng KK, Chen SF, Xu KW, Zhou RC, Li MW, Dhamala MK, Liao WB, and Fan Q. 2021. Phylogenomic analyses based on genome-skimming data reveal cyto-nuclear discordance in the evolutionary history of *Cotoneaster* (Rosaceae). *Molecular Phylogenetics and Evolution* 158: 1070–1083. https://doi.org/10.1016/j. ympev.2021.107083
- MORGAN DR, SOLTIS DE, and ROBERTSON KR. 1994. Systematic and evolutionary implications of *rbcL* sequence variation in Rosaceae. *American Journal of Botany* 81(7): 890–903. https://doi.org/10.1002/j.1537–2197.1994.tb15570.x
- OYBAK DÖNMEZ E. 2008. Pollen morphology in Turkish *Crataegus* (Rosaceae). *Botanica Helvetica* 118: 59–70. https://doi.org/10.1007/s00035-008-0823-5
- Perveen A, and Qaiser M. 2014. Pollen Flora of Pakistan-LXXI. Rosaceae. *Pakistan Journal of Botany* 46(3): 1027–1037.

- Phipps JB, Robertson KR, Smith PG, and Rohrer JR. 1990. Checklist of the subfamily Maloideae (Rosaceae). *Canadian Journal of Botany* 68 (10): 2209–2269. https://doi.org/10.1139/b90-288
- POTTER D, ERIKSSON T, EVANS RC, OH S, SMEDMARK JEE, MORGAN DR, KERR M, ROBERTSON KR, ARSENAULT M, DICKINSON TA, and CAMPBELL CS. 2007. Phylogeny and classification of Rosaceae. *Plant Systematics and Evolution* 266: 5–43. https://doi.org/10.1007/s00606-007-0539-9
- Punt W, Hoen PP, Blackmore S, Nilsson S, and Le Thomas A. 2007. Glossary of pollen and spore terminology. *Review of Palaeobotany and Palynology* 143(1–2): 1–81. https://doi.org/10.1016/j.revpalbo.2006.06.008
- Punt W, and Hoen PP. 2009. The Northwest European Pollen Flora, 70: Asteraceae-Asteroideae. *Review of Palaeobotany and Palynology* 157(1–2): 22–183. https://doi.org/10.1016/j.revpalbo.2008.12.003
- RAEI NIAKI NA, ATTAR F, MIRTADZADINI M, and MAHDIGHOLI K. 2020. Pollen and floral micromorphological studies of the genus *Cotoneaster* Medik. (Rosaceae) and its systematic importance. *Caryologia* 73(3): 133–151. https://doi.org/10.13128/caryologia-569
- R CORE TEAM 2020. R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna.
- Reitsma TJ. 1966. Pollen morphology of some European Rosaceae. *Acta Botanica Neerlandica* 15(2): 290–307.
- Richards AJ. 1990. *Plant Breeding Systems*, Unwin Hyman. London.
- ROTHLEUTNER JJ, FRIDDLE MW, and CONTRERAS RN. 2016. Ploidy levels, relative genome sizes, and base pair composition in *Cotoneaster*. *Journal of the American Society for Horticultural Science* 141(5): 457–466. https://doi.org/10.21273/JASHS03776-16
- Sáez & Rosselló (Rosaceae), a new species from Major-

- ca (Balearic Islands, Spain). *Candollea* 67(2): 243–253. https://doi.org/10.15553/c2012v672a5
- Sax HJ. 1954. Polyploidy and apomixis in *Cotoneaster*. *Journal of the Arnold Arboretum* 35(4): 334–365.
- Song JH, Moon HK, and Hong, SP. 2016. Pollen morphology of the tribe Sorbarieae (Rosaceae). *Plant Systematics and Evolution* 302: 853–869. https://doi.org/10.1007/s00606-016-1303-9
- Song JH, Oak MK, Roh HS, and Hong SP. 2017. Morphology of pollen and orbicules in the tribe Spiraeeae (Rosaceae) and its systematic implications. *Grana* 56(5): 351–367. https://doi.org/10.1080/00173134.2016.1274334
- The Angiosperm Phylogeny Group. 2016. An update of the Angiosperm Phylogeny Group classification for the orders and families of flowering plants: APG IV. *Botanical Journal of the Linnean Society* 181: 1–20. https://doi.org/10.1111/boj.12385
- ULLAH F, GAO YD, ZAMAN W, and GAO XF. 2022. Pollen Morphology of Rosa sericea complex and their taxonomic contribution. Diversity 14(9): 705. https://doi.org/10.3390/d14090705
- Van Bergen MA, Van Der Ham RWJM, and Turner H. 1995. Morphology and evolution of *Arytera* pollen (Sapindaceae-Cupanieae). *Blumea: Biodiversity, Evolution and Biogeography of Plants* 40(1): 195–209.
- Wodehouse RP. 1935. Pollen Grains: Their structure, identification and significance in science and medicine. Mc Grew-Hill, New York.
- Zamani A, Attar F, and Maroofi H. 2010. Pollen morphology of the genus *Pyrus* in Iran. *Acta Biologica Szegediensis* 54(1): 51–56.
- Zhou LH, Wei ZX, and Wu ZY. 1999. Pollen morphology of Prunoideae of China (Rosaceae). *Acta Botanica Yunnan-ica* 21: 207–211.
- ZHOU L, WEI Z, and WU Z. 2000. Pollen morphology of Maloideae of China (Rosaceae). Acta Botanica Yunnanica 22(1): 47–52.