

Morphologic, Palynologic, and Phylogenetic Relationships of *Acantholimon* Species (Plumbaginaceae) Sharing Similar Habitats

Mevlüde Alev ATEŞ¹, Bahar KAPTANER İĞCİ*², Tuğrul Selçuk KÖRÜKLÜ³, Ahter FİŞNE², Zeki AYTAÇ²

¹Department of Agricultural Biotechnology, Faculty of Agriculture, Kırşehir Ahi Evran University, Kırşehir, Turkey;

²Department of Biology, Faculty of Science, Gazi University, Ankara, Turkey

³Department of Biology, Faculty of Science, Ankara University, Ankara, Turkey

ORCID ID: Mevlüde Alev ATEŞ: <https://orcid.org/0000-0002-2141-5438>; Bahar KAPTANER İĞCİ: <https://orcid.org/0000-0001-7102-6337>; Tuğrul Selçuk KÖRÜKLÜ: <https://orcid.org/0000-0002-7440-6818>; Ahter FİŞNE: <https://orcid.org/0000-0001-6895-8870>; Zeki AYTAÇ: <https://orcid.org/0000-0003-3244-3183>

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Abstract: The genus *Acantholimon* Boiss. is comprised of approximately 200 species worldwide and represented with 71 taxa 56% of which are endemic in Turkey. *Acantholimon anatolicum*, *A. ulicinum* var. *ulicinum*, *A. gemicianum*, *A. riyatguelii* and *A. acerosum* var. *acerosum* are discussed in the current study due to their similar habitats. Although *Acantholimon anatolicum*, *A. ulicinum* var. *ulicinum*, *A. gemicianum* are morphologically and taxonomically very similar to each other, *Acantholimon riyatguelii* and *A. acerosum* var. *acerosum* are different than other species with scale (2-3 not 4-5) and scapes lengths. To understand the relationships among these taxa, morphological, palynological, and molecular structures were studied. Both LM and SEM were used to obtain the morphological characters and ITS nuclear gene region was amplified for molecular characteristics of taxa. According to the data, in addition to the morphological similarities there are also palynological and molecular similarities.

Keywords: Ecology, Flora, Systematic, Turkey.

Benzer Yaşam Alanlarını Paylaşan *Acantholimon* Türlerinin (Plumbaginaceae) Morfolojik, Palinolojik ve Filogenetik İlişkileri

Öz: *Acantholimon* Boiss. cinsi. Dünya genelinde yaklaşık 200 tür ve 71 taksonla temsil edilir, bunların %56'sı Türkiye için endemiktir. Bu çalışmada *Acantholimon anatolicum*, *A. ulicinum* var. *ulicinum*, *A. gemicianum*, *A. riyatguelii* ve *A. acerosum* var. *acerosum* benzer habitatları paylaşmaları nedeniyle karşılaştırılmıştır. *Acantholimon anatolicum*, *A. ulicinum* var. *ulicinum*, *A. gemicianum* morfolojik ve taksonomik olarak birbirlerine çok benzerdir. Ancak *A. riyatguelii* ve *A. acerosum* var. *acerosum* diğer türlerden skalalarının 2-3 (4-5 değil) ve çiçeklenme boylarının yapraklardan uzun oluşuyla farklıdır. Bu taksonların ilişkilerini anlamak için morfolojik, palinolojik ve moleküler çalışmalar yapılmıştır. Morfolojik karakterleri elde etmek için hem ışık hem de elektron mikroskopu kullanılmış ve taksonların moleküler özellikleri için ITS nükleer gen bölgesi çalışılmıştır. Morfolojik benzerlikler verilerine ek olarak palinolojik ve moleküler veriler de benzerlik göstermektedir.

Anahtar kelimeler: Ekoloji, Flora, Sistematik, Türkiye.

1. Introduction

The genus *Acantholimon* Boiss. is comprised of approximately 200 species worldwide and most of them derived from its main diversity center Irano-Turanian Phytogeographic Region (Bokhari, 1970). The genus is represented with 25 species in Turkey. Moreover, 11 imperfectly known and unclearly recorded taxa were named as *Acantholimon* in Flora of Turkey (Bokhari and Edmondson, 1982). In recent years, the number of taxa has been increased up to 71 and forty of them were accepted as endemic. Therefore, the endemism ratio of the genus has become 56% in Turkey with recent studies (Akaydın & Doğan, 2012; Doğan & Akaydın, 2007; Yıldırım, 2009; Yıldırım, 2015; Kaptaner İğci, Körüklü, & Aytaç, 2017; Akaydın, 2018).

In taxonomical studies, *Acantholimon* species are classified by using their morphological features like perennial, laxly or densely pulvinate subshubs forming thorn cushions. Furthermore, the genus members have got ecological and ornamental importance with nicely colored

and long-lasting flowering periods (Muvaffak, Doğan, & Bilgin, 2001). Due to its importance, there were many studies about the taxa. Although one of the revision studies was performed in Turkey about the breed resulted in a new taxon, members of the genus has not been clearly classified yet (Doğan and Akaydın, 2007).

In the current study, our purpose is to investigate the taxonomic positions of *A. anatolicum* Yıld., *A. gemicianum* Kaptaner İğci, Körüklü & Aytaç, and *A. ulicinum* (Willd. ex Schultes) Boiss. var. *ulicinum* *A. acerosum* (Willd.) Boiss. var. *acerosum* and *A. riyatguelii* Yıldırım by using their morphological, palynological, and molecular structures through their shared habitats and their similar morphological structures.

The members of genus *Acantholimon* taxa usually preferred steppes, calcareous soils but some of them grow on serpentine main rocks. In this study, taxa grown in narrow areas and adapted to the gypsum soils were examined. *A. anatolicum*, *A. gemicianum* and *A. riyatguelii* have been identified in the last 10 years and they grow in

*Corresponding author: kaptanerbahar@yahoo.com

gypsum soils. *A. ulicinum* var. *ulicinum* and *A. acerosum* var. *acerosum* grow in both calcareous and gypsum soils and share the same habitats. Therefore, similarities of these 5 taxa, relations, and morphological structures were aimed to be examined.

2. Material and Methods

2.1. Morphological Studies

For morphological and palynological analysis, the materials were collected from different parts of Ankara and Eskişehir provinces (Turkey) during flowering and fruiting times (Tab.1, Fig.1). The morphologic characters were investigated by light and scanning electron microscope. Until studying they were stored in GAZI.

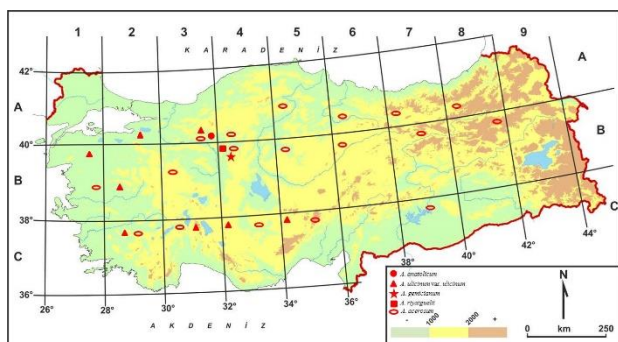


Figure 1: Distribution of the taxa in Turkey.

Tab. 1: Locations where the samples were collected and used in the studies.

Species	Localities	Collector number
<i>A. anatolicum</i>	Ankara: Beypazarı, Doğankonak village, 518 m, 13. 07. 2017, calcareous-gypsum steppe.	Körüklü 22015 & Aytaç
<i>A. ulicinum</i> var. <i>ulicinum</i>	Ankara: Beypazarı-Sekli road, 690 m, 13. 07. 2017, calcareous steppe.	Körüklü 22018 & Aytaç
<i>A. gemicianum</i>	Eskişehir: Sivrihisar, Balıkdamu, Ahiler-Kurtseyh, 14. 07. 2017, calcareous-gypsum steppe.	Körüklü 22023 & Aytaç
<i>A. riyatguelii</i>	Eskişehir: Sivrihisar, Balıkdamu, above Ertuğrulköy, 900 m, 14. 07. 2017, calcareous-gypsum steppe.	Körüklü 22021 & Aytaç
<i>A. acerosum</i>	Ankara: Beypazarı, Doğankonak village, 13. 07. 2017, calcareous-gypsum steppe.	Körüklü 22016 & Aytaç

2.2. Palynological Studies

Pollen grains of five taxa were studied by using LM (Light Microscopy) and SEM (Scanning Electron Microscopy). Firstly, for the LM analysis, pollen grains were treated with 70% alcohol to remove oily substances around grains and they were embedded in glycerin jell and stained with basic fuchsin by following the method of Wodehouse (1959). At least 30 fully developed grains per sample were measured under a Leica ICC50 HD microscope (1000×) and polar axis (P), equatorial axis (E), and exine thickness (Ex) were analyzed. For all data, minimum, maximum, and mean \pm standard deviations were provided.

Additionally, for SEM analyses, pollen grains were transferred directly to stubs with double-sided adhesive tape and their micrographs were obtained using Jeol-6060 SEM at Gazi University at an accelerating voltage of 15 kV. Diameter of lumina and thickness of muri were also

measured. For all analysis, the reference materials were stored at GAZI and ANK.

2.3. Molecular Studies

For molecular studies, leaf samples were obtained from natural habitats and optimized 2XCTAB method was used for total DNA isolation (Doyle 1990). The amplification of ITS region (ITS1+5.8S+ITS2) was done by using the primers of Hslao, Chatterton, Asay, & Jensen, K (1995). For PCR amplifications 5x FIREPol® Master Mix Ready to Load was used. The PCR reactions were performed with the steps of 95 °C for 5 min initial denaturation, followed by 30 cycle 95 °C for 30 sec, annealing temperature for 30 sec, 72 °C for 90 sec and 72 °C for 10 min as final extension. All the products were checked by using agarose gel (1.5%) and all purification and sequencing of samples were performed by RefGEN Biotechnology company (Ankara). All sequences were formed for the analysis by editing null bases composed of unsuccessful nucleotide peaks from raw data that were examined by Finch TV software Version 1.4.0-manufactured by Geopiza Research Team; (Patterson, Chamberlain, & Thayer, 2004-2006). The sequences were aligned by MUSCLE (Multiple Sequence Comparison by Log Expectation) tool (Edgar, 2004) of MEGA (Molecular Evolutionary Genetics Analysis) 7.0.9 software (Kumar, Stecher, & Tamura, 2016). Additionally, for constructing the phylogenetic tree, neighbor-joining (NJ) method (Saitou & Nei, 1987) which is the basic type of minimum evolutionary (ME) method, was used with bootstrap test analysis in MEGA software.

For all the samples studied, plants belonging to the collector number in the table are given.

3. Results

3.1. Morphological Features

The general morphological characteristic features of investigated species are being pulvinate, sub-shurblet, and glaucous (Fig. 2). Although *A. anatolicum*, *A. ulicinum* var. *ulicinum* and *A. gemicianum* are very similar to each other in morphological appearance; *A. anatolicum* leaves are glabrous and ciliate on their margins. Moreover, margins are glabrous or sparsely pubescent and scabrid in *A. ulicinum* var. *ulicinum* whereas they are puberulent in *A. gemicianum*. Leaf scapes are 5–8 cm long, branched, and represented in number 4–5 in *A. anatolicum* scapes. Although the scales are absent in *A. ulicinum* var. *ulicinum*, they are shorter in *A. gemicianum*. Even though the number of spikletes is 6–12 in *A. anatolicum*; it is 3–7 in *A. ulicinum* var. *ulicinum* and 3–5 in *A. gemicianum*. Additionally, spikletes' numbers are 7–20 in *A. riyatguelii* whereas 5–15 in *A. acerosum* var. *acerosum* (Tab. 2).

Unlike all the other species, *A. acerosum* var. *acerosum* has a wide range of habitat as steppe, calcareous soil, rocky igneous slopes, and sandy soil due to its large ecological tolerance. In the shed light on these, intra-species variations are quite extensive in the species. Although other species are in narrow habitats, intra-species variation is not seen between them.

3.1.1. Pollen Morphology

The size range of polar axis (P) varies between 46 μ m (in *A. ulicinum* var. *ulicinum*) and 81,6 μ m (in *A. Acerosum* var. *acerosum*), while the size range of equatorial axis (E) varies

gemicianum and *A. anatolicum*. Although *A. ulicinum* var. *ulicinum* and *A. anatolicum* species were located at the same subcluster, *A. gemicianum* was not very distinct than others. This means that, there were low number of base

substitutions resulted in the separation of this subcluster (Fig. 4). Moreover Moharrek et al. (2014) reported that the genetic divergence between the *Acantholimon* species was very close to each other according to ITS sequences (0.02).

Table 2: Morphological comparison of species.

	<i>A. anatolicum</i>	<i>A. ulicinum</i> var. <i>ulicinum</i>	<i>A. gemicianum</i>	<i>A. riyatguelii</i>	<i>A. acerosum</i>	
Habitus	Densely pulvinate, glaucous shrublet	Densely pulvinate, calcareous-punctate shrublet	Densely pulvinate, glaucous shrublet	Laxly dwarf caespitose, pale-green to glaucous Swollen-fleshy, linear to	Densely pulvinate, glaucous shrublet	
Leaves	Linear-triquetrous, 5–12 × 0.5–1 mm, glabrous, ciliate on margins	Linear-triquetrous, 5–20 × 1–1.2 mm, glabrous or sparsely pubescent	Linear-triquetrous, 7–10 × 0.5–1 mm, puberulent, calcareous punctate	oblong-linear, 3–8 × 1.5–2 mm, mucronate, pilose (mostly on margins), calcareous-punctate Simple, 0.8–1.5 cm, densely puberulent, as long as or slightly exceeding leaves	Linear or plano-triquetrous, 15–60 × 0.8–2.2 mm	
Scapes	5–8 cm long, branched	Very short or ± absent	2–5 cm long, branched	2–3, pilose, not overlapping, the lowermost shorter than internodes	Simple, longer than leaves	
Scales	4–5, shorter than internodes, strigillose all over	Absent	2–3, shorter than internodes, strigillose all over	Solitary, densely distichous, 15–40 mm long	Usually laxly distichous, longer than leaves	
Spikes	2–3, 15–20 mm long, imbricate or terminal	1–2, imbricate or densely distichous	3–5, 15–20 mm long, imbricate	7–20, 12–14 mm long	5–15	
Spikelets	6–12 in each spike, 1-flowered, 12–13 mm	3–7	3–5 in each spike, 1-flowered, 12–13 mm	6–7 mm long, ovate, cuspidate, densely pilose, calcareouspunctate, narrow hyaline margins	Equal to shorter than inner, triangular-lanceolate, 5–10 mm, with narrow hyaline margin	
Outer bracts	Unequal, puberulent all over, 4–5 mm (including aristate point), ovate, narrowly hyaline on margin	Glabrous or pubescent, oblong-lanceolate	Unequal, puberulent purplish apex, 4–5 mm (including aristate point), ovate, narrowly hyaline on margin	5–7 mm long, oblonglanceolate to narrowly lanceolate, cuspidate, pilose on midrib, hyaline except ribs	Oblong-lanceolate, to narrowly lanceolate, obtuse to acute cuspidate, with broad hyaline margin	
Inner bracts	7–7.5 mm (including aristate point c. 1 mm), oblong-lanceolate, obtuse, cuspidate with narrowly hyaline margin	Acuminate, long cuspidate cuspidate, hyaline except for the dark brown vein	7–7.5 mm (including aristate point c. 1 mm), oblong-lanceolate, obtuse cuspidate with narrowly hyaline margin	8–10 mm long, sparsely pilose on ribs; limb 5-lobed	Calyx tube pillose on veins, limb white or pale flesh color, usually 5 lobe	
Calyx	10–11 mm, tube densely pilose; limb 10-lobed, white; veins 5- brownish, pilose, expanded towards margins, not excurrent	11–12 mm, tube densely pilose; limb obscurely 5-10 lobed; veins purple, expanded towards margin, excurrent	10–11 mm, tube sparsely pilose; limb 10-lobed, white; veins 5-brownish, pilose, expanded towards margins, not excurrent	Deep marl-gypsum-rich soil	Gypsum-rich soils	Calcareous steppe
Habitat	Deep sandy gypsum-rich soil	Deep marl-gypsum, eroded hills	Deep marl-gypsum-rich soil			

Table 3: Summary of pollen morphological data for the *Acantholimon* taxa examined.

Taxa	P (µm)	E (µm)	Pollen shape	Exine thickness	Diameter of lumina	Thickness of muri
<i>A. anatolicum</i>	(53,7-) 57 ± 2,3 (-62,4)	(46-) 59,8 ± 3,4 (-65,2)	oblate - spheroidal	(6,7-) 8,1 ± 0,8 (-9,6)	(5,4-) 6,5 ± 1,5 (-11,3)	(2-) 2,1 ± 0,1 (-2,2)
<i>A. ulicinum</i> var. <i>ulicinum</i>	(46-) 58 ± 5,8 (-68,1)	(51,8-) 63,9 ± 6,4 (-79,6)	oblate - spheroidal	(5,7-) 7 ± 0,9 (-8,6)	(5,5-) 7,7 ± 1,6 (-10,6)	(2-) 2,1 ± 0,1 (-2,3)
<i>A. gemicianum</i>	(55,6-) 58 ± 1,7 (-61,4)	(59,5-) 62,6 ± 1,4 (-66,2)	oblate -spheroidal	(3,4-) 5,7 ± 0,8 (-7,6)	(1-) 1,8 ± 0,6 (-4,1)	(0,4-) 0,6 ± 0,1 (-0,8)
<i>A. riyatguelii</i>	(47-) 52,9 ± 2,5 (-57)	(51,8-) 57,3 ± 3,2 (66,6)	oblate - spheroidal	(5,7-) 7 ± 0,9 (-8,6)	(5,9-) 9,8 ± 2,5 (-15,1)	(0,8-) 1,1 ± 0,1 (-1,6)
<i>A. acerosum</i>	(66,2-) 71,4 ± 4,6 (-81,6)	(72-) 77,9 ± 4,8 (-92,1)	oblate - spheroidal	(4,8-) 5,5 ± 0,6 (-6,7)	(1,7-) 2,7 ± 0,5 (-4)	(0,4-) 0,8 ± 0,2 (-1,7)

4. Discussion

4.1. Morphological Studies

As a result of the obtained data, it could be said that *Acantholimon anatolicum*, *A. gemicianum*, and *A. ulicinum* var. *ulicinum* were very close to each other morphologically. Only some small differences in numbers of some morphological features separated these taxa from each other. For instance, *A. ulicinum* var. *ulicinum* was separated from other two taxa by its strigillose bracts,

strongly glaucous leaves, and 2-3 in number of scales on scape and *A. anatolicum* was distinguished from *A. gemicianum* with 4-5 in number of scales on scape and ciliate margin with glabrous leaves (not puberulous and calcareous punctate).

4.2. Palynological and Molecular Studies

On the other hand, the pollen grains did not give significant criteria for their distinctions in taxa. Therefore, pollen characters could be partly used as taxonomic

characters in order to distinguish the species in the taxa (Muvaffak et al. 2001). Moreover, it is seen that genetic similarities between *A. anatolicum* and *A. gemicianum* are very close to each other. In a phylogenetic study on the Plumbaginaceae family, the genus *Acantholimon* is seen as a separate clade (Koutroumpa et al. 2018). Moharrek, Osaloo, & Mostafa (2014) reported that genetic divergence between *Acantholimon* species was very close to each other according to ITS sequences (0.02). In our study, *A. anatolicum* and *A. gemicianum* were first studied phylogenetically. It is difficult to taxonomically classify those two taxa, which shared similar habitats and can be expected to have exchanged genes through hybridization. It could be more useful if more genetic regions belonging to the species that are morphologically close are studied in the future to understand the classification of the taxa.

When looking at the dendrogram, *A. riyatguelii* is located in the same branch with *A. acerosum* var. *acerosum*, although it grows in gypsum soils like *A. anatolicum* and *A. gemicianum*. This finding suggests that genetic similarities are not dependent to the gypsum rock type. So that, it should be investigated whether these species have selective characteristics related to main rock types which they grow.

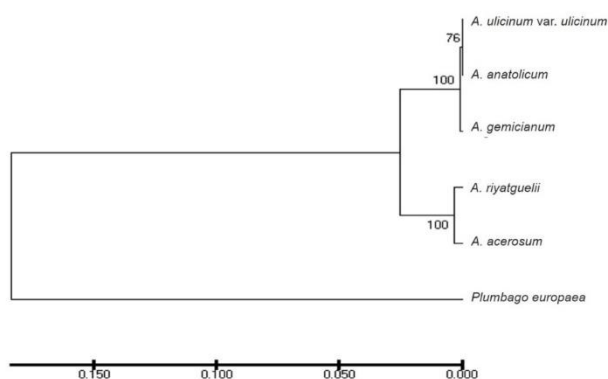


Figure 4: Phylogenetic dendrogram of ITS region drawn by using NJ method

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