## Morphological phylogenetic analysis of the genera Fragaria and Duchesnea in Iran Received: 06.10.2018 / Accepted: 19.12.2018

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#### **Abstract**

The phylogenetic relationships of the genera Fragaria and Duchesnea, and their closely related species are reconstructed using 17 morphological characters. Data matrix consisted of 11 taxa belonging to two species of Fragaria, two species of Duchesnea, two species of Potentilla, and also five species of Drymocallis rupstris, D. poteriifolia, Geum urbanum, Rosa centifilia, and Filipendula vulgaris. The two latter species were used as outgroups. The phylogenetic analysis was carried out using maximum parsimony (MP) and Bayesian analysis. The result showed that, Fragaria and Duchesnea are monophyletic taxa, and Potentilla reptans is the closest relative of Duchesnea. Two theca anthers, terminal to subterminal style and yellow petals are synapomorphic for Duchesnea indica and Potentilla reptans. In addition, six characters in Fragaria were genus specific. An identification key was also presented based on the studied morphological traits.

Keywords: Cladistics analysis, maximum parsimony, mock strawberry, related species, Rosaceae, strawberry

# مطالعه فیلوژنتیک ریختشناختی جنسهای Fragaria و Duchesnea از تیره گلسرخیان در ایران <sup>\*</sup> دریافت: ۱۳۹۷/۰۹/۲۸ پذیرش: ۱۳۹۷/۰۹/۲۸

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خلاصه

در این تحقیق، روابط فیلوژنتیکی دو جنس Fragaria و Duchesnea و گونههای خویشاوند وابسته نزدیک آنها متعلق به تیره گل سرخیان با استفاده از ۱۷ صفت ریختشناختی مورد بررسی قرار گرفت. ماتریس دادهها شامل ۱۱ آرایه متعلق به دو گونه گل سرخیان با استفاده از ۱۷ صفت ریختشناختی مورد بررسی قرار گرفت. ماتریس دادهها شامل ۱۱ آرایه متعلق به دو گونه اخیر، به عنوان برون گروه مورد استفاده قرار گرفتند. Fragaria میباشد. دو گونه اخیر، به عنوان برون گروه مورد استفاده قرار گرفتند. آنیجه نشان داد که Fragaria و Fragaria و Fragaria و جرهای و خامه آنالیز دادهها به دو روش بیشینه پارسیمونی (MP) و بایزین مورد تجزیه و تحلیل قرار گرفتند. نتیجه نشان داد که Duchesnea و خامه انتهایی تکنیا و گونه و گونه برای دو گونه اصلاح نزدیک ترین خویشاوند Duchesnea است. بساک دو حجرهای و خامه انتهایی تا تقریبا انتهایی از صفات مشترک پیشرفته برای دو گونه Duchesnea و Duchesnea و Potentilla reptans هستند. همچنین، شش صفت تفکیک کننده جنس Fragaria تشخیص داده شدند. براساس صفات ریختی گونههای مورد مطالعه کلید شناسایی ارایه شده است.

**واژههای کلیدی**: بیشینه پارسیمونی، تحلیل کلادیستیک، توت فرنگی، توت فرنگی کاذب، گل سرخیان، گونههای خویشاوند

#### Introduction

Fragaria L. and Duchesnea Sm. are two representative genera of the tribe Potentilleae (Hutchinson 1964, Dobeš & Paule 2010, Soják 2008), under the subfamily Rosoidea (Morgan 1994, Eriksson et al. 1998, 2003, Potter et al. 2007) and the family Rosaceae. The two genera resemble each other on their common names (Fragaria: strawberry and Duchesnea: Mock strawberry, Indian strawberry or false strawberry) (Zare et al. 2007), and also taxonomic complexities bases in their history of classification. Fragaria comprises about 30 species (Soják 2008) world widely including herbaceous plants, distributed in the north temperate and holarctic zones (Rousseau-Gueutin et al. 2008). Primarily, Linnaeus (1753) introduced Fragaria as one of the five relevant genera (Potentilla L., Comarum L., Fragaria L., Sibbaldia L., and Tormentilla L.) (Eriksson et al. 1998). However, many changes were made to this classification. Adanson (1763) separated Fragaria and Comarum from the former group. Wolf (1908) isolated plant having white petal, ternate leaves in the genus Fragaria. The genus comprises three species (including: Nubicola Lind., F. viridis Weston and F. vesca L.) in both Flora Iranica (Schonbeck-Temesy 1969), and Flora of Iran (including: Fragaria viridis, F. vesca, and F.  $\times$  ananassa (Weston) Duchesne ex Rozier (Khatamsaz 1993). Among them,  $F. \times ananassa$  is a hybrid of F. chiloensis (L.) Mill. and F. virginiana Mill. (Purgar et al. 2017).

The genus *Duchesnea* consists of about six herbaceous representatives growing in East Asia and North America (Eriksson *et al.* 2003). These plants are similar to strawberry in terms of life form but differ by their inedible dry fruits. For this reason, Smith (1818) separated and classified them under the genus *Duchesnea*.

However, taxonomic delimitation of the genus is influenced by several controversies and often is in concurrent with that of *Fragaria* (Syn.: *Fragaria indica* Andrews 479: 1807), *Fragaria malayana* Roxb.

520: 1832), and Potentilla [Syn: Potentilla trifida Lehm. 263: 1851, Potentilla indica (Andrews) Wolf 661: 1904] (Wolf 1908, Bate-Smith 1968, Kalman 1988). Wolf (1908) placed Potentilla indica close to P. reptans under Tormentillae group. Juzepchuk (1941) and Schonbeck-Temesy (1969) classified it with short rootstock; creeping stems; ternate leaves, solitary flowers; toothed (3-5) episepals; yellow petals; 20-25 stamens; numerous ovaries, fruit not separating from hypanthium; lateral or nearly terminal styles in a distinct genus Duchesnea. In addition, Soják (2004) based on style position (terminal style) and anther structure (two-theca anther), placed Duchesnea, under the subtribe Potentillinea. His main aim of this study was to conduct a phylogenetic analysis of Fragaria and Duchesnea based on the morphological traits to identify important traits that can provide efficient identification key at generic and species level.

Duchesnea comprises one species in the Flora Iranica, reported from Afghanistan and Pakistan (Schonbeck-Temesy 1969). However, some researches confirm presence of *D. indica* in Mazandaran (Zare et al. 2003) and Guilan provinces (Pourebrahim et al. 2018) also. The most important researches conducted in the genus Fragaria are those of Bringhurst & Senanayake (1966), Harrison et al. (1997), Hancock (1999), Staudt et al. (2003, 2006), Darrow (1966), Oda & Nishimura (2009), Retamales et al. (2005), Rousseau-Gueutin et al. (2009), Olbricht et al. (2012), DiMeglio et al. (2014), and Dujmovi'c Purgar et al. (2017). However, regarding the genus Duchesnea, only a few researches have been performed (Parado et al. 2012, Naruhashi & Sugimoto 1996).

The present survey, however, tries to answer the following questions:

- 1. Fragaria and Duchesnea; are they monophyletic?
- 2. Which of the two genera *Fragaria* and *Potentilla* are the closest relatives of *Duchesnea*?

#### **Materials and Methods**

In the current survey, 17 morphological characters belonging to 11 taxa were used in the cladistics analysis. These evidences obtained through the current morphological analyses and several populations from different species were examined. This includes 23 population of the genus Fragaria (including 17 populations of Fragaria vesca, and six populations of F. viridis), two species of the genus Duchesnea (Duchesnea indica and D. Chrysantha), and Potentilla reptans and P. micrantha as their closely related species. To this, we added two species Drymocallis (Drymocallis rupstris D. poteriifolia), Geum urbanum, Rosa centifolia, and Filipendula vulgaris. The two latter species considered as outgroups. Taxa selection follows the sequences of Schonbeck Temesy (1969), Soják (2012), and previous molecular phylogenetic studies of Eriksson et al. (2003), Potter et al. (2007), Dobeš & Paule (2010), and Faghir et al. (2014). Both dry and freshly collected samples (during 2016-17) were used for morphological evaluation. The herbarium specimens deposited in the herbaria of the Guilan University (GUH), Natural Resources Research Center of Guilan province (GILAN), and Tehran University (TUH). The voucher specimens of newly collected materials were deposited in the Guilan University [GUH. For identification purpose, Schönbeck-Temesy (1969), Khatamsaz (1993), and Soják (2004, 2012) were used].

Species used in this survey are presented in table 1.

- Data analysis and cladogram construction

Phylogenetic reconstruction was inferred using both maximum parsimony (MP) and Bayesian methods.

- Maximum parsimony analysis (MP)

Phylogenetic analyses of the data matrix (Table 3) were carried out using maximum parsimony analysis (MP) incorporated in the PAUP Ver. 4.0b10 software program, installed in Windows (Swofford 2002). The polarity of characters was determined using the outgroup method (Maddison et al. 1984). Heuristic search settings followed by all characters 'unord' type, having equal weight. Starting trees obtained via stepwise addition, branchswapping algorithm: tree-bisection-reconnection (TBR). Initial 'MaxTrees' setting = 100. 'MulTrees' option were turned on and steepest descent option were not affected. For character-state, optimization accelerated transformation (ACCTRAN) was performed. For improving, the trees' indices, successive re-weighting strategy (Farris 1969) were conducted. The reweighing of the traits continued until the trees' indices did not change in two separate analyzes after two replications. The bootstrap values (Felsenstein 1985) were calculated from 1000 and 20000 replicates. Re-scaled consistency index (RC), consistency index (CI), and retention index (RI) were also calculated.

Table 1. Plant samples used in the current study along with related data

Taxon	Collection data (all samples are from Iran)	Accession No.			
1. Fragaria	•				
F. viridis	Mazandaran province: Haraz Road, 2.4.2011, Ebrahimzadeh	5779 (GUH)			
F. vesca	Guilan province: Langeroud, Darvizhe, 13.4.2008	5794 (GUH)			
2. Duchesnea					
Duchesnea indica	Guilan province: Pirbazar, 29.6.2015, Ashouri	5793 (GUH)			
	Guilan province: Roodbar, 5.6.2011, Assadi	64085 (GILAN)			
D. chrysantha	Tehran province: Campus of Tehran University, 9.8.2016, Hadirzadeh and Rahmani	46799 (TUH)			
3. Potentilla					
Potentilla reptans	Guilan province: Siahkal, 16.6.2006, Faghir	36639 (TUH)			
P. micrantha	Guilan province: Asalem to Khalkhal, 6.6.2006, Faghir	36602 (TUH)			
4. Drymocallis					
Drymocallis rupestris	Mazandaran province: Firouzkooh, 12.6.2006, Attar	39000 (TUH)			
D. poteriifolia	Chaharmahal-o-Bakhtiari province: Yasouj, 17.6.2006, Attar and Zamani	36674 (TUH)			
5. Geum urbanum	Golestan province: 11 km to south Shahpasand, 10.5.1966, Papo	7521 (TARI)			
6. Rosa centifolia	Guilan province: Rasht, 15.4.2017, Pourebrahim	7540 (GUH)			
7. Filipendula vulgaris	Azerbaijan province: Kalibar, 21.6.1992, Ghahreman and Mozaffarian	17532 (TUH)			

Table 2. Morphological characteristics used in the current cladistics analysis (abbreviations and character state are presented)

	Character	Character state
1.	Stem height	0 = Tall (over 20 cm)  1 = Short (8-20 cm)
2.	Stem type	0 = Erect, not stoloniferous 1 = creeping, stoloniferous
3.	Leaf shape	0 = Pinnate, 1 = trifoliate, 2 = 5-foliate
4.	Leaflets shape	0 = Middle leaflet oblong, acute, 1 = middle leaflet widely ovate, obtuse
5.	Number of flower in each inflorescence	0 = Many, $1 = a few (2-4)$ , $2 = 1$
6.	Inflorescence type	0 = Not solitary (corymbiform, cyme)  1 = solitary
7.	Direction of sepals in the mature fruits	0 = Ascending-erect, $1 = $ recurvate
8.	Presence or absence of epicalyx (outer sepal)	0 = -, 1 = +
9.	Epicalyx number of teeth	0 = Entire, 1 = forked, 2 = trifide
10	. Receptacle enlargement after anthesis	0 = Receptacle protruding from calyx, 1 = receptacle not protruding from calyx
11	. Receptacle type	0 = Dry or flesh but dries at ripening $1 =$ fleshy at ripening
12	. Petal color	0 = White-yellow or pink, $1 = $ yellow
13	. Pistil hairy or glabrous	0 = Hairy, 1 = glabrous
14	. Anther type	0 = Single theca, $1 = $ two theca
15	. Filament type	0 = Not broadened, flat, $1 = broadened$ , flat
16	. Style position	0 = Lateral, 1 = terminal
17	. Fruit type	0 = Follicle, 1 = achene, 2 = fleshy aggregate

Table 3. Data matrix of studied species in the present study

Taxon	Character																
1 GAUII		2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17
1. Fragaria																	
Fragaria viridis	1	1	1	0	1	0	0	1	0	1	1	0	1	0	1	0	2
F. vesca	1	1	1	0	1	1	1	1	0	1	1	0	0	0	1	0	2
2. Duchesnea																	
Duchesnea indica	1	1	1	0	2	1	0	1	2	0	0	1	1	1	0	1	1
D. chrysantha	1	1	1	1	2	1	0	1	2	0	0	1	1	1	0	1	1
3. Potentilla								1									
P. reptans	1	1	2	0	1	1	0	1	0	0	0	1	1	1	0	1	1
P. micrantha	1	0	1	0	1	0	0	1	0	0	0	0	0	1	0	1	1
4. Drymocallis																	
Drymocallis rupestris	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1
D. poteriifolia	0	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0	1
5. Geum urbanum	0	0	0	0	0	0	0	1	0	0	0	1	0	?	0	1	1
6. Rosa centifolia.	0	0	0	0	0	0	0	0	0	0	0	1	0	?	0	0	1
7. Filipendula vulgaris	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0

#### - Bayesian method

Bayesian analyses were performed by MrModeltest Ver. 2.3 programs (Nylander 2004). The discrete approach to the morphological data matrix was adopted. All 17 characters were treated as discrete variables and treated as ordered (Table 1). Lewis (2001) bases the model used by MrBayes for morphological (standard) discrete data on the Markov model. Secondary probabilities were also determined. Analyses were repeated for several million generations and MrBayes performed two simultaneous analyses starting with different random trees (Nruns = 2). Four Markov chains were established for each tree and a tree from every 100 generations was sampled. After linear phase (3 million generations), burn-in period samples (including 25% of initial trees sampled) were removed and a tree with maximum 50% (majority rule consensus tree) was plotted. The values of posterior probability (PP) were measured and the final tree was plotted.

#### Results

The phylogenetic analysis based upon equally weighted characters resulted 100 most-parsimonious trees of 26 steps with consistency index (CI) = 0.8889 and retention index (RI) = 0.9189, Homoplasy index (HI) = 0.1308 and Rescaled consistency index (RC) = 0.8069. All characters were parsimony-informative. The strict consensus tree of obtained from morphological characters was shown in figure 1. It comprises one independent clade Filipendulla vulgaris and two lower small monophyletic groups (A and B) at the base of the tree. Group A includes Rosa centifolia and Geum urbanum and group B comprises two species of Drymocallis species. Species of Fragaria, Duchesnea, and Potentilla were united in a monophyletic group (O), from which two clades (O1 and O2) are derived. In clade O1 (with BP = 71%), Duchesnea indica and D. chrysantha formed a monophyletic group, and P. reptans and P. micranhta formed independent paraphyletic clades while clade O2 (with BP = 94%) comprised a monophyletic group of F. viridis and F. vesca.

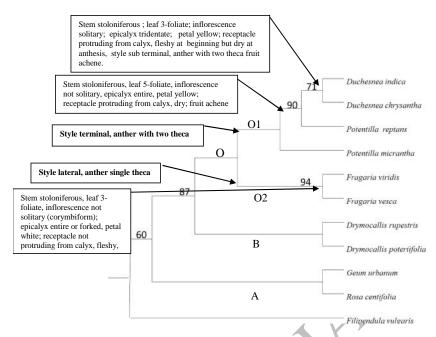


Fig. 1. A strict consensus tree obtained from 100 most parsimony trees of 17 morphological traits. Numbers above branches are bootstrap supports over 50%. Consistency index (CI) = 0.8889, and retention index (RI) = 0.9686. Clades characteristic features indicated in squares.

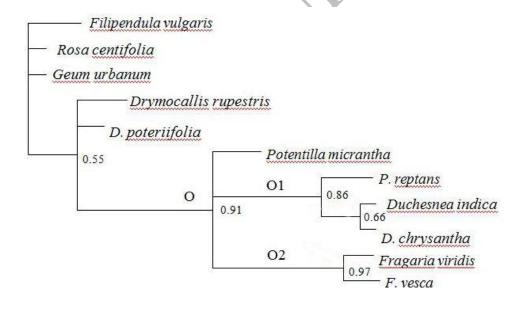


Fig. 2. Fifty percent majority-rule consensus tree obtained from a Bayesian analysis 17 morphological characters. Posterior probability (PP) indicated near each branch.

In the Bayesian analysis, 50% majority rule consensus tree was obtained (Fig. 2). At the base of this tree, *Filipendula vulgaris*, *Geum urbanum*, *Rosa centifolia*, *Drymocallis rupstris*, and *D. poteriifolia* formed paraphyletic independent branches while species

0.1

of Fragaria, Duchesnea, and Potentilla units are in a monophyletic group (O).

The representatives of this group form a tritomy including three species of *Duchesnea indica*, *D. chrysantha*, and *Potentilla reptans* in group Q1, two

species of *Fragaria vesca* and *F. viridis* in group O2, and *P. micrantha* in an independent branch.

#### Discussion

In the present cladistics analysis, the two species of *Drymocallis*, three species of *Geum urbanum*, *Rosa centifolia* and *Filipendula vulgaris* formed either monophyletic groups (in MP tree, Fig. 1) or paraphyletic independent branches (in Bayesian tree, Fig. 2). This corresponds to synplesiomorphic morphological characters such as stem height and type, pinnate leaf, more number of flower, and inflorescent not solitary (characters Nos 1–7, 10–11, 14 and 15) shared between these species.

The present phylogenetic analysis based on morphological traits showed monophyly of the two genera Fragaria and Duchesnea. This result supports former molecular researches (Eriksson et al. 2003, Töpel et al. 2011). The two genera have identical stem height and type (characters Nos 1 and 2), leaf shape (character No. 3, state 1), and direction of epicalyx (character No. 7, state 1). Using these evidences, some previous studies (Schonbeck-Temesy 1969, Juzepchuk 1941) classified the two genera under subfamily Rosoideae. However, based on Soják's (2004, 2012) morphological survey, Fragaria and Duchesnea have been separated in to two distinct subtribes (Fragarinea and Potentillinea), for having two theca anther (character No. 14, state 1) and subterminal style (character No. 16, state 1). The current result is in agreement with former molecular researches (Dobes & Paule 2010, Faghir et al. 2014, Zhang et al. 2017, Xiang et al. 2016, Heo et al. 2019).

Fragaria is characterized by its six specific characters as following: 3-foliate leaf (character No. 3, state 1), inflorescence not solitary (corymbiform) (character No. 6, state 0), entire or forked epicalyx (character No. 9, state 1), white petal (character No. 12, state 0), receptacle protruding from calyx (character No. 10, state 1), and fruit fleshy aggregate (character No. 17, state 2). The genus composed of euploid chromosome numbers form diploid (2n=2x=14), tetraploid (2n=4x)

=28), hexaploid (2n=6x=42), octoploid (2n=8x=56), and decaploid (2n=10x=70 (DiMeglio *et al.* 2014). Among them, *Fragaria vesca* is self-compatible, sympodial-running diploid, native from northern Europe and North American continent (Hummer *et al.* 2011). The later species, in Europe overlaps in distribution with another diploid, *F. viridis*, resulting a hybrid species ( $F. \times bifera$ ) (Staudt *et al.* 2003). However, direction of sepals in the mature fruits (character No. 7) and hairy or glabrous pistil (character No. 13), are importance characters isolating F. vesca from F. viridis.

In this analysis, Duchesnea with its terminal to subterminal style (character No. 16, state 1); two-theca anther (character No. 14, state 1), yellow petal (character No. 12, state 1), dentate epicalyx (character No.9, state 1), solitary flower (character No. 6, state 1), and positioned far from Fragaria and units with Potentilla species in O1 clades in both MP and Bayesian trees. The result supports previous molecular analyses based on both nrDNA ITS and chloroplast trnL-F sequence data studies (Eriksson et al. 2003, Faghir et al. 2014). Two theca anthers, terminal to subterminal style and yellow petals are the most informative characters that put the two species D. indica and P. reptans close to each other. These two characters are synapomorphic for both species. Adding D. chrysantha to the present analysis resulted a small monophyletic group composed of Duchesnea indica and D. chrysantha. The two morphologically species are very similar (Debes et al. 2011), but they can be recognized based on their receptacle enlargement (character No. 10), and middle leaflet shape (character No. 4). Duchesnea indica, is originated India with 2n=84, from while D. chrysantha has been reported in Japan, China, India, Korea, Taiwan, Philippines, and Indonesia, having chromosome number of 2n=14 (Naruhashi et al. 1999).

In conclusion, the current phylogenetic morphological based study revealed the phylogenetic relationship between two genera of *Fragaria* and *Duchesnea*. It was shown that, *Fragaria* and *Duchesnea* are monophyletic taxa and *Potentilla reptans* is the

closest relative of *Duchesnea*. Among the studied evidences, anthers structure, style positions are

synapomorphic traits that clearly separates the two genera *Duchesnea* and *Potentilla* from *Fragaria*.

### Identification key based on studied morphological traits for Fragaria and Duchesnea, and their allied species

1. Anthers with two thecae, style terminal to subterminal	2
Anthers with one thecae, style lateral	5
2. Leaf trifoliate, episepals tridentate	
Leaf 3- and 5-foliate, episepals entire	4
3. Receptacle not protruding from calyx, middle leaflet -oblong, acute	Duchesnea indica
Receptacle protruding from calyx, middle leaflet widely ovate, obtuse	D. chrysantha
4. Stem not stoloniferous, leaves tri-foliate, petal white	Potentilla micrantha
Stem stoloniferous leaves 5-foliate, petal yellow	P. reptans
5. Sepals in mature fruit ascending-erect	Fragaria viridi
Senals in mature fruit recurved below	F. vesco

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#### References

- Adanson, M. 1763. Familles des plantes. Paris. Chez Vincent. Imprimeur-Librarie de Mgr le. Comte de Provence, rue S. Servin.
- Bringhurst, R.S. & Senanayake, Y.D.A. 1966. The evolutionary significance of natural *Fragaria* chiloensis× F. vesca hybrids resulting from unreduced gametes. American Journal of Botany 53(10): 1000–1006.
- Darrow, G.M. 1966. The Strawberry: History, Breeding and Physiology. Holt, Rinehart and Winston, New York.
- DiMeglio, L.M., Staudt, G., Yu, H. & Davis, T.M. 2014.

  A phylogenetic analysis of the genus *Fragaria* (strawberry) using intron-containing sequence from the ADH-1 gene. PLoS One 9(7): e102237.
- Debes, M.A., Arias, M.E., Grellet-Bournonville, C.F., Wulff, A.F., Martínez-Zamora, M.G., Castagnaro, A.P. & Díaz-Ricci, J.C. 2011. White-fruited *Duchesnea indica* (*Rosaceae*) is impaired in ANS gene expression. American Journal of Botany 98(12): 2077–2083.

- Dobeš, C. & Paule, J. 2010. A comprehensive chloroplast DNA-based phylogeny of the genus *Potentilla* (*Rosaceae*): Implications for its geographic origin, phylogeography and generic circumscription. Molecular Phylogenetic and Evolution 56: 156–175.
- Domac, R. 1994. Flora Hrvatske. Priru nik za odre ivanje bilja. Školska knjiga, Zagreb.
- Eriksson, T., Donoghue, M.J. & Hibbs, M.S. 1998.

  Phylogenetic analysis of *Potentilla* using DNA sequences of nuclear ribosomal internal transcribed spacer (ITS), and its implications for the classification of *Rosoideae* (*Rosaceae*). Plant Systematic and Evolution 11: 155–179.
- Eriksson, T., Hibbs, M.S., Yoder, A.D., Delwiche, C.F. & Donoghue, M.J. 2003. The phylogeny of *Rosoideae* (*Rosaceae*) based on sequences of the internal transcribed spacers (ITS) of nuclear ribosomal DNA and the trnL/F region of chloroplast DNA. Intentional Journal of Plant Science 164: 197–211.

- Faghir, M.B., Attar, F., Farazmand, A. & Kazempur Osaloo, Sh. 2014. Phylogeny of the genus *Potentilla* (*Rosaceae*) in Iran based on nrDNA ITS and cpDNAtrnL-F sequences with a focus on leaf and style characters' evolution. Turkish Journal of Botany 38: 417–429.
- Farris, J.S. 1969. A successive approximation approach to character weighting. Systematic Zoology 18: 374–385.
- Felsenstein, J. 1985. Confidence limits on phylogenies: An approach using the bootstrap. Evolution 39: 783–791.
- Hancock, J.F., Lavín, A. & Retamales, J.B. 1999. Our southern strawberry heritage: *Fragaria chiloensis* of Chile. Horticultural Science 34(5): 814–816.
- Harrison, R.E., Luby, J.J. & Furnier, G.R. 1997. Chloroplast DNA restriction fragment variation among strawberry (*Fragaria* spp.) Taxa. Journal of the American Society for Horticultural Science 122(1): 63–68.
- Hebda, R.J. & Chinnappa, C.C. 1994. Studies on pollen morphology of *Rosaceae*. Acta Botanica Gallica 141(2): 183–193.
- Heo, K., Kim, Y., Maki, M. & Park, J. 2019. The complete chloroplast genome of mock strawberry, Duchesnea indica (Andrews) Th. Wolf (Rosoideae). Journal of Mitochondrial DNA Part B. Resources 4(1): 560–562. https://doi.org/10.1080/23802359.2018.1553527.
- Hutchinson, J. 1964. The genera of flowering plants, Vol. 1, Dicotyledons. Clarendon Press, Oxford.
- Hummer, K.E., Nathewet, P. & Yanagi, T. 2009.
  Decaploidy in *Fragaria iturupensis (Rosaceae)*.
  American Journal of Botany 96(3): 713–716.
- Hummer, K.E. & Janick, J. 2009. *Rosaceae*: taxonomy, economic importance, genomics. Pp. 1–17. *In*: Genetics and genomics of *Rosaceae*. Springer, New York.
- Hummer, K.E., Bassil, N. & Njuguna, W. 2011. Fragaria In Wild Crop Relatives: Genomic and

- Breeding Resources. Springer, Berlin, Heidelberg. Pp. 17–44.
- Juzepczuk, S.W. 1941. Alchemilla. In: Komarov, V.L. (ed.). Flora of USSR, Vol. 10, Izd Akad Nauk SSSR, Moskva-Leningrad. Pp. 289–410.
- Khatamsaz, M. 1993. Flora of Iran. *Rosaceae*. Research Institute of Forests and Rangeland Press. Tehran 6: 88–140.
- Linnaeus, C. 1753. *Fragaria (Rosaceae)*. P. 492. *In*: Species Plantarum, Vol. 2. Laurentius Salvius, Stockholm.
- Lei, J., Li, Y., Du, G., Dai, H. & Deng, M. 2005. A natural pentaploid strawberry genotype from the Changbai Mountains in northeast China. Horticultural Science 40: 1194–1195.
- Lewis, P.O. 2001. A likelihood approach to estimating phylogeny from discrete morphological character data. Systematic Biology 50: 913–925.
- Metcalfe, C.R. & Chalk, L. 1957. Anatomy of dicotyledons II. Claredon Press, Oxford.
- Maddison, W.P., Donoghue, M.J. & Maddison, D.R. 1984. Outgroup analysis and parsimony. Systematic Zoology 33: 83–103.
- Morgan, D.R., Soltis, D.E. & Robertson, K.R. 1994.
  Systematic and evolutionary implication of rbCL sequence variation in *Rosaceae*. American Journal of Botany 12: 890–903.
- Naruhashi, N. & Sugimoto, M. 1996. The floral biology of *Duchesnea (Rosaceae)*. Plant Species Biology 11(3): 173–184.
- Naruhashi, N., Iwatsubo, Y. & Peng, C.I. 1999. Cytology, flower morphology and distribution of *Fragaria hayatai* Makino (*Rosaceae*). Journal of Phytogeography and Taxonomy 47: 139–143.
- Nylander, J.A., Ronquist, J.P., Huelsenbeck, F. & Nieves-Aldrey, J.L. 2004. Bayesian phylogenetic analysis of combined data. Systematic Biology 53: 47–67.
- Oda, Y. & Nishimura, K. 2009. Ichigo-Japanese name of strawberry- its origin and historical change of

- phonogram for strawberry in Japan. Acta Horticulturae 842: 611–618.
- Olbricht, K., Ludwig, A., Ulrich, D., Spangenberg, R., Guenther, M. & Neinhuis, C. 2012. Leaf morphology and anatomy in the genus *Fragaria*: implications for resistances. VII International Strawberry Symposium 1049. Pp. 269–273.
- Pardo, E.M., Grellet, C.F., Salazar, S.M., Castagnaro, A.P., Diaz Ricci, J.C. & Arias, M.E. 2012. Histopathology of the resistance to *Colletotrichum gloeosporioides* of wild strawberries and species related to commercial strawberry. Australian Journal of Crop Science 6(7): 1147–1153.
- Potter, D., Luby, J.J. & Harrison, R.E. 2000.

  Phylogenetic relationships among species of 
  Fragaria (Rosaceae) inferred from non-coding 
  nuclear and chloroplast DNA sequences.

  Systematic Botany 25(2): 337–348.
- Potter, D., Eriksson, T., Evans, R.C., Oh, S., Smedmark,
  J.E.E., Morgan, D.R. & Campbell, C.S. 2007.
  Phylogeny and classification of *Rosaceae*. Plant
  Systematics and Evolution 266: 5–43.
- Pourebrahim, S., Faghir, M.B. & Attar, F. 2018. Report of new species of *Duchesnea indica* from *Rosaceae* for Guilan province. The International Conference on Agricultural Science. Medicinal Plants and Traditional Medicine. Mashhad, Iran.
- Purgar, D.D., Duralija, B., Vokurka, A., Todori, M. & Židovec, V. 2017. The Genus *Fragaria* in Croatia. Die Gattung *Fragaria* in Kroatien. Erwerbs-Obstbau 59(3): 227–233.
- Retamales, J.B., Caligari, P.D.S., Carrasco, B. & Saud, G. 2005. Status of the Chilean native strawberry and the research needs to convert the species into a commercial crop. Horticultural Science 40: 1633–1644.
- Rousseau-Gueutin, M., Gaston, A., Aïnouche, A., Aïnouche, M.L., Olbricht, K., Staudt, G. & Denoyes-Rothan, B. 2009. Tracking the evolutionary history of polyploidy in *Fragaria* L. (strawberry): new insights from phylogenetic

- analyses of low-copy nuclear genes. Molecular Phylogenetic and Evolution 51(3): 515–530.
- Schönbeck-Temesy, H. 1969. *Duchesnea (Rosaceae)*. Pp. 77–78. *In*: Flora Iranica, Vol. 66 (Rechinger, K.H., ed.). Akademische Druck u. Verlagsanstalt, Graz.
- Soják, J. 2004. Potentilla L. (Rosaceae) and related genera in the former USSR. Notes on Potentilla XVI. Botanische Jahrbücher für Systematik 125: 253–340.
- Soják, J. 2008. Notes on *Potentilla* 21. A new division of the tribe *Potentilleae* (*Rosaceae*) and notes on generic delimitations. Botanische Jahrbücher für Systematik 127: 349–358.
- Soják, J. 2010. Argentina hill, a genus distinct from *Potentilla* (*Rosaceae*). Thaiszia Journal of Botany Košice 20: 91–97.
- Soják, J. 2012. *Potentilla* L. (*Rosaceae*) and related genera in Asia (excluding the former USSR), Africa and New Guinea-Notes on *Potentilla* XXVIII. Plant Diversity and Evolution 130: 7–157.
- Staudt, G. 1989. The species of *Fragaria*, their taxonomy and geographical distribution. Acta Horticulturae 265: 23–33.
- Staudt, G. 2003. Notes on Asiatic *Fragaria* species: III. *Fragaria orientalis* Losinsk and *Fragaria mandshurica* sp. nov. Botanische Jahrbücher für Systematik 124(4): 397–419.
- Staudt, G. 2006. Himalayan species of *Fragaria* (*Rosaceae*). Botanische Jahrbücher für Systematik 126(4): 483–508.
- Staudt, G. 2008. Strawberry Biogeography, Genetics and Systematics. VI International Strawberry Symposium, International Society for Horticultural Science. Huelva, Spain.
- Staudt, G.S., Schneider, S., Scheewe, P. Ulrich, D. & Olbricht, K. 2008. *Fragaria iturupensis*, a new source for strawberry improvement. VI International Strawberry Symposium 842: 479–482.

- Töpel, M., Lundberg, M., Eriksson, T. & Eriksen, B. 2011. Molecular data and ploidal levels indicate several putative allopolyploidization events in the genus *Potentilla* (*Rosaceae*). PLoS Currents 3.
- Wolf, F.T. 1908. Monographie der Gattung *Potentilla*. Bibliotheca Botanica 16: 1–713.
- Xiang, Y., Huang, C., Hu, Y., Wen, J., Li, S., Yi, T., Chen, H., Xiang, J. & Ma, H. 2016. Evolution of *Rosaceae* fruit types based on nuclear phylogeny in the context of geological times and genome duplication. Molecular Biology and Evolution 34(2): 262–281.
- Zare, H., Ramezani, K.E. & Amini, T. 2007. A record of Duchesnea indica (Rosaceae) in Iran, its westernmost distributional limit in Asia. Iranian Journal of Botany 13(2): 93–94.
- Zhang, S.D., Jin, J.J., Chen, S.Y., Chase, M.W., Soltis, D.E., Li, H.T., Yang, J.B., Li, D.Z. & Yi, T.S. 2017. Diversification of *Rosaceae* since the late cretaceous based on plastid phylogenomics. New Phytologist 1–13.

