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A phylogenetic analysis of *Linaria* (Plantaginaceae) species from Iran based on ITS sequence data

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ABSTRACT

Linaria Miller With more than 150 taxa is the largest genus of the tribe Antirrhineae. This study was carried out on the species of *Linaria* that growing in Iran. Some of these species are native to Iran. Internal transcribed spacer (ITS) sequences were obtained for 37 samples representing 6 sections of *Linaria* recognized by recent taxonomic treatments from Iran. In addition, we used 82 previously ITS sequences from GenBank to test the monophyly of *Linaria*. Phylogenetic analysis were conducted using Bayesian inference and maximum likelihood. The results indicate that *Linaria* species from Iran constituted a monophyletic group within the Antirrhineae. data analysis indicates that the classification of species according winged or wingless seeds so not true, so seed morphology especially seed wing that have been considered as useful for the taxonomy of *Linaria* species, appears to be a homoplasious character in *Linaria*. Bayesian inference and maximum parsimony analyses confirmed Two sections of *Linaria* including *Macrocentrum* and *Versicolores* from Iran are monophyletic, while monophyly of sections *Linaria*, *Diffusae*, *Supinae* and *Speciosae* are unsupported by this results. To determine the evolution of *Linaria* use of morphological characteristics coupled with molecular data will be most effective.

Keywords: Iran, *Linaria*, phylogeny, ITS.

INTRODUCTION

Linaria Miller With more than 150 taxa is the largest genus of the tribe Antirrhineae [28]. distributed in the Northern hemisphere, its species living in Europe, Asia and North Africa [19]. *Linaria* was identified as a taxonomic being as soon as the time of pre-Linnaean botanists [22, 29]. At first Linnaeus (1753) identified *Linaria* species inner genus *Antirrhinum* [20]. Miller (1754) first logical description for *Linaria* supplied [21]. *Linaria* species are annual or perennial herbs, with heteromorphic shoots, sessile leaves and racemose inflorescences [26]. for the taxonomy of *Linaria* species, seed morphology was applicable. Small seeds of *Linaria* were encircled by capsules, they were winged seeds or wingless seeds. The wingless seeded species by Viano [32, 33], and the winged seeded species by Valde's (1970) established [30]. The winged seeds species designated *Discoideae* and the wingless seeds species introduced *Oblongae* by Boissier (1879)[3]. Viano's theory called wingless and winged seeds formed

distinct natural sister lineages in *Linaria* species [32, 33]. Sutton(1988) recognized seven sections mainly based on seed morphology within *Linaria* widely accepted today [28]. Species of sections *Linaria*, *Supinae* and *Pelisserianae* have discoid and usually winged seeds. While species of sections *Versicolores*, *Speciosae*, *Diffusae* and *Macrocentrum* have nondiscoid and wingless seeds. In Iran, 35 species of *linaria* belonging to the sections *Linaria* (20 spp.), *Speciosae* (6 spp.), *Supinae* (4 spp.), *Macrocentrum* (2 spp.), *Diffusae* (2 spp.) and *Versicolores* (1 spp.) are present [3, 23, 28, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17]. *Linaria* species that distribution in different regions of Iran, showed in Table 1 [7]. The aims in this study were to test the naturalness of the seven sections of Sutton's (1988) classification that six sections of them are present in Iran and to test the hypothesis of a basal divergence between two lineages of *Linaria*, one with winged seeds and the other with wingless seeds [28].

Table 1. Distribution of Linuria species in Iran

Geographical abbreviations: C, center Iran; S, southern Iran; SE, south-eastern Iran; N, northern Iran; NE, north-eastern Iran; NW, north-western Iran; E, eastern Iran; W, western Iran [7].

sections	Taxon	Distribution in Iran	Major Features of morphological traits of Linaria [28].	
Linaria	<i>L. vulgaris</i> Miller	C	Seed form: Discoid Seed wing: Wing Duration: Perennial Stigma: Entire	
	<i>L. odora</i> (Bieb) Fischer	C, NE		
	<i>L. khorasanensis</i> S. M. M. Hamdi and M. Assadi	NE		
	<i>L. leptoceras</i> Kuprian	NE		
	<i>L. striatella</i> Kuprian	C, NE		
	<i>L. baminaica</i> Patzak in Koie and Rech. f	NE		
	<i>L. michauxii</i> Chavannes	C, W, NW, E, SE,NE		
	<i>L. farsensis</i> S. M. M. Hamdi and M. Assadi	C		
	<i>L. pyramidalis</i> (Vent) F. G. Dietr.	Subsp. <i>pyramidalis</i>		NW
		Subsp. <i>Koptedaghensis</i>		NE
	<i>L. fastigiata</i> Chavannes	C, W, NW		
	<i>L. nurensis</i> Boiss. and Hausskn.	C		
	<i>L. remotiflora</i> Patzak in Rech. f.	C		
	<i>L. kurdica</i> Boiss. and Hohen. In Boiss.	Subsp. <i>kurdica</i>		W, NW
		Subsp. <i>Pycnophylla</i>		W, NW
	<i>L. lineolata</i> Boiss.	C, NW		
<i>L. karajensis</i> S. M. M. Hamdi and M. Assadi	C			
<i>L. elymatica</i> (Boiss.) Kuprian	N, W, C			
<i>L. azerbaijanensis</i> S. M. M. Hamdi and M. Assadi	NW			
<i>L. khalkhalensis</i> S. M. M. Hamdi and M. Assadi	N, NW			
<i>L. shahroudensis</i> S. M. M. Hamdi and M. Assadi	C			
<i>L. guilanensis</i> S. M. M. Hamdi and M. Assadi	N			
Diffusae	<i>L. bousherensis</i> S. M. M. Hamdi and M. Assadi	S	Seed form: Nondiscoid Seed wing: No wing Duration: Annual or perennial Stigma: Entire	
	<i>L. albifrons</i> (Sibth. and Sm.) Steudel.	S		
Speciosae	<i>L. mazandaranensis</i> S. M. M. Hamdi and M. Assadi	N	Seed form: Nondiscoid Seed wing: No wing Duration: Perennial Stigma: Entire	
	<i>L. genistifolia</i> (L.) Miller	N, NW		
	<i>L. golestanensis</i> S. M. M. Hamdi and M. Assadi	N		
	<i>L. dalmatica</i> (L.) Miller	N, W, NW		
	<i>L. grandiflora</i> Desf, Ann.	W, NW		
<i>L. orientalis</i> S. M. M. Hamdi and M. Assadi	N			
Macrocentrum	<i>L. chalepensis</i> (L.) Miller	Subsp. <i>chalepensis</i>	N, NW, W, C, NE, S	Seed form: Nondiscoid Seed wing: No wing Duration: Annual Stigma: Entire
		subsp. <i>gorganensis</i>	N	
	<i>L. armenica</i> Chavannes.	NW		
Supinae	<i>L. arvensis</i> (L.) Desf.	S	Seed form: Discoid Seed wing: Wing Duration: Annual or perennial Stigma: Entire	
	<i>L. simplex</i> (Willd) DC.	N, NW, C, NE		
	<i>L. kavirensis</i> S. M. M. Hamdi and M. Assadi.	C, NE		
	<i>L. micrantha</i> (Cav.) Hoffmanns. and Link.	W, S		
Versicolores	<i>L. iranica</i> S. M. M. Hamdi and M. Assadi	S	Seed form: Nondiscoid Seed wing: No wing Duration: Annual or perennial Stigma: Divided	

Table 2. List of taxa investigated in our analysis and herbaria where the vouchers are deposited with Genbank accession numbers
 TARI= herbarium of Research Institute of Forests and Rangelands, IAUH= Islamic Azad University Avicennia herbarium

Species	Origin, voucher	ITS Genbank accession numbers
<i>L. vulgaris</i>	Iran: prov. Tehran; Lawasan, Galendoak, 1700m, Mozzafarian, (52344 TARI).	KJ747021
<i>L. odora</i>	Iran: prov. Khorasan: Neshabbour, Dizbade-Olia, 1800m, Pariab and Abbasi, (8825 TARI).	KJ747015
<i>L. korasanensis</i>	Iran: prov. Khorasan, Mashad, SW of Moghan, 2300 m, Assadi and hamdi, (85531 TARI).	KJ747008
<i>L. striatella</i>	Iran: prov. Tehran, 10 km Roudehen to Firouzkouh, 2200m, Rahmani, IAUH without herbariums number.	KJ747020
<i>L. bamanica</i>	Iran: prov. Khorasan: Torbate-Jam, North of Torbate-Jam, 10 km to Bani-Tak, 1050 m, Jouharchi, (30649 TARI).	KJ747001
<i>L. michauxii</i>	Iran: prov. Tehran; 10km Tehran- Karaj highway, 1400m, Rahmani, IAUH without herbariums number.	KJ747013
<i>L. farsensis</i>	Iran: prov. Fars; Abadeh, Soghad, Dashte Ayon, 2100-2270 m, Wendelbo and Foroughi, (17870 TARI).	KJ747003
<i>L. pyramidalis subsp. pyramidalis</i>	Iran: Prov. Azerbaijan; between Urmia and Salmas, 1900m, Assadi, (78929 TARI).	KJ747017
<i>L. pyramidalis subsp. kopetdaghensis</i>	Iran: prov. Khorasan: Bojnurd; 40km western-south Bojnurd, Kuh-Sakuh, 2700, Mehregan, (13934 TARI).	KJ747016
<i>L. fastigiata</i>	Iran; prov. Hamedan; Ghahavand, Biukabad, to Shahbodagh, Aghdash maintain, slope of western, 1800-2300m, Mozzafarian, (64470 TARI).	KJ747004
<i>L. remotiflora</i>	Iran; prov. Kogiloueh and Bouyerahmad; between Yasouj and Dehdasht, Dilgoon, Saverz maintain, 2200-2300m, Assadi and Abouhamzeh, (46411 TARI).	KJ747018
<i>L. nurensis</i>	Iran: prov. Fars; Nourabad, 44km next Fahelian, toward Reshk, 1800m, Mozzafarian, (45932 TARI)	KJ747014
<i>L. kurdica subsp. kurdica</i>	Iran; prov. Kourdestan; Sanandaj, 25km sout-east of Naran village, 2200-2600m, Assadi, (60498 TARI).	KJ747009
<i>L. kurdica subsp. pycnophylla</i>	Iran: Prov. Azerbaijan; Jolfa, 25km sout-east Jolfa, Gheshlagh village, Kuh-Gholenj, 2100-2700m, Assadi and Shahsavari, (65736 TARI).	KJ747010
<i>L. lineolata</i>	Iran: prov. Tehran; Lawasan between Oshan and Lawasan, 1750m, Rahmani, without herbariums number.	KJ747012
<i>L. karajensis</i>	Iran; prov. Tehran; shahrestanac, Kuh- Touchal, 3150 m, Riazi, (8470 TARI).	KJ747006
<i>L. elymatica</i>	Iran; prov. Kerman; Kerman, Lalehzar maintain, 3200 m, Froughi and Assadi, (16360 TARI).	KJ747002
<i>L. azerbaijanensis</i>	Iran: Prov. Azerbaijan; Tabriz, beginning of road to Tehran, near Ghori gol, pass of Shebeli, 2300 m, Assadi(85348 TARI).	KJ747000
<i>L. khalkhalensis</i>	Iran: Prov. Azerbaijan; Khalkhal, 48 km Masooleh to Khalkhal, 1200 m, Assadi (86496TARI).	KJ747007
<i>L. shahroudensis</i>	Iran: Prov. Semnan, Shahroud, 15 km of Tash to Gorgan, 2800m, Assadi and hamdi (85676 TARI).	KJ747019
<i>L. guilanensis</i>	Iran: Prov. Gazvin; 50 km of Gazvin-rasht route, 1700m, Rahmani, IAUH without herbariums number.	KJ747005
<i>L. bousherensis</i>	Iran: Prov. Boushehr, 43 km to Ganaveh, 180 m, 29.02.1972, Foroughi (3107 TARI).	KJ747025
<i>L. albifrons</i>	Iran: Prov. Boushehr, 30 km to Bandare-Amir, 20 m, Runemark and Mozzafarian (26985 TARI).	KJ747022
<i>L. mazandaranensis</i>	Iran: Prov. Mazandaran; Ghaem-shahr, toward Firouz-Kuh, 8km west-sout of pole-sefid, 1270m, Moosavi (33702 TARI).	KJ747033
<i>L. genistifolia</i>	Iran: Prov. Mazandaran, Ramsar, Javaherdeh, western-south, 2500-2750 m, Wendelbo and Maassoumi, (20889 TARI).	KJ747028
<i>L. golestanensis</i>	Iran: Prov. Golestan; east-south Maraveh-Tapeh, Shalmi mountain 1200m, Faghih-Nia and Zangouii (32889 TARI).	KJ747029
<i>L. dalmatica</i>	Iran: Prov. Azerbaijan; Sarab, Saraban Kouh, Termeh (39071 TARI).	KJ747027
<i>L. grandiflora</i>	Iran: Prov. Azerbaijan; Ahar, 11km North of Azghan, 1840-2200mm Termeh and moosavi (38822 TARI).	KJ747030
<i>L. orientalis</i>	Iran: Prov. Semnan, Shahroud, to azadshahr, Khosh Yielagh, Riazi (8496 TARI).	KJ747035
<i>L. chalepensis</i>	Iran; prov. Kogiloueh and Bouyerahmad; Dogonbadan, 11km to Charam, 1100-1400 m, Assadi and Abouhamzeh, (38592 TARI).	KJ747026
<i>L. armenica</i>	Iran: Prov. Azerbaijan; 5km Khoy to Siah Cheshmeh, 1880m, Hamdi (82225 TARI).	KJ747023
<i>L. arvensis</i>	Iran: Prov. Khuzestan; Dehdez, Gharun-Kuh, 1000-1500m, Mozzafarian, (74494 TARI).	KJ747024
<i>L. simplex</i>	Iran: Prov. Guilan; Bandar Anzali, 30- 50 m, Mozzafarian, (75120 TARI).	KJ747036
<i>L. kavirensis</i>	Iran: Prov. Yazd, Taft, Taft-kuh, 1000-1300m, Dehghanzadeh, (26052 TARI).	KJ747032
<i>L. micrantha</i>	Iran: Prov. Khuzestan; Behbahan, 320m, Mozzafarian, (62494 TARI).	KJ747034
<i>L. iranica</i>	Iran: Prov. Kerman; Jiroft, 1000-1500m, Mirtajeddiny without herbariums number.	KJ747031

MATERIALS AND METHODS

Samples were collected in the field and dried in silica gel or obtained from herbaria in Iran (TARI, IAUH), [18]. Phylogenetic reconstructions were performed in 37 samples of *Linaria* presented in six sections from Iran. Table 2 lists all taxa used in this study and summarizes sources, voucher specimen data and GenBank accession numbers. Total DNA was extracted using the DNeasy Plant Mini kit (Qiagen, Germany). We amplified the ITS region (ITS1-5.8S-ITS2) of the nuclear ribosomal DNA using primer combinations AB101 and AB102 primers: a forward primer AB101 annealing, 5'-ACG AAT TCA TGG TCC GGT GAA GTG TTC G-3', and a reverse primer (AB102) annealing, 5'-TAG AAT TCC CCG GTT CGC TCG CCG TTA C-3'. The PCR protocol for ITS region included: 25 cycles of 1 min denaturation (94°C), 1 min annealing (54°C), and 2 min, 30 s elongation (72° C), with two additional seconds elongation per cycle [5].

PHYLOGENETIC ANALYSIS

Phylogenetic reconstructions were performed in 37 samples of *Linaria* in six sections from Iran. In this study We used the ITS sequence of 34 species of *Linaria*, 37 species of *Antirrhinum*, 4 species of *Galvezia*, 2 species of *Maurandya* and 2 species of *Kickxia* from GenBank. List of non-Iranian taxa used in our analysis with GenBank accession numbers showed in Table 3. We also used the ITS sequence of *Lafuentea rotundifolia*, from GenBank as the outgroup [1]. Matrices were analyzed with PAUP*4.0b10, with the following options: heuristic search with 1,000 random-addition-sequence replicates; tree bisection-reconnection (TBR) branch swapping; "collapse zero length branches;" saving all most parsimonious trees. Character state changes were treated as equally weighted. Nonoverlapping parsimony informative indels were coded as binary characters and added to the end of the data matrix. Relative clade support was estimated using 1,000 bootstrap replicates in PAUP* via "full heuristic" searches and simple taxon addition. The consistency index (CI) and retention index (RI) were used to assess the amount of homoplasy present in the data. The best-fitting substitution model (GTR+I+G) was determined under the Akaike Information Criterion (AIC) in Model selected [24]. BI was performed in MrBayes ver. 3.1.2 [25]. A 50% majority-rule consensus tree with Bayesian posterior probabilities (PPs) of clades was calculated after removing the first 10% generations as burn in.

RESULTS AND DISCUSSION

The data set of the ITS region included 611 characters, 224 of them potentially parsimony informative. Strict consensus phylogeny trees, with 1014 steps was included consistency index (CI)=0.4872, retention index (RI)=0.8971 and homoplasy index (HI)= 1.83. Within the *Antirrhineae* four major clades were identified which have been given the name of one typical genus: the *Antirrhinum* clade (2 genera; PP = 0.99; BS = 60%), the *Maurandya* clade (2 genera; PP = 0.99; BS = 90%), the *Kickxia* clade (2 genera; PP = 1; BS = 100%), and all sampled species of *Linaria* were formed *Linaria* clade (PP = 0.99; BS = 100%). A little supported relationships between major clades were existed. *Linaria* clade included five major clades (A–E). Clade A included two species (*L. chalepensis* and *L. armenica* from Iran) of sect. *Macrocentrum* (PP = 1; BS = 100%). Clade B included one Iranian species (*L. iranica*) and two other species of sect. *Versicolores* (PP = 0.99; BS = 100%). *Pelisserianae* constituted clade C (PP = 0.69; BS = 71%). *L. triornithophora* is sister to clade B. This is not recognized in Iran. All sampled species of sect. *Supinae* (*L. arvensis*, *L. simplex*, *L. kavirensis* and *L. micrantha* from Iran) and three species of sect. *Diffusae* were formed Clade D (PP = 0.99; BS = 79%). Clade E (PP = 0.98; BS = 60%) was formed by all sampled species of sect. *Linaria* (*L. vulgaris*, *L. odora*, *L. khorasanensis*, *L. striatella*, *L. farsensis*, *L. bamianica*, *L. michauxii*, *L. pyramidalis subsp. pyramidalis*, *L. pyramidalis subsp. kopetdaghensis*, *L. fastigiata*, *L. remotiflora*, *L. nurensis*, *L. kurdica subsp. kurdica*, *L. kurdica subsp. pycnophylla*, *L. lineolata*, *L. karajensis*, *L. elymatica*, *L. azerbaijanensis*, *L. khalkhalensis*, *L. shahroudensis* and *L. guilanensis* from Iran), all sampled species of sect. *Speciosae* (*L. mazandaranensis*, *L. genistifolia*, *L. golestanensis*, *L. dalmatica*, *L. grandiflora* and *L. orientalis* from Iran), and six species of sect. *Diffusae* (*L. bousherensis1*, *L. bousherensis2* and *L. albifrons* from Iran). A little supported relationships among main clades were existed. Best support was occurred for two sister-group relationship between clades D and E (PP = 0.98; BS = 100%). Relationship between two sister-group clades B and C was supported by BI (PP = 0.69; BS = 71%). On the other side, relationships among other clades remained indefinite in our results.

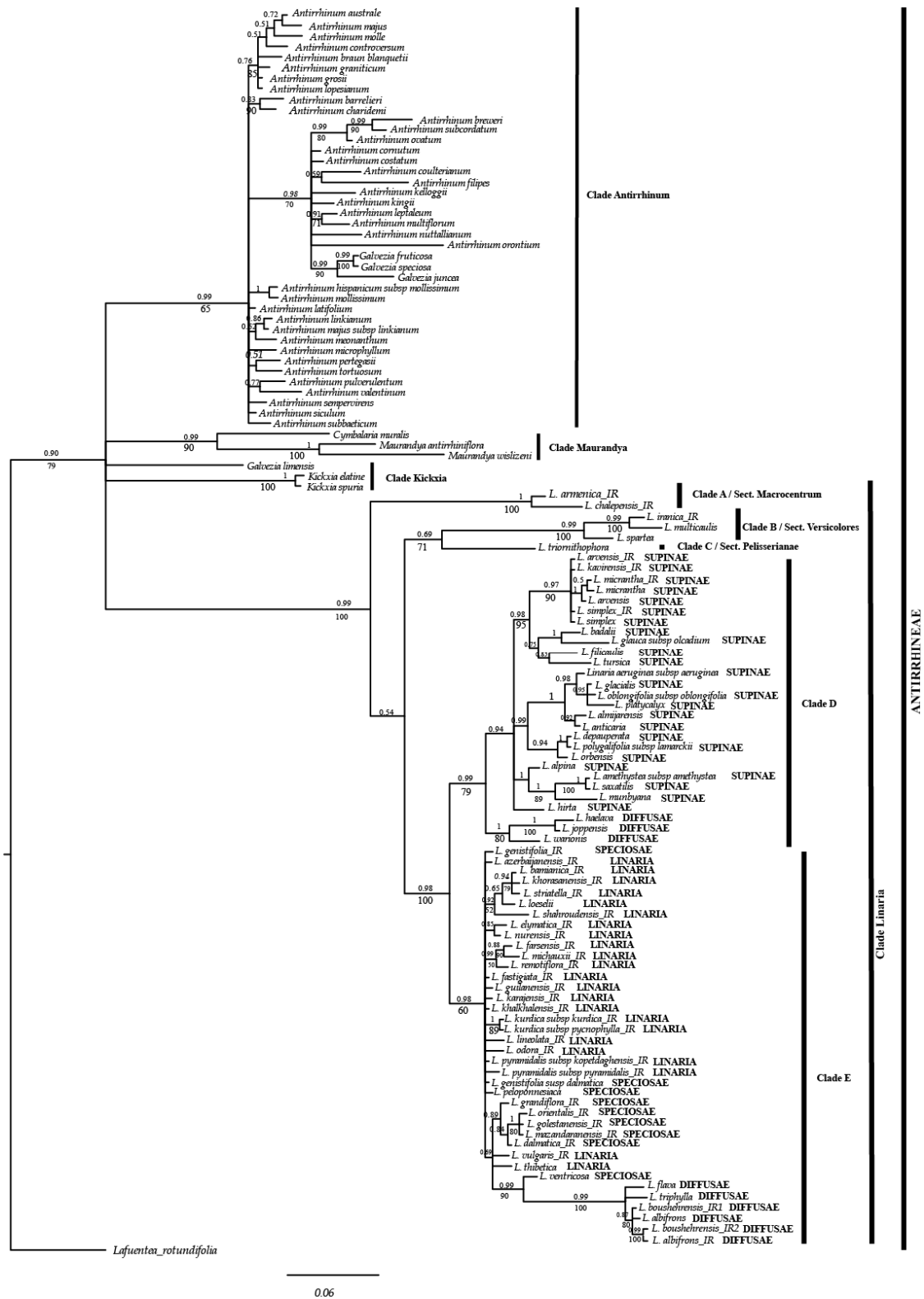
Table 3. List of non-Iranian taxa used in our analysis with GenBank accession number

Species	Genbank accession numbers
<i>Antirrhinum_barrelieri</i>	AY880240
<i>Antirrhinum_australe</i>	EU677195
<i>Antirrhinum_braun_blanquetii</i>	EU677196
<i>Antirrhinum_breweri</i>	AY880229
<i>Antirrhinum_charidemi</i>	EU677199
<i>Antirrhinum_controversum</i>	EU677206
<i>Antirrhinum_cornutum</i>	AY880230
<i>Antirrhinum_costatum</i>	AF513893
<i>Antirrhinum_coulterianum</i>	AF513890
<i>Antirrhinum_filipes</i>	AF513896
<i>Antirrhinum_graniticum</i>	EU677209
<i>Antirrhinum_grosii</i>	EU677210
<i>Antirrhinum_hispanicum_mollissimum</i>	EU677224
<i>Antirrhinum_kelloggii</i>	AF513904
<i>Antirrhinum_kingii</i>	AF513903
<i>Antirrhinum_latifolium</i>	EU677212
<i>Antirrhinum_leptaleum</i>	AF513906
<i>Antirrhinum_linkianum</i>	AY731278
<i>Antirrhinum_lopesianum</i>	EU677217
<i>Antirrhinum_majus_linkianum</i>	EU677215
<i>Antirrhinum_majus</i>	AF513888
<i>Antirrhinum_meonanthum</i>	AF513887
<i>Antirrhinum_microphyllum</i>	EU677221
<i>Antirrhinum_molle</i>	AY731268
<i>Antirrhinum_mollissimum</i>	AF513886
<i>Antirrhinum_multiflorum</i>	AY880228
<i>Antirrhinum_nuttallianum</i>	AF513895
<i>Antirrhinum_orontium</i>	AF513889
<i>Antirrhinum_ovatum</i>	AY880233
<i>Antirrhinum_pertegasii</i>	EU677228
<i>Antirrhinum_pulverulentum</i>	EU677230
<i>Antirrhinum_sempervirens</i>	EU677236
<i>Antirrhinum_siculum</i>	AY731276
<i>Antirrhinum_subbaeticum</i>	AY731287
<i>Antirrhinum_subcordatum</i>	AF513902
<i>Antirrhinum_tortuosum</i>	AY731285
<i>Antirrhinum_valentinum</i>	EU677243
<i>Galvezia_fruticosa</i>	AY731252
<i>Galvezia_junceae</i>	AY880243
<i>Galvezia_limensis</i>	AY492104
<i>Galvezia_speciosa</i>	AY316309
<i>Cymbalaria_muralis</i>	AY492101
<i>Kickxia_elatine</i>	AY731265
<i>Kickxia_spuria</i>	AF513880
<i>Maurandya_antirrhiniiflora</i>	AY492110
<i>Maurandya_wislizeni</i>	AY878930
<i>Lafuentea_rotundifolia</i>	AF509816
<i>Linaria_triornithophora</i>	AY731248
<i>Linaria_alpina</i>	AY731243
<i>Linaria_hirta</i>	AY731244
<i>Linaria_micrantha</i>	AY731242
<i>Linaria_aeruginea_aeruginea</i>	JQ814486
<i>Linaria_almijarensis</i>	JQ814492
<i>Linaria_amethystea_amethystea</i>	JQ814490
<i>Linaria_anticaria</i>	JQ814491
<i>Linaria_arvensis</i>	JQ814493
<i>Linaria_badalii</i>	JQ814495
<i>Linaria_depauperata</i>	JQ814499
<i>Linaria_filicaulis</i>	JQ814500
<i>Linaria_glacialis</i>	JQ814505
<i>Linaria_glauca_olcadium</i>	JQ814506
<i>Linaria_haelava</i>	JQ814507
<i>Linaria_joppensis</i>	JQ814508

<i>Linaria_loeselii</i>	JQ814511
<i>Linaria_multicaulis</i>	JQ814514
<i>Linaria_munbyana</i>	JQ814515
<i>Linaria_oblongifolia_subsp_oblongifolia</i>	JQ814516
<i>Linaria_orbensis</i>	JQ814518
<i>Linaria_platycalyx</i>	JQ814520
<i>Linaria_polygalifolia_lamarckii</i>	JQ814522
<i>Linaria_saxatilis</i>	JQ814526
<i>Linaria_simplex</i>	JQ814528
<i>Linaria_sparteae</i>	JQ814529
<i>Linaria_thibetica</i>	JQ814531
<i>Linaria_triphylla</i>	JQ814532
<i>Linaria_albifrons</i>	JQ814488
<i>Linaria_flava</i>	JQ814501
<i>Linaria_tursica</i>	JQ814533
<i>Linaria_ventricosa</i>	JQ814534
<i>Linaria_warionis</i>	JQ814536
<i>Linaria_genistifolia_dalmatica</i>	JQ814498
<i>Linaria_peloponnesiaca</i>	JQ814519

In this study we provide the first phylogenetic analysis about *Linaria* from Iran. At first the monophyly of *Linaria* on the eight sampled species exponent the seven sections of Sutton's classification, proposed by Vargas et al. [31]. Also in recent phylogenetic study based on ITS sequences by Fernánde z et al. (2013), Monophyly of *Linaria* was supported [6]. This analysis include 37 sampels of *Linaria* from Iran. Phylogenetic relationships between *Linaria* species in this study are according to those obtained by Vargas et al. (2004) and Fernánde z et al. (2013) based on the ITS sequences [6,31]. In this ITS phylogeny study, our species of *Linaria* organized a intensely supported monophyletic group (fig.1). morphological characters that are not ocured elsewhere in the Antirrhineae tribe and basic chromosome number ($x=6$) affirmed the monophyly of *Linaria*. [31, 28]. According to our obtained data the genus *Linaria* was determined as a paraphyletic group (fig. 1). This results as Fernánde z et al. (2013) finding, Viano's theory, separation of natural sister lineages in *Linaria* species based on with wingless and winged seeds forms, were refused. [32, 33]. Our analysis represent Sections *Macrocentrum* and *Versicolores* to be monophyletic (fig. 1). Other sections including *Linaria*, *Speciosae*, *Supinae* and *Diffusae* do not constitute as monophyletic groups. Species of sect. *Macrocentrum* (*L. chalepensis* and *L. armeniaca*, clade A; fig. 1) describe with some characters including the adaxial lobe of the calyx is shorter than abaxial lobes and lateral appendage present at the base of each stamen filament is small. Sutton (1980), based on these characters are specific between *Linaria*, separated them from sect. *Versicolores*. [27]. Section *Versicolores* is including just one species in Iran (*L. iranica*). Sutton (1988) was defined morphological character for sect. *Versicolores* that not found elsewhere in the Antirrhineae tribe [27]. This character was including separated style with distinct stigmatic region. The species *L. iranica*, that is endemic of Iran, has a bifid style. Champagnat (1961) With specific pattern of seedling progression is affirmed the naturalness of sect. *Versicolores*[4]. Four sections of *Linaria* including *Speciosae*, *Diffusae*, *Supinea* and *Linaria*, conversely sects. *Macrocentrum*, *Pelisserianae* and *Versicolores*, were not determined as monophyletic in this study. Relationships between species of sects. *Speciosae* and *Linaria* were showed in clade E. Previously Sutton (1988) was shown a identical morphological relationship among species of sections *Linaria* and *Speciosae* based on the perennial duration, erect stems, alike leaf, flower, and capsule morphology [28]. But there is a remarkable difference in seed form among species of both sections, also sect. *Linaria* have winged seeds and sect. *Speciosae* have wingless seeds. Three species of sect. *Diffusae* were also included in clade D (*L. haelava*, *L. joppensis* and *L. warionis*). This species formed a monophyletic group within clade D. Clade D also included species of sect. *Supinae* (*L. arvensis*, *L. simplex*, *L. kavirensis* and *L. micrantha* from Iran and other species of sect. *Supinae*). This results was supported the monophyly of species of sect. *Supinae*. The naturalness of sect. *Supinae* was affirmed union of both plastid and nuclear sequences in a merging-based analysis permitted the discovery of hybridization and imperfect lineage grouping [2]. Another species of sect. *Diffusae*, including *L. bousherensis1*, *L. bousherensis2*, *L. albifrons_IR*, *L. albifrons*, *L. flava*, and *L. triphylla*, constituted a monophyletic group within clade E. The main traits species of sect. *Speciosae* and sect. *Diffusae* including wingless seeds, entire style, normal adaxial lobe of calyx. Actually, separated species from both of sections is difficult. But chiefly morphological differences between species of sect. *Diffusae* and sect. *Speciosae* including annuals duration with ascending stems in sect. *Diffusae* and perennials duration with erect stems in sect. *Speciosae*. At first Valde's (1970) was proposed that sect. *Diffusae* was apparently polyphyletic, But Sutton (1988) believed that it was constituted a heterogeneous group [28]. Because of low phylogenetic resolution within clade E, we required extra markers for expose the phylogenetic relationships.

Fig. 1 Phylogenetic relationships of 37 samples of *Linaria* from Iran, 34 species of *Linaria*, 37 species of *Antirrhinum*, 4 species of *Galvezia*, 2 species of *Maurandya* and 2 species of *Kickxia* on the basis of the analysis of internal transcribed spacer sequences. Numbers above branches are Bayesian posterior probabilities. Numbers below branches are maximum likelihood percentage bootstrap values



CONCLUSION

The results indicate that *Linaria* species from Iran constituted a monophyletic group within the Antirrhineae. data analysis indicates that the classification of species according winged or wingless seeds so not true, so seed morphology especially seed wing that have been considered as useful for the taxonomy of *Linaria* species, appears to be a homoplasious character in *Linaria*. Two sections of Macrocentrum and Versicolores based on conformity special morphological traits and ITS sequences organized separated developmental lineages, therefor should be retained in classification of *Linaria*. But, naturalness of sections Supinae, *Linaria*, *Speciosae*, and *Diffusae* were not supported. For carefully determined of *Linaria* location we need to more molecular markers and more *Linaria* sampling from all sections.

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