

# Karyological Studies of Six Endemic Species of *Stachys* (Lamiaceae) Subsect. *Fragiles* from Turkey

Esra Martin<sup>1</sup>, Fahim Altınordu<sup>2</sup>, Özal Güner<sup>3\*</sup> and Ekrem Akçiçek<sup>3</sup>

<sup>1</sup>Department of Biotechnology Science Faculty, Necmettin Erbakan University, Konya, Turkey

<sup>2</sup>Department of Biology Science Faculty, Selçuk University, Konya, Turkey

<sup>3</sup>Department of Biology Education, Necatibey Education Faculty Balıkesir University, Balıkesir, Turkey

Received April 30, 2015; accepted April 14, 2016

**Summary** Several chromosome numbers are reported for *Stachys* taxa but little information is available about chromosome morphology. Thus, this study aims to investigate the karyotypes of six endemic *Stachys* species from Turkey. Two different chromosome numbers are reported as  $2n=30$  (*Stachys chamosericea* and *S. pinardii*) and  $2n=34$  (*S. euadenia*, *S. buttleri*, *S. pseudopinardii* and *S. longiflora*). The chromosome types of the studied taxa are metacentric and submetacentric. Karyological features and asymmetry indices ( $M_{CA}$ ,  $CV_{CI}$  and  $CV_{CL}$ ) of the six endemic species are identified for the first time, and karyological relationships are determined. Also we presented published chromosome data of *Stachys* along with new counts. These data can be used for chromosome number evolution for the genus and helpful for understanding its evolution.

**Key words** Chromosome, Comparative karyology, *Fragiles*, *Stachys*.

The genus *Stachys* L., one of largest genera in the Lamiaceae, comprises about 300 species. The genus is distributed mainly in the warm temperate regions of the Mediterranean and southwestern Asia as well as North America, South America and southern Africa (Bhattacharjee 1980). It consists of annual and perennial herbs and subshrubs. A great number of *Stachys* species grow in various ecological conditions such as forests, rocky places, on limestone, banks of streams. In Turkey, *Stachys* was represented 115 taxa belonging to 15 sections and two subgenera. Of the 115 taxa, 54 (47%) are endemic to Turkey and are mostly eastern Mediterranean elements (Bhattacharjee 1982, Davis *et al.* 1988, Gemici and Leblebici 1998, Duman 2000, İlçim *et al.* 2008, Dirmenci *et al.* 2011, Akçiçek *et al.* 2012).

The section *Fragilicaulis* R. Bhattacharjee contains 30 taxa and is distributed only in Turkey, North Iraq and West Iran. All species are suffrutescent saxatile perennials and their stems are fragile at base. This section is distinguished from the other section of *Stachys* by this feature. In Turkey, the section has 21 taxa. Endemism rate of this section are 71.4%. A majority of these species are concentrated in Irano-Turanian phytogeographical region (Bhattacharjee 1982, Rechinger 1982, Davis *et al.* 1988, Duman 2000).

Chromosome analysis of the sect. *Eriostomum* was studied in Turkey by Martin *et al.* (2011) and somatic chromosome numbers of these species were reported as  $2n=30$ . Besides, other karyological research conducted

on taxa of the genus *Stachys* showed that chromosome numbers were found to be from  $2n=10$  to  $2n=102$  (Pogán *et al.* 1980, Van Loon and Kieft 1980, Bhattacharjee 1982, Strid and Franzen 1983, Papanicolaou 1984, Mulligan and Munro 1989, Baden 1991, Baltisberger 1991a, b, 2006, Falciani and Fiorini 1996, Carr 1998, Wagner *et al.* 1999, Weller and Sakai 1999, Baltisberger and Widmer 2004) (Appendix 1). Generally, chromosome numbers  $2n=30$ , 34 and 66. Our reports are congruent with the literature.

Recently, there are some studies on the taxonomy, palynology, karyological, anatomy and micromorphology of *Stachys* (Dirmenci *et al.* 2011, Salmaki *et al.* 2011, Martin *et al.* 2011, Akçiçek *et al.* 2012, Satıl *et al.* 2012, Salmaki *et al.* 2012, Dündar *et al.* 2013). There has been no investigation of karyotype analysis of these endemic species yet. With this study, somatic chromosome numbers of subsect. *Fragiles* were identified. These species are *S. longiflora* Boiss. and Bal., *S. euadenia* P.H. Davis, *S. pinardii* Boiss., *S. buttleri* R.Mill., *S. pseudopinardii* R. Bhattacharjee & Hub.-Mor. and *S. chamosericea* Ayaşlıgil & P.H. Davis. Also we determined the karyotype asymmetry indices, statistically correct parameters such as  $M_{CA}$ ,  $CV_{CI}$  and  $CV_{CL}$ , and demonstrated the karyologic relationships among *Stachys* taxa. We performed an extensive literature review to show all published chromosome data of *Stachys* using *The Chromosome Counts Database* (CCDB) (Rice *et al.* 2014) and internet resources (*e.g.*, ISI Web of Science, Google Scholar).

\*Corresponding author, e-mail: ozalgnr57@hotmail.com

DOI: 10.1508/cytologia.81.231

## Materials and methods

*Stachys* specimens were collected from Antalya, Karaman and Mersin provinces between 2012 and 2014 in Turkey. Localities of the collected specimens were presented (Table 1).

For chromosome preparation and karyological observations, we considered Martin *et al.* (2011) and Altınordu *et al.* (2014) respectively. Chromosomes were classified using the nomenclature of Levan *et al.* (1964) as median (m), submedian (sm), subterminal (st) and terminal point (T). For karyotype asymmetry, we measured  $CV_{CI}$  (Coefficient of Variation of Centromeric Index),  $CV_{CL}$  (Coefficient of Variation of Chromosome Length) (Paszko 2006) and  $M_{CA}$  (Mean Centromeric Asymmetry) (Peruzzi and Eroğlu 2013).

A new approach, proposed by Peruzzi and Altınordu (2014), was used to reconstruct the karyological relationships among *Stachys* taxa. To perform cluster analysis, a similarity matrix was created using Gower's (1971) general coefficient similarity to summarize the relationships among taxa (Sneath and Sokal 1973) in Past 3.03 software (Hammer *et al.* 2001, Hammer 2013).

## Results

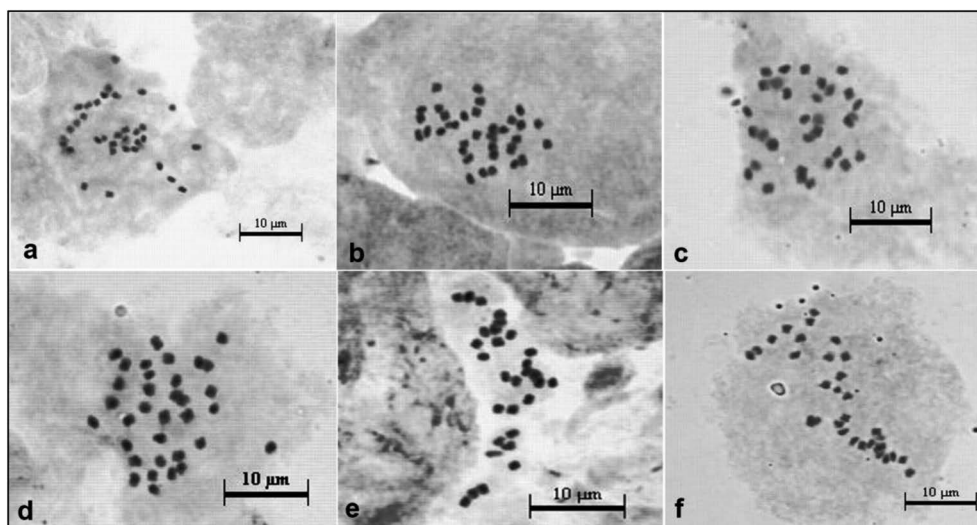
We report here on somatic chromosome numbers of five species and karyological features of six species first. *Stachys chamosericea* and *S. pinardii* are  $2n=30$ , *S. euadenia*, *S. butleri*, *S. pseudopinardii* and *S. longiflora* are  $2n=34$  (Fig. 1). *S. butleri* has the shortest chromosome with a length of  $0.90\ \mu\text{m}$ , whereas *S. pinardii* has the longest chromosome with a length of  $2.12\ \mu\text{m}$ . Length of the largest haploid chromosome was observed ( $22.3\ \mu\text{m}$ ) in *S. pinardii*. Length of the shortest haploid chromosome was observed ( $19.01\ \mu\text{m}$ ) in *S. longiflora*.

## Discussion

In plants, karyotype asymmetry is good evidence for karyotype morphology (Zarco 1986) and one of the most popular, inexpensive and widely used approaches, especially by botanists (Peruzzi and Eroğlu 2013). Researchers have developed a variety of methods for analyzing karyotype asymmetry in chromosome sets. Stebbins (1971), Huziwara (1962), Arano (1963), Greilhuber and Speta (1976), Zarco (1986), Lavania and Srivastava (1992), Watanabe *et al.* (1999), and Paszko (2006)

**Table 1.** Localities of *Stachys* specimens.

Localities of taxa and Collector and voucher numbers	
<i>S. longiflora</i>	C5 Mersin: Işıktepe villages, Kızıl Dere (Güzel dere) limestone, mouth of cave, 200–250 m, $36^{\circ}52'560''\text{N}/34^{\circ}33'382''\text{E}$ , 13.07.2013, Ö.Güner 2364, Akçiçek & Dirmenci
<i>S. euadenia</i>	C4 Karaman: Ermenek, Hamitseydi Boğaz, Kirboğazi place, limestone crevices, $36^{\circ}30'202''\text{N}$ , $032^{\circ}47'219''\text{E}$ , 1700 m, 12.07.2013, Ö.Güner 2374, Akçiçek & Dirmenci
<i>S. pinardii</i>	C3 Antalya: Road of Feslikan, Geyik Bayırı, Küllien cave, limestone crevices, 230 m, $36^{\circ}52'657''\text{N}/30^{\circ}28'596''\text{E}$ , 20.04.2013, 24.05.2013, Ö.Güner 2325 & Akçiçek
<i>S. butleri</i>	C3 Antalya: Düden waterfall, damp rocks, $36^{\circ}57'49''\text{N}/30^{\circ}3'41''\text{E}$ , 88 m, 24.05.2013, Ö.Güner 2327 & Akçiçek
<i>S. pseudopinardii</i>	C4 Mersin: Silifke, Cennet cave, 0–50 m, $36^{\circ}27'08''\text{N}$ , $34^{\circ}06'22''\text{E}$ , 18.08.2013, Ö.Güner 2390 & Akçiçek
<i>S. chamosericea</i>	C3 Antalya: Manavgat, Beşkonak, east of Yarış, the upper region of Karadağ, Karamuşar (Karapınar), limestone crevices, $37^{\circ}08'807''\text{N}$ , $031^{\circ}15'525''\text{E}$ , 825 m, 11.07.2013, Ö.Güner 2371, E. Akçiçek & Dirmenci



**Fig. 1.** Metaphase plates of *Stachys*. a: *S. longiflora*; b: *S. euadenia*; c: *S. butleri*; d: *S. pseudopinardii*; e: *S. chamosericea*; f: *S. pinardii*.

have developed a variety of methods for analyzing the karyotype variation of a chromosome complement (Zuo and Yuan 2011). For this aim, we considered three of them ( $M_{CA}$ ,  $CV_{CI}$  and  $CV_{CL}$ ) and discarded TF%, AsK%, AsI%, Syi, A1, CG (for details and references see Peruzzi and Eroğlu 2013; Peruzzi and Altınordu 2014). Lowest  $CV_{CI}$  values indicate chromosome complements with the most homogeneous centromere position, observed in *S. euadenia*. On the other hand, the most heterogeneous centromere position occurs in *S. chamosericea*. Among species,  $CV_{CL}$  values range between 11.6 and 17.44, and *S. buttlerei* shows highest interchromosomal asymmetry with 17.44 value. As to intrachromosomal asymmetry ( $M_{CA}$ ), *S. buttlerei* has highest value as 22.34 and *S. euadenia* has lowest value as 14.63. Detailed karyotype asymmetry indices and variability of symmetry indices among species are given (Table 2). Also, variabilities of  $CV_{CL}$  and  $CV_{CI}$  values for each taxon are illustrated by the boxplots (Fig. 2).

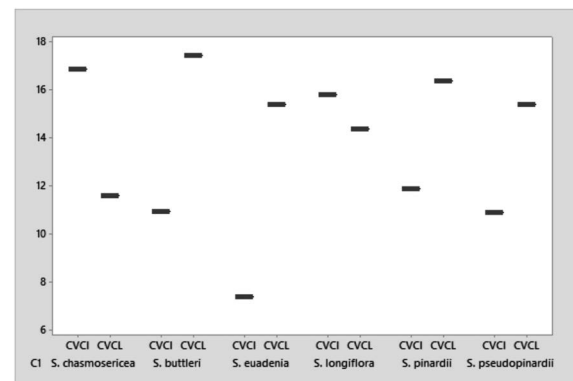
To reconstruct the karyological relationships among *Stachys* taxa, we used six quantitative features such as characters; chromosome number ( $2n$ ), basic chromosome number ( $x$ ), THL,  $M_{CA}$ ,  $CV_{CL}$  and  $CV_{CI}$  (Peruzzi and Altınordu 2014), and formed karyological tree according to these characters. Moreover, we superimposed the

idiograms onto this tree for clear relationships (Fig. 3).

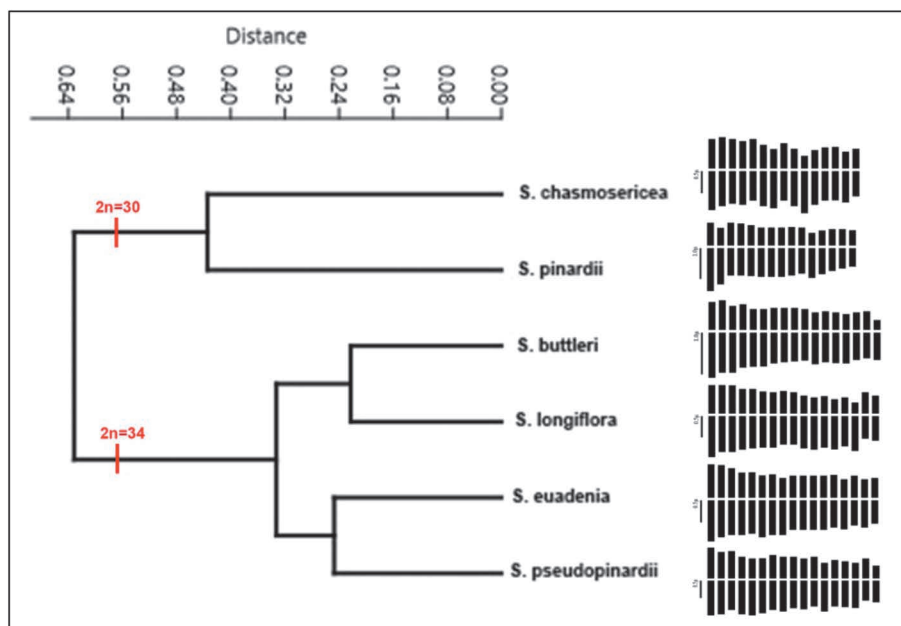
According to literature, there are several data on the basis of chromosome count for *Stachys* taxa (Appendix 1). However, chromosome morphologies could not be analysed because of their short lengths which were not suitable for karyotype analysis. As such, there is little data on the karyology of the genus *Stachys*. Khadivi-Khub and Aghaei (2014) investigated *Stachys lavandulifolia* Vahl from four Iranian populations. Chromosome number of this species reported as  $2n=60$ . The presence of 60 chromosomes in all populations corresponds to tetraploidy ( $2n=4x=60$ ) levels. The same taxa were investigated by Asghari Zakaria and Zare (2013) and chromosome numbers reported as  $2n=68$ , tetraploidy level. When Turkish and Iranian *Stachys* taxa were compared based on chromosome number, it was revealed that our studied taxa were diploid while Iranian taxa were tetraploid. When compared with their chromosome morphology, for each study, chromosome lengths varied from 1.30 to 1.48  $\mu\text{m}$  and 0.93 to 2.45  $\mu\text{m}$ , respectively. The lengths of chromosomes in the present study are

**Table 2.** Karyological features of studied taxa.

Taxa	$CV_{CI}$	$CV_{CL}$	$M_{CA}$	THL	$2n$	$x$
<i>Stachys longiflora</i>	15.79	14.38	21.51	19.01	34	17
<i>S. euadenia</i>	7.41	15.4	14.63	19.31	34	17
<i>S. buttlerei</i>	10.93	17.44	22.34	19.64	34	17
<i>S. pseudopinardii</i>	10.89	15.37	16.63	22.11	34	17
<i>S. chamosericea</i>	16.87	11.6	18.32	19.1	30	15
<i>S. pinardii</i>	11.87	16.38	20.53	22.3	30	15



**Fig. 2.** Variabilities of  $CV_{CL}$  and  $CV_{CI}$  values among taxa.



**Fig. 3.** Idiograms superimposed onto karyological tree.

congruent with these reports and vary between 0.90 and 2.12  $\mu\text{m}$ . Khadivi-Khub and Aghaei (2014) reported the  $CV_{CI}$  index between 1.22 and 3.44. In this study  $CV_{CI}$  index was found between 7.41 and 16.87. Lowest  $CV_{CI}$  values implies chromosome complements with the most homogeneous centromere position, so our studied species have more heterogeneous centromere positions than Iranian *Stachys*. Also, low  $CV_{CL}$  values imply low interchromosomal asymmetry. We found  $CV_{CL}$  values between 11.6 and 17.44. Khadivi-Khub and Aghaei (2014) reported the  $CV_{CL}$  index between 5.12 and 14.20. Thus, we can say that Iranian species is more symmetrical than Turkish species.

According to previous reports, the most common ploidy levels of *Stachys* species are diploid. Martin *et al.* (2011) studied chromosome numbers of 26 *Stachys* taxa from Turkey and reported the chromosome numbers of all taxa as  $2n=30$  and diploid (Appendix 1). Chehregani-Rad *et al.* (2012) investigated 13 populations of *Stachys inflata* Benth. from Iran and reported chromosome numbers of  $2n=2x=16$  and  $2n=4x=32$ , both diploid and tetraploid. Also, *S. aspera* Michx., *S. hispida* Pursh, *S. pilosa* Nutt. and *S. turcomanica* Trautv. were reported at tetraploid level (Mulligan and Munro 1989; Chuksanova and Kaplanbekova 1971). In the present study, six endemic *Stachys* were studied and reported as  $2n=30$  and 34 as diploid.

In conclusion, our results showed that the basic chromosome numbers of studied *Stachys* taxa are  $x=15$  and  $x=17$ . Chromosome numbers correspond to the diploid level and when compared with available studied taxa from other countries, the karyotype of Turkish taxa is more asymmetrical than others.

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## Appendix 1.

Taxa	2n	References	Taxa	2n	References
<i>Stachys aculeolata</i> Hook.	2n=46 and 52	Morton 1993.	<i>S. candida</i> Bory and Chaub.	2n=34	Baltisberger and Lenherr 1984.
<i>S. aegyptiaca</i> Pers.	2n=34	CCDB	<i>S. canescens</i> Bory and Chaub.	2n=34	Baltisberger and Lenherr 1984.
<i>S. affinis</i> Bunge	2n=16	Krestovskaya and Vassiljeva 1998.	<i>S. chamissonis</i> Benth.	2n=64	Mulligan and Munro 1989.
<i>S. agraria</i> Schldl. and Cham.	2n=32	Mulligan and Munro 1989.	<i>S. chinensis</i> Bunge ex Benth.	2n=c.80	Sokolovskaya <i>et al.</i> 1986.
<i>S. ajugoides</i> Benth.	2n=66	Mulligan and Munro 1989.	<i>S. chrysantha</i> Boiss. and Heldr.	2n=34	Baltisberger and Lenherr 1984.
<i>S. albanica</i> Markgr.	2n=34	Baltisberger and Lenherr 1984.	<i>S. circinata</i> L'Hér.	x=10	Ruiz de Clavijo 1991.
<i>S. albens</i> A. Gray	2n=66	Mulligan and Munro 1989.	<i>S. coccinea</i> Ortega	x=42	Spellenberg 1986.
<i>S. alopecuros</i> (L.) Benth.	2n=16	Baltisberger 1991.	<i>S. cordata</i> Riddell	2n=34	Mulligan and Munro 1989.
<i>S. alpina</i> L.	2n=30	Strid and Franzen 1983.	<i>S. corsica</i> Pers.	2n=18	Verlaque <i>et al.</i> 1992.
<i>S. angustifolia</i> M. Bieb.	2n=34	Sekovski and Jovanovska 1983.	<i>S. cretica</i> L.	2n=30	Baltisberger 1987.
<i>S. anisochila</i> Vis. and Pancic	2n=34	Baltisberger and Lenherr 1984.	<i>S. discolor</i> Benth.	2n=16	Gagnidze and Gviniashvili 1997.
<i>S. annua</i> (L.) L.	2n=34	Uhrliková and Schwarzová 1980.	<i>S. drummondii</i> Benth.	2n=c.80	Mulligan and Munro 1989.
<i>S. arvensis</i> (L.) L.	2n=10	Ruiz de Clavijo 1990.	<i>S. eplingii</i> J. B. Nelson	2n=34	Mulligan and Munro 1989.
<i>S. aspera</i> Michx.	2n=68	Mulligan and Munro 1989.	<i>S. euboica</i> Rech.	2n=34	Baltisberger and Lenherr 1984.
<i>S. atherocalyx</i> K. Koch	2n=34	Guinochet and Lefranc 1981.	<i>S. floccosa</i> Benth.	x=15	Gill 1984.
<i>S. balansae</i> Boiss. and Kotschy ssp. <i>balansae</i>	2n=30	Martin <i>et al.</i> 2011.	<i>S. floridana</i> Shuttlew. ex Benth.	2n=34	Mulligan and Munro 1989.
<i>S. bayburtensis</i> R. Bhattacharjee and Hub.-Mor.	2n=30	Martin <i>et al.</i> 2011.	<i>S. germanica</i> L.	2n=30	Pogan <i>et al.</i> 1980.
<i>S. beckeana</i> Dörf. and Hayek	2n=34	Baltisberger and Lenherr 1984.	<i>S. gilliesii</i> Benth.	x=16	De Fernandes and De Sarmiento 1973.
<i>S. bergii</i> G. A. Mulligan and D. B. Munro	2n=34	Mulligan and Munro 1989.	<i>S. glutinosa</i> L.	2n=34	Villa 1978.
<i>S. betoniciflora</i> Rupr.	2n=16	Astanova 1984.	<i>S. heraclea</i> All.	2n=30	Baltisberger 1988.
<i>S. bullata</i> Benth.	2n=66	Mulligan and Munro 1989.	<i>S. hyssopifolia</i> Michx.	2n=34	Mulligan and Munro 1989.
<i>S. byzantina</i> K. Koch	2n=30	Gill 1981.	<i>S. hissarica</i> Regel	2n=30	Astanova 1981.
<i>S. hispida</i> Pursh	2n=68	Mulligan and Munro 1989.	<i>S. mollissima</i> Willd.	2n=34	Baltisberger 1991.
<i>S. huber-morathii</i> R. Bhattacharjee	2n=30	Martin <i>et al.</i> 2011.	<i>S. spinosa</i> L.	x=17	De Montmollin, 1984.
<i>S. huetii</i> Boiss.	2n=30	Martin <i>et al.</i> 2011.	<i>S. spinulosa</i> Sm.	2n=18	CCDB
<i>S. inflata</i> Benth.	2n=32	Cartier 1983.	<i>S. spreitzenhoferi</i> Heldr.	2n=34	Baltisberger 2006.
<i>S. ionica</i> Halácsy	2n=34	Baltisberger and Lenherr 1984.	<i>S. spruneri</i> Boiss.	2n=34	Baltisberger and Lenherr 1984.
<i>S. iva</i> Griseb.	2n=34	Baltisberger and Lenherr 1984.	<i>S. stebbinsii</i> G. A. Mulligan and D. B. Munro	2n=66	Mulligan and Munro 1989.
<i>S. komarovii</i> Knorrning	2n=32	Gurzenkov 1973.	<i>S. stricta</i> Greene	2n=66	Mulligan and Munro 1989.
<i>S. latidens</i> Small	x=17	Mulligan and Munro 1989.	<i>S. swainsonii</i> Benth.	2n=34	Baltisberger, 2006.
<i>S. lavandulifolia</i> Vahl	2n=60	Chuksanova and Kaplanbekova 1971.	<i>S. tenuifolia</i> Willd.	2n=34	Mulligan and Munro 1989.
<i>S. leucoglossa</i> Griseb.	2n=34	Baltisberger and Lenherr 1984.	<i>S. tetragona</i> Boiss. and Heldr.	2n=34	Baltisberger and Lenherr 1984.
<i>S. libanotica</i> Benth. var. <i>minor</i> Boiss.	2n=30	Martin <i>et al.</i> 2011.	<i>S. thirkei</i> K. Koch	2n=30	Falciani and Fiorini 1996.
<i>S. macrantha</i> (K. Koch) Stearn	2n=32	Baltisberger 1989.	<i>S. thracica</i> Davidov	2n=30	Martin <i>et al.</i> 2011.
<i>S. macrostachys</i> (Wender.) Briq.	2n=16	Pogosyan 1974.	<i>S. tmolea</i> Boiss.	2n=30	Martin <i>et al.</i> 2011.
<i>S. maritima</i> Gouan	2n=34	Baltisberger 1991.	<i>S. tournefortii</i> Poir.	x=15	Montmollin 1982.
<i>S. marrubifolia</i> Viv.	2n=16	Contandriopoulos 1962.	<i>S. turcomanica</i> Trautv.	2n=60	Chuksanova and Kaplanbekova 1971.
<i>S. melissifolia</i> Benth.	x=15	Saggo 1983.	<i>S. tymphaea</i> Hausskn.	2n=30	Krestovskaya and Vassiljeva 1998.
<i>S. menthifolia</i> Vis.	2n=34	Baltisberger 1991.	<i>S. viticina</i> Boiss.	2n=30	Martin <i>et al.</i> 2011.
<i>S. menthoides</i> Kotschy and Boiss.	x=17	CCDB	<i>S. vuralii</i> Yıldız, Dirmenci and Akçiçek	2n=30	Martin <i>et al.</i> 2011.
<i>S. mexicana</i> Benth.	x=32	Gill 1981.			