

日本産アキギリ属の核形態学的比較

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Tsuneo Funamoto, Masatoshi Zushi, Tomoko Harana and Takuzo Nakamura : **Comparative karyomorphology of the Japanese species of *Salvia* L. (Lamiaceae)**

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Abstract

Ten species and one variety of *Salvia* in Japan had common karyomorphology of the complex chromocenter type in the resting stage, the proximal type in the mitotic prophase chromosomes, and the chromosome number of $2n=16$. The chromosome number of *S. koyamae*, *S. glabrescens*, *S. lutescens* f. *lobato-crenata*, *S. isensis*, *S. ranzaniana*, *S. pygmaea*, *S. omerocalyx* var. *omerocalyx* and *S. omerocalyx* var. *prostrata* were counted here for the first time, while that of *S. nipponica*, *S. plebeia* and *S. japonica* verified the previous reports. Two largest chromosomes of *S. plebeia* were median-centromeric and were different from those of the other ten taxa which were subterminal-centromeric. The two largest chromosomes in each of the ten taxa had commonly satellites on the short arm.

Key words : Japan, karyomorphology, Lamiaceae, *Salvia*.

Introduction

Genus *Salvia* L., which consists of over 900 species (Standley and Williams 1973), is the largest of the Lamiaceae (Labiales) and is found in subtropical and temperate regions of the world. Among the ten species of the genus distributed in Japan, eight species are endemic (Murata and Yamazaki 1993).

Chromosome studies have been made in about 210 non-Japanese species of the genus, and reported 35 different chromosome numbers of $2n=12$ up to $2n=72+2B$ as aneuploid and polyploid (e.g., Scheel 1931; Stewart 1939; Delestaing 1954; Epling et al. 1962; Gadella et al. 1966; Gill 1971, 1984; Chuksanova and Kaplanbekova 1971; Afzal-Rafii 1971, 1972, 1976, 1980; Patudin et al. 1975; Wu and Huang 1975; Magulaev 1976; Haque and Ghoshal 1980; Markova and Ivanova 1982; Rosúa and Blanca 1985, 1988; Palomino et al. 1986; Mercado et al. 1989; Cherian and Kuriachan 1990). In contrast, chromosome studies of only three species of Japanese *Salvia* have been made; $2n=16$ (Morinaga et al. 1929; Scheel 1931; Carlson and Stuart 1936) in *S. nipponica*, $n=16, 18$ (Wu and Huang 1975) and $2n=16$ (Choshi 1968) in

S. japonica, $n=8$ (Mehra and Gill 1968; Vij and Kashyap 1975, 1976; Bir et al. 1978; Saggoo and Bir 1983), $n=8+1B$ (Gill 1984), $2n=16$ (Bir and Sidhu 1980) and $2n=32$ (Ayyangar and Vembu 1984) in *S. plebeia*.

This paper reports on karyomorphology of the species of *Salvia* in Japan.

Materials and methods

Eighty plants of ten species and one variety of Japanese *Salvia* were collected in 24 localities (Table 1) and were cultivated in pots in the experimental garden of the Showa Pharmaceutical University. Karyomorphological observations were made in fresh root-tip cells. Root tips were pretreated in 2 mM 8-hydroxyquinoline for 4 h at ca. 20°C, fixed in 45% acetic-acid for 10 min at ca. 2°C, macerated in a mixture of 45% acetic-acid and 1 N hydrochloric-acid (1:1) for 20-23 sec at ca. 60°C, and then stained and squashed in 2% aceto-orcein.

Karyomorphological expression of the resting stage and the mitotic prophase chromosomes were classified according to Tanaka (1971, 1977) and the mitotic metaphase chromosomes by the centromeric position accepted by Levan et al.

Table 1. Localities, samples and chromosome numbers of Japanese *Salvia*

Species	Locality	Sample number	Chromosome number (2n)
<i>S. koyamae</i> Makino	Nagano Pref., Minamisaku-gun, Usuda-machi, Mizuochi	6	16
<i>S. nipponica</i> Miq.	Niigata Pref., Nishikanbara-gun, Yahiko-mura, Mt. Yahiko	3	16
	Yamanashi Pref., Nakakoma-gun, Ashiyasu-mura, Yashajin-toge	2	16
	Kumamoto Pref., Kikuchi City, Kikuchi-keikoku	4	16
<i>S. glabrescens</i> Makino	Fukui Pref., Katsuyama City, Iwaya	3	16
	Fukui Pref., Ono-gun, Izumi-mura, Shimohanbara	4	16
	Gifu Pref., Ibi-gun, Fujihashi-mura, Tokuyama	3	16
<i>S. plebeia</i> R. Br.	Gunma Pref., Annaka City, Yanase	4	16
<i>S. japonica</i> Thunb.	Chiba Pref., Awa-gun, Amatsukominato-machi, Mt. Kiyosumi	2	16
	Toyama Pref., Kaminiikawa-gun, Oyama-machi, Arimineko	2	16
	Gifu, Pref., Kaidu-gun, Nannou-cho, Ishizu	1	16
	Shizuoka Pref., Fujieda City, Nishikata	2	16
	Kyoto Pref., Funai-gun, Hiyoshi-cho, Kamisegi	2	16
	Okayama Pref., Maniwa-gun, Yatsuka-son, Basari-toge	3	16
<i>S. lutescens</i> (Koidz.) <i>Koidz. f. lobato-crenata</i> (Makino) Murata	Aichi Pref., Kitashitara-gun, Shitara-cho, Onakura	4	16
	Aichi Pref., Kitashitara-gun, Inabu-cho, Noiri	2	16
<i>S. isensis</i> Nakai ex H. Hara	Shizuoka Pref., Inasa-gun, Inasa-cho, Kurumegi	4	16
	Shizuoka Pref., Inasa-gun, Inasa-cho, Ihei	3	16
<i>S. ranzaniana</i> Makino	Mie Pref., Taki-gun, Miyagawa-mura, Mobarra	3	16
	Mie Pref., Owase City, Kuchisubo	4	16
<i>S. pygmaea</i> Matsum.	Okinawa Pref., Yaeyama-gun, Taketomi-cho, Urauchi	3	16
	Okinawa Pref., Yaeyama-gun, Taketomi-cho, Komi	8	16
<i>S. omerocalyx</i> Hayata var. <i>omerocalyx</i>	Tottori Pref., Yazu-gun, Wakasa-cho, Mt. Ogino-sen	5	16
<i>S. omerocalyx</i> Hayata var. <i>prostrata</i> Satake	Fukui Pref., Mikata-gun, Mikata-cho, Kitamaekawa	3	16

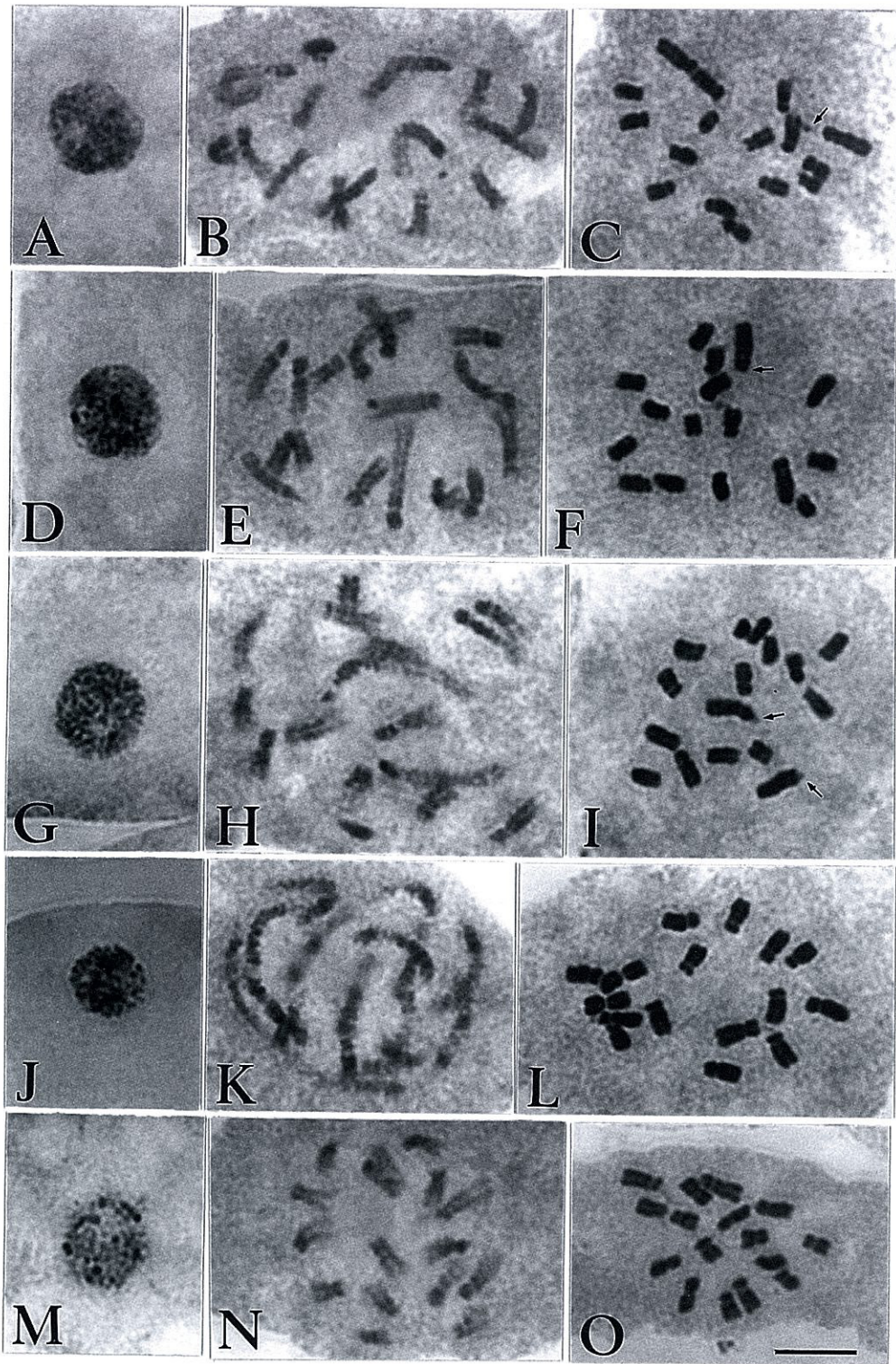


Fig. 1. Karyomorphology of Japanese species of *Salvia*.

A-C: *S. koyamae*; D-F: *S. nipponica*; G-I: *S. glabrescens*; J-L: *S. plebeia*; M-O: *S. japonica*. Nuclei with resting chromosomes (A, D, G, J and M). Mitotic prophase chromosomes (B, E, H, K and N). Mitotic metaphase chromosomes, $2n=16$ (C, F, I, L and O). Arrows represent satellites. Bar=5 μ m.

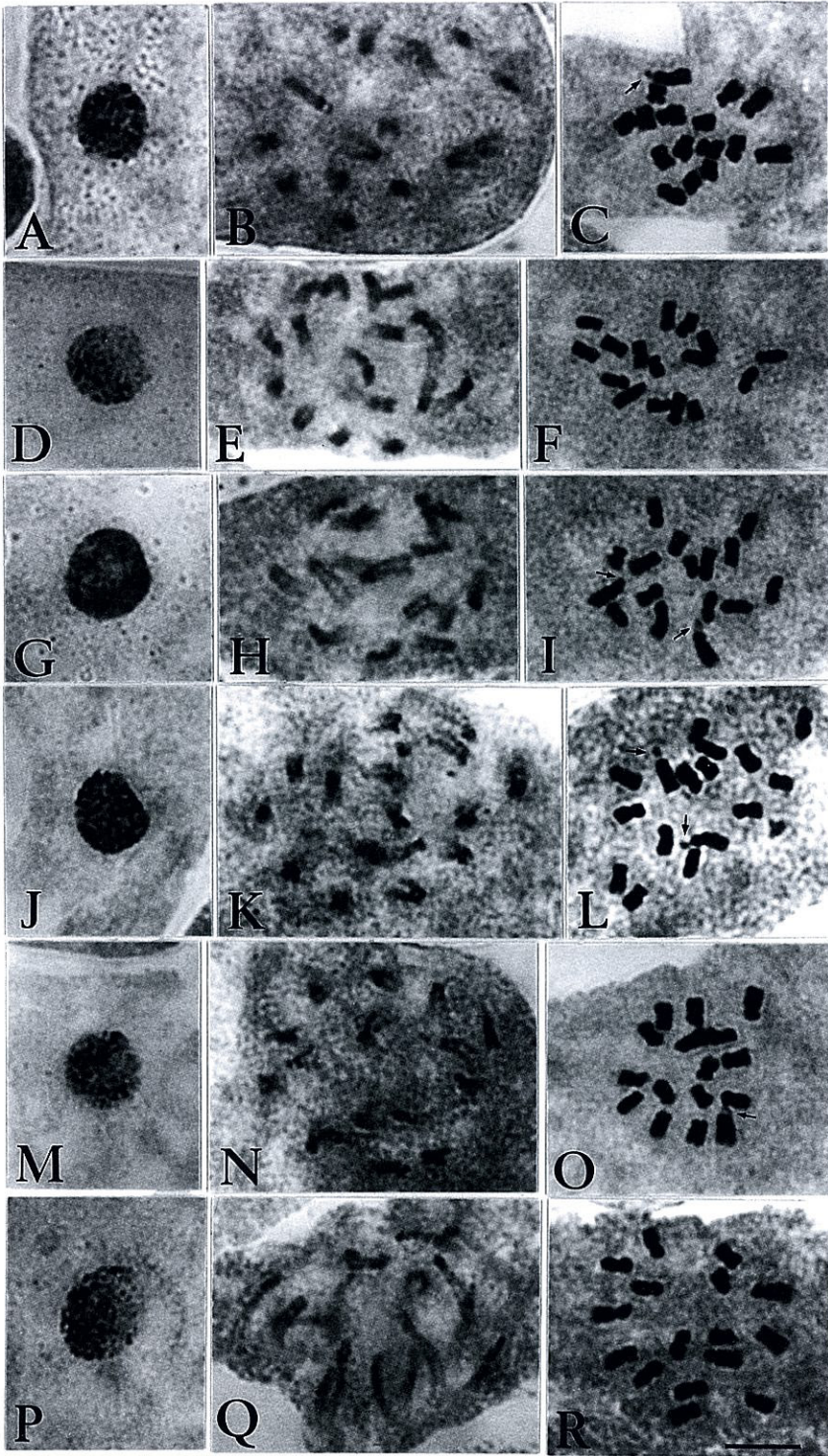


Fig. 2. Karyomorphology of Japanese species of *Salvia*.

A-C: *S. lutescens* f. *lobato-crenata*; D-F: *S. isensis*; G-I: *S. ranzaniana*; J-L: *S. pygmaea*; M-O: *S. omerocalyx* var. *omerocalyx*; P-R: *S. omerocalyx* var. *prostrata*. Nuclei with resting chromosomes (A, D, G, J, M and P). Mitotic prophase chromosomes (B, E, H, K, N and Q). Mitotic metaphase chromosomes, $2n=16$ (C, F, I, L, O and R). Arrows represent satellites. Bar=5 μ m.

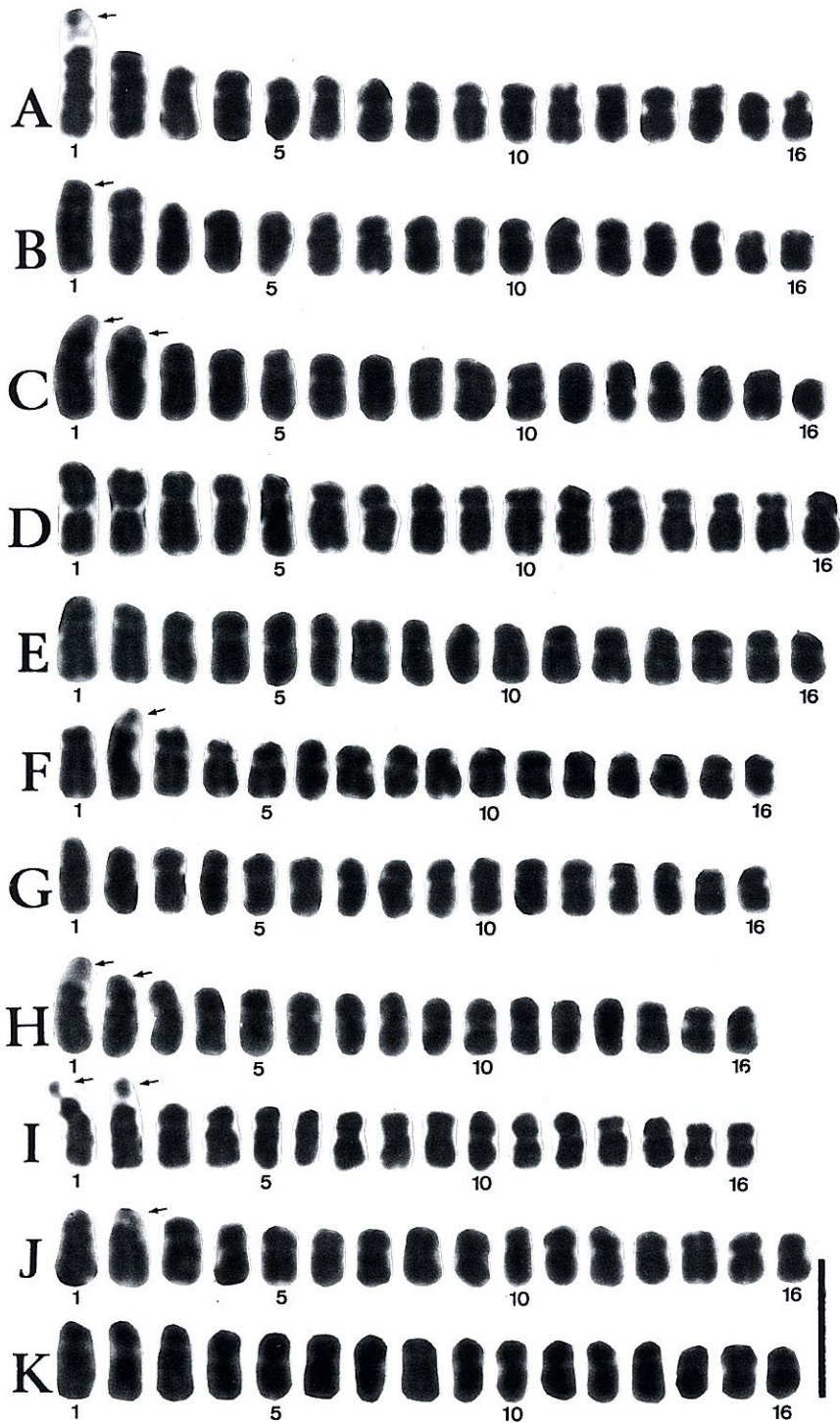


Fig. 3. Karyotypes of Japanese species of *Salvia*.

A: *S. koyamae*; B: *S. nipponica*; C: *S. glabrescens*; D: *S. plebeia*; E: *S. japonica*; F: *S. lutescens* f. *lobato-crenata*; G: *S. isensis*; H: *S. ranzianiana*; I: *S. pygmaea*; J: *S. omerocalyx* var. *omerocalyx*; K: *S. omerocalyx* var. *prostrata*. Arrows represent satellites. Bar=5 μ m.

(1964). Taxonomical treatment followed Murata and Yamazaki (1993). The voucher specimens were deposited in the Showa Pharmaceutical University.

Results and discussion

The ten species and one variety of *Salvia* in Japan investigated had in common the following karyomorphology; the complex chromocenter type of the resting stage which had many chromomeric granules and chromocentral small blocks scattered in the whole region (Fig. 1 A, D, G, J, M and Fig. 2 A, D, G, J, M, P) and the proximal type of the mitotic prophase chromosomes in which the early condensed segments were confined to the proximal part of the both short and long arms or only to the short arm (Fig. 1 B, E, H, K, N and Fig. 2 B, E, H, K, N, Q). The 11 taxa had commonly $2n=16$ chromosomes (Fig. 1 C, F, I, L, O and Fig. 2 C, F, I, L, O, R). Neither intraspecific polyploid nor aneuploid were observed in the taxa (Table 1). The somatic chromosome numbers of *S. koyamae* Makino, *S. glabrescens* Makino, *S. lutescens* (Koidz.) Koidz. f. *lobato-crenata* (Makino) Murata, *S. isensis* Nakai ex H. Hara, *S. ranzaniana* Makino, *S. pygmaea* Matsum., *S. omerocalyx* Hayata var. *omerocalyx* and *S. omerocalyx* Hayata var. *prostrata* Satake were reported here for the first time, while those of *S. nipponica* Miq., *S. plebeia* R. Br. and *S. japonica* Thunb. verified the previous reports (e.g., Morinaga et al. 1929; Choshi 1968; Mehra and Gill 1968; Bir and Sidhu 1980). These taxa of the genus had similar karyotypes; gradual decrease in chromosome length from the longest to the shortest chromosomes, and the chromosome complement of $2n=16$ consisted of median-, submedian- and subterminal-centromeric chromosomes. The two longest chromosomes of *S. plebeia* were median-centromeric (Fig. 3 D), while those of the other ten taxa were subterminal-centromeric. Satellite was observed basically in the short arm of the two longest chromosomes (Fig. 3 A-C, E-K). This difference in centromeric position was correlated with life span; annual or biennial life span in *S. plebeia* and perennial life span in the other ten taxa.

Variations of chromosome numbers in *Salvia*

were recognized in respective continent; polyploid and aneuploid with mainly $2n=12, 14, 16, 20, 22$ in *Salvia* in Europe and South, Central and South-east Asia continents (e.g., Gadella et al. 1966; Bothmer 1970; Gill 1971; Patudin et al. 1975; Haque and Ghoshal 1980; Markova and Ivanova 1982; Cherian and Kuriachan 1990), polyploid and aneuploid with $2n=12, 16, 20, 22$ in *Salvia* in North and Central American continents (Stewart 1939; Epling et al. 1962; Palomino et al. 1986; Mercado et al. 1989), polyploid and aneuploid with $2n=14, 16, 18, 20, 22$ in *Salvia* in Africa continent (e.g., Delestaing 1954; Hedge 1974). However, such variation of chromosome numbers was not well differentiated in Japanese species of *Salvia*. The Japanese species of *Salvia* may not be chromosomally diversified.

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船本常男・図師正敏・原名智子・中村卓造：日本産アキギリ属の核形態学的比較

日本産アキギリ属10種1変種について核形態の比較を行った。11 taxa はともに静止期は複雑染色中央粒型、分裂期前期染色体は基部型であった。分裂期中期においても染色体数は11 taxa とも $2n=16$ であり、倍数体、異数体は全く観察されなかった。シナノアキギリ、アキギリ、ナツノタムラソウ、シマジタムラソウ、ハルノタムラソウ、ヒメタムラソウ、タジマタムラソウ、ハイタムラソウの染色体数は今回始めて観察され、キバナアキギリ、ミゾコウジュ、アキノタムラソウの染色体数は今までの報告と一致した。11 taxa とも染色体は最大染色体から最小染色体まで勾配的な長さの変化を示し、中部型、次中部型及び次端部型染色体で構成されていた。唯一の違いは最大染色体対がミゾコウジュは中部型、他の10 taxa は次端部型で、短腕部に付随体が観察された。

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