Phylogenetic relationships among *Roegneria amurensis*, *R. ciliaris* and *R. japonensis* (Poaceae: Triticeae)

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To evaluate the phylogenetic relationships among *Roegneria amurensis*, *R. ciliaris* and *R. japonensis*, hybridizations among them were carried out. In the hybrids of *R. amurensis* \times *R. ciliaris*, *R. japonensis* \times *R. amurensis* and *R. ciliaris* \times *R. japonensis*, average chromosome associations were 0.38 I + 13.83 II, 0.18 I + 13.90 II and 0.15 I + 13.91 II, respectively. All the hybrids were partially fertile with stainable pollen grains and the fertility was 82%, 88% and 95%, respectively. The results indicated that the three species share the same basic genomes, StStYY and they are closely related. Combined with morphology and geographic distribution, *R. amurensis* and *R. japonensis* are treated as a subspecies and a variety of *R. ciliaris*, respectively.

Key words: chromosome pairing, genome, morphology, Poaceae, *Roegneria amurensis*, *R. ciliaris*, *R. japonensis*, Triticeae

Introduction

Roegneria is a very large and widely distributed genus in the tribe Triticeae (Poaceae). It includes about 120 species worldwide and nearly 80 species in China (Keng & Chen 1963, Kuo 1987, Baum et al. 1991, Cai 1997), and many subspecies or varieties are recognized. The morphological limits among some species are not clear, and as such the taxonomy within the genus is very difficult. Roegneria was once included in Elymus (Tzvelev 1976, Dewey 1984, Löve 1984), but many taxonomists consider it a valid genus (Keng & Chen 1963, Kuo 1987, Baum et al. 1991, Cai 1997, Zhou et al. 1999a).

Roegneria amurensis, R. ciliaris and R. japonensis are three tetraploid (2n = 4x = 28) taxa (Dewey 1984, Lu et al. 1988a, Wei et al. 2005). Both R. ciliaris and R. japonensis contain StStYY genomes (Dewey 1984, Lu et al. 1988b, Wang et al. 1994), while the genome constitution of R. amurensis is still unknown. Morphologically, R. amurensis differs slightly from R. ciliaris and R. japonensis in its pubescent leaves and leaf sheaths. Roegneria ciliaris differs slightly from R. japonensis in its pubescent glumes and erect spikes (Kuo 1987, Lu et al. 1988a). Cytologically, Wei et al. (2005) reported that the chromosomes of the three taxa are mainly metacentric and share similar karyotypes.

It is difficult to classify the three species due to the fact that their morphology differs only slightly. Tzvelev (1976) treated R. amurensis as Elymus ciliaris ssp. amurensis. Similarly, Löve (1984) treated R. amurensis and R. japonensis as subspecies of Elymus ciliaris. Based on cytological studies, Lu et al. (1988b) described R. japonensis as Roegneria ciliaris var. japonensis. Cai (1997) considered R. amurensis a valid species, and described R. japonensis as R. ciliaris var. hackeliana based on morphological comparison. Therefore, the taxonomic treatment and phylogenetic relationships of these three taxa are still in dispute. The main question is whether R. amurensis and R. japonensis are valid species, subspecies of R. ciliaris, or varieties of R. ciliaris.

The objectives of this study are: (1) to investigate the genomic constitution of *R. amurensis*; (2) to evaluate the phylogenetic relationships among *R. amurensis*, *R. ciliaris* and *R. japonensis* based on morphological comparison, meiotic study and fertility analysis; and (3) to provide a taxonomic treatment of the three taxa in *Roegneria*.

Material and methods

The material used in this study is listed in Table 1. *Roegneria amurensis* that originated in the Russian Federation was kindly provided by American National Plant Germplasm System (Pullman, Washington, USA). Other accessions were collected by the authors of this paper. All species are currently growing at Triticeae Research Institute, Sichuan Agricultural University, China (SAUTI).

The procedures of hybridization, fixation and staining of the cytological material and

meiotic preparation followed those reported by Zhou *et al.* (1999a). Chromosome pairing behavior at meiotic metaphase-I (MI) of parents and F_1 hybrids was observed in at least 50 pollen mother cells (PMCs). The count of mean pairing frequency (C = the mean frequency with which two related chromosome arms pair) was calculated according to Alonso and Kimber (1981). When mean C > 0.50, the two genomes in the hybrids are considered identical (Wang 1992). Microphotographs were taken from permanent meiosis preparations with an Olympus BX-51 camera. Pollen grains from mature anthers were stained in an aqueous I_2 -KI solution for pollen fertility studies.

Results

Intraspecific cross

The crosses R. amurensis $(Q) \times R$. ciliaris (Q), R. japonensis $(Q) \times R$. amurensis (Q) and R. ciliaris (Q) $\times R$. japonensis (O) produced hybrids without the aid of embryo rescue (Table 2). In R. amurensis \times R. ciliaris and R. japonensis \times R. amurensis, the seed sets were 22% and 20%, respectively. The seed set of R. ciliaris \times R. japonensis was 50%, thus much higher than that of the hybrids including R. amurensis (Table 2). Most of the seeds from each cross germinated easily and developed into vigorous hybrid plants. Six plants of the eleven seeds in R. amurensis × R. ciliaris, three plants of the nine seeds in R. japonensis \times R. amurensis and two plants of the two seeds in R. ciliaris \times R. japonensis were obtained. All the hybrid plants developed vigorously and set spikes later.

Table 1. Material used in this study.

Taxa	2n	Genomes	Origin	Accession no.
Roegneria amurensis	28	_	Russian Federation	PI 547303
Roegneria ciliaris	28	StY	Wugong, Shanxi, China Yaan, Sichuan, China	88-89-227 88-89-229
Roegneria japonensis	28	StY	Yibin, Sichuan, China Haerbin, Heilongjiang, China	88-89-253 88-89-257

Morphological comparisons among the parental species and the hybrids

Twenty-one morphological characters of the parents and their hybrids were measured (Table 3). Only few characters distinguished the three Roegneria taxa. The leaves and leaf sheaths of R. amurensis were pubescent, while they were glabrous in R. ciliaris and R. japonensis. The mean length of internodes is clearly longer and the number of florets per spikelet in R. amurensis is higher than in R. ciliaris and R. japonensis. Roegneria amurensis had smooth lemma tops, while R. ciliaris and R. japonensis had toothed ones. Roegneria ciliaris is readily distinguished from R. amurensis and R. japonensis by its longer top internodes. The general appearances of the F₁ hybrids were intermediate between their parents or near to one of the parental species. There was a tendency toward female parents in R. amurensis \times R. ciliaris and R. japonensis \times R. amurensis, especially in the number of spikelets per spike, number of florets per spikelet and length of spikelet (Table 3 and Fig. 1).

Meiosis in parents and hybrids

Chromosome pairing at MI of PMCs in the three parental species and their F_1 hybrids is listed in Table 4. The meiotic pairing of the parents was quite normal with 14 ring bivalents (Fig. 2A–C). Univalents were observed only occasionally and multivalent was not found in parents.

The patterns of chromosome pairing at MI were different in the three combinations. In the hybrid of R. amurensis \times R. ciliaris, the mean pairing configuration was 0.38 I + 13.83 II and the chiasma frequency was 24.99 with C = 0.89. In R. japonensis \times R. amurensis, the average

meiotic configuration was 0.18 I + 13.90 II and the chiasma frequency was 25.95 with C = 0.93. As for the hybrid of R. ciliaris $\times R$. japonensis, the mean pairing configuration was 0.15 I + 13.91 II and the chiasma frequency was 25.98 with C = 0.93. The frequencies of 14 bivalents observed in MI cells of the three combinations were 83%, 91% and 93%, respectively. In the three combinations, the lagging chromosomes were found at anaphase-I cells and micronuclei were also observed in the tetrads of the three crosses (Fig. 2H, I).

Fertility

The pollen fertilities and seed sets of the parental species and their hybrids were examined (Table 5). Most pollen grains of the parental species as well as their hybrids were viable. The average pollen fertilities of R. amurensis, R. ciliaris and R. japonensis were 89%, 95% and 92%, respectively. In the hybrids of R. amurensis \times R. ciliaris, R. japonensis \times R. amurensis and R. ciliaris \times R. japonensis, the mean pollen fertilities were 82%, 88% and 95%, respectively. The seed set of the parental plants was higher than that of their F₁ hybrids. Under open pollination, the respective average seed sets of R. amurensis, R. ciliaris and R. japonensis were 91%, 91% and 92%. In the three hybrids of R. amurensis \times R. ciliaris, R. japonensis \times R. amurensis and R. ciliaris \times R. japonensis, the mean opening seed sets were 53%, 64% and 84%, respectively.

Discussion

There is slight disparity among R. ciliaris, R. japonensis and R. amurensis in morphology.

Table 2. The cross combinations carried out in this study

Combinations ($Q \times \mathcal{O}$)	Number of emasculated	Se	ed set	P	Plants
,	florets	Number	Percentage	Number	Percentage
R. amurensis (PI 547303) × R. ciliaris (88-89-227)	50	11	22	6	54.55
R. japonensis (88-89-257) × R. amurensis (PI 54730	3) 45	9	20	3	33.33
R. ciliaris (88-89-229) × R. japonensis (88-89-253)	4	2	50	2	100.00

Table 3. Comparison of morphological characters in *Roegneria amurensis, R. ciliaris, R. japonensis* and their F₁ hybrids (cm, mean ± S.D)

Characters	R. amurensis	R. ciliaris (88-89-227)	R. japonensis (88-89-257)	R. amurensis×R. ciliaris	R. amurensis × R. ciliaris R. japonensis × R. amurensis	R. ciliaris× R. japonensis
Height	72.38 ± 5.33	77.86 ± 14.49	70.10 ± 4.47	67.24 ± 6.38	62.34 ± 8.39	71.40 ± 7.75
Length of flag leaf	19.32 ± 3.66	17.82 ± 3.15	14.86 ± 2.97	19.24 ± 3.21	16.27 ± 3.10	15.44 ± 2.03
Width of flag leaf	0.87 ± 0.11	1.14 ± 0.22	0.92 ± 0.08	1.02 ± 0.22	0.91 ± 0.09	0.87 ± 0.06
Length of top internode	37.58 ± 1.95	46.86 ± 6.36	32.88 ± 4.25	40.12 ± 5.37	33.28 ± 2.04	36.26 ± 3.25
Length of internode	3.42 ± 0.69	2.22 ± 0.38	1.82 ± 0.16	3.06 ± 0.42	2.26 ± 0.23	1.65 ± 0.12
Length of spike	20.36 ± 1.26	18.22 ± 4.10	16.44 ± 1.51	19.22 ± 2.13	18.43 ± 1.87	16.61 ± 4.46
No. of spikelets per spike	19.60 ± 2.97	13.60 ± 3.65	14.00 ± 1.22	19.40 ± 2.58	14.20 ± 1.24	15.80 ± 3.00
No. of florets per spikelet	12.00 ± 1.41	9.60 ± 1.14	8.60 ± 0.55	11.00 ± 2.18	9.80 ± 0.62	14.40 ± 0.50
Length of spikelet	4.06 ± 0.34	3.16 ± 0.36	3.24 ± 0.13	4.01 ± 0.22	3.28 ± 0.26	4.15 ± 0.15
Length of lower glume*	0.61 ± 0.05	0.74 ± 0.05	0.67 ± 0.03	0.65 ± 0.08	0.64 ± 0.05	0.70 ± 0.11
Length of upper glume*	0.70 ± 0.07	0.84 ± 0.05	0.78 ± 0.03	0.78 ± 0.05	0.76 ± 0.05	0.75 ± 0.12
Length of lemma**	0.86 ± 0.19	0.80 ± 0.26	0.84 ± 0.41	0.85 ± 0.21	0.90 ± 0.22	0.90 ± 0.01
Length of lemma awn	1.46 ± 0.22	1.42 ± 0.26	1.58 ± 0.39	1.41 ± 0.24	1.48 ± 0.22	1.48 ± 0.23
Length of palea	0.57 ± 0.06	0.68 ± 0.02	0.68 ± 0.03	0.64 ± 0.04	0.62 ± 0.03	0.67 ± 0.03
Lemma vestiture	rough	rough	glabrous	rough	rough	rough
Lemma top	smooth	toothed	toothed	toothed	toothed	toothed
Palea top	obtuse	optnse	obtuse	obtuse	obtuse	obtuse
Hairs on leaf	+++	I	I	I	I	I
Hairs on leaf sheath	++	I	1	I	I	I
Hairs on glume	+	+	I	+	+	+
Hairs on Iemma	+	+++	+	+	+	+

*including glume awn length, **excluding lemma awn length. + pubescent along margin, ++ pubescent on both surfaces and margin, +++ pubescent on both surfaces and margin, - hairless.

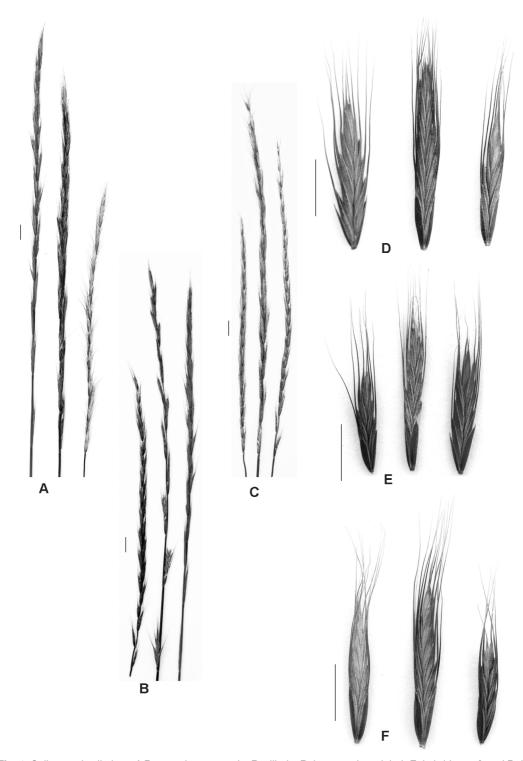


Fig. 1. Spikes and spikelets of *Roegneria amurensis*, *R. ciliaris*, *R. japonensis* and their F_1 hybrids. — **A** and **D**: Left *R. amurensis*, middle *R. amurensis* × *R. ciliaris*, right *R. ciliaris* (88-89-227). — **B** and **E**: Left *R. japonensis* (88-89-257), middle *R. japonensis* × *R. amurensis*, right *R. amurensis*. — **C** and **F**: Left *R. ciliaris* (88-89-229), middle *R. ciliaris* × *R. japonensis*, right *R. japonensis* (88-89-253). Bars = 1 cm.

Table 4. Chromosome pairing at MI of PMCs in the parental species of Roegneria and their F, hybrids

-		-	-	-				
Parents and hybrids	2n	No. of cells	_	Ch	Chromosome pairing II		Chiasma	O
		opserved		Total	Rod	Ring	ies Jed	
R. ciliaris (88-89-227)	28	89	0.03 (0–2)	13.99 (13–14)	0.60 (0–6)	13.39 (8–14)	27.38 (22–28)	0.98
R. amurensis (PI 547303)	28	106	0.08 (0-2)	13.85 (13–14)	1.27 (0–6)	12.58 (7–14)	26.43 (20–28)	0.94
R. japonensis (88-89-257)	28	136	0.07 (0-2)	13.97 (13–14)	1.10 (0-5)	12.87 (9–14)	26.84 (23–28)	96.0
R. amurensis × R. ciliaris	28	126	0.38 (0-4)	13.83 (12–14)	2.67 (0-5)	11.16 (9–14)	24.99 (23–28)	0.89
R. japonensis × R. amurensis	28	100	0.18 (0-2)	13.90 (13–14)	1.85 (0–5)	12.05 (9–14)	25.95 (23–28)	0.93
R. ciliaris × R. japonensis	28	123	0.15 (0-4)	13.91 (12–14)	1.84 (0–6)	12.07 (8–14)	25.98 (20–28)	0.93

Traditionally, they were distinguished from each other by three morphological characters: the hairs on glumes, the erect or slightly vertical spikes and the hairs on leaves and leaf sheaths. In fact, the variation of hairs on glumes is continuous and the erect spikes become vertical as the plants are growing up and seeding (Lu *et al.* 1988a).

The karyotype formula of R. ciliaris and R. japonensis was 2n = 4x = 28 = 18m (4SAT) + 6sm as shown by Lu et al. (1988a). Sun et al. (1992) suggested that the karyotype formula of R. ciliaris was 2n = 4x = 28 = 22m (2SAT) + 6sm (2SAT). Wei et al. (2005) reported that the karyotype formula of R. amurensis was 2n = 4x = 28 = 24m (2SAT) + 4sm and the karyotype formula slightly varied in different populations of R. ciliaris and R. japonensis. Therefore, the variations of karyotype within different populations of one species also made a sound taxonomy difficult.

The level of chromosome pairing in interspecific or intergeneric hybrids has played an important role in the study of biosystematic relationships among genera or species in Triticeae (Kihara & Nishiyama 1930, Sakamoto & Muramatsu 1966, Kimber 1983, Dewey 1984, Jauhar & Crane 1989, Lu et al. 1990, Wang 1992, Zhou et al. 1999a, 1999b). Lu et al. (1988b) reported that R. ciliaris and R. japonensis were closely related and had the same StStYY genomes. Combining morphological comparison with chromosome pairing study, Lu et al. (1988b) treated R. japonensis as Roegneria ciliaris var. japonensis.

In the present study, chromosome pairings in R. amurensis \times R. ciliaris, R. japonensis \times R. amurensis and R. ciliaris \times R. japonensis were quite high. The bivalents observed per cell varied between 12-14, 13-14 and 12-14 and the chiasma frequencies per cell were 23-28, 23-28 and 20-28, respectively. It indicated that R. amurensis contains a genome similar to that of R. ciliaris and R. japonensis, i.e. StStYY. The frequency of univalents is higher in both R. amurensis \times R. ciliaris and R. japonensis \times R. amurensis, while C, fertility and especially seed set are lower than in R. ciliaris \times R. japonensis (Tables 4 and 5). Therefore, it is concluded that the phylogenetic relationship between R. amurensis and R. ciliaris or R. japonensis is more

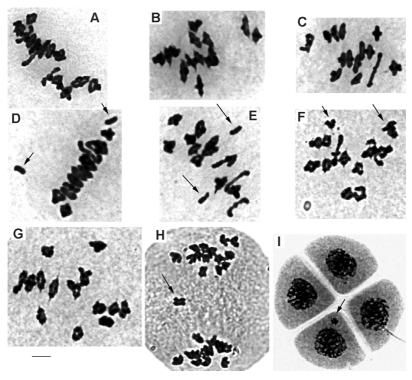


Fig. 2. Chromosome pairings at MI in three parental species of *Roegneria* and their F_1 hybrids. — **A**: *R. amurensis* with 14 bivalents (11 rings + 3 rods). — **B**: *R. ciliaris* (88-89-227) with 14 bivalents (14 rings). — **C**: *R. japonensis* (88-89-257) with 14 bivalents (12 rings + 2 rods). — **D**: *R. amurensis* × *R. ciliaris* with two univalents (indicated with arrows) and 13 bivalents (11 rings + 2 rods). — **E**: *R. japonensis* × *R. amurensis* with two univalents (indicated with arrows) and 13 bivalents (10 rings + 3 rods). — **F**: *R. ciliaris* × *R. japonensis* with two univalents (indicated with arrows) and 13 bivalents (11 rings + 2 rods). — **G**: *R. ciliaris* × *R. japonensis* with 14 bivalents (14 rings). — **H** and **I**: Lagging chromosomes at anaphase-I and micronuclei in tetrads; **H**: lagging chromosomes at anaphase-I of *R. japonensis* × *R. amurensis* (indicated with an arrow). **I**: a tetrad of *R. amurensis* × *R. ciliaris* showing micronuclei (indicated with an arrow). Bar = 5 µm.

distant than that between R. ciliaris and R. japonensis.

Roegneria amurensis is distributed in the northeastern parts of China and eastern parts of the Russian Federation, while *R. ciliaris* and *R. japonensis* are mainly growing widely in China

as well as in Japan and Korea. According to the results of morphological comparison, chromosome pairing study, pollen fertility and seed set in the hybrids including *R. amurensis*, it is reasonable to treat *R. amurensis* as a subspecies of *R. ciliaris* and *R. japonensis* as a variety of *R.*

Table 5. Pollen fertility and seed set in the parental species of *Roegneria* and their F, hybrids.

Parents and hybrids	No. of	No. of	Fertility (%)	No. of	Se	ed set
	pollen grains observed	fertile pollen grains		florets observed	Number	Percentage
R. ciliaris (88-89-227)	663	628	95	128	116	91
R. amurensis (PI 547303)	827	739	89	120	109	91
R. japonensis (88-89-257)	786	724	92	132	121	92
R. amurensis × R. ciliaris	1035	852	82	300	160	53
R. japonensis × R. amurensis	1100	973	88	280	180	64
R. ciliaris × R. japonensis	1287	1218	95	380	320	84

ciliaris. This combination of *R. amurensis* as a subspecies of *R. ciliaris* has not previously been made.

Roegneria ciliaris (Trin.) Nevski ssp. amurensis (Drob.) Y.H. Zhou, H.Q. Yu & X. H. Wei, comb. nov.

Agropyron amurensis Drob., Tr. Bot. Muz. AN 12: 50. 1914. — Roegneria amurensis (Drob.) Nevski, Fl. SSSR 2: 606. 1934. — Elymus ciliaris (Trin.) Tzvelev ssp. amurensis (Drob.) Tzvelev, Nov. Sist. Vyssch. Rast. 9: 61. 1972.

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