

THE KARYOTYPE OF THE AMU DARYA STURGEON, *PSEUDOSCAPHIRHYNCHUS KAUFMANNI* (ACTINOPTERYGII: ACIPENSERIFORMES: ACIPENSERIDAE)

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Background. Karyological studies of acipenserid fishes are of great importance because they present the only direct method to evaluate their ploidy levels for further research on polyploid evolution in these fishes. They are also important for prediction of the results of interspecific hybridizations in sturgeon aquaculture. None of the species of the genus *Pseudoscaphirhynchus* has hitherto been studied karyologically. The aim of this paper was to present the first data on the karyotype of the dwarf form of *Pseudoscaphirhynchus kaufmanni* (Kessler, 1877).

Materials and methods. Three females of the dwarf form of *Pseudoscaphirhynchus kaufmanni* of the total body length 19–23 cm were caught in the Vakhsh River (Amu Darya River drainage), Tadzhikistan, in 2012. The chromosome slides were prepared by using previously published karyological method of Vasil'ev and Sokolov. Totally, 14 metaphase plates were analyzed.

Results. The karyotype of *Pseudoscaphirhynchus kaufmanni* consists of 118–120 chromosomes and includes about 18–20 large bi-armed chromosomes, about 32–34 small bi-armed chromosomes, from one to two pairs of large acrocentrics, and about 64 small acrocentrics or microchromosomes.

Conclusion. The karyological study revealed that *Pseudoscaphirhynchus kaufmanni* belongs to low-chromosome acipenserid group with about 120 chromosomes. Its karyotype demonstrates noticeable differences from the karyotype of the shovelnose sturgeon, *Scaphirhynchus platorhynchus* (Rafinesque, 1820), in the number of large acrocentrics, thereby, karyological data confirms polyphyletic origin of the subfamily Scaphirhynchinae (or tribe Scaphirhynchini).

Keywords: acipenserids, karyology, ploidy levels, polyphyly, shovelnose sturgeons

INTRODUCTION

The order Acipenseriformes is an ancient group of vertebrate animals represented by two fossil families: Peipiaosteidae, extending back to the Upper Jurassic, and Chondrosteidae, dating from the Lower Jurassic, and extant representatives in two families: Acipenseridae and Polyodontidae (see Findeis 1997, Nelson 2006). Both extant families have fossil records with Mesozoic dating. Most primitive and the oldest paddlefishes (Polyodontidae) are known from the Lower Cretaceous of China (*Protopsephurus*) and North America (*Paleopsephurus*) (Grande and Bemis 1991); the only two living freshwater species represent two distinct phyloge-

netic lineages and genera (Hilton 2004): *Polyodon spathula* (Walbaum, 1792) in United States and *Psephurus gladius* (Martens, 1862) in China. Both the oldest sturgeon (Acipenseridae) species dated back to the Late Cretaceous in Montana are treated as taxa with an uncertain position, namely plesion *Protoscaphirhynchus* Wilimovsky, 1956, *insertae sedis*, and plesion *Priscosturion* Grande et Hilton, 2009, *sedis mutabilis* (see Hilton et al. 2011). The phylogenetic relations and taxonomic states of recent 25 sturgeon species also are the subject of a long-term discussion.

Usually, living Acipenserid fish are subdivided into two subfamilies. Some authors (Findeis 1997, Nelson 2006)

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recognize two species, namely *Huso huso* (Linnaeus, 1758) and *Huso dauricus* (Georgi, 1775), as a separate subfamily Husinae. (The recent taxonomic revision—Vasil'eva et al. 2009—proved, by using recent molecular genetic, karyological, and morphological evidences, that *Huso* Brandt et Ratzeburg, 1833 is not valid genus, but a junior synonym of *Acipenser* Linnaeus, 1758). The remaining species from the subfamily Acipenserinae they divide into two tribes—Acipenserini, and Scaphirhynchini—the latter tribe including genera *Scaphirhynchus* and *Pseudoscaphirhynchus* (see Grande and Bemis 1996). Another taxonomic hypothesis accepts Scaphirhynchinae as a separate subfamily opposed to all other Acipenserids (Berg 1905, Sokolov and Berdichevskii 1989). Both of these taxonomic concepts do not agree with recent molecular data (Ludwig et al. 2001, Dillman et al. 2007, Krieger et al. 2008).

The genus *Scaphirhynchus* includes three species occurring in the freshwaters of North America: *Scaphirhynchus albus* (Forbes et Richardson, 1905), *Scaphirhynchus platorhynchus* (Rafinesque, 1820), and *Scaphirhynchus suttkusi* Williams et Clemmer, 1991. Three species of the genus *Pseudoscaphirhynchus* used to inhabit rivers of the Aral Sea basin in Central Asia. The endemic Syr Darya shovelnose sturgeon, *Pseudoscaphirhynchus fedtschenkoi* (Kessler, 1872), has not been recorded since the 1960s (Birstein 1997) and most probably is extinct. The second species, the small Amu Darya shovelnose sturgeon *Pseudoscaphirhynchus hermanni* (Kessler, 1877), endemic in the middle and lower reaches of the Amu Darya drainage, is a very rare, critically endangered fish. Only the third species, the large Amu Darya shovelnose sturgeon, *Pseudoscaphirhynchus kaufmanni* (Kessler, 1877), despite of its disappearance from the most part of historic area, is still recorded in the Vakhsh River and in the middle reaches of the Amu Darya River (Birstein 1997). *Pseudoscaphirhynchus kaufmanni* is represented by two forms with unclear taxonomic relations, namely the common- and the dwarf forms (Sagitov 1969). In the early 1990s, the dwarf form predominated in the population (Birstein 1997).

The comparative karyological studies of Acipenserid fishes are of great importance for many reasons. First of all, the karyological studies in Acipenseridae revealed three ploidy levels in these fishes (Fontana et al. 2008a, Vasil'ev 2009, Vasil'ev et al. 2010), treated as diploid–tetraploid–hexaploid or tetraploid–octoploid–12-ploid (Vasil'ev 2009). Therefore—the polyploidization events in different Acipenserid evolutionary branches as well as the phylogenetic relations between groups with different ploidy levels—call for further studies. Moreover, the knowledge on acipenserid karyotypes is essential in sturgeon aquaculture, as a tool for predicting fertile artificial hybrid lines, because their fertility depends on the ploidy level of the hybridized original parental species. However, the karyotypes of *Pseudoscaphirhynchus* species have never been studied. It should be mentioned that Birstein et al. (1993) presented data on the nuclear DNA content in *P. kaufmanni*, but

this method, as well as the microsatellite analyses—in terms of ploidy level evaluation—are considered indirect (and therefore less accurate) comparative methods. Consequently, they often lead to erroneous conclusions about the ploidy state in some sturgeon species. For example, Ludwig et al. (2001) erroneously estimated octoploid level for *Acipenser mikadoi* Hilgendorf, 1892 and diploid level for *Acipenser dauricus* Georgi, 1775 by the analysis of microsatellite allelic distributions. Thus, they repeat the same mistaken conclusions obtained by Birstein et al. (1993) based on the nuclear DNA content values. In contrast to these inferences, the direct karyological study at first proved about 250 chromosome numbers (tetraploid ploidy level) both for *A. mikadoi* and *A. dauricus* (Vasil'ev et al. 2009).

Therefore, the main goal of this research was to evaluate the ploidy level of *Pseudoscaphirhynchus kaufmanni* by direct karyological study.

MATERIALS AND METHODS

The work was carried out within the Program of EAR-AZA (Euro Asian Regional Association of Zoos and Aquariums) entitled “Conservation of the rarest sturgeons of Eurasia” (<http://earaza.ru/?p=449>). In this Program, we develop methods for artificial propagation of the Amu Darya shovelnose sturgeon, *Pseudoscaphirhynchus kaufmanni*, which is a protected species in Tadzhikistan.

Three specimens used in this study were caught in the Vakhsh River (tributary of the Amu Darya River), Tadzhikistan in April 2012 under the permission of the Tadzhikistan Government. These females, with total body length of 19–23 cm and weighing 50–75 g, were identified as the dwarf form of *Pseudoscaphirhynchus kaufmanni* because they possessed its species-specific characters, namely the tail filament and developed snout thorns (Berg 1948). All females were injected with 0.5–0.75 mL colchicine solution (0.5%–0.6%) depending on their weight. After 4.5–10.5 h cells of the head lymphoid organ were used for chromosome slide preparing by using previously published karyological method (Vasil'ev and Sokolov 1980). The study was conducted by using anesthesia of fishes with a solution of MS-222. Metaphase chromosomes stained in 4% Giemsa solution in phosphate buffer (pH 6.8) were counted with PC software Quick Photo Micro. For karyological description we selected metaphase plates in which the chromosome number can be count accurate to two chromosomes. Totally, 14 metaphase plates were detailed analyzed: five, six, and three plates, respectively, from each female. These numbers are suitable for the primary karyotype description. Chromosome morphology in chromosomes with large and medium sizes was determined according to Levan et al. (1964). The classification of microchromosomes was not possible. Karyotype was arranged using PC software and Adobe Photoshop®CS5.

RESULTS AND DISCUSSION

The karyotype of the large Amu Darya shovelnose sturgeon, *Pseudoscaphirhynchus kaufmanni*, consists of 118–120 chromosomes and includes about 18–20 large bi-armed chromosomes, about 32–34 small bi-armed chromosomes, one–two pairs of large acrocentrics (26th–27th pairs) and about 64 small acrocentrics or microchromosomes (Fig. 1). Among large metacentrics, the smallest last (8th–10th pairs) are about equal or longer than the arm of the first pair of metacentrics, while, the largest first small metacentrics (11th–12th pairs) are distinctly shorter than the arm of the first pair of large metacentrics. The large acrocentrics have their length similar to the length of the arm of chromosomes from the 10th pair of metacentrics.

To date, 18 species from the genus *Acipenser* are karyologically studied, and the only non-karyotyped species is *Acipenser dabryanus* Duméril, 1869. The karyotype of *Scaphirhynchus platorhynchus* is also described (Ohno et al. 1969). All karyotyped *Acipenser*ids are divided into

three discrete groups. The first group includes 120-chromosome species (Table 1). The second group combines sturgeon species with 250–270 chromosomes, which are apparently represented by two subgroups with different karyotypes, namely:

- European species (*Acipenser gueldenstaedtii* Brandt et Ratzeburg, 1833; *Acipenser persicus* Borodin, 1897; and *Acipenser naccarii* Bonaparte, 1836) and Siberian sturgeon, *Acipenser baerii* Brandt, 1869, with about 250 chromosomes;
- Species distributed in America and Eastern Asia (*Acipenser transmontanus* Richardson, 1836; *Acipenser fulvescens* Rafinesque, 1817; *Acipenser medirostris* Ayres, 1854; *Acipenser schrenckii* Brandt, 1869; *Acipenser sinensis* Gray, 1835; *A. mikadoi*; and *A. dauricus*) with about 264–270 chromosomes.

It should be emphasized that the grouping of 250–270-chromosome species correlates with the separation of Atlantic and Pacific clades of *Acipenser*ids in phylogenetic

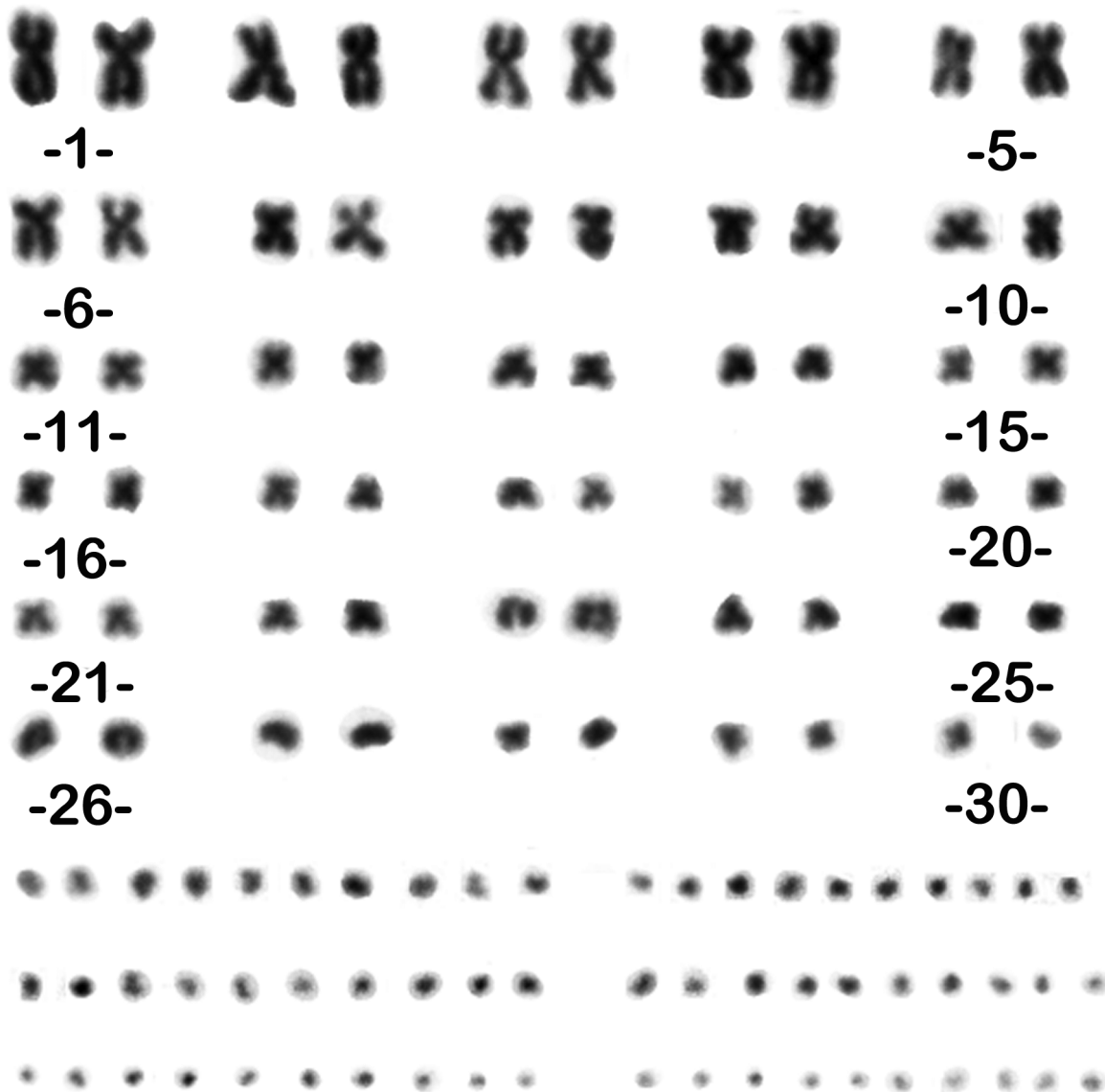


Fig. 1. The karyotype of the Amu Darya sturgeon, *Pseudoscaphirhynchus kaufmanni*; 1–30 = the numeration of the first 30 pairs of the largest chromosomes

Table 1

Karyological characters of the low-chromosome (2n~120) acipenseriform species

Species	2n	LB	SB	LA	M	NF	Reference
<i>Polyodon spathula</i>	120	18–20	20–22	8	~72	~160	Dingerkus and Howell 1976
<i>Scaphirhynchus platorhynchus</i>	~112	18–20	30–32	8–10	~52	~162	Ohno et al. 1969
<i>Pseudoscaphirhynchus kaufmanni</i>	~118–120	18–20	32–34	2–4	~64	~172	Presently reported study
<i>Acipenser sturio</i>	~121	20–22	48–50	2	~48	~191	Fontana and Colombo 1974
<i>Acipenser nudiventris</i>	~118	16–18	42–44	2	~56	~178	Sokolov and Vasil'ev 1989
<i>Acipenser ruthenus</i>	~118	18–20	36–38	2	~58	~174	Vasil'ev 1985 Ráb 1986 Birstein and Vasiliev 1987
<i>Acipenser stellatus</i>	~118	20–22	34–36	4 4–6	~58	~174	Vasil'ev 1985 Birstein and Vasiliev 1987
<i>Acipenser oxyrinchus</i>	~121	24	54	0–2	~41	~199	Fontana et al. 2008b
<i>Acipenser huso</i>	~118	16–18	44–66	2	~36–50	~180–202	Fontana et al. 1998 Birstein and Vasiliev 1987

2n = chromosome number, LB = number of large bi-armed chromosomes, SB = number of small bi-armed chromosomes, LA = number of large acrocentric chromosomes, M = number of microchromosomes, NF = approximate number of chromosome arms (all microchromosomes are classified as acrocentrics); Note: the reported differences in microchromosome numbers in the karyotype of *Acipenser huso* and the increased number of chromosome arms (NF = 202) were caused by very weak chromosome condensation in the karyogram, presented by Fontana et al. (1998); this results in the possibility to define morphology of some microchromosomes.

trees based on molecular genetic data (Ludwig et al. 2001, Krieger et al. 2008). The third group includes the only species *Acipenser brevirostrum* Lesueur, 1818 with 370–372 chromosomes (Kim et al. 2005, Fontana et al. 2008a). According to karyological data obtained in the presently reported study, *Pseudoscaphirhynchus kaufmanni* belongs to the first Acipenserid group of low-chromosome species with about 120 chromosomes.

The detailed analysis of chromosome morphology in the metaphase plates with approximately similar chromosome condensation, demonstrates intraspecific variability in the numbers of different types of chromosomes in low-chromosome species (Table 1). Although slight differences in numbers of large metacentrics can be related to the classification accuracy of these chromosomes in different species, the most noticeable differences are observed in the numbers of small and medium bi-armed chromosomes, as well as large acrocentrics. Exactly, *Polyodon spathula* and *Scaphirhynchus platorhynchus* both have 8–10 large acrocentrics, whereas the other 120-chromosome species have only 2–6 large acrocentrics (Table 1). *Pseudoscaphirhynchus kaufmanni* has 2–4 large acrocentrics (Table 1; Fig. 1); thereby, karyological data does not support close relation between *Scaphirhynchus* and *Pseudoscaphirhynchus* postulated by morphological studies and resulted in their accepted taxonomic integration in the same tribe or subfamily. The same results were obtained by molecular genetic researches, not supported sister relation between *Scaphirhynchus* and *Pseudoscaphirhynchus*, but demonstrated *Pseudoscaphirhynchus* as a sister taxon for the genus *Acipenser* (see Birstein et al. 1997) or for some of *Acipenser* species (Dillman et al. 2007, Krieger et al. 2008).

Thus, monophyly of the subfamily Scaphirhynchinae, or tribe Scaphirhynchini, has been consistently rejected by all phylogenetic reconstruction methodologies with the molecular character set, while monophyly of both genera (*Scaphirhynchus* and *Pseudoscaphirhynchus*) was firmly supported (Dillman et al. 2007).

In a concluding remark, we would like to state that our data on the karyotype structure of *Pseudoscaphirhynchus kaufmanni* confirm polyphyletic origin for the subfamily Scaphirhynchinae (or tribe Scaphirhynchini) previously revealed by molecular genetic studies.

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