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| 4 | Chinese tetraploid loach, Misgurnus anguillicaudatus |
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| 6 | Ya-Juan Li \cdot Zhuo Yu \cdot Ming-Zhao Zhang, \cdot Cong Qian \cdot Syuiti Abe \cdot Katsutoshi Arai |
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| 18 | Abstract When eggs from the Chinese tetraploid loach which had 100 chromosomes |
| 19 | were fertilized with UV irradiated sperm, we obtained viable gynogenetic progeny |

20without any additional treatment for the duplication of maternal chromosomes, which 21survived beyond first-feeding towards adult stage of development. Gynogenetic progeny 22were determined to be diploid since they possessed 50 chromosomes, along with two 23chromosomes bearing nucleolar organizing regions (NORs), detected by silver nitrate 24staining (Ag-NORs), chromomycin-A₃ (CMA₃) positive sites and fluorescence in situ 25hybridization (FISH) signals for rDNA loci. In contrast, when gynogens were induced 26using eggs from diploid loach fertilized by UV irradiated sperm, but without 27chromosome doubling, we found that all resultant progeny were non-viable haploid 28gynogens with 25 chromosomes, along with one NOR-bearing chromosome detected by 29Ag-NOR, CMA₃ and FISH. These observations demonstrate the true genetic tetraploid 30 nature of the Chinese loach possessing 100 chromosomes and the potential use of this 31tetraploid as a source of functional diploid gametes for further ploidy manipulation 32experiments.

- 33
- Keywords Chromosome · Gametes · FISH · Polyploid 34
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37 Introduction

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39Diploid gametes represent key resources in order to expand the chromosome 40 manipulation for improved breeding techniques. If we are able to use diploid gametes 41 for breeding and subsequent manipulation, then it would allow the mass production of 42sterile auto- or allo-triploid lines, synthesis of new amphidiploids or allotetraploid lines, 43recovery of androgenotes from cryopreserved diploid sperm, and other applications for 44aquaculture (Arai 2000, 2001; Arai et al. 2010). However, successful examples of induced tetraploid fish as a source of diploid gametes have been limited due to the 4546 technical difficulties in producing tetraploids by inhibition of early cleavage coupled 47with very low survival rates of the resultant tetraploid progeny (Chourrout et al. 1986; 48Nam et al. 2001; Zou et al. 2004; Sakao et al. 2006). An alternative way would be to use 49the diploid gametes of natural tetraploid lines in certain species or hybrids.

50In dojo loach or Oriental weatherfish Misgurnus anguillicaudatus (Teleostei: 51Cobitidae), most individuals of Japanese wild populations are diploid with 50 52chromosomes. However, tetraploids possessing 100 chromosomes have often been 53identified amongst specimens recovered from Japanese markets (Ojima and Takai 1979; 54Arai et al. 1991a). Using chromosome manipulation techniques, Arai et al. (1991b, 1993) 55concluded that loaches possessing 100 chromosomes were true tetraploids possessing 56four sets of homologous chromosomes, because gynogenetic progeny which were 57artificially induced by fertilizing eggs from these market-sourced specimens with 58genetically-inert UV irradiated sperm were viable without requiring any treatment to 59duplicate the maternal chromosome set. In a similar manner, viable androgenetic 60 progeny were successfully induced by fertilizing UV irradiated eggs with the sperm 61from loach possessing 100 chromosomes (Arai et al. 1995). Gynogenetic and 62androgenetic progeny produced from the gametes of normal diploid fish are non-viable 63 due to the expression of abnormalities collectively referred as 'haploid syndrome', 64 observed in many species of fish including loach (Suzuki et al. 1985; Arai et al. 1992; 65Fujimoto et al. 2007), common carp (Nagy et al. 1978) and salmonids (Chourrout et al. 66 1980; Chourrout 1982; Onozato 1982; Onozato and Yamaha 1983).

In central parts of China, natural tetraploid loach possessing 100 chromosomes co-exist, in the same habitats, with sympatric diploid loach possessing 50 chromosomes (Li et al. 2008). Cytogenetic studies including differential staining with silver nitrate for nucleolar organizing regions (Ag-NOR), fluorochromes such as chromomycin A₃ (CMA₃)/ Distamycin (DA)/ 4',6-diamidino-2-phenylindole (DAPI) and fluorescence *in situ* hybridization (FISH) using human 5.8S+28S rDNA sequences suggested that a loach

73possessing 100 chromosomes should be a tetraploid, based upon a karyotype showing 74quartets of chromosomes and the presence of four FISH signals (Li et al. 2010). However, 75specific genetic characteristic of these Chinese loaches have not yet been determined 76 using the chromosome manipulation approach reported in Arai et al. (1991b, 1993, 771995). If Chinese loaches possessing 100 chromosomes are genetic tetraploids with four 78sets of homologous chromosomes, then it should be possible to use these as a source of 79diploid gametes in order to expand and improve existing breeding techniques. In the 80 present study, we artificially induced gynogenesis using eggs from putative tetraploid 81 females collected from the Chang Jiang River, Hubei Province, China by fertilizing with 82 UV irradiated sperm and then carefully determined survival rates and chromosome 83 numbers in the resultant gynogenetic progeny.

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85 Materials and methods

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| or rish and gametee | 87 | Fish | and | gametes |
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89 Diploid (4 females, 4 males) and tetraploid (4 females) loach were collected in Chang 90 Jiang River in Hubei Province, China and transported to Dalian Ocean University. 91 Identification of tetraploid and diploid specimens was determined in accordance to Li et 92al. (2008). Artificial ovulation was induced by the injection of hCG (human Chorionic 93 Gonadotropin, 20 IU per body gram weight, Aska Pharmaceutical Co. Ltd., Tokyo) 94 according to Suzuki and Yamaguchi (1975). Sperm was collected into a capillary hematocrit 95tube by gently squeezing the male's abdomen (Morishima et al. 2002). Sperm was then diluted 1: 96 100 with Kurokura solution (Kurokura et al. 1984).

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98 Fertilization and gynogenesis

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Eggs taken from diploid (2n) and tetraploid (4n) females were fertilized with sperm from 2n males and then, diploid female x diploid male (control cross, 2n x 2n) and tetraploid female x diploid male (triploid cross, 4n x 2n) crosses were performed. Eggs of 2n and 4n females were then inseminated with UV irradiated sperm from a 2n male (2n x UV and 4n x UV, respectively) to induce gynogenetic development. UV-irradiation of sperm from 2n males was performed according to Suzuki et al. (1985). Fertilized and treated eggs from each cross were reared in a Petri dish (90mm diameter x 15mm depth) containing freshwater (aged tap water) at room temperature.

107 Fertilization rate was calculated as the proportion of cleaved eggs relative to the initial number 108 of eggs. Hatching rate was calculated as the proportion of hatched larvae relative to the number of fertilized eggs. The normal rate was calculated as the proportion of normal larvae relative to the number of hatched larvae. Survival rate at 7 days after hatching (dah) was calculated as the proportion of surviving larvae relative to the number of hatched larvae. Water was changed every day and larvae were first fed *Artemia* from 3 days after hatching. Larvae were incubated at a temperature of 20°C.

- 114
- 115 Chromosome preparation and observation
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Embryos at the optic vesicle stage (15 to 20 h after fertilization at 20°C) were used for chromosome preparation. In physiological saline, the chorion was mechanically removed and embryos treated for 45 min by soaking them in 0.0025% demecolcine (Sigma). Hypotonic treatment was performed with 0.8% citric acid for 20 min and samples were fixed with Carnoy's fixative (3 parts of methanol: 1 part of aceteic acid). Chromosome preparations were prepared from developing embryos as described by Inokuchi et al. (1994). Chromosome slides were stained with Giemsa (Merck) for microscopic observation and karyotyping carried out in accordance with Levan et al. (1964).

- Differential staining with CMA₃/DA/ DAPI (Schweizer 1976; Schweizer et al.1978) and the Ag-NOR method (Howell and Black 1980) was applied in accordance with Li et al. (2010). FISH using human 5.8S+28S rDNA sequences as a probe was applied in accordance with Li et al. (2010).
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128 **Results**

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130 Viability of gynogenetic progeny

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All control crosses (2n x 2n) resulted in high rates of fertilization (83-94%), hatching (75-96%),
normal fry (84-97%) and survival at 7 dah (72-95%; Table 1). Triploid crosses (4n x 2n) also
exhibited high rates of fertilization (77-98%), hatching (51-99%), normal fry (81-99%) and survival
at 7 dah (77-90%; Table 1). Most control and triploid larvae exhibited normal appearance (Fig. 1A).

In gynogenetic crosses, differences were observed between 2n x UV and 4n x UV crosses. Gynogenetic progeny from 2n x UV crosses produced lower rates of fertilization (43-84%) and hatching (3.2-56%) than control and triploid crosses and no normal fry were apparent (Table 1). All resultant fry exhibited abnormalities characteristic of haploidy (Fig. 1B). In contrast, 4n x UV crosses produced high rates of fertilization (69-95%) and hatching (81-99%). Whilst 37 to 67% of resultant larvae were normal (Fig. 1C) or malformed (Fig. 1D), 46-60% fry survived to 7 dah (Table

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144 Chromosomal nature of gynogenetic progeny

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146 As shown in Table 2 and Fig. 2AC, the modal chromosome number of embryos from 2n x 2n crosses 147was diploidy (2n=50), while the mode of embryos from $4n \ge 2n$ crosses was triploidy (3n=75). The 1482n x UV embryos exhibited a haploid range of chromosomes with a modal number of 1n=25 (Table 1492, Fig. 2B). In contrast, 4n x UV embryos exhibited a diploid range of chromosomes with a modal 150number of 2n=50 (Table 2, Fig. 2D). Consequently, most embryos from crosses 2n x 2n, 4n x 2n, 2n 151x UV and $4n \times UV$ crosses were considered diploid (2n=50), triploid (3n=75), gynogenetic haploid 152(1n=25) and gynogenetic diploid (2n=50), respectively. However, hypo- and hyper-chromosome 153numbers were observed in each ploidy group (Table 2). The 2n x UV embryos possessed one 154Ag-NOR/ CMA₃ positive region and FISH signal indicating haploidy (Fig 3A-C), while 4n x UV 155embryos demonstrated two Ag-NOR/ CMA3 positive regions and FISH signals indicating diploidy 156(Fig. 3 D-F).

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158 Discussion

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160 In the present study, viable gynogenetic progeny were produced without chromosome duplication 161from the eggs of Chinese tetraploid loach females fertilized with UV-irradiated loach sperm. 162Progeny possessed 50 chromosomes, equivalent to one half of the 100 chromosomes found in 163 parental tetraploid loach specimens (Li et al. 2010). Gynogenetic fry possessing 25 chromosomes 164that were induced from the eggs of diploid females could not survive beyond hatching or feeding due 165to abnormalities. Diploidy of viable gynogenetic progeny was also demonstrated by the presence of 166 two Ag-NORs, CMA₃ positive sites and FISH signals, whereas the haploid nature of non-viable 167gynogens was indicated by one site and one signal. Consequently, the production of viable 168 gynogenetic progeny suggests that Chinese tetraploid loach are true tetraploids, possessing four sets 169of homologous chromosomes with the ability of generating diploid gametes. Thus, the tetraploid 170nature of the Chinese loach possessing 100 chromosomes was conclusively demonstrated by 171observing the viability of UV-induced gynogenetic progeny. Similar results have already been 172obtained in gynogenetic and androgenetic progeny induced from the gametes of loach possessing 173100 chromosomes obtained from a Japanese market (Arai et al. 1991b, 1993, 1995), and in those 174induced from the gametes of natural tetraploid population in a different cobitid species Cobitis biwae 175(Kusunoki et al. 1994). Therefore, the overall genetic characteristics of Chinese tetraploid loach are 176essentially the same as those of tetraploid loach found in Japanese markets (Arai et al. 1991ab, 1993, 1771995; Arai 2001, 2003). These results suggest the possibility that the tetraploid loach found in 178Japanese markets should be transported from China as food or fishing bait. 179An important application of genetically tetraploid fish is as a source of diploid

180 gametes for the expansion and improvement of existing ploidy manipulation techniques.

181 In Chinese loach, it is also possible to expand ploidy manipulation using diploid 182gametes as reported in our previous studies using natural tetraploid loach found in 183Japanese markets (Arai 2000, 2001, 2003). The mass production of triploids was first realized by crossing tetraploid and diploid fish (Matsubra et al. 1995; Zhang and Arai 184 1851996) whilst a neo-tetraploid strain was produced by inhibiting second polar body 186 release in the 2n x 4n cross (Zhang and Arai 1996; Fujimoto et al. 2010). Fertile 187 hexaploid lines were also produced by inhibiting second polar body release just after 188 fertilization in 4n x 4n crosses. Second generation hexaploid, pentaploid and tetraploid 189 fish were then produced using fertile triploid gametes from hexaploid loaches (Arai et al. 190 1999). Diploid sperm from natural tetraploids is particularly useful for producing viable 191androgenotes by fertilizing UV-irradiated eggs (Arai et al. 1995; Yasui et al. 2010). 192Consequently, the Chinese tetraploid loach represents a valuable resource for inducing 193 various levels of polyploids, gynogens and androgenotes for aquaculture. Breeding 194 programs using diploid gametes are already implemented in allotetraploid crucian carp 195x common carp hybrids (Liu et al. 2001).

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325 Legends for figures

Figure 1 Normal larvae from 2n x 2n (A), abnormal larvae from 2n x genetically inactivated sperm with UV irradiation (UV) (B), normal larvae from 4n x UV (C) and malformed larvae from 4n x UV (D). Bars indicate 1 mm.

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Figure 2 Metaphase spread with (A) 50 chromosomes from a 2n x 2n embryo, (B) 25
chromosomes from a 2n x UV embryo, (C) 75 chromosomes from a 4n x 2n embryo, and
(D) 50 chromosomes from a 4n x UV embryo (D). Bars indicate 10 µm.

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334 Figure 3 Metaphase spreads in 2n x UV (A, B, C) and 4n x UV (D, E, F) dojo loach

335 embryos. Ag-NORs (A, D), CMA₃/DA/DAPI staining positive sites (B, E) and FISH

336 signals detected with 5.8S+28S rDNA probe (C, F) are shown. Arrows indicate NORs

337 shown by Ag staining (A, D), CMA3 positive sites (B, E) or FISH signals (C, F). Bars

indicate 10 μm.







| | No. of | Fertilization | Hatching | Normal rate | Survival rate | | | | |
|----------|--------|------------------|------------|------------------|-----------------|--|--|--|--|
| Cross | NO. 01 | rate (%) | rate (%) | (%) | at 7 days after | | | | |
| | eggs | | | | hatch (%) | | | | |
| 2nx2n-1 | 412 | 92.95 | 79.36 | 91.14 | 75.79 | | | | |
| 2nx2n-2 | 716 | 83.58 | 74.51 | 96.7 | 94.86 | | | | |
| 2nx2n-3 | 448 | 66.07 | 95.61 | 91.17 | 71.73 | | | | |
| 2nx2n-4 | 414 | 93.96 | 78.41 | 84.26 | 73.77 | | | | |
| mean±SD | | 84.14±0.13 | 81.97±0.09 | 90.82±0.05 | 79.04±0.11 | | | | |
| 2nxUV-1 | 701 | 75.89 | 3.2 | 0 | 0 | | | | |
| 2nxUV-2 | 766 | 56.27 | 9 | 0 | 0 | | | | |
| 2nxUV-3 | 433 | 42.96 | 27.96 | 0 | 0 | | | | |
| 2nx2nUV- | 565 | 83.89 | 56.33 | 0 | 0 | | | | |
| mean±SD | | 64.75±0.19 | 24.12±0.24 | 0 | 0 | | | | |
| 4nx2n-1 | 1210 | 97.69 | 51.02 | 88.72 | 76.78 | | | | |
| 4nx2n-2 | 850 | 77.41 | 91.79 | 98.68 | 89.74 | | | | |
| 4nx2n-3 | 778 | 89.2 | 99.14 | 80.52 | 77.91 | | | | |
| 4nx2n-4 | 524 | 90.65 | 94.32 | 84.38 | 84.38 | | | | |
| mean±SD | | 88.74 ± 0.08 | 84.06±0.22 | 88.07 ± 0.08 | 82.20±0.06 | | | | |
| 4nxUV-1 | 480 | 95.21 | 81.33 | 66.67 | 54.53 | | | | |
| 4nxUV-2 | 489 | 73.62 | 98.89 | 44.1 | 59.55 | | | | |
| 4nxUV-3 | 695 | 68.78 | 96.23 | 37.39 | 49.13 | | | | |
| 4nxUV-4 | 744 | 70.97 | 92.42 | 48.57 | 45.7 | | | | |
| mean±SD | | 77.14±0.12 | 92.22±0.08 | 49.18±0.13 | 52.23±0.06 | | | | |

Table 1. Number of eggs, fertilization rate, hatching rate, normal rate and survival rateat 7 days after hatch in different crosses

| | Embryo | Cell | Chromosome number frequency | | | | | | | | | | | | | | | | | | |
|-------------------------------|--------|--------|-----------------------------|----|----|--------|------|----|----|----|----|----|----|----|----|----|----|----|----|----|----|
| Cross | number | number | 23 | 24 | 25 | 26 | 27 | | 40 | 41 | 42 | 43 | 44 | 45 | 46 | 47 | 48 | 49 | 50 | 51 | 52 |
| 2n x 2n | 17 | 72 | 0 | 0 | 0 | 0 | 0 | | 1 | 3 | 2 | 2 | 2 | 1 | 4 | 4 | 5 | 2 | 41 | 2 | 3 |
| 2n x UV | 11 | 39 | 5 | 6 | 26 | 1 | 1 | | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| 4n x 2n | 16 | 77 | 0 | 0 | 0 | 0 | 0 | | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| 4n x UV | 17 | 100 | 0 | 0 | 0 | 0 | 0 | | 0 | 0 | 0 | 2 | 1 | 0 | 3 | 10 | 9 | 3 | 44 | 3 | 6 |
| Cross Chromosome number frequ | | | | | | freque | ency | | | | | | | | | | | | | | |
| Closs | | | 53 | 54 | 56 | 57 | | 66 | 67 | 68 | 69 | 70 | 71 | 72 | 73 | 74 | 75 | 76 | 77 | 78 | 79 |
| 2n x 2n | | | 0 | 0 | 0 | 0 | | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| 2n x UV | | | 0 | 0 | 0 | 0 | | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| 4n x 2n | | | 0 | 0 | 0 | 0 | | 1 | 0 | 2 | 4 | 5 | 3 | 8 | 5 | 2 | 31 | 5 | 1 | 4 | 6 |
| 4n x UV | | | 5 | 11 | 2 | 1 | | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |

Table 2. Metaphase chromosome count in Dojo loach embryos originating from different crosses

 $2n \times 2n$: control cross from fertilization of eggs of diploid female with sperm of diploid male, $2n \times UV$:gynogenetic cross from fertilization of eggs of diploid female with UV-irradiated sperm, $4n \times 2n$: cross from fertilization of eggs of tetrap female with sprem of diploid male, $4n \times UV$: gynogenetic cross from fertilization of eggs of tetraploid female with UV-irradiated sperm