

Chromomycin A₃ and DAPI Staining of Chromosomes of Three Endemic *Pseudophoxinus* Bleeker, 1860 (Teleostei: Leuciscidae) Species from Anatolia

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Research Article

Received 30 December 2019; Accepted 10 April 2020; Release date 01 June 2020.

How to Cite: Karasu Ayata, M. (2020). Chromomycin A₃ and DAPI staining of chromosomes of three endemic *Pseudophoxinus* Bleeker, 1860 (Teleostei: Leuciscidae) species from Anatolia. *Acta Aquatica Turcica*, 16(2), 283-289. <https://doi.org/10.22392/actaquatr.667595>

Abstract

The karyotypes and other chromosomal markers of representatives of the genus *Pseudophoxinus* have been little studied. Therefore, this study documents chromosomal NOR phenotypes as revealed using Chromomycin A₃ (CMA₃) and DAPI chromosomal stainings in three species *Pseudophoxinus elizavetae* Bogutskaya, Küçük & Atalay, 2007, *P. firati* Bogutskaya, Küçük & Atalay, 2007 and *P. hittitorum* Freyhof & Özuluğ, 2010, endemic in Anatolia. Simple NOR phenotype with only two CMA₃ positive signals were observed in the karyotype of *P. firati* whereas derived NOR phenotype with four CMA₃ positive signals were observed in those of *P. elizavetae* and *P. hittitorum*, respectively. No DAPI positive signal was detected in karyotypes in all three species. This study described new NOR phenotypes, i.e. number and position of major rDNA genes, in genomes of three *Pseudophoxinus* species contributing thus to known diversity of NOR phenotypes in otherwise karyotypically highly conservative leuciscid fishes.

Keywords: Fish cytotaxonomy, chromosome banding, chromosomal markers, major rDNA genes

Anadolu'da Yaşayan Endemik Üç *Pseudophoxinus* Bleeker, 1860 (Teleostei: Leuciscidae) Türünün CMA₃ ve DAPI Boyalı Kromozomları

Özet

Pseudophoxinus cinsindeki balıkların kromozom ve diğer sitotaksonomik markerları konusundaki çalışma sayısı azdır. Bu çalışmada, Anadolu'ya endemik üç leuciscine türü *Pseudophoxinus elizavetae* Bogutskaya, Küçük & Atalay, 2007, *P. firati* Bogutskaya, Küçük & Atalay, 2007 ve *P. hittitorum* Freyhof & Özuluğ, 2010'da kromozomal NOR fenotipleri CMA₃ ve DAPI kromozomal boyama ile çalışılmıştır. *P. firati*'nin karyotipinde iki, *P. elizavetae* ve *P. hittitorum*'un karyotiplerinde ise dört CMA₃ pozitif sinyal gözlenmiştir. Üç türün karyotipinde de DAPI pozitif sinyal gözlenmemiştir. Bu çalışma üç *Pseudophoxinus* türünün genomunda majör rDNA genlerinin sayısı ve pozisyonu gibi yeni NOR fenotiplerini tanımlamıştır. Böylece karyotipik olarak yüksek derecede konservatif bilinen leuciscid NOR fenotip çeşitliliğine katkıda bulunur.

Anahtar kelimeler: Balık sitotaksonomisi, kromozomal bantlama, kromozom markerları, majör rDNA genleri

INTRODUCTION

The genus *Pseudophoxinus* Bleeker, 1860 belongs to family Leuciscidae and its subfamily Leuciscinae (Tan and Ambruster, 2018). The genus *Pseudophoxinus* contains at least 30 locally endemic species in Anatolia and other countries of Near and Middle East (Froese and Pauly, 2020). This genus has 21 endemic species in the freshwaters of Anatolia (Güçlü and Küçük, 2017; Saç et al., 2019). Karyological studies of representatives of this genus have started just recently (Karasu, 2009). In, the last decade, *Pseudophoxinus antalyae*, *P. battalgilae*, *P. burduricus*, *P. crassus*, *P. egridiri*, *P. elizavetae*, *P. evliyaie*, *P. fahrettini*, *P. firati*, *P. hittitorum*, *P. maeandri* and *P. zekayi* have been studied cytogenetically (Table 1). In these studies, karyotypes, C-banding and Ag-NOR stainings have been performed. Also, G- and Q-banding patterns in the karyotype of *P. antalyae* have been reported (Ergene et al., 2010).

Table 1. Cytogenetic data for the representatives of the genus *Pseudophoxinus* from Anatolia, Turkey

Species	2n	Chromosome morphology	FN	Ag-NOR number	References
<i>P. antalyae</i>	50	16m+14sm+12st+8a	92	2	Ergene et al., 2010
<i>P. firati</i>	50	38m-sm+12st	88	4	Karasu et al., 2011
<i>P. elizavetae</i>	50	8m+34sm+8st	92	4	Gaffaroğlu et al., 2014
<i>P. crassus</i>	50	12m+30sm+8st-a	92	2	Ünal et al., 2014
<i>P. hittitorum</i>	50	14m+26sm+10st-a	90	2	Ünal et al., 2014
<i>P. battalgilae</i>	50	16m+28sm+6st-a	94	4	Ayata et al., 2016
<i>P. burduricus</i>	50	18m+26sm+6st-a	94	2	Ayata et al., 2016
<i>P. egridiri</i>	50	14m+28sm+8st-a	92	2	Ayata et al., 2016
<i>P. evliyae</i>	50	14m+30sm+6st-a	94	4	Ayata et al., 2016
<i>P. fahrettini</i>	50	16m+26sm+8st-a	92	2	Ayata et al., 2016
<i>P. maeandri</i>	50	10m+32sm+8st-a	92	4	Ayata et al., 2016
<i>P. zekayi</i>	50	16m+26sm+8st-a	92	2	Ünal and Gaffaroğlu, 2016

2n: diploid chromosome number, FN: fundamental number, m: metacentric, sm: submetacentric, st-a: subtelo-acrocentric

The number and locations of NORs have been used as a cytotoxic character in fish cytotoxicology (Rábová et al., 2015). Ag-NOR staining is a very common cytotoxic technique (Rábová et al., 2015). However, this method identifies only transcriptionally active rDNA genes. Also, the number and location of NORs have been determined using CMA₃ staining (Gornung, 2013; Rábová et al., 2015). This staining is based on GC-specific fluorochrome and detects both active and inactive NORs in fish (Gornung, 2013). Both Ag- and CMA₃ stainings have been done in many fish species (Arai, 2011; Gornung, 2013). Fluorescence staining with 4',6-diamidino-2-phenyl-indole (DAPI) dyes detect chromosomal regions that contain AT-rich DNA (Kapuscinski, 1995). This method is studied on many fish species too (Mayr et al., 1986; Valic et al., 2010).

A presence of four Ag-NORs on submetacentric (sm) chromosomes in karyotype of *P. elizavetae* (Gaffaroğlu et al., 2014) and four Ag-NORs on sm-subtelocentric (st) chromosomes in that of *P. firati* (Karasu et al., 2011) have previously been reported. Additionally, Ünal et al. (2014) observed two Ag-NORs on sm chromosome pair in karyotype of *P. hittitorum*. The present study was carried out to determine NOR phenotypes using CMA₃ staining and DAPI stainings in *P. elizavetae*, *P. firati* and *P. hittitorum*, endemic in Anatolia.

MATERIALS and METHODS

The analyzed individuals of *P. elizavetae*, *P. firati* and *P. hittitorum* were collected by electrofishing from Anatolia, Turkey (Table 2, Figure 1). The specimens were carried alive to the laboratory. The chromosomes were obtained from head kidney cells according to the air drying technique of Bertollo et al. (2015). Slides were stained with CMA₃/DAPI stainings according to Rábová et al. (2015). For this technique, slides were treated with an ice-cold ethanol series (70%, 90%, 100%, 2 minutes in each). After this step, work continued in the dark. Slides were incubated in McIlvaine buffer with MgCl₂ for 10 minutes at room temperature. Then stained with CMA₃ in a wet chamber for 15 minutes, at room temperature. Slides were covered with a coverslip and washed up shortly in McIlvaine buffer and allowed the coverslip to detach. After this step, slides were incubated in McIlvaine buffer with Methyl Green for 15 minutes at room temperature. Slides were washed shortly in McIlvaine buffer again. Then, slides were stained with DAPI in a wet chamber for 20 minutes at room temperature. Slides were washed shortly in McIlvaine buffer and drops were removed by touching slide, mounted with antifade (glycerol-propylgalate), and sealed the margins with nail polish. Sealed slides were placed on a tray covered by aluminum foil and kept in the fridge several days before the examination. Slides were screened in an Olympus BX 60 epifluorescence microscope equipped with a DP50 Olympus CCD camera. At least 10 metaphases were examined per individual. Karyotypes from CMA₃ stained metaphases were arranged manually.

Table 2. General information regarding collected individuals

Species	Locality	Coordinates	Date	Number
<i>P. elizavetae</i>	Sultan Swamps, Develi, Kayseri	38°22'N, 35°21'E	2010	15
<i>P. firati</i>	Tohma Creek, Gürün, Sivas	38°47'N, 36°57'E	2009	20
<i>P. hittitorum</i>	Spring Eflatun, Beyşehir, Konya	37°52'N, 31°34'E	2010	8

**Figure 1.** The map of Turkey with three collection sites of *Pseudophoxinus* species. The map was created using the software Google Earth Pro

RESULTS and DISCUSSION

Karyotypes of examined species corresponded to previous reports, i.e. $2n = 50$ and karyotypes composed of 8 m chromosomes, 34 sm chromosomes and 8 st chromosomes in *P. elizavetae*; 38 m-sm chromosomes and 12 st chromosomes in *P. firati*; 14 m chromosomes, 26 sm chromosomes, and 10 st-a chromosomes in *P. hittitorum* (Karasu et al., 2011; Gaffaroğlu et al., 2014; Ünal et al., 2014).

Four CMA₃ positive signals were observed in chromosomes of *P. elizavetae* (Figure 2A) and those of *P. hittitorum* (Figure 2E) whereas two CMA₃ positive signals were observed in the genome of *P. firati* (Figure 2C). CMA₃ staining revealed their location in the whole short (p) arms of medium-sized sm chromosome pair in karyotypes of all three species. No DAPI positive signal was detected in the genomes of examined species (Figures 2B, D, F).

Molecular methods are a highly efficient tool for providing species-specific markers in many areas of fish biology. As a result, the comprehensive use of both cytogenetic and molecular methods increase the level of certainty also in fish taxonomy (Kirtiklis et al., 2010; Gornung, 2013). Among the cytogenetic markers, NOR phenotypes, i.e. number and position of major rDNA sites on chromosomes, have played an important role in fish comparative cytogenetics. The number and localization of NORs have mainly been studied using silver nitrate (AgNO₃), Mithramycin (MM), or CMA₃ staining (e.g. Mani et al., 2009; Gornung, 2013; Rábová et al., 2015).

This study is the first attempt to characterize the CMA₃ and DAPI characteristics in Anatolian freshwater fishes. The result of CMA₃ staining confirmed Ag-NOR phenotype observed in *P. elizavetae* (Gaffaroğlu et al., 2014). In this study, CMA₃ positive signals were located in no. 7 and no. 9 in karyotype of *P. elizavetae* (Figure 3A). However, two more positive NOR sites were detected in chromosomes of *P. hittitorum* by CMA₃ staining. CMA₃ positive signals were located in no. 9 and no. 13 in karyotype of *P. hittitorum* (Figure 3C). This polymorphism could be the result of translocation of rDNA sites. Only two CMA₃ positive signals were observed in no. 5 in genome of *P. firati* (Figure 3B). In this sense, four Ag-NORs that reported by Karasu et al. (2011) should be a polymorphism in this species. Consequently, our data suggest that major rDNA genes have been conserved in sm chromosomes in the genomes of three endemic species of the genus *Pseudophoxinus*.

Characteristics of CMA₃ staining in *P. firati* resembles that in other leuciscins for instance *Leuciscus borysthenticus* (Ráb et al., 1996), *Pachychilon macedonicum* (Ráb et al., 2000), *Vimba vimba* and *V. elongata* (Rábová et al., 2003), some *Rutilus* and *Scardinius* species (Bianco et al.,

2004), *Squalius pyrenaicus* (Gromicho and Collares-Pereira, 2004), *S. aradensis* and *S. torgalensis* (Nabais et al. 2013).

Moreover, results of CMA₃ staining in *P. elizavetae* and *P. hittitorum* resembles that of *Leuciscus idus*, *L. cephalus* and *L. leuciscus* (Boron et al., 2009), *Iberochondrostoma almacai* and *I. lusitanicum* (Monteiro et al., 2009) and *Telestes ukliwa* (Valic et al., 2010).

On the other hand, staining with DAPI did not show any positive signals along *P. elizavetae*, *P. firati* and *P. hittitorum* chromosomes. The same results were observed in chromosomes of other cyprinoid species (Mayr et al., 1986; Valic et al., 2010). Valic et al. (2010) reported that that this could be another common feature for genomes of cyprinids.

Also, FISH experiments have to be included in future studies to determine the exact number and position of NORs of the studied species. In conclusion, this study revealed the details of the NOR phenotypes in the three endemic species using CMA₃ and DAPI stainings.

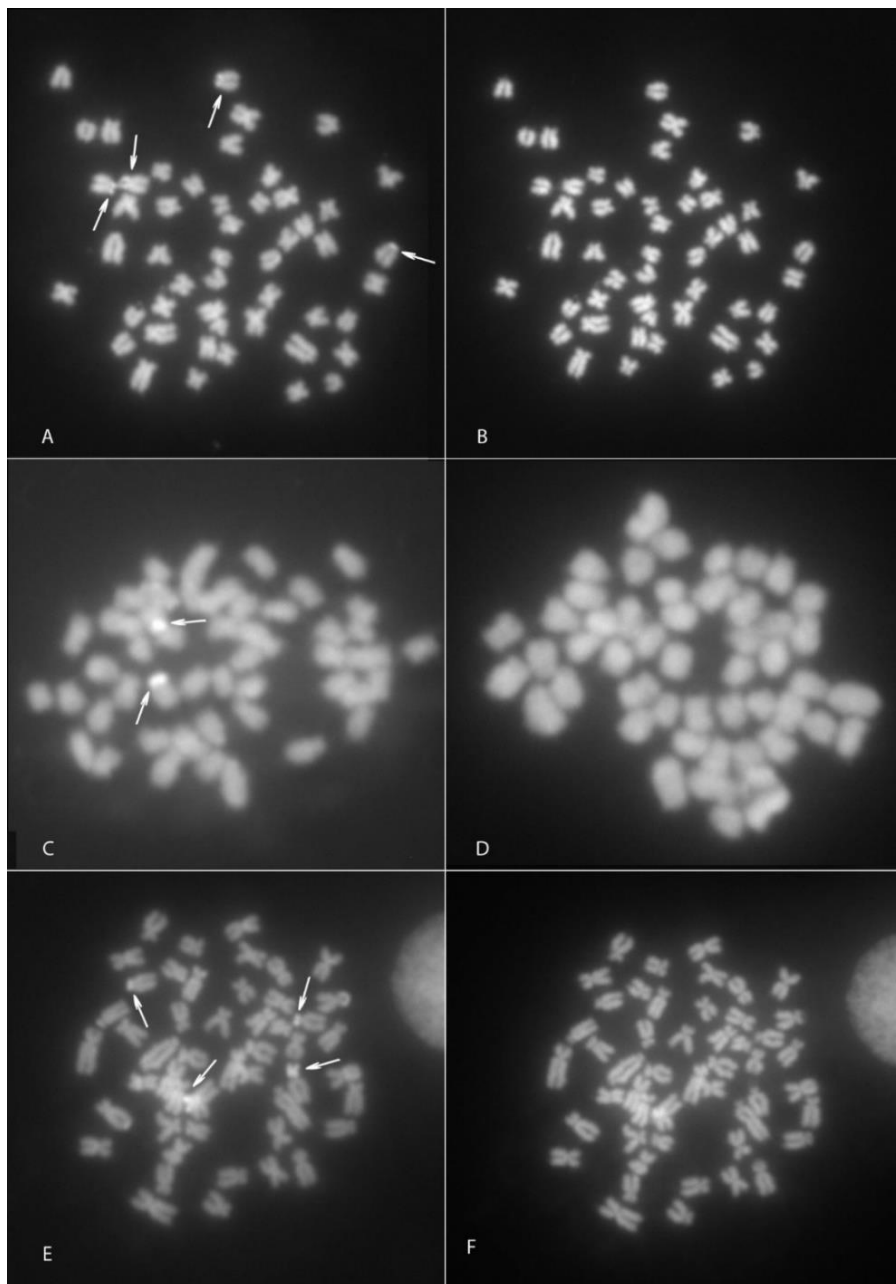


Figure 2. Metaphase of (A) *P. elizavetae*, (C) *P. firati*, (E) *P. hittitorum* after CMA₃ staining. Arrows indicate the CMA₃ positive signals. Metaphases of (B) *P. elizavetae*, (D) *P. firati*, (F) *P. hittitorum* after DAPI staining

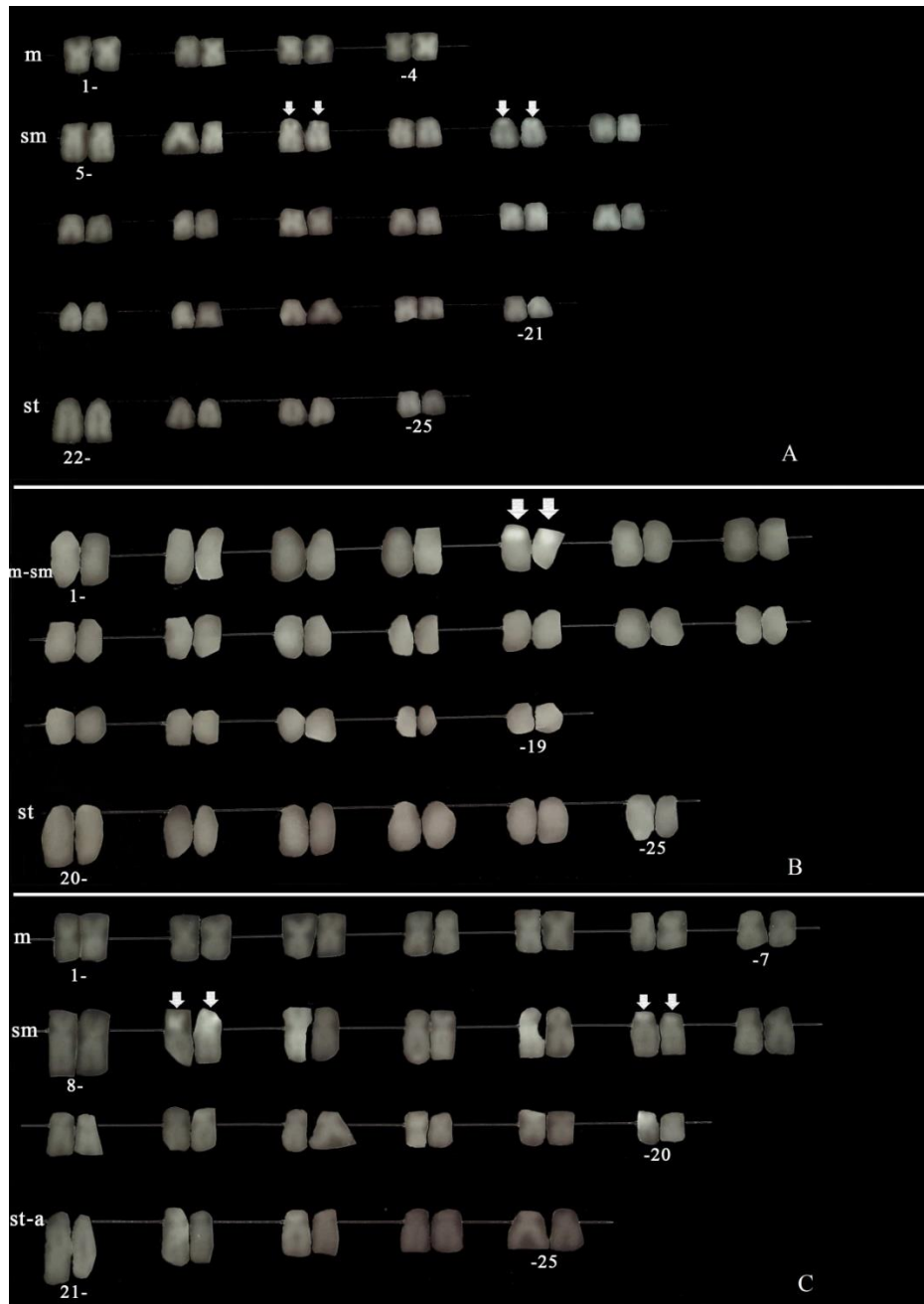


Figure 3. Arranged karyotypes of (A) *P. elizavetae*, (B) *P. firati*, (C) *P. hittitorum* after CMA₃ staining. Arrows indicate the CMA₃ positive signals

Acknowledgments: This study was supported by Kırşehir Ahi Evran University Scientific Research Projects (Project code: SYO.A4.18.001). The author is thankful to Prof. Muhammet Gaffaroğlu (Kırşehir Ahi Evran University) and Dr. Sevgi Ünal (Bartın University) for helping in the field and to Prof. Petr Ráb (Laboratory of Fish Genetics, Czechia) for his laboratory support.

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