

Polymorphism of white muscle myosin and parvalbumins in the genus *Barbus* (Teleostei: Cyprinidae)

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Muscle proteins were investigated in two large European barbels, *Barbus barbus* and *B. meridionalis*, and in four small tropical barbels native to SE Asia: *B. conchoniis*, *B. tetrazona*, *B. sachsi* and *B. itteya*. Polyacrylamide gel electrophoresis was used to analyse myosin heavy and light chains and parvalbumin isotypes from white trunk muscle. Each species could be biochemically identified. The myosin subunit and parvalbumin isotype patterns obtained for the two European barbels were similar. The Asian barbels, on the other hand, not only differed from the European species but displayed a greater diversity within their group. These biochemical results are largely in agreement with morphological and genetic data, but fail to substantiate suggested close relationships between Asian barbel species.

Key words: myosin; light chains; heavy chains; parvalbumins; fish; *Barbus*.

I. INTRODUCTION

The genus *Barbus* (Cyprinidae) currently comprises some 800 species in Europe, Africa and Asia. These species present considerable diversity. The common species *Barbus barbus* (L.) is the main representative in Europe and has the largest distribution area, from the Atlantic to the Black Sea. It is taxonomically related to another European barbel, the Mediterranean barbel *Barbus meridionalis* R. (Almaça, 1984). These two river fishes have been found to hybridize naturally in southern France, producing fertile descendants (Berrebi *et al.*, 1987). Berrebi *et al.* (1990) have shown, by genetic differentiation, that African barbels form two distinct lineages: large, tetraploid species and small, diploid ones. For these authors, the large African barbels resemble, in many respects, their European counterparts, which are also tetraploid. The small African and Asian specimens, on the other hand, would appear to constitute another, less homogeneous group. The heterogeneity of the Asian barbels is further illustrated by the fact that they are often subdivided into three genera (*Puntius*, *Capoeta* and *Barbodes*) according to the number of barbs (Berrebi *et al.*, 1990).

Myosin from fish muscles exhibits a subunit structure and isozymic distribution resembling those of mammalian muscle myosin. Fish white-fibre myosin is composed of two heavy chains (HC, 200 kD) and three types of light chains (LC₁, LC₂ and LC₃, 16 to 25 kD). Light chains in various teleosts have been extensively studied by polyacrylamide gel electrophoresis (Focant *et al.*, 1976; Huriaux & Focant, 1977, 1985; Watabe *et al.*, 1982; Huriaux *et al.*, 1983, 1990; Focant & Pequeux, 1985; Rowlerson *et al.*, 1985; Ochiai *et al.*, 1988, 1990; Martinez *et al.*, 1990*a, b*). LC₁ and LC₃, especially the latter, show significant species-related

characteristics. A single isoform of myosin heavy chain has generally been detected (Karasinsky & Kilarsky, 1989; Martinez *et al.*, 1990*a, b*). Fish sarcoplasm contains low-molecular-weight calcium-binding proteins, the parvalbumins. These are abundant in white fibres and frequently present several isotypes (for a review, see Gerday, 1982). Electrophoretic analysis of myosin light chains and parvalbumins has made it possible to discriminate between closely related species of Gobiidae (Focant & Joyeux, 1988), Serranidae (Focant *et al.*, 1988), Labridae (Focant *et al.*, 1990) and Cichlidae (Focant & Vandewalle, 1991). So far, however, it has not been possible, by this means, to establish any phylogenetic correlations within a genus.

We have recently identified the myosin subunits and parvalbumin isotypes of several trunk and head muscles from *Barbus barbus* (Huriaux *et al.*, 1990, 1991*a, b*; Focant *et al.*, 1992). The white muscle myosin light chains exhibit species-specific properties as well as ones common to fish. A single type of heavy chain was found in trunk muscle but two isotypes were detected in head muscles. Three parvalbumin isotypes were identified in the dorso-lateral muscle; their distribution varies with fish size and according to the antero-posterior myotomal location, but is influenced neither by the growth rate nor by life style.

To gain a more detailed view of biochemical polymorphism in the genus *Barbus* we have compared, on polyacrylamide gels, the electrophoretic mobility of myosin heavy and light chains and parvalbumin isotypes from the trunk white fibres of the common barbel *Barbus barbus* (Linné, 1758), with the corresponding proteins from the Mediterranean barbel *Barbus meridionalis* Risso, 1826 and four tropical barbels native to regions of SE Asia, rosy barbel *Barbus conchoni* (Hamilton-Buchanan, 1822), tiger barbel *Barbus tetrazona* (Bleeker, 1855), gold barbel *Barbus sachs* Ahl, 1923 and cherry barbel *Barbus titteya* (Deraniyagala, 1929). These Asian species were previously chosen by Frankel for studies on the tissue-specificity and ontogenetic expression of isozymes in the genus *Barbus* (Varma & Frankel, 1980; Frankel, 1983, 1985; Frankel & Wilson, 1984, 1985).

II. MATERIALS AND METHODS

MUSCLE SAMPLES

B. meridionalis [seven specimens; standard length (S.L.) 12–14.5 cm] were caught in the River Lergue (area of Montpellier, France). The Asian barbels were obtained from a local retail distributor in Liège. Three sets of experiments were performed; one *B. conchoni* (S.L. 4 cm), two *B. tetrazona* (S.L. 3 cm), three *B. sachs* (S.L. 3 cm) and three *B. titteya* (S.L. 2.4 cm) were examined in each set. The whole dorso-lateral white muscle was dissected from the Asian barbels. In the case of *B. meridionalis*, only the anterior part of this white muscle located in front of the dorsal fin was used. The corresponding muscle samples from *B. barbus* have previously been analysed (Huriaux *et al.*, 1990, 1991*a, b*) and were used as references in this work.

The various muscles were minced and kept in 10 vol. of a solution containing 0.01 M Tris-HCl, 0.05 M KCl, 0.01 M DTT, 50% glycerol (v/v), 0.005% NaN₃ (w/v), pH 7.5, at –20°C for one to several weeks.

PREPARATION OF PROTEINS

Actomyosin preparation, isolation of parvalbumin-containing sarcoplasmic extract, and incubation of samples were performed according to Huriaux *et al.* (1990).

POLYACRYLAMIDE GEL ELECTROPHORESIS

Vertical-slab polyacrylamide gel (18 × 8 × 0.2 cm) electrophoresis (PAGE) was used under several sets of conditions.

Myosin light chains were separated in an 8 M urea, 10% acrylamide gel at pH 8.6 (Perrie & Perry, 1970) or in a sodium dodecyl sulphate (SDS)-discontinuous-buffer system (separating gel: 20% acrylamide, pH 8.4) according to Laemmli (1970) with the modifications described by Huriaux *et al.* (1990). Myosin heavy chains were discerned in a SDS-discontinuous-buffer system (separating gel: 6% acrylamide) according to Danieli-Betto *et al.* (1986), but with further inclusion of 30, 40 or 50% glycerol (w/v) at pH 8.8 or 8.4 (Huriaux *et al.*, 1991*b*). Parvalbumin isotypes were analysed in 10% glycerol (w/v), 10% acrylamide gel at pH 8.6 (Focant *et al.*, 1981).

Conditions for staining with Coomassie Brilliant Blue R-250 and destaining have been previously described (Huriaux & Focant, 1977). Densitometry was performed with a Helena Quick-Scan apparatus (Beaumont, Texas).

III. RESULTS

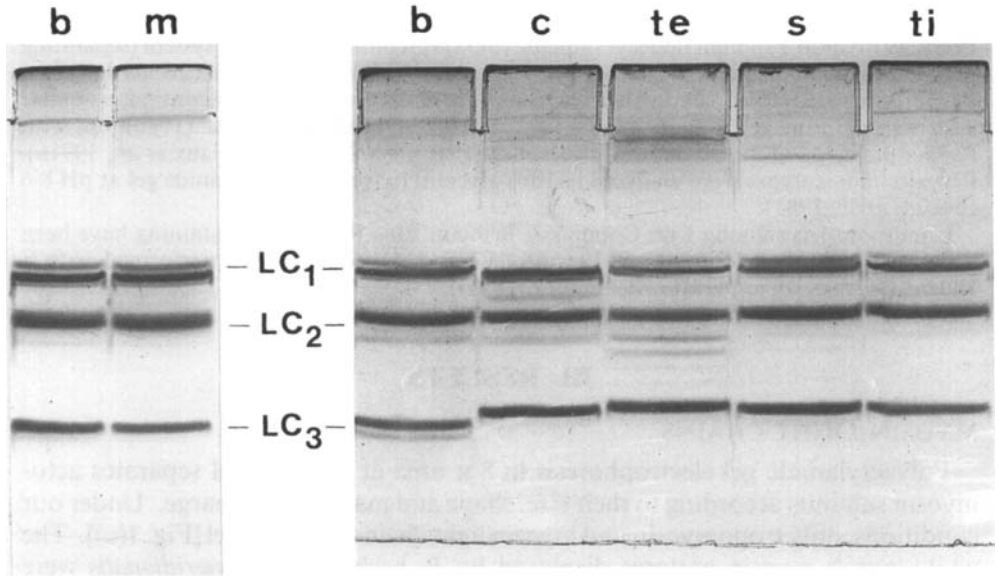
MYOSIN LIGHT CHAINS

Polyacrylamide gel electrophoresis in 8 M urea at alkaline pH separates actomyosin subunits according to their size, shape and mainly their charge. Under our conditions, only tropomyosin and myosin light chains enter the gel [Fig. 1(a)]. The white-muscle myosin patterns displayed by *B. barbus* and *B. meridionalis* were similar, while the patterns obtained with the four Asian barbels presented some peculiarities: LC₁ from *B. conchoni* migrated a little faster; for all four Asian fishes the mobility of LC₃ was the same, but inferior to that of the corresponding light chain from both European specimens. All LC₂ looked similar, sometimes yielding faster, minor bands which could reflect their level of phosphorylation or degradation.

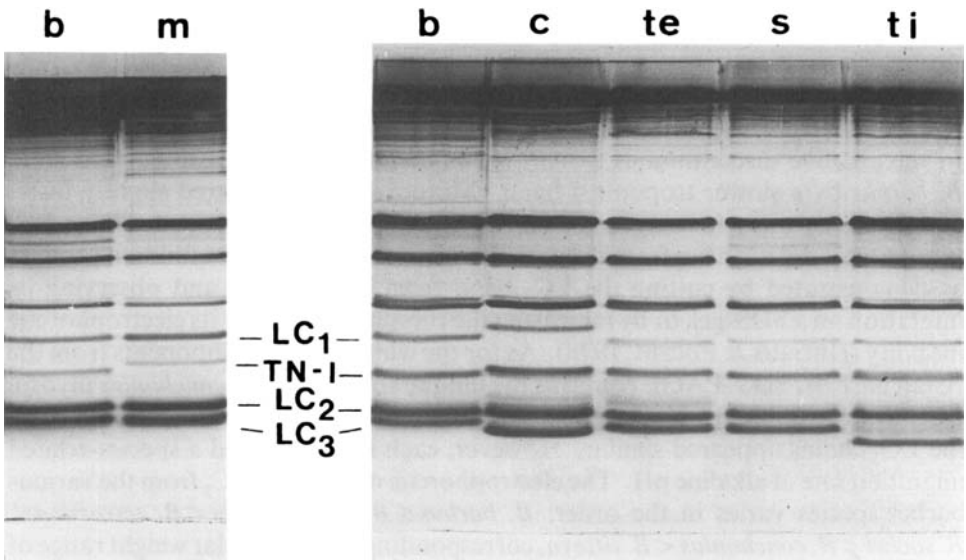
In SDS-PAGE experiments, where proteins are separated according to their molecular weight, the various myofibrillar components can be observed [Fig. 1(b)]. In the alkaline discontinuous system, *B. meridionalis* can be distinguished from *B. barbus* by a slower troponin-I band. Moreover, LC₃ appeared slightly faster and distinctly more heterogeneous in the seven specimens examined. These LC₃ peculiarities did not arise from the presence of a co-migrating contaminant protein, as demonstrated by cutting the LC₃ band from the urea gel and observing its migration on a SDS-gel, or by increasing the running pH to alter its electrophoretic mobility (Huriaux & Focant, 1978). As for the white-muscle components from the Asian barbels, SDS-PAGE confirms the unique structure of *B. conchoni* myosin LC₁, which exhibited a higher apparent molecular weight. As in the urea gels, all the LC₂ chains appeared similar. However, each LC₃ displayed a species-related migration rate at alkaline pH. The electrophoretic mobility of LC₃ from the various barbel species varies in the order: *B. barbus* ≤ *B. meridionalis* < *B. tetrazona* ≤ *B. sachsi* < *B. conchoni* < *B. titteya*, corresponding to a molecular weight range of 17 000–16 300.

MYOSIN HEAVY CHAINS

High-porosity SDS-PAGE of white-muscle myosin heavy chains revealed a single band for each barbel species (Fig. 2). Under all electrophoretic conditions investigated, the heavy chains from *B. barbus* and *B. meridionalis* were very similar, while heavy-chain mobility was a little higher for the Asian fishes. These differences were confirmed by densitometric scanning of gels on which *B. barbus* myosin was allowed to comigrate with each Asian barbel myosin (Fig. 3). This procedure



(a)



(b)

FIG. 1. (a) Urea and (b) SDS-PAGE of actomyosin from *B. meridionalis* (m), *B. conchonus* (c), *B. tetrazona* (te), *B. sachsi* (s) and *B. titteya* (ti) compared with *B. barbus* (b).

further allowed the detection of some very slight specific variations among the latter fishes. The relative mobility of the various heavy chains increased in the order: *B. barbus* = *B. meridionalis* < *B. sachsi* < *B. titteya* < *B. tetrazona* < *B. conchonus*.

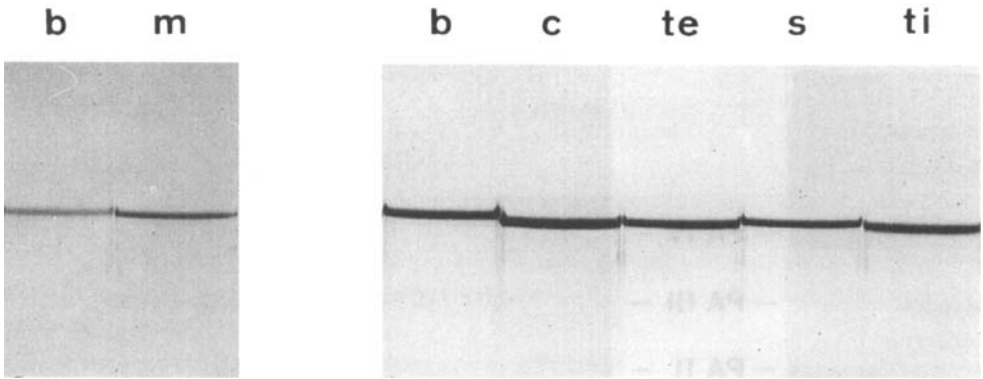


FIG. 2. SDS-PAGE at pH 8.8 and 30% glycerol of myosin heavy chains from *B. meridionalis* (m), *B. conchoniuis* (c), *B. tetrazona* (te), *B. sachsi* (s) and *B. titteya* (ti) compared with *B. barbus* (b).

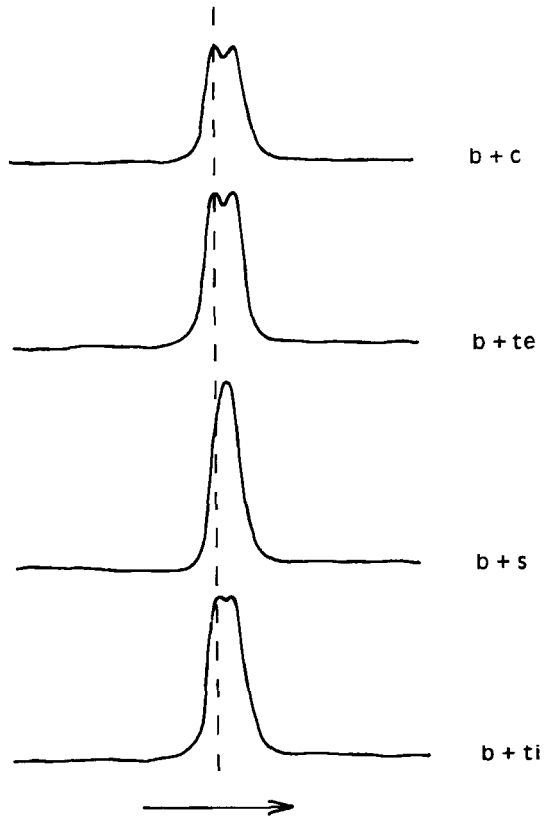


FIG. 3. Densitometric traces of comigrating myosin heavy chains from *B. barbus* (b) with *B. conchoniuis* (c), *B. tetrazona* (te), *B. sachsi* (s) or *B. titteya* (ti) in SDS-PAGE at pH 8.8 and 30% glycerol. The vertical dotted line indicates the *B. barbus* heavy-chain location.

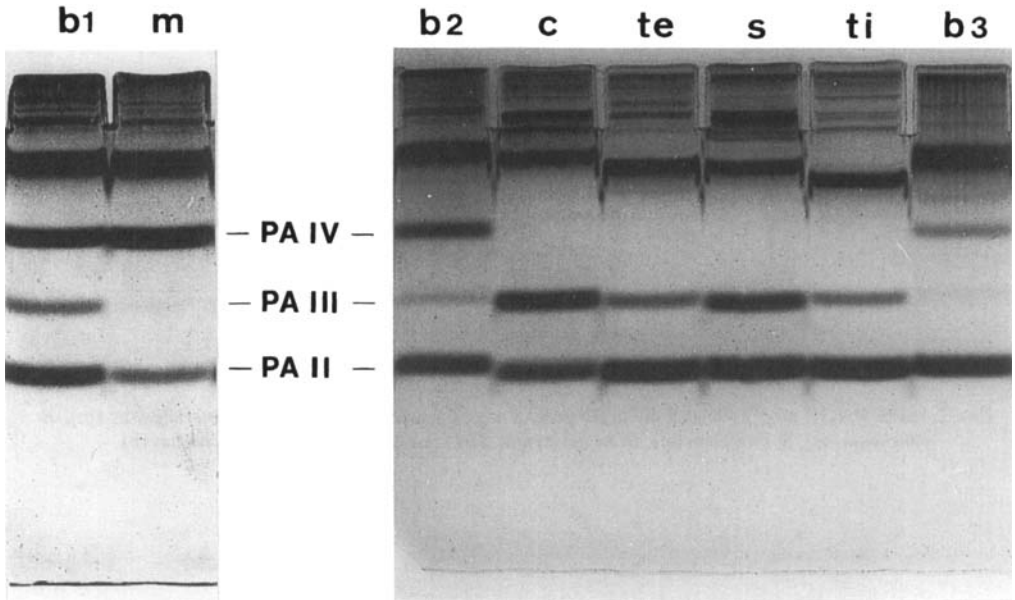


FIG. 4. Glycerol-PAGE of sarcoplasmic extracts from *B. meridionalis* (m), *B. conchoniis* (c), *B. tetrazona* (te), *B. sachsi* (s) and *B. titteya* (ti) compared with *B. barbus* (b) of same size (b 1: 11.5 cm S.L.; b 2: 3.9 cm; b 3: 2.4 cm).

PARVALBUMINS

Sarcoplasm from *B. barbus* white trunk muscle contains three electrophoretic components displaying the mobility of carp parvalbumins II, III and IV (Huriaux *et al.*, 1990). As the parvalbumin distribution varies according to fish size and the muscle-sample location, in the dorso-lateral muscle of *B. barbus*, any quantitative comparison requires suitable references. The anterior part of the *B. meridionalis* trunk muscle differed from the corresponding muscle portion from *B. barbus* specimens of the same size by the near absence of PA III and a lesser amount of PA II (Fig. 4 and Table I). In the case of the Asian barbels, the whole dorso-lateral muscle was used because of the small amount of tissue available. Only PA II and PA III isotypes appeared in the electrophoretic patterns of sarcoplasmic extracts from the four species, the stoichiometry of the two isotypes being species-specific except between *B. tetrazona* and *B. titteya*, where it appeared similar (Fig. 4 and Table I). This parvalbumin distribution is also very different from that observed in the *B. barbus* specimens of the same size: these exhibit little PA III but a noticeable amount of PA IV increasing with fish size.

IV. DISCUSSION

Various morphological and genetic data demonstrate the considerable species variability of the genus *Barbus* (Almaça, 1984; Berrebi *et al.*, 1990). This made it interesting to compare the electrophoretic properties of muscle proteins from various representatives of the genus. To this end, we chose to examine two large European barbels and four small Asian ones.

TABLE I. Percentage of parvalbumin isotypes in the dorso-lateral muscle from *B. meridionalis* (m), *B. conchoniuis* (c), *B. tetrazona* (te), *B. sachsi* (s) and *B. titteya* (ti) compared with *B. barbus* (b) of same size

Species	Standard length (cm)	PA I	PA II	PA III	PA IV
<i>B. meridionalis</i> *	12–14.5	—	18 ± 4	1.5 ± 1	80.5 ± 4
<i>B. barbus</i> †	11.5	—	41	15	44
<i>B. conchoniuis</i> ‡	4	—	40 ± 3	60 ± 3	—
<i>B. tetrazona</i> ‡	3	—	84 ± 3	16 ± 3	—
<i>B. sachsi</i> ‡	3	—	60 ± 4.5	40 ± 4.5	—
<i>B. titteya</i> ‡	2.4	—	83 ± 4	17 ± 4	—
<i>B. barbus</i> §	4	—	59	7	34
<i>B. barbus</i> §	3	—	67	4	29
<i>B. barbus</i> §	2.4	—	78	3	19

*Anterior part of the dorso-lateral muscle; average of 21 scans.

†Anterior part of the dorso-lateral muscle; from Huriaux *et al.* (1990).

‡Whole dorso-lateral muscle; average of 10 scans.

§Whole dorso-lateral muscle; from Huriaux *et al.* (1991a).

It is now well established that SDS-PAGE patterns of myosin light chains constitute a useful biochemical tool for characterizing fish species (Focant *et al.*, 1976; Huriaux & Focant, 1985; Rowleron *et al.*, 1985; Ochiai *et al.*, 1988, 1990; Martinez *et al.*, 1990a, b). These studies generally failed to reveal any phylogenetic correlations between electrophoretic patterns of the various marine or freshwater teleosts examined. Nevertheless, Ochiai *et al.* (1988, 1990), studying the physico-chemical and immunochemical properties of light chains from several marine teleosts, reported structural similarities within some families.

The myosin light chains of the barbel species examined in this work exhibited the electrophoretic pattern typical of fish. Light and heavy chains from *B. meridionalis* behaved in the same way as the corresponding chains from *B. barbus* (Huriaux *et al.*, 1990, 1991b), apart from the minor migration difference exhibited by LC₃ in SDS-PAGE. The present biochemical results are in agreement with morphological and genetic data which indicate that the two species are closely related. It is interesting to note that each European barbel showed a specific troponin-I in SDS-PAGE; this regulatory protein might be used as an additional criterion for species differentiation. As for the Asian barbels, several differences were found in the myosin subunits, not only compared with the reference species *B. barbus* but also within the group itself. *B. conchoniuis* LC₁ differed from all other electrophoretically similar LC₁ chains. On urea gels, the Asian barbels all exhibited a same LC₃ band, distinct from the European counterpart. On SDS gels, however, the myosin LC₃ of each Asian fish exhibited a specific pattern, again different from that of the corresponding *B. barbus* chain. As previously reported (Huriaux & Focant, 1985; Rowleron *et al.*, 1985; Ochiai *et al.*, 1990), LC₃ is the light chain showing the greatest variability on SDS-PAGE with an alkaline discontinuous buffer system. It thus appears to be the most characteristic of a species. In the present work, each barbel species could be identified according to the mobility of this light chain. This electrophoretic diversity may be due to a variable artefact affecting LC₃ mobility at

alkaline pH (Huriaux & Focant, 1978), but, as in the case of *B. conchoni* LC₁, it does seem to reflect structural differences between the LC₃ chains of the various barbel species. It seems that any size or age effect on the LC₃ mobility very likely can be ruled out; in the case of *B. barb*, this myosin light chain has been shown to be electrophoretically identical from the embryo to the adult stage (Focant *et al.*, 1992). Unlike LC₃, LC₂ was very similar in all species studied, suggesting it could be a protein that was highly conserved in the course of evolution (Focant *et al.*, 1976; Huriaux & Focant, 1985; Ochiai *et al.*, 1988, 1990). On the other hand, the electrophoretic patterns of the myosin heavy chains from the four Asian barbels were also species-specific although unrelated to the light-chain patterns, and distinct from those of the European barbels.

Parvalbumins appear as another biochemical criterion for distinguishing related fish species, especially since they present several isotypes in Cyprinidae. Interspecific comparisons, however, must be made with caution: the qualitative and quantitative distribution of isotypes have been shown to vary, in the dorso-lateral muscle of *B. barb*, with fish size, especially in the first developmental stages, and according to the myotomal location (Huriaux *et al.*, 1990, 1991a; Focant *et al.*, 1992). *B. meridionalis* parvalbumins were shown to resemble their *B. barb* counterparts qualitatively but to differ quantitatively. The Asian barbels, on the other hand, clearly diverge from the European species in that only two parvalbumin isotypes were detected. The stoichiometry of these isotypes differed from one species to another, except for *B. tetrazona* and *B. titteya*. The absence of a PA IV isotype does not seem to be imputable to the small size of the specimens: in the case of *B. barb*, PA IV was detected in the sarcoplasm of embryos 1 cm long (5 days post-fertilization). On the other hand, the variable stoichiometry of PA II and PA III among Asian fishes was not related to their size.

The PAGE patterns of myosin subunits and parvalbumin isotypes from white trunk muscle can thus be used to differentiate six species of the genus *Barbus*, even the very near species *B. barb* and *B. meridionalis*. On the whole, these biochemical results agree with the above-mentioned morphological, physiological, and genetic data: the two large European species appear closely related while the four small Asian species not only differ from the European fishes but also vary among themselves. On the basis of morphological features, *B. conchoni* and *B. sachsi* have sometimes been grouped in the genus *Puntius* while *B. tetrazona* and *B. titteya* were classified in the genus *Capoeta*. Frankel, however, pointed out that *B. tetrazona* and *B. conchoni* are closely related species, whereas a greater genetic distance exists between *B. tetrazona* and *B. sachsi* or *B. titteya* (Frankel, 1985; Frankel & Wilson, 1985). Our biochemical results fail to confirm these close relationships. We find no correlation between the characteristics of myosin light chains, myosin heavy chains, and parvalbumins among any of the Asian species studied here, this excludes any phylogenetic comparison.

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