

ESTERASE ISOZYMES PATTERN IN *PANGASIVS PANGASIVS* AND *P. SUTCHI*

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The electrophoretic banding pattern of esterase isozymes was examined in two fish species of Bangladesh. Samples were taken from both the flesh and intestine of *Pangasius sutchi* (Thai pangus) and *Pangasius pangasius* (Native pangus) for investigation. A maximum number of different esterase bands observed in the genus *Pangasius* were ten of which, eight different bands were observed in *P. sutchi* distributed in six loci (*viz.* Est.-2, Est.-3, Est.-5, Est.-6, Est.-7, Est.-8) and seven bands were found in *P. pangasius* distributed in seven loci (*viz.* Est.-2, Est.-3, Est.-4, Est.-5, Est.-6, Est.-7, Est.-8). Among these, Est.-5 was found to be polymorphic in *P. sutchi*. These isozymes showed a species specific, tissue specific and intensity variation in their expression. The electrophoretic patterns of esterase isozymes appear to be genetically controlled and are unaffected by physiological or environmental factors.⁽¹⁾ Electrophoretic pattern of isozymes in fish is species specific which could be used for the identification of fish species.^(2,4) During the last decade, large amount of genotype and allele frequency data has been obtained from a number of fish species primarily through the means of protein electrophoresis which enabled identification of many complex species.^(3,4) In this study, we observed variation in the pattern and intensity of esterase bands between *P. pangasius* and *P. sutchi*. *P. pangasius* is known to be more delicious and *P. sutchi* is first growing. By using the electrophoretic pattern of esterase isozymes we could possibly select for the best hybrid pangus that would be delicious, fast growing and could be cultured in the pond.⁽⁵⁾

Pangasius pangasius (Native pangus) and *Pangasius sutchi* (Thai pangus) were investigated to study flesh and intestine. Vertical polyacrylamide gel electrophoresis was used to verify the genotype of esterase isozymes in flesh and intestine following the method of Hames⁽⁶⁾ (with some modifications). Electrophoresis was carried out using a Tris-Borate-EDTA buffer, pH 8.9. α and β -naphthyl acetate was used as an esterase substrate.

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Substrate buffer pH 6.4 was made by mixing 50 ml of 0.2 M sodiumphosphate (monobasic) to 10 ml of 0.2 M sodium phosphate (dibasic) making final volume to 100 ml with distilled water. 0.042 g α -naphthyl acetate was dissolved in 1 ml acetone and 1 ml distilled water at 45°C. 0.042 g β -naphthyl acetate was dissolved in 2 ml acetone at 70°C. Finally, 2 ml α -naphthyl acetate solution and 2 ml β -naphthyl acetate solution were added to 100 ml of staining buffer. This solution was known as substrate mixture. Bands were scored from the stained gel. Relative mobilities were estimated from the commonly found band (Est-5^{1.0}). The esterase isozyme loci were assigned to an increase number based on the increasing mobility from the origin following Trebatoski and Craig.⁽⁷⁾

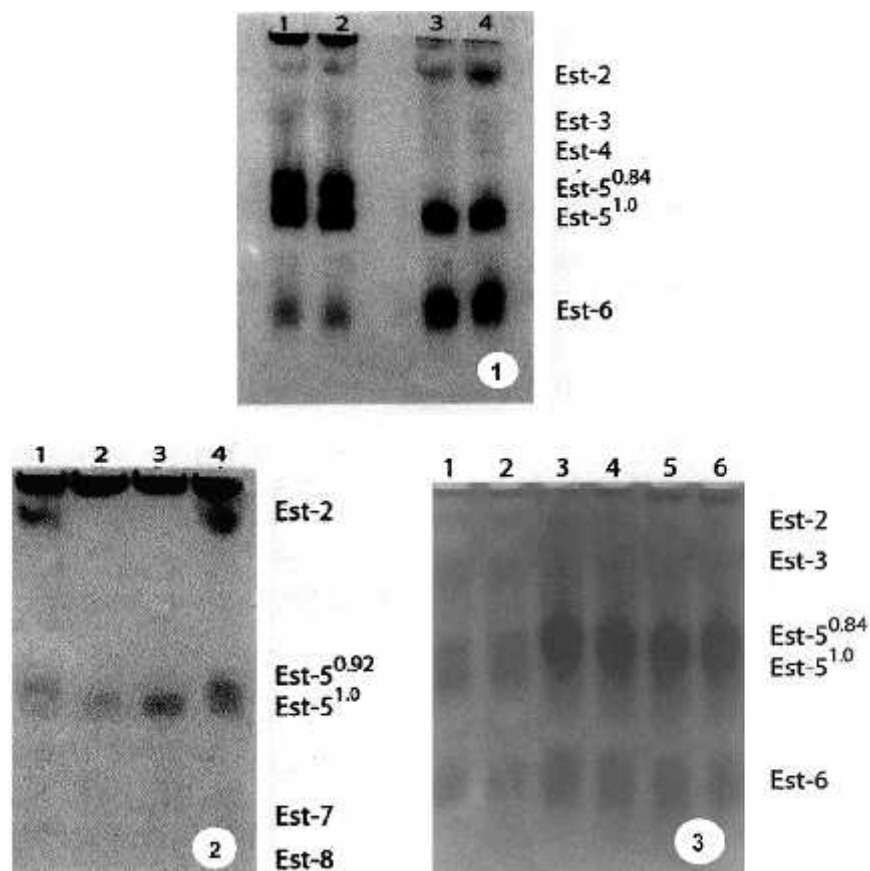
Table 1. The intensity variation of esterase isosymes bands in different tissues of *P. sutchi* and *P. pangasius*.

Relative mobility	Locus	<i>P. sutchi</i>		<i>P. pangasius</i>	
		Intestine	Flesh	Intestine	Flesh
0.18 ± 2	Est.-2	MS	DS	MS	-
0.44 ± 2	Est.-3	MDS	-	MDS	-
0.63 ± 2	Est.-4	-	-	FS	-
0.84 ± 2		DS	-	-	-
0.92 ± 2	Est.-5	-	MDS	-	-
1.0 ± 2		DS	MDS	DS	MDS
1.43 ± 2	Est.-6	MDS	-	DS	-
1.5 2 ± 2	Est.-7	-	FS	-	FS
1.75 ± 2	Est.-8	-	FS	-	FS

- = Absent; DS = Deep stained; MDS = Medium deep stained; MS = Medium stained; FS = Faint stained.

Variation was observed in the pattern and intensity of the esterase isozymes band between *P. sutchi* and *P. pangasius* as shown in Figs. 1, 2 and 3. Intensity of esterase bands differed from tissue to tissue and species to species.⁽⁸⁾ In Thai pangas, Est.-2 was stained deep in flesh (tail) but stained medium in intestine (hind parts) (Figs. 1 and 2). Est.-5^{0.84} and Est.-5^{1.0} was more intense in the middle and hind part of intestine of same species (Fig. 3). Species specific intensity was observed in the Est.-6 and Est.-5^{0.84}. In intestine, Est.-6 was stained medium-deep in *P. sutchi* and stained deep in *P. pangasius* (Fig. 1). The intensity variation of the bands is summarized in Table 1. Est.-2 and Est.-5 only were found in flesh of Thai pangas but it was not found in that of native pangas. Again, Est. 5^{0.84} was observed only in the intestine of Thai pangas. On the other hand, Est.-4 was only observed in the native pangas. Est.-5 was observed as a polymorphic locus in *P. sutchi* having three alleles. Two polymorphic loci were found in two species and four sub-species of the fish of the genus *Xiphophorus* (Poeciliidae) having eight alleles in the Est-2 and five alleles in the Est-3.⁽⁹⁾

Isozymes analysis by electrophoresis has a great significance in the field of fish genetics and fish technology particularly in taxonomy for identification of species and in breeding technology.^(2,5) In the present investigation, ten esterase bands were observed in flesh and intestine of *P. pangasius* and *P. sutchi*. All the ten bands were



Figs. 1 - 3: 1. Comparison of esterase isozymes loci in the intestine (posterior part) between *P. sutchi* (1, 2) and *P. pangasius* (3, 4). 2. Comparison of esterase isozymes loci in the flesh (tail region) between *P. sutchi* (1, 4) and *P. pangasius* (2, 3). 3. Comparison of esterase isozymes banding pattern among the fore-intestine (1, 2) mid-intestine (3, 4) and hind-intestine (5, 6) of *P. sutchi*.

not observed in the same tissue and in the same species. Maximum five different bands were found in the intestine of both species. Eight esterase bands were reported to find in *Oreochromis aureus* in brain, eye, heart, muscle and liver after electrophoretic analysis.⁽¹⁰⁾ In this study *P. pangasius*, *P. sutchi* and their hybrid from the natural source could be accurately identified by using the esterase pattern as a biochemical marker. The hybrid larvae of *Penaeus monodon* and *Penaeus esculentus*, was identified by using such biochemical marker.⁽⁵⁾ We need to continue

further such research on the above mentioned fish species and others to obtain concrete data that could be used for identification of fish species, selection of the best hybrid and above all technology - based improvement of fish farms.

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