Arch. Anim. Breed., 59, 435–444, 2016 www.arch-anim-breed.net/59/435/2016/ doi:10.5194/aab-59-435-2016 © Author(s) 2016. CC Attribution 3.0 License.





# Characterization of morphological and meristic traits and their variations between two different populations (wild and cultured) of *Cichlasoma festae*, a species native to tropical Ecuadorian rivers

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Received: 7 September 2016 - Revised: 13 October 2016 - Accepted: 18 October 2016 - Published: 31 October 2016

**Abstract.** This study was carried out to determine morphometric and meristic characteristics of two populations (wild and cultured) of *Cichlasoma festae* and to establish whether populations could be discriminated based on morphometric variability. Twenty-two morphometric and four meristic characters were used to test the hypothesis differentiation. Univariate analysis of variance (ANOVA) from 100 adult specimens showed significant differences (p < 0.05) for 21 standardized morphometric measurements out of 26 characters among the means of the wild and cultured *Cichlasoma festae* populations tested. Cross correlation amongst certain morphometric variables (i.e. body weight, total length, standard length, pre-ventral length, AC1, LC1 and P1) were medium-strong ( $r \ge 0.5$ ), while the remaining were weakly correlated (r < 0.5). The length–weight relationship parameter *b* and condition factor (K) values were respectively 2.21 and 1.97 (indicating allometric growth) for cultured fish groups and 2.86 and 4.07 (p < 0.05) for wild fish groups. The condition factor values were significantly different from each other and showed that feeding of cultured fish should be improved. Both groups were accurately separated (> 80 % success rate) by linear discriminant functions that included only four morphometric measures.

# 1 Introduction

In Ecuador, human communities in coastal as well as inland areas greatly depend on fishery for their incomes and as their source of animal protein (Espinosa-Lemus et al., 2009). Environmental degradation and habitat destruction have caused the decline in the production of fishery resources from the wild, which have diminished greatly (Ajah et al., 2006). Therefore, the domestication of certain fish species is necessary for intensive cultivation in captivity.

The morphometric study of fish is a powerful tool for characterizing strains and/or stocks of the same species which involves the detection of subtle variation in shape, independent of size. These examinations require exact measurements and counts of fin ray elements. For morphological study, morphometric (referring to measurable structures such as fin length, head length, eye diameter, or ratios between such measurements) and meristic (including almost any countable structure, such as fin rays, scales and gill rakers) characters are used. The morphometric characters are classified into genetically (narrow range) controlled, intermediate (moderate range) and environmentally (vast range) controlled characters (Johal et al., 1994). Despite the advent of techniques which directly examine biochemical or molecular genetic variation, the morphometric or meristic methods continue to play an important role in stock identification even today (Swain and Foote, 1999).

The phenotypic plasticity of fish is very high, with greater variances in morphological traits both within and between populations than any other vertebrates. The cause of variation in the morphometric and meristic characters can be partly attributed to intraspecific variability, which is under the influence of environmental parameters (Wimberger, 1992). Fish are very sensitive to environmental changes and quickly adapt by changing necessary morphometric character (Cabral et al., 2003; Hossain et al., 2010). Morphometric variation between stocks may be applicable for studying short-term environmentally induced variation (Pinheiro et al., 2005). In addition, while both morphometric and meristic characters respond to changes in environmental factors, their responses are different in some situations and can differ from species to species. Finally, is important to farmers to know the differences between cultured and wild fish of different species; this could lead them to understand the chemical, physical, nutritional and sensorial profiles of the wild animal and try to reproduce these in their cultured products (Orban et al., 2003).

In Ecuador, fisheries contribute 7% to the total supply of animal protein, estimated at 391 700 t catches made by capture fisheries in 2011 (FAO, 2011). These catches are made by artisanal fishermen in areas of rivers, lakes, ponds, lagoons, gorges and dams. This activity is performed throughout the year in areas of rivers (Muñoz et al., 2014) or between May and January in other inland areas. Cichlasoma *festae*, among the freshwater fish (Boulenger, 1899), is a teleost fish (Luna-Figueroa, 2000), native to the continental South America, with a high presence in Ecuador. It is among the nine commercially important species that inhabit the inland waters of Ecuador, Colombia and Peru (Revelo and Elias, 2004). It can be found in rivers, lakes, ponds and dams (Pacheco and Chicaiza, 2008) and noted for its white meat, excellent taste and high acceptance in local cuisine (Barnhill et al., 1974).

In order to produce and preserve this native species, the state administration created the Cachari Experimental Station, located in Babahoyo in the province of Los Ríos, where a conservation programme for native species is currently being developed by the Subsecretaría de Acuacultura of Ministerio de Agricultura, Ganadería, Acuacultura y Pesca (MA-GAP). At this experimental station, fingerlings were produced for distribution to farmers and to repopulate the rivers. According to MAGAP, the cultivation of *Cichlasoma festae* is becoming more and more popular due to its good growth rate, fecundity, ease of manipulation, ability to grow under suboptimal environmental conditions, disease resistance and good consumer acceptance.

Understanding the morphometrics of the fish species will enhance the development of cost-effective aquaculture protocols and thus increase in productivity. Although comparisons of the morphology between cultured and wild fishes from several species have already been carried out by a number of authors (Swain et al., 1991; Ponton and Mérigoux, 2000; Solem et al., 2006; Solomon et al., 2015), there is a lack of information on the level of this variation for most tropical fish species. Difference among cultured and wild *Cichlasoma festae* stocks based on morphological characters have not yet been studied and, to the best of our knowledge, this is the first such study that has focused on examining the extent of their morphological variations in cultured and wild environments.

Since this information is vital for the proper management of the fisheries and for optimum utilization of the resources, the aim of the current research was to assess the morphological and meristic traits of *Cichlasoma festae* caught in different habitats (cultured and wild). This will help in planning further breeding and conservation strategies for this fish and improving productivity.

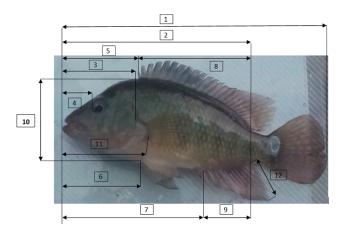
# 2 Material and methods

## 2.1 Data and sampling

The study included three areas of the Babahoyo River and a fish farm in the province of Los Ríos (Ecuador). The climate of the area is tropical with an average temperature of 25 °C, an annual rainfall of 2400 mm and a relative humidity of 82 %. The salinity of water, both in the river and the fish farm, did not exceed 0.1 %; the pH was between 7.0 and 7.29; the temperature ranged between 19.7 °C in the river and 24.7 °C in cultured fish; and dissolved oxygen was between 6.8 and 8.9 mg L<sup>-1</sup> in the river and fish farm, respectively. The conductivity values were about 145 mS cm<sup>-1</sup>.

One hundred matured fish samples (following the rules described by Frost and Kipling, 1980; Chávez-Lomelí et al., 1988; Konings, 1989) of Cichlasoma festae, comprising 50 individuals from natural habitat (wild population) and 50 from a cultured environment (private fish farms, cultured stock), were collected at dawn over the month of May 2016 with the help of standard fishing gears such as cast and hand nets. Since males and females could not be differentiated morphologically, sexing of the sampled fish was not carried out. Specimen collection was performed weekly by purchasing representative samples of the two selected populations from local fishermen (wild fish) or a fish farm (cultured fish). Wild fish were caught from three different locations within their natural geographic distributions in Babahoyo River (Los Ríos province, Ecuador). Cultured fish were collected from the fish farm. Directly after catching, the fish were placed at the same time in a mixture of 40 L of ice and 40 L of water  $(0.8 \,^{\circ}\text{C})$  until their apparent stunning  $(20 \,\text{min})$ was over. After confirmation of their death, the fish were identified and weighed, and then morphometric measurements and meristic counts were performed.

The study was carried out according to Ecuadorian national recommendations for the management of fish, taking into consideration the rules on animal welfare.



**Figure 1.** The morphometric measurements registered in each analysed organism (source: own elaboration). 1: total length (TL); 2: standard length (SL); 3: head length (HL); 4: pre-orbital length (Pre-OL); 5: pre-dorsal length (Pre-DL); 6: pre-ventral length (Pre-VL); 7: pre-anal length (Pre-AL); 8: dorsal fin length (DFL); 9: pharyngeal bone length (PhBL); 10: maximum height body (MaxBH); 11: pectoral fin length (PFL); 12: anal fin length (AFL).

## 2.2 Body measurements

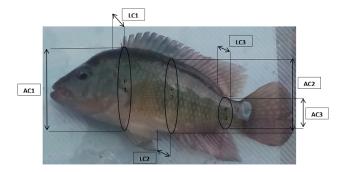
Lineal morphometric measurements were taken on the left side of fish, by the same person in order to minimize artificial error, and most of the morphometric characters were measured following the conventional method described by Morales et al. (1998) and Diodatti et al. (2008). The fish were measured using a measuring board, measuring tape and digital callipers graduated in millimetres and then weighed with an electronic weighing balance up to the nearest 0.1 g (Figs. 1 and 2). Meristic characteristics were examined according to Froese and Pauly (2007). A total of 26 body measurements were used, including 21 morphometric variables and 4 meristic variables (Table 1).

# 2.3 Fulton condition factor (K)

The Fulton condition factor (*K*), which is defined as the well-being of the fish, was calculated. *K* is a useful index for monitoring of feeding intensity, age and growth rates. The *K* was calculated with the following equation:  $K = (100 \times BW)/SL^3$ , where BW refers to body weight of fish in grams and SL is the standard length of fish in centimetres.

## 2.4 Length-weight relationship

Length–weight relationships were calculated using the allometric regression analysis (Sasi and Berber, 2012). Length– weight was expressed as  $BW = a \times SL^b$ , the logarithm transformation of which gives the linear equation  $\log BW = a + (b \times \log SL)$ , where BW refers to body weight of fish in grams, SL is the standard length of fish in centimetres, *a* is the constant for the initial growth index and *b* is the growth



**Figure 2.** The morphometric measurements registered in each analysed organism (source: own elaboration). AC1: body depth at the first ray of the dorsal fin; AC2: body depth at the level of the first ray of the anal fin; AC3: body depth at the level of the first radius of the caudal fin; P1: body perimeter of the body at the level of the first ray of the dorsal fin; P2: body perimeter at the level of the first radius of the anal fin; P3: body perimeter at the level of the last ray of the dorsal fin; LC1: head width; LC2: trunk width; LC3: tail width.

coefficient. Constants a and b represent the point at which the regression line intercepts the y axis and the slope of the regression line, respectively.

## 2.5 Statistical analyses

All statistical analyses were performed using SAS University Edition 3.5 (SAS Institute, Cary, NC). Each collection site was considered a priori as a discrete group. To evaluate whether the data have equal variances, a Bartlett test was done prior to further analyses. Means, standard deviation (SD) and coefficient of variation (CV %) were recorded for each population.

The morphometric (continuous) and meristic (discrete) data were analysed separately. Since meristic characters are independent of size and did not change during growth (Turan et al., 2006), the raw data were used in analysis. However, to avoid possible biases produced by size effects on the morphometric variables, all morphometric characters were standardized by the following equation (Elliott et al., 1995):  $M_{\rm adj} = M(L_{\rm s}/L_{\rm o})^b$ , where M is the original morphometric measurement,  $M_{adj}$  the size adjusted measurement,  $L_0$  the standard length of fish and  $L_s$  the overall mean of standard length for all fish from all samples for each variable. The parameter b was estimated for each character from the observed data as the slope of the regression of  $\log M$  on  $\log L_0$ , using all specimens. This method normalizes the individuals in a sample to a single, arbitrary size, common to all samples and, at the same time, maintains the individual variation (Tudela, 1999). It has been successfully used by many researchers in recent years (Ibañez-Aguirre and Lleonart, 1996; Salini et al., 2004; Turan et al., 2006). The efficiency of the size-adjustment transformations was assessed by testing the significance of the correlation between a transformed variable and the SL.

Tab	le	1.	Body	measurements.
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Trait		Description			
Morphometric variables					
Body weight	BW	Measured as total weight including gut and gonads			
Total length	TL	Measured from the middle of the upper lip of the mouth to the caudal end of the caudal fin			
Standard length	SL	Measured between the central portion of the upper lip of the mouth and the base of the caudal fin			
Head length	HL	Distance between the most cranial point of the upper lip of the mouth and the rear end of the operculum			
Pre-orbital length	Pre-OL	Distance between the most cranial point of the lower lip of the mouth and the cranial edge of the eye			
Pre-dorsal length	Pre-DL	Distance between the most cranial point of the lower lip and the start of the first dorsal spine			
Pre-ventral length	Pre-VL	Distance between the most cranial point of the lower lip and the start of the first spine of the ventral fin			
Pre-anal length	Pre-AL	Distance between the most cranial point of the lower lip and the beginning of the anal orifice			
Dorsal fin length	DFL	Distance from the most cranial point of the base of the fin to the caudal end of the dorsal fin			
Pharyngeal bone length	PhBL	Distance from the most cranial point of the base of the fin to the caudal end of the anal fin			
Maximum height body	MaxBH	Distance between the most cranial point of the pectoral fin and the lateral line			
Pectoral fin length	PFL	Distance between the base point cranial flap to the rear end of greater radii			
Anal fin length	AFL	Distance from the most cranial point of the base of the fin to the end of anal fin			
Body depth	AC1	Measured with a calliper, at the first ray of the dorsal fin			
Body depth	AC2	Measured with a ruler, at the level of the first ray of the anal fin			
Body depth	AC3	Measured with a calliper, at the level of the first radius of the caudal fin			
Head width	LC1	Distance from side to side at the level of the flow side of the head			
Trunk width	LC2	Distance from side to side at the level of the most cranial point of the anal fin side			
Tail width	LC3	Distance from side to side at the level of the last thorn on the back side			
Body perimeter	P1	Measured with measuring tape, at the level of the first ray of the dorsal fin			
Body perimeter	P2	Measured with measuring tape, at the level of the first radius of the anal fin			
Body perimeter	P3	Measured with measuring tape, at the level of the last ray of the dorsal fin			
Meristic variables					
Dorsal fin rays	DFR	Count of thorns has the dorsal fin from start to finish			
Radius dorsal fin	RDF	Count of cartilage found in the space between thorns from start to finish			
Anal fin rays	AFR	Count of thorns has the anal fin from start to finish			
Radius anal fin	RAF	Count of cartilage found in the space between thorns from start to finish			

Size-adjusted morphometric data and meristic characters were compared by univariate analysis of variance (ANOVA procedure) and Kruskal–Wallis test (NPAR1WAY procedure), respectively, using the group (cultured or wild) as the fixed effect. In addition, the DISCRIM procedure was used to perform a canonical discriminant analysis of both sizeadjusted morphometric data and meristic characters. The variables that would be included as predictors in the canonical discriminant function were previously selected with the STEPDISC procedure. The probabilities to enter and to stay in the model were both set at 0.05.

# 3 Results

## 3.1 Morphometric characters

Morphometric and meristic traits mean values of *Cichlasoma festae* from cultured and wild specimens are shown in Table 2.

Among the morphometric characters, the most used are the body weight (BW), total length (TL), standard length (SL) and head length (HL). The mean BW of *Cichlasoma festae* from all data ranged from 55.8 to 152.0 g with a mean value of 90.45  $\pm$  18.2 g. The value of TL ranged between 12.5 and 25.0 cm with a mean value of 18.27  $\pm$  1.75 cm, SL ranged between 9.8 and 19.0 cm with a mean value of 14.14  $\pm$  1.58 cm, and HL ranged between 4.4 and 6.5 cm with a mean value of 5.35  $\pm$  0.48 cm. Cultured fish were larger than those coming from a natural habitat, so weight and most morphometric variables showed higher mean values, except for LC3. The mean Pre-OL, AFL, LC2, LC3, AFR and RAF of *Cichla*-

#### M. A. González et al.: Characterization of morphological and meristic traits

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Table	2. Descriptive statistics	of the morphometric an	d meristic characters	(original data) fro	om Cichlasoma festae.

	All data		Cultured		Wild		
	Mean	SD	CV %	Mean	CV %	Mean	CV %
Body weight (g)	90.45	18.16	20.07	101.84 <sup>a</sup>	16.43	79.06 <sup>b</sup>	13.94
Fulton condition factor, K	3.32	0.92	27.80	3.01 <sup>a</sup>	22.41	3.62 <sup>b</sup>	28.65
Total length (cm)	18.27	1.75	9.60	19.40 <sup>a</sup>	7.17	17.14 <sup>b</sup>	7.54
Standard length (cm)	14.14	1.58	11.18	15.12 <sup>a</sup>	8.00	13.15 <sup>b</sup>	9.64
Head length (cm)	5.35	0.48	9.03	5.57 <sup>a</sup>	8.52	5.14 <sup>b</sup>	7.66
Pre-orbital length (cm)	2.18	0.44	20.07	2.27 <sup>a</sup>	23.43	2.10 <sup>a</sup>	14.40
Pre-dorsal length (cm)	5.37	0.67	12.44	5.69 <sup>a</sup>	7.81	5.05 <sup>b</sup>	13.93
Pre-ventral length (cm)	5.83	0.57	9.71	6.20 <sup>a</sup>	7.94	5.45 <sup>b</sup>	6.11
Pre-anal length (cm)	9.05	0.92	10.22	9.28 <sup>a</sup>	11.98	8.83 <sup>b</sup>	7.08
Pectoral fin length(cm)	8.01	0.88	10.96	8.25 <sup>a</sup>	9.91	7.78 <sup>b</sup>	11.32
Pharyngeal bone length (cm)	3.40	0.43	12.67	3.52 <sup>a</sup>	12.10	3.27 <sup>b</sup>	12.25
Maximum body height (cm)	3.90	0.60	15.26	4.16 <sup>a</sup>	12.91	3.64 <sup>b</sup>	14.82
Dorsal fin length (cm)	5.99	0.69	11.52	6.40 <sup>a</sup>	10.41	5.58 <sup>b</sup>	7.53
Anal fin length (cm)	4.58	1.02	22.27	4.78 <sup>a</sup>	22.15	4.39 <sup>a</sup>	21.66
AC1 (cm)	5.46	0.45	8.24	5.75 <sup>a</sup>	7.04	5.16 <sup>b</sup>	4.85
AC2 (cm)	4.93	0.40	8.22	5.18 <sup>a</sup>	6.53	4.67 <sup>b</sup>	6.27
AC3 (cm)	1.94	0.25	12.86	2.03 <sup>a</sup>	9.73	1.84 <sup>b</sup>	14.05
LC1 (cm)	2.31	0.27	11.77	2.41 <sup>a</sup>	8.79	2.21 <sup>b</sup>	13.06
LC2 (cm)	1.56	0.36	23.03	1.57 <sup>a</sup>	27.45	1.55 <sup>a</sup>	17.73
LC3 (cm)	0.71	0.23	32.72	0.69 <sup>a</sup>	24.43	0.74 <sup>a</sup>	38.49
P1 (cm)	13.24	1.00	7.55	13.72 <sup>a</sup>	8.11	12.76 <sup>b</sup>	4.43
P2 (cm)	11.36	0.68	6.01	11.78 <sup>a</sup>	5.01	10.95 <sup>b</sup>	4.51
P3 (cm)	4.73	0.44	9.41	4.87 <sup>a</sup>	9.63	4.58 <sup>b</sup>	8.11
Dorsal fin rays	27.04	0.98	3.64	27.32 <sup>a</sup>	3.09	26.76 <sup>b</sup>	3.89
Radius dorsal fin	26.12	1.54	5.89	26.52 <sup>a</sup>	7.07	25.72 <sup>b</sup>	3.77
Anal fin rays	13.70	0.76	5.54	13.80 <sup>a</sup>	5.86	13.60 <sup>a</sup>	5.15
Radius anal fin	12.82	0.85	6.59	12.80 <sup>a</sup>	6.31	12.84 <sup>a</sup>	6.92

AC1: body depth at the first ray of the dorsal fin; AC2: body depth at the level of the first ray of the anal fin; AC3: body depth at the level of the first radius of the caudal fin; LC1: head width between the right and left point at the level of the flow side of the head; LC2: trunk width between the right and left at the level of the most cranial point of the anal fin side; LC3: tail width between the right and left at the level of the first ray of the dorsal fin; P2: body perimeter at the level of the first ray of

the dorsal fin.

<sup>a,b</sup> Within a row, means without a common superscript are different (p < 0.05).

*soma festae* from the two populations were not significantly different from each other.

The TL, HL, Pre-VL, AC1, AC2, P1, P2 and P3 showed a coefficient of variation lower than 10%; SL, Pre-DL and Pre-AL, PFL, PhBL, MaxBH, DFL, AC3 and LC1 showed a coefficient of variation between 10 and 20%; and the BW, Pre-OL, AFL, LC2 and LC3 showed coefficients of variation greater than 20%. The coefficients of variation of different morphometric characters were not significantly (p < 0.05) different between populations, except for Pre-OL, Pre-DL, Pre-AL, AC3, LC2 and LC3.

The meristic characters showed mean values of  $27.04 \pm 1.0$ ,  $26.12 \pm 1.5$ ,  $13.70 \pm 0.8$  and  $12.82 \pm 0.9$  for DFR, RDF, AFR and RAF, respectively, with no significant difference (p < 0.05) among populations. The

coefficients of variation were very low (< 7%) and similar between populations.

Dorsal fin rays (DFR) and radius dorsal fin (RDF) ranged from 24 to 28, with most in the range of 27–28 (82%) and 26–27 (70%), respectively. In anal fin rays (AFR) and radius dorsal fin (RDF) ranged from 11 to 15, presenting most of the 13–14 (82%) and 12–13 (76%) respectively. Significant differences (p < 0.05) between cultured and wild were found (data not shown). The range of the dorsal fin characters was higher for wild (W) than cultured (C) fishes, although 27 (56% for C and 36% for W) and 26 (48% for C and 18% for W) were the most frequent classes for DFR and RDF, respectively. Conversely, the range for anal fin characters (AFR and RAF) was higher for C (12–15 and 11–14) than W (13–15 and 12–14), and the most frequent class differed between populations (14 = 44 % for C and 13 = 52 % for W, and 13 = 52 % for C and 12 = 44 % for W).

The mean BW / SL ratio was  $6.39 \pm 0.98$ . HL SL, MaxBH, body depth (AC1, AC2, AC3), body width (LC1, LC2, LC3) and body perimeter (P1, P2, P3) represented 38 %, 28 %, 39 to 14 %, 16 to 5 % and 94 to 34 %, respectively. The ratios of TL, Pre–VL, Pre–VL, DFL, AC1, AC2, P1 and P2 with SL showed a coefficient of variation lower than 10 %; ratios BW, HL, Pre–DL, Pre–AL, PFL, PhBL, MaxBH, AC3, LC1 and P3 with SL showed a coefficient of variation between 10 and 20 %, while ratios of Pre–OL, AFL, LC2 and LC3 showed coefficients of variation greater than 20 %. In general, the coefficients of variation of the indices are slightly lower than those recorded in the corresponding morphological measurements.

Among populations, the BW / SL was significantly higher (p < 0.05) in the cultured population, while the ratios of HL / SL, Pre-AL / SL, PFL / SL, PhBL / SL, AC1 / SL, AC2 / SL, LC1 / SL, LC2 / SL, LC3 / SL, P1 / SL, P2 / SL and P3 / SL were significantly higher (p < 0.05) in the wild population. Based on these relationships, wild fish were proportionately more profound at the cranial level than cultured, without significant differences (p > 0.05) at the caudal level. Likewise, at the cranial and caudal levels, they were proportionally wider. All of this caused the body perimeter / SL ratios, both at cranial and caudal levels significantly, to be lower (p < 0.05) in cultured fish.

After standardizing according to Elliot et al. (1995), the mean values of BW, TL, SL and HL were  $90.38 \pm 1.87$ ,  $18.32 \pm 0.13$ ,  $14.14 \pm 0.16$  and  $5.36 \pm 0.06$  cm, respectively. The habitat had a significant effect (p < 0.05) in some of the morphometric characters evaluated. BW, TL, SL, HL, Pre-VL, DFL, AC1, AC2, AC3 and P2 were significantly higher (p < 0.05) in cultured specimens. AFL, LC1 and P1 tended to be higher (p < 0.1) in the cultured population.

## 3.2 Fulton condition factor

The mean value of the condition factor *K* was  $3.32 \pm 0.9$  (Table 2) for the original data set, with mean values of 3.01 and 3.62 for cultured and wild populations, respectively. The coefficient of variation was high (27.8%). Once the data were adjusted to SL (Elliot et al., 1995), the mean value of the condition factor *K* was  $3.47 \pm 0.15$ , with significantly higher values (p < 0.001) in the wild than in the cultured population, where the values were  $4.07 \pm 0.24$  and  $2.86 \pm 0.12$ , respectively.

#### 3.3 Length-weight relationship

The parameter *b* of the fish studied ranged from a minimum of 1.57 to a maximum of 2.46, with a mean value of  $2.096 \pm 0.078$ , and with a slightly higher average value in cultured fish when compared with wild fish (2.21 vs. 1.97).

**Table 3.** Fisher's discriminant functions for morphometric variables.

	Cultured	Wild
Constant	-441.77	-375.03
SL	19.77	17.80
Pre-VL	41.88	38.83
AC2	57.95	54.36
AFL	8.19	7.47

SL: standard length; Pre-VL: pre-ventral length; AC2: body depth at the level of the first ray of the anal fin; AFL: anal fin length.

#### 3.4 Relationships between morphometric characters

The morphometric relationships between numerous body parts of fish can be used to determine possible difference between separate unit stocks of the same species (King, 2007). Several significant (p < 0.05) positive correlations were found between the morphometric and meristic characters of the two populations (data not shown). Most correlation coefficients were between 0.3 and 0.5. The results reveal that the size effect was almost entirely eliminated in the populations during analysis as there were no significant correlations between TL and SL, with most of the remaining parameters measured with the analysed characters. Meristic characters, except for RDF, are not significantly related (p > 0.05) to each other or other morphometric characters.

# 3.5 Discriminant analysis

Four morphometric variables (SL, Pre-VL, AC2, AF1) out of 23 were selected as predictors in the canonical discriminant analysis (Table 3). Wilks' lambda (0.39; p < 0.001) indicated that the data were appropriate for discriminant analysis, whereas the eigenvalue (1.54) and canonical correlation (0.78) showed that the canonical function had very good discrimination ability.

The Mahalanobis squared distance between the cultured and wild populations was 6.03, and the F test of the distance was highly significant (p < 0.001). SL, followed at some distance by Pre-VL, AC2 and AFL, had the greatest discriminating ability and the highest correlation value with the canonical discriminant function, according to the standardized canonical coefficients and the pooled within-canonical structure, respectively. Fisher's linear discriminant functions are shown in Table 2. In the original classification matrices, eight cases were misclassified in the cultured group and four cases were misclassified in the wild group. In cross-validated classification matrices, nine cases were misclassified in the cultured group and seven cases were misclassified in the wild group. As a result, 88 and 84 % of the original grouped cases were classified correctly in the original and cross-validated classification matrices, respectively.

Regarding meristic variables, the only RDF was selected as predictor and, despite the Wilks' lambda statistical significance (p < 0.01), the eigenvalue and the canonical correlation were very low (0.09 and 0.29, respectively). The obtained Fisher's linear discriminant functions correctly classified 61 and 58 % of the original grouped cases in the original and cross-validated classification matrices, respectively.

# 4 Discussion

#### 4.1 Morphometric characters

According to Turan et al. (2006), the introduction and domestication of a fish species (especially those from the wild) leads to high adaptation to a wide range of geographical locations, which leads to phenotypic variations with respect to the pure stock (strains) of the brood stock. In order to know the ecological variation and to evaluate morphological differences between wild and cultured fish of the same species, different authors have used morphometric and meristic variables (Narváez et al., 2005; Fagbuaro et al., 2015; Solomon et al., 2015) to quantify biological variation and identify and explain adaptive processes of different populations of the same species.

On the basis of the classification of Negi and Nautiyal (2002), of the morphological characters studies from *Barilius bendelisis* and *Barilius vagra*, 12 characters were genetically controlled, 8 characters were intermediate and 7 characters were environmentally controlled. Twenty-one characters have been studied in percentage of standard fish length, from which seven characters were genetically controlled, nine characters were intermediate and five characters were environmentally controlled.

In the current study, it has been observed that the meristic counts did not change with increasing or decreasing body weight and length of the fish. Similar variations in meristic characters were reported in many fishes such as *Nematalosa nasus* (Al-Hassan, 1987), *Pseudobagrus ichikawai* (Watanabe, 1998), *Pterophyllum scalare* (Bibi-Koshy et al., 2008), *Garra gotyla gotyla* (Gray) (Brraich and Akhter, 2015).

This study recorded significant differences (p < 0.05) between populations in 11 morphometric parameters, in agreement with Fagbuaro et al. (2005) and Solomon et al. (2015). Barriga-Sosa et al. (2004), after analysing morphometric characters in natural and domesticated populations of Nile tilapia (*Oreochromis niloticus*), reported morphological differences among these populations. Likewise, Narváez et al. (2005) found significant differences between the two populations (wild and cultured) of *Oreochromis niloticus* in northern Colombia; differences were attributed to food, environmental conditions and the type of habitat (wild and cultured). However, in the present study, not all meristic characters registered showed significant differences between populations, in contrast to the results obtained by Solomon et al. (2005) in *Clarias gariepinus*. The discrepancy between results could be attributed to the characters studied in each work.

In the present study, TL / SL and DFL / SL ratios were not significantly (p > 0.05) different between cultured and wild specimens, in contrast to results obtained by El-Zaeem et al. (2012). While there is overlap between the two works in the differences between populations (cultured and wild) in the ratio between the standard length and depth and width of the body. These authors point out that the highest mean value of TL / SL in Nile tilapia (Oreochromis niloticus) was recorded by cultured population and differed significantly (P < 0.05) from that of the wild population. Also, the mean value of HL / AC1 ratio was not significantly different between populations (wild and cultured), in contrast to the results offered by Narváez et al. (2005), who observed that domesticated individuals were characterized by sharper heads than those of naturalized fish. Solomon et al. (2015) recorded significant differences in the ratio HL / SL in wild (23.7) and cultured (26.6) populations of C. gariepinus. Similarly, Vreven et al. (1998) and Barriga-Sosa et al. (2004) indicated that the biggest differences between wild and cultured populations were presented at the head. The value of this relationship and other relationships between morphometric characters is closely linked to the species, so it is not surprising that differences can be registered between studies. Thus, Van der Bank et al. (1989) reported mean values from 0.29 to 0.34 for HL / SL and 0.31 to 0.45 for body depth/SL in fifteen cichlid fish species endemic to southern Africa, whereas in our study means the values for these ratios were 0.38 and 0.39, respectively. Brraich and Akter (2015) in Garra gotyla gotyla (Gray) recorded mean values of 0.27 and 0.18, respectively. According Vreven et al. (1998) the confinement of domesticated fish affects their growth rate, without allowing elongate the body, which would result in a higher K value. Contrary to this, in our work the value of K is higher in wild specimens.

#### 4.2 Fulton condition factor (*K*)

Condition factor is a useful index for the monitoring of feeding intensity, age and growth rates in fish (Oni et al., 1983). It is strongly influenced by both biotic and abiotic environmental conditions and can be used as an index to assess the status of the aquatic ecosystem in which fish live.

The condition factor values of *Cichlasoma festae* from the current study (3.32) were comparable to those registered by Chukwuemeka et al. (2014) in *Tilapia aurea, Tilapia galilaea* and *Auchenoglanis occidentalis* and lower than those reported by Anene (2005) in four cichlid fish (4.9). However, Fagbuaro et al. (2015) recorded significantly lower values (0.68) in *Clarias gariepinus* fish. The correlation coefficients between the factor *K* and the total length or standard length are negative (-0.488 and -0.774 for cultured fish and -0.557 and -0.873 for wild fish, respectively) and statistically significant (p < 0.01 and 0.001), indicating that a shortened factor occurs with increasing size of the fish. These re-

sults are consistent with those obtained in four cichlid species by Anene (2005), who registered a significant and progressive decrease (p < 0.05) between the size range of 120 and 150 mm. Sasi and Berber (2012) recorded increases in condition factor until the age of 5 years (from 1.6 to 2.5) and a drop below.

In disagreement with Fagbuaro et al. (2015), the condition factor K was higher in the wild population. This implies that the fish from the cultured population may not have been fed to the required level.

# 4.3 Length-weight relationship

In the present study, the length–weight relationship parameter *b* is lower than in many studies (Abdallah, 2002; Bayhan et al., 2008; Sasi and Berber, 2012) and close (2.27-2.46)to that obtained by Fagbuaro et al. (2015), although it is located in the range of values (1.51-3.49) indicated by Bok et al. (2011). This study shows that the fish from both the cultured and wild fish population have exhibited no isometric relative growth, which does not maintain their specific body shape throughout their life. These results also showed that both wild and cultured habitats do not provide enough food to maintain an isometric growth.

In contrast to the results obtained by Fagbuaro et al. (2015) (2.27 for farmed fish and 2.46 for wild fish), in the present study the parameter b was higher in the cultured population.

#### 4.4 Correlation among morphometric variables

Out of 26 characters, 2 characters show high values of correlation coefficient and 24 characters show moderate to low correlation coefficient. In Cichlasoma festae, BW was found to be the most correlated part. In general, the correlation coefficients between morphological variables were slightly higher among wild fish, and clearly lower than those recorded by Brraich and Akhter (2015) in Garra gotyla gotyla. Chukwuemeka et al. (2014) recorded correlation coefficients between live weight and standard length of 0.76 to 0.94 in Tilapia galilaea, Tilapia aurea and Auchenoglanis occidentalis from Tagwai Lake (Nigeria). The variations observed in correlation coefficients of the morphometric and meristic data for wild and cultured Cichlasoma festae, aligned with the results obtained by Solomon et al. (2015), could be strongly linked to feeding pattern, environmental conditions and genetic variability. Also, there is sufficient evidence to prove the influence of habitat on fish morphology (Turan et al., 2006).

#### 4.5 Discriminant analysis

Canonical discriminant analysis demonstrated a clear influence of origin in the morphometric variables and a low effect in the meristic characters measured in the present work. The fact that only four morphometric variables were needed to separate the two groups suggests that Fisher's linear discriminant could be useful to identify the origin of stocks on a commercial basis. However, Van der Bank et al. (1989) attribute less value to the morphologic variables than to meristic counts in the differentiation of populations of the same species. The meristic counts showed a very low variability and overlapped broadly, showing no divergence among the populations, in agreement with several authors (Gacitúa et al., 2008; El-Zaeem et al., 2012; Solomon et al., 2015). These characters, due to their relative stability, cannot give the necessary variability in measurements which is essential for multivariate analysis and stock discrimination studies.

Although the causes of morphological differences between populations are often quite difficult to explain, the morphometric differences between the cultured and wild *Cichlasoma festae* could have been linked to environmental factors; furthermore, breeding over several years may have diluted the initial gene pool of the domesticated fish, leading to genetic variation (translating to morphological differences) (Solomon et al., 2015).

# 5 Conclusions

Our results show that the rearing system significantly influences most of the analysed morphometric and meristic characteristics of the two populations (wild and cultured) of Cichlasoma festae. Twenty-two morphometric and four meristic characters were used to test the hypothesis differentiation. Univariate analysis of variance showed significant differences for 21 standardized morphometric measurements out of 26 characters among the means of the wild and cultured populations tested. The condition factor values were significantly different from each other and showed that feeding could be improved in the farms. Both groups were accurately separated by linear discriminant functions that included only four morphometric measures. These results are of vital importance for the Ecuadorian population because they will allow for planning of further breeding and conservation strategies for this native fish and improving productivity.

## 6 Data availability

The original data are available upon request to the corresponding author.

Edited by: M. Mielenz Reviewed by: J. Rivas and one anonymous referee

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