Stock delineation in *Clupisoma garua* (Hamilton, 1822) populations of Ganga riverine system using truss network analysis

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ABSTRACT

Stock identification of fish plays a significant role in management of fishery resources. It is considered as the precursor to the species conservation and imperative for declining fish populations. The stock structure of *Clupisoma garua* (Hamilton, 1822) populations of Ganga riverine system was studied using truss network analysis. A total of 306 fish specimens obtained from river Ganga in Allahabad, Farakka and Triveni were used to construct the truss network. The truss variables were subjected to principal component analysis (PCA) in which the first component explained 46.71% and the second 14.66% representing body depth and caudal regions of fish, respectively. Bivariate plots of PCA and discriminant function (DF) analysis revealed that it is a single continuous stock in the study area of the river Ganga and the misclassification of DF indicated only a marginal morphological difference. The results indicated the need for a uniform management strategy appropriate for population of *C. garua* in the river Ganga.

Keywords: Catfish, Conservation, Ganga River, PCA, Stock assessment

Introduction

Presently, the concept of fish stock identification is getting popular in terms of sustainability. Stocks are defined as a self-sustaining population which comes from the same gene pool, more or less isolated from the same species. Inadequate knowledge on population differentiation and connectivity may impact the accuracy of stock assessment, whereby management decision could result in recruitment failures, loss of productivity and reduced yields (Cadrin and Friedland, 2005).

One of the most frequently used and cost effective method for identification of stock is analysing morphometric variation in fish. Morphometric variables are measured from the distance based on anatomical landmarks to study the phylogeny, phenotypic plasticity and fish condition. The conventional morphometric analysis mostly relied on the measurement along the longest axis of the body, thus making it difficult to measure the actual aspect of body shape (Bookstein *et al.*, 1985). In contrast, truss network method is not dominated by excessive measurement along single axis which presents a comprehensive characterisation of shape and thus it is considered as a powerful tool to describe morphological differentiation (oblique, horizontal and vertical) for

individual classification and intraspecific identification (Strauss and Bookstein, 1982; Turan, 1999).

The species *Clupisoma garua* (Hamilton, 1822) along the Ganga River commonly known as *Garua bachcha* or river catfish belongs to family Schilbeidae. It is a subtropical freshwater catfish inhabiting the bottom niche in riverine ecosystem of Gangetic plains having high percentage of fat content (Jafri *et al.*, 1964; Talwar and Jhingran, 1991). *C. garua* commonly feeds on molluscs, insects, crustaceans, decaying matter and sometimes algal matters (Hora, 1929). The species was considered as a Least Concern species under IUCN Red List of Threatened Species (IUCN, 2017) but it was reported as Vulnerable in River Betwa (Lakra *et al.*, 2010; Dubey *et al.*, 2013). Moreover, *C. garua* is recognised as vulnerable species (CAMP, 1998; CAFF, 2006) and a gradual reduction in natural population has been observed (Saraswat *et al.*, 2013). The catch rate of *C. garua* has declined by about 30.4% during 1960-2000 in Sundarbans and 27.7% in south-western Bengal (Patra *et al.*, 2005; Mishra *et al.*, 2009). Insufficient information regarding its distribution was noticed in other localities (Verma *et al.*, 2014; Siddik *et al.*, 2017; Biswas *et al.*, 2018).

To determine stock structure of a single species several complimentary techniques need to be assessed

The findings of Biswas *et al.* (2018) using conventional morphometric variations and meristic counts revealed multiple stocks of *C. garua* in the middle and lower Gangetic regions. Therefore, to facilitate stock assessment, it was felt essential to comprehend the stock structure of *C. garua* along the Ganga River using truss network by considering complete body archive without any information loss. Though several fish faunal studies were conducted in River Ganga and its tributaries (Hamilton, 1822; Hora, 1929; Sarkar *et al.*, 2012) sufficient data on Schilbid freshwater catfish was lacking. Hence, the aim of the present study was to assess the stock structure of *C. garua* in the middle and lower stretches of the Ganga River by emphasising on truss network.

Materials and methods

Collection of samples

A total of 306 specimens of *C. garua* were obtained from three locations of the river Ganga, viz., Allahabad (n=100), Farakka (n=109) and Triveni (n=97) in India during March and May, 2016 (Fig. 1). The sampling was done prior to breeding season (June-August) of the species for putative stock identification (Hora, 1938; Chondar, 1999). The specimens were collected from commercial fish catch in gillnet and hook and line, followed by screening for any physical damages. Simple random sampling was adopted to collect the fish from the landing centre. The collected specimens were preserved in ice in an insulated box and transported to the laboratory for further analysis.

Digitisation of samples

Digital images of the samples were taken after thawing the samples in running tap water. After thawing,

the samples were wiped well and placed on a platform with horizontal and vertical grids forming square of 1 cm² size to calibrate the coordinates of the digital images. The body and fins of the samples were stretched to natural position for facilitating the visibility of insertion points. Homologous anatomical landmarks across specimens were marked by piercing round head pins to get an effective discrimination among the group (Winans, 1987). Digital camera mounted on a leveling tripod was utilised to capture the images (Nikon Coolpix L840 16.0 MP). For perfect alignment, bubble level was monitored to provide a complete structural archive of the sample (Cadrin and Friedland, 1999).

Measurement of truss distance

To convert the images from JPEG (*.jpeg) to TPS (*.tps), tpsUtil V1.38 (Rohlf, 2006a) program was used. The truss distances were extracted from the digitised image using tpsDig2 V2.1 (Rohlf, 2006b) and Paleontological STatistics (PAST) (Hammer *et al.*, 2001) to construct an inter linked network of thirteen landmarks to form 27 truss measurements (Fig. 2).

Analysis of truss morphometric data

All data of truss measurement were transformed and its normality tested by SAS PROC UNIVARIATE procedure (SAS, 2008). An allometric approach was employed for size correction (Ihssen *et al.*, 1981; Hurlbut and Clay, 1998).

$$M_{\text{trans}} = (SL_{\text{mean}}/SL)^b \quad (1)$$

where, M_{trans} = Transformed truss measurement, M = Original truss measurement, SL = Standard length of fish, SL_{mean} = Overall mean standard length and b = Within-group slope regressions of the logM against logSL.

Fig 1. Sampling locations of *C. garua* along the river Ganga

Fig. 2. Truss network of *C. garua*

Correlation coefficient was calculated between the transformed data and SL of the fish to test whether data transformation is effective in removing the effect of size. Twenty seven size corrected truss measurements were exposed to principal component analysis (PCA) by PROC FACTOR procedure in SAS (SAS, 2008). Classification of individuals through cross-validation of the discriminant analysis was then exerted using PROC DISCRIM procedure of SAS (SAS, 2008).

Results

None of the standard truss measurements showed correlation with SL, which indicated that the body length effect has been removed by allometric transformation of data. The component loadings of the 27 truss distances (variables) are listed in Table 1 and it shows the variance explained by the variable on that particular component. The variables highlighted (with asterisk) are the variables with meaningful loading.

First two components of the PCA in the present study accounted for 61.37% of total morphometric variation. The characters belonging to depth of body and belly region were loaded highly on PC1 and the caudal portions were loaded on PC2 which accounted for 46.71% and 14.66% of total variance respectively. Variables loaded on first factor were all positive, indicating a positive correlation between variables within PC1. The truss distances with meaningful loading on PC1 were 2-12, 3-12, 4-11, 5-11, 5-12 and 12-13 (Fig. 3a) and were found to be significant. All these six distances characterised the anterior half of the fish body with depth measurements and horizontal abdominal position. PC2 explained 14.66% of total variation with two significant loadings which were 6-7 and 6-9 truss distances (Fig. 3b). In PC2, both positive and negative correlations were observed on trait loading.

The populations of Allahabad, Farakka and Triveni (Fig. 4) showed a single cluster due to marginal separation. Cross-validation of the discriminant analysis correctly classified 82.00, 88.07 and 85.57% of individuals of

Table 1. Variable loadings in principal component analysis of truss distance

Truss distances (cm)	PC1	PC2
01-02	0.1084	-0.1488
01-13	0.0701	-0.1272
02-03	0.1034	0.0480
02-12	0.2749*	0.1412
02-13	0.1067	-0.0638
03-04	0.0510	-0.0540
03-12	0.2934*	0.1754
03-13	0.1934	-0.0276
04-05	0.1365	0.1219
04-11	0.3068*	0.1599
04-12	0.2965	0.2030
05-06	0.2034	-0.4776
05-10	0.2837	-0.3459
05-11	0.2766*	0.1093
05-12	0.2795*	0.1602
06-07	0.1159	0.2826*
06-09	0.1426	0.2119*
06-10	0.1325	0.0721
06-11	0.2784	-0.3272
07-08	0.0560	-0.0479
07-09	0.1038	0.0056
07-10	0.1413	0.1693
08-09	0.0874	-0.0155
09-10	0.0598	0.1685
10-11	0.2313	-0.3665
11-12	0.1173	0.0240
12-13	0.2021*	0.0446

Allahabad, Farakka and Triveni populations, respectively. In this study, 85.21% individuals were well classified (Table 2).

Discussion

Stock identification and its quantification is essential to assess fishing pattern and state of the fishery resources (Gulland, 1971; Hilborn and Walters, 1992) to obtain the optimum yield over a long period of time. Generally, fishes show a high degree of variation within and among

Fig. 3. Distances with meaningful loadings on (a) PC1 and (b) PC2 components in truss network analysis of *C. garua*

Fig 4. Bivariate plot of scores on the two components extracted from 13 point truss measurements of *C. garua*

populations than other vertebrates and furthermore, the local habitats influence the morphological variations due to multiple stock reproduction and migration, which assists in distinguishing fish populations (Wimberger, 1992). Improper knowledge of such variations results in stock structure obscurity and genetic diversity loss, thus highlighting the importance of stock identification and its delineation.

Table 2. Percentage of fish from each location (in rows) classified in each location (in columns) in the cross validation of the discriminant analysis

Location	Allahabad	Farakka	Triveni
Allahabad	82.00	8.00	10.00
Farakka	6.42	88.07	5.50
Triveni	2.06	12.37	85.57
Total rate of classification (%)		85.21	
Total rate of misclassification (%)		14.79	

The present study on truss-based morphometrics exposed the phenotypic homogeneity among the populations of *C. garua* in three localities of Ganga River which may be due to inevitable migratory behaviour of the species, and as a result reproductive isolation was not noticed (Kumar *et al.*, 2012). PCA is a data reduction technique which facilitates identification of variables with high loading for a given component which is then subjected to scratching procedure as described by Hatcher (2003). Eigen values obtained from the covariance matrix using PCA, explained variations of about 46.71% from first component and 14.66% from the second representing body depth and caudal regions of fish, respectively caused

by different components. All the six distances that showed meaningful loadings on PC1 were positively correlated with major vertical size dimensions on the anterior side of fish body which has also been reported in Serranids (Cavalcanti *et al.*, 1999) and in *Megalaspis cordyla* (Sajina *et al.*, 2011). On the other hand, traits loaded on PC2 belonged to caudal region represented with positive and negative correlation. Variation in caudal region was observed in *M. cordyla* populations between the Arabian Sea and the Bay of Bengal (Sajina *et al.*, 2011) and in brook charr (Imre *et al.*, 2002). These observations could be a consequence of phenotypic plasticity due to hydrological variations including water turbulence and swimming ability.

The coefficients of same sign in PCs are indicative of “size variation” while positive and negative reveal “shape variation” (Sundberg, 1989). However in the present study, both size effect and size variation on morphometric characters was highly predominant. Identical studies showed head and caudal region as the major morphological characters for stock delineation in Rasbora (Muchlisin, 2013) as well as in *Channa punctata* (Khan *et al.*, 2013). Slight variations among populations of a species prevail due to differences in food preferences and swimming activities, but cannot be reflected as stock heterogeneity (Wimberger, 1992). Contradicting this, studies on *C. garua* on the basis of univariate analysis found significant variations with varying degree of difference in five populations viz., Yamuna, Betwa, Gomti, Son and Tons rivers (Verma *et al.*, 2014).

Further cross-validation of the morphometrics by DFA exhibited moderate percentage of classification confirming marginal separation in *C. garua* population. DFA is used to determine the continuous variables that discriminate between two or more naturally occurring groups. The analysis revealed that Allahabad population as the most well-defined group with only 8.48% of misclassified individuals whereas Farakka showed the highest discriminant value of classification among the three locations. Likewise misclassification report on *Heteropneustes fossilis* distinctly supported the similarity between its populations (Khan *et al.*, 2012). However, minimal phenotypic differences may be attributed due to water temperature, salinity, food availability and swimming pattern while experiencing specific environmental conditions (Rahman *et al.*, 2014).

The present study of truss network system successfully revealed a single continuous stock of *C. garua* population in three different locations of the Ganga River. Therefore, a uniform management strategy by developing suitable mesh size/hook size based on this morphometric study for sustainability of *C. garua* could be provided to

policy makers. The findings can further be validated by additional studies employing molecular techniques.

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