

Research Article

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# Meristics and Morphometrics of Four Populations of *Clarias gariepinus* (Burchell 1822) Fish, in the Sudan



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## Abstract

The intraspecific variation of *Clarias gariepinus* across four geographically distinct populations in Sudan was evaluated using meristic counts and morphometric measurements. Fish samples were obtained from Sinnar Dam area, Blue Nile (Site 1), Khor Abu Gassaba, White Nile (Site 2), Tordat Al Rahad an inland lake, mid Kordofan (Site 3) and Khashm El Girba, Atbara River (Site 4). Analysis of variance revealed highly significant differences ( $p < 0.01$ ) among populations in four of five meristic traits and 26 of 27 morphometric measurements. Post hoc comparisons (LSD test) showed that the Site 1 population exhibited the highest mean values ( $p < 0.05$ ) for all meristic and morphometric characters. In contrast, Site 3 population showed the lowest means in three of five meristic counts, while Site 2 individuals recorded the lowest morphometric means, except for orbital frontal width. Twelve morphometric variables and several meristic traits contributed substantially to total phenotypic variation, indicating their diagnostic utility in population differentiation. Head measurements were strongly and positively correlated with head length. Most morphometric traits exhibited high positive correlations ( $r > 0.65$ ), whereas others showed moderate to weak ( $r < 0.40$ ) associations. Standard length and body weight demonstrated strong positive correlations with most morphometric variables. The length-weight relationship was highly significant ( $p < 0.0001$ ), with correlation coefficients ranging from 0.766 to 0.979. Principal Component Analysis grouped the populations into two clusters, with partial segregation of Site 1 population, suggesting relatively greater intra-population variability. The findings indicate a close phylogenetic affinity among the four populations, likely derived from a common clade, while identifying key morphological traits relevant for taxonomic discrimination and selective breeding programmes.

**Keywords:** *Clarias gariepinus*; Meristics; Morphometrics; Sites; Sudan

## Introduction

The African catfish, *Clarias gariepinus*, is widely distributed across Africa, Asia, and Southeast Asia [1] and is particularly common in the Nile and its tributaries [2,3]. In Sudan, studies on freshwater fish taxonomy initially focused on descriptive morphological characteristics [4,5]. This approach later expanded to include meristic counts and morphometric measurements [6-8]. More recently, molecular techniques have been applied

to refine the classification and identification of freshwater fish species [7,9,10], although traditional morphometric and meristic methods remain central tools in fish taxonomic studies.

Phenotypic diversity in *C. gariepinus* were investigated in Nigeria [11,12] and in Turkey [13,14], revealing significant variation among populations. These differences have been attributed to genetic variability, environmental influences, or both.

The freshwater fish fauna in Sudan is similarly diverse, highlighting the importance of assessing meristic and morphometric traits in natural populations to determine patterns of homogeneity or heterogeneity among populations.

The objective of this work in *C. gariepinus* is to compare homogeneity and heterogeneity of meristic counts and morphometric characters three populations from three tributaries from the Nile and one from an inland lake.

### Material and Methods

#### Source of specimens and identification

One hundred twenty-three specimens of *C. gariepinus* fish were obtained from 4 sites. These were Site 1 (Sinnar Dam area, Blue Nile, n=30); Site 2 (Khor Abu Gassaba, White Nile, n=30); Site 3 Tordat Al-Rahad, N. Kordofan State, n=33) and Site 4 (Khashm Al-Girba, Atbara River, n=30). Fish identification followed Abu Gideiri [4] and Bailey [5].

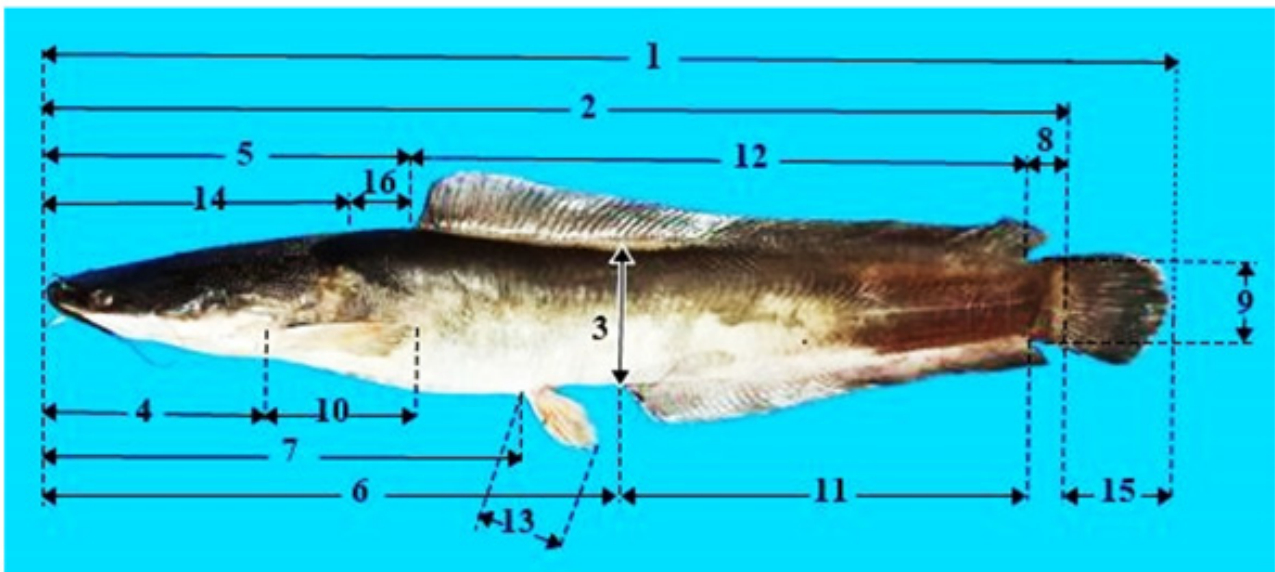


Figure 1: *Clarias gariepinus* lateral view.

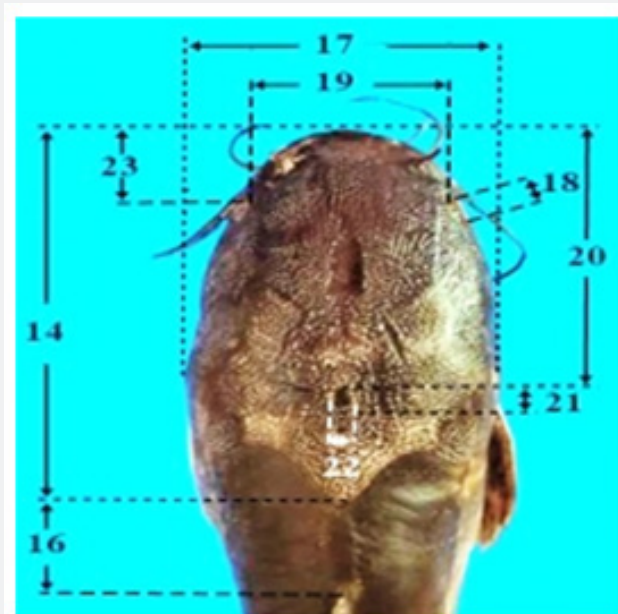


Figure 2: *Clarias gariepinus* top view of head.

**Meristic counts and morphometric measurements**

Five meristic counts (The number of rays in the: dorsal fin

(DFR); anal fin (AFR); pectoral fin (PFR); pelvic fin (PEFR) and caudal fin (CFR). Twenty-six morphometric measurements (Figures 1 & 2, Table 1) were measured following Ebraheem [7].

**Table 1:** Measured morphometric character of *Clarias gariepinus* and their acronym.

Morphometric character and measurement unit	Acronym	Code
Body weight was obtained and recorded in grams	BW	
Total length: distance from the tip of snout to the end of caudal fin (cm)	TL	1
Standard Length: from the tip of snout to the caudal fin base (cm)	SL	2
Body depth: from the mid dorsal fin to the anal opening (cm)	BD	3
Pre-pectoral distance: from the tip of snout to base of the first pectoral fin ray (cm)	PPD	4
Pre-dorsal Distance: from the rostral tip of the upper jaw to the most rostral point of the dorsal fin base (cm)	PDD	5
Pre-anal Distance: from the rostral tip of the upper jaw to the most rostral point of the anal fine base (cm)	PAD	6
Pre-Ventral Distance: from the upper lip to the origin of pelvic fin (cm)	PVD	7
Distance between dorsal and caudal fin (cm)	DDCF	8
Caudal peduncle depth: vertical depth of caudal peduncle (cm)	CPD	9
Pectoral fin length: length of pectoral fin (cm)	PFL	10
Anal fin length: horizontal length of the anal fin base (cm)	AFL	11
Dorsal fin length: from its most anterior to its most posterior point (cm)	DFL	12
Pelvic fin length: length of pelvic fin (cm)	PSL	13
Head length: rom the tip of snout to posterior margin of operculum (cm)	HL	14
Caudal peduncle length: length of caudal peduncle (cm)	CPL	15
Distances between occipital process and dorsal fin (cm)	DODF	16
Head width: as measured at level of posterior edge of frontal (cm)	HW	17
Eye diameter: maximum diameter of the eye (mm)	ED	18
Inter-orbital distance was the distance between the orbits (mm)	ID	19
Distance between snout and occipital process (mm)	DSO	20
Length of occipital fontanelle (mm)	OFL	21
Width of occipital fontanelle (mm)	OFW	22
Snout length from the tip of snout bony anterior margin of eye (cm)	SNL	23
Maxillary barbell length (cm)	MBL	
Nasal barbell length (cm)	NBL	
Inner mandibular barbell length (cm)	IMBL	

**Table 2:** The ANOVA of mean meristic counts of the four sites and the four populations.

Mean count						
Traits	DF	AFR	BEFR	PFR	CFR	DFR
Location	3	35.71**	1.21*	2.13**	3.50**	199.19**
Error	119	10.11	0.36	0.22	0.64	14.75
Total	122					
Character	Site 1	Site 2	Site 3	Site 4	5% LSD	
AFR	51.37 <sup>a</sup>	50.47 <sup>ab</sup>	49.15 <sup>b</sup>	49.17 <sup>b</sup>	1.63	
PEFR	8.63 <sup>a</sup>	8.63 <sup>a</sup>	8.24 <sup>c</sup>	8.37 <sup>b</sup>	0.31	
PFR	5.93 <sup>a</sup>	5.7 <sup>ab</sup>	5.55 <sup>b</sup>	5.3 <sup>c</sup>	0.24	
CFR	18.57 <sup>a</sup>	18.27 <sup>ab</sup>	18.03 <sup>c</sup>	17.77 <sup>c</sup>	0.41	
DFR	70.53 <sup>a</sup>	67.37 <sup>b</sup>	64.88 <sup>c</sup>	65.47 <sup>c</sup>	1.96	

\*Significant difference (p<0.05), \*\*highly significance (p<0.01), different superscript in a raw indicates significant differences.

**Table 3:** ANOVA of the morphometric characters in four populations of *Clarias gariepinus*.

Trait	DF	BW	TL	ST	PFL	AFL	DFL	PSL
Site	3	939939.9**	1233.5**	894.2**	22.8**	166.6**	405.6**	12.8**
Error	119	32514.11	26.77	22.80	0.39	3.71	7.10	0.26
Total				122				
F calc.		28.91	46.08	39.22	57.96	44.89	50.72	49.96
Trait	DF	PDD	DODF	PPD	PVD	PAD	PDA	DOCF
Site	3	120.6**	2.4**	45.5**	231.9**	326.2**	31.48**	9.78**
Error	119	2.77	0.21	1.12	4.61	6.81	1.43	0.15
Total				122				
F calc.		43.58	11.00	40.44	50.31	47.92	22.06	65.42
Trait	DF	CPD	CPL	MBL	NBL	OMB	IMBL	ED
Site	3	6.7**	22.5**	60.3**	5.47**	47.3**	16.2**	23.4**
Error	119	0.21	0.61	1.51	0.57	3.47	0.68	1.13
Total				122				
F calc.		31.91	37.05	40.04	9.58	13.62	23.99	20.71
Trait	DF	SNL	OFL	OFW	HW	DSO	HL	ID
Site	3	246.2**	5.2*	3.5**	2731**	4125**	99.3**	1333**
Error	119	10.08	1.38	0.77	87.3	110.49	2.06	31.91
Total				122				
F calc.		24.42	3.76	4.52	31.28	37.33	48.23	41.78

**Table 4:** ANOVA of the morphometric characters of *Clarias gariepinus* from four sites.

Character	Site 1	Site 2	Site 3	Site 4	5% LSD
Morphometric measurements					
TL	42.98 <sup>a</sup>	28.27 <sup>d</sup>	35.28 <sup>c</sup>	39.97 <sup>b</sup>	2.640
SL	37.40 <sup>a</sup>	24.97 <sup>c</sup>	31.07 <sup>b</sup>	35.15 <sup>a</sup>	2.440
PFL	4.32 <sup>a</sup>	2.27 <sup>d</sup>	2.94 <sup>c</sup>	3.48 <sup>b</sup>	0.320
DFL	24.12 <sup>a</sup>	15.59 <sup>d</sup>	18.98 <sup>c</sup>	21.73 <sup>b</sup>	1.440
AFL	15.89 <sup>a</sup>	10.56 <sup>d</sup>	12.43 <sup>c</sup>	14.59 <sup>b</sup>	0.980
PSL	3.37 <sup>a</sup>	1.82 <sup>d</sup>	2.45 <sup>c</sup>	2.84 <sup>b</sup>	0.260
PDD	13.57 <sup>a</sup>	8.93 <sup>d</sup>	11.32 <sup>c</sup>	12.59 <sup>b</sup>	0.850
DODF	2.13 <sup>b</sup>	1.66 <sup>c</sup>	1.94 <sup>b</sup>	2.32 <sup>a</sup>	0.240
PPD	8.34 <sup>a</sup>	5.52 <sup>d</sup>	6.83 <sup>c</sup>	7.76 <sup>b</sup>	0.540
PVD	18.50 <sup>a</sup>	12.14 <sup>d</sup>	15.3 <sup>c</sup>	17.3 <sup>b</sup>	1.100
PAD	21.79 <sup>a</sup>	14.53 <sup>c</sup>	18.50 <sup>b</sup>	21.13 <sup>a</sup>	1.330
PDA	7.40 <sup>a</sup>	5.18 <sup>c</sup>	6.56 <sup>b</sup>	7.30 <sup>a</sup>	0.610
DDCF	1.16 <sup>b</sup>	0.58 <sup>d</sup>	0.77 <sup>c</sup>	1.86 <sup>a</sup>	0.200
CPD	2.96 <sup>a</sup>	1.95 <sup>c</sup>	2.52 <sup>b</sup>	2.92 <sup>a</sup>	0.230
CPL	5.31 <sup>a</sup>	3.30 <sup>d</sup>	4.21 <sup>c</sup>	4.82 <sup>b</sup>	0.400
MBL	11.063 <sup>a</sup>	7.61 <sup>c</sup>	9.04 <sup>b</sup>	9.163 <sup>b</sup>	0.630
NBL	4.973 <sup>a</sup>	3.937 <sup>c</sup>	4.34 <sup>b</sup>	4.423 <sup>b</sup>	0.390
OMB	8.43 <sup>a</sup>	5.44 <sup>c</sup>	6.62 <sup>b</sup>	7.33 <sup>a</sup>	0.950

IMBL	5.65 <sup>a</sup>	3.91 <sup>a</sup>	4.42 <sup>b</sup>	4.53 <sup>b</sup>	0.420
ED	6.343 <sup>a</sup>	4.553 <sup>c</sup>	5.103 <sup>b</sup>	6.257 <sup>a</sup>	0.544
ID	44.53 <sup>a</sup>	28.3 <sup>c</sup>	35.94 <sup>b</sup>	37.6 <sup>b</sup>	2.88
SNL	20.4 <sup>a</sup>	14.4 <sup>c</sup>	20.182 <sup>a</sup>	17.167 <sup>b</sup>	1.623
OFL	6.543 <sup>ab</sup>	6.19 <sup>ab</sup>	6.6152 <sup>a</sup>	5.7167 <sup>b</sup>	0.600
OFW	3.943 <sup>a</sup>	3.767 <sup>a</sup>	3.170 <sup>b</sup>	3.646 <sup>a</sup>	0.450
HL	11.32 <sup>a</sup>	7.13 <sup>d</sup>	8.967 <sup>c</sup>	10.343 <sup>b</sup>	0.734
HW	63.243 <sup>a</sup>	40.233 <sup>c</sup>	52.546 <sup>b</sup>	55.267 <sup>b</sup>	4.777
DSO	77.75 <sup>a</sup>	50.233 <sup>d</sup>	64.121 <sup>c</sup>	70.733 <sup>b</sup>	5.374

Different superscripts in a raw are statistically different.

**Table 5:** Correlation between body weight (BW), standard length (SL) and morphometric measurements.

Traits	Site 1		Site 2		Site 3		Site 4	
	BW	SL	BW	SL	BW	SL	BW	SL
TL	0.964	0.965	0.895	0.977	0.977	0.610	0.757	0.941
ST	0.939	0.954	0.871	0.983	0.969	0.974	0.766	0.762
PFL	0.689	0.619	0.684	0.730	0.901	0.571	0.642	0.760
DFL	0.959	0.936	0.821	0.910	0.952	0.601	0.706	0.889
AFL	0.940	0.926	0.793	0.831	0.947	0.605	0.701	0.903
PSL	0.828	0.814	0.561	0.645	0.900	0.674	0.642	0.883
BD	0.794	0.728	0.800	0.715	0.697	0.689	0.802	0.873
CPD	0.696	0.729	0.671	0.751	0.884	0.890	0.614	0.830
CPL	0.725	0.730	0.679	0.587	0.631	0.623	0.531	0.719
PPD	0.892	0.921	0.729	0.878	0.947	0.962	0.609	0.951
PDD	0.954	0.905	0.827	0.729	0.939	0.958	0.738	0.826
PAD	0.926	0.927	0.852	0.934	0.948	0.966	0.750	0.957
PVD	0.933	0.934	0.885	0.895	0.957	0.971	0.761	0.970
DODF	0.572	0.587	0.680	0.401	0.427	0.659	0.464	0.562
DDCF	0.361	0.427	0.048	0.690	0.648	0.028	0.620	0.759
MBL	0.796	0.777	0.281	0.109	0.144	0.347	0.572	0.547
NBL	0.689	0.640	0.347	-0.205	-0.181	0.389	0.488	0.499
OMB	0.782	0.727	0.516	0.420	0.464	0.464	0.190	0.198
IMBL	0.802	0.731	0.069	0.332	0.384	0.080	0.271	0.424
SNL	0.777	0.804	0.696	0.777	0.649	0.690	0.271	0.388
HL	0.901	0.901	0.815	0.815	0.949	0.949	0.749	0.749
HW	0.939	0.939	0.936	0.936	0.967	0.967	0.343	0.343
ID	0.924	0.924	0.791	0.791	0.925	0.925	0.343	0.343
ED	0.546	0.546	0.642	0.642	0.429	0.429	0.448	0.448
DSO	0.987	0.987	0.815	0.815	0.932	0.932	0.574	0.574
OFL	0.073	0.073	0.179	0.179	-0.106	-0.106	0.372	0.372
OFW	0.087	0.087	0.371	0.371	0.353	0.353	0.198	0.198

**Table 6:** Correlation coefficient of Head length of *C. gariepinus* with different morphometric traits of head from the four populations.

Traits	Site 1	Site 2	Site 3	Site 4
HW	0.888	0.858	0.972	0.548
SNL	0.853	0.737	0.686	0.307
ED	0.522	0.566	0.420	0.447
ID	0.847	0.770	0.952	0.911
ST	0.906	0.873	0.980	0.936
DSO	0.898	0.954	0.981	0.783
OFL	0.189	0.092	0.004	0.298
OFW	0.050	0.300	0.291	0.246

r>0.65 highly, r>0.40 to r<0.65 moderately and r<0.40 weakly correlated.

**Table 7:** Similarity Matrix Index within the four populations.

Site	Lowest value	Highest value	High %	Moderate %	Weak %
Within populations					
Site 1 vs Site 1	0.289	8.512	80.0	13.0	7.0
Site 2 vs Site 2	0.094	3.567	85.0	14.0	1.0
Site 3 vs Site 3	0.171	6.473	71.9	20.47	7.5
Site 4 vs Site 4	0.290	6.095	85.8	10.7	3.3
Between populations					
Site 1 vs Site 2	0.353	17.898	78.00	13.77	8.2
Site 1 vs Site 3	0.453	13.372	86.00	8.30	5.3
Site 1 vs Site 4	0.370	13.722	89.33	9.50	1.1
Site 2 vs Site 3	0.281	9.719	82.80	13.20	3.9
Site 2 vs Site 4	0.567	8.245	69.00	19.66	11.3
Site 3 vs Site 4	0.395	6.773	68.80	25.95	5.2

Morphometric measurements were taken from the left side of the 123 *C. gariepinus* fish, using a measuring board, a tape and a Vernier calliper to the nearest 0.01 cm accuracy.

### Length-Weight Relationship

The Body weight (BW) was recorded in grams using a Docebel Braun balance. The relationship between Standard Length (SL) and Body Weight (BW) was obtained by linear regression analysis following the equation:  $SL = a + b BW$ ; Where: *a* = the intercept and *b* = growth mode.

### Statistical Analysis

Data were analyzed using analysis of variance (ANOVA) in Microsoft Excel after normality testing. Means were compared by the Least Significant Difference test (p<0.05). Correlation coefficients were estimated for traits. Regression of BW and SL was performed in GraphPad QuickCalcs.

## Results

### Meristic Counts

ANOVA results indicated that the meristic counts DFR, AFR,

PFR and CFR four of (Table 2) were highly significantly different (p<0.01) among the populations of the 4 sites. The PEFR was just significantly different (p<0.05) among the 4 populations. The highest means of the meristic counts were found among Site 1 population followed by Site 2 samples with no significant difference (p>0.05) between them in AFR, PEFR, CFR and PEFR. The DFR means in Site 1 was significantly different (p<0.05) from the other three sites.

### Morphometric measurements

The ANOVA (Table 3) shows that:

- The Sites significantly affected all measured traits. Most traits are highly significant at p< 0.01, while OFL is significant at p<0.05.
- The high F-calculated values show strong between-population variability relative to within-population variability. This indicates morphological heterogeneity among the four populations.
- ANOVA of TL, SL and BW suggest significant differential growth performance among sites.

d) All fin measurements (PFL, AFL, DFL, PSL) show highly significant differences ( $p < 0.01$ ) suggesting adaptation to different water regimes.

e) All PDD, DODF, PPD, PVD, PAD, PDA, DOC traits show significant site effects related to body proportions.

There is strong statistical evidence of morphometric variation among the four populations of *C. gariepinus*. High F-values in structural traits imply possible adaptive divergence rather than random variation.

The morphometric characters measured (Table 3), indicated highly significant differences ( $p < 0.01$ ) in all 26 characters among the fish populations in the 4 sites. OFL was significant ( $p < 0.05$ ) among the 4 populations. Site 1 *C. gariepinus* population scored the highest mean values with the exception of two characters, DODF and DDCF.

Table 4 shows that:

a) The 5% LSD suggests significant differences ( $p < 0.05$ ) and morphological heterogeneity among the four populations.

b) Site 1 consistently recorded the highest mean values for most morphometric characters and all fin-related parameters, while Site 2 had the lowest values across almost all characters. The environmental suitability for the populations followed the order: Site 1 > Site 4 > Site 3 > Site 2.

c) TL, SL and HL were significantly highest at Site 1.

d) Site 1 and Site 4 populations exhibited more robust body structures, whilst Site 2 individuals were comparatively slender and smaller.

e) Cranial and sensory structures ED, ID, SNL, OFL, OFW were significantly larger in Site 1.

f) DODF, DDCF were highest in Site 4 suggesting differences in microhabitat and/or water flow.

### Correlations between BW, SL and morphometric measurements

The correlation coefficients (Table 5) show that:

a) Strong positive relationships ( $r > 0.65$ ) between BW and most morphometric measurements in all sites. The strength of association varies by trait and location.

b) Site 4 tends to display comparatively lower (r-value) for many traits, though the relationships remain positive and moderate to strong for key measurements such as BD, head HL, and PVD.

c) DODF, DDCF, MBL, NBL, OFL, and OFW show low correlations ( $r < 0.40$ ), and in some cases negative values (e.g., NBL and OFL at Site 3). These weak associations suggest that these measurements contribute less to variations in body weight.

d) TL, PDD, PAD), PVD, AFL and DFL showed very high correlations ( $r \geq 0.90$ ) with SL across most sites. This proportional growth reflects coordinated somatic development.

e) BD, CPD, CPL, PPD, SNL and PSL exhibited moderate to strong positive correlations ( $r = 0.60-0.89$ ). Suggesting size-dependent flexible growth.

f) Measurements such as DODF, DDCF, MBL, NBL, OMB and IMBL showed weak, inconsistent, or even negative correlations (e.g., NBL at Site 3),  $r < 0.60$ ).

g) The regression of SL and BW of *C. gariepinus* from the four populations were:

Site 1;  $SL = 48.15BW - 1228$ ,  $r = 0.965$ ; Site 2;  $SL = 19.44BW - 321$ ,  $r = 0.871$ .

Site 3;  $SL = 32.39BW - 561.1$ ,  $r = 0.979$ ; Site 4;  $SL = 22.31BW - 347.1$ ,  $r = 0.766$  site 4.

The relation is very highly significantly correlated ( $p < 0.0001$ ).

The majority of the measurements of the head, in the 4 populations, were highly and positively associated with head length (Table 6). In sites 3 and 4 the correlation is moderate in ED. OFL from all sites and OFW from site 1, showed weak positive correlation.

### Similarity Matrix Index (SMI)

The SMI was used to quantify the structural resemblance within and between the four populations (Table 7). The SMI within-population displayed high values (71.9–85.8%); with highest homogeneity in Site 4 (85.8%) and Site 2 (85.0%). Site 3 shows comparatively lower SMI of 71.9% indicating heterogeneity. The SMI between-population comparisons show high readings (68.8–89.33%). The strongest resemblance occurs between Site 1 and Site 4 (89.33%), suggesting similar environmental conditions. In contrast, Site 3 vs Site 4 (68.8%) and Site 2 vs Site 4 (69.0%) display a relatively lower similarity percentages and higher moderate/weak components, implying greater compositional divergence.

### Principal Component Analysis

PCA of morphometric traits showed 80.7% accuracy across populations, ranging from 71% at Site 3 to 85.8% at Site 4, with variation at Site 1 and least at Site 2. The PCA (Figure 3) revealed that:

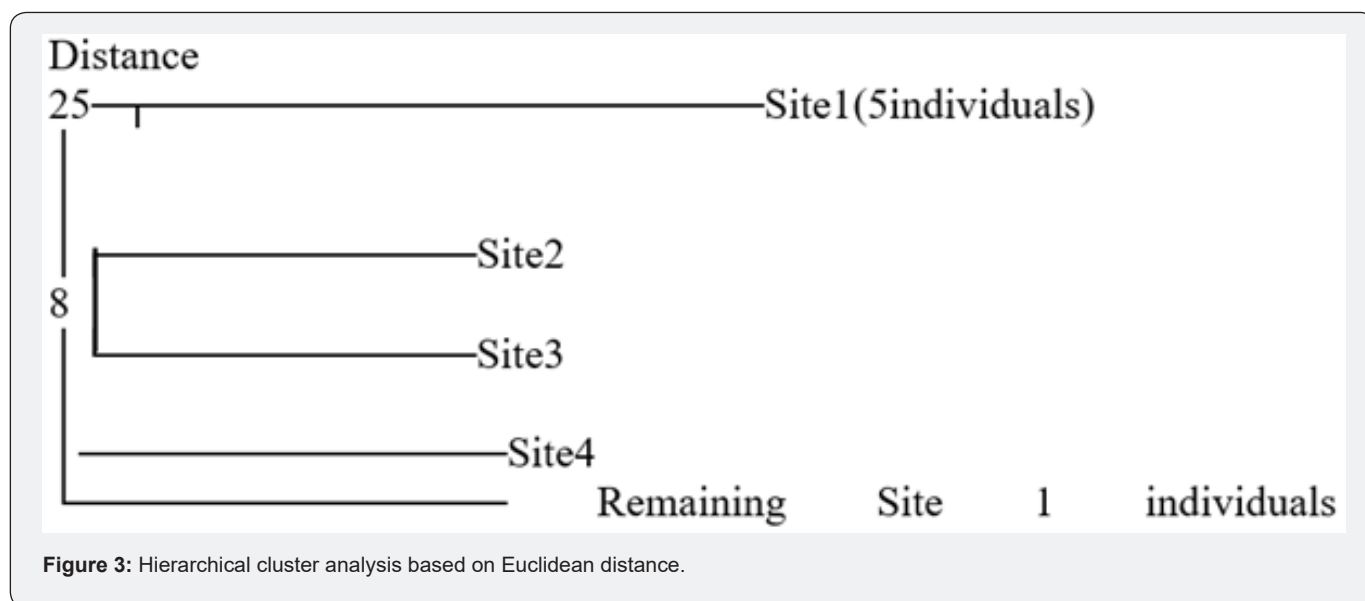
a) Sites, with lower values along the Y-axis indicate higher similarity.

b) At a distance of 25, Site 1 comprised five individuals forming a distinct cluster, with four individuals separating from the fifth, indicating intra-site variation.

c) At a distance of 8, Sites 2 and 3 clustered together, reflecting their high similarity.

d) Site 4, joining at a distance slightly below 8, exhibited moderate similarity to Sites 2 and 3.

e) The remaining individuals from Site 1 clustered separately, highlighting their marked dissimilarity from the other sites, indicating internal variation within Site 1.



### Discussion

*Clarias gariepinus*, an ecologically significant and economically valuable freshwater fish, is widely distributed throughout Africa and extending into parts of Asia and Southeast Asia [1]. It is abundant in the Nile [2,3]. Its geographic range, high tolerance, and importance in capture fisheries and aquaculture have made it a focal species in studies of taxonomy, population structure, and phenotypic plasticity.

In Sudan, descriptions made by [4,5] provided base knowledge of freshwater fishes, but they were often limited in resolving intraspecific variation. Subsequent investigations incorporated quantitative meristic counts and morphometric measurements to enhance taxonomy and population comparisons [6,7,9]. These methods allowed researchers to statistically evaluate differences in meristic and morphometric traits. More recently, molecular techniques have been introduced to complement morphological analyses and refine species identification and phylogenetic relationships [7,9,10].

The ANOVA demonstrated highly significantly ( $p < 0.01$ ), spatial variation in meristic traits among the four studied populations in DFR, AFR, PFR, and CFR, while PEFR showed a lower but still significant difference ( $p < 0.05$ ). Site 1 exhibited the highest mean meristic counts, followed by Site 2, with no significant differences ( $p > 0.05$ ) in AFR, PEFR, and CFR between them. However, DFR in Site 1 differed significantly ( $p < 0.05$ ) from the other sites, indicating localized divergence. Evidence on usefulness of meristic counts variation in morphological characterization, and population differentiation in *C. gariepinus* align with previous studies [15-21]

and supporting the taxonomic relevance of meristic traits. Using PCA, Umoh et al. [22] revealed that hybrids *C. gariepinus* from selected fish farms differed meristically in pectoral fin-ray, caudal fin-ray, anal fin-ray, dorsal fin-ray and pelvic fin-ray.

The present investigation indicated that the fish population in Site 1 recorded the highest mean value in all morphometric measurements (26 out of 27), and Site 2 the lowest means value (24 out of 28 traits, 88.9%). DDCF, OFL and OFW characters showed no correlation with SL and BW. The means, of TL, PFL, AFL, DFL, PSL, PDD, PPD, PVD, DDCF, HL and DSO as well as BW are most likely to be the characters that greatly contributed to the total variation in the four populations. Studies on *C. gariepinus* from the African freshwater bodies showed significant morphometric variations among populations from different environments [16,19,11,21,23]. Similarly, studies conducted in Turkey by Ihsan [13], Turan et al. [14] reported measurable differences in morphometric traits among geographically disconnected populations. These variations are attributable to environmental influences such as water flow [24], food availability [25], and habitat [26,27]. To this genetic divergence may accumulate due to geographic isolation or restricted gene flow [28] and consequently impact morphometrics.

The regression coefficient of the SL and the BW relationship of *C. gariepinus* from Site 1, Site 2, Site 3 and Site 4 populations indicates that the incremental change in (b) of the body weight is a consequence of change in SL. The highest b slope was among Site 1 (48.2) and the lowest (19.4) was recorded from Site 2. The result suggests that the SL has a high positive correlation with the BW and that the two variables can be used as indicators for

selecting a high body weight candidate for selective breeding. Anderson and Gutreuter (1983) stated that it is of paramount importance to determine this relation to establish the production and the biomass estimations of the species. Site 1 represented the highest mean BW value, and it has the highest and positive SL relationships. This finding indicated that the heavier the fish are the better its environmental conditions as reported by Idris and Mahmoud [6] for *Labeo niloticus*. The present study showed a positive length-weight relationship in the four populations, between the individuals of the four populations and between the sexes among the four populations. It seems that the conditions existing in the freshwater habitat, in the Sudan, are in favour for the propagation and the growth of this species. Another important morphological character is the measurements of the head. Most measurements of the head in the four populations are highly and positively associated with the head length are important in the characterization and identification of the species. This supports the findings of Turan et al. [14], in Turkey, in six populations of *C. gariepinus*. However, the ED reveals moderate correlation in the four population. On the other hand, OFW and OFL, in the four populations and the SNL in Site 4 population, showed weak positive correlation. It is concluded that the measurements of the head are important in the characterization of the fish.

This study demonstrated that PCA of morphometric traits in four *C. gariepinus* populations yielded a correct classification rate (80.7%), indicating significant phenotypic structuring among sampling sites. Multivariate approaches such as PCA have been widely applied in *Clarias* systematics and stock discrimination to detect subtle morphological divergence attributable to environmental and/or genetic factors. Studies of *Clarias* species across West African basins emphasized the importance of morphometric differentiation and attributed this to geographic variation.

The highest correct assignment observed in Site 4 (85.8%) suggests greater morphological distinctiveness of this population, possibly reflecting ecological specialism. Morphological divergence in *C. gariepinus* populations has been associated with habitat heterogeneity, and local adaptation [29]. Conversely, the lowest classification rate in Site 3 (71%) may indicate greater overlap in morphometric traits with adjacent populations, suggesting phenotypic plasticity and/or environmental pressures shaping body form. Such overlap has been reported in populations of *C. gariepinus* from Nigeria [11].

The minimal variation detected in Site 2 suggests relative genetic homogeneity or environmental stability. This reduced morphometric variability within populations of *Clarias* is in line with Fagbua et al. [30] and is likely to be linked to ecological effects. In contrast, Site 1 exhibited the greatest morphological variability indicating mixed stock origin and/or ecological heterogeneity. Turan et al. [14] related within-population to fluctuating environmental conditions promoting phenotypic

flexibility.

LSD analyses revealed strong affinities among Sites 1, 3, and 4, suggesting comparable ecological regimes as suggested by Teugels (1992). However, the comparatively lower similarity between Abu Gassba and Site 4 (69%), as well as between Site 3 and Site 4 (68.8%), indicates measurable divergence. The causes of such differentiation in *Clarias* populations are likely related to hydrological isolation, anthropogenic influences, or differential.

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