



# International Journal of Fisheries and Aquatic Studies

E-ISSN: 2347-5129

P-ISSN: 2394-0506

(ICV-Poland) Impact Value: 5.62

(GIF) Impact Factor: 0.549

IJFAS 2018; 6(4): 115-123

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www.fisheriesjournal.com

Received: 13-05-2018

Accepted: 16-06-2018

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## Seasonal distribution patterns of fish assemblage in the lagoon of Somone natural reserve of communal interest in Senegal: influence of abiotic factors

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### Abstract

Fish assemblage was investigated for patterns in seasonal structure in Somone lagoon, Senegal. Seasonal sampling from three stations with beach seine was performed in 2017. Abiotic factors such as temperature and salinity were measured during each season at all stations. Ecological and trophic classification was carried out in order to determine the fish fauna nature of this lagoon. Multivariate analysis like factorial correspondence analysis (FCA) and hierarchical classification analysis (HCA) were carried out to study temporal organization of fish assemblage. Canonical correspondence analysis (CCA) was performed to assess the influence of abiotic factors on temporal fish assemblage distribution. The fish assemblage of Somone lagoon consisted of 29 species from 19 families. The most abundant species were Mugilidae and Cichlidae species, accounting for 82.87% of total abundance and 87.53% of the total biomass. In terms of ecological and trophic guilds, the fish assemblage was dominated by species with estuarine affinity (19 species representing 94.75% of total individuals and 94.69% of total biomass) and herbivorous species (8 species accounting for 82.80% of total individuals and 87.17% of total biomass). The multivariate analysis indicated that temperature and salinity played an important role on the temporal distribution of fish assemblage in Somone lagoon.

**Keywords:** Canonical Correspondence Analysis, Fish assemblage, Somone lagoon

### Introduction

There is a growing interest on lagoon and estuarine ecosystems due to their ecological and socioeconomic importance. Interface between marine systems and continental environments, lagoons provide a variety of habitat types for many species, function as nursery areas and feeding grounds for marine opportunistic species as well [29, 45, 52]. Some of them support important fishery activities.

However, overfishing, climate change, pollution, lack of management, among other factors, are considered as major concerns for both the structure and the functioning of these sensitive coastal ecosystems [27, 53]. In this context, for their sustainability, it is necessary to understand and protect these critical ecosystems by documenting the communities they support and by investigating the factors that influence the distribution and abundance of associated species.

Therefore, several studies investigating fish assemblage structure were carried out within estuarine and lagoon ecosystems [2, 6, 7, 21, 23, 27, 37, 42, 54, 61, 63]. Most of these studies suggested that variations in time of fish assemblage could be caused by variation of environmental conditions. The main environmental parameters used to assess habitat conditions on fish assemblage distribution in estuarine ecosystems are temperature, salinity, dissolved oxygen, pH and turbidity.

In Senegal, studies on fish assemblage in lagoon ecosystems are scarce. This study dealing with the lagoon of Somone in the Natural Reserve of Communal Interest, aims to describe the fish assemblages in terms of their taxonomic and functional composition and to describe their seasonal variations relative to environmental fluctuations. In this study, fish assemblage patterns were investigated by clustering the different species according to the sampling seasons relative to the similarities in species composition and their abundance. The canonical correspondent analysis was applied to assess environmental influences on temporal fish assemblage distribution.

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## 2. Materials and Methods

### 2.1 Study area

Somone lagoon is located in Somone Natural Reserve of Communal Interest (RNICS) which encompasses three rural settlements: Guereo, Thiafoura and Sorokhassap (Sindia) and commune of Somone, in the region of Thiès (Fig. 1). The RNICS covering an area of about 7 km<sup>2</sup> was established in 1999. Somone lagoon with a surface of 2.04 km<sup>2</sup>, is covered with mangrove, mainly *Rhizophora racemosa* and *Avicennia africana* which serve as nursery areas and feeding grounds for opportunistic marine species [29, 45, 52]. It communicates with the sea via approximately 0.5 to 3-m-deep channel and receives freshwater from Somone river (Fig. 1). The climate in the RNICS is tropical Sahelo-Sudanian which consists of dry season, cool from November to April, and warm from May to June, and by a short warm and rainy season from July to October. Total annual rainfall is estimated at 760 mm with a maximum (400 mm) in August [12].

### 2.2 Sampling protocol

Samples were collected at three sampling stations (Fig. 1). At each station, samples were collected in Cold Season (CS) in April, transition Cold to Warm season (CW) in June, Warm Season (WS) in August and transition Warm to Cold season (WC) in December in 2017. The seasons correspond to the four main hydro-climatic periods in Senegal [24, 57, 58]. Fish Sampling was performed using a beach seine. Environmental parameters such as temperature and salinity were measured during each fish sampling with a multi-probe kit. After each fishing haul, fish were identified at the species level, counted, sized and weighed by species. In the case of large number of individuals, a sub-sample of 30 individuals per species was analyzed.

### 2.3 Data analysis

The relative abundance indices (AI) and the biomass indices (BI) were calculated as followed:

$$AI = \log\left(\frac{\text{Number of individuals for a given species}}{\text{Number of total individuals}} + 1\right) \quad (1)$$

$$BI = \log\left(\frac{\text{Biomass for a given species}}{\text{Total biomass}} + 1\right) \quad (2)$$

The logarithm function was applied to address the assumptions of normality and homogeneity of variance. Species richness (the total number of species caught in each station or during each season) was calculated. Species richness and abundance were compared between station and sampling seasons.

Species were classified according to their habitats and diet preferences. The ecological classification proposed by Albaret [7] was used in this study. This method classified species on several ecological guilds according to their degree of euryhalinity and the characteristics of their bio-ecological cycle in different estuarine environments. Five ecological categories were sampled in the GMPA: Strictly estuarine species (Es), Estuarine species from marine origin (Em), Marine-estuarine species (ME), Marine species which are accessory in estuaries (Ma) and Marine species that are occasional in estuaries (Mo). Concerning their feeding behavior, seven trophic guilds were identified: Scavenger or grazer herbivores (he-de), Herbivores mainly feeding on phytoplankton or micro-phytoplankton (he-ph), First level predators mainly benthophagous (p1-bt), First level generalist

predators mainly feeding on macro-crustaceans or insects (p1-mc), First level predators mainly feeding on zooplankton (p1-zo), Second level generalist predators mainly feeding on fish, shrimps and crabs (p2-ge) and Second level piscivorous predators mainly feeding on fish (p2-pi) [61].

### 2.4 Statistical analysis

An ANOVA was used to test for significant differences in environmental variables among seasons. Similar analysis was performed to examine significant difference between species richness, abundance and biomass between seasons. Multivariate analysis techniques such as factorial and automatic classification analysis methods that allow to resume the temporal organization of data from a complex picture whose structure is difficult to pin down clearly [50], were applied here. Factorial correspondence analysis (FCA) was carried out using fish abundance indices to investigate the pattern of species assemblage among seasons. The Hierarchical Classification Analysis (HCA) was also used to group species according to their seasonal affinity or similarity [44]. The dendrograms were performed using the Euclidean distance and the Ward minimum variance clustering method [28].

Canonical correspondence analysis (CCA) was performed to assess the influence of environmental variables on temporal fish assemblage structure [67]. This method (CCA) allows to assess the relative importance of environmental parameters to the distribution of each species. The relative length of the vector indicates the importance of the environmental parameter. More the vector is longer, greater is its influence. Concerning the species, the closer are two species, the more similar is their distribution; for the vectors [54, 67] as well. Species or groups that are highly influenced by two parameters are on the axes generated by the corresponding vectors of these parameters rather than at the end of any single vector [67]. The canonical analysis concerned only the most commonly occurring species (Mugilidae and Cichlidae species) in order to reduce effects of rare species. The Variance Inflation Factors (VIFs) were calculated for all environmental variables in order to detect possible high dimensional collinearities [74]. In fact, it was suggested by these authors that covariates with VIFs >5 are highly collinear.

## 3. Results

### 3.1 Fish assemblage

A total of 29 fish species belonging to 19 families from 2,114 specimens with a total biomass of 95.48 kg were identified in Somone lagoon (Table 1). Of the 29 species captured, the Mugilidae (6 species with 43.33% of total abundance and 53.19% of total biomass) and the Cichlidae (2 species with 39.54% and 34.34% of the total abundance and biomass) dominated the fish assemblage (Table 1). *Chelon dumerili* (22.80% of total abundance and 21.81% of total biomass), *Coptodon guineensis* (21.71% and 19.51% of total abundance and biomass) and *Sarotherodon melanotheron* 17.83% of total individuals and 14.83% of total biomass) were the dominant species.

According to their habitat preference, the estuarine species from marine origin (Em) dominated the fish assemblage in terms of species richness, abundance and biomass (10 species representing 50.57% and 56.45% of total abundance and biomass, respectively) (Fig. 2). The strictly estuarine species with 4 species, account for 41.87% of total individuals and

36.15% of total biomass precede the Em species. The marine-estuarine were the second most important ecological guilds in terms of species richness (6 species). The marine-occasionally species (5 species) were the lowest as well as in abundance and biomass (1.42% of total individuals and 0.93% of total biomass).

As for their diet preference, the scavenger or grazer herbivores species (he-de), 8 species accounting for 82.80% of the total abundance and 87.17% of the total biomass, were the most encountered (Fig. 3). Height species were benthophagous (p1-bt), comprising 9.40% of the total specimens and 3.98% of the total biomass. Ten species were second level predators (p2-ge and p2-pi, 5 species each one), but they comprised low number of the fish collected (5.24% of total abundance and 8.09% of total abundance). The omnivorous and phytoplanktivores species represented each by one species were the less abundant, less than 1% of total individuals and biomass.

The distribution of species richness, abundance and biomass was not significantly different among seasons ( $P > 0.05$ ). However, highest species richness (22 species), abundance (48.69% of total individuals) and biomass (37.97%) occurred in WS (Fig. 4). The lowest species richness, abundance and biomass (14 species accounting for 6.91% and 8.61% of total biomass) were recorded in CS. The clustering analysis based on species abundance allowed to identify three different groups (Fig. 5a). The first group (15 species accounting for 61.92% of total individuals and 56.21% of total biomass) was associated with WS (Fig. 5b). The second group consisted of the most abundant species in cold and WC (4 species accounting for 27.44% of total abundance and 28.99% of total biomass). The third group with 10 species, accounting 10.64% of total abundance and 14.83% of total biomass, was related to CS and CW.

### 3.2 Abiotic factors

The studied abiotic factors showed a clear seasonal pattern (Fig. 6). Temperature ranged from 19.51°C in CS to 28.03°C in WS (mean  $\pm$  SD = 23.41  $\pm$  4.01°C) (Fig. 6a). The differences in temperature between the four sampling seasons were not statistically significant ( $F=0.07$ ;  $P=0.80$ ). Salinity varying between 29.14‰ in WC and 45.5‰ in CW (29.83 $\pm$ 0.48‰) (Fig. 6b), was not significant between seasons ( $F=8.09$ ;  $P=0.09$ ).

### 3.3 Environmental influence on temporal fish assemblage distribution

The CCA analysis based on species abundance indices revealed that axis 1 (57.06%) and axes 2 (36.36%) explained 93.42% of variance of the seasonal species-environment relation. Temperature and salinity were positively and highly correlated with axis 1 (Fig. 7). According to the relative length of the vectors, both factors played an important environmental in the seasonal distribution of fish species in Somone lagoon, even if the influence of salinity was slightly more important. Temperature was positively and negatively correlated with *Chelon dumerili* and *Neochelon falcipinnis*, respectively. In other words, season distribution of both species was mainly influenced by temperature. *Mugil bananensis* and *Parachelon grandisquamis* were positively and strongly correlated with salinity. *Sarotherodon melanotheron* located near the origin either did not show a strong relation to any of the factors, while *Coptodon guineensis* was strongly influenced by both factors,

temperature and salinity. Salinity negatively influenced the abundance distribution of *Mugil curema*.

## 4 Discussion

### 4.1 Fish assemblage

Fish assemblage in Somone lagoon was characterized by moderate species diversity, 29 species belonging to 19 families. This species richness can be equal, higher or lesser than that of other lagoon or estuaries: 108 species in St Lucia lagoon [18], more than 70 in Laguna Madre, Texas [35], 26 in Ghar El Melh in Tunisia, 17 in Lake Manzala in Egypt [41], 13 in the Ichkeul Lagoon in Tunisia [62] and 11 in Mellah lagoon [27]. By comparison, in North-West Africa, 153 species from 71 families were identified in Ebrié lagoon in Côte d'Ivoire [5] and 15 fish species were recorded in Nador lagoon in Morocco [37]. However, caution should be taken when comparing fish diversity between these cited studies. In fact, difference in fish diversity could be related to the characteristics of the lagoon (surface area, depth, connection with the sea), hydrological parameters (e.g., tidal range, temperature, salinity etc.), the sampling effort, as well as the fishing gear type [2, 29, 45].

The most abundant species were the Mugilidae (*Chelon dumerili*, *Mugil bananensis*, *Neochelon falcipinnis*, *Mugil curema* and *Parachelon grandisquamis*) and the Cichlidae (*Coptodon guineensis* and *Sarotherodon melanotheron*) probably due to their high adaptation capacity in high salinity variations [3, 4, 19, 36, 51, 55, 63, 68]. In terms of richness, abundance and biomass, the fish assemblage of Somone lagoon was dominated estuarine species forms of marine origin (Em) whose life cycle is spread across both marine and estuarine environments [8, 25, 26]. Concerning the trophic guilds, herbivores were dominant in terms of species richness, abundance and biomass. Similar results were found in Palmarin Communal Natural Reserve [22]. Sadio *et al.* [61] reported that grazer herbivores were the most abundant in Bamboung marine protected area.

### 4.2 Spatial and temporal variation of fish assemblage in relation with environmental parameters

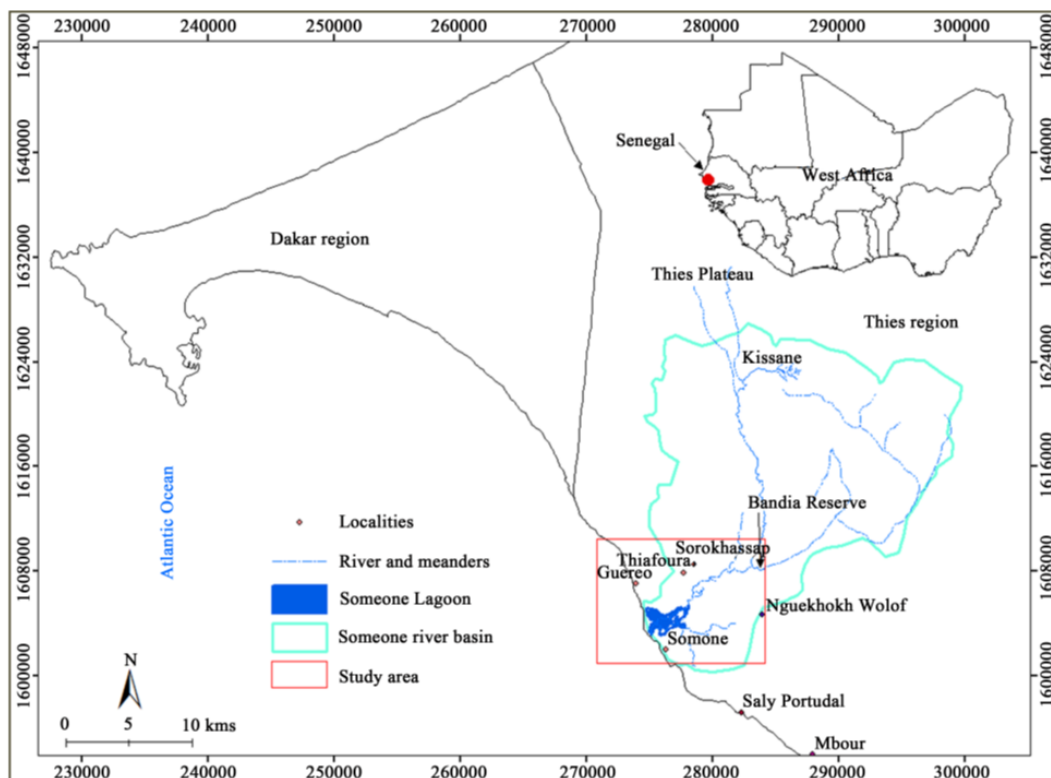
The fish assemblages differed among seasons both in terms of fish diversity, abundance and biomass. This temporal variation in total abundance reflects fluctuations in the most dominant species, the Mugilidae and Cichlidae (82.87% of the total sampled species and 87.53% of the total abundance). High abundance of these species within the lagoon might result of high tolerant to fluctuating environmental conditions [19, 36, 51, 63, 68]. In fact, it has been suggested that fish assemblage structure in lagoons and estuaries is influenced by both abiotic and biotic factors [1, 11, 17, 30, 31, 46, 48, 56, 59, 69, 71]. In this study, the CCA revealed that the temporal organization of the fish assemblages in Somone lagoon was greatly influenced by studied abiotic factors, temperature and salinity. These two abiotic factors have been postulated to be important determinants of spatial and temporal assemblage structure [1, 33, 34]. In fact, temperature always affects fish species at different stages of their life cycles, including during spawning and the development and survival of the eggs and larvae, as well as influencing their distribution, diet, migration pattern and schooling behavior [32, 43, 66]. Concerning salinity, several authors suggested that it influences reproduction, larval dispersal and recruitment, geographical distribution, and behavior of many species [10, 14, 17, 38, 64, 65]. As example, it has been showed that juvenile mullets prefer oligohaline

waters, while adults prefer euryhaline waters [20]. It's worth highlighting that, even if the abiotic factors tested in this study had an important influence on fish assemblage distribution in Somone lagoon, they could not fully explain the temporal of the assemblage. In fact, it has been suggested

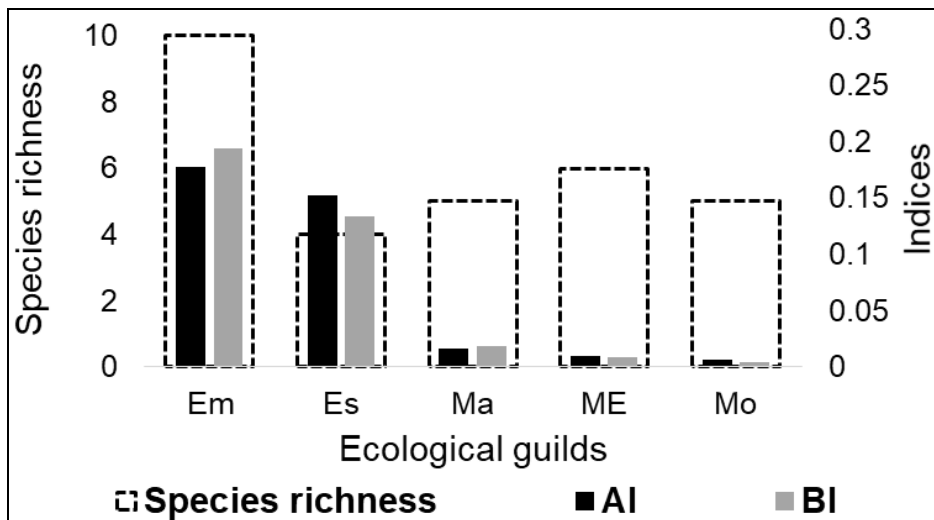
that other factors such as aquatic vegetation [1, 39, 60, 70, 73], food availability [15, 40, 59], status of the estuarine mouth (open or intermittently open) [16, 72], and biological interrelationships [49] were associated with fish assemblage structure in other lagoon and estuarine ecosystems.

**Table 1:** List of the 29 species observed in Somone lagoon in 2017. Percentage of abundance and biomass with the name of the family, species labels, ecological and trophic guilds.

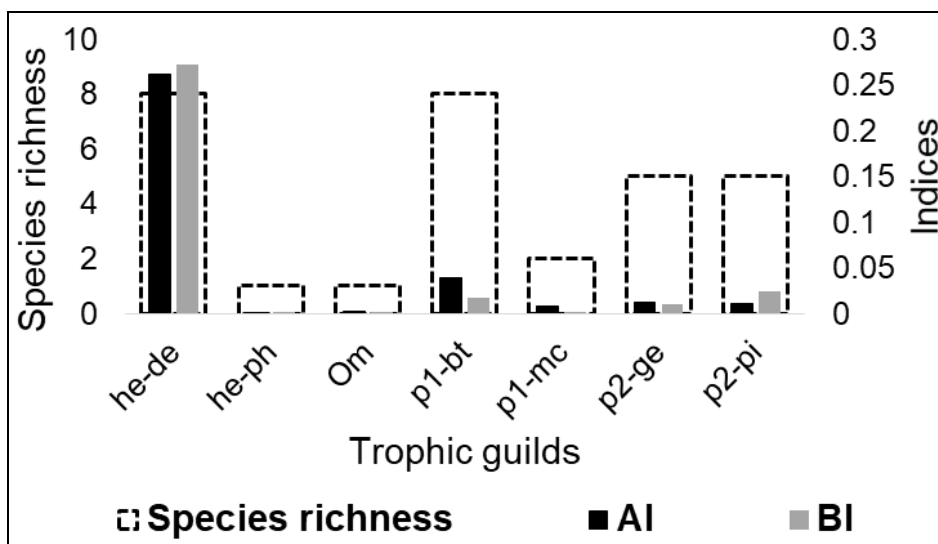
Family	Species	Labels	Ecological guilds	Trophic guilds	Abundance (%)	Biomass (%)
Carangidae	<i>Caranx hippos</i>	CHI	ME	p2-ge	0.19	0.02
	<i>Lichia amia</i>	LAM	Ma	p2-ge	0.14	0.21
Cichlidae	<i>Coptodon guineensis</i>	CGU	Es	he-de	21.71	19.51
	<i>Sarotherodon melanotheron</i>	SME	Es	he-de	17.83	14.83
Clupeidae	<i>Ethmalosa fimbriata</i>	EFI	Em	he-ph	0.14	0.21
Elopidae	<i>Elops lacerta</i>	ELA	ME	p2-pi	0.47	1.78
Gerreidae	<i>Eucinostemus melanopterus</i>	EME	ME	p1-mc	1.37	0.1
	<i>Gerres nigri</i>	GNI	Es	p1-mc	0.61	0.28
Haemulidae	<i>Pomadasys Jubelini</i>	PJU	Em	p1-bt	7	2.77
	<i>Pomadasys incisus</i>	PIN	Ma	p1-bt	1.8	0.42
	<i>Plectorhinchus macrolepis</i>	PMA	Em	p2-ge	0.05	0.1
Lutjanidae	<i>Lutjanus agennes</i>	LAG	Ma	p2-pi	1.66	3.56
	<i>Lutjanus dentalus</i>	LDA	Ma	p2-pi	0.19	0.1
	<i>Lutjanus fulgens</i>	LFU	Ma	p2-pi	0.05	0.1
Monodactylidae	<i>Monodactylus sebae</i>	MSE	Es	p2-ge	1.75	1.68
Moronidae	<i>Dicentrarchus punctatus</i>	DPU	Mo	p2-ge	0.71	0.47
Mugilidae	<i>Chelon dumereli</i>	CDU	Em	he-de	22.8	21.81
	<i>Mugil bananensis</i>	MBA	Em	he-de	6.05	9.17
	<i>Neochelon falcipinnis</i>	NFA	Em	he-de	5.96	8.49
	<i>Mugil curema</i>	MCU	Em	he-de	5.16	8.85
	<i>Parachelon grandisquamis</i>	PGR	Em	he-de	2.98	4.35
	<i>Mugil cephalus</i>	MCE	Em	he-de	0.38	0.52
Paralichthyidae	<i>Syacium micrurum</i>	SMI	ME	p1-bt	0.19	0.09
Serranidae	<i>Epinephelus aeneus</i>	EAE	ME	p2-pi	0.05	0.12
Soleidae	<i>Synaptura cadenati</i>	SSP	Mo	p1-bt	0.09	0.15
Sparidae	<i>Diplodus cervinus</i>	DCE	Mo	Om	0.43	0.16
	<i>Diplodus vulgaris</i>	DVU	Mo	p1-bt	0.14	0.16
	<i>Diplodus sargus</i>	DSA	Mo	p1-bt	0.05	0.01
Tetraodontidae	<i>Ephippion guttifer</i>	EGU	ME	p1-bt	0.05	0.01



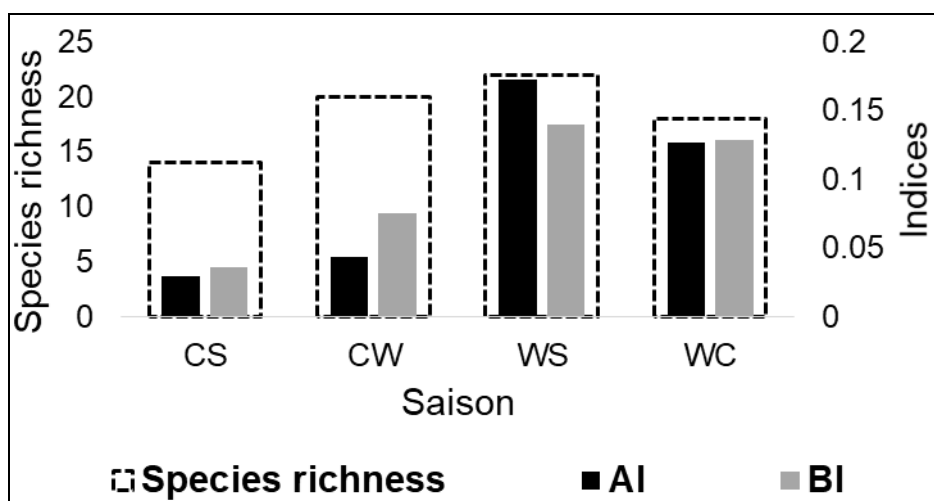
**Fig 1:** Location of the study area (Barry et al. [12]).



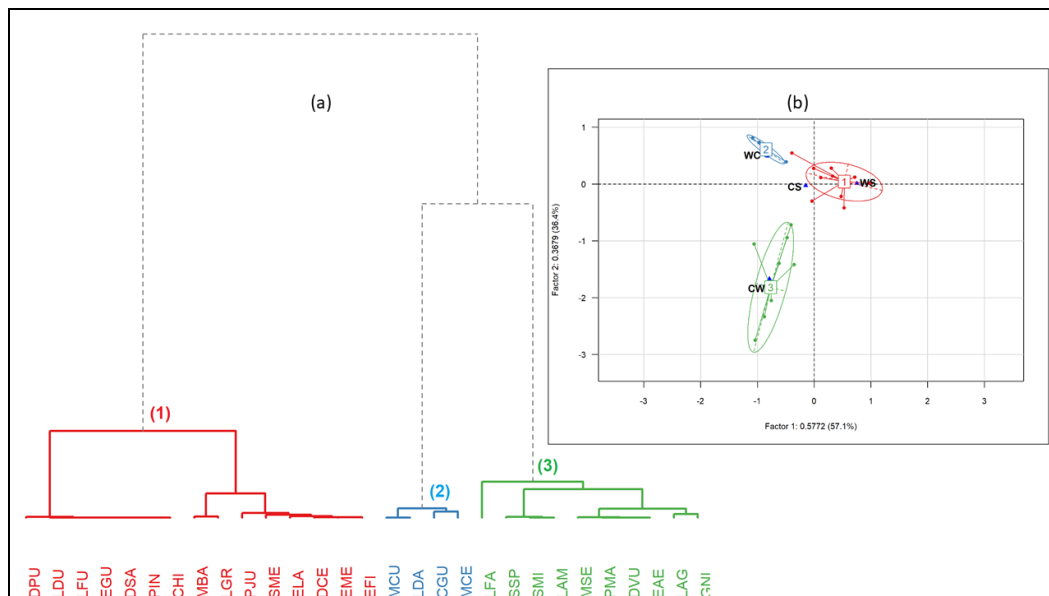
**Fig 2:** Species richness abundance and biomass of the different ecological and trophic guilds in Somone lagoon. AI = abundance indices; BI= biomass indices. Es = Strictly estuarine species, Em = Estuarine species from marine origin, ME = Marine-Estuarine species, Ma = Marine species, accessory in estuaries and Mo = Marine species, occasional in estuaries (Albaret *et al.*, 1999) [7].



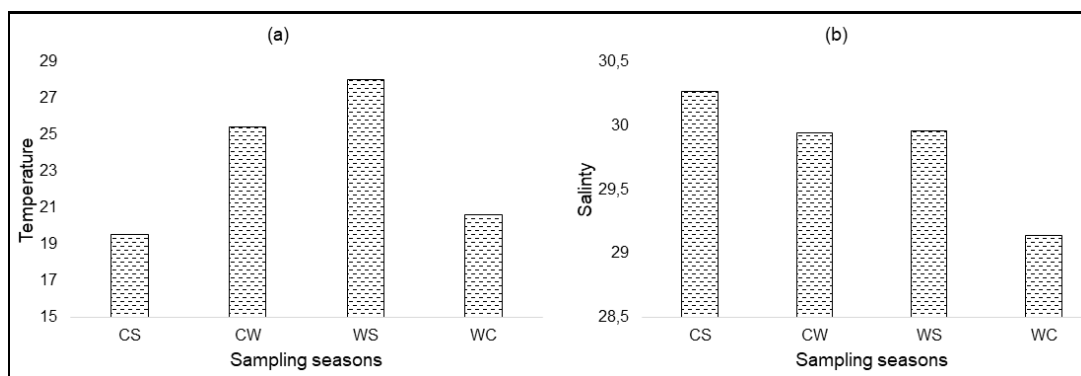
**Fig 3:** Species richness abundance and biomass of the different ecological and trophic guilds in Somone lagoon. AI = abundance indices; BI= biomass indices. he-de = scavenger or grazer herbivores, he-ph = herbivores mainly feeding on phytoplankton or micro-phytoplankton, Om = omnivorous species, p1-bt = first level predators mainly benthophagous (mollusks, cockles, marine worms), p1-mc = first level generalist predators mainly feeding on macro-crustaceans or insects, p2-ge = second level generalist predators mainly feeding on fish, shrimps and crabs, p2-pi = second level piscivorous predators mainly feeding on fish.



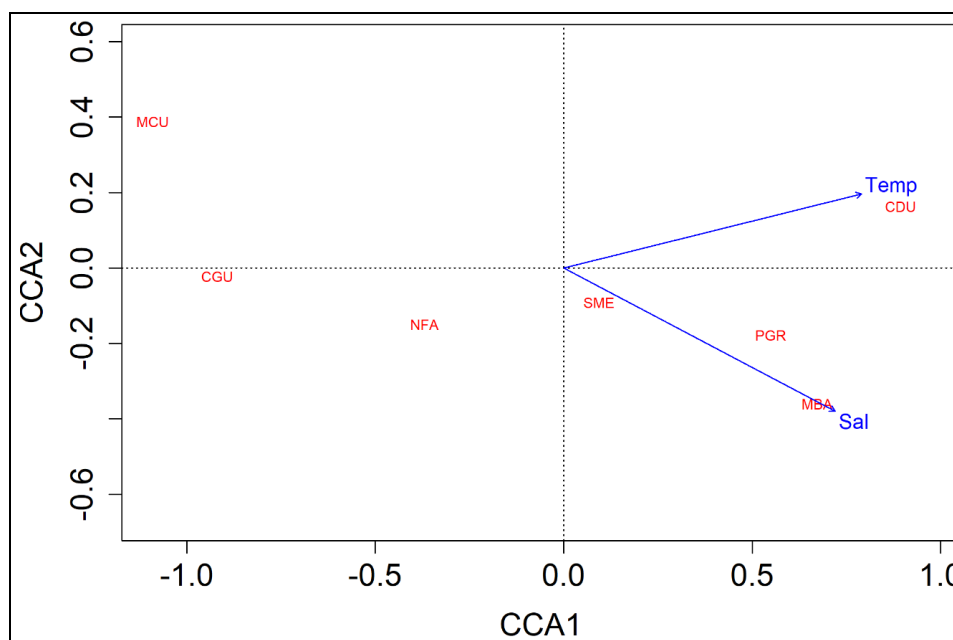
**Fig 4:** Temporal fluctuations in species richness, abundance and biomass in Somone lagoon in 2017. CS= Cold season; CW transition season from Cold to Warm season; WS= Warm season; WC transition from Warm to season Cold.



**Fig 5:** Factorial correspondence analysis (FCA) performed using seasonal abundance of fishes: a) is the dendrogram showing the groups of species, b) is the correspondence between groups and seasons. CS = Cold season, CW = Cold to Warm transition, WS = Warm season and WC = Warm to Cold transition. See Table 1 for species labels.



**Fig 6:** Temporal fluctuations of abiotic factors in 2017 Somone lagoon. CS= Cold season; CW transition season from Cold to Warm season; WS= Warm season; WC transition from Warm to season Cold.



**Fig 7:** Plot of canonical correspondence analysis relating season, fish taxa abundance and abiotic factors correlated with axes; response variables (species abundance) are plotted by species, arrows represent quantitative explanatory variables (environmental variables: Temp = temperature, Sal = salinity). See Table 1 for species labels.

## 5. Conclusion

Overall this study, the first, revealed that moderate species richness composed of five ecological guilds and seven trophic guilds, was the general features of Somone lagoon. The most abundant species were species with high adaptation capacities in strong salinity variations. The temporal organization of fish assemblage in this lagoon was greatly influenced by both temperature and salinity. Results from this study might serve as the reference point of the fish assemblage in Somone lagoon.

## Acknowledgement

We would like acknowledge the members of the Management Comity of the RNICS and Fishermen for their active collaboration. We are also thankful to the research institutions such as CRODT (Centre de Recherche Océanographique de Dakar-Thiaroye) and IUPA (Aquaculture and Fisheries Institute of the University of Dakar) for their support in the elaboration and the setting of this monitoring protocol.

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