

Metazoan Parasites of the Mediterranean Garfish Belone belone gracilis (Teleostei: Belonidae) as a Tool for Stock Discrimination

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Abstract

The aim of the present study was to provide a current survey of metazoan parasites of the Mediterranean garfish Belone belone gracilis Lowe, 1839 off Tunisian coast and their potential use as biological tags for discriminate stocks. Parasitological analysis of 453 specimens of B. b. gracilis along the eastern Tunisian coast revealed the presence of ten species of metazoan parasites: one monogenean Axine belones Abildgaard, 1794; four digeneans Lecithostaphylus retroflexus (Molin, 1859), Tergestia acanthocephala (Stossich, 1887) Stossich, 1899, Aponurus laguncula Looss, 1907 and Condylocotyla pilodora (metacercaria) Pearson et Prévot, 1985; one copepod Bomolochus bellones Burmeister, 1835; two isopods Irona nana Schioedte et Meinert, 1884 and Nerocila orbignyi (Guérin-Méneville, 1832); 1 acanthocephalan Telosentis exiguus (Von Linstow, 1901) and one nematod Anisakid larvae. Discriminant analysis using parasites as biological tags, allowed identifying two stocks of B. b. gracilis. Digenea L. retroflexus, A. laguncula and Anisakid larvae were the most important species in determining the position of sampled fishes from the central coast off Tunisia. Metacercaria C. pilodora characterized specimens from the Southern coast off Tunisia. These results were corroborated by comparing parameters of prevalence and mean abundance between zones. Discriminant analysis used for the separation of *B. b. gracilis* between seasons in both localities showed that a seasonal variation affected mainly specimens from the center suggesting seasonal move of fishes.

Keywords

Belone belone gracilis, Metazoan Parasites, Stock Discrimination, Tunisian Coast, **Mediterranean Sea**

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1. Introduction

Garfish, *Belone belone* (Linnaeus, 1761) is an epipelagic teleost marine fish widely distributed in the northeastern Atlantic Ocean, the Mediterranean and the Black Sea. Three subspecies have been recognised according to their geographic distribution: *B. b. belone* (Linnaeus, 1761) restricted to the north-eastern Atlantic; *B. b. euxini* Günther, 1866 founded in the Black Sea and *B. b. gracilis* Lowe, 1839 mainly distributed in the Mediterranean Sea [1].

Parasite fauna of *Belone belone* has received considerable taxonomic interest. Nine metazoan parasite species have been therefore reported in the Mediterranean subspecies *B. b. gracilis* [2]-[11]. Recently Radujković and Šundić (2014) listed six flatworm's species in the Adriatic Sea [12]. Oktener (2005) and Keser *et al.* (2007) recorded two species of parasites in the Black sea subspecies *B. b. euxeni* [13] [14]. Rynkiewicz (1970) and Grabda (1981) signalized eleven metazoan parasite species in the north east Atlantic subspecies *B. b. belone* and a geographic variation of its parasites were also shown by Grabda (1981) [15] [16].

At regional scale, parasite composition of *B. b. gracilis* remains unknown in the southern shore of the Mediterranean Sea; most studies on the metazoan parasites of *B. b. gracilis* were carried out in the north western and central Mediterranean Sea. Understanding geographical variation in the composition, abundance and diversity of parasite is important to identify and delineate populations of fish species in different regions of the world [17]-[20]. Therefore, in this study we try to list for the first time regionally the parasite fauna of *B. b. gracilis* in the south western Mediterranean Sea. Secondly, at local scale we use spatial variability of parasites between the central and southern coast off Tunisia for eventual identification of different stocks. Thus, parasites can be used as a model for the discrimination of fish stocks. It's well known that parasites have been used as biological tags in the stock discreteness, migratory movements of marine fish as well as indicators of host phylogenetics and systematics [21]-[27].

Knowledge of the number of fish stocks of *B. b. gracilis* is a basic requirement to ensure sustainable fishing. Indeed, overexploitation of benthic fish stock [28], currently in danger [29] (Anonymous, 2002), led to other little or no exploited resources. These resources are mainly the stock of small pelagic fishes and secondary economic interest fishes such belonid fishes. In eastern Tunisian coast, belonids landings represent only 0.3% of the national production in 2010 which province from fishing area at the central and the southern zones (almost 90%) [30].

The aims of this paper were to identify the metazoan parasites of *B*. *b*. *gracilis* in the south western Mediterranean Sea and to assess using spatial and temporal variability of metazoan parasites at local scale for stocks discrimination.

2. Materials and Methods

2.1. Study Area and Examination of Samples

A total of 453 specimens of *Belone belone gracilis* were randomly sampled from local fishermen using gillnets in common fishing areas off eastern Tunisian coast: the central zone (Gulf of Hammamett) and the southern zone (Gulf of Gabes) (Table 1).

Fishes were identified using Bauchot (1987) [31]. Sampled fishes fresh and frozen were subsequently examined for ecto- and endoparasites. Parasitological examination using a stereomicroscope under incident light included the skin, fins, gills, eyes, body cavity and visceral organs (stomach, intestine, liver, the pericardial cavity, swimbladder, gallbladder and gonads). Platyhelminthes were fixed between slide and coverslip in 70% al-

 Table 1. The number of samples of Belone belone gracilis caught from two zones off the eastern coast of Tunisia (NT: total number, TL: total length of fish samples).

Locolity	Season					TL (cm)
Locality -	Autumn	winter	Spring	Summer		
Centre	59	19	48	79	205	24.3 - 55
South	36	22	98	92	248	24.2 - 53.7

cohol or in Bouin's fluid. Fixed specimens were stained with Semichon's acetic carmine, dehydrated using a graded ethanol series then cleared in clove oil and mounted in Canada balsam. Other parasites (copepod, isopod, nematod and acanthocephalan) were directly fixed in 70% for later examination. Parasites were determined to the lowest taxonomic level possible.

2.2. Data Analysis

Prevalence (P) and mean abundance (MA) were determined following Bush *et al.* (1997) [32]. Non parametric analyses were performed to evaluate characteristics of the infection at population and community levels. Univariate analysis was performed to evaluate the infections at infrapopulation levels only for parasites with prevalence > 5% [32]. χ^2 analyses and a posteriori multiple comparisons for proportions were used to test for significant differences of prevalence between zones [33]. Student test was used to examine effect of geographic variation on mean abundance of parasites after log10(*x* + 1) transformed data.

Discriminant analyses, based on Mahalanobis distances were used to find differences between zones and between seasons in both zones (eight groups: every season associated to every locality) and to identify which parasite species were responsible for these differences [34].

3. Results

A total of 395 (87%) fish was found to be parasitized by at least one of parasite species. There were 12434 specimens of metazoan parasites collected, with mean of 32 ± 53 parasite/fish (1 - 408). Ten parasite species were identified: one monogenean *Axine belones* Abildgaard, 1794; four digenans *Lecithostaphylus retroflexus* (Molin, 1859), *Tergestia acanthocephala* (Stossich, 1887) Stossich, 1899, *Aponurus laguncula* Looss, 1907 and *Condylocotyla pilodora* metacercaria Pearson et Prévot, 1985; one nematod Anisakid larvae; one acanthocephalan *Telosentis exiguus* (Von Linstow, 1901); one copepod *Bomolochus bellones* Burmeister, 1835 and two isopods *Irona nana* Schioedte et Meinert, 1884 and *Nerocila orbignyi* (Guérin-Méneville, 1832) (**Table 2**). Infection site, prevalence and mean abundance of parasite species were also shown in **Table 2**. Digenean parasites were the most abundant and accounted 85% of collected material (**Figure 1**). *Condylocotyla pilodora* was the most abundant species with 9661 specimens collected (77.69% of all parasites). Only common parasites with prevalence > 5% in at least one station were included in further analyses as recommended by Bush *et al.* (1997).

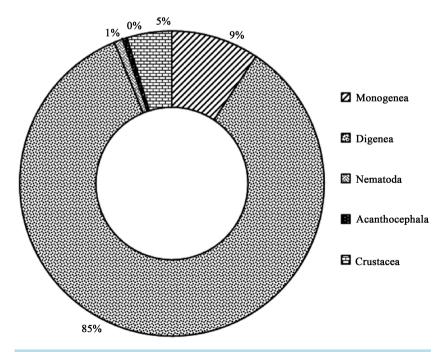


Figure 1. Participation of particular taxons of metazoan parasites in *B. b. gracilis* off Tunisian coast.

Parasite	SI	Р %	МА
Monogenea			
Axine belones	G	59	2.57
Digenea			
Condylocotyla pilodora (metacercria)	PS, DT	50	21.42
Lecithostaphylus retroflexus	DT	11	0.21
Tergestia acanthocephala	DT	24	1.17
Aponurus laguncula	DT	12	0.5
Crustacea			
Copepoda			
Bomolochus bellones	G	42	1.23
Isopoda			
Irona nana	G	7	0.07
Nerocila orbygini	Fi	1	0.01
Acanthocephala			
Telosentis exiguus	DT	4	0.13
Nematoda			
Anisakid larvae	DT, L	9	0.26

 Table 2. Prevalence, mean abundance and infection site of the metazoan parasites of *Belone belone gracilis*. SI: site of infection, P: prevalence and MA: mean abundance, DT: digestive tract, G: gills, L: liver, PS: Pericardial sac, Fi: Fins.

3.1. Geographic Variation

Comparisons of prevalence and mean abundance showed that five parasites differ significantly between two sampled zones. *A. laguncula* and *L. retroflexus* were more prevalent and abundant in the central zone. Anisakid larvae were more abundant in the center. *C. pilodora* and *T. exiguus* were more abundant in the southern zone (Table 3).

A multivariate analysis used for the separation of *B. b. gracilis* between localities showed that the first discriminant functions explained 100% of the variance (eigenvalue = 0.120). There was a significant overall effect of group (Wilks' lambda = 0.893, F₄₄₆; p < 0.01). Individual fishes were distributed mainly along the first axe. Using dimensionality tests showed that the two studied zones were significantly separated in both dimensions ($\chi^2 = 50.676$, d.f. = 4, p < 0.01). Each fish was classified correctly to the two component communities with an accuracy of 64.3% (Table 4). Scores of individual fishes showed a clear discrimination between two groups, one including fishes from the center and one including specimens from the south. The importance of each parasite species with respect to discrimination between groups, evaluated as the contribution of each variable to the total sum Mahalanobis distances, showed that Digenea *L. retroflexus*, *A. laguncula* and Nematoda Anisakid larvae were the most important species in determining the position of samples from the central zone. Metacercaria *C. pilodora* was characteristic of specimens from the south.

3.2. Seasonal Variation

A multivariate analysis used for the separation of *B. b. gracilis* between seasons in both localities showed that the first two discriminant functions explained 79.8% of the variance, contributing to 51.4% (eigenvalue = 0.448) and 28.4.3% (eigenvalue = 0.247), respectively. There was a significant difference between groups (Wilks' lambda = 0.467; $F_{8.408}$; p < 0.01) (Figure 2(a)). Dimensionality tests showed that seasons were significantly separated in both dimensions (χ^2 =328.482, d.f = 42, p < 0.01). Results showed a clear discrimination between

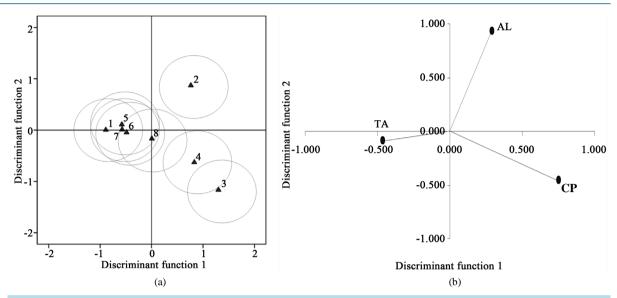


Figure 2. (a) Sample scores of the first two discriminant functions for specimens of *B. b. gracilis* from center and south/ seasons: 1, samples from center in Autumn; 2, samples from center in summer; 3, samples from center in winter, 4 samples from center in spring; 5, samples from south in Autumn; 6, samples from south in summer; 7, samples from south in winter; 8, samples from south in spring. Lettering on the axes represent groups averages surrounded by circles 90% to show tolerances; (b) canonical correlations between the first tow discriminant functions and three parasites species between zone/seasons; *Aponurus laguncula* (AL), *Tergestia acanthocephala* (TA) and *Condylocotyla pilodora* (CP).

	Prevalence			Mean abundance				
	Center	South	χ²	р	Center	South	t	р
Axine belones	61	58	0.07	0.79	2.65	2.50	0.83	0.40
Condylocotyla pilodora (metacercaria)	56	46	0.92	0.34	0.76	1.50	4.32	0.00^{*}
Lecithostaphylus retroflexus	19	4	8.63	0.00^{*}	0.35	0.08	4.84	0.00^{*}
Tergestia acanthocephala	20	27	0.88	0.35	0.87	0.19	-1.56	0.11
Aponurus laguncula	20	5	7.82	0.01^*	34.70	10.48	4.97	0.00^{*}
Bomolochus bellones	7	7	0.00	0.95	0.36	0.24	0.86	0.38
Irona nana	5	9	0.64	0.42	0.08	0.40	-1.72	0.08
Telosentis exiguus	6	1	2.32	0.12	0.05	0.10	2.30	0.02^*
Anisakid larvae	5	12	1.86	0.16	0.19	0.07	-2.60	0.01^{*}

Table 3. Comparison of prevalence and mean abundance of selected metazoan parasites from *B. b. gracilis* among two zones in eastern Tunisian Coast.

 Table 4. Discriminant analysis classifications showing the numbers and percentages of fish classified in each zone (rows correspond to group memberships).

	Center	South	Per cent
Center	96	107	52.7%
South	54	194	78.2%

three groups. One including fishes from the south in all seasons and fishes from the center in autumn, one including those from the center in winter and spring and one including samples from the center in summer. *T. acanthocephala* was the most important species in determining samples from the first group. *C. pilodora* was the most important species identifying specimens of winter and spring from the second group. *A. laguncula* characterized specimens from the third group (Figure 2(b)).

4. Discussion

The present study increases our knowledge on the metazoan parasite of the Mediterranean subspecies *B. b.* gracilis. This is the first survey on the entire parasite fauna of *B. b. gracilis* from southern shore of the Mediterranean Sea. Previous records of parasites from *B. b. gracilis* come from taxonomic studies of individual parasite species such as cestod *Ptychobothrium belones*; copepod *B. bellones*; digeneans *L. retroflexus*, *T. acanthocephala*, *A. laguncula* and *C. pilodora* metacercaria; isopods *I. nana*; monogenean *A. belones* and acanthocephalan *T. exiguus* [2]-[11]. In the central Mediterranean Sea, Radujković and Šundić (2014) listed six parasite species on *B. b. gracilis* (Table 5). In our study, remaining parasite species excepting cestod were found on *B. b.*

Host	Host Belone belone gracilis			Belone belone belone			
Locality	Adriatic Sea	Tunisian coast	Baltic sea	Southern Baltic sea	Courtmacsherry Bay, Co Cork, Ireland	Çanakkale Coasts, Turkey	
Parasites/author	Radujković and Šundić (2014)	Present study	Rynkiewicz (1970)	Grabda (1981)	Dorman (1991)	Oktener (2005), Keser <i>et al.</i> (2007)	
Monogenea	Axine belones	Axine belones	-	-	-	Axine belones	
Digenea	-	Condylocotyla pilodora	Diplostomum spathaceum	-	-	Lecithostaphylus retroflexus	
	Lecithostaphylus retroflexus	Lecithostaphylus retroflexus	Tylodelphys sp.	-	-	-	
	Theledera acanthocephala	Tergestia acanthocephala		-	-	-	
	Aponurus laguncula	Aponurus laguncula		-	-	-	
Cestoda	Lacistorhynchus tenuis	-	Scolex pleuronectis	Lacistorhynchus tenuis	-	-	
	Ptychobothrium belones	-	-	-	-	-	
Copepoda	-	Bomolochus bellones	-	-	Caligus belones	-	
	-		-	-	Caligus elongatus	-	
					Caligus pelamydis	-	
Isopoda	-	Irona nana	-	-	Nerocila orbygini	-	
	-	Nerocila orbygini	-	-	-	-	
Acanthocephala	-	Telosentis exiguus	Pomphorynchus laevis	Pomphorynchus laevis	-	-	
	-	-	Neoechinorhynchus rutili	Neoechinorhynchus rutili	-	-	
	-	-	-	Echinorhynchus gadi	-	-	
Nematoda	-	Hysterotylacium sp.	Contracaecum aduncum	Anisakis simplex	-	-	
	-	Contracaecum sp.	Cuculanus truttae	Thynnascaris adunca	-	-	

Table 5. Comparaison of parasitism of B. belone subspecies.

gracilis alongside specimens of nematod Anisakid larvae and isopod *N. orbignyi*. Latest species have been reported from *B. b. belone* in the north east Atlantic by Dorman (1991) [35], but not found previously in Belonids from Tunisian coast. *N. orbignyi* was collected attached mainly to fins of Mugilids, rarely on *Solea solea* Linnaeus, 1758 and *Dicentrarchus labrax* Linnaeus, 1758 [9]. The low value of prevalence of *N. orbignyi* (1%) compared to that reported by Cheikhrouha *et al.* (2000) in Mugilids (47%) calls for an accidental infestation.

A comparison of the parasite fauna composition of B. b. gracilis with those of B. b. euxeni revealed similarity between parasite species of the Mediterranean and the Black Sea subspecies of B. belone (Table 5). Indeed, B. b. euxeni from the Black Sea hosted A. belones and L. retroflexus [13] [14]. These parasite species could be with a wide distribution, present in the Mediterranean and the Black Seas. Nevertheless, the metazoan parasite fauna of B. b. gracilis was totally different from that reported on B. b. belone in the north east Atlantic [15] [16]. Indeed, ten parasites taxa can be identified as proper to the Mediterranean Sea subspecies B. b. gracilis and seven parasites taxa can be identified as proper to the Atlantic subspecies B. b. belone were founded. It appears that variation in infection with parasites on *B. belone* could be influenced by geographical distribution of these different subspecies of hosts or by specificity of parasites in the host subspecies. B. b. gracilis, B. b. euxeni and B. b. belone are recognized by Collette (2003) as different subspecies that lives in completely different and separate ecosystems [1]. More detailed studies involving the qualitative and the quantitative aspect of the metazoan parasites of B. b. euxeni are necessary for providing useful information on the use of parasites for separating subspecies of *B. belone*. These results could reinforce the usefulness of parasites as adequate tools to clarify the taxonomic status of their hosts. In fact, parasites have been used successfully not only as tools to understand the population structure of the host species but also to clarify doubtful systematic relations between closely related hosts [36]. In this context, Oliva and Ballon (2002) evaluated also the use of metazoan parasites of the hake, Merlucciuus gayii (Guichenot, 1848) as biological tags for stocks identification of fish subspecies [37]. Indeed, result of examined specimens of Chilean hake subspecies Merluccius gayi gayi (Guichenot, 1848) in the central and southern Chile as well as analysis of previously published data on Peruvian hake subspecies Merluccius gayi peruanus (Guichenot, 1848) from Peru revealed presence of 12 species found only in the Chilean hake and seven species exclusively found in the Peruvian hake. Thus, differences in infection with parasite can be due to the geographic distribution of the host; phylogeny and also to variation of biological and ecological factors [38]-[42]. These factors can act individually or together altering the parasite distribution patterns in their hosts [43].

Our findings showed that Digenea were the most abundant parasites in the viewpoint of species richness (number of parasites species) and number of individuals collected from *B. b. gracilis*. Diversity of digenean parasites appears to be an index of the diversity of intermediate hosts ingested and therefore a varied diet of the host. Garfish *B. belone* is carnivorous based mainly on crustaceans (brachyura, copepod and decapod), followed by teleost fish and other items of terrestrial origin, including insects [44] [45]. Indeed, piscivorous fish occupy the top of the food web and may present more chance of infection [46]. Among digenean parasites, metacercaria *Condylocotyla pilodora* is the most abundant species found on *B. b. gracilis*. The high infection with *C. pilodora* could be explained by the long life of encysted parasites which may be longer than those of intestinal parasites. Indeed, encysted parasites, mainly in muscles and abdominal cavity cannot be released from the host and are therefore accumulated in the organism [47]. The Shoaling behaviour and the pelagic mode of life of this Belonida could also explain the success of *C. pilodora* to reach its definitive host. The life-cycle of *C. pilodora* involves the first intermediate hosts a snail *Cerithium vulgatum* Bruguière, 1792, the metacercariae in the pericardial sac of the second intermediate hosts a fish *Belone belone*, and adults in the definitive host a herring gull *Larus argentatus* [7].

Nine species of parasites among ten, reaching a prevalence > 5% in at least one of the studied zones, were selected as biological tags. The significant differences of both infection parameters prevalence and mean abundance between areas and results of discriminant analysis demonstrate the potential value of selected parasites as biological tags. Two stocks can be identified, one in the central and one in the southern Tunisian coast.

Distribution patterns of parasites showed difference between fishes from the central zone and those from the southern zone. Indeed, specimens of *B. b. gracilis* from the center were characterized by the highest abundance of Adult Digenea *A. laguncula* and *L. retroflexus*. Whereas, specimens from the southern coast off Tunisia were characterized by the highest abundance of Digenea metacercaria *C. pilodora*. It appears that central coast of Tunisia is probably the endemic area of the first intermediate host (mollusc) of *A. laguncula* and *L. retroflexus* but in the southern coast environmental conditions can be more suitable for transmission of the first intermediate

host of the digenean metacercaria *C. pilodora* which is the snail *Cerithium vulgatum* that prefers areas of low wave intensity [48]. This could reflect the different oceanographic characteristics of both ecosystems. Central coast off Tunisia (Gulf of Hammamet) is characterized by the existence of a relatively strong and continuous current (Atlantic), which would be added to littoral currents and combined to currents generated by the wind [49]; whereas in the southern coast (Gulf of Gabes) characterized by the extended continental shelf, the Atlantic current losing influence due to the presence of local permanent cyclonic circulation and hydrodynamics generally under the control of winds in deep areas and tidal along the coastline [50]-[52].

Specimens of *B. b. gracilis* from the central coast were characterized by the highest abundance of Nematod Anisakid larvae. This difference could be related to quantitative and qualitative change of the diet of *B. b. gracilis* between the center and the south off Tunisia and therefore variation in the intermediate host's involved. Transmission pathways of ascaridoid Nematoda are habitat-dependent and usually involve a broad spectrum of invertebrates and intermediate or paratenic fish hosts [53]. Several larval stages could therefore be affected differentially by environment factors [54]. The paucity of Anisakid larvae in the south may be also a consequence of the unstable environment due to the pollution by various industrial and urban activities in the Gulf of Gabes which disrupts the stability of the ecosystem. The damage of this ecosystem can generate consequently a variation of the intermediate host's distribution (invertebrates, fish).

These results were corroborated by those of discriminant analysis which showed that fishes from the two areas represented two units with a mean accuracy of correctly classifying fishes to their respective zone (64.3%). Fishes from the south were more classified in their origin area with an accuracy of (78.2%) than fishes from the centre (52.7%). This is the result of distance separating samples, a common phenomenon that usually emerges as the best predictor of similarity in the analyses of parasites communities in relation to characteristics of the habitat [55] [56]. Results of geographic variation can be confirmed by those from discriminant analysis of seasonal variation in both localities. Indeed, specimens from the south were clumped together as a single stock. Fishes from the center were separated in three groups according to seasons where Digenea T. acanthocephala and A. laguncula were the most responsible for these differences. Seasonal variation of Digenea with short life span could be explained by the seasonal variation of the host's diet. The change in the diet composition may be associated with ecological characters of hosts such as feeding behavior and seasonal changes [47]. Additionally, seasonal variability of the distribution of the fish hosts can be related to the seasonal variation of the activity of the fishing fleets. The accuracy of classifying fishes incorrectly in their origin zone can be explained by migratory movements of B. b. gracilis between inshore and offshore waters in search of optimal conditions for living, feeding and breeding. Parasites can be also used to follow seasonal migrations of fish to and from spawning and feeding grounds [57]. In the Baltic Sea, parasites have been successfully used by Grabda (1981) for stock discrimination of the garfish B. belone. B. belone spawn in areas of low salinity and migrate seasonally to feeding grounds farther west in areas of higher salinity. These migratory stocks can be identified on the spawning grounds by their infections of the characteristically marine anisakid nematode larvae, adult hemiurid digeneans and trypanorynch metacestodes.

5. Conclusion

In summary, a complete list of metazoan parasites of *B. b. gracilis* from Tunisian coast was established for the first time. Findings of different parasite taxa in the Mediterranean subspecies *B. b. gracilis* compared to those of the Atlantic subspecies *B. b. belone* could reinforce the usefulness of parasites as adequate tools to clarify the taxonomic status of their hosts. Studies of spatial and seasonal variability of metazoan parasites in the garfish *B. gracilis* off Tunisia allowed for the identification of discrete stocks in the central coast characterized by Nematoda Anisakid larvae and in the southern coast characterized by digenean metacercaria *Condylocoltyla pilodora*.

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