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SHORT COMMUNICATION

Identifying copepod functional groups from species functional traits

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We gathered information on the functional traits of the most representative copepod species in the Mediterranean Sea. Our database includes 191 species described by 7 traits encompassing diverse ecological functions: minimal and maximal body length, trophic group, feeding type, spawning strategy, diel vertical migration and vertical habitat. Cluster analysis in the functional trait space revealed that Mediterranean copepods can be separated into groups with distinct ecological roles.

KEYWORDS: copepods; functional trait; zooplankton; Mediterranean Sea

Functional traits are phenotypic characteristics of organisms that impact their fitness and are relevant to ecosystem function (Violle *et al.*, 2007). For zooplankton, traits can be classified according to ecological functions feeding, growth/reproduction, survival— and types morphological, physiological, behavioural, life history (Litchman *et al.*, 2013). As organisms have to allocate energy among ecological functions, traits relate to one another through trade-offs (Kiørboe *et al.*, 2015). Litchman *et al.* (Litchman *et al.*, 2013) recently advocated the implementation of a comprehensive matrix of key functional traits for zooplankton taxa. However, these functional traits have mainly been documented from laboratory experiments, thereby only encompassing a small number of species (Barnett *et al.*, 2007; Saiz and Calbet, 2007; Barton *et al.*, 2013; Kiørboe *et al.*, 2015). For application in marine ecology, traits are needed at the species level and for the largest possible number of species, which requires the gathering of a huge amount of specific information. Such a challenge should be tackled step by step, focusing on some taxa and/or ocean basins at first. Functional traits could be used to gather species with

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similar traits into functional groups (i.e. sets of species with similar effects on ecosystem functions and/or similar response to environmental conditions; Gitay and Noble, 1997) or to describe functional diversity of zoo-plankton communities (Barnett *et al.*, 2007; Vogt *et al.*, 2013; Pomerleau *et al.*, 2015). Identifying and describing zooplankton functional groups should then increase our understanding of zooplankton ecological roles in marine ecosystems.

In this study, we developed a trait database for the most commonly sampled and abundant Mediterranean copepod species. Copepods are abundant in marine pelagic ecosystems (e.g. Siokou-Frangou *et al.*, 2010) where they constitute the main trophic link between primary producers and higher trophic levels, such as small pelagic fishes (Costalago *et al.*, 2015). Copepods are also relatively well documented in terms of distribution and biology (Razouls *et al.*, 2005–2015). From this trait database, the aim of this study was to estimate how many functional groups could be identified among Mediterranean copepods, to characterize them and to discuss their ecological significance.

We considered 191 copepod species (Table I) that are the most representative of the Mediterranean copepod communities, in terms of both abundance and presence (Siokou-Frangou *et al.*, 2010; Mazzocchi *et al.*, 2014). More details on how this list of species has been gathered are available in Supplementary data, Material S1. We used the following traits covering various types and ecological functions (Litchman *et al.*, 2013), known to be ecologically meaningful (Kiørboe and Sabatini, 1994; Kiørboe, 2011; Kiørboe *et al.*, 2015) and commonly used for zooplankton (Barnett *et al.*, 2007; Barton *et al.*, 2013; Pomerleau *et al.*, 2015):

- two morphological traits which relate to many ecological traits: minimum and maximum adult body (cephalothorax) length (mm);
- one physiological trait defining the species' trophic group (Carnivore, Omnivore–Carnivore, Omnivore, Omnivore–Herbivore, Omnivore–Detritivore);
- one behavioural trait depicting feeding strategy, classified into three classes (Kiørboe, 2011): active ambush feeding, cruise feeding and filter feeding, mixed feeding (for species that can switch between the three strategies);
- one life history trait related to reproduction, defining the egg-spawning strategy (broadcast-spawner, sac-spawner);
- one behavioural trait related to diel vertical migration (DVM) behaviour, classified into four classes according to the intensity of the observed migration: Non-migrant, Weak migrant (DVM occurs within tens of metres), Strong migrant (over several hundreds of metres), Reverse migrant (for species that migrate deeper at night).

All body sizes were obtained from Razouls *et al.* (Razouls *et al.*, 2005–2015), whereas the other traits were obtained from an extensive literature review (see Supplementary data, Material S4 and Table SII for the full list of references). Additionally, to discuss the potential role of each functional group in the pelagic ecosystem, the species' preferential depth layer was established (epi-/meso-/bathypelagic). We were able to determine at least 5 of the 7 functional traits for 171 species. The trait database for the 191 copepod species is available as Supplementary Table SII and can also be downloaded from PANGAEA (http://doi.pangaea.de/10.1594/PANGAEA.854331).

In order to identify functional groups, we performed a multiple correspondence analysis (MCA) on the trait matrix. MCA is an ordination method in reduced space for the multivariate analysis of categorical variables (Husson et al., 2010). The computation of the MCA functional space was performed on four traits: class of maximum body length (Size_1: 0.501.80 mm, Size_2: 1.89-2.85 mm, Size 3: 3.00-5.70 mm, Size 4: 6.10-11.0 mm), binary trophic group (Carnivore, Omnivore, Herbivore, Detritivore), feeding type and spawning strategy. Indeed, the minimum body length was highly correlated to the maximum body length ($R^2 = 0.866, n = 191$) and DVM behaviour was not taken into account since it tends to be very plastic for most species, meaning that it is known to vary greatly according to the environmental fluctuations and species' ontogeny (see Pomerleau et al., 2015). A preliminary MCA incorporating these traits showed they had no impact in the definition of functional groups. Species for which the four traits were not fully defined were used as supplementary objects, meaning that they are associated with a group a posteriori, from their informed traits (see Supplementary data, Material S2 for more details on the MCA). The Euclidean distance among the 191 species in the functional space was computed using their coordinates along the four significant axes of the MCA (70.77% of the variance). Hierarchical agglomerative clustering was performed on this distance matrix using a synoptic aggregation method (Ward's; Husson et al., 2010). Depending on the cutting level, two, three or six clusters could be identified (Fig. 1). The first level distinguished species according to the trophic group (carnivore vs. non-carnivore). Among non-carnivore species, the second level separated broadcasters from sacspawners. Then, each of these groups was divided into two subgroups with different size and/or feeding type. Since higher cut levels could not be clearly related to functional traits, six functional groups were retained (Tables I and II). We will now detail each functional group and discuss their ecological role in the Mediterranean pelagic food web.

1		2		3		4		5		6	
Number	Species	Number	Species	Number	Species	Number	Species	Number	Species	Number	Species
5	Aetideopsis armata	6	Aetideus armatus	1	Acartia clausi	8	Anomalocera patersoni	69	Diaixis pygmaea	45	Clausocalanus arcuicornis
10	Augaptilus longicaudatus	7	Aetideus giesbrechti	2	Acartia danae	12	Calanoides carinatus	70	Disco minutus	46	Clausocalanus furcatus
11	Augaptilus spinifrons	29	Candacia bispinosa	3	Acartia discaudata	13	Calanopia elliptica	71	Distioculus minor	47	Clausocalanus jobei
27	Candacia armata	31	Candacia giesbrechti	4	Acartia negligens	14	Calanus helgolandicus	96	Homeognathia brevis	50	Clausocalanus parapergens
28	Candacia bipinnata	34	Candacia simplex	9	Archescolecithrix auropecten	15	Calocalanus adriaticus	113	Monothula subtilis	51	Clausocalanus paululus
30	Candacia ethiopica	35	Candacia tenuimana	37	Centropages chierchiae	16	Calocalanus contractus	118	Neomormonilla minor	52	Clausocalanus pergens
32	Candacia longimana	36	Candacia varicans	38	Centropages furcatus	17	Calocalanus elegans	119	Oithona atlantica	79	Euchirella messinensis
33	Candacia norvegica	56	Corycaeus anglicus	39	Centropages hamatus	18	Calocalanus elongatus	120	Oithona brevicornis	80	Euchirella rostrata
54	Copilia quadrata	57	Corycaeus brehmi	40	Centropages kroyeri	19	Calocalanus longisetosus	121	Oithona decipiens	81	Euchirella truncata
75	Euchaeta acuta	58	Corycaeus clausi	41	Centropages ponticus	20	Calocalanus neptunus	122	Oithona linearis	82	Euterpina acutifrons
76	Euchaeta marina	59	Corycaeus flaccus	42	Centropages typicus	21	Calocalanus pavo	123	Oithona longispina	85	Goniopsyllus rostratus
77	Euchaeta media	60	Corycaeus furcifer	43	Centropages violaceus	22	Calocalanus pavoninus	124	Oithona nana	97	lsias clavipes
78	Euchaeta spinosa	61	Corycaeus giesbrechti			23	Calocalanus plumatus	125	Oithona parvula	106	Macrosetella gracilis
86	Haloptilus acutifrons	62	Corycaeus latus			24	Calocalanus plumulosus	126	Oithona plumifera	107	Mecynocera clausi
87	Haloptilus angusticeps	63	Corycaeus limbatus			25	Calocalanus styliremis	128	Oithona similis	109	Microcalanus pygmaeus
89	Haloptilus mucronatus	64	Corycaeus minimus			26	Calocalanus tenuis	129	Oithona tenuis	110	Microsetella norvegica
90	Haloptilus ornatus	65	Corycaeus ovalis			44	Chiridius poppei	130	Oithona vivida	111	Microsetella rosea
91	Haloptilus oxycephalus	66	Corycaeus speciosus			48	Clausocalanus lividus	188	Vettoria granulosa	127	Oithona setigera
92	Haloptilus tenuis	67	, Corycaeus typicus			49	Clausocalanus mastigophorus	189	Vettoria longifurca	131	Oncaea curta
93	Heterorhabdus abyssalis	72	Euaugaptilus hecticus			53	Copilia mediterranea	190	Vettoria parva	132	Oncaea media
95	Heterorhabdus spinifrons	83	Farranula carinata			55	Copilia vitrea			133	Oncaea mediterranea
143	Paraeuchaeta hebes	84	Farranula rostrata			68	Ctenocalanus vanus			134	Oncaea ornata
144	Paraeuchaeta norvegica	88	Haloptilus longicornis			73	Eucalanus elongatus			135	Oncaea scottodicarloi

Table I: List of the 191 Mediterranean copepod species whose traits have been described

Continued

Table I: Continued

Functional group

4

1		2		3		4		5		6	
Number	Species	Number	Species	Number	Species	Number	Species	Number	Species	Number	Species
147	Phaenna spinifera	94	Heterorhabdus papilliger			74	Eucalanus hyalinus			136	Oncaea venusta
158	Sapphirina angusta	99	Lubbockia aculeata			98	Labidocera wollastoni			137	Oncaea waldemari
160	Sapphirina gemma	100	Lubbockia squillimana			101	Lucicutia clausi			142	Paracartia latisetosa
161	Sapphirina intestinata	138	Pachos punctatum			102	Lucicutia flavicornis			155	Pseudocalanus elongatus
163	Sapphirina metallina	145	Parapontella brevicornis			103	Lucicutia gaussae			170	Scaphocalanus curtus
164	Sapphirina nigromaculata	153	Pontellina plumata			104	Lucicutia gemina			171	Scaphocalanus invalidus
165	Sapphirina opalina	154	Pontellopsis villosa			105	Lucicutia ovalis			172	Scolecithricella abyssalis
166	Sapphirina ovatolanceolata	159	Sapphirina auronitens			108	Mesocalanus tenuicornis			173	Scolecithricella dentata
167	Sapphirina sali	162	Sapphirina lactens			112	Monacilla typica			174	Scolecithricella tenuiserrata
168	Sapphirina scarlata	169	Sapphirina vorax			114	Mormonilla phasma			175	Scolecithricella vittata
						115	Nannocalanus minor			176	Scolecithrix bradyi
						116	Neocalanus gracilis			177	Scolecithrix danae
						117	Neocalanus robustior			178	Spinocalanus abyssalis
						139	Paracalanus denudatus			179	Spinocalanus Iongicornis
						140	Paracalanus nanus			183	Triconia conifera
						141	Paracalanus parvus			184	Triconia dentipes
						146	Pareucalanus attenuatus			185	Triconia minuta
						148	Pleuromamma abdominalis			186	Triconia similis
						149	Pleuromamma borealis			187	Triconia umerus
						150	Pleuromamma gracilis			191	Xanthocalanus agili
						151	Pleuromamma xiphias				0
						152	Pontella mediterranea				
						156	Rhincalanus cornutus				
						157	Rhincalanus nasutus				
						180	Subeucalanus crassus				
						181	Subeucalanus monachus				
						182	Temora stylifera				

The species are numbered by alphabetical order but gathered by functional groups, as revealed by the hierarchical clustering on the first four axes of the MCA on functional traits (see Fig. 1). The species with bold numbers are the 99 species used in the MCA space calculation.

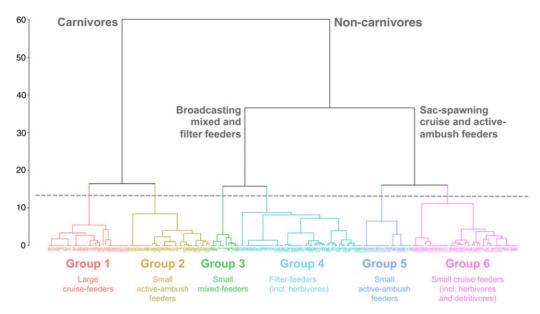


Fig. 1. Identification of functional groups among the 191 most representative copepod species of the Mediterranean Sea from hierarchical clustering on the first four axes of the MCA based on four functional traits (class of maximum body length, binary trophic group, feeding type and spawning strategy). Depending on the cutting level, two, three or six clusters could be identified. The first level distinguished species according to trophic group (carnivore vs. non-carnivore). Among non-carnivore species, the second level discriminates broadcasters vs. sac-spawners. Then, each of these groups can be divided into two subgroups with different size and/or feeding type. Since higher cut levels could not be clearly related to functional traits, six functional groups were retained (Table II). Copepod species are indicated by numbers (Table I).

Group 1 is composed of large carnivores that primarily feed by cruising. There is no unique reproductive strategy, though the species are mainly sac-spawners. Many species have a broad vertical distribution, ranging from the epipelagic to the bathypelagic. Representative genera are Calanoids such as *Candacia* spp., *Haloptilus* spp., *Heterorhabdus* spp., members of the Euchaetidae family and Cyclopoids of the *Sapphirina* genus. These species are known to prey on smaller copepods, as well as other zooplanktonic taxa, such as doliolids (Takahashi *et al.*, 2013), larvaceans (Ohtsuka and Onbé, 1989) and even fish larvae (Yen, 1987).

Group 2 is defined by smaller carnivore species, all active ambush feeders and mainly sac-spawners, belonging to the Corycaeidae family. These are small visual predators that prey on nanozooplankton, nauplii, younger stages of copepodites through active ambush tactics (Landry *et al.*, 1985), on wide depth intervals. Together, Groups 1 and 2 mainly contribute to the top-down control of mesozooplankton, including the other copepod functional groups.

Group 3 gathers a lesser number of species and consists of Calanoids of the genera *Centropages* and *Acartia*. They are small omnivorous broadcasters, but phytoplankton can become an important component of their diet. They exhibit mixed feeding strategies, depending on the available food items. *Acartia* spp. and *Centropages* spp. are generally restricted to the epipelagic and are affiliated with neritic environments (Siokou-Frangou *et al.*, 2010).

Group 4 is the largest group and comprises almost all filter-feeding species, spanning all size classes, with a clear tendency towards herbivory. The species of this group for which reproductive strategy could be found were mainly broadcasters. This group contains not only small-bodied calanoids that are numerically very important in the Mediterranean epipelagic (Clausocalanus spp., Calocalanus spp., Temora stylifera; Mazzocchi et al., 2014), but also larger calanoids, some of which are strong vertical migrants, such as Calanus helgolandicus, Pleuromamma spp. or Neocalanus spp. (Andersen et al., 2001, 2004). The small surface calanoids are the target prey for larval and juvenile pelagic fish (Borme et al., 2013; Costalago et al., 2015), whereas larger calanoids are the preferential prey of Mediterranean mesopelagic fishes (Palma, 1990). Additionally, the strong calanoid migrants might play a differential key role in carbon cycling as they graze upon microalgae in the euphotic zone, and then migrate below the permanent thermocline where they excrete their lipid reserves (lipid pump hypothesis; Jónasdóttir et al., 2015). Therefore, Groups 3 and 4 are crucial for the transfer of energy from photoautotrophs to higher trophic levels, both in neritic (Group 3) and in oceanic environments (Group 4). Also, the latter group might comprise species that play a potentially underestimated role for the carbon flux (Jónasdóttir et al., 2015).

Group 5 consists essentially of *Oithona* spp. These are small active ambush-feeding omnivores that carry their

		Functiona	Tatal assessments an						
Functional trait	Category	Group 1	Group 2	Group 3	Group 4	Group 5	Group 6	Total number of species	
Class of maximum	Size_1 (0.50–1.80 mm)	0	12	6	16	20	32	86	
body length	Size_2 (1.89–2.85 mm)	0	20	6	17	0	8	51	
	Size_3 (3.00–5.70 mm)	30	1	0	10	0	1	42	
	Size_4 (6.10–11.0 mm)	3	0	0	7	0	2	12	
Trophic group	Carnivore	25	29	0	0	0	0	54	
	Omnivore	0	0	6	12	11	4	33	
	Omnivore-carnivore	6	1	1	0	0	0	8	
	Omnivore-detritivore	0	0	0	0	0	28	28	
	Omnivore-herbivore	0	0	4	36	0	11	51	
	NA	2	3	1	2	9	0	17	
Feeding type	Active ambush	0	16	0	0	11	1	28	
	Cruise	11	5	0	3	0	15	34	
	Filter	4	0	0	43	0	8	55	
	Mixed	0	0	11	0	0	2	13	
	NA	18	12	1	4	9	17	61	
Spawning strategy	Broadcaster	7	7	11	24	0	3	52	
	Sac-spawner	16	20	0	3	17	28	84	
	NA	10	6	1	23	3	12	55	
Type of DVM	No DVM	13	8	6	13	2	15	57	
	Weak DVM	0	1	0	0	2	2	5	
	Strong DVM	2	1	0	8	0	2	13	
	Reverse DVM	5	12	2	15	2	14	50	
	NA	13	11	4	14	14	10	66	
Vertical distribution	Epipelagic (0–200 m)	10	2	11	21	7	14	65	
	Epimesopelagic (0–1000 m)	11	14	0	17	7	13	62	
	Epibathypelagic (0–4000 m)	11	15	0	9	2	9	46	
	Mesopelagic (200–1000 m)	0	1	0	0	0	2	3	
	Mesobathypelagic (200–4000 m)	1	0	1	3	2	4	11	
	NA	0	1	0	0	2	1	4	
Mean minimum body	2.27	1.18	1.07	1.47	0.67	0.93	-		
Mean maximum body	4.47	2.03	1.82	2.95	1.11	1.74	-		
Total number of speci		33	33	12	50	20	43	191	

Table II:	Traits charac	terization of	f the six	identified	functional	groups	(Fig.	<u>1</u>)

The number of species recorded within each trait's class and within each functional group is reported. The groups are based on hierarchical clustering on the first four axes of the MCA based on four functional traits: class of maximum body length, binary trophic group, feeding type and spawning strategy. For information, the type of DVM, the vertical distribution, the average minimum body length (mm) and the average maximum body length (mm) are also indicated (in grey).

NA, not available; DVM, diel vertical migration.

eggs. It is difficult to assign a particular ecological function to such a group, since Oithona spp. are a major component of the global ocean's plankton, independently of environmental conditions (Gallienne and Robins, 2001). Feeding and trophic group were unknown for the other species of Group 5. Therefore, they are related to Oithona spp. only because of they are small (<1.8 mm) sacspawners. Group 6 also comprised small sac-spawning omnivores, but these are mainly cruising detritivores (Oncaea spp., Microsetella spp.) or herbivores (Clausocalanus spp.). The former usually exhibit a wide vertical distribution, while the latter are epipelagic. Oncaea spp. and Microsetella spp. are known associates of appendicularian houses (Alldredge, 1972; Steinberg et al., 1994). Several calanoid species of this group (Euchirella spp., Scolecitrichidae) are also deepwater detritivores. Consequently (and as suggested by a higher cut level on dendrogram Fig. 1), Group 6 could be sensibly divided into two subgroups: (i) deep-water dwelling detritivores that actively participate in the recycling of particulate organic matter and (ii) small cruising grazers contributing to epipelagic secondary production.

By focusing on Mediterranean copepods, we were able to gather information on functional traits for 191 species, with at least 6 traits described for 135 species and 7 for 66 species. Using 4 of these traits, functional groups with different ecological roles were described. Although the definition of these groups was robust (similar groups were found using a K-means partitioning method, or accounting for all traits), the main limitation, as for any traitbased approach, remains the scarcity of trait descriptions at the species level. Compiling a trait database for a larger number of zooplankton species remains challenging, but the present initiative demonstrates the usefulness of this endeavour.

The present study also confirmed or revealed tradeoffs among zooplankton functional traits (Litchman *et al.*, 2013, Kiørboe, 2011). For instance, small carnivores and small omnivores were active ambush feeders, while large

carnivores were cruise feeders. This relationship between size and feeding strategy could be explained by the differences in metabolic requirements. Indeed, whereas cruise feeders have to swim actively to encounter their prey, ambush feeders passively encounter them (Kiørboe, 2011), which requires less energy. As metabolic rates scale with body size (Kiørboe and Hirst, 2014), this difference in metabolic requirement could explain their difference in size. We also found that sac-spawners were active ambush feeders or cruise feeders, whereas broadcasters were mainly filter feeders or mixed feeders. This could be due to optimal resource allocation: egg-carrying ambush-feeding copepods have longer lifespans and lower fecundity rates than broadcasting active feeders (Kiørboe and Sabatini, 1994; Kiørboe et al., 2015). We also found that three guarters of the carnivorous species were sac-spawners; hence, these exhibit a higher degree of parental care and avoid predation on their own eggs. Conversely, for epipelagic omnivorous filter feeders, which are heavily preved upon by other zooplankters and fishes, broadcasting is likely to be favoured to avoid being eaten together with the eggs, as an adaptation to the elevated mortality of ovigerous females (Kiørboe and Sabatini, 1994). Together, these results call for a better understanding of the mechanistic processes that lead to such trade-offs across traits.

The functional trait database compiled here can be used to estimate the functional diversity of zooplankton communities (Vogt *et al.*, 2013; Pomerleau *et al.*, 2015) and test whether traits can be related to environmental variables (Barton *et al.*, 2013). Finally, the existence of several functional groups encourages the integration of more diverse planktonic assemblages in ecosystem models.

SUPPLEMENTARY DATA

Supplementary data can be found online at http://plankt. oxfordjournals.org.

DATA ARCHIVING

The trait database for the 191 copepod species can be downloaded from PANGAEA (http://doi.pangaea.de/10.1594/PANGAEA.854331).

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