

How do harpacticoid copepods colonize detrital seagrass leaves?

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Abstract An experiment was carried out investigating the colonization ability and specific pattern of copepods towards a provisional benthic habitat. Since copepods are known to disperse passively and actively, the experiment aimed to investigate the pool of colonizers of macrophytodetritus and the species-specific active colonization pathways. The experiment was performed in a Mediterranean seagrass *Posidonia oceanica* meadow on defaunated macrophytodetritus accumulations (mainly dead seagrass leaves) for two time intervals (24 and 96 h). Active colonization by copepods, independently of their adjacent potential source pool habitat (bare sandy sediments, *P. oceanica* canopy, water column and macrophytodetritus) occurred within 24 h. Natural densities (as in the control treatments) were only reached by active colonization through the water column. Both neither diversities nor species composition of natural macrophytodetritus were ever reached by one single migratory pathway, therefore only a combination of interstitial migration and water column migration can explain the species occurrence under natural condition. Moreover, every potential adjacent source pool habitat contributed species to the newly colonized macrophytodetritus. However, the main colonizers were mostly species with good swimming capabilities. The diverse pool of species present in the newly colonized macrophytodetritus underlines the

complex communities and dispersion capabilities of copepods. Hence, macrophytodetritus possesses the potential ability to be a colonizer source pool for every adjacent habitat and thus behaves as a copepod hub for the entire seagrass ecosystem.

Introduction

Dispersion and colonization of new habitats by meiofauna (i.e. the benthic fauna belonging to size class 38 μm –1 mm) are highly variable in space and time. This variability is caused by complex interactions between habitat structures, species-specific biological traits, hydrodynamics, resource availability, predation pressure and environmental deterioration (Armonies 1994; Commito and Tita 2002; Bostrom et al. 2010). Since meiofauna lack planktonic larvae (Hagerman and Rieger 1981; Huys and Boxshall 1991), the dispersion mode of adults might be crucial for population dispersion. On small scales (up to a metre), early studies revealed horizontal migration through the sediments interstices as primary mean for meiofauna colonization (i.e. infaunal migration) (McIntyre 1969). However, most of the meiofauna lives on the sediment–water interface being capable of active swimming and infaunal migration on short distances during low water flow (Fleeger et al. 1995). These are thus susceptible to passive erosion and are therefore classified as passive dispersers on larger scales (Palmer 1988; Armonies 1994; Sun and Fleeger 1994; Fleeger et al. 1995). Similar trends for passive dispersal are seen in lotic freshwater ecosystems (Palmer and Gust 1985) and soft-bottom coastal ecosystems with regular tidal currents and strong hydrodynamic forces (Sedlacek and Thistle 2006). Nevertheless, in hydrodynamic calm environments (low flow or with biogenic structures reducing the flow),

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meiobenthic organisms were found suspended in the near-bottom waters revealing active emergence, especially on a diurnal cycle at the onset of dusk (Fleeger et al. 1984; Hicks 1986; Walters 1991; Teasdale et al. 2004). Morphological characteristics were put forward to endorse the emergence availability of phytal and epibenthic meiofauna (Bell et al. 1987; Thistle and Sedlacek 2004; Sedlacek and Thistle 2006). Lower taxonomic classification seems thus to be irrelevant in predicting the habitat utilization of copepods (Sedlacek and Thistle 2006). Noodt (1971) attempted a provisional classification of Copepoda based on the variety of evolved ecological forms (Remane 1952). He highlighted different trends in specialization of eco-morphological characteristics, distinguishing various types of copepod adapted to certain conditions of various habitats (e.g. sediment-living, phytal-living, pelagic). Subsequently, a preferred habitat could be deduced from eco-morphological characteristics. However, a classification of specific copepod colonization abilities is still missing, conversely to nematodes (Bongers 1990). Nowadays, phytal and epibenthic copepods are mainly classified as active dispersers and sedimentary copepods as passive dispersers (Hicks 1986; Kurdziel and Bell 1992).

Though, in case of colonization of new habitat or in the habitat connectivity context, the exact habitat source pool of the colonizers is often unknown. Several studies documented copepods' colonization ability within relatively stable habitats, such as coral fragments (Gheerardyn et al. 2009; Callens et al. 2012), hard substrates (Thomsen et al. 2011), coastal soft sediments (Thielemans and Heip 1984; Scheef and Marcus 2010) and deep-sea sediments (Thistle 1978; Guidi-Guilvard et al. 2009). However, few studies tackled the colonization of provisional habitats such as temporary ponds (Frisch and Green 2007), marine snow aggregates (Kiorboe 2000; Koski et al. 2005) or floating vegetal material (Faust and Gullledge 1996; Ólafsson et al. 2001). The present in situ experiment investigated the colonization of provisional dead seagrass detritus, hereafter referred to as macrophytodetritus. The majority of the macrophytodetritus accumulates on bare sand patches close to seagrass meadow and is decomposed within a few days to several months, depending on the chemical composition and biotic and abiotic fragmentation speed (Romero et al. 1992; Mateo and Romero 1997). These accumulations thereby support high values of secondary production in the receiving communities (Vetter 1995; Mateo and Romero 1997). The structurally complex macrophytodetritus accumulations seem to facilitate the development of meiofaunal communities in coastal marshes (Sanmarti and Menendez 2007), mangrove forests (Torres-Pratts and Schizas 2007) and seagrass beds (Hicks 1980; Coull and Wells 1983; Mascart et al. 2013). In terms of copepod community, connectivity between these accumulation and other adjacent

habitats (i.e. seagrass meadow, water column and sediment) is still unstudied, as well as the colonizing process and specific pattern of this colonization.

To try to tackle these three questions, a field experiment was deployed in order to understand the mode of copepod's colonization of Neptune grass *Posidonia oceanica* macrophytodetritus. The objectives here were threefold: (1) to assess the species composition, densities and diversity of the colonist's source pool (i.e. the sediment, the water column, the *P. oceanica* canopy or other macrophytodetritus patches); (2) to investigate the rate of active colonization of defaunated seagrass macrophytodetritus from adjacent habitats; and (3) to contribute to our knowledge species-specific colonization characteristics.

Materials and methods

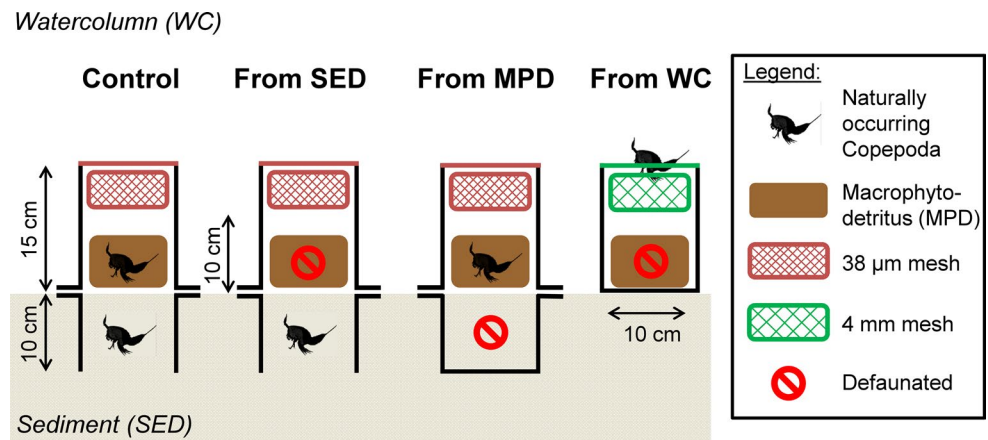
Experimental design and sampling site

The experimental site was located in the Gulf of Calvi, Corsica, northwest Mediterranean (42°35'N, 8°43'E) near the oceanographic station of STARESO (Station de Recherches Sous-marines et Océanographiques, University of Liège). The site consisted of a sand patch (about 100–200 m²) at a depth of 12 m, inside a *P. oceanica* seagrass meadows, partly covered with macrophytodetritus accumulation (MPD). Macrophytodetritus accumulation (10–20 cm thick) was mostly composed of dead *P. oceanica* leaves and living uprooted seagrass shoots, typical for the Bay area and the time of the year (Mascart et al. 2015).

The in situ experiment, comprised of cylindrical experimental PVC units (inner diameter of 10 cm), was set up by scuba divers on 26 October 2012 for 24 h of incubation and on 2 November 2012 for 96 h of incubation. During the experiment, the site was characterized by a constant salinity of 38, calm weather conditions and weak currents (4–5 cm s⁻¹). Temperatures varied between 18 and 21 °C and light intensities were highest (1200–4000 lux) between 11 a.m. and 14 p.m. (HOBO® Onset Computer Corporation).

The cylindrical experimental PVC units were divided into two parts: (1) an upper compartment containing on average 12 ± 6 gDW fresh macrophytodetritus (i.e. slightly degraded dead seagrass leaves) (MPD) and (2) a lower compartment containing sandy sediment (SED) (Fig. 1). The upper compartment had a height of 15 cm, as used in emergence traps (Walters and Bell 1986) in order to exclude water flow-driven effects and random contamination of mesopsammic copepods. At 10 cm height, a circumference window was made and together with the open top of the tube, both were covered with a 38-µm mesh, in order to exclude any contamination and predation by macrofauna

Fig. 1 Experimental design representing the four treatments, showing the dimensions, the two mesh sizes used, the defaunated habitats and the two compartments: *upper* (on top of the sediments) and *lower* (inside the sediments). *MPD* macrophytodetritus. *SED* sediments, *WC* water column



but to allow water and oxygen exchange. The lower compartment was inserted 10 cm deep into the sediment. This depth was chosen since vertical penetration of copepods happens in surficial layer (Danovaro and Fraschetti 2002; Kotwicki et al. 2005; Giere 2009), rarely exceeding 5–10 cm depth and therefore contamination from the surrounding sediments is excluded. The bottom of the tube remained open. Both compartments were placed on top of each other and fitted to stabilizing plates at the water–sediment interface providing support. The stabilizing plates offered guidance for the insertion of the splitter plate used to collect upper or lower compartment at the end of the incubation (Fig. 1).

The experiment was conducted twice. A first short-term incubation of 24 h (T_{24}) lasting a full diurnal cycle was carried out to rule out any pattern of vertical migration due to the change in light intensity (Walters and Bell 1986). A second mid-term incubation of 96 h (T_{96}) was performed to give a chance to the colonizing community to stabilize, however, not too long to avoid potential intra-specific competition and thus a colonization-competition trade-off (Chandler and Fleeger 1983; De Troch et al. 2005). Both experiments started at midday, but were not set up simultaneously to avoid disturbance during collection of an experiment. Each experiment was preceded by a reference sampling (T_0) in order to define the natural species composition and variability in the adjacent, potential source pool, namely the natural bare sandy sediments (T_0 SED), the natural macrophytodetritus (T_0 MPD), the adjacent *P. oceanica* canopy (T_0 POS) and the water column 1 m above the site (T_0 WC). These four habitats were sampled with, respectively, sediment meiocores (De Troch et al. 2001), detritus-cores (Mascart et al. 2015), plastic bags (Lepoint et al. 2006; Mascart et al. 2013) and 50- μ m-mesh hand towed horizontal plankton nets (Hamner and Carleton 1979), respectively.

The experimental design consisted of four treatments in quadruplicates (Fig. 1). The first treatment ‘Control’ was

set to test for possible effects of the deployed units on meiofauna MPD and sediment natural community. The upper and lower compartments were filled with natural macrophytodetritus and sediments, respectively. The second treatment ‘From SED’ tested the colonization of copepods from natural bare sediments towards defaunated macrophytodetritus positioned above it. The upper and lower compartments were filled with defaunated macrophytodetritus and natural sediments, respectively. The third treatment ‘From MPD’ tested the opposite colonization direction from natural macrophytodetritus towards defaunated sediments. The upper compartment contained natural macrophytodetritus, while the lower compartment contained defaunated sediments. The lower compartment was closed to suppress colonization by the sediment. The fourth treatment ‘From WC’ only consisted of an upper compartment filled with defaunated macrophytodetritus and was closed with a 4-mm mesh to allow larger planktonic copepods to enter the system, while excluding macrofaunal predators (e.g. juvenile fishes, amphipod crustaceans and shrimps). This last treatment tested for copepod colonization towards the macrophytodetritus from the surrounding water column.

Defaunation of macrophytodetritus and sediment was performed prior to the experiment. For this purpose, additional natural samples of macrophytodetritus were taken and were rinsed thoroughly with fresh water and an 8 % $MgCl_2$ -solution (Hulings and Gray 1971) on a 1-mm sieve in order to stun and remove attached organisms, while keeping the loss of epiphytes living on the dead leaves surface as minimal as possible. Additional sediment samples were collected and defaunated by, a gentle defaunation technique (in contrast to the destructive methods applied by Chandler and Fleeger 1983; Chertoprud et al. 2005) to prevent loss of attractiveness for the potential colonizing copepods. The sediments were bathed in freshwater for several minutes (to detach copepods), mixed by hand and decanted. A control subsample was taken and analysed under a stereomicroscope to confirm the successful defaunation. The

process was repeated on average five times until no copepods remained.

Sample collection and treatment

At the end of the incubation, prior to sampling, both compartments were isolated by inserting a splitter plate in between the stabilizing plates. Each compartment together with its content was subsequently transferred to a closed plastic zip bag to avoid contamination and any loss of material. The upper compartments containing macrophytodetritus were afterwards rinsed with an 8 % MgCl_2 -solution and fresh water in order to separate meiofauna from the macrophytodetritus. The rinsed samples were sieved over a 1-mm and 38- μm -mesh sieve to exclude macrophytodetritus and retain copepods, respectively. The copepods were preserved in a 4 % formaldehyde seawater solution. The separated macrophytodetritus was dried at 60 °C for 4 days (Mascart et al. 2015) to measure dry weight for further standardization of copepod densities. The lower compartments comprising sediments were subsampled with meiocores and afterwards preserved in a 4 % formaldehyde seawater solution. The copepods present in the sediment were subsequently extracted by centrifugation with Ludox HS40 (specific density of 1.18 g dm^{-3}). The copepods were stained with Rose Bengal, counted, picked out and mounted on slides for microscopic identification to species level based on identification keys of Lang (1948) and Boxshall and Hasley (2004). Densities of copepods in the macrophytodetritus were standardized towards the dry weight of macrophytodetritus (indiv. g^{-1} DW) and towards the sediment surface in the case of benthic samples (indiv. 10 cm^{-2}).

Data analysis

To analyse the structure of the community, five species diversity metrics were used reflecting the different elements of biodiversity (see Magurran (2004) and Magurran and McGill (2011) for an overview): S = number of species observed (species number); d = Margalef's corrected number of species for N number of individuals (species richness); H' = the Shannon's diversity index based on natural logarithm (species diversity); E_H = Heip's evenness index sensitive to rare species (species evenness); and N_1 = number of species that would have been found in the sample when all species would be equally common (dominance metrics).

In order to identify the most typifying copepod species primarily providing the discrimination between and within factors, a SIMPER (similarity percentages) routine was used after an ANOSIM (analysis of similarity) difference test. To visualize the community structure a principal coordinate analysis (PCO) based on the zero-adjusted

Bray–Curtis similarity resemblance matrix of the log-transformed relative multivariate data of copepod species abundances was performed.

The analysis of variance in univariate or multivariate data were examined using a PERMANOVA routine and post hoc pairwise comparisons with fixed factors Treatment (Control, From SED, From MPD and From WC), Habitat (MPD, SED) or Time (T_0 , T_{24} and T_{96}). PERMANOVA allows us to perform an ANOVA with P values obtained by permutation (Anderson et al. 2008), thus avoiding the assumption of normality. Prior to run two-way PERMANOVA's, assumption of homogeneity of dispersion was tested with a PERMDISP and distances amongst centroids calculated on the interaction level (Quinn and Keough 2002). Euclidean distance and Bray–Curtis-based resemblance matrices were used, respectively, for univariate and multivariate measures. Pairwise tests type III and Monte-Carlo P values were used since sometimes the total number of unique permutations did not exceed a hundred, whereas 4999 unique permutation is favourable (Anderson et al. 2008).

The calculated resemblance matrixes were based on a zero-adjusted log-transformed data with the addition of one dummy variable, for the reason that defaunated compartments contained no species at the start of the incubation. Prior to the analysis, both experiments' reference samples (T_0 samples of T_{24} and T_{96}) were checked for significant differences in species composition for the factors Habitat and Time. No differences were found between both time references sample for each habitat ($P = 0.471$). Within each duration, all four reference habitats were significantly different in species composition ($P < 0.001$). Therefore, both incubations reference samples were pooled into one T_0 reference per habitat in order to have a higher replication and thus higher statistical results. Due to different standardization methods within each unit compartment (habitat MPD per gram dry weight vs. SED per surface area), differences in total absolute copepod densities were examined per compartment using factors Time and Treatment. Variance in species compositions between the start (T_0) and the end of incubation (T_{24} or T_{96}) were investigated using the fixed factors Treatment and Habitat.

All the above-mentioned analyses were performed with the Primer 6.1.11 software (Clarke and Gorley 2006) with PERMANOVA add-on software (Anderson et al. 2008). A significance level of $P < 0.05$ was used in all tests. Graphs were constructed in GraphPad 5.03 for Windows (GraphPad Software, San Diego California USA).

Results

Within 24 and 96 h of incubation, all defaunated habitats were colonized by copepods; therefore, no repulsive effect

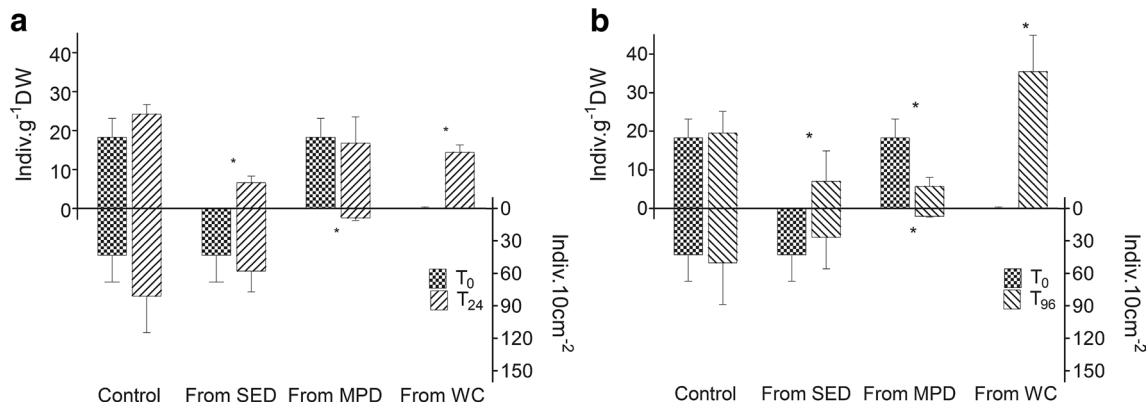


Fig. 2 Mean Copepoda densities per treatment for the 24-h incubation experiment (a) and the 96-h incubation experiment (b). Upper part represents the upper compartment with macrophytodebris (MPD) standardized per indiv. g⁻¹DW (dry weight) on the left y axis. Lower part represents the lower compartment with the sediments

(SED) standardized per indiv. 10 cm⁻² on the right y axis. Error bars represent the standard deviation. $N = 4$ per treatment and WC water column. Asterisk shows a significant difference ($P < 0.05$) in total densities between start and end incubation

of the experimental set-up was found (all pairwise $P > 0.1$) (Fig. 2). Over all samples, a total of 58 different species were identified belonging to three Copepoda orders. The majority of the species (50) belonged to the order harpacticoida, representing 83.8 ± 2.1 % (average \pm SD, henceforth used as notation) of the encountered species. Five species belonged to the order Cyclopoida and three species to the calanoida order (Table 1). Noodt (1971) attempted a provisional classification of Copepoda based on the variety of morphological forms which are adapted to special conditions in various habitats. All eco-morphological types except parasitic types were present in this study: (M) Mesopsammic types, primarily sediment living; (P) Phytal types, clinging to phytal structures; (E) Epibenthic types, benthic-swimmers; (W) Water column types, pelagic free-swimmers. A complete classification per species was made here (Table 1) in accordance with former studies and morpho-ecological traits (Lang 1948; Noodt 1971; Bell et al. 1987; Higgins and Thiel 1988; Bodin and Leguellec 1992; Thistle and Sedlacek 2004). Following the obtained results, a classification in terms of active colonization was added per species: (I) infaunal colonizers, interstitial dispersal pathway; (S) suspension colonizers, water-bound dispersal pathway; and (-) non-active colonizer, persists in its initial habitat (Table 1). In order to keep a comprehensive overview, the results are presented per factor Time and succinctly by factor Treatment.

Reference samples T_0

All T_0 reference samples in sediment, water column, *P. oceanica* canopy and macrophytodebris were significantly different from each other in terms of species composition (ANOSIM, global $R = 0.994$, $P < 0.001$). The

top four species typifying a reference habitat were always distinct with the exception of *Sarsamphiascus tenuiremis* (Miraciidae family) which is omnipresent in all reference habitats, except for the water column (Table 2). Since a high similarity between replicates was found in each habitat, the dissimilarity between pairs of habitats is as expected high (>81 %), except between the macrophytodebris and *P. oceanica* canopy where the dissimilarity is reduced to 46.2 %. The average evenness E_H was 0.27 ± 0.04 for all samples. The highest number of species and species richness was accounted for macrophytodebris with an S of 25.3 ± 1.0 and $d = 5.1 \pm 0.2$, which is quite similar to the number of species in the *P. oceanica* canopy with $S = 23.5 \pm 1.7$ and $d = 4.9 \pm 0.3$. The sediments had a lower number of species $S = 14.3 \pm 0.6$ and richness $d = 3.6 \pm 0.3$. The lowest species number and richness was found in the water column with $S = 8.0 \pm 1.4$ and $d = 1.6 \pm 0.3$. The evenness was similar in all four samples. The total copepod density was 42.7 ± 24.5 indiv. 10 cm⁻² for the bare sediments (T_0 SED), 18.3 ± 4.9 indiv. g⁻¹ DW for the natural macrophytodebris (T_0 MPD), 8.1 ± 2.4 indiv. g⁻¹ DW for the *P. oceanica* canopy (T_0 POS) and 120.2 ± 4.6 indiv. m⁻³ for the water column (T_0 WC).

All the species found in the seagrass meadow were found in macrophytodebris, with the exception of *Sacodiscus littoralis* (family Tisbidae). It was the only non-colonizing species exclusively found in the *P. oceanica* canopy. The species *Ambunguipes rufocincta*, *Probosciphontodes stellate*, *Syngastes cornalinus*, *Tegastes calcaratus*, *Rhynchothalestris helgolandica* and *Xouthous laticaudatus* were only present in the macrophytodebris habitat, while the following species were exclusively present in the sediment: *Arenosetella tenuissima*, *Canuella*

Table 1 Cumulative presence list of Copepoda species sorted per order and per family based on four replicates

Active Colonizer	Ecologic. Type	Species	Time:		T ₀								T ₂₄								T ₉₆							
			Treatment:	Habitat:				Control		From SED		From MPD		From WC		Control		From SED		From MPD		From WC						
				SED	MPD	POS	WC	MPD	SED	MPD	SED	MPD	MPD	SED	MPD	SED	MPD	MPD	SED	MPD	SED	MPD	SED	MPD				
Harpacticoida																												
Ameiridae																												
I/S	M	<i>Ameira longipes</i> (Boeck, 1865)			X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X			
S	M	<i>Ameiropsis nobilis</i> (Sars G.O., 1911)	X	X						X	X											X	X					
Ancorabolidae																												
I/S	P	<i>Laophontodes bicornis</i> (Scott A., 1896)			X	X				X					X						X	X	X		X			
-	P	<i>Probosciphontodes stellata</i> (Fiers, 1988)			X					X											X		X					
Canuelidae																												
-	M	<i>Canuella furcigera</i> (Sars G.O., 1903)	X							X	X									X	X							
Cletodidae																												
I/S	M	<i>Cletodes timicola</i> (Brady, 1872)	X	X						X		X	X															
Dactylopusiidae																												
S	P	<i>Dactylopusia tisboides</i> (Claus, 1863)			X	X				X		X			X							X		X				
-	P	<i>Diarthrodes minutus</i> (Claus, 1863)			X	X				X		X			X							X						
-	P	<i>Paradactylopusia brevicornis</i> (Claus, 1866)			X	X				X		X			X							X						
Ectinosomatidae																												
I/S	M	<i>Ectinosoma dentatum</i> (Steuer, 1940)	X	X						X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X			
-	M	<i>Arenosetella tenuissima</i> (Klie, 1929)			X					X		X										X						
-	M	<i>Pseudobradia hirsuta</i> (Scott T. & A., 1896)			X					X		X									X							
S	W	<i>Microsetella norvegica</i> (Boeck, 1865)					X							X											X			
Hamondiidae																												
-	P	<i>Ambungapes rufocincta</i> (Norman in Brady, 1880)			X					X			X							X		X						
Harpacticidae																												
S	P	<i>Harpacticus littoralis</i> (Sars G.O., 1910)			X	X				X			X													X		
Laophontidae																												
I	M	<i>Asellopsis duboscqui</i> (Monard, 1926)	X	X							X	X	X	X							X					X		
S	P	<i>Esola longicauda</i> (Edwards, 1891)													X											X		
I	M	<i>Laophonte cornuta</i> (Philippi, 1840)	X							X	X	X			X						X		X	X	X	X		
I/S	M	<i>Laophonte elongata elongata</i> (Boeck, 1873)	X	X						X	X	X			X						X	X	X	X	X	X		
I	M	<i>Laophontina posidoniae</i> (Fiers, 1986)	X							X	X	X								X	X	X	X	X	X	X		
I/S	P	<i>Paralaophonte brevivirostris</i> (Claus, 1863)	X	X	X					X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
Leptastacidae																												
I	M	<i>Leptastacus laticaudatus</i> (Nicholls, 1935)	X							X		X	X							X	X	X	X	X	X	X		
Leptopontiidae																												
I	M	<i>Leptopontia curvicauda</i> (Scott, 1902)	X	X						X	X	X			X					X	X	X	X	X	X	X		
Longipediidae																												
S	P	<i>Longipedia minor</i> (Scott T. & A., 1895)	X	X						X			X		X					X	X					X		
Metidae																												
I	P	<i>Metis ignea</i> (Philippi, 1843)			X	X				X	X	X		X	X					X								
Miraciidae																												
I/S	M	<i>Amphiascoides debilis</i> (Giesbrecht, 1881)	X	X	X					X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
I/S	P	<i>Diosaccus tenuicornis</i> (Claus, 1863)	X	X	X					X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
I/S	M	<i>Amphiascus minutus</i> (Claus, 1863)	X	X	X					X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
I/S	M	<i>Sarsamphiascus tenuiremis</i> (Brady, 1880)	X	X	X					X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
I/S	M	<i>Delavalia normani</i> (Scott T., 1905)	X	X						X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
Rhizotrichidae																												
I	M	<i>Rhizotrix curvatum</i> (Brady, 1880)	X	X						X		X		X						X	X		X					
Paramesochridae																												
I	M	<i>Wellsoipsyllus (Scott) robertsoni</i> (Scott T. & A., 1895)	X	X						X		X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
I	M	<i>Wellsoipsyllus (Inter.) intermedius</i> (Scott T. & A., 1895)	X							X		X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
Peltidae																												
S	P	<i>Alteutha depressa</i> (Claus, 1863)			X	X				X			X	X	X					X		X		X		X		
Porcellidiidae																												
S	P	<i>Porcellidium ovatum</i> (Haller, 1879)			X	X				X			X	X	X					X		X		X		X		
S	P	<i>Porcellidium fimbriatum</i> (Claus, 1863)			X	X				X			X	X	X					X		X		X		X		
Pseudotachidiidae																												
I/S	P	<i>Dactylopodella flava</i> (Claus, 1866)	X	X						X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
S	P	<i>Xouthos laticaudatus</i> (Thompson I.C. & Scott A., 1903)	X							X					X					X	X	X	X	X	X	X		
Tegastidae																												
S	P	<i>Parategastes sphaericus</i> (Claus, 1863)			X							X	X	X							X		X		X	X		
S	P	<i>Syngastes cornalinus</i> (Monard, 1924)			X					X		X	X	X														
-	P	<i>Tegastes calcaratus</i> (Sars G.O., 1910)			X							X	X	X														
S	P	<i>Tegastes satyrus</i> (Claus, 1860)			X	X				X		X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
Tetragonicepsidae																												
I/S	P	<i>Tetragoniceps laevis</i> (Wiley, 1930)				X				X	X			X						X						X		
I/S	M	<i>Phyllopodopsyllus bradyi</i> (Scott T., 1892)	X	X							X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
I	M	<i>Tetragoniceps scotti</i> (Sars G.O., 1911)	X							X			X	X	X	X	X	X	X	X	X	X	X	X	X	X		
Thalestridae																												
S	P	<i>Rhynchothalestris helgolandica</i> (Claus, 1863)			X							X								X		X		X		X		
Tisbidae																												
S	E	<i>Tisbe elegantula</i> (Sars G.O., 1905)			X	X				X		X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
S	E	<i>Tisbe ensifer</i> (Fischer, 1860)			X	X				X		X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
S	E	<i>Tisbe furcata</i> (Baird, 1837)			X	X				X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
-	P	<i>Sacodiscus littoralis</i> (Sars G.O., 1904)			X																							
Calanoida																												
Clausocalanidae																												
S	W	<i>Clausocalanus arcuicornis</i> (Dana, 1849)			X	X	X			X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
Lucicutiidae																												
S	W	<i>Lucicutia magna</i> (Wolfenden, 1903)			X	X				X	X				X					X		X		X		X		
Paracalanidae																												
-	W	<i>Paracalanus parvus parvus</i> (Claus, 1863)					X																					
Cyclopoida																												
Cyclopinidae																												
S	W	species 1			X	X	X	X		X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X		
S	W	species 2													X						X					X		
-	W	species 3																										
Oithonidae																												
S	W	<i>Oithona nana</i> (Giesbrecht, 1893)			X	X									X					X	X	X	X	X	X	X		
S	W	<i>Oithona similis</i> (Claus, 1866)			X	X									X					X	X	X	X	X	X	X		

X = presence. Blank cells = absence. The active colonization pathway in the outer left column are *I* infaunal colonizers, interstitial dispersal pathway, *S* suspension colonizers, water-bound dispersal pathway, – non-active colonizer, persists in its initial habitat. The ecological types presented in the second column are *M* mesopsammic types, primarily sediment living, *P* phytal types, clinging to phytal structures, *E* epibenthic types, benthic-swimmers, *W* water column types, pelagic free-swimmers

Table 2 Multivariate PERMANOVA and SIMPER results representing the typifying species of the reference samples

Species typifying reference habitats		
Species	%	cum.%
<i>T₀</i> SED (54.4 % similarity)		
<i>Sarsamphiascus tenuiremis</i>	43.0	43.0
<i>Leptastacus laticaudatus</i>	11.6	54.6
<i>Wellsopsyllus (Scottopsyllus) robertsoni</i>	8.1	62.7
<i>Amphiascoides debilis</i>	8.1	70.8
<i>Ectinosoma dentatum</i>	8.0	78.7
<i>T₀</i> MPD (74.4 % similarity)		
<i>Dactylopodella flava</i>	22.9	22.9
<i>Ameira longipes</i>	12.3	35.2
<i>Dactylopusia tisbooides</i>	9.0	44.2
<i>Sarsamphiascus tenuiremis</i>	8.0	52.2
<i>Amphiascus minutus</i>	6.7	58.9
<i>T₀</i> POS (73.1 % similarity)		
<i>Harpacticus littoralis</i>	14.3	14.3
<i>Diosaccus tenuicornis</i>	12.2	26.5
<i>Sarsamphiascus tenuiremis</i>	9.6	36.1
<i>Amphiascus minutus</i>	7.6	43.6
<i>Porcellidium ovatum</i>	6.4	50.0
<i>T₀</i> WC (90.0 % similarity)		
<i>Paracalanus parvus parvus</i>	36.3	36.3
<i>Clausocalanus arcuicornis</i>	27.2	63.5
<i>Oithona similis</i>	13.3	76.8
Cyclopinae sp. 1	9.7	86.5
<i>Microsetella norvegica</i>	7.8	94.3

T₀ first five contributing species are shown, *SED* sediments, *MPD* macrophytodebris, *POS* *Posidonia oceanica* canopy, *WC* water column

furciger, *Pseudobradya hirsuta* and *Wellsopsyllus (Intermediopsyllus) intermedius*. All the water column habitat species were present in the newly colonized macrophytodebris, except for Cyclopinae sp. 3 and *Paracalanus parvus parvus* (Table 1).

The 24-h experiment

In the first experiment after an incubation of 24 h, all defaunated habitats were colonized independently of the treatment (Fig. 2). The multivariate two-way PERMANOVA was significant for both factors (Time and Habitat) and interaction factor (Table 3). PERMDISP's for the interaction factor turned out to be significantly different, indicating that the variation within all factors and interactions was due to the dispersion and location effect, mainly because of the large number of zeros present. Pairwise comparisons revealed no significance difference between

the *T₂₄* 'Control' habitats and the *T₀* reference samples (Table 3). In comparison with other treatments, the 'Control' treatment showed only significant differences for the treatment 'From SED' SED habitat (Pairwise: $t = 2.39$, $P_{(MC)} = 0.013$) and for 'From WC' MPD habitat (Pairwise: $t = 2.94$, $P_{(MC)} = 0.003$) (Table 3).

In the 'From SED' treatment, the defaunated MPD compartment was colonized by sediment copepods reaching a density of 6.6 ± 1.7 indiv. $g^{-1}DW$ (Fig. 2a). A dissimilarity of 65.0 % (SIMPER) was found between the 'From SED' and 'Control' treatment MPD habitat, although species composition was not significantly different (Table 3). In the 'From SED' treatment, the PCO clearly separates SED from MPD habitats, each were, however, similar at 65 % (Fig. 3a). Diversity in this treatment is lower than the 'Control' sample (Table 5) due to the high level of dominance (>56 %) of *S. tenuiremis* (Table 4). The source pool community for this colonization, the SED habitat of the 'From SED' treatment had a lower diversity, evenness (Table 5) and a distinct species assemblage (Fig. 3a). Even though *Sarsamphiascus tenuiremis* remained the dominant species (Table 4).

The reverse treatment ('From MPD') tested a possible active migration from the natural macrophytodebris to the defaunated sediments. The defaunated SED habitat was colonized with a density of 8.6 ± 2.4 indiv. 10 cm^{-2} and reached a diversity comparable to the 'Control' SED habitat. The PCO displayed an assemblage almost identical to the 'Control'. No significant difference in composition was found and four species of the top five most contributing species to the similarity were identical. *S. tenuiremis* was the dominant species (Table 4). The species assemblage had a low dissimilarity of 37.2 % and *Dactylopodella flava* remained the dominant species (Table 4). The emigration to the sediments from the MPD compartment showed no significant impact on the latter's initial density (Fig. 2a) and diversity (Table 5), displaying the high variability present within the source pool.

Colonization from the surrounding water column was unambiguous since the density raised from zero to 14.4 ± 1.9 indiv. $g^{-1}DW$ (Fig. 2a). The species composition was dominated by a Cyclopid species from the Cyclopinae family and the calanoida, *Clausocalanus arcuicornis* (Table 4). The diversity metrics were in the same order of magnitude as in the 'Control' MPD habitat; however, the diversity and evenness were very low (Table 5). In comparison with treatment 'From SED', where the sediments exclusively serve as source pool a higher influx of individuals and species occurred. The species composition present after MPD colonization through the water column was significantly different from all four adjacent reference habitat samples (Table 3).

Table 3 Two-way multivariate PERMANOVA of species composition and post hoc pairwise tests

Factors and interaction	T_{24}		T_{96}	
Habitat (Ha)	$F'_{(1,2)} = 12.9; P = 0.001^{**}$		$F'_{(1,2)} = 18.4; P = 0.001^{**}$	
Treatment (Tr)	$F'_{(3,2)} = 4.1; P = 0.001^{**}$		$F'_{(3,2)} = 3.2; P = 0.001^{**}$	
Ha × Tr	$F'_{(3,2)} = 2.5; P = 0.011^*$		$F'_{(3,2)} = 2.8; P = 0.003^{**}$	
Pairwise comparisons	T_{24}		T_{96}	
	SED	MPD	SED	MPD
T_0 versus control	$t = 1.39; P_{(MC)} = 0.137$	$t = 1.30; P_{(MC)} = 0.166$	$t = 1.97; P_{(MC)} = 0.035^*$	$t = 2.09; P_{(MC)} = 0.012^{**}$
From SED versus control	$t = 2.39; P_{(MC)} = 0.013^*$	$t = 1.69; P_{(MC)} = 0.065$	$t = 1.86; P_{(MC)} = 0.039^*$	$t = 1.61; P_{(MC)} = 0.063$
From MPD versus control	$t = 0.98; P_{(MC)} = 0.431$	$t = 1.11; P_{(MC)} = 0.293$	$t = 2.98; P_{(MC)} = 0.004^{**}$	$t = 1.30; P_{(MC)} = 0.169$
From WC versus control	–	$t = 2.94; P_{(MC)} = 0.003^{**}$	–	$t = 2.01; P_{(MC)} = 0.011^{**}$
T_0 SED versus from WC	–	$t = 4.06; P = 0.038^*$	–	$t = 5.13; P = 0.032^*$
T_0 MPD versus from WC	–	$t = 3.48; P = 0.031^*$	–	$t = 5.60; P = 0.035^*$
T_0 POS versus from WC	–	$t = 3.55; P = 0.029^*$	–	$t = 5.36; P = 0.032^*$
T_0 WC versus from WC	–	$t = 6.60; P = 0.020^*$	–	$t = 8.77; P = 0.033^*$

F' pseudo- F value, $P_{(MC)}$ Monte-Carlo P value, * $0.05 > P > 0.01$ = significant; ** $0.01 > P > 0.001$ = highly significant. On the left side, the 24-h incubation experiment and on the right the 96-h incubation experiment

SED sediments, MPD macrophytodebris, POS *Posidonia oceanica* canopy, WC water column

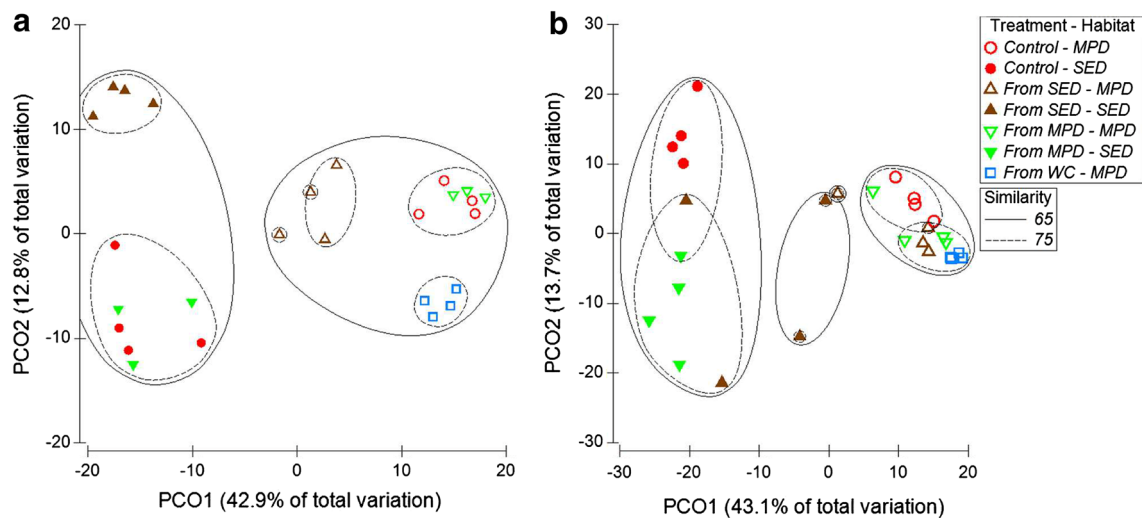


Fig. 3 Principal coordinate analysis (PCO) based on a Bray–Curtis similarity resemblance matrix on log-transformed data of species abundance of (a) 24-h incubation experiment and (b) 96-h incubation experiment. Filled symbols represent the sediments (SED) habitat and the un-filled symbols the macrophytodebris (MPD) habitat. Differ-

ent treatments are represented by symbols: circles control; triangles from SED; reverse triangles from MPD; squares from WC (water column). Full line represents 65 % similarity and dashed line represents 75 % similarity

The 96-h experiment

The copepod density in both ‘Control’ compartments of T_{96} was not significantly different from the respective T_0 reference samples habitat. A density of 19.5 ± 5.7 indiv. $g^{-1}DW$ and 50.3 ± 38.4 indiv. 10 cm^{-2} was, respectively, found for the ‘Control’ MPD and SED compartment (Fig. 2b). The diversity (S) was significantly lower (Table 5) in both

‘Control’ compartments after 96 h of incubation (Pairwise: $t = 2.92, P_{(MC)} = 0.025$ for MPD and $t = 2.68, P_{(MC)} = 0.031$ for SED). In terms of species composition, pairwise comparisons revealed significant differences between the T_{96} ‘Control’ habitats and the T_0 reference samples (Table 3). The dissimilarity between the T_0 and the T_{96} ‘Control’ reached 53.5 % for the MPD and 52.9 % for the SED habitats (SIMPER).

Table 4 Similarity percentages (SIMPER) for both experiments with factors *treatment* and *habitat* for copepod species contributions

24-h incubation experiment		From SED		From MPD		From WC	
Upper compartment (habitat MPD)							
Species (65.7 % similarity)	%	cum.%	Species (43.7 % similarity)	%	cum.%	Species (72.8 % similarity)	%
<i>Dactylopodella flava</i>	23.7	23.7	<i>Sarsamphiascus tenuiremis</i>	56.6	56.6	<i>Dactylopodella flava</i>	14.7
<i>Sarsamphiascus tenuiremis</i>	17.2	41.0	<i>Diosaccus tenuicornis</i>	9.6	66.2	<i>Dactylopusia tisbooides</i>	13.0
<i>Ameira longipes</i>	13.4	54.3	<i>Ectinosoma dentatum</i>	9.6	75.8	<i>Sarsamphiascus tenuiremis</i>	12.9
<i>Amphiascus minutus</i>	11.1	65.4	<i>Ameira longipes</i>	9.6	85.3	<i>Tegastes satyrus</i>	7.8
<i>Diosaccus tenuicornis</i>	6.6	72.0	<i>Amphiascus minutus</i>	4.8	90.2	<i>Paradactylopodia brevicornis</i>	7.2
Lower compartment (habitat SED)							
Species (59.8 % similarity)	%	cum.%	Species (61.1 % similarity)	%	cum.%	Species (61.7 % similarity)	%
<i>Sarsamphiascus tenuiremis</i>	38.5	38.5	<i>Sarsamphiascus tenuiremis</i>	33.1	33.1	<i>Sarsamphiascus tenuiremis</i>	23.9
<i>Rhizothrix curvatum</i>	14.1	52.6	<i>Leptopontia curvicauda</i>	24.5	57.6	<i>Wellsopsyllus (Inter.) intermedius</i>	16.4
<i>Leptopontia curvicauda</i>	10.0	62.6	<i>Canuella furcigera</i>	18.0	75.6	<i>Rhizothrix curvatum</i>	16.4
<i>Amphiascoides debilis</i>	10.0	72.6	<i>Delavatia normani</i>	16.0	91.6	<i>Leptopontia curvicauda</i>	11.0
<i>Wellsopsyllus (Inter.) intermedius</i>	9.1	81.6	<i>Phyllopoposyllus bradyi</i>	2.0	93.6	<i>Ectinosoma dentatum</i>	9.3
96-h incubation experiment							
Upper compartment (habitat MPD)							
Species (50.4 % similarity)	%	cum.%	Species (34.4 % similarity)	%	cum.%	Species (56.1 % similarity)	%
<i>Ameira longipes</i>	17.0	17.0	<i>Ectinosoma dentatum</i>	47.9	47.9	<i>Ectinosoma dentatum</i>	37.6
<i>Ectinosoma dentatum</i>	13.2	30.2	<i>Amphiascus minutus</i>	17.2	65.2	<i>Oithona nana</i>	15.8
<i>Tisbe furcata</i>	13.2	43.4	<i>Ameira longipes</i>	13.5	78.6	<i>Clausocalanus arcuicornis</i>	15.7
<i>Dactylopodella flava</i>	10.3	53.7	<i>Tisbe furcata</i>	11.6	90.2	<i>Dactylopodella flava</i>	6.1
<i>Diosaccus tenuicornis</i>	9.0	62.7	<i>Sarsamphiascus tenuiremis</i>	3.8	94.1	<i>Tegastes satyrus</i>	6.1
Lower compartment (habitat SED)							
Species (66.9 % similarity)	%	cum.%	Species (24.0 % similarity)	%	cum.%	Species (59.8 % similarity)	%
<i>Leptopontia curvicauda</i>	28.7	28.7	<i>Ectinosoma dentatum</i>	30.0	30.0	<i>Leptastacus laticaudatus</i>	36.1
<i>Sarsamphiascus tenuiremis</i>	28.6	57.3	<i>Leptastacus laticaudatus</i>	25.4	55.4	<i>Rhizothrix curvatum</i>	30.8
<i>Amphiascoides debilis</i>	11.7	69.0	<i>Canuella furcigera</i>	17.1	72.5	<i>Wellsopsyllus (Scott.) robertsoni</i>	20.3
<i>Rhizothrix curvatum</i>	9.6	78.7	<i>Leptopontia curvicauda</i>	11.8	84.3	<i>Leptopontia curvicauda</i>	6.2
<i>Tetragoniceps scotti</i>	6.7	85.3	<i>Laophontina postidoniae</i>	6.2	90.5	<i>Sarsamphiascus tenuiremis</i>	2.6

First five contributing species are shown in percentage (%) and cumulative percentage (cum.%)

Table 5 Diversity metrics (average \pm SD) based on $N = 4$ per treatment per compartment for both incubation experiments

	24-h incubation experiment				
	T_0	T_{24}	T_{24}	T_{24}	T_{24}
	Reference	Control	From SED	From MPD	From WC
Upper compartment (habitat MPD)					
S	25.25 \pm 0.96	23.75 \pm 0.96	8 \pm 0.82	23.67 \pm 2.08	21 \pm 1.83
d	5.07 \pm 0.23	4.32 \pm 0.18	2.43 \pm 0.09	4.37 \pm 0.33	4.39 \pm 0.42
H'	2.82 \pm 0.07	2.87 \pm 0.06	2.37 \pm 0.26	2.9 \pm 0.06	1.64 \pm 0.1
E_H	0.25 \pm 0.01	0.29 \pm 0.02	0.58 \pm 0.15	0.3 \pm 0.03	0.1 \pm 0.01
N_1	16.78 \pm 1.13	17.74 \pm 1.08	10.94 \pm 3.06	18.27 \pm 1.15	5.18 \pm 0.49
Lower compartment (habitat SED)					
S	14.33 \pm 0.58	12.64 \pm 5.1	10.25 \pm 0.5	14 \pm 1.73	
d	3.6 \pm 0.29	3.03 \pm 0.92	2.31 \pm 0.19	3.12 \pm 0.35	
H'	2.24 \pm 0.13	2.14 \pm 0.46	1.9 \pm 0.06	2.76 \pm 0.07	
E_H	0.26 \pm 0.04	0.32 \pm 0.11	0.27 \pm 0.03	0.45 \pm 0.05	
N_1	9.49 \pm 1.21	9.19 \pm 4.09	6.73 \pm 0.39	15.79 \pm 1.1	
	96-h incubation experiment				
	T_0	T_{96}	T_{96}	T_{96}	T_{96}
	Reference	Control	From SED	From MPD	From WC
Upper compartment (habitat MPD)					
S	25.25 \pm 0.96	17 \pm 4.4	8 \pm 2.83	17 \pm 3.92	19 \pm 2.71
d	5.07 \pm 0.23	4.38 \pm 0.65	2.53 \pm 0.81	3.78 \pm 0.55	3.59 \pm 0.45
H'	2.82 \pm 0.07	2.62 \pm 0.24	1.84 \pm 0.34	2.37 \pm 0.31	2.16 \pm 0.21
E_H	0.25 \pm 0.01	0.32 \pm 0.02	0.37 \pm 0.09	0.26 \pm 0.05	0.18 \pm 0.02
N_1	16.78 \pm 1.13	13.99 \pm 3.34	6.57 \pm 1.99	11.12 \pm 3.85	8.82 \pm 1.83
Lower compartment (habitat SED)					
S	14.33 \pm 0.58	10.75 \pm 0.96	6 \pm 2	9.25 \pm 2.22	
d	3.6 \pm 0.29	2.74 \pm 0.25	1.92 \pm 0.25	2.08 \pm 0.6	
H'	2.24 \pm 0.13	2.06 \pm 0.13	1.6 \pm 0.16	1.65 \pm 0.23	
E_H	0.26 \pm 0.04	0.3 \pm 0.06	0.39 \pm 0.08	0.24 \pm 0.04	
N_1	9.49 \pm 1.21	7.91 \pm 0.98	4.87 \pm 0.69	5.33 \pm 1.22	

S number of species observed, d Margalev's species richness, H' Shannon species diversity, E_H Heip's evenness index and N_1 abundance of the most dominant species. *MPD* macrophytodebris, *SED* sediments and *WC* water column

Colonization of defaunated macrophytodebris from the sediments (treatment 'From SED') took place. After 96 h, 7.2 ± 7.0 indiv. g^{-1} DW were found; however, a large variability amongst the replicates was present (Fig. 2b). The species composition present in the newly colonized MPD habitat showed no significant difference (Table 3) compared to the 'Control'. Nonetheless, a dissimilarity of 68.6 % (SIMPER) was present. The newly colonized habitat was dominated by *Ectinosoma dentatum* (Ectinosomatidae family) (Table 4). In terms of species number, diversity and richness, lower values were noted compared to the 'Control'. However, a larger evenness was found (Table 5). In the natural sediment, the colonisers' source compartment, two species dominated after 96 h: the *E. dentatum* and *Leptastacus laticaudatus* (Table 4). Again a low species number, diversity and richness were combined with a high evenness (Table 5).

Colonization of the defaunated sediments by copepods from the natural macrophytodebris was effective within

96 h. The copepod densities of the colonized habitat in the treatment 'From MPD' increase to 7.4 ± 0.8 indiv. 10 cm^{-2} with species diversity metrics, showing no significant differences with the 'Control' (Fig. 2b). However, the species composition was significantly different (Table 3). Two species were most abundant in the SED compartment: *L. laticaudatus* and *Rhizotrix curvatum* (Table 4). The colonisers' source MPD compartment showed, in spite of a decrease in density towards 5.8 ± 2.3 indiv. g^{-1} DW, a similar diversity and species composition as the 'Control' MPD habitat. Worth noticing are the two non-harpacticoid species *Oithona nana* and *Clausocalanus arcuicornis* present in the top five contributors of the similarity (Table 4).

The colonization of defaunated macrophytodebris from the water column, treatment 'From WC', displayed a strong increase in copepods density from zero to 35.4 ± 9.4 indiv. g^{-1} DW, which is more than double of the densities after 24 h (Fig. 2b). After 96-h incubation,

similar species numbers and richness were found, however, with a higher evenness and diversity compared to the 24-h incubation. The species composition changed and another Cyclopoid copepod, *O. nana*, became dominant closely followed by *E. dentatum* (Table 4).

Discussion

Macrophytodebris colonization

Our results showed colonization of all defaunated habitats by species originating from adjacent habitats. These adjacent habitats were composed of very specific communities and species compositions defining each adjacent habitat. The densities and diversities encountered during the experiment were congruent with previous studies in the area (Dauby 1980; Heip et al. 1983; Mascart et al. 2013, 2015).

Colonization occurred from adjacent habitats within the first 24 h, which is corresponding to recovery times found after a physical disturbance (Sun and Fleeger 1994; Fleeger et al. 1995). Meiobenthic copepods exert colonization and active habitat selection via two pathways: infaunal dispersion through the interstitial spaces and dispersion through suspension in the water column. The majority of adjacent habitat-specific typifying species were present in the macrophytodebris, therefore showing traits of dispersion and making the macrophytodebris a diverse copepod hub. Moreover, some species seemed to be exclusively associated with macrophytodebris. It could thus be hypothesized that species have more than one generation in these provisional accumulations, highlighting the ecological role of macrophytodebris to the overall seagrass system.

When defaunated macrophytodebris covered natural bare sediments, an almost immediate interstitial colonization through the boundary layer from the sediment community occurred. Nonetheless, the formed assemblage did not fully resemble the possible macrophytodebris assemblage as found in the control. Similar patterns were found in the colonization from the water column towards defaunated macrophytodebris. On the contrary, during colonization from the natural macrophytodebris towards defaunated sediments, the formed assemblage in the sediment resembled the control as shown in the PCO (Fig. 3a). It can therefore be concluded, firstly, that species defining the sedimentary assemblage were present in the natural macrophytodebris and crossed the boundary layer downwards. Secondly, that infaunal colonization through the interstitial spaces played a major role, in the occurrence of two contiguous habitats. In case two habitats were not contiguous, recruitment occurred through dispersal through the water column. Thirdly, assemblages in macrophytodebris were a mixture of the surrounding habitat assemblages.

Subsequently, not only habitat-specific species actively migrate as suggested by Hicks (1986). This study concluded that ecologically different copepods originating from diverse habitats were conspicuous dispersers and actively migrated towards defaunated habitats using their species-specific preferred dispersal pathways.

Dispersion and colonization drivers

In our study, hydrodynamic flow effects were excluded, due to the experimental set-up shielding the macrophytodebris. Nonetheless, in natural environments, the recruitment or settlement of water-bound benthic organisms is defined by landscape attributes and hydrodynamic processes (Armonies 1994; Commito and Tita 2002). Palmer (1984) states that meiofauna inhabiting unvegetated habitats avoid the benthic boundary layer during high flow disturbances and frequent the sediment surface during reduced flow. Since above-ground structures locally reduce the hydrodynamic disturbance and diminish the predation risk associated with freely swimming in the water column (Coull and Wells 1983; Palmer and Gust 1985), macrophytodebris stabilizes the hydrodynamic flow and serves as refuge from predators. However, macrophytodebris is only deposited on the seafloor and is not rooted. Therefore, during low hydrodynamic flow, it indeed provides shelter from flow and predation, to the contrary during high hydrodynamic flow or storms, the macrophytodebris and the upper layer of sediment passively (re)suspends in the water column together with their associated organisms (Thistle et al. 1995). Subsequently, the suspended material resettles randomly on different sand patches, driven by hydrodynamic flows. Hence, rafting or drifting on macrophytodebris triggered by storms should be considered as a dispersal method at larger spatial scales (Bonsdorff 1992; Norkko and Bonsdorff 1996; Ólafsson et al. 2001).

High disturbances have an adverse effect on densities; however, intermediate disturbances could have an opposing effect, corresponding to the intermediate disturbance hypothesis (Connell 1978; Cadotte 2007). It states that high diversity is a consequence of continually changing conditions. As a result, disturbance is put forward as the explanation for coexistence of species and often high disturbance resets the local succession pathway (Connell 1978). Under sheltered conditions, macrophytodebris interstitial water slowly changes from water-column-like chemical conditions to oxygen-poor conditions, due to bacterial respiration and reduced compound advection from sediment (Mascart et al. 2015). Community structure significantly changed between the T_0 reference samples and the T_{96} 'Control' samples, without showing a drop in densities. For macrofauna, Gallmetzer et al. (2005) and Remy (unpublished data) found a dominance shift towards low-oxygen tolerant

species, reducing the overall diversity in the macrophytodebris accumulation under stable conditions. Therefore, the experimental units are not expected to have a negative impact on a short-term interval. However, on the long term, a plausible oxygen drop could occur and perturb the initial community structure. Hence, the T_{96} incubations displayed the variability of the community structure and its sensitivity to a potential drop in oxygen levels.

The habitat selectivity of the copepod resettlement (following a passive erosion or active emergence) depends on the chemical and microbiological signals perceived at small distances (Hicks 1977; Fleeger et al. 1995). Decho and Fleeger (1988) stated that copepods exhibit a preference for food-enriched microhabitats over their initial habitat food availability. Macrophytodebris are abundantly colonized by microepiphytes (diatoms, bacteria, fungi, protists) (Lepoint et al., 2006) degrading detrital material or using remineralized nutrients. Macrophytodebris could support a higher microbial biomass and production than living material, since it may have reduced levels of polyphenolic compounds, which in living leaves deter both bacterial colonization and herbivory (Dethier et al. 2014). Degrading seagrass material could be also a minor source of food for few species (Mascart et al. 2013).

In conclusion, our study is congruent with Bostrom and Bonsdorff (2000), whom stated that structurally complex plant assemblages (like macrophytodebris) may trap or attract organisms more efficiently than structurally simple leaf canopies or bare sediments. Structural complexity and dispersion drivers, such as hydrodynamics (i.e. disturbances), habitat complexity (i.e. shelter effect) and food availability, control the colonization ability of a copepod population.

Species-specific behaviour and ecological types

This study rigorously tried to quantify and qualify the species-specific active colonization of copepods from adjacent habitats towards defaunated habitats. Our results showed that the dominant *Sarsamphiascus tenuiremis* (Miracidae family) rapidly colonized macrophytodebris, using two different dispersal pathways, one via the water column and another via the sediment and macrophytodebris interstitial spaces. In order to be able to use the former pathway, one needs to actively emerge from the substrate's surface (e.g. seagrass leaves or sediment) and possess well-developed swimming abilities to than disperse (Thistle and Sedlacek 2004). Other genera, for instance, *Ambunguipes*, *Ameira*, *Amphiascoides*, *Amphiascus*, *Dactylopodella*, *Diosaccus* and *Ectinosoma* seemed to have a similar behaviour in our study. On the other hand, the genera *Arenosetella*, *Canuella*, *Leptastacus*, *Leptopontia*, *Rhizotrix*, *Tetragoniceps* and *Wellsopsyllus* did

not seem to behave in an emergent way and preferably stayed in the sediment where they can interstitially migrate between bordering substrata. This is congruent with the study of Hockin and Ollason (1981) whom differentiated two sets of species, one with superior and one with inferior dispersal abilities. Indeed, several of the above-mentioned species were exclusively present in the sediments, showing no tendency to migrate and therefore can be tagged as species with inferior dispersal abilities. Regarding the species with superior dispersion abilities Kurdziel and Bell (1992) defined seven true phytal-dwelling families on *Thalassia testudinum* in accordance with previous research by Hicks and Coull (1983). Although studying a temperate water seagrass meadow, our results were analogous as those families were all present in the *P. oceanica* canopy and both studies share the same dominant genus, *Harpacticus*. The *P. oceanica* phytal-dwelling species contributed for 50 % to the species composition of naturally occurring macrophytodebris. The prominent role of seagrass species can partially be explained by the origin of macrophytodebris. Phytal copepods have prehensile legs, allowing them to cling to a substrate (Hicks and Coull 1983); therefore, it can be assumed that some species are passively transported attached on the falling dead leaves.

Besides, Kurdziel and Bell (1992) found free-swimming Cyclopoids in the *T. testudinum* canopies. Our study found similar copepods in the macrophytodebris together with calanoids. Hence, free-swimming species can play an important role and represent up to 30–40 % in the densities of the macrophytodebris assemblage. It is known that these copepods migrate vertically to the bottom for shelter, reproduction and feeding purposes (Teasdale et al. 2004). Seen their high abundance in the 'From WC' treatment, especially with time, it should not be excluded that macrophytodebris has an important ecological role for free-swimming copepods from coastal area.

Consequently, macrophytodebris species assemblages are composed of passive leaf clingers, active infaunal dispersers, active suspended dispersers and free-swimming copepods. All those different dispersion modes of adult copepods seem to confirm their importance to population dispersion. Chandler and Fleeger (1983) concluded in their colonization experiment, which suspended water column dispersion is more important than interstitial infaunal transport. However, the latter should not be neglected. Every species with good swimming capacities and capability of emergence (e.g. Armonies, 1994; Kurdziel and Bell 1992; Thistle and Sedlacek, 2004) colonized or showed a predisposition to migrate to new or provisional habitats. Therefore, it would be preferable to refine the provisional classification of Noodt (1971) with a more ecological and mobility-based grouping. Subsequently, in Table 1, the active colonization capabilities of some species were incorporated.

Hicks (1986) predicted the epibenthic capabilities of many phytal species, however, he agreed upon Palmer and Gust (1985) ideas of passive erosion being the only pathway to suspend benthic copepods from unvegetated habitats. The presented results prove that active emergence and migration occur from sedimentary habitats, water column habitats and phytal habitats. Therefore, it would be desirable to correct the idea that exclusively phytal components actively contribute members to the water column, as the results proved that not only phytal-dwelling copepods possess well-developed swimming abilities.

The formation of the communities during low disturbance conditions seems to distinguish two stages. The initial stage is characterized by colonizer species and subsequently after a few days competitor species arrive on site, impacting the subsequent settlement (Sun and Fleeger 1994; Jacobi and Langevin 1996). Colonization experiments over time (e.g. Chertoprud et al. 2005; De Troch et al. 2005) generally observed a stage shift after more than 4 days, the duration of the second incubation experiment. However, an initiation of a stage shift seemed visible in our 96-h experiment. *Ectinosoma dentatum* became the most dominant species in the newly colonized macrophytodetritus after 96 h. Hence, a colonization-competition trade-off seemed to be present between *S. tenuiremis* and *E. dentatum*. However, in the reference samples, *S. tenuiremis* is more abundant than *E. dentatum*. During a seasonal characterization study in the same bay (Mascart et al. 2015), effects of oxygen gradients were measured for *Diosaccus tenuicornis*, a species from the same Miraciidae family as *S. tenuiremis*. To the contrary, *E. dentatum* seemed not to be impacted by the oxygen level present in the macrophytodetritus. De Troch et al. (2005) confirmed the definition of Hicks and Coull (1983) of Ectinosomatidae as ‘jacks of all habitats’, so-called r-strategist. The experiment revealed thus a sensibility of *S. tenuiremis* towards a potential drop in oxygen gradient induced by the experimental setup after 96 h. This could explain the difference between the control and the reference samples after 96 h. Therefore, it can be concluded that the experiment duration was not long enough to see a stage shift, but an effect of drop in oxygen was visible. Nonetheless, it confirms the important role and flexibility of *E. dentatum* within the macrophytodetritus accumulations and the main colonizer role for *S. tenuiremis*. Both *E. dentatum* and *S. tenuiremis* used the infaunal and suspension pathway for active colonization.

Conclusions

In conclusion, free-living harpacticoid copepods actively colonize adjacent defaunated macrophytodetritus and sediment within 24 h. Eco-morphologically different copepods

originating from diverse habitats (sediment, phytal and pelagic) were conspicuous dispersers and actively migrated towards defaunated habitats using their species-specific preferred dispersal pathways. Two pathways occur for active colonization: infaunal migration through the boundary layer and migration suspended via the water column. Eco-morphological characteristics can be used to predict the potential preferred habitat of copepods. However, copepods are not obligate residents of their preferred habitat since phytal and sedimentary components actively contribute to the water column. Therefore, copepods are opportunistic and are capable of assembling dynamic communities in provisional habitats. Macrophytodetritus offers a structurally complex substitute habitat proposing shelter and food supply, therefore serving as a hub for all copepods of the surrounding habitats. Due to the high diversity of associated fauna, the macrophytodetritus assemblage has a certain resilience which can cope and restore swiftly from disturbances. Swimming abilities, structural complexity and dispersion drivers, such as hydrodynamics, habitat complexity and food availability, control the colonization ability of a copepod population.

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