# ORIGINAL PAPER

# Recently established *Crassostrea*-reefs versus native *Mytilus*-beds: differences in ecosystem engineering affects the macrofaunal communities (Wadden Sea of Lower Saxony, southern German Bight)

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Received: 4 January 2008/Accepted: 8 January 2009/Published online: 30 January 2009 © Springer Science+Business Media B.V. 2009

Abstract Since 1998 the non-indigenous Pacific oyster Crassostrea gigas (Thunberg 1793) has been invading the Wadden Sea of Lower Saxony, southern German Bight. C. gigas settles predominantly on intertidal Mytilus-beds (M. edulis L.) and subsequently create rigid reef-like structures. Both bivalve species are ecosystem engineers in sedimentary tidal flats. They provide hard substrate for sessile species, mobile organisms find refuge within the habitat matrix of dense suspension feeders, and biodeposits enrich the sediments with organic matter. The transformation of Mytilus-beds into Crassostrea-reefs gives rise to the question whether the invader may affect the native community. We investigated two parts of a changing bivalve bed in the backbarrier area of the island of Juist in March 2005. One part was still dominated by *M. edulis* whereas the other part was already densely colonized by C. gigas. Crassostrea-reefs compensate for the conceivable loss of Mytilus-beds in the intertidal of the Wadden Sea by replacing the ecological function of M. edulis. There was no

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I. Kröncke e-mail: ingrid.kroencke@senckenberg.de indication of a suppression of indigenous species. This even applied to *M. edulis*, which persisted at the site invaded by *C. gigas*. The associated macrofaunal community showed increased species richness, abundance, biomass, and diversity in the *Crassostrea*-reef. The latter particularly favored sessile species like anthozoans, hydrozoans, and barnacles. Higher abundance and biomass for vagile epizoic species like the shore crab *Carcinus maenas* and the periwinkle *Littorina littorea* also occurred among oysters. Abundance of deposit feeding oligochaetes was enhanced by oysters as well. More opportunistic, facultative filter-feeding polychaetes occurred in the *Crassostrea*-reef.

**Keywords** Bioinvasion · Biozoenosis · *Crassostrea* gigas · Diversity · Ecosystem engineering · Macrofauna · *Mytilus edulis* · Neozoa · North Sea

# Introduction

More and more marine species have been transported across natural barriers in recent times (Carlton and Geller 1993; Carlton 1987; Nehring 1998; Gollasch 2002). When they become established, the nonindigenous species may threaten native biodiversity as well as ecosystem functions of the region (Kolar and Lodge 2001). However, introduced species may also create habitat structures which locally may enhance biological diversity (Crooks 1998).

For the purpose of aquaculture, the Pacific oyster Crassostrea gigas (Thunberg 1793) was first imported 1964 into the Easter Scheld estuary (The Netherlands) which is connected to the open North Sea. Natural reproduction was not expected due to low water temperatures. But after spatfalls in the warm summers of 1975, 1976, and 1982, C. gigas established itself outside the culture plots in the Easter Scheld estuary (Drinkwaard 1999). In 1983 the first Pacific oyster was found in the Dutch Wadden Sea near Texel (West Frisian Island), which appeared to have been deliberately introduced (Dankers et al. 2004). First specimens in the East Frisian Wadden Sea (Wadden Sea of Lower Saxony, southern German Bight) were recorded in 1998 from the backbarrier area of the island of Baltrum most likely originating from eastward larval drift via Texel (Wehrmann et al. 2000; Brandt et al. 2008). Massive spatfalls of Crassostrea in 2003 and 2004 resulted in an exponential increase in abundance. In 2005, up to 700 individuals  $m^{-2}$  were found in the backbarrier of the island of Juist (Wehrmann et al. 2006).

In the Wadden Sea, the colonization by invasive Pacific oysters occurred predominantly on intertidal *Mytilus*-beds (*M. edulis* L.), which provide secondary hard substrate for the settlement of *C. gigas* larvae (Reise 1998; Wehrmann et al. 2000; Diederich 2005). The indigenous but extirpated European oyster *Ostrea edulis* L. was found in subtidal environments of the Wadden Sea (Möbius 1877). Thus, *C. gigas* is not occupying the open niche of the native oyster but intertidal areas already occupied and affected by *M. edulis* (Reise 1998).

Mytilus edulis is an ecosystem engineer in the sediment dominated tidal flats of the Wadden Sea. Compared to ambient tidal flats, Mytilus-beds are "Islands of biodiversity" (Buschbaum and Nehls 2003). They offer habitats for several sessile species like barnacles, anthozoans, hydrozoans, bryozoans and macroalgae. They provide shelter and offer food resources to various mobile epibenthic organisms like crustaceans and molluscs. Mytilus-beds have been termed a biosedimentary system (Hertweck and Liebezeit 1996). Biodeposits enrich the sediments of patches and open spaces with organic matter (Dittmann 1987), enhancing infaunal diversity. Even areas adjacent to a Mytilus-bed are influenced by biogenic deposition (Kröncke 1996; Bergfeld 1999; Hild and Günther 1999). Additionally, Mytilus-beds are an important food resource for many fish species, a variety of bird species and humans.

According to Linke (1954) mature *Mytilus*-beds are composed of *Mytilus*-covered "patches" and interspersed bare sediments as *Mytilus*-free "open spaces". *M. edulis* creates flexible three-dimensional structures as multiple individuals connect and disconnect their shells by byssus threads. The *Mytilus*meshwork covers the sediment of the patches in horizontally oriented layers. Due to their mobility, the individuals mostly manage to stay above the sediment.

In contrast, *Crassostrea*-reefs constitute a new biogenic structure in the Wadden Sea (Reise et al. 2005). *C. gigas* larvae prefer settling on conspecifics (Arakawa 1990). The process of multiple settlement leads to the creation of stable clusters as the individuals stay cemented to their substrate of settlement. In patches of high oyster density, the rigid reef-like structures tend to be vertically oriented. Over time, the lower part in *Crassostrea*-patches becomes buried with sediment whereas the upper part remains exposed. By constructing reefs and producing biodeposits, *C. gigas* may influence habitat features in a similar way like *M. edulis* (Reise 2002; Ruesink et al. 2005). However, differences in ecosystem engineering may alter the resident biocoenosis.

The aim of this study was to investigate whether and how the bioinvasion of the Pacific oyster affects the recipient biota. To analyze community effects we chose one bivalve bed in the backbarrier area of the island of Juist (Fig. 1) where patches of high *Crassostrea* density co-occurred with still remaining *Mytilus*-patches. We focused on the following question: are there differences in species number, abundance, biomass and diversity of the associated macrofauna between *Mytilus*-beds and *Crassostrea*reefs?

#### Methods

Study site, sampling, and sample treatment

The Wadden Sea of Lower Saxony covers an area of  $1200 \text{ km}^2$ . 108 intertidal bivalve beds are recorded which cover approx. 1% of the tidal flats. Sampling took place in March 2005 on one 37 ha large intertidal bivalve bed, located in the backbarrier tidal

Fig. 1 Top: Study area in the Wadden Sea of Lower Saxony, south of the East Frisian Island of Juist, southern German Bight, North Sea. Blow-up shows the bivalve bed and the locations of the sampling sites S (sand flat, 100 m outside the bivalve bed), M (Mytilus-bed) and C (Crassostrea-reef). Bottom: Sampling design shows the sampling station S (sand flat), Mytilus-patch (covered by Mytilus), M<sub>COV</sub> (underlying sediment) and  $M_{\rm OS}$  (bare open spaces between the Mytiluspatches), Crassostrea-patch (covered by Crassostrea),  $C_{\rm COV}$  (underlying sediment),  $C_{OS}$  (bare open spaces between the Crassostrea-patches)



sand flats south of the East Frisian Island of Juist, southern German Bight, North Sea (N 53°38'; E 006°56'; Fig. 1). In the area we find a salinity of ~30 psu, an average temperature of 10.9°C (04/ 2004-03/2005) and an average tidal range of 2.4 m. Compared to other intertidal bivalve beds in the area this bed is characterized by a consistently high occurrence of *M. edulis* over the last decades (Herlyn and Millat 2004). First individuals of *C. gigas* were found in 1999 but a massive spread occurred not till 2003. The positive feedback in *Crassostrea*  settlement (Diederich 2005; Wehrmann et al. 2006) led to a rapid reef formation in the western part of the bivalve bed in 2003/2004 (*Crassostrea*-reef, site C, Fig. 1), while at the time of the investigation the eastern part still remained dominated by *M. edulis* (*Mytilus*-bed, site M, Fig. 1). Over a distance of 700 m from one site to the other a continuous transition in *C. gigas* density occurred. Tidal elevation with a mean exposure time at low tide of about 3.5 h per tidal cycle and sediment characteristics were similar at both sites; thus no obvious differences between the sites occurred which could entail changes in associated fauna. Both sites consist of bivalve-covered patches ("*Mytilus*-patches", "*Crassostrea*-patches") and of interspersed epibenthic bivalve-free bare sediment ("open spaces").

We took epibenthos samples in Mytilus-patches and Crassostrea-patches (Fig. 1) with a 15 cm diameter tube ( $\sim 177 \text{ cm}^{-2}$ ). Directly after clearing the patches from epibenthic bivalve coverage, the underlying sediment (M<sub>COV</sub>, C<sub>COV</sub>) was taken with a 5.9 cm diameter tube ( $\sim 27 \text{ cm}^{-2}$ ) down to 30 cm sediment depth (Fig. 1). Additionally, we took sediment samples with a 5.9 cm diameter tube down to 30 cm sediment depth in the epibenthic bivalve-free open spaces at site M and site C (M<sub>OS</sub>, C<sub>OS</sub>) and in the epibenthic bivalve-free adjacent sand flat (site S) (Fig. 1). The latter was located 100 m outside the bivalve bed area. Tidal elevation and sediment characteristics of the sand flat were similar as at site M and C. By taking three replicates per station (Mytilus-patch, Crassostrea-patch, M<sub>COV</sub>, C<sub>COV</sub>, M<sub>OS</sub>, C<sub>OS</sub> and S) a total of 21 samples were analyzed.

All samples were fixed in 4% buffered formaldehyde and stored for at least 3 months to level off weight losses (Hamilton and Kingston 1985; Böttger and Schnack 1986; Brey 1986). The samples were then sieved through 0.5 mm mesh size in the laboratory. All remaining organisms were collected from the recovered material. Shell lengths of M. edulis and C. gigas specimens were measured to the nearest mm. The flesh of each specimen was stored separately in 70% ethanol before determining ash free dry weight in gram (g AFDW) after drying and combusting in a furnace using standard methods (Rumohr 1990). All other macrobenthic fauna of the epibenthos samples and of the sediment samples were determined to species level, counted and stored in 70% ethanol. Biomass (g AFDW) was calculated from wet weight using published conversion factors (Brey 2001; Ricciardi and Bourget 1998). For barnacles a conversion factor was calculated based on data of this study.

#### Data analysis

In this survey individuals of *M. edulis* <6 mm shell length were defined as juveniles of the previous spatfall (late summer 2004) and individuals of *C. gigas* <21 mm were defined as juveniles of the spatfall in early autumn 2004. To evaluate differences between the sites M and C, we quantified abundance and biomass (g AFDW) of adult *M. edulis* and adult *C. gigas* as arithmetic means per m<sup>2</sup> patch with standard deviation ( $\pm$ ), calculated mean shell length (mm) of both adult bivalve species (Table 1) as well as the shell length frequency distribution of *M. edulis* in *Mytilus*-patches and *Crassostrea*-patches.

Mytilus edulis and Crassostrea gigas were excluded from the analysis of the associated macrofauna of the epibenthos samples. Species richness results from total species number of all replicates, respectively. Species were categorized after their mode of life as sessile and vagile epifauna and infauna (Tables 2, 3). Colonies forming sessile taxa were grouped separately, noted as present/absent and were excluded from quantitative analysis. Abundance and biomass are presented as arithmetic means per  $m^2$ (rounded values). To detect similarities in community structure ordination (Multi-Dimensional Scaling, MDS) and classification (Hierarchical cluster) methods (Kruskal and Wish 1978) were applied, using the Bray-Curtis similarity coefficients (Bray and Curtis 1957). Diversity of Mytilus-patches and Crassostreapatches was determined by the Shannon-Wiener Index (Weaver and Shannon 1949) and by the Evenness (Pielou 1966), calculated for biomass data.

To evaluate differences between all stations, we compared the infauna of the sand flat (S) with the infauna of the Mytilus-bed (M) and the Crassostreareef (C), sites M and C subdivided into the stations "patch" and "open space", respectively. C. gigas clusters were sticking up to 5 cm in the sediment of the patches. Therefore, we picked up more sediment by taking the epibenthos samples in Crassostreapatches than in Mytilus-patches. To obtain comparable data for the infauna analysis, we added the infauna data of the epibenthos samples to the infauna data of the sediment samples, which we took beneath the Mytilus- and Crassostrea-cover in the patches (M<sub>COV</sub>, C<sub>COV</sub>), respectively. Species richness of the infauna communities at the five stations results from total species numbers, respectively. Abundance of infauna is presented as arithmetic means per m<sup>2</sup> (rounded values). We classified infauna species into higher taxonomical groups. Polychaetes, the main group of the macrobenthic species assemblages at all stations, were categorized into feeding modes (Fauchald and Jumars 1979; Hartmann-Schröder 1996).

#### Results

Ecosystem engineers: *Mytilus edulis* and *Crassostrea gigas* 

Total mean abundance of both ecosystem engineers in *Crassostrea*-patches was almost three times higher and total mean biomass was more than twice higher than in *Mytilus*-patches (Table 1).

*Mytilus*-patches were dominated by its ecosystem engineer *Mytilus* which accounted for 92% of all adult individuals and 95% of the total biomass (Fig. 2). In *Mytilus*-patches, adult *Crassostrea* stayed low in abundance and due to small individuals as well biomass of the oysters was low (Table 1).

In contrast, *Crassostrea*-patches were built up by 54% adult oysters and 46% adult blue mussels. Due to large individuals (Table 1), *Crassostrea* dominated the *Crassostrea*-patches in biomass (Fig. 2). The largest *C. gigas* specimen in *Crassostrea*-patches was 179 mm in length.

The abundance of adult *Mytilus* in *Crassostrea*patches was higher than in *Mytilus*-patches but was represented by smaller individuals (Table 1; Figs. 2, 3). At both stations, *Mytilus* was found in all size classes up to 70 mm but we detected a distinct gap in medium size classes between 26 and 45 mm in *Mytilus*-patches whereas these size classes were abundant in *Crassostrea*-patches (Fig. 3). Hence, biomass of *Mytilus* was lower in *Crassostrea*-patches than in *Mytilus*-patches (Fig. 2). Juvenile *Mytilus* were abundant at both stations but *Mytilus*-patches had 1.5 times more juveniles than *Crassostrea*patches. Juveniles accounted for 73% of the *Mytilus* population in *Mytilus*-patches (3597  $\pm$  887 juvenile *Mytilus* m<sup>-2</sup>) and 55% of the *Mytilus* population in *Crassostrea*-patches (2335  $\pm$  974 juvenile *Mytilus* m<sup>-2</sup>).

Associated macrofaunal communities

In total we identified 59 macrofaunal taxa (Tables 2, 3). According to their mode of life, 31 taxa were assigned to the infauna and 28 taxa to the epifauna. The latter were subdivided in 12 vagile and 16 sessile taxa. The sessile epibiont community included 9 colonies forming taxa (ascidian, bryozoans, hydrozoans), which were noted as present/absent and are excluded from quantitative analysis (Table 2). Most common were the polychaetes with 21 taxa.

Epibenthos samples: species richness, abundance and biomass of associated macrofauna in Crassostrea- and Mytilus-patches

We identified 28 macrofaunal taxa in *Mytilus*-patches and 45 in *Crassostrea*-patches. Six species were found exclusively in *Mytilus*-patches and 23 exclusively in *Crassostrea*-patches, while 22 taxa were found at both stations (Table 2). Infauna had highest species numbers at both stations, followed by vagile epifauna, colonies forming sessile taxa and other sessile epifauna (Fig. 4). The higher total number of species in *Crassostrea*-patches was reflected in all groups. Infauna increased from 14 to 19 species, vagile epifauna from 7 to 10 species, colonies forming taxa from 4 to 9 species and other sessile epifauna from 3 to 7 species. Infauna and vagile

Table 1	Mean shell ler	igth (SL) ii	n mm, mean	abundance	e and mean l	biomass in g	gram ash f	free dry w	eight (g AFD	$W) \pm SD pe$	r m² of
adult ind	ividuals of the	two ecosy	stem engine	ers Mytilu.	s edulis and	Crassostre	a gigas in	n Mytilus-	patches and	Crassostrea-	patches

Ecosystem engineers	SL (mm)	Individuals $m^{-2}$	g AFDW m <sup>-2</sup>
<i>Mytilus</i> -patch		1412 ± 539	$283.9 \pm 74.8$
Mytilus edulis	$41 \pm 23$	$1318 \pm 534$	$271.4 \pm 67.7$
Crassostrea gigas	$30 \pm 27$	$94 \pm 86$	$12.5 \pm 11.0$
Crassostrea-patch		$4011 \pm 1314$	624.9 ± 152.5
Mytilus edulis	$30 \pm 15$	$1883 \pm 690$	$204.5\pm74.1$
Crassostrea gigas	$67 \pm 34$	$2128 \pm 625$	$420.4 \pm 86.1$

Total abundance/biomass in bold

Fig. 2 Mean abundance (left) and biomass (right) in gram ash free dry weight (g AFDW) with standard deviation per m<sup>2</sup> of adult Mytilus edulis and adult Crassostrea gigas in Mytilus-patches and Crassostrea-patches; proportion of total values inside bars





Mvtilus-patch

57.55 56.60 67.<sub>65</sub> 66.<sub>20</sub> 2 2 2 C ? (mm) (mm) shell length Crassostrea-patch shell length

epifauna decreased in relative species numbers in Crassostrea-patches, whereas the sessile epibiont species assemblage was emphasized (Fig. 4). Anthozoans were exclusively found in Crassostreapatches (Table 2).

Total mean abundance and total mean biomass in Crassostrea-patches were twice as high as they were in Mytilus-patches (Table 2; Fig. 5). An increased abundance was detected for all functional groups in Crassostrea-patches whereas the increase in biomass was related to vagile epifauna.

Sessile epifauna was most abundant at both stations and dominated the associated macrofauna community with over 80% at both stations (Fig. 5). In Crassostreapatches, 1.4 times more sessile epifauna individuals were found. Vagile epifauna had lowest abundances at both stations but as well 1.4 times more individuals were found in Crassostrea-patches. Whereas total biomass of associated macrofauna was still dominated by sessile epifauna in Mytilus-patches (75%), sessile epifauna biomass only made up 45% of the total biomass in Crassostrea-patches (Fig. 5). Despite of more but smaller barnacles, sessile epifauna had a lower biomass in Crassostrea-patches than in Mytiluspatches. The higher total biomass in Crassostreapatches (Table 2; Fig. 5) was related to higher abundances of large individuals of vagile epifauna species like the periwinkle Littorina littorea and the shore crab Carcinus maenas as well as the exclusive occurrence of the polychaete *Harmothoe imbricata* and the starfish Asterias rubens (Table 2). Biomass of vagile epifauna was four times higher in Crassostrea-patches and contributed with 50% to the biomass of the associated macrofauna (Table 2; Fig. 5).

Infauna showed the highest increase in abundance. 3.4 times more infaunal individuals were encountered in Crassostrea-patches than in Mytilus-patches. In contrast, infauna biomass was only twice higher in Crassostrea-patches. Due to the high occurrence of oligochaetes, Tubificoides benedii being dominant, infauna abundance gained in importance in Crassostrea-patches and reached 16% of associated macrofauna individuals (Fig. 5).

# Epibenthos samples: multivariate analyses and diversity

Multi-Dimensional Scaling performed with abundance data after 4th root transformation resulted in a separation of the species assemblages "Mytilus-patch"

67%

33%

Crassostrea-

patch

n = 4219

adult Mytilus edulis

Associated macrofauna	Epibenthos samples					
	<i>Mytilus</i> -patch		Crassostrea-patch			
Taxon ID species	Mode of life	Individuals $m^{-2}$	g AFDW m <sup>-2</sup>	Individuals m <sup>-2</sup>	g AFDW m <sup>-2</sup>	
Anthozoa						
1 Actinia equina	s epi			170	0.2333	
2 Anthozoa sp.	s epi			19	0.0004	
3 Corynactis viridis	s epi			19	0.0047	
4 Sagartia elegans	s epi			113	1.9344	
Amphipoda	-					
5 Melita palmata	v epi			132	0.0349	
Bivalvia	-					
6 Angulus tenuis	in			19	2.5550	
7 Bivalvia sp.	in			19	0.0002	
8 Cerastoderma edule	in	19	0.0002	38	0.0196	
9 Macoma balthica	in	19	2.5446			
10 Scrobicularia plana	in	19	0.0006			
11 Venerupis senegalensis	in	19	0.0023			
Cirripedia						
12 Balanus crenatus	s epi	57	0.4225	396	0.9488	
13 Elminius modestus	s epi	63247	59.4731	88995	55.3937	
14 Semibalanus balanoides	s epi	641	5.1841	245	1.2576	
Collembola						
15 Anurida maritima	v epi	75	0.0015			
Decapoda	-					
16 Carcinus maenas	v epi	113	0.8835	717	18.9747	
Echinodermata	-					
17 Asterias rubens	v epi			19	0.7851	
Isopoda	-					
18 Jaera albifrons	v epi	472	0.0483	547	0.0664	
Nemertea	-					
19 Lineus ruber	in	264	0.1068	226	0.1926	
20 Malacobdella sp.	v epi	1056	0.0113	264	0.0030	
21 Tetrastemma sp.	in	19	0.0094	38	0.0026	
22 Nemertini sp.	in			94	0.0283	
Oligochaeta						
23 Oligochaeta sp.	in	245	0.0130	2075	0.2861	
24 Oligochaeta spp.	in	736	0.0302	1943	0.1368	
25 Tubificoides benedii	in	1867	0.1886	9884	1.7931	
Opisthobranchia						
26 Aeolidia papillosa	v epi			19	0.0217	
Polychaeta						
27 Ampharete acutifrons	in			19	0.1409	
28 Aphelochaeta marioni	in	1226	0.3327	1056	0.4516	
29 Capitella capitata	in	585	0.1217	151	0.2494	

**Table 2** Mean abundance and mean biomass in gram ash free dry weight (g AFDW) per  $m^2$  of associated macrofauna species of epibenthos samples in *Mytilus*-patches and *Crassostrea*-patches

Table 2 continued

Associated macrofauna	Epibenthos samples					
	Mode of life	Mytilus-patch		Crassostrea-patch		
Taxon ID species		Individuals $m^{-2}$	g AFDW $m^{-2}$	Individuals $m^{-2}$	g AFDW m <sup>-2</sup>	
30 Eulalia viridis	v epi			19	0.0051	
31 Harmothoe imbricata	v epi			75	2.0079	
32 Heteromastus filiformis	in			57	0.1332	
33 Lanice conchilega	in			38	0.2773	
34 Microphthalmus sczelkowii	in			302	0.0043	
35 Neanthes succinea	in			472	0.3982	
36 Phyllodoce mucosa	v epi	19	0.0419			
37 Polychaeta sp.	in	38	0.0004	207	0.0036	
38 Polydora ciliata	in	189	0.0147	773	0.0685	
39 Polydora cornuta	in			755	0.1168	
40 Pygospio elegans	in	38	0.0040			
Polyplacophora						
41 Lepidochitona cinereus	v epi	207	1.2299	132	0.3567	
Prosobranchia						
42 Littorina littorea	v epi	604	15.5750	1603	45.2208	
Total species number		24		36		
Total abundance/biomass		71774	86.2403	111650	134.1073	
Ascidiacea						
43 Perophora listeri	col			+		
Bryozoa						
44 Chartella cf. papyracea	col			+		
45 Conopeum reticulum	col	+		+		
46 Electra pilosa	col			+		
Hydrozoa						
47 Coryne sp.	col			+		
48 Hydrozoa spp.	col			+		
49 Laomedea flexuosa	col	+		+		
50 Obelia cf. dichotoma	col	+		+		
51 Tubularia sp.	col	+		+		
Species richness		28		45		

Listed are species ID and mode of life as infauna (in), vagile epifauna (v epi), sessile epifauna (s epi) or colonies forming taxon (col), the latter noted as present (+)/absent

Total species number, total abundance/biomass and species richness per station in bold

and "*Crassostrea*-patch". The corresponding cluster analysis confirmed this grouping. The cluster of the three replicates in *Crassostrea*-patches had a similarity of 70% and a similarity of even 80% in *Mytilus*patches (Fig. 6).

With reference to a mean density of  $26.3 \pm 1.2$  species in *Crassostrea*-patches, the Evenness for biomass was  $0.44 \pm 0.06$ . In *Mytilus*-patches mean species density was only  $17.7 \pm 1.2$ , Evenness was

 $0.35 \pm 0.07$ . The mean Shannon–Wiener index was higher in *Crassostrea*-patches (1.43  $\pm$  0.21) than in *Mytilus*-patches (0.99  $\pm$  0.24; Fig. 7).

# Sediment samples: species richness and abundance of infauna

In total we identified 31 infaunal taxa. The sand flat and stations covered by epibenthic bivalves, the



ecosystem engineers *Mytilus edulis* or *Crassostrea* gigas, tended to be higher in species number than the sediments in the open spaces of the *Mytilus*-bed and the *Crassostrea*-reef. Endobenthic amphipods were only found in the sand flat, with *Urothoe poseidonis* being dominant. Nemertines only occurred at stations covered by the epibenthic bivalves, with *Lineus ruber* being dominant. Only the sediments in the open spaces of the *Mytilus*-bed and the *Crassostrea*-reef were free of endobenthic bivalves. Annelids were found at all stations. With 16 species polychaetes were the most diverse infaunal group and dominated the species assemblages of all stations (Table 3; Fig. 8).

In terms of total infauna abundance, we detected highest values in patches and open spaces of the *Crassostrea*-reef. Abundance was conspicuously lower in the open spaces of the *Mytilus*-bed and in the sand flat, whereas the *Mytilus*-patches had an intermediate density (Table 3; Fig. 8). Annelids were the main taxa in structuring infauna abundance. They had a share of over 95% at all stations but the sand flat. Here amphipodes achieved a high proportion of 35%. Endobenthic bivalves and nemertines stayed low in abundance at all stations of occurrence.

Oligochaetes dominated stations covered by the epibenthic bivalves. The relative abundance of oligochaetes increased from the sand flat (5%) via the open spaces ( $M_{OS}$  20% und  $C_{OS}$  39%) to the stations covered by the epibenthic bivalves ( $M_{COV}$  53%,  $C_{COV}$  76%). Highest infauna abundance was found in *Crassostrea*-patches. A comparable lower abundance as in *Mytilus*-patches was encountered in the open spaces of the *Crassostrea*-reef. Lowest oligochaete abundance was found in the open spaces of the *Mytilus*-bed and in the sand flat (Table 3; Fig. 8). At all stations *Tubificoides benedii* was the most abundant species.

Polychaetes dominated the stations without epibenthic bivalve coverage (Fig. 8). The highest Fig. 6 Multi-Dimensional Scaling (MDS) and cluster analysis for associated macrofauna based on abundance data (after 4th root transformation); plotted are the three replicates in Mytiluspatches (M1-3) and in Crassostrea-patches (C1-3)

Fig. 7 Mean species numbers (left), Shannon-Wiener indices (middle) and Evenness (right) of associated macrofauna calculated for biomass of the three replicates in Mytilus-patches and in Crassostrea-patches, respectively; mean values as numerical data

abundance of polychaetes was found in the open spaces of the Crassostrea-reef. Almost all polychaete species had highest abundances at this station (Table 3). In the open spaces of the Mytilus-bed we found the lowest abundance of all stations. An intermediate abundance of polychaetes was found in Crassostrea-patches, in Mytilus-patches and in the sand flat (Table 3; Fig. 8). The latter was dominated by Pygospio elegans. All stations of the bivalve bed were dominated by the abundance of Aphelochaeta marioni (Table 3). Highest abundance of A. marioni was detected in the open spaces of the Crassostreareef. Capitella capitata was a common species in the sand flat, in Mytilus-patches and in the open spaces of the Crassostrea-reef. Neanthes succinea as an exclusive species in Crassostrea-patches was of notable importance. The predatory polychaete ranked third after A. marioni and the spionid polychaetes of the genus Polydora. Scoloplos armiger was only encountered at stations without epibenthic bivalve coverage. Highest abundance was found in the sand flat (Table 3).

#### Polychaetes: feeding mode

Considering infaunal and epifaunal polychaetes, surface deposit feeders dominated all stations. Facultative filter feeders, resp. Polydora cornuta and P. ciliata, reached a high proportion of 44% in Crassostrea-patches. Subsurface deposit feeders attained a marked percentage only in the sand flat (Fig. 9).

#### Discussion

Ecosystem engineers: Mytilus edulis and Crassostrea gigas

The Mytilus-patches were dominated by M. edulis. In contrast, the Crassostrea-patches were characterized by almost equal abundances of both ecosystem engineers. We even found more Mytilus in Crassostrea-patches than in Mytilus-patches but we detected differences in shell length frequency distribution of M. edulis at the two stations.



MDS: abundance (4th root)

M1

C1

C3

C2

Crassostrea-

patch

Stress: 0

M2

**M**3

Mvtilus-

patch

Cluster: abundance (4th root/group average)

60

70

80

90

100

M3

M2 M1

Mytilus-

patch

C3

C1 C2

patch

Crassostrea-

patch

similarity

**Table 3** Mean abundance per  $m^2$  of infauna species of the sediment samples in the sand flat (S), in *Mytilus* covered areas (M<sub>COV</sub>), in *Crassostrea* covered areas (C<sub>COV</sub>) and in the respective bare open spaces of the bivalve bed (M<sub>OS</sub> and C<sub>OS</sub>)

Associated macrofauna	Sediment samples (individuals m <sup>-2</sup> )						
Taxon ID infauna species	S	M <sub>OS</sub>	M <sub>COV</sub>	C <sub>OS</sub>	C <sub>COV</sub>		
Amphipoda	2439						
52 Amphipoda sp.	122						
53 Bathyporeia nana	122						
54 Urothoe poseidonis	2195						
Bivalvia	244		76		76		
6 Angulus tenuis					19		
7 Bivalvia sp.					19		
8 Cerastoderma edule	122		19		38		
9 Macoma balthica	122		19				
10 Scrobicularia plana			19				
11 Venerupis senegalensis			19				
Nemertea			283		358		
19 Lineus ruber			264		226		
21 Tetrastemma sp.			19		38		
22 Nemertini sp.					94		
Oligochaeta	366	488	5531	5974	14390		
23 Oligochaeta sp.	122		733	610	2075		
24 Oligochaeta spp.	122		2199	975	1943		
25 Tubificoides benedii	122	488	2599	4389	10372		
Polychaeta	3903	1951	4515	9389	4195		
27 Ampharete acutifrons				122	19		
28 Aphelochaeta marioni	610	1219	2445	6340	1056		
55 Arenicola marina	122						
29 Capitella capitata	610		951	975	151		
56 Hediste diversicolor		122					
32 Heteromastus filiformis			122		179		
33 Lanice conchilega	244		244	244	38		
34 Microphthalmus sczelkowii	366	122	366	610	302		
35 Neanthes succinea					472		
57 Nephtys hombergii	122	122		122			
37 Polychaeta sp.			38	244	207		
38 Polydora ciliata			189		895		
39 Polydora cornuta					876		
40 Pygospio elegans	1219	122	160	366			
58 Scoloplos armiger	610	122		366			
59 Spionida sp.		122					
Total abundance	6952	2439	10405	15363	19019		
Species richness	16	8	17	12	19		

Listed are species ID and abundance of higher taxa, total abundance and species richness per station in bold



**Fig. 8** Total species numbers (*left*) and mean abundance with standard deviation per  $m^2$  (*right*) of infauna in the sand flat (S), in *Mytilus* covered areas (M<sub>COV</sub>), in *Crassostrea* covered areas (C<sub>COV</sub>) and in the respective bare open spaces of the bivalve

This study revealed a spatfall of *M. edulis* across the entire bivalve bed. Whereas Diederich (2005) stated that recruitment of *Mytilus* larvae was similar in *Crassostrea*-reefs and in *Mytilus*-beds, we found a lower abundance of juvenile *Mytilus* in the *Crassostrea*-patches. Troost et al. (2004) reported a more diffuse inhalant feeding current of *C. gigas*. Low swimming velocities of *Mytilus* veligers might increase the chance of being filtered by *C. gigas*.

However, mortality of juvenile *Mytilus* after successful recruitment seems to be lower in *Crassostrea*-invaded areas than in *Mytilus*-dominated areas. Hiding-places between individuals of *C. gigas* may have protected *M. edulis* from predation. However, up to the time of the investigation, birds as main predators were avoiding biogenic structures dominated by *C. gigas* (Wehrmann et al. 2006). The absence of medium size classes of *M. edulis* in *Mytilus*-beds was reported by Zens et al. (1997) as

bed ( $M_{OS}$  and  $C_{OS}$ ); taxonomical composition based on species number (*left*), based on abundance (*right*) and proportional occurrence in circle diagrams

related to increasing bird populations, as well as to increasing predation caused by a decrease of *Mytilus*bed coverage in the area (Herlyn and Michaelis 1996). Specimens of *Mytilus* <20 mm and >50 mm are rejected by oystercatchers (Haematopus ostralegus) (Meire and Ervynck 1986; Meire 1996; Zwarts et al. 1996). Birds like oystercatchers, hunting by vision control (Zwarts et al. 1996), might have been subjected to optical delusion. At first glance, Mytilus seems to be absent in Crassostrea-patches. Therefore, predation by birds which appears to be stronger during winter (Meire 1996) coincides with the higher abundance of medium sized M. edulis in the Crassostrea-patches and the sparse occurrence of 25-50 mm Mytilus in the Mytilus-patches (Fig. 3). In the meantime, oystercatchers in the Wadden Sea of Lower Saxony are frequently observed on the reefs and now prey predominantly on Crassostrea (unpublished data).

Fig. 9 Frequency of polychaetes categorized into feeding modes in the sand flat (S), in *Mytilus* covered areas ( $M_{COV}$ ), in *Crassostrea* covered areas ( $C_{COV}$ ) and in the respective bare open spaces of the bivalve bed ( $M_{OS}$  and  $C_{OS}$ )





#### Associated macrofaunal communities

Crassostrea-reefs compensate for the conceivable loss of Mytilus-beds in the intertidal of the Wadden Sea by replacing the ecological function of *M. edulis*. The reefs seem to be an alternative habitat for species typically found in former Mytilus-beds. At the time of the investigation we did not observe a displacement of native macrofaunal species by the invasion of the nonindigenous Pacific oyster C. gigas. Besides C. gigas, no new invasive species was observed within this study. Except for Corynactis viridis, all species identified in this study have already been reported for Mytilus-beds of tidal flats in the German Wadden Sea. However, the literature data about macrofaunal species associated with Mytilus in the intertidal of the Wadden Sea originates from different references and is a sum of occurring species over the last 20-30 years (e.g. Dörjes 1978, 1992; Grotjahn 1987; Herlyn and Millat 2004). The composition of the community associated with "Mytilus-beds" has been shown to be highly variable over time. Whereas species are found in high abundances in 1 year they are absent in other years. In contrast, Crassostrea-reefs seem to provide a suitable habitat for a synchronous appearance of more common species. We found a statistical separation of the macrofaunal communities "Mytilus-patch" and "Crassostrea-patch". We calculated higher values for species richness, abundance, biomass and diversity in Crassostrea-patches. In the following, these findings are discussed in terms of differences in ecosystem engineering by C. gigas versus M. edulis.

The *Crassostrea*-reef might have influenced the frequency of epibenthic organisms by providing a more complex habitat matrix with an extended hard substrate surface. Increased biodeposition might have led to an enhanced enrichment of the sediments in the *Crassostrea*-patches and in the open spaces between the patches, affecting the infauna.

# Availability of hard substrate

The tidal flats of the Wadden Sea are dominated by soft sediments. Thus, *M. edulis* and *C. gigas* function as ecosystem engineers, offering a suitable hard substrate for the settlement of sessile organisms like barnacles (Buschbaum and Nehls 2003). The triple density of both ecosystem engineers in the *Crassostrea*-reef and the mean shell length of 67 mm of

*C. gigas* implies an extended shell surface area and therefore led to the higher abundance of sessile organisms, e.g. of the barnacle *E. modestus*. We found anthozoans only in the *Crassostrea*-reef and hydrozoans as well as bryozoans were more diverse. The extended surface area of available hard substrate in the newly created habitat may as well have promoted the higher abundance of the periwinkle *L. littorea*.

# Habitat matrix

The topography and structural complexity of biogenic habitats affects the ecological role of a habitat (Bell 1991). At low tide bivalve beds represent a moist habitat which diminishes the risk of dehydration. Compared to the flexible Mytilus-meshwork, the rigid structure of coalesced Crassostrea shells form new stable microhabitats. Regarding the Crassostreapatches, the topography and in particular a vertical orientation of Crassostrea shells may amplify alteration in physical properties. Within the matrix of vertically oriented surfaces the influence of solar radiation decreases with increasing depth. This may explain the higher abundance of C. maenas as well as the exclusive occurrence of anthozoans, M. palmata, H. imbricata and A. rubens in the Crassostrea-reef (Table 2).

The geometry of the Crassostrea shells offers various cryptic microhabitats most suitable for colonization by several vermicular organisms. Sediment gets caught up in the petalled margins caused by shell proliferation. Through the settlement of Crassostrea larvae on conspecifics, sessile barnacles on these individuals happen to be overgrown by the extent of shell accretion of the young oyster. Sediment gets washed into the remaining space between oyster shells and plates of dead barnacles. In these crevices, between coalesced oyster shells and in the petalled margins of the oyster shells, we found T. benedii, N. succinea, and Polydora-species (Fig. 10). This may have resulted in the high abundance of oligochaetes, the exclusive and marked occurrence of N. succinea and the shift in the composition of polychaetes to more opportunistic facultative filter feeders in the Crassostrea-patches.

In contrast to mainly horizontal surfaces which occur in *Mytilus*-patches, vertically oriented *Crass-ostrea* shells show complex patterns of current flow.



Fig. 10 a Polydora-species in sediment-filled crevices of the petalled shell margins of Crassostrea gigas b traces of Tubificoides benedii; two coalesced Crassostrea shells broken apart, honeycomb-like structures display the basal plates of overgrown barnacles

Soniat et al. (2004) reported that flow between vertically oriented shells was reduced to zero, whereas higher values were measured at the lateral extent of the shells and above the surface. Higher current velocities can favor food ingestion and growth of oysters as well as food ingestion and growth of associated filter feeders like *Polydora*-species (Lenihan et al. 1996). Also barnacles, anthozoans, hydrozoans and ascidians may benefit from these environmental conditions resulting in the enhanced colonization by sessile suspension feeding organisms in the *Crassostrea*-reef and, hence, may have resulted in a higher diversity of the epibiont community within this study.

Flow rates increase from bottom level towards the top of an oyster reef (Lenihan 1999). Accordingly, sedimentation near the reef base is favored by a strong reduction of hydrodynamic energy, whereas the upper parts of the shells remain free of sediment deposition due to high turbulence. *M. edulis* are more frequently affected by burial (Soniat et al. 2004). They are able to move back to the surface but attached organisms may suffer. The epifaunal reef community is continuously protected against sedimentation because of a permanently sediment-free upper shell surface in the *Crassostrea*-patch. This may have as well contributed to the richer epibenthos in *Crassostrea*-patches.

According to Asmus (1987), grazers and suspension feeders hold the highest share in total biomass of the associated macrofauna of intertidal *Mytilus*-beds. Predatory species appear to be less important. We found a similar distribution in *Mytilus*-patches. In contrast, we detected a shift in the distribution of sessile and vagile epifauna in the *Crassostrea*-reef. The strong increase in total biomass in Crassostreapatches was associated with the vagile epifauna which contributed to 50% to the biomass, predominantly provided by L. littorea and C. maenas. Thus, suspension feeders like barnacles and anemones were of minor importance regarding their biomass, whereas predators (e.g. C. maenas, H. imbricata, A. rubens) became more important. While providing refuge to intermediate predators, habitat complexity appears to reduce interference competition among predators (Grabowski and Powers 2004). The evolving Crassostrea-reefs may promote the establishment of communities characterized by organisms of higher trophic levels. Already Dittmann (1990) reported a variety of trophic groups co-occurring in a Mytilusbed as an attribute of structural heterogeneity of epibenthic bioherms.

#### Biodeposition

Walne (1972) reported higher clearance rates for *C. gigas* compared with equal sized *M. edulis*. Other studies revealed a higher clearance rate for *M. edulis* in springtime (Deslous-Paoli et al. 1987; May 2006; Wehrmann et al. 2006). However, the clearance rate and the corresponding biodeposit production depend on the size, respectively on the biomass of a bivalve. Biomass of both ecosystem engineers in the *Crass-ostrea*-patches was twice as high as it was in the *Mytilus*-patches. This implies a higher biodeposit production in *Crassostrea*-patches than in *Mytilus*-patches within this study.

The presence of biodeposits favors colonization by endobenthic organisms (Dittmann 1987). Kröncke (1996) showed that the amount of organic material in

sediments constitutes the main factor in structuring endobenthic macrofaunal communities. The high food supply to the sediment benefits organisms feeding on organic particles or organic-rich sediments from the surface (surface deposit feeders). Mixing of the sediment column, for example by bioturbation, results in an enrichment of even deeper sediment layers with organic matter. Thus, subsurface deposit feeders are also favored. Resuspension transports biodeposits into the adjacent sandflats of Mytilusbeds (Kröncke and Bergfeld 1996; Hild and Günther 1999). Compared to a Mytilus-bed, the superficial structure of a Crassostrea-reef increases bottom roughness and water turbulences (Reise 2002). Thus, more biodeposits could have been exported from Crassostrea-patches than from Mytilus-patches.

The organic enrichment of sediments by biodeposits causes an increase in bacteria density (Dittmann 1987; Villbrandt et al. 1999) altering the geochemical sedimentary environment. High oxygen consumption by bacterial degradation of organic material leads more and more to anoxic conditions and causes an increase in toxic H<sub>2</sub>S produced by sulphur bacteria even in upper sediment layers (Little-Gadow 1978; Kristensen 2000). Dittmann (1990) and Kröncke (1996) showed that sand flats are dominated by polychaetes, whereas the underlying sediments of Mytilus-patches are dominated by oligochaetes. Likewise, in this study the sand flat and the sediments in the open spaces of the bivalve bed were dominated by polychaetes whereas the stations covered by epibenthic bivalves were dominated by oligochaetes. T. benedii, the most common oligochaete of this study is known to be indicative for areas of extremely high organic pollution (Pearson and Rosenberg 1978; Dittmann 1987). The detoxification of sulfides allows T. benedii to live in environments characterized by high H<sub>2</sub>S content (Hartmann-Schröder 1996). The strong appearance of oligochaetes in Crassostrea-patches and in the open spaces between these patches indicates increasing oxygen depletion and H<sub>2</sub>S production resulting from maximum exploitation of biodeposits.

Several authors (Dittmann 1987; Kröncke 1996; Bergfeld 1999; Villbrandt et al. 1999) predict that polychaete composition is affected by the accumulation of biodeposits and related changes in sediment chemistry. Dittmann (1987) reported that *A. marioni*, *C. capitata* and *Microphthalmus* sp. indicate enrichment of sediments by biodeposits. *A. marioni* prefers sandy sediments rich in fine grained muddy material. Enhanced organic enrichment of the sediments, especially in the open spaces between the Crassostrea-patches, may be responsible for the dominance of A. marioni. Besides C. capitata, M. sczelkowii and S. armiger frequently occurred in the open spaces between the Crassostrea-patches, also indicating a high concentration of organic material. According to Dittmann (1987) polychaetes respond very sensitively to epibenthic bivalve coverage. In her field experiments the composition of polychaetes changed within one year at stations artificially covered by Mytilus. At the beginning of the experiment A. marioni and S. armiger were the most abundant species. The species composition after one year was dominated by C. capitata, whereas the abundance of A. marioni and P. elegans decreased markedly. The frequencies of polychaetes in Mytilus-patches of our study did not correspond to the distribution found by Dittmann (1987) after one year. We counted more individuals of A. marioni and P. elegans in Mytiluspatches than in the respective open spaces. However, C. capitata became more important.

The influence of the C. gigas coverage on the sediment chemistry and subsequently on the composition of the infauna in Crassostrea-patches seems to be more distinct than the influence of the M. edulis cover in the Mytilus-patches. All polychaetes were found in low abundances in the Crassostrea-patches. Even C. capitata was rare and P. elegans was absent, perhaps due to enhanced accumulation of organic material. A. marioni dominated in Crassostreapatches, even though with very low abundance. The internal fabric of the Crassostrea-patches, characterized by various microhabitats and a specific flow regime through vertical oriented shells (see section 'Habitat matrix'), may result in a highly variable small-scale distribution pattern of sediment types differing in geochemistry. This is supported by the high species number but low abundance of polychaetes found in Crassostrea-patches. Most common were Polydora-species, added values for P. ciliata and P. cornuta reached 43% of the total number of polychaetes, which we found predominantly in the petalled margins of Crassostrea shells. N. succinea was also common, exclusively found between coalesced oyster shells. This may imply a distribution of these species independent of the geochemical conditions of the underlying sediments and may more likely be controlled by the availability of suitable habitats associated with the structure and formation of the oyster shells in the *Crassostrea*-patches.

The sand flat was predominantly characterized by the absence of nemertines and the common occurrence of S. armiger, A. marina and amphipods of the genus Bathyporeia, all of them typical residents of sandflats (Dörjes 1978). The latter, along with all other identified infaunal amphipods, were found exclusively in the sand flat. The dominant species, however, was P. elegans, known to prefer sandy tidal flats (Bergfeld 1999). C. capitata, an indicator species of nutrient-rich sediments, was found in moderate abundance. The sediments in the sand flat 100 m outside the bivalve bed seem to be influenced by biodeposits produced within the bivalve bed. This is also supported by the occurrence of large individuals of C. edule and M. *balthica*, both typical representatives of mudflats with intermediate mud content.

# Outlook

According to Jones et al. (1994) ecosystem engineers are organisms that directly or indirectly modulate the availability of resources to other species. Major impact originates in high abundance of an ecosystem engineer influencing vast areas over a long period. The biodiversity of a habitat increases with its structural complexity (MacArther and MacArther 1961). Tsuchiya and Nishihira (1986) reported higher species richness and higher biodiversity with increasing structural complexity even within Mytilus-beds. The conceivable change from intertidal Mytilus-beds to Crassostrea-reefs in the Wadden Sea may promote the indigenous fauna. Long-term stability of a Crassostrea-reef and progressing habitat modulation as a consequence seems to be likely. This may lead to further increase in diversity.

Acknowledgments The study was carried out within the framework of the project "Bioinvasion" which is financially supported by the SNG Senckenbergische Naturforschende Gesellschaft and the Niedersächsische Wattenmeer Stiftung. Thanks are due to Torsten Janßen for helping during field work. We are grateful to Norbert Dankers, Frouke Fey and Karsten Reise for valuable remarks which considerably improved an earlier version of the manuscript. The paper benefits essentially from constructive comments of two anonymous reviewers. Thanks to Charles Ver Straeten (New York State Museum) for correcting the English.

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