Water salinity and benthic macrophyte communities are the key variables defining the distribution pattern of benthic faunal assemblages in the shallow water areas of the Gulf of Riga

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Abstract. The distribution of shallow water macrozoobenthic assemblages of the Gulf of Riga is analysed in relation to water current velocity, salinity, sediment type, ice disturbance, eutrophication, and benthic macrophytes. Salinity and benthic macrophyte communities described most of the variability in both the abundance and biomass structure of benthic invertebrates. In addition, surface chlorophyll $a$ also contributed to the variability of benthic invertebrate biomass. Benthic invertebrate assemblages in the vicinity of the Daugava and Pärnu rivers experience higher nutrient loading and were therefore characterized by a lower species number and higher biomass of benthic suspension feeders. Benthic invertebrate communities in other regions of the Gulf of Riga were typical for the mildly disturbed areas of the northern Baltic Sea. Compared to the situation in the 1960s, no substantial increase in the abundance and biomass values of macrozoobenthos was observed as could be expected from the basinwide trend of increasing eutrophication.

Key words: Baltic Sea, distribution, benthic invertebrates, benthic macrophytes.

INTRODUCTION

Contrary to the intertidal areas of the oceanic coasts (e.g. Terlizzi & Schiel, 2009), the large-scale patterns of benthic invertebrate communities in shallow water macrophyte habitats are not well studied in the Baltic Sea range. The majority of published papers focus on sedimentation and deeper areas (e.g. Laine et al., 2007; Kotta et al., 2007). When shallow water macrophyte communities are studied emphasis is only rarely placed on benthic macrophytes (Kautsky & Van der Maarel, 1990; Kautsky et al., 1999). This is quite surprising considering the specificity of the Baltic Sea ecosystem and the potential uniqueness of how environmental variables shape its local populations.

The Gulf of Riga is a relatively shallow and closed water basin in the eastern Baltic Sea characterized by a fairly high riverine input from its extensive drainage area. Therefore, the communities are composed of a specific mixture of freshwater, brackish-water, and marine species (Kotta et al., 2008). Although the nutrient loading into the Gulf of Riga is high and the basin is considered to be
among the most eutrophied areas in the Baltic Sea (Suursaar, 1995; Ojaveer, 1997; Kotta et al., 2008), the coastal areas of the Gulf of Riga support healthy macrophyte assemblages with the biomass of perennial species exceeding the values reported elsewhere in the Baltic Sea range (Kautsky et al., 1999).

Benthic invertebrates have been often used to assess the state of an aquatic environment because they are long-lived, sessile, and easy to collect and determine (e.g. Leppäkoski, 1975; HELCOM, 1993). The earliest quantitative surveys of benthic invertebrates in the Gulf of Riga were carried out in the 1950s (Shurin, 1953, 1960, 1961). These investigations were followed by the profound studies by Järvekülg (1961, 1962, 1979). Both series of investigations concentrated mainly on the macrozoobenthos inhabiting the soft substrate, that is the deeper part of the basin. Since then no concise work about the benthic invertebrates of the whole gulf has been published and most papers deal with the impact of the Pärnu or Daugava rivers (e.g. Lagzdīns, 1975; Kotta & Kotta, 1995; Kotta et al., 2009b). Hence, the extensive coastal areas of the Gulf of Riga are almost overlooked in terms of benthic invertebrates and we are still not aware how sensitive these communities are to changes in human induced pressures and climate conditions.

The objective of this investigation is to provide the first quantitative data about the distribution of benthic invertebrates in the shallow water range of the Gulf of Riga. As the coastal zone controls exchange, storage, and transformation processes of materials, knowledge about the structure and functions of benthic assemblages in this zone is fundamental for understanding the functioning of the system as a whole. Scuba diving made it possible to describe quantitatively the zoobenthic assemblages on hard substrate, but also contributed to better distinction between true sediment- and macrophyte-dwelling species. The distribution of benthic invertebrates was related to environmental parameters such as water current velocity, salinity, sediment type, ice disturbance, eutrophication, and benthic macrophytes. Most studies dealing with the shallow water benthic fauna have neglected benthic macrophytes. Finally, the results will be compared with the historical data in the area.

**MATERIAL AND METHODS**

Sampling was performed in August 1995 and 1996 on 10 transects located around the whole coastline of the Gulf of Riga (Fig. 1). The northern part of the gulf is characterized by a wide coastal zone with a diverse bottom topography and macrophyte community, and the southern side by a narrower sandy coastline and hard bottom substrate in the deeper parts of the littoral zone.

The samples were collected with a Tvärminne sampler (Kangas, 1972) on sand and gravel bottoms and a suction sampler (Hiscock & Hoare, 1973) on hard bottom types like boulders and solid limestone floor. The sampling area of both samplers is 314 cm². Three replicate samples were generally taken from 10 transects each containing 3 sites (a total of 94 samples), of which 19 were
unvegetated (56 samples), 6 sites had vascular plants (18 samples), and 7 sites algal (filamentous algae, *Fucus vesiculosus* L. or *Furcellaria lumbricalis* J. V. Lamour) vegetation (20 samples). The sampling depth varied from 0.1 to 6.0 m. Samples below 0.5 m were taken by SCUBA diving. Sampling sites were chosen to cover all possible sediment types and macrophytobenthic communities. In each site the type of substrate (stone, boulder, gravel, sand, silt, clay, or mixed), depth, and dominant plant species were recorded.
Material was sieved through a net of 0.25 mm mesh size and preserved in 4% buffered seawater formalin solution. In the laboratory animals were counted under binocular microscope. Dry weights were obtained (to the nearest 0.1 mg) after drying the material at 70°C for 60 hours. Molluscs were weighed with shells.

The values of water temperature, salinity, and water current velocity were obtained from the results of hydrodynamic model calculations from the early April 1995 to the early August 1996. The calculations were based on the COHERENS model, which is a primitive equation ocean circulation model. The model was formulated with spherical coordinates on a 1' × 1' minute horizontal grid and 30 vertical sigma layers. The model was forced with hourly meteorological fields of 2 m air temperature, wind speed, wind stress vector, cloud cover, and relative humidity. The meteorological fields were obtained from an operational atmospheric model. The model was validated against water level, temperature, salinity, and water velocity measurements from the study area (Bendtsen et al., 2009).

The Finnish Meteorological Institute provided the long-term averages of ice cover and thickness over the study area. Ice cover and thickness were produced on daily basis at a nominal resolution of 500 m and were based on the most recent available ice chart and a synthetic aperture radar (SAR) image. The ice regions in the ice charts were updated according to a SAR segmentation, and new ice parameter values were assigned to each SAR segment based on the SAR back-scattering and the ice thickness range at that location.

As a proxy of local eutrophication in the study area, the long-term average and maximum of surface chlorophyll \( a \) values obtained from SeaWiFS chlorophyll composites were used. Each image had a spatial resolution of 11 km × 11 km. Cloud gaps were filled by an interpolation procedure over a maximum distance of 100 km.

The contribution of site, sediment type, and dominant phytobenthic species to the abundance and biomass of macrozoobenthos was tested by one-way ANOVA after checking for normality of the data (Kolmogorov–Smirnov test for goodness of fit) and homogeneity of variance (Bartlett’s and Hartley’s tests) (Sokal & Rohlf, 1981).

Multivariate data analyses were performed by the statistical program ‘PRIMER’ version 6.1.5 (Clarke & Gorley, 2006). In order to increase the signal-to-noise ratio in the multivariate analysis an average of three replicates in each site was used to construct similarity matrices. A zero-adjusted Bray–Curtis similarity measure was used. Data was log-transformed to reduce the contribution of abundant species to similarity measure. Ordination was made by nonmetric multi-dimensional scaling (MDS) (Clarke & Green, 1988). Environmental (abiotic and plant) variables were normalized and then the Euclidean distance matrix was calculated. BEST analysis (BIOENV procedure) was used to relate the similarity matrices of environmental variables to the similarity matrices of the abundance and biomass of benthic invertebrates. A global BEST match permutation test was run to examine the statistical significance of the observed relationships between environmental variables and biotic patterns. The statistical differences in invertebrate
assemblages between defined groups were obtained by the ANOSIM permutation test (Clarke & Green, 1988; Clarke, 1993). The contribution of each species to dissimilarities was investigated using the similarities percentages procedure (SIMPER) (Clarke, 1993).

RESULTS

The total abundance and biomass of macrozoobenthos varied largely between the transects of the study area (Table 1, Fig. 1). The highest mean abundance was found in the outermost part of the Gulf of Riga (transect 4; 5838 ind m\(^{-2}\)) and the lowest on the adjacent area across the Irbe Strait (transect 9; 71 ind m\(^{-2}\)). The highest mean biomass was recorded in the easternmost part of the gulf (transect 6; 71 g dw m\(^{-2}\)) and the lowest adjacent to the Daugava River mouth and in the coastal zone of Ruhnu Island (transects 7 and 10, respectively; 7 g dw m\(^{-2}\) in both regions). The distribution pattern of abundance and biomass values along the depth gradient was not consistent. Depending on the location of the transect, denser populations were observed in the shallower areas (transects 4 and 5), at moderate depths (transects 1, 2, 7, 8, and 9), or in the deeper areas (transects 3, 6, and 10).

BIOENV analysis showed that salinity and vegetation type were the environmental variables predicting best the abundance of benthic invertebrate communities (Spearman $\rho_{\text{salinity+vegetation}} = 0.332$; separate contributions of environmental variables were $\rho_{\text{salinity}} = 0.243$ and $\rho_{\text{vegetation}} = 0.178$; $p = 0.04$). As for biomass, chlorophyll $a$ and sediment type also contributed to the model (Spearman $\rho_{\text{salinity+vegetation+sediment+chlorophyll}} = 0.326$; separate contributions of environmental variables were $\rho_{\text{salinity}} = 0.216$, $\rho_{\text{vegetation}} = 0.224$, $\rho_{\text{chlorophyll}} = 0.110$, and $\rho_{\text{sediment}} = 0.265$; $p = 0.02$).

The univariate measures of benthic invertebrate abundance and biomass also varied along the salinity gradient and among different bottom and vegetation types (Fig. 2). Vascular plant communities were characterized by a higher abundance of insects and gastropods of freshwater origin (*Lymnaea* sp., *Bithynia*...)

Table 1. One-way ANOVAs of the effects of sediment and vegetation types on the abundance and biomass values of different trophic groups of benthic invertebrates. Significant differences ($p < 0.05$) are in bold.

<table>
<thead>
<tr>
<th>Model</th>
<th>Total</th>
<th>Suspension Feeder</th>
<th>Herbivore</th>
<th>Carnivore</th>
<th>Detritivore</th>
<th>Omnivore</th>
</tr>
</thead>
<tbody>
<tr>
<td>Sediment</td>
<td></td>
<td></td>
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</tr>
<tr>
<td>Abundance</td>
<td>$&lt;0.001$</td>
<td>0.069</td>
<td>$&lt;0.001$</td>
<td>$&lt;0.001$</td>
<td>$&lt;0.001$</td>
<td>$&lt;0.001$</td>
</tr>
<tr>
<td>Biomass</td>
<td>0.573</td>
<td>0.809</td>
<td>$&lt;0.001$</td>
<td>0.988</td>
<td>0.209</td>
<td>$&lt;0.001$</td>
</tr>
<tr>
<td>Vegetation</td>
<td></td>
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</tr>
<tr>
<td>Abundance</td>
<td>$0.025$</td>
<td>0.001</td>
<td>$&lt;0.001$</td>
<td>0.202</td>
<td>0.660</td>
<td>$&lt;0.001$</td>
</tr>
<tr>
<td>Biomass</td>
<td>$&lt;0.001$</td>
<td>$&lt;0.001$</td>
<td>$&lt;0.001$</td>
<td>$&lt;0.001$</td>
<td>0.170</td>
<td>$&lt;0.001$</td>
</tr>
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</table>
tentaculata (L.)), while biomass was dominated by burrowing mussels (*Macoma balthica* L. and *Cerastoderma glaucum* Bruguière). Unvegetated bottom types were dominated by crustaceans (mainly *Gammarus* spp. and *Corophium volutator* (Pallas)), followed by polychaetes (*Hediste diversicolor* (O. F. Müller) and *Marenzelleria neglecta* Sikorski & Bick) and oligochaetes. The abundance was highest on shallow (<1 m) sandy bottoms in the north-eastern part of the gulf (3700 ind m⁻²) while the shallow areas of the south-eastern part were very poor in macrofauna (only 160 ind m⁻²). The biomass was mainly formed by burrowing mussels, *M. balthica* and *C. glaucum* in shallow areas and *M. balthica* and *Mya arenaria* L. in deeper and coarser bottoms. The biomass was highest in deep (5–6 m) gravel bottoms (76 g dw m⁻²). Hard bottom algal communities were composed mainly of *Theodoxus fluviatilis* (L.), *Jaera albifrons* Leach, *Idotea*
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*chelipes* (Slabber), *I. baltica* (Pallas), *Gammarus* spp., and the sedentary mussels *Dreissena polymorpha* (Pallas) and *Mytilus trossulus* Gould. The last two mussel species were found in the same samples at four sites.

Five different trophic groups of macrozoobenthos were distinguished in the study range: omnivores, detrivores, carnivores, herbivores, and suspension feeders (Fig. 3). Detrivores (mainly *M. balthica*) contributed significantly to the total abundance in the northern part of the gulf. The proportion of herbivores was higher in the eastern part of the gulf (*T. fluviatilis* in the south-eastern part and *Hydrobia* spp. and *Lymnaea* spp. in the north-eastern part of the gulf). As to the biomass, detrivores (*M. balthica*) were still the dominant feeding type in the northern part of the gulf, with a significant proportion of suspension feeders (*D. polymorpha*) in the vicinity of Pärnu Bay. The share of macrozoobenthos of different feeding types was similar in the southern part of the gulf. The proportion of suspension feeders was highest on transects 6 and 8. Transect 6 was unique representing either the communities of northern type (by biomass) or southern type (by abundance).

**Fig. 3.** Proportion of different trophic groups in terms of abundance and biomass at different transects.
Salinity explained a large part of the observed variability of the benthic invertebrate communities in the Gulf of Riga (Fig. 4). Of the three groups of species (marine, brackish-water, and freshwater) brackish and freshwater species had a higher proportion in the southernmost (transects 7 and 8) and the northeastern part of the gulf (transect 2). These transects are influenced by the inflow of either the Daugava or the Pärnu River.

According to one-way ANOVA (Table 1), sediment type affected the total abundance values of macrozoobenthos whereas dominant plant species was significant in describing both the total abundance and biomass values of macrozoobenthos. Higher abundances were found on stone, sandy gravel, and gravelly clay bottoms, lower on clayey sand and sand bottoms. The lowest abundance and biomass were found on *Zannichellia palustris* L. community, the highest on *F. vesiculosus* and *Cladophora rupestris* Kütz. community. The density of herbivores and omnivores was significantly higher on stone bottoms, that of detrivores on sandy gravel and gravelly clay bottoms. Suspension feeders preferred communities dominated by *F. vesiculosus*, *F. lumbricalis*, and *Myriophyllum spicatum* L., herbivores the *F. vesiculosus* community, carnivores *F. lumbricalis* and *M. spicatum*

![ABUNDANCE](image1.png)

![BIOMASS](image2.png)

**Fig. 4.** Proportion of marine, brackish-water, and freshwater species in terms of abundance and biomass at different transects.
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communities, and omnivores *C. rupestris* and *F. vesiculosus* communities. The abundance and biomass of detrivores were independent of the type of plant communities around the sampling area.

Ordination of log-transformed abundance and biomass data of macrozoobenthos did not produce very distinguished groups (Fig. 5). Considering the biomass values, transect 2 was clearly separated from transects 8, 9, and 10 (ANOSIM, *p* < 0.05), transect 3 from transects 8 and 9 (*p* = 0.03), and transect 4 from transect 7 (*p* = 0.03). The most of dissimilarities were due to *M. balthica*, *C. glaucum*, *Bithynia tentaculata* L., *H. diversicolor*, *D. polymorpha*, and *T. fluviatilis*. The first three species had a higher biomass in the northern part of the gulf. On the contrary, *T. fluviatilis* was more common at the southern coasts of the gulf. At the north-eastern and southern coasts *D. polymorpha* thrived whereas *H. diversicolor* occurred with a higher biomass in the western side of the gulf. The ordination of macrozoobenthos by abundance shows the transects to be even more similar to one another.

Surprisingly, there was only a weak relationship between bottom types and the structure of macrozoobenthic communities. Significant differences were observed comparing very different substrate types such as stone bottoms with gravel–sand bottoms (ANOSIM, *p* < 0.01). Stone bottoms were characterized by the highest density of *D. polymorpha*, *T. fluviatilis*, Turbellaria, *Gammarus oceanicus* Segerstråle, *G. salinus* Spooner, *G. zaddachi* Sexton et Spooner, and *C. glaucum*. The highest abundance of *M. neglecta* and Oligochaeta was observed on sandy gravel bottoms.

There was a significant difference between the areas devoid of vegetation and the *Z. palustris* community (calculations based on macrozoobenthic biomass data, ANOSIM, *p* = 0.05) and between *F. vesiculosus* and *Potamogeton pectinatus* L.

![Fig. 5. MDS ordination of sampling sites based on benthic invertebrate log-transformed abundance (A) and biomass (B).](image-url)
communities (abundance data, \( p = 0.03 \)). In the former case, \( T. \) fluvialitis, \( M. \) balthica, and \( H. \) diversicolor contributed to the differences between these two community types. The last two species had a higher biomass on unvegetated areas and \( T. \) fluvialitis in the \( Z. \) palustris community. Comparison of \( F. \) vesiculosus and \( P. \) pectinatus communities showed that \( J. \) albifrons, \( G. \) spp., \( T. \) fluvialitis, \( I. \) neglecta, and \( M. \) balthica had higher abundance in the \( Fucus \) belt whereas only Chironomidae of the \( P. \) pectinatus community outnumbered those of the \( F. \) vesiculosus community.

**DISCUSSION**

The study showed that the large-scale distribution of benthic invertebrate communities in the Gulf of Riga was most likely driven by changes in salinity and benthic macrophyte communities. The benthic biomass was additionally affected by water quality. Although the role of salinity in determining the large-scale invertebrate patterns was demonstrated earlier in the whole Baltic Sea range (Bonsdorff & Pearson, 1999), the relative contribution of macrovegetation and eutrophication has not been assessed. Locally, however, the role of macrophytes in providing structured habitats for benthic invertebrates may have been well established (e.g. Kotta & Orav, 2001). Besides providing the habitat, benthic macrophytes fulfil multiple ecological functions, including improvement of sediment stability, food regime, and water quality as well as protection from wave disturbance (e.g. Orav-Kotta & Kotta, 2004; Torn et al., 2010).

Our results clearly demonstrated that in the Gulf of Riga the abundance of benthic invertebrates was fairly insensitive to variability in eutrophication (i.e. long-term average of water chlorophyll \( a \) content) whereas their biomass was strongly related to changes in water quality. Specifically, strong gradients were observed in water chlorophyll \( a \) along the major river mouths; moreover, such gradients matched with the biomass patterns of the suspension-feeding bivalves. This advocates the use of invertebrate biomass rather than abundance for the assessment of the status of the Gulf of Riga coastal areas in the frame of The European Union Water Framework Directive (Kotta et al., 2012).

The abundance values of macrozoobenthos found in this study were of the same magnitude as those found in other coastal areas in the northern Baltic Sea (e.g. Elmgren & Ganning, 1974; Skult, 1977; Orav et al., 2000). However, the values of biomass found in the Gulf of Riga were much lower than those in the Baltic Proper. This corresponds to the scarcer \( M. \) trossulus population due to lower salinity values in the Gulf of Riga. The biomass of the brackish-water \( D. \) polymorpha was not high enough to compensate for this.

The proportion of suspension feeders was fairly low in the vicinity of the mouth of the Daugava River, the main pollution source of the Gulf of Riga (transect 7). At the same time the abundance and biomass of suspension feeders (mainly \( D. \) polymorpha) were very high on both sides of the Daugava River mouth (transects 6 and 8). A lack of the suspension feeders in the vicinity of the Daugava
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River mouth is by no means related to the water quality but reflects the prevalence of soft substrate in the area, not suitable for any suspension-feeding mussels, including *D. polymorpha*, inhabiting the south-eastern Gulf of Riga.

Higher biomass of detrivores in the northern part of the gulf is related to different coastal morphology. Due to the wider coastal zone and lower input of wave energy, sedimentation is more intensive in the northern part as compared to the southern part of the gulf, hence favouring the development of dense populations of e.g. *M. balthica*.

Comparison of the proportions of benthic invertebrates of different feeding type in unvegetated and vegetated areas revealed that the latter were characterized by a much smaller proportion of suspension feeders in the vicinity of the Daugava and Pärnu river mouths but not in the coastal sea of Saaremaa Island. In general, macrophytes and mussels are expected to compete for space and an inverse relationship between these trophic guilds is expected. Specifically, macroalgae are known to outcompete benthic suspension feeders at shallow depths, and lush macrophyte communities are therefore often characterized by low densities of suspension feeders (Janke, 2006). Although this may be a plausible mechanism in the vicinity of river mouths where the fast-growing phanerogams such as *Potamogeton* spp. prevail, in the northern Gulf of Riga with the presence of small-sized macroalgae the mussels are likely facilitating algal growth in this dynamic and physically disturbed habitat (Kotta et al., 2009a).

Shallow-water benthic invertebrate communities are expected to face strong disturbances by ice and waves. It is therefore surprising that the ice effects were negligible in all statistical models applied. It is plausible, though, that strong seasonality of the shallow water ecosystem of the Gulf of Riga is behind the lack of a such relationship (Kotta et al., 2008). In the study area virtually the whole benthic invertebrate biomass is produced during the spring–summer season and thus communities respond to environmental variables in this period (i.e. water temperature, salinity, nutrients) rather than to any ice-mediated effects.

According to Järvekülg (1961, 1979), *M. balthica* dominated the whole sublittoral zone of the Gulf of Riga in the 1960s. Nowadays this is also true and only locally *M. neglecta*, *H. diversicolor*, *D. polymorpha*, or *T. fluviatilis* may take the leading role. Unfortunately, due to the differences in methods, it is hard to compare how the species diversity has changed. Nevertheless, earlier studies demonstrate that the present-day communities are more uniform (shown by lower dissimilarities/distances between the stations) compared to the past communities (Kotta et al., 2012). Besides, the decline of phytophilous species in the northern part of the gulf was documented by Kotta & Kotta (1997). In the 1960s the average abundance and biomass values were respectively 840 ind m⁻² and 3 g ww m⁻² at 0.1–0.4 m depth and 2210 ind m⁻² and 62 g ww m⁻² at 0.5–9 m (Järvekülg, 1979). We measured 2640 ind m⁻² and 18 g dw m⁻² in the shallower areas and 2475 ind m⁻² and 35 g dw m⁻² in the deeper areas, respectively. The average ratio of wet weight to dry weight in the study area was about 2.8. Figure 6 compares the average abundance and biomass data of benthic
invertebrates at different sediment types between the 1960s and this study. As the methods used by Järvekülg underestimated the abundance and biomass of macrozoobenthos on stone and gravel bottoms (especially at 0.1–2 m), the magnitude of the actual changes was smaller than presented in the figure. Hence, it is likely that the biomass and abundance of benthic invertebrates have not significantly increased in the shallow water coastal range of the Gulf of Riga during the last 30 years.

CONCLUSIONS

On the basis of shallow water benthic invertebrate assemblages, the Gulf of Riga may be broadly divided into southern and northern areas. Among abiotic variables salinity, macrophyte communities, and nutrient input by rivers explain a significant proportion of variation in the abundance and biomass of macrozoobenthos. Despite the heavy nutrient loading into the Gulf of Riga the biomass and abundance of benthic invertebrates have not substantially increased during the last 30 years. The overall condition of benthic invertebrate assemblages seems to be good compared to the similar areas in the Northern Baltic Sea.
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