

## **Final report of the ICES Science fund project "Holistic approach to analyse benthic fauna communities on the whole Baltic Sea (BSW)"**

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### **Background**

The focus of the initiative is the identification, description and mapping of major benthic macrofauna communities, its distribution and functional features on the Baltic Sea wide scale. A number of works were previously published, describing and mapping the benthic macrofauna communities in different sub-basins of the Baltic Sea. Nevertheless, no synthesis or joint data analyses have been done for the entire Baltic Sea region in the recent time.

### **Aims**

The project aims to perform the inventory of the recent benthic macrofaunal communities in the entire Baltic Sea, thereby adding to the investigation of living resources in the ICES region. This development can serve as an evidence based advice for assessment of ecosystem status, be useful for environmental scientists to account for benthic macrofauna in implementation of ecosystem approach and helpful for decision makers in marine spatial planning and fishery management.

This work will set a baseline for future studies for better recognition of links between benthic fauna and ecosystem functioning.

Sampling efforts from all countries around the Baltic Sea are aggregated, harmonized and analyzed to extract ecologically significant groups shaped by pronounced environmental heterogeneity of the Baltic Sea.

Present report describes the results of analysis to delineate 10 major communities inhabiting the Baltic Sea. This is an arbitrary level of classification chosen to provide an overall global picture. The work will continue with further refinement of presented results and description of finer community units in exemplary regions.

### **Workshop**

The two days' workshop devoted to BSW Baltic Sea (BS) benthic fauna communities took place 27-28 May 2014. It was hosted by the IOW in Warnemünde, Germany. There were 9 attendees representing 7 institutes from Germany, Denmark, Finland, Lithuania, Estonia, Latvia and Poland.

The workshop was aimed to finalize the procedure of joining together the national datasets and summarize the experiences of participants in the community analysis. Following methodological issues were addressed:

- It was agreed that a global picture is desired on the cost of details: therefore inner waters, lagoons and small enclosed bays should be excluded from the data analysis due to high diversity and variability as we are aiming at defining and mapping the distribution of BS "regular/widespread/representative" communities; alternatively they can be treated as separate class of "diverse/patchy/variable" community.
- The relevance ("life span") of the map was discussed – temporal validity is limited by regional variability of benthic communities.

- The reasonable approach for spatially explicit mapping based on point data is to model key species distributions separately and derive communities by overlapping results.
- New round of data inventory, including the submission of all available biomass data (for conversion of wet weight to AFDW regional conversion factors can be used), was considered necessary during the May workshop and completed by the end of August 2014. To increase the coverage and resolution of data Alexey Maximov (Zoological Institute of Russian Academy of Science) was contacted and has contributed data for 25 stations from the eastern part of the Gulf of Finland. These data was included in the dataset.
- The range of sizes of the gridding cell to be considered for the final product (including estimation of variability within cell): EU grid, 1 km, 1nm, 2 km (HELCOM), 5 km.
- If community classification is based on relative dominance of key species, differences in sampling gear, sampling area, number of replicate in combined data set are acceptable, but for absolute abundance/biomass based community classification methodological differences can lead to comparability problems and data should be treated with caution for application of predictive modelling techniques.
- Distribution of oxygen minimum zone can be used in order to map regions with no macrofauna. This is a problematic issue since data from such sampling events/locations rarely enter the databases on benthic macrofauna due to their structure specifications and are therefore poorly documented.
- There are also community changes driven by biological pressure e.g. invasive species whose distribution dynamics can rapidly lead to significant community shifts.
- Taxonomic inconsistencies were discussed: e.g. taxonomic units higher than species level should be removed when one or several species belonging to those units already is present from the same sampling event/spatial cell, all alternatively all aggregated to genus level (as done below).

## Methods

The data on the distribution of benthic macrofauna is collected together in one GIS database to adequately cover the whole Baltic Sea basin. Most of the data was collected with van Veen grab, mostly 2 to 3 replicates per sampling event, sieved on a 1 mm screen, preserved in 4 % buffered formaldehyde-seawater solution and identified in the laboratory to the lowest taxonomic level possible. More details including specifications and the area covered by each laboratory is provided below and in Appendix 1. All species abundance and biomass data was recalculated to the area of square meters. For biomass data conversion factors available at the IOW were used.

Baltic Sea can be viewed as a gradient system, where species richness increase with salinity towards the southwest and community composition changes, also responding to differences in substrates (generally dominated by fine sand and mud) and oxygen conditions (with seasonal and even permanent oxygen depletion occurring in deeper parts). Therefore, characteristic (dominant) species were first identified separately for each HELCOM sub-basin, accounting for their frequency and relative abundance at location, to extract the list of species feasible for the overall community analysis. Distinct benthic assemblages discriminated by particular species are defined, related to different spatial subarea and characterized by a certain variability of environmental parameters by means of various statistical methods (hierarchical clustering, indicator value, MDS).

As a starting point all available infaunal data was pooled together. For analyses we have used a regular 5 km grid (matching with the EU 10 km grid). Shallow lagoon regions were omitted not to blur the analysis due to high variability of infauna. Average abundance for taxonomic units was calculated accounting for all sampling events within one grid cell.

PRIMER v6 program package (Clarke and Warwick, 2001) was used to perform cluster analysis (group-average linking) and multidimensional scaling for abundance data to reveal similarities between stations. The cluster

analysis on a large number of species including patchy occurring rare species may tend to confuse the picture. The number of species was reduced, retaining only species ranking for up to 90% of cumulative abundance per each sampling event for cluster analysis. Some characteristic Baltic Sea taxa were considered at genus or even higher taxonomic level (*Diastylis*, *Harmothoe*, *Exogone*, *Hidrobidae*, *Idotea*, *Nephtys*, *Ophiura*, *Phoronis*, *Polydora*, *Diptera*) to overcome the taxonomic inconsistencies and as the global picture of characteristic communities distribution is of interest at this stage of the analysis. For list of taxa considered see Appendix 2. First abundance data was subject to fourth-root transformation. Similarities were calculated using the Bray-Curtis coefficient with a virtual dummy variable being 1 for all objects. Similarity percentage analysis (SIMPER) and Indicator species analysis (De Cáceres and Legendre, 2009) were used to identify the main taxa that were responsible for differences in community structure.

## Results

*Structure of the dataset.* The gathered and harmonised dataset is comprising over thousand taxa at over 7 thousand locations (17 thousand visit events) sampled in years 2000-2013 for species abundance and partly for biomass (Figures 1-2).

Overall 1176 taxonomic units (accepted in World register of Marine Species WoRMS) are included in the final joint dataset (774, i.e. 53% are unique to individual dataset). Harmonization to the lowest taxonomic level possible was done in order to reduce dimensionality for later statistical processing. Biomass data is limited: fresh weight is only available for 2943 (39%) locations, ash free dry weight - for 2241 (29.7%).

*Resent spatial distribution of characteristic macrofaunal communities.*

Based on cluster analysis of abundance data 10 major communities inhabiting the Baltic Sea were identified (cut off at 23%, clusters with minimum number of 10 cells considered as major communities, see the legend of Fig. 3 for characteristic species). Their distribution is shown in Fig. 4. This is an arbitrary level of classification chosen to provide an overall global picture. Results of Indicator species and SIMPER analyses revealed species primarily contributing to observed pattern and helped determine if the group is distinctly different from other. The outcome was also confirmed by expert knowledge. Characteristics of major communities are listed in Table 1.

Community 1 is the main type in the whole northern area, which isn't surprising when looking at the characteristic species of it (*Monoporeia affinis*, *Marenzelleria*, *Macoma balthica*). The Finnish samples are mainly from muddy sediments so that also explains a bit. There are some yellow spots of community 2 as well, those are the sandy areas with characteristic species *Hydrobiidae*, *Pygospio elegans*, *Cerastoderma glaucum*. This community is mainly found in sandy habitats in the south-western Baltic Sea. The red spots, group 4, in Gulf of Finland and central Baltic down to Arkona Basin, must be seen as "when not anoxic" communities, as those areas are often anoxic and the motile *Bylgides sarsi* is the one to arrive first when conditions are better. Communities in the southern part are much more diverse. Bay of Mecklenburg in shallower areas is mostly inhabited by sandy community 2 and community 3 (with 3 *Diastylis*, *Corbula gibba*, *Dipolydora quadrilobata*, *Arctica islandica*, *Aricidea suecica*, *Abra alba*) in the deeper part. Communities 5 (characterised by *Amphiura*, *Abra nitida*, *Galathowenia oculata*, *Ennucula tenuis*, *Thyasira flexuosa*, *Nucula nitidosa*, *Diplocirrus glaucus*), 7 (with *Phoronis*, *Tellina fabula*, *Thracia phaseolina*, *Ophelia borealis*, *Spiophanes bombyx*, *Branchiostoma lanceolatum*, *Spio arndti*) and 8 (*Angulus tenuis*, *Ensis directus*, *Haustorius arenarius*, *Lamprops fasciatus*) are mostly found in Kattegat region. Community 9 (with *Lagis koreni*, *Cerastoderma edule*, *Polydora*, *Halicryptus spinulosus*) is typical for Flensburg Fjord, whereas community 10 (characterised by *Echinocyamus pusillus*, *Harmothoe*, *Bittium reticulatum*, *Oligochaeta*, *Alitta virens*, *Turritella communis*, *Asterias rubens*) is nearly restricted to the Great Belt region. Results of SIMPER analysis can be found in Appendix 3.

Figure 1. Sampling station in the joint database and data providers

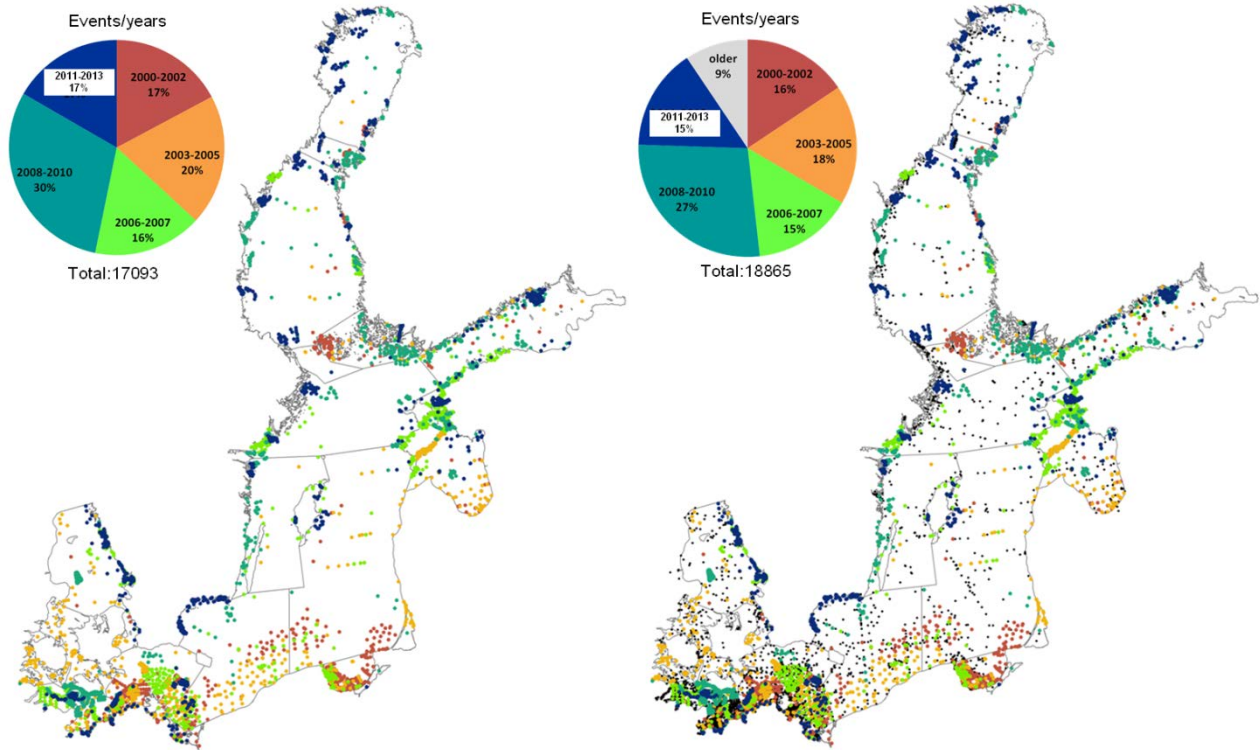


Figure 2. Data from different years

Examples of distribution maps for 5 exemplary bivalve species dominating abundance and biomass are given in Appendix 4.

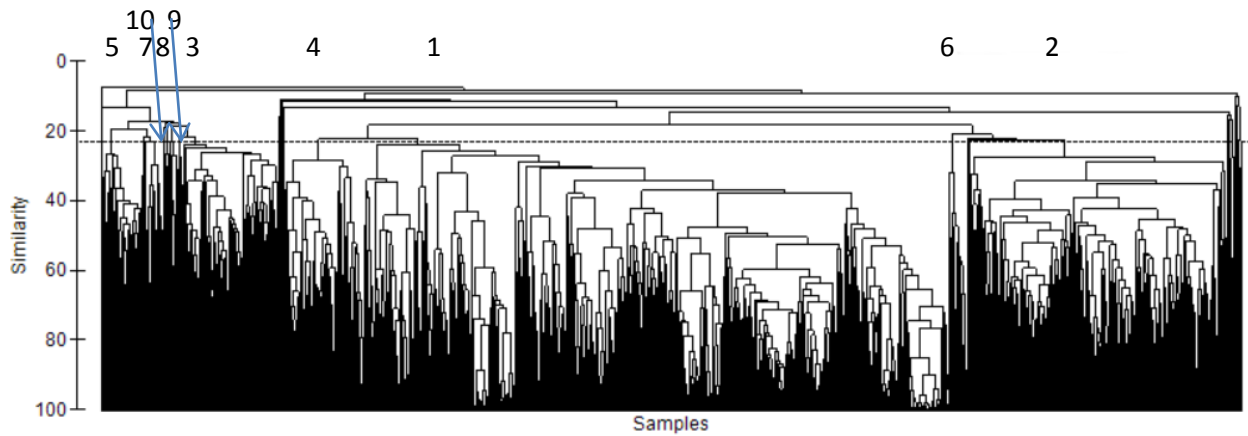
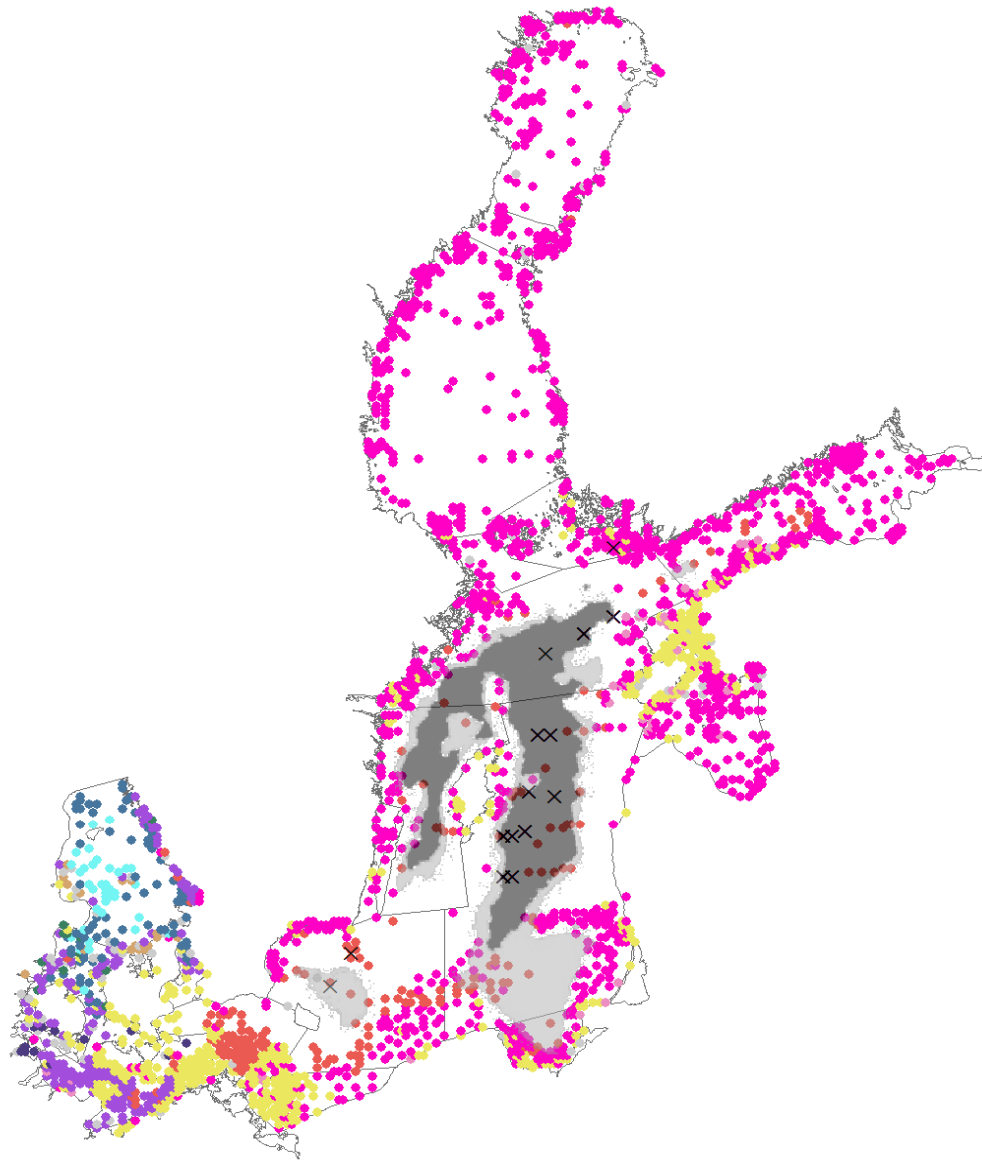


Figure 3. Dendrogram resulting from cluster analysis of benthic samples based on species abundance data. The horizontal dotted line represents the cut off which was at 23% similarity.

Table 1. Characteristics of major communities

Community	N cells	N visits	N species/ station	Mean abundance/ station (ind/ m <sup>2</sup> )	Mean biomass/ station (g/m <sup>2</sup> )	Depth	Av. salinity (min-max)	Sediment class (BALANCE)
1	1415	9977	5	1345	83	5-65	6.1 (0.9-19.4)	mud to sand
2	557	3827	14	4797	193	<30	9.5 (0.1-27.7)	sand
3	208	1434	38	3288	358	10-25	20.7 (13.5-30.8)	sand
4	205	855	7	485	54	20-120	12 (2.4-24.9)	mud
5	84	600	44	1903	204	3-40	28.3 (19.1-34.7)	mud
6	39	77	7	2986	20	2-70	7.6 (4.1-21.4)	hard bottom complex
7	38	127	29	1508	466	8-40	30 (25.4-34.1)	sand
8	17	22	21	1542	44	<25	26.4 (20.1-32.8)	sand
9	12	45	11	1385	41	<20	18.9 (13-22.3)	sand
10	11	12	29	1719	16	2-20	24.2 (19.4-28.1)	gravelly
other	70							
No fauna	50						11.3 (5.9-15.4)	mud



### Community

- 1 *Monoporeia.affinis*, *Marenzelleria*, *Macoma.balthica*
- 2 *Hydrobiidae*, *Pygospio.elegans*, *Cerastoderma.glaucum*
- 3 *Diastylis*, *Corbula.gibba*, *Dipolydora.quadrilobata*, *Arctica.islandica*, *Aricidea.suecica*, *Abra.alba*
- 4 *Bylgides.sarsi*, *Pontoporeia.femorata*
- 5 *Amphiura*, *Abra.nitida*, *Galathowenia.oculata*, *Ennucula.tenuis*, *Thyasira.flexuosa*, *Nucula.nitidosa*, *Diplocirrus.glaucus*
- 6 *Mytilus*, *Amphibalanus.improvisus*
- 7 *Phoronis*, *Tellina.fabula*, *Thracia.phaseolina*, *Ophelia.borealis*, *Spiophanes.bombyx*, *Branchiostoma.lanceolatum*, *Spio.arndti*
- 8 *Angulus.tenuis*, *Ensis.directus*, *Haustorius.arenarius*, *Lamprops.fasciatus*
- 9 *Lagis.koreni*, *Cerastoderma.edule*, *Polydora*, *Halicryptus.spinulosus*
- 10 *Echinocyamus.pusillus*, *Harmothoe*, *Bittium.reticulatum*, *Oligochaeta*, *Alitta.virens*, *Turritella.communis*, *Asterias.rubens*
- other

Figure 4. 10 major communities with characteristic species according to hierarchical clustering, SIMPER and Indicator species analysis. Crosses indicate sampled cells with no benthic infauna recorded (mainly dataset PL).

Light gray and dark gray areas mask out the deep water hypoxic and anoxic oxygen conditions, correspondingly (winters 2014 and 2015, after Aranda's monitoring cruise, SYKE) – when oxygen depletion takes place these areas are usually uninhabited by macrofauna.

Among the major outcomes of the project is the gathered and harmonised dataset that can be used as a baseline for multiple analyses of benthic fauna distribution and features. Data harmonisation is a continuous iterative process and often is the most time consuming step, as in this study.

### **Outlook and follow up**

Presented results are the big global picture, and of course when going more to detail more groups and species would appear important. The work will continue with further refinement of presented results and description of finer community units in exemplary regions.

As an outlook different methodological techniques/possibilities are considered for development of an appropriate approach for spatially explicit delineation of communities. Modelling the abundance of all possible taxonomic groups using Random Forest technique followed by overlay of results using measured and expected averaged abundance and biomass dominance is found to be most promising for delineation of communities. At the moment available environmental predictors are those available from IOW database, HELCOM Baltic Sea Data and Map service and EUSeaMap, including relevant physical and ecological features such as bathymetry, salinity, substrate type data, eutrophication related variables etc. Recently, results of the developed at IOW GETM model with horizontal resolution of the model grid 1x2 nautical miles became available for the project (will be used in the future work), providing data for near bottom temperature, salinity and water speed averaged over the period 2000-2014 (Ulf Gräwe, personal communication).

On this basis further predictions using species distribution modeling will be done in areas where gaps in sampling effort remain. Here several methods will be tested to find the most appropriate for that particular purpose (including random forest, GLM and extensions).

The research paper aims to set a baseline for future studies for better recognition of links between benthic fauna and ecosystem functioning. The ICES Journal of Marine Science can be considered as an appropriate alternative for submission of the manuscript.

As an outlook based on the derived map in the further study the spatial distribution of specific macrozoobenthos functions (e.g. biodeposition and bioturbation) can be estimated. This will help to better understand the possible effects of natural or human-induced changes in macrofaunal communities on the hosting ecosystem.

It is essential to emphasize the unique chance to join all the data on benthic macrofauna around the Baltic Sea to finally create a Baltic Sea Benthic Atlas, first targeted on the community level.

Even though raw data engaged is partly not publicly assessable, all parties involved are willing to contribute to sound, reliable, transparent and credible scientific advice.

### **Financial overview**

The total amount of 3524 EUR was spent for travel/workshop organization, leaving the possibility for another meeting at the later stage of the project, with the remaining 2076 EUR left from the initial money received from ICES Science Fund.

Participation in ICES BEWG Meeting to present and consult the results and further steps is planned for two project co-workers – Michael L. Zettler and Mayya Gogina.



## **Presentations**

The idea of the project, preliminary joint dataset and first results were presented at the IMBER Future Ocean OSC in Bergen, Norway, 23-27 June 2014. The presentation has shown that there is quite a large interest in the aims of the map project. Major questions were on whether, when and where the data/map will be available, as the work is still in progress this remain open until now.

### **Summary of status and challenges:**

The biological data acquisition is completed, showing that full-coverage environmental data is limited.

Methods and algorithms are under development and following issues are to be clarified: linking biomass and abundance based community analysis; interpolation from point to surface is needed for a good quality high-resolution abiotic predictor layers. How to tackle temporal changes of community structure (2000-2013), the causality and how to assess uncertainties should be further addressed.

The manuscript/atlas structure is yet to be established.

## **References**

Clarke, K.R., Warwick, R.M., 2001. Change in Marine Communities: An Approach to Statistical Analysis and Interpretation, second ed. PRIMER-E, Plymouth.

De Cáceres, M., Legendre, P., 2009. Associations between species and groups of sites: indices and statistical inference. *Ecology* 90, 3566-3574.

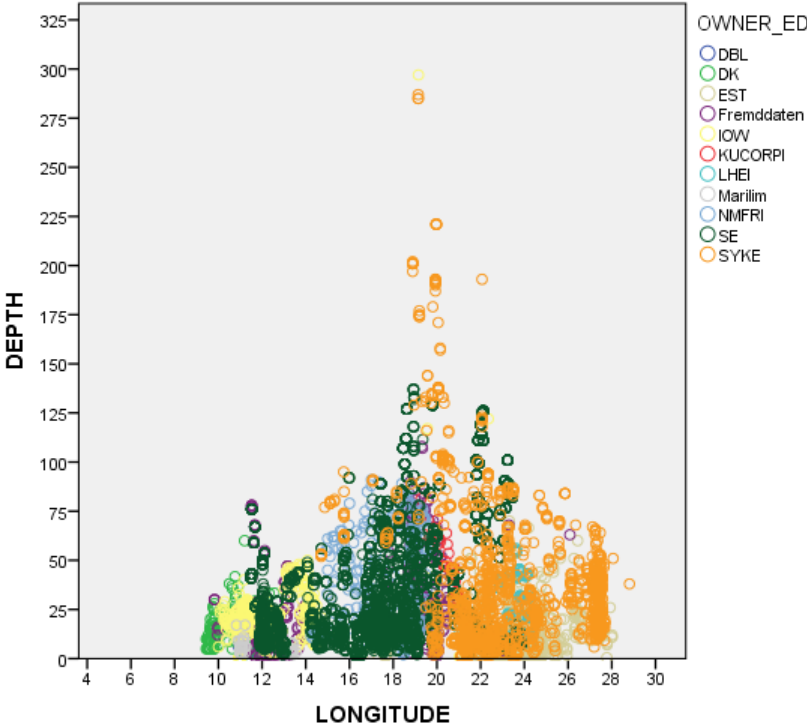
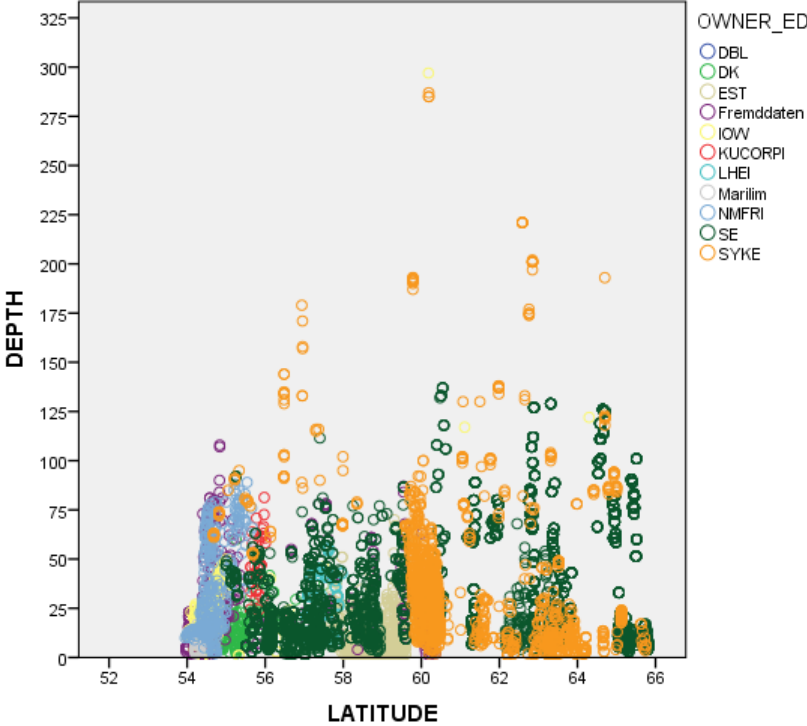
## Appendix 1. Sources of data

An overview of benthic fauna data considered at the beginning of the project.

DB	INSTITUTE	CONTACT PERSON	N VISITS	N VISITS PER AREA
DK	Department of Bioscience, Aarhus University, Denmark	Alf Bernhard Josefson	165	Arkona Basin, 8; Bay of Mecklenburg, 2; Flensburg Fjord, 32; Great Belt, 60; Kattegat, 16; Kiel Bay, 3; Little Belt, 34; The Sound, 10;
EST	Estonian Marine Institute, University of Tartu, Estonia	Jonne Kotta	1108	Eastern Gotland Basin, 35; Gulf of Finland, 296; Gulf of Riga, 689; Northern Baltic Proper, 87; Western Gotland Basin, 1
IOW	The Leibniz Institute for Baltic Sea Research, Warnemünde, Germany	Michael L. Zettler	1731	
KUCORPI	Coastal Research and Planning Institute, Klaipeda University, Lithuania	Darius Daunys	40	Eastern Gotland Basin, 40
LHEI	Latvian Institute of Aquatic Ecology, Latvia	Vadim Yermakov	195	Eastern Gotland Basin, 40; LHEI, 195; Eastern Gotland Basin, 6; Gulf of Riga, 189
NMFRI	National Marine Fisheries Research Institute, Poland	Jan Warzocha	319	Bornholm Basin, 114; Eastern Gotland Basin, 36; Gulf of Gdansk, 169
SE	Hafok AB, Sweden	Mats Blomqvist	4848	Arkona Basin, 52; Bornholm Basin, 390; Bothnian Bay, 1154; Bothnian Sea, 1544; Eastern Gotland Basin, 101; Kattegat, 533; Northern Baltic Proper, 418; The Quark, 104; The Sound, 70; Western Gotland Basin, 482
SYKE	Finnish Environment Institute SYKE /	Henrik Nygård	1383	Aland Sea, 14; Archipelago Sea, 207; Bornholm Basin, 33; Bothnian Bay, 233; Bothnian Sea, 238;

	Marine Research Centre, Finland			Eastern Gotland Basin, 43; Gulf of Finland, 496; Northern Baltic Proper, 22; The Quark, 97
Fremddaten	Literature based collection	Michael L. Zettler	1882	Aland Sea, 8; Archipelago Sea, 69; Arkona Basin, 408; Bay of Mecklenburg, 498; Bornholm Basin, 125; Bothnian Sea, 5; Darss-Zingst Lagoon, 1; Eastern Gotland Basin, 102; Flensburg Fjord, 6; Greifswald Lagoon, 37; Gulf of Finland, 6; Gulf of Gdansk, 212; Gulf of Riga, 1; Kattegat, 159; Kiel Bay, 10; Kiel Fjord, 20; Little Belt, 24; Northern Baltic Proper, 85; Rugia Lagoons, 3; Szczecin Lagoon, 7; The Sound, 9; Vistula Lagoon, 2; Warnow Estuary, 19; Western Gotland Basin, 31; Wismar Bay, 35; IOW, 1731; Aland Sea, 1; Arkona Basin, 414; Bay of Mecklenburg, 316; Bornholm Basin, 142; Bothnian Bay, 1; Bothnian Sea, 1; Curonian Lagoon, 23; Darss-Zingst Lagoon, 61; Eastern Gotland Basin, 2; Eckernfoerde Bay, 1; Great Belt, 22; Greifswald Lagoon, 21; Gulf of Gdansk, 11; Kattegat, 6; Kiel Bay, 438; Kiel Fjord, 2; Northern Baltic Proper, 1; Rugia Lagoons, 13; Szczecin Lagoon, 114; Trave Estuary, 20; Warnow Estuary, 54; Western Gotland Basin, 2; Wismar Bay, 65
Total			12548	

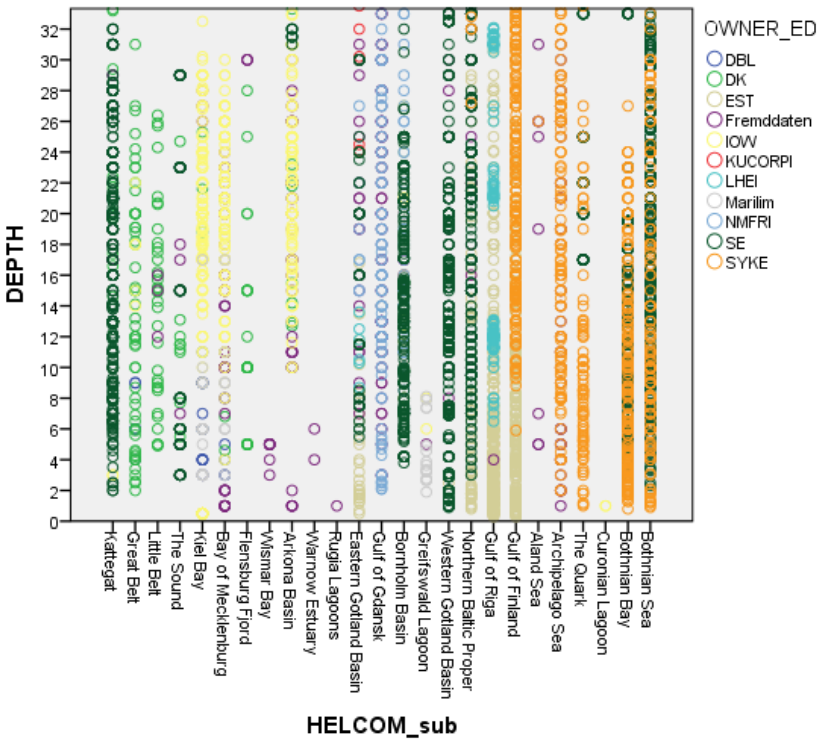
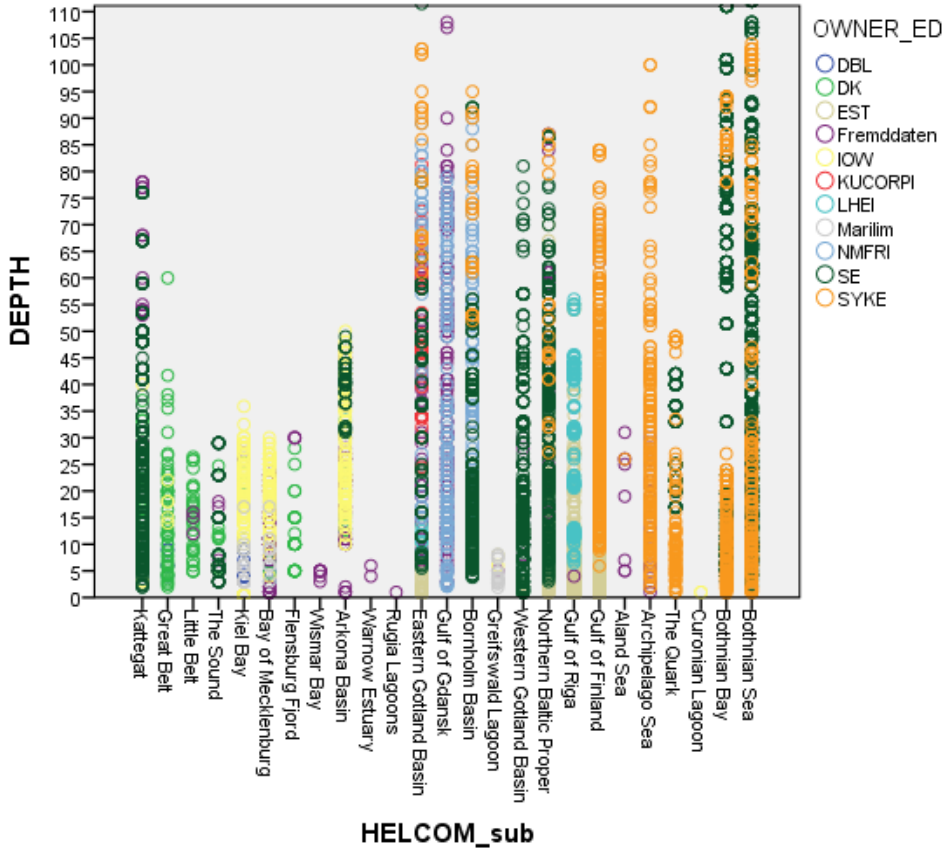
Sampling intensity as function of latitude, longitude and depth is shown (pattern dictated by bathymetry of the Baltic Sea and areas of interest of contributing partners).



REGION	N_RECORDS	%TOTAL
Kiel Bay	25864	17.964
Kattegat	22728	15.786
Bay of Mecklenburg	22438	15.584
Arkona Basin	14212	9.871
Bornholm Basin	7744	5.379
Bothnian Sea	7446	5.172
Gulf of Riga	7273	5.051
Western Gotland Basin	4399	3.055
Northern Baltic Proper	4172	2.898
Bothnian Bay	3883	2.697
Gulf of Finland	3811	2.647
Great Belt	3235	2.247
The Sound	3193	2.218
Gulf of Gdansk	2708	1.881
Eastern Gotland Basin	2522	1.752
Wismar Bay	1633	1.134
Little Belt	1282	0.890
Archipelago Sea	1264	0.878
Greifswald Lagoon	963	0.669
The Quark	827	0.574
Warnow Estuary	625	0.434
Szczecin Lagoon	514	0.357
Darss-Zingst Lagoon	441	0.306
Flensburg Fjord	308	0.214
Trave Estuary	255	0.177
Rugia Lagoons	119	0.083
Aland Sea	74	0.051
Eckernfoerde Bay	38	0.026
Kiel Fjord	7	0.005

Sampling equipment used for different datasets included and availability of biomass

<b>Row Labels</b>	<b>EQUIPMENT</b>	<b>SAMPLERARE</b>	<b>N_VISITS</b>	<b>BIOMASS</b>
<b>DK</b>	Haps	0.0143	191	To species
	van Veen grab	0.1	313	
<b>EST</b>	Birge	0.021	2153	Dry weight, to species
	Boxcore	0.04	19	
	Lenz	0.021	627	
	Petersen	0.044	344	
	van Veen grab	0.1	427	
<b>Fremddaten</b>	van Veen grab	0.1	1536	Partly, to species
<b>IOW</b>	van Veen grab	0.1	1370	To species
<b>ICES</b>	van Veen grab	0.1	40	-
<b>KUCORPI</b>	van Veen grab	0.1	313	Fresh weight
<b>LHEI</b>	van Veen grab	0.1 - 0.118	195	-
<b>ZIN</b>	van Veen grab	0.1	25	To species
<b>NMFRI</b>	Reineck	0.0225	48	To species
	van Veen grab	0.1	298	
<b>SE</b>	Smith-McIntyre grab	0.1	21	To species
	van Veen grab	0.0977 - 0.1221	6791	
<b>SYKE</b>	Boxcore	-	57	To species
	Ekman	0.0225 - 0.1445	247	
	van Veen grab	0.1 - 0.1153	1425	



## Appendix 2

List of taxa considered for communities classification

Original taxa	Included as	on at least one sampling event		
		Max cumulative abundance	Max relative abundance	N of stations
<i>Abra alba</i>	<i>Abra alba</i>	0.90	0.86	792
<i>Abra nitida</i>	<i>Abra nitida</i>	0.90	0.64	292
<i>Abra prismatica</i>	<i>Abra prismatica</i>	0.88	0.10	2
<i>Akera bullata</i>	<i>Akera bullata</i>	0.79	0.19	5
<i>Alderia modesta</i>	<i>Alderia modesta</i>	0.88	0.05	1
<i>Alitta succinea</i>	<i>Alitta succinea</i>	0.90	0.65	141
<i>Alitta virens</i>	<i>Alitta virens</i>	0.90	0.14	17
<i>Alkmaria romijni</i>	<i>Alkmaria romijni</i>	0.84	0.50	6
<i>Ampelisca brevicornis</i>	<i>Ampelisca</i>	0.90	0.71	115
<i>Ampelisca diadema</i>	<i>Ampelisca</i>	0.89	0.07	11
<i>Ampelisca macrocephala</i>	<i>Ampelisca</i>	0.79	0.14	4
<i>Ampelisca sp.</i>	<i>Ampelisca</i>	0.90	0.11	6
<i>Ampelisca tenuicornis</i>	<i>Ampelisca</i>	0.90	0.32	198
<i>Ampharete acutifrons</i>	<i>Ampharete acutifrons</i>	0.89	0.16	10
<i>Ampharete baltica</i>	<i>Ampharete baltica</i>	0.90	0.43	393
<i>Ampharete finmarchica</i>	<i>Ampharete finmarchica</i>	0.90	0.50	37
<i>Amphibalanus improvisus</i>	<i>Amphibalanus improvisus</i>	0.90	0.88	224
<i>Amphictene auricoma</i>	<i>Amphictene auricoma</i>	0.90	0.13	123
<i>Amphitrite cirrata</i>	<i>Amphitrite cirrata</i>	0.90	0.07	9
<i>Amphiura chiajei</i>	<i>Amphiura</i>	0.90	0.44	251
<i>Amphiura filiformis</i>	<i>Amphiura</i>	0.90	0.72	555
<i>Amphiura sp.</i>	<i>Amphiura</i>	0.89	0.04	8
<i>Ampithoe rubricata</i>	<i>Ampithoe rubricata</i>	0.90	0.36	38
<i>Angulus pygmaeus</i>	<i>Angulus pygmaeus</i>	0.88	0.33	31
<i>Angulus tenuis</i>	<i>Angulus tenuis</i>	0.89	0.88	49
<i>Anobothrus gracilis</i>	<i>Anobothrus gracilis</i>	0.90	0.15	111
<i>Aonides paucibranchiata</i>	<i>Aonides paucibranchiata</i>	0.90	0.25	13
Aoridae	Aoridae	0.88	0.09	10
<i>Apherusa bispinosa</i>	<i>Apherusa bispinosa</i>	0.90	0.29	24
<i>Apocorophium lacustre</i>	<i>Apocorophium lacustre</i>	0.87	0.08	3
<i>Aporrhais pespelecani</i>	<i>Aporrhais pespelecani</i>	0.89	0.12	4
<i>Arctica islandica</i>	<i>Arctica islandica</i>	0.90	0.88	635
<i>Arenicola marina</i>	<i>Arenicola marina</i>	0.90	0.30	30
<i>Aricidea cerrutii</i>	<i>Aricidea cerrutii</i>	0.89	0.43	19
<i>Aricidea minuta</i>	<i>Aricidea minuta</i>	0.90	0.07	55
<i>Aricidea suecica</i>	<i>Aricidea suecica</i>	0.90	0.19	269
<i>Artacama proboscidea</i>	<i>Artacama proboscidea</i>	0.90	0.09	29
<i>Asellus aquaticus</i>	<i>Asellus aquaticus</i>	0.90	0.90	69
<i>Astarte</i>	<i>Astarte</i>	0.90	0.81	665



Asterias rubens	Asterias rubens	0.90	0.75	155
Autonoe longipes	Autonoe longipes	0.90	0.08	5
Balanus crenatus	Balanus crenatus	0.89	0.72	65
Bathyporeia	Bathyporeia	0.90	0.88	610
Bithynia tentaculata	Bithynia tentaculata	0.90	0.77	47
Bittium reticulatum	Bittium reticulatum	0.90	0.88	140
Branchiostoma lanceolatum	Branchiostoma lanceolatum	0.89	0.25	56
Bylgides elegans	Bylgides elegans	0.87	0.09	4
Bylgides sarsi	Bylgides sarsi	0.90	0.90	612
Calliopijs laeviusculus	Calliopijs laeviusculus	0.89	0.30	42
Capitella	Capitella	0.90	0.88	155
Caprella linearis	Caprella linearis	0.90	0.07	27
Caulleriella killariensis	Caulleriella killariensis	0.90	0.15	35
Cerastoderma edule	Cerastoderma edule	0.90	0.86	48
Cerastoderma glaucum	Cerastoderma glaucum	0.90	0.89	906
Ceratopogonidae	Diptera	0.90	0.50	7
Chaetoderma nitidulum	Chaetoderma nitidulum	0.90	0.07	32
Chaetozone setosa	Chaetozone setosa	0.90	0.46	260
Chamelea gallina	Chamelea gallina	0.90	0.06	6
Chamelea striatula	Chamelea striatula	0.90	0.06	37
Cheirocratus sundevalli	Cheirocratus sundevalli	0.90	0.19	53
Chironomidae	Chironomidae	0.90	0.90	1514
Ciona intestinalis	Ciona intestinalis	0.90	0.14	19
Cochlodesma praetenuae	Cochlodesma praetenuae	0.90	0.14	21
Corbula gibba	Corbula gibba	0.90	0.89	703
Corophium volutator	Corophium volutator	0.90	0.89	735
Crangon crangon	Crangon crangon	0.89	0.17	23
Crassicorophium bonellii	Crassicorophium bonellii	0.89	0.10	15
	Crassicorophium			
Crassicorophium crassicornae	crassicornae	0.90	0.47	160
Crenella decussata	Crenella decussata	0.89	0.13	28
Cyanophthalma obscura	Cyanophthalma obscura	0.90	0.84	59
Cyathura carinata	Cyathura carinata	0.90	0.38	69
Cylichna cylindracea	Cylichna cylindracea	0.90	0.05	89
Dendrodoa grossularia	Dendrodoa grossularia	0.90	0.79	159
Diaphana minuta	Diaphana minuta	0.88	0.10	4
Diastylis	Diastylis	0.85	0.33	3
Diastylis bradyi	Diastylis	0.75	0.03	2
Diastylis lucifera	Diastylis	0.90	0.20	108
Diastylis rathkei	Diastylis	0.90	0.88	1187
Diplocirrus glaucus	Diplocirrus glaucus	0.90	0.12	232
Dipolydora caulleryi	Dipolydora caulleryi	0.90	0.03	7
Dipolydora coeca	Dipolydora coeca	0.90	0.56	43
Dipolydora quadrilobata	Dipolydora quadrilobata	0.90	0.69	254
Diptera	Diptera	0.90	0.84	81

<i>Dreissena polymorpha</i>	<i>Dreissena polymorpha</i>	0.90	0.81	38
<i>Dyopedos monacantha</i>	<i>Dyopedos monacantha</i>	0.90	0.24	28
<i>Echinocardium cordatum</i>	<i>Echinocardium cordatum</i>	0.90	0.42	79
<i>Echinocyamus pusillus</i>	<i>Echinocyamus pusillus</i>	0.90	0.41	133
<i>Echiurus echiurus</i>	<i>Echiurus echiurus</i>	0.82	0.09	4
<i>Ecrobia ventrosa</i>	Hydrobiidae	0.90	0.84	145
<i>Edwardsia danica</i>	<i>Edwardsia danica</i>	0.90	0.32	352
<i>Einhornia crustulenta</i>	<i>Einhornia crustulenta</i>	0.89	0.25	22
<i>Ennucula tenuis</i>	<i>Ennucula tenuis</i>	0.90	0.27	130
<i>Ensis directus</i>	<i>Ensis directus</i>	0.90	0.07	5
Ephemeroptera	Ephemeroptera	0.89	0.50	7
<i>Ericthonius punctatus</i>	<i>Ericthonius punctatus</i>	0.90	0.34	22
<i>Eteone longa</i>	<i>Eteone longa</i>	0.90	0.45	199
<i>Euchone papillosa</i>	<i>Euchone papillosa</i>	0.90	0.05	16
<i>Eudorella emarginata</i>	<i>Eudorella emarginata</i>	0.90	0.06	57
<i>Eudorella truncatula</i>	<i>Eudorella truncatula</i>	0.90	0.05	23
Eumida	Eumida	0.64	0.06	2
<i>Euspira nitida</i>	<i>Euspira nitida</i>	0.90	0.08	20
Exogone	Exogone	0.51	0.07	1
<i>Exogone hebes</i>	Exogone	0.90	0.11	16
<i>Exogone naidina</i>	Exogone	0.89	0.09	21
<i>Exogone sp.</i>	Exogone	0.89	0.21	6
<i>Fabricia stellaris</i>	<i>Fabricia stellaris</i>	0.90	0.21	15
<i>Fabriciola baltica</i>	<i>Fabriciola baltica</i>	0.90	0.14	27
<i>Facelina bostoniensis</i>	<i>Facelina bostoniensis</i>	0.88	0.14	4
<i>Galathowenia oculata</i>	<i>Galathowenia oculata</i>	0.90	0.42	263
<i>Gammarellus homari</i>	<i>Gammarellus homari</i>	0.89	0.08	3
Gammaridae	Gammaridae	0.90	0.86	949
<i>Gastrosaccus spinifer</i>	<i>Gastrosaccus spinifer</i>	0.90	0.23	102
<i>Gattyana cirrhosa</i>	<i>Gattyana cirrhosa</i>	0.85	0.09	2
<i>Glycera alba</i>	<i>Glycera alba</i>	0.90	0.07	163
<i>Glyphohesione klatti</i>	<i>Glyphohesione klatti</i>	0.89	0.07	5
<i>Goniada maculata</i>	<i>Goniada maculata</i>	0.90	0.07	175
<i>Halcampa duodecimcirrata</i>	<i>Halcampa duodecimcirrata</i>	0.90	0.08	29
<i>Halicryptus spinulosus</i>	<i>Halicryptus spinulosus</i>	0.90	0.85	329
<i>Haploops sp.</i>	<i>Haploops sp.</i>	0.71	0.08	3
<i>Harmothoe imbricata</i>	<i>Harmothoe</i>	0.90	0.29	114
<i>Harmothoe impar</i>	<i>Harmothoe</i>	0.90	0.55	129
<i>Harmothoe sp.</i>	<i>Harmothoe</i>	0.88	0.33	14
<i>Haustorius arenarius</i>	<i>Haustorius arenarius</i>	0.88	0.08	2
<i>Hediste diversicolor</i>	<i>Hediste diversicolor</i>	0.90	0.80	945
<i>Heteromastus filiformis</i>	<i>Heteromastus filiformis</i>	0.90	0.84	610
<i>Heterotanais oerstedii</i>	<i>Heterotanais oerstedii</i>	0.90	0.33	6
<i>Hyala vitrea</i>	<i>Hyala vitrea</i>	0.90	0.32	224
<i>Hydrobia acuta</i>	Hydrobiidae	0.87	0.02	1

Hydrobiidae	Hydrobiidae	0.90	0.83	360
Idotea balthica	Idotea	0.90	0.43	97
Idotea chelipes	Idotea	0.90	0.53	72
Idotea granulosa	Idotea	0.88	0.12	9
Idotea sp.	Idotea	0.90	0.38	20
Ischyrocerus anguipes	Ischyrocerus anguipes	0.89	0.08	5
Jaera sp.	Jaera sp.	0.90	0.63	110
Kurtiella bidentata	Kurtiella bidentata	0.90	0.73	1183
Labidoplax buskii	Labidoplax buskii	0.90	0.10	54
Laeospira corallinae	Laeospira corallinae	0.89	0.35	8
Lagis koreni	Lagis koreni	0.90	0.85	509
Lamprops fasciatus	Lamprops fasciatus	0.89	0.09	3
Lanice conchilega	Lanice conchilega	0.90	0.12	27
Laonome kroyeri	Laonome kroyeri	0.90	0.54	31
Lekanesphaera hookeri	Lekanesphaera hookeri	0.89	0.10	12
Lepidonotus squamatus	Lepidonotus squamatus	0.89	0.11	5
Lepidoptera	Lepidoptera	0.90	0.37	4
Leptocheirus hirsutimanus	Leptocheirus hirsutimanus	0.89	0.40	26
Leptocheirus pilosus	Leptocheirus pilosus	0.89	0.66	16
Leptochiton asellus	Leptochiton asellus	0.89	0.18	33
Leucon nasica	Leucon nasica	0.89	0.35	103
Levinsenia gracilis	Levinsenia gracilis	0.90	0.57	130
Lineus ruber	Lineus ruber	0.90	0.19	27
Littorina littorea	Littorina littorea	0.90	0.70	40
Littorina saxatilis	Littorina saxatilis	0.90	0.22	46
Lymnaea stagnalis	Lymnaea stagnalis	0.81	0.07	1
Lysilla loveni	Lysilla loveni	0.90	0.12	24
Macoma balthica	Macoma balthica	0.90	0.90	6113
Macoma calcarea	Macoma calcarea	0.90	0.50	80
Magelona alleni	Magelona alleni	0.90	0.13	35
Magelona mirabilis	Magelona mirabilis	0.89	0.08	11
Maldane sarsi	Maldane sarsi	0.90	0.64	147
Manayunkia aestuarina	Manayunkia aestuarina	0.90	0.38	31
Mangelia attenuata	Mangelia attenuata	0.42	0.42	2
Marenzelleria	Marenzelleria	0.90	0.90	2482
Medicorophium affine	Medicorophium affine	0.89	0.08	12
Megamphopus cornutus	Megamphopus cornutus	0.89	0.11	28
Melita palmata	Melita palmata	0.89	0.10	12
Metridium senile	Metridium senile	0.89	0.18	10
Microdeutopus anomalus	Microdeutopus anomalus	0.89	0.07	7
Microdeutopus gryllotalpa	Microdeutopus gryllotalpa	0.90	0.46	284
Modiolus modiolus	Modiolus modiolus	0.89	0.14	16
Molgula manhattensis	Molgula manhattensis	0.90	0.22	20
Monocorophium insidiosum	Monocorophium insidiosum	0.90	0.50	162
Monoporeia affinis	Monoporeia affinis	0.90	0.90	2202

Musculus discors	Musculus discors	0.90	0.34	50
Musculus niger	Musculus niger	0.90	0.14	51
Musculus subpictus	Musculus subpictus	0.90	0.12	77
Mya arenaria	Mya arenaria	0.90	0.79	897
Mya truncata	Mya truncata	0.90	0.50	30
Mytilus	Mytilus	0.90	0.90	2206
Nassarius	Nassarius	0.90	0.28	71
Nemertea	Nemertea	0.90	0.20	450
Neoamphitrite figulus	Neoamphitrite figulus	0.88	0.49	8
Nephtys caeca	Nephtys	0.90	0.24	256
Nephtys ciliata	Nephtys	0.90	0.47	216
Nephtys hombergii	Nephtys	0.90	0.53	508
Nephtys incisa	Nephtys	0.90	0.39	134
Nephtys longosetosa	Nephtys	0.90	0.17	51
Nephtys paradoxa	Nephtys	0.88	0.02	1
Nephtys pente	Nephtys	0.86	0.02	8
Nephtys sp.	Nephtys	0.90	0.17	12
Neptunea antiqua	Neptunea antiqua	0.44	0.18	1
Nereididae	Nereididae	0.90	0.46	24
Nereimyra punctata	Nereimyra punctata	0.90	0.29	74
Nereis pelagica	Nereis pelagica	0.88	0.22	5
Nicolea zostericola	Nicolea zostericola	0.90	0.07	36
Nicomache minor	Nicomache minor	0.87	0.14	10
Notomastus latericeus	Notomastus latericeus	0.90	0.08	37
Nucula nitidosa	Nucula nitidosa	0.90	0.68	225
Nucula sulcata	Nucula sulcata	0.90	0.45	44
Nymphon brevistre	Nymphon brevistre	0.89	0.07	11
Odostomia scalaris	Odostomia scalaris	0.90	0.38	38
Oligochaeta	Oligochaeta	0.90	0.90	3015
Onchidoris muricata	Onchidoris muricata	0.56	0.14	3
Onoba semicostata	Onoba semicostata	0.90	0.28	114
Ophelia borealis	Ophelia borealis	0.89	0.47	106
Ophelia limacina	Ophelia limacina	0.89	0.25	18
Ophelia rathkei	Ophelia rathkei	0.90	0.70	39
Ophelina acuminata	Ophelina acuminata	0.90	0.09	23
Ophiodromus flexuosus	Ophiodromus flexuosus	0.90	0.04	11
Ophiura albida	Ophiura	0.90	0.50	226
Ophiura ophiura	Ophiura	0.80	0.02	2
Ophiura sp.	Ophiura	0.90	0.17	88
Orbinia sertulata	Orbinia sertulata	0.84	0.06	3
Owenia fusiformis	Owenia fusiformis	0.89	0.79	49
Palaemon adspersus	Palaemon adspersus	0.61	0.07	4
Pallaseopsis quadrispinosa	Pallaseopsis quadrispinosa	0.86	0.26	11
Paraonis fulgens	Paraonis fulgens	0.90	0.39	28
Parvicardium hauniense	Parvicardium hauniense	0.90	0.06	6

Parvicardium minimum	Parvicardium minimum	0.90	0.06	17
Parvicardium pinnulatum	Parvicardium pinnulatum	0.90	0.68	370
Parvicardium scabrum	Parvicardium scabrum	0.89	0.08	33
Pectinaria auricoma	Pectinaria auricoma	0.90	0.06	12
Peringia ulvae	Hydrobiidae	0.90	0.90	2423
Perioculodes longimanus	Perioculodes longimanus	0.89	0.05	6
Phaxas pellucidus	Phaxas pellucidus	0.90	0.38	86
Pherusa plumosa	Pherusa plumosa	0.90	0.11	65
Philine aperta	Philine aperta	0.90	0.17	53
Pholoe assimilis	Pholoe assimilis	0.90	0.09	110
Pholoe baltica	Pholoe baltica	0.90	0.25	451
Pholoe inornata	Pholoe inornata	0.90	0.21	142
Phoronis muelleri	Phoronis	0.90	0.53	362
Phoronis sp.	Phoronis	0.90	0.33	100
Phoxocephalus holbolli	Phoxocephalus holbolli	0.90	0.30	181
Phtisica marina	Phtisica marina	0.90	0.03	16
Phyllodoce groenlandica	Phyllodoce groenlandica	0.89	0.12	23
Phyllodoce maculata	Phyllodoce maculata	0.86	0.06	2
Phyllodoce mucosa	Phyllodoce mucosa	0.90	0.10	63
Physa fontinalis	Physa fontinalis	0.89	0.07	1
Pisidium sp.	Pisidium sp.	0.90	0.90	76
Platynereis dumerilii	Platynereis dumerilii	0.90	0.06	9
Podocoryna carnea	Podocoryna carnea	0.87	0.59	12
Polycirrus medusa	Polycirrus medusa	0.90	0.29	122
Polydora ciliata	Polydora	0.90	0.82	78
Polydora cornuta	Polydora	0.90	0.20	55
Polydora sp.	Polydora	0.89	0.43	5
Polynoidae	Polynoidae	0.83	0.06	4
Pontocrates arenarius	Pontocrates arenarius	0.89	0.09	29
Pontoporeia femorata	Pontoporeia femorata	0.90	0.90	441
Potamopyrgus antipodarum	Potamopyrgus antipodarum	0.90	0.89	282
Priapulus caudatus	Priapulus caudatus	0.90	0.17	31
Prionospio fallax	Prionospio fallax	0.90	0.43	270
Prionospio steenstrupi	Prionospio steenstrupi	0.66	0.14	2
Protomedeia fasciata	Protomedeia fasciata	0.90	0.23	46
Psammechinus miliaris	Psammechinus miliaris	0.90	0.08	5
Psolus phantapus	Psolus phantapus	0.87	0.06	4
Pusillina inconspicua	Pusillina inconspicua	0.90	0.25	38
Pusillina sarsii	Pusillina sarsii	0.90	0.34	14
Pygospio elegans	Pygospio elegans	0.90	0.87	1904
Radix balthica	Radix balthica	0.90	0.82	71
Retusa obtusa	Retusa obtusa	0.90	0.07	12
Retusa truncatula	Retusa truncatula	0.90	0.26	101
Rhodine gracilior	Rhodine gracilior	0.90	0.39	99
Rhodine loveni	Rhodine loveni	0.90	0.06	22

Rissoa membranacea	Rissoa membranacea	0.89	0.06	14
Sabellidae	Sabellidae	0.85	0.06	2
Saduria entomon	Saduria entomon	0.90	0.84	301
Saxicavella jeffreysi	Saxicavella jeffreysi	0.90	0.13	10
Scalibregma inflatum	Scalibregma inflatum	0.90	0.72	243
Scolecopsis foliosa	Scolecopsis foliosa	0.89	0.02	3
Scoloplos armiger	Scoloplos armiger	0.90	0.90	1680
Scrobicularia plana	Scrobicularia plana	0.81	0.15	4
Seraphsidae	Seraphsidae	0.84	0.06	5
Sphaerodorum gracilis	Sphaerodorum gracilis	0.90	0.12	103
Spio arndti	Spio arndti	0.90	0.11	48
Spio filicornis	Spio arndti	0.90	0.76	38
Spio goniocephala	Spio goniocephala	0.90	0.24	201
Spio martinensis	Spio martinensis	0.88	0.12	6
Spiophanes bombyx	Spiophanes bombyx	0.90	0.44	113
Spiophanes kroyeri	Spiophanes kroyeri	0.90	0.08	63
Spirorbis spirorbis	Spirorbis spirorbis	0.89	0.34	9
Spisula subtruncata	Spisula subtruncata	0.90	0.19	32
Stenothoe monoculoides	Stenothoe monoculoides	0.82	0.37	6
Streblospio shrubsolii	Streblospio shrubsolii	0.90	0.16	9
Streptosyllis websteri	Streptosyllis websteri	0.90	0.21	33
Syllidae	Syllidae	0.88	0.10	8
Tanaissus lilljeborgi	Tanaissus lilljeborgi	0.90	0.27	51
Tellimya ferruginosa	Tellimya ferruginosa	0.90	0.13	25
Tellimya tenella	Tellimya tenella	0.90	0.14	46
Tellina fabula	Tellina fabula	0.90	0.86	91
Tenellia adpersa	Tenellia adpersa	0.88	0.08	1
Terebellides stroemii	Terebellides stroemii	0.90	0.65	586
Theodoxus fluviatilis	Theodoxus fluviatilis	0.90	0.83	367
Thracia phaseolina	Thracia phaseolina	0.90	0.48	79
Thyasira flexuosa	Thyasira flexuosa	0.90	0.65	123
Thysanocardia procera	Thysanocardia procera	0.74	0.06	1
Travisia forbesii	Travisia forbesii	0.90	0.24	64
Trichoptera	Trichoptera	0.90	0.17	11
Trochochaeta multisetosa	Trochochaeta multisetosa	0.90	0.67	106
Turbellaria	Turbellaria	0.90	0.05	16
Turritella communis	Turritella communis	0.90	0.22	92
Urothoe elegans	Urothoe elegans	0.89	0.10	13
Urothoe grimaldii	Urothoe grimaldii	0.78	0.15	5
Urothoe poseidonis	Urothoe poseidonis	0.90	0.14	7
Valvata macrostoma	Valvata macrostoma	0.80	0.40	4
Valvata piscinalis	Valvata piscinalis	0.88	0.39	9
Virgularia mirabilis	Virgularia mirabilis	0.89	0.05	5

### Appendix 3

#### Results of SIMPER analysis for 10 major communities

##### Group 2

Average similarity: 36.33

Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%
Hydrobiidae	5.07	13.86	1.59	38.16	38.16
Mytilus	2.89	4.62	0.67	12.71	50.87
Pygospio elegans	2.47	3.69	0.56	10.15	61.02
Macoma balthica	2.17	3.54	0.66	9.73	70.75
Oligochaeta	1.81	2.20	0.53	6.07	76.82
Mya arenaria	1.73	2.10	0.49	5.79	82.61
Cerastoderma glaucum	1.55	1.68	0.40	4.61	87.22
Hediste diversicolor	1.25	1.25	0.39	3.44	90.66

##### Group 3

Average similarity: 29.52

Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%
Scoloplos armiger	2.70	3.72	1.02	12.62	12.62
Nephtys	1.97	2.94	1.00	9.96	22.58
Abra alba	2.45	2.78	0.77	9.42	31.99
Kurtiella bidentata	2.48	2.75	0.97	9.32	41.32
Diastylis	2.39	2.75	0.76	9.30	50.62
Corbula gibba	2.14	2.69	0.79	9.11	59.74
Lagis koreni	1.73	1.34	0.55	4.52	64.26
Arctica islandica	1.43	1.16	0.53	3.93	68.19
Hydrobiidae	1.50	1.02	0.44	3.46	71.65
Pygospio elegans	1.36	0.70	0.41	2.38	74.03
Heteromastus filiformis	1.02	0.64	0.41	2.18	76.21
Macoma balthica	1.14	0.60	0.39	2.05	78.26
Mytilus	1.41	0.60	0.40	2.04	80.30
Terebellides stroemii	1.16	0.54	0.34	1.84	82.14
Ampharete baltica	0.86	0.41	0.39	1.39	83.53
Parvicardium pinnulatum	1.04	0.37	0.34	1.25	84.78
Oligochaeta	0.83	0.31	0.31	1.04	85.82
Dipolydora quadrilobata	0.87	0.29	0.31	0.99	86.81
Bylgides sarsi	0.58	0.28	0.27	0.96	87.77
Astarte	0.91	0.28	0.30	0.93	88.70
Aricidea suecica	0.74	0.27	0.30	0.91	89.61
Scalibregma inflatum	0.65	0.22	0.17	0.74	90.35

##### Group 1

Average similarity: 27.01

Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%
Macoma balthica	2.66	12.51	0.79	46.31	46.31
Monoporeia affinis	1.67	5.39	0.46	19.94	66.25
Marenzelleria	1.70	4.80	0.45	17.78	84.03
Oligochaeta	1.06	1.97	0.31	7.29	91.32

##### Group 4

Average similarity: 32.29

Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%
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Macoma balthica	2.60	9.98	1.00	30.91	30.91
Bylgides sarsi	1.77	9.85	0.82	30.51	61.42
Diastylis	1.84	4.13	0.59	12.78	74.19
Scoloplos armiger	1.55	3.67	0.46	11.36	85.56
Pontoporeia femorata	0.97	1.77	0.39	5.49	91.04

Group 6

Average similarity: 46.31

Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%
Mytilus	5.26	43.47	2.40	93.87	93.87

Group 9

Average similarity: 31.46

Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%
Lagis koreni	3.87	16.62	1.41	52.83	52.83
Abra alba	1.94	3.28	0.57	10.43	63.26
Nephtys	1.25	2.50	0.65	7.94	71.20
Corbula gibba	0.95	1.67	0.47	5.30	76.50
Harmothoe	1.01	1.66	0.51	5.27	81.77
Halicryptus spinulosus	0.90	1.27	0.40	4.05	85.82
Mytilus	1.23	0.77	0.30	2.43	88.25
Kurtiella bidentata	0.76	0.63	0.28	2.01	90.26

Group 5

Average similarity: 34.17

Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%
Amphiura	3.77	5.14	1.10	15.06	15.06
Kurtiella bidentata	2.84	3.94	1.48	11.54	26.59
Nephtys	1.63	1.93	0.83	5.64	32.23
Phoronis	1.92	1.89	0.77	5.54	37.77
Abra nitida	1.68	1.52	0.75	4.46	42.23
Ophiura	1.47	1.32	0.62	3.87	46.10
Abra alba	1.50	1.27	0.57	3.73	49.83
Prionospio fallax	1.57	1.27	0.67	3.71	53.54
Pholoe baltica	1.34	1.18	0.61	3.45	56.99
Ampelisca	1.21	1.01	0.68	2.94	59.93
Thyasira flexuosa	1.34	0.99	0.53	2.90	62.83
Galathowenia oculata	1.41	0.98	0.58	2.87	65.70
Scalibregma inflatum	1.39	0.93	0.55	2.73	68.43
Ennucula tenuis	1.21	0.86	0.45	2.52	70.95
Nucula nitidosa	1.28	0.86	0.49	2.52	73.47
Pholoe inornata	1.25	0.83	0.49	2.43	75.90
Corbula gibba	1.02	0.77	0.49	2.24	78.14
Nemertea	0.93	0.68	0.51	1.99	80.14
Heteromastus filiformis	0.96	0.54	0.40	1.57	81.71
Hyala vitrea	0.89	0.50	0.29	1.45	83.17
Diplocirrus glaucus	0.83	0.46	0.39	1.35	84.51
Glycera alba	0.72	0.42	0.39	1.22	85.73
Arctica islandica	0.68	0.39	0.32	1.14	86.87
Rhodine gracilior	0.81	0.36	0.32	1.04	87.92
Goniada maculata	0.63	0.34	0.37	1.00	88.92
Chaetozone setosa	0.65	0.33	0.36	0.98	89.90
Maldane sarsi	0.84	0.32	0.30	0.95	90.85



Group 10

Average similarity: 31.92

Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%
Oligochaeta	2.73	5.98	1.27	18.73	18.73
Scoloplos armiger	2.24	5.73	1.28	17.96	36.68
Kurtiella bidentata	2.78	4.59	0.96	14.38	51.06
Mytilus	2.35	4.06	0.74	12.71	63.77
Harmothoe	1.98	2.89	0.76	9.06	72.83
Bittium reticulatum	1.62	1.79	0.46	5.62	78.45
Echinocyamus pusillus	1.91	1.79	0.45	5.62	84.07
Microdeutopus gryllotalpa	1.07	0.68	0.35	2.13	86.20
Asterias rubens	1.03	0.68	0.34	2.12	88.31
Alitta succinea	0.86	0.62	0.24	1.93	90.25

Group 7

Average similarity: 35.30

Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%
Phoronis	2.95	3.60	1.14	10.19	10.19
Spiophanes bombyx	2.55	3.42	1.24	9.68	19.87
Scoloplos armiger	2.43	3.31	1.28	9.38	29.24
Tellina fabula	2.39	2.53	0.78	7.17	36.41
Nephtys	1.77	2.19	0.96	6.20	42.62
Ophelia borealis	1.72	1.74	0.65	4.93	47.54
Bathyporeia	1.80	1.70	0.63	4.80	52.35
Ampelisca	1.78	1.55	0.67	4.39	56.74
Kurtiella bidentata	1.72	1.55	0.69	4.38	61.12
Edwardsia danica	1.54	1.52	0.73	4.31	65.43
Echinocyamus pusillus	1.58	1.38	0.59	3.90	69.34
Spio arndti	1.48	1.31	0.71	3.72	73.06
Arctica islandica	1.32	1.04	0.62	2.94	76.00
Thracia phaseolina	1.30	0.91	0.51	2.57	78.57
Nemertea	1.07	0.87	0.57	2.47	81.03
Chaetozone setosa	0.99	0.71	0.48	2.01	83.05
Amphiura	0.96	0.53	0.46	1.52	84.56
Branchiostoma lanceolatum	0.85	0.47	0.37	1.32	85.89
Lanice conchilega	0.85	0.39	0.28	1.11	87.00
Corbula gibba	0.83	0.38	0.25	1.08	88.07
Ophiura	0.75	0.37	0.28	1.04	89.11
Pholoe baltica	0.77	0.32	0.34	0.89	90.00

Group 8

Average similarity: 34.11

Species	Av.Abund	Av.Sim	Sim/SD	Contrib%	Cum.%
Angulus tenuis	4.57	16.78	1.75	49.20	49.20
Scoloplos armiger	1.87	4.85	0.90	14.23	63.43
Kurtiella bidentata	1.94	3.54	0.65	10.38	73.81
Corbula gibba	1.52	2.57	0.62	7.53	81.34
Pygospio elegans	1.15	1.38	0.40	4.03	85.37
Bathyporeia	1.15	1.26	0.32	3.69	89.06
Nephtys	0.93	1.05	0.41	3.08	92.14

**Appendix 4**

**Abundance (ind/m<sup>2</sup>)**

**Biomass fresh weight (g/m<sup>2</sup>)**

*Abra alba*

*Arctica islandica*

*Mya arenaria*

*Cerastoderma glaucum*

*Macoma balthica*