The Arctic Traits Database – A repository of arctic benthic invertebrate traits

Renate Degen[1] & Sarah Faulwetter[2, 3]

[1] Department of Limnology and Bio-Oceanography, University of Vienna, 1090 Vienna, Austria
[2] Department of Zoology, University of Patras, 26504 Rio, Greece
[3] Institute of Oceanography, Hellenic Centre for Marine Research, 19013 Anavyssos, Greece

Correspondence to: Renate Degen (renate.degen@hotmail.com)

Abstract. The recently increased interest in marine trait-based studies highlights one general demand – the access to standardized, reference-based trait information. This demand holds especially true for polar regions, where the gathering of ecological information is still challenging. The Arctic Traits Database is a freely accessible online repository (https://doi.org/10.25365/phaidra.49; https://www.univie.ac.at/arctictraits) that fulfils these requests for one important component of polar marine life, the Arctic benthic macroinvertebrates. It accounts for 1) obligate traceability of information (every entry is linked to at least one source), 2) exchangeability among trait platforms (use of most common download formats), 3) standardization (use of most common terminology and coding scheme), and 4) user friendliness (granted by an intuitive web-interface and rapid and easy download options, including for the first time the option to download a fuzzy coded trait matrix). The combination of these aspects makes the Arctic Traits Database the currently most sophisticated online accessible trait platform in (not only) marine ecology and a role-model for prospective databases of other marine compartments or other (also non-marine) ecosystems. At present the database covers 19 traits (80 trait categories) and holds altogether 14242 trait entries for 1911 macro- and megabenthic taxa. Thus, the Arctic Traits Database will foster and facilitate trait-based approaches in polar regions in the future and increase our ecological understanding of this rapidly changing system.

1 Introduction

The interest in trait-based approaches – i.e. such that consider the life history, morphological, physiological and behavioral characteristics (i.e. traits) of species – in the marine realm has been growing tremendously in the last decades (reviewed in Degen et al., 2018) (Fig. 1). Reasons for the increasing popularity of these approaches are that they offer a variety of additional options to solely species-based methods: Traits can be analyzed across wide geographical ranges and across species pools (Bernhardt-Römermann et al., 2011), they can be used to calculate a variety of functional diversity indices (Schleuter et al., 2010), to estimate functional redundancy (Darr et al., 2014), or be used as indicators of ecosystem functioning (Bremner et al., 2006). Given the rapid changes we observe in many marine regions of the world, and especially in the Arctic Ocean (Wassmann et al., 2011), the potential to indicate vulnerability to climate change and biodiversity loss, or to estimate climate change effects on ecosystem functions is another inherent advantage of trait-based approaches (Foden et al., 2013; Hewitt et al., 2016).

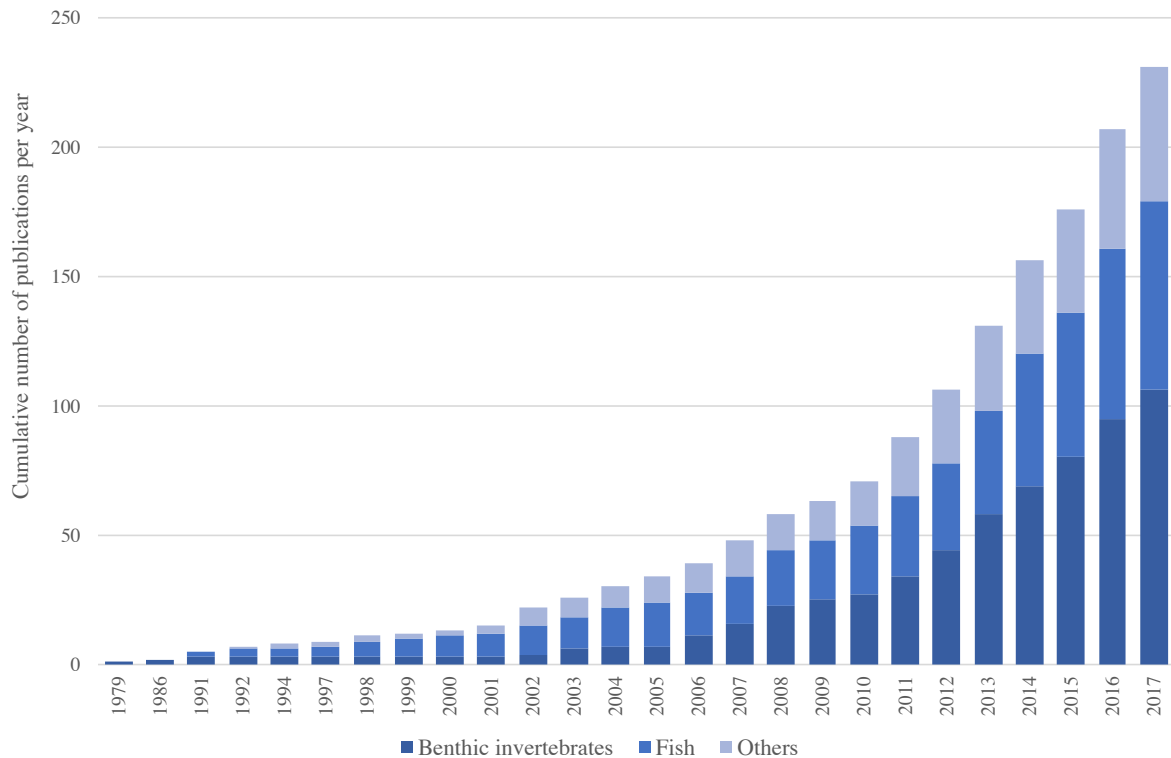



Figure 1. Cumulative number of marine trait-based studies based on the literature review of 233 studies from the marine realm by Degen et al. (2018).

Although the methodical diversity and complexity of trait-based approaches has broadened in the last years (Beauchard et al., 2017; Kleyer et al., 2012), the underlying data are always species traits. Trait information, however, is often not easy to find, and its collation requires a time and labor intensive survey of literature, databases, field data, and expert knowledge. This holds especially true for the polar regions, as ecological information for many polar marine taxa is still scarce, and only few publications supplement traceable resources of trait information (e.g. Kokarev et al., 2017). An additional obstacle is that existing trait repositories focus mainly on species from temperate regions. The increasing variability in terminology that surrounds traits is another challenge, and recent publications stress the importance of standardization in order to facilitate meta-analyses and comparison of results (Costello et al., 2015; Degen et al., 2018). Several online accessible trait databases specialize in specific taxonomic groups such as fish, polychaetes, or copepods, while others cover a wider part of the marine community (Table 1). The number of traits included and the form of access varies considerably among the different repositories. The database for marine copepods (Brun et al., 2017) contains 14 traits, whereas Fishbase (http://www.fishbase.org), polytraits (Faulwetter et al., 2014) and BIOTIC (http://www.marlin.ac.uk/biotic) contain more than 40 traits. Some repositories allow only for online browsing, while others enable different forms of download that range from spread sheets to different matrix formats (Table 1). No traits repository explicitly comprising polar species exists so far.

Table 1. List of marine trait databases or repositories. "Component" indicates the organism group targeted, "Access options" indicates in which forms the data can be accessed. Reference and web links are provided.

Component	Access options	Publication, web links
Copepoda	Download of excel workbook via PANGAEA, traits provided as original values or binary code (0/1), references per trait provided.	Brun et al. (2017) https://doi.pangaea.de/10.1594/ PANGAEA.862968

Polychaeta	Download of full database or specified subsets in various formats (references and partly original quote and page number provided), online via browsing the *Polychaetes Scratchpads*	Faulwetter et al. (2014) http://polytraits.lifewatchgreece.eu http://polychaetes.lifewatchgreece.eu
Benthos	Download of trait information in several matrix formats; as text and for certain traits as binary (0/1) code, also browsing online	Biological Traits Information Catalogue (BIOTIC) http://www.marlin.ac.uk/biotic
Fish	Browse online, programmatically via Application Programming Interface (API) and R package rfishbase	Froese, R. and D. Pauly. Editors. 2018. FishBase. www.fishbase.org, version (02/2018)
Benthos	Browse online	Marine Macrofauna Genus Trait Handbook, http://www.genustraithandbook.org.uk
Corals	Browse online, download as *.csv file, traits provided as original values or text information, references provided.	https://coraltraits.org/
Phytoplankton (coastal)	Download of excel workbook, traits provided as original values or binary code (0/1).	Klais et al. (2017) https://www.riinaklais.com/phytotraits
All marine	Browse online	Marine Species Traits, www.marinespecies.org/traits
All marine	Browse online	Sea Life Base, http://www.sealifebase.org
Fossil groups	Browse online	Neogene Marine Biota of Tropical America (NMiTA) http://eusmilia.geology.uiowa.edu
All biota	Browse online, programmatically via API	Encyclopedia of Life (EoL), http://www.eol.org

55 With the here presented Arctic Traits Database we aim to bridge some of the above-mentioned issues for one important compartment of marine life: the Arctic macro- and megabenthic invertebrates. In order to fulfil the communities' demand for standardization and comparability only those traits and trait categories are included, that are most frequently used in topical publications or which are already provided in freely accessible trait databases (Table 1). Regarding download options and traceability we follow the successful example given in Faulwetter et

60 al. (2014) and provide download of trait data in different tabular formats (i.e. data in columns, once following a database-specific format and once DarwinCore) (Wieczorek et al., 2012). The use of these formats guarantees that the included trait information can be easily shared between trait repositories and that the content is fully exploitable both by humans and computers. Every trait code is backed up by at least one reference, and where possible the original quote and page number are provided. In addition to above mentioned formats, for the first time trait

65 information is made available also in a fuzzy-coded and ready-to-use matrix format, that can be directly incorporated into appropriate analysis software.

By providing the Arctic Traits Database to the community of benthic ecologists we aim to provide a sound basis for prospective trait-based approaches in polar regions which will in return aid our overall understanding of these unique and rapidly changing ecosystems.

70 **2 Data**

2.1 Taxon data

The current taxa in the database are a subset of the dataset compiled in the frame of the "Arctic Traits Project" (Austrian Science Fund FWF, T801-B29), with focus on pan-Arctic benthic invertebrate macro- and megafauna. This dataset comprises species lists from published studies of collaborators (Blanchard et al., 2013a, 2013b;

75 Grebmeier et al., 2015), but also from so far unpublished sampling campaigns (e.g. field courses of the University

Center in Svalbard, UNIS, 2007-2017). The regional coverage currently comprises the Chukchi Sea and the Svalbard area. At present stage mainly species in the macrofauna size class have been uploaded.

2.2 Trait data

80 Currently we consider 19 traits and 80 trait categories that reflect the morphology, life history, and the behavior of Arctic benthic invertebrates (Table 3). All traits are in categorical format, i.e. belonging to one out of up to six clearly defined trait categories (see Table 3). The three continuous traits included (body size, longevity, and depth distribution) are converted into categories, but the associated text information assures accessibility to users also in their original, numerical or continuous format.

The choice of which traits to include in the database is based on the following considerations: 1) trait
85 information should be available for and applicable to all benthic taxa (Costello et al. 2015), 2) traits used in previous studies and databases should be favored to enable comparisons across studies (Degen et al. 2018), and 3) the traits should be usable across a wide geographical area (Bremner et al. 2006). In order to fulfil this last precondition, the trait body size is provided as "maximum body size as adult" (see also Table 3). While clearly a tradeoff in regard to the detection of intraspecific plasticity, it enables the use of this trait across large spatial scales.
90 Recent trait-based studies emphasize the importance of standardized traits and trait terminology to ensure that data can be integrated more easily in the future (Costello et al 2015, Degen et al. 2018, Faulwetter et al. 2014). To meet these requirements of the scientific community, the Arctic Traits Database includes seven of the ten traits prioritized in Costello et al. (2015): "depth range", "substratum affinity", "mobility", "skeleton", "diet", "body size" and "reproduction" (Table 3). The remaining three traits emphasized in Costello et al. (2015) – taxonomic
95 identity, environment, and geography – are not included. For taxonomic traits, every species in the database is bidirectionally deep linked (i.e. connected via a hyperlink) to the World register of Marine Species (WoRMS Editorial Board 2017; http://www.marinespecies.org/). For more detailed biogeographic information we refer users to the Global Biodiversity Information System (GBIF; http://www.gbif.org/) or the Ocean Biogeographic Information System (OBIS; http://www.iobis.org). We do include, however, the trait "zoogeography", which
100 enables a differentiation between typical arctic and boreal or cosmopolitan taxa. Of the 19 traits used here, 17 are also identical to those used by the BIOTIC database (MarLIN 2006, Table 1), one of the most comprehensive databases on biological traits of marine organisms. BIOTIC also includes the trait "salinity". We cover salinity preferences within the trait "tolerance", which accounts also for temperature and pollution tolerance (see Table 3 for details). Traits we include in addition are "skeleton", and "mobility" (i.e. the relative degree of movement).
105 Although physiological traits are of high interest in trait-based studies, we do not include them as they are not easily retrieved for many (arctic) benthic taxa (one of the preconditions for inclusion in the database as stated above). In addition, physiological traits (e.g. growth rate, respiration rate, ingestion rate) depend on body mass and temperature (Brown et al., 2004), which can vary tremendously among Arctic regions, contradicting that the provided traits information should be usable across a wide geographical area.

110 **Table 2.** Trait terminology as used in the Arctic Traits Database, BIOTIC, Costello et al. 2015, and in "other" marine trait-based studies (i.e. studies reviewed in Degen et al. 2018, list non-exhaustive, see Appendix 1 of Degen et al. 2018 for total trait list and corresponding literature references). Be aware that the Arctic Traits Database and BIOTIC consider only benthic taxa, while Costello et al. (2015) and the studies summarized in "Other" cover all marine groups.

Arctic Traits Database	BIOTIC	Costello et al. (2015)	Other

Body size	Body size	Body size	Body size/length/height, Largest radius, Biovolume, Coverage
Body form	Growth form	–	Body form, Body design, Body shape, Growth form, Growth type, Functional form group, Morphology
Fragility	Fragility	–	Fragility, Structural robustness, Shell strength
Skeleton	–	Skeleton	Skeletal composition/ thickness/material/density
Sociability	Sociability	–	Sociability, Schooling, Gregariousness, Social group size, Social behavior
Reproduction	Reproductive type	Reproduction	Reproduction, Reproduction type, Reproductive method/strategy/type/technique
Larval development	Developmental mechanism	–	Larval development, Larvae type, Larval feeding, Larval development location, Developmental mode/type/mechanism/technique
Life span	Life span	–	Longevity, Age, Life span, Maturity, Life duration, Generation time
Environmental position	Environmental position	–	Environment, Environmental position, Habitat, Vertical distribution, Sediment position, Living position, Life zone
Living habit	Living habit	–	Living habit, Habit, Life habit, Life form, Habitat, Living mode, Habitat structure
Mobility	–	Mobility	Mobility, Relative mobility, Degree of mobility, Mobility within sediment
Adult movement	Mobility/Movement	–	Adult movement, Mobility, Movement method/type, Locomotion
Feeding habit	Feeding habit	–	Feeding habit/behavior/method/type/apparatus, Resource capture method, Trophic mode, Oral gape position/height/surface, Protrusion
Trophic level	Typical food types	Diet	Trophic level, Diet, Food type, Trophic group, Dietary group
Bioturbation	Bioturbation	–	Bioturbation mode/type/potential, Sediment movement/reworking/transport, Direction of sediment transport, Reworking mode, Fecal deposition, Irrigation
Tolerance	Salinity	–	Tolerance, Tolerance limits, Salinity tolerance, Survival salinity/temperature, Temperature optimum, Thermal affinity, Hypoxia tolerance, Tolerance to pollutants, Ecological group, Resilience, Condition index
Zoogeography	Biogeographic range	–	Biogeography, Geographical range/distribution, Range size, Native region, Median latitude
Depth range	Biological zone	Depth range	Depth range/regime, Diving depth
Substratum affinity	Substratum affinity	Substratum affinity	Substratum affinity, Habitat, Habitat preference/type/specifity/complexity, Preferred substrate, Substrate type, Living location

One common approach to use traits is as indicators of ecosystem functions (effect traits) or of changes in the environment (response traits) (Hooper et al., 2005). An overview of how each of the 19 traits that are currently included in the database may relate to ecosystem functions or respond to environmental changes or pressures is given in Table 3.

Table 3. Detailed information on the 19 biological traits currently included in the Arctic Traits Database, clustered into morphology traits (5), life history traits (3), and behavioral traits (11). For every trait and its categories, the definition as used in the Arctic Traits Database is given. Abbreviations of each category are given (e.g. S1, S2) as these are used in files downloaded from the website. The relation of the respective trait to benthic ecosystem functions or responses (i.e. its role as effect or response trait) are given via specific examples and underlying literature sources are displayed.

MORPHOLOGY

Body size			
Definition	Maximum body size as adult given in mm, as individual or colony and excluding appendages. Can be height in rather upright animals (e.g. corals), body width or diameter in rather round animals (e.g. crabs), or body length in elongated animals (e.g. worms).		
Categories	S1	small	< 10 mm
	S2	small-medium	10-50 mm
	S3	medium	50-100 mm
	S4	medium-large	100-300 mm
	S5	large	> 300 mm

Function	Size has a direct effect on productivity, the amount of habitat structuring and facilitation, and is important for the amount of oxygen and nutrient flux across the sediment-water interface. It correlates with food web structure, trophic levels, and energy flow in ecosystems.		
Detail	Smaller animals are faster growing, usually show a higher productivity and are less affected by trawling as they are more likely to fit through the net of trawling gear, thus often replacing larger slow-growing fauna in trawl-impacted areas. A clear majority of small-bodied species may be indicative for environments with high instability or be the result of environmental or anthropogenic disturbances. Larger taxa usually show a lower productivity but higher carbon fixation and have a higher effect on fluxes of nutrients, energy and matter. They usually grow slower, reproduce later, and are more affected by trawling and other disturbances.		
References	Bolam and Eggleton, 2014; Bremner, 2008; Costello et al., 2015; Emmerson, 2012; Micheli and Halpern, 2005; Norkko et al., 2013; van der Linden et al., 2016		

Body form			
Definition	The external characteristic of an organism.		
Categories	BF1	globulose	Round or oval (e.g. sea urchin, sponge, some bivalves)
	BF2	vermiform	Wormlike
	BF3	dorso-ventral compressed	Species that are flat, or encrusting (e.g. starfish, sponge)
	BF4	laterally compressed	Thin (e.g. isopods, amphipods, some bivalves)
	BF5	upright	E.g. coral, basket star, sponge
Function	The body form can be indicative for the ecological role of species in an ecosystem (e.g. if it is habitat-forming), and for its vulnerability to mechanical disturbances (e.g. bottom trawling). Species with an upright body form will be more affected than vermiform or flat ones. Sets restrictions to habitat use and migration capability. Vermiform taxa can be a proxy for litter quality/decomposition.		
Remark	Often simply a proxy of taxonomy (e.g. vermiform > polychaetes, laterally compressed > amphipods).		
References	Beauchard et al., 2017; Bolam and Eggleton, 2014; Costello et al., 2015; Törnroos and Bonsdorff, 2012; Wiedmann et al., 2014		

125

Fragility			
Definition	The degree to which an organism can withstand physical impact.		
	F1	fragile	Likely to crush, break, or crack as a result of physical impact (e.g. brittle star, soft worms, smaller crustaceans, mollusks with thin shells)
	F2	intermediate	Liable to suffer minor damage, chips or cracks as result of physical impacts (e.g. mollusks with thicker shells, animals with harder cuticle like some echinoderms)
	F3	robust	Unlikely to be damaged as a result of physical impacts, e.g. hard or tough enough to withstand impact, or leathery or wiry enough to resist impact (e.g. starfish, sponges, tunicates)
Function	Determines sensitivity to physical disturbance (e.g. bottom trawling) and to predatory aggression. Softer/fragile bodies are stronger affected by trawling. Indicative for prey accessibility and ease of ingestion.		
References	Beauchard et al., 2017; Bolam and Eggleton, 2014; Weigel et al., 2016		

Skeleton			
Definition	Presence and type of supporting structures in the animal body.		
Categories	SK1	calcareous	Skeleton material aragonite or calcite (e.g. bivalves)
	SK2	siliceous	Skeleton material silicate (e.g. siliceous sponges)
	SK3	chitinous	Skeleton material chitin (e.g. arthropods)
	SK4	cuticle	No skeleton but a protective structure like a cuticle (e.g. sea-squirts)
	SK5	none	No form of protective structure (e.g. sea slugs)
Function	Indicates vulnerability (trawling, ocean acidification), resistance to predation (proxy of palatability), and ecosystem engineering (provision of habitat, increased heterogeneity). Large calcifying taxa contribute most to inorganic carbon sequestration.		
References	Costello et al., 2015; Frid and Caswell, 2016, 2015; Spitz et al., 2014		

Sociability			
Definition	The degree to which species aggregate.		
Categories	SO1	solitary	Single individual
	SO2	gregarious	Single individuals forming groups; growing in clusters (e.g. barnacles)
	SO3	colonial	Living in permanent colonies (e.g. stony corals, Bryozoa, Synascidia)

Function	Determines sensitivity to physical disturbance (e.g. bottom trawling) and can indicate if a species can increase habitat heterogeneity or is habitat forming. If yes, then it affects habitat creation, nursery, refuge, facilitation, and sediment oxygenation.
References	Beauchard et al., 2017; Costello et al., 2015

130 **LIFE HISTORY TRAITS**

Reproduction			
Definition	The way species reproduce, here including information about where fertilization occurs and whether propagules are released or not.		
Categories	R1	asexual	Budding and fission (e.g. sponges, cnidarians)
	R2	sexual – external	Fertilization external, eggs & sperm deposited on substrate or released into water (broadcast spawners) (e.g. echinoderms, cnidarians)
	R3	sexual – internal	Fertilization internal, but no brooding, eggs deposited on substrate, indirect or direct development (e.g. gastropods)
	R4	sexual – brooding	Fertilization internal or external, Eggs or larvae are brooded, indirect or direct development (e.g. amphipods, isopods, echinoderms)
Function	Indicates the ability of a species to disperse, become invasive, or recover from a population decline. Can indicate if carbon is transported from the benthic to the pelagic realm or stays locally bound. Animals without a planktonic stage that perform brooding and parental care might have a higher tolerance against some forms of stress (e.g. ocean acidification), but may be higher vulnerable to local disturbances (biotic or abiotic).		
References	Bremner, 2008; Costello et al., 2015; Lucey et al., 2015		

Larval development			
Definition	Larval development and feeding type.		
Categories	LD1	pelagic/planktotrophic	High fecundity, larvae feed and grow in water column, generally pelagic for several weeks (e.g. echinoderms, bivalves)
	LD2	pelagic/lecitotrophic	Medium fecundity, larvae with yolk sac, pelagic for short periods (e.g. tunicates)
	LD3	benthic/direct	Larvae have benthic or direct development (no larval stage, eggs develop into miniature adults)
Function	Ability of a species to disperse, become invasive, or recover from a population decline. Indicator for long-term sensitivity (ability to recolonize disturbed areas). Planktonic stages indicate productivity and elemental transport from benthos to pelagos.		
References	Bolam and Eggleton, 2014; Cardeccia et al., 2018; Törnroos and Bonsdorff, 2012		

Life span			
Definition	The maximum reported life span of the adult stage in years.		
Categories	A1	short	<2 years
	A2	medium	2-5 years
	A3	medium-long	5-20 years
	A4	long	>20 years
Function	Long lived animals are more susceptible to disturbance and need longer to recover (while short-lived species can recover fast and may increase in richness and abundance as disturbance increases). An indicator for population stability over time, carbon fixation, productivity.		
Detail	Indicates the relative investment of energy in somatic rather than reproductive growth and the relative age of sexual maturity. A proxy for relative r- and k-strategy.		
References	Bolam and Eggleton, 2014; Bremner, 2008; Cain et al., 2014; Costello et al., 2015		

135 **BEHAVIORAL TRAITS**

Environmental position			
Definition	The position of the animal relative to the sediment.		
Category	EP1	infauna	Lives in the sediment
	EP2	epibenthic	Lives on the surface of the seabed
	EP3	hyper-benthic	Living in the water column, but (primarily/occasionally) feeds on the bottom; bentho-pelagic
Function	Affects carbon fixation and transport within the sediment, between aerobic and anaerobic layers, or from pelagos to benthos. Can indicate facilitation (e.g. for microbial communities in the sediment) and sensitivity to perturbation (e.g. bottom trawling, infauna less affected than epifauna,		

			hyper-benthic taxa might be able to escape). Endobenthic life style effects the sediment biogeochemistry. Epibenthic and shallow sediment-dwelling taxa are more vulnerable to predation. Hyper-benthic taxa are involved in transport of carbon from benthos to pelagos.
References			Bolam et al., 2014; Bremner et al., 2008; Frid and Caswell, 2016; Törnroos & Bonsdorff, 2012

Living habit

Definition			The mode of living, ranging from free over tube or burrow dwelling to permanently attached.
Categories	LH1	free living	Not limited to any restrictive structure at any time. Able to move freely within and/or on the sediments
	LH2	crevice dwelling	Adults are typically cryptic, inhabiting spaces made available by coarse/rock substrate and/or biogenic species or algal holdfasts
	LH3	tube dwelling	Tube may be lined with sand, mucus or calcium carbonate, tube can also be in a burrow
	LH4	burrowing	Species inhabiting permanent or temporary burrows in the sediment, or are just burrowing in the sediment
	LH5	epi/endo zoic/phytic	Living on or in other organisms
	LH6	attached	Adherent to a substratum
Function			Attached species are more vulnerable to predation and perturbations (e.g. bottom trawling). Burrowing, crevice and tube dwelling taxa affect sediment biogeochemistry, carbon transport, elemental cycling, and are less affected by strong hydrodynamic disturbance, anoxic conditions and water pollution. Tube building can add to local storage of chemicals and waste materials. Microbial processes are facilitated and microbial biomass promoted by deep-dwelling fauna. Burrowing and irrigation generally facilitates life of associates. Burrowing or attached living habit can be related to habitat creation and facilitation.
References			Aller, 1983; Bolam and Eggleton, 2014; Bremner, 2008; Bremner et al., 2006; Costello et al., 2015; Törnroos and Bonsdorff, 2012; van der Linden et al., 2016

Mobility

Definition			Degree or intensity of movement.
Categories	MO1	none	No movement as adult (sponge, coral)
	MO2	low	Slow movement (e.g. anemones, snails)
	MO3	medium	Medium movement (e.g. starfish, brittle stars)
	MO4	high	High movement, swimmer or fast crawler (e.g. amphipods, shrimp)
Function			Slowly or non-moving species are more vulnerable to predation, perturbations and decrease in food input, while mobile taxa are more flexible and may evade trawl gear or predators. High percentage of non-moving organisms can indicate high amount of food, while high percentage of highly mobile taxa may indicate food patchiness or scarcity. Indicative for dispersal potential and ability to recolonize.
References			Costello et al., 2015; Micheli and Halpern, 2005; Tyler et al., 2012

Adult movement

Definition			Type of movement as an adult.
Categories	MV1	sessile/none	No movement as adult (sponge, coral)
	MV2	burrower	Movement in the sediment (e.g. annelids, echinoderms, crustaceans, bivalves)
	MV3	crawler	An organism that moves along on the substratum via movements of its legs, appendages or muscles (e.g. crabs, snails)
	MV4	swimmer (facultative)	Movement above the sediment (e.g. amphipods)
Function			Indicates the dispersal and recolonization potential, and the invasiveness of an organism. Related to nutrient cycling (burrowing taxa contribute most to nutrient cycling and regeneration, burrows increase the total sediment surface area available for exchange with the water column), carbon deposition (sessile calcifying taxa), facilitation of microbial and other fauna (either via burrowing or via constructing biogenic habitats), and habitat stability. Swimmers may escape predators and trawling gear.
Remark			Closely linked to the trait mobility.
References			Aller, 1983; Bremner, 2008; Bremner et al., 2006; Costello et al., 2015; Frid and Caswell, 2016

Feeding habit

Definition			The mode of food uptake.
Categories	FH1	surface deposit feeder	Active removal of detrital material from the sediment surface. Includes species which scrape and/or graze algal matter from surfaces
	FH2	subsurface deposit feeder	Removal of detrital material from within the sediment matrix (e.g. Echinocardium)

	FH3	filter/suspension feeder	Sponge, coral, hydrozoa, bivalves
	FH4	opportunist/scavenger	An organism that can use different types of food sources/an organism that feeds on dead organic material (e.g. crabs, whelks)
	FH5	predator	An organism that feeds by preying on other organisms (e.g. starfish)
	FH6	parasite/commensal	An organism that lives in or on another living organism (the host), from which it obtains food and other requirements
Function	Can indicate hydrodynamic conditions (suspension feeders in turbulent, deposit feeders in calmer water), carbon transport between pelagos and benthos (suspension feeders) and backwards (predators), and vulnerability (e.g. surface deposit feeders and suspension feeders are more sensitive to trawling). Impacts resource utilization and facilitation (e.g. deposit feeders facilitate microbes that further decompose organic carbon). Effects the depth of oxygen and detritus penetration and can enhance organic matter decomposition and nutrient recycling/regeneration. Control of other species in the assemblage.		
References	Bremner, 2008; Bremner et al., 2006; Dolbeth et al., 2009; Frid et al., 2008; Kröncke, 1994; Oug et al., 2012; Rosenberg, 1995; Tyler et al., 2012; van der Linden et al., 2016		

140

Trophic level

Definition	Rank of an animal according to how many steps it is above the primary producers at the base of the food web.		
Categories	TL1	1	Primary producer
	TL2	2	Primary consumers – Herbivore / Deposit Feeder /Suspension Feeder
	TL3	3	Secondary consumers – Carnivore
	TL4	4	Tertiary consumers
	TL5	5	Quaternary consumers – Apex predator
Function	Determines the role of an organism in energy transfer within the food web. Control of other species abundance in the assemblage.		
References	Costello et al., 2015; Micheli and Halpern, 2005; Renaud et al., 2011		

Bioturbation

Definition	Biogenic modification of sediments through living, movement and feeding habits of organisms.		
Categories	B1	diffusive mixing	Surficial movement of sediment and/or particles, resulting from movement or feeding activities on the surface
	B2	surface deposition	Deposition of particles at the sediment surface resulting from e.g. defecation or egestion (pseudofaeces) by, for example, surface deposit feeding organisms (e.g. holothuroids, bivalves, tubiculous polychaetes)
	B3	conveyor belt transport (upward)	Translocation of sediment and/or particulates from depth within the sediment to the surface during subsurface deposit feeding or burrow excavation
	B4	downward (reverse) conveyor	The subduction of particles from the surface to some depth by feeding or defecation
	B5	none	No bioturbation (e.g. sessile animals on hoard bottom)
Function	Impacts sediment biogeochemistry (oxygen, pH and redox gradients, elemental carbon), organic matter regeneration, nutrient cycling, sediment granulometry, pollutant release, microbial composition, abundance and diversity and in general provision and maintenance of habitats for other organisms.		
References	Chen et al., 2017; Frid et al., 2008; Gogina et al., 2017; Lacoste et al., 2018; Mermillod-Blondin, 2011; Pearson, 2001; Queirós et al., 2013; Solan et al., 2012		

Tolerance

Definition	Degree to which a species reacts to changes in its environment.		
Categories	T1	low	Species reacts sensitive to changes in the environment like organic enrichment, pollution, temperature or salinity changes; AMBI group I
	T2	intermediate	Species react indifferent or no information available; AMBI group II
	T3	high	Species tolerates organic enrichments, pollution, temperature or salinity changes; AMBI groups III-IV
Function	Indicates vulnerability or resistance/resilience of a species towards pollution or climate change induced changes in water biogeochemistry.		
References	Borja and Franco, 2000; Gusmao, 2017; Marchini et al., 2008; Piló et al., 2016		

Zoogeography			
Definition	Spatial distribution of a species in relation to commonly used zoogeographic regions.		
Categories	Z1	arctic	Confined to Arctic regions.
	Z2	arctic-boreal	Arctic, sub-Arctic and North Atlantic/North Pacific distribution.
	Z3	boreal	North Atlantic and/or North Pacific distribution; potentially sub-Arctic regions such as Southern Barents Sea or Bering Sea.
	Z4	cosmopolite	Cosmopolite distribution
Function	Indicates vulnerability (arctic species may be more vulnerable to changes than species with an arctic-boreal or cosmopolite distribution) or potential of a species to become invasive.		
References	Fetzer, 2005; Fetzer and Arntz, 2008; Piepenburg, 2000; Weslawski et al., 2003		

Depth range			
Definition	Species distribution related to water depth.		
Categories	DR1	shallow	0-20 m
	DR2	shelf	20-200 m (some shelves can extend to 500 m)
	DR3	shelf-slope	200-1000 m (sometimes the slope starts deeper, e.g. 500-)
	DR4	slope-basin	> 1000 m
Function	Can be used – along substratum affinity – for habitat classification. Can depict depth distribution of other traits.		
Detail	Shallow water and shelf taxa face a higher exposure to predation of marine mammals and to physical disturbance such as iceberg scouring and to coastal processes and pollution.		
References	Costello et al., 2015; Gutt, 2001		

145

Substratum affinity			
Definition	Type of substratum that organisms (preferential) live on.		
Categories	SA1	soft	Soft substrata, sand or mud
	SA2	hard	Hard substrata, rock, gravel
	SA3	biological	Epizoic or epiphytic life style
	SA4	none	Species is hyper/supra benthic and has no affinity for a certain substrate, but it might prefer one for hunting/scavenging (this category should not occur too often, as we work with benthos)
Function	Can be used – along depth range – for habitat classification. Can depict potential substrate specificity of other traits.		
References	Costello et al., 2015		

2.3 Sources of trait information

Sources of trait information are research papers, books, databases and online repositories (Table 1), but also grey literature such as cruise reports. Trait information can also result from onsite measurements (e.g. for the trait body size), personal observations, or be transmitted via communication with experts for a specific taxonomic group. In

150 any case, the source is indicated as precise as possible, for published literature with complete reference and DOI (if available), in case of expert communication the name and contact details of the respective expert are given. Wherever possible the original quote from literature and page numbers are given to ensure the traceability of the provided trait information. Although literature sources targeting the Arctic are used preferably (and for exclusively Arctic species are the only option) we do not restrict source information for arctic-boreal or cosmopolite taxa to

155 stem from Arctic regions. This bears the risk that the assigned trait information is not accurate, as polar taxa might differ in their expression of certain traits from their relatives at lower latitudes (Degen et al. 2018). However, this is an issue for now not resolved, as trait information from the high latitudes is often scarce, and we recommend the user to consider the source of trait information when interpreting results.

2.4 Fuzzy coding of traits

160 The fuzzy coding procedure indicates to which extent a taxon exhibits each trait category (Chevenet et al., 1994). This method has the advantage that it enables us to analyze diverse kinds of biological information derived from a variety of sources (as those included in the Arctic Traits Database, see Sect. 2.3), and that also intermediate scenarios (i.e. when a taxon does not clearly fall into one category or the other) can be accounted for (Chevenet et

al. 1994). We use the 0–3 coding scheme (details in Table 4 below) as it is the most widely used (which facilitates comparisons and exchange of trait information) and provides a compromise between binary codes and many not clearly delineated graduations (Degen et al. 2018).

Table 4. Explanation of fuzzy codes as used in the Arctic Traits Database.

Fuzzy code	Explanation
3	Taxon has total and exclusive affinity for a certain trait category, all other categories do not apply and must be coded with "0".
2	Taxon has a high affinity for a certain trait category, but other categories can occur with equal (2) or lower (1) affinity.
1	Taxon has a low affinity for a certain trait category.
0	Taxon has no affinity for a certain trait category.

Table 5. Two coding examples for the trait "Feeding habit" which has six trait categories (FH1 – FH6, see also Table 3). Species 1 is a surface deposit feeder, but can switch facultative to suspension feeding, while species 2 is an exclusive suspension feeder.

Feeding habit	Abbreviation	Species 1	Species 2
Surface deposit feeder	FH1	2	0
Subsurface deposit feeder	FH2	0	0
Filter/suspension feeder	FH3	1	3
Opportunist/scavenger	FH4	0	0
Predator	FH5	0	0
Parasite/commensal	FH6	0	0

Table 6. This is how the above example would appear in the matrix downloaded from the Arctic Traits Database. In the download matrix format species are rows, trait categories are columns, and the fuzzy codes are the values. Due to the database structure zero codes ("0") are only displayed when they are backed up by a specific reference (e.g. for the trait category LH3/tube dwelling: "No species within the family Polynoidae is tubiculous").

	FH1	FH2	FH3	FH4	FH5	FH6
Species 1	2		1			
Species 2			3			

While the coding might for some traits and taxa be pretty straight forward, in some cases a decision might be drawn not so easily. As one of the clearer cases, we point out the coding of the trait "body size" for the star fish *Crossaster papposus*. A literature reference states that the body size can range "Up to 340 mm in diameter" (Hayward and Ryland, 2012, p. 668). This size fits into the category "large" (S5, > 300 mm), thus the taxon is coded "3" for this size class, and "0" for all other categories (S1 – S4). The trait "mobility" is trickier. A literature reference (Himmelman and Dutil, 1991), p. 68) states the following: "*Crossaster papposus* and *Solaster endeca* are highly mobile; large individuals can cover distances of more than 5 meters in 12 hours". Here we have to keep in mind that the particular reference frame in this publication are subtidal sea stars in the northern Gulf of St. Lawrence (West Atlantic). The reference of the Arctic Traits Database however are all benthic invertebrates, and the trait category "high mobility" is defined here for taxa which are "swimmers or fast crawlers", such as some amphipods and shrimp (see Table 2). Accordingly, the correct coding for *C. paposus* in the reference system of the Arctic Traits Database is the category "medium" mobility (MO3). Users of the Arctic Traits Database should bear this reference system in mind when downloading only the fuzzy coded trait data and aiming to apply it to another reference system. But as the detailed literature quote that lead to the coding of a trait is always provided (see Sect. 2.3), the trait information can easily be adjusted by the user.

There will always be a certain degree subjectivity related to the fuzzy coding procedure. To find out how strong the coding might differ among scientists a small experiment at the Arctic Traits Workshop in Vienna (December 2016) was performed (Degen et al. 2018). Participants coded 27 trait categories of three common

Arctic benthic species, and found the final trait matrices to be to 83% identical. We are confident that the sophisticated structure of the Arctic Traits Database (see Sect. 3) and the provided information and instructions will support a more consistent coding of benthic traits in the future.

3 Database

In order to collect trait information and to disseminate it among users, a web-based database was created. The database features a public interface (Sect. 3.1) and an entry interface that is accessible only for registered collaborators (Supplement). The public interface (Fig. 2, a) allows to browse the traits and references online ("Data per taxon" in the top menu bar), to view background information ("About" and "Trait definitions") and to download either the entire species, trait and literature information or specified subsets in several formats ("Download data") (see Sect. 3.1). Registered collaborators – i.e. those users that actively contribute trait information to the Arctic Traits Database – can access the interactive part of the database via the log in button on the public page (Fig. 2a). This access offers additional options (Fig. 2b): browsing the existing information also per traits ("Traits" in the top menu bar), uploading new taxa, trait and source information, or adding trait information, references and comments to already existing taxa in the database ("Taxa"). As several users can work on the same taxa, a flagging system is used to highlight and discuss potentially conflicting sources and opinions. The "References", "Statistics", and "Tools" sections are equally accessible only for registered users (Fig. 2, b; Supplement). Every scientist working in the field of Arctic benthic ecology aiming to share trait information can become a registered user by getting in touch with the editor and retrieving a user login. Credit to the registered collaborators is given in the "About" section on the public site and also on taxon pages after each trait entry they conduct. A detailed manual for registered users is provided in the supplementary material to this publication (Supplement), or can alternatively be accessed via the public web interface ("About"). Collaborators who want to share trait information without registering to the database can alternatively be provided with an upload template (.xls).

Figure 2. Screenshots of the start page of the Arctic Traits Database. Toolbar of the public page with Login button for the registered user (a), and toolbar in the area for registered users (b).

3.1 Public access and download options

The public access enables to browse the database online and to download the complete set of data as well as the bibliography, or specified subsets. Taxon traits can be visually inspected online via the "Data per taxon" button from the top menu bar and "Browse taxa" or "Search taxa". Taxa can be browsed and selected via the taxonomic tree, as indicated for the asteroid *Crossaster papposus* in Fig. 3. Alternatively, the "Search taxa" panel allows to type in and search a specific taxon.

Figure 3. Screenshot of the taxon page of the asteroid *Crossaster papposus* selected from the classification tree on the left.

The completeness of trait information can be inspected via "Data completeness" (Fig. 4), equally accessible via "Data per taxon" on the top menu bar. This option shows an alphabetic list of all taxa in the database for which

230 trait information is available. The bar on the right side indicates the information coverage for each taxon and trait, blue color indicates that trait information is present.

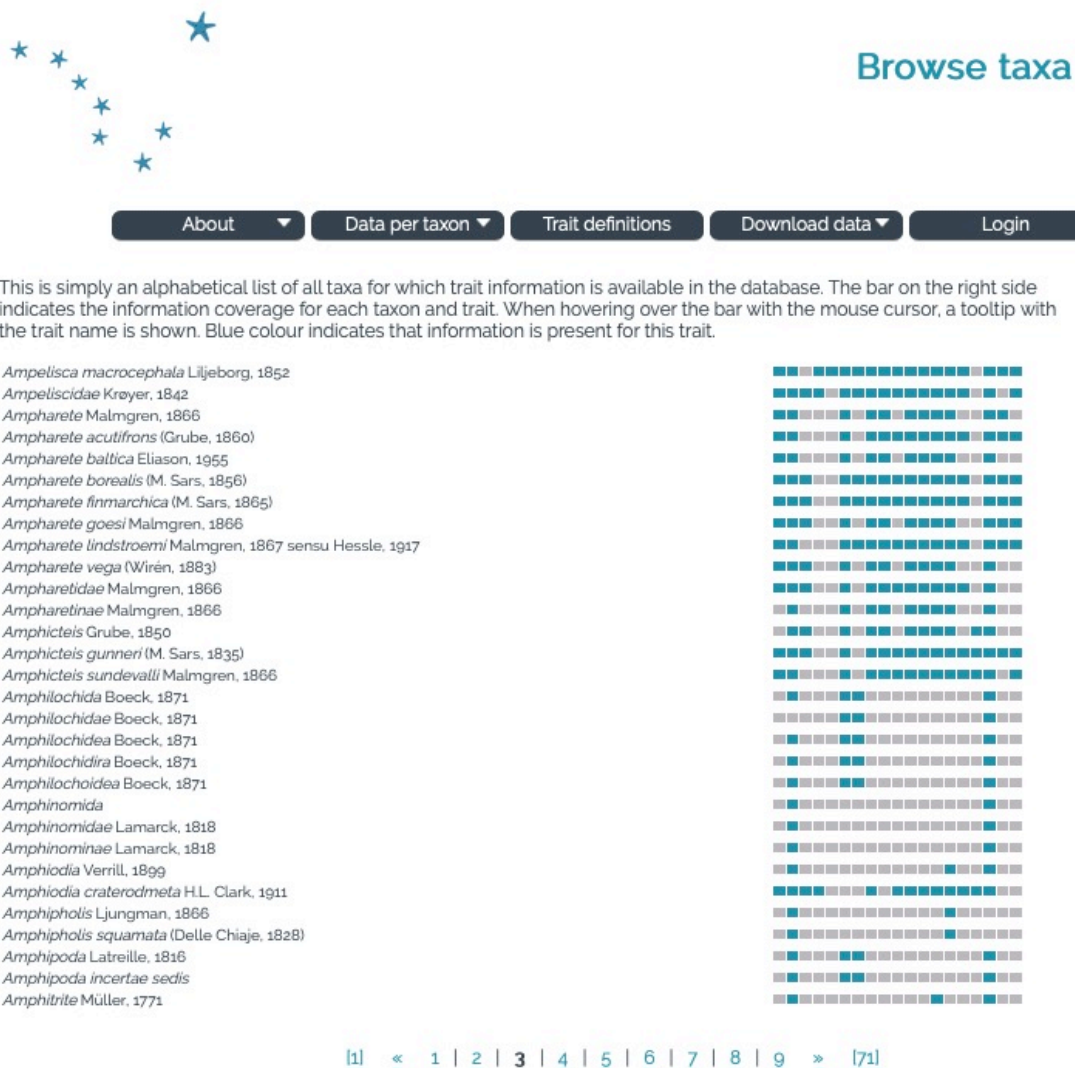



Figure 4. Screenshot of data completeness.

The download section can be accessed via the "Download data" button on the top menu bar (Fig. 2, a; Fig. 3; Fig. 4). Download is enabled in three different computer readable formats: 1) as data in columns (*.csv) (Table 7), 2) in DarwinCore format (Table 8), and 3) as fuzzy coded trait matrix which some users might prefer (see Sect. 2.4 and Fig. 5). Also, the entire bibliography is available for download. Before the download commences the user is asked whether to download a) all data in the database, b) only data for an uploaded list of taxon names, c) only data for an uploaded list of AphiaIDs, or d) only the data selected from a classification tree. In the last option, entire phyla or sub-groups can be easily selected from the tree. By default, all 19 traits are exported, but if the user is interested only in one or a few specific traits, the option to select these from the total list of 19 traits is available. As the fuzzy coded trait matrix (download option 3) contains only the fuzzy codes per trait category but no literature sources, we recommend to also download the "Data in columns" (download option 1) for the same taxa, where the detailed source per species and trait category is included. Details on the structure of the first two download options are given below in Table 7 and Table 8. A clipping from a downloaded fuzzy coded trait matrix is shown in Fig. 5. The database can also be accessed programmatically via a REST API (documented at https://www.univie.ac.at/arctictraits/download-api).

Table 7. List of fields returned by the Arctic Traits Database when "Data as columns" (*.csv) is chosen as an export option from the download section.

Column label	Column description
Taxon	The taxon for which the information was recorded.
Author	The author and year of the *Taxon* for which the information was recorded.
Rank	Rank of the taxon for which the information was recorded.
Valid taxon	Currently accepted name of the *Taxon* (as stored in the Arctic Traits Database - information might not be up to date with the WoRMS or the latest taxonomic literature in some cases). Users should check all taxa against WoRMS before use. If *Taxon* is currently accepted, this field contains the same value as *Taxon*).
Valid author	Currently accepted name of the *Author* (as stored in the Arctic Traits Database - information might not be up to date with the WoRMS or the latest taxonomic literature in some cases). Users should check all taxa against WoRMS before use. If *Taxon* is currently accepted, this field contains the same value as *Author*.
Taxonomic status	The status of the use of the Taxon (e.g. objective synonym, subjective synonym) as stored in the Arctic Traits database.
Source of synonymy	Literature reference for synonymy of taxon (if present).
Parent taxon	The *Taxon*'s direct parent in the taxonomic classification (as stored in the Arctic Traits Database).
Trait	The biological trait for which information is available (e.g. "Feeding habit").
Category	The sub-category of the *Trait* for which information is available (e.g. "Predator").
Category abbreviation	An abbreviated version of the often verbose trait category - useful as a label in further analyses of the data (e.g. "FH(6)").
Traitvalue	Describes the affinity of the *Taxon* to the *Category*. Values range from 0–3: "0"= no affinity for a certain trait category; "1"= low affinity for a certain trait category; "2"= high affinity for a certain trait category, but other categories can occur with equal (2) or lower (1) affinity; "3"= total and exclusive affinity for a certain trait category.
Reference	Literature reference leading to the assignment of the *Traitvalue* to the *Category* for the *Taxon*.
DOI	Digital Object Identifier (where available) of the *Reference*.
Value creator	Person who assigned the *Traitvalue* to the *Category* for the *Taxon*, supported by a *Reference*.
Value creation date	Date and time when the above information was entered into the database.
Value modified by	Person who last modified the Traitvalue. Empty if no modifications were done.
Value modification date	Date and time when the Traitvalue was last modified. If no modification was done since the first entry, this has the same value as Value creation date.
Text Excerpt	A quotation of the original text passage from the literature source that led to the assignment of assignment of the *Category*/*Traitvalue* to the *Taxon*. Empty if information has not been recorded yet.
Text Excerpt creator	Person who entered the *Text excerpt*. Only present if *Text Excerpt* is present.
Text Excerpt creation date	Date and time when the *Text Excerpt* was entered into the database. Only present if *Text Excerpt* is present.
Text Excerpt modified by	Person who last modified the *Text excerpt*. Empty if no modifications were done.
Text Excerpt modification date	Date and time when the *Text Excerpt* was last modified. If no modification was done since the first entry, this has the same value as *Text Excerpt creation date*.

250 **Table 8.** List of fields returned by the Arctic Traits Database when "Darwin Core" is chosen as an export option from the download section. DarwinCore does not provide the same granularity as the "Data as columns" format. The output file consequently contains fewer details.

Column label	Column description
scientificName	The taxon for which the information was recorded
scientificNameAuthorship	The author and year of the taxon for which the information was recorded
taxonRank	Rank of the taxon for which the information was recorded.
acceptedNameUsage	Currently accepted name and authorship of the *scientificName* (as stored in the *arctictraits* database – information might not be up to date with the latest taxonomic literature in some cases.)
Taxonomic Status	The status of the use of the *scientificName* (e.g. objective synonym, subjective synonym) as stored in the *arctictraits* database. Empty if *scientificName* is the currently accepted name.
MeasurementOrFact	Trait name and trait category, separated by a colon (e.g. Size:small)
measurementValue	Value from 0–3, describing the affinity of the taxon to a trait category. Coding of values as described in Table 7 "Traitvalue".
dcterms:bibliographicCitation	Full literature reference (including Digital Object Identifier (DOI) where present) supporting the trait information for the current taxon.

measurementRemarks	A quotation of the original text passage containing the trait information for the current taxon
measurementDeterminedBy	Person who entered the trait information for this taxon into the database.
measurementDeterminedDate	Date the trait information was entered into the database or last modified.

#	Taxon	Valid_name	S1	S2	S3	S4	S5	W1	W2	W3	W4	W5	BF1	BF2	BF3	BF4	BF5	SK1	SK2	SK3	SK4	SK5	F1	F2	F3	SO1	SO2	SO3	R1	R2	R3	R4	LD1
1	Taxon	Valid_name																															
4	Acanthonotozoma	Acanthonotozoma														3				3												3	
5	Acanthonotozomatidae	Acanthonotozomatidae														3				3												3	
6	Acanthostepheia	Acanthostepheia														3				3												3	
7	Acanthostepheia malmgreni	Acanthostepheia malmgreni														3				3												3	
8	Aceroides	Aceroides														3				3												3	
9	Aceroides (Aceroides)	Aceroides (Aceroides)														3				3												3	
10	Aceroides (Aceroides) latipes	Aceroides (Aceroides) latipes	3													3				3	3		3									3	
11	Acmaeidae	Acmaeidae													3																		
12	Actiniaria	Actiniaria																			1	2			3								
13	Adapedonta	Adapedonta														3																3		3
14	Admete	Admete														3																		
15	Admete viridula	Admete viridula														3																		
16	Admetinae	Admetinae														3																		
17	Aglaophamus	Aglaophamus										3								3											2			3
18	Aglaophamus malmgreni	Aglaophamus malmgreni			3							3								3	3										3			3
19	Akanthophoreidae	Akanthophoreidae																	3														3	2
20	Akanthophoreus	Akanthophoreus																	3														3	2
21	Akanthophoreus gracilis	Akanthophoreus gracilis																	3														3	2
22	Alcyonidiidae	Alcyonidiidae																										3					
23	Alcyonidiina	Alcyonidiina																										3					
24	Alcyonidioidea	Alcyonidioidea																										3					
25	Alcyonidium	Alcyonidium																										3					
26	Alcyonidium gelatinosum	Alcyonidium gelatinosum	3												3													3	1		2		
27	Allantactis	Allantactis																															

Figure 5. A clipping from the fuzzy coded trait matrix returned by the Arctic Traits Database when the "Data in matrix format" is chosen as export option from the download section. Species are rows ("Valid_name" refers to the currently accepted taxonomy in WoRMS), abbreviated trait categories are columns. For abbreviations of trait categories see Table 3. Due to the database structure zero codes ("0") are not displayed (see Table 6).

3.3 Database specification

The website runs on an Apache 2.2. server, the database is implemented in MySQL 5. PHP 5 is used as a scripting language. Web technologies used are HTML4, CSS and JavaScript/Jquery. A code package to create such a web-based trait database including a README file with instructions for installation is provided at figshare, https://doi.org/10.6084/m9.figshare.7491869.

4 Results

4.1 Taxonomic data coverage

At present, the database contains 1911 Arctic marine benthic invertebrate taxa. Thereof 686 are on species level, 516 on genus level, and 274 on family level. The remaining 435 taxa are higher taxonomic levels or intermediate ranks. The largest taxonomic group in the database at present stage are the Arthropoda with 557 taxa (186 entries on species level), followed by the Annelida with 489 taxa (218 entries on species level) and the Mollusca with 418 taxa (146 entries on species level) (Fig. 6).

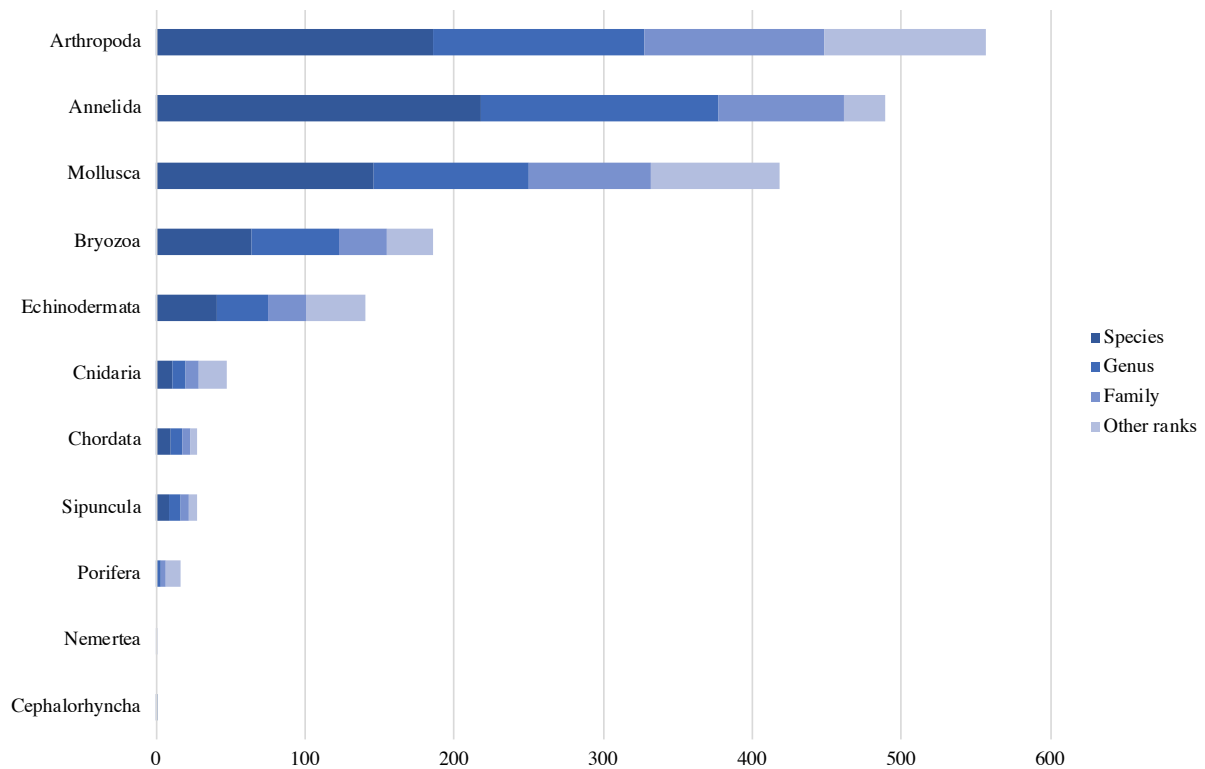



Figure 6. Taxonomic data coverage. "Other ranks" include higher taxonomic levels and intermediate ranks.

4.2 Trait data coverage

At present, the database contains 19 traits and 80 trait categories with in total currently 14242 entries of trait information. The trait for which most entries exist is "Skeleton" (1837 entries), followed by "Reproduction" (1328 entries) and "Body form" (1151 entries) (Fig. 7). The phylum with most entries are the Annelida (6130 entries, 43 %), followed by Arthropoda (2968 entries, 21 %) and Mollusca (2177 entries, 15 %). Regarding the taxonomic level, most trait information was added on the species level (48 %), less on the genus (25 %) and family level (17 %).

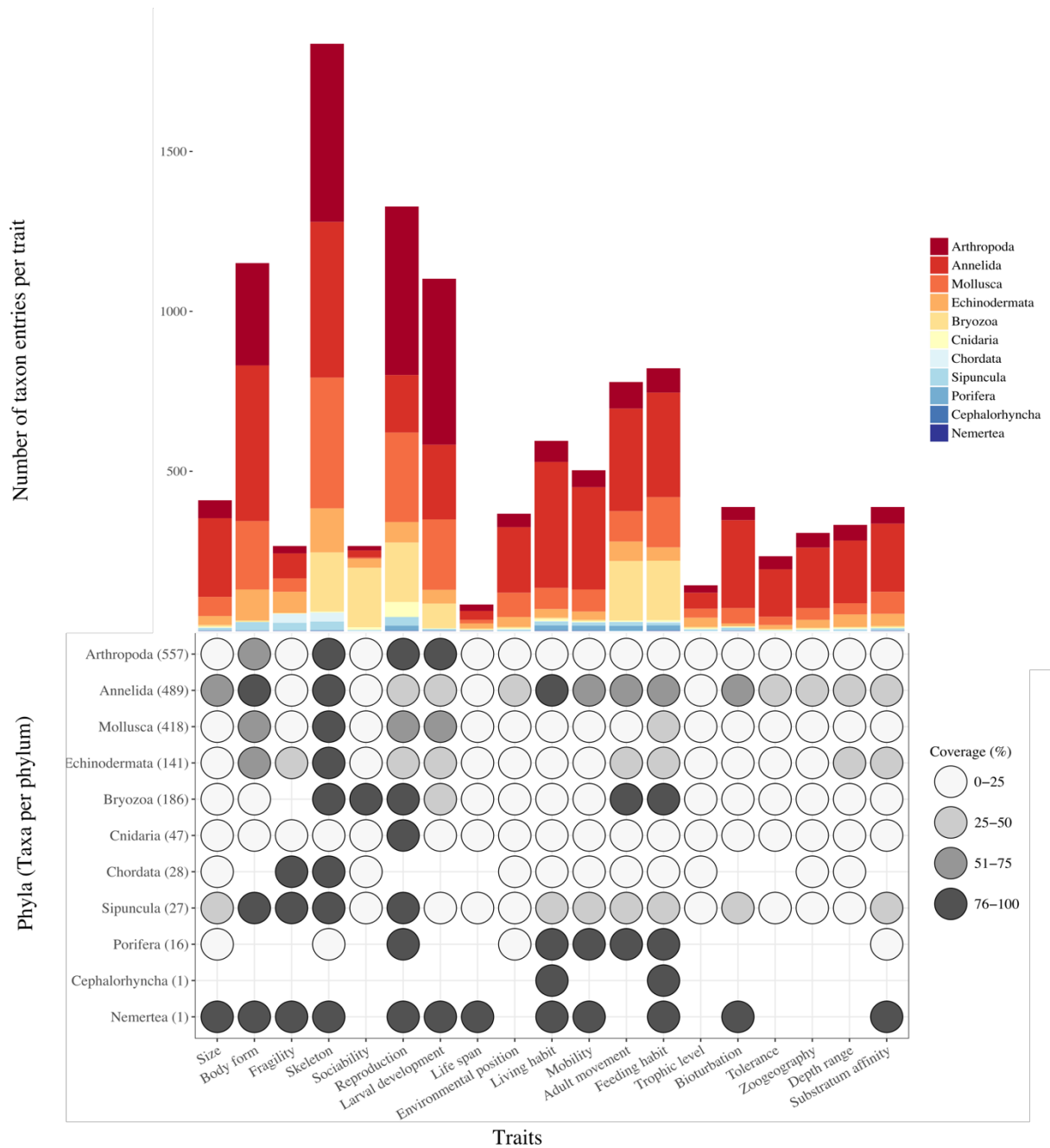
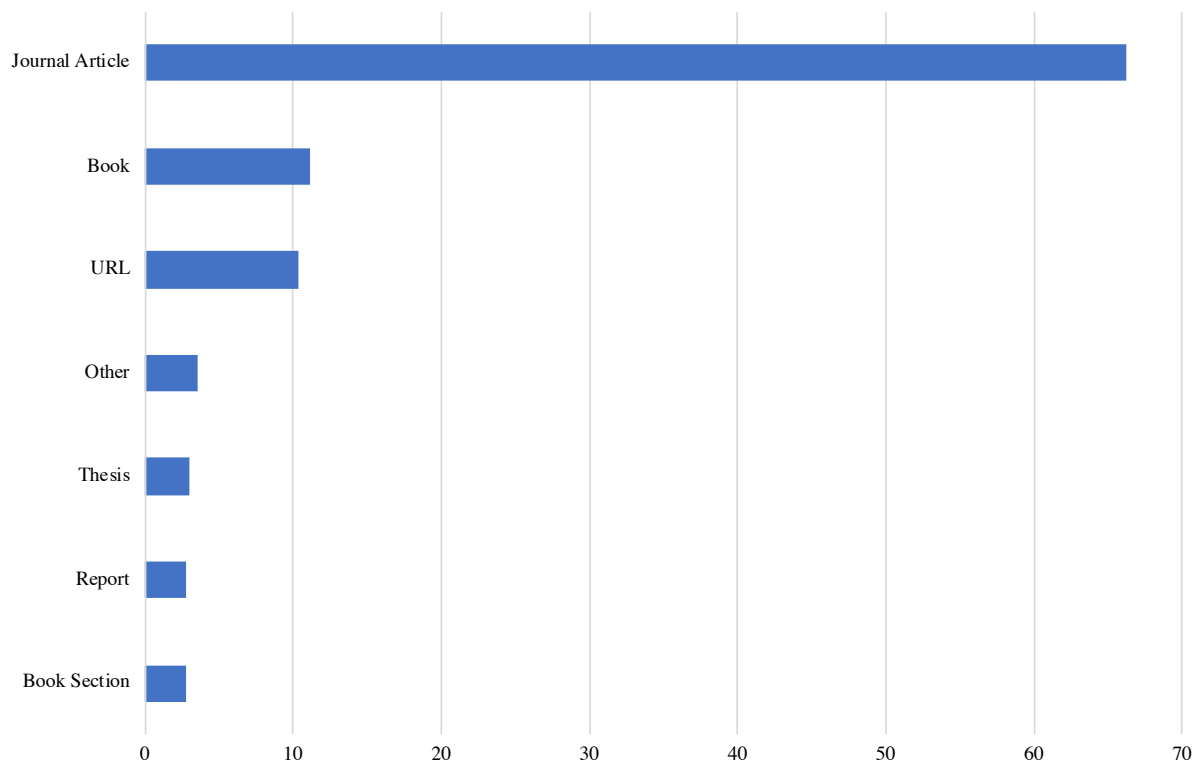



Figure 7. Scheme visualizing the taxon entries per trait (bar chart), the number of taxa per phylum (brackets), and the data coverage per trait per phylum (dot plot).

4.3 Bibliography

285   The Arctic Traits Database currently includes 394 sources of trait information. Thereof 66 % scientific papers, 11 % are books, 10 % webpages, and 4 % are expert communications and personal observation ("Other"). Theses, book sections, and reports each make up around 3 %. Most sources were used for the phylum Echinodermata and Annelida (33 % each), followed by Arthropoda (29 %).

 Figure 8. Relative amount (%) of trait source types.

5 Discussion

Although the Arctic Traits Database is still growing as new taxa and trait information are added, certain trends in data completeness or scarceness, respectively, became apparent (Fig. 7). Thus, the database is not only a valuable tool for collecting and providing information, but also for pointing out where more research might be needed. Regarding the 19 traits included at the present stage, it shows that our knowledge on e.g. the live span of many Arctic benthic species is still limited (information only for < 5 % of species). This lack of data on species longevity is astonishing, as polar taxa are traditionally depicted as slow growing and long-lived compared to their relatives from lower latitudes. Accordingly, one might have expected that more studies and measurements are available for a variety of Arctic taxa, which is not the case for many groups. Other traits that are currently underrepresented are trophic level (< 8 %) and tolerance (<13 %).

Regarding our interest to identify knowledge gaps, a special strength of the database is the implemented flagging system (described in detail in the supplement). As registered users continue to upload trait information, also more "conflicts" – i.e. cases where the sources or observations added by different users point towards different trait categories – may arise. Such cases are then indicated by a red flag and can be easily filtered for. Monitoring and statistical evaluation of these cases will grant important information on where conflicts exist and for which taxa or traits future research is needed. Such evaluation will also aid to identify which traits are more robust (i.e. are never flagged), and which show a higher plasticity (frequent flagging). This kind of information is of tremendous value as it can aid the choice as of which traits to include in prospective trait-based studies. Apart from clearly diverging source information, also different levels of experience or customs in fuzzy coding might lead to red flags in the system. Here the editorial team will take care for consistency by solving the conflicts according to the database standard, by that also fostering a standardized way of coding within the community. In addition,

repetitively occurring discrepancies in the coding of certain traits might also point towards a need for revision of these trait categories or their definitions, or maybe even the adding of a new trait, in that way improving the quality of the database.

315   In addition to the above discussed knowledge gaps surrounding certain traits, also the data coverage among taxonomic groups varies considerable (Fig. 7). This potentially mirrors the sampling design of the underlying datasets. Some taxonomic groups such as the polychaetes clearly dominate many benthic soft-bottom communities, while other taxa such as the shrimp/caridea are highly mobile and might be permanently undersampled with sampling gears like grabs, box corers, or bottom trawls (Eleftheriou and McIntyre, 2007). This

320   points toward the need to include also datasets derived from video and still image analysis in the future development of the database. These methods – despite certain disadvantages (discussed in Degen et al. 2018, Supplementary file 3) – have the great benefit that also traits of hard bottom communities can be analyzed, ecosystems which are at present stage underrepresented in the Arctic Traits Database.

6 Data availability

325   The Arctic Traits Database is hosted at the University of Vienna (Austria) and can be accessed via https://www.univie.ac.at/arctictraits/ (https://doi.org/10.25365/phaidra.49). A code package to create a web-based trait database including a README file with instructions for installation is provided at figshare, https://doi.org/10.6084/m9.figshare.7491869.

7 Conclusions

330   The Arctic Traits Database provides an easy accessible and sound knowledge base of traits of Arctic benthic invertebrates and will thus facilitate prospective trait-based studies for a variety of benthic ecologists at all career stages. Its sophisticated structure accounts for the most commonly raised demands to contemporary trait databases: 1) obligate traceability of information (every entry is linked to at least one source), 2) exchangeability among platforms (use of most common download formats), 3) standardization (use of most common terminology and

335   coding scheme), and last but not least 4) user friendliness (granted by an intuitive web-interface and rapid and easy download options). The combination of these aspects makes the Arctic Traits Database a cutting-edge tool for (not only) the marine realm and a role-model for prospective databases.

Author contribution

RD designed the project and performed the trait data collection. SF performed database and webpage development

340   and design. RD prepared the manuscript with contributions from SF.

Competing interests

The authors declare that they have no conflict of interest.

Acknowledgements

345   The authors wish to thank all collaborators that support the Artic Traits Project, especially Bodil Bluhm, Jackie Grebmeier, Lauren Sutton, Dieter Piepenburg, and Arny Blanchard. This work was supported by the Austrian Science Fund (FWF; T 801-B29) to RD.

References

Aller, R. C.: The importance of the diffusive permeability of animal burrow linings in determining marine
350   sediment chemistry, J. Mar. Res., 41(2), 299–322, doi:10.1357/002224083788520225, 1983.

Beauchard, O., Veríssimo, H., Queirós, A. M. and Herman, P. M. J.: The use of multiple biological traits in marine community ecology and its potential in ecological indicator development, Ecol. Indic., 76, 81–96, doi:10.1016/j.ecolind.2017.01.011, 2017.

Bernhardt-Römermann, M., Gray, A., Vanbergen, A. J., Bergès, L., Bohner, A., Brooker, R. W., Bergès, L.,
355   Bohner, A., Brooker, R. W., De Bruyn, L., De Cinti, B., Dirnböck, T., Grandin, U., Hester, A. J., Kanka, R., Klotz, S., Loucougaray, G., Lundin, L., Matteucci, G., Mészáros, I., Oláh, V., Preda, E., Prévosto, B., Pykälä, J., Schmidt, W., Taylor, M. E., Vadineanu, A., Waldmann, T. and Stadler, J.: Functional traits and local environment predict vegetation responses to disturbance: a pan-European multi-site experiment, J. Ecol., 99(3), 777–787, doi:10.1111/j.1365-2745.2011.01794.x, 2011.

360   Blanchard, A. L., Parris, C. L., Knowlton, A. L. and Wade, N. R.: Benthic ecology of the northeastern Chukchi Sea. Part I. Environmental characteristics and macrofaunal community structure, 2008-2010, Cont. Shelf Res., 67, 52–66, doi:10.1016/j.csr.2013.04.021, 2013a.

Blanchard, A. L., Parris, C. L., Knowlton, A. L. and Wade, N. R.: Benthic ecology of the northeastern Chukchi Sea. Part II. Spatial variation of megafaunal community structure, 2009–2010, Cont. Shelf Res., 67, 67–76,
365   doi:10.1016/j.csr.2013.04.031, 2013b.

Bolam, S. G. and Eggleton, J. D.: Macrofaunal production and biological traits: Spatial relationships along the UK continental shelf, J. Sea Res., 88, 47–58, doi:10.1016/j.seares.2014.01.001, 2014.

Borja, A., Franco, J. and Pérez, V.: A Marine Biotic Index to Establish the Ecological Quality of Soft-Bottom Benthos Within European Estuarine and Coastal Environments, Mar. Pollut. Bull., 40(12), 1100–1114, 2000.

370   Bremner, J.: Species' traits and ecological functioning in marine conservation and management, J. Exp. Mar. Bio. Ecol., 366(1–2), 37–47, doi:10.1016/j.jembe.2008.07.007, 2008.

Bremner, J., Rogers, S. I. and Frid, C. L. J.: Methods for describing ecological functioning of marine benthic assemblages using biological traits analysis (BTA), Ecol. Indic., 6(3), 609–622, doi:10.1016/j.ecolind.2005.08.026, 2006.

375   Brown, J. H., Gillooly, J. F., Allen, A. P., Van Savage, M. and West, G. B.: Toward a metabolic theory of ecology, Ecology, 85(7), 1771–1789, 2004.

Brun, P., Payne, M. R. and Kiørboe, T.: A trait database for marine copepods, Earth Syst. Sci. Data, 9(1), 99–113, doi:10.5194/essd-9-99-2017, 2017.

Cain, M. L., Bowman, W. D. and Hacker, S. D.: Ecology, Third Edit., Sinauer Associates., 2014.

380   Cardeccia, A., Marchini, A., Occhipinti-Ambrogi, A., Galil, B., Gollasch, S., Minchin, D., Narščius, A., Olenin, S. and Ojaveer, H.: Assessing biological invasions in European Seas: Biological traits of the most widespread non-indigenous species, Estuar. Coast. Shelf Sci., 201, 17–28, doi:10.1016/j.ecss.2016.02.014, 2018.

Chen, X., Andersen, T. J., Morono, Y., Inagaki, F., Jørgensen, B. B. and Lever, M. A.: Bioturbation as a key driver behind the dominance of Bacteria over Archaea in near-surface sediment, Sci. Rep., 7(1), 1–14, doi:10.1038/s41598-017-02295-x, 2017.

Chevenet, F., Dolédec, S. and Chessel, D.: A fuzzy coding approach for the analysis of long-term ecological data, Freshw. Biol., 31(3), 295–309, doi:10.1111/j.1365-2427.1994.tb01742.x, 1994.

Costello, M. J., Claus, S., Dekeyzer, S., Vandepitte, L., Tuama, É. Ó., Lear, D. and Tyler-Walters, H.: Biological and ecological traits of marine species., PeerJ, 3, e1201, doi:10.7717/peerj.1201, 2015.

Darr, A., Gogina, M. and Zettler, M. L.: Functional changes in benthic communities along a salinity gradient- a western Baltic case study, J. Sea Res., 85, 315–324, doi:10.1016/j.seares.2013.06.003, 2014.

Degen, R., Aune, M., Bluhm, B. A., Cassidy, C., Kędra, M., Kraan, C., Vandepitte, L., Włodarska-Kowalczuk, M., Zhulay, I., Albano, P. G., Bremner, J., Grebmeier, J. M., Link, H., Morata, N., Nordström, M. C., Shojaei, M. G., Sutton, L. and Zuschin, M.: Trait-based approaches in rapidly changing ecosystems: A roadmap to the future polar oceans, Ecol. Indic., 91(April), 722–736, doi:10.1016/j.ecolind.2018.04.050, 2018.

Dolbeth, M., Teixeira, H., Marques, J. C. and Pardal, M. Â.: Feeding guild composition of a macrobenthic subtidal community along a depth gradient, Sci. Mar., 73(2), 225–237, doi:10.3989/scimar.2009.73n2225, 2009.

Eleftheriou, A. and McIntyre, A.: Methods for the Study of Marine Benthos: Third Edition, Blackwell Science Ltd., 2007.

Emmerson, M. C.: The importance of body size, abundance, and food-web structure for ecosystem functioning, in Marine Biodiversity and Ecosystem Functioning: Frameworks, methodologies, and integration, edited by M. Solan, R. J. Aspden, and D. M. Paterson, p. 240, Oxford University Press, Oxford., 2012.

Faulwetter, S., Markantonatou, V., Pavloudi, C., Papageorgiou, N., Keklikoglou, K., Chatzinikolaou, E., Pafilis, E., Chatzigeorgiou, G., Vasileiadou, K., Dailianis, T., Fanini, L., Koulouri, P. and Arvanitidis, C.: Polytraits: A database on biological traits of marine polychaetes, Biodivers. Data J., 2, e1024, doi:10.3897/BDJ.2.e1024, 2014.

Fetzer, I.: Reproduction strategies and distribution of larvae and juveniles of benthic soft-bottm invertebrates in the Kara Sea (Russian Arctic), Universtiy of Bremen., 2005.

Fetzer, I. and Arntz, W. E.: Reproductive strategies of benthic invertebrates in the Kara Sea (Russian Arctic): Adaptation of reproduction modes to cold water, Mar. Ecol. Prog. Ser., 356(1878), 189–202, doi:10.3354/meps07271, 2008.

Foden, W. B., Butchart, S. H. M., Stuart, S. N., Vié, J. C., Akçakaya, H. R., Angulo, A., DeVantier, L. M., Gutsche, A., Turak, E., Cao, L., Donner, S. D., Katariya, V., Bernard, R., Holland, R. A., Hughes, A. F., O'Hanlon, S. E., Garnett, S. T., Şekercioğlu, Ç. H. and Mace, G. M.: Identifying the world's most climate change vulnerable species: a systematic trait-based assessment of all birds, amphibians and corals, edited by S. Lavergne, PLoS One, 8(6), e65427, doi:10.1371/journal.pone.0065427, 2013.

Frid, C. L. J. and Caswell, B. A.: Is long-term ecological functioning stable: The case of the marine benthos?, J. Sea Res., 98, 15–23, doi:10.1016/j.seares.2014.08.003, 2015.

Frid, C. L. J. and Caswell, B. A.: Does ecological redundancy maintain functioning of marine benthos on centennial to millennial time scales ?, Mar. Ecol., 37(2), 392–410, doi:10.1111/maec.12297, 2016.

Frid, C. L. J., Paramor, O. A. L., Brockington, S. and Bremner, J.: Incorporating ecological functioning into the designation and management of marine protected areas, edited by J. Davenport, G. Burnell, T. Cross, M. Emmerson, R. McAllen, R. Ramsay, and E. Rogan, Hydrobiologia, 606, 69–79, doi:10.1007/978-1-4020-8808-

7_7, 2008.

425 Gogina, M., Morys, C., Forster, S., Gräwe, U., Friedland, R. and Zettler, M. L.: Towards benthic ecosystem functioning maps: Quantifying bioturbation potential in the German part of the Baltic Sea, Ecol. Indic., 73, 574–588, doi:10.1016/j.ecolind.2016.10.025, 2017.

Grebmeier, J., Bluhm, B., Cooper, L., Denisenko, S., Iken, K., Kedra, M. and Serratos, C.: Time-Series Benthic Community Composition and Biomass and Associated Environmental Characteristics in the Chukchi Sea During 430 the RUSALCA 2004–2012 Program, Oceanography, 28(3), 116–133, doi:10.5670/oceanog.2015.61, 2015.

Gusmao, J. B.: Sediments and Functional Traits : Applying a Functional Trait Approach To Assess Marine Macrobenthic Function, Univeridade Federal do Parana., 2017.

Gutt, J.: On the direct impact of ice on marine benthic communities, a review, Polar Biol., 24(8), 553–564, doi:10.1007/s003000100262, 2001.

435 Hayward, P. J. and Ryland, J. S.: Handbook of the marine fauna of North-West Europe, edited by P. J. Hayward and J. S. Ryland, Oxford University Press, Oxford, New York, Tokyo., 2012.

Hewitt, J. E., Norkko, J., Kauppi, L., Villnäs, A., Norkko, A. and Peters, D. P. C.: Species and functional trait turnover in response to broad-scale change and an invasive species, Ecosphere, 7(3), doi:10.1002/ecs2.1289/supinfo, 2016.

440 Himmelman, J. H. and Dutil, C.: Distribution, population structure and feeding of subtidal seastars in the northern Gulf of St. Lawrence, Mar. Ecol. Prog. Ser., 76, 61–72, doi:10.3354/meps076061, 1991.

Hooper, D. U., Chapin, F. S., Ewel, J. J., Hector, A., Inchausti, P., Lavorel, S., Lawton, J. H., Lodge, D. M., Loreau, M., Naeem, S., Schmid, B., Setälä, H., Symstad, A. J., Vandermeer, J. and Wardle, D. A.: Effects of biodiversity on ecosystem functioning: A consensus of current knowledge, Ecol. Monogr., 75(1), 3–35, 445 doi:10.1890/04-0922, 2005.

Klais, R., Norros, V., Lehtinen, S., Tamminen, T. and Olli, K.: Community assembly and drivers of phytoplankton functional structure, edited by E. Carrington, Funct. Ecol., 31(3), 760–767, doi:10.1111/1365-2435.12784, 2017.

Kleyer, M., Dray, S., Bello, F., Lepš, J., Pakeman, R. J., Strauss, B., Thuiller, W. and Lavorel, S.: Assessing 450 species and community functional responses to environmental gradients: Which multivariate methods?, edited by O. Wildi, J. Veg. Sci., 23(5), 805–821, doi:10.1111/j.1654-1103.2012.01402.x, 2012.

Kokarev, V. N., Vedenin, A. A., Basin, A. B. and Azovsky, A. I.: Taxonomic and functional patterns of macrobenthic communities on a high-Arctic shelf: A case study from the Laptev Sea, J. Sea Res., 129(April), 61–69, doi:10.1016/j.seares.2017.08.011, 2017.

455 Kröncke, I.: Macrobenthos composition, abundance and biomass in the Arctic Ocean along a transect between Svalbard and the Makarov Basin, Polar Biol., 14(8), 519–529, doi:10.1007/BF00238221, 1994.

Lacoste, É., Piot, A., Archambault, P., McKindsey, C. W. and Nozais, C.: Bioturbation activity of three macrofaunal species and the presence of meiofauna affect the abundance and composition of benthic bacterial communities, Mar. Environ. Res., 136, 62–70, doi:10.1016/j.marenvres.2018.02.024, 2018.

460 van der Linden, P., Marchini, A., Dolbeth, M., Patrício, J., Veríssimo, H. and Marques, J. C.: The performance of trait-based indices in an estuarine environment, Ecol. Indic., 61, 378–389, doi:10.1016/j.ecolind.2015.09.039, 2016.

Van Der Linden, P., Borja, A., Rodríquez, J. G., Muxika, I., Galparsoro, I., Patrício, J., Veríssimo, H. and Marques, J. C.: Spatial and temporal response of multiple trait-based indices to natural- and anthropogenic

465   seafloor disturbance (effluents), Ecol. Indic., 69, 617–628, doi:10.1016/j.ecolind.2016.05.020, 2016.

Lucey, N. M., Lombardi, C., DeMarchi, L., Schulze, A., Gambi, M. C. and Calosi, P.: To brood or not to brood: Are marine invertebrates that protect their offspring more resilient to ocean acidification?, Sci. Rep., 5(January), 12009, doi:10.1038/srep12009, 2015.

Marchini, A., Munari, C. and Mistri, M.: Functions and ecological status of eight Italian lagoons examined using

470   biological traits analysis (BTA), Mar. Pollut. Bull., 56(6), 1076–1085, doi:10.1016/j.marpolbul.2008.03.027, 2008.

Mermillod-Blondin, F.: The functional significance of bioturbation and biodeposition on biogeochemical processes at the water–sediment interface in freshwater and marine ecosystems, J. North Am. Benthol. Soc., 30(3), 770–778, doi:10.1899/10-121.1, 2011.

475   Micheli, F. and Halpern, B. S.: Low functional redundancy in coastal marine assemblages, Ecol. Lett., 8(4), 391–400, doi:10.1111/j.1461-0248.2005.00731.x, 2005.

Norkko, A., Villnäs, A., Norkko, J., Valanko, S. and Pilditch, C.: Size matters: implications of the loss of large individuals for ecosystem function., Sci. Rep., 3, 2646, doi:10.1038/srep02646, 2013.

Oug, E., Fleddum, A., Rygg, B. and Olsgard, F.: Biological traits analyses in the study of pollution gradients and

480   ecological functioning of marine soft bottom species assemblages in a fjord ecosystem, J. Exp. Mar. Bio. Ecol., 432, 94–105 [online] Available from: http://apps.webofknowledge.com/full_record.do?product=CCC&search_mode=GeneralSearch&qid=7&SID=W 29TQeqCuCsBtQBk9hp&page=1&doc=9 (Accessed 28 April 2016), 2012.

Pearson, T. H.: Functional group ecology in soft-sediment marine benthos: the role of bioturbation, Oceanogr.

485   Mar. Biol. Annu. Rev., 39(1), 78–94, 2001.

Piepenburg, D.: Arctic Brittle Stars (Echinodermata: Ophiuroidea), Oceanogr. Mar. Biol. Annu. Rev., 38, 189–256, 2000.

Piló, D., Ben-Hamadou, R., Pereira, F., Carriço, A., Pereira, P., Corzo, A., Gaspar, M. B. and Carvalho, S.: How functional traits of estuarine macrobenthic assemblages respond to metal contamination?, Ecol. Indic., 71, 645–

490   659, doi:10.1016/j.ecolind.2016.07.019, 2016.

Queirós, A. M., Birchenough, S. N. R., Bremner, J., Godbold, J. A., Parker, R. E., Romero-Ramirez, A., Reiss, H., Solan, M., Somerfield, P. J., Van Colen, C., Van Hoey, G. and Widdicombe, S.: A bioturbation classification of European marine infaunal invertebrates., Ecol. Evol., 3(11), 3958–85, doi:10.1002/ece3.769, 2013.

Renaud, P., Tessmann, M., Evenset, A. and Christensen, G.: Benthic food-web structure of an Arctic fjord

495   (Kongsfjorden, Svalbard), Mar. Biol. Res., 7(1), 13–26, doi:10.1080/17451001003671597, 2011.

Rosenberg, R.: Benthic marine fauna structured by hydrodynamic procesess and food availability, Neth. J. Sea. Res., 34(4), 303–317, 1995.

Schleuter, D., Daufresne, M., Massol, F. and Argillier, C.: User's guide to functional diversity indices., Ecol. Monogr., 80(3), 448–469, doi:10.1890/08-2225.1, 2010.

500   Solan, M., Aspden, R. J. and Paterson, D. M.: Marine biodiversity & ecosystem functioning, First edit., edited by M. Solan, R. J. Aspden, and D. M. Paterson, Oxford University Press, Oxford., 2012.

Spitz, J., Ridoux, V. and Brind'Amour, A.: Let's go beyond taxonomy in diet description: Testing a trait-based approach to prey-predator relationships, J. Anim. Ecol., 83(5), 1137–1148, doi:10.1111/1365-2656.12218, 2014.

Törnroos, A. and Bonsdorff, E.: Developing the multitrait concept for functional diversity: Lessons from a

505   system rich in functions but poor in species, Ecol. Appl., 22(8), 2221–2236, doi:10.1890/11-2042.1, 2012.

Tyler, E. H. M., Somerfield, P. J., Berghe, E. Vanden, Bremner, J., Jackson, E., Langmead, O., Palomares, M. L. D. and Webb, T. J.: Extensive gaps and biases in our knowledge of a well-known fauna: Implications for integrating biological traits into macroecology, Glob. Ecol. Biogeogr., 21(9), 922–934, doi:10.1111/j.1466-8238.2011.00726.x, 2012.

510   Wassmann, P., Duarte, C. M., Agustí, S. and Sejr, M. K.: Footprints of climate change in the Arctic marine ecosystem, Glob. Chang. Biol., 17(2), 1235–1249, doi:10.1111/j.1365-2486.2010.02311.x, 2011.

Weigel, B., Blenckner, T. and Bonsdorff, E.: Maintained functional diversity in benthic communities in spite of diverging functional identities, Oikos, 125(10), 1421–1433, doi:10.1111/oik.02894, 2016.

Weslawski, J., Wlodarska-Kowalczuk, M. and Legezynska, J.: Occurrence of soft bottom macrofauna along the

515   depth gradient in High Arctic, 79 N, Pol. Polar Res, 24(1), 73–88 [online] Available from: http://polish.polar.pan.pl/ppr24/ppr24-073.pdf, 2003.

Wieczorek, J., Bloom, D., Guralnick, R., Blum, S., Döring, M., Giovanni, R., Robertson, T. and Vieglais, D.: Darwin core: An evolving community-developed biodiversity data standard, PLoS One, 7(1), doi:10.1371/journal.pone.0029715, 2012.

520   Wiedmann, M., Aschan, M., Certain, G., Dolgov, A., Greenacre, M., Johannesen, E., Planque, B. and Primicerio, R.: Functional diversity of the Barents Sea fish community, Mar. Ecol. Prog. Ser., 495, 205–218, doi:10.3354/meps10558, 2014.

Web links / Trait databases

Encyclopedia of Life: http://www.eol.org/, last access: 29 June 2018.

525   Froese, R. and D. Pauly. Editors. 2018. FishBase. World Wide Web electronic publication. www.fishbase.org, version (02/2018).

Functional trait compilation for marine phytoplankton: https://www.riinaklais.com/phytotraits, last access: 28 January 2019.

GBIF.org (2018), *GBIF Home Page*: https://www.gbif.org/, last access: 29 June 2018.

530   Marine Macrofauna Genus Trait Handbook: http://www.genustraithandbook.org.uk/, last access: 27 June 2018.

MarLIN, 2006. *BIOTIC - Biological Traits Information Catalogue.* Marine Life Information Network. Plymouth: Marine Biological Association of the United Kingdom: www.marlin.ac.uk/biotic/, last access: 18 April 2018.

Marine Species Traits editorial board (2018). Marine Species Traits: http://www.marinespecies.org/traits/, last

535   access: 27 June 2018.

Neogene Marine Biota of Tropical America (NMiTA): http://eusmilia.geology.uiowa.edu/, last access: 29 June 2018.

Ocean Biogeographic Information System (OBIS): http://www.iobis.org/, last access: 27 June 2018.

Sea Life Base: http://www.sealifebase.org/, last access: 29 June 2018.

540   The Polychaetes scratchpad: http://polychaetes.lifewatchgreece.eu/, last access: 27 June 2018.

WoRMS Editorial Board (2017). World Register of Marine Species: http:// www.marinespecies.org/, last access: 27 June 2018. doi:10.14284/170.