

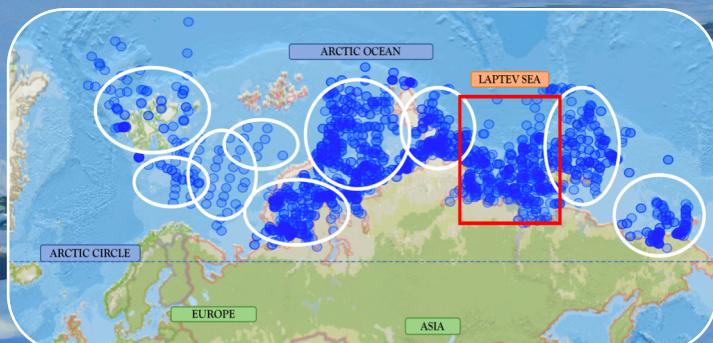
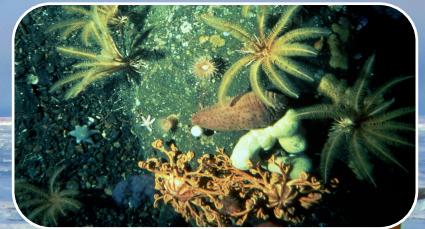
Spatial Variation in Distributional Patterns of Arctic Benthic Biodiversity



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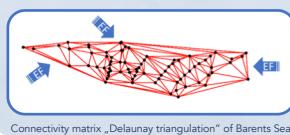
Background

Marine benthic organisms living in shelf seas of the Siberian Arctic are impacted severely by the pronounced consequences of climate change. Polar ecosystems are used to stable conditions and even small changes could influence benthic communities, food webs, and ecosystem functions. Regional patterns in community composition result from the interaction between marine organisms and their local environment. In this study, these relationships are investigated in order to understand how climate change affect Arctic biodiversity in particular.



Approach

- Modelling the regional distribution and composition of Arctic biota
- Identification of spatial scales and influencing environmental factors (EF)
- Multivariate modelling techniques, e.g. Moran's eigenvector mapping (MEM)



Connectivity matrix „Delaunay triangulation“ of Barents Sea

Limitations

- “Zero-Problem”: Siberian Arctic biota display low occurrences
- Data transformations solved nothing
- Result: insignificant data outputs

Broad Scale environ. data	variables	order	R2	R2Cum	AdjR2Cum	F	p-value
bat	5	0.14131599	0.141316	0.137516	38.19342	0.001	
sea_ice_mean	1	0.02774553	0.1690615	0.1656754	7.5121	0.005	
sea_ice_max	4	0.03014955	0.1992112	0.1884863	8.43358	0.001	
sea_ice_sd	2	0.04130085	0.240512	0.2268889	12.12679	0.0001	
sea_ice_min	3	0.0286491	0.2691611	0.2527008	8.70246	0.002	
bat_squared	6	0.0420276	0.2831839	0.2637228	4.323327	0.0073	

Example: significant data output of environmental factors

Data

Biotic

Panarctic Biota “Panabio” Databank

- 1800 – 2014
- 10322 samples
- 49 regions
- 2746 species



- 1992 – 2014
- 228 samples
- 10 regions
- 412 species

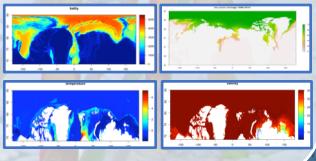
Example species:
Stegophura nodosa;
Ampelisca macrocephala;
Acanthostephia malmgreni;
Cylchna alba



Abiotic

Public Online Databases

- Bathymetry: IBCAO
- Sea ice cover: NSIDC
- Temperature: NOAA WOA 2018
- Salinity: NOAA WOA 2018



Alternative

MaxEnt - Maximum Entropy distribution modelling

- Modelling species distribution from presence-only records
- Contrasting species presences against background locations with unknown p/a → absences allow the prediction of presence probability
- Input: presence-only; environmental factors; landscape; ecological traits
- Output: Relative Occurrence Rate; biodiversity hotspots and coldspots; trait relevance important for species composition



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