

A unified typology for small European rivers

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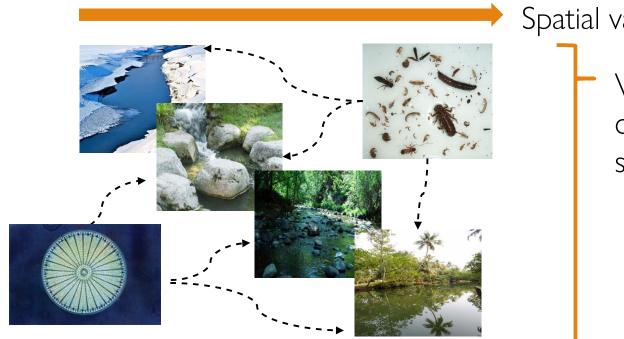


Adverse effects of Chemicals on Biodiversity

- In 2018, 84.1 Million tons of environmentally hazardous chemicals were produced in the EU (EUROSTAT, 2020)
- Their release into aquatic ecosystems can lead to adverse effects on biodiversity (Floehr et al. 2015; Schäfer et al. 2016; Vethaak et al. 2005)
- Ecological Risk Assessment (ERA) tries to prevent or limit such effects
- In Europe, ERA practices follow a one-size-fits-all approach regulatory thresholds do not differ between recipient systems But should they?

Spatial variation in sensitivity

- How does chemical sensitivity vary across European streams?
- And how should this be considered in ERA?



Spatial variation

Variation in chemical sensitivity?

Get real!

To investigate these questions we started the GetReal project

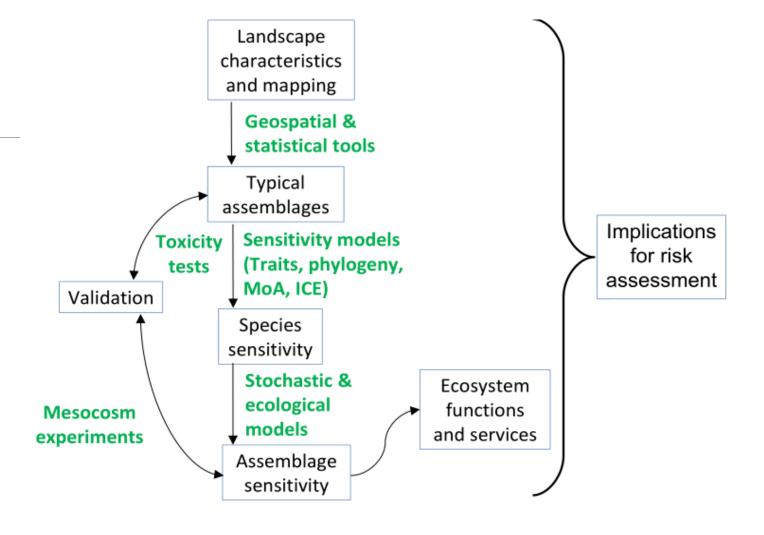


Figure 1: Research Program of GetReal

GetReal is a collaborative effort of: Ralf Schäfer, Martin Entling, Paul van den Brink, Lorraine Maltby, Frederik De Laender, Sebastian Scheu, Javier Jarillo, Sanne van den Berg, Tomás Duque and Jonathan Jupke



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In this presentation we will focus on the first step - a stream typology

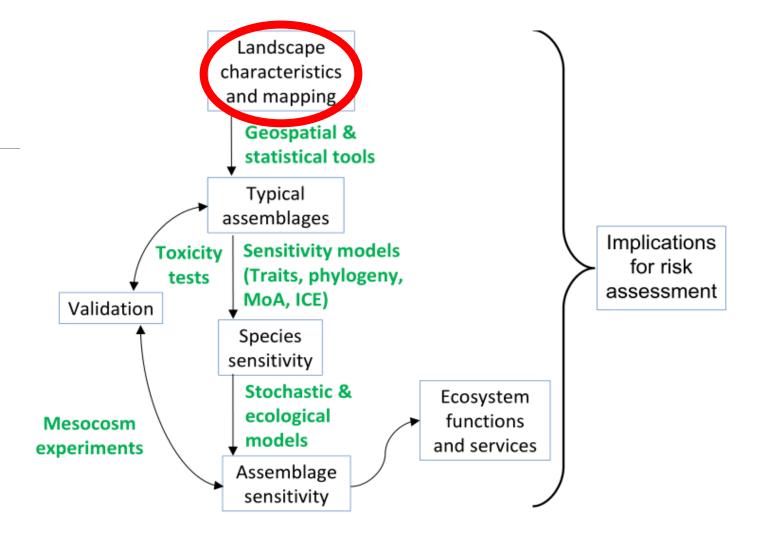


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Rationale

- For this project, we selected eight countries, which represent macroecological gradients like climate and geology
- •We further focused on small rivers which host a proportionally higher share of biodiversity and are at higher risk from chemical inputs (Lorenz et al. 2017; Link et al. 2017)
- A pan-European typology will be available later this year

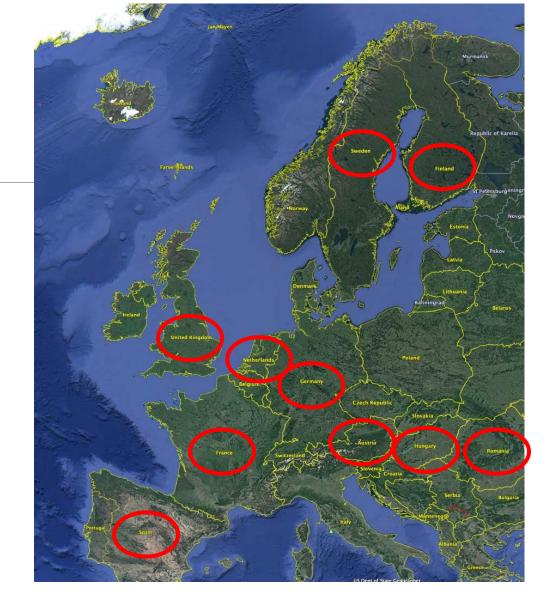


Figure 2: Map of selected Countries



Catchment characteristics and modelling

- The typology is based on CCM2 (Vogt et al. 2007)
- CCM2 is a Pan-European stream-network with associated catchments and environmental variables

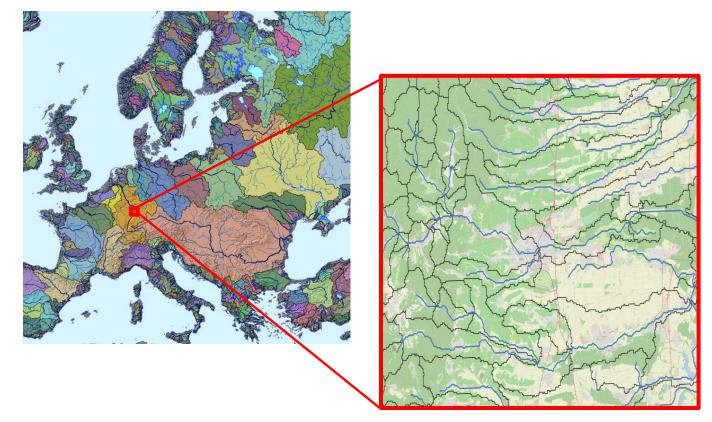


Figure 3: Extend of CCM2 with enlarged region showing streams (blue) and catchments (black)



Overview of variables

Abiotic Descriptor	Spatial Unit	Source
Catchment area	Catchments	CCM2.1 [7]
Upstream catchment area	Catchments	CCM2.1 [7]
Size class	River Segment	Derived from discharge
Sinuosity	River Segment	CCM2.1 [7]
Distance to Source	River Segments	CCM2.1 [7]
Elevation	Catchment	CCM2.1 [7]
Elevation heterogeneity	Catchment	CCM2.1 [7]
Maximal Elevation	Catchment	CCM2.1 [7]
Air temperature	Catchment	Worldclim [8]
Temperature range	Catchment	Worldclim [8]
Upstream catchment area	Catchment	CCM2.1 [7]
Share of Ice-covered Area	Catchment	Vegetation at last glacial maximum[9]
Share of steppe	Catchment	Vegetation at last glacial maximum [9]
Share of forest	Catchment	Vegetation at last glacial maximum [9]

Abiotic Descriptor	Spatial Unit	Source
Share of boreal vegetation	Catchment	Vegetation at last glacial maximum [9]
Precipitation	Catchment	CHELSA [10]
Precipitation Seasonality	Catchment	CHELSA [10]
Share of Calcareous/Silicate/Sediment	Catchment	IHME 1500 v1.1 [11]
Mean Discharge	River Segment	CHELSA [10]; GRDC [12]; NRFA [13]
Coefficient of Variation Discharge	River Segment	CHELSA [10]; GRDC [12]; NRFA [13]
Skewness discharge	River Segment	CHELSA [10]; GRDC [12]; NRFA [13]
Kurtosis discharge	River Segment	CHELSA [10]; GRDC [12]; NRFA [13]
the autoregressive lag-one correlation coefficient	River Segment	CHELSA [10]; GRDC [12]; NRFA [13]
Amplitude of seasonal signal	River Segment	CHELSA [10]; GRDC [12]; NRFA [13]
Phase of seasonal signal	River Segment	CHELSA [10]; GRDC [12]; NRFA [13]

Discharge Timeseries

- We modeled daily discharge for each stream segment from 01.01.2000 until 31.12.2013
- Following Irving et al. 2018, we used upstream precipitation as predictor variable
- Gauging data from Global Runoff data center and the National River Flow archive were used
- Resulting discharge time series were summarized as: mean, coefficient of variation, skewness, kurtosis, autoregressive lag-one correlation coefficient, amplitude, and phase of the seasonal signal (Archfield *et al.*, 2014)
- All stream segments with mean discharge > 10 m³/s were categorized as big and omitted from the subsequent clustering



Weighting Variables

- We use Generalized Dissimilarity Models (GDM) to weigh the variables prior to clustering (Ferrier *et al.*, 2007)
- GDMs are GLMs in which a distance matrix between sites is modeled through the differences in I-spline smooth functions of the environmental variables. The splines are scaled by their explanatory power in the model. The smooth functions from the GDM can be applied to a data set with the same variables but additional sites
- Using a large database of aquatic invertebrate occurrences compiled for the second part of GetReal, we fit a GDM and transformed the variables for all stream segments

Clustering

- We dropped variables until no correlation was >= 0.7
- We computed tree-based dissimilarity (d4) between all stream segments (Buttrey & Whitaker, 2015) and clustered the resulting distance-matrix with CLARA (Kaufmann & Rousseeuw, 1986)
- •CLARA is a non-hierarchical clustering approach, hence the number of clusters must be supplied by the researcher. We computed all clusterings with group sizes from 2 to 30 and selected the minimum group size that was within 5% silhouette with of the group with the absolute minimum silhouette width.
- For the unweighted variables this resulted in 15 stream types and for the weighted variables in 11.

Stream Typologies

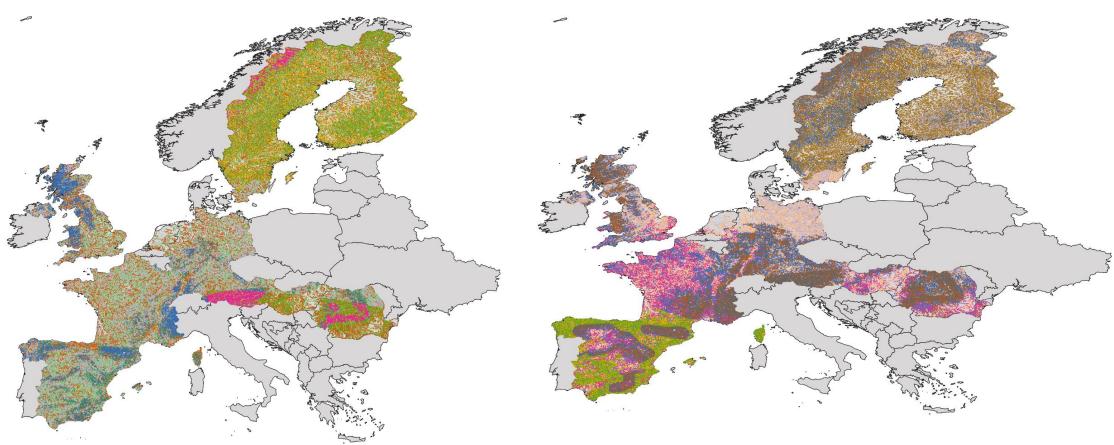
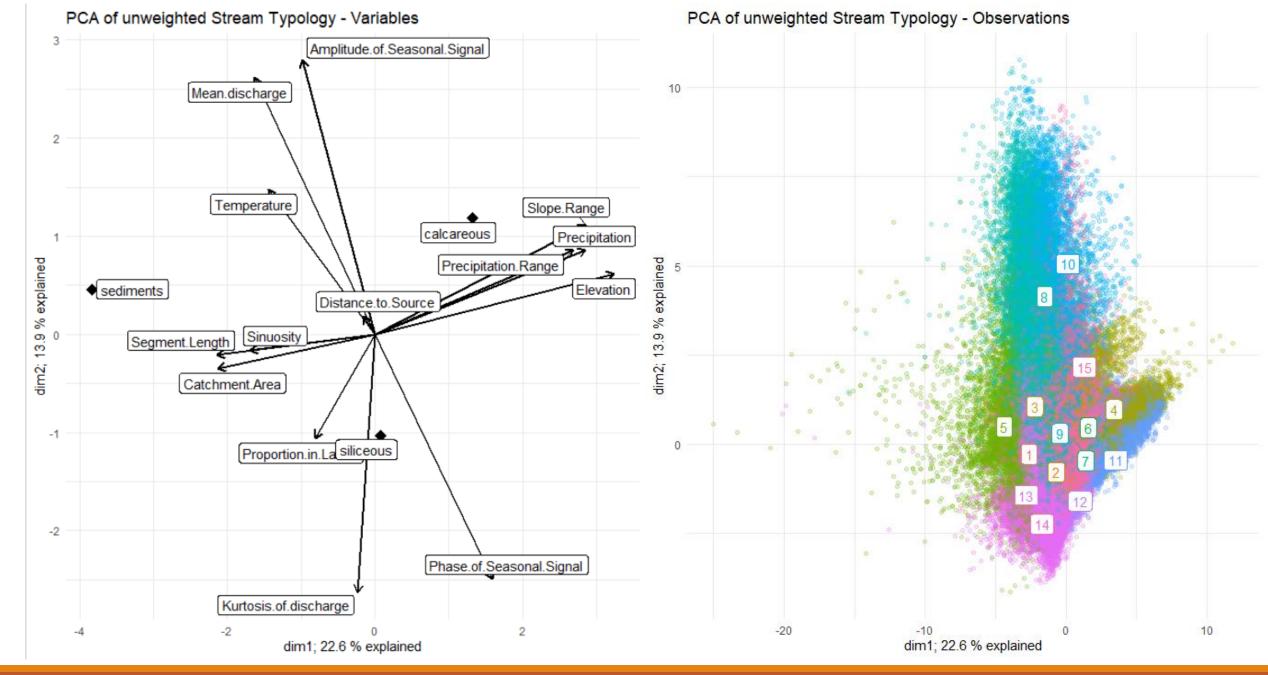


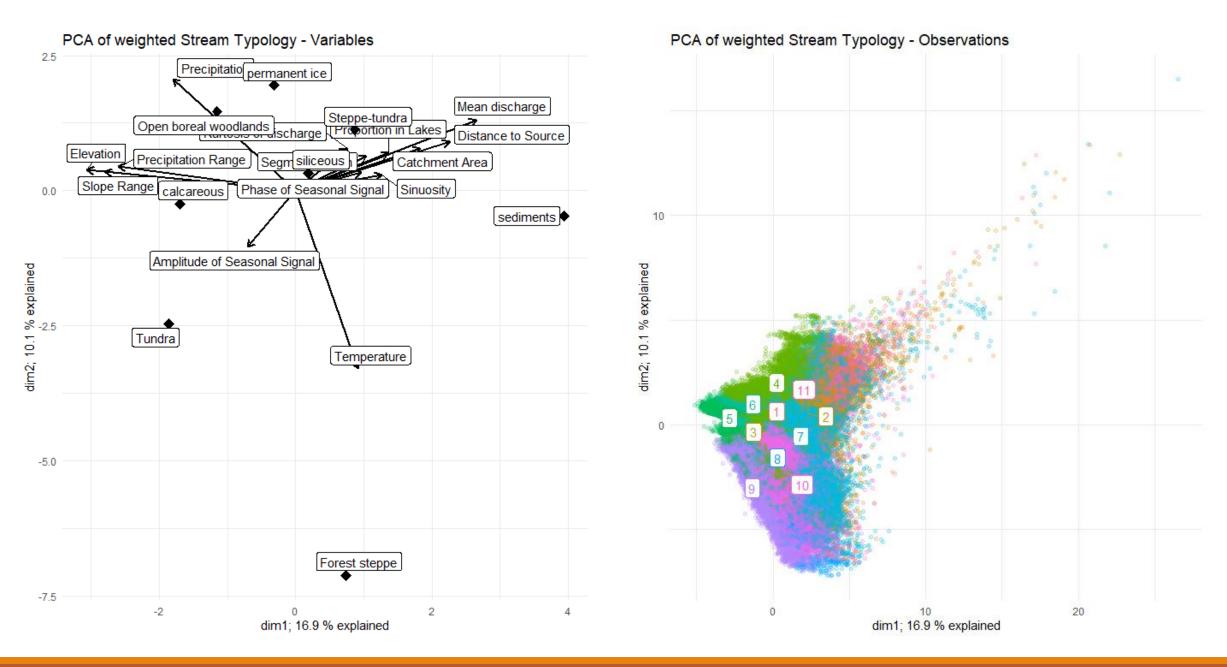
Figure 4: Unweighted stream typology

Figure 5: Weighted stream typology

Ordination of input variables

- On the next two slides you will see biplots of first the unweighted and then the weighted typology
- To declutter these images we separated the variables (left) from the observations (right)
- Please note, that axis length differs between the variable and the observation plot and that the variable coordinates have been stretched by a factor of four to improve legibility
- The color of observations together with a label in the same color indicates stream type
- •The ordination is a mixed ordination employing Principal Component Analysis for continuous variables and Multiple Correspondence Analysis for categorical variables. See PCAmix in Chavent *et al.* (2017) for details





Potential use cases

- Besides our aim to predict typical assemblages our typology can be applied to:
 - develop stratification schemes for monitoring programs
 - analyze gaps in existing or planned conservation endeavors
 - Species Distribution Models as a predictor variable (McManamay et al. 2018)
 - guide restoration efforts by identifying reference conditions within each type

Thanks to





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