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# Exploring patterns of diatom assemblages from circumpolar Arctic lakes & streams

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# Background

The assessment of spatial and temporal trends in Arctic freshwater biota and their physical and biogeochemical habitat in response to a warming climate requires a large-scale harmonized monitoring program. The presented work, performed by the freshwater group of the Circumpolar Biodiversity Monitoring Plan (Arctic Council: Conservation of Arctic Flora and Fauna), is one of the first steps towards such a harmonized monitoring program. Using circum-Arctic diatom assemblage data, specific goals of this study include: 1) establishing current and pre-industrial environmental conditions as reference points to guide future environmental monitoring programs, and 2) understanding the historical context of diatom distributions. Large-scale assessments of diatom distributions in Arctic regions are currently scarce, and detailed taxonomic studies tracking species compositional changes in both lakes and streams have yet to be undertaken. We used contemporary stream and lake data from ongoing monitoring programs, spot data from research projects, as well as surface sediment (modern) diatoms from paleolimnological studies to provide a spatial assessment of species distributions of the circum-Arctic region. Additionally, we analysed a large set of dated and “top-bottom” paleolimnological records to better describe historical background conditions on a circum-Arctic scale.

More information: [caff.is/freshwater](http://caff.is/freshwater)

# Background

The presented work, performed by the **freshwater** group of the Circumpolar Biodiversity Monitoring Plan (CBMP) (Arctic Council: Conservation of Arctic Flora and Fauna, **CAFF**), is one of the first steps towards a harmonized circumpolar monitoring program.

Goals of this study, aimed at guiding the monitoring plan development:

- 1) evaluating existing data and identifying gaps in monitoring efforts and scientific knowledge of Arctic freshwaters
- 2) establishing current and pre-industrial environmental conditions as reference points
- 3) understanding the historical context of diatom distributions



## Circumpolar diatom database

**5 different types of diatom data** included in our analyses, from monitoring programs & academic research studies:

427 Stream samples (Canada, Finland, Norway, Sweden, USA)

727 Lake surface samples (Canada, Finland, Iceland, Sweden, Russia, USA)

118 Lake shoreline samples (Canada, Sweden) *Modern samples*

116 Top/bottom of sediment cores (Canada, Finland, Russia, USA)

52 Full cores (Canada, Finland, Russia) *Pre-industrial era samples*

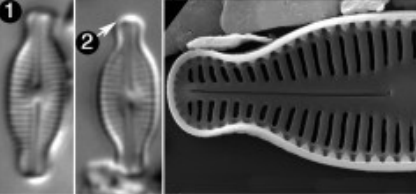
- The Circumpolar diatom database is managed centrally by CAFF, and is depending on the CAFF member countries' input of data
- Our study focuses on
  - diatom data expressed as relative abundances
  - dated paleo cores and the last ~ 200 year interval

# Circumpolar diatom database

Final taxa group: *Achnanthes acares/ricula/carissima*

NOMENCLATURE\_DIATOMS


Class	TaxonName	TaxonNameFin	GenusFinal	Species
Bacillariophyce	Achnanthes cf. carissima	Achnanthes cf. carissima	Achnanthes	cf. cari
Bacillariophyce	Achnanthes cf. rricula	Achnanthes cf. rricula	Achnanthes	cf. rricu
Bacillariophyce	Achnanthes rricula	Achnanthes rricula	Achnanthes	rricula
Bacillariophyce	Diadensis sp. 1 (Achnanthes cariss)	Achnanthes cf. carissima	Achnanthes	cf. cari
Bacillariophyce	Diadensis sp. aff. contenta (Navicu	Navicula cf. schmassmannii		cf. schi
Bacillariophyce	Karayevia carissima	Achnanthes carissima		carissii
Bacillariophyce	Navicula cf. schmassmannii	Navicula cf. schmassmannii		schma
Bacillariophyce	Navicula schmassmannii	Navicula schmassmannii		schma
Bacillariophyce	Navicula schmassmannii	Navicula schmassmannii		schma
Bacillariophyce	Navicula schmassmannii	Navicula schmassmannii		schma
Bacillariophyce	Navicula schmassmannii	Navicula schmassmannii		schma
Bacillariophyce	Navicula schmassmannii	Navicula schmassmannii		schma
Bacillariophyce	Navicula schmassmannii	Navicula schmassmannii		schma
Bacillariophyce	Navicula schmassmannii	Navicula schmassmannii		schma
Bacillariophyce	Navicula schmassmannii Hustedt	Navicula schmassmannii		schma
Bacillariophyce	Navicula schmassmannii Hustedt 1	Navicula schmassmann		schma
Bacillariophyce	Navicula schmassmanni	Navicula schmassmann		schma
Bacillariophyce	Navicula schumassmannii	Navicula schmassmann		schma
Bacillariophyce	Naviculadicta schmassmannii	Navicula schmassmann		schma
Bacillariophyce	Achnanthes brevipes	Achnanthes brevipes		brevip
Bacillariophyce	Achnanthes brevipes v. intermedi	Achnanthes brevipes va		brevip
Bacillariophyce	Achnanthes cf. coarctata	Achnanthes coarctata		coarcta



*Humidophila schmassmannii*

Length: < 9 µm

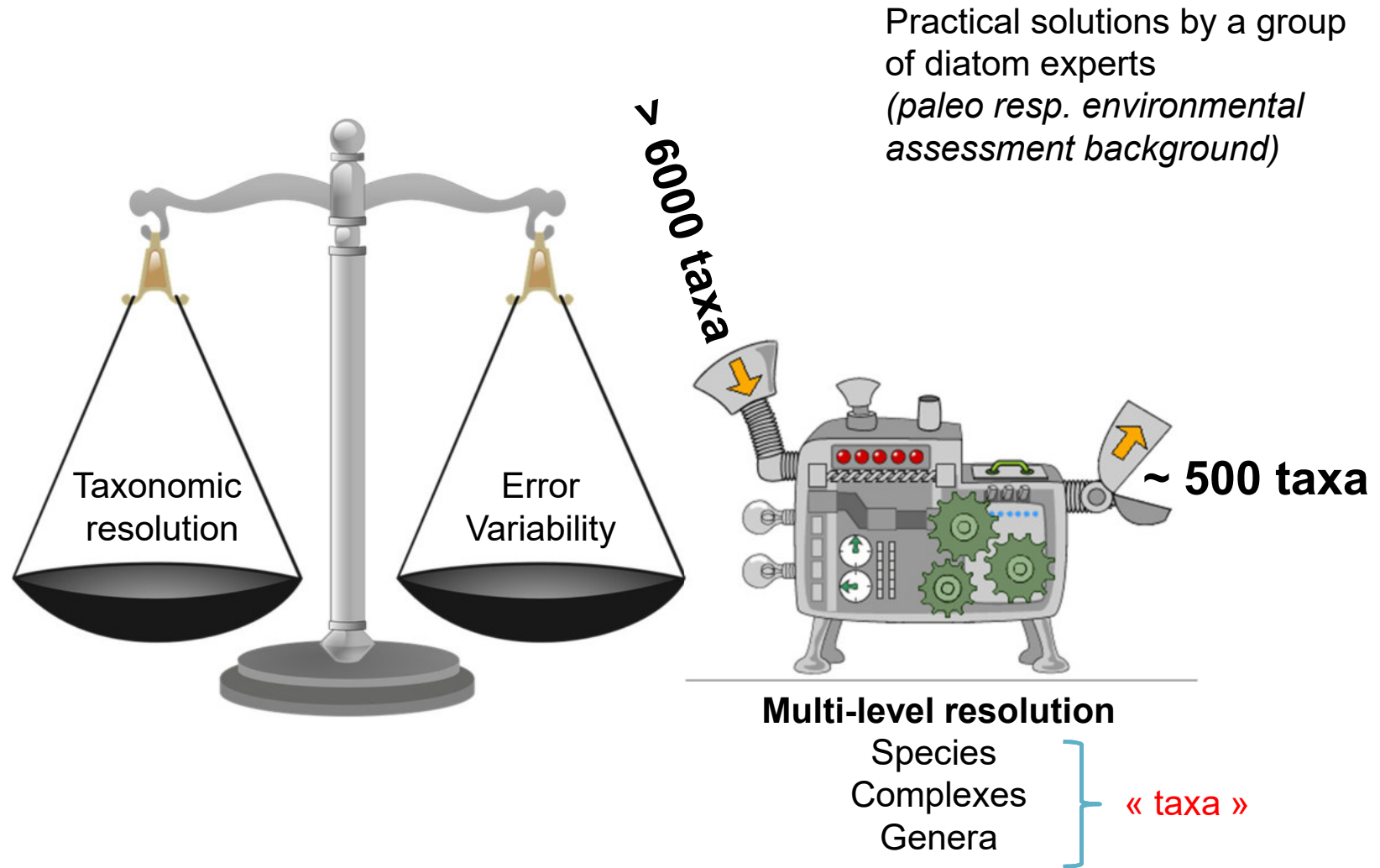
<https://diatoms.org/>



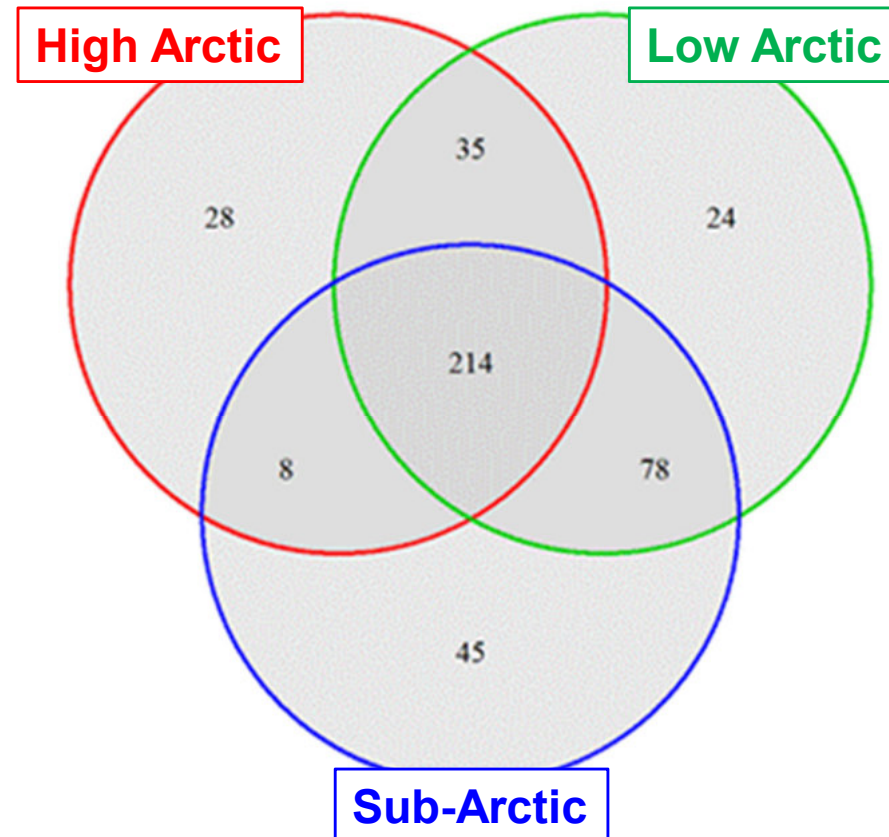
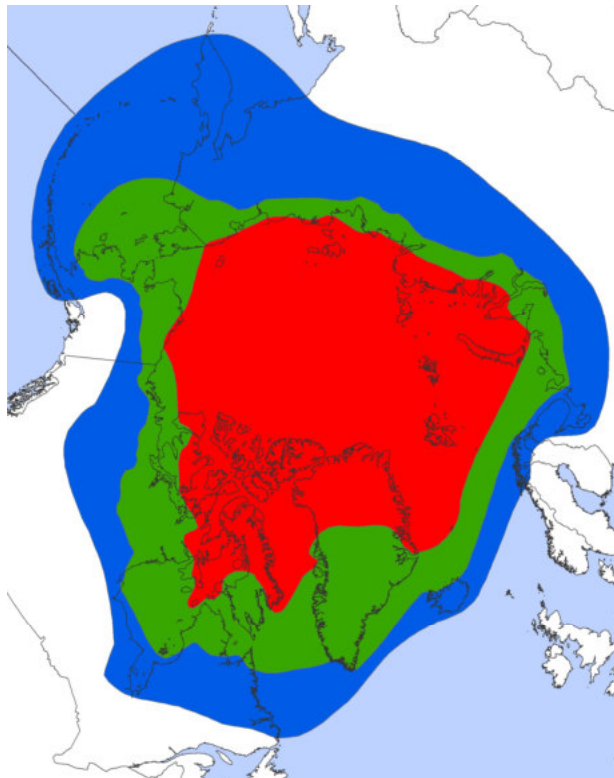
Record: 1 of 6148



## Taxonomic harmonization

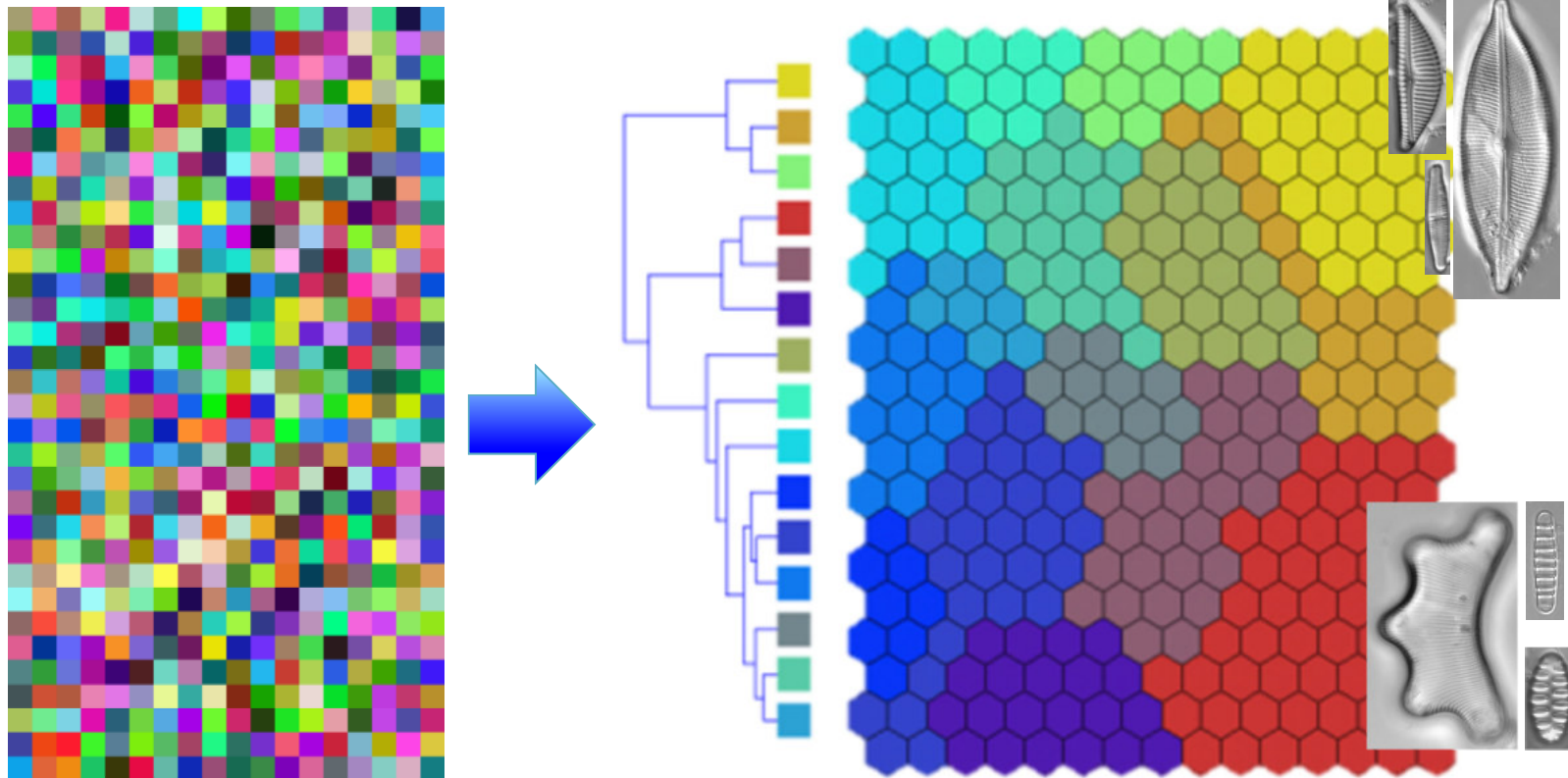


Most taxa are common in all Arctic regions (latitude and longitude)

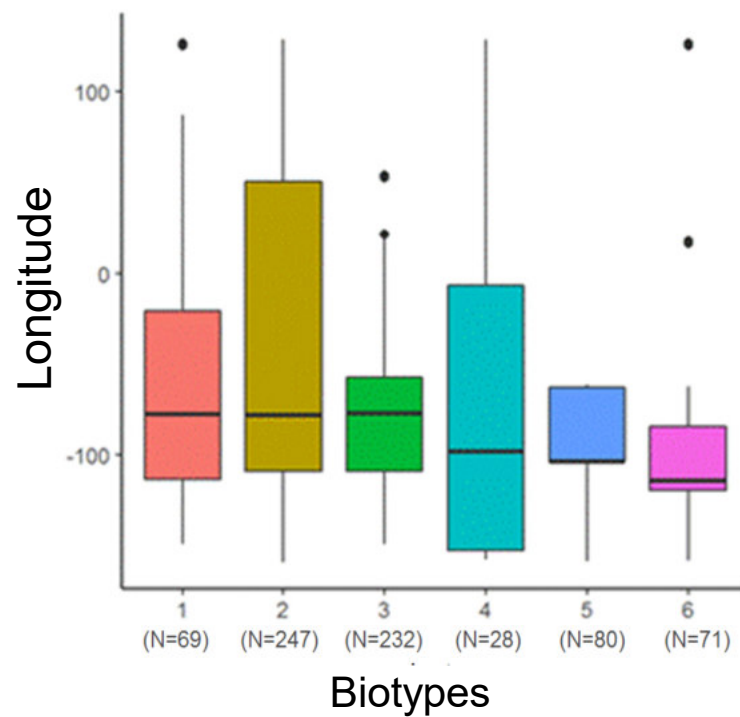


Diatom taxa among Arctic regions  
for lake surface sediments

**“Biotypes”** created using self-organizing maps (SOM) ... clusters in short!





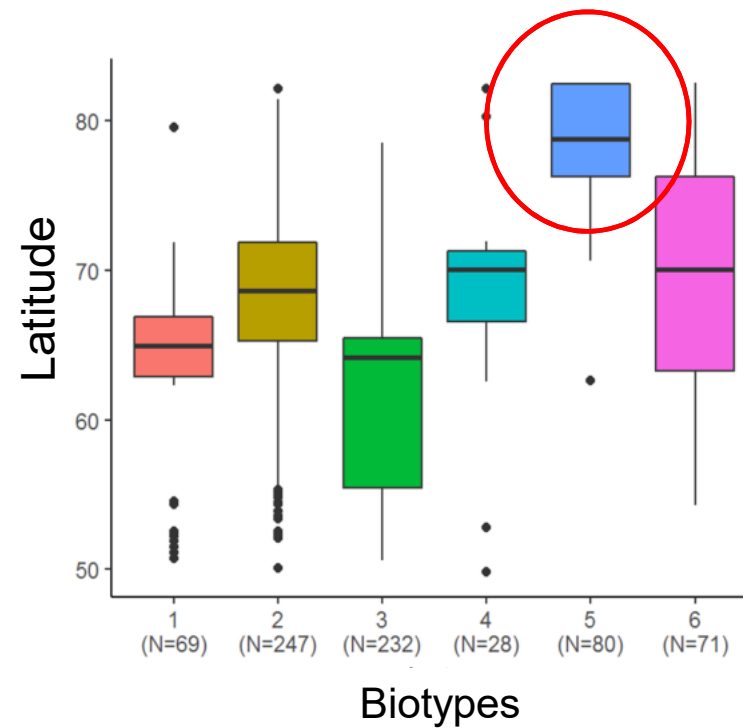
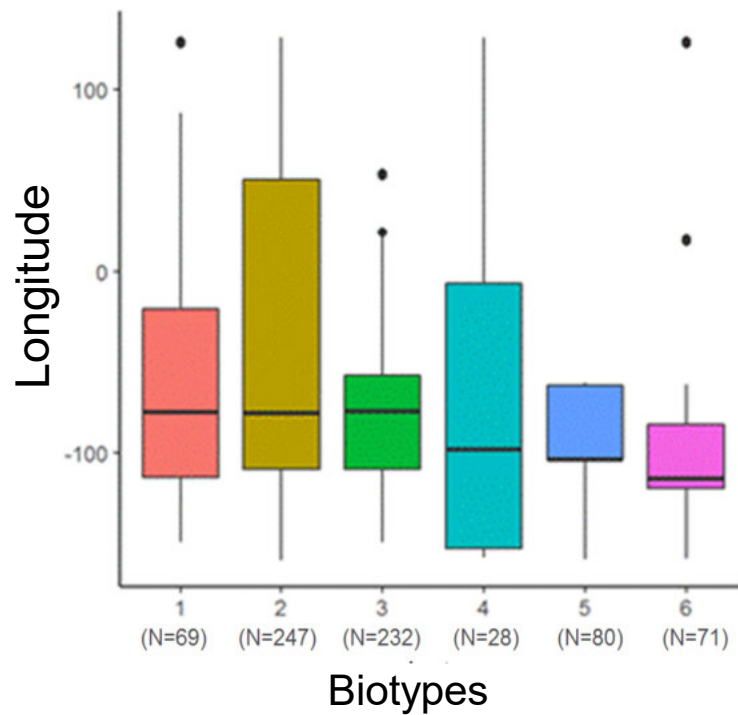
Lake surface sediments, n = 727**6 “Biotypes”**

# Lake surface sediments



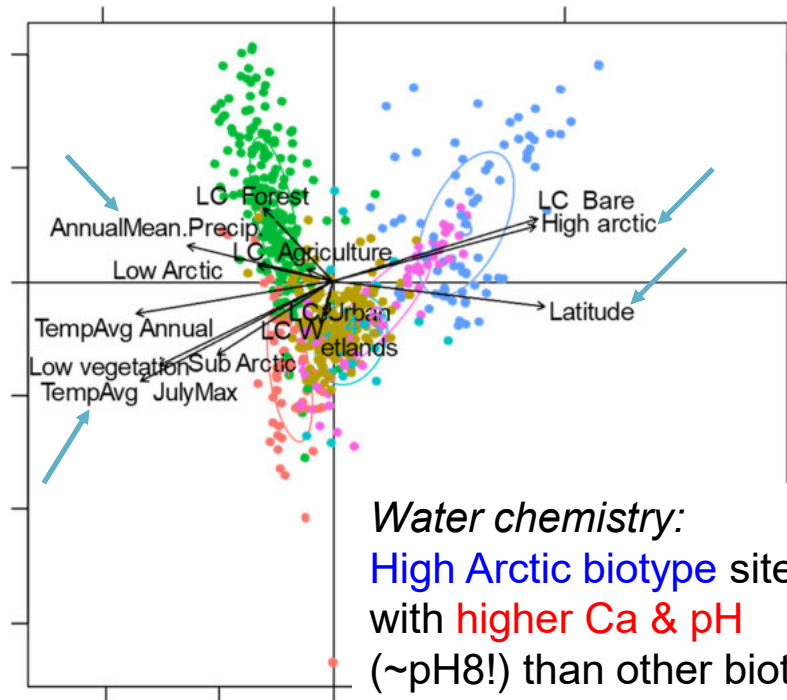
## Lake surface sediments, n = 727

### 6 “Biotypes”

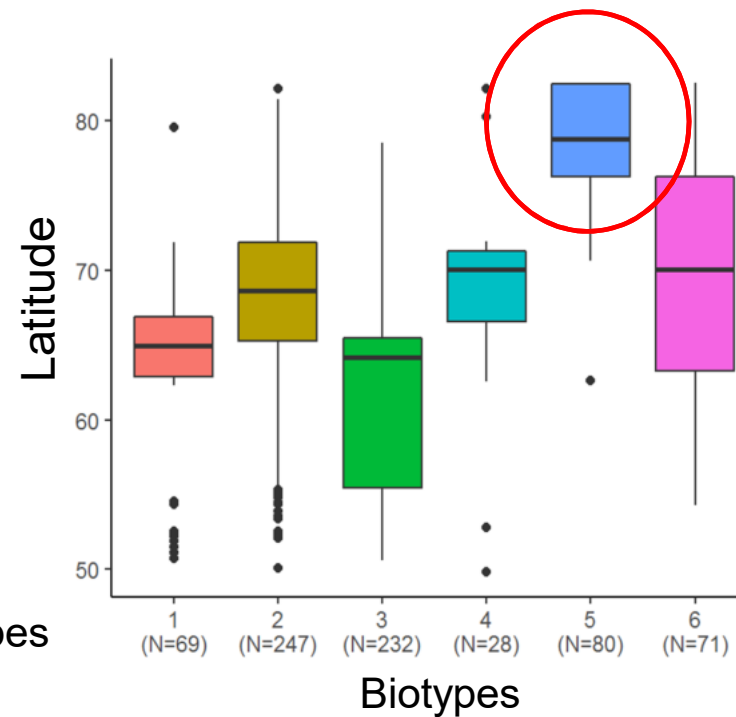


## Lake surface sediments, n = 727

Canonical correspondence analysis  
Samples color-coded according to biotypes

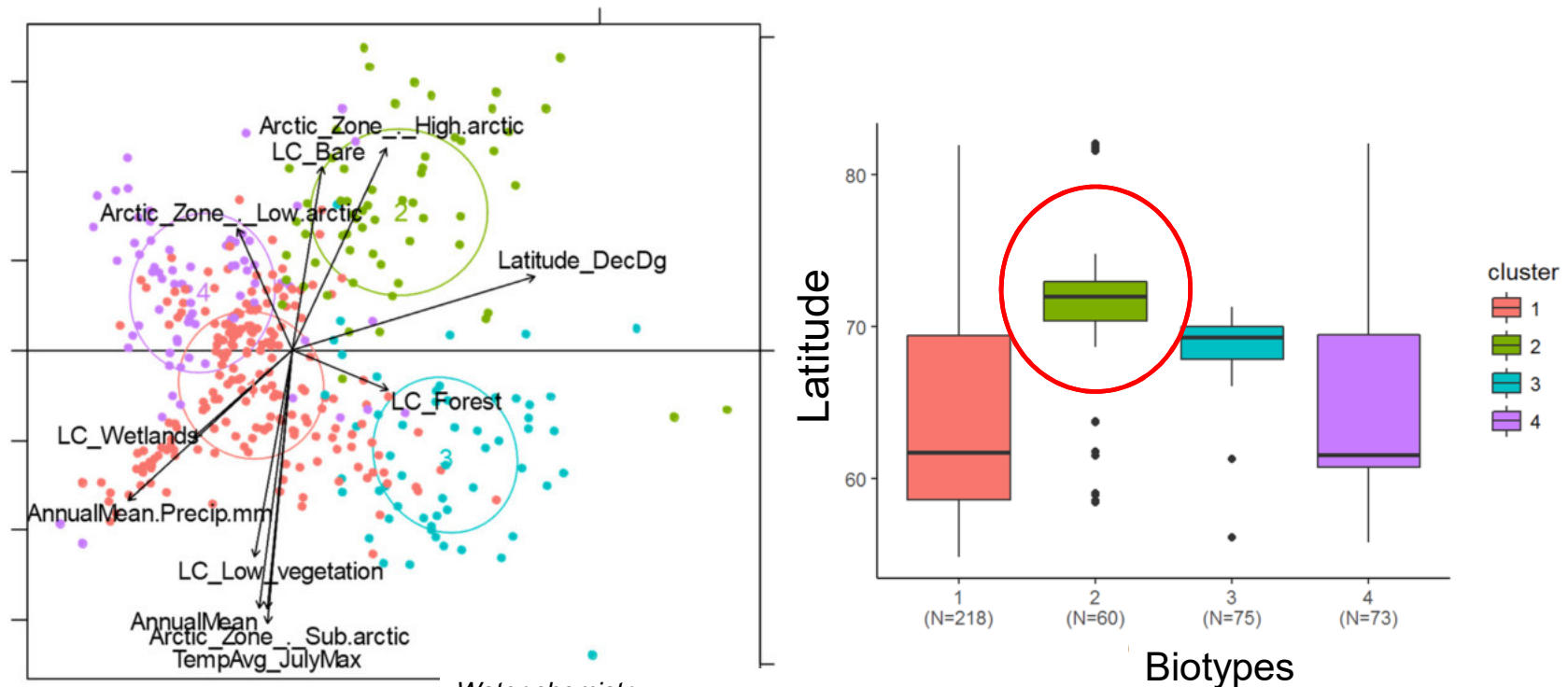


*Water chemistry:*  
High Arctic biotype sites  
with higher Ca & pH  
(~pH8!) than other biotypes  
[not available for all sites,  
n=253]



## Streams, n = 427

### 4 “Biotypes”



Water chemistry:  
 High Arctic biotype sites with higher Ca & pH (~pH8!) than other biotypes  
 [not available for all sites, n=174]



Streams, n=427

*Modern samples*

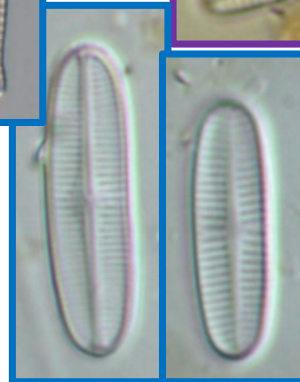
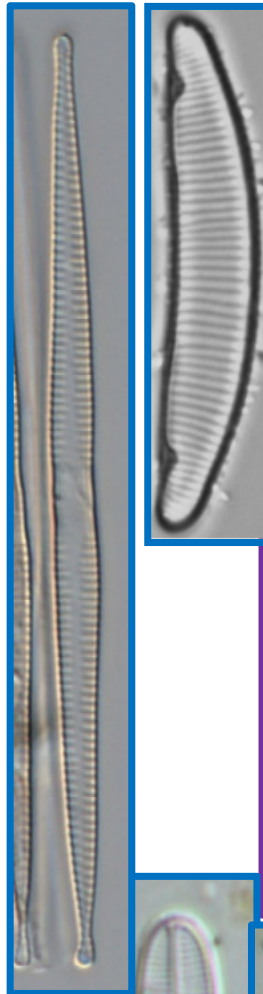
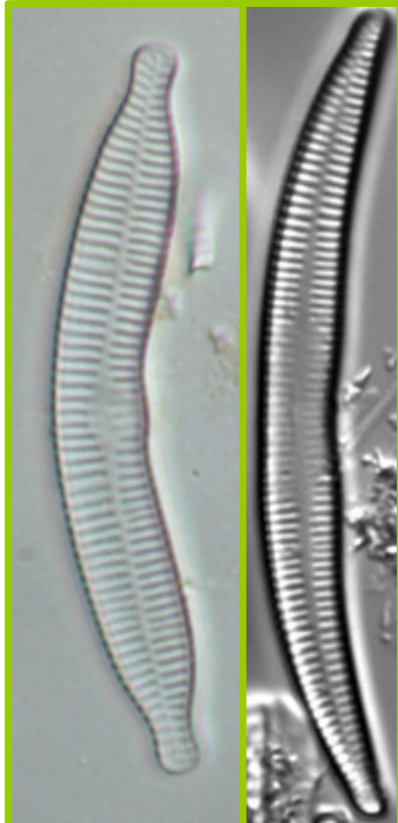


rel. abu. [%]	1	2	3	4	sum abund
<b>Achnanthidium minutissimum</b>	0,45	0,19	0,12	0,1	<b>0,86</b>
Fragilaria capucina complex	0,09	0,1	0,05	0,18	<b>0,42</b>
<b>Tabellaria flocculosa Complex</b>	0,07	0,01	0,03	0,21	<b>0,32</b>
<b>Syndedra cyclopum/Hannaea arcus</b>	0,02	0,19	0,05	0,01	<b>0,27</b>
<b>Diatoma moniliformis/tenuis</b>	0,02	0,19	0,01	0	<b>0,22</b>
<b>Achnanthidium kriegeri</b>	0,03	0	0,01	0,13	<b>0,17</b>
<b>Rossithidium pusillum/anastasiae</b>	0,02	0	0,1	0	<b>0,12</b>
Nitzschia perminuta Complex	0,02	0,03	0,02	0,04	<b>0,11</b>
<b>Fragilaria tenera complex</b>	0,02	0,01	0,08	0	<b>0,11</b>
Eunotia.incisa.Complex	0	0	0,02	0	<b>0,02</b>
<b>Psammothidium.marginulatum.scoticum.la</b>	0	0	0	0,03	<b>0,03</b>

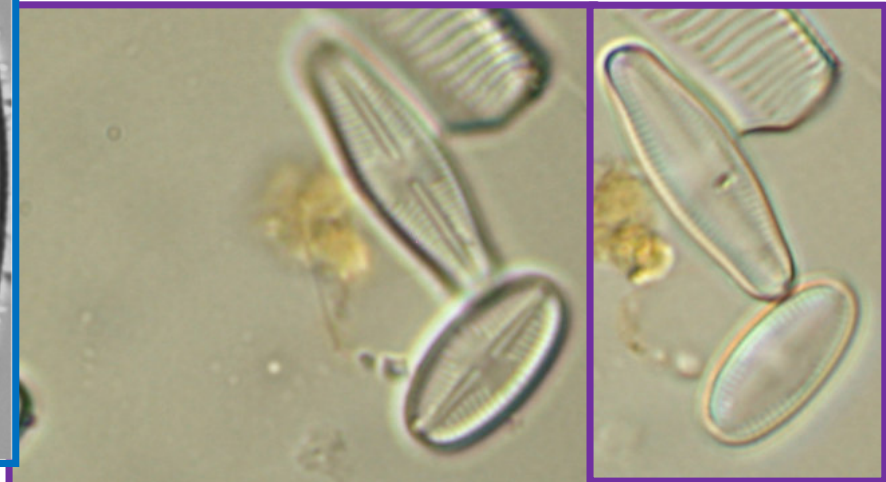
IndVal analysis							
	1	2	3	4			
<b>Achnanthidium.minutissimum</b>	<b>0.53</b>	Diatoma.m	<b>0.72</b>	Rossithid	<b>0.71</b>	Tabellaria.flocculos	<b>0.6</b>
Brachysira.neoexilis	<b>0.33</b>	Syndedra.c	<b>0.56</b>	Fragilaria	<b>0.61</b>	Achnanthidium.krie	<b>0.59</b>
Fragilaria.capucina.complex	0.2	Encyonema	<b>0.43</b>	Eunotia.in	<b>0.51</b>	Psammothidium.ma	<b>0.51</b>
Tabellaria.flocculosa.Complex	0.19	Cymbella.ci	<b>0.35</b>	Staurosire	<b>0.49</b>	Fragilaria.capucina.c	<b>0.41</b>
Brachysira.brebissonii	<b>0.17</b>	Fragilaria.ca	0.23	Eunotia.in	<b>0.49</b>	Brachysira.procera	<b>0.41</b>
Delicata	<b>0.17</b>	Encyonopsi	<b>0.22</b>	Gomphor	<b>0.4</b>	Eunotia.arcus.muco	<b>0.38</b>
Encyonema.silesiacum.minutum.lange.be	0.16	Achnanthid	0.21	Navicula.	<b>0.33</b>	Eunotia.Complex	<b>0.37</b>
Nitzschia.perminuta.Complex	0.13	Nitzschia.pi	<b>0.21</b>	Navicula.	<b>0.32</b>	Frustulia.rhomboide	<b>0.37</b>
Achnanthidium.deflexum.Complex	<b>0.12</b>	Nitzschia.pi	0.2	Karayevia	<b>0.32</b>	Nitzschia.perminuta	<b>0.29</b>
<b>Gomphonema.ariostriatum</b>	<b>0.11</b>	Nitzschia.in	<b>0.19</b>	Psammot	<b>0.28</b>	Brachysira	<b>0.24</b>
Rossithidium.pusillum.anastasiae	0.11						



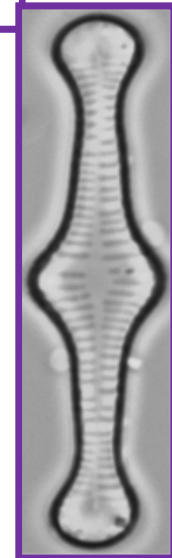
High Arctic biotype



Modern samples

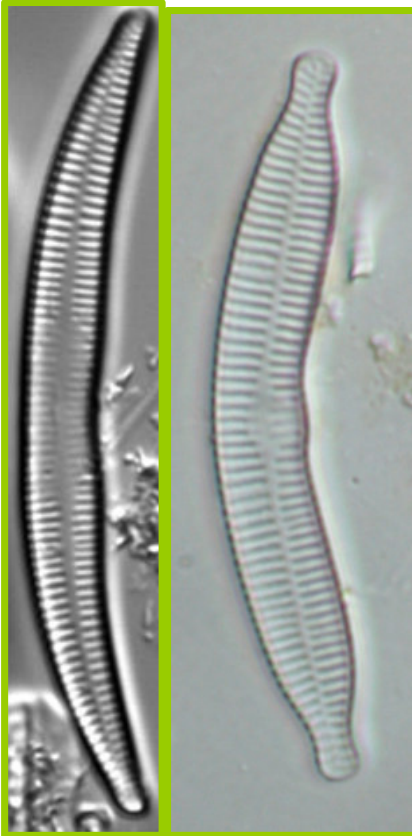


DSCN4668 *Psammothidium marginulatum* & *acidoclinatum*



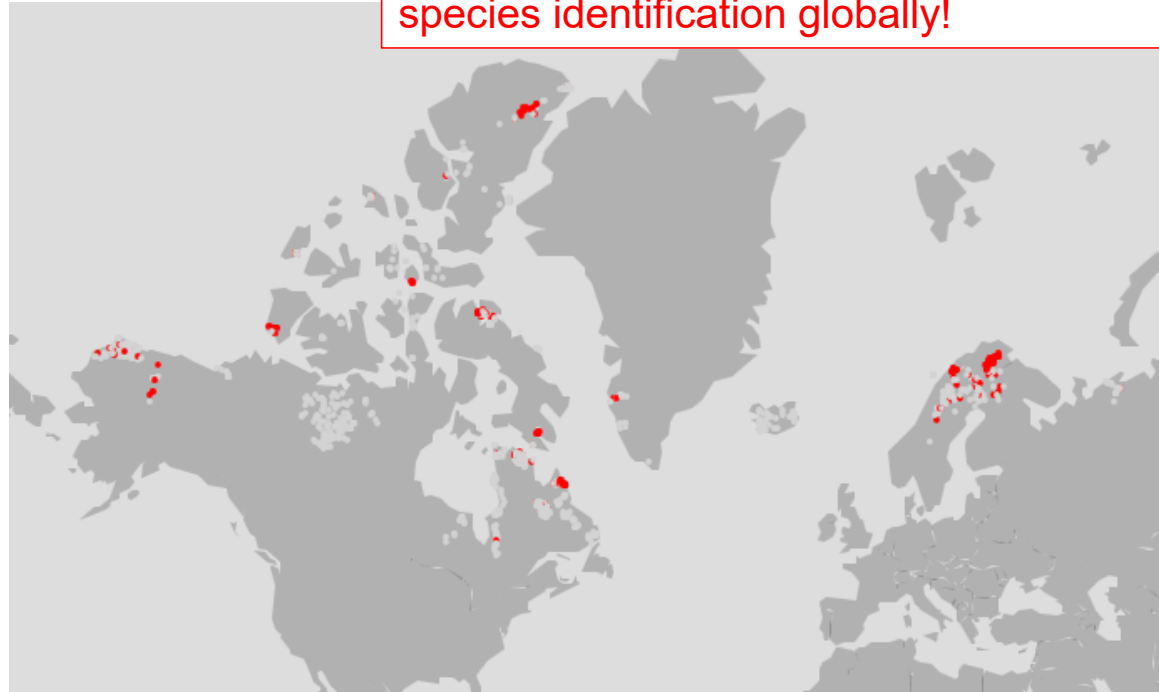
## Distribution of complex *Synedra cyclopus*/*Hannaea arcus*

Global distribution of single taxa not easy to analyse because of the need to pool species -> **we need a more harmonized species identification globally!**



*Synedra  
cyclopus*  
Brutschy

*Hannaea  
arcus*  
(Ehrenb.)  
R.M.Patrick

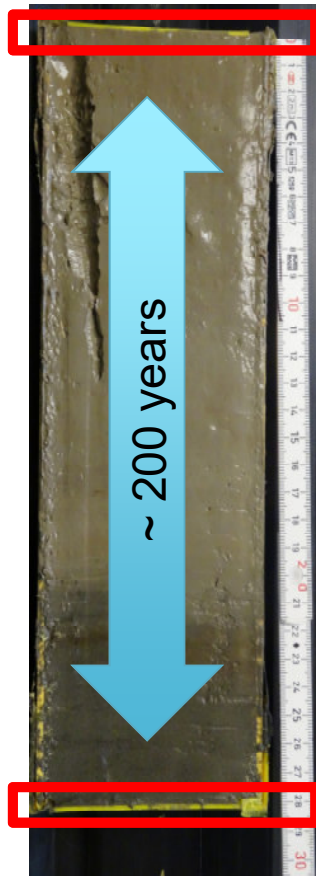


*Synedra cyclopus* is uncommon and widely dispersed in the plankton of lakes and reservoirs (and in streams below lakes and reservoirs) in the western US. Patrick and Reimer (1966) report that it is often attached to crustacea in cool water. In Europe, Krammer and Lange-Bertalot (1991) report it from lakes of medium electrolyte content, where it lives on copepods.

*Hannaea arcus* is found in cool flowing water in mountainous regions across the US and Europe. Typical for streams.

Top/Bottom, n = 116

*Pre-industrial (~200 years)  
versus modern sediments*

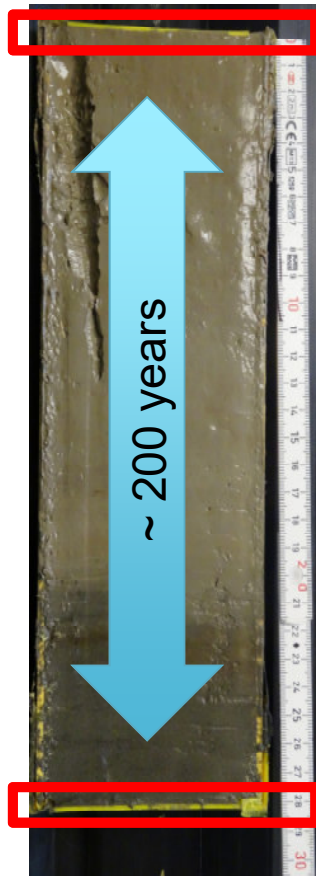


<https://www.bas.ac.uk/project/identification-of-glacial-time-sources-for-antarctic-deep-and-bottom-water-masses/>

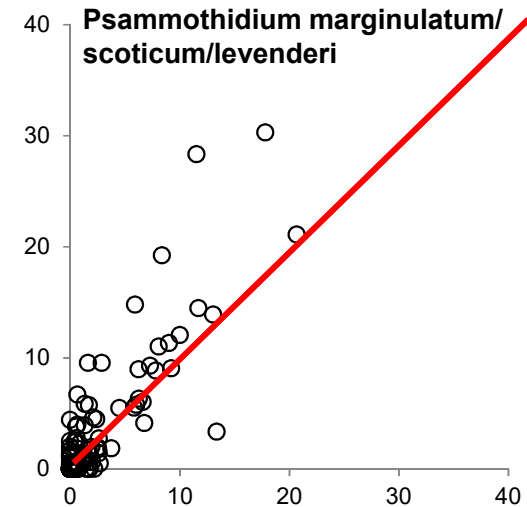
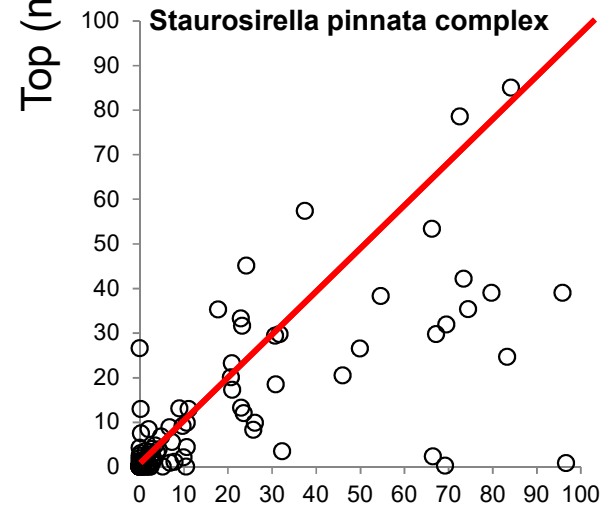
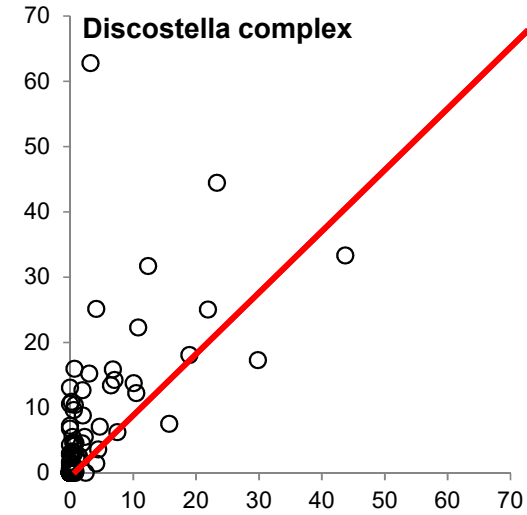
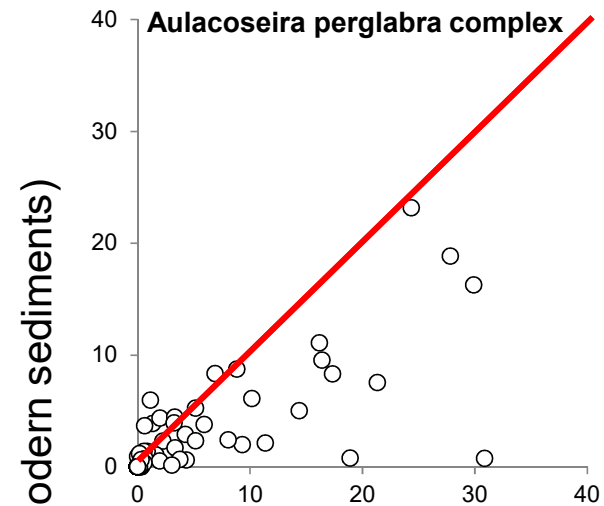


Top/Bottom, n = 116

*Pre-industrial (~200 years)  
versus modern sediments*



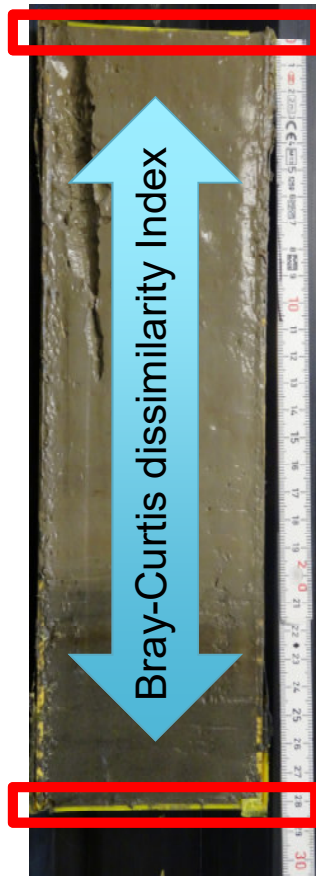
<https://www.bas.ac.uk/project/identification-of-glacial-time-sources-for-antarctic-deep-and-bottom-water-masses/>



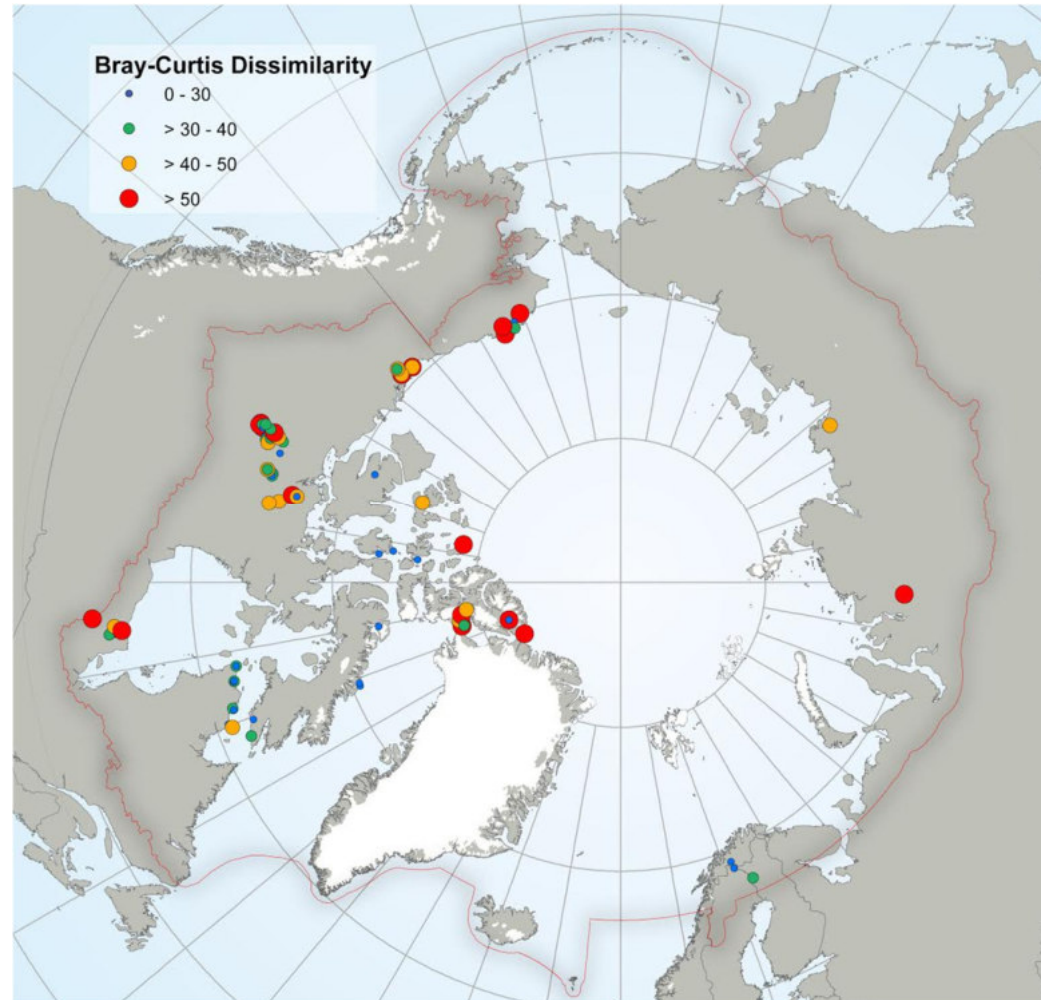
Bottom (pre-industrial sediments)

Top/Bottom, n = 116

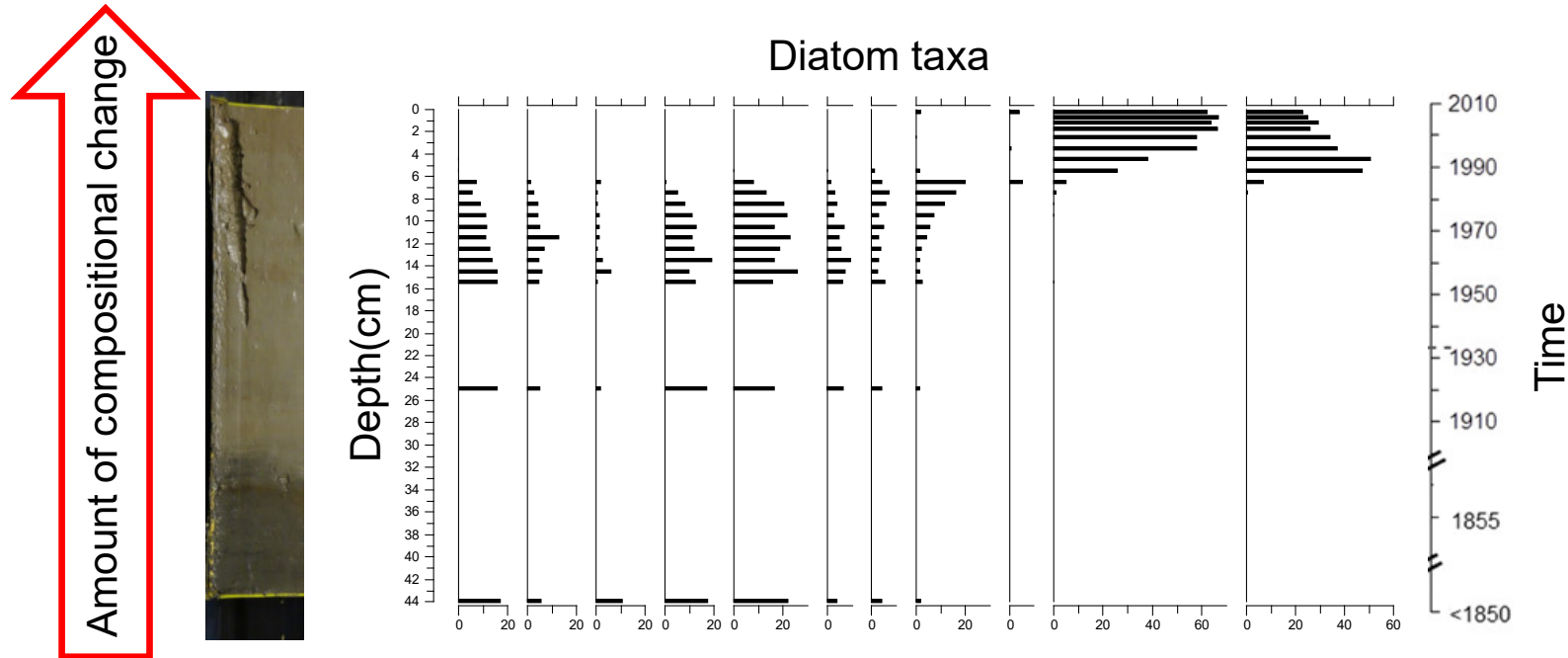
*Pre-industrial (~200 years)  
versus modern sediments*



<https://www.bas.ac.uk/project/identification-of-glacial-time-sources-for-antarctic-deep-and-bottom-water-masses/>



Sediment cores,  $n = 52$  *Pre-industrial (~200 years)*  
*Full cores*



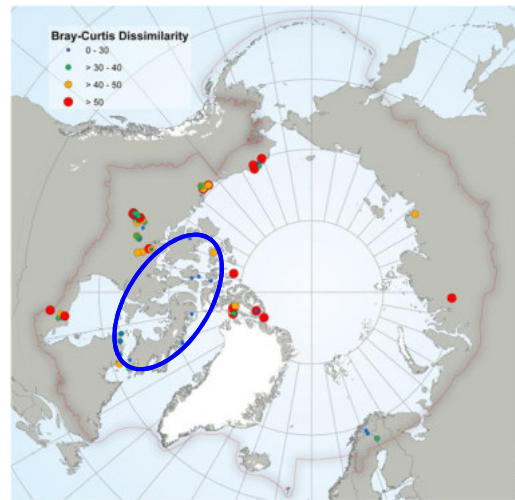
Detrended canonical correspondence analysis (DCCA) to estimate diatom compositional turnover as beta-diversity, scaled in SD units (Smol et al., 2005; PNAS)

# Sediment cores, $n = 52$

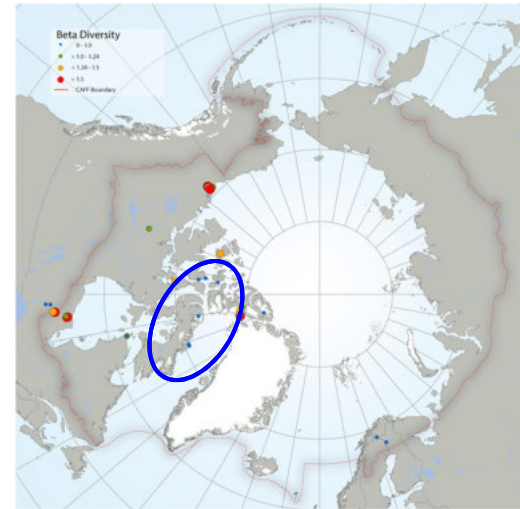
*Pre-industrial (~200 years)*  
*Full cores*



Top/Bottom



Full cores



- Eastern Canada\* least turnover, i.e. maybe least affected by global change
- (!) Results quite general, because community turnover is depending on lake characteristics (depth, area, morphology, etc.)) and core age (*both pooled in this study*)

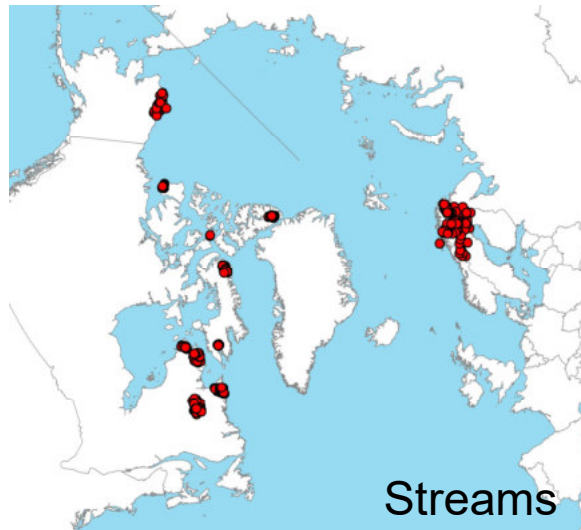
\*and Finland



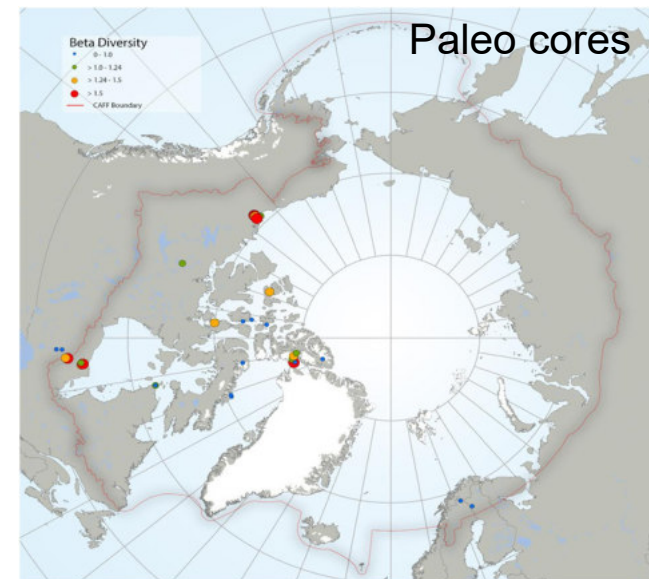
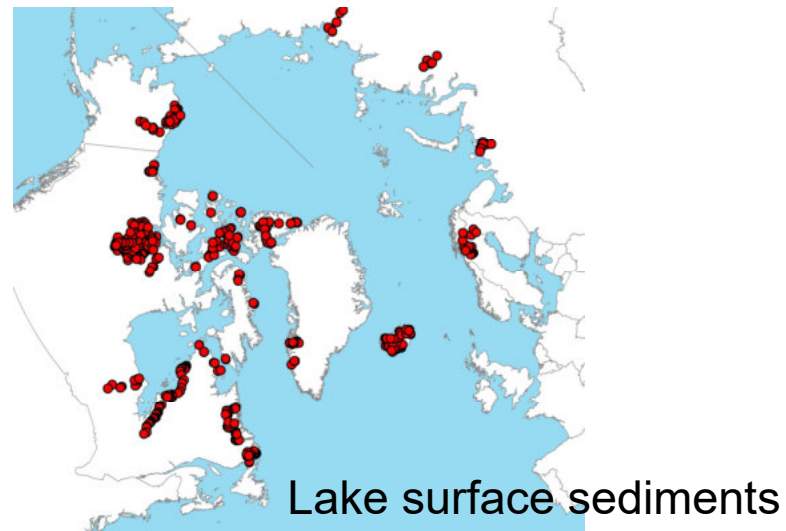
## General conclusions

- *Modern samples:*
  - gradual shifts of taxa across the Arctic
  - still typical community compositions found
  - a **High Arctic** biotype found for both lakes & streams
  - driving factors were: low temp., low precipitation, sparse vegetation, bedrock (high Ca & pH), lake depth, substrate (sand)
- *Pre-industrial era samples:* Changes in lake diatom assemblages least in Eastern Canada (and Finland)  
Causes for change found in other studies:
  - High Arctic: shorter ice cover -> changes in thermal stratification and light penetration
  - Sub-Arctic: changes in the terrestrial vegetation, increase of humic substances (color) and nutrients

## General conclusions



- Clear lack of stream sampling  
Important baseline for future assessment!
- Most paleo cores from Canada, need more from other countries



## General conclusions

- Only Finland and Sweden have established long-term diatom-based monitoring programs (streams only)
- Many research based data available (especially lakes), but those sites not included in a routine follow-up
- Low agreement on diatom species identification -> problems when using data from different countries/ laboratories
  - Need for circumpolar diatom harmonization/identification activities
  - In the future: metabarcoding



More information: [caff.is/freshwater](http://caff.is/freshwater)



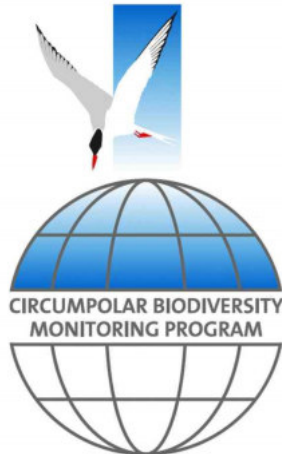
Swedish Agency  
for Marine and  
Water Management

Laboratory for  
Paleoclimatology and  
Climatology  
uOttawa

INRS  
UNIVERSITÉ DE RECHERCHE  
CENTRE  
EAU  
TERRE  
ENVIRONNEMENT

Thank you

NIVA  
Norsk institutt for vannforskning



CAFF  
Conservation of Arctic Flora and Fauna



More information: [caff.is/freshwater](http://caff.is/freshwater)

Points over global mean richness in red.

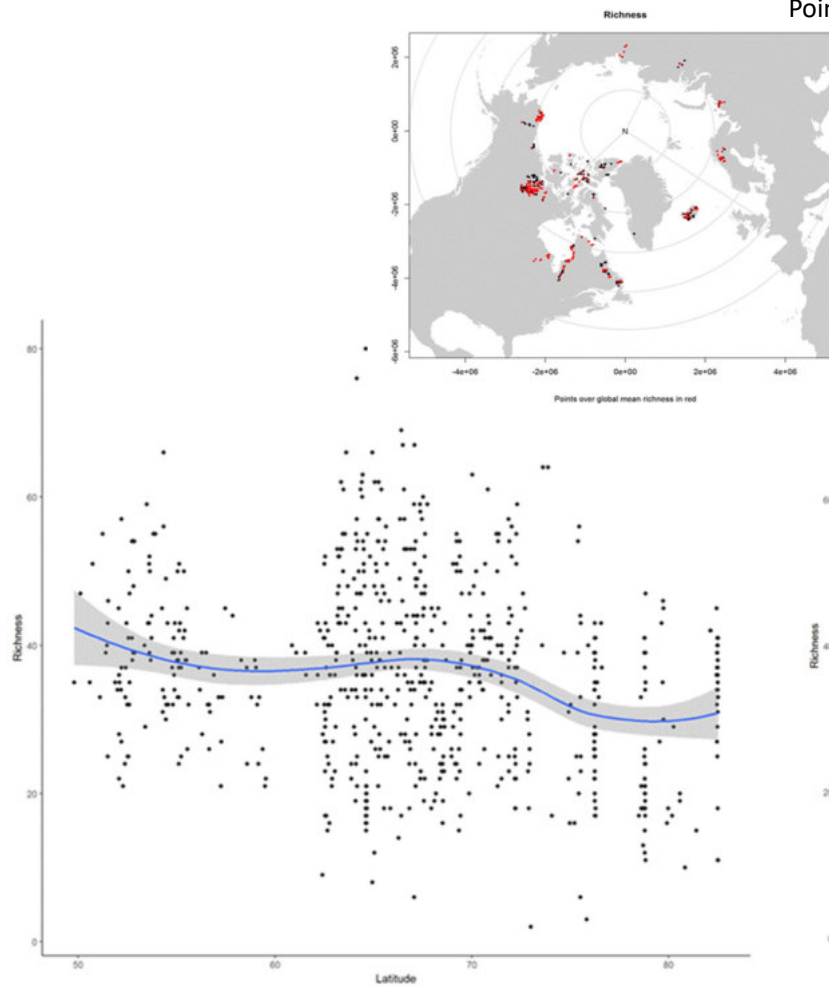


Fig 2b Local species richness of diatom communities in Arctic lakes vs. latitude.

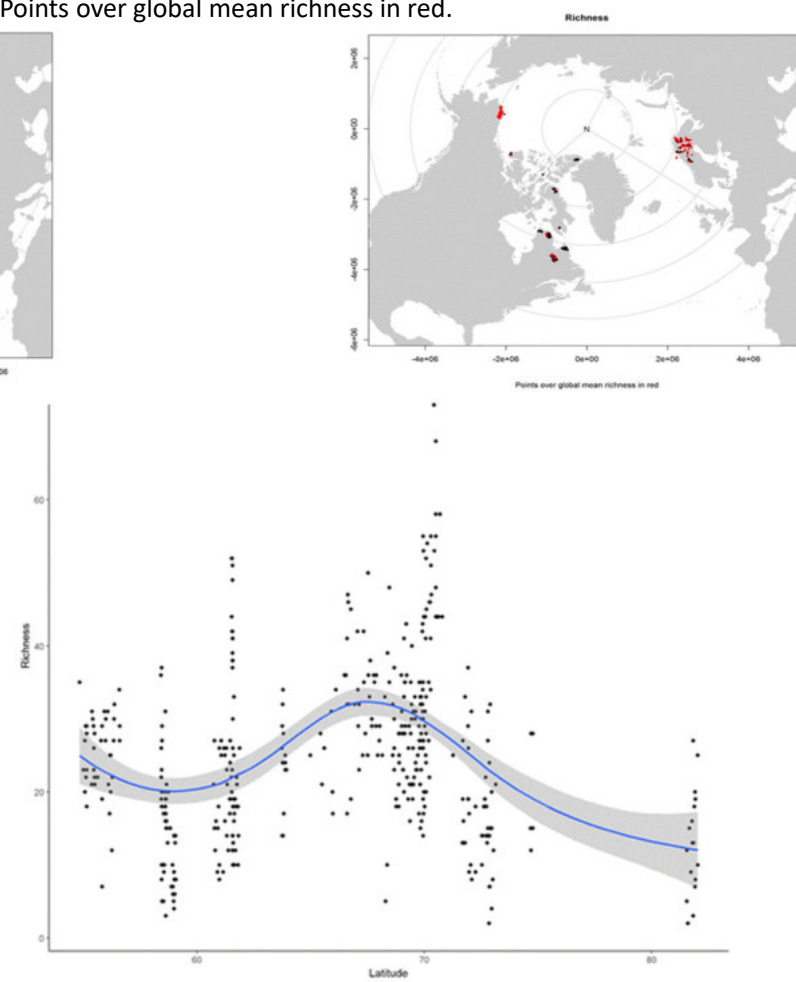


Fig 3b Local species richness of diatom communities in Arctic streams vs. latitude.



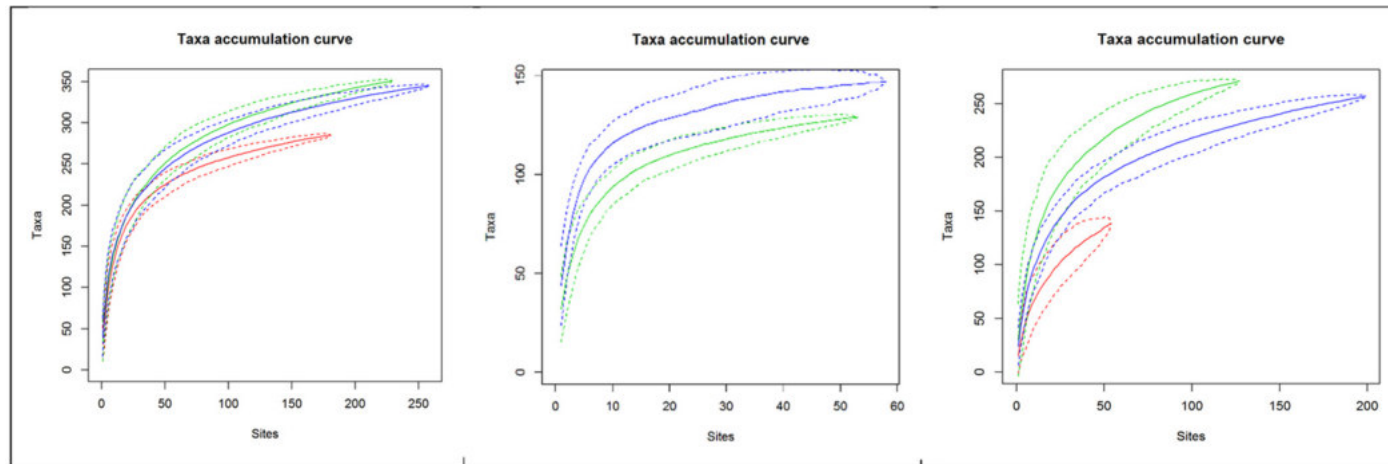


Fig 4. Taxa accumulation curves for the different habitats and arctic zones. Left: Lakes (top sediments), Middle: Lakes (scrapes). Right: Streams. Red: High arctic zone, green: low arctic, blue: sub-arctic.

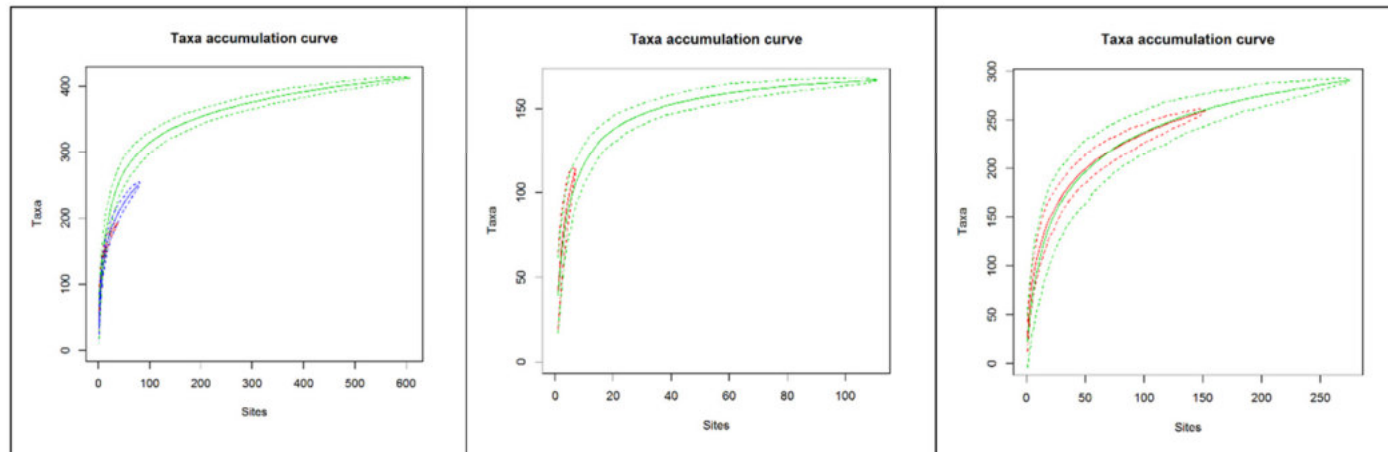


Fig 5. Taxa accumulation curves. Left: Lakes (top sediments), Middle: Lakes (scrapes). Right: Streams. Green: America, red: Europe, blue: Asia.