

### CNL(24)76

### Using eDNA to estimate the distribution of pink salmon

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# What is eDNA?



«Conservation in a cup of water»







### What is eDNA?





# eDNA results and inference



- 1. Presence (NOT absence!)
  - Rivers with unknown or uncertain status
- 2. Distribution
  - Upper limit
  - Tributaries
- 3. Abundance
  - Changes over time?



X Population size, age structure, sex ratio



# NINA and eDNA



River pearl mussel



Bees and pollination



Gyrodactylus salaris



Reindeer and CWD



Y 1 1 1 2 9

Northern pike



Great-crested newts





Insects

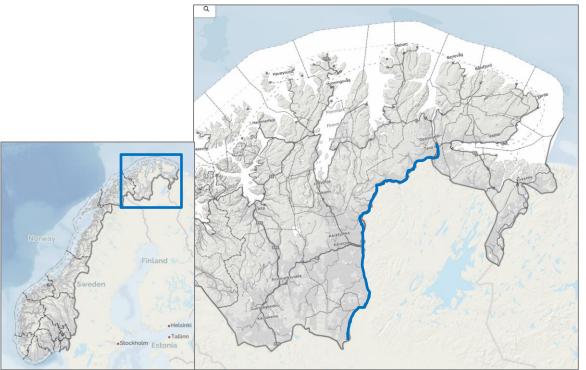


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# Pink salmon and eDNA



- 1. eDNA in the Tana River, Norway
- 2. PINKTRACK

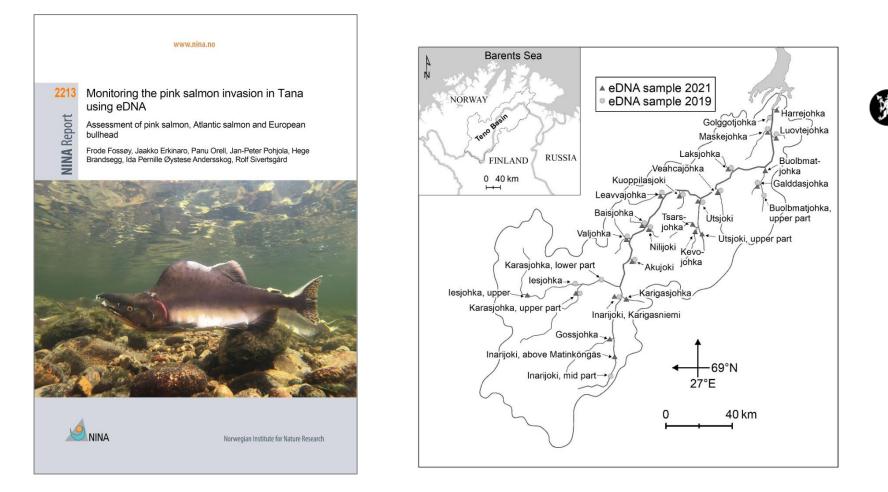








# Tana eDNA sampling



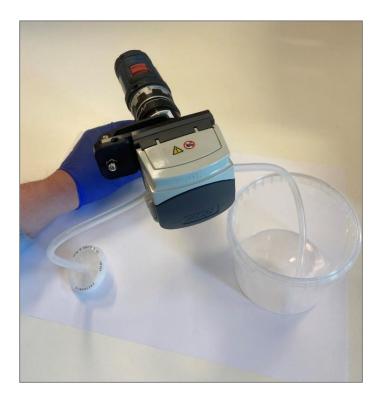
Statsforvalteren i Troms og Finnmark







# Tana eDNA sampling



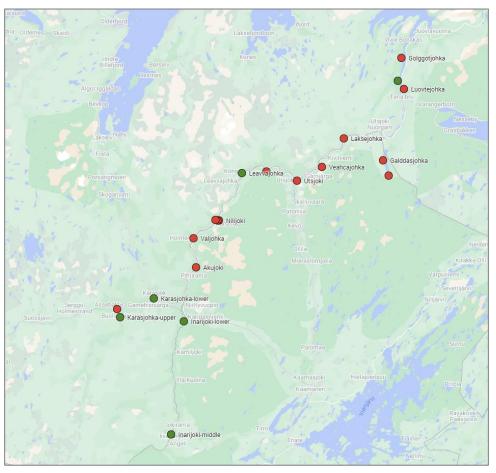
Year	Date	Water volum (L) mean (range)	Temperature (°C) mean (range)
2019	August 12-14	4.9 (3.5-5)	11.1 (8.6-13.8)
2021	August 9-13	5.0	15.2 (11.5-17.7)
2022	August 24-29	4.3 (1.5–5)	12.6 (9.6-15.8)
2023	August 8-17	4.0 (2.5-5)	15.5 (11.9-18.5)

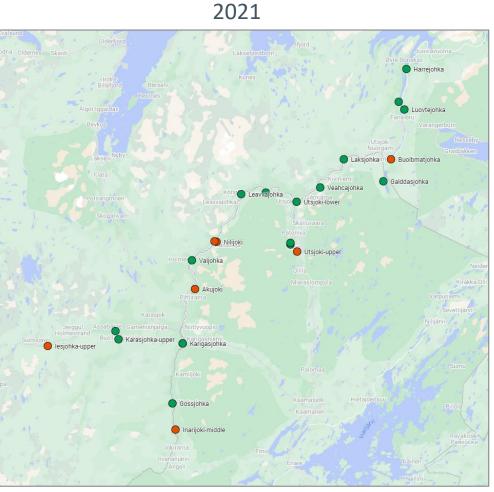


### Tana river – Pink salmon



2019



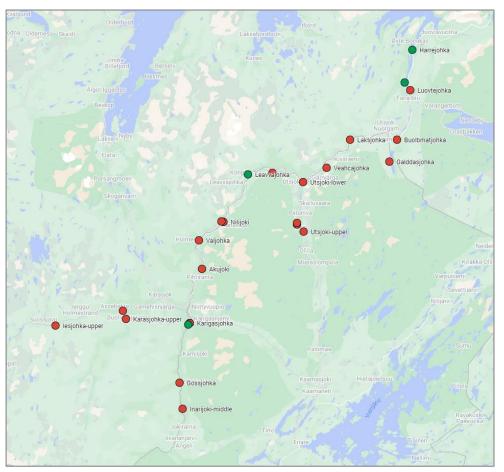


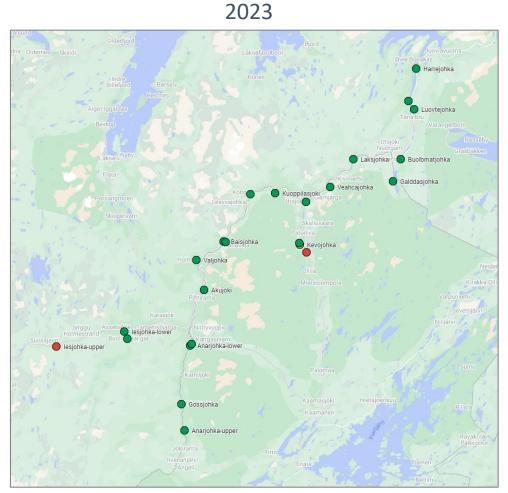




## Tana river – Pink salmon

2022





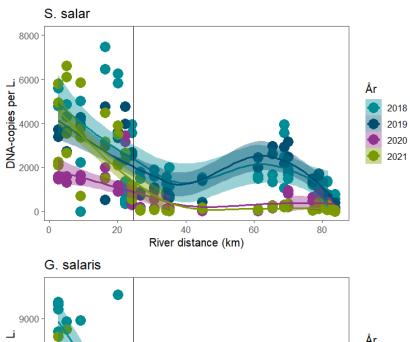
Norwegian Institute for Nature Research

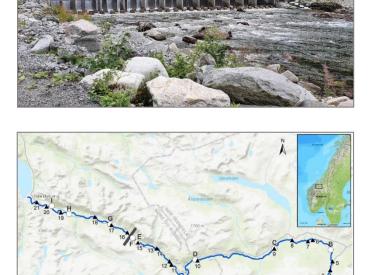
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# eDNA and biomass in river Driva



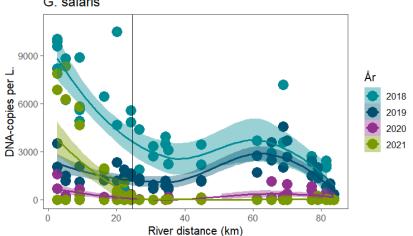




Legend O eDNA-sampling site ▲ Electrofishing sites Migration barrier County borders 5 10

20 KJ

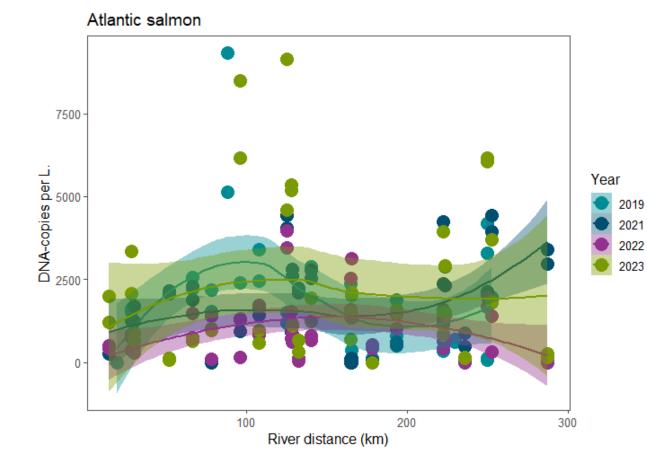








### Tana river – Atlantic salmon





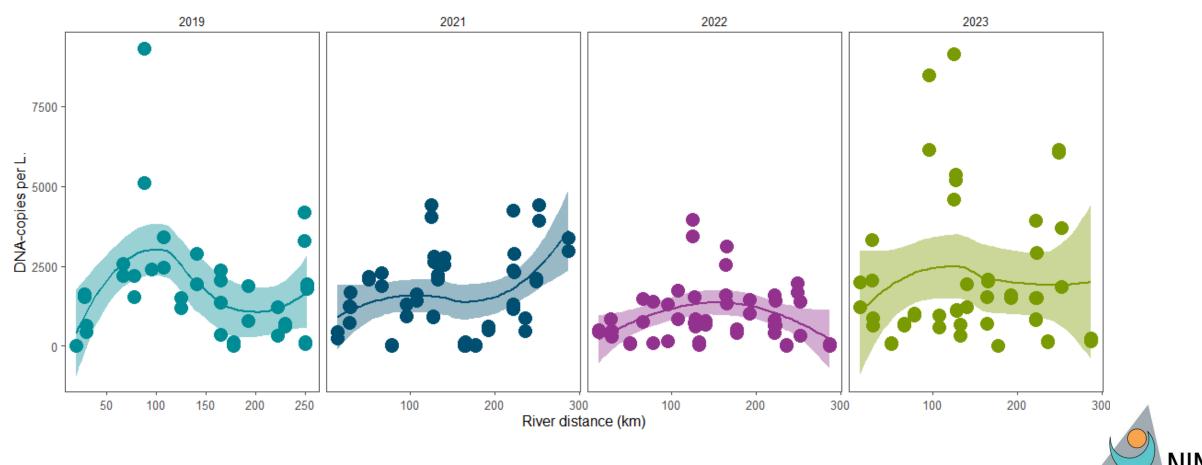
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Norwegian Institute for Nature Research

### Tana river – Atlantic salmon

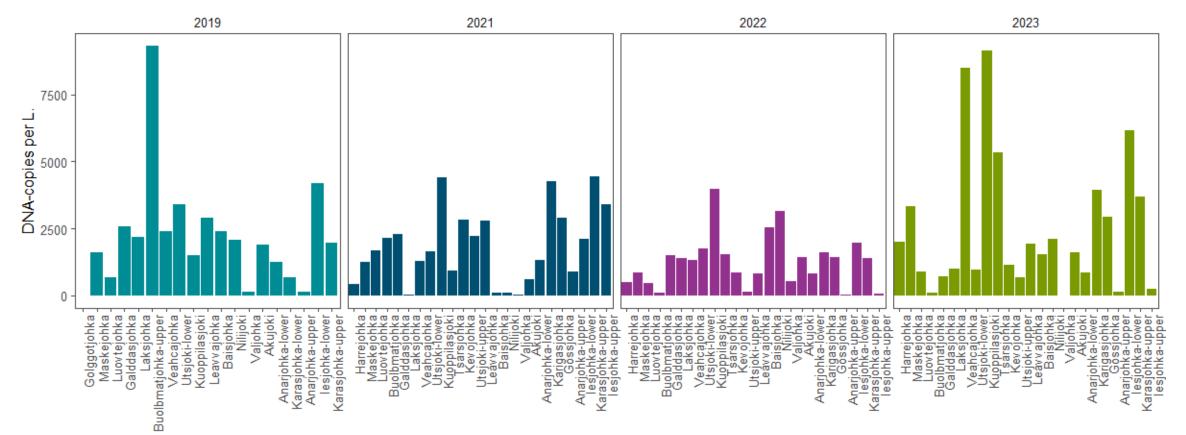
Atlantic salmon





### Tana river – Atlantic salmon

Atlantic salmon

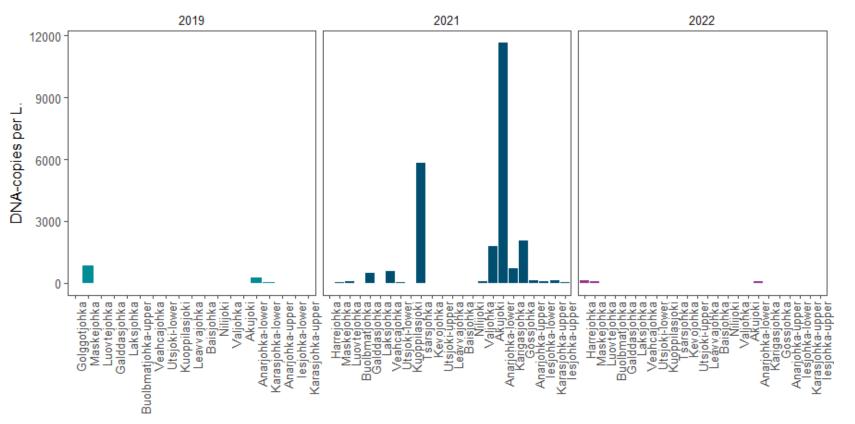




### Tana river – Pink salmon



Pink salmon

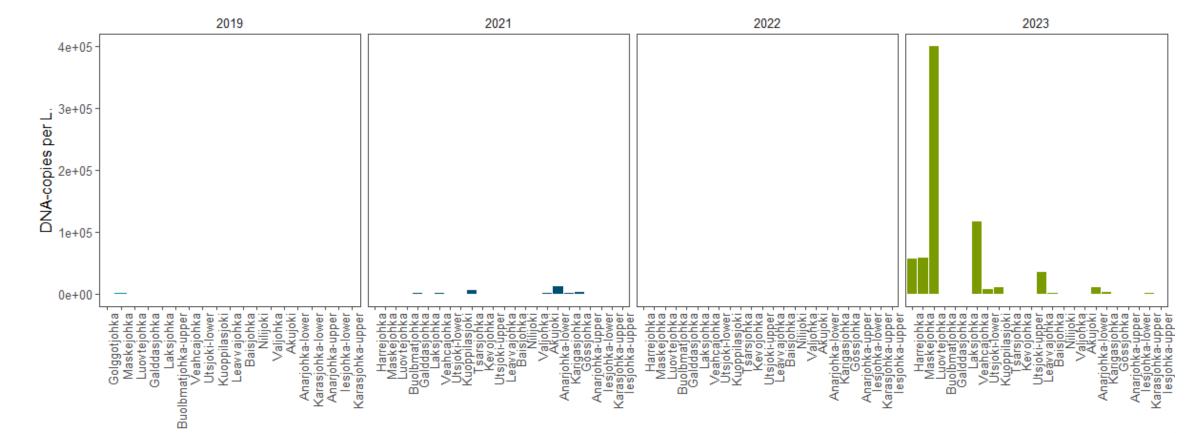








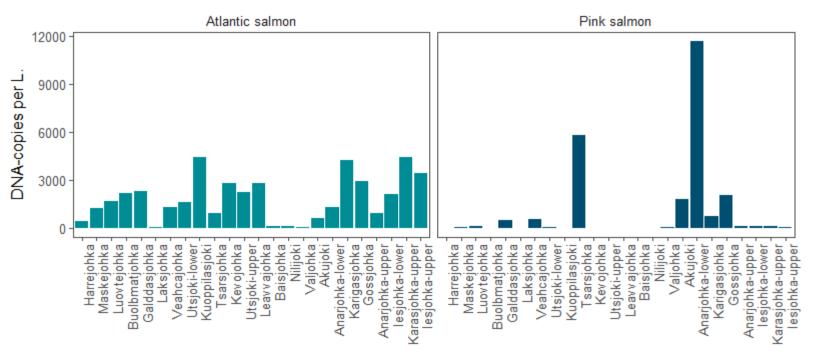
#### Pink salmon







### Tana river – eDNA

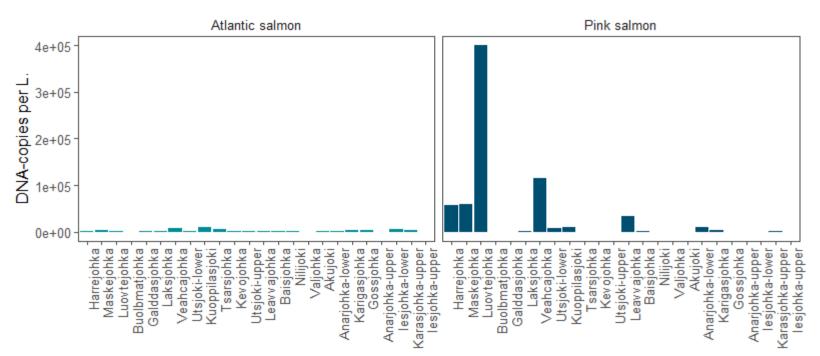


Tana 2021





### Tana river – eDNA



Tana 2023





### Co-ordinated environmental DNA (eDNA) surveillance programme for pink salmon in the EU

#### **Country Nominations**

France: Laurent Beaulaton Sweden: Thomas Staveley Germany: Marko Freese Denmark: Niels Jepsen Finland: Jaakko Erkinaro Ireland: Michael Millane

#### Experts: Norway: Frode Fossøy

Ireland: Jens Carlsson

#### **Observers:**

*Portugal:* Isabel Flores Figueira *Spain:* Julián García Baena

Cathal Gallagher Michael Millane







### • Objective 1 (Work Package 1: eDNATrack)

Development of standardised protocols for eDNA sampling and standardised approaches for the analyses of eDNA samples for the detection of pink salmon in EU MS with the intention that such methods can continue to be utilised in routine national monitoring programmes after the project concludes. This includes preparatory work to evaluate different approaches to sampling and analyses and their effect on the results for detection. Establishment of a repository of eDNA samples collected during the project and in subsequent years to provide valuable material for future assessments as analytical technologies develop.





### Objective 2 (Work Package 2: SurveillTrack)

Establishment and undertaking of an eDNA sampling programme for detection of pink salmon in EU MS in order to elucidate temporal and geographic patterns of spread and provide an 'early warning system' of their presence to inform appropriate management responses with the intention that this programme of work can provide a basis for continued routine national programmes after the project concludes.



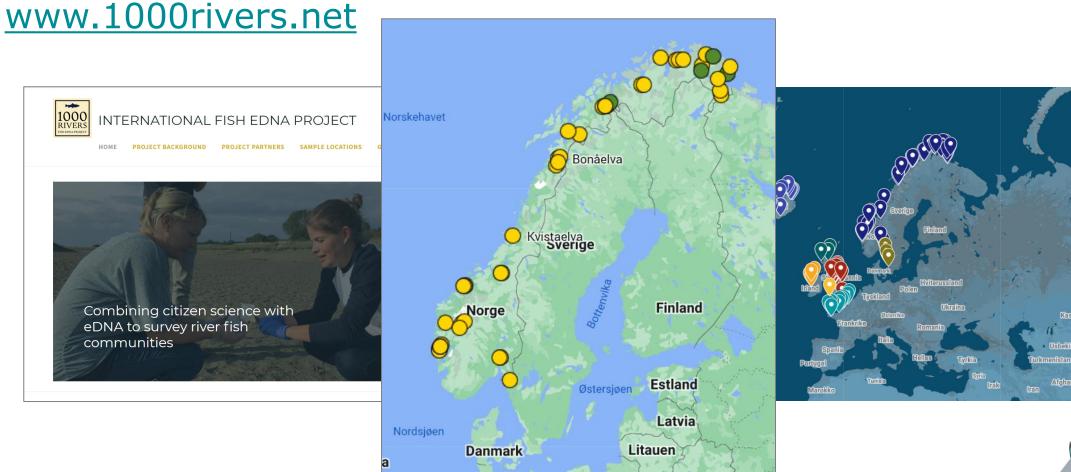


- Objective 3 (Work Package 3: ProjectTrack)
  - Organisation and hosting of a two day virtual project workshop after project commencement, where partners will meet and discuss the development and execution of the sampling programme, protocols and methodologies for sampling and analyses and the dissemination of results. NASCO staff will be encouraged to participate to discuss how NASCO can participate in the dissemination activities. A project wrap-up virtual workshop will also be held as the project concludes to review outputs and inform their further adoption by EU MS.





### 1000 rivers project – citizen science

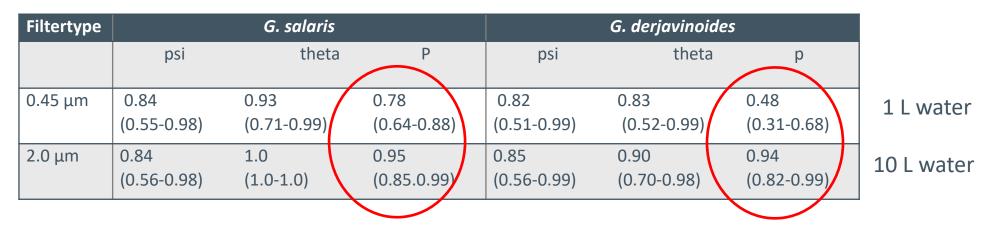




# Gyrodactylus salaris in Driva



### eDNA occupancy modelling





Frode Fossøy 回 Hege Brandsegg Rolf Sivertsgård | Oskar Pettersen | Brett K. Sandercock | Øyvind Solem | Kjetil Hindar | Tor Atle Mo





DNA

# PINKTrack - filtertest



### 2024:

- Objective 1 (Work Package 1: eDNATrack)
  - Development of standardised protocols for eDNA sampling and standardised approaches for the analyses of eDNA samples for the detection of pink salmon

Filter type	Pore siz	e Membrane
Sterivex	0.45	PES
Merck millipore glassfiber	2.0	Glass fiber
Waterra	1.2	PES
Sylphium	0.8	PES
Sylphium	5.0	PES
Smith-Root glassfiber	1.5	Glass fiber
Smith-Root self preserving	1.2	PES



# PINKTrack - filtertest



- 6 countries
  - Norway, Sweden, Denmark, Ireland, Germany and France
- 2 rivers and 4 samples per filter per country
- 7 filter types: 24 filters per type across 6 countries
- 3 labs analysing the samples from 2 countries each



# PINKTrack - filtertest



- DNA detection and concentration will be modelled
  - Filter type
  - Country/river
  - Lab
  - River size and flow
  - Water volume
  - Water turbidity
  - Water temperature
  - Water pH

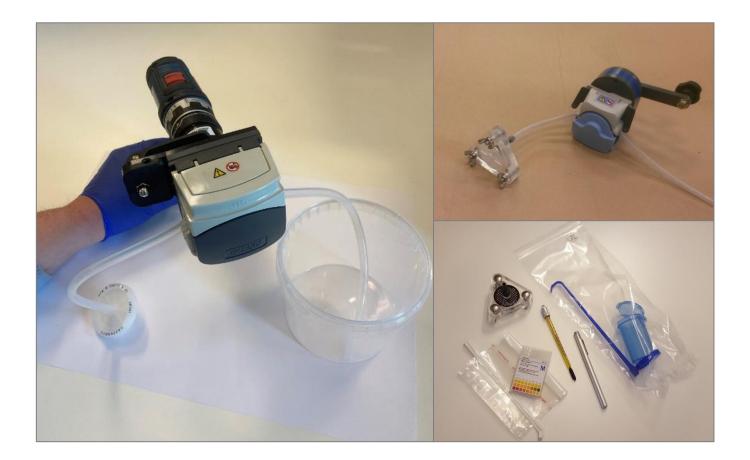






## NINA eDNA kit

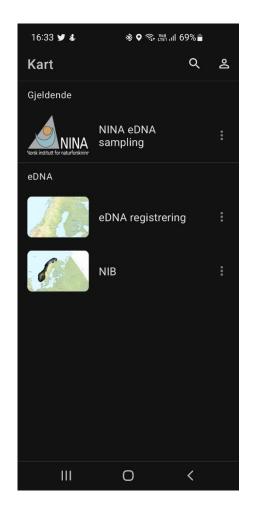


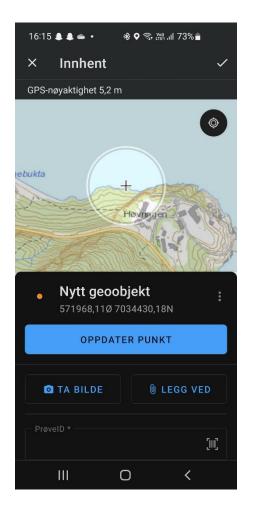






### NINA eDNA phone app





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Prøvetype * Water		$\otimes$
Lokalitetstype * Marine		$\otimes$
Filternavn * NatureMetrics Gf	5.0/PES 0.8	3µr ⊗
Innsamlingsmetode *		$\otimes$
Prøvetaker *	0	<



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- Move from monitoring status to consequence
   DNA samples are time-capsules of biodiversity
- Ecological status reuse eDNA samples
   Fish diversity
  - Benthic macro-invertebrates
  - Benthic algae
  - Key species such as freshwater pearl mussels



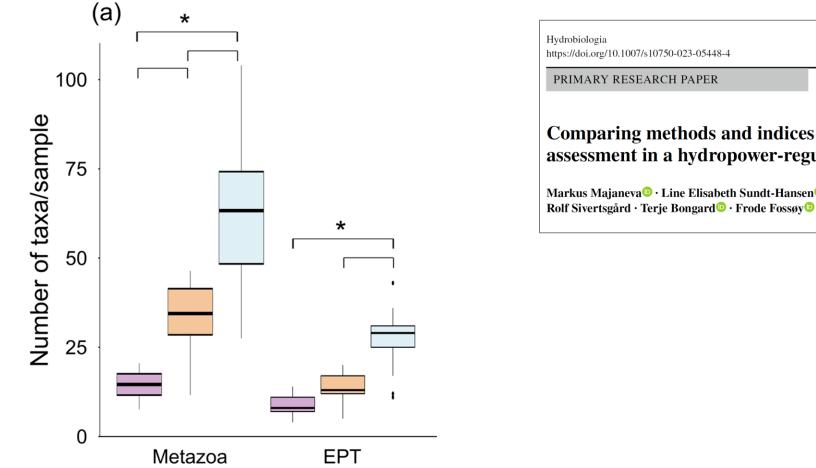


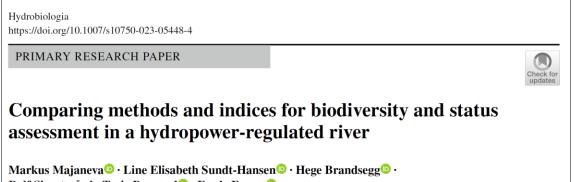






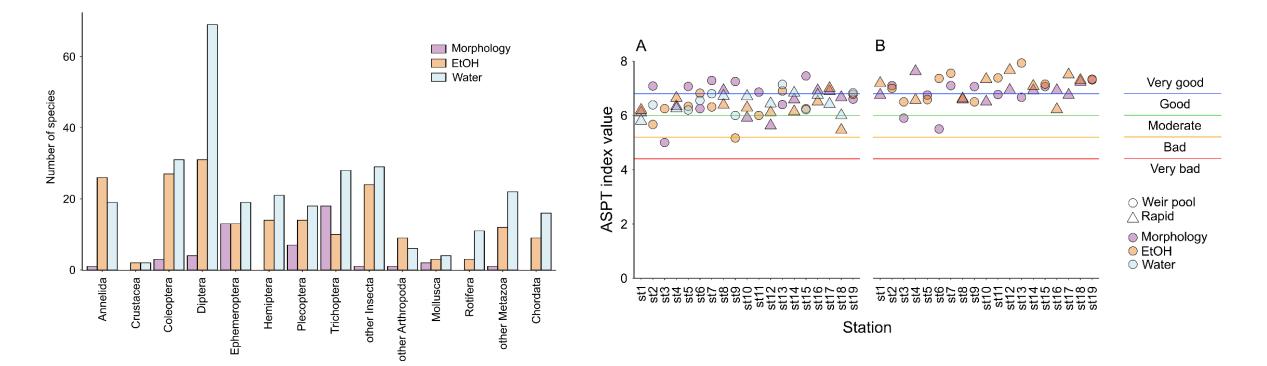
















(a) (b) 1.25 High nEQR (ASPT index) 1.00 Good Moderate  $\bigcirc$ 0.75 Poor Bad 0.50 0.25 + Average nEQR  $\bigcirc$  Weir pool  $\triangle$  Rapid 0.00 Morphology
 EtOH
 Water (d) (c) 1.25 nEQR (IBIBI index) 1.00 \* \* High 0.75 Good  $\frown$ Moderate 0.50 Poor 0.25 0 Bad  $\bigcirc$  $\cap$ 0.00 st12 st13 st15 st15 st15 st16 st17 st18 st18 st1 st2 st5 st5 st5 st7 st8 st9 st10 £ Station



ASPT





