

Nye metoder for miljøovervåking for fremtidens oppdrett

Nigel Keeley, Kathy Dunlop, et al.

Havforskningsinstituttet



NS: 9410:2016

Analytical types

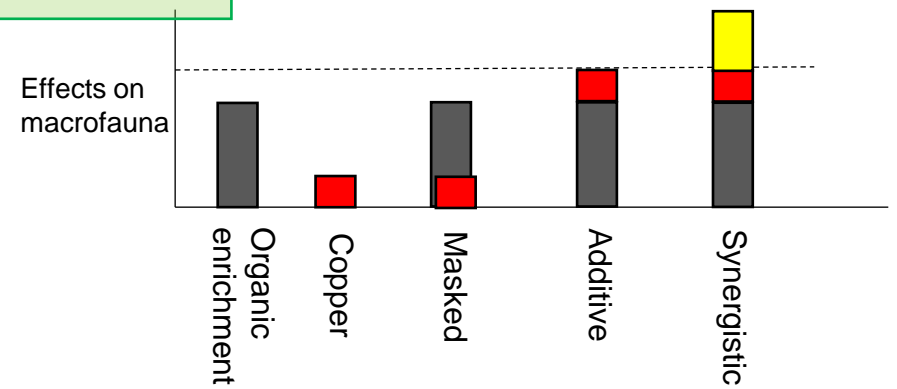
		B-investigation					C-investigation											
Ecological compartment		Sensory (Farge, luft, konsistens etc.)					Macrofauna					Kjemiske parametere (in sediments)						
Discharge type		Farge	Luft	Konsistens	Grabvolum	Tykkelse	Redox /pH	S	N	NQ1	NSH'	TOM	nTOC	N	C:N	P	Zn	Cu
Soft sediment footprint																		
	Organic enrichment	~	~	~		~	✓/?	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓
	Biofouling drop off			~				~	~	~	~	~	~	~	~	~	~	~
	Antifouling chemicals: Cu							~	~	~	~	~	~	~	~	~	~	✓
	All new products							?										
	Infeed therapeutants Zn							~	~	~	~	~	~	~	~	~	~	✓
	All others							?										
	New feed composition (eg terrestrial)							?						~				
	Bath treatments							?										

✓ = Quantitative and useful
 ~ = Qualitative / some relevance
 ? = Potential relevance / not well understood

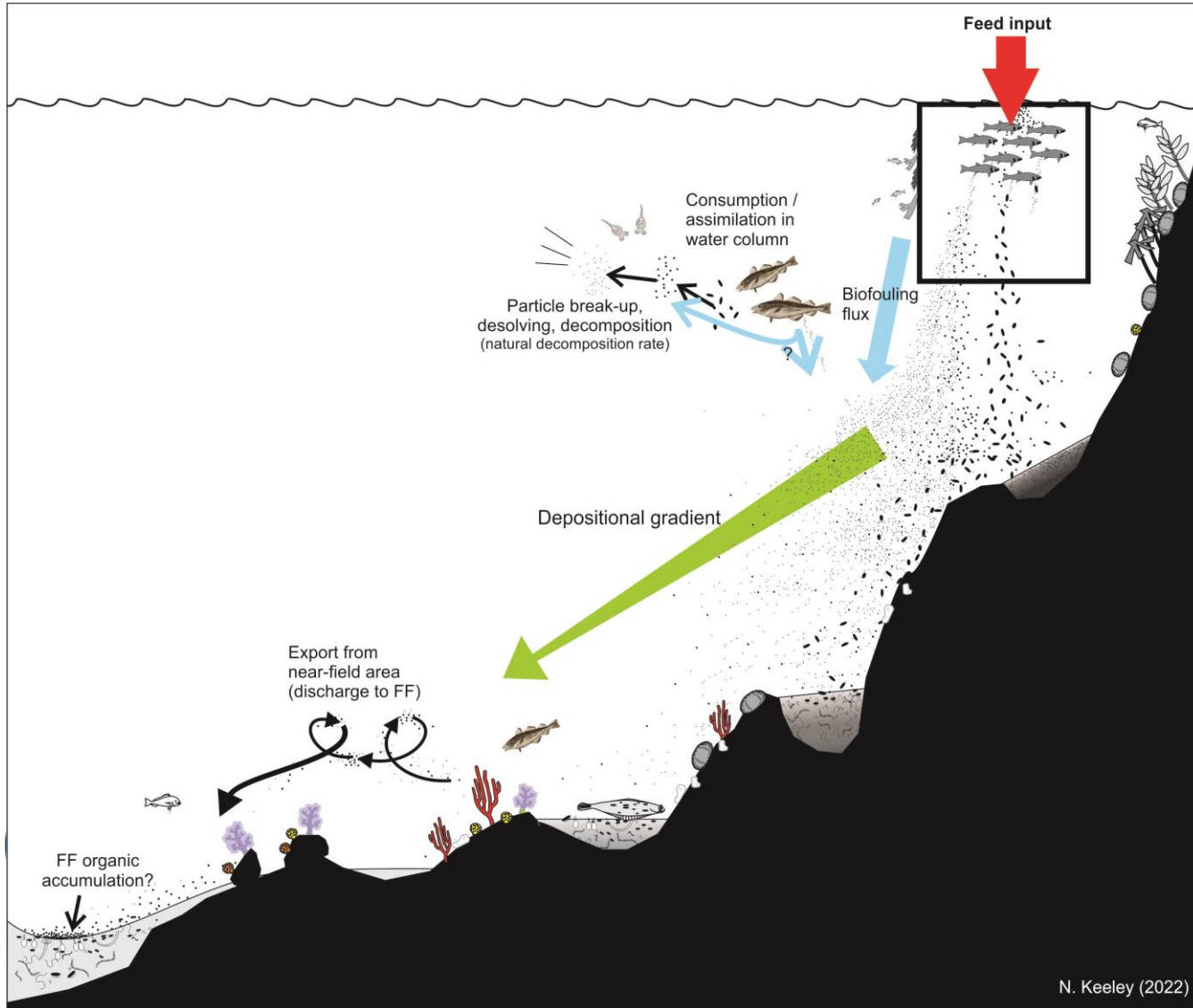


Additional /independent /synergistic effects?

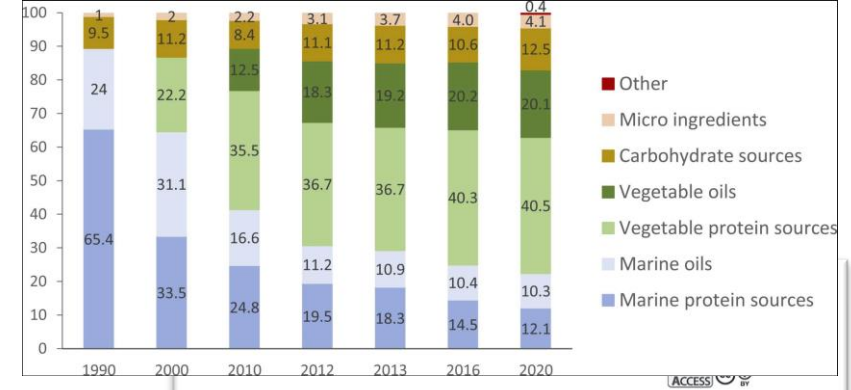
Lacking (W.I.P.)



Avfallssporingsmetoder



N. Keeley (2022)



FEATURE ARTICLE

Combining biochemical methods to trace organic effluent from fish farms

S. H. Woodcock^{1,*}, C. Troedsson^{2,3,4}, T. Strohmeier¹, P. Balseiro^{2,4}, K. Sandnes Skaar¹, Ø. Strand¹

¹Nordnesgaten 50, 5065 Bergen, Norway
²Jormhøllensgate 55, 5008 Bergen, Norway
³veien 232, 5258 Blomsterdalen, Norway
⁴Box 7803, Thormøhlensgate 53 A/B, 5006 Bergen, Norway

Ecological Indicators
 journal homepage: www.elsevier.com/locate/ecolind

Fate and longevity of terrestrial fatty acids from caged fin-fish aquaculture in dynamic coastal marine systems
 S.H. Woodcock¹, S. Meier, N.B. Keeley, R.J. Bannister
 Institute of Marine Research, Postboks 1870 Nordnes, 5817 Bergen, Norway

ABSTRACT
 Fin fish production continues to expand with the switch in diet from traditional fish meals and oils to terrestrial sourced components. Terrestrial crops can be traced into the marine environment due to their high abundance of C18 fatty acids (FA). Here we compare two FA extraction techniques on coastal sediments: Folch lipid extraction and direct methylation. We found differences in the FA classes which could be extracted using these methods and that direct methylation extracted more of the terrestrial biomarkers. We applied the direct methylation method to detect organic waste dispersal from a fin-fish farm along a transect in sedimentation and sediment samples across a production cycle, along with polychaetes collected during peak production. We traced organic waste from the fin-fish farm as far as 1100 m by mid production. Furthermore, we found polychaetes to have FA profiles reflective of the sediment, indicating they are assimilating aquaculture waste.

REVIEWS IN Aquaculture

Reviews in Aquaculture (2019) 11, 133–148

doi: 10.1111/raq.12230

Terrestrial fatty acids as tracers of finfish aquaculture waste in the marine environment

Camille A. White^{1,2}, Skye H. Woodcock¹, Raymond J. Bannister² and Peter D. Nichols⁴

- ¹ Institute for Marine and Antarctic Studies, University of Tasmania, Hobart, TAS, Australia
- ² Sustainable Aquaculture Laboratory - Temperate and Tropical, School of Biosciences, University of Melbourne, Parkville, Vic., Australia
- ³ Institute of Marine Research, Bergen, Norway
- ⁴ Oceans and Atmosphere, Commonwealth Scientific and Industrial Research Organization, Hobart, TAS, Australia

Correspondence

Camille A. White, Institute for Marine and Antarctic Studies, University of Tasmania, Private Bag 49, Hobart TAS 7000, Australia.
 Email: camille.white@utas.edu.au

Received 11 July 2017, accepted 30 October 2017.

Abstract

Waste from open-cage aquaculture flows directly into the marine environment from uneaten feeds, faecal material and dissolved nutrients. Sustainable management outcomes are regularly based on the dispersal patterns of the waste, with biochemical tracing a key tool in understanding the footprint of aquaculture. We examined the use of fatty acid (FA) analysis to trace aquaculture waste for this purpose, with the aim of identifying specific biomarkers for environmental applications, as well as identifying challenges that are regularly encountered. Overall, the widespread use of terrestrial-based oils in the production of marine aquaculture feeds has increased the use of FA biomarkers to trace aquaculture waste across benthic and pelagic systems, in vertebrates, invertebrates and environmental samples such as sediment and seston. A combination of linoleic acid (LA), oleic acid (OA) and γ -linolenic acid (ALA), which are dominant C₁₈ FA in terrestrial seed and animal oils, is the most commonly used biomarkers, along with

*Corresponding author.
 E-mail address: skye.woodcock@utas.edu.au (S.H. Woodcock).

<https://doi.org/10.1016/j.ecolind.2019.03.007>
 Received 1 December 2018; Received in revised form 27 March 2019; Accepted 30 March 1470-166X/© 2019 Published by Elsevier Ltd.

Avfallssporingsmetoder

Tracing feed signatures into environment

- Terrestrial fatty acids (LA, OA, ALA)
- Soy DNA (other candidates?)
- Stable isotopes ($\delta N:\delta C$)
- Others? – new feed compositions...

Ecological Indicators 103 (2019) 43–54

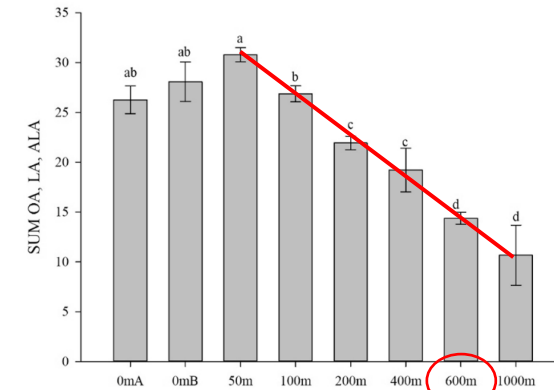


Fig. 7. Comparison of sum of OA, LA and ALA (average \pm standard error) of infauna polychaetes sampled during peak production along a transect moving away from a fin-fish farm. Different letters represent significant difference in the sum of OA, LA and ALA.

Sediments (sinks)
Bivalves
Crustaceans
Echinoderms
Fish

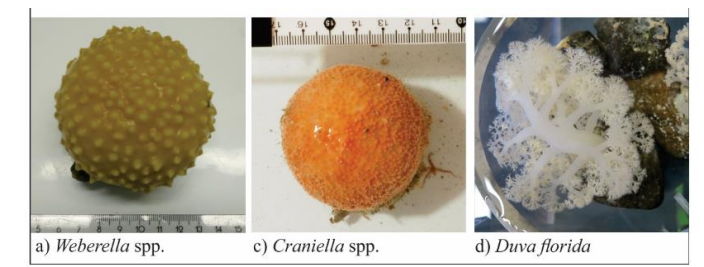
...

Corals
Sponges

...



×/?



Benthic microbial eDNA – sediments

Dowle, E., Pochon, X., Keeley, N., Wood, S.A., 2015. Assessing the effects of salmon farming seabed enrichment using bacterial community diversity and high-throughput sequencing. *FEMS Microbiology and Ecology* 91.

‘Sustain Aqua’: NFR No. 267829

Received: 21 September 2020 | Revised: 31 March 2021 | Accepted: 4 May 2021
DOI: 10.1111/1375-0998.13426

ORIGINAL ARTICLE

MOLECULAR ECOLOGY
RESOURCES WILEY

Beyond taxonomy: Validating functional inference approaches in the context of fish-farm impact assessments

Olivier Laroche^{1,2} | Xavier Pochon^{2,3} | Susanna A. Wood² | Nigel Keeley^{1,2}

frontiers
in Microbiology

ORIGINAL RESEARCH
published: 29 April 2021
doi: 10.3389/fmicb.2021.637811



Global Trends of Benthic Bacterial Diversity and Community Composition Along Organic Enrichment Gradients of Salmon Farms

Larissa Frühe¹, Verena Dully¹, Dominik Forster¹, Nigel B. Keeley^{2,3}, Olivier Laroche², Xavier Pochon^{2,4}, Shawn Robinson⁵, Thomas A. Wilding¹ and Thorsten Stoeck^{1*}

frontiers
in Marine Science

ORIGINAL RESEARCH
published: 14 April 2021
doi: 10.3389/fmars.2021.627687



A Substrate-Independent Benthic Sampler (SIBS) for Hard and Mixed-Bottom Marine Habitats: A Proof-of-Concept Study

Nigel Keeley^{1*}, Olivier Laroche¹, Murray Birch² and Xavier Pochon^{2,4}

Aquaculture Environment Interactions Published January 23, 2025
Volume 17:1–20
https://doi.org/10.3354/aei00499



Disaggregation rates of salmon feces and microbial inoculation of sediments: new insight for particle dispersion modelers

Nigel Keeley^{1*}, Katherine Dunlop¹, Olivier Laroche², Ellie Watts¹, Pål Sævik³, Jon Albretnsen³

Wood, S.A., Tremblay, L.A., Laroche, O., Pochon, X., Lear, G., Ellis, J.I., 2018. Incorporating molecule-based functional and co-occurrence network properties into benthic marine impact assessments. *FEMS Microbiology Ecology* 94.

Cordier, T., Alonso-Sáez, L., Apothéloz-Perret-Gentil, L., Aylagas, E., Bohan, D.A., Bouchez, A., Chariton, A., Creer, S., Frühe, L., Keck, F., Keeley, N., Laroche, O., Leese, F., Pochon, X., Stoeck, T., Pawlowski, J., Lanzén, A., 2020. Ecosystems monitoring powered by environmental genomics: a review of current strategies with an implementation roadmap. *Molecular Ecology*, 1-22.

Pochon, X., Wood, S., Atalah, J., Laroche, O., Zaiko, A., Keeley, N., 2020. A validated protocol for benthic monitoring of New Zealand's salmon farms using environmental DNA.

Pawlowski, J., Esling, P., Martins, C.I.M., Kvalvik, A., Staven, K., 2018. Salmon farms in Norway using foraminifera. *Marine Interactions* 8, 371-380.

Cordier, T., Esling, P., Martins, C.I.M., Frühe, L., Pawlowski, J., 2017. Environmental DNA metabarcoding of benthic bacterial communities indicates the benthic footprint of salmon aquaculture. *Marine Pollution Bulletin* 127, 139-149.

Cordier, T., Forster, D., Dufresne, Y., Martins, C.I.M., Stoeck, T., Pawlowski, J., 2018. Supervised machine learning outperforms taxonomy-based environmental DNA metabarcoding applied to biomonitoring. *Molecular Ecology Resources* 0.

Pearman, J.K., Keeley, N., Wood, S.A., Laroche, O., Zaiko, A., Thomson-Laing, G., Biessy, L., Atalah, J., Pochon, X., 2020. Comparing sediment DNA extraction methods for assessing organic enrichment associated with marine aquaculture. *PeerJ* 8:e10231.

Frühe, L., Dully, V., Forster, D., Keeley, N.B., Laroche, O., Pochon, X., Robinson, S., Wilding, T.A., Stoeck, T., 2021. Global Trends of Benthic Bacterial Diversity and Community Composition Along Organic Enrichment Gradients of Salmon Farms. *Frontiers in Microbiology* 12.

‘AQUAed’: NFR No. 320076

Marine Genomics 66 (2022) 100991



Contents lists available at ScienceDirect

Marine Genomics

journal homepage: www.elsevier.com/locate/margen



Bimodal distribution of seafloor microbiota diversity and function are associated with marine aquaculture

R. Pettersen^a, I. Ormaasen^b, I.L. Angell^b, N.B. Keeley^c, A. Lindseth^d, L. Snipen^b, K. Rudi^{b,*}

Phillip et al. *BMC Bioinformatics* (2024) 25:237
<https://doi.org/10.1186/s12859-024-05837-z>

BMC Bioinformatics

RESEARCH

Open Access

METASEED: a novel approach to full-length 16S rRNA gene reconstruction from short read data



Melcy Phillip¹, Knut Rudi¹, Ida Ormaasen¹, Inga Leena Angell¹, Ragnhild Pettersen², Nigel B. Keeley³ and Lars-Gustav Snipen¹

ISME

ISME Communications, 2024, 4(1), ycae071
<https://doi.org/10.1093/ismeco/ycae071>
Advance access publication: 9 May 2024
Original Article

Swarm and UNOISE outperform DADA2 and Deblur for denoising high-diversity marine seafloor samples

Tonje Nilsen¹, Lars-Gustav Snipen¹, Inga Leena Angell¹, Nigel Brian Keeley², Sanna Majaneva³, Ragnhild Pettersen³, Knut Rudi^{1,*}

¹Faculty of Chemistry, Biotechnology and Food Science (CBM), Norwegian University of Life Sciences (NMBU), Chr. M. Falnesvei 1B, Biotechnolog Building

Nilsen T, Pettersen R, Keeley NB, Ray JL, Sanna Majaneva S, Stokkan M, Hervik A, Angell IL, Snipen LG, Sundt MØ, Rudi K. **In Review**. Key roles of microbial sulfur and ammonium oxidizers for the coastal seafloor ecological state.

Keeley N*, Dunlop K, Laroche O, Hansen PK, Rudi K. In Prep. **An approach for quantifying the influence of fish farm waste on hard-bottom habitats.**

Mikrobiell eDNA

- Converged on microbial
- Repeatedly proven for organic enrichment
- Sensitive / tracer equivalent
- Uptake for compliance is slow*

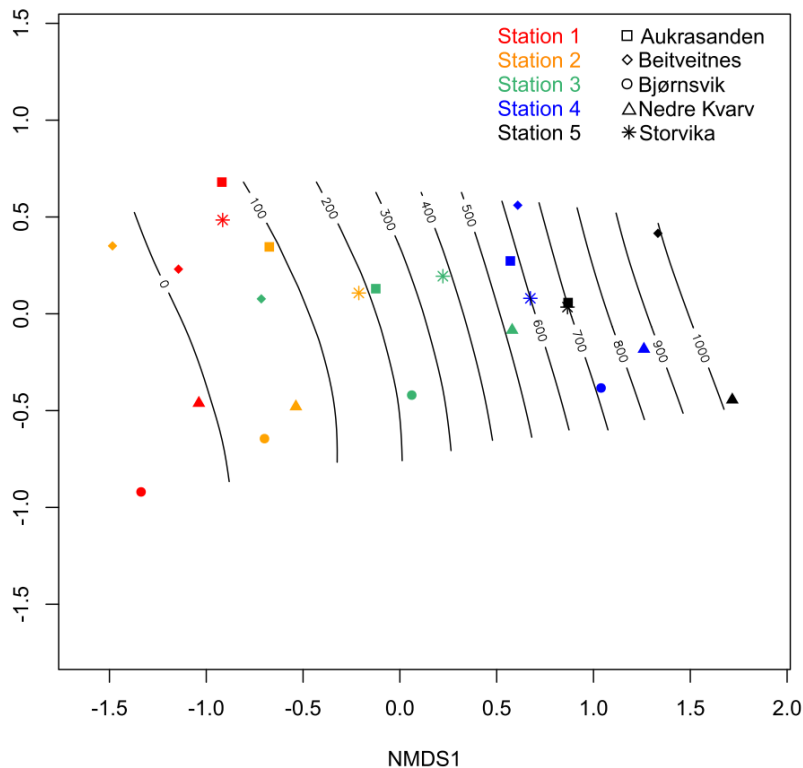
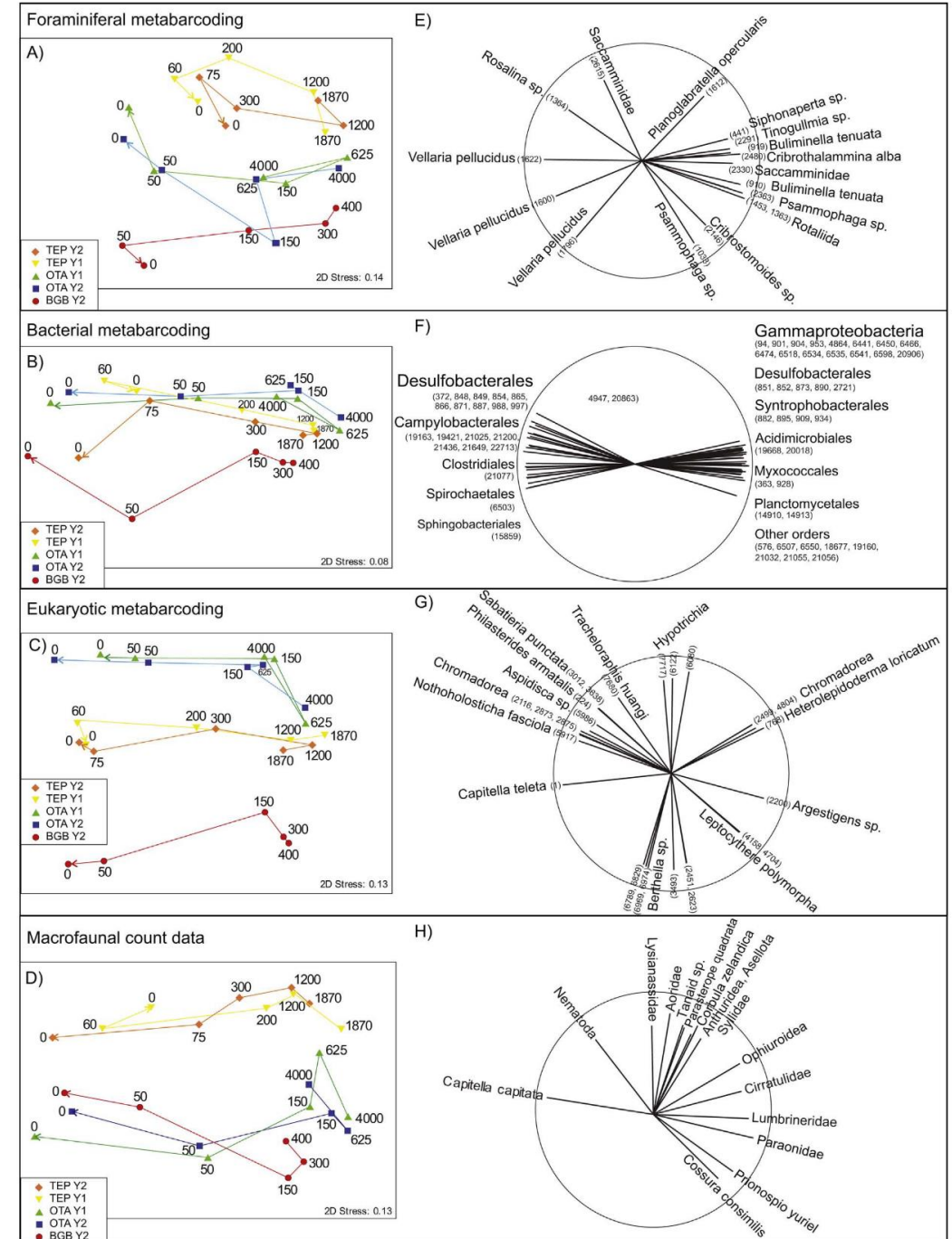


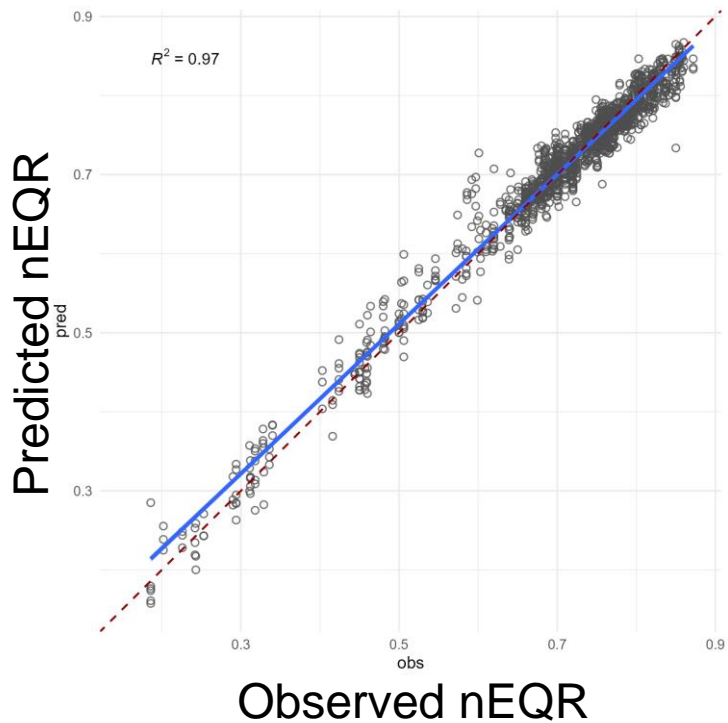
Fig. 5. Plot of non-metric multidimensional scaling (NMDS) analyses of bacterial communities at the different farms and sampling sites with model-fitted distance contour. Correlation results of envfit-analyses for NMDS axes 1 and 2 are shown in Table 2. Each symbol in the plot originates from three pooled replicates of the same sampling sites, accounting for potential spatial variation (patchiness) of bacterial communities at a specific sampling site. An NMDS plot for all individual replicates is provided as Supplemental File.

(Stoeck et al. 2018, MPB127)

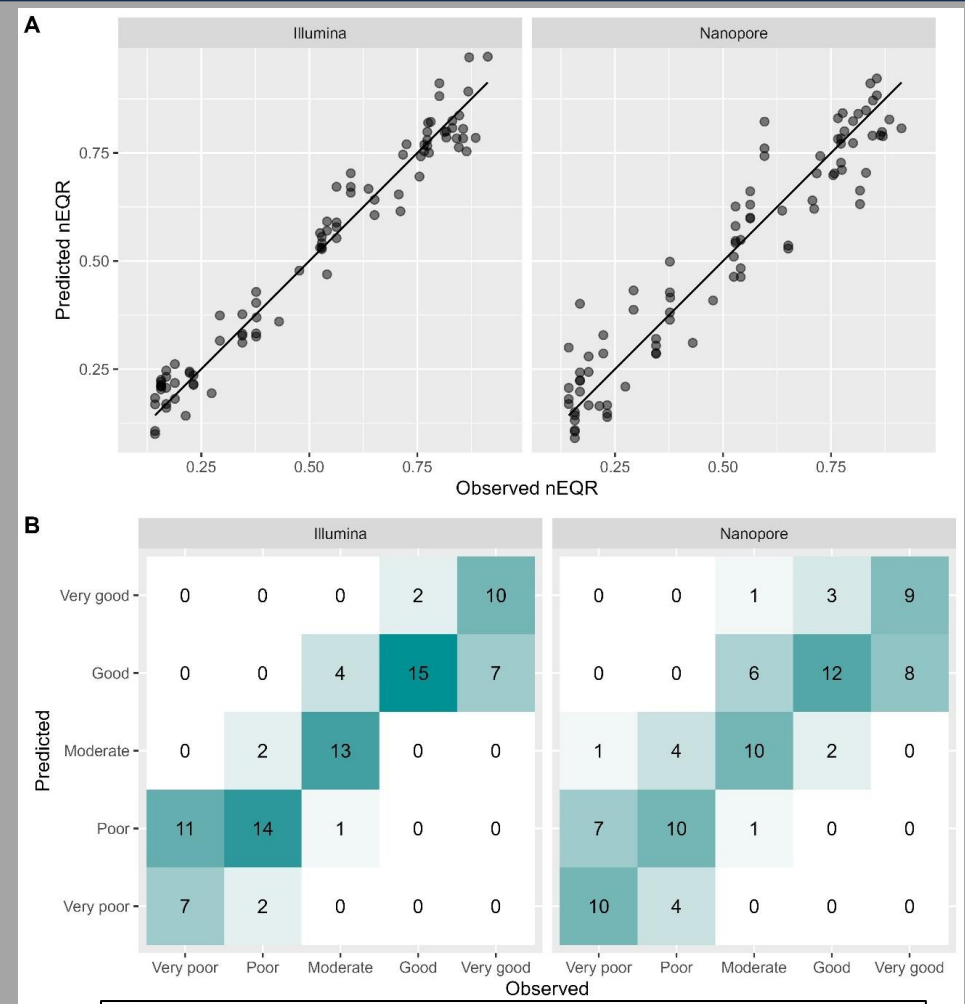
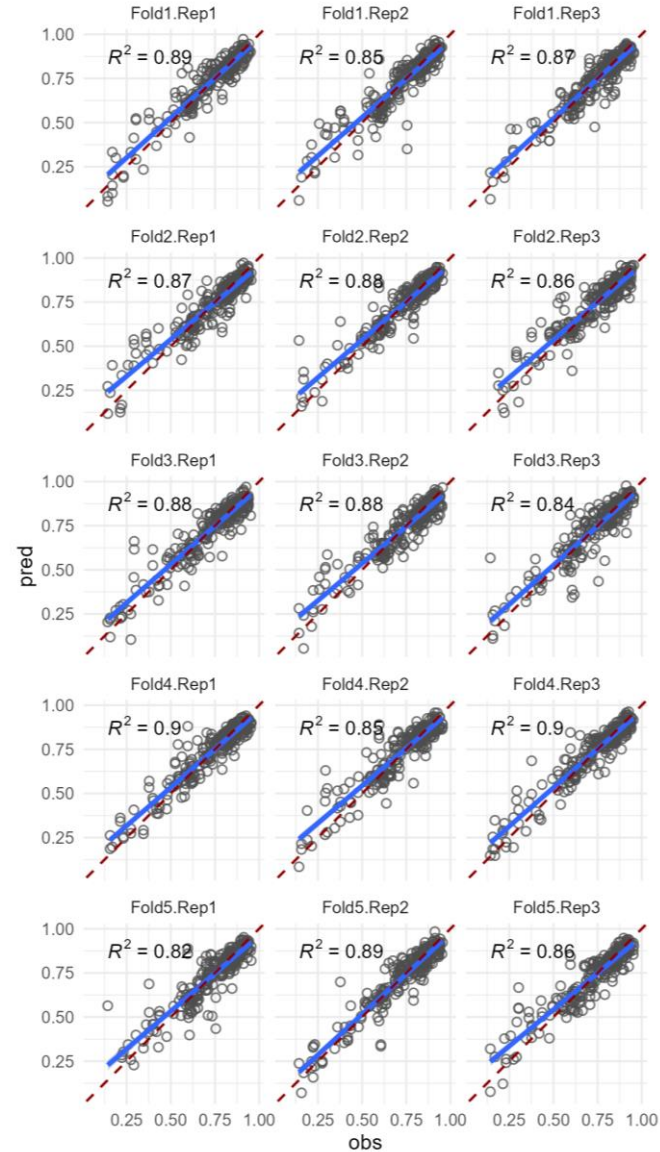


Mikrobiell eDNA

- Norway-specific AQUAed database
- Can be used to assess benthic enrichment *now*
- Question of exact method and how to implement



nEQR



CSH Cold Spring Harbor Laboratory

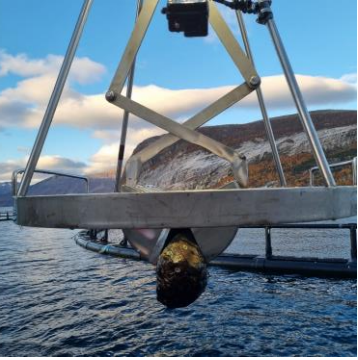
bioRxiv THE PREPRINT SERVER FOR BIOLOGY

New Results [Follow this preprint](#)

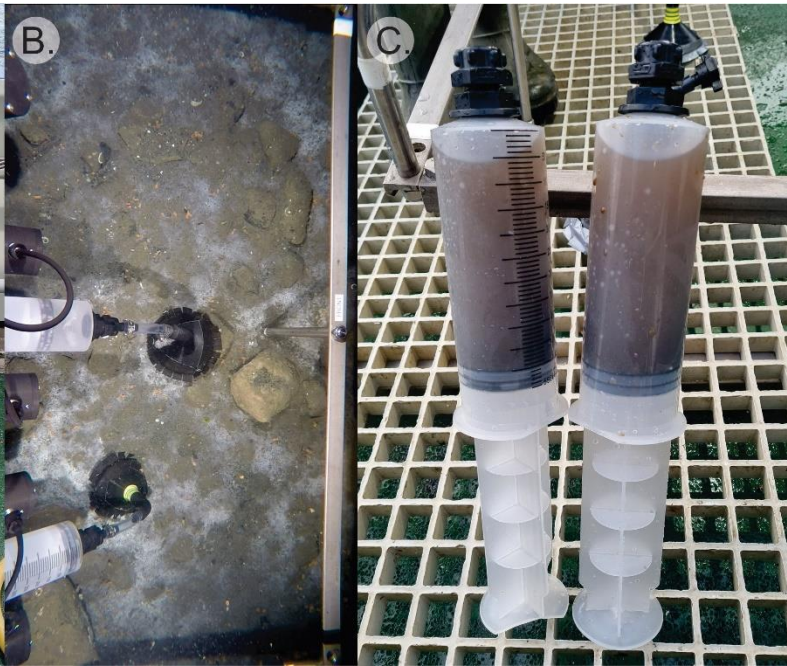
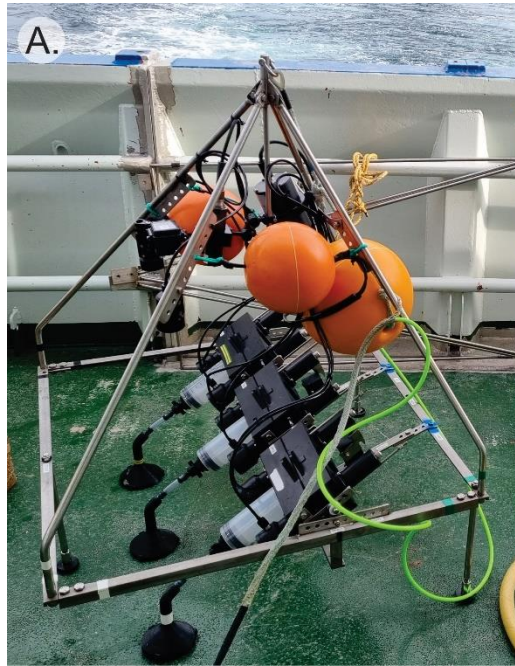
Enhanced Prediction of Seafloor Ecological State Using 16S Nanopore Sequencing

Melcy Philip, Tonje Nilsen, Sanna Majaneva, Ragnhild Pettersen, Morten Stokkan, Jessica Louise Ray, Nigel Keeley, Knut Rudi, Lars-Gustav Snipen

doi: <https://doi.org/10.1101/2024.10.25.620171>

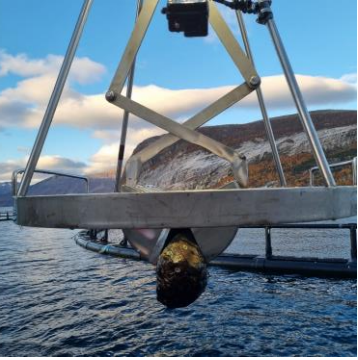


Hardbunnsovervåking



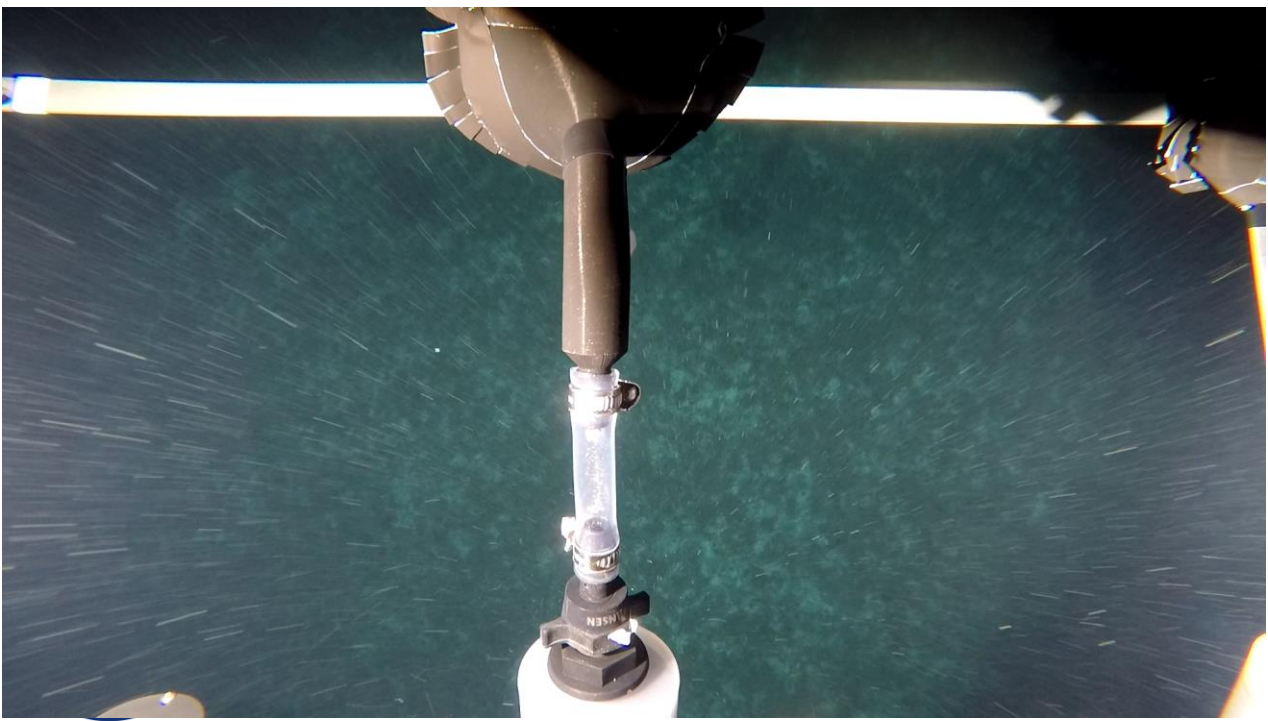
Funded by: Akvakyst (HI), 'Sustain Aqua': NFR No. 267829, 'AQUAed': NFR No. 320076



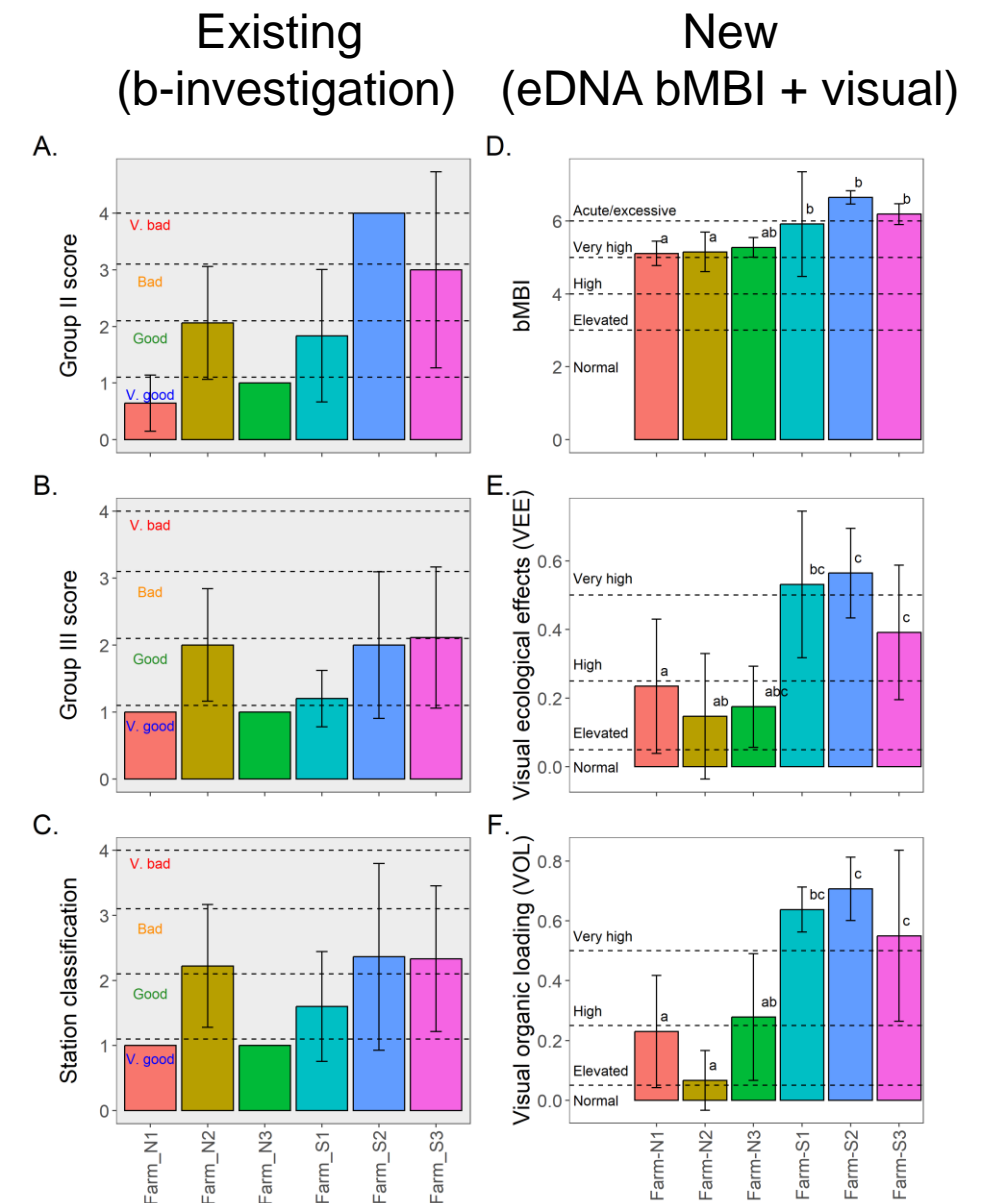


Hardbunnsovervåking

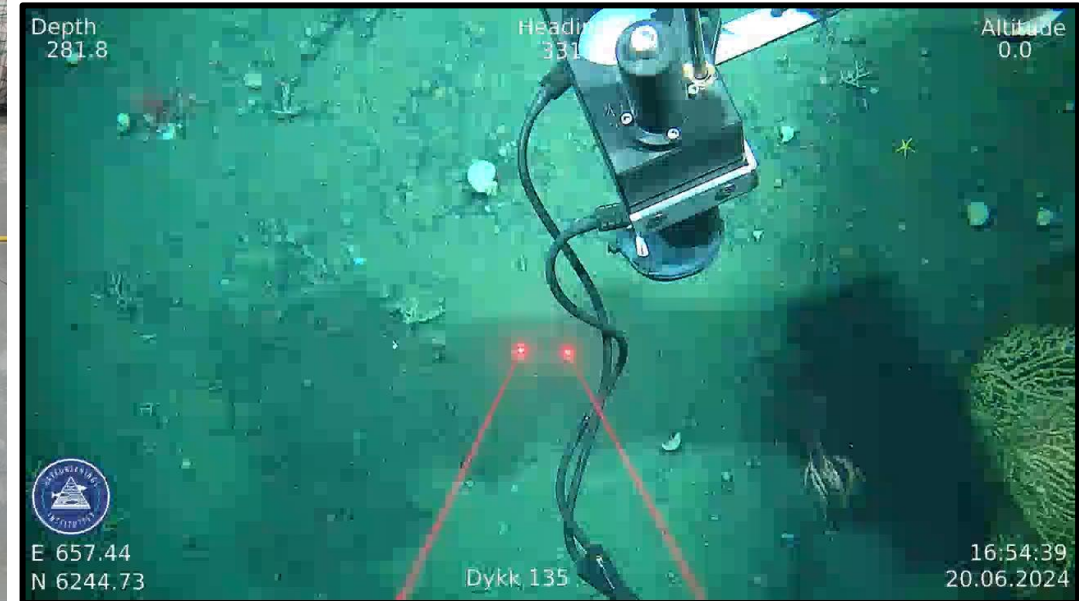
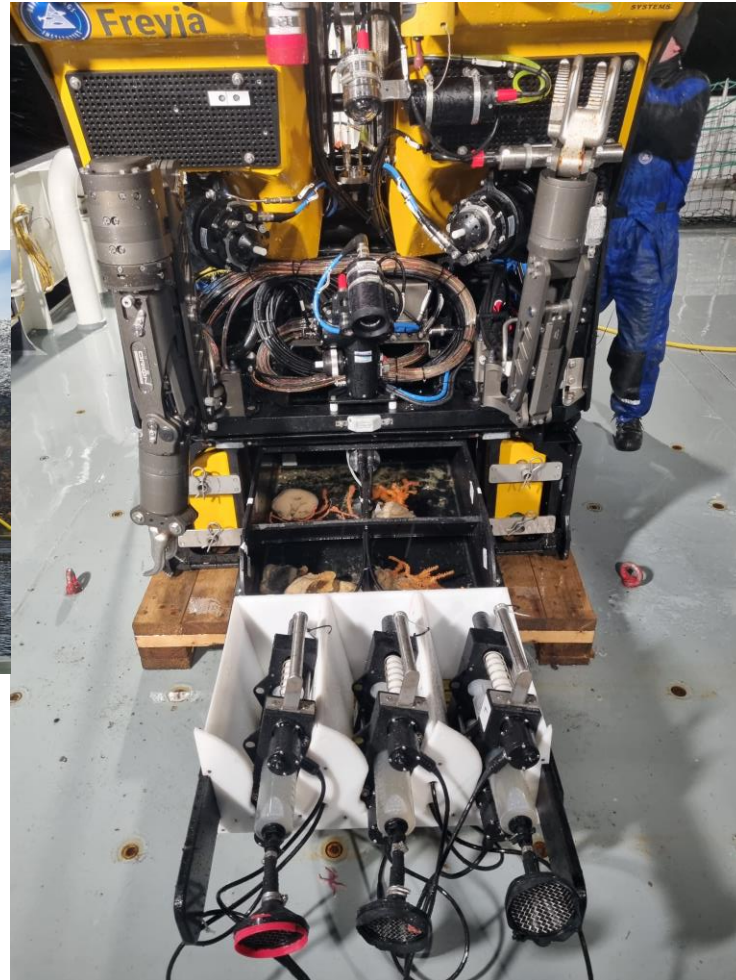
- Combining 16S microbial eDNA + visual indicators
- Cable mounted sample 0-170m depth
- Triplicate samples
- Relatively rapid, quantitative, sig. cheaper than macrofauna



Keeley et al. In Review. An approach for quantifying the influence of fish farm waste on hard-bottom habitats. Ecological Indicators



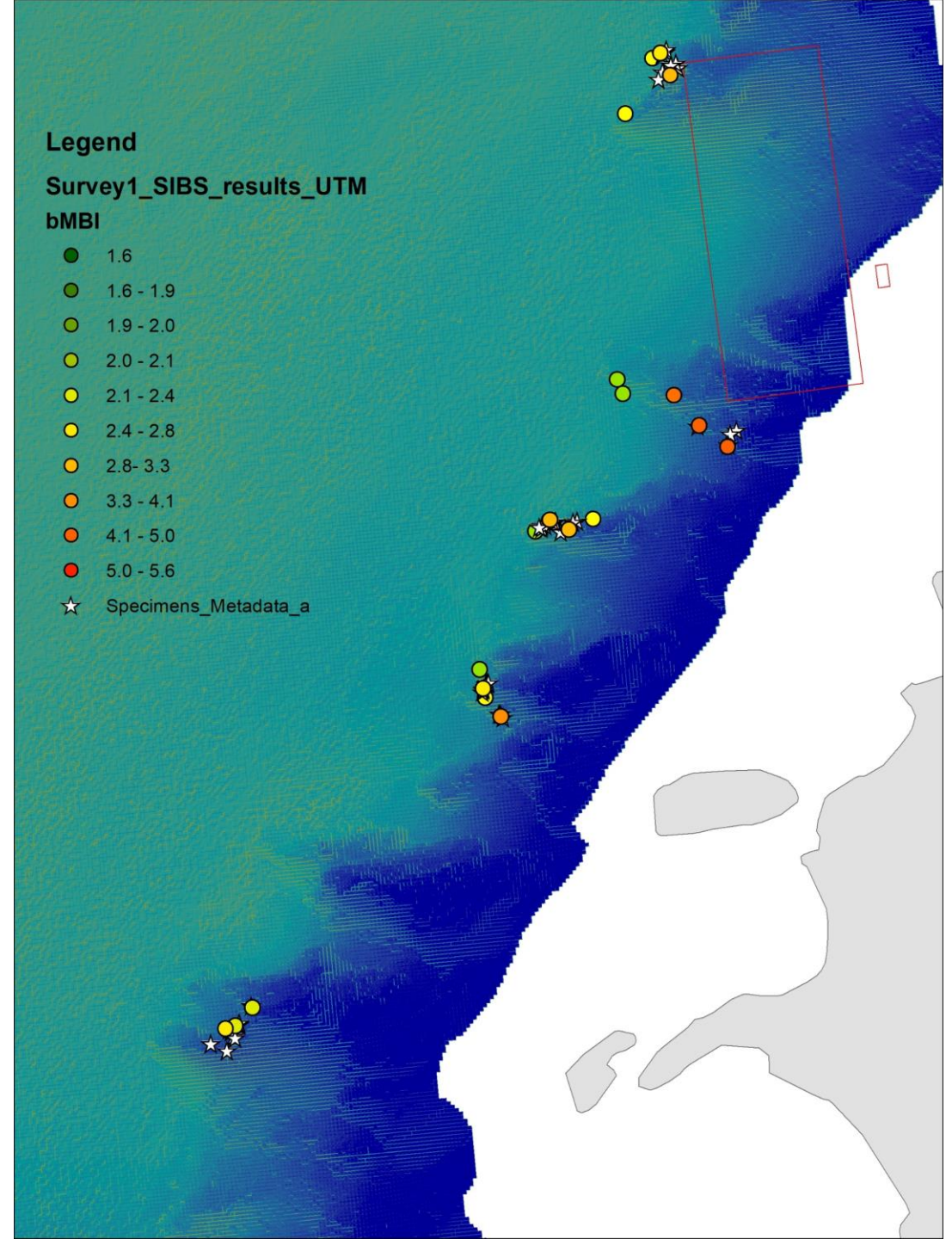
Hardbunnsovervåking - dype lokaliteter



Hardbunnsovervåking - dype lokaliteter

(FHF: VDWS Transition)

- Triplicate samples per dive
- Control over positioning
- Potential for benthic enrichment monitoring at depth
- Scope for properly mapping waste distribution
- Scope for studying waste exposure on sessile reef species



UV-sulfider

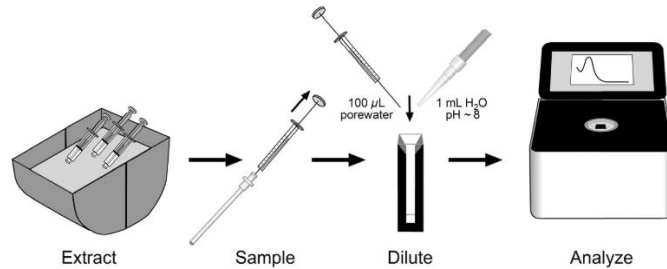


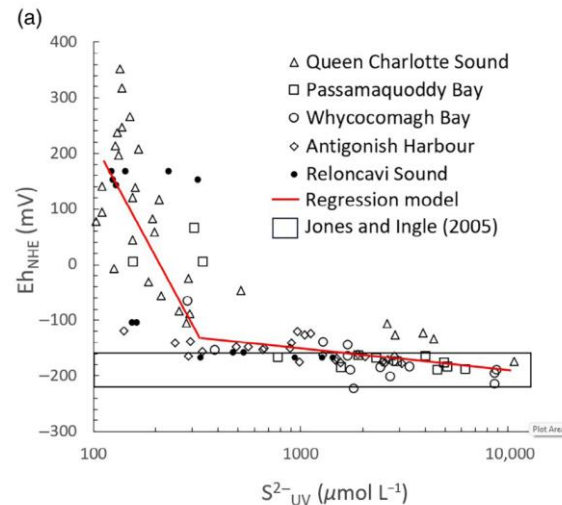
Fig. 1. Illustration of the steps involved in the collection and analysis of sediment porewater for total free sulfide analysis by simplified direct UV spectrophotometry. The depicted materials are described in the text.

Cranford et al (DFO, Canada)

- New UV sulfides method
- Resolved some historical issues with TFS method
- Rapid, practical, needs more testing in Norway
- Arguable 'better' than redox



Cranford



A simple and rapid method for measuring total free sulfides in marine sediments

Peter J. Cranford

Emeritus Marine, St. Andrews, New Brunswick, Canada

Abstract

The quantitatively most important process by which organic matter in marine sediments is mineralized is performed by sulfate-reducing bacteria, resulting in the accumulation of total dissolved (free) sulfide ($S^{2-} = H_2S + HS^- + S^{2-}$) in porewater. S^{2-} is toxic to benthic animals and vascular plants and measurements serve as a proxy for the deleterious effects of organic enrichment on benthic habitat, biodiversity, and ecosystem function. Methodologies for measuring S^{2-} in water have been pursued for at least a century, and standard approaches employ colorimetry (methylene blue and iodometric titration) and potentiometry. These standard methods require between 1 and 200 mL of porewater, which can be laborious to obtain. The ion-selective electrode method is widely employed as a practical approach for sediment S^{2-} analysis but lacks analytical robustness and is highly prone to measurement biases that misinform research and environmental management decisions. A technically simple method is described, based on direct UV spectrophotometry, for the near real-time field analysis of small porewater samples. The procedure prevents known measurement biases associated with particulate sulfide interference, S^{2-} volatilization and oxidation, and represents a practical approach for monitoring organic enrichment and classifying benthic ecological quality status. Porewater concentrations between 200 and 15,000 $\mu\text{mol L}^{-1}$ can be measured and instrument calibration is highly stable. The method has the capacity to rapidly process and analyze sediment samples at low cost, which helps resolve the problem of chronic under-sampling associated with the use of traditional S^{2-} methods.

Anoxic conditions occur at some depth in all marine sediments. In areas where the vertical flux of organic matter exceeds the supply of oxygen required for aerobic carbon mineralization, the anoxic zone can reach the sediment surface and even into the water column. Sulfate is plentiful in seawater and its reduction to sulfide by heterotrophic bacteria is the quantitatively dominant process in the diagenesis of organic matter in marine sediments (Fenchel 1987). Sulfide accumulates in porewater primarily in the form of HS^- but also includes H_2S and S^{2-} depending on pH. These three dissolved sulfide forms are collectively referred to as total free sulfide or S^{2-} .

The ecological implications of sediment organic enrichment from coastal eutrophication and industry effluents (e.g., aquaculture, wood pulp, seafood waste, and offshore drilling) are of interest to scientists and managers worldwide. S^{2-} is toxic to benthic macrofauna (Bagarinao 1992; Grieshaber and Völkel 1998; Wang and Chapman 1999; Gray et al. 2002; Hargrave et al. 2008) and some vascular plants that provide critical coastal habitat (Lamers et al. 2013). S^{2-} measurements have been recommended as a more practical indicator of

sediment toxicity than traditional benthic community taxonomic analysis, which requires greater technical expertise, much higher analysis and interpretation costs, and long delays to obtain results (Wildish et al. 2001). Subsequently, S^{2-} measurements are used in several countries to monitor and classify habitat quality and the health status of benthic communities exposed to organic enrichment (Hargrave et al. 2008; Cranford et al. 2020, 2022).

The measurement of S^{2-} in water and sediments has been the topic of research for many decades. Established analytical methods for S^{2-} analysis of water include methylene blue colorimetry, potentiometry with an ion-selective electrode, and iodometric titration (APHA 2017; Methods 4500- S^{2-} -D, 4500- S^{2-} -G, and 4500- S^{2-} -F). These classical methods require between 5- and 200-mL sample volumes although laboratory adaptations of the methylene blue method can reduce the required sample volume to 1 mL for measuring sulfide concentrations below 30 $\mu\text{mol L}^{-1}$. Extraction of this volume of porewater from sediments is often impractical for research and environmental management applications that require large numbers of samples. The ISE method was adapted for measuring S^{2-} directly in 5-mL sediment/porewater slurries and is relatively simple and inexpensive to perform compared with the

*Correspondence: peter.cranford@outlook.com

Sensitive deepwater species monitoring

Desmophyllum pertusum



Paramuricea placomus



Paragorgia arborea



Primnoa resedaeformis



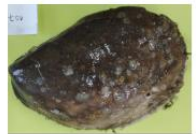
Phakellia ventilabrum



Geodia barretti



Acesta excavata



- SustainAqua (NFR 2017-2020)
- Akvakyst (HI, on-going)
- VDWS Transition (FHF, 2023-2025)

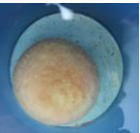
Isidella lofotensis



Duva florida



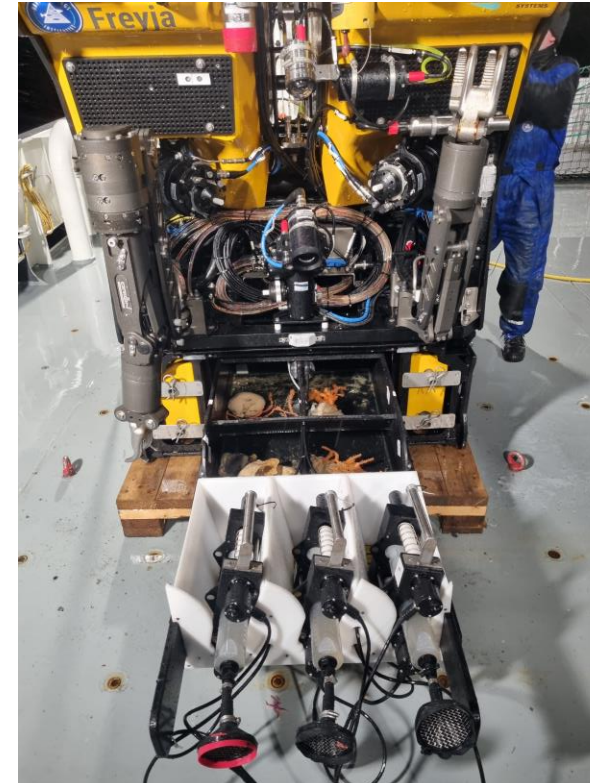
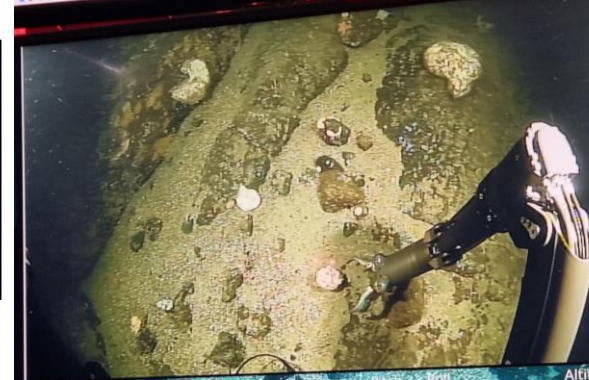
Craniella sp.



Weberella sp.



Hormathia diitata



Analytical processes

(VDWS Transition + Akvakyst)

Culled specimens:

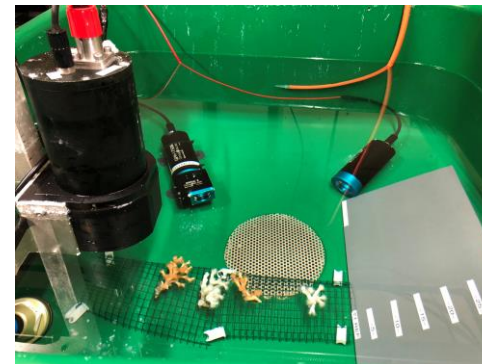
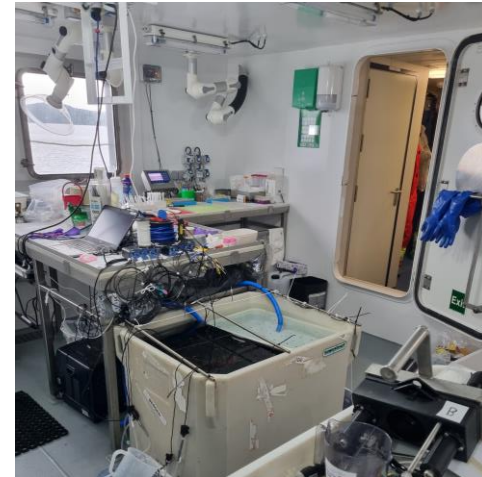
- In-situ visual assessment
- On-board respirometry
- On-board LMS
- Multiple types of tissue samples for in-depth analysis for stress indicators* ('scatter gun' approach)

Live specimens:

- Laboratory exposure trials
- Visual / hyper spectral assessment

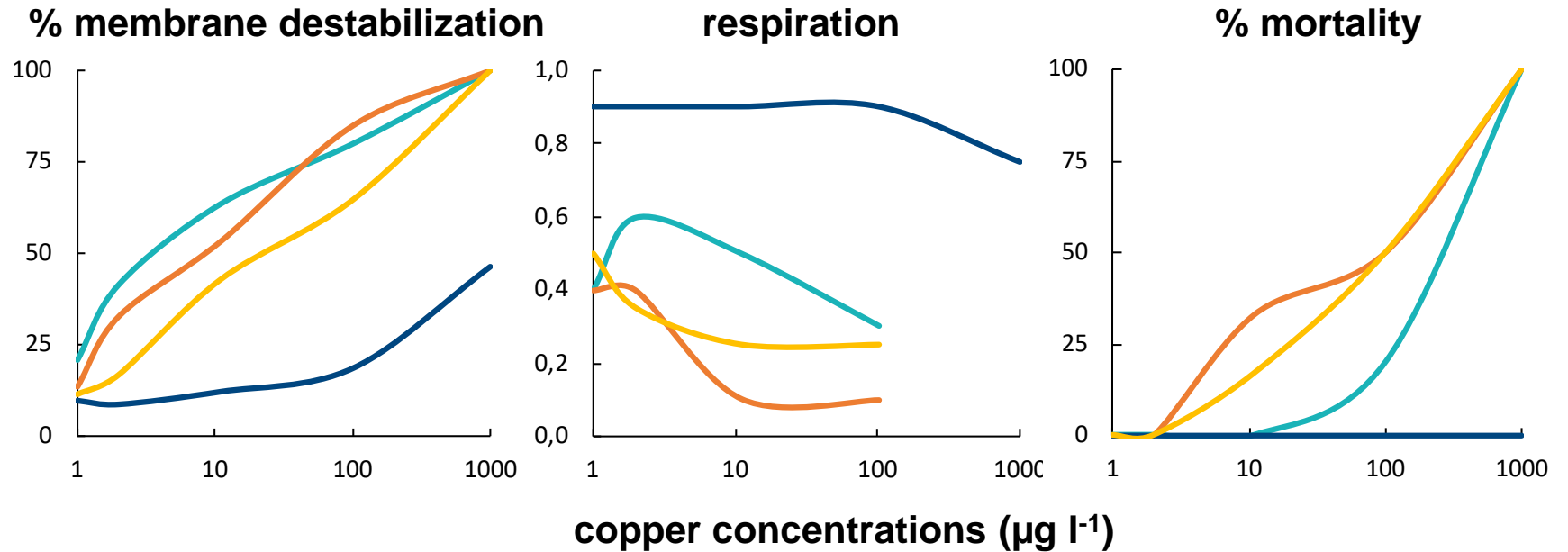
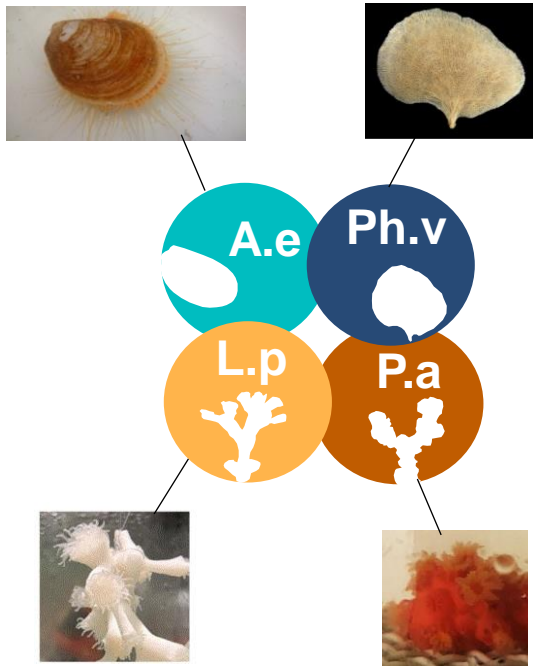
Evaluate 'specimen-specific' waste exposure

- Distance from farm (as the crow flies)
- Sedimentation measurements
- Modelled particle dispersion
- Waste influence on seabed (microbial eDNA)



Sensitive deepwater species monitoring

Effects of antifouling agents on selected deep-sea foundation species - Exposure of 4 deep-sea foundation species to **dissolved copper**. Tina Kutti, Vivian Husa, James Kar Hei Fang, Gry Hunvik, Bjørn Einar Grøsvik, Ketil Hylland



- Large differences in lethal and sub-lethal thresholds for Cu exposure between exposed species
- High sensitivity to Cu detected in the two corals, likely linked to limited mucus production capacity and high metabolic rates
- Rapid mobilization of cellular defence system in the bivalves, possibly causing enhanced tolerance to copper exposure
- No mortality and high tolerance to copper detected in the sponge *Phakellia*



Stressors / contaminants

W.I.P.

Deepwater species

	In-situ exposure	Organic enrichment*	Zn	Cu	Hydrogen-peroxide	Azametifos	Diflubenzuron	Florfenicol	Teflubenzuron	Praziquantel	Emamectin	New-a	New-b
<i>Desmophyllum pertusum</i> 	✓/~			✓							✓		
<i>Paramuricea placomus</i> 	✓/~												
<i>Paragorgia arborea</i> 	✓/~			✓							✓		
<i>Primnoa resedaeformis</i> 	✓/~												
<i>Phakellia ventilabrum</i> 	✓/~			✓							✓		
<i>Geodia barretti</i> 	✓/~			✓							✓		
<i>Acesta excavata</i> 	✓/~			✓							✓		
<i>Duva florida</i> 	✓/~			✓									
<i>Craniella sp.</i> 	✓/~												
<i>Isidella lofotensis</i> 	✓/~												
<i>Isidella lofotensis</i> 	✓/~	✓		✓							✓		

For each:

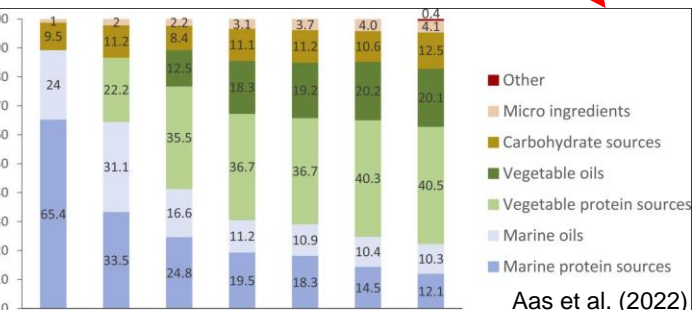
- Relative sensitivity (eg LC50, waste tolerance)
- Species specific stress indicators
- General knowledge of distributions
- General knowledge of functions and ecological value

Clear need to prioritize important stressors based on level of use and anticipated use in future

✓ = Quantitative and useful
 ~ = Qualitative / some relevance
 ? = Potential relevance / not well understood

Potential of new technologies

Ecological compartment	Analytical types						C-investigation				Microbes (eDNA)	Targeted, adaptive contaminant monitoring (>> toxicity testing)	Stress indicators in sensitive species	Specific tracers FA / isotope /genetics					
	B-investigation						Kjemiske parametere (in sediments)												
	Sensory						Macrofauna												
Discharge type	Farge	Luft	Konsistens	Grabvolum	Tykkelse	Redox /pH	S	N	NQI1	NSI	H'	TOM	nTOC	N	C:N	P	Zn	Cu	
Soft sediment footprint																			
Organic enrichment	~	~	~			~	✓/!	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓
Biofouling drop off			~				~	~	~	~	~	~			~				✓
Antifouling chemicals: Cu							~	~	~	~	~							✓	?
All new products							?												?
Infeed therapeutants Zn							~	~	~	~	~						✓		?
All others							?												?
New feed composition (eg terrestrial)							?							~					✓
Bath treatments							?												✓
Subtle far-field effects							~	~	~	~	~								✓
Information from hard substrates?																			✓
Deep water, hard substrata fauna (corals?)																			✓/!
Mobile, demersal, pelagic species?																			✓



NB: Fish feed does not start and end with the salmon!

More sensitive, farm-specific options available



Recommendations for future emphasis

1. Expand toxicity testing for new and emerging contaminants – *don't wait for the natural environment to be the testing ground...*
2. Continue multi-species eco-tox & stress indicator work with 'sensitive' species.
3. Continue to develop new waste tracers based on new feed compositions.
4. Attempt to separate effects from organic enrichment per se, versus that of associated contaminants.
5. Obtain consensus on 16s eDNA monitoring methods. Decide on optimum method and implementation pathway. Implement.
6. Revision of existing monitoring strategy to ensure it is optimally focused and using best available (and cost effective) methods

