

Microbial and viral communities of the deep seafloor sediment and manganese nodules from the CCZ, Pacific

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Background & Summary Main Goals

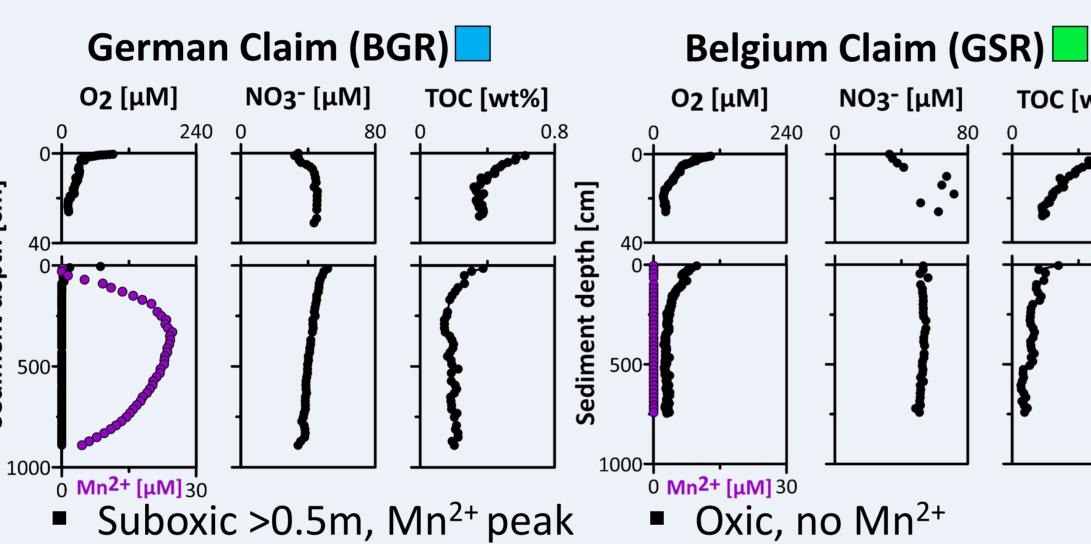
- Manganese (Mn) nodules contain Ni, Co, Cu, Mn, Fe, and rare earth elements.
- The environmental impacts of large-scale deep-sea nodule mining are currently unknown.
- In Feb.-May 2019 (RV SONNE cruise SO268, Mining Impact II) the Belgian & German licence area in the Clarion-Clipperton Zone (CCZ; Eastern Pacific) were studied to obtain baseline characteristics of the > 4000 m deep habitat.
- Research aspects: i) characterization of the distinct present & active microbial and viral communities of bottom water (> 4000 m deep), (< 5 m deep) seafloor sediment, and Mn nodules, ii) diversity and distribution of potential deep-sea cable bacteria and metal-cycling microorganisms, iii) enrichment of Mn-/Fe-cycling bacteria, iv) investigation of deep-sea microbial metabolisms by metagenomic/-transcriptomic, v) quantification of microbial extracellular enzyme activity & cell number, and vi) analysis of (bio-/)geochemistry
- In summary, the potential consequences associated with removal of Mn nodules and resuspension of sediments during mining could help to evaluate the environmental risks.

- To quantify microbial & viral community composition by Illumina 16S rRNA sequencing (RNA & DNA based) from ≤ 100 Mn nodules and eight 5 m long gravity cores from the CCZ.
- To follow the distribution, quantify the abundance, detect the diversity and activity of relevant metal-cycling microorganisms, and as well of potential deep seafloor cable bacteria.
- To enrich potential Mn- and Fe-cycling microorganisms from "live" sediment & "live" Mn nodules under (an)oxic conditions.
- To evaluate microbial activity by extracellular enzyme activity.
- To investigate microbial metabolism of potential Mn- and Fecycling microbes by metagenomics /metatranscriptomics.
- To compare metal-cycler of different locations (a) Mn nodule areas (CCZ/DISCOL), (b) massive sulfide deposit (Indian Ocean).

Deployments during SO268

Field Sites – Clarion-Clipperton Fracture Zone (Eastern Pacific Ocean)

Sampling locations of cruises SO239 & SO268 Area of Particular



- POC flux: 2 mg m⁻² d⁻¹
- NO₃ reduction
- Oxic, no Mn²⁺
- POC flux: 1.5 mg m⁻² d⁻¹

TOC [wt%]

■ Constant NO₃

Manganese (Mn) Nodules

and different depth (~0.5 cm)

Differences in volume (17 - 640 cm³),

weight (30 - 960 g), shape (round or

Subsamples from on top of the nodule

surface (a), bottom side (b), inside of

the nodule core (c), 3 samples per

nodule for DNA & RNA extraction;

enzyme activity tests; cell counts

elliptical), porosity, attached fauna

■ ≤ 100 Mn nodules from BGR & GSR

Gravity Core Sediment

- 5 m long gravity cores (GC) from eight different locations of the CCZ
- Differences in O₂ penetration depth (around 2-3.5 m; oxic and suboxic layers)
- 8 to 27 samples per GC core, taken at 0-5 m depth; stored at -20°C and -80°C
- Preliminary enzyme activity results of an oxic GC (GSR) from a carst region: Aminopeptidase activity until 40 ± 5 cm depth→ detectable active C_{org} degradation

Manganese Concentration

Preliminary Results – Microbial Activity of Manganese Nodules

e.g. β-Glucosidase and Chitobiase ctivity (µM/h) **Bottom** enzyme 0.5 -

BGR Ref. BGR Trial GSR Ref. GSR Trial

- **Activity:** surface layer > bottom layer
- Ref.: > 1.6 μ M/h; nodule core: 0.3 μ M/h
- **Activity:** deep-sea sediment > Mn nodule
- **Bottom water** Mn nodule surface Sponge Ophiuroid Cellulose (polymer of β -(1-4)linked glucosyl (polymer of β-(1-4)residues) linked N-acetyl-βglucosamine) Mn nodule bottom Sediment
- > High unexpected activity in & on Mn nodules!
- β-Glucosidase: degradation of oligosaccharides
- Chitobiase: degradation of chitin
- Micro-habitat of Mn nodules may be a potential major location of microbial C_{org} degradation
- Highest activity on nodule surface of BGR \triangleright Implication: what happens after mining operations when Mn nodules are removed?
 - > Open question: who are the active players which are responsible for C_{org} degradation in &on nodules?

Working on board during SO268