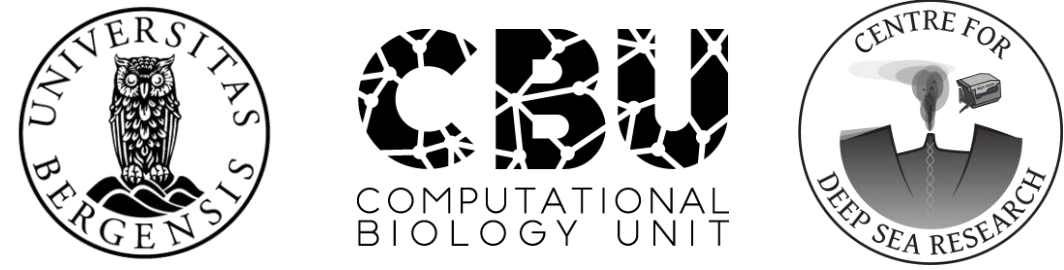


Interpretation of microbial food webs from metagenomic data in deep-sea hydrothermal vents across the Arctic Mid-Ocean Ridge



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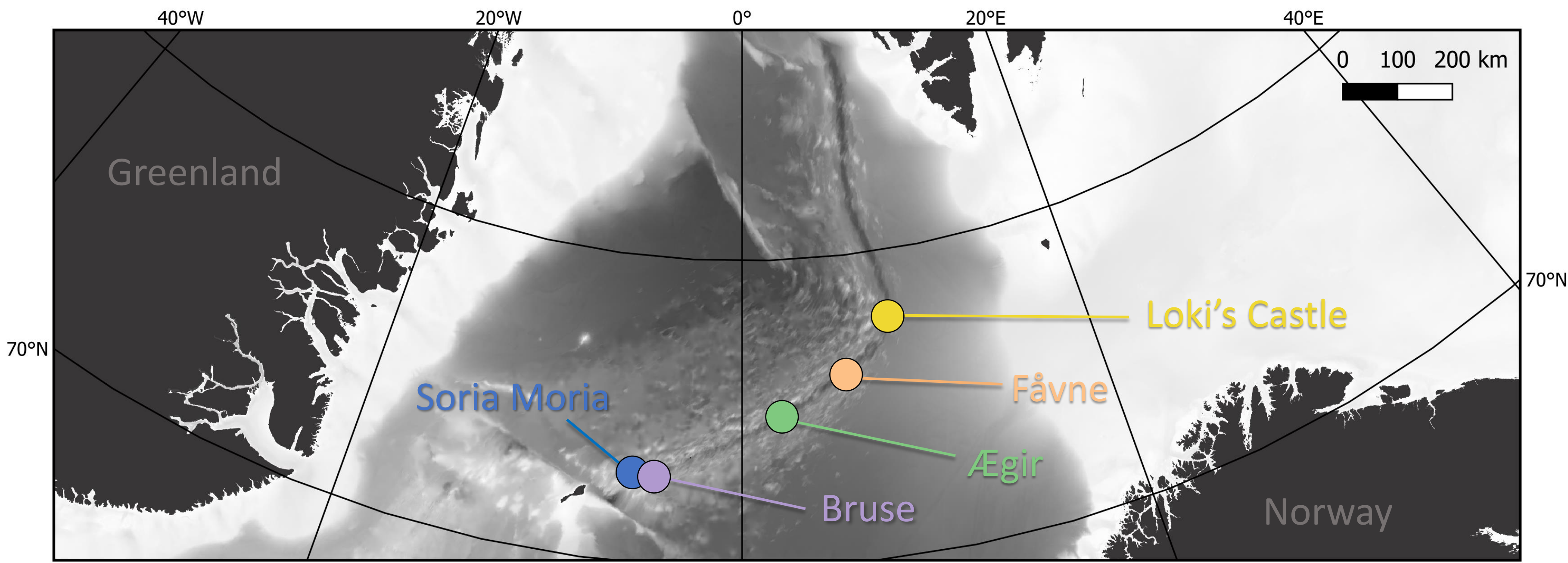
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Study Motivation

Energy availability from different redox reactions varies considerably between and within the studied hydrothermal vent fields due to variations in geological and chemical setting. Do the same primary producers occupy available niches or do different organisms fill these roles within Arctic Mid-Ocean Ridge hydrothermal food webs?

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Hydrothermal Vent Sites



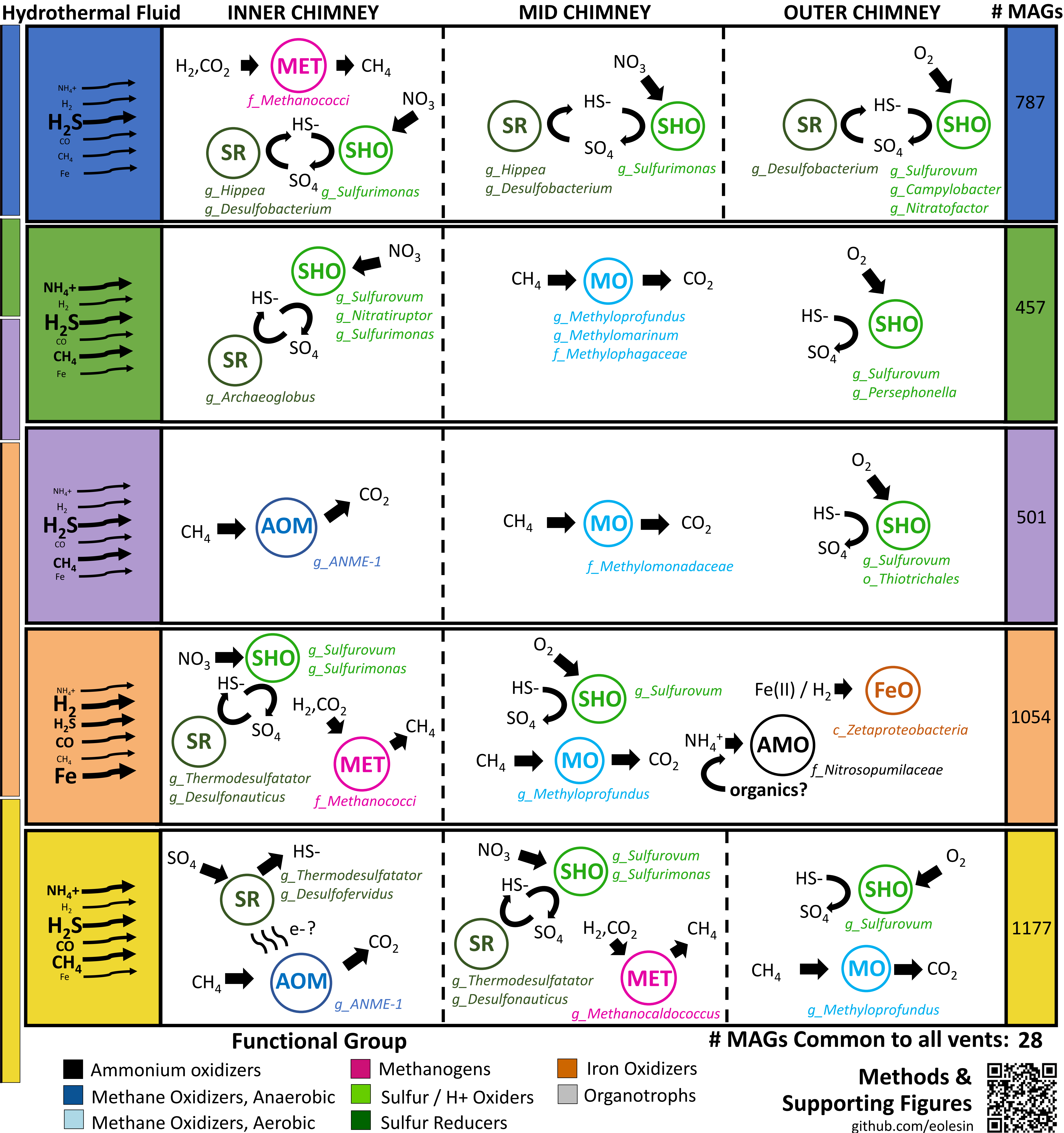
Key Findings

- Food webs are different between sites
- Food webs are shaped by energy landscapes
- Few genomes are ubiquitous

Next Steps

- Comparative genomics between highly-related metagenome-assembled genomes (MAGs)
- Investigate CRISPR spacers to describe viral predator diversity

Results



Methods & Supporting Figures
github.com/eolesin

