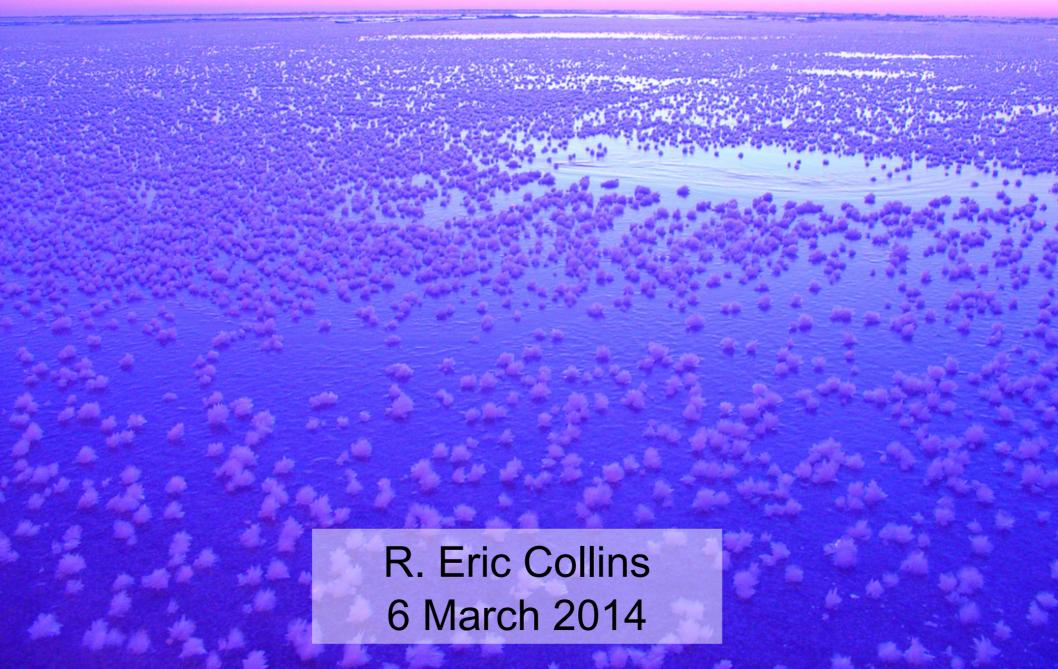
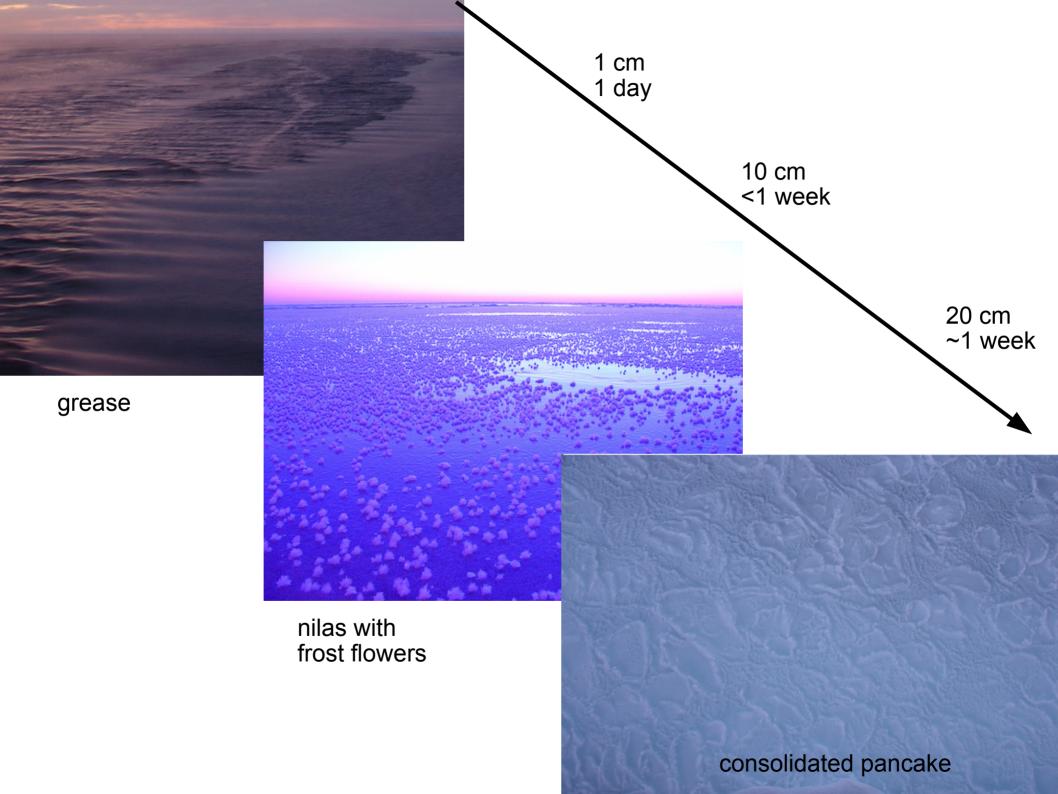
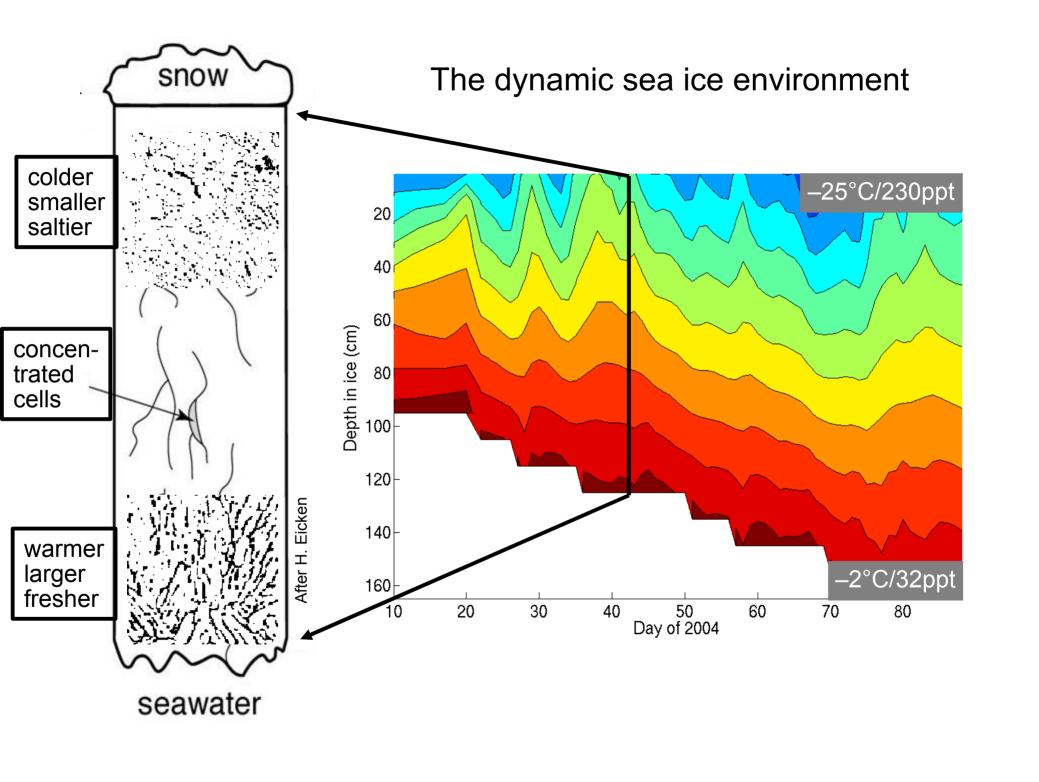
Oil Biodegradation Potential of Sea Ice Microbial Communities

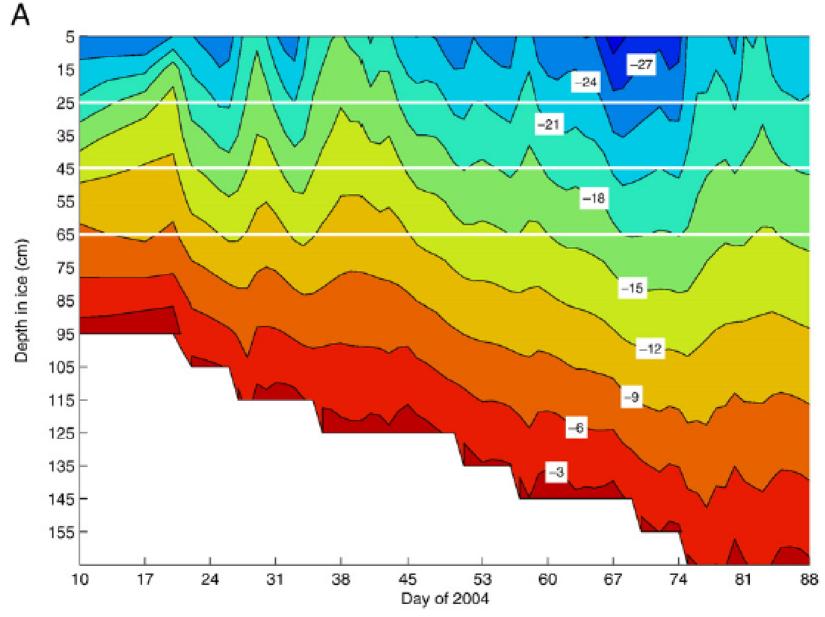




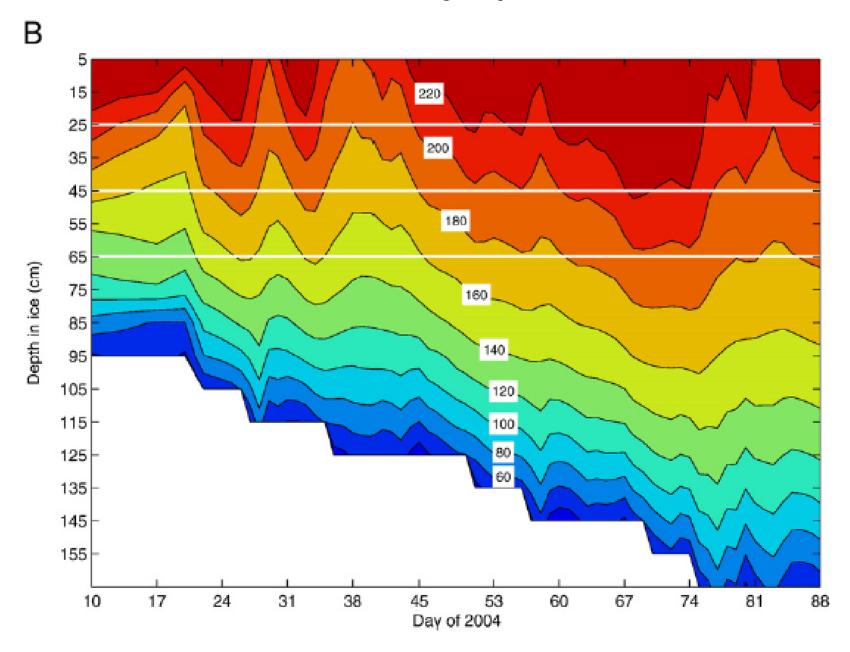




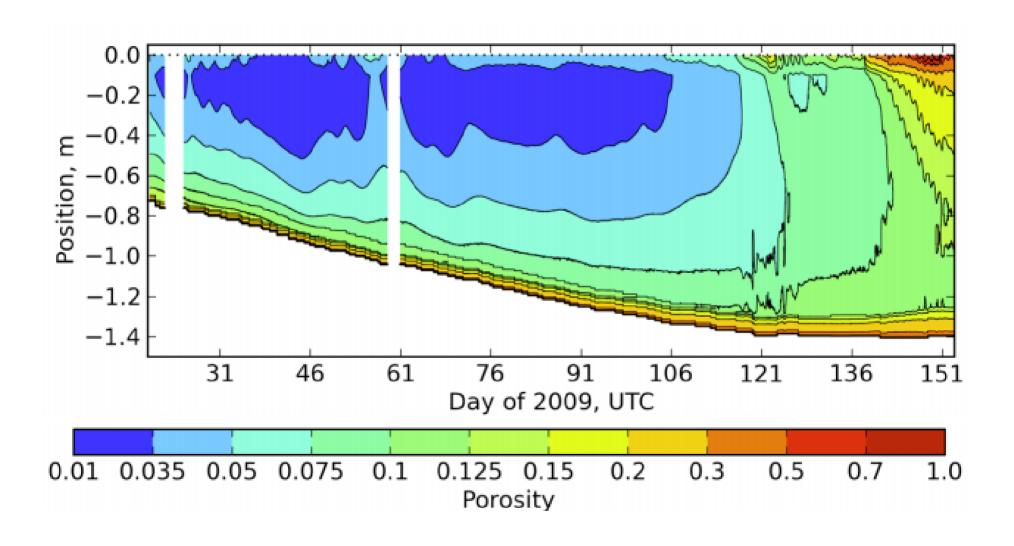
Ice temperature (measured, deg C)



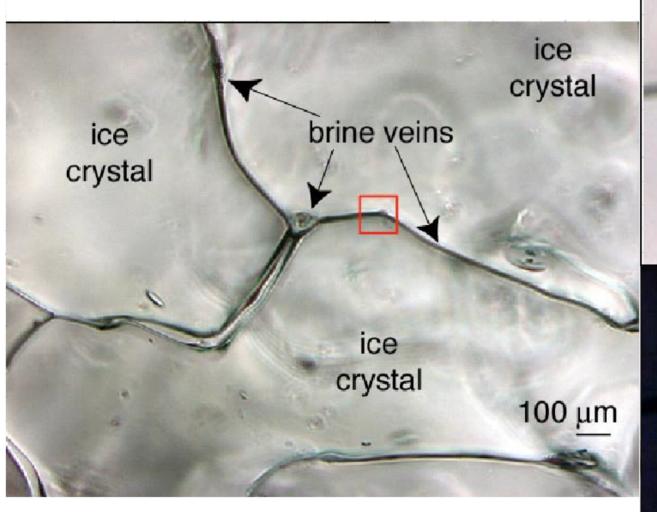
In situ brine salinity (calculated, ‰)

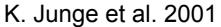


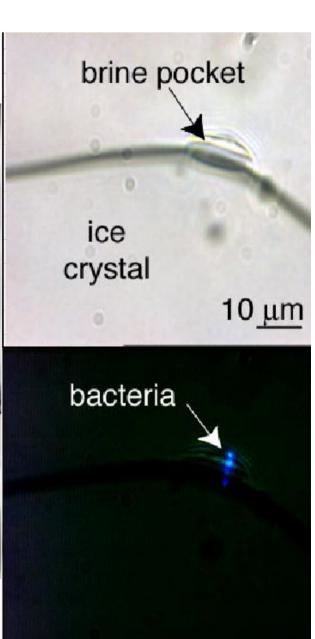
Brine volume fraction/porosity



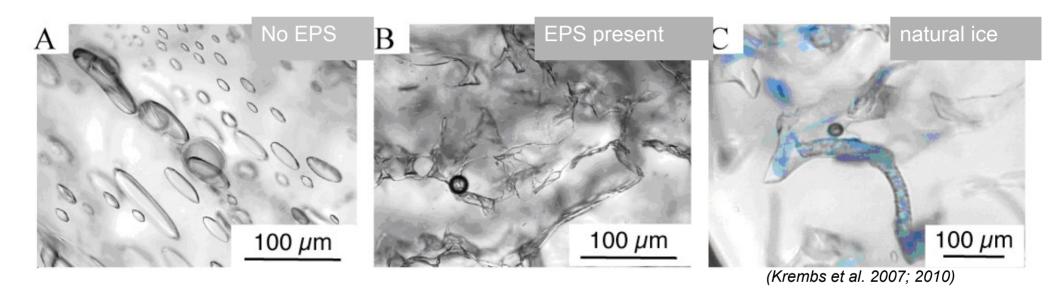
Microbial habitats





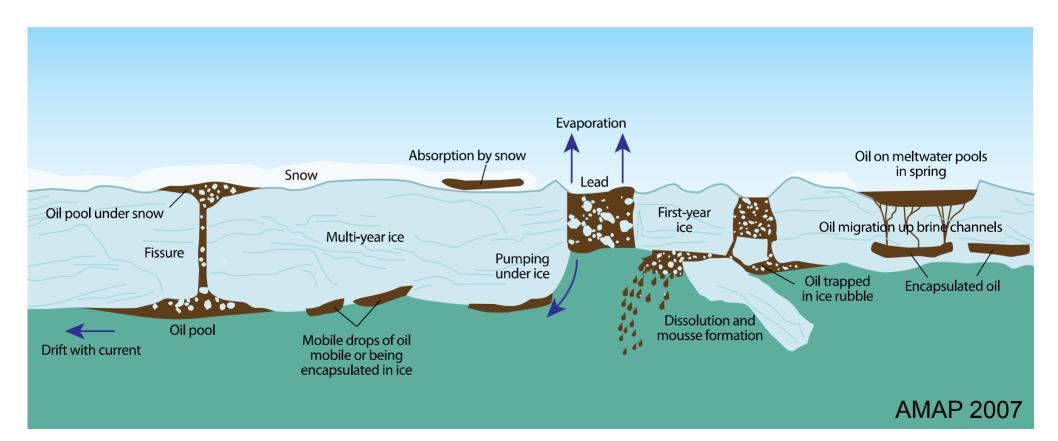


Microbial habitats

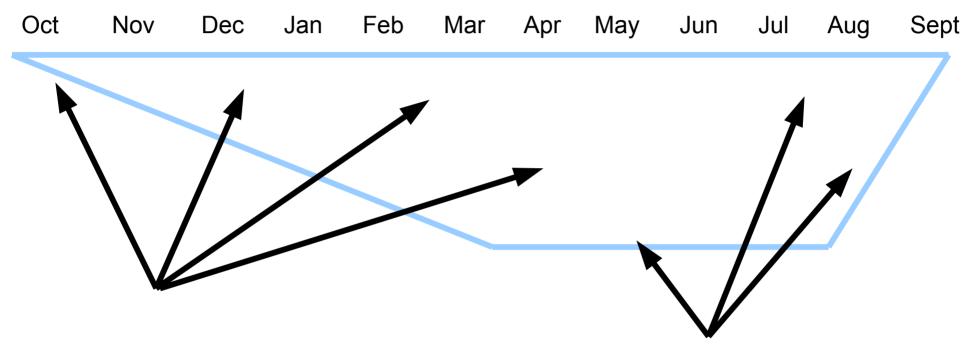


Oil in Ice Scenarios

- Effects of oil on microbial communities?
- Effects of microbial communities on oil (biodegradation)?

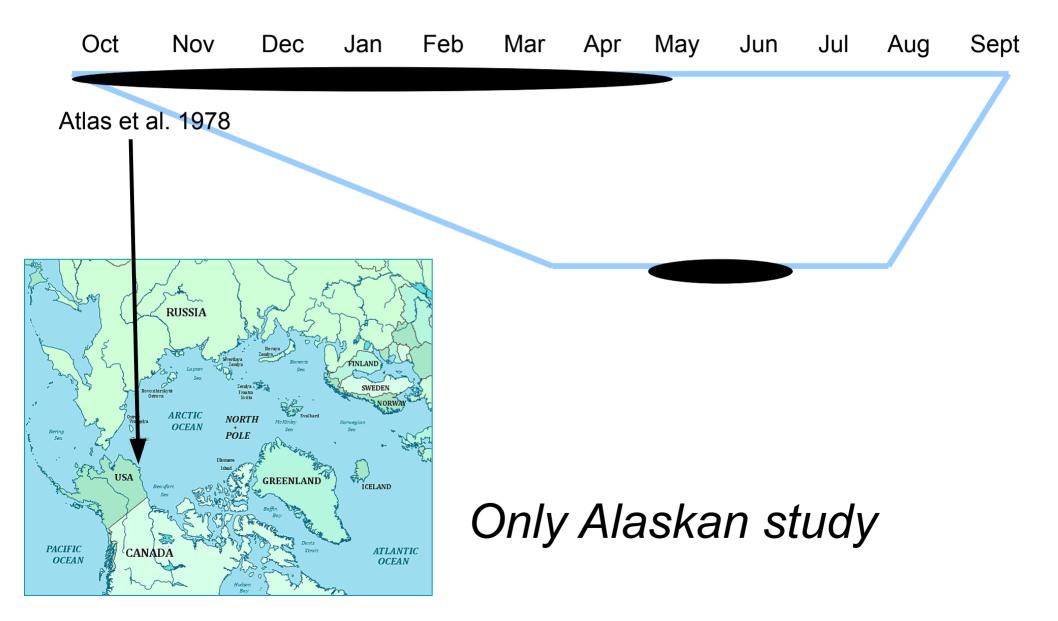


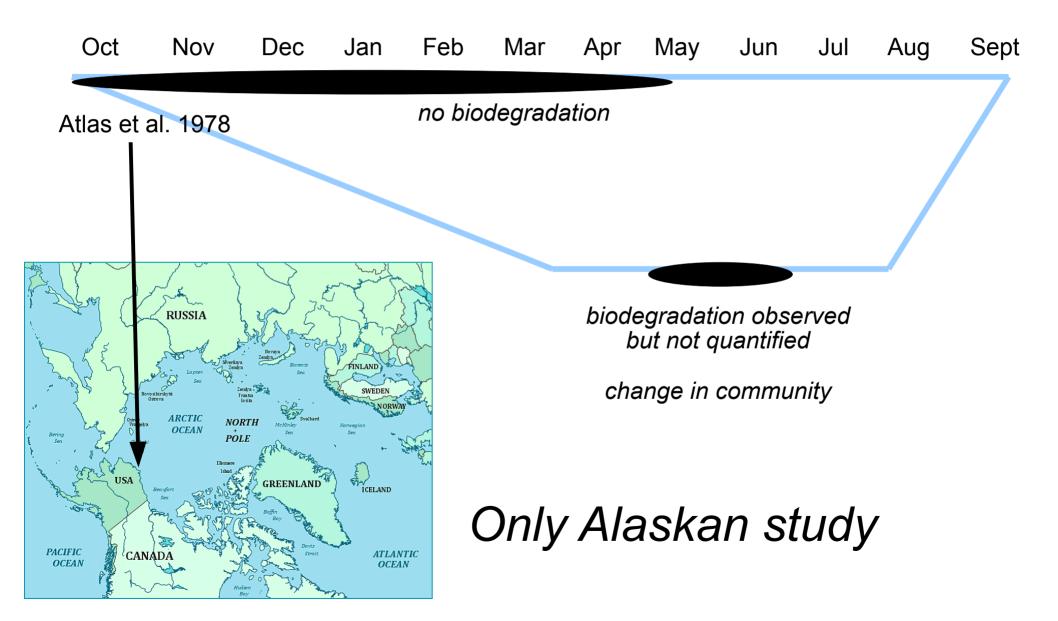
Sea ice microbial community evolution

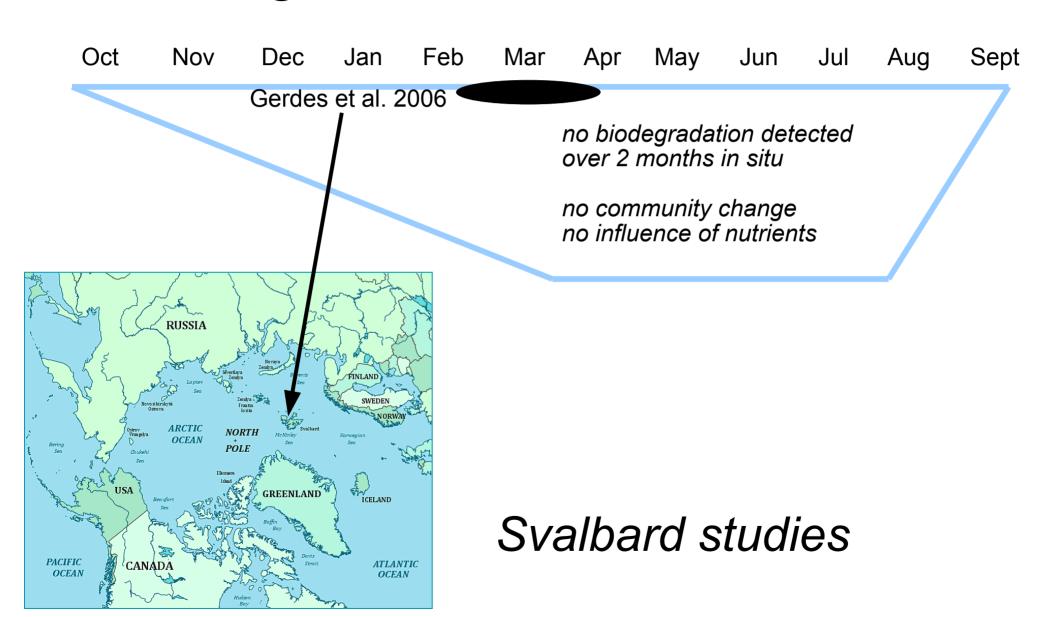


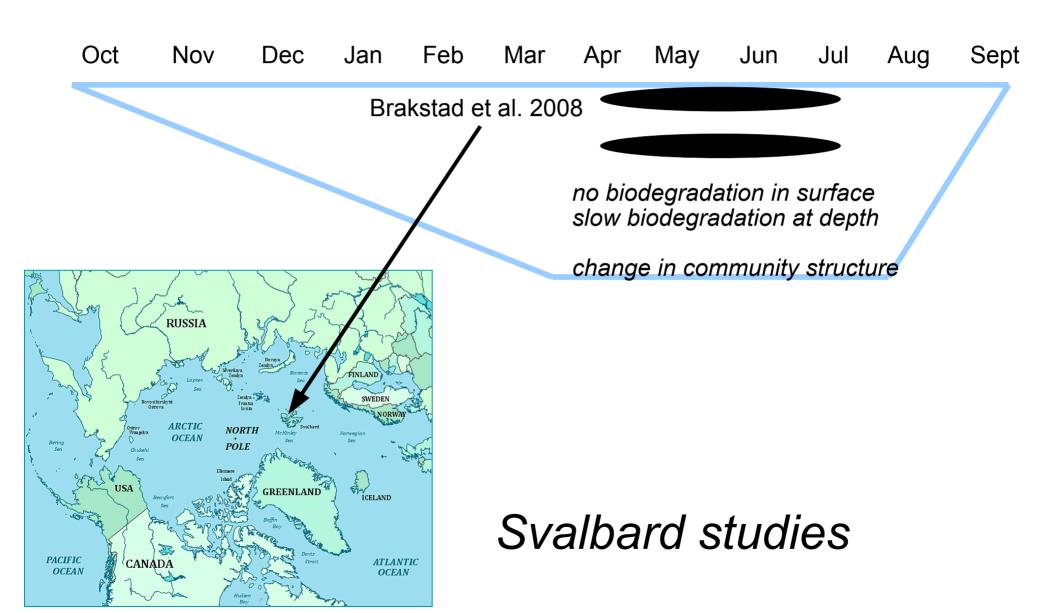
incorporated from seawater
prefer low nutrients
grow slowly
no algal production

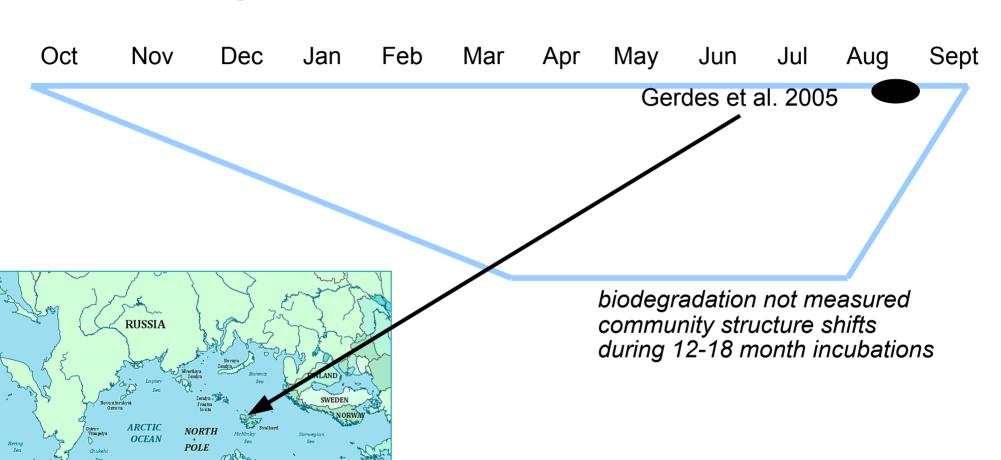
selective growth of 'weeds'
prefer high nutrients
grow quickly
intense algal production











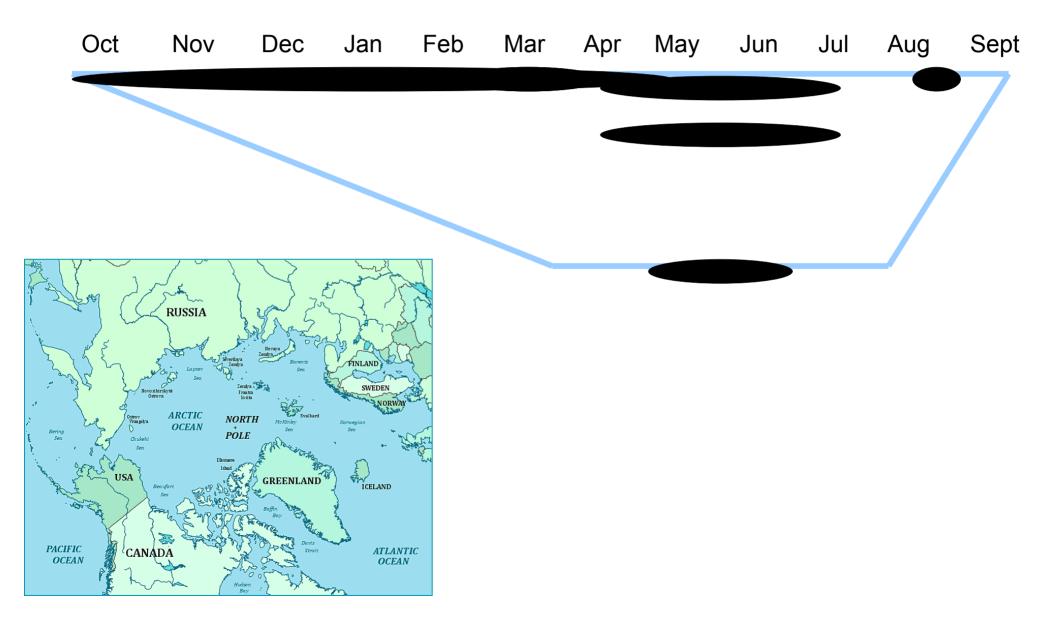
GREENLAND

PACIFIC

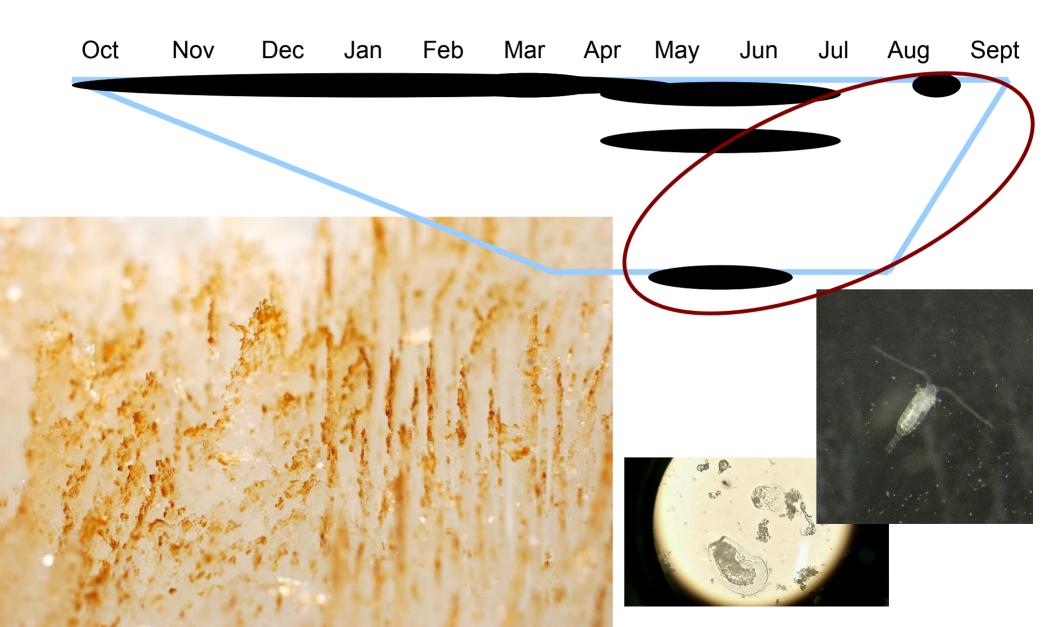
OCEAN

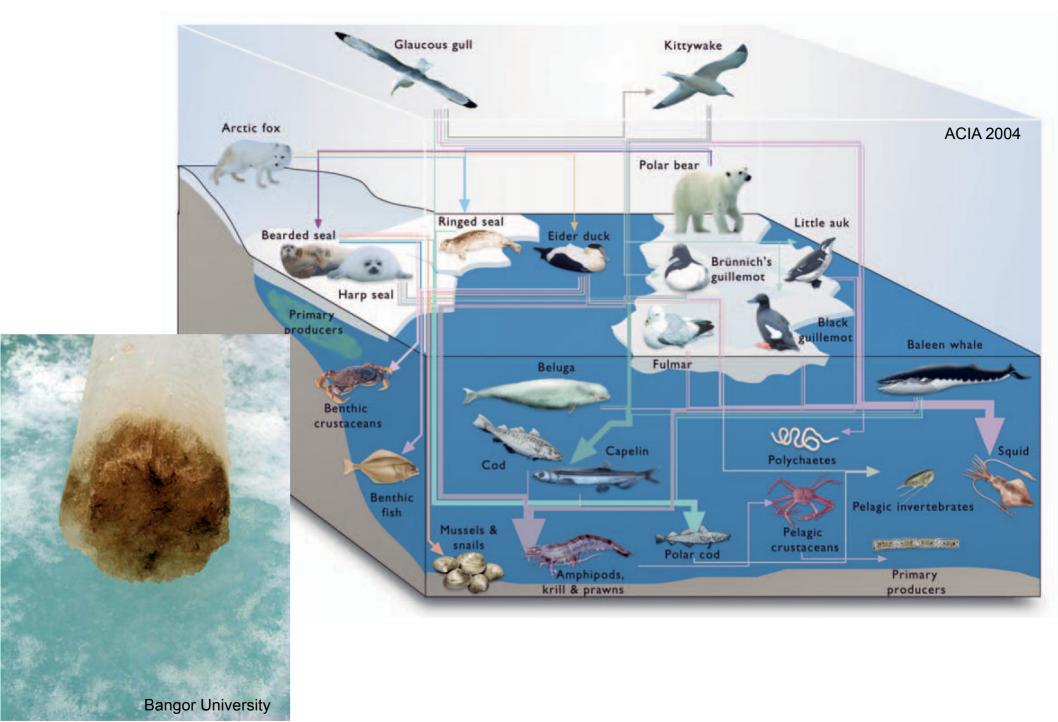
CANADA

Svalbard studies

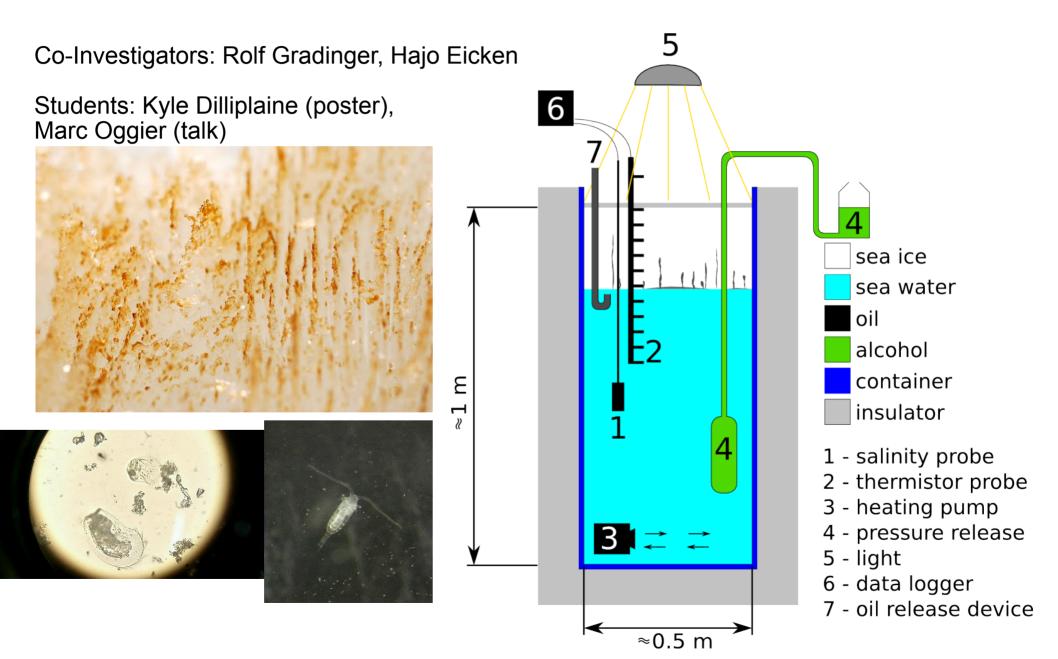


Gap 1: Lack of studies on critical ice biota habitat (bottom of spring ice + summer ice)

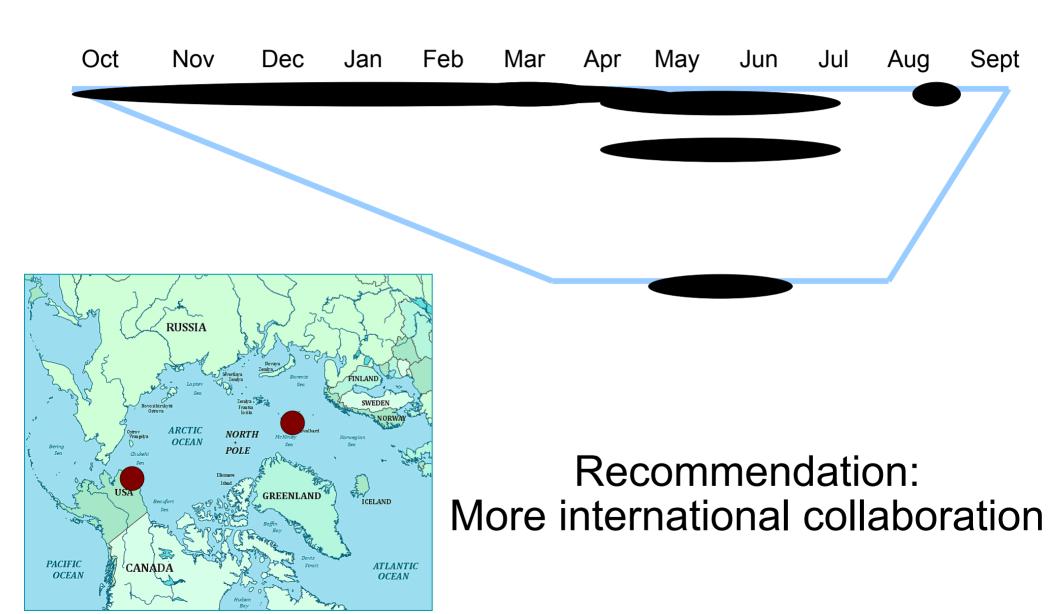




CMI: Crude oil infiltration and movement in first-year sea ice -- Impacts on ice-associated biota and physical constraints



Gap 2: Minimal spatial coverage



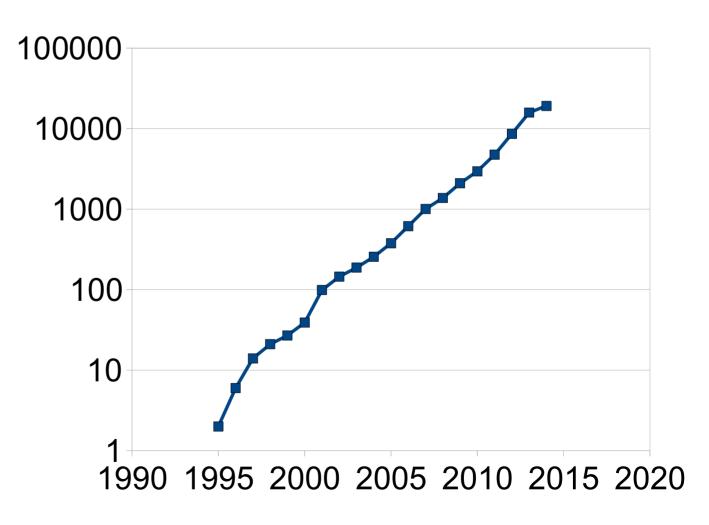
Gap 3: Absence of modern molecular methods (genomics and metagenomics)

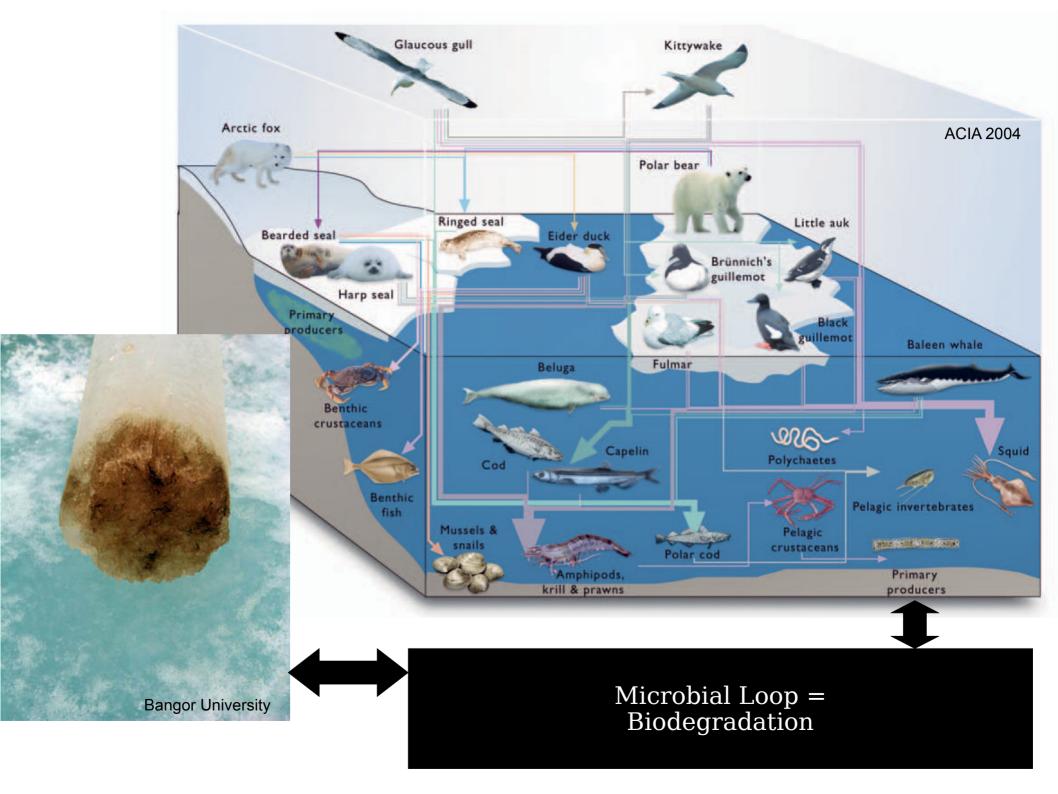
Bacterial Genomes

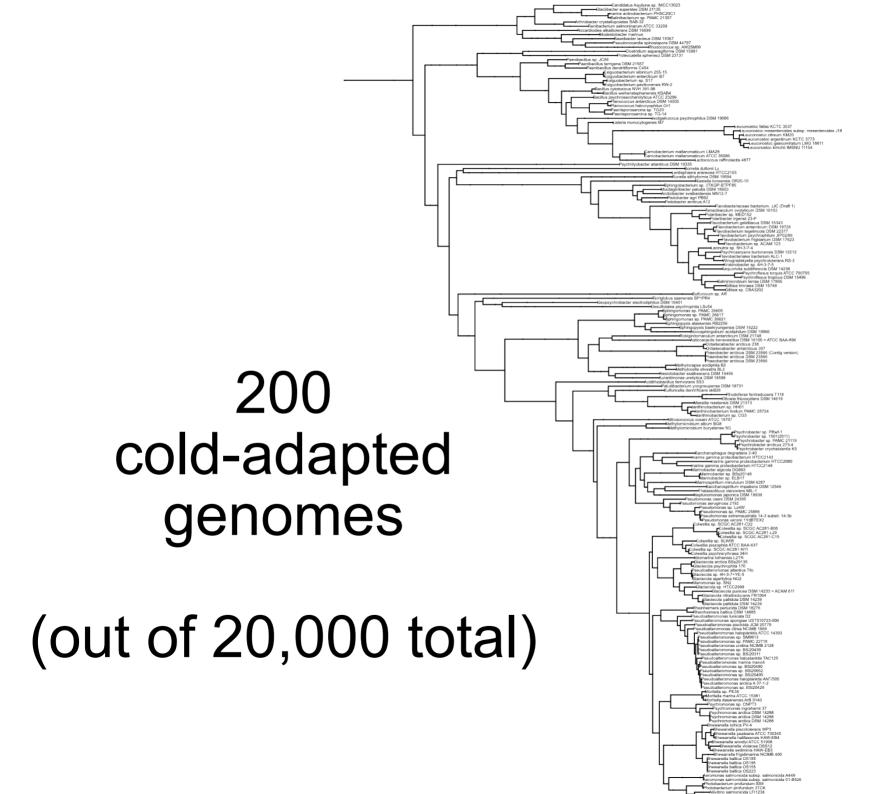
Culturing: Atlas et al. 1978

Fingerprinting: Gerdes et al. 2005 Gerdes et al. 2006 Brakstad et al. 2008

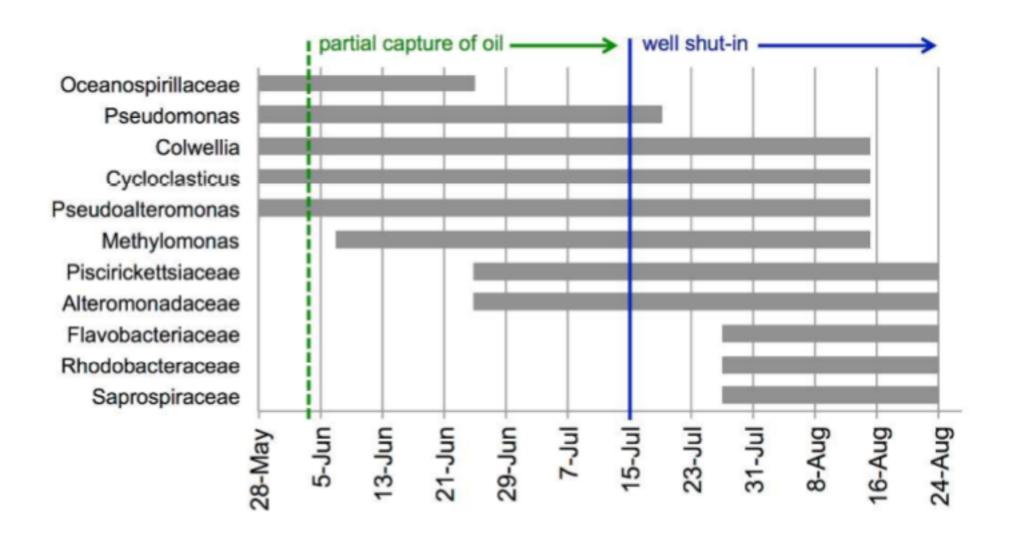
Sequencing: none to date

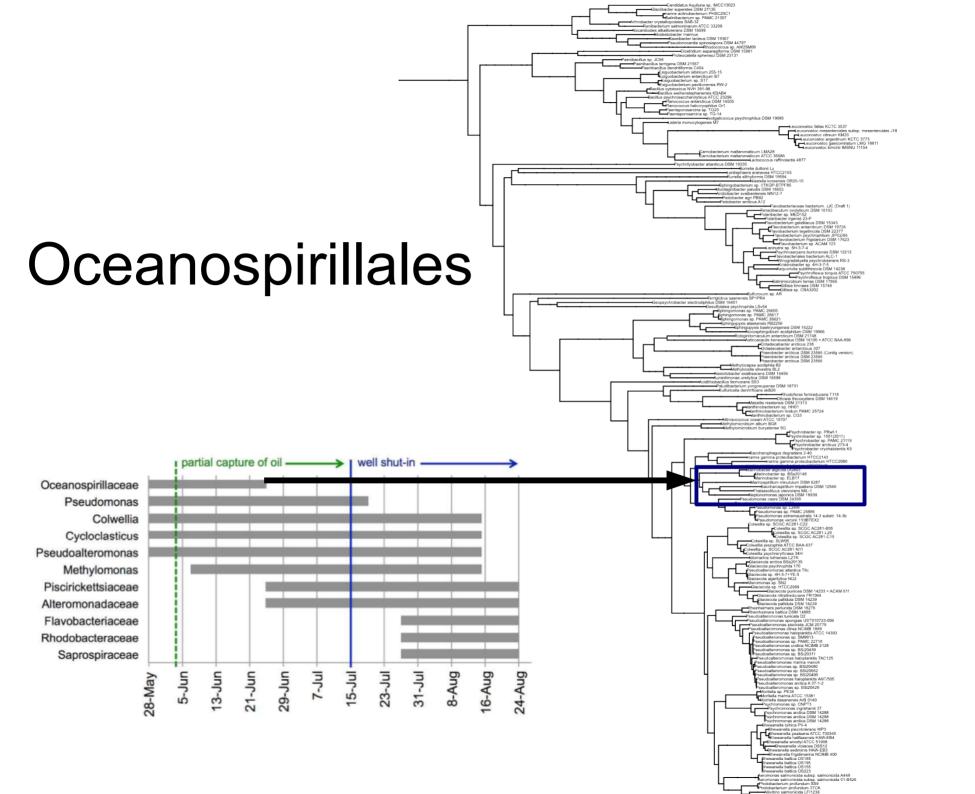


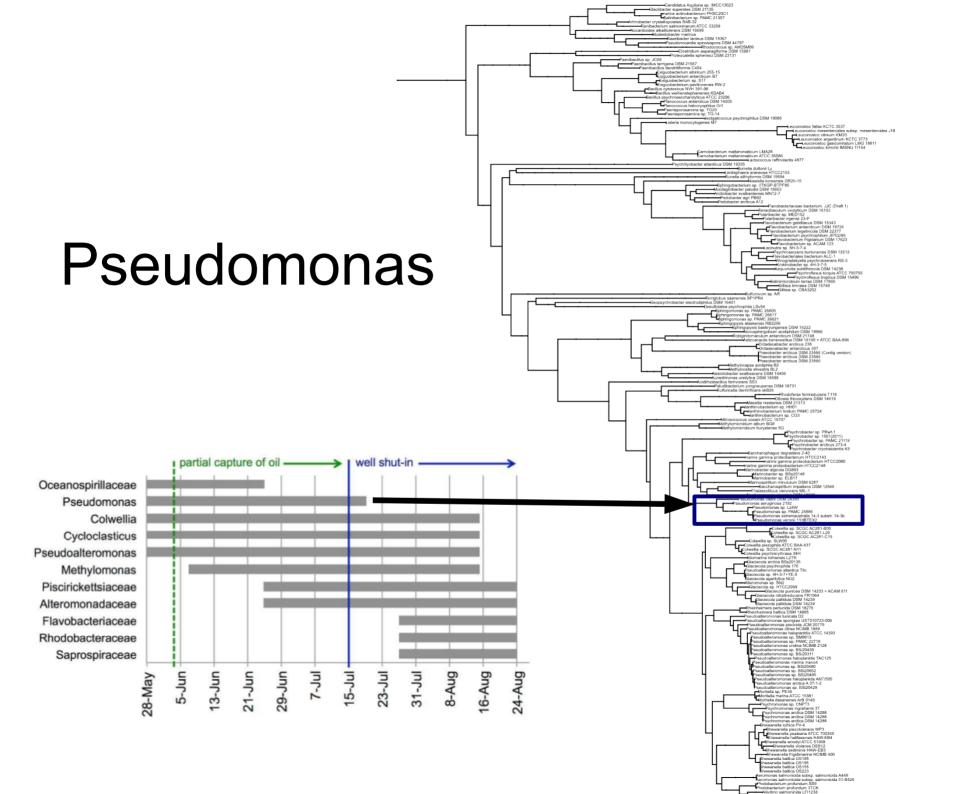


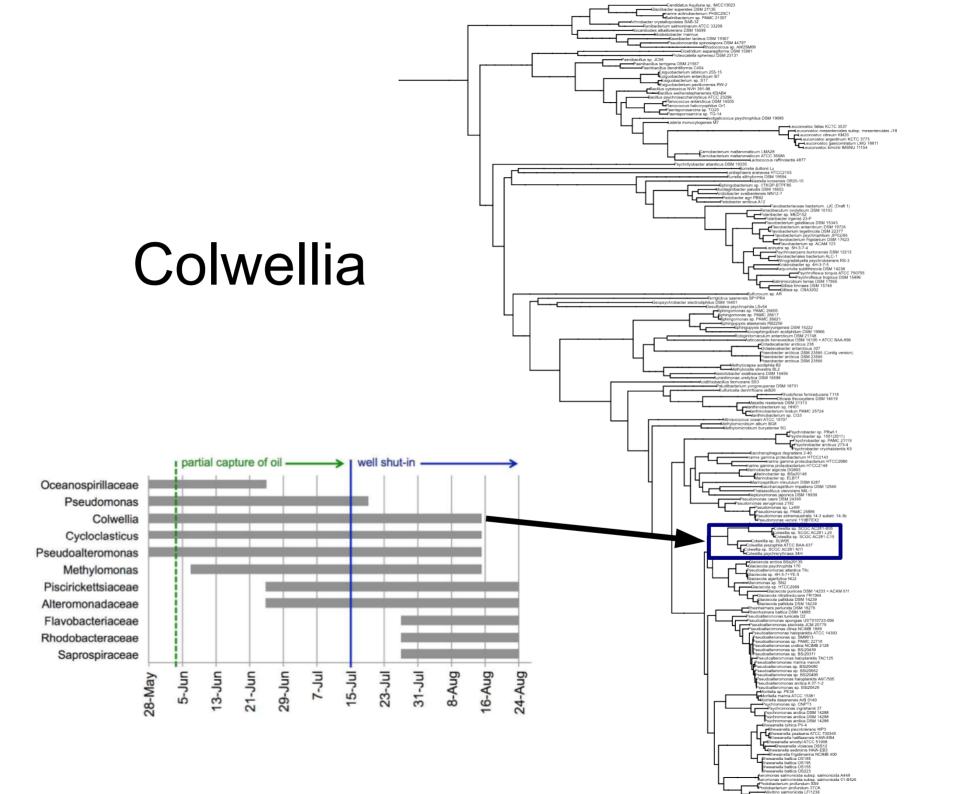


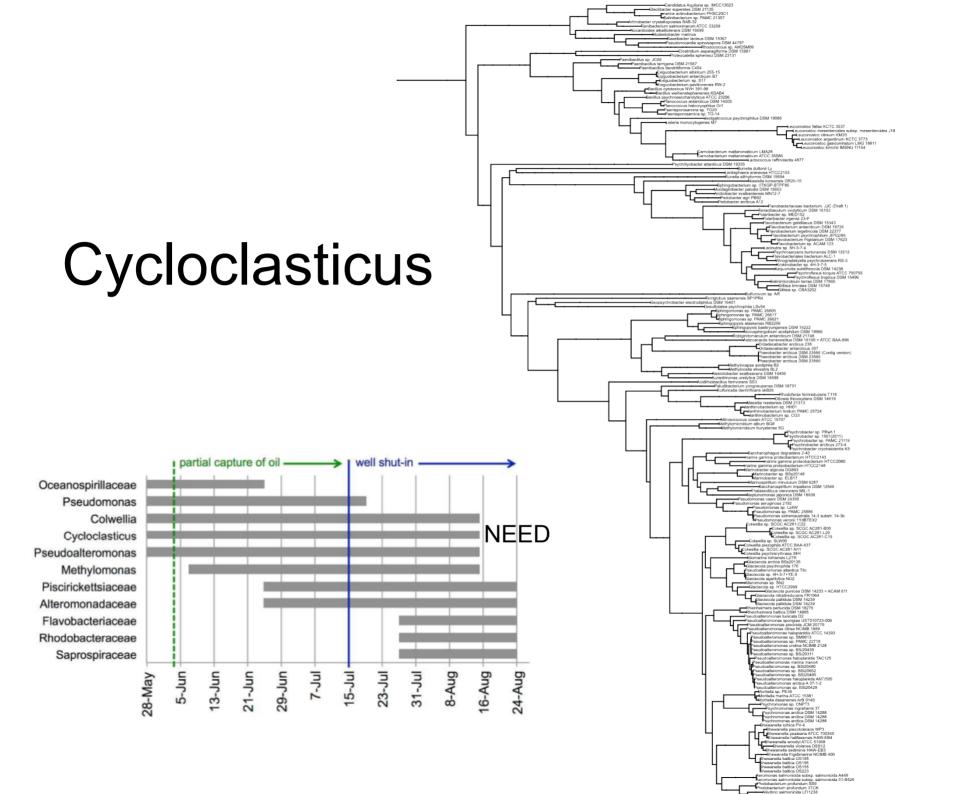
Deepwater Horizon Succession

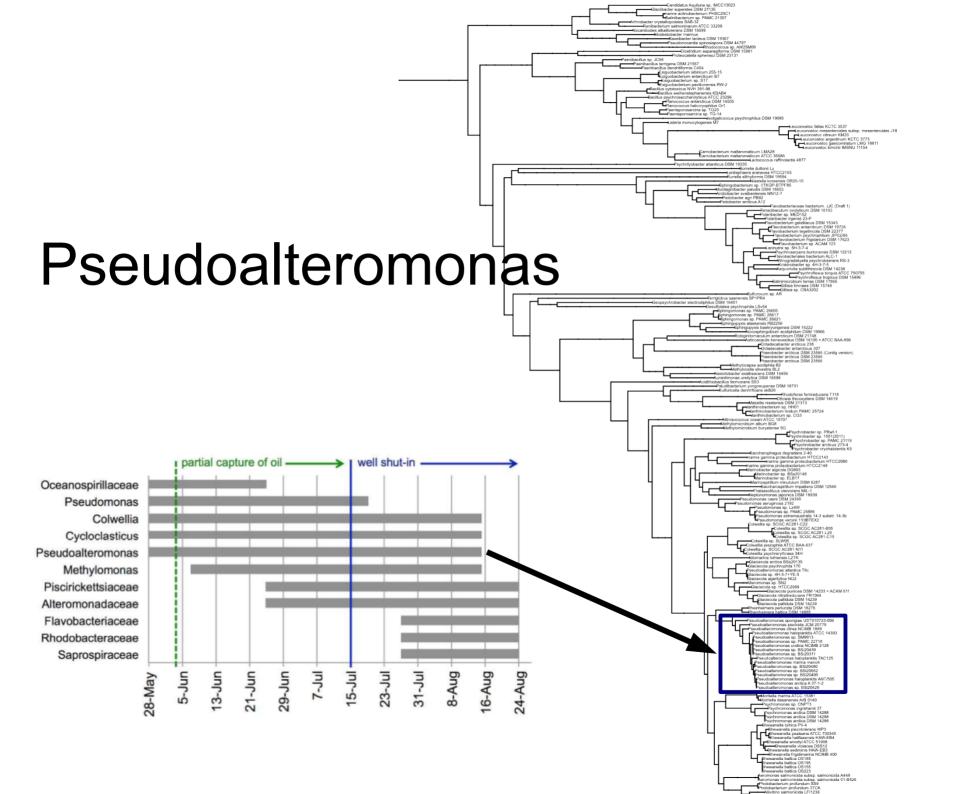


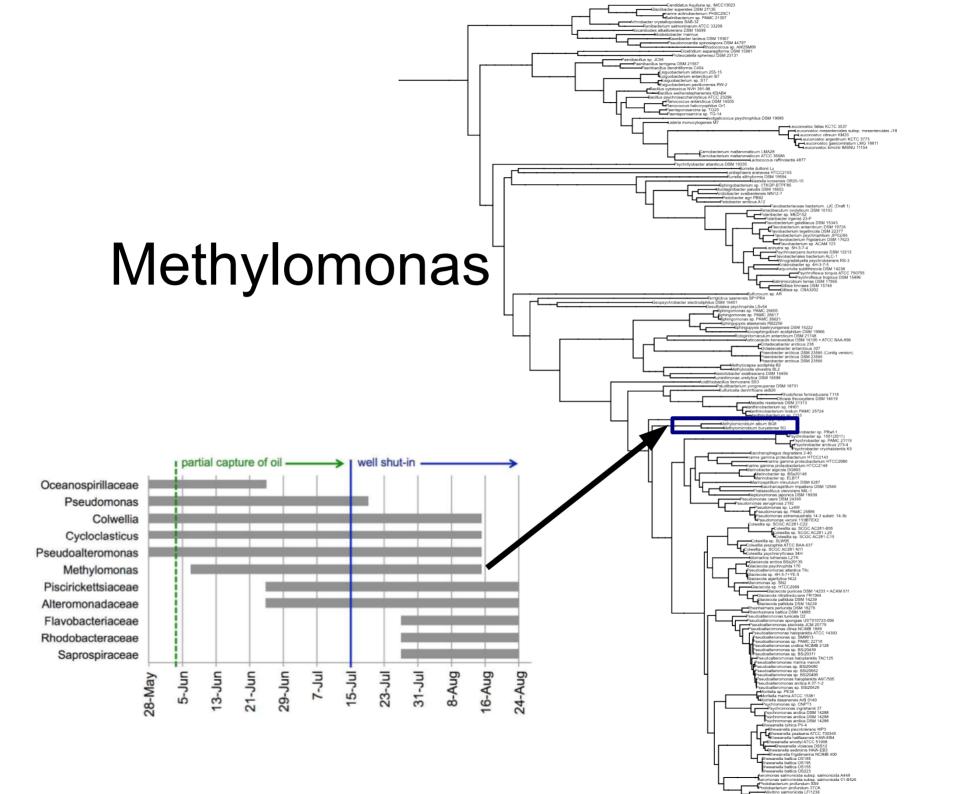




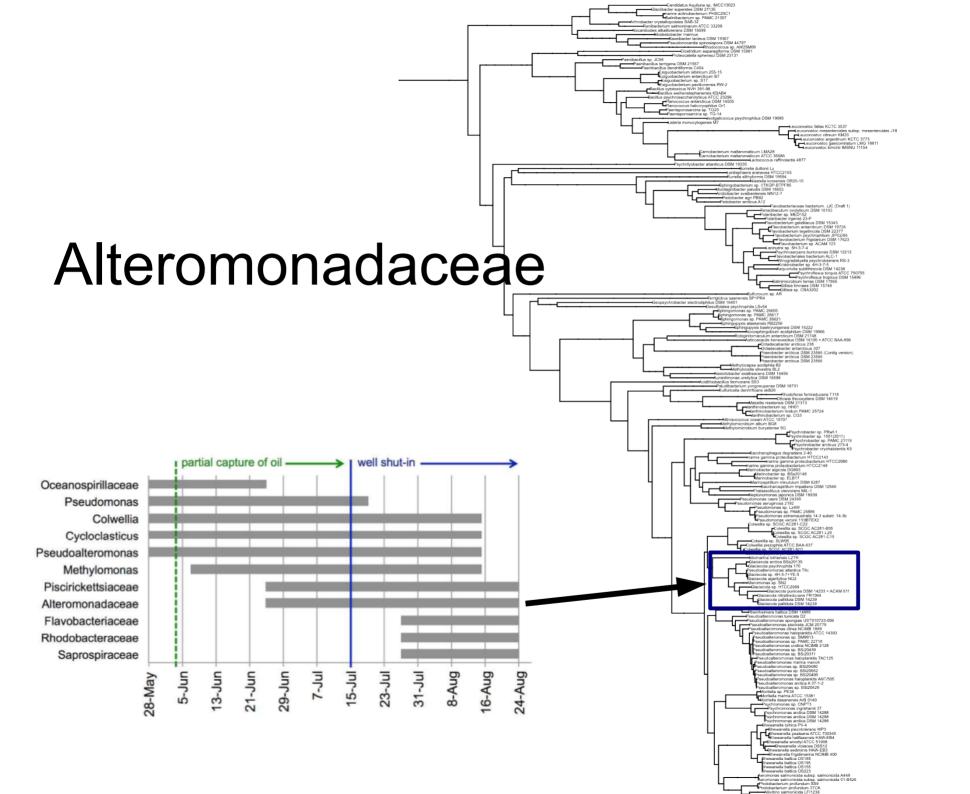


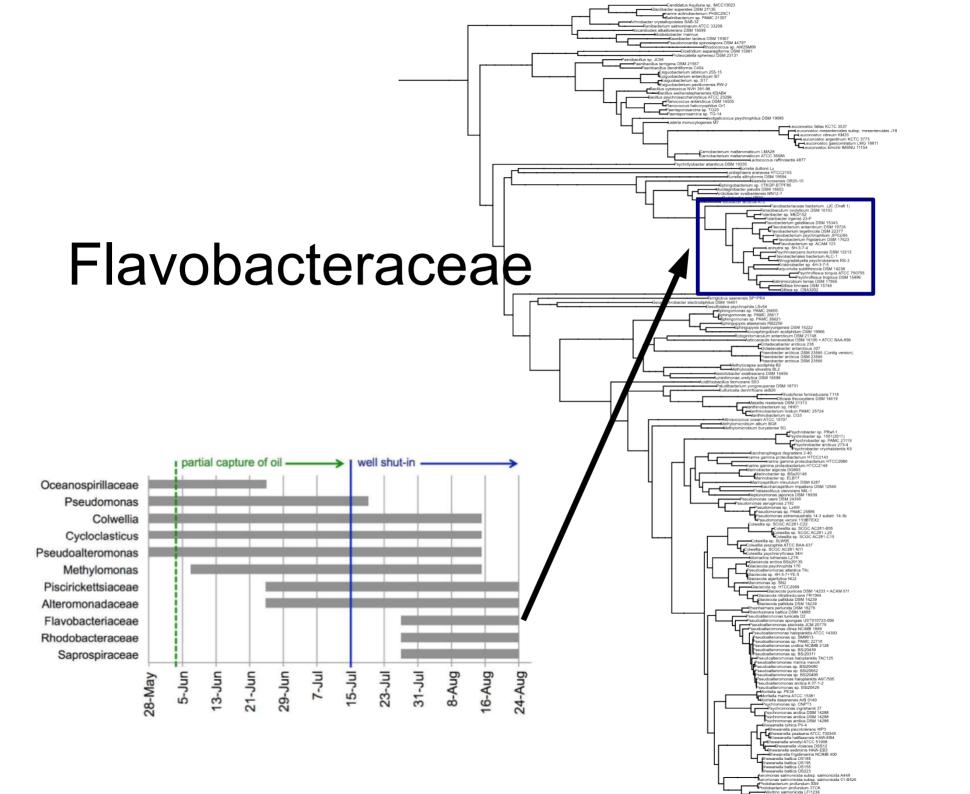


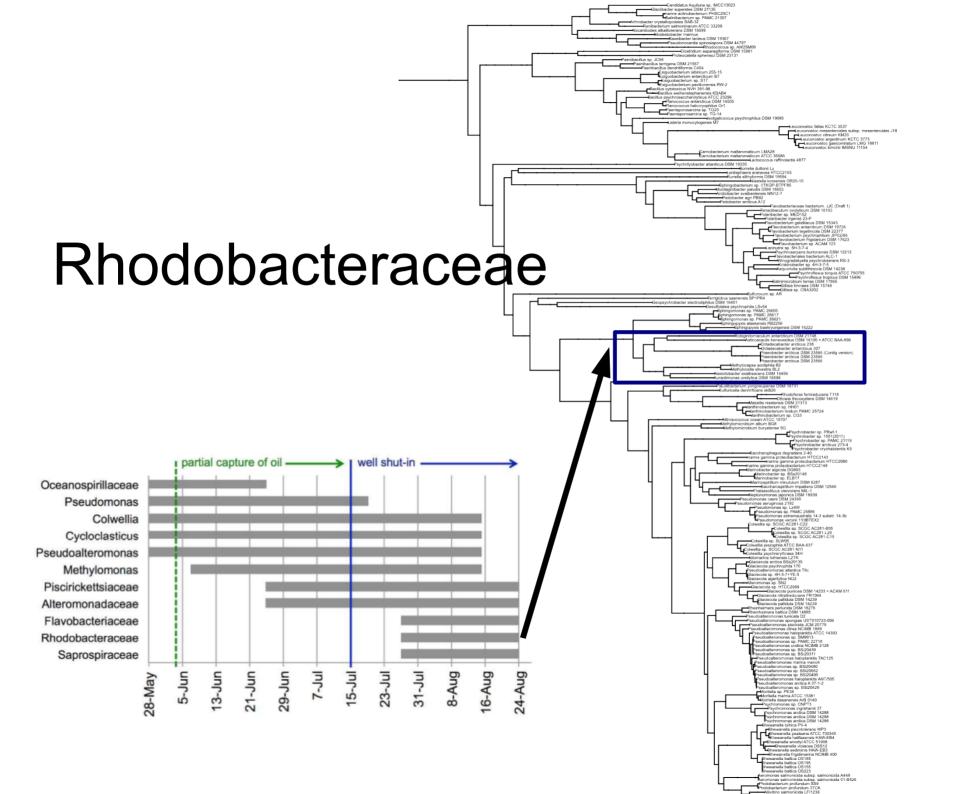


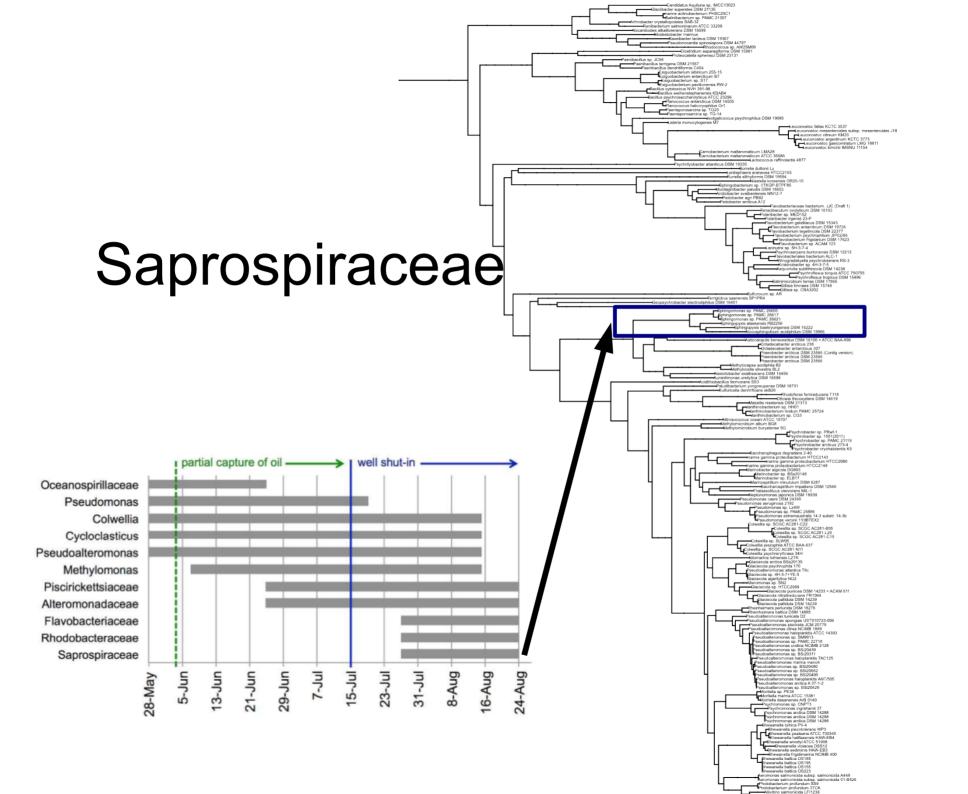






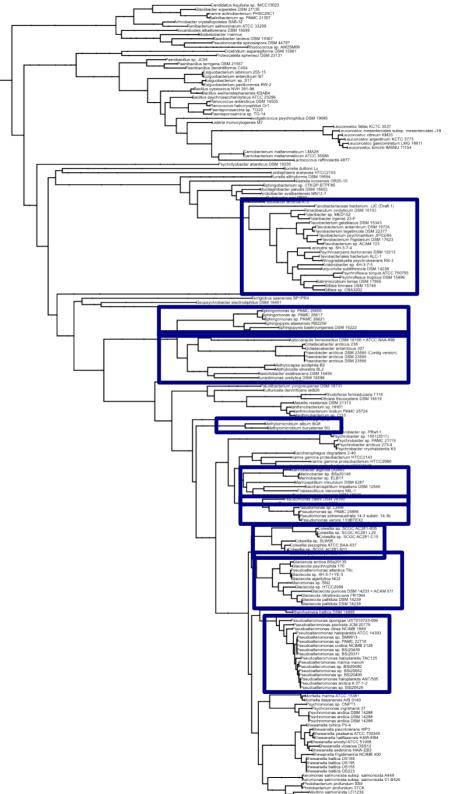






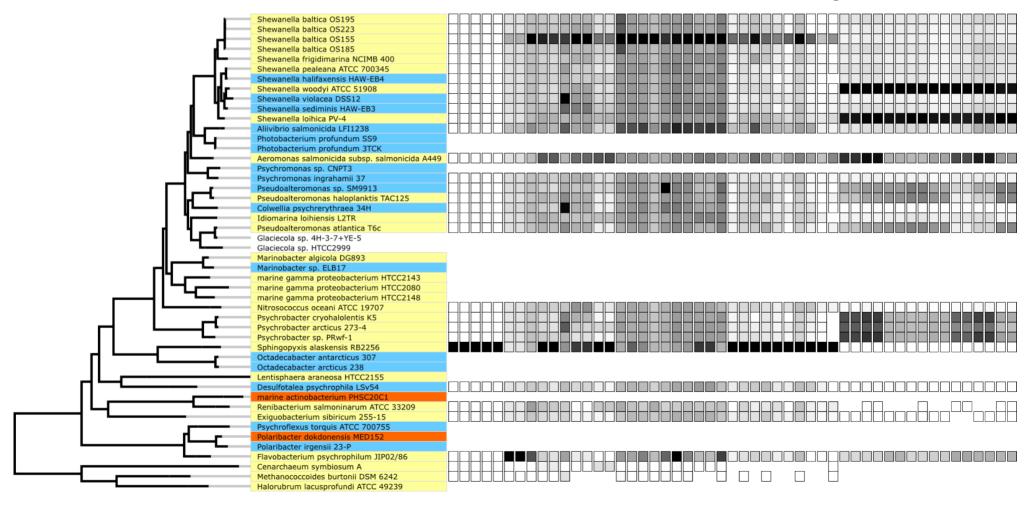
Most cold-adapted species are in oil-degrading groups

Cold ocean is 90% by volume but cold-adapted genomes are only 200/20,000 = 1%



Psychrophile Genome Biodegradation Potential

Genes for contaminant biodegradation



Recommendations:

- Application of modern sequencing & bioinformatics techniques to determine biodegradation potential of sea ice communities
- Integration of biodegradation rate measurements with biological community analysis
- Increase the relevance of lab experiments to natural communities
- Rigorous analysis of natural variability and uncertainty in environmental conditions
- Forecast of future changes in microbial community structure as a result of global warming-induced sea ice loss

