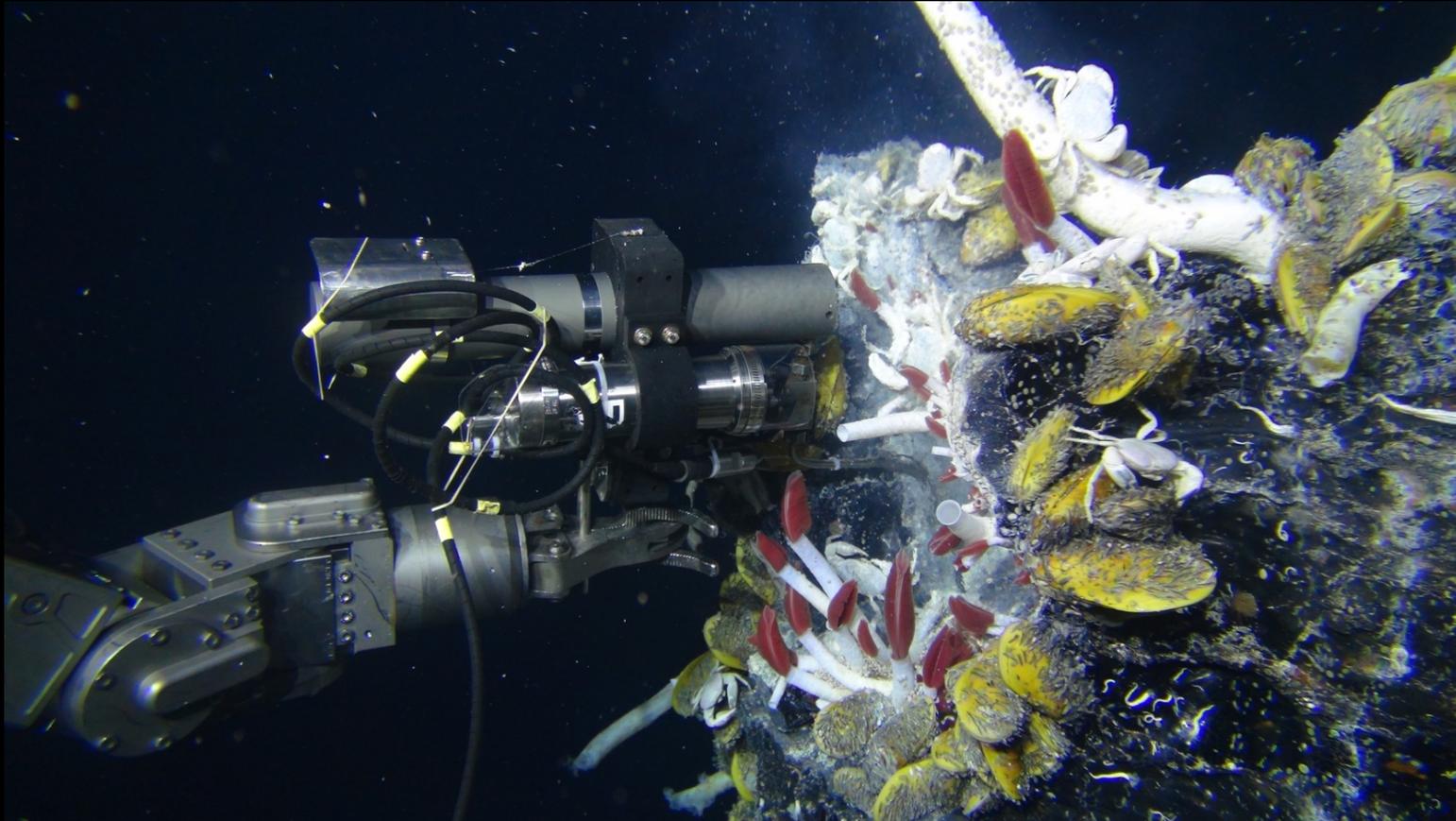


Exploring Life at Deep-Sea Hydrothermal Vents ... and what does it have to do with Medicine?

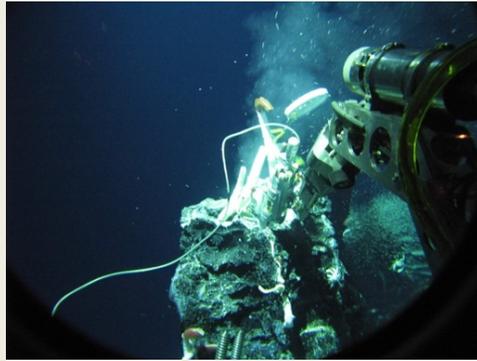


Stefan M. Sievert
Woods Hole Oceanographic Institution
ssievert@whoi.edu
<http://www.whoi.edu/people/ssievert>

SIEVERT Lab for Dark Life



Salt marsh



Deep-sea vents



Shallow-water vents



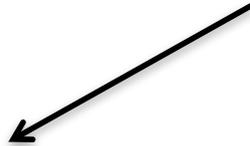
Oxygen minimum zones

CHEMOSYNTHESIS

Chemical Energy + Carbon Dioxide



Living cells



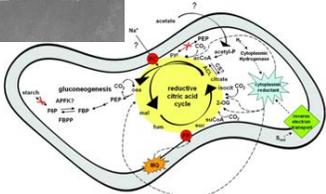
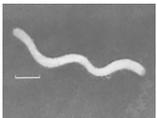
Who is there?



Who is active?



What process(es) are they mediating and at what rate(s)?

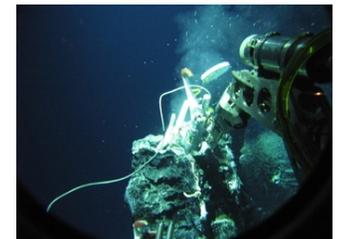


Organism



Ecosystem

Physiology
Biochemistry
Genetics
Genomics
Proteomics



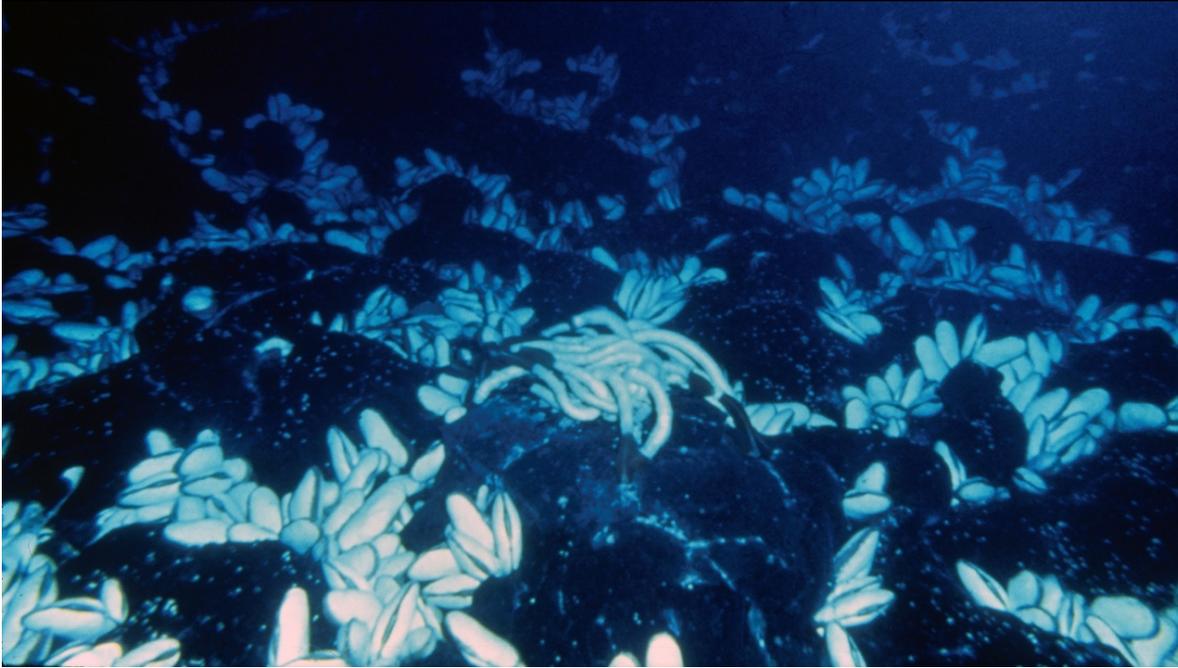
Abundance, Diversity
Functions (Meta 'Omics')
Rates
Geochemistry
Interactions (biotic, abiotic)

<http://www.whoi.edu/groups/sievertlab>

Outline

- Introduction to Deep-Sea Vents
 - Geological and Physical Setting
 - Chemosynthesis
- Microbial Processes and Symbiosis – Link to Medicine
- Fieldwork at 9°N East Pacific Rise
 - Study site and approach
 - Productivity of subseafloor biosphere
- Synthesis

Discovery of Deep-Sea Hydrothermal Vents only ~40 years ago in 1977



Submarine Thermal Springs on the Galápagos Rift

16 March 1979, Volume 203, Number 4385

SCIENCE

John B. Corliss, Jack Dymond, Louis I. Gordon,
John M. Edmond, Richard P. von Herzen, Robert D. Ballard,
Kenneth Green, David Williams, Arnold Bainbridge,
Kathy Crane, Tjeerd H. van Andel

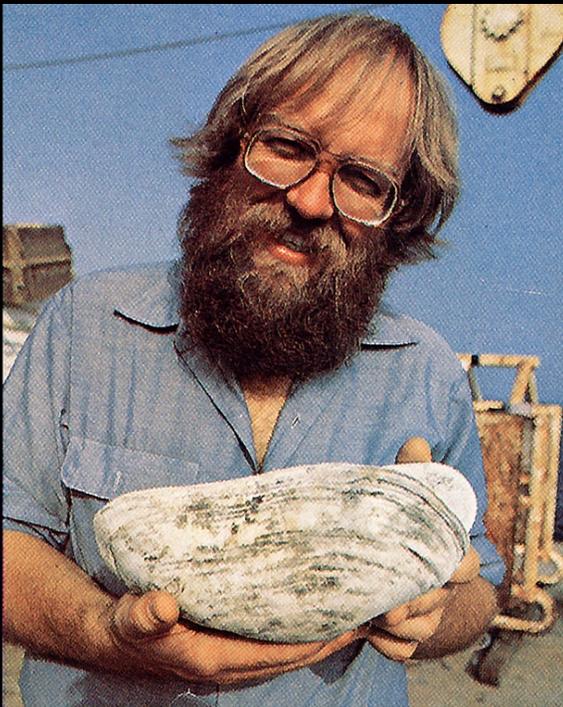


John (Jack) Corliss
(OSU)

Bob Ballard
(WHOI)



Discovery of Deep-Sea Hydrothermal Vents only ~40 years ago in 1977



John (Jack) Corliss
(OSU)

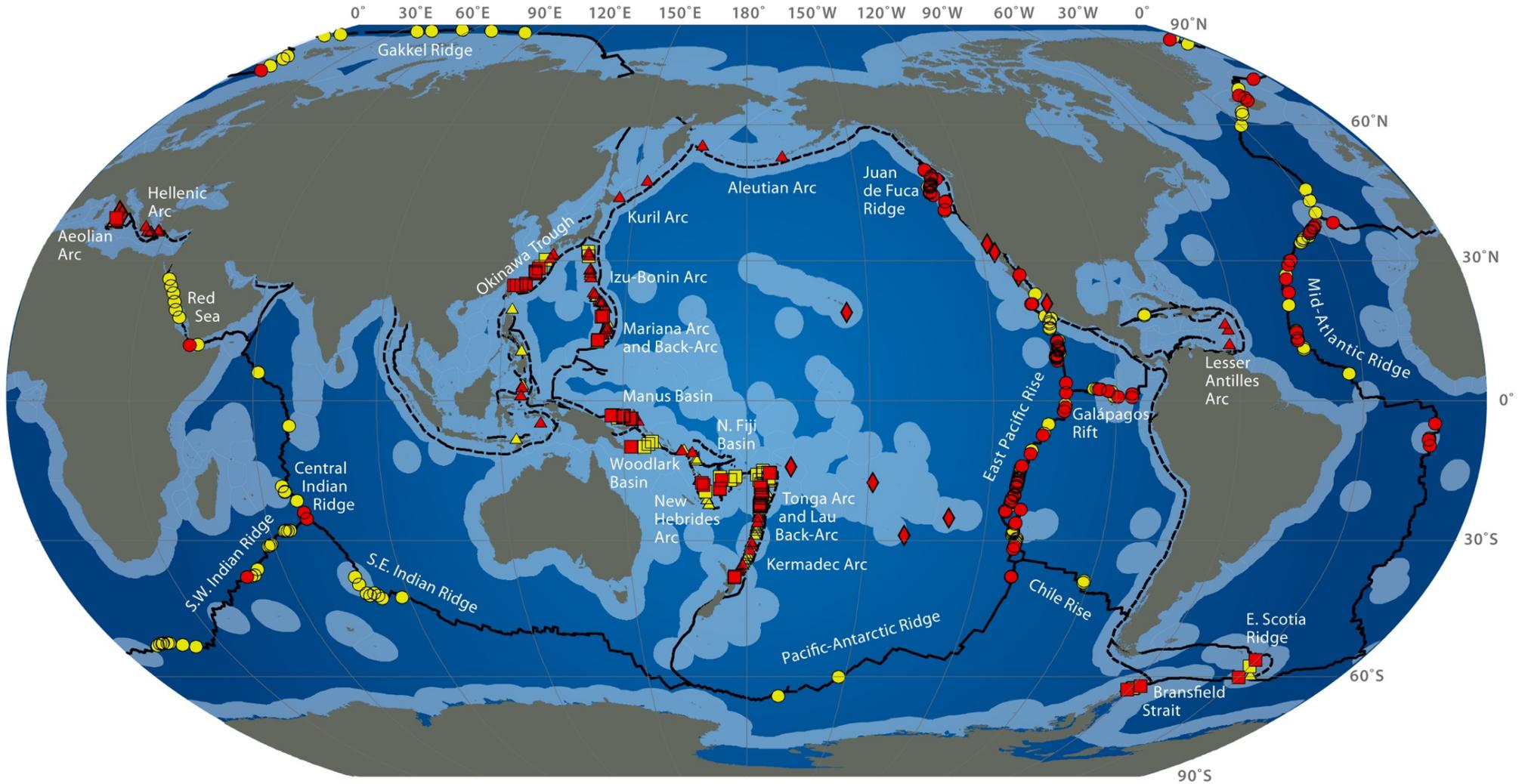


Bob Ballard
(WHOI)

Global Distribution of Hydrothermal Vent Fields



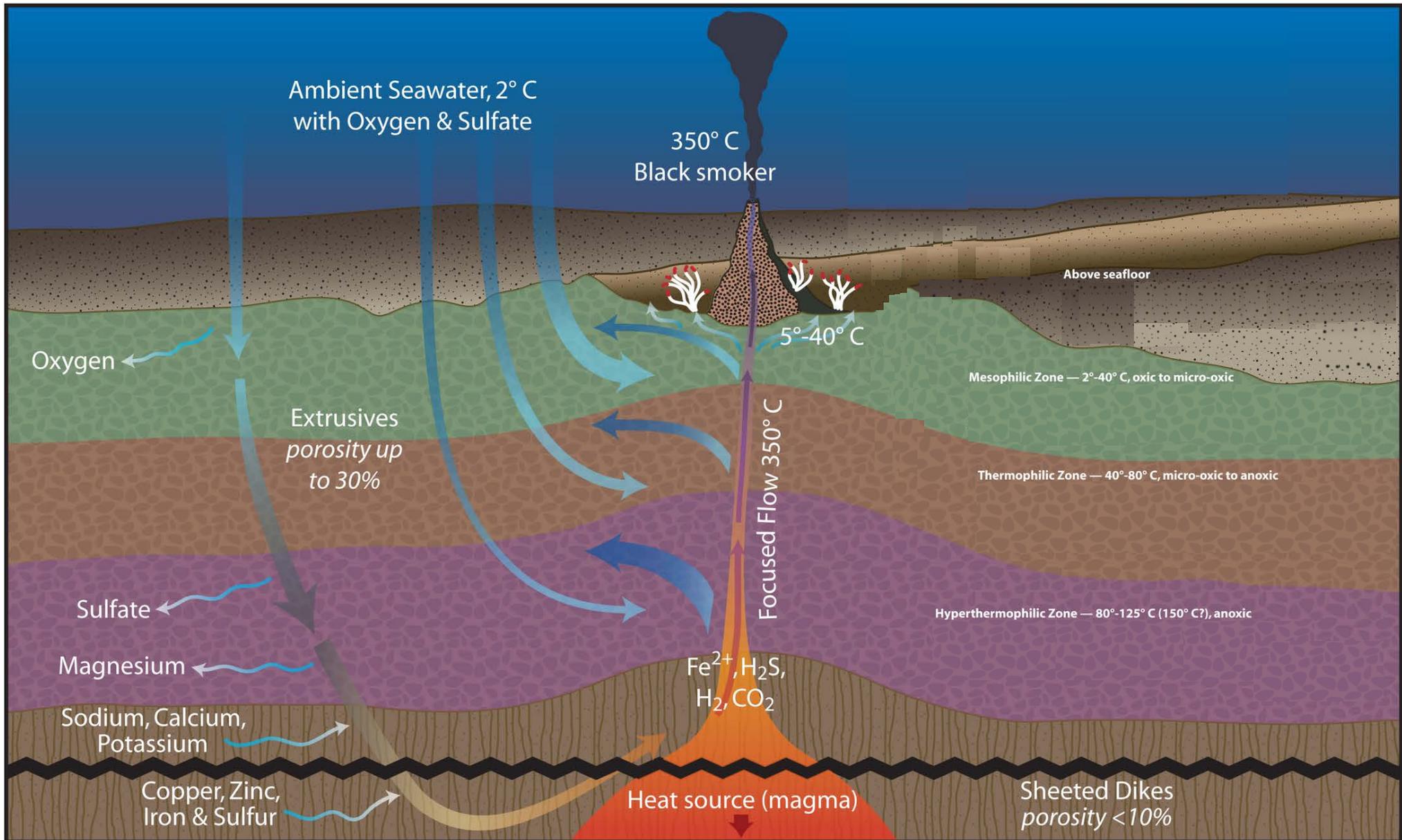
Global Distribution of Hydrothermal Vent Fields



Mid-ocean ridge	Arc volcano	Back-arc spreading center	Intra-plate volcano & Other	Ridge & Transform
● Active	▲ Active	■ Active	◆ Active	— Trench
● Unconfirmed	▲ Unconfirmed	■ Unconfirmed		● Exclusive Economic Zones

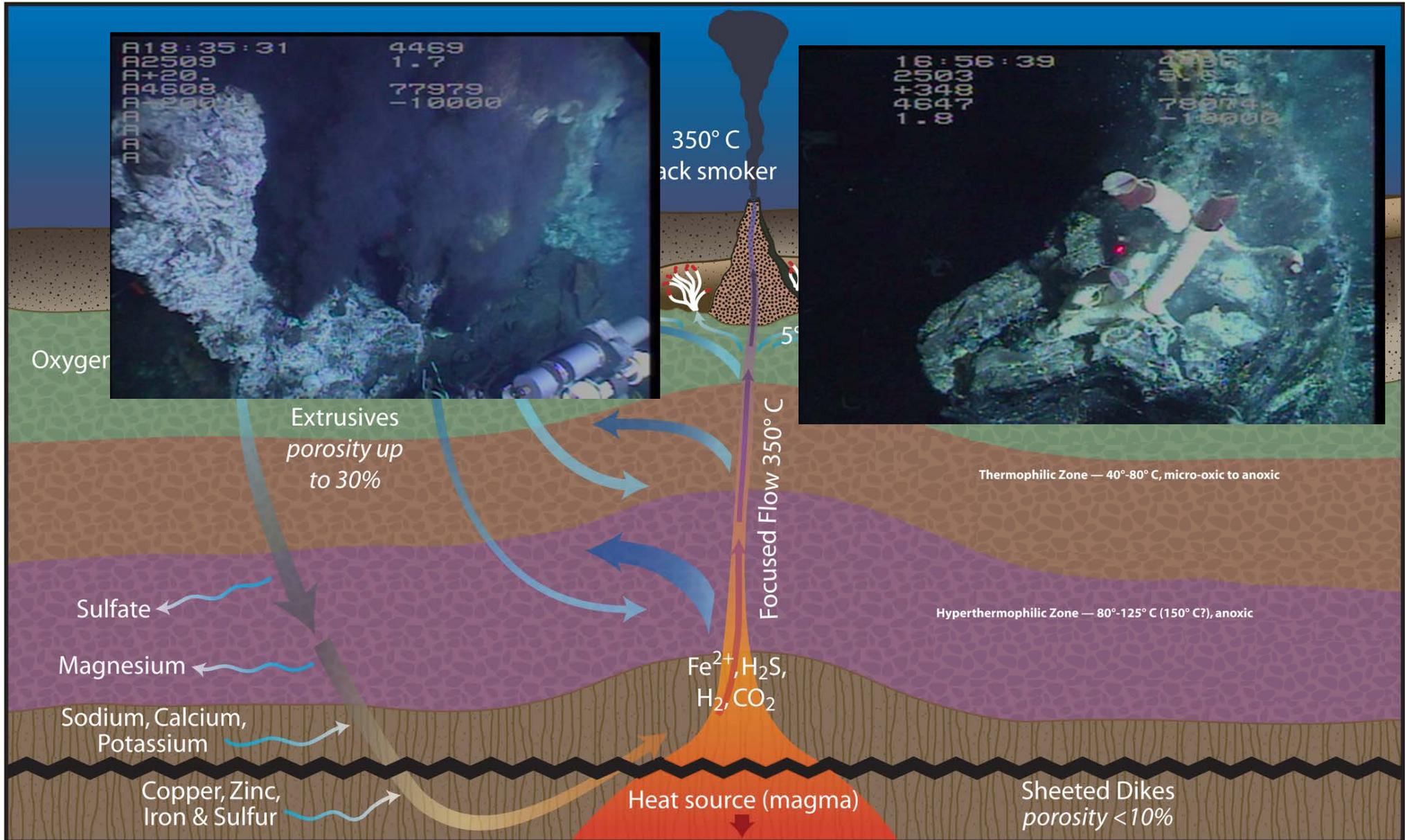


Physical Setting

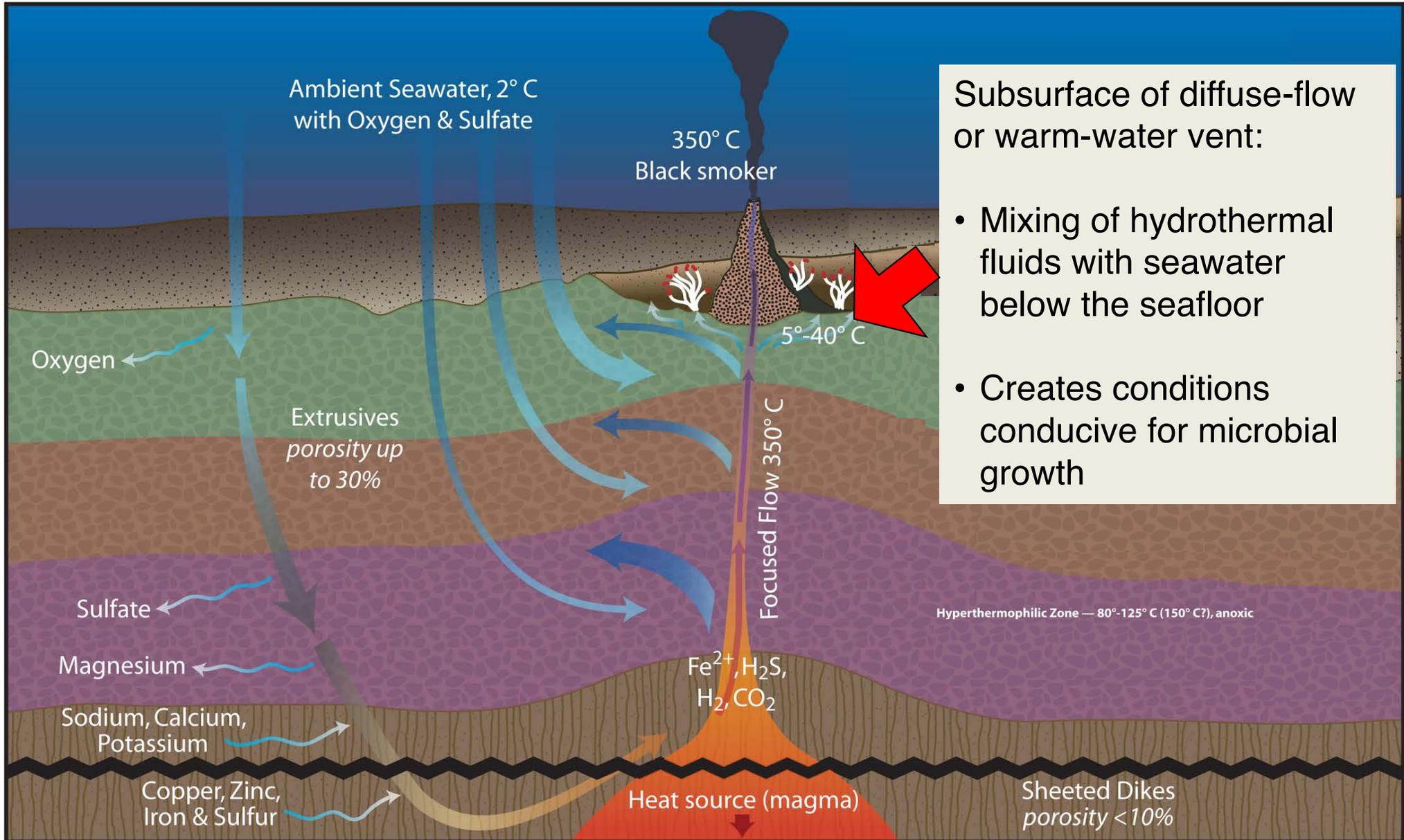


Sievert et al., 2008

Physical Setting



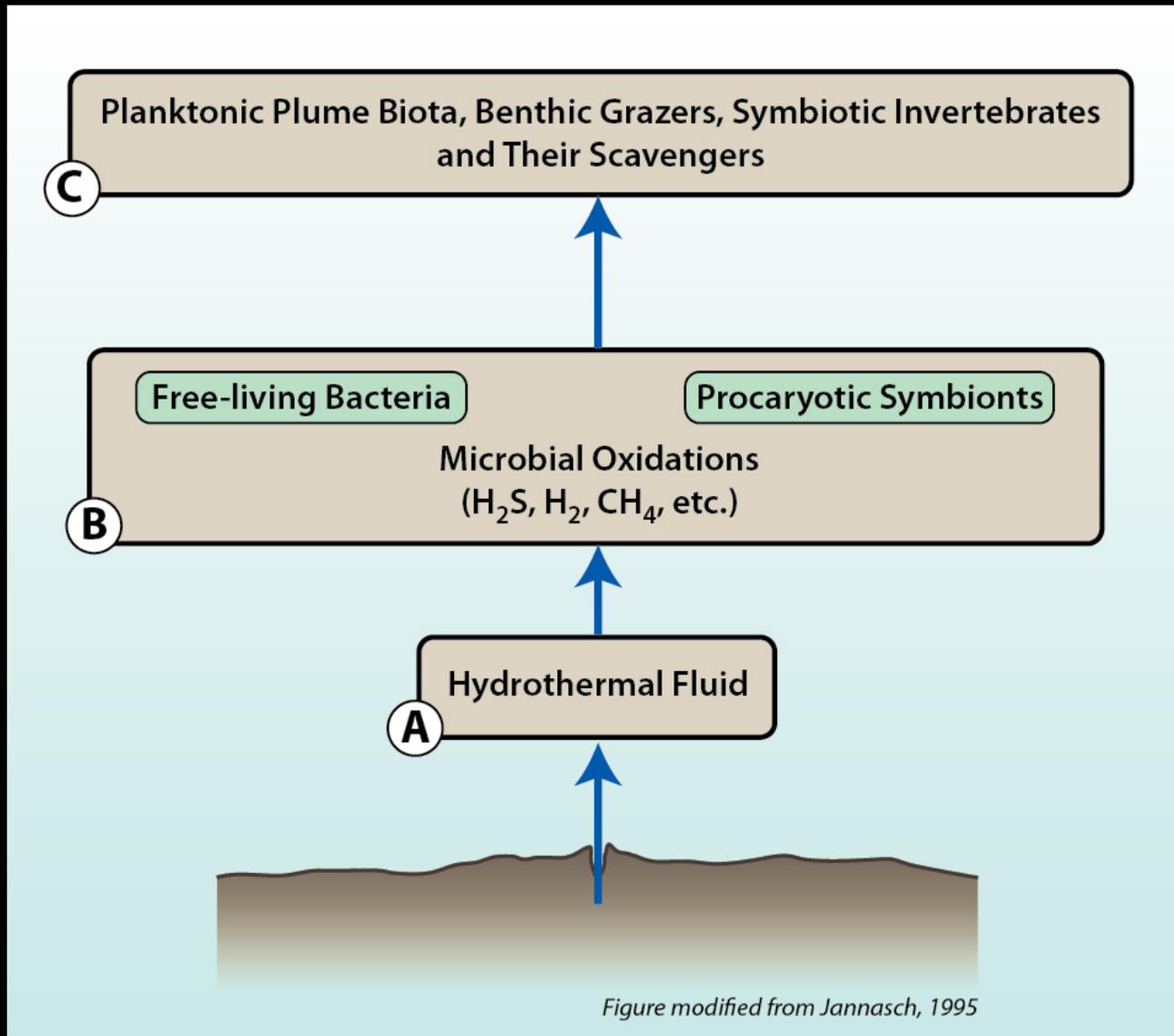
Physical Setting



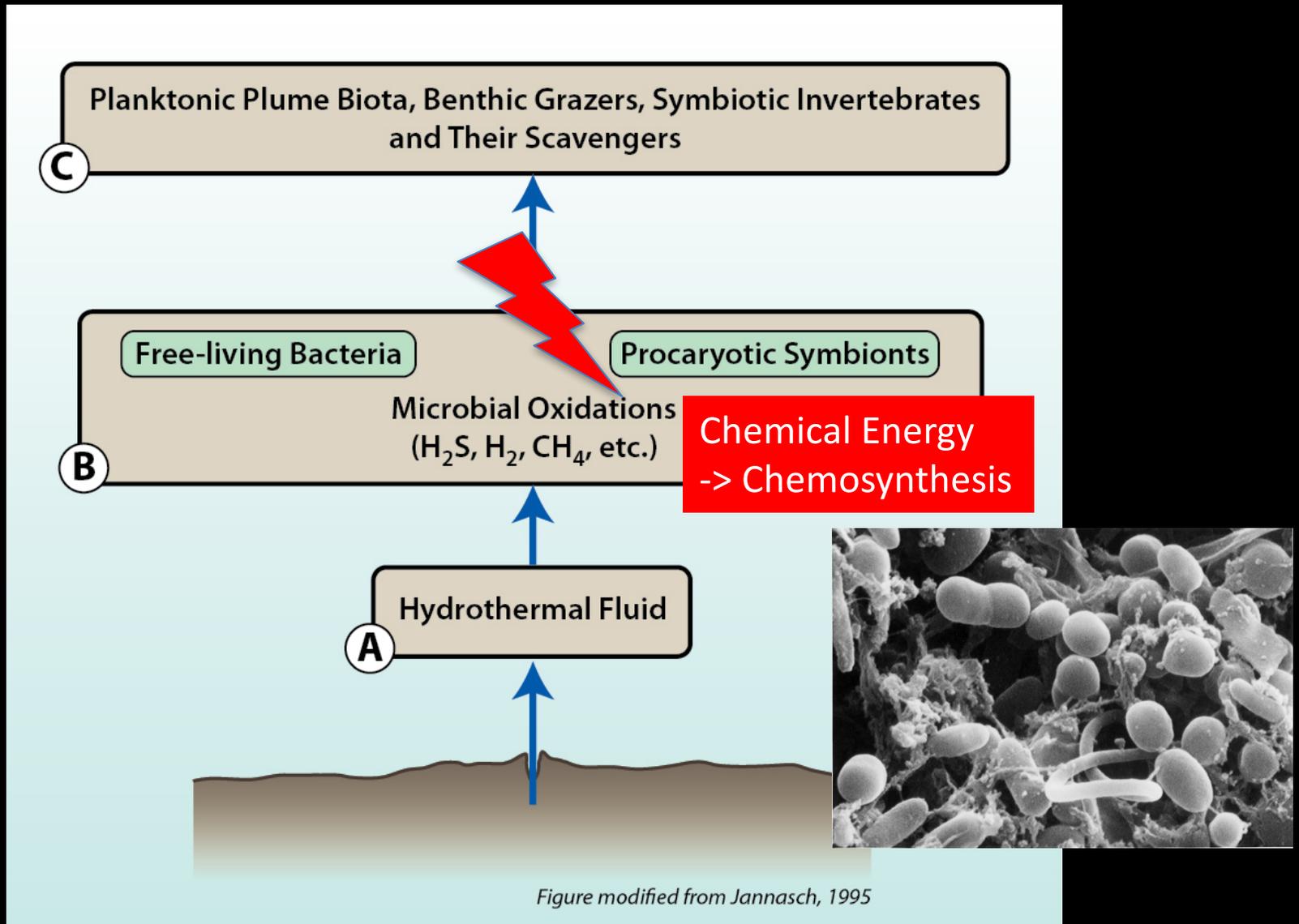
Subsurface of diffuse-flow
or warm-water vent:

- Mixing of hydrothermal fluids with seawater below the seafloor
- Creates conditions conducive for microbial growth

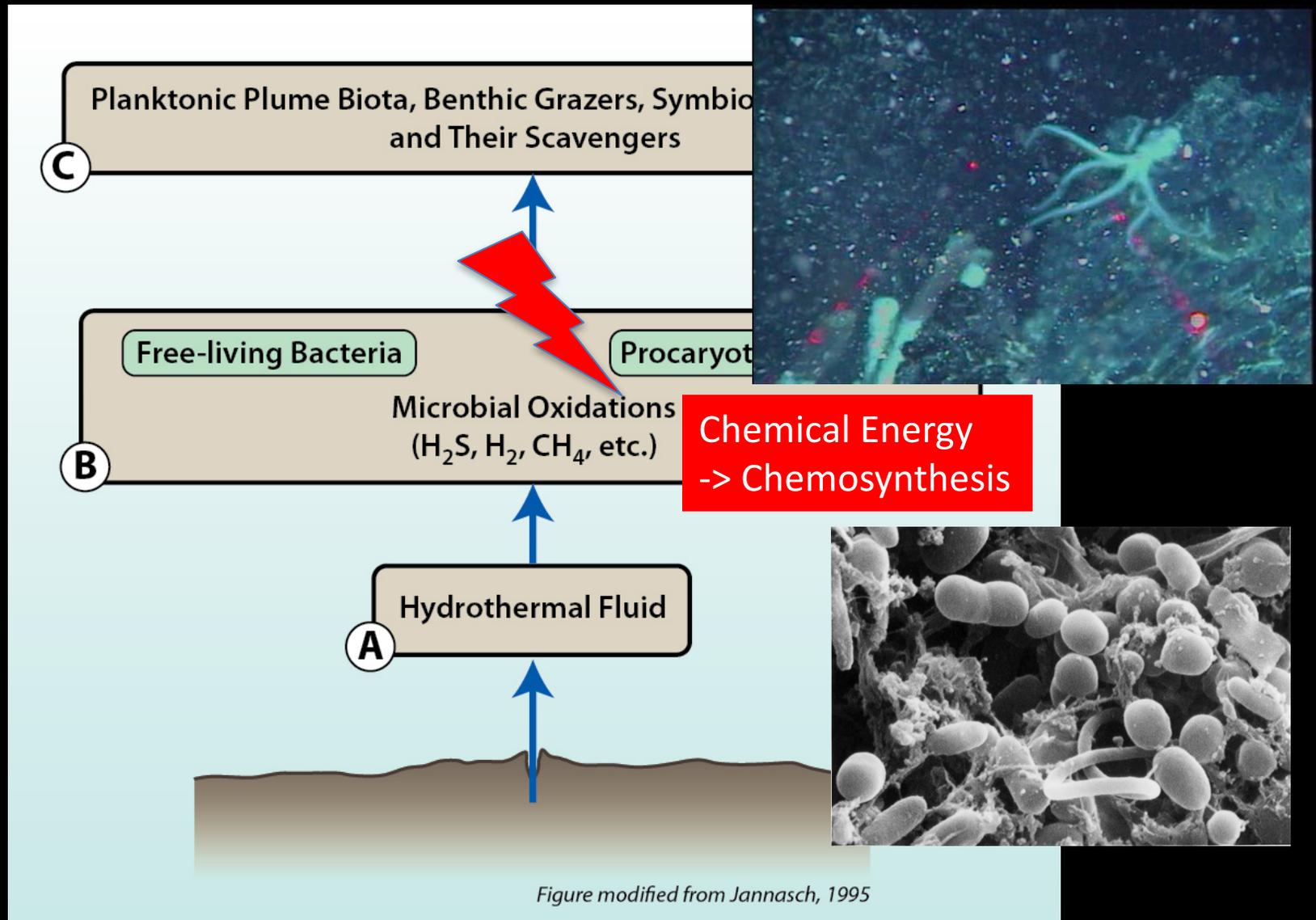
“FOOD CHAIN” AT DEEP-SEA VENTS: CONCEPTUAL FRAMEWORK



"FOOD CHAIN" AT DEEP-SEA VENTS: CONCEPTUAL FRAMEWORK

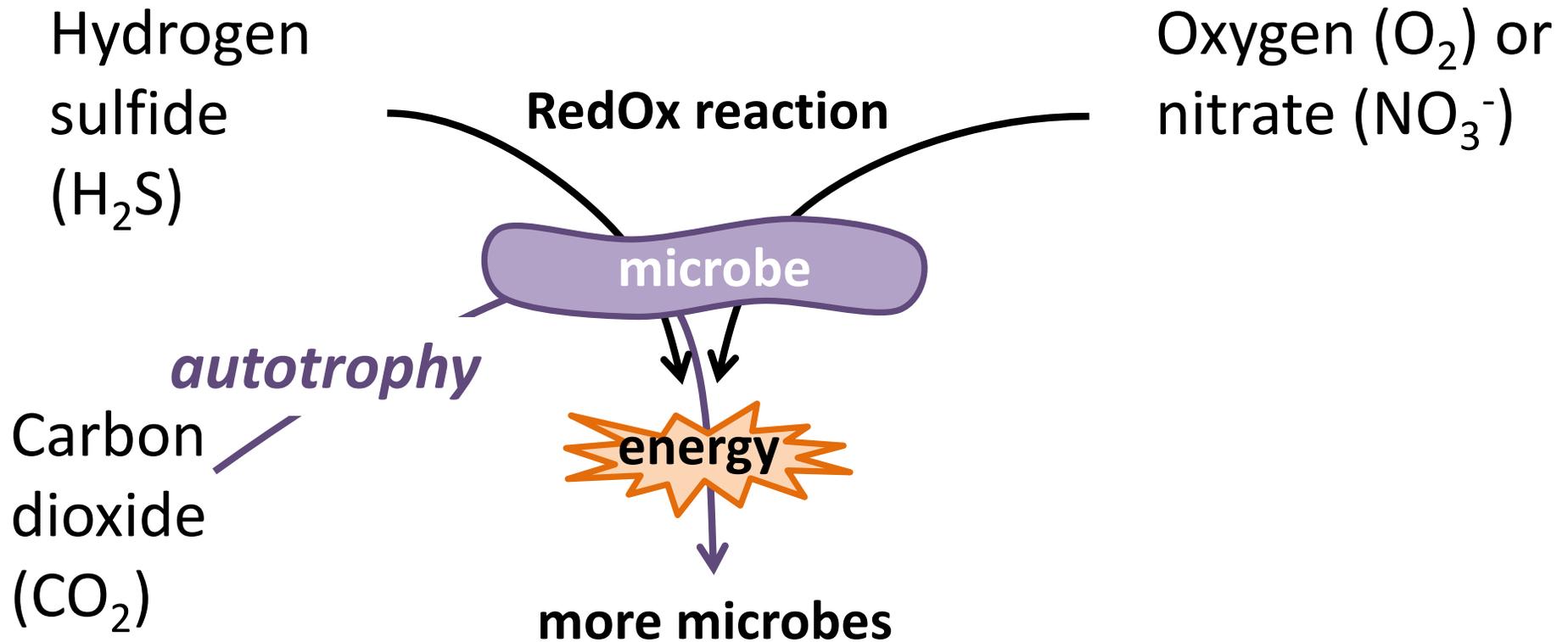


“FOOD CHAIN” AT DEEP-SEA VENTS: CONCEPTUAL FRAMEWORK



Microbes are at the Base of Ecosystem by Mediating the Transfer of Energy from Geothermal Source to Higher Trophic Levels

Chemosynthesis



Photosynthesis – Foundation for life on land and surface ocean

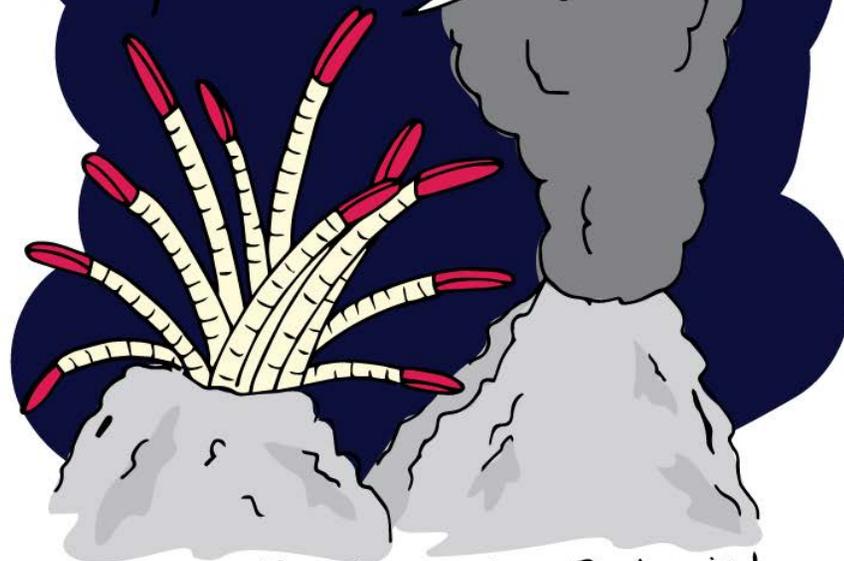
Chemosynthesis – Foundation for life at hydrothermal vents

Meanwhile, at the bottom
of the ocean...

Do you think life
exists at the
surface of the ocean?

Nah. There are no
hydrothermal vents
up there. What would
they use for energy?

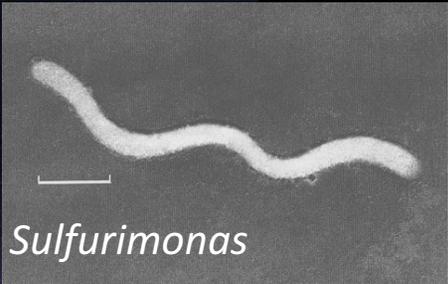
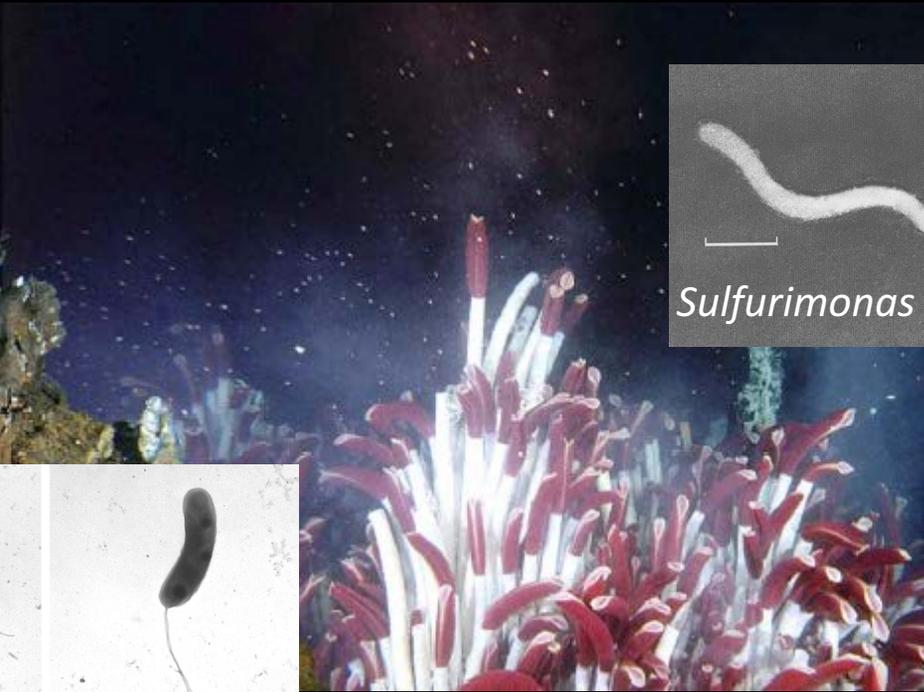
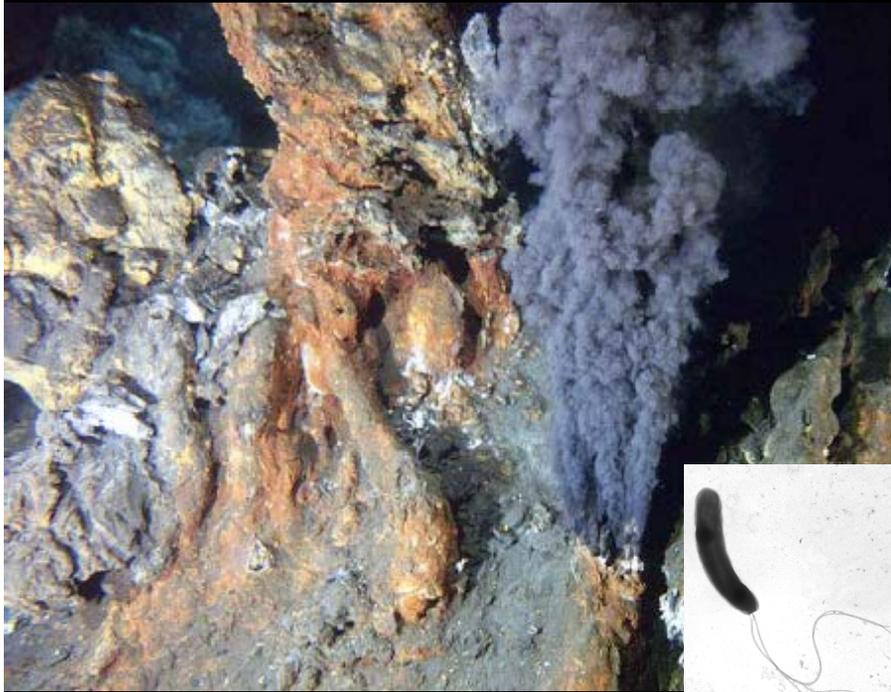
Mmm, true.



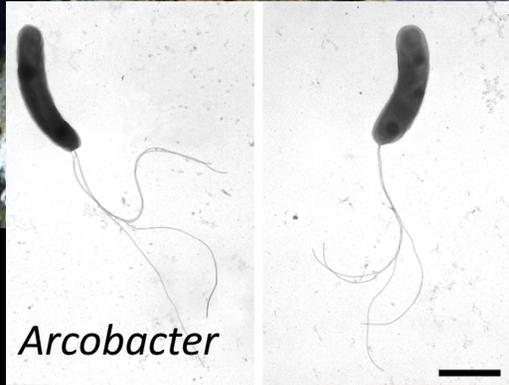
Beatrice the Biologist

Katie McKissick

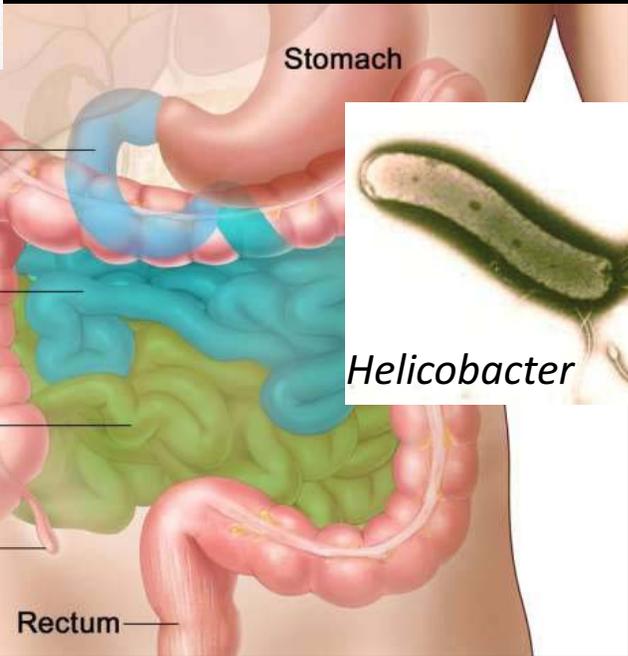
<http://www.beatricebiologist.com>



Sulfurimonas



Arcobacter



Stomach

Duodenum

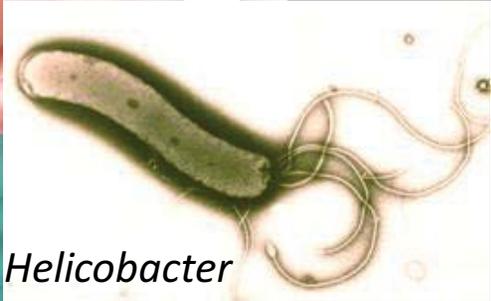
Small intestine

Jejunum

Ileum

Appendix

Rectum



Helicobacter

What's in common?

Campylobacteria

Formerly known as *Epsilonproteobacteria*



Campylobacter

The 5300-year-old *Helicobacter pylori* genome of the Iceman

Frank Maixner,^{1*†} Ben Krause-Kyora,^{2†} Dmitriy Turaev,^{3†} Alexander Herbig,^{4,5†} Michael R. Hoopmann,⁶ Janice L. Hallows,⁶ Ulrike Kusebauch,⁶ Eduard Egarter Vigl,⁷ Peter Malfertheiner,⁸ Francis Megraud,⁹ Niall O’Sullivan,¹ Giovanna Cipollini,¹ Valentina Coia,¹ Marco Samadelli,¹ Lars Engstrand,¹⁰ Bodo Linz,¹¹ Robert L. Moritz,⁶ Rudolf Grimm,¹² Johannes Krause,^{4,5†} Almut Nebel,^{2†} Yoshan Moodley,^{13,14†} Thomas Rattei,^{3†} Albert Zink^{1*‡}

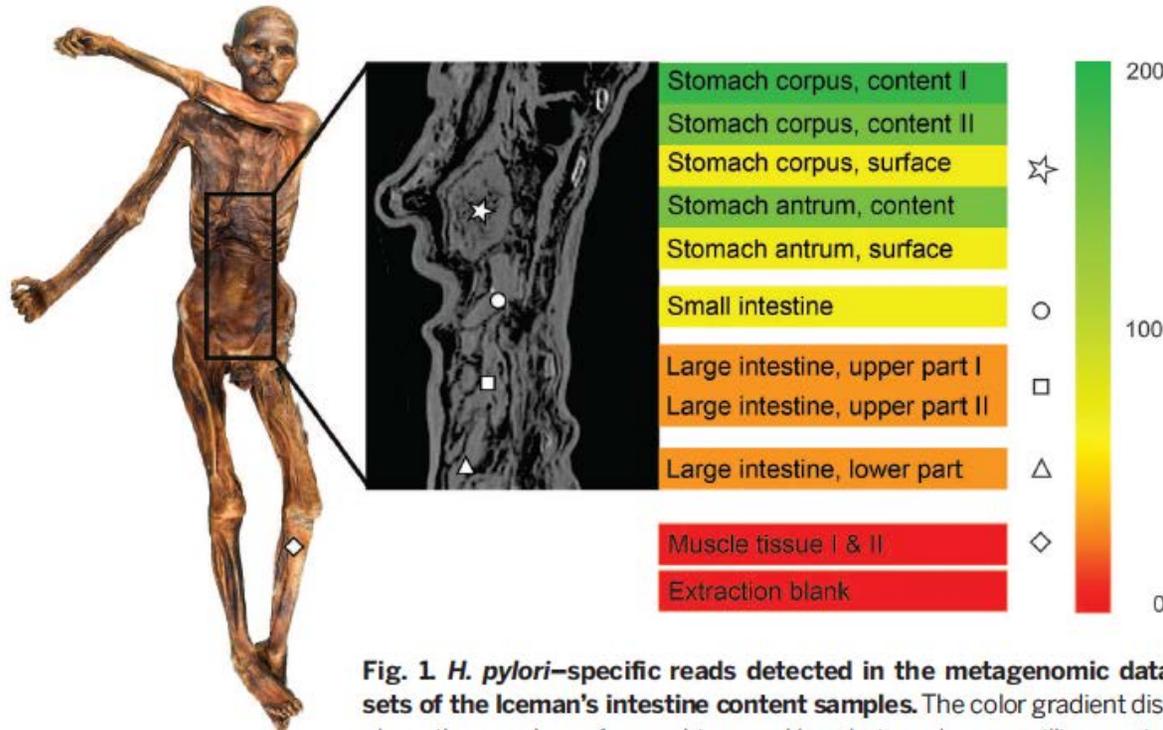
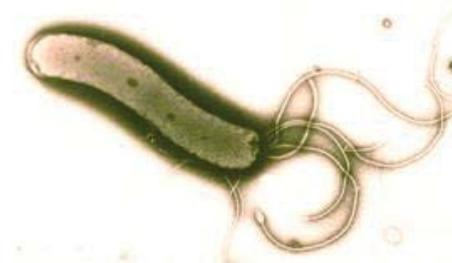


Fig. 1 *H. pylori*-specific reads detected in the metagenomic data sets of the Iceman's intestine content samples. The color gradient displays the number of unambiguous *H. pylori* reads per million meta-

genomic reads. Control metagenomic data sets of the Iceman's muscle tissue and of the extraction blank were included in the analysis. The different intestinal content sampling sites are marked in the radiographic image by the following symbols: asterisk, stomach content; circle, small intestine; square, upper large intestine; triangle, lower large intestine. The sampling site of the muscle control sample is highlighted in the Iceman overview picture (diamond).



Eduard Egarter-Vigl (left) and Albert Zink (right) taking a sample from the Iceman in November 2010. Credit: © EURAC/Marion Lafogler

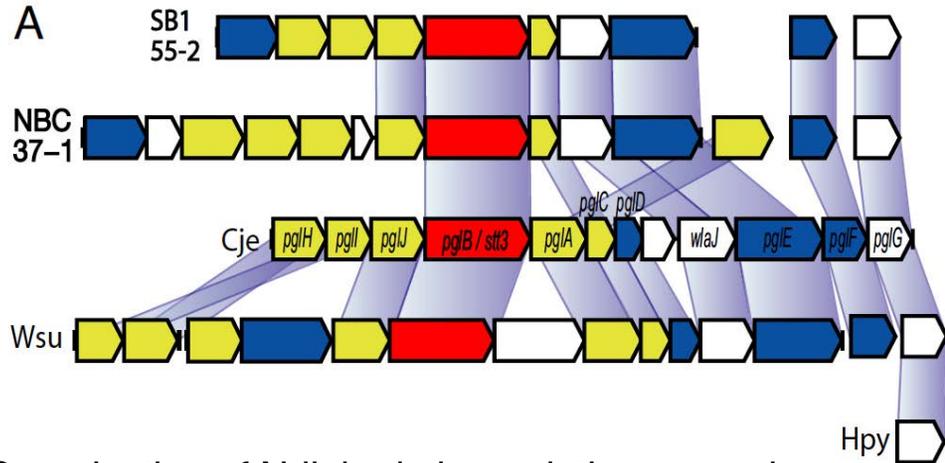
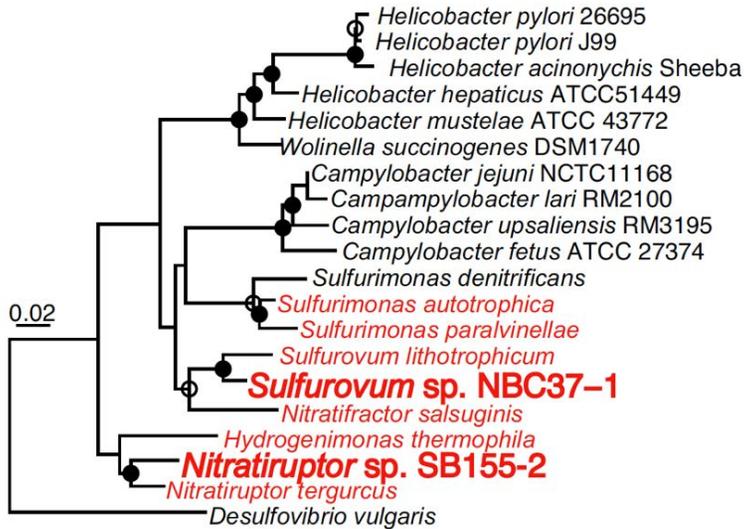




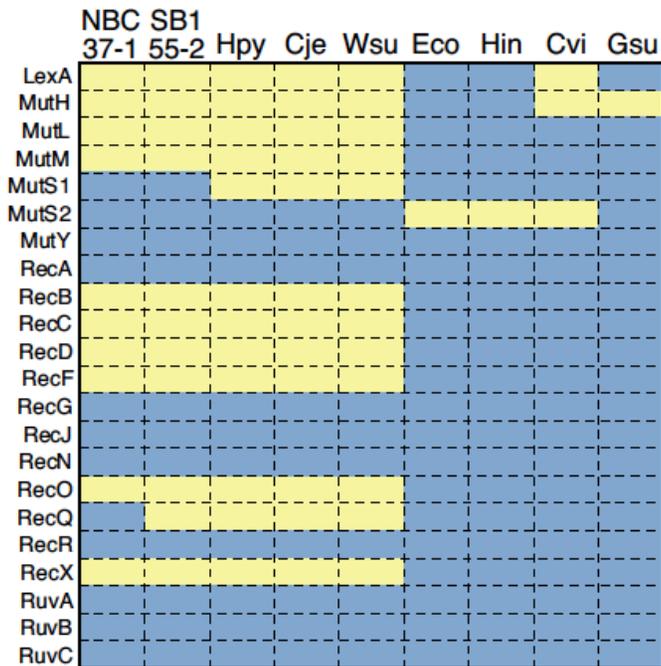
Deep-sea vent ϵ -proteobacterial genomes provide insights into emergence of pathogens

12146–12150 | PNAS | July 17, 2007 | vol. 104 | no. 29

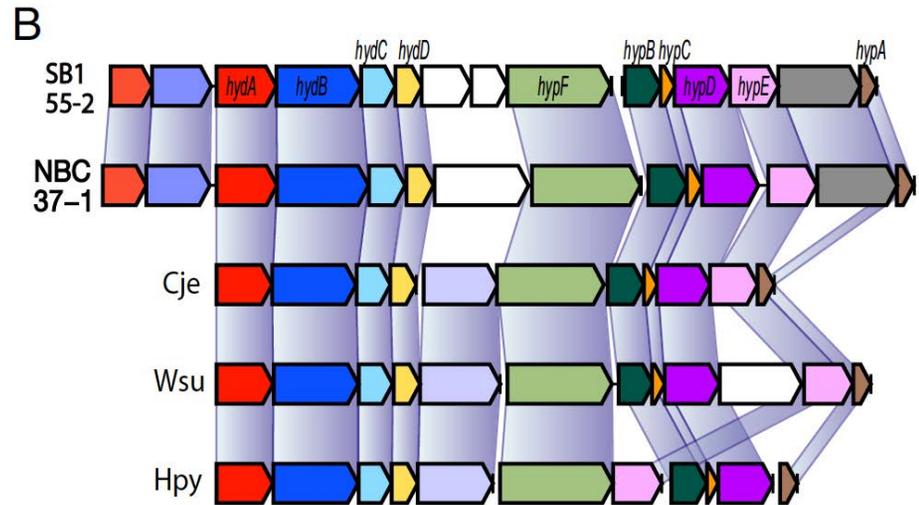
Satoshi Nakagawa^{1,2}, Yoshihiro Takaki⁵, Shigeru Shimamura⁵, Anna-Louise Reysenbach¹, Ken Takai⁴, and Koki Horikoshi⁵



Organization of N-linked glycosylation gene clusters



DNA repair genes



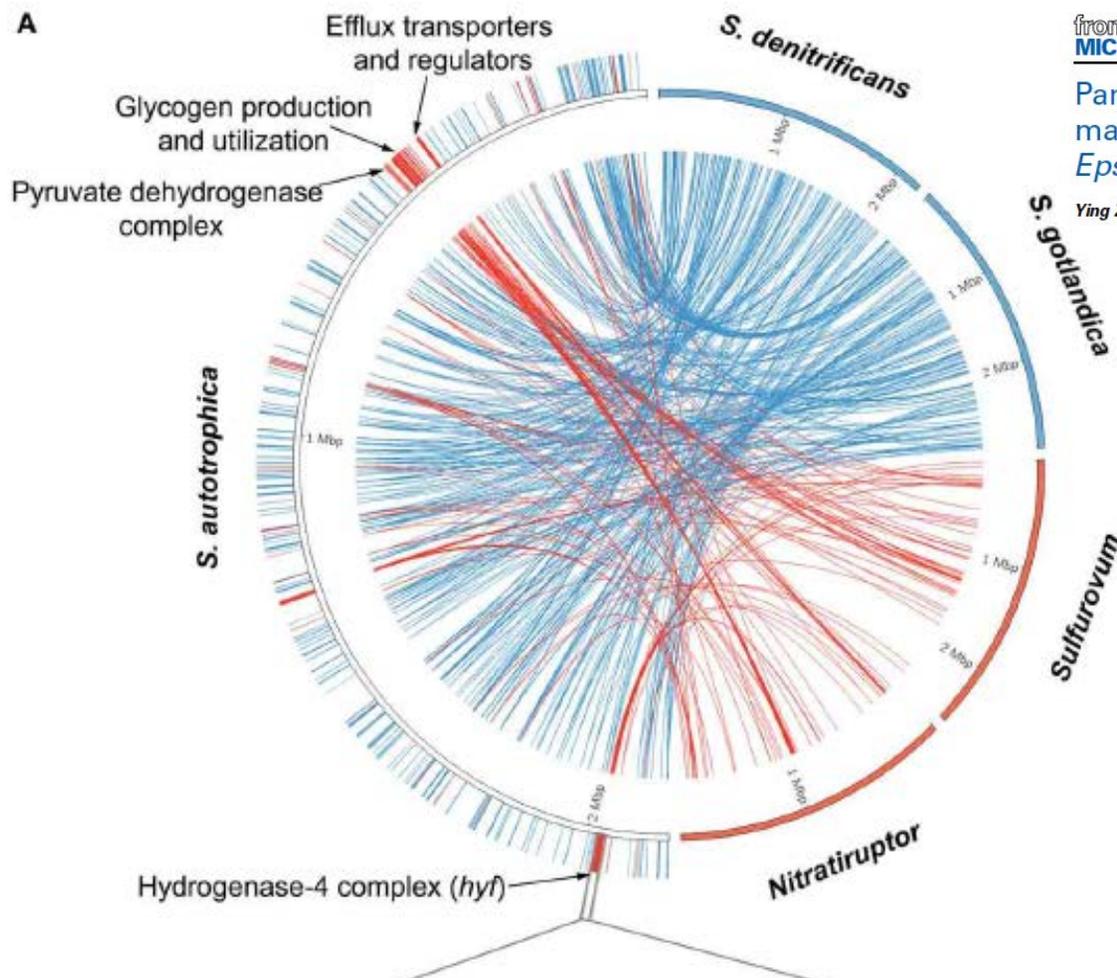
Organization of H_2 -uptake and -sensing hydrogenase gene clusters

Deep-sea vent chemoautotrophy has provided the core of virulence for important human/animal pathogens!

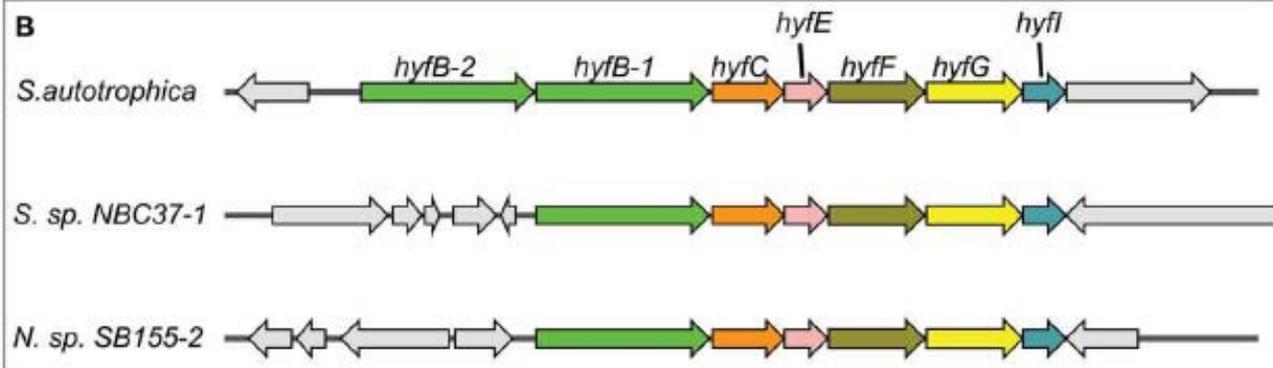


Pan-genome analyses identify lineage- and niche-specific markers of evolution and adaptation in *Epsilonproteobacteria*

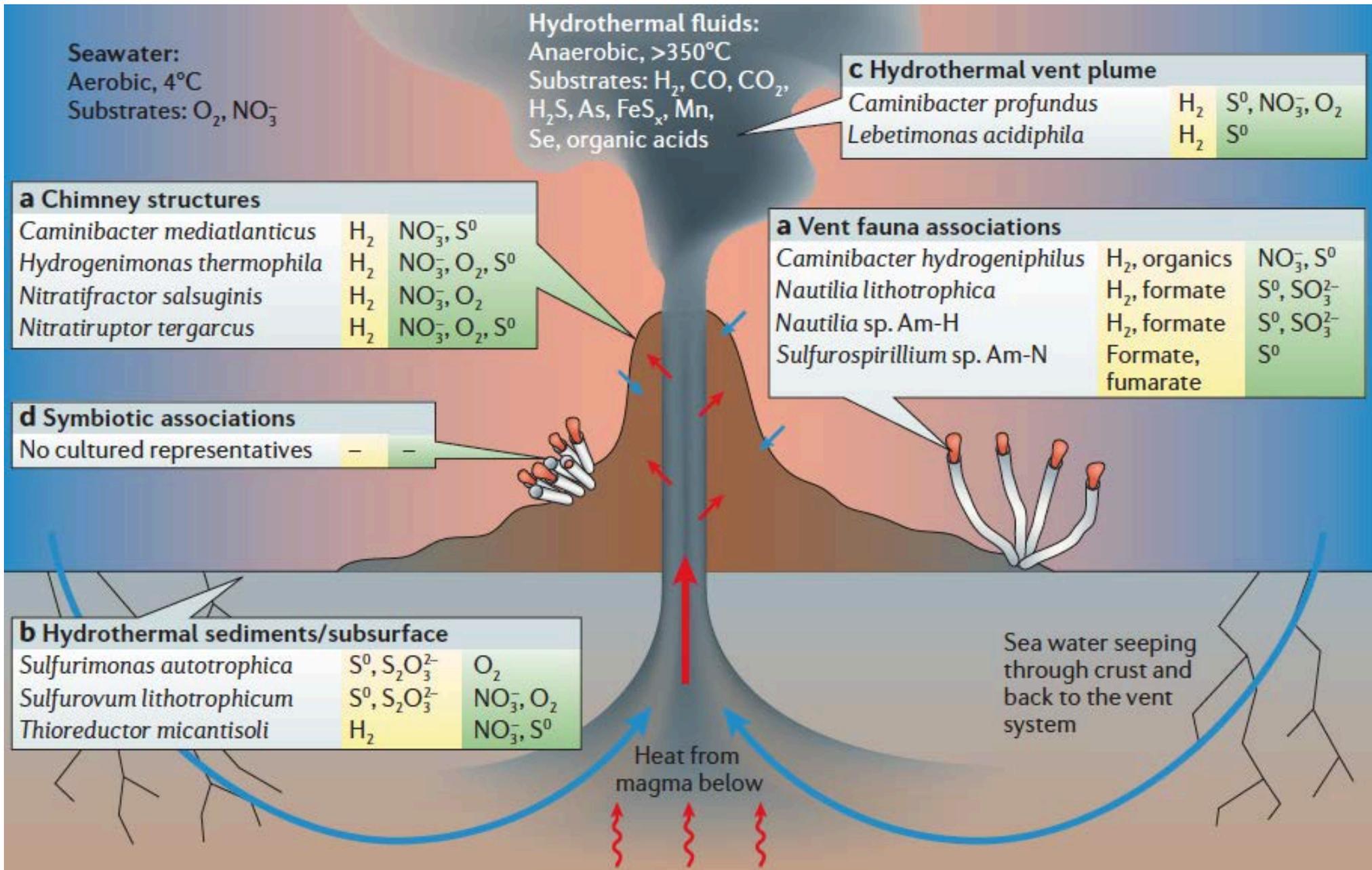
Ying Zhang*† and Stefan M. Sievert



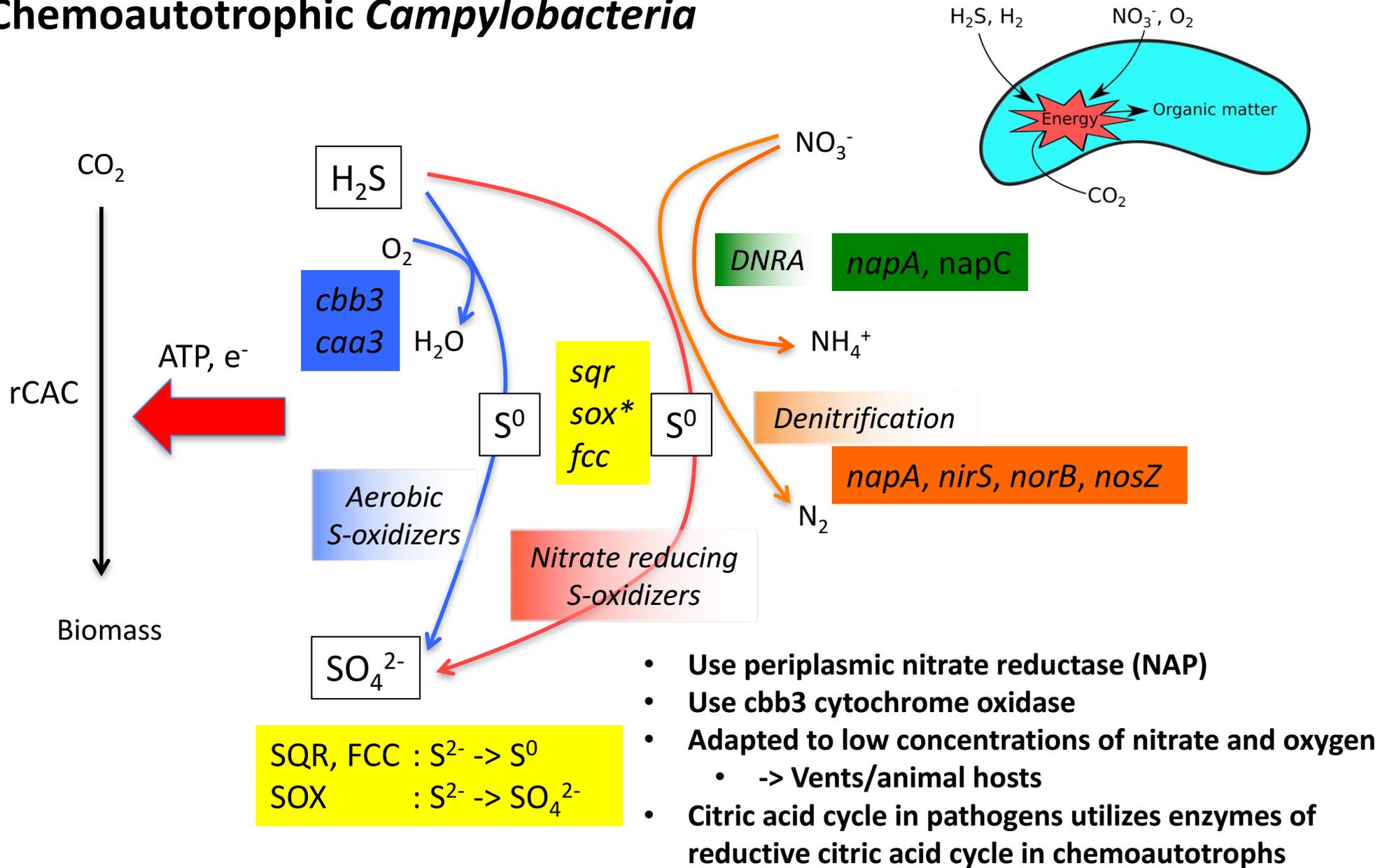
- Genome comparison of
 - 3 species of same genus
 - 3 species from different genera, but same habitat
- Most genes are lineage specific
- Select habitat specific genes



Campylobacteria inhabit a variety of niches at deep-sea hydrothermal vents



Oxidation of Sulfide Coupled to Reduction of O_2 and NO_3^- in Chemoautotrophic *Campylobacteria*

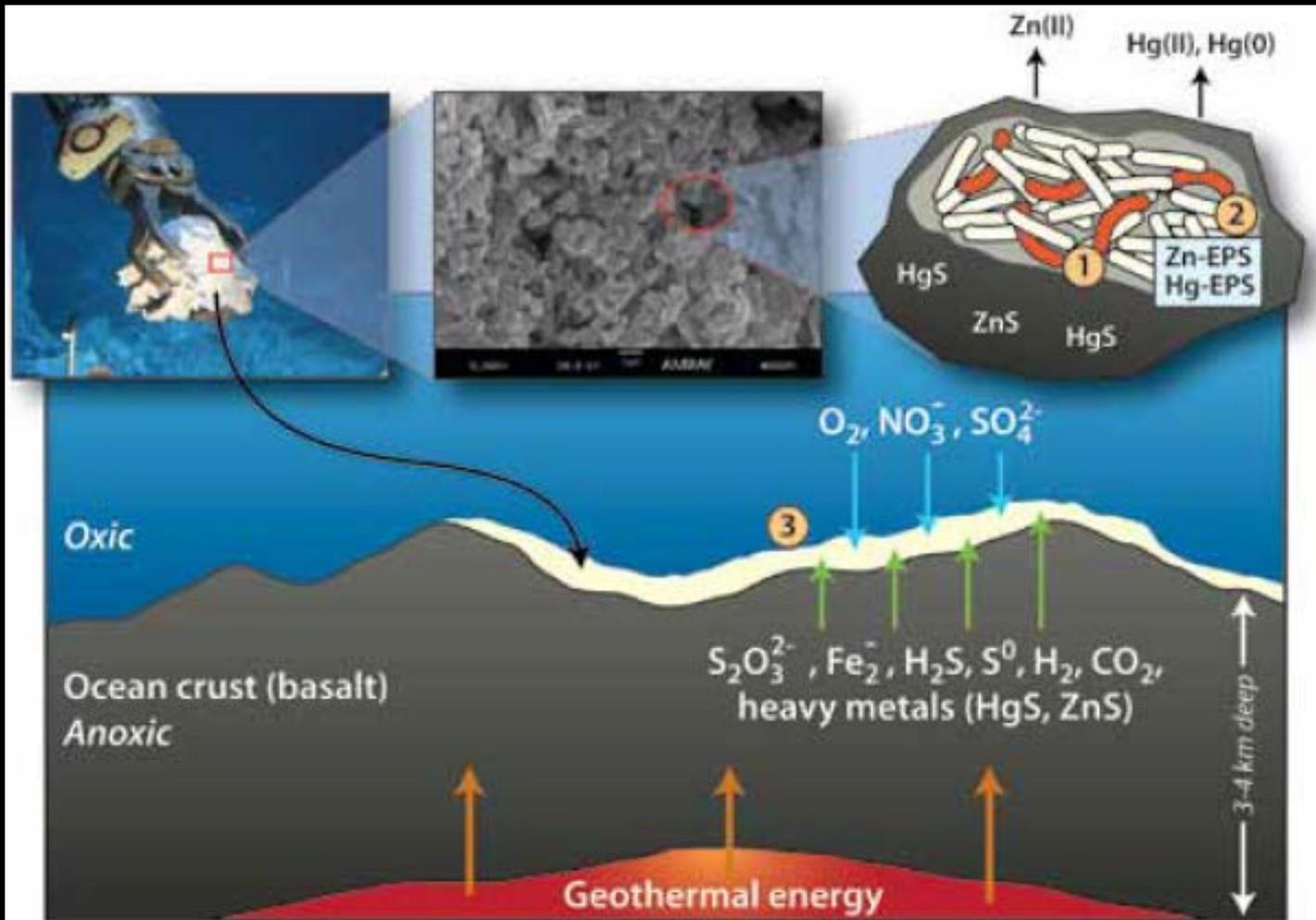


*soxABCDXYZ



Colonization of animate and inanimate surfaces at deep-sea vents

- Facilitation of larval settlement
- Trophic interactions
- Symbiotic associations

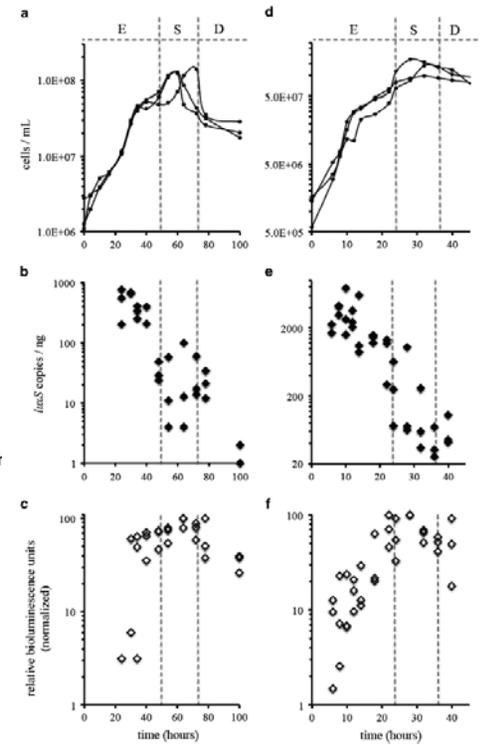
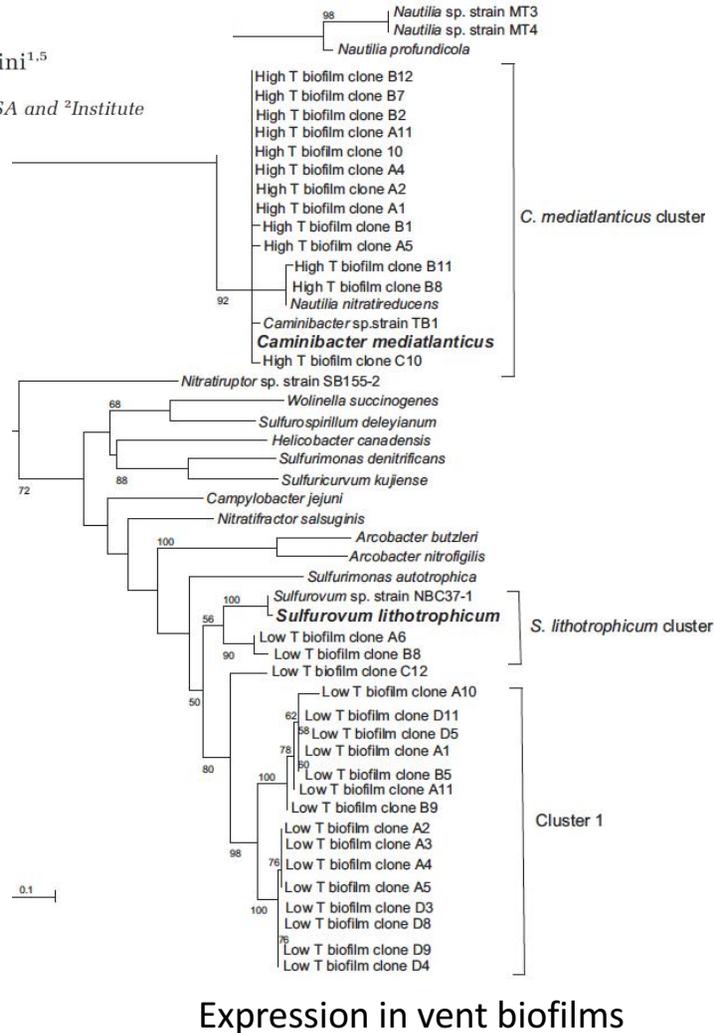
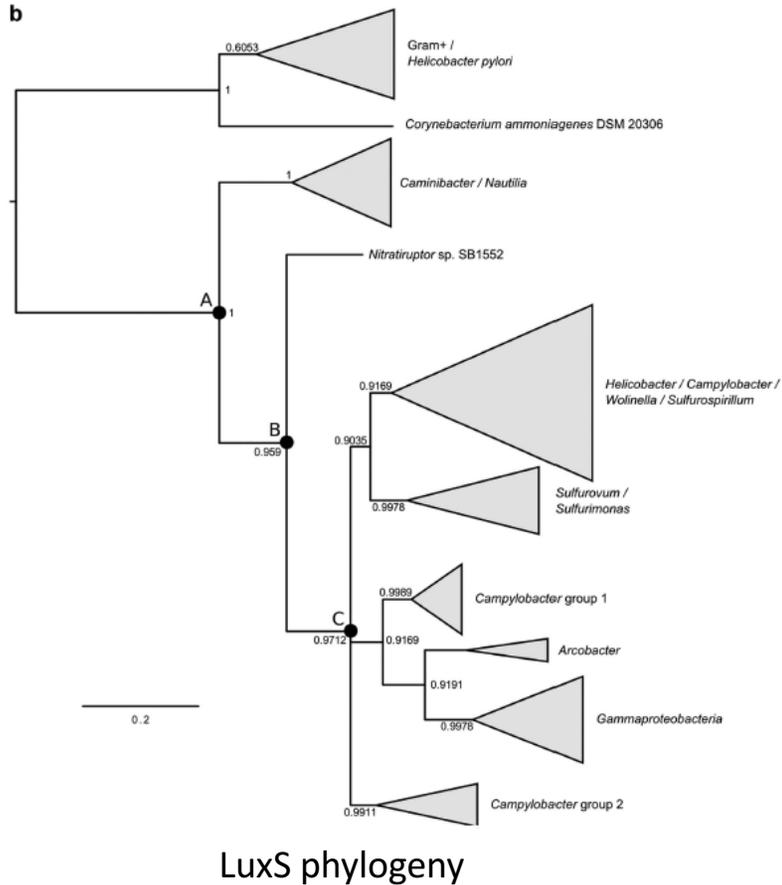


ORIGINAL ARTICLE

From deep-sea volcanoes to human pathogens:
 a conserved quorum-sensing signal in
Epsilonproteobacteria

Ileana Pérez-Rodríguez^{1,2,3}, Marie Bolognini^{1,2}, Jessica Ricci^{1,2,4}, Elisabetta Bini^{1,5}
 and Costantino Vetriani^{1,2}

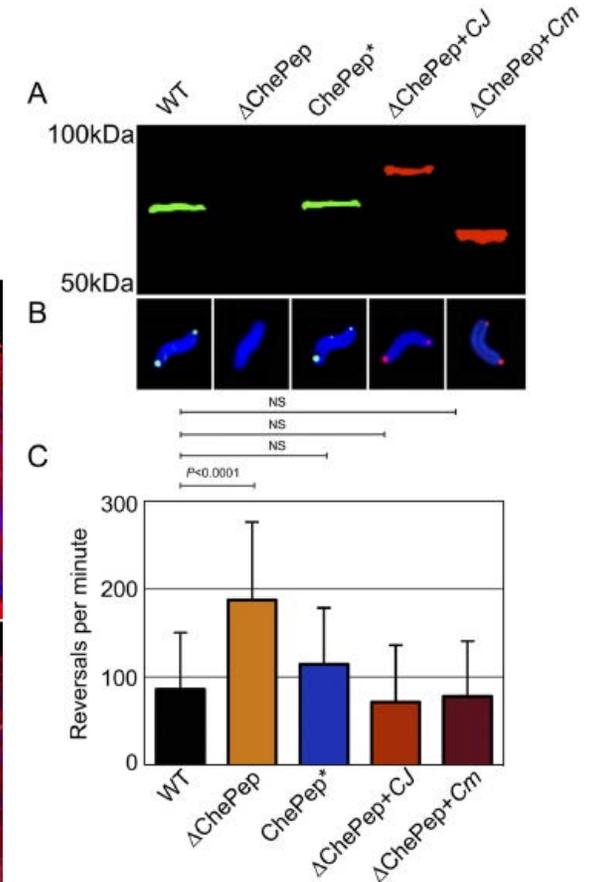
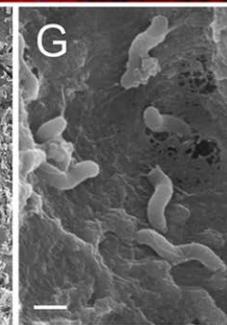
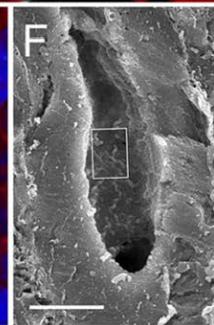
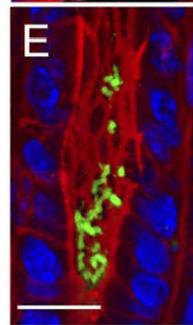
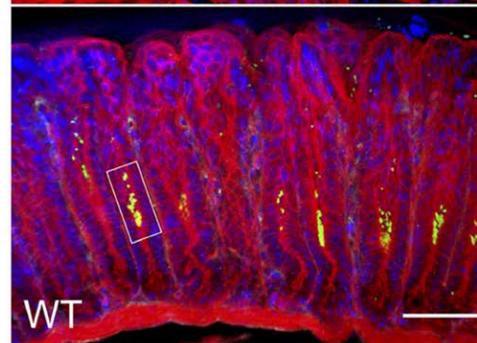
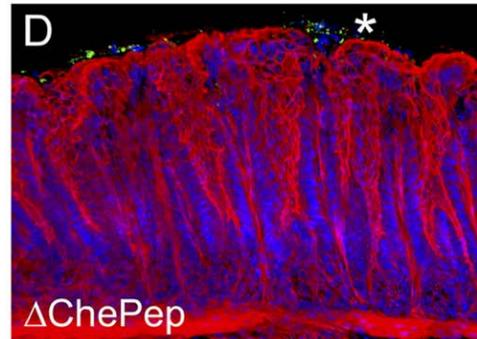
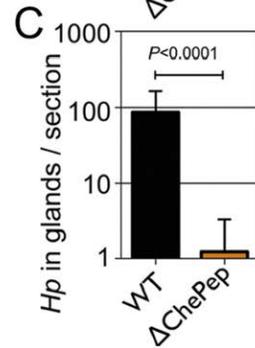
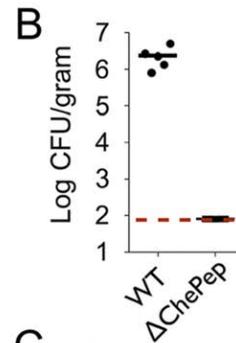
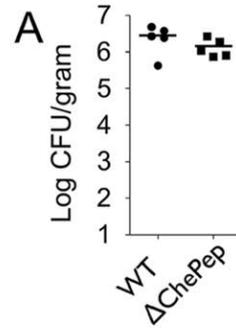
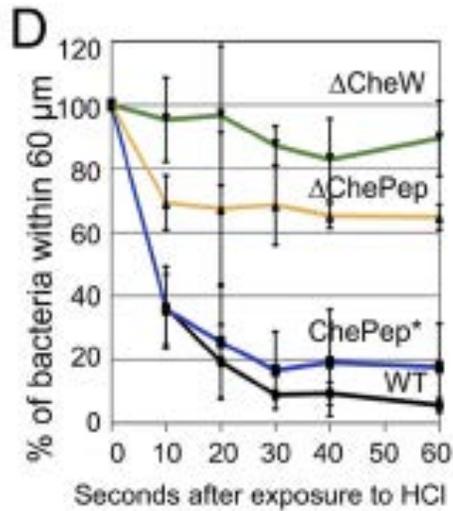
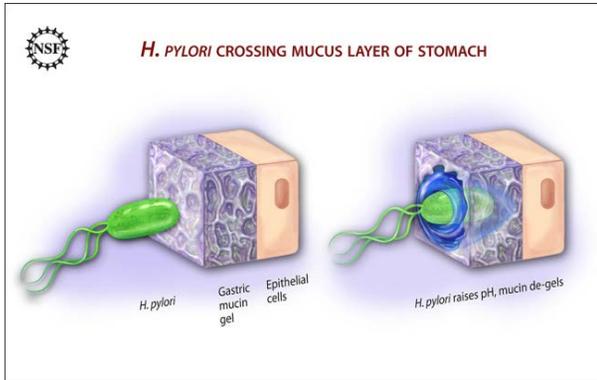
¹Department of Biochemistry and Microbiology, Rutgers University, New Brunswick, NJ, USA and ²Institute of Marine and Coastal Sciences, Rutgers University, New Brunswick, NJ, USA



- Use LuxS/AI-2 as quorum sensing mechanism
- Expressed *in situ* in vent biofilms
- Stability under harsh conditions?

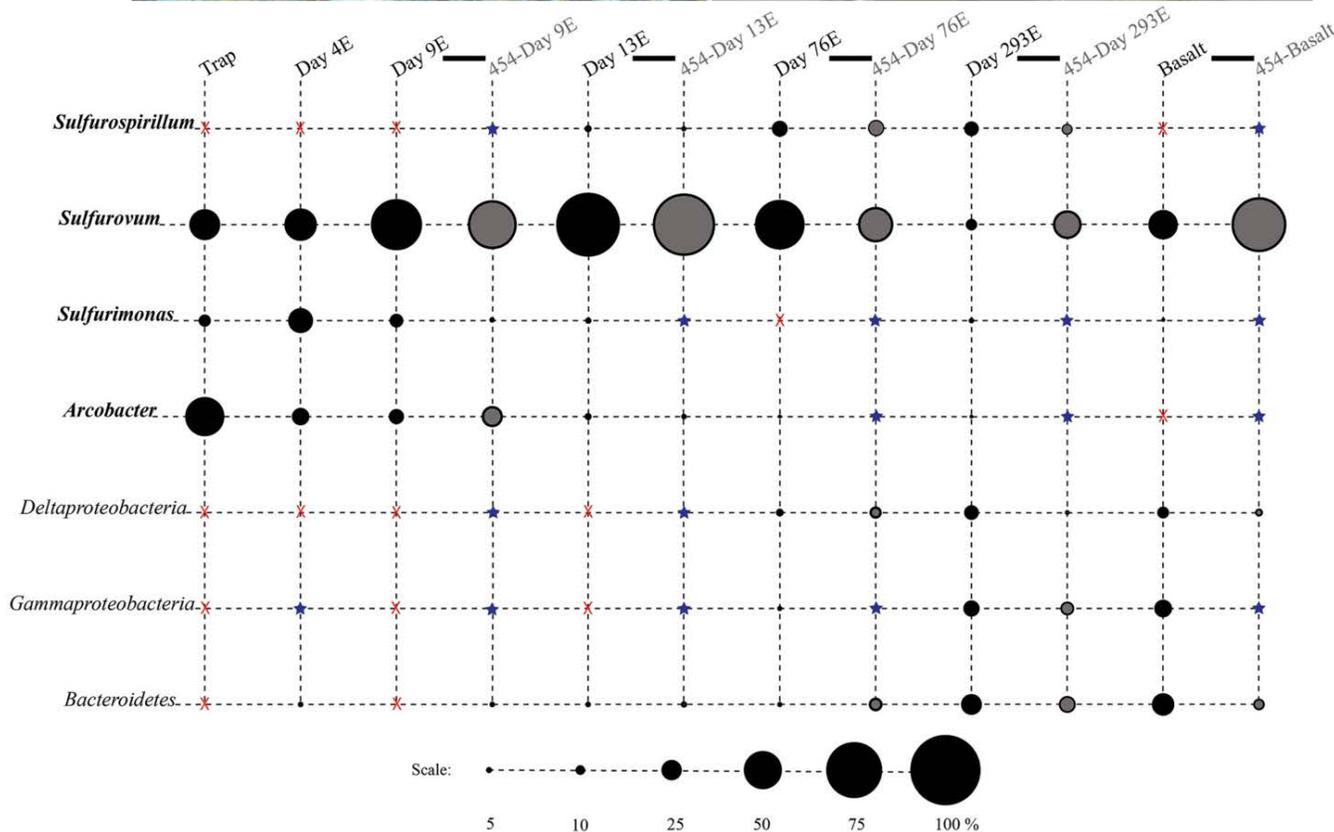
ChePep Controls *Helicobacter pylori* Infection of the Gastric Glands and Chemotaxis in the *Epsilon*proteobacteria

Michael R. Howitt,^a Josephine Y. Lee,^a Paphavee Lertsethtakarn,^b Roger Vogelmann,^{a*} Lydia-Marie Joubert,^c Karen M. Ottemann,^b and Manuel R. Amleiva^{a,d}



Colonization of gastric glands

Microbial succession on basalt rock: Foundation for Ecosystem



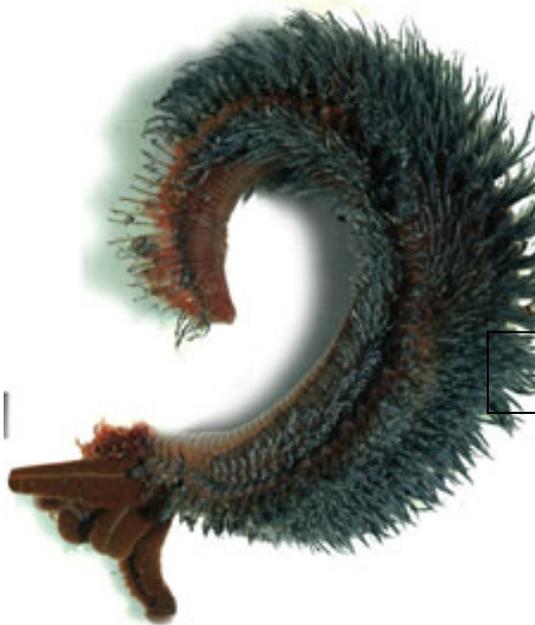
biofilm radula scrapping

Bacterial diversity and successional patterns during biofilm formation on freshly exposed basalt surfaces at diffuse-flow deep-sea vents

Lara K. Gulmann¹, Stacey E. Beaulieu¹, Timothy M. Shank¹, Kang Ding², William E. Seyfried² and Stefan M. Sievert^{1*}

Gulmann et al, 2015

Cooperative Associations: *Alvinella pompejana* and *Campylobacteria* episymbionts



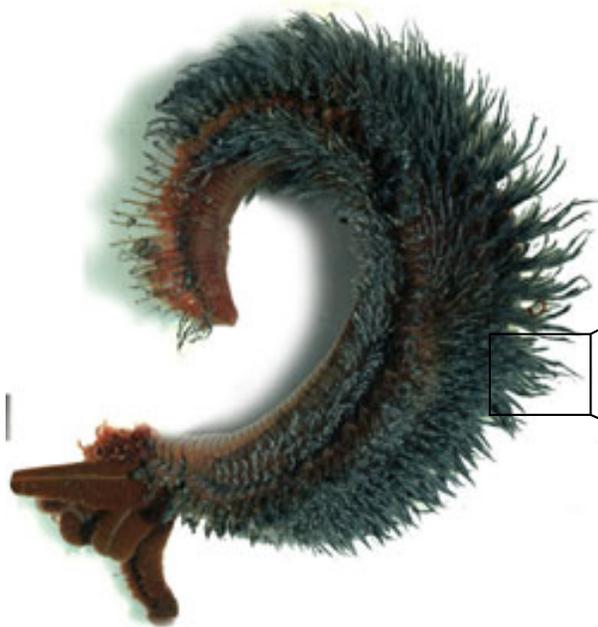
Alvinella pompejana



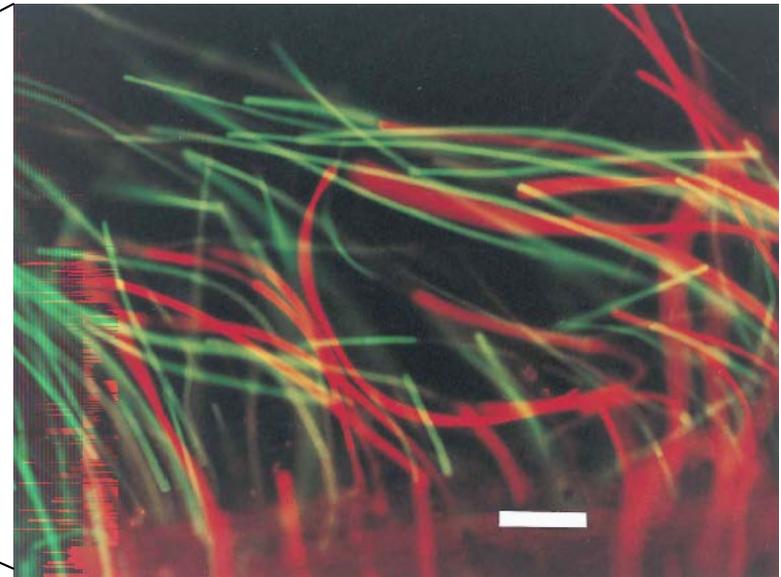
- Worms colonize outside of black-smoker chimneys
- Most eurythermal animal known, can tolerate 60°C
- Colonized by highly specific assemblage of *Campylobacteria*

Cary et al, 1997

Cooperative Associations: *Alvinella pompejana* and *Campylobacteria* episymbionts



Alvinella pompejana

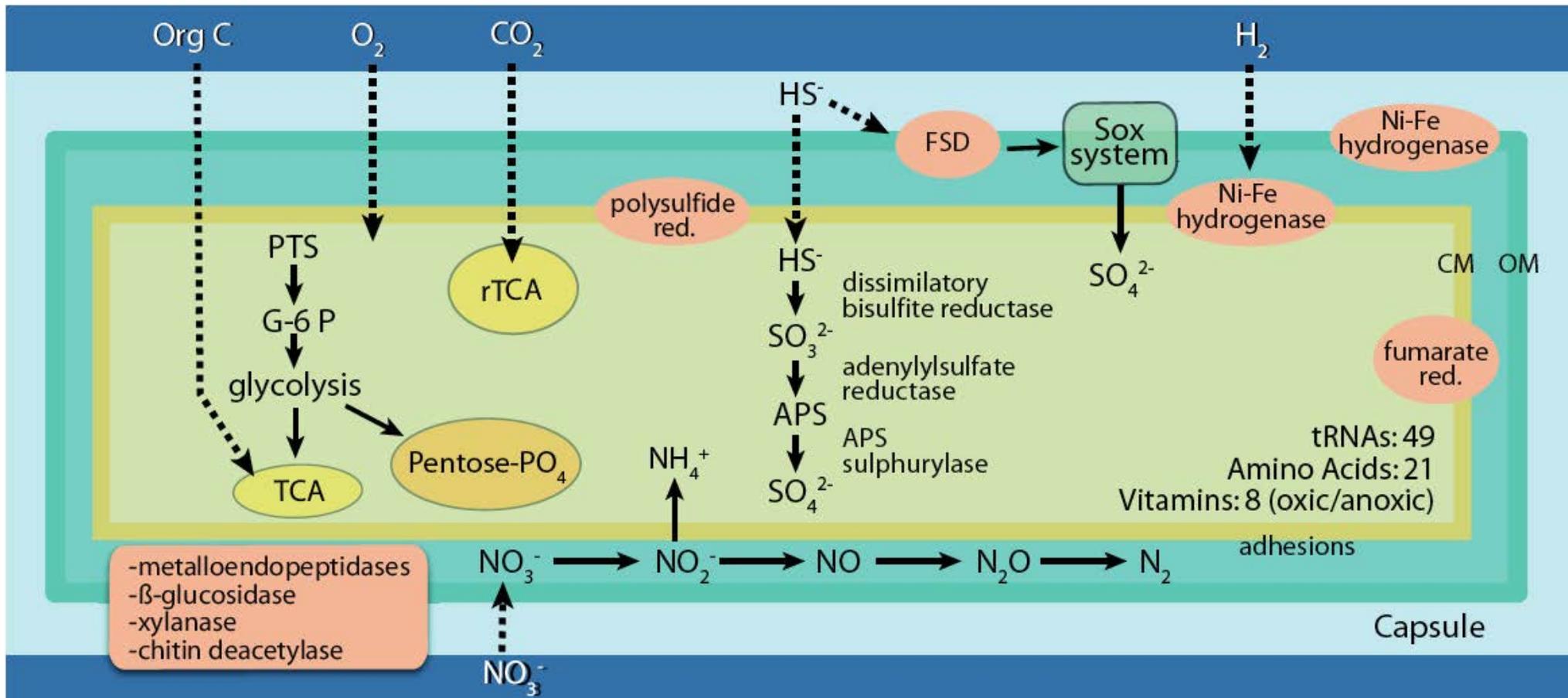


FISH analysis of *Epsilonproteobacteria* attached to the hair structures of *Alvinella pompejana*

- Worms colonize outside of black-smoker chimneys
- Most eurythermal animal known, can tolerate 60°C
- Colonized by highly specific assemblage of *Campylobacteria*

Cary et al, 1997

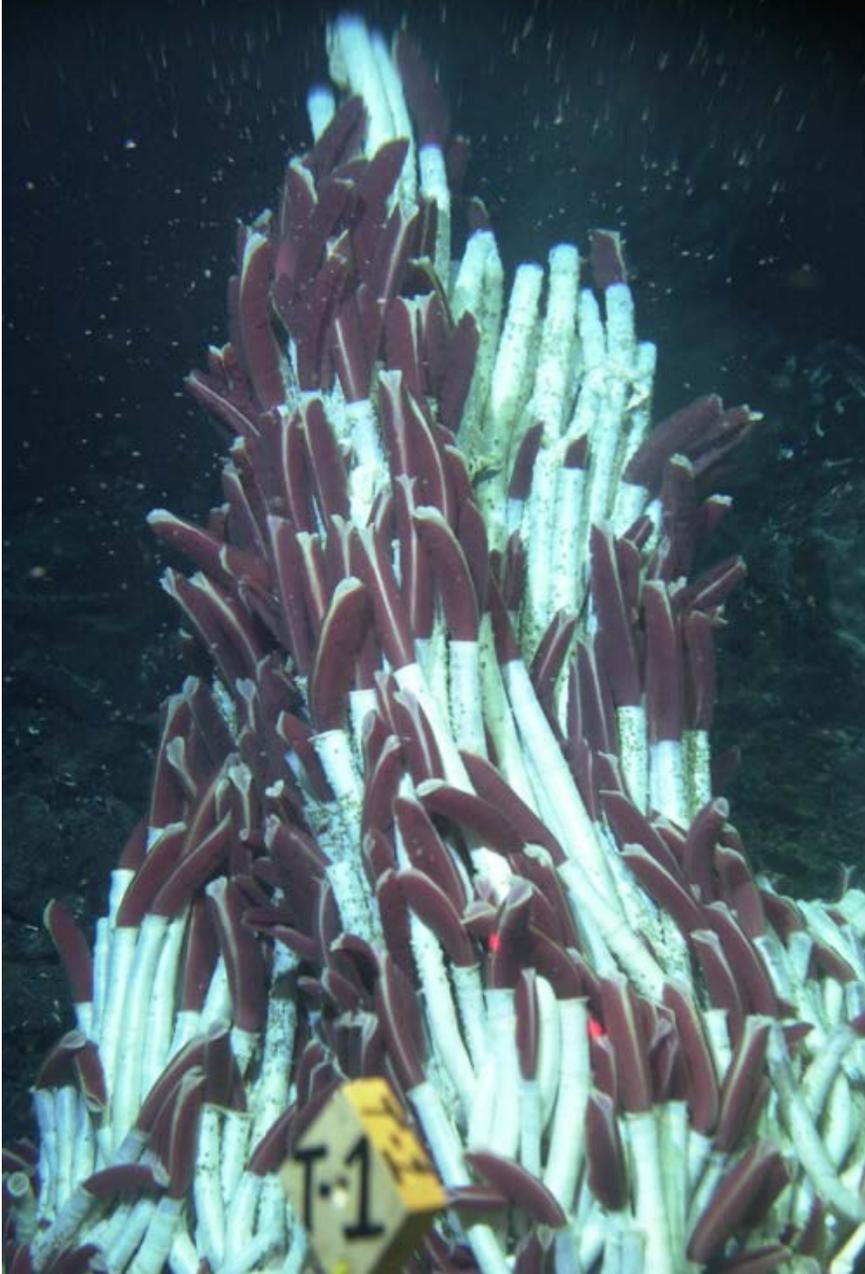
Model of **predicted** metabolic processes in the episybiont community based on annotation of the episybiont metagenome



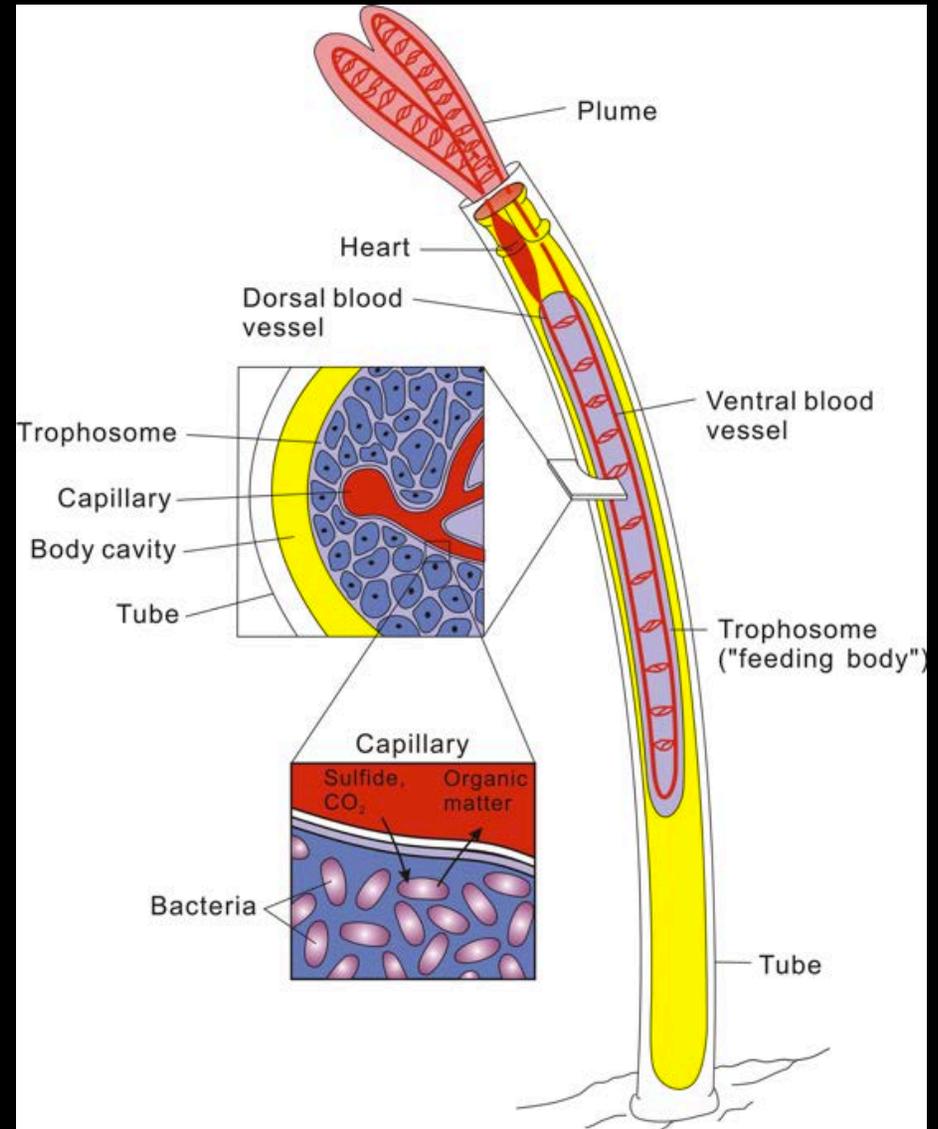
- Filamentous bacteria provide food and can provide protection from high levels of sulfide and toxic metals (arsenic, cadmium, copper).
- Highly specific association, requiring microbe-host recognition

Grzymski et al., 2008. Proc. Natl. Acad. Sci. USA 105:17516-17521

Riftia pachyptila: A unique symbiosis



Plume: Takes up sulfide, oxygen, and carbon dioxide to deliver to symbiotic chemosynthetic bacteria which produce sugars to feed the worm

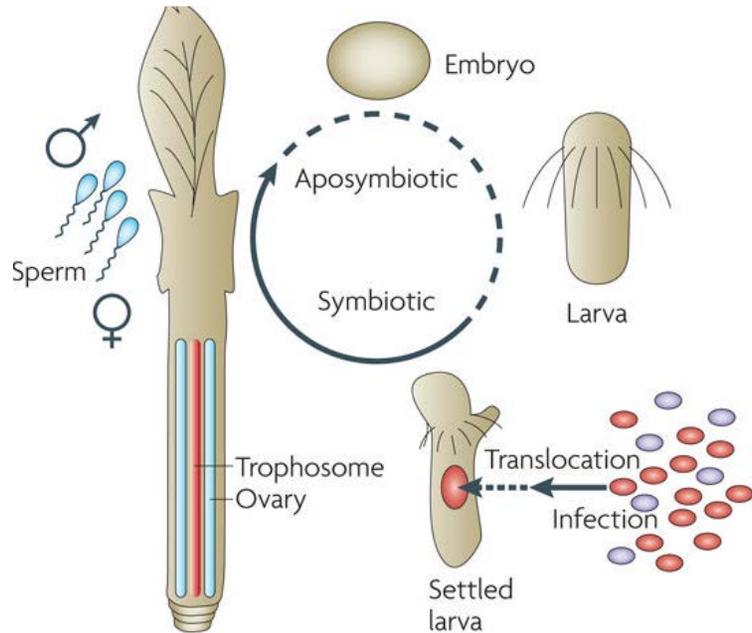


Tube: Provides protection

Trophosome: Houses symbiotic bacteria

Horizontal endosymbiont transmission in hydrothermal vent tubeworms

Andrea D. Nussbaumer¹, Charles R. Fisher² & Monika Bright¹



- Symbiont is newly acquired from environment w/ each generation
- Symbiont acquisition reminiscent of infection
- Symbionts penetrate skin of settled tubeworms
- 'Infected' tissue develops into trophosome

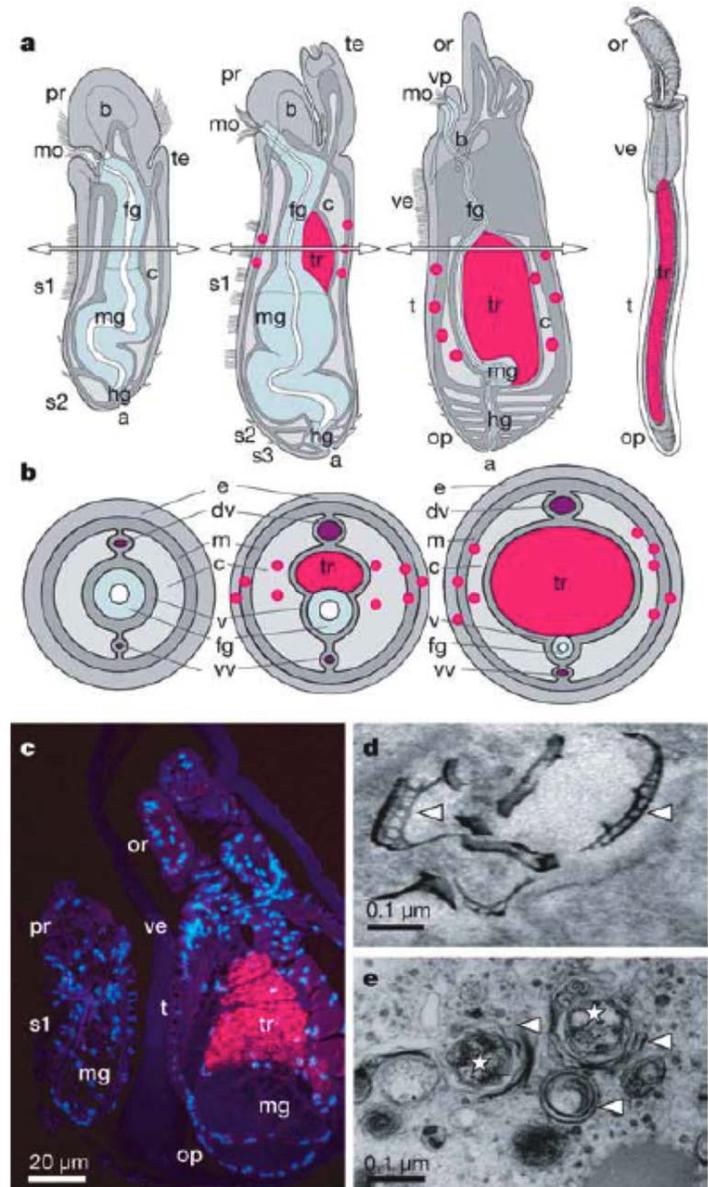


Figure 2 | Symbiont acquisition and early development of recently settled vestimentiferans. Schematic sagittal drawings of animals reconstructed

Activity and Productivity of the Subseafloor Biosphere

Submarine Thermal Springs on the Galápagos Rift

16 March 1979, Volume 203, Number 4385

SCIENCE

John B. Corliss, Jack Dymond, Louis I. Gordon,
John M. Edmond, Richard P. von Herzen, Robert D. Ballard,
Kenneth Green, David Williams, Arnold Bainbridge,
Kathy Crane, Tjeerd H. van Andel

**First comprehensive paper on deep-sea vent
geology, chemistry, and biology based on 1977
expedition**

“This flux of bacteria from the vents must be supported by the production of a large population of bacteria living within the rock mass, lining the walls of fissures (and fractures) through which the hydrogen sulfide-laden fluids ascend.”

“and that they may significantly influence the chemistry of the system.”

“The generation time for these populations of sulfur-oxidizing bacteria in situ is not known, but, unlike phytoplankton blooms, their productivity is, presumably, continuous.”

- Corliss et al, 1979, *Science*, 1979

**Deep-Sea Primary Production at the
Galápagos Hydrothermal Vents**

SCIENCE, VOL. 207, 21 MARCH 1980

D. M. KARL
*Department of Oceanography,
University of Hawaii, Honolulu 96822*
C. O. WIRSEN, H. W. JANNASCH
*Woods Hole Oceanographic Institution,
Woods Hole, Massachusetts 02543*

One of first studies to measure primary production at deep-sea vents

In 1980, after follow up study

“In view of the complexity of the entire vent system and the limited amount of sampling possible, a **useful quantification of deep-sea primary production is quite out of reach at this time.**”

-Karl, Wirsen and Jannasch, *Science*, 1980

Today

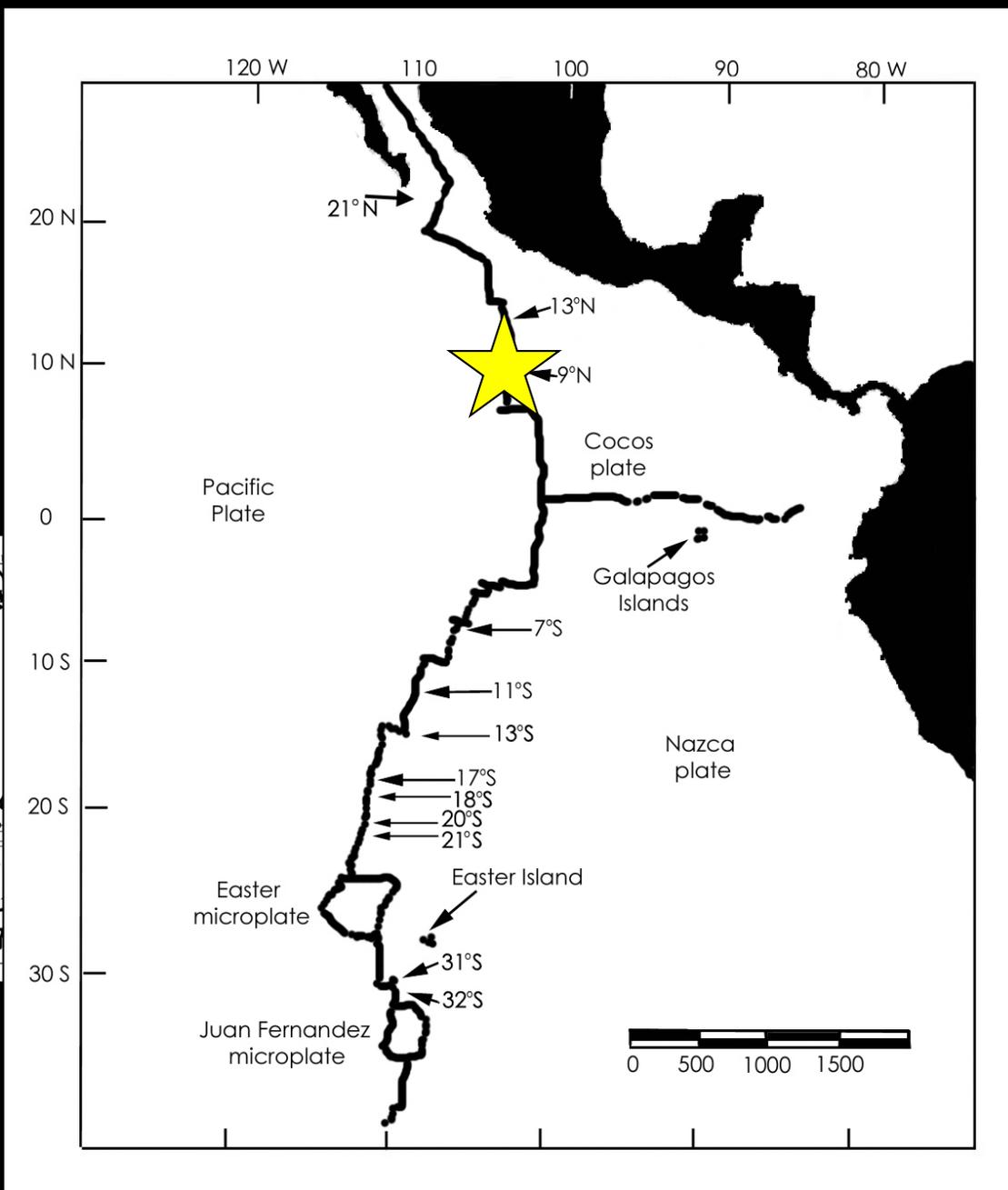
“In view of the complexity of the entire vent system and the limited amount of sampling possible, **a useful quantification of deep-sea primary production is quite out of reach at this time.**

-Karl, Wirsen and Jannasch, *Science*, 1980

This goal is now within reach

'CRAB SPA' AT 9°50'N ON EAST PACIFIC RISE

A MODEL SYSTEM TO STUDY CHEMOAUTOTROPHIC PROCESSES AT DEEP-SEA VENTS



Crab Spa at Tica

Temp: ~24 (30°C in '07)

H₂S: 300 μM (1 mM in '07)

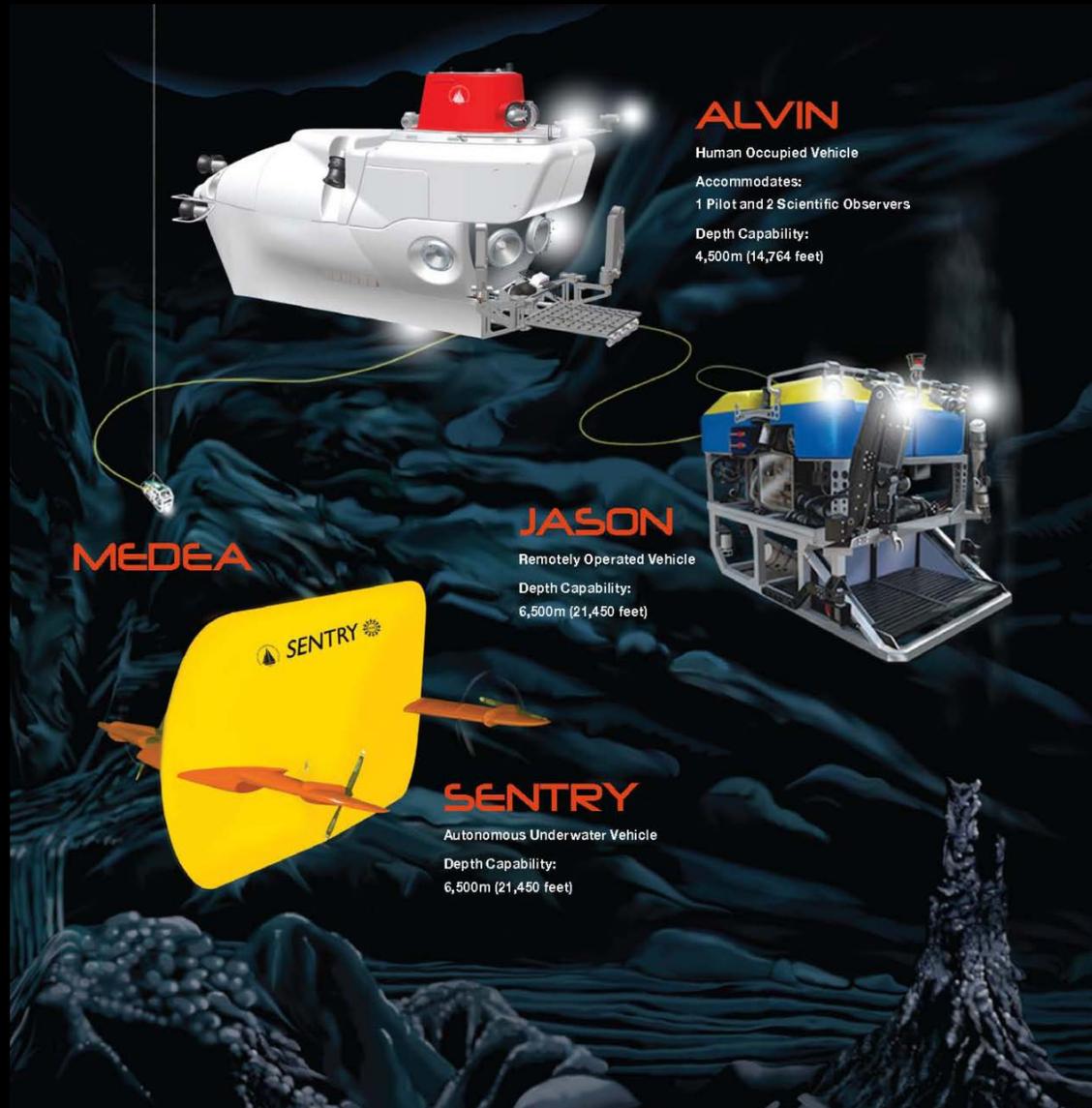
H₂: ~5 μM

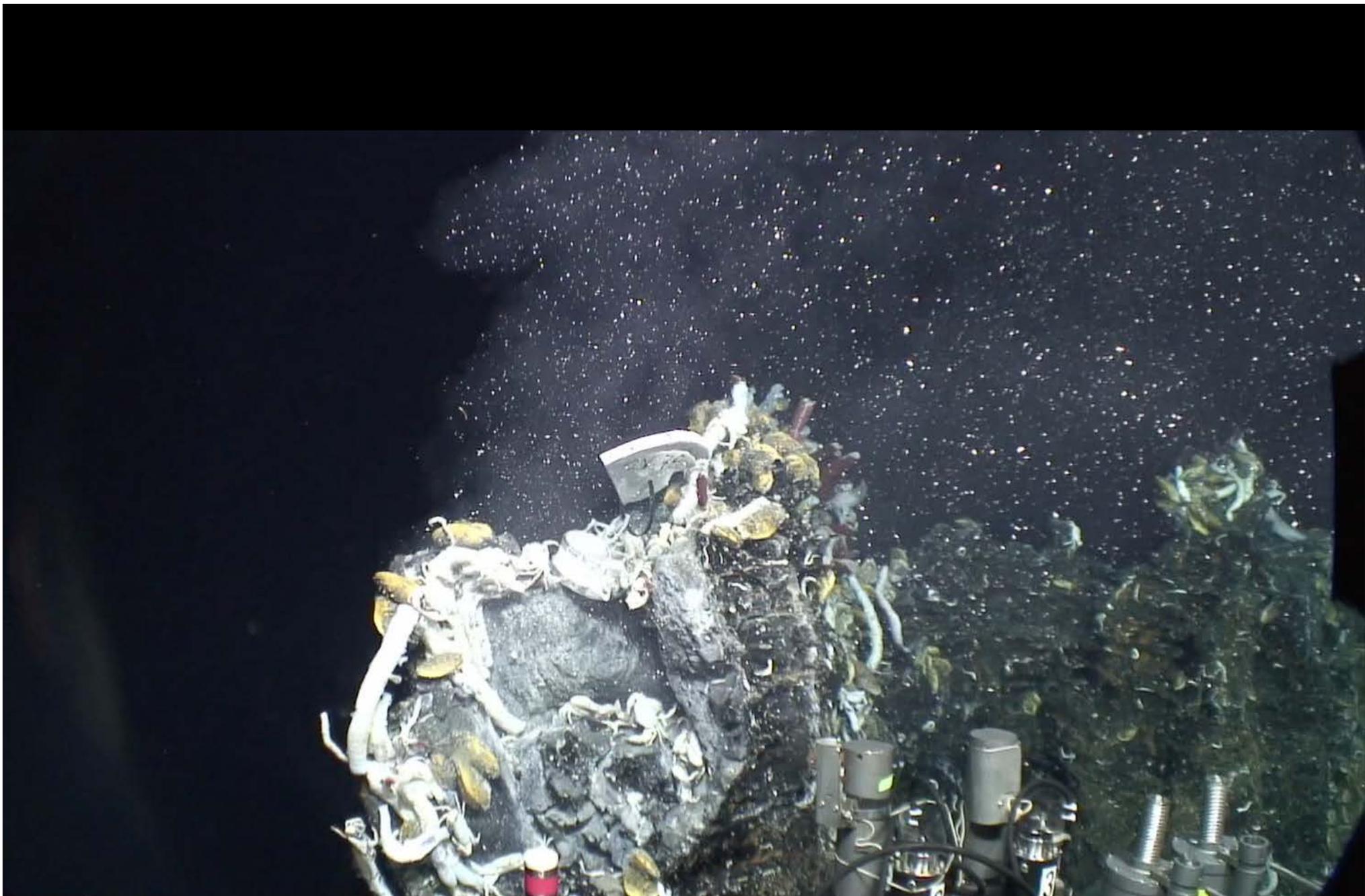
pH: 5.8

O₂: <2 μM, NO₃²⁻: <7 μM

Sampled in Jan '07, Jan '08, Oct '08,
May '10, May '12, Jan '14, Nov '14,
May '17

NATIONAL DEEP SUBMERGENCE FACILITY AT WHOI: ALVIN, JASON & SENTRY

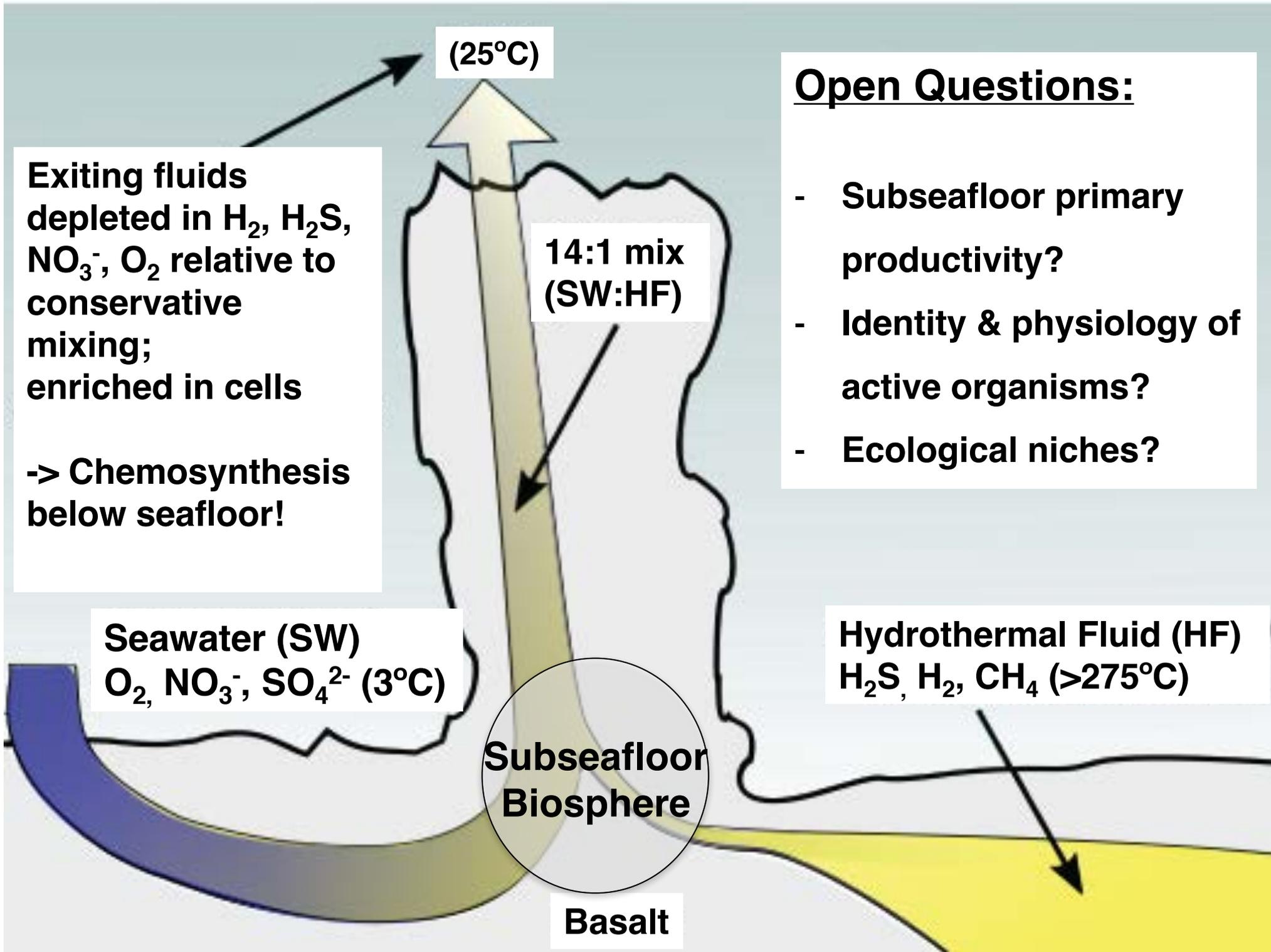




'Crab Spa'



'Snow Blower'



Exiting fluids depleted in H₂, H₂S, NO₃⁻, O₂ relative to conservative mixing; enriched in cells

-> Chemosynthesis below seafloor!

- Open Questions:**
- **Subseafloor primary productivity?**
 - **Identity & physiology of active organisms?**
 - **Ecological niches?**

**Seawater (SW)
O₂, NO₃⁻, SO₄²⁻ (3°C)**

**Hydrothermal Fluid (HF)
H₂S, H₂, CH₄ (>275°C)**

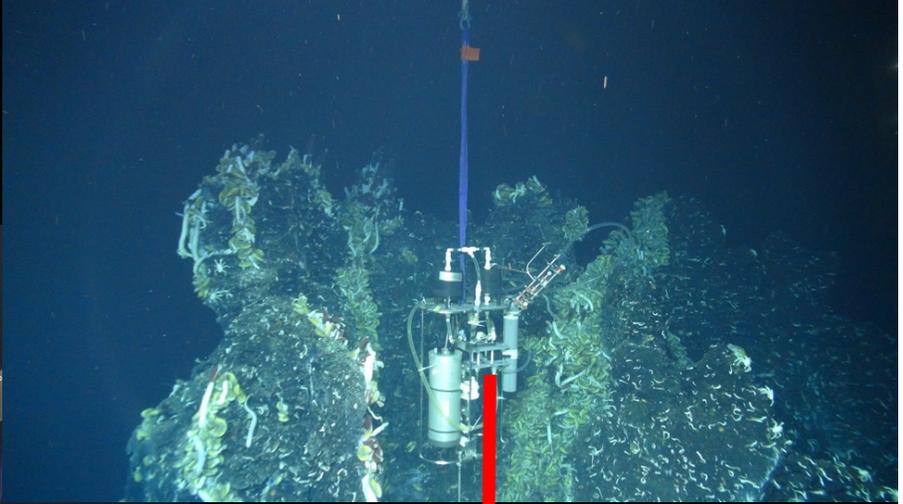
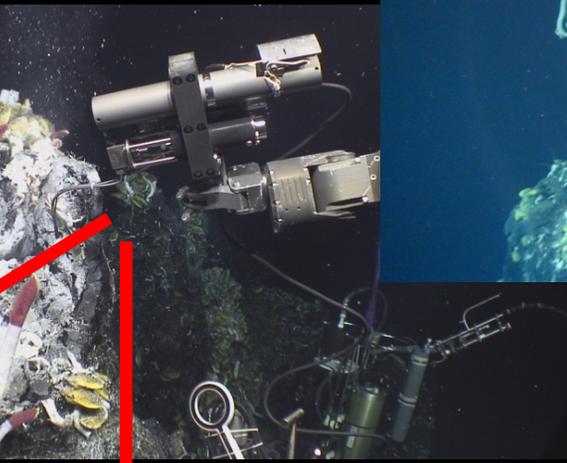
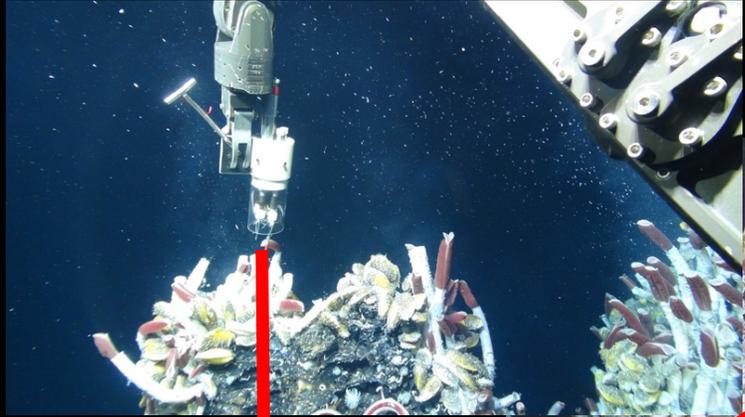
Subseafloor Biosphere

Basalt

(25°C)

**14:1 mix
(SW:HF)**

Multifaceted Approach to Study Crab Spa Microbiome



Geochemistry

1. *In situ*
2. Lab based

In situ filtration w/ LVP

1. 16S rRNA pyrotags
2. Metagenome
3. Metatranscriptome
4. **Metaproteome**
5. Lipids

Samples w/ IGT fluid samplers

1. **Shipboard incubations**
2. CARD-FISH
3. Single-cell Amplified Genomes (SAG)

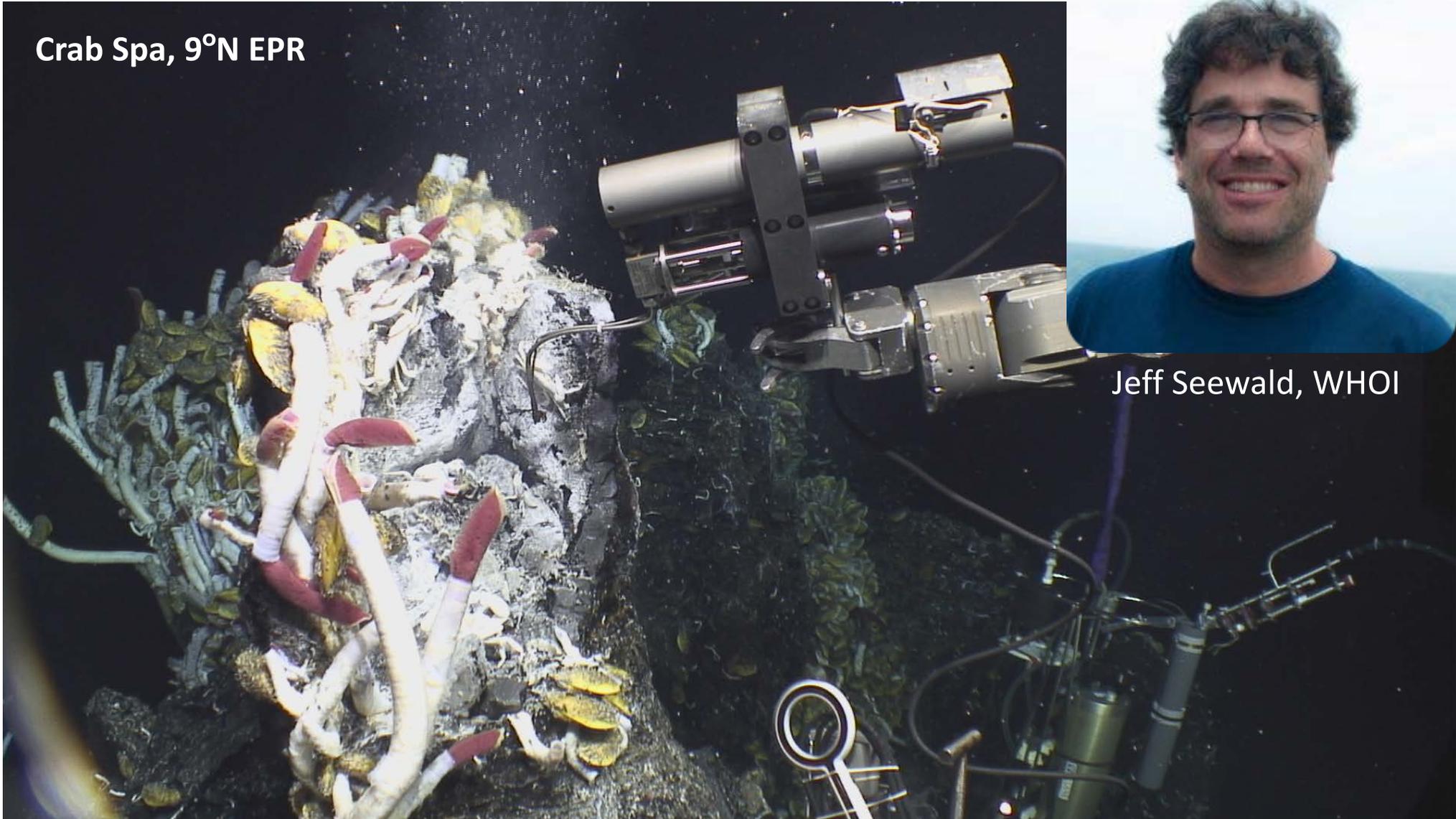


Jesse McNichol, François Thomas



Florian Götz

Crab Spa, 9°N EPR



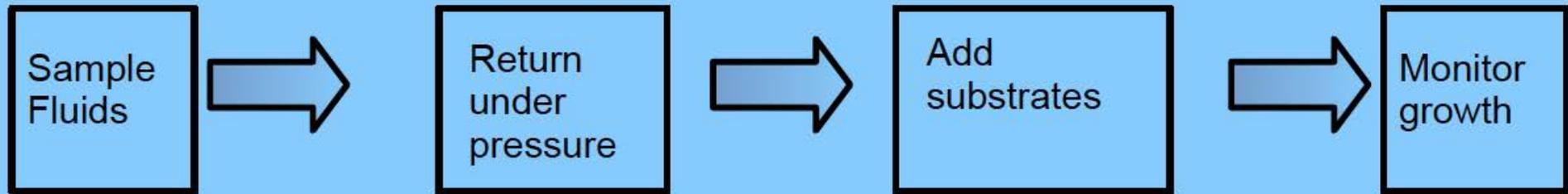
Jeff Seewald, WHOI

Isobaric Gas Tight Sampler (IGT) to collect fluids for chemical analyses and investigating microbial activity at *in-situ* pressure and temperature

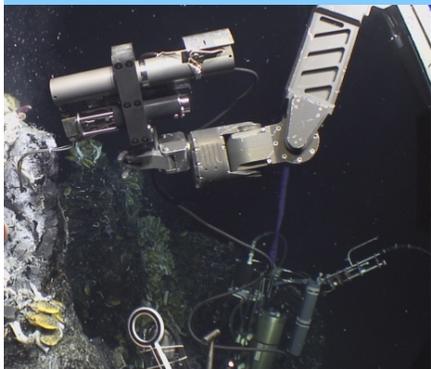
SHIPBOARD INCUBATIONS WITH HYDROTHERMAL FLUIDS UNDER SIMULATED *IN SITU* CONDITIONS



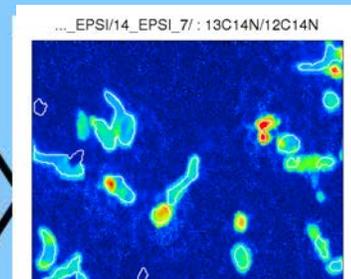
Jesse McNichol



IGT

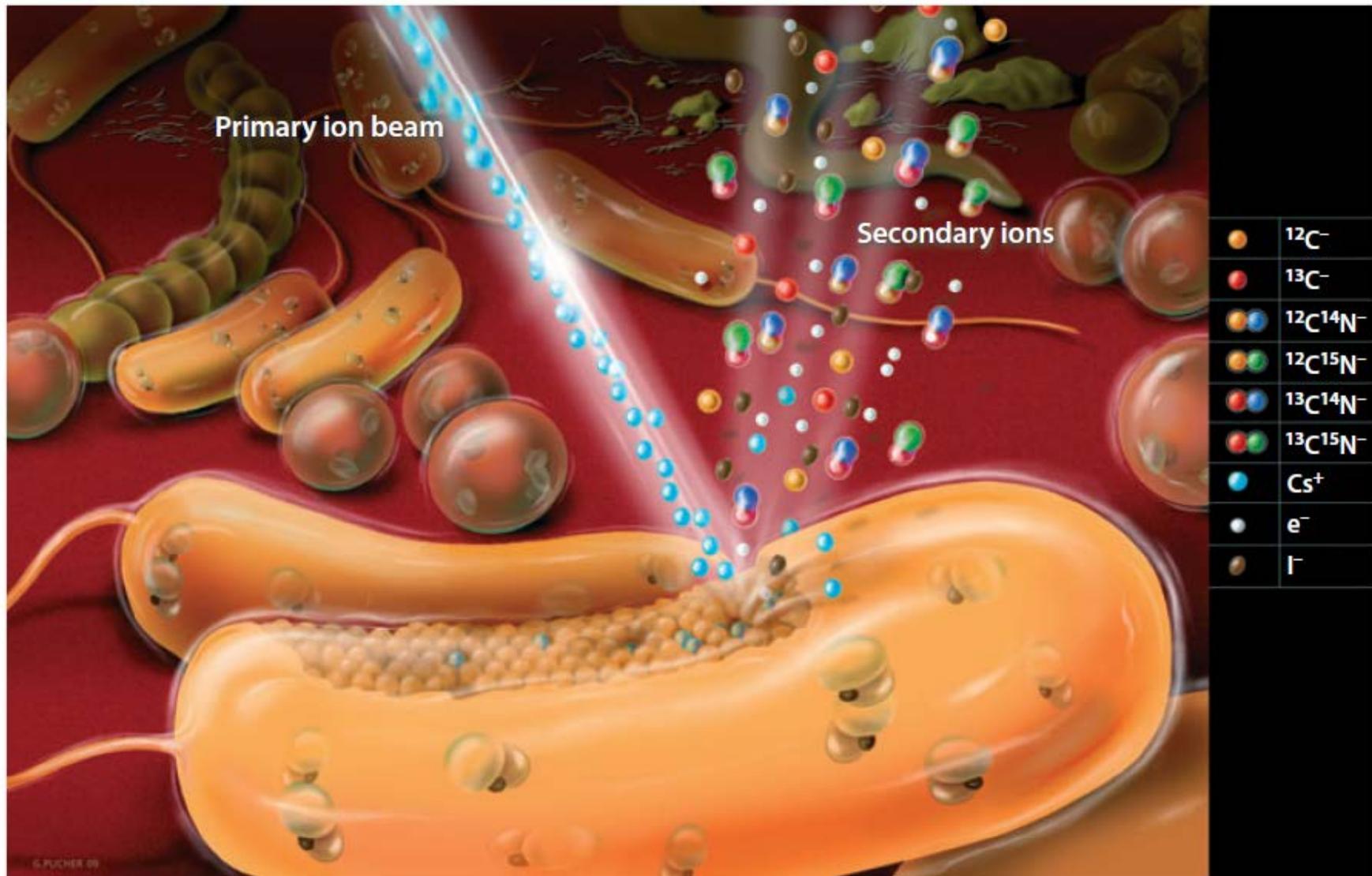


nano-SIMS



- Added ^{13}C -labeled bicarbonate to incubations conducted at *in situ* pressure, different temperatures and conditions
- Analyses at single cell level with HISH-SIMS: Nanoscale-Secondary Ion Microprobe coupled with Halogen In Situ Hybridization

Nano-SIMS is mass spectrometry on individual cells

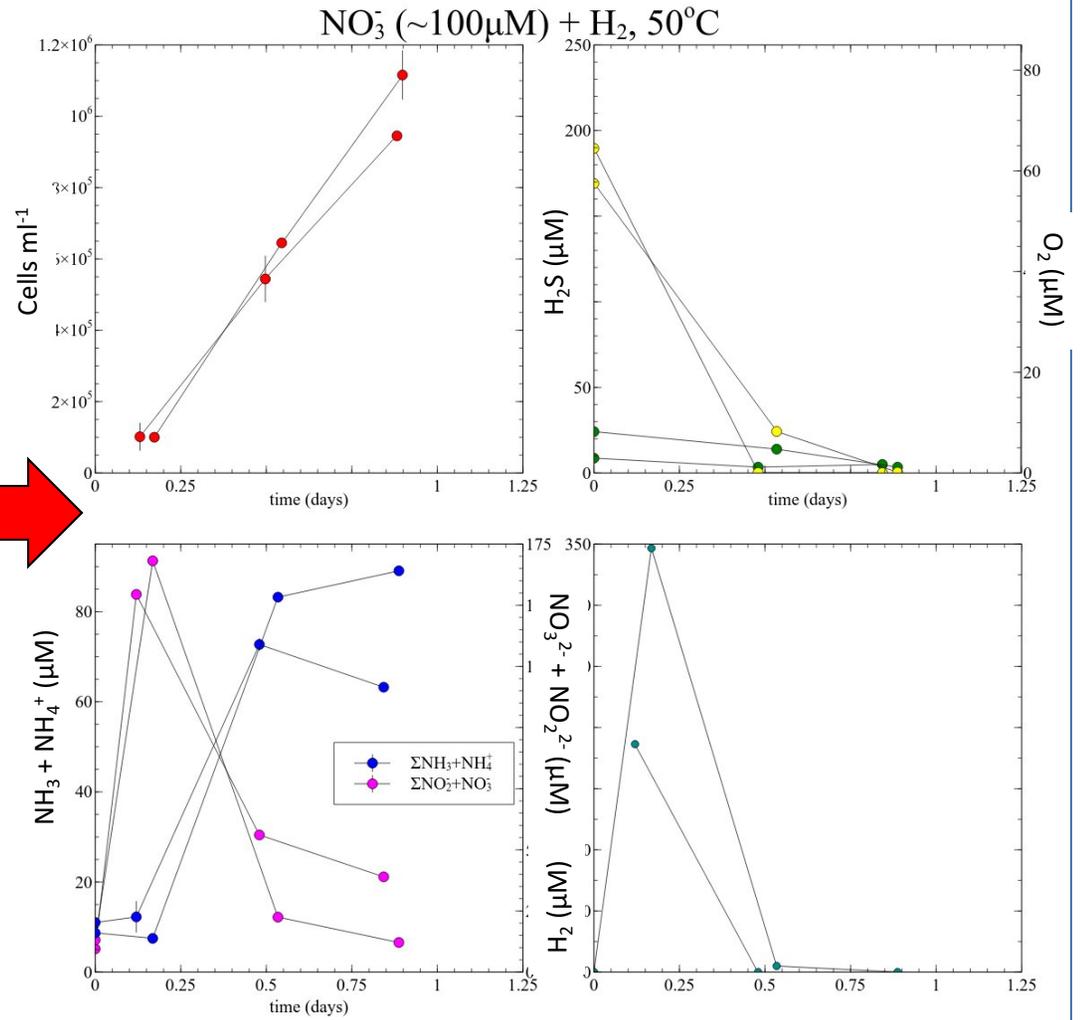


Wagner, 2009, Annual Review of Microbiology

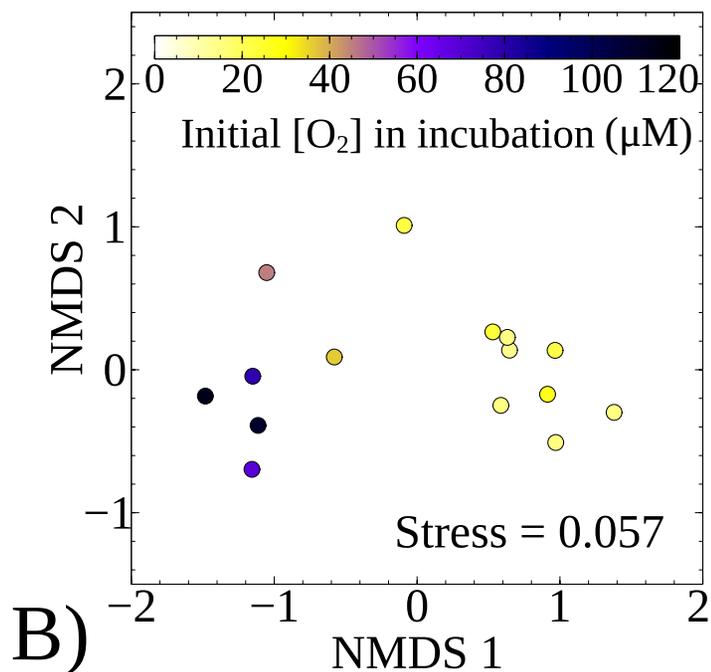
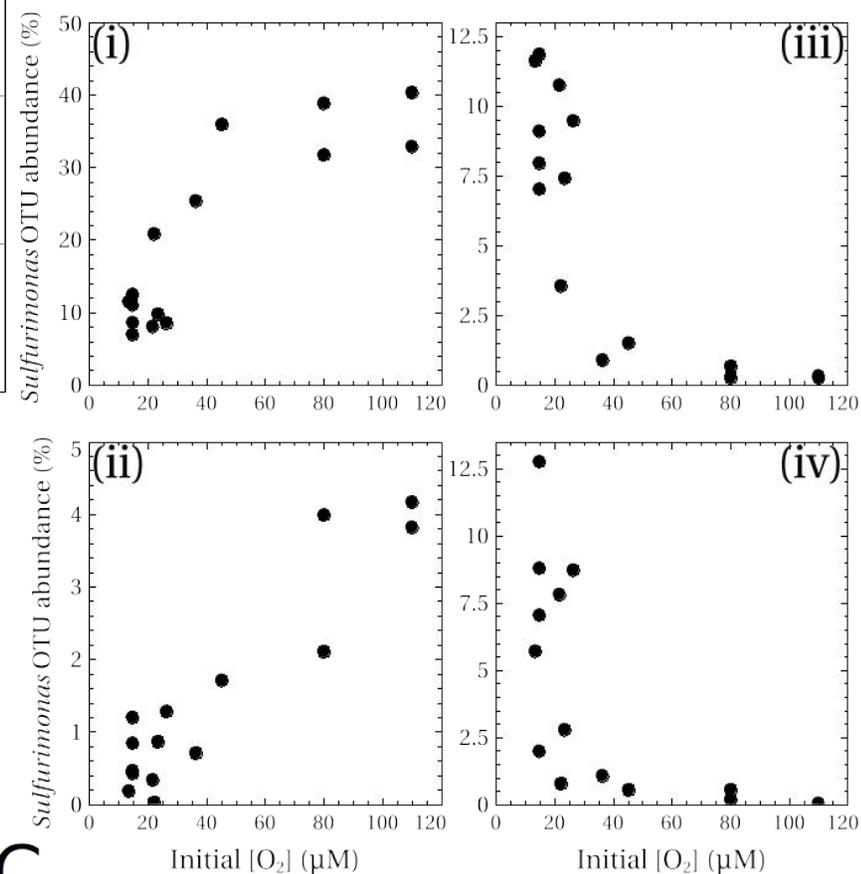
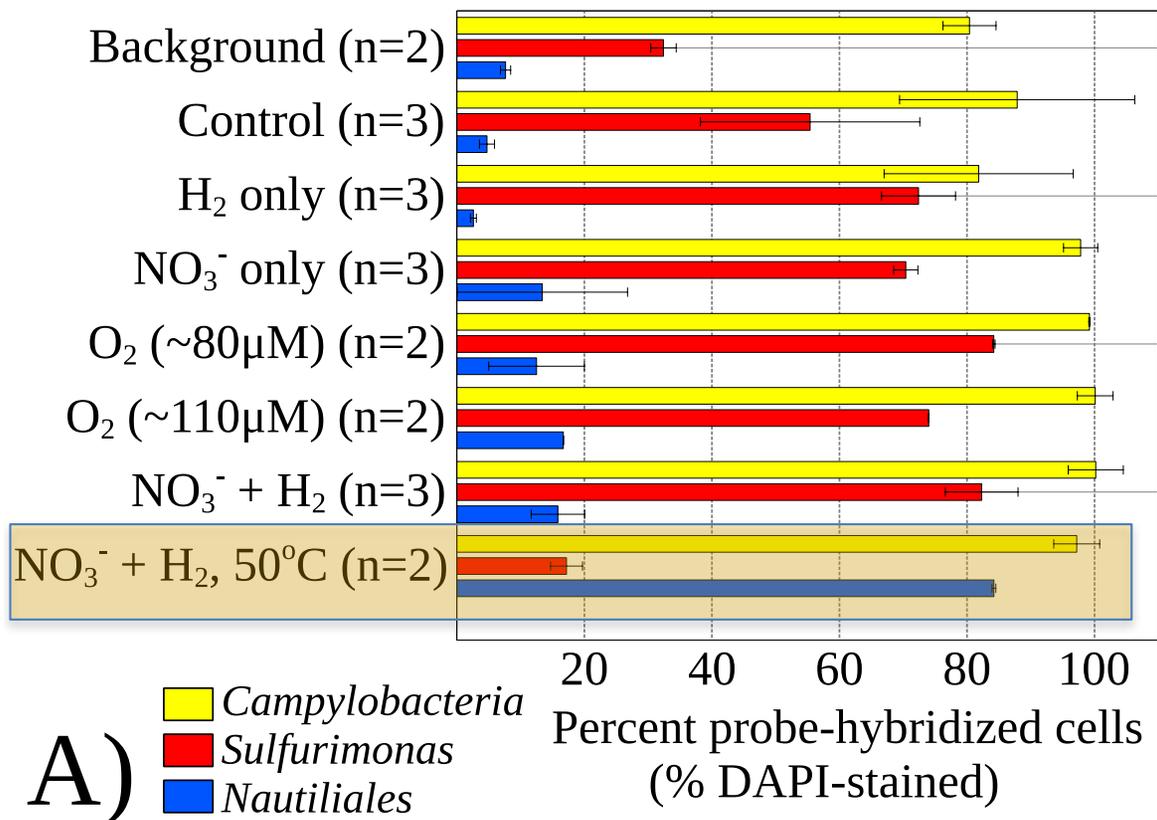
SHIPBOARD GENERATED DATA



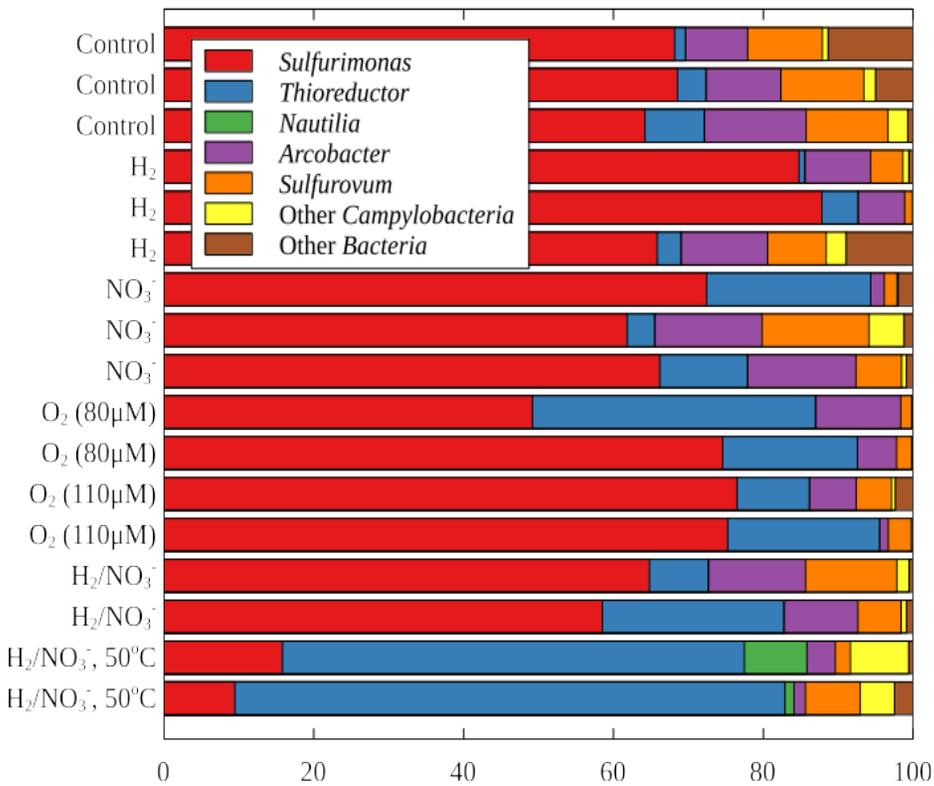
Jesse McNichol & François Thomas



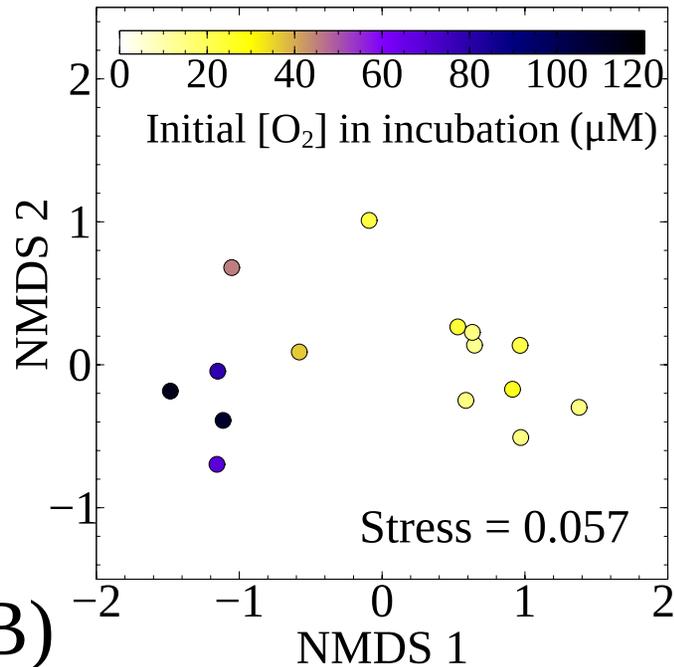
Bacterial community composition during incubations



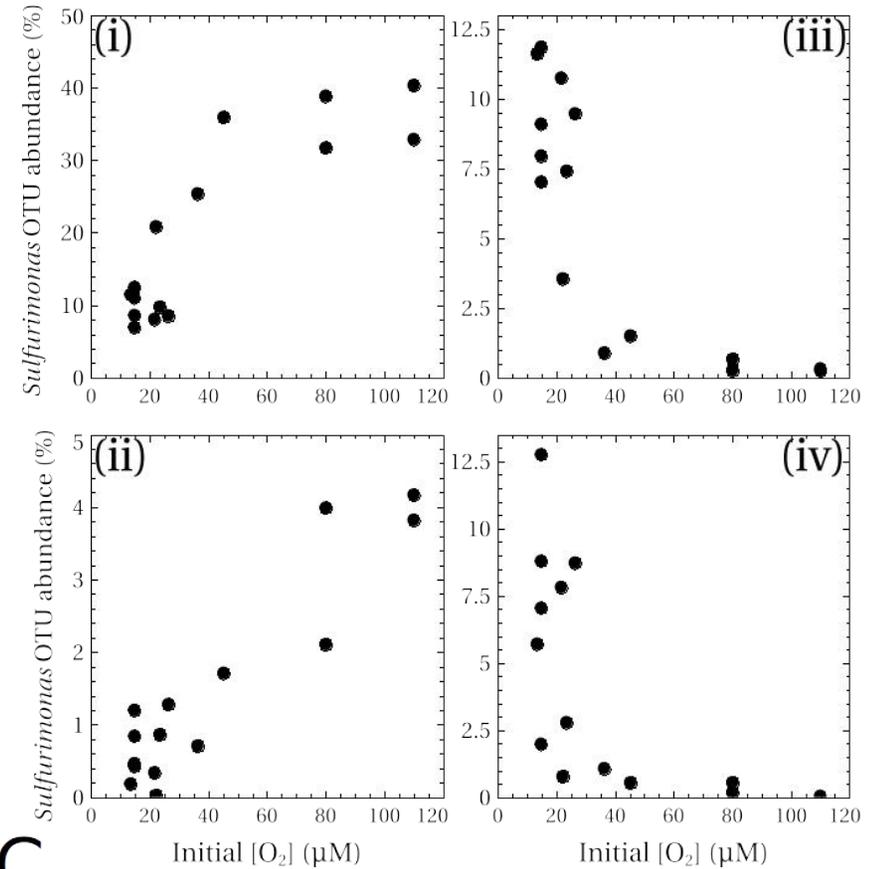
Bacterial community composition during incubations



A) Percent of 16S amplicon sequences (27Fmod/519Rmodbio)



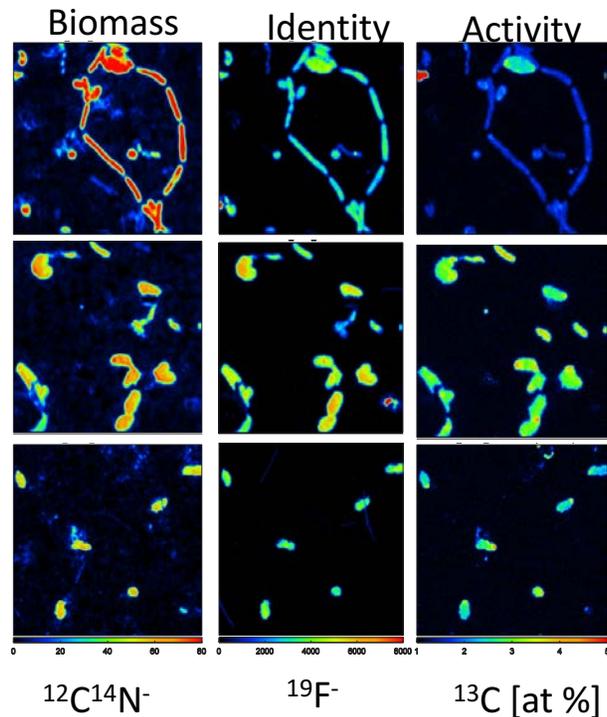
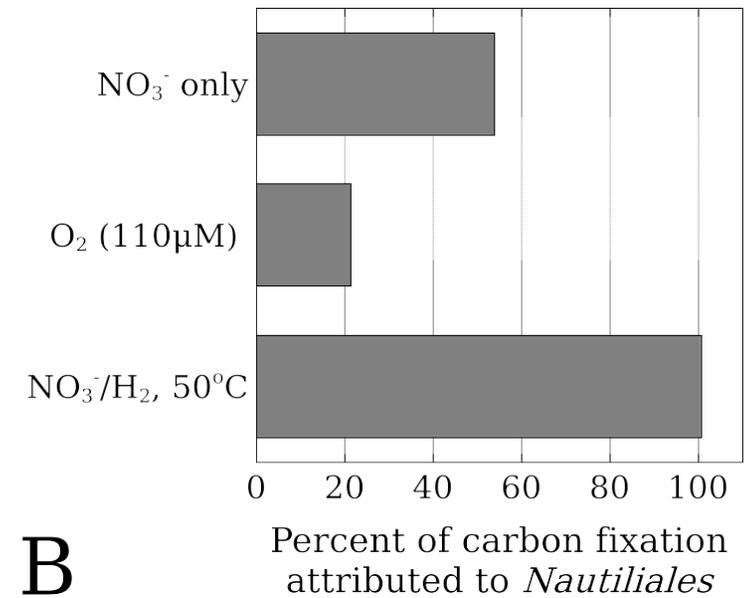
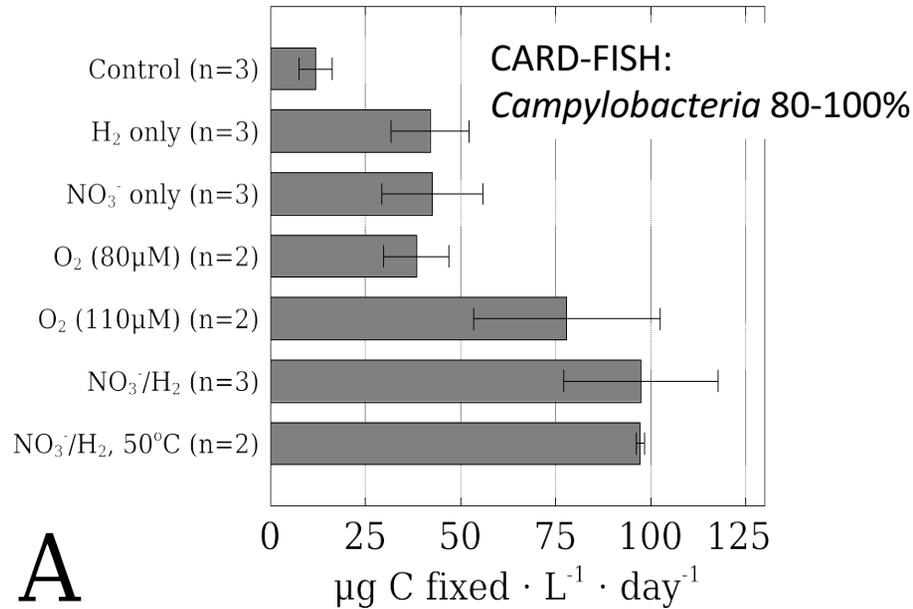
B)



C

***Sulfurimonas* OTUs respond to differences in O₂-concentration -> niche partitioning?**

Estimations of Primary Productivity in Incubations of Hydrothermal Vent Fluids at *In Situ* Temperature and Pressure Determined by HISH-SIMS



- *Campylobacteria* dominate carbon fixation!
- Amendments increased carbon fixation
- Active carbon fixation (and DNRA) also by *Nautiliales* (-> *Thioreductor*) at lower temperatures, w/o H₂, and under oxic conditions
- Evidence for physiological versatility not currently reflected in cultures

Constraints on Subseafloor Productivity, Standing Stock, and Turnover from Measurements of CGE

Parameter	Lower bound	Upper bound	Units
Absolute carbon fixation rates ¹	17.3	321.4	$\mu\text{g C} \cdot \text{L}^{-1} \cdot \text{day}^{-1}$
Chemosynthetic growth efficiency ¹	0.06	0.13	Fraction electron equivalents to Carbon fixation
Estimated in situ carbon fixation ²			
(per L Crab Spa mixed fluid):	104	253	$\mu\text{g C} \cdot \text{L}^{-1}$
(per L Crab Spa end-member fluid):	1.4×10^3	3.5×10^3	
Estimated annual productivity ³ of:			
Crab Spa vent ⁴	6.1×10^3	1.5×10^4	$\text{g C} \cdot \text{y}^{-1}$
Surrounding vent field ⁵	3.8×10^6	9.3×10^6	
Global diffuse-flow vents ⁶	4.5×10^{10}	1.4×10^{12}	
Standing stock ⁷ , Crab Spa	28.6	NA	g C
Biomass residence time ⁸ , Crab Spa	17	41	hours
Global standing stock ⁶	1.4×10^9	2.7×10^9	g C

Synthesis

Ambient Deep-Sea Water:

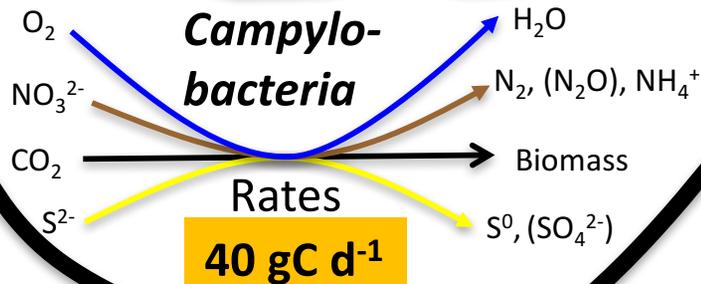
- 2°C
- 2.3 mM DIC
- 115 μM O₂
- 40 μM NO₃⁻
- 0 μM S²⁻, H₂
- 10⁴ cells/ml
- ~0% *Campylobacteria*

Diffuse-Flow Fluid:

- 25°C
- 8.2 mM DIC
- < 3.6 μM O₂
- < 6 μM NO₃⁻
- ~12 μM NH₄⁺
- ~ 200 μM S²⁻
- < 2 μM H₂
- 2-5 * 10⁵ cells/ml
- ~80% *Campylobacteria*

Water

Crust



Flow rate: 162 m³ d⁻¹

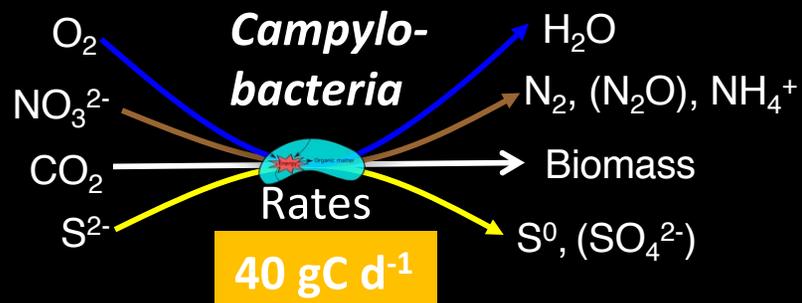
Growth of *Campylobacteria* under e⁻ acceptor limitation

- **Sulfide Oxidation w/ O₂ and NO₃⁻ to S⁰**
 - Mesophiles
 - Oxygen reduction: CBB3
 - Nitrate reduction: **NAP, NIR, NOR, NOS**
 - Sulfide oxidation: **SQR, SOX** (incomplete)
- **Hydrogen Oxidation w/ NO₃⁻ and S⁰**
 - moderate thermophiles, some mesophiles
 - DNRA and denitrification
 - Ni/Fe hydrogenase

Hydrothermal Fluid:

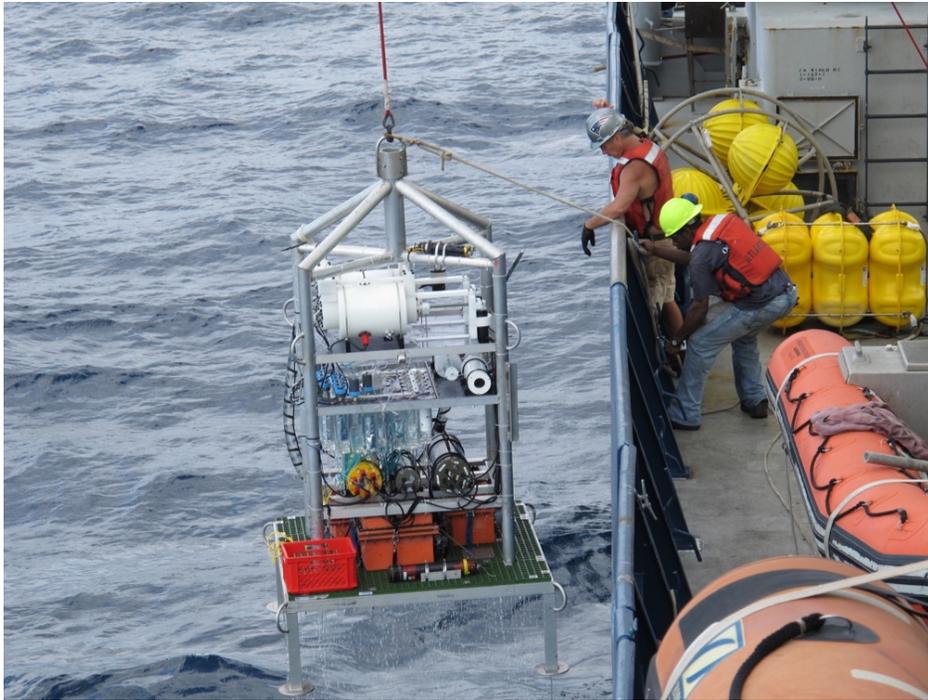
- >275°C
- 85 mM DIC
- 7.7 mM S²⁻
- 410 μM H₂
- 0 μM O₂
- 0 μM NO₃⁻
- 0 Cells

Subseafloor productivity rivals above seafloor production by symbiotic associations!

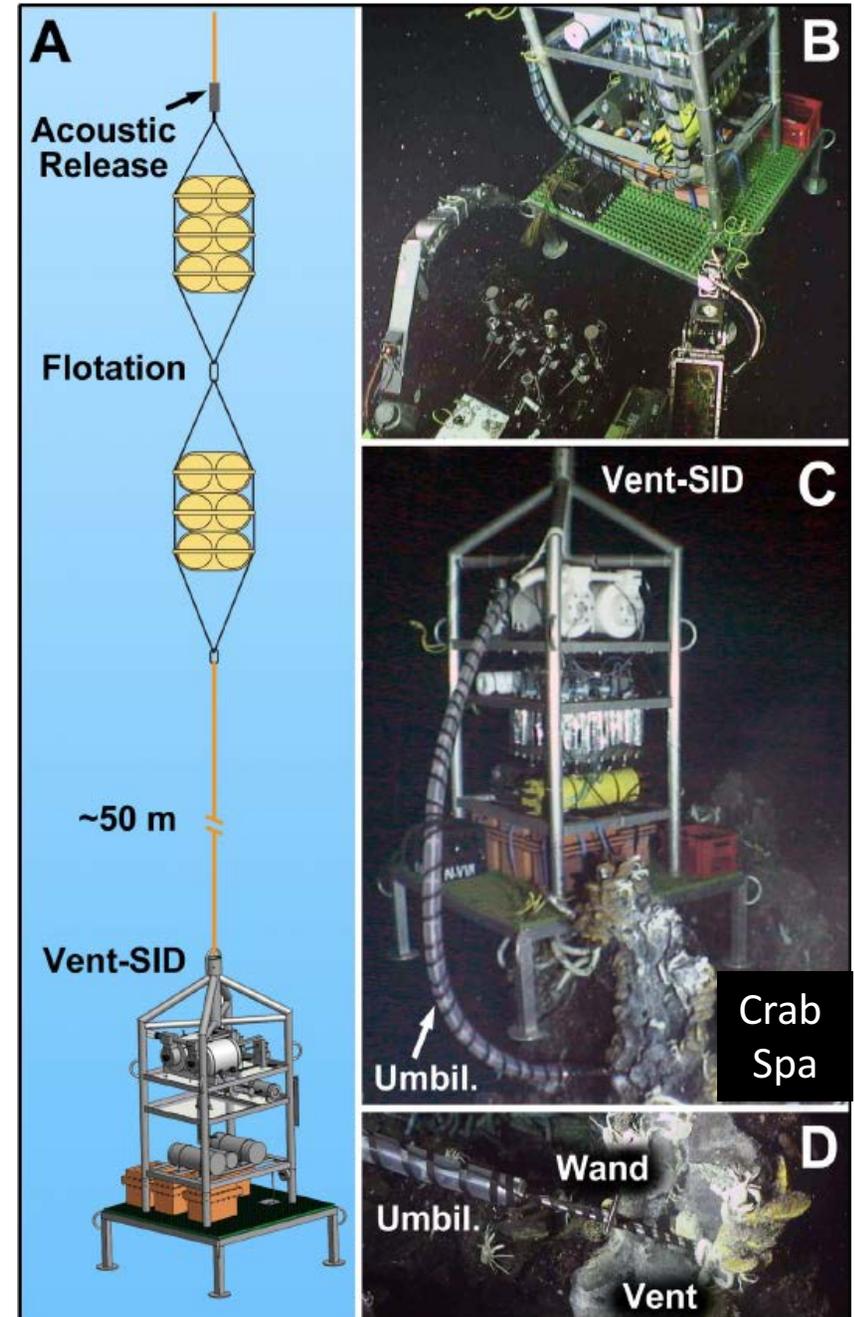


- Diversity
- Pathways

Next Step: Incubations at the Seafloor



- Deployed and tested at Crab Spa, 9°N EPR (2014, 2017)
- Directly correlate the measurement of these processes with community composition and gene expression
- Can perform repeated incubations



CONCLUSIONS

- *Campylobacteria* are dominant members of microbial communities at deep-sea hydrothermal vents
- Have conserved pathways and mechanisms that have allowed them to colonize a variety of environments, including humans
- Connection between environmental microbiology and medical microbiology, one can inform the other
- Deep-sea vents are potentially rich resource for bioactive compounds: high density, productivity, competition, etc.
- *Campylobacteria* dominate chemoautotrophy at Crab Spa
- Activity and rate measurements indicate very active community
- Processes thought to operate separately co-occur: aerobic/anaerobic, denitrification/DNRA, sulfide/hydrogen oxidation
- First direct estimates of subseafloor productivity, standing stock, and turnover!
- Subseafloor productivity rivals above seafloor production by symbiotic associations

MANY THANKS TO ...

SIEVERT lab

Florian Götz, Lara Gulmann,
Jesse McNichol, François
Thomas

WHOI

Jeff Seewald
Craig Taylor
Sean Sylva
Kerry McCulloch

University of Greifswald

Thomas Schweder
Stephanie Markert
Dörte Becher
Stephan Fuchs

UFZ Leipzig

Niculina Musat
Hryhoriy Stryhanyuk
Sabrina Lübke

Bigelow Lab

Ramunas Stepanauskas
Maria Pachiadaki
Jessica Labonte

Rutgers University

Gostantino Vetriani
Ashley Grosche
Donato Giovanelli

UPMC Banyuls-sur-Mer

Nadine Le Bris
Erwan Peru

Shanghai JiaoTong Univ.

Fengping Wang
Xiao Xiang

University of Maine

Jeremy Rich
Sean O'Neill

Carnegie Geophysical Lab

Dionysis Foustoukos
Ileana Perez-Rodriguez
(now UPenn)

Georgia Tech

Leonid Germanovich

Joint Genome Institute

Tanja Woyke

Shipboard scientific parties (AT15-28, AT15-38, AT26-10, AT26-23, AT37-12)
Captain, crew, and pilots of R/V ATLANTIS and DSRV ALVIN and ROV Jason-II



Thank you!

Questions?

