

# Impact of heavy metal and nutrient pollution on the microbiota of temperate corals from the Mediterranean Sea

Jeroen A.J.M. van de Water<sup>§</sup>, Christian R. Voolstra<sup>¥</sup>, Denis Allemand<sup>§</sup> & Christine Ferrier-Pagès<sup>§</sup>

<sup>§</sup> Centre Scientifique de Monaco, 8 Quai Antoine 1er, MC 98000, Principality of Monaco - Contact: [jvdewater@centrescientifique.mc](mailto:jvdewater@centrescientifique.mc)  
<sup>¥</sup> Red Sea Research Center, King Abdullah University of Science and Technology (KAUST), Thuwal 23955-6900, Saudi Arabia

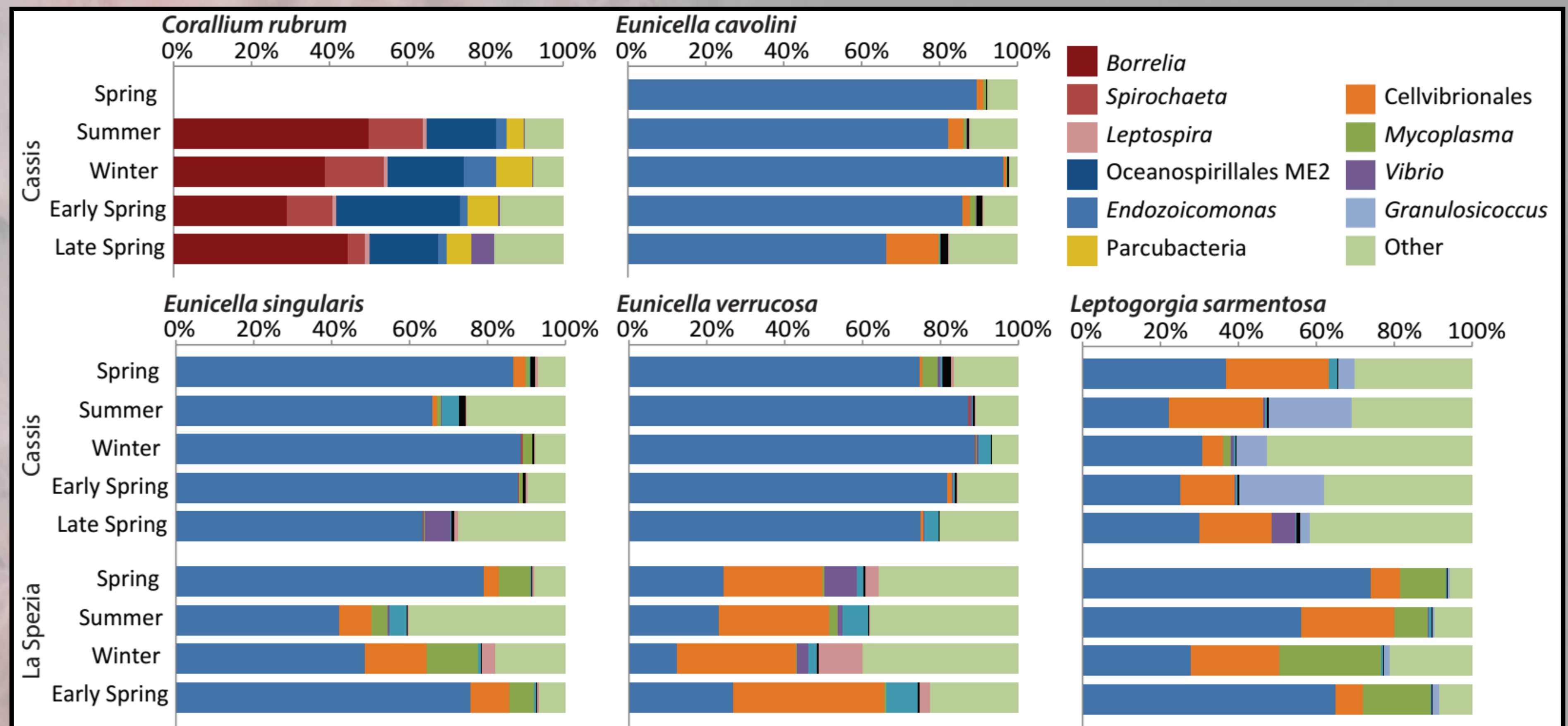
## Introduction

- Gorgonians are key habitat-forming species of temperate benthic communities<sup>1</sup>.
- Dramatic population declines due to local human impacts and mass mortality events caused by high temperatures and disease outbreaks<sup>2</sup>.

Recently, we have shown<sup>3,4,5</sup>:

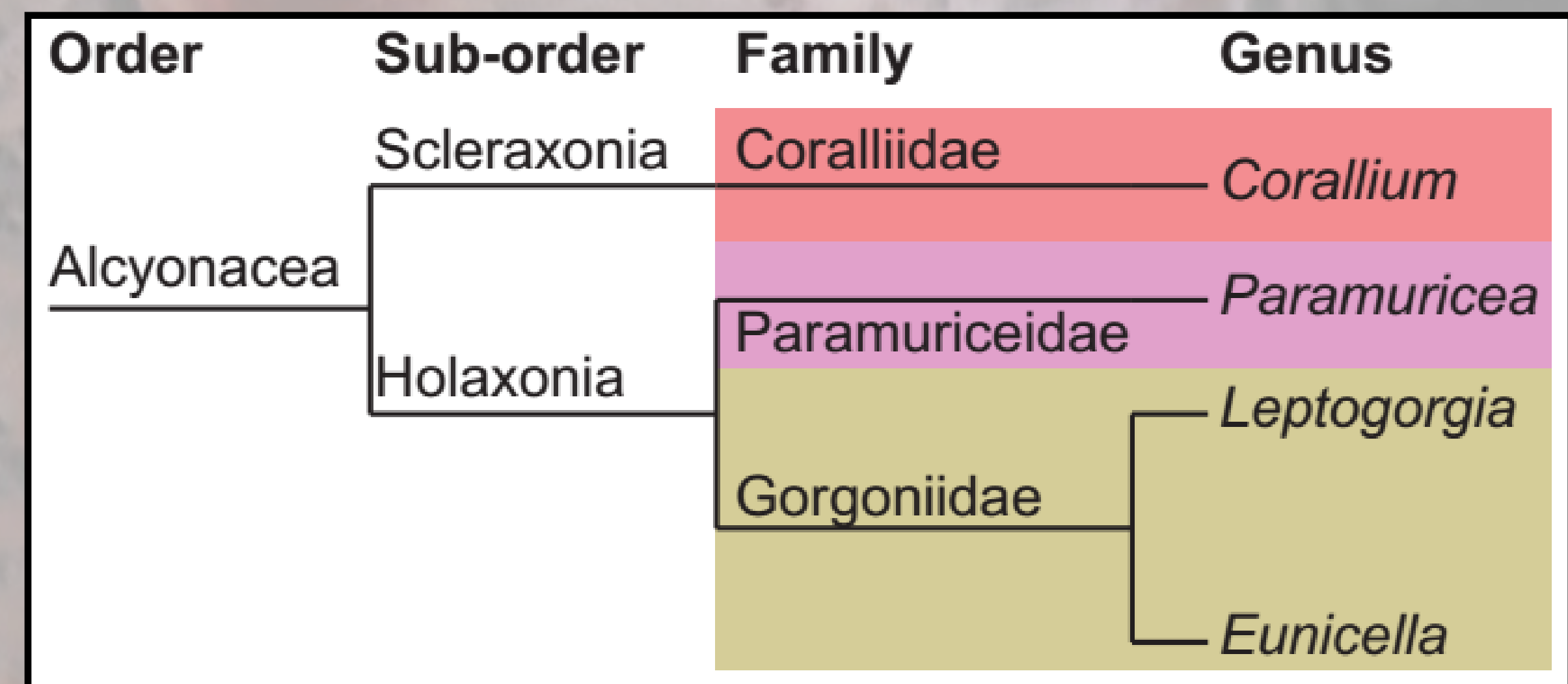
- Ancient host-microbe associations conserved through evolutionary times.
- Divergence in microbiome composition clear along distant phylogenetic lines.
- Highly structured and relatively stable gorgonian-associated bacterial communities on both temporal and seasonal scales
- Unique composition of *Corallium rubrum* microbiota
- Microbiome impacted by / acclimated to local environmental conditions.

**OBJECTIVES** – Understand functional differences in the microbiota of gorgonians and how it is impacted by the most common pollutants in the Mediterranean Sea: copper and nutrients.



**Figure 1** - Composition of the bacterial communities associated with five different gorgonians over time at two different locations – Cassis and La Spezia.

**Figure 2** - Schematic overview of Mediterranean gorgonian taxonomy. The different colours identify taxa harbouring distinct core microbiomes.



## Material & Methods

**Study species (encompassing 3 families, 2 sub-orders)**

- Eunicella cavolini*
- Paramuricea clavata*
- Corallium rubrum*

**Experimentally exposed to copper pollution and eutrophication for 18 hours (n=6 per condition)**

**Metatranscriptomics analysis**

- Library preparation – rRNA and polyA mRNA-depleted Sequencing on Illumina HiSeq4000
- Data analysis: SAMSA2 pipeline<sup>6</sup> and DESeq2<sup>7</sup>

## Discussion & Conclusions

- Eutrophication has no effect on the microbiota of temperate gorgonians
- Bacteria limit toxicity of copper by upregulation of heavy metal transporters
- Copper may affect host health due to increased bacterial virulence and inactivation of phages<sup>8</sup>  
→ Long term effect microbiota composition?
- Major functional differences in the microbiota of sympatric gorgonian species  
→ *C. rubrum*: protection against bacteria and phages, amino acid production, iron uptake.

## Results

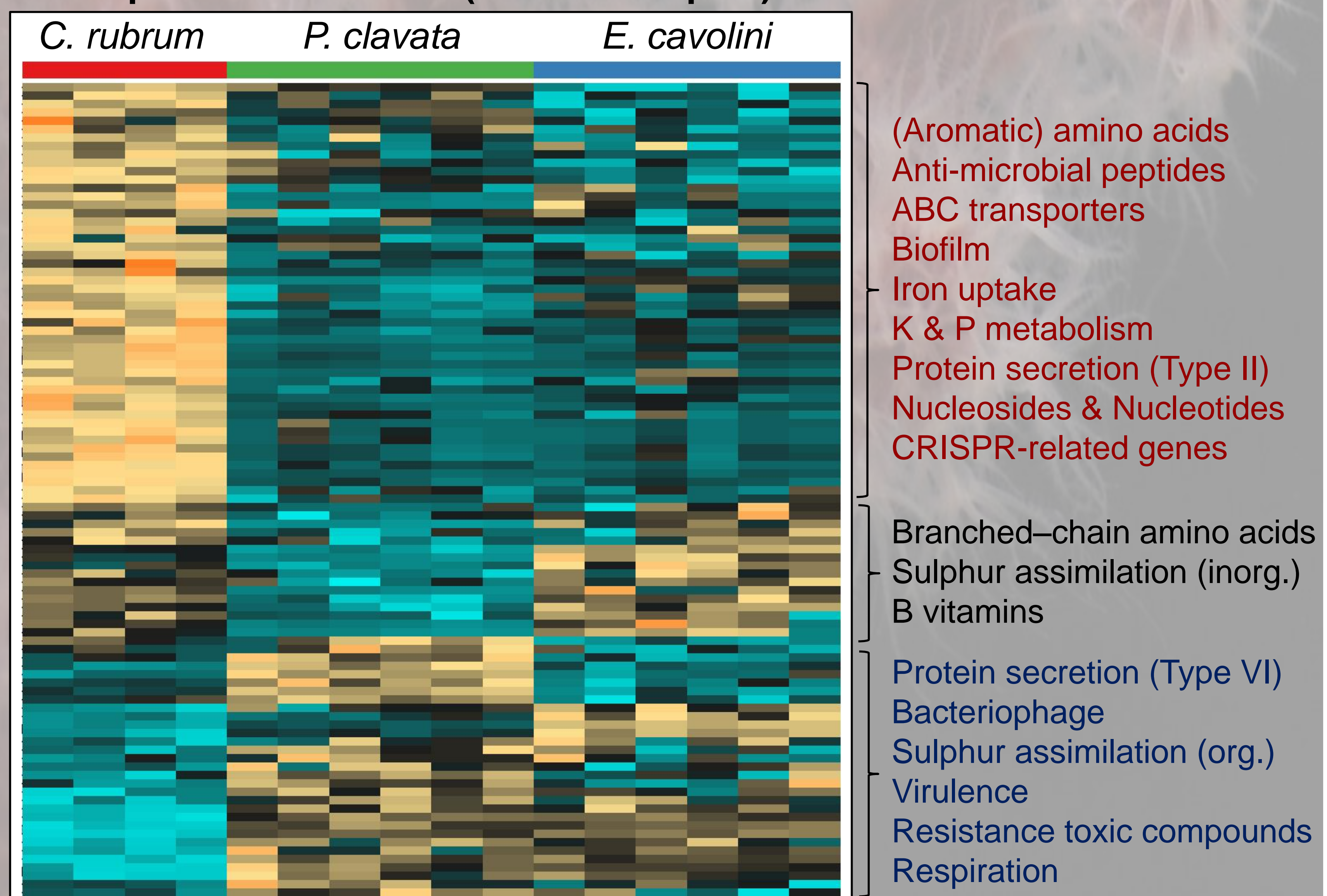
**Eutrophication:** No short-term effect

**Copper pollution:** No effect on *C. rubrum* microbiota

Effect on microbiota of *P. clavata* and *E. cavolini* is similar

- > upregulation copper and heavy metal transporters, and chaperone DnaK
- > increased virulence and reduced phage-related processes

**Inter-species differences (control samples):**



### References

- Ballesteros E. *Oceanography and Marine Biology - An Annual Review*. 2006: 123-195.
- Bally M & Garrabou J. *Global Change Biology* 2007, 13 (10):2078-2088.
- van de Water JAJM et al. *Scientific Reports*. 2016 June; 6: 27277
- van de Water JAJM et al. *Microbial Ecology*. 2016 October; 73(2): 466-478

### References

- van de Water JAJM et al. *Microbial Ecology*. 2018 January; 75(1): 274-288
- Westreich ST et al. *BMC Bioinformatics*. 2018 May; 19:175
- Love MI et al. *Genome Biology*. 2014 December, 15(12): 550
- Li J & Dennehy JJ. *Applied & Environmental Microbiology*. 2011 October; 77(19)

### Acknowledgements

- Eric Tambutti - background photo ©
- Funding provided by Fondation Paul Hamel & KAUST