

Supplementary Material

Bacterial Community Composition in the Sea Surface Microlayer off the Peruvian Coast

Birthe Zäncker^{1*}, Michael Cunliffe^{2,3}, Anja Engel¹

¹GEOMAR – Helmholtz Centre for Ocean Research Kiel, Kiel, Germany

²Marine Biological Association of the UK, The Laboratory, Citadel Hill, Plymouth, UK

³Marine Biology and Ecology Research Centre, School of Biological and Marine Sciences, Plymouth University, Plymouth, UK

***Correspondence:**

Birthe Zäncker

bzaencker@posteo.de

1 Supplementary Figure

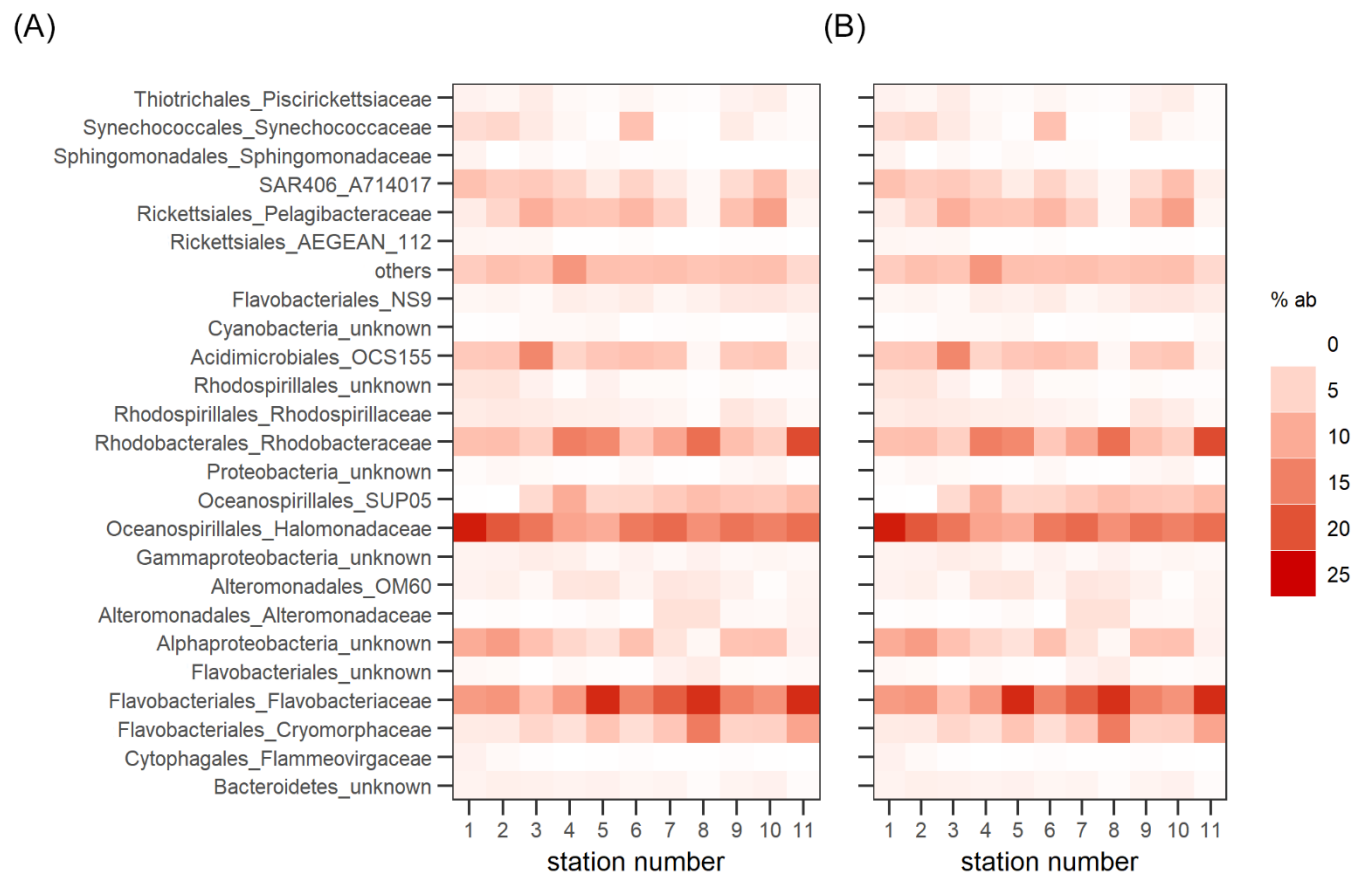


Figure S1: Relative abundance of bacterial families that make up at least 1 % of the total bacterial community in at least two stations. Abundance at each individual station in the SML (A) and the ULW (B). White indicates no sequences of a certain bacterial family were found, the intensity of red indicates the relative abundance of each bacterial family.