

EGU21-5614

<https://doi.org/10.5194/egusphere-egu21-5614>

EGU General Assembly 2021

© Author(s) 2022. This work is distributed under the Creative Commons Attribution 4.0 License.



Understanding Bacterial Ice Nucleation

Ralph Schwidetzky¹, Max Lukas¹, Anna T. Kunert², Ulrich Pöschl², Janine Fröhlich-Nowoisky², Mischa Bonn¹, and Konrad Meister^{1,3}

¹Molecular Spectroscopy Group, MPI for Polymer Research, Mainz, Germany

²Multiphase Chemistry, MPI for Chemistry, Mainz, Germany

³Arts and Science, University of Alaska Southeast, Juneau, USA

Bacterial ice-nucleating proteins (INPs) promote heterogeneous ice nucleation better than any known material. On the molecular scale, bacterial INPs are believed to function by organizing water into ice-like patterns to enable the formation of embryonic crystals. However, the details of their working mechanism remains largely elusive. Here, we report the results of comprehensive evaluations of environmentally relevant effects such as changes in pH, the presence of ions and temperature on the activity, three-dimensional structure and hydration shell of bacterial ice nucleators using ice affinity purification, high-throughput ice nucleation assays and surface-specific sum-frequency generation spectroscopy.

[1] Lukas, Max, et al. "Electrostatic Interactions Control the Functionality of Bacterial Ice Nucleators." *Journal of the American Chemical Society* 142.15 (2020): 6842-6846.

[2] Lukas, Max, et al. "Interfacial Water Ordering Is Insufficient to Explain Ice-Nucleating Protein Activity." *The Journal of Physical Chemistry Letters* 12 (2020): 218-223.