

Antifreeze Proteins from a Sea Ice Diatom

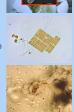
Maddalena Bayer-Giraldi, Ilka Weikusat, Gerhard Dieckmann Alfred Wegener Institute for Polar and Marine Research Bremerhaven, Germany



Alfred-Wegener-Institut für Polar- und Meeresforschung in der Helmholtz-Gemeinschaft

INTRODUCTION

Polar sea ice is an extreme habitat with regard to its ablotic constraints as low temperatures and high brine salinities. However, several microorganisms proliferate within the ice and on its peripheries, thus creating an ecosystem of global significance. The diatom *Fragilariopsis cylindrus* is a dominant species in sea ice communities, and its recently found antifreeze proteins (AFPs) probably contribute to its success in ice. AFPs are proteins that bind to ice and influence its growth, lower the freezing point (freezing hysteresis) and inhibit recrystallization¹. Here we present a characterization of an AFP from *F. cylindrus* (fcAFP) in a crystallographic frame and interpret our data in the context of sea ice.



ANTIFREEZE ACTIVITY

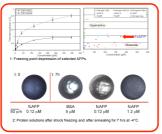
We used a fcAFP isoform shown to be active under conditions typical for sea ice and recombinantely expressed in *E. coll.*

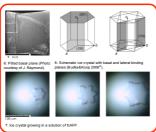
The protein showed moderate freezing hysteresis activity, further increased by salt addition (1). Moreover, FcAFP caused strong inhibition of recrystallization in shock frozen polycrystalline ice (2). The effect was monitored for 7 hours at -4°C.

INTERACTION WITH ICE

Submersion of an ice crystal into a solution of fcAFP at temperatures below the freezing point caused dense pitting formation on the basal plane (6). Crystals growing in

the presence of fcAFPs were star-like (7), while control showed a disclike habit. These results suggest that fcAFPs bind on the basal plane and the lateral faces of ice crystals (8).

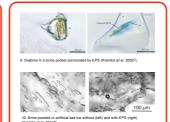




ICE MICROSTRUCTURE

Frozen fcAFP solutions showed marked influence of the protein on ice microstructure. FcAFP caused structures with oriented striations (3-4).

Image: Constraint of the second se



A mushy structure, possibly with inclusions with brine and unbound proteins, may have formed due to irregularities in ice growth caused by the bound fcAFPs.

Moreover, the effect of fcAFPs caused gradual c-axes transitions and irregular grain boundaries (5).

FcAFP IN SEA ICE

Physiological protein concentration measured by immuno-blotting in cells was too low for an effective freezing hysteresis and recrystallization inhibition.

FcAFPs may accumulate around cells in an envelope of exopolymeric substance (EPS), made of polysaccharides and proteins (9). The protein fraction was shown to be effective in influencing sea ice structure, increasing porosity and retaining brine (10)⁴.

CONCLUSIONS

FcAFPs have strong antifreeze activity, that may be of interest for industrial applications. In sea ice, physiological concentrations are too low to have any relevant antifreeze activity. However, fcAFPs may accumulate in EPS and reach high concentrations. In contact with ice, fcAFPs may create a mushy structure as in picture 3, forcing the formation of brine pores and thus creating an environment where sea ice organisms can locate and survive.

Kristiansen, Zachariassen (2005): The mechanism by which fish antifreeze proteins cause thermal hysteresi, Cryobiology;

Budie, Koop (2008): be receptibilization hibition and molecular recognition of be based by Polylinyl elochidi, ChemPhysChem; Yamba, Elden, Junga, Daming (2002): High cooperatations of expopyments abstances in writeratime area to:: implications for the base of a cooperative and acyoprotection of distoms, Daep-Sea Res Kometa. Damina. Tealan: 2011: Economic alterations of thebase increasing of a gas be and implications for the habitability and biogeochemistry in a warmer Actio, FNAS.