## Metadata of the chapter that will be visualized online

Chapter Title	Thermal Hysteresis	
Copyright Year	2020	
Copyright Holder	pringer Nature Switzerland AG	
Corresponding Author	Family Name Kristiansen	
	Particle	
	Given Name Erlend	
	Suffix	
	Organization NTNU University Library	
	Address Trondheim, Norway	
	Email erlendkr@ntnu.no	
Abstract	•	
Keywords (separated by '-')	Thermal hysteresis - Antifreeze activity - Kelvin effect - Critical radius - Hyperactive - Hyperactivity - Protein size - Protein solubility - Hofmeister series - Additive - Enhancement - Adsorption - Desorption - Heterogeneous nucleation - Ice nucleator - Ice nucleating agent	

Chapter 6	
<b>Thermal</b>	Hysteresis

1 2

3

10

11

12

13

14

15

16

17

18

19

20

21

22

Erlend Kristiansen

#### 6.1 Introduction

Thermal hysteresis refers to the phenomenon where antifreeze proteins (AFPs) or 4 antifreeze glycoproteins (AFGPs) cause a separation of the freezing and melting 5 temperature of existing ice crystals in solution. This ability to separate the melting 6 and freezing temperature of ice is limited in that on sufficient cooling the ice crystal 7 undergoes a sudden and rapid ice growth. Ramsay (1964) when studying a mech- 8 anism of water reabsorption in the beetle Tenebrio molitor first reported the phe- 9 nomenon. In a footnote, he states:

When small ice crystals are observed under the microscope, as in the freezing-point method of Ramsay and Brown, one notices that large crystals grow at the expense of small ones and that the edges of the crystals are rounded—the natural consequences of surface tension at the water-ice interface. The change of state between solid and liquid is perfectly temperaturereversible.....By contrast, the crystals which appear in fluid from the anterior perinephric space tend to have jagged outline and large crystals do not grow at the expense of smaller ones. Furthermore, the system is not temperature-reversible. As the temperature is raised the crystals decrease in size, but as the temperature is lowered they do not increase in size. After the temperature has been lowered by a few degrees the crystal suddenly begins to grow rapidly. On occasion undercooling of the order of 10 °C was observed (in the continued presence of small crystals) and then suddenly the whole sample appeared to solidify instantaneously.

The temperature interval between the melting and freezing temperatures is 23 referred to as the hysteresis gap, and the lower temperature where rapid ice growth 24 is initiated is termed the hysteresis freezing point. The quantitative difference 25 between the melting temperature and the hysteresis freezing point is termed the 26 hysteresis activity, or antifreeze activity. 27

E. Kristiansen (⊠)

NTNU University Library, Trondheim, Norway

e-mail: erlendkr@ntnu.no



Thermal hysteresis reflects the role of AF(G)Ps as protectors against ice nucleation in the supercooled body fluids of freeze avoiding animals. Their presence enables hypoosmotic fish to occupy ice-laden polar waters (DeVries 1971, 1982; Raymond and DeVries 1977), and allow terrestrial arthropods, such as insects, spiders, and collembolans, to remain year-round in the cold temperate and polar areas. The body temperatures of such terrestrial animals may in some cases drop well below –30 °C in winter (Zachariassen and Husby 1982; Duman 2001; Duman et al. 2004; Graham and Davies 2005). Within the animal, AF(G)Ps are known to act by inactivating structures in the body fluids that could initiate freezing, so-called ice-nucleating agents (INAs), and by preventing ice from penetrating through the body wall (Olsen and Duman 1997a, b; Olsen et al. 1998; Duman 2002).

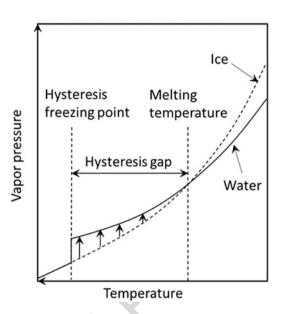
AF(G)Ps are categorized as being moderately active or hyperactive, based on the hysteresis activity they cause at equimolar concentrations. This distinct difference in antifreeze potency is accompanied by distinct shapes of the ice crystals that form in their presence; moderately active AF(G)Ps cause bipyramidal crystals to develop, a shape that only exposes a single crystal plane to the surrounding solution. In the presence of hyperactive AF(G)Ps, ice crystals express several crystal planes, usually in the form of hexagonal discs. A number of different factors affect the hysteresis activity, including their size and the addition of large organic macromolecules and inorganic ions. This chapter outlines current understanding of the modus operandi of AF(G)Ps. An attempt is made to provide some simple explanations to the antifreeze potency of AF(G)Ps, including their characteristics as moderately active or hyperactive, and how their antifreeze potency is affected by their size and by different additives. Some characteristics of INAs and their relevance in cold tolerance are also examined briefly.

## 6.2 A Hysteresis Mechanism: The Kelvin Effect

The vapor pressure of bulk ice is lower than that of water. Thus, below the melting point a net transfer of water molecules from the bulk water to ice occurs and the ice mass grows. However, it follows from the observable fact that ice crystals in the presence of AF(G)Ps remain unchanged within a temperature interval, that the AF (G)Ps somehow causes vapor pressure equilibrium between ice and water at all temperatures within the hysteresis gap. This must be so, since the rate by which water molecules adds onto the crystal surface must equal the rate by which they leave. Otherwise, net transfer of water molecules would result, from solution to ice or vice versa and the crystal would visibly change volume. AF(G)Ps do not lower the vapor pressure of water any more than other solutes do (Westh et al. 1997). Thus, they must act by elevating the vapor pressure of the ice to correspond to the higher vapor pressure of the surrounding solution. The difference between the vapor pressure of water and ice increases with temperature departure below the equilibrium melting temperature. Thus, the effect of the AF(G)Ps on the vapor pressure of ice

### 6 Thermal Hysteresis

Fig. 6.1 Vapor pressure equilibrium within a temperature interval near the melting temperature. For the ice crystal to be stable within the hysteresis gap, the AF(G)Ps must elevate the vapor pressure of the ice surface to correspond to that of the surrounding supercooled solution. This elevation of the vapor pressure must increase with decreasing temperature. Adapted from Kristiansen and Zachariassen (2005)



69

75

85

must be temperature dependent and increase with decreasing temperature, see 68 Fig. 6.1.

Raymond and DeVries (1977) proposed that the AF(G)Ps act by changing the 70 microscopic growth pattern of the ice surface. Since this is achieved by the AF(G)Ps 71 becoming irreversibly adsorbed onto the ice surface, they coined the mechanism the 72 adsorption-inhibition mechanism. Since then, several investigators have had similar 73 approaches to explaining the phenomenon by irreversible adsorption, including 74 Wilson (1993) and Kristiansen and Zachariassen (2005).

Using fluorescently tagged AFPs, Celik et al. (2013) exchanged the slightly 76 supercooled solution surrounding an ice crystal. The ice surface of the supercooled 77 crystals remained fluorescent following the exchange of the surrounding solution, 78 showing that AFPs were adsorbed onto the crystal surface. Further, the removal of 79 AFPs in the surrounding solution by the exchange process did not weaken the 80 hysteresis effect. These observations provide the most unequivocal evidence to 81 date to show that AF(G)Ps become irreversibly adsorbed onto the ice surface and 82 that the phenomenon is caused only by the surface-bound AF(G)Ps. Also, Chao et al. 83 (1995) and DeLuca et al. (1998) found that AF(G)Ps principally operate as mono- 84 meric units.

Elevation of the vapor pressure of the ice by the changed microscopic surface 86 growth pattern could occur by the so-called Kelvin effect. In the following, a brief 87 historical outline of the Kelvin effect is provided. This is followed by a description of 88 how the Kelvin effect is thought to operate at the ice surface. 89



## 6.2.1 The Kelvin Effect: Vapor Pressure at a Curved Interface

In 1871, Prof. William Thomson, later to become first Baron Kelvin, pointed out that 91 92 the vapor pressure of water at a concave and a convex surface must be lower and higher, respectively, than at a plane surface of the water (Thomson 1871). This was 93 deduced by considering the rise and fall of liquids in a capillary tube as a function of 94 the curvature of the meniscus; in an atmosphere saturated with vapor, the vapor 95 pressure decreases with height above the surface of a liquid. Consequently, since a 96 concave interface in a capillary causes the liquid to come to rest at some fixed height 97 above the liquid body, Thomson deduced that the vapor pressure at the elevated 98 concave meniscus is reduced relative to the vapor pressure at the lower plane surface 99 and must correspond to the lowered saturated atmospheric vapor pressure at that height. Otherwise, a perpetual net directional motion of water molecules would develop, as there would be continuous net evaporation at the elevated meniscus 102 and consequently net condensation at the lower plane surface. Such perpetual motion 103 of water molecules would violate the fundamental law of thermodynamics. Convex 104 interfaces must have the opposite effect on the vapor pressure, as such an interface 105 comes to rest below the plane liquid body where the saturated vapor pressure is higher. The effect of a surface curvature on the vapor pressure has since become 107 known as the Kelvin effect.

### 109 6.2.1.1 The Critical Radius of Curvature

A decade later, Prof. John Henry Poynting (1881) recognized that the effect of a surface curvature on the resultant vapor pressure in Thomson's capillary is caused by a change in the bulk pressure in the water in the capillary; a concave interface evokes a lower pressure inside the liquid water, as evident from the rise in the capillary, and hence to a lower vapor pressure, and vice versa for a convex interface. Thus, the underlying cause of the changing vapor pressure with changing curvature of an interface is an accompanying curvature-induced change in bulk pressure within the curved volume.

118 Poynting applied his reasoning to the melting temperature of ice. He inferred that if the bulk pressure of ice alone was elevated, then the resultant elevated vapor 119 pressure of the ice would depress the temperature at which the vapor pressures of ice 120 and water coincides, i.e., a pressure-induced depression of the melting temperature 121 of the ice surface. By extension, since the pressure-elevating effect of a convexity 122 123 increases with decreasing radius, there must be a convexity with a radius small enough to cause a pressure great enough for ice/water vapor pressure equilibrium to 124 develop at any temperature below the normal melting point. The radius of this convexity at a specific temperature is referred to as the critical radius of curvature at that temperature.

AU1

145

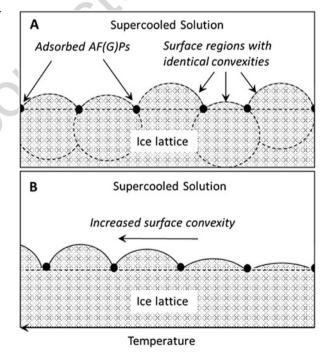
## The Kelvin Effect at the Ice Surface

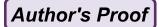
It follows from the above paragraphs that AF(G)Ps that are irreversibly adsorbed 129 onto the ice surface could evoke the Kelvin effect by causing the ice surface to grow out as many tiny convex interfaces between them. These convex interfaces would 131 elevate the vapor pressure of the ice surface and, hence, eliminate the difference 132 between the vapor pressures at different temperatures, as illustrated in Fig. 6.1.

The Kelvin effect implies that, at any temperature below the normal melting 134 temperature, the growth of the convex surface zones between the adsorbed AF(G)Ps 135 will halt when they obtain a curvature with a radius corresponding to the critical 136 radius at that temperature. Thus, at any temperature where the phenomenon is expressed, the surface of the entire ice crystal is covered by spherical growth regions with identical convexities, i.e., identical local vapor pressures. This causes the entire ice crystal surface to be in vapor pressure equilibrium with the surrounding 140 supercooled solution, and hence the crystal surface is at its melting temperature, 141 see Fig. 6.2. A. Such a crystal could in principle remain unchanged indefinitely. 142 Crystals in supercooled solutions of AF(G)Ps have been observed for many days 143 without expressing any visible growth (DeVries 1971; Raymond and DeVries 1977; 144 Graether et al. 2000; Fletcher et al. 2001).

As the temperature is lowered further, the many tiny surface zones expand until 146 their convex interfaces again cause vapor pressure equilibrium with the surrounding 147

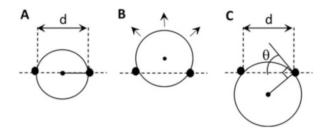
**Fig. 6.2** The convexities of the growth zones within the hysteresis gap. (a) All growth zones must have the same convexity at a specific temperature within the hysteresis gap. (b) The convexities increases with decreasing temperature and elevates the vapor pressure of the ice surface in a temperature-dependent manner, as seen in Fig. 6.1. Adapted from Kristiansen and Zachariassen (2005)





151

152



**Fig. 6.3** At the hysteresis freezing point. (a) When one of the convexities has reached the shape of a half-sphere it has reached its maximum convexity. (b) Any further growth of this structure will cause the convexity to decrease and cause spontaneous growth. (c) The relation between adsorbent spacing, d, and the angle,  $\theta$ . Adapted from Kristiansen and Zachariassen (2005)

solution. In this manner ice/water vapor pressure equilibrium is maintained across a temperature interval, the hysteresis gap, see Figs. 6.1 and 6.2 b.

There is a limit to how much such a crystal can be cooled, i.e., how convex the tiny curved interfaces may become; no surface zone can become more convex than that of a half-sphere. Once such a shape is reached, then any further cooling will result in the convexity of the structure to decrease on growth. The resultant drop in vapor pressure due to the reduced convexity will result in spontaneous growth. This is illustrated in Fig. 6.3a and b. This temperature is the hysteresis freezing point.

## 66 6.3 Hysteresis Activity

157 In the following paragraphs, an attempt is made to explain what fundamentally determines the hysteresis freezing point, based on the theory outlined above. This explanation is then extended to incorporate the characteristic difference in activity 160 between moderately active and hyperactive kinds of AF(G)Ps.

# 161 6.3.1 The Largest Intermolecular Adsorbent Gap Determines 162 Hysteresis Activity

163 If only a single one of all the tiny growth zones that protrude out at the crystal surface 164 should fail, then the hysteresis phenomenon is terminated. Hence, the hysteresis 165 freezing point is determined by the single growth zone that reaches the shape of a 166 half-sphere at the highest temperature. Any further growth of this single growth 167 zone, i.e., any further cooling, will only result in a reduction in its convexity and, 168 consequently, the phenomenon is terminated.

Since all the surface growth zones have the same convexity, it will be the single one growth zone with the widest diameter that will reach the shape of a half-sphere at

### 6 Thermal Hysteresis

the highest temperature. Thus, the hysteresis freezing point, and therefore the 171 hysteresis activity, is determined by the single largest intermolecular adsorbent 172 spacing between AF(G)Ps that comprise a single growth zone at the crystal surface. 173 Mathematically, the hysteresis activity ( $\Delta T$ ) as a function of the largest such 174 adsorbent spacing, d, may be expressed as (Kristiansen and Zachariassen 2005): 175

$$\Delta T = \frac{4\gamma T_{\rm E} \sin \theta}{\Delta H_{\rm d}};\tag{6.1}$$

181

182

186

where d is the spacing in units of cm,  $\gamma$  is the ice/water interfacial tension (taken to 176 be 32 ergs/cm<sup>2</sup>), T<sub>E</sub> is the normal melting temperature for a plane interface (units of 177 K), and  $\Delta H$  is the heat of fusion of water (3.3 · 10<sup>9</sup> ergs/cm<sup>3</sup>).  $\theta$  is an angle 178 describing the situation if a curvature fails before reaching the shape of a halfsphere. For a half-sphere,  $\theta$  is 90° and, hence, the term (sin  $\theta$ ) is 1. See Fig. 6.3. C for an illustration of the angle  $\theta$ .

#### Moderately Active and Hyperactive AF(G)Ps 6.3.2

There is a great difference in the hysteresis activities caused by different AF(G)Ps. Based on their activities at equimolar concentrations and the shape of the crystals they form in solution, they fall into two categories: hyperactive and moderately 185 active.

Marshall et al. (2004a) found that moderately and hyperactive AFPs accumulate 187 in ice to a similar extent. Also, experimentally determined estimates of average 188 adsorbent spacings between AF(G)Ps on the surface of ice crystals are quite similar 189 in the case of moderately and hyperactive AF(G)Ps; Drori et al. (2015) estimated the 190 average adsorbent distance between hyperactive TmAFP to 7.6–35.2 nm at concentrations ranging from 31.4 to 0.4 µM. Comparable results were obtained by Celik 192 et al. (2013) for the same protein. For the moderately active type III AFP, Drori et al. 193 (2015) estimated the average adsorbent distance to be 8.7 to 24.7 nm at concentrations ranging from 19.8 to 1.2 µM. Others have estimated similar values for 195 moderately active AF(G)Ps (Wilson et al. 1993; Grandum et al. 1999; Zepeda 196 et al. 2008). Thus, the principal cause of the great difference in the activities of 197 moderately and hyperactive AF(G)Ps do not seem to be due to differences in their 198 preference for ice. Rather, it is likely that the distinct difference between them is the 199 result of the single largest adsorbent gap at the ice surface for some reason is much 200 larger in the case of moderately active AF(G)Ps. 201



224

225

226

227

228

229

230

231

232

## 6.3.2.1 Moderate or Hyperactive: Caused by Plane Specificity and Adsorption Pattern?

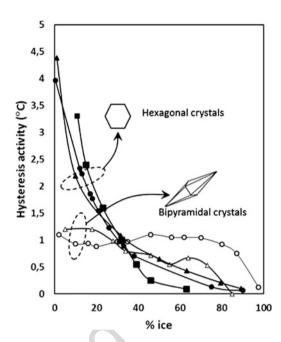
**Moderate Activity** A characteristic feature of moderately active AF(G)Ps is that 204 they only adsorb onto a single crystal plane in the ice structure. Notably, none of the 205 moderately active AF(G)Ps adsorb onto the basal plane of crystals, only onto a single 206 prism or pyramidal plane (Knight and DeVries 1988; Knight et al. 1991). This plane-207 specific adsorption is apparently a consequence of structural features of their 208 ice-binding sites (IBSs), that restricts these AF(G)Ps to only become irreversibly 209 adsorbed onto a single plane and orientation. Laursen et al. (1994) showed this by 210 observing that the moderately active chiral L-AFP I and D-AFP I variants resulted in adsorption on mirror image directions on the ice surface. The result of such a specific 212 preference for a single crystal plane is a crystal that only expresses this single 213 protected crystal plane toward the surrounding supercooled solution. Consequently, in the presence of moderately active AF(G)Ps crystals obtain a bipyramidal shape, as 215 this is the only possible crystal shape whose entire surface consists of a single plane. 216 At the hysteresis freezing point, these bipyramidal crystals freeze out from their 217 apexes (Raymond and DeVries 1977; Jia and Davies 2002). The fact that they 218 characteristically grow out of their apexes at the hysteresis freezing point strongly 219 suggests that the antifreeze potency of moderately active AF(G)Ps are limited by a 220 large intermolecular spacing at the apex of the bipyramidal crystal (Jia and Davies 221 2002). This must arise from the fact that these proteins only adsorb onto a single 222 crystal plane. 223

The surface area involved in determining the hysteresis activity for moderately active AF(G)Ps is only that miniscule fraction of the total surface area of the crystal that comprises the two apexes of the bipyramid. Consequently, the hysteresis activity in the presence of moderately active AF(G)Ps should not be much affected by changing the total surface area of the ice. Consistent with this, the hysteresis activity of moderately active AF(G)Ps are reportedly rather insensitive to the amount of ice present in the sample; large variations in the ice content, i.e., large variations in total ice crystal surface area, does not appreciably affect the hysteresis activity, see Fig. 6.4 (Hansen et al. 1991; Wöhrmann 1996; Sørensen and Ramløy 2001).

**Hyperactivity** In contrast to the moderately active AF(G)Ps, the hyperactive AF 233 (G)Ps have been shown to adsorb to several crystal planes that differ greatly in their 234 orientation, such as both prism and basal planes (Graether et al. 2000; Liou et al. 235 2000). Structural studies have shown that hyperactive AFPs have IBS that afford the 236 protein freedom to adsorb in different orientations and on different planes. Their 237 ability to adsorb onto multiple crystal planes, and most notably the basal plane, is a 238 feature that separates them from the moderately active AF(G)Ps. Basal plane 239 adsorption has been implicated as a key feature that causes them to be hyperactive 240 (Graether et al. 2000; Liou et al. 2000; Pertaya et al. 2008). Because of their ability to 241 adsorb onto multiple planes, crystals formed in the presence of hyperactive AF(G)Ps 242 expresses multiple planes to the surrounding supercooled solution and usually take the form of hexagonal discs (Graether et al. 2000; Liou et al. 2000).

### 6 Thermal Hysteresis

Fig. 6.4 The dependency of the hysteresis activity on the % ice in the sample. Filled symbols: hyperactive AF(G)Ps. Open symbols: moderately active AFGP. (Filled square) PAGP, a hyperactive AFGP from the nototheniid Pleuragramma Antarcticum (Wöhrmann 1996). (Filled circle) Hemolymph from Tenebrio molitor (Hansen and Baust 1988). (Filled triangle) Hemolymph from Rhagium inquisitor (Zachariassen et al. 2002). (Open circle) Serum from P. antarcticum. (Open triangle) AFGP from P. antarcticum (Wöhrmann 1996). For explanation, see text



256

Because of their ability to adsorb onto different planes and at different orienta- 245 tions, hyperactive AFPs likely become spread out across the crystal surface in a 246 rather random adsorption pattern. Such a random pattern should, by chance alone, 247 result in the largest adsorption gap increasing with increasing surface area. Conse-248 quently, the hysteresis activity of hyperactive AF(G)Ps should decrease with increasing crystal surface area. Consistent with this, several investigators have reported 250 strong dependence of hyperactive AF(G)Ps on the amount of ice present in the 251 sample, see Fig. 6.4 (Zachariassen and Husby 1982; Hansen and Baust 1988; 252 Wöhrmann 1996). As can be seen from the figure, "hyperactivity" is apparently a 253 consequence of using small ice crystals in the experiment, since hyperactive AF(G) 254 Ps have a lower hysteresis activity than their moderately active counterparts at higher 255 contents of ice in the samples.

The Shape of the Bipyramidal Apexes When bipyramidal crystals form in the 257 presence of moderately active AF(G)Ps, the ice crystal grows out from the basal 258 planes. Once this bipyramidal shape is formed the crystal stops growing and it 259 remains stable within the hysteresis gap. What is the physical shape of the apex 260 interfaces? Since the moderately active AF(G)Ps do not adsorb onto the basal plane, 261 is the apex a tiny unprotected flat basal plane? If so, then one could envision 262 two-dimensional curved interfaces protruding out only in the direction of the 263 prism planes that form the surrounding edge of the exposed apex basal plane 264 (Raymond and DeVries 1977). The effect of these 2D curvatures that are in the 265 prism plane direction must then also elevate the vapor pressure beyond the base of 266 the curvature toward the center of the flat basal plane in order for vapor pressure 267



equilibrium to persist between the flat apex interface and the surrounding solution.
Another, and perhaps simpler, approach is to assume that the apexes are threedimensional spheres protruding out in the basal plane direction. In any event, it is
these areas of the bipyramidal crystal that apparently determines the hysteresis
activity of the moderately active AF(G)Ps.

## 273 6.4 Factors That Affect the Hysteresis Activity

In the above paragraphs the categorization of AF(G)Ps into moderately active and hyperactive were ascribed to consequences of irreversible adsorption to the ice surface that arises from features of their IBS. In the following, differences in 276 hysteresis activity within each of these categories will be ascribed to the situation that exist prior to the AF(G)Ps becoming irreversibly adsorbed. It will be argued that, 278 while the ice crystal is held at the equilibrium melting temperature, AF(G)Ps acquire 279 an equilibrium distribution between the crystal surface melting region and the 280 surrounding solution. Then, following a cooling event, AF(G)Ps within this surface 281 region freeze onto the solidifying crystal surface and, hence, become irreversibly 282 adsorbed (Kristiansen and Zachariassen 2005). Any change in this distribution 283 pattern prior to the cooling event will result in changes in the surface density of 284 irreversibly adsorbed AF(G)Ps after the cooling event and, hence, to changes in the 285 observed hysteresis activity. Differences in hysteresis activity among hyperactive or 286 among moderately active AF(G)Ps, may be attributed to differences in the solubility 287 of the AF(G)Ps in the solution; a lowered solubility results in a shift in the 288 distribution of the AF(G)Ps toward the ice surface region prior to the cooling 289 event, and hence, to increased hysteresis activity (Kristiansen and Zachariassen 2005; Kristiansen et al. 2008). 291

## 292 **6.4.1** The Factors

Several investigators have reported that the size of the AF(G)Ps can have a profound 293 effect on their capacity to cause thermal hysteresis. For structurally similar isoforms, their potency reportedly increases with molecular size for both moderately active 295 AFGPs (Schrag et al. 1982; Chao et al. 1996; Miura et al. 2001; Baardsnes et al. 296 297 2003; Nishimiya et al. 2003) and hyperactive AFPs (Leinala et al. 2002; Marshall et al. 2004b; Liu et al. 2005; Mok et al. 2010; Friis et al. 2014). Synthetic oligomers 298 of moderately active AFPs also reportedly have increased potency (Nishimiya et al. 299 2005; Holland et al. 2008; Can and Holland 2011, 2013; Stevens et al. 2015). In all 300 the cases mentioned above, the increased size is accompanied by an increased IBS or 301 302 the addition of multiple IBSs. Other investigators have reported that AFPs are potentiated by ligation to, or interaction with, large non-ice binding structures 303 (Deluca et al. 1998; Hakim et al. 2013; Wu and Duman 1991, Wu et al. 1991;



### 6 Thermal Hysteresis

Horwath et al. 1996; Wang and Duman 2005, 2006). In these cases, the IBS is 305 unchanged. 306

In addition to the effect of molecular size, several authors have reported that the 307 hysteresis activity is also elevated in the presence of various low-mass co-solutes. These low-mass solutes include sugars, polyols, salts, amino acids, salts of 309 polycarboxylates, and NADH. The effect has been reported for both moderately active AF(G)Ps (Kerr et al. 1985; Caple et al. 1986; Evans et al. 2007; Gong et al. 311 2011) and hyperactive AFPs (Li et al. 1998; Kristiansen et al. 2008; Amornwittawat 312 et al. 2008; Wang et al. 2009a, b; Amornwittawat et al. 2009; Wen et al. 2011; Liu 313 et al. 2015). 314

There is one thing that variations in molecular size and additives have in 315 common; they change the solubility of proteins in solution. Moreover, they reportedly enhance the hysteresis activity in manners predicted by their general effects on 317 protein solubility. In the following section, the potential importance of the solubility of AF(G)Ps to their antifreeze potency is briefly explored. 319

320

321

333

334

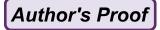
### 6.4.2 The Solubility of the AF(G)Ps: A General Concept to Explain Variability?

Several authors have in various ways implicated protein solubility as a relevant 322 factor in antifreeze potency (Kristiansen and Zachariassen 2005; Evans et al. 2007; 323 Kristiansen et al. 2008; Wang et al. 2009a). Solubility of AF(G)Ps have also 324 inadvertently been implicated in the manner the AF(G)Ps are thought to orient 325 toward the ice; these proteins are somewhat amphipathic, were the more hydrophobic side that contains the IBS orient toward the ice (Yang et al. 1988; Sönnichsen 327 et al. 1996; Haymet et al. 1998, 1999). In other words, the less soluble side of the 328 molecule orients toward the ice whereas the more soluble side orients toward the 329 water. The logical extension of this is that a less soluble AFP would have a greater 330 affinity toward the ice surface than a more soluble AFP. In the following paragraphs, 331 a brief examination of the significance of this common denominator, the solubility of 332 the AF(G)Ps, to their potency is presented.

#### 6.4.2.1 The AF(G)P/Ice Interaction Is Temperature Dependent

The ice surface in equilibrium with surrounding liquid water is not distinct but a 335 transition region where the configuration of the water molecules changes from the ordered crystal structure of the ice lattice to the random distribution of the bulk water 337 in the surrounding solution. This change occurs across a 1-2 nm deep region called 338 the interfacial region or the melting/freezing region (Hayward and Haymet 2001).

As stated in the introductory quote by Ramsay (1964), AF(G)Ps act at tempera- 340 tures below the equilibrium melting temperature of the ice, not at temperatures above 341



it, i.e., the ice crystal does not grow below this temperature but melts above it (but see also next section concerning superheating of ice crystals). This suggests that the AF(G)Ps are irreversibly adsorbed onto the ice crystal surface only at temperatures below the melting temperature. A simple explanation to this is that AF(G)Ps freeze onto the crystal surface as the temperature is lowered to within the hysteresis gap and then melt off the ice when the temperature is raised to the melting temperature (Kristiansen and Zachariassen 2005). Such a temperature-dependent behavior of freezing onto (adsorption) and melting off (desorption) would explain why ice crystals in the presence of AF(G)Ps typically melt at the equilibrium temperature irrespective of any colligative variation in this temperature. It also provides an intuitive and simple explanation to the long-standing conundrum of the origin of the necessary bond strength to achieve irreversible adsorption (Wen and Laursen 1992; Knight et al. 1993; Chao et al. 1995); the bond strength between the irrevers-ibly adsorbed AF(G)P and the ice surface corresponds to those between water molecules in bulk ice at that temperature. Recently, Garnham et al. (2011a) showed that the hydration water of a hyperactive AFP has a clathrate-like configuration and is firmly embedded by extensive H-bonds to the backbone of the protein. Hence, this crystalline-like water at the IBS appears to be prone to fuse together with the solidifying crystalline interface once the temperature is lowered and melt off when the interface disintegrates into chaos on warming to the equilibrium melting tem-perature. Molecular dynamics studies support this contention (Chakraborty and Jana 2019; Zanetti-Polzi et al. 2019).

Pertaya et al. (2008) reported on the fluorescence associated with an ice crystal in a solution containing fluorescently tagged AFP. When slowly melting a crystal at a temperature just above that of equilibrium the crystal showed no fluorescence, indicating no adsorbed AFPs. When cooled to within the hysteresis gap the crystal surface became fluorescent, indicating irreversible adsorption. Similar results were reported by Pertaya et al. (2007), who used a technique of photo-bleaching of fluorescently tagged AFPs to study the AFP/ice association at the crystal surface at temperatures within, and just above, the hysteresis gap. Bleached AFPs at the surface were not replaced within the hysteresis gap but were replaced at temperatures just above, showing that the AFPs were irreversibly adsorbed within the hysteresis gap and desorbed off the ice at the melting temperature.

While in the desorbed state, at the melting temperature of the crystal surface, there must be a distribution of AF(G)Ps between the melting/freezing region and the bulk solution. It is this distribution pattern that presumably becomes affected by changes in the solubility of the AF(G)Ps; a lowered protein solubility means that the AF(G)P has an increased tendency to move away from the solution and toward the melting/ freezing region. This results in more AF(G)P molecules being at the ice/water interfacial region and available to freeze onto the solidifying crystal surface the instant the temperature is lowered. Consequently, lowered solubility of an AF(G)P should result in greater surface density of the AF(G)P below the melting temperature and, hence, to greater hysteresis activity (Kristiansen and Zachariassen 2005).

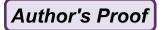
### 6 Thermal Hysteresis

**Superheating of Ice Crystals** Several investigators have reported that ice crystals 385 in solutions of AF(G)Ps may superheat slightly (Celik et al. 2010; Cziko et al. 2014). 386 Celik et al. (2010) reported that tiny ice crystals became superheated by  $0.04\,^{\circ}$ C and 387  $0.44\,^{\circ}$ C in the presence of several hyperactive AFPs. In the case of moderately active 388 AFPs, superheating up to  $0.02\,^{\circ}$ C was reported at high AFP concentrations. The 389 observed superheating reflects the presence of concave surface regions developing 390 between irreversibly adsorbed AF(G)Ps at temperatures above the equilibrium 391 temperature (Knight and DeVries 1989). These observations potentially contradict 392 the notion of an equilibrium distribution of AF(G)Ps developing between the 393 solution and the ice surface region at the equilibrium temperature, as outlined above. 394

The samples that expressed this superheating also expressed hysteresis activities 395 ranging from  $1.7 \,^{\circ}$ C to  $4.1 \,^{\circ}$ C. The hysteresis activity increases approximately as a 396 function of the square root of the surface density of AF(G)Ps (Raymond and DeVries 397 1977; Kristiansen and Zachariassen 2005). Thus, apparently only a small fraction of 398 the AFPs that was originally frozen onto the surface and caused these high hysteresis 399 activities was subsequently involved in the comparatively much lower superheating. 400 That is, most AFPs melted off the ice surface.

The superheating phenomenon requires a cooling event to occur; when Celik 402 et al. (2010) melted out ice in solutions with high concentrations of moderate AFPs 403 or low concentrations of hyperactive AFPs, they observed that the many small 404 crystals decreased uniformly in size. If the melting process was briefly halted, then 405 the crystals began to show slight superheating. This change in melting behavior 406 following a brief cooling event suggests that AFPs in solution do not adsorb 407 irreversibly to the ice surface unless there is a cooling event, i.e., the adsorption is 408 a freezing of the AFPs onto the ice surface. The subsequent desorption as the 409 temperature is raised is for some of the adsorbed AFPs a delayed process.

Why do some of the AF(G)Ps not simply melt off the surface as the temperature is 411 raised to the melting point? The freezing of the AF(G)Ps onto the ice surface imply 412 that the hydration water at the IBS becomes part of the crystal lattice. Above its 413 equilibrium melting temperature, ice melts from its surface, as lattice water molecules are released to the fluid hydrogen-bonding network of the surrounding solu- 415 tion. However, if no liquid water is in contact with the lattice that is to be melted, 416 e.g., in the interior of a crystal, the lattice structure may superheat extensively before 417 a melting nucleation event occurs (Turnbull 1950; Chalmers 1964; Lu and Li 1998). 418 Consequently, if the crystalline water at the IBS of an adsorbed AFP is shielded from 419 the surrounding liquid solution, then the melting process at the IBS is prevented and 420 the AFP will remain adsorbed onto the crystal surface at temperatures above the 421 melting point. The distinct difference in the capacities of moderately and hyperactive 422 AFPs to cause superheating reported by Celik et al. (2010) presumably reflect 423 differences in their respective capacities to shield the crystalline water at the IBS 424 from the surrounding liquid water when adsorbed onto the ice. They observed that in 425 the presence of hyperactive AFPs, crystals sporadically disappeared over time up to 426 4 h, showing that this situation can be quite stable if it develops. Since the phenom- 427 enon is very weak compared to the hysteresis activity, it might be that only those AF 428



465

(G)Ps with certain rare orientations at the crystal surface is able to postpone the initiation of the melting process at the IBS.

#### 6.4.3 Basic Concepts in Solubility Theory

The solubility of a protein in water reflects its energetic state in water (Reynolds et al. 432 1974). Once present in the water, the solubility of a protein is determined by two opposing effects acting on structural features of the protein; favorable attractive 434 forces such as van der Waals- and dipole-type forces lower the energy state of the 435 protein and therefore increase its solubility. This is opposed by an energetic cost 436 associated with occupying a cavity within the water that increases its energetic state 437 and therefore lowers its solubility (Uhlig 1937; Tolls et al. 2002). In the latter case, 438 the presence of the protein in the water effectively adds additional high-energy water 439 surface at the water/protein boundary of the cavity occupied by the solute. The 440 presence of nonpolar surface regions of the protein restricts hydrogen bond forma-441 tion between water molecules in the surface boundary, and consequently reduces the 442 freedom of these local water molecules to orientate. This structuring of water at the 443 protein/water boundary is known as the hydrophobic effect. 444

According to Uhlig (1937), the solubility (S) of a dissolved molecule may be 445 expressed as: 446

$$RT \ln(S) = -A\gamma + E \tag{6.2}$$

where R and T are the universal gas constant and the absolute temperature, respectively. The first term on the right side of Eq. (6.2),  $A\gamma$ , represents the "hydrophobic" effect that lowers the solubility of a molecule. This effect is a function of the nonpolar surface area, A, of the molecule in contact with water, and the energetic 450 state of the water at this surface, expressed as the water surface tension,  $\gamma$ . This 451 hydrophobic effect is opposed by the second term on the right-hand side of Eq. (6.2), 452 the favorable "electrostatic" effect, E, that raises the solubility of the dissolved 453 454 molecule (Reynolds et al. 1974; Melander and Horváth 1977).

Changing the size of the AF(G)Ps, for instance by adding or removing repetitive 455 peptide segments, inadvertently also changes the nonpolar surface area, A, of the 456 protein and consequently its solubility. Also, for structurally similar isoforms of 457 different size, their nonpolar surface areas, and hence, their solubility, correlate with 458 459 their size. The small mass solutes that reportedly enhance the hysteresis activity, such as salts, sugars, polyols, and amino acids are known to elevate the surface 460 tension, γ, of water (Washburn 1929; Melander and Horváth 1977; Kaushik and Bhat 1998; Landt 1931; Matubayasi and Nishiyama 2006; Bull and Breese 1974). 462 Thus, their reported enhancement effect may simply be the result of the solutes 463 464 lowering the solubility of the AF(G)P by elevating  $\gamma$  at the protein/water interface.

The basic framework outlined above may be useful when interpreting natural and induced variations in the antifreeze potency among moderately active and among



### 6 Thermal Hysteresis

hyperactive AF(G)Ps. In the following paragraphs, standard solubility theory will be 467 applied to examine some of the reported effects small co-solutes and variations in 468 size have on hysteresis activity. 469

## 6.4.4 Low-Mass Additives, Solubility, and Antifreeze Potency

The effects of salts on the hysteresis activity in relation to solubility theory will be 471 exemplified by the effects of salts on the hyperactive AFP, RiAFP, from the 472 cerambycid beetle *Rhagium inquisitor* (Kristiansen et al. 2008). Wang et al. 473 (2009a) also had a quite similar approach to this issue. It will be shown that these 474 effects are entirely consistent with being caused by salt-induced lowered solubility 475 of the RiAFP molecules. To support this claim, the nonpolar surface area and the 476 dipole moment of RiAFP is derived from the effects of salts on its antifreeze 477 potency.

## 6.4.4.1 The Salting-Out Constant, $K_s$

As mentioned above, salts are known to lower the solubility of proteins. This effect 480 is termed "salting-out." The salting-out effect is qualitatively similar for different 481 kinds of proteins and different kinds of salts in that the solubility of the protein 482 changes in a log-linear manner with the concentration of salt (Cohn 1925; Melander 483 and Horváth 1977):

$$\ln(S) = \beta - K_s m$$
(6.3)

470

479

491

492

where S is the solubility of the protein (mg/ml),  $\beta$  is the solubility of the protein in the 485 absence of salts (mg/ml), m is the concentration of the salt (molal), and  $K_s$  is known 486 as the salting-out constant (molal<sup>-1</sup>).  $K_s$  is an expression of the sensitivity of the 487 solubility of a particular protein to the presence of a particular salt. The value of  $K_s$  488 depends on both the salt and the protein and is experimentally determined as the 489 slope of the linear relationship between  $\ln(S)$  and m.

## 6.4.4.2 Obtaining Salting-Out Constants from Measurements of Hysteresis Activity

Since the presence of salts increases the hysteresis activity, adding salts is equivalent 493 to increasing the concentration of the AF(G)P. Since solubility is in units of 494 concentration, Eq. (6.3) should describe the salt-induced apparent changes in the 495 concentration of AF(G)P. Thus, the salting-out constant,  $K_s$ , in the presence of a 496 particular salt may be obtained from the hysteresis measurements as follows; the 497 actual concentration of AF(G)P in the samples is kept unchanged during the 498



procedure. An "apparent" concentration of AF(G)P in the presence of different concentrations of salts is then obtained by converting the observed enhanced hysteresis activity in the presence of salts to the equivalent concentration of AF(G)P needed to cause this activity in the absence of salt. The value of  $K_s$  for that salt is then obtained simply as the slope of the linear relationship obtained by plotting the natural logarithm of the "apparent" concentration of AF(G)P in the samples as a function of the concentration of salt.

Kristiansen et al. (2008) used this method to determine  $K_s$  for each of ten different salts from the salt-induced enhancement of the hysteresis activity for RiAFP. As predicted by Eq. (6.3) all "apparent" concentrations were log-linear functions of the concentrations of the different salts tested.

## 10 6.4.4.3 The Hofmeister Series and Its Linearity

An experimentally determined salting-out constant,  $K_s$ , is an expression of the two opposing effects acting on structural features of the protein, the favorable "electrostatic" effect that increases the solubility of the protein and the unfavorable "hydrophobic" effect that lowers its solubility, as outlined in Eq. (6.2). The net observed salting-out constant,  $K_s$ , in Eq. (6.3) is given by (Melander and Horváth):

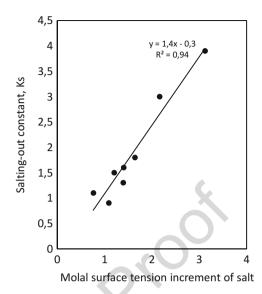
$$K_s = \Omega \sigma - \Lambda \tag{6.4}$$

where  $\Omega$  is a protein-specific intrinsic salting-out constant (cm dyn<sup>-1</sup>),  $\sigma$  is the molal surface tension increment of the salt ( $10^{-3}$  dyn g/cm mol), and  $\Lambda$  is a protein-specific intrinsic salting-in constant (molal<sup>-1</sup>).

By arranging salts according to their ability to lower the solubility of proteins, the 519 so-called Hofmeister series of salts is obtained. The arrangement of different salts in 520 the Hofmeister series may be understood from Eq. (6.4); for a specific protein, the 521 molal surface tension increment,  $\sigma$ , is the only variable in the equation. Thus, for any 522 single protein the arrangement of salts according to their ability to lower the 523 solubility of that protein is similar for all proteins and dictated by the molal surface 525 tension increment,  $\sigma$ , of the different salts. For example, the following eight salts listed in descending order according to their ability to lower protein solubility form 526 series as (value for σ in parenthesis): 527  $(NH_4)_2SO_4$  (2.16) > NaCl (1.64) > KCl (1.40) > NH<sub>4</sub>Cl (3.12) >528  $(1.39) > \text{NaNO}_3$  (1.20) > NaI,  $(1.09) > \text{N(CH}_3)_4\text{Cl}$  (0.76). Since the value of  $\sigma$  is 529 530 actually the surface tension increment of the water/air interface and not the protein/ water interface, there are slight differences between the predicted and observed 531 Hofmeister series. However, this general arrangement of salts varies little for 532 different proteins. Hence, if the salt-induced enhancement of the hysteresis activity 533 is caused by salt-induced reduction in the solubility of RiAFP, then the 534 535 enhancement-effect of the different salts should reflect the Hofmeister series. In the case of RiAFP the experimentally determined  $K_s$  values were arranged as (value 536 for  $\sigma$  in parenthesis): Na<sub>3</sub>C<sub>6</sub>H<sub>6</sub>O<sub>7</sub> (3.12) > (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> (2.16) > NaCl (1.64) > KCl

### 6 Thermal Hysteresis

Fig. 6.5 The linearity of the Hofmeister series. Different salting-out constants,  $K_s$ , determined from the hysteresis activity in the presence of different kinds of salts, versus the surface tension increment of the salts used. The slope of the linear line is the intrinsic salting-out constant,  $\Omega$ , for the protein. The intercept value is the intrinsic saltingin constant,  $\Lambda$ , for the protein. These two proteinspecific constants may be used to determine the nonpolar surface area and the dipole moment of the protein. Adapted from Kristiansen et al. (2008)

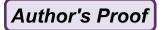


552

559

 $(1.40) > \text{NaNO}_3 (1.20) > \text{NH}_4\text{Cl} (1.39) > \text{N(CH}_3)_4\text{Cl} (0.76) > \text{NaI} (1.09)$ . This 538 arrangement is in close agreement with the Hofmeister series. An examination of the 539 results of Wang et al. (2009a) also shows that arrangement of the salts according to 540 their ability to enhance a hyperactive AFP, DAFP, from the beetle Dendroides 541 canadensis corresponds well with the Hofmeister series. For those salts where the 542 value for  $\sigma$  is known, they are listed as (value for  $\sigma$  in parenthesis); NaCl 543 (1.64) > KCI (1.40) > KBr (1.31) > NaBr (1.32) > KI (0.84) > NaI 544 $(1.09) > \text{NaClO}_4$  (0.55). In their extensive study, Li et al. (1998) found that, 545 among all the different compounds tested, citrate was the strongest enhancer of the 546 antifreeze potency of DAFP. Citrate has among the highest known surface tension 547 increments ( $\sigma$  of 3.12) and was also the strongest enhancer in the case of RiAFP. 548 Evans et al. (2007) did not find differences in the efficacies of LiCl, NaCl, and KCl to 549 enhance the antifreeze activity of different kinds of fish AF(G)Ps. This is consistent 550 with the fact that the molal surface tension increments of these salts are very similar, 551 1.63, 1.64, and 1.40, respectively.

Since both  $\Omega$  and  $\Lambda$  of Eq. (6.4) are constant features of the protein, the  $K_s$  values 553 obtained for that protein will be a linear function of the molal surface tension increment,  $\sigma$ , of the different salts. Figure 6.5 shows that the salting-out constants 555 of RiAFP, determined from the salt-induced enhancement of the hysteresis activity, vary as a linear function of  $\sigma$ , consistent with this prediction of Eq. (6.4). Thus, the 557 linear relationship depicted in Fig. 6.5 is a quantitative representation of the 558 Hofmeister series for RiAFP.



## 6.4.4.4 Quantitative Predictions of Protein Properties from Salt-Induced Enhancement

The protein-specific salting-out constant,  $\Omega$ , and the protein-specific salting-in constant,  $\Lambda$ , of Eq. (6.4) reflect physicochemical properties of the protein (Melander and Horváth 1977). Thus, if the concept of solubility-induced enhancement is correct, then it should be possible to use the information derived from the antifreeze measurements to predict features of the protein that are reflected by these constants.

The protein-specific salting-out constant,  $\Omega$ , of Eq. (6.4) represents the hydrophobic properties of the protein and is a function of its nonpolar surface area,  $\phi$ . The numeric value of  $\phi$ , in units of square Ångstrøm, may be obtained from  $\Omega$  as (Melander and Horváth 1977):

$$\phi = 411\Omega - 12\tag{6.5}$$

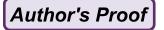
According to Eq. (6.4),  $\Omega$  is given by the slope of the linear relationship depicted in Fig. 6.5. Using the value of 1.4 cm dyn<sup>-1</sup> for  $\Omega$  in Eq. (6.5) gives a value for  $\phi$  for RiAFP of 563 Å<sup>2</sup>. This is about 20% of the total surface area of the protein (Kristiansen et al. 2008). According to Melander and Horváth (1977),  $\phi$  is typically between 20% and 40% of the total surface area of proteins. The sensitivity of RiAFP to become enhanced by salts therefore seems to correspond well with the expected salt sensitivity of a protein of its size.

The intrinsic protein salting-in constant,  $\Lambda$ , of Eq. (6.4) reflects the favorable electrostatic forces acting to enhance the solubility of the protein and is a function of its dipole moment,  $\mu$ . The dipole moment,  $\mu$ , may be numerically obtained in units of Debye from  $\Lambda$  using the formula (Melander and Horváth 1977):

$$\mu = -578\Lambda \tag{6.6}$$

According to Eq. (6.4), the value of  $\Lambda$  is given by the intercept of the linear relationship depicted in Fig. 6.5 and has the value of  $-0.3 \text{ molal}^{-1}$ . This gives a predicted dipole moment for RiAFP of 173 Debye. Since the original study was published (Kristiansen et al. 2008), the crystal structure of RiAFP has become available (Hakim et al. 2013). The structure file (PDB 4DT5) contains two molecules, A and B, which, when submitted to the online Protein Dipole Moments Server (Felder et al. 2007) has predicted dipole moments of 182 Debye and 125 Debye, respectively. It is noteworthy that the dipole moment of RiAFP, derived from its molecular structure, coincides within a few percentage points with the dipole moment derived from the effects of salts on the antifreeze potency of the protein.

Considering the above presented relations, it appears obvious that salts enhance the antifreeze potency by lowering the solubility of AF(G)Ps. Since the other small mass solutes known to enhance the antifreeze potency of AF(G)P, i.e., polyols, amino acids, sugars etc., act on protein solubility in a manner similar to that of salts, they are all likely to operate by the same mechanism.



### 6 Thermal Hysteresis

## Molecular Size, Solubility, and Antifreeze Potency

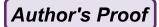
Several explanations are provided for the effect of size on the potency of AF(G)Ps. 598 In those cases where the IBS does not vary with the size of the protein, the size effect 599 is ascribed to the larger AFP-macromolecule complex covering a larger surface area 600 than the AFP alone. This larger coverage effectively reduces the intermolecular 601 adsorbent gap between adsorbed AFPs at the ice surface, thereby displacing the 602 hysteresis freezing point to a lower temperature (Wu et al. 1991). When the variation 603 in molecular size of the protein involves changes in the size of the IBS, then the 604 effect has additionally been ascribed to various aspects of their ice-binding ability (Leinala et al. 2002; Mok et al. 2010; Chao et al. 1996; Liu et al. 2005). The 606 increased potency reported for a natural and several synthetic intramolecular 607 multimers of AFPs is ascribed to an overall greater likelihood of successful adsorption due to the presence of multiple IBSs (Miura et al. 2001; Nishimiya et al. 2003) 609 or to increased overall ice-binding area (Baardsnes et al. 2003).

Although some, or even all, of these explanations may contribute to some extent 611 to the observed effect, there are nevertheless problems associated with their applicability. For instance, Marshall et al. (2004b) pointed out that, explanations relying 613 on differences in interaction energies at the IBS are not likely to be correct, since AF 614 (G)Ps are irreversibly adsorbed onto the ice surface, i.e., it is an all-or-none situation. 615 As alluded to above (Sect. 6.4.2), if the AF(G)Ps become irreversibly adsorbed by 616 freezing onto the interface, then they are as strongly adsorbed to the ice as any piece 617 of ice is to the surface of ice. Thus, changing the size of the IBS, or the like, should 618 not make any difference. In the case of the added surface cover explanation provided 619 by Wu et al. (1991), it is intuitively logical and could well be a satisfactory 620 explanation. However, as pointed out by the original authors, experimentally there 621 is no correlation between the size of the enhancer and the enhancement effect 622 (Wu and Duman 1991). The enhancers, identified by Wu and Duman (1991), 623 range according to efficiency as 70 kDa (endogen enhancer) > 70 kDa (protein ice 624 nucleator) > 800 kDa (lipoprotein ice nucleator) > 150 kDa (antibody) > gelatin 625 (80–375 kDa) > agar (average 120 kDa). The effectiveness of all these enhancers is 626 surpassed by a 28 kDa endogenous enhancer (Wang and Duman 2006). Also, 627 Horwath et al. (1996) reported that an efficient endogenous enhancer from the beetle 628 Tenebrio molitor was 12 kDa, about the same size as the AFP. Thus, there seem to be 629 little experimental support for the otherwise logical contention that the enhancement 630 effect of size arises from added surface cover of the adsorbent complex. 631

Equation (6.2) provides a general explanation to the size effect; variations in size 632 is inevitably accompanied by variations in the nonpolar surface area, A, of the 633 protein and probably also variations in the electrostatic forces, E, acting between 634 the protein and the solution. Such size-induced differences in solubility is consistent 635 with the gradual increase in antifreeze potency with size that are reported for 636 structurally similar variants of both hyperactive and moderately active AF(G)Ps. This approach also provides an explanation as to why there is no correlation between 638 antifreeze potency and size for macromolecules that are very different; if the 639

597

610



structures are different, then differences in their nonpolar surface areas and the strength of the electrostatic forces acting between the structure and the surrounding water do not vary with size. In other words, the solubility of structurally different compounds does not vary with molecular size. This would explain why a 28 kDa protein is a far more efficient enhancer than a protein of 800 kDa; the smaller is simply less soluble.

Some complicated and intriguing findings have been reported that ties in well with the concept of solubility-induced enhancement; Wang and Duman (2005) found that certain of the isoforms of hyperactive AFPs, DAFPs, from D. canadensis interact, and the association results in greater activities. This greater activity may be ascribed to a reduced solubility due to the overall larger nonpolar surface area,  $\phi$ , of the complex (Eqs. 6.2, 6.4 and 6.5). But further, they found that the additive glycerol only acted as an enhancer if the isoforms interacted. This may also be understood from Eqs. (6.4) and (6.5); the sensitivity of a protein to some additive increases with increased nonpolar surface area,  $\phi$ . It should be noted that several of the polyols, glycerol included, actually reduces the surface tension of water. Nevertheless, Gekko and Timasheff (1981) found that glycerol lowered the solubility of proteins by the same mechanism as salts, i.e., polyols act differently at the air/water interface than at the protein/water interface.

Amornwittawat et al. (2008) found that many carboxylates enhanced DAFPs and ascribed the effect to aggregation of DAFPs. As in the case with Wang and Duman (2005), such aggregation results in lowered solubility due to increased overall nonpolar surface area, which could explain the increased activity. Wang et al. (2009b) identified the binding sites for these carboxylates to be specific arginine residues in the DAFP structure, since blocking these residues abolished the effect. With intact such residues the monomeric DAFP aggregated in the presence of carboxylates and the complex was more sensitive to other additives. This situation is similar to that of Wang and Duman (2005) described above, i.e., the greater nonpolar surface area,  $\phi$ , of the complex makes the overall complex more sensitive to additives than the monomers alone (Eqs. 6.2, 6.4 and 6.5).

As have been outlined above, ascribing variations in antifreeze potency to variations in protein solubility explain many aspects of hysteresis activity, including the significance of size and how additives enhance AF(G)Ps. This approach also explains why interactions between isoforms cause enhancement and the increased sensitivity to additives when isoforms interact. Ascribing variability of antifreeze potency to variations in protein solubility give a plausible explanation to the natural variability reported among AF(G)Ps that are either hyperactive or moderately active. Both qualitative and quantitative agreements with predictions based on established theory support this approach.

The presence of a 1 molal solution of sodium citrate has the effect on the hysteresis activity of RiAFP equivalent to elevating its concentration 50-fold (Kristiansen et al. 2008). Thus, aside from the categorization into hyperactive and moderately active, which are consequences of structural aspects of their IBS, the physicochemical property of solubility is probably the most dominant determinant of AF(G)P potency.



6 Thermal Hysteresis

#### 6.5 AF(G)Ps and Ice Nucleation

The convex surface zones that grow out at the ice surface within the hysteresis gap 686 are developing ice nuclei with their critical radius at that temperature. At the 687 hysteresis freezing point, the phenomenon is terminated by a surface nucleation 688 event, as one of these surface nuclei initiates nucleation. Apart from causing thermal 689 hysteresis by controlling the development of nuclei at the ice surface, AFPs also 690 interact with structures in the body fluids that can trigger an ice nucleation event. 691 Such a structure is known as an ice-nucleating agent, INA, and the nucleation 692 process triggered by INAs is referred to as heterogeneous nucleation. This is to 693 distinguish this kind of nucleation from that which occur by spontaneous ordering of 694 water, so-called homogenous nucleation. Evidence suggest that the ice nucleation 695 sites of INAs are structurally related to the IBSs of AFPs. Thus, the mechanism of ice 696 nucleation by INAs may be very similar to the mechanism of adsorption of AFPs 697 to ice. 698

#### 6.5.1 Biological Relevance of INAs

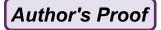
Freeze-avoiding species die if their body fluids freeze out. Consequently, they rely 700 on extensive supercooling of their body fluids to survive subfreezing temperatures. 701 Any incidental INAs in the body fluids of such an organism would therefore be 702 potentially lethal. Freeze-avoiding insects are known to remove or reduce the 703 amount of such incidental INAs that could pose a threat (Neven et al. 1986; Olsen 704 and Duman 1997a, b). In addition, AFPs prevent incidental INAs from initiating 705 freezing by physically interacting with such structures (Olsen and Duman 1997a, b; 706 Duman 2002). By removal of INAs from their body fluids and by producing high 707 concentrations of AFPs, the supercooling points of freeze avoiding larvae of the 708 pychroid beetle *Dendroides canadensis* changes from about -7 °C in the summer to 709 below -30 °C during winter (Olsen and Duman 1997a, b).

Freeze-tolerant species, that adaptively allow their body fluids to freeze out, often 711 produce INAs and allocate them to the extracellular fluid. The principal function of 712 such adaptive INAs in freeze tolerance is to prevent harmful cellular freezing by 713 initiating a preemptive nucleation event outside the cells at a temperature above the 714 nucleation temperature of any incidental harmful cellular INAs (Zachariassen and 715 Hammel 1976). Since solutes are excluded from the growing ice mass, the extracel-716 lular freezing event causes the remaining unfrozen extracellular fluid fraction to 717 become increasingly concentrated. This in turn initiates a concomitant osmotic 718 efflux of water out from the cells. The extracellular freezing process and consequent 719 efflux of cell water continues until the melting point of the remaining unfrozen fluid 720 fraction is colligatively depressed to the environmental temperature, at which point 721 the danger of harmful cellular freezing is eliminated.

685

699

722

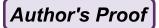


## 723 6.5.2 Overall Structural Aspects of INAs

It is vital for the functionality of adaptive INAs found in freeze-tolerant species that their nucleation temperature is above that of any incidental harmful cellular INAs. The efficiency of INAs to initiate nucleation depends on their size. This may be understood from Eq. (6.1); the larger the diameter of the INA, d, the less supercooling,  $\Delta T$ , is required to initiate nucleation. Consistent with this, adaptive INAs found in freeze tolerant species are very large structures. It is likely that the great potency of biologically adaptive INAs results from association between monomeric INA molecules; it has been shown that an adaptive 800 kDa INA from freeze-tolerant larvae of the cranefly, *Tipula trivittata*, form long chains of 800 kDa monomers, akin to pearls-on-a-string, and that two such chains align side by side into extended dimers (Yeung et al. 1991). This association apparently relies on the presence of phosphatidylinositol, PI, at the surface of the INA, as enzymatic removal of PI depressed the nucleation temperature (Neven et al. 1989). PI has also been shown to anchor highly active bacterial INAs to the bacterial membrane (Kozloff et al. 1991) and thereby possibly causing them to cooperate. The proposed structure of a large repetitive segment of the 123 kDa INA from Pseudonomas borealis suggests that the operating INA consists of at least two monomers (Garnham et al. 2011b).

AFPs are known to physically interact with INA molecules (Wu and Duman 1991). A simple explanation to how AFPs depress the nucleation temperature of INAs would be if they act by preventing them from forming larger associations, analogous to the effect of reducing the diameter of growing surface nuclei at the ice surface (Eq. 6.1). A peculiar aspect of this AFP/INA association is that it apparently does not involve the IBS of the AFP (Duman 2001). This is evident from the fact that the hysteresis activity, which requires the IBS to be free to adsorb onto the ice surface, is enhanced by the AFP/INA interaction (Wu and Duman 1991). It remains unclear if AFPs contain some secondary functional surface-site outside the IBS dedicated to the interaction of structures other than ice (Duman 2001).

Although the details of how INAs trigger freezing is not entirely identified, it is likely that they do so by structuring their hydration water to mimic that of ice. It has been shown that the hydration water at the IBS of  $\beta$ -helical hyperactive AFPs are clathrate-like, and this structured water has been implicated in the process of adsorption (Garnham et al. 2011a). Large internal repetitive parts of several bacterial INAs have been modeled to fold into  $\beta$ -helixes (Graether and Jia 2001; Garnham et al. 2011b). The structural similarity between the IBS of the  $\beta$ -helical AFPs and the suspected nucleation sites of the INAs suggest they share a similar mode of operation. Supporting this contention, Kobashigawa et al. (2005) reported that a recombinant protein corresponding to an internal part of one of these bacterial INAs shape ice crystals into hexagonal bipyramids. Similar results were also reported by Xu et al. (1998), who found that a 164 kDa molecule with INA activity shaped ice crystals into hexagonal bipyramids. Apparently, these bacterial INAs have some kind of internal IBS. It is not clear if the part of the INA responsible for the observed



### 6 Thermal Hysteresis

structuring of the ice, an IBS, corresponds to the site that causes nucleation. Another 766 aspect is the shape of the ice crystals in the presence of the INAs reported by 767 Kobashigawa et al. (2005); these INAs are those of the species Pseudonomas 768 syringae, the same INA modeled as a β-helix by Graether and Jia (2001). The ice 769 crystals in the presence of all known β-helical AFPs express multiple ice crystal 770 planes, e.g., in the form of hexagonal discs. These INAs, on the other hand, shape ice 771 into hexagonal bipyramids, as seen in the presence of the monoplane-specific AF(G) 772 Ps of fish.

If the IBS of AF(G)Ps is structurally comparable to the nucleation sites of INAs, 774 then why are AF(G)Ps not INAs? The explanation may in part rely on differences in 775 the structure of the hydration water at the IBS/nucleation site and in part be due to the 776 large difference in size of AFPs and INAs. What is clear is that β-helical insect AFPs 777 do not act as INAs within the supercooling range of the freeze avoiding insects, i.e., 778 down to about -30 °C, or even below.

773

790

801

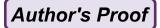
6.6 **Conclusions** 780

This chapter has dealt with the modus operandi of AF(G)Ps. The characteristic 781 prevention of ice growth within the hysteresis gap is explained by ice/water vapor 782 pressure equilibrium being maintained by the Kelvin effect as the ice surface grows 783 out as microscopic curvatures between adsorbed AF(G)Ps. The different potencies 784 of moderately and hyperactive AF(G)Ps are ascribed to differences in their adsorp-785 tion habits, whereas variations in antifreeze potencies within each of these categories 786 are ascribed to variations in their solubilities. In the latter case, experimental proof of 787 concept is discussed in the context of basic solubility theory. Some characteristics of 788 ice-nucleating agents (INAs) in relation to AF(G)Ps and their relevance in cold 789 tolerance was also briefly examined.

AF(G)Ps as a group are defined by their shared capacity to prevent ice in solution 791 from growing at temperatures below the melting point. However, another wide- 792 spread trait observed for many of these proteins when at very low concentrations 793 occurs at the melting temperature; they inhibit the spontaneous process by which 794 larger ice crystals grow at the expense of smaller crystals. This trait is not an 795 exclusive property of AF(G)Ps but are also found among non-antifreeze proteins 796 and organic solutes. This fascinating phenomenon of recrystallisation inhibition is 797 both biologically and commercially important and is the topic of the next chapter. 798

References 799

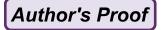
Amornwittawat N, Wang S, Duman JG, Wen X (2008) Polycarboxylates enhance beetle antifreeze 800 protein activity. Biochim Biophys Acta 1784:1942-1948



- Amornwittawat N, Wang S, Banatlao J, Chung M, Velasco E, Duman JG, Wen X (2009) Effects of
   polyhydroxy compounds on beetle antifreeze protein activity. Biochim Biophys Acta
   1794:341–346
- Baardsnes J, Kuiper MJ, Davies PL (2003) Antifreeze protein dimer. When two ice binding faces
   are better than one. J Biol Chem 278:38942–38947
- 807 Bull HB, Breese K (1974) Surface tension of amino acid solutions: a hydrophobicity scale of amino
   808 acid residues. Arch Biochem Biophys 161:665–670
- 809 Can O, Holland NB (2011) Conjugation of type I antifreeze protein to polyallylamine increases 810 thermal hysteresis activity. Bioconjug Chem 22:2166–2171
- 811 Can O, Holland NB (2013) Utilizing avidity to improve antifreeze protein activity: a type III 812 antifreeze protein trimer exhibits increased thermal hysteresis activity. Biochemistry 813 52:8745–8752
- Caple G, Kerr WL, Burcham TS, Osuga DT, Yeh Y, Feeney RE (1986) Superadditive effects in
   mixtures of fish antifreeze glycoproteins and polyalcohols or surfactants. J Colloid Interface Sci
   111:299–304
- 817 Celik Y, Graham LA, Mok Y-F, Bar M, Davies PL, Braslavsky I (2010) Superheating of ice crystals
   818 in antifreeze protein solutions. Proc Natl Acad Sci 107:5423–5428
- 819 Celik Y, Drori R, Pertaya-Braun N, Altan A, Barton T, Bar-Dolev M, Groisman A, Davies PL,
   820 Braslavsky I (2013) Microfluidic experiments reveal that antifreeze proteins bound to ice
   821 crystals suffice to prevent their growth. Proc Natl Acad Sci 110:1309–1314
- 822 Chakraborty S, Jana B (2019) Ordered hydration layer mediated ice adsorption of a globular
   823 antifreeze protein: mechanistic insight. Phys Chem Chem Phys 21:19298–19310
- 824 Chalmers B (1964) Principles of solidification. Wiley, New York
- 825 Chao H, DeLuca CL, Davies PL (1995) Mixing antifreeze protein types changes ice crystal
   826 morphology without affecting antifreeze activity. FEBS Lett 357:183–186
- 827 Chao H, Hodges RS, Kay CM, Gauthier SY, Davies PL (1996) A natural variant of Type I
   828 antifreeze protein with four ice-binding repeats is a particularly potent antifreeze. Protein Sci
   829 5:1150-1155
- 830 Cohn EJ (1925) The physical chemistry of the proteins. Physiol Rev 5:349–437
- 831 Cziko PA, DeVries AL, Evans CW, Cheng C-HC (2014) Antifreeze protein-induced superheating
   832 of ice inside Antarctic notothenioid fishes inhibits melting during summer warming. Proc Natl
   833 Acad Sci 111:14583–14588
- 834 DeLuca CI, Comley R, Davies PL (1998) Antifreeze proteins bind independently to ice. Biophys J
   835 74:1502–1508
- 836 DeVries AL (1971) Glycoproteins as biological antifreeze agents in Antarctic fishes. Science 837 172:1152–1155
- 838 DeVries AL (1982) Biological antifreeze agents in Coldwater fishes. Comp Biochem Physiol A
   839 73:627-640
- B40 Drori R, Davies PL, Braslavsky I (2015) Experimental correlation between thermal hysteresis
   activity and the distance between antifreeze proteins on an ice surface. RSC Adv 5:7848–7853
- B42 Duman JG (2001) Antifreeze and ice nucleator proteins in terrestrial arthropods. Annu Rev Physiol
   63:327–357
- Buman JG (2002) The inhibition of ice nucleators by insect antifreeze proteins is enhanced by
   glycerol and citrate. J Comp Physiol B 172:163–168
- Buman JG, Bennett V, Sformo T, Hochstrasser R, Barnes BM (2004) Antifreeze proteins in
   Alaskan insects and spiders. J Insect Physiol 50:259–266
- 848 Evans PE, Hobbs RS, Goddard SV, Fletcher GL (2007) The importance of dissolved salts to the 849 *in vivo* efficacy of antifreeze proteins. Comp Biochem Physiol A 148:556–561
- 850 Felder CE, Prilusky J, Silman I, Sussman JL (2007) A server and database for dipole moments of 851 proteins. Nucleic Acids Res 35:W512–W521. http://dipole.weizmann.ac.il/
- 852 Fletcher GL, Hew CL, Davies PL (2001) Antifreeze proteins in teleost fishes. Annu Rev Physiol
   853 63:359–390

## 6 Thermal Hysteresis

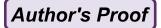
Friis DS, Kristiansen E, von Solms N, Ramløv H (2014) Antifreeze activity enhancement by site	854
directed mutagenesis on an antifreeze protein from the beetle Rhagium mordax. FEBS Lett	855
588:1767–1772	856
Garnham CP, Campbell RL, Davies PL (2011a) Anchored clathrate waters bind antifreeze proteins	857
to ice. Proc Natl Acad Sci 108:7363–7367	858
Garnham CP, Campbell RL, Walker VK, Davies PL (2011b) Novel dimeric $\beta\text{-helical}$ model of an	859
ice nucleation protein with bridged active sites. BMC Struct Biol 11:36	860
Gekko K, Timasheff SN (1981) Mechanism of protein stabilization by glycerol: preferential	861
hydration in glycerol-water mixtures. Biochemistry 20:4667–4676	862
Gong HS, Croft K, Driedzic WR, Ewart VK (2011) Chemical chaperoning action of glycerol on the antifreeze protein of rainbow smelt. J Therm Biol 36:78–83	863 864
Graether SP, Jia Z (2001) Modeling <i>Pseudomonas syringae</i> ice-nucleation protein as a β-helical protein. Biophys J 80:1169–1173	865 866
Graether SP, Kuiper MJ, Gagné SM, Walker VK, Jia Z, Sykes BD, Davies PL (2000) β-helix	867
structure and ice-binding properties of a hyperactive antifreeze protein from an insect. Nature 406:325–328	868 869
Graham LA, Davies PL (2005) Glycine-rich antifreeze proteins from snow fleas. Science 310:461	870
Grandum S, Yabe A, Nakagomi K, Tanaka M, Takemura F, Kobayashi Y, Frivik P-E (1999)	871
Analysis of ice crystal growth for a crystal surface containing adsorbed antifreeze proteins. J Cryst Growth 205:382–390	872 873
Hakim A, Nguyen JB, Basu K, Zhu DF, Thakral D, Davies PL, Isaacs FJ, Modis Y, Meng W (2013)	874
Crystal structure of an insect antifreeze protein and its implications for ice binding. J Biol Chem 288:12295–12304	875 876
Hansen TN, Baust JG (1988) Serial dilution of Tenebrio molitor haemolymph: analysis of anti-	877
freeze activity by differential scanning calorimetry. Cryo-Letters 9:386–391	878
Hansen TN, DeVries AL, Baust JG (1991) Calorimetric analysis of antifreeze glycoproteins of the	879
polar fish, Dissostichus mawsoni. Biochim Biophys Acta 1079:169–173	880
Haymet ADJ, Ward LG, Harding MM, Knight CA (1998) Valine substituted winter flounder	881
'antifreeze': preservation of ice growth hysteresis. FEBS Lett 430:301-306	882
Haymet ADJ, Ward LG, Harding MM (1999) Winter flounder "antifreeze" proteins: synthesis and	883
ice growth inhibition of analogues that probe the relative importance of hydrophobic and	884
hydrogen-bonding interactions. J Am Chem Soc 121:941-948	885
Hayward JA, Haymet ADJ (2001) The ice/water interface: molecular dynamics simulations of the	886
basal, prism, 2021 and 2110 interfaces of ice Ih. J Chem Phys 114:3713-3726	887
Holland NB, Nishimiya Y, Tsuda S, Sönnichsen FD (2008) Two domains of RD3 antifreeze protein	888
diffuse independently. Biochemistry 47:3955–3941	889
Horwath KL, Easton CM, Poggioli GJ Jr, Myers K, Schnorr IL (1996) Tracking the profile of a	890
specific antifreeze protein and its contribution to the thermal hysteresis activity in cold hardy	891
insects. Eur J Entomol 93:419–433	892
Jia Z, Davies PL (2002) Antifreeze proteins: an unusual receptor-ligand interaction. Trends Biochem Sci 27:101–106	893 894
Kaushik JK, Bhat R (1998) Thermal stability of proteins in aqueous polyol solutions: role of the	895
surface tension of water in the stabilizing effect of polyols. J Phys Chem B 102:7058-7066	896
Kerr WL, Burcham TS, Osuga DT, Yeh Y, Feeney RE (1985) Synergistic depression of the freezing	897
temperature in solutions of polyhydroxy compounds and antifreeze glycoproteins. Cryo-Letters	898
6:107–114	899
Knight CA, DeVries AL (1988) The prevention of ice crystal growth from water by "antifreeze	900
proteins". In: Wagner PE, Vali G (eds) Atmospheric aerosols and nucleation. Lecture notes in	901
physics 309. Springer, Berlin	902
Knight CA, DeVries AL (1989) Melting inhibition and superheating of ice by an antifreeze	903
glycopeptide. Science 245:505–507	904
Knight CA, Cheng CC, DeVries AL (1991) Adsorption of α-helical antifreeze peptides on specific	905
ice crystal surface planes. Biophys J 59:409–418	906



- 907 Knight CA, Driggers E, DeVries AL (1993) Adsorption to ice of fish antifreeze glycopeptides 7 and
   908 8. Biophys J 64:252–259
- 909 Kobashigawa Y, Nishimiya Y, Miura K, Ohgiya S, Miura A, Tsuda S (2005) A part of ice 910 nucleation protein exhibits the ice-binding ability. FEBS Lett 579:1493–1497
- 911 Kozloff LM, Turner MA, Arellano F, Lute M (1991) Phosphatidylinositol, a phospholipid of 912 ice-nucleating bacteria. J Bacteriol 173:2053–2060
- 913 Kristiansen E, Zachariassen KE (2005) The mechanism by which fish antifreeze proteins cause 914 thermal hysteresis. Cryobiology 51:262–280
- 915 Kristiansen E, Pedersen SA, Zachariassen KE (2008) Salt-induced enhancement of antifreeze 916 protein activity: a salting-out effect. Cryobiology 57:122–129
- 917 Landt E (1931) The surface tension of solutions of various sugars. Z Ver Dtsch Zucher-Ind 918 81:119–124
- 919 Laursen RA, Wen D, Knight CA (1994) Enantioselective adsorption of the D- and L-forms of an 920 α-helical antifreeze polypeptide to the {2021} planes of ice. J Am Chem Soc 116:12057–12058
- 921 Leinala EK, Davies PL, Doucet D, Tyshenko MG, Walker VK, Jia Z (2002) β-Helical antifreeze
   922 protein isoform with increased activity. J Biol Chem 277:33349–33352
- 923 Li N, Andorfer C, Duman JG (1998) Enhancement of insect antifreeze protein activity by solutes of
   924 low molecular mass. J Exp Biol 201:2243–2251
- 925 Liou YC, Tocilj A, Davies PL, Jia Z (2000) Mimicry of ice structure by surface hydroxyls and water
   926 of a beta-helix antifreeze protein. Nature 406:322–324
- 927 Liu K, Jia Z, Chen G, Tung C, Liu R (2005) Systematic size study of an insect antifreeze protein and
   928 its interaction with ice. Biophys J 88:953–958
- 929 Liu Z, Li H, Pang H, Me J, Mao X (2015) Enhancement effect of solutes of low molecular mass on
   930 the insect antifreeze protein ApAFP752 from *Anatolica polita*. J Therm Anal Calorim
   931 120:307-315
- 932 Lu K, Li Y (1998) Homogeneous nucleation catastrophe as a kinetic stability limit for superheated
   933 crystal. Phys Rev Lett 80:4474–4477
- 934 Marshall CB, Tomczak MM, Gauthier SY, Kuiper MJ, Lankin C, Walker VK, Davies PL (2004a)
   935 Partitioning of fish and insect antifreeze proteins into ice suggests they bind with comparable
   936 affinity. Biochemistry 43:148–154
- 937 Marshall CB, Daley ME, Sykes BD, Davies PL (2004b) Enhancing the activity of a β-helical 938 antifreeze protein by the engineered addition of coils. Biochemistry 43:11637–11646
- Matubayasi N, Nishiyama A (2006) Thermodynamic quantities of surface formation of aqueous
   electrolyte solutions VI. Comparison with typical nonelectrolytes, sucrose and glucose. J
   Colloid Interface Sci 298:910–913
- 942 Melander W, Horváth C (1977) Salt effects on hydrophobic interactions in precipitation and
   943 chromatography of proteins: an interpretation of the lyotropic series. Arch Biochem Biophys
   944 183:200–215
- Miura K, Ohgiya S, Hoshino T, Nemoto N, Suetake T, Miura A, Spyracopoulos L, Kondo H, Tsuda
   S (2001) NMR analysis of Type III antifreeze protein intramolecular dimer. Structural basis for
   enhanced activity. J Biol Chem 276:1304–1310
- 948 Mok Y-F, Lin F-H, Graham LA, Celik Y, Braslavsky I, Davies PL (2010) Structural basis for the
   949 superior activity of the large isoform of snow flea antifreeze protein. Biochemistry
   950 49:2593–2603
- 951 Neven LG, Duman JG, Beals JM, Castellino FJ (1986) Overwintering adaptations of the stag beetle,
   952 Ceruchus piceus: removal of ice nucleators in winter to promote supercooling. J Comp Physiol
   953 B 156:707–716
- Neven L, Duman JG, Low MG, Sehl LC, Castellino FJ (1989) Purification and characterization of
   an insect hemolymph lipoprotein ice nucleator: evidence for the importance of
   phosphatidylinositol and apolipoprotein in the ice nucleator activity. J Comp Physiol B
   159:71–82
- 958 Nishimiya Y, Ohgiya S, Tsuda S (2003) Artificial multimers of the type III antifreeze protein.
   959 Effects on thermal hysteresis and ice crystal morphology. J Biol Chem 278:32307–32312

## 6 Thermal Hysteresis

Nishimiya Y, Sato R, Takamichi M, Miura A, Tsuda S (2005) Co-operative effect of the isoforms of	960
type III antifreeze protein expressed in Notched-fin eelpout, Zoarces elongatus Kner. FEBS J	961
272:482–492	962
Olsen TM, Duman JG (1997a) Maintenance of the supercooled state in overwintering pyrochroid	963
beetle larvae, <i>Dendroides canadensis</i> : role of hemolymph ice nucleators and antifreeze proteins.	964
J Comp Physiol B 167:105–113	965
Olsen TM, Duman JG (1997b) Maintenance of the supercooled state in the gut fluid of	966
overwintering pyrochroid beetle larvae, <i>Dendroides canadensis</i> : role of ice nucleators and	967
antifreeze proteins. J Comp Physiol B 167:114–122	968
Olsen TM, Sass SJ, Li N, Duman JG (1998) Factors contributing to seasonal increases in inocu-	969
lative freezing resistance in overwintering fire-colored beetle larvae <i>Dendroides canandensis</i>	970
(Pyrochroidae). J Exp Biol 201:1585–1594	971
Pertaya N, Marshall CB, DiPrinzio CL, Wilen L, Thomson ES, Wettlaufer JS, Davies PL,	972
Braslavsky I (2007) Fluorescence microscopy evidence for quasi-permanent attachment of	973
antifreeze proteins to ice surfaces. Biophys J 92:3663–3673	974
Pertaya N, Marshall CB, Celik Y, Davies PL, Braslavsky I (2008) Direct visualization of spruce	975
budworm antifreeze protein interacting with ice crystals: basal plane affinity confers hyperac-	976
tivity. Biophys J 95:333–341	977
Poynting JH (1881) Change of state: solid–liquid. Philos Mag 5th series 12:32–48	978
Ramsay JA (1964) The rectal complex of the mealworm Tenebrio molitor L. (Coleoptera,	979
Tenebrionidae). Philos Trans R Soc B 348:279–314	980
Raymond JA, DeVries AL (1977) Adsorption inhibition as a mechanism of freezing resistance in	981
polar fishes. Proc Natl Acad Sci 74:2589–2593	982
Reynolds JA, Gilbert DB, Tanford C (1974) Empirical correlation between hydrophobic free energy	983
and aqueous cavity surface area. Proc Natl Acad Sci 71:2925–2927	984
Schrag JD, O'Grady SM, DeVries AL (1982) Relationship of amino acid composition and	985
molecular weight of antifreeze glycopeptides to non-colligative freezing point depression.	986
Biochim Biophys Acta 717:322–326	987
Sönnichsen FD, DeLuca CI, Davies PL, Sykes BD (1996) Refined solution structure of type III	988
antifreeze protein: hydrophobic groups may be involved in the energetics of the protein-ice	989
interaction. Structure 4:1325–1337	990
Sørensen TF, Ramløv H (2001) Variations in antifreeze activity and serum inorganic ions in the	991
eelpout <i>Zoarces viviparus</i> : antifreeze activity in the embryonic state. Comp Biochem Physiol A	992
30:123–132	993
Stevens CA, Drori R, Zalis S, Braslavsky I, Davies PL (2015) Dendrimer-linked antifreeze proteins	994
have superior activity and thermal recovery. Bioconjug Chem 26:1908–1915	995
Thomson W (1871) On the equilibrium of vapour at a curved surface of liquid. Philos Mag	996
42:448–452	997
Tolls J, van Dijk J, Verbruggen EJM, Hermens JLM, Loeprecht B, Schüürmann G (2002) Aqueous	998
solubility-molecular size relationships: a mechanistic case study using C10- to C19-alkanes. J	999
Phys Chem A 106:2760–2765	1000
Turnbull D (1950) Kinetics of heterogenous nucleation. J Chem Phys 18:198–203	1001
Uhlig HH (1937) The solubilities of gases and surface tension. J Phys Chem 41:1215–1225	1002
Wang L, Duman JG (2005) Antifreeze proteins of the beetle <i>Dendroides canadensis</i> enhance one	1003
another's activities. Biochemistry 44:10305–10312	1004
Wang L, Duman JG (2006) A thaumatin-like protein from larvae of the beetle <i>Dendroides</i>	100
canadensis enhances the activity of antifreeze proteins. Biochemistry 45:1278–1284	1006
Wang S, Amornwittawat N, Banatlao J, Chung M, Kao Y, Wen X (2009a) Hofmeister effects of	1007
common monovalent salts on the beetle antifreeze protein activity. J Phys Chem B	1008
113:13891–13894	1009
Wang S, Amornwittawat N, Juwita V, Kao Y, Duman JG, Pascal TA, Goddard WA, Wen X	1010
(2009b) Arginine, a key residue for the enhancing ability of an antifreeze protein of the beetle	101
Dendroides canadensis. Biochemistry 48:9696–9703	1012



- 1013 Washburn EW (1929) International critical tables of numerical data, physics, chemistry and technology, vol 4. McGraw-Hill, New York
- 1015 Wen D, Laursen RA (1992) A model for binding of an antifreeze polypeptide to ice. Biophys J 1016 63:1659–1662
- 1017 Wen X, Wang S, Amornwittawat N, Houghton EA, Sacco MA (2011) Interaction of reduced 1018 nicotinamide adenine dinucleotide with an antifreeze protein from *Dendroides canadensis*:
- 1019 mechanistic implication of antifreeze activity enhancement. J Mol Recognit 24:1025-1032
- 1020 Westh HP, Ramløv H, Wilson PW, DeVries AL (1997) Vapor pressure of aqueous antifreeze 1021 glycopeptide solutions. Cryo-Letters 18:277–282
- 1022 Wilson PW (1993) Explaining thermal hysteresis by the Kelvin effect. Cryo-Letters 14:31-36
- 1023 Wilson PW, Beaglehole D, DeVries AL (1993) Antifreeze glycopeptide adsorption on single crystal 1024 ice surfaces using ellipsometry. Biophys J 64:1878–1884
- 1025 Wöhrmann APA (1996) Antifreeze glycopeptides and peptides in Antarctic fish species from the 1026 Weddell Sea and the Lazarev Sea. Mar Ecol Prog Ser 130:47–59
- 1027 Wu DW, Duman JG (1991) Activation of antifreeze proteins from larvae of the beetle *Dendroides* 1028 canadensis. J Comp Physiol B 161:279–283
- 1029 Wu DW, Duman JG, Xu L (1991) Enhancement of antifreeze protein activity by antibodies. 1030 Biochim Biophys Acta 1076:416–420
- 1031 Xu H, Griffith M, Patten CL, Glick BR (1998) Isolation and characterization of an antifreeze protein
   1032 with ice nucleation activity from the plant growth promoting rhizobacterium *Pseudomonas* 1033 putida GR12-2. Can J Microbiol 44:64–73
- 1034 Yang DSC, Sax M, Chakrabartty A, Hew CL (1988) Crystal structure of an antifreeze polypeptide 1035 and its mechanistic implications. Nature 333:232–237
- 1036 Yeung KL, Wolf EE, Duman JG (1991) A scanning tunneling microscopy study of an insect1037 lipoprotein ice nucleator. J Vac Sci Technol B 9:1197–1201
- 1038 Zachariassen KE, Hammel HT (1976) Nucleating agents in the haemolymph of insects tolerant to 1039 freezing. Nature 262:285–287
- 1040 Zachariassen KE, Husby JA (1982) Antifreeze effect of thermal hysteresis agents protects highly 1041 supercooled insects. Nature 298:865–867
- 1042 Zachariassen KE, DeVries AL, Hunt B, Kristiansen E (2002) Effect of ice fraction and dilution 1043 factor on the antifreeze activity in the hemolymph of the cerambycid beetle *Rhagium inquisitor*.
- 1044 Cryobiology 44:132–141
- 1045 Zanetti-Polzi L, Biswas AD, Del Galdo S, Barone V, Daidone I (2019) Hydration shell of antifreeze 1046 proteins: unveiling the role of non-ice-binding surfaces. J Phys Chem B 123:6474–6480
- 1047 Zepeda S, Yokoyama E, Uda Y, Katagiri C, Furukawa Y (2008) *In situ* observation of antifreeze
- 1048 glycoprotein kinetics at the ice interface reveals a two-step reversible adsorption mechanism.
- 1049 Cryst Growth Des 8:3666–3672



# **Author Query**

Chapter No.: 6 385597\_1\_En

Query Refs.	Details Required	Author's response
AU1	Please check and confirm the edit made in the sentence beginning "Such perpetual motion of water molecules would"	