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Site-specific conjugation of antifreeze proteins onto polymer-stabilized nanoparticles†

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Antifreeze proteins (AFPs) have many potential applications, ranging from cryobiology to aerospace, if they can be incorporated into materials. Here, a range of engineered AFP mutants were prepared and site-specifically conjugated onto RAFT polymer-stabilized gold nanoparticles to generate new hybrid multivalent ice growth inhibitors. Only the SNAP-tagged AFPs lead to potent 'antifreeze' active nanomaterials with His-Tag capture resulting in no activity, showing the mode of conjugation is essential. This versatile strategy will enable the development of multivalent AFPs for translational and fundamental studies.

Antifreeze (glyco)proteins (AF(G)Ps), are found in a diverse range of organisms¹ including polar fish,² insects³ and plants,⁴ as well as ice-binding polysaccharides in insects.⁵ There is significant interest in the use of these proteins for low temperature applications from aeronautical engineering to wind turbines and for the cryopreservation of donor cells and tissues.^{6,7} For this to be a reality, the AFPs need scalable syntheses (or mimics^{8,9}) and methods to incorporate them into more complex devices or coatings.^{10,11} AFPs have three main effects, of ice recrystallisation (growth) inhibition (IRI), non-colligative depression of the freezing point leading to a thermal hysteresis (TH) gap and dynamic ice shaping (DIS). To enable interaction with the dynamic surface of ice, most AFPs (but not the more flexible antifreeze glycoproteins¹²) have a defined ice-binding face¹ which can anchor them directly or *via* ordered clathrate water.^{13–16} A distinct class of AFPs are the hyperactive AFPs.^{17,18} The increased TH activity of hyperactive AFPs is linked to their binding of both prism and basal planes of ice, compared to just the prism plane for standard AFPs.^{19,20} Increasing the concentration of particular type I

AFPs has been observed to lead to oligomerization to a tetramer and the onset of hyperactivity, linked to its supramolecular assembly.²¹ Davies and coworkers assembled 6–11-mers of AFP type III on PAMAM dendrimers to mimic this oligomerization. On a per-protein basis there was only a small increase in IRI, but the ability to span multiple ice faces increased the TH activity.²² Synthetic mimics of AFPs also show strong molecular weight dependence on activity, with longer polymers (such as poly(vinyl alcohol) having significantly higher IRI activity^{23–25} as do supramolecular safranin-O based mimics.²⁶

Despite the evidence for increasing the valency of AFPs to modulate activity there remain few reports of multivalent display, in part due to the challenges of site-specific protein conjugation.^{10,27,28} Traditional approaches to combine polymers with proteins involve targeting unpaired cysteine residues.^{29,30} Unnatural amino acids for bioconjugation can be incorporated in a site-specific fashion by the AMBER stop codon.³¹ Johnsson and co-workers have developed recombinantly expressible 'SNAP-tags' based on O⁶alkylguanine-DNA alkyltransferase.³² By attaching the tag as a fusion protein a covalent bond can be formed to any surface bearing benzylguanine without any unnatural amino acids. The commonly used hexa-histidine purification tag can also facilitate ionic conjugation.³³

Here AFPs are conjugated, by site-specific methods, onto nanoparticles to generate hybrid ice growth inhibiting materials to mimic the multivalent presentation of hyperactive antifreeze proteins and aid the application of these exciting proteins (Fig. 1).

RAFT (reversible addition fragmentation chain transfer) polymerization was employed to synthesize telechelic poly(hydroxyethyl acrylamide), pHEA, bearing a pentafluorophenyl (PFP) ester at the α -terminus and a trithiocarbonate at the ω -terminus.³⁴ These polymers were characterized by SEC, ¹H, ¹⁹F NMR and IR (Table 1). The PFP group was substituted by addition of amino-benzylguanine (BG) (for SNAP conjugation³²) or tris-NTA amine (for His-Tag capture). Successful conjugation was confirmed by ¹⁹F NMR (Fig. 2B) as well as by

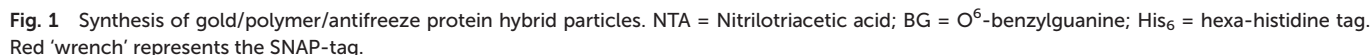
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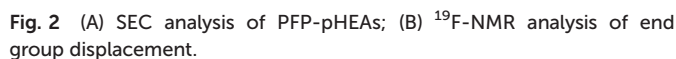
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Polymer	[M]/[CTA] (–)	Conv. ^a (%)	DP _n (–)	M _{n(THEO)} ^a (g mol ^{–1})	M _{n(SEC)} ^b (g mol ^{–1})	D ^b (–)
PFP-pHEA ₉₅	103	93	95	11 000	15 000	1.3
PFP-pHEA ₄₅	47	94	45	5700	8300	1.1
PFP-pHEA ₁₇	26	64	17	2500	3000	1.1

^a Determined by ¹H NMR. ^b Determined by SEC against PMMA standards.

Recombinant AFP type III (from ocean pout) were produced by expression in *Escherichia coli*, and purified by nickel affinity chromatography and HPLC. Three mutants were produced bearing N-terminal modifications of hexa-histidine (His-AFP), SNAP (SNAP-AFP) and cysteine (Cys-AFP). Direct conjugation of the SNAP-AFP was achieved by incubation with the BG-pHEA₄₅@AuNP₄ in buffer. NTA-pHEA₁₇@AuNP₄ was first activated by addition of NiCl₂ followed by His-AFP. Direct Cys-AFP immobilization onto AuNPs was unsuccessful (see ESI†). Particles were further purified by centrifugal dialyses and removal of non-conjugated residual protein confirmed by nanodrop absorption analysis on residual washings. Following



Fig. 3 Nanoparticle characterization. (A–C) Transmission electron micrographs of citrate@Au₄, SNAP-AFP-BG-pHEA₄₅@Au₄ and His-AFP-Ni-NTA-pHEA₁₇@Au₄ respectively; (D) N 1s peak from XPS upon conjugation of SNAP-AFP; (E) diameter from dynamic light scattering.

Table 2 Nanoparticle characterization

Particle	D_h^a (nm)	D_{TEM}^b (nm)	ζ potential (mV)
Citrate@Au ₄	4.3	3.9 ± 0.7	−19.4 ± 2.7
NTA-pHEA ₁₇ @Au ₄	8.5	3.7 ± 0.8	−15.9 ± 4.8
Ni-NTA-pHEA ₁₇ @Au ₄	7.7	3.9 ± 0.8	−6.16 ± 1.1
His-AFP-Ni-NTA-pHEA ₁₇ @Au ₄	8.9	4.3 ± 0.9	−7.03 ± 1.3
BG-pHEA ₄₅ @Au ₄	16.5	3.9 ± 1.0	−10.2 ± 1.9
SNAP-AFP-BG-pHEA ₄₅ @Au ₄	8.92	4.0 ± 1.1	−4.35 ± 0.9

^a Hydrodynamic diameter from DLS. ^b Gold core diameter, average of >100 particles by TEM.

conjugation the zeta-potential became less negative and XPS confirmed a further increase in the nitrogen concentration upon protein conjugation (Table 2).

With these site-specifically attached AFP/nanoparticle conjugates to hand, their IRI activity could be assessed by the 'splat' assay. In brief, a polynucleated wafer of ice crystals is seeded from a solution containing the compounds of interest and the ice crystals annealed at −8 °C for 30 minutes. After this time the average ice crystal size is evaluated and compared to a negative control of buffer alone: smaller ice crystal sizes signify more IRI activity. The two AFP mutants were very potent inhibitors, preventing all growth below 0.01 mg mL^{−1}. There was no significant difference between the two mutants, showing the fusions were tolerated in terms of retention of function. His-AFP-Ni-NTA-pHEA₁₇@Au₄ was found to be surprisingly inactive as an ice growth inhibitor with no activity up to 1 mg mL^{−1} (AFP concentration). This could be due to the dynamic nature of the His-NTA interaction or incorrect orientation of the AFP on the surface. This also shows that designing a multivalent AFP is non-trivial, with the correct chemistry



Fig. 4 (A, C, D) IRI activity of His-AFP and conjugates by MLGS (mean length grain size); (B, E, F) IRI activity of SNAP-AFP and conjugates; (G) thermal hysteresis showing freezing point depression in SNAP-AFP and conjugate samples; ice crystal morphology of (H) SNAP-AFP; (I) SNAP-AFP-BG-pHEA₄₅@Au₄. Scale bar = 100 μm.

being essential. Previous studies suggest excess charge does not affect AFP activity and was ruled out as the cause.³⁶ In contrast, the covalent hybrid nanoparticle SNAP-AFP-BG-pHEA₄₅@Au₄ was found to be a very potent IRI, Fig. 4B. In terms of total AFP concentration (to enable multivalent effects to be seen) the nanoparticles essentially identical activity compared to SNAP-AFP alone. If considered on a molar basis (as has been done by Davies *et al.* for dendrimer AFPs²²) then the particles are more active. AFPs are known to become hyperactive when they oligomerize, but such enhancement was not seen here due to the different 3-D placement of the AFPs.²¹ Using a nanoliter osmometer the thermal hysteresis (TH) gap (non-colligative freezing point depression) was determined. TH is closely linked to the ability to bind specific ice faces (in this case the prism plane) hence providing additional information about the functionality of the particles; the TH is unique to antifreeze proteins, and is defined as the difference between the freezing and melting points. It was observed that the AFP and SNAP-AFP nanoparticles have very similar TH gaps, of 1.4 °C at 7.6 mg mL^{−1}. As with the IRI measurements, this confirms that all activity is retained despite the major structural modifications and that the placement of the AFPs *via*-polymeric linkers is tolerated. As the effective AFP concentration in solution when immobilized on NPs is lower, this suggests that the particles can bind multiple planes of ice simultaneously. To probe this dynamic ice shaping experiments were conducted in sucrose solution (as described previously²⁸). Compared to SNAP-AFP alone (Fig. 4H), the SNAP-AuNP conju-



gates resulted in more faceting of the ice crystals, consistent with them being able to bind more ice faces simultaneously.²²

Conclusions

In conclusion, we report the synthesis of hybrid nanoparticle–AFP conjugates, making use of polymer-stabilized gold nanoparticles and protein engineering to enable precision-conjugation. Hetero-telechelic polymers were obtained by RAFT, enabling a nanoparticle immobilization moiety (thiol) and distal protein capture (nickel NTA or O⁶-benzyl-guanine) moiety to be installed. The polymer coating is essential to enable conjugation in buffer, avoiding nanoparticle aggregation. It was found that the SNAP-tag (covalent) approach led to the most active particles compared to oligohistidine capture. The observed ice recrystallisation inhibition and thermal hysteresis activity was comparable to free antifreeze proteins on a valency-corrected basis confirming retention of activity in this format. Other conjugation methods based on ionic interactions did not give stable particles or resulted in loss of activity. These results demonstrate that antifreeze protein can be incorporated into more complex assemblies to facilitate their application in colloid science or for fundamental studies exploiting nanoparticle cores for bioimaging and tracking.

Data access statement

The research data supporting this publication can be found at <http://wrap.warwick.ac.uk>.

Conflicts of interest

There are no conflicts to declare.

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