Cation-induced shape programming and morphing in proteinbased hydrogels

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ABSTRACT: Smart materials that are capable of memorizing a temporary shape, and morph in response to a stimulus, have the potential to revolutionize medicine and robotics. Here we introduce a novel method to program protein-hydrogels and to induce shape-changes in aqueous solutions at room temperate. We demonstrate our approach using hydrogels made from serum albumin, the most abundant protein in the blood plasma, which are synthesized in a cylindrical or flower shape. These gels are then programmed into a spring or a ring shape, respectively. The programming is done through a dramatic change in stiffness (of up to 17-fold), induced by adsorption of Zn²⁺ or Cu²⁺ cations. We show that these programmed biomaterials can then morph back into their original shape, as the cations diffuse outside the hydrogel material. The approach demonstrated here represents a novel strategy to program protein-based hydrogels, which naturally harvest from the biodiversity and large functional space of these molecules, and morph between two configurations through a chemo-mechanical triggering stimulus.

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INTRODUCTION

Dynamic biomaterials that undergo conformational changes can enable artificial tissue structures, that could experience morphological transformations, and soft robotics, that could react and change based on their environment. Currently, most common shapemorphing materials are based on polymers and require to switch between a stiff and a soft phase (1). These materials typically rely on two or more network skeletons, sharing the same 3-dimensional (3D) space (2), or have a chemical response to small ions (3, 4). Programming, defined as the capability of fixing a temporary shape in a material, requires a dramatic increase in stiffness, in a reversible manner (5). The initial shape recovery entails a switch from a stiff to a soft phase, and is typically realized by compromising the integrity of the secondary network, by changing the temperature (1, 6), pH (7), solvent (8, 9), or through photo-switching (10, 11).

Protein-based hydrogels utilize a protein as their primary network, in a water-rich environment (12). These hydrogels retain many of the characteristics displayed by the polymer based materials, but can harvest from a much more diverse bio-functional library. Proteins accomplish many life-supporting functions, from structural role to enzymatic reactivity and their function is in most cases directly related to their folded 3D structure. While there is a significant diversity for the starting material, when compared to polymers, proteins are stable and functional in a much narrower range of temperatures, pH, or salt conditions, and require a water-based environment. Since the mechanical response of a material depends directly on the concentration of its constituent network nodes (13-15), the range of obtainable stiffness for protein-based hydrogels is extremely limited: the protein needs to be above the critical gelation concentration to be turned into a biomaterial, and below its specific solubility limit (15-17). This narrow range allows for a change in stiffness of only ~10-30 %, depending on the starting protein. Furthermore, well-defined cross-linked network connections are critical to ensure high shape recovery ratio (18). Unlike most linear polymer molecules, which can entangle like spaghetti in a bowl, globular proteins have well-defined 3D structures, acting as hard spheres. This

structural integrity provides an excellent control over the cross-linking points and density, while preserving the tertiary structure of the network nodes.

Recently, we have introduced a new method to obtain shape memory in proteinbased hydrogels by stiffening them through adsorbed polyelectrolytes (19). Our approach relies on producing protein hydrogels from bovine serum albumin (BSA), which is the homologous of human serum albumin, the most abundant protein in the blood plasma. BSA solutions can be turned into completely covalently cross-linked hydrogel biomaterials when starting from solutions with protein concentrations above 1 mM (17, 20). Below 1 mM, BSA hydrogels show irreversible plastic deformation under force, indicative of incomplete cross-linking (17). Due to the overall charge of BSA, these protein hydrogels have been programmed by stiffening, induced by the secondary network made from positively-charged polyelectrolytes (19). The shape change was induced in this case via the unfolding response of the protein domains in chemical denaturants, and complete recovery upon removal of the denaturing solution. In the unfolding phase transition responsible for shape memory, the proteins lose their tertiary structure and show a remarkable decrease in stiffness. This transition is highly repeatable, as protein folding has been tightly controlled by billions of years of evolution. However, the polyelectrolytes adsorption is irreversible and, due to their large size, the loading capacity is relatively limited, resulting in a change in stiffness of up to ~6.5 folds. Here we explore the viability of using divalent cations to stiffen protein-based hydrogels toward programming them in various shapes. We then manage to morph back the protein materials from the temporary programmed shape to the initial permanent shape, through simple diffusion. It has been previously shown, for other protein hydrogels (21) and peptide-based hydrogels (22), that divalent cations can reinforce their network. It is also well known that BSA can bind cations at various exposed amino acids (23, 24). For example, divalent cations such as Ni²⁺, Cu²⁺ and Zn²⁺ were shown to bind histidine (25, 26), tryptophan (24) and to bridge cysteine amino acids (27). Adsorption of cations was also associated with an increase in the mechanical stability of proteins at the molecular level (28, 29). Here we explore this mechanical change to induce the increase in stiffness needed to program protein-based biomaterials in various shapes (Figure 1). The small ions have the advantage of allowing much higher loading of positive charges, leading to a 17-fold stiffening, and the capacity

to diffuse outside the programmed material. This novel implementation of programming protein-based hydrogels with positively charged ions is an important step toward obtaining biocompatible materials that can adjust their shape.

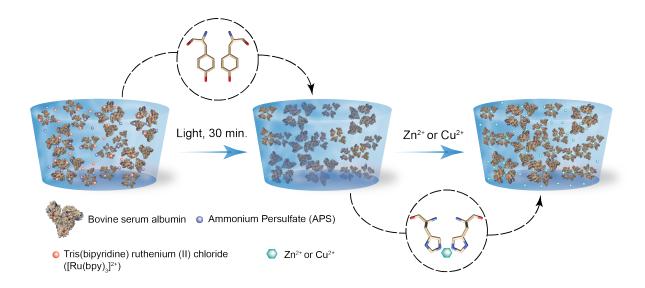


Figure 1. Schematics of the fixation process. (Left): BSA-based protein hydrogels are fabricated using a light activated reaction, in the presence of APS and $[Ru(bpy)_3]^{2^+}$. (Right): Following synthesis, the protein hydrogels are exposed to Zn^{2^+} or Cu^{2^+} , which reversibly increases their stiffness by up to 200 folds. This stiffening effect can be used for shape-programming.

RESULTS AND DISCUSSION

Protein-based hydrogels can be obtained using various cross-linking strategies, such as treatment with glutaraldehyde (20), enzymatic reactions (30), using protein-based lock-and-key ends (31, 32), or photo-activation (12). Here we use photo-activation via $[Ru(bpy)_3]^{2+}$ to obtain protein-based hydrogels made from BSA (Figure 1). This reaction was shown to produce covalent carbon-carbon bonds at the exposed tyrosine amino acid sites between adjacent protein domains (12). The advantage of using light to trigger the start of the cross-linking reaction is that it allows us to load the reaction mixture in the desirable shape, without any change in viscosity before light exposure.

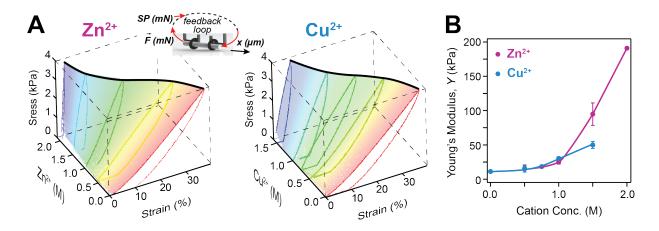


Figure 2. Cation-based stiffening of protein-based hydrogels. A) Chemo-mechanical changes induced by adsorption of various concentrations of Zn^{2+} (left) and Cu^{2+} (right) by protein hydrogels made from 2 mM BSA. The mash highlights the force-loading part, used to assess the change in stiffness, and the thick, black curve follows the final strain at 4 kPa stress. Inset: Schematics of a hydrogel tube pulled under a feed-back controlled force, where the set-point SP was increased and decreased linearly with 40 Pa/s. B) Change in measured Young's modulus as a function of cation concentrations. Both Zn^{2+} and Cu^{2+} induce stiffening when adsorbing to BSA-based hydrogels. Lines are eye-guides. Error bars are S.D. (n = 3).

Our first step was to explore the range of concentrations of two positively charged ions, that can potentially increase the stiffness of protein hydrogels and allow for shape

programming. The change in stiffness of protein-based hydrogels in the presence of positively charged ions was quantified using our force-clamp rheometry apparatus (17, 33). We first produced cylindrical-shaped hydrogels staring from 2 mM BSA, using PTFE tubes with an inner diameter of 0.56 mm. We chose 2 mM as staring concentration, as our polymerization method produces complete cross-linking for BSA, and the hydrogels show reversible behavior and no plastic deformations in the sampled force range (17). Using surgical thread, these gels were then attached in our force-clamp rheometer via two metal hooks connected to the voice-coil motor and force sensor, respectively (Figure 2 inset, see also the Methods section). Following incubation for 30 min in a phosphate buffer with the desired cation concentration, the mechanical response of treated BSA hydrogel was measured in the 0-4 kPa range (Figure 2). The change in stress as a function of strain, as the applied force is increased linearly with time, can be used to assess the stiffness of the material, as the slope in the rising part of the trace directly reports on the dynamic Young's modulus. Typically, hydrogels made from globular proteins such as BSA also show hysteresis in the stress-stain curves (Figure 2A). This hysteresis disappears for BSA-hydrogels when exposed to chemical denaturants, which break down the tertiary structure of the protein domains forming the hydrogel network (12, 17). This hysteresis was related to the imbalance between the forces where unfolding and refolding take place, with the unfolding transition occurring at much higher forces than the refolding (34), and can allow for large energy dissipation before failure (35, 36). BSA hydrogels show an up to 5-fold increase in stiffness when treated with Cu²⁺ (50 kPa in 1.5 M Cu²⁺ from 11 kPa) and a 17-fold increase in stiffness in the presence of Zn²⁺ (191 kPa in 2 M Zn²⁺) (Figure 2B). The stiffening of BSA-hydrogels in 2 M Zn²⁺ is several orders of magnitude greater than that reported for the same gels when treated with polyelectrolytes (19) and should allow for more complex programmed shapes. The stiffening effect seems to depend more on the solution concentration of cations rather than their nature (Figure 2B). The main advantage of Zn²⁺ over Cu²⁺ is its higher solubility in water, which allows us to prepare solutions with higher concentrations and observe greater stiffening.

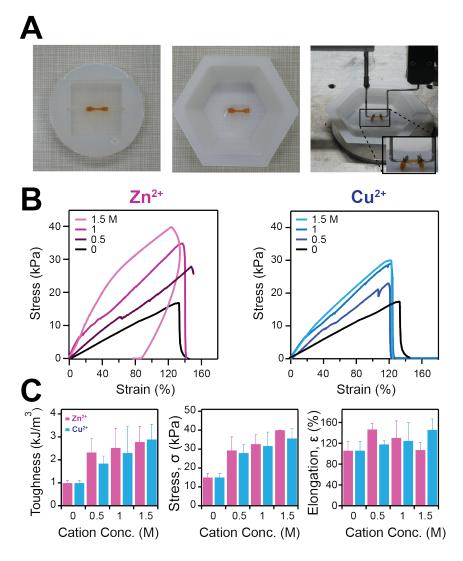


Figure 3. Mechanical characterization of BSA-hydrogels immersed in cations of various concentrations. A) Picture of a gel casted using a bone-shaped silicone mold (left and middle) and attached in the force-clamp rheometer (right). B) Stress vs strain of BSA hydrogels immersed in Zn^{2+} (left) and Cu^{2+} (right) and pulled until breaking. C) Toughness (left), failure stress (middle) and maximum elongation (right) of BSA hydrogels incubated with increasing concentrations of Zn^{2+} (magenta) and Cu^{2+} (blue). Toughness increased from 1.0 ± 0.1 kJ/mol to 2.8 ± 0.7 kJ/mol in the presence of 1.5 M Zn^{2+} and to 2.9 ± 0.7 kJ/mol in the presence of 1.5 M Cu^{2+} . The breaking stress increase from 15 ± 2 . kPa to 33 ± 5 kPa when BSA-hydrogels were treated with 1.5 M Zn^{2+} and to 36 ± 5 kPa when treated with 1.5 M Cu^{2+} . The failing strain show little variation ($106 \pm 18\%$ vs $107 \pm 15\%$ in Zn^{2+} 1.5 M and $146 \pm 21\%$ in Cu^{2+} 1.5 M). Error bars are S.D. (n=3) (photo credit: Luai R. Khoury, UWM; Marina Slawinski, UWM).

Apart for the stiffening effect, incubation of protein hydrogels in solutions with high concentrations of cations improves their mechanical failure properties (22). For these tests, we use a typical bone shape, where the BSA hydrogels were extended until failure or maximum force range of our force sensor. BSA-based hydrogels show an increase in both toughness and failure stress with increasing cation concentration. The measured toughness, which represents the ability of a material to absorb energy and deform without fracturing and is derived from the area under the pulling stress-strain curve (Figure 3B and C left), increased from 1 kJ/mol to 2.8 kJ/mol with cation concentration. The failure stress increased from 15 kPa to 33 kPa. Interestingly, the maximum elongation did not show a significant variation with cation concentrations. This behavior suggests that, while the stiffness is probably given by the increase in the mechanical stability of protein domains (27) and non-covalent bridging (22) at ~ 120 % the primary network of the BSA hydrogels is the one starting to fail, experiencing irreversible breaking of covalent bonds. Hence, the cross-linking geometry is the limiting factor for extensions, and for further improvements in maximum elongation the primary hydrogel network would require adjustments.

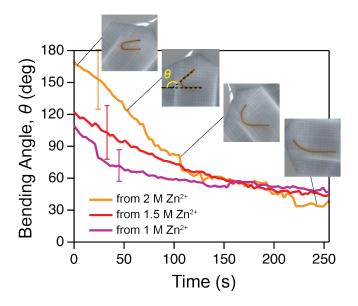


Figure 4. Changes in the hydrogel shape due to cation diffusing outside the hydrogel. Measured programmed angle of a U-shape gel, θ , as a function of time, upon immersion from Zn^{2+} into regular TRIS buffer. Inset: Pictures of the hydrogel recovering from a U-shape at four

different time points. Second inset shows how the angle is measured. The error bars represent S.D. (n = 3). Supplementary Movie 3 accompanies this figure (photo credit: Luai R. Khoury, UWM; Marina Slawinski, UWM).

While stiffening through a secondary effect is necessary for programing in-shape of the hydrogel material, the dynamics of morphing from the programmed to the initial shape will directly depend on the diffusion of the cations outside the hydrogel. To monitor this effect, we first programmed a cylindrical hydrogel in a U-shape (Figure 4). BSA hydrogels were mounted in a U-shape configuration, corresponding to a bending angle of 180 deg (as defined in Figure 4 Inset), and incubated for 30 min in three different concentrations of Zn²⁺ (8, 37). The fixity, R_f, which reports on the degree of the programmed hydrogel to maintain its shape when taken out of the mold used during the programming phase, varied from 96 \pm 3 % in 2 M Zn²⁺ to 58 \pm 15 % in 1 M Zn²⁺ (Supplementary Figure 2). The fixity of protein hydrogels depends on the amount of stiffening that can be induced when dosing with the cation concentration. Hence the ~17-fold increase in stiffness when incubating BSA-hydrogels with 2 M Zn²⁺ produces a programmed bending degree closer to the Ushape mold than the equivalent ~2-fold stiffening in 1 M Zn2+ (Figure 4 and Supplementary Figure 2). These values are comparable with those measured for BSA hydrogels programmed with PEI (19) and with other polymeric materials using double network or polymer-ion interactions (37). As shown in Fig. 4, when a programmed hydrogel is immersed in regular TRIS buffer, the bending angle decreases to a final value of ~45 deg in ~3 min. The mechanism behind obtaining the temporary shape involves a combination of ionic cross-linking (4) and the stabilization due to divalent cations of BSA domains (38). The shape morphing results from the diffusion of these divalent ions in the surrounding medium (Figure 4).

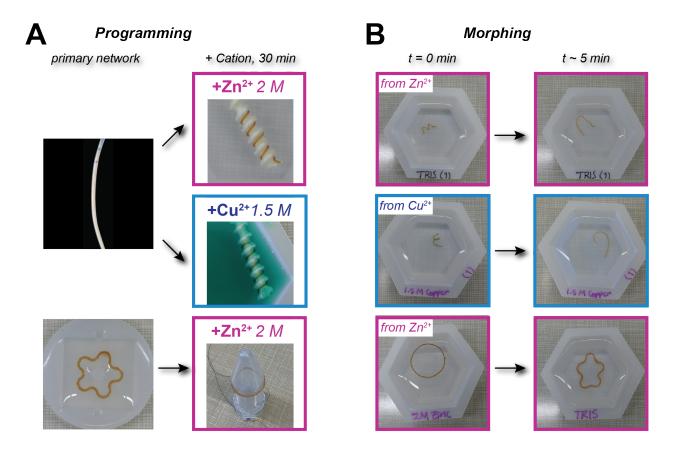


Figure 5. Programming of protein hydrogels with cations and morphing via chemomechanical changes. A) BSA hydrogels were casted in cylindrical shape in PTFE tubes (top left) and programmed in a spring shape using a negative cast, by immersion in 2 M Zn²⁺ solution (top right) or 1.5 M Cu²⁺ solution (middle right) for 30 min; BSA hydrogels were produced in a flower shape using a silicone mold (bottom left) and programmed into a ring, by immersion in 2 M Zn²⁺ solution for 30 min (bottom right). B) Morphing from the programmed shape into the casted shape of BSA hydrogel upon immersion in regular TRIS buffer at time 0 (left) and 5 min (right) for the hydrogels from panel A. Supplementary Movie 1 and 2 accompany this figure (photo credit: Luai R. Khoury, UWM; Marina Slawinski, UWM).

Using the large change in stiffness of BSA hydrogels, induced by immersion in Zn^{2+} and Cu^{2+} solutions, we programmed cylindrically casted biomaterials into a spring shape, and flower casted materials into a ring shape (Figure 5 top and Supplementary Figure 3). As shown with polyelectrolytes, a ~6.5-fold increase in stiffness already suffices to program BSA hydrogels into a spring shape, and both Zn^{2+} and Cu^{2+} induce a strong enough stiffening (up to ~17 fold). The main advantage of the small ions over polyelectrolytes is

their diffusion, which happens relatively fast (< 5 min). Furthermore, in the present case, the shape morphing is driven by simple diffusion, and does not require compromising the primary protein network with the help of chemical denaturant. Another advantage of the increase in Young's modulus of BSA hydrogels in the presence of cations is that more complex shapes can be obtained. For example, we demonstrate the morphing from a ring to a flower shape (Figure 5 bottom and Supplementary Figure 3). Compared to the spring shape, in this case there are no free ends that can release any torsional tension. To obtain this complex shape morphing, we first casted the hydrogel in a flower-like shape using a silicone mold. Following the light-activated cross-linking reaction, we then programmed the hydrogel into a ring shape by mounting it onto a plastic tube, which was then immersed in the cation solution for 30 min. When removed from the plastic tube into a Zn^{2+} solution with the same concentration, the hydrogel maintains the ring shape (Figure 5B bottom-left). However, when immersed into a regular TRIS buffer, the ring shape quickly morphs into the original flower shape (Figure 5B bottom-left and Supplementary Movie 1).

CONCLUSIONS

Polymer-based hydrogels have found various applications for shape memory and shape morphing. While these approaches can be made biocompatible, polymers cannot reach the same diversity and control over the sequence and structure as proteins. The approach demonstrated here enables shape morphing in protein-based hydrogels, which could harvest the best of both worlds. This approach relies on the stiffening induced by Zn²⁺ and Cu²⁺ to program a permanent shape into a new temporary configuration, and the diffusion of these ions outside the material enables the recovery of the original shape. While we demonstrate this approach with both Zn²⁺ and Cu²⁺, we envision that Zn²⁺ will enable more biologically relevant applications, as it is significantly less toxic than Cu²⁺. The main advantages of cation-programmed protein hydrogels are that the attainable stiffness is much higher than in regular buffer (~17 fold), enabling programming in complex shapes, and that the fast diffusion of the small ions leads to fast irreversible

morphing (<5 min). Furthermore, the shape change is taking place in an aqueous environment at room temperature, which is compatible with conditions present in the human body. Permanent shape morphing based on protein hydrogels and cations could find applications for various implants (39) and injectable hydrogels (40). Also, the shape morphing method demonstrated here, from a temporary to a permanent profile, does not require denaturation of the tertiary structure of protein domains inside the hydrogels. Hence this approach allows the preservation of the functionality of proteins forming the skeleton of the hydrogel. In conclusion, the approach presented here provides a remarkable combination of biological diversity and programming capability.

MATERIALS AND METHODS

Materials. Bovine Serum Albumin (BSA, M.W. ~ 66.5 kDa) was purchased from Rocky Mountain Biologicals (RMBIO); Sodium Phosphate, Monobasic, Anhydrous (NaHPO₄) (M.W. 119.98 g/mol) was obtained from Research Products International. Sodium Chloride (NaCl) (M.W. 58.44 g/mol) purchased from Fisher Scientific. Ammonium persulfate (APS) (NH₄)₂S₂O₈ (M.W. 228.20 g/mol, 1 M) solution, Tris(bipyridine) ruthenium(II) chloride [Ru(bpy)₃]²⁺ (M.W. 748.62 g/mol, 6.67 mM) solution, Trizma base NH₂C(CH₂OH)₃ (M.W. 121.14 g/mol), Hydrochloric acid (HCl) (37%), Zinc sulfate monohydrate (ZnSO₄·H₂O) (purity ≥99.9%) (M.W. 179.47 g/mol), Copper(ii) chloride dihydrate (CuCl₂·2H₂O purity ≥99.99%) (M.W. 170.48 g/mol), and Sigmacote were purchased from Sigma-Aldrich. TRIS (Tris-NaCl 20 mM, NaCl 150 mM, pH~7.4) and PBS (NaHPO₄ 10 mM, NaCl 150 mM, pH~3) were used as buffers. DDH₂O was used in all solution preparations.

Synthesis of BSA-based hydrogels. Three different shapes of BSA-based hydrogels were prepared in this study. First, a BSA-based hydrogel mixture was prepared by mixing a 2 mM BSA solution with 1 M APS and 1 M [Ru(bpy)₃]²⁺ in a volume ratio of 15:1:1. To prepare the hydrogels with the cylindrical shape, we followed the same procedure as described in our previous studies (*17*, *33*). Briefly, polytetrafluoroethylene (PTFE) tubes (I.D. 0.022 in, Cole-Parmer) were passivated with Sigmacote for 5-10 min and dried thoroughly. The solution mix was then injected and the tube was placed under a light source for 30 min. The Bone- and Flower-like shape hydrogels were prepared starting

from a custom-made silicone rubber mold made of Dragon Skin 30 (purchased from Smooth-On) (Supplementary Figure 1). The bone-shape hydrogel samples possessed an overall length of 9 mm. The gauge length and width are 5 mm and 1 mm, respectively and a 1 mm thickness (Supplementary Figure 1A). The flower-like shape sample has a width of 0.8 mm and 1 mm thickness (Supplementary Figure 1B). Thereafter, the molds were passivated with Sigmacote for 10 min. The hydrogel mixture was casted into the slot and covered with a glass coverslip to reduce evaporation. The loaded molds were then placed under a 100 W mercury lamp for 30 min, after which the hydrogel samples were removed from the molds and immersed in TRIS solution.

Mechanical studies. The mechanical investigation of hydrogel samples was carried out using a force-clamp machine, as described in previous studies (*17*, *33*). A force-ramp protocol with a rate of 0.04 kPa/s was applied on the hydrogel sample while immersed in TRIS solution at room temperature. Afterwards the hydrogel sample was immersed for 30 minutes at room temperature into one of two cations solutions: Zn²⁺, Cu²⁺ which were dissolved in phosphate buffer at different concentrations. We chose 30 min incubation time, as for the gels used here (with cylindrical shape and diameter of 0.56 mm, or with square cross-section of 0.5x0.5 mm) the material fully equilibrates with the solvent environment. Then, same force protocol was applied on the treated hydrogel sample. The Young's modulus, toughness, and breaking stress and strain were calculated from stress-strain curves.

Shape programming and morphing. The shape programming and morphing of the BSA-based hydrogel samples were performed by starting from the tube, or the flower shapes. First, a 2 mM BSA-based hydrogel was synthesized inside the plastic tube or the silicone mold. Then, the hydrogels obtained in the tubes were fixed in a 3D spiral-like shape or U-shape, while the flower-shaped hydrogels were fixed in a circular-shape around a 5 mL plastic tube. Following treatment with Zn²⁺ or Cu²⁺ for 30 min at room temperature, at various concentrations, the gels were removed from the fixing mold and placed into the same solution buffer. The shape morphing was then induced by transferring the fixed gels from the Zn²⁺ or Cu²⁺ solution into a regular TRIS buffer. Recordings were done with a Panasonic digital camera, in the time-lapse video mode, with a frame saved every three seconds.

Characterizations. All data acquisition and analysis of the mechanical behavior of hydrogels was done in Igor Pro (Wavemetrics). All image-processing for measuring bending angle of the U-shape gels was accomplished in ImageJ. The stress values were calculated by dividing the measured force (mN) to the cross-section area, while the strain values were obtained from the measured extension, in respect to the tethered gel length.

To quantify the ability of our protein hydrogels to memorize their temporary shape and the dynamics of recovery from the temporary programmed shape into the initial permanent shape, we used the U-shape recovery method (37, 41). In this approach, the hydrogel is programmed into a U-shape and a bending angle θ is defined as the difference between the original orientation (in our case 180 deg) and the programmed orientation (the angle between the arms of the programmed shape). The shape fixity, R_f , reports on how well the temporary programmed shape can memorize the geometry of the casting mold in the presence of cations, and was calculated as:

$$R_f = \frac{\theta_o}{\theta_p} \times 100$$

where θ_0 was the measured bending angle at time zero, right after the gel was removed from the programming process. θ_p is the expected programmed angle (which for the U-shape is 180 deg).

The dynamics of switching from the temporary programmed shape into the initial permanent shape was also quantified by monitoring θ (t), the bending angle of the U-shape hydrogel in TRIS buffer, at various times t.

SUPPLEMENTARY MATERIALS

Supplementary Figure 1. Schematics showing the two different shapes used in this study to synthetize hydrogels

Supplementary Figure 2. Changes in the measured fixity ratio of U-shape hydrogel as a function of Zn²⁺ concentration.

Supplementary Figure 3. Additional Examples of cation-induced shape morphing for protein hydrogels.

Supplementary Movie 1. Movie showing the morphing of a BSA-hydrogel casted as a ring and programmed into a flower in 2 M Zn²⁺ for 30 min and immersed in PBS buffer.

Supplementary Movie 2. Movies showing the morphing of BSA-hydrogels casted as a cylinder and programmed as a spring shape in 2 M Zn²⁺.

Supplementary Movie 3. Movie showing the morphing of BSA-hydrogels casted as a cylinder and programmed in a U-shape in 2 M Zn²⁺.

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Data and materials availability:

All data needed to evaluate the conclusions in the paper are present in the paper and/or the Supplementary Materials. Additional data related to this paper may be requested from the authors.