## **Supporting Information**

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A T7 His6 Bio Elastin-Like Region 3 B								
	Elastin-Like Protein	Amino Acid Sequence						
	Component	Sequence						
	T7-tag	MASMTGGQQMG						
	Polyhistidine-tag	ННННН						
	<b>Bio-Active Region</b>	TVYAVTGRGDSPASS						
	Elastin-Like Region	(VPGIV)2(VPGKG)(VPGIG)2						

- 1 Figure S1. Elastin-like protein amino acid sequence information: (A) A schematic representation of the
- 2 elastin-like protein (ELP) used in the present manuscript. Our ELP contains three key-components: (1) a
- 3 tag region with a T7-tag and Polyhistidine-tag, (2) a bio-active domain and (3) an elastin-like region. (B)
- 4 The amino acid sequence for these components are included for reference.

### NMR Quantification:

5 Hydrazine functionalization: dissolve ~10 mg of the Boc-protected ELP-Hydrazine intermediate in 6 ~750  $\mu$ L deuterated DMSO and analyze by 1H NMR (500 Hz, DMSO-d6): Tetramethyl silane 7 (TMS; 0). Efficiency of modification is determined by comparing the integrated signal of the Boc 8 protons (27H;  $\delta$  1.5-1.35) to the aromatic protons of tyrosine residues on ELP (8H each; centered 9 at  $\delta$  7.00 and 6.62).



- 10 Figure S2. NMR quantification of modification efficiency for ELP-Hydrazine: NMR of Boc-protected
- 11 ELP indicates approximatly 12.28 hydrazine moities are present per ELP chain.



Figure S3. Summary of conjugation and oxidation reactions for hyaluronan: (A) A schematic representation of our different hyaluronan (HA) components. Three variants were produced by oxidizing 14 1.5 MDa HA for 24 hr (HA24), 16 hr (HA16), and 8 hr (HA8). The other two HA variants were produced via 2-part bio-conjugation reaction wherein 100 kDa HA was first reacted to have 12% of the HA repeat groups 16 modified with an alkyne group (HA-Alkyne). A secondary reaction then conjugated a small molecule with 17 either a pendant aldehyde (HA-A) or benzaldehyde group (HA-B) to the alkyne through a copper-click 18 reaction. (B) A summary of our HA groups and relevant properties have been summarized.

#### 19 2,4,6-Trinitrobenzene Sulfonic Acid (TNBS) Assay:

The degree of HA oxidation for our oxidized HA variants (HA24, HA16, and HA8) was measured 20 using a modified TNBS assay, as previously described [1]. HA samples were dissolved in ultra 21 pure DI water at 0.6% (w/v) overnight. The following morning, 25 µL of HA was mixed with 25 µL 22 23 tert-butyl carbazate (t-BC; 30 mM; 1% aqueous trichloroacetic acid; Sigma, B91005) and left to react for 24 hr at room temperature in the dark. The following day, 5 µL of sample was diluted 24 with 995 µL sodium bicarbonate buffer (0.1M, pH 8.5). 500 µL of diluted sample was then mixed 25 with 250 µL aqueous TNBS (0.1%, diluted with 0.1M sodium bicarbonate; Thermo, 28997) and 26 27 left to react with the excess t-BC for 2 hr at 37 °C. The reaction was stopped by the careful addition 28 of 125 µL of 1 M hydrochloric acid, and 100 µL of the mixture was finally transferred into a 96 well assay plate. The absorbance at 340 nm was measured using a SpectraMax M2 microplate 29 reader. A standard calibration curve made from the aqueous t-BC solutions (1 - 20 mM; diluted 30 with 0.1M sodium bicarbonate) was used to determine the amount of unreacted t-BC and 31 to convert the result into di-aldehyde content. All measurements were repeated in triplicate. 32



Figure S4. TNBS assay for measuring degree of oxidation: The dialdehyde content of our oxidized variants (HA24, HA16, and HA8) were determined through a modified, two-part, TNBS assay. First, the aldehyde groups were reacted with a known quantity of tert-butyl carbazate (t-BC) to form a stable carbazone. The remaining unreacted, t-BC content was determined by subsequent reaction with TNBS, which forms a chromogenic derivative that can be detected at  $\lambda = 340$  nm. (A) Our measurements (in triplicate) are plotted against a standard curve. (B) The results are further summarized in a table below for ease of comparison.

#### NMR Quantification:

40 Aldehyde functionalization: dissolve ~10 mg of HA-Aldehyde (HA-A) in ~750  $\mu$ L of D<sub>2</sub>O and 41 analyze by 1H NMR (500 Hz, D<sub>2</sub>O): TMS (0); Acetyl group (3H,  $\delta$  ~1.8); water ( $\delta$  ~4.7); triazole 42 (1H;  $\delta$  ~7.85). To quantify the degree of modification, compare the integrated peak of the triazole 43 group to the acetyl group.

Benzaldehyde functionalization: dissolve ~10 mg of HA-Benzaldehyde (HA-B) in ~750  $\mu$ L of D<sub>2</sub>O and analyze by 1H NMR (500 Hz, D<sub>2</sub>O): TMS (0); Acetyl group (3H,  $\delta$  ~1.8); water ( $\delta$  ~4.7); triazole (1H) and aldehyde group (1H  $\delta$  ~9.8 ~ 9.9). To quantify the degree of modification, compare the integrated peak of the aldehyde proton to the acetyl group.



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- 48 Figure S5: NMR quantification of modification efficiency for aldehyde (HA-A) and benzaldehyde
- 49 (HA-B) modified HA: NMR of both (A) HA-A and (B) HA-B show approximately 12% of the available
- 50 carboxylic acid sites have been modified with eiteher aldehyde or benzaldehyde groups, respectively.

#### 51 Gel Permeation Chromatography:

Gel permeation chromatography (GPC) was carried out using a Dionex Ultimate 3000 instrument 52 53 (including pump, autosampler, and column compartment). Detection consisted of an Optilab TrEX (Wyatt Technology Corporation) refractive index detector operating at 658 nm and a HELEOS II 54 55 light scattering detector (Wyatt Technology Corporation) operating at 659 nm. The column used was a Superose 6 increase 10/300 GL. The eluent was PBS buffer with 30% (w/v) NaN<sub>3</sub>, 137 mM 56 NaCl, 0.0027 mM KCl, 10 mM Phosphate pH 7.3, at 0.75 mL min<sup>-1</sup> at RT. HA samples of 1.5 57 MDa, 1 MDa at 1 mg mL<sup>-1</sup>, and other analyte samples at 3 mg mL<sup>-1</sup> were dissolved in ultra-pure 58 59 DI water overnight and filtered through Millipore Express PES membrane filter with 0.22 pore size 60 prior to injection. A refractive index increment (dn/dc) value of 0.165 was applied for all samples. The time frame for integration was determined by running multiple unmodified HA samples (20 61 kDa, 40 kDa, 60 kDa, 100 kDa, 500 kDa, 1 MDa; (Lifecore), and 1.5 MDa (Sigma)) as a reference 62 and then applying the same integration time for our experimental samples (HA24, HA16, HA8, 63 HA-A, and HA-B). Values represent the approximated weight average molecular weight ( $M_{\mu}$ ). 64 65 number average molecular weight ( $M_n$ ) and the polydispersity index (PDI).

#### 66 Dynamic Light Scattering for Measuring Hydrodynamic Radius:

50  $\mu$ L of each HA sample were dissolved in PBS at a final concentration of 1% (w/v). Then, 40  $\mu$ L 67 of sample were transferred into disposable cuvettes (BrandTech, 759200) and sealed with a cap 68 69 (VWR, 47744-636) to prevent any evaporation during measurements. We then performed a size measurement using the Zetasizer Nano NS, with three replicate measurements per sample. From 70 71 these measurements, and assuming the viscosity of water, we were able to determine the z-72 average size for each sample from which we could then determine an approximate hydrodynamic radius (R<sub>h</sub>). The R<sub>h</sub> values provided presently represent the average of three independent, 73 samples. 74

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Α					ВС		
	Gel Permeation Chromatography				Unmodified HA	Experimental HA	
1	HA-Variant	Mw, obs.	Mn, obs.	PDI	i M	$\frown$	
	1.5 MDa	1.24 MDa	1.23 MDa	1.01	an		
	1 MDa	1.11 MDa	1.05 MDa	1.05	1.5 MD	Да НА-В	
	500 kDa	345.0 kDa	324.0 kDa	1.07	1.0 ME	Da	
	100 kDa	101.0 kDa	95.2 kDa	1.15			
	60 kDa	60.4 kDa	42.2 kDa	1.43		HA-A	
	40 kDa	59.7 kDa	44.0 kDa	1.36			
	20 kDa	19.8 kDa	14.2 kDa	1.40		Da	
	HA-B*	141.0 kDa	119.0 kDa	1.19	100 kD	Da HA-Alkyne	
	HA-A*	221.0 kDa	161.0 kDa	1.37			
	HA8	177.0 kDa	143.0 kDa	1.24	40 kDa	HAIO	
	HA16	80.1 kDa	57.3 kDa	1.41	20 kDa		
	HA24	56.8 kDa	43.9 kDa	1.30	5 10 15 20 25 30	5 10 15 20 25 30	
	HA-Alkyne	113.0 kDa	86.0 kDa	1.30	Elution Time (min)	Elution Time (min)	
D					E F		
	Dynamic Light Scattering				Rh VS Mw	log(Rh) vs log(Mw)	
	HA-Variant	HA-Variant Hydrodynamic Radius			<b>50</b> 1 <b>.</b>	<b>2.0</b> ]	
	1.5 MDa	45.3	9 nm ± 2.645	nm	<b>40</b>	- 4 5	
	1 MDa	1 MDa      26.69 nm ± 0.497 nm        500 kDa      15.83 nm ± 0.632 nm		E 20			
	500 kDa			nm	<u> </u>	∑ 1.0 • •	
	100 kDa	10.50	6 nm ± 0.123	nm	<u>v</u> 20		
	60 kDa	9.73 nm ± 0.518 nm			10	0.5  y = 0.411x + 0.214	
	40 kDa	9.61	1 nm ± 0.183	nm	0	0.0	
	20 kDa	8.10 nm ± 0.705 nm 11.51 nm ± 0.871 nm			0 500 1000 1500	1 2 3 4	
	HA-B*				Mw, obs. (kDa)	Log (Mw, obs.)	
	HA-A*	14.48	3 nm ± 0.234	nm			
	HA8	12.91	nm ± 1.36 n	т			
	HA16	7.45	5 nm ± 0.379	nm			
	HA24	7.09	0 nm ± 0.109	nm			
	HA-Alkyne	10.95	5 nm ± 0.526	nm			

75 Figure S6. Gel permeation chromatography (GPC) measurements of hyaluronans: To characterize 76 the molecular weight  $(M_w)$  of our different hyaluronan (HA) variants, we used gel-permeation 77 chromatography (GPC) on both reference unmodified H, and our experimental HA groups. (A) Tabulated 78 results showing the observed weight-average molecular weight ( $M_{w}$ , obs.), number-average molecular weight ( $M_n$ , obs.), and polydispersity index (PDI) are shown. GPC measurements were able to accurately 79 80 determine the molecular weight of our unmodified control groups. HA modified by copper click chemistry 81 (HA-B\* and HA-A\*) appeared to increase in molecular weight, but this is likely due to the presence of bulky side groups increasing their hydrodynamic radii and inflating the measured value. As expected, oxidized 82 HA samples (HA8, HA16, and HA24) decreased greatly from their starting molecular weight (1.5 MDa) as 83

- 84 a function of oxidation time. Representative GPC curves have been provided for (B) unmodified HA and
- 85 our (C) experimental HA groups. Note: we used a specific refractive index increment (dn/dc) value of 0.165
- 86 to calculate the approximate values for all samples. (D) Using dynamic light scattering (DLS) we measured
- 87 the hydrodynamic radii of unmodified HA controls and our modified HA groups. The hydrodynamic radius
- 88 ( $R_h$ ) and the  $M_w$ , obs. have been plotted both as (E) linear plots and (F) log-log plots.



89 Figure S7. Representative time sweeps: Following mixing, our five gel formulations (all: 2% (w/v) ELP,

90 1% (w/v) HA) were crosslinked for 30 min at room temperature. All five formulations began to gel almost

- 91 immediately following mixing, and all formulations (except HA-B) had a storage modulus (G') > loss modulus
- 92 (G") prior to the start of measurement. Typical plateau moduli following crosslinking were as follows: ~450
- 93 Pa for HA24; ~730 Pa for HA16; ~2500 Pa for HA8; ~590 Pa for HA-A; ~ 980 Pa for HA-B.



94Figure S8. Summary of Matrigel mechanics: Summary mechanics of Matrigel using our prescribed95mechanical testing protocols. We found that Matrigel has a shear modulus of ~39 Pa and a stress96relaxation,asdenotedby $t_{0.5}$ ,of~67.0s.

#### 97 Measuring Lower Critical Solution Temperature:

The lower-critical solution temperatures (LCST) of our proteins were measured by taking 98 absorbance measurements ( $\lambda$  = 300 nm) in a circular dichroism (CD) spectrometer (J-1000, 99 100 JASCO). Briefly, a 250 µL sample of 1% (w/v; 10 mg mL-1) ELP both pre- and post-101 functionalization was made by dissolving lyophilized protein in 1X PBS overnight at 4 °C on a 102 constant rotator. The following morning, 200 µL of freshly dissolved sample was loaded into a 103 pre-chilled (4 °C) 1-mm band-width guartz cuvette and then loaded into the CD spectrometer 104 sample chamber. The absorbance spectra were measured at 1 °C increments over the 105 temperature range 4 °C - 65 °C. To ensure adequate incubation, the temperature was elevated 106 at 1 °C min<sup>-1</sup> and held at each degree increment for 30 seconds. To approximate the LCST for 107 each sample, we first subtracted away a blank (PBS only) scan collected using the same protocol. Then, we normalized the curves by dividing each absorbance measurement collected by the 108 109 maximum absorbance value over the length of the measurement. The LCST temperature range 110 is defined as the approximate temperature where absorbance (norm.) = 0.5.



111 Figure S9. Lower critical solution temperature (LCST) shift: (A) The lower critical solution temperature

112 measurements of ELP pre- and post-modification with hydrazine moieties. (B) Shear moduli of crosslinked

- gels at ambient (23 °C) and body (37 °C) temperatures. 2-Way ANOVA multiple comparison's test,  $\alpha$  = 0.05,
- 114 post-hoc Bonferroni test, \*\* p < 0.01, ns = not significant.

# 115 Note: These videos have been uploaded as separate files. A brief description is listed116 below.

117	Figure S10. Representati	ve injection videos:	(A) Representative vide	eo of injecting 50 μ	IL of HA24 (dyed
118	red with food coloring for e	ase of visibility), (B) I	HA16 (dyed purple with	food coloring for e	ease of visibility),
119	(C) HA8 (dyed blue with fo	ood coloring for ease	of visibility), (D) HA-A	(dyed yellow with	food coloring for
120	ease of visibility), and (E)	HA-B (dyed green	with food coloring for	ease of visibility)	through a 30-G,
121	manually	hooked,	needle	by	hand

#### 122 Note: This video has been uploaded as a separate file. A brief description is listed below.

Figure S11. Catheter injection video of HA16: To screen the potential for translatability of our hyaluronan and elastin-like protein (HELP) gel system, we tested the injection of 700  $\mu$ L HA16 (dyed dark green for ease of visibility) through a 150-cm catheter. Injections were done by hand. Note: the beginning of the video has been accelerated (x8). Total injection time is 3 min and 8 sec. For reference, the 30-G needles used in our study had an injection length of 0.8 cm and an 0.159 mm inner diameter compared to the Codman catheter which was 150 cm in length and had an inner diameter of 0.75 mm.

#### 129 **Post-Injection Mechanics Test:**

130 50 µL of hydrogel were prepared as previously described and loaded into the back of a 30-G syringe. Hydrogels were crosslinked for 30 min and then injected, by hand, onto an ARG2 131 rheometer (TA Instruments). Post-injection recovery was monitored at 1% strain and radial 132 133 frequency of 1 rad s<sup>-1</sup> over the course of 24 hr. Heavy mineral oil was used to fill the gap between the rheometer geometry and external environment to ensure hydration over the course of all 134 measurements. Immediately following the 24-hr recovery period, the shear modulus for each 135 sample was measured at 23 °C using a non-destructive frequency sweep over the oscillatory 136 137 frequency range of 0.1 - 10 rad s<sup>-1</sup> at a fixed strain of 10%.



138 Figure S12. Recovery experiment for HA16 and HA24: 24-hr recovery experiment of 50  $\mu$ L of (A) HA16

139 and (B) HA24 post-injection at 1% strain and a frequency of 1 rad s<sup>-1</sup>.



Figure S13. Post-recovery frequency sweep: After 24-hr of recovery post-injection we measured shear stiffness of (A) HA16 and (B) HA24 via a frequency sweep at 10% strain and over a range of 0.1 rad s<sup>-1</sup> to 10 rad s<sup>-1</sup>, as detailed in the methods section.



НА-В

143 Figure S14. Representative stress and strain curves: Representative failure stress measurement tests 144 for (A) HA24, (B) HA16, (C) HA8, (D) HA-A, and (E) HA-B showing the measured strain over successive stress steps (demarcated by a semi-transparent bar and corresponding stress) and relaxation steps. The 145

- 146 x-axis, for all plots indicates time (s), plotted linearly. The stress required to induce fracture is indicated by
- 147  $\sigma_F$ . Note that the y-axis is broken into two parts: (1) 0 100 % strain and (2) 100 200% strain. The oxidized
- 148 HELP (HA24, HA16, and HA8) groups reached much lower strain values (< 50%) prior to failure than the
- 149 bio-conjugated variants, HA-A and HA-B, which experienced % strain > 100% prior to failure. Additionally,
- 150 both HA-A and HA-B had evidence of a yielding behavior (% strain >> 0 post-relaxation ( $\sigma = 0$ )) and are
- 151 denoted with a  $\sigma_y$ .

#### 152 Labeling ELP with Cyanine 5

To aid with the visualization of our HELP hydrogels post-injection, we labeled a single lysine group 153 on a portion of our ELPs with a cyanine 5 (Cy5) NHS molecule (excitation: 651, emission: 670; 154 Sigma 679011) via standard NHS-ester chemistry. Briefly, ELP was dissolved in anhydrous 155 156 DMSO at an initial concentration of 6% (w/v) at room temperature (RT) under constant rotation. 157 After dissolving, the reaction volume was diluted further with anhydrous DMF, dropwise, to a final concentration of 2% (w/v). Next, a 1 molar equivalent of activated Cy5-NHS ester was dissolved 158 in anhydrous DMF (100 mg mL<sup>-1</sup>) and added dropwise to the 2% (w/v) ELP reaction mixture. Once 159 160 added, the reaction vile was purged with nitrogen gas for 5 ~ 10 min. After purging, the reaction 161 vile was covered with aluminum foil to prevent photo bleaching and the reaction was allowed to proceed overnight at RT. 162

163 To isolate the modified ELP-Cy5, the reaction volume was added, dropwise, into ice-cold diethyl ether to a final volumetric ratio 1:5 (reaction: ether) in a solvent-safe centrifugation tube. 164 ELP-Cy5 was then collected by centrifugation (>18000 x g) at 4 °C for 30 min, decanted, and 165 166 dried overnight. The dried ELP-Cy5 product is then dissolved in ultra-pure DI water at a final concentration of 2% (w/v) and dialyzed against 4 L of ultra-pure DI. The 4 L of dialysis water was 167 168 routinely refreshed every 12 hr over the course of 3 days. The final dialyzed product was then sterilized using a 0.22-µm syringe filter, frozen, and finally lyophilized for 3 days in a sterile, filtered 169 50-mL tube. To verify the degree of functionalization, ELP-Cy5 can be characterized via NMR 170 171 (below). If the ELP molecule has been sufficiently modified (~ 1 Cy5 molecule per ELP), ELP-Cy5 172 can be further modified with a hydrazine molecule (ELP-Cy5-Hyd) following the previously listed 173 protocol and incorporated into standard HELP gels to reach a desired level of fluorescence.

174 NMR Quantification:

175 Cy5 Functionalization: dissolve ~10 mg of ELP-Cy5 in ~750 μL of deuterated DMSO and analyze 176 by 1H NMR (500 Hz, DMSO-d6): Tetramethyl silane (TMS; 0). Efficiency of modification is

- 177 determined by comparing the integrated signal of the lysine peak (δ 2.75-1.65; unmodified 28H)
- 178 to the aromatic protons of tyrosine residues on ELP (8H each; centered around  $\delta$  7.00 and 6.62,).



181 Figure S15. Ex vivo hydrogel fluorescence and microsphere concentration determination: To identify 182 an appropriate level of ELP-cyanine-5 labeling and microsphere concentration for subsequent in vivo 183 experiments, we prepared HELP gels with varying levels of (A) volume percent labeled ELP and (B) 184 concentration of fluorescent microparticles. From our ex vivo screening, we opted to use 20% (v/v) ELP-185 Cy5-Hydrazine and 1.25 mg mL<sup>-1</sup> 10-µm fluorescent particles (selections are outlined in red). Based on this 186 image, we further estimated that a 50- $\mu$ L injection would have approximately ~70,110 microspheres. Note: 187 all images represent 20-µm thick z-stacks to emulate the intended tissue thickness; all scale bars denote 188 250 μm.

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All data previously shown

Figure S16. Summary of test group mechanics: Summary of the (A) shear moduli and (B) stress relaxation rates of HA16, HA24, and Matrigel. (C) Average values are reported for ease of comparison. Note: these data have been previously shown, but are recompiled here for ease of direct comparison. Collectively, these gels established "low" (~40 Pa), "medium" (~500 Pa), and "high" (~1500 Pa) stiffness gels for our in vivo retention study. Statistical test: one-way ANOVA,  $\alpha = 0.05$ , post-hoc Tukey test. \* p < 0.05; \*\*\* p < 0.001; \*\*\*\* p < 0.0001.



195 Figure S17. Representative injection images: Representative images showing injections of 50  $\mu$ L of (A)

196 fluorescently labeled HA16, (B) fluorescently labeled HA24, and (C) Matrigel, each with a final concentration

- 197 of 1.25 mg mL<sup>-1</sup> 10-µm fluorescent microspheres. Arrows indicate evidence of material injection of the
- 198 material, indicated by the formation of a bolus and slight discoloration of the tissue.

#### **199 Injection Scoring Procedure:**

200 For injection scoring, we devised a 4-part scoring system that assessed clinically relevant 201 concerns in cardiothoracic surgical injection paradigms: (1) the observable amount of material 202 reflux (on a scale of 0 - 3 points ranging from no visibly remaining material to no material reflux), 203 (2) ease of injection (on a scale of 0 - 3 points ranging from no smoothness in injection to 204 controlled smooth injection), (3) degree of bleeding post-injection (on a scale of 0 - 3 points 205 ranging from excessive bleeding with risk of death to no spot bleeding), and (4) correct location of the injection site (on a binary scale of 0 - 1 point with the injection visually being in the incorrect 206 207 or correct location). These scores were tabulated for each gel group at both time points for a total 208 possible score of 10 points. Scores were assigned by an external observer, blind to the injection 209 group and time point.



210 Figure S18. Injection scoring: To probe the possibility that retention could be influenced by the quality of 211 our injections, we devised a 4-part scoring system that looked at: (1) the observable amount of reflux, (2) 212 ease of injection, (3) degree of bleeding post-injection, and (4) correct location of the injection site. (A) 213 These scores were tabulated for each gel group for both cohorts, 1 and 2, which correspond to 1 and 7 day 214 time points post-injection, respectively. Scores were assigned by an external observer, blind to the injection group and time point. (B – D) A breakdown of the scoring scheme has been provided for reference. Notably, 215 216 we found no significant differences in the quality of our injections between our groups. Statistical test: two-217 way ANOVA,  $\alpha = 0.05$ , post-hoc Tukey test, not significant = n.s.



**Figure S19. Tissue sectioning schematic:** *To maintain a consistent and unbiased survey of our different* test groups, all tissues were sectioned and sampled according to the above scheme. Briefly, 15 gelatincoated slides were collected from each animal for analysis. Serial slices (20-µm thick) of heart tissue were made, and every seventh slice was collected; i.e., slides were prepared from tissue slices spaced

222 approximately ~120-µm apart. Importantly, the exact number of tissue slices between each collected tissue 223 sample was recorded and used in the final quantification. In the above diagram, we have numbered each 224 tissue section (1 - 45) to reflect the general ordering of sample collection, where the first 15 samples (1-15)225 were placed at the top of each slide, the next 15 samples (16 – 30) were collected in the middle row, and 226 the last 15 were collected on the bottom row (31 - 45). By sectioning tissue this way, each slide spanned 227 approximately 4 mm in height and provided a rapid way of surveying a large amount of tissue with relatively 228 few slides needing to be imaged. From the 15 total slides, 8 (slides 1, 3, 5, 7, 9, 11, 13, and 15) were 229 collected from each animal and imaged for microspheres and elastin-like protein (ELP) according to our 230 described methods. To select a slide for Hematoxylin & Eosin (H&E) staining, we designated from those 8 231 slides one that was 'representative' (having qualities numerically and morphologically indicative of that 232 particular injection) and a slide adjacent to that was then selected for histological staining.



233 Figure S20. Example images of microsphere retention quantification: Tabulation of microsphere count 234 was carried out by automated quantification of fluorescent images. (A) First, 45, 20-µm thick, tissue 235 sections were extracted semi-periodically with a cryostat from a larger 5-mm section of rat heart collected 236 and fixed according to our previously stated methods (see Figure S19). Tissue sections were imaged with 237 aid of an inverted fluorescent microscope for evidence of injected cargo: ELP and/or fluorescent 238 microspheres. (B) From composite images we isolated the microsphere channel and saved them from each 239 tissue slice. (C) These images were then converted to binary images (using a constant threshold value) 240 and the number of microspheres were tallied for each slice using a particle counter in ImageJ. To generate 241 more accurate counts, tallies from each image slice were then checked for instances of 'clustering' (multiple 242 microspheres being counted as one) by a secondary Python script that divided the cross-sectional area of 243 each microsphere by the average area of an individual microsphere, and extra counts were added

- 244 accordingly for each slice. Note: scale bar on low magnification images represents 1 mm, scale bar on high-
- 245 magnification images represents 100 μm.

## 246 Note: This file has been uploaded separately. A brief description is listed below:

- 247
- 248 **Figure S21. Python script for post-processing microsphere count:** A Python script was written and
- 249 used to account for microsphere 'clustering' which superficially undercounted the number of microspheres
- 250 in our captured images.



Figure S22. Sex-based differences in microsphere retention: To investigate the possibility of any sexrelated differences, we plotted the average microsphere count for each hydrogel, by sex, for both (A) 1 day and (B) 7 days post-injection. Importantly, we did not observe any significant differences based on sex. Generally, it appeared that male rats had a greater degree of retention, but not in a statistically significant manner. (C) The results have been additionally summarized for reference. Statistical test: two-way ANOVA,  $\alpha = 0.05$ , post-hoc Tukey test, not significant = n.s.

# 257 Supplemental References:

Y.-C. Chen, W.-Y. Su, S.-H. Yang, A. Gefen, F.-H. Lin, In situ forming hydrogels composed of
 oxidized high molecular weight hyaluronic acid and gelatin for nucleus pulposus regeneration,
 Acta Biomater. 9 (2013) 5181–5193. https://doi.org/10.1016/j.actbio.2012.09.039.

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