

1 Supporting Information

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**Tuning Polymer Hydrophilicity to Regulate Gel Mechanics and Encapsulated Cell Morphology**

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**Supporting Information**

Table S1.....	2
Figure S1.....	3
Figure S2.....	3
Figure S3.....	4
Figure S4.....	5
Figure S5.....	6
Figure S6.....	7

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2  
3  
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<b>hMSC</b>			
<b>Temperature Profile</b>	<b>Median Actin Area (<math>\mu\text{m}^2</math>)</b>	<b>Avg. Actin Area (<math>\mu\text{m}^2</math>)</b>	<b>Standard Deviation (<math>\mu\text{m}^2</math>, N = gels)</b>
I	733.82	997.53	91.95
II	635.66	803.08	52.29
III	462.73	636.85	99.34

<b>HUVEC</b>			
<b>Temperature Profile</b>	<b>Median Actin Area (<math>\mu\text{m}^2</math>)</b>	<b>Avg. Actin Area (<math>\mu\text{m}^2</math>)</b>	<b>Standard Deviation (<math>\mu\text{m}^2</math>, N = gels)</b>
I	271.09	525.52	33.06
II	236.04	387.33	61.05
III	243.05	361.90	74.56

<b>hNPC</b>			
<b>Temperature Profile</b>	<b>Median Actin Area (<math>\mu\text{m}^2</math>)</b>	<b>Avg. Actin Area (<math>\mu\text{m}^2</math>)</b>	<b>Standard Deviation (<math>\mu\text{m}^2</math>, N = gels)</b>
I	52.11	70.94	2.04
II	54.43	76.74	5.40
III	57.90	87.73	8.48

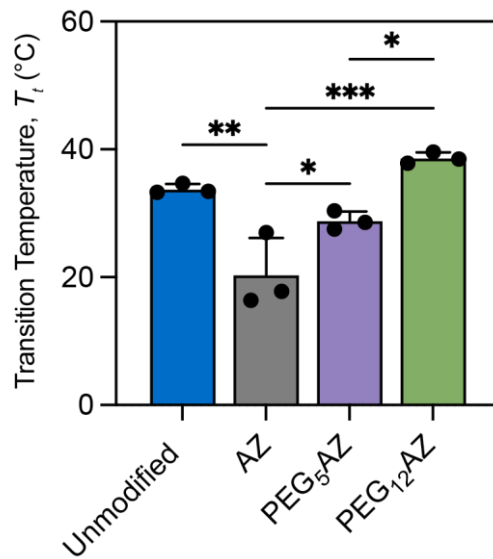
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**Supplemental Table S1.** Quantification and error measurements of actin area depicted in Figure 6. Standard deviation ( $N = \text{gels}$ ) represents the variance between gel replicates. For hMSCs,  $n = 1,063, 823,$  and  $899$  (**I**, **II**, and **III**) cells examined over  $N = 3$  independent gel samples. For HUVECs,  $n = 386, 448,$  and  $503$  (**I**, **II**, and **III**) cells examined over  $N = 3$  independent gel samples. For hNPCs,  $n = 3,357, 3,013,$  and  $3,067$  (**I**, **II**, and **III**) cells examined over  $N = 3$  independent gel samples.

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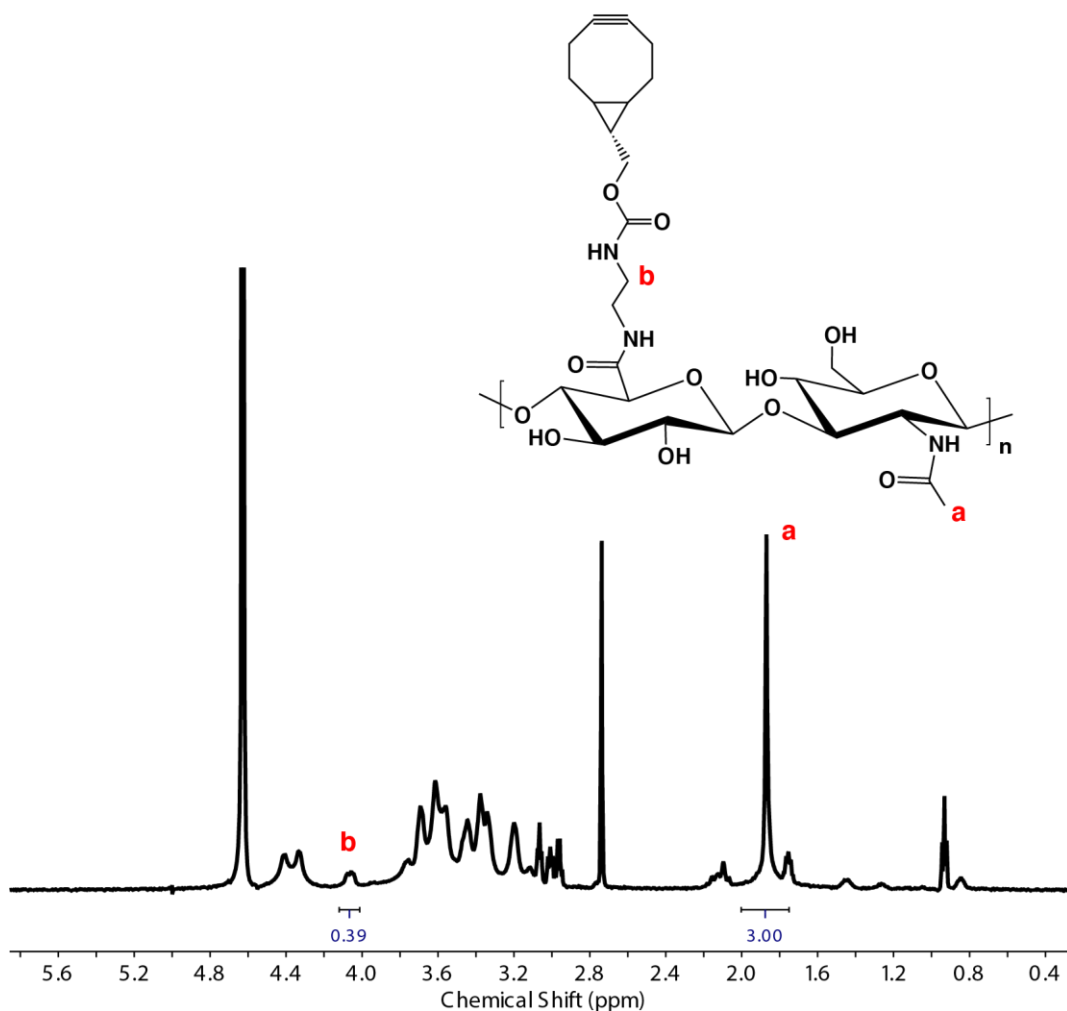


**Supplemental Figure S1.** Full amino acid sequence of cell-adhesive RGD-ELP with tag domains consisting of a T7 tag (MASMTGGQQMGG), a histidine tag (HHHHHH), and an enterokinase cleavage site (DDDDK) to permit tag removal if desired. All ELP proteins were kept intact with no tag removal for these studies.



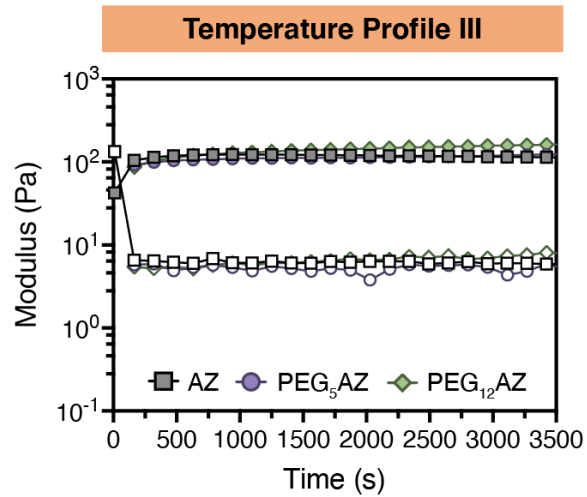
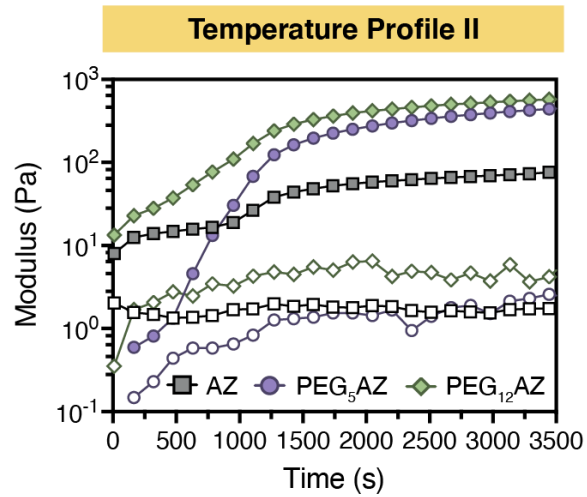
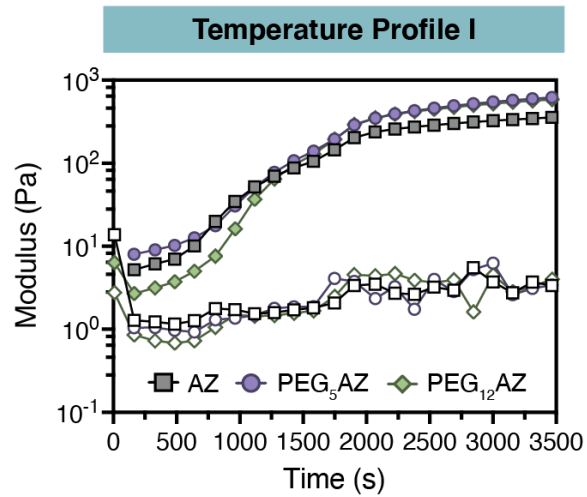
**Supplemental Figure S2.** Quantification of ELP transition temperatures before and after coupling reaction ( $n = 3$ , data are averages  $\pm$  standard deviation, \* $p < 0.1$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ , one-way ANOVA with Tukey *post hoc* test).

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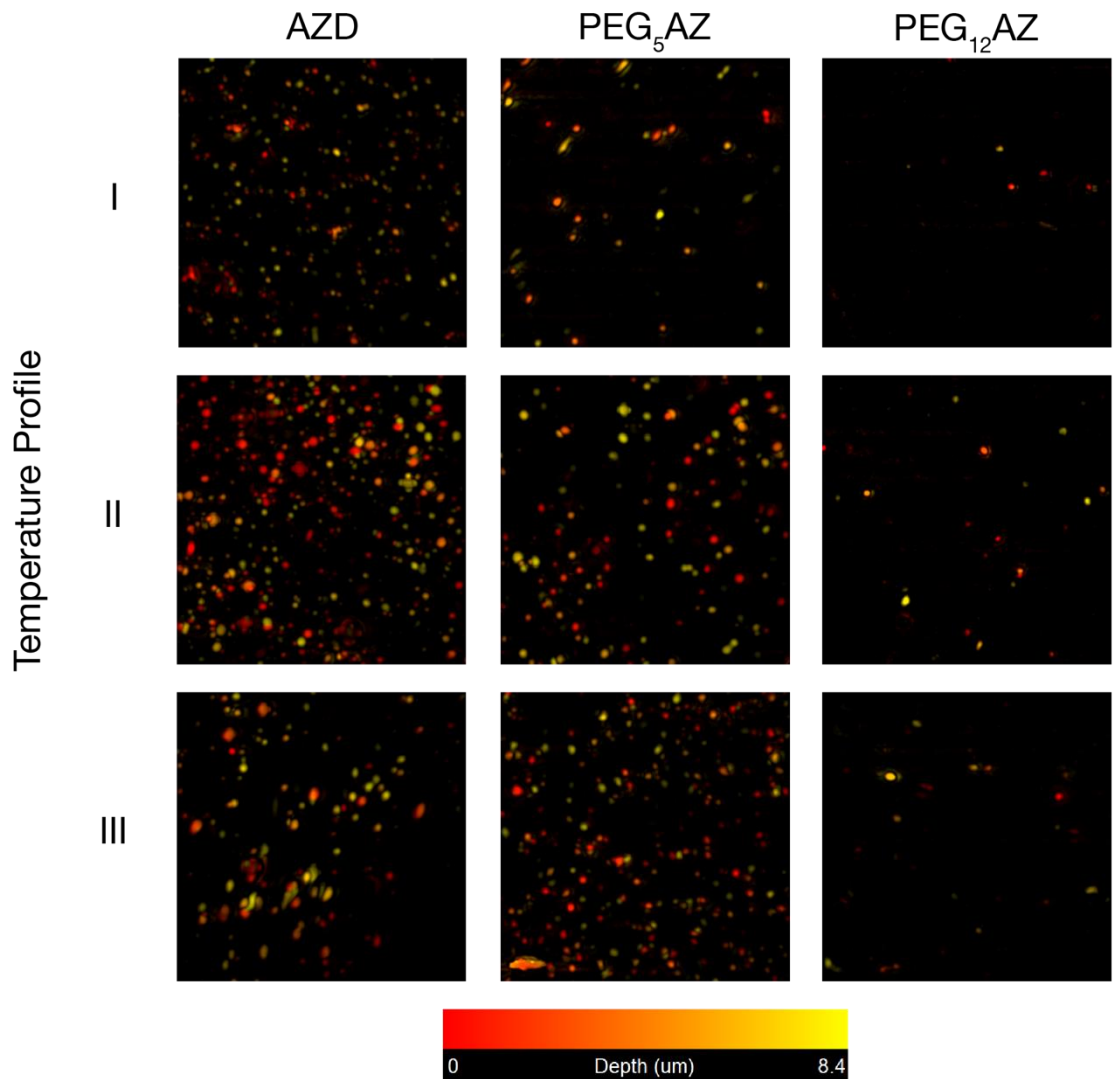
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**Supplemental Figure S3.** <sup>1</sup>H NMR 600 MHz spectrum of hyaluronic acid conjugated with bicyclononyne.



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 2 **Supplemental Figure S4.** Representative gelation time sweep data showing storage moduli  
 3 ( $G'$ , filled symbols) and loss moduli ( $G''$ , open symbols) during SPAAC-crosslinking of HELP-  
 4 SPAAC hydrogels for the three temperature profiles and each of the HELP-SPAAC gel  
 5 formulations.

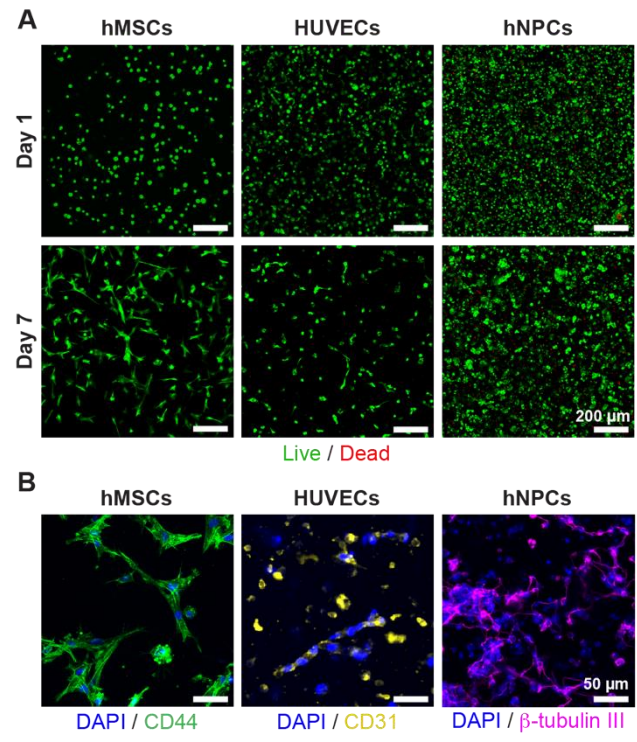
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**Supplemental Figure S5.** Representative maximum projections of CARS images of HELP-SPAAC gels crosslinked with all three temperature profiles. Regions of bright intensity are protein-rich ELP aggregates.

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**Supplemental Figure S6.** HELP-SPAAC hydrogels support 3D cell culture. A) Live/Dead staining of hMSCs, HUVECs, and hNPCs cultured within HELP-SPAAC hydrogels (formulated with PEG<sub>5</sub>AZ and temperature profile I) indicated high viability for all cell types 1 and 7 days post-encapsulation. B) Representative confocal images of immunostained hMSCs, HUVECs, and hNPCs cultured within HELP-SPAAC hydrogels (formulated with PEG<sub>5</sub>AZ and temperature profile I). After 7 days in culture, hMSCs stain positive for the MSC surface marker CD44 (green), HUVECs stain positive for the endothelial marker CD31 (yellow), and hNPCs stain positive for the neuronal marker β-tubulin III (pink). Nuclei are counter-stained with DAPI (blue).