Supporting Information

From 4-Arm Star Proteins to Diverse Stimuli-Responsive Molecular

Networks Enabled by Orthogonal Genetically Encoded Click Chemistries

Hong Kiu Francis Fok, ^{‡a} Zhongguang Yang, ^{‡a} Bojing Jiang, ^a and Fei Sun ^{*ab}

^aDepartment of Chemical and Biological Engineering, The Hong Kong University of Science and Technology, Clear Water Bay, Kowloon, Hong Kong SAR, China. ^bGreater Bay Biomedical InnoCenter, Shenzhen Bay Laboratory, Shenzhen 518036, China.

Name	Molecular	Protein sequence
	Weight (Da)	
BPB	47244	MKGSSHHHHHHVDIPTTENLYFQGAMVDTLSGLSSEQGQSGD MTIEEDSATHIKFSKRDEDGKELAGATMELRDSSGKTISTWISDG QVKDFYLYPGKYTFVETAAPDGYEVATAITFTVNEQGQVTVNGK ATKGDAHIDGPQGIWGQLDGHGVGVPGVGVPGVGVPGVGVPG GVGVPGVGVPGVGVPG
BQB	59482 (pACYCDuet-1 construct), 58497 (pQE80I	pACYCDuet-1 construct: MGSSHHHHHHSQDPHHHHHVDIPTTENLYFQGAMVDTLSGLS SEQGQSGDMTIEEDSATHIKFSKRDEDGKELAGATMELRDSSGK TISTWISDGQVKDFYLYPGKYTFVETAAPDGYEVATAITFTVNEQ GQVTVNGKATKGDAHIDGPQGIWGQLDGHGVGVPGVGVPGV GVPGEGVPGVGVPGVGVPGVGVPGVGV
	construct)	HPDYPDIYGAIDQNGTYQNVRTGEDGKLTFKNLSDGKYRLFENS EPAGYKPVQNKPIVAFQIVNGEVRDVTSIVPQDIPATYEFTNGKH YITNEPIPPKMHTSVPGVGVPGVGVPGEGVPGVGVPGVGVPGV GVPGVGVPGEGVPGVGVPGV

Table S1. Amino acid sequences of proteins used in this study.

		KGDAHIDGPQGIWGQLEWKK
		pOF80l construct
		MKGSSHHHHHHVDIPTTENI YEOGAMVDTI SGI SSEOGOSGD
		MTIEEDSATHIKESKRDEDGKEI AGATMEI RDSSGKTISTWISDG
		GVGVPGVGVPGVGVPGVGVPGFGVPGVGVPGVGVPGVGV
		GEGVPGVGVPGVGVPGVGVPGVGVPGVGVPGVGVPGVGV
ΔΔ	19708	
	15700	
		GVGVPGEGVPGVGVPGVGVPGVGVPGEGVPGVGVPG
A-SNAC-A	19927	MKGSSHHHHHHVDAHIVMVDAYKPTKLDGHGVGVPGVGVPG
		VGVPGEGVPGVGVPGVGVPGVGVPGEGVPGVGVPGV
		GVPGVGVPGVGVPGEGVPGVGVPGVGELGSHHWTSVPGVGV
		PGVGVPGEGVPGVGVPGVGVPGVGVPGVGVPGEGVPGVGVP
		GVGVPGVGVPGVGVPGEGVPGVGVPGVGVPGGLLDAHIVMV
		DAYKPTKLEWKK
AEC	32182	MKGSSHHHHHHVDAHIVMVDAYKPTKLDGHGVGVPGVGVPG
		VGVPGEGVPGVGVPGVGVPGVGVPGVGVPGEGVPGVGVPGV
		GVPGVGVPGVGVPGEGVPGVGVPGVGELPEDLGTGLLEALLRG
		DLAGAEALFRRGLRFWGPEGVLEHLLLPVLREVGEAWHRGEIGV
		AEEHLASTFLRARLQELLDLAGFPPGPPVLVTTPPGERHEIGAML
		AAYHLRRKGVPALYLGPDTPLPDLRALARRLGAGAVVLSAVLSEP
		LRALPDGALKDLAPRVFLGGQGAGPEEARRLGAEYMEDLKGLAE
		ALWLPRGPEKEAI



Figure S1. SDS-PAGE analysis of reconstitution of purified BPB and BQB.



Figure S2. Time-sweep tests of Snoop-Spy hydrogels (8 and 15 wt %). The shear frequency and strain were fixed at 1 rad/s and 10%, respectively.



Figure S3. Strain-sweep tests of Snoop-Spy hydrogels (8 and 15 wt %). The shear frequency was fixed at 10 rad/s.



Figure S4. Additional representative z slice images of encapsulated 3T3 fibroblasts (Live (green)/Dead (red) staining) by Snoop-Spy hydrogels. Data are shown as mean \pm SD (n = 3). *, p \leq 0.05. Scale bar: 100 µm.



Figure S5. Encapsulation of 3T3 fibroblasts by Snoop-Spy Hydrogels for 5 days. Live (green)/Dead (red) staining was performed to assess cell viability. Representative 3D rendering and *z* slice are shown.



Figure S6. SDS-PAGE analysis of 4-arm star proteins including (SpyCatcher)₄Snoop and $(CarH_c)_4$ Snoop. $(CarH_c)_4$ Snoop was obtained as a crude product from the reaction of (SpyCatcher)_4Snoop + SpyTag-ELP-CarH_c at a 1:5 molar ratio. AEC, SpyTag-ELP-CarH_c.



Figure S7. Frequency- and strain-sweep tests of $(CarH_c)_4$ Snoop hydrogels (7 wt %). The shear frequency and strain were fixed at 10 rad/s and 10%, respectively.

Video S1. Light-induced release of cells from (CarH_c)₄Snoop hydrogel (20X speed).