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## 2 **Tuning the Hydrophobicity of a Hydrogel by** 3 **Self-assembly of Polymer Cross-linkers**

4 **Hee-Jin Kim**<sup>1</sup>, **Sungwoo Cho**<sup>1</sup>, **Seung Joo Oh**<sup>1</sup>, **Sung Gyu Shin**<sup>1</sup>, **Hee Wook Ryu**<sup>1</sup>, and **Jae Hyun**  
5 **Jeong**<sup>1,\*</sup>

6 <sup>1</sup> Department of Chemical Engineering, Soongsil University, 369, Sangdo-Ro, Dongjak-Gu, Seoul, 06978,  
7 Republic of Korea; kimhj0706@ssu.ac.kr (H.-J.K.); som113@ssu.ac.kr (S.C.); ohsj0610@ssu.ac.kr (S.J.O.);  
8 whitegd45@ssu.ac.kr (S.G.S.); hwryu@ssu.ac.kr (H.W.R.)

9 \* Correspondence: nfejhh@ssu.ac.kr; Tel.: +82-2-828-7043

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11 **Abstract:** Hydrogels incorporated with hydrophobic motifs have received considerable attention to  
12 recapitulate the cellular microenvironments, specifically for the bio-mineralization of a 3D matrix.  
13 Introduction of hydrophobic motifs into a hydrogel often results in irregular arrangement of the  
14 motifs, and further phase separation of hydrophobic domains, but limited efforts have been made  
15 to resolve this challenge in the hydrophobically-modified hydrogel. Therefore, this study presents  
16 an advanced integrative strategy to incorporate hydrophobic domains regularly in a hydrogel by  
17 self-assembling of polymer cross-linkers, building blocks of a hydrogel. Self-assemblies between  
18 polymer cross-linkers were examined as micro-domains to incorporate hydrophobic motifs in a  
19 hydrogel. The self-assembled structures in a pre-gelled solution were confirmed with the  
20 fluorescence analysis and the hydrophobicity of a hydrogel could be tuned by incorporating the  
21 motifs in a controlled manner. Overall, the results of this study would greatly serve to tuning  
22 performance of a wide array of hydrophobically-modified hydrogels in drug delivery, cell  
23 therapies and tissue engineering.

24 **Keywords:** Hydrogel; Hydrophobicity; Self-assembly; Degree of swelling

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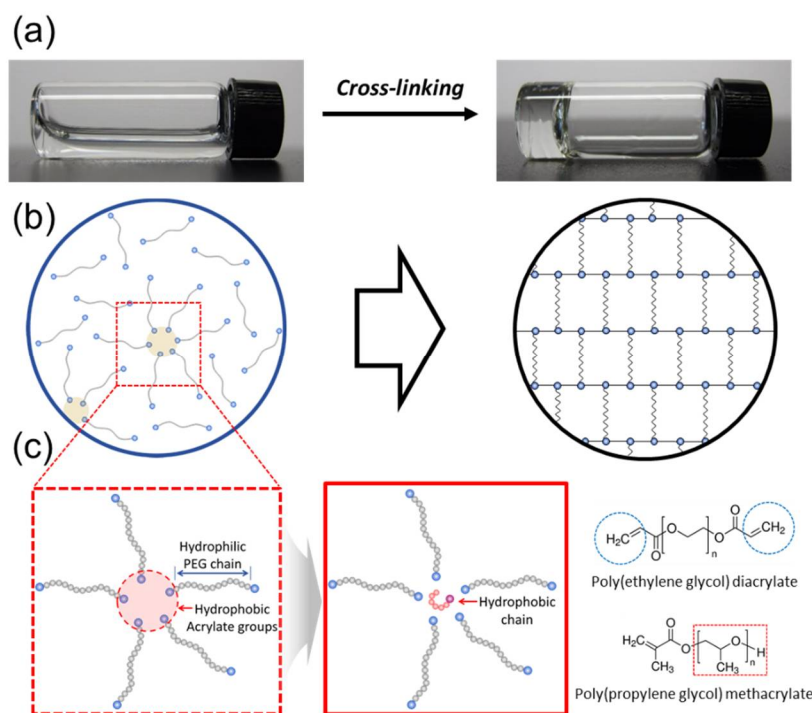
### 26 **1. Introduction**

27 Hydrogels have been extensively studied for use in various biomedical applications including  
28 drug delivery, tissue engineering and recently, Bio-MEMS(bio-microelectromechanical system)  
29 [1-3]. The successful use of the hydrogels in these applications greatly relies on their physical and  
30 chemical properties for maximizing their functionality [4-7]. For example, mineralized hydrogel  
31 systems are being increasingly studied to understand bio-mineralization processes related to the  
32 development, repair, regeneration and remodeling of bone tissue [8,9]. In here, the hydrogels  
33 incorporated with hydrophobic motifs are required to recapitulate the hydrophobic/hydrophilic  
34 microenvironments in mineralized bone tissue [10,11]. However, incorporating of hydrophobic  
35 motifs into a hydrogel often results in irregular arrangement of the motifs, and further phase  
36 separation of hydrophobic domains, but limited efforts have been made to resolve this challenge in  
37 the hydrophobically-modified hydrogel.

38 Therefore, we hypothesized that self-assemblies of polymer cross-linkers in a pre-gelled  
39 solution would allow us to incorporate hydrophobic motifs regularly as micro-domains in a  
40 hydrogel. This hypothesis was examined using a model system for a hydrophobically-modified  
41 hydrogel formed from the cross-linking of poly(ethylene glycol) diacrylate (PEGDA). The  
42 self-assembled structures of PEGDAs in a pre-gelled solution were confirmed with the fluorescence  
43 analysis (Fig. 1(b)). Then, poly(propylene glycol) methacrylate (PPGMA) with varying of mass  
44 fraction was used as model hydrophobic motif (Fig. 1(c)). The effects of the hydrophobic domains  
45 incorporated into a hydrogel were studied by measuring swelling ratio and contact angle of a  
46 hydrogel. The underlying mechanism by which micro-domains provided by self-assembling of

47 polymer cross-linkers tuned the hydrophobicity in a hydrogel was examined by evaluating average  
 48 pore size of a hydrogel and characterizing by fluorescence analysis. Overall, this study demonstrates  
 49 a novel strategy to create a hydrogel incorporated with hydrophobic chains in a controlled manner  
 50 by self-assembling of polymer cross-linkers.

51 **Figure 1.** Schematic description of hydrogel forming from polymer cross linkers in a pre-gelled  
 52 solution (a). The internal structures of self-assemblies associated with polymer cross-linkers in a



53 pre-gelled solution (b) and incorporating of hydrophobic chains using the self-assemblies (c).

## 54 2. Materials and Methods

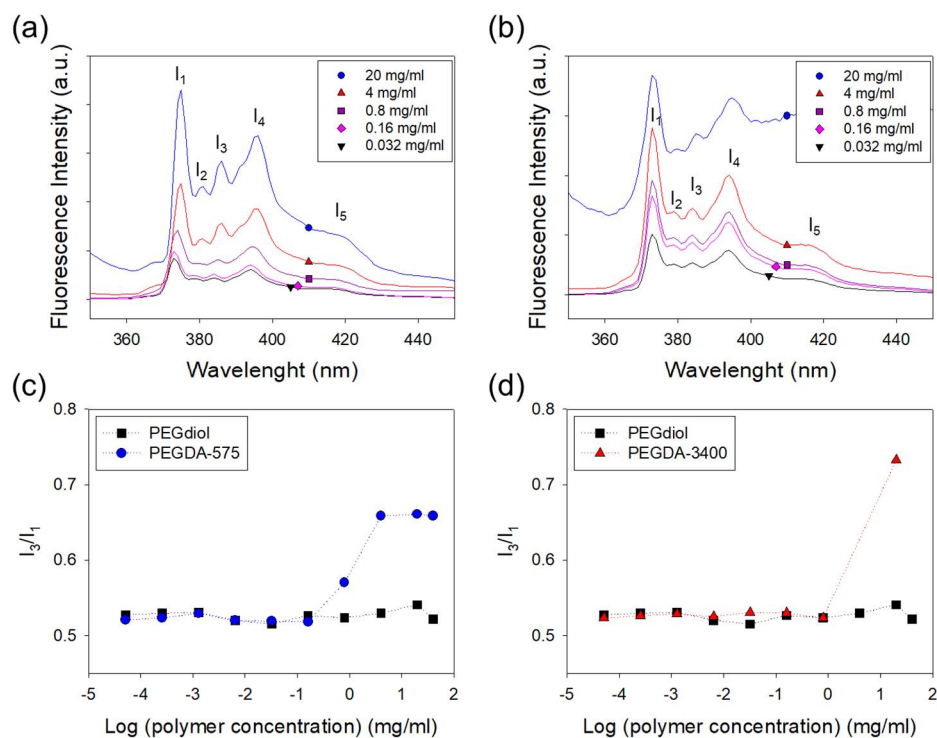
### 55 2.1. Fluorescent analysis of self-assembling of polymer cross-linkers in a solution

56 The self-assemblies of polymer cross-linkers in a pre-gelled solution were investigated using a  
 57 pyrene probe. [12,13]. The pyrene (Sigma) was dissolved in acetone to prepare a stock solution with  
 58 concentration of  $6.0 \times 10^{-4}$  M. Polymer cross-linker pre-gelled solutions were prepared in DI  
 59 (deionized) water (2 mL) by poly(ethylene glycol) diacrylate of  $M_n$  575 g/mol (PEGDA-575, Sigma)  
 60 and  $M_w$  3400 g/mol (PEGDA-3400, Sigma) respectively. In parallel, acrylate group-free poly(ethylene  
 61 glycol) of  $M_n$  3350 g/mol (PEGdiol, Sigma) were dissolved in DI water as a control. Then, the pyrene  
 62 solution was dropped into the polymer solutions with varying of polymer concentrations. The  
 63 mixture of the polymer solution and pyrene was sonicated for 10 min to ensure dispersion of pyrene  
 64 in the polymer solution. The mixture was further incubated at room temperature for at least 12 hour  
 65 in the dark, so the pyrene was preferentially associated with hydrophobic domains of polymers. The  
 66 mixture loaded in a quartz cuvette was excited at wavelength of 330nm and resulting emission  
 67 spectrum was obtained using photo luminescence (QM40, Photon Technology International). The  
 68 band-width was adjusted to 2.0 nm for both excitation and emission.

### 69 2.2. Hydrogel preparation

70 Pre-gelled solutions were prepared by mixing 10 wt% PEGDA-575 which has hydrophilic  
 71 property and poly(propylene glycol) methacrylate (PPGMA,  $M_n$  of 375 g/mol, Sigma), acting  
 72 hydrophobic chain, by increasing of mass fraction under fixed total polymer concentration, 10 wt%.  
 73 The PEGDA was dissolved in DI water at 40 wt% stock solution, and PPGMA was dissolved in  
 74 dimethyl sulfoxide (DMSO, Sigma Aldrich) at 20 wt% stock solution. The photo-initiator,

75 2-hydroxy-4'-(2-hydroxyethoxy)-2-methylpropiophenone (Irgacure 2959, Sigma) was dissolved  
 76 DMSO at 10 wt% stock solution and added to the 1mL pre-gelled solution to form 0.2 wt% as the  
 77 final concentration. First, the 1mL of pre-gelled solution mixed by vortex mixer. Second the mixed  
 78 pre-gelled solution was cast between glass plates with spacer of 1mm of thickness. Then, the cast  
 79 pre-gelled solution was exposed by UV lamp (365 nm, VL-4.LC, VILBER LOURMAT) for gelation  
 80 about 10 min, after then, gel was punched with 8mm diameter. Gel disks were immersed DI water to  
 81 remove unreacted hydrophilic residuals and then, subsequently immersed DMSO to residual  
 82 hydrophobic polymers in the cross-linked gel disk. Finally, gel disk was immerged to exchange  
 83 DMSO to DI water and further incubated in DI water for 24 h before characterizations described  
 84 below.  
 85



86  
 87 **Figure 2.** Fluorescence emission spectra of pyrene loaded in the (a) PEGDA-575 solution, (b)  
 88 PEGDA-3400 solution were captured at various polymer concentrations. The ratio of third-to-first  
 89 vibrational fine structure ( $I_3/I_1$ ) in PEGDA increased while that of PEGdiol kept constant. The  
 90 increase of  $I_3/I_1$  ratio of pyrene in the presence of polymer indicated that polymers formed  
 91 aggregation. In (c) and (d), ● represents the solution of PEGDA-575, ▲ the solution of PEGDA-3400,  
 92 and ■ the solution of PEGdiol.

### 93 2.3. Hydrogel characterization

94 The swelling ratio of the hydrogel at equilibrium was determined by measuring the weight of  
 95 the hydrated gel and that of the dried gel. The degree of swelling ( $Q$ ), defined as the reciprocal of the  
 96 volume fraction of a polymer in a hydrogel ( $v_2$ ), was calculated from the following equation (1),

$$Q = v_2^{-1} = \rho_p \left[ \frac{Q_m}{\rho_s} + \frac{1}{\rho_p} \right] \quad (1)$$

97 where  $\rho_s$  is the density of water,  $\rho_p$  is the density of polymer and  $Q_m$  is the swelling ratio, the mass  
 98 ratio of swelled gel to the dried gel.

99 The average pore size ( $\xi$ ) of hydrogel was calculated from the polymer volume fraction ( $v_{2,s}$ )  
 100 and the unperturbed mean-square end-to-end distance of the monomer unit ( $r_o^2$ ) using equation (2)  
 101 and (3):

$$\xi = (v_{2,s}^{-1/3})(r_0^{-2})^{1/2} \quad (2)$$

$$(r_0^{-2}) = l(2\frac{\bar{M}_c}{\bar{M}_r})^{1/2} C^{1/2} = l(2n)^{1/2} C^{1/2} \quad (3)$$

102

103 where  $l$  is the average value of the bond length between C–C and C–O bonds in the repeatable unit  
 104 of PEG [–O–CH<sub>2</sub>–CH<sub>2</sub>–], which is taken as 1.46 Å ;  $M_c$  is the average molecular mass between  
 105 cross-links in the network;  $M_r$  is the molecular mass of the PEG repeating unit;  $n$  is the number of  
 106 repeat unit, which is taken as 7 (PEGDA  $M_n$  of 575 g/mol);  $C$  is the characteristic ratio for PEG, which  
 107 is taken here as 4 [14].

108

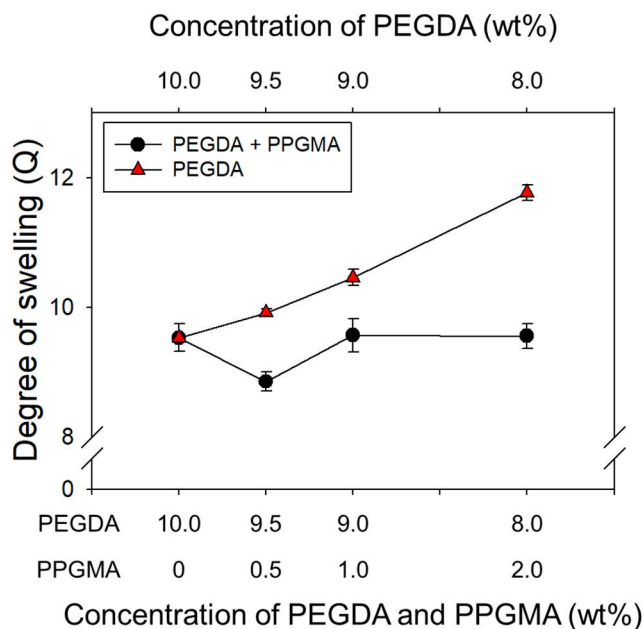
109 The inner micro-structures of the prepared final hydrogels were analyzed with SEM  
 110 (FE-Scanning electron microscope, JEOL-7001F). Values of the water contact angle ( $\theta_w$ ) on the surfaces  
 111 of the hydrogels were measured by depositing a drop of water (4.0  $\mu$ L) under atmospheric condition  
 112 with DSA100 (KRÜSS). Also, bovine serum albumin (BSA, Sigma) was used as a model protein to  
 113 evaluate the protein release rate from the hydrophobically-modified hydrogels. BSA amount released  
 114 from the hydrogel was quantitatively measured using Pierce™ BCA protein assay kit (Thermo  
 115 Scientific) according to the manufacturer's instructions. The absorbance was measured at 562 nm using  
 116 ELISA (Multiskan GO, Thermo Scientific, USA). The BSA release profile obtained was fitted to the  
 Ritger-Peppas equation [15].

$$\frac{M_t}{M_\infty} = k \cdot t^n \quad (4)$$

117

118 where  $M_t$  is the cumulative amount of protein released at the time,  $t$ ;  $M_\infty$  is the total amount of protein  
 119 in the hydrogels;  $k$  is the kinetic rate constant; and  $n$  is the exponent related to the release mechanism.

119



120

121 **Figure 3.** The degree of swelling ( $Q$ ) of hydrogel was calculated 10% PEGDA-575 hydrogels with  
 122 decrease mass fraction of PEGDA ( $\blacktriangle$ ), and with increase mass fraction of PPGMA while keeping  
 123 total polymer concentration constant ( $\bullet$ ). The hydrophobic chain in the hydrogel was the major  
 124 cause of decreasing degree of swelling from 10 to 9.5%.

125

126 **3. Results and Discussion**127 *3.1. Analysis of self-assembling of polymer cross-linkers in a solution*

128 This study presents an effective method to incorporate hydrophobic domains regularly in a  
 129 hydrogel by self-assembling of polymer cross-linkers. First, the self-assembled structures formed in  
 130 a pre-gelled solution were confirmed with the fluorescence analysis. The hydrophobic association  
 131 between the polymer cross-linkers with acrylate groups which are slightly hydrophobic was  
 132 examined with a ratio of the third-to-first vibrational fine structure ( $I_3/I_1$ ) in the fluorescence  
 133 spectrum of pyrene probe (Fig. 2). Generally, the  $I_1$  peak arises from the transition that can be  
 134 enhanced by the distortion of the  $\pi$ -electron cloud [12,13]. Therefore, as the microenvironment of  
 135 pyrene becomes more polar, the  $I_1$  peak becomes more notable at the expense of other peaks ( $I_3$ ).  
 136 That means the ratio of  $I_3/I_1$  represents the degree of self-assemblies between acrylate groups linked  
 137 to the polymer cross-linkers.  $I_3/I_1$  of the PEGDA polymer cross-linker solutions increased as the  
 138 PEGDA concentration exceeded a critical concentration (critical aggregation concentration, CAC)  
 139 which means the self-assemblies are formed at this point, as shown in Figure 2. In contrast,  $I_3/I_1$  of the  
 140 PEGdiol polymer without acrylate groups was independent of PEGdiol concentration. The  $I_3/I_1$  of  
 141 the PEGdiol solution was approximately 0.5, which is characteristic value for pyrene dispersed in  
 142 water. Therefore, this fluorescent analysis demonstrated that polymer cross-linkers with acrylate  
 143 groups are self-assembled in an aqueous solution because of the hydrophobic association between  
 144 acrylate groups [13].  
 145  
 146

**Table 1.** Composition and characterization of the hydrophobically-modified hydrogel.

	Composition		Theoretical pore-size (Å)	Experimental degree of swelling	Contact angle (°)
	<sup>2</sup> PEGDA	<sup>3</sup> PPGMA			
<sup>1</sup> HMH-1	10.0	0.0	23.16	9.53 ± 0.21	44.30 ± 4.00
HMH-2	9.5	0.5	23.47	8.85 ± 0.15	53.40 ± 0.96
HMH-3	9.0	1.0	23.89	9.56 ± 0.26	54.84 ± 0.30
HMH-4	8.0	2.0	24.85	9.55 ± 0.19	56.87 ± 1.05

147 <sup>1</sup> Hydrophobically modified hydrogel.

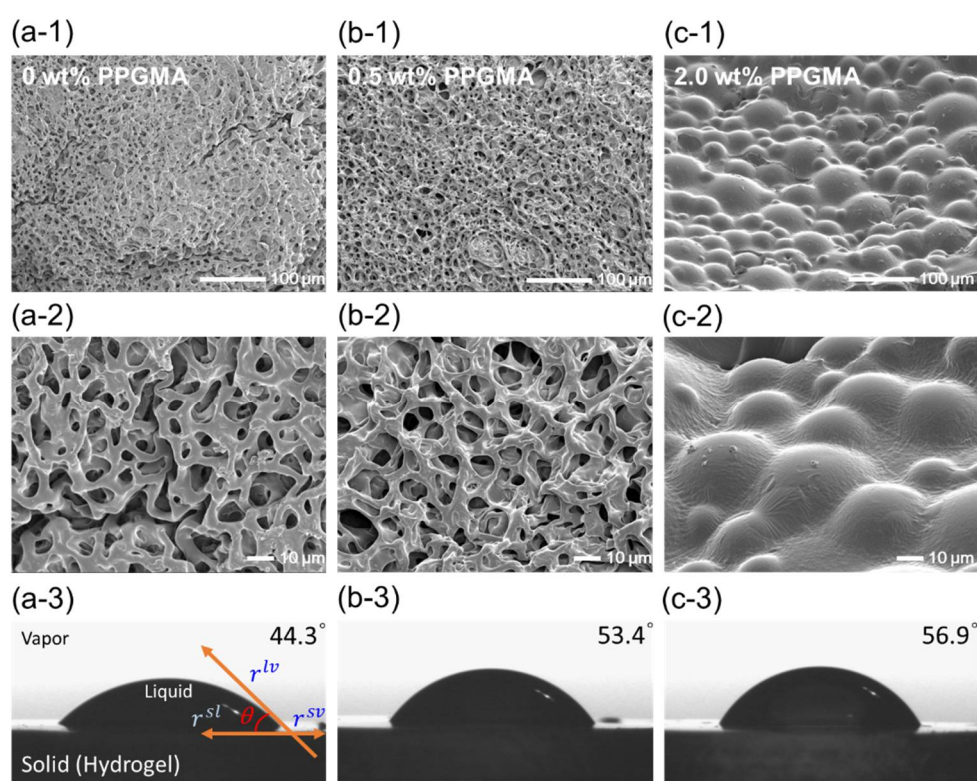
148 <sup>2</sup> Poly(ethylene glycol) diacrylate (wt%).

149 <sup>3</sup> Poly(propylene glycol) methacrylate (wt%).

150 *3.2. Effects of the hydrophobic domains incorporated into a hydrogel*

151 The hydrophobically-modified hydrogels were prepared *via* in situ radical polymerization of  
 152 self-assembled PEGDAs and PPGMAs with varying of mass fraction. As pointed out above, the  
 153 concentration of polymer cross-linkers used to form hydrogel was higher than the CACs of  
 154 cross-linkers as shown in Fig. 2(c). Figure 3 shows that in general, the degree of swelling ( $Q$ ) of  
 155 hydrogels increases as the concentration of pure PEGDAs decreases due to the increase of average  
 156 pore-size ( $\xi$ ) of hydrogel. For example, 2.31 nm of  $\xi$  in 10 wt% of PEGDAs increases to 2.48 nm of  
 157 that in 8.0 wt% of PEGDAs as shown in Table 1, calculated using the eqn. (1). However,  
 158 interestingly, in case of the hydrophobically-modified hydrogels, the  $Q$  of hydrogel was decreased  
 159 with increasing PPGMA portion from 0 to 0.5 wt%, despite of the decrease of PEGDA concentration.  
 160 And then, the  $Q$  was increased with increasing PPGMA from 0.5 to 2.0 wt%. These results indicated  
 161 that PPGMA regulates the degree of swelling of the hydrogel at a mass fraction of less than 0.5 wt%  
 162 of PPGMA as a hydrophobic repulsion. In contrast, above 0.5 wt% of the mass fraction of PPGMAs,  
 163 the decreased mass fraction of PEGDAs has a predominant role to the  $Q$  of hydrogel. Note that  
 164 PPGMAs are not related to regulate the  $\xi$  of hydrogel, because PPGMA molecule have a single  
 165 acrylate group which is not acting as a cross-linker. However, the  $\xi$  of hydrogel from 0 to 0.5 wt% of  
 166 PPGMAs was decreased despite of the decrease of PEGDAs' concentration. As addressed above, this  
 167 result is attributed to hydrophobic repulsion by PPGMA.

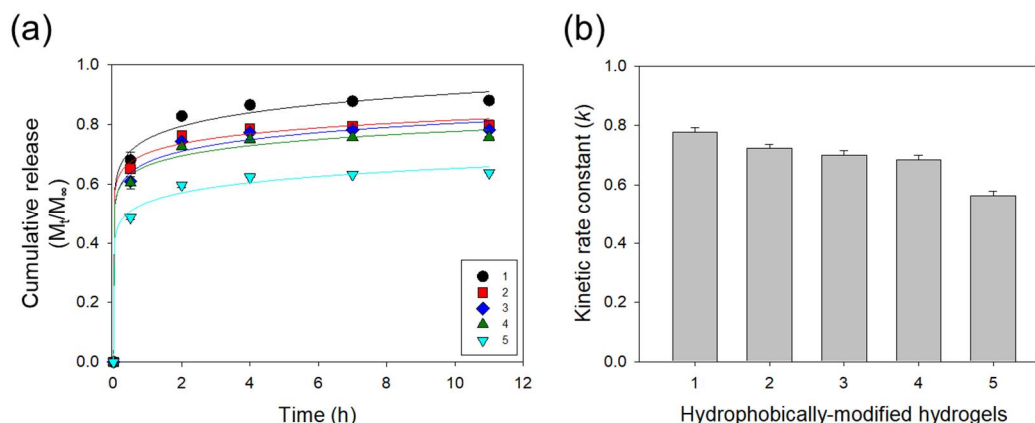
168 The inner micro-structures of hydrophobically-modified hydrogels (HMHs) were examined  
 169 with freeze-dried gels by SEM. Figure 4 shows the images of hydrogels depending on the  
 170 introduced amounts of PPGMAs. Structural difference between pure PEGDA hydrogel (HMH-1)  
 171 and HMH-2 at 0.5 wt% of PPGMA was not significant and their 3D networked microstructures were  
 172 both well formed. However, HMH-4 at 2.0 wt% of PPGMAs exhibited relatively large hydrophobic  
 173 micro-domains (Fig. 4a-3 & 4b-3). It was seen to be taken place of a phase separation between  
 174 hydrophilic PEGDA and hydrophobic PPGMA molecules. It means that a certain amount of  
 175 PPGMA in hydrogel has functioned as hydrophobic repulsion. Hydrophobic parts, PPGMAs over a  
 176 certain amount which is likely the loading capacity of the self-assemblies of PEGDAs in pre-gelled  
 177 solution, caused the phase separation. Also, water contact angles ( $\theta_w$ ) measured at the surface of  
 178 hydrogel increased with increasing of incorporating PPGMAs (Fig. 4). As a result, the  
 179 hydrophobicity of a hydrogel could be tuned by incorporating the hydrophobic motifs into the  
 180 self-assembled structures in a pre-gelled solution.  
 181



182 **Figure 4.** FE-SEM images and water contact angles of (a) pure PEGDA hydrogel (HMH-1),  
 183 hydrophobically-modified hydrogel, HMH-2 (b), and HMH-4 (c) at 2.0 wt% of PPGMAs.

184 BSA was encapsulated into the HMHs using PEGDA-3400 to evaluate the effects of the  
 185 hydrophobic motifs on protein release rate. The release rate of BSA from the HMH gel was  
 186 quantified by measuring the amount of BSA released into the incubation media on a daily basis  
 187 over ten days (Fig. 5). Increasing the hydrophobicity from 0 to 2 wt% of PPGDAs significantly  
 188 decreased the amount of BSA initially released from the HMH gels (Fig. 5(a)). The cumulative mass  
 189 fraction of BSA released from HMHs over 12 hours,  $M_t/M_\infty$ , was fitted with the Ritger-Peppas  
 190 equation to calculate a kinetic rate constant ( $k$ ) (Fig. 5(b)). The  $k$  of the HMHs decreased by 30% as  
 191 PPGMAs was increased from 0 to 2 wt%. The hydrogels incorporated with hydrophobic motifs  
 192 would be useful to study for releasing and secreting of soluble and insoluble factors in the cellular  
 193 microenvironment. In summary, the self-assemblies formed with polymer cross-linkers in a  
 194 pre-gelled solution could be a place of incorporation of hydrophobic motifs regularly and stably in  
 195 thermodynamics upon the critical range. However, above this critical point, the self-assemblies

196 would not provide enough room for the hydrophobic parts, so that the phase will be separated and  
 197 consequently irregular hydrophobic macro-domains in size and arrangement will be generated.  
 198



199 **Figure 5.** (a) Cumulative release profiles of BSA from the hydrogels. The solid lines represent the  
 200 fitting curves used to quantify the kinetic rate constant ( $k$ ) of release presented in (b).

## 201 5. Conclusions

202 Taken together, this study presents a new strategy to incorporate hydrophobic domains  
 203 regularly in a hydrogel by self-assembling of polymer cross-linkers, building blocks of a hydrogel.  
 204 Self-assemblies between polymer cross-linkers were examined as micro-domains to incorporate  
 205 hydrophobic motifs in a hydrogel. The self-assembled structures in a pre-gelled solution were  
 206 confirmed with the fluorescence analysis and the hydrophobicity of a hydrogel could be tuned by  
 207 incorporating the motifs in a controlled manner. Overall, the results of this study would greatly  
 208 serve to tuning performance of a wide array of hydrophobically-modified hydrogels in drug  
 209 delivery, cell therapies and tissue engineering.  
 210

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 212 Jeong; formal analysis, H.J. Kim, S.J. Oh, S.G. Shin; data curation, H.J. Kim, S.J. Oh, J.H. Jeong; writing—original  
 213 draft preparation, H.J. Kim, J.H. Jeong; writing—review and editing, X.X.; project administration, J.H. Jeong;  
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