

Supporting Information

**Hydrophilic Packaging of Iron Oxide Nanoclusters for Highly
Sensitive Imaging**

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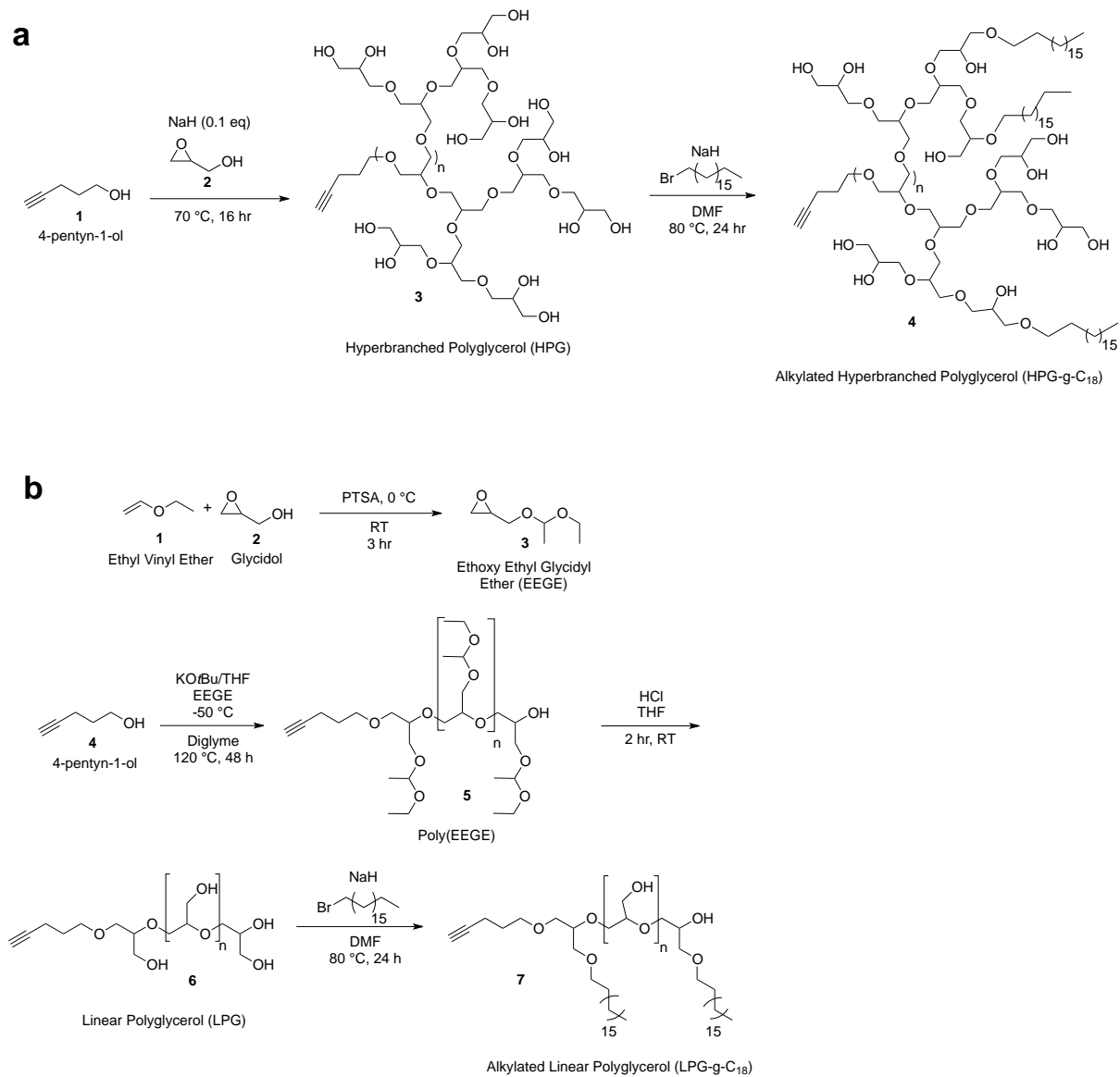


Figure S1. Synthesis of (a) alkylated HPG and (b) alkylated LPG.

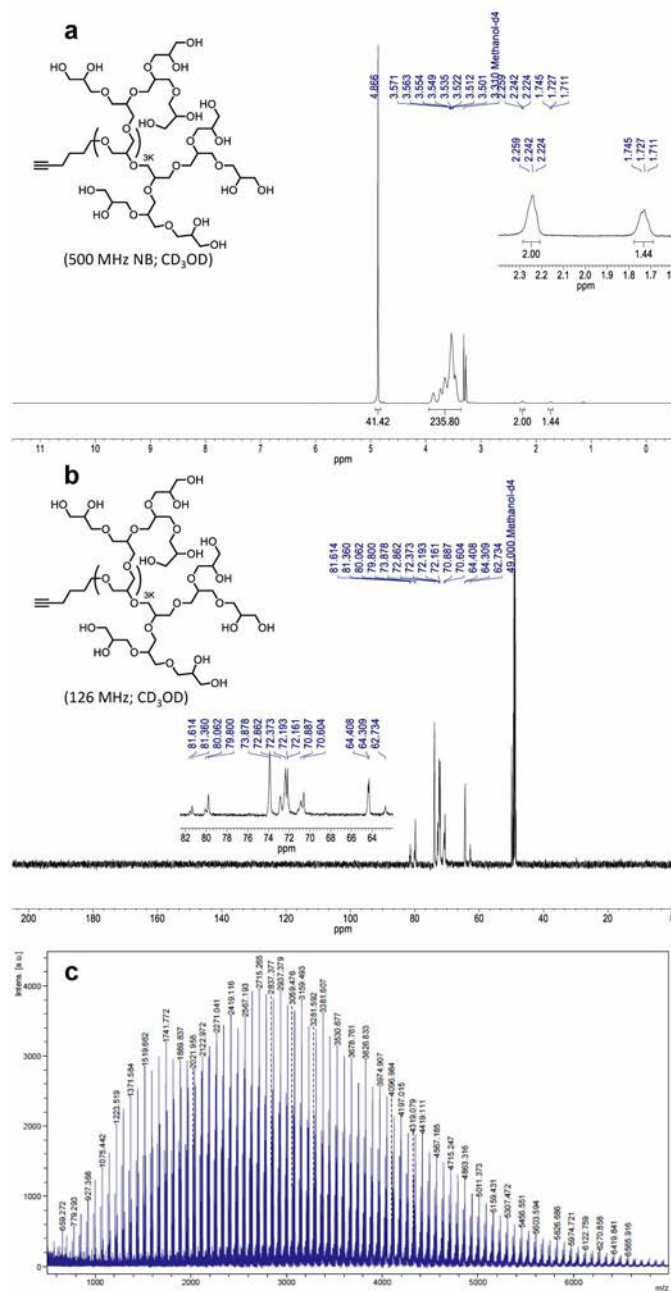


Figure S2. Characterization of HPG_{3k} by (a) ¹H NMR, (b) ¹³C NMR, and (c) MALDI-TOF. HPG_{3k} represents HPG with a molecular weight (MW) of 3,000 g/mol. ¹H NMR (400 MHz, methanol-*d*₄) δ 4.87 (s, 41H), 3.94 – 3.37 (m, 236H), 2.24 (t, *J* = 7.0 Hz, 2H), 1.78 – 1.69 (m, 1H). ¹³C NMR (126 MHz, methanol-*d*₄) δ 81.61, 81.36, 80.06, 79.80, 73.88, 72.86, 72.37, 72.19, 72.16, 70.89, 70.60, 64.41, 64.31, 62.73. MS (MALDI) *m/z* ~3000, peak separation 74 *m/z* units.

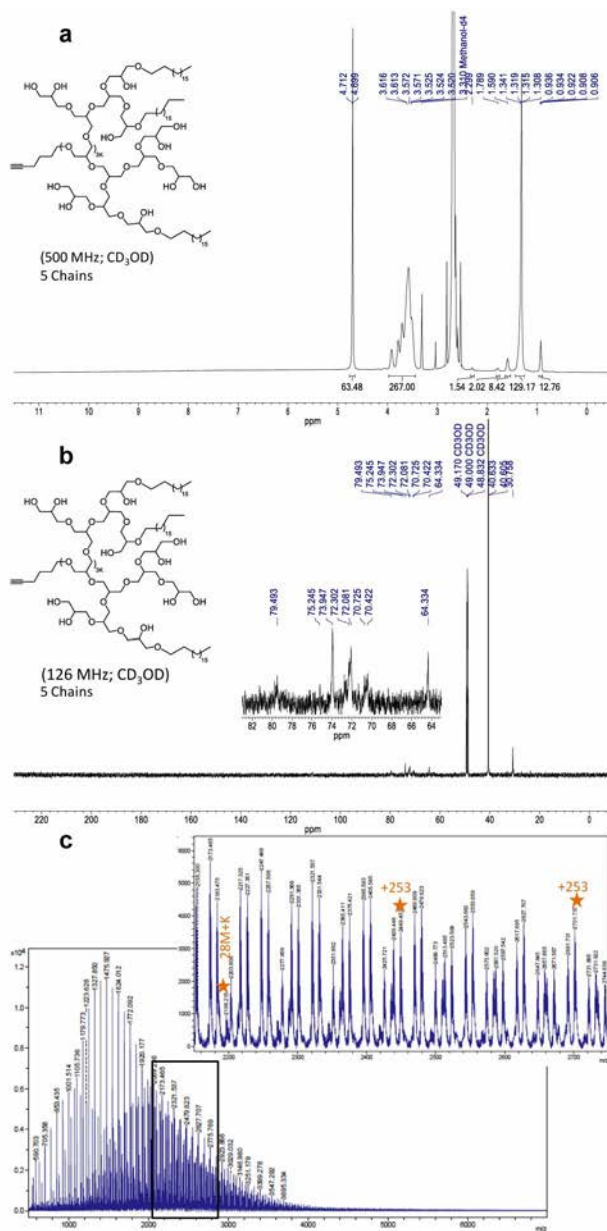


Figure S3. Characterization of HPG_{3k}-g-C₁₈(5) by **(a)** ¹H NMR, **(b)** ¹³C NMR, and **(c)** MALDI-TOF. Protons of the C₁₈ chain are from 1-2 ppm in the ¹H NMR spectrum. HPG_{3k}-g-C₁₈(5) represents HPG (MW ~3,000 g/mol) substituted with 5 C₁₈ chains. ¹H NMR (500 MHz, methanol-*d*₄) δ 4.71 (d, *J* = 6.6 Hz, 63H), 3.97 – 3.44 (m, 267H), 2.30 (s, 2H), 1.79 (s, 2H), 1.59 (s, 8H), 1.43 – 1.25 (m, 129H), 0.96 – 0.89 (m, 13H). ¹³C NMR (126 MHz, methanol-*d*₄) δ 79.49, 73.95, 72.30, 72.08, 70.73, 70.42, 64.33, 40.63, 40.61, 30.76. MS (MALDI) peak separation 253 m/z units.

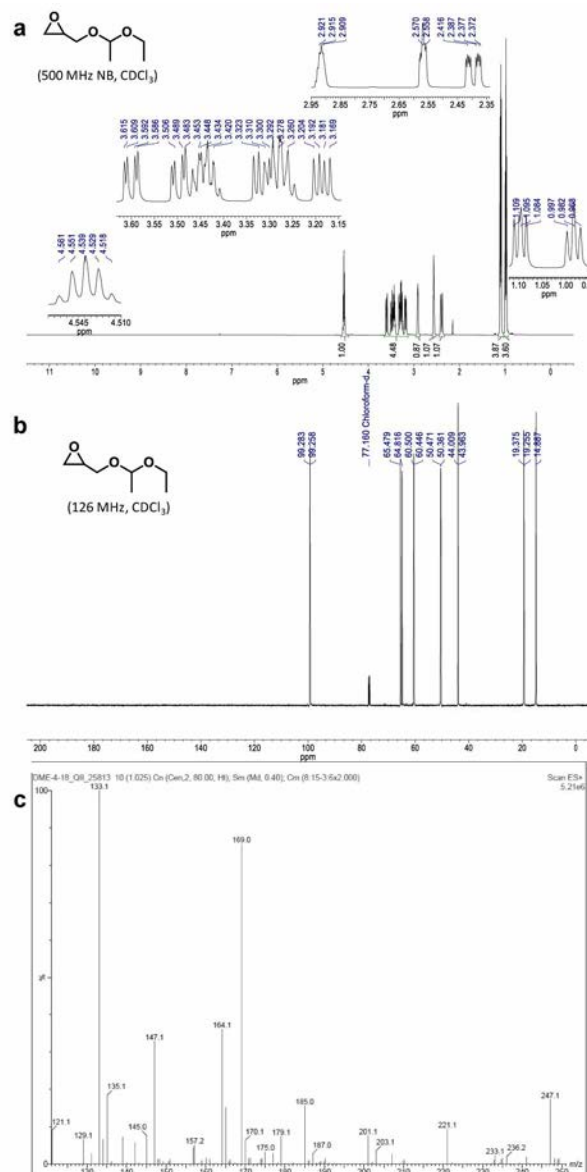


Figure S4. Characterization of EEGE, an LPG synthetic intermediate, by (a) ^1H NMR, (b) ^{13}C NMR, and (c) ESI mass spectrometry, calculated 146 m/z, experimental 147.1 (M+1), 169.0 m/z (M+Na). ^1H NMR (500 MHz, chloroform-*d*) δ 4.58 – 4.49 (m, 1H), 3.64 – 3.14 (m, 5H), 2.95 – 2.88 (m, 1H), 2.57 (ddt, $J = 5.9, 3.1, 1.5$ Hz, 1H), 2.43 – 2.35 (m, 1H), 1.10 (ddd, $J = 6.5, 5.4, 1.1$ Hz, 4H), 0.98 (tt, $J = 7.1, 1.3$ Hz, 4H). ^{13}C NMR (126 MHz, chloroform-*d*) δ 99.28, 99.26, 65.48, 64.82, 60.50, 60.45, 50.47, 50.36, 44.01, 43.96, 19.37, 19.25, 14.89.

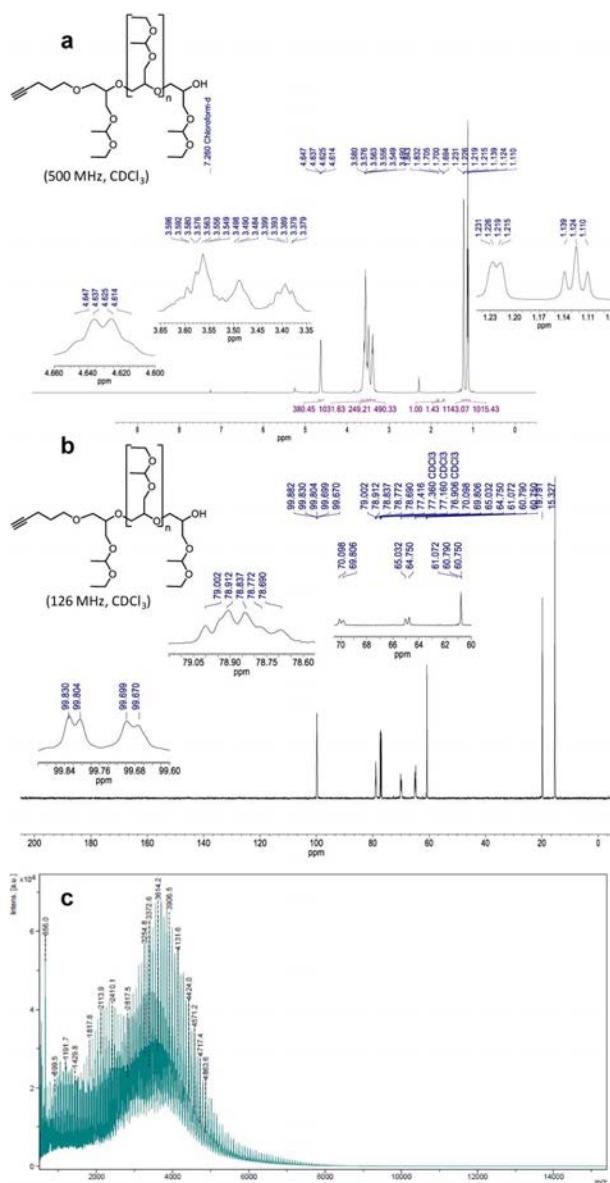


Figure S5. Characterization of poly(EEGE), an LPG synthetic intermediate, by (a) ^1H NMR, (b) ^{13}C NMR, and (c) MALDI-TOF. ^1H NMR (500 MHz, chloroform-*d*) δ 4.63 (q, $J = 5.5$ Hz, 380H), 3.65 – 3.51 (m, 1032H), 3.49 (q, $J = 4.2$ Hz, 249H), 3.44 – 3.34 (m, 490H), 1.84 (d, $J = 5.6$ Hz, 1H), 1.70 (t, $J = 2.6$ Hz, 1H), 1.26 – 1.18 (m, 1143H), 1.12 (t, $J = 7.1$ Hz, 1015H). ^{13}C NMR (126 MHz, CDCl_3) δ 99.88, 99.83, 99.80, 99.70, 99.67, 79.00, 78.91, 78.84, 78.77, 78.69, 77.42, 70.10, 69.81, 65.03, 64.75, 62.38, 61.07, 61.02, 60.79, 60.75, 19.79, 15.33. MS (MALDI) 3600 m/z, peak separation of 147 m/z units.

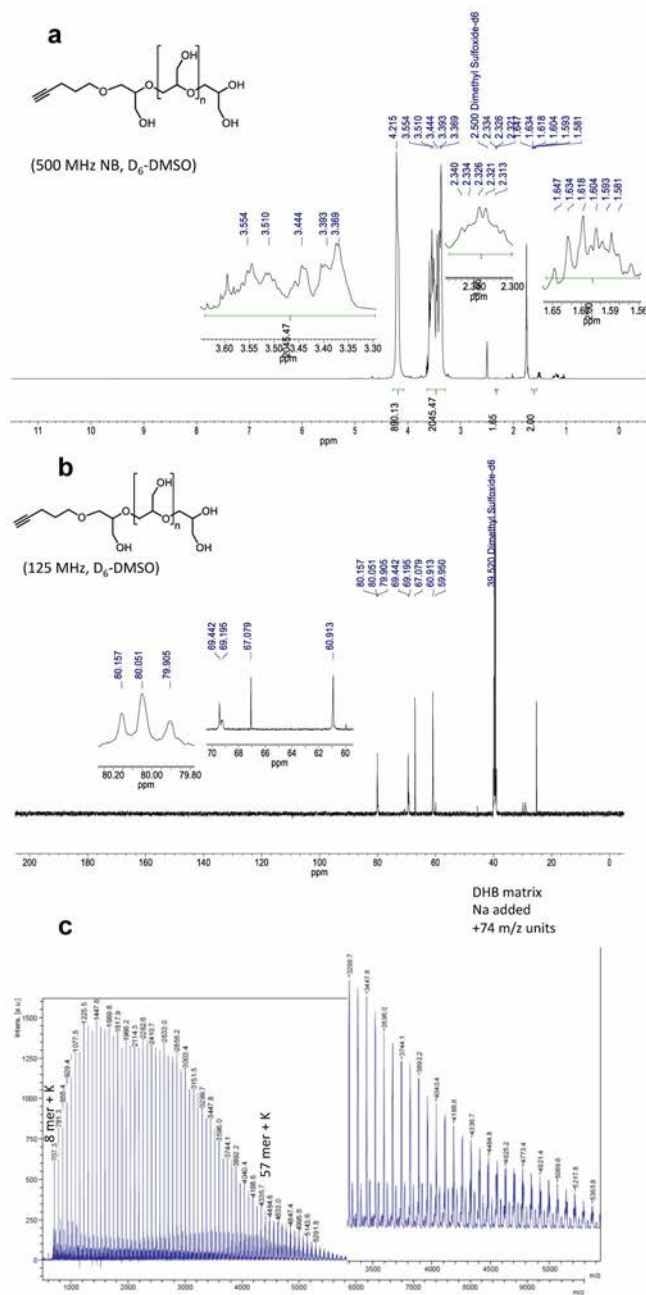


Figure S6. Characterization of LPG_{3k} by (a) 1H NMR, (b) ^{13}C NMR, and (c) MALDI-TOF. LPG_{3k} represents LPG with MW of $\sim 3,000$ g/mol. 1H NMR (500 MHz, $DMSO-d_6$) δ 4.29 – 4.11 (m, 1044H), 3.62 – 3.32 (m, 1897H), 2.32 (td, $J = 6.9, 3.0$ Hz, 1H), 1.65 – 1.58 (m, 2H). ^{13}C NMR (126 MHz, $DMSO-d_6$) δ 80.16, 80.05, 79.91, 69.45, 69.29, 67.08, 60.92, 59.95, 40.02, 39.94, 39.86, 39.78, 39.69, 39.60, 39.52, 39.35, 39.27, 39.19, 39.02. MS (MALDI) ~ 3000 m/z, separation of 74 m/z units.

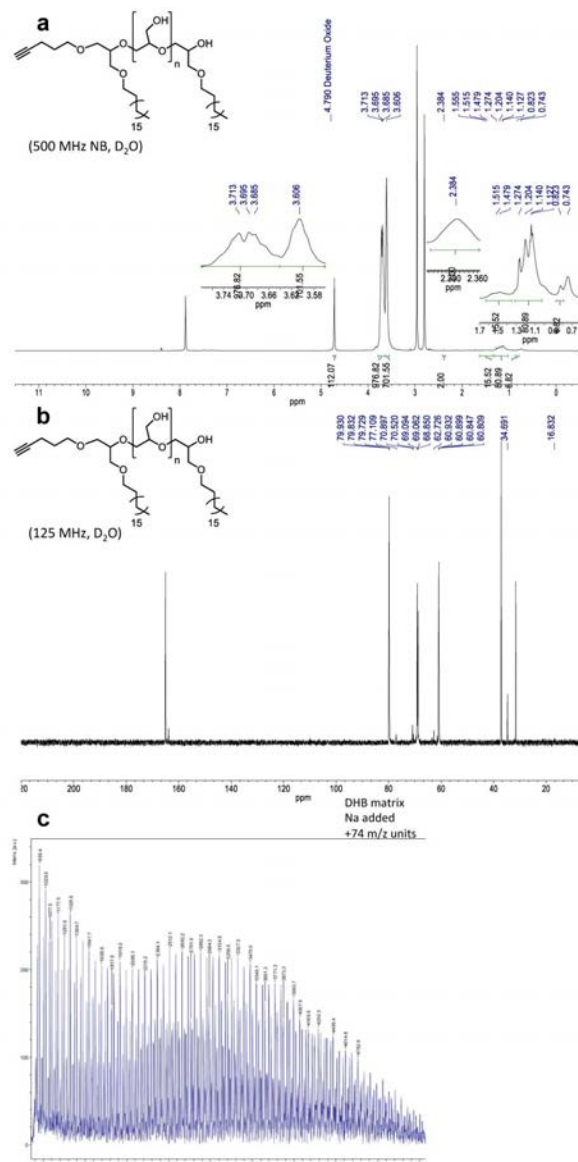


Figure S7. Characterization of LPG_{3k}-g-C₁₈(2) by (a) ¹H NMR, (b) ¹³C NMR, and (c) MALDI-TOF. Protons of the C₁₈ chain are from 1-2 ppm in the ¹H NMR spectrum. LPG_{3k}-g-C₁₈(2) represents LPG (MW ~3,000 g/mol) substituted with 2 C₁₈ chains. ¹H NMR (500 MHz, deuterium oxide) δ 3.70 (dd, *J* = 15.5, 9.7 Hz, 977H), 3.61 (s, 702H), 2.38 (s, 2H), 1.52 (m, 16H), 1.24 – 0.98 (m, 80H), 0.77 (s, 6H). ¹³C NMR (127 MHz, deuterium oxide) δ 79.93, 79.83, 79.73, 77.11, 70.90, 70.52, 69.09, 69.06, 68.85, 62.73, 60.93, 60.90, 60.85, 60.81, 37.08, 34.69, 31.57, 16.83. MS (MALDI) ~3500 m/z, separation of 74 m/z units.

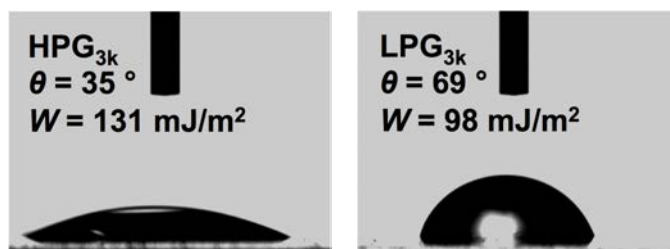


Figure S8. Contact angles and corresponding surface energies of water droplets on films of HPG_{3k} and LPG_{3k}.

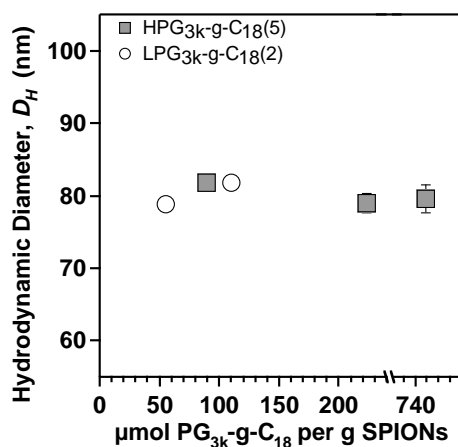


Figure S9. Z-average hydrodynamic diameter of SPION nanoclusters produced by various concentrations of HPG_{3k}-g-C₁₈(5) (■) and LPG_{3k}-g-C₁₈(2) (○). HPG_{3k}-g-C₁₈(5) and LPG_{3k}-g-C₁₈(2) represent HPG substituted with 5 C₁₈ chains and LPG substituted with 2 C₁₈ chains, respectively. Molecular weights of both HPG and LPG were 3,000 g/mol. Error bars, though partially obscured by data point markers, represent standard deviation of three measurements.

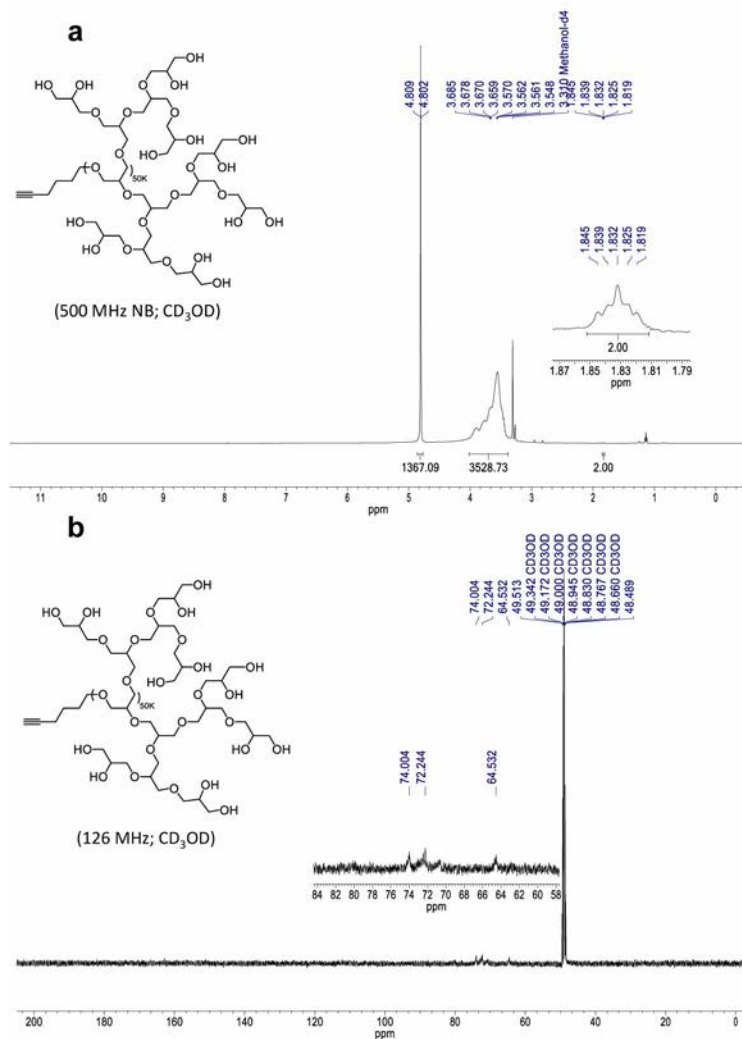


Figure S10. Characterization of HPG_{50k} by (a) ¹H NMR and (b) ¹³C NMR. HPG_{50k} represents HPG with MW of ~50,000 g/mol. ¹H NMR (500 MHz, methanol-*d*₄) δ 4.86 – 4.77 (m, 1367H), 4.02 – 3.39 (m, 3529H), 1.85 – 1.81 (m, 2H). ¹³C NMR (126 MHz, methanol-*d*₄) δ 74.00, 72.24, 64.53.

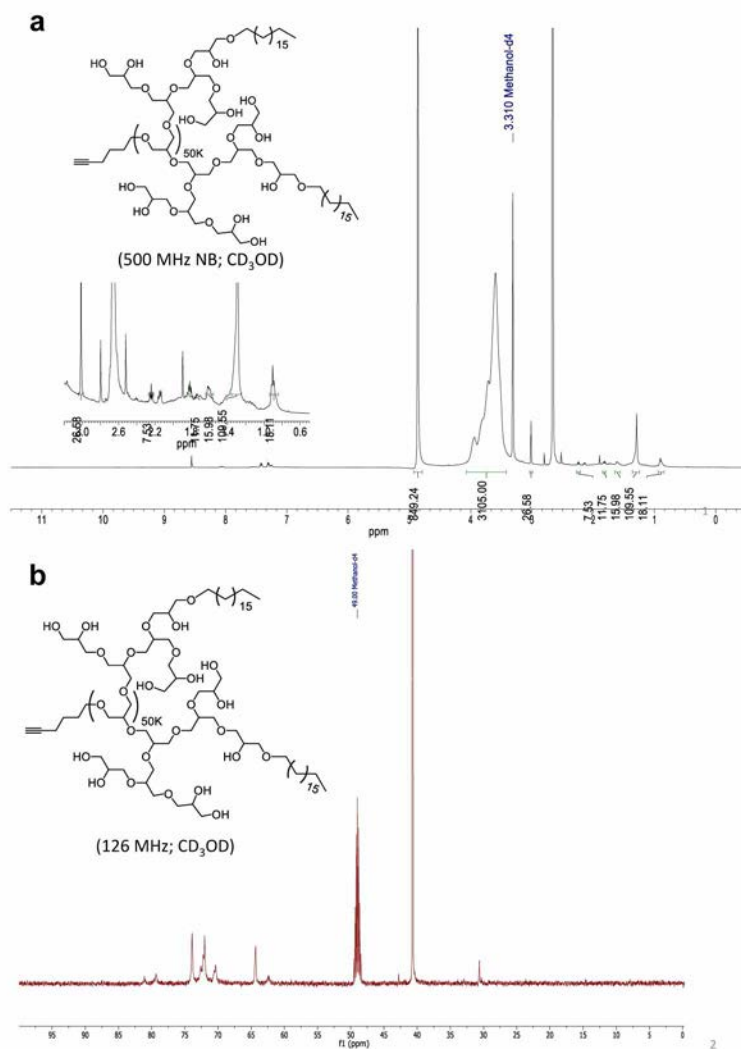


Figure S11. Characterization of HPG_{50k}-g-C₁₈(2) by **(a)** ¹H NMR and **(b)** ¹³C NMR. Protons of the C₁₈ chain are from 1-2 ppm in the ¹H NMR spectrum. HPG_{50k}-g-C₁₈(2) represents HPG (MW ~50,000 g/mol) substituted with 2 C₁₈ chains. **¹H NMR** (500 MHz, methanol-*d*₄) δ 4.88 (s, 849H), 4.10 – 3.37 (m, 3105H), 3.03 (d, *J* = 3.0 Hz, 26H), 2.26 (t, *J* = 7.3 Hz, 8H), 1.88 – 1.78 (m, 12H), 1.67 – 1.56 (m, 16H), 1.32 (d, *J* = 9.5 Hz, 110H), 0.91 (dt, *J* = 10.0, 6.0 Hz, 18H). **¹³C NMR** (126 MHz, methanol-*d*₄) δ 78.69, 72.84, 71.25, 71.04, 69.49, 63.36, 47.91, 47.84, 47.74, 47.56, 47.45, 47.39, 41.44, 39.28, 29.59.

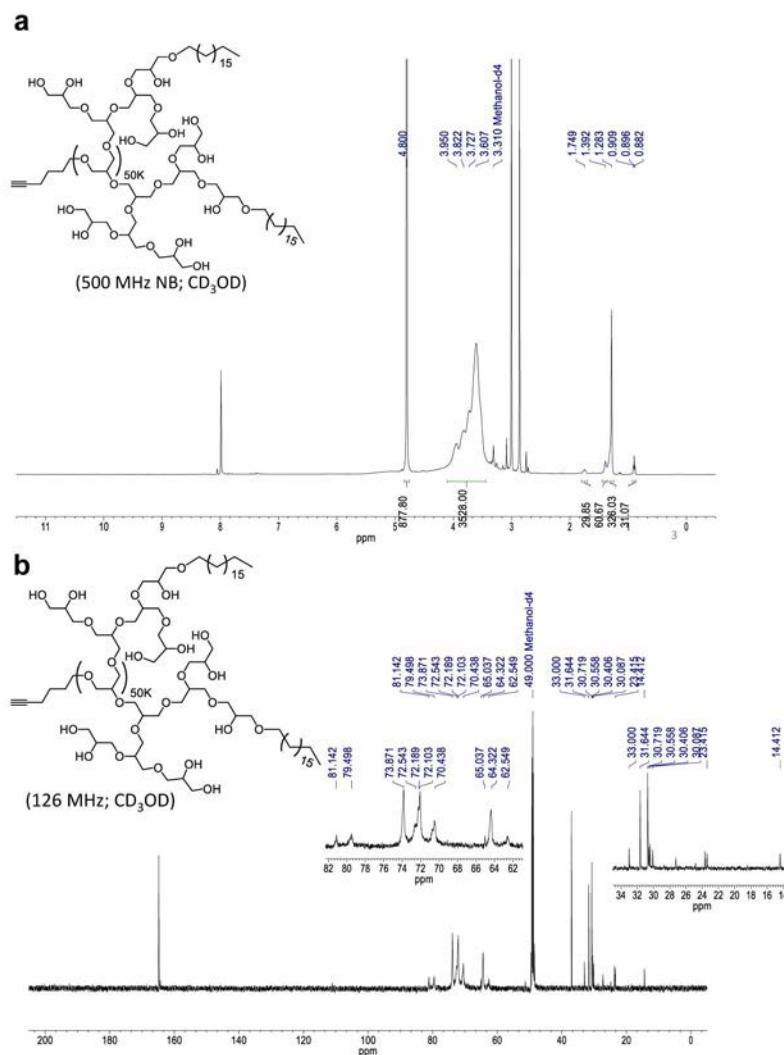
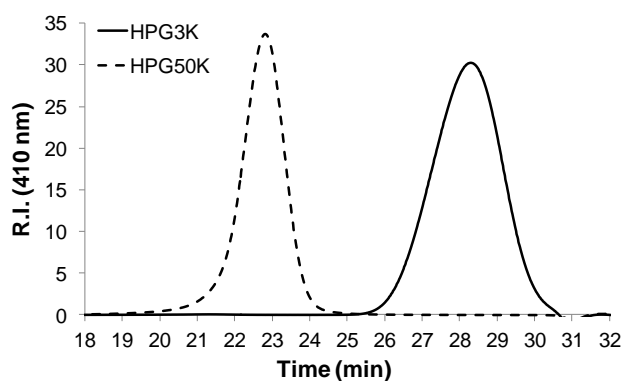


Figure S12. Characterization of HPG_{50k}-g-C₁₈(10) by (a) ¹H NMR and (b) ¹³C NMR.

Protons of the C₁₈ chain are from 1-2 ppm in the ¹H NMR spectrum. HPG_{50k}-g-C₁₈(10) represents HPG (MW ~50,000 g/mol) substituted with 10 C₁₈ chains. **¹H NMR** (500 MHz, methanol-*d*₄) δ 4.80 (s, 878H), 3.95 – 3.61 (m, 3528H), 1.75 (dd, *J* = 10.9, 5.9 Hz, 30H), 1.40 (d, *J* = 5.7 Hz, 60H), 1.28 (s, 326H), 0.91 (t, *J* = 6.7 Hz, 31H). **¹³C NMR** (126 MHz, methanol-*d*₄) δ 81.14, 79.50, 73.87, 72.54, 72.19, 72.10, 70.44, 65.04, 64.32, 62.55, 30.00, 31.64, 30.72, 30.56, 30.41, 30.09, 23.42, 14.41.



Compounds Synthesized	SEC DMF MW ^a	PDI
HPG _{3k}	28.28 min (3019 MW)	1.20
HPG _{50k}	22.80 min (46,025 MW)	1.46

^a PEG standards

Figure S13. Characterization of molecular weights of HPG_{3k} and HPG_{50k} by GPC. The molecular weight of HPG_{3k} is in agreement with that determined by MALDI-TOF.

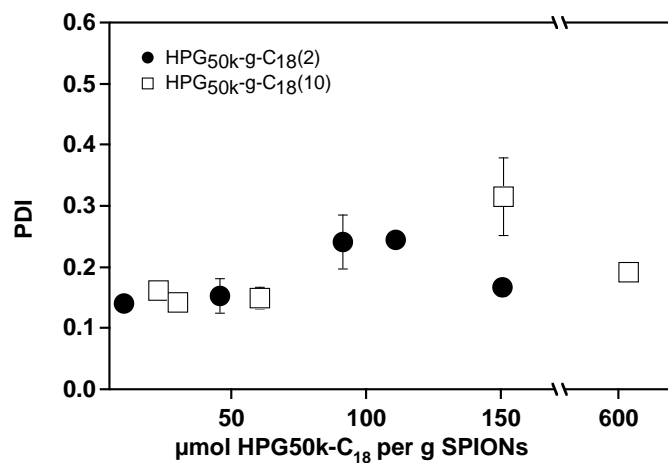


Figure S14. Polydispersity index (PDI) for the size-tunable clusters shown in Figure 2 of the text. Data are the average of three replicate measurements, with error bars representing standard deviation partially obscured by data point markers.

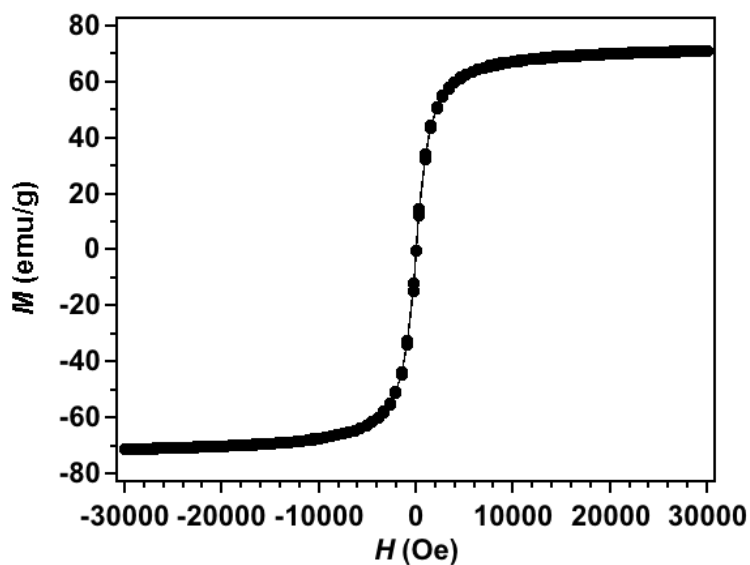


Figure S15. Magnetization curve of oleic acid-capped SPIONs at 300 K. A saturation magnetization, M_s , of 71 emu/g, determined from the plateau of the magnetization curve, was used to estimate theoretical maximum relaxivity in Eq. (3) of the text.

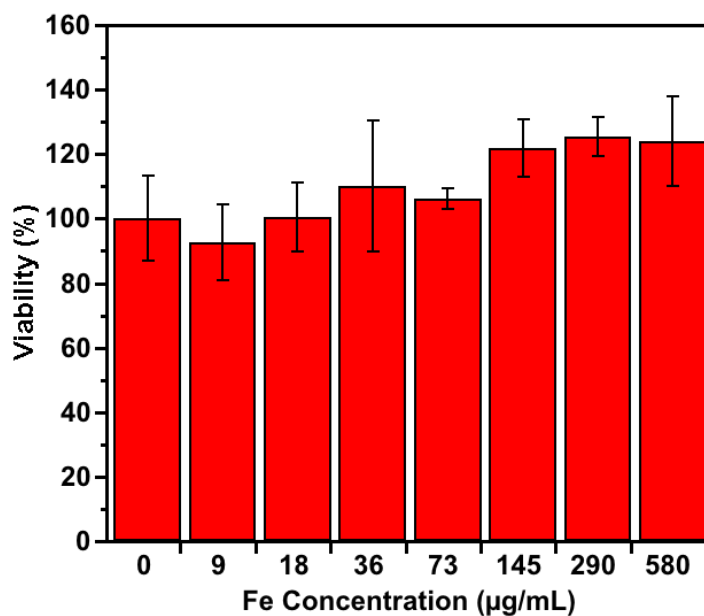


Figure S16. MTT assay to assess effects of HPG-SPION nanoclusters on metabolic activity of C166 endothelial cells. Percent viability was determined by normalization of absorbance values after blank subtraction to that of a control group of cells incubated without HPG-SPIONs. The concentration of Fe in the cell culture media was varied by altering the mass of SPION clusters. Values are the average of 3 replicates and error bars represent standard deviation.

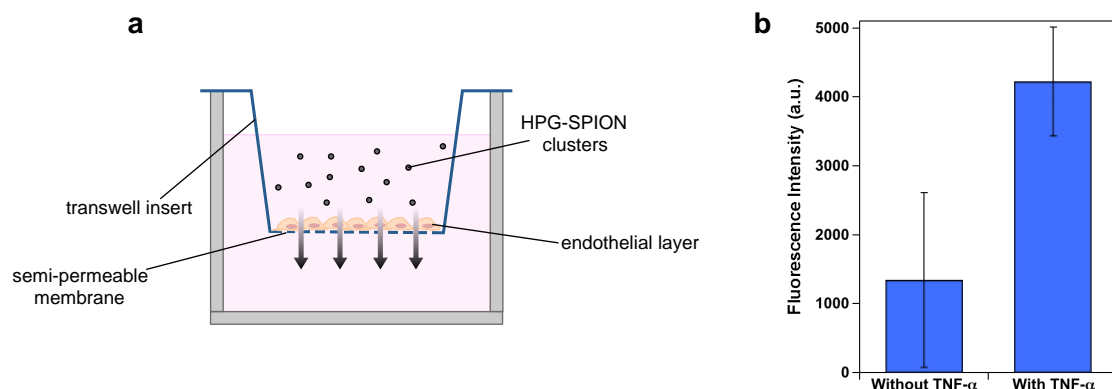


Figure S17. Endothelial transwell migration assay. **(a)** Schematic of the transwell insert. **(b)** Fluorescence intensity of the receiving well after incubation of FITC-labeled HPG-SPIONs and C166 cells with and without exposure to inflammation-inducing TNF- α . The diffusion of HPG-SPION nanoclusters was more than 3-fold higher through the endothelial layer exposed to TNF- α . Error bars represent standard deviation of three replicates.

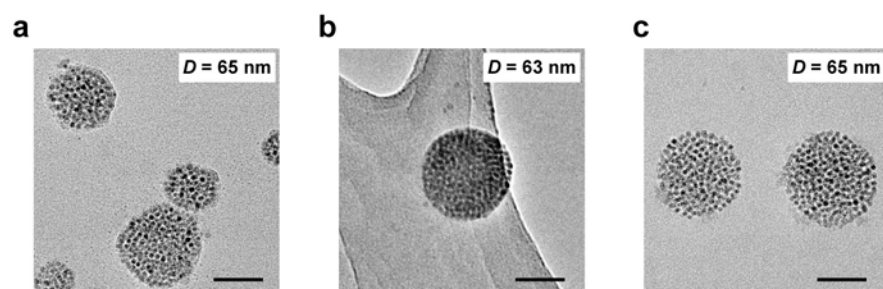


Figure S18. TEM micrographs of HPG-SPIONs before and after incubation with 50% human serum in PBS. The HPG-SPIONs with a hydrodynamic diameter of 122 nm and relaxivity of $719 \text{ mM}^{-1} \text{ s}^{-1}$ were analyzed, and average core diameter is indicated on each figure. **(a)** TEM and **(b)** cryo-TEM images of the SPION clusters before serum incubation. **(c)** After incubation with 50% human serum in PBS for 2 h at 37 °C, the morphology and size remained unchanged, according to TEM micrographs. Scale bars represent 50 nm.

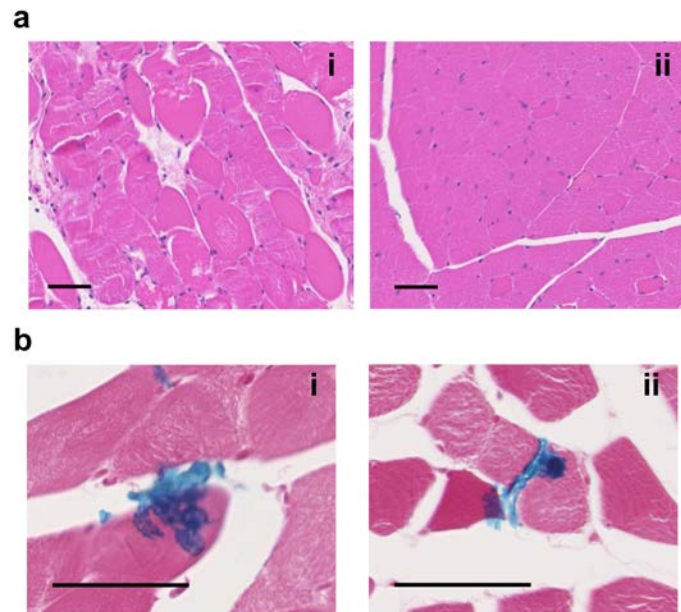


Figure S19. Histological analysis of the hindlimb ischemia model. **(a)** H&E staining of (i) the thigh of the ischemic left hindlimb, indicating tissue damage as a result of ischemia and (ii) the thigh of the uninjured, right hindlimb. **(b)** Prussian blue staining of the left, ischemic hindlimb sections containing (i) HPG-SPION nanoclusters and (ii) unclustered SPIONS, indicating the presence of contrast agent in both cases. In all figures, scale bars represent 50 μm .