

Figure S1. Proton nuclear magnetic resonance

Methods: Proton nuclear magnetic resonance (^1H NMR) was performed whereby 10 mg/mL Hep^N MAm and PEGDA samples were each dissolved in deuterated H_2O (Sigma), run on a Bruker Avance III spectrometer at 400 Hz, and analyzed using iNMR software.³⁵ Percent modification was determined by dividing the integral of the methacrylamide peak by the heparin peak for Hep^N MAm and the acrylate peaks by the PEG peak for PEGDA.

Results: Using ^1H NMR analysis, PEGDA was determined to be ~55% functionalized (Figure S1A) and Hep^N methacrylamide was determined to be 22-28% functionalized (Figure S1B).

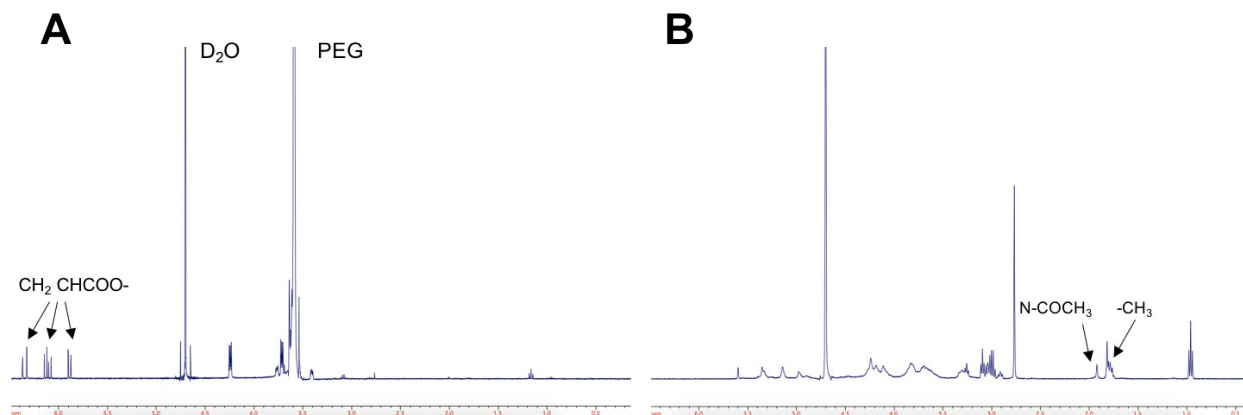


Figure S1. ^1H NMR of poly (ethylene glycol) diacrylate and N-desulfated heparin methacrylamide. (A) Characteristic peaks for acrylate groups were present in the PEGDA spectra and (B) characteristic peaks for methacrylamide peaks were present in the N-desulfated heparin methacrylamide.

Figure S2. Microparticle size and morphology before and after TSG-6 loading

Methods: TSG-6 was loaded onto microparticles (MPs) using the same method for all other *in vitro* and *in vivo* studies. Briefly, 0.6 mg 10 wt% Hep^N MPs were incubated with 1.0 µg TSG-6 for 2 hours at 4°C. Phase microscopy and ImageJ software were used to image and quantify the diameter of microparticles before and after the loading protocol.

Results: MPs appear to have similar morphology (Figure S2A-B) and exhibit no difference in size distribution (Figure S2C) before and after loading.

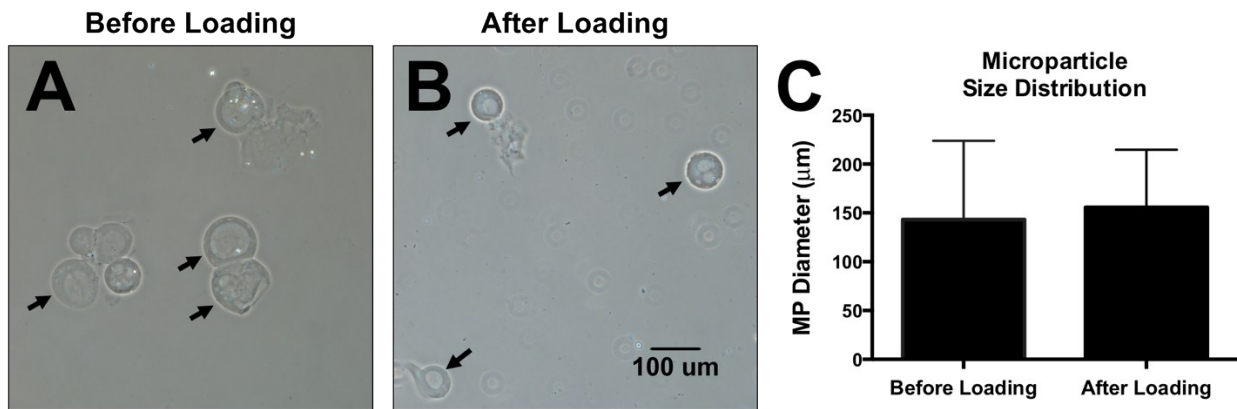


Figure S2. Microparticle size and morphology before and after TSG-6 loading. (A-B) 10 wt% Hep^N MPs appeared similar in morphology before and after TSG-6 loading, and (C) were not significantly different in average size distribution. Before loading and after loading group were not significantly different; $p \leq 0.05$; two-tailed t-test; $n > 30$ microparticles per group; data shown as mean \pm SD.