Supplementary Information for

Grafting of Cationic Molecules to Hyaluronic Acid Improves Adsorption and Cartilage Lubrication

Gavin Gonzales¹, Jiaul Hoque², Colin Kaeo¹, Stefan Zauscher³, and Shyni Varghese^{1, 2, 3*} ¹Department of Biomedical Engineering, Duke University, Durham, NC 27710, USA

²Department of Orthopaedic Surgery, Duke University School of Medicine, Durham, NC 27710, USA

³Department of Mechanical Engineering and Materials Science, Duke University, Durham, NC 27710, USA

*To whom correspondence should be addressed:

E-mail: <u>shyni.varghese@duke.edu</u>, Tel: +1-919-660-5273

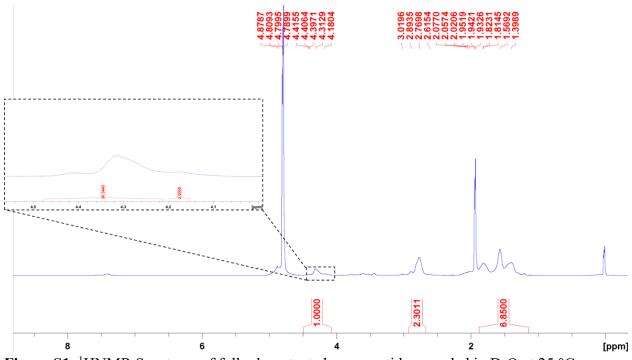


Figure S1. ¹HNMR Spectrum of fully deprotected core peptide recorded in D_2O at 25 °C.

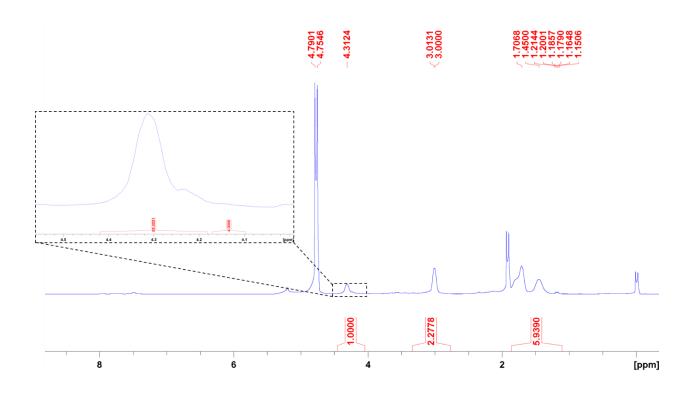


Figure S2. ¹HNMR Spectrum of fully deprotected G0 peptide recorded in D₂O at 25 °C.

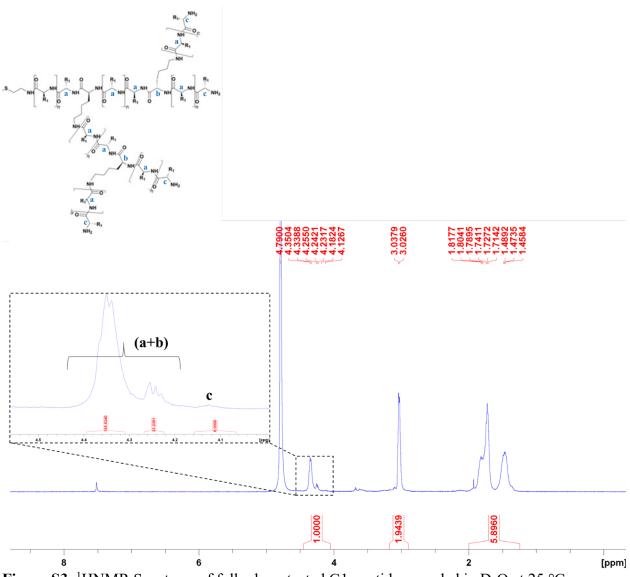


Figure S3. ¹HNMR Spectrum of fully deprotected G1 peptide recorded in D₂O at 25 °C.

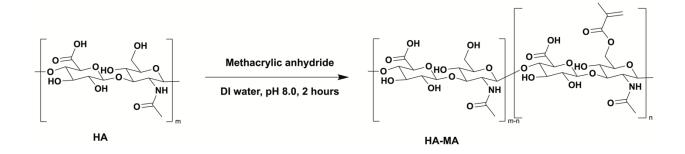


Figure S4. Synthesis of HA-MA from HA. Reaction scheme showing the incorporation of methacrylate groups in the HA by reacting HA with methacrylic anhydride.

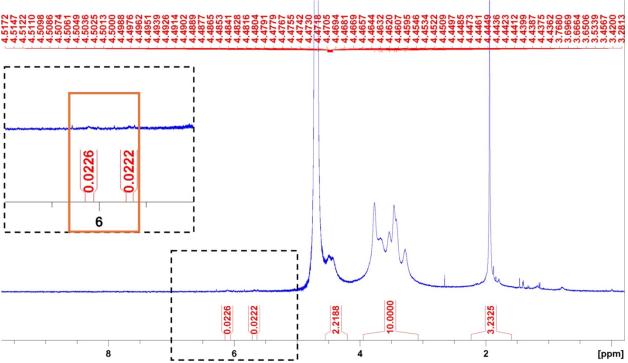


Figure S5. ¹HNMR Spectrum of HA-MA recorded in D₂O at 25 °C. The box shows an expansion of the region from 4.25-6.5ppm with the methacrylate protons of the HA-MA marked by the orange box.

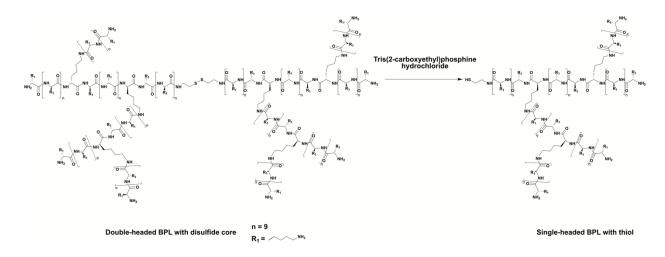


Figure S6. Reduction of disulfide bond in the double-headed BPL molecule.

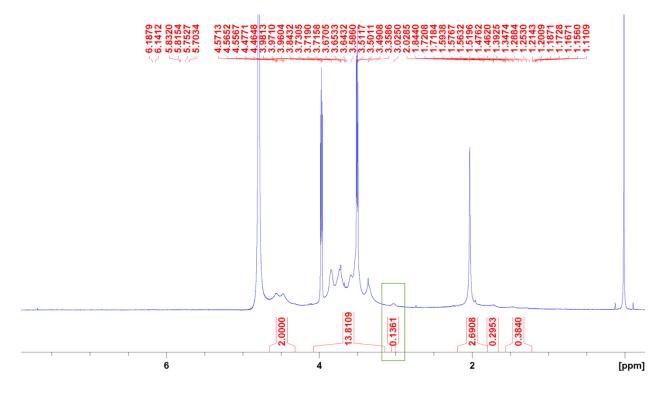


Figure S7. ¹HNMR Spectrum of HA-2BPL recorded in D₂O at 25 °C. Peak at 3.025 ppm (marked with a green colored box) indicates the appearance of methylene protons adjacent to NH₂ group (-CH₂CH₂CH₂CH₂CH₂NH₂) of lysine unit from fully deprotected BPL.

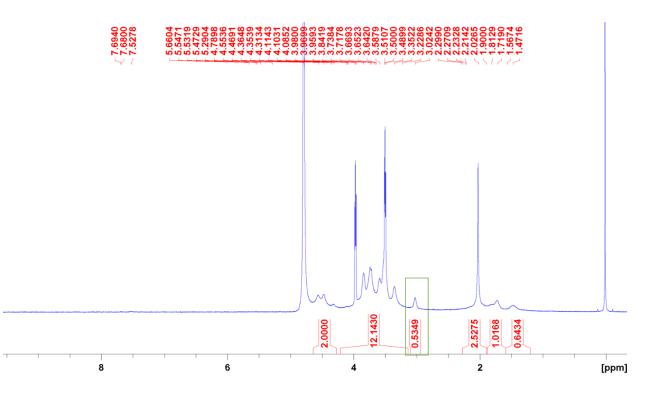


Figure S8. ¹HNMR Spectrum of HA-8BPL recorded in D₂O at 25 °C. Peak at 3.024 ppm (marked with a green colored box) indicates the appearance of methylene protons adjacent to NH₂ group (-CH₂CH₂CH₂CH₂CH₂NH₂) of lysine unit from fully deprotected BPL.

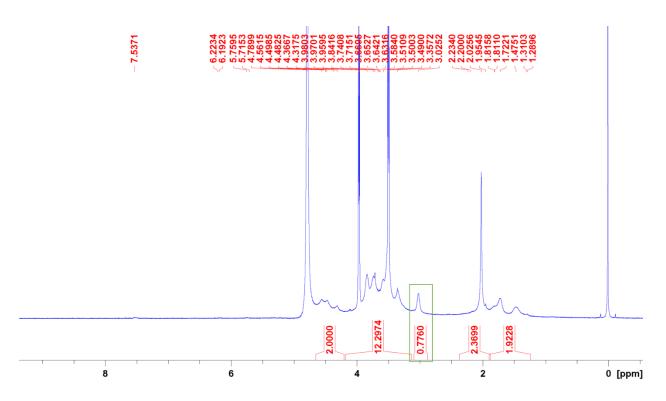


Figure S9. ¹HNMR Spectrum of HA-12BPL recorded in D₂O at 25 °C. Peak at 3.025 ppm (marked with a green colored box) indicates the appearance of methylene protons adjacent to NH_2 group (-CH₂CH₂CH₂CH₂CH₂NH₂) of lysine unit from fully deprotected BPL.

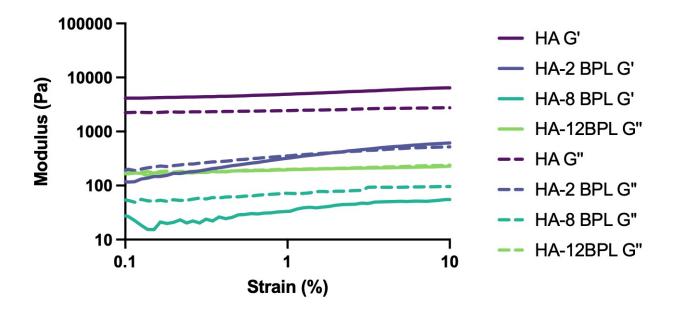


Figure S10. Strain sweep of HA and HA-BPL molecules.

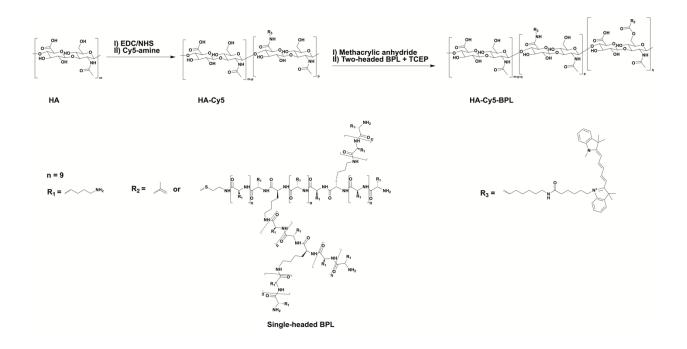


Figure S11. Synthesis of Cy5 containing HA-BPL molecules.

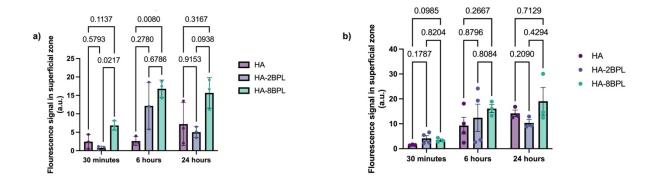


Figure S12. Adsorption of HA-BPL molecules to cartilage as a function of time. a) Penetration of HA BPL molecules in healthy cartilage explants. b) Penetration of HA BPL molecules in OA-mimetic cartilage explants.

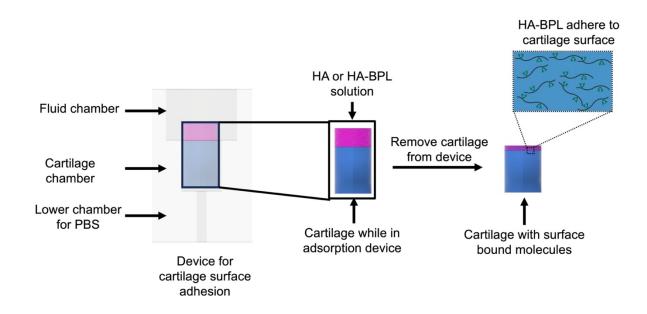


Figure S13. Schematic of device used to assess adsorption of HA and HA-BPL molecules to cartilage.

Table S1.	Molecular	characteristics	of different	generations
-----------	-----------	-----------------	--------------	-------------

	Number of terminal lysine residues	Total number of branches/arms	^a DP _{branch}	^b DP _{total}
Core	2	2	10	20
G0	4	6	10	60
Gl	8	14	10	140

^{a,b} Theoretical estimation of average number of lysine repeating units per branch (DP_{branch}) and the total number average degree of polymerization (DP_{total})

^a Yields determined gravimetrically after precipitation, filtration, and freeze-drying of the polypeptides.

¹ HNMR Integrals							
	$I(H_a) + I(H_b)$	$I(H_c)$	$[I(H_a)+I(H_b)+I(H_c)]/I(H_c)$	DPbranch	DP _{total}		
Core	20.3	2	11.15	~11.1	22		
G0	85.0	4	22.25	~11.1	89		
G1	178.76	8	23.34	~11.6	187		

Table S2. Results of the ¹HNMR characterization of BPL molecules

 $I(H_a)$ = Integral for methine protons of the terminal lysine units from the core peptide $I(H_b)$ = Integral for methine protons of the terminal lysine units from the G0 peptide $I(H_c)$ = Integral for methine protons of the terminal lysine units from the G1 peptide