

Electronic Supplementary Information

Induction of a higher-ordered architecture in glatiramer acetate improves its biological efficiency in an animal model of multiple sclerosis

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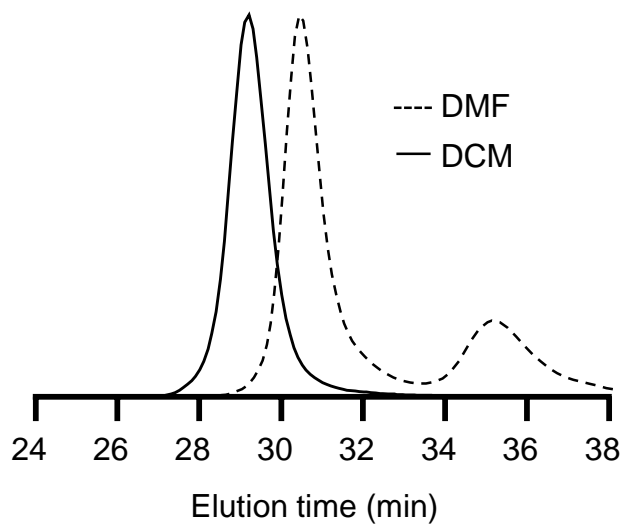


Fig. S1. Normalized GPC-dRI traces of the resulting polypeptides from PAMAM-initiated polymerization of BLG-NCA in DMF ($M_n = 65.0$ kDa, $\mathcal{D} = 3.22$) and DCM ($M_n = 345$ kDa, $\mathcal{D} = 1.10$). $[M]_0/[I]_0 = 50$. $[M]_0 = 400$ and 100 mM for the polymerization in DMF and DCM, respectively.

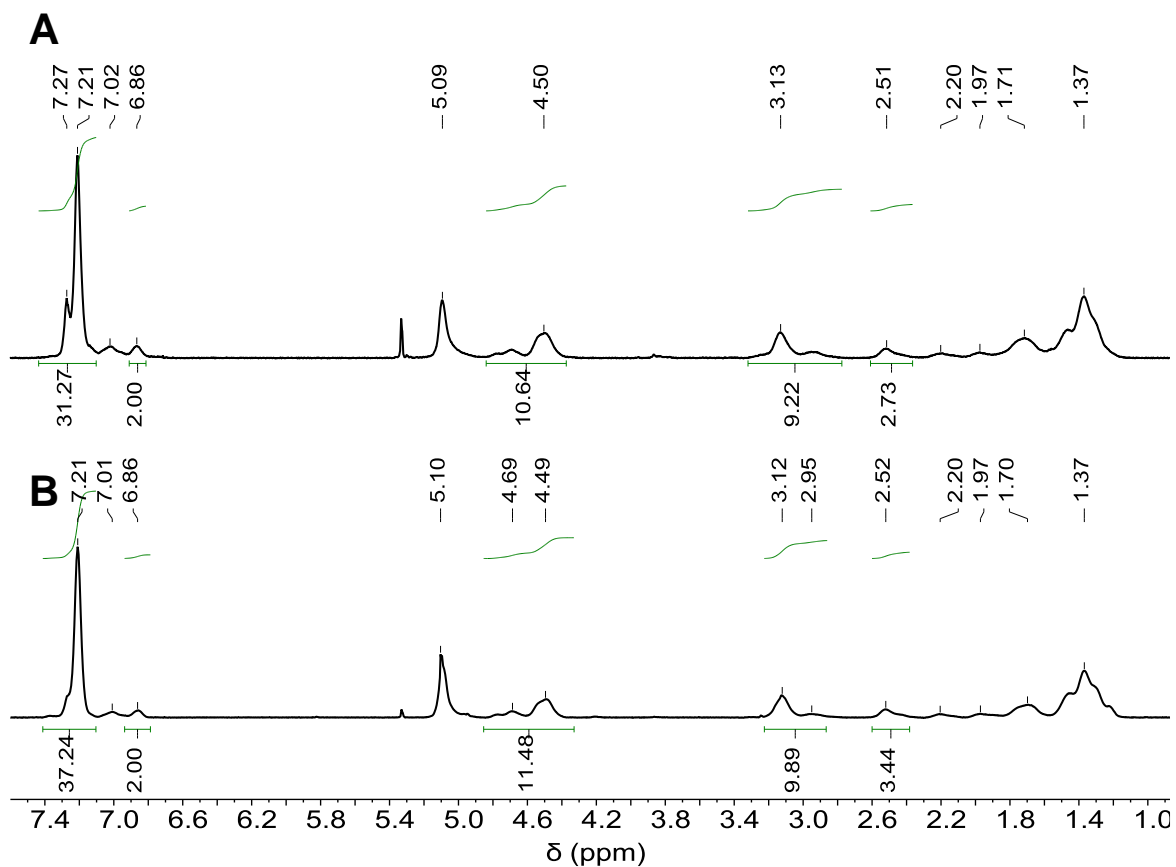


Fig. S2. ¹H NMR spectra (500 MHz) of the polypeptide precursors for sGA (A) and GA (B) in TFA-*d*. The fractions of amino acid residues were calculated based on the integration of peaks at 6.86 (Ar-H of BLT residues), 4.85–4.30 (α-H of all residues), 3.25–2.80 (ε-H of ZLL residues and β-H of BLT residues), and 2.51 (γ-H of BLG residues) ppm.

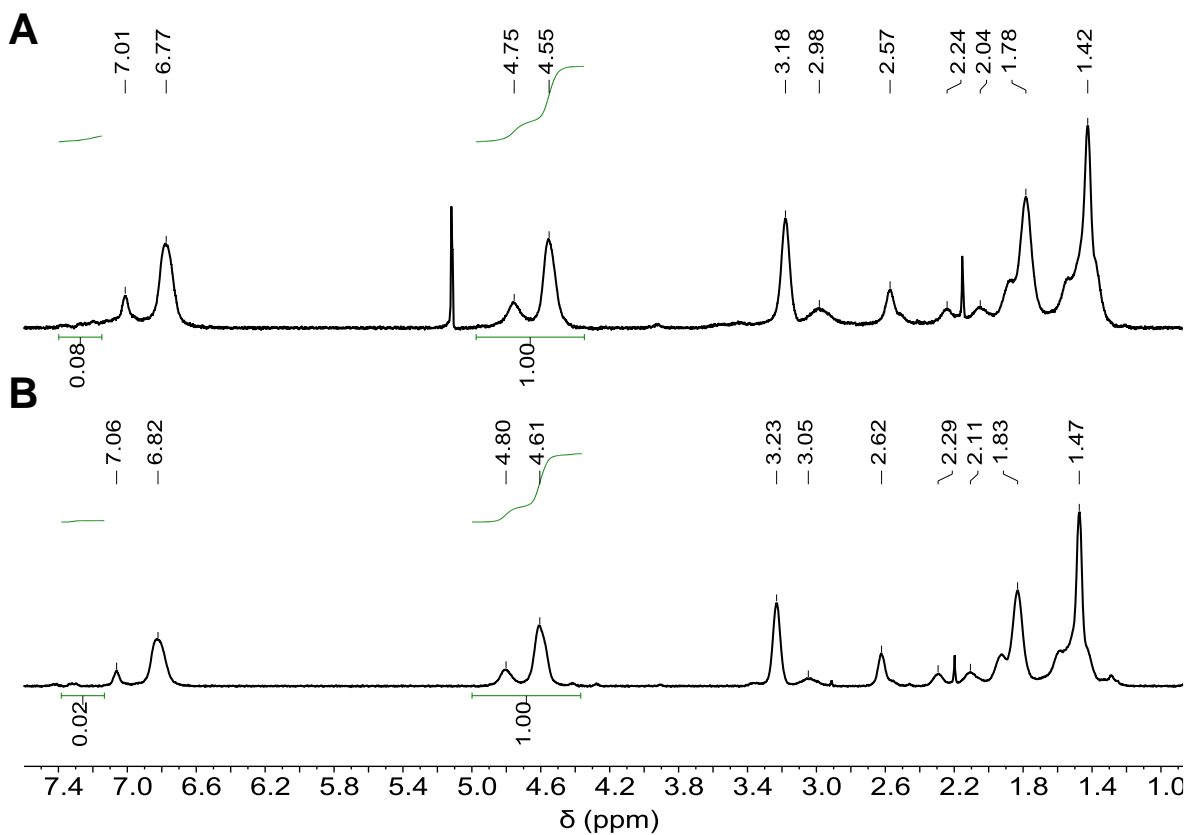


Fig. S3. ^1H NMR spectra (500 MHz) of sGA (A) and GA (B) in $\text{TFA-}d$. The deprotection efficiency was determined by comparing the integration of peaks at 7.4–7.1 ppm (Ar-H of protecting groups) before and after deprotection.

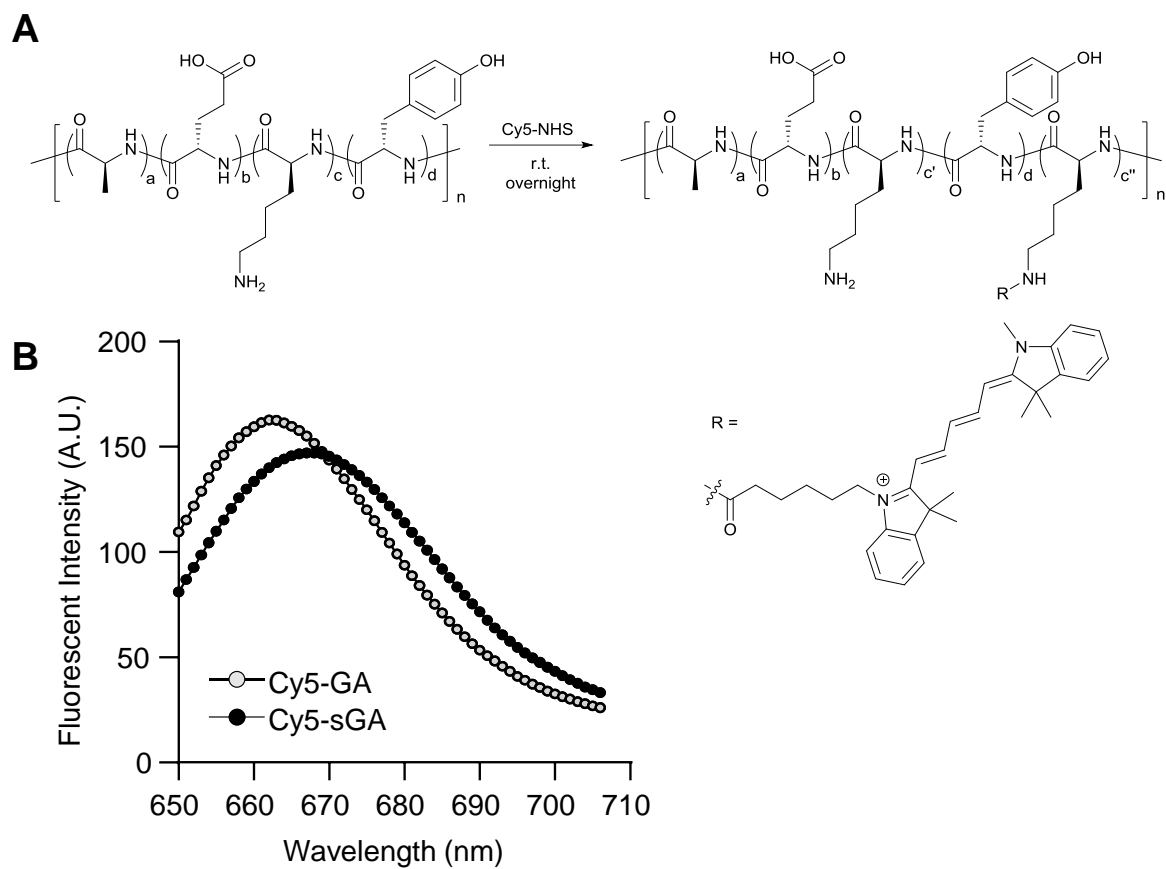


Fig. S4. Fluorescent labelling of sGA and GA with Cy5. (A) Synthetic route to Cy5-labelled sGA and GA. (B) Representative fluorescent spectra of Cy5-labelled sGA and GA at 20 $\mu\text{g/mL}$ in a PBS buffer.