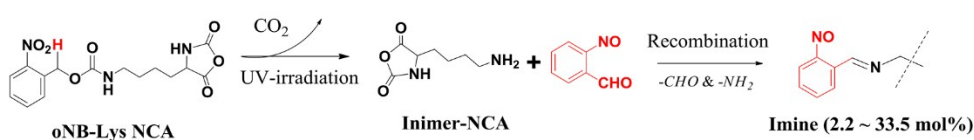


Hyperbranched Polypeptides Synthesized from Phototriggered ROP of a Photocaged N_ϵ -[1-(2-nitrophenyl)ethoxycarbonyl]-L-Lysine- N -Carboxyanhydride: Microstructures and Effects of Irradiation Intensity and Nitrogen Flow Rate

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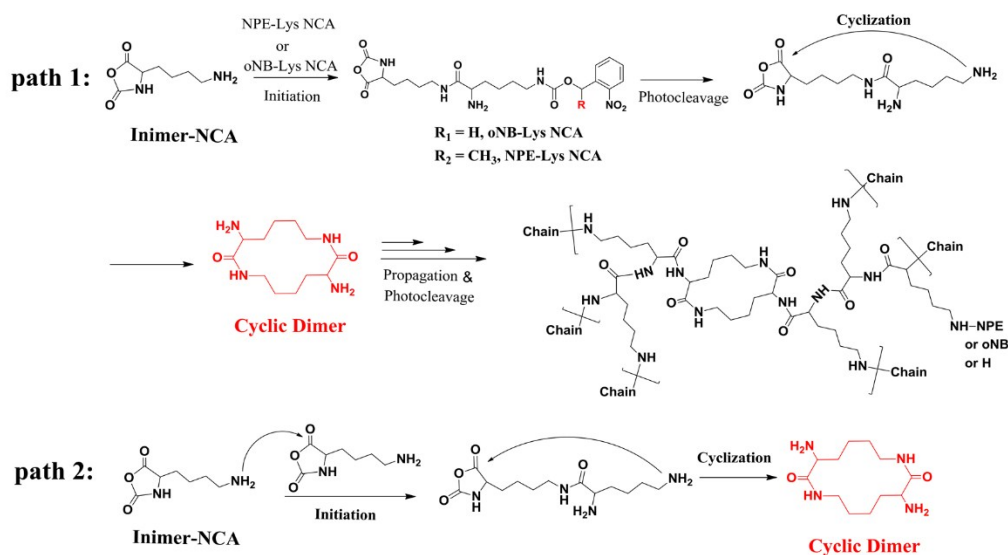
A Our previous work (*ACS Macro Lett.* 2017, 6, 292-297.)



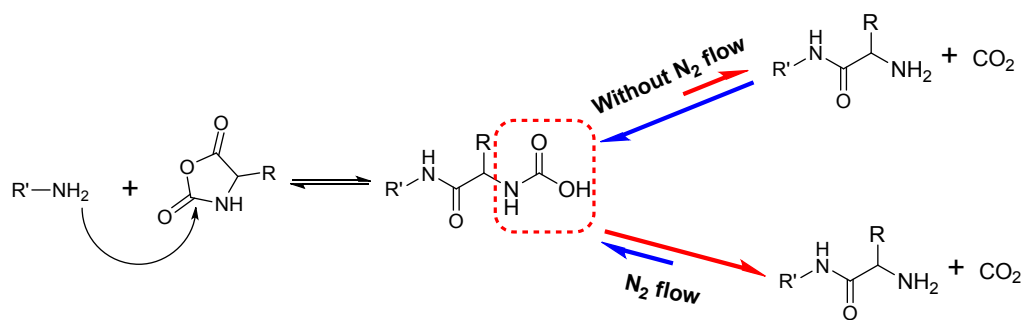
In this work



B



Scheme S1. The imination side reaction between photo-byproduct and amino groups during UV-triggered ROP (A) and the postulated pathways for forming the cyclic dimer initiating species during UV-triggered ROP (B).



Scheme S2. The promoted normal amine mechanism of NCA ROP under N_2 flow.

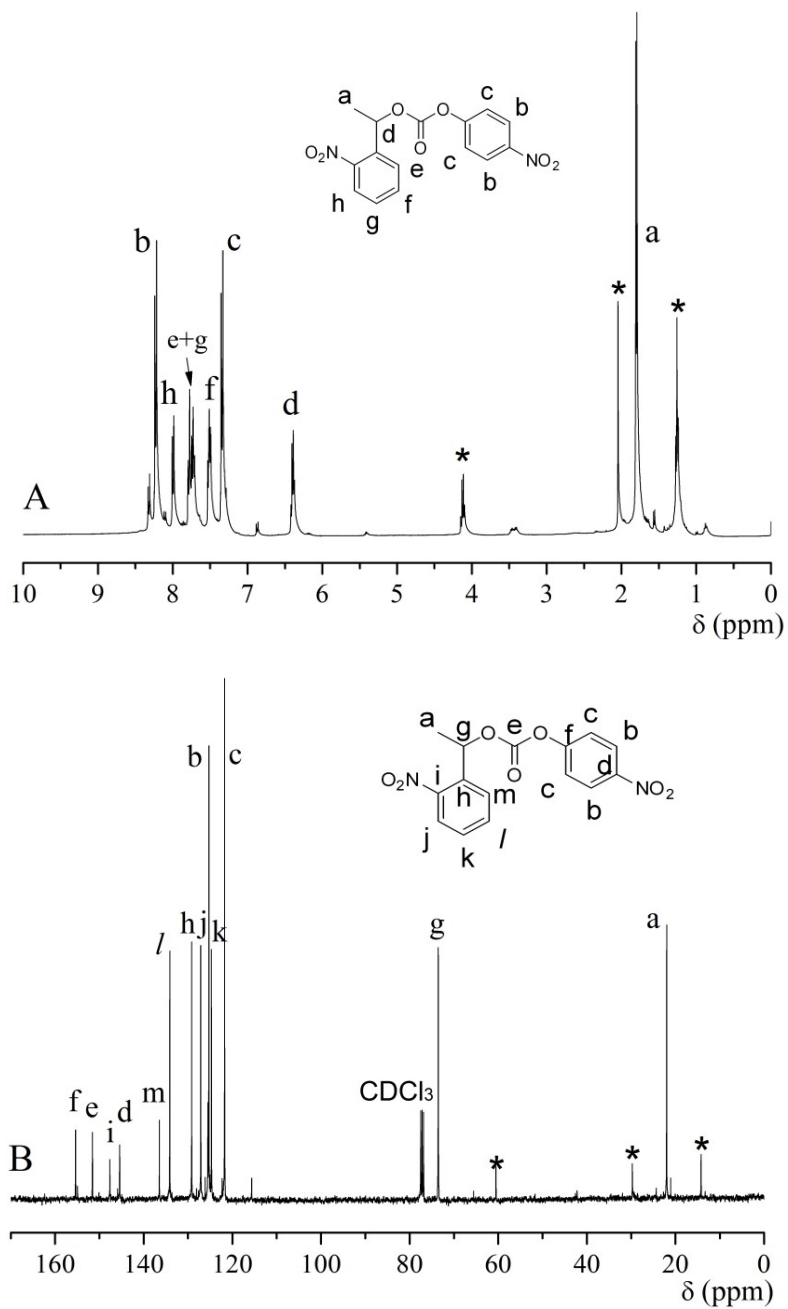


Figure S1. 1H NMR (A) and ^{13}C NMR spectra of NPE-NPC ("*" denotes the EtOAc solvent peaks).

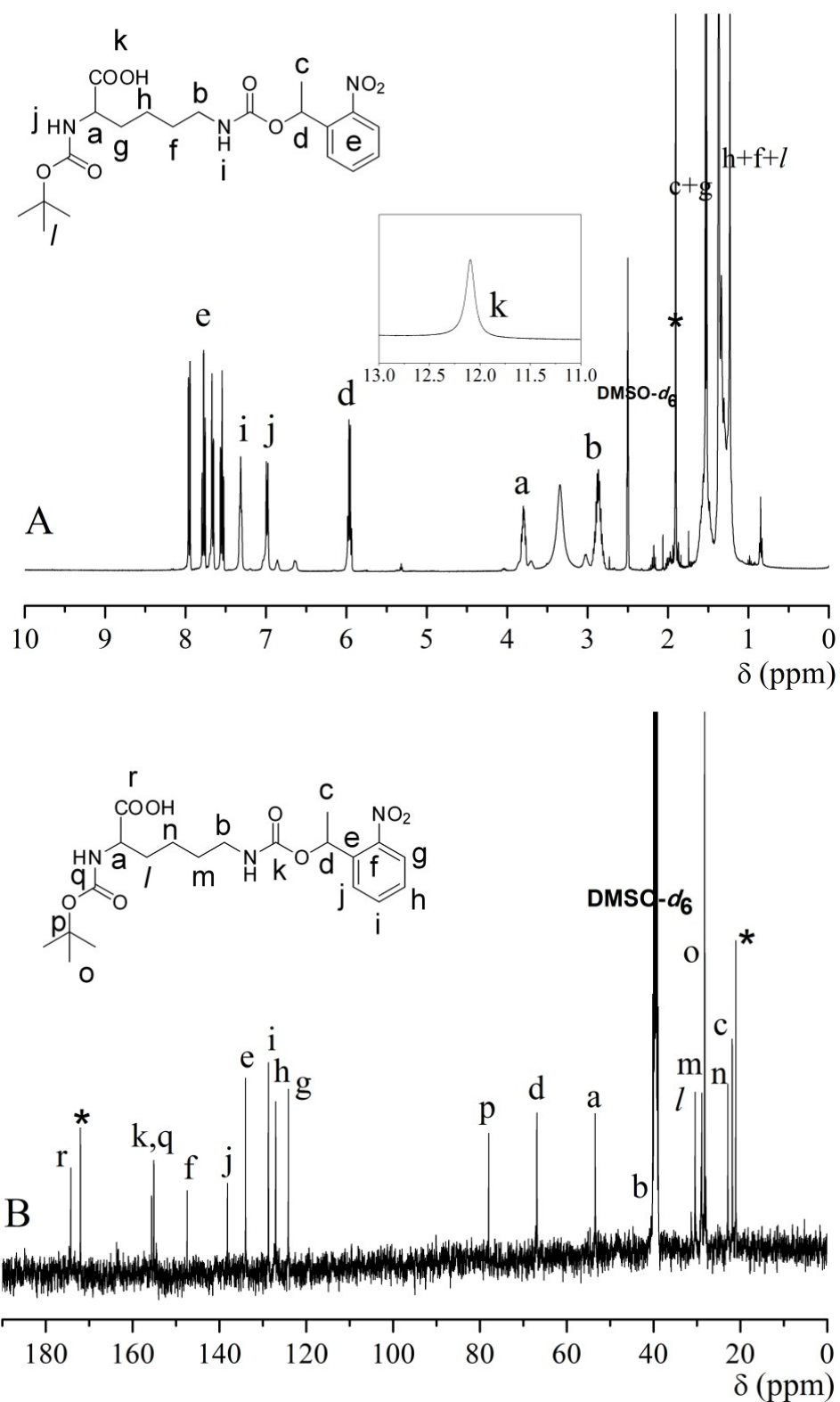


Figure S2. ^1H NMR (A) and ^{13}C NMR spectra of Boc-Lys(NPE)-OH (“*” denotes the AcOH solvent peaks).

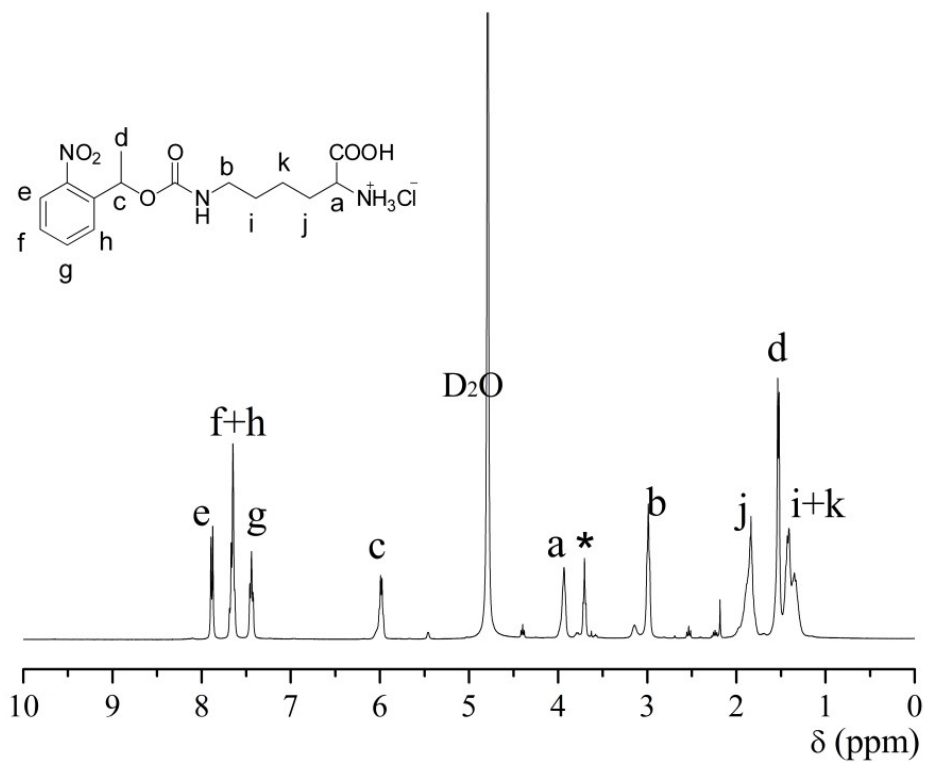


Figure S3. ¹H NMR spectrum of H-Lys(NPE)-OH·HCl (“*” denotes the THF solvent peaks).

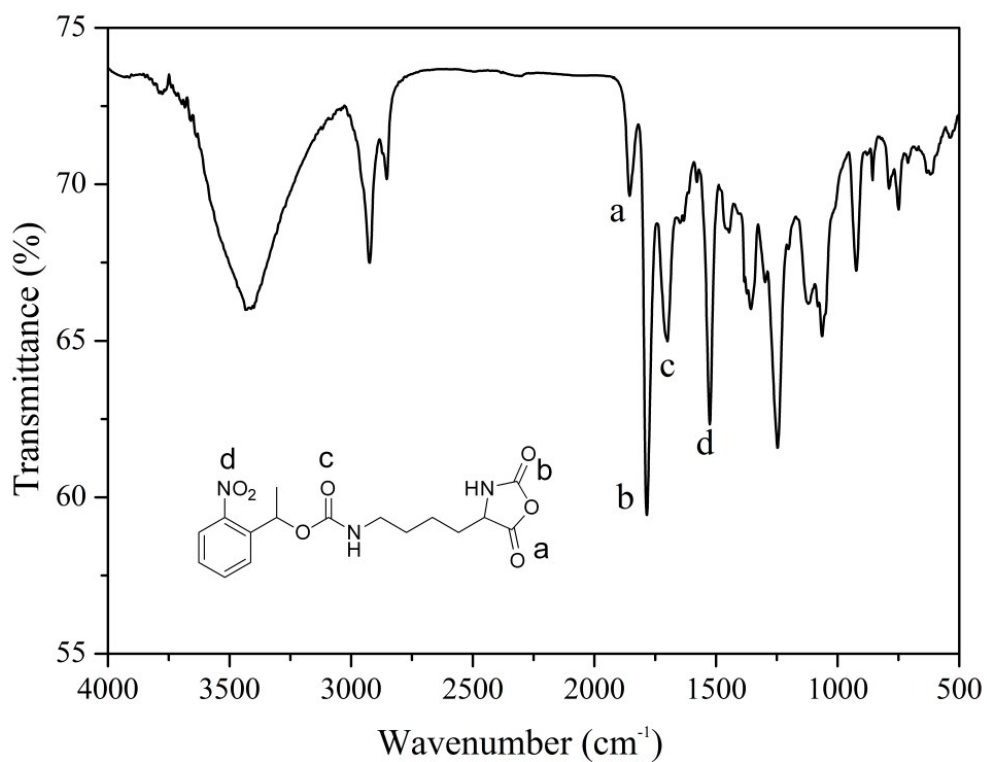


Figure S4. FT-IR spectrum of NPE-Lys NCA.

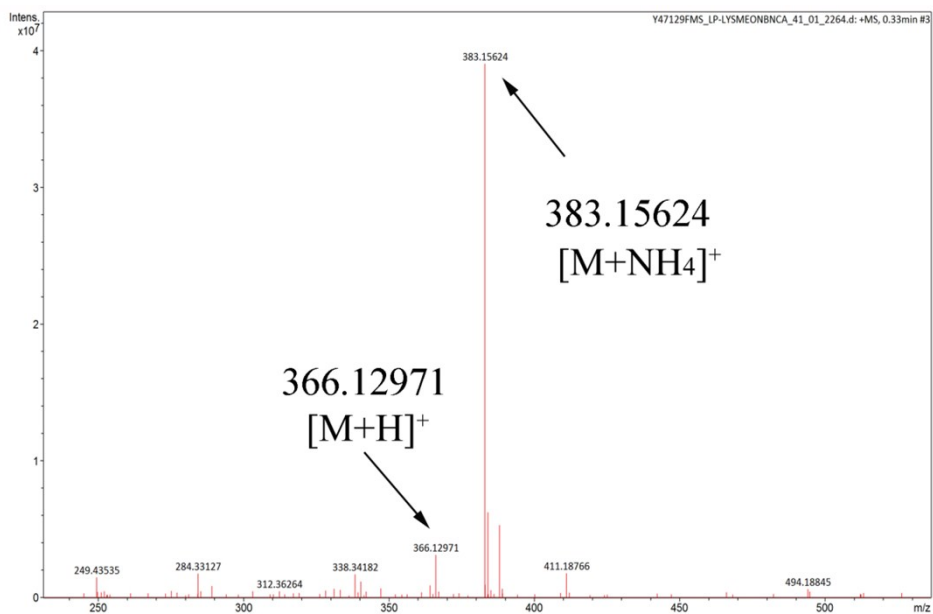


Figure S5. ESI-MS spectrum of NPE-Lys NCA.

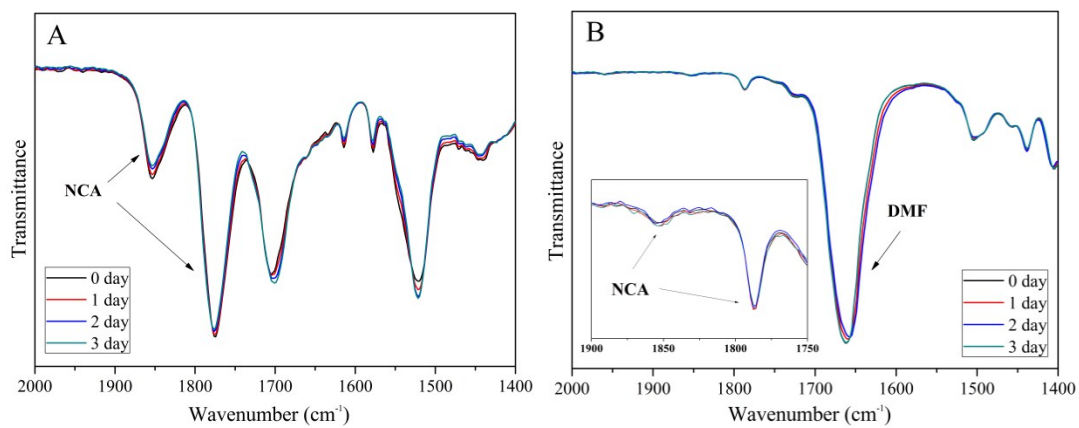


Figure S6. FT-IR monitoring the NPE-Lys NCA monomer consumption in the dark at 25 °C in both acetone (A) and DMF solution (B).

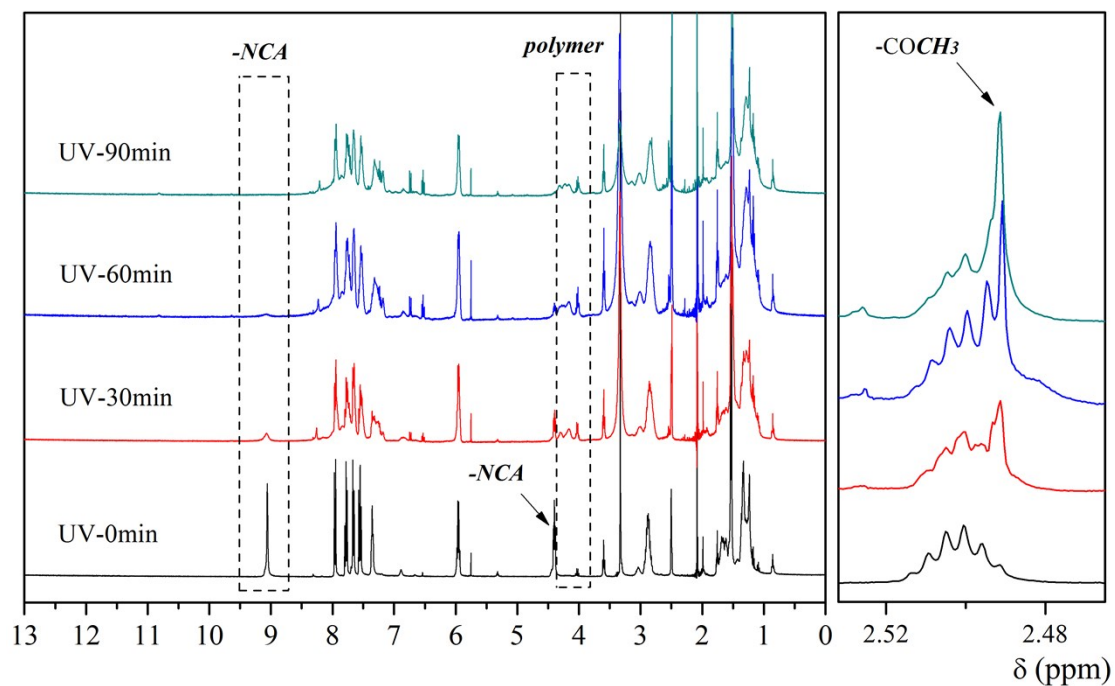


Figure S7. ^1H NMR spectra of NPE-Lys NCA in acetone upon 365 nm UV-irradiation at 16 mW/cm^2 for different irradiation times.

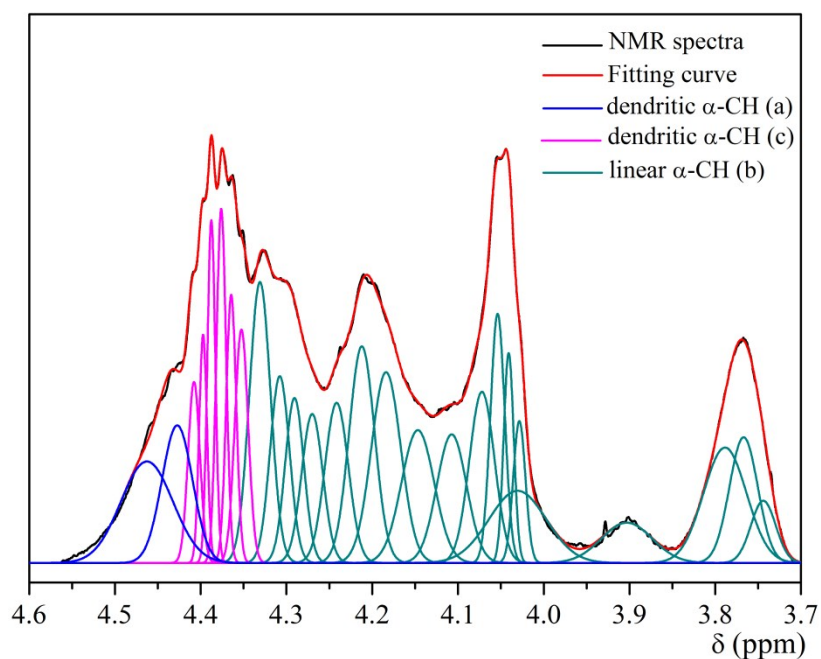


Figure S8. Peak fitting for $\alpha\text{-CH}$ in the ^1H NMR spectrum of sample P12 to calculate DB value by using software PeakFit (v 4.12).

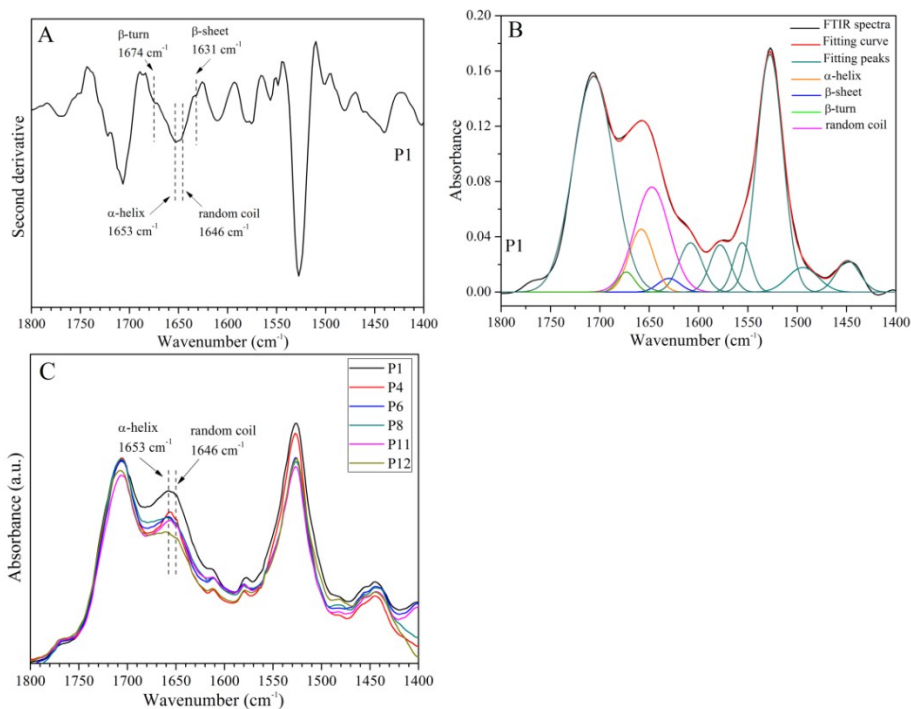


Figure S9. Second derivative (A) and FT-IR peak fitting results (B) of P1 in the solid state; and FT-IR spectra of P1, P4, P6, P8, P11 and P12 (C).

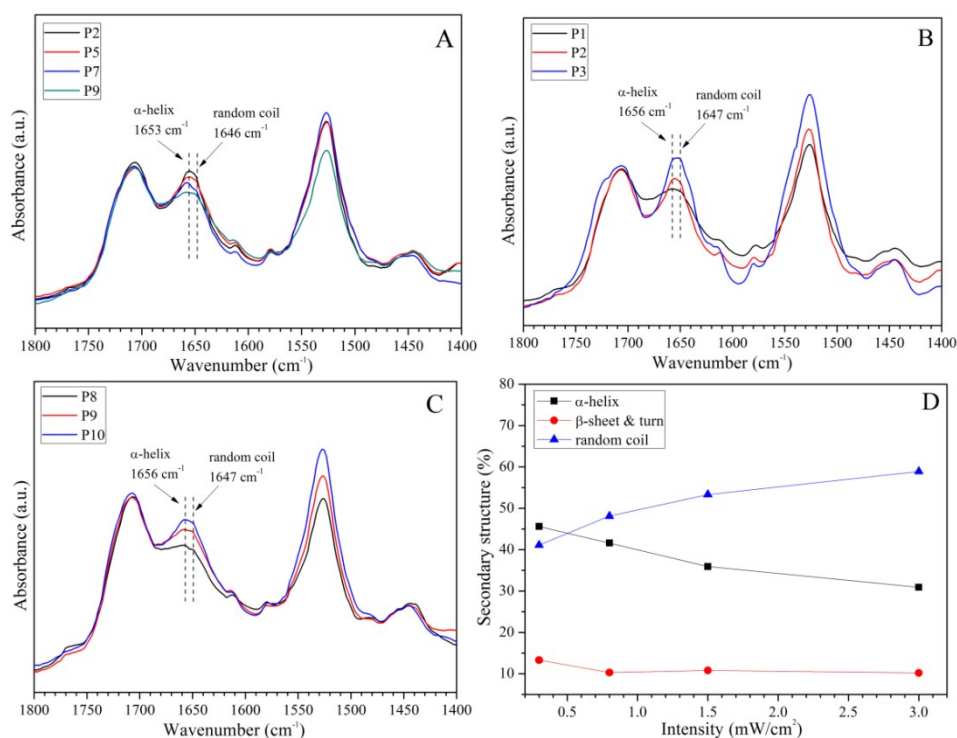


Figure S10. FT-IR spectra of P2, P5, P7 and P9 (A), P1-P3 (B) and P8-P10 (C) in the solid state, and the secondary structures of P2, P5, P7 and P9 synthesized at different irradiation intensities (D).

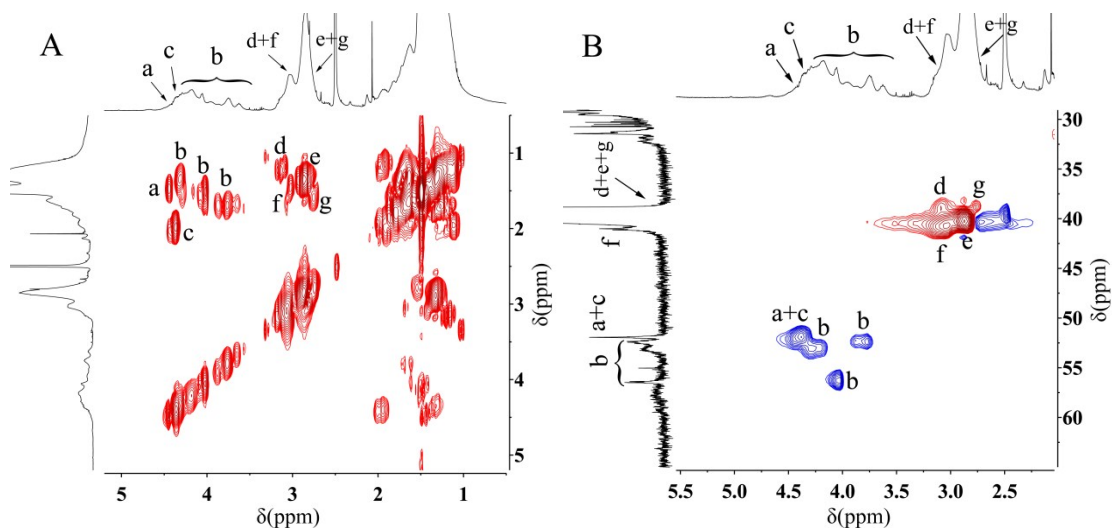


Figure S11. ^1H , ^1H -COSY spectrum (A) and ^1H , ^{13}C -HSQC spectrum (B) of P10.

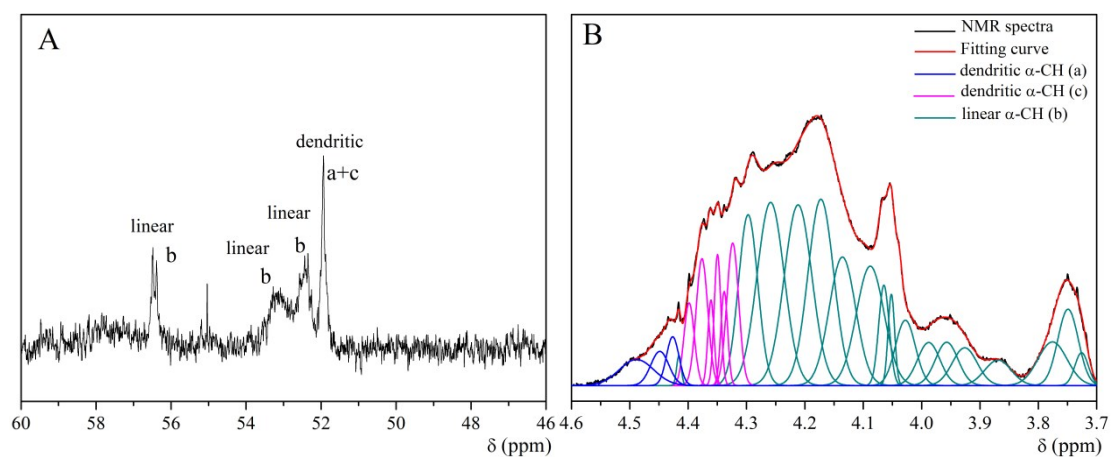


Figure S12. Quantitative ^{13}C NMR spectrum of P10 (A) and the peak fitting for α -CH in the ^1H NMR spectrum of P10 to calculate DB value (B).

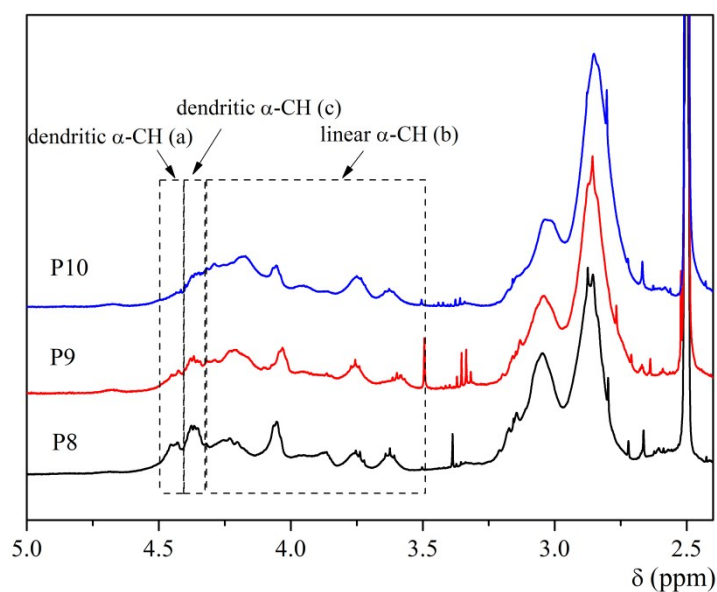


Figure S13. ^1H NMR spectra of P8-P10 synthesized under different N_2 flow rates.

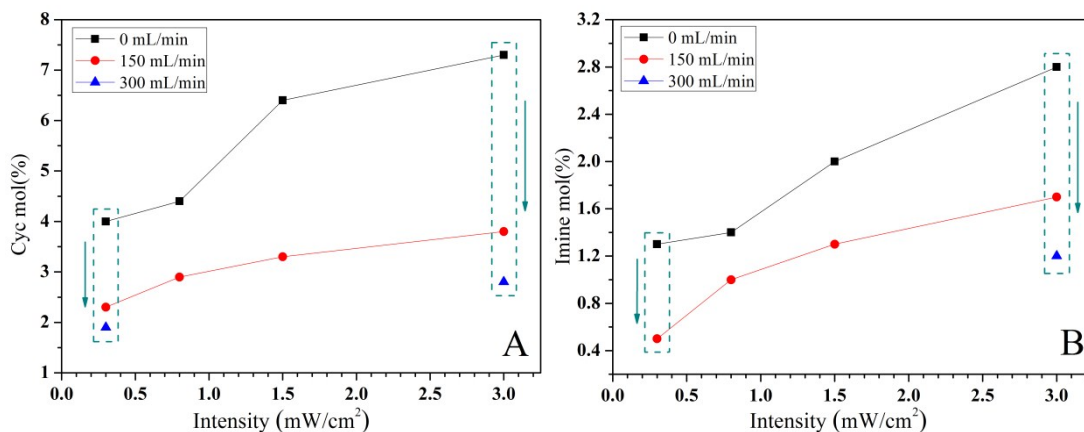


Figure S14. The effects of the irradiation intensity and/or N₂ flow rate on the contents of the cyclic dimer (A) and the imine (B).

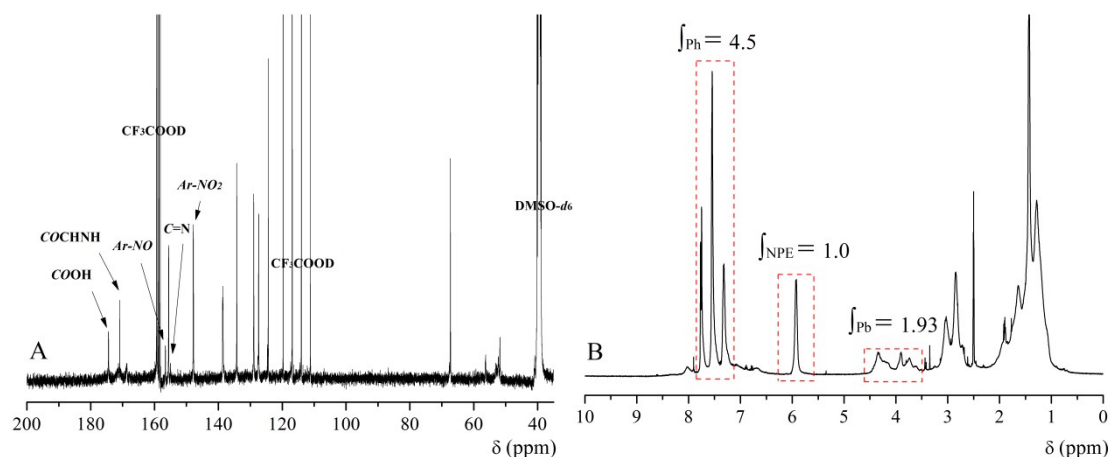


Figure S15. Both ¹³C NMR spectrum (A) and ¹H NMR spectrum in mixed solvents of DMSO-*d*₆ and CF₃COOD (v:v = 3:2) of P12.

Note: Imine molar percentage is calculated according to the following procedure. The integration of peak at 5.93 ppm attributable to the methine proton of NPE group is set as the reference $\int_{\text{NPE}} = 1.0$; the theoretical integrated area of phenylic region 7.77~7.31 ppm should be $\int_{\text{Ph}} = 4.0$ if without imination, so the extra area compared to the theoretical integration ($\Delta = 4.5 - 4.0 = 0.5$) comes from the phenylic signals of imination.

Therefore, imine mole percentage in the whole polypeptide backbone, $\int_{\text{Pb}} = 1.93$

$$f = \frac{(\int_{\text{NPE}} - \int_{\text{Ph}})/4}{\int_{\text{Pb}}} = \frac{(4.5 - 4)/4}{1.93} = 6.5 \text{ mol\%}.$$