

Physical Gelation of α -Helical Copolypeptides

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Statistical Distribution of AG and BLG Units in P(BLG_x-co-AG_{1-x})_n Copolypeptides

Using ^1H NMR, the ratio of BLG to AG units was measured throughout the polymerization reaction of different $\text{P}(\text{BLG}_x\text{-co-AG}_{1-x})_n$. For each polymerization reaction, a sample was taken out at regular intervals, precipitated and washed in methanol and measured by ^1H NMR in deuterated TFA ($\text{TFA-}d_1$). This ratio remained approximately constant during the polymerization reaction, which indicates that $\text{P}(\text{BLG}_x\text{-co-AG}_{1-x})_n$ are statistical copolymers.

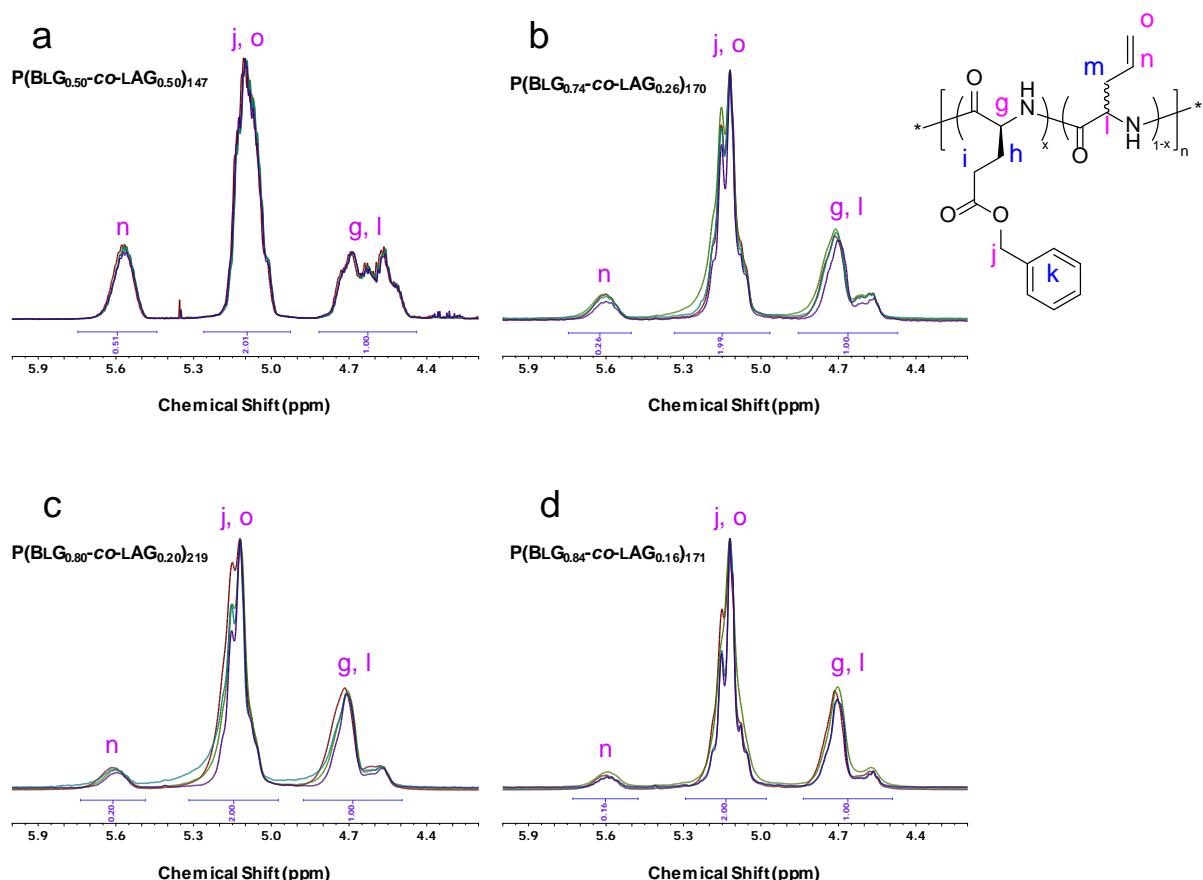


Figure S1. [4.2 - 6 ppm] sections of ^1H NMR spectra corresponding to polymerization samples taken at (a) (violet) 10 h, (blue) 24 h, (green) 32 h and (red) after termination; (b, c, d) (violet) 24 h, (blue) 48 h, (green) 72 h and (red) after termination; spectra were normalized to the 5.0-5.3 ppm peak

Homopolyptide PBLG₅₁ (originally presented in a former study¹)

¹H NMR, 400 MHz, TFA-*d*₁, δ (ppm) 7.26-7.24 (br, 255H), 5.60-5.13 (br, 102H), 4.69-4.66 (br, 51H), 3.29 (br, 2H), 2.58-1.94 (br, 593H), 1.53-1.50 (br, 2H), 1.36-1.34 (br, 6H), 0.86-0.83 (br, 3H)

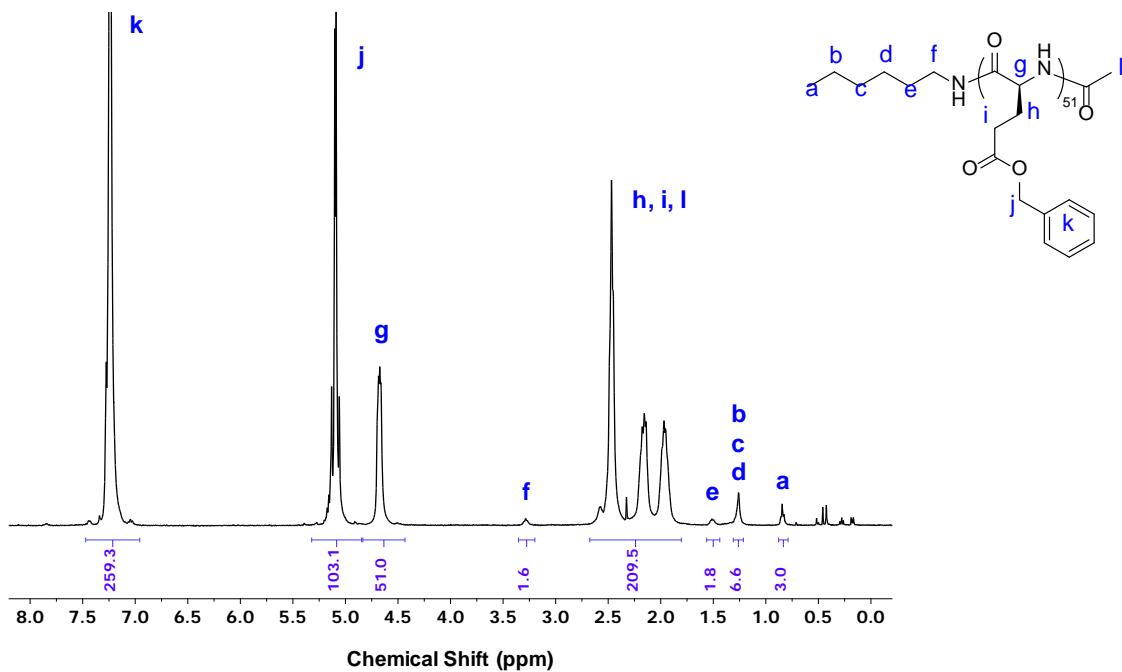


Figure S2. ^1H NMR spectrum (400 MHz) of PBLG₅₁ in TFA- d_1

Statistical Copolyptide P(BLG_{0.89}-co-DLAG_{0.11})₅₃ (originally presented in a former study¹)

¹H NMR, 400 MHz, TFA-*d*₁, δ (ppm) 7.25-7.23 (br, 235H), 5.59 (br, 6H), 5.16-5.02 (br, 106H), 4.71-4.53 (br, 53H), 3.27 (br, 2H), 2.53-1.94 (br, 203H), 1.51-1.48 (br, 2H), 1.33-1.25 (br, 6H), 0.85-0.82 (br, 3H)

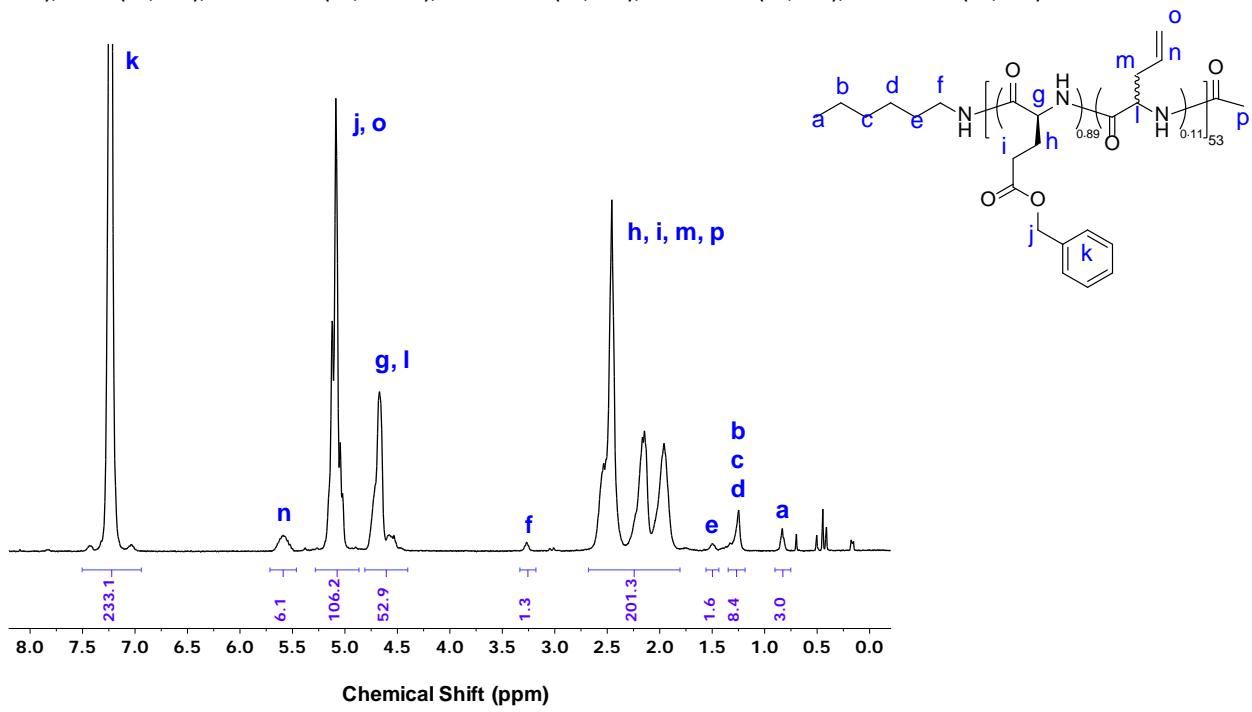


Figure S3. ^1H NMR spectrum (400 MHz) of $\text{P(BLG}_{0.89}\text{-}co\text{-DLAG}_{0.11}\text{)}_{53}$ in TFA- d_1

Statistical Copolypeptide P(BLG_{0.76}-co-DLAG_{0.24})₅₉ (originally presented in a former study¹)

¹H NMR, 400 MHz, TFA-*d*₁, δ (ppm) 7.28-7.26 (br, 225H), 5.62 (br, 14H), 5.19-5.06 (br, 118H), 4.74-4.57 (br, 59H), 3.29 (br, 2H), 2.57-1.98 (br, 211H), 1.54-1.51 (br, 2H), 1.36-1.28 (br, 6H), 0.88-0.85 (br, 3H)

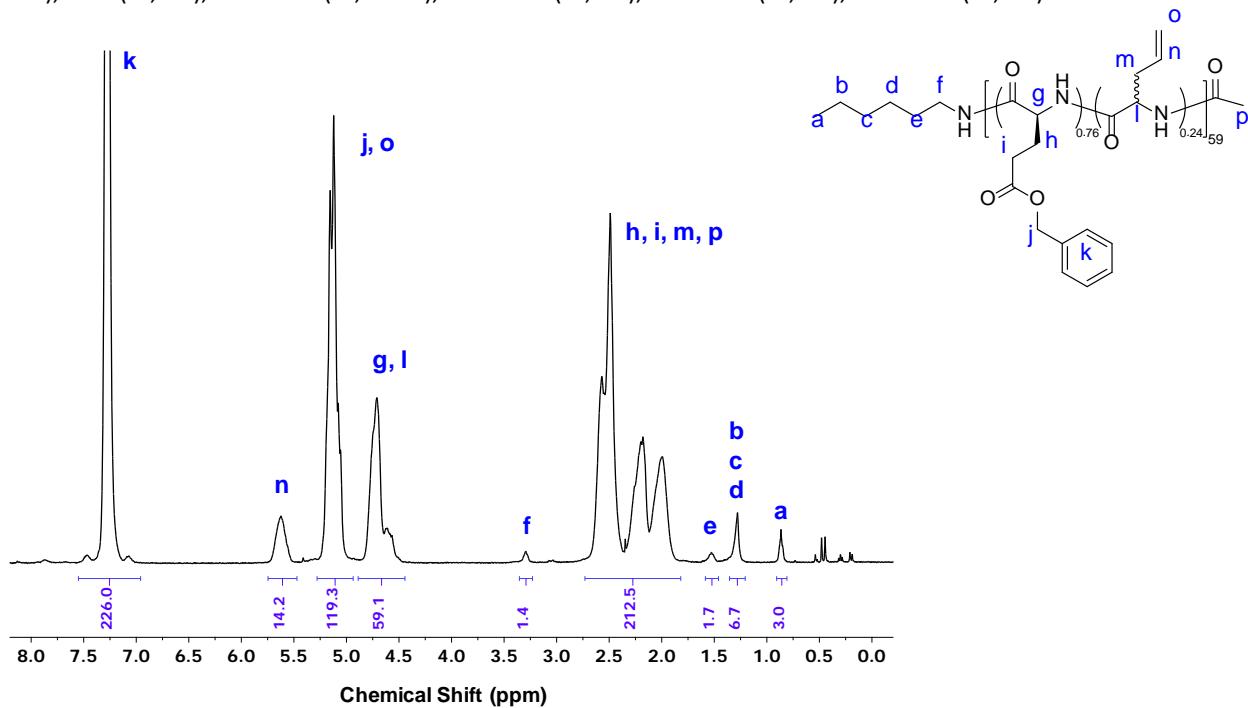


Figure S4. ¹H NMR spectrum (400 MHz) of P(BLG_{0.76}-co-DLAG_{0.24})₅₉ in TFA-*d*₁

Statistical Copolypeptide P(BLG_{0.77}-co-LAG_{0.23})₅₇ (originally presented in a former study¹)

¹H NMR, 400 MHz, TFA-*d*₁, δ (ppm) 7.29-7.27 (br, 220H), 5.60 (br, 13H), 5.19-5.06 (br, 114H), 4.74-4.57 (br, 57H), 3.30 (br, 2H), 2.59-1.99 (br, 205H), 1.55-1.51 (br, 2H), 1.37-1.28 (br, 6H), 0.88-0.85 (br, 3H)

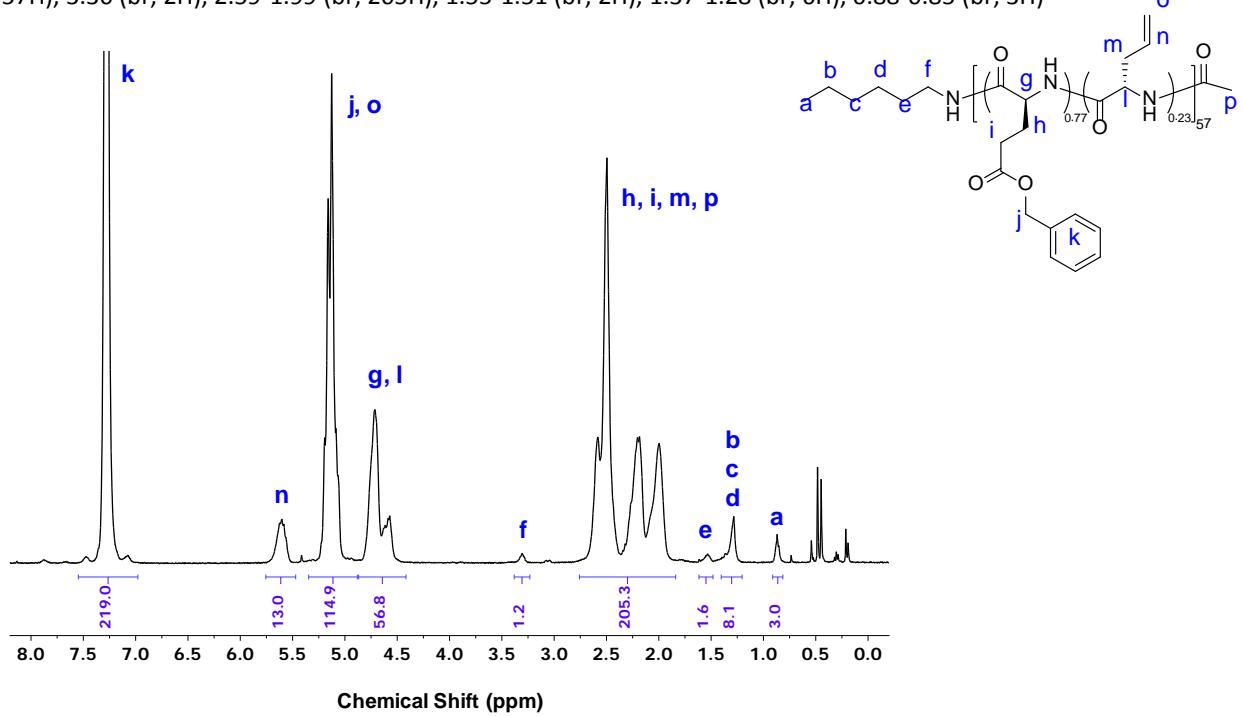


Figure S5. ¹H NMR spectrum (400 MHz) of P(BLG_{0.77}-co-LAG_{0.23})₅₇ in TFA-*d*₁

Statistical Copolyptide P(BLG_{0.74}-co-LAG_{0.26})₉₁ (originally presented in a former study)¹

¹H NMR, 400 MHz, TFA-*d*₁, δ (ppm) 7.22 (br, 335H), 5.56 (br, 24H), 5.15-5.02 (br, 182H), 4.73-4.52 (br, 91H), 3.23 (br, 2H), 2.55-1.94 (br, 319H), 1.54-1.48 (br, 2H), 1.33-1.23 (br, 6H), 0.81 (br, 3H)

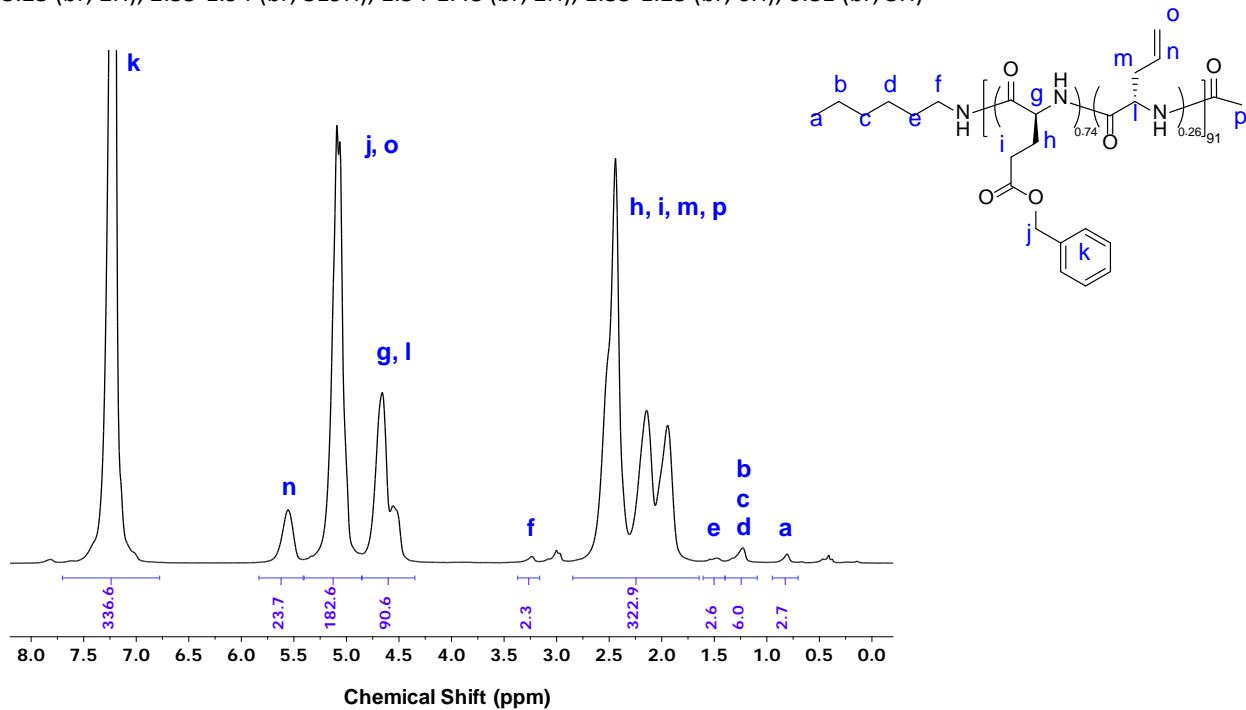


Figure S6. ¹H NMR spectrum (400 MHz) of P(BLG_{0.74}-co-LAG_{0.26})₉₁ in TFA-*d*₁

Statistical Copolyptide P(BLG_{0.78}-co-LAG_{0.22})₉₆

¹H NMR, 400 MHz, TFA-*d*₁, δ (ppm) 7.20 (br, 375H), 5.54 (br, 21H), 5.14-5.00 (br, 192H), 4.70-4.50 (br, 96H), 3.23 (br, 2H), 2.51-1.93 (br, 345H), 1.53-1.47 (br, 2H), 1.34-1.22 (br, 6H), 0.80 (br, 3H)

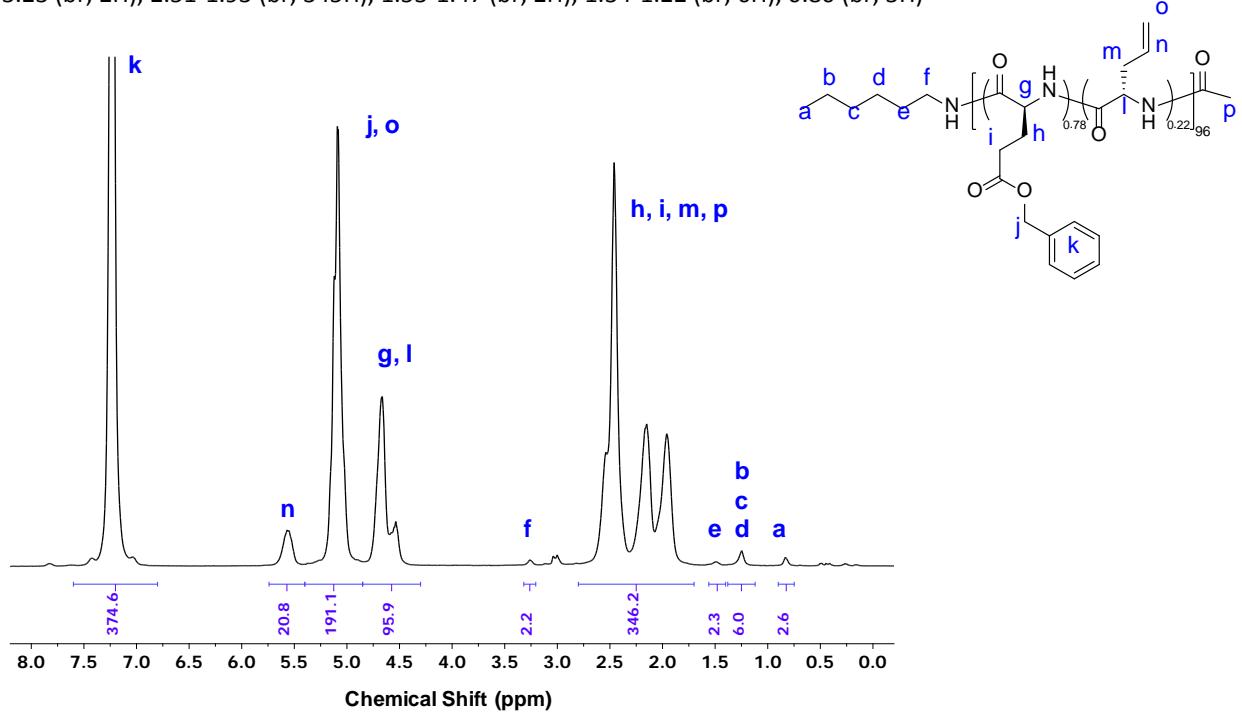


Figure S7. ¹H NMR spectrum (400 MHz) of P(BLG_{0.78}-co-LAG_{0.22})₉₆ in TFA-*d*₁

Statistical Copolyptide P(BLG_{0.74}-co-LAG_{0.26})₁₇₀ (originally presented in a former study)¹

¹H NMR, 400 MHz, TFA-*d*₁, δ (ppm) 7.20 (br, 625H), 5.54 (br, 45H), 5.14-5.00 (br, 340H), 4.70-4.50 (br, 170H), 3.23 (br, 2H), 2.51-1.93 (br, 593H), 1.53-1.47 (br, 2H), 1.34-1.22 (br, 6H), 0.80 (br, 3H)

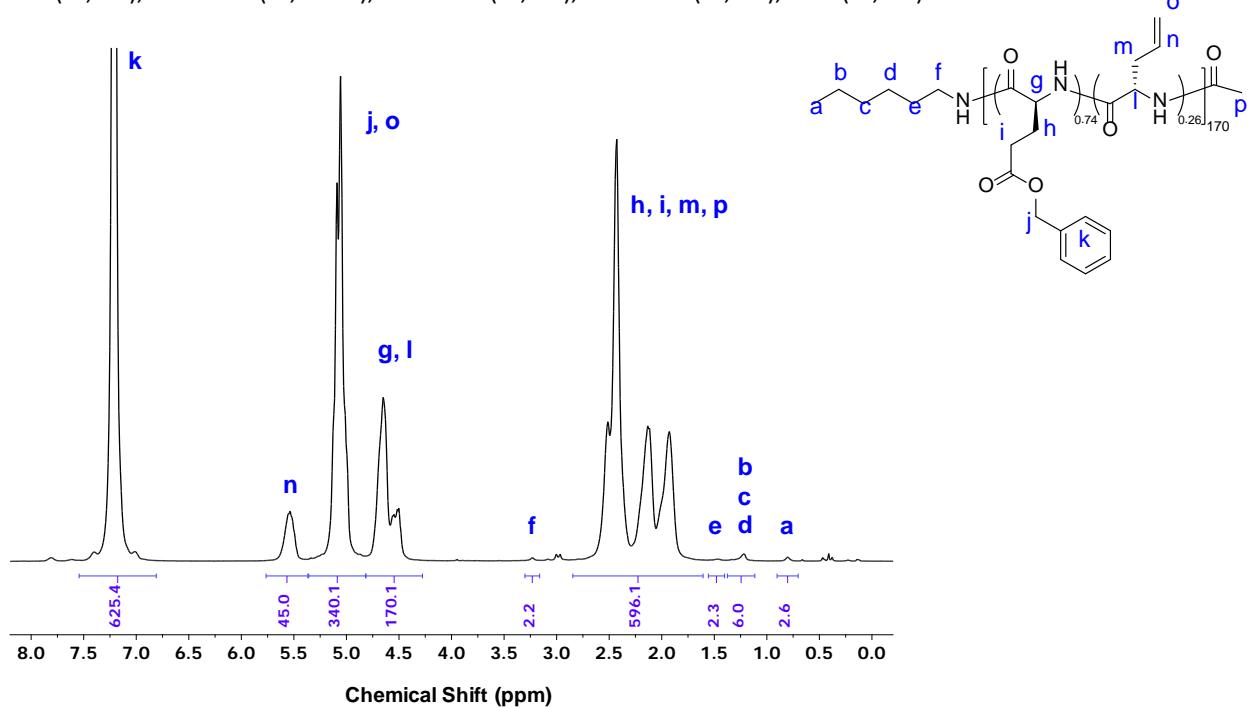


Figure S8. ¹H NMR spectrum (400 MHz) of P(BLG_{0.74}-co-LAG_{0.26})₁₇₀ in TFA-*d*₁

Statistical Copolyptide P(BLG_{0.80}-co-LAG_{0.20})₂₁₉

¹H NMR, 400 MHz, TFA-*d*₁, δ (ppm) 7.20 (br, 875H), 5.54 (br, 44H), 5.14-5.00 (br, 438H), 4.70-4.50 (br, 219H), 3.23 (br, 2H), 2.51-1.93 (br, 791H), 1.53-1.47 (br, 2H), 1.34-1.22 (br, 6H), 0.80 (br, 3H)

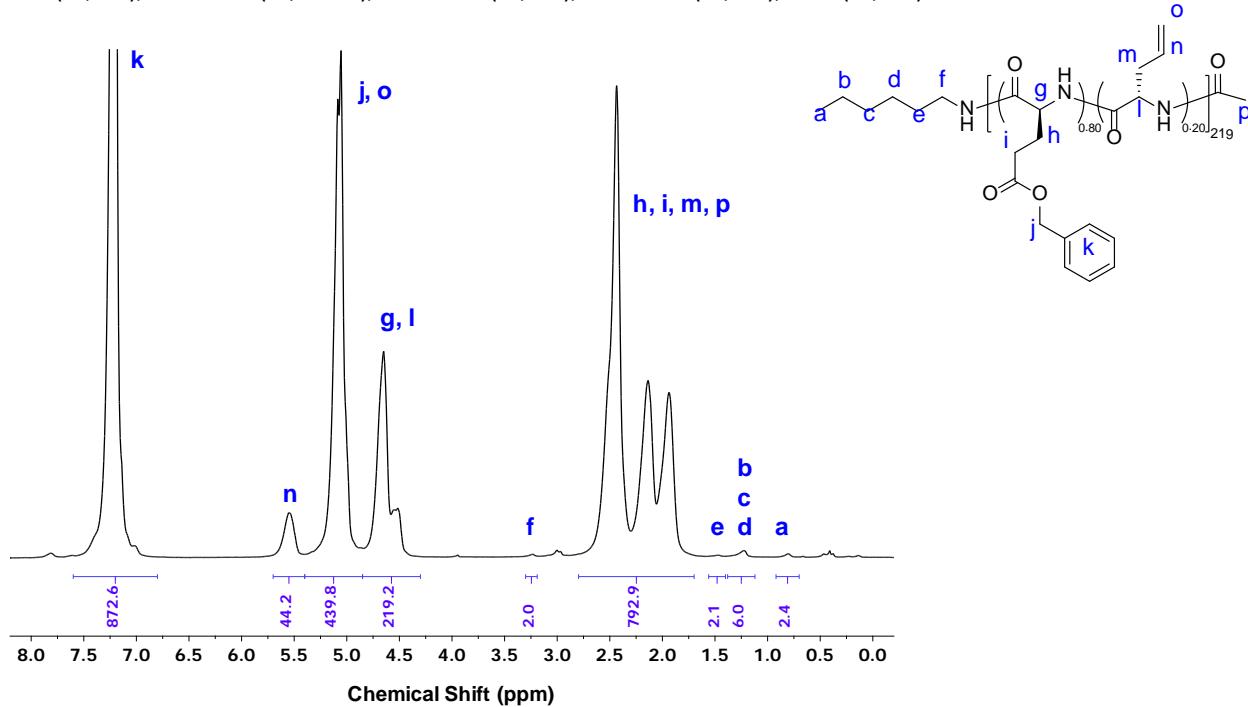


Figure S9. ¹H NMR spectrum (400 MHz) of P(BLG_{0.80}-co-LAG_{0.20})₂₁₉ in TFA-*d*₁

Statistical Copolypeptide P(BLG_{0.84}-co-LAG_{0.16})₁₇₁

¹H NMR, 400 MHz, TFA-*d*₁, δ (ppm) 7.20 (br, 715H), 5.54 (br, 28H), 5.14-5.00 (br, 342H), 4.70-4.50 (br, 171H), 3.23 (br, 2H), 2.51-1.93 (br, 631H), 1.53-1.47 (br, 2H), 1.34-1.22 (br, 6H), 0.80 (br, 3H)

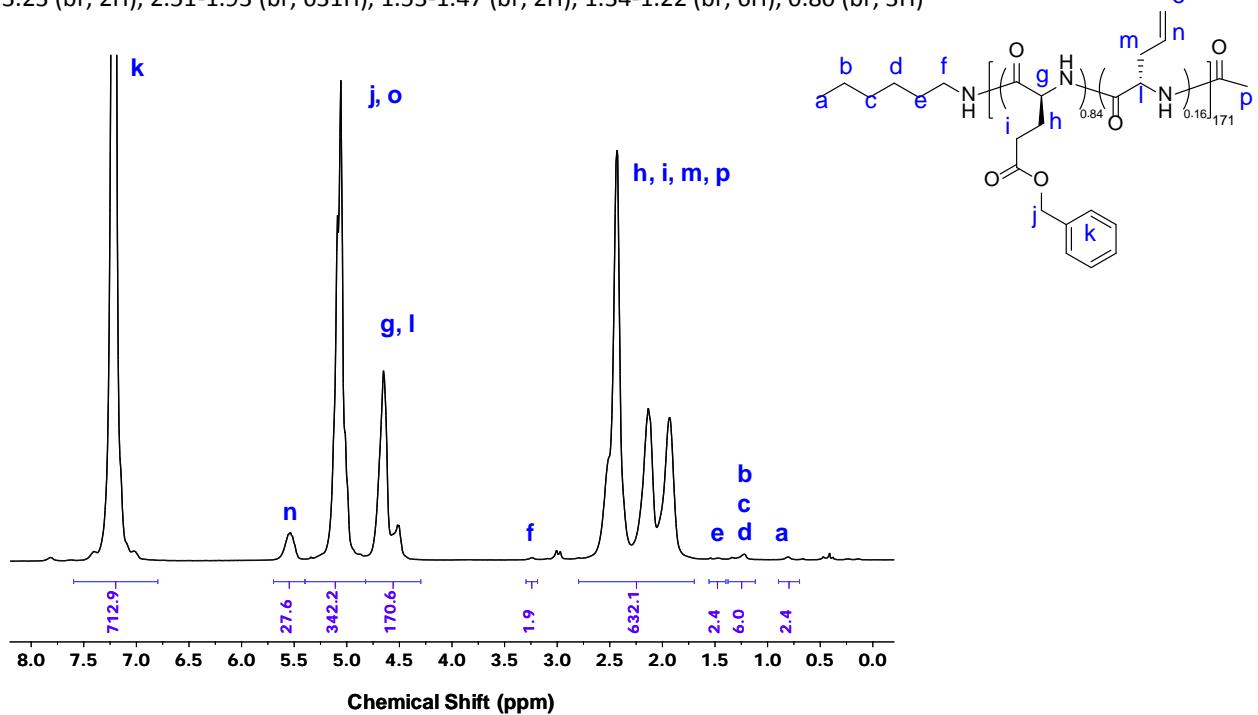


Figure S10. ¹H NMR spectrum (400 MHz) of P(BLG_{0.84}-co-LAG_{0.16})₁₇₁ in TFA-*d*₁

Size Exclusion Chromatography (SEC)

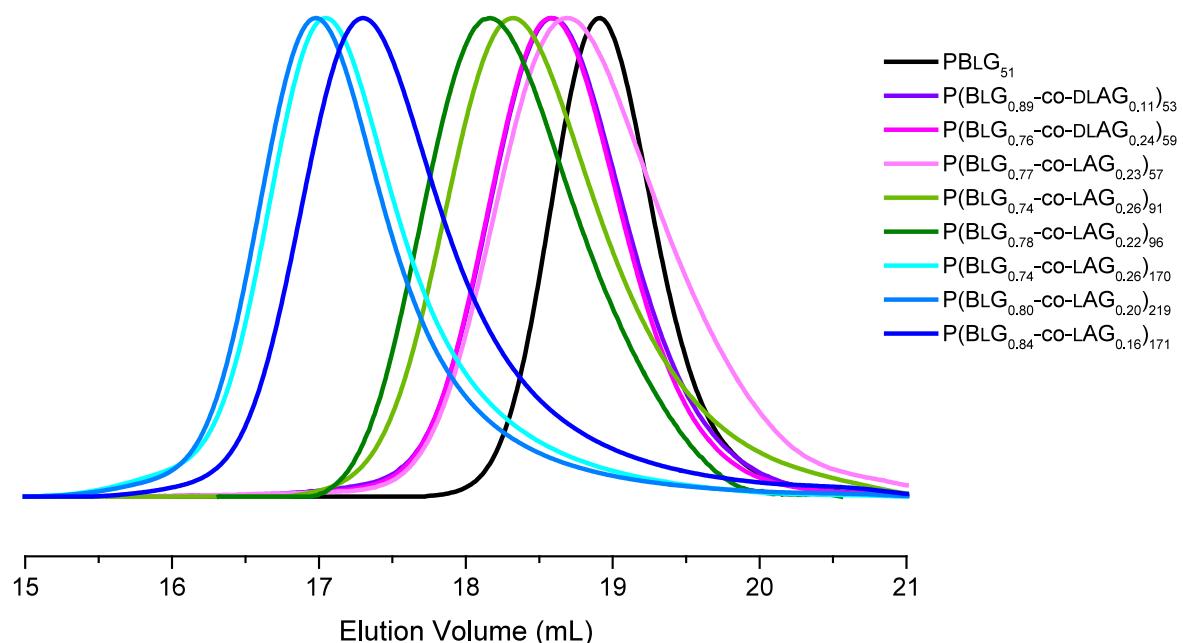


Figure S11. SEC-RI traces (eluent: NMP+LiBr) of PBLG and P(BLG_x-co-AG_{1-x})_n copolyptides

FTIR Spectroscopy

Like the Raman spectra, the FTIR spectra of $P(BLG_x\text{-}co\text{-}AG_{1-x})_n$ copolypeptides show an Amide I band composed of two peaks: a main α -helical peak at 1650 cm^{-1} (FTIR) and at 1653 cm^{-1} (Raman), and a secondary peak (noted α') at 1640 cm^{-1} (FTIR) and at 1643 cm^{-1} (Raman).

Relative to the α peaks, the α' peaks have a greater intensity in Raman than in FTIR spectroscopy. This can be explained by the different quantum mechanical selection rules for IR and Raman processes.² As a result of the deconvolution process performed on FTIR and Raman spectra, smaller peaks are likely to be tinged with a greater margin of error compared with larger peaks. This was a major driver for the choice of Raman spectra to help interpret rheological results.

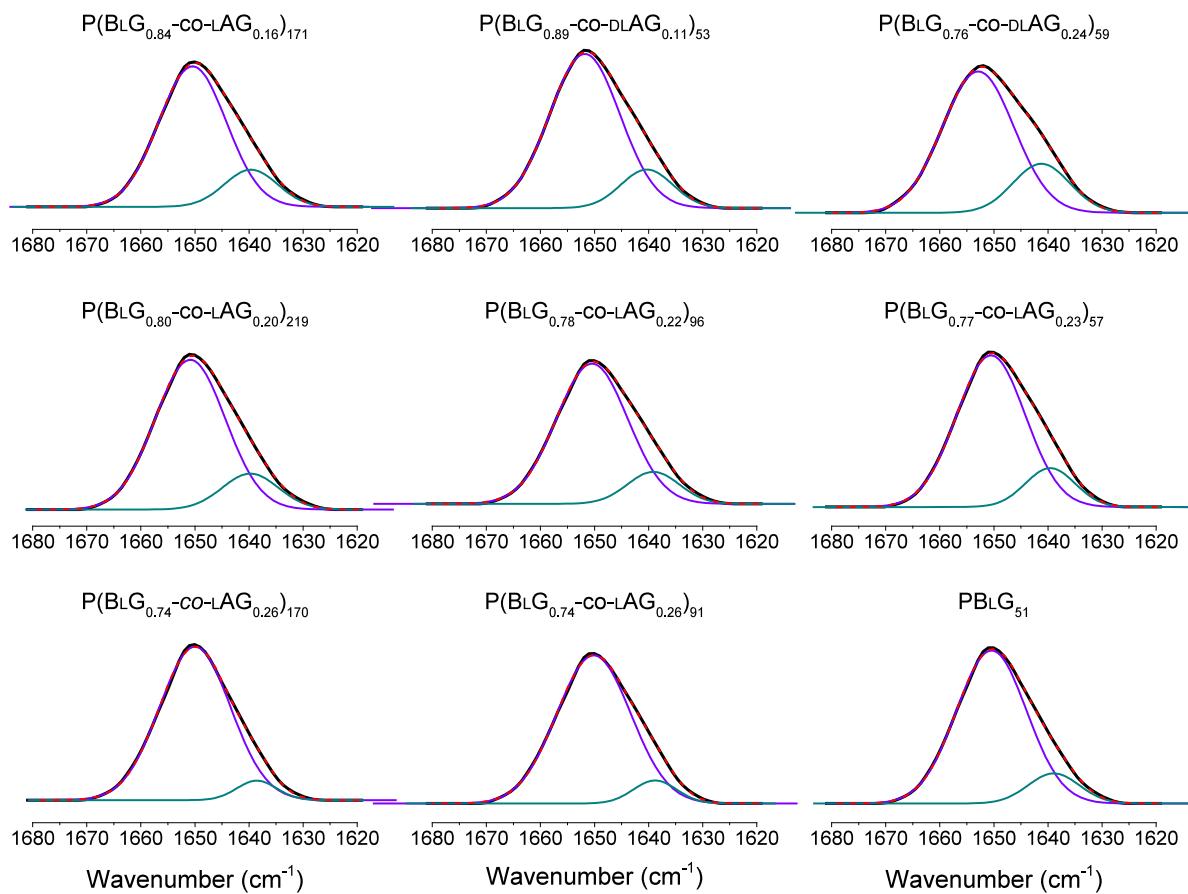


Figure S12. Amide I bands from FTIR spectra of PBLG and $P(BLG_x\text{-}co\text{-}AG_{1-x})_n$ copolypeptides, composed of a main peak (α peak, purple) at approximately 1650 cm^{-1} that corresponds to an α -helical conformation, and a secondary peak (α' peak, green) at approximately 1640 cm^{-1} , whose exact nature is discussed in this study

Copolypeptide Solution Properties

Table S1. Properties of PBLG and P(BLG_x-co-AG_{1-x})_n copolyptides studied in toluene (rheology) or toluene-*d*₈ (¹H NMR) at 20 g·L⁻¹

Copolypeptide	Crossover Modulus (Stiffness) ^a	Crossover Temperature (T _{gel}) ^a	Complex Viscosity (η^*) at 20 °C ^a	Temperature at which benzyl signal disappears in ¹ H NMR (± 5 °C) ^b
	(Pa)	(°C)	(mPa s)	(°C)
PBLG ₅₁	5	39	5900	10
P(BLG _{0.89} -co-DLAG _{0.11}) ₅₃	26	-26	520	-40
P(BLG _{0.76} -co-DLAG _{0.24}) ₅₉	27	-38	630	< -40
P(BLG _{0.77} -co-LAG _{0.23}) ₅₇	30	-27	150	-40
P(BLG _{0.74} -co-LAG _{0.26}) ₉₁	96	-35	780	< -40
P(BLG _{0.78} -co-LAG _{0.22}) ₉₆	54	-23	620	-40
P(BLG _{0.74} -co-LAG _{0.26}) ₁₇₀	158	-23	2270	-35
P(BLG _{0.80} -co-LAG _{0.20}) ₂₁₉	444	-12	3000	-20
P(BLG _{0.84} -co-LAG _{0.16}) ₁₇₁	331	-8	1450	-20

^a Determined by rheometry. ^b Determined by ¹H NMR.

Temperature-sweep ¹H NMR Spectroscopy

During a sol-gel transition, gelators are incorporated in a supramolecular solid-like network, which causes line broadening, loss of spectral resolution, and signal disappearance in ¹H NMR measurements.³⁻⁵ This phenomenon was observed in a temperature-sweep ¹H NMR experiment for dilute solutions of our polypeptides in toluene-*d*₈ (Figure S13 and S14). A good correlation was found between the disappearance of the benzene peak, noted k, and the gelation temperature measured by rheology (Table S1).

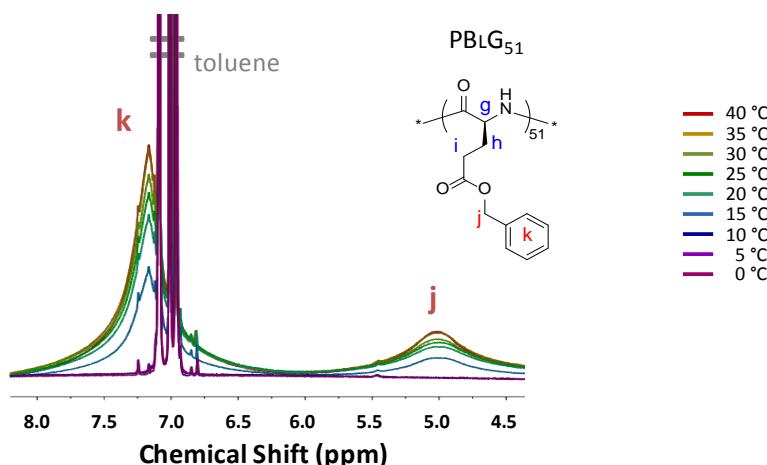


Figure S13. [4.5 - 8 ppm] sections of ¹H NMR spectra of a 20 g·L⁻¹ solution of PBLG₅₁ in toluene-*d*₈, taken at incrementally decreasing temperatures, from 40 °C to 0 °C; all spectra were normalized to the toluene signal at 7.01 ppm

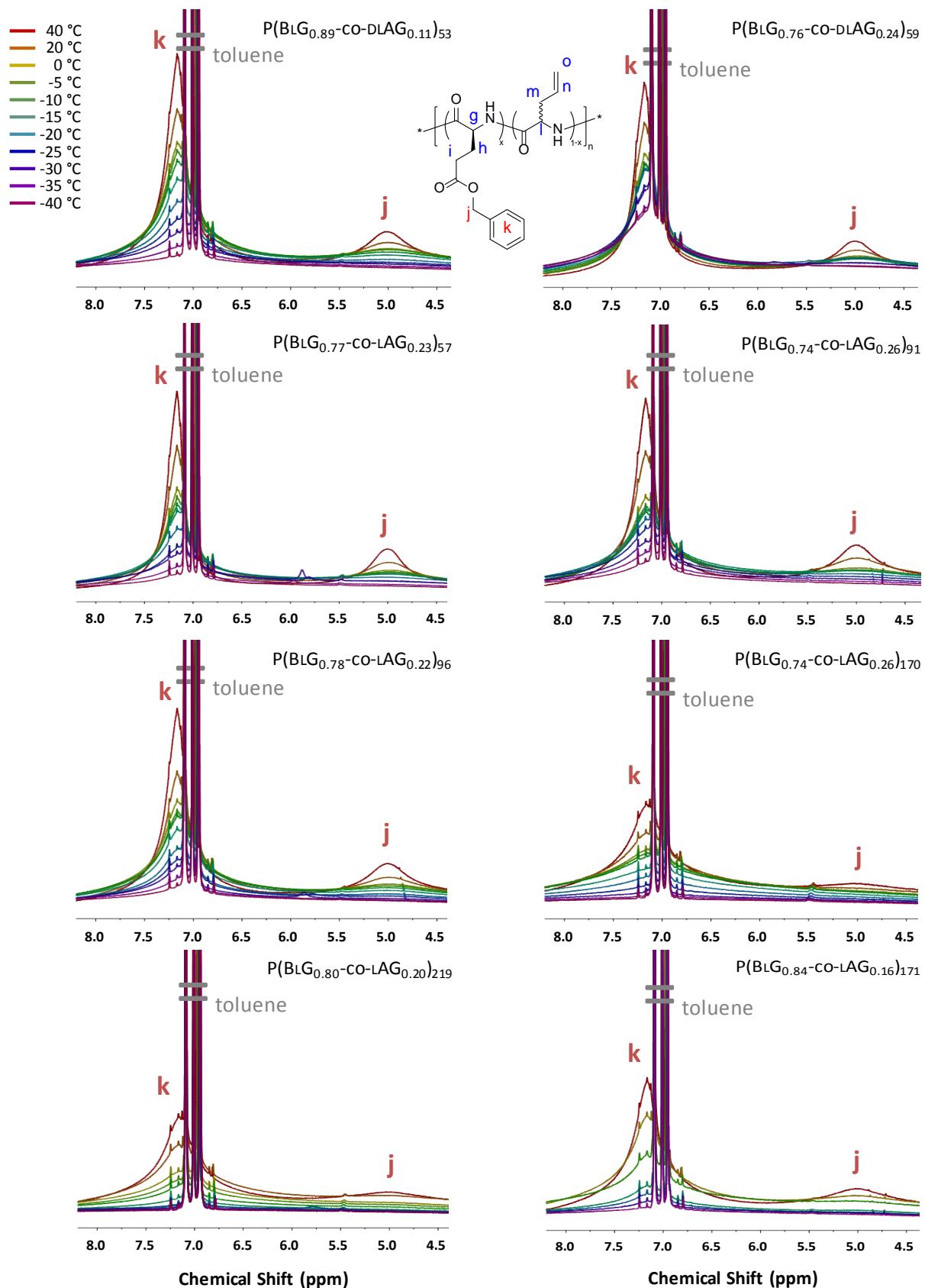


Figure S14. [4.5 - 8 ppm] sections of ^1H NMR spectra of 20 g·L $^{-1}$ solution of P(BLG $_x$ -co-AG $_{1-x}$) $_n$ copolypeptides in toluene- d_8 , taken at incrementally decreasing temperatures, from 40 °C to -40 °C; all spectra were normalized to the toluene signal at 7.01 ppm

Circular Dichroism (CD) Spectroscopy

1,1,1,3,3,3-Hexafluoroisopropanol (HFIP) is known to be a helicogenic solvent for PBLG and was therefore used for CD measurements.⁶ CD spectra of three selected $P(BLG_x\text{-}co\text{-}AG_{1-x})_n$ copolyptides and homopolymer $PBLG_{51}$ showed two minima at 208 nm and 222 nm, which are indicative of an α -helical conformation.

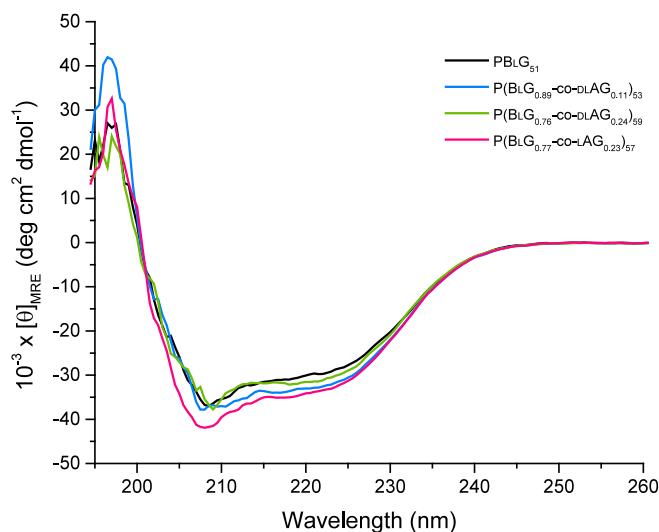


Figure S15. CD spectra of PBLG and $P(BLG_x\text{-}co\text{-}AG_{1-x})_n$ copolyptides in HFIP at $0.2 \text{ g}\cdot\text{L}^{-1}$

Raman Spectroscopy

The following spectra (Figure S16 and S17) were normalized to the 1610 cm^{-1} peak, which corresponds to the $\nu(\text{C-C})$ of the phenyl group of the BLG moiety.⁷ Since the Amide I band ($1620 - 1680 \text{ cm}^{-1}$) of our copolyptides comprises of the $\nu(\text{C=O})$ of both BLG and AG residues, its area, relative to the 1610 cm^{-1} peak, increases with increasing AG content.

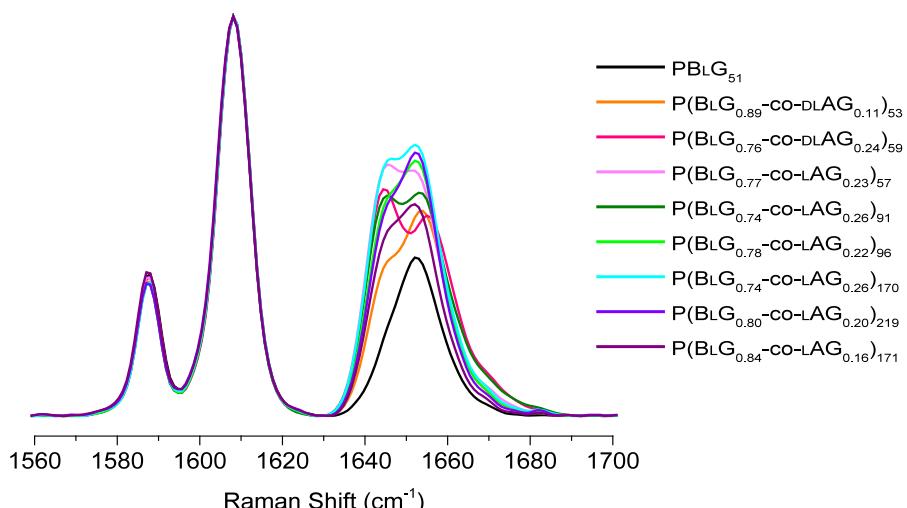


Figure S16. Amide I band of dry PBLG and $P(BLG_x\text{-}co\text{-}AG_{1-x})_n$ copolyptides after baseline correction and normalization

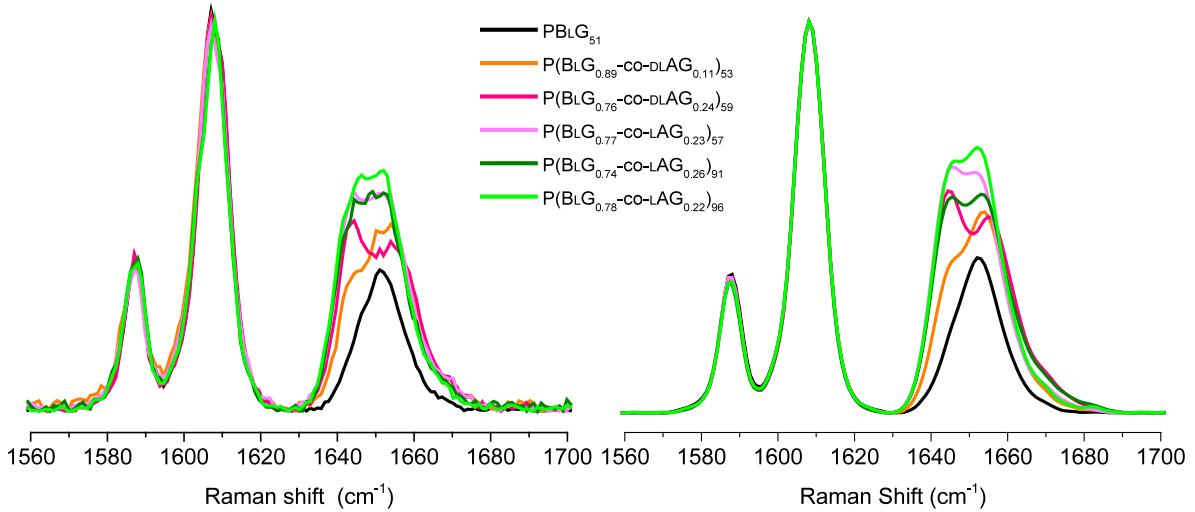


Figure S17. Amide I band of dry PBLG and P(BLG_x-co-AG_{1-x})_n copolypeptides (left) non compacted (originally presented in a former study¹), (right) compacted and showing an improved signal-to-noise ratio (present study)

Wide Angle X-Ray Scattering (WAXS)

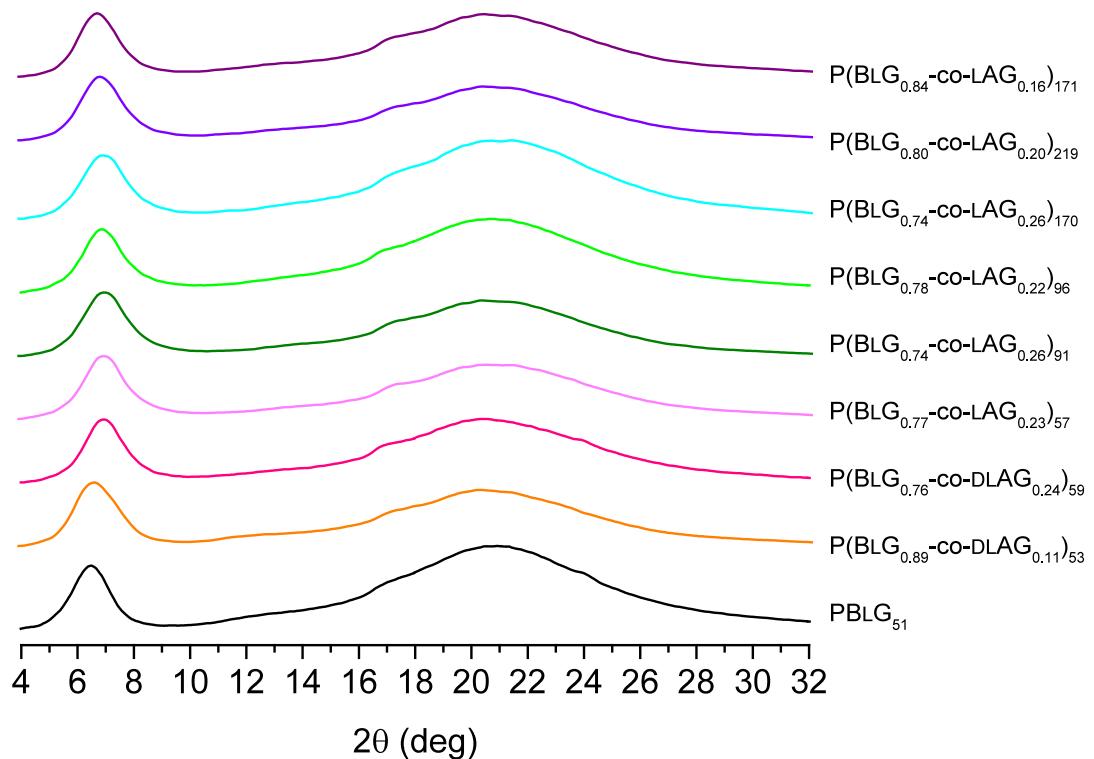


Figure S18. Full WAXS spectra of PBLG and P(BLG_x-co-AG_{1-x})_n copolypeptides, freeze-dried in dioxane; the peak at 6–7° corresponds to the d-spacing of ca. 1.3 nm from the pseudo-hexagonal arrangement of α -helical polymers; the shoulder at 17° corresponds to the pitch of α -helices (ca. 0.5 nm); the broad peak centered around 20–21° is the amorphous halo, resulting from the glass capillary (sample holder) and the measured polymers

References

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