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Kirkham, Steven, Castelletto, Valeria, Hamley, Ian William, Reza, Mehedi, Ruokolainen, Janne, Hermida-Merino, Daniel, Bilalis, Panayiotis and Iatrou, Hermis (2016) Self-assembly of telechelic tyrosine end-capped PEO and poly(alanine) polymers in aqueous solution. Biomacromolecules, 17 (3). pp. 1186-1197. ISSN 1525-7797 doi: https://doi.org/10.1021/acs.biomac.6b00023 Available at https://centaur.reading.ac.uk/63954/

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Published version at: http://dx.doi.org/10.1021/acs.biomac.6b00023
To link to this article DOI: http://dx.doi.org/10.1021/acs.biomac.6b00023

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Self-Assembly of Telechelic Tyrosine End-Capped PEO and Poly(alanine) Polymers in Aqueous Solution

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ABSTRACT: The self-assembly in aqueous solution of three novel telechelic conjugates comprising a central hydrophilic polymer and short (trimeric or pentameric) tyrosine end-caps has been investigated. Two of the conjugates have a central poly(ethylene oxide) (PEO) central block with different molar masses. The other conjugate has a central poly(α-lanine) (PAla) sequence in a purely amino-acid based conjugate. All three conjugates self-assemble into β-sheet based fibrillar structures, although the fibrillar morphology revealed by cryogenic-TEM is distinct for the three polymers—in particular the Tyr5-PEO6k-Tyr5 forms a population of short straight fibrils in contrast to the more diffuse fibril aggregates observed for Tyr5-PEO2k-Tyr5 and Tyr3-PAla-Tyr3. Hydrogel formation was not observed for these samples (in contrast to prior work on related systems) up to quite high concentrations, showing that it is possible to prepare solutions of peptide—polymer-peptide conjugates with hydrophobic end-caps without conformational constraints associated with hydrogelation. The Tyr5-PEO6k-Tyr5 shows significant PEO crystallization upon drying in contrast to the Tyr5-PEO2k-Tyr5 conjugate. Our findings point to the remarkable ability of short hydrophobic peptide end groups to modulate the self-assembly properties of polymers in solution in model peptide-capped “associative polymers”. Retention of fluidity at high conjugate concentration may be valuable in potential future applications of these conjugates as bioresponsive or biocompatible materials, for example exploiting the enzyme-responsiveness of the tyrosine end-groups.

INTRODUCTION

Conjugation of polymers with peptides enables the unique properties of each component to be combined, leading to novel biomacromolecules with a diversity of potential applications.1−8 In particular, polymers can be synthesized inexpensively and attached to peptides, modifying their solubility and aggregation properties. On the other hand, modification of polymers with peptides provides biofunctionality of many types, depending on the peptide and on the mode of incorporation, i.e., whether laterally attached where the peptide motif forms part of the polymer repeating unit, or whether attached at one or both ends of the polymer. Novel materials can result from the combination of polymer and peptide including lyotropic liquid crystal phases in aqueous solution9−11 or hydrogels.12−17

Hydrophilic polymers may be modified with hydrophobic end groups to create telechelic polymers, which among other properties resulting from the noncovalent association of the end groups, possess useful rheological behavior. This is exploited in associative polymers18−20 widely used as thickeners in a variety of applications, and in commercially available polyethylene glycol (PEG)-based polymers such as reverse Pluronics,20 for example. In the field of peptide biomaterials, it has been proposed that polymers such as PEG or PEO (poly(ethylene oxide)) may be used as inert spacers within peptide sequences to improve stability and circulation time of the biomolecules for applications such as gene delivery,21 or cell signaling.22 In the field of purely peptide/protein systems (i.e., systems not incorporating synthetic polymers), the properties of gels of genetically expressed proteins comprising a hydrophilic random coil-like middle block and collagen-like end blocks have been investigated by the groups of Tirrell12,23 and Cohen-Stuart.24−26

There are still few papers investigating the properties of self-assembled telechelic peptide—polymer—peptide conjugates. In a recent paper, the Hamley group reported on the gel−sol transition of conjugates of PEO with hydrophobic dipeptides,
including dityrosine and diphenylalanine. The conjugates were synthesized by coupling the dipeptides to bis-aminopropyl PEO of molar mass 1500 g mol$^{-1}$. The Fmoc (N-fluorenyl-9-methoxycarbonyl) protecting group (used in the peptide synthesis) was retained in some of the conjugates prepared, at one or both termini. This influences the hydrophile–lipophile balance and hence has a significant effect on the self-assembly behavior. Remarkably, one conjugate comprising dityrosine end-caps and with C-terminal Fmoc was found to undergo a gel–sol transition near body temperature, and this was shown to be driven by the loss of β-sheet secondary structure, associated with the formation of extended fibrils. The Lendlein group later prepared similar telechelic conjugates of desaminotyrosine (DAT) or desaminotyrosyl-tyrosine (DATT) with a linear PEG3000 midblock and also a form arm-PEG conjugate. A critical aggregation concentration was noted for the linear midblock conjugate, pointing toward self-assembly at high concentration. The rheological properties of hydrogels formed by linear and star-shaped oligo-ethylene glycol/DATT conjugates were examined. The same group also investigated the structural, conformational and rheological properties of gelatin (which has a collagen-based helical structure) functionalized with DAT or DATT at lysine residues. Micelles have been reported for PLL−PEG−PLL copolymers, where PLL denotes poly(L-leucine), the PLL adopting an α-helical conformation. Release of encapsulated drug was also investigated and sustained release over times 100−200 h was observed. Fibrillar hydrogels have recently been reported for a PAla−PGlu−PAla [PGlu: poly(L-glutamic acid)] conjugate self-assembling in aqueous buffer solution.

In the present paper, we investigate the self-assembly in aqueous solution of conjugates comprising telechelic polymers capped with short (pentameric or trimeric) tyrosine sequences. The creation of tyrosine-functionalized polymers is of interest in the development of novel aqueous biomaterials and hydrogel systems. In contrast to our previous study, the polymers are prepared by living NCA polymerization techniques, which give greater flexibility to tailor block length and which enable here the creation of novel purely peptidic conjugates with a central poly(L-alanine) block. The central core polymer in these designed telechelic conjugates is either PEO with molar mass of 2 kg mol$^{-1}$ or 6 kg mol$^{-1}$ or poly(L-alanine). According to most hydrophobicity scales, tyrosine is considered hydrophobic (although not according to the Kyte−Doolittle scale and capable of undergoing aromatic stacking interactions. Tyrosine has interesting functionality, as it is involved in cell signaling and is responsive to enzymes including kinases and phosphatases, which add and delete phosphate groups, respectively. It also has interesting fluorescence properties. Alanine is somewhat hydrophilic, at least according to the White−Wimley hydrophobicity scale, and so the Tyr3-PAla-Tyr3 conjugate has a hydrophobic−hydrophilic−hydrophobic architecture analogous to that of the two PEO conjugates, although with a purely amino acid based architecture (which may be of relevance to future applications where bioactivity is important). We find using a comprehensive suite of characterization measurements that all three novel conjugates form β-sheet-based fibrillar structures via self-assembly in aqueous solution, although the fibril morphology is distinct for the three conjugates. Unexpectedly, hydrogel formation was not observed up to rather high polymer concentration, enabling the production of high polymer/peptide conjugate content liquids. The influence of PEO crystallization on drying films is examined for the Tyr5-PEO2k-Tyr5 and Tyr5-PEO6k-Tyr5 conjugates, representing the first study to our knowledge on crystallization in this class of polymer/peptide conjugate that provides insight into PEO crystallization when constrained within a telechelic architecture.

**EXPERIMENTAL SECTION**

**Materials.** *N*-(tert-Butoxycarbonyl)-L-tyrosine, Boc-Tyr(tBu)-OH, (99%) was purchased from Bachem. Poly(ethylene oxide) bis(amine) with molecular weights of 5.9 × 10^{3} g/mol and 2.0 × 10^{3} g/mol were purchased from Aldrich and were used as difunctional macroinitiators.
The purity of Tyr(tBu)-NCA was condensed overnight, under high vacuum, to a white solid precipitate. Tyr(tBu)-NCA was isolated by being poured into 600 mL of hexane under vigorous stirring. Finally, the yellow suspension. The suspension was filtered and then transferred to a round-bottom flask. A quantity 40 mL of ethyl acetate was distilled giving a clear yellow solution. Then the solution was poured into 700 mL of hexane, and a yellow solid was precipitated. The solid was filtered and then transferred to a round-bottom flask. A quantity 40 mL of ethyl acetate was distilled giving a yellow suspension. The suspension was filtered, and the filtrate was poured into 600 mL of hexane under vigorous stirring. Finally, the white solid precipitate Tyr(tBu)-NCA was isolated by filtration and dried under high vacuum, to afford 1.85 g (7.0 mmol, 79% yield). The purity of Tyr(tBu)-NCA was confirmed by 'H NMR spectroscopy, shown in Supporting Information Figures S1 and S2, respectively.

Synthesis of N-Carboxy Anhydride of l-Alanine (Ala-NCA). The synthesis process of the l-alanine NCA (Ala-NCA) was accomplished according to a previously published method. The synthetic method followed for the Tyr5-PEO6k-Tyr5 was similar to that described above. Briefly, 0.95 g (0.16 mmol) of PEO bis(amine) with Mₙ = 5.9 × 10³ g/mol and 0.40 g (1.52 mmol) of Tyr(tBu)-NCA were used. The reaction takes 3–4 h.

The PEO precursors as well as the final copolymers containing PEO with Mₙ = 2.0 × 10³ g/mol or 5.9 × 10³ g/mol were characterized by 'H NMR spectroscopy (Figure 1) and size exclusion chromatography featuring a two-angle laser light scattering detector (Figure 2).

Synthesis of Poly(l-tyrosine)–Poly(l- alanine)–Poly(l-tyro- sine). Polymerization was carried out with ethylendiamine as the initiator. The sequential addition methodology was used for the synthesis of the triblock polypeptide. Ala-NCA (0.36 g) was diluted in 10 mL of DMF, and 2 mL of ethylendiamine solution in DMF (1.89

Figure 1. 'H NMR spectrum of the Tyr₅-PEO₂k-PTyr₅ triblock copolymer. Two sharp solvent peaks are assigned to DMSO at 2.54 ppm and one from water in DMSO at 3.33 ppm. Ethylenediamine (≥99%) was obtained from Aldrich and was left to dry over a sodium mirror for 24 h. It was then distilled into precalibrated ampules, diluted with purified DMF to the appropriate concentration, subdivided into ampules, and stored under high vacuum at −20 °C. Thionyl chloride (99.7%, Acros Organics) was distilled prior to use. The solvent purification was performed using standard high vacuum techniques reported elsewhere.

Synthesis of Tyr₅-PEO₂k-PTyr₅. The synthetic method followed for the Tyr₅-PEO₂k-PTyr₅ was similar to that described above. Briefly, 0.95 g (0.16 mmol) of PEO bis(amine) with Mₙ = 5.9 × 10³ g/mol and 0.40 g (1.52 mmol) of Tyr(tBu)-NCA were used. The reaction takes 3–4 h.

The PEO precursors as well as the final copolymers containing PEO with Mₙ = 2.0 × 10³ g/mol or 5.9 × 10³ g/mol were characterized by 'H NMR spectroscopy (Figure 1) and size exclusion chromatography featuring a two-angle laser light scattering detector (Figure 2). From the FTIR spectrum (Supporting Information Figure S2), it is obvious that the peak of the Boc- protection of the amine groups at 1736 cm⁻¹ has completely disappeared after the formation of the NCA, while the carbonyl groups of the cyclic N-carboxy anhydride.

Synthesis of Poly(l-tyrosine)–Poly(l- alanine)–Poly(l-tyro- sine). Polymerization was carried out with ethylendiamine as the initiator. The sequential addition methodology was used for the synthesis of the triblock polypeptide. Ala-NCA (0.36 g) was diluted in 10 mL of DMF, and 2 mL of ethylendiamine solution in DMF (1.89
× 10^{-5} \text{ mol/mL} \) was added. An aliquot of the solution was removed after the completion of the polymerization (2 days) for characterization of the poly(l-alanine) block. Then, 0.05 g of Tyr(tBu)-NCA was then added and was left to react until all monomer was completely consumed. After complete consumption of Tyr(tBu)-NCA (2 days), the final triblock copolymer was precipitated in diethyl ether, filtered, and dried to constant weight. For the deprotection of tBu groups, the same procedure as above was followed. Finally, poly(L-tyrosine)–poly(L-alanine)–poly(L-tyrosine) was produced as a white powder.

The reactions used are given in Scheme 2. The final copolypeptide was characterized only by \(^1\)H NMR spectroscopy (Figure 3), since it was not soluble in DMF.

**Methods.** **Size Exclusion Chromatography.** Size-exclusion chromatography (SEC) was used to determine the \(M_n\) and \(M_w/M_n\) values. The analysis was performed using a system that was composed of a Waters 600 high pressure liquid chromatographic pump, Waters Ultrastyragel columns (HR-2, HR-4, HR-5E, and HR-6E), a Waters 410 differential refractometer detector and a Precision PD 2020 two angle (15°, 90°) light scattering detector at 60 °C. A 0.1N LiBr DMF solution was used as an eluent at a rate of 1 mL/min (SEC-TALLS).

**NMR and FTIR Spectroscopy.** \(^1\)H NMR spectroscopy (300 MHz) was performed using a Varian Unity Plus 300/54 spectrometer. The spectra of the polymers were measured either in deuterated DMSO (NCA and poly(l-tyrosine)–PEO–poly(l-tyrosine)) or in CF\(^3\)COOD (the Tyr\(_7\)-PAla-Tyr\(_7\) copolypeptide). FTIR measurements were performed with a PerkinElmer Spectrum One instrument, in KBr pellets at room temperature, spectra being recorded over the range 450–4000 cm\(^{-1}\).

**Sample Preparation.** The solubility of samples dissolved into water was found to be poor but pH adjustment to 12 (above the pK\(_a\) of tyrosine\(^{37}\)) using an NaOH solution led to satisfactory solubility, therefore this was the pH value selected for all measurements for samples Tyr\(_5\)-PEO\(_6\)k-Tyr\(_5\) and Tyr\(_3\)-PAla-Tyr\(_3\). Since sample Tyr\(_5\)-PEO\(_6\)k-Tyr\(_5\) was found to show good solubility in water, it was also studied at its native pH, pH 5.69.

**Circular Dichroism.** CD spectra were recorded using a Chirascan spectropolarimeter (Applied Photophysics, UK) in the wavelength range 190–320 nm. The samples, 0.5% in water raised to pH 12 or native pH for Tyr\(_5\)-PEO\(_6\)k-Tyr\(_5\), were placed in a coverslip cuvette (0.1 mm thick), with spectra presented with absorbance A < 2 at any measured point with a 0.5 nm step, 1 nm bandwidth, and 1 s collection time per step. Samples were acclimatized at each increasing 10 °C temperature point for 5 min, and 30 min for the 60–20 °C change, before measurements were taken. The CD signal from the background was subtracted from the CD spectra of the sample solutions.

**FTIR.** For the secondary structure studies, spectra were recorded using a Thermo Scientific Nicolet IIS and a Nexus-FTIR spectrometer, both equipped with a DTGS detector. A 40 μL drop of the sample (in D\(_2\)O) was sandwiched between two CaF\(_2\) plate windows, with a 0.006 mm thick Mylar spacer) in a Specac GS20500 sample cell holder. Cells were heated using a Specac 4000 series high
stability heating controller, and a Specac electrical heating jacket. The sample was acclimatized at each temperature point for 10 min before measurements were taken. Spectra were scanned 128 times over the range of 900–4000 cm⁻¹.

**X-ray Diffraction.** X-ray diffraction was performed on a peptide stalk prepared by drawing a fiber of 1% conjugate solution between the ends of wax-coated capillaries, after separation and drying a stalk was left on the end of one capillary. For some samples, the capillary was mounted vertically onto the four axis goniometer of a RAXIS IV++ X-ray diffractometer (Rigaku) equipped with a rotating anode generator (CuKα, radiation, wavelength λ = 1.54 Å). The XRD data was collected using a Saturn 992 CCD camera. The sample-to-detector distance was 50 mm. Other measurements were performed on stalks mounted (vertically) onto the goniometer of an Oxford Instruments Gemini X-ray diffractometer, equipped with a Sapphire 3 CCD detector. The sample to detector distance was 45 mm.

**Small-Angle X-ray Scattering.** For the solutions, experiments were performed on ID02 at the ESRF, Grenoble France. Samples were flowed through a quartz capillary. After the sample was injected in the capillary and reached the X-ray beam, the flow was stopped during the SAXS data acquisition. The sample–detector distance was 2 m (data were also obtained at 10 m). Data processing (background subtraction, radial averaging) was performed using the dedicated beamline software or SAXSUtilities (http://www.sztucki.de/SAXSutilities/).

For the solid state Tyr₅-PEO₆k-Tyr₅ sample, measurements were performed on BM26B (DUBBLE) at the ESRF (Grenoble, France). Time-resolved DSC, SAXS, and WAXS measurements were performed. The sample-to-SAXS detector distance was ca. 1476 mm using a wavelength of 1.033 Å. A Dectris-Pilatus 1 M detector with a resolution of 981 × 1043 pixels and a pixel size of 172 μm². The wavenumber q = (4π/λ)sinθ scale for the WAXS scale was done using alpha aluminum. The samples were introduced in modified DSC pans pierced in order to produce X-ray windows which were placed in a DSC 600 heating stage from Linkam. The modified DSC pan windows are made of mica, and the capsules were sealed, avoiding sample loss. The sample was heated from −20 to 80 °C to −20 to 80 °C at 5 °C/min, SAXS/WAXS measurements being taken every 30 s.

**Rheology.** Dynamic shear moduli were measured using a controlled stress AR-2000 rheometer from TA Instruments. A cone-and-plate geometry (20 mm diameter, 1° angle) was used for a sample of Tyr₅-PEO₆k-Tyr₅ (20 wt % in water). Stress sweeps were performed first to determine the linear viscoelastic regime. Frequency sweeps were performed with a stress of 4.75 Pa. The angular frequency range was 0.1–622.8 rad s⁻¹, although data are only presented up to 50 rad s⁻¹ due to device inertia at high frequency.

The viscosity of the sample was measured performing varying stress experiments in the range 0.0056 to 16880 s⁻¹.

**Polarized Optical Microscopy.** Images were obtained with an Olympus BX41 polarized microscope by placing the sample between crossed polarizers. Films were prepared by drying 40 μL of 1 wt % peptide. Dry films were placed between a glass slide and a coverslip before capturing the images with a Canon G2 digital camera.

**RESULTS**

The molecular characteristics of the polymers synthesized are shown in Table 1. In the case of the two PEO-containing polymers, the molecular weights were obtained by SEC-TALLS. The composition of the final copolymers obtained was in excellent agreement with the stoichiometric molecular weights as well as the composition obtained by ¹H NMR spectroscopy. The molecular weight of the PTyr₅-b-PTyr₅ copolymer was obtained from the ratio of integral of the NMR signal from the protons of the initiator and those of the poly(L-tyrosine) and poly(L-alanine). Due to the low molecular weight of the blocks, it was possible to obtain the accurate...
molecular weights, which were in good agreement with the stoichiometric ones. This reveals the copolymer contains three tyrosine residues at each terminus, so the polymer is henceforth named Tyr$_3$-PAla-Tyr$_3$.

So far, the benzyl protecting group has been used for the synthesis of poly(L-tyrosine) through the NCA approach.$^{40-44}$ The benzyl groups are usually deprotected utilizing HBr in acetic acid. These conditions are so acidic that almost all protecting groups that are treated under acidic conditions are cleaved, and therefore, these conditions cannot be utilized selectively if an amphiphilic copolypeptide bearing a protected as well as a free residue is to be synthesized. It is well known that the tert-butyl group can be cleaved under mild conditions in the presence of TFA.$^{36}$ Although during the synthesis of the NCA of tyrosine HCl is produced, we found that it did not influence the tert-butyl protective group. We found that TFA was an effective reagent to quantitatively remove the tert-protecting group of the hydroxyl group of tyrosine, while, under these conditions, several other protective groups remain intact. Therefore, the novel monomer that we present in this work can be used effectively for the selective deprotection of tyrosine (Supporting Information, Scheme S1).

The secondary structure of the peptide components of the conjugates was examined using a combination of circular dichroism (CD) and FTIR spectroscopy. Figure 4 shows CD spectra. The temperature dependence of the CD data was examined in the range 20°–60°C, and no significant changes in the CD spectra were noted for either Tyr$_5$-PEO$_{2k}$-Tyr$_5$ or Tyr$_5$-PEO$_{6k}$-Tyr$_5$ (Figure 4a,b). For Tyr$_5$-PEO$_{6k}$-Tyr$_5$ at native pH (pH 5.69), the spectra (Figure 4a) show a minimum at 218 nm, which is a signature of $\beta$-sheet structure. The maximum at 228 nm is due to the absorbance of the tyrosine residues, similar features having been observed for a Tyr$_2$-PEO$_{1.5k}$-Tyr$_2$ conjugate, YYKLVF-PEG conjugates with PEG$_{1k}$ or PEG$_{3k}$, as well as dityrosine containing peptides.$^{46}$ The spectra for Tyr$_5$-PEO$_{6k}$-Tyr$_5$ at pH 12 (Figure 4b) are qualitatively similar in shape to those at native pH, but with reduced molar ellipticity values pointing to reduced secondary structure content. For Tyr$_5$-PEO$_{2k}$-Tyr$_5$ the spectra shown in Figure 4c do not extend as low in wavenumber as those for Tyr$_5$-PEO$_{6k}$-Tyr$_5$, due to observed absorption effects but the shape resembles that for the latter conjugate, but with a red shift such that the minimum is at 220 nm and the maximum due to the tyrosine aromatic units is at 244 nm. The spectra for Tyr$_3$-PAla-Tyr$_3$ in Figure 4d show a minimum at 226 nm, which can also be assigned to a red-shifted $\beta$-sheet structure.$^{47}$ There is no maximum at higher wavelength due to the tyrosine units, which is presumably because the contribution from these groups is overwhelmed by the poly(L-alanine) midblock with.

| polymer         | $M_n(\text{PEO}) \times 10^3$ | $M_n(\text{PAla}) \times 10^3$ | $M_n(\text{Tyr}) \times 10^3$ | $\text{Tyr units}$ |
|-----------------|-------------------------------|-------------------------------|-------------------------------|-------------------|
| 1 Tyr$_5$-PEO$_{2k}$-Tyr$_5$ | 2.0                          | -                             | 1.70                          | 10$^3$            |
| 2 Tyr$_5$-PEO$_{6k}$-Tyr$_5$ | 5.9                          | -                             | 1.75                          | 11$^+$            |
| 3 Tyr$_3$-PAla-Tyr$_3$      | -                             | 5.7 x 10$^3$                   | 1.00                          | 6$^b$             |

$^a$Obtained by SEC-TALLS. $^b$Obtained by $^1$H NMR. $^c$Obtained by SEC-TALLS and confirmed with $^1$H NMR.

![Figure 4](image_url)
approximately 64 alanine repeats. There is a reduction in β-sheet content on heating leading to a conformational state retained on cooling as the sample approaches its equilibrium state. There is no evidence for helical structure of PAla in the CD spectra, which is consistent with the analysis of the XRD data (vide infra) from this sample.

The FTIR spectra measured for all three polymers presented in Figure 5 show features in the amide II’ region (NH deformation) around 1500 cm⁻¹ and, most informatively, in the amide I’ region which provides information on peptide secondary structure via the C=O deformation modes. The spectrum for Tyr5-PAla-Tyr3 shows a peak at 1623 cm⁻¹, consistent with the β-sheet structure of polyalanine-rich peptides (previously observed at 1626 cm⁻¹ for an arginine-capped alanine oligomer), with a β-sheet assignment also for the 1635 cm⁻¹ peak for Tyr5-PEO2k-Tyr5. For Tyr5-PEO6k-Tyr5 there are no defined peaks in the amide I’ region (for samples at either native pH or at pH 12), which prohibits assignment of secondary structure. It is believed that this is due to the absorbance of the longer PEO chain in this compound overwhelming the signal from the peptide end groups. For all samples, the observed peak near 3400 cm⁻¹ is consistent with −OH stretch deformations of hydrogen-bonded carbonyl groups in PEO or poly(−alanine) and the peak at 2930 cm⁻¹ is assigned to PEO −CH₂ or poly(−alanine) backbone −CH₂ deformation modes.

Fiber X-ray diffraction on dried stalks was used to investigate the molecular ordering. The fiber X-ray diffraction patterns from all samples were found to be isotropic and were reduced to one-dimensional form. The resulting intensity profiles are shown in Figure 6. These show peaks at 7.49, 5.27, 4.36 and 3.71 Å for Tyr5-PAla-Tyr3. These are very similar to those we have reported for oligo(alanine) peptides and are assigned to the unusually tightly packed β-sheet structure of alanine-rich peptides, with a β-strand spacing of 4.36 Å and an intersheet spacing of 5.27 Å. The XRD data in Figure 6 indicate that there is also β-sheet content in Tyr5-PEO2k-Tyr3, due to the presence of the 4.69 Å reflection, along with some others that are assigned to semicrystalline PEO. That PEO is semicrystalline is shown by the presence of a broad peak due to amorphous polymer, upon which the crystal peaks are superposed. The expected d-spacings for the strong reflections (indicated Miller indices) from crystalline PEO are: d(101, s) = 7.89 Å, d(101, s) = 5.11 Å, d(120, vs) = 4.63 Å, d(202, s) = 3.94 Å, d(112) = 3.86 Å, d(031, vs) = 3.81 Å, and d(322/212, vs) = 3.79 Å. The data for Tyr5-PEO2k-Tyr3 are consistent with a very high degree of PEO crystallinity, showing strong peaks at 4.69 and 3.83 Å. In studies of PEO−peptides comprising KLVFF-based peptides and PEO of different molar mass, we have found that PEO crystallization is modulated by the fibril-forming propensity of the peptide, for instance, in conjugates with a fixed PEO molar mass (PEO3k), crystallization was observed for the weak fibrilizing peptide KLVFF, but not for the strong fibrilizer FFKLWVF. These studies, together with an investigation that compared the structures of FFKLWVF−PEOs with different PEO molar mass (1k, 2k, and 10k), suggest that PEO2k may be close to the molar mass threshold to exhibit some crystallinity. This is consistent with the data in Figure 3 for Tyr5-PEO2k-Tyr5, which shows some PEO crystallinity (but to a low extent) along with β-sheet structure from the Tyr5 end groups. On the other hand, the peaks in the X-ray intensity profiles for Tyr5-PEO6k-Tyr5 in Figure 3 are not consistent with β-sheet order (there is no peak that can be assigned to the β-sheet stacking distance) and instead can be assigned to peaks from highly crystalline PEO. In fact, there are shifts from the expected peak positions listed above, which is assigned to the influence of the Tyr5 end groups on the PEO unit cell dimensions. A detailed examination of this interesting effect whereby PEO crystallization is modified by the presence of small peptide end groups is outside the scope of the present paper. The findings concerning PEO crystallization for the samples containing PEO6k are consistent with expectations based on the PEO molar mass, and with images of dried films of the samples to be discussed shortly. The same XRD pattern is obtained for stalks dried from Tyr5-PEO6k-Tyr3 at native pH or at pH 12 (data not shown).

Cryo-TEM was used to image self-assembled nanostructure. Figure 7 shows representative images. For Tyr5-PEO6k-Tyr3, the presence of short, very straight fibrils approximately 5 nm in diameter extending to a few hundred nm in length is clear, these coexisting with shorter (and nearly spherical) nanostructures of similar diameter. For Tyr5-PEO2k-Tyr3, cryo-TEM reveals clusters of short fibrils. SAXS data (Figure 8) clearly supports the presence of fibrillar structures. A similar structure of local clusters of fine fibrils/filaments was also observed for Tyr5-PAla-Tyr3. These observations are consistent with the presence of β-sheet structures (which are typically present in extended fibrils) for these samples, as revealed by spectroscopic methods, as discussed above. The difference in appearance of the fibrils of Tyr5-PEO6k-Tyr3 is striking as they are extremely straight. Similar straight fibrils have previously been observed for Phe₄PEOSk and suggest that such short straight fibrils are
favored when the PEO chain length is in the range PEO5k–PEO6k, whereas more flexible fibrils, also termed "wormlike micelles", are observed for diblock conjugates of PEO (molar mass 0.5–1.8 kg mol$^{-1}$) to Phe4.59.

Small-angle X-ray scattering measurements were performed to provide further information on the dimensions of the cylindrical fibrils. The measured data along with form factor fits are shown in Figure 8. The data were fitted to a cylindrical core–shell form factor since there is a noticeable change in slope of the intensity profiles near $q = 0.3$ Å$^{-1}$. The fit parameters are listed in Table 2. The core radius for both Tyr$_5$-PEO2k-Tyr$_5$ and Tyr$_5$-PEO6k-Tyr$_5$ is $R = 1.03$ nm and $R = 2.28$ nm respectively, which is consistent with the core comprising Tyr$_5$ plus some of the inner part of the PEO chains, since the estimated length of a Tyr$_5$ sequence in a $\beta$-strand is $5 \times 0.32$ nm = 1.6 nm only (using the 3.2 Å translation per residue expected for a parallel $\beta$-sheet structure$^{37}$). This radius is consistent with that estimated from the cryo-TEM images for Tyr$_5$-PEO6k-Tyr$_5$ (Figure 7).

As expected, the shell radius is lower for Tyr$_5$-PEO2k-Tyr$_5$ compared to Tyr$_5$-PEO6k-Tyr$_5$ ($\Delta R = 1.92$ vs 2.20 nm). The estimated radius of gyration of PEO6k is $R_g = 2.6$ nm and of PEO2k is $R_g = 1.5$ nm (based on a segment length $b = 5.6$ Å$^{38-40}$ Hammouda and Ho report $b = 5.8$ Å$^{63}$). The values of $\Delta R$ from the core–shell cylinder form factor fit are close to these values, but are not identical due to the fact that the core radius $R$ includes some of the inner PEO chains, as mentioned above, and there may also be some expansion of the PEO2k chains, in particular in the fibril corona.

We also examined possible gel formation by inspection of samples, followed by shear rheometry of concentrated liquids. It was observed that neither Tyr$_5$-PEO6k-Tyr$_5$ (native pH or pH 12) nor Tyr$_5$-PEO2k-Tyr$_5$ form a gel up to 20 wt % (sample Tyr$_3$-PAla-Tyr$_3$ is not soluble at high concentration so gel formation is precluded). Figure 9 shows measurements of

| sample | $N$  | $\sigma$/nm | $R$/nm | $\Delta R$/nm | $L$/nm | $\eta$(core) | $\eta$(shell) | $\eta$(solv) | BG |
|--------|-----|-------------|--------|---------------|-------|-------------|---------------|--------------|-----|
| PEO2k  | 0.06| 2.48        | 1.03   | 1.92          | 50    | $1.6 \times 10^{-5}$ | $-7.1 \times 10^{-5}$ | $2.4 \times 10^{-4}$ | 1.3 $\times 10^{-4}$ |
| PEO6k  | 0.015| 2.66        | 2.28   | 2.20          | 50    | $4.1 \times 10^{-4}$ | $-1.7 \times 10^{-3}$ | $1.0 \times 10^{-4}$ | 3 $\times 10^{-4}$ |

"The fit was performed using the long cylindrical shell form factor using software SASfit.$^{72}$ $N =$ scaling factor. $\sigma =$ Gaussian polydispersity parameter. $R =$ cylinder radius. $\Delta R =$ shell thickness. $L =$ Length of cylinder (fixed). $\eta$(core) = scattering length density of core. $\eta$(shell) = scattering length density of shell. $\eta$(solv) = scattering length density of solvent. BG = background (fixed).
dynamic shear modulus for a viscous 20 wt % solution of Tyr5-PEO6k-Tyr5. Stress sweeps (Figure 9a) showed approximately linear viscoelastic behavior up to a stress $\sigma = 10$ Pa. Frequency sweeps were then performed, and the results at two different stress values are shown in Figure 9b. These reveal frequency-dependent moduli $G'' > G'$ (and low values of both moduli) consistent with a predominantly fluid-like response. This conclusion is supported by the observed terminal scalings of the moduli $G' \sim \omega^2$ and $G'' \sim \omega^{1.64}$ The lack of observed gel formation by Tyr5-PEO6k-Tyr5 even with conjugate concentration up to 20 wt % in aqueous solution may be contrasted with the prior observation of hydrogel formation (at 6 wt %) by NH$_2$-Tyr2-PEO1.5k-Tyr2-Fmoc; however, the Fmoc (N-fluorenyl-9-methoxycarbonyl) unit has a substantial influence on the hydrophobicity of this conjugate and hence on its ability to form a hydrogel. Also hydrogel formation has been observed for a 10 wt % solution of PAla-PGlu-PAla (PGlu: poly(L-glutamic acid)) in buffer solution although this has a charged midblock and different block sequence to our conjugates.

Figure 9c shows the shear rate dependence of the viscosity for the Tyr5-PEO6k-Tyr5 conjugate. This shows a Newtonian regime up to approximately $\dot{\gamma} = 0.1$ s$^{-1}$, followed by a shear-thinning regime with an initial power-law Ostwald-de-Waele behavior, $\eta \sim \dot{\gamma}^{0.5}$. Similar shear-thinning behavior is observed for conventional associative polymers and solutions of wormlike micelles.

As well as examining the self-assembly of the conjugates in aqueous solution, the behavior of the Tyr5-PEO6k-Tyr5 sample that exhibits PEO6k crystallization on drying was investigated in more detail in the solid state. Figure 10 shows results from synchrotron SAXS/WAXS samples on a dry sample. SAXS and WAXS (Figure 10a,b) show that the first melting of PEO occurs at 52 °C, which is a typical value for the melting temperature $T_m$ for PEO with a molar mass of 6 kg mol$^{-1}$. On cooling, crystallization is observed to start at 50 °C and the second melting at 52 °C. These results indicate a low degree of hysteresis in the melting/crystallization behavior. After the second crystallization, the SAXS data (Figure 10c) shows a broad peak centered at $q^* = 0.5$ nm$^{-1}$ consistent with the presence of crystalline lamellae but there was no evidence for microphase separation in the melt state. The crystalline lamellar period is 12.6 nm, which indicates significant extension of the PEO chains compared to the conformation in aqueous solution discussed above. Actually, the chain length of PEO in Tyr5-PEO6k-Tyr5 can be estimated as $l = 0.095z_{EO}$ where 0.095 Å is the length per atom along the PEO helix and $z_{EO}$ is the number of atoms (here 136 × 3, where $N = 136$ is the degree of polymerization), yielding $l = 38.8$ nm. Comparison of this to the measured value of the crystal period points to chain folding of the PEO. Figure 10d shows the WAXS profile in the crystalline state for comparison to the fiber XRD data shown in Figure 6.

These findings concerning PEO crystallization were supported by polarized optical microscopy (POM) images from dried films, shown in Figure 11. For Tyr5-PEO6k-Tyr5, under native pH conditions there is clear evidence for the presence of crystalline spherulites for the dried sample whereas the sample dried from pH 12 exhibits much smaller and less extensive crystallites. Partial crystallinity with poorly developed spherulites was observed for Tyr5-PEO2k-Tyr5 at pH 12, consistent with the WAXS data.

**SUMMARY AND CONCLUSIONS**

All three oligo(tyrosine)—polymer—oligo(tyrosine) conjugates self-assemble into $\beta$-sheet fibrillar structures as shown by a combination of spectroscopic, scattering and microscopy methods. The amide I’ FTIR spectrum for Tyr5-PEO6k-Tyr5 did not show the typical peak for $\beta$-sheet structure (probably due to the dominant contribution from unordered PEO6k), but
the CD data confirmed the assignment of this structure. The CD spectra for this sample and the Tyr₅-PEO₂k-Tyr₅ sample contain peaks also from the Tyr₅ end groups. The SAXS and cryo-TEM data provide clear evidence for the formation of fibrillar structures. A schematic of the proposed fibrillar structure is shown in Figure 12. One remarkable result from

![Figure 11. Polarized optical microscopy images of dried samples. (a) Tyr₅-PEO₆k-Tyr₅, native pH, (b) Tyr₅-PEO₆k-Tyr₅, pH 12, (c) Tyr₅-PEO₂k-Tyr₅, pH 12.](image)

![Figure 12. Schematic self-assembled structure showing tyrosine fibrillar network linked by PEO or PAla chains with bridging and looping conformations.](image)

our study is that even polymers with end-groups as short as three tyrosine residues can self-assemble to form tyrosine fibrils. The self-assembly of the Tyr₅-PAla-Tyr₅ conjugate is significantly influenced by the PAla secondary structure and as discussed above oligo- or poly-(l-alanine) adopts a very interesting tightly packed β-sheet structure with a substantially smaller intersheet spacing and interstrand spacing than formed by peptides containing larger side groups. For Tyr₅-PEO₆k-Tyr₅ the presence of the longer PEO chain compared to that in Tyr₅-PEO₂k-Tyr₅ enables dissolution in water at native pH rather than above the tyrosine pKₐ (pKₐ = 10.1⁵⁷); however, interestingly, this has little difference on the observed secondary structure formation.

The fibril morphology is distinct comparing the two PEO-based conjugates; whereas irregular clusters of short flexible fibers are observed for Tyr₅-PEO₂k-Tyr₅, the conjugate with PEO₆k self-assembles to form a population of very straight and well-defined (narrow dispersity in width) fibrils. This difference in morphology is consistent with prior reports on the formation of straight fibril-like structures in “diblock” polymer/peptide conjugates with PEO molar mass around 5k.⁵⁸,⁵⁹ In the present conjugates, which differ in having a telechelic “triblock” architecture, the longer PEO chain will have greater conformational flexibility, which may have several effects: (i) The greater PEO chain flexibility may enable tighter packing of the Tyr end-groups in the fibril reducing the tendency for fibril curvature; (ii) It may permit more looped conformations, which could act to stabilize short straight fibrils via coverage of fibril sides (reducing fibril width dispersity) or ends (preventing very long fibrils from assembling).

The presence of PEO means that PEO crystallization can occur upon drying. Indeed, the XRD pattern for Tyr₅-PEO₆k-Tyr₅ is dominated by strong peaks from crystalline PEO, whereas Tyr₅-PEO₂k-Tyr₅ contains a substantial amorphous fraction in the solid state. The SAXS pattern for Tyr₅-PEO₆k-Tyr₅ in the solid state contains a peak due to the formation of semicrystalline PEO lamellae. These findings are supported by observation of the spherulites formed in dried films of these samples, which are extensive and sample-spanning for Tyr₅-PEO₆k-Tyr₅ but not Tyr₅-PEO₂k-Tyr₅.

This manuscript reports a comprehensive study of the self-assembly of these telechelic conjugates capped with short tyrosine blocks. In future work, it will be interesting to examine the bioactivity and potential cytocompatibility of these conjugates resulting from the tyrosine functionality or to investigate responsiveness, induced for example by enzymatic triggering of tyrosine phosphorylation using kinases. Tyrosine phosphorylation plays an important role in many cell signaling processes, and there is the scope to develop novel biomaterials to tune cell signaling processes based on polymer/tyrosine conjugates. In a similar vein, it will also be interesting to explore the conditions where (hydro)gelation can be induced.

**ASSOCIATED CONTENT**

Supporting Information
The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acs.biomac.6b00023.

Reaction scheme for the Tyrosine NCA, ¹H NMR spectrum of the novel monomer Tyr(tBu)-NCA, and FTIR spectra monitoring the synthesis of the Tyr(tBu)-NCA (PDF)

**AUTHOR INFORMATION**

Notes
The authors declare no competing financial interest.

**ACKNOWLEDGMENTS**

This work was supported by an STFC Futures Studentship (ST/L502480/1) that supported the work of S.K. and EPSRC Platform Grant EP/L020599/1. We are grateful to the ESRF (Grenoble, France) for the award of beamtime on ID02 (ref SC-4063) and to Theyencheri Narayanan for assistance. Use of facilities in the Chemical Analysis Facility at the University of Reading is gratefully acknowledged.

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