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Amyloid-Like Fibrils from a Diphenylalanine Capped with an Aromatic Fluorenyl

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ABSTRACT

The self-assembly behavior of a diphenylalanine amphiphile blocked at the C-terminus with a 9-fluorenylmethyl ester and stabilized at the N-terminus with a trifluoroacetate (TFA) anion, TFA·FF-OFm, has been examined. At low peptide concentration (0.5 mg/mL), long amyloid-like fibrils, which come from the fusion of two or more helical ribbons and/or thinner fibrils, organized in bundles or as individual entities are detected. Microbeam synchrotron radiation infrared spectroscopy have shown that TFA·FF-OFm molecules in amyloid-like fibrils arrange forming antiparallel β -sheets. Alteration of the experimental conditions to prioritize the thermodynamic contribution with respect to the kinetic one in the self-assembly process inhibits the organization of amyloid-like structures in favor of the formation of conventional fibrous structures. On the basis of experimental observations, a structural model where the individual antiparallel β -sheets are oriented in parallel has been proposed for TFA·FF-OFm amyloid-like fibrils.

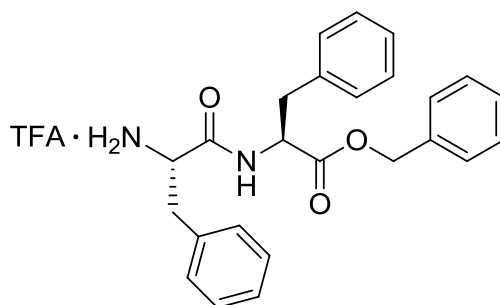
INTRODUCTION

Controlling the self-assembly of organic molecules into highly ordered nano- and microstructures is a very interesting topic in supramolecular chemistry and materials science.¹ In particular, due to the unique properties, one-dimensional morphologies, such as tubes and fibers have attracted much attention.^{2,3} Within this issue, the diphenylalanine dipeptide, L-Phe-L-Phe (FF), which is the core recognition motif of Alzheimer's A β peptides is a captivating building block that spontaneously self-assembles into nanotubes and nanofibers through well-ordered and consistent processes.⁴⁻⁷ The stability of these structures was attributed to the directionality offered by a combination of hydrogen bonding and repeated phenyl stacking interactions. Consequently, FF displays more well-defined structures than longer homo-oligomers, as for example FFF⁸ and FFFF.⁹ In addition, FF has attracted increasing interest as building block due to its biocompatibility, functional molecular recognition, and unique biological and electronic properties.¹⁰⁻¹³

Due to its inherent self-assembly capacity, the FF has given place to the development of a new class of biomaterials that is based on the addition of N- and/or C-terminal capping groups to the aromatic FF or, even, on chemical modification of the own F residues.⁸⁻¹⁷ The amphiphile Fmoc-FF, where Fmoc refers to fluorenylmethoxycarbonyl, is among the most studied capped FF-based biomaterials. This short peptide forms gels under very different experimental conditions, neighboring molecules interacting through hydrogen bonds and weak offset π - π interactions.¹⁸⁻²⁷ Similarly, Nap-FF (Nap= naphthalene) and Cbz-FF (Cbz= benzyloxycarbonyl) hydrogels showed fibrous structures made of β -sheet arrangements.²⁸ On the other hand, the Fmoc-FF-OFm peptide, capped with Fmoc and 9-fluorenylmethyl ester (OFm) at the N- and C-terminus, respectively, exhibited a great variety of polymorphic self-

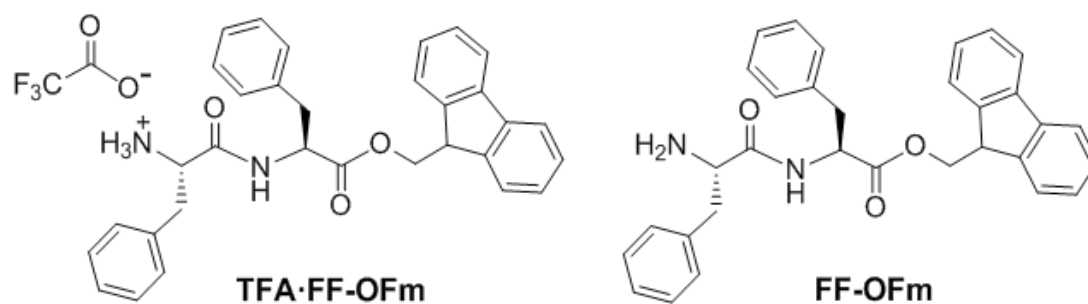
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3 assembled microstructures (*e.g.* doughnut, stacked-braids, dendritic and microtubes)
4 depending on the solvents used to promote self-assembly processes.²⁹ Advances on
5 structural trends of assembled FF-based amphiphiles have been periodically reviewed
6 along the last decade.³⁰⁻³³

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12 In a recent study, we examined the self-assembly of a FF-based amphiphile that
13 exhibited three main differences with respect to Fmoc-FF:³⁴ (1) the size and (2) the
14 position of the aromatic capping group, and (3) the stability of the free (charged)
15 terminus. In this new compound, named TFA·FF-OBzl (Scheme 1), the C-terminus was
16 capped with a benzyl ester (OBzl) group and the protonated amino group was stabilized
17 forming an ion pair with trifluoroacetate (TFA). Obviously, aromatic interactions in
18 TFA·FF-OBzl were weaker than on Fmoc-FF and Fmoc-FF-OFm, while the dominant
19 role played by intermolecular electrostatic interactions in the parent FF peptide
20 decreased considerably because of the charge shielding induced by TFA. Results
21 showed that polar aqueous environments tend to promote the growth of TFA·FF-OBzl
22 fibers, which coexisted with branched-like microstructures when milli-Q water was
23 replaced by diluted KCl aqueous solutions (50 mM). In contrast, non-polar
24 environments obtained by mixing organic solvent gave place to peptide assemblies
25 organized in plates and spherulites.³⁴



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Scheme 1. Chemical structure of dipeptide TFA·FF-OBzl.

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3 In this work we investigate the self-assembly of a new amphiphile that, apparently,
4 can be considered as an intermediate between Fmoc-FF and TFA·FF-OBzl. More
5 specifically, in this peptide the C-terminal OBzl group of TFA·FF-OBzl has been
6 replaced by OFm. Thus, the resulting peptide, named TFA·FF-OFm (Scheme 2), only
7 differs from Fmoc-FF in (1) the position and size of the aromatic capping group and (2)
8 the stabilization of the free end through the TFA anion. Self-assembly studies have been
9 conducted considering polar environments formed by 1,1,1,3,3,3-hexafluoro-2-propanol
10 (HFIP):water mixtures and different peptide concentrations, temperatures, ionic
11 strengths, and incubation times for the feeding peptide solution. Results show that the
12 TFA·FF-OFm amphiphile self-associates into amyloid-like fibrils under well-defined
13 conditions of concentration and temperature. The formation of these twisted structures
14 is kinetically driven while the conventional fibers typically adopted by FF-derivatives
15 are thermodynamically favored. Besides, the elimination of the TFA stabilizing anion to
16 produce FF-OFm (Scheme 2) results in bundles of fibers.



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46 Scheme 2. Chemical structure of dipeptide FF-OFm (with either protonated or
47 neutral amino terminus)
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50 51 52 53 **RESULTS AND DISCUSSION**

54 55 **Peptide synthesis**

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57 The compounds studied in this work, TFA·FF-OFm and FF-OFm, were synthesized
58 according to the route shown in Figure 1. Two residues of L-Phe functionalized in the
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3 amino or carboxylic acid terminus as *N-tert*-butoxycarbonyl (Boc) or 9-fluorenylmethyl
4 (Fm) ester, respectively, were coupled to give the corresponding dipeptide Boc-L-Phe-L-
5 Phe-OFm (Boc-FF-OFm). Removal of the *N*-Boc protection by treatment with
6 trifluoroacetic acid (TFA) provided the dipeptide with a protonated amino terminus as
7 the trifluoroacetate salt, TFA·FF-OFm. Further neutralization afforded the compound
8 with a free neutral amino group, FF-OFm. The ¹H and ¹³C NMR spectra of TFA·FF-
9 OFm and FF-OFm are displayed in Figures S1-S4, the FTIR spectra in Figure S5-S6,
10 and the high resolution mass spectrum of TFA·FF-OFm in Figure S7.
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23 **Self-assembly of TFA·FF-OFm: from fibers to amyloid fibrils**

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26 A stock solution (5 mg/mL) of TFA·FF-OFm in HFIP was diluted with milli-Q
27 water, which acted as a co-solvent, to a final peptide concentration of 2 mg/mL (4:6
28 HFIP:water). As displayed in Figure 2, disordered fibrous peptide assemblies were
29 obtained after slow solvent evaporation at 4 °C. These consisted of fibers bundles tens
30 to even hundreds micrometers long and variable width, which grew forming highly
31 developed 3D entangled networks. Magnified SEM micrographs show that fibers of
32 micrometric width were actually constituted by a dense packing of nanofibers with
33 width of 98±17 nm, which arranged in a *braided rope*-like configuration. Thus,
34 nanofibers twisted around each other forming very homogeneous and compact braided
35 microfibrils (width: 1.3±0.4 μm) without topographic imperfections, as is evidenced in
36 the corresponding AFM images. Importantly, high magnification SEM micrograph and
37 AFM images clearly show that nanofibers do not fuse to form the braided microfibrils
38 but they maintain the own identity.
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56 Reduction of the peptide concentration to 0.5 mg/mL by increasing the amount of
57 water in the solvent:co-solvent mixture (1:9 HFIP:water) resulted in a drastic
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3 morphological change. Thus, TFA·FF·OFm molecules aggregate into long amyloid-like
4 fibrils, which can be detected forming part of relatively dense bundles (Figure 3a) or as
5 individual entitites (Figure 3b). The width of the fibril is variable, even though it
6 typically ranges from 1.5 to 2.5 μm . Fibrils clearly come from the fusion of two or more
7 thinner helical ribbons, as is clearly displayed in Figures 3c-d (red marks). More
8 specifically, two different situations were identified in SEM micrographs: (i) two
9 strands intertwine to form a ribbon (blue arrow in Figure 3c) that, subsequently,
10 intertwines with another ribbon to form a fibril (red arrows in Figure 3c); or (ii) multiple
11 strands and/or ribbons intertwine simultaneously to form the fibril (red arrows in
12 Figures 3c-d). The fusion of strands and/or helical ribbons into microfibrils is an
13 important difference with respect to assemblies discussed above for the 2 mg/mL
14 peptide concentrations, in which nanofibers maintained their own identity and
15 individuality when they entangled into microfibers (Figure 2).

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33 Amyloid-like fibrils were never detected for TFA·FF·OBzl,³⁴ which was investigated
34 using experimental conditions identical to those employed in the present work for
35 TFA·FF·OFm. Thus, 2.0 mg/mL (4:6 HFIP:water) TFA·FF·OBzl solutions formed
36 peptide nanofibers with a diameter (ϕ) of ~ 250 nm, which aligned and packed into well-
37 defined microfibers of up to ~ 10 μm . At the same time, such microfibers organized into
38 very dense aggregates with a spike-like morphology. When the TFA·FF·OBzl
39 concentration was reduced by increasing the polarity of the HFIP:water (*i.e.* adding
40 water), the density of aggregated microfibers decreased and a very porous mesh of
41 randomly oriented (bundled) fibers coexisted with the spike-like supramolecular
42 structure. In any case, no amyloid-like fibril was detected, even when HFIP was
43 replaced by other organic solvent, as for example dimethylsulfoxide and
44 dimethylformamide.³⁴ This observation evidences the important role played by the
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3 strength of π - π stacking interactions in the nucleation and growing of amyloid fibrils
4 formed by amphiphilic peptides, which have been achieved by replacing the OBzl C-
5 terminal capping group of TFA·FF-OBzl by OFm. Thus, the very high hydrophobicity
6 and aromaticity of the latter capping group with respect to the former one promote the
7 association of the small peptide molecules into unique amyloid-like structures.
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15 Parallel and antiparallel molecular arrangements in a β -sheet structure can be
16 distinguished by FTIR spectroscopy. It is well known that intermolecular hydrogen
17 bonding interactions can be established by analyzing the amide I band, which is
18 associated to the stretching vibration of the main chain carbonyl groups. The
19 wavenumber of this band is comprised between 1600 cm^{-1} and 1700 cm^{-1} , the exact
20 position depending on the formation of intra- or inter-molecular specific interactions,
21 such as hydrogen bonds.^{35,36} The antiparallel β -sheet arrangement exhibits two Amide I
22 bands at around 1685 cm^{-1} (weak) and 1632 cm^{-1} (strong), whereas a single band at
23 around 1645 cm^{-1} is characteristic of the parallel β -sheet disposition. The random
24 molecular conformation is distinguished by a single band around 1654 cm^{-1} .
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38 The FTIR spectrum, which was recorded for a powder sample recovered from a
39 preparation of TFA·FF-OFm fibrils obtained using a peptide concentration of 0.5
40 mg/mL, is displayed in Figure 4a. Typical amide bands indicative of hydrogen bonding
41 interactions (*e.g.* amide A, amide B, amide I and amide II bands at 3310, 3064, 1671,
42 and 1554 cm^{-1} , respectively) can be detected together with free carboxylate bands at
43 1716 (C=O stretching) and 1100 cm^{-1} (CO-O stretching), respectively. Amide I band
44 appears overlapped with the C=O stretching of the carbamate unit giving rise to a strong
45 signal at 1671 cm^{-1} . These results suggest that TFA·FF-OFm adopts a β -sheet structure
46 with an antiparallel disposition of hydrogen bonded molecules.
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3 Spectra of individual crystals could be acquired using polarized synchrotron
4 radiation. Specifically, individual crystals with a specific orientation were selected to
5 record the corresponding spectra. Figure 4b shows an optical micrograph of a selected
6 region where typical spherulitic aggregates (Figure 4c) and twisted amyloid-like fibrils
7 (Figure 4d) can be detected. Analyses were performed with a minimum of 10 crystals,
8 which were selected with an orientation parallel to the plane of polarization (*i.e.*
9 horizontal disposition for a 0° polarized spectrum). Figure 4e clearly shows a strong
10 intensity for the 1674 cm⁻¹ band, suggesting that C=O bonds of both amide and
11 carbamate groups are oriented parallel to the helical axis. On the contrary, the spectrum
12 taken with a polarization of 90° shows the practical disappearance of the indicated band
13 and also a clear enhancement of a band at 1657 cm⁻¹. The latter can be related to TFA
14 carboxylate groups, which seems to be more randomly distributed. It is worth noting
15 that at 90° polarization the band at 1674 cm⁻¹ splits into two bands at 1678 cm⁻¹
16 (carbamate) and 1668 cm⁻¹ (Amide I). Spectra recorded for single crystals allow
17 demonstrating the antiparallel arrangement of molecules. To complement microscopy
18 and microbeam synchrotron radiation infrared spectroscopy, Congo red staining was
19 performed for 0.5 mg/mL TFA·FF·OFm structures. As observed by polarized
20 microscopy (Figure 3e), the birefringence associated to such staining, which is a
21 diagnosis for amyloid fibril formation,³⁷ was interfered by the birefringence of highly
22 oriented fibers.

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49 In order to obtain information about the structure of the aggregates in the first stages
50 of the assembly process, circular dichroism (CD) spectra were recorded at very diluted
51 peptide solution (0.05 mg/mL) in 1:99 HFIP:water. Figures 5a and 5b displays the CD
52 spectra in a heating run from 30 °C to 90 °C and in a cooling run from 90 °C to 10 °C,
53 respectively. As it can be seen, the spectra exhibited very small variations with the
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3 temperature. The negative bands at 194 and 209 nm suggests a mixture of random and
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5 turn structures while the positive maxima at around 220 nm indicates the presence of π -
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7 π stacking, which has been attributed to the OFm capping units. Regarding to the latter
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9 band, it is worth noting that it was found to be much less intense for FF nanotubes, in
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11 which π - π stacking interactions only involve the phenyl side groups, and completely
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13 inexistent for non-self-assembling Ala-Ala dipeptide.¹⁵ Thus, the potential for aromatic
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15 stacking interactions introduced by the OFm is expected to play a crucial role in the
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17 peptide assembly, including the formation of amyloid-like fibrils. Moreover, at very
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19 diluted concentrations (*i.e.* 0.05 mg/mL in 1:99 HFIP:water), in which amyloid-like
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21 fibrils were not detected, the π - π stacking could be even the driving force. This is
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23 consistent with the poorly defined fiber-like agglomerates, with a length of a few tens of
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25 micrometers long, heterogeneously distributed onto the surface observed after solvent
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27 drying (Figure S8).
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33 34 35 **Kinetics *versus* thermodynamic stability of TFA·FF·OFm amyloid-like fibrils**

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38 In the previous section we showed that TFA·FF·OFm molecules formed twisted
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40 amyloid-like fibrils when the peptide concentration decreased from 2.0 mg/mL, which
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42 provided well-defined microfibers, to 0.5 mg/mL. This feature suggests that the
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44 nucleation and growth of amyloid structures comes from the competency between
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46 different parameters that affect the kinetics and the thermodynamics of the self-
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48 assembly process. In order to provide a deeper understanding of the factors that favor
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50 the association of TFA·FF·OFm molecules into well-organized supramolecular
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52 amyloid-like fibrils, the structures formed after altering selected experimental
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54 conditions have been analyzed.
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3 Figures 6a-b display the supramolecular structures obtained when a 0.5 mg/mL
4 TFA·FF-OFm solution in 1:9 HFIP:water was stored at rest for 12 days at room
5 temperature, before placing the solution drops on microscope coverslips for drying at 4
6 °C. Thus, during the first period the self-assembly of TFA·FF-OFm was
7 thermodynamically controlled, while solvent evaporation imposes a kinetic control after
8 deposition onto the glass substrate. As it can be seen in Figure 6a, long plates of
9 approximately 2 μm width that, apparently, come from the hierarchical assembly of
10 nanowires were obtained. Although some of these plates resemble deformed microfibers
11 because of their dimensions, twisted fibrils were completely absent when the peptide
12 solution was incubated to promote the pre-assembly of thermodynamically favored
13 structures. This feature indicates that the twisted assembly of the small amphiphile
14 under study is kinetically driven when the medium evaporates from low concentration
15 solutions.
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33 On the other hand, stacked braid-like microstructures were observed at the edges of
34 glass coverslips (Figure 6b). Although the length and width of these braids was variable,
35 they tend to organize in very compact aggregates. The dipeptide capped with two
36 fluorenyl functionalities, Fmoc-FF-OFm, also self-assembled into compact stacked
37 braid-like microstructures from HFIP:water solutions with the peptide concentrations
38 ranging from 0.5 to 2 mg/mL.²⁹ Moreover, highly aromatic peptides with a higher
39 number of F-residues, as for example Fmoc-FFF-OFm,²⁹ Fmoc-FFFF-OFm²⁹ and
40 Fmoc-FFFF-OBzl,⁹ also exhibited very similar morphologies. These observations
41 suggest that aromatic capping groups play a crucial role in the interfacial mechanism
42 that systematically stabilizes such unique microstructure at the edges of the coverslip.
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56 In order to corroborate that TFA·FF-OFm amyloid-like fibrils are not
57 thermodynamically stable, their temporal stability was evaluated. For this purpose,
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3 microstructures like those displayed in Figure 3 were prepared as obtained in the
4 previous subsection (*i.e.* using a 0.5 mg/mL peptide 1:9 HFIP:water solution). The
5 samples were stored at rest during six weeks at 4 °C. After this time, the twist of the
6 fibrils decreased or even disappeared, as is reflected in the representative SEM
7 micrographs displayed in Figure 6c. These results indicate the thermodynamic stability
8 of TFA·FF-OFm helical ribbons and amyloid-like fibrils is lower than that of fibers,
9 even though their formation is kinetically promoted under controlled conditions.
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21 **Elimination of the stabilizing TFA anion**

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23 Disruption of the TFA·FF-OFm ion pair was achieved by neutralizing with 5%
24 aqueous NaHCO₃, which yielded FF-OFm (Figure 1). The CD spectrum at room
25 temperature (Figure 5c) shows important differences with respect to TFA·FF-OFm,
26 suggesting changes in secondary structure due to the elimination of the stabilizing
27 counteranions. Thus, the CD spectrum, which shows a positive maximum at 197 nm
28 while a negative shoulder and peak appear at 208 and 222 nm, respectively, indicates
29 the complicated secondary structure of the neutralized peptide in diluted (0.05 mg/mL)
30 1:99 HFIP:water solution. This may be not uniform β -sheet and/or α -helical.
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42 The self-assembly behavior of the resulting N-uncapped compound was examined at
43 4 °C considering both 4:6 and 1:9 HFIP:water solutions, which correspond to 2 and 0.5
44 mg/mL peptide concentrations (Figure 7a and 7b, respectively). As it can be seen,
45 solvent evaporation from the solution with the highest peptide concentration resulted in
46 assemblies similar to those described for TFA·FF-OFm (Figure 2). These consist of
47 disordered clusters of relatively long fibers and varying thicknesses. Fibers of
48 micrometric thickness are constituted by well-defined nanofibers that assemble
49 hierarchically forming very compact 3D structures (Figure 7a). The similarity with
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3 TFA·FF-OFm structures is not maintained when the peptide concentration decreases to
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5 0.5 mg/mL. While TFA · FF-OFm forms amyloid fibrils (Figure 3), FF-OFm self-
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7 assembles into disordered groups of nanofibers of thickness ranging from ~50 nm to a
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9 few hundred nanometers (Figure 7b). Each of these fibers is an individual entity rather
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11 than a hierarchical organization of thinner sub-entities, as occurred in assemblies
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13 derived from 2 mg/mL peptide solutions. Moreover, the characteristic regular twist of
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15 helical ribbons and amyloid fibrils is not detected in FF-OFm fibers.
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19 Structural analyses by infrared micro-spectroscopy were easier for FF-OFm than for
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21 TFA·FF-OFm since signals corresponding to carboxylate salts are not present. The
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23 region selected for this study corresponds to a group of fibers with dimensions of
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25 several microns long and less than 50 nm wide, as is displayed in the SEM micrographs
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27 shown in Figure 7c. More specifically, zones bearing horizontally and vertically
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29 oriented fibers, which are marked in the optical micrograph (Figure 7d), were both
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31 examined by synchrotron microbeam polarized spectroscopy. Figure 7e indicates that
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33 the spectra of zones with horizontally- and vertically oriented fibers are clearly
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35 different. Thus, a predominant band at 1674 cm^{-1} and a clear shoulder at 1663 cm^{-1} were
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37 characteristic of fibers with a horizontal orientation when observed with 0° polarized
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39 light. On the contrary, the intensity of the main band clearly diminished when observed
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41 with 90° polarized light. These observations clearly indicate a parallel arrangement of
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43 carbamate C=O groups to the crystal long axis and a sheet structure with an antiparallel
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45 disposition of molecules, as above discussed for the TFA·FF-OFm derivative (Figure
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47 4e). Note that the carboxylate band at 1657 cm^{-1} logically disappeared after the basic
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49 treatment. For the sake of completeness, it is worth noting that vertically oriented fibers
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51 observed under a polarization of 0° gave rise to a spectrum with again a low intense
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3 band at 1674 cm^{-1} , which supported the high orientation of these groups in the formed
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5 crystals.
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10 **Structural model of amyloid-like fibrils**

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12 As was shown above, TFA·FF·OFm assembles into amyloid-like fibrils under
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14 appropriated conditions (Figure 3). Moreover, the periodicities, as estimated from SEM
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16 micrographs, are similar for one fiber that for a set of different fibers (*i.e.* $11.7\pm 0.5\ \mu\text{m}$
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18 and $12.2\pm 0.7\ \mu\text{m}$), supporting that the assembly is the same in all cases. Considering
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20 that microbeam synchrotron radiation infrared spectra demonstrate an antiparallel β -
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22 sheet arrangement of the peptide molecules and that CD spectroscopy results prove
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24 interactions between Fmoc units (Figure 4e and Figures 5a-b respectively), a rough
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26 structural model has been proposed for such hierarchical assembly (Figure 8). For this
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28 purpose, the TFA·FF·OFm molecule has been divided in five zones, which are
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30 schematically displayed in Figure 8a. The two zones located at the TFA counterion
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32 correspond to the trifluoromethyl and the carboxylate, which should establish favorable
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34 intermolecular van der Waals and electrostatic interactions, respectively, while the three
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36 at the peptide molecule are the charged NH_3^+ at the N-terminus, the two Phe residues
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38 and the OFm capping group. Intermolecular interactions involving the NH_3^+ and Fmoc
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40 groups should be electrostatic and π - π stacking, respectively, whereas Phe residues are
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42 expected form both hydrogen bonds and π - π stacking. Kinetically driven molecular
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44 association processes are expected to be guided by unspecific electrostatic interactions,
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46 which are much stronger than specific hydrogen bonding and π - π stacking interactions.
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54 In order to favor attractive $\text{COO}^- \cdots \text{NH}_3^+$ inter-sheet interactions avoiding strong both
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56 $\text{NH}_3^+ \cdots \text{H}_3\text{N}^+$ and $\text{COO}^- \cdots \text{OOC}^-$ repulsions, successive antiparallel β -sheets are
57
58 proposed to be laterally assembled with a spatial displacement along the peptide-axis
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3 (Figure 8b). It is worth noting that both recuperative and progressive shearing of the
4 sheets promote favorable electrostatic interactions, discrimination between such two
5 models being not possible with available experimental data. In both cases, the inter-
6 sheet cohesive forces involve attractive electrostatic, π - π stacking and van der Waals
7 forces. On the other hand, the change along the register of each individual antiparallel
8 sheet is responsible of the twist characteristic of the amyloid-like fibrils (Figure 8c).
9 This shift in the register of individual sheets facilitates the intertwined assembly and
10 subsequent fusion of helical ribbon or strands to form micrometric fibrils (Figure 8d). In
11 spite of this, it should be remarked that the effective interaction area is always higher for
12 cylindrical fibers (Figure 8e) than for helical ribbons (Figure 8d). Therefore, the
13 attraction between helical ribbons is weak as compared with that between cylindrical
14 fibers. In addition, depending on the sequence and/or blocking groups, cylindrical fibers
15 made of short peptides are compatible with different molecular packings,^{9,14,38,39} which
16 may also enhance the interactions among associated structures. These observations are
17 consistent with the experiments discussed above about the kinetics versus the
18 thermodynamic stability of amyloid fibrils. More specifically, the stronger interactions
19 between laterally associated cylindrical fibers agree with their tendency to aggregate
20 and precipitate from the solution, whereas the association of helical ribbons is favored
21 from a thermodynamic point of view and, therefore, amyloid fibrils do not correspond
22 to the lowest free energy.

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49 It is worth noting that helical ribbons and/or amyloid-like fibrils are not formed when
50 the OFm capping group is replaced by OBzl.³⁴ Therefore, π - π stacking interactions
51 between aromatic OFm groups seems to be crucial for the formation of such structures.
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The model proposed in Figure 8, stacked OFm ladders are formed among laterally
assembled molecules in the direction perpendicular to the molecular axis and the fibril

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3 axis. In those ladders, the distance between stacked OFm groups is expected to be
4
5 defined by the protuberating Ph side groups of Phe residues, which probably are also
6
7 involved in intermolecular π - π stacking interactions with the side groups of neighboring
8
9 molecules. Obviously, for the same molecular conformation the role of such stacked
10
11 ladders would be much weaker for TFA·FF-OBzl, especially considering that the
12
13 distance among stacked OBzl groups should be the same that in the proposed model.
14
15 Indeed, TFA·FF-OBzl was found to crystallize in a folded conformation, which allowed
16
17 a unique structure with a segregated distribution of hydrophilic and hydrophobic
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19 regions, rather than in a conventional β -strand conformation within a β -sheet. This
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21 conformation and supramolecular organization was incompatible with the formation of
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23 amyloid-like fibrils.
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31 CONCLUSIONS

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33 Herein, we show the influence that increasing aromaticity has on the self-assembly
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35 behavior of amphiphilic dipeptides of sequence TFA-FF-X. In particular, we study the
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37 dipeptide capped with X= OFm, comparing the results with those reported for X= OBzl
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39 using identical experimental conditions.³⁴ While TFA·FF-OBzl molecules adopted a
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41 segregated organization in which the hydrophilic core of each molecule (carboxylate
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43 from TFA, amide and ester groups) was surrounded by the hydrophobic core of
44
45 neighboring molecules, which explained the observed self-assembled structures,
46
47 TFA·FF-OFm adopts an antiparallel β -sheet structure that allows the formation of
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49 amyloid-like fibrils at appropriate peptide concentration (0.5 mg/mL). However, this
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51 structure, which results from a kinetically driven self-aggregation process, is
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53 thermodynamically less stable than non-twisted regular fibers. As it is usual for Phe-
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55 containing peptides, the self-assembly of TFA·FF-OFm displayed concentration-
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3 dependent polymorphism. This observation indicates that π - π stacking interaction play a
4
5 decisive role in the self-assembly process, including the amyloid fibril formation. The
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7 self-assembly of TFA·FF-OFm into amyloid fibrils opens new doors for the design of
8
9 FF-based materials.
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15 **SUPPORTING INFORMATION**

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17 The Supporting Information is available free of charge on the ACS Publications
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19 website.
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21 Experimental methods, ^1H -, ^{13}C - and ^{19}F -NMR spectra, FTIR spectra, high resolution
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23 mass spectra and additional SEM micrographs.
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41 Generalitat de Catalunya.
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CAPTIONS TO FIGURES

Figure 1. Synthesis of TFA·FF-OFm and FF-OFm (F, L-phenylalanine; Fm, 9-fluorenylmethyl; TFA, trifluoroacetate). Reagents and conditions: (a) EDC, HOBT, DIPEA, CH₂Cl₂, 0 °C 30 min, rt, 24 h; (b) TFA, CH₂Cl₂, rt, 1 h; (c) 5% aqueous NaHCO₃.

Figure 2. Representative SEM micrographs with (a) low and (b) high magnifications and AFM (c) height and (d) 3D topographic images (40×40 μm², 20×20 μm² and 5×5 μm² in left, center and right, respectively) of assemblies obtained from 2 mg/mL TFA·FF-OFm solutions in 4:6 HFIP:water at 4 °C.

Figure 3. Representative SEM micrographs of helical ribbons and amyloid fibrils obtained from 0.5 mg/mL TFA·FF-OFm solutions in 1:9 HFIP:water at 4 °C: (a) dense bundle of amyloid fibrils; (b) individual amyloid fibril; (c) transition from two strands to an intertwined helical ribbon (blue arrow) and from two helical ribbons to a fused fibril (red arrows); and (d) transition from multiple strands and/or helical ribbons to a fused fibril (red arrows). (e) Polarized optical microscope images of Congo red stained structure.

Figure 4. (a) FTIR spectrum of TFA·FF-OFm with labeling of main signals. (b) Micrograph showing selected crystals obtained from 0.5 mg/mL TFA·FF-OFm solutions in 1:9 HFIP:water at 4 °C with a horizontal orientation. (c, d) SEM micrographs showing (c) spherulitic aggregates and (d) twisted amyloid-like fibrils. (e) Microbeam synchrotron radiation infrared (SRIR) spectrum taken with polarization at 0° and 90°.

Figure 5. CD spectra of TFA·FF-OFm at temperatures ranging (a) from 30 °C to 90 °C (heating run) and (b) from 90 °C to 10 °C (cooling run). (c) CD spectra of FF-OFm at 30 °C. In all cases a 0.05 mg/mL peptide solution in 1:99 HFIP:water was considered.

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3 **Figure 6.** Representative (a,d) SEM micrographs and both (b) height and (c) 3D
4 topographic AFM images ($50 \times 50 \mu\text{m}^2$) of the structures obtained when a 0.5 mg/mL
5 TFA·FF-OFm solution in 1:9 HFIP:water was stored at rest for 12 days at room
6 temperature, before placing on microscope coverslips until dryness at 4 °C: (a-c)
7 nanowires of micrometric length and plates; and (d) stacked braid-like microstructures
8 located at the coverslips edges. (e) SEM of the structures obtained after store at rest for
9 6 weeks the amyloid-like fibrils formed from 0.5 mg/mL TFA·FF-OFm solutions in 1:9
10 HFIP:water at 4 °C (Figure 3).
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21 **Figure 7.** Representative SEM micrographs of structures obtained from: (a) 2
22 mg/mL and (b) 0.5 mg/mL FF-OFm solutions (4:6 and 1:9 HFIP:water, respectively) at
23 4 °C. (c) SEM micrographs showing the region used for synchrotron-based infrared
24 micro-spectroscopy measurements. (d) Optical micrograph displaying horizontally and
25 vertically oriented FF-OFm fibers (blue and red, respectively) selected for
26 measurements. (e) Microbeam synchrotron radiation infrared (SRIR) spectra of FF-
27 OFm fibers with horizontal and vertical orientations, which were taken with
28 polarizations of 0° and 90° (only for the horizontal orientation).
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40 **Figure 8.** Structural model of TFA·FF-OFm amyloid-like fibrils: (a) Scheme
41 displaying the five fragments used to represent TFA·FF-OFm molecules, which have
42 been selected as a function of the chemical identity and capacity to form intermolecular
43 interactions; (b) Recuperative (left) and progressive (right) models for the lateral
44 assembly of antiparallel β -sheets; (c) Model for the shearing along each antiparallel β -
45 sheet, which explains the twisting in the resulting structure; (d) Model for the fibril
46 explaining the fusion of twisted strands or helical ribbons. (e) Lateral assembly between
47 cylindrical fibers.
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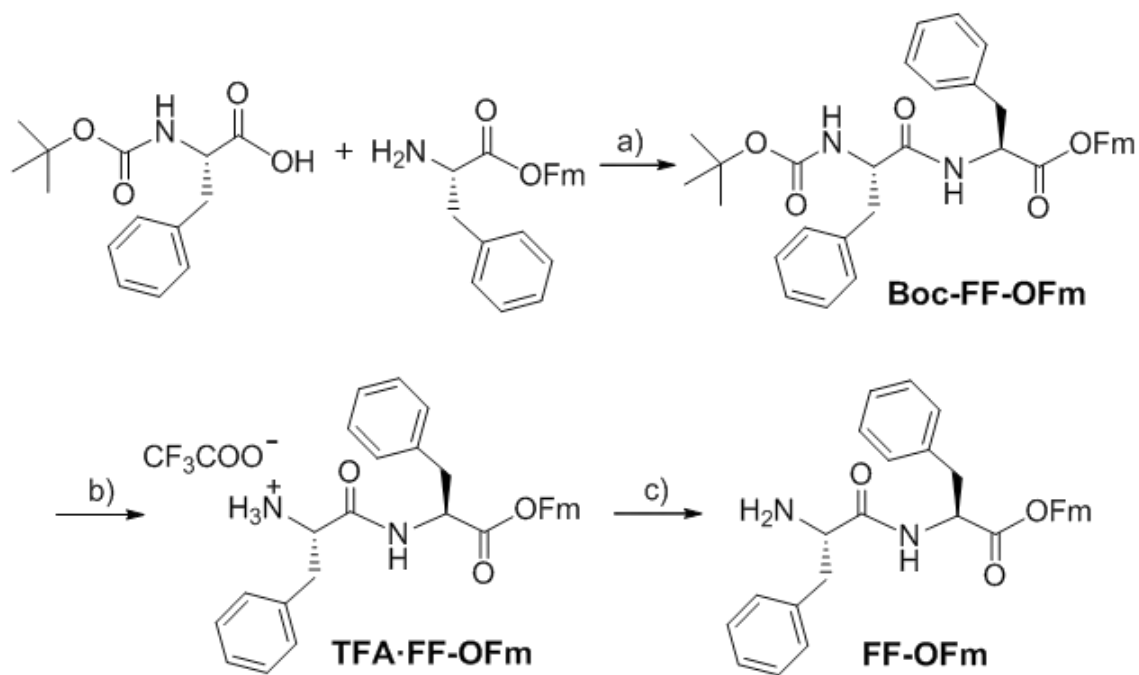


Figure 1

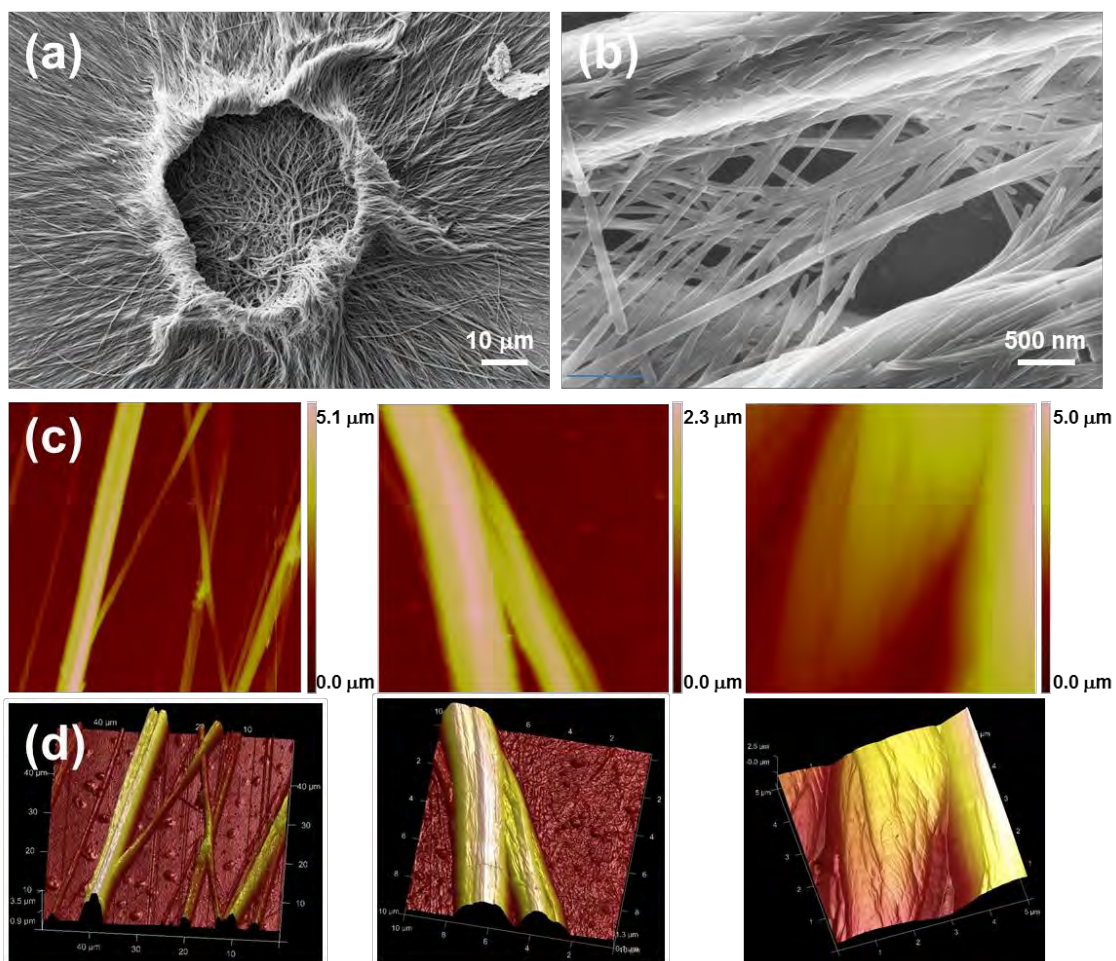


Figure 2

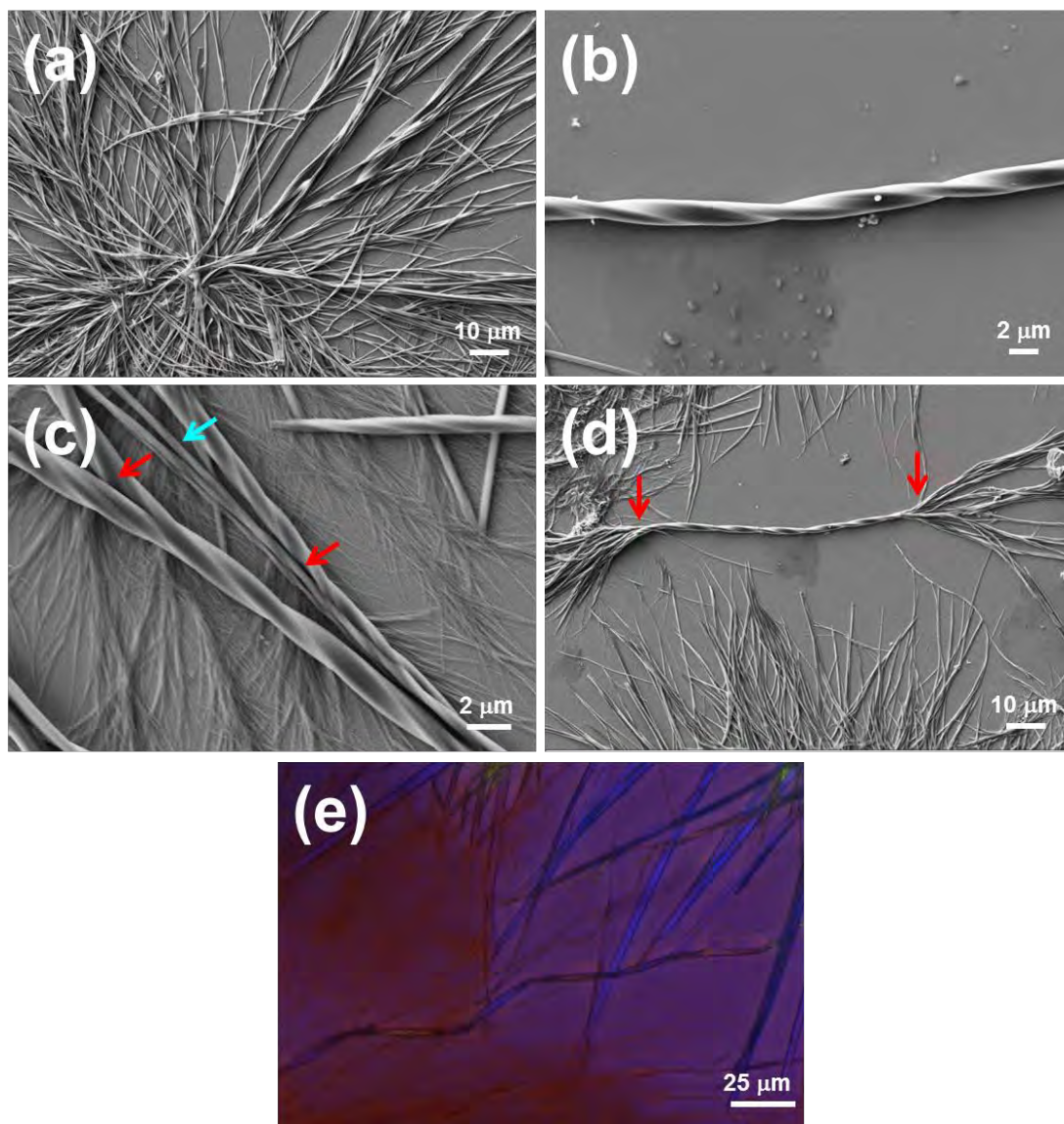


Figure 3

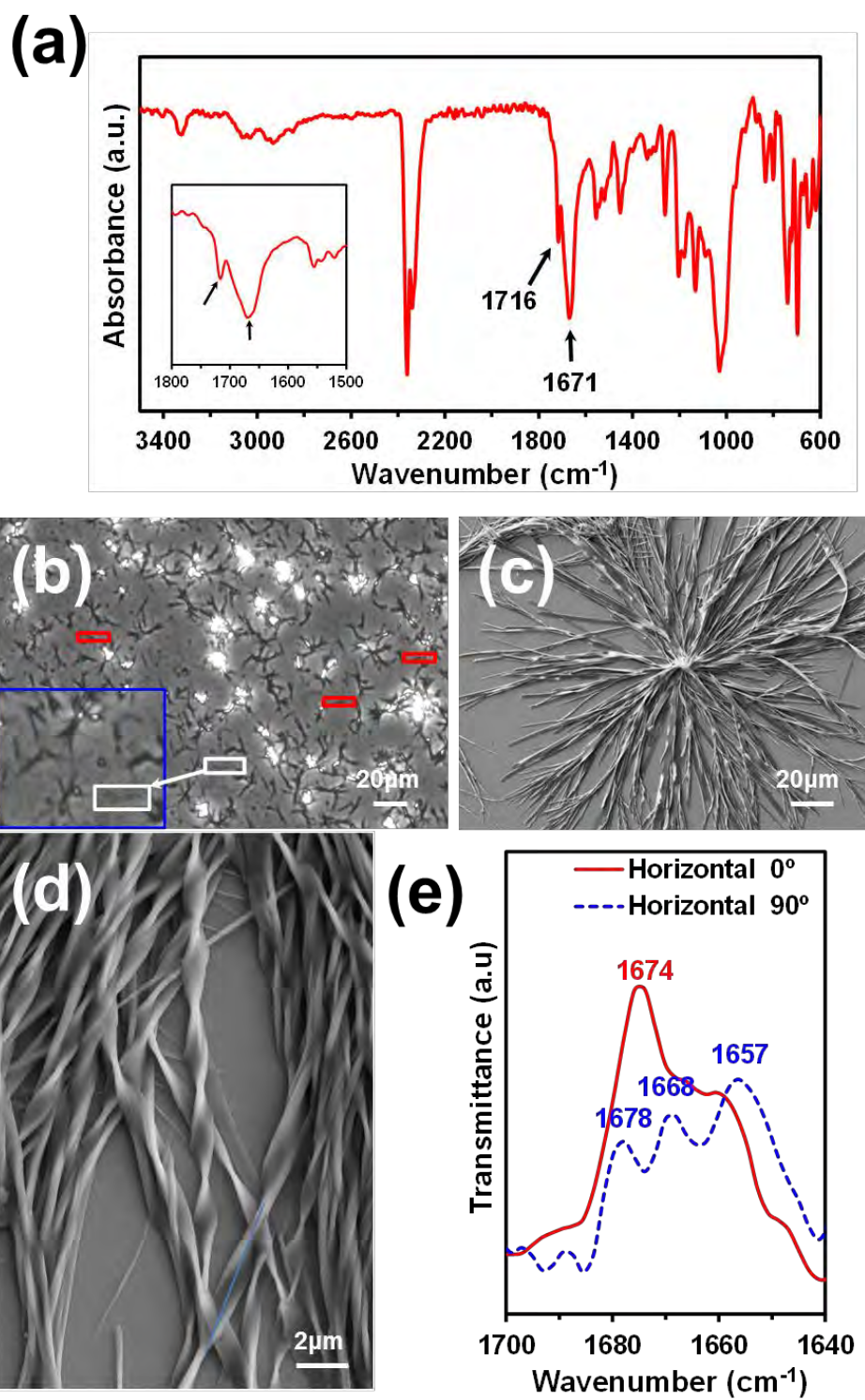


Figure 4

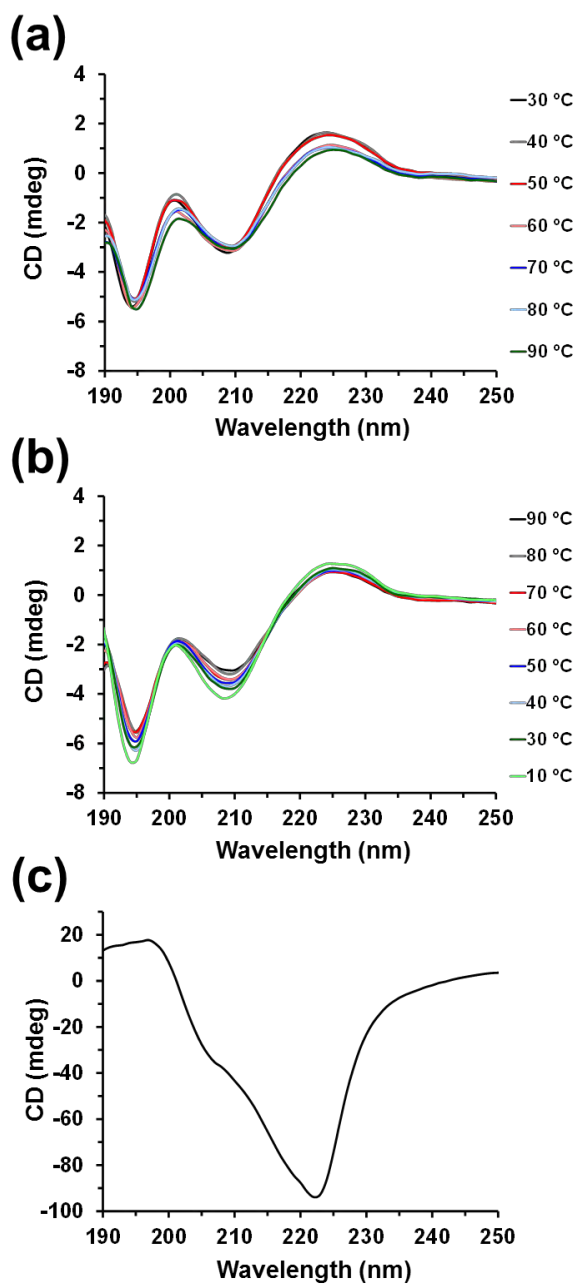


Figure 5

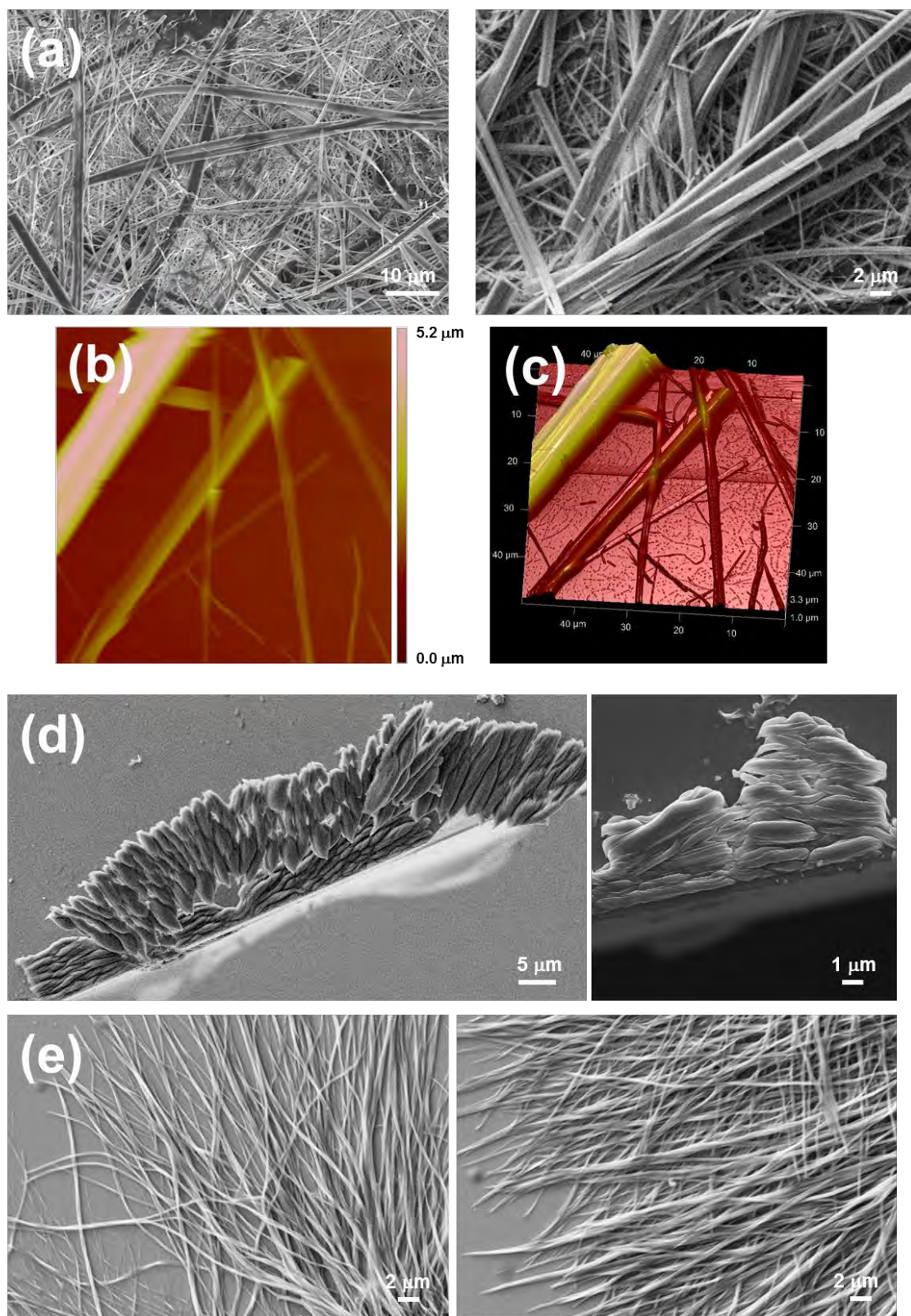


Figure 6

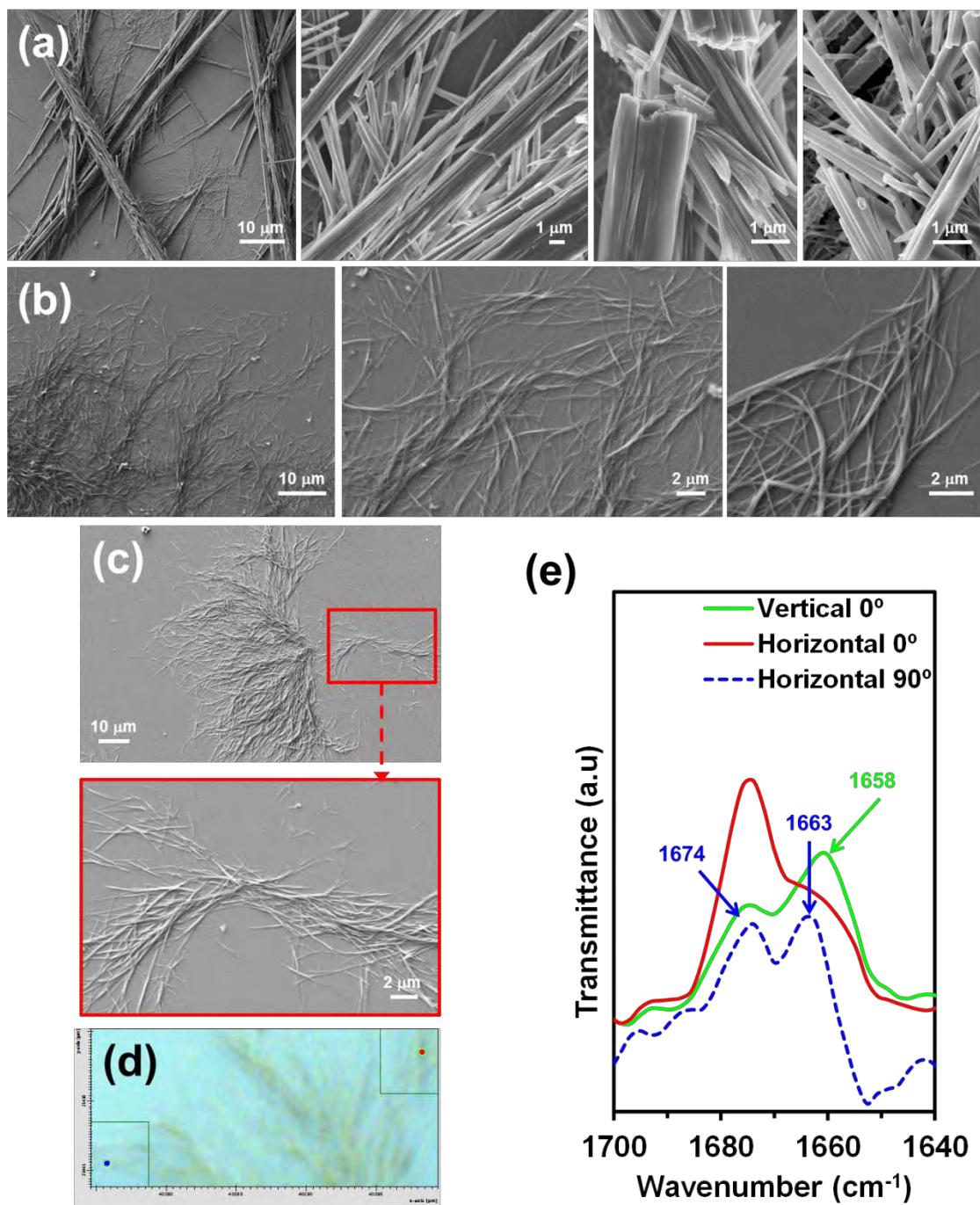


Figure 7

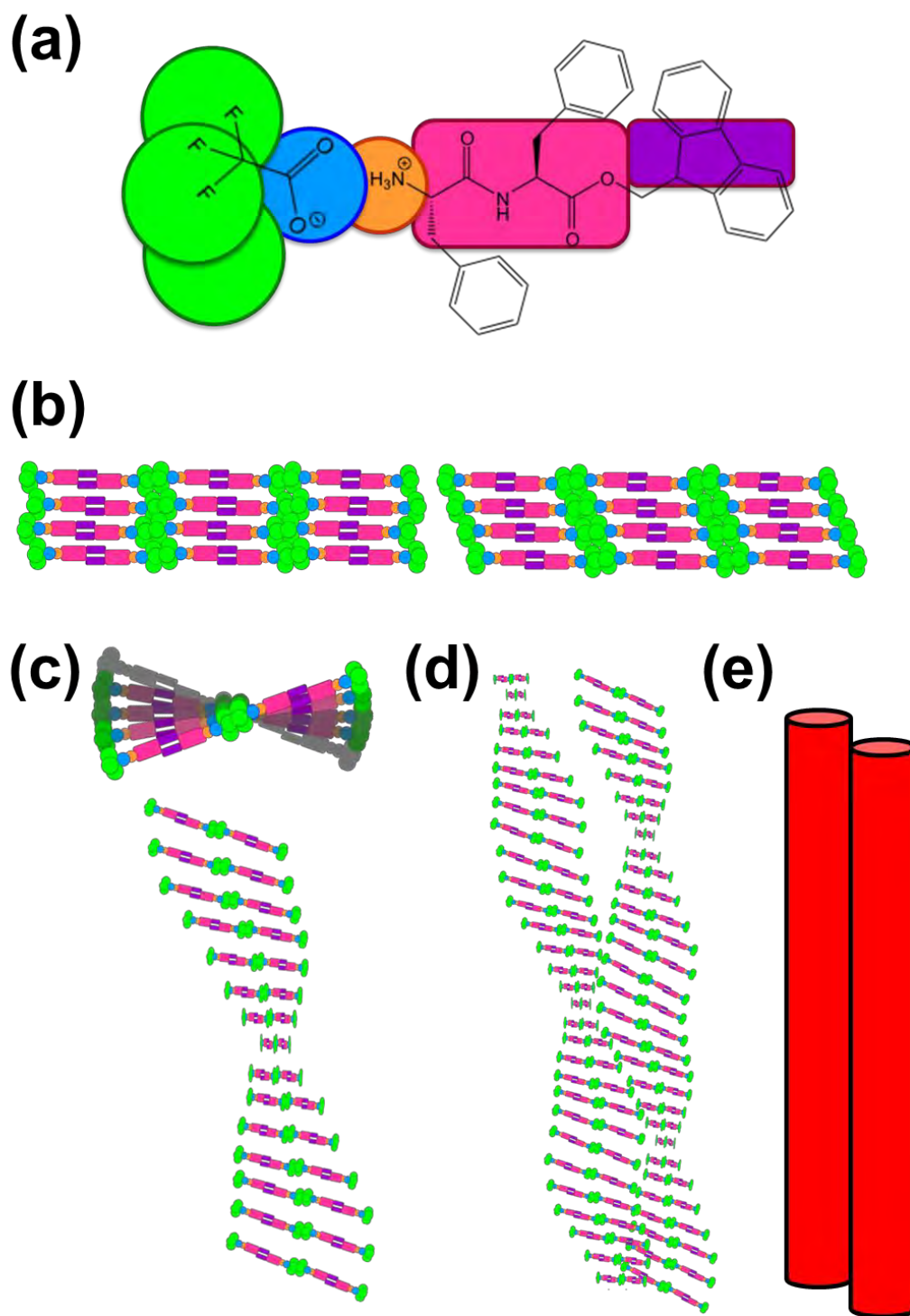


Figure 8

TOC

