Effect of pH on the Conformational Transition of Silk Fibroin in Aqueous Solution Monitored by Thioflavin-T Fluorescence

Ben Jia · Lan Jia · Jingxin Zhu

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Abstract
In this work, the potential application of the fluorescence dye Thioflavin-T (ThT), which can specifically bind to amyloid, as a powerful tool for monitoring secondary structural transitions of silk fibroin (SF) induced by pH in low solution concentrations was examined. Results showed that ThT emission intensities substantially increased when pH decreased from 6.8 to 4.8. This increase may be ascribed to conformational transitions from random coil to β-sheet. The morphology and secondary structure of SF were also investigated via TEM, AFM and circular dichroism spectroscopy. The information obtained herein can be utilized not only for the development of convenient and efficient noninvasive method for monitoring the assembly behavior of SF in aqueous solution but also for in vitro fluorescence imaging.

Keywords Conformational transition · Fluorescence spectroscopy · Protein structures · Silk fibroin · Thioflavin-T

Introduction
The self-assembly of natural proteins or designed peptides into fibrillar structures based on β-sheet conformation is a ubiquitous and important phenomenon [1–3]. These structures typically include conserved peptide fragments from silk proteins and critical peptide fragments from amyloid proteins related to degenerative diseases. Owing to the adjustable mechanical properties and biodegradability of silk protein-based materials that are related closely to the content of β-sheet crystallinity, they are promising candidates in material sciences and biotechnology [4, 5].

Various methods, such as Fourier transform infrared spectroscopy (FTIR) [6, 7], Raman spectroscopy [8], circular dichroism (CD) [9], nuclear magnetic resonance (NMR) spectroscopy [9, 10], X-ray scattering [11], rheology [12, 13], and fluorescence spectroscopy [14], have been used to study the transition from random coil and/or helix to β-sheet in fibroin solutions. However, most of the studies that adopted these methods yielded static rather than dynamic data as they only described the conformations of silk protein before and after natural spinning or under specified conditions. A more useful endeavor is to follow protein conformational changes continuously over time while they are induced by various factors, such as shear force, pH, metallic ions, temperature change, and low dielectric constant solvents. This undertaking would yield important data on the kinetics of conformation transitions that cannot be obtained by static experiments [15]. Thus, a detailed understanding of the sequence of events that occurs as random coil and/or helix conformation changes to β-sheet in silk proteins can be achieved.

Fluorescence spectroscopy is widely used to interpret protein structure in real time. For example, the intrinsic fluorescence relative changes in tryptophan (Trp) and tyrosine (Tyr) in silk fibroin (SF) can monitor conformational states among silk solution, gel, and scaffold samples [16]. Moreover, 8-anilino-1-naphthalene-sulfonic acid (ANS) is utilized as an external probe to investigate the effects of environmental factors in aqueous solutions [17].

Since its first description in 1959 [18], Thioflavin-T (ThT) has become the “gold standard” for selectively staining and identifying amyloid fibrils because its fluorescence emission intensifies upon binding to fibrils [19]. Given the similarities in self-assembly into fibrillar structures between SF and amyloid protein, the fluorescent dye ThT, which specifically binds to amyloid protein, was utilized herein as an external probe to monitor conformational transitions of SF under different pH conditions.
The design rationale is illustrated in Scheme 1. The enhancement of ThT fluorescence emission is associated with elevated β-sheet contents as pH decreases from 6.8 to 4.8. This enhancement makes ThT a particularly convenient and sensitive tool for monitoring conformational transitions in low solution concentrations. Intrinsic Trp residues and another commonly used hydrophobic dye, namely, ANS, were also investigated for comparison. The morphologies were observed via transmission electron microscopy (TEM) and atomic force microscopy (AFM). FTIR and CD spectroscopy were performed to monitor the secondary structures of SF.

Materials and Methods

Materials

ThT and ANS were purchased from Sigma Co. (USA). Tris (hydroxymethyl) aminomethane (Tris) and 2-(N-morpholino)ethanesulfonic acid (MES) were obtained from Shanghai Sangon (Shanghai, China). All other reagents were of analytical grade and used as received. All aqueous solutions were prepared with Milli-Q purification system.

Preparation of SF Solutions

Aqueous SF solutions were prepared following the method described in a previous report [13]. In brief, degummed fibers were dissolved in 9.3 M LiBr solutions, and SF solution was dialyzed against distilled water using Slide-a-Lyzer dialysis cassettes (MWCO 3500, Pierce) for 3 d (triple-distilled water for the last day) until the conductivity of the dialyzed water was close to that of distilled water.

Optimization of the Probe

ThT stock solution at 1 mM concentration was prepared by Milli-Q water and filtered through a 0.22 μm filter. Afterward, 2 mL of SF solutions (10 mg mL^{-1}) were mixed with different amounts of ThT stock solution (the final ThT concentration was 2.5, 5, 7.5, 10, 20, and 40 μM). The mixtures were diluted to 4 mL with water and kept at room temperature in the dark for 2 h.

The Effect of pH on the Structure Change of the SF

To investigate the conformation change of the SF solutions with respect to pH, the mixtures of SF and ThT were added to 20 mM Tris–MES buffer with different pH (4.8, 5.0, 5.2, 5.6 and 6.8). The final concentrations of SF and ThT were 5 mg mL^{-1} and 20 μM, respectively.

Characterizations

Fluorescence measurements were performed using a Fluoromax-4 fluorescence spectrophotometer (HORIBA, USA). For intrinsic fluorescence spectra, the excitation wavelength was set as 280 nm and emission spectra were recorded ranged form 300 to 500. For ThT and ANS fluorescence measurements, the samples were excited at 420 nm and 390 nm, the emission spectra were recorded in the range of 430–650 nm and 400–650 nm respectively. Protein concentration for all the fluorescence experiments was 5 mg mL^{-1}, using slit width of 5 nm for both excitation and emission.

TEM was performed on a JEM-1200EX (JEOL, Japan). Samples for TEM measurements were prepared by placing a drop of colloidal dispersion on a carbon-coated copper grid, followed by evaporation of the solvent. To prepare the samples for AFM imaging, 2 μL of the diluted SF solution was cast on silicon wafer and was allowed to dry in ambient air for 20 min. The morphology of silk fibroin was observed by NX10 AFM (Park Systems, Korea) in air. AFM scan rate was 1.0 Hz and imaging resolution was 256 × 256 pixel.

Samples were freeze-dried completely before being grinded into powder and pressed into pellets with KBr for measuring infrared spectra with a Fourier transform infrared spectrophotometer (FTIR) (Bruker VERTEX80 V, Germany).
Circular dichroism (CD) spectra were recorded on a MOS-500 spectropolarimeter (Bio-Logic, France) by using a quartz cell with a 1 cm path length under a nitrogen atmosphere. The spectra of sample solutions (5 mg mL$^{-1}$) were obtained from 190 to 260 nm with a 0.2 nm step and 2 s collection times per step, taking three averages.

The fibroin aqueous solutions under different pHs were kept at ambient temperature for 5 days to ensure the conformational transition for TEM, AFM, FTIR and CD measurements.

Results and Discussion

Optimization of the Probe

Fluorescence spectroscopy is widely used to interpret protein structure at atomic resolution. Aromatic amino acids, e.g., Trp, Tyr, and phenylalanine, offer intrinsic fluorescent probes for monitoring protein conformations, dynamics, and intermolecular interactions. Among them, Trp is the most popular probe for investigating protein structural changes because it is sensitive to the environment [20]. The fluorescence spectroscopy of Trp in SF was evaluated under different pH conditions to indicate changes in SF conformation. However, as shown in Fig. S1, the emission spectrum barely changed with pH. This result may be attributed to the fact that Trp is more sensitive to the water content of SF because most Trp residues are distributed on the surface of proteins, a property that induces poor sensitivity in aqueous solution. Thus, the external probe might be more suitable for our system.

ThT is perhaps the most widely used amyloid dye as this reagent specifically binds to the β-sheet structure of fibrils [21, 22]. In this study, ThT was used as an external probe to monitor the conformational state of SF. As shown in Fig. 1a, the fluorescence intensity of ThT at 485 nm substantially increased upon binding to the hydrophobic region of SF compared with free ThT. Thus, this phenomenon was qualified for fluorescence assay. The relationship between ThT concentration and fluorescence intensities at 485 nm of the SF solution is depicted in Fig. 1b. The highest intensity was obtained with 20 µM ThT, and it decreased when the concentration exceeded 30 µM probably because of the self-quenching effect of excessive ThT. Hence, 20 µM ThT was chosen as the optimum concentration and used for further analysis.

Effect of pH on Structural Changes in SF

pH is clearly an important factor in the formation of supramolecular assemblies of Bombyx mori SF because it has an influence on electrostatic interactions [12]. The fluorescence emission spectra of SF at pH 4.8, 5.0, 5.2, 5.6, and 6.8 are given in Fig. 2. These pH levels represent the conditions at different parts of the silk gland of Bombyx mori: pH 4.8 for the anterior division and pH 6.8 for the posterior division [23].

Emission intensity increased by about 3.6-fold when pH decreased from 6.8 to 4.8, indicating the β-sheet conformation of SF in aqueous solution. Photographs of SF solutions with the ThT probe at pH 4.8, 5.2, and 6.8 displayed the same trend. The fluorescence strategy could be used as a real-time method for monitoring the assembly process. The emission intensities at 485 nm initially increased with

![Fig. 1](image-url) (a) Fluorescence emission spectra of ThT (20 µM) and SF+ThT solutions in water. (b) Fluorescence intensity as a function of ThT concentration at 485 nm
incubation time at pH 4.8 and 5.0 within 5 days and then decreased over time. This result might be ascribed to the aggregated formation and the self-quenching of the ThT dye. By contrast, the intensity scarcely increased at pH 6.8. The pH and time dependence of ThT fluorescence was also determined. Results showed that the effect of pH and incubation time were negligible for pure ThT (Fig. S2). The conformational transformation of SF in low concentration solution is not easily detected as the quite slow process, and the low β-sheet content leads to low signal strength. The results showed that ThT probe was a sensitive and effective fluorescent probe in this study.

ANS is also a frequently used extrinsic fluorescence probe for exploring changes in hydrophobic regions during conformational transitions of SF. Its fluorescence behavior upon binding to SF was also investigated for comparison. The fluorescence spectrum of pure ANS exhibited a wide band with very low fluorescence intensity and an emission maximum of 542 nm in polar environments (Fig. S3a). The fluorescence intensity of ANS evidently increased in the presence of SF, which was accompanied by shifting of the emission maximum to 471 nm. The emission intensity of ANS at 470 nm also increased when pH decreased from 6.8 to 4.8. However, compared with ANS, the ThT probe showed better sensitivity as the fluorescence enhancement was more obvious. Compared with free ThT, the fluorescence of the SF solution with the ThT probe increased by about 25-fold at pH 4.8, whereas the fluorescence of the SF solution with the ANS probe increased by about sevenfold only (Fig. S3b). The difference in this phenomenon might be ascribed to the
possibility that ANS was sensitive to the hydrophobic environment, whereas ThT was bound to the β-sheet structure. Moreover, SF (PI ~4.5) is negatively charged at pH 4.8 to 6.8, whereas ANS has a negative sulfonate ion in aqueous solution [24]. The repulsion force between SF and ANS also made them difficult to be closely bound.

The assembly process was further investigated via TEM. As pH was decreased, the morphology of SF transitioned from spherical to dendritic aggregates. At pH 4.8, the aggregates formed by rod-like structures appeared (Fig. 3a, b), whereas at pH 6.8, heterogeneously sized micelles with diameters ranging from several tens of nanometers to hundreds of nanometers were observed (Fig. 3d). Previous studies identified micelles at neutral pH [23]. These structures might be assembled extrinsically by the aggregation of solubilized globular protein chains driven by their amino acid hydrophobicity. At pH 5.2, irregular and patchy aggregates can be seen in the image (Fig. 3c).

The results of AFM were basically consistent with those of TEM. At pH 4.8, SF proteins assembled into granules and displayed a tendency to join into bundles of fibrils (Fig. 4a). Moreover, the length of the fibrils reached micron scale. At pH 5.2 and 4.8, only particle-like aggregates can be seen, and the proportion of larger aggregates was higher at pH 5.2 (Fig. 4).

Differences among the FTIR spectra of SF under different pH values were not remarkable. As shown in Fig. S4, the peak at 1650 (amide I) and 1545 cm\(^{-1}\) (amide II) were associated with the random coil of SF, whereas the peak around 1630 (amide I), 1530 (amide II), and 1245 cm\(^{-1}\) (amide III) were characteristic of β-sheet structure. Both random coil and β-sheet structure existed in the solutions under different pH values. The peak at 1245 cm\(^{-1}\) was more obvious at pH 4.8 than at pH 6.8. This result may be attributed to the low β-sheet content in the aqueous solution.

CD spectra were used to verify the second structure of SF under different pH values, as shown in Fig. 5. At pH 6.8, the CD spectra demonstrated a strong negative band centered around 193 nm and a negative one at 215 nm, indicating the dominance of random-coil structure. However, when pH decreased to 4.8, the two bands drastically inverted to a positive band at 195 nm and to a negative band at ~215 nm, indicating the emergence of β-sheet conformation. The CD
spectra indicated a decrease in the content of random coil and an increase in β-sheet.

**Conclusion**

In summary, the conformational transitions of SF in aqueous solution under different pH values were investigated using ThT, a fluorescence dye that specifically binds to β-sheet. A substantial fluorescence enhancement was demonstrated as pH decreased. The time dependence of ThT fluorescence was also investigated under different pH values. The level of fluorescence attributed to β-sheet content substantially increased. ThT was more efficient than ANS and intrinsic Trp residues because it showed a more substantial fluorescence enhancement ratio. This result may be attributed to the ability of ThT to specifically bind to β-sheet. The morphological transition from spherical micelles to rod-like aggregates and fibrils due to a reduction in pH was observed via TEM and AFM, respectively. The transition of the secondary structure from random coil to β-sheet during this change was confirmed via CD spectroscopy. Thus, the fluorescence external probe may provide a direct and suitable noninvasive tool for monitoring the assembly behavior of proteins with conformational transitions in low solution concentrations. Owing to its ability to specifically bind to β-sheet, ThT has the potential for bioimaging of silk-based materials in vitro.

**Supplementary Information** The online version contains supplementary material available at https://doi.org/10.1007/s10895-021-02841-x.

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**Declarations**

**Ethics Approval** No applicable.

**Consent to Participate** No applicable.

**Consent for Publication** No applicable.

**Conflict of Interest** The authors declare that they have no competing interests.

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