

# Long-Term Super-Resolution Imaging of Amyloid Structures Using Transient Binding of Thioflavin T

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**Abstract:** Amyloids are implicated in Alzheimer’s disease but cannot be well resolved by standard light microscopy. We developed a tool to directly image native amyloid structures and dynamics at nanometer resolution over minutes to days. © 2019 The Authors

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Nanometer-sized aggregation intermediates are the main culprits in amyloid toxicity [1] in Alzheimer’s disease (AD), but they cannot be resolved by conventional light microscopy. Single-molecule (SM) super-resolution (SR) fluorescence microscopy techniques, such as (d)STORM [2, 3], overcome the optical diffraction barrier (~250 nm for visible light) and visualize structures with nanoscale resolution. However, biological targets must be densely labeled with fluorescent molecules [4, 5], which can potentially alter the structure of interest. Furthermore, photobleaching of tagged fluorescent molecules limits measurement time and prevents long-term imaging of targets.

A related SM-SR technique, called PAINT [6], uses combinations of fluorophore binding and unbinding, diffusion into and out of the imaging plane, and/or spectral shifts upon binding to generate flashes of SM fluorescence. When in the immediate vicinity of their target, these probes either become fluorescent, temporarily bind to the target, or both, thereby creating a “flash” of fluorescence that is used to locate the target of interest. These blinking events are recorded over time, and image-processing algorithms [7] measure the position of each bright molecule with high precision to reconstruct a SR image.

Herein, we report a technique to image native amyloid structures using Thioflavin T (ThT) on the nanometer scale, called Transient Amyloid Binding (TAB) imaging [8]. ThT fluorescence increases upon binding to amyloid proteins, transforming dark ThT in solution into its bright state [9,10]. The molecules emit fluorescence until they photobleach or dissociate from the structure, enabling us to record movies of “blinking” ThT molecules, localize their positions, and reconstruct the amyloid structure (Fig. 1A, B, C).

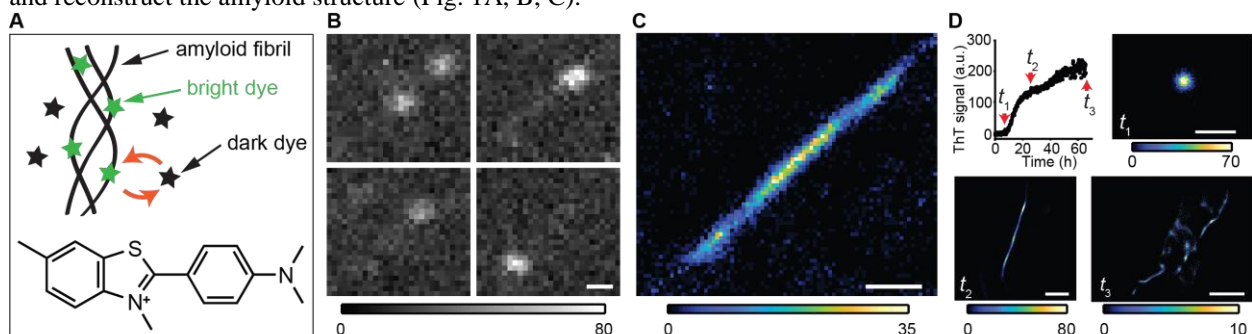


Fig 1. Demonstration of TAB imaging. (A) TAB mechanism explained: unbound dye molecules are dark while bound dyes are bright. (B) ThT blinking events on an Aβ42 fibril using TAB. Scale bar: 300 nm. Grey scale: photons per pixel. (C) Aβ42 fibril SR image reconstructed from the raw data in B. Scale bar: 300 nm. Color scale: localization per bin. (D) Top-left depicts the Aβ40 aggregation kinetic denoting  $t_1$  (8 h),  $t_2$  (24h), and  $t_3$  (66 h). SR images corresponding these time points show an oligomer, a fibril, and a fibril cluster. Scale bar: 0.5 ( $t_1$ ), 1 ( $t_2$ ), and 2.5  $\mu\text{m}$  ( $t_3$ ).

Accurate imaging of oligomeric structures is crucial to capturing the dynamics of Aβ aggregation and could open the door for future applications in cellular imaging of oligomeric structures. To demonstrate the versatility of TAB imaging, we imaged Aβ42 fibrils and Aβ40 aggregates with various morphologies (Fig. 1C, D), including oligomers, fibrils, and fibril clusters.

To image amyloid dynamics, it is essential to have a robust tool that can follow a single aggregate over hours or more. The continuous binding and unbinding of ThT from its amyloid target enables TAB imaging to maintain a

constant localization rate over 24 hours when imaging a single A $\beta$ 42 fibril (Fig. 2A, B). Therefore, TAB imaging can produce multiple time-lapse SR images with minute-scale temporal resolution over extended periods.

We next validated the capability of TAB for dynamic SR imaging. The time-lapse images (Fig. 2C-E) show the dissolution and remodeling of A $\beta$ 42 fibrils by epi-gallocatechin gallate (EGCG) [11]. Remarkably, TAB imaging captured the structural dynamics of amyloid fibrils for ~2 days, thus allowing us to observe remodeling over tens of micrometers with ~16 nm precision. In this experiment, we observed dynamics that were slower than at 37°C in solution [11], most likely due to the lack of agitation of fibrils that were adsorbed to the glass surface and to incubation at room temperature.

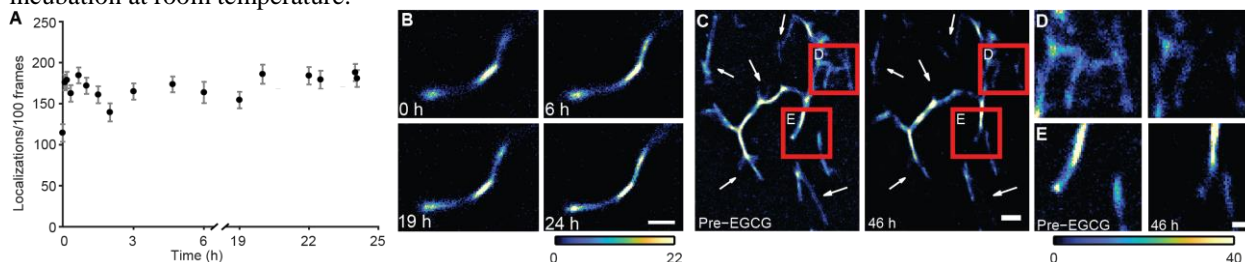


Fig 2. Long-term TAB imaging. (A) Average number of localizations/100 frames for each time point captured of a single A $\beta$ 42 fibril. (B) SR images of the A $\beta$ 42 fibril at the time points in A. Scale bar: 500 nm. (C) A $\beta$ 42 fibrils before addition of EGCG and 46 hours after adding EGCG. Scale bar: 500 nm. (D, E) Zoom-ins of portions of the structure in C showing fibril remodeling. Scale bar: 200 nm.

Previous SR studies had limited observation times due to photobleaching and due to the permanent attachment of probes to the amyloid fibril [12]. Because ThT probes continuously dissociate and new ThT molecules rebind to amyloid, TAB imaging is inherently resistant to photobleaching. The success of these experiments demonstrates the ability of TAB imaging to follow the dynamics of amyloid structures with nanometer resolution. This capability will be essential for visualizing drugs acting on amyloid structures in order to gain insight into their molecular-scale interactions with these structures.

In summary, TAB SR microscopy is a flexible imaging technique that can provide images of amyloid structures with nanometer resolution over observation times of hours. It is capable of imaging various stages of amyloid aggregation as well as dynamic imaging of fibrillar remodeling by an anti-amyloid drug. Nanoscale imaging of aggregation intermediates will provide a clearer understanding of which structures are toxic to cells and will pave the road for further study into molecular mechanisms of AD and other amyloid diseases.

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