



Optical Digital Assay for High-Sensitivity Quantification of Protein Protofibril Disassembly

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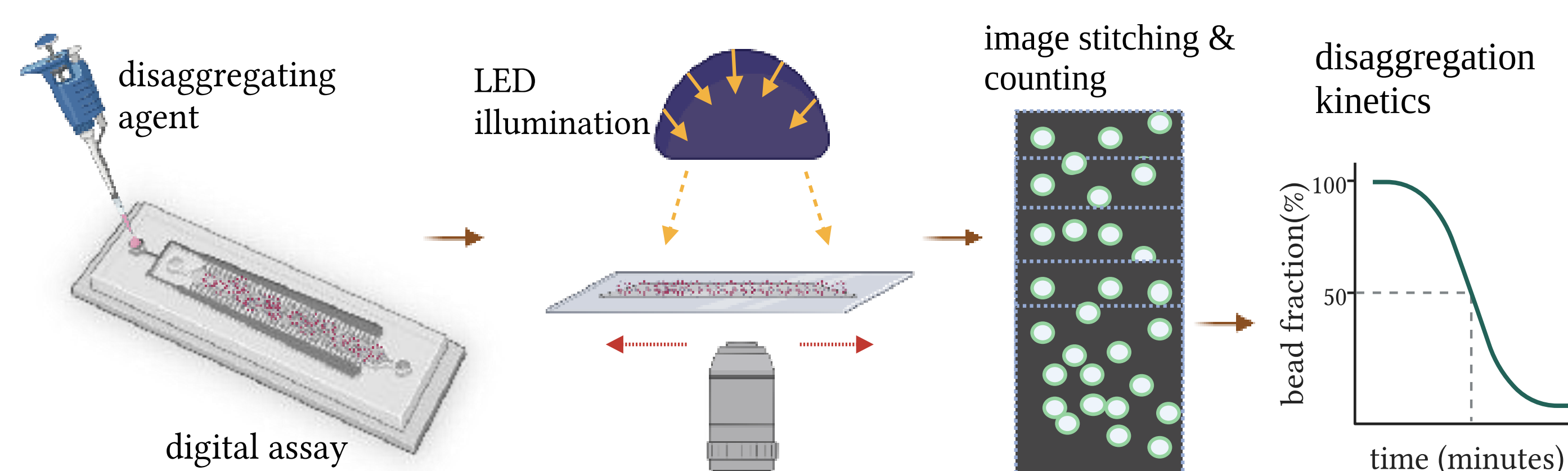
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Introduction

We present a latex bead-based digital molecular assay for sensitive monitoring of small molecule treatment-induced disaggregation of surface-tethered early-stage A β_{42} aggregates. The assay provides a digital readout of aggregate detachment kinetics, enabling quantitative detection of early disaggregation events that are difficult to resolve using conventional ensemble-averaged assays.

Schematic of the Latex Bead Assay



Methodology

- ❖ Early-stage A β_{42} aggregates were covalently immobilized onto latex beads.
- ❖ Aggregates were treated with the small molecules EPPS, HEPES, and PIPES.
- ❖ Dark-field mosaicking was used to generate large-field-of-view datasets for single-bead level digital bead counting.
- ❖ Bead-retention kinetics were determined by measuring the time-dependent decrease in retained bead counts using digital bead counting.
- ❖ Unbinding times and spatiotemporal dynamics of aggregate detachment were characterized.

Results - Digital Bead Assay

Representative dark-field images and progressive bead loss

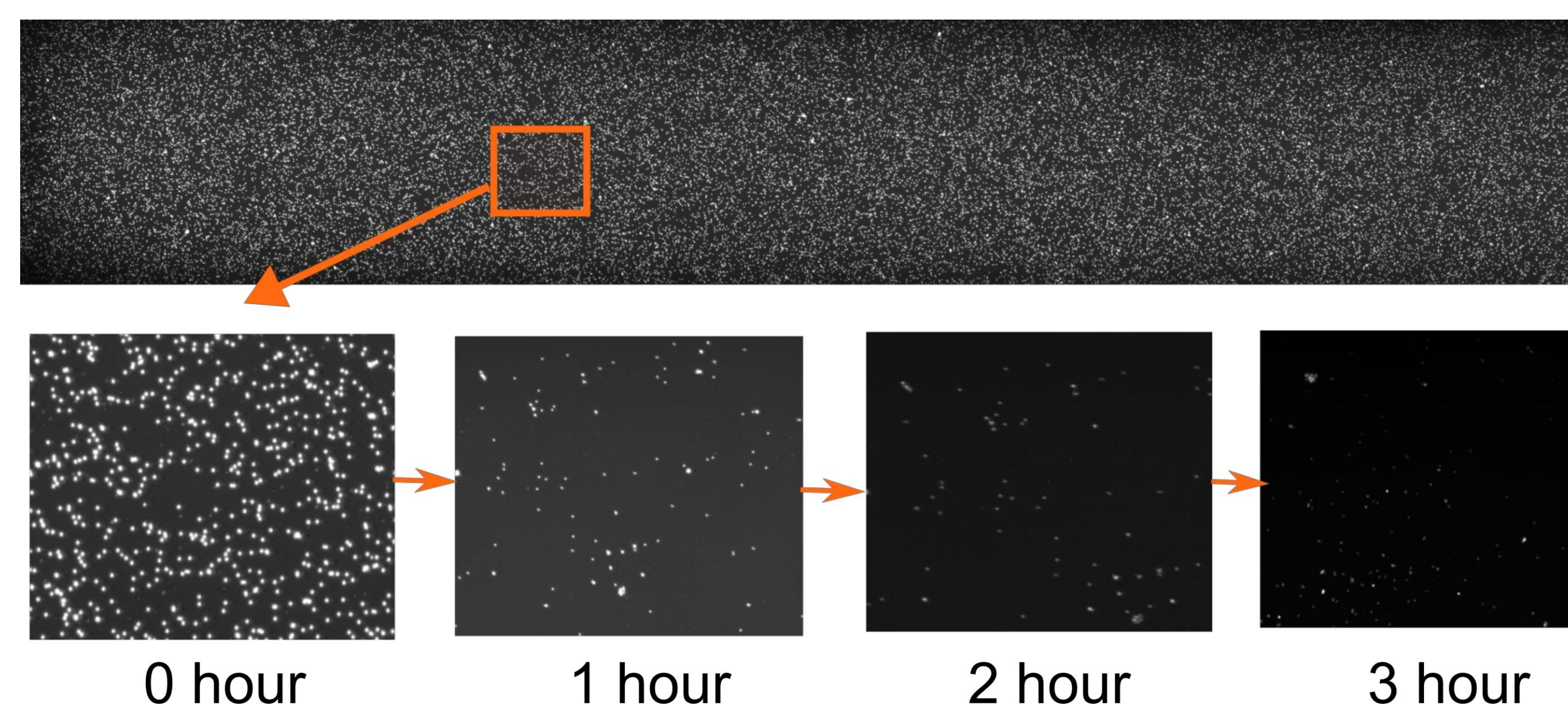
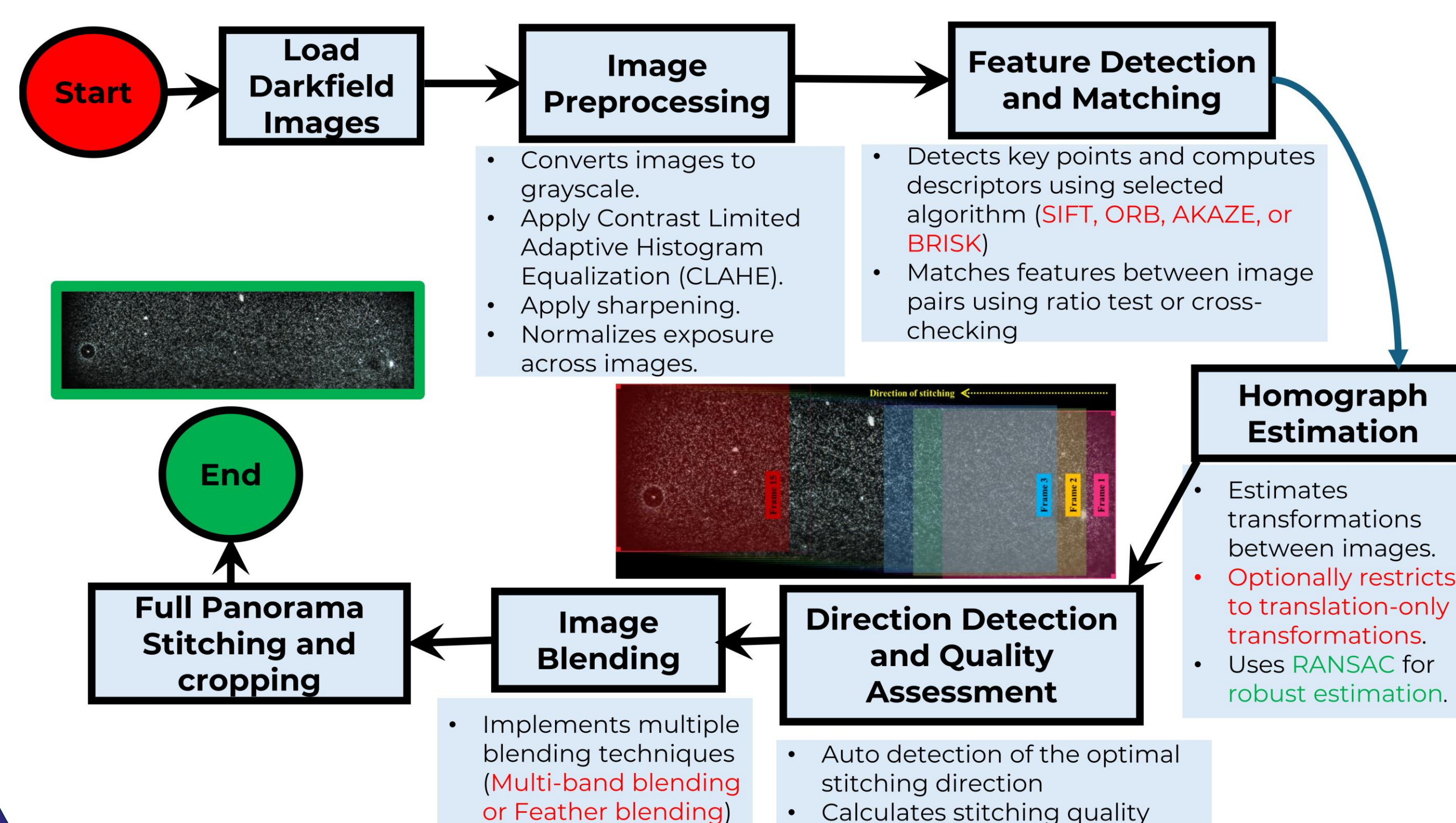


Image stitching workflow

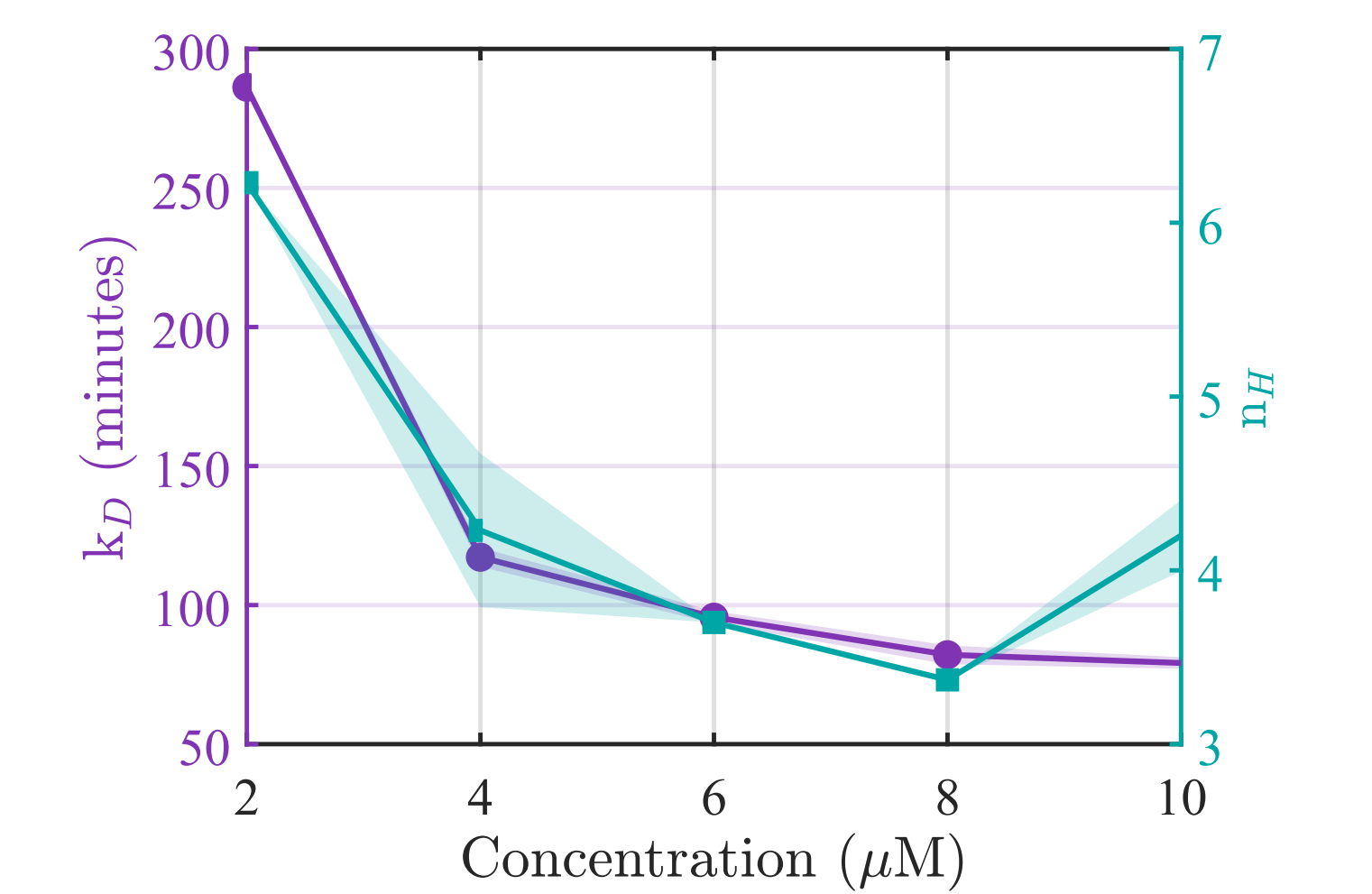
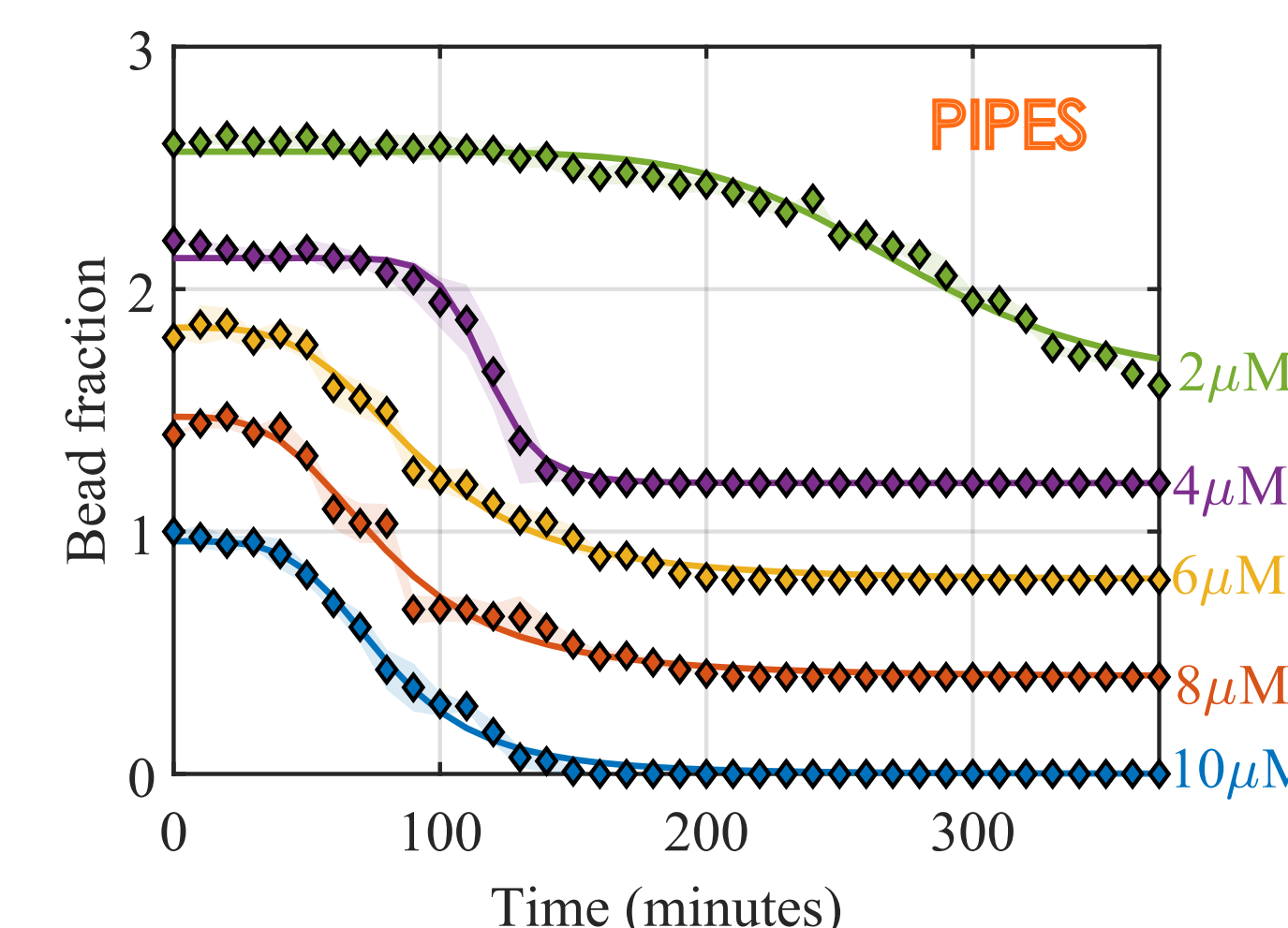
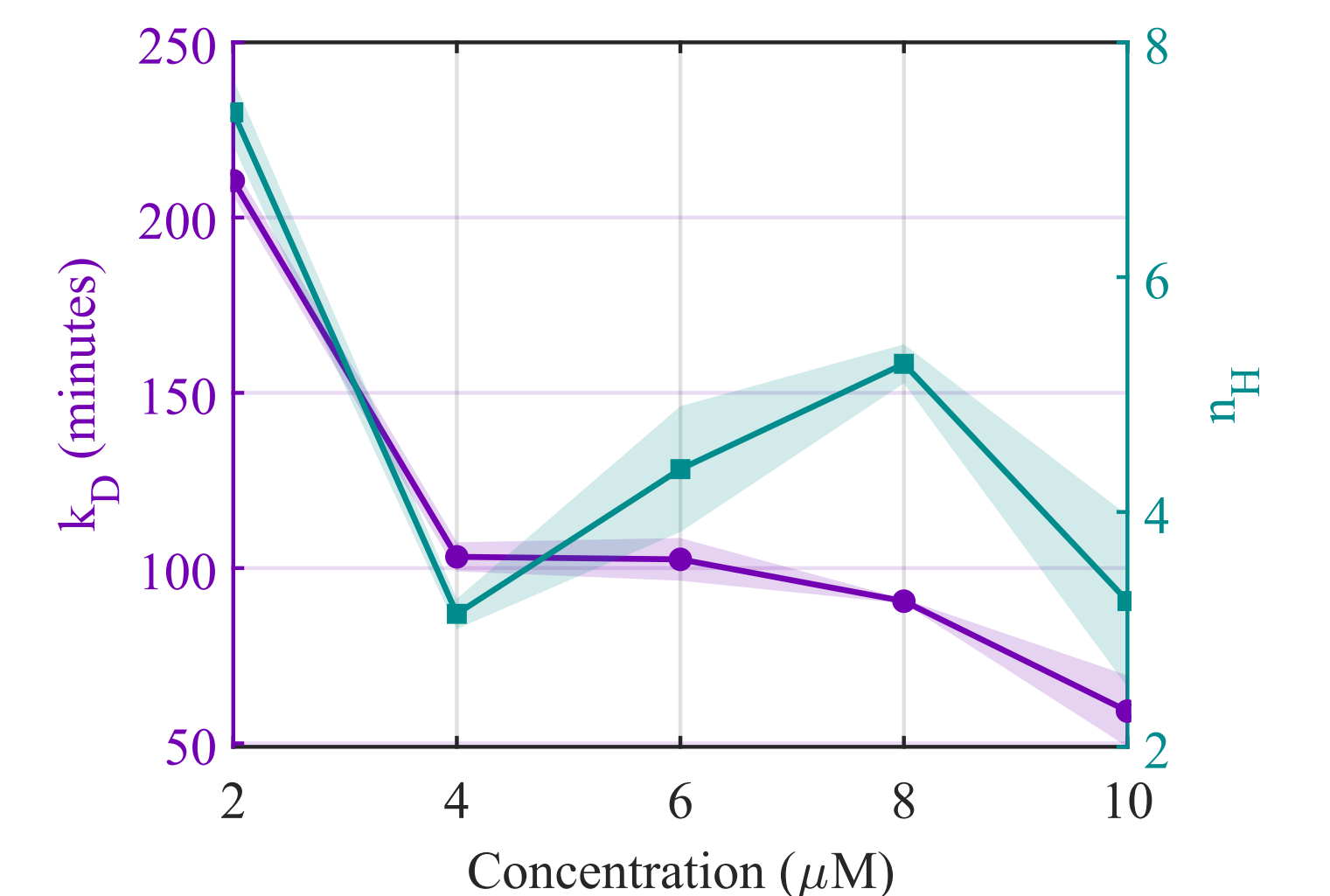
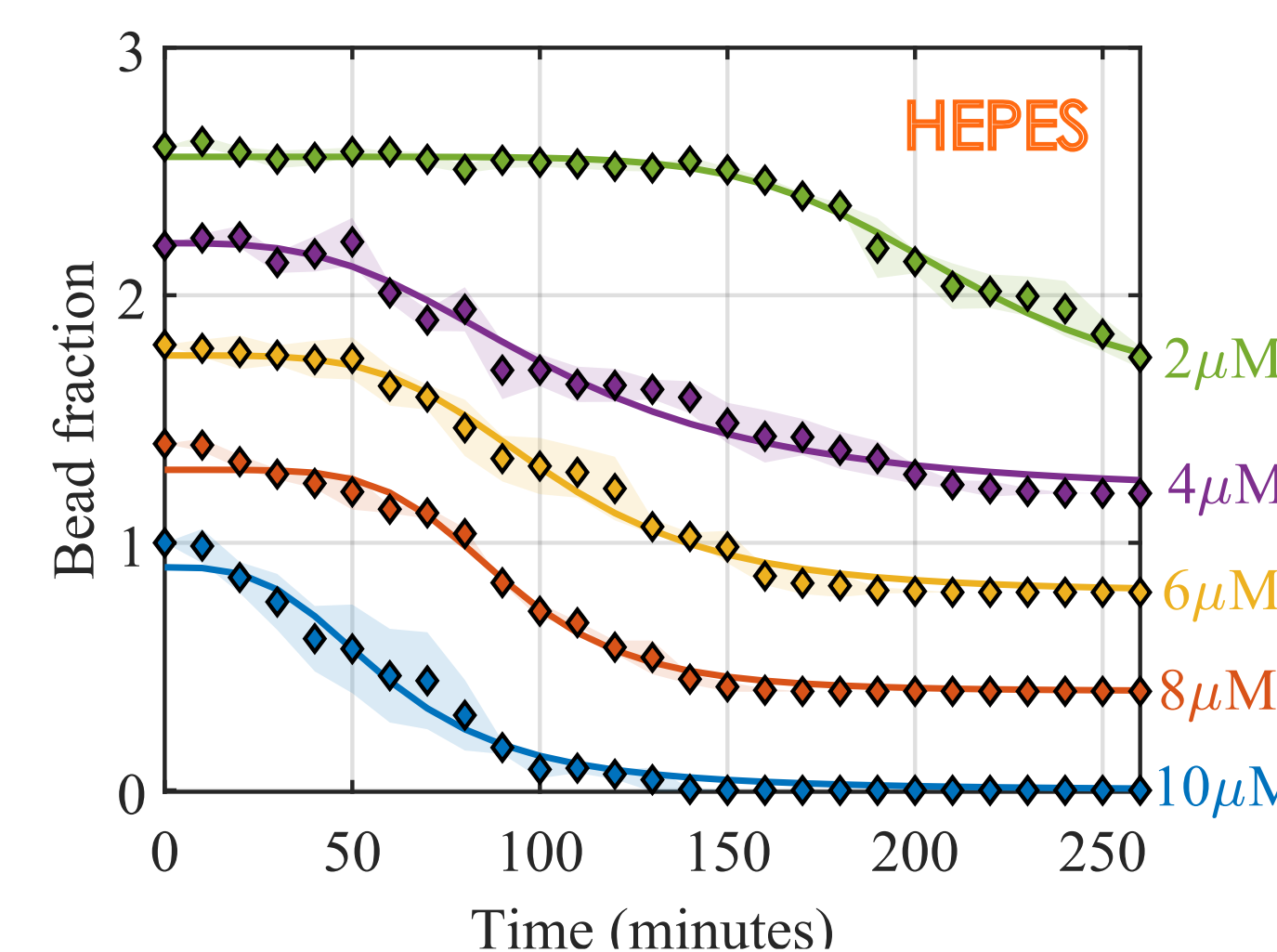
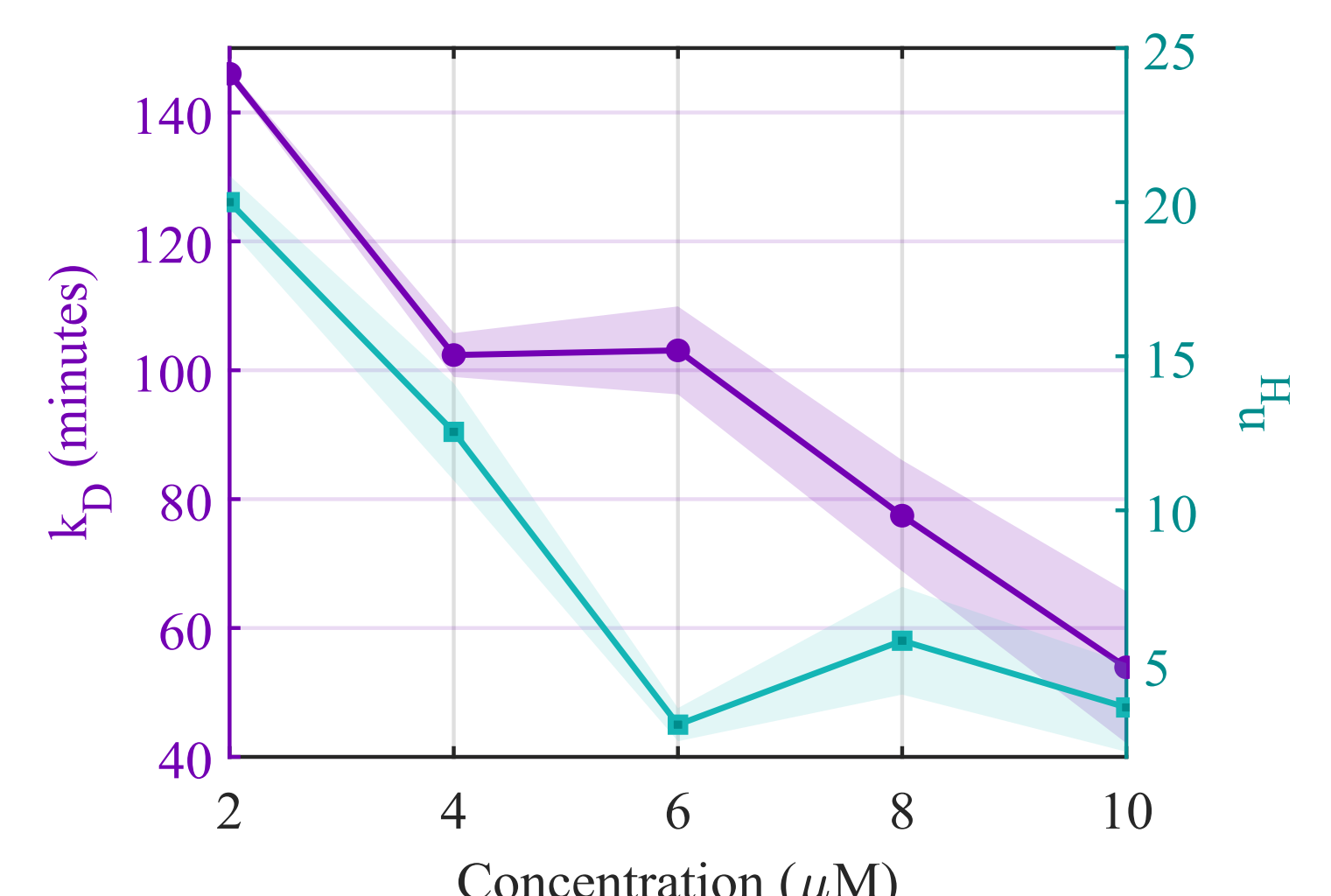
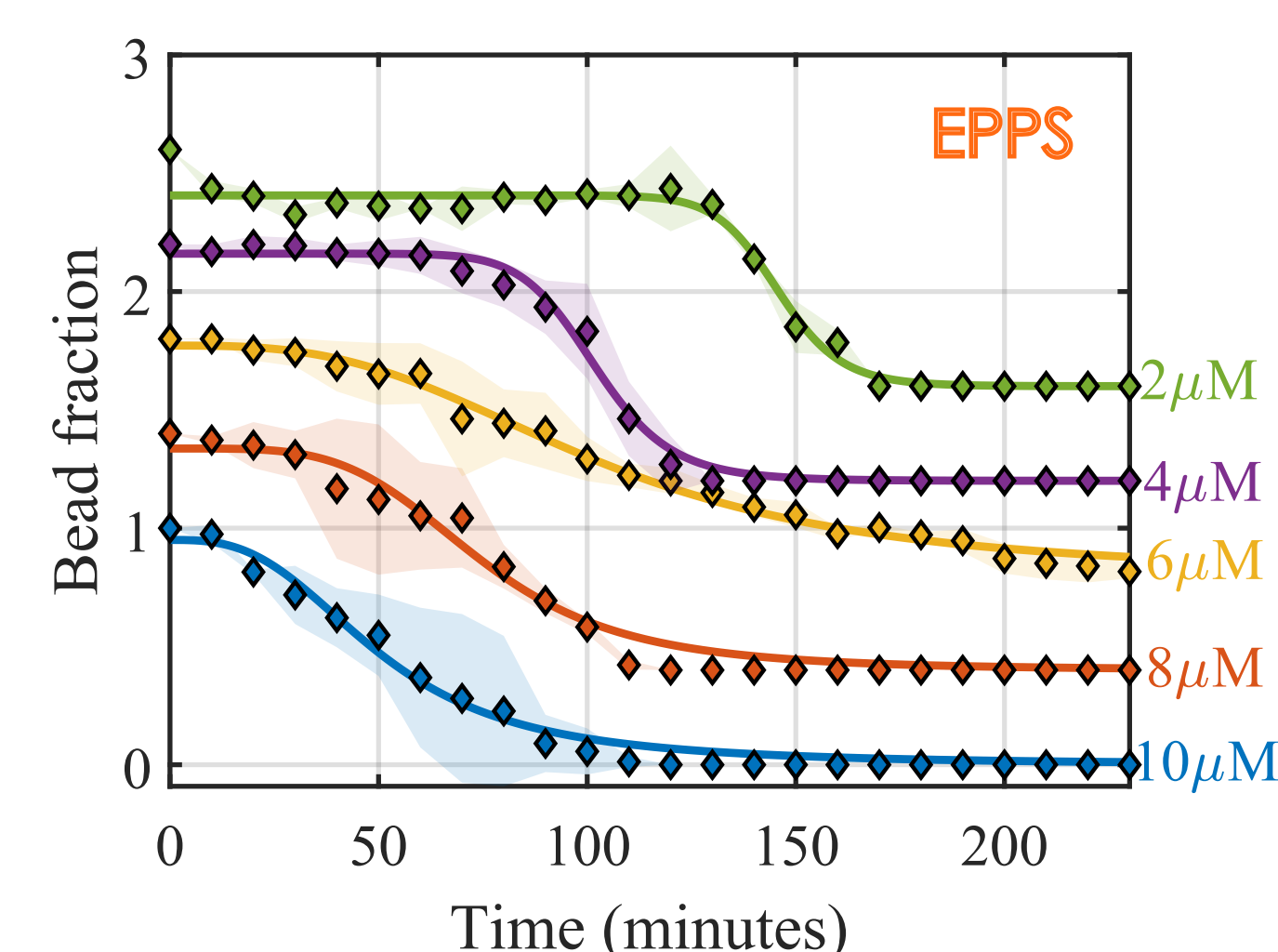


Results - Bead Detachment Analysis

$$\text{Hill Model: } F(t) = F_{\text{final}} + (1 - F_{\text{final}}) \left(\frac{1}{1 + (t/k_d)^{n_H}} \right)$$

Hill fits of temporal bead loss at different disaggregation agent concentrations

Extracted Hill parameters
 k_d = half time
 n_H = cooperativity



Conclusion

- ❖ Developed a latex bead-based digital molecular assay for sensitive monitoring of treatment-induced changes in surface-tethered early-stage A β_{42} aggregates.
- ❖ Enabled time-resolved digital analysis of bead detachment, revealing unbinding kinetics and spatiotemporal characteristics of aggregate destabilization.
- ❖ Captured early aggregate disaggregation in attomole quantities of surface-bound A β .
- ❖ Provides a versatile platform for quantitative studies of amyloid disaggregation induced by small molecules.

References

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