

## **Webinar**

### **Biophysical Characterization of Interactions between Apolipoprotein E and Amyloid- $\beta$**

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Alzheimer's disease (AD) is one of the most prevalent forms of dementia among the elderly population in the world. It is characterized primarily by the deposition of Amyloid- $\beta$  ( $A\beta$ ) in the brain. However, the major genetic risk factor of AD is apolipoprotein E4. Apolipoprotein E (ApoE) is the major lipoprotein present in the human brain. It exists as three isoforms viz., apoE2 (Cys 112, Cys 158), apoE3 (Cys 112, Arg 158), and apoE4 (Arg 112, Arg 158). ApoE plays a critical role in the modulation of  $A\beta$  metabolism in an isoform-dependent manner. However, its interaction with  $A\beta$ , causing AD, remains poorly understood.

In this talk, first I will discuss the molecular mechanism of interaction between apoE and  $A\beta$ , by comparing the effects of different domains of apoE on the aggregation of  $A\beta$ . We find all the domains of apoE delay the kinetics of aggregation of  $A\beta$  in substoichiometric concentration. A competitive binding assay further shows that the binding affinity of the full-length apoE to  $A\beta$  is much higher compared to the individual domains. Taken together, we hypothesize that a high affinity of the apoE- $A\beta$  interaction is achieved due to multivalent binding between oligomeric  $A\beta$  and full-length apoE. Further, we characterize the apoE- $A\beta$  complexes using the Two-colour coincident detection (TCCD) method. We find the complexes formed are heterogeneous in nature. Finally, I will discuss, how apoE affects various microscopic processes viz., primary nucleation, elongation, and secondary nucleation involved in the pathway of aggregation of  $A\beta_{42}$  using total internal reflection fluorescence microscopy (TIRFM). We find that all the isoforms of apoE inhibit the elongation and secondary nucleation of the fibrils.

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***4:00 PM***