

# The Dynamics of $\beta$ -Amyloid Aggregation

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$\beta$ -amyloid ( $A\beta$ ) aggregates are the presumed neurotoxins leading Alzheimer's Disease (AD). Aggregation of  $A\beta$  and other amyloidogenic proteins yields polymorphic products, both soluble oligomers and fibrils. This contrasts with normal protein folding, in which a single amino acids sequences leads to a unique protein fold. In this talk, I will discuss the polymorphism and dynamics of  $\beta$ -amyloid aggregates. As time permits, I will briefly consider up to four related topics: 1) Structure and polymorphism of fibrils formed by seeding of  $A\beta$  solutions by authentic AD brain amyloid. 2) Modelling of these fibrils as aberrant aqueous pores. 3) "Chaotic" aggregation of an internal fragment of  $A\beta$ ,  $A\beta_{16-34}$ . 4) Formation of  $A\beta$  soluble oligomers as a second-order phase separation.