



Biophysical aspects of amyloid fibrils

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Abstract:

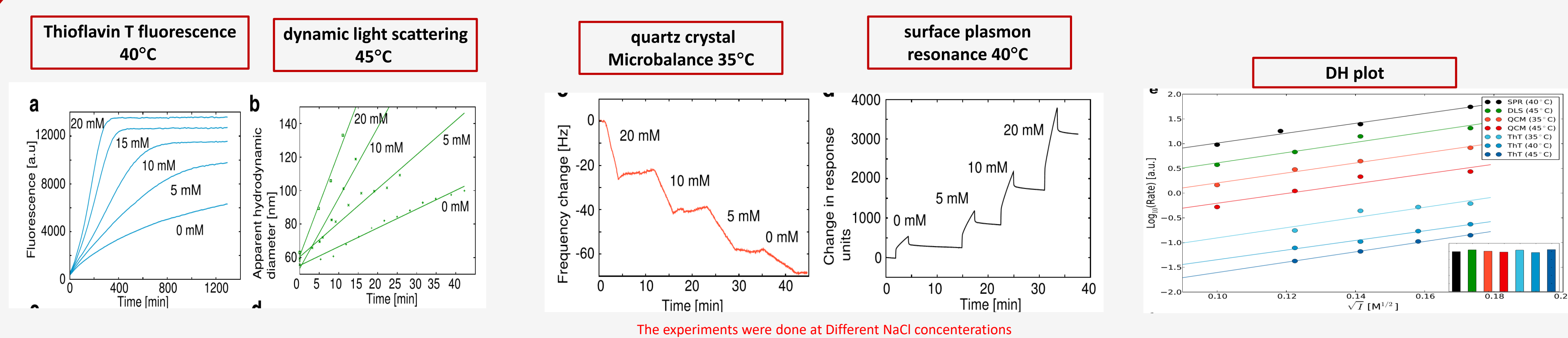
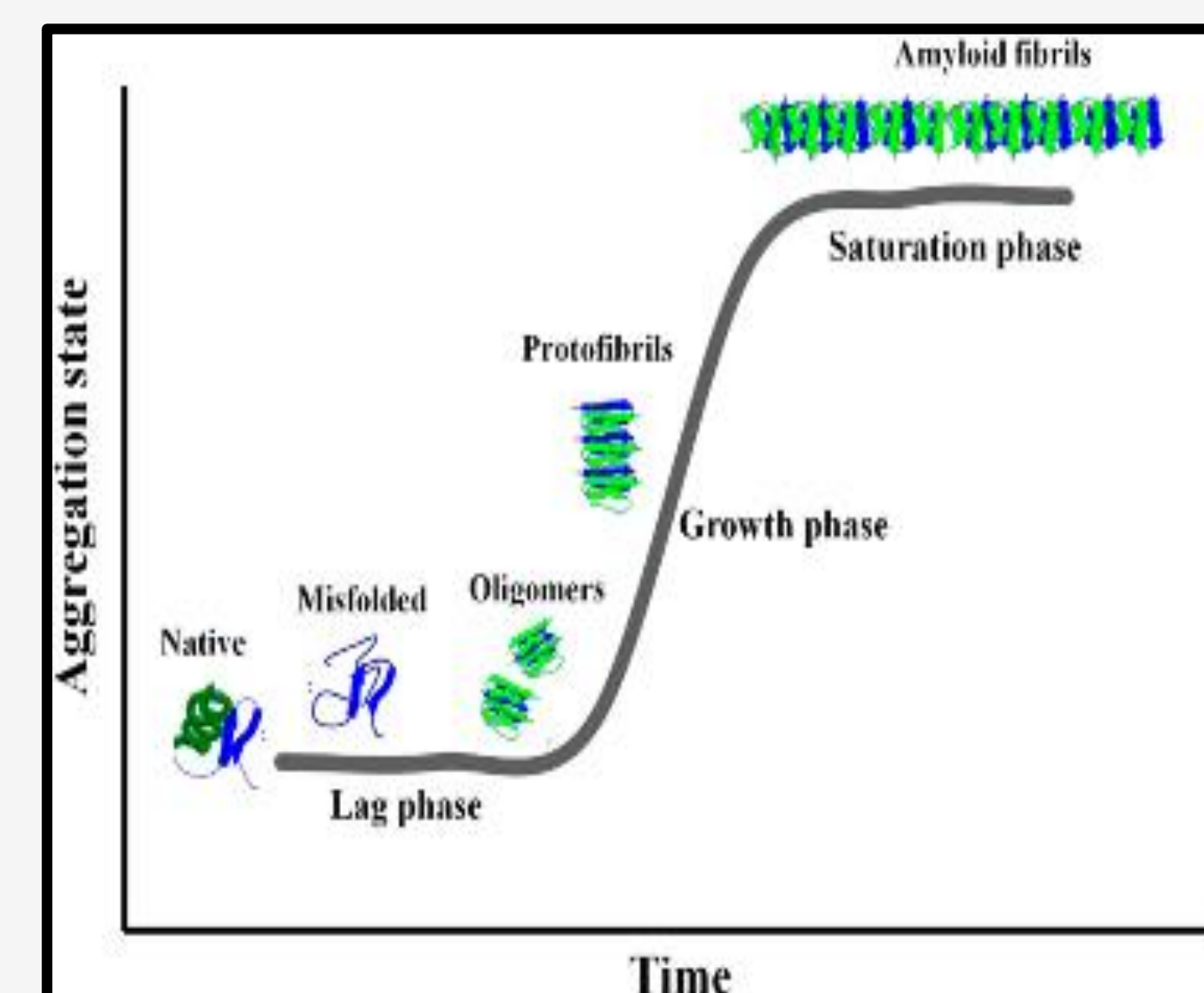
Introduction: Amyloid fibrils are associated with lots of human diseases such as **Alzheimer, Parkinson, spongiform encephalopathies and diabetes**. Various protofibrils assemblies together and form ordered packed B-parallel structures. Their crucial role in initiation of toxic process in neurodegenerative disorders is well known. Divers sequences of amino acids can form the amyloid state of proteins.

Methods: In this study, the spectroscopy methods were **NMR spectroscopy, CD spectroscopy and Infrared spectroscopy**. Kinetic measurements of amyloid fibril elongation were done using Thioflavin-T (**ThT**) and quartz crystal microbalance (**QCM**). Solution small angle x-ray scattering (**SAXS**) and differential scanning calorimetry (**DSC**) combined with simulated annealing of the protein were applied. Computational studies help to understand the application and analysing the obtained results.

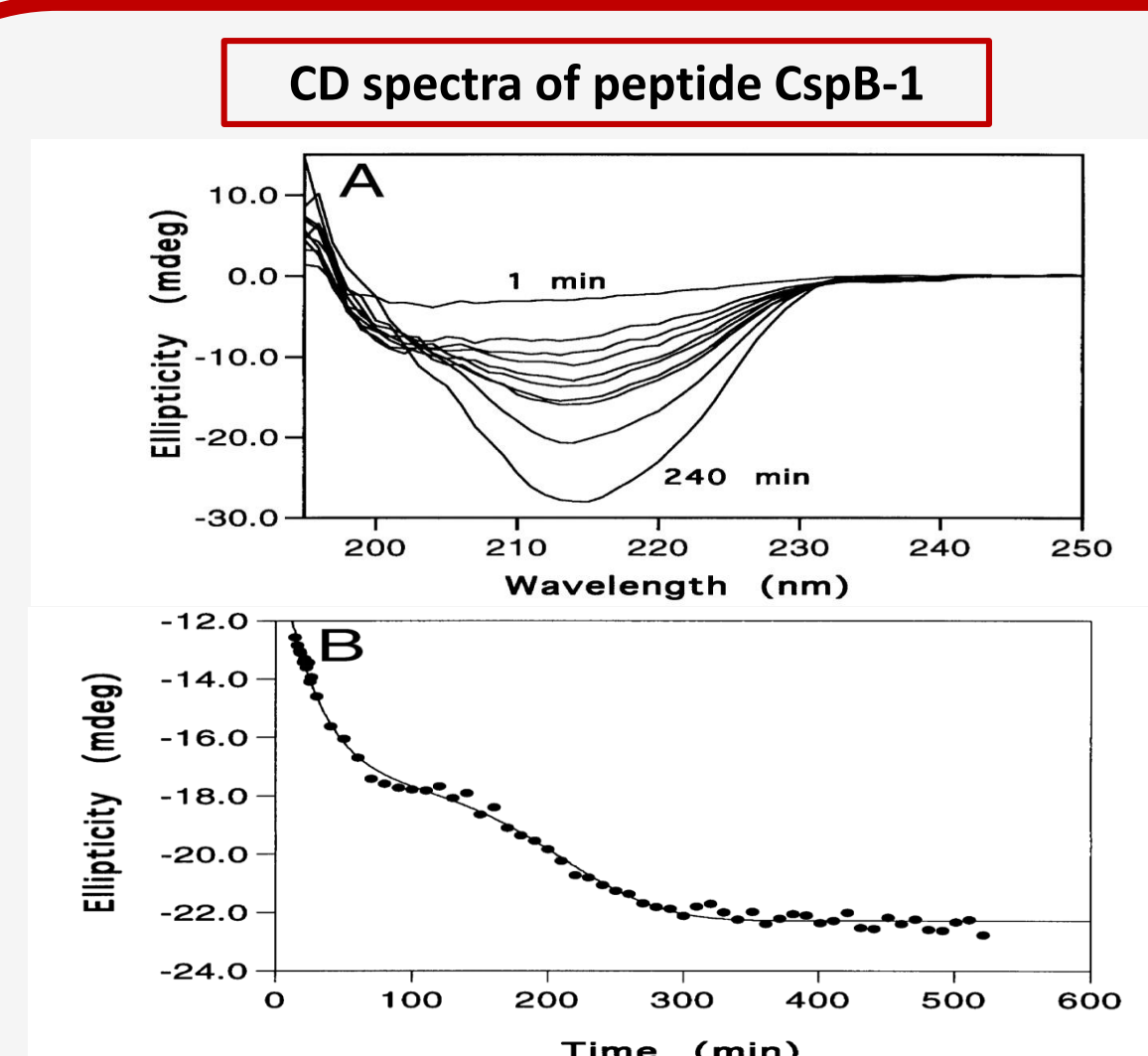
Results and discussion: Experimental kinetic measurements and theoretical analysis, showed that **electrostatic effects control protein aggregation**. Furthermore, the magnitude of binding of a variety of ions to protein molecules was determined. Our results suggest that **longer amyloid fibrils are more stable**. Our spectroscopy techniques confirmed that the formation of amyloid fibrils are a **generic property** of polypeptide chains, and for different peptides and proteins, the mechanism of formation is similar.

Conclusion: amyloid fibrils cause **neurotoxic effects** in neurodegenerative disorder. Detailed biophysical studies of amyloid fibrils can elucidate new aspects and features of this diseases and help our investigating on the **prevention of these fibril formation** and treatment of these diseases to be more effective.

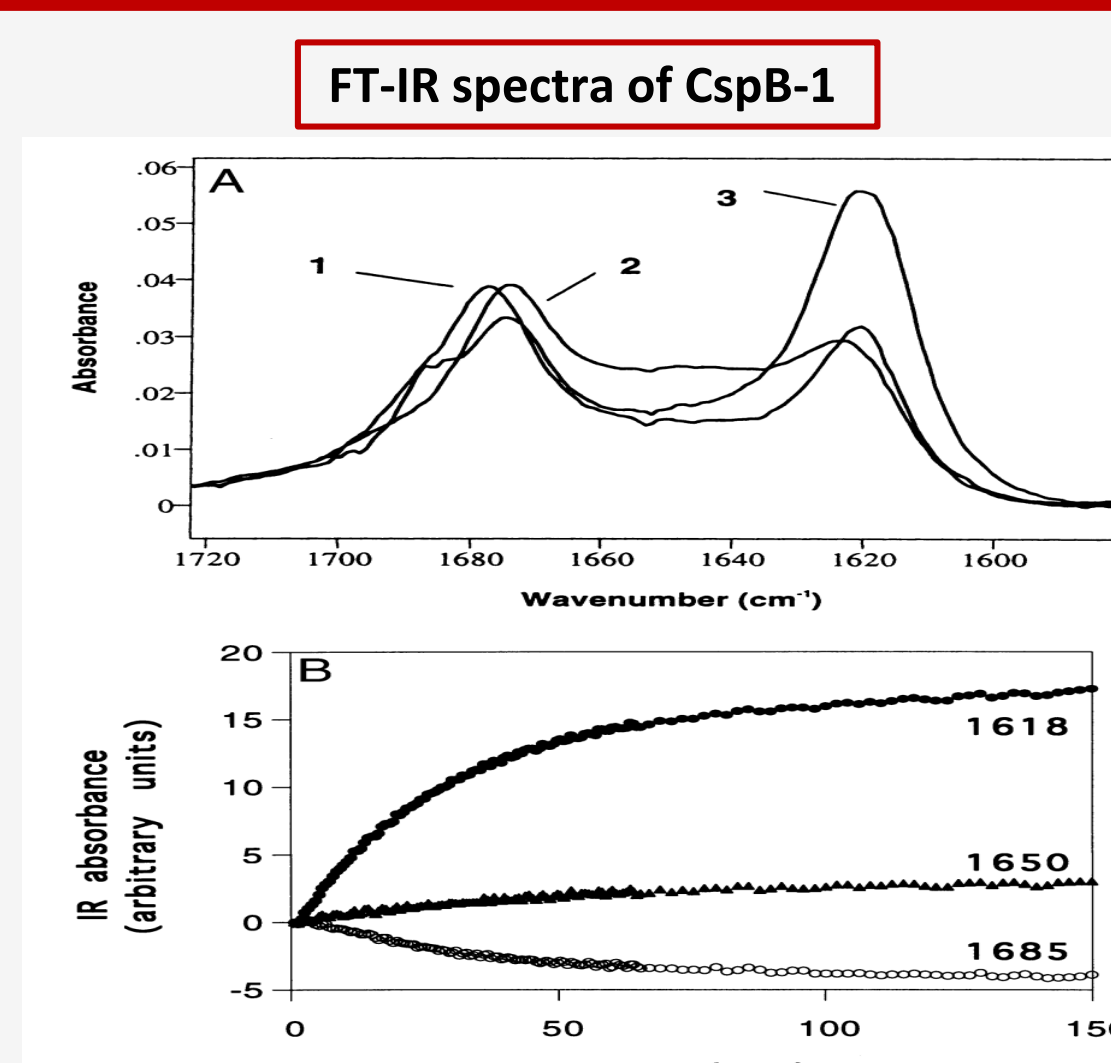
- ❖ Amyloid fibrils are highly ordered protein aggregates.
- ❖ They play a crucial role in the initiation and progression of various neurodegenerative disorders, including Alzheimer's disease, Parkinson's disease, type 2 diabetes.
- ❖ Under certain conditions these disease proteins can undergo structural rearrangements resulting in misfolded proteins that can lead to the formation of aggregates with a fibrillar amyloid-like structure



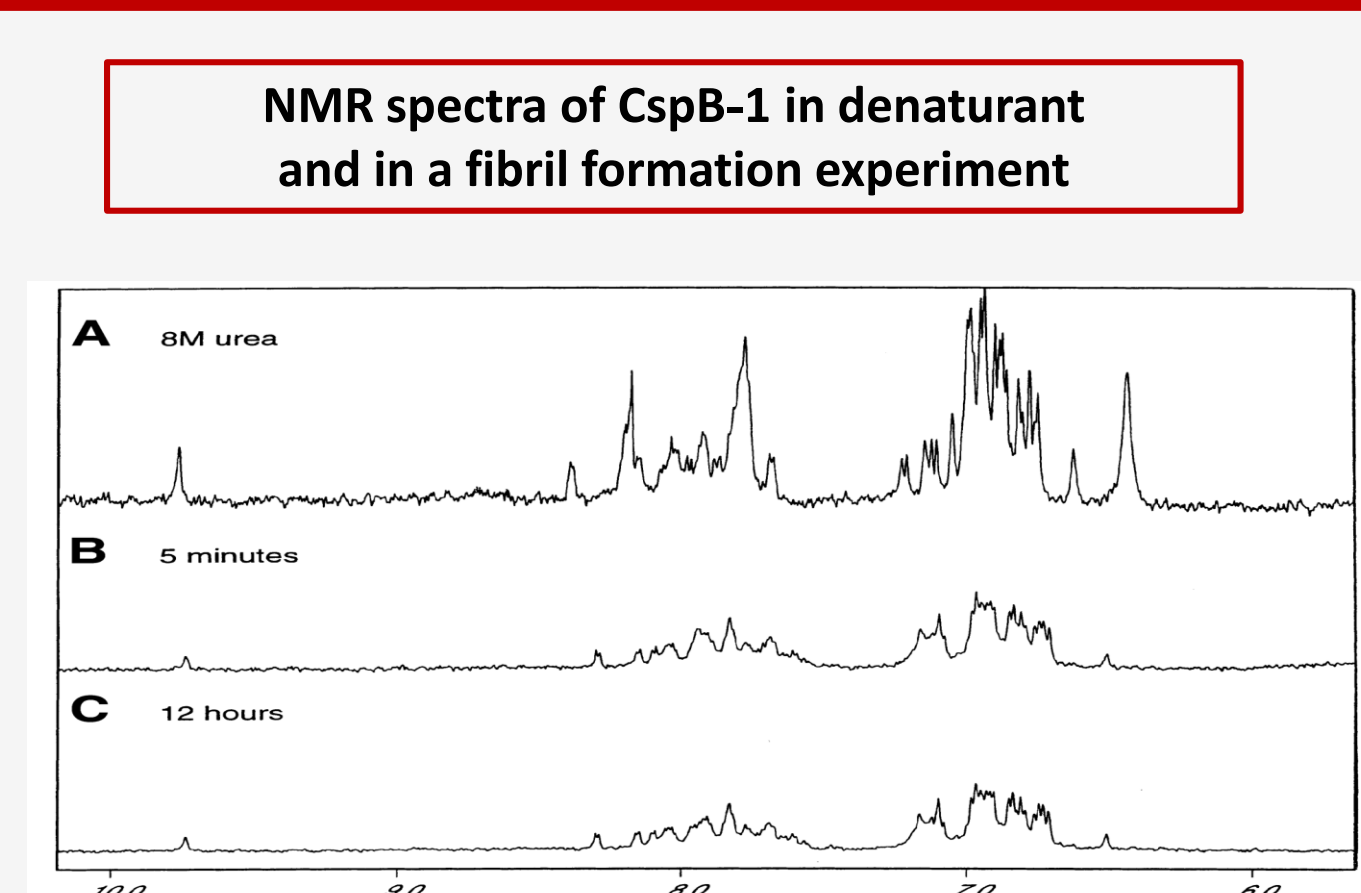
- ❖ The excellent agreement between the four techniques confirms that the process of fibril elongation along a surface is equivalent to the elongation in bulk solution
- ❖ the plots illustrate that the observed screening effects have no detectable temperature dependence in the investigated temperature rang



(A) a kinetic CD experiment over 4 h
(B) typical timecourse of a CD experiment recording the ellipticity at 215 nm over 5 h

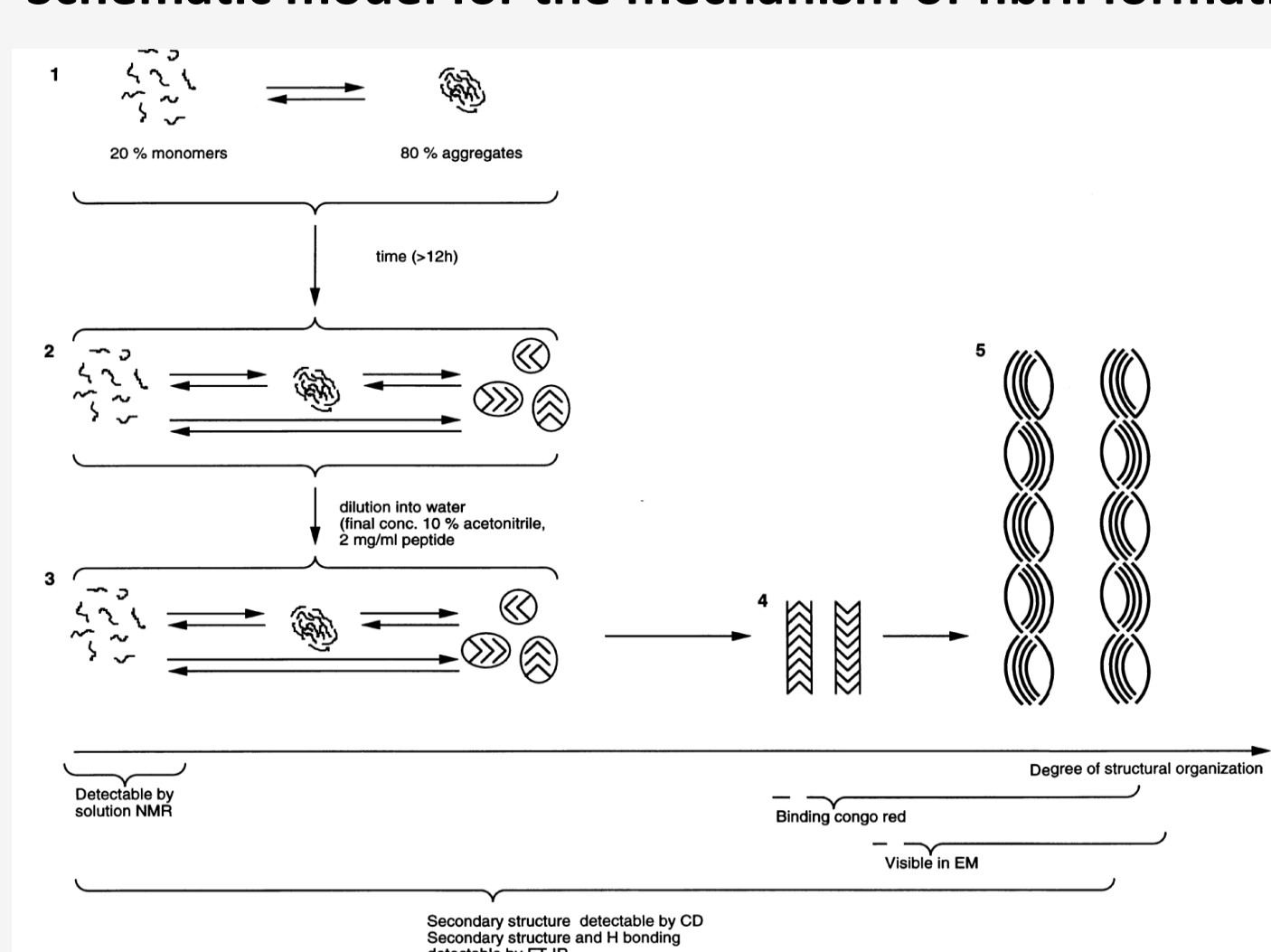


(A) FT-IR spectra of CspB-1
(B) timecourses derived from the FT-IR intensity changes at three different wavelengths



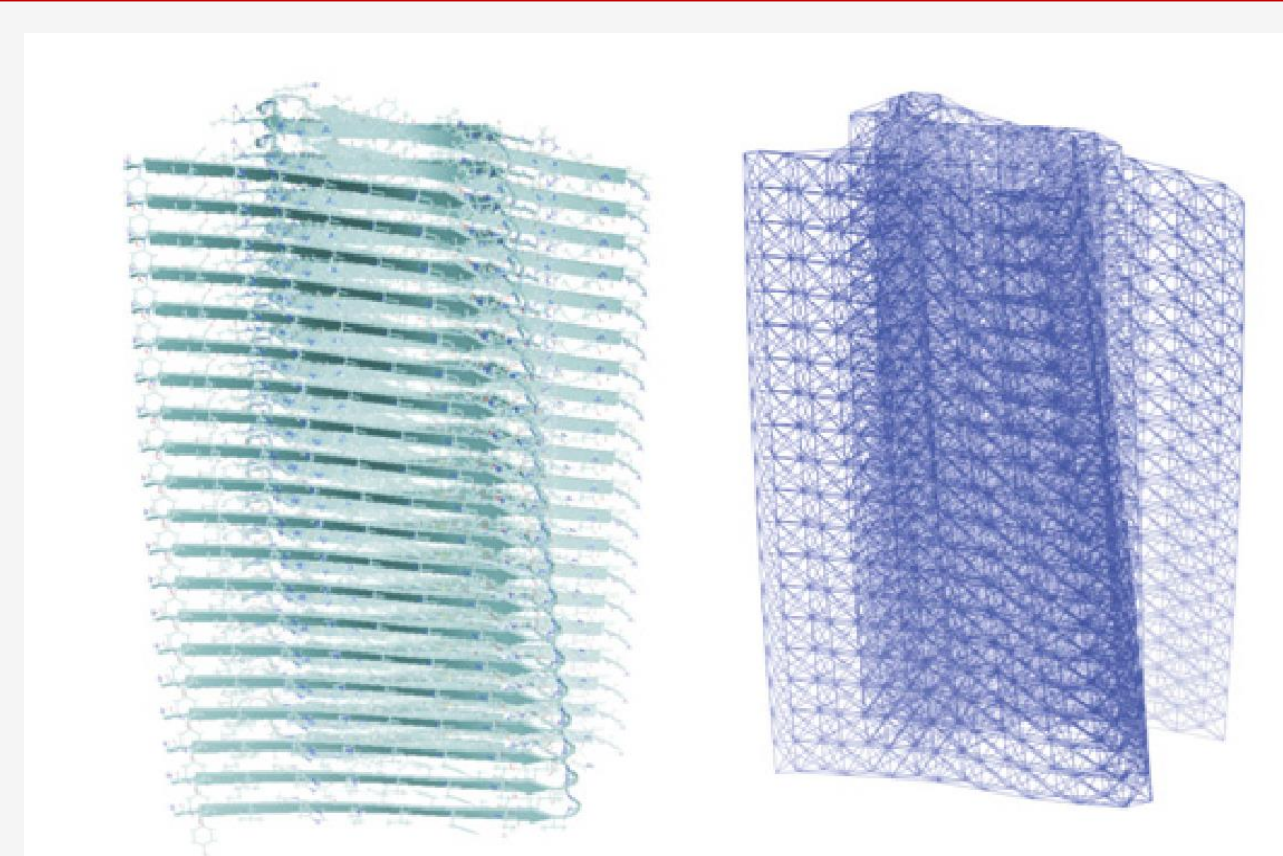
(A) Spectrum of CspB-1 in 8M urea
(B) spectrum recorded 5 min after initiation of fibril formation
(C) spectrum from the same experiment after 12 h

Schematic model for the mechanism of fibril formation

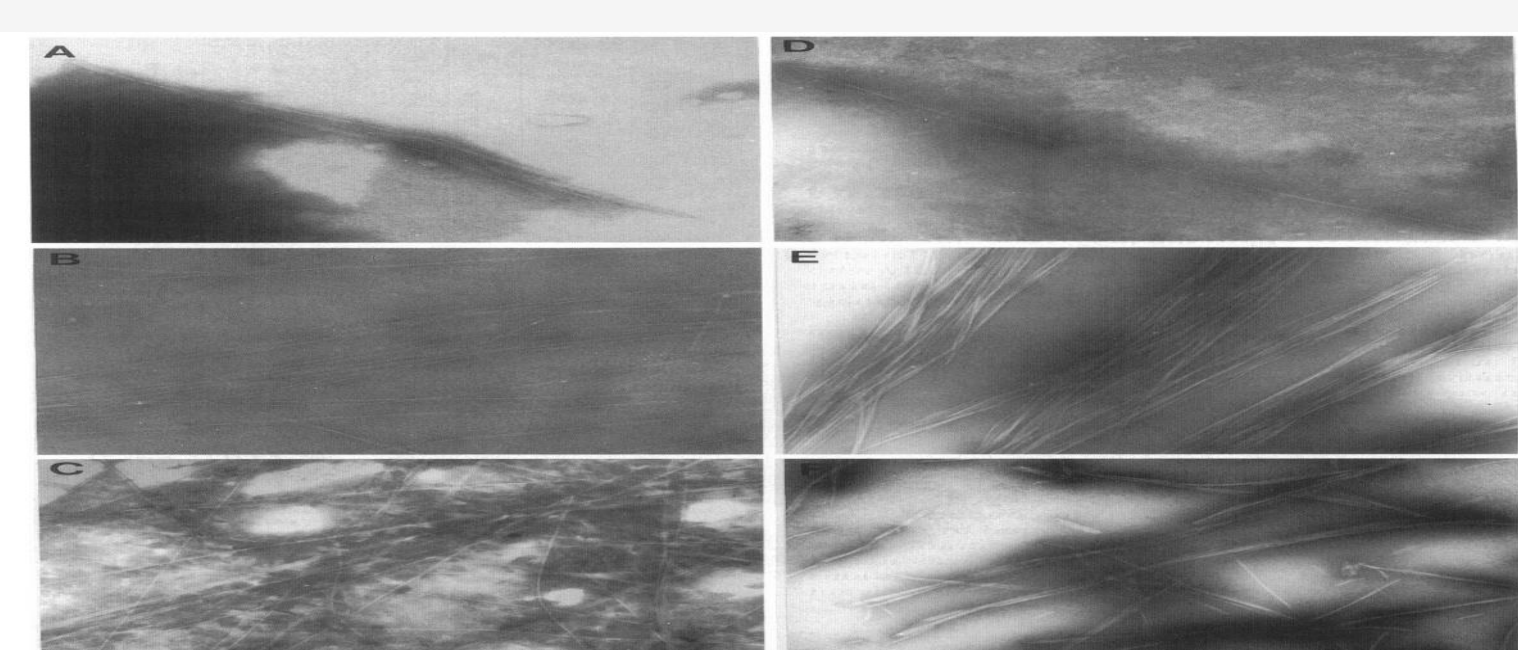


- (1) Unstructured monomers and disordered aggregates are present in solution after dissolving the peptide in 50% CH₃CN.
- (2) After the 12 h ageing step, the species initially present coexist with ordered aggregates.
- (3) Following the solvent shift to 10% CH₃CN, some of the ordered aggregates redissolve, but there are still the same three species in solution
- (4) complete fibrils.

Full-atomistic and corresponding ENM representations of a twofold symmetric amyloid fibril

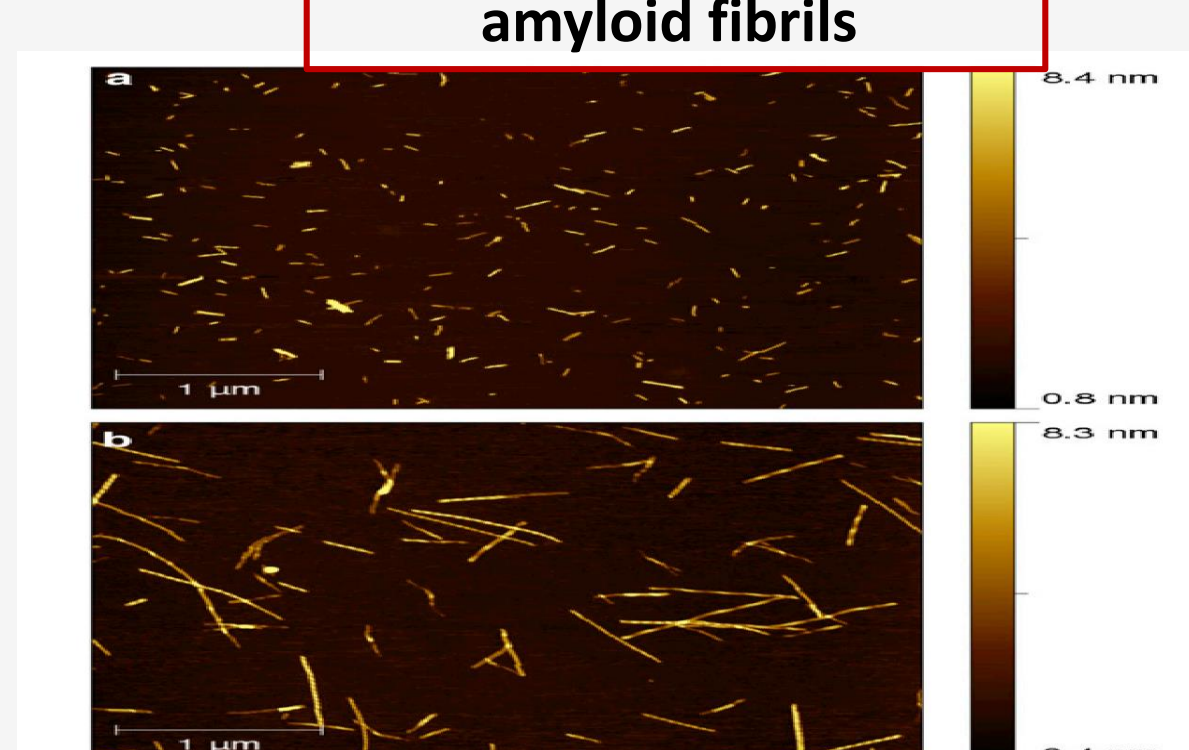


Electron micrographs of B(1-28)



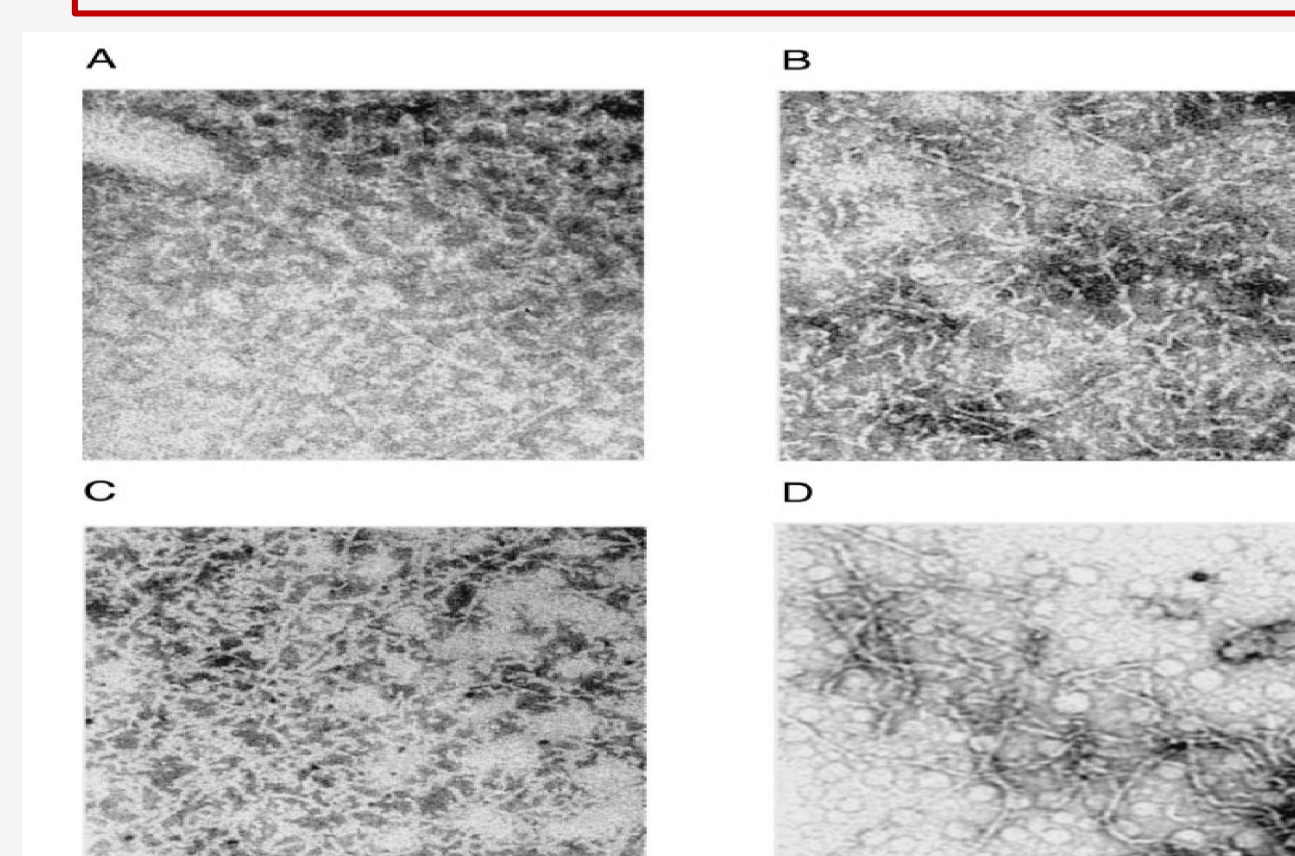
(A) 1 hour phosphate buffer, (B) 21 hours phosphate buffer
(C) 3 month phosphate buffer, (D) 3 hours in PBS
(E) 7 weeks in PBS, (F) pH= 4.4

AFM images of insulin amyloid fibrils



dees detacinoS (a) fibrils
(b) Elongatd fibrils

Electron micrographs of CspB-1 amyloid fibrils observed



(A) 30 min
(B) 67 min
(C) 137 min
(D) 1 week

- ❖ electrostatic effects control protein aggregation
- ❖ longer amyloid fibrils are more stable
- ❖ the formation of amyloid fibrils are a generic property of polypeptide chains
- ❖ magnitude of binding of a variety of ions to protein molecules

Discussion

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3. Shen, C.-L., et al., *Light scattering analysis of fibril growth from the amino-terminal fragment beta (1-28) of beta-amyloid peptide*. Biophysical journal, 1993. 65(6): p. 2115-2121.
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