

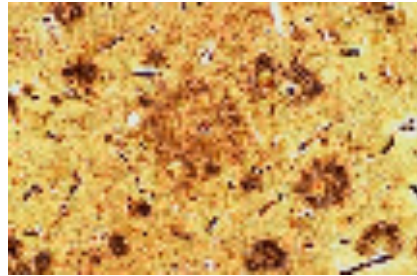
The role of amyloid beta peptide variability toward progress of Alzheimer disease

Kerensa Broersen
Brain Disorders and Therapeutics
London, August 26



Alzheimer's disease

Amyloid β peptide



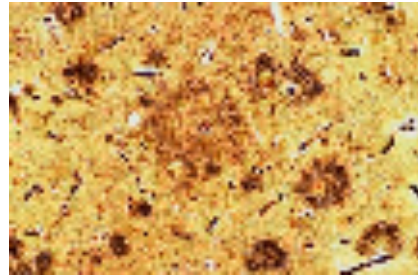
Alzheimer's disease

Amyloid β peptide

Genetic preposition

Neuroinflammation

Tau



Apolipoprotein E isoform

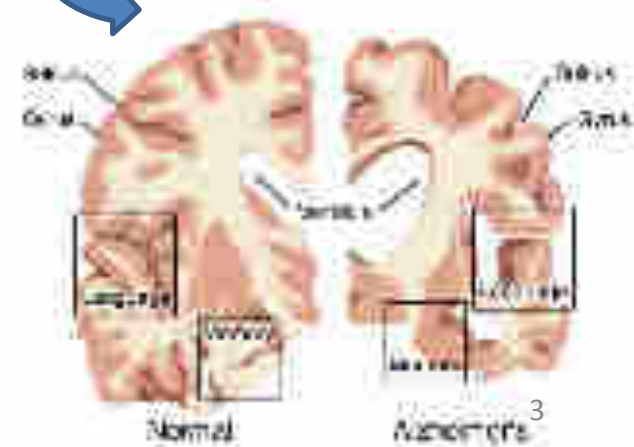


Mitochondrial dysfunction

Aging

Environmental factors

Hyperphosphorylated tau



The aggregation process of A β is characterized by intermediates that may have toxic activity

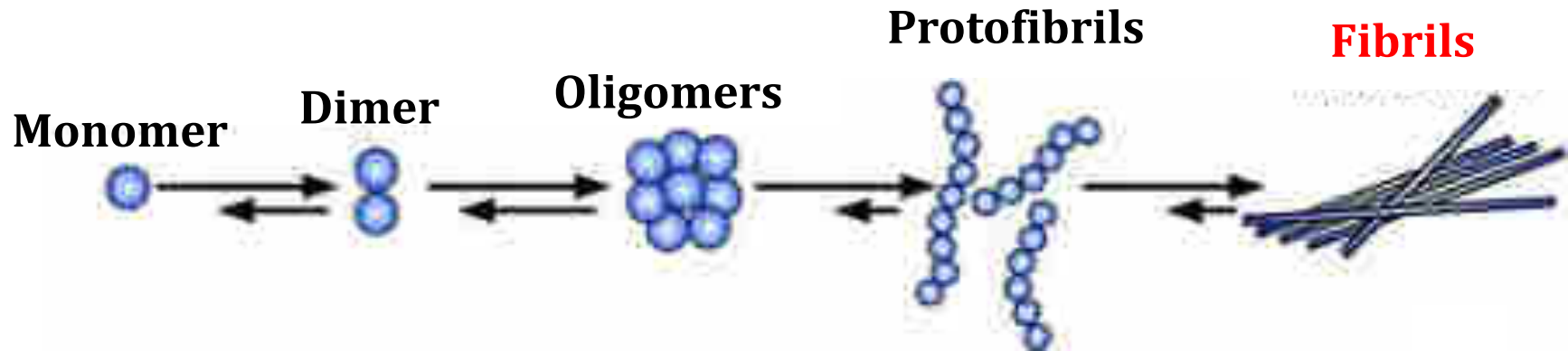


Figure adapted from Kumar *et al*
EMBO J (2011)

The aggregation process of A β is characterized by intermediates that may have toxic activity

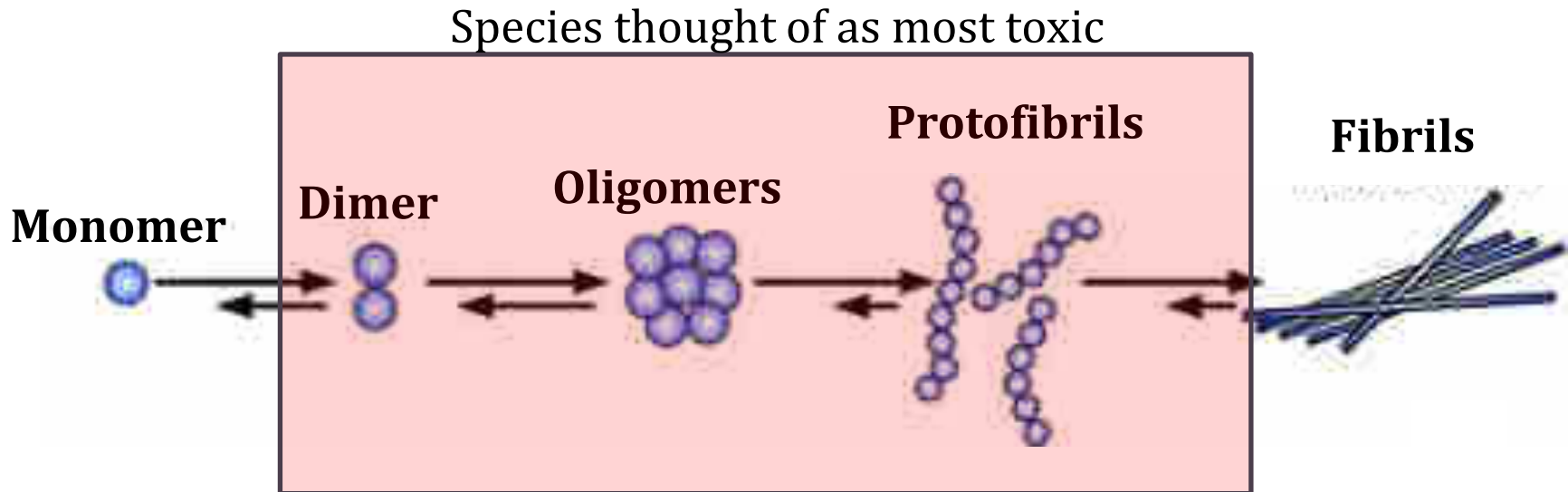


Figure adapted from Kumar *et al*
EMBO J (2011)

A β variation – result of heterogeneous γ -secretase activity

A β 37

A β 38

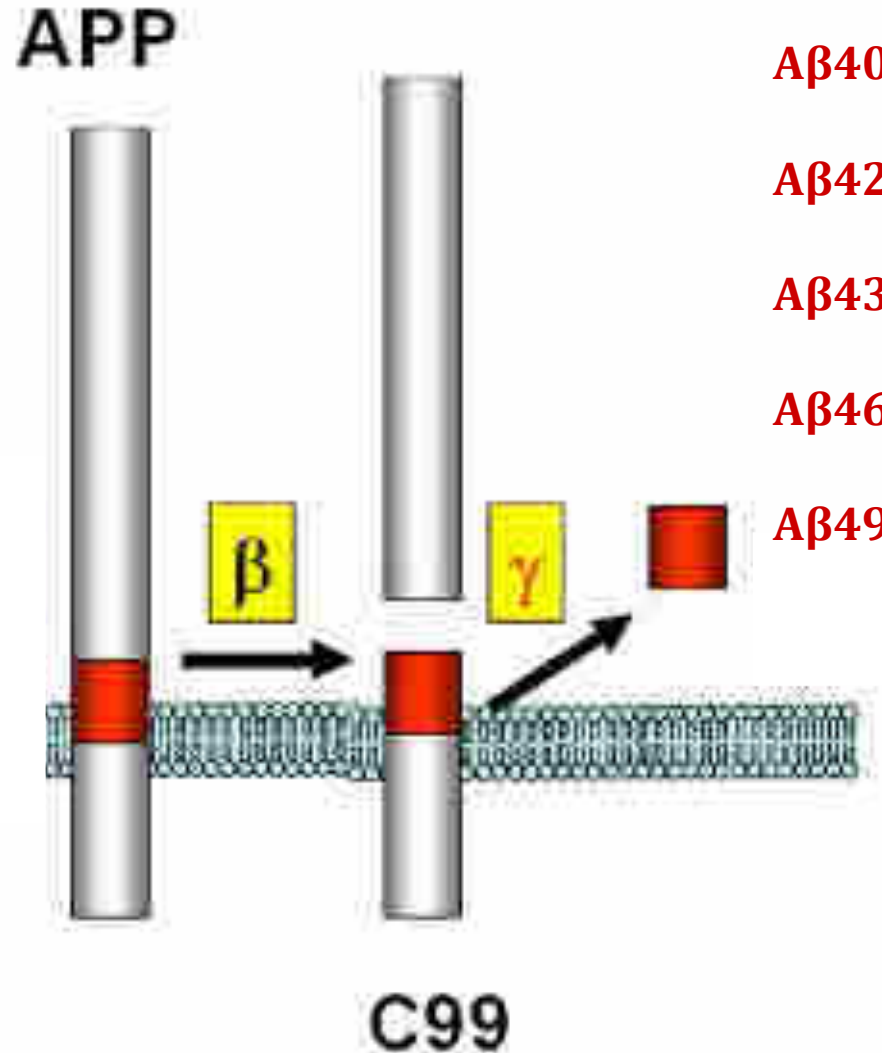
A β 40

A β 42

A β 43

A β 46

A β 49



A β variation – result of heterogeneous γ -secretase activity **A β 37**

A β 38

~ 90%

A β 40

A β 42

~ 10%

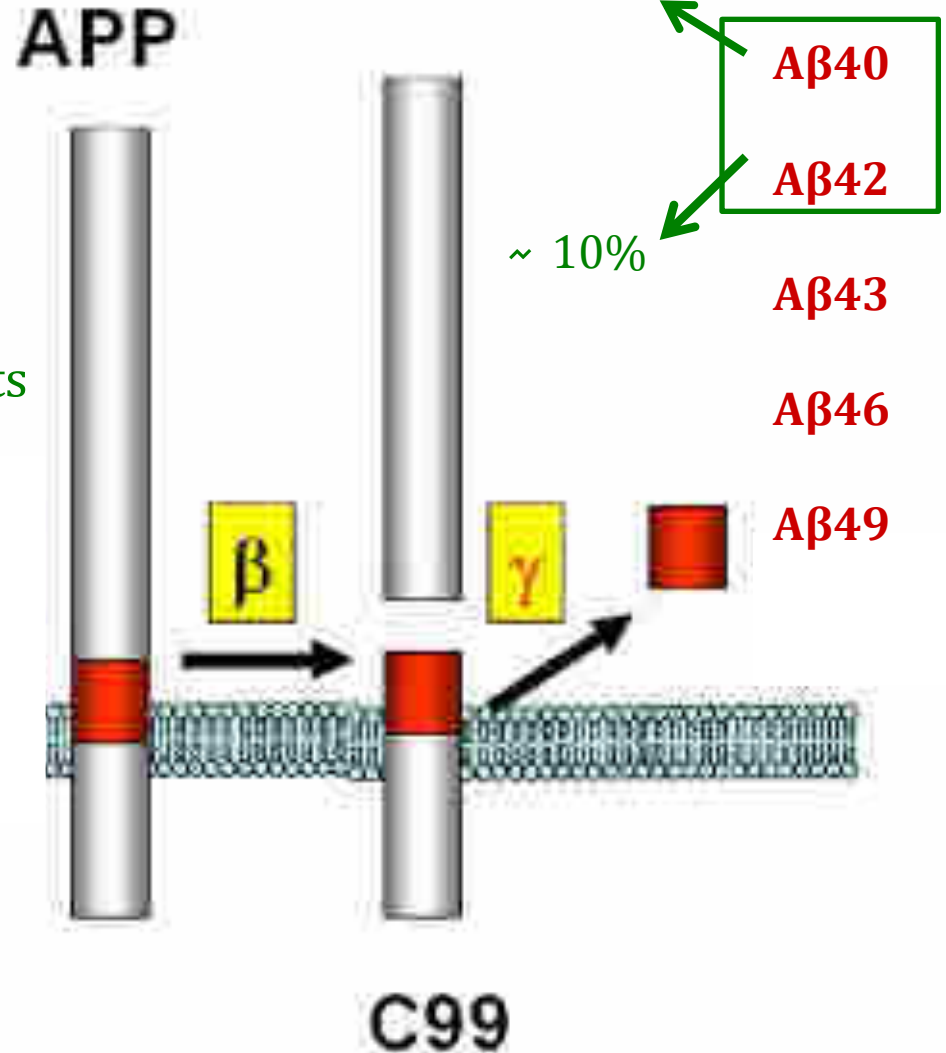
A β 43

A β 46

A β 49

A β production in healthy aged subjects

A β 42: A β 40 = 1:9



A β variation – result of heterogeneous γ -secretase activity **A β 37**

A β 38

~ 90%

A β 40

A β 42

~ 10%

A β 43

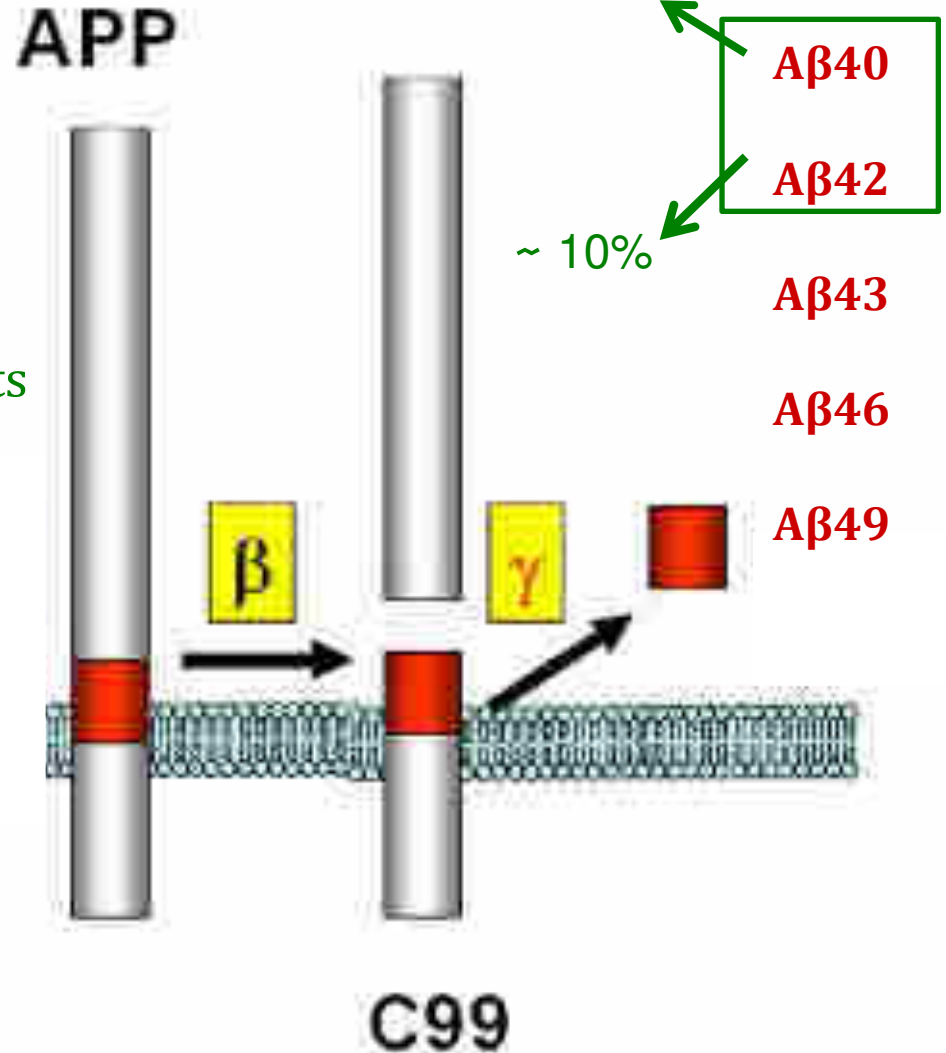
A β 46

A β 49

A β production in healthy aged subjects

A β 42: A β 40 = 1:9

Highly amyloidogenic



A β variation – result of heterogeneous γ -secretase activity **A β 37**

A β 38

~ 90%

A β 40

A β 42

~ 10%

A β 43

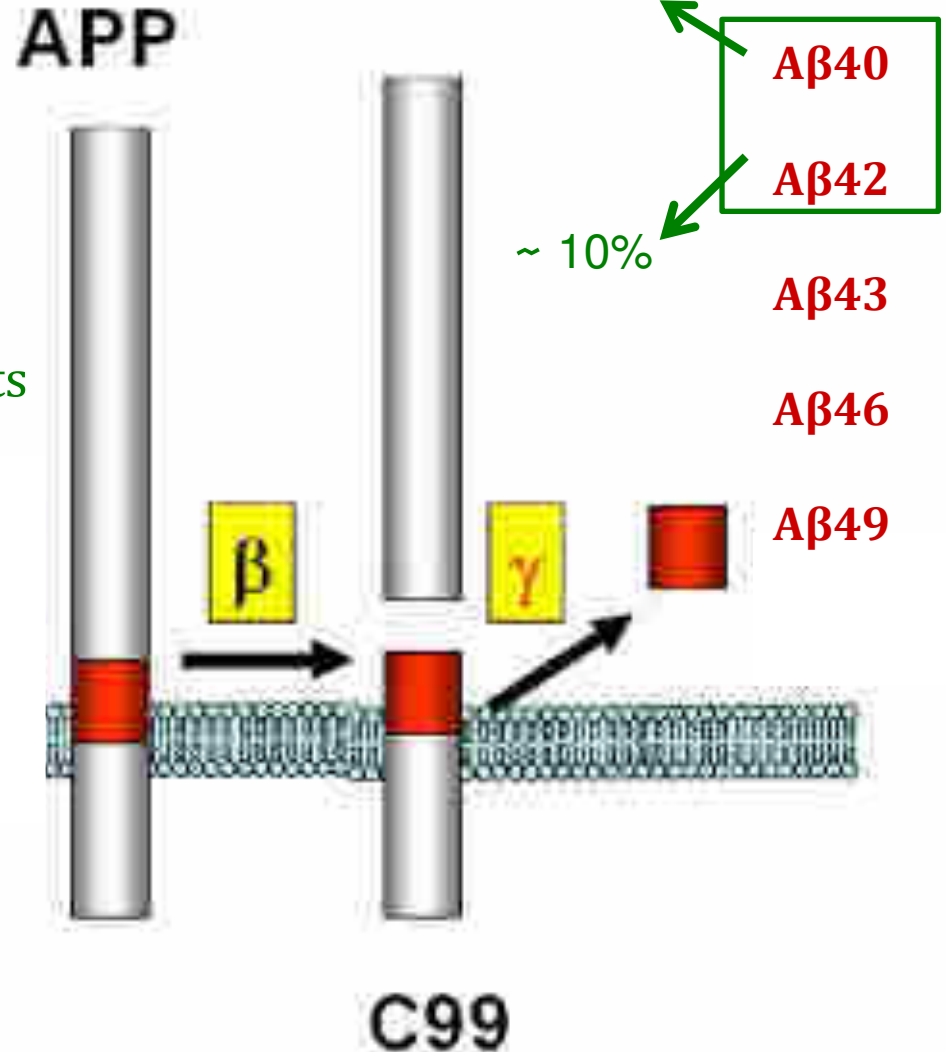
A β 46

A β 49

A β production in healthy aged subjects

A β 42: A β 40 = 1:9

Little amyloidogenic



A β variation – result of heterogeneous γ -secretase activity **A β 37**

A β 38

~ 70%

A β 40

A β 42

~ 30%

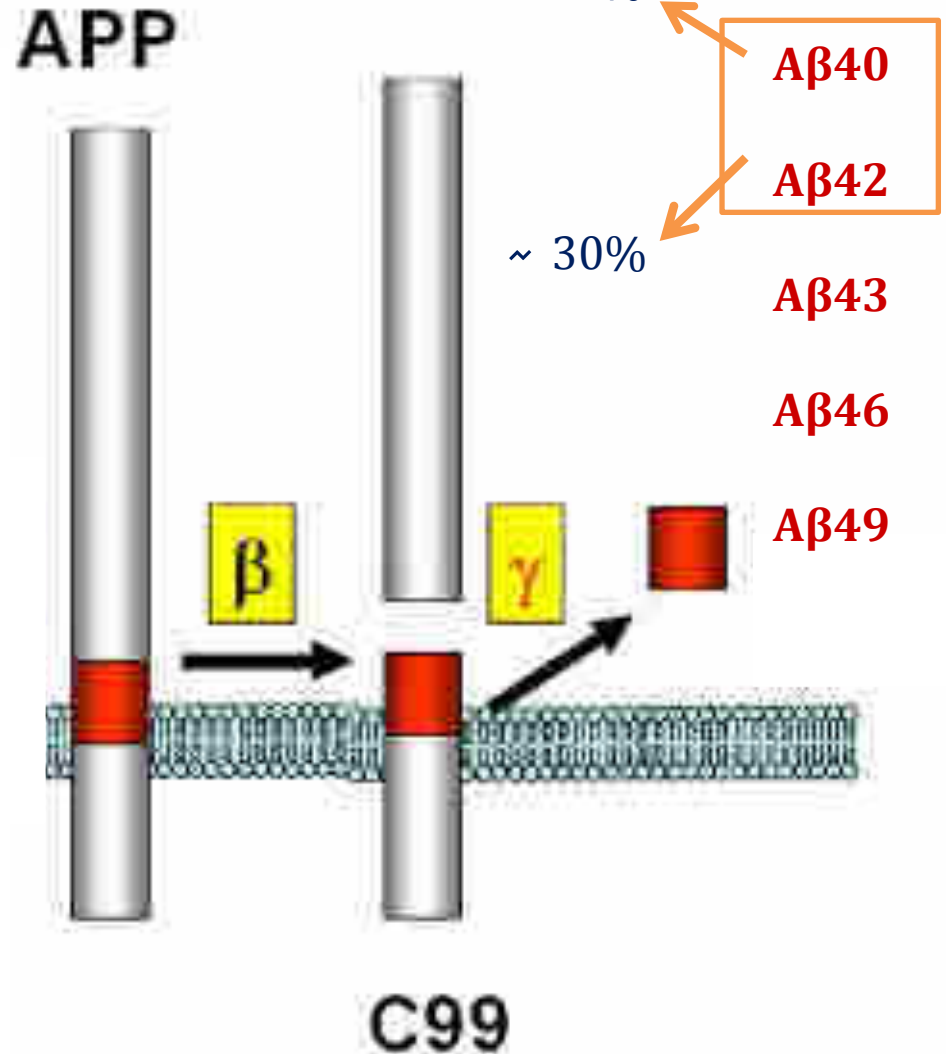
A β 43

A β 46

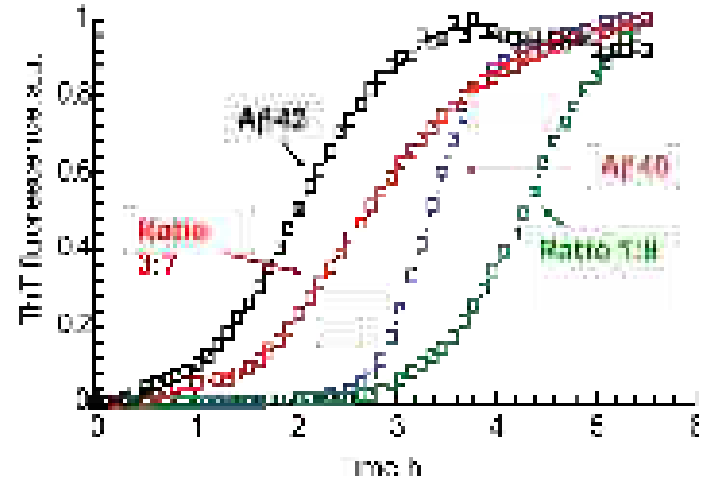
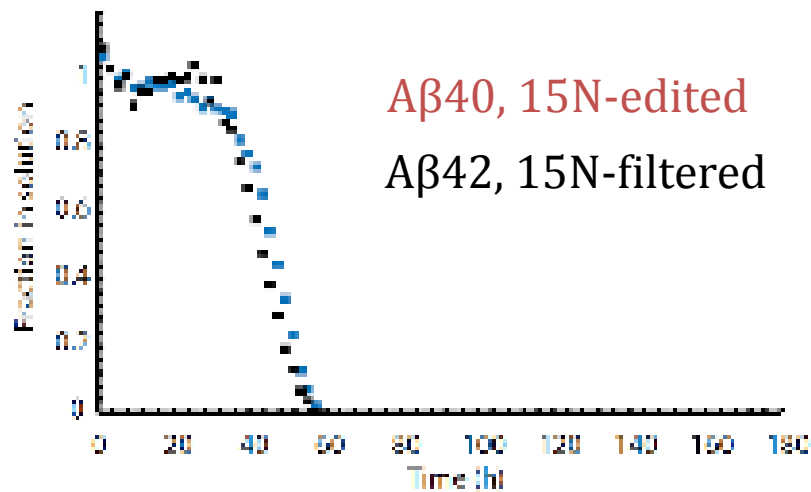
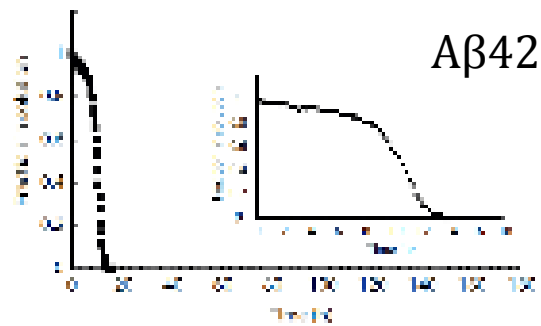
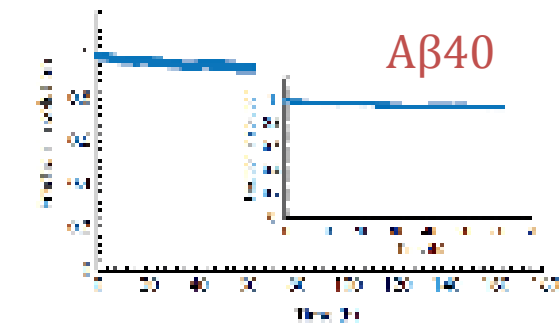
A β 49

A β production in aged subjects with Alzheimer pathology

$$A\beta_{42}:A\beta_{40} = 3:7$$

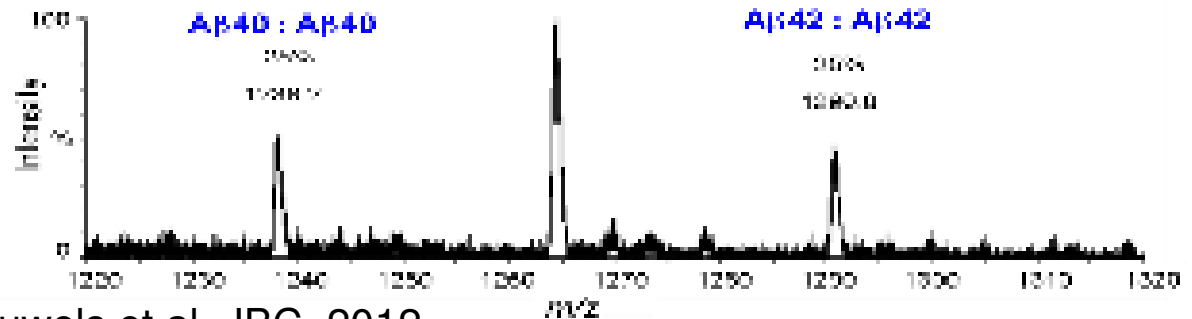


A β heterogeneity affects aggregation



A β 40 : A β 42
A β 42 : A β 40

50%
100%

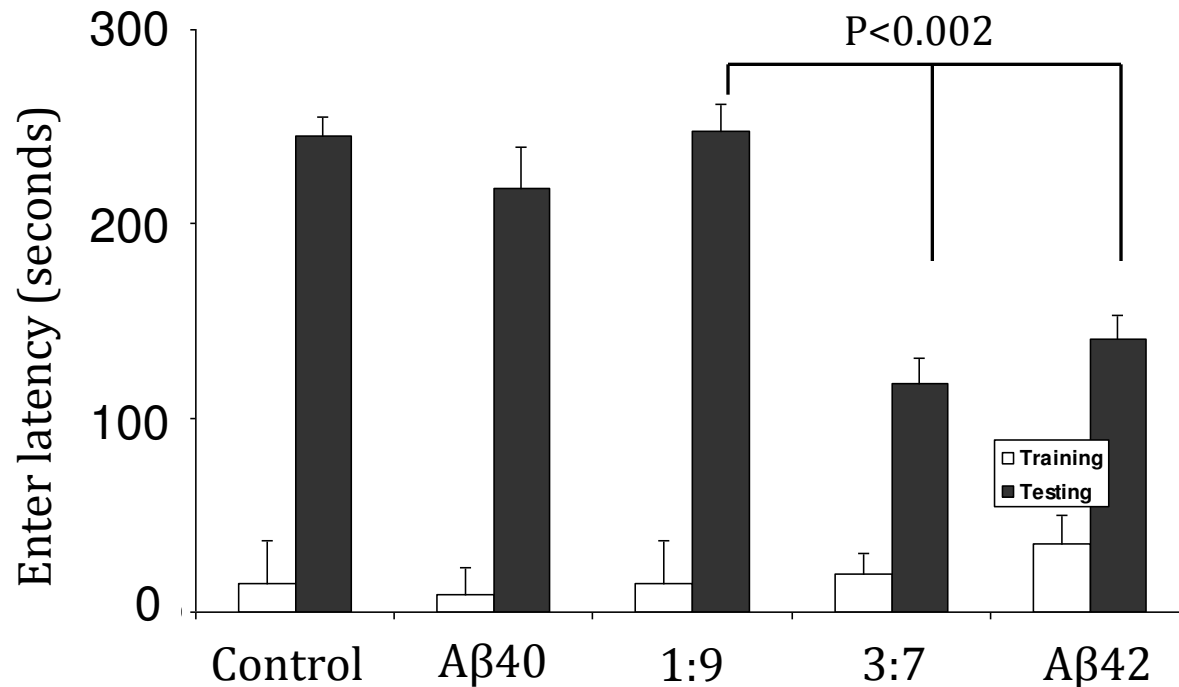
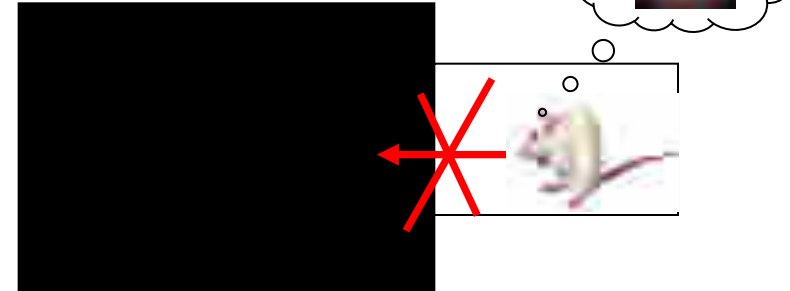


A β heterogeneity affects toxicity

Day 1-training (shock)



Day 2-memory testing

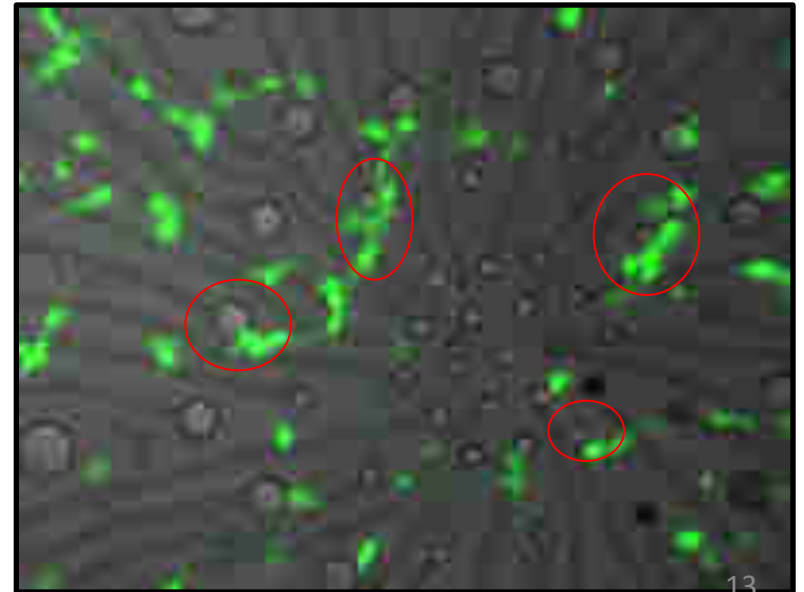
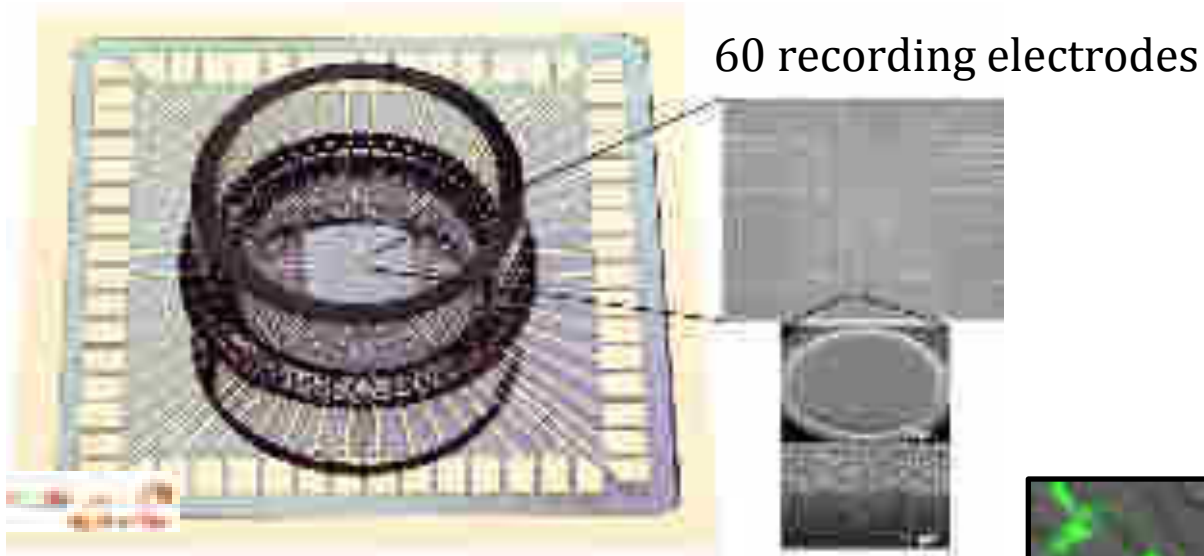


3:7 A β 42:A β 40 ratio
causes memory
and learning impairment in
mice

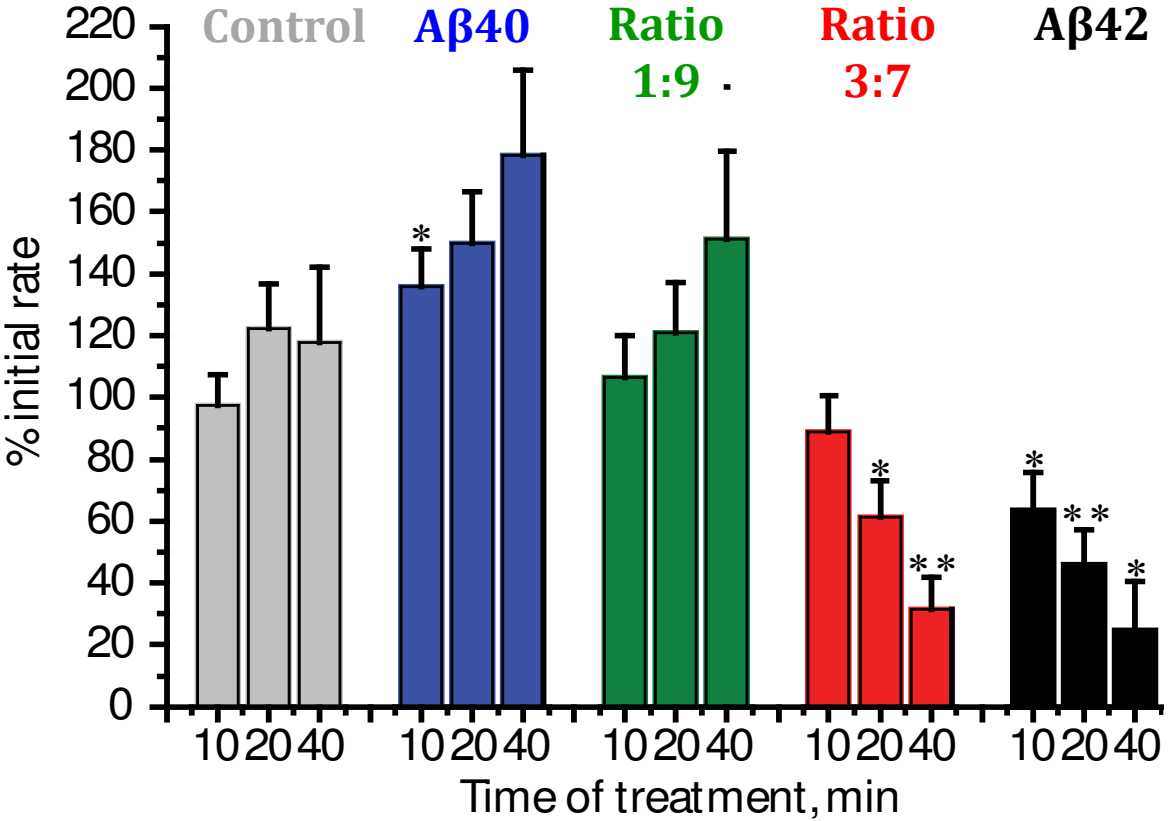
Passive avoidance response

A β heterogeneity affects toxicity

Microelectrode array: synaptic activity recordings in neuronal network grown on chip



Pathological mixtures of A β 40 and A β 42 induce acute synaptotoxicity



A β variation – result of heterogeneous γ -secretase activity

A β 37

A β 38

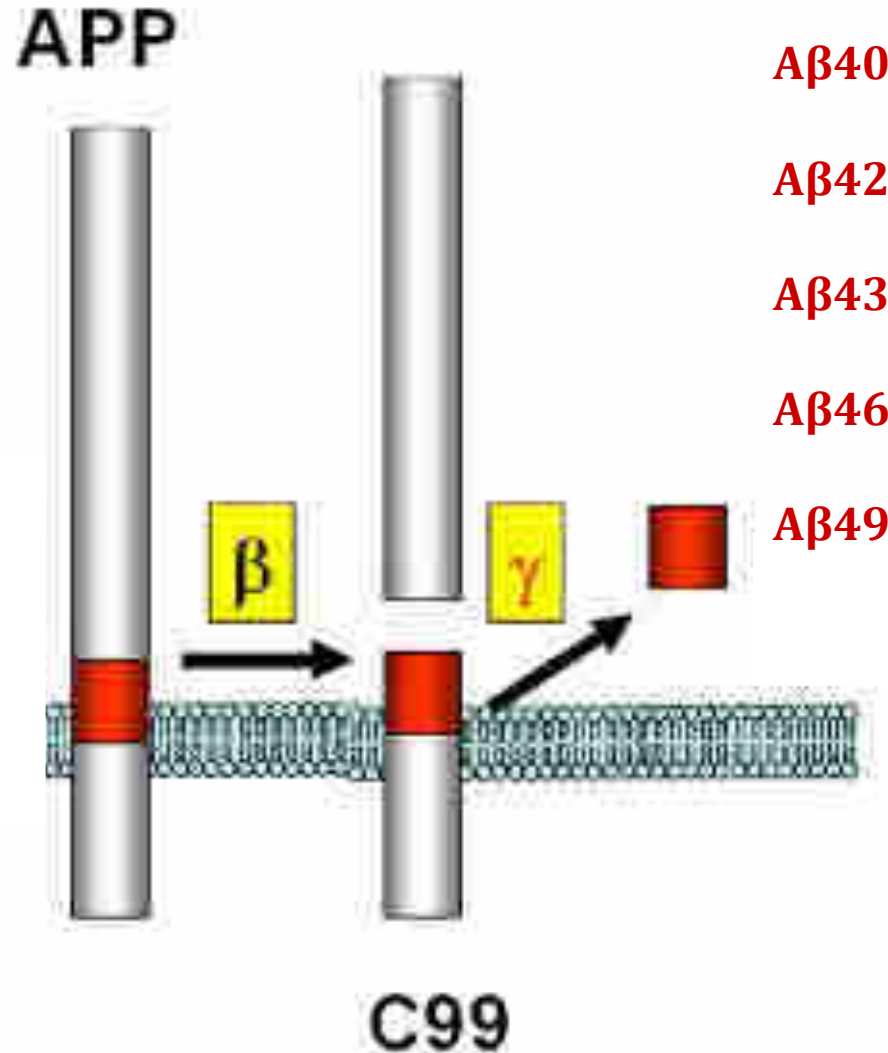
A β 40

A β 42

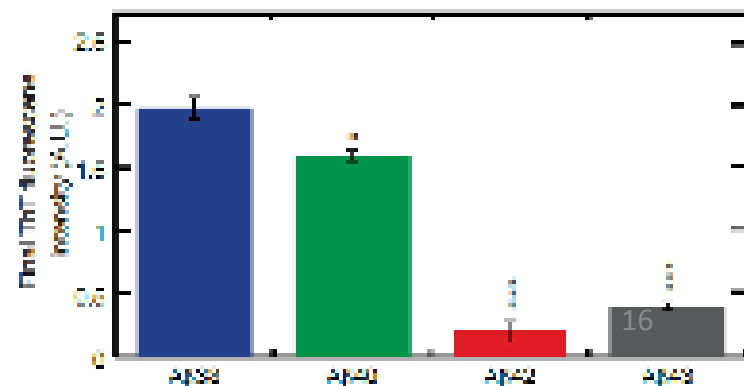
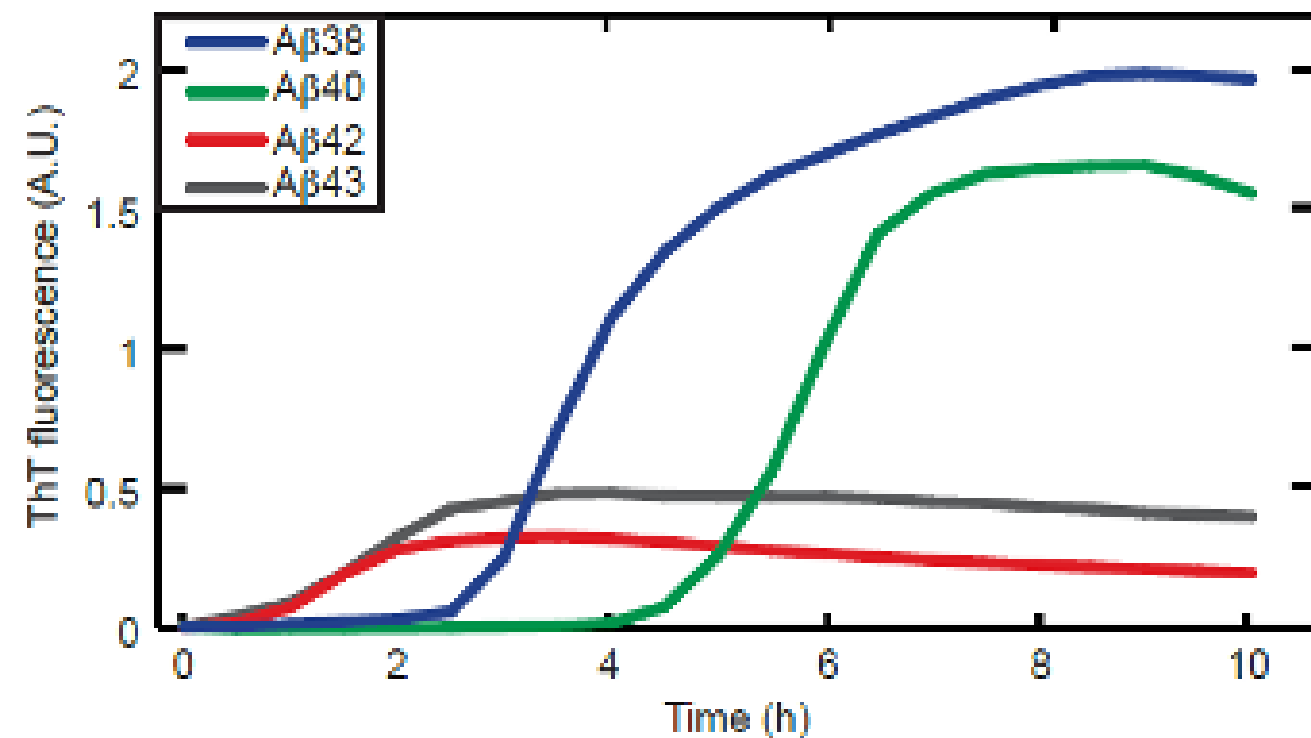
A β 43

A β 46

A β 49

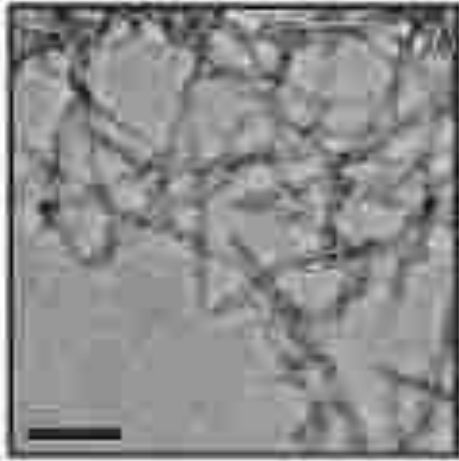


A β peptide length affects aggregation rate

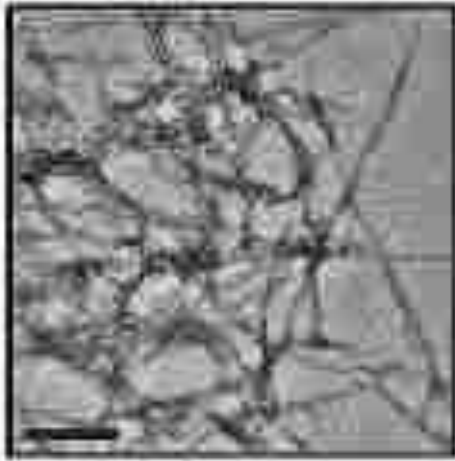


Tendency to form amyloid fibrils is independent of A β peptide length
... but morphological differences

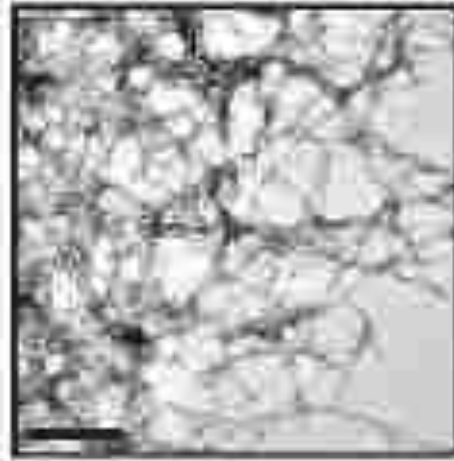
A β 38



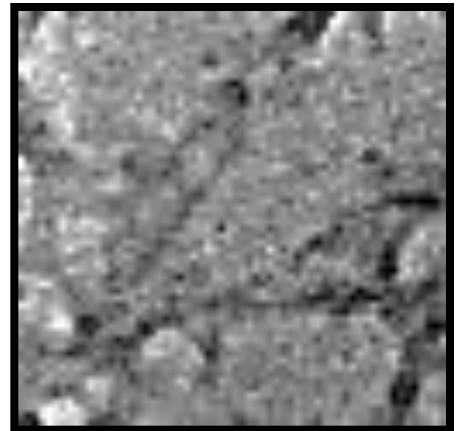
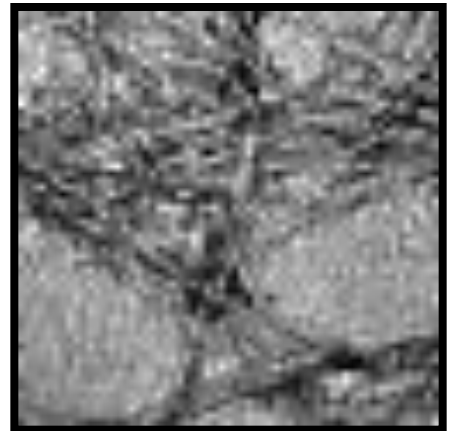
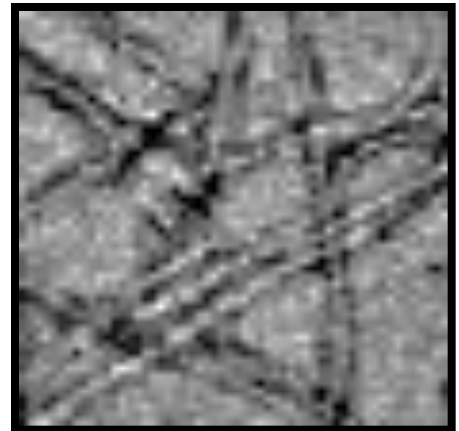
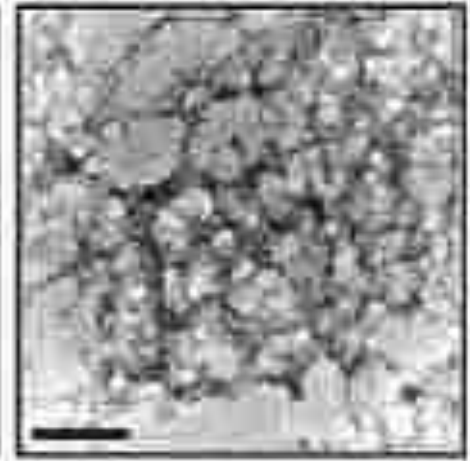
A β 40



A β 42



A β 43



All tested A β peptide lengths form oligomers but with different structural properties.

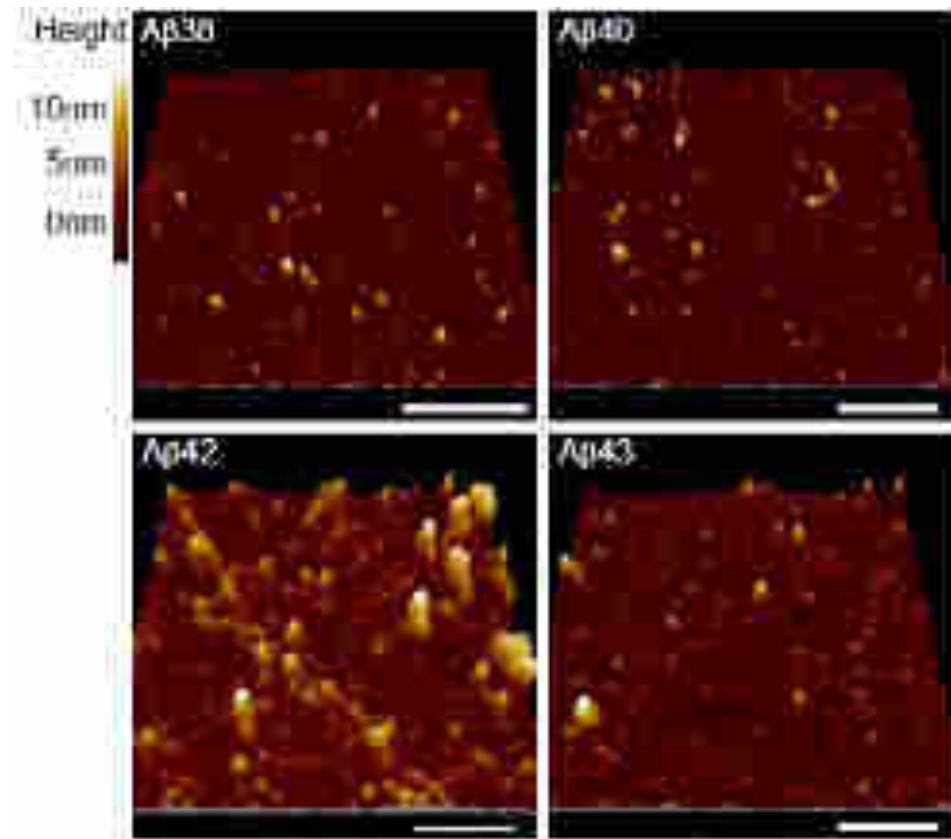
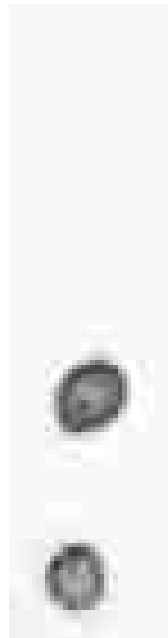
Recognition of oligomeric A β by **A11 antibody**.

A β 38

A β 40

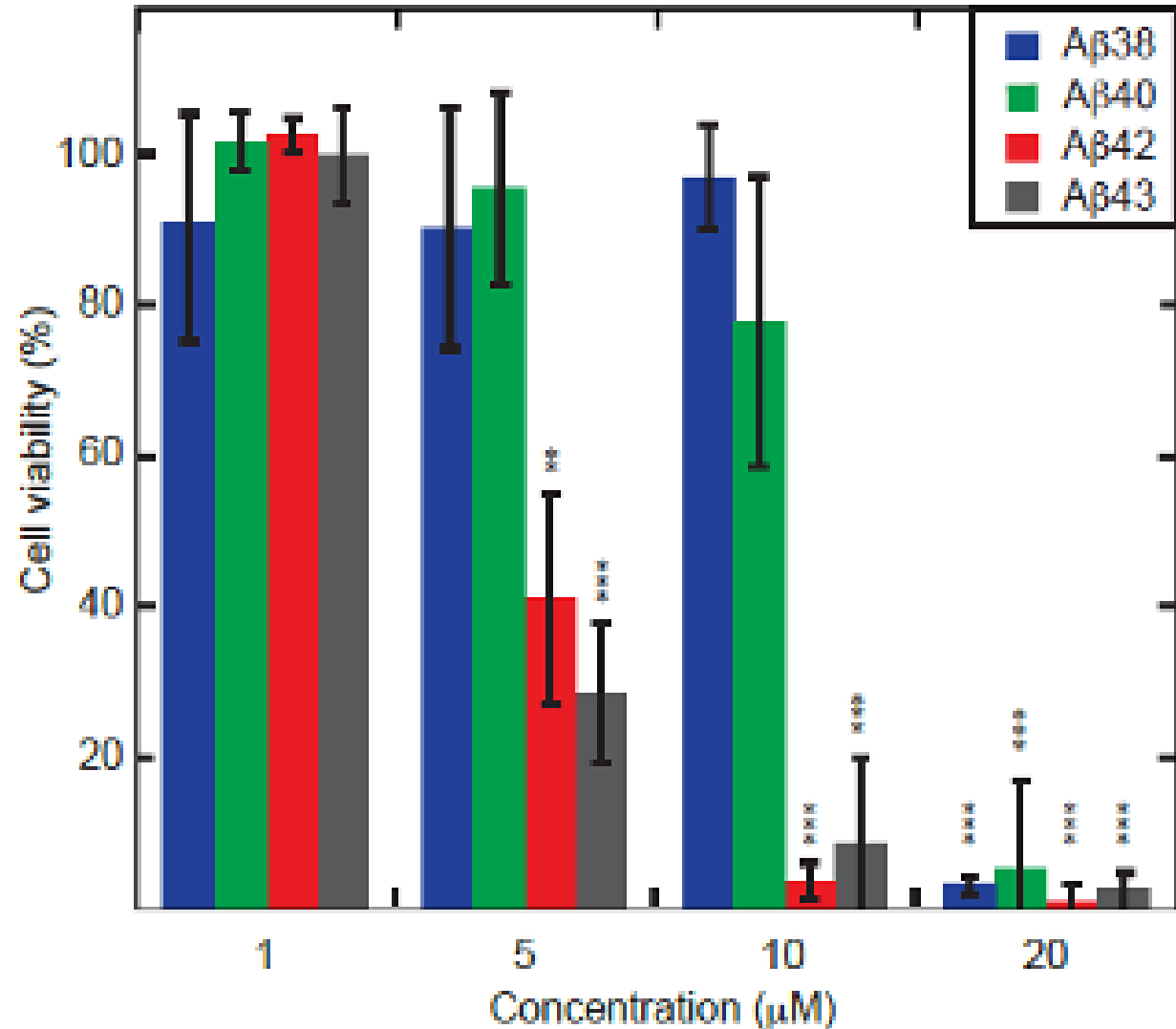
A β 42

A β 43

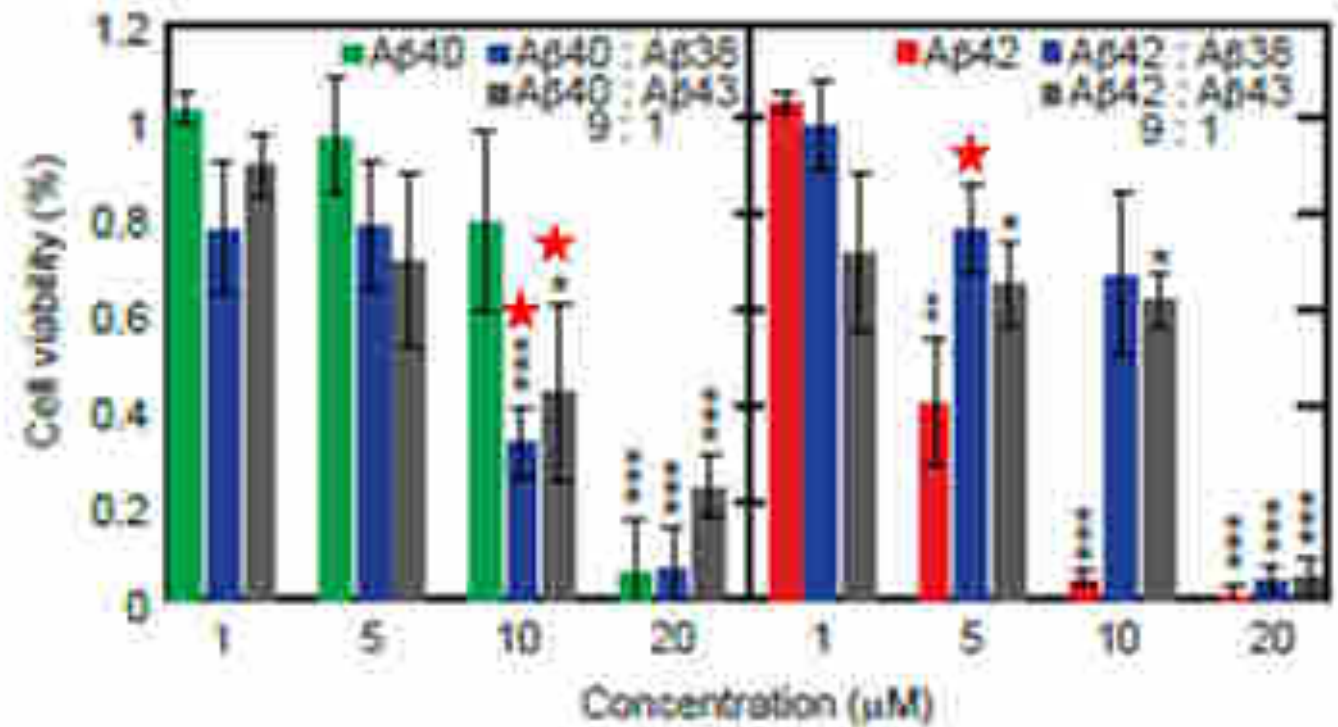
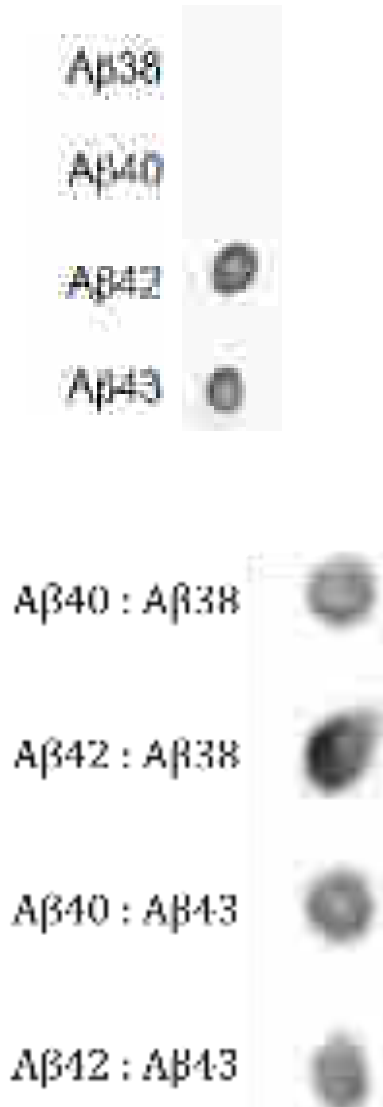


A β peptide length affects cytotoxicity: unexpected toxicity of A β 38

Cell titer blue **cell viability assay** of neuroblastoma SHSY-5Y cells by incubation with oligomeric A β .

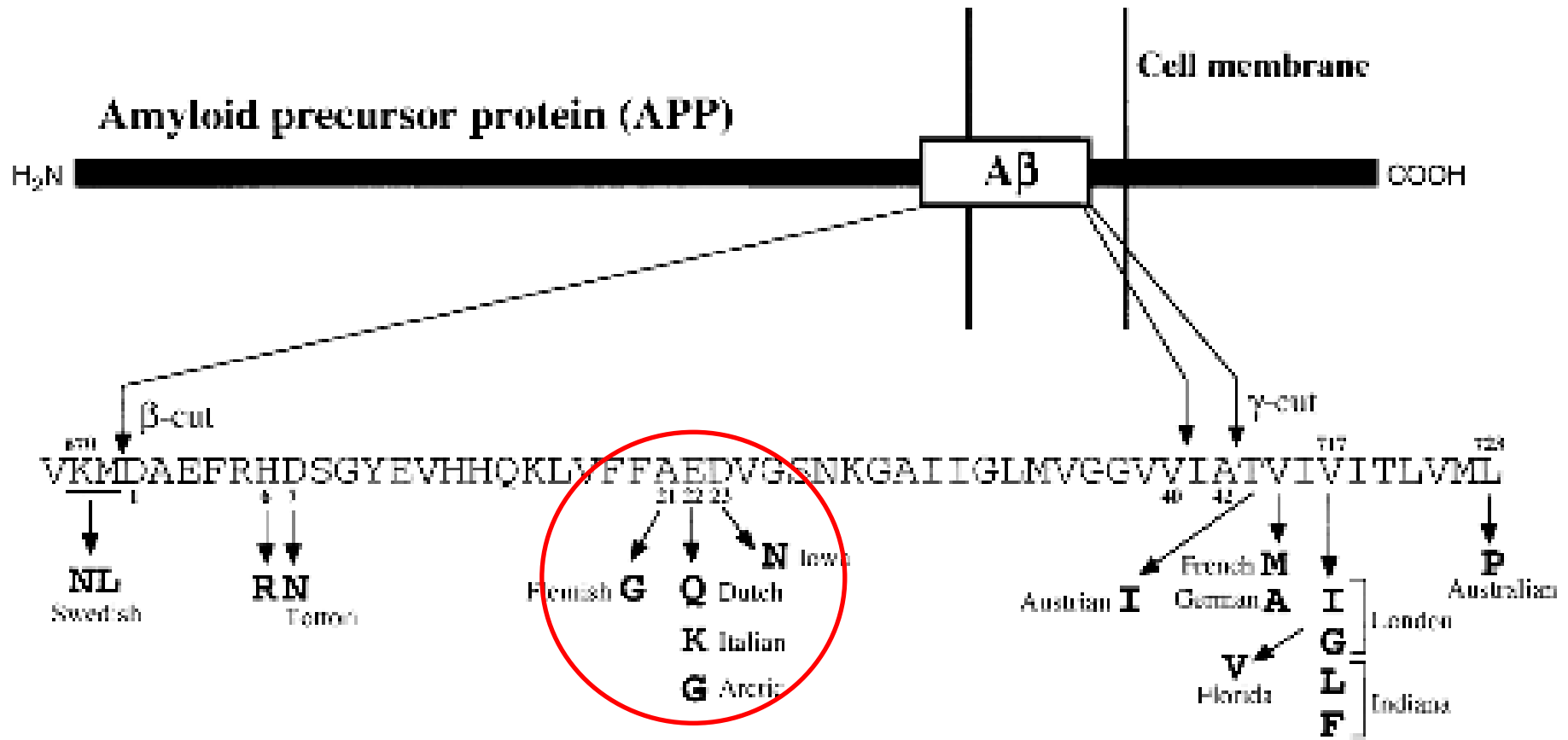


A β peptide length affects cytotoxicity: unexpected toxicity of A β 38

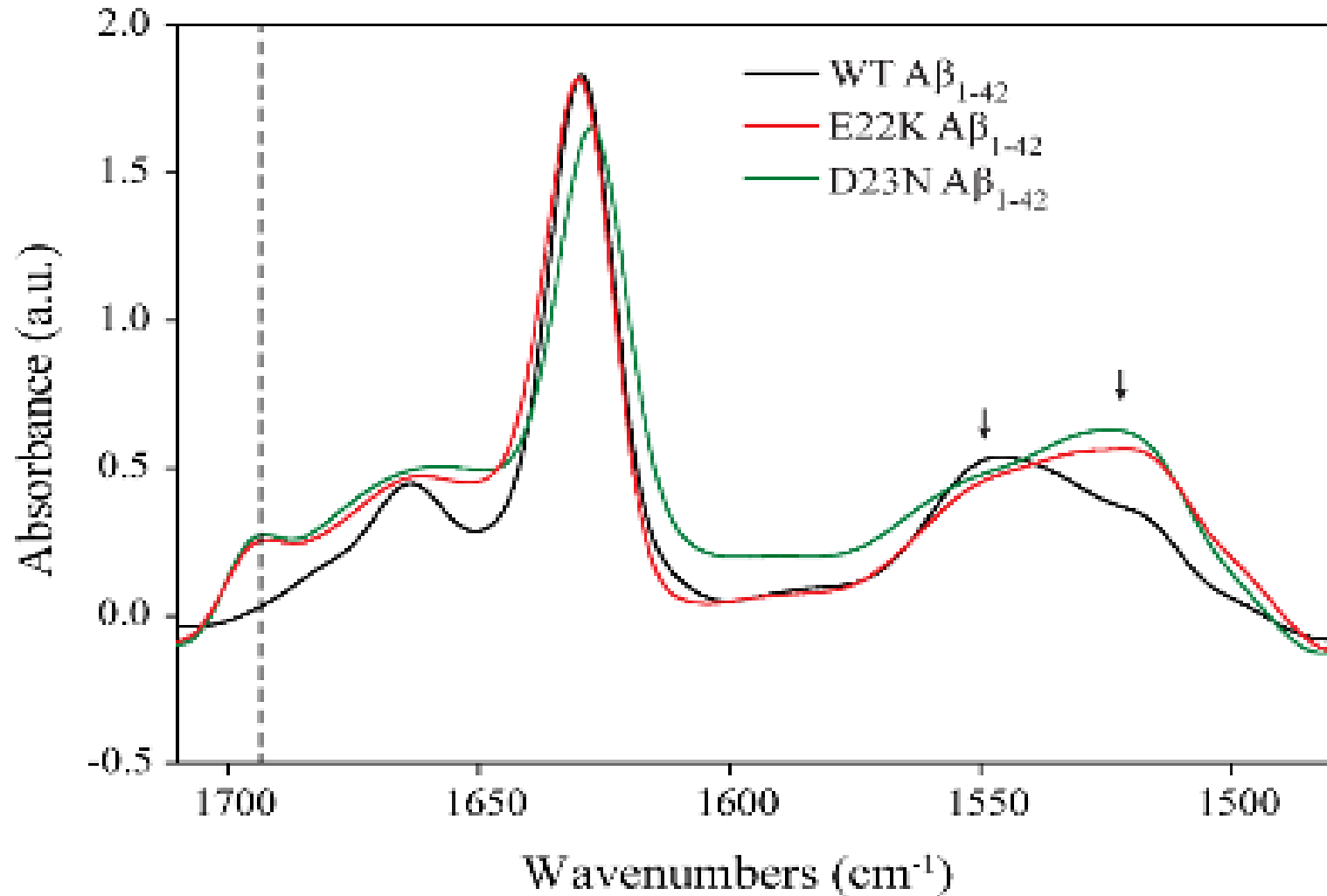


Cell titer blue **cell viability assay** of neuroblastoma SHSY-5Y cells by incubation with oligomeric A β .

A β mutations



D23N (Iowa mutant) and E22K (Italian mutant) A β 42 form antiparallel β -sheet fibrils



Alzheimer's disease – Cerebral amyloid angiopathy (CAA)

CAA is a common clinical symptom of early-onset familial AD

BRAIN

Amyloid-beta ($A\beta$) plaques



Bugiani et al 2010.

BLOOD

$A\beta$ deposits in cerebral blood vessel walls

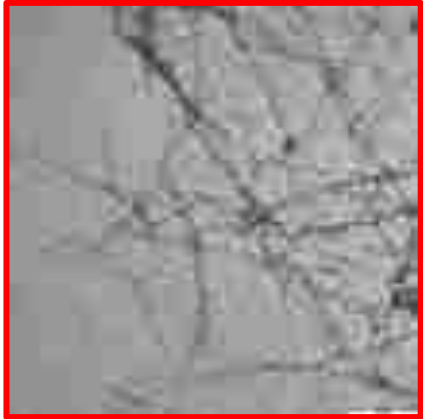
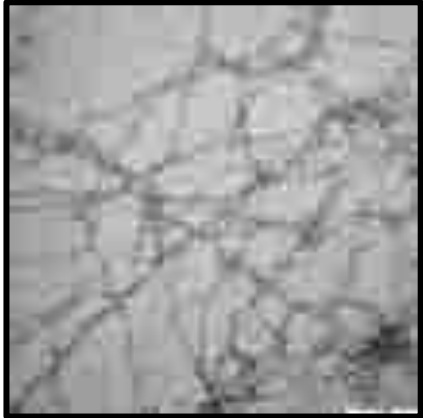
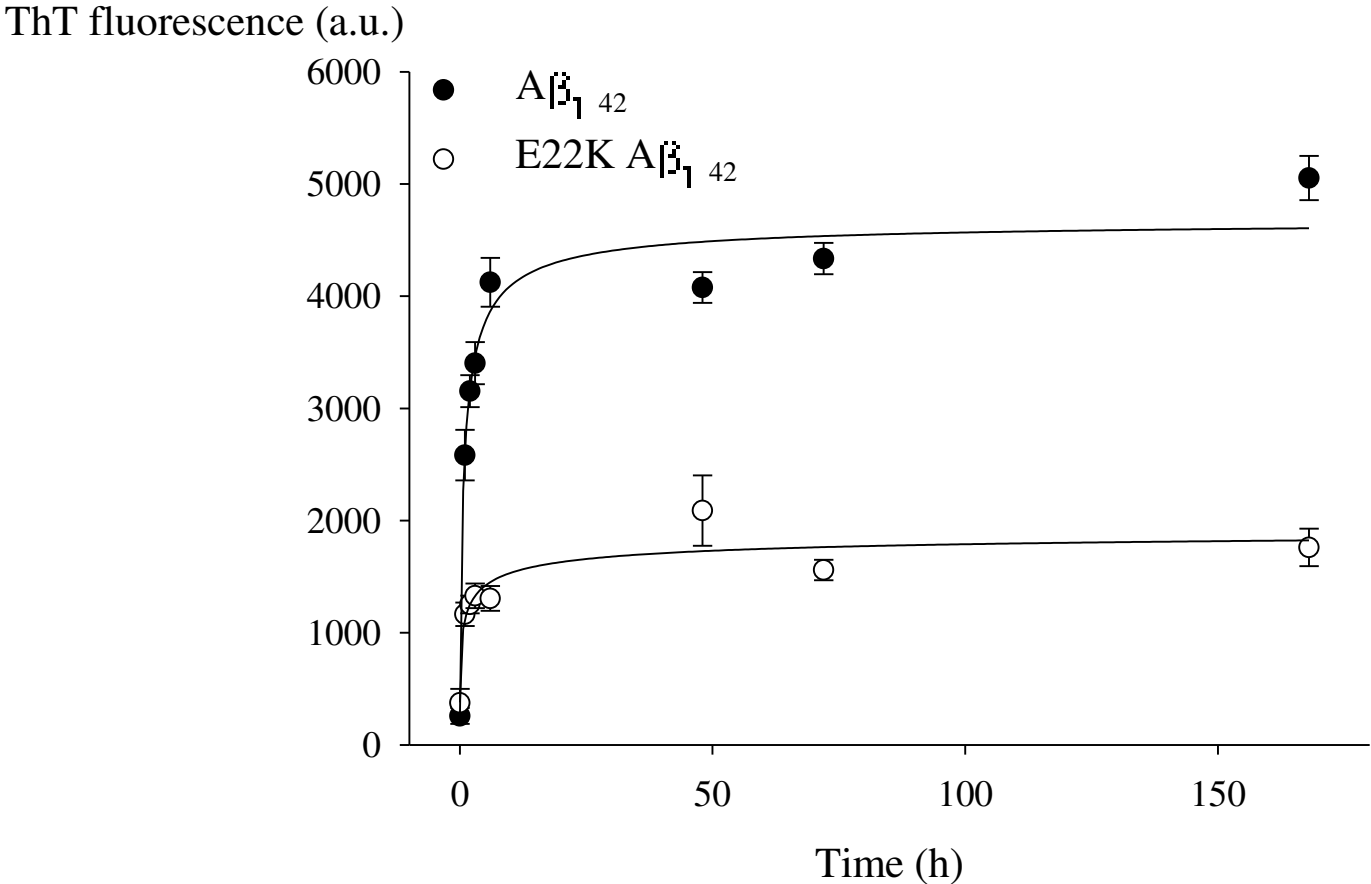


Vessel walls prone to rupture
Narrows lumina



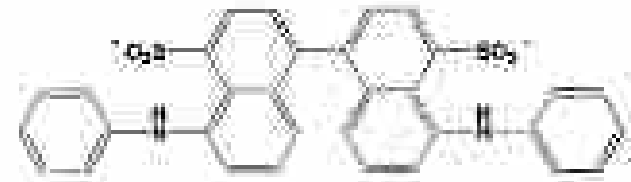
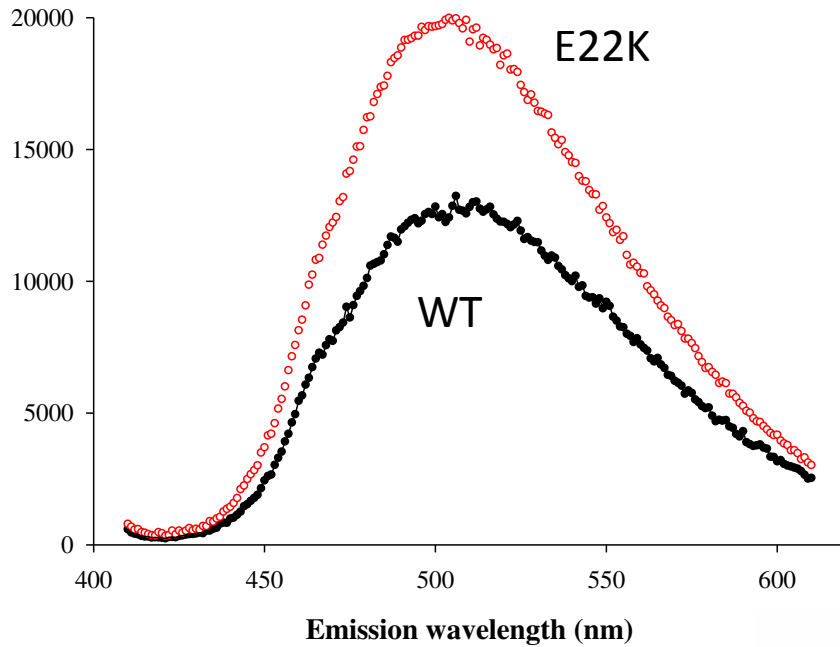
Intracerebral and subarachnoid bleeding
Infarcts
Periventricular oedema

ThT fluorescence intensity of fibrils varies with mutation



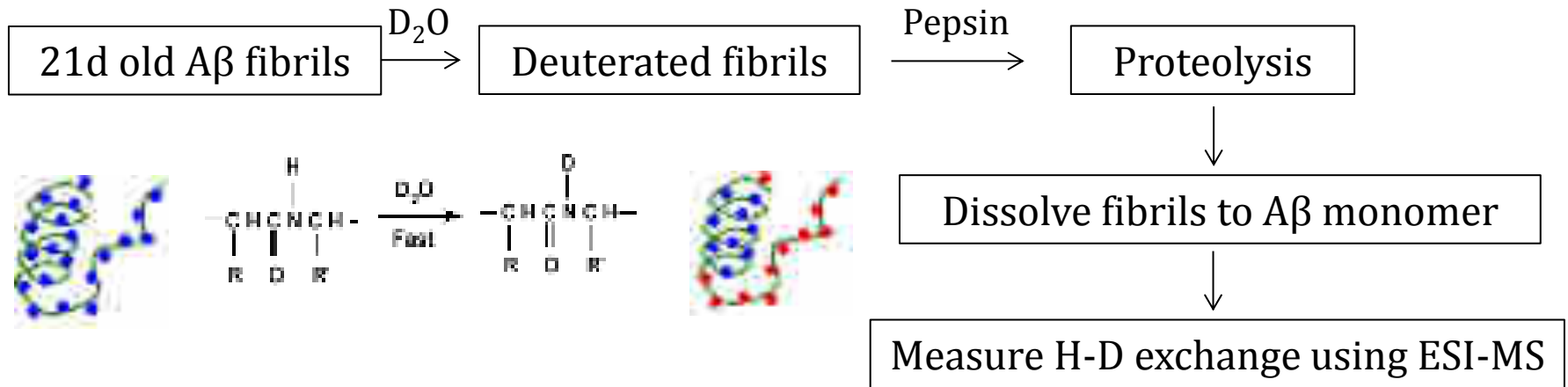
Wild type and E22K $A\beta_{1-42}$ fibrils interact differently with conformation-dependent fluorescent probes (ThT and bis-ANS)

Bis-ANS fluorescence (a.u.)



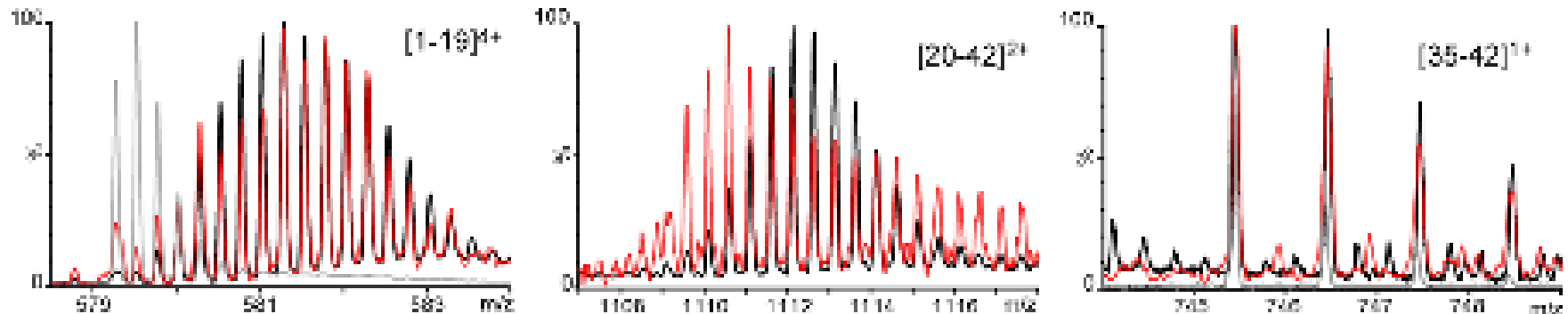
Bis-ANS: binds to hydrophobic patches
LeVine H. (2002)

Wild type and E22K A β_{1-42} differ in solvent accessibility in the A β region comprising residues [20-34]

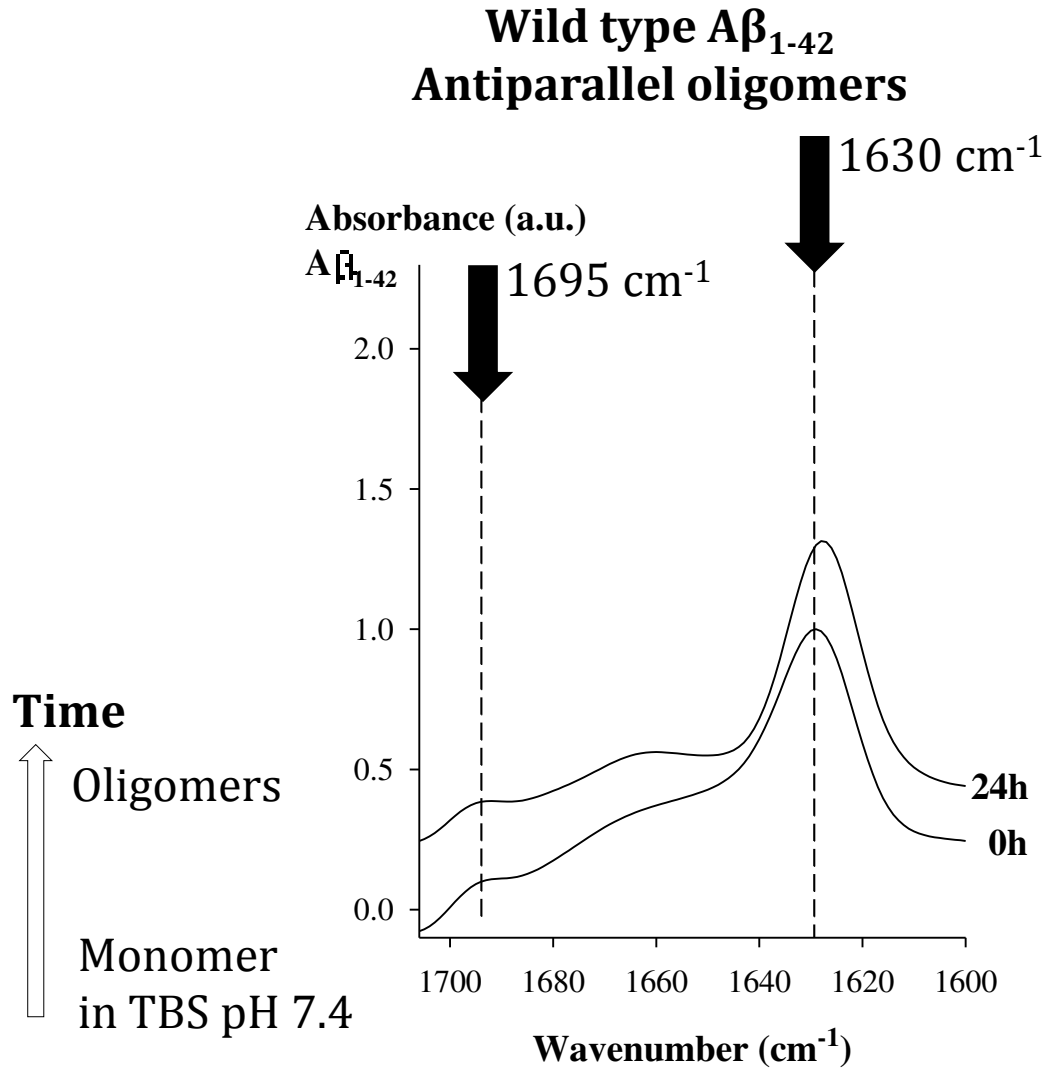


H/D exchange - Mass Spectrometry with proteolytic fragmentation

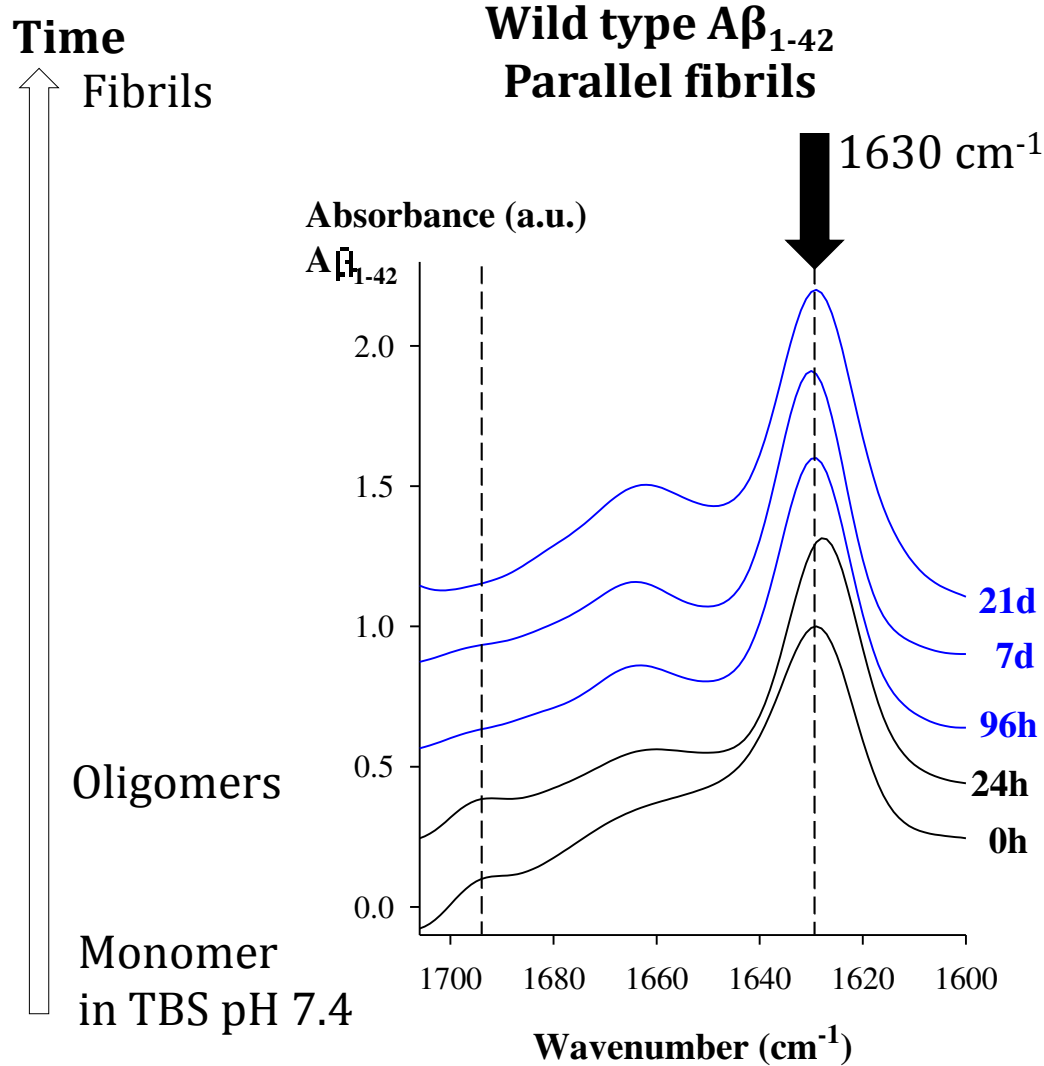
WT/E22K WT (HDX) E22K (HDX)



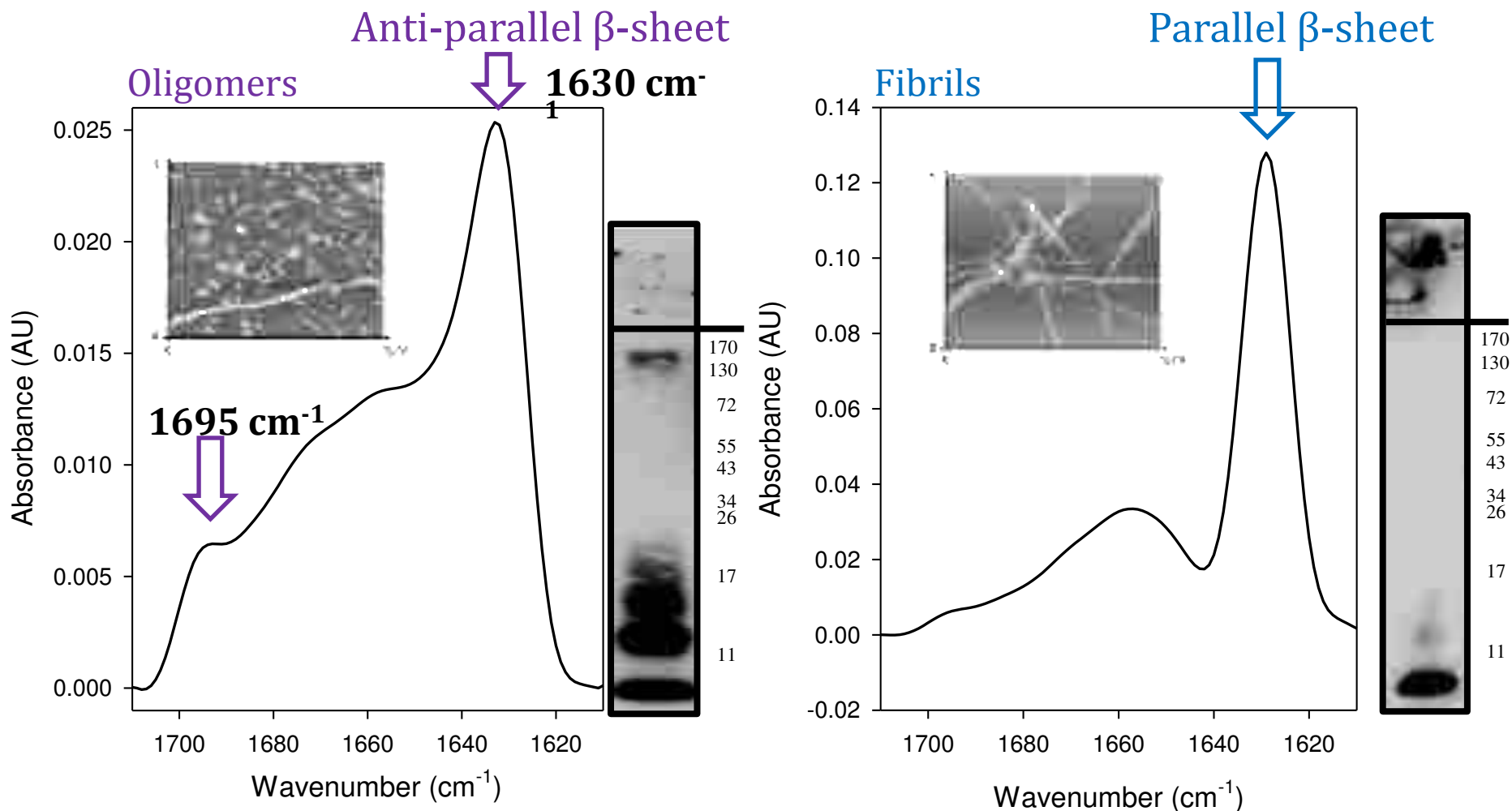
Changes in secondary structure during A β aggregation can be monitored by ATR-FTIR



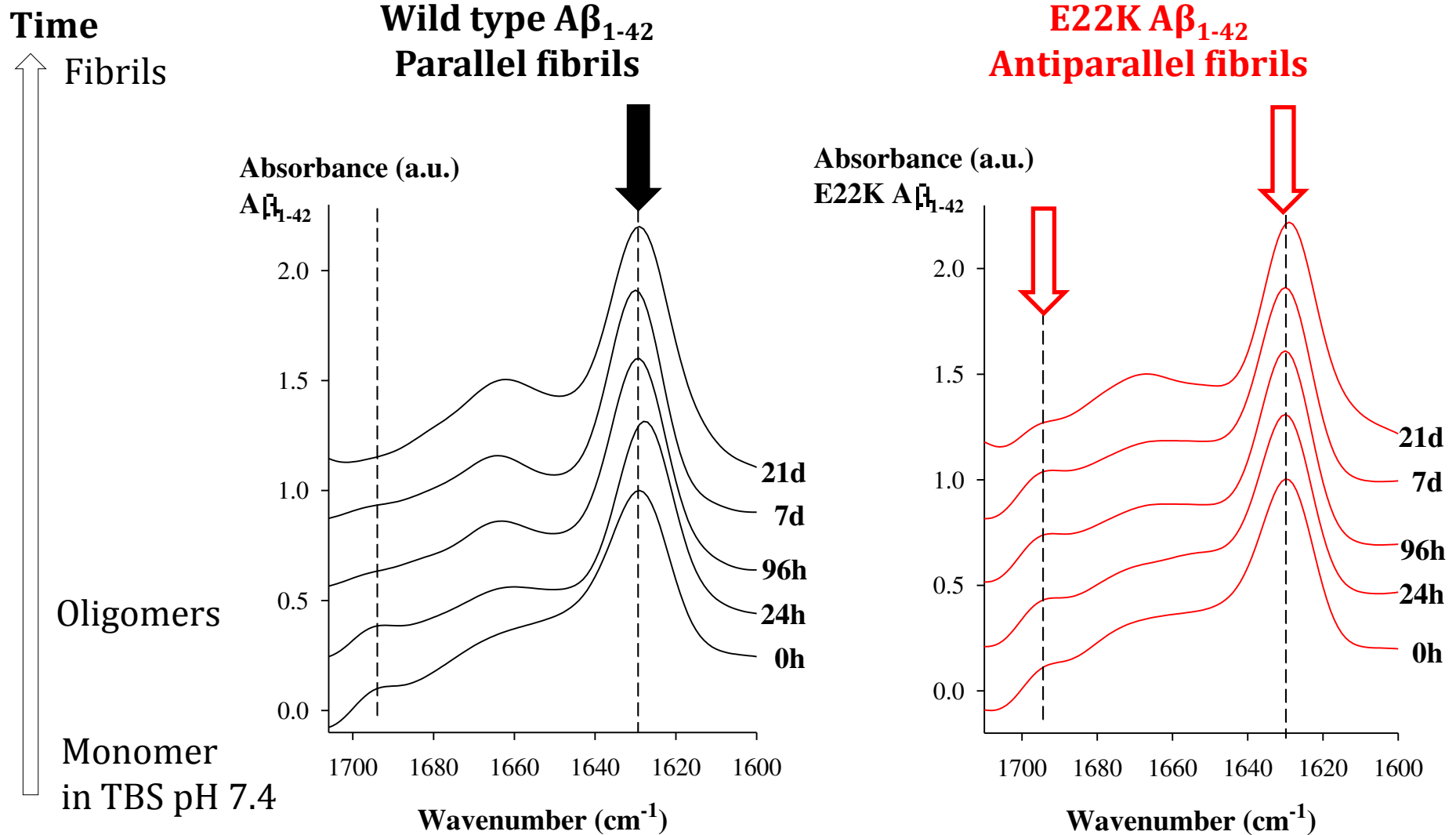
Changes in secondary structure during A β aggregation can be monitored by ATR-FTIR



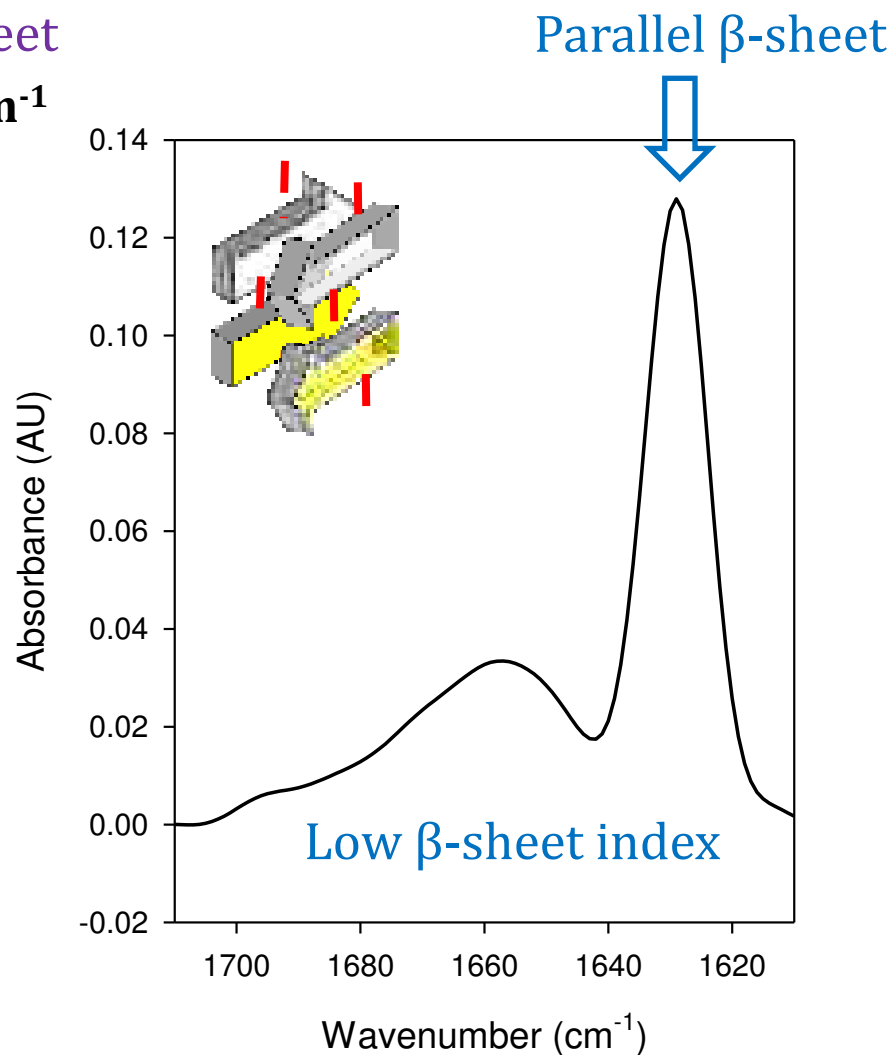
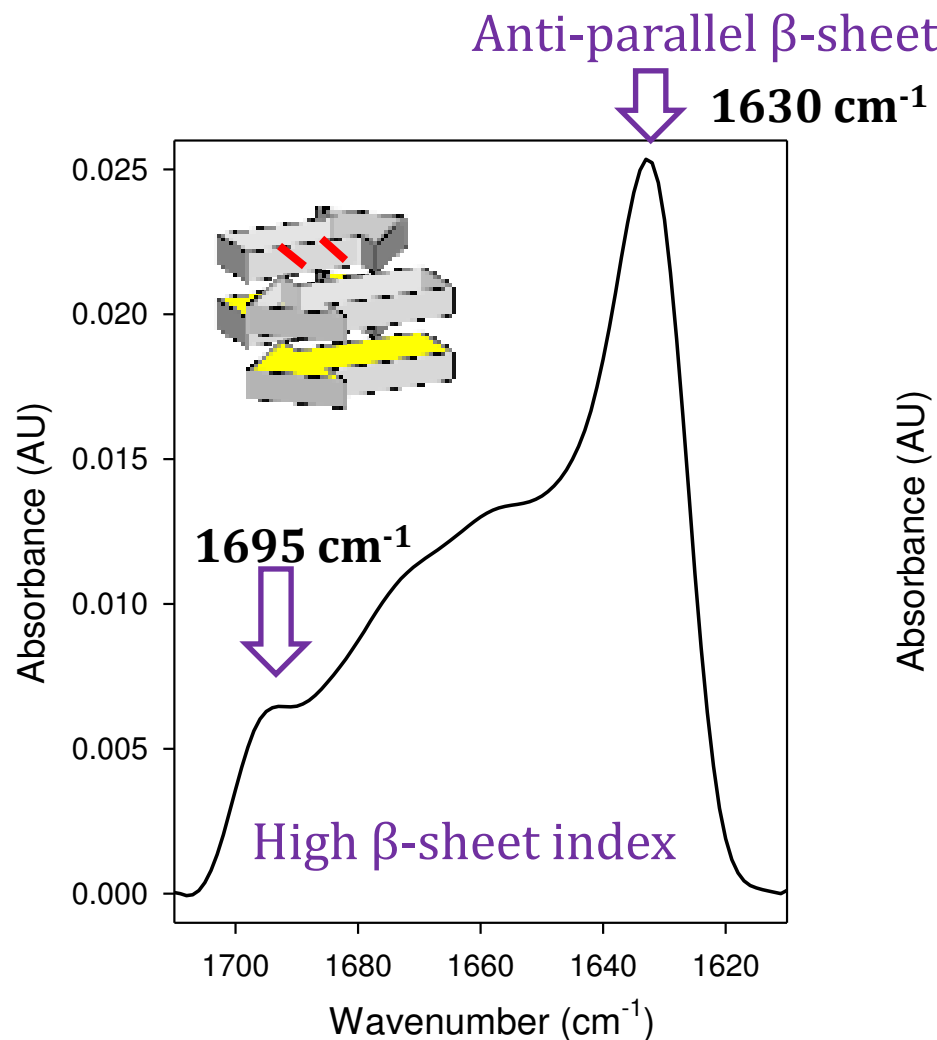
ATR-FTIR reveals major change in secondary structure during transformation of wild type A β oligomers into fibrils



A minor peak around 1695 cm^{-1} , representative of antiparallel β sheets, persists during E22K $\text{A}\beta_{1-42}$ aggregation

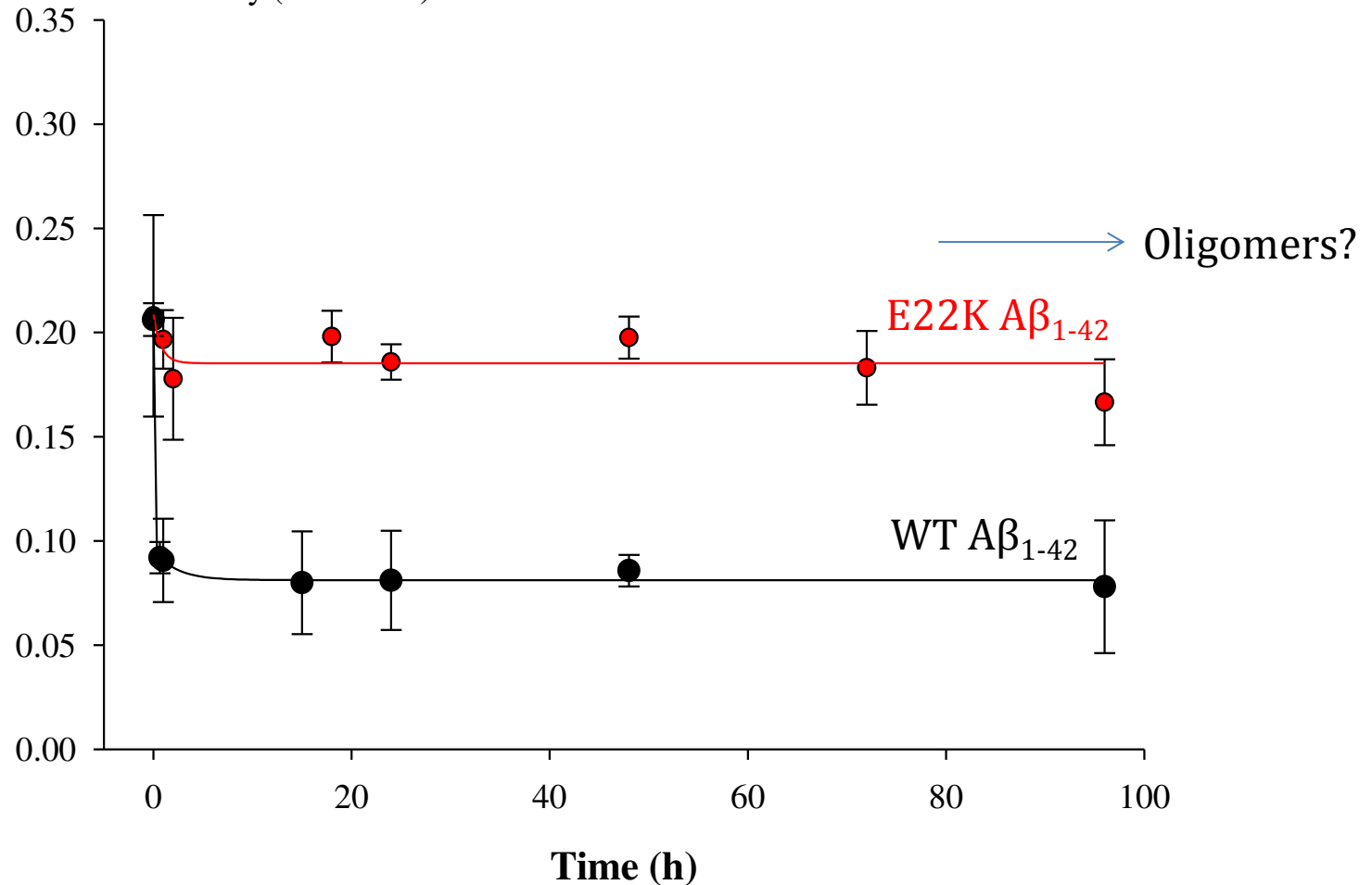


The β -sheet index ($1695/1630$ intensities ratio) is proportional to the percentage of antiparallel arrangement of β -strands in a β -sheet

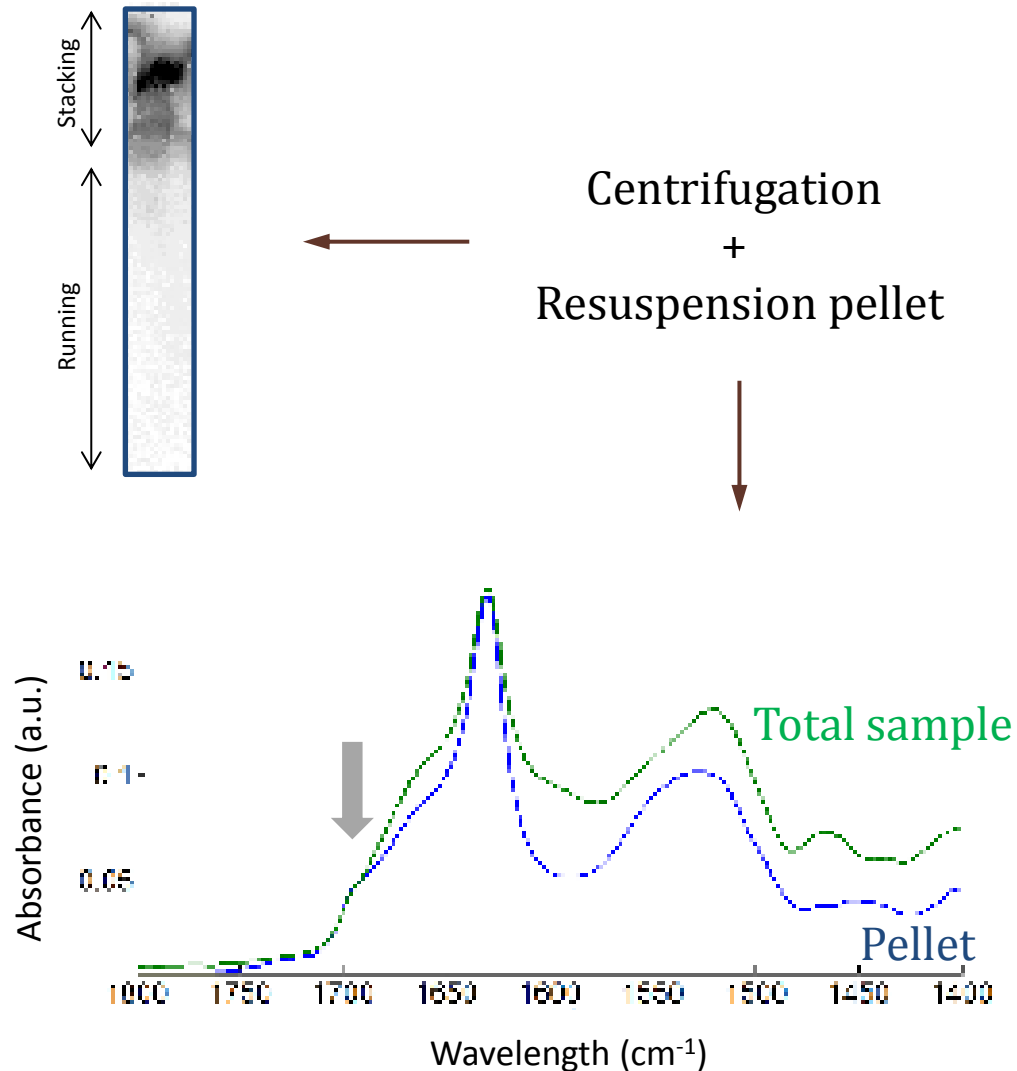


The β -sheet index (1695/1630 intensities ratio) is proportional to the percentage of antiparallel arrangement of β -strands in a β -sheet

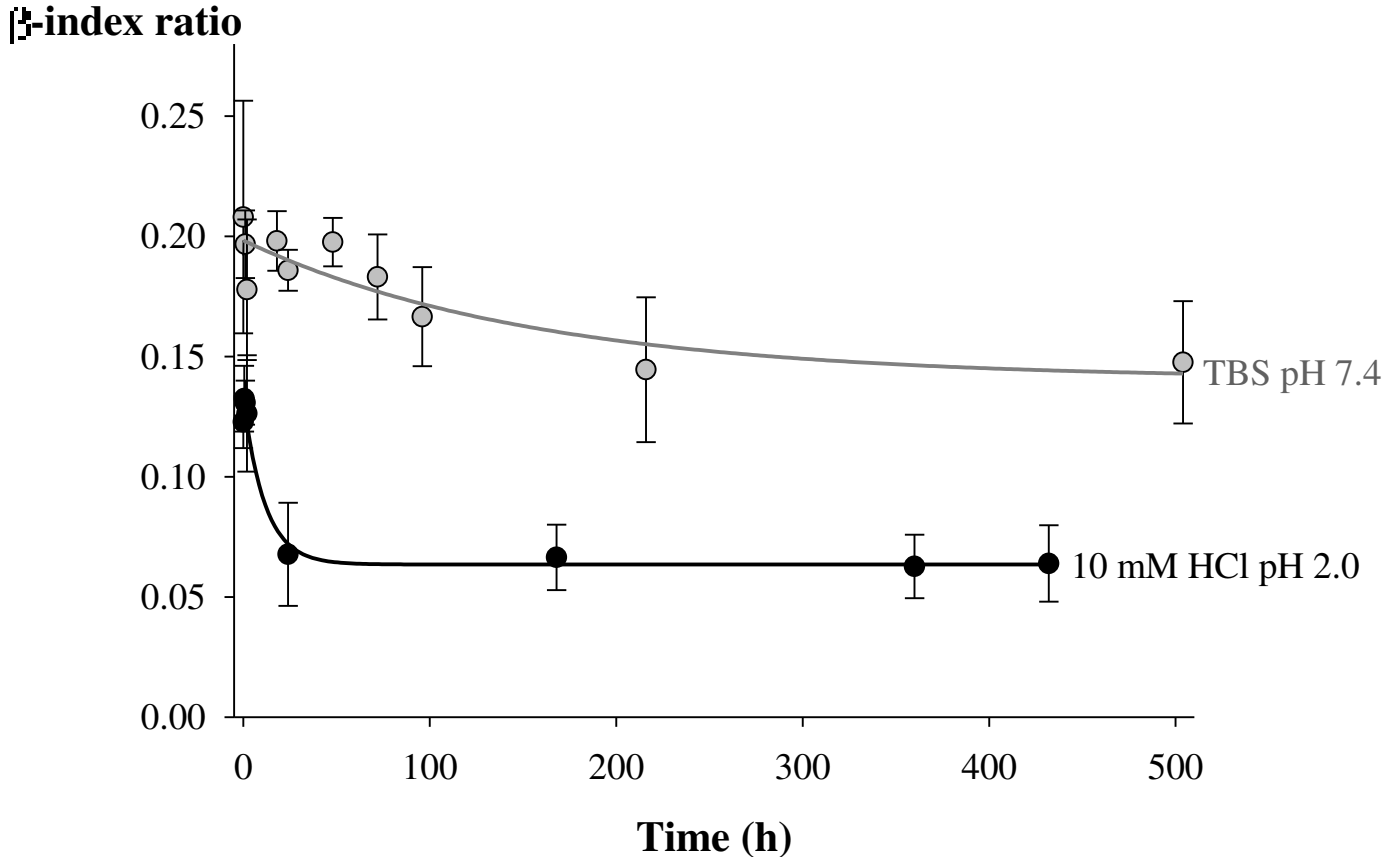
$$\beta\text{-index ratio} = \frac{\text{Intensity (1695 cm}^{-1}\text{)}}{\text{Intensity (1630 cm}^{-1}\text{)}}$$



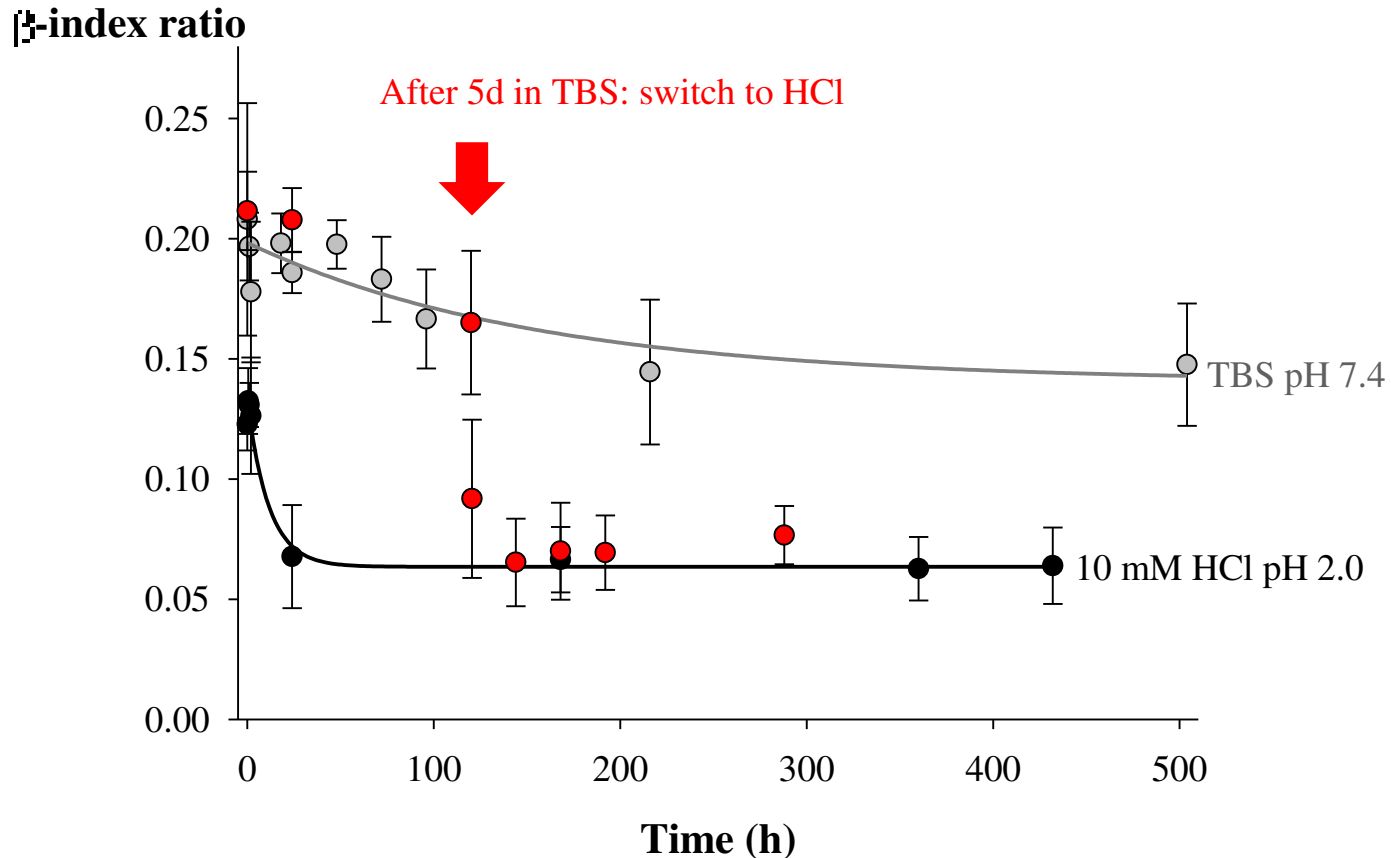
But: Fibrils and not remaining oligomers are responsible for observed antiparallel β -sheet conformation



E22K A β_{1-42} fibrils can convert from antiparallel to parallel β -sheet structure in an acidic environment (switch from pH 7.4 to pH 2.0)



E22K A β_{1-42} fibrils can convert from antiparallel to parallel β -sheet structure in an acidic environment (switch from pH 7.4 to pH 2.0)



Conclusions

'Physiologically' relevant mixtures of A β peptides may behave differently from what you would expect based on their isolated characteristics.

What is the perfect in vitro read-out for AD-related toxicity?

Antiparallel β -sheet fibrils exist, are remarkably stable but can also be converted upon changing environmental conditions.

Acknowledgements

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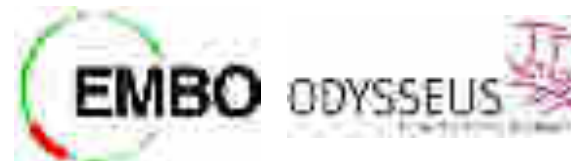
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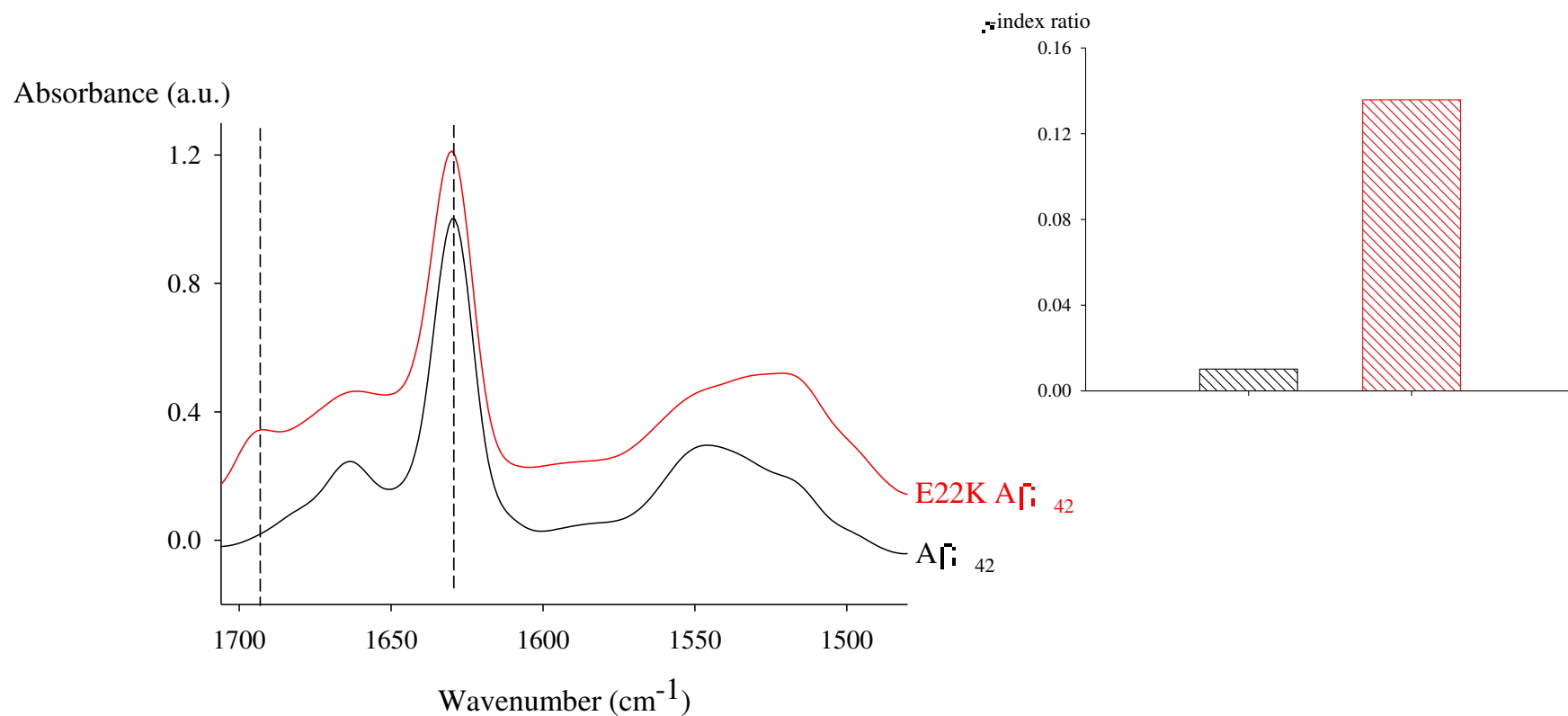
University of Sussex

Louise Serpell
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Thomas Williams


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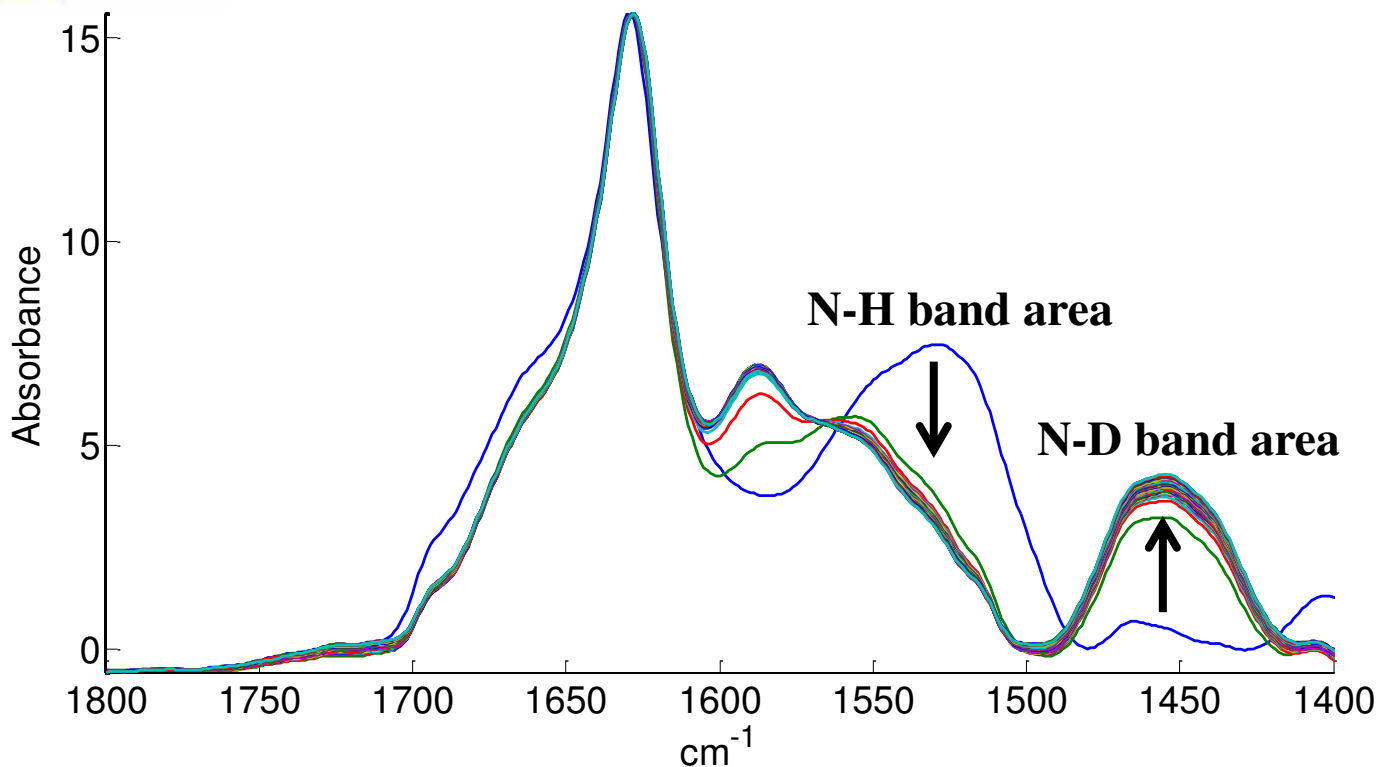
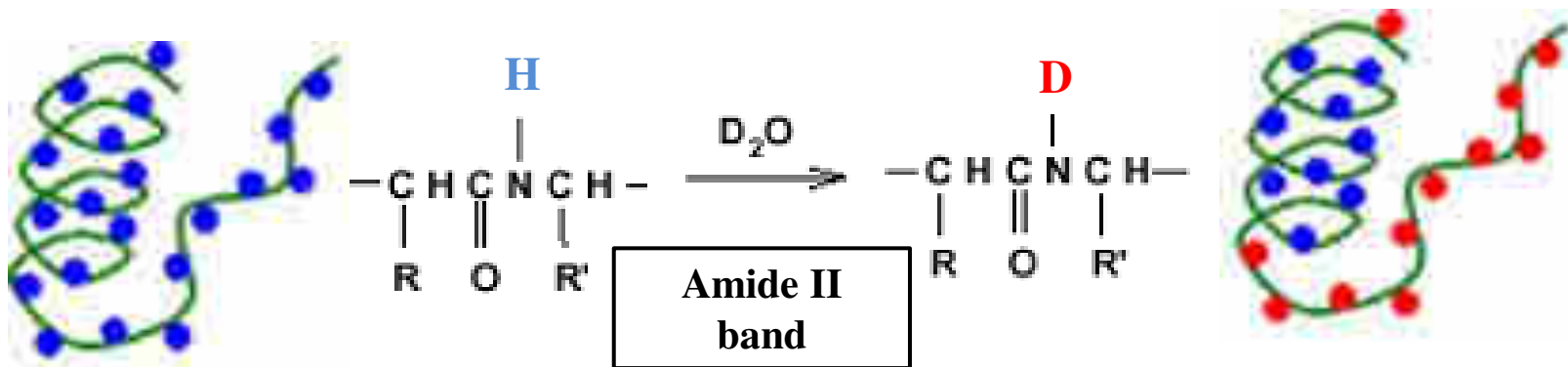
E22K A β : more random coil and turn at the expense of β -structure.



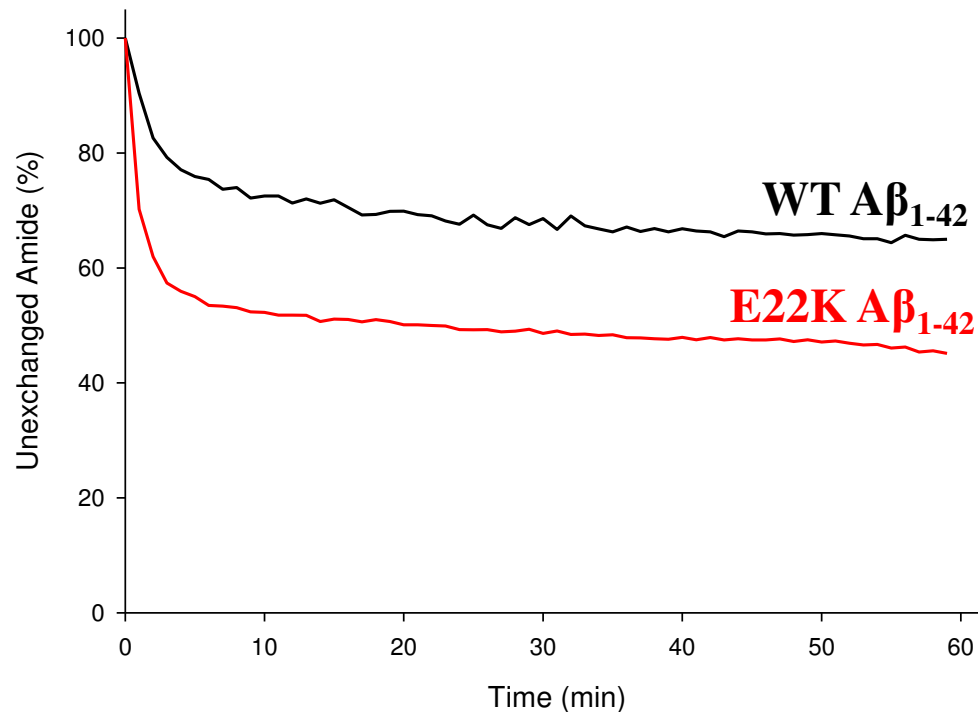
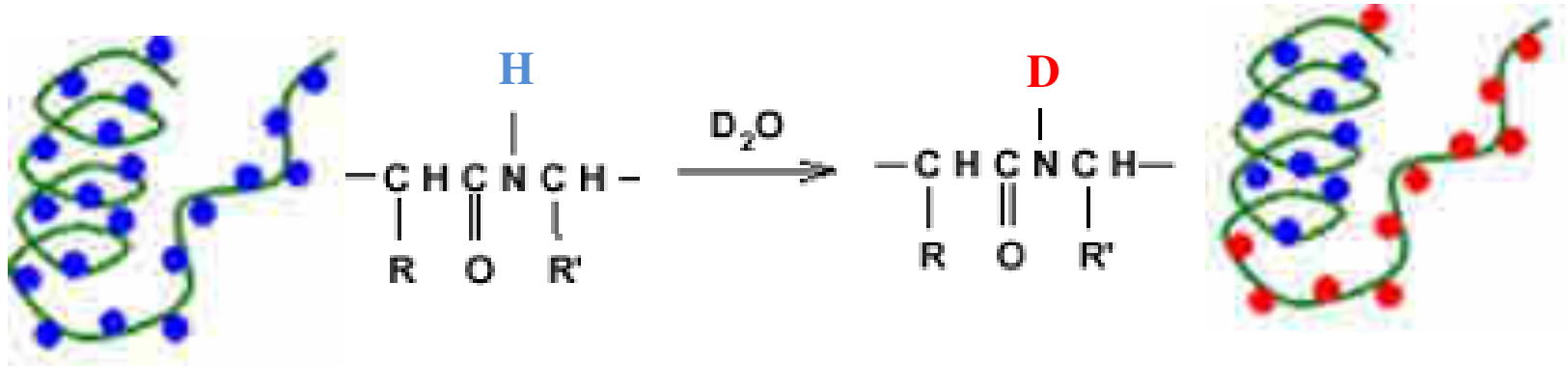
	A β ₁₋₄₂	E22K A β ₁₋₄₂
β -structure	73	59
α -helix	8	10
random	11	18
turn	8	13

Approx. 8 residues 

H/D exchange shows 20 % difference in accessibility of backbone amide hydrogens between wild type and E22K A β_{1-42} fibrils



H/D exchange shows 20 % difference in accessibility of backbone amide hydrogens between wild type and E22K A β_{1-42} fibrils



H/D exchange monitored by ESI-MS and pepsin proteolysis reveals a higher solvent accessibility for the Italian mutant in the central region

DAEFRHDSGYEVHHQKLV**FFAEDVGSNKGAIIGLMVGGVVIA**

K

Peptide		Backbone amide protons exchanged	Protected NH/total
[1-19]	WT	14.0 ± 0.5	4/18
	E22K	14.3 ± 0.4	3.7/18
[20-42]	WT	7.3 ± 0.7	14.7/22
	E22K	9.6 ± 0.8	12.4/22
[35-42]	WT	3.1 ± 0.5	3.9/7
	E22K	3.4 ± 0.3	3.6/7

pepsin



pepsin



DAEFRHDSGYEVHHQKLV**FFAEDVGSNKGAIIGLMVGGVVIA**

[1-19]

[20-42]

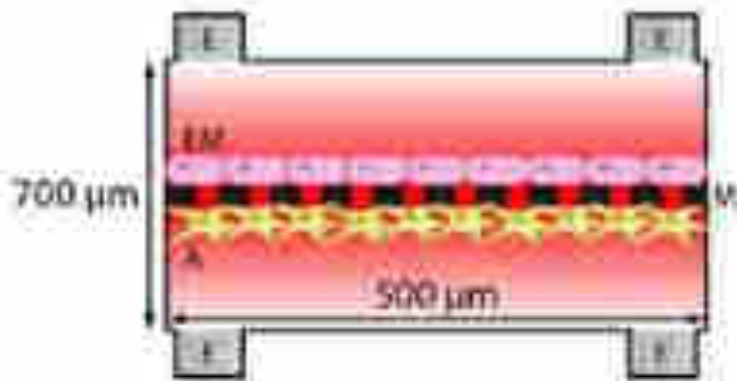
[35-42]

Conclusions & perspectives: is there a link between the underlying β -sheet orientation of A β fibrils and disease-associated pathology?

- E22K A β 42 (and D23N A β 40 and A β 42) arrange into anti-parallel β -sheet fibrils.
- E22K fibrils are structurally different from WT A β 42.
- Antiparallel β -sheet fibrils from E22K A β 42 can rapidly (< 30 min) converse into parallel by changing the pH.

Work in progress – relation with CAA:

Do antiparallel β -sheet fibrils disintegrate the blood-brain barrier \rightarrow blood-brain-barrier-on-a-chip technology.

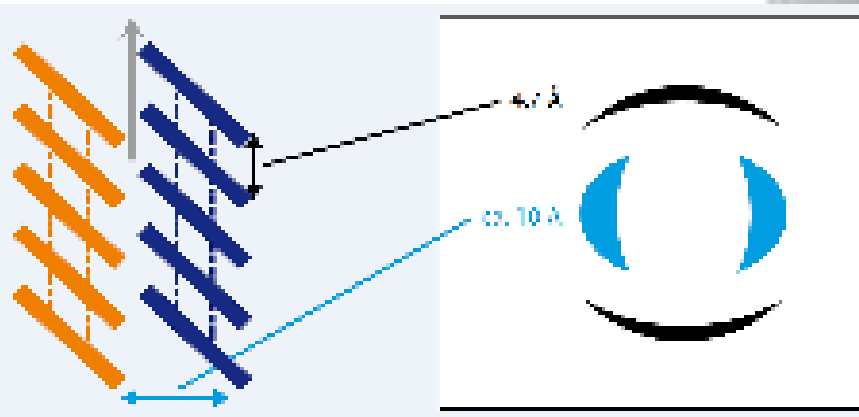
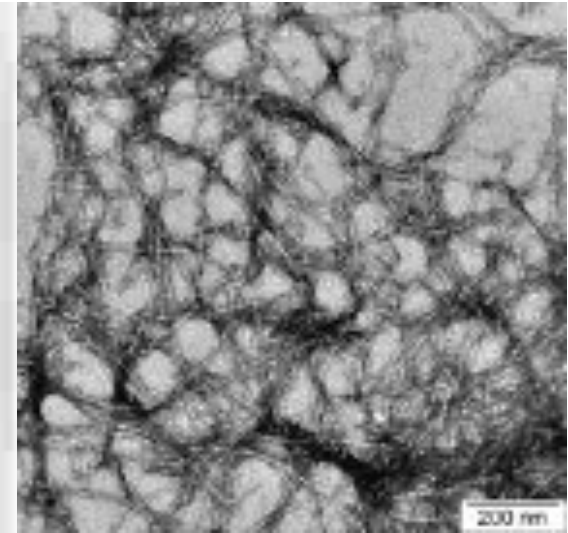
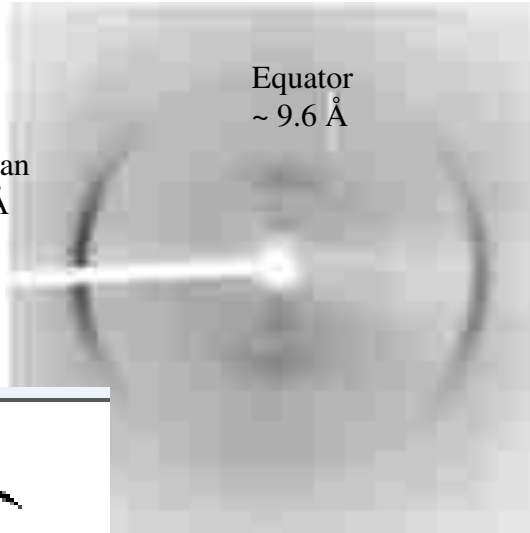


In our search for additional confirmation ...

$A\beta_{1-42}$

Meridian
~ 4.7 Å

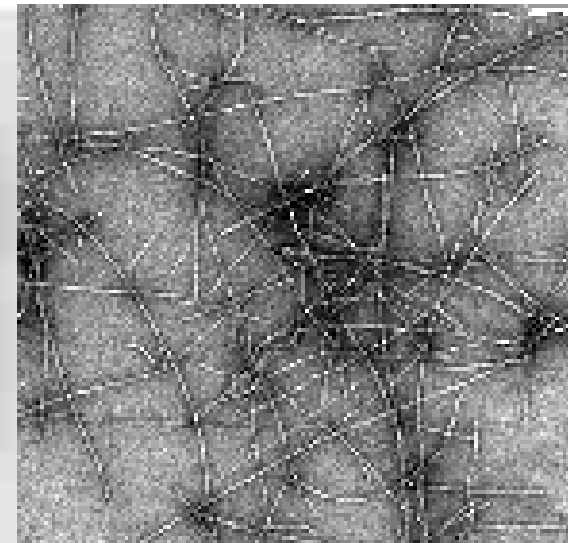
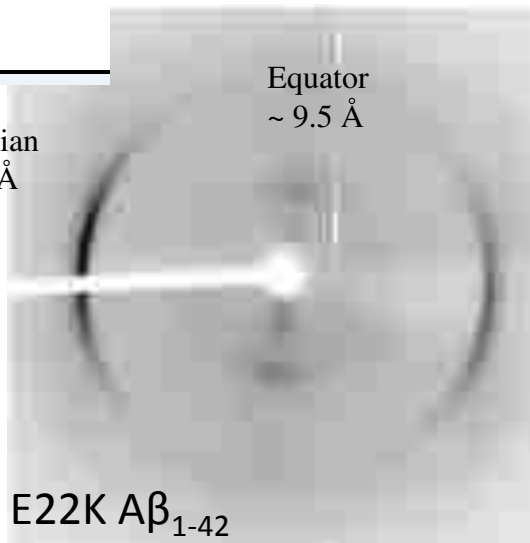
Equator
~ 9.6 Å



X-ray fiber diffraction → two reflections but no differences in structure.

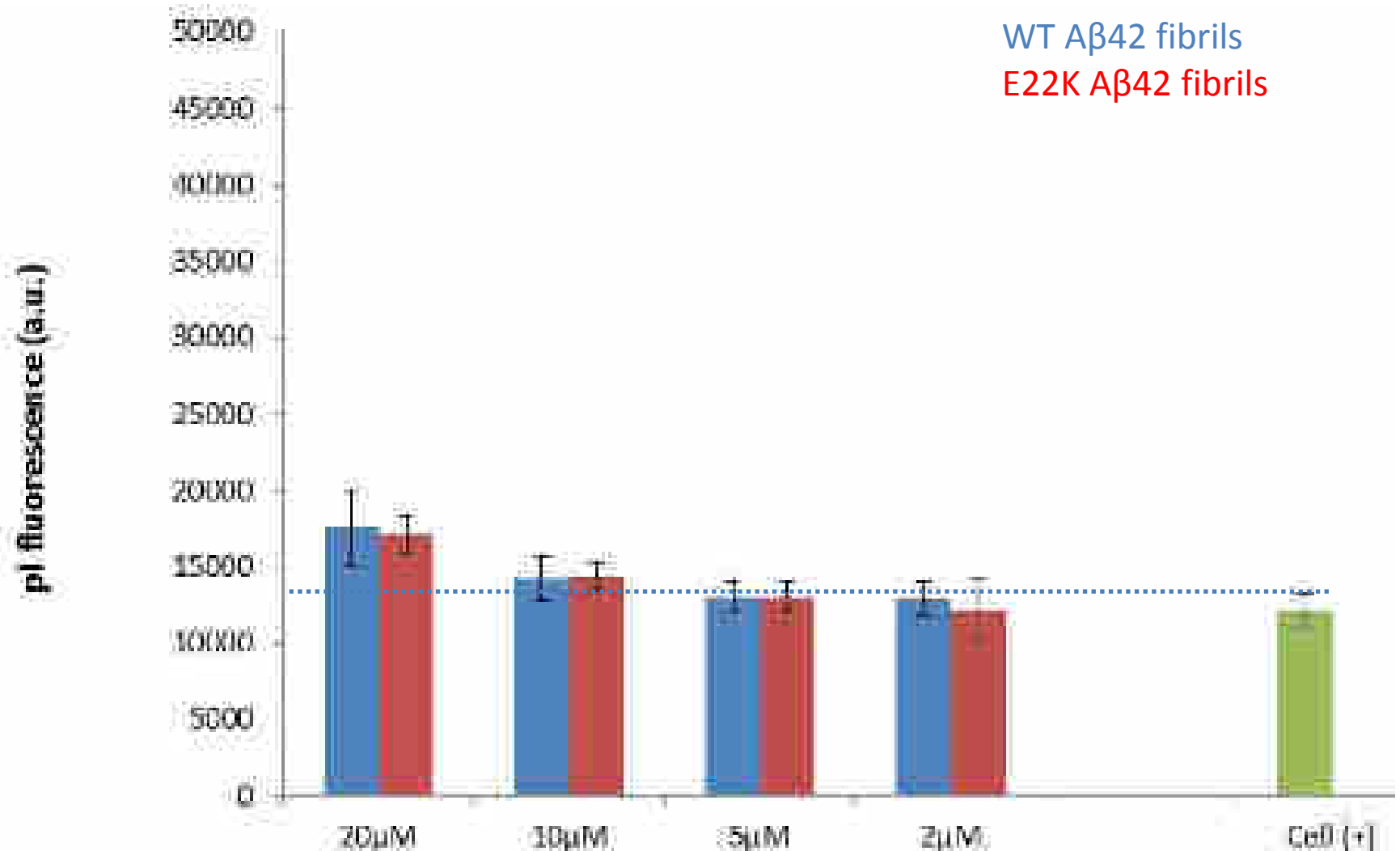
Meridian
~ 4.7 Å

Equator
~ 9.5 Å

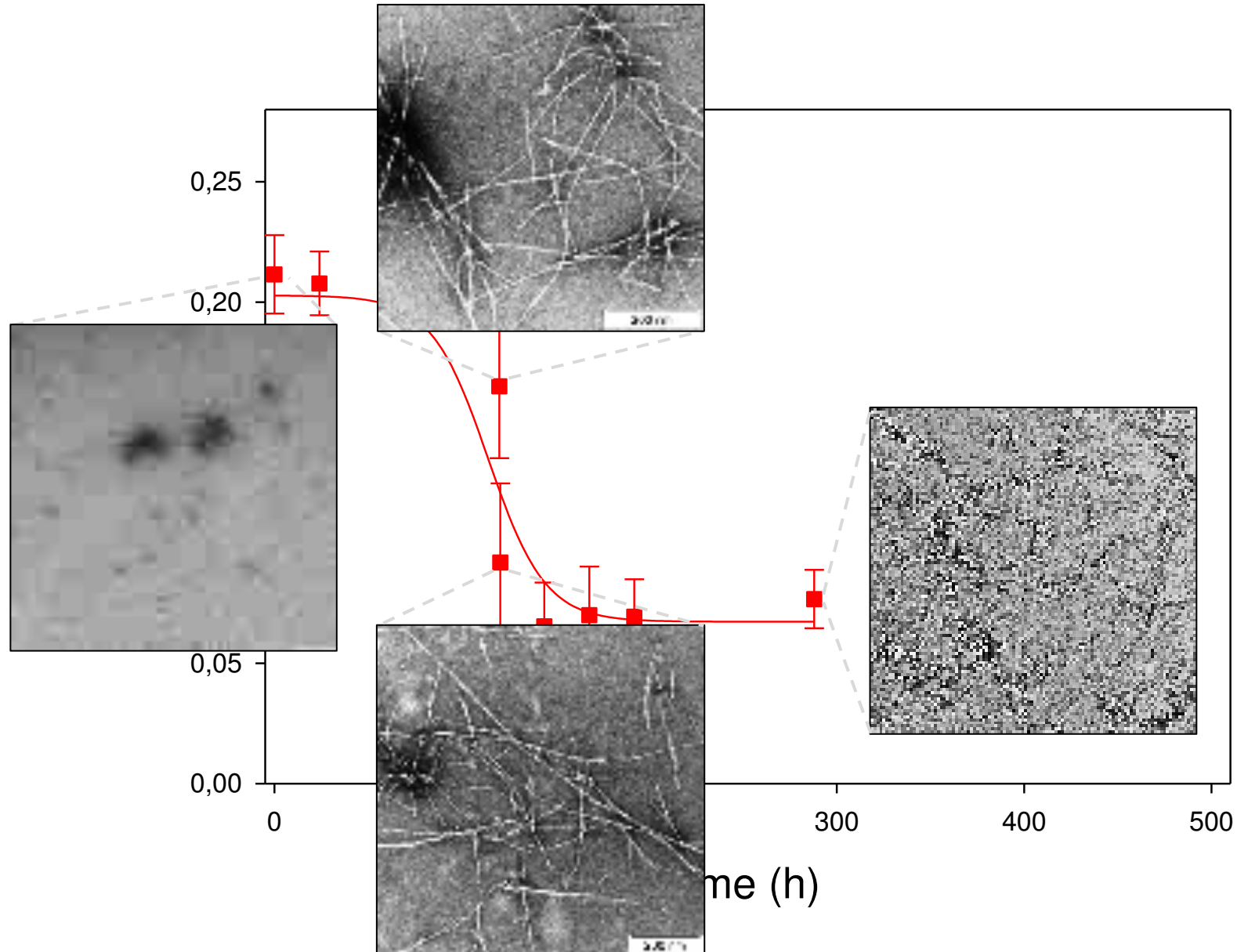


E22K $A\beta_{1-42}$

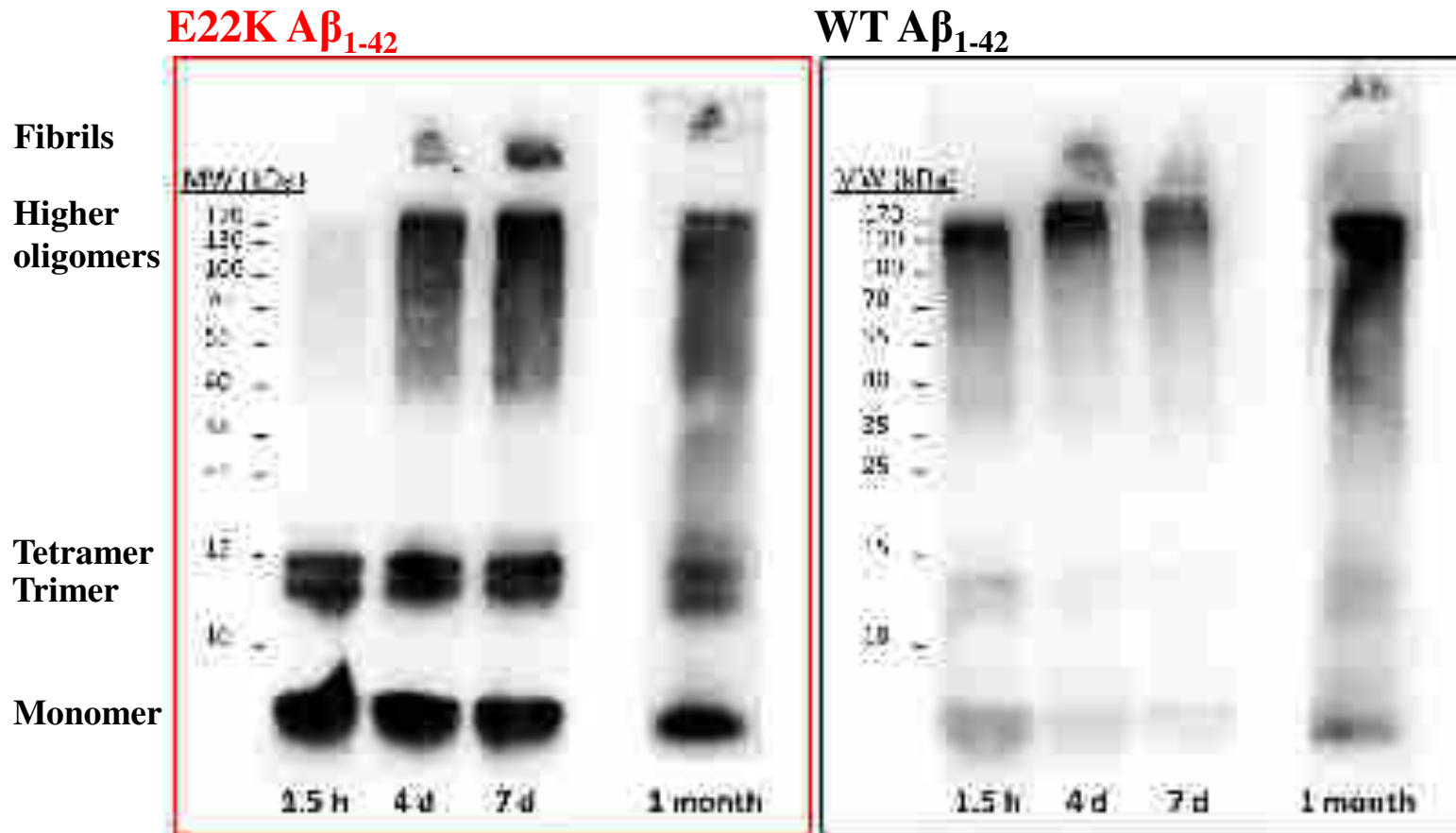
Fibrils from E22K and WT A β 42 are similarly 'toxic' to astrocytes (and neuroblastoma cells).



Mature E22K $A\beta_{1-42}$ fibrils can convert from antiparallel to parallel structure

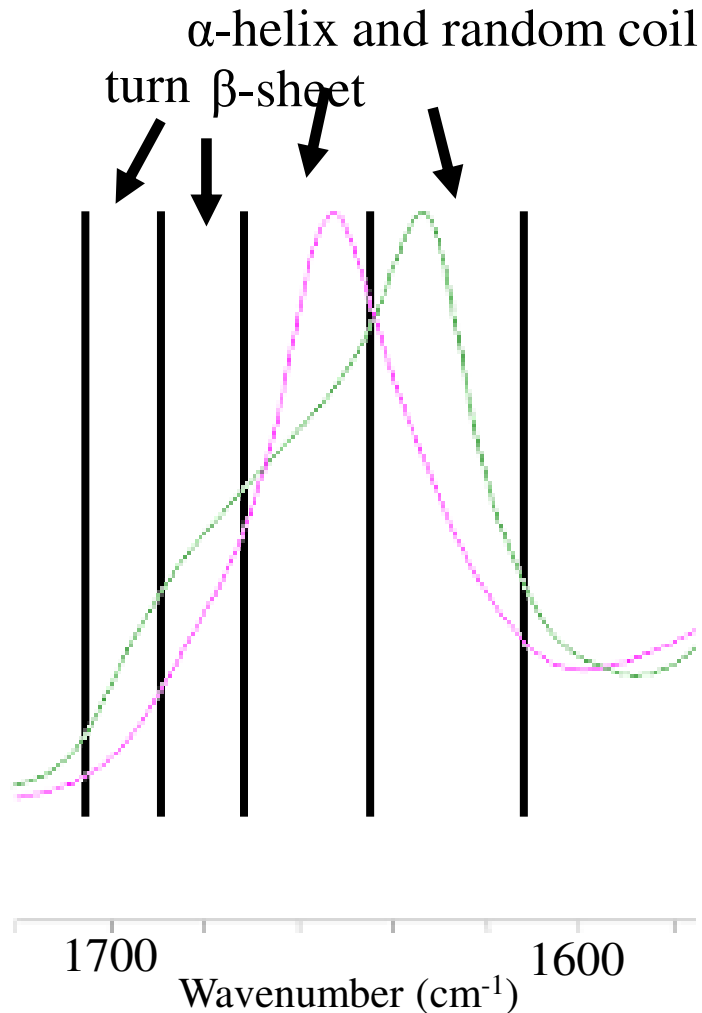


E22K A β_{1-42} shows sustained (SDS-stable) oligomer formation.



SDS-PAGE/WB (antibody 6E10)

Quantitative analysis consists of deconvolution of the amide I band into its primary components, followed by a curve fitting

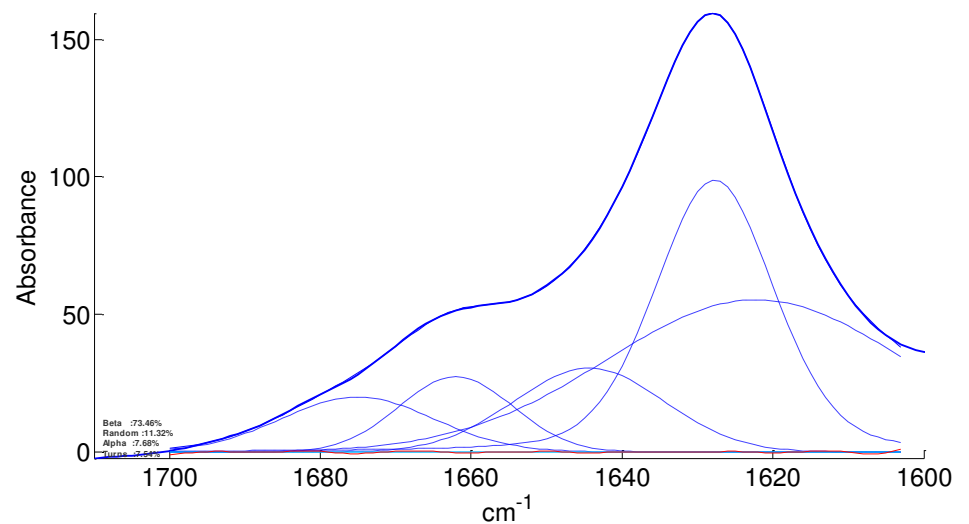


Parallel β -sheets:

1 peak at 1630 cm^{-1}

Antiparallel β -sheets:

peaks at 1630 cm^{-1} and 1695 cm^{-1}

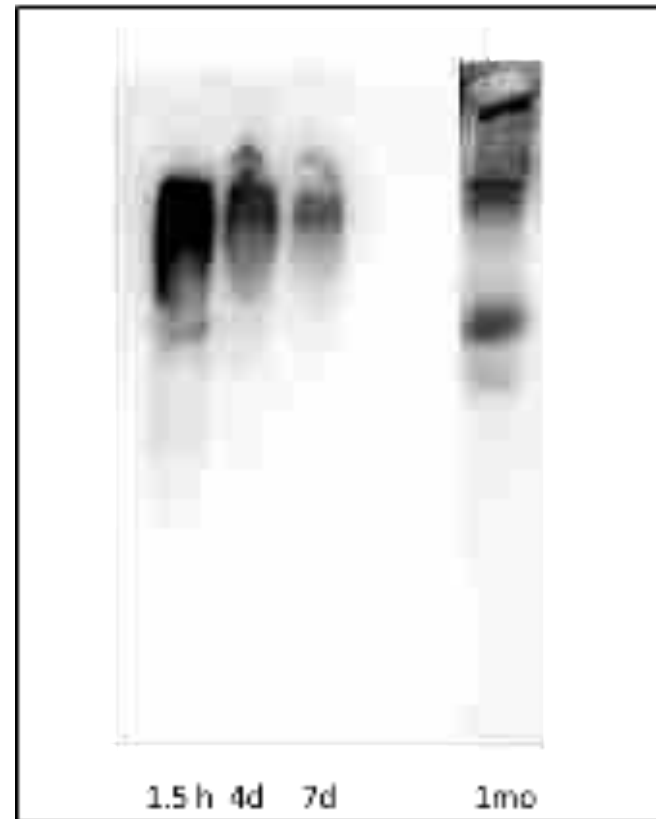


E22K A β_{1-42} shows sustained (SDS-stable) oligomer formation.

E22K A β_{1-42}

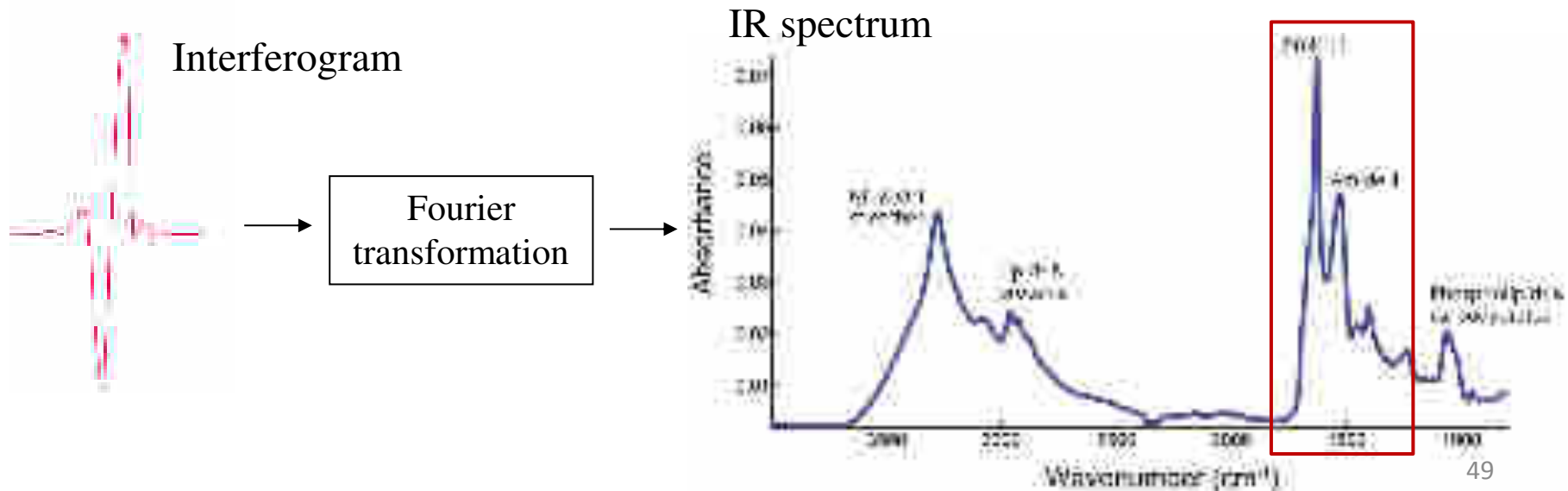


WT A β_{1-42}

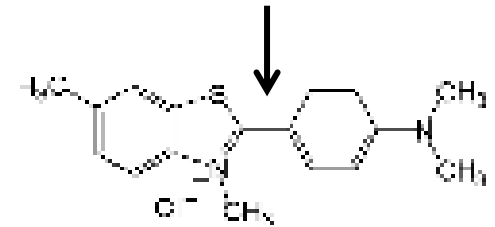
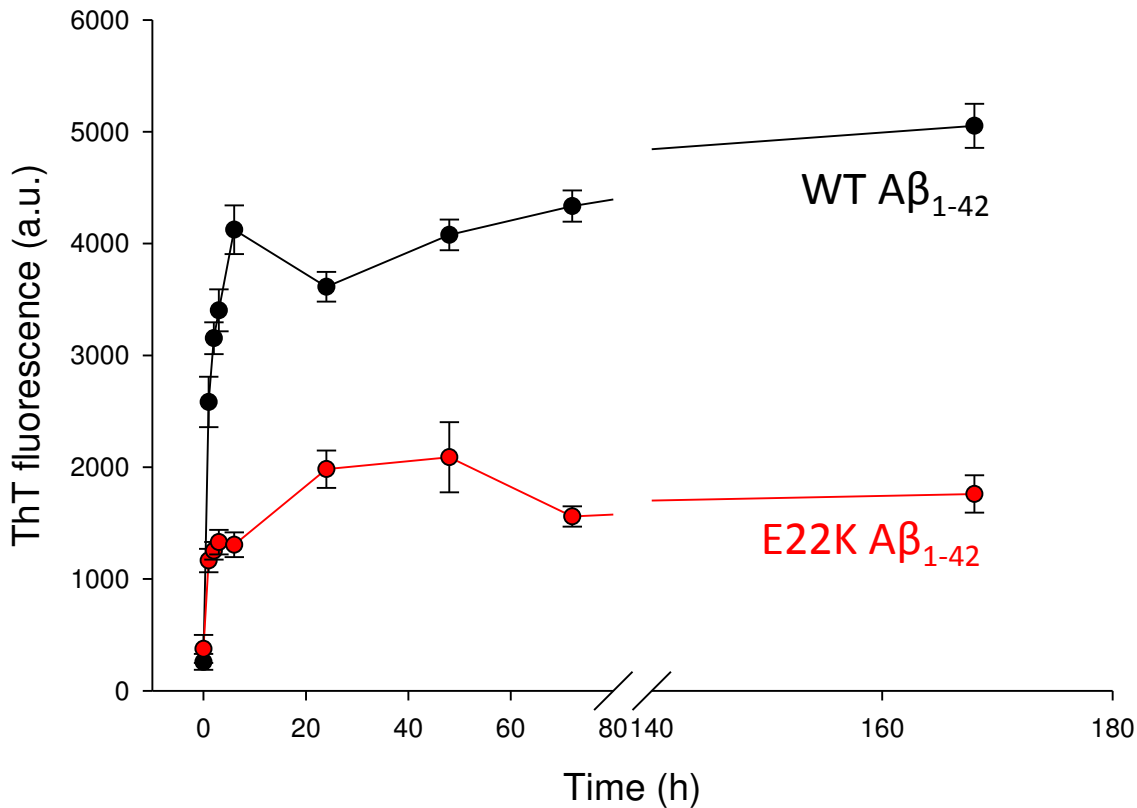


Native PAGE/WB (antibody 6E10)

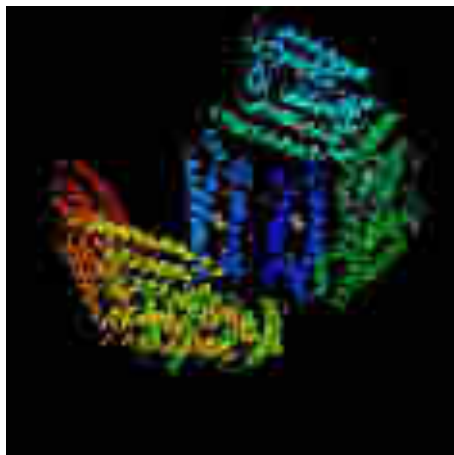
Attenuated Total Reflectance (ATR) – FTIR spectroscopy has the capability to identify functional groups (C=O, C-H, N-H, ...)



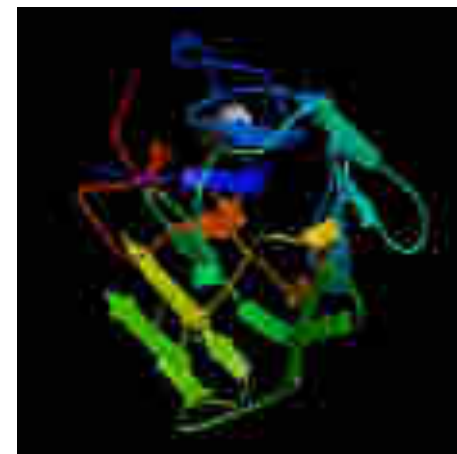
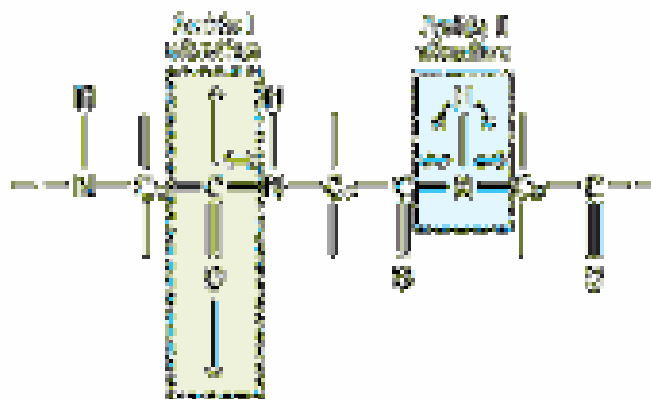
Thioflavin T (ThT) is less easily locked in its excited conformation in antiparallel fibrils, leading to a lower final ThT fluorescence intensity



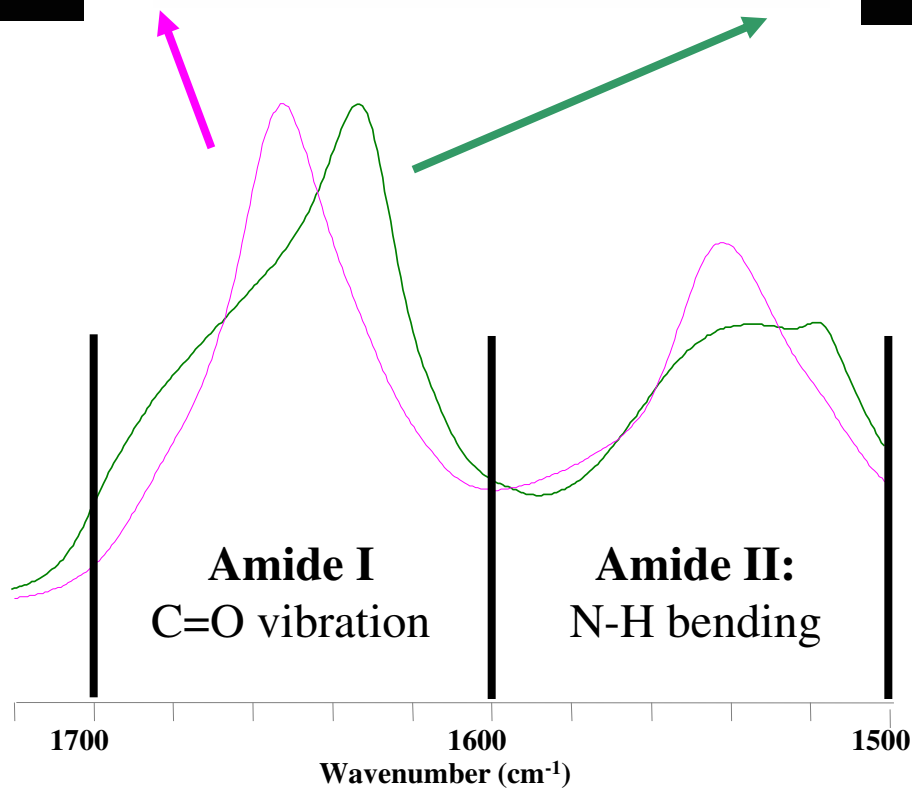
Amide I and Amide II bands contain secondary structure information



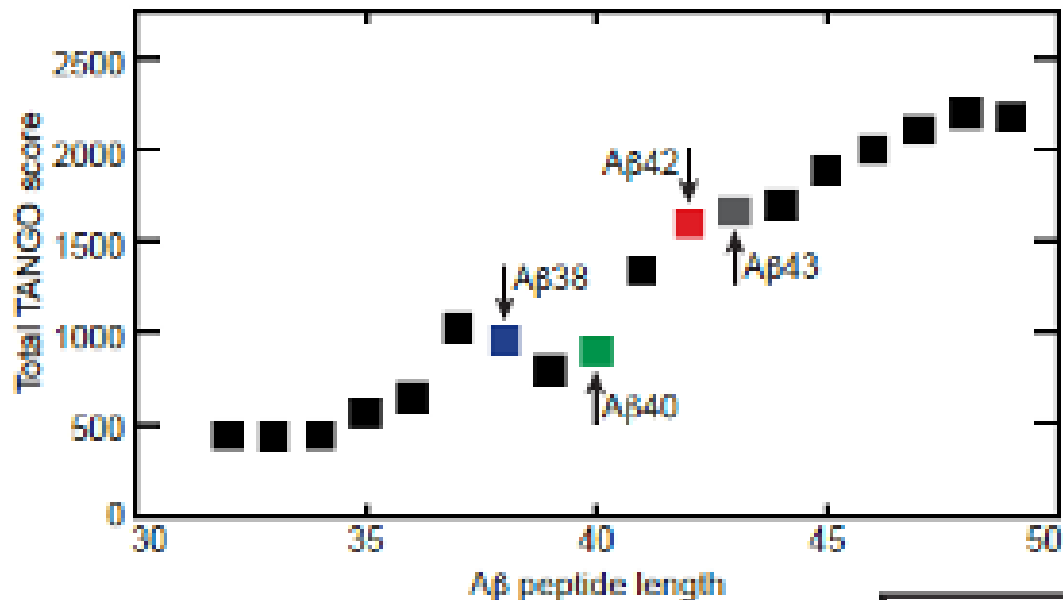
Ferritin



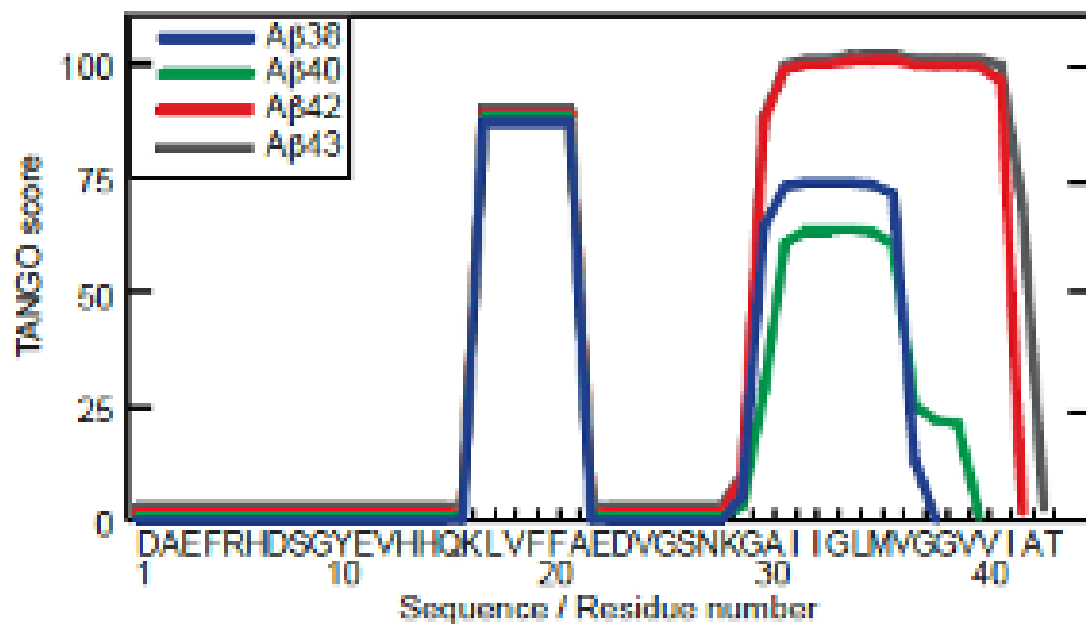
Concanavalin A



Aggregation of A β peptide is predicted to vary with peptide length



TANGO is an algorithm designed to predict the aggregation propensity of peptide sequences.
www.tango.switchlab.org



Antiparallel β -sheet architecture of Italian mutant E22K $\Delta\beta_{1-42}$ fibrils

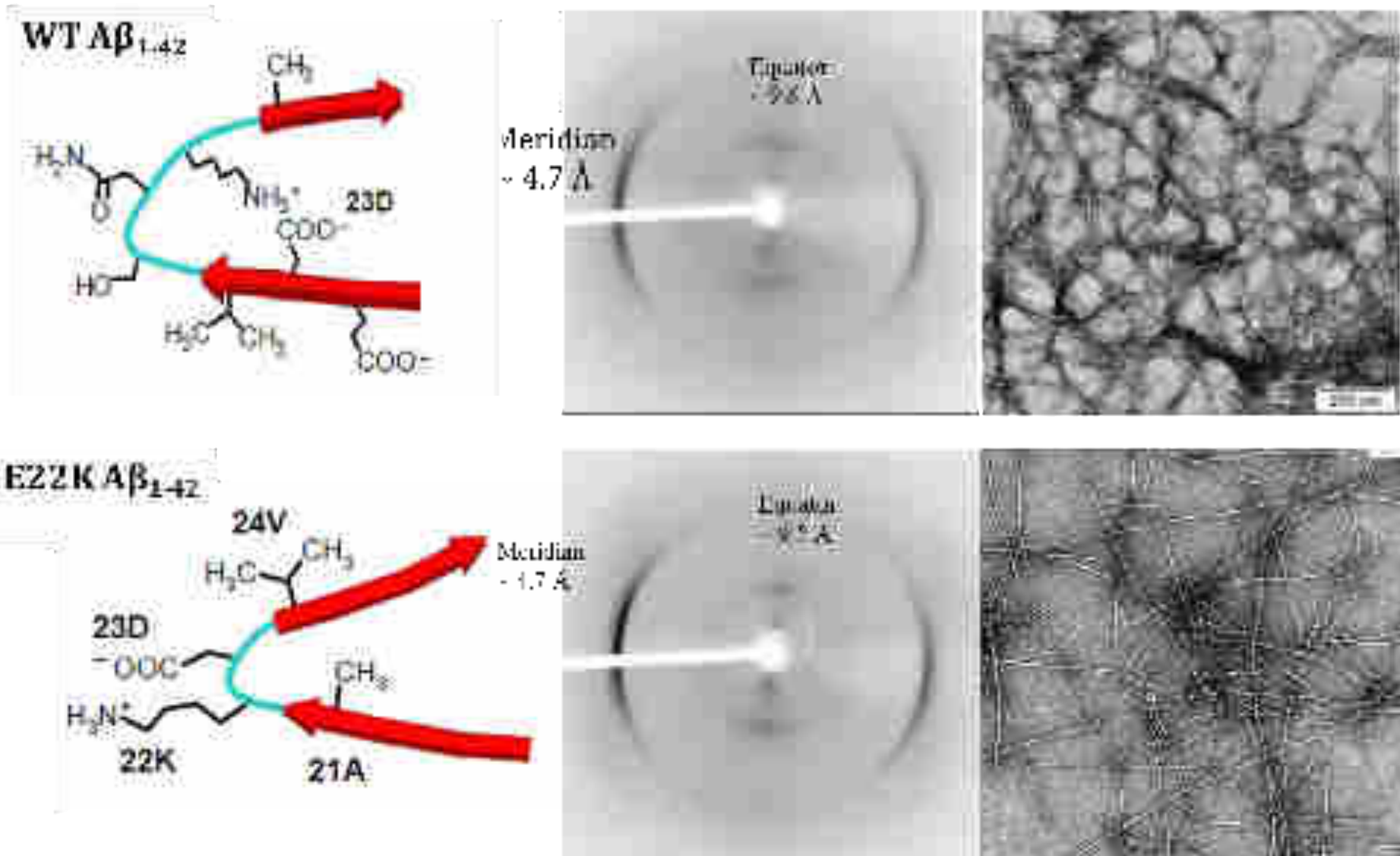


Figure adapted from Masuda *et al.* Bioorg & Med Chem (2005), FDX: Louise Serpell, Sussex University, UK