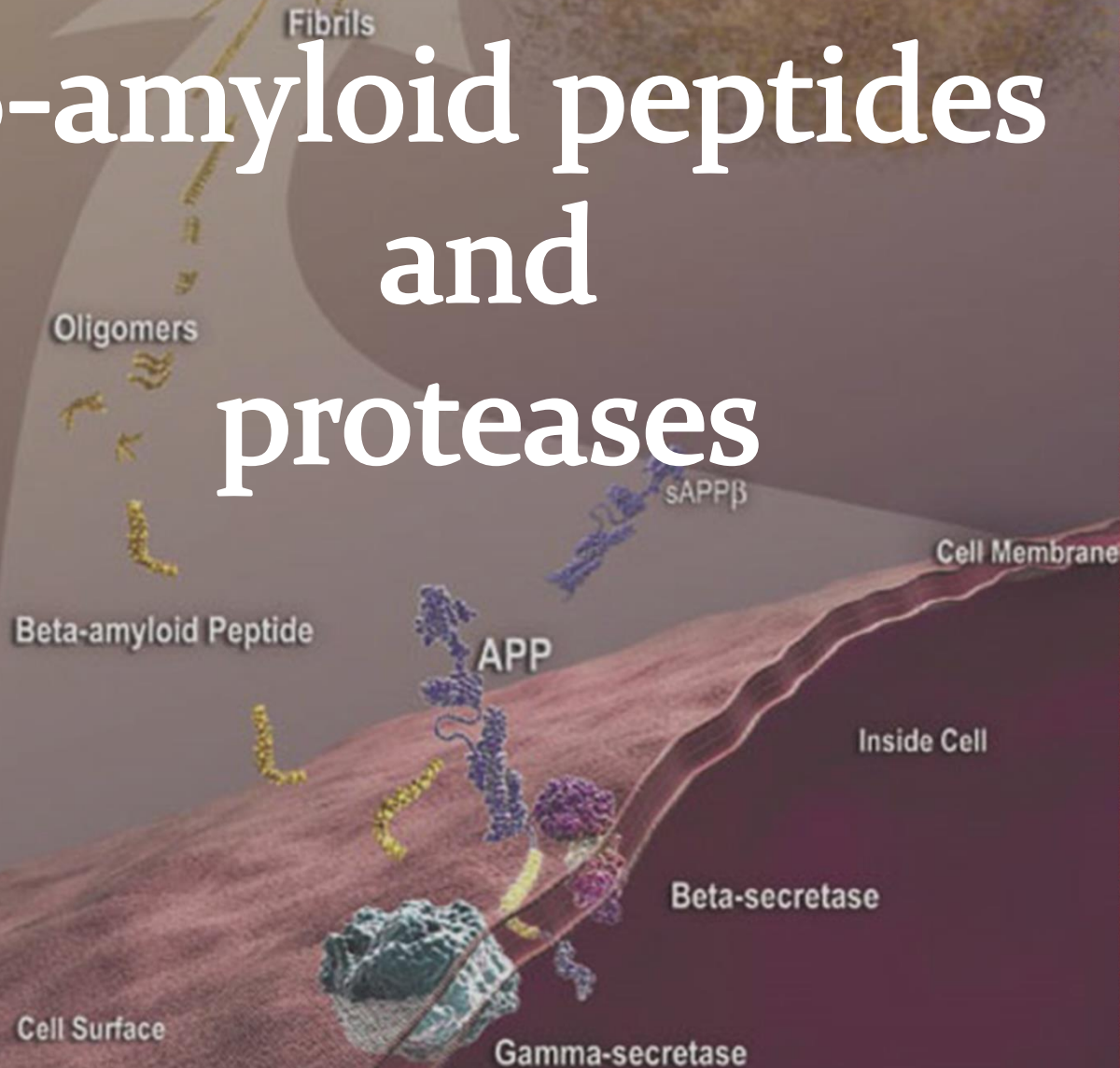




Università degli Studi di Ferrara  
Laurea in Scienze Biomolecolari e dell'Evolutione  
Corso di Macromolecole Biologiche

PLAQUE

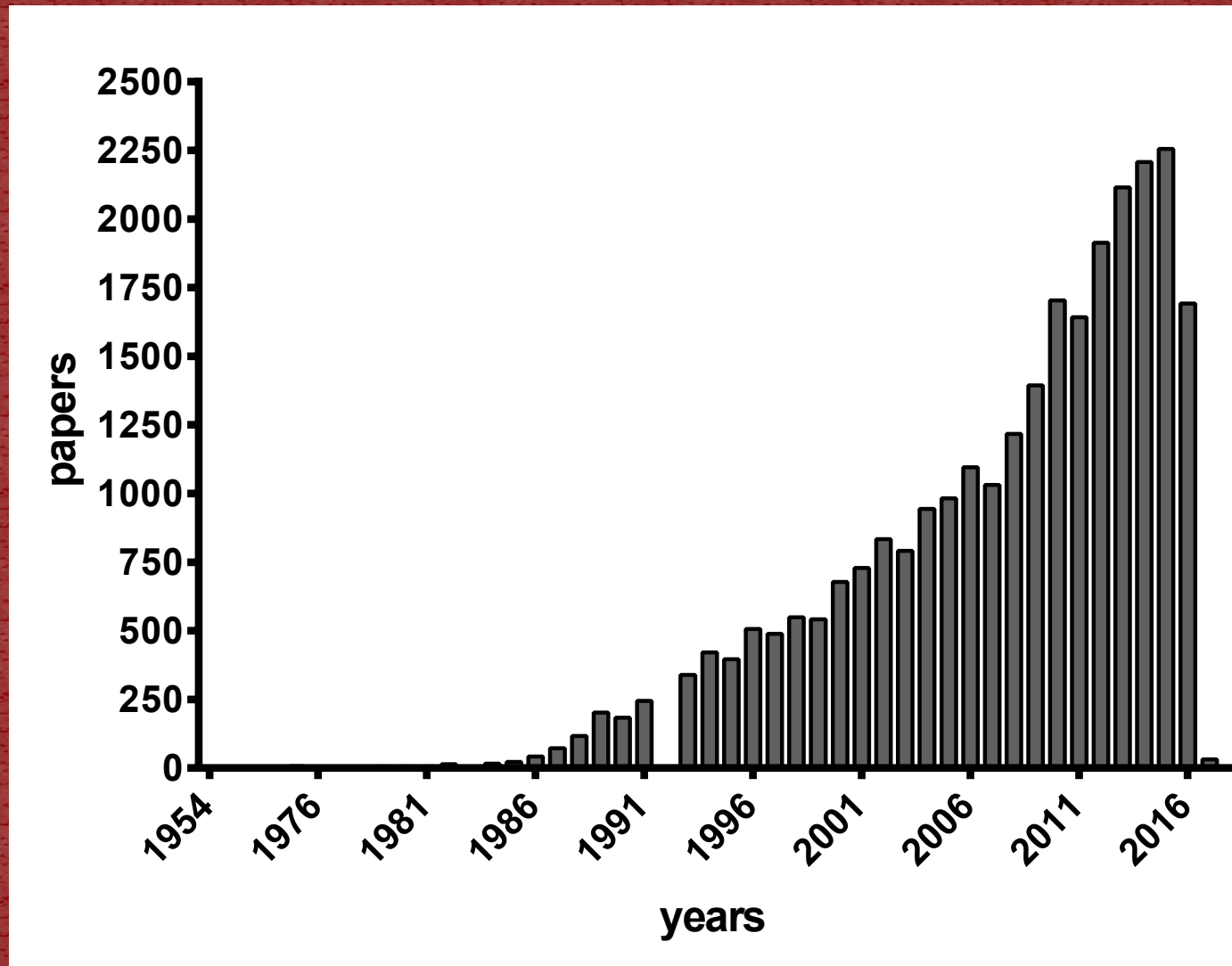
# $\beta$ -amyloid peptides and proteases



04/04/2017

Nicole Ziliotto

# Articles about “Alzheimer amyloid” over the years



# Problem statement

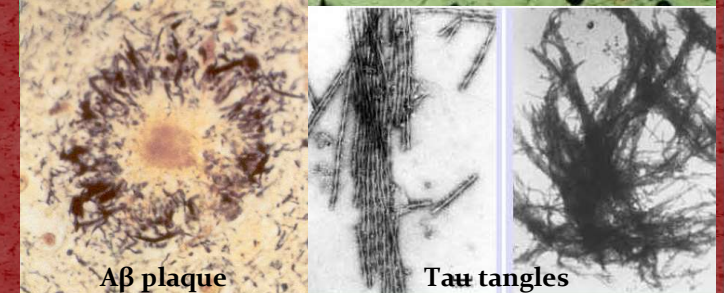
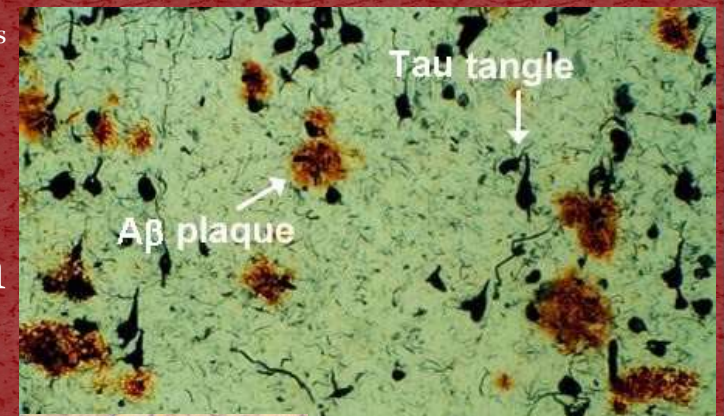


- 46.8 million people living with dementia in 2015 , this number is projected to reach 131.5 million by 2050.
- One new case of dementia every 3 seconds
- 1 in 10 people over age 65 and nearly half of people over 85 have Alzheimer's disease (AD).
- AD is the most common form of dementia with 60 to 80% of cases.
- Curative therapies are absent.

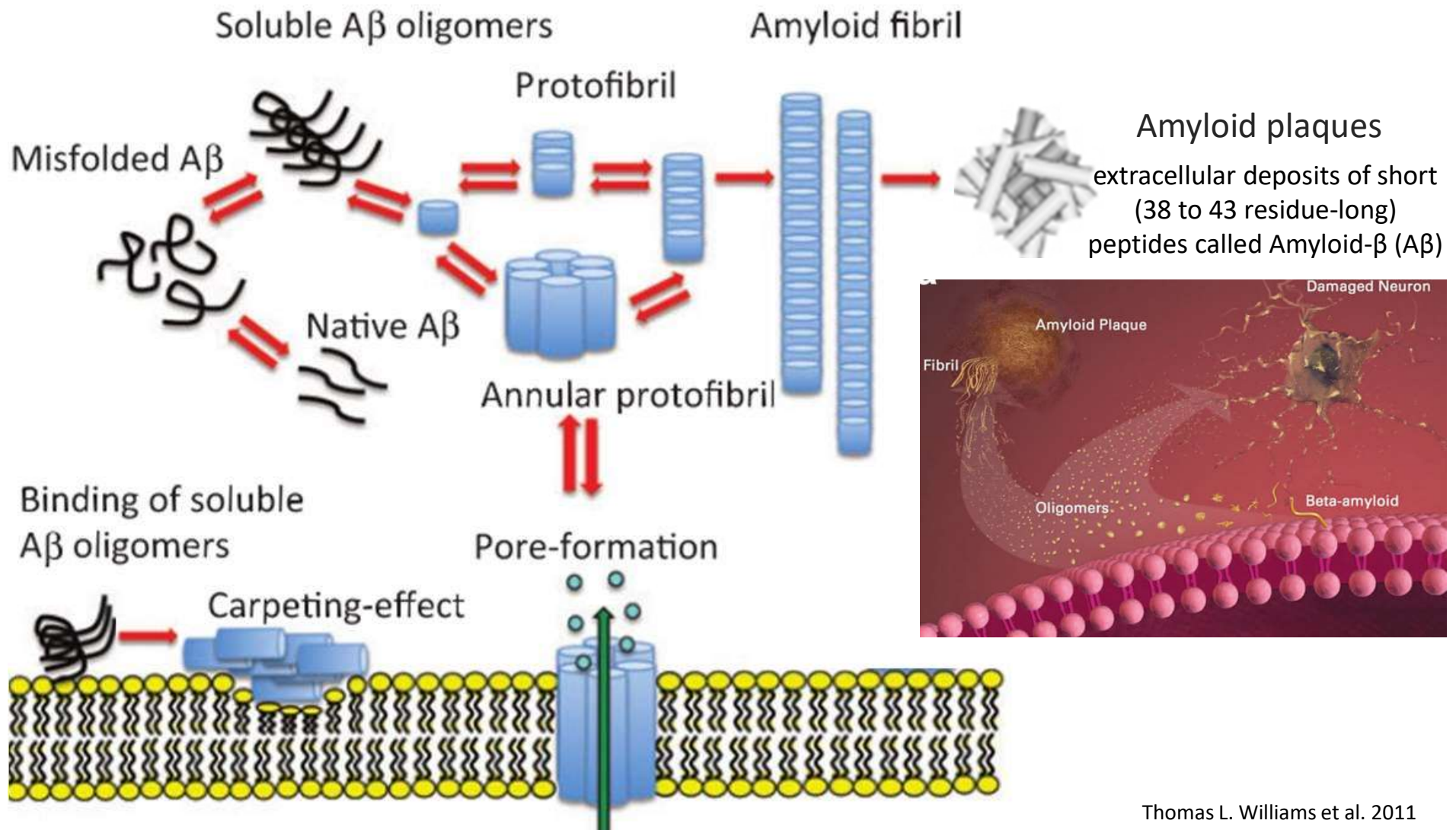
World Alzheimer Report 2016  
2016 Alzheimer's Disease facts and figures

Abundance of two abnormal structures in the brains of people with AD:

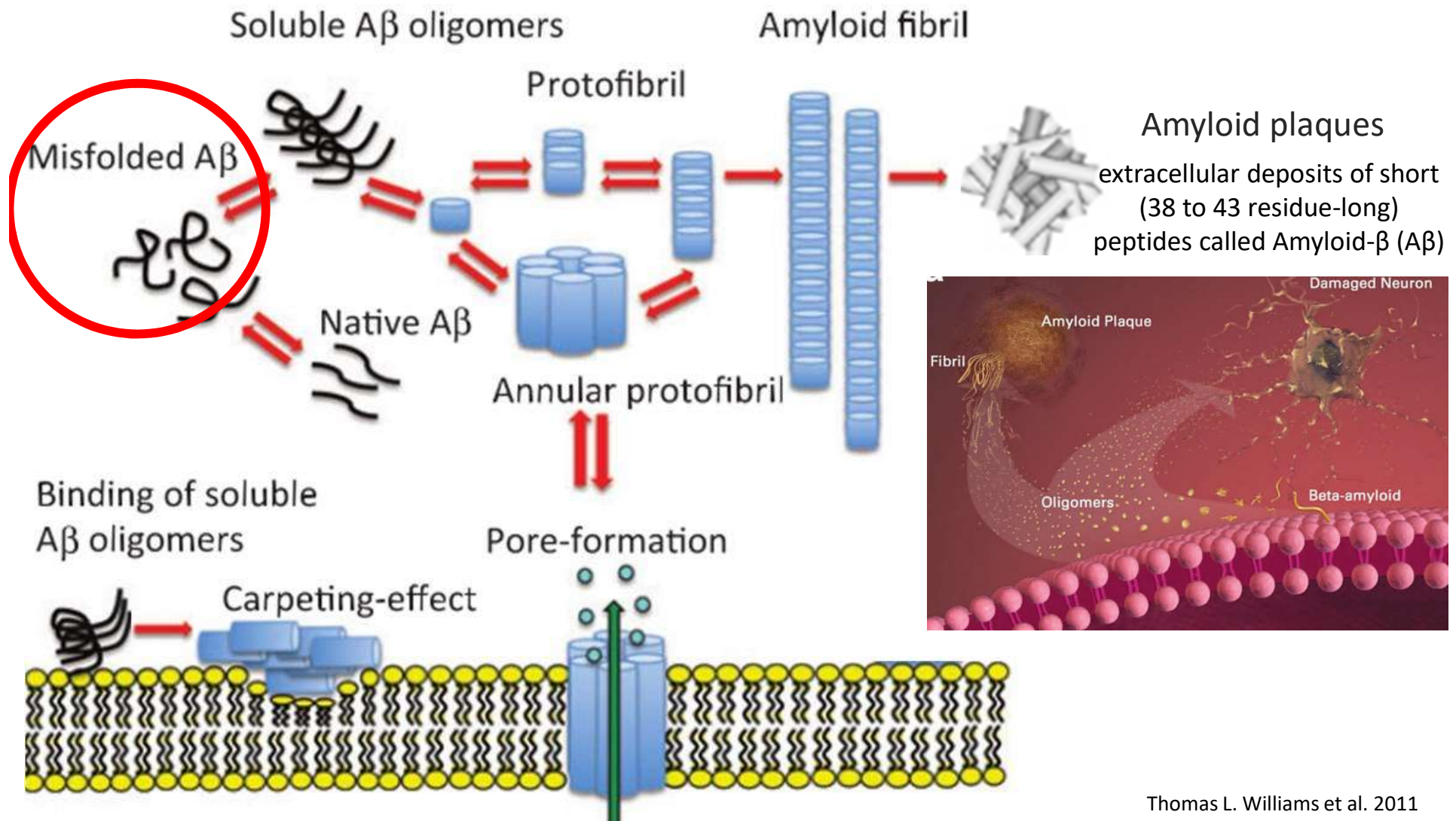
- Amyloid- $\beta$  ( $A\beta$ ) plaques, which are dense deposits of protein and cellular material that accumulate outside and around nerve cells
- Neurofibrillary TAU tangles, which are twisted fibers that build up inside the nerve cell



# $\beta$ -amyloid self-aggregating structure



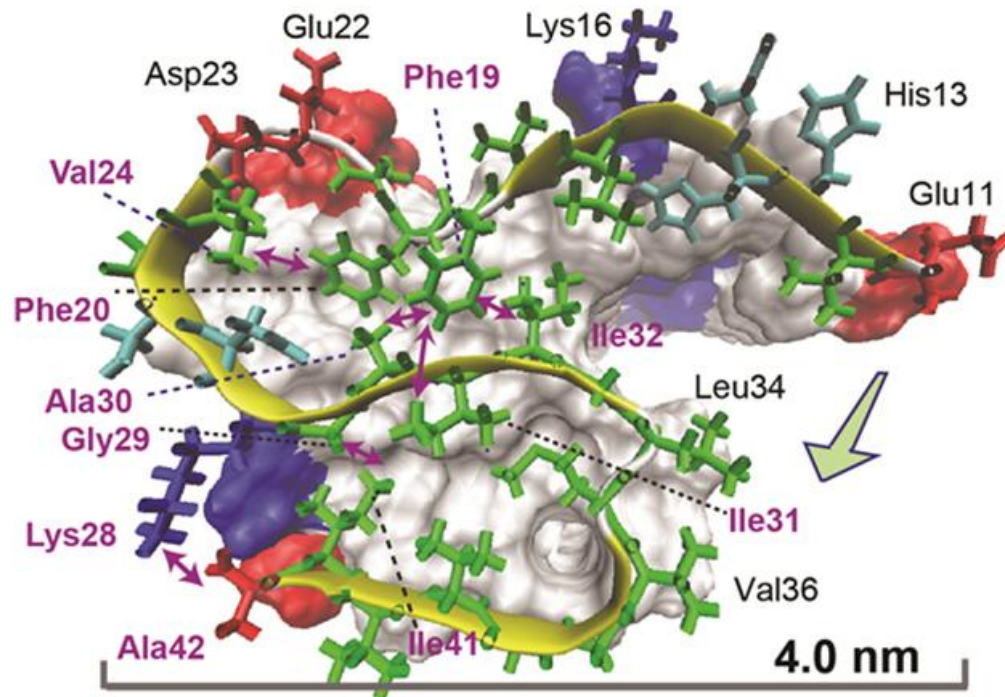
# $\beta$ -amyloid self-aggregating structure



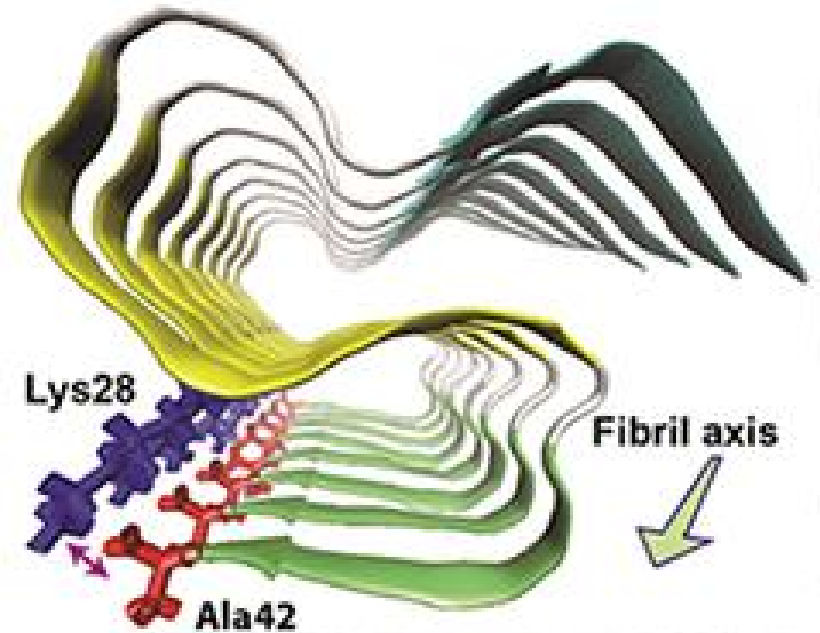
# Atomic structure model of A $\beta$ (1–42) fibrils

A $\beta$ 1-42 aggregates at a faster rate than A $\beta$ 1–40 due to its highly hydrophobic isoleucine and alanine at C-terminus

4 KDa



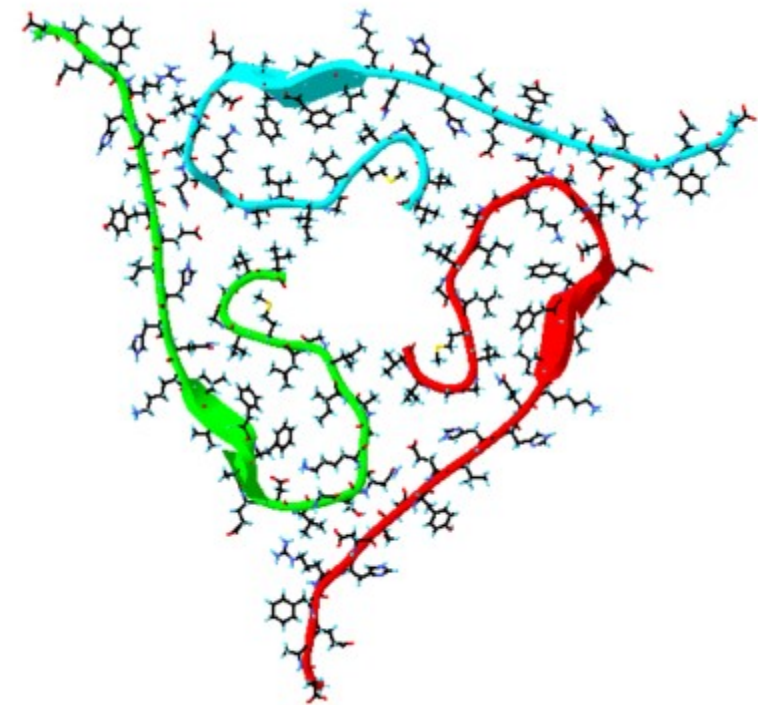
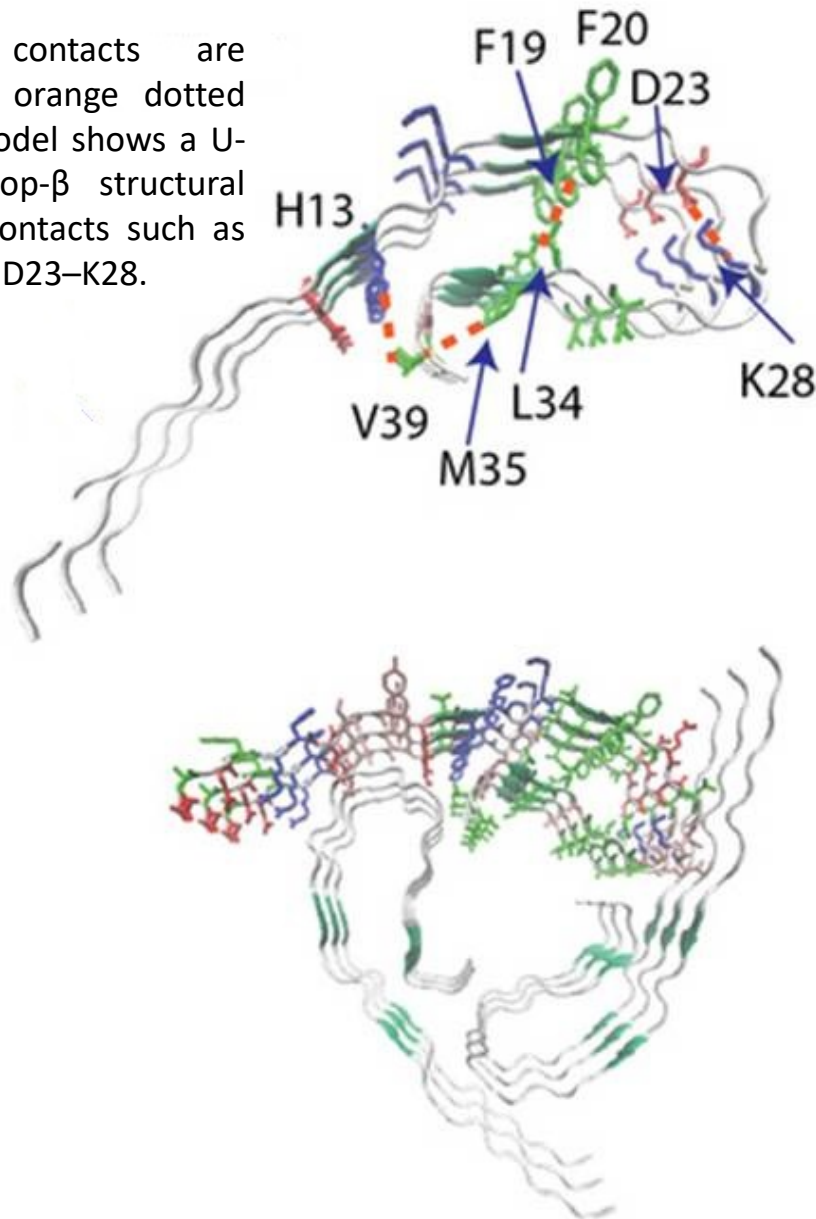
An atomic view of an A $\beta$ 42 molecule within a fiber reveals intramolecular connections (double-headed purple arrows) between residues in different parts of the S-shaped structure. Residues are colored as green, hydrophobic; cyan, polar; red, acidic; and blue, basic.



Three  $\beta$ -strand regions (cyan, residues 12–18; yellow, 24–33; green, 36–40) connected by two short coil or turn (white) regions. A salt bridge between Ala42 and Lys28 stabilizes the structure.

# Atomic structure model of A $\beta$ (1–40) fibrils

Side-chain contacts are denoted by orange dotted lines. The model shows a U-shaped  $\beta$ -loop- $\beta$  structural motif with contacts such as F19–L34 and D23–K28.

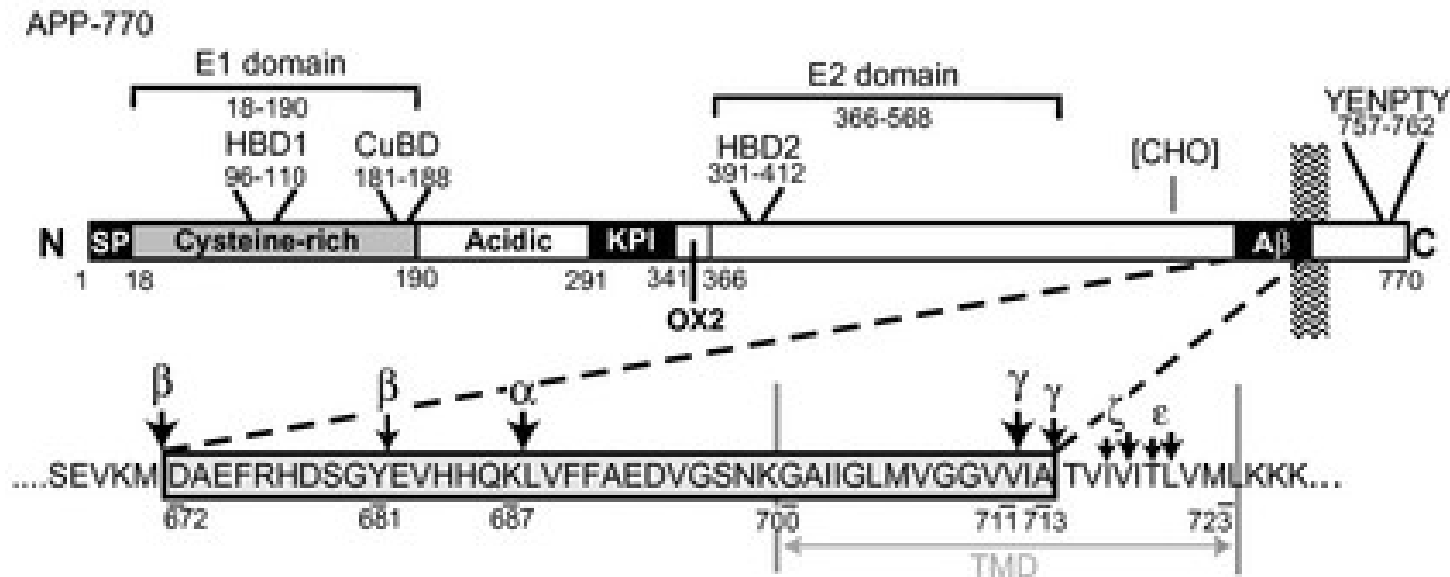


A $\beta$ 40 peptide monomers tend to aggregate in oligomers multiple of three units (trimers, hexamers, nonamers and dodecamers), where the N-termini are exposed to the solvent, while the hydrophobic C-termini, are buried in the trimer core.

# Amyloid Protein Precursor (APP)

## Protein Function:

- Cellular proliferation and differentiation
- Neurite outgrowth
- Synaptogenesis
- Synaptic plasticity
- Inhibition blood coagulation
- Signal transduction
- Gene regulation
- Trafficking



HBD: Heparin-binding domain  
 KPI: Kunitz protease inhibitor  
 CHO: Copper-binding domain

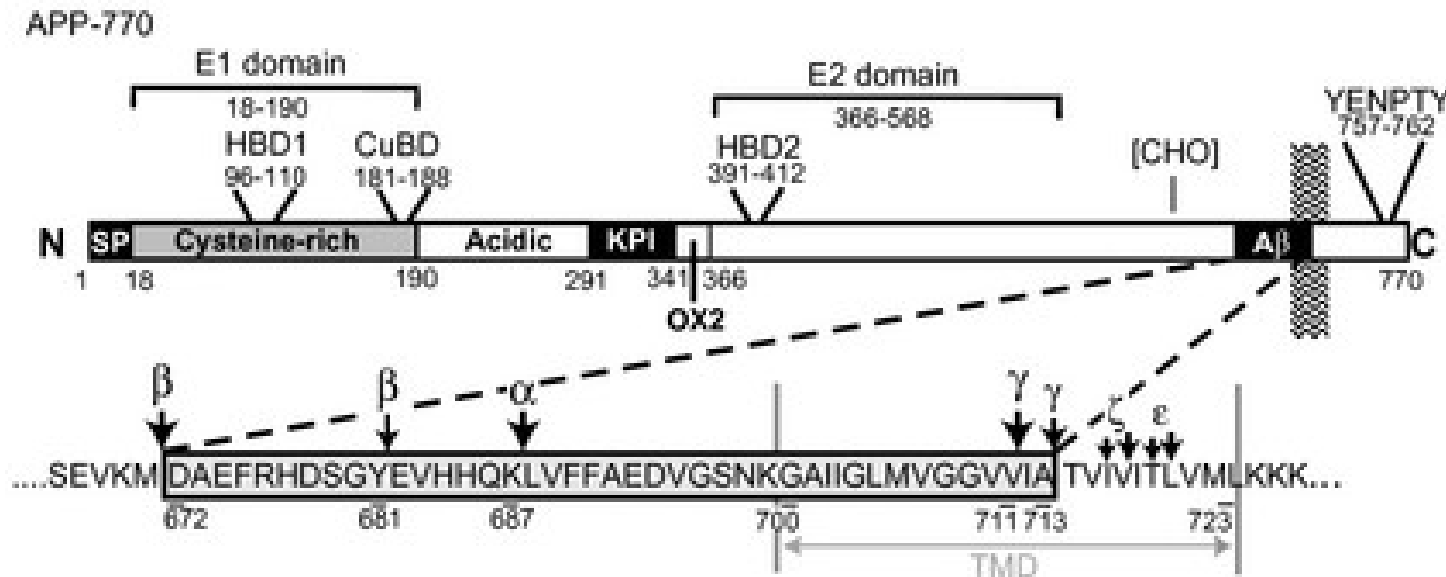
110-135 KDa



# Amyloid Protein Precursor (APP)

## Protein Function:

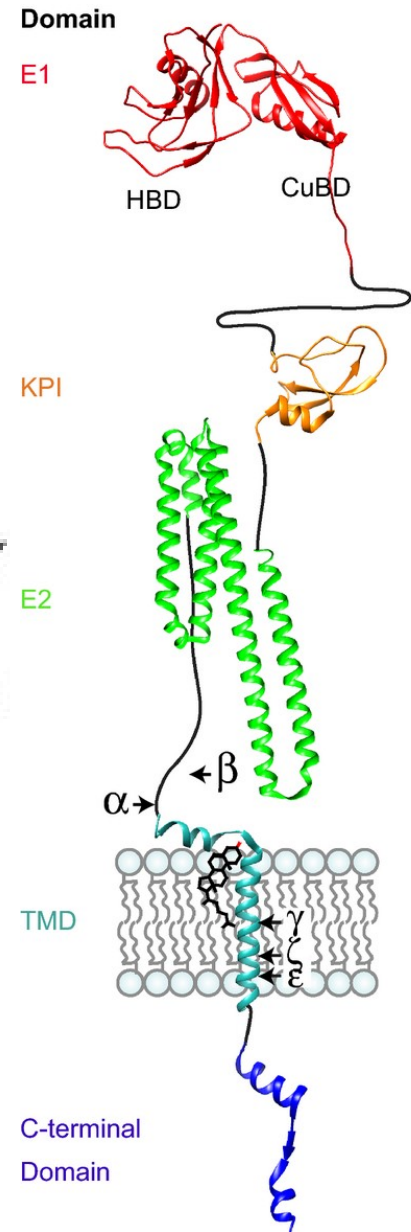
Cellular proliferation and differentiation  
 Neurite outgrowth  
 Synaptogenesis  
 Synaptic plasticity  
 Inhibition blood coagulation  
 Signal transduction  
 Gene regulation  
 Trafficking



**mutations in APP  
 increase Aβ deposition  
 (e.g. Swedish mutation)**

HBD: Heparin-binding domain  
 KPI: Kunitz protease inhibitor  
 CHO: Copper-binding domain

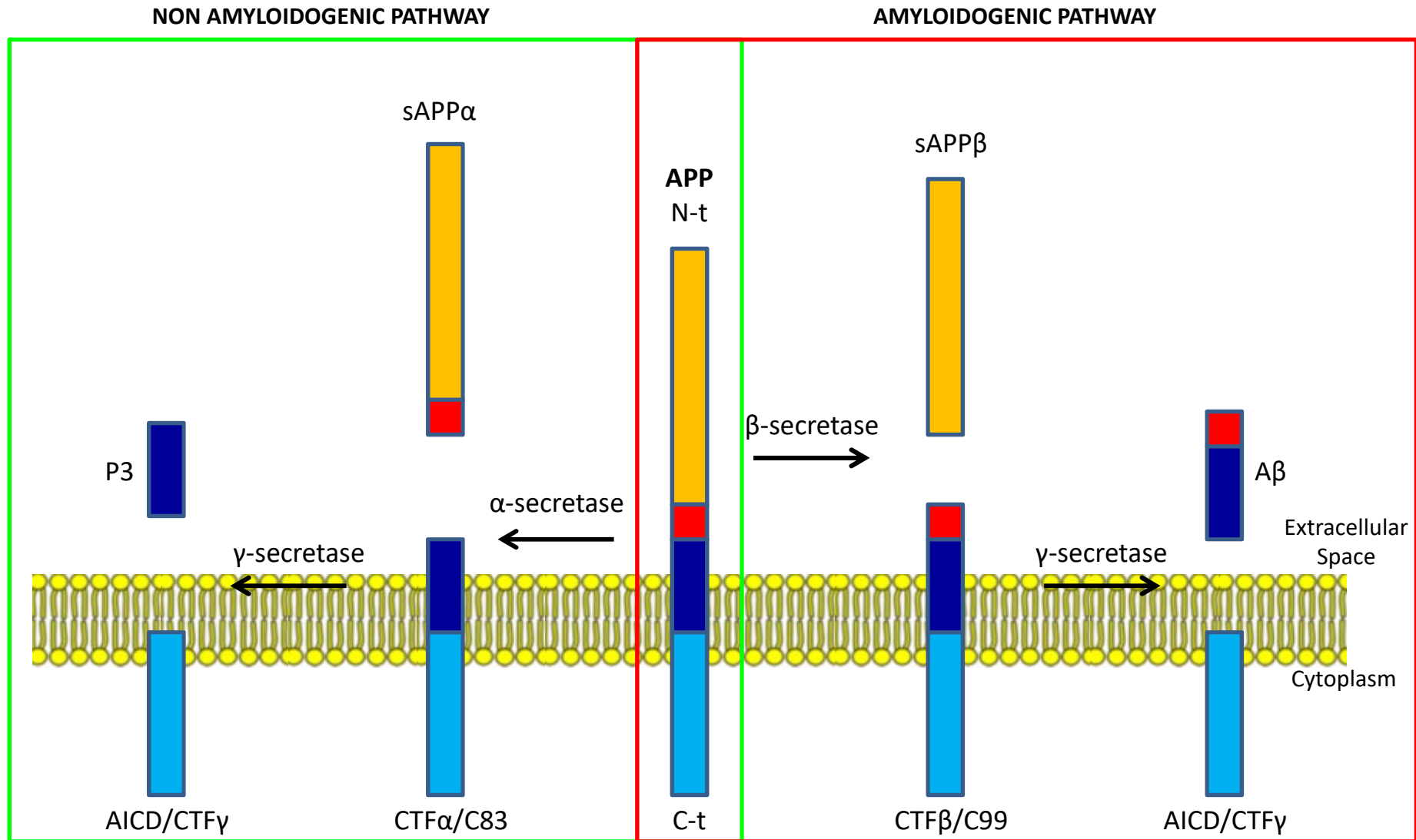
110-135 KDa



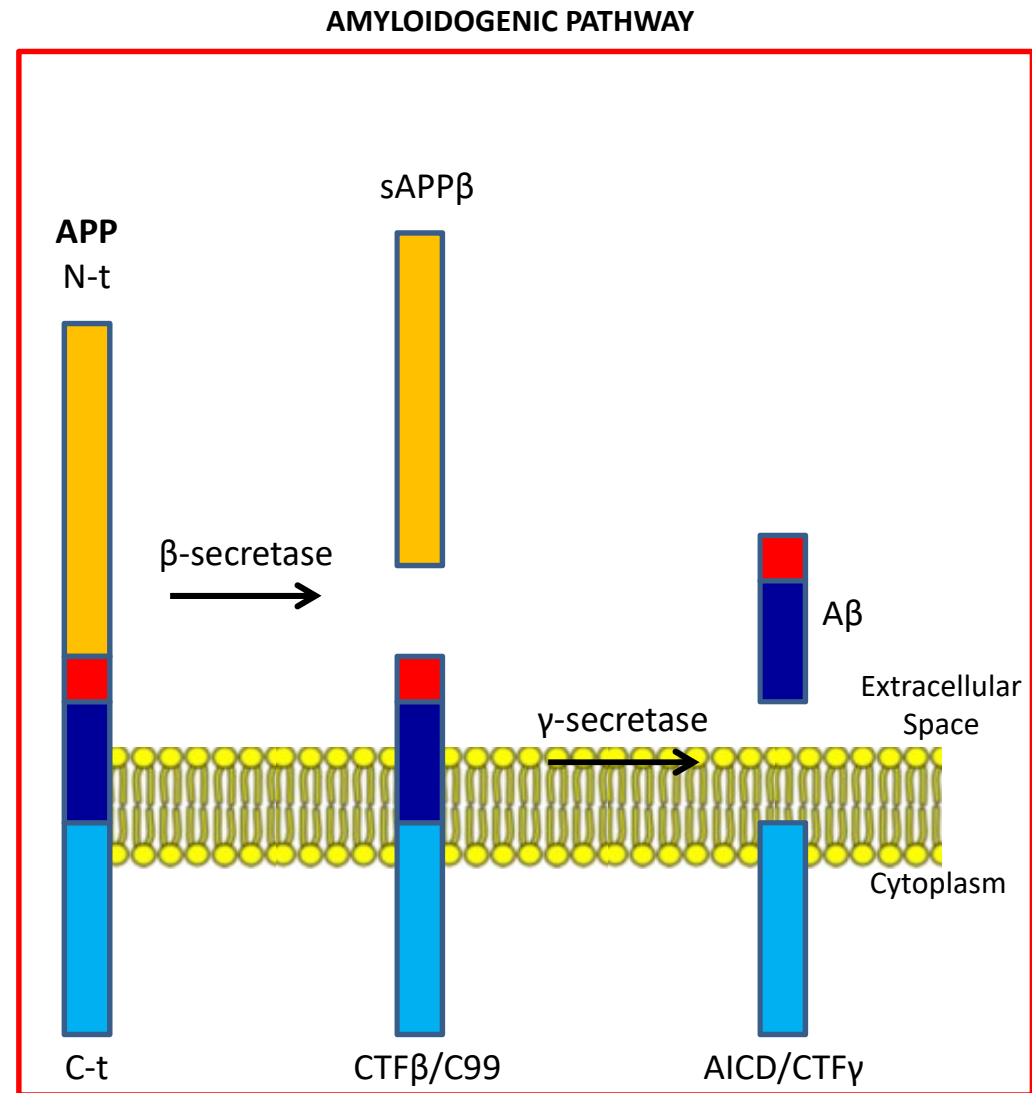
# Amyloid Plaque Formation

- Amyloid plaques are extracellular deposits of short (38 to 43 residue-long) peptides called amyloid- $\beta$  ( $A\beta$ ).
- $A\beta$  peptides derived from amyloid precursor protein (APP).
- APP is a membrane glycoprotein that normally behave in the brain as a cell surface signaling molecule.
- The hydrophobicity, net charge and the sequence propensity to form secondary structures, have been shown to modulate amyloidogenicity. In fact  $A\beta_{42}$  aggregates at a faster rate than  $A\beta_{40}$ .
- Neurotoxic  $A\beta$  assemblies contain a high level of  $\beta$ -sheet conformation.
- $A\beta$  oligomers are capable of seeding their own replication and may be analogous to different strains of prions.

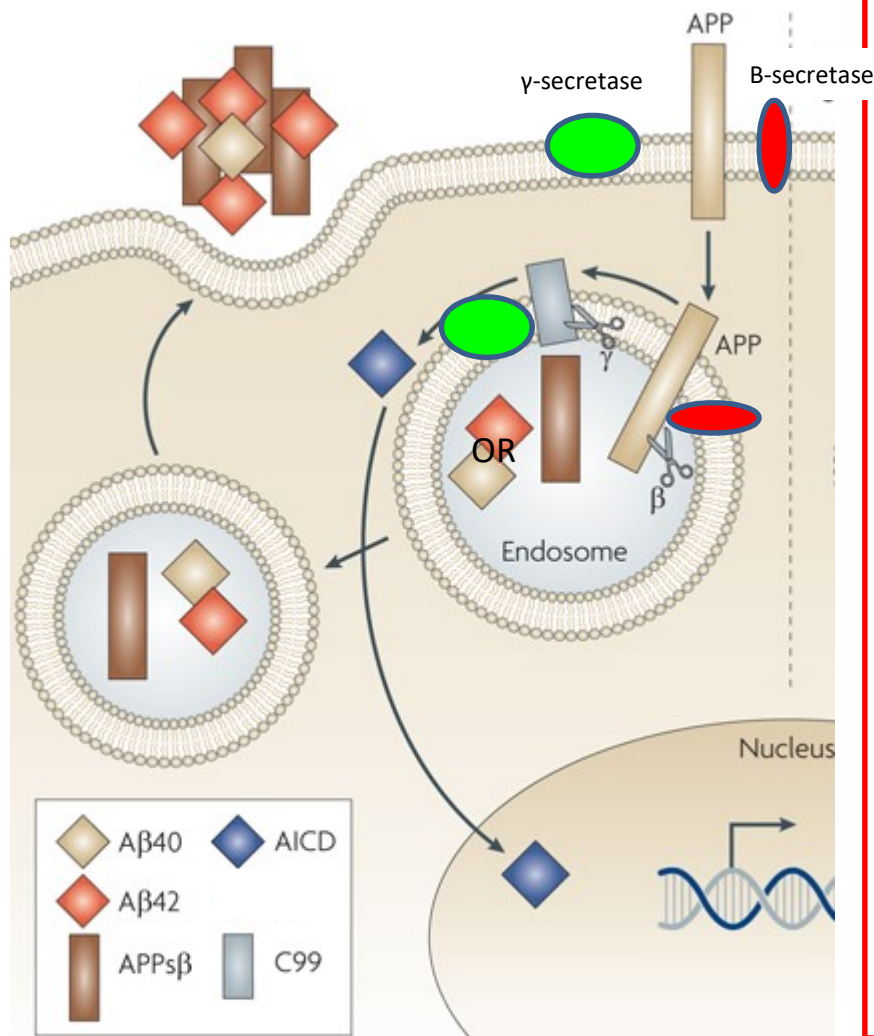
# Amyloid Hypothesis



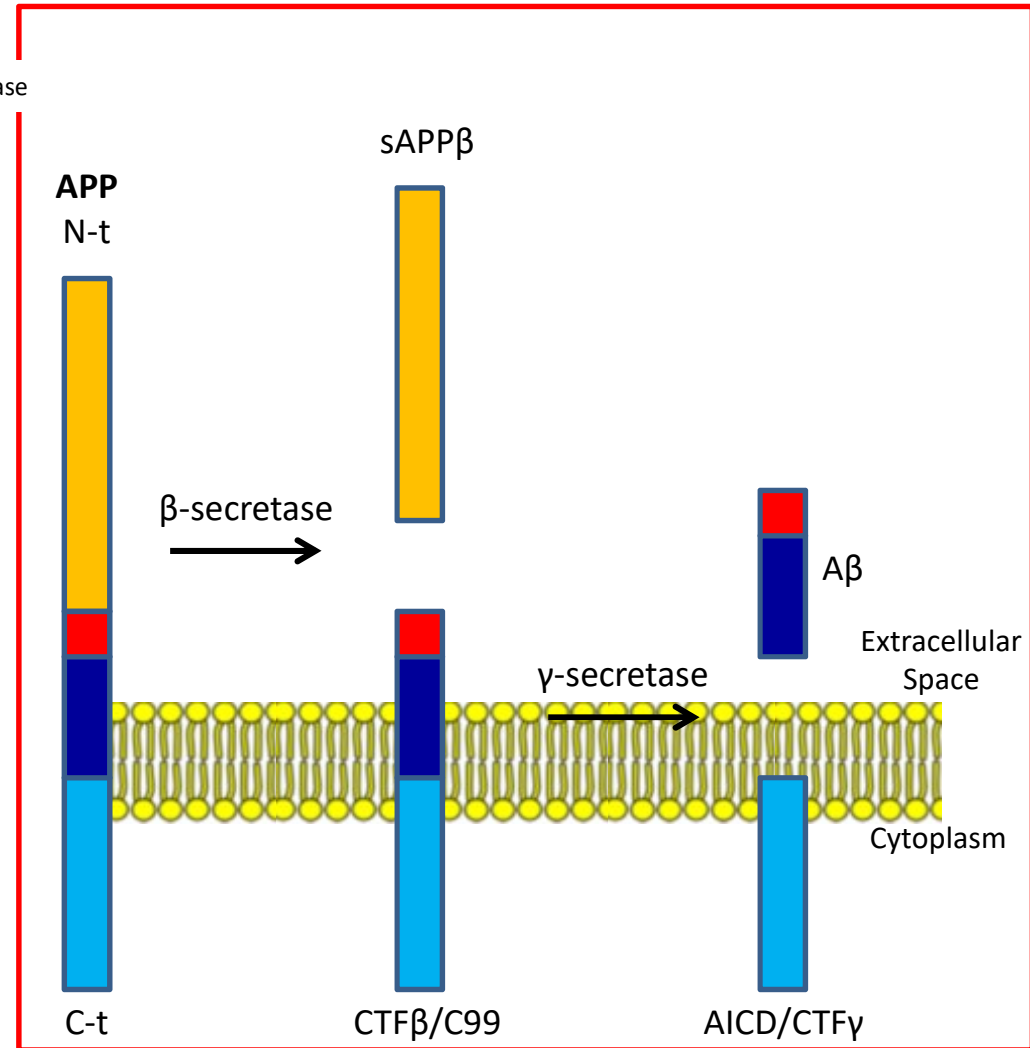
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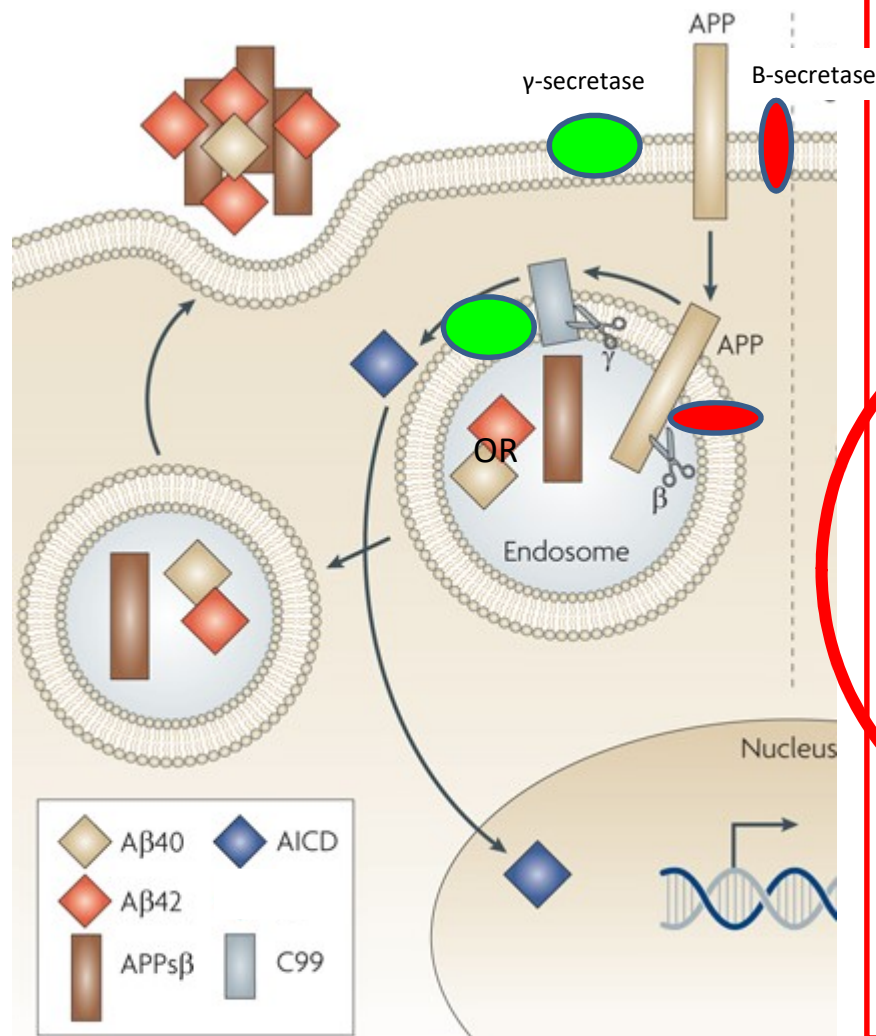
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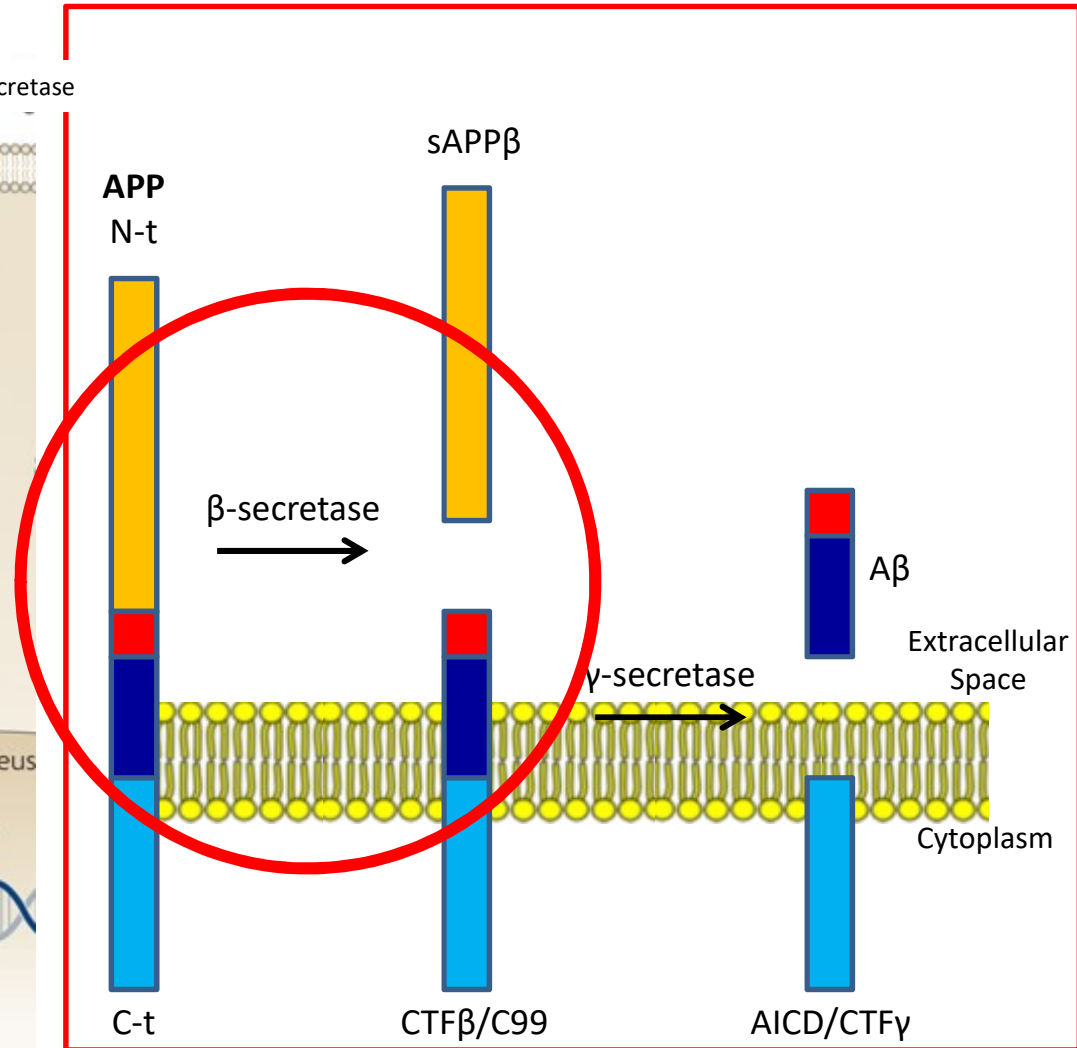
AMYLOIDOGENIC PATHWAY



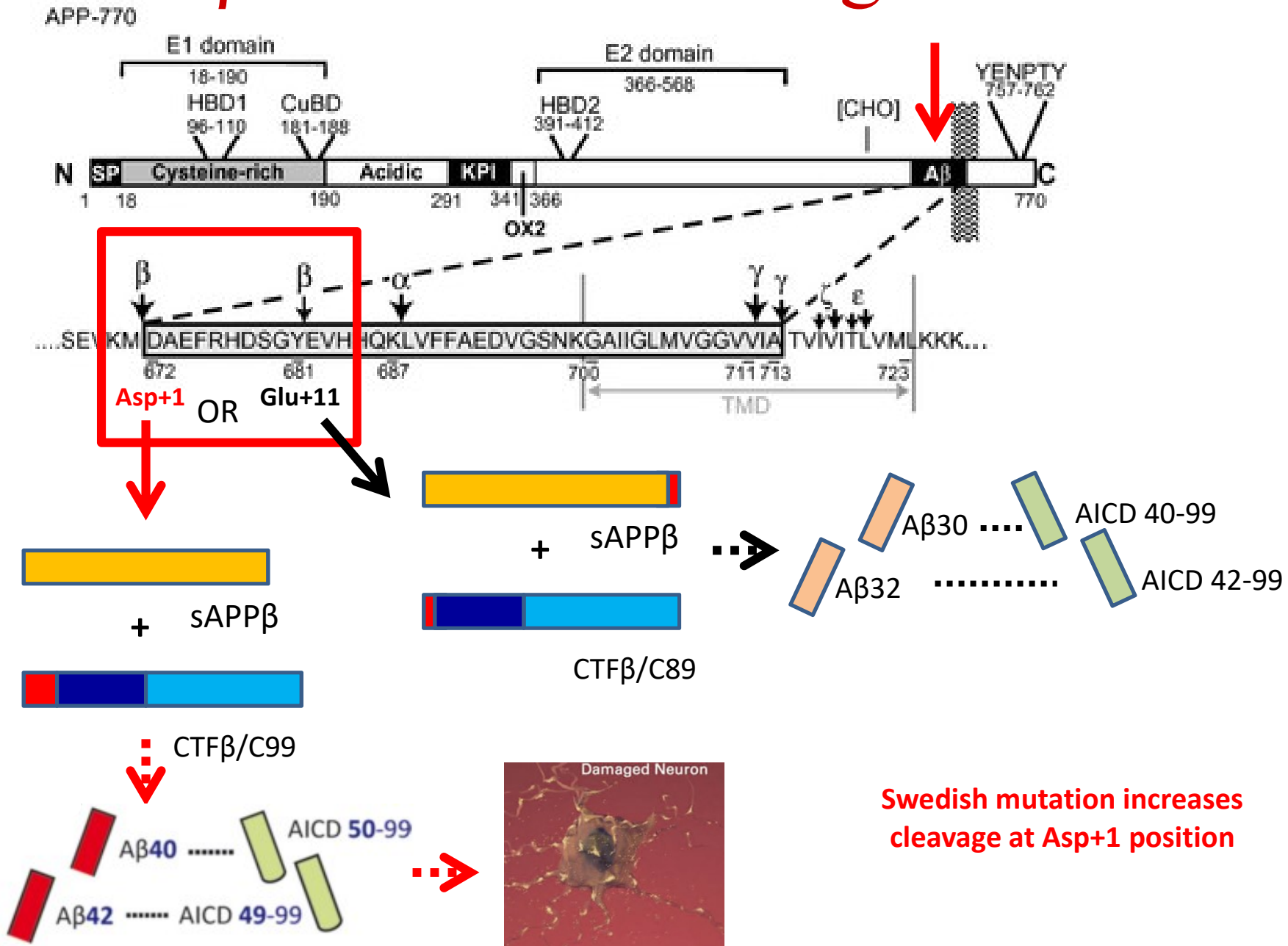
# Amyloid Hypothesis



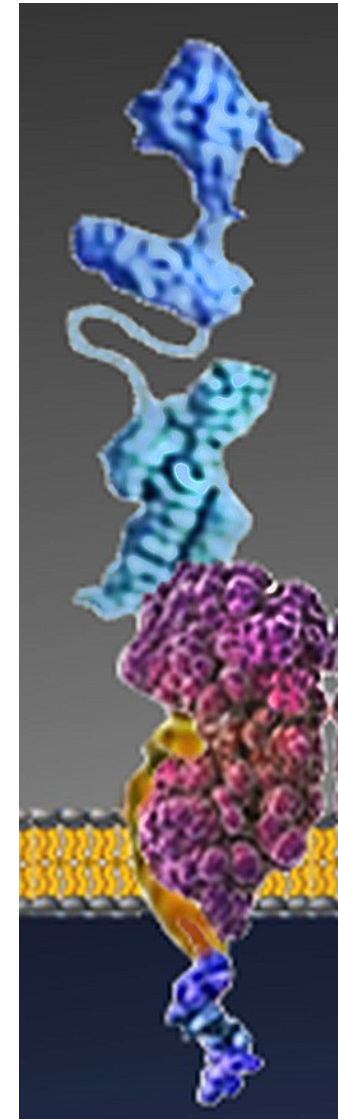
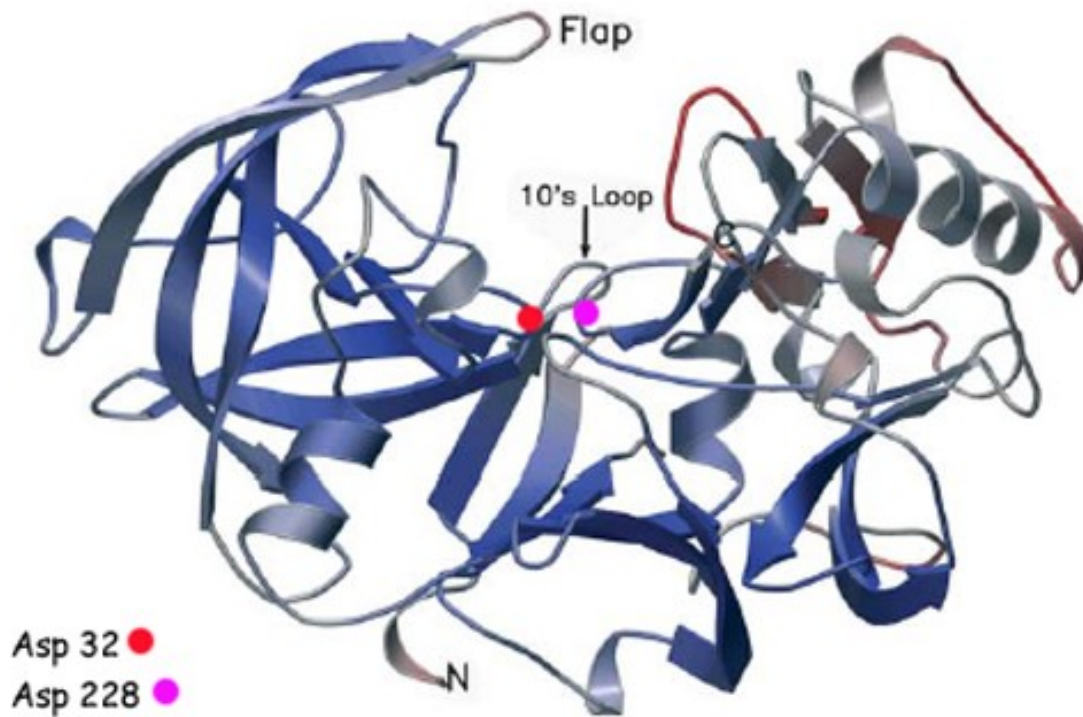
AMYLOIDOGENIC PATHWAY



# $\beta$ -secretase cleavage of APP

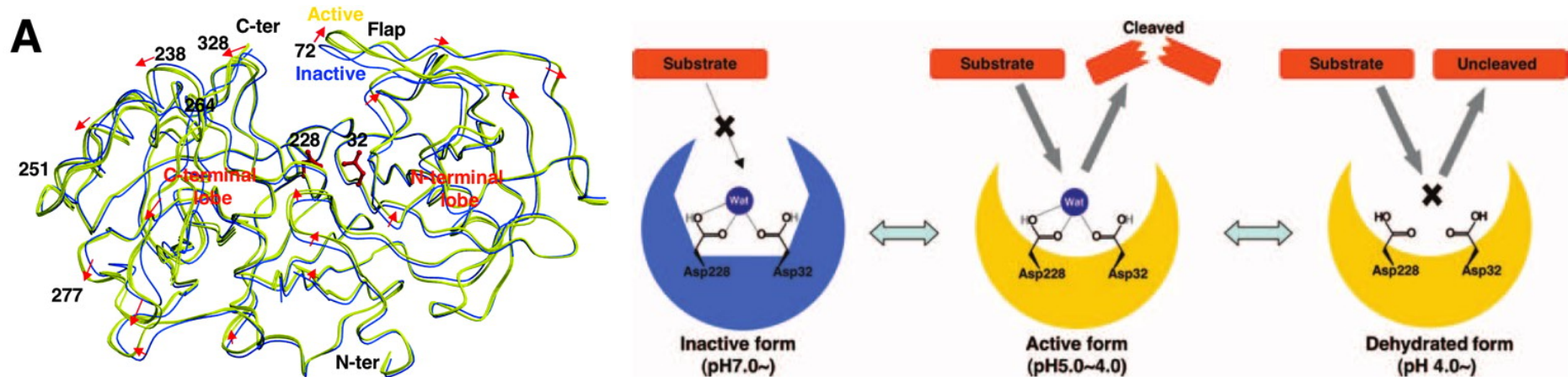


# $\beta$ -Secretase: Beta Amyloid-site-Cleaving Enzyme 1 (BACE1)





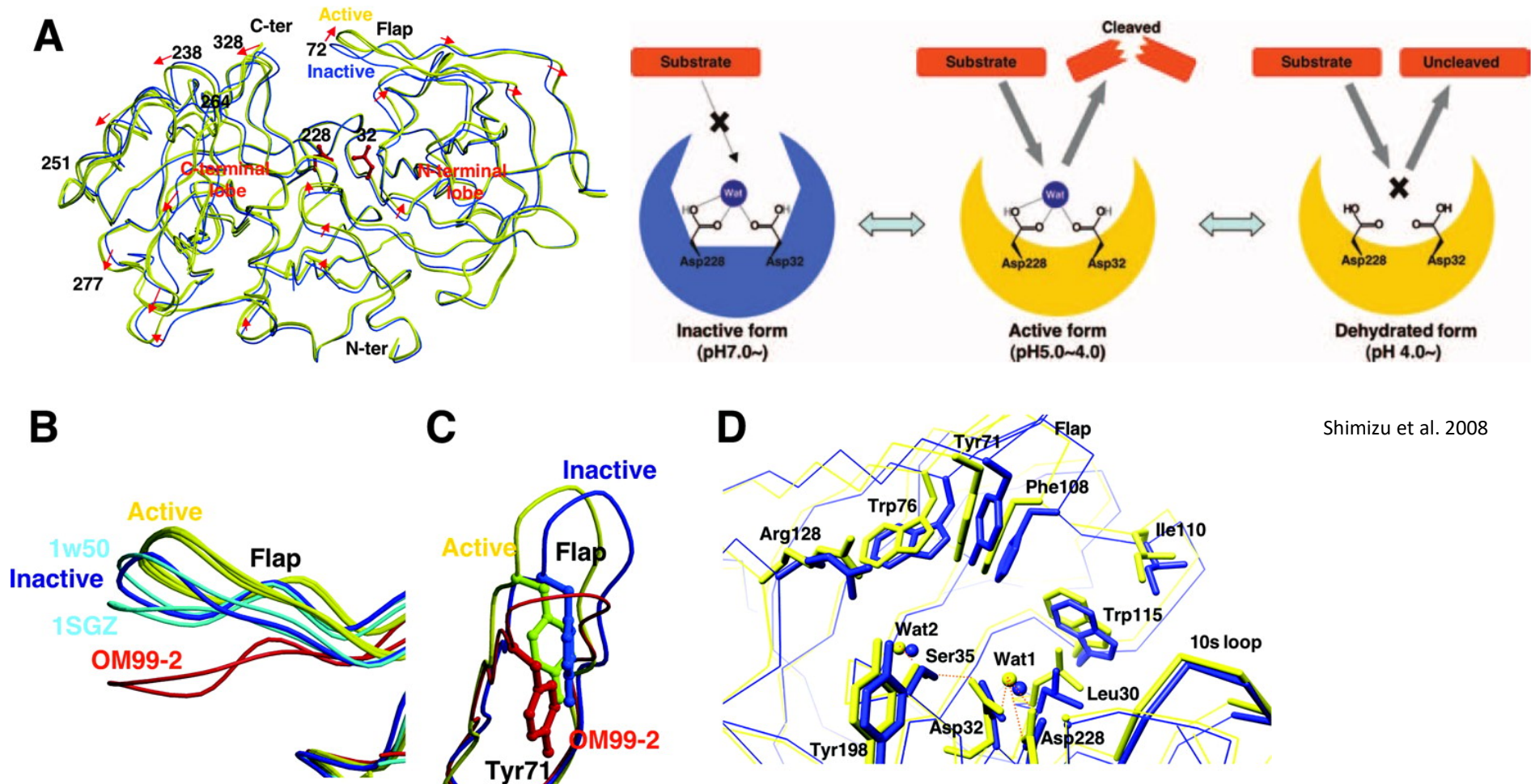
# Conformational changes associated with activation of BACE1



Shimizu et al. 2008

In the aspartic proteases there are two conserved water molecules. The first water molecule (Wat1) is located between the Asp pair of Asp32 and Asp228 of BACE1. The second water molecule (Wat2) is involved in the hydrogen bond, with a conserved Tyr residue in the flap. Wat2 also participates in a conserved hydrogen-bonding network Wat2-Ser35-Asp32-Wat1-Asp228 and was proposed to assist in the catalytic reaction.

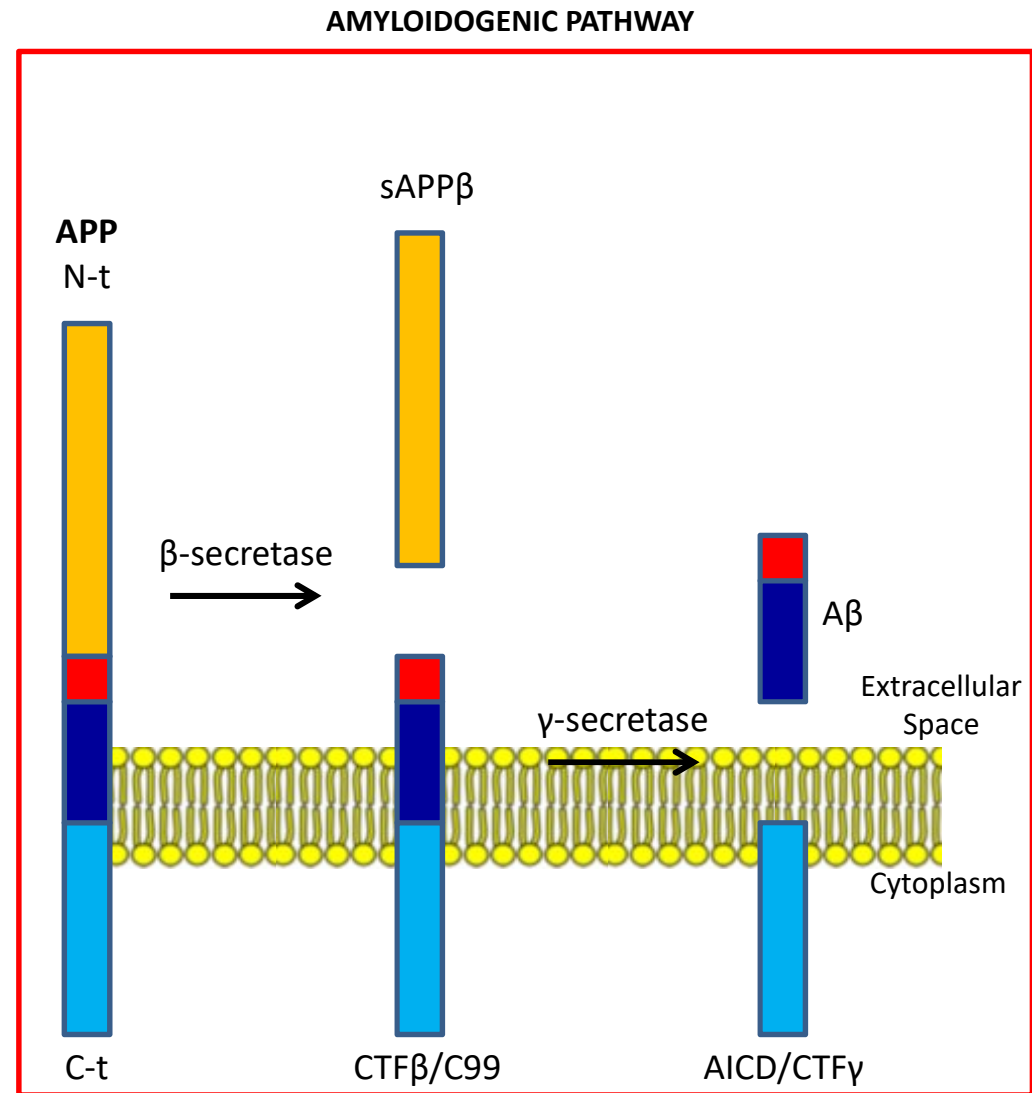
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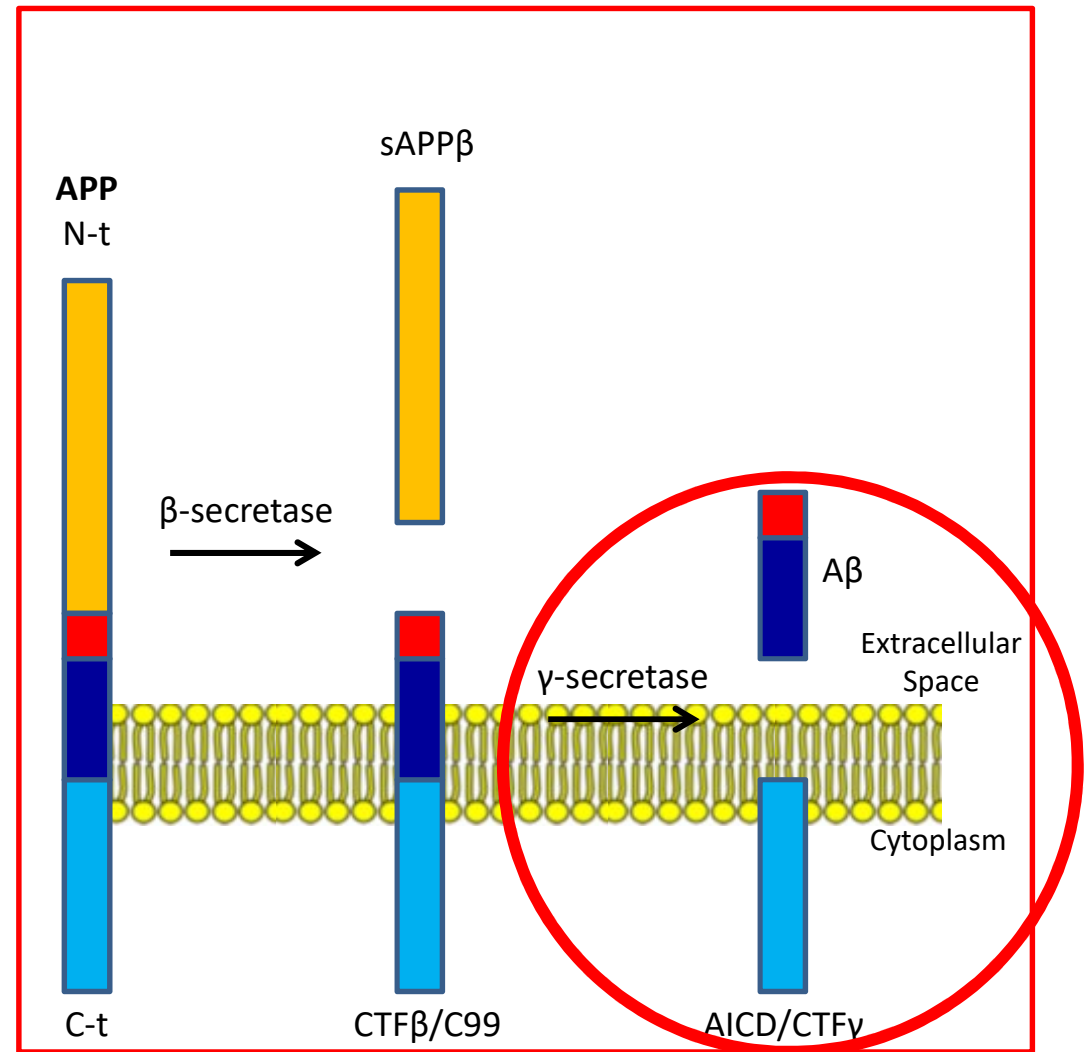
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# Amyloid Hypothesis

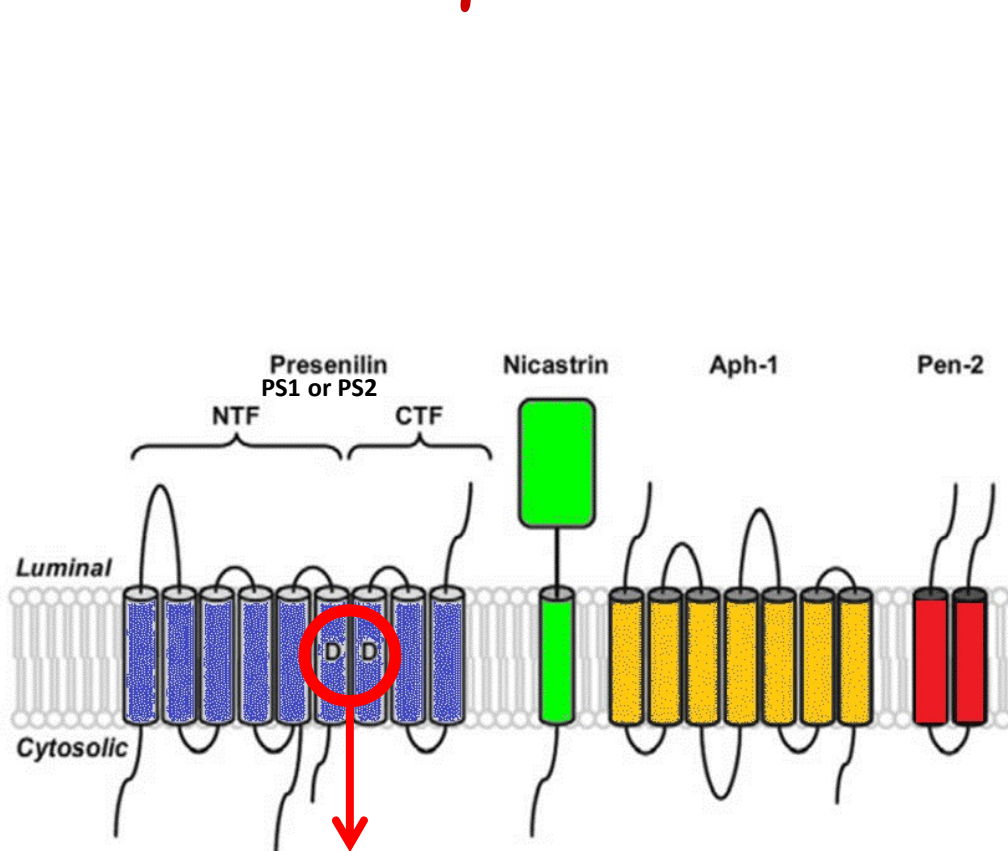


# Amyloid Hypothesis

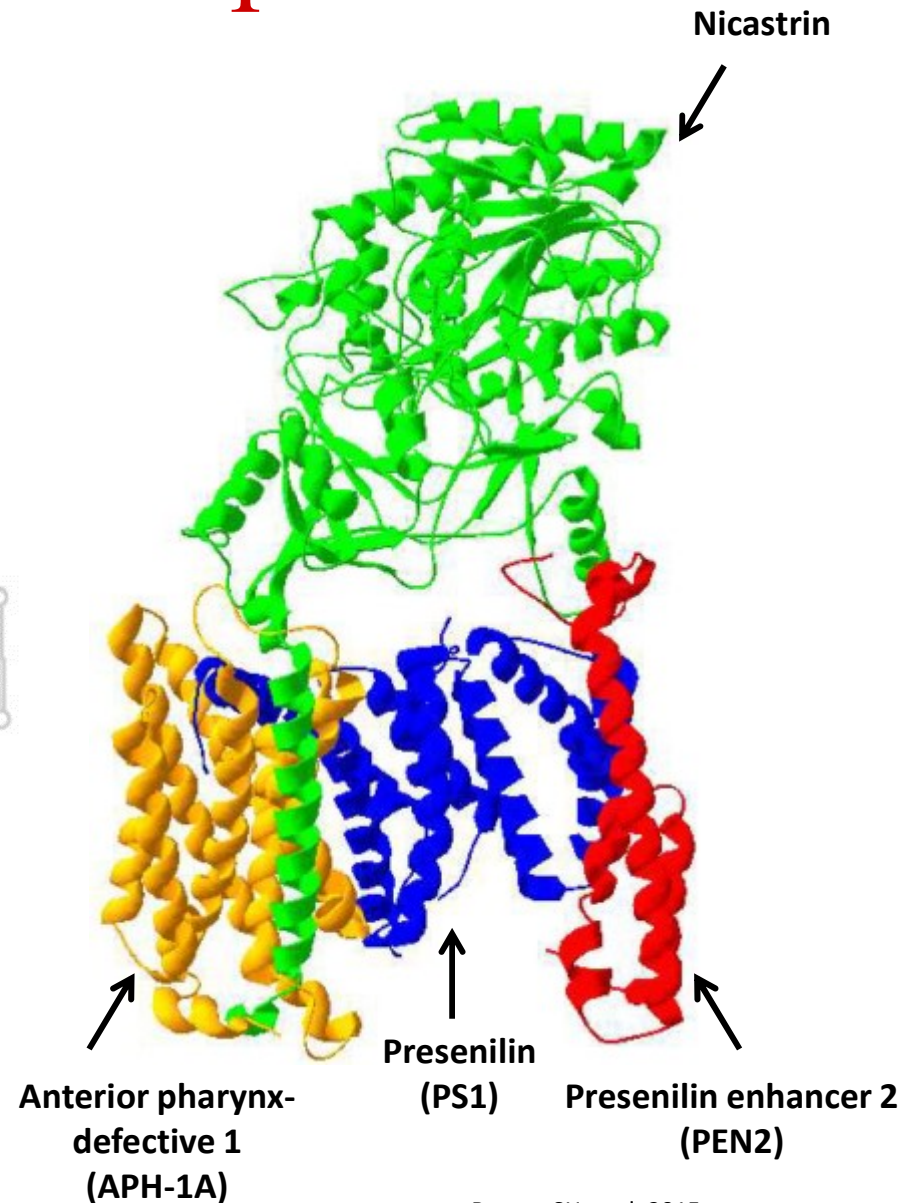
AMYLOIDOGENIC PATHWAY



# $\gamma$ -Secretase Complex

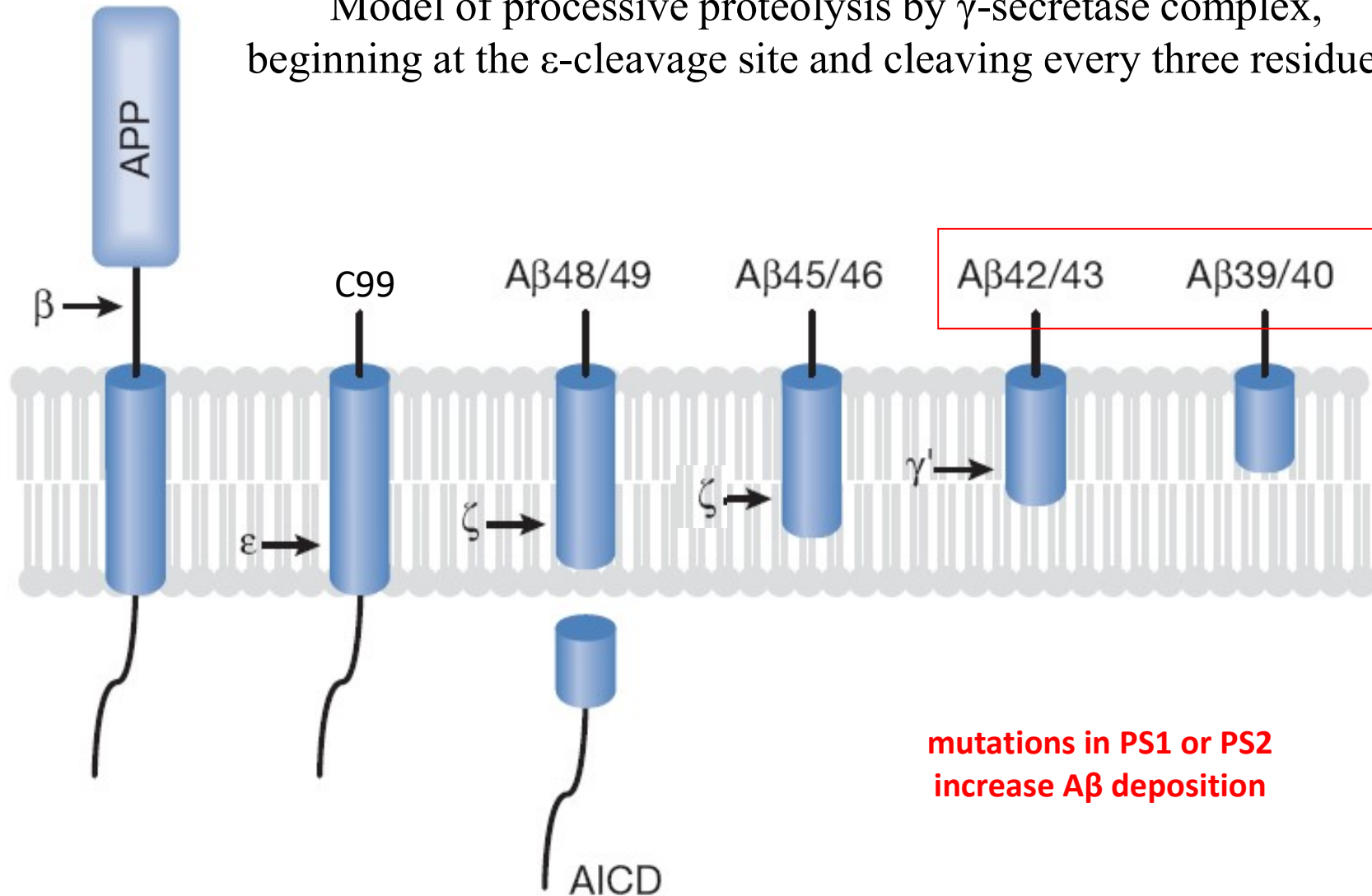


Asp257 and Asp385 are the PS1 catalytic residue

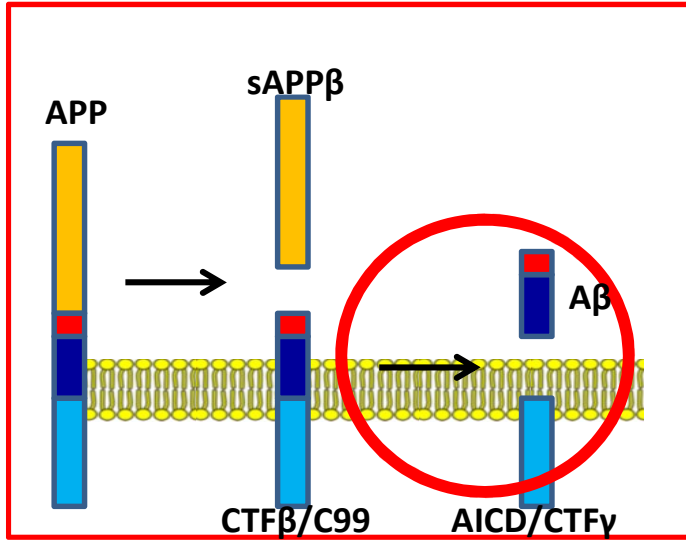


# $\gamma$ -Secretase cleavage

Model of processive proteolysis by  $\gamma$ -secretase complex, beginning at the  $\epsilon$ -cleavage site and cleaving every three residues.



AMYLOIDOGENIC  
PATHWAY

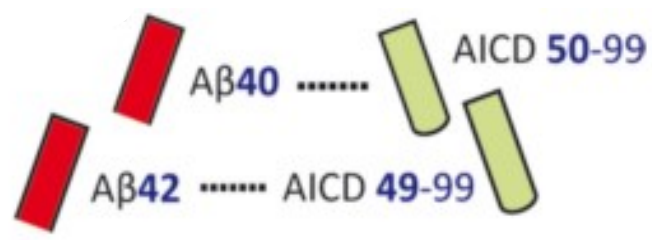
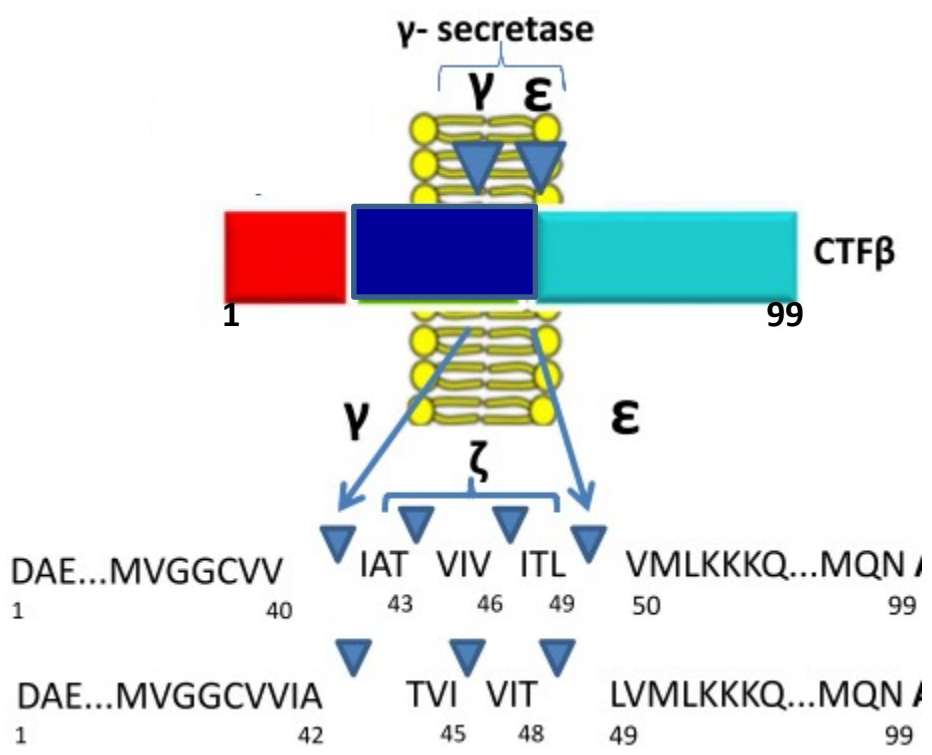


# γ-Secretase cleavage

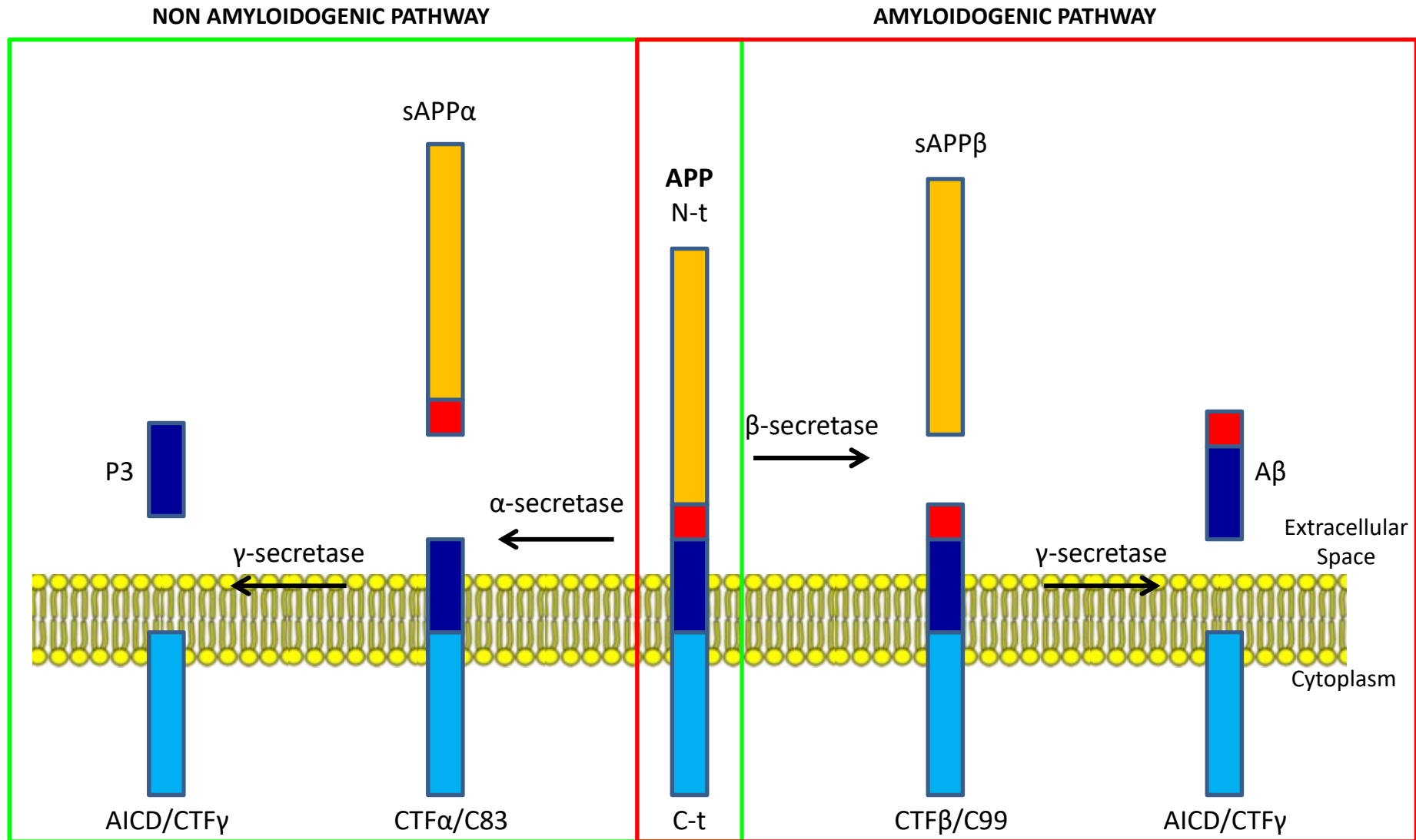
Three cleavages:  
1<sup>st</sup> the ε  
2<sup>nd</sup> the ζ  
3<sup>rd</sup> the γ



Alternative products:

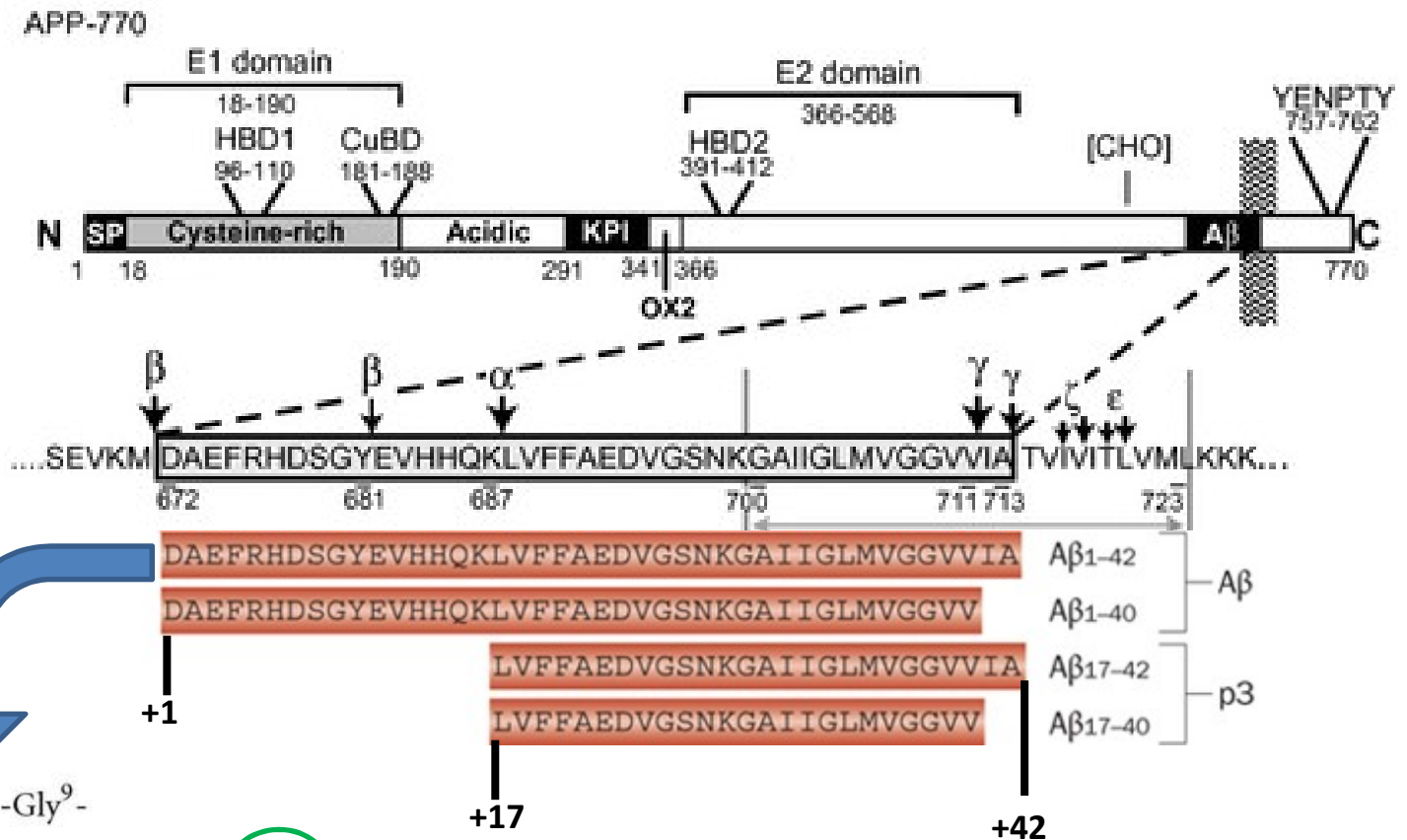


# Amyloid Hypothesis





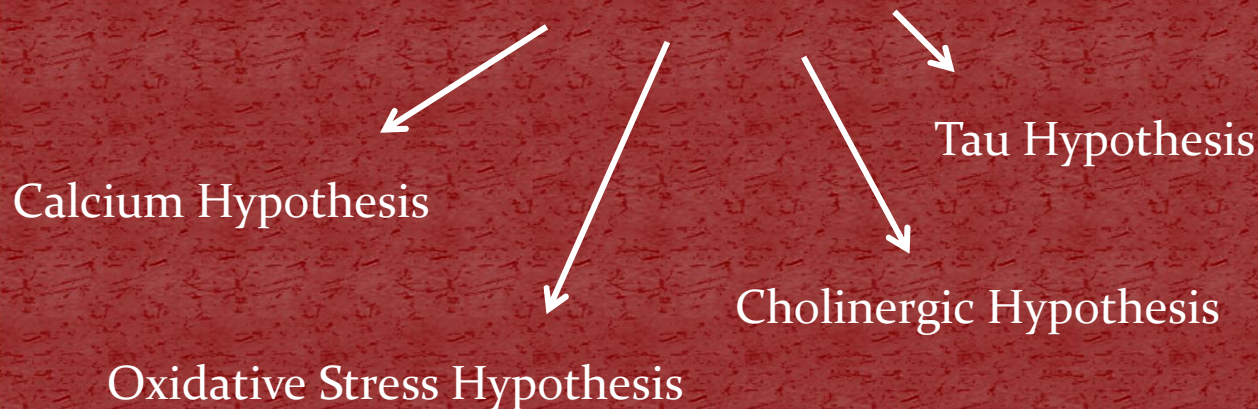
# Comparison between fragments



Aβ(1-42)

H<sub>2</sub>N-Asp<sup>1</sup>-Ala<sup>2</sup>-His<sup>6</sup>-Asp<sup>7</sup>-Ser<sup>8</sup>-Gly<sup>9</sup>-  
 Tyr<sup>10</sup>-Glu<sup>11</sup>-Val<sup>12</sup>-His<sup>13</sup>-His<sup>14</sup>-Gln<sup>15</sup>-Lys<sup>16</sup>-Leu<sup>17</sup>-  
 Val<sup>18</sup>-Phe<sup>19</sup>-Phe<sup>20</sup>-Ala<sup>21</sup>-Glu<sup>22</sup>-Asp<sup>23</sup>-Val<sup>24</sup>-Gly<sup>25</sup>-  
 Ser<sup>26</sup>-Asn<sup>27</sup>-Lys<sup>28</sup>-Gly<sup>29</sup>-Ala<sup>30</sup>-Ile<sup>31</sup>-Ile<sup>32</sup>-Gly<sup>33</sup>-  
 Leu<sup>34</sup>-Met<sup>35</sup>-Val<sup>36</sup>-Gly<sup>37</sup>-Gly<sup>38</sup>-Val<sup>39</sup>-Val<sup>40</sup>-Ile<sup>41</sup>-Ala<sup>42</sup>-COOH

# ALZHEIMER'S DISEASE



## Amyloid Hypothesis

### Changes in A $\beta$ metabolism

- Increase in total A $\beta$  production
- Increase in the A $\beta$ 42/A $\beta$ 40 ratio
- Reduced A $\beta$  degradation/clearance

Oligomerization of A $\beta$ 42 and initial (diffuse) A $\beta$ 42 deposits

Subtle effects of soluble A $\beta$ 42 oligomers on synaptic function

Inflammatory responses (microglial and astrocytic activation) and amyloid plaque formation

Progressive synaptic/neuronal injury

Altered neuronal ionic homeostasis & oxidative injury

Aberrant oligomerization and hyperphosphorylation of tau

Widespread neuronal dysfunction and cell death associated with neurotransmitter deficits

**Dementia with plaque and tangle pathology**

# Problems with the amyloid hypothesis

- In some cases, individuals without symptoms of AD have many cortical A $\beta$  deposits. However, in these cases, these are diffuse amyloid plaques that are not associated with surrounding necrotic and glial pathology.
- The degree of dementia appears to correlate with soluble A $\beta$  species. Several lines of evidence demonstrate that soluble A $\beta$  oligomers, instead of monomers or insoluble amyloid fibrils, may be responsible for synaptic dysfunction in the brains of AD patients and in animal models.

