

# Molecular Dynamics Simulation of Amyloid Beta in Alzheimer's Disease

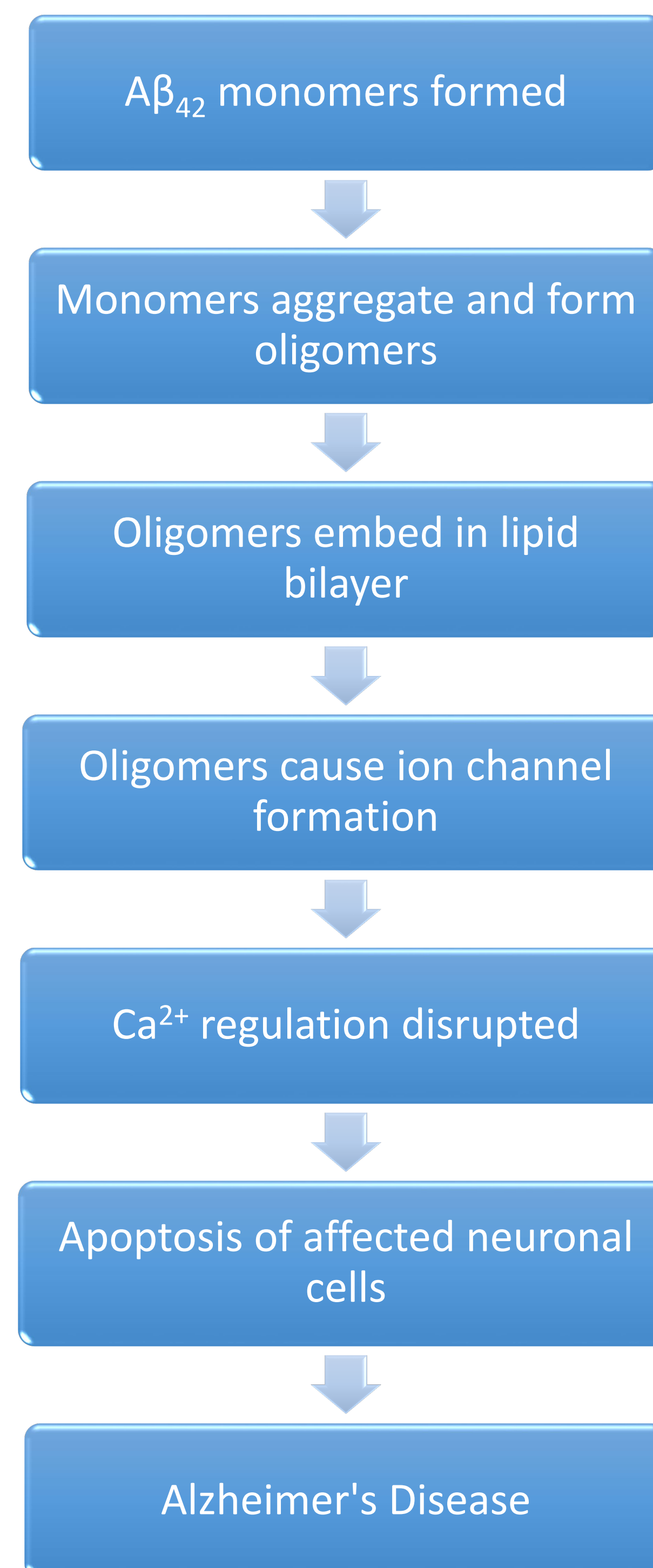
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## Background

- Amyloid beta ( $A\beta$ ) peptides are formed by improper cleavage<sup>1</sup>
  - Most toxic form is 42 amino acids
  - Plaques are a hallmark of Alzheimer's Disease, but oligomers cause disease and progression

## Ion Channel Hypothesis of Alzheimer's Disease<sup>2</sup>

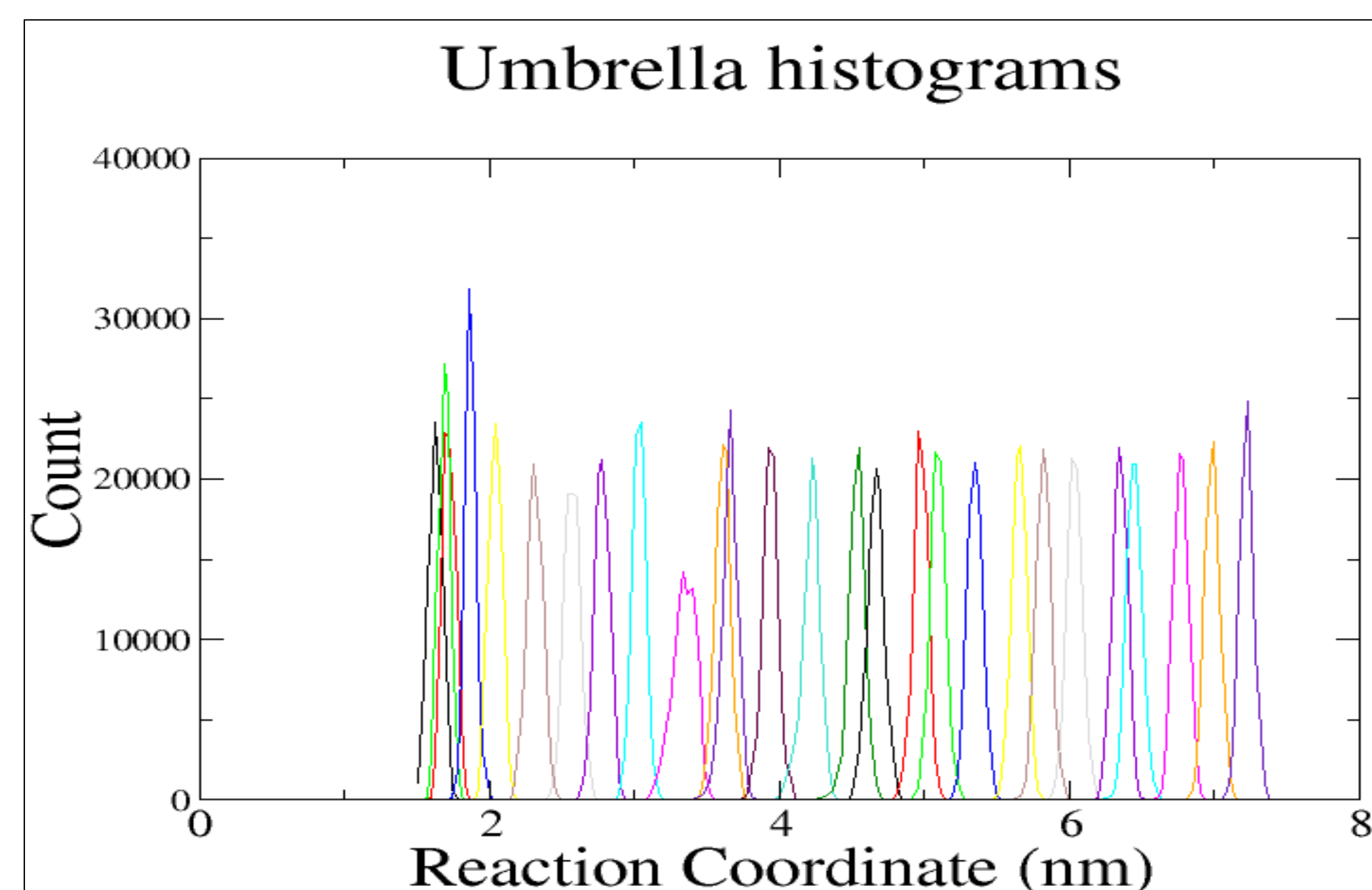


## Introduction

- Modeling the neuronal membrane and  $A\beta_{42}$  oligomers
- Simulating pore formation in the membrane
- Replicate results found by Xiang, et al.<sup>3</sup>
- Run additional simulations
- Find binding energies

## Methods

- Utilize CHARMM-GUI membrane builder
  - Model the membrane
  - Implant oligomer
- Run simulation using GROMACS
- Visualize using VMD
- Use umbrella sampling
  - Pulling simulation now frames
  - Individual simulations for frames



## Simulation

- Pulls  $A\beta_{42}$  oligomer away from the center of the mass of the membrane
- Force generated using a spring with a constant of  $1000 \frac{kJ}{mol \cdot nm^2}$
- Spring expanded at a constant rate  $0.01 \frac{nm}{ps}$

## Simulation Components

- $A\beta_{42}$ 
  - PDB: 2MXU<sup>4</sup>
  - Homo-oligomer
  - Colored regions are amino acids
- Membrane
  - POPC<sup>3</sup>
  - Not showing lipid tails
  - Red is O, blue is N, gold is P
- Solution
  - 0.15M KCl<sup>3</sup>
  - Water shown as blue points
  - KCl ions not shown

## Results

Figure 1

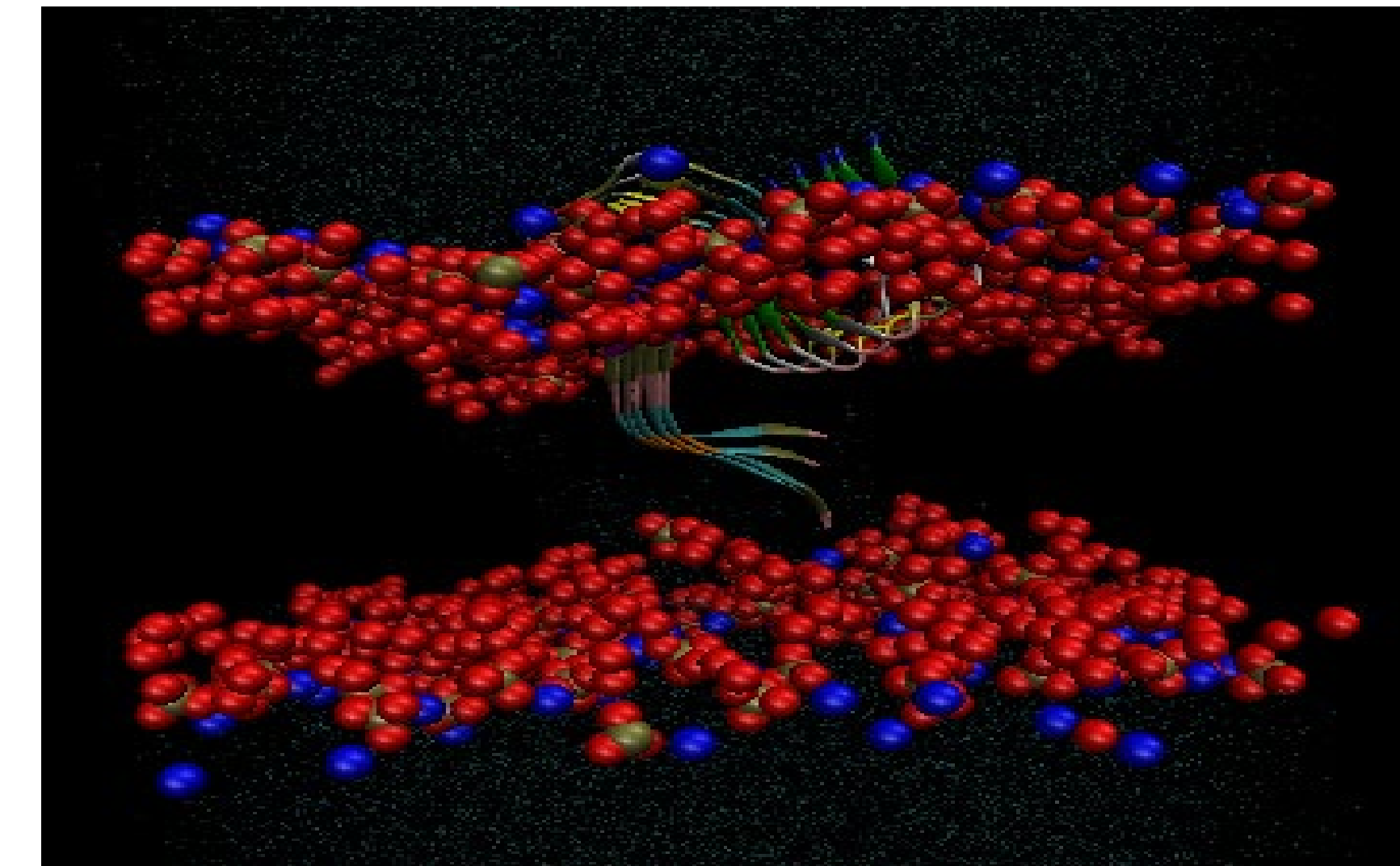


Figure 2

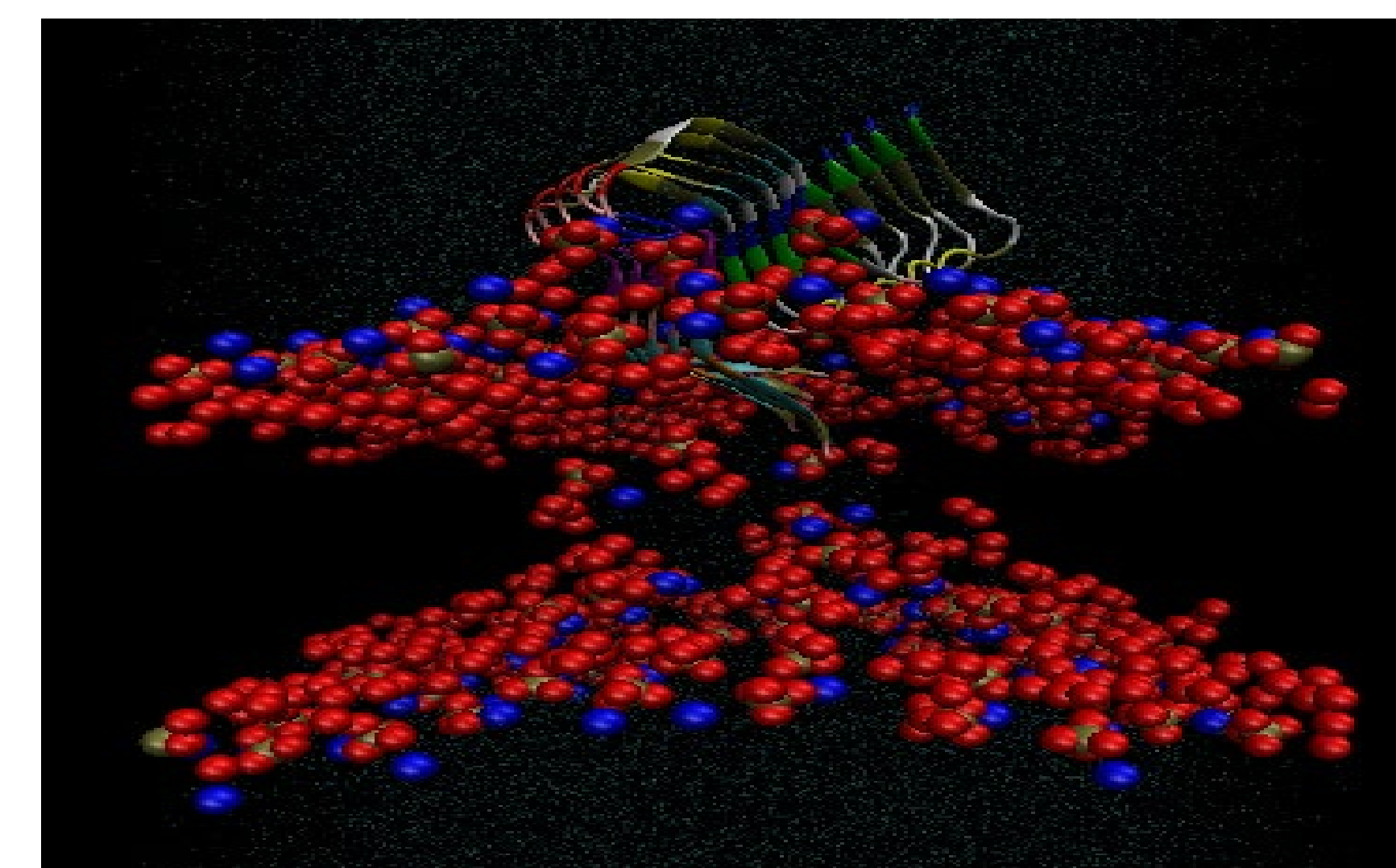


Figure 3

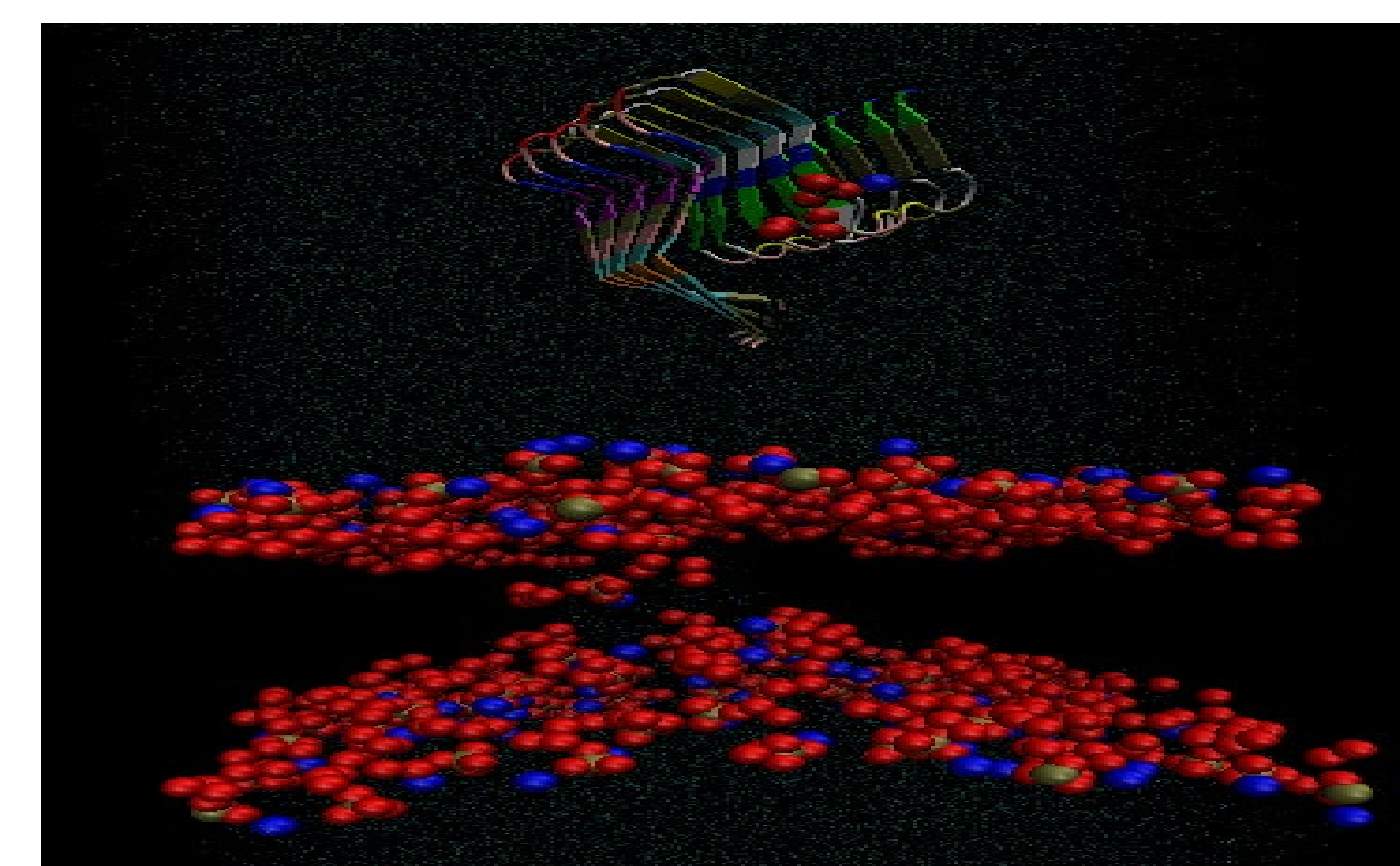
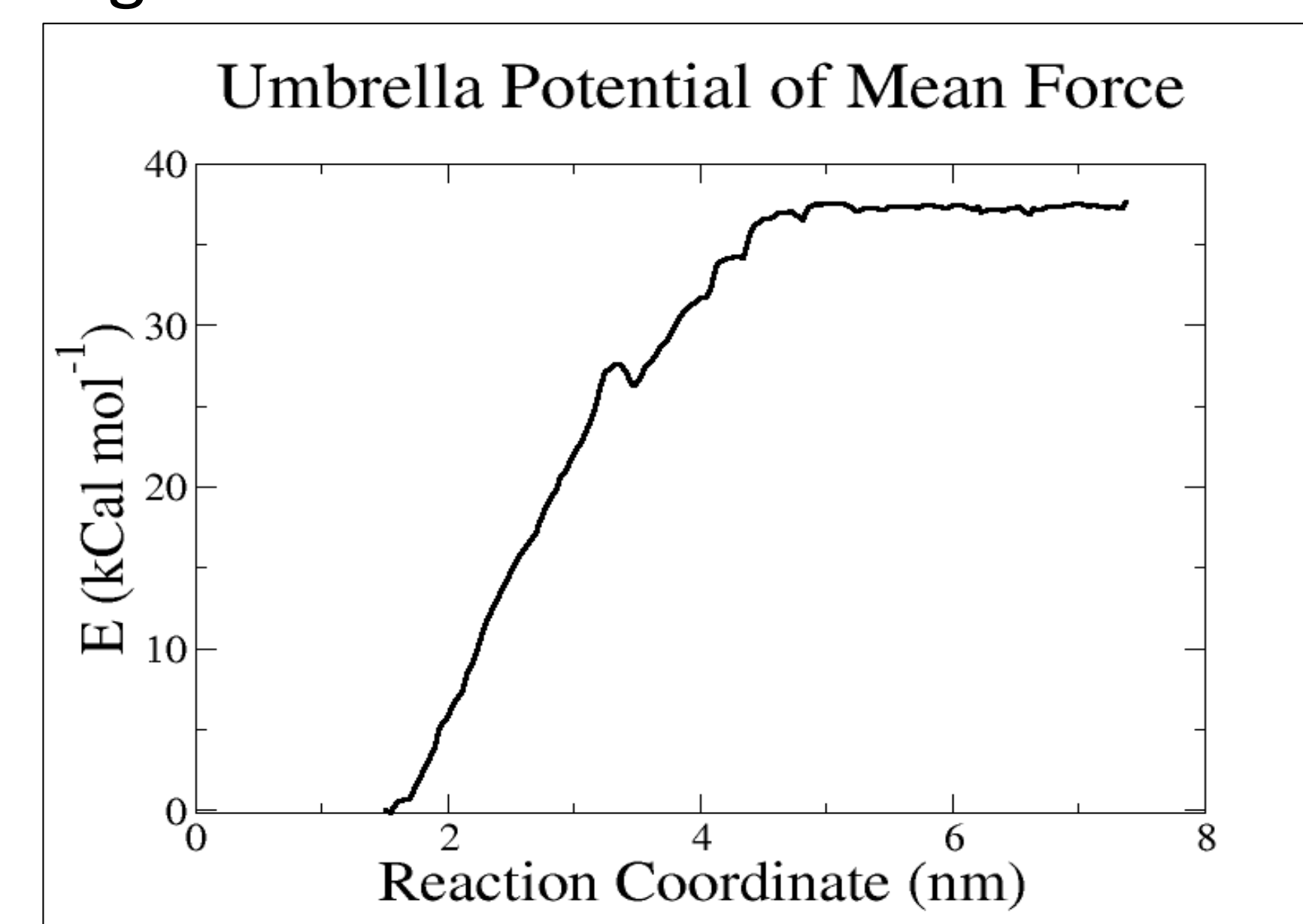


Figure 4



## Conclusions

- 2MXU caused deformation in all configurations
  - Largest deformation with amine terminus inside membrane, shown progressing from fig. 1 – fig. 3
- Binding energy of 2MXU to membrane,  $\Delta G_{bind}$  in fig. 4
  - $\Delta G_{bind} = 37.3 \pm 0.2 \frac{kCal}{mol}$

## Future Work

- Literature search on binding energies of  $A\beta_{42}$
- Apply umbrella sampling to all  $A\beta_{42}$  configurations that have completed the pulling simulation

## References

1. Pearson, H. A., & Peers, C. (2006). Physiological roles for amyloid beta peptides. *The Journal of physiology*, 575(Pt 1), 5–10. doi:10.1113/jphysiol.2006.111203
2. Shirwany, N. A., Payette, D., Xie, J., & Guo, Q. (2007). The amyloid beta ion channel hypothesis of Alzheimer's disease. *Neuropsychiatric disease and treatment*, 3(5), 597–612.
3. Xiang, N., Lyu, Y., Zhu, X., & Narsimhan, G. (2018). *PCCP*, 20(10), 6817-6829.
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