

# Physical-chemistry methods for study of protein-protein interactions involved in neurodegenerative diseases

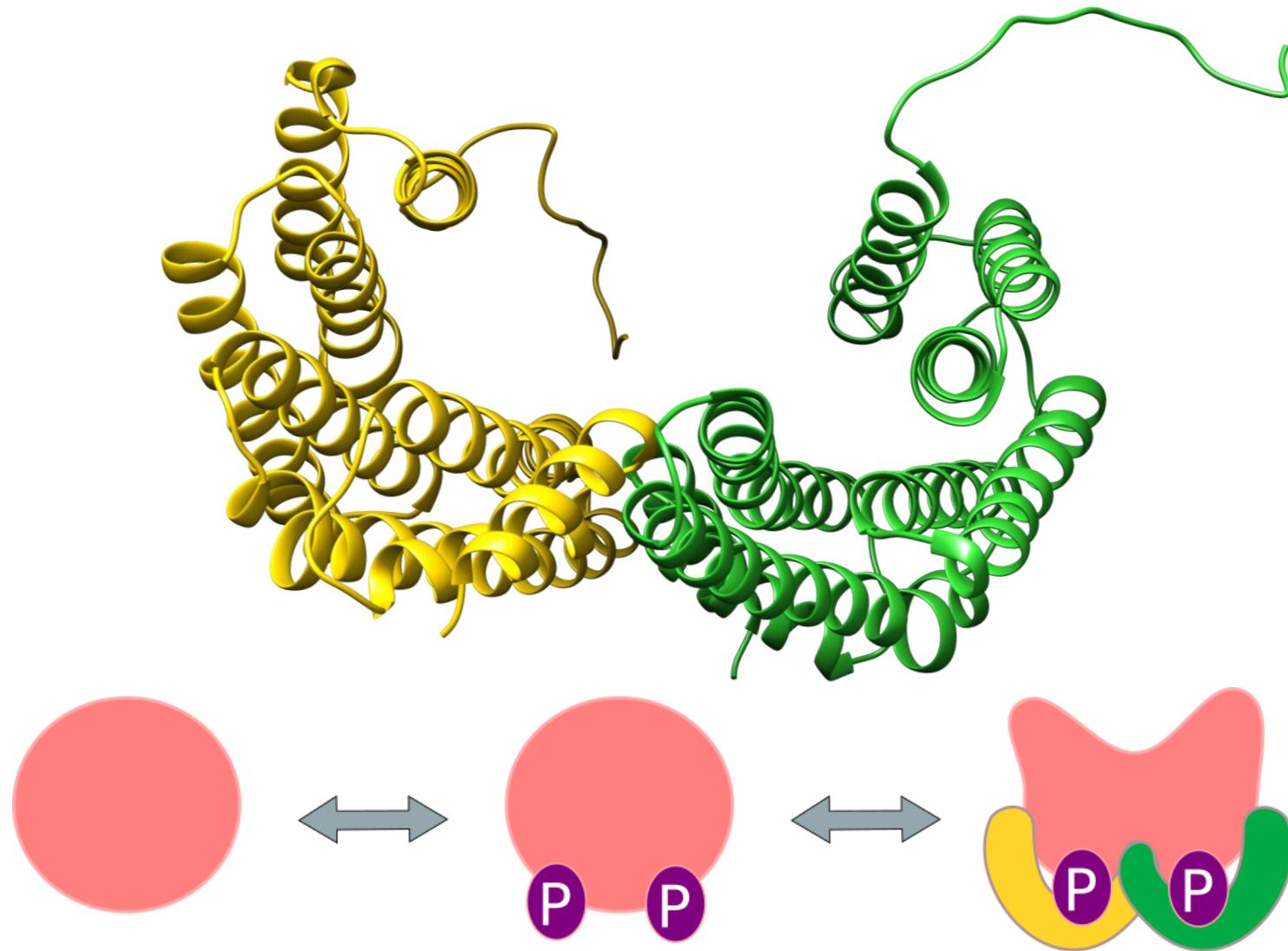
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Biophysical Chemistry

# RESEARCH OBJECT: 14-3-3/complexes



# NMR: REGULATORY DOMAIN

Biophysical Chemistry 223 (2017) 25–29



Contents lists available at ScienceDirect

Biophysical Chemistry

journal homepage: <http://www.elsevier.com/locate/biophyschem>



## Phosphorylation of the regulatory domain of human tyrosine hydroxylase 1 monitored using non-uniformly sampled NMR



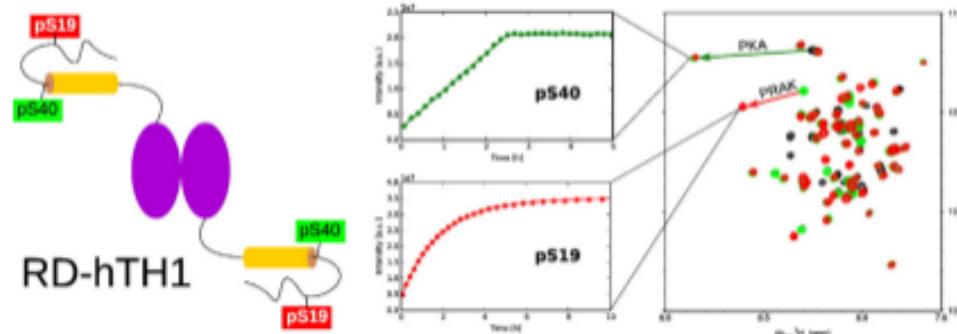
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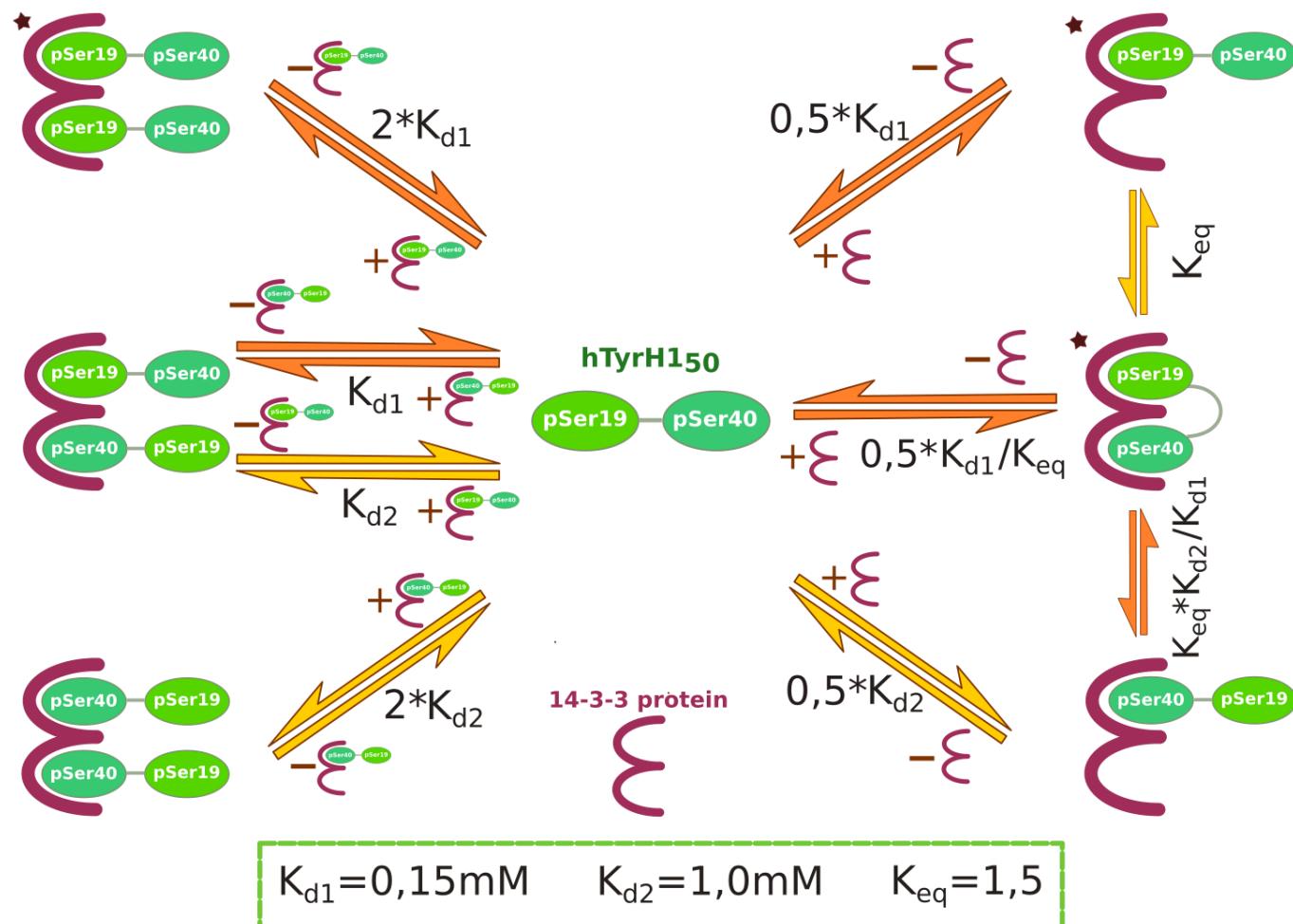
### HIGHLIGHTS

- Disordered part of regulatory domain of human tyrosine hydroxylase 1 was assigned.
- Transient alpha-helices are present next to phosphorylation sites S40 and S19.
- The secondary structure does not change after phosphorylation.
- The phosphorylation kinetic rates were measured efficiently using time resolved NMR.

### GRAPHICAL ABSTRACT

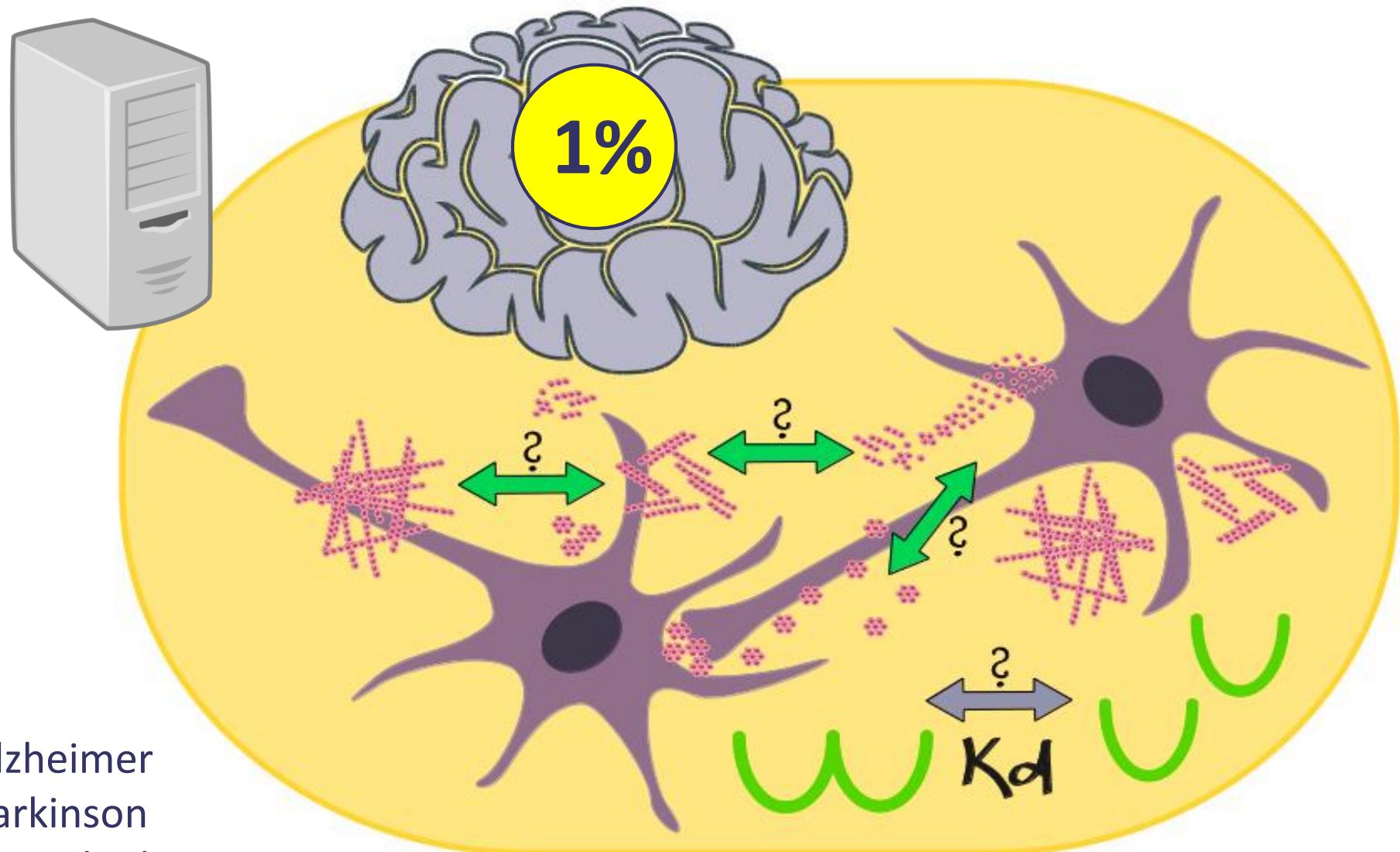


## 14-3-3: BINDING MODES

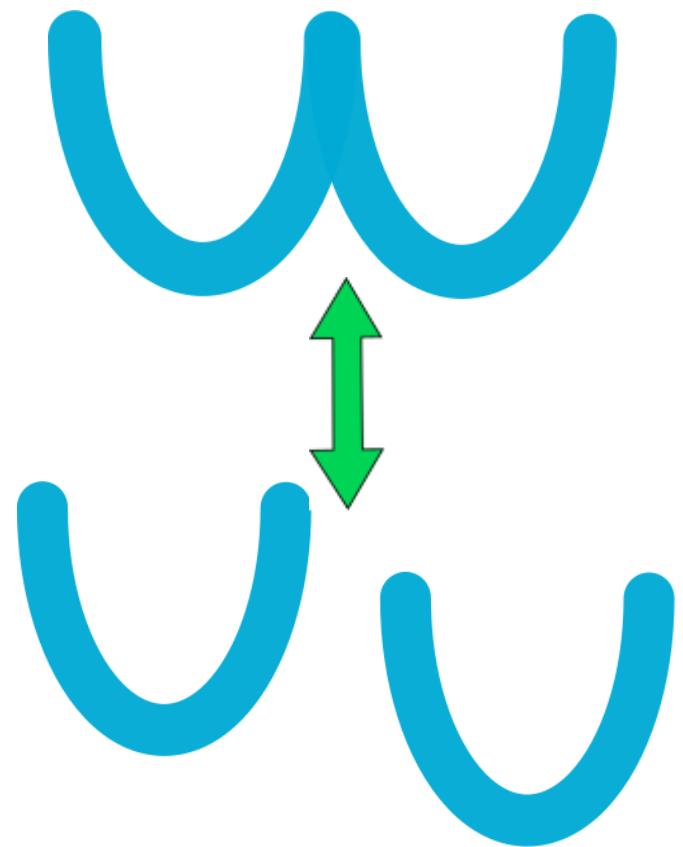
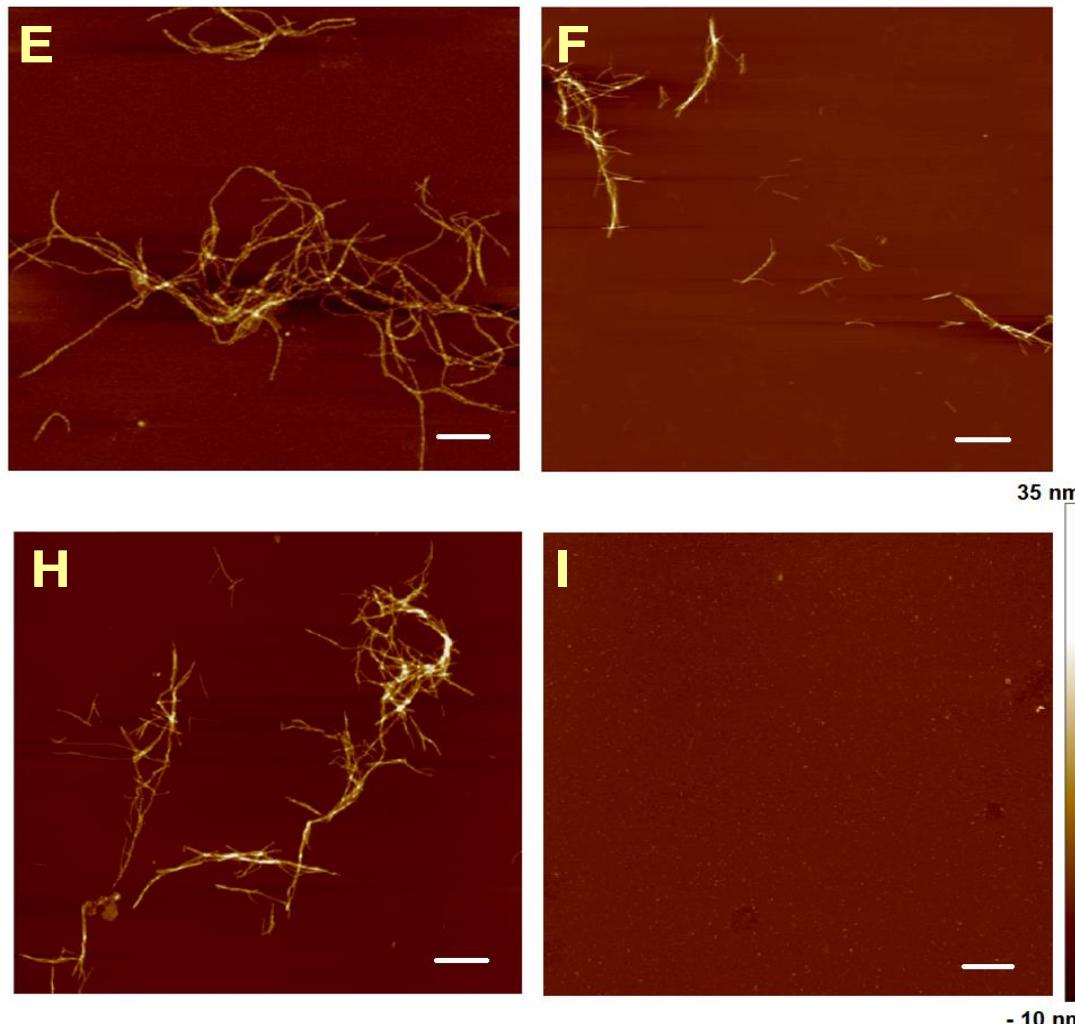


Hritz J.; Byeon I-J.; Krzysiak T.; Martinez A.; Sklenář V.; Gronenborn A.M. Dissection of binding between a phosphorylated tyrosine hydroxylase peptide and 14-3-3 $ζ$ : a complex story elucidated by NMR. *Biophys. J.* 2014, 107, 2185-2194

## 14-3-3: NEURODEGENERATION

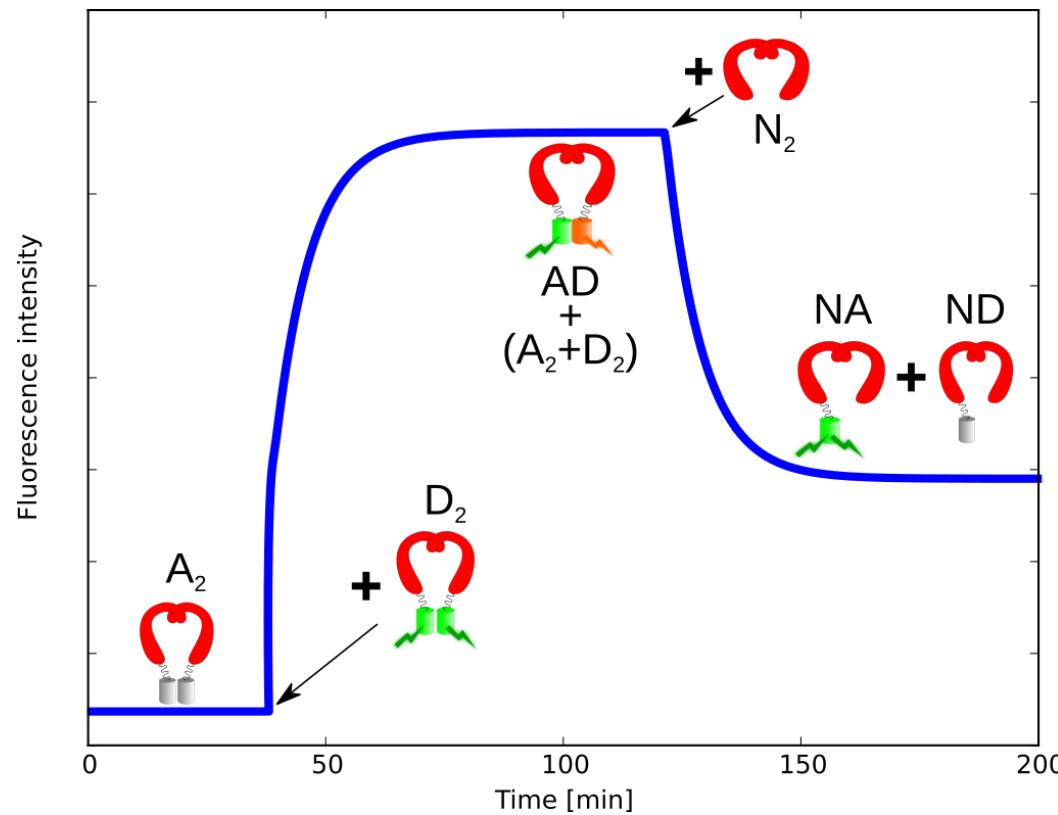
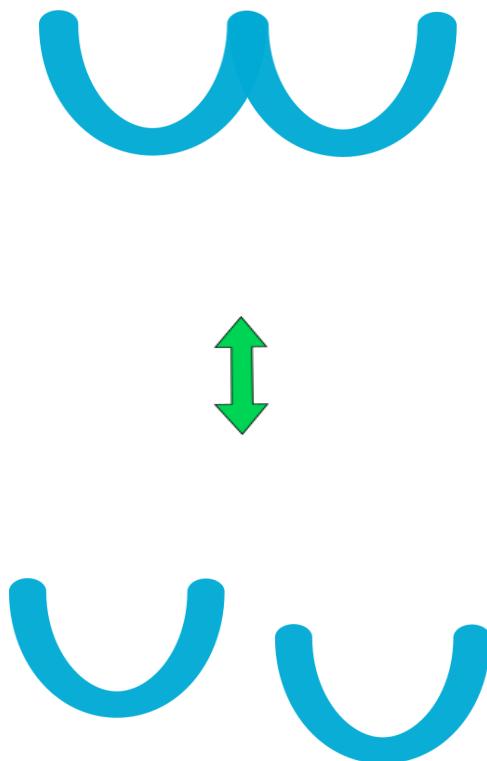


# ALZHEIMER DISEASE: 14-3-3+A $\beta$



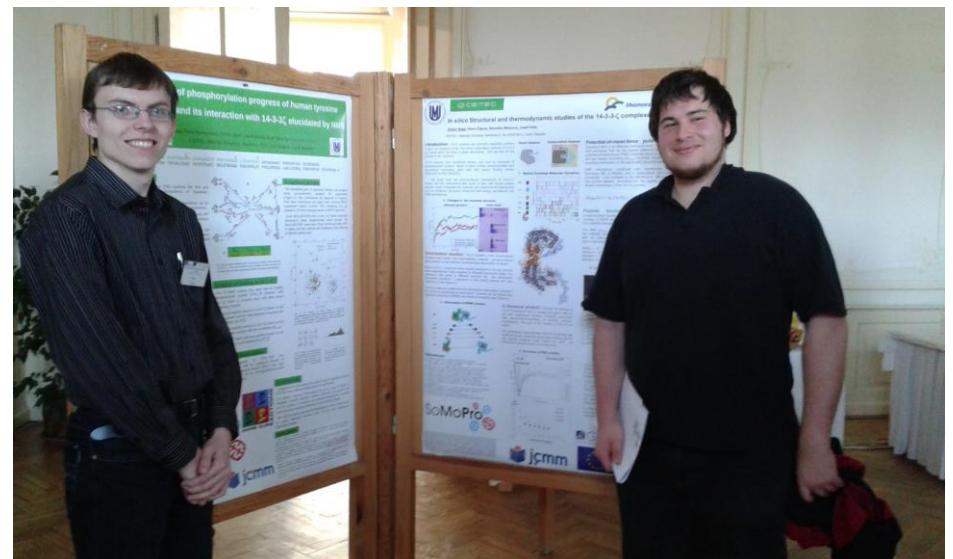
**BIOPHYSICAL  
CHEMISTRY**

## 14-3-3 KINETIC ASSAY: FRET



- Experimental Biophysical Chemistry – fluorescence assays
- Computational Biophysical Chemistry – structural and free energy calculations of dimer/monomer equilibria

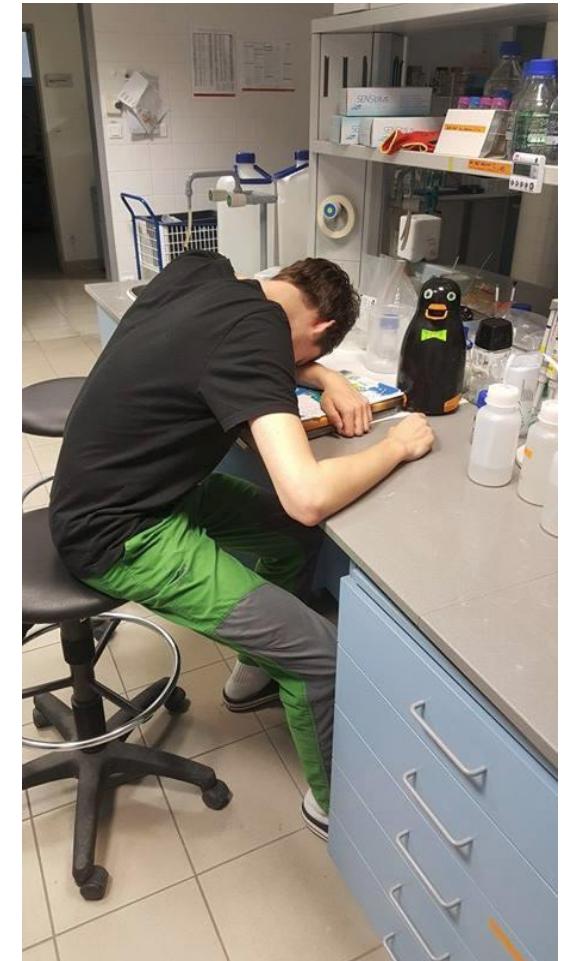
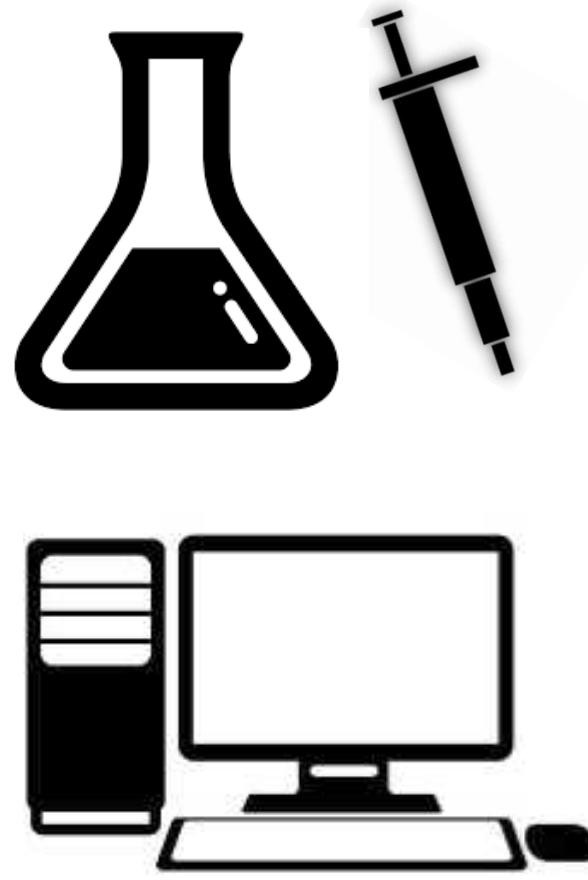
# HANDS ON EXPERIENCE CONFERENCES AND INTERNSHIPS



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# SUITABLE CANDIDATES OPEN POSITIONS



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