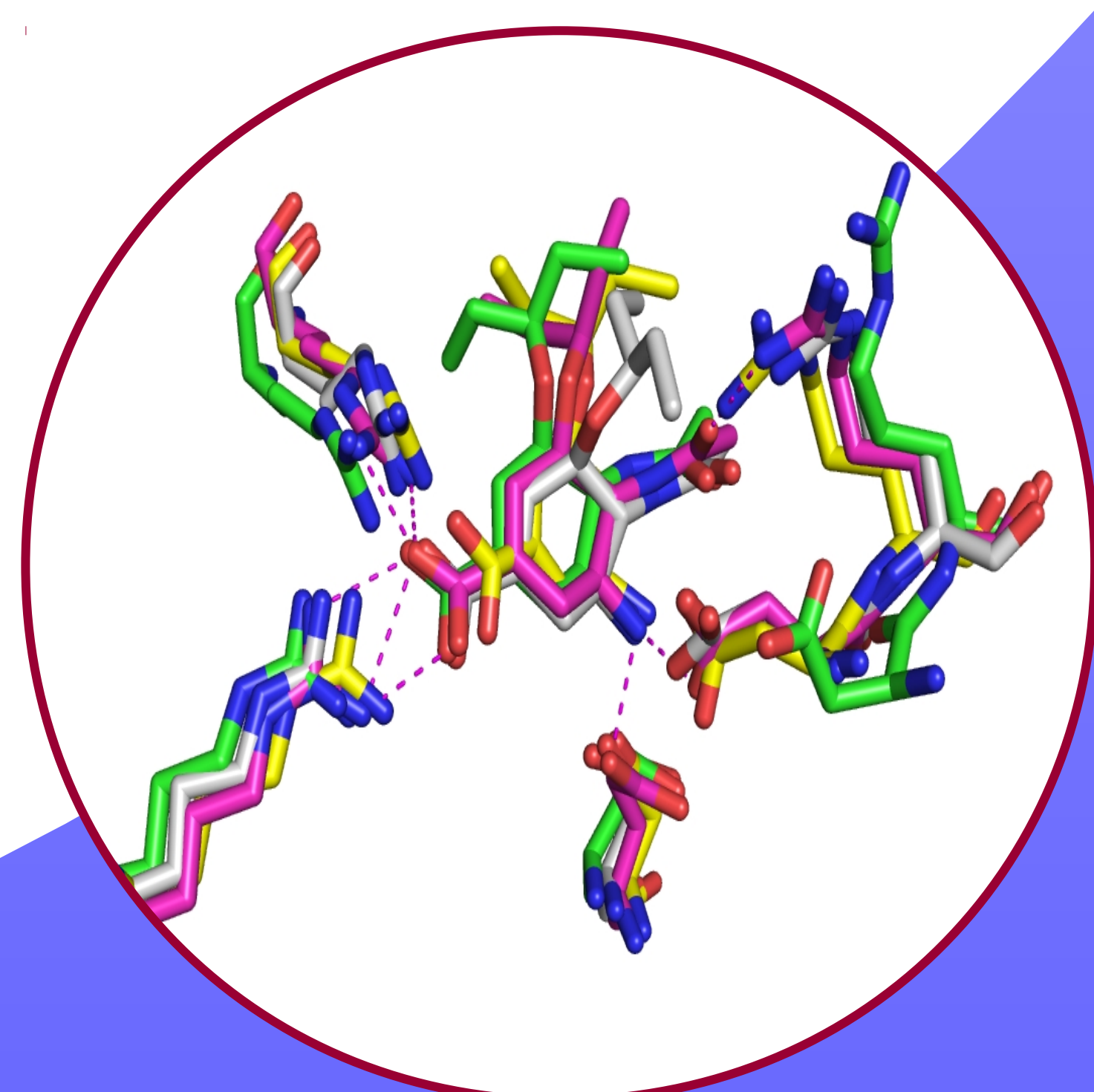
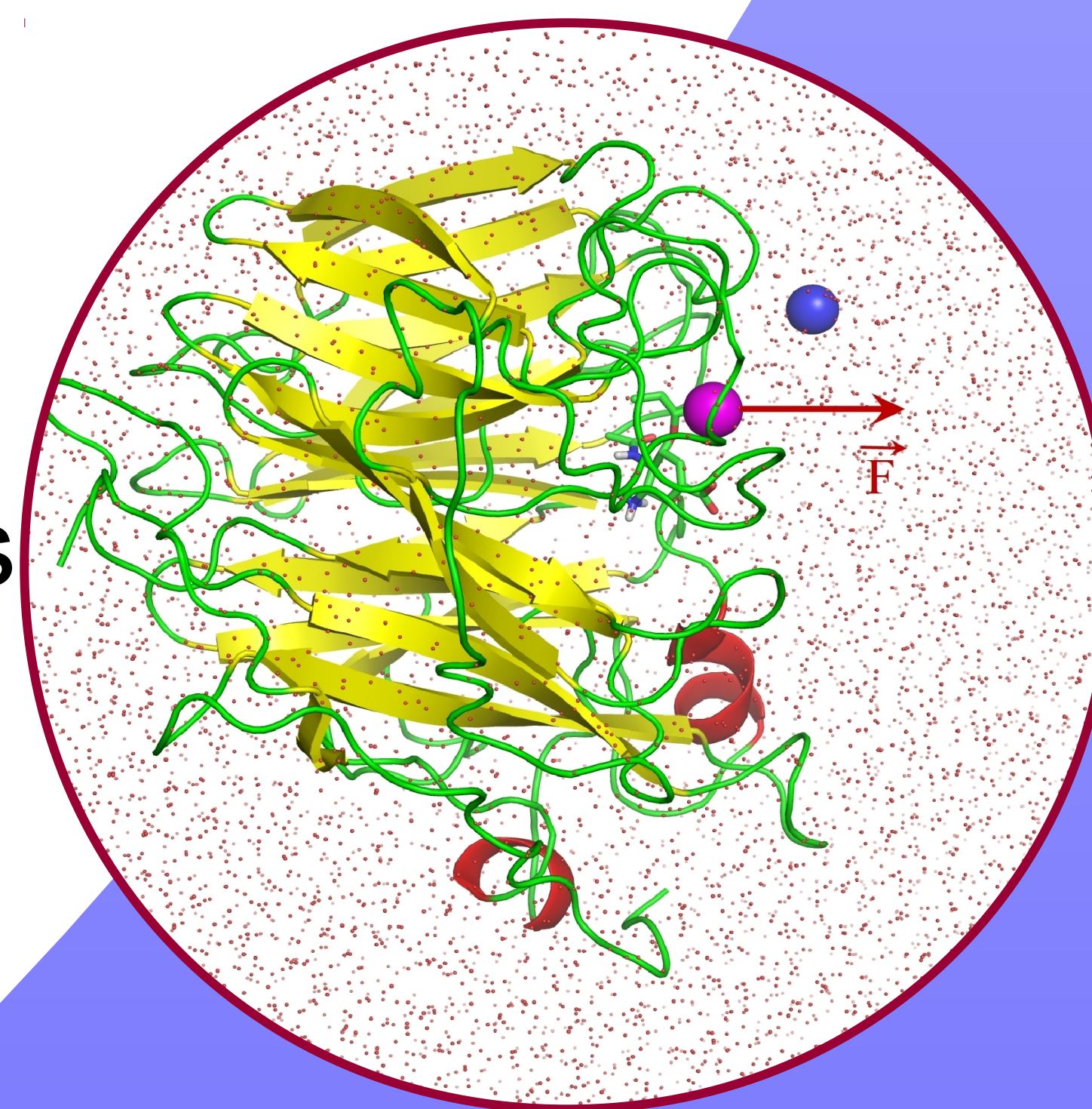
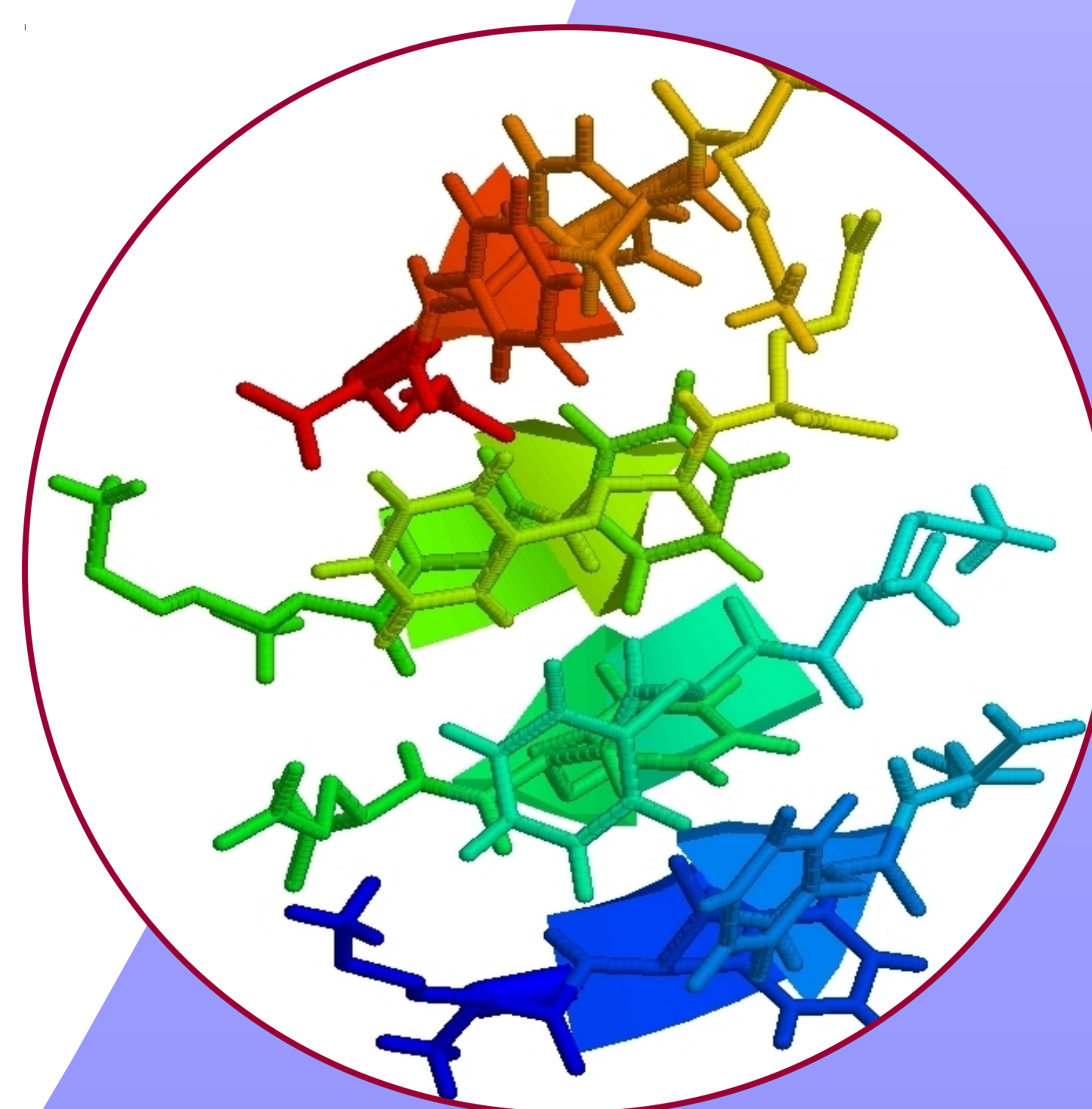
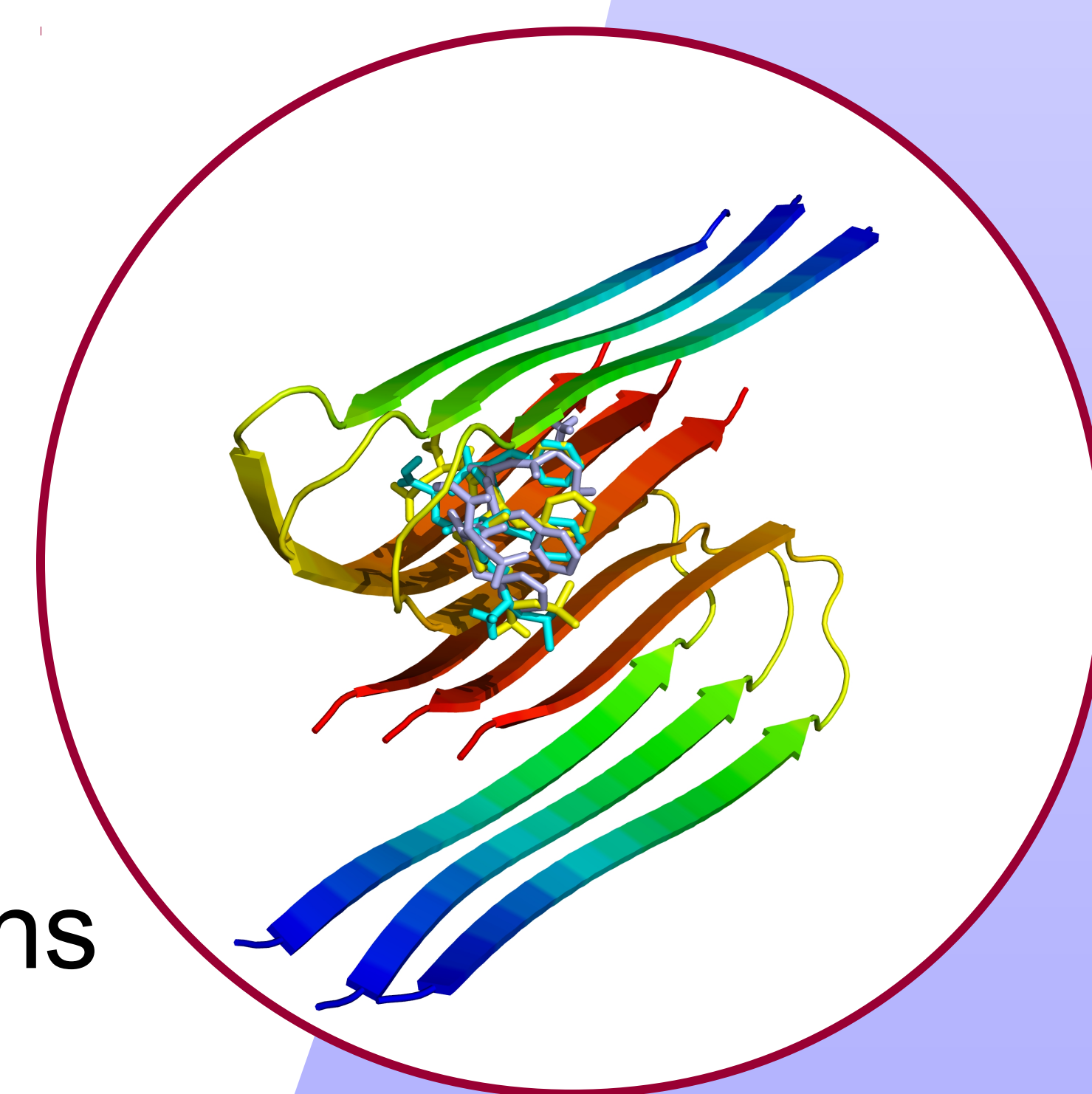




PROTEIN DAY

December 16, 2010

- 8:25-8:30** Welcome
- 8:30-9:10** **Trinh Xuan Hoang**, Center for Computational Physics,
VAST, Hanoi
Protein folding: unified perspective from the tube model
- 9:10-9:50** **Jacek Lulek**, University of Zielona Góra, Poland
Constructing of molecular phylogenetic trees by
different approaches
- 9:50-10:30** **Le Thi Ly**, ICST, Ho Chi Minh City
Computer-aided drug design for Diabetes type II
- 10:30-11:00** Break
- 11:00-11:30** **Mai Suan Li**, Polish Acad Sci and ICST, HCM City
Alzheimer's disease: Insights from computer simulations
- 11:30-11:45** **Nguyen Truong Co**, ICST, HCM City
Factors governing fibrillogenesis of polypeptide chains
revealed by lattice models
- 11:45-12:00** **Ngo Son Tung**, ICST, HCM City
Inhibition of aggregation by beta-sheet
breaker peptides
- 12:00-12:15** **Mai Khanh Binh**, ICST, HCM City
Study of influenza virus by
steered molecular dynamics
- 12:15-12:30** **Nguyen Truc Trang**,
ICST, HCM City
Estimation of binding free
energies by different force fields
- 12:30-14:00** Lunch and Discussion



Place Room 204. Institute for Computational
Science and Technology
Quarter 6. Linh Trung Ward,
Thu Duc District. HCMC

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