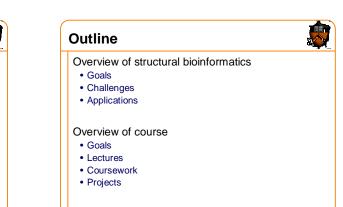


Thomas Funkhouser Princeton University Fall 2005



# **Bioinformatics**

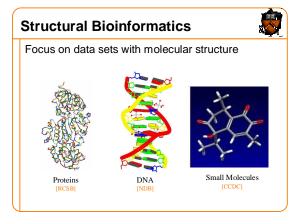


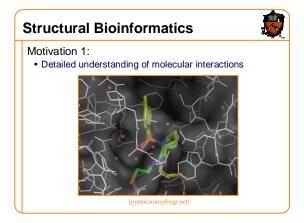
Definition:

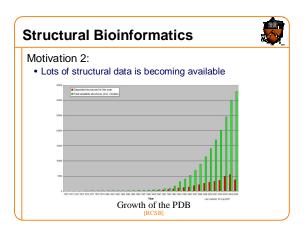
"The collection, archiving, organization, and interpretation of biological data." [Orengo, 2003]

Generally speaking:

- Large biological data sets
- Computer representations and algorithms
- Storage, analysis, visualization, prediction, and design





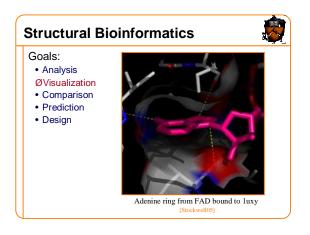


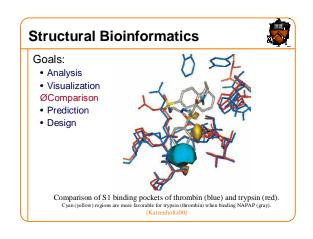
# Structural Bioinformatics

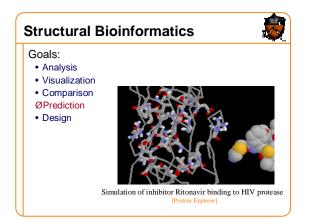
Goals:

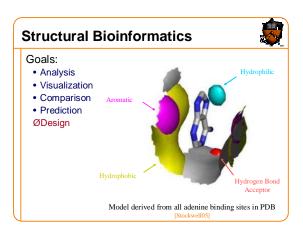
- Analysis
- Visualization
- ComparisonPrediction
- Design

Structural Bioinformatics Coals: ØAnalysis 0 Visualization 0 Prediction 0 Design 0 Design 0 Comparison 0 Prediction 0 Design 0 Comparison 0 Design 0 D







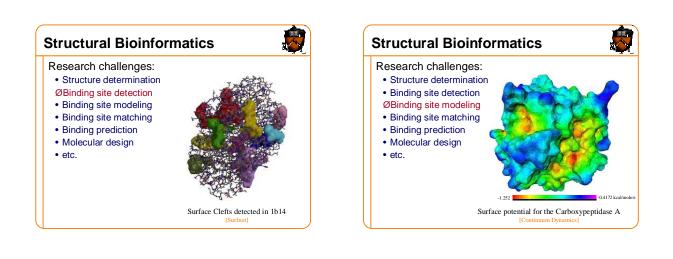


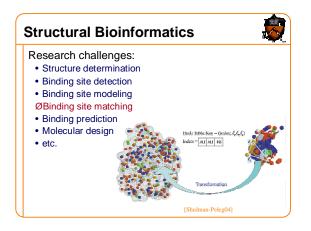
# **Structural Bioinformatics**

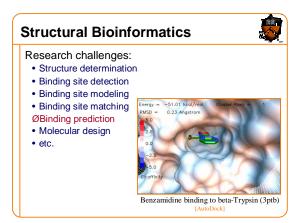
Research challenges:

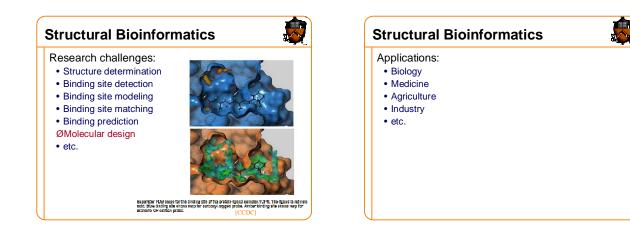
- Structure determination
- Binding site detection
- Binding site modeling
- Binding site matching
- Binding prediction
- Molecular design
- etc.

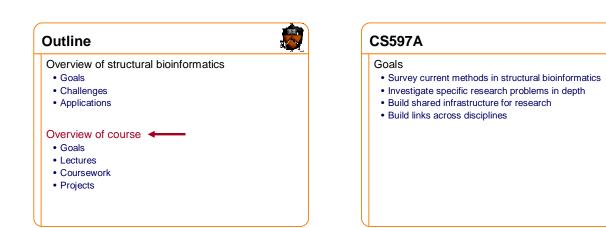
Structural Bioinformatics Research challenges: ØStructure determination Binding site detection Binding site modeling Binding prediction Binding prediction Molecular design • etc.

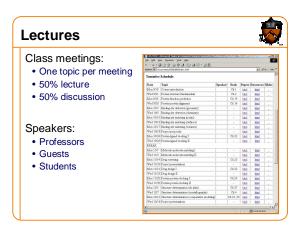














• Lead discussion afterwards

# Readings

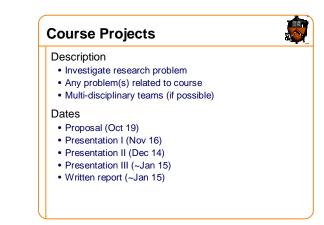
### Book

• P.E. Bourne, H. Weissig, "Structural Bioinformatics," Wiley-Liss, 2003.

## Papers

- Read ~1 paper for each lecture
- Be prepared for class discussion

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# Course Projects Wrap Up Example projects • Develop new method • Develop new method • Hand in survey questionaire • Evaluate/compare existing methods • Gombine methods in system • Apply method • Sign up for in-class presentations • Key elements • Investigation • Investigation • Evaluation

# 5