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here are the details for the invited ECMI Minisymposium on Molecular Modeling that I was asked to organize.

Arnold Neumaier, University of Vienna,
Global Optimization and Protein Structure Prediction

Burkhard Rost, European Molecular Biology Laboratory Heidelberg,
Learning from Evolution to Predict Protein Structure

Christodoulos A. Floudas, Princeton University,
Structure Prediction in Computational Chemistry, Biology, and Immunology: Protein Folding and Peptide Docking

(in this order, each talk 40 minutes including discussion)

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