Near-term Potential Practical Wins for Modeling and Simulation of Microbes

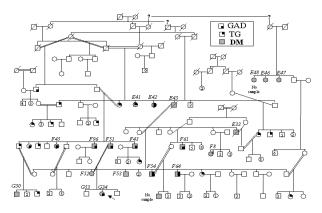
Bioinformatics

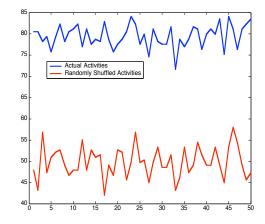
- Predicting Domain-Ligand Interaction using Signature Kernel Support Vector Machines
- Natural Language Processing
- Gene Finding, Phylogeny

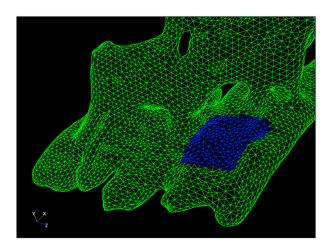
Hardware & Operating Systems Research

 What does the architecture of the computer look like that can solve these problems?











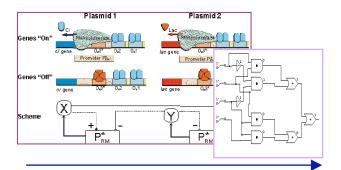
Near-term Potential Practical Wins for Modeling and Simulation of Microbes

Computational Molecular Biophysics

• +40ns Simulation of Rhodopsin Membrane Protein System for Insight into the determination of the light-adapted structure

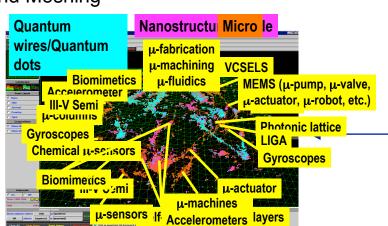
Complex Systems

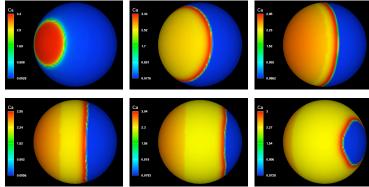
Network Modeling



Complex Systems

 Massively Parallel Finite Elements and Meshing





Computational Technologies

Parallel Algorithm Development, Optimization, Data
 Mining and Management and Visualization, Frameworks & User Interfaces