

HPC, Computational Science & Engineering, *Shake-and-Bake*, and 21st Century Academia

Russ Miller

Cyberinfrastructure Lab

The State University of New York at Buffalo



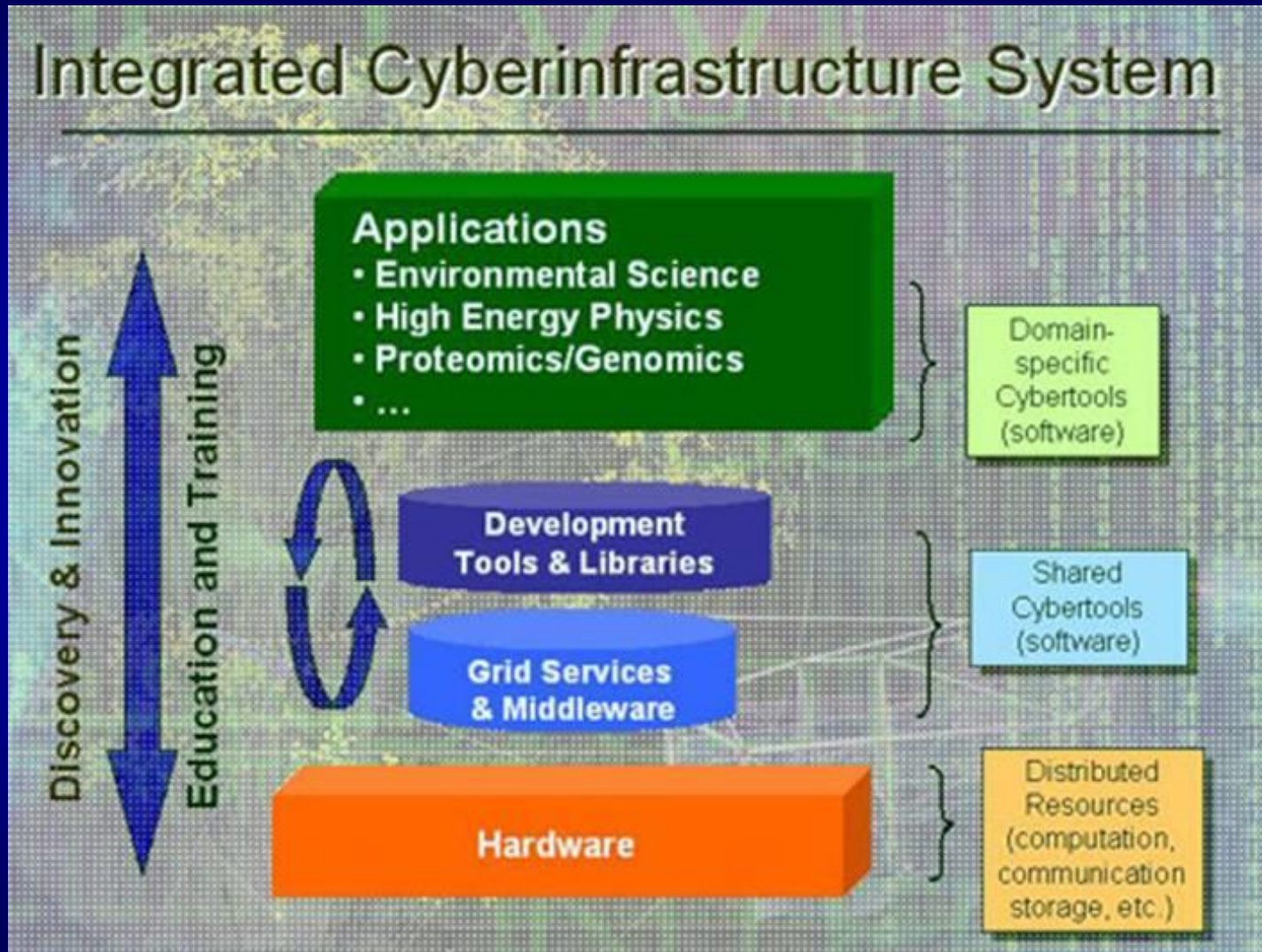
Academia in the 21st Century: High-Level View

- Empower students to compete in knowledge-based economy
- Embrace digital data-driven society
- Accelerate discovery and comprehension
- Enhance virtual organizations
- Provide increased education, outreach, and training
- Enhance and expand relationships between academia and the corporate world

Academia in the 21st Century: Medium-Level View

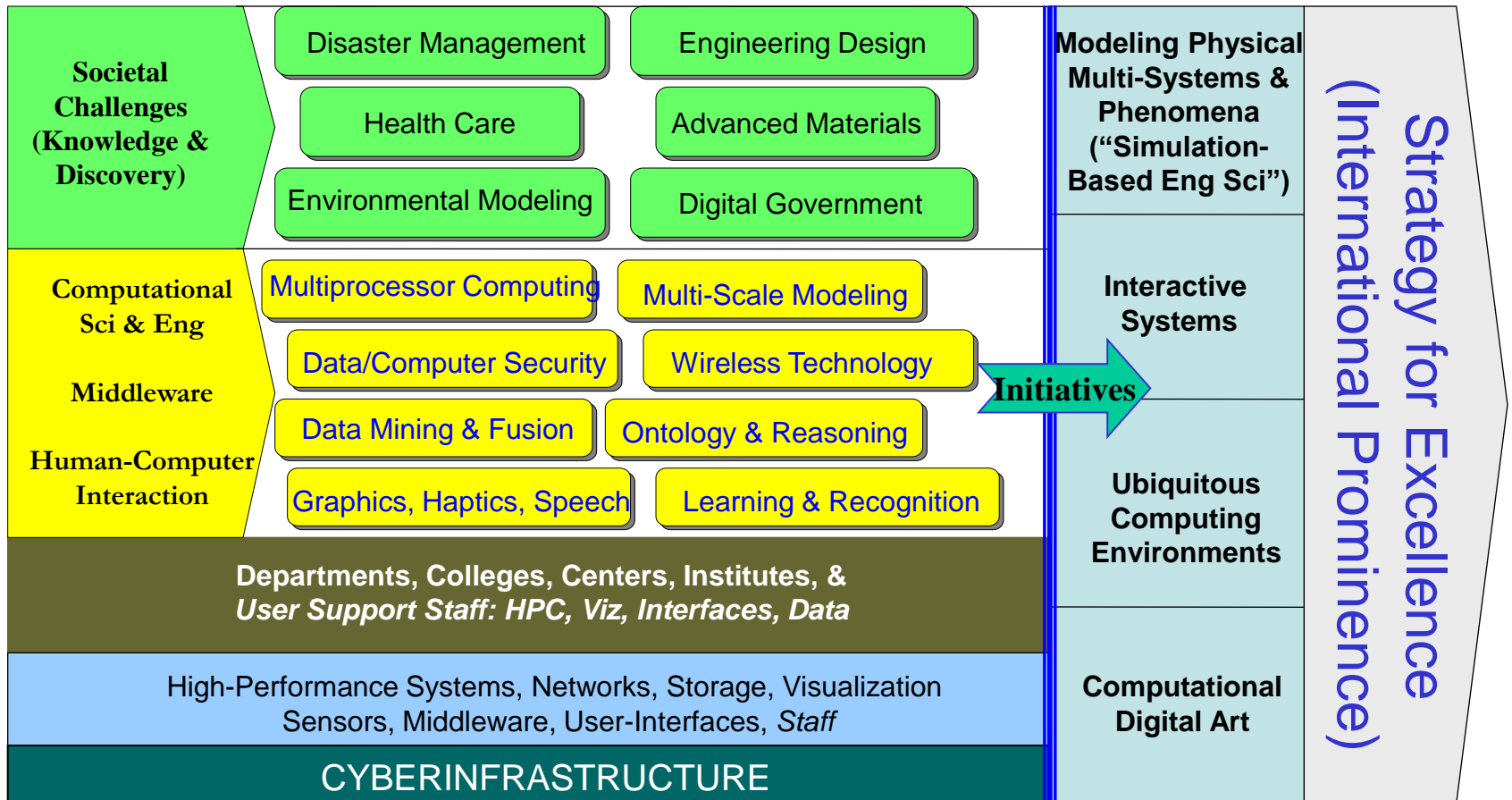
- Create links between enabling technologists and disciplinary users
- Improve efficiency of knowledge-driven applications in myriad disciplines
 - New Techniques
 - New Algorithms
 - New Interactions (people & systems)
- Support HPC infrastructure, research, and applications
- Deliver high-end cyberinfrastructure to enable efficient
 - Collection of data
 - Management/Organization of data
 - Distribution of data
 - Analysis of data
 - Visualization of data

NSF Integrated Cyberinfrastructure



NSF Director Arden L. Bement: "leadership in cyberinfrastructure may determine America's continued ability to innovate – and thus our ability to compete successfully in the global arena."

Academic Computing Initiative: Inverted Umbrella (Sample)



Academic Computing Initiative: Organization

- **Must be Pervasive Across the Entire University**
- **Must Remove Barriers**
- **Groups Must Interact**
 - **Research Groups**
 - **Support Staff**
 - **Students**
 - **Departments**
 - **Colleges**
- **Issues**
 - **Tenure & Promotion**
 - **University vs Colleges vs Departments vs Faculty vs Centers/Institutes vs Degrees vs Courses**
- **Details are University Dependent**

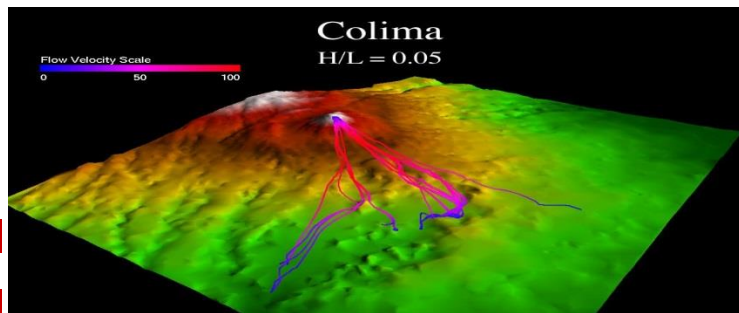
Center for Computational Research (CCR)

- Founding Director (1998-2006)

- Facts & Figures

- ❑ Top Academic HPC Center in World
- ❑ Top 25 HPC System
- ❑ Massive High-End Storage
- ❑ Significant Visualization
- ❑ Special-Purpose Systems
- ❑ ~30 FTEs Staff
- ❑ 140 Projects Annually

- EOT

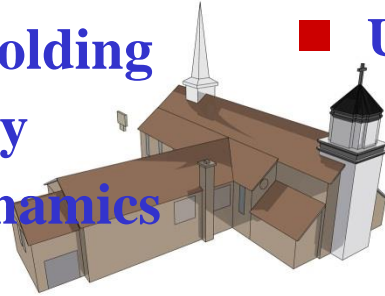
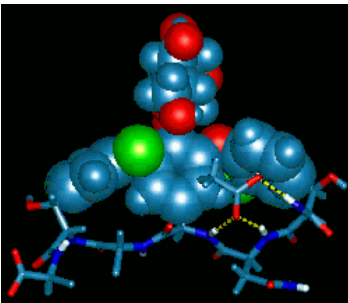


CCR Highlights (1998-2006)

- Provide HE-Comp
- Provide HE-Vis + AGN
- Special Purpose Systems
 - Bioinformatics
 - Data Warehouse / Mining
- Support Local/National Efforts – Industry + Acad
- Create jobs in WNY
- Certificate Program
- Workshops + Tours
 - Campus, Industry
 - High-School
- Urban Planning & Design
- MTV Videos
- Peace Bridge, Med Campus
- Olmsted Parks, Thruway
- NYS Agencies
- Elected Officials
- Magnet on Campus
- Significant Funds
- Numerous Awards
- Significant Publicity

CCR Research & Projects

- Archaeology
- Bioinformatics/Protein Folding
- Computational Chemistry
- Computational Fluid Dynamics
- Data Mining/Database
- Earthquake Engineering
- Environ Modeling & Simulation
- Grid Computing
- Molecular Structure Determination
- Physics



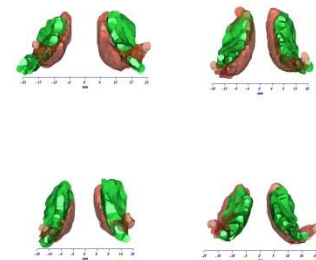
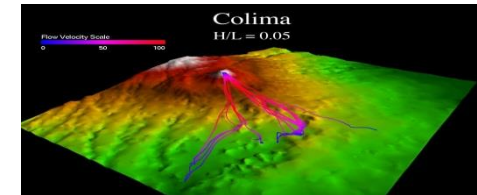
- Videos: MTV
- Urban Simulation and Viz

- StreetScenes
- I-90 Toll Barrier
- Medical Campus
- Peace Bridge



- Accident Reconstruction
- Scientific Viz

- Dental
- Surgery
- MRI/CT Scan
- Confocal Microscopy
- Crystallization Wells
- Collaboratories



Real-Time Visualization

StreetScenes: Real-Time 3D Traffic Simulation

- Accurate local landmarks: Bridges, Street Signs, Business, Homes
- Can be viewed from driver's perspective
- Real-Time Navigation
- Works with
 - Corsim
 - Synchro
- Generate AVI & MOV
- Multiple Simultaneous
 - Traffic Loads
 - Simulation
 - Varying POV



Real-time Simulation



- Key Receptor Sites
- Multiple Viewpoints
- Fully Interactive
- Aerial Photography

Animation & Simulation

Rendered Scenes

Visualization in Planning Studies



Williamsville Toll Barrier Improvement Project



Initial Photo Match incorporating real and computer-generated components

Peace Bridge Visualization: Animation & Simulation

■ Proposed Options

- ❑ Relocate US plaza
- ❑ Build a 3-lane companion span & rehab existing bridge



PHOTO AND STORY BY BRUCE JACKSON

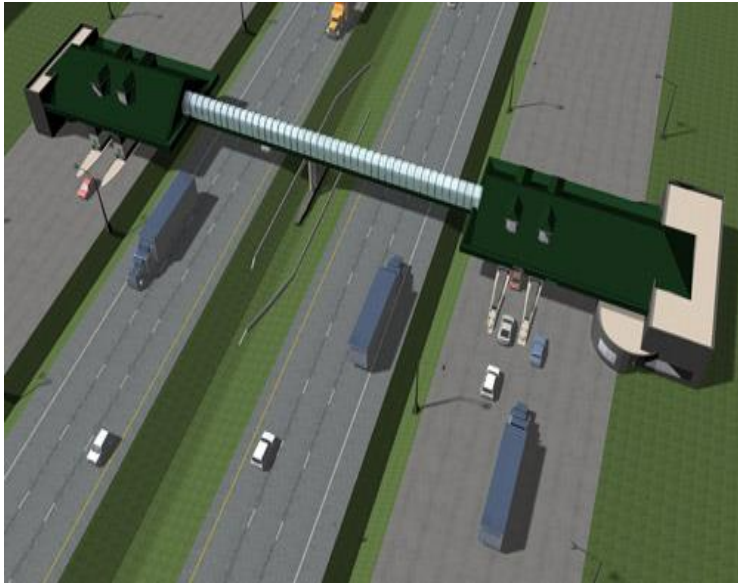


Thruway HOT Lanes Animation



Urban Modeling & Visualization

- Peace Bridge Gateway Improvement Project
- Olmsted Park Conservancy
- Williamsville Toll Barrier Relocation
- Buffalo Niagara Medical Campus



M. Innus, A. Koniak, A. Levesque, T. Furlani

CCR Model Development

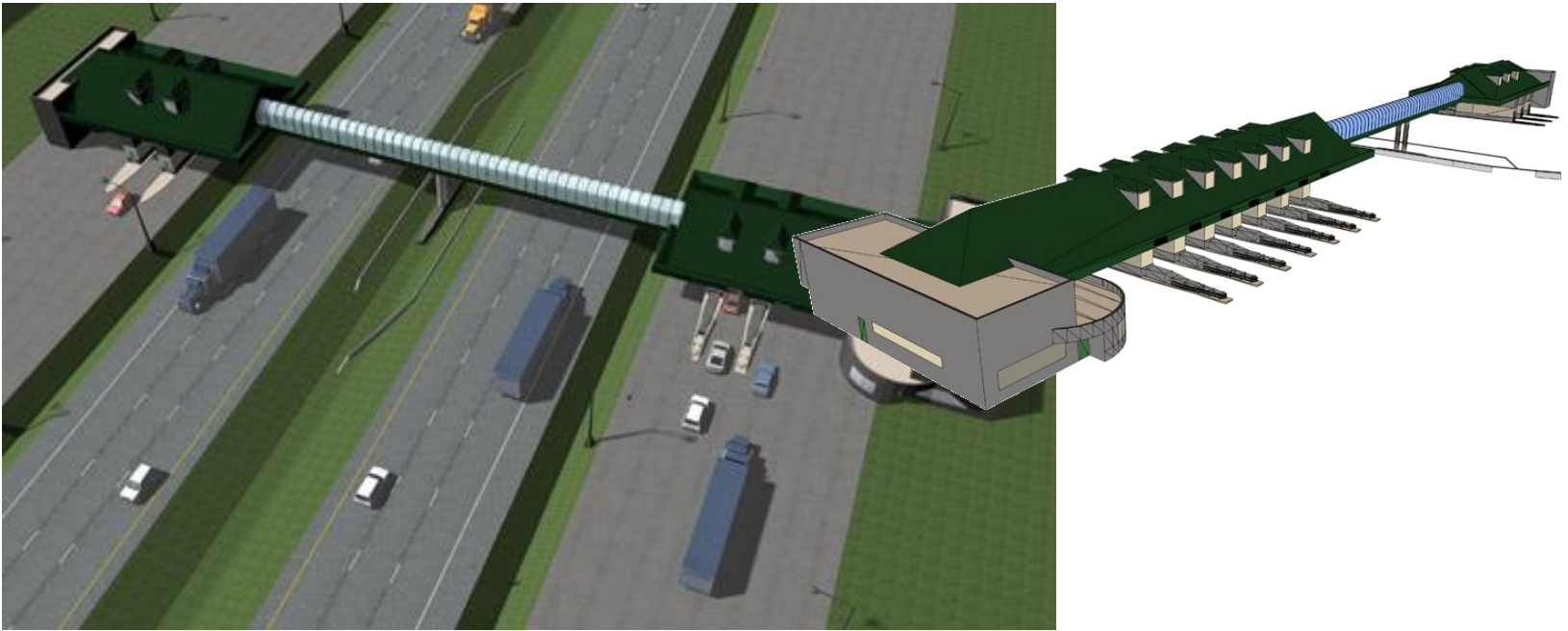
- *StreetScenes*[®] is a Virtual Reality (VR) software solution for 3D interactive visualization of surface traffic.
- Import data from most traffic simulation packages
 - Corsim
 - Synchro
 - Vissim



H. Bucher

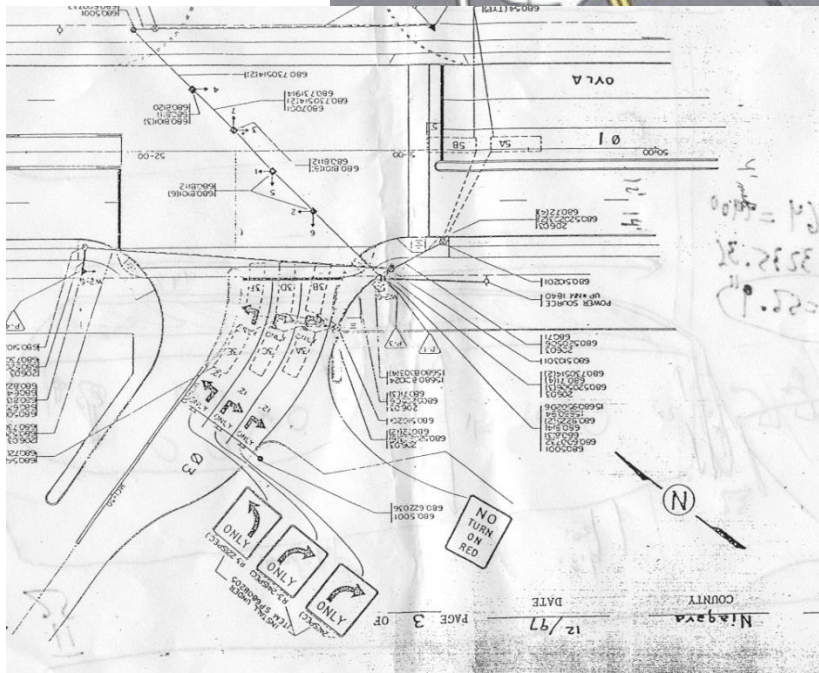
Urban Modeling & Visualization

- High Speed EZPass
- Planning tool for NYS Thruway Authority
- Visualization of real traffic data
- Interactive model for public meetings and demonstrations



M. Innus, A. Koniak, A. Levesque, T. Furlani

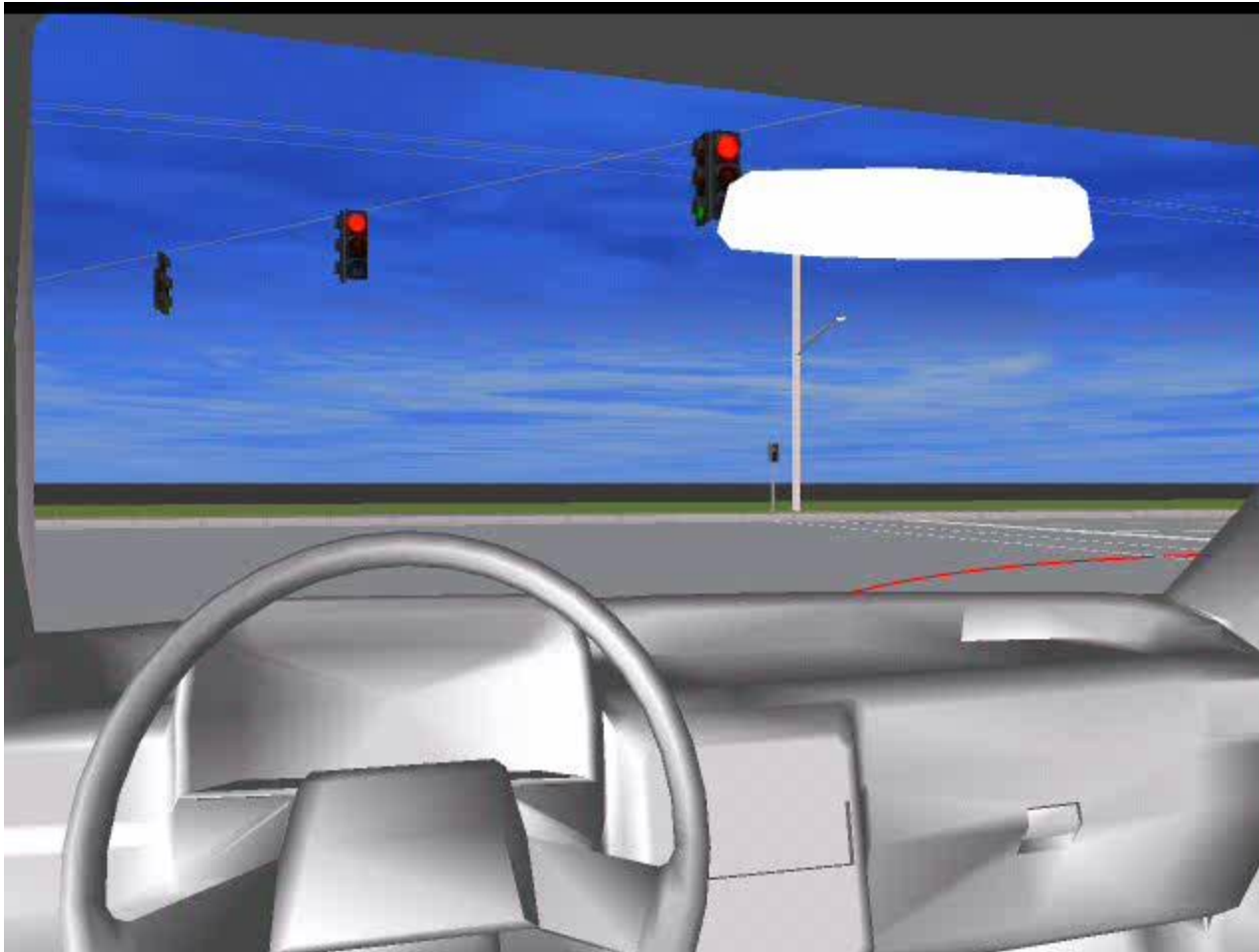
Accident Reconstruction



The Accident



Accident Animation (Driver's View)



StreetScenes[®]

3D Traffic Simulation

- *StreetScenes*[®] is a Virtual Reality (VR) software solution for 3D visualization of surface traffic
- 3D model of proposed soccer stadium in Rochester
- Used *StreetScenes*[®] to import output file from Synchro traffic simulation



MTV

Song: I'm OK (I Promise)

Band: Chemical Romance

IBC Digital & CCR Gaming Environment: Death Jr.



Public Forum



Virtual Reality

Alive on the Grid: PAAPAB

■ Networked art application for CAVE

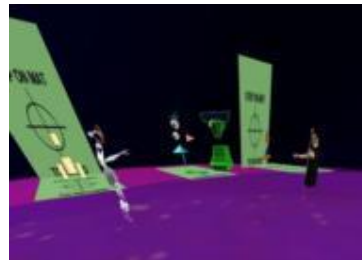
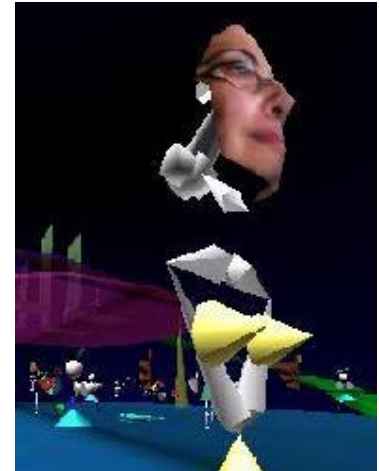
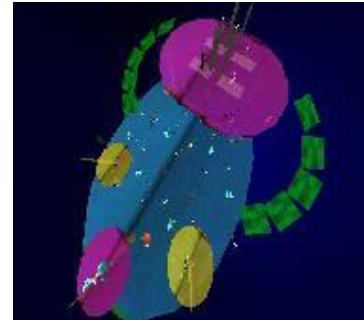
- Users from around the world
- First performance 2001

■ Dance-floor environment

- Inhabited by life-size puppets
- Dance with each other
- Synchro

■ Recording Booth

- User enters booth
- User dances
- System records dance from tracking on head and hands
- Dance mapped to Avatar



J. Anstey

The Thing Growing

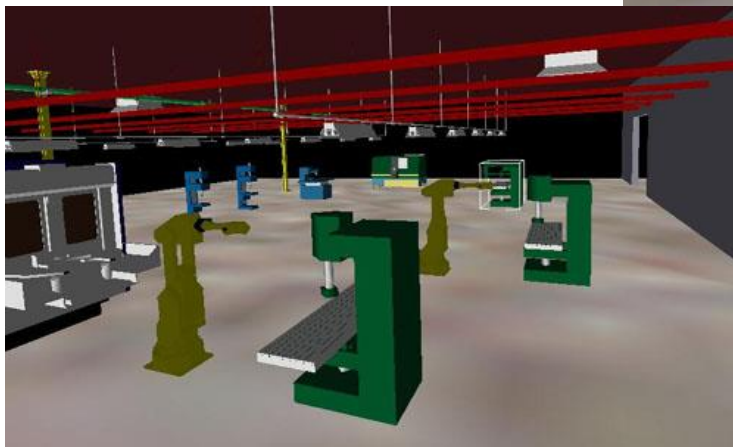
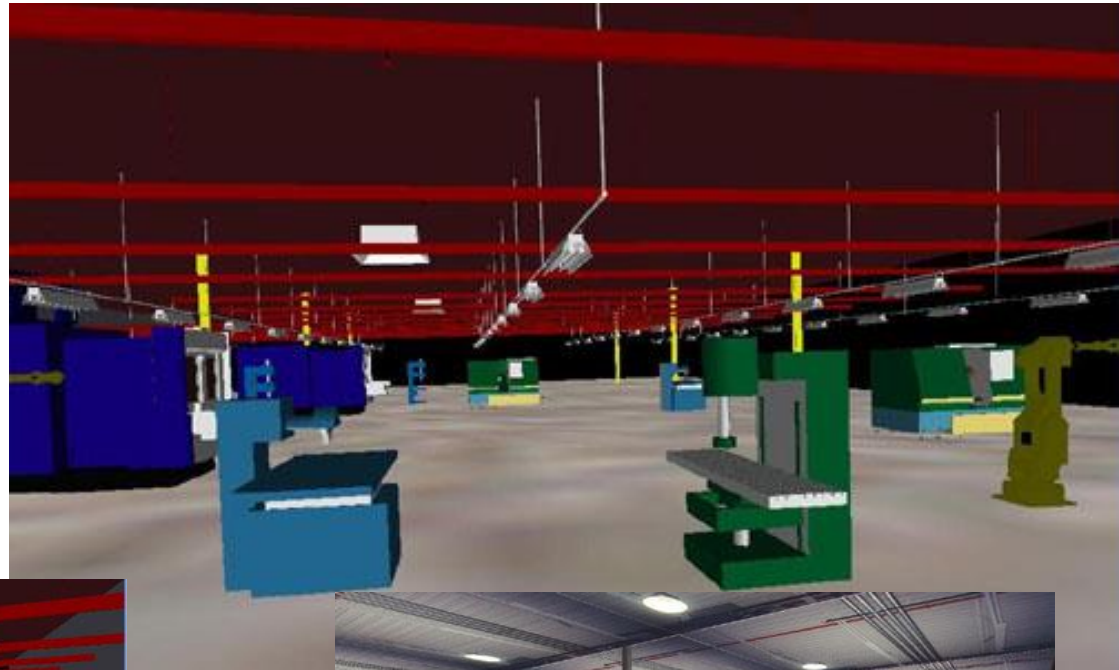
- VR work of fiction build for CAVE at EVL 1997-2000
- Users is protagonist
- User interacts with computer controlled characters
- Based on short story of J. Anstey



J. Anstey

VR-Fact!

- Interactive virtual factory
- Creates digital mock-up of factory
- Drag & place modular machines
- Mathematical algorithms for consistency checks



Kesh

Collaborative Visualization Environments

- Enable distributed collaboration via software developed at CCR
- Enable visualization and interaction with data across a geographically disparate network topology
- Integrate multiple data sources:
 - ❑ Scientific
 - ❑ Multimedia
- Research Topics
 - ❑ Distributed databases
 - ❑ OpenGL 3D programming
 - ❑ 3D Modeling
 - ❑ Character animation
 - ❑ User interaction
 - ❑ Virtual Reality



A. Ghadersohi, R. Miller, M. Green

Western New York

Some Facts



Buffalo, New York



- **The Queen City: 2nd Largest City in NYS**
- **City of Lights**
 - ❑ **First U.S. city to have electric street lights**
 - ❑ **Pan American Exposition (1901)**
 - **Pres. McKinley Shot**
- **Architecture**
 - ❑ **Frederick Law Olmsted**
 - ❑ **Frank Lloyd Wright**
- **Underground Railroad**
 - ❑ **Slaves escaped to freedom in Canada**
- **Four straight Super Bowl appearances**
- **Culinary Delights**
 - ❑ **Beef on Weck, Pizza, Fish Fries**
 - ❑ **(Buffalo) Wings: Anchor Bar, 1964**
- **Health Problems**
 - ❑ **Heart Disease/Stroke**
 - ❑ **Multiple Sclerosis**



Recent Biomedical Advances (Buffalo, NY)

■ PSA Test (screen for Prostate Cancer)

■ Avonex: Interferon Treatment for Multiple Sclerosis

■ Artificial Blood

■ Nicorette Gum

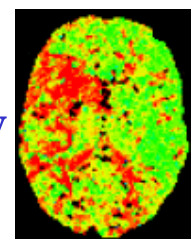
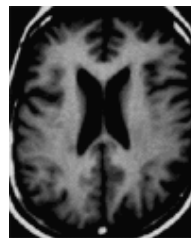
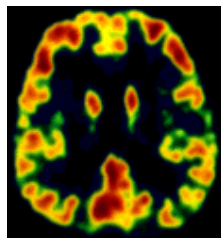
■ Fetal Viability Test

■ Edible Vaccine for Hepatitis C

■ Timed-Release Insulin Therapy

■ Anti-Arrhythmia Therapy

□ Tarantula venom



■ Direct Methods Structure Determination

□ Listed on “Top Ten Algorithms of the 20th Century”

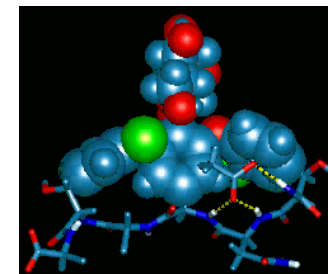
■ Vancomycin

■ Gramacidin A

■ High Throughput Crystallization Method: Patented

■ NIH National Genomics Center: Northeast Consortium

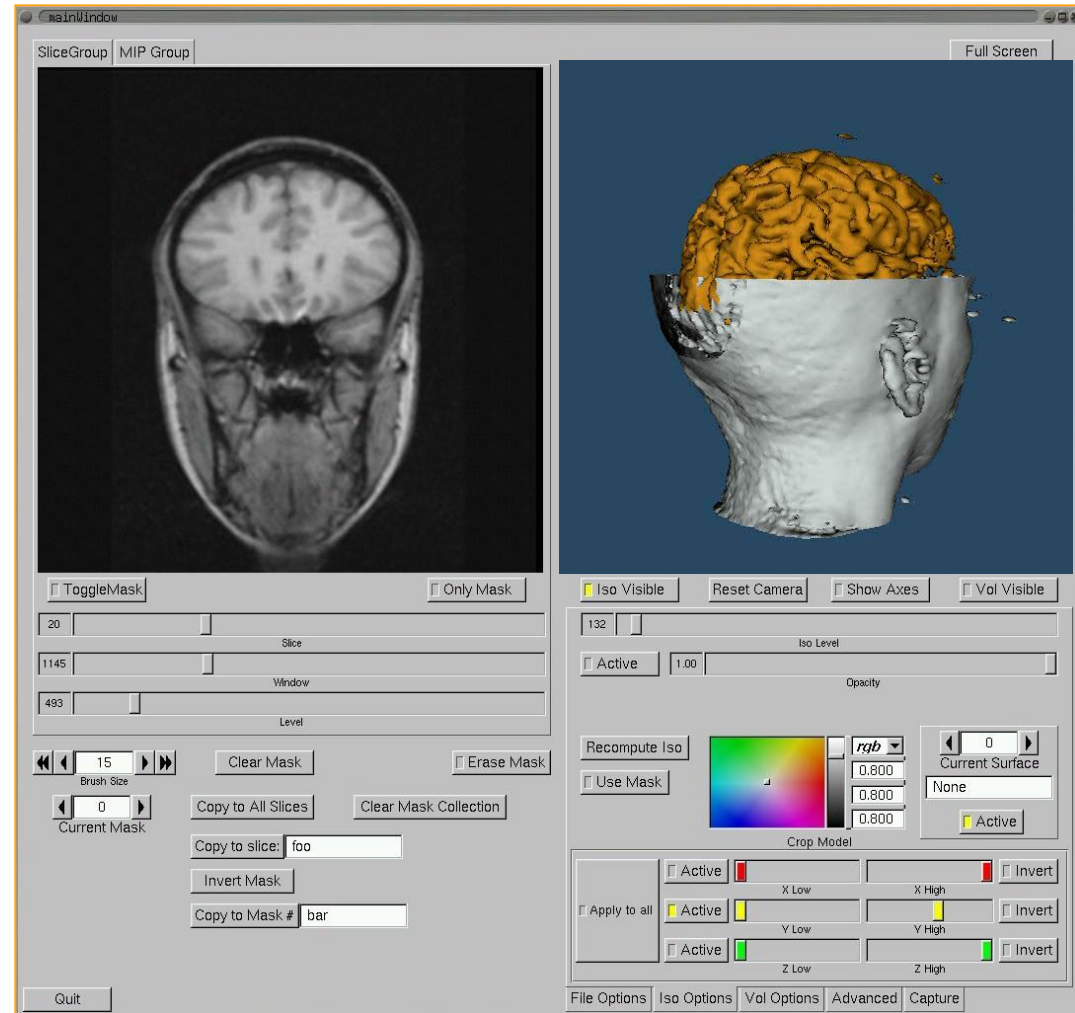
■ Howard Hughes Medical Institute: Center for Genomics & Proteomics



Scientific Visualization

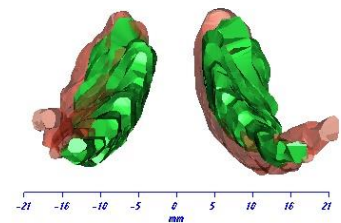
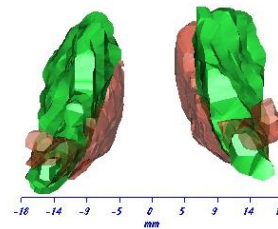
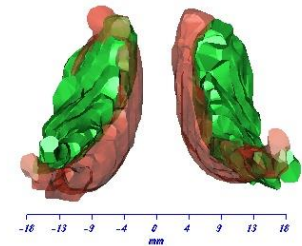
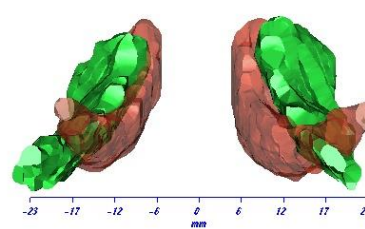
Multiple Sclerosis Project

- Collaboration with Buffalo Neuroimaging Analysis Center (BNAC)
 - Developers of Avonex, drug of choice for treatment of MS
- MS Project examines patients and compares scans to healthy volunteers



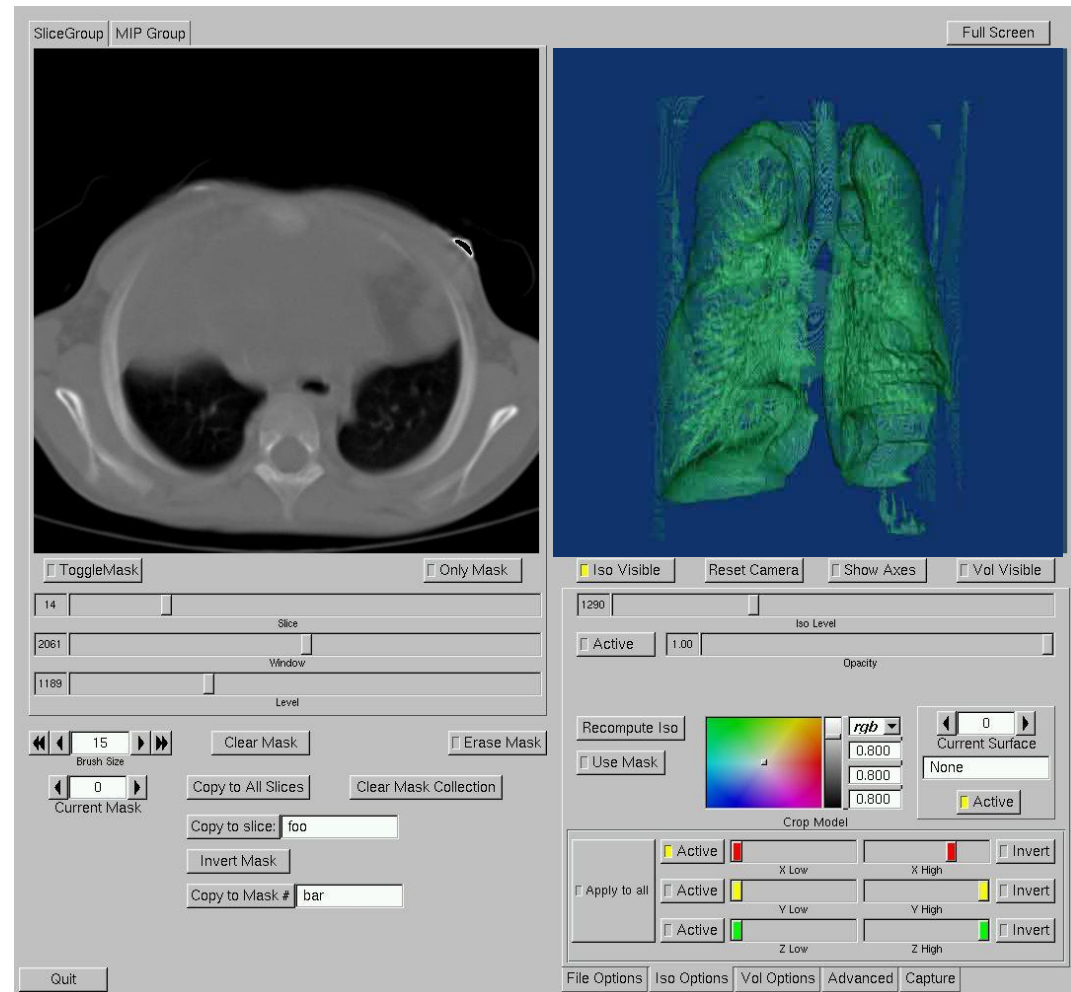
Multiple Sclerosis Project

- Compare caudate nuclei between MS patients and healthy controls
- Looking for size as well as structure changes
 - Localized deformities
 - Spacing between halves
- Able to see correlation between disease progression and physical structure changes



3D Medical Visualization

- Reads data output from a CT or MRI Scan
- Collaboration with Children's Hospital
- Visualize multiple surfaces and volumes
- Export images, movies or CAD file
- Pre-surgical planning
- Runs on a PC



M. Innus

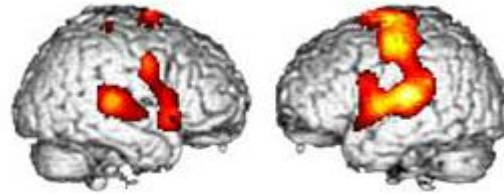
Mapping Brain Activity

Positron emission tomography (PET), shows sites activated and deactivated as subjects decide whether a sound is a target or not.

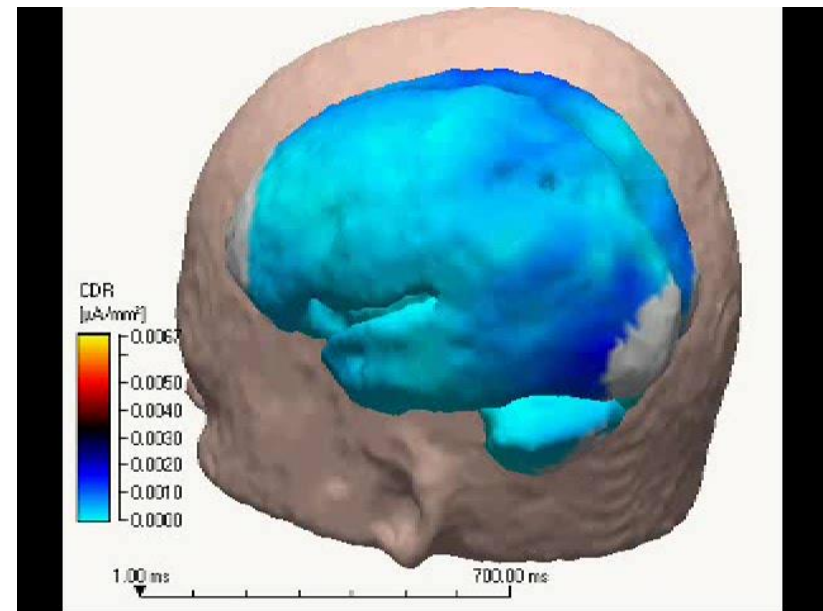
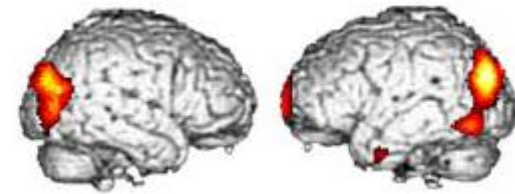
Current density maps of brain surface (1–700 ms after target) show dynamic pattern of brain activity during decision-making process.

A. Lockwood

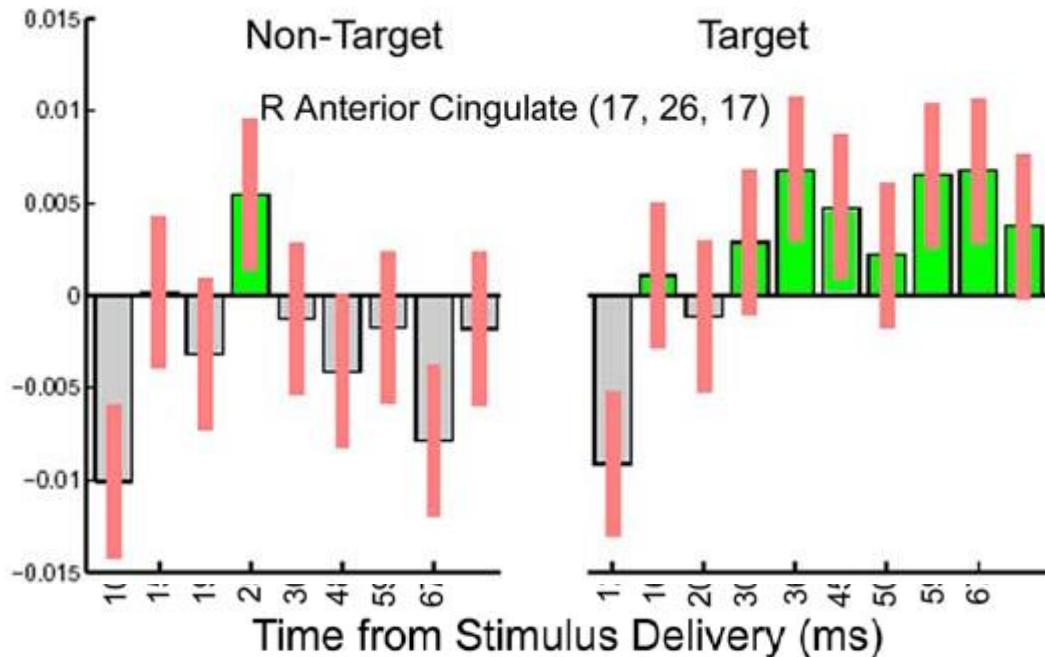
Sites Activated



Sites Deactivated



Mapping Brain Activity

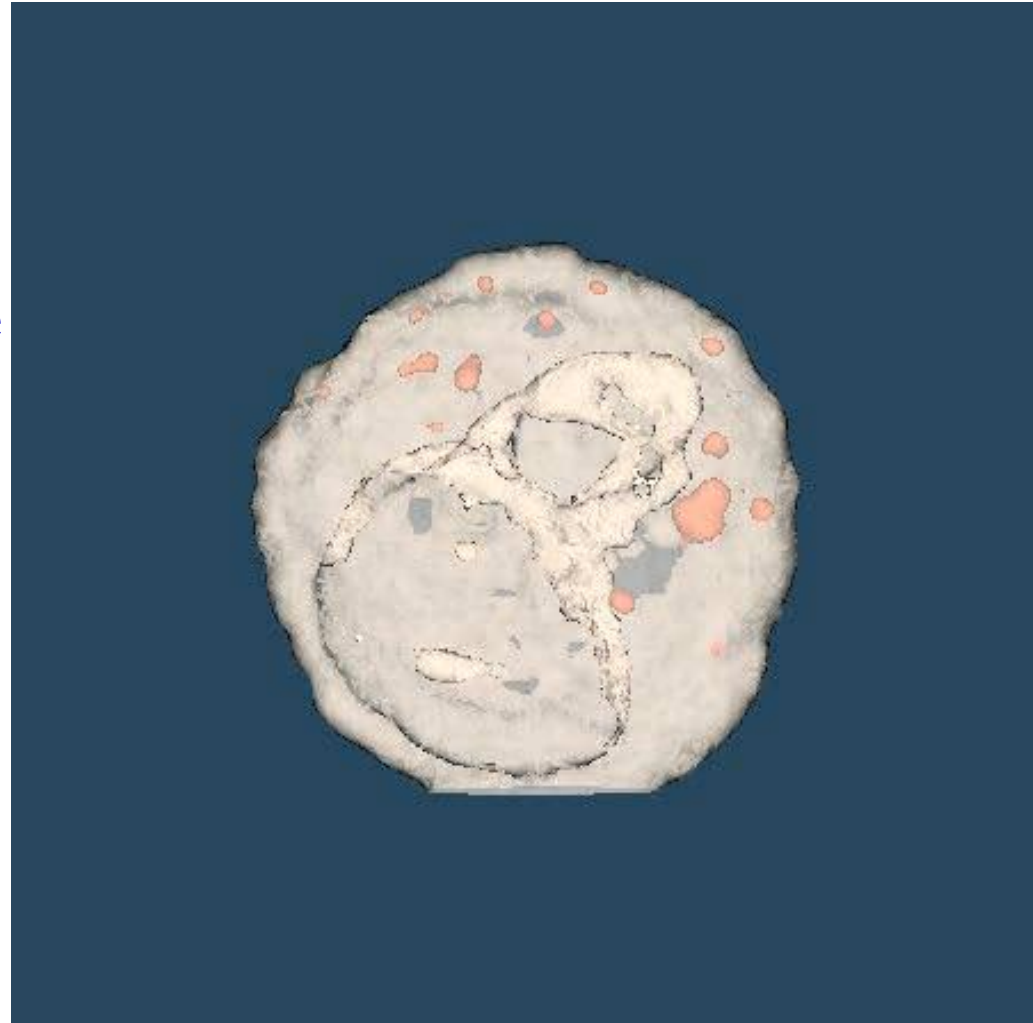


Temporal sequence of anterior cingulate cortex activation in response to targets and non-targets. This brain region controls attention-related neural activity. Green bars indicate significant differences compared to $T = 0$, the time of stimulus presentation.

A. Lockwood

Confocal Microscopy

- **3D Reconstruction of an Oral Epithelial Cell**
- **Translucent White Surface Represents the Cell Membrane**
- **Reddish Surface Represents Groups of Bacteria**

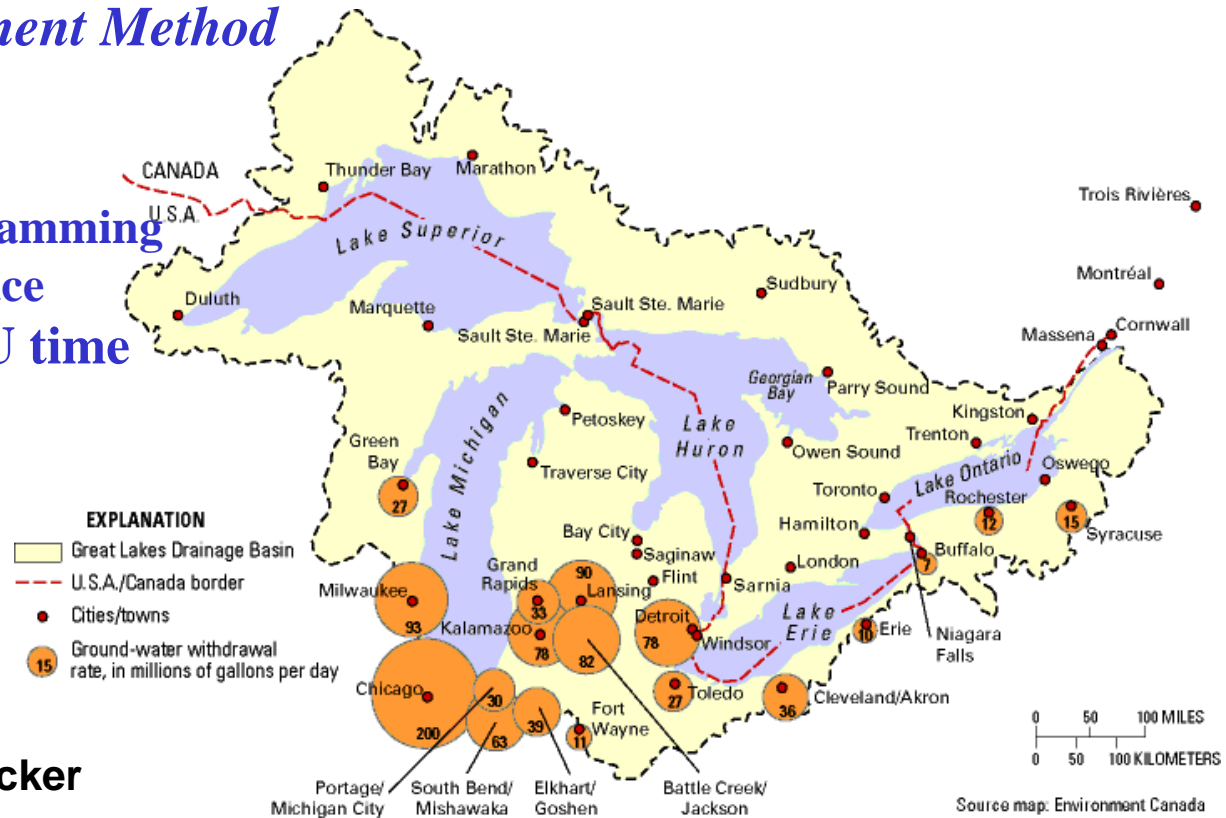


Science & Engineering

Small Subset of Projects

Groundwater Flow Modeling

- Regional scale modeling of groundwater flow and contaminant transport (Great Lakes)
- Ability to include all hydrogeologic features as independent objects
- Based on *Analytic Element Method*
- Key features:
 - Highly parallel
 - Object-oriented programming
 - Intelligent user interface
- Utilized 42 years of CPU time on CCR computers in 1 calendar year

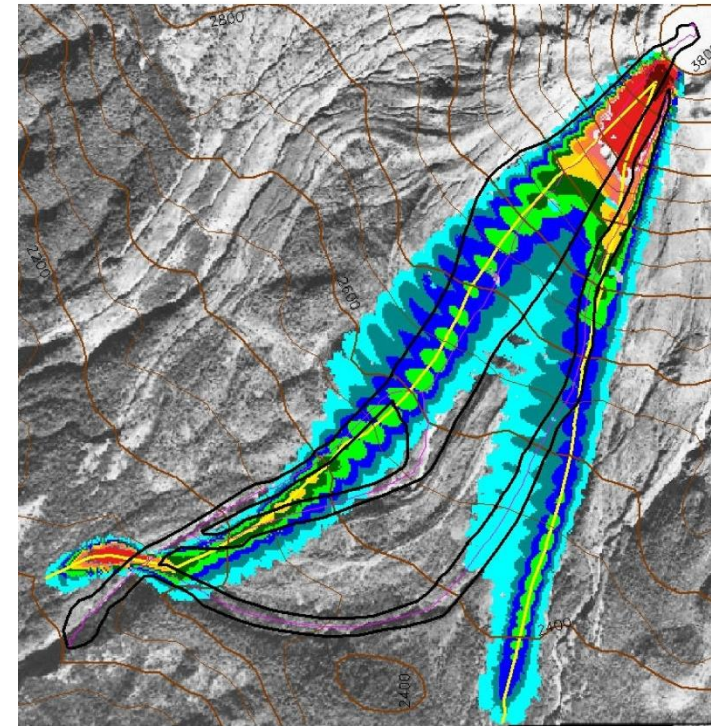


A. Rabideau, I. Jankovic, M. Becker

Avalanches, Volcanic and Mud Flows

Geology, Engineering

- Modeling of Volcanic Flows, Mud flows (flash flooding), and avalanches
- Integrate information from several sources
 - ❑ Simulation results
 - ❑ Remote sensing
 - ❑ GIS data
- Present information to decision makers using custom visualization tools local & remote
- GRID enabled for remote access
- Key Features
 - ❑ Parallel Adaptive Computation
 - ❑ Integrated with GIS System for flows on natural terrain

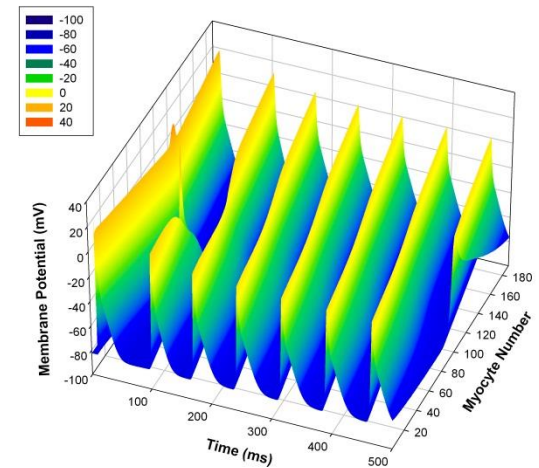
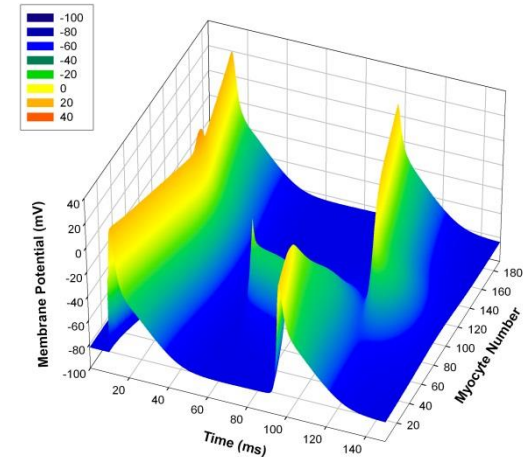


Flow models of Colima volcano
In Mexico – courtesy Rupp et. al.'06

A. Patra, B. Pitman, M. Sheridan, M. Jones

Cardiac Arrhythmia

- Comprehensive models of cardiac cells
- Modeling multicellular cardiac tissues and mechanisms of arrhythmias in the heart
- Simulation of genetic heart disease and arrhythmia suppression by drug application

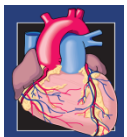
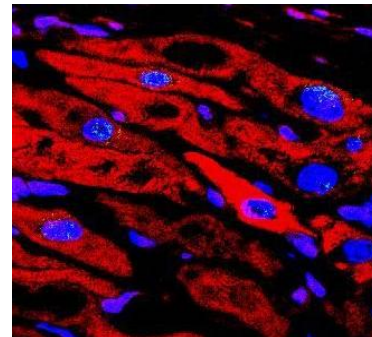
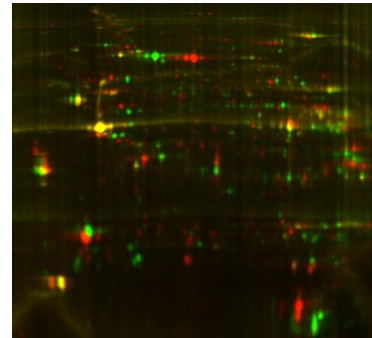
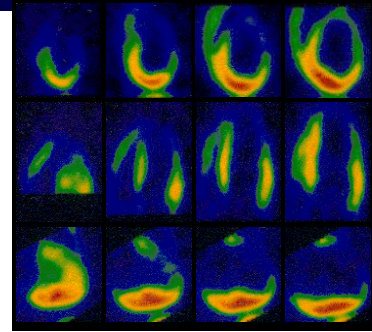


**Non-sustained and
sustained arrhythmia**

Center for Cellular and Systems Electrophysiology

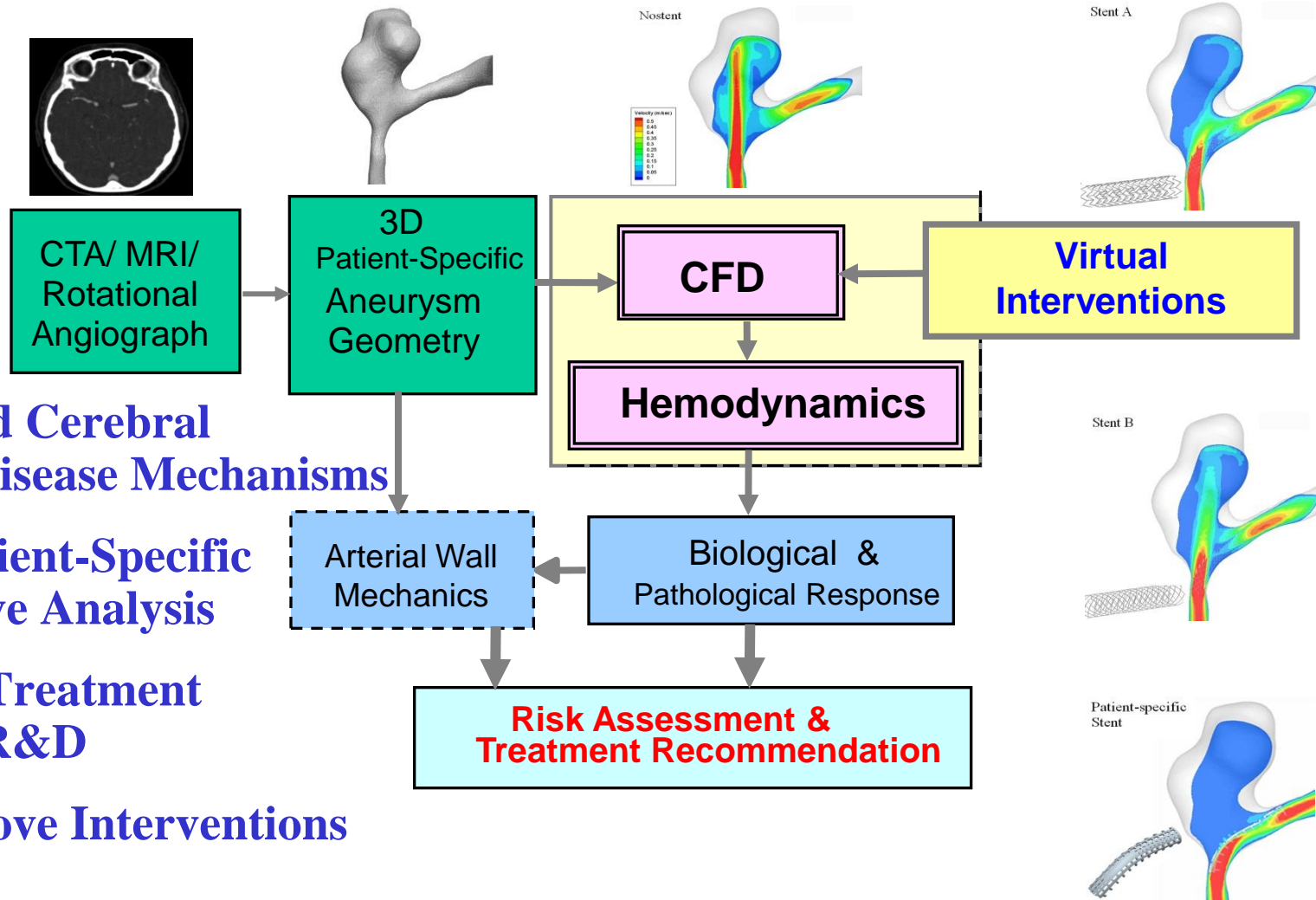
Cardiovascular Research

- **Molecular Imaging – PAREPET Clinical Study**
Analysis of cardiac PET (Positron Emission Tomography) scans aims to revolutionize assessment of an individual's risk for sudden cardiac death.
- **High-Throughput Discovery – Proteomics and Genomics**
- **Protein and gene expression profiling using differential in-gel electrophoresis and microarray technology provides a blueprint for the cellular mechanisms involved in hibernating myocardium.**
- **Translate results to identify gene and other therapeutic targets aimed at improving heart function and survival.**



Center for Research in
Cardiovascular Medicine

Cerebral Aneurysm: Virtual Intervention



■ Understand Cerebral Vascular Disease Mechanisms

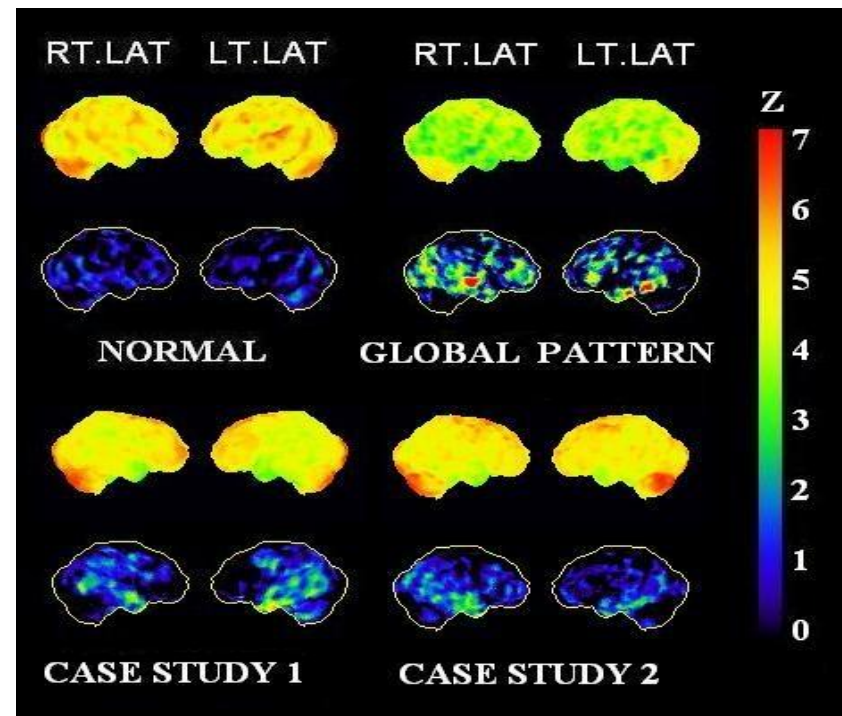
■ Enable Patient-Specific Quantitative Analysis

■ Diagnose, Treatment Planning, R&D

■ Help Improve Interventions

Vascular Dementia Imaging

- Early diagnosis of dementia from cerebral small vessel disease using computer analysis of SPECT Images
- Collaboration between Nuclear Medicine, CCR, Neurology, and Kaleida Stroke Center
- Funded by the Pfeiffer Foundation
- Fractal scores:
 - ❑ Normal 0.75
 - ❑ Global Pattern 1.13
 - ❑ Case Study 1 0.96
- Case Study 1 Moderate white matter and cortical hypoperfusion with visual memory, speed of processing, and verbal fluency deficits



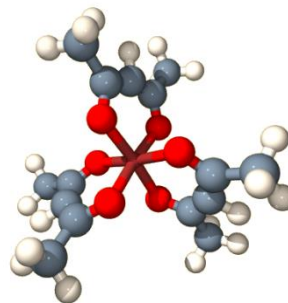
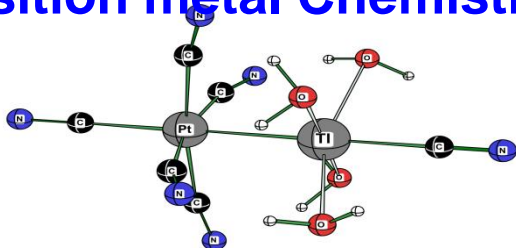
J. Baker, M. Innus

Theoretical and Computational Chemistry

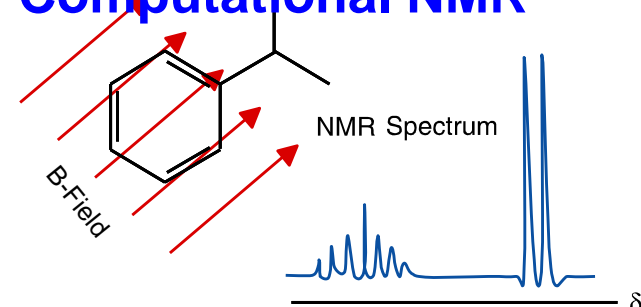
Chemistry

Applied to:

- Polypeptides
- Carbon Nanotubes
- Fullerenes
- Cluster Compounds
- Transition metal Chemistry



Computational NMR



Magnetic Properties of Molecules



Software Development

```
do ispin = 1, nspin
  nmo1 = nmo1(ispin)
  nmo2 = nmo2(ispin)/2
  noccv = noccv(ispin)

  if (dgnat) call moutp (hmat(1,ispin), nmo1, 'Delta V Matrix'//ispin)

  read constant part of h1 matrix

  clabel = 'fcf=90'
  call csadd (clabel, ..., ipert)
  call csadd (clabel, ..., nspin(ispin))
  call kfrdr (sufc, clabel, work, noccv, 1)

  compute f0cyc

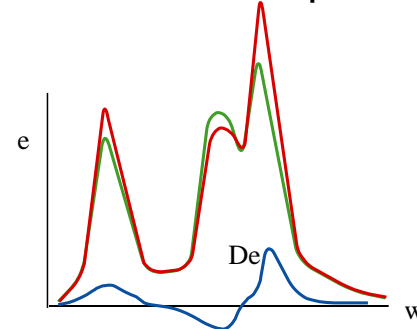
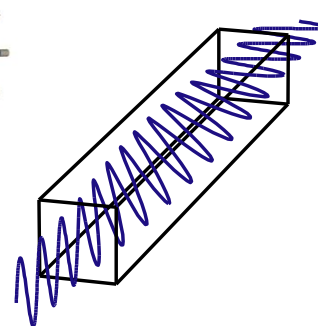
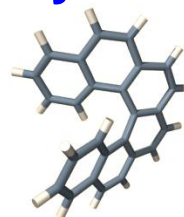
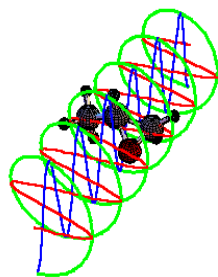
  if (.not. lnewoc .or. lastit) then
    rtemp = zero
    do ll = 1, noccv
      rtemp = rtemp + hmat(ll,ispin)**2
    end do
    f0cyc = f0cyc + rtemp
    if (ispin=1) f0cyc = f0cyc*two
  end if

  if (lastit .or. lconvt) then
    constant part of h1 matrix
    fcp1b,990 (f90 CVS:1,2)=1802-10-563
```

Theory and Calculation of Optical Activity

Optical Rotation

Circular Dichroism Spectra



J. Autschbach

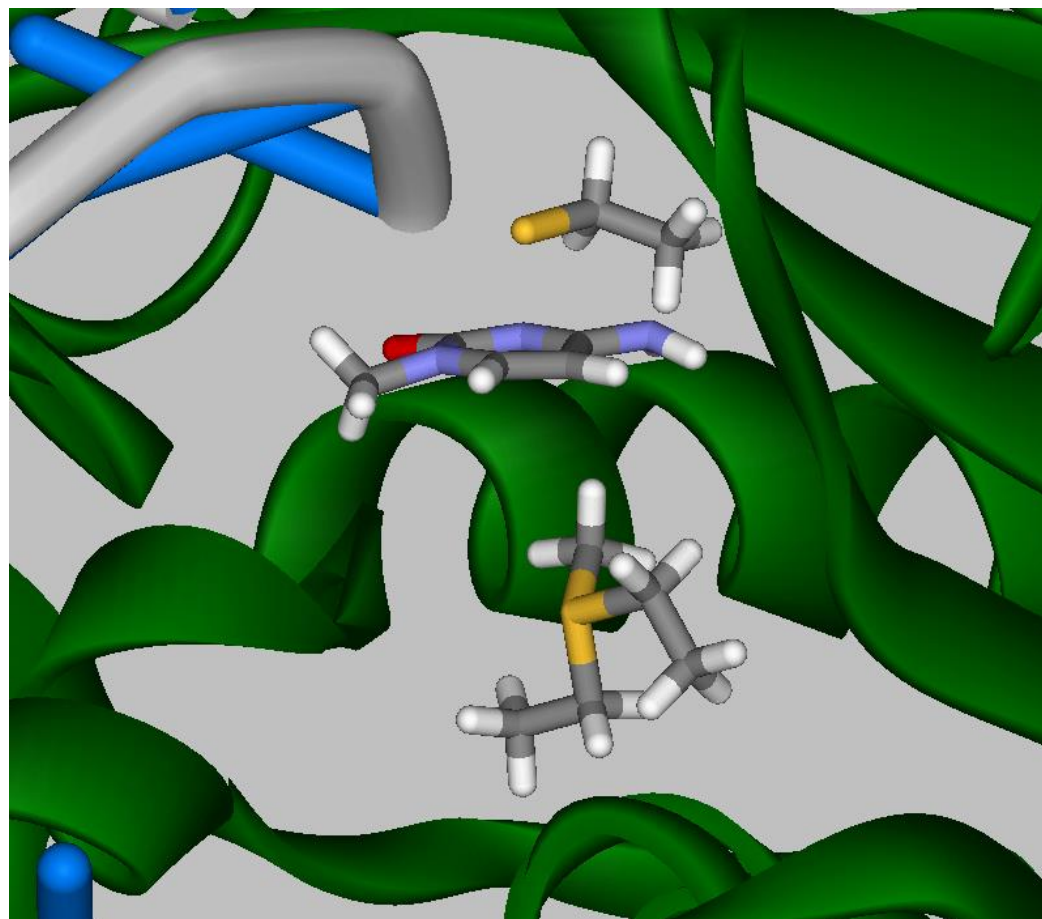
Understanding How Proteins Work

Collaboration with Merck Pharmaceutical Company

Modeling:

- DNA-Protein Interaction (understanding cancer)
- Drug-Protein Interaction (understanding blood clotting)

Movie shows a chemical reaction between a protein and DNA, which is responsible for some types of cancer.



M. Freindorf, T. Furlani

Computational Chemistry

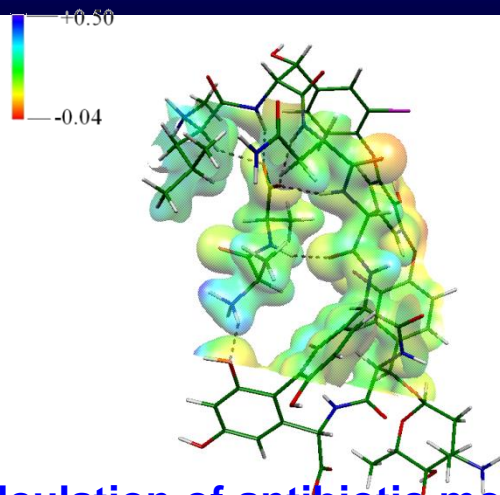
- UB Software development in Quantum Chemistry
 - Q-Chem – development of combined QM/MM methods for large molecular systems such as proteins
 - ADF – development of algorithms to calculate magnetic and optical properties of molecules
- Used to determine
 - 3D Molecular Structure
 - Electronic Spectra
 - Chemical Reactivity
- Applications
 - Pharmaceutical Drug Design
 - Industrial Catalysis
 - Materials Science
 - Nanotechnology



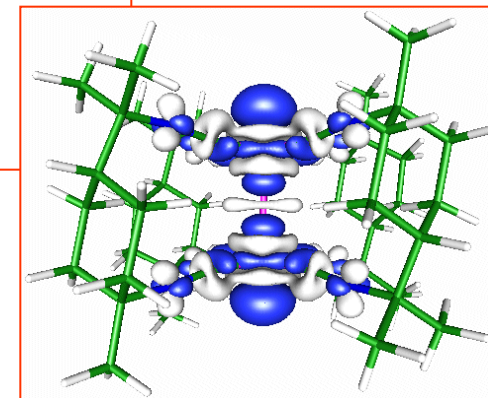
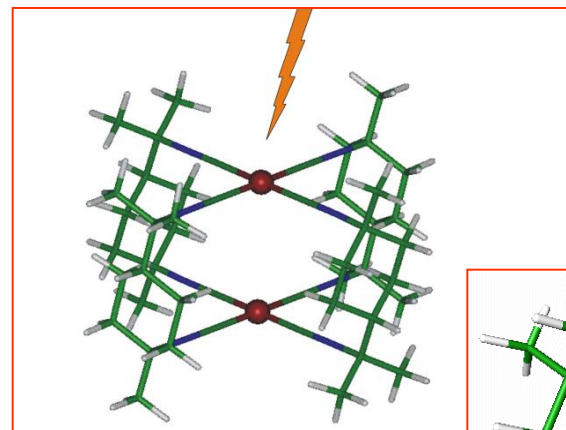
T. Furlani, J. Autschbach, M. Freindorf

Understanding Large Molecules and Fleeting Species

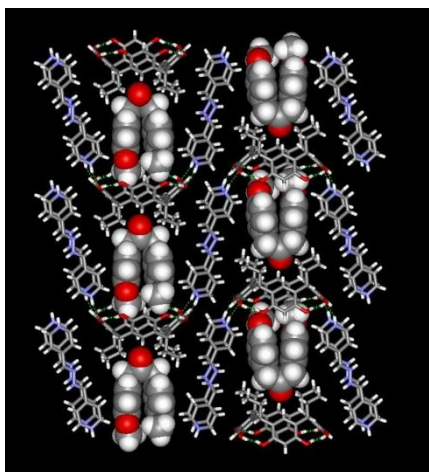
Chemistry



Calculation of antibiotic molecules:
electrostatic potential of vancomycin



A molecule changes on excitation by light



A supramolecular solid

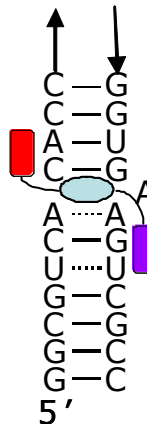
P. Coppens

Prediction of RNA Structure to Facilitate Design of Drugs Targeting RNA

Chemistry

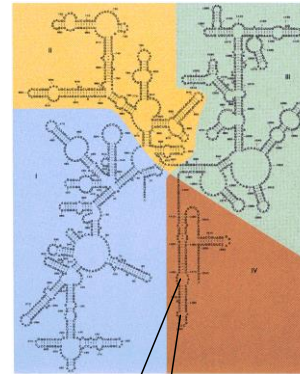
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 ggtcctgaa gaaagctgaa gacgacgag ttgataggcc ggtgtgttaa gcgcaogcat
 cgttgagct aaccgtact aatgaacctg gaggctaac ct

23s rRNA Sequence



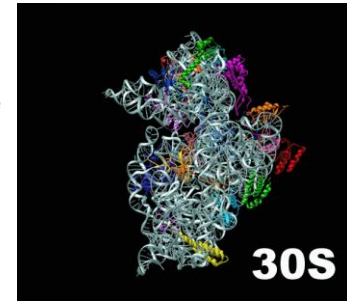
$$\Delta G_{\text{ligand}} = \Delta G_1 + \Delta G_2 + \Delta G_3 + \Delta G_{\text{linker(s)}}$$

RNA Secondary Structure Prediction

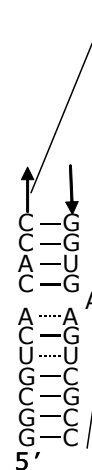


23s rRNA 2D Structure

RNA 3D Structure

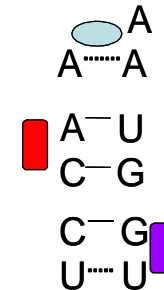


Modular Assembly to increase affinity



A-site
Aminoglycoside target

Search Small-Molecule-RNA Ligand Database for RNA motifs to which we have identified an organic ligand



Results from ligand database search

M. Disney

3D Structure of Proteins

- **Direct Methods for Crystal Structure Determination**
 - Listed on “Top Ten Algorithms of the 20th Century”
- **UB/HWI collaborative software development**
 - **SnB** – determine protein heavy-atom substructures
(<http://www.hwi.buffalo.edu/SnB/>)
 - **BnP** – determine complete protein structures
(<http://www.hwi.buffalo.edu/BnP/>)
- **Applications to drug design**
 - AIDS
 - Arthritis
 - Cancer
 - Heart disease
 - SARS



R. Miller, C. Weeks



Hauptman-Woodward Institute

Determining 3D Protein Structure

Structural Biology

- NMR-based Structural Biology and Structural Genomics
- Bio-NMR Methodology
- NMR-based Metabonomics in Cancer Research

Structural Biology

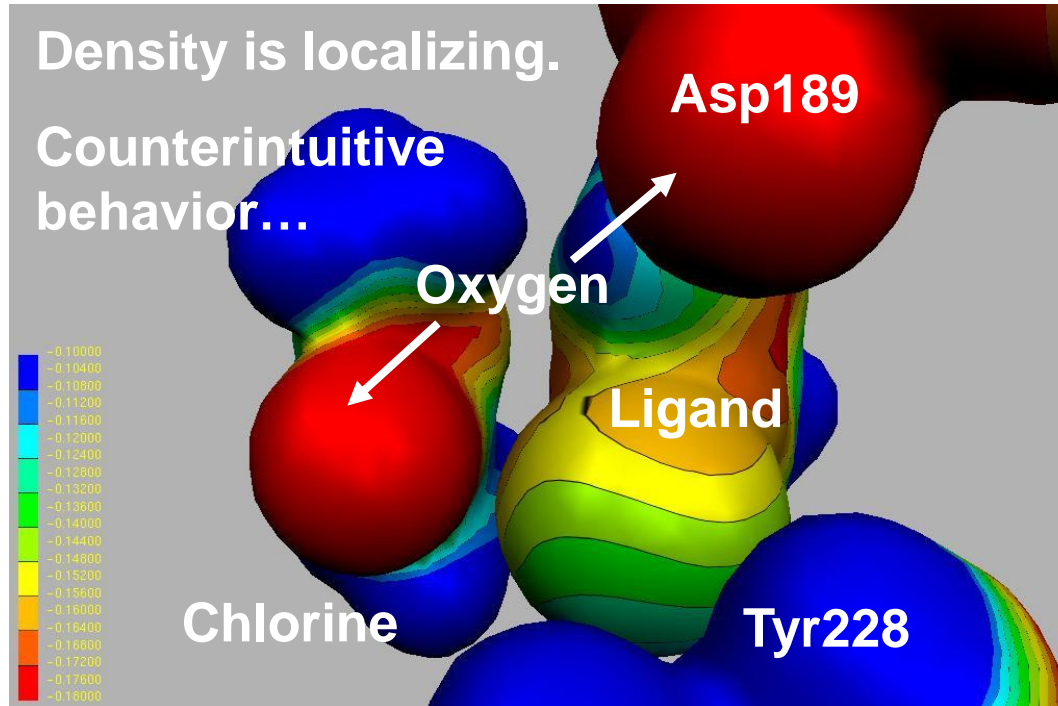


Propelled by Recent Advances, NMR Moves Into the Fast Lane

A speedy new NMR technique could finally help structural genomics groups achieve their goal of devising factory-style approaches to mapping protein structures at high speeds

T. Szyperski

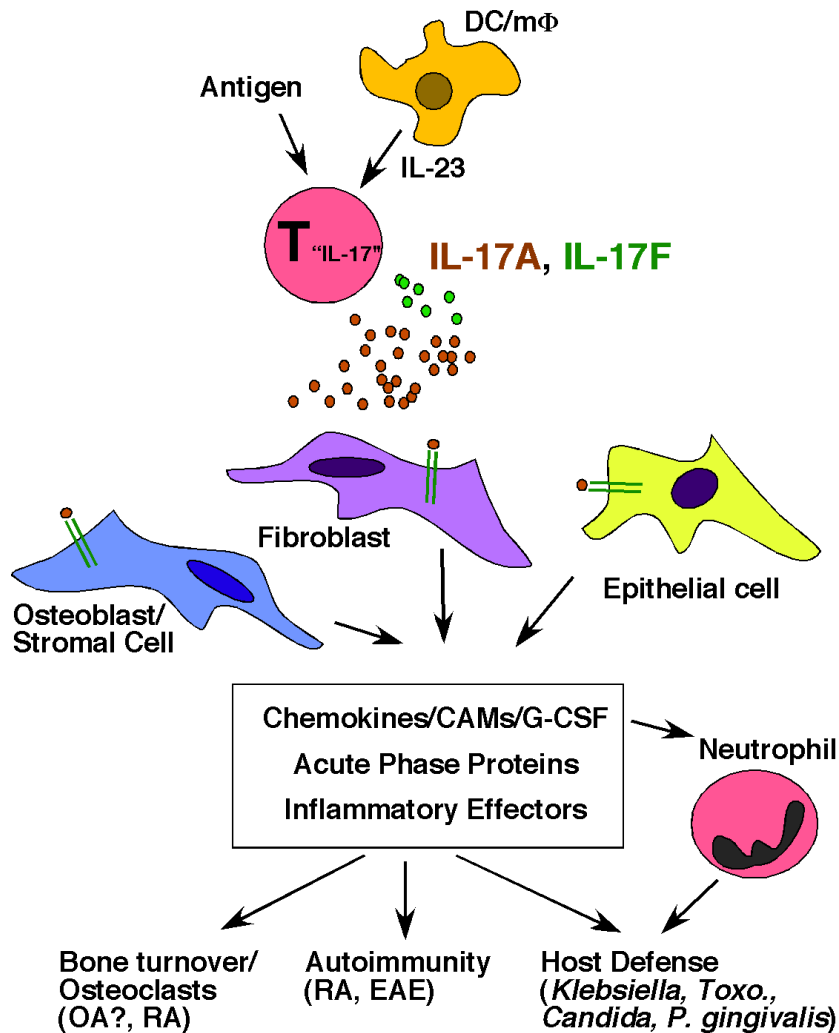
Binding in a Drug-Receptor Complex



- Ligand docked with residues of the active site of thrombin. Electrostatic potential map superimposed onto the electron density isosurface.
- The goal is to elucidate the thermodynamics of molecular recognition in binding.

D. Hangauer, M. Freindorf

Defining Cytokine Signaling Mechanisms



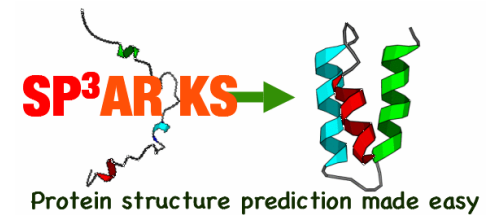
S. Gaffen, Z. Hu

- T cells secrete cytokines such as IL-17 to promote host defense and/or autoimmunity
- Microarrays used to define IL-17 gene targets in various cell types
- Computational and statistical approaches used to compare the promoters of IL-17 target genes in mouse and human genomes to identify conserved transcription factor binding sites (TFBS), with the ultimate goal of understanding how IL-17 mediates molecular signals
- IL-17 target promoters contain conserved TFBSs, including NF- κ B and C/EBP

Computational Biology & Bioinformatics

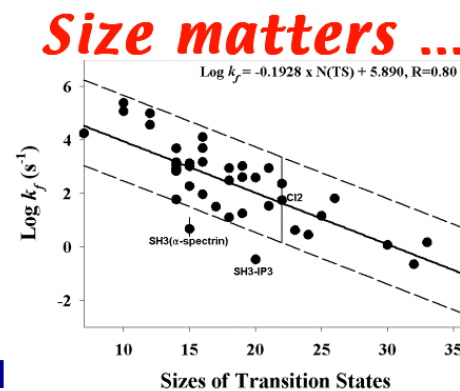
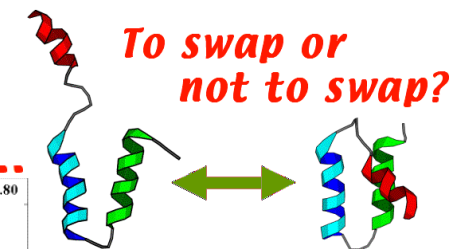
■ Development of Bioinformatic Tools

- ❑ **SPEM** – align multiple sequences for discovering hidden evolution information of genes.
- ❑ **SPARKS/SP³** – predict three-dimensional structures of proteins by matching a query sequence with known structural templates.
- ❑ **DFIRE** – predict binding affinities of protein-protein, protein-ligand, and protein-DNA complexes for structure-based drug design.



■ Mechanistic study of protein folding and binding

■ <http://theory.med.buffalo.edu>

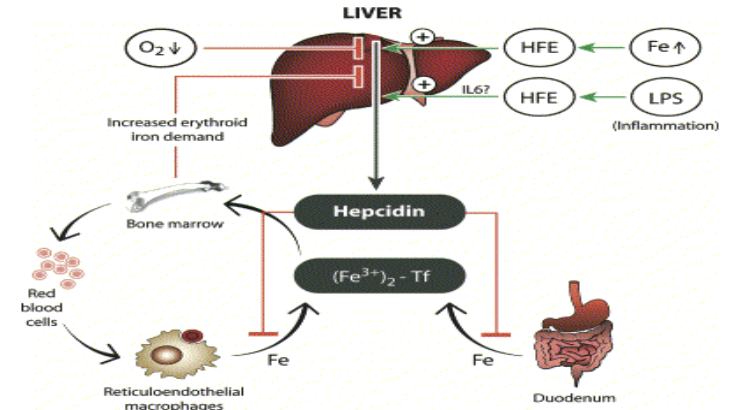


M. Halfon <http://theory.med.buffalo.edu>

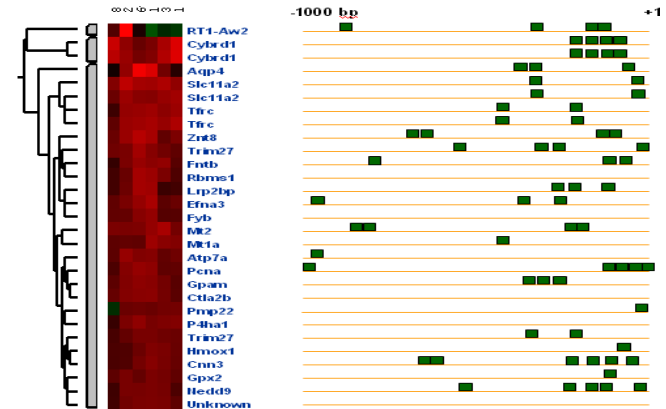
Genome-Wide Study of Iron Homeostasis

Bioinformatics

- Physiological and Heparin-hormonal regulators are involved in iron homeostasis
- Intestinal iron transport controls overall body iron homeostasis
- Computational biology to discover new genes involved in iron homeostasis
- Systems biology to reveal regulatory pathway responding to iron status
- Known and novel genes are regulated according to iron status
- Sp1 or related TFs may be involved in regulating expression of some genes during iron-deficiency



Hentze M.W., et. al. Cell 117(3):285-97, 2004



Distribution of SP1 on promoters of genes induced during iron-deficiency

J. Collins, Z. Hu

Regulation of Gene Expression

- **REDfly** (Regulatory Element Database for Fly) Database of verified transcriptional regulatory elements
- Over 650 entries
- Most comprehensive resource of animal regulatory elements
- Fully searchable, has DNA sequence and gene expression data, links to other databases

The image displays three overlapping screenshots of the REDfly website. The largest screenshot on the left shows the homepage with the title "REDfly Regulatory Element Database for Drosophila" and a "Welcome to REDfly" message. Below the welcome message, there is a paragraph describing the database as a curated collection of known *Drosophila* transcriptional cis-regulatory modules (CRMs). A second paragraph explains that the database includes all experimentally verified fly CRMs along with their DNA sequence, associated genes, and expression patterns. A third paragraph provides information on how to contribute or correct data. At the bottom, it identifies the project as a collaboration between the Halfon lab and the Center for Computational Research at the State University of New York at Buffalo.

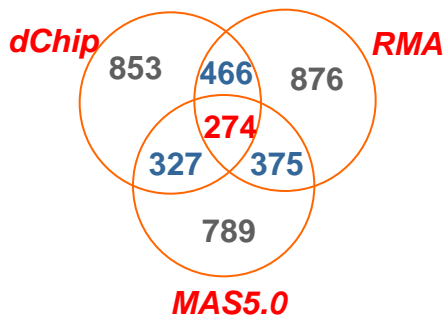
The middle screenshot shows a search interface with a search bar and a list of search results. The results list various CRM entries with their IDs (e.g., 375.8, 579.3) and associated gene names (e.g., ap, brain, BS1.0, BS1.1, BS2.0, BS3.0, CE7002, CE7003, CE7004). There are checkboxes for "Exact Match" and "Base Pairs".

The rightmost screenshot shows a detailed view of a specific CRM entry. The "Species" is *Drosophila*, the "Element Name" is UEG.0, and the "Gene Name" is ey. The "Location" is 4:719644..721428. The "Evidence Term" is "reporter construct (in vivo)". The "Sequence Source Term" is "sequence inferred from restriction map - left end estimated/uncertain". The "Includes Promoter" is "No". The "PubMed Id" is 14568101. The "Sequence" is provided as a text block. The "Expression Terms" list various developmental stages and tissues: FBtc:00001060 embryonic brain, FBtc:00005662 embryonic central brain, FBtc:00005666 embryonic central brain mushroom body, FBtc:00001063 embryonic protocerebral neuromere, and FBtc:00001919 larval central nervous system. The "Notes" field is empty.

M. Halfon, S. Gallo

Data Mining and Analysis

- GeneChips hybridized with cRNAs of biological interest
- Many probe set algorithms for summarizing expression intensity
- Significant differences of differentially expressed genes generated by different algorithms from the same dataset



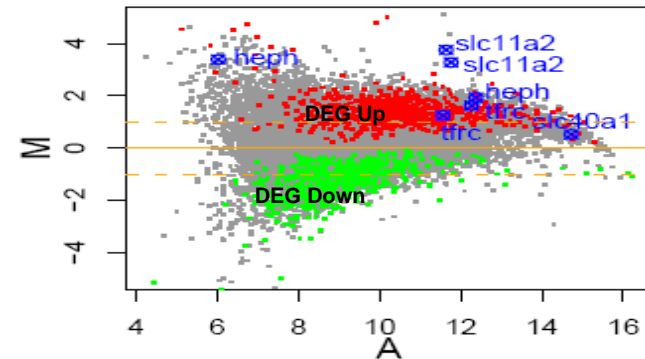
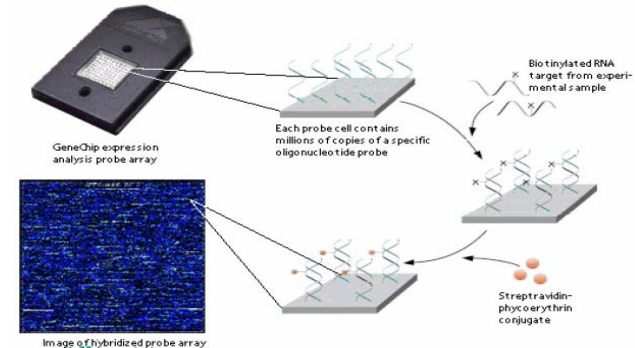
31% - 55% overlap

Which algorithm is best?

Great impact on subsequent expression data analysis

- Novel statistical approach for data variance and result bias analyses
- No external reference data needed
- Algorithm evaluation with direct applications to experimental datasets of interest

Z. Hu, G. Willisky



BMC Bioinformatics



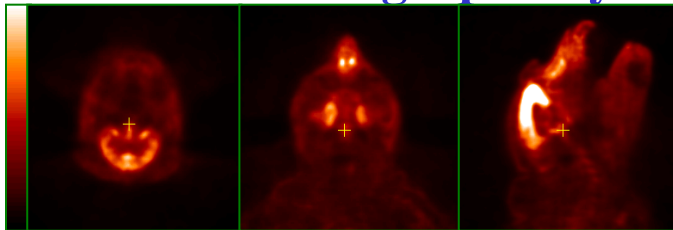
Methodology article

Open Access

Utilization of two sample t-test statistics from redundant probe sets to evaluate different probe set algorithms in GeneChip studies

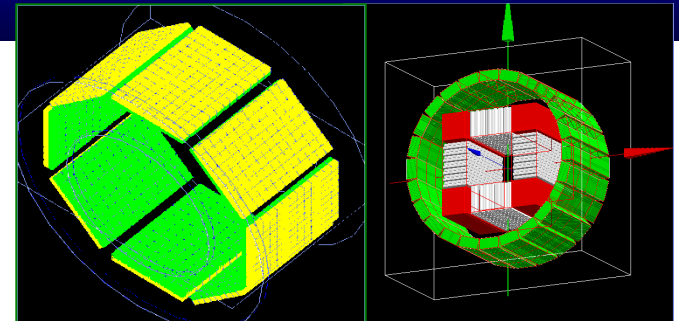
Nuclear Medicine

- **Monte Carlo simulation**
 - for modeling imaging system characteristics, optimizing system design, and validating data correction algorithms.
- **Image reconstruction**
 - for development of high resolution image reconstruction algorithm and software for both human and animal nuclear emission tomographic systems.

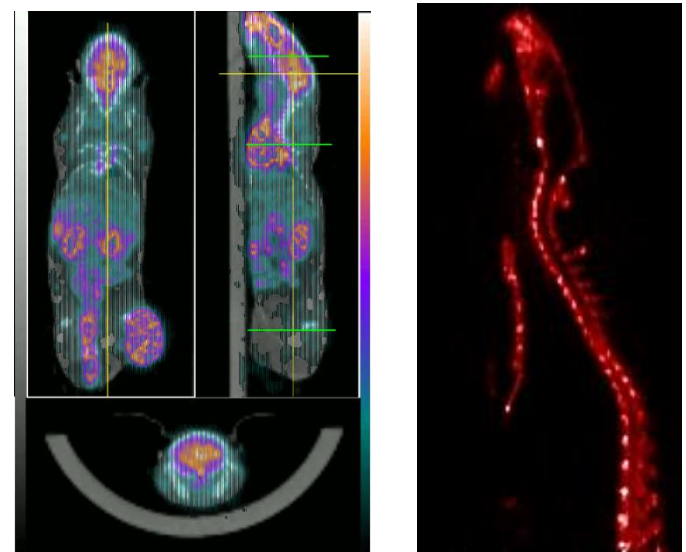


Transverse, coronal and sagittal views of a monkey brain scanned on a dedicated brain PET using the radioligand ^{18}F -FCWAY.

Y. Yao, M. Jones



Two virtual imagers simulated for system modeling and design evaluation.



Co-registered ^{18}F -FDG PET and CT mouse images (left) and a ^{18}F -Fluorine bone image of a 250 gram rat.

Parallel Algorithms

- String pattern matching searches for word processors, Web, molecular biology
- Image processing
- Computational geometry
- Fundamental operations

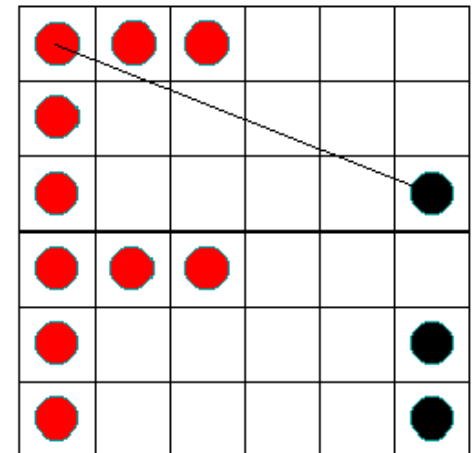
ALGORITHMS
SEQUENTIAL
& PARALLEL
A UNIFIED APPROACH
Second Edition



Computer Engineering Series

ROSS MILLER / LAURENCE BOXER

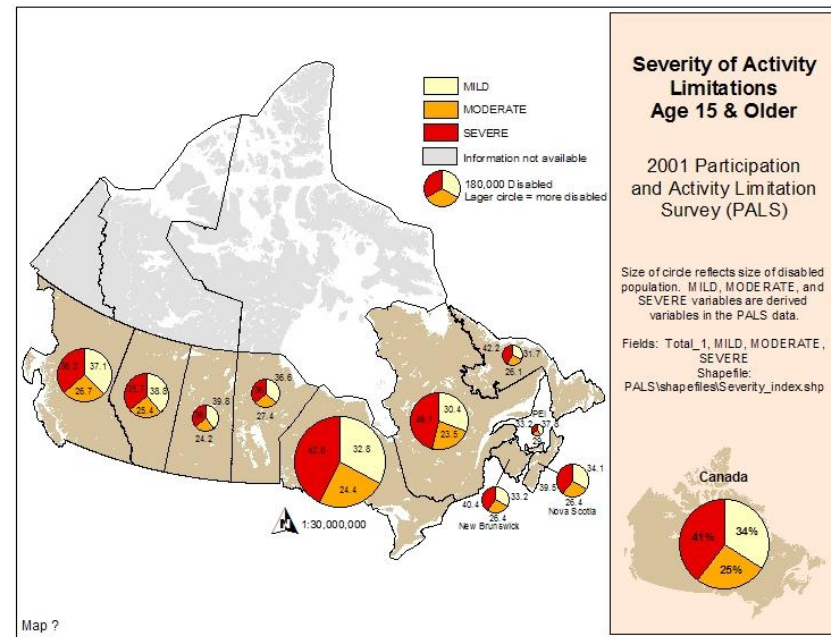
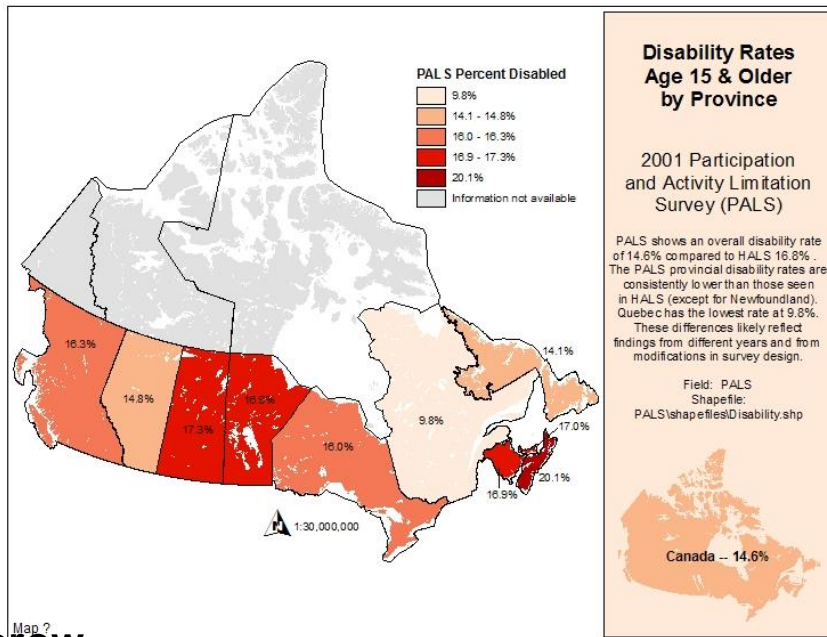
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T:	A	l	g	o	r	i	t	h	m	i	c	M	a	t	h	e	m	a	t	i	c	s	
P:		M	a	t																			
			M	a	t																		
				M	a	t																	



L. Boxer, R. Miller

Literacy & Disability in Canada

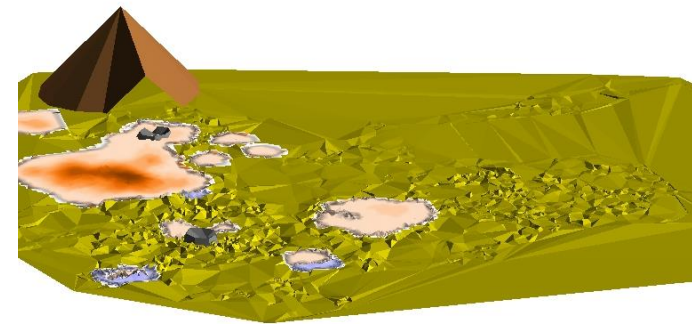
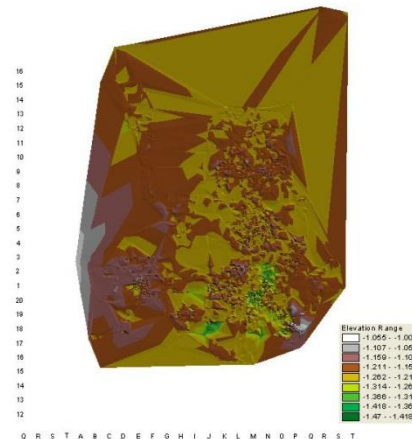
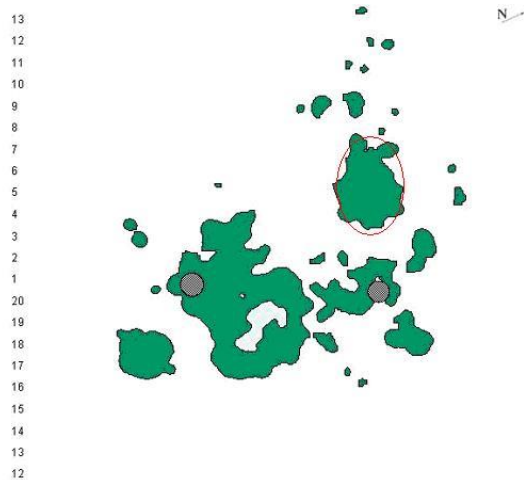
- Exploring the relationship between illiteracy & disability across the Canadian landscape
- Social Systems GIS Lab in the Dept. of Anthropology is working with researchers from York University & the Canadian Abilities Foundation.
- Sponsored by The Adult Learning & Literacy Directorate of the Ministry of Human Resources & Social Development Canada.



E. Zubrow

Verberie Paleolithic Site in France

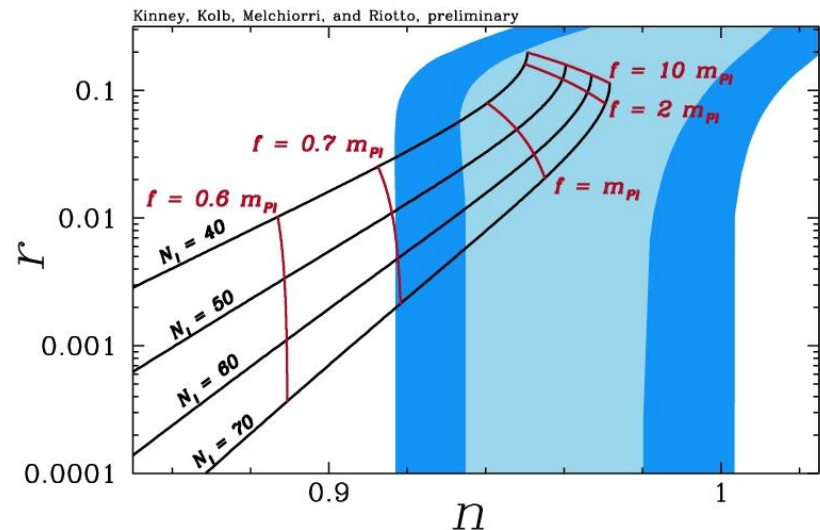
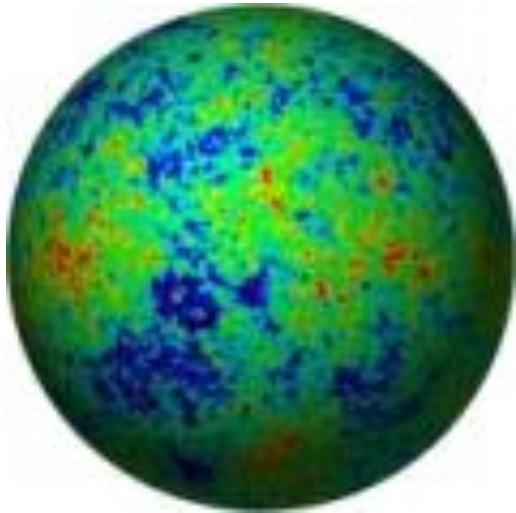
- Intrasite spatial analysis and 3D modeling of the a Late Upper Paleolithic archaeological site in the Paris Basin of France
- Social Systems GIS Lab in the Dept. of Anthropology is working with researchers from the CNRS in Paris
- Sponsored by the National Science Foundation



E. Zubrow

Cosmological Parameter Estimation

- Wealth of new precision cosmological data
- WMAP Cosmic Microwave Background Measurement
- Sloan Digital Sky Survey: 3-D map of a million galaxies
- Interpret implications of data for models of the first trillionth of a second of the universe: *inflation*
- *Monte Carlo Markov Chain data analysis: stochastic exploration of many-dimensional parameter spaces*

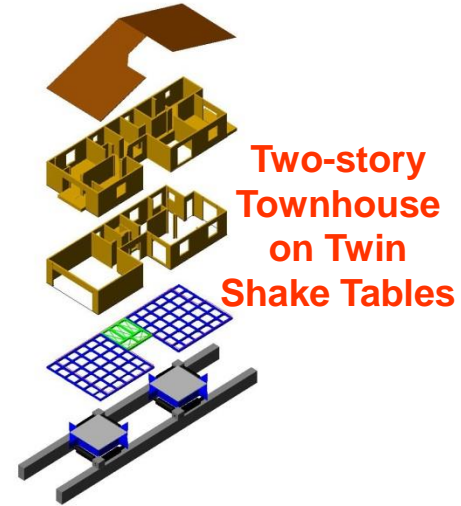


W. Kinney

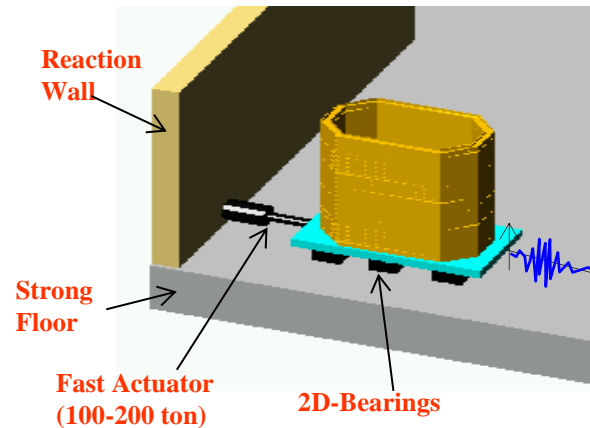
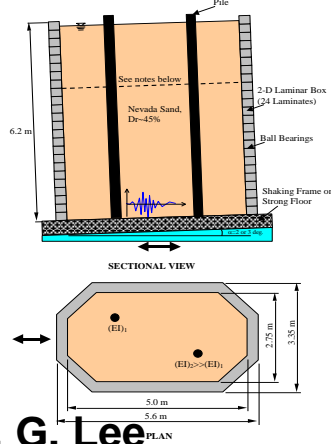
UB's Structural Engineering and Earthquake Simulation Laboratory (SEESL)

Structural Engineering

NEESWood:
Development of a
Performance-Based
Seismic Design for
Woodframe
Construction:



Two-story
Townhouse
on Twin
Shake Tables

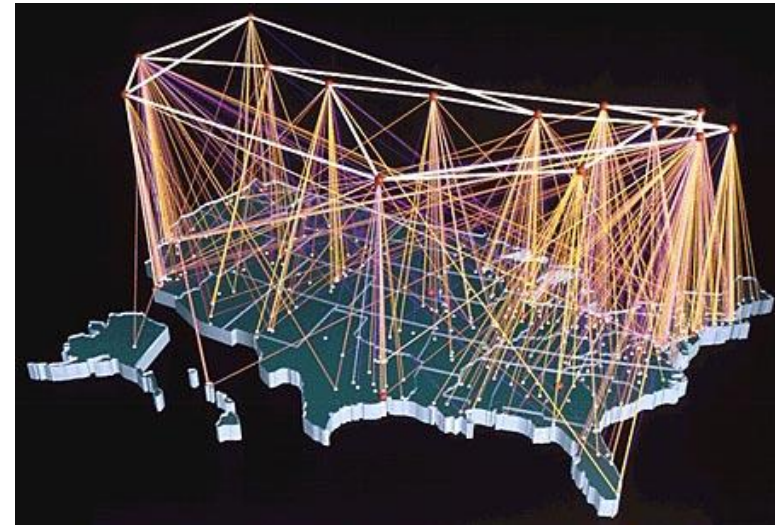


2-D
Geotechnical
Laminar Box
Tests of Pile
Foundations
Subjected to
Soil
Liquefaction

M. Bruneau, A. Reinhorn, G. Lee

Cyberinfrastructure in STEM Education

- Developing a scalable, multi-site cyber-infrastructure for Science, Technology, Engineering and Mathematics (STEM) education and training called *MyDesignSpace*.
- Implementing a digital design repository to enhance instruction and learning in STEM education.
- *MyDesignSpace* will also help bridge existing gaps between secondary and collegiate STEM education.



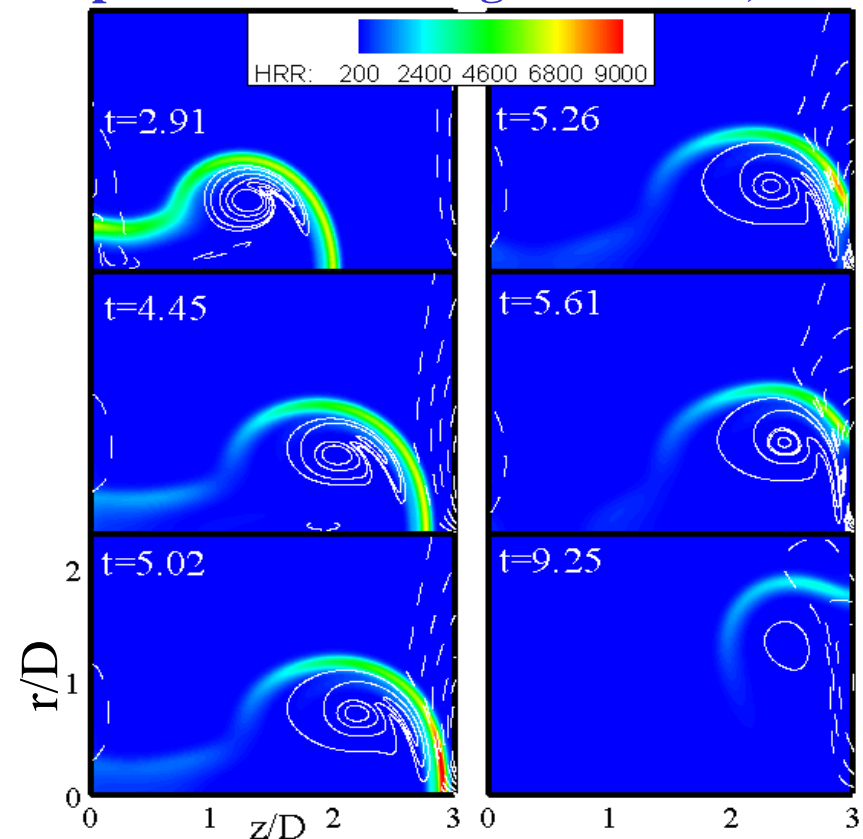
NYSCEDII

New York State Center for Engineering
Design and Industrial Innovation

www.nyscedii.buffalo.edu

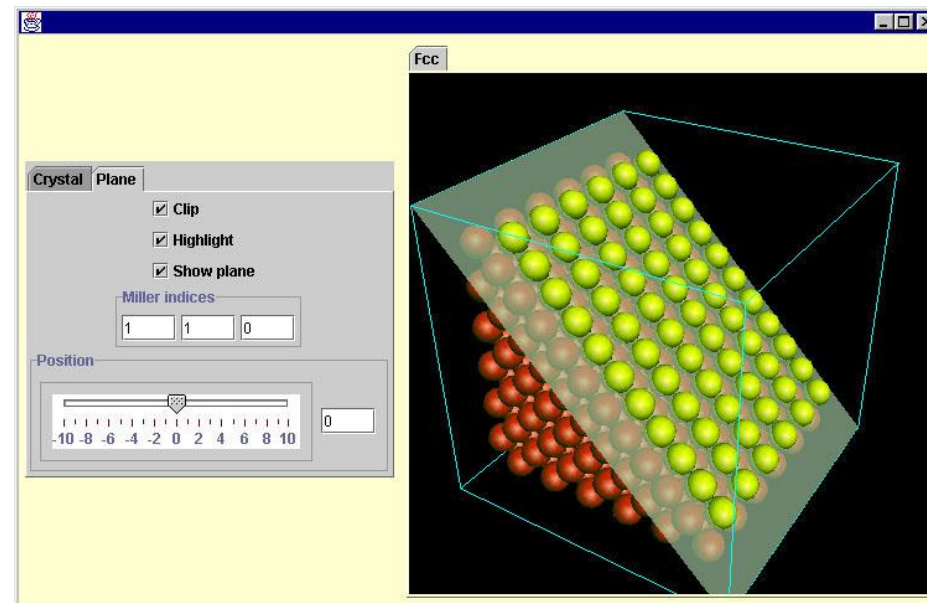
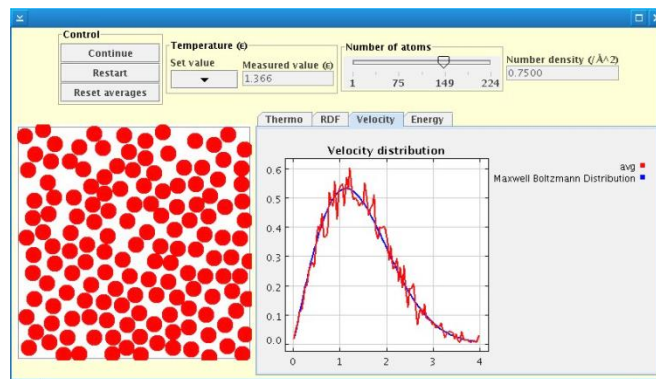
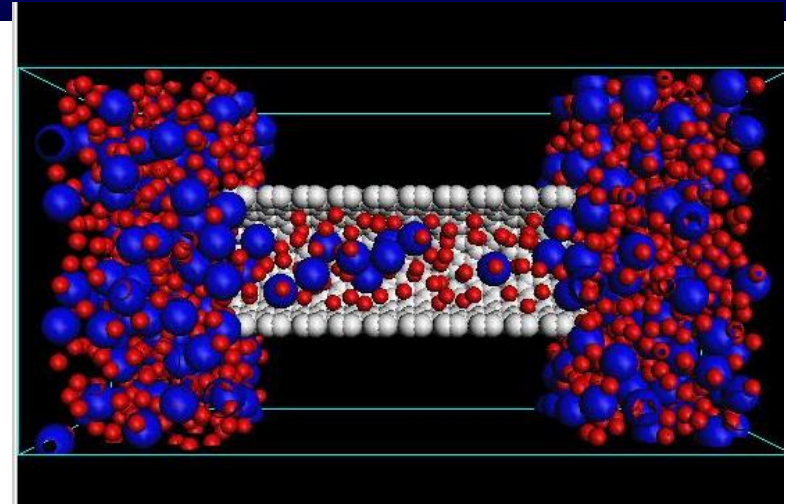
Understanding Combustion

- Flame-wall interaction modeling for a non-premixed flame propelled by a vortex ring.
- In this figure different time instants are shown during the interaction. White line contours and color contours represent vortex ring and flame, respectively.
- Key Features:
 - Modeling of Detailed GRI3. Mechanism for Methane Combustion
 - Parallel algorithm using mpi
 - 85-90% Parallel efficiency for up to 64 processors
- FWI study is important to determine
 - Engine Design
 - Quenching Distances
 - Flame Structure
 - Unburned hydrocarbon
 - Maximum Wall heat fluxes



Molecular Simulation Software

- Molecular simulation has wide application in existing and emerging technologies
- Recent advances in information technology make simulation more broadly accessible
- *Etomica* development environment permits easy construction of simulations
- Object-oriented, Extensible, Interactive, Portable and Adaptable
- Stand-alone simulations can be constructed as a teaching tools



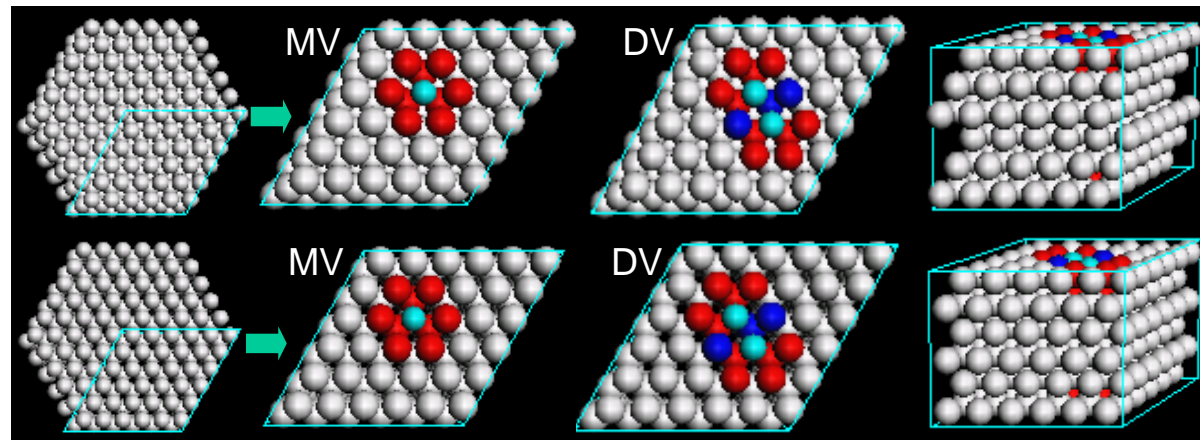
D. Kofke

Computational Materials Science

- Molecular and mesoscale modeling used to understand the behavior of materials
- Example application: **Electromigration**
- Strong electrical currents cause movement of atoms in metal
- Result is large defects that lead to failure of electrical connection
- Consequences can be catastrophic
- Interdisciplinary experimental/ modeling studies leading to understanding of behavior



Photos of metal lines that have developed voids (above) and hillocks (below) due to electromigration. (source: www.nd.edu)

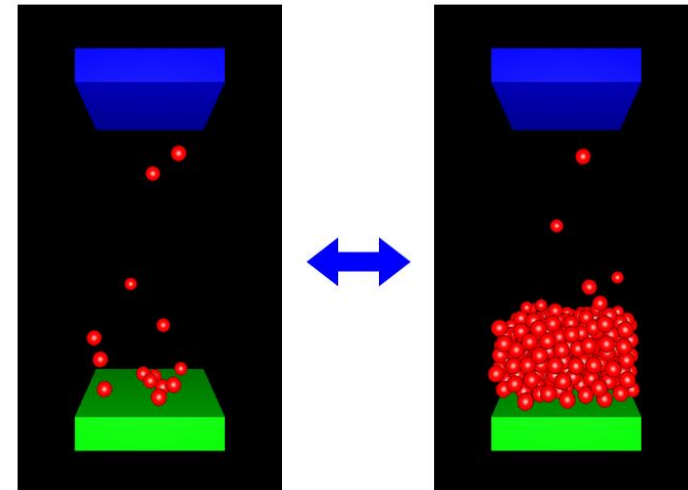
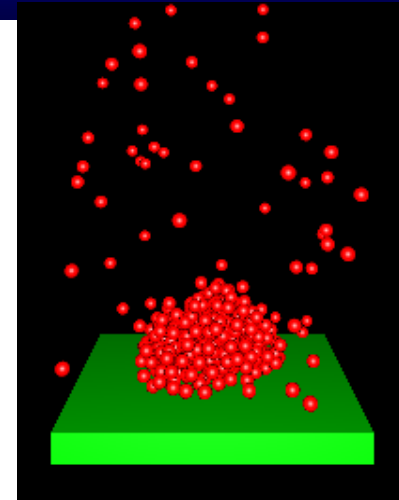


Simulation cells of solids with mono- and di-vacancies (light blue spheres), highlighting atoms neighboring the defects.

D. Kofke

Nano Confinement of Fluids

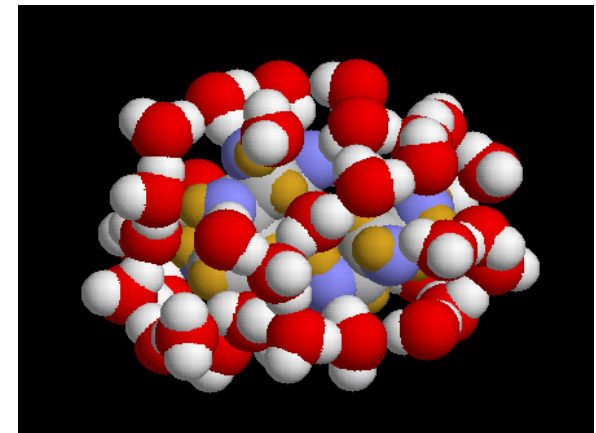
- Fluids in the presence of one or more surfaces exhibit rich phase behavior that can be strikingly different than that observed for bulk fluids
- A fundamental understanding of the relationship between a system's microscopic interactions and the phase behavior of a system is essential for the development of novel materials
- Molecular simulation is a useful tool for developing these relationships through the use of model systems that mimic the behavior of real fluids



J. Errington

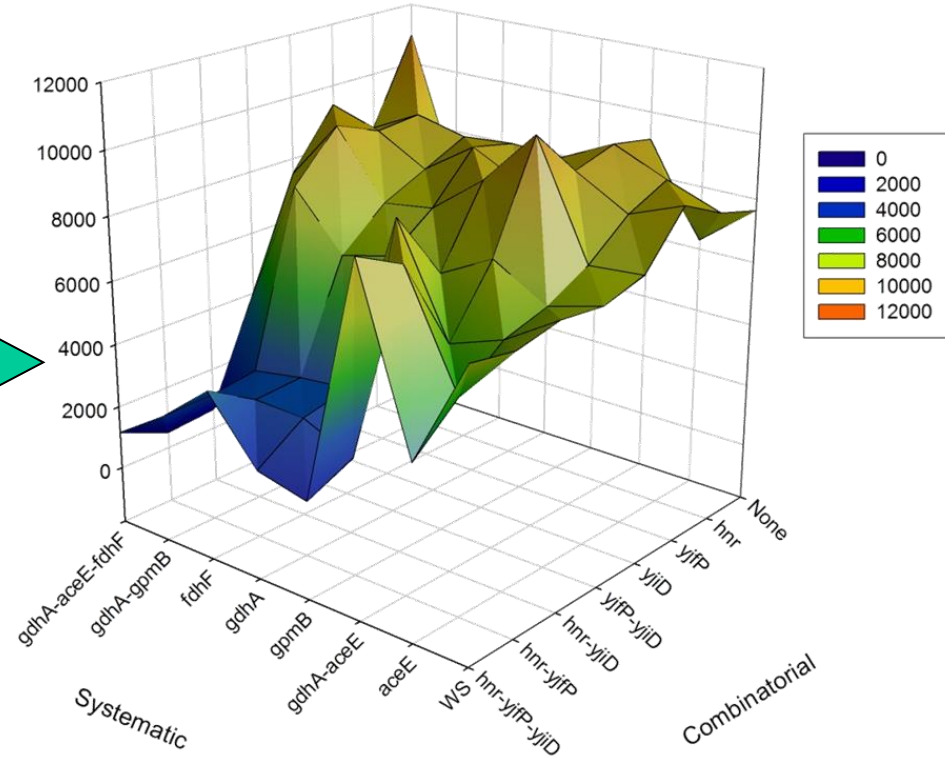
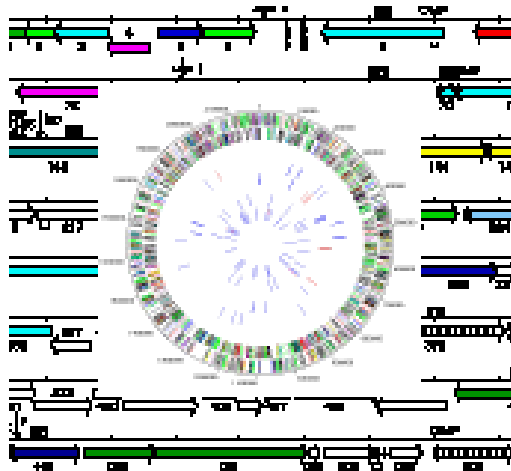
Aqueous Solutions

- The behavior of water and aqueous mixtures plays a key role in biology, chemistry, physics, and the design of many chemical and biological processes
- To gain a fundamental understanding of aqueous solutions, one must consider the effect the microscopic hydrogen-bond network has on the macroscopic properties of the system
- The goal of our program is to obtain a more complete understanding of aqueous systems using this molecular approach
- The diagram is a snapshot from a molecular dynamics simulation that depicts the organization of water molecules within 3.5 \AA of a trehalose molecule



J. Errington

Designing Cellular Phenotypes

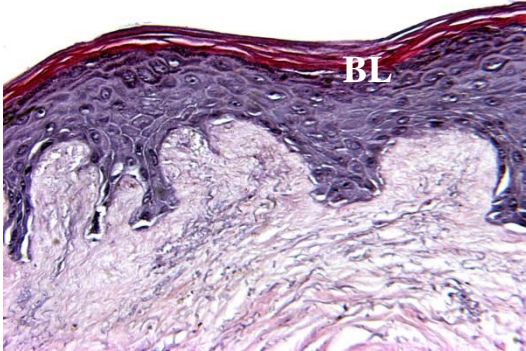


- Genome-wide metabolic models of sequenced microorganisms
- Optimization of metabolic and cellular phenotypes
- Goal is to design biocatalysts for the production of pharmaceuticals and high-value chemicals

M. Koffas

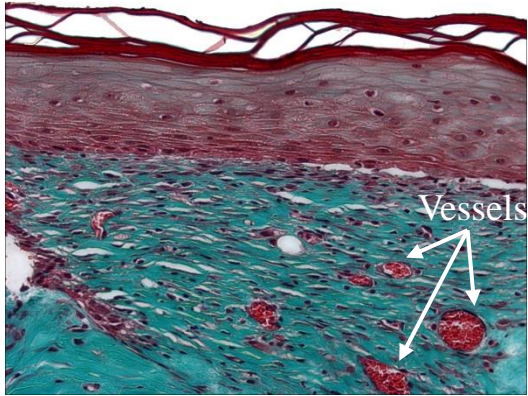
Stem Cells and Tissue Engineering

Bioengineered Skin

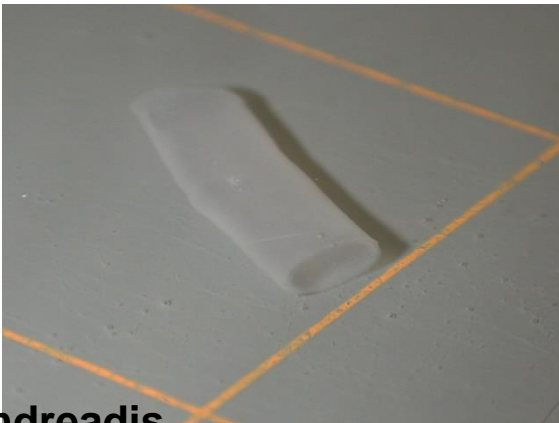


- Genetically modified tissue engineered skin:
1. Wound healing e.g. burns, chronic wounds
 2. Insulin delivery for treatment of diabetes
 3. Development of a model to study tumor invasion

Bioengineered Skin Transplanted onto mouse

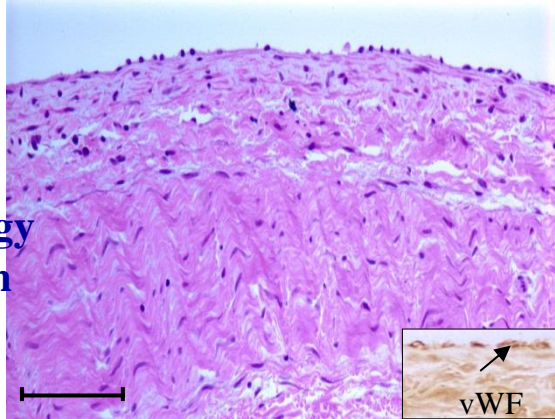


Bioengineered Blood Vessel



- Tissue engineered blood vessels (TEV):
1. Bypass surgeries
 2. Model to study mechano transduction and vascular biology
 3. TEVs from bone marrow stem cells

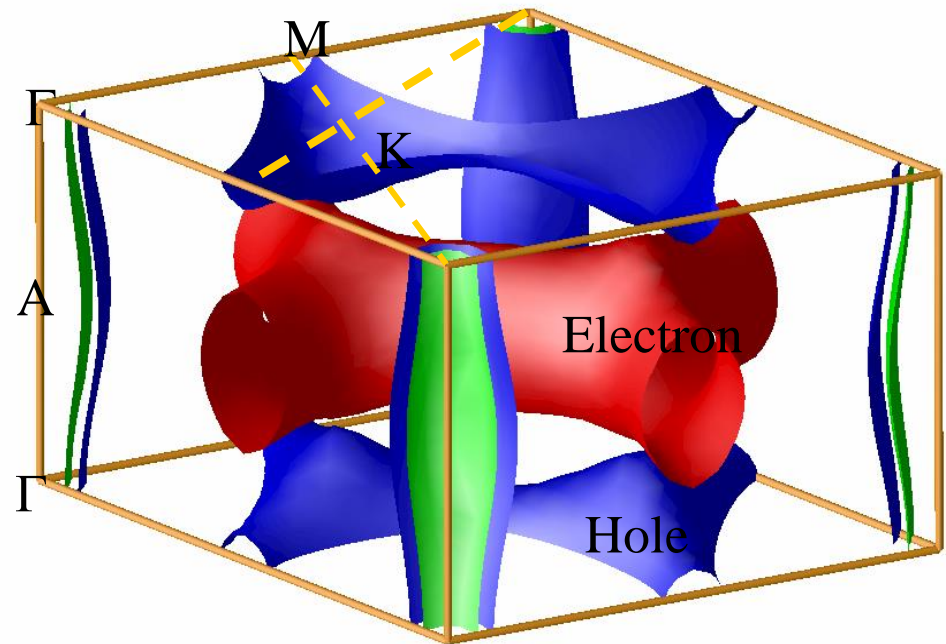
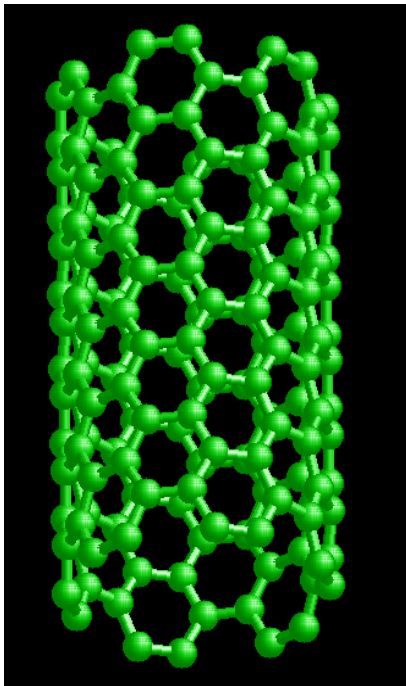
TEV Transplanted in jugular vein of lambs



S. Andreadis

Designing New Materials

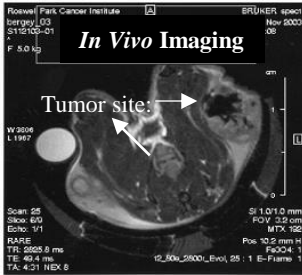
- Understand and predict materials properties
- Materials design from first-principles
- Development of new theoretical and computational techniques



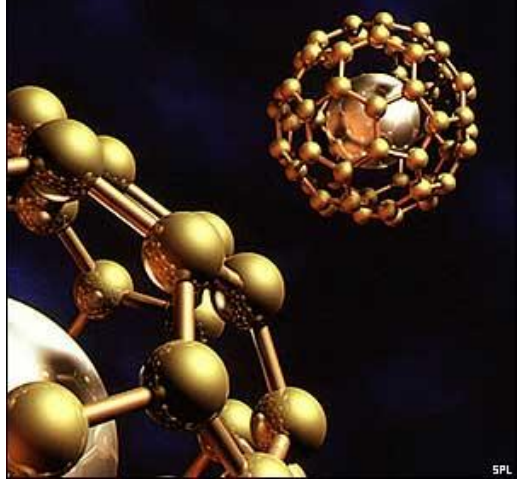
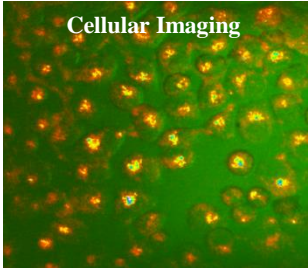
P. Zhang

Nanomedicine Program

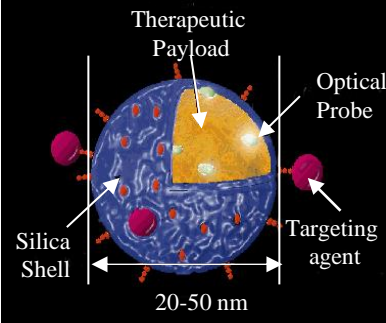
World class Research Program Melding
Nanotechnology with Biomedical Sciences



Multi-Modal
Imaging



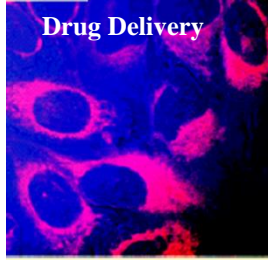
BioCompatibility/
Distribution



In Vivo
Sensing



Targeted Therapy



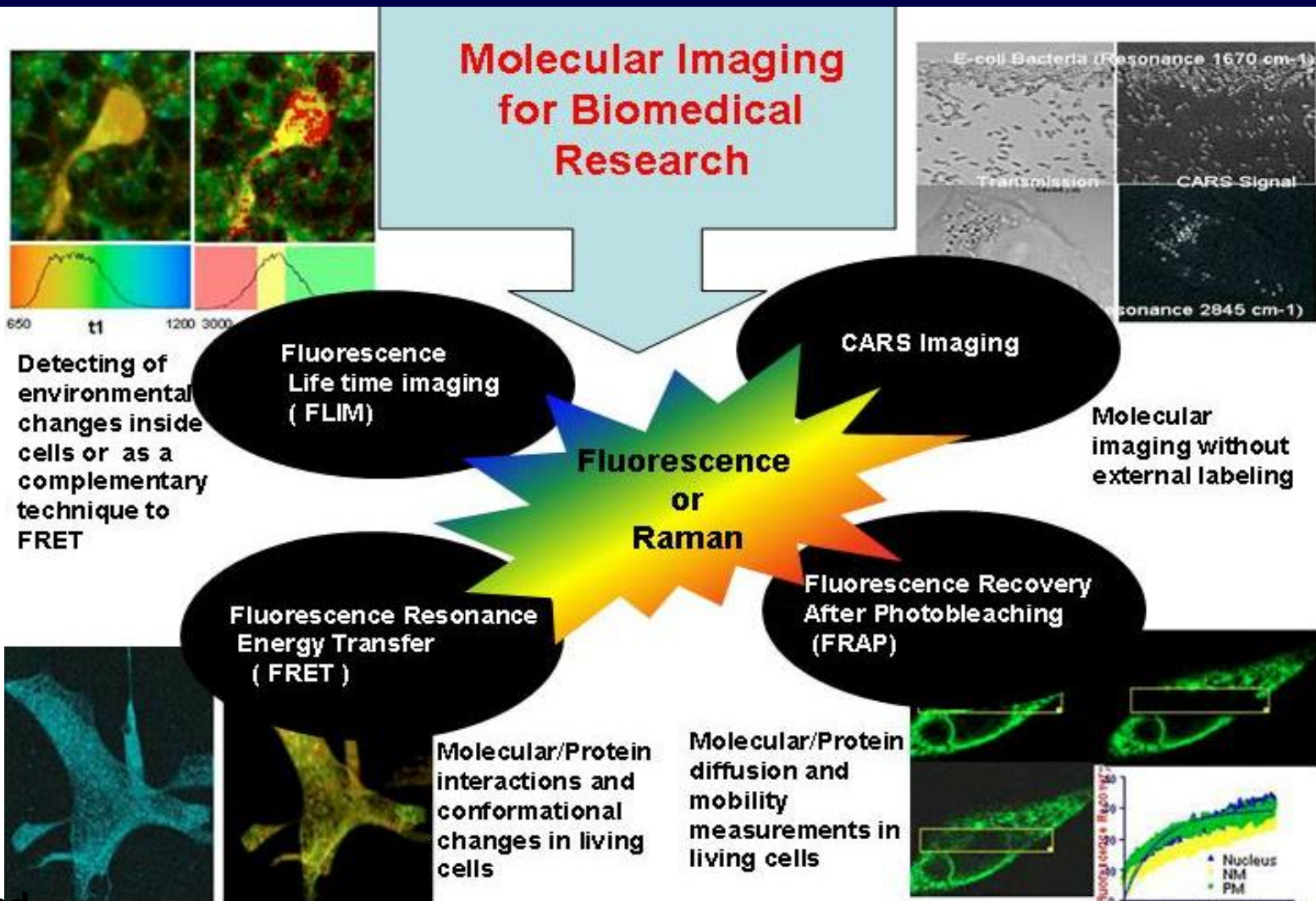
Building from the Bottom Up

State of the Art Molecular Imaging
and Nanocharacterization Facilities

- Multiphoton Laser Scanning System
- Confocal Imaging including FRET, FLIM & FRAP analysis
- Coherent Anti-Stokes Raman Imaging
- Optical Trapping/Dissection
- Advanced Laser Systems

P. Prasad
www.biophotonics.buffalo.edu

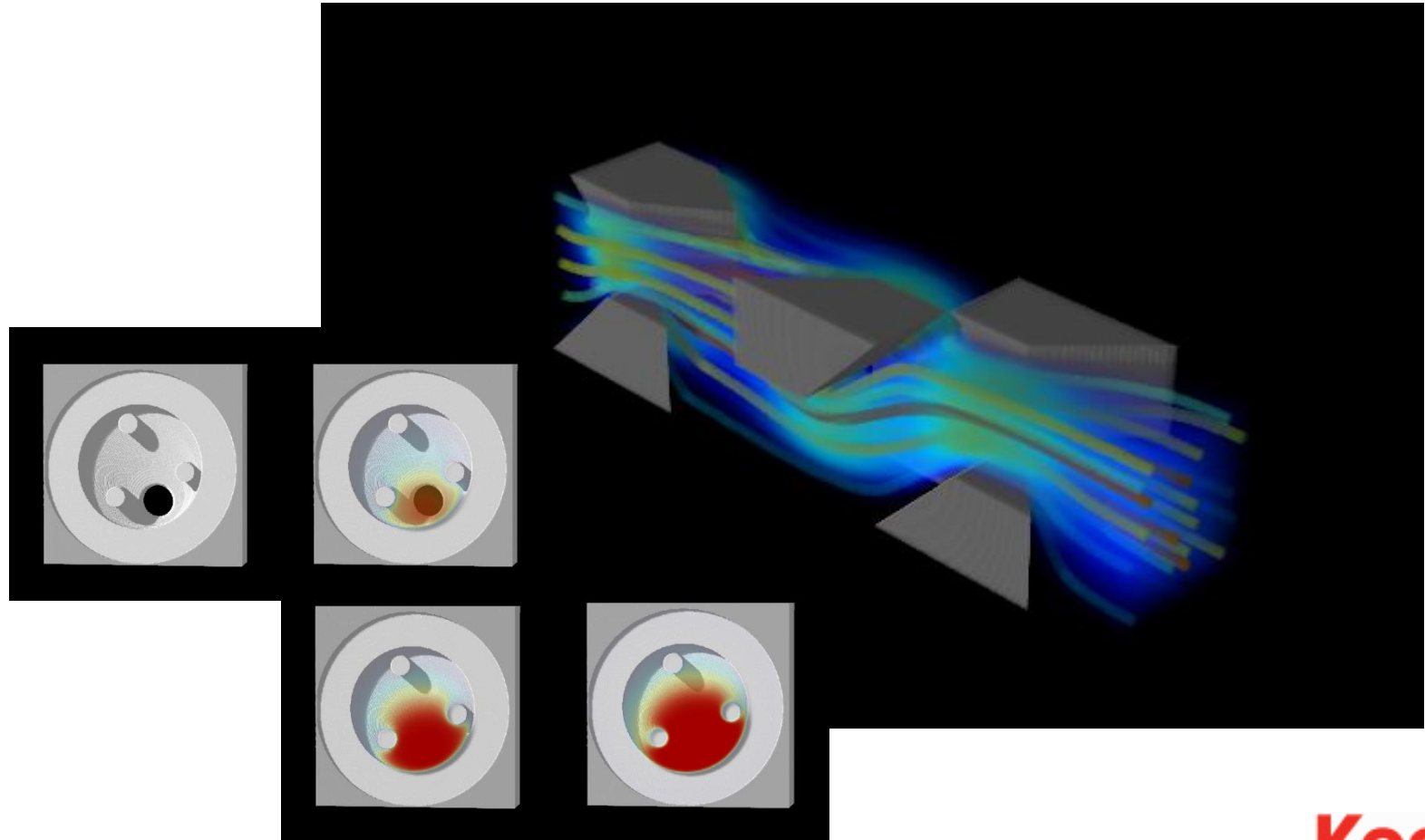
Molecular Imaging



P. Prasad

Industrial 3D Flow Analysis

- Modeling of Complex 3D and Mixing Flows for Part Analysis and Design



Shake-and-Bake

**Molecular Structure Determination
from X-Ray Crystallographic Data**

Molecular Structure Determination via *Shake-and-Bake*

■ *SnB* Software by UB/HWI

- ❑ IEEE “Top Algorithms of the Century”

■ Worldwide Utilization

■ Critical Step

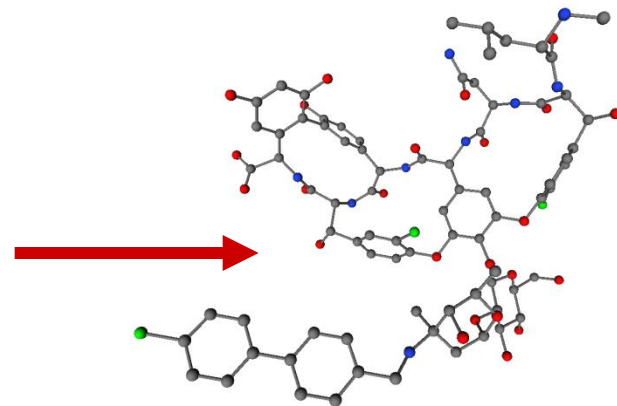
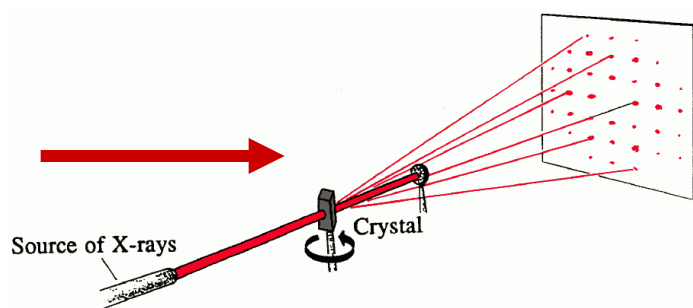
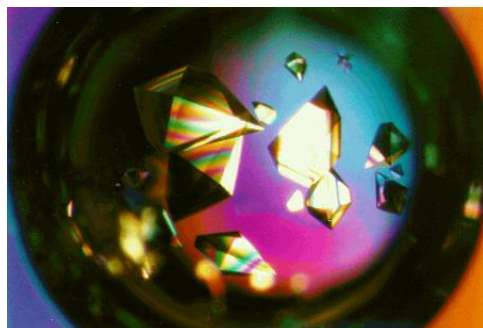
- ❑ Rational Drug Design
- ❑ Structural Biology
- ❑ Systems Biology

■ Vancomycin

- ❑ “Antibiotic of Last Resort”

■ Current Efforts

- ❑ Grid
- ❑ Collaboratory
- ❑ Intelligent Learning



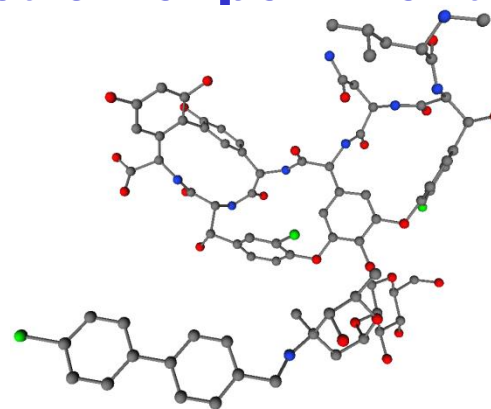
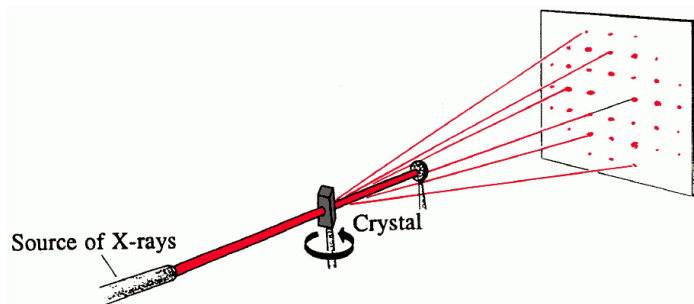
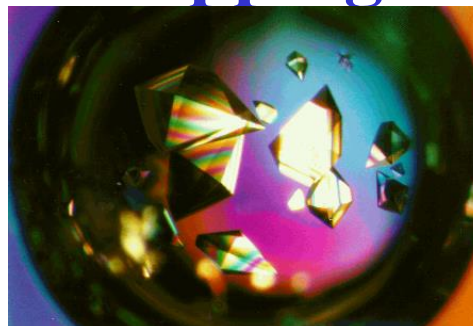
1. Isolate a single crystal
2. Perform the X-Ray diffraction experiment
3. Determine the crystal structure

X-Ray Crystallography

- **Objective:** Provide a 3-D mapping of the atoms in a crystal.

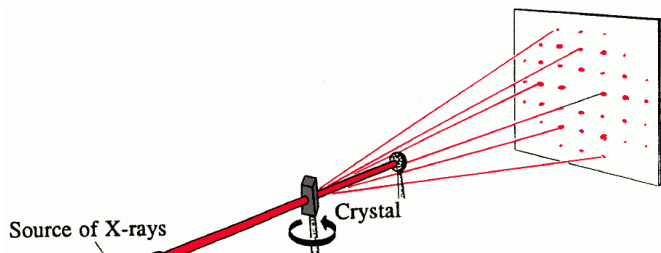
- **Procedure:**

1. Isolate a single crystal.
2. Perform the X-Ray diffraction experiment.



3. Determine molecular structure that agrees with diffraction data.

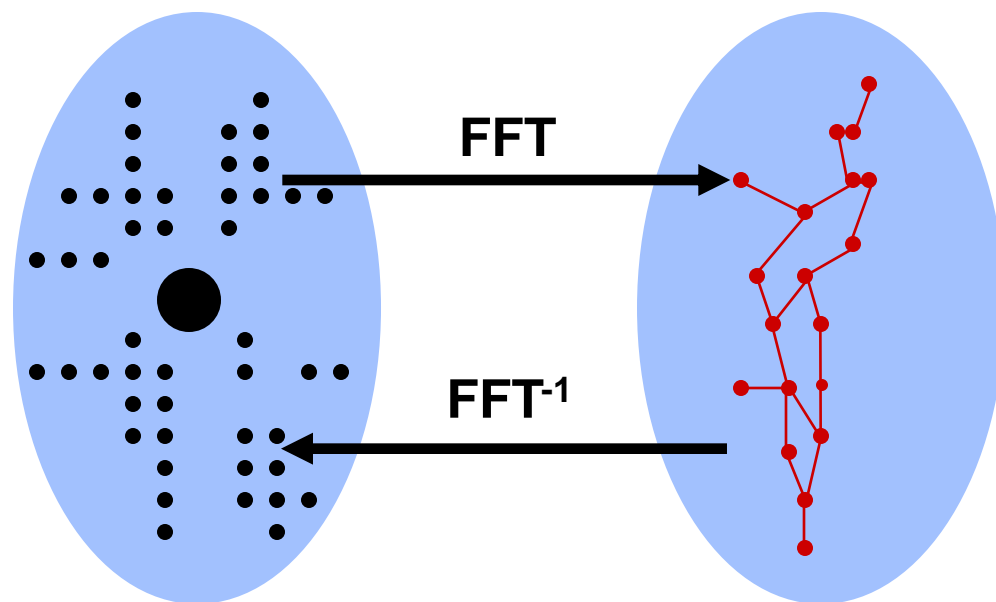
X-Ray Data & Corresponding Molecular Structure



- Experiment yields reflections and associated intensities.
- Underlying atomic arrangement is related to the reflections by a 3-D Fourier transform.
- *Phase angles are lost in experiment.*
- *Phase Problem: Determine the set of phases corresponding to the reflections.*

Reciprocal or
“Phase” Space

Real Space



X-Ray Data

Molecular Structure

Overview of Direct Methods

- Probability theory gives information about certain linear combinations of phases.
 - In particular, the triples $\phi_H + \phi_K + \phi_{-H-K} = 0$ with high probability.
- Probabilistic estimates are expressed in terms of normalized structure factor magnitudes ($|E|$).
- Optimization methods are used to extract the values of individual phases.
- A multiple trial approach is used during the optimization process.
- A suitable figure-of-merit is used to determine the trials that represent solutions.

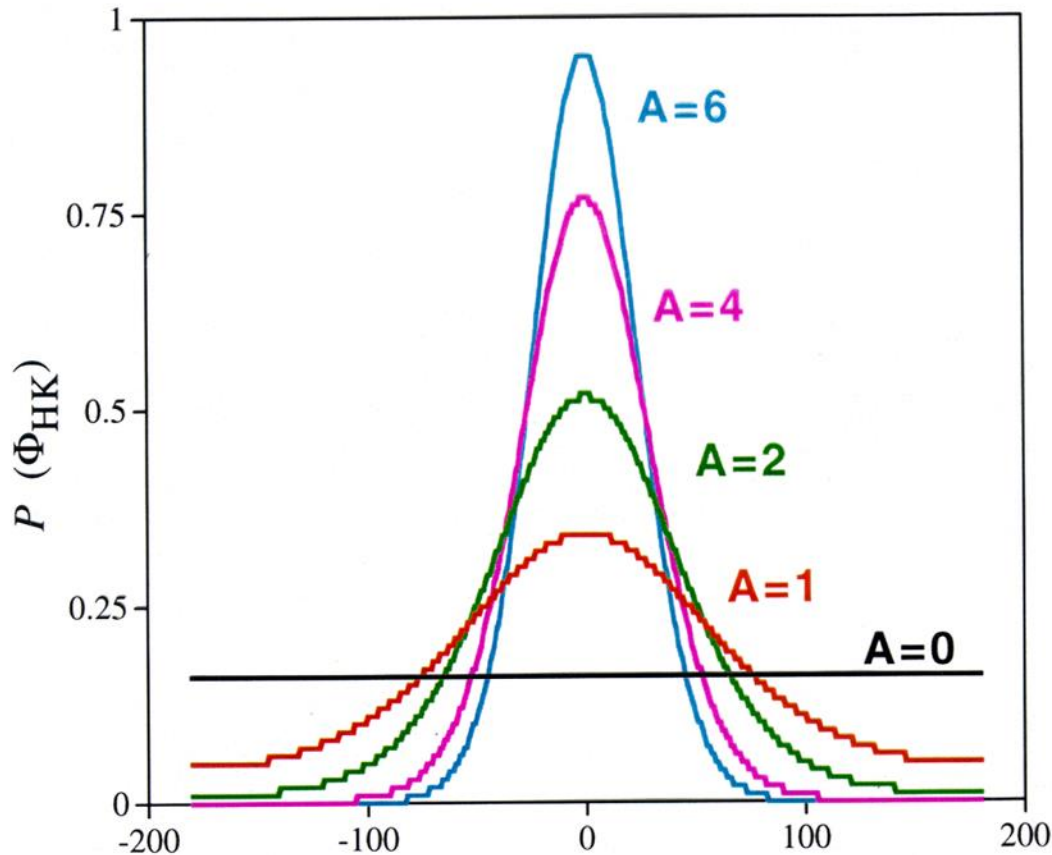
Normalized Structure-Factor Magnitudes: $|E_H|$

$$E_H = |E_H| \exp(i\phi_H)$$

$$|E_H| = \frac{|F_H|}{\langle |F_H|^2 \rangle^{1/2}} = \frac{k \langle \exp[-B_{iso} (\sin \theta)^2 / \lambda^2] \rangle^{-1} |F_H|_{meas}}{\left(\varepsilon_H \sum_{j=1}^N f_j^2 \right)^{1/2}}$$

- $\langle |E| \rangle$ constant for concentric resolution shells.
- $\langle |E| \rangle$ constant regardless of reflection class (ε_H correction factor).
- The *renormalization* condition, $\langle |E|^2 \rangle = 1$ is always imposed

Cochran Distribution



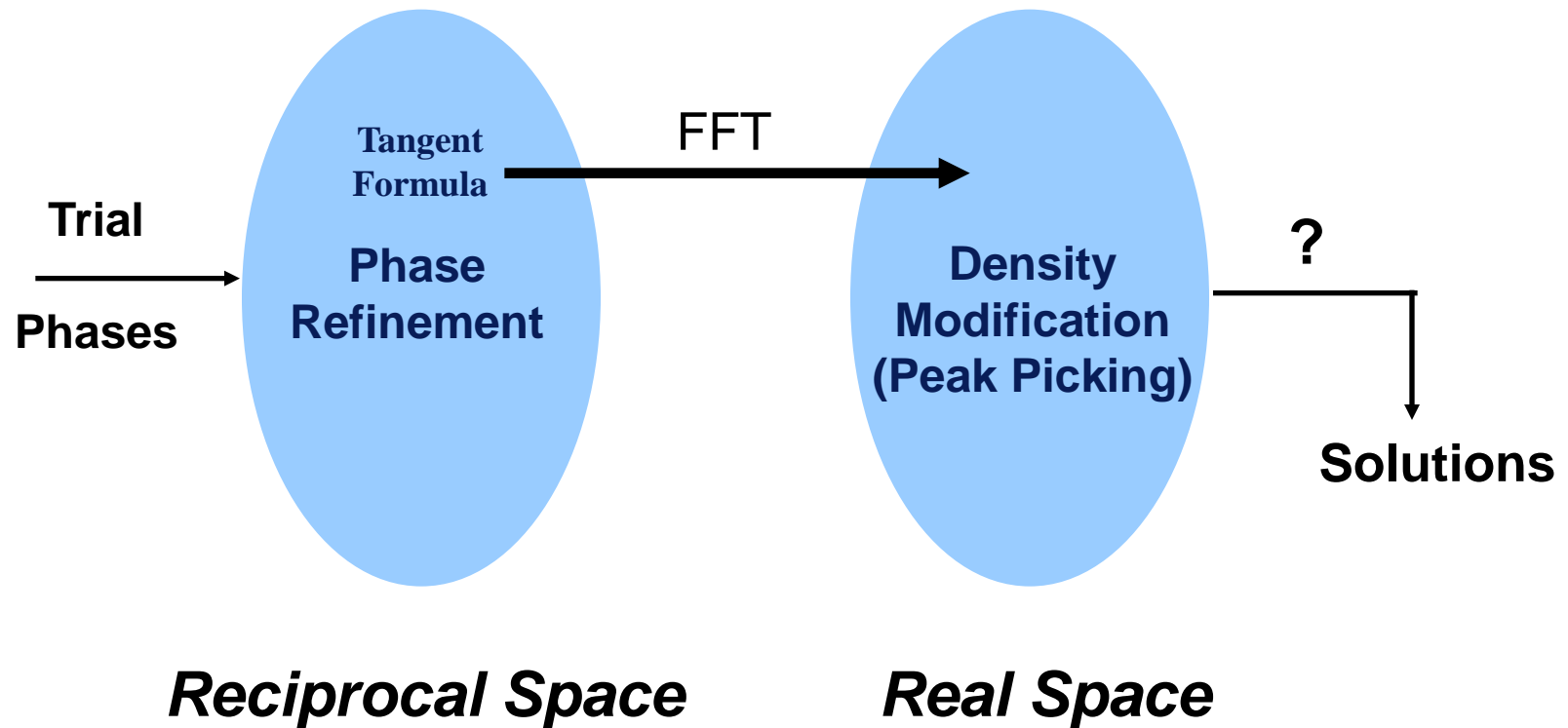
$$\Phi_{HK} = \phi_H + \phi_K + \phi_{-H-K}$$

- N = non-H atoms in unit cell
- Each triplet of phases or structure invariant, Φ_{HK} , has an associated parameter

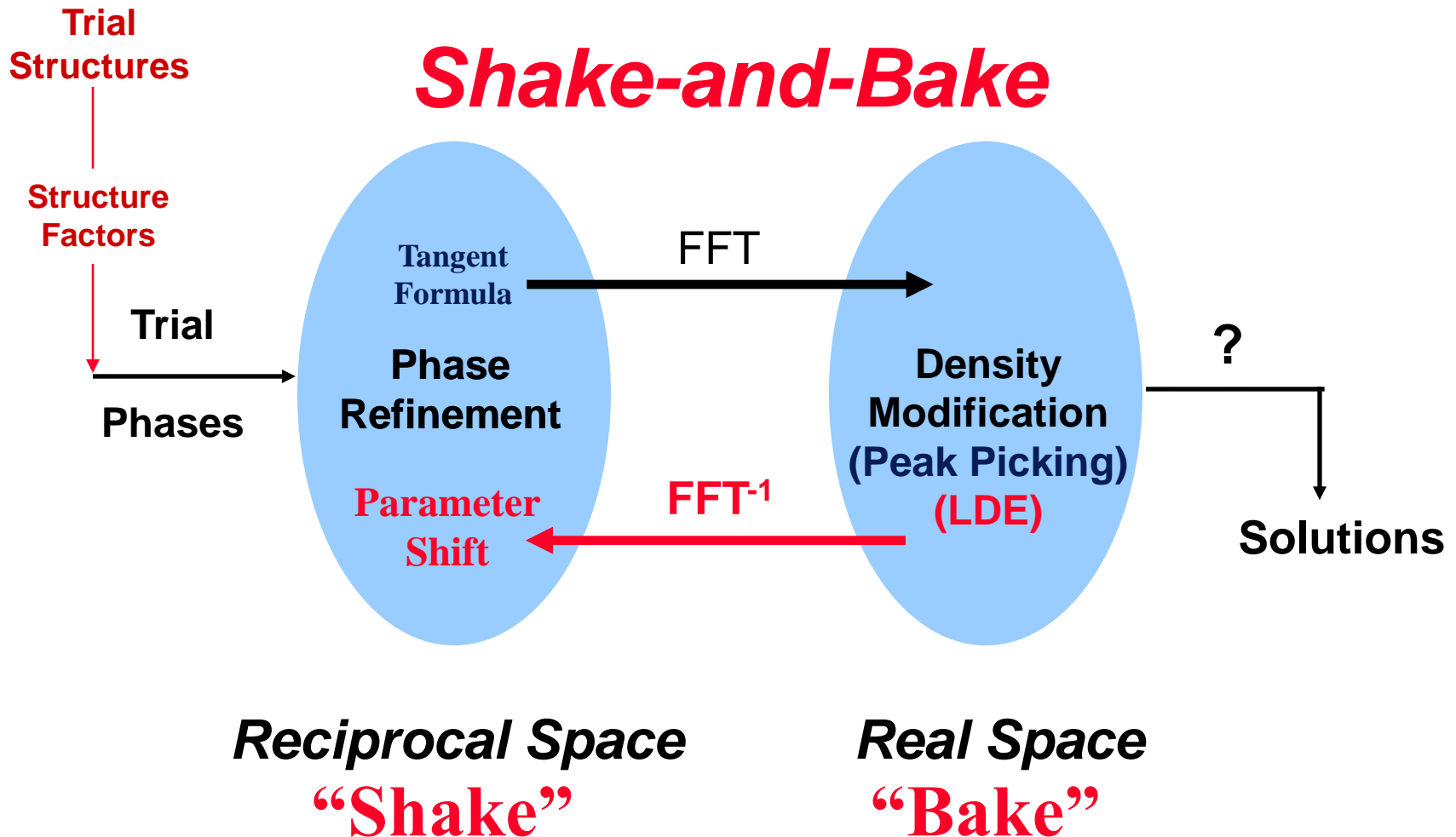
$$A_{HK} = 2|E_H E_K E_{-H-K}| / N^{1/2}$$

- A_{HK} is large if
 - $|E_H|$, $|E_K|$, $|E_{-H-K}|$ are large
 - N is small
- If A_{HK} is large, $\Phi_{HK} \approx 0$

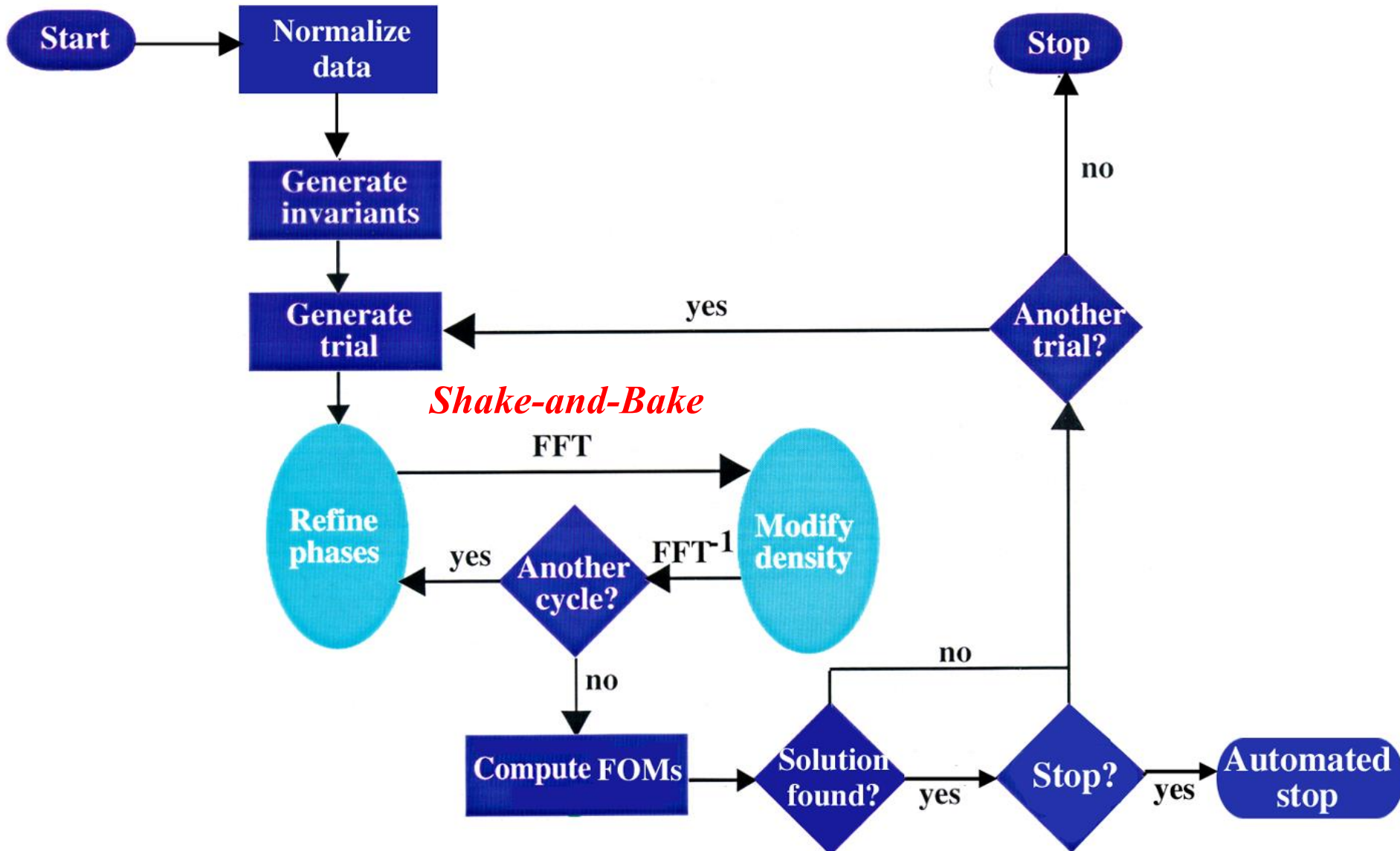
Conventional Direct Methods



Shake-and-Bake Method: Dual-Space Refinement



A Direct Methods Flowchart



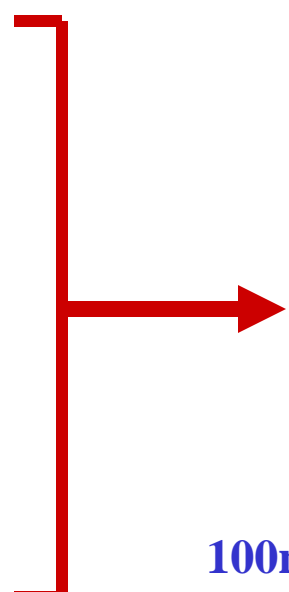
Generate Triplet Invariants

Reflections

Rank	h	k	l	E
1	0	3	4	4.65
2	0	7	30	3.67
3	5	1	1	3.67
4	8	8	5	3.26
5	6	0	1	3.15
⋮	⋮	⋮	⋮	⋮
10n=840	7	0	3	1.33

Triplets

Rank	H	K	-H-K	A
1	1	4	45	3.90
2	1	3	165	3.52
3	3	5	17	3.37
4	1	3	289	3.16
5	1	28	40	3.09
⋮	⋮	⋮	⋮	⋮
100n=840	19	259	734	0.71



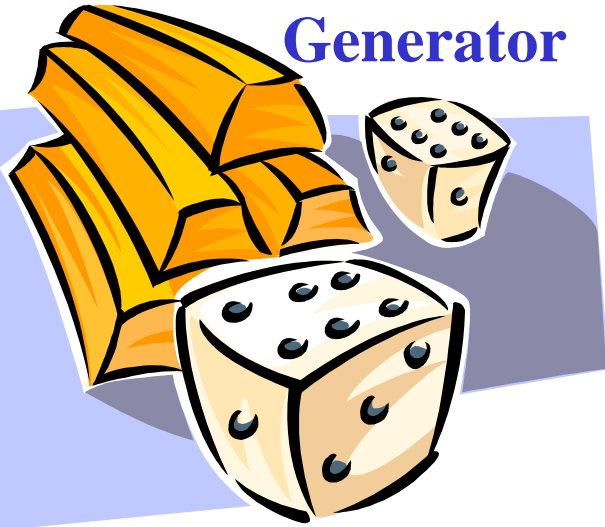
841 2 4 30 1.33

8401 142 179 283 0.71

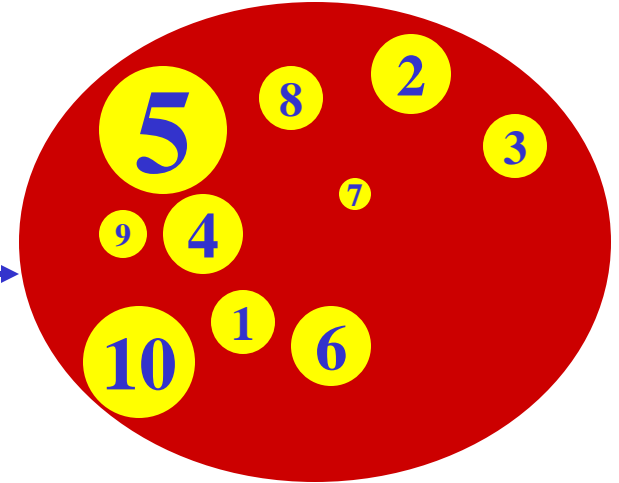
$n = 84$ unique atoms

Getting Started: Random Atoms

Random Number
Generator



$n = 10$ atoms
(30 coordinates)



Structure Factor
Calculation

ϕ_1 ϕ_2
 ϕ_3 ϕ_4
 ϕ_5 ϕ_6
 ϕ_7 ϕ_8
 ϕ_9 ϕ_{10}

Useful Relationships for Multiple Trial Phasing

Tangent
Formula

$$\tan \phi_H = \frac{-\sum_K |E_K E_{-H-K}| \sin(\phi_K + \phi_{-H-K})}{\sum_K |E_K E_{-H-K}| \cos(\phi_K + \phi_{-H-K})}$$

Parameter Shift
Optimization

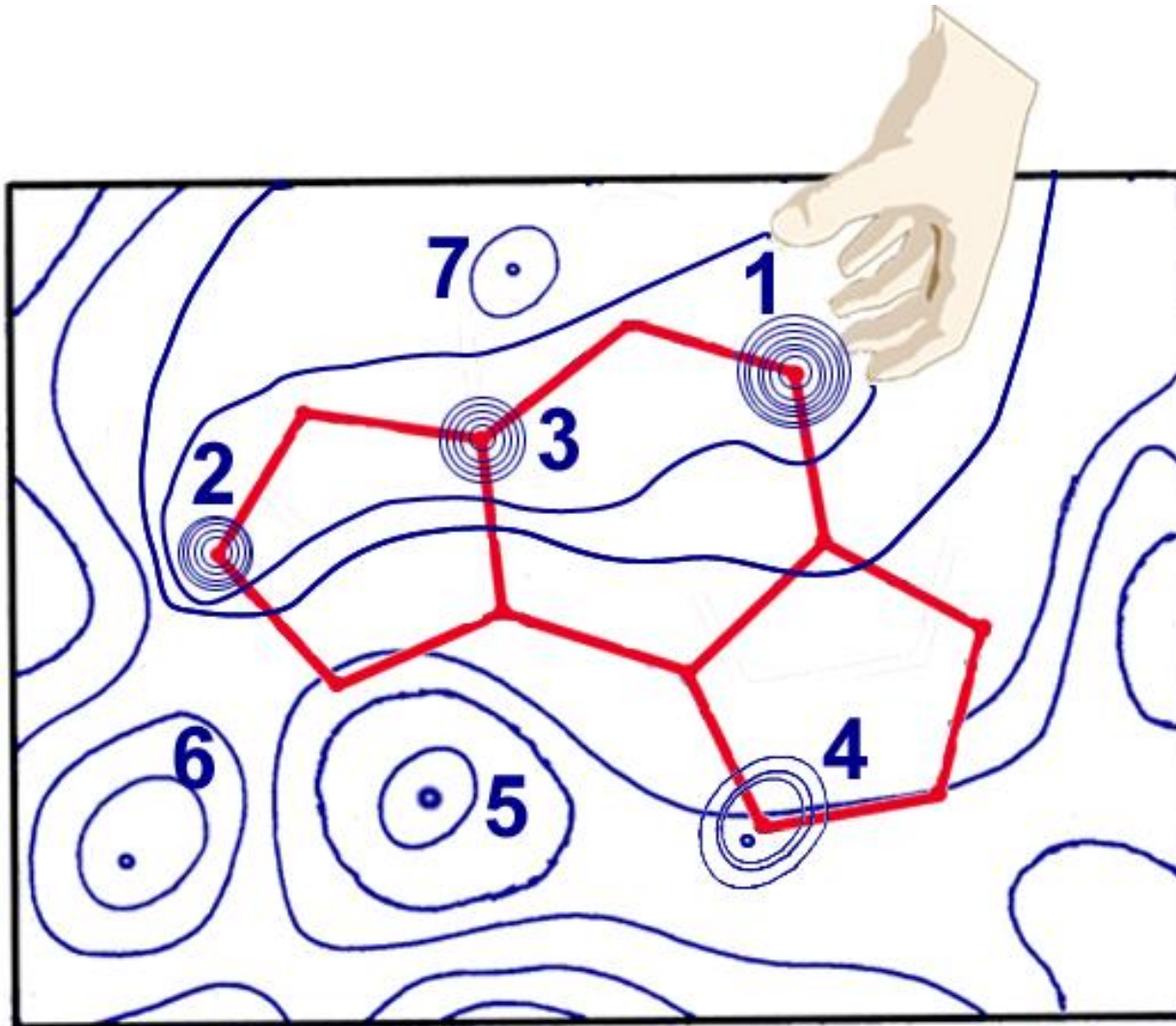
$$R(\phi) = \frac{1}{\sum_{H,K} W_{HK}} \sum_{H,K} W_{HK} \left(\cos \Phi_{HK} - \frac{I_1(W_{HK})}{I_0(W_{HK})} \right)^2$$

where $|E_H| \propto |F_H|$ normalized in resolution shells

Invariants: $\Phi_{HK} = \phi_H + \phi_K + \phi_{-H-K} \approx 0$

Weights: $W_{HK} = A_{HK} = 2N^{-1/2} |E_H E_K E_{-H-K}|$

Peak Picking




Sorted Trials

<i>Sorted Trial Data</i>							
Trial	Cycle	Refl Phased	Rmin	R Cryst.	CC	R Ratio	Peak Ratio
97	56	836	0.349	0.27	0.45	0.05	1.2
51	56	836	0.350	0.26	0.43	0.03	1.1
82	56	836	0.350	0.26	0.44	0.03	1.1
30	56	836	0.351	0.26	0.45	0.03	1.0
56	56	836	0.351	0.27	0.48	0.03	1.1

93	56	836	0.506	0.36	0.36	0.08	1.0
81	56	836	0.515	0.38	0.37	0.18	2.3
69	56	836	0.522	0.37	0.39	0.21	2.6
63	56	836	0.523	0.37	0.39	0.21	2.5
16	56	836	0.525	0.39	0.43	0.21	2.7

Solutions



Nonsolutions

Scoring Trial Structures: *SnB* FOMs

1. The minimal function ($R(\Phi)$ or R_{min})

$$2. R_{cryst} = \sum ||E_o| - k|E_c|| / \sum |E_o|$$

where the scale factor $k = \sum |E_o| / \sum |E_c|$

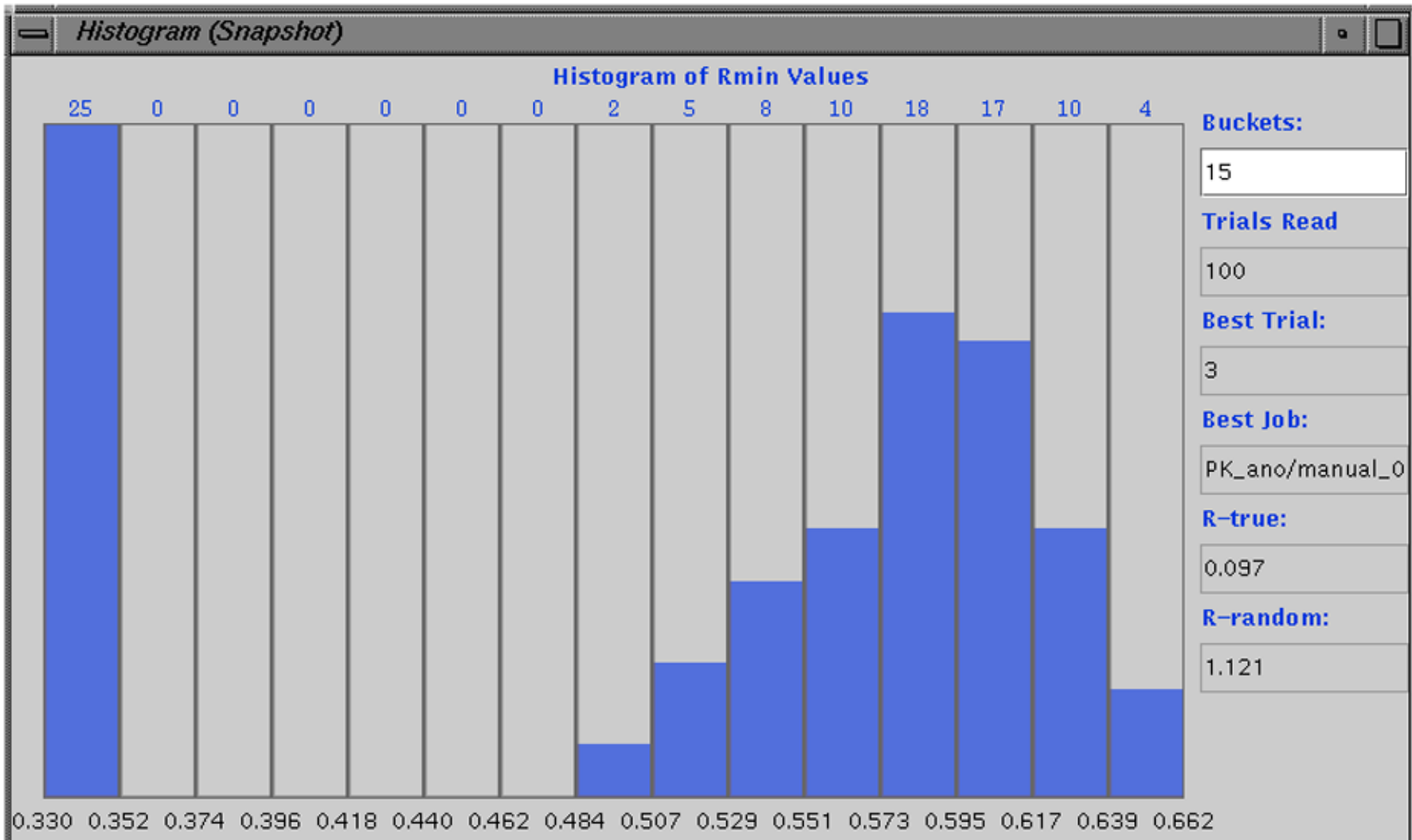
3. Correlation Coefficient (CC)

$$CC = [\sum wE_o^2 E_c^2 \sum w - \sum wE_o^2 \sum wE_c^2] /$$

$$\{[\sum wE_o^4 \sum w - (\sum wE_o^2)^2][\sum wE_c^4 \sum w - (\sum wE_c^2)^2]\}^{1/2}$$

where weights $w = 1/[0.04 + \sigma^2(E_o)]$

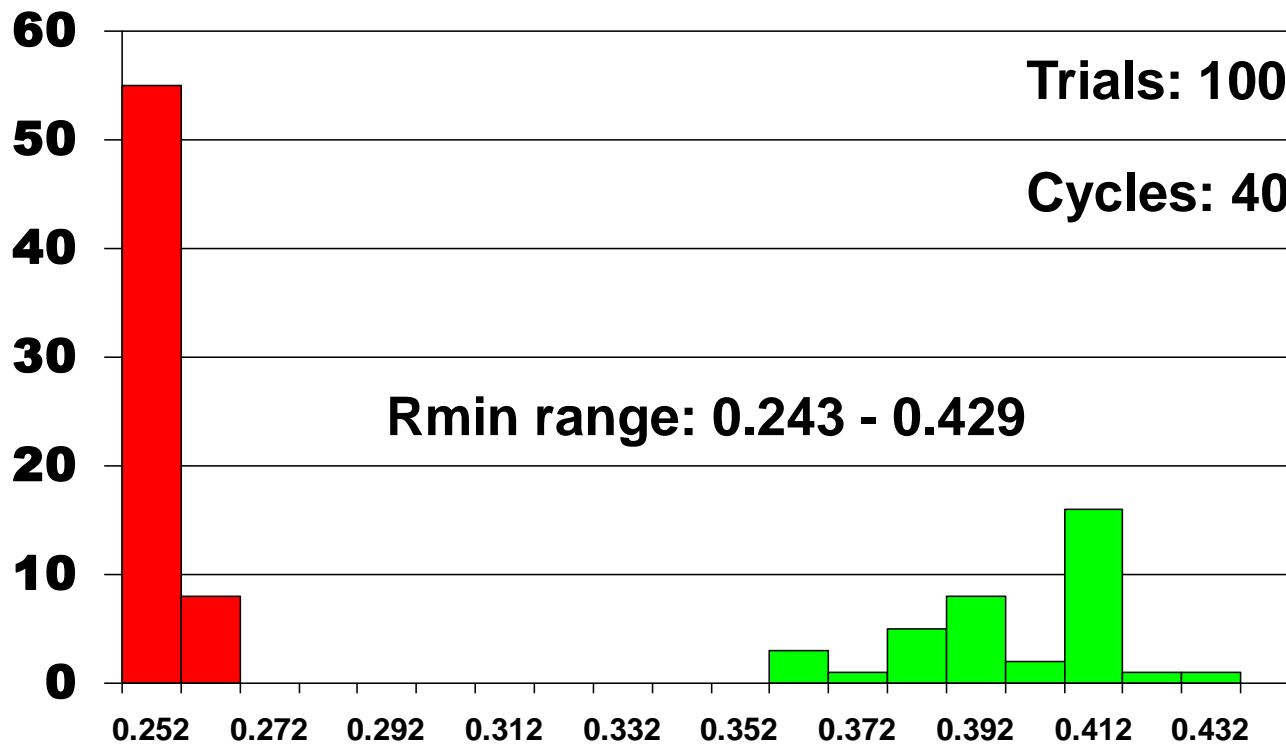
Ph8755: *SnB* Histogram



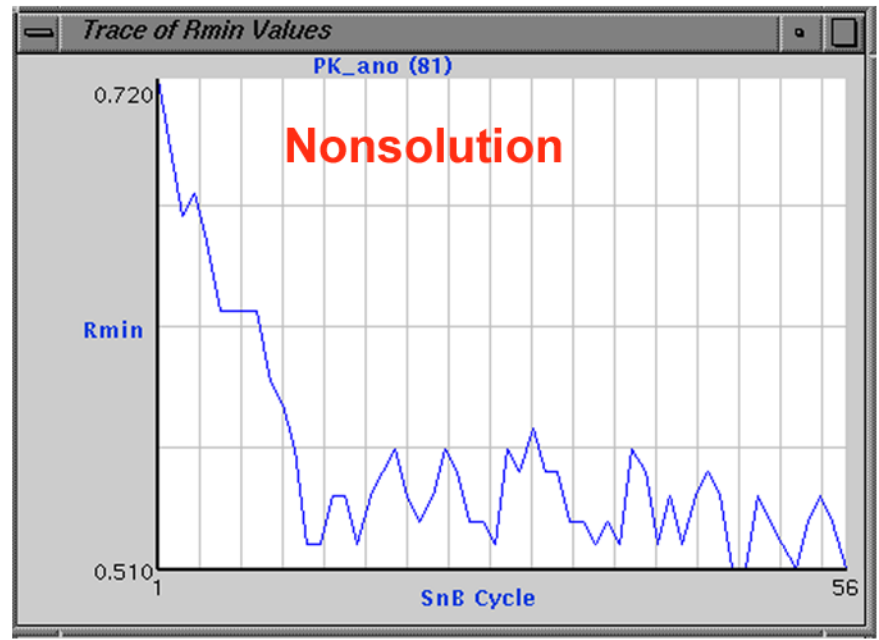
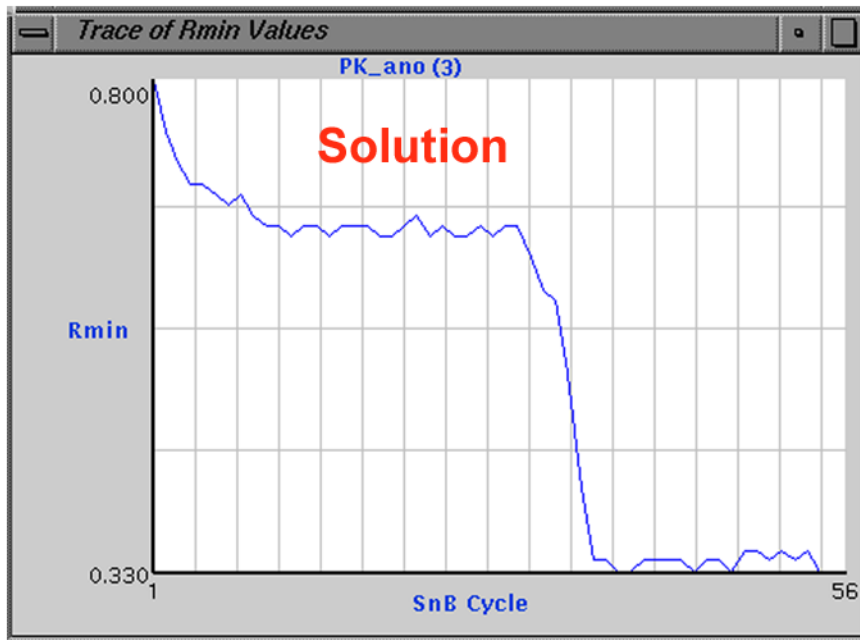
Ph8755: *SnB* Histogram

Atoms: 74
Space Group: P1

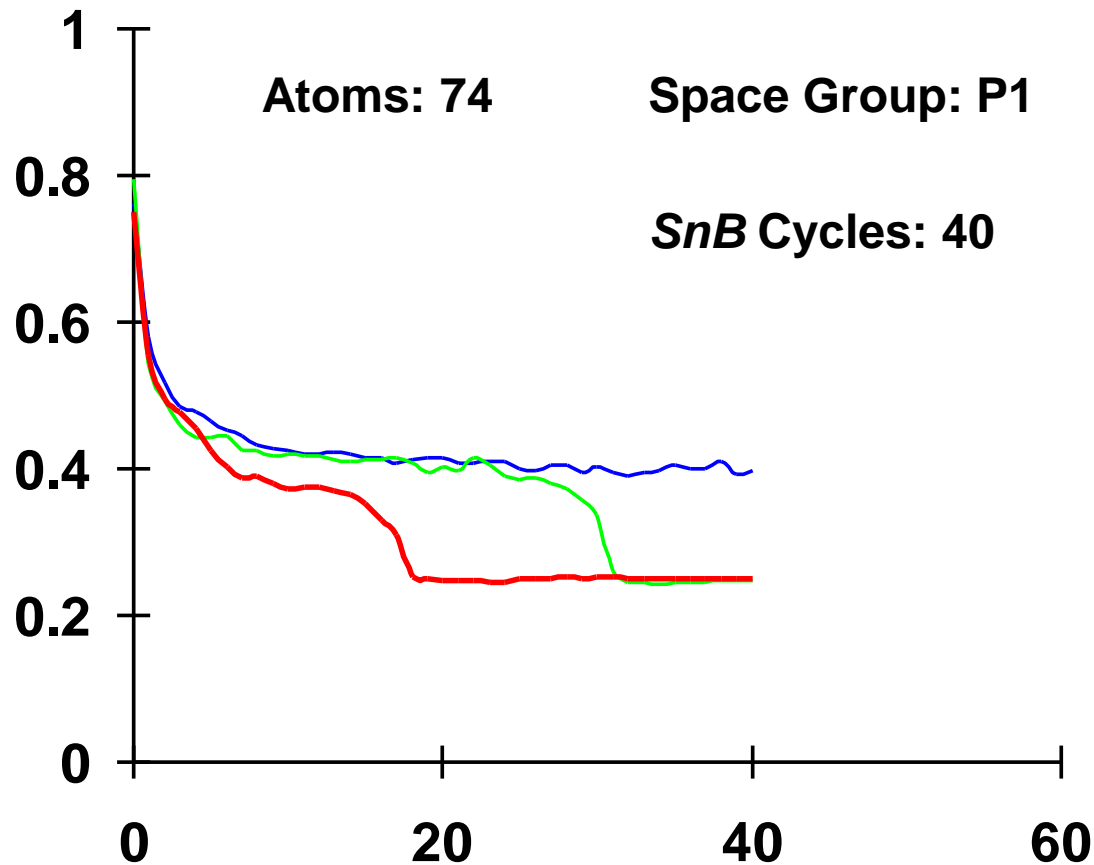
Phases: 740
Triples: 7,400



Minimal Function Traces



Ph8755: Trace of *SnB* Solution



Default *SnB* Parameters (given n atoms)

<u>Parameter</u>	<u>Full Structures</u>	<u>Substructures</u>
Phases	$10n$	$30n$
Triplet Invariants	$100n$	$300n$
Cycles		
$n < 100$	$n/2$	$2n$
$n > 100$	n	$2n$
Peaks		
$n < 100$	n	n
$n > 100$	$0.8n$	$0.8n$

Phasing and Structure Size

Se-Met with *Shake-and-Bake*



Se-Met



567 kDa (160 Se)

Multiple Isomorphous Replacement



Shake-and-Bake



Conventional Direct Methods



Vancomycin



Number of Atoms in Structure

Shake-and-Bake Applications: Structure Size and Data Resolution

■ **Basic Data (Full Structure)**

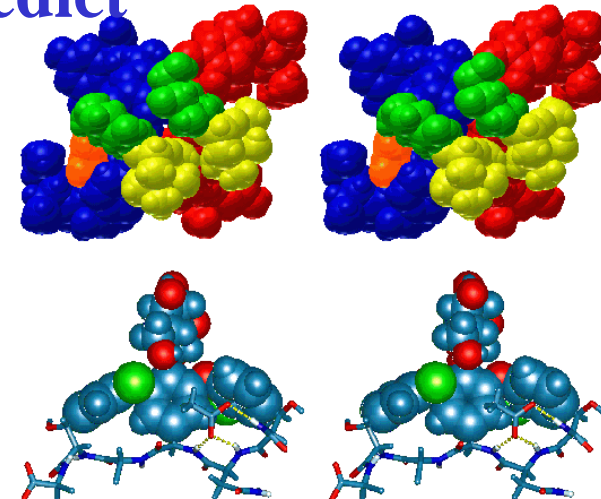
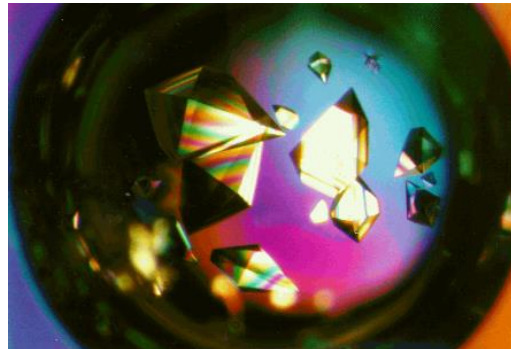
- ~750 unique non-H atoms (equal)
- ~2000 such atoms including 8 Fe's
- 1.1-1.2Å data (equal atom)
- 1.3-1.4Å data (unequal atoms, sometimes)

■ **SAS or SIR Difference Data (substructures)**

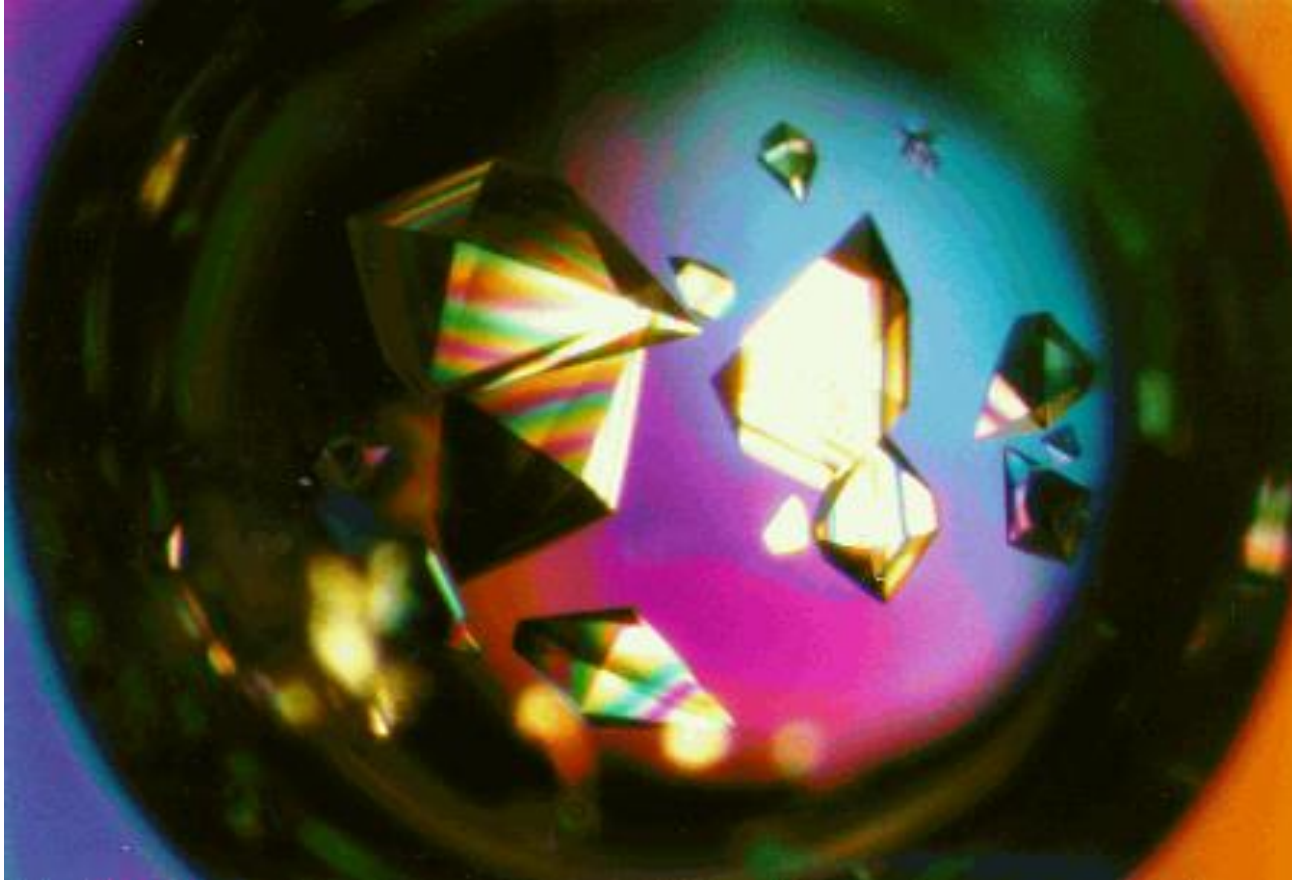
- 160 Se (567 kDa / ASU)
- 3-4Å data
- 5Å truncated data have also worked

Vancomycin

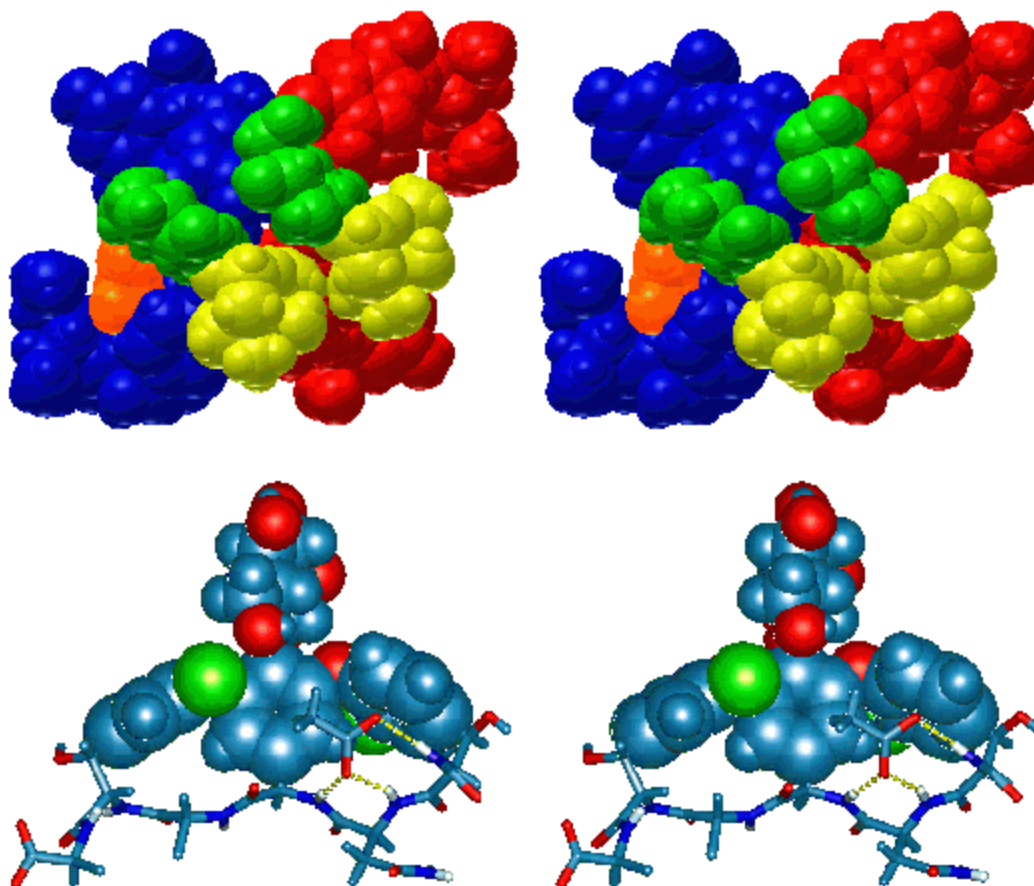
- Interferes with formation of bacterial walls
- *Last line of defense* against deadly
 - streptococcal and staphylococcal bacteria strains
- Vancomycin resistance exists (Michigan)
- Can't just synthesize variants and test
- Need structure-based approach to predict
- Solution with *SnB (Shake-and-Bake)*
 - Pat Loll
 - George Sheldrick



Vancomycin Crystal (courtesy of P. Loll)



Vancomycin Crystal Structure Views (courtesy of P. Loll & P. Axelsen)



Grid Computing

iVD gL

NEESgrid (NSF logo)

Data GRID

GLOBAL GF

GriPhyN
Data Intensive Science

European GRID Forum

TERA GRID
SDSC/UCSD • NCSA/UIUC • Caltech • ANL
NSF PACI

EURO GRID

United States virtual observatory

APAN
Asia-Pacific Advanced Network

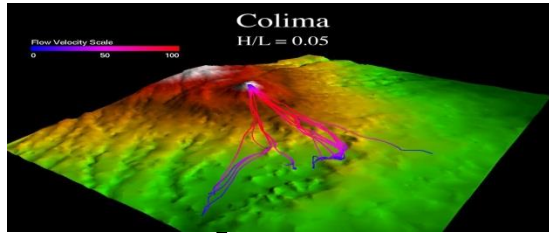
PDB
PROTEIN DATA BANK

Open Science Grid

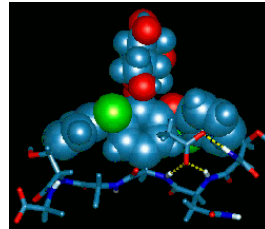
GRID

Center for Computational Research Data Center

Grid Computing Overview



Data Acquisition



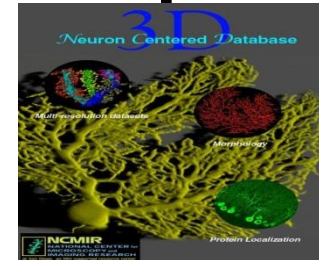
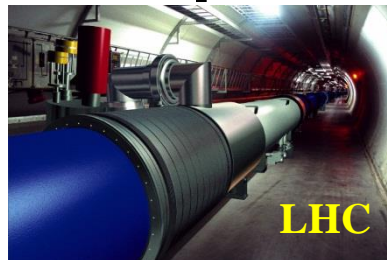
Advanced Visualization



Analysis



Imaging Instruments



Large-Scale Databases

- Coordinate Computing Resources, People, Instruments in Dynamic Geographically-Distributed Multi-Institutional Environment
- Treat Computing Resources like Commodities
 - ❑ Compute cycles, data storage, instruments
 - ❑ Human communication environments
- No Central Control; No Trust

“Middleware”

- **Intermediate Software Layer between Application Codes and Grid Resources**
- **Required for applications, users, and resource providers to operate effectively in a manner transparent to the user**
- **Security; Resource Management; Data Access; Policies; Accounting;**
- **Globus; Condor**
- **Checks availability of Resources**
 - CPUs; Storage; Networking; Render Farms; etc.
- **Scheduling / Workload Management System**
- **Resource Broker**
 - Evaluates Job and Breaks Up/Submits

NSF Middleware Initiative (NMI)

- Develop, improve, and deploy a suite of reusable software components for use in national-scale “cyberinfrastructure”.
- APST, Condor, CPM, DataCutter, DataCutter STORM, Globus Toolkit, GPT, Gridconfig, GridPort, GridSolve, GSI OpenSSH, Inca, KX.509/KCA, Look, MPICH-G2, MyProxy, Network Weather Service, OpenSAML, PERMIS, PyGlobus, Shibboleth, SRB Client, UberFTP, and WebISO (Web Initial Sign-on).

Grid Issues

- **High-Throughput Computing**
- **Transparent Integration of Data, Computing, Sensors/Devices, Networking**
- **Heterogeneous Resources**
- **Standards (Grid, Data)**
- **Major User Communities**
 - **High-Energy Physics and Astrophysics**
 - **Medicine and Biological Sciences**
 - **Earth Sciences**
- **Public Funding Still Critical**
- **Grids are in their Infancy**

Major Grid Initiatives

- **TeraGrid (NSF)**
 - ❑ Integrates High-End Resources
 - ❑ High-Performance (Dedicated) Networks
 - ❑ 9 Sites (?); 250TF & 30PB (?)
 - ❑ 100+ Databases Available
- **OSG (DOE, NSF)**
 - ❑ High-Throughput Distributed Facility
 - ❑ Open & Heterogeneous
 - ❑ Biology, Computer Science, Astrophysics, LHC
 - ❑ 57 Compute Sites; 11 Storage Sites;
 - ❑ 10K CPUS; 6PB
- **EGEE: Enabling Grids for E-Science (European Commission)**
 - ❑ Initial Focus on CERN (5PB of Data/Year)
 - High-Energy Physics and Life Sciences
 - ❑ Expanded Focus Includes Virtually All Scientific Domains
 - ❑ 200 Institutions; 40 Countries
 - ❑ 20K+ CPUs; 5PB; 25,000 jobs per day!

Open Science Grid

Courtesy of Paul Avery

Open Science Grid *Applications, Infrastructure, and Facilities*

Applications

BaBar,
STAR, PHENIX
etc

Biology

Computer
Science

Astrophysics

Run 2
CDF, D0

LHC
Atlas, CMS
Alice

Persistent Grid
Infrastructure

User Support
Center

Middleware
Providers

Certificate
Authorities

Service
Providers

Grid Operations
Center

Database
Operators

Facilities

General Facility
for any
Community e.g.
TeraGrid

Laboratory
Serving Multiple
Communities
e.g. Fermilab,
BNL, NERSC

Community
Facility
e.g. US ATLAS
or CMS
Tier-1/Tier-2

University
Facility e.g.
UFlorida,
Buffalo

University
Community
Facility e.g.
GLOW

Cyberinfrastructure

- **Foster & Kesselman:** “a domain-independent computational infrastructure designed to support science.”
- **NSF:** “comprehensive phenomenon that involves creation, dissemination, preservation, and application of knowledge”
- **Generic:** transparent and ubiquitous application of technologies central to contemporary engineering and science
- **NSF Cyberinfrastructure (OCI)**
 - ❑ HPC Hardware and Software
 - ❑ Data Collections
 - ❑ Science Gateways/Virtual Organizations
 - ❑ Support of Next Generation Observing Systems

Miller's Cyberinfrastructure Lab

- CI sits at core of modern simulation & modeling
- CI allows for new methods of investigation to address previously unsolvable problems
- Focus on development of *algorithms, portals, interfaces, middleware*
- Free end-users to do disciplinary work
- Funding (2001-pres): NSF ITR, NSF CRI, NSF MRI, NYS, Fed
- Experimental Equipment (Dell/Lenovo): 1.25 TF Clusters, 140 Cores (Intel/AMD), 4 TB Internal Storage, GigE, IB, Condor Flock (35 Intel/AMD), 22 TB Storage (2)
- Production Equipment (Dell): Workstations, 15 TB Storage, CCR equipment

Evolution of CI Lab Projects

■ ACDC-Grid

- ❑ Experimental Grid: Globus & Condor
- ❑ Integrate Data & Compute, Monitor, Portal, Node Swapping, Predictive Scheduling/Resource Management
- ❑ GRASE VO: Structural Biology, Groundwater Modeling, Earthquake Eng, Comp Chemistry, GIS/BioHazards
- ❑ Buffalo, Buffalo State, Canisius, Hauptman-Woodward

■ WNY Grid

- ❑ Heterogeneous System: Hardware, Networking, Utilization
- ❑ Buffalo, Geneseo, Hauptman-Woodward, Niagara

■ NYS Grid

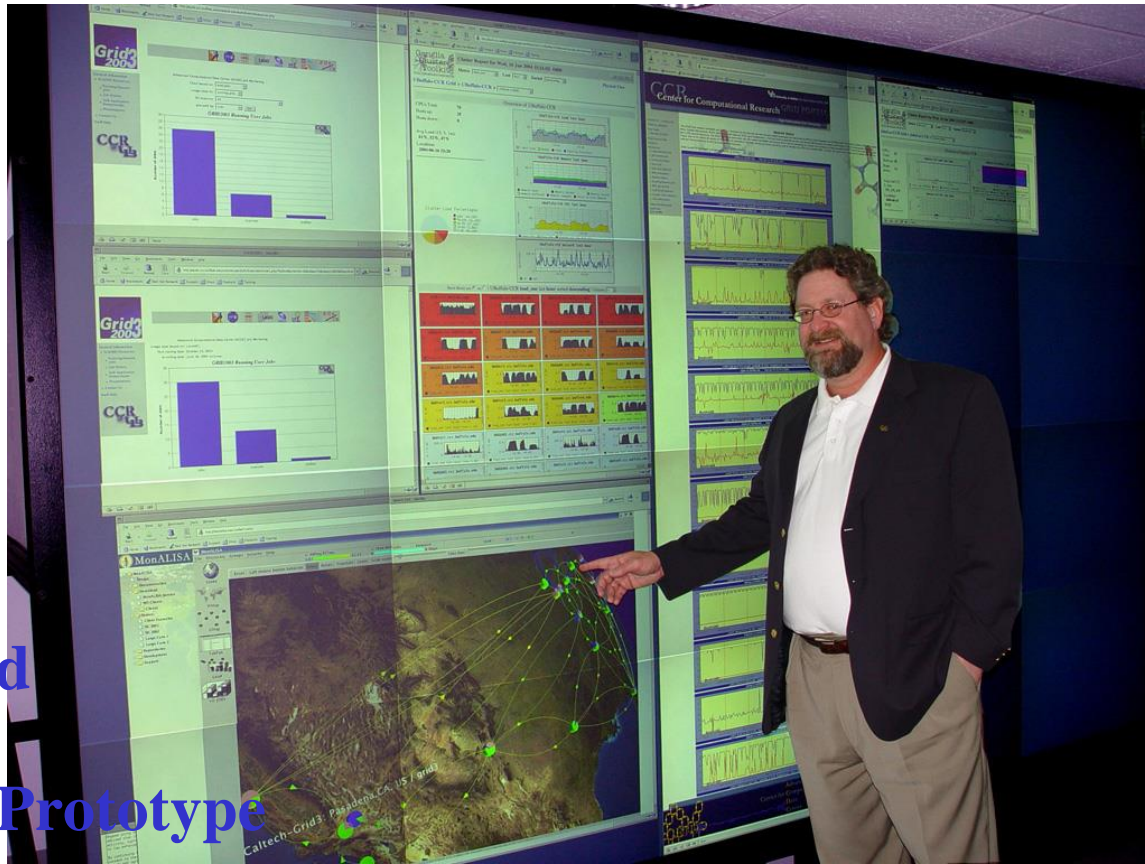
- ❑ Extension to Hardened Production-Level System State-Wide
- ❑ Albany, Binghamton, Buffalo, Geneseo, Canisius, Columbia, HWI, Niagara, [Cornell, NYU, RIT, Rochester, Syracuse, Marist], {Stony Brook, RPI, Iona}

NYS Grid Resources

- Albany: 8 Dual-Processor Xeon Nodes
- Binghamton: 15 Dual-Processor Xeon Nodes
- Buffalo: 1050 Dual-Processor Xeon Nodes
- Cornell: 30 Dual-Processor Xeon Nodes
- Geneseo State: Sun/AMD with 128 Compute Cores
- Hauptman-Woodward Institute: 50 Dual-Core G5 Nodes
- Marist: 9 P4 Nodes
- Niagara University: 64 Dual-Processor Xeon Nodes
- NYU: 58 Dual-Processor PowerPC Nodes
- RIT: 4 Dual-Processor Xeon Nodes
- Syracuse: 8 Dual-Processor Xeon Nodes

CI Lab Collaborations

- High-Performance Networking Infrastructure
- Grid3+ Collaboration
- iVDGL Member
 - Only External Member
- Open Science Grid
 - GRASE VO
- NYSGrid.org
 - NYS CI Initiative
 - Executive Director
 - Various WGs
- Grid-Lite: Campus Grid
 - HP Labs Collaboration
- Innovative Laboratory Prototype
 - Dell Collaboration



ACDC-Grid Collaborations II

■ Grass Roots NYS Grid (pre-NYSGrid.org)

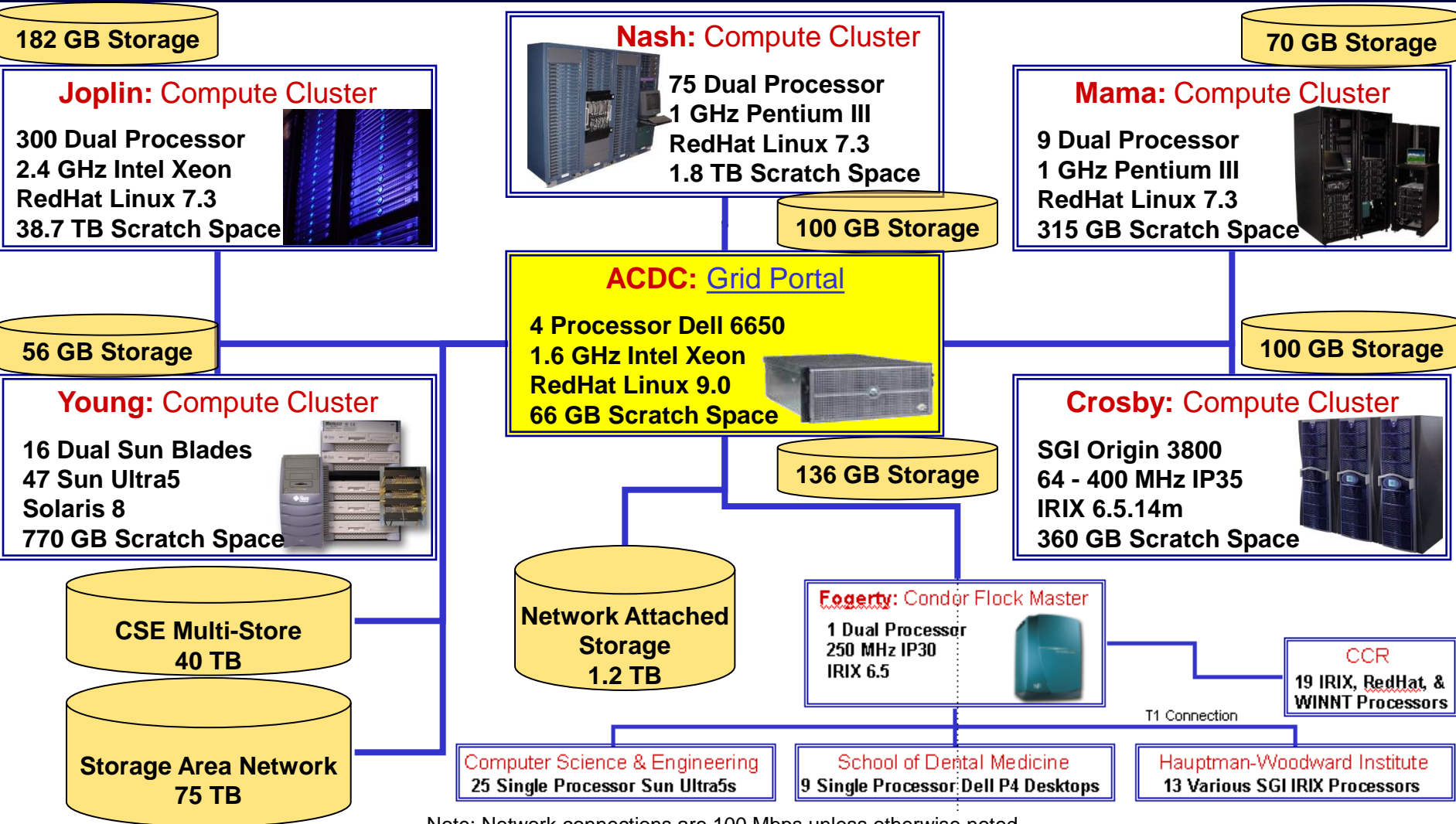
- SUNY-Albany
- SUNY-Binghamton
- SUNY-Buffalo
- SUNY-Geneseo
- Canisius College
- Columbia
- Hauptman-Woodward Inst.
- Niagara University

■ GRASE VO: Grid Resources for Advanced Science and Engineering Virtual Organization

- (Non-Physics Research)
- Structural Biology
- Groundwater Modeling
- Earthquake Engineering
- Computational Chemistry
- GIS/BioHazards

ACDC Data Grid Overview

(Grid-Available Data Repositories)

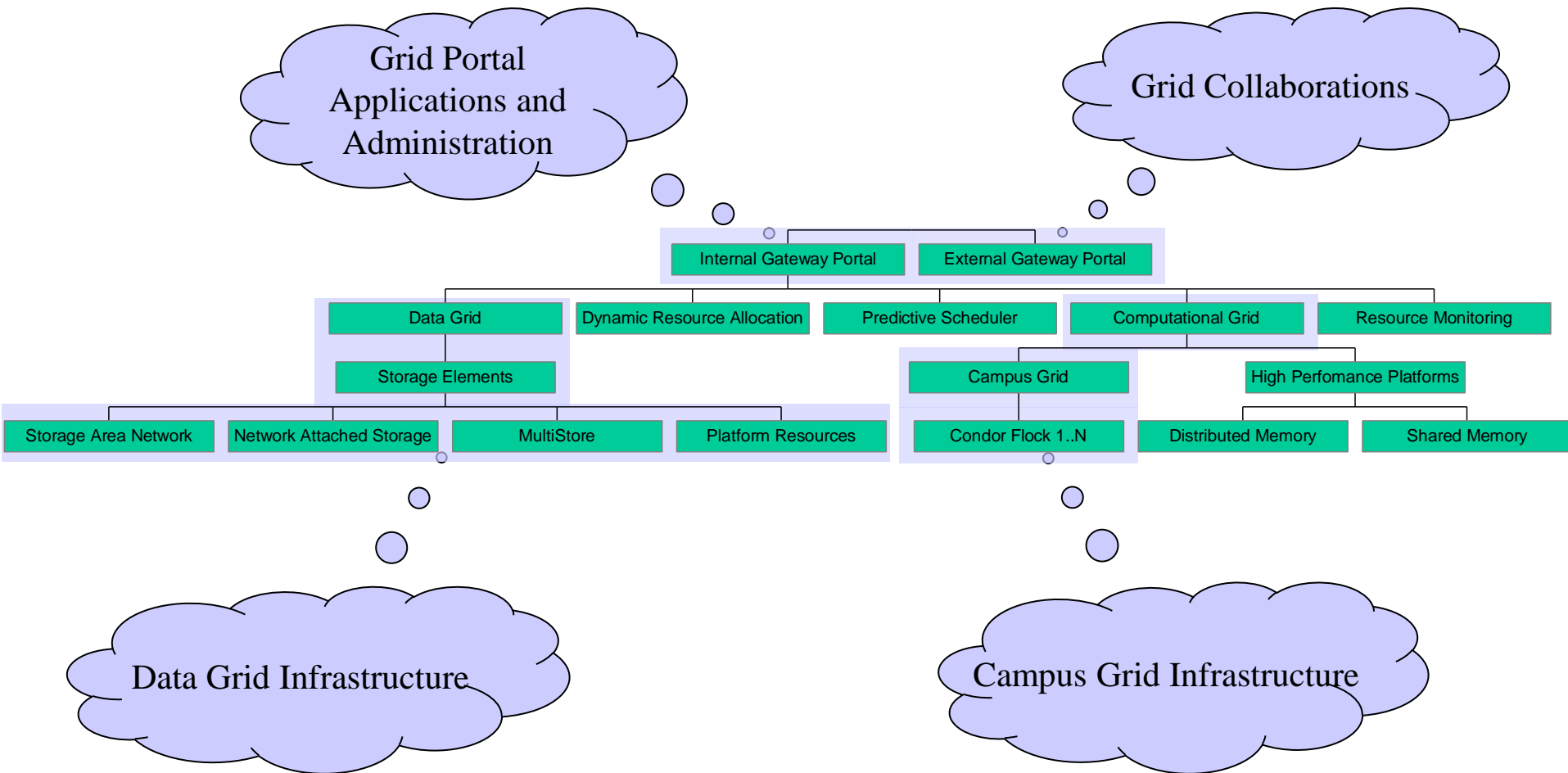


Note: Network connections are 100 Mbps unless otherwise noted.

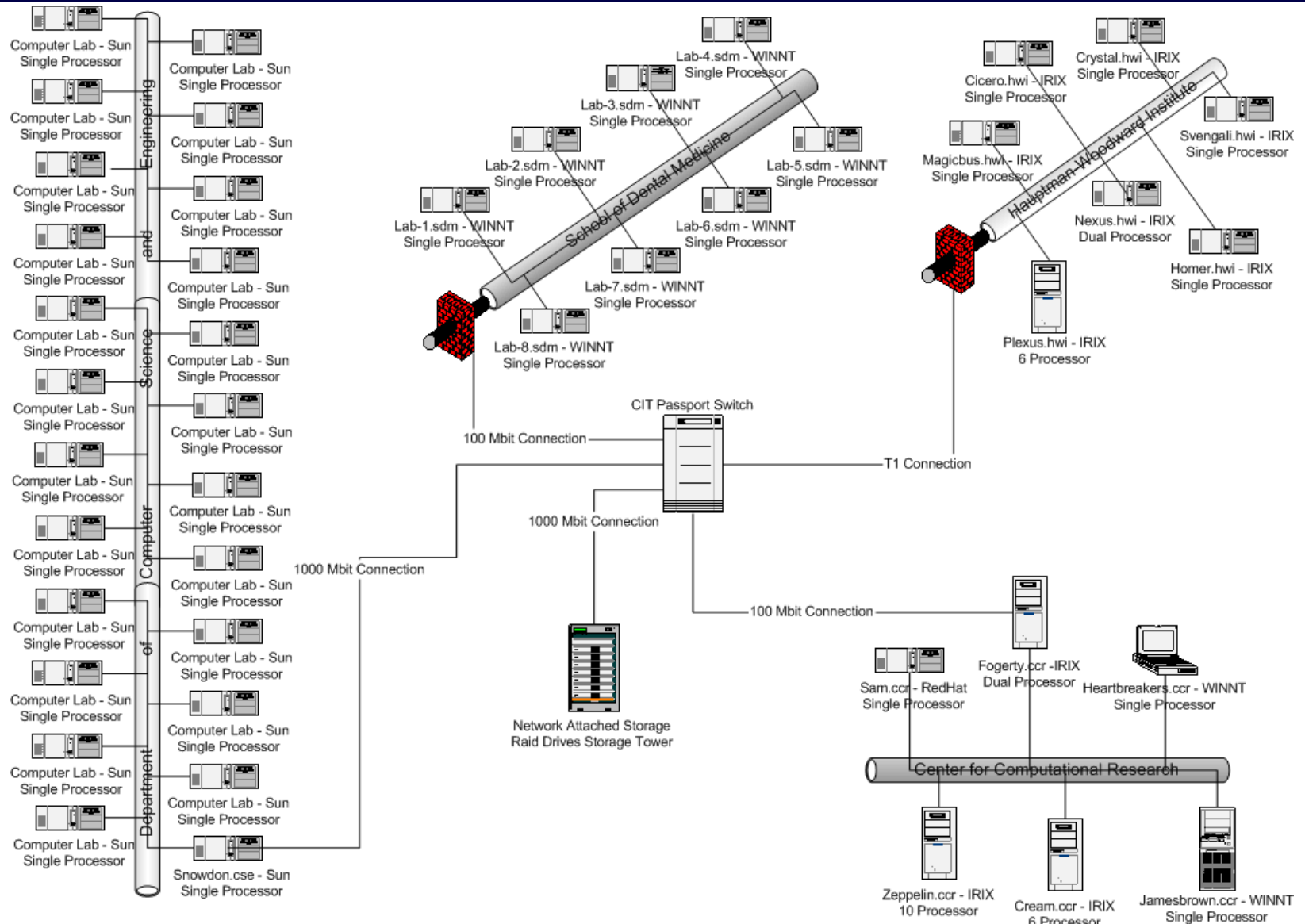
CI Lab Projects

- **Lightweight Grid Monitor (Dashboard)**
- **Predictive Scheduler**
 - Define quality of service estimates of job completion, by better estimating job runtimes by profiling users.
- **Dynamic Resource Allocation**
 - Develop automated procedures for dynamic computational resource allocation.
- **High-Performance Grid-Enabled Data Repositories**
 - Develop automated procedures for dynamic data repository creation and deletion.
- **Integrated Data Grid**
 - Automated Data File Migration based on profiling users.
- **Grid Portal**

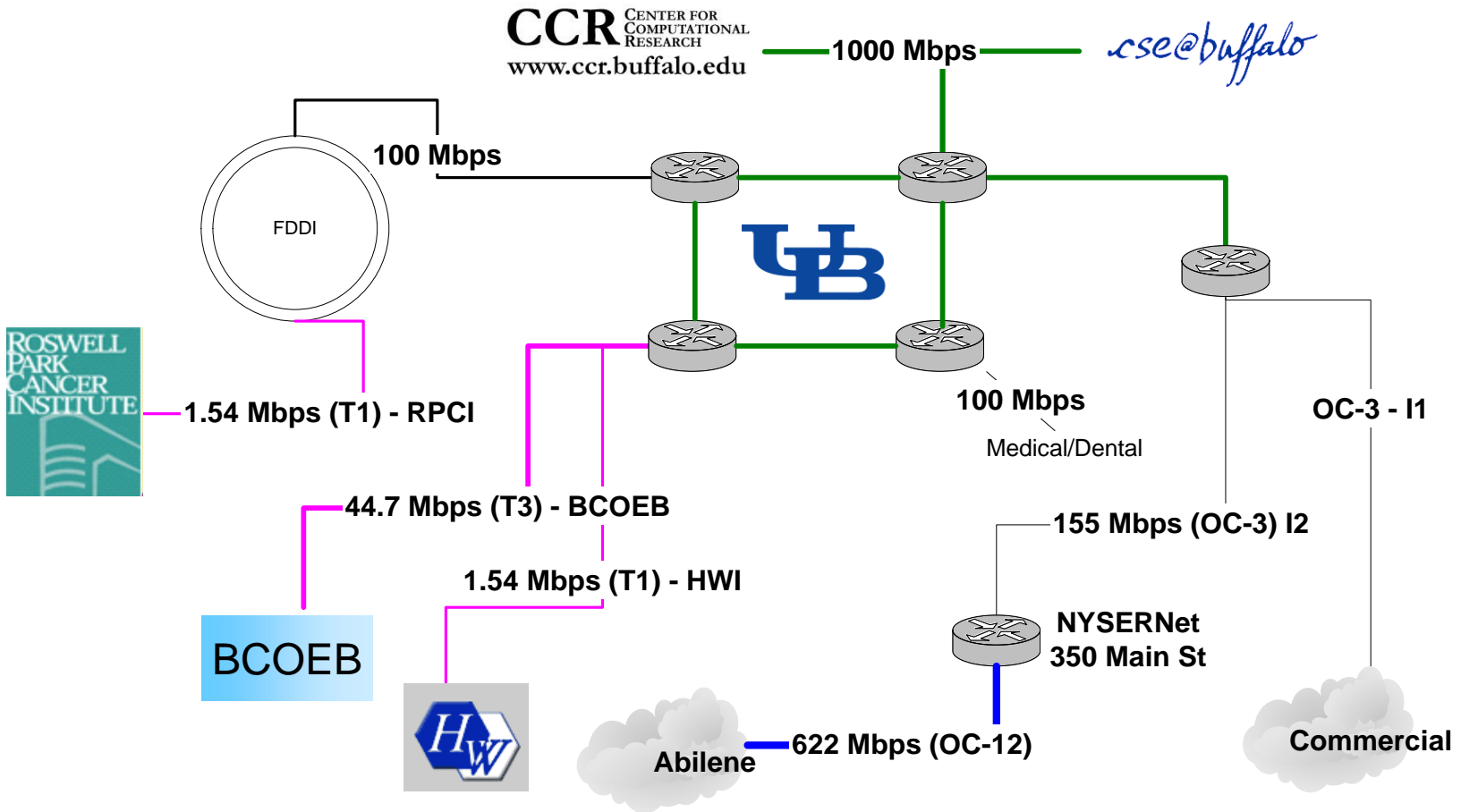
ACDC-Grid System Architecture



Initial ACDC Campus Grid



Network Connections

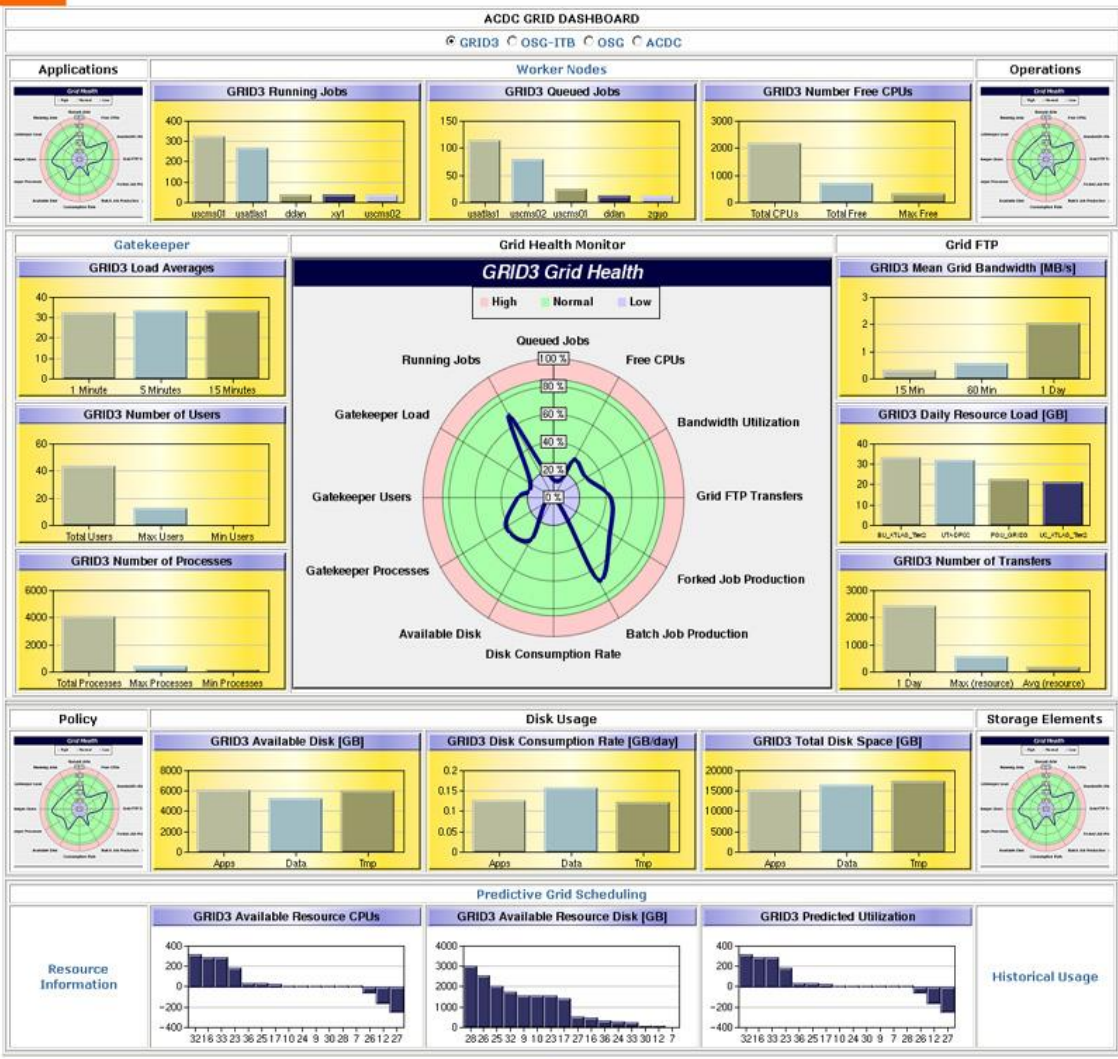




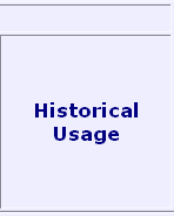
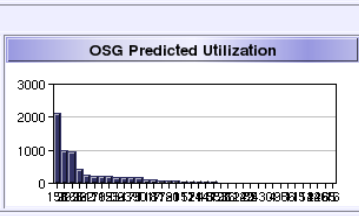
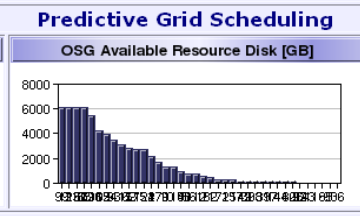
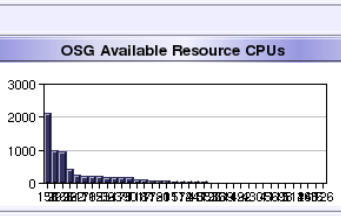
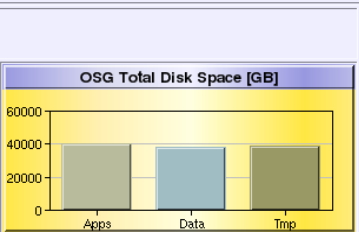
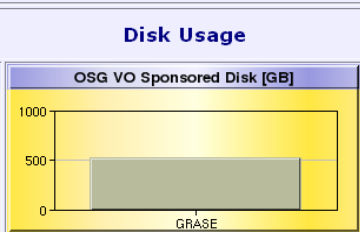
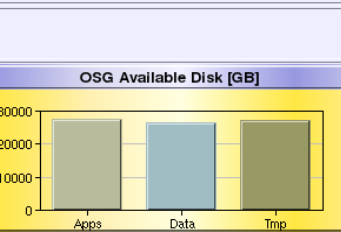
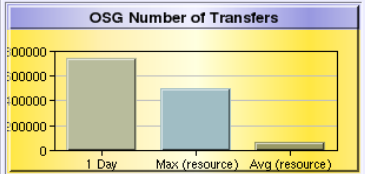
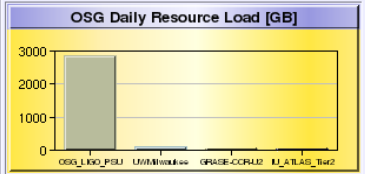
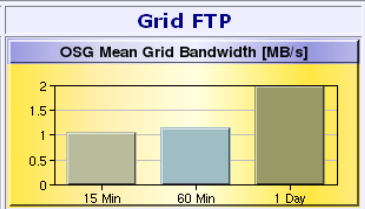
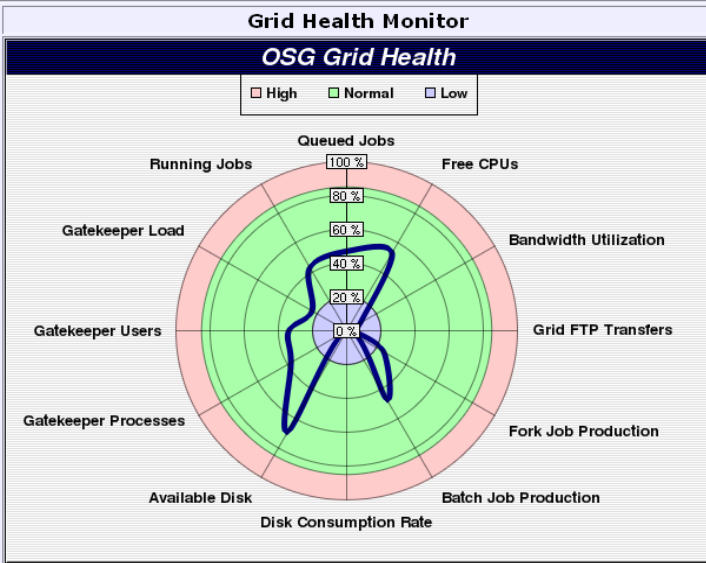
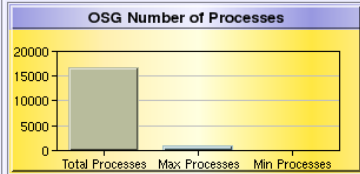
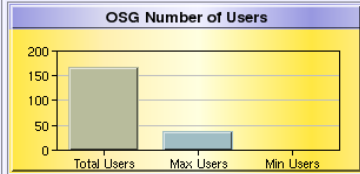
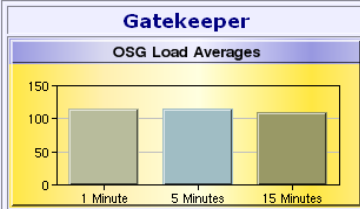
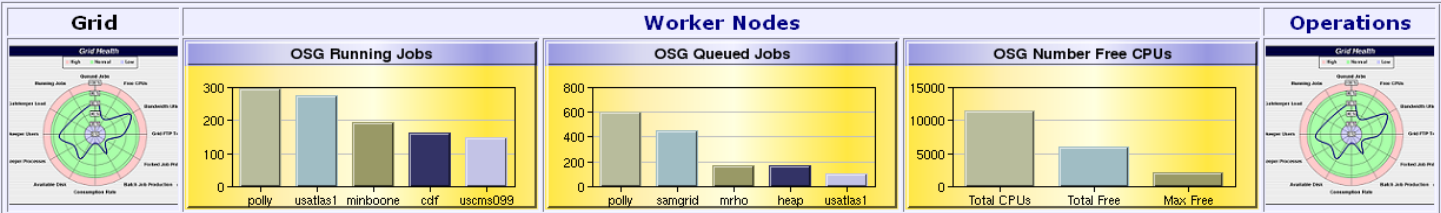
- Grid Resources
- ACDC Monitoring
- ACDC Grid Dashboard
- Running/Queued Jobs
- Job History
- Detailed Job History
- Detailed GridFTP History
- Resource Queue Visualization
- Resource User Visualization
- Sell Application Demonstrator
- Presentations
- ACDC Site Status
- Contact Us
- Staff Only



ACDC-Grid Monitoring: The ACDC-Grid DASHBOARD



- CI Lab Grid Monitor
- Grid Dashboard
- Operations Dashboard
- Historical Dashboard
- Running/Queued Jobs
- Job History
- Detailed Job History
- VO Sponsor CPUs
- Free/Running/Queued CPUs
- VO Support Matrix
- Current Bandwidth Matrix
- Historical Bandwidth Matrix
- Current Latency Matrix
- Historical Latency Matrix
- Resource Queue Visualization
- Resource User Visualization
- SnB Application Demonstrator
- ACDC Grid Dashboard
- Site Status
- ACDC Grid Dashboard Tutorial
- GRASE VO
 - Overview
 - Request Membership
 - Request Help
- Staff Only
- Contact Us / Staff
- CI Lab

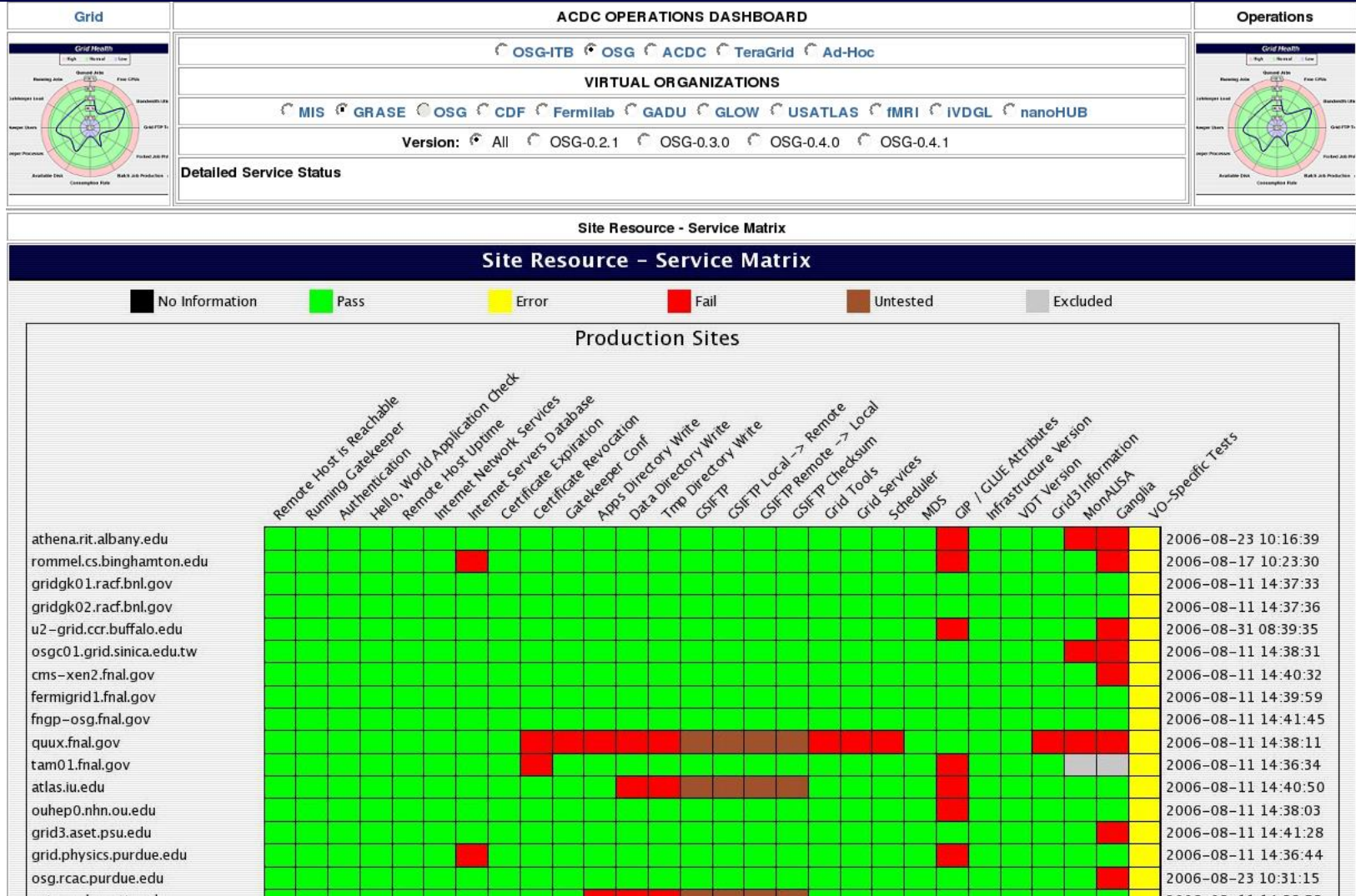


Supported by the National Science Foundation and the Department of Energy

CI Lab Grid Monitor: <http://osg.ccr.buffalo.edu/>

ACDC Monitor

<http://osg.ccr.buffalo.edu/operations-dashboard.php?grids=3&vos=10>



ACDC-Grid

CCR Grid Computing Services - Microsoft Internet Explorer

CCR Center for Computational Research GRID PORTAL High Performance Grid Computing

UNIVERSITY AT BUFFALO The State University of New York

Welcome to Grid Computing Services

University at Buffalo Center for Computational Research is currently forming the first Western New York computational grid. The computational grid consist of many supercomputers located at the Center and several other networked supercomputers throughout the Western New York region. These resources will be shared by many researchers from several departments working on a diverse suite of problems including Bioinformatics, Computational Chemistry, and Medical Imaging to name a few.

We also provide grid computing support for the University's Center for Computational Research learning & teaching and research activities plus the infrastructure for both high performance computing and grid enabled software.

Got your "Grid Computing Guide"?

PORTAL LOGIN
Grid General Info
About ACDC Grid
Computational Grid
Data Grid
Publications
Technical Papers
Presentations
Contact Us
Grid Account Request
Grid Account Support
Events
News
Projects
Resources
Education/Outreach
Staff Only
CCR HOME

CCR Grid Computing Services: Data Management - Microsoft Internet Explorer

CCR Center for Computational Research GRID PORTAL High Performance Grid Computing

UNIVERSITY AT BUFFALO The State University of New York

VIEW: Group: **GROUP:miller** UserList: rappleye

PORTAL LOGOUT
User Tools
Manage Account
Grid General Info
Projects
Resources
Computational Grid
Job Submission
Job/Queue Status
Data Grid
Network Status
Running/Queued Jobs
PBS Job History
Grid Portal Statistics
Conder Flock Statistics
User Info
Education/Outreach
Staff Only
CCR HOME

File tree view showing folders: rappleye, KeyMaster, Morpheus, Tank, Agent, Rabbit, Tank, Morpheus, Oracle.m, Neo.

Browser view of "miller" group files published

CCR Grid Computing Services: Grid Admin - Microsoft Internet Explorer

CCR Center for Computational Research GRID PORTAL High Performance Grid Computing

UNIVERSITY AT BUFFALO The State University of New York

View statistics for: disk_space
Data based on: group
from starting date: January 1 2000
to ending date: September 13 2003 inclusive
for: Grid_Portal resources OK

Datagrid Historical Group Disk Space Usage

PORTAL LOGOUT
User Tools
Manage Account
Grid General Info
Projects
Resources
Computational Grid
Job Submission
Job/Queue Status
Data Grid
Network Status
Running/Queued Jobs
PBS Job History
Grid Portal Statistics
Conder Flock Statistics
User Information
Education/Outreach
Staff Only
CCR HOME

CCR Grid Computing Services: Grid Admin - Microsoft Internet Explorer

CCR Center for Computational Research GRID PORTAL High Performance Grid Computing

UNIVERSITY AT BUFFALO The State University of New York

View statistics for: disk_space
Data based on: user
from starting date: January 1 2000
to ending date: September 13 2003 inclusive
for: Grid_Portal resources OK

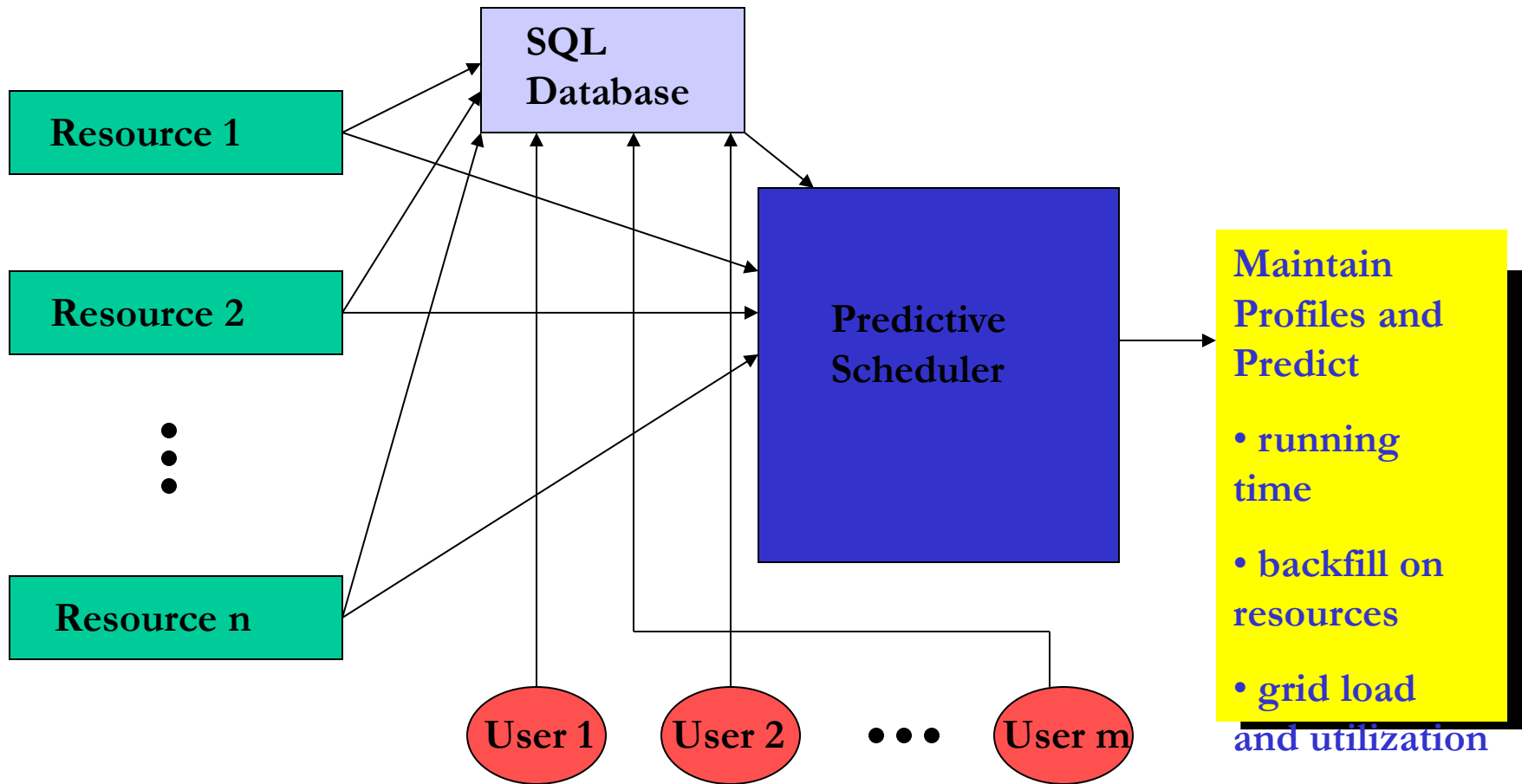
File_num	File_ID	Filename	Dir_ID	Resource_ID	Owner	Groupname	Type
1	56033	Cypher.txt	52831	10	mlgreen	griddev	txt
2	56034	Cypher.sh	52858	10	mlgreen	griddev	sh
3	56035	Oracle.asc	52958	10	mlgreen	griddev	asc
4	56036	Cypher.sh	52634	10	mlgreen	miller	sh
5	56037	Rabbit.dat	52830	10	mlgreen	ccrstaff	dat
6	56038	Agent.exe	53064	10	mlgreen	griddev	exe
7	56039	Dozer.sh	52852	10	mlgreen	griddev	sh
8	56040	Neo.asc	52187	10	mlgreen	mlgreen	asc
9	56041	Agent.mpg	52833	10	mlgreen	mlgreen	mpg
10	56042	Tank.txt	52188	10	mlgreen	mlgreen	txt
11	56043	Smith.xls	52298	10	mlgreen	ccrstaff	xls
12	56044	KeyMaster.csh	52186	10	mlgreen	miller	csh
13	56045	Oracle.csh	52622	10	mlgreen	griddev	csh
14	56046	Dozer.xls	52808	10	mlgreen	mlgreen	xls
15	56047	Cypher.exe	52204	10	mlgreen	griddev	exe
16	56048	Rabbit.ppt	52861	10	mlgreen	miller	ppt
17	56049	Neo.dat	52217	10	mlgreen	ccrstaff	dat
18	56050	Cypher.asc	53086	10	mlgreen	ariddev	asc

PORTAL LOGOUT
User Tools
Manage Account
Grid General Info
Projects
Resources
Computational Grid
Job Submission
Job/Queue Status
Data Grid
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PBS Job History
Grid Portal Statistics
Conder Flock Statistics
User Information
Education/Outreach
Staff Only
CCR HOME

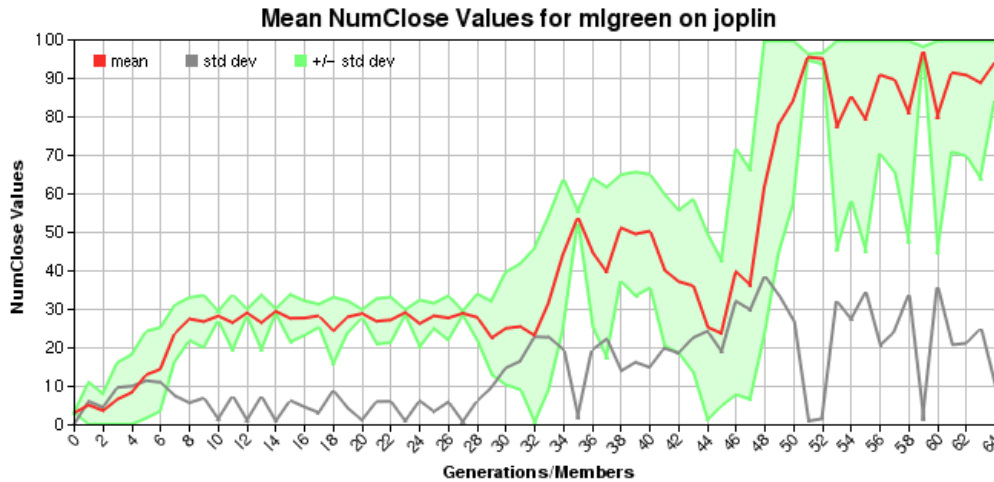
Predictive Scheduler

- Build profiles based on statistical analysis of logs of past jobs
 - Per User/Group
 - Per Resource
- Use these profiles to predict runtimes of new jobs
- Make use of these predictions to determine
 - Resources to be utilized
 - Availability of Backfill

System Diagram



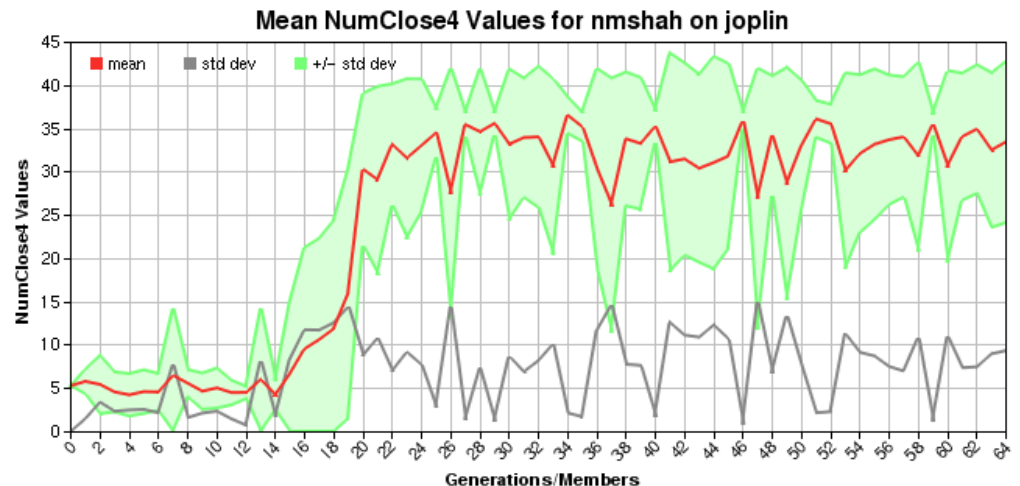
Preliminary GA results



Percent of estimates within 5% of actual values



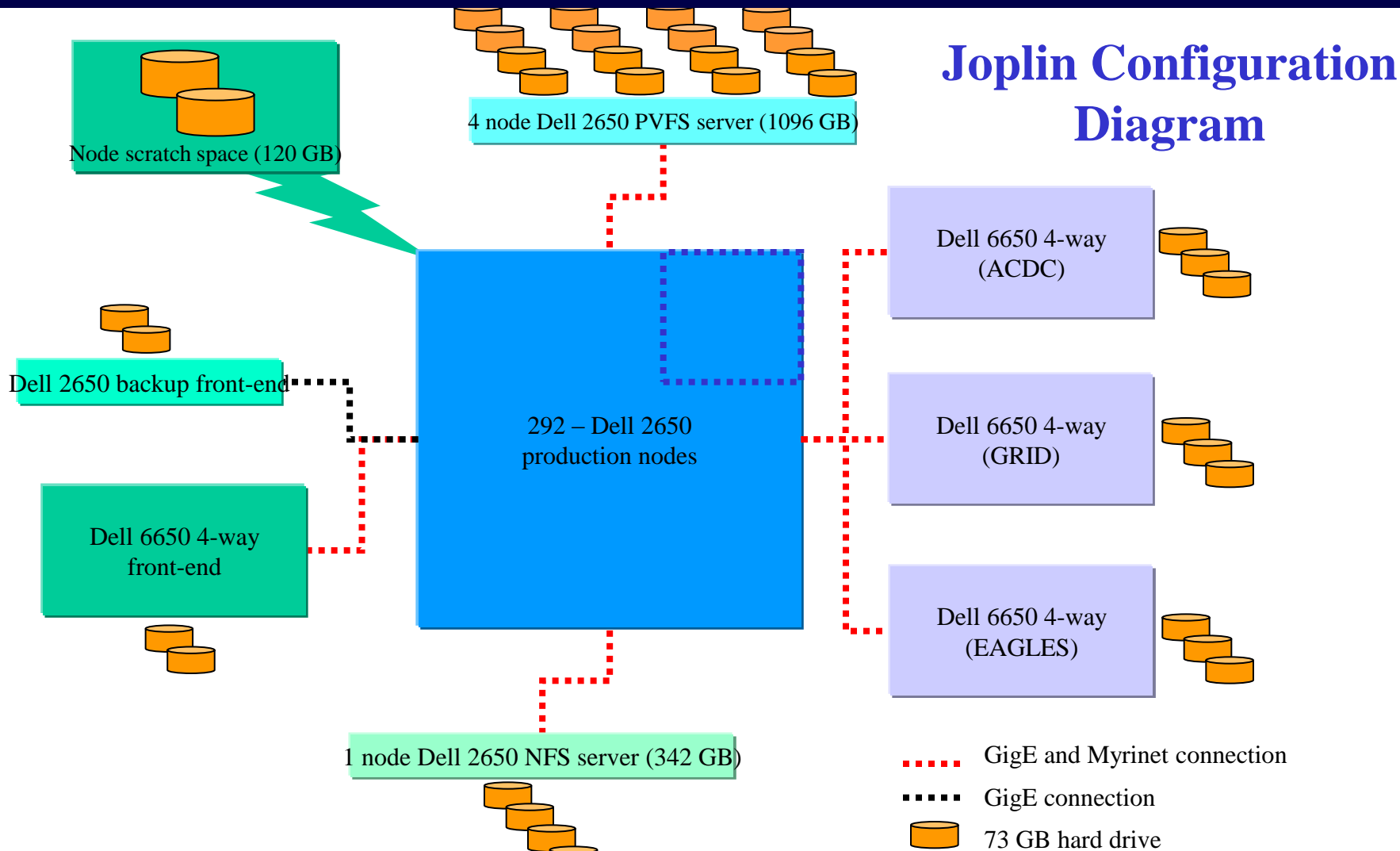
Percent of estimates within 20% of actual values



ACDC-Grid Dynamic Resource Allocation at SC03 with Grid3

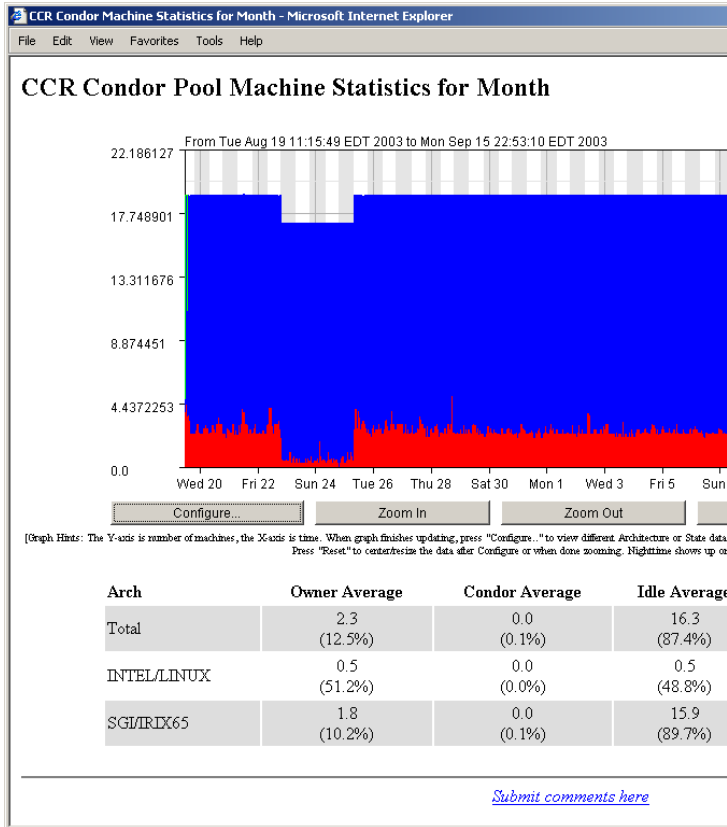
- Small number (40) of CPUs were dedicated at night
- An additional 400 CPUs were dynamically allocated during the day
- No human intervention was required
- Grid applications were able to utilize the resources and surpassed the Grid3 goals

ACDC-Grid Dynamic Resource Allocation



ACDC-Grid Portal Condor Flock

CondorView integrated into ACDC-Grid Portal



CCR Grid Computing Services: Grid Admin - Microsoft Internet Explorer

CCR Center for Computational Research GRID PORTAL High Performance Grid Computing

PORTAL LOGOUT

User Tools

- » Manage Account
- Grid General Info
- Projects
- Resources
 - » Computational Grid
 - » Job Submission
 - » Job/Queue Status
 - » Data Grid
 - » Data Grid Statistics
 - » Network Status
 - » Running/Queued Jobs
 - » PBS Job History
 - » Grid Portal Statistics
 - » Condor Flock Statistics
 - » User Information
- Education/Outreach
- Staff Only
- CCR HOME

Condor High Throughput Computing

Condor Pool Statistics for CCR

Pool Resource (Machine) Statistics

- For the past hour
- For the past day
- For the past week
- For the past month
- For the month of [Jan] [Feb] [Mar] [Apr] [May] [Jun] [Jul] [Aug] [Sep] [Oct] [Nov] [Dec]

Pool User (Job) Statistics

- For the past hour
- For the past day
- For the past week
- For the past month
- For the month of [Jan] [Feb] [Mar] [Apr] [May] [Jun] [Jul] [Aug] [Sep] [Oct] [Nov] [Dec]

Submit comments here

Advanced Center for Computational Research Data Center

GRID

Grid Administration

The collage displays four screenshots of the CCR Grid Portal interface:

- Grid Site Administration:** Shows a sidebar with navigation options like 'Users', 'Groups', and 'Organizations'. The main content area lists administrative tasks for these categories.
- Generate Globus grid-mapfile:** A form for generating a grid-mapfile, including fields for 'Optional include file' and 'Optional grid-mapfile path', and a checkbox for 'Do not stage the file to the grid nodes'.
- Create New Database Job:** A form to create a new database job, with fields for 'Job Name', 'Full Path To Script', 'Accepts Arguments', 'Run Script', and 'Run As User'.
- MDS Resource Update Status:** A table showing the status of various resources, including their names, last update times, next update intervals, and overall status (OK).

Data Grid

■ Motivation:

- ❑ Large data collections are emerging as important community resources.
- ❑ Data Grids complement Computational Grids.

■ Definition: *A data grid is a network of distributed storage resources, including archival systems, caches, and databases, which are linked logically to create a sense of global persistence.*

■ Goal: Design and implement transparent management of data distributed across heterogeneous resources.

ACDC-Grid Data Grid

The screenshot shows a web browser window titled "CCR Grid Computing Services: Data Management - Microsoft Internet Explorer". The page header includes the University at Buffalo logo and the text "Center for Computational Research GRID PORTAL High Performance Grid Computing".

On the left side, there is a "PORTAL LOGOUT" section with a list of links: "User Tools" (Manage Account), "Grid General Info", "Projects", "Resources" (Computational Grid, Job Submission, Job/Queue Status, Data Grid, Network Status, Running/Queued Jobs, PBS Job History, Grid Portal Statistics, Conder Flock Statistics, User Information), and "Education/Outreach" (Staff Only, CCR HOME).

The main content area displays a file tree for the user "rappleye". At the top, there are controls: "VIEW Group", "GROUP miller", and "UserList rappleye". The file tree shows a hierarchy of folders: KeyMaster, Morpheus, Tank, Agent, Rabbit, Tank, Morpheus, Oracle.m, Neo, Neo, Cypher, Neo, Morpheus, and Oracle. A 3D molecular model is overlaid on the file tree. A yellow callout bubble points to the "Oracle.m" file with the text: "Browser view of 'miller' group files published by user 'rappleye'".

At the bottom of the page, there is a footer with the text "Advanced Center for Computational Research Data" and a stylized "GRID" logo.

ACDC-Grid Data Grid Functionality

- Basic file management functions are accessible via a platform-independent web interface.
- User-friendly menus/interface.
- File Upload/Download to/from the Data Grid Portal.
- Simple Web-based file editor.
- Efficient search utility.
- Logical display of files (user/ group/ public).
- Ability to logically display files based on metadata (file name, size, modification date, etc.)

ACDC-Grid

Data Grid File Migration

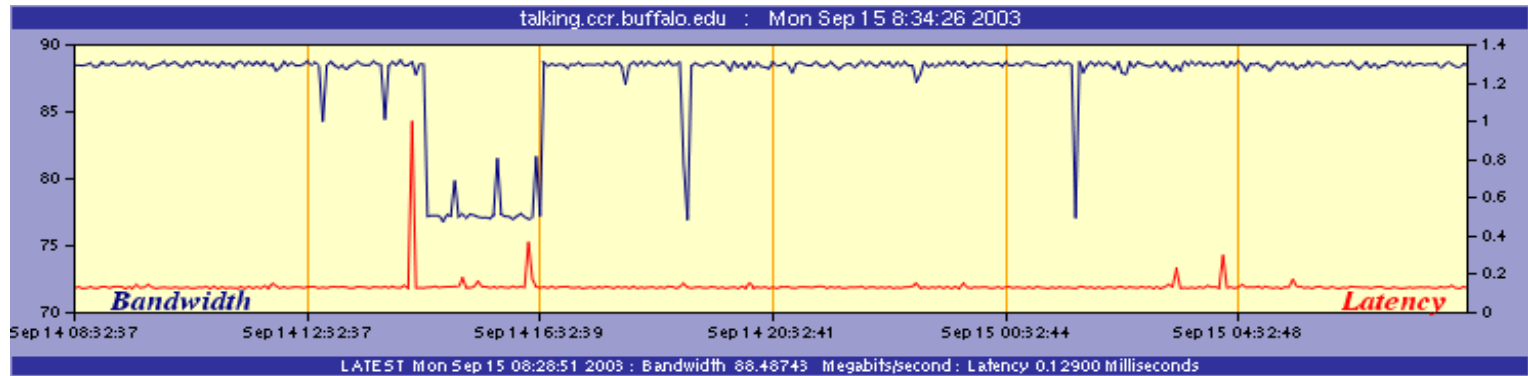
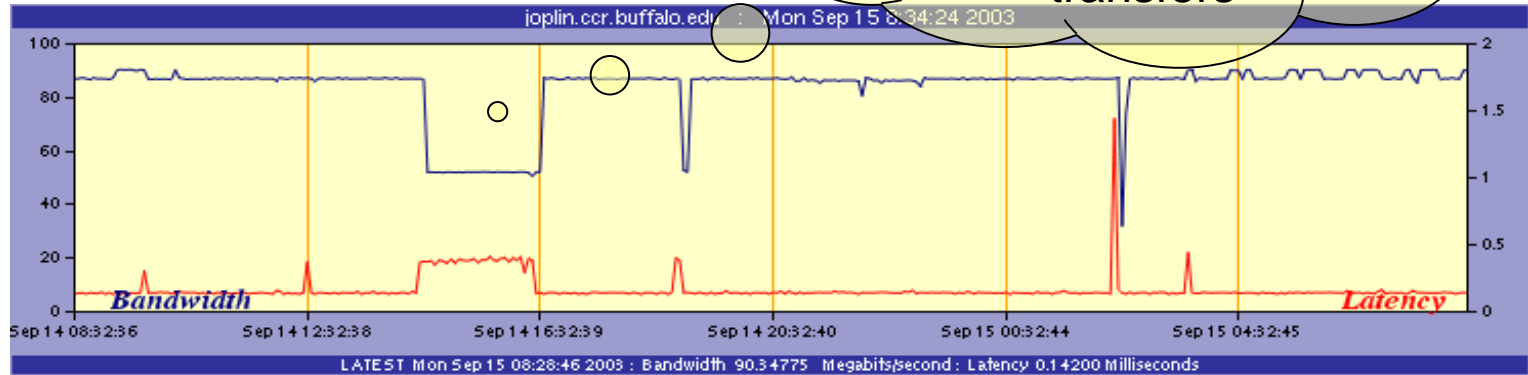
- Migration Algorithm dependent on
 - User access time
 - Network capacity at time of migration
 - User profile
 - User disk quotas on various resources

Data Grid File Aging

- For a given user, the average of the `file_aging_local_param` attributes of all files should be close to 1.
 - Operating tolerance before action is taken is within the range of 0.9 – 1.1.
- In this way, the user `file_aging_global_param` can be a function of this average.
 - If the average `file_aging_local_param` attribute > 1.1 , then files of the user are being held too long before being migrated.
 - The `file_aging_global_param` value should be decreased.
 - If the average `file_aging_local_param` attribute < 0.9 , then files of the user are being accessed at a higher frequency than the `file_aging_global_param` value.
 - The `file_aging_global_param` value should be increased.

Data Grid Resource Info

Both platforms have reduced bandwidth available for additional transfers



Grid Services and Applications

ACDC-Grid
Computational
Resources



Applications

Shake-and-Bake

Apache

MySQL

Oracle

High-level Services and Tools

Globus
Toolkit

NWS

MPI

MPI-IO

C, C++, Fortran, PHP

globusrun

Core Services

Metacomputing
Directory
Service

Globus
Security
Interface

GRAM

GASS

ACDC-Grid
Data
Resources



Local Services

Condor

Stork

MPI

RedHat Linux

WINNT

LSF

PBS

Maui Scheduler

TCP

UDP

Irix

Solaris

Adapted from Ian Foster and Carl Kesselman

Grid-Enabling Application Templates (GATs)

■ Structural Biology

- ❑ *SnB* and *BnP* for Molecular Structure Determination/Phasing

■ Groundwater Modeling

- ❑ *Ostrich*: Optimization and Parameter Estimation Tool
- ❑ *POMGL*: Princeton Ocean Model Great Lakes for Hydrodynamic Circulation
- ❑ *Split*: Modeling Groundwater Flow with Analytic Element Method

■ Earthquake Engineering

- ❑ *EADR*: Evolutionary Aseismic Design and Retrofit; Passive Energy Dissipation System for Designing Earthquake Resilient Structures

■ Computational Chemistry

- ❑ *Q-Chem*: Quantum Chemistry Package

■ Geographic Information Systems & BioHazards

- ❑ *Titan*: Computational Modeling of Hazardous Geophysical Mass Flows

Grid Enabled *SnB*

■ Problem Statement

- Use all available resources for determining a single structure

■ Grid Enabling Criteria

- Run on heterogeneous set of resources
- Store results in *SnB* database
- Mine database (and automagically deploy new jobs) to improve parameter settings

■ Runtime Parameters Transparent to User

- Assembling Necessary Files
- Number of Processors
- Trials per Processor
- Appropriate Queue and Running Times

Middleware

■ Grid (Computational and Data)

- ❑ Globus Toolkit 2.2.4 → direct upgrade WSRF
- ❑ Condor 6.6.0
- ❑ Network Weather Service 2.6
- ❑ Apache2 HTTP Server
- ❑ PHP 4.3.0
- ❑ MySQL 3.23
- ❑ phpMyAdmin 2.5.1

■ Collaboratory

- ❑ OpenGL (LibDMS, DevIL, GLUT)
- ❑ Windows, IRIX, Mac OS X, Linux
- ❑ CAVE, Desktop

Grid Enabled *SnB*

■ Required Layered Grid Services

□ Grid-enabled Application Layer

- *Shake – and – Bake* application
- Apache web server
- MySQL database

□ High-level Service Layer

- Globus, NWS, PHP, Fortran, and C

□ Core Service Layer

- Metacomputing Directory Service, Globus Security Interface, GRAM, GASS

□ Local Service Layer

- Condor, MPI, PBS, Maui, WINNT, IRIX, Solaris, RedHat Linux

Cyberinfrastructure Laboratory

Grid Portal

Dr. Russ Miller

UB Distinguished Professor of Computer Science & Engineering



CI Lab

Grid Portal Info

- Overview
- Portal Login
- Grid Account Info

Computational Grid

- Job Submission
- Job/Queue Status
- MDS Information
- Network Status
- Running/Queued Jobs
- PBS Job History
- Condor Flock Statistics
- GAT/Resource Matrix

Data Grid

- Data Grid Tree
- Data Grid Upload
- Data Grid Download
- Data Grid File Manager
- Data Grid Replica Manager
- Data Grid Simulator
- Data Grid Admin Tools
- Data Grid Admin File Tools

Contact Us / Staff

- CI Lab
- Staff Only

Welcome to the Cyberinfrastructure Laboratory Grid Portal

The **Cyberinfrastructure Laboratory**, in conjunction with the **Center for Computational Research**, has created an integrated Data and Computational Grid. This site is devoted to a Grid Portal that provides access to applications that can be run on a variety of grids. A related site contains a **Grid Monitoring System** designed by the Cyberinfrastructure Laboratory.

Applications may be run on the Cyberinfrastructure Laboratory's **ACDC Grid**, **Western New York Grid**, and **New York State Grid**, which includes computational and data storage systems from dozens of institutions throughout the State of New York.

The applications available to the users cover a variety of disciplines, including Bioinformatics, Computational Chemistry, Crystallography and Medical Imaging, to name a few.

The grids developed by the CI Lab support teaching and research activities, as well as providing infrastructure that includes high-end data, computing, imaging, grid-enabled software, all of which relies on the New York State Research Network (**NYSERNet**).

This work is funded by the National Science Foundation (ITR, MRI, CRI), three program projects from The National Institutes of Health, and the Department of Energy.



Software : BnP

Field : Protein crystal structure determination

Startup Screen for CI Lab Grid Job Submission

Advanced Computational Data Center Grid Job Submission Instructions

The grid-enabling application templates used on the ACDC-Grid are created from the application developers grid user profiles that contain the users standard information uid, name, organization, address, etc., and more specific information such as group id and access level information for each of grid-enabled applications. This information is stored in a database for each of the grid-enabled applications and can be accessed through selected queries throughout the ACDC-Grid Web Portal.

Additionally, each grid-enabled scientific application profile contains information about specific execution parameters, required data files, optional data files, computational requirements, etc. and statistics on application historical ACDC-Grid jobs for predictive runtime estimates. MySQL provides the speed and reliability required for this task and it is currently being used as the ACDC-Grid Web Portal database provider.

The grid-enabled versions of many well-defined scientific and engineering applications have very similar general requirements and core functionality that are require for execution in the ACDC-Grid environment. We have identified that sequentially defining milestones for the grid user to complete intuitively guides them through the application workflow.

- Software Application:** Grid user chooses a grid-enabled software application.
- Template:** Grid user selects the required and/or optional data files from the ACDC Data Grid. User defined computational requirements are input or a template defined computational requirement runtime estimate is selected.
- Job Definition:** Grid user defines application specific runtime parameters or accepts default template parameter definitions.
- Review:** Grid user accepts the template complete job definition workflow or corrects any part of job definition.
- Execution Scenario:** The grid user has the ability to input an execution scenario or select a ACDC-Grid determined template defined execution scenario.
- Grid Job Status:** The grid user can view specific grid job completion status, grid job current state (COMPLETE, RUNNING, QUEUED, BLOCKED, FAILED, ETC.), detailed information on all running or queued grid jobs and grid-enabled application specific intermediate and post processing grid job graphics, plots and tables.

Each item of the job definition workflow is then stored in the ACDC-Grid Web Portal database so the grid user may use/modify any previously created workflow in creating new job definitions. The job definitions can also be accessed via batch script files for executing hundreds of similar workflows in an automated fashion. For example, a grid user would first define/save a relatively generic job workflow template for the grid-enabled application and then use the batch script capabilities to change the job definition workflow data files or application parameters and execute a series of new grid jobs.

Instructions and Description for Running a Job on ACDC-Grid

Software → Template → General Information → Detailed Information → Job Definition → Review → Execution Scenario

Select a GAT: BnP Auto Run

- BnP Auto Run
- EADR
- Ostrich
- POM
- Q-Chem
- SnB**
- SnB DREAR
- Split
- snb-dev

Continue

Reset Current Stage

Cancel

Return to the

Software Package Selection

Software → Template → General Information → Detailed Information → Job Definition → Review → Execution Scenario

Enter structure definition manually

Select structure from Data Grid:

Select Config File

Continue

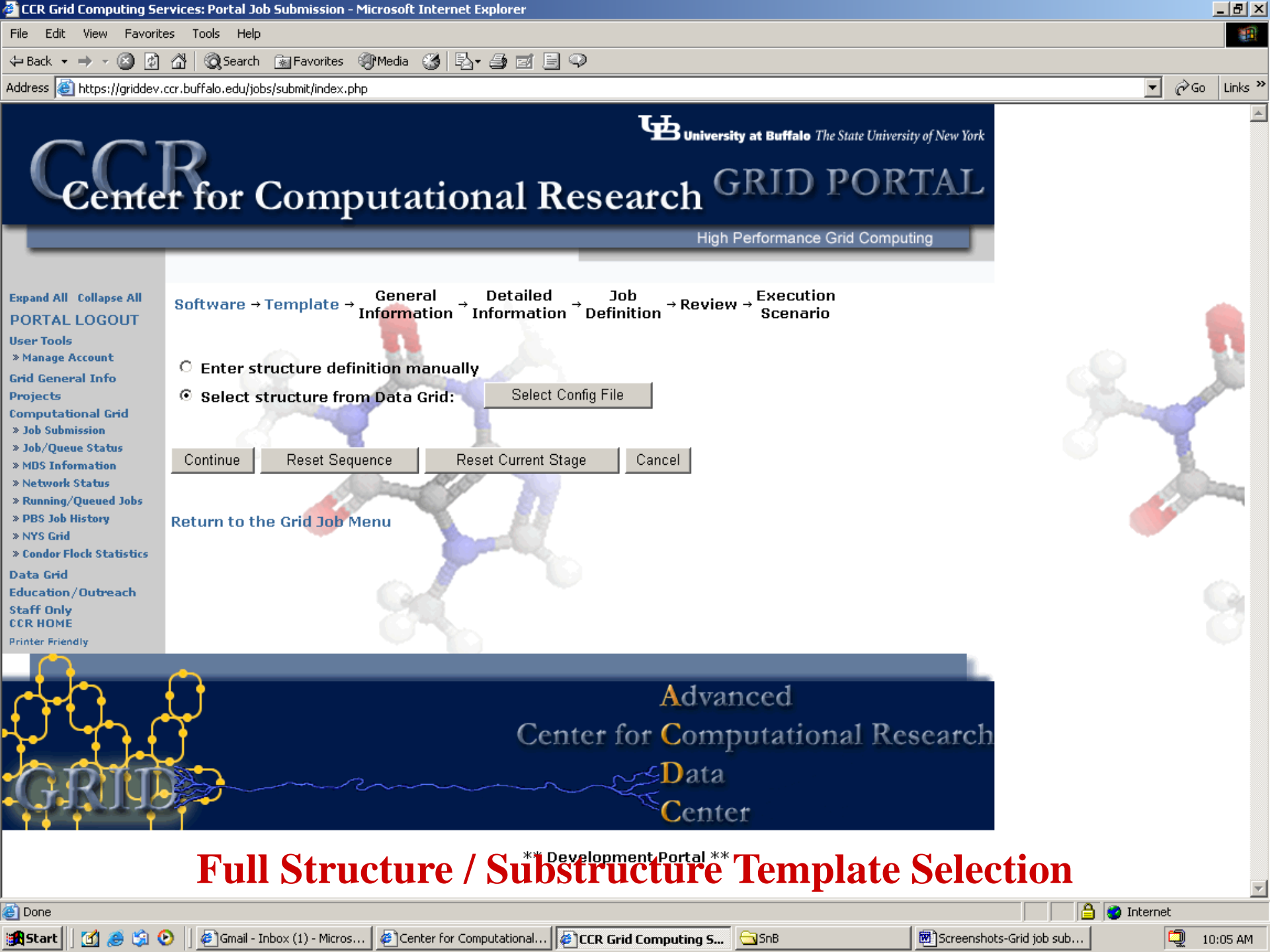
Reset Sequence

Reset Current Stage

Cancel

Return to the Grid Job Menu

Full Structure / Substructure Template Selection



CCR Center for Computational Research GRID PORTAL

High Performance Grid Computing

Software -> **Template** -> General Information -> Detailed Information -> Job Definition -> Review -> Execution Scenario

- Enter structure definition manually
- Select structure from Data Grid:

[Return to the Grid Job Menu](#)



Full Structure / Substructure Template Selection
** Development Portal **

- USER TOOLS
- » Manage Account
- Grid General Info
- Projects
- Computational Grid
- » Job Submission
- » Job/Queue Status
- » MDS Information
- » Network Status
- » Running/Queued Jobs
- » PBS Job History
- » NYS Grid
- » Condor Flock Statistics
- Data Grid
- Education/Outreach
- Staff Only
- CCR HOME
- Printer Friendly

General Information

Structure Information

Title:

Structure ID :

Space Group :

Cell Constants and Cell Errors (Cell Errors optional)

A: +/-

B: +/-

C: +/-

Alpha: +/-

Beta: +/-

Gamma: +/-

Native Asymmetric Unit Contents

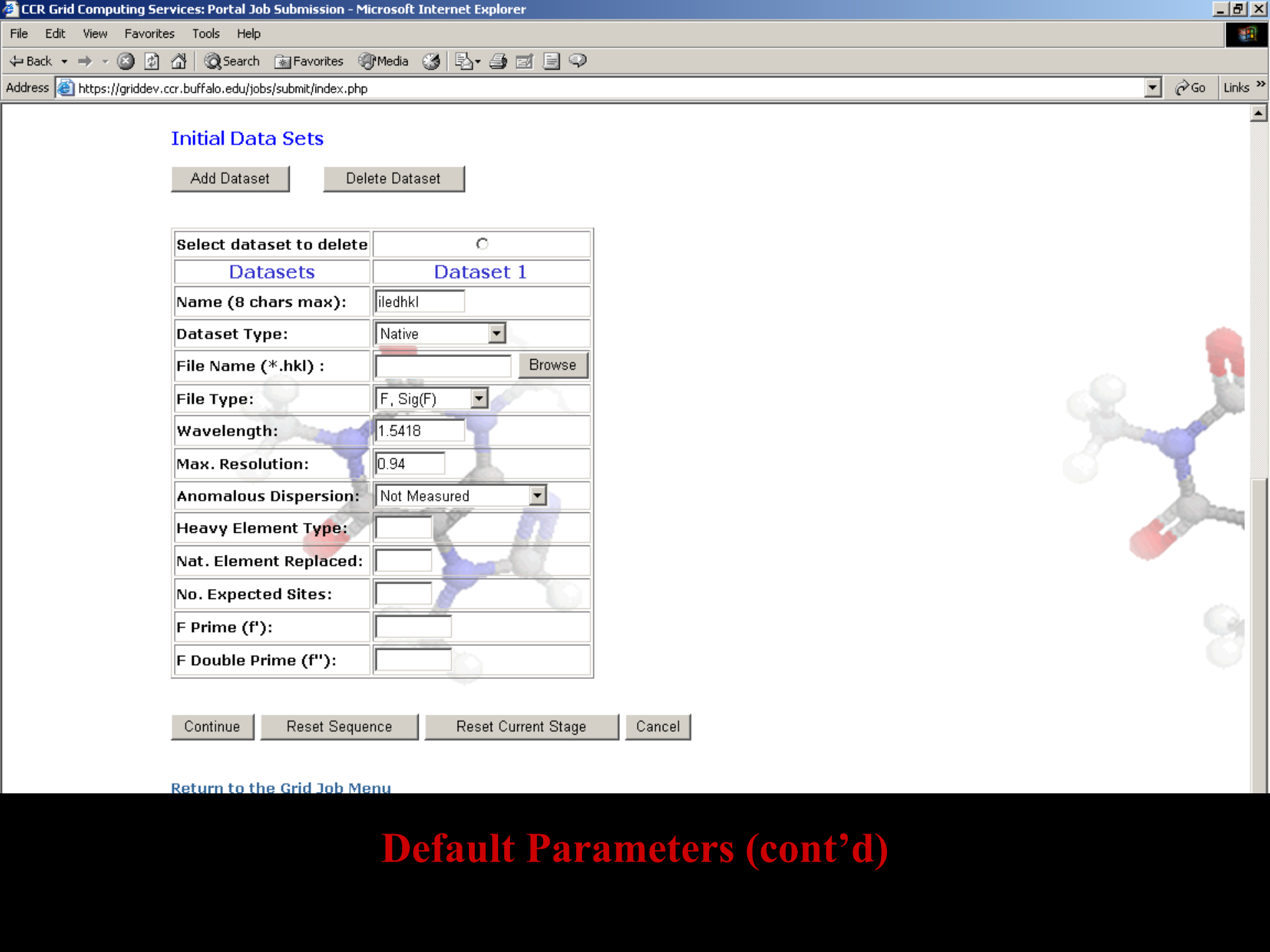
No Residues (Optional):

ASU Contents : (examples: C6H12O6 OR C6 H12 O6)

Initial Data Sets

Select dataset to delete	<input type="text" value="○"/>
Datasets	Dataset 1
Name (8 chars max):	<input type="text" value="iledhkl"/>

Default Parameters Based on Template



Initial Data Sets

Add Dataset

Delete Dataset

Select dataset to delete	Dataset 1
Datasets	Dataset 1
Name (8 chars max):	<input type="text" value="iledhkl"/>
Dataset Type:	<input type="text" value="Native"/>
File Name (*.hkl) :	<input type="text"/> <input type="button" value="Browse"/>
File Type:	<input type="text" value="F, Sig(F)"/>
Wavelength:	<input type="text" value="1.5418"/>
Max. Resolution:	<input type="text" value="0.94"/>
Anomalous Dispersion:	<input type="text" value="Not Measured"/>
Heavy Element Type:	<input type="text"/>
Nat. Element Replaced:	<input type="text"/>
No. Expected Sites:	<input type="text"/>
F Prime (f\')	<input type="text"/>
F Double Prime (f'')	<input type="text"/>

Continue

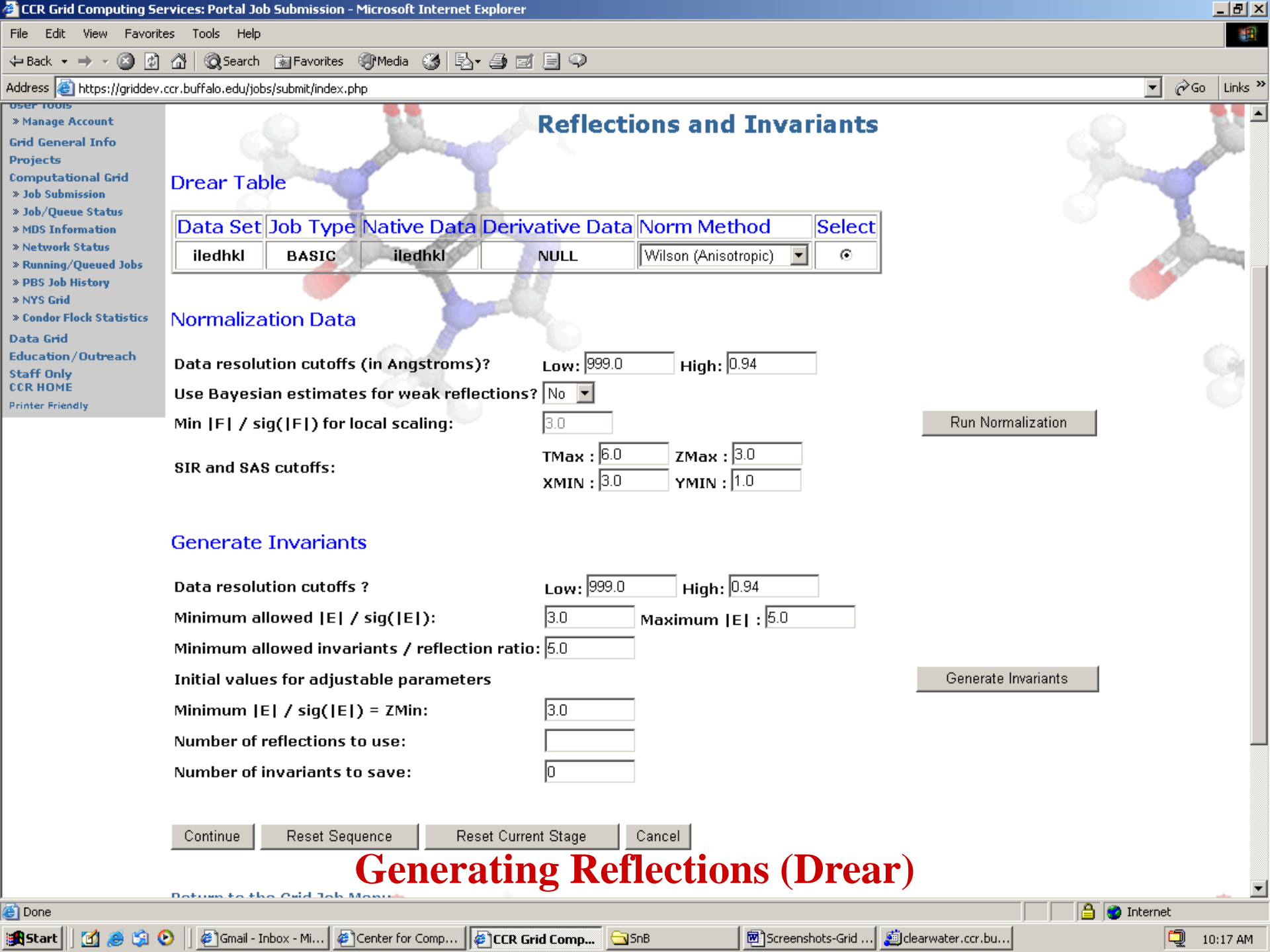
Reset Sequence

Reset Current Stage

Cancel

[Return to the Grid Job Menu](#)

Default Parameters (cont'd)



- USER TOOLS
- > Manage Account
- Grid General Info
- Projects
- Computational Grid
- > Job Submission
- > Job/Queue Status
- > MDS Information
- > Network Status
- > Running/Queued Jobs
- > PBS Job History
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Reflections and Invariants

Drear Table

Data Set	Job Type	Native Data	Derivative Data	Norm Method	Select
iledhkl	BASIC	iledhkl	NULL	Wilson (Anisotropic)	<input type="radio"/>

Normalization Data

Data resolution cutoffs (in Angstroms)? Low: High:

Use Bayesian estimates for weak reflections?

Min |F| / sig(|F|) for local scaling:

SIR and SAS cutoffs:

TMax : ZMax :

XMIN : YMIN :

Run Normalization

Generate Invariants

Data resolution cutoffs ? Low: High:

Minimum allowed |E| / sig(|E|): Maximum |E| :

Minimum allowed invariants / reflection ratio:

Initial values for adjustable parameters

Minimum |E| / sig(|E|) = ZMin:

Number of reflections to use:

Number of invariants to save:

Generate Invariants

Continue

Generating Reflections (Drear)

- User Tools
- » Manage Account
- Grid General Info
- Projects
- Computational Grid
- » Job Submission
- » Job/Queue Status
- » MDS Information
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- » Running/Queued Jobs
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Reflections and Invariants

Declar Table

Data Set	Job Type	Native Data	Derivative Data	Norm Method	Select
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Normalization Data

Data resolution cutoffs (in Angstroms)? Low: High:

Use Bayesian estimates for weak reflections?

Min |F| / sig(|F|) for local scaling:

SIR and SAS cutoffs: TMax : ZMax :
 XMIN : YMIN :

Generate Invariants

Data resolution cutoffs ? Low: High:

Minimum allowed |E| / sig(|E|): Maximum |E| :

Minimum allowed invariants / reflection ratio:

Initial values for adjustable parameters

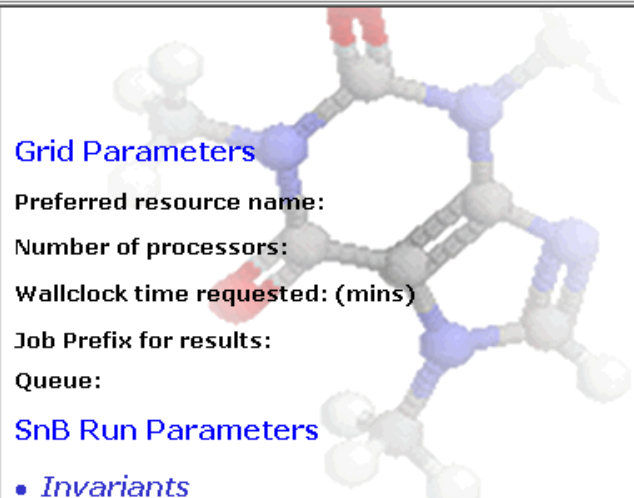
Minimum |E| / sig(|E|) = ZMin:

Number of reflections to use:

Number of invariants to save:

Invariant Generation

- User Tools
 - » Manage Account
- Grid General Info
- Projects
- Computational Grid
 - » Job Submission
 - » Job/Queue Status
 - » MDS Information
 - » Network Status
 - » Running/Queued Jobs
 - » PBS Job History
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SnB Setup

Grid Parameters

Preferred resource name:

Number of processors:

Wallclock time requested: (mins)

Job Prefix for results:

Queue:

SnB Run Parameters

• Invariants

Number of triplet invariants to use:

• Trials To Process

Starting phases from:

Random seed (prime):

Number of Trials:

Starting Trial:

Input Phase File:

Input Atom File:

Keep complete (every trial) peak file? :

• Cycles Information

Number of Shake-and-Bake cycles:

Keep complete (every cycle) trace file? :

Terminate trials failing the R-Ratio test? :

R-Ratio cutoff:

• Phase Refinement Method

SnB Setup

CCR Grid Computing: Portal Job Submission - Microsoft Internet Explorer

File Edit View Favorites Tools Help

Address <https://griddev.ccr.buffalo.edu/jobs/submit/index.php> Go Links

- *Phase Refinement Method*
Phase Refinement Method :
Number of passes through phase set:
Phase shift:
Number of shifts:
- *Real-Space Constraints*
Number of peaks to select:
Minimum interpeak distance:
Minimum distance between symmetry-related peaks:
Number of special position peaks to keep:
Fourier grid size:
Perform extra cycles with more peaks? :
Number of extra cycles :
Number of peaks :
- *Twice Baking*
Trials for E-Fourier filtering (fourier refinement)? :
Number of cycles :
Number of peaks :
Minimum |E| :
- *Automatic solution identification criteria*
Rmin Improvement (%):
Rcryst Imporvement (%):

SnB Setup (cont'd)

Done Start Gmail - Inbox - Mi... Center for Comp... CCR Grid Comp... SnB Screenshots-Grid ... clearwater.ccr.bu... Internet 10:18 AM

- User Tools
- » Manage Account
- Grid General Info
- Projects
- Computational Grid
- » Job Submission
- » Job/Queue Status
- » MDS Information
- » Network Status
- » Running/Queued Jobs
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SnB Job Review

Grid Job ID:	447
Selected resource:	clearwater.ccr.buffalo.edu
Number of processors:	5
Wallclock time requested:	720
Number of triplet invariant to use:	8400
Start Phases From:	Random Atoms
Random seed (prime):	11909
Number of trials:	1000
Starting Trial:	1
Input Phase File:	Unused
Input Atom File:	Unused
Keep complete (every trial) peak file? :	Yes
Number of Shake-and-bake cycles:	20
Keep complete (every cycle) trace file? :	No
Terminate trials failing the R-Ratio test? :	No
R-Ratio cutoff:	Unused
Phase Refinement Method:	Parameter Shift(Fast)
Number of passes through phase set:	3
Phase shift:	90.0
Number of shifts:	2
Number of peaks to select:	84
Minimum interpeak distance:	3
Minimum distance between symmetry-related peaks:	3.0
Number of special position peaks to keep:	0
Fourier grid size:	0.31
Perform extra cycles with more peaks? :	No
Number of extra cycles:	Unused
Number of peaks:	Unused
Trials for E-Fourier filtering (fourier refinement)? :	None
Number of cycles:	Unused
Number of peaks:	Unused
Minimum E :	Unused

***SnB* Review (Grid job ID: 447)**

Details for Grid Job 447 - iledhkl

Job Detail Information

Status: **RUNNING**

Rmin Min: 0.344 Rmin Max: 0.56

Last Updated: 15-Mar-2005 10:22:00

Total Trials: 1000

Complete Trials: 285

Resource: clearwater.ccr.buffalo.edu Processors: 5

Best Trial Number: 34

Best Trial Rmin: 0.344

Trial Summary

Grid Job 447 Trial Summary

Number of Trials Complete: 285 (28.5%)



Walltime Summary

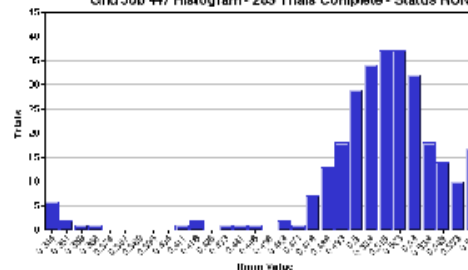
Grid Job 447 Walltime Summary

Walltime Consumed: 2 (0.3%)



Grid Job Trial Histogram

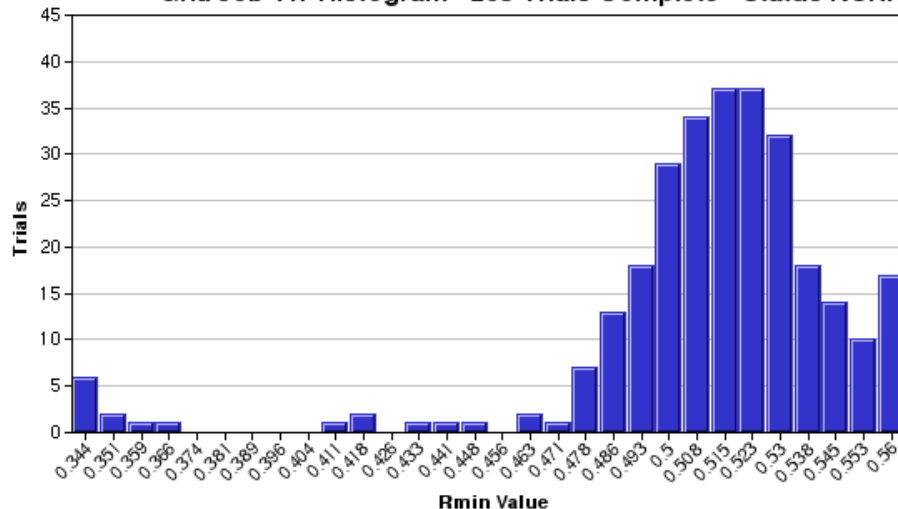
Grid Job 447 Histogram - 285 Trials Complete - Status RUNNING



Click on image for enlarged view.

Graphical Representation of Intermediate Job Status

Grid Job 447 Histogram - 285 Trials Complete - Status RUNNING



Histogram of Completed Trial Structures

Expand All Collapse All

PORTAL LOGOUT

User Tools

» Manage Account

Grid General Info

Projects

Computational Grid

» Job Submission

» Job/Queue Status

» MDS Information

» Network Status

» Running/Queued Jobs

» PBS Job History

» NYS Grid

» Condor Flock Statistics

Data Grid

Education/Outreach

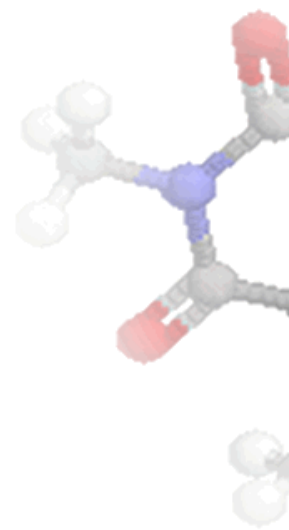
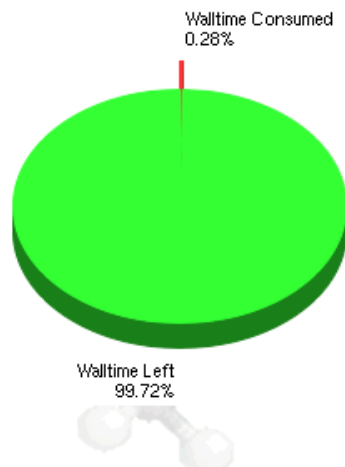
Staff Only

CCR HOME

Printer Friendly

Grid Job 447 Walltime Summary

Walltime Consumed: 2 (0.3%)



Walltime Summary Chart

Expand All Collapse All

PORTAL LOGOUT

User Tools

» Manage Account

Grid General Info

Projects

Computational Grid

» Job Submission

» Job/Queue Status

» MDS Information

» Network Status

» Running/Queued Jobs

» PBS Job History

» NYS Grid

» Condor Flock Statistics

Data Grid

Education/Outreach

Staff Only

CCR HOME

Printer Friendly

Grid Job Status

15-Mar-2005 10:23:49

Job Filter Criteria

Show GATs

BnP Auto Run
EADR
Ostrich
POM
Q-Chem
SnB
SnB DREAR

Job State

DEFINITION
STAGING
QUEUED
RUNNING
UPLOADING
COMPLETE
INCOMPLETE

Sort By









Job Id
Job Name
Resource
Num Procs
Status
Percent Complete
Last Update

Descending

Ascending

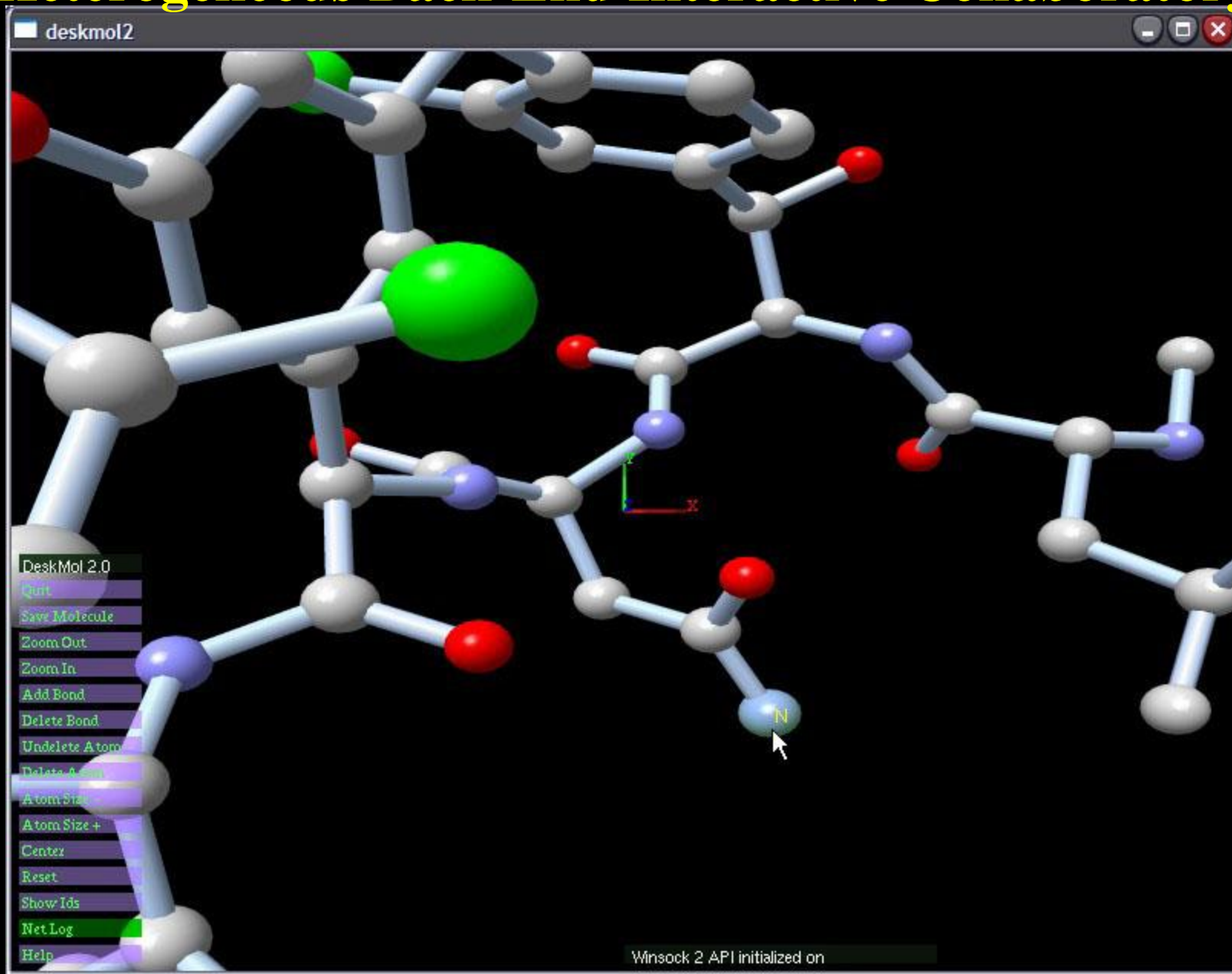
Filter Job List

SnB

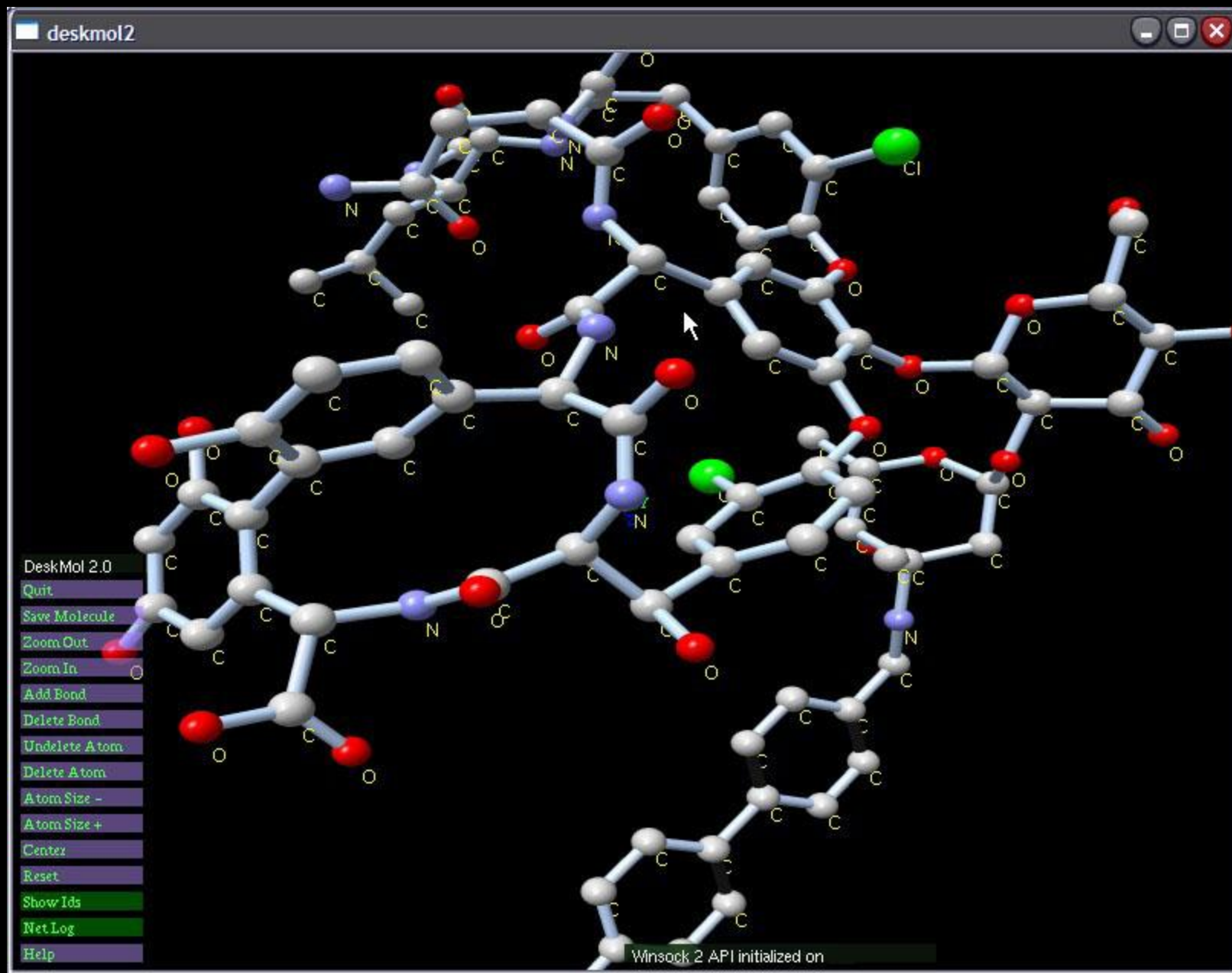
Job Id	Job Name	Resource	Num Procs	Status	Percent Complete	Last Update	Cancel Job	Drilldown
447	iledhkl	clearwater.ccr.buffalo.edu	5	RUNNING	28.5	15-Mar-2005 10:22:00	<input type="checkbox"/>	
446	trilys	clearwater.ccr.buffalo.edu	10	RUNNING	1	15-Mar-2005 10:22:00	<input type="checkbox"/>	
444	64chkl	nash.ccr.buffalo.edu	3	COMPLETE	100	14-Mar-2005 22:00:01		
443	trilys	clearwater.ccr.buffalo.edu	10	COMPLETE	100	10-Mar-2005 22:48:00		
442	pr435hkl	nash.ccr.buffalo.edu	5	COMPLETE	100	10-Mar-2005 17:26:01		
441	vancohkl	clearwater.ccr.buffalo.edu	10	COMPLETE	100	10-Mar-2005 18:08:01		
434	16chkl	clearwater.ccr.buffalo.edu	5	COMPLETE	100	10-Mar-2005 14:42:01		
433	16chkl	clearwater.ccr.buffalo.edu	5	COMPLETE	100	10-Mar-2005 14:38:01		

Status of Jobs

Heterogeneous Back-End Interactive Collaboratory



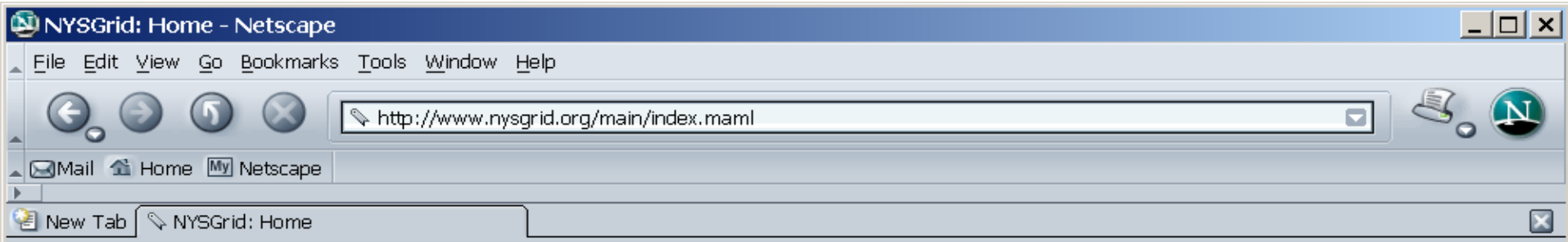
User starts up – default image of structure.



Molecule scaled, rotated, and labeled.

NYSGrid.org

- **Grass-Roots Cyberinfrastructure Initiative in NYS**
- **Open to academic, research, government, and industrial organizations.**
- **Goal is to allow transparent collection, management, organization, analysis, and visualization of data, while ignoring location.**
- **Enable Research, Scholarship, and Economic Development in NYS.**
- **Mission Stmt: To create and advance collaborative technological infrastructure that supports and enhances the research and educational missions of institutions in NYS.**



New York State Grid

Cyberinfrastructure for the 21st Century

[Home](#)
[Contact Us](#)

[About Us](#) [Organization](#) [Infrastructure](#) [User Information](#) [Events](#) [Media & News](#) [Grid Access](#)

In the 21st century, leading academic institutions will embrace our digital data-driven society and empower students to compete in this knowledge-based economy. In order to support research, scholarship, education, and community outreach, a grass-roots cyberinfrastructure initiative has been formed in New York State that will integrate research in disciplinary domains, including science, engineering, and biomedicine, with research in enabling technologies and interfaces. This initiative will allow students and scientists to transparently collect, manage, organize, analyze, and visualize data without having to worry about details such as where the data is stored, where the data is processed, where the data is rendered, and so forth. This ease of use and high availability of data and information processing tools will allow for revolutionary advances in all areas of science, engineering, and beyond.

Mission:

To create an advanced collaborative technological infrastructure that supports and enhances the research and educational missions of institutions in New York State.

Cyberinfrastructure sits at the core of modern simulation and modeling, which allows for entirely new methods of investigation that allow scholars to address previously unsolvable problems. Specifically, the development of necessary software, algorithms, portals, and interfaces that will enable research and scholarship by freeing end-users from dealing with the complexity of various computing environments is critical to extending the reach of high-end computing, storage, networking, and visualization to the general user community.

The Cyberinfrastructure Initiative consists of resources at institutions throughout the state. The initiative is open to all interested parties and more information can be found on some of the accompanying pages.

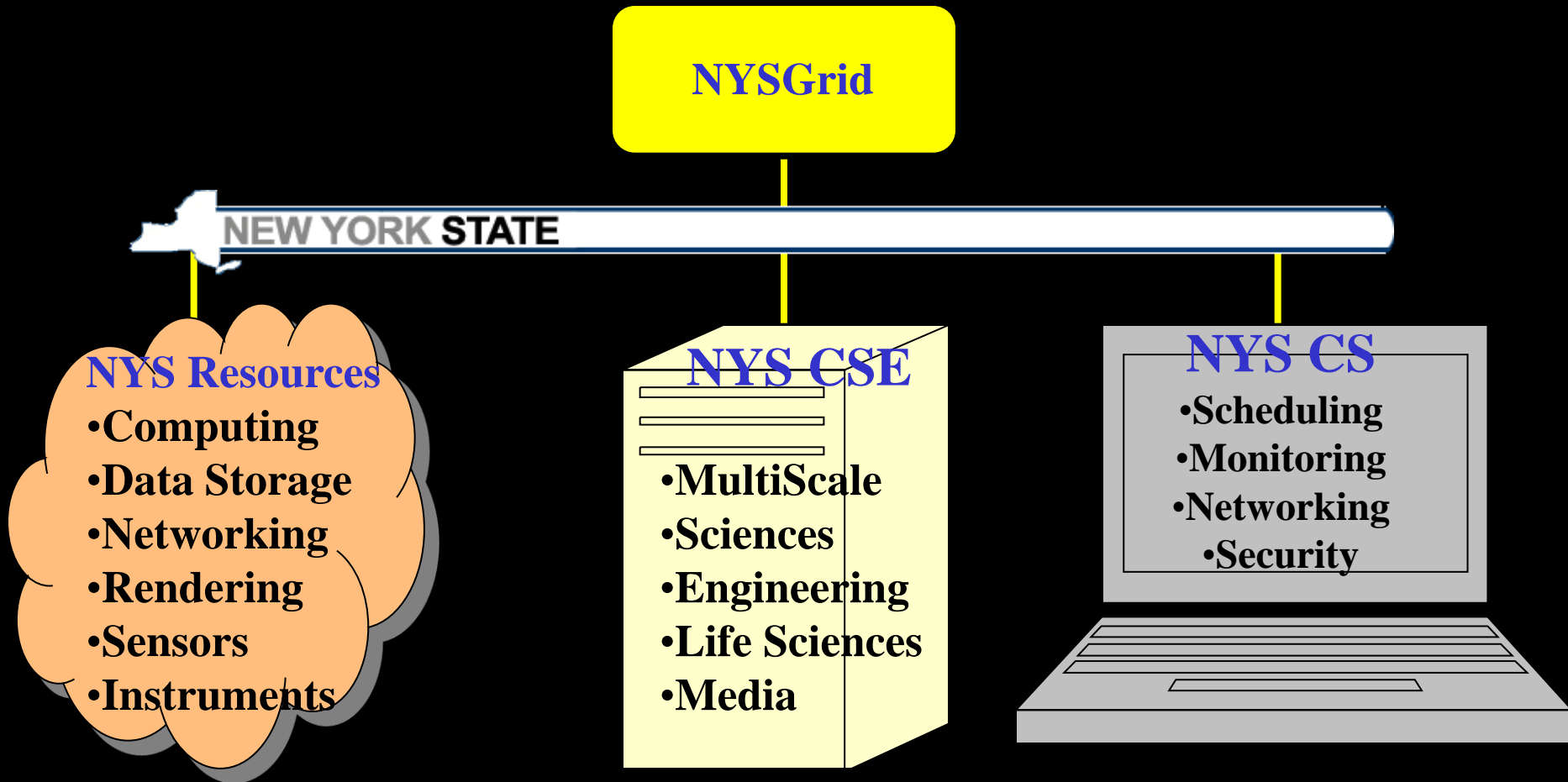
Current NYS Grid Participation

- Albany • ✓
- Alfred
- Binghamton • ✓
- Brookhaven
- Buffalo • ✓
- Columbia •
- Cornell • ✓
- Geneseo • ✓
- Hauptman-Woodward • ✓
- Iona •
- Marist • ✓
- Memorial Sloan-Kettering
- NYU • ✓
- Niagara • ✓
- RIT • ✓
- Rochester • ✓
- RPI •
- Stony Brook • ✓
- Syracuse • ✓
- NYSERNet

• - expressed interest in NYS Grid

✓ - on NYS Grid

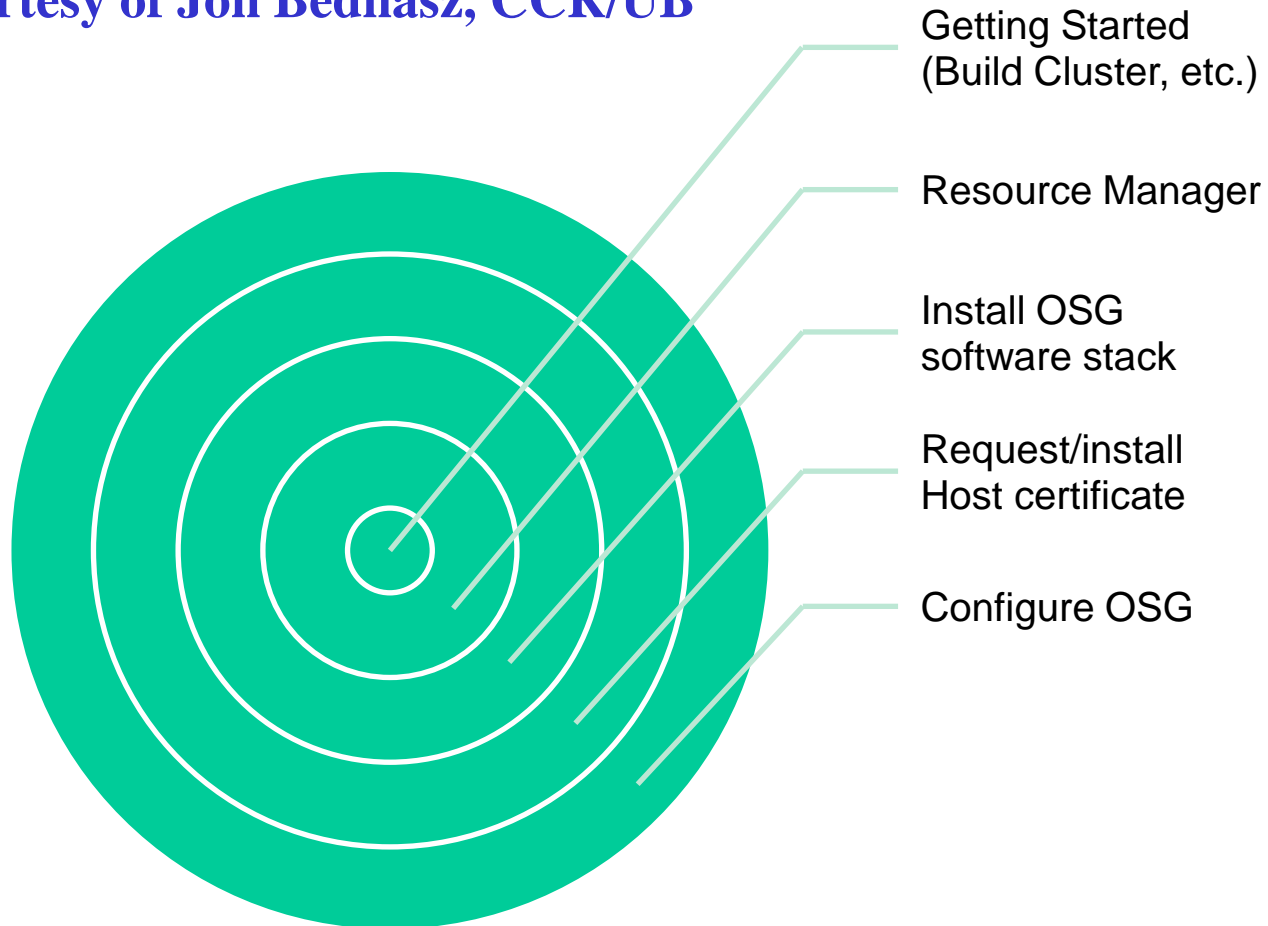
NYSGrid.org Organization



NSF Director Arden L. Bement: "leadership in cyberinfrastructure may determine America's continued ability to innovate – and thus our ability to compete successfully in the global arena."

NYS Grid Implementation Details

Figure Courtesy of Jon Bednasz, CCR/UB



Getting Started

(Courtesy of Jon Bednasz & Steve Gallo, CCR/UB)

- **Physically build a cluster**
 - ❑ 1 head node
 - ❑ 4+ compute nodes
- **Install Cluster Software**
 - ❑ Operating System (Red Hat)
 - ❑ Drivers for Interconnect (Myrinet, Infiniband, etc.)
 - ❑ Resource Manager (PBS, LSF, Condor, SGE)
- **Identify Gatekeeper Node for OSG Software**
 - ❑ Either stand alone machine or co-resident on Head Node
 - ❑ 5GB of space in /opt/grid
 - ❑ 5GB of space in /grid-tmp
- **Need to have ability to adjust firewalls**
- **Need to have ability to add users**

Installing OSG Stack on Gatekeeper

■ Installs are done via PACMAN

- ❑ `wget http://physics.bu.edu/pacman/sample_cache/tarballs/pacman-3.16.1.tar.gz`

■ Install OSG software

- ❑ `pacman -get OSG:ce`

■ Install (1) Package for your Resource Manager

- ❑ `pacman -get OSG:Globus-Condor-Setup`

- ❑ `pacman -get OSG:Globus-PBS-Setup`

- ❑ `pacman -get OSG:Globus-LSF-Setup`

- ❑ `pacman -get OSG:Globus-SGE-Setup`

NYSGrid.org Technical Group

- Jon Bednasz, Buffalo, Chair
- Steve Gallo, Buffalo
- Eric Warnke, Albany
- Steaphan Greene, Binghamton
- Ken Smith, Columbia
- Resa Alvord, Cornell
- Kirk Anne, Geneseo
- Steve Potter, Hauptman-Woodward
- Robert Schiaffino, Iona
- Earle Nietzel, Marist
- Ann Rensel, Niagara
- Chris Grim, NYU
- Rick Bohn, RIT
- Bill Webster, Rochester
- Lindsay Todd, RPI
- Ajay Gupta, Stony Brook
- Jorge González Outeiriño, Syracuse

NYSGrid.org Activities & Board

Activities

- **Technical Working Group**
- **Middleware**
- **User Support and Services / EOT**
- **Communications**
- **Infrastructure**
- **Resource Providers**
- **Funding**

Board

- **Russ Miller**
- **Gurcharan Khanna**
- **Linda Callahan**
- **Mark Shephard**
- **Tim Lance**
- **(Heather Stewart)**
- **Jim Davenport**
- **Chris Haile**

Technical WG Current Efforts

(Led by Steve Gallo and Jon Bednasz)

- **NYS Grid is Available**
- **OSG Jobs Running on NYS Grid**
- **CCR/UB & CTC/Cornell**
 - **Streamline bringing users onto NYS Grid**
 - **Documentation**
 - **Recommendations**
- **Need Early Adopters**
 - 1. Current Grid Users**
 - 2. New Users to Grid with HPC Needs**

Middleware WG Current Efforts

- Discussions on current state of Middleware at Buffalo, Binghamton, & RPI
 - Scheduling
 - Portals
 - Monitoring
 - Fault Tolerance
 - Checkpoint/Restart

CCR Outreach

- **HS Summer Workshops in Computational Science**
 - **Chemistry, Visualization, Bioinformatics**
 - **10-14 HS Students Participate Each Summer for 2 weeks**
 - **Project-Based Program**



CCR Outreach

■ Pilot HS Program in Computational Science

- Year long extracurricular activity at Mount St. Mary's, City Honors, and Orchard Park HS
- Produce next generation scientists and engineers
- Students learn Perl, SQL, Bioinformatics
- \$50,000 startup funding from Verizon, PC's from HP



Acknowledgments

- Mark Green
 - Cathy Ruby
 - Amin Ghadersohi
 - Naimesh Shah
 - Steve Gallo
 - Jason Rappleye
 - Jon Bednasz
 - Sam Guercio
 - Martins Innus
 - Cynthia Cornelius

 - George DeTitta
 - Herb Hauptman
 - Charles Weeks
 - Steve Potter
- Alan Rabideau
 - Igor Janckovic
 - Michael Sheridan
 - Abani Patra
 - Matt Jones

 - NSF ITR
 - NSF CRI
 - NSF MRI
 - NYS
 - CCR

