CCR CENTER FOR COMPUTATIONAL RESEARCH



- Enable Research at UB and in WNY
 - 90 Research Groups/25 Departments
- Industrial Outreach
- Education, Outreach & Training
 - High Schools and Middle Schools
 - Local Colleges
- Total Aggregate Computing Capacity
 - 13 Teraflops (13 Trillion operations/sec)
 - Can do in 1 day what a PC would take 6 years to do



CCR Computing Resources (13 Tflop)

Dell Linux Cluster (10TF)

- 1600 P4 Processors (3.2 GHz)
- Myrinet High-Speed Interconnect
- 2000 GB RAM; 60 TB Local Disk
- 30 TB SAN

• Dell Linux Cluster (3.0TF)

- 512 P4 Processors (3.0 GHz)

EMC SAN

- 35 TB Storage
- HP/Compaq SAN
 - 75 TB Disk; 190 TB Tape
 - 64 Alpha Processors (400 MHz)
 - 32 GB RAM; 400 GB Disk

SGI Altix3700 (0.4TF)

- 64 Processors (1.3GHz ITF2)
- 256 GB RAM,2.5 TB Disk

BioACE: Bioinformatics System

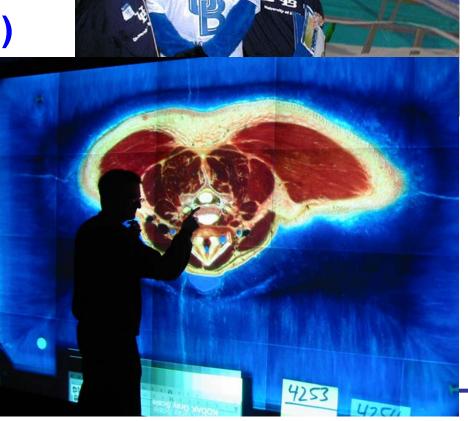
- Sun V880 (3), Sun 6800
- Sun 280R (2)
- Intel P4's
- Sun 3960: 7 TB Disk Storage



CCR Visualization Resources

Tiled-Display Wall

- 20 NEC projectors: 15.7M pixels
- Screen is 11'×7'
- Dell PCs with Myrinet2000
- Access Grid Nodes (2)
 - Group-to-Group Communication
 - Commodity components
- 3D Passive Stereo Display
 - VisDuo ceiling mounted system



Groundwater Flow Modeling

- Regional scale modeling of groundwater flow and contaminant transport (Great Lakes)
- Ability to include all hydrogeologic features as independent objects

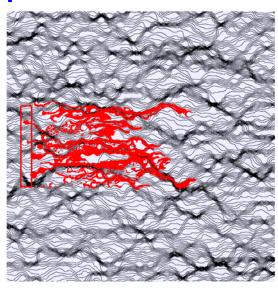
Based on Analytic Element Method

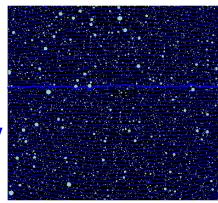


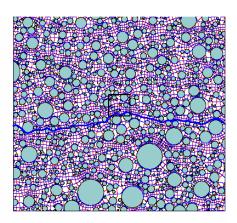


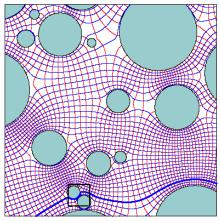
Groundwater Dispersion

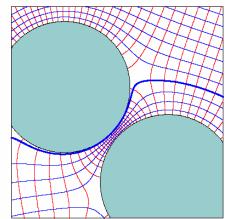
- Basic research on contaminant spreading in aquifers
- Collaboration with leading scientists from Israel and Italy
- Designed to improve characterization of existing and future contaminant plumes







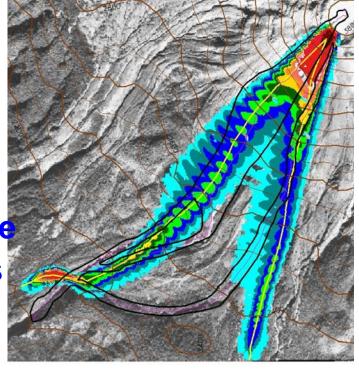




Analytic and numerical modeling with up to 10 billion cells in 2D and 3D

Avalanches, Volcanic and Mud Flows

- 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

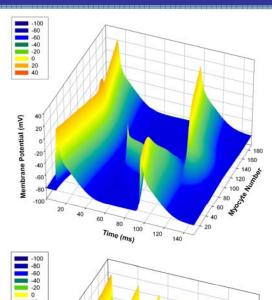


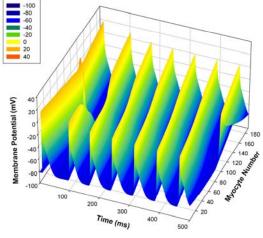
Flow models of Colima volcano In Mexico – courtesy Rupp et. al.'06

Cardiac Arrhythmia

Center for Cellular and Systems Electrophysiology

- 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



Cardiovascular Research



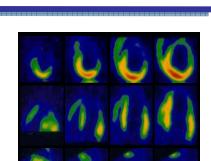
Molecular Imaging – PAREPET Clinical Study

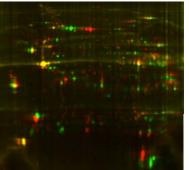
Analysis of cardiac PET (Positron Emission Tomography) scans aims to revolutionize assessment of an individual's risk for sudden cardiac death.

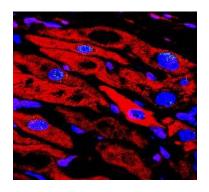


Protein and gene expression profiling using differential in-gel electrophoresis and microarray technology provides a blueprint for the cellular mechanisms involved in hibernating myocardium.

Translate results to identify gene and other therapeutic targets aimed at improving heart function and survival.

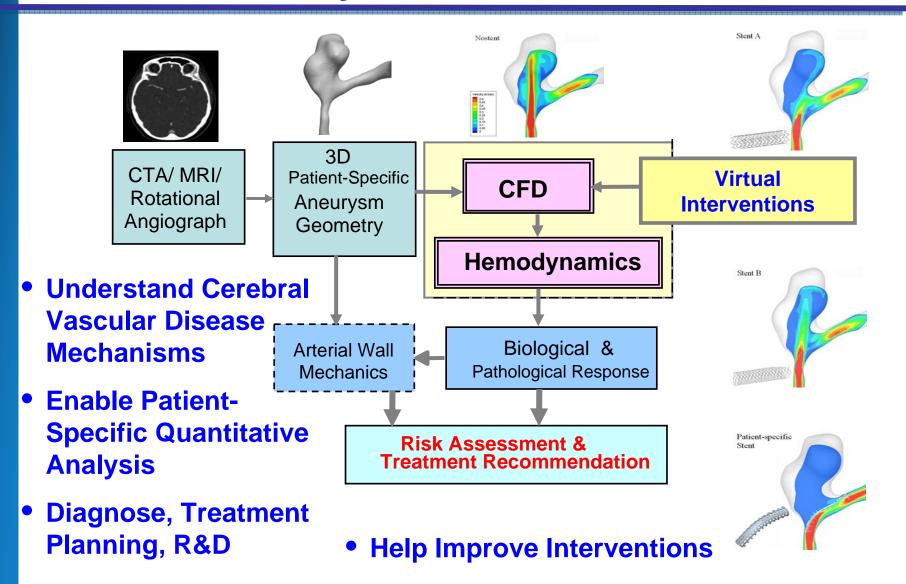








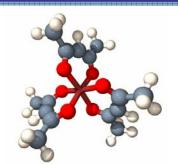
Cerebral Aneurysm: Virtual Intervention

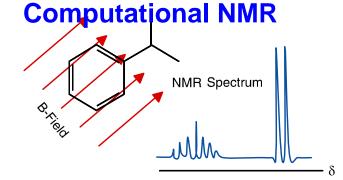


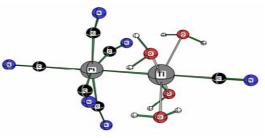
Theoretical and Computational Chemistry

Applied to:

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



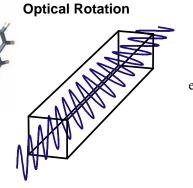


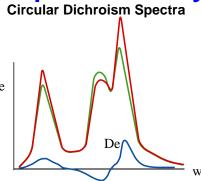


Magnetic Properties of Molecules

Software Development

Theory and Calculation of Optical Activity





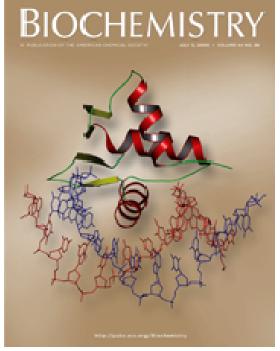
J. Autschbach

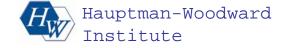
3D Structure of Proteins

- 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





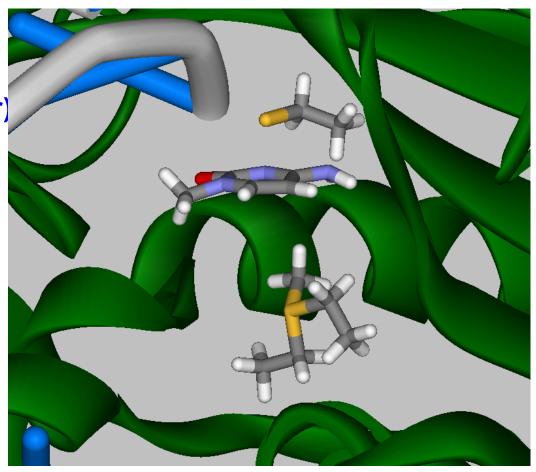
Understanding How Proteins work

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.



Computational Biology & Bioinformatics

Development of Bioinformatic Tools

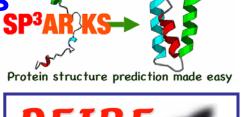
SPEM – align multiple sequences for discovering hidden evolution information of genes.

SPARKS/SP³ – predict threedimensional structures of proteins by matching a query sequence with known structural templates.

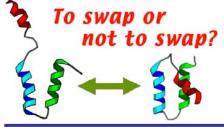
DFIRE – predict binding affinities of protein-protein, protein-ligand, and protein-DNA complexes for structure-based drug design.

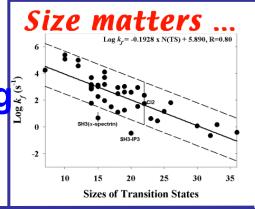
Mechanistic study of protein folding
 and binding

http://theory.med.buffalo.edu









Computational Chemistry

- UB Software development in Quantum Chemistry
 - Q-Chem development of combined QM/MM methods for large molecular systems such as proteins

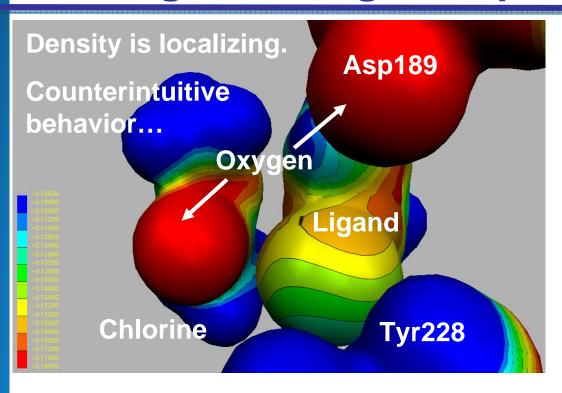
ADF – development of algorithms to calculate magnetic and

optical properties of molecules

- Used to determine
 - 3D Molecular Structure
 - Electronic Spectra
 - Chemical Reactivity
- Applications
 - Pharmaceutical Drug Design
 - Industrial Catalysis
 - Materials Science
 - Nanotechnology

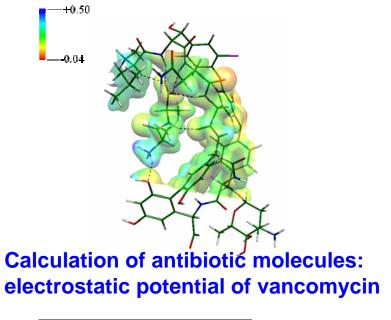


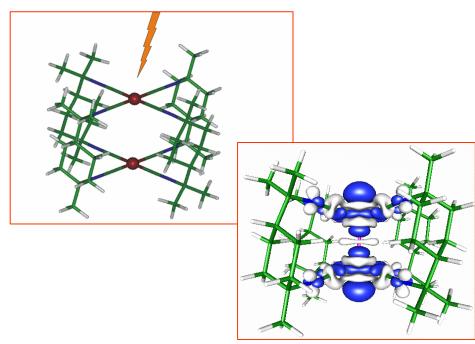
Binding in a Drug-Receptor Complex

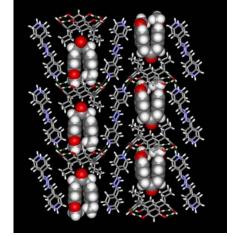


- Ligand docked with residues of the active site of thrombin. Electrostatic potential map superimposed onto the electron density isosurface.
- The goal is to elucidate the thermodynamics of molecular recognition in binding.

Understanding Large Molecules and Fleeting Species





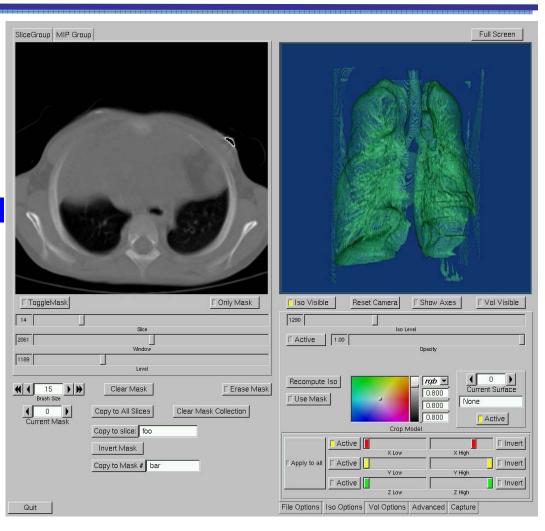


A molecule changes on excitation by light

A supramolecular solid

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

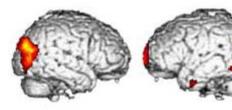


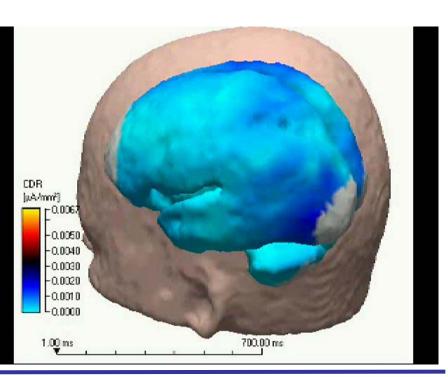
Mapping Brain Activity

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

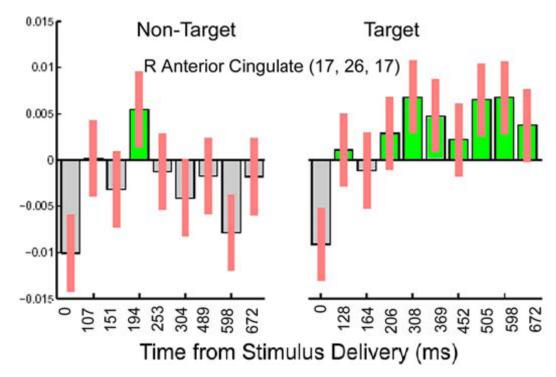
Current density maps of brain surface (1–700 ms after target) show dynamic pattern of brain activity during decision-making process.

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.

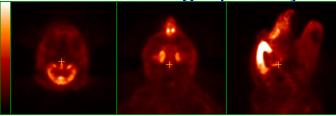
Nuclear Medicine

Monte Carlo simulation

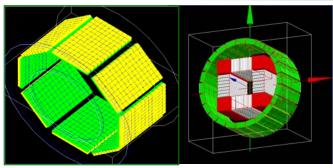
 for modeling imaging system characteristics, optimizing system design, and validating data correction algorithms.

Image reconstruction

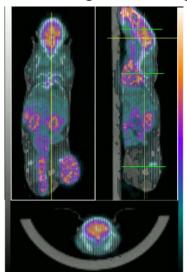
 for development of high resolution image reconstruction algorithm and software for both human and animal nuclear emission tomographic systems.

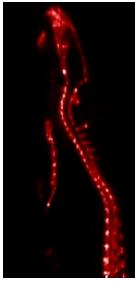


Transverse, coronal and sagittal views of a monkey brain scanned on a dedicated brain PET using the radioligand ¹⁸F-FCWAY.



Two virtual imagers simulated for system modeling and design evaluation.





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

Vascular Dementia Imaging

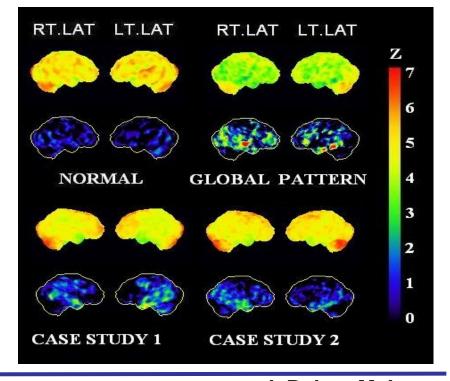
- Early diagnosis of dementia from cerebral small vessel disease using computer analysis of SPECT Images
- Collaboration between Nuclear Medicine, CCR, Neurology, and Kaleida Stroke Center
- Funded by the Pfeiffer Foundation
- Fractal scores:

_	Normal	0.75

Global Pattern 1.13

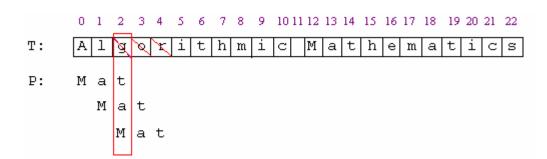
Case Study 1 0.96

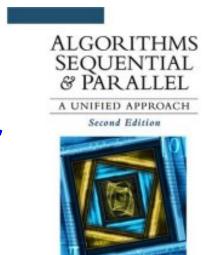
Case Study 1 Moderate
 white matter and cortical
 hypo-perfusion with visual
 memory, speed of
 processing, and verbal
 fluency deficits

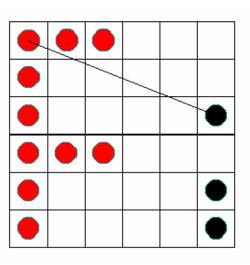


Parallel Algorithms

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







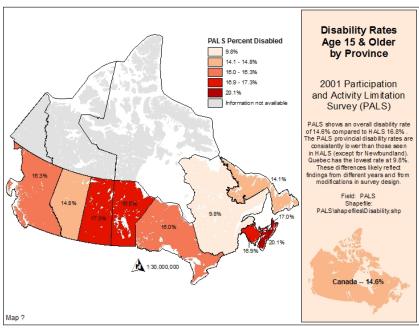
Collaborative Visualization Environments

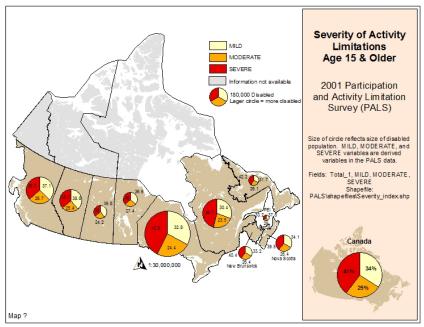
- Enable distributed collaboration via software developed at CCR
- Enable visualization and interaction with data across a geographically disparate network topology
- Integrate multiple data sources:
 - Scientific, Multimedia
- Research topics:
 - Distributed databases
 - OpenGL 3D programming
 - 3D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



Literacy & Disability in Canada

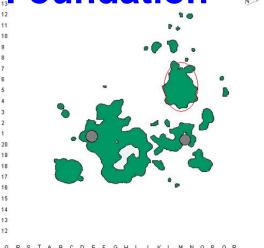
- Exploring the relationship between illiteracy & disability across the Canadian landscape
- Social Systems GIS Lab in the Dept. of Anthropology is working with researchers from York University & the Canadian Abilities Foundation.
- Sponsored by The Adult Learning & Literacy Directorate of the Ministry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

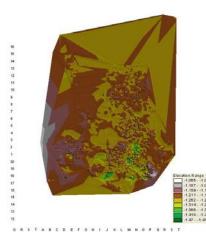


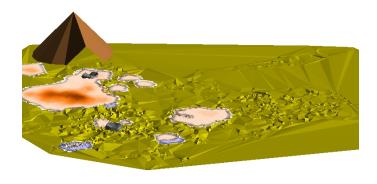


Verberie Paleolithic Site in France

- 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

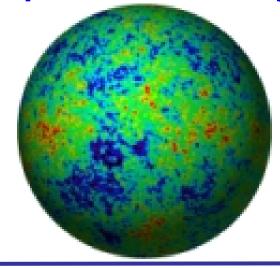


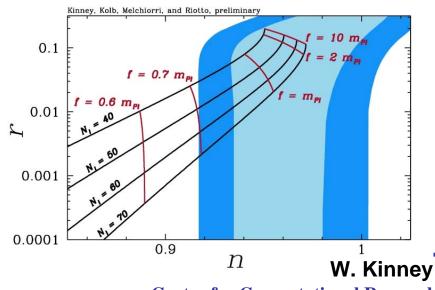




Cosmological Parameter Estimation

- 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





- World Class Center for Earthquake Engineering
- Flagship lab for the (National) Multidisciplinary Center for Earthquake Engineering Research (N)MCEER since 1986
- Upgraded in 2000-2004 under the NSF George E. Brown Network for Earthquake Engineering Simulation (NEES): (\$21.2M)
- Commissioned in 2004 and funded through 2014 under NSF's 10-Year "Management, Operation and Management" Award (\$15M)

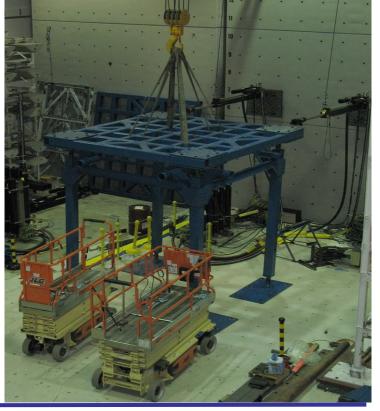




Equipment

- Twin 6-DOF shake tables (50 m-ton capacity); 7mx7m platforms,
- Three 100-ton dynamic actuators and two 200-ton static actuators
- Non-structural component Simulator (NCS)
- 2-D Geotechnical Laminar Box







Projects



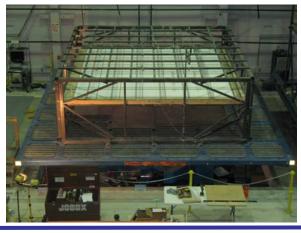
Seismic Qualification and Fragility Tests of a 230KV-3 Phase Switch Assembly



Seismic Fragility Tests of a Centrifugal Liquid Chiller and Restraint System



Dynamic Seismic Response of Steel Braced Frames



Suspended Ceiling Tile Tests



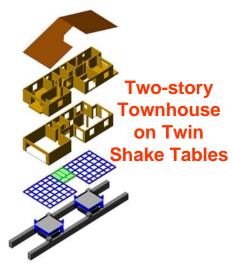
Ridge-U-Rak Inventory Storage System Tests

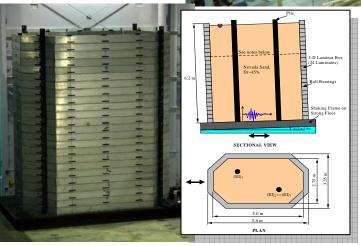


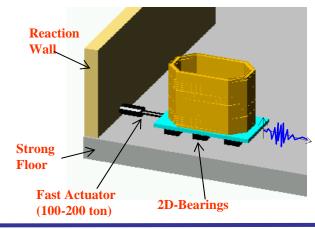
Projects

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





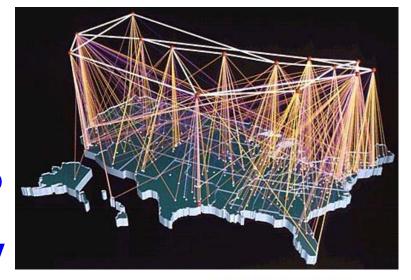




2-D
Geotechnical
Laminar Box
Tests of Pile
Foundations
Subjected to
Soil
Liquefaction

Cyberinfrastructure in STEM Education

- Developing a scalable, multi-site cyberinfrastructure 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





Determining 3D Protein Structure

- NMR-based Structural Biology and
 Structural Genomics
 TITELY
- Bio-NMR Methodology

HIGH TECHNOLOGY

Breakthrough at UB

 NMR-based Metabonomics in Cancer Research

Structural Biology

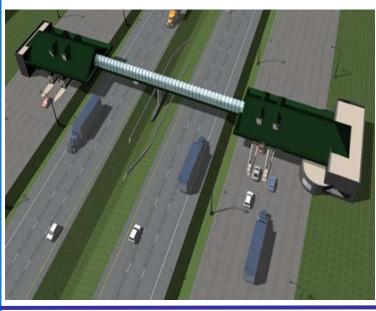


Propelled by Recent Advances, NMR Moves Into the Fast Lane

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

Urban Modeling & Visualization

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







CCR Model Development

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

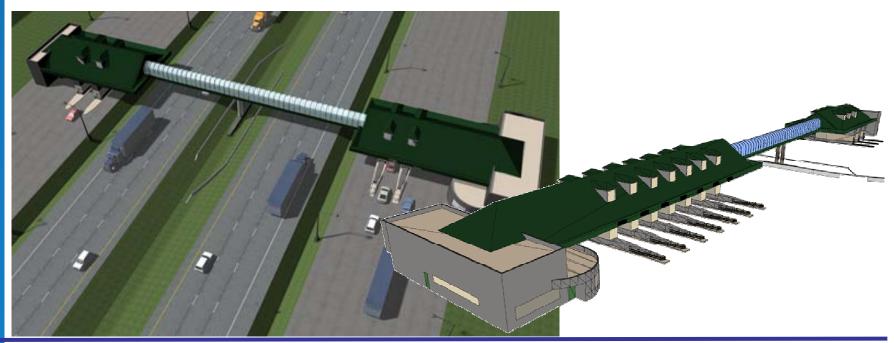
packages

- Corsim
- Synchro
- Vissim



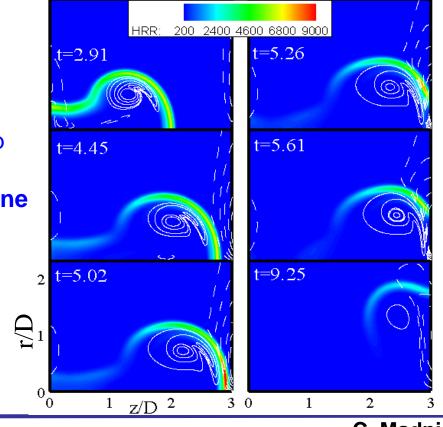
Urban Modeling & Visualization

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



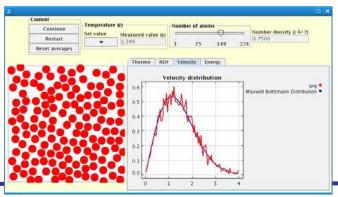
Understanding Combustion

- Flame-wall interaction modeling for a non-premixed flame propelled by a vortex ring. CFD lab, SUNY, Buffalo.
- 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 GRI3.0
 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

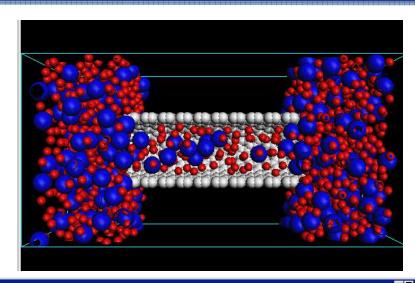


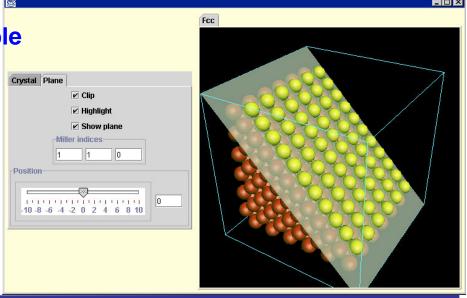
Molecular Simulation Software

- 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



University at Buffalo
The State University of New York



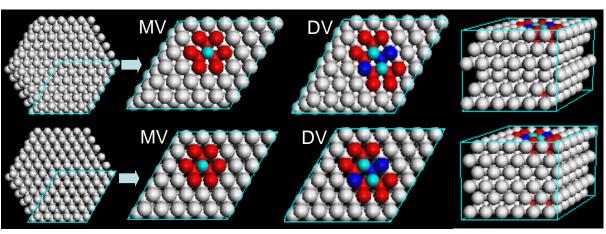


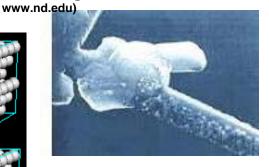
Computational Materials Science

- 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



hillocks (below) due to electromigration. (source:

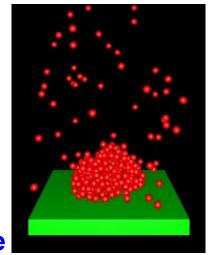




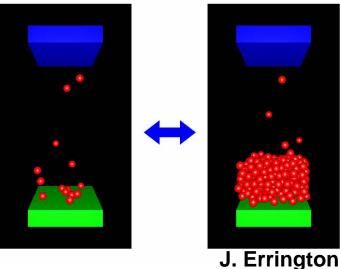
Simulation cells of solids with mono- and di-vacancies (light blue spheres), highlighting atoms neighboring the defects.

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



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 Å of a trehalose molecule



UB/WNY Biomedical Advances

- PSA Test for Prostate Cancer
- Avonex: Interferon Treatment for Multiple Sclerosis
- Artificial Blood
- Nicorette Gum
- Fetal Viability Test
- Implantable Pacemaker
- Edible Vaccine for Hepatitis
 C
- Timed-Release Insulin Therapy
- Anti-Arrythmia Therapy
 - Tarantula venom

- Direct Methods for Crystal Structure Determination
 - Listed on "Top Ten Algorithms of the 20th Century"
 - Vancomycin, Gramacidin A
- Patented High Throughput Crystallization Method
- NIH National Genomics Center: Northeast Consortium
- Howard Hughes Medical Institute: Center for Genomics & Proteomics



Genome-wide Study of Iron Homeostasis

 Physiological and Hepcidin-hormonal regulators are involved in iron homeostasis

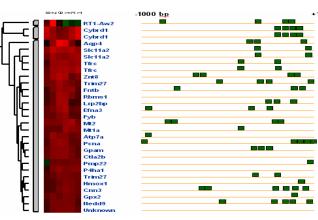
Intestinal iron transport controls overall body iron homeostasis

• 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

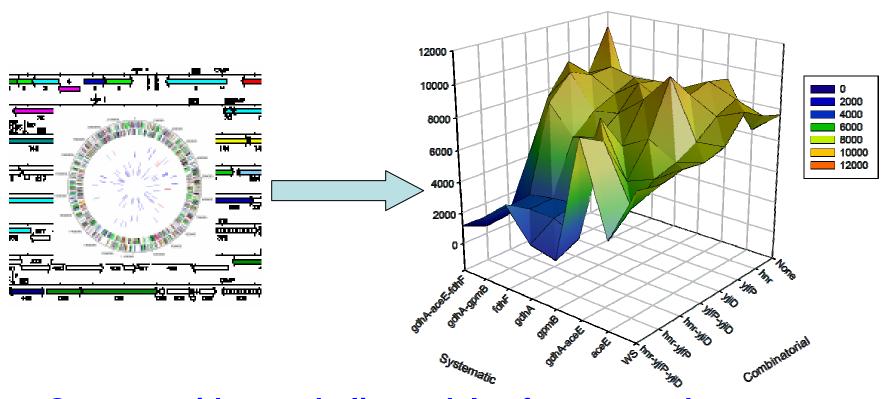


Hepcidin

(Fe3+)2 - Tf

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

Designing Cellular Phenotypes

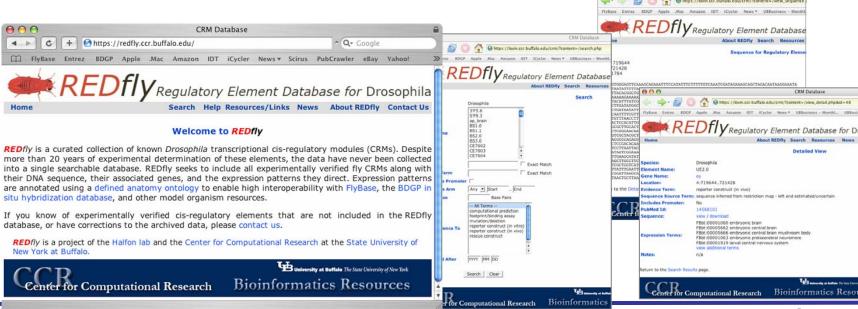


- Genome-wide metabolic models of sequenced microorganisms
- Optimization of metabolic and cellular phenotypes
- Goal is to design biocatalysts for the production of pharmaceuticals and high-value chemicals



Regulation of Gene Expression

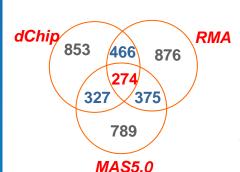
- REDfly (Regulatory Element Database for Fly) Database of verified transcriptional regulatory elements
- Over 650 entries
- Most comprehensive resource of animal regulatory elements
- Fully searchable, has DNA sequence and gene expression data, link-outs to other databases





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



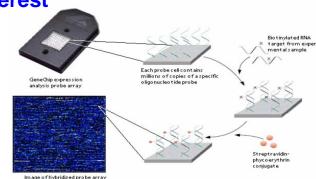
31% - 55% overlap

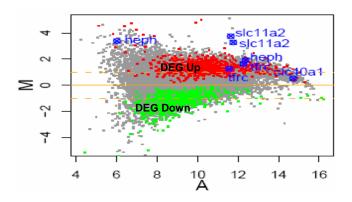
Which algorithm is best?

Great impact on subsequent expression data analysis



- No external reference data needed
- Algorithm evaluation with direct applications to experimental datasets of interest





BMC Bioinformatics

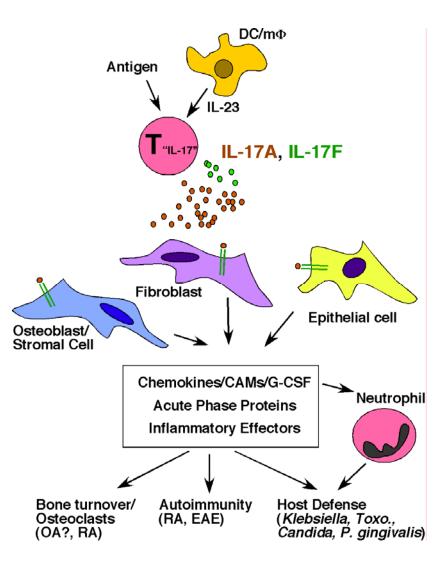


Methodology article

Open Access

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

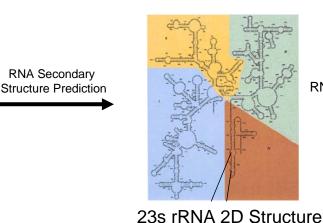
Defining Cytokine Signaling Mechanisms

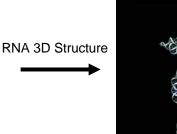


- -T cells secrete cytokines such as IL-17 to promote host defense and/or autoimmunity
- -Microarrays used to define IL-17 gene targets in various cell types
- -Computational and statistical approaches used to compare the promoters of IL-17 target genes in mouse and human genomes to identify conserved 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-kB and C/EBP

Prediction of RNA Structure to Facilitate Design of Drugs Targeting RNA

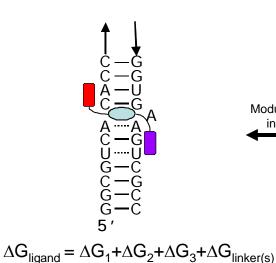
aaaaacaca gcactgtgca aacacgaaag tggacgtata cggtgtgacg cctgcccggt gccggaaggt taattgatgg ggttagcgca agcgaagctc ttgatcgaag ccccggtaa cggcggccgt aactataacg gtcctaaggt agcgaaattc cttgtcgggt aagttccgac ctgcacgaat ggcgtaatga tggccaggct gtctccaccc gagactcagt gaaattgaac agettgacae tgaacattga geettgatgt gtaggatagg tgggaggett tgaagtgtgg acgccagtct gcatggagcc gaccttgaaa taccaccctt taatgtttga tgttctaacg tectaaagag taacggagga geacgaaggt togetaatee togteggaea teaggaggt agtgcaatgg cataagccag cttgactgcg agcgtgacgg cgcgagcagg tgcgaaagca ggtcatagtg atccggtggt tctgaatgga agggccatcg ctcaacggat aaaaggtact ccggggataa caggctgata ccgcccaaga gttcatatcg acggcggtgt ttggcacct gatgtcggct catcacatcc tggggctgaa gtaggtccca agggtatggc tgttcgccat ttaaagtggt acgcgagctg ggtttagaac gtcgtgagac agttcggtcc ctatctgccg tgggcgctgg agaactgagg ggggctgctc ctagtacgag aggaccggag tggacgc actggtgttc gggttgtcat gccaatggca ctgcccggta gctaaatgcg gaagagataa gtgctgaaag catctaagca cgaaacttgc cccgagatga gttctccctg accctttaag ggtcctgaag gaacgttgaa gacgacgacg ttgataggcc gggtgtgtaa gcgcagcc gcgttgagct aaccggtact aatgaaccgt gaggcttaac ctt

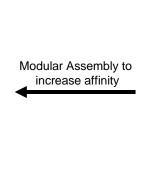


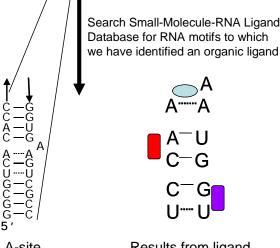




23s rRNA Sequence





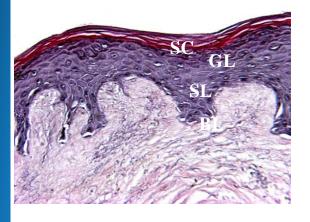


A-site
Aminoglycoside target

Results from ligand database search

Stem Cells and Tissue Engineering

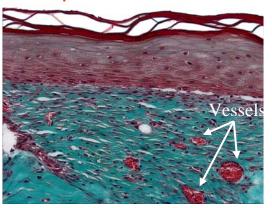
Bioengineered Skin



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



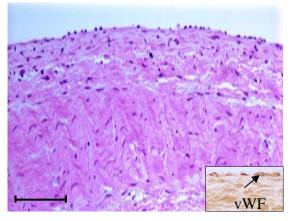
Bioengineered Blood Vessel



Tissue engineered blood vessels (TEV):

- 1. Bypass surgeries
- 2. Model to study mechano transduction and vascular biology
- 3. TEVs from bone marrow stem cells

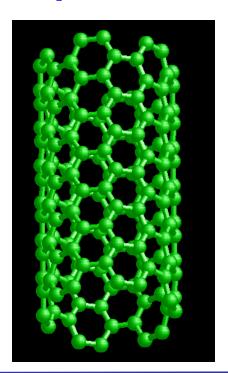
TEV Transplanted in jugular vein of lambs

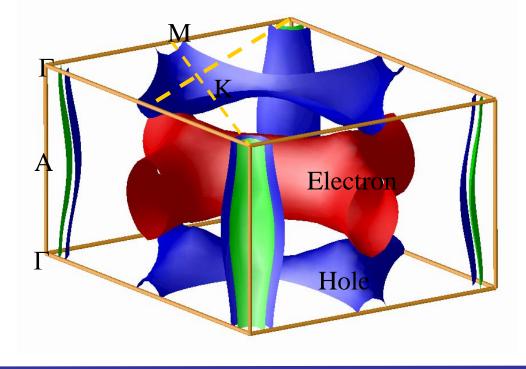




Designing new materials

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

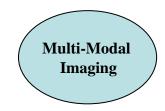


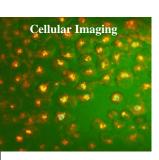


Nanomedicine Program

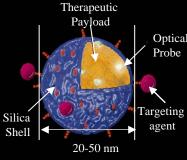
World class Research Program Melding Nanotechnology with Biomedical Sciences



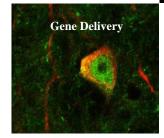


















Building from the Bottom Up

State of the Art Molecular Imaging and Nanocharacterization Facilities

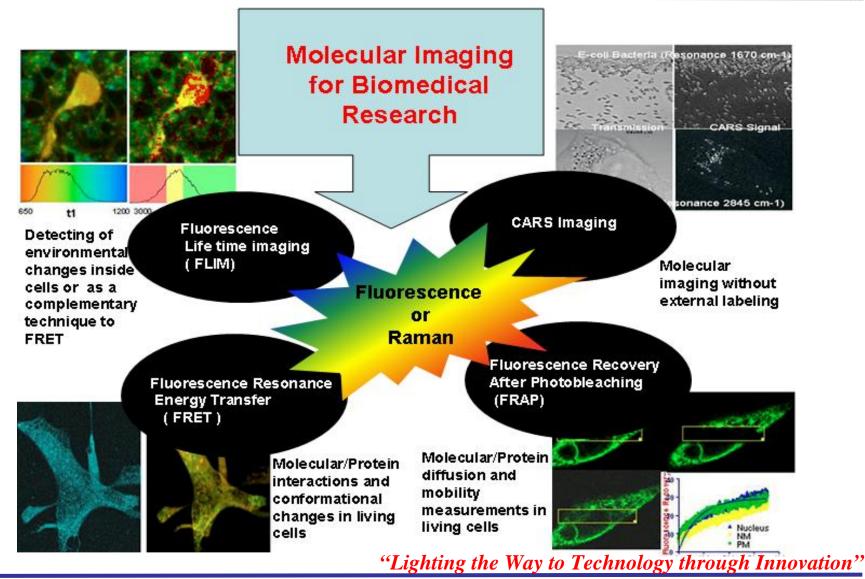
- Multiphoton Laser Scanning System
- Confocal Imaging including FRET, FLIM & FRAP analysis
- Coherent Anti-Stokes Raman Imaging
- Optical Trapping/Dissection
- Advanced Laser Systems

"Leading the Way to Technology through Innovation"



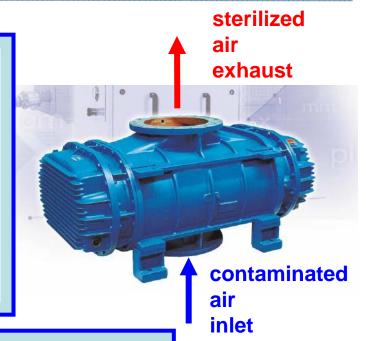


Molecular Imaging



Complete Air Sterilization





compressive heating of flowing air destroys ANY & ALL airborne pathogens, 99.9999% killed in a single pass through the BioBlower air pump

Can completely protect any enclosed area (plane, train, bus, building) from:

anthrax spores, smallpox, SARS, MARS, influenza, tuberculosis, avian flu, mold, fungi, allergens, etc

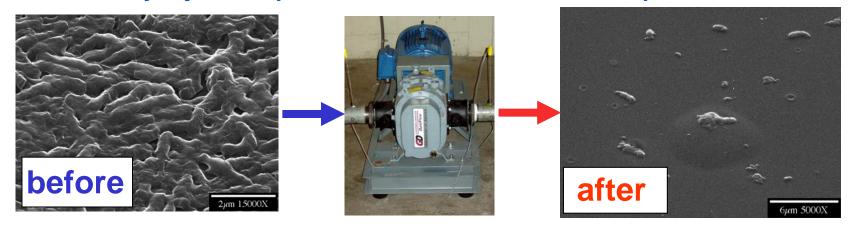


Complete Air Sterilization



compressive heating of flowing air destroys ANY & ALL airborne pathogens, 99.9999% killed in a single pass through the BioBlower air pump

heat treated *Bacillus globigii* (*Bg*) spores, an anthrax surrogate, are continuously injected upstream of BioBlower, then sampled downstream



"Everything from hospitals, first-responder units, and postal facilities -- to government buildings and mass-transit systems could benefit enormously from the security and peace of mind generated by this device" Rep. L. Slaughter, Business First, 8/16/05

Industrial 3D Flow Analysis

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

