



Biomedical Informatics Research Network
a testbed for a biomedical knowledge infrastructure

BIRN Overview

- Federated database of neuroimaging data
- Fusion of diverse data sources
- Grid access to computational resources
- Datamining
- Scalable, to extend to all NCRR Centers for a full range of biomedical studies

Challenges

- Large data sets
 - ISCAR - information storage, curation, archiving, and retrieval
 - Bandwidth issues
 - Quality control of data only analyzed by machine (audit requirements)
- Data sharing issues (privacy, etc.)

More Challenges

- Federated DB
 - Integration of disparate sources
 - Accessing legacy systems
 - Knowledge distillation
- Project Coordination
 - Encouraging collaboration
 - Balance centralization vs distribution
 - Role of standards
 - Sociology (ie., credit, ownership, intellectual property)
 - Project subdivision and optimization

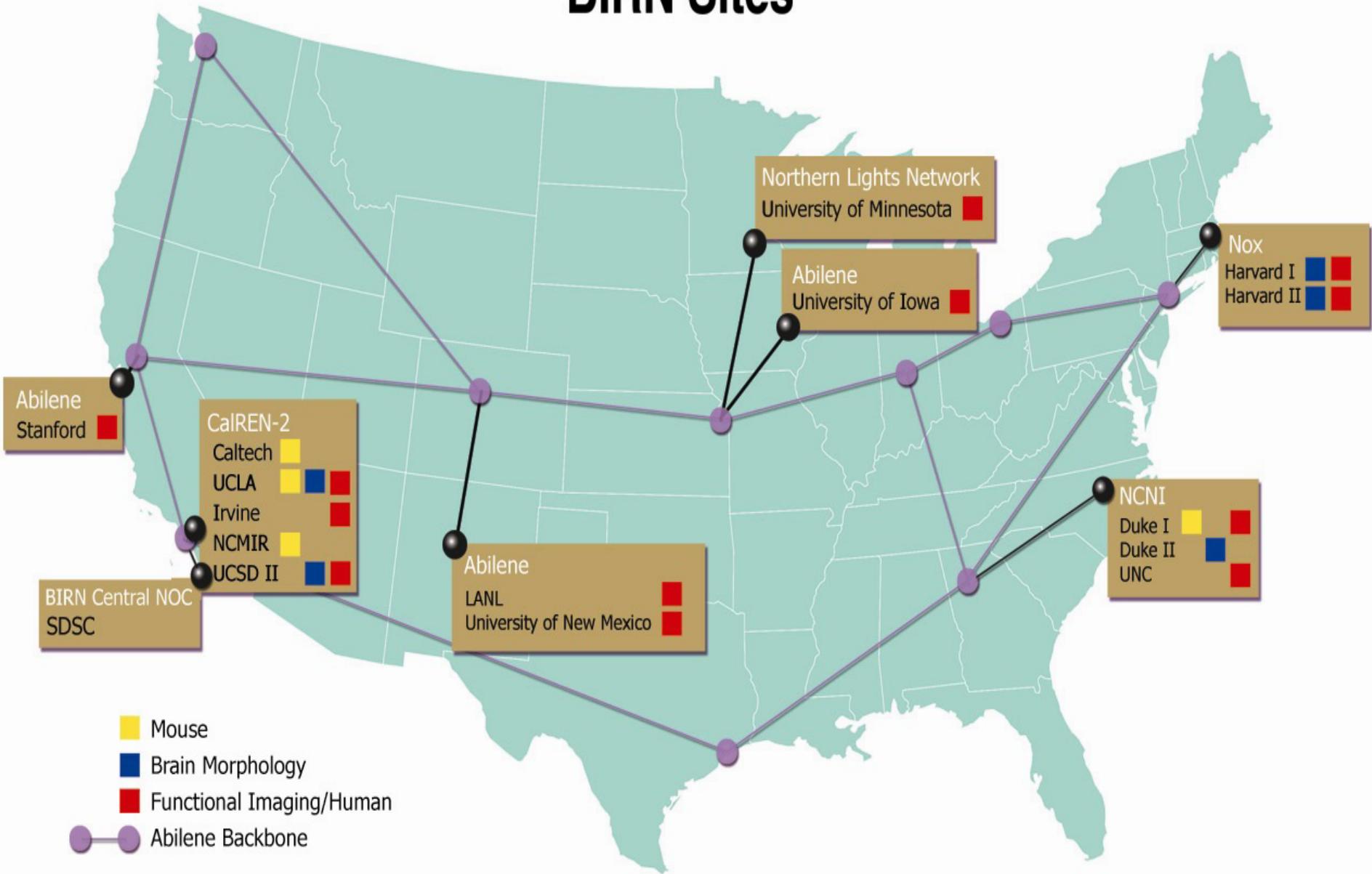
BIRN Project Objectives:

- 1) establish a stable high performance network linking key NIH / NCRR-supported Biotechnology Research Resources (P41) and General Clinical Research Center (GCRC) sites - using Internet 2 & beyond
- 2) establish distributed and linked data collections for the test bed projects
- 3) enable distributed heterogeneous "grid-based" computing resources
- 4) enable data mining from multiple data collections
- 5) develop a stable software and hardware infrastructure that will allow centers to accumulate larger studies than can be carried out at one site.

BIRN Phases

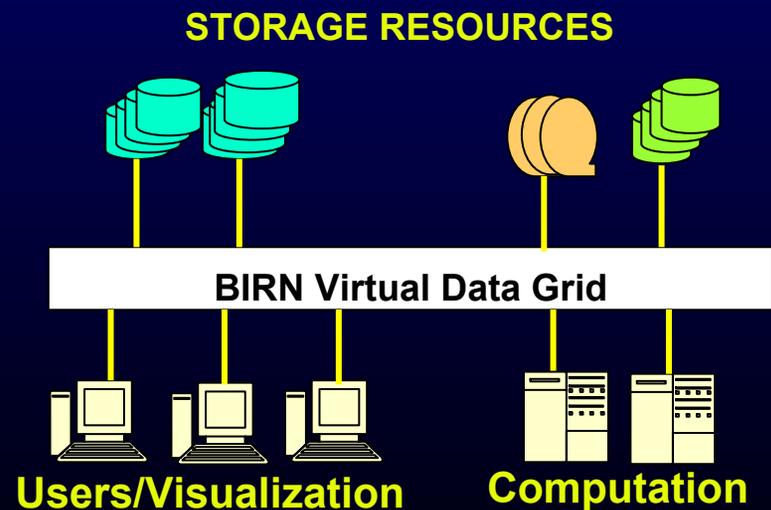
- Phase I — neuroimaging federation
 - Mouse BIRN
 - Morph BIRN
 - FIRST BIRN
- Phase II — Neuroscience, including physiology, genetics, proteomics
- Phase III— Biomedical Research with broad participation of 25 NIH funding Institutes

BIRN Sites



BIRN Virtual Data Grid w/SRB inside

- Defines a Distributed Data Handling System
- Integrates Storage Resources in the BIRN network
- Integrates Access to Data, to Computational and Visualization Resources
- Acts as a Virtual Platform for Knowledge-based Data Integration Activities
- Provides a Uniform Interface to Users



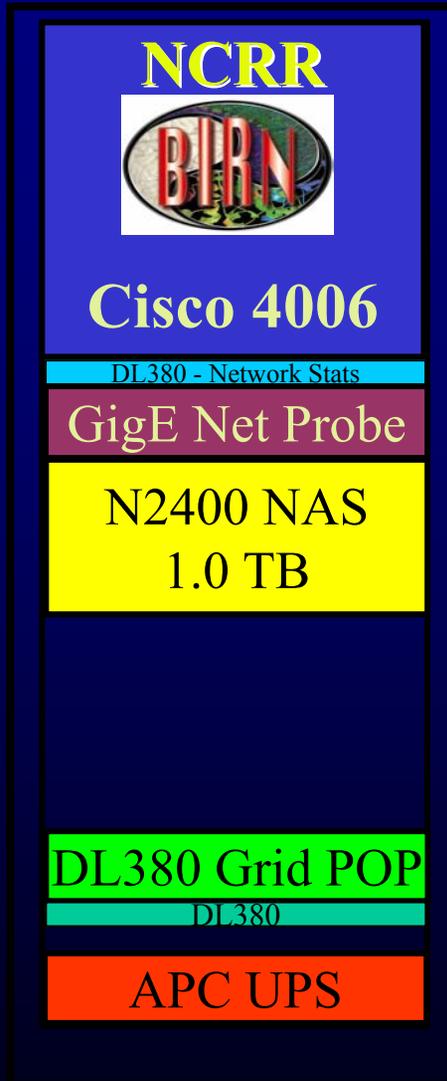
Integrating Across Data Sources

- The idea
 - Different models capture correlated but distinct aspects of biological reality
 - How can we express and evaluate queries that compute data across models
- The approach
 - For each source create a knowledge-base of the anatomy of the observations
 - Attach the data of each source to the respective knowledge-base
 - “bridge” the sources by a simple ontological mapping
 - Compute the query across bridged sources

Getting down to work - BIRN must:

- 1) establish Network Infrastructure..
- 2) establish policies and protocols for use of the BIRN infrastructure and operating guidelines.
- 3) build a Virtual Data Grid (VDG) for the initial projects.
- 4) develop the data modeling and mediation tools for complex queries.

Uniform Hardware for BIRN Sites



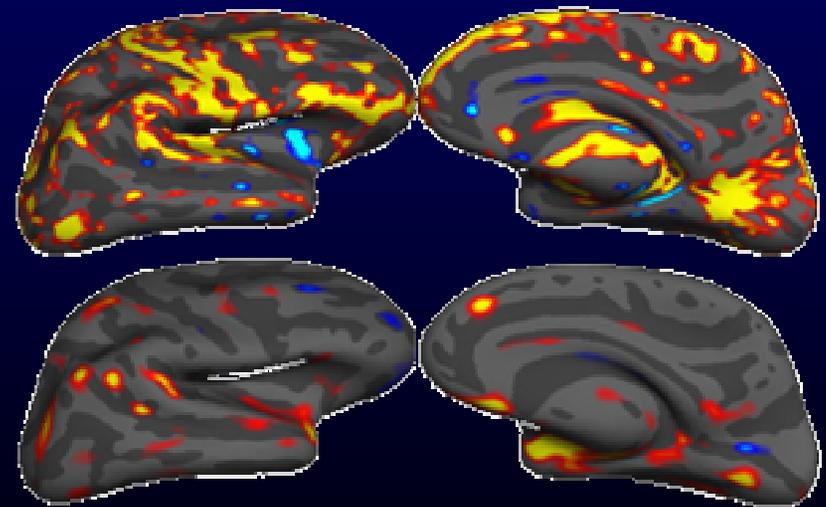
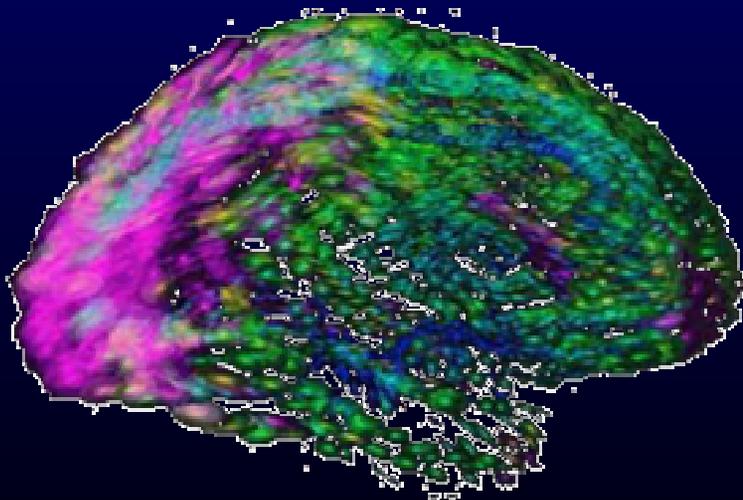
- Gigabit/10/100 Network Switch – Cisco4006
- Network Statistics System
- Gigabit Ethernet Network Probe
- Network Attached Storage – Gigabit Ether
 - 1.0 to 8.0 TB
- Grid POP
 - SRB, Globus
 - Dual Processor Linux w/ 1GB memory
- DL 380 General purpose
 - Specialized encryption
- UPS for Rack

Two active BIRN project partnerships, so far:

- **Mouse BIRN** - *Animal Models of Disease / Multi Scale/Multi Method - MS Mouse and DAT KOM (a schizophrenic and otherwise interesting mouse animal model)*
- **Brain Morphology BIRN** - *Neuroanatomical correlates of neuropsychiatric illness (Unipolar Depression, mild Alzheimer's Disease (AD), mild cognitive impairment (MCI))*

Morphology **BIRN**

- Combining data from multiple acquisition sites
- Increasing the statistical power for studying relatively rare populations
- Harvard (MGH and BWH), Duke, UCLA, UC San Diego



Brain Morphology BIRN

Institutional Participants:

- Center for Neuroimaging Technologies, Harvard (Lead Institution)
Bruce Rosen, PI
- Neuroimaging Analysis Center, Harvard
Ron Kikinis PI
- Laboratory of Neuro Imaging, University of California Los Angeles
Arthur Toga, PI
- GCRC @ Duke University, Durham NC
Ranga Krishnan, PI
- GCRC @ University of California San Diego
Ed Holmes, PI

Brain Morphometry BIRN

- Clinical Specific Aims

Attempt to address questions of the following type:

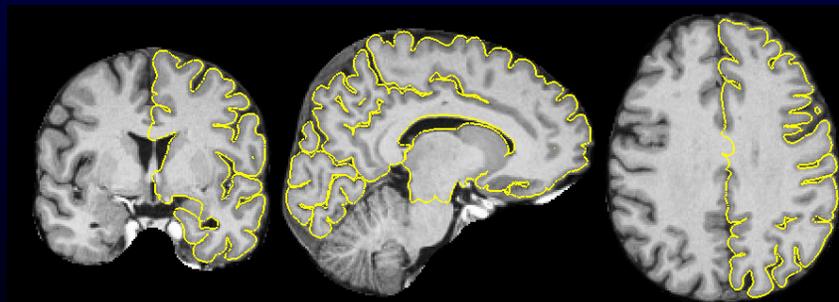
- Do structural differences contribute to specific symptoms such as memory dysfunction or depression independent of diagnosis?
- Do specific structural differences distinguish specific diagnostic categories?

- Technological Specific Aims

Attempt to overcome obstacles to the use of neuroimaging data as quantitative outcome measures for clinical investigation including the issues raised by longitudinal and multi-site studies. In other words- to make the Clinical Specific Aims possible to address.

Harvard

UCLA



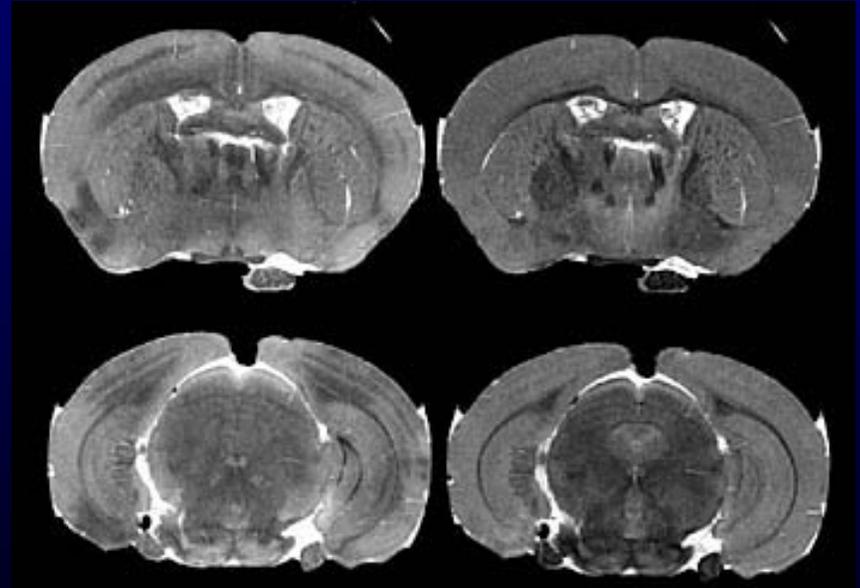
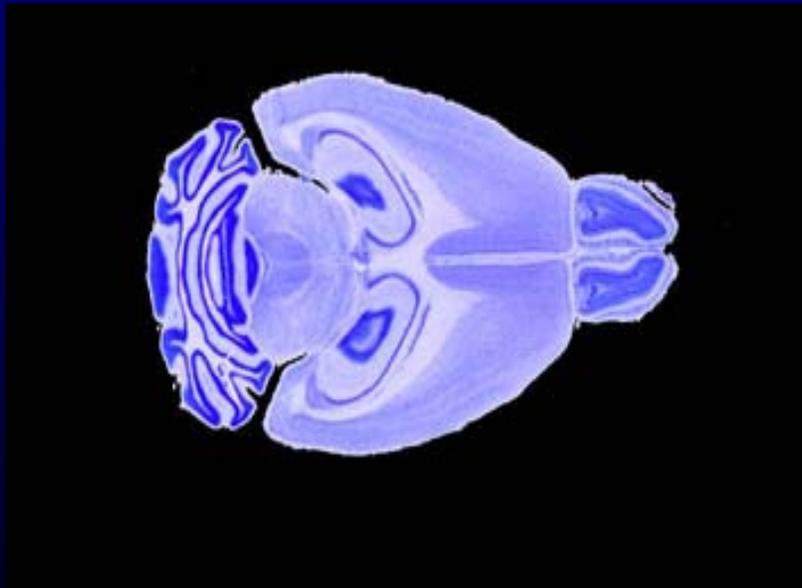
Duke

UCSD

NCRR

Mouse BIRN

- Looking at disease models within and across individuals
- Looking at different resolutions by combining data from multiple modalities

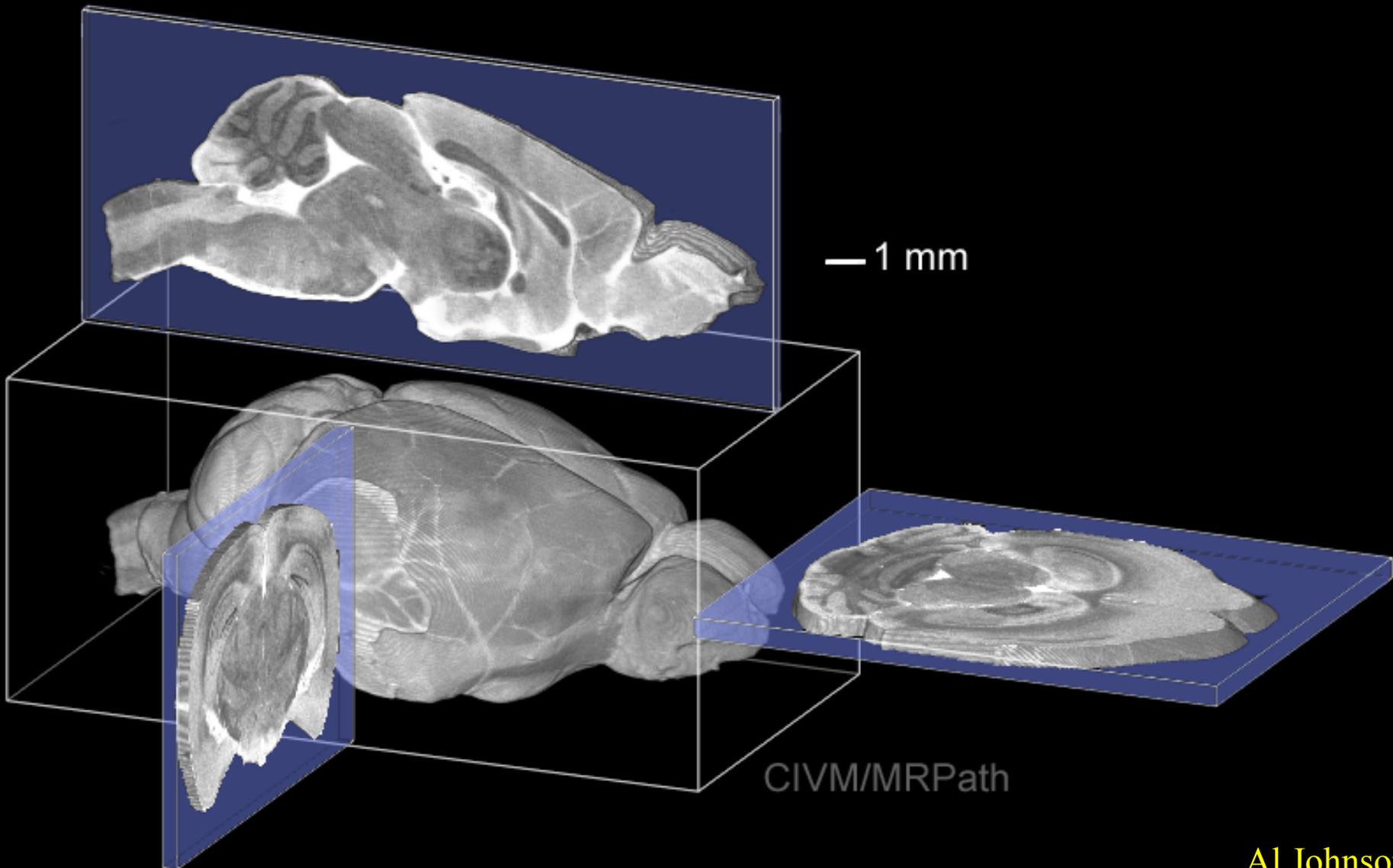


Duke, UCLA, UC San Diego, Cal Tech

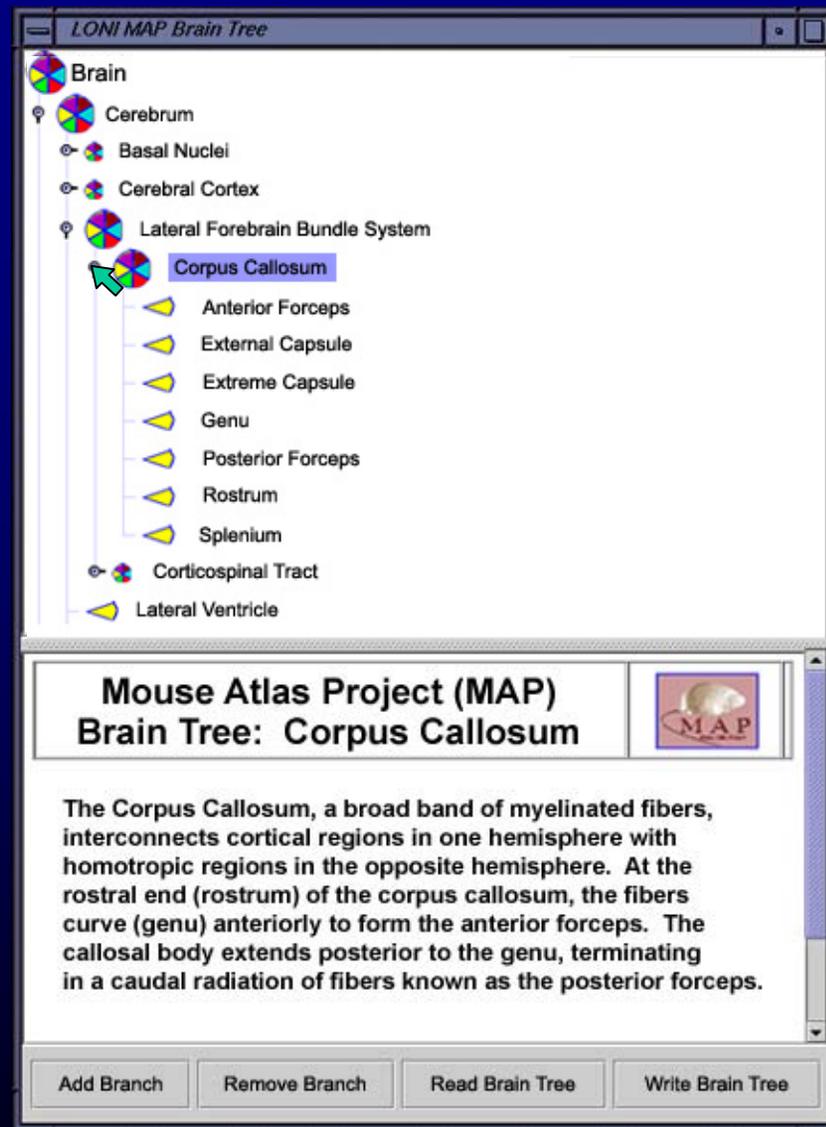
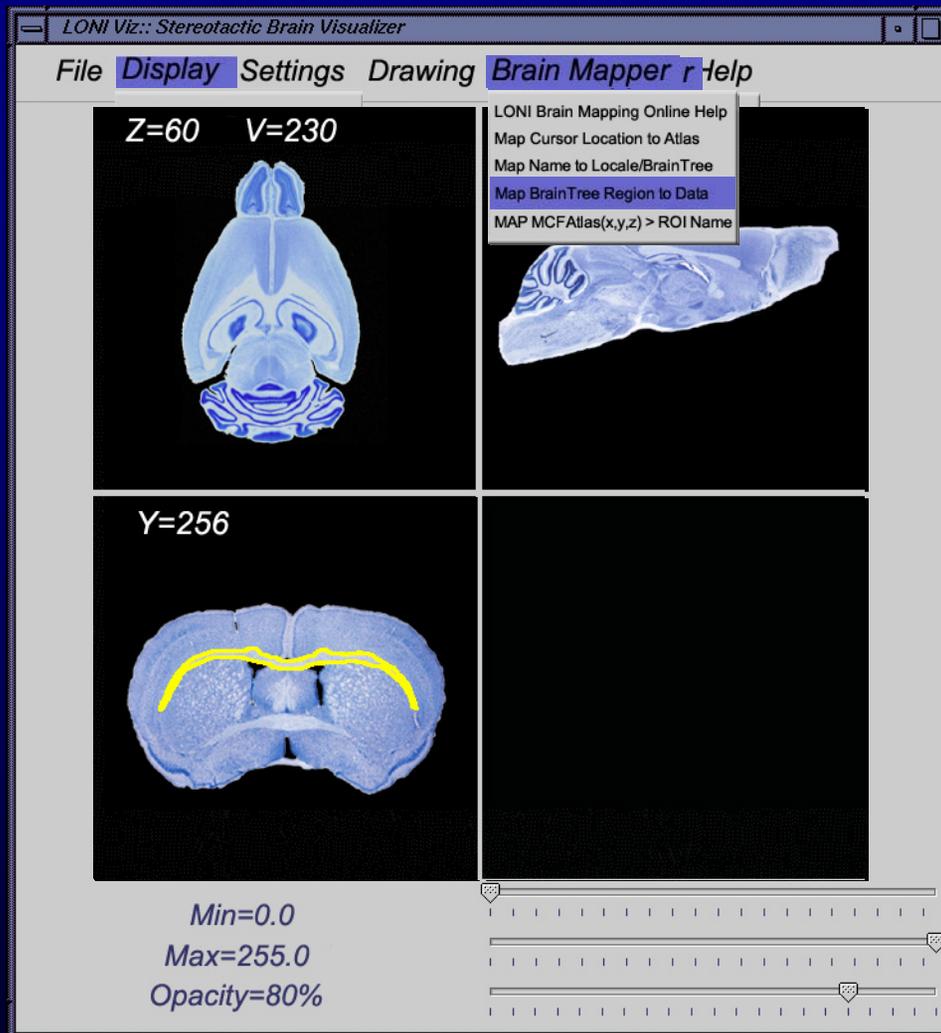
Mouse BIRN Project

Institutional Participants:

- Center for In Vivo Microscopy, Duke Univ. (Lead Institution)
Allen Johnson, PI
- Laboratory of Neuro Imaging, University of California Los Angeles
Arthur Toga, PI
- MRI Center @ Caltech's Beckman Institute, CalTech
Russell Jacobs, PI
- National Center for Microscopy and Imaging Research, UCSD
Mark Ellisman, PI



Al Johnson

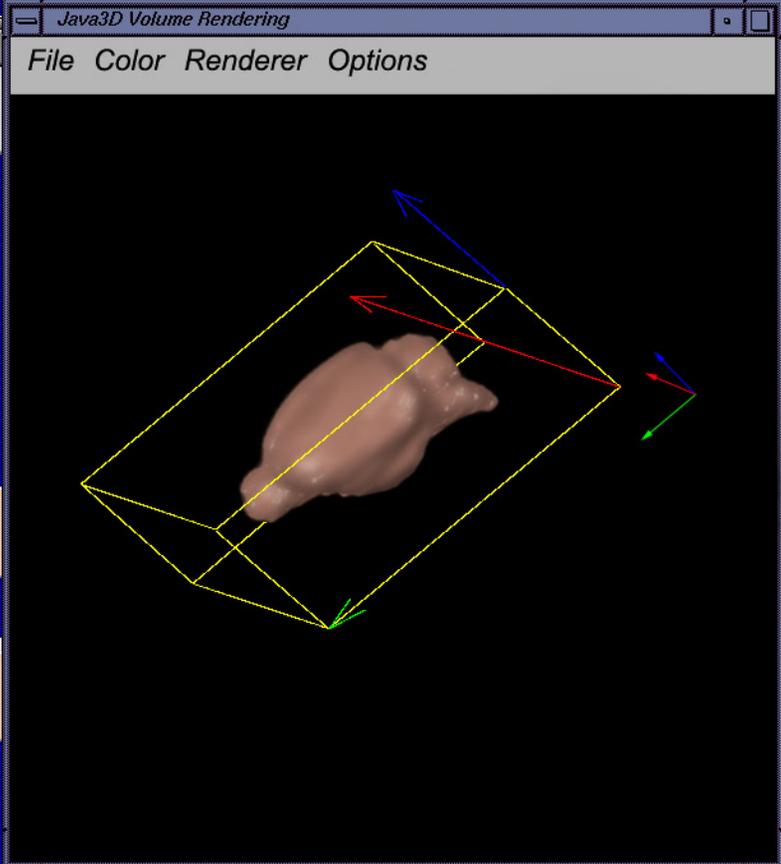
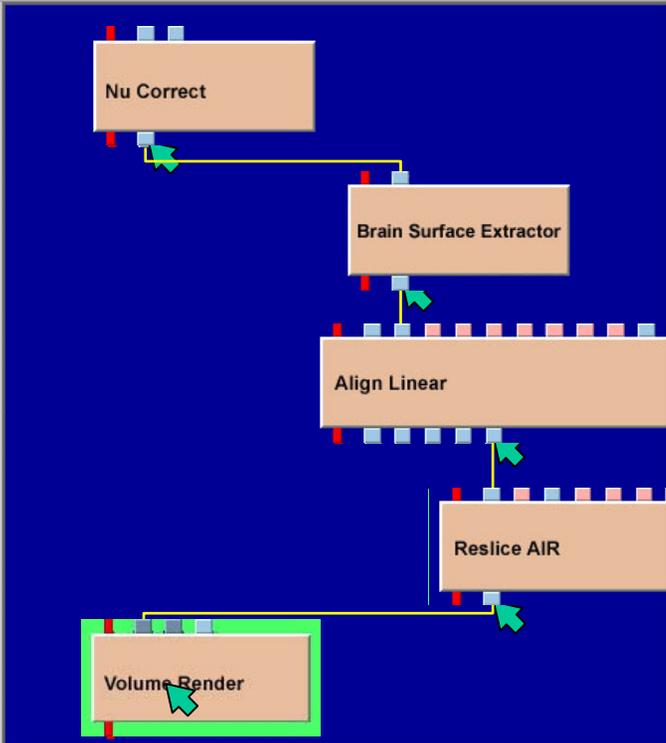


LONI PIPELINE

File Edit Execution Server Windows Help

System Module Li...

- Converters
- LONI Tools
 - CVS
 - 2D Colorix
 - Nu Correct
 - Elastic Warp Flat Maps
 - Brain Surface Extractor
 - Plane Objects to UCF
 - Parametric 3D UCF
 - Volume Render
 - UNIX Commands
 - AIR Tools
 - Align Linear
 - Combine AIR
 - Dual Editor
 - Gaussian Smooth
 - Invert AIR
 - Reslice AIR
 - Soft Mean



User's Module List:

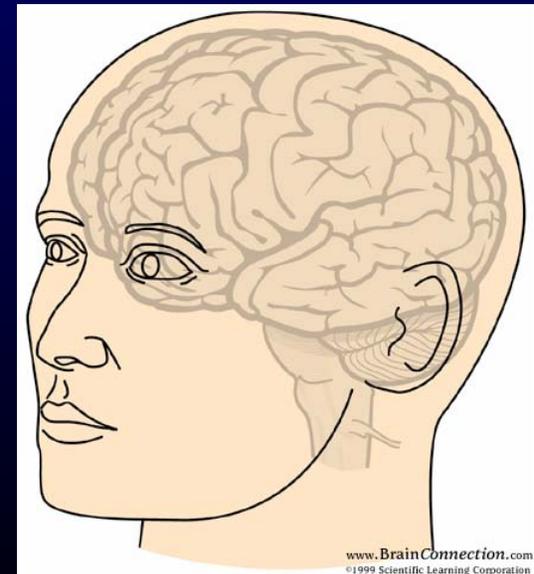
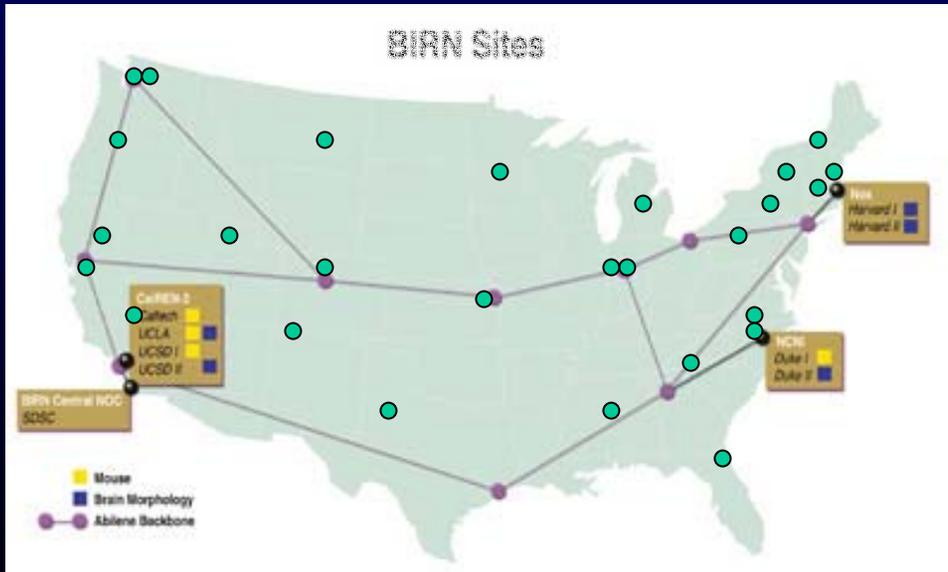
Name	Volume Render
Location	/usr/lshare/java/LONI_Vis
Command	VolRend_driver -InFile1 -InFile2 -InFile3
Stdin	

Ok Cancel



BIRN Future

- Addition of new sites in a rapid, well-defined fashion
- Cross-species integration of data
- Characterization of treatment at multiple levels
- Expand model to other areas of biomedical research



www.BrainConnection.com

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NCRR