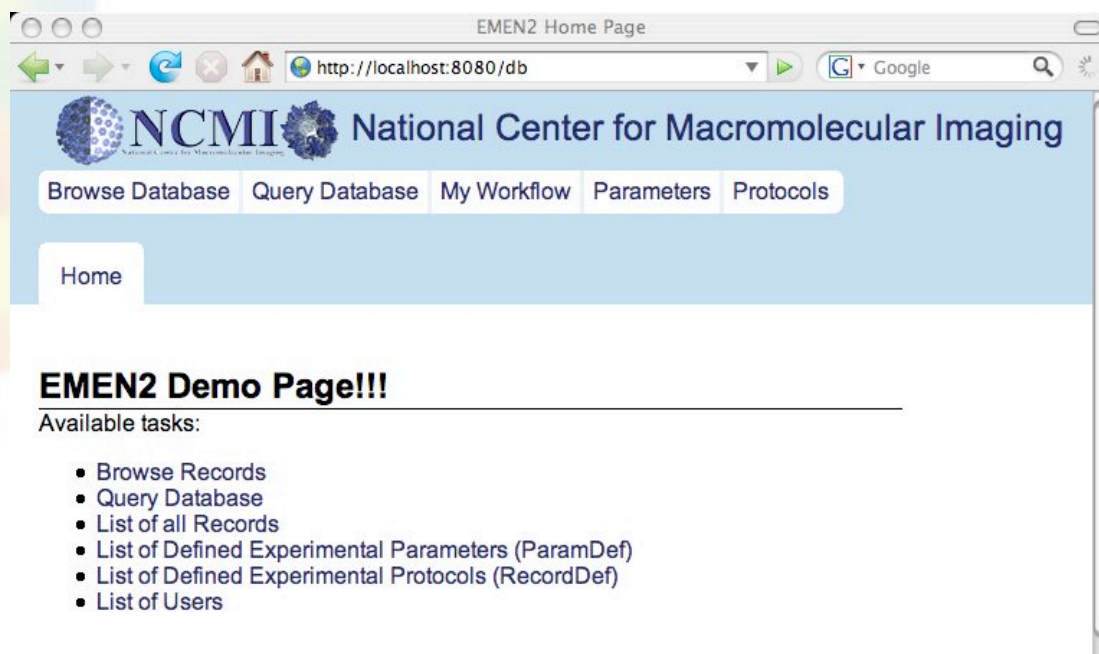


EMEN2: An Extensible, Collaborative, Electronic Lab Notebook



Problem:

Traditional databases lack the flexibility needed for an electronic lab notebooks. They usually rely on fixed schemas that can be difficult to extend, foresight in preparing tables, and designs optimized for millions of similar records instead of the lower-volume but somewhat heterogeneous data of a labnotebook.

However, unstructured documents like traditional notebooks lack the consistency and searchability of databases.

Is there a way to provide an environment where neither flexibility nor consistency is compromised?

EMEN2 Goals:

Data sharing internally and externally

Organization in a complex, changing environment

Standardization of user practices

User friendly and extensible by “normal” people

Flexibility and adaptability to a variety of lab types

Security, in-line viewing of record histories

Instrument integration

Structure through flexible hierarchies

Data-mining and analysis

Overview:

Parameters:

Well defined, semantically correct variable names to store experimental parameters in a consistent way

Protocols:

Free-form, human readable description of an experiment, listing what should be done and what should be recorded

Records:

An instance of a protocol and collected values

Parameters:

All data is associated with a Parameter type that describes what the value represents, what kind of data is stored, default units or possible value choices, and how the Parameter is related to other Parameters.

The use of well-defined Parameters helps ensure the goal of global consistency which is necessary for meaningful data mining. The use of semantically meaningful Parameter names, the Parameter hierarchy, and concise descriptions helps the user choose the most appropriate Parameter type to record a value.

Protocols:

Simple description of an experiment, similar to lab notebook:

A prepared `$grid_prep` was placed in a pair of forceps and loaded into the plunger. `$cryogen` was prepared below the plunger. `$grid_volume` of specimen was deposited on the front of the grid using a pipette. The grid was then blotted on `$grid_blot_side` using `$filter_paper_type` and the plunger was triggered after a `$grid_plunge_delay` to rapidly submerge the grid in the cryogen.

Parameters are embedded into the description and filled-in with each new record created.

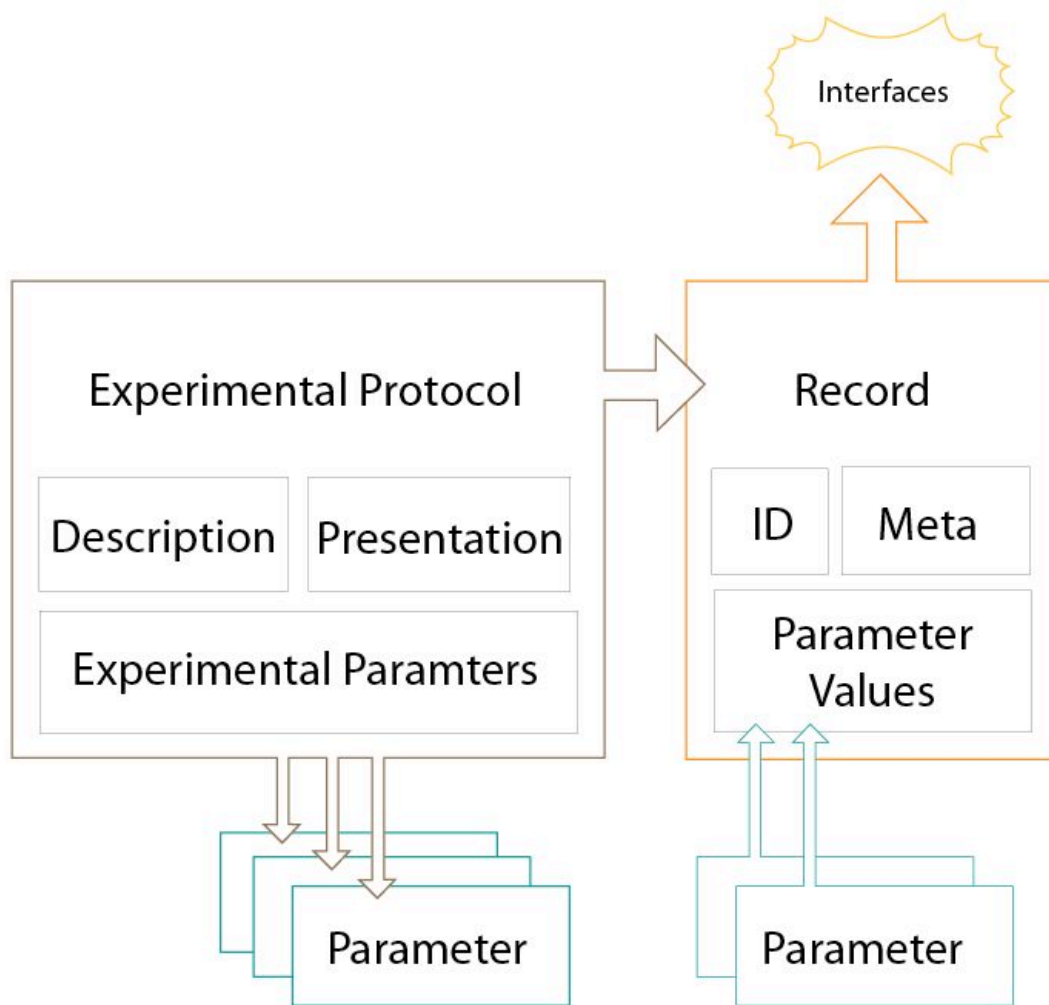
Protocols also define their own presentation options in the user-interface using a similar syntax. A hierarchy is also used to structure protocols to enhance searching and organization.

Records:

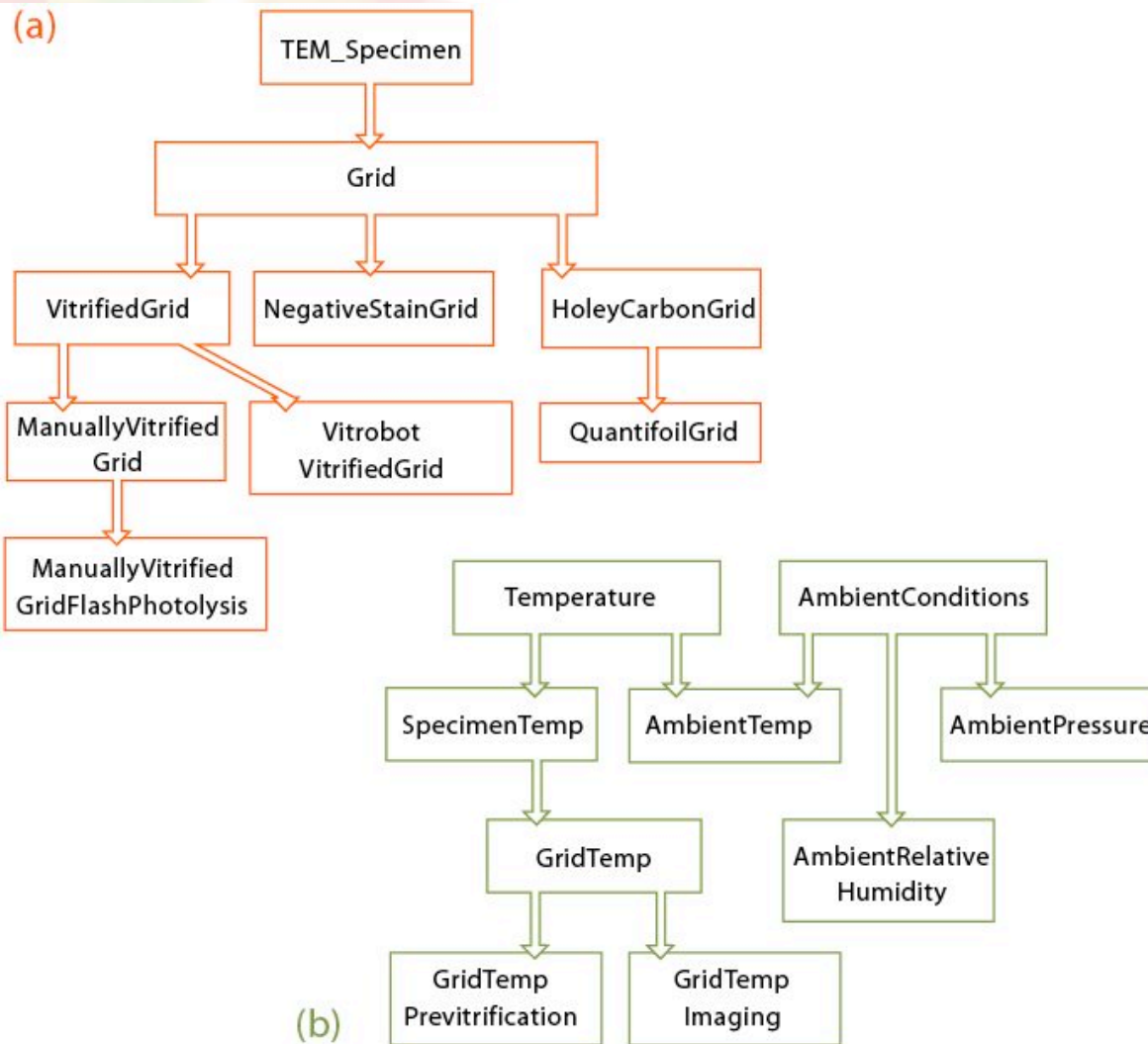
Each record is an instance of an experimental protocol, like a typical page in a lab notebook. The user (or instrument) enters all the data collected during the experiment as defined by the Protocol.

A key to EMEN2's flexibility is the ability to attach additional parameters and values at any time, beyond what is defined by the protocol. These are fully indexed and integral to the record.

Organization of data and projects is provided through the use of multiply-linked, user defined hierarchies.



Each record inherits base parameters from an Experimental Protocol. Additional parameters may be attached to the record at any time by the user. The Experimental Protocol also defines various layouts for the presentation of the information.



Examples of sections of Experimental Protocol and Parameter hierarchies. Well defined hierarchies are helpful in creating new parameters and protocols while preserving the maintainability and structure of the database.

Database Core:

Berkeley DB

BSDDB Python Library

Stores Pickled Python objects

Native Python data types

Security checked before all operations

Flexible indexing and storage options for large
binary files

Interface:

Python

Twisted.Web base library for handling HTTP

AJAX used where appropriate to increase
responsiveness

Friendly, efficient web-based interface with quick access to common tools and information



National Center for Macromolecular Imaging

[Browse Database](#) [Query Database](#) [My Workflow](#) [Parameters](#) [Protocols](#)

EMEN2 → All Groups → NCMI

NCMI » All Children » project (140)

Address of user	One Baylor Plaza N420 Alkek Building
City	Houston
Contact email	webadmin@blake.bcm.tmc.edu
Date data entered	2001/07/05
Date of Experiment	2001/07/05
Fax number	713 798-8682
Homepage	ncmi.bcm.tmc.edu
Institution	Baylor College of Medicine
Organization	Verna and Marrs McLean Department of Biochemistry & Molecular Biology
Phone number	713 798-6989
Record Title	NCMI
Source of financial support	NCRR, NIGMS, NEI, NIAID, R. Welch Foundation

Created: 2001/07/05 laurien	Modified: 2001/07/05 harid
- Comments and history:	
+ Add comment	
+ User permissions and emails:	
+ Child history	
+ Attach files	
+ Add child	
Views: params default protocol	

Hierarchical organization, useful table summaries

EMEN2 → All Groups → NCI → Project → Project → Wah Chiu

Project → Wah Chiu

» All Children

» labnotebook (6)

» microscopy (6)

» purification (3)

microscopy

(0-6 of 6)

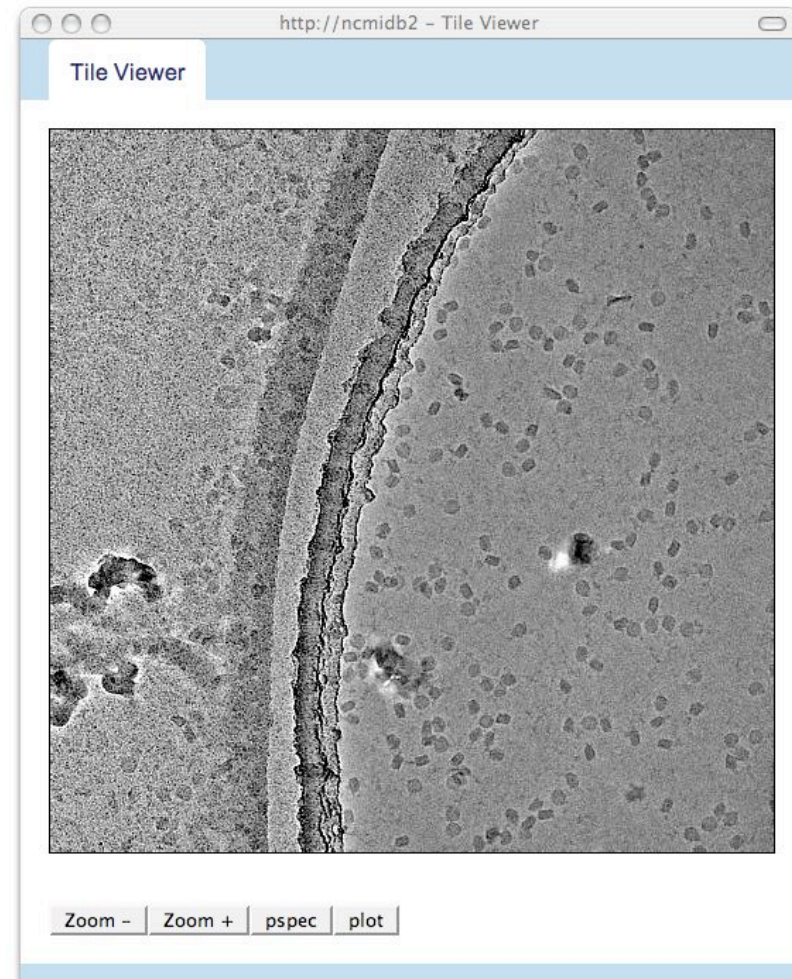
Date of Experiment	Link to microscope	Data entered by	Freezing session ID	Condenser used	Size of electron beam spot	Size of aperture	Magnification	Methods used	Ice thickness	m: childcount
2004/10/20				70	1.0	40	40.0	FasTEM MDS	About right	34
	Purpose of Experiment: collecting data									
2004/03/24				70	1.0	40	50.0	Yoshi box	About right	102
	Purpose of Experiment: collecting data									
2004/10/28				150	2.0	60	40.0	MDS	About right	281
	Purpose of Experiment: collecting data									
2004/03/31				70	1.0		50.0	Yoshi box	too thin to about right	136
	Purpose of Experiment: collecting data, Sample E2 box									
2004/04/05				70	1.0	40	50.0	Yoshi box	About right	427
	Purpose of Experiment: collecting data									

Integrated Viewing of Data:

Image Browser:

Google-Maps Style
interactive viewing
of image data with
fast zooming and
panning

Quick access to
data quality
evaluation tools



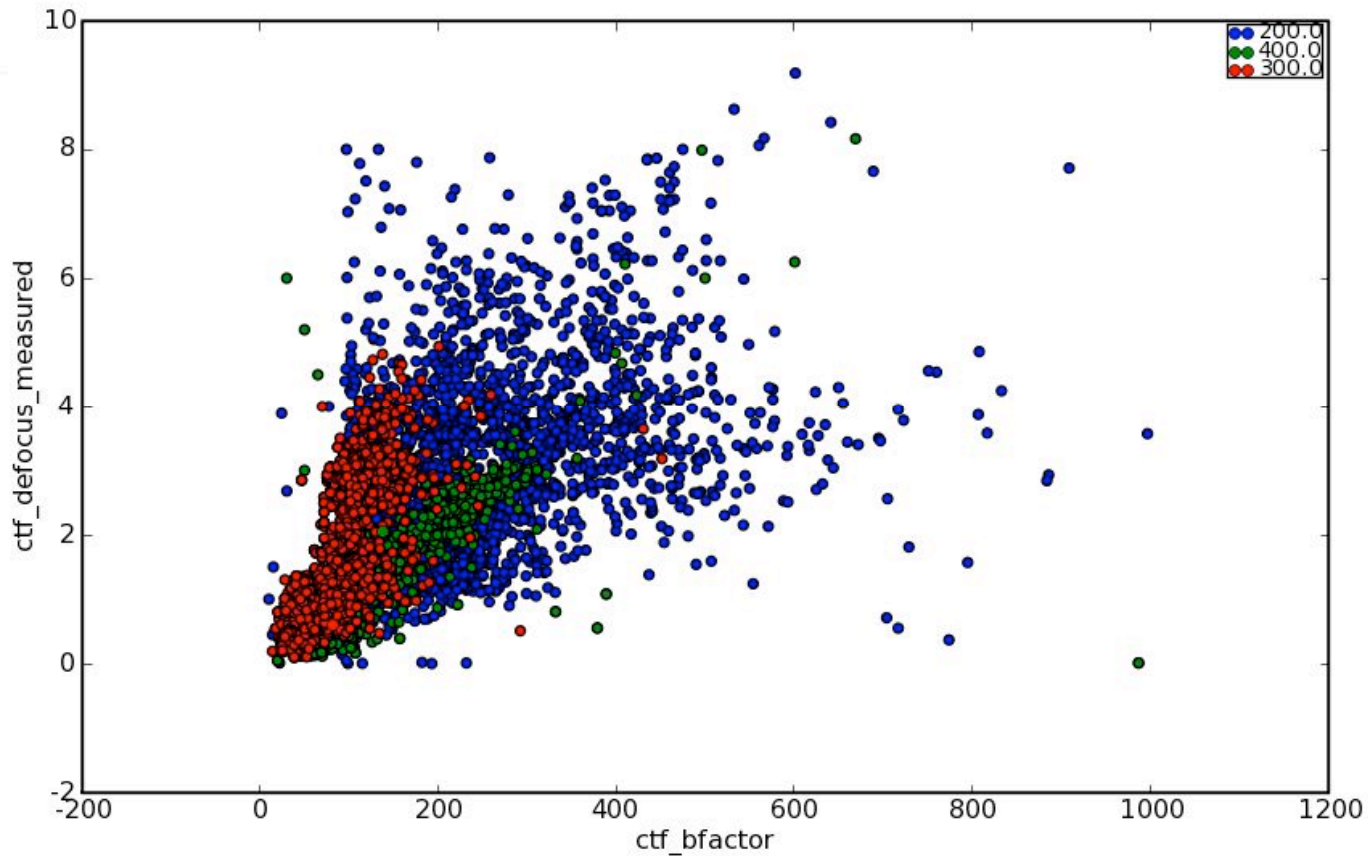
Flexible query language with integrated graphing

Database Query

```
plot ctf_defocus_measured vs ctf_bfactor ctf_bfactor is between 10 and 1000 and  
ctf_defocus_measured > 0 group by tem_voltage
```

Query:

Submit



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