

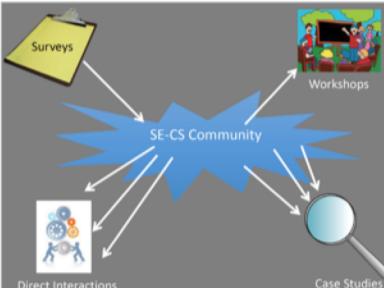
# Making Software Engineering Work for Computational Science & Engineering: An Integrated Approach

SE4Science

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- Publications
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Software Engineering for Science

The increase in the importance of Scientific & Engineering Software motivates the need to identify and understand which software engineering (SE) practices are appropriate. Because of the uniqueness of the scientific & engineering software domain, existing SE tools and techniques developed for the business/IT community are often not efficient or effective. Appropriate SE solutions must account for the salient characteristics of the scientific & engineering software development environment. To identify these solutions, members of the SE community must interact with members of the scientific & engineering software community.



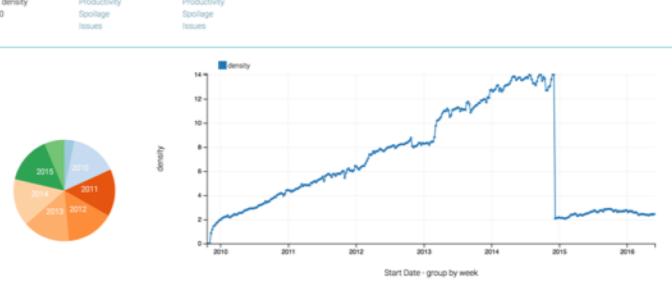
Public Dashboard - Golang

Live updates for tracked project

Project: Golang  
name: Golang  
frequency: week  
metric: density  
filter: > 0

Weekly Metrics: Defect Density, KLOC, Productivity, Coverage, Issues

Monthly Metrics: Defect Density, KLOC, Productivity, Coverage, Issues

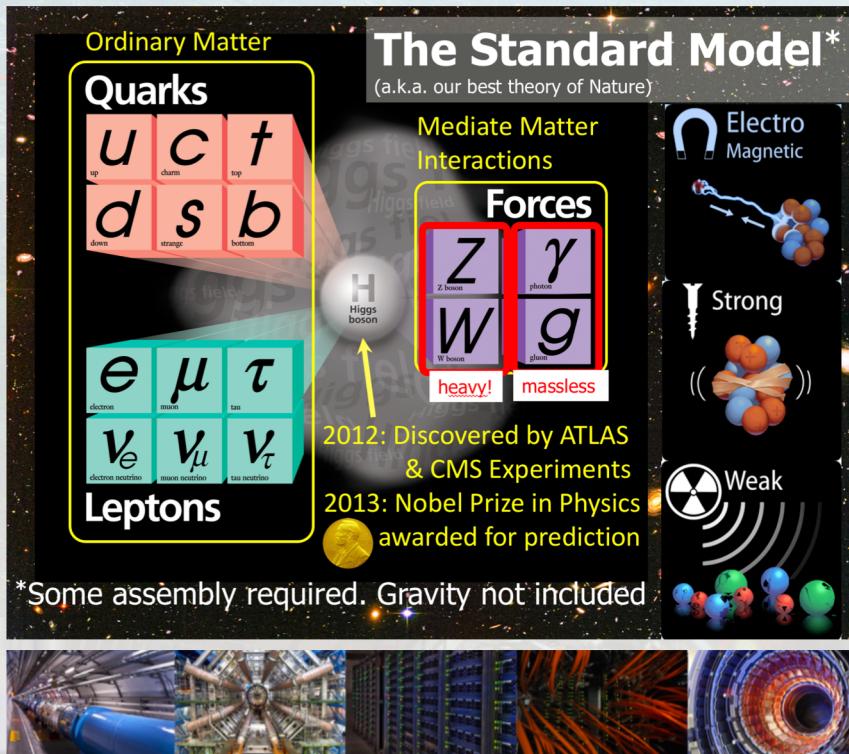
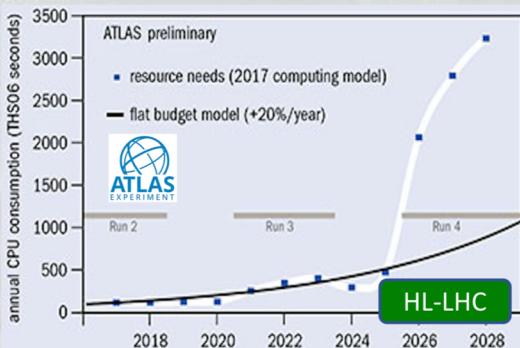
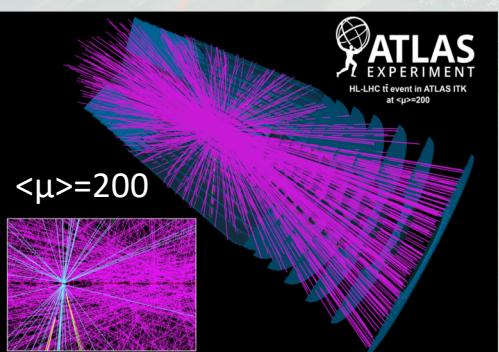


Jeffrey C. Carver  
George K. Thiruvathukal

1445344  
1445347

# Conceptualization of a Software Institute for High Energy Physics

- **High-energy Physics: Software & Computing enables our science**
- **Big challenges ahead for the High-Luminosity (HL-)LHC era**
  - x10 projected shortfall of CPU & storage



- **Advances in hardware will not get us there → need advances in Software!**
- **Community process → Strategic Plan for a HEP Software Institute**



Strategic Plan for a  
Scientific Software Innovation Institute ( $S^2I^2$ )  
for High Energy Physics

Peter Elmer (Princeton University)  
Mark Neubauer (University of Illinois at Urbana-Champaign)  
Michael D. Sokoloff (University of Cincinnati)

December 20, 2017

**M. Neubauer**  
Supported by ACI-1558233

**P. Elmer, M. Sokoloff**  
ACI-1558216, ACI-1558219

# Automated synchronization and boundary condition application for the Cactus framework

Samuel Cupp, Steven Brandt, Peter Diener  
Louisiana State University

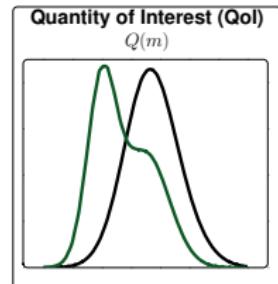
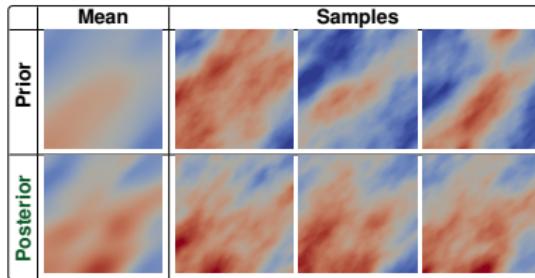
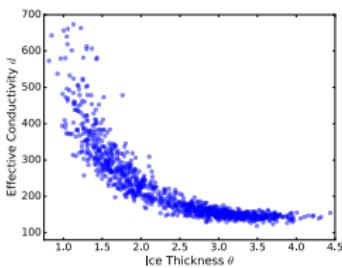
- Cactus Framework is an open-source environment for numerically solving Cauchy problems in parallel
- Current ghost zone synchronization and boundary condition application requires non-trivial, manual scheduling by programmers
- PreSync project replaces old system with an automated scheme
  - Tracks region of validity for grid functions (*interior* or *everywhere*)
  - Schedules synchronization and boundary conditions as needed
- PreSync reduces burden on users and programmers to understand inner workings of the Cactus Framework

We are supported by NSF Grant #1550551.



# Integrating Data with Complex Predictive Models under Uncertainty: An Extensible Software Framework for Large-Scale Bayesian Inversion

Noisy + Uncertain Models → Informed Decisions



- Software framework (Python/c++) for large-scale Bayesian inference
- Easy to use for both users and algorithm developers
- Combined capabilities of MUQ and hIPPYlib

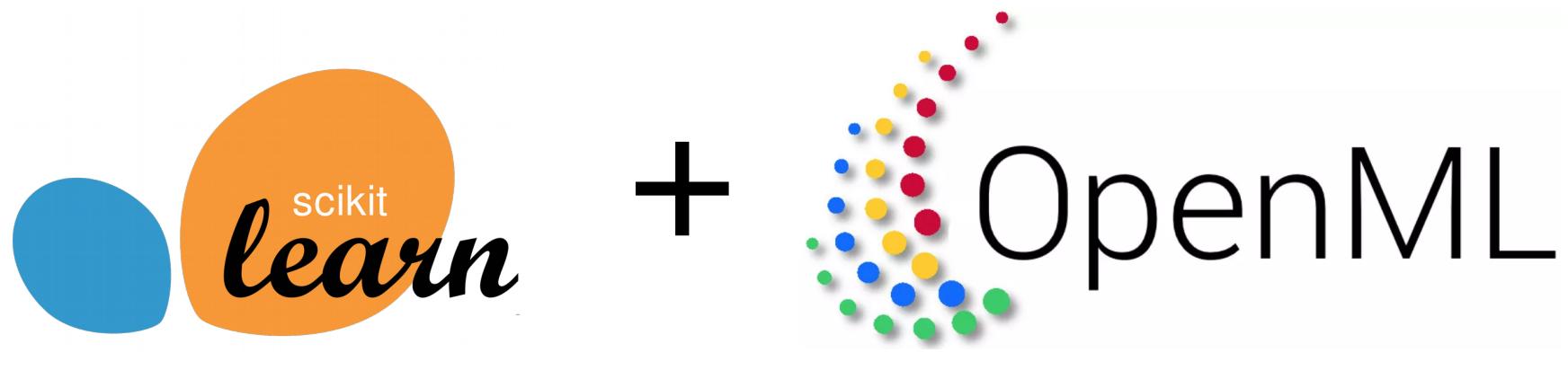
This work was partially supported by National Science Foundation grants ACI-1550487, ACI-1550547, and ACI-1550593.



COLUMBIA  
UNIVERSITY

# SI2-SSE: Improving Scikit-learn usability and automation

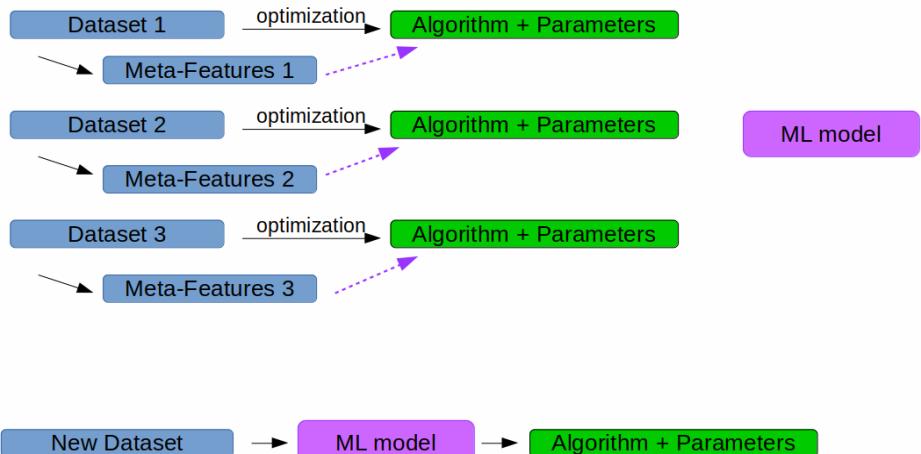
Andreas C. Müller, Columbia Data Science Institute

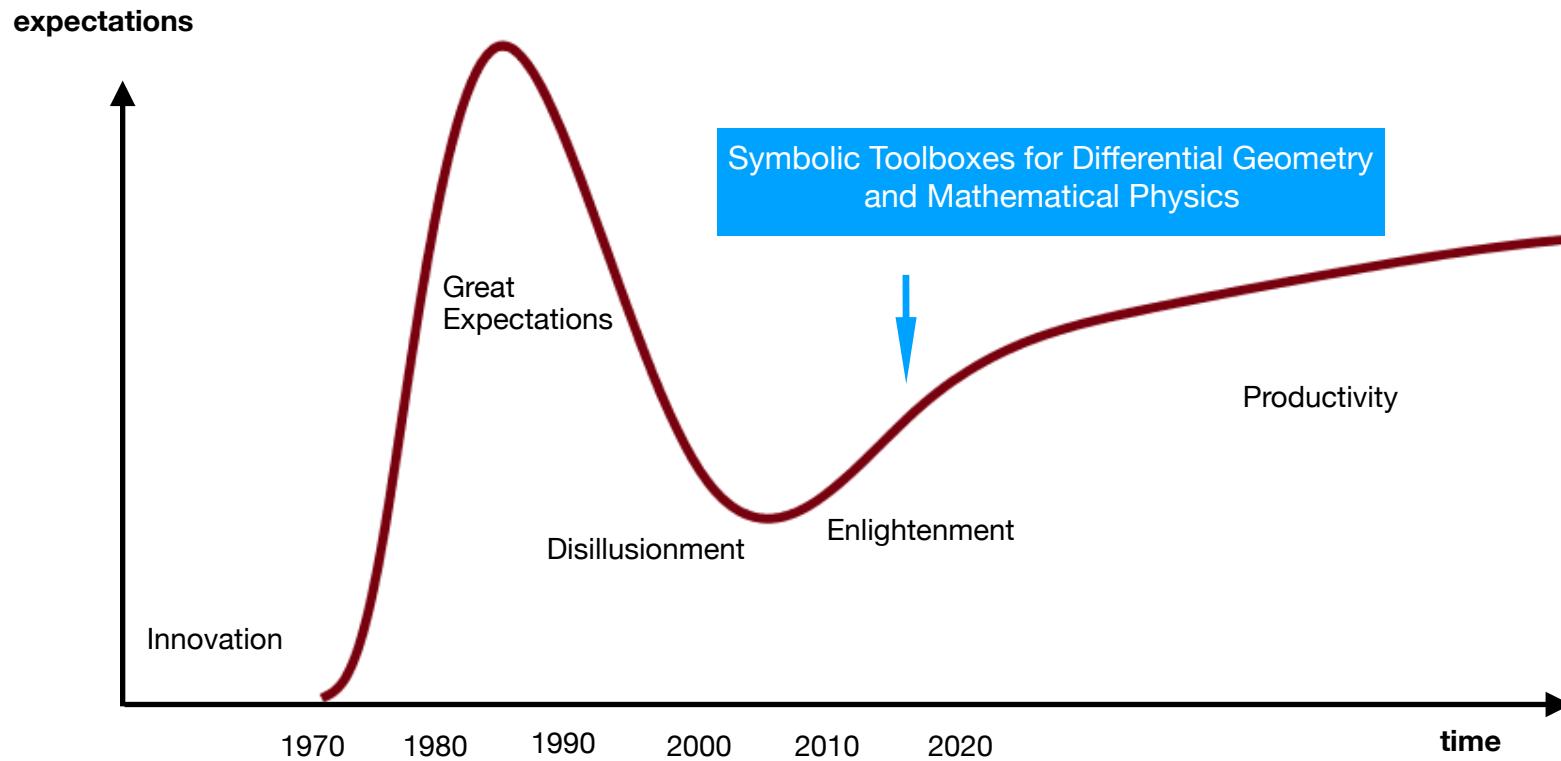


> 150 models

20.000 datasets

# Meta learning





## Project Highlights

- Symbolic computation – computer algebra
- Expert Systems: toolboxes and libraries for domain scientists and educators
- Multiple domains: Differential geometry, Lie theory, general relativity and field theory, geometry of differential equations
- Vertically integrated, interdisciplinary curriculum development

# SI2-SSE: Development of a Software Framework for Formalizing ForceField Atom-Typing for Molecular Simulation

Christopher R. Iacovella<sup>1</sup>, Peter Volgyesi<sup>2</sup> and Janos Sallai<sup>2</sup>

<sup>1</sup> Department of Chemical and Biomolecular Engineering, Vanderbilt University,

<sup>2</sup> Institute for Software Integrated Systems, Vanderbilt University



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School of Engineering



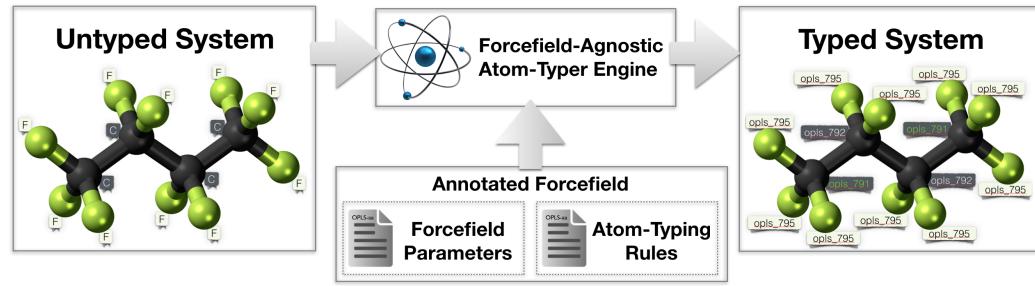
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MUMS  
Multi-scale Modeling and Simulation

- **Challenge:** Develop a general scheme to encode and apply forcefield parameter rules
  - Forcefields describe the way atoms and collections of atoms interact via a set of adjustable parameters
    - *Can contain thousands of sets that are differentiated by the chemical context of an atom, e.g.:*
      - *number of bonds, identity of bonded neighbors, local-environment of bonded neighbors, etc.*
  - Rules for usage are typically hard-coded into software as a deeply nested hierarchy with specific rule order
    - *This approach can be difficult to debug, extend, and disseminate*
- Defining parameter usage via SMARTS and overrides
  - Encode chemical context using the SMARTS language for defining molecular patterns

```
opls_135 = [C;X4](C)(H)(H)H
```
  - Set rule precedence via “overrides”

```
opls_148 = [C;X4]([C;X3])(H)(H)H overrides=opls_135
```
  - Rules are both **human and machine readable** and can be tested for accuracy and completeness
- Foyer: General Python library for applying forcefields
  - Atom types assigned using matching patterns determined by performing a subgraph isomorphism on the system graph
  - Rules can be evaluated in any order
    - *Uses a fixed point iterative scheme that creates white- and blacklists, rather than rigid hierarchy*
  - Source code does not change when rules change
    - *Allows for easier testing, validation, versioning and dissemination*



# SI2-SSI: Integrated Molecular Design Environment for Lubrication Systems (iMoDELS)

Peter Cummings<sup>1</sup>, Clare McCabe<sup>1</sup>, Ákos Lédeczi<sup>1</sup>, Gabor Karsai<sup>1</sup>, Adri van Duin<sup>2</sup>, Paul Kent<sup>3</sup>  
<sup>1</sup>Vanderbilt University, <sup>2</sup>Pennsylvania State University, and <sup>3</sup>Oak Ridge National Laboratory



VANDERBILT  
School of Engineering

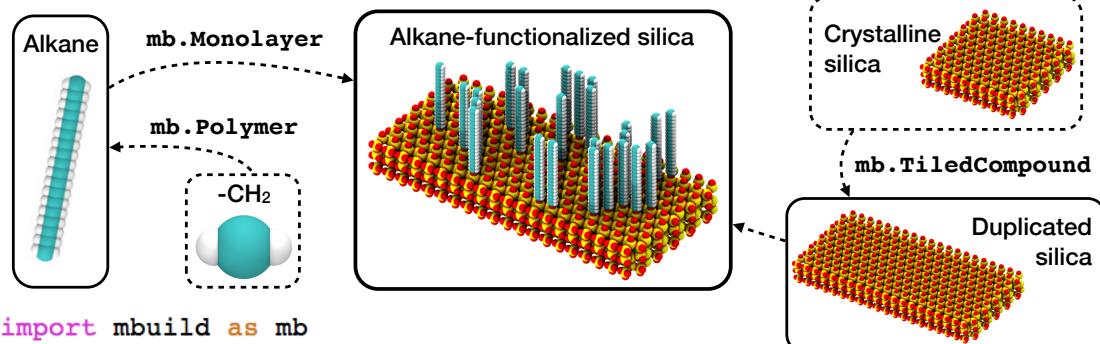


INSTITUTE FOR SOFTWARE  
INTEGRATED SYSTEMS  
MUMS  
Multiscale Modeling and Simulation

- Challenge:** Improved lubrication strategies required for devices with nanoscale separations
  - Molecular simulation can be used to understand lubrication at the molecular level
  - Use this to screen for relationships between chemistry and tribology (i.e., lubrication properties)

- mBuild:** a Hierarchical, Component Based Molecule Builder written in Python

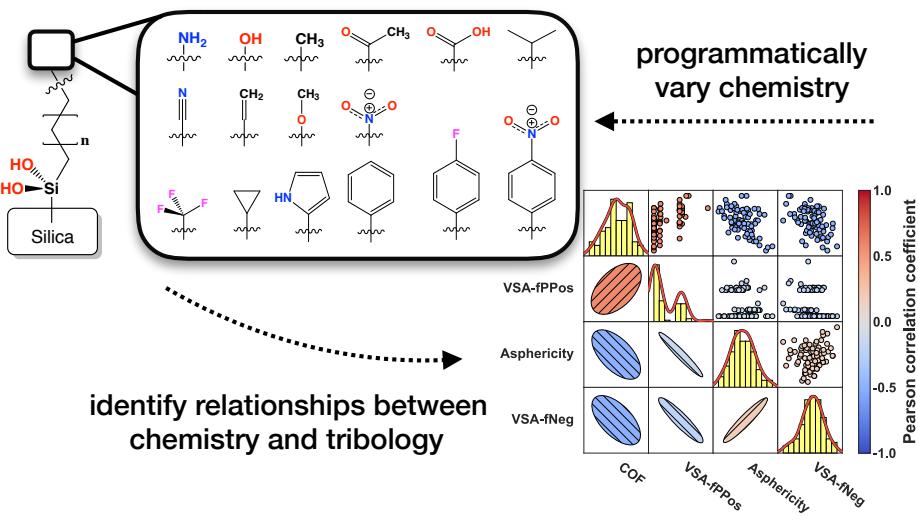
- Construct complex systems from smaller, interchangeable pieces
- Enable programmatic variation of chemistry, required for screening



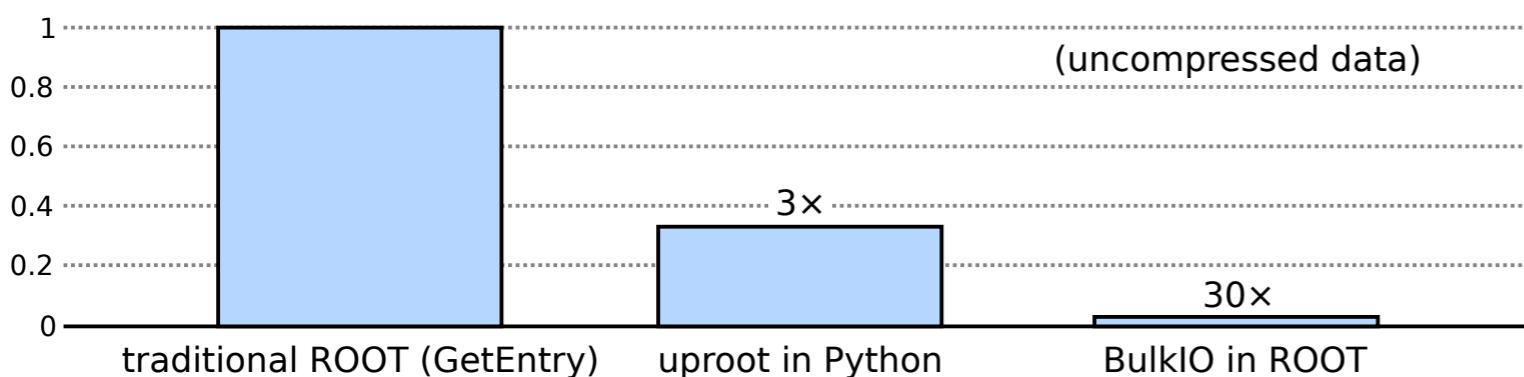
- metaMDS:** define parameter landscape for screening

```
# Initialize simulation with a template and some metadata
sim = mds.Simulation(name='monolayer',
                      template=configure_run_script,
                      output_dir='output')

chain_lengths = [8, 12, 16, 20]
for length in chain_lengths:
    parameters = {'chain_length': length,
                  'n_molecules': 100,
                  'forcefield': 'OPLS-aa',
                  'build_func': build_monolayer}
    # Parameterize our simulation template
    sim.parametrize(**parameters)
```

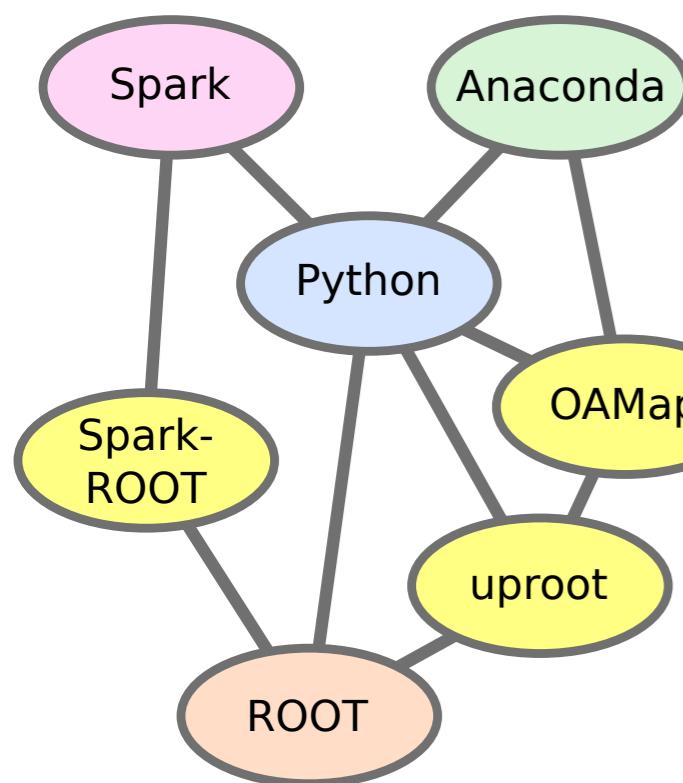


# Advancing Analysis for HEP



## Improved Performance

To reduce the time to scientific discovery and to enable more in-depth analyses, we are increasing the rate of access to ROOT data files. This includes streamlined access to simpler data types (uproot and BulkIO) and faster compression algorithms (LZ4 and ZSTD). These efforts have already provided factors-of-several improvements.



## Bridging to Big Data

“Big Data” software in industry, such as the Spark and scientific Python ecosystems, both complement and reproduce functionality of HEP software developed. To provide more options and reduce maintenance burdens, DIANA is building bridges between HEP software and the Big Data ecosystems: Spark-ROOT to Spark and uproot/OAMap to Numpy, Numba, and Dask.

## Statistical Techniques

We are developing tools and methods for statistical analysis in HEP, including research for simulator-based inference (Carl), machine learning for particle physics (Scikit-Optimize), and software for efficient numerical computations.



## High Level Tools

We are therefore striving to present HEP analysis with higher-level interfaces. Scikit-HEP incorporates HEP techniques in Pythonic idioms, uproot provides access to ROOT data as Numpy and Pandas abstractions, and OAMap compiles object-centric user code into fast array operations.

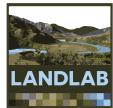


# uproot



# A Landlab-built cellular automaton model of hillslope evolution

Gregory E. Tucker<sup>1,2</sup>, Scott W. McCoy<sup>3</sup>, Daniel E.J. Hobley<sup>4</sup>



a python toolkit for modeling earth-surface processes



1 - CIRES and Department of Geological Sciences, University of Colorado, Boulder

2 - Community Surface Dynamics Modeling System (CSDMS)

3 - Department of Geological Sciences and Engineering, University of Nevada, Reno

4 - School of Earth and Ocean Sciences, Cardiff University, Cardiff, UK

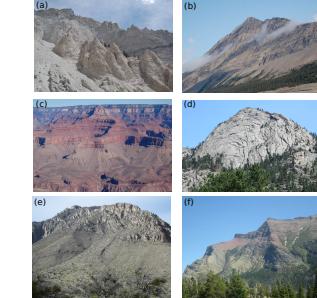
**Abstract:** This poster describes and explores a new continuous-time stochastic cellular automaton model of hillslope evolution. The software was written using Landlab, a Python package for rapidly creating and modifying 2D numerical models of various sorts. The Grain Hill model provides a computational framework with which to study slope forms that arise from stochastic disturbance and rock weathering events. The model can reproduce a range of common slope forms, from fully soil mantled to rocky or partially mantled, and from convex-upward to planar shapes. With the addition of a rule for large blocks, the model captures the morphology of hogbacks, scarps, and similar features. Model parameters have a direct link with corresponding parameters in continuum theory, and can reproduce observed slope forms at the correct scale.

**Landlab** is a Python package that supports building and exploring 2D grid-based computational models. Key capabilities include:

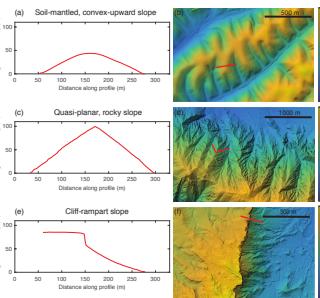
- Set up a structured or unstructured **grid** with just a few lines of code
- Use, create, and/or combine standardized **components** that encapsulate numerical simulations of individual processes

For more about **Landlab** see:  
<http://landlab.github.io>

We seek to explain the form and evolution of rocky hillslopes like these:

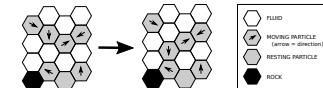


Common slope forms include parabolic, planar, and cliff-rampart:

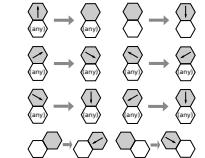


We start with a lattice-grain cellular model (Tucker et al., 2016):

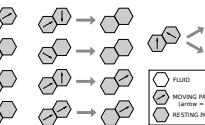
State	Description
0	Fluid
1	Grain moving upward
2	Grain moving up and right
3	Grain moving down and right
4	Grain moving down
5	Grain moving down and left
6	Grain moving up and left
7	Resting grain
8	Rock
(9)	Block (optional)



Transition rules are used to represent gravity ...

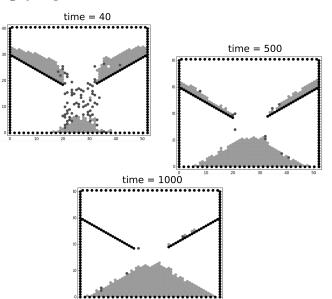


... and frictional collisions:

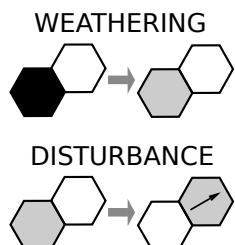


Transport- vs. weathering-limited behavior is reflected in scaling of gradient and fraction soil (regolith) cover

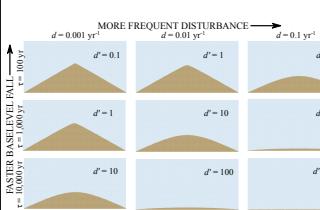
Example of granular dynamics simulation: emptying of a silo



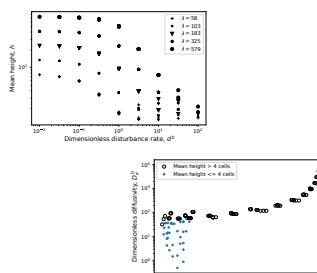
For the Grain Hill mode, we add rules for periodic soil disturbance and soil formation by weathering



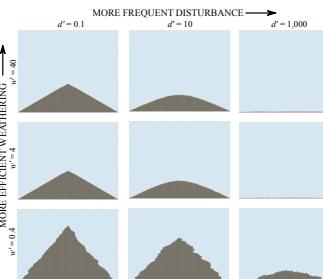
When the hill is 100% soil, the height and form depend on the ratio of disturbance frequency to uplift rate



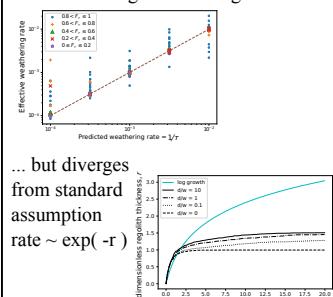
We can interrogate the scaling of height and effective diffusivity



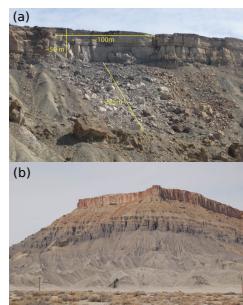
When rock is present, behavior ranges from transport- to weathering-limited



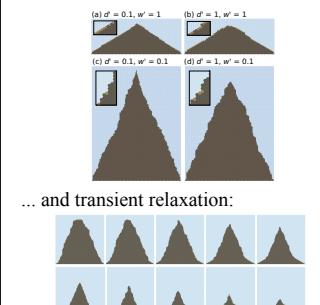
Model captures influence of fractional soil cover on average weathering rate ...



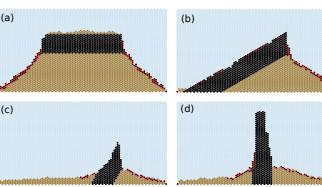
What about cliff-rampart morphology?



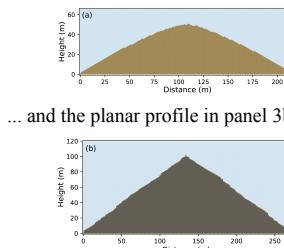
With "collapse rule": quasi-steady forms ...



Adding a rule for "blocks" allows us to capture mesas, hogbacks, and rocky ridges



With the right parameter mapping, model captures specific case studies, such as the convex-up slope in panel 3a ...



## ACKNOWLEDGMENTS:

Landlab was supported by a SI2-SSI award ACI-1450409. Hillslope evolution research was supported by EAR-1349390. Additional support was provided by the Community Surface Dynamics Modeling System (CSDMS), EAR-1226297.

## REFERENCES:

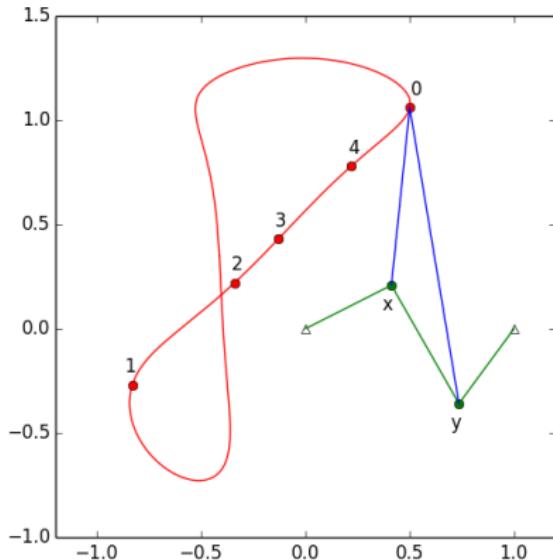
- Hobley, D. E., Adams, J. M., Nudurupati, S. S., Hutton, E. W., Gasparini, N. M., Istanbulluoglu, E., & Tucker, G. E. (2017) Creative computing with Landlab: an open-source toolkit for building, coupling, and exploring two-dimensional numerical models of Earth-surface dynamics. *Earth Surface Dynamics*. doi:10.5194/esurf-5-21-2017.
- Tucker, G.E., Hobley, D.E., Hutton, E., Gasparini, N.M., Istanbulluoglu, E., Adams, J.M., and Nudurupati, S.S. (2016) CellLab-CTS 2015: Continuous-time stochastic cellular automaton modeling using Landlab. *Geoscientific Model Development*, v. 9, p. 823-839, doi:10.5194/gmd-9-823-2016.
- Tucker, G.E., McCoy, S.W., & Hobley, D.E.J. (2018) A lattice grain model of hillslope evolution. *Earth Surface Dynamics Discussions*. doi:10.5194/esurf-2018-4.

# Solving Polynomial Systems with PHCpack and phcpy

PHCpack is software for Polynomial Homotopy Continuation

phcpy is a new Python package, available at [www.phcpack.org](http://www.phcpack.org)

use case from the phcpy tutorial:



reproduces J. Mech. Design paper

jupyter bbsolvesnippet Logout

SageMath 8.0

File Edit View Insert Cell Kernel Help PHCpy

In [2]:

```
f = ['x*y^2 + y - 3;', 'x^3 - y + 1']
from phcpy.solver import solve
sols = solve(f)
for sol in sols: print sol
```

total degree : 9  
2-homogeneous Bezout number : 7  
with partition : { x }{ y }  
general linear-product Bezout number : 7  
based on the set structure :  
{ x }{ y }{ y }  
{ x y }{ x }{ x }  
mixed volume : 7  
stable mixed volume : 7  
t : 1.00000000000000E+00 7.13177119756522E+00  
m : 1  
the solution for t :  
x : -1.14928524947248E+00 -4.33149270057445E-01  
y : 1.28839810793789E-01 -1.63511747105322E+00  
== err : 1.650E-16 = rco : 3.038E-01 = res : 2.220E-16  
=  
t : 1.00000000000000E+00 2.72148344088863E+00  
m : 1  
the solution for t :  
x : -1.14928524947248E+00 4.33149270057445E-01  
y : 1.28839810793789E-01 1.63511747105322E+00



# CRESCAT

## A Computational Research Environment for Scientific Collaboration on Ancient Topics

PI: David Schloen, University of Chicago

### Goals

- Support all 5 stages of data** for multi-disciplinary collaborative research
- Automate data transfers and transformations** from one stage to the next via high-level GUI
- Accommodate heterogeneity** of data sources, types, and schemas while preserving the original ontologies
- Seamless scalability** for data management and algorithmic analyses
- Ensure sustainability** of software maintenance and technical support
- Test and document** with complex use cases from
  - ❖ Archaeology
  - ❖ Paleontology
  - ❖ Historical linguistics
  - ❖ Ancient economics
  - ❖ Population genetics
  - ❖ Paleoceanography etc.

### 1. Acquisition

- External curated data repositories** via live links using their Web APIs
- Instruments and data files** with support for many data types and file formats (2D images, 3D models, audio, video, geospatial, etc.)
- Manual entry** with offline mode for field input and automated syncing of data when back online

### 4. Publication

- REST API** exposes published data as XML with XSLT stylesheets to render it as JSON/HTML
- Sample Web apps** provided for various research domains, to be customized as needed
- Secure, password-protected data** controlled and published by owners

### 2. Integration

- Ontology-agnostic data warehouse** stores both data and the ontologies inherent in the data
- XQuery DBMS** optimized for hierarchies of atomic keyed data objects representing spatial, temporal, linguistic, and taxonomic relationships
- Automatic parsing** of source data to populate the integrated warehouse

### 5. Archiving

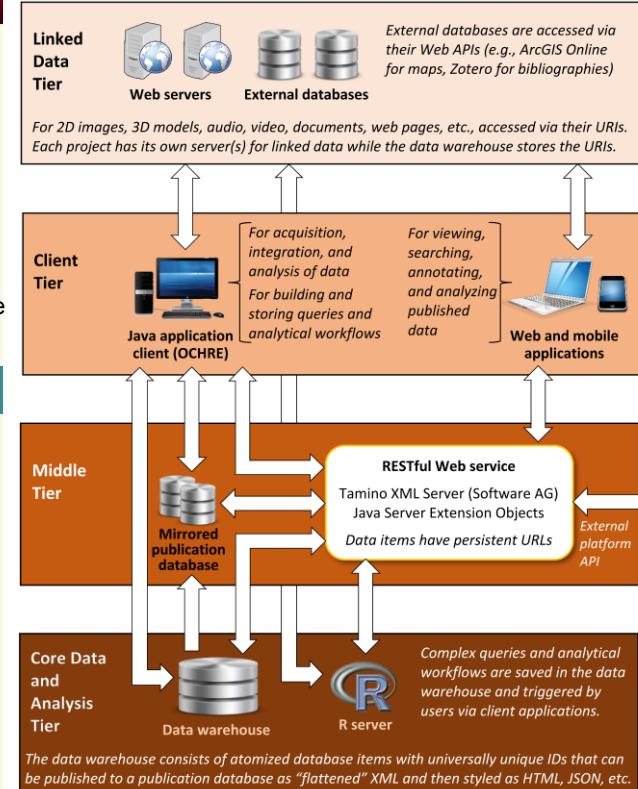
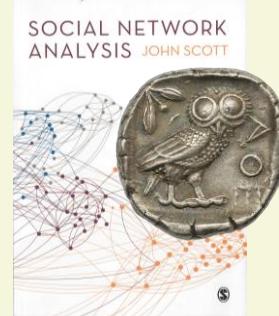
- OWL-RDF ontology specification** documents the top-level (upper) ontology underlying the data warehouse
- Can export RDF triples** conformant to the OWL ontology, preserving all distinctions and relationships in the data, for use in other graph databases

### 3. Analysis

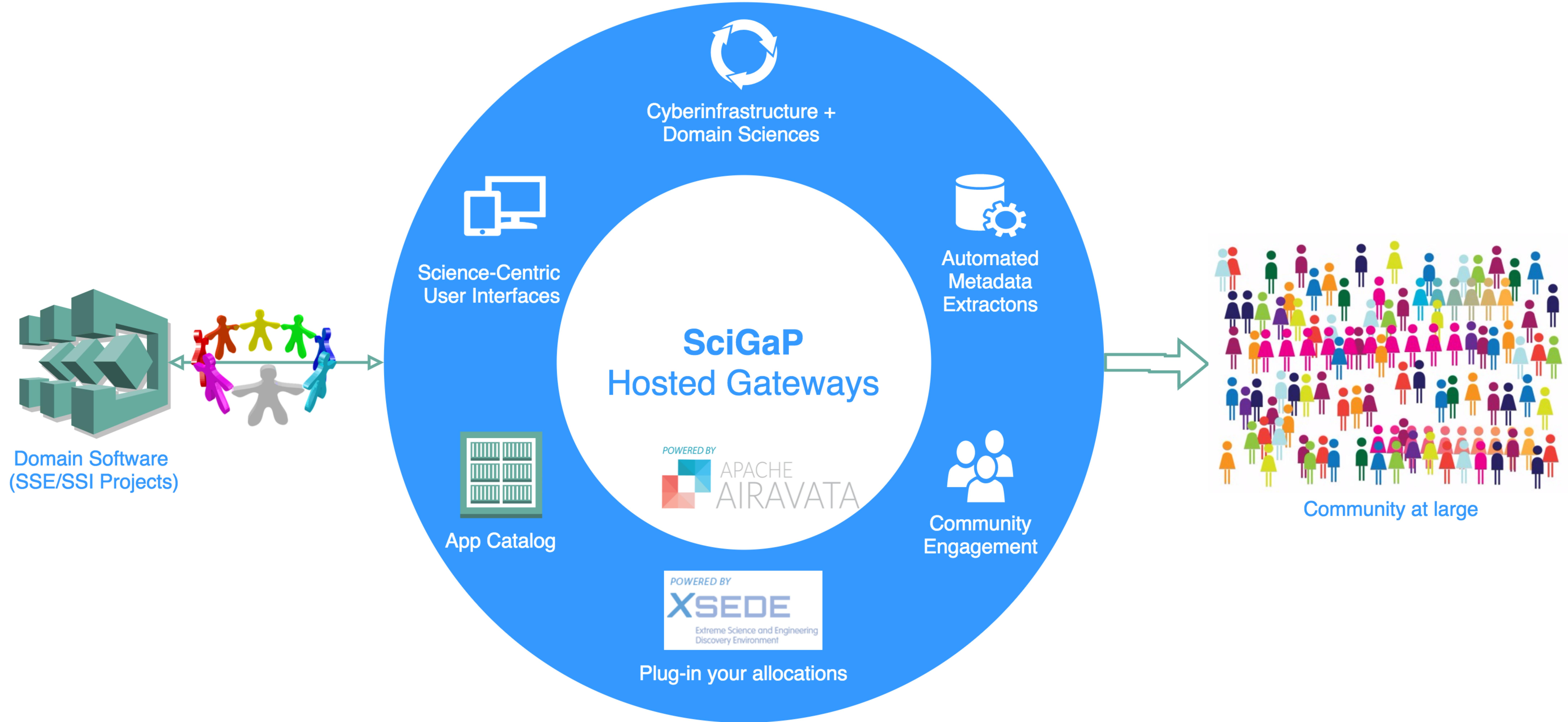
- Complex queries** use hierarchical taxonomies with semantic inheritance
- Statistical analysis and visualization** via tightly integrated R server with data-aware console
- Geospatial mapping and analysis** via ArcGIS Online and ESRI components

### Example Use Case

Ancient Greek economy  
via network analysis of thousands of coin hoards

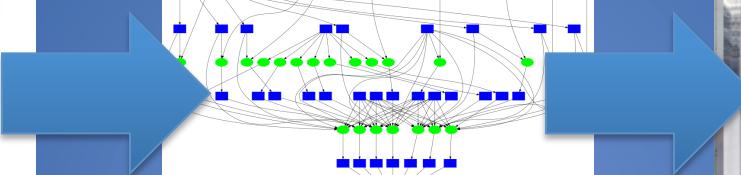


Funded by NSF SI2-SSI award 1450455



# SI2-SSE: Scaling Up Science with the Cooperative Computing Tools

Douglas Thain, University of Notre Dame



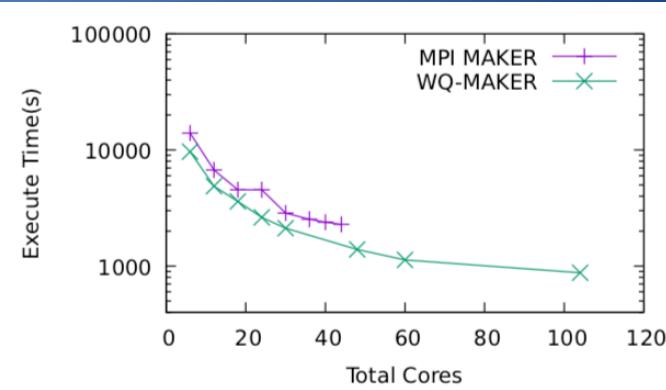
Portable Workflow  
Expression

```
{ "command" :  
  "mysim.exe -p " + x*2 +  
  " input.txt > output." + x + ".txt",  
  "outputs" : [ "output" + x + ".txt" ],  
  "inputs" : [  
    "input.dat",  
    "mysim.exe"  
  ]  
} [ for x in range(1,100) ]
```

Experiment  
Management

Table 2. U-Net Hyperparameter Search Spaces	
Parameter	Values
Min. Kernels	16, 32, 64, 128
Kernel Size	1, 3, 5, 7, 9
Activations	sigmoid, tanh, relu, elu, PReLU, LeakyReLU, ThresholdedReLU
Initializers	zeros, ones, glorot_normal, he_normal
Regularizers	l1, l2, l1_l2
Dropout Rate	uniform distribution over [0, 1]
Learning Rate	uniform distribution over [10 <sup>-4</sup> , 1]

Scalability  
and Robustness





# NIMBLE: Programmable Statistical Modeling for Hierarchical/ Graphical Models

## What do we want to do with hierarchical models?

### 1. More and better MCMC

- Many different samplers
- Better adaptive algorithms

### 2. Numerical integration

- Laplace approximation
- Adaptive Gaussian quadrature
- Hidden Markov models

### 3. Maximum likelihood estimation

- Monte Carlo EM
- Data cloning
- Monte Carlo Newton-Raphson

### 4. Sequential Monte Carlo

- Auxiliary Particle Filter
- Ensemble Kalman Filter
- Iterated Particle Filter

### 5. Normalizing constants

- Importance sampling
- Bridge sampling
- Others

### 6. Model assessment

- Bootstrapping
- Calibrated posterior predictive checks
- Cross-validation
- Posterior re-weighting

### 7. Idea combinations

- PF + MCMC
- MCMC + Laplace/quadrature

## NIMBLE Components

### 1. Domain-specific language (DSL) for statistical models

- We adopt and extend the widely-used BUGS language

### 2. Domain-specific language embedded within R for model-generic algorithms

### 3. Code-generator (compiler) that generates C++ from the model and algorithms DSLs.

- C++ objects are managed from R by dynamically-generated interface classes

### 4. Algorithm library (MCMC, SMC, etc.)

## Core Team

Perry de Valpine (PI); UC Berkeley

Christopher Paciorek (co-PI); UC Berkeley

Daniel Turek; Williams College

Nicholas Michaud; UC Berkeley

Duncan Temple Lang; UC Davis



Journal of Computational and Graphical Statistics

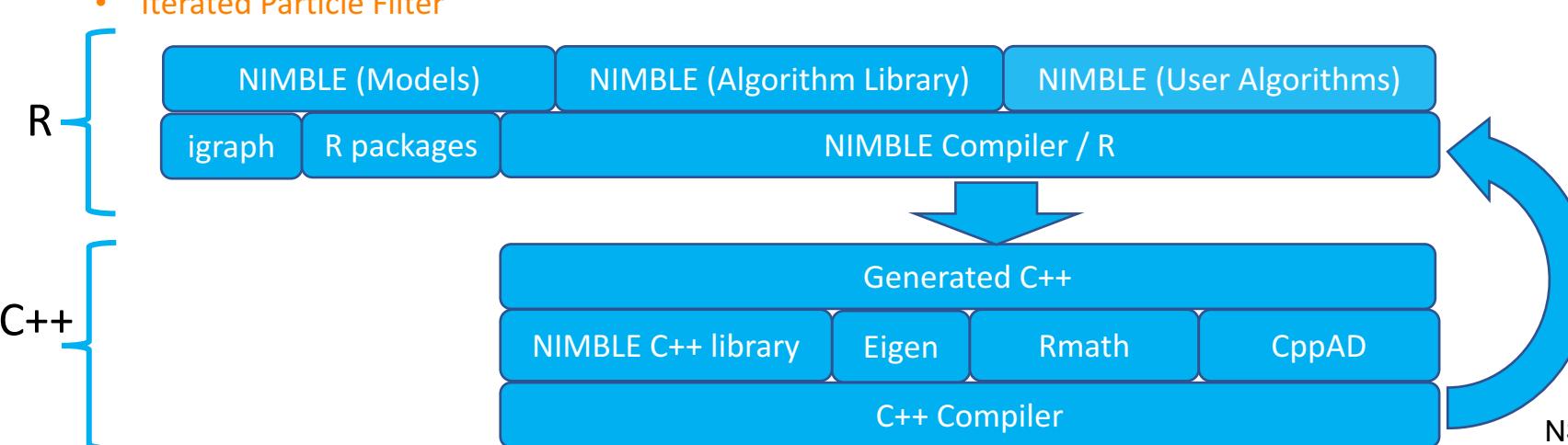
ISSN: 1061-8600 (Print) 1537-2715 (Online) Journal homepage: <http://www.tandfonline.com/loi/ucgs20>

Programming With Models: Writing Statistical Algorithms for General Model Structures With NIMBLE

Perry de Valpine, Daniel Turek, Christopher J. Paciorek, Clifford Anderson-Bergman, Duncan Temple Lang & Rastislav Bodík

<https://r-nimble.org>

NSF ACI-1550488, DBI-1147230 (completed), DMS-1622444



# SI2-SSE: High Performance Low Rank Approximation for Scalable Data Analytics

R. Kannan (ORNL), G. Ballard (WFU), B. Drake (GTRI), and H. Park (GAtech) <https://github.com/ramkikannan/nmflibrary>

**Constrained Low Rank Approximation (CLRA)** for  
Modeling Key Data Analytics problems of clustering,  
topic modeling, community detection, and hybrid  
clustering

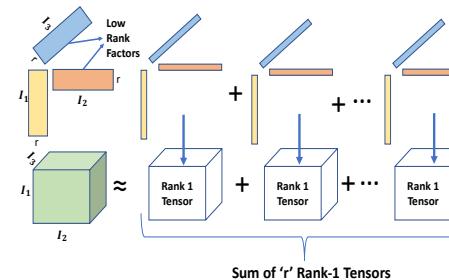
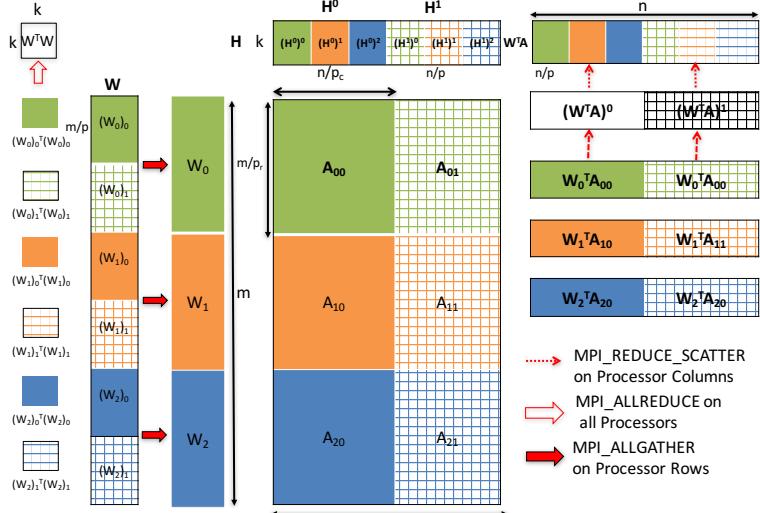
**Our current focus:** Nonnegative Matrix/Tensor  
Factorization (NMF and NTF) and other Variants (e.g.  
Sparse NMF, SymNMF, and JointNMF)

**Why CLRA** such as NMF and NTF?

Utilize advances in numerical linear algebra  
algorithms and software, Behavior of algorithm  
easier to understand and analyze, Facilitates design  
of MPI based algorithms for scalable solutions

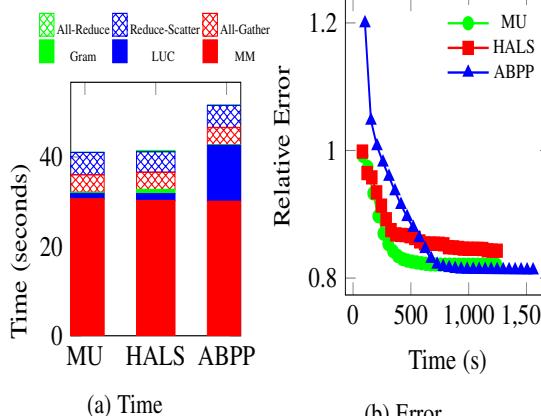
*PPoPP'16, TKDE'18, PPoPP'18, IPDPS'18, JGO'18*

Fast Alternating Updating NMF/NTF (FAUN) Framework:



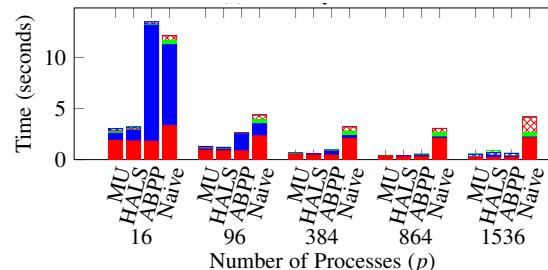
Titan – Dense Matrix, Low Rank 50,  
100 Iterations, 12650 Nodes,  
202500 Cores,

Matrix Size	Algos	NMF Time (in Secs)
2.7 million x 2.7 million	MU	554
	HALS	197.75
	ANLS /BPP	219.8
3.03 million x 3.03 million (72 TBs)	MU	554
	HALS	197.75
	ANLS /BPP	219.8

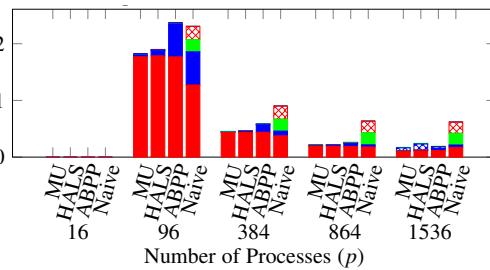


NMF on 118 million Web-graph

Sparse Webbase – 1M Vertex



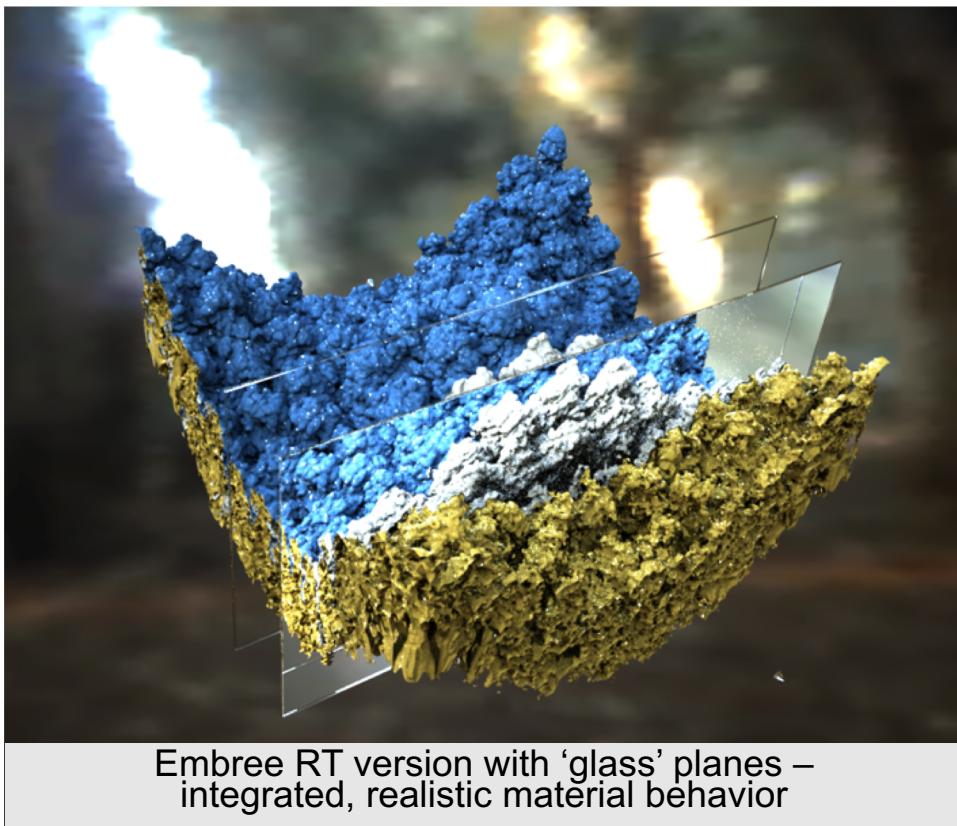
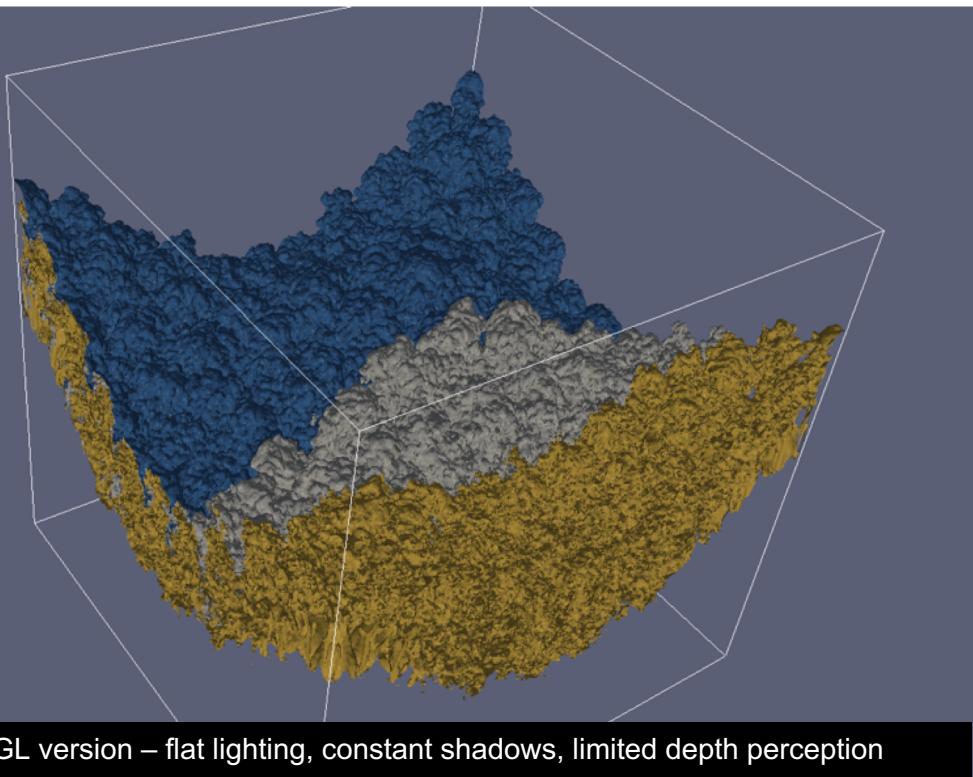
Dense real world-video



# GraviT Distributed Ray Tracing Framework

ACI-1339863 (TACC) ACI-1339840 (Oregon) ACI-1339881 (Utah)

Bring photo-quality rendering to your large-data visualizations through ray tracing, and now integrated into the SI2 yt project!



gvt  
+  
yt

# SI2-SSE: Collaborative Research: Extending the Practicality and Scalability of LibMesh-Based Unstructured, Adaptive Finite Element Computations

**Paul Bauman**



# A MACHINE LEARNING GATEWAY FOR SCIENTIFIC WORKFLOW DESIGN

NSF SI2-SSE #1740151



Akos Ledeczi (PI) · Brian Broll · Tamas Budavari · Peter Volgyesi



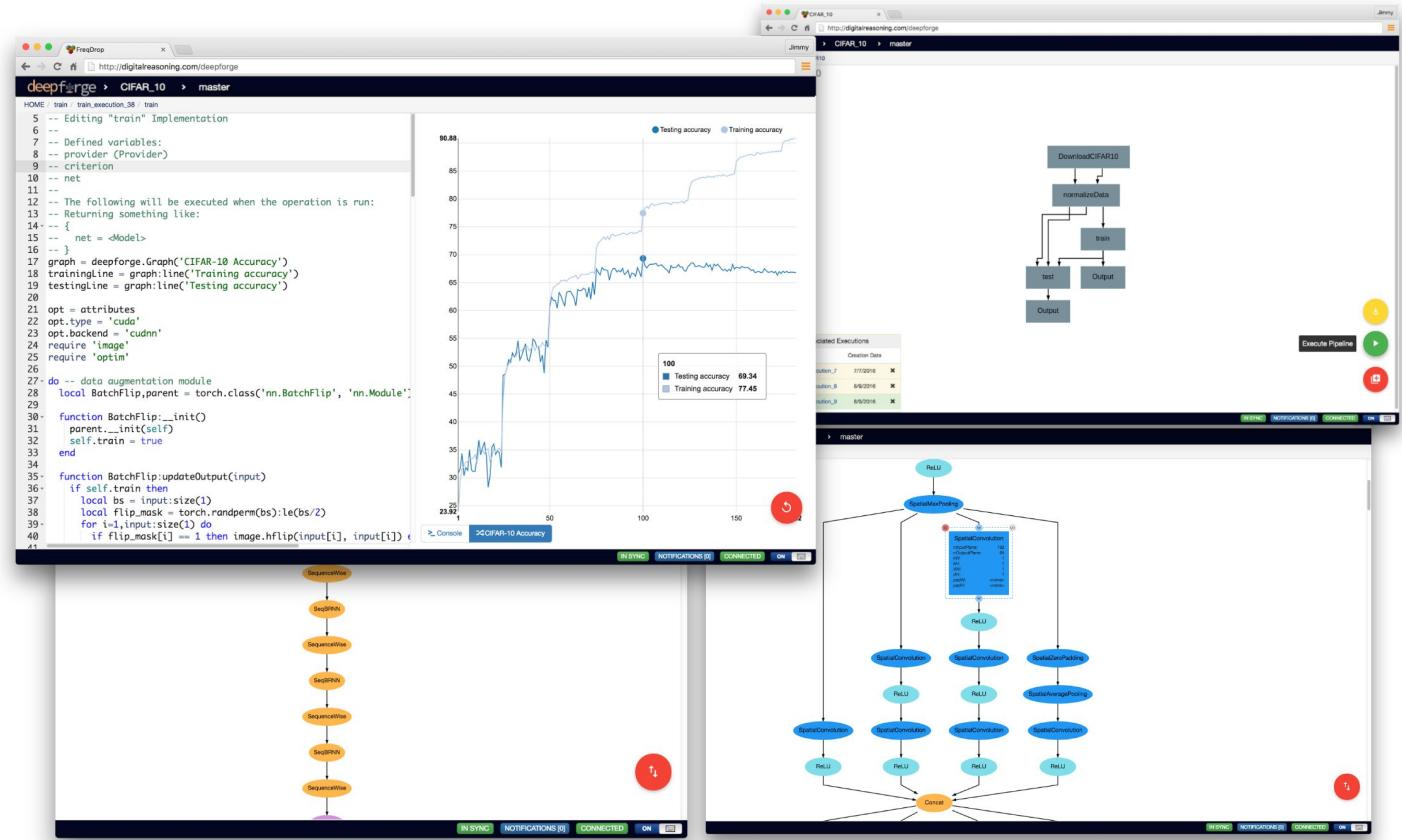
## TENSORFLOW

- DNN architectures
- custom workflows
- alternative engines:  
CNTK, Theano



## WEBGME

- web application
- model-based design
- online collaboration
- project history
- job management



**DeepForge** is an open source platform for **deep learning** designed for promoting **reproducibility, simplicity and rapid development** within diverse scientific domains.



# Why you must visit our poster!

## A modern dense linear algebra software stack

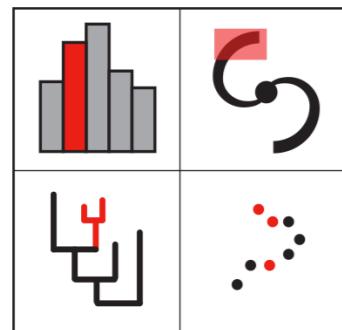
- BLIS: Framework for Rapid Instantiation of BLAS-like functionality
- libflame: LAPACK functionality
- TBLIS: A C++ tensor contraction library

## Effective outreach

- Professional development for scientific software scientists
- Massive outreach through Massive Open Online Courses. (145,000 participants to date)
- Cultivation of external contributors



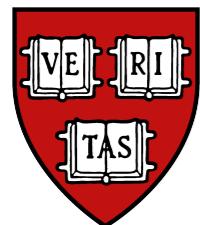
@glueviz



# glue

multidimensional data exploration

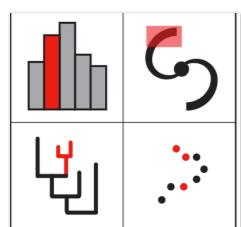
glueviz.org



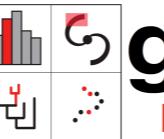
**Alyssa Goodman, PI**  
Harvard University  
[@AlyssaAGoodman](#)



**Michelle Borkin, PI**  
Northeastern University  
[@michelle\\_borkin](#)



**Thomas Robitaille**  
Lead Architect  
[@astrofrog](#)

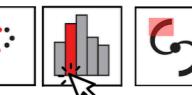


# glue

multidimensional data exploration

glueviz.org [github.com/glue-viz](#) [glueviz.slack.com](#)

glue's modular design


standard 1D, 2D & 3D plots

**built-in**

- standard data loaders
- Data File 1
- Data File 2
- ⋮
- Data File N

**plugin**

- custom data loaders

**options**

- define new variables, import/export insights, interactive plots for the web, save state, all from GUI
- link data files' attributes
- highlight live or algorithmic selections with Boolean logic
- custom buttons, features


standard 1D, 2D & 3D plots

**?**

custom plots

**jupyter**

- user config.py file (loaders, colors, plot types, +)
- access to all matplotlib functions through built-in IPython terminal

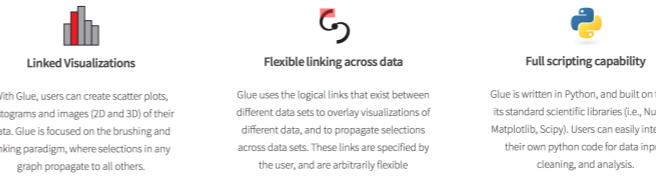

run & interact with glue from Jupyter notebook & other tools

**Want to plug-in your project or tool?**

Consider joining us for **glue-con**, right after JupyterCon, August 27-29, 2018, at Harvard.

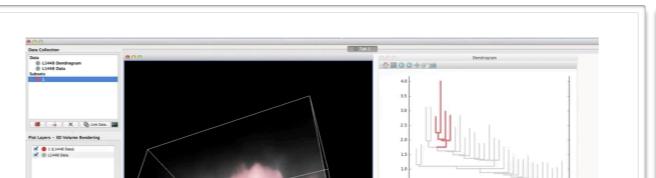

glue-con 2018, CAMBRIDGE, MA

[projects.iq.harvard.edu/gluecon](#)



**Linked Visualizations**  
Flexible linking across data

With Glue, users can create scatter plots, histograms and images (2D and 3D) of their data. Glue is focused on the brushing and linking paradigm, where selections in one graph propagate to all others.



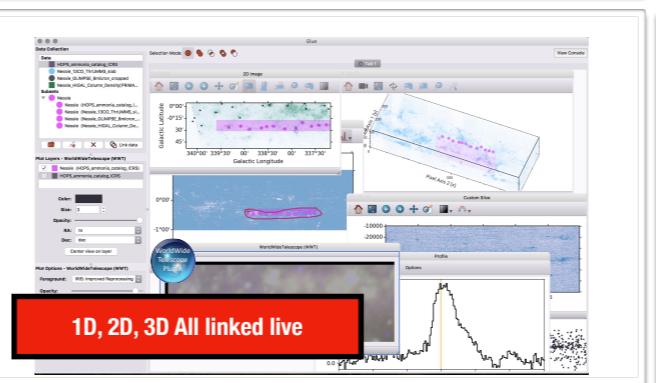
**Full scripting capability**

Glue is written in Python, and built on top of its standard scientific libraries (i.e., Numpy, Matplotlib, Scipy). Users can easily integrate their own python code for data input, cleaning, and analysis.



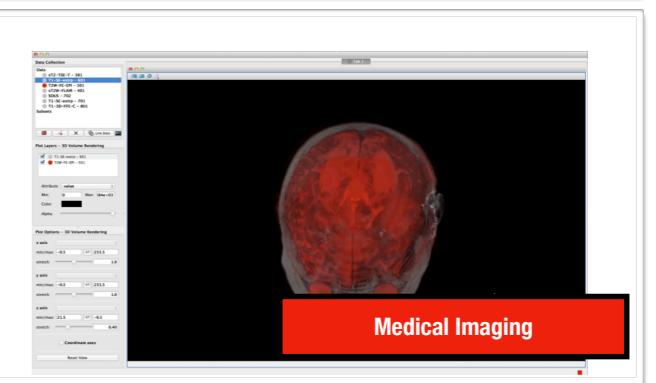
**GIS compatible**

Glue supports various geospatial data formats and can be used for geographic information systems (GIS) applications.



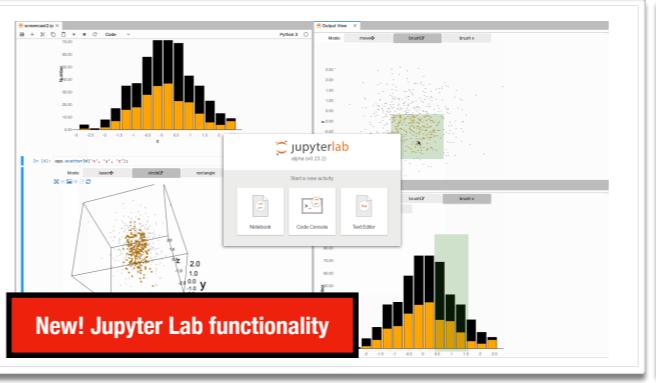
**1D, 2D, 3D All linked live**

Glue allows for live linking between 1D, 2D, and 3D data sets, enabling users to explore complex datasets in multiple dimensions simultaneously.



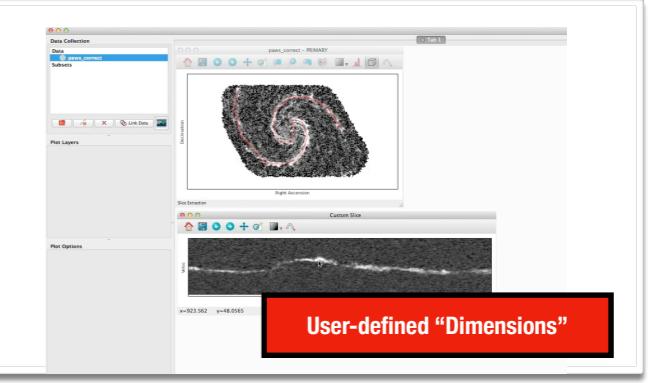
**Medical Imaging**

Glue can handle medical imaging data, such as MRI scans, and provide interactive visualization and analysis tools.



**New! Jupyter Lab functionality**

Glue now integrates with Jupyter Lab, providing a powerful environment for data exploration and analysis.



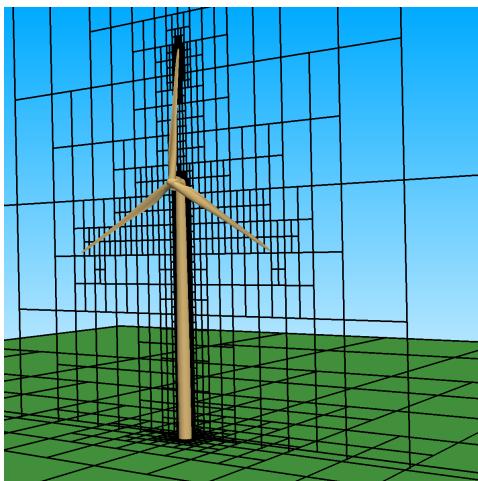
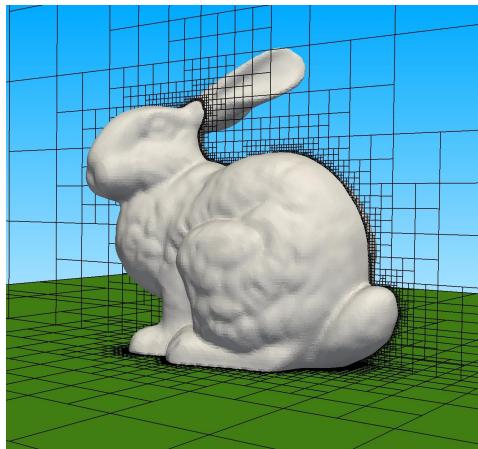
**User-defined Dimensions**

Users can define their own dimensions for data sets, making it easier to work with complex multi-dimensional data.

# Massively Parallel Solvers for Computational Fluid Dynamics on Block Structured Cartesian Grids

Jaber Hasbestan, Scott Aiton, Brenton Peck, Donna Calhoun, Inanc Senocak, Grady Wright\*; <https://github.com/GEM3D>

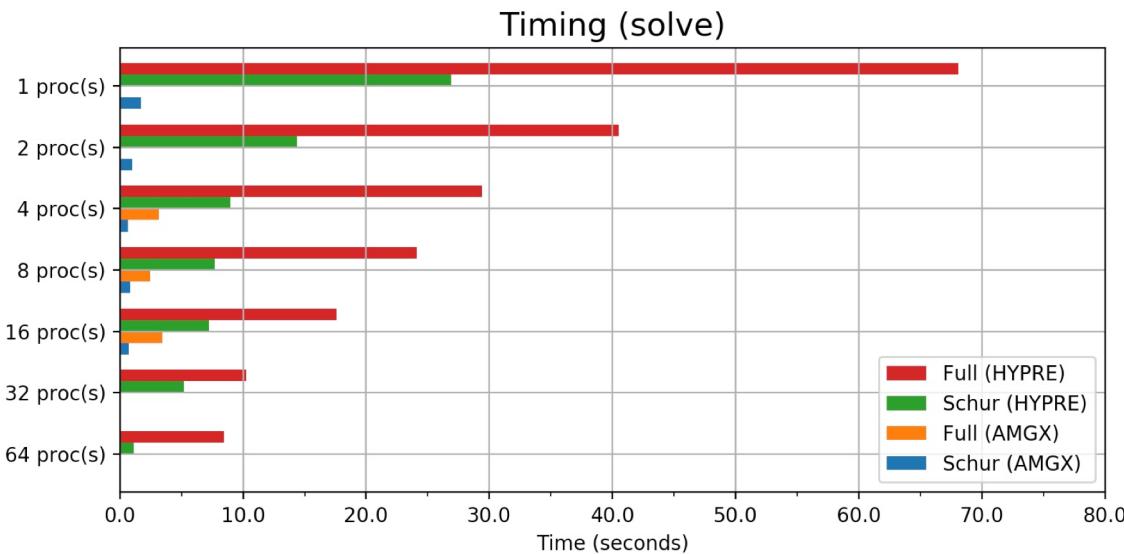
Block Structured Cartesian AMR



Highly scalable red-black binarized-octree for generating and managing the adaptively refined grids

Schur complement domain decomposition method for the **Pressure Poisson** equation on block structured grids

$$\text{Solve: } \begin{cases} \nabla^2 p = f & \text{on } \Omega \\ \mathbf{n} \cdot \nabla p = 0 & \text{on } \partial\Omega \end{cases} \quad \text{for } p \text{ given } f,$$



Refinement level	10	12	14	16
Red-black tree	3.3	21.14	69.75	145.87
Z-curve enhanced hash function	3.7	26.85	87.85	180.01
C++ STL default hash function	4.89	39.95	138.19	272.97



# Automated Detection and Repair of Errors in Event-Driven Applications



Frank Tip

College of Computer and Information Science, Northeastern University  
[www.franktip.org](http://www.franktip.org)

- modern applications rely on event handling for, e.g., user input, network communication
- key operations: register event handlers, emit events, call-back to event handler
- programmer errors are common, and lead to hard-to-debug failures
  - e.g., event race errors depending on nondeterministic scheduling of event handlers
- research goal: provide programmers with better tools to detect and repair such errors
  - based on static & dynamic program analysis



Fig. 4. (i) The AlertRadar report for <https://www.apple.com>, where the magnified part shows the warning message flagged by AlertRadar. (ii) The screen resulting from clicking on the search icon after the page has loaded, which is the normal behavior. (iii) The screen resulting from clicking on the search icon after the page has fully loaded.

# GeoVisuals Software: Capturing, Managing, and Utilizing GeoSpatial Multimedia Data for Collaborative Field Research

**Ye Zhao**

# Collaborative RAPID

BUILDING INFRASTRUCTURE TO PREVENT DISASTERS LIKE HURRICANE MARIA

PUBLIC ACCESS INFORMATION	OBJECTIVE 01 <b>Water Quality Sampling Campaign</b>	OBJECTIVE 02 <b>Data Archive</b>	OBJECTIVE 03 <b>Cyberinfrastructure Advances</b>	<b>Expected Science Outcomes</b>
	Drinking water samples from public streams  Spatially aggregated anonymized information of the impact zone	Baseline assessment: Population Health Data, Healthcare Providers and supporting organizations, natural system environmental variables, Public Water System location and infrastructure status.  Hurricane Maria health and environmental data from public data repositories and Luquillo CZO instruments in El Yunque National Park	LANDLAB raster model grid and diverse data formats  Observation Data Model (ODM2)	
PRIVACY PROTECTED INFORMATION	PRASA Utility, community operated tank system, household data  Teacher collection of student health data (IRB)	Water samples with personal information  De-identified water samples that can be geo-located	Population health researcher user-testing  Water quality professionals and researchers user testing  Individual data owners user testing	<b>DISASTER:</b> Contamination, drought, landslides, bio-diversity  <b>DRINKING WATER:</b> Geographic location and use data  <b>HUMAN IMPACT:</b> Spatial distribution of contamination or drought

# PAPI-EX

Performance Application Programming Interface for Extreme-scale Environments

SI2-SSI-1450122

- Performance Measurement Library
- Cross-platform
- Widely used in Supercomputing Environments
- Find Bottlenecks in your code!
- Measure raw performance, architectural effects  
(Cache, Branch Predictor, etc.), Power and Energy
- Supports most modern computing hardware
- Companion tools: PAPI-ex, Counter Inspection Toolkit



Jack Dongarra, Heike Jagode, Anthony Danalis

University of Tennessee



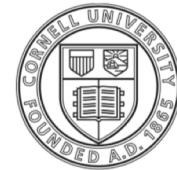
Vince Weaver

University of Maine



# SI<sup>2</sup>-SSE: Foundations for MATPOWER as an Extensible Tool for Power Systems Research and Education

Ray Zimmerman, Cornell University



## What is MATPOWER?

- Set of free, open-source, Matlab language tools
  - *compatible with MATLAB® and GNU Octave*
- For steady-state power system simulation and optimization, including:
  - power flow (PF)
  - extensible, optimal power flow (OPF)

### MATPOWER's Unique Combination

- free, open-source license (BSD)
- code that is easy to understand, customize
- state-of-the-art, high performance solvers
- ready-to-use realistic data included



### MATPOWER boosted to de facto standard

- benchmark platform for power systems research
- educational tool for power systems engineers and optimization

## Project Overview

Expand MATPOWER's future impact as a successful **research-enabling tool** for the problems of the power systems of the future by providing the **project infrastructure** and **core software architecture** needed to facilitate ongoing community-supported growth.

### MATPOWER Project Infrastructure

- Transition to fully collaborative **open development paradigm** with, public code repository, issue tracker, user and developer forums, contributor guidelines, public list of project descriptions

### MATPOWER Core Software Architecture

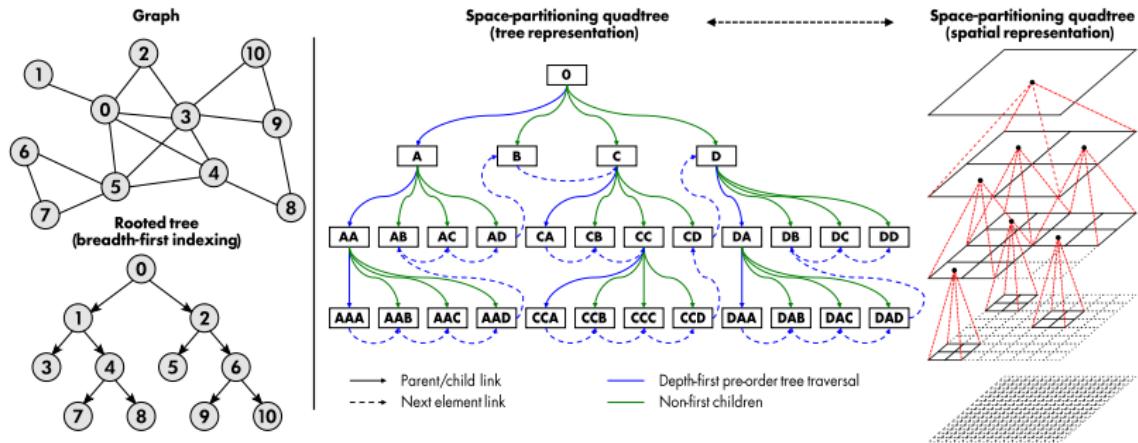
- Redesign core software around a general **modular architecture** to enable more flexible **user customization** and facilitate significant **user contributions**, while retaining and enhancing the **simplicity** that makes it attractive in education

CRII: OAC: A Hybrid Finite Element and Molecular  
Dynamics Simulation Approach for Modeling  
Nanoparticle Transport in Human Vasculature

**Ying Li**

## Summary

A software architecture challenge to design a unified set of high performance tree abstractions.

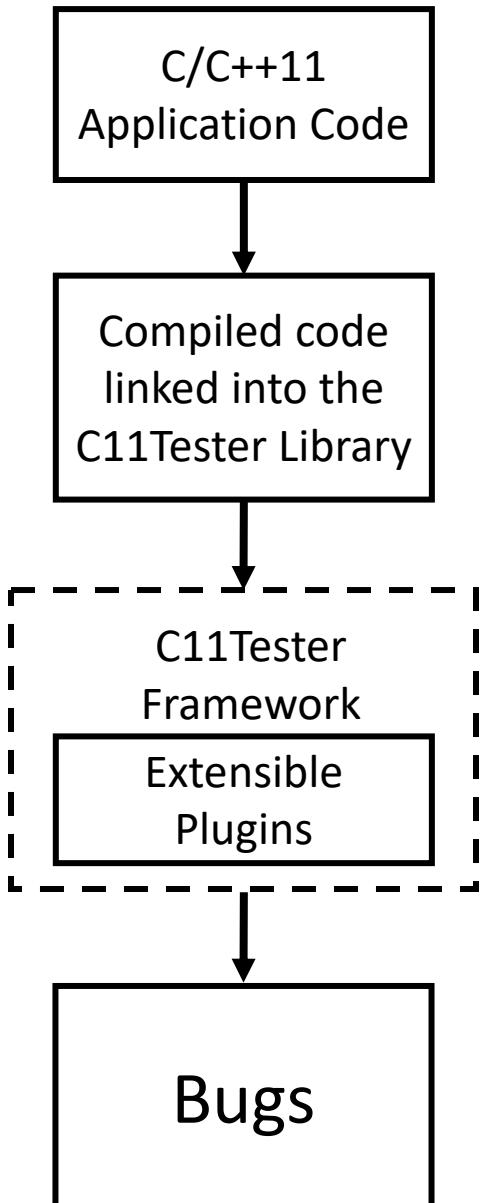


Representation	Depth	Arity	Traversal	Ordering
<ul style="list-style-type: none"> <li>■ Explicit</li> <li>■ Implicit</li> </ul>	<ul style="list-style-type: none"> <li>■ Static</li> <li>■ Dynamic</li> </ul>	<ul style="list-style-type: none"> <li>■ Static</li> <li>■ Dynamic</li> <li>■ Level dependent</li> </ul>	<ul style="list-style-type: none"> <li>■ Breadth first</li> <li>■ Depth first pre-order</li> <li>■ Depth first post-order</li> <li>■ Leaves only</li> </ul>	<ul style="list-style-type: none"> <li>■ Ordered</li> <li>■ Unordered</li> </ul>

# SI2-SSE: C11Tester: Scaling Testing of C/C++11 Atomics to Real-World Systems

Brian Demsky, University of California, Irvine

- Modern programming languages provide atomic operations
- Atomic operations:
  - Make it possible to build faster, more scalable data structures with stronger guarantees
  - Expose developers to complex behaviors that arise from CPU & compiler optimization
  - Are extremely difficult to use correctly
- C11Tester project is building tools to help developers effectively test code with atomic operations



*Bringing higher end computational tools to the bench scientist to accelerate the discovery process.*

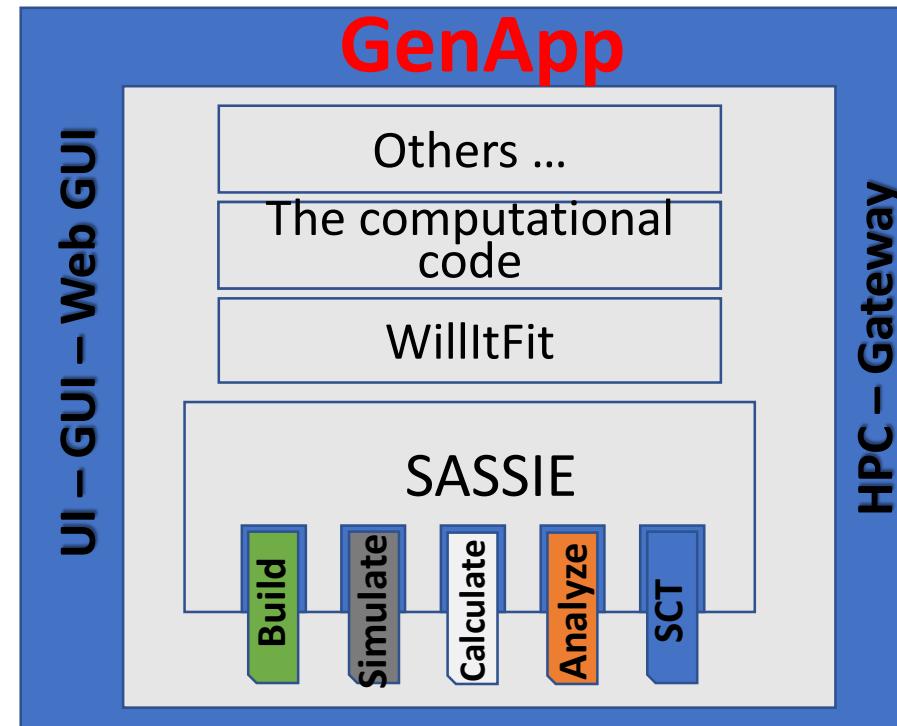
Current grant coming to a close.

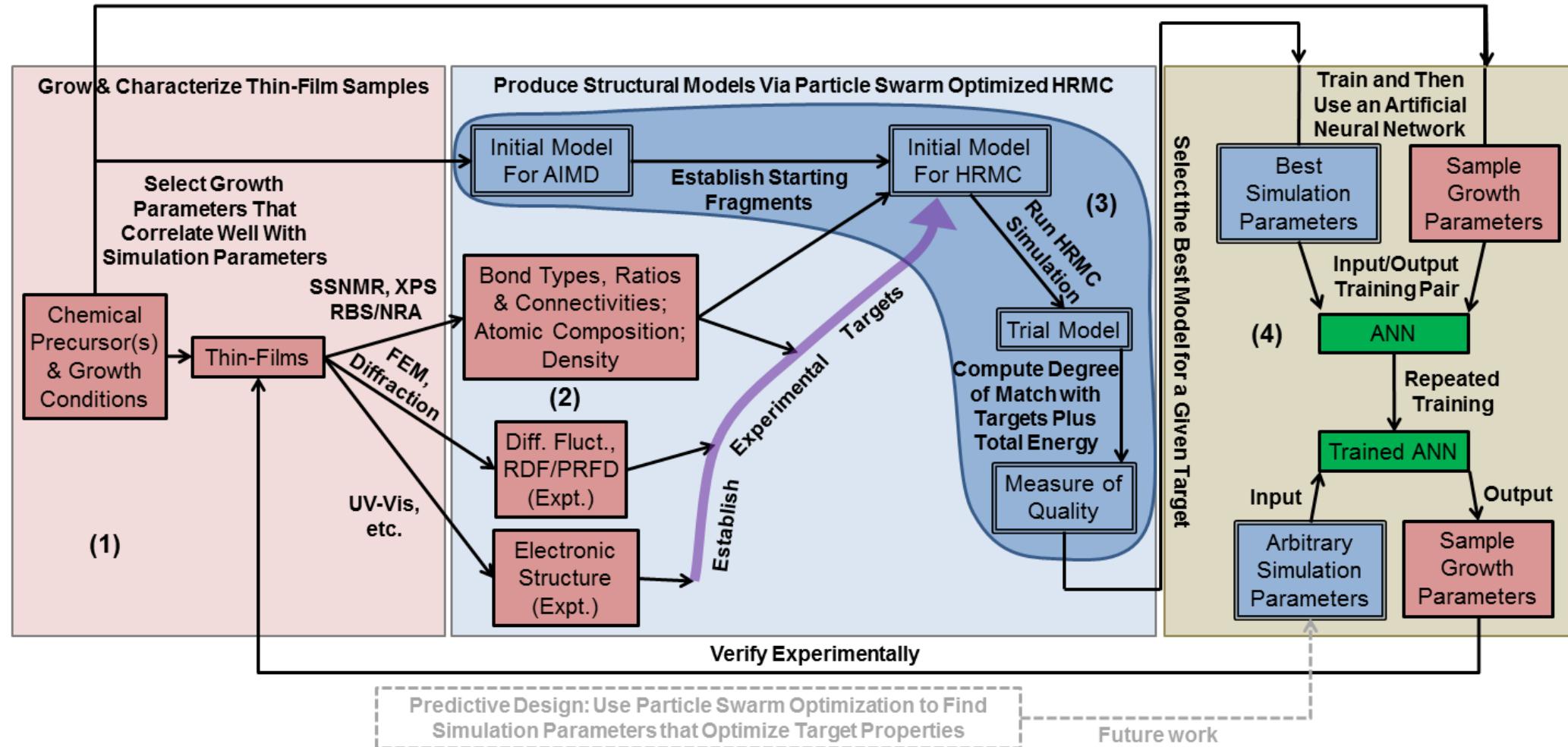
Working on transitioning to Community efforts and how to coordinate new efforts going forward.

Several new projects being proposed -- plus

- Networking/workshop grants
  - COST action proposal in Europe
  - looking at various US opportunities
  - Nurture involvement from major facilities

**OTHER IDEAS WELCOME**





The project includes (left) growth and characterization, (middle) iterative modeling, and (right) design training and validation. Single-frame red boxes represent experimental samples and data, while double-framed blue boxes represent computational products. The shaded region in the middle represents the application of particle swarm optimization. The general flow can be understood as: (1) growth of samples varied by composition and growth procedures; (2) experimental structural characterization; (3) iterative model simulation using characterization data; (4) ANN training to link simulation and growth parameters followed by predictive application of the ANN.

# Hearing the Signal through the Static: Realtime Noise Reduction in the Hunt for Binary Black Holes and other Gravitational Wave Transients

Sydney Chamberlin<sup>1</sup>, Reed Essick<sup>2</sup>, Patrick Godwin<sup>1</sup>, Chad Hanna<sup>1</sup>, Erik Katsavounidis<sup>3</sup>, Duncan Meacher<sup>1</sup>, Madeline Wade<sup>4</sup>

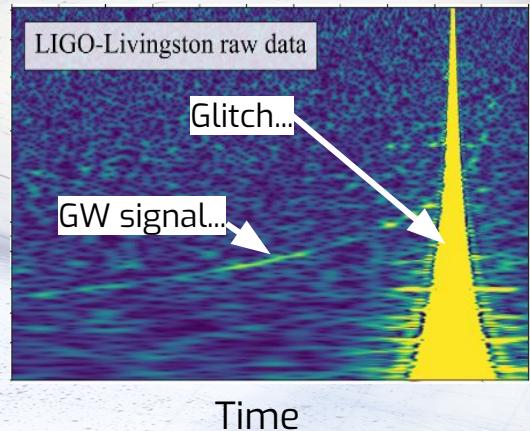
<sup>1</sup>The Pennsylvania State University, University Park, PA, 16801

<sup>2</sup>University of Chicago, Chicago, IL 60637

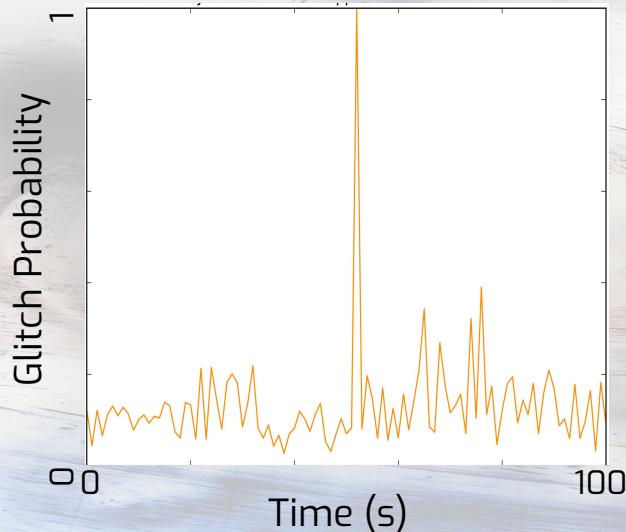
<sup>3</sup>Massachusetts Institute of Technology, Cambridge, MA 02139

<sup>4</sup>Kenyon College, Gambier, OH 43022

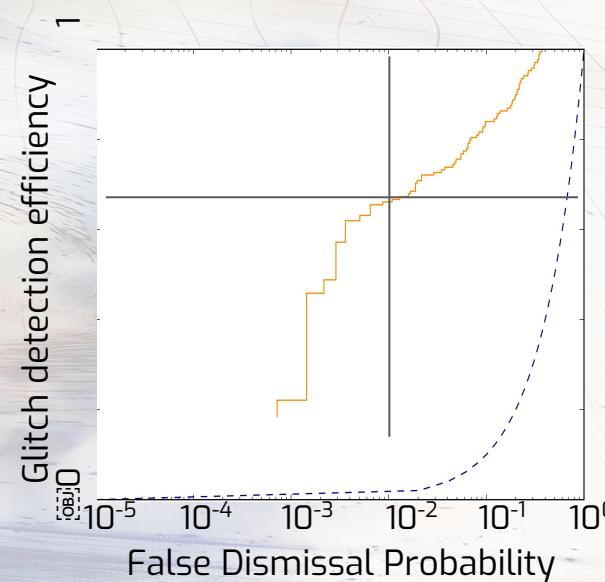
Real-time GW searches are plagued by "glitches". E.g., GW170817 - a binary neutron star merger - had a delayed alert because we had to deal with data quality issues.



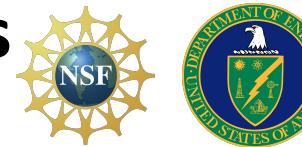
Goal: use machine learning to classify glitches in real-time in based on auxiliary information like seismometers, magnetometers, etc.



Currently can reject  $\frac{2}{3}$  of the glitches with a 1% false dismissal. Working to make it even better.



# RADICAL-Cybertools: Building Blocks for Workflow System Middleware



## Overview

**Motivation:** Sophisticated and scalable workflows have become essential for advances in computational science. In spite of the many successes of workflow systems, there is an absence of a reasoning framework for end-users to determine **which** systems to use, **when** and **why**. Workflows are increasingly a manifestation of the algorithmic and methodological advances; workflow users and workflow system developers are often the same. Workflow systems must be easily extensible so as to support diverse functionality and the proverbial “last mile customization”.

We advance the science of workflows and prevent workflow system “vendor lock-in” by formulating a **building blocks** approach to middleware for workflow systems grounded on four design principles of **self-sufficiency**, **interoperability**, **composability**, and **extensibility**. A building block has: (i) one or more modules implementing functionalities to operate on a set of explicitly defined entities; and (ii) two well-defined and stable interfaces, one for input and one for output.

### Properties of building blocks

- **Self-sufficiency:** design does not depend on the specificity of other building blocks
- **Interoperability:** can be used in diverse system architectures without semantic modifications
- **Composability:** its interfaces enable communication and coordination with other building blocks
- **Extensibility:** its functionalities and entities can be extended to support new requirements or capabilities

## RADICAL-Cybertools: An implementation of the Building Block Approach to Middleware

RADICAL-Cybertools are designed and implemented in accordance with the building block approach, spanning four functional levels:

(L4) **Workflow and Application Description:** Requirements and semantics of applications and workflows.

(L3) **Workload Management System (WLMS):** Applications devoid of semantic context are expressed as workloads.

(L2) **Task Runtime System (TRS):** Execution of the tasks of a workload.

(L1) **Resource:** Capabilities, availability and interfaces required by the tasks to be executed.

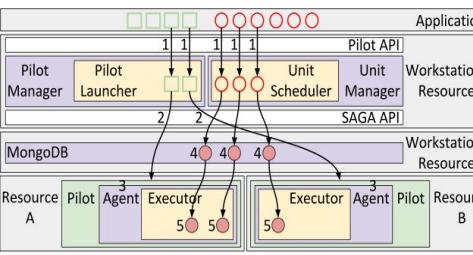
RADICAL-Cybertools are used at each level to support scalable, efficient and effective use of high-performance and distributed computing.

## (L2-L1) Interface to Resource

**RADICAL-SAGA (Simple API for Grid Applications):** Provides an interoperability layer that lowers the complexity of using distributed infrastructure whilst enhancing sustainability of distributed applications, services, and tools in the form of a Python API. By abstracting away the heterogeneity of the underlying systems, RADICAL-SAGA **simplifies access** to many distributed cyberinfrastructures such as XSEDE and OSG.

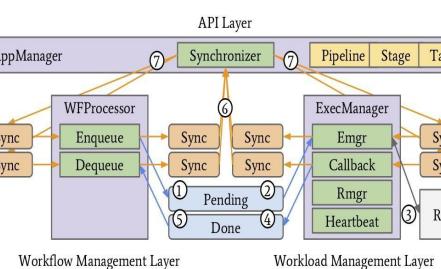
## (L2) Task Runtime Management

**RADICAL-Pilot:** Scalable pilot system for the simple and versatile execution of concurrent and distributed many-task applications on clusters, grids, and clouds. RADICAL-Pilot offers users a lightweight Python API to handle a variety of workloads—including MPI, multiprocess, multithreaded, CPU, and GPU tasks—and scheduling O(10k) tasks while marshalling O(10k) distributed cores.



## (L3) Workload Management

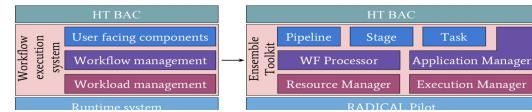
**Ensemble Toolkit:** Provides the ability to execute flexible combinations of **ensemble-based applications** on high-performance distributed computing resources. Ensemble Toolkit takes charge of where and how the workload is distributed: users only have to worry about what to run and when.



## (L4) Applications and Scientific Workflows

**ExTASY:** Enables sampling of complex macromolecules with molecular dynamics. It supports **high-performance** and **high-throughput** execution of molecular dynamic calculations, and analysis tools that provide runtime control over a simulation.

**HTBAC:** Enables the scalable, adaptive and automated calculation of the binding free energy on high-performance computing resources.



**RepEx:** Enables performing Replica Exchange simulations at a scale which is not attainable by stand-alone molecular dynamics applications. It uses RADICAL-Pilot for workload execution.

**ICEBERG:** Enables scalable image analysis on high-performance distributed computing for geoscience research. It provides a library based on extensible building-blocks that allows the integration of frameworks and algorithms seamlessly.

## Integration with existing systems

### SeisFlows

- Supports seismic inversion workflows on HPC machines, at scale
- We integrated **SeisFlow**
  - with **RADICAL-SAGA** (L1) to execute compute jobs
  - with **RADICAL-EnTK** (L3) to orchestrate tasks and data staging

### Atlas (Panda and Harvester)

- **Panda** is a WMS designed to support the distributed execution of workflows via pilots.
- **Harvester** is a service which provides pilot and workload management to Panda
- We integrated **Panda** and **RADICAL-Pilot** to improve its scaling on large HPC resources, and integrated **Harvester** and **RADICAL-Pilot** to provide scalable task execution on HPC machines

### Swift

- **Swift** is a language and a runtime system to execute workflows.
- We integrated **Swift** with **RADICAL-WLMS** (L3) to execute workloads concurrently on HPC and HTC resources.

### Fireworks

- **Fireworks** is a system that enables material science workflows
- We integrate **Fireworks** and **RADICAL-Pilot** (L2) to improve its scaling on HPC resources

# Large-Scale Causal Structure Learning

**W CDS&E: Statistical Methods for Discrete-Valued High-Dimensional Time Series**

UNIVERSITY of  
WASHINGTON

Ali Shojaie

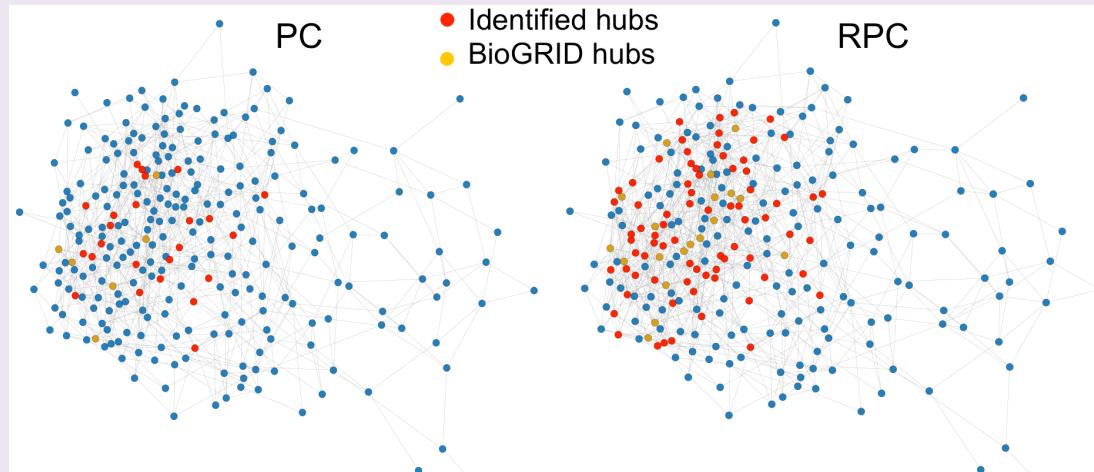


## Learning Directed Acyclic Graphs (DAGs)

- Generally NP-hard
- Not scalable
- Stringent assumptions

Large Network Properties

- Polynomial ( $n^3$ ) Alg.
- Scalable to large systems
- Relaxed assumptions



# SI2-SSE: ShareSafe: A Framework for Researchers and Data Owners to Help Facilitate Secure Graph Data Sharing

**Raheem Beyah**