

Broadening Access to Geospatial Capabilities

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Great opening!

80/20

Open

Virtual environment

Collaborative environment

Humanities 2.0

Toolkit

Participatory

Volunteer (VGI)

Multidisciplinary

Broader community needs



Acknowledgement

- Venkatesh Merwade
- Nelson Villoria
- Lan Zhao
- Larry Biehl
- Mike McLennan
- George Howlett
- Rob Campbell
- Rajesh Kalyanam
- Many others



Outline

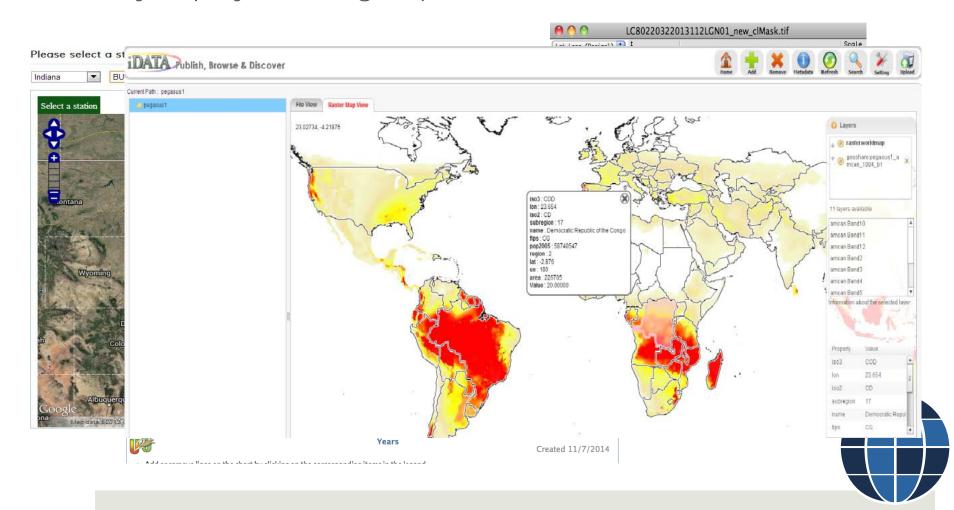
- Needs from user community
- Introduction of the GABBs project
- Technology
 - HUBzero
 - New geospatial additions
- Examples

Why do you care?



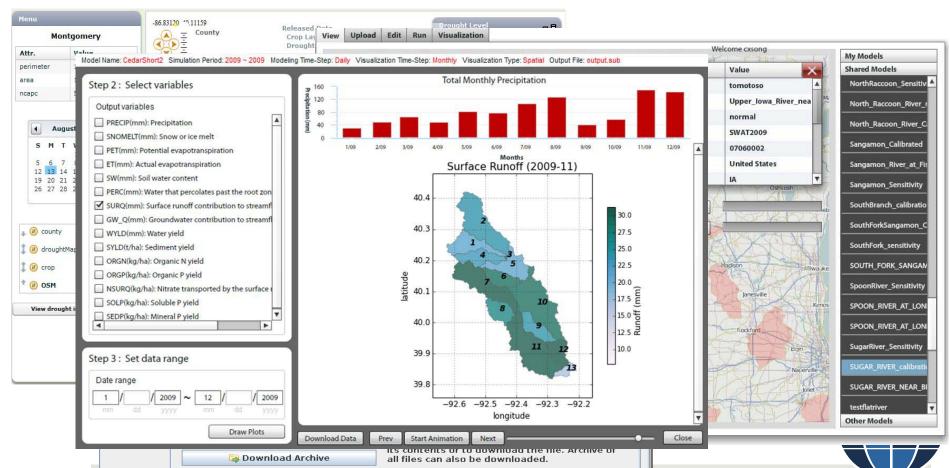
Driving Use Cases

Easy deployment of geospatial tools



Driving Use Cases (cont'd)

Multi-scale and multi-disciplinary data and modeling for addressing hydrologic and ag economic issues



The GABBs Project

Geospatial Modeling and Data Analysis Building Blocks

Making it easy to:

- create geospatial tools
- share these tools
- share geospatial data
- use other people's data



Key Outcomes

- The rapid tool development library RAPPTURE will support
 - geo-referenced data objects (maps, images, etc)
 - Easy way to share geospatial data, both in raw data, and visually and interactively
 - Easy way to share interactive tools that uses, and produces geospatial data
- Tool builder that supports geospatial data to further lower the barrier of creating interactive online tools
- Service interfaces to upload and share geospatial and other types of data in HUBzero
- Service interfaces to link tools and data
- Geospatial capabilities as part of core HUBzero open sor release

Funding

- A National Science Foundation grant
- Data Infrastructure Building Blocks (DIBBs) program
- □ GABBs: 1 of 4 implementation awards in 2013
- □ \$4.5M, 4 years (10/2013 9/2017)
- Collaboration with other DIBBs and DataNet awards



Team (14+)

Carol Song, PI & Project Director

Larry Biehl (CoPI, image processing and visualization)

Venkatesh Merwade (CoPI, hydrologic modeling)

Nelson Villoria (agricultural economics, sustainability)

Betsy Hillery (project manager)

Michael McLennan (HUBzero architect)

Rob Campbell (sr developer)

Leif Delgass (sr developer)

George Howlett (sr developer)

Lan Zhao (research scientist)

Rajesh Kalyanam (graduate student)

Hou-Jen Ko (graduate student)

Graduate students in scientific domains

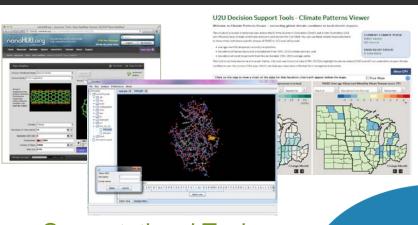


Building on prior work

- HUBzero (Rappture, graphics rendering, collaborative tools)
- iData (tool for self service data sharing and management)
- Multispec (tool for analyzing multispectral/hyperspectral image data)
- Geospatial hub projects (DRINET, Geoshare, WaterHUB, U2U etc)
- Leveraging software developed elsewhere
 - iRODS federated data management
 - Globus reliable transfer of large data



Platform for Scientific Collaboration



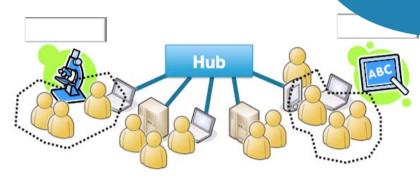
Download (JPG) Additional materials available (5) Version 1.0 - published on May 09, 2013 doi:10.4231/D32J68452 - cite this

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Computational Tools

hubzero

Databases / Publications



Learning Management

 $\nu(E) \sim E^{1+(d/\alpha)}$

 $v(E) \sim E^{1-(1/\alpha)}$

Who's Using HUBzero?

Supporting Purdue's largest research projects:



NEES: NSF \$105M - earthquake engr data (Ramirez)



NCN: NSF \$18M - nanotechnology (Klimeck/Lundstrom)



C3Bio: DoE EFRC \$20M - biofuels (McCann)



PRISM: DoE \$17M - mems devices (Murthy/Strachan)

Supporting Purdue infrastructure



Purdue University Research Repository (PURR) – data mgmt



PURDUE PurdueNExT / nanoHUB-U - online education

Supporting many other Purdue Projects









Outside Institutions



60+ Hubs for many disciplines

visitors

users

330,251	689,743
112,862	343,350
32,763	64,131
4,669	59,517
14,646	56,355
23,088	47,967
12,643	46,710
5,372	44,723
5,396	41,689
8,207	40,289
6,362	39,188
7,933	39,134



nanoHUB.org



nees.org



pharmaHUB.org



HABRIcentral.org



V_{HUB} vhub.org



GlobalHUB.org



cceHUB.org



PURR



iemhub.org



StemEdHub.org



ciHUB.org



molecularH

~1,500,000 visitors total

Courtesy of M. McLennan, Purdue University

Tool page

AgMIP Tool: A GEOSHARE tool for aggregating outputs from the AgMIP's Global Gridded Crop Modeling Initiative (Ag-GRID)

By Nelson Benjamin Villoria, Joshua Elliott, Christoph MÄ_ller, Jaewoo Shin 1, Lan Zhao 1

1. Purdue University

Aggregate the yield shocks provided by the AgMIP project from their original 30x30 min resolution to any user specified level.

Launch Tool

Version **1.2.4** - published on 05 Nov 2014

Open source: license | download

View All Supporting

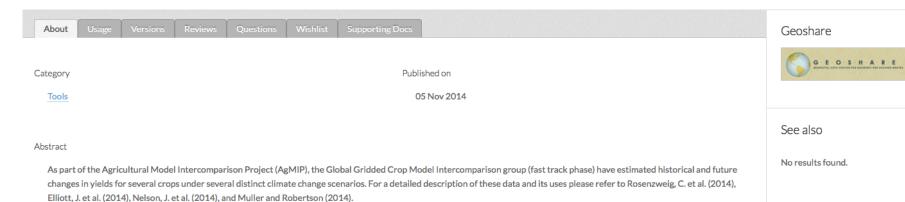
Documents

II 40 users, detailed usage

★ 0 review(s) (Review this)

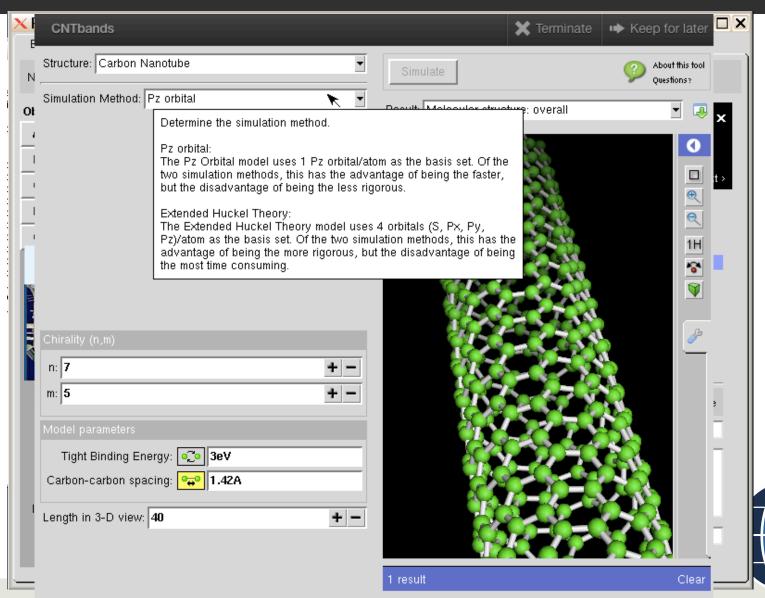
Q questions (Ask a question)

Ø wish(es) (Add a new wish)



The output archive comprises time series (1971-2099) generated by seven crop models (EPIC, GEPIC, pDDSAT, LPJmL, IMAGE-AEZ, PEGASUS, LPJ-GUESS), under a number of temperature trajectories from a suite of five global climate models (HadGEM2-ES, IPSL-CMSA-LR, MIROC-EXM-CHEM, GFDL-ESM2M, NorESM1-M) and five representative

Rappture Builder



Publication

Localization of MLL1 morphemes in mouse mm9 genomic DNA

By Minou Bina¹, Phillip Wyss¹, Daidong Wang¹, Xiaohui Song²

 Purdue University, Department of Chemistry 2. Purdue University, Academic Technologies and Rosen Center for Advanced Computing

Supplementary materials for the publication entitled "Discovery of MLL1 binding units, their localization to CpG Islands, and their potential function in mitotic chromatin." Bina M, Wyss P, et al. BMC Genomics. 14:927 (2013)

Listed in Datasets



Version **1.0** - published on Mar 31, 2014 doi:10.4231/R7KW5CXF - cite this

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About Supporting Docs Versions Reviews Questions Citations

Abstract

Mixed Lineage Leukemia 1 (MLL or MLL1) plays central roles in the regulation of protein-coding genes. Previous studies of mice have established that MLL1 controls key developmental pathways including the formation of body plan during the early stages of embryogenesis. Even though MLL1 is best known for catalyzing methylation of lysine 4 in histone H3 (H3K4), reported data indicate that MLL1 performs histone methyltransferase-independent functions including gene-bookmarking during mitosis, favoring genes that were highly transcribed during interphase, to accelerate reactivation of transcription following mitotic exit (Blobel et al., Mol Cell. 2009).

By analyzing results of reported DNA binding and chromatin immunoprecipitation assays, we have uncovered the DNA sequence elements that bind the MT-domain in MLL1: Bina et. al, BMC Genomics 14:927 (2013). We describe these elements as minimal units or morphemes: the smallest 'words' in DNA that selectively bind the MLL1 MT-domain. This publication provides a file (BINA_MLL_morphemes_mm9.bed) to display the position of MLL1 morphemes in the genome browser at UCSC. You can obtain a copy of the bed file (in text format) by clicking on download. Alternatively, you can use the link below to view the position of MLL1 morphemes directly on a track in the browser at UCSC.

While using the genome browser, select Dense to view morpheme positions in genomic DNA. Select pack when zooming to short DNA segments. If you use the data in your research, please cite the primary publication by Bina et al., published in BMC Genomics 14:927 (2013).

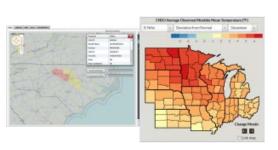
Open the position of MLL1 morphemes in the mouse built mm9 genomic DNA:

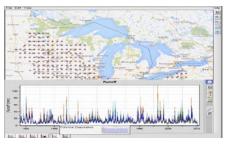


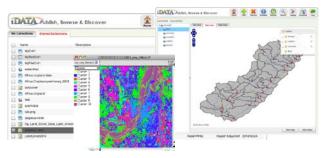
Building on HUBzero

- Extend RAPPTURE Toolkit
- Integrate map and spatial data rendering
- Adding new data objects to Rappture Builder
- Support geospatial data viewing and metadata in iData
 - Data management
 - Data sharing
 - Data quick view
 - Common processing
- Link data and tools in Hubzero
 - Dataset -> invoke tools
 - Tools -> result published and shared









Geospatial computation

Data presentation

Data Sharing

Data management Rendering Tool builder Remote servers

Layers

Geo-processing Data-Tool linkage

Maps

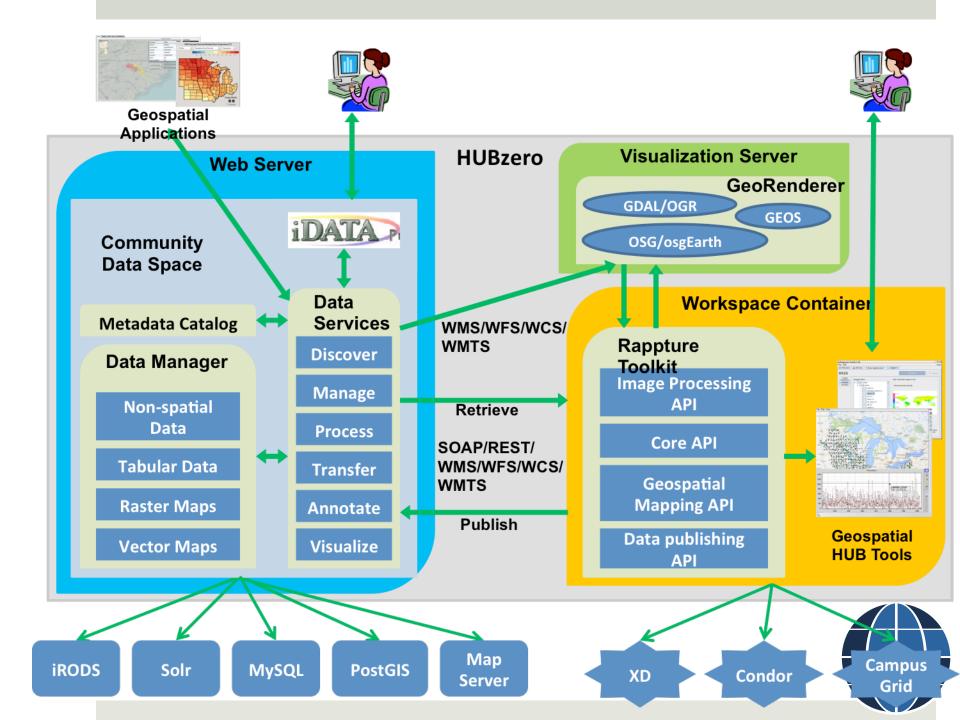
Data sharing Data presentation



Computation tools and online databases, Content publishing, Collaboration (group, project), Learning (courses, self-help), Support (tickets, Q&A), Community (forum, review, calendar)



Supercomputers, storage, map servers, geo-processing engines

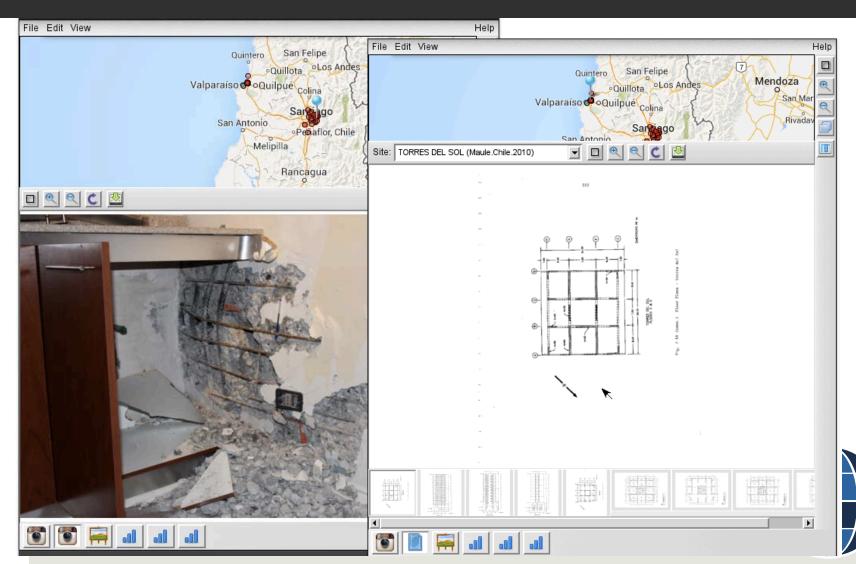


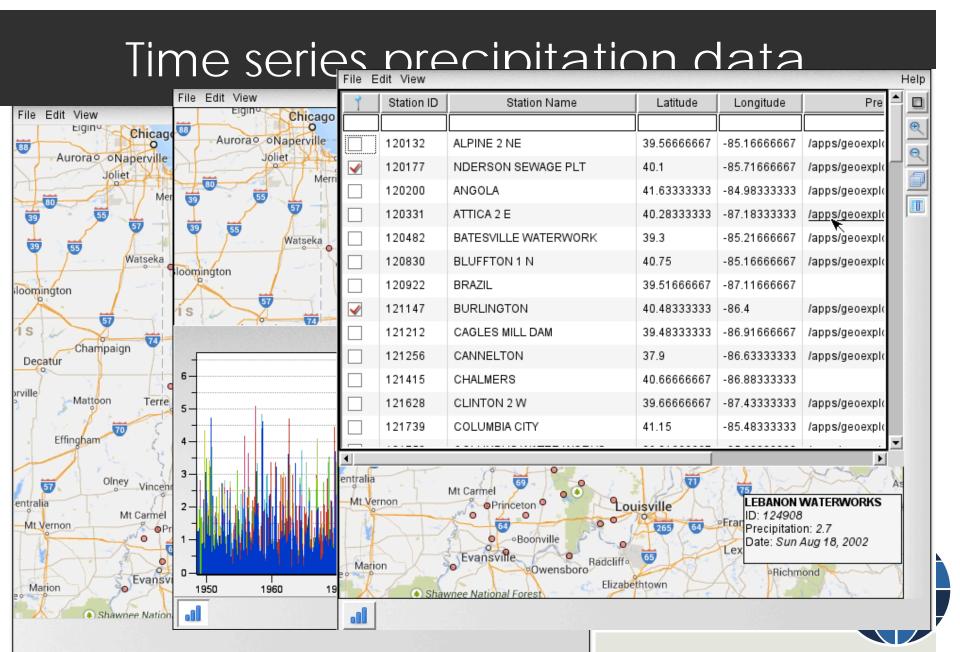
Examples

- Accessible via web browser
- Interactive
- Demonstrate capabilities

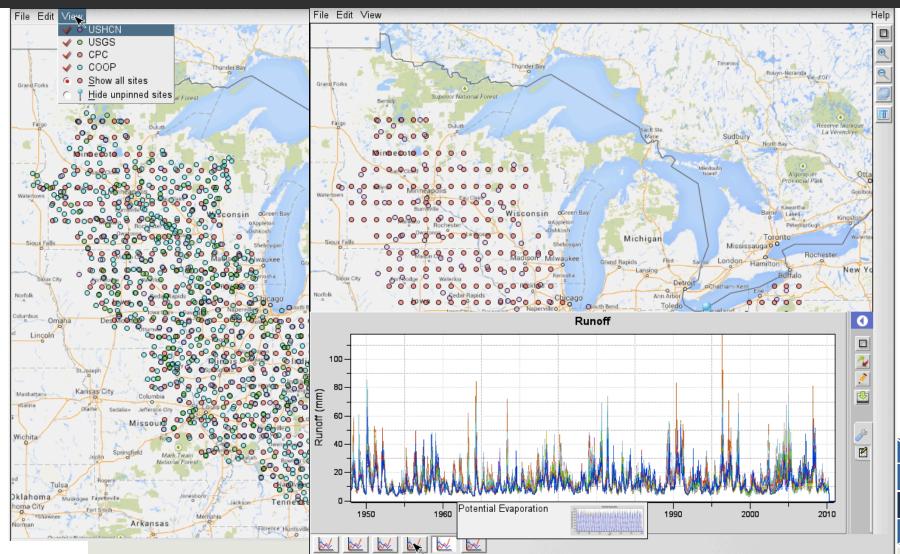


Chile Earthquake data



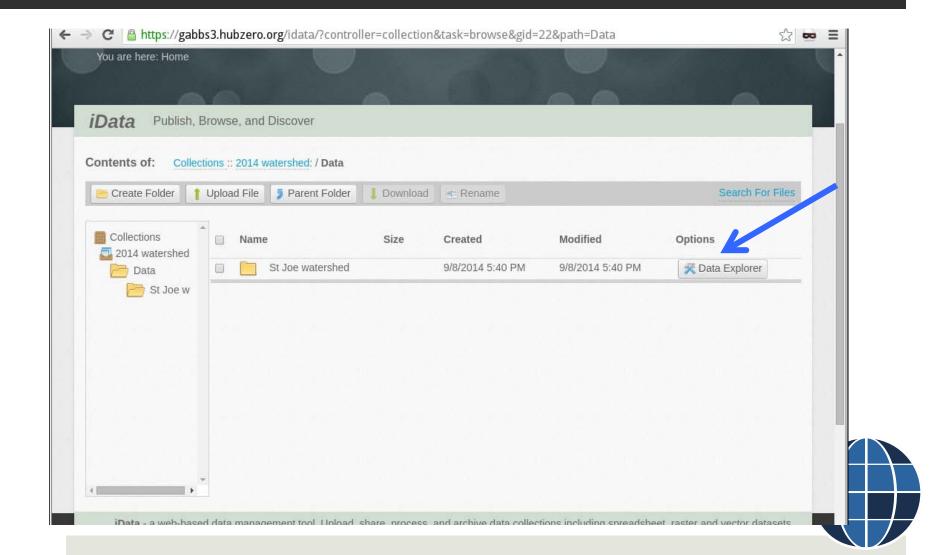


Upper MS River Basin & OH River Basin Data

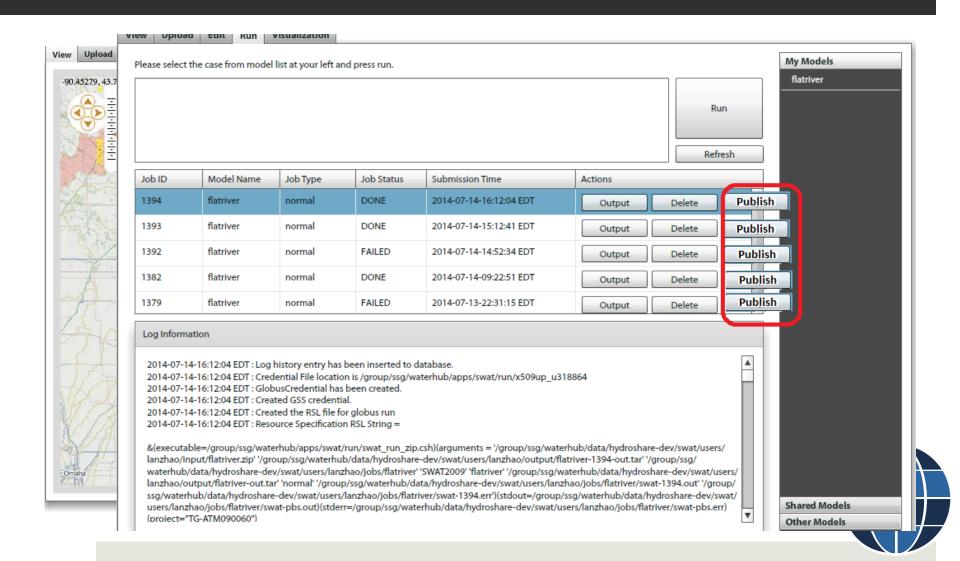




Start tools from datasets



Publish data from tools



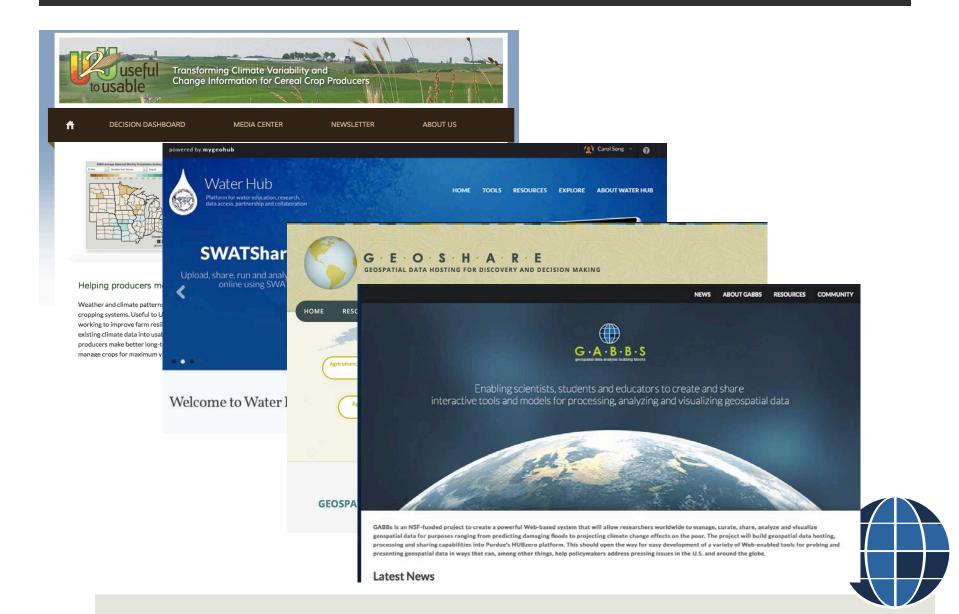
MyGeoHUB.org

- A hub for geospatial modeling, data analysis and visualization
- Hosting tools, datasets, groups and projects
- Free to sign up
- GABBs software will be installed on MyGeoHub.org
- Demonstration applications will be hosted
- User feedback and wishes are welcome!
- Build your own project, group, and community with Supergroups!





Super Groups



Collaboration

- Use cases
 - Datasets
 - applications
- Feedback, suggestions
- Participation
 - Share your work on MyGeoHub.org
- Ask questions

