

# Virtual Imaging Platform

## Enabling distributed computing for medical image simulation

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### Context

#### Goals

- Provide online access to medical image simulators
- Annotate data to facilitate retrieval and usage
- Support computing and data requirements

#### Challenges in biomed VO

- Hardware heterogeneity
- Job failures
- Data transfers (e.g. result merging)

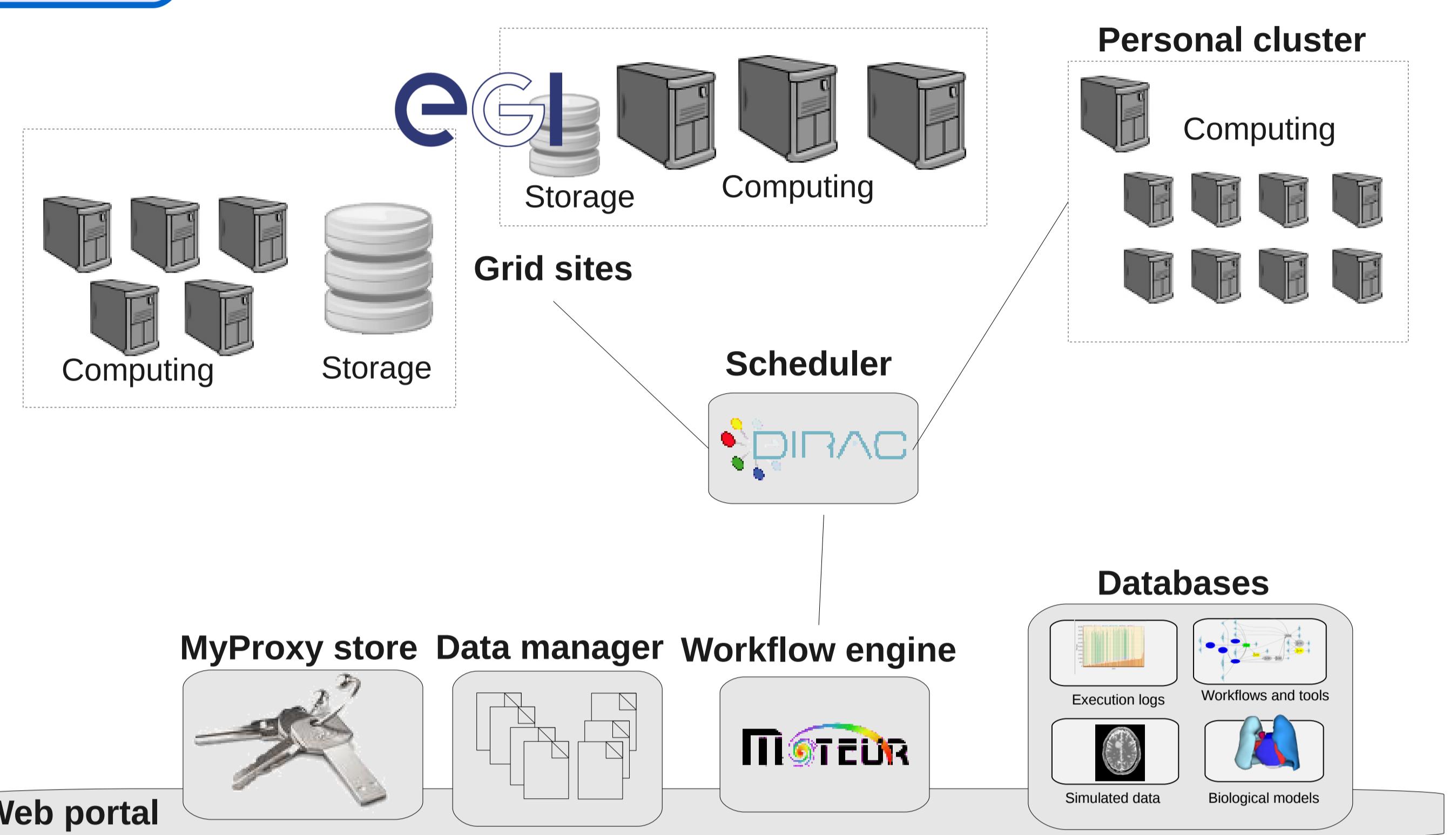
### Platform outline

#### Computing

- DIRAC pilot-job system
- Local personal clusters can be used
- Applications are MOTEUR workflows

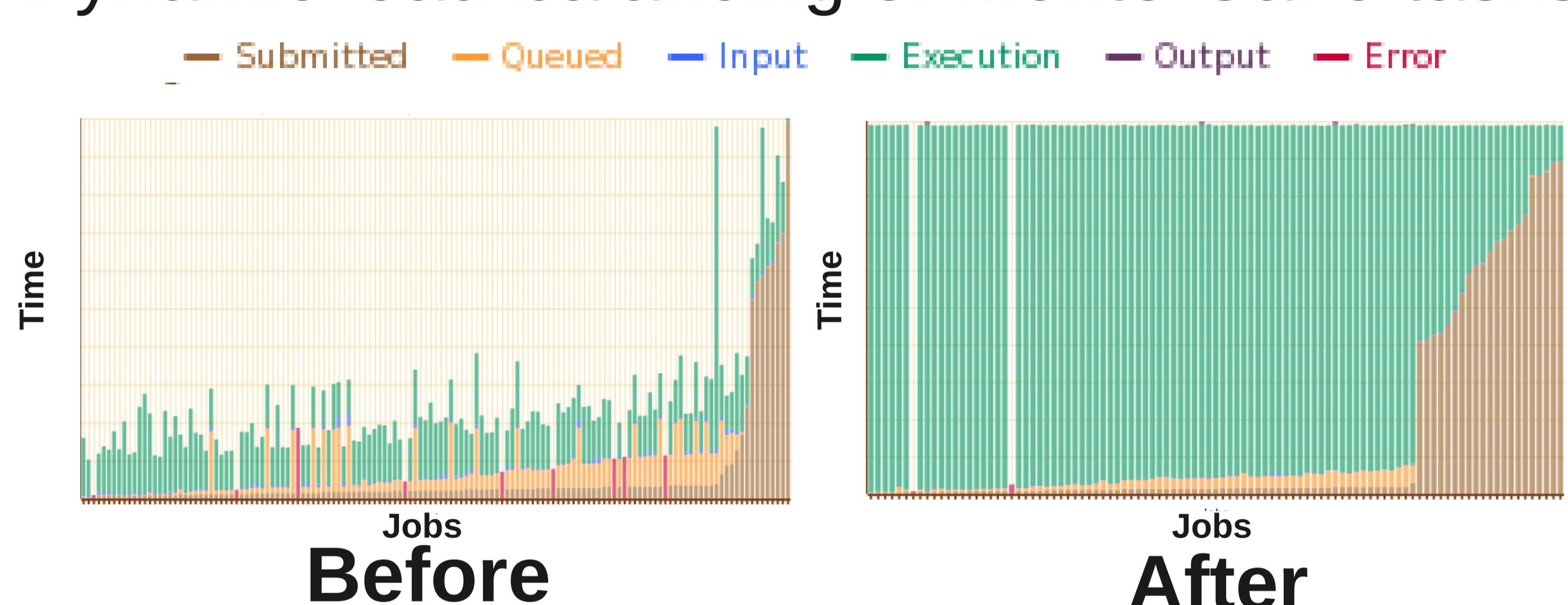
#### Data files

- Indexed in the Logical File Catalog
- Replicated on platform's Storage Element
- Cached in portal and worker nodes
- Biological models are semantically indexed



### Optimization

Dynamic load-balancing of Monte-Carlo tasks:



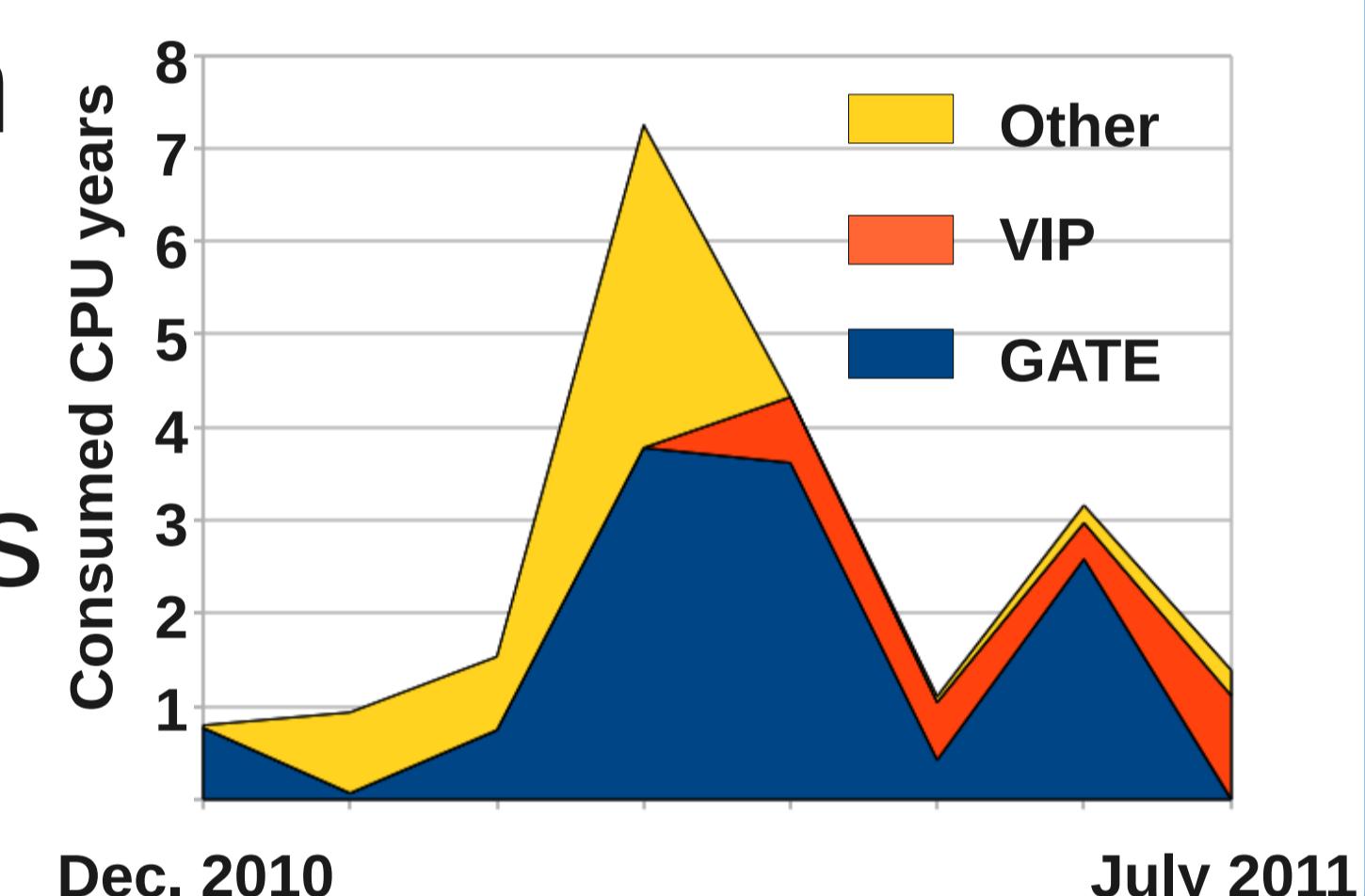
### Usage report

#### Achievements

- 9 regular users, 8 applications
- > 1 CPU year / month

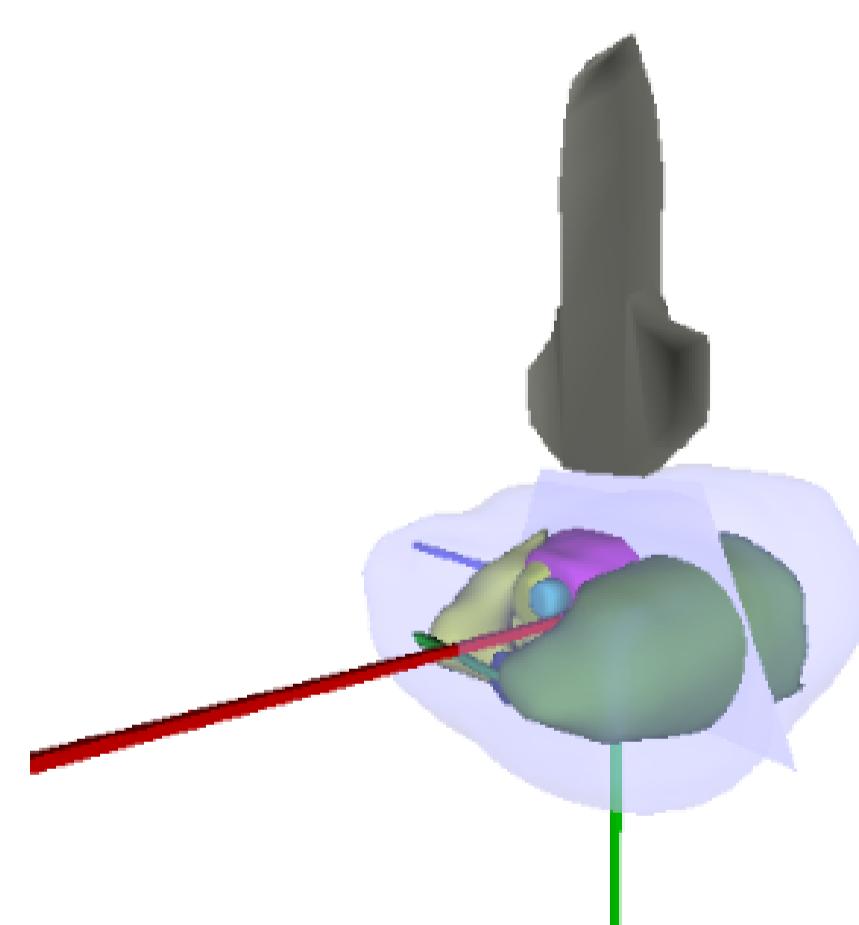
#### Main issues

- Scalability, robustness
- Administration
- X509 authentication

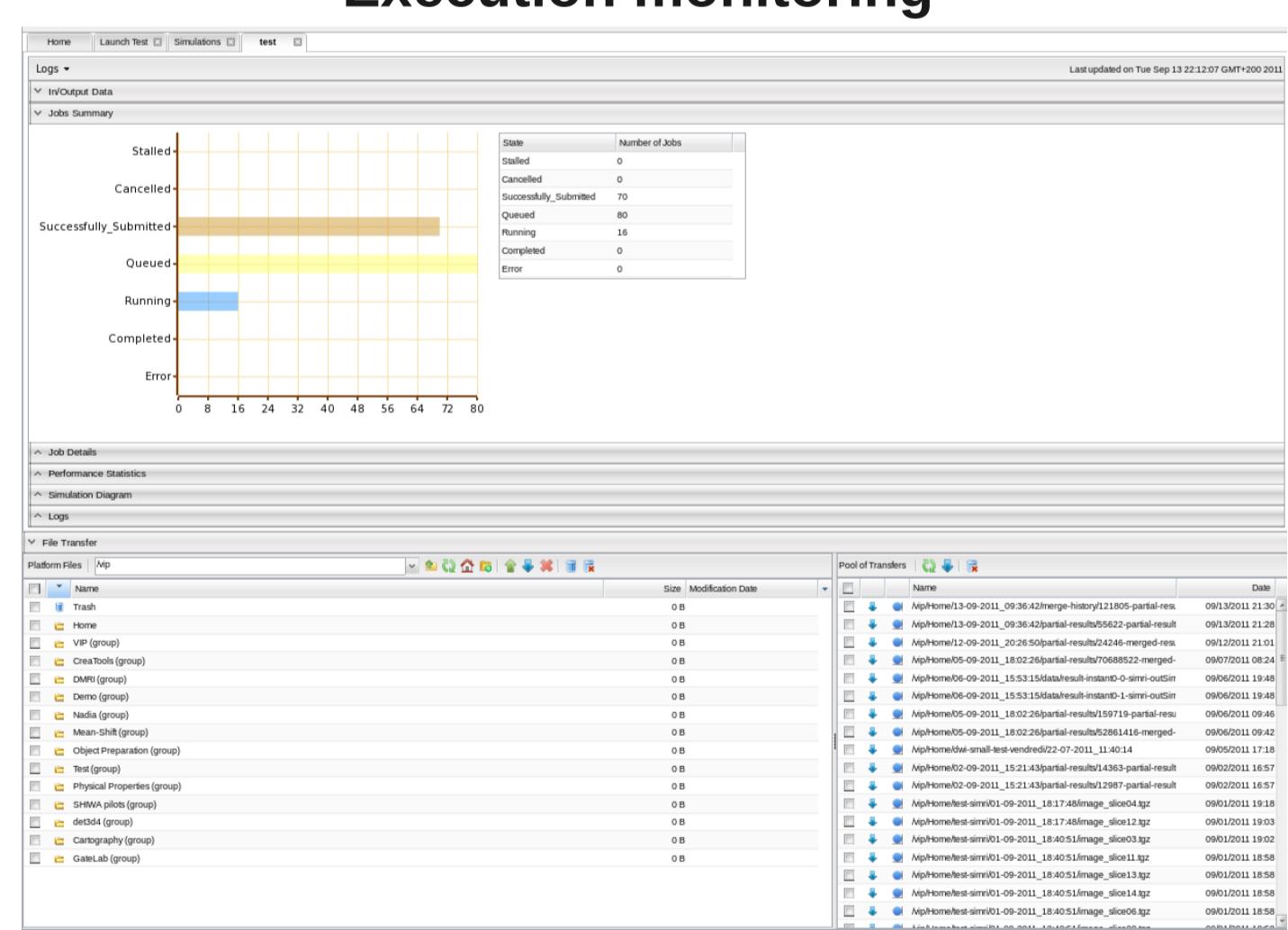


### Snapshots

#### Simulation definition



#### Execution monitoring



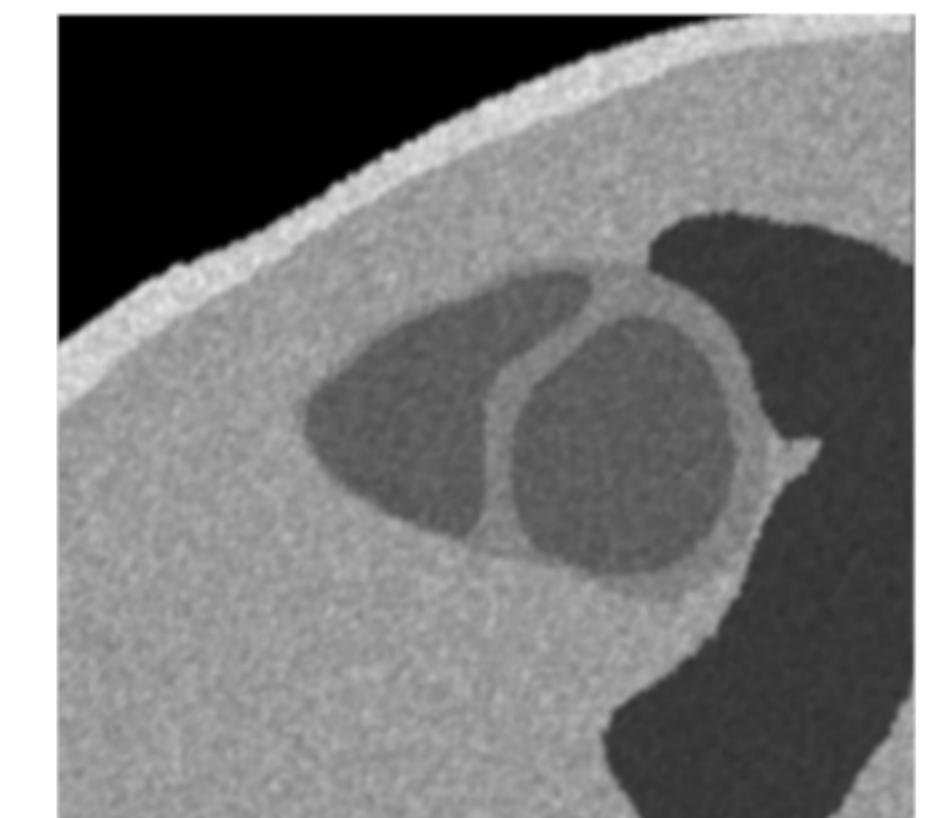
#### Simulated ultrasound image



#### Simulated X-Ray image



#### Simulated MRI



### On-going work

#### Grid-related

- Checkpoint / restart workflows
- Improved merging of job results
- Assisted administration

#### Semantic annotations

- Simulated data
- Simulation tools
- Integration in workflow designer

### Credits

Platform development: Sorina Camarasu-Pop, Rafael Ferreira da Silva, Tristan Glatard, Baptiste Grenier, David Manset, Jérôme Revillard

Semantic repositories: Germain Forestier, Alban Gaignard, Bernard Gibaud, Johan Montagnat, Javier Rojas Balderrama

Simulators and applications: Christian Barillot, Hugues Benoit-Cattin, Frédéric Cervenansky, Patrick Clarysse, Denis Friboulet,

Patrick Hugonnard, Carole Lartizien, Hervé Liebgott, Adrien Marion, Joachim Tabary