

Pegasus Workflow Management System

Karan Vahi

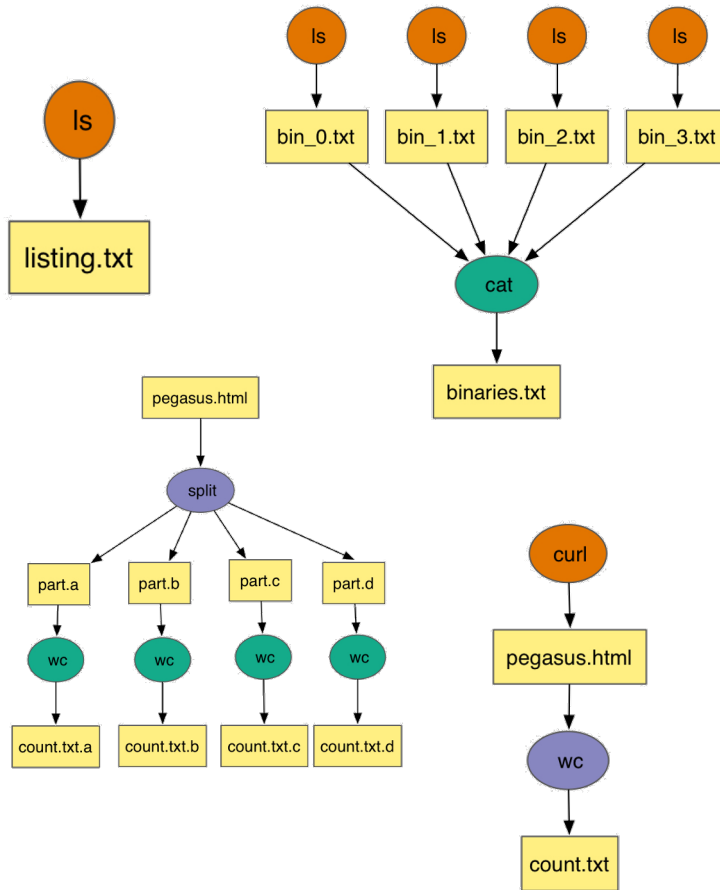
USC Information Sciences Institute



Benefits of Scientific Workflows

(from the point of view of an application scientist)

- Conducts a series of computational tasks.
 - Resources distributed across Internet.
- Chaining (outputs become inputs) replaces manual hand-offs.
 - Accelerated creation of products.
- Ease of use - gives non-developers access to sophisticated codes.
 - Avoids need to download-install-learn how to use someone else's code.
- Provides framework to host or assemble community set of applications.
 - Honors original codes. Allows for heterogeneous coding styles.
- Framework to define common formats or standards when useful.
 - Promotes exchange of data, products, codes. Community metadata.
- Multi-disciplinary workflows can promote even broader collaborations.
 - E.g., ground motions fed into simulation of building shaking.
- Certain rules or guidelines make it easier to add a code into a workflow.



Challenges of Workflow Management

Challenges across domains

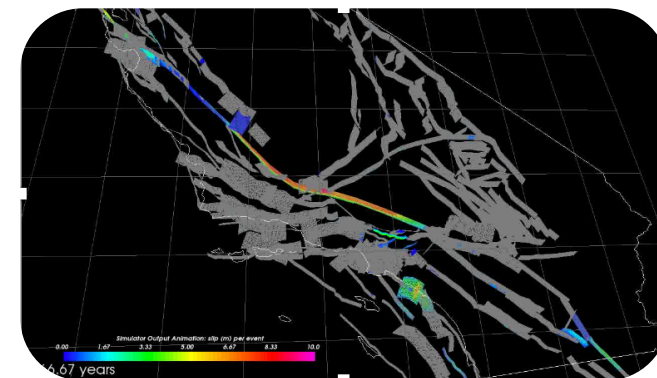
- Need to describe complex workflows in a simple way
- Need to access distributed, heterogeneous data and resources (heterogeneous interfaces)
- Need to deal with resources/software that change over time

Our focus

- Separation between workflow description and workflow execution
- Workflow planning and scheduling (scalability, performance)
- Task execution (monitoring, fault tolerance, debugging)
- Provide additional assurances that a scientific workflow is not accidentally or maliciously tampered with during its execution.



Sky mosaic, IPAC, Caltech



Earthquake simulation, SCEC, USC

Pegasus Workflow Management System

- Operates at the level of files and individual applications
- Allows scientists to describe their computational processes (workflows) at a logical level
- Without including details of target heterogeneous CI (portability)
- Scalable to $O(10^6)$ tasks, TBs of data
- Captures provenance and supports reproducibility
- Includes monitoring and debugging tools

Workflow Listing



Show results for all 1

Workflow Label	Submit Host	Submit Directory	State	Submitted On
split	workflow.isl.edu	/ifs/cog3/cog/home/pegasus01/learningsplit/pegasus01/pegasus01/run0008	Running	Fri, 23 Oct 2015 16:04:00
split	workflow.isl.edu	/ifs/cog3/cog/home/pegasus01/learningsplit/pegasus01/pegasus01/run0004	Failed	Fri, 23 Oct 2015 15:05:01
diamond	workflow.isl.edu	/ifs/cog3/cog/home/pegasus01/learningsplit/pegasus01/pegasus01/diamond/run0002	Successful	Fri, 23 Oct 2015 15:05:17
split	workflow.isl.edu	/ifs/cog3/cog/home/pegasus01/learningsplit/pegasus01/pegasus01/run0003	Failed	Fri, 23 Oct 2015 15:04:15
split	workflow.isl.edu	/ifs/cog3/cog/home/pegasus01/learningsplit/pegasus01/pegasus01/run0002	Successful	Fri, 23 Oct 2015 15:04:44
process	workflow.isl.edu	/ifs/cog3/cog/home/pegasus01/learningsplit/pegasus01/pegasus01/process/run0001	Successful	Fri, 23 Oct 2015 15:00:28
pipeline	workflow.isl.edu	/ifs/cog3/cog/home/pegasus01/learningsplit/pegasus01/pegasus01/pipeline/run0001	Successful	Fri, 23 Oct 2015 15:00:28
merge	workflow.isl.edu	/ifs/cog3/cog/home/pegasus01/learningsplit/pegasus01/pegasus01/merge/run0001	Successful	Fri, 23 Oct 2015 15:00:15
diamond	workflow.isl.edu	/ifs/cog3/cog/home/pegasus01/learningsplit/pegasus01/pegasus01/diamond/run0001	Successful	Fri, 23 Oct 2015 15:00:06
split	workflow.isl.edu	/ifs/cog3/cog/home/pegasus01/learningsplit/pegasus01/pegasus01/run0001	Successful	Fri, 23 Oct 2015 14:59:50

Showing 1 to 10 of 10 entries

Statistics

Workflow Wall Time	12 mins 23 secs
Workflow Cumulative Job Wall Time	9 mins 34 secs
Cumulative Job Walltime as seen from Submit Side	9 mins 35 secs
Workflow Cumulative Badput Time	9 mins 23 secs
Cumulative Job Badput Walltime as seen from Submit Side	9 mins 20 secs
Workflow Retries	1

Workflow Statistics

Type	Succeeded	Failed	Incomplete	Total	Retries	Total + Retries
Tasks	5	0	0	5	0	5
Jobs	16	0	0	16	2	18
Sub Workflows	0	0	0	0	0	0

Type	Succeeded	Failed	Incomplete	Total	Retries	Total + Retries
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Job Breakdown Statistics

Job Statistics

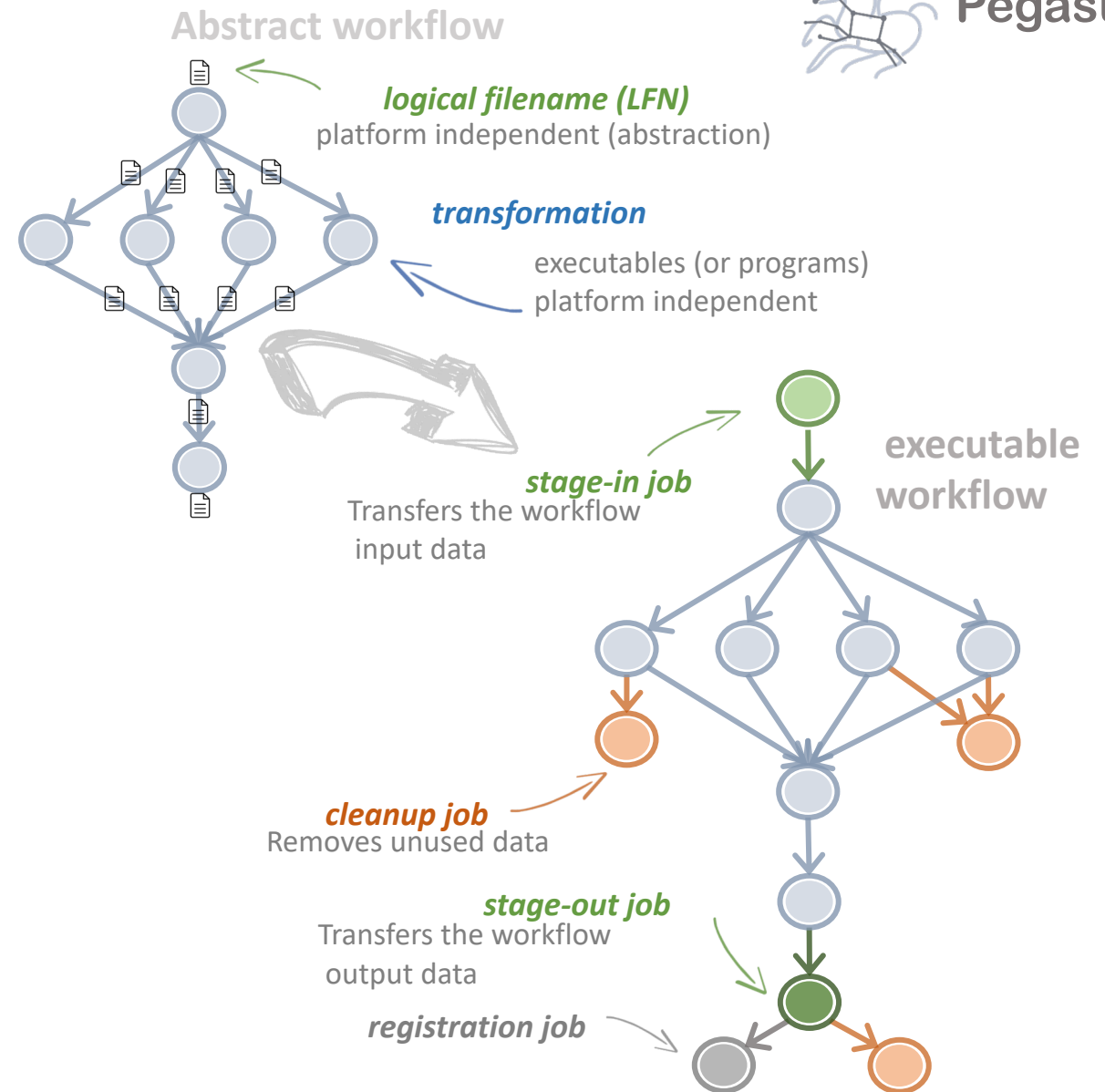
Composition in Python, R, Java, Perl, Jupyter Notebook

hubzero

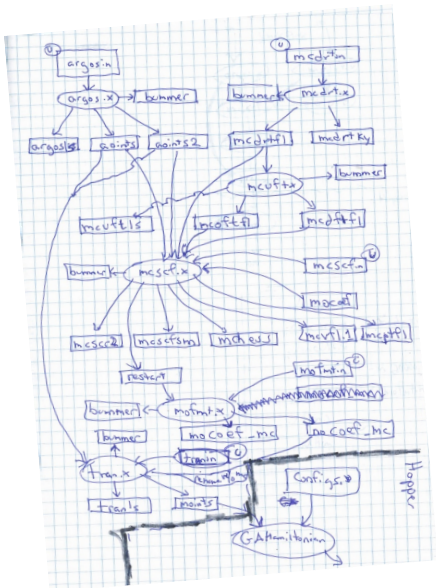
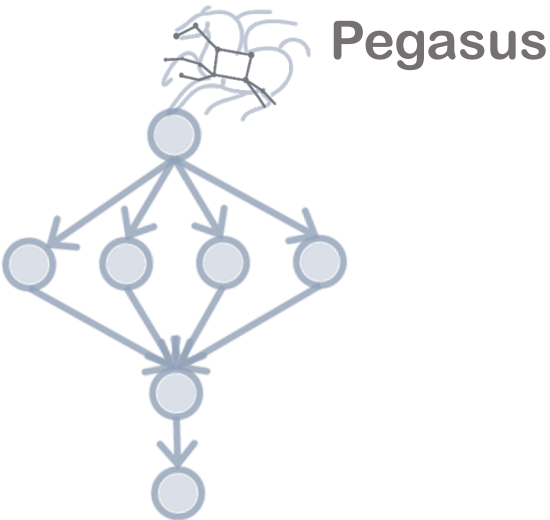


Pegasus Concepts

- Users describe their pipelines in a **portable** format called Abstract Workflow, **without worrying** about **low level execution** details.
- Workflows are DAGs
 - Nodes: jobs, edges: dependencies
 - No while loops, no conditional branches
 - Jobs are standalone executables
- Pegasus takes this and **generates an executable workflow** that
 - has **data management** tasks added
 - **transforms** the workflow for **performance** and **reliability**



Pegasus also provides tools to generate the workflow descriptions



```
#!/usr/bin/env python
from Pegasus.DAX3 import *
import sys
import os

# Create an abstract dag
dax = ADAG("hello_world")

# Add the hello job
hello = Job(namespace="hello_world",
            name="hello", version="1.0")
b = File("f.b")
hello.uses(a, link=Link.INPUT)
hello.uses(b, link=Link.OUTPUT)
dax.addJob(hello)

# Add the world job (depends on the hello job)
world = Job(namespace="hello_world",
            name="world", version="1.0")
c = File("f.c")
world.uses(b, link=Link.INPUT)
world.uses(c, link=Link.OUTPUT)
dax.addJob(world)

# Add control-flow dependencies
dax.addDependency(Dependency(parent=hello,
                              child=world))

# Write the DAX to stdout
dax.writeXML(sys.stdout)
```



```
<?xml version="1.0" encoding="UTF-8"?>
<!-- generator: python -->
<adag xmlns="http://pegasus.isi.edu/schema/DAX"
      version="3.4" name="hello_world">

  <!-- describe the jobs making
  up the hello world pipeline -->
  <job id="ID0000001" namespace="hello_world"
       name="hello" version="1.0">

    <uses name="f.b" link="output"/>
    <uses name="f.a" link="input"/>
  </job>

  <job id="ID0000002" namespace="hello_world"
       name="world" version="1.0">

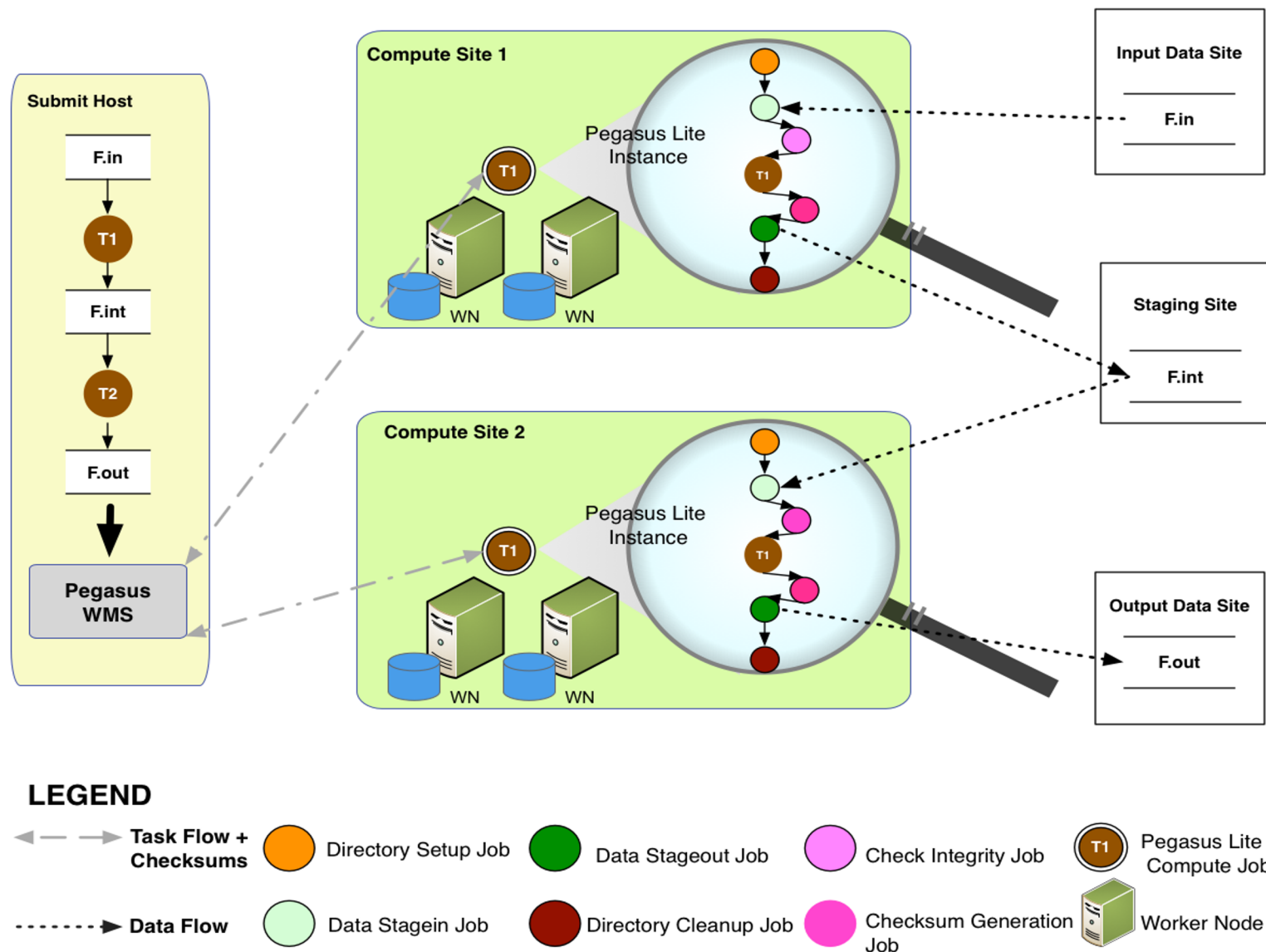
    <uses name="f.b" link="input"/>
    <uses name="f.c" link="output"/>
  </job>

  <!-- describe the edges in the DAG -->
  <child ref="ID0000002">
    <parent ref="ID0000001"/>
  </child>
</adag>
```

DAX

Pegasus Deployment

- **Workflow Submit Node**
 - Pegasus WMS
 - HTCondor
- **One or more Compute Sites**
 - Compute Clusters
 - Cloud
 - OSG
- **Input Sites**
 - Host Input Data
- **Data Staging Site**
 - Coordinate data movement for workflow
- **Output Site**
 - Where output data is placed



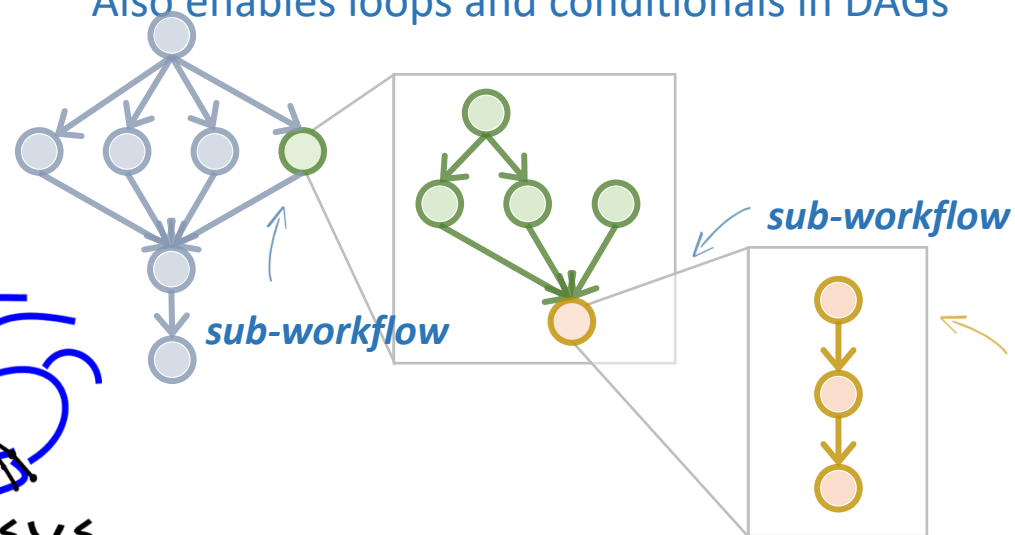
Pegasus Optimizations



Pegasus

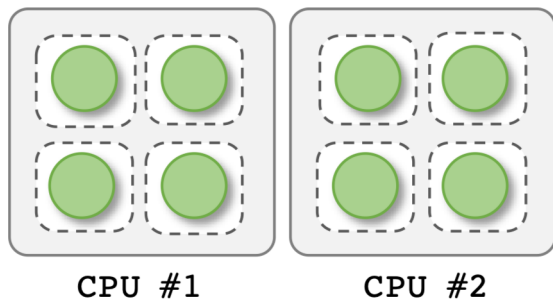
Hierarchical workflows

Enacts the execution of **millions of tasks**
Also enables loops and conditionals in DAGs

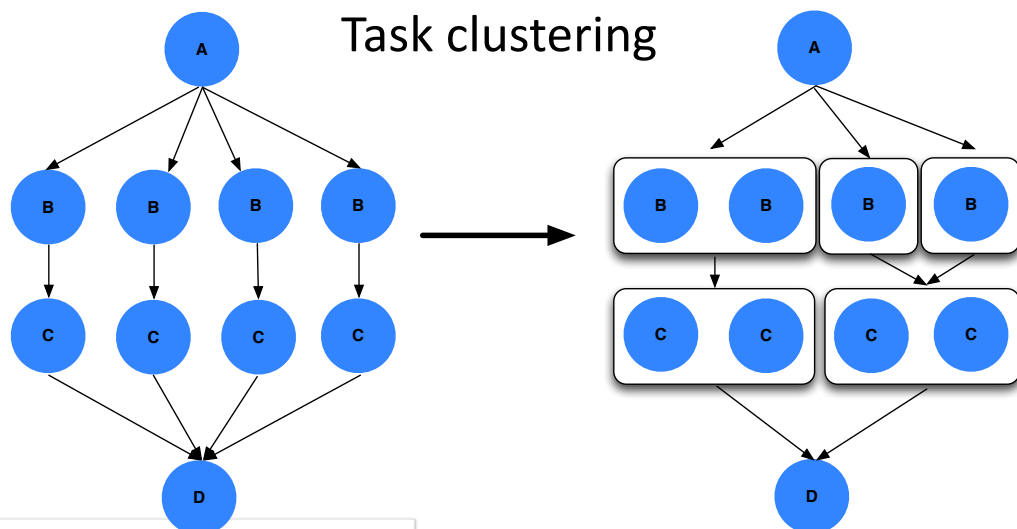


pegasus

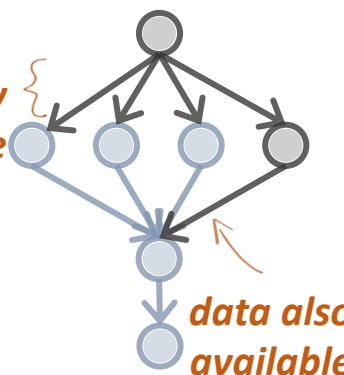
Task-resource co-allocation



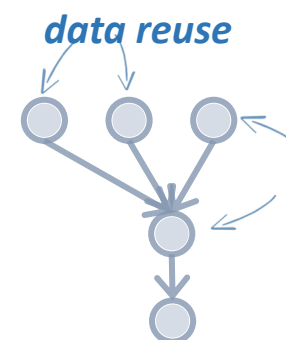
Task clustering



data already available



workf
low
reduc
tion



Jobs which output data is already available are pruned from the DAG

modern workflow optimizations

well-known optimizations

HTCondor I/O (HTCondor pools, OSG, ...)

Worker nodes do not share a file system

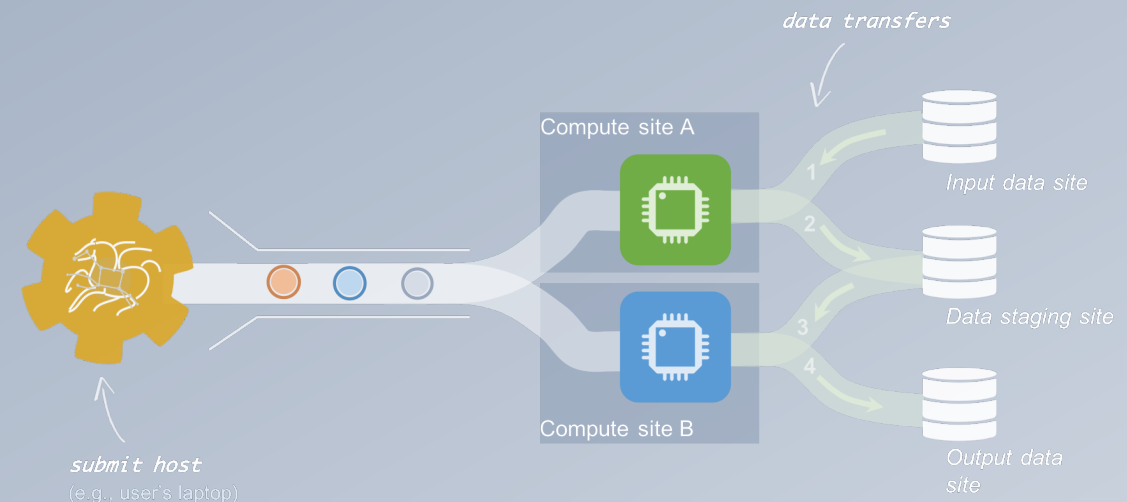
Data is pulled from / pushed to the submit host via HTCondor file transfers

Staging site is the submit host

Non-shared File System (clouds, OSG, ...)

Worker nodes do not share a file system

Data is pulled / pushed from a staging site, possibly not co-located with the computation



Shared File System (HPC sites, XSEDE, Campus clusters, ...)

I/O is directly against the shared file system

pegasus-transfer

Pegasus' internal data transfer tool with support for a number of different protocols



Pegasus

Directory creation, file removal

If protocol can support it, also used for cleanup

Two stage transfers

e.g., GridFTP to S3 = GridFTP to local file, local file to S3

Parallel transfers

Automatic retries

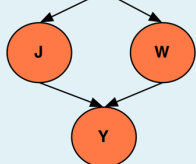
Credential management

Uses the appropriate credential for each site and each protocol (even 3rd party transfers)

HTTP
SCP
GridFTP
Globus
Online
iRods
Amazon S3
Google
Storage
SRM
FDT
Stashcp
Rucio
Webdav
cp
ln -s

Data Flow for LIGO Pegasus Workflows in OSG

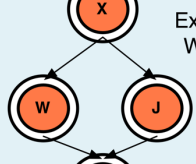
SUBMIT HOST **x** Abstract Workflow



Pegasus Planner

Workflow Setup Job

Workflow Stagein Job



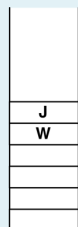
Executable Workflow

Workflow Stageout Job

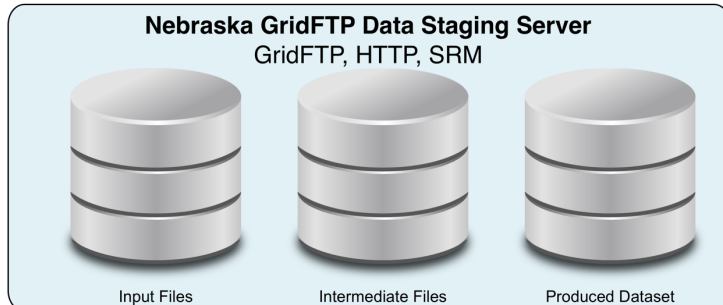
Data Cleanup Job

Condor Schedd Queue

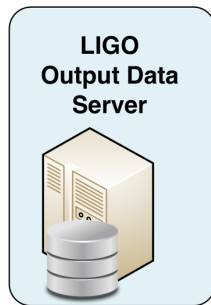
Condor DAGMan



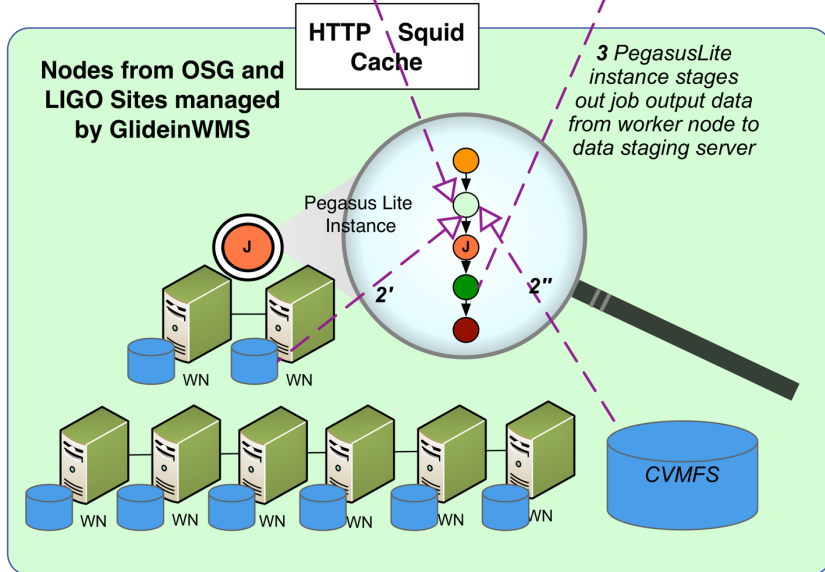
1 Workflow Stagein Job stages in the input data for workflow from user server



2 PegasusLite instance looks up input data on the compute node/ CVMFS If not present, stage-in data from remote data staging server



4 Workflow Stageout Job stages produced data from data staging server to LIGO Output Data Server



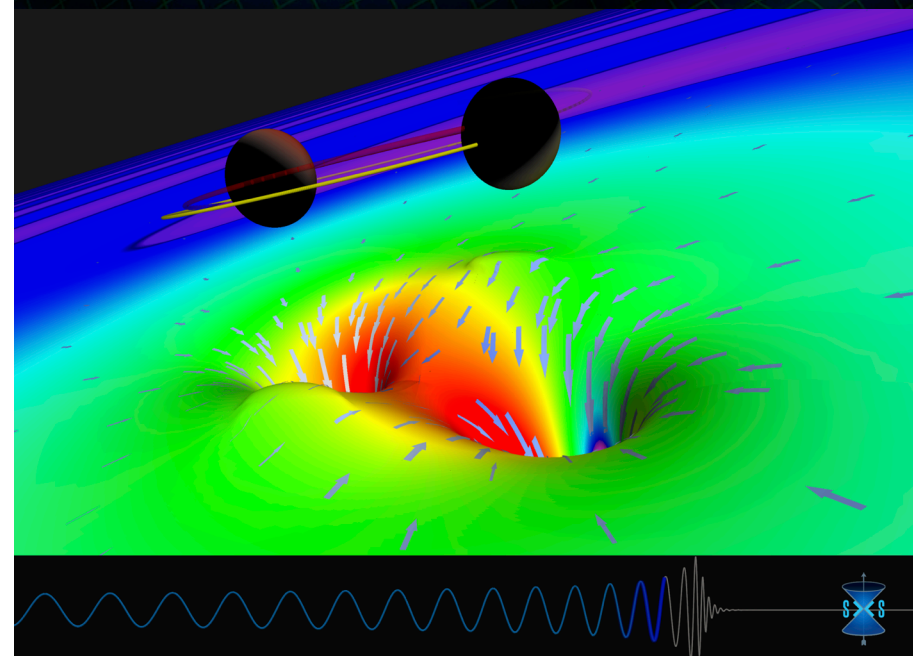
LEGEND

- Directory Setup Job
- Data Stageout Job
- Pegasus Lite Compute Job
- Data Stagein Job
- Directory Cleanup Job
- Worker Node

Advanced LIGO – Laser Interferometer Gravitational Wave Observatory

First gravitational wave detection:
21k Pegasus Workflows
107M tasks

Executed on LIGO Data Grid,
Open Science Grid and XSEDE



Challenges to Scientific Data Integrity

Modern IT systems are not perfect - errors creep in.

At modern “Big Data” sizes we are starting to see checksums breaking down.

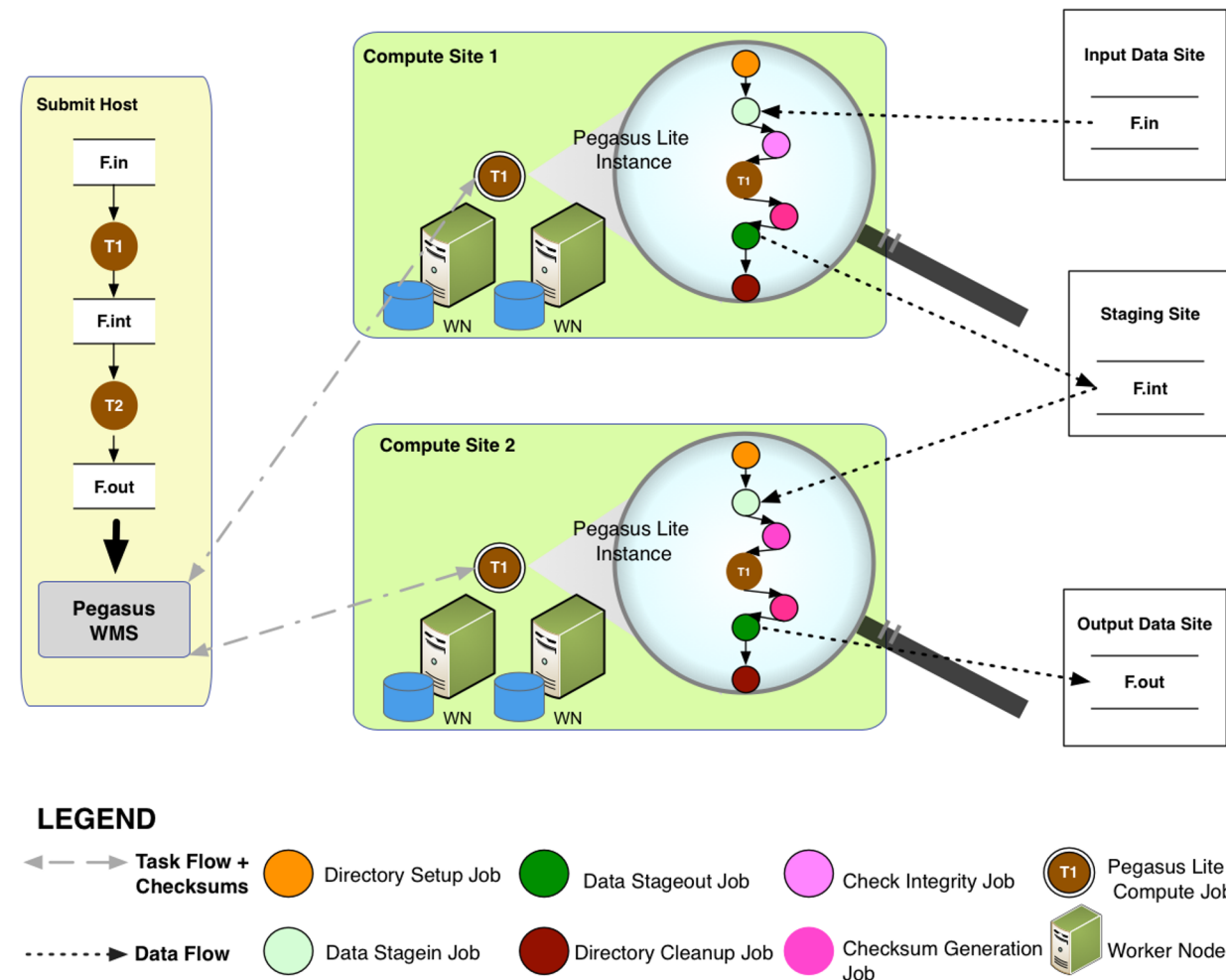
Plus there is the threat of intentional changes: malicious attackers, insider threats, etc.

User Perception: “Am I not already protected? I have heard about TCP checksums, encrypted transfers, checksum validation, RAID and erasure coding – is that not enough?”

Automatic Integrity Checking in Pegasus

Pegasus performs integrity checksums on input files right before a job starts on the remote node.

- For raw inputs, checksums specified in the input replica catalog along with file locations
- All intermediate and output files checksums are generated and tracked within the system.
- Support for sha256 checksums



Job failure is triggered if checksums fail

Pegasus: Containers Data Management

- Treat containers as input data dependency
 - Needs to be staged to compute node if not present
- Users can refer to container images as
 - Docker Hub or Singularity Library URL's
 - Docker Image exported as a TAR file and available at a server , just like any other input dataset.
- If an image is specified to be residing in a hub
 - The image is pulled down as a tar file as part of data stage-in jobs in the workflow
 - The exported tar file is then shipped with the workflow and made available to the jobs
 - Motivation: Avoid hitting Docker Hub/Singularity Library repeatedly for large workflows
- Symlink against a container image if available on shared filesystem
 - For e.g. CVMFS hosted images on Open Science Grid

Pegasus: Container Representation

Described in Transformation Catalog

- Maps logical transformations to physical executables on a particular system

container

Reference to the container to use.

Multiple transformation can refer to same container

type

Can be either docker or singularity or shifter

image

URL to image in a docker|singularity hub OR
to an existing docker image exported as a
tar file or singularity image

mount

Mount information to mount host directories
into container



Pegasus

```
- transformations
```

```
- namespace: "example"
  name: "keg"
  version: 1.0
```

```
site:
```

```
- name: "isi"
  arch: "x86"
  os "linux"
  pfn "/usr/bin/pegasus-keg"
  container "centos-pegasus"
```

```
# INSTALLED means pfn refers to path in the container.
```

```
# STAGEABLE means the executable can be staged into the container
type "INSTALLED"
```

```
- cont:
```

```
- name: "centos-pegasus"
```

```
# can be docker, singularity or shifter
```

```
type: "docker"
```

```
# URL to image in docker|singularity hub or shifter repo URL or
```

```
# URL to an existing image exported as a tar file or singularity image file
```

```
image: "docker:///centos:7"
```

```
# mount information to mount host directories into
```

```
# container format src-dir:dest-dir[:options]
```

```
mount:
```

```
- "/Volumes/Work/lfs1:/shared-data/:ro"
```

```
# environment to be set when the job is run in the container
```

```
# only env profiles are supported
```

```
profile:
```

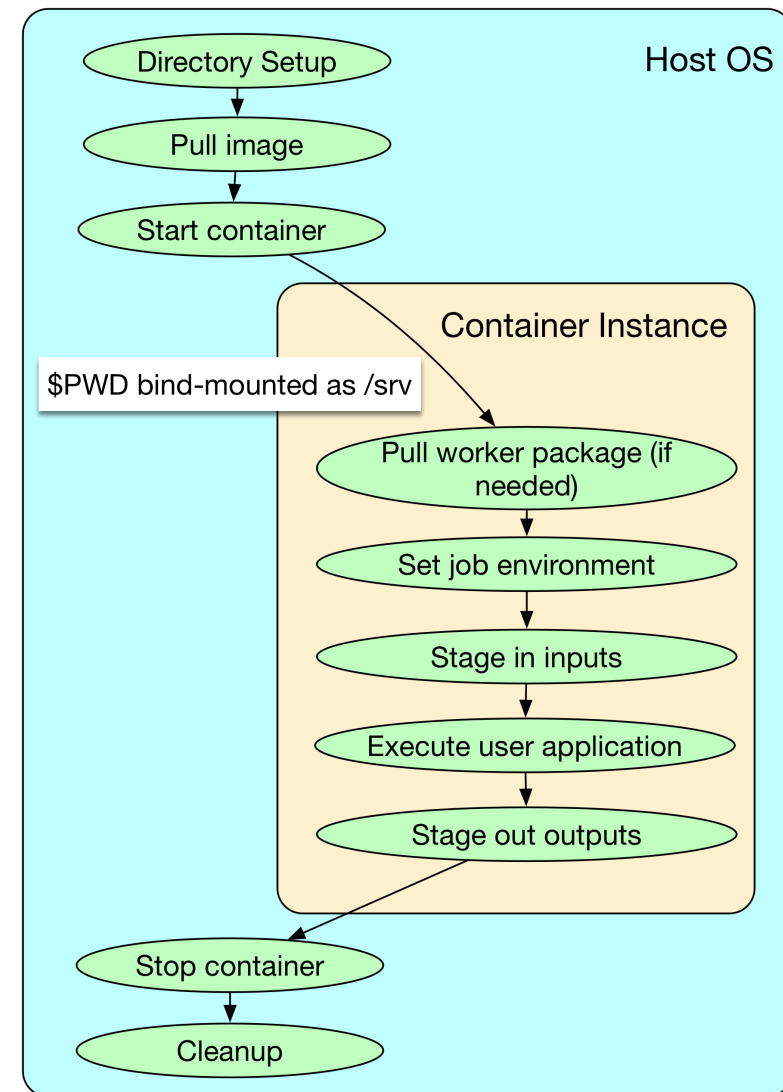
```
- env:
```

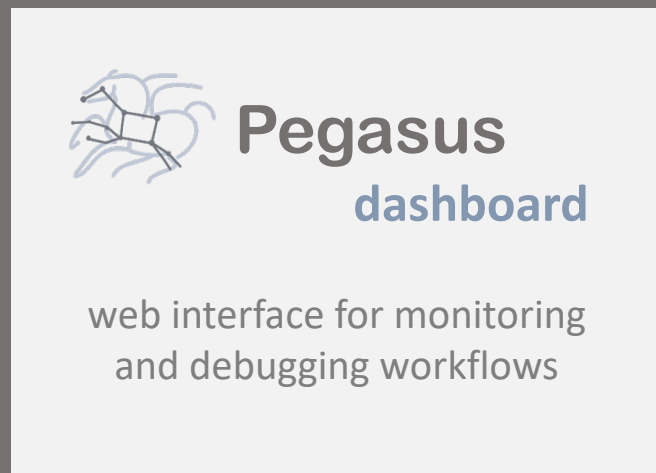
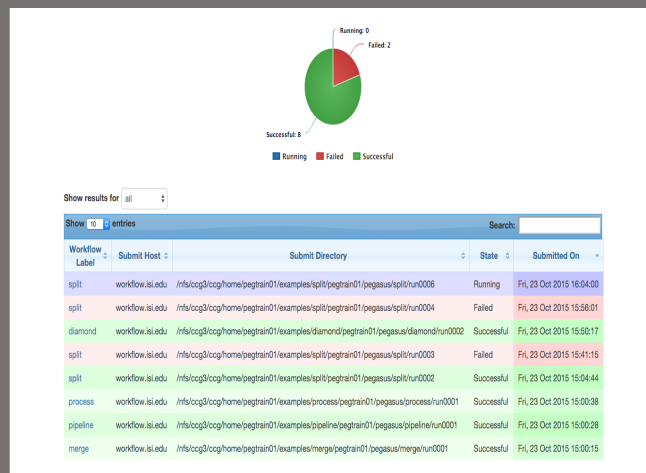
```
"JAVA_HOME" "/opt/java/1.6"
```



Pegasus: Container Execution Model

- Containerized jobs are launched via Pegasus Lite
 - Container image is put in the job directory along with input data.
 - Loads the container if required on the node (applicable for Docker)
 - Run a script in the container that sets up Pegasus in the container and job environment
 - Stage-in job input data
 - Launches user application
 - Ship out the output data generated by the application
 - Shut down the container (applicable for Docker)
 - Cleanup the job directory





Statistics

Workflow Wall Time	12 mins 23 secs
Workflow Cumulative Job Wall Time	9 mins 34 secs
Cumulative Job Walltime as seen from Submit Side	9 mins 35 secs
Workflow Cumulative Badput Time	9 mins 23 secs
Cumulative Job Badput Walltime as seen from Submit Side	9 mins 20 secs
Workflow Retries	1

Workflow Statistics

Type	Succeeded	Failed	Incomplete	Total	Retries	Total + Retries
Tasks	5	0	0	5	0	5
Jobs	16	0	0	16	2	18
Sub Workflows	0	0	0	0	0	0

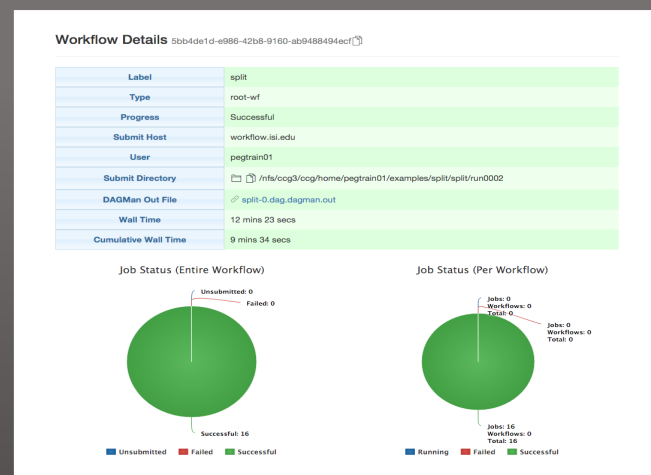
Entire Workflow

Type	Succeeded	Failed	Incomplete	Total	Retries	Total + Retries
Tasks	5	0	0	5	0	5
Jobs	16	0	0	16	2	18
Sub Workflows	0	0	0	0	0	0

Job Breakdown Statistics

Job Statistics

Real-time monitoring of workflow executions. It shows the status of the workflows and jobs, job characteristics, statistics and performance metrics. Provenance data is stored into a relational database.



Real-time Monitoring
Reporting
Debugging
Troubleshooting
RESTful API



Pegasus dashboard

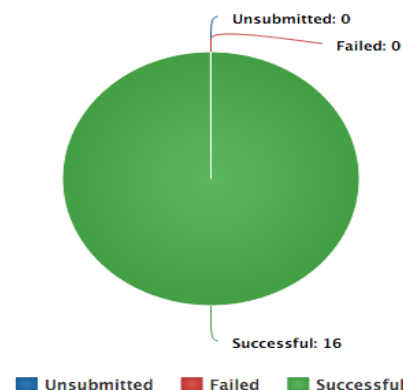
web interface for monitoring
and debugging workflows

Real-time monitoring of
workflow executions. It shows
the status of the workflows and
jobs, job characteristics, statistics
and performance metrics.
Provenance data is stored into a
relational database.

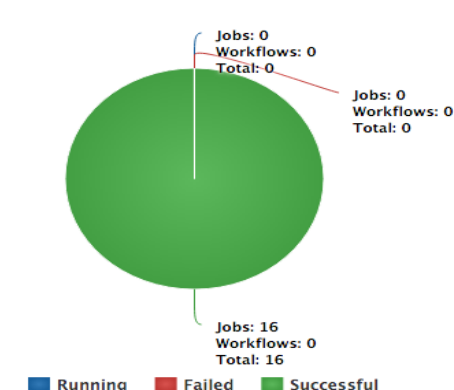
Workflow Details 5bb4de1d-e986-42b8-9160-ab9488494ecf

Label	split
Type	root-wf
Progress	Successful
Submit Host	workflow.isi.edu
User	pegtrain01
Submit Directory	/nfs/ccg3/ccg/home/pegtrain01/examples/split/split/run0002
DAGMan Out File	split-0.dag.dagman.out
Wall Time	12 mins 23 secs
Cumulative Wall Time	9 mins 34 secs

Job Status (Entire Workflow)



Job Status (Per Workflow)





Pegasus 5.0

Automate, recover, and debug scientific computations

- Reworked Python API to compose, submit and monitor workflows and configure catalogs
- New Catalog Formats
- Python 3 support
 - All Pegasus tools are Python 3 compliant.
 - 5.0 release will require Python 3 on workflow submit node
 - Python PIP packages for workflow composition and monitoring
- Zero configuration required to submit to local HTCondor pool.
- Data Management Improvements
 - New output replica catalog that registers outputs including file metadata such as size and checksums
 - Improved support for hierarchal workflows
- Revamped Documentation

```
#!/usr/bin/env python3
import logging
import sys

from Pegasus.api import *

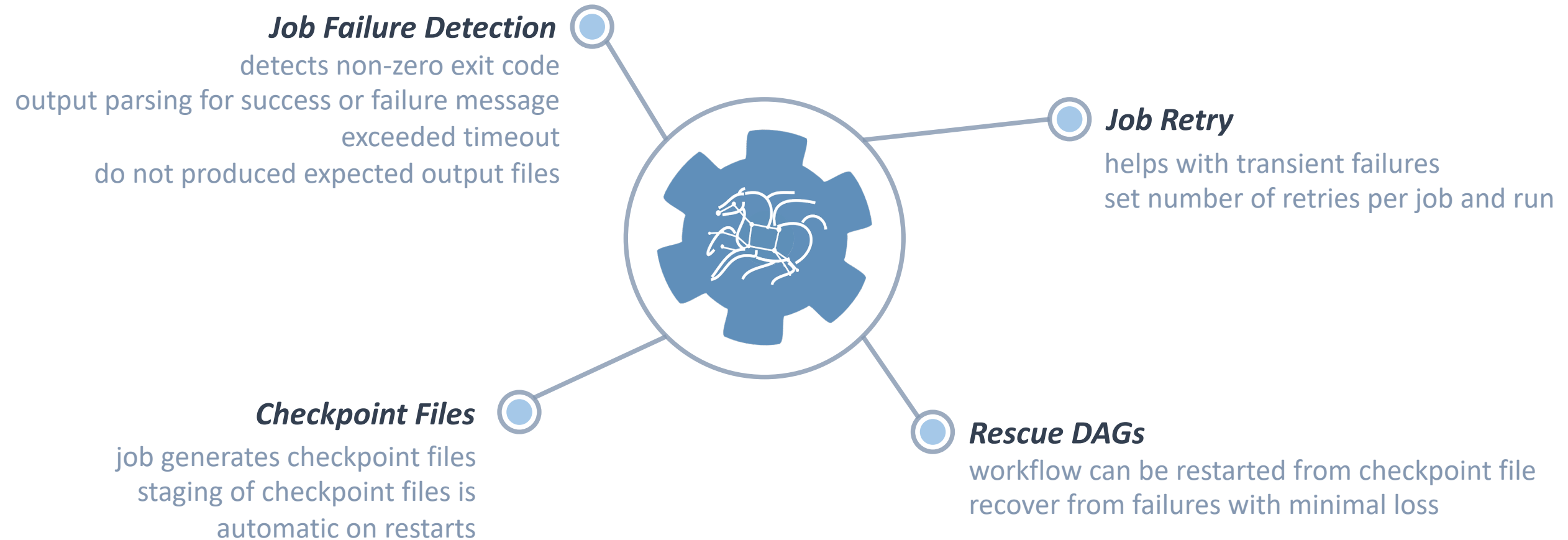
# logs to be sent to stdout
logging.basicConfig(level=logging.DEBUG, stream=sys.stdout)

# --- Transformations -----
echo = Transformation(
    "echo",
    pfn="/bin/echo",
    site="condorpool"
)

tc = TransformationCatalog()\
    .add_transformations(echo)

# --- Workflow -----
Workflow("hello-world", infer_dependencies=True)\
    .add_jobs(
        Job(echo)
        .add_args("Hello World")
        .set_stdout("hello.out")
    ).add_transformation_catalog(tc)\
    .plan(submit=True)\
    .wait()
```

And if a job fails?



Job Submissions

Local

Submit Machine

Personal HTCondor

Local Campus Cluster accessible via Submit Machine **

HTCondor via BLAHP

**** Both Glite and BOSCO build on HTCondor BLAHP**

Currently supported schedulers:

SLURM SGE PBS MOAB

Remote

BOSCO + SSH**

Each node in executable workflow submitted via SSH connection to remote cluster

BOSCO based Glideins**

SSH based submission of glideins

PyGlidein

IceCube glidein service

OSG using glideinWMS

Infrastructure provisioned glideins

CREAMCE

Uses CondorG

Globus GRAM

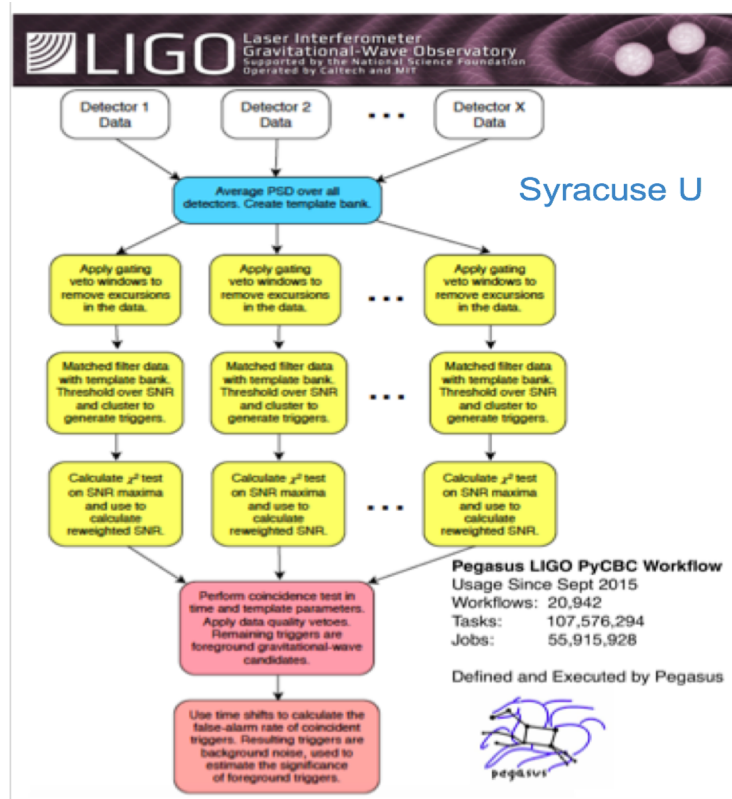
Uses CondorG

Pegasus Workflow Management System, Production Use

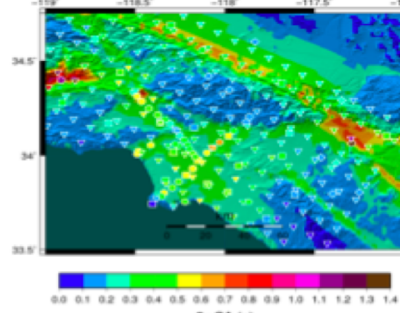


Last 12 months: Pegasus users ran **240K** workflows, **145M** jobs

Majority of these include data transfers, using LAN, the Internet, local and remote storage



Southern California Earthquake Center, USC



Nek2 Kinase

PDB ID: 2DKA

Chain: A

Binding Cavity

0.1 0.2

Rank-Ordered Compounds

Binding Mode

Predicted Additional Targets

Other Proteins in Crystal Structure (Chain)

No other chains found for this PDB structure

Other Proteins in Crystal Structure (Chain)

No other chains found for this PDB structure

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Other Proteins in Crystal Structure (Chain)

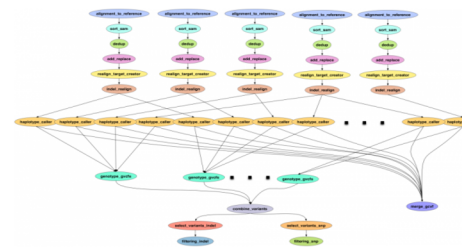
No other chains found for this PDB structure

Other Proteins in Crystal Structure (Chain)

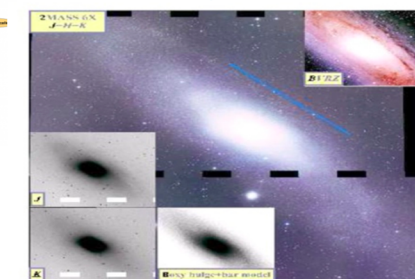
No other chains found for this PDB structure

Other Proteins in Crystal Structure (Chain)

No other chains found for this PDB structure



Bioinformatics: Protein interactions, IU



Montage, Caltech

Bioinformatics: SoyKB, University of Arizona

Questions?



Pegasus

est. 2001

Automate, recover, and debug scientific computations.

Get Started

Pegasus Website

<https://pegasus.isi.edu>

Users Mailing List

pegasus-users@isi.edu

Support

pegasus-support@isi.edu

Pegasus Online Office Hours

<https://pegasus.isi.edu/blog/online-pegasus-office-hours/>

Bi-monthly basis on second Friday of the month, where we address user questions and also apprise the community of new developments

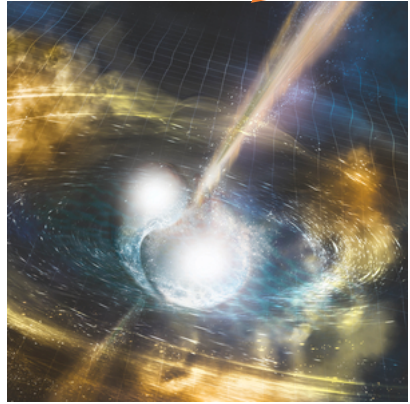
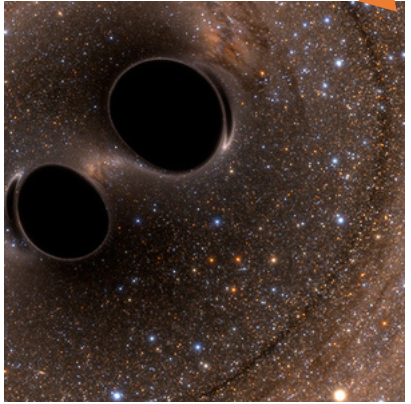
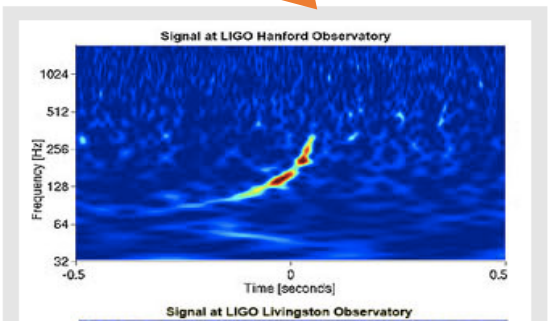
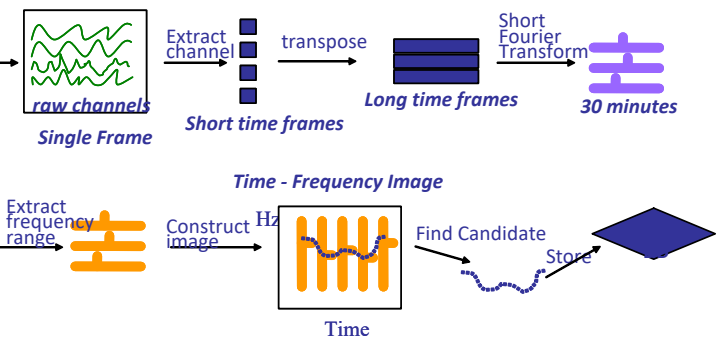
Extra Slides: User Stories

Pegasus: Grounding Research and Development

Nobel Prize



Working with LIGO



**First Pegasus
prototype for
LIGO Pulsar
Searches**

Blind injection detection

**First detection of
black hole collision**

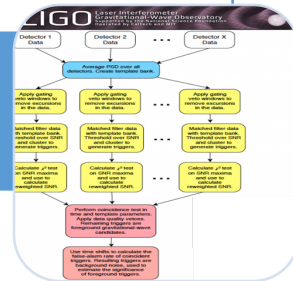
**Multi-messenger
neutron star
merger
observation**

Complexity of LIGO Workflows

First GW detection: $\sim 21\text{K}$ Pegasus workflows, $\sim 107\text{M}$ tasks

Analysis measures the statistical significance of collected data

Science Workflow



Efficient,
scalable, and
robust execution
of tasks and
data access

Automation



LIGO, Open
Science Grid,
XSEDE, Blue
Waters

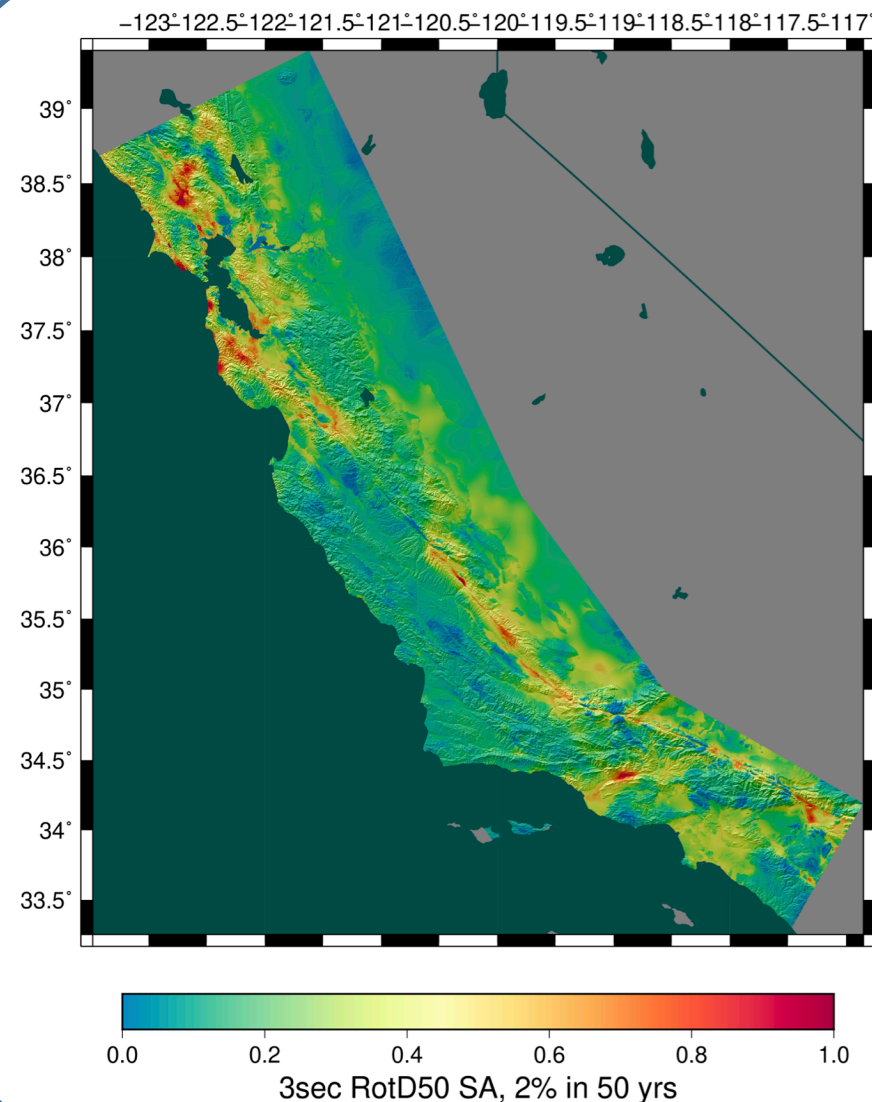
Distributed Power



2015/16

Supporting Heterogeneous Workflows

**SCEC's
CyberShake:
What will the
peak
earthquake
motion be
over the next
50 years?**



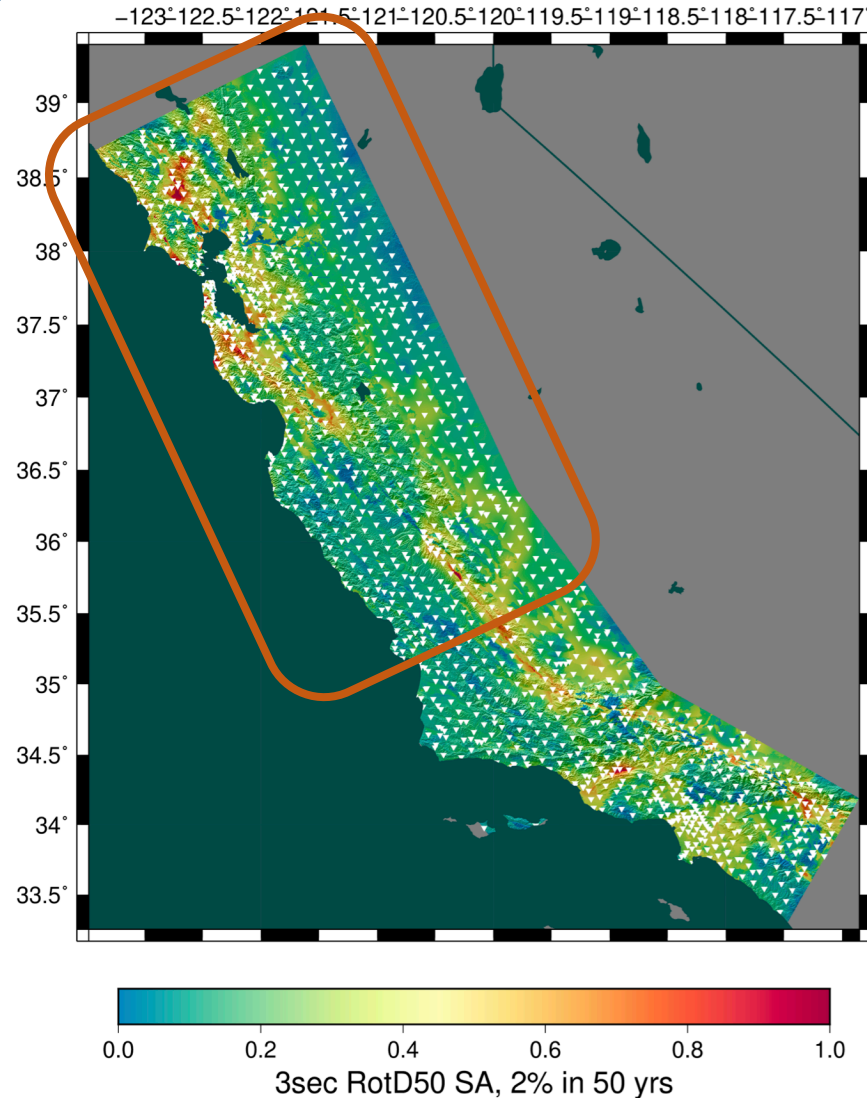
Useful information for:

- Building engineers
- Disaster planners
- Insurance agencies

Slide credit: Southern California
Earthquake Center

Supporting Heterogeneous Workflows

2018-2019 Mapping Northern California



- 120 million core-hours
- 39,285 jobs
- 1.2 PB of data managed
- 157 TB of data automatically transferred
- 14.4 TB of output data archived

- NCSA *Blue Waters*
- OLCF *Titan*

Total map:
170 million core hours
> 19,407 core years

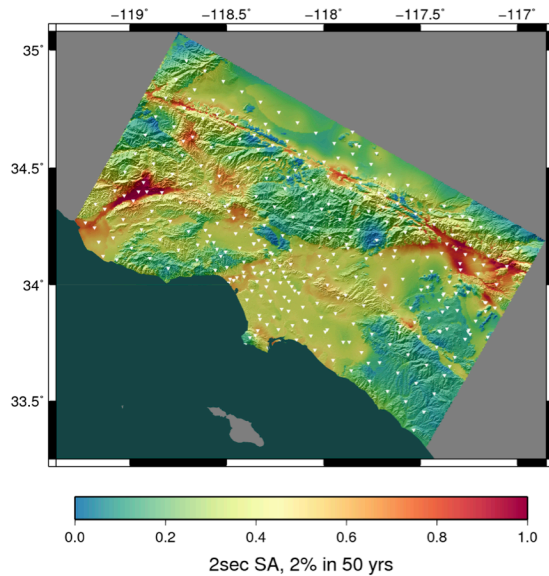
Mix Workloads on Heterogeneous/ Changing CI

Since 2007: 215 million core-hours (24,543 years)

9 different supercomputers

Pegasus Optimizations:

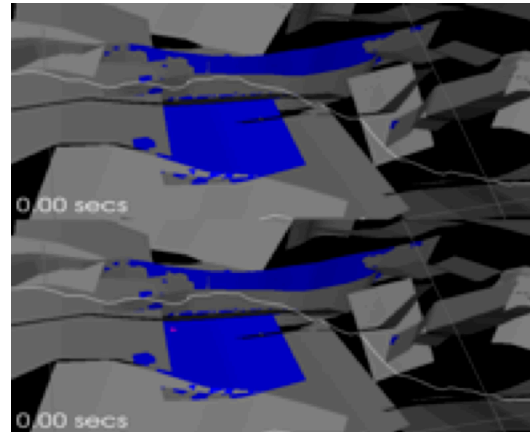
- Task clustering
- MPI-based workflow engine



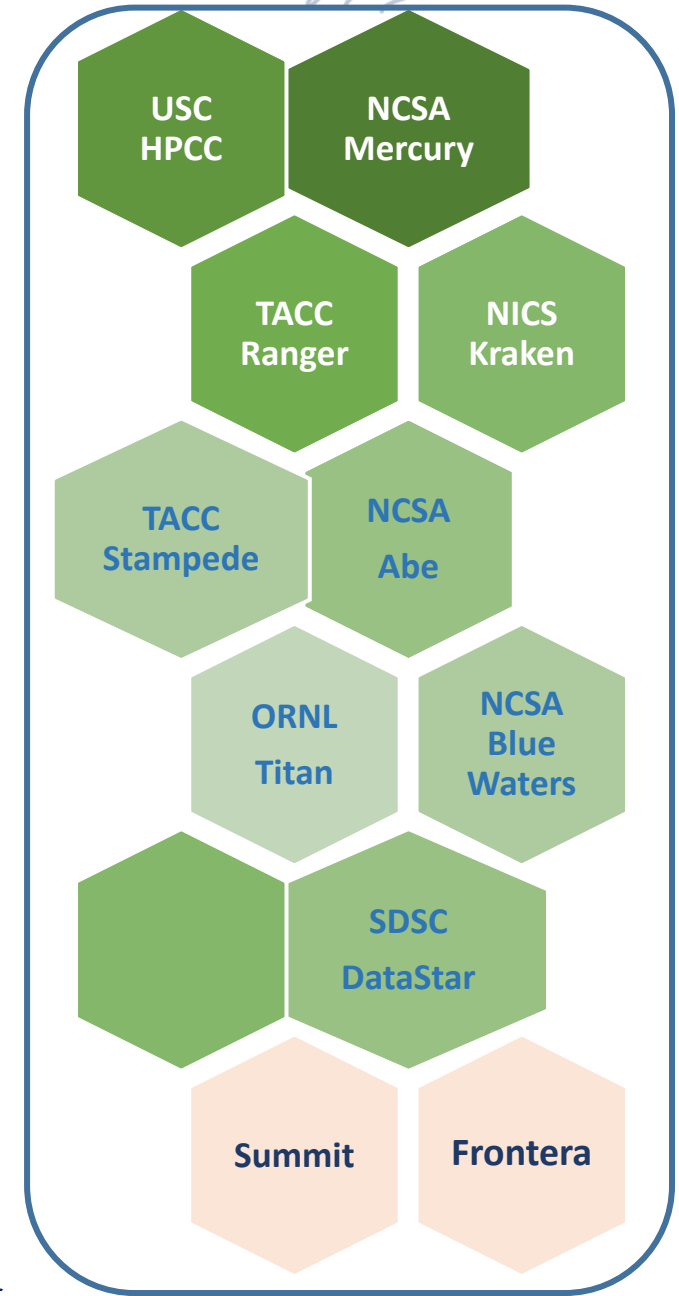
2010: World's first physics-based probabilistic seismic hazard map

Application Optimizations:

- Workflow restructuring
- MPI/code tuning
- Porting to GPUs

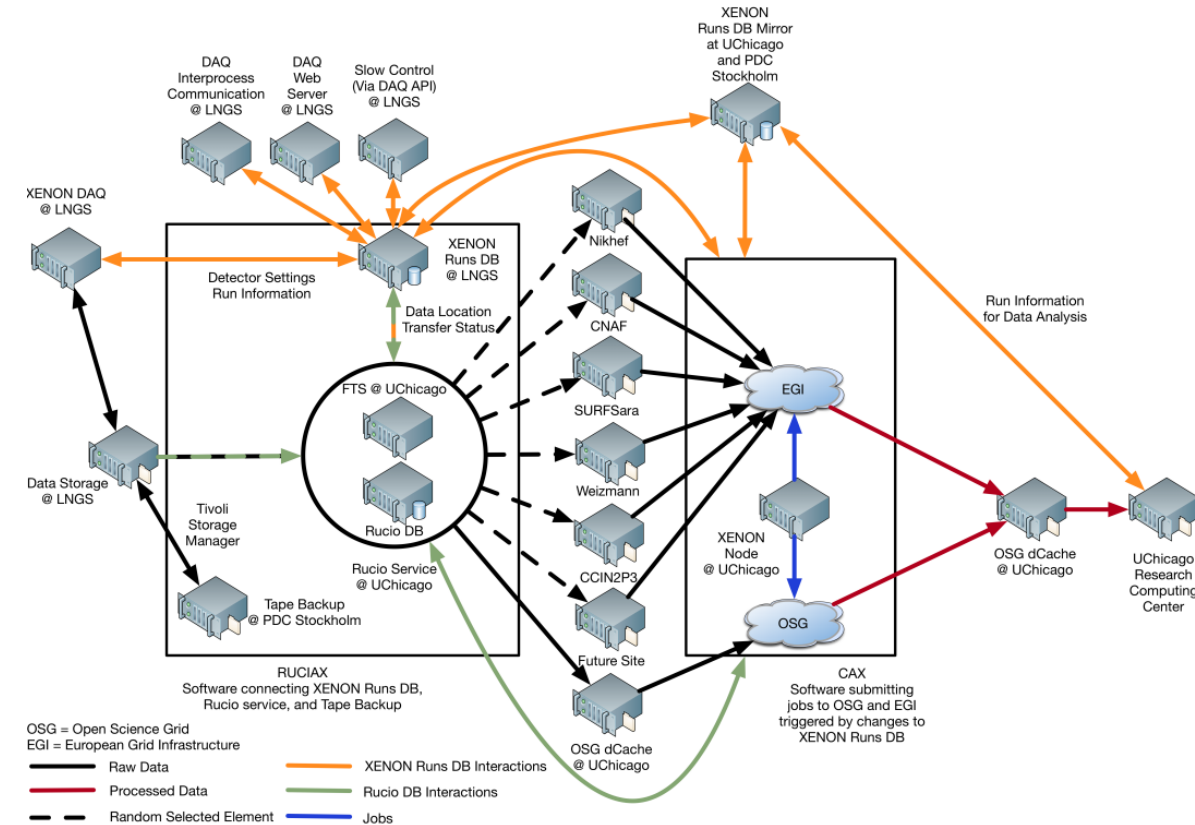


2018: Incorporating earthquake simulator with a 1 million-year catalog of California seismicity



XENONnT - Dark Matter Search

Detector at Laboratori Nazionali del Gran Sasso (LNGS) in Italy. Data is distributed world-wide with Rucio. Workflows execute across Open Science Grid (OSG) and European Grid Infrastructure (EGI)



Type	Succeeded	Failed	Incomplete	Total	Retries	Total+Retries
Tasks	4000	0	0	4000	267	4267
Jobs	4484	0	0	4484	267	4751
Sub-Workflows	0	0	0	0	0	0

Workflow wall time : 5 hrs, 2 mins
 Cumulative job wall time : 136 days, 9 hrs
 Cumulative job wall time as seen from submit side : 141 days, 16 hrs
 Cumulative job badput wall time : 1 day, 2 hrs
 Cumulative job badput wall time as seen from submit side : 4 days, 20 hrs

Main processing pipeline for XENONnT

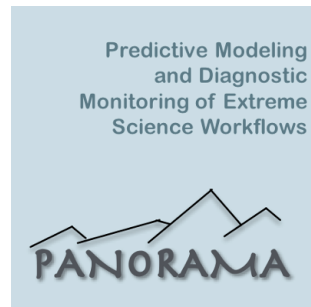
Impact on DOE Science

Enabled cutting-edge domain science (e.g., drug delivery) through collaboration with scientists at the DoE **Spallation Neutron Source (SNS)** facility

A Pegasus workflow was developed that confirmed that **nanodiamonds** can enhance the dynamics of tRNA

It compared SNS neutron scattering data with MD simulations by calculating the epsilon that best matches experimental data

Ran on a Cray XE6 at NERSC using 400,000 CPU hours, and generated 3TB of data.

The screenshot shows the Oak Ridge National Laboratory website. The header includes the ORNL logo and navigation links like "Visit ORNL", "News", "Events", "Careers", "Find People", "Retirees & Staff", and "Index". A search bar is on the right. The main content area features a news article titled "Diamonds that deliver" with a sub-headline "Neutrons, simulation analysis of tRNA-nanodiamond combo could transform drug delivery design principles". To the right of the article are "Related Topics" links for "Advanced Materials" and "Neutron Science". Below the text is a large 3D visualization of a nanodiamond sphere with water molecules (small red and white spheres) and tRNA molecules (colored spheres) attached to its surface.

OAK RIDGE
National Laboratory

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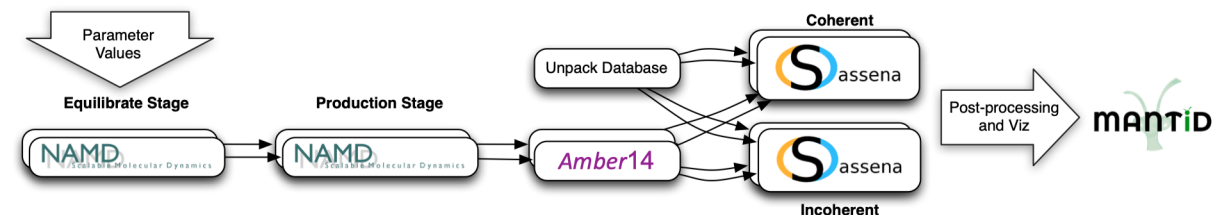
Home | News | Diamonds that deliver

Diamonds that deliver

Neutrons, simulation analysis of tRNA-nanodiamond combo could transform drug delivery design principles

Related Topics:
[Advanced Materials](#)
[Neutron Science](#)

Water is seen as small red and white molecules on large nanodiamond spheres. The colored tRNA can be seen on the nanodiamond surface.
(Image Credit: Michael Mattheson, OLCF, ORNL)



An automated analysis workflow for optimization of force-field parameters using neutron scattering data. V. E. Lynch, J. M. Borreguero, D. Bhowmik, P. Ganesh, B. G. Sumpter, T. E. Proffen, M. Goswami, Journal of Computational Physics, July 2017.