

Pegasus Workflows on XSEDE

Mats Rynge <rynge@isi.edu>

USC Information Sciences Institute



Information Sciences Institute

XSEDE ECSS Workflow Community Applications Team

https://www.xsede.org/web/guest/ecss-workflows

The XSEDE Workflow Community Applications Team's charter is to assist researchers to use scientific workflow technologies on XSEDE to solve challenging scientific problems involving parameter sweeps, multiple applications combined in dependency chains, tightly coupled applications, and similar execution patterns that require multiple applications and multiple XSEDE resources. The workflow team accomplishes its mission through the use of third party workflow software in collaboration with the workflow developers, service providers and XSEDE Extended Collaborative Support Services.

Community mailing list: workflows@xsede.org

 To subscribe, email <u>majordomo@xsede.org</u> with "subscribe workflows" in the body of the message

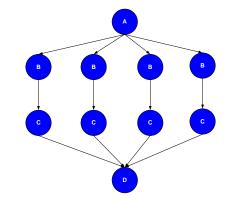
SFI

More Information:

Marlon Pierce (<u>mpierce@iu.edu</u>) Mats Rynge (<u>rynge@isi.edu</u>) Suresh Marru (<u>smarru@iu.edu</u>)

Pegasus Workflow Management System

- NSF funded project and developed since 2001 as a collaboration between USC Information Sciences Institute and the HTCondor Team at UW Madison
- Builds on top of HTCondor DAGMan.
- Abstract Workflows Pegasus input workflow description
 - Workflow "high-level language"
 - Only identifies the computation, devoid of resource descriptions, devoid of data locations
- Pegasus is a workflow "compiler" (plan/map)
 - Target is DAGMan DAGs and HTCondor submit files
 - Transforms the workflow for performance and reliability
 - Automatically locates physical locations for both workflow components and data
 - Collects runtime provenance



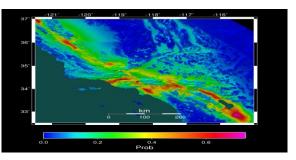


Southern California Earthquake Center's "CyberShake" workflow

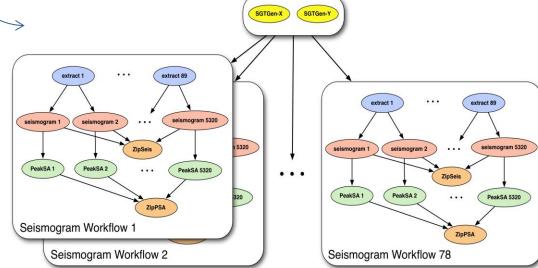


Description

- Builders ask seismologists: "What will the peak ground motion be at my new building in the next 50 years?"
- Seismologists answer this question using Probabilistic Seismic Hazard Analysis (PSHA)
- Hierarchal workflow with mix of large MPI jobs (top) and large number of serial tasks (post processing)
- The size of the workflow depends on the number of sites and sampling frequency
- For each of the 600 sites in the input map, generate a hazard curve
- Each site has a sub-workflow with 820,000 tasks



School of Engineering



- SGT (Strain Green Tensor MPI jobs) output: 15.6 TB (40 * 400 GB)
- Final outputs: 500 million files (820,000/site x 600 sites)
 5.8 TB (600 * 10 GB)

Probability of exceeding 0.1g in 50 yrs

4

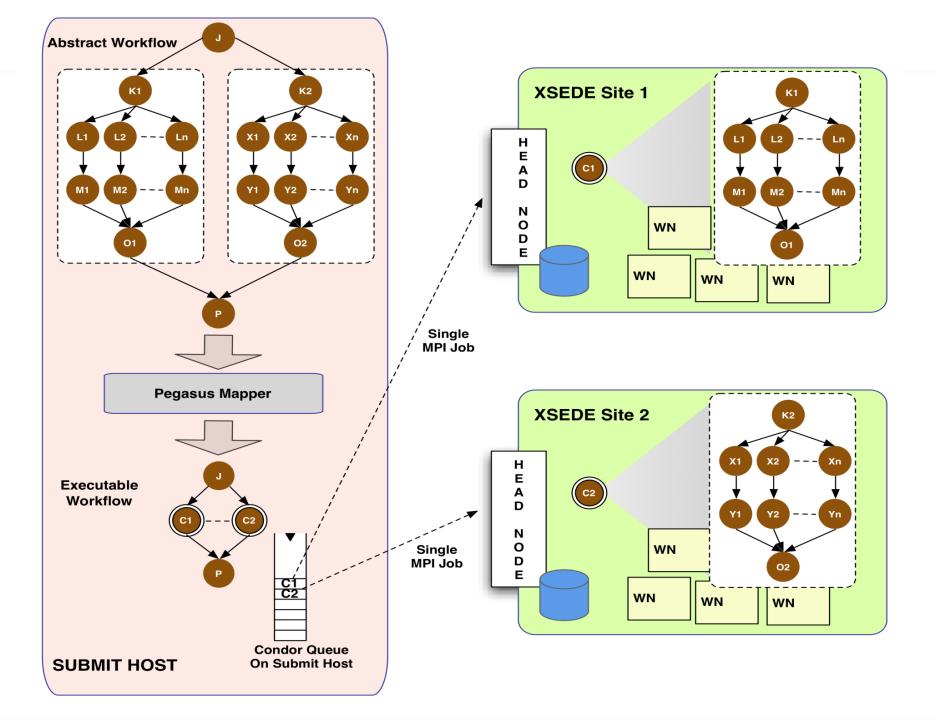


pegasus-mpi-cluster

- Master/worker paradigm
- Master manages the subgraph tasks, handing out work to the workers
- Efficient scheduling / handling of input/outputs
- Subgraph described in a DAG-similar format
- Failure management / rescue DAG

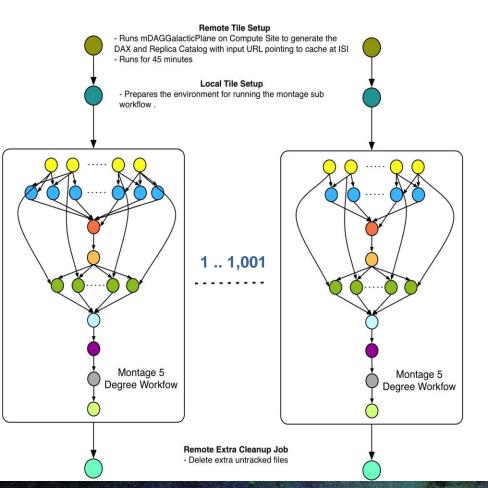






GALACTIC PLANE WORKFLOW

Montage Galactic Plane Workflow



- Galactic Plane for generating mosaics from the NASA Telescope Missions like Spitzer etc.
- Used to generate tiles 360 x 40 degrees around the galactic equator
- A tile 5 x 5 with 1 overlap with neighbors
- Output datasets to be potentially used in NASA Sky and Google Sky
- Each per band workflow
 - 1.6 million input files
 - 10.5 million tasks
 - Consumes 34,000 CPU hours
 - Generates **1,001 tiles** in FITS format

× 16

- Ran on XSEDE, moved to Amazon EC2
- Run workflows corresponding to each of the 16 bands (wavelengths)
- Total Number of Data Files 18 million
- Potential Size of Data Output 45 TB

Acknowledgements: Bruce Berriman, John Good at IPAC - http://montage.ipac.caltech.edu/gallery.html





SoyKB – Soybean Knowledge Base

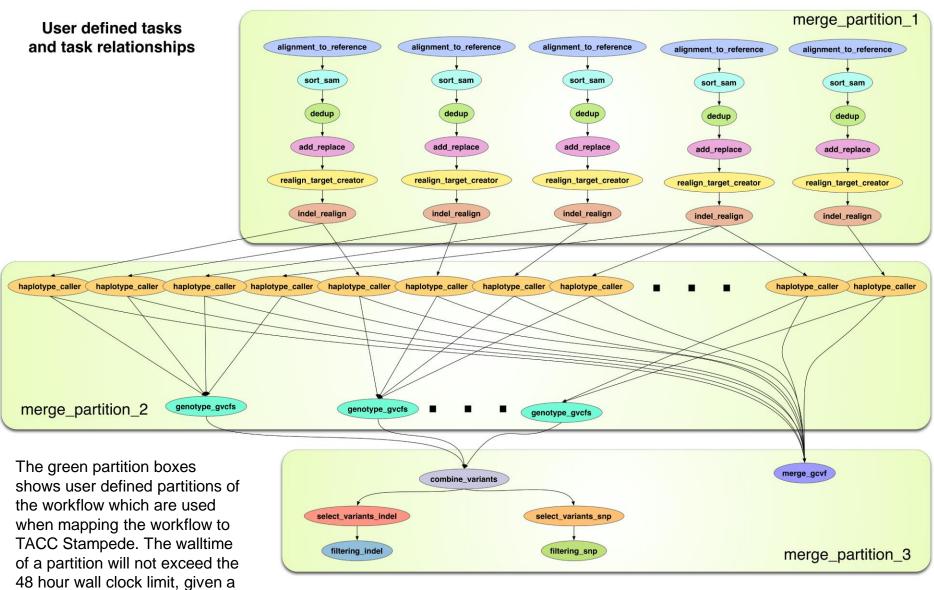
- Bioinformatics analysis of 1000+ resequenced soybean germplasm lines selected for major traits including oil, protein, soybean cyst nematode resistance (SCN), abiotic stress resistance (drought, heat and salt) and root system architecture.
- Sequencing pipeline using standard tools: Picard, BWA, GATK

Environment

- iPlant Data Store for inputs and outputs
- iPlant Atmosphere for VMs for workflow management
- TACC Stampede as execution environment





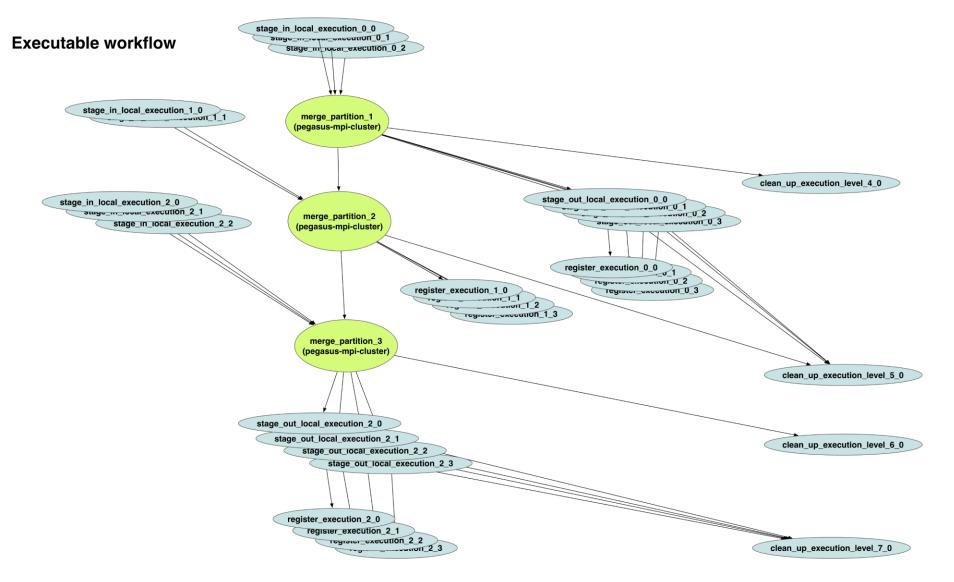


of a partition will not exceed 48 hour wall clock limit, given certain number of compute nodes.

USCViterbi

School of Engineering





The green jobs are MPI container jobs of the partitions shown in the earlier slide. These are run on the remote supercomputer. The blue auxiliary tasks are run on the workflow submit host, and handle things like data staging and cleanup.



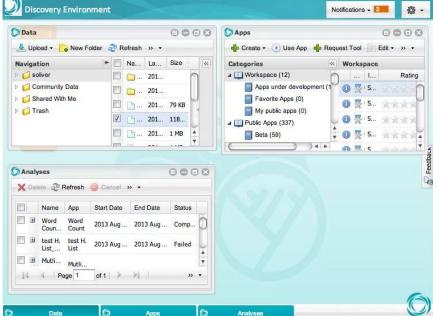


SoyKB - Highlights

- Workflow can be mapped to a distributed or a supercomputer execution environment
 - Distributed environment is used for testing the pipeline
 - The supercomputer environment is used for production runs, and tasks are packed up in MPI container jobs (pegasus-mpi-cluster)

11

- Input data is fetched from iPlant Data Store, or if already replicated, from the TACC iPlant Data Store node for close to computation access
- Outputs are automatically put back into the Data Store for easy access and further analysis in the iPlant Discovery Environment



iPlant

Collaborative



http://pegasus.isi.edu

Tutorial and documentation: http://pegasus.isi.edu/wms/docs/latest/

Mats Rynge rynge@isi.edu



