

First Brazil-France Workshop On High Performance Computing and Scientific Data Management Driven by Highly Demanding Applications

COPPE UFRJ
Instituto Alberto Luis Coimbra de Pós-Graduação e Pesquisa de Engenharia

EXPLORING PROVENANCE IN HIGH PERFORMANCE SCIENTIFIC COMPUTING

Marta Mattoso
COPPE/Federal University of Rio de Janeiro



15/09/12

Group 2 at COPPE/UFRJ **COPPE UFRJ**

- Marta Mattoso (P.I. Distributed Database Group)
- Post-Docs:
 - ✓ Kary Ocana, D.Sc 2010 - Bioinformatics
 - ✓ Eduardo Ogasawara, DSc 2011 - Workflow engine
 - ✓ Daniel Oliveira, DSc 2012 - Cloud Workflows
- DSc Students:
 - ✓ Jonas Dias
 - ✓ Flavio Costa

15/09/12

Group 2 at COPPE/UFRJ **COPPE UFRJ**

- **On going Collaborations** (now also in HOSCAR)
- INRIA:
 - ✓ Patrick Valduriez
 - CAPES, CNPq, FAPERJ and INRIA Equipes Associées
- LNCC:
 - ✓ Fabio Porto
 - ✓ Luiz Gadelha Jr

informatics mathematics
inria

National Laboratory Scientific Computing

15/09/12

Group 2 at COPPE/UFRJ **COPPE UFRJ**

- **On going Collaborations** (now also in HOSCAR – GROUP 1)
- COPPE- Numerical Methods:
 - ✓ Alvaro Coutinho
 - Renato Elias
 - José Camata
 - ✓ Fernando Rochinha
 - Gabriel
 - ✓ Luiz Landau
 - Josias

NACAD
High Performance Computing Center

15/09/12

Group 2 at COPPE/UFRJ

Other International Collaborations (US)

- U.Chicago/ Argonne:
 - ✓ Ian Foster
 - ✓ Mike Wilde
- New York - Polytechnic Institute of NYU:
 - ✓ Juliana Freire

15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting

HOSCAR - Objectives

- “...taking full benefits of the processing capabilities of future high performance massively **parallel architectures** in the framework of very **large-scale datasets** and **numerical simulations**...”
- **Group 3** (parallel architectures): basis for everyone
- Intergroup collaboration **Group 1** (numerical simulations) and **Group 2** (large-scale datasets)

15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting

HOSCAR - Objectives

- “...taking full benefits of the processing capabilities of future high performance massively **parallel architectures** in the framework of very **large-scale datasets** and **numerical simulations**...”
- Intergroup collaboration **Group 1** (numerical simulations) and **Group 2** (large-scale datasets)
- Why, how to collaborate ?
 - ✓ Some of our (large datasets) previous experiences collaborating with (simulations) COPPE& Petrobras
 - ✓ Manage scientific resources (files, programs, images)

15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting


1st Workshop on High-Performance Computing Meets Databases


SC11
Seattle, WA

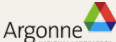
Exploring Provenance in High Performance Scientific Computing

Luiz Gadelha,
Marta Mattoso
COPPE, Federal Univ Rio de Janeiro,
Brazil

Michael Wilde,
Ian Foster
CI, U. Chicago / ANL, USA

 **COPPE**
Instituto Alberto Luis Coimbra de Pós-Graduação e Pesquisa de Engenharia UFRJ

 **THE UNIVERSITY OF CHICAGO**

 **Argonne**
NATIONAL LABORATORY

1st International Workshop on Scalable Workflow Enactment Engines and Technologies **SWEET'12**
Colocated with SIGMOD'12

EVALUATING PARAMETER SWEEP WORKFLOWS IN HIGH PERFORMANCE COMPUTING

Fernando Chirigati*, Vitor Silva, Eduardo Ogasawara, Daniel de Oliveira, Jonas Dias, Fábio Porto, Patrick Valduriez and Marta Mattoso

COPPE / Federal University of Rio de Janeiro, Rio de Janeiro, Brazil
LNCC, Rio de Janeiro, Brazil
INRIA & LIRMM, Montpellier, France

* Currently at Polytechnic Institute of NYU


CCSA 2012 @ CCGrid
2nd Cloud Computing and Scientific Applications (CCSA)

Cloud-based Phylogenomic Inference of Evolutionary Relationships: A Performance Study*

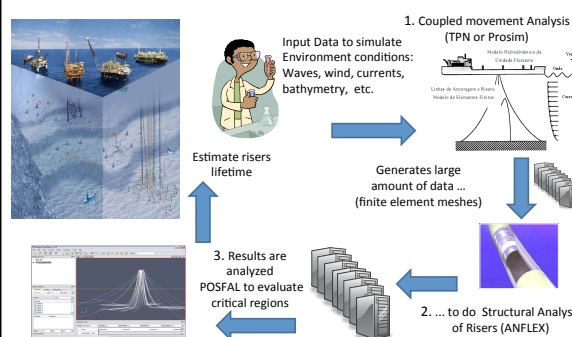
Daniel de Oliveira, Kary Ocaña, Eduardo Ogasawara, Jonas Dias, João Gonçalves and Marta Mattoso

Federal University of Rio de Janeiro
Rio de Janeiro, Brazil

* Best paper award



Risers' Fatigue Analysis in Ultra-Deep Waters



1. Coupled movement Analysis (TPN or Prosim)
Input Data to simulate Environment conditions: Waves, wind, currents, bathymetry, etc.
Generates large amount of data ... (finite element meshes)

2. ... to do Structural Analysis of Risers (ANFLEX)

3. Results are analyzed POSFAL to evaluate critical regions
Estimate risers lifetime

Marta Mattoso – HOSCAR, Brazil Meeting

HPC problems potentially amenable to database techniques

- as HPC computations become longer-running, there is a need to make them more interactive.
- Ad hoc science questions such as “Which regions generated by experiment X and parameter Y exhibit critic above some threshold?”
- Ad hoc monitoring questions such as “Which regions should I visualize?”

- Modeled, expressed, and optimized as queries
- Using Provenance / Scientific Workflow approach

COPPE UFRJ

Marta Mattoso – HOSCAR, Brazil Meeting

Scientific Computing with Workflows

COPPE UFRJ
Centro de Processamento de Dados
 Instituto de Física e Química

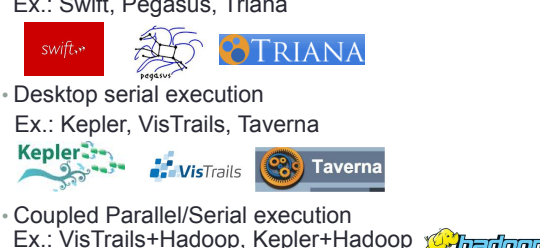
- Scientific experiments are often defined by a composition of many computational tasks linked through data flow
- These are also called *scientific workflows*
- Data sets manipulated can be large in size and quantity
- Analyzing the outcome of these large-scale computations can be difficult
- Provenance support is the key

15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting

Scientific Workflow Management Systems

COPPE UFRJ
Centro de Processamento de Dados
 Instituto de Física e Química

- Native parallel execution
 Ex.: Swift, Pegasus, Triana
- Desktop serial execution
 Ex.: Kepler, VisTrails, Taverna
- Coupled Parallel/Serial execution
 Ex.: VisTrails+Hadoop, Kepler+Hadoop



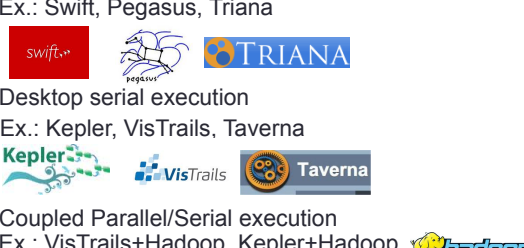
15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting

Scientific Workflow Management Systems

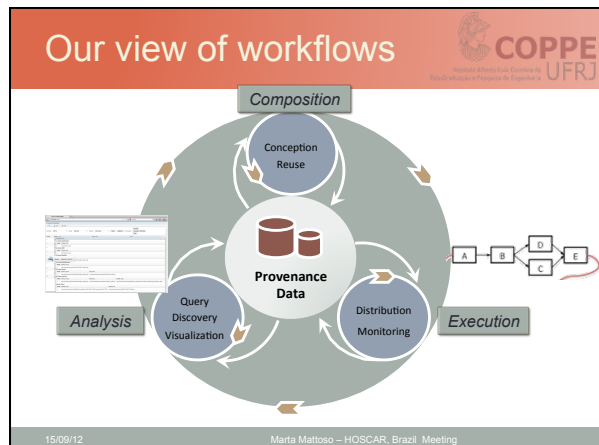
COPPE UFRJ
Centro de Processamento de Dados
 Instituto de Física e Química

- Native parallel execution
 Ex.: Swift, Pegasus, Triana
- Desktop serial execution
 Ex.: Kepler, VisTrails, Taverna
- Coupled Parallel/Serial execution
 Ex.: VisTrails+Hadoop, Kepler+Hadoop

Provenance is only available after the execution



15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting



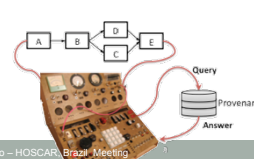
Data explorations with Wf

- The execution of variations upon the same workflow repeatedly
 - ✓ Parameter Sweeps
 - ✓ Fine-tuning (tolerance, solver options, ...)
 - ✓ Iterative methods
 - Stop criteria defined by the scientist evaluation
- Interfere in the dataflow
 - ✓ Input parameters
 - ✓ Workflow configurations
- Dynamic nature of simulations/ large data

15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting

Objectives

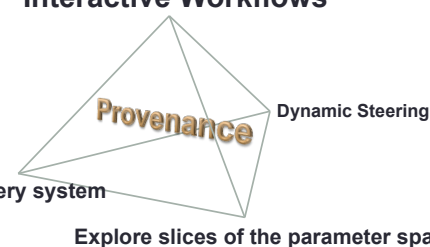
- Interactive Workflows
- Adaption and user-steering on workflow execution
 - ✓ Parameter space slicing
 - To compare results from different explored slices
 - ✓ Workflow adjustments
 - Human intervention
 - Dynamic steering
- Improvements
 - ✓ Quality of results
 - ✓ Execution efficiency
 - ✓ Scientific Data analysis



15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting

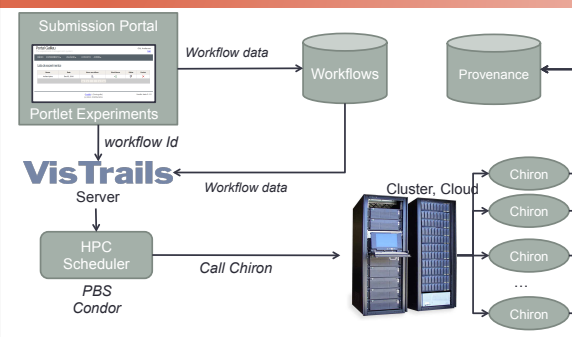
Using provenance

Interactive Workflows

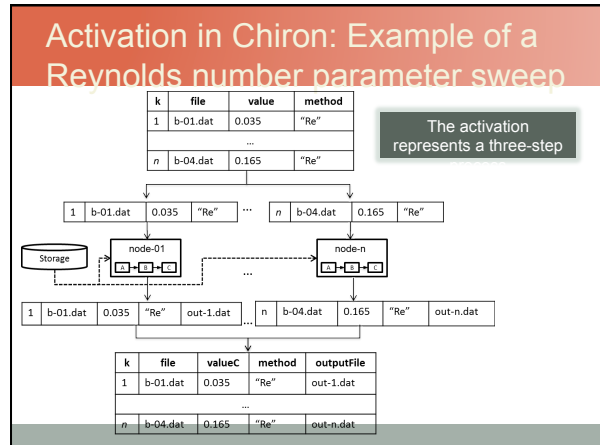
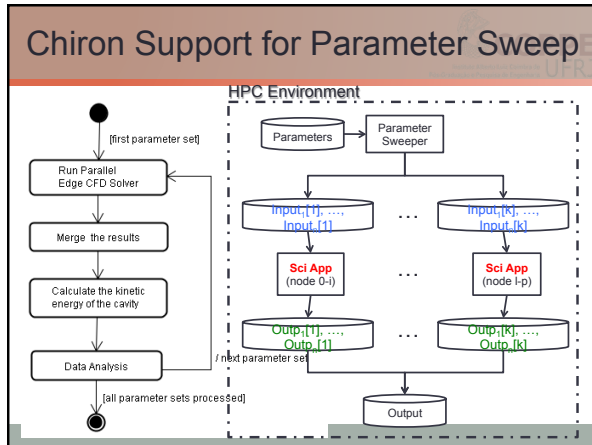


15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting

Chiron Workflow Execution



15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting



Provenance querying

COPPE UFRJ

```

SELECT t.workspace
FROM Task t, ParameterValue pv, Parameter p
WHERE p.ParameterID = pv.ParameterID
AND pv.TaskID = t.TaskID
AND p.ParameterName = 'Reynolds_Number'
AND pv.NumericValue > 0.090
AND pv.NumericValue < 0.110;
    
```

Provenance DB:

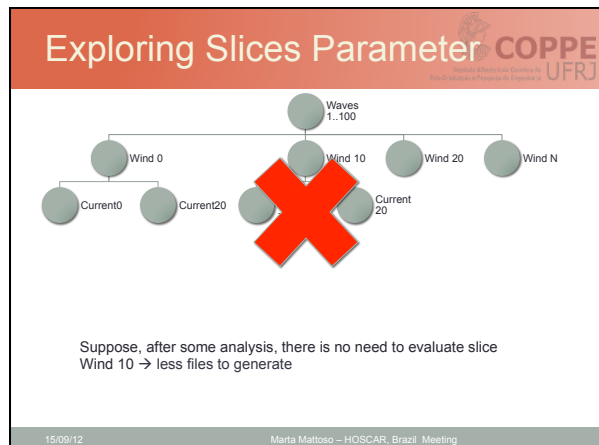
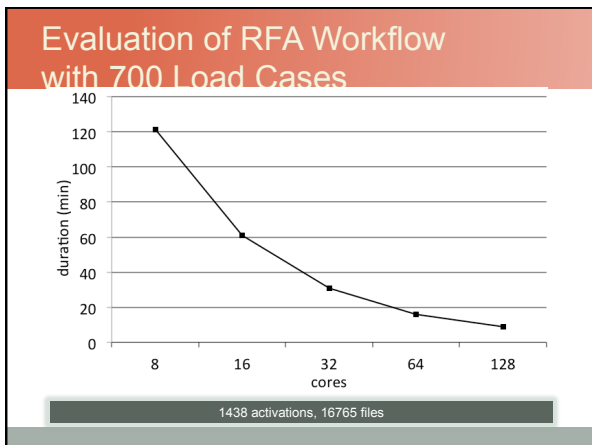
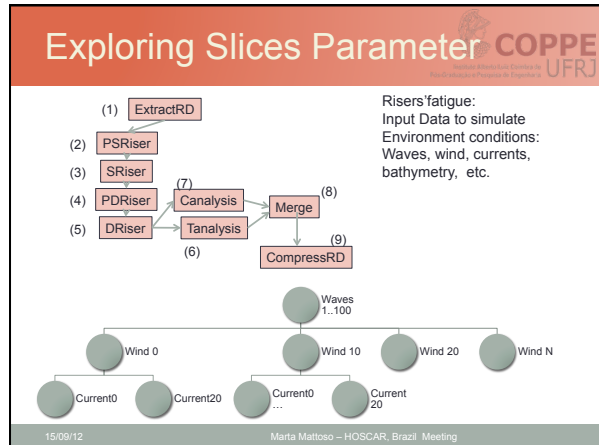
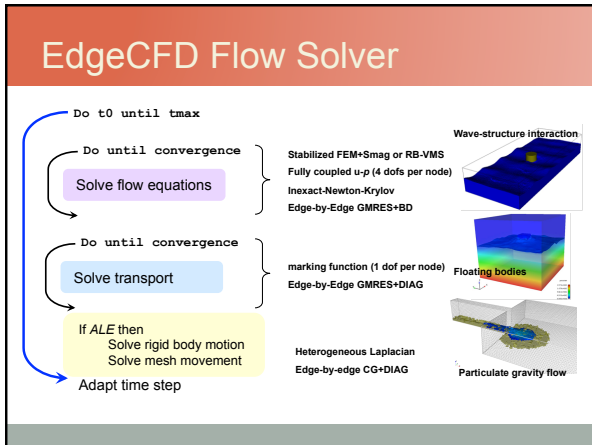
- files consumed and produced, standard output and standard error,
- execution time and the computer node that executed the program,
- used in real time to verify if any deterministic solver failed

Large Eddy Simulation Workflow Parallel Execution in Chiron

```

./edgesolver cav1
./out_merge cav1
./stats cav1
./extract cav1 .3956..1724,..4709
./rm -rf *.out
    
```

- Provenance gathers data to be used to evaluate statistical moments
- We can track the error between MC and SGC and determine an interpolation level to satisfy a minimum error
- Provenance capabilities in Chiron can automatically increase the interpolation level and resubmit the experiment to obtain statistical moments with a prefixed error



Large-Scale Provenance-enriched Visualization

1. Visualizes experiment data through the workflow
 - ✓ Displays provenance data for information browsing
 - ✓ Provenance data collected by **Chiron** [5] or **SciCumulus** [6] workflow engines
2. Allows for selections and filtering
 - ✓ **Let scientists select just the results that interest them**
3. Stage-out only the data the scientists selected
4. Tracks which input parameters produced a given result
5. Display the results locally or on a display environment
 - ✓ Integration with tiled wall displays technologies
 - ✓ Currently implemented to use **SAGE** [7]
 - ✓ Can be extended to other platforms such as **CGLX** or **DisplayCluster**

15/09/12

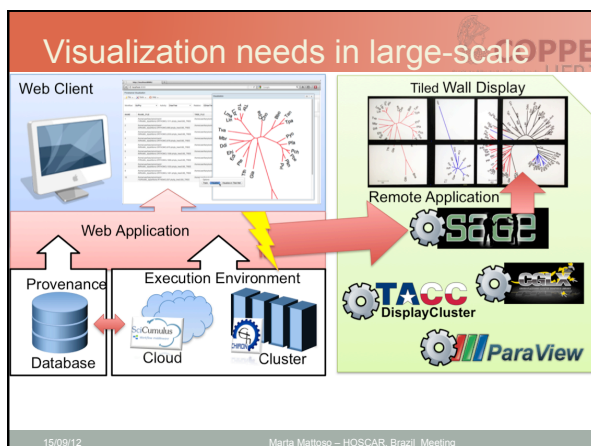
Marita Mattoso – HOSCAR, Brazil Meeting

Visualization needs in large-scale

1. Go fast to the results you need to analyze
 - ✓ Easy navigation over workflow data and provenance data
2. Filter results and stage out only the data you need to analyze
 - ✓ Staging data out can be costly (remote clusters, clouds)
3. Use high-resolution display environments
 - ✓ Visualize multiple results to establish comparisons
 - On parameters exploration scenarios
 - ✓ Analyze highly detailed images and simulations
 - ✓ **Can take advantage of available tiled display technologies such as SAGE, CGLX, TACC DisplayCluster and Paraview**

15/09/12

Marita Mattoso – HOSCAR, Brazil Meeting



15/09/12

Marita Mattoso – HOSCAR, Brazil Meeting


Data explorations with Wf

- The execution of variations upon the same workflow repeatedly
 - ✓ Parameter Sweeps
 - ✓ Fine-tuning (tolerance, solver options, ...)
 - ✓ Iterative methods
 - Stop criteria defined by the scientist evaluation
- Interfere in the dataflow
 - ✓ Input parameters
 - ✓ Workflow configurations
- Dynamic nature of simulations/ large data

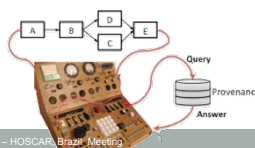
15/09/12

Marita Mattoso – HOSCAR, Brazil Meeting

Objectives




- Interactive Workflows
- Adaption and user-steering on workflow execution
 - ✓ Parameter space slicing
 - To compare results from different explored slices
 - ✓ Workflow adjustments
 - Human intervention
 - Dynamic steering
- Improvements
 - ✓ Quality of results
 - ✓ Execution efficiency
 - ✓ Scientific Data analysis

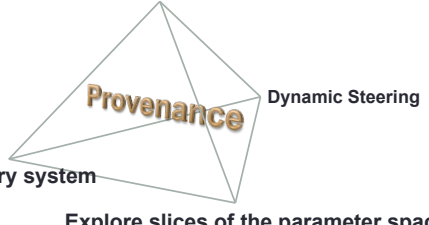


15/09/12 Marita Maltoso – HOSCAR, Brazil Meeting

Using provenance




Interactive Workflows

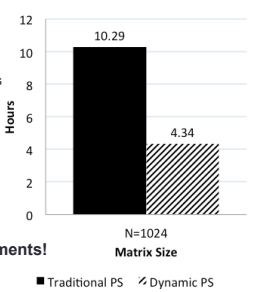


15/09/12 Marita Maltoso – HOSCAR, Brazil Meeting

Dynamic Experiments




- Optimization problems
 - ✓ Genetic Algorithms
- Uncertainty Quantification
 - ✓ Adaptive sparse grid colocation methods
- Reduced order models
 - ✓ Numerical solutions for heat conduction
- Bioinformatics
 - ✓ Comparative genomics
 - ✓ Phylogeny
- **Good results on preliminary experiments!**
 - ✓ (Dias et al., WORKS at SC 2011)



Method	Hours
Traditional PS	10.29
Dynamic PS	4.34

15/09/12 Marita Maltoso – HOSCAR, Brazil Meeting

Parameter Space Slicing



- Workflow executes choosing different slices of the parameter space at run time
 - ✓ To take initial decisions
 - ✓ To compare results
- We can make the slices addressable
- Parameter Space = Relation
 - ✓ Slices can be represented as horizontal fragments

$$R_i = \sigma_{F_i}(R), 1 \leq i \leq w$$

where F_i is the fragmentation predicate or a mintern

15/09/12 Marita Maltoso – HOSCAR, Brazil Meeting

Example

R			
IID	ModelFile	Flexibility	Curvature
1	I-01	3.2	0.8
2	I-02	4.5	0.9
3	I-03	5.6	1.2
4	I-04	5.2	1.1

$$M = \{ m_1, m_2 \mid \begin{array}{l} m_1 : Curvature < 1.0, \\ m_2 : Curvature \geq 1.0 \end{array} \}$$

$T \leftarrow \text{SlicedMap}(ga, M, R)$

T					
IID	Mãe	Pai	ModelFile	Flexibility	Curvature
1	2	2	I-02M	3.1	0.5
2	1	1	I-01M	5.1	1.4
3	3	3	I-03	5.6	1.2
4	4	4	I-04	4.9	1.0
5	3	4	I-03x04	6.1	1.5

15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting

Evaluating our approach

- Real experiments as **Algebraic Workflows**
 - ✓ Bioinformatics experiments
 - Working with the biochemist Kary Ocaña
 - ✓ Numerical Simulations
 - Partnership with prof. Alvaro Coutinho's team
 - ✓ Uncertainty Quantification
 - Partnership with Prof. Fernando Rochinha's team
 - ✓ Oil and Gas experiments
 - Petrobras
 - ✓ Characterization of PS Workflows - Evaluation Framework
 - PS workflow patterns • workload configuration • performance metrics

15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting

Scientific Workflows in Clouds

- Many scientists are migrating their experiments to clouds, including bioinformatics ones
- They are not required to assemble expensive computational infrastructure to execute their experiments
- They do not need to configure many pieces of software
- However, clouds are changing environments and they may be susceptible to performance fluctuations during the execution course of the workflow

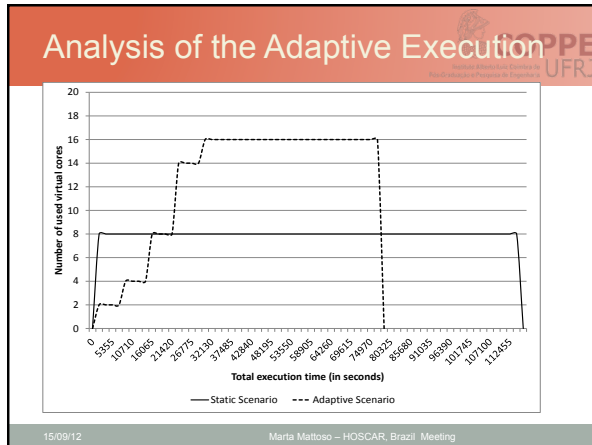
15/09/12 HOSCAR

SciCumulus Workflow Engine

<https://sourceforge.net/projects/scicumulus/>

- Workflow engine that manages the parallel execution of workflow activities in cloud environments
- Parameter sweep
- Data parallelism
- Static and adaptive modes

15/09/12 OLIVEIRA, D.; OGASAWARA, E.; BAIÃO, F.; MATTOSO, M. L. O. SciCumulus: A Lightweight Cloud Middleware to Explore Many Task Computing Paradigm in Scientific Workflows. In: Proc. of the 3rd IEEE CLOUD, 2010, p. 378-385



Querying provenance data

• "Get workflow ,executions, activity names that executed without error"

```

SELECT w.tag,
a.tag,
t.execution_status,
t.processor,
t.workspace,
t.status,
t.endtime,
t.starttime,
extract ('epoch' from (t.endtime-t.starttime)) || ', as duration
from hworkflow w, hactivity a, hexecution e
where w.wkfid = a.wkfid
and a.actid = t.actid
and not exists (select * from hactivity
where a2.actid = t.actid
and a2.exists)
order by w.wkfid
    
```

hactivity	actid	processor	workspace	status	endtime	starttime	duration
hactivity	actid	processor	workspace	status	endtime	starttime	duration

15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting

Other real Bio Experiments... (one more in E-Science 2012)

Adaptive Execution and cost model

OLIVEIRA, D., Ocaña, K. A. C. S., Ogasawara, E., Dias, J., Goncalves, J., Mattoso, M., (2012), "Cloud-based Phylogenomic Inference of Evolutionary Relationships: A Performance Study". 2nd International Workshop on Cloud Computing and Scientific Applications (CCSA), Ottawa.

Phylogenomic analysis

Parallel execution with blocking activities

Ocaña, K. A. C. S., OLIVEIRA, D. de, Horta, F., Dias, J., Ogasawara, E., Mattoso, M., (2012), "Exploring Molecular Evolution Reconstruction Using a Parallel Cloud-based Scientific Workflow". Brazilian Symposium on Bioinformatics (BSB 2012), Campo Grande, MT.

Evolutionary analysis

Adaptive Execution


OCANA, K.; OLIVEIRA, D.; DIAS, J.; OGASAWARA, E.; MATTOSO, M. L. Q. . Optimizing Phylogenetic Analysis Using SciHm Cloud-based Scientific Workflow. 7th IEEE e Science conference. IEEE Computer Society, 2011.

Comparative genomics

15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting

- ### Conclusions
- Exploratory nature of experiments
 - ✓ Parameter explorations
 - ✓ Iterative methods
 - Algebraic approach for workflows
 - ✓ Data uniformity
 - ✓ Workflow execution optimizations
 - Interactive workflows
 - ✓ Fine-tuning adjustments during Wf execution
 - Parameter space slicing
 - Iterative support
 - Real time provenance
 - ✓ Experiment Optimization
- 15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting

Acknowledgements




- CNPq, CAPES, FAPERJ and Petrobras
- INRIA
- at COPPE / UFRJ:
 - ✓ NACAD's support team and CPU time

**Matisse series of parallel paintings
"execution"**



Matisse-paires-et-series- George Pompidou, 2012

15/09/12 Marta Mattoso – HOSCAR, Brazil Meeting



**EXPLORING PROVENANCE IN HIGH
PERFORMANCE SCIENTIFIC COMPUTING**

THANKS!
marta@cos.ufrj.br