# A METHOD TO BUILD AND ANALYZE SCIENTIFIC WORKFLOWS FROM PROVENANCE THROUGH PROCESS MINING

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#### MOTIVATION

- Before creating scientific workflows, the provenance can only be captured from provenance enabled applications.
- It is often very hard to manually create and maintain scientific workflows.
- Can we leverage existing provenance to build scientific workflows automatically?

### **MOTIVATION (CONT)**

- Process mining has become an active research area in the past decade,
- Process mining synthesizes a process model from event logs,
- We aim to automatically generate a scientific workflow model from provenance using established process mining techniques
  - + Offers an effective approach for creating an initial scientific workflow model,
  - + Facilitates analysis techniques such as simulation and verification for detecting potential scientific workflow design problems,
  - Helps to discover hidden dependencies among different scientific workflows,
  - + Supports automated synthesis of existing scientific workflows

#### A METHOD TO BUILD AND ANALYZE SCIENTIFIC WORKFLOWS FROM PROVENANCE

- Issues when applying process mining in the context of scientific workflow
  - + Control flow mining
    - In this paper we focus on control flow mining
  - + Data dependency
  - + Incremental mining



# TOOLS

- ProM is a generic open-source framework for implementing process mining tools in a standard environment.
- The ProM framework accepts input logs in the <u>XES</u> or <u>MXML</u> format.
- The ProM framework has plugins for process mining, analysis, moni toring and conversion.
  - + Conversion from event logs in relational databases to XES or MXML.
  - + We have converted provenance in Taverna and Kepler to XES / MXML.



http://prom.sourceforge.net/

#### **RESULTS OF PROCESS DISCOVERY ALGORITHMS**

|                    | Description  | Result  |
|--------------------|--|---|
| Fuzzy Miner        | Provides a <b>zoomable view</b> of<br>scientific workflows by<br>controlling significance cutoff<br>to show tasks at different<br>importance levels.                               | Under certain significance cutoff, the fuzzy miner<br>successfully gives the changed part and<br>unchanged part. The fuzzy miner gets most<br>dependency correctly in the original scientific<br>workflows, but includes some non-existent<br>dependency. |
| Alpha Miner        | Provides a view of <b>direct</b><br><b>succession</b> between tasks in<br>provenance.  | Assuming the completeness of direct succession,<br>the alpha miner fails to give a view close to the<br>original scientific workflow.   |
| Genetic<br>Miner   | Provides a view of <b>frequency</b><br>for both tasks and succession<br>between tasks, and discovers<br>all common control-flow<br>structures assuming the<br>existence of noises. | The genetic miner gets a good view of structures<br>and frequencies, yet gives some wrong<br>dependencies which do not exist in both the<br>original scientific workflows and the results of the<br>fuzzy miner.  |
| Heuristic<br>Miner | Provides a view of scientific<br>workflows by considering <b>long</b><br><b>distance dependency</b> .  | The heuristic miner gives long distance<br>dependency successfully, but gives too much<br>dependency for some tasks such as<br>ReadCSVFileColumnNames.  |

## CONCLUSION

- A method using process mining to build and analyze scientific workflows
- The method can be applied to provenance data in many different forms
  - + it is quite straight forward to translate the provenance to XES format acceptable to process mining tools