

# WORKFLOW COMPOSITION AND DESCRIPTION TOOL

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Keywords: Workflow, Workflow description language, Directed acyclic graph.

Abstract: In this paper, we present a new approach for creating workflow. The workflow is represented as sequence of tasks with explicitly defined input/output data. Parallelism between tasks is implicitly defined by the data dependence. The workflows described in our approach are easily converted to other format. Nested and parameterized workflows are also supported.

## 1 INTRODUCTION

With the advance of computational technologies, the scientific applications running on modern distributed systems became more and more complex. Each execution of the applications is usually a workflow of several connected steps, where the output of the previous steps are the input of the next steps. Therefore, the tasks have to be executed in the correct order and the data need to be transferred between tasks in order to get the correct results.

At the moment, there are many existing workflow management systems, each system has its own language for describing the workflows. The way how the workflows are described in current systems are rather complex and inflexible. Some systems come also with graphical editors for creating the workflows easier.

In this paper, we present a new approach for creating workflows for scientific applications. Also our approach is applicable elsewhere, we primarily focus on distributed systems, where each task is an execution of a program (script, binary executable) on target hardware platforms. Most of grid workflow management systems have the same characteristics, so we will compare our approach with these workflow managers.

## 2 OVERVIEW OF WORKFLOW DESCRIPTION APPROACHES

Each workflow description consists from two parts: description of tasks and description of dependences

between tasks. Each task may have several properties like execution code, input/output data, command-line arguments, requirements on hardware and so on. There are two main approaches to describe these properties of tasks: in a plain text form as pairs of property name and value (e.g. *CPUNumber = 4*), or in XML language where task properties are elements or attributes.

Beside the task description, the dependence between tasks in the workflows must be also described in the workflow languages. There are two main ways to describe dependence between tasks in workflows: using parallel/sequence instructions and using directed acyclic graphs.

In the first approach, a workflow is consisted of (nested) parallel or sequential blocks of tasks. Tasks that can be executed in parallel are placed in blocks with parallel instruction, otherwise, in a block with sequential instruction; the tasks must be executed in the order as they are defined in the block. An example of workflow described in this way is as follows:

```
SEQ
  Task1
  PAR
    Task2
    Task3
  Task4
```

In this example, the workflow has four tasks named *Task1*, ..., *Task4*. The first task *Task1* must finish before *Task2* and *Task3* can start. *Task2* and *Task3* can be executed in parallel (or in any order), and *Task4* must wait until both tasks finish. For example Karajan (Gregor, et al., 2007) in Cog Kit (Gregor, et al., 2001) uses this approach for describing

workflows.

In the second approach, the dependences between tasks are described as by parent-child pair. Children tasks must wait until all parent tasks finish before starting. The workflow above can be described in this approach as follows:

```
PARENT Task1 CHILD Task2, Task3
PARENT Task2, Task3 CHILD Task4
```

Majority of scientific workflows use this approach for describing dependence. Typical examples are JDL (Job Description Language) (E. Laure, et al., 2006) which are used by gLite (gLite, 2011), DAGMan (J. Frey, 2002) in Condor (Condor project, 2011), SCULF (Kostas, et al., 2004) in Taverna (D. Hull, et al., 2006), Pegasus (K. Lee, et al., 2008). The main advantage of this approach is that it can describe more complex workflows than the first approach. The dependences can be visualized as directed acyclic graphs (DAG), where tasks are represented by nodes of the graphs and the directed edges show the parent-child relationships.

### 3 PROGRAMMABLE WORKFLOW DESCRIPTION

In this section we will describe our approach for describing workflow. We will start with basic ideas and gradually to more complex cases.

#### 3.1 Basic Ideas

Therefore, we use a simpler way to describe workflows as follows:

- A task in a workflow is described by triple: its code, a set of input data and a set of output data.
- A workflow is described as a sequence of tasks.
- Dependence and parallelism among tasks are implicitly defined by the input/output of tasks.

An example of a workflow is follows:

```
My_workflow(input, result)
  Task(code1, input, data1)
  Task(code2, data1, data2)
  Task(code3, data1, data3)
  Task(code4, [data2, data3], result)
```

In the code above, *input* and *result* are the lists of input and output data of the workflow. The items in the lists are usually the names of files containing corresponding data. The first task uses code in file *code1* for processing data from *input* and produces data stored in files in *data1*. Similarly, second and third tasks use *data1* as input and generate *data2*, or

*data3* respectively. Finally the last task use data from *data2* and *data3* for create *result*, which is also the output of whole workflow.

As it is shown in the example above, we only describe tasks, not the dependences among tasks. The dependence is implicitly defined by the input/output data of tasks. For example, second task use data produced by first task, so it must wait until the first task finishes.

We can prove the equivalence of workflows in our approach and workflows described by directed acyclic graphs by following statements:

- Every workflow represented in DAG can be described in our approach.
- Every workflow described in our approach can be converted to DAG with linear complexity  $O(N)$ .

So, in our approach, we can omit the part describing dependence between tasks and save cost of creating workflows. However, the main advantages of our approach are in the following sections.

#### 3.2 Nested Workflows

In the example above, the workflow has the same structure as the task: the code (the body of the workflow), input and output data. Therefore, we can define sub-workflows and use them in the way like tasks.

```
My_sub_workflow(input, output)
  Task(code5, input, data1)
  ...
My_workflow(input, result)
  Task(code1, input, data1)
  Workflow(My_sub_workflow, data1,
  data2)
  Task(code3, data1, data3)
  Task(code4, [data2, data3], result)
```

In the example above, the second task is replaced by a sub-workflow. The syntax is similar to calling sub-programs/functions in high-level programming language: the sub-workflow command in the main workflow will replace the formal input/output parameters of the sub-workflow by actual data in the main workflows.

Nested workflows are very useful for defining workflows with repeated patterns (a group of tasks doing the same actions with different input/output data). Like functions/subprograms in classical languages, they also make abstractions of task groups and make the workflows more readable.

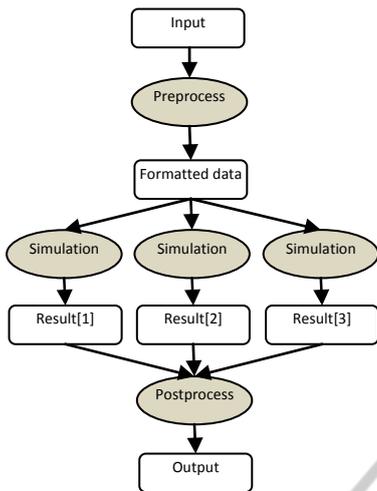


Figure 1: Graphical presentation of workflow.

### 3.3 Scripting a Parameterized Workflows

In some cases, workflows have parameters and the final forms of the workflows are known only after giving values to the parameters. The typical example is fork-join workflows, where the number of forks is a parameter given at runtime. Fork-join workflows are widely used in application with Monte-Carlo simulations or parametric studies.

We allow users to define workflows as scripts with loops and other control constructions. An example of fork-join workflows is as follows:

```
Myworkflow(input, output, N)
  Task(preprocess, input, local)
  for i = 1 to N
    Task(simulation, local,
result[i])
  Task(postprocess, result, output)
```

In the example above, we expand the workflow definition by adding other literal parameters beside the lists of input and output data. Unlike data in the input/output list, the values of the parameters must be known at the time the workflows are created.

### 3.4 Task Properties

As it was said in Section 2, tasks may have several properties like command-line arguments, hardware and software requirements, virtual organizations and so on. Some properties are usually common for all tasks in the workflows, other are specific for every task.

Users can define common properties of tasks by setting default values for these properties. Users can set values for properties of specific task using refe-

rence to the task. An example is as follows

```
My_workflow(input, result)
  default. set("VO", "egee")
  Task(code1, input, data1)
  Workflow(My_sub_workflow, data1,
data2)
  Task(code3, data1, data3)
  t = Task(code4, [data2,data3], result)
  t.set("CPU", "4")
```

In the example above, we set default virtual organization for all tasks to "egee" and special hardware requirements for the last task to 4 CPUs. The list of all valid properties is dependent to the execution backend of the workflows.

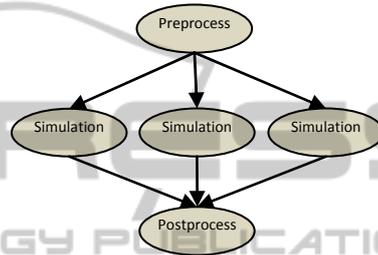


Figure 2: Graphical presentation of workflow in DAG.

### 3.5 Visualization

Workflow described in our approach can be visualized as graph of tasks and data. Each task has directed edges from/to their input/output data. Figure 1 shows the graphical representation of the fork-join example in Section 3.3 with N=3. In comparison with the classical DAGs (Figure 2), the graph in Figure 1 is more informative. If we remove nodes representing data items by connecting edges to them with edges from them, the graph in Figure 1 is identical with the DAG in Figure 2.

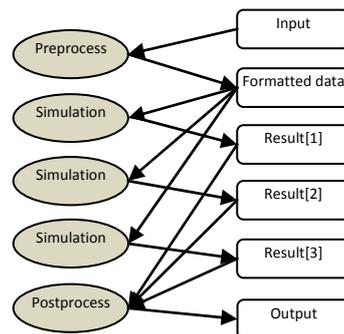


Figure 3: Internal structure of workflow.

It is worth to note that the graph in Figure 1 can be generated with linear complexity. We don't have to analyze every pair of tasks to know if they have

data dependence, but just read the task description and make connection to the its data. As every task is read only once, the complete graph can be generated with linear complexity. As the graph in Figure 1 is easily converted to DAG like in Figure 2, we can prove the statement in Section 3.1 that the workflow description in our approach can be converted to DAG with linear complexity.

## 4 IMPLEMENTATION DETAILS

We use Python scripting language for implementing a workflow composition tool for processing workflow description in our approach. With Python, we can easily write define workflows with loops and/other control instructions.

Internally, the workflow internally consists of two lists: list of tasks and list of data. Each task in the workflow is an object in memory with references to its data. Figure 3 shows the internal memory structures of workflows: the list of tasks on the left side, the list of data on the right side, and references between tasks and data.

It is interesting to see that the internal data structure of the workflow in Figure 3 is exactly the graphical representation of the workflow like in Figure 1. It means that, once the workflow description is processed, we have already graphical representation of the workflow in DAG form in memory. Therefore, it is easy to export the workflow description to any other formats compatible with DAG.

The workflow composition tool can execute the workflow in three ways:

- Execute the workflow locally: The aim of this mode is for experimentation, verification and debugging of the composed workflow.
- Use other workflow manager as backend: For executing workflows in Grid, we have implemented the export of our workflow description to JDL format.
- Use native workflow manager: a native workflow manager is still in development. It would exploit the explicit data declaration for performing direct data transfer among tasks.

## 5 CONCLUSIONS AND FUTURE WORK

In this paper, we have presented our approach for flexible workflow description. Workflows are

described as a list of tasks with input/output data explicitly defined. This approach supports nested workflows, loops and other control instructions in Python scripting language, which is used in our implementation. The workflows then can be easily exported to other formats with low complexity.

In the near future, we are developing a distributed workflow management system natively based on our approach of workflow description. The native workflow manager would allow direct data transfer among tasks according to the input/output data, without use of storage elements, what would minimize data transfer and increase performance. The workflow management system is designed, implementation is going on.

## ACKNOWLEDGEMENTS

This work is supported by projects SMART ITMS: 26240120005, SMART II ITMS: 26240120029, VEGA 2/0184/10.

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