



# **JUBE Documentation**

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## JUBE TUTORIAL

This tutorial is meant to give you an overview about the basic usage of *JUBE*.

### 1.1 Installation

Requirements: *JUBE* needs **Python 2.7** or **Python 3.2** (or higher)

To use the *JUBE* cmdline tool the `PHYTONPATH` must contain the position of the *JUBE* package

- You can use the **installation tool** to copy all files to the right position (preferred):

```
>>> python setup.py install --user
```

This will install the *JUBE* package and the binary to your `$HOME/.local` directory.

- You can also add **parent folder path** of the *JUBE* package-folder to the `PHYTONPATH` environment variable:

```
>>> export PHYTONPATH=<parent folder path>:$PHYTONPATH
```

- You can move the *JUBE* package by hand to an existing Python package folder like `site-packages`

To use the *JUBE* cmdline tool like a normal cmdline command you can add it to the `PATH`:

```
>>> export PATH=$HOME/.local/bin:$PATH
```

### 1.2 Hello World

In this example we will show you the basic structure of a *JUBE* input file and the basic cmdline options.

The files used for this example can be found inside `examples/hello_world`.

The input file `hello_world.xml`:

```
1 <?xml version="1.0" encoding="UTF-8"?>
2 <jube>
3   <benchmark name="hello_world" outpath="bench_run">
4     <comment>A simple hello world</comment>
5
6     <!-- Configuration -->
7     <parameterset name="parameterset">
8       <parameter name="hello_str">Hello World</parameter>
9     </parameterset>
10
11    <!-- Operation -->
12    <step name="say_hello">
13      <use>parameterset</use> <!-- use existing parameterset -->
14      <do>echo $hello_str</do> <!-- shell command -->
```

```
15     </step>
16   </benchmark>
17 </jube>
```

Every *JUBE* input file starts (after the general *XML* header line) with the root tag `<jube>`. This root tag must be unique. *XML* does not allow multiple root tags.

The first tag which contains benchmark specific information is `<benchmark>`. `hello_world` is the benchmarkname which can be used to identify the benchmark (e.g. when there are multiple benchmarks inside a single input file, or when different benchmarks using the same run directory).

The `outpath` describes the benchmark run directory (relative to the position of the input file). This directory will be managed by *JUBE* and will be automatically created if it doesn't exist. The directory name and position are very important, because they are the main interface to communicate with your benchmark, after it was submitted.

Using the `<comment>` you can store some benchmark related comment inside the benchmark directory. You can also use normal *XML*-comments to structure your input-file:

```
<!-- your comment -->
```

In this benchmark a `<parameterset>` is used to store the single `<parameter name="hello_str">` parameter. The name of the parameterset must be unique (relative to the current benchmark). In further examples we will see that there are more types of sets, which can be distinguished by their names. Also the name of the parameter must be unique (relative to the parameterset).

The `<step>` contains the operation tasks. The name must be unique. It can use different types of existing sets. Only sets, which are explicitly used, are available inside the step! The `<do>` contains a single **shell command**. This command will run inside of a sandbox directory environment (inside the `outpath` directory tree). The step and its corresponding parameterspace is named *workpackage*.

**Available** parameters can be used inside the shell commands. To use a parameter you had to write

```
$parametername
```

or

```
${parametername}
```

The brackets must be used if you want some variable concatenation. `$hello_strtest` will not be replaced, `${hello_str}test` will be replaced. If a parameter doesn't exist or isn't available the variable will not be replaced! If you want to use `$` inside your command, you had to write `$$` to mask the symbol. Parameter substitution will run before the normal shell substitution!

To run the benchmark just type:

```
>>> jube run hello_world.xml
```

This benchmark will produce the following output:

```
#####
# benchmark: hello_world
```

```
A simple hello world
```

```
#####
```

```
Running workpackages:
```

```
##### ( 1/ 1)
```

```
  stepname | all | open | wait | done
-----+-----+-----+-----+-----
say_hello |   1 |   0 |   0 |   1
```

```
>>>> Benchmark information and further useful commands:
```

```
>>>>     id: 0
```

```
>>>>     dir: bench_run
```

```
>>>> analyse: jube analyse bench_run --id 0
>>>> result: jube result bench_run --id 0
>>>> info: jube info bench_run --id 0
#####
```

You see, that inside the benchmark execution there was a single step `say_hello` which run one shell command `echo Hello World`.

The **id** is (in addition to the benchmark directory) an important number. Every benchmark run will get a new unique **id** inside the benchmark directory.

Inside the benchmark directory you will see the following structure:

```
bench_run          # the given outpath
|
+- 000000          # the benchmark id
|
+- configuration.xml # the stored benchmark configuration
+- workpackages.xml # workpackage information
+- 000000_say_hello # the workpackage
|
+- done            # workpackage finished marker
+- work            # user sandbox folder
|
+- stderr          # standard error messages of used shell commands
+- stdout          # standard output of used shell commands
```

`stdout` will contain `Hello World` in this example case.

## 1.3 Help

*JUBE* contains a commandline based help functionality:

```
>>> jube help <keyword>
```

With this command you will have direct access to all keywords inside the *glossary*.

Another useful command is the `info` command. It will show you information concerning your existing benchmarks:

```
1 # display a list of existing benchmarks
2 >>> jube info <benchmark-directory>
3 # display information about given benchmark
4 >>> jube info <benchmark-directory> -- id <id>
5 # display information about a step inside the given benchmark
6 >>> jube info <benchmark-directory> -- id <id> --step <stepname>
```

The third, but very important, functionality is the **logger**. Every run, continue, analyse and result execution will produce a new log file inside your execution directory. This file contains much useful debugging output.

## 1.4 Parameterspace creation

In this example we will show you an important feature of *JUBE*: The automatic parameterspace generation.

The files used for this example can be found inside `examples/parameterspace`.

The input file `parameterspace.xml`:

```
1  <?xml version="1.0" encoding="UTF-8"?>
2  <jube>
3    <benchmark name="parameterspace" outpath="bench_run">
4      <comment>A parameterspace creation example</comment>
5
6      <!-- Configuration -->
7      <parameterset name="parameterset">
8        <!-- Create a parameterspace out of two template parameter -->
9        <parameter name="number" type="int">1,2,4</parameter>
10       <parameter name="text" separator=";">Hello;World</parameter>
11     </parameterset>
12
13     <!-- Operation -->
14     <step name="say_hello">
15       <use>parameterset</use> <!-- use existing parameterset -->
16       <do>echo "$text $number"</do> <!-- shell command -->
17     </step>
18   </benchmark>
19 </jube>
```

Whenever a parameter contains a , (this can be changed using the `separator` attribute) this parameter becomes a **template**. A step which **uses the parameterset** containing this parameter will run multiple times to iterate over possible parameter combinations. In this example the step `say_hello` will run 6 times:

```
stepname | all | open | wait | done
-----+-----+-----+-----+-----
say_hello | 6 | 0 | 0 | 6
```

Every parameter combination will run in its own sandbox directory.

Another new keyword is the `type` attribute. The parameter type isn't used inside the substitution process, but it is used for sorting operation inside the result creation. The default type is `string`. Possible basic types are `string`, `int` and `float`.

## 1.5 Step dependencies

If you start writing a complex benchmark structure, you want to have dependencies between different *steps*. For example between a compile and the execution step. *JUBE* can handle these dependencies and will also preserve the given parameterspace.

The files used for this example can be found inside `examples/dependencies`.

The input file `dependencies.xml`:

```
1  <?xml version="1.0" encoding="UTF-8"?>
2  <jube>
3    <benchmark name="dependencies" outpath="bench_run">
4      <comment>A Dependency example</comment>
5
6      <!-- Configuration -->
7      <parameterset name="parameterset">
8        <parameter name="number" type="int">1,2,4</parameter>
9      </parameterset>
10
11     <!-- Operations -->
12     <step name="first_step">
13       <use>parameterset</use> <!-- use existing parameterset -->
14       <do>echo $number</do> <!-- shell command -->
15     </step>
16
17     <!-- Create a dependency between both steps -->
18     <step name="second_step" depend="first_step">
```



```

19         <do>cat first_step/stdout</do> <!-- shell command -->
20     </step>
21 </benchmark>
22 </jube>

```

In this example we create a dependency between `first_step` and `second_step`. After `first_step` finished, the corresponding `second_step` will start. You are able to have multiple dependencies (separated by `,` in the definition), but circular definitions will not be resolved. A dependency is a unidirectional link!

To communicate between a step and its dependency there is a link inside the work directory pointing to the corresponding dependency step work directory. In this example we use

```
cat first_step/stdout
```

to write the `stdout`-file content of the dependency step into the `stdout`-file of the current step.

Because the `first_step` uses a template parameter which creates three execution runs. There will also be three `second_step` runs each pointing to a different `first_step`-directory:

stepname	all	open	wait	done
first_step	3	0	0	3
second_step	3	0	0	3

## 1.6 Loading files and substitution

Every step runs inside a unique sandbox directory. In normal cases you need external files inside this directory (e.g. the source files) and in some cases you want to change a parameter inside the file based on your current parameterspace. there are two addition set-types which handle this behaviour inside of *JUBE*.

The files used for this example can be found inside `examples/files_and_sub`.

The input file `files_and_sub.xml`:

```

1  <?xml version="1.0" encoding="UTF-8"?>
2  <jube>
3      <benchmark name="files_and_sub" outpath="bench_run">
4          <comment>A file copy and substitution example</comment>
5
6          <!-- Configuration -->
7          <parameterset name="parameterset">
8              <parameter name="number" type="int">1,2,4</parameter>
9          </parameterset>
10
11         <!-- Files -->
12         <fileset name="files">
13             <copy>file.in</copy>
14         </fileset>
15
16         <!-- Substitute -->
17         <substituteset name="substitute">
18             <!-- Substitute files -->
19             <iofile in="file.in" out="file.out" />
20             <!-- Substitute commands -->
21             <sub source="#NUMBER#" dest="$number" />
22         </substituteset>
23
24         <!-- Operation -->
25         <step name="sub_step">
26             <use>parameterset</use> <!-- use existing parameterset -->
27             <use>files</use> <!-- use existing fileset -->
28             <use>substitute</use> <!-- use existing substituteset -->

```

```
29     <do>cat file.out</do>    <!-- shell command -->
30   </step>
31 </benchmark>
32 </jube>
```

The content of file `file.in`:

```
Number: #NUMBER#
```

Inside the `<fileset>` the current location (relatively seen towards the current input file) of files is given. `<copy>` specify that the file should be copied to the sandbox directory when the fileset is used. Also a `<link>` option is available to create a symbolic link to the given file inside the sandbox directory.

The `<substituteset>` describe the substitution process. The `<iofile>` contains the input and output file. The path is relatively seen towards the sandbox directory. Because we do/should not know that location we used the fileset to copy `file.in` inside the directory.

The `<sub>` specify the substitution. All occurrence of `source` will be substituted by `dest`. As you can see, you can use Parameter inside the substitution to use your current parametersapce.

In the `sub_step` we use all available sets. The use order isn't relevant. The normal execution process will be:

1. Parameterspace expansion
2. Copy/link files
3. File substitution
4. Run shell operations

The resulting directory-tree will be:

```
bench_run          # the given outpath
|
+- 000000          # the benchmark id
|
  +- configuration.xml # the stored benchmark configuration
  +- workpackages.xml  # workpackage information
  +- 000000_say_hello  # the workpackage ($number = 1)
  |
    +- done           # workpackage finished marker
    +- work           # user sanbox folder
    |
      +- stderr        # standard error messages of used shell commands
      +- stdout        # standard output of used shell commands (Number: 1)
      +- file.in       # the file copy
      +- file.out      # the substituted file
  +- 000001_say_hello  # the workpackage ($number = 2)
  |
    +- ...
  +- ...
```

## 1.7 Creating a result table

Finally, after running the benchmark, you will get several directories. *JUBE* allows you to parse your result files to extract relevant data (e.g. walltime information) and create a result table.

The files used for this example can be found inside `examples/result_creation`.

The input file `result_creation.xml`:

```
1 <?xml version="1.0" encoding="UTF-8"?>
2 <jube>
3   <benchmark name="result_creation" outpath="bench_run">
```

```

4      <comment>A result creation example</comment>
5
6      <!-- Configuration -->
7      <parameterset name="parameterset">
8          <!-- Create a parameterspace out of two template parameter -->
9          <parameter name="number" type="int">1,2,4</parameter>
10     </parameterset>
11
12     <!-- Regex pattern -->
13     <patternset name="pattern">
14         <pattern name="number_pat" type="int">Number: $jube_pat_int</pattern>
15     </patternset>
16
17     <!-- Operation -->
18     <step name="write_number">
19         <use>parameterset</use> <!-- use existing parameterset -->
20         <do>echo "Number: $number"</do> <!-- shell command -->
21     </step>
22
23     <!-- Analyse -->
24     <analyzer name="analyse">
25         <use>pattern</use> <!-- use existing patternset -->
26         <analyse step="write_number">
27             <file>stdout</file> <!-- file which should be scanned -->
28         </analyse>
29     </analyzer>
30
31     <!-- Create result table -->
32     <result>
33         <use>analyse</use> <!-- use existing analyzer -->
34         <table name="result" style="pretty" sort="number">
35             <column>number</column>
36             <column>number_pat</column>
37         </table>
38     </result>
39 </benchmark>
40 </jube>

```

Using `<parameterset>` and `<step>` we create three *workpackages*. Each writing `Number: $number` to `stdout`.

Now we want to parse these `stdout` files to extract information (in this example case the written number). First of all we had to declare a `<patternset>`. Here we can describe a set of `<pattern>`. A `<pattern>` is a regular expression which will be used to parse your result files and search for a given string. In this example we only have the `<pattern>` `number_pat`. The name of the pattern must be unique (based on the usage of the `<patternset>`). The `type` is optional. It is used when the extracted data will be sorted. The regular expression can contain other pattern or parameter. The example uses `$jube_pat_int` which is a *JUBE* given default pattern matching integer values. The pattern must contain a group, given by brackets `(...)`, to declare the extraction part (`$jube_pat_int` already contains these brackets).

To use your `<patternset>` you had to specify the files which should be parsed. This can be done using the `<analyzer>`. It uses relevant patternsets, and inside the `<analyse>` a step-name and a file inside this step is given. Every workpackage file combination will create its own result entry.

To run the analyse you had to write:

```
>>> jube analyse bench_run
```

The analyse data will be stored inside the benchmark directory.

The last part is the result table creation. Here you had to used an existing analyzer. The `<column>` contains a pattern or a parameter name. `sort` is the optionl sorting order (separated by `,`). The `style` attribute can be `csv` or `pretty` to get different ASCII representations.

To create the result table you had to write:

```
>>> jube result bench_run
```

The result table will be written to `STDOUT` and into a `result.dat` file inside `bench_run/<id>/result`.

Output of the given example:

number		number_pat
1		1
2		2
4		4

This was the last example of the basic *JUBE* tutorial. Next you can start the [advanced tutorial](#) to get more information about including external sets, jobssystem representation and scripting parameter.

## ADVANCED TUTORIAL

This tutorial will show more detailed functions and tools of *JUBE*. If you want a basic overview you should read the general *JUBE tutorial* first.

### 2.1 Schema validation

To validate your input files you can use DTD or schema validation. You will find `jube.dtd` and `jube.xsd` inside the `schema` folder. You had to add these schema information to your input files which you want to be validated.

DTD usage:

```
1 <?xml version="1.0" encoding="UTF-8"?>
2 <!DOCTYPE benchmarks SYSTEM "<jube.dtd path>">
3 <benchmark>
4 ...
```

Schema usage:

```
1 <?xml version="1.0" encoding="UTF-8"?>
2 <benchmarks xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"
3   xsi:noNamespaceSchemaLocation="<jube.xsd path>">
4 ...
```

Example validation tools:

- eclipse
- xmllint:
  - For validation (using the DTD):

```
>>> xmllint --noout --valid <xml input file>
```
  - For validation (using the DTD and Schema):

```
>>> xmllint --noout --valid --schema <schema file> <xml input file>
```

### 2.2 Scripting parameter

Sometimes you want to create parameter based on other parameter. In this cases you can use scripting parameter.

The files used for this example can be found inside `examples/scripting_parameter`.

The input file `scripting_parameter.xml`:

```
1 <?xml version="1.0" encoding="UTF-8"?>
2 <jube>
3   <benchmark name="scripting_parameter" outpath="bench_run">
4     <comment>A scripting parameter example</comment>
5
6     <!-- Configuration -->
7     <parameterset name="parameterset">
8       <!-- Normal template -->
9       <parameter name="number" type="int">1,2,4</parameter>
10      <!-- A template created by a scripting parameter-->
11      <parameter name="additional_number" mode="python" type="int">",".join(str(a*${number}) for a in range(1,${number}))</parameter>
12      <!-- A scripting parameter -->
13      <parameter name="number_mult" mode="python" type="float">${number}*${additional_number}</parameter>
14      <!-- Normal template -->
15      <parameter name="text">Number: $number</parameter>
16    </parameterset>
17
18    <!-- Operation -->
19    <step name="operation">
20      <use>parameterset</use> <!-- use existing parameterset -->
21      <!-- shell commands -->
22      <do>echo "number: $number, additional_number: $additional_number"</do>
23      <do>echo "number_mult: $number_mult, text: $text"</do>
24    </step>
25  </benchmark>
26 </jube>
```

In this example we see four different parameter.

- `number` is a normal template which expands to three different *workpackages*.
- `additional_number` is a scripting parameter which creates a new template and bases on `number`. The mode is set to the scripting language (python and perl are allowed). The additional type is optional and declare the result type after evaluating the expression. The `int` type is only used by the sort algorithm in the result step. It is not possible to create a template of different scripting parameter. Because of this second template we will get six different *workpackages*.
- `number_mult` is a small calculation. You can use any other existing parameter (which is used inside the same step).
- `text` a normal parameter which uses the content of another parameter. For simple concatenation parameter you need no scripting parameter.

For this example we will find the following output inside the `jube.log`-file:

```
===== operation =====
>>> echo "number: 1, additional_number: 1"
>>> echo "number_mult: 1, text: Number: 1"
===== operation =====
>>> echo "number: 1, additional_number: 2"
>>> echo "number_mult: 2, text: Number: 1"
===== operation =====
>>> echo "number: 2, additional_number: 2"
>>> echo "number_mult: 4, text: Number: 2"
===== operation =====
>>> echo "number: 2, additional_number: 4"
>>> echo "number_mult: 8, text: Number: 2"
===== operation =====
>>> echo "number: 4, additional_number: 4"
>>> echo "number_mult: 16, text: Number: 4"
===== operation =====
>>> echo "number: 4, additional_number: 8"
>>> echo "number_mult: 32, text: Number: 4"
```

Scripting inside the `<do>` or any other position is not possible. If you want to use some scripting expressions you had to create a new parameter.

## 2.3 Jobsystem

In most cases you want to submit jobs by *JUBE* to your local machines jobsystem. You can use the normal file access and substitution system to prepare your jobfile and send it to the jobsystem. *JUBE* also provide some additional features.

The files used for this example can be found inside `examples/jobsystem`.

The input jobsystem file `job.run.in` for *Torque/Moab* (you can easily adapt your personal jobscript):

```

1  #!/bin/bash -x
2  #MSUB -l nodes=#NODES#:ppn=#PROCS_PER_NODE#
3  #MSUB -l walltime=#WALLTIME#
4  #MSUB -e #ERROR_FILEPATH#
5  #MSUB -o #OUT_FILEPATH#
6  #MSUB -M #MAIL_ADDRESS#
7  #MSUB -m #MAIL_MODE#
8
9  ### start of jobscript
10
11 #EXEC#
12 touch #READY#

```

The *JUBE* input file `jobsystem.xml`:

```

1  <?xml version="1.0" encoding="UTF-8"?>
2  <jube>
3    <benchmark name="jobsystem" outpath="bench_run">
4      <comment>A jobsystem example</comment>
5
6      <!-- benchmark configuration -->
7      <parameterset name="parameterset">
8        <parameter name="number" type="int">1,2,4</parameter>
9      </parameterset>
10
11     <!-- Job configuration -->
12     <parameterset name="executeset">
13       <parameter name="submit_cmd">msub</parameter>
14       <parameter name="job_file">job.run</parameter>
15       <parameter name="nodes" type="int">1</parameter>
16       <parameter name="walltime">00:01:00</parameter>
17       <parameter name="ppn" type="int">4</parameter>
18       <parameter name="ready_file">ready</parameter>
19       <parameter name="mail_mode">abe</parameter>
20       <parameter name="mail_address"></parameter>
21       <parameter name="err_file">stderr</parameter>
22       <parameter name="out_file">stdout</parameter>
23       <parameter name="exec">echo $number</parameter>
24     </parameterset>
25
26     <!-- Load jobfile -->
27     <fileset name="files">
28       <copy>${job_file}.in</copy>
29     </fileset>
30
31     <!-- Substitute jobfile -->
32     <substituteset name="sub_job">
33       <iofile in="${job_file}.in" out="$job_file" />
34       <sub source="#NODES#" dest="$nodes" />

```

```
35     <sub source="#PROCS_PER_NODE#" dest="$ppn" />
36     <sub source="#WALLTIME#" dest="$walltime" />
37     <sub source="#ERROR_FILEPATH#" dest="$err_file" />
38     <sub source="#OUT_FILEPATH#" dest="$out_file" />
39     <sub source="#MAIL_ADDRESS#" dest="$mail_address" />
40     <sub source="#MAIL_MODE#" dest="$mail_mode" />
41     <sub source="#EXEC#" dest="$exec" />
42     <sub source="#READY#" dest="$ready_file" />
43 </substituteset>
44
45 <!-- Operation -->
46 <step name="submit" work_dir="$WORK/jobsystem_bench_${jube_benchmark_id}_${jube_wp_id}" >
47     <use>parameterset</use>
48     <use>executeset</use>
49     <use>files,sub_job</use>
50     <do done_file="$ready_file">$submit_cmd $job_file</do> <!-- shell command -->
51 </step>
52 </benchmark>
53 </jube>
```

As you can see the jobfile is very general and several parameter will be used for replacement. By using a general jobfile and the substitution mechanism you can control your jobsystem directly out of your *JUBE* input file.

The submit command is a normal *Shell* command so there are no special tags needed.

There are two new attributes:

- `done_file` inside the `<do>` allows you set a filename/path which should be used by the jobfile to mark the end of execution. *JUBE* doesn't when the job ends. normally it will return when the *Shell* command was finished. When using a jobsystem we had to wait until the jobfile was executed. If *JUBE* found a `<do>` containing a `done_file` attribute *JUBE* will return directly and will not continue automatically until the `done_file` exists. If you want to check the current status of your running steps and continue the benchmark process if possible you can type:

```
>>> jube continue benchmark_run
```

This will continue your benchmark execution.

- `work_dir` can be used to change the sandbox work directory of a step. In normal cases *JUBE* checks that every work directory get a unique name. When changing the directory the user must select a unique name by his own. For example he can use `$jube_benchmark_id` and `$jube_wp_id` which are *JUBE* internal parameter and will expand to the current benchmark and workpackage id. Files and directories out of a given `<filesset>` will be copied to the new work directory. Other automatic links, like the dependency links, will not be created!

You will see this Output after running the benchmark:

```
stepname | all | open | wait | done
-----+-----+-----+-----+-----
submit  |   3 |   0 |   3 |   0
```

and this output after running the `continue` command (after the jobs where executed):

```
stepname | all | open | wait | done
-----+-----+-----+-----+-----
submit  |   3 |   0 |   0 |   3
```

Every workpackage continue, but because there are no more additional operations there is no more work to be done.

You had to run `continue` multiple times if not all `done_file` were written when running `continue` for the first time.



## 2.4 Include

As you seen in the example before a benchmark can be become very long. To structure your benchmark using multiple files and to reuse existing sets there are three different include features inside of *JUBE*.

The files used for this example can be found inside `examples/include`.

The include file `include_data.xml`:

```

1  <?xml version="1.0" encoding="UTF-8"?>
2  <jube>
3    <parameterset name="parameterset">
4      <parameter name="number" type="int">1,2,4</parameter>
5    </parameterset>
6
7    <parameterset name="parameterset2">
8      <parameter name="text">Hello</parameter>
9    </parameterset>
10
11   <dos>
12     <do>echo Test</do>
13     <do>echo $number</do>
14   </dos>
15 </jube>

```

All files which contains data that should be included use the *XML*-format. The include files can have a user specific structure (there can be none valid *JUBE* tags like `<dos>`), but the structure must be allowed by the searching mechanism (see below). The resulting file must have a valid *JUBE* structure.

The main file `main.xml`:

```

1  <?xml version="1.0" encoding="UTF-8"?>
2  <jube>
3    <benchmark name="include" outpath="bench_run">
4      <comment>A include example</comment>
5
6      <!-- use parameterset out of an external file and add a additional parameter -->
7      <parameterset name="parameterset" init_with="include_data.xml">
8        <parameter name="foo">bar</parameter>
9      </parameterset>
10
11     <!-- Operation -->
12     <step name="say_hello">
13       <use>parameterset</use> <!-- use existing parameterset -->
14       <use from="include_data.xml">parameterset2</use> <!-- out of an external file -->
15       <do>echo $foo</do> <!-- shell command -->
16       <include from="include_data.xml" path="dos/do" /> <!-- include all available tag -->
17     </step>
18   </benchmark>
19 </jube>

```

In these file there are three different include types.

The `init_with` can be used inside any set definition. Inside the given file the searching mechanism will search for the same set (same type, same name), will parse its structure (this must be *JUBE* valid) and copy the content to `main.xml`. Inside `main.xml` you can add additional values or can overwrite existing ones. If your include-set use a different name inside your include file you can use `init_with="filename.xml:new_name"`.

The second method is the `<use from="...">`. This is mostly the same like the `init_with` structure but in this case you are not able to add or overwrite some values. The external set will be used directly. There is no set-type inside the `<use>`, because of that, the setname must be unique inside the include-file.

The last method is the most generic include. By using `<include />` you can copy any nodes you want to your main-XML file. The path is optional and can be used to select a specific nodeset (otherwise the root-node will

be included). The `<include />` is the only include-method that can be used to include any tag you want. The `<include />` will copy all parts without any changes. The other include types will change pathnames, which were relative to the include-file position.

To run the benchmark you can use the normal command:

```
>>> jube run main.xml
```

It will search for include files inside four different positions:

- inside the same directory of your `main.xml`
- inside a directory given over the commandline:

```
>>> jube run --include-path some_path another_path -- main.xml
```

- inside any path given with the `JUBE_INCLUDE_PATH` environment variable:

```
>>> export JUBE_INCLUDE_PATH=some_path:another_path
```

- inside any path given by a `<include-path>`-tag:

```
1  <?xml version="1.0" encoding="UTF-8"?>
2  <benchmarks>
3    <include-path>
4      <path>some_path</path>
5      <path>another_path</path>
6    </include-path>
7    ...
```

## GLOSSARY

**analyse** Analyse an existing benchmark. The analyzer will scan through all files given inside the configuration by using the given patternsets.

If no benchmark id is given, last benchmark found in directory will be used. If benchmark directory is missing, current directory will be used.

**analyzer\_tag** The analyzer describe the steps and files which should be scanned using a set of pattern.

```
<analyzer name="...">
  <use from="...">...</use>
  ...
  <analyse step="...">
    <file>...</file>
  </analyse>
  ...
</analyzer>
```

- you can use different patternsets to analyse a set of files
- only patternsets are useable
- using patternsets `<use>set1,set2</use>` is the same as `<use>set1</use><use>set2</use>`
- the from-attribute is optional and can be used to specify an external set source
- any name must be unique, it is not allowed to reuse a set
- the step-attribute contains an existing stepname
- each file using each workpackage will be scanned separately

**benchmark\_tag** The main benchmark definition

```
<benchmark name="..." outpath="...">
  ...
</benchmark>
```

- container for all benchmark information
- benchmark-name must be unique inside input file
- outpath contains the path to the root folder for benchmark runs
  - multiple benchmarks can use the same folder
  - every benchmark and every (new) run will create a new folder (named by an unique benchmark id) inside this given outpath
  - the path will be relative to input file location

**comment** Add or manipulate the comment string.

If no benchmark id is given, last benchmark found in directory will be used. If benchmark directory is missing, current directory will be used.

**continue** Continue an existing benchmark. Not finished steps will be continued, if they are leaving pending mode.

If no benchmark id is given, last benchmark found in directory will be used. If benchmark directory is missing, current directory will be used.

**copy\_tag** A copy can be used to copy a file or directory from your normal filesystem to your sandbox work directory.

```
<copy directory="..." name="..." rel_path_ref="...">...</copy>
```

- `directory` is optional, it can be used if you want to copy several files inside the same directory
- `name` is optional, it can be used to rename the file inside your work directory
- `rel_path_ref` is optional
  - `external` or `internal` can be chosen, default: `external`
  - `external`: rel.-paths based on position of xml-file
  - `internal`: rel.-paths based on current work directory (e.g. to link files of another step)
- each copy-tag can contain a list of filenames (or directories), separated by ,
  - if `name` is present, the lists must have the same length
- you can copy all files inside a directory by using `directory/*`
  - this can't be mixed using `name`
- in the execution step the given files or directories will be copied

### **directory\_structure**

- every (new) benchmark run will create its own directory structure
- every single workpackage will create its own directory structure
- user can add files (or links) to the workpackage dir, but the real position in filesystem will be seen as a blackbox
- general directory structure:

```
benchmark_runs (given by "outpath" in xml-file)
|
+- 000000 (determined through benchmark-id)
  |
  +- 000000_compile (step: just an example, can be arbitrary chosen)
    |
    +- work (user environment)
    +- done (workpackage finished information file)
    +- ... (more jube internal information files)
  +- 000001_execute
    |
    +- work
      |
      +- compile -> ../../000000_compile/work (automatic generated link for depending
    +- wp_done_00 (single "do" finished, but not the whole workpackage)
    +- ...
  +- 000002_execute
  +- result (result data)
  +- configuration.xml (benchmark configuration information file)
  +- workpackages.xml (workpackage graph information file)
  +- analyse.xml (analyse data)
+- 000001 (determined through benchmark-id)
  |
  +- 000000_compile (step: just an example, can be arbitrary chosen)
  +- 000001_execute
  +- 000002_postprocessing
```

**fileset\_tag** A fileset is a container to store a bundle of links and copy commands.

```
<fileset name="..." init_with="...">
  <link>...</link>
  <copy>...</copy>
  ...
</fileset>
```

- `init_with` is optional
  - if the given filepath can be found inside of the `JUBE_INCLUDE_PATH` and if it contains a fileset using the given name, all link and copy will be copied to the local set
  - the name of the external set can differ to the local one by using `init-with="filename.xml:external_name"`
- link and copy can be mixed within one fileset (or left)
- filesets can be used inside the step-command

### general\_structure

```
<?xml version="1.0" encoding="UTF-8"?>
<jube>
  <!-- optional additional include pathes -->
  <include-path>
    <path>...</path>
    ...
  </include-path>
  <!-- optional benchmark selection -->
  <selection>
    <only>...</only>
    <not>...</not>
    ...
  </selection>
  <!-- global sets -->
  <parameterset>...</parameterset>
  <substitutionset>...</substitutionset>
  <fileset>...</fileset>
  <patternset>...</patternset>
  ...
  <benchmark>
    <!-- optional benchmark comment -->
    <comment>...</comment>
    <!-- local benchmark parametersets -->
    <parameterset>...</parameterset>
    ...
    <!-- files, which should be used -->
    <fileset>...</fileset>
    ...
    <!-- substitution rules -->
    <substituteset>...</substituteset>
    ...
    <!-- pattern -->
    <patternset>...</patternset>
    ...
    <!-- commands -->
    <step>...</step>
    ...
    <!-- analyse -->
    <analyzer>...</analyzer>
    ...
    <!-- result -->
```

```
<result>...</result>
...
</benchmark>
...
</jube>
```

**include\_tag** Include *XML*-data from an external file.

```
<include from="..." path="..." />
```

- `<include>` can be used to include an external *XML*-structure into the current file
- can be used at every position (inside the `<jube>`-tag)
- path is optional and can be used to give an alternative xml-path inside the include-file (default: root-node)

**info** Show info for the given benchmark directory, a given benchmark or a specific step.

If benchmark directory is missing, current directory will be used.

**iofile\_tag** A iofile declare the name (and path) of a file used for substitution.

```
<iofile in="..." out="..." />
```

- in and out filepath are relative to the current work directory for every single step (not relative to the path of the inputfile)
- in and out must be different

**jube\_pattern** List of available jube pattern:

- \$jube\_pat\_int: integer number
- \$jube\_pat\_nint: integer number, skip
- \$jube\_pat\_fp: floating point number
- \$jube\_pat\_nfp: floating point number, skip
- \$jube\_pat\_wrd: word
- \$jube\_pat\_nwrld: word, skip
- \$jube\_pat\_bl: blank space (variable length), skip

**link\_tag** A link can be used to create a symbolic link from your sandbox work directory to a file or directory inside your normal filesystem.

```
<link directory="..." name="..." rel_path_ref="...">...</link>
```

- directory is optional, it can be used if you want to link several files inside the same directory
- name is optional, it can be used to rename the file inside your work directory
- rel\_path\_ref is optional
  - external or internal can be chosen, default: external
  - external: rel.-pathes based on position of xml-file
  - internal: rel.-pathes based on current work directory (e.g. to link files of another step)
- each link-tag can contain a list of filenames (or directories), separated by ,
  - if name is present, the lists must have the same length
- in the execution step the given files or directories will be linked

**parameter\_tag** A parameter is a usable configuration option.

```
<parameter name="..." mode="..." type="..." separator="...">...</parameter>
```

- a parameter can be seen as variable: Name is the name to use the variable, and the text between the tags will be the real content
- name must be unique inside the given parameterset
- type is optional (only used for sorting, default: string)
- mode is optional (used for script-types, default: text)
- separator is optional, default: ,
- if the text contains the given (or the implicit) separator, a template will be created
- use of another parameter:
  - inside the parameter definition, a parameter can be reused: ... \$nameofparameter ...
  - the parameter will be replaced multiply times (to handle complex parameter structures; max: 5 times)
  - the substitution will be run before the execution step starts with the current parameter space. Only parameters reachable in this step will be useable for substitution!
- Scripting modes allowed:
  - mode="python": allow python snippets (using `eval <cmd>`)
  - mode="perl": allow perl snippets (using `perl -e "print <cmd>"`)
- Templates can be created, using scripting e.g.: `" , ".join([str(2**i) for i in range(3)])`

**parameterset\_tag** A parameterset is a container to store a bundle of parameter.

```
<parameterset name="..." init_with="...">
  <parameter>...</parameter>
  ...
</parameterset>
```

- parameterset-name must be unique (can't be reuse inside substitutionsets or filesets)
- init\_with is optional
  - if the given filepath can be found inside of the `JUBE_INCLUDE_PATH` and if it contains a parameterset using the given name, all parameters will be copied to the local set
  - local parameters will overwrite imported parameters
  - the name of the external set can differ to the local one by using `init-with="filename.xml:external_name"`
- parametersets can be used inside the step-command
- parametersets can be combined inside the step-tag, but they must be compatible:
  - Two parametersets are compatible if the parameter intersection (given by the parameter-name), only contains parameter based on the same definition
  - These two sets are compatible:

```
<parameterset name="set1">
  <parameter name="test">1,2,4</parameter>
  <parameter name="test2">foo</parameter>
</parameterset>
<parameterset name="set2">
  <parameter name="test">1,2,4</parameter>
  <parameter name="test3">bar</parameter>
</parameterset>
```

- These two sets aren't compatible:

```
<parameterset name="set1">
  <parameter name="test">1,2,4</parameter>
  <parameter name="test2">foo</parameter>
</parameterset>
<parameterset name="set2">
  <parameter name="test">2</parameter> <!-- Template in set1 -->
  <parameter name="test2">bar</parameter> <!-- Other content in set2 -->
</parameterset>
```

**remove** The given benchmark will be removed.

If no benchmark id is given, last benchmark found in directory will be removed.

Only the *JUBE* internal directory structure will be deleted. External files and directories will stay unchanged.

If no benchmark id is given, last benchmark found in directory will be used. If benchmark directory is missing, current directory will be used.

**result** Create a result table.

If no benchmark id is given, last benchmark found in directory will be used. If benchmark directory is missing, current directory will be used.

**run** Start a new benchmark run by parsing the given **JUBE** input file.

**selection\_tag** Select benchmarks by name.

```
<selection>
  <only>...</only>
  <not>...</not>
  ...
</selection>
```

- select or unselect a benchmark by name
- only selected benchmarks will run (when using the `run` command)
- multiple `<only>` and `<not>` are allowed
- `<only>` and `<not>` can contain a name list divided by ,

**step\_tag** A step give a list of *Shell* operations and a corresponding parameter environment.

```
<step name="..." work_dir="..." shared="...">
  <use from="...">...</use>
  ...
  <do stdout="..." stderr="..." active="...">...</do>
  <do done_file="...">...</do>
  <do shared="true">...</do>
  ...
</step>
```

- parametersets, filesets and substitutionsets are useable
- using filesets and substitutesets `<use>set1,set2</use>` is the same as `<use>set1</use><use>set2</use>`
- using parametersets `<use>set1</use><use>set2</use>` means: use both; `<use>set1,set2</use>` means: use in one case the first set and in second case the other set
- the `from` attribute is optional and can be used to specify an external set source
- any name must be unique, it is **not allowed to reuse** a set
- `work_dir` is optional and can be used to switch to an alternative work directory



- the user had to handle **uniqueness of this directory** by his own
- no automatic parent/children link creation
- `shared` is optional and can be used to create a shared folder which can be accessed by all workpackages based on this step
  - a link, named by the attribute content, is used to access the shared folder
  - the shared folder link will not be automatically created in an alternative working directory!
- `do` can contain any *Shell*-syntax-snippet (parameter will be replaced ... `$nameofparameter` ...)
- `stdout-` and `stderr-`filename are optional (default: `stdout` and `stderr`)
- `active` is optional
  - can be set to `true` or `false` to enable or disable the single command
  - parameter are allowed inside this attribute
- `done_file`-filename is optional
  - by using `done_file` the user can mark `async`-steps. The operation will stop until the script will create the named file inside the work directory.
- `shared="true"`
  - can be used inside a step using a shared folder
  - `cmd` will be **executed inside the shared folder**
  - `cmd` will run once (synchronize all workpackages)
  - `$jube_wp_... -` parameter can't be used inside the shared command

**sub\_tag** A substitution expression.

```
<sub source="..." dest="..." />
```

- `source`-string will be replaced by `dest`-string
- both can contain parameter: ... `$nameofparameter` ...

**substituteset\_tag** A `substituteset` is a container to store a bundle of subs.

```
<substituteset name="..." init_with="...">
  <iofile/>
  ...
  <sub/>
  ...
</substituteset>
```

- `init_with` is optional
  - if the given filepath can be found inside of the `JUBE_INCLUDE_PATH` and if it contains a `substituteset` using the given name, all `iofile` and `sub` will be copied to the local set
  - local `iofile` will overwrite imported ones based on `out`, local `sub` will overwrite imported ones based on `source`
  - the name of the external set can differ to the local one by using `init-with="filename.xml:external_name"`
- `substitutesets` can be used inside the step-command

**workpackage** A workpackage is the combination of a *step* (which contains all operations) and one parameter setting out of the expanded parameterspace.

Every workpackage will run inside its own sandbox directory!



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