

## **JUBE Documentation**

Release 2.0.0

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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 commandline 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 commandline tool like a normal commandline 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 commandline options.

The files used for this example can be found inside examples/hello\_world.

The input file hello\_world.xml:

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

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 follwing output:

You see, that inside the benchmark execution there was a single step <code>say\_hello</code> which run one shell command echo <code>Hello</code> <code>World</code>.

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 follwing 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 sanbox 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:

```
# display a list of existing benchmarks

| >>> jube info <benchmark-directory>
| display information about given benchmark  
| >>> jube info <benchmark-directory> -- id <id>  
| display information about a step inside the given benchmark  
| >>> 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.

You can also use the debugging mode:

```
>>> jube --debug <command> [other-args]
```

This mode avoid any shell execution but will generate log files.

## 1.4 Parameterspace creation

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

1.3. Help 3

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

The input file parameterspace.xml:

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

```
<?xml version="1.0" encoding="UTF-8"?>
   <jube>
     <benchmark name="dependencies" outpath="bench_run">
       <comment>A Dependency example
       <!-- Configuration -->
       <parameterset name="parameterset">
         <parameter name="number" type="int">1,2,4</parameter>
       </parameterset>
9
10
       <!-- Operations -->
11
       <step name="first_step">
12
         <use>parameterset</use> <!-- use existing parameterset -->
13
         <do>echo $number</do> <!-- shell command -->
       </step>
```

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:

## 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:

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

```
<use>parameterset</use> <!-- use existing parameterset -->

<use>files</use> <!-- use existing fileset -->

<use>substitute</use> <!-- use existing substituteset -->

<use>cdo>cat file.out</do> <!-- shell command -->

</step>

</benchmark>

</jube>
```

The content of file file.in:

```
Number: #NUMBER#
```

Inside the <fileset> the current location (relativly 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 a 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 relativly 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)
                      # workpackage finished marker
      +- done
       +- 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:

```
<?xml version="1.0" encoding="UTF-8"?>
   <jube>
2
     <benchmark name="result_creation" outpath="bench_run">
       <comment>A result creation example
4
       <!-- Configuration -->
6
       <parameterset name="parameterset">
         <!-- Create a parameterspace out of two template parameter -->
         <parameter name="number" type="int">1,2,4</parameter>
       </parameterset>
10
11
       <!-- Regex pattern -->
12
       <patternset name="pattern">
13
         <pattern name="number_pat" type="int">Number: $jube_pat_int</pattern>
14
       </patternset>
15
16
       <!-- Operation -->
17
18
       <step name="write_number">
         <use>parameterset <!-- use existing parameterset -->
19
         <do>echo "Number: $number"</do> <!-- shell command -->
20
21
       </step>
22
       <!-- Analyse -->
23
       <analyzer name="analyse">
24
         <use>pattern</use> <!-- use existing patternset -->
25
         <analyse step="write_number">
26
           <file>stdout</file> <!-- file which should be scanned -->
27
         </analyse>
       </analyzer>
29
30
31
       <!-- Create result table -->
32
       <result>
         <use>analyse</use> <!-- use existing analyzer -->
33
         34
           <column>number</column>
35
           <column>number_pat</column>
36
         37
       </result>
38
     </benchmark>
39
   </jube>
```

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 anlayse 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 optional 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, jobsystem representation and scripting parameter.

## **ADVANCED TUTORIAL**

This tutorial will show you 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 validate.

#### DTD usage:

```
<?xml version="1.0" encoding="UTF-8"?>
<!DOCTYPE benchmarks SYSTEM "<jube.dtd path>">
<benchmark>
Schema usage:
 <?xml version="1.0" encoding="UTF-8"?>
 <benchmarks xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"</pre>
   xsi:noNamespaceSchemaLocation="<jube.xsd path>">
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 a parameter which based on the value of another parameter. In this case you can use scripting parameter.

The files used for this example can be found inside examples/scripting\_parameter.

The input file scripting\_parameter.xml:

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

In this example we see four different parameter.

- number is a normal template which will be expanded 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 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 do not need 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):

```
#!/bin/bash -x
   #MSUB -1 nodes=#NODES#:ppn=#PROCS_PER_NODE#
   #MSUB -1 walltime=#WALLTIME#
   #MSUB -e #ERROR FILEPATH#
   #MSUB -o #OUT_FILEPATH#
   #MSUB -M #MAIL ADDRESS#
   #MSUB -m #MAIL_MODE#
   ### start of jobscript
10
   #EXEC#
11
   touch #READY#
12
   The JUBE input file jobsystem.xml:
   <?xml version="1.0" encoding="UTF-8"?>
   <jube>
2
     <benchmark name="jobsystem" outpath="bench_run">
       <comment>A jobsystem example
       <!-- benchmark configuration -->
       <parameterset name="parameterset">
         <parameter name="number" type="int">1,2,4</parameter>
       </parameterset>
       <!-- Job configuration -->
11
       <parameterset name="executeset">
12
         <parameter name="submit_cmd">msub</parameter>
13
         <parameter name="job_file">job.run</parameter>
14
         <parameter name="nodes" type="int">1</parameter>
15
         <parameter name="walltime">00:01:00</parameter>
16
         <parameter name="ppn" type="int">4</parameter>
17
         <parameter name="ready_file">ready</parameter>
18
         <parameter name="mail_mode">abe</parameter>
19
         <parameter name="mail_address"></parameter>
20
         <parameter name="err_file">stderr</parameter>
21
         <parameter name="out_file">stdout</parameter>
22
          <parameter name="exec">echo $number</parameter>
23
       </parameterset>
24
25
       <!-- Load jobfile -->
26
       <fileset name="files">
27
         <copy>${job_file}.in</copy>
28
       </fileset>
29
30
31
       <!-- Substitute jobfile -->
       <substituteset name="sub_job">
32
         <iofile in="${job_file}.in" out="$job_file" />
33
         <sub source="#NODES#" dest="$nodes" />
34
```

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

There are two new attributes:

• done\_file inside the <do> allows you set a filename/path to a file which should be used by the jobfile to mark the end of execution. *JUBE* doesn't know 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. the position of the done\_file is relativly seen to the work directory.

• 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 <fileset> 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:

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

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 external data

As you seen in the example before a benchmark can become very long. To structure your benchmark youe can use multiple files and 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:

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

All files which contains data that should be included must 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:

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

In these file there are three different inleude 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 update 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 (in the following order):

- 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:

## 2.5 Tagging

Tagging is a easy way to hide selectable parts of your input file.

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

The input file tagging.xml:

```
<?xml version="1.0" encoding="UTF-8"?>
   <jube>
2
     <benchmark name="tagging" outpath="bench_run">
3
       <comment>A simple tagging example
       <!-- Configuration -->
6
       <parameterset name="parameterset">
         <parameter name="hello_str" tag="eng">Hello</parameter>
         <parameter name="hello_str" tag="deu">Hallo</parameter>
         <parameter name="world_str" tag="!deu">World</parameter>
10
       </parameterset>
11
12
       <!-- Operation -->
13
       <step name="say_hello">
14
         <use>parameterset</use> <!-- use existing parameterset -->
15
         <do>echo '$hello_str $world_str'</do> <!-- shell command -->
17
       </step>
     </benchmark>
18
   </jube>
```

When running this example:

```
>>> jube run tagging.xml
```

all <tags> which contain a special tag="..." attribute will be hidden. !deu stands for not deu so this tag will not be hidden when running the command.

The result inside the stdout file will be

```
$hello_str World
```

because there was no alternative to select the \$hello\_str.

When running this example using a specific tag:

```
>>> jube run tagging.xml --tag eng
```

the result inside the stdout file will be

```
Hello World
```

The tag attribute or the commandline expression can also contain a list of different names. A hidden <tag> will be ignored completely! If there is no alternative this can produce a wrong execution behaviour!

The tag attribute can be used inside every <tag> inside the input file (except the <jube>).

## 2.6 Platform independent benchmarking

If you want to create platform independent benchmarks you can use the include features inside JUBE.

All platform related sets must be declared in a includeable file e.g. platform.xml. There can be multiple platform.xml in different directories to allow different platforms. By changing the include-path the benchmark changes its platform specific data.

An example benchmark structure bases on three include files:

- The main benchmark include file which contain all benchmark specific but platform independent data
- A mostly generic platform include file which contain benchmark independent but platform specific data (this can be created once and placed somewhere central on the system, it can be easily accessed using the <code>JUBE\_INCLUDE\_PATH</code> environment variable.
- A platform specific and benchmark specific include file which must be placed in a unique directory to allow inlcude-path usage

To avoid writting of long include-pathes every time you run a platform independent benchmark you can store the include-path inside your input file. This can be mixed using the tagging-feature:

Now you can run your benchmark using:

```
>>> jube run filename.xml --tag plat1
```

## 2.7 Multiple benchmarks

Often you only have one benchmark inside your input file. But it is also possible to store multiple benchmarks inside the same input file:

All benchmarks can use the same global (as a child of <jube>) declared sets. Often it might be better to use an include feature instead. *JUBE* will run every benchmark in the given order. Every benchmark gets an unique benchmark id.

To select only one benchmark you can use:

```
>>> jube run filename.xml --only-bench a
or:
>>> jube run filename.xml --not-bench b
```

This information can also be stored inside the input file:

## COMMANDLINE DOCUMENTATION

Here you will find a list of all available *JUBE* commandline options. You can also use:

```
jube -h
```

to get a list of all available commands.

Because of the *shell* parsing mechanism take care if you write your optional arguments after the command name before the positional arguments. You **must** use -- to split the ending of an optional (if the optional argument takes multiple input elements) and the start of the positional argument.

### 3.1 run

Run a new benchmark.

```
jube run [-h] [--only-bench ONLY_BENCH [ONLY_BENCH ...]]
      [--not-bench NOT_BENCH [NOT_BENCH ...]] [-t TAG [TAG ...]]
      [--hide-animation] [--include-path INCLUDE_PATH [INCLUDE_PATH ...]]
      [-a] [-r] [-m COMMENT] FILE [FILE ...]
```

- -h, --help show command help information
- --only-bench ONLY\_BENCH [ONLY\_BENCH ...] only run specific benchmarks given by benchmark name
- --not-bench NOT\_BENCH [NOT\_BENCH ...]] do not run specific benchmarks given by benchmark name
- -t TAG [TAG ...], --tag TAG [TAG ...] use specific tags when running this file. This will be used for tagging
- --hide-animation hide the progress bar animation (if you want to use JUBE inside a scripting environment)
- --include-path INCLUDE\_PATH [INCLUDE\_PATH ...] add additional include pathes where to search for include files
- -a, --analyse run analyse after finishing run command
- -r, --result run result after finishing run command (this will also start analyse)
- -m COMMENT, --comment COMMENT overwrite benchmark specific comment

```
FILE [FILE ...] input XML file
```

### 3.2 continue

Continue an existing benchmark.

```
jube continue [-h] [-i ID [ID ...]] [--hide-animation] [-a] [-r] [DIRECTORY]
```

- -h, --help show command help information
- -i ID [ID ...], --id ID [ID ...] select benchmark id, default: last found inside the benchmarks directory
- --hide-animation hide the progress bar animation (if you want to use JUBE inside a scripting environment)
- -a, --analyse run analyse after finishing run command
- -r, --result run result after finishing run command (this will also start analyse)

**DIRECTORY** directory which contain benchmarks, default: .

## 3.3 analyse

Run the analyse procedure.

- -h, --help show command help information
- -i ID [ID ...], --id ID [ID ...] select benchmark id, default: last found inside the benchmarks directory
- -u UPDATE\_FILE, --update UPDATE\_FILE use given input XML file to update patternsets, analyzer and result before running the analyse
- --include-path INCLUDE\_PATH [INCLUDE\_PATH ...] add additional include pathes where to search for include files (when using --update)
- -t TAG [TAG ...], --tag TAG [TAG ...] use specific tags when running this file. This will be used for tagging (when using --update)

**DIRECTORY** directory which contain benchmarks, default: .

#### 3.4 result

Run the result creation.

- -h, --help show command help information
- -i ID [ID ...], --id ID [ID ...] select benchmark id, default: last found inside the benchmarks directory
- -a, --analyse run analyse before running result command
- -u UPDATE\_FILE, --update UPDATE\_FILE use given input XML file to update patternsets, analyzer and result before running the analyse
- --include-path INCLUDE\_PATH [INCLUDE\_PATH ...] add additional include pathes where to search for include files (when using --update)
- -t TAG [TAG ...], --tag TAG [TAG ...] use specific tags when running this file. This will be used for tagging (when using --update)

-o RESULT\_NAME [RESULT\_NAME ...], -only RESULT\_NAME [RESULT\_NAME ...] only create specific results given by name

**DIRECTORY** directory which contain benchmarks, default: .

### 3.5 comment

Add or manipulate the benchmark comment.

```
jube comment [-h] [-i ID [ID ...]] [-a] comment [DIRECTORY]
```

- -h, --help show command help information
- -i ID [ID ...], --id ID [ID ...] select benchmark id, default: last found inside the benchmarks directory
- -a, --append append new comment instead of overwrite existing one

comment new comment

**DIRECTORY** directory which contain benchmarks, default: .

#### 3.6 remove

Remove an existing benchmark

```
jube remove [-h] [-i ID [ID ...]] [-f] [DIRECTORY]
```

- -h, --help show command help information
- -i ID [ID ...], --id ID [ID ...] select benchmark id, default: last found inside the benchmarks directory
- -f, --force do not prompt

**DIRECTORY** directory which contain benchmarks, default: .

## 3.7 info

Get benchmark specific information

```
jube info [-h] [-i ID [ID ...]] [-s STEP [STEP ...]] [DIRECTORY]
```

- -h, --help show command help information
- -i ID [ID ...], --id ID [ID ...] show benchmark specific information
- -s STEP [STEP ...], --step STEP [STEP ...] show step specific information

**DIRECTORY** show directory specific information

## 3.8 help

#### Command help

```
jube help [-h] [command]
```

-h, --help show command help information

command command to get help about

3.5. comment 19

## **FOUR**

### **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.

- 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 seperatly

## 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.

comment\_tag Add a benchmark specific comment. These comment will be stored inside the benchmark directory.

```
<comment>...</comment>
```

**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 filesytem 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.-pathes based on position of xml-file
  - internal: rel.-pathes 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)
```

**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>
    . . .
```

include-path\_tag Add some include pathes where to search for include files.

```
<include-path>
  <path>...</path>
   ...
</include-path>
```

• the additional path will be scanned for include files

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

jube\_variables List of available jube variables:

- Benchmark:
  - \$jube\_benchmark\_name: current benchmark name
  - \$jube\_benchmark\_id: current benchmark id
  - \$jube\_benchmark\_home: original input file location
- Step:
  - \$jube\_step\_name: current step name

- \$jube\_step\_iteratuions: number of step iterations (default: 1)
- Workpackage:
  - \$jube\_wp\_id: current workpackage id
  - \$jube\_wp\_iteration: current iteration number (default: 0)
  - \$jube\_wp\_parent\_<parent\_name>\_id: workpackage id of selected parent step
  - \$jube\_wp\_abspath: absolute path to workpackage work directory

**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="test">1,2,4</parameter>
    <parameter name="test3">bar</parameter>
</parameterset></parameterset></parameter></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset></parameterset>
```

- These two sets aren't compatible:

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.

- · 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.

- 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 substition 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

**tagging** Tagging is a simple way to include or exclude parts of your input file.

- Every available <tag> (not the root < jube>-tag) can contain a tag-attribute
- The tag-attribute can contain a list of names: tag="a, b, c" or "not" names: tag="a, !b, c"
- When running JUBE, multiple tags can be send to the input-file parser:

```
jube run <filename> --tag a b
```

- <tags> which doesn't contain one of these names will be hidden inside the include file
- <tags> which doesn't contain any tag-attribute will stay inside the include file
- "not" tags are more important than normal tags: tag="a,!b,c" and running with a b will hide the <tag> because the !b is more important than the a

**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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