

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 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.

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:

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```
<!-- 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
       <!-- Operation -->
13
       <step name="say_hello">
         <use>parameterset</use> <!-- use existing parameterset -->
         <do>echo "$text $number"</do> <!-- shell command -->
       </step>
17
     </benchmark>
18
  </jube>
19
```

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:

```
===== say_hello ======

>>> echo "Hello 1"

===== say_hello ======

>>> echo "Hello 2"

===== say_hello ======

>>> echo "World 1"

===== say_hello ======

>>> echo "World 2"

===== say_hello ======

>>> echo "World 4"
```

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">
7
         <parameter name="number" type="int">1,2,4</parameter>
       </parameterset>
10
       <!-- Operations -->
11
       <step name="first_step">
12
         <use>parameterset</use> <!-- use existing parameterset -->
13
         <do>echo $number</do> <!-- shell command -->
14
       </step>
15
16
```

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:

```
===== first_step ======

>>> echo 1

===== first_step ======

>>> echo 2

===== first_step ======

>>> echo 4

===== second_step ======

>>> cat first_step/stdout

===== second_step ======

>>> cat first_step/stdout

===== second_step ======

>>> cat first_step/stdout
```

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">
3
       <comment>A file copy and substitution example
4
5
       <!-- Configuration -->
6
       <parameterset name="parameterset">
         <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
17
       <substituteset name="substitute">
         <!-- Substitute files -->
```

```
<iofile in="file.in" out="file.out" />
19
         <!-- Substitute commands -->
20
         <sub source="#NUMBER#" dest="$number" />
21
       </substituteset>
22
23
       <!-- Operation -->
24
       <step name="sub_step">
25
         <use>parameterset</use> <!-- use existing parameterset -->
         <use>files</use> <!-- use existing fileset -->
27
         <use>substitute</use> <!-- use existing substituteset -->
28
         <do>cat file.out</do> <!-- shell command -->
29
       </step>
30
     </benchmark>
31
  </jube>
32
```

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 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:

```
# the given outpath
bench_run
+- 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
                       # user sanbox folder
     +- work
        +- stderr
                       # standard error messages of used shell commands
        +- stdout
                       # standard output of used shell commands (Number: 1)
                      # the file copy
        +- file.in
         +- 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>
     <benchmark name="result_creation" outpath="bench_run">
       <comment>A result creation example
       <!-- Configuration -->
       <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
       <step name="write_number">
18
         <use>parameterset</use> <!-- use existing parameterset -->
19
         <do>echo "Number: $number"</do> <!-- shell command -->
20
       </step>
21
22
       <!-- Analyse -->
23
       <analyzer name="analyse">
24
         <use>pattern</use> <!-- use existing patternset -->
25
         <analyse step="write_number">
           <file>stdout</file> <!-- file which should be scanned -->
27
         </analyse>
28
       </analyzer>
29
30
       <!-- Create result table -->
31
       <result>
32
         <use>analyse</use> <!-- use existing analyzer -->
33
         34
35
           <column>number</column>
36
           <column>number_pat</column>
37
         38
       </result>
39
     </benchmark>
   </jube>
40
```

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 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:

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:

```
<?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 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 get the following output:

```
===== 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>
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 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 <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:

```
===== submit =====

>>> msub job.run

Waiting for file "ready" ...

===== submit =====

>>> msub job.run

Waiting for file "ready" ...

===== submit =====

>>> msub job.run

Waiting for file "ready" ...
```

and this output after running the continue command:

```
===== submit ======
===== submit ======
```

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.

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THREE

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
- $\bullet\,$ 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 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)
+- 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 -->
```

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_nwrd: 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></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="set2">
        <parameter name="test">2</parameter> <!-- Template in set1 -->
        <parameter name="test">2</parameter> <!-- Other content in set2 -->
</parameterset></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.

- 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

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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