

Package ‘Rcwl’

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Title Wrap Command Tools and Pipelines Using CWL

Version 1.0.14

Description The package can be a simple and user-friendly way to manage command line tools and build data analysis pipelines in R using Common Workflow Language (CWL).

Depends R (>= 3.6), yaml, methods, S4Vectors

Imports utils, stats, BiocParallel, batchtools, DiagrammeR, shiny, R.utils, codetools

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+ , cwlStepParam , stepParam-method
Pipeline

Description

To build a pipeline by connecting multiple ‘stepParam’ to a ‘cwlStepParam’ object.

Usage

```
## S4 method for signature 'cwlStepParam,stepParam'
e1 + e2
```

Arguments

e1	A ‘cwlStepParam’ object.
e2	A ‘stepParam’ object.

Value

A ‘cwlStepParam’ object.

See Also

[cwlStepParam](#)

Description

The main CWL parameter class and constructor for command tools. More details: <https://www.commonwl.org/v1.0/CommandLineTool.html>

Usage

```
cwlParam(cwlVersion = "v1.0", cwlClass = "CommandLineTool",
         baseCommand = character(), requirements = list(), hints = list(),
         arguments = list(), id = character(), label = character(),
         inputs = InputParamList(), outputs = OutputParamList(),
         stdout = character(), expression = character(),
         extensions = list())
```

Arguments

cwlVersion	CWL version
cwlClass	"CommandLineTool"
baseCommand	Specifies the program or R function to execute
requirements	A list of Requirement lists that apply to either the runtime environment or the workflow engine.
hints	Any or a list for the workflow engine.
arguments	Command line bindings which are not directly associated with input parameters.
id	The unique identifier for this process object.
label	A short, human-readable label of this process object.
inputs	A object of ‘InputParamList’.
outputs	A object of ‘OutputParamList’.
stdout	Capture the command’s standard output stream to a file written to the designated output directory.
expression	Javascripts for ExpressionTool class.
extensions	A list of extensions and metadata

Details

<https://www.commonwl.org/v1.0/CommandLineTool.html>

Value

A ‘cwlParam’ class object.

Examples

```
input1 <- InputParam(id = "sth")
echo <- cwlParam(baseCommand = "echo", inputs = InputParamList(input1))
```

cwlShiny*cwlShiny***Description**

Function to generate shiny app automatically for a ‘cwlParam‘ object.

Usage

```
cwlShiny(cwl, inputList = list(), upload = FALSE, ...)
```

Arguments

<code>cwl</code>	A <code>cwlParam</code> object.
<code>inputList</code>	a list of choices for the inputs of <code>cwl</code> object. The name of the list must match the inputs of the <code>cwl</code> object.
<code>upload</code>	Whether to upload file. If <code>FALSE</code> , the <code>upload</code> field will be text input (file path) instead of file input.
<code>...</code>	More options for ‘runCWL’.

Value

A shiny webapp.

Examples

```
input1 <- InputParam(id = "sth")
echo <- cwlParam(baseCommand = "echo", inputs = InputParamList(input1))
echoApp <- cwlShiny(echo)
```

cwlStepParam-class*cwlStepParam***Description**

A workflow steps parameter, which connect multiple command line steps into a workflow. More details: `stepInParamList`.

Usage

```
cwlStepParam(cwlVersion = "v1.0", cwlClass = "Workflow",
            requirements = list(), id = character(), hints = list(),
            arguments = list(), extensions = list(), inputs = InputParamList(),
            outputs = OutputParamList(), stdout = character(),
            steps = stepParamList())
```

Arguments

<code>cwlVersion</code>	CWL version
<code>cwlClass</code>	"Workflow".
<code>requirements</code>	Requirements that apply to either the runtime environment or the workflow engine.
<code>id</code>	The unique identifier for this process object.
<code>hints</code>	Any or a list for the workflow engine.
<code>arguments</code>	Command line bindings which are not directly associated with input parameters.
<code>extensions</code>	A list of extensions and metadata.
<code>inputs</code>	A object of 'InputParamList'.
<code>outputs</code>	A object of 'OutputParamList'.
<code>stdout</code>	Capture the command's standard output stream to a file written to the designated output directory.
<code>steps</code>	A list of 'stepParamList'.

Value

An object of class 'cwlStepParam'.

Examples

```
input1 <- InputParam(id = "sth")
echo1 <- cwlParam(baseCommand = "echo",
                   inputs = InputParamList(input1))
input2 <- InputParam(id = "sthout", type = "File")
echo2 <- cwlParam(baseCommand = "echo",
                   inputs = InputParamList(input2),
                   stdout = "out.txt")
i1 <- InputParam(id = "sth")
o1 <- OutputParam(id = "out", type = "File", outputSource = "echo2/output")
wf <- cwlStepParam(inputs = InputParamList(i1),
                    outputs = OutputParamList(o1))
s1 <- Step(id = "echo1", run = echo1, In = list(sth = "sth"))
s2 <- Step(id = "echo2", run = echo2, In = list(sthout = "echo1/output"))
wf <- wf + s1 + s2
```

Description

cwlParam methods
 cwlVersion CWL document version
 cwlClass
 cwlClass
 baseCommand
 baseCommand

```
arguments
arguments
hints
hints
requirements
requirements
stdout of cwlParam
stdout of cwlParam
Extensions and metadata of cwlParam
```

Usage

```
cwlVersion(cwl)

cwlVersion(cwl) <- value

cwlClass(cwl)

cwlClass(cwl) <- value

baseCommand(cwl)

baseCommand(cwl) <- value

arguments(cwl, step = NULL)

arguments(cwl, step = NULL) <- value

hints(cwl)

hints(cwl) <- value

requirements(cwl)

requirements(cwl) <- value

stdOut(cwl)

stdOut(cwl) <- value

extensions(cwl)

extensions(cwl) <- value
```

Arguments

<code>cwl</code>	A ‘cwlParam’ object.
<code>value</code>	Assign value to the ‘cwlParam’ object.
<code>step</code>	To specify a step ID when ‘cwl’ is a workflow. It can be multiple levels of steps separated by “/” for nested workflow.

Value

cwlVersion: cwl version
 cwlClass: CWL Class
 baseCommand: CWL baseCommand
 arguments: CWL arguments
 hints: CWL hints
 requirements: CWL requirements
 stdOut: CWL stdout
 extensions: A list of extensions or metadata

InputArrayParam-class InputArrayParam

Description

Parameters for array inputs. To specify an array parameter, the array definition is nested under the type field with 'type: array' and items defining the valid data types that may appear in the array. More details: <https://www.commonwl.org/v1.0/CommandLineTool.html#CommandInputArraySchema>

Usage

```
InputArrayParam(label = "", type = "array", items = character(),
  prefix = "", separate = TRUE, itemSeparator = character(),
  valueFrom = character())
```

Arguments

label	A short description for this object
type	Must be "array".
items	Defines the type of the array elements.
prefix	Command line prefix to add before the value.
separate	If true (default), then the prefix and value must be added as separate command line arguments; if false, prefix and value must be concatenated into a single command line argument.
itemSeparator	Join the array elements into a single string with separator.
valueFrom	String or Expression.

Value

An object of class 'InputArrayParam'.

Examples

```
InputArrayParam(items = "string", prefix="-B=", separate = FALSE)
```

InputParam-class *Input parameters InputParam*

Description

parameter for a command tool. More details: <https://www.commonwl.org/v1.0/CommandLineTool.html#CommandInputParam>

Usage

```
InputParam(id, label = "", type = "string", doc = character(),
secondaryFiles = character(), streamable = logical(),
format = character(), loadListing = character(),
loadContents = logical(), position = 0L, prefix = "",
separate = TRUE, itemSeparator = character(),
valueFrom = character(), shellQuote = logical(),
default = character(), value = character())

## S4 method for signature 'cwlParam'
x$name

## S4 replacement method for signature 'cwlParam'
x$name <- value
```

Arguments

<code>id</code>	The unique identifier for this parameter object.
<code>label</code>	A short, human-readable label of this object.
<code>type</code>	valid types of data that may be assigned to this parameter.
<code>doc</code>	A documentation string for this type.
<code>secondaryFiles</code>	Only valid when type: File or is an array of items: File. Provides a pattern or expression specifying files or directories that must be included alongside the primary file.
<code>streamable</code>	Only valid when type: File or is an array of items: File. A value of true indicates that the file is read or written sequentially without seeking.
<code>format</code>	Only valid when type: File or is an array of items: File.
<code>loadListing</code>	Only valid when type: Directory or is an array of items: Directory.
<code>loadContents</code>	Only valid when type: File or is an array of items: File.
<code>position</code>	The position for this parameter.
<code>prefix</code>	Command line prefix to add before the value.
<code>separate</code>	If true (default), then the prefix and value must be added as separate command line arguments; if false, prefix and value must be concatenated into a single command line argument.
<code>itemSeparator</code>	Join the array elements into a single string with the elements separated by by itemSeparator.
<code>valueFrom</code>	String or Expression.
<code>shellQuote</code>	If ShellCommandRequirement is in the requirements for the current command, this controls whether the value is quoted on the command line (default is true).

default	The default value for this parameter
value	Assigned value for this parameter
x	A ‘cwlParam‘ object.
name	One one of input list

Value

An object of class ‘InputParam‘.

Examples

```
input1 <- InputParam(id = "sth")
```

InputParamList-class *InputParamList*

Description

InputParamList
InputParamList A list of InputParam
inputs

Usage

```
InputParamList(...)  
inputs(cwl)
```

Arguments

... The InputParam objects.
cwl A cwlParam object

Value

An object of class ‘InputParamList‘.
inputs: A list of ‘InputParam‘.

Examples

```
input1 <- InputParam(id = "sth")  
InputParamList(input1)  
## Inputs  
input1 <- InputParam(id = "sth")  
echo <- cwlParam(baseCommand = "echo", inputs = InputParamList(input1))  
inputs(echo)
```

OutputArrayParam-class

*Output array parameters***Description**

Parameters for array outputs. More details: <https://www.commonwl.org/v1.0/CommandLineTool.html#CommandOutput>

Usage

```
OutputArrayParam(label = character(), type = "array",
  items = character(), glob = character(), loadContents = logical(),
  outputEval = character())
```

Arguments

label	A short, human-readable label of this object.
type	Must be "array".
items	Defines the type of the array elements.
glob	Pattern to find files relative to the output directory.
loadContents	Read text from globbed file.
outputEval	Evaluate an expression to generate the output value.

Value

An object of class ‘OutputArrayParam’.

Examples

```
b <- OutputParam(id = "b", type = OutputArrayParam(items = "File"), glob = "*.txt")
```

OutputParam-class

*Output parameters***Description**

An output parameter for a Command Line Tool. More details: <https://www.commonwl.org/v1.0/CommandLineTool.html#CommandOutput>

Usage

```
OutputParam(id = "output", label = character(), doc = character(),
  type = "stdout", format = character(),
  secondaryFiles = character(), streamable = logical(),
  glob = character(), loadContents = logical(),
  outputEval = character(), outputSource = character())
```

Arguments

<code>id</code>	The unique identifier for this parameter object.
<code>label</code>	A short, human-readable label of this object.
<code>doc</code>	A documentation string for this object, or an array of strings which should be concatenated.
<code>type</code>	Specify valid types of data that may be assigned to this parameter.
<code>format</code>	Only valid when type: File or is an array of items: File. This is the file format that will be assigned to the output File object.
<code>secondaryFiles</code>	Provides a pattern or expression specifying files or directories. Only valid when type: File or is an array of items: File.
<code>streamable</code>	A value of true indicates that the file is read or written sequentially without seeking. Only valid when type: File or is an array of items: File.
<code>glob</code>	Pattern to find files relative to the output directory.
<code>loadContents</code>	Read text from globbed file.
<code>outputEval</code>	Evaluate an expression to generate the output value.
<code>outputSource</code>	Specifies one or more workflow parameters that supply the value of to the output parameter.

Value

An object of class ‘OutputParam’.

Examples

```
o1 <- OutputParam(id = "file", type = "File", glob = "*.txt")
```

OutputParamList-class *OutputParamList***Description**

`OutputParamList`
`OutputParamList #'` A list of `InputParam`
`outputs` The outputs of a `cwlParam` object

Usage

```
OutputParamList(out = OutputParam(), ...)  

outputs(cwl)
```

Arguments

<code>out</code>	The default stdout parameter.
<code>...</code>	The <code>InputParam</code> objects.
<code>cwl</code>	A <code>cwlParam</code> object

Value

An object of class ‘OutputParamList’.
 outputs: A list of ‘OutputParam’.

Examples

```
o1 <- OutputParam(id = "file", type = "File", glob = "*.txt")
OutputParamList(o1)
input1 <- InputParam(id = "sth")
echo <- cwlParam(baseCommand = "echo", inputs = InputParamList(input1))
outputs(echo)
```

plotCWL

*plotCWL***Description**

Function to plot cwlStepParam object.

Usage

```
plotCWL(cwl, ...)
```

Arguments

cwl	A cwlStepParam object to plot
...	other parameters from ‘mermaid’ function

Value

A mermaid workflow plot.

Examples

```
input1 <- InputParam(id = "sth")
echo1 <- cwlParam(baseCommand = "echo",
                   inputs = InputParamList(input1))
input2 <- InputParam(id = "sthout", type = "File")
echo2 <- cwlParam(baseCommand = "echo",
                   inputs = InputParamList(input2),
                   stdout = "out.txt")
i1 <- InputParam(id = "sth")
o1 <- OutputParam(id = "out", type = "File", outputSource = "echo2/output")
wf <- cwlStepParam(inputs = InputParamList(i1),
                    outputs = OutputParamList(o1))
s1 <- Step(id = "echo1", run = echo1, In = list(sth = "sth"))
s2 <- Step(id = "echo2", run = echo2, In = list(sthout = "echo1/output"))
wf <- wf + s1 + s2
plotCWL(wf)
```

Rcwl

Rcwl

Description

An R package to wrap command line tools and build pipelines with Common Workflow Language.

See Also

[cwlParam](#)

[runCWL](#)

readCWL

Read CWL Function to read CWL command or workflow files.

Description

Read CWL Function to read CWL command or workflow files.

Usage

`readCWL(cwlfile)`

Arguments

`cwlfile` The cwl file to read.

Value

A object of class ‘cwlParam’ or ‘cwlStepParam’.

Examples

```
input1 <- InputParam(id = "sth")
echo <- cwlParam(baseCommand = "echo",
                  inputs = InputParamList(input1))
tf <- tempfile()
writeCWL(echo, tf)
readCWL(paste0(tf, ".cwl"))
```

`runcWL``run cwlParam`

Description

Execute a `cwlParam` object with assigned inputs.

Usage

```
runCWL(cwl, prefix = tempfile(), cwlRunner = "cwltool",
       cwlTemp = FALSE, outdir = ".", Args = character(), stdout = TRUE,
       stderr = TRUE, docker = TRUE, ...)
```

Arguments

<code>cwl</code>	A ‘ <code>cwlParam</code> ‘ or ‘ <code>cwlStepParam</code> ‘ object.
<code>prefix</code>	The prefix of ‘ <code>cwl</code> ‘ and ‘ <code>yml</code> ‘ file to write.
<code>cwlRunner</code>	The path to the ‘ <code>cwltool</code> ‘ or ‘ <code>cwl-runner</code> ‘. If not exists, the <code>cwltool</code> package will be installed by ‘ <code>reticulate</code> ‘.
<code>cwlTemp</code>	Whether to keep temporary files. If true, all temporary files will be kept in a “temp” folder of current output directory.
<code>outdir</code>	Output directory, default current directory.
<code>Args</code>	The arguments for ‘ <code>cwltool</code> ‘ or ‘ <code>cwl-runner</code> ‘. For example, “ <code>-debug</code> ” can work with ‘ <code>cwltool</code> ‘ to show debug information.
<code>stdout</code>	standard output from ‘ <code>system2</code> ‘.
<code>stderr</code>	standard error from ‘ <code>system2</code> ‘. By setting it to “”, the detailed running logs will return directly.
<code>docker</code>	Whether to use docker.
...	The other options from ‘ <code>writeCWL</code> ‘ and ‘ <code>system2</code> ‘.

Value

A list of outputs from tools and logs from `cwltool`.

Examples

```
input1 <- InputParam(id = "sth")
echo <- cwlParam(baseCommand = "echo",
                  inputs = InputParamList(input1))
echo$sth <- "Hello World!"
## res <- runCWL(echo)
```

runCWLBatch	<i>run CWL with batchtools</i>
-------------	--------------------------------

Description

run CWL with batchtools

Usage

```
runCWLBatch(cwl, outdir = getwd(), inputList, paramList = list(),
BPPARAM = BatchtoolsParam(workers = lengths(inputList)[1]), ...)
```

Arguments

cwl	A ‘cwlParam’ or ‘cwlStepParam’ object.
outdir	Directory to output results
inputList	An input list to run in parallel. The list names must be in the inputs of cwl. Jobs will be submitted in parallel for each element in the list. The output directory of each job will be made using the name of each element under the ‘outdir’.
paramList	A parameter list for the cwl. The list names must be in the inputs of cwl.
BPPARAM	The options for ‘BiocParallelParam’.
...	The options from runCWL.

Value

Results from computing nodes and logs from cwltool.

runs	<i>runs</i>
------	-------------

Description

The function to access all runs of a cwlStepParam object

Usage

```
runs(object)
```

Arguments

object	A cwlStepParam object.
--------	------------------------

Value

cwlParam objects or paths of CWL file.

Examples

```
s1 <- cwlStepParam()
runs(s1)
```

<code>short</code>	<i>short</i>
--------------------	--------------

Description

The function to show short summary of cwlParam or cwlStepParam

Usage

```
short(object)
```

Arguments

<code>object</code>	An cwlParam or cwlStepParam object
---------------------	------------------------------------

Value

A short summary of an object of cwlParam or cwlStepParam.

Examples

```
s1 <- cwlStepParam()
short(s1)
```

<code>Step</code>	<i>Step function</i>
-------------------	----------------------

Description

Function to assign value to ‘stepParam‘ object.

Usage

```
Step(id, run = cwlParam(), In = list(), scatter = character(),
     scatterMethod = character())
```

Arguments

<code>id</code>	The id of ‘stepParam‘ object.
<code>run</code>	A ‘cwlParam‘ object for command tool, or path to a CWL file.
<code>In</code>	one or two layers of list.
<code>scatter</code>	character or a list. The inputs to be scattered.
<code>scatterMethod</code>	required if scatter is an array of more than one element. It can be one of "dot-product", "nested_crossproduct" and "flat_crossproduct". Details: https://www.commonwl.org/v1.0/

Value

An object of ‘stepParam‘.

See Also

[cwlStepParam](#)

stepInParam-class *stepInParam*

Description

The input parameter of a workflow step. More details: <https://www.commonwl.org/v1.0/Workflow.html#WorkflowStepInput>

Usage

```
stepInParam(id, source = character(), linkMerge = character(),
            default = character(), valueFrom = character())
```

Arguments

id	A unique identifier for this workflow input parameter.
source	Specifies one or more workflow parameters that will provide input to the underlying step parameter.
linkMerge	The method to use to merge multiple inbound links into a single array.
default	The default value for this parameter to use if either there is no source field, or the value produced by the source is null.
valueFrom	value from string or expression.

Value

An object of class ‘stepInParam’.

Examples

```
s1 <- stepInParam(id = "s1")
```

stepInParamList-class *stepInParamList*

Description

```
stepInParamList  
stepInParamList
```

Usage

```
stepInParamList(...)
```

Arguments

... A list of ‘stepInParam’ objects.

Value

An object of class ‘stepInParamList’.

Examples

```
s1 <- stepInParam(id = "s1")
stepInParamList(s1)
```

stepParam-class

*stepParam***Description**

A workflow step parameters. More details: <https://www.commonwl.org/v1.0/Workflow.html#WorkflowStep>

Usage

```
stepParam(id, run = cwlParam(), In = stepInParamList(), Out = list(),
scatter = character(), scatterMethod = character())
```

Arguments

<code>id</code>	The unique identifier for this workflow step.
<code>run</code>	A ‘cwlParam‘ object or the path of a cwl file.
<code>In</code>	A ‘stepInParamList‘.
<code>Out</code>	A list of outputs
<code>scatter</code>	character or a list. The inputs to be scattered.
<code>scatterMethod</code>	required if scatter is an array of more than one element. It can be one of "dot-product", "nested_crossproduct" and "flat_crossproduct". Details: https://www.commonwl.org/v1.0/

Value

An object of class ‘stepParam‘.

Examples

```
s1 <- stepParam(id = "s1")
```

stepParamList-class

*stepParamList***Description**

```
stepParamList
stepParamList
```

Usage

```
stepParamList(...)
```

Arguments

<code>...</code>	A list of ‘stepParam‘.
------------------	------------------------

Value

An object of class ‘stepParamList’.

Examples

```
s1 <- stepParam(id = "s1")
stepParamList(s1)
```

steps

Steps

Description

Function to extract step slots

Usage

```
steps(cwl)
steps(cwl) <- value
```

Arguments

cwl	A cwlStepParam object.
value	A list of steps.

Value

steps: A list of stepParam objects.

See Also

[cwlStepParam](#)

writeCWL

Write CWL

Description

write ‘cwlParam’ to cwl and yml.

Usage

```
writeCWL(cwl, prefix, docker = TRUE, ...)
```

Arguments

cwl	A ‘cwlParam’ or ‘cwlStepParam’ object.
prefix	The prefix of ‘cwl’ and ‘yml’ file to write.
docker	Whether to use docker.
...	Other options from ‘yaml::write_yaml’.

Value

A CWL file and A YML file.

Examples

```
input1 <- InputParam(id = "sth")
echo <- cwlParam(baseCommand = "echo",
                  inputs = InputParamList(input1))
tf <- tempfile()
writeCWL(echo, tf)
```

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