# Package 'pysd2r'

October 14, 2022

Title API to 'Python' Library 'pysd'

Version 0.1.0
<b>Description</b> Using the R package 'reticulate', this package creates an interface to the 'pysd' toolset. The package provides an R interface to a number of 'pysd' functions, and can read files in 'Vensim' 'mdl' format, and 'xmile' format. The resulting simulations are returned as a 'tibble', and from that the results can be processed using 'dplyr' and 'ggplot2'. The package has been tested using 'python3'.
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<b>Depends</b> R (>= $3.3$ )
Encoding UTF-8
LazyData true
Imports knitr, reticulate, tibble
Suggests dplyr, ggplot2, testthat
RoxygenNote 6.1.0
VignetteBuilder knitr
<b>SystemRequirements</b> 'python3' needs to built for the same architecture R is built for (32 or 64 bit).
NeedsCompilation no
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get\_doc

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get\_doc

Formats a table of variable names

### Description

```
get_doc() Get mode variable names
```

### Usage

```
get_doc(o)
```

#### **Arguments**

0

is the ipysd S3 object

#### Value

tibble

```
## Not run:
target <- system.file("models/vensim", "Population.mdl", package = "pysd2r")
py <- pysd_connect()
py <- read_vensim(py, target)
mdoc <- get_doc(py)
## End(Not run)</pre>
```

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get\_final\_time

Gets the final time from the model

### **Description**

get\_timestep uses pysd to fetch the time step from the model

#### Usage

```
get_final_time(o)
```

#### **Arguments**

0

is the ipysd S3 object

#### **Details**

As it's a generic function, this call is dispatched to set\_component.isdpy

### Value

The finaltime

### **Examples**

```
## Not run:
target <- system.file("models/vensim", "Population.mdl", package = "pysd2r")
py <- pysd_connect()
py <- read_vensim(py, target)
final_time <- get_final_time(py)
## End(Not run)</pre>
```

get\_initial\_time

Gets the initial time from the model

### **Description**

```
\ensuremath{\mbox{get\_initial\_time}} uses pysd to fetch the time step from the model
```

### Usage

```
get_initial_time(o)
```

### **Arguments**

0

is the ipysd S3 object

get\_python\_info

### **Details**

As it's a generic function, this call is dispatched to set\_component.isdpy

### Value

The initial time

### **Examples**

```
## Not run:
target <- system.file("models/vensim", "Population.mdl", package = "pysd2r")
py <- pysd_connect()
py <- read_vensim(py, target)
initial_time <- get_initial_time(py)
## End(Not run)</pre>
```

get\_python\_info

Gets the current python configuration for reticulate

### **Description**

get\_python\_info returns information on what version of python is being used with reticulate

#### Usage

```
get_python_info()
```

### Value

python information

```
## Not run:
get_python_info()
## End(Not run)
```

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get\_timestep

Gets the time step (DT) from the model

### Description

get\_timestep uses pysd to fetch the time step from the model

#### Usage

```
get_timestep(o)
```

#### **Arguments**

0

is the ipysd S3 object

#### **Details**

As it's a generic function, this call is dispatched to set\_component.isdpy

#### Value

The simulation time step

#### **Examples**

```
## Not run:
target <- system.file("models/vensim", "Population.mdl", package = "pysd2r")
py <- pysd_connect()
py <- read_vensim(py, target)
time_step <- get_timestep(py)
## End(Not run)</pre>
```

pysd\_connect

Creates an object to facilitate interaction with pysd

### Description

pysd\_connect returns a ipysd object to the calling program. This object will contain a link variable to pysd and will subsequently store a reference to the simulation model in pysd.

### Usage

```
pysd_connect()
```

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### **Details**

### Link to pysd

The result is used as a parameter for read\_vensim() & read\_xmile() functions

#### Value

An S3 object of class ipysd

### **Examples**

```
## Not run:
py pysd_connect()
## End(Not run)
```

read\_vensim

Loads a Vensim simulation file (mdl)

### Description

read\_vensim() calls pysd.read\_vensim() and stores the object for further use. This is a key object, as it relates to a model and it can support a number of functions (e.g. model run, parameter changes)

### Usage

```
read_vensim(o, file)
```

### Arguments

o is the ipysd S3 object

file is the filename and path for the Vensim mdl file that needs to be simulated

### **Details**

The result is used as a parameter for simulation calls.

As it's a generic function, this call is dispatched to read\_vensim.isdpy

#### Value

An S3 object of class ipysd that will contain a reference to the model

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#### **Examples**

```
## Not run:
target <- system.file("models/vensim", "Population.mdl", package = "pysd2r")
py <- pysd_connect()
read_vensim(py, target)
## End(Not run)</pre>
```

read\_xmile

Loads a XMILE simulation file (.xmile)

### Description

read\_xmile() calls pysd.read\_xmile() and stores the object for further use. This is a key object, as it relates to a model and it can support a number of functions (e.g. model run, parameter changes)

### Usage

```
read_xmile(o, file)
```

#### **Arguments**

o is the ipysd S3 object

file is the filename and path for the Vensim mdl file that needs to be simulated

#### **Details**

The result is used as a parameter for simulation calls.

As it's a generic function, this call is dispatched to read\_xmile.isdpy

### Value

An S3 object of class ipysd that will contain a reference to the model

```
## Not run:
target <- system.file("models/xmile", "Population.xmile", package = "pysd2r")
py <- pysd_connect()
read_xmile(py, target)
## End(Not run)</pre>
```

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reload\_model

Reloads the model from original mdl file

### **Description**

```
reload_model() Reloads the model
```

### Usage

```
reload_model(o)
```

### **Arguments**

0

is the ipysd S3 object

#### Value

ipysd object

### **Examples**

```
## Not run:
target <- system.file("models/vensim", "Population.mdl", package = "pysd2r")
py <- pysd_connect()
py <- read_vensim(py, target)
set_time_values(py,0,10,0.5)
py<-reload_model(py)
## End(Not run)</pre>
```

run\_model

Runs a simulation model

### **Description**

run\_model() calls run in pysd and returns all the simulation output in tidy data format (tibble)

### Usage

```
run_model(o)
```

### **Arguments**

0

is the ipysd S3 object

### **Details**

As it's a generic function, this call is dispatched to run\_model.isdpy

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### Value

tibble containing the simulation results

#### **Examples**

```
## Not run:
target <- system.file("models/vensim", "Population.mdl", package = "pysd2r")
py <- pysd_connect()
py <- read_vensim(py, target)
results <- run_model(py)
## End(Not run)</pre>
```

set\_components

Changes a model parameter

### **Description**

set\_components() calls .set\_components() and changes a resulting parameter in the model

### Usage

```
set_components(o, vals)
```

### Arguments

o is the ipysd S3 object

vals contains a list with the parameter and value to be changed

#### **Details**

As it's a generic function, this call is dispatched to set\_component.isdpy

```
## Not run:
target <- system.file("models/vensim", "Population.mdl", package = "pysd2r")
py <- pysd_connect()
py <- read_vensim(py, target)
results <- run_model(py)
1 <- list("Growth Fraction"=0.02)
set_components(py,1)
out2 <- run_model(py)
## End(Not run)</pre>
```

set\_time\_values

set\_time\_values

Sets the initial time, final time, and timestep

### Description

```
set_time_values1() sets the simulation times and DT
```

### Usage

```
set_time_values(o, init, final, DT)
```

### Arguments

```
o is the ipysd S3 object init is the initial time final is the final time DT is the time step
```

```
## Not run:
target <- system.file("models/vensim", "Population.mdl", package = "pysd2r")
py <- pysd_connect()
py <- read_vensim(py, target)
set_time_values(py,0,10,0.5)
## End(Not run)</pre>
```

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