

# Package ‘pysd2r’

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**Title** API to 'Python' Library 'pysd'

**Version** 0.1.0

**Description** 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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**Depends** R (>= 3.3)

**Encoding** UTF-8

**LazyData** true

**Imports** knitr, reticulate, tibble

**Suggests** dplyr, ggplot2, testthat

**RoxygenNote** 6.1.0

**VignetteBuilder** knitr

**SystemRequirements** 'python3' needs to be built for the same architecture  
R is built for (32 or 64 bit).

**NeedsCompilation** no

**Author** Jim Duggan [aut, cre]

**Maintainer** Jim Duggan <jim.duggan@nuigalway.ie>

**Repository** CRAN

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## Contents

get_doc . . . . .	2
get_final_time . . . . .	3
get_initial_time . . . . .	3
get_python_info . . . . .	4

get_timestep . . . . .	5
pysd_connect . . . . .	5
read_vensim . . . . .	6
read_xmile . . . . .	7
reload_model . . . . .	8
run_model . . . . .	8
set_components . . . . .	9
set_time_values . . . . .	10

<b>Index</b>	<b>11</b>
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get_doc	<i>Formats a table of variable names</i>
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## Description

get\_doc() Get mode variable names

## Usage

```
get_doc(o)
```

## Arguments

o is the ipysd S3 object

## Value

tibble

## Examples

```
## 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)
```

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get_final_time	<i>Gets the final time from the model</i>
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**Description**

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

**Usage**

```
get_final_time(o)
```

**Arguments**

o 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)
```

---

get_initial_time	<i>Gets the initial time from the model</i>
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---

**Description**

get\_initial\_time uses pysd to fetch the time step from the model

**Usage**

```
get_initial_time(o)
```

**Arguments**

o is the ipysd S3 object

**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)
```

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`get_python_info`*Gets the current python configuration for reticulate*

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**Description**

`get_python_info` returns information on what version of python is being used with reticulate

**Usage**

```
get_python_info()
```

**Value**

python information

**Examples**

```
## Not run:
get_python_info()

## End(Not run)
```

---

get_timestep	<i>Gets the time step (DT) from the model</i>
--------------	---

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**Description**

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

**Usage**

```
get_timestep(o)
```

**Arguments**

o 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)
```

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pysd_connect	<i>Creates an object to facilitate interaction with pysd</i>
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**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()
```

**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)
```

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read_vensim	<i>Loads a Vensim simulation file (mdl)</i>
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**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

## Examples

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

## End(Not run)
```

---

read_xmile	<i>Loads a XMILE simulation file (.xmile)</i>
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## 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

## Examples

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

## End(Not run)
```

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reload_model	<i>Reloads the model from original mdl file</i>
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**Description**

reload\_model() Reloads the model

**Usage**

```
reload_model(o)
```

**Arguments**

o 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)
```

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run_model	<i>Runs a simulation model</i>
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**Description**

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

**Usage**

```
run_model(o)
```

**Arguments**

o is the ipysd S3 object

**Details**

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

**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)
```

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set_components	<i>Changes a model parameter</i>
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**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

**Examples**

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

## End(Not run)
```

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set_time_values	<i>Sets the initial time, final time, and timestep</i>
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**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

**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)  
  
## End(Not run)
```

# Index

`get_doc`, [2](#)  
`get_final_time`, [3](#)  
`get_initial_time`, [3](#)  
`get_python_info`, [4](#)  
`get_timestep`, [5](#)  
  
`pysd_connect`, [5](#)  
  
`read_vensim`, [6](#)  
`read_xmile`, [7](#)  
`reload_model`, [8](#)  
`run_model`, [8](#)  
  
`set_components`, [9](#)  
`set_time_values`, [10](#)