

# Package ‘CaPO4Sim’

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**Type** Package

**Title** A Virtual Patient Simulator in the Context of Calcium and Phosphate Homeostasis

**Version** 0.2.1

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**Description** Explore calcium (Ca) and phosphate (Pi) homeostasis with two novel 'Shiny' apps, building upon a previously published mathematical model written in C, to ensure efficient computations. The underlying model is accessible here <<https://pubmed.ncbi.nlm.nih.gov/28747359/>>.

The first application explores the fundamentals of Ca-Pi homeostasis, while the second provides interactive case studies for in-depth exploration of the topic, thereby seeking to foster student engagement and an integrative understanding of Ca-Pi regulation.

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arrow_lighting	<i>Highlight arrows for steady state events</i>
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## Description

Use inside in the [networkCaPO4](#). Nothing is returned except that the network is updated via [vis-NetworkProxy](#).

## Usage

```
arrow_lighting(edges, simulation, counter, session)
```

## Arguments

edges	A dataframe of edges provided by <a href="#">generate_edges</a> .
simulation	Which disease is currently selected. See <a href="#">extract_running_sim</a> .
counter	To determine which notification to display. We expect a counter returned by the <a href="#">networkCaPO4</a> module.
session	Session object.

---

CaPO4Sim	<i>CaPO4Sim</i>
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---

## Description

Explore calcium (Ca) and phosphate (Pi) homeostasis with two novel 'Shiny' apps, building upon on a previously published mathematical model written in C, to ensure efficient computations. The underlying model is accessible here <<https://www.ncbi.nlm.nih.gov/pubmed/28747359>>. The first application explores the fundamentals of Ca-Pi homeostasis, while the second provides interactive case studies for in-depth exploration of the topic, thereby seeking to foster student engagement and an integrative understanding of Ca-Pi regulation. These applications are hosted at <<https://rinterface.com/AppsPhysiol.html>>.

## Author(s)

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- Diane de Zélicourt [copyright holder]
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- Kidney NCCR.CH [funder]
- The Interface Group (Hosting Group) [copyright holder]
- RinteRface (R/HTML Templates) [copyright holder]

diseaseCheckBox

*Create a checkbox for diseaseSelectUi***Description**

Create a [prettyCheckbox](#).

**Usage**

```
diseaseCheckBox(inputId, label)
```

**Arguments**

inputId	Checkbox Input id.
label	Checkbox label.

diseaseSelect

*Create a disease selector server logic***Description**

Only returns inputs associated with php1, hypopara, hypoD3

**Usage**

```
diseaseSelect(input, output, session)
```

**Arguments**

input	Shiny inputs
output	Shiny Outputs
session	Session object.

---

**diseaseSelectUi***Create a disease selector UI module*

---

**Description**

Contains php1, hypopara, hypoD3

**Usage**

```
diseaseSelectUi(id)
```

**Arguments**

<b>id</b>	module id.
-----------	------------

---

**extract\_running\_sim***Extract the current running simulation*

---

**Description**

Simulations are currently php1, hypoD3 and hypopara. Takes diseases as input given by the [diseaseSelect](#) module.

**Usage**

```
extract_running_sim(diseases)
```

**Arguments**

<b>diseases</b>	Shiny input disease selector. See <a href="#">diseaseSelect</a> .
-----------------	---

---

**fullScreen***Create a fullScreen server logic*

---

**Description**

Nothing is contained inside for now...

**Usage**

```
fullScreen(input, output, session)
```

**Arguments**

<b>input</b>	Shiny inputs
<b>output</b>	Shiny Outputs
<b>session</b>	Session object.

---

<code>fullScreenUI</code>	<i>Create a fullScreen UI module</i>
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---

### Description

Trigger a fullScreen mode. Based on <https://stackoverflow.com/questions/42371164/how-to-run-r-shiny-app-in-full-sized-window>

### Usage

```
fullScreenUI(id)
```

### Arguments

<code>id</code>	module id.
-----------------	------------

---



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<code>generate_edges</code>	<i>CaPO4 Edges Generator</i>
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---

### Description

Generate edges for the CaPO4 network

### Usage

```
generate_edges(
  components,
  organs,
  regulations,
  diseases,
  organs_edges_size,
  hormones_edges_size
)
```

### Arguments

<code>components</code>	Shiny input CaPO4 component selector. See <a href="#">networkOptions</a> .
<code>organs</code>	Shiny input to toggle organs display. See <a href="#">networkOptions</a> .
<code>regulations</code>	Shiny input to toggle hormone display. See <a href="#">networkOptions</a> .
<code>diseases</code>	Shiny input disease selector. See <a href="#">diseaseSelect</a> .
<code>organs_edges_size</code>	Shiny input for organs edges size. See <a href="#">networkOptions</a> .
<code>hormones_edges_size</code>	Shiny input for hormones edges size. See <a href="#">networkOptions</a> .

---

generate\_network      *CaPO4 Network Generator*

---

### Description

Create a CaPO4 network taking nodes and edges as inputs

### Usage

```
generate_network(nodes, edges, usephysics = FALSE, isMobile)
```

### Arguments

nodes	A dataframe of nodes provided by <a href="#">generate_nodes</a> .
edges	A dataframe of edges provided by <a href="#">generate_edges</a> .
usephysics	Whether to use physic. FALSE by default. A visNetwork API parameter.
isMobile	Shiny input checking if the app is running on a cellphone/tablet.

---

generate\_nodes      *CaPO4 Nodes Generator*

---

### Description

Generate nodes for the CaPO4 network

### Usage

```
generate_nodes(  
  components,  
  organs,  
  regulations,  
  background,  
  diseases,  
  organs_nodes_size,  
  hormones_nodes_size  
)
```

### Arguments

components	Shiny input CaPO4 component selector. See <a href="#">networkOptions</a> .
organs	Shiny input to toggle organs display. See <a href="#">networkOptions</a> .
regulations	Shiny input to toggle hormone display. See <a href="#">networkOptions</a> .
background	Shiny input background selector. See <a href="#">networkOptions</a> .

**diseases** Shiny input disease selector. See [diseaseSelect](#).  
**organs\_nodes\_size** Shiny input for organs node size. See [networkOptions](#).  
**hormones\_nodes\_size** Shiny input for hormones node size. See [networkOptions..](#)

**generate\_notification** *Notifications Generator for CaPO4 animations*

### Description

Generate sequential notification as a function of the selected diseases. All notifications are in the notifications.R file in the inst/entry\_level app folder. Used in the [infos](#) module.

### Usage

```
generate_notification(simulation, counter, allowed)
```

### Arguments

**simulation** Which disease is currently selected. See [extract\\_running\\_sim](#).  
**counter** To determine which notification to display. We expect a counter returned by the [networkCaPO4](#) module.  
**allowed** Whether to allow simulations. Expect logical value. See [infos](#) module.

**generate\_userFields** *Generate user fields*

### Description

Use inside in the [userInfo](#). Function that helps in generating 4 users fields, image, stat1, stat2 and stat3, so as to reinject them in the header userMenu

### Usage

```
generate_userFields(diseases, sliderDisease)
```

### Arguments

**diseases** Shiny input disease selector. See [diseaseSelect](#).  
**sliderDisease** Shiny slider input related to the current disease severity. See [plotBox](#).

---

*glossaryCaP04*      *CaPO4 glossary server module*

---

### Description

Create a CaPO4 glossary

### Usage

```
glossaryCaP04(input, output, session)
```

### Arguments

input	Shiny inputs
output	Shiny Outputs
session	Session object.

---

*glossaryCaP04Ui*      *CaPO4 glossary UI module*

---

### Description

Create a CaPO4 glossary

### Usage

```
glossaryCaP04Ui(id)
```

### Arguments

id	module id.
----	------------

---

`helpCaP04`*Help server module*

---

**Description**

Create the help section

**Usage**

```
helpCaP04(input, output, session)
```

**Arguments**

input	Shiny inputs
output	Shiny Outputs
session	Session object.

---

`helpCaP04Ui`*Help UI module*

---

**Description**

Create a help button

**Usage**

```
helpCaP04Ui(id)
```

**Arguments**

id	module id.
----	------------

---

**infos***Info server module*

---

**Description**

Create modals, alerts, ...

**Usage**

```
infos(input, output, session, diseases, animation_counter, regulations)
```

**Arguments**

input	Shiny inputs
output	Shiny Outputs
session	Session object.
diseases	Shiny input disease selector. See <a href="#">diseaseSelect</a> .
animation_counter	Give the current temporal state of the animation. See <a href="#">networkCaPO4</a> .
regulations	Shiny input to toggle hormone display. See <a href="#">networkOptions</a> .

---

**infosUi***Info UI module*

---

**Description**

Create modals, alerts, ...

**Usage**

```
infosUi(id)
```

**Arguments**

id	module id.
----	------------

**infoSwitch***Create a switch input for [infosUi](#)***Description**

Create a [prettySwitch](#).

**Usage**

```
infoSwitch(inputId, label)
```

**Arguments**

<code>inputId</code>	Checkbox Input id.
<code>label</code>	Checkbox label.

**make\_plot\_hypoD3***Produce plots related to vitamin D3 deficiency (hypoD3)***Description**

Use inside the [plotBox](#) module.

**Usage**

```
make_plot_hypoD3(sliderVal, isMobile)
```

**Arguments**

<code>sliderVal</code>	Shiny slider input related to the current disease severity. See <a href="#">plotBox</a> .
<code>isMobile</code>	Shiny input useful to scale elements based on the device screen size.

**make\_plot\_hypopara***Produce plots related to hypoparathyroidism (hypopara)***Description**

Use inside the [plotBox](#) module.

**Usage**

```
make_plot_hypopara(sliderVal, isMobile)
```

**Arguments**

<code>sliderVal</code>	Shiny slider input related to the current disease severity. See <a href="#">plotBox</a> .
<code>isMobile</code>	Shiny input useful to scale elements based on the device screen size.

---

`make_plot_php1`*Produce plots related to primary hyperparathyroidism (php1)*

---

## Description

Use inside the [plotBox](#) module.

## Usage

```
make_plot_php1(sliderVal, isMobile)
```

## Arguments

<code>sliderVal</code>	Shiny slider input related to the current disease severity. See <a href="#">plotBox</a> .
<code>isMobile</code>	Shiny input useful to scale elements based on the device screen size.

---

`myCarousel`*carousel container*

---

## Description

Creates a carousel. Adapted from shinydashboardplus to also allow control of the carousel animation

## Usage

```
myCarousel(
  ...,
  id,
  indicators = TRUE,
  width = 6,
  .list = NULL,
  data.interval = 5000,
  data.ride = "carousel"
)
```

## Arguments

<code>...</code>	Slot for <a href="#">carouselItem</a>
<code>id</code>	Carousel id. Must be unique.
<code>indicators</code>	Whether to display left and right indicators.
<code>width</code>	Carousel width. 6 by default.
<code>.list</code>	Should you need to pass <a href="#">carouselItem</a> via <a href="#">lapply</a> or similar, put these item here instead of passing them in ...

`data.interval` specify data-interval in ms. 5000ms by default, set to "false" to prevent automated animation of the slides.  
`data.ride` specify data-ride. "carousel" by default.

**networkCaPO4***CaPO4 Network server module***Description**

Create a CaPO4 network

**Usage**

```
networkCaPO4(
  input,
  output,
  session,
  isMobile,
  components,
  organs,
  regulations,
  background,
  diseases,
  organs_nodes_size,
  hormones_nodes_size,
  organs_edges_size,
  hormones_edges_size,
  help
)
```

**Arguments**

<code>input</code>	Shiny inputs
<code>output</code>	Shiny Outputs
<code>session</code>	Session object.
<code>isMobile</code>	Shiny input checking if the app is running on a cellphone/tablet.
<code>components</code>	Shiny input CaPO4 component selector. See <a href="#">networkOptions</a> .
<code>organs</code>	Shiny input to toggle organs display. See <a href="#">networkOptions</a> .
<code>regulations</code>	Shiny input to toggle hormone display. See <a href="#">networkOptions</a> .
<code>background</code>	Shiny input background selector. See <a href="#">networkOptions</a> .
<code>diseases</code>	Shiny input disease selector. See <a href="#">diseaseSelect</a> .
<code>organs_nodes_size</code>	Shiny input for organs node size. See <a href="#">networkOptions</a> .

---

hormones_nodes_size	Shiny input for hormones node size. See <a href="#">networkOptions</a> .
organs_edges_size	Shiny input for organs edges size. See <a href="#">networkOptions</a> .
hormones_edges_size	Shiny input for hormones edges size. See <a href="#">networkOptions</a> .
help	Help input.

---



---

networkCaPO4Ui	<i>CaPO4 Network UI module</i>
----------------	--------------------------------

---

## Description

Create a CaPO4 network

## Usage

```
networkCaPO4Ui(id)
```

## Arguments

id	module id.
----	------------

---

networkOptions	<i>CaPO4 Network Options server module</i>
----------------	--

---

## Description

Create a CaPO4 network options

## Usage

```
networkOptions(input, output, session, mobile)
```

## Arguments

input	Shiny inputs
output	Shiny Outputs
session	Session object.
mobile	Whether we are on cellphone/tablets or not. Slot for input\$ismobile().

---

networkOptionsUi

*CaPO4 Network Options UI module*

---

## Description

Options for the network

## Usage

```
networkOptionsUi(id)
```

## Arguments

id	module id
----	-----------

---

plotBox

*plot box server module*

---

## Description

Create modals, alerts, ...

## Usage

```
plotBox(input, output, session, diseases, help, isMobile)
```

## Arguments

input	Shiny inputs
-------	--------------

output	Shiny Outputs
--------	---------------

session	Session object.
---------	-----------------

diseases	Shiny input disease selector. See <a href="#">diseaseSelect</a> .
----------	---

help	Help input.
------	-------------

isMobile	Shiny input useful to scale elements based on the device screen size.
----------	---

---

**plotBoxUi***plot box UI module*

---

**Description**

Create modals, alerts, ...

**Usage**

```
plotBoxUi(id)
```

**Arguments**

**id** module id.

---

---

**run\_CaP04Sim***Launch the virtual patient simulator*

---

**Description**

Unleash the virtual patient simulator

**Usage**

```
run_CaP04Sim(context = c("introduction", "virtual-patient"))
```

**Arguments**

**context** Choose between c("introduction", "virtual-patient").

**Examples**

```
if (interactive()) {  
  run_CaP04Sim(context = "introduction")  
  run_CaP04Sim(context = "virtual-patient")  
}
```

---

**skinSelect***Dashboard skin selector, server side*

---

**Description**

Select the shinydashboard skin you want

**Usage**

```
skinSelect(input, output, session)
```

**Arguments**

input	Shiny inputs
output	Shiny Outputs
session	Session object.

---

**skinSelectUi***Dashboard skin selector, ui side*

---

**Description**

Select the shinydashboard skin you want

**Usage**

```
skinSelectUi(id)
```

**Arguments**

id	module id.
----	------------

---

**userInfo***CaPO4 user info server module*

---

**Description**

Create a CaPO4 user info card

**Usage**

```
userInfo(input, output, session, diseases, sliderDisease, help)
```

**Arguments**

input	Shiny inputs
output	Shiny Outputs
session	Session object.
diseases	Shiny input disease selector. See <a href="#">diseaseSelect</a> .
sliderDisease	Shiny input disease severity selector. See <a href="#">plotBox</a> .
help	Help input.

---

**userInfoUi***CaPO4 user info UI module*

---

**Description**

Create a CaPO4 user info card

**Usage**

```
userInfoUi(id)
```

**Arguments**

id	module id.
----	------------

---

video	<i>Create a video server logic</i>
-------	------------------------------------

---

**Description**

Nothing is contained inside for now...

**Usage**

```
video(input, output, session)
```

**Arguments**

input	Shiny inputs
output	Shiny Outputs
session	Session object.

---

videoUi	<i>Create a movie UI module</i>
---------	---------------------------------

---

**Description**

Contains php1, hypopara, hypoD3

**Usage**

```
videoUi(id, data)
```

**Arguments**

id	module id.
data	Video data.

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