



**gamar: an R interface to
the GAMA agent-based
simulation platform**

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Population dynamics modeling

Increased level of detail

population-based

differential equations

`bvpSolve`

`dde`

`deSolve`

`diffEQr`

`PBSddesolve`

individual-based

Gillespie algorithms

`adaptivetau`

`GillespieSSA`

agent-based



Agents are

- autonomous
- heterogeneous
- active
- adaptive

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Agent-based modeling platforms

Performance

since 1999



ccl.northwestern.edu/netlogo

since 2007



gama-platform.github.io

since 2000



repast.github.io

Ease of use

NetLogoR
RNetLogo

rrepast

Agent-based modeling platforms

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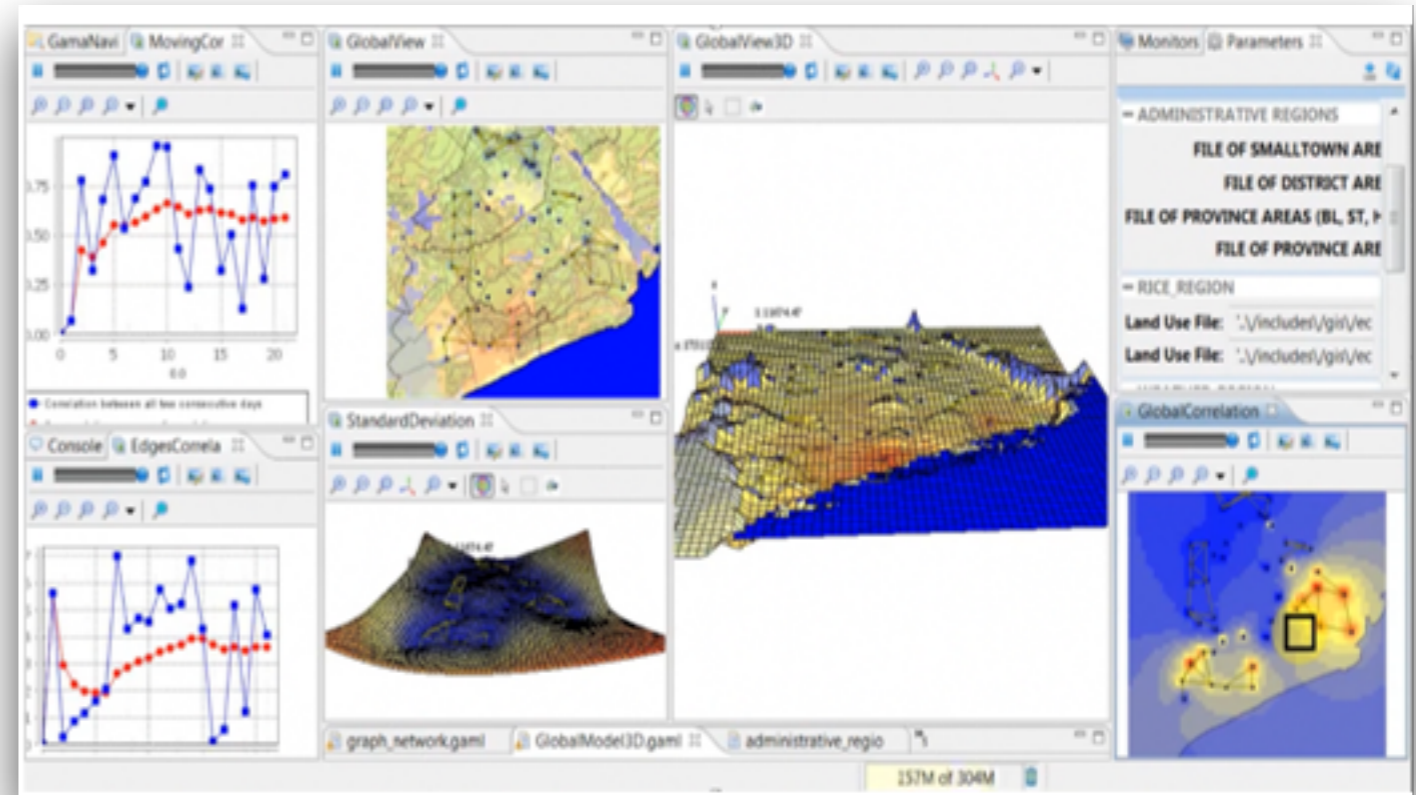
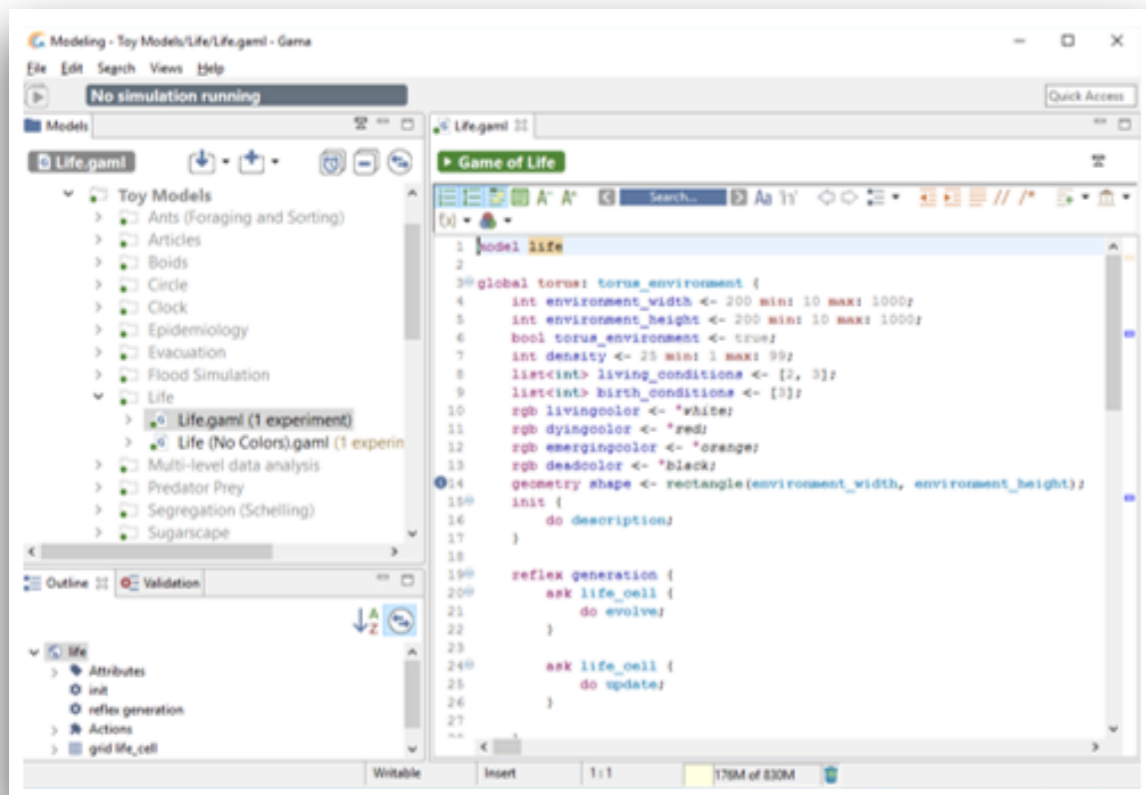
rrepast

The GAMA simulation platform



- a language: GAML
- a user-interface
- a fast and parallelized engine
- integration with GIS data

<https://gama-platform.github.io>

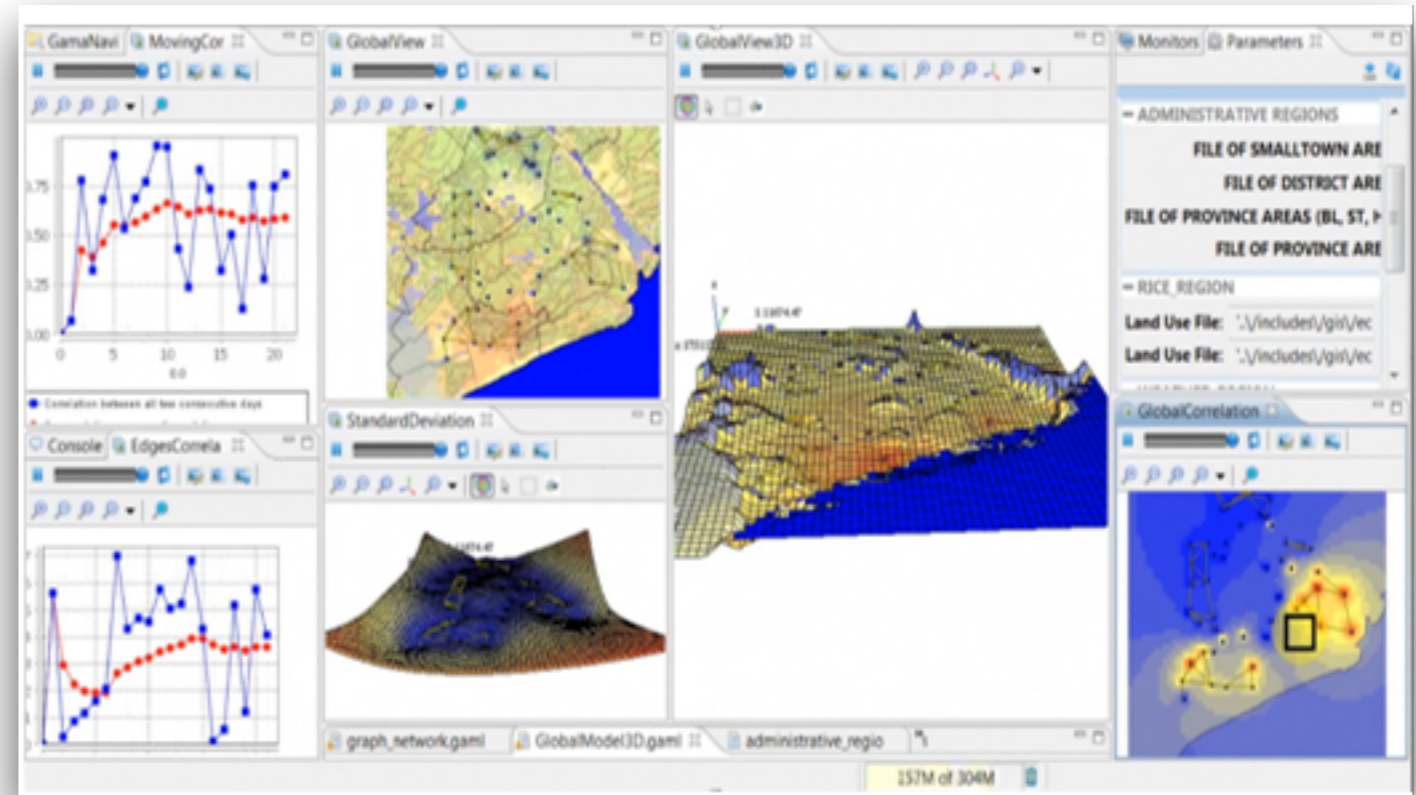
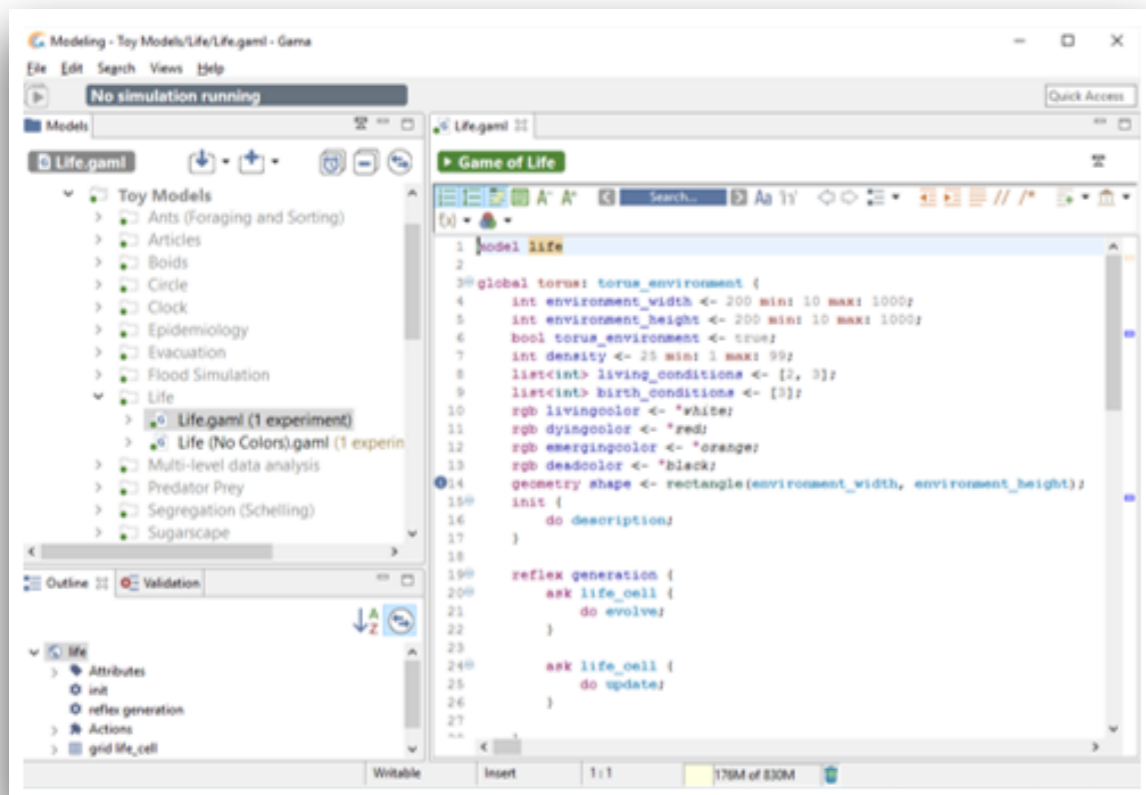


The GAMA simulation platform

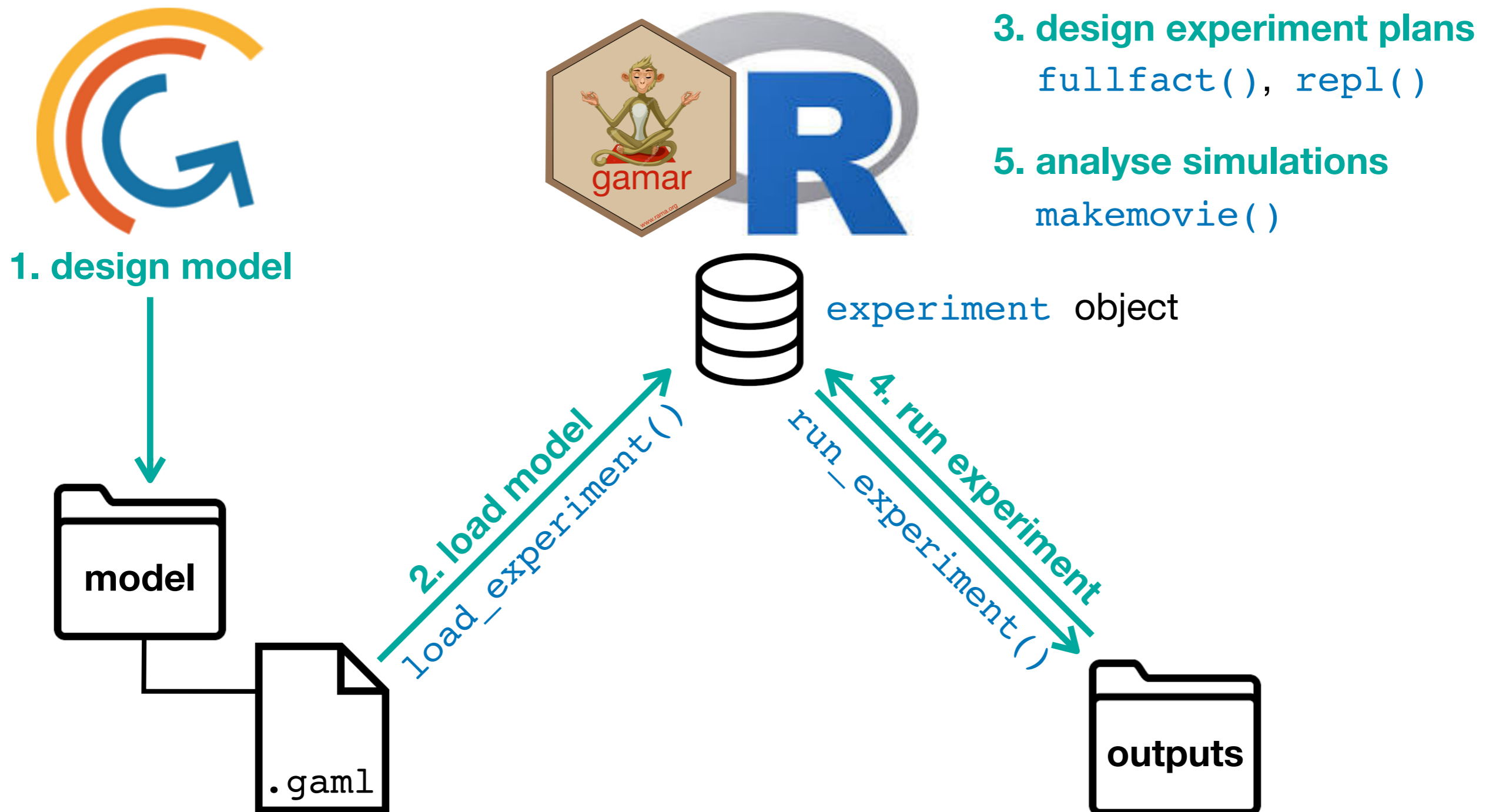


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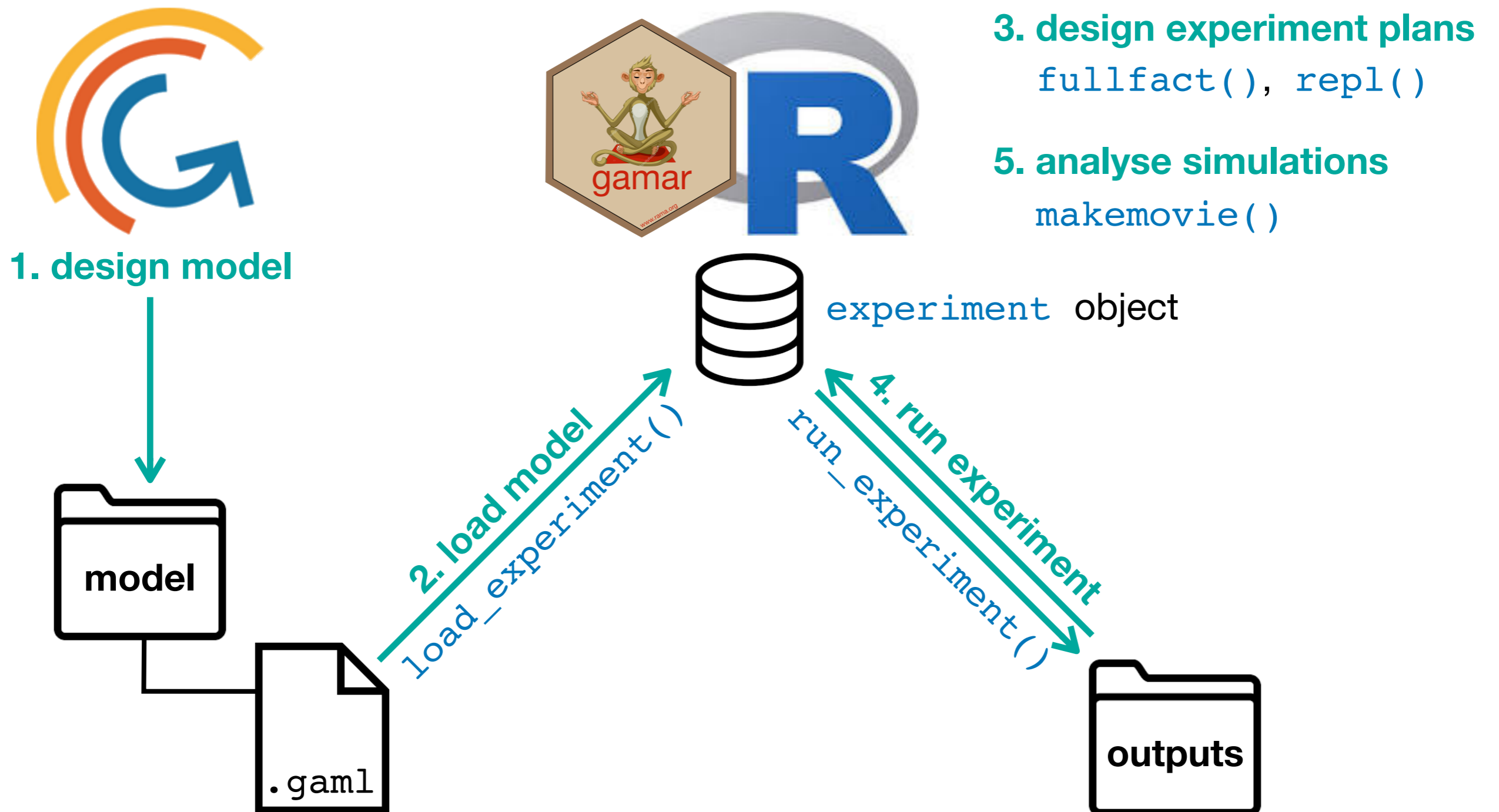
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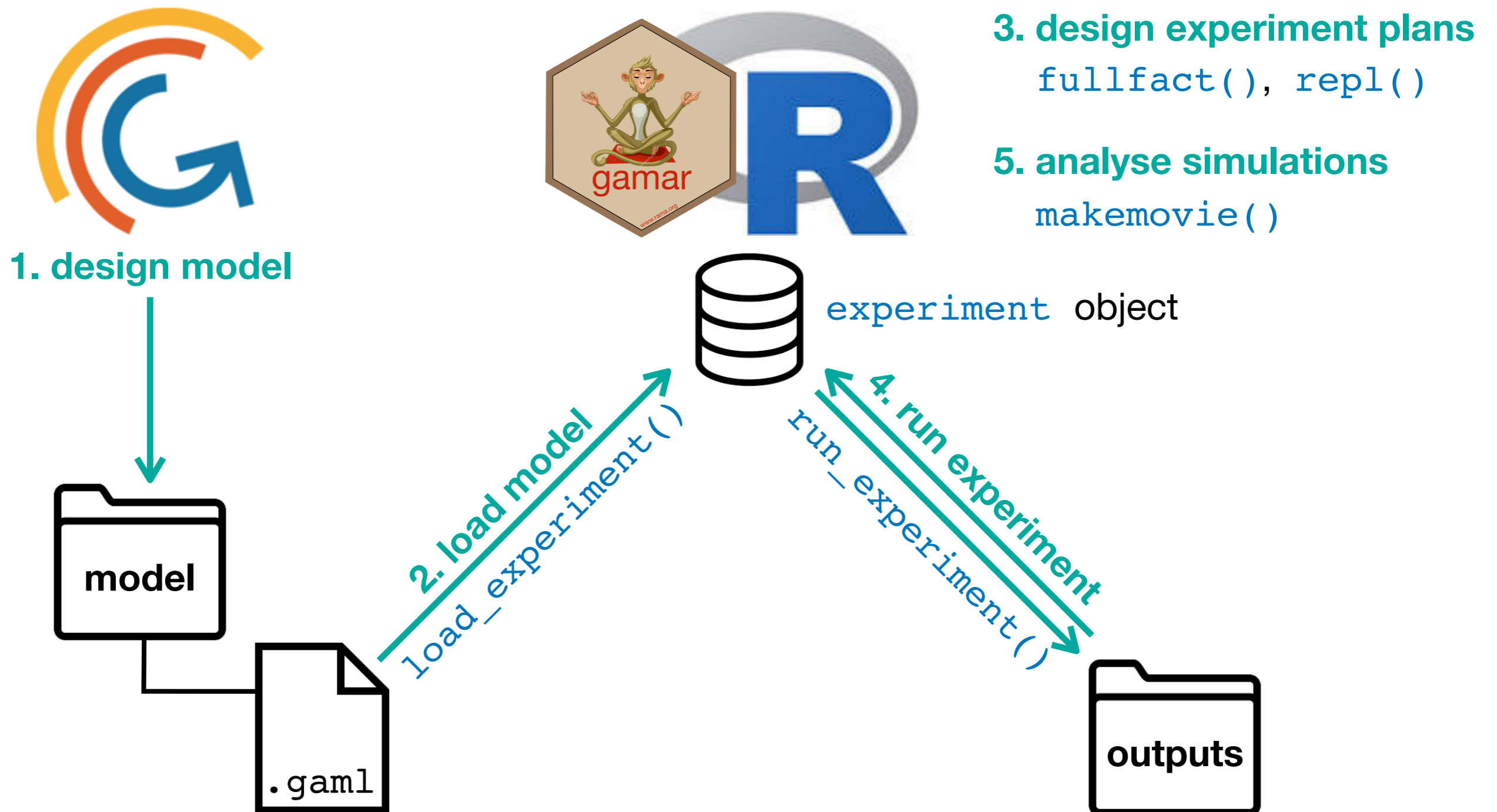
The `gamar` R package



The `gamar` R package



The `gamar` R package



The experiment class

Experiment with 30 simulations of 5 parameters and 3 observed variables

experiment name: sir
input gaml file: /Users/gamar/models/sir/sir.gaml
output directory: /Users/gamar/models/sir/output
model parameters: p_S0, p_I0, p_R0, p_beta, p_gamma
observed variables: r_S, r_I, r_R

Experiment overview:

	p_S0	p_I0	p_R0	p_beta	p_gamma	r_S	r_I	r_R	tmax	seed	output
1	999	1	0	1.5	0.15	1	1	1	1000	1	NA
2	999	1	0	1.5	0.15	1	1	1	1000	1	NA
3	999	1	0	1.5	0.15	1	1	1	1000	1	NA
4	999	1	0	1.5	0.15	1	1	1	1000	1	NA
5	999	1	0	1.5	0.15	1	1	1	1000	1	NA
6	999	1	0	1.5	0.15	1	1	1	1000	1	NA
.
.
.
25	999	1	0	1.5	0.15	1	1	1	1000	1	NA
26	999	1	0	1.5	0.15	1	1	1	1000	1	NA
27	999	1	0	1.5	0.15	1	1	1	1000	1	NA
28	999	1	0	1.5	0.15	1	1	1	1000	1	NA
29	999	1	0	1.5	0.15	1	1	1	1000	1	NA
30	999	1	0	1.5	0.15	1	1	1	1000	1	NA

one column for simulations durations
- always penultimate position
- named **tmax**

one list-column for simulations outputs
- always last position
- named **output**

one column for seeds values
- always penultimate position
- named **seed**

several columns for parameters values
- always first columns
- names start with **p_**

several columns for monitored variables
- just after the parameters columns
- names start with **r_**

The experiment class

Experiment with 30 simulations of 5 parameters and 3 observed variables

experiment name: sir
input gaml file: /Users/gamar/models/sir/sir.gaml
output directory: /Users/gamar/models/sir/output
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Experiment overview:

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1	999	1	0	1.5	0.15	1	1	1	1000	1	<data.frame[1000,3]>
2	999	1	0	1.5	0.15	1	1	1	1000	1	<data.frame[1000,3]>
3	999	1	0	1.5	0.15	1	1	1	1000	1	<data.frame[1000,3]>
4	999	1	0	1.5	0.15	1	1	1	1000	1	<data.frame[1000,3]>
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6	999	1	0	1.5	0.15	1	1	1	1000	1	<data.frame[1000,3]>
.
.
.
25	999	1	0	1.5	0.15	1	1	1	1000	1	<data.frame[1000,3]>
26	999	1	0	1.5	0.15	1	1	1	1000	1	<data.frame[1000,3]>
27	999	1	0	1.5	0.15	1	1	1	1000	1	<data.frame[1000,3]>
28	999	1	0	1.5	0.15	1	1	1	1000	1	<data.frame[1000,3]>
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30	999	1	0	1.5	0.15	1	1	1	1000	1	<data.frame[1000,3]>

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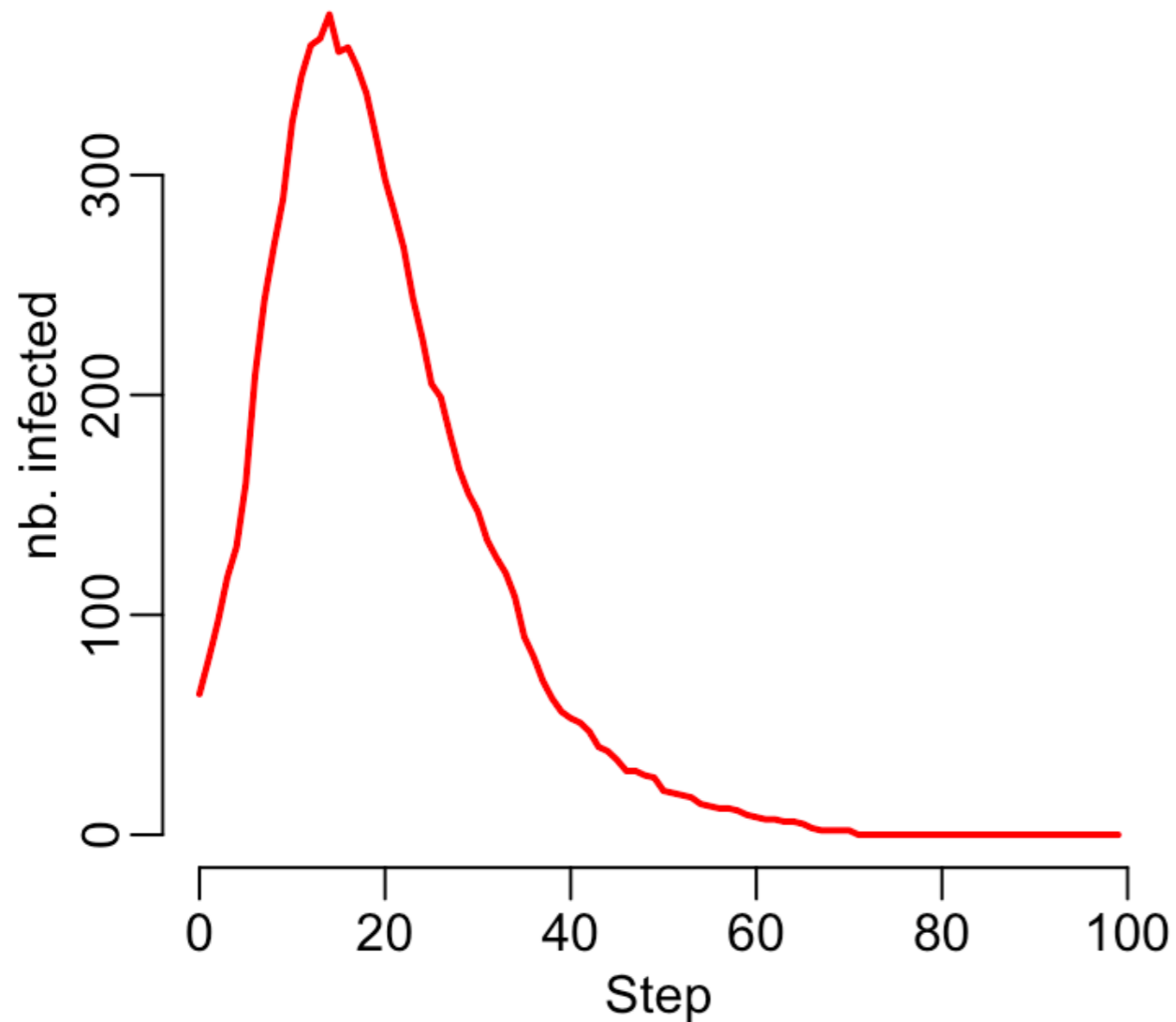
one column for seeds values
- always penultimate position
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several columns for parameters values
- always first columns
- names start with **p_**

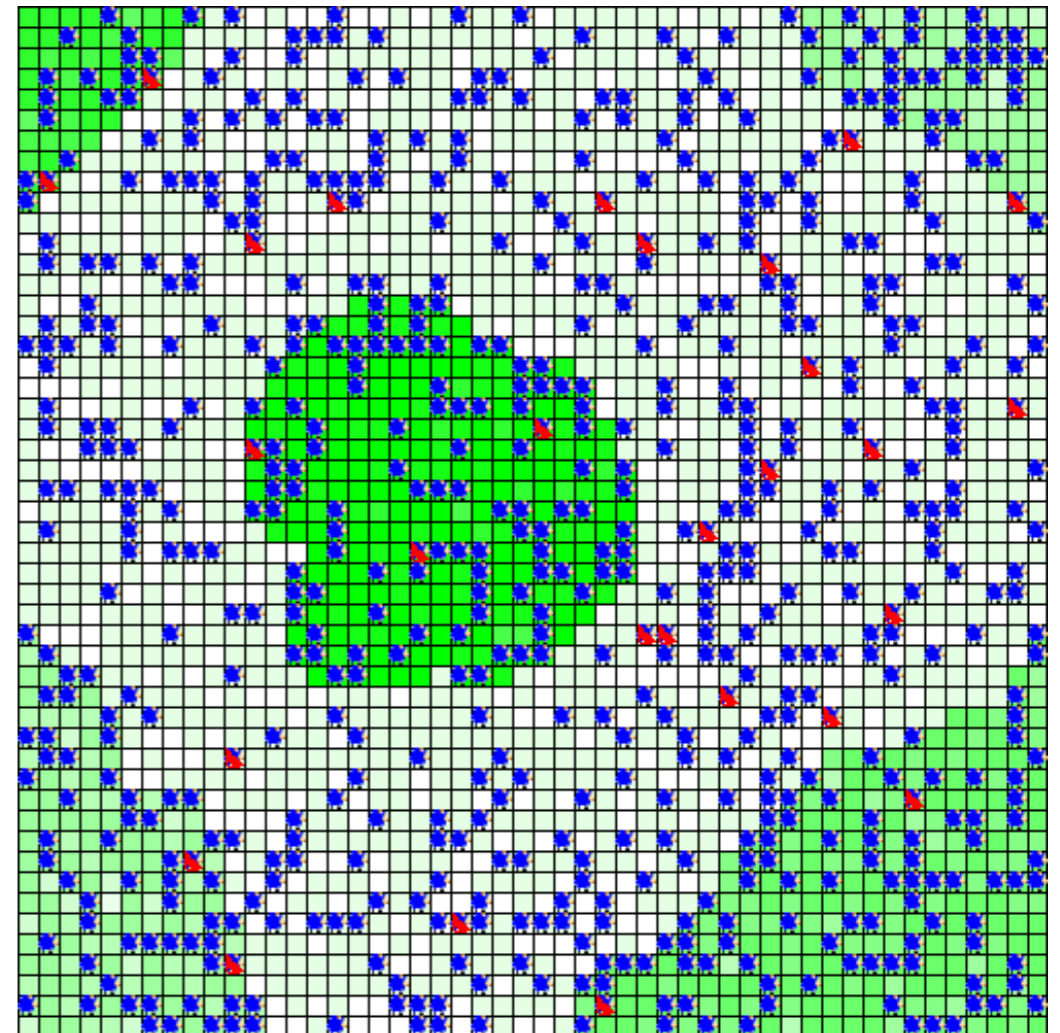
several columns for monitored variables
- just after the parameters columns
- names start with **r_**

Visualizing simulation results

```
with(sir$output[[1]],  
     plot(Step, r_I,  
          ylab = "nb. infected"))
```



```
path_to_movie <-  
  make_movie(pp$output[[1]],  
            "r_main_display")
```



<https://r-and-gama.github.io/gamar>

rama **0.1.0** [Home](#) [Get started](#) [Tutorials](#) [Examples](#) [Reference](#)

rama is an R interface to the **GAMA agent-based** simulation platform. It allows to

- **read** an experiment of a model defined in a `.gaml` file,
- **manipulate** this experiment, including generate experiment plans and
- **run** the simulations defined in an experiment plan.

An **experiment** is a group of simulations. A **simulation** is an execution of a given **model** with

- a given set of **parameters values**,
- a given set of **observed variables**,
- a **duration** of simulation and
- a **seed** value.

All the simulations of an experiment relate to the same model. In R, an experiment belongs to the class `experiment` that is an extension of the class `data.frame`. The creation and manipulation of experiments can thus efficiently be performed with the `data.frame` methods. The class `experiment` is also **tidyverse**-compliant, which allows its insertion into **pipelines** (or workflows). Outputs of `experiment` runs are in an object of class `experiment` too, with fields corresponding to the simulation outputs, typically data frames of time series of observed variables and / or links to snapshots that can subsequently be assembled into movies. The R environment allows to

- create **experimental designs** (for example with the `expand.grid()` function),
- **statistically explore** results of simulation (how the parameters values influence the dynamics of the variables),
- perform **sensitivity analysis** of model's parameters (how much each parameter quantitatively influences the outputs),
- **estimate parameters values** (model calibration) if real data are available for the model's state variables.

In addition to above-mentioned data frame, an object of class `experiment` contains a link to a `.gaml` file containing the GAML model (**input**) and a link to a folder containing the **outputs** of simulations. It is possible to change these links but potentially dangerous and not advised. The `.gaml` file can be visualized in R but is not supposed to be modified by the user in R. Instead, a safe practice is to develop the model in the **GAMA** software and to reserve the use of `rama` to the design and exploitation of experiments' simulations as outlined above.

Links

Browse source code at <https://github.com/r-and-gama/rama>

Report a bug at <https://github.com/r-and-gama/rama/issues>

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Dev status

lifecycle **experimental**

CRAN **not published**

build **passing**

build **passing**

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Dev status

lifecycle **experimental**

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