

# Package ‘goldfish’

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**Title** Statistical Network Models for Dynamic Network Data

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**License** GPL (>= 3)

**Description** Tools for fitting statistical network models to dynamic network data. Can be used for fitting both dynamic network actor models ('DyNAMs') and relational event models ('REMs').  
Stadtfeld, Hollway, and Block (2017a) <[doi:10.1177/0081175017709295](https://doi.org/10.1177/0081175017709295)>,  
Stadtfeld, Hollway, and Block (2017b) <[doi:10.1177/0081175017733457](https://doi.org/10.1177/0081175017733457)>,  
Stadtfeld and Block (2017) <[doi:10.15195/v4.a14](https://doi.org/10.15195/v4.a14)>,  
Hoffman et al. (2020) <[doi:10.1017/nws.2020.3](https://doi.org/10.1017/nws.2020.3)>.

**URL** <https://snlab-ch.github.io/goldfish/>

**BugReports** <https://github.com/snlab-ch/goldfish/issues>

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goldfish-package	<i>goldfish package</i>
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## Description

The goldfish Project is an R package that allows to fit statistical network models (such as DyNAM and REM) to dynamic network data.

## Details

The **goldfish** package in R allows the study of time-stamped network data using a variety of models. In particular, it implements different types of Dynamic Network Actor Models (DyNAMs), a class of models that is tailored to the study of actor-oriented network processes through time. Goldfish also implements different versions of the tie-oriented Relational Event Model by Carter Butts.

## References

- Stadtfeld, C. (2012). Events in Social Networks: A Stochastic Actor-oriented Framework for Dynamic Event Processes in Social Networks. *KIT Scientific Publishing*. doi:10.5445/KSP/1000025407
- Stadtfeld, C., and Block, P. (2017). Interactions, Actors, and Time: Dynamic Network Actor Models for Relational Events. *Sociological Science* 4 (1), 318-52. doi:10.15195/v4.a14
- Stadtfeld, C., Hollway, J., and Block, P. (2017). Dynamic Network Actor Models: Investigating Coordination Ties Through Time. *Sociological Methodology* 47 (1). doi:10.1177/0081175017709295
- Hollway, J. (2020). Network embeddedness and the rate of international water cooperation and conflict. In *Networks in Water Governance*, edited by Manuel Fischer and Karin Ingold. London: Palgrave, pp. 87-113.
- Hoffman, M., Block P., Elmer T., and Stadtfeld C. (2020). A model for the dynamics of face-to-face interactions in social groups. *Network Science*, 8(S1), S4-S25. doi:10.1017/nws.2020.3

## See Also

[estimate\(\)](#)

---

defineDependentEvents *Define dependent events for a model*

---

## Description

The final step in defining the data objects is to identify the dependent events.

## Usage

```
defineDependentEvents(
  events,
  nodes,
  nodes2 = NULL,
  defaultNetwork = NULL,
  envir = environment()
)
```

## Arguments

- |                |  |
|----------------|--|
| events         | a data frame containing the event list that should be considered as a dependent variable in models.  |
| nodes          | a data frame or a <code>nodes.goldfish</code> object containing the nodes used in the event list.  |
| nodes2         | a second nodeset in the case that the events occurs in a two-mode network.   |
| defaultNetwork | the name of a <code>network.goldfish</code> object.  |
| envir          | An <a href="#">environment</a> object where the nodes-set and default network objects are defined. The default value is <code>environment()</code> . |

## Details

Before this step is performed, we have to define: the nodeset ([defineNodes\(\)](#)), the network ([defineNetwork\(\)](#)) and the link the event list to the network ([linkEvents\(\)](#)).

During the definition as a dependent event, some checks are done to ensure consistency with the default network and the nodeset. In particular, consistency of the labels of nodes in the events with the nodes' labels in the network and the nodeset is done.

It is possible to define as a dependent event a different set of events to the ones link to the default network. This is useful to model different type of events where the event dynamic is driven by different effects or its weight differs. [Fisheries\\_Treaties\\_6070](#) has an example of it, the relational event modeled are fisheries treaties between countries. The `bilatchanges` data frame contains information of creation and dissolution of treaties. `vignette(teaching2)` shows how to model just the creation of treaties conditional on creation and dissolution.

## Value

an object with additional class `dependent.goldfish` with attributes:

<code>nodes</code>	a character vector with the names of the nodes set that define the dimensions of the <code>defaultNetwork</code> . <code>nodes</code> and <code>nodes2</code> arguments.
<code>defaultNetwork</code>	A character value with the name of the network object when this is present. <code>defaultNetwork</code> argument.
<code>type</code>	A character value that can take values <code>monadic</code> or <code>dyadic</code> depending on the arguments used during the definition.

The object can be modified using methods for data frames.

## See Also

[defineNodes\(\)](#), [defineNetwork\(\)](#), [linkEvents\(\)](#)

## Examples

```
actors <- data.frame(
  actor = 1:5, label = paste("Actor", 1:5),
  present = TRUE, gender = sample.int(2, 5, replace = TRUE)
)
actors <- defineNodes(nodes = actors)
calls <- data.frame(
  time = c(12, 27, 45, 56, 66, 68, 87),
  sender = paste("Actor", c(1, 3, 5, 2, 3, 4, 2)),
  receiver = paste("Actor", c(4, 2, 3, 5, 1, 2, 5)), increment = rep(1, 7)
)
callNetwork <- defineNetwork(nodes = actors)
callNetwork <- linkEvents(
  x = callNetwork, changeEvent = calls, nodes = actors
)

# Defining the dependent events:
callDependent <- defineDependentEvents(
  events = calls, nodes = actors, defaultNetwork = callNetwork
```

)

---

defineGlobalAttribute *Define a global time-varying attribute*

---

### Description

This function allows to define a global attribute of the nodeset (i.e a variable that is identical for each node but changes over time).

### Usage

```
defineGlobalAttribute(global)
```

### Arguments

`global` a data frame containing all the values this global attribute takes along time.

### Details

For instance, seasonal climate changes could be defined as a changing global attribute. Then, this global attribute can be linked to the nodeset by using [linkEvents\(\)](#)

### Value

an object of class `global.goldfish`

### Examples

```
seasons <- defineGlobalAttribute(data.frame(time = 1:12, replace = 1:12))
```

---

```
defineGroups_interaction
```

*To define the second mode of a DyNAM-i model*

---

### Description

This function create all objects necessary to the estimation of a DyNAM-i model `model = "DyNAMi"` from dyadic interaction records and an actor set. It first creates a nodeset for the second mode of the interaction network that will be modeled, i.e. the interaction groups set, and an event list that indicates when groups are present or not through time. It then creates a list of interaction events, between actors and groups, in which an actor either joins or leaves a group. It is decomposed in an list of dependent events (that should be modeled) and a list of exogenous events (that should not be modeled). For example when an actor leaves a group and joins her own singleton group, only the leaving event is modeled but not the joining one, and vice versa when an actor belonging to a singleton group joins another group.

**Usage**

```
defineGroups_interaction(
  records,
  actors,
  seed.randomization,
  progress = getOption("progress")
)
```

**Arguments**

records	an object of class <code>data.frame</code> that is a list of rows of type <code>node A</code> , <code>nodeB</code> , <code>Start</code> , <code>End</code> , where <code>nodeA</code> and <code>nodeB</code> indicate the actors involved in a dyadic interaction, and <code>Start</code> and <code>End</code> indicating the starting and ending time of their interaction.
actors	a object of class <code>nodes.goldfish</code> that defines the actors interacting (labels in records and actors should be identical).
seed.randomization	an integer used whenever there should be some random choice to be made.
progress	logical weather detailed information of intermediate steps should be printed in the console.

**Details**

It is important to notice that sometimes some random decisions have to be made regarding who joined or left a group, for example when two actors start interacting but we do not know who initiated the interaction. To test for the robustness of such a procedure, one can use different randomization seeds and run the model several times.

**Value**

a list with the following data frames

**interaction.updates** containing all joining and leaving events

**groups** containing the nodeset corresponding to interaction groups (the second mode of the network)

**dependent.events** for the events that should be modeled

**exogenous.events** that are not modeled (for example when an actor leaves a group and joins its own singleton group, only the leaving event is modeled but not the joining event)

**composition.changes** that is an events list that should be attached to the groups nodeset to indicate when a group is present or not

## Description

The function defines a network object either from a nodeset or from a matrix (sociomatrix or adjacency matrix). If a matrix is used as input, `defineNetwork()` returns a network filled with the same values as the ones present in the provided network. If the nodeset is the only argument, `defineNetwork()` returns an empty network with the number of columns and rows corresponding to the size of the nodeset. These networks are static, but they can be turned into dynamic networks by linking dynamic events to the network objectw using `linkEvents()`.

## Usage

```
defineNetwork(  
  matrix = NULL,  
  nodes,  
  nodes2 = NULL,  
  directed = TRUE,  
  envir = environment()  
)
```

## Arguments

<code>matrix</code>	An initial matrix (optional), and object of class <code>matrix</code> .
<code>nodes</code>	A node-set (see <code>defineNodes()</code> ).
<code>nodes2</code>	A second optional node-set for the definition of two-mode networks.
<code>directed</code>	A logical value indicating whether the network is directed.
<code>envir</code>	An <code>environment</code> object where the nodes-set objects are defined. The default value is <code>environment()</code> .

## Details

If a matrix is used as input, its dimension names must be a subset of the nodes in the nodeset as defined with the `defineNodes()` and the order of the labels in rows and columns must correspond to the order of node labels in the nodeset. The matrix can be directed or undirected (as specified with the `directed` argument).

If the network is updated over time (e.g., a new wave of friendship data is collected), these changes can be added with the `linkEvents()` - similar to link changing attribute events to a nodeset. This time, the user needs to provide the network and the associated nodeset. If no matrix is provided, goldfish only considers the nodeset and assumes the initial state to be empty (i.e., a matrix containing only 0s).

**Value**

an object with additional class `network.goldfish` with attributes:

`nodes` a character vector with the names of the nodes set objects used during the definition. `nodes` and `nodes2` arguments.

`directed` Logical value indicating whether the network is directed. `directed` argument

`events` An empty character vector. `linkEvents()` is used to link event data frames.

The object can be modified using methods for matrix.

**See Also**

`defineNodes()`, `linkEvents()`

**Examples**

```
# If no initial matrix is provided
data("Social_Evolution")
callNetwork <- defineNetwork(nodes = actors)

# If a initial matrix is provided
data("Fisheries_Treaties_6070")
bilatnet <- defineNetwork(bilatnet, nodes = states, directed = FALSE)
```

---

defineNodes

*Defining a node set with (dynamic) node attributes.*

---

**Description**

The `defineNodes()` function processes and checks the data.frame passed to the `nodes` argument. This is a recommended step before the definition of the network.

**Usage**

```
defineNodes(nodes)
```

**Arguments**

`nodes` a data.frame object with the nodes attributes with the following reserved names

- label** character variable containing the nodes labels (mandatory)
- present** logical variable indicating if the respective node is present at the first time-point (optional)



## Details

Additional variables in the nodes data frame object are considered as the initial values of the nodes attributes. Those variables must be of class `numeric`, `character`, `logical`.

It is important that the initial definition of the node set contain all the nodes that could be potential senders or receivers of events. In case that all the nodes are not available at all times, the `present` variable can be used to define compositional changes. Therefore, the initial node set would contain all the potential senders and receivers nodes and the variable `present` will indicate all the nodes present at the beginning as senders or receivers. Using `linkEvents()` is possible to link events where the composition of available nodes changes over time, see `vignette("teaching2")`.

For the attributes in the nodeset to become dynamic, them can be linked to a dynamic event-list data frames in the initial state object by using the `linkEvents()`. A new call of `linkEvents()` is required for each attribute that is dynamic.

Objects of class `tbl_df` from the tibble package frequently use in the tidyverse ecosystem and objects of class `data.table` will produce errors in later steps for goldfish. Current implementation of goldfish relies on the subsetting behavior of data frames objects. The previous mentioned objects classes change this behavior producing errors.

## Value

an object with an additional class `nodes.goldfish` with attributes:

`events` An empty character vector. `linkEvents()` is used to link event data frames.

`dynamicAttributes`

An empty character vector. `linkEvents()` is used to link event data frames and their related attribute.

The object can be modified using methods for data frames.

## See Also

`defineNetwork()`, `linkEvents()`

## Examples

```
nodesAttr <- data.frame(
  label = paste("Actor", 1:5),
  present = c(TRUE, FALSE, TRUE, TRUE, FALSE),
  gender = c(1, 2, 1, 1, 2)
)
nodesAttr <- defineNodes(nodes = nodesAttr)

# Social evolution nodes definition
data("Social_Evolution")
actors <- defineNodes(actors)

# Fisheries treaties nodes definition
data("Fisheries_Treaties_6070")
states <- defineNodes(states)
```

estimate

*Estimate a model***Description**

Estimates parameters for a dynamic network model via maximum likelihood implementing the iterative Newton-Raphson procedure as describe in Stadtfeld and Block (2017).

**Usage**

```
estimate(
  x,
  model = c("DyNAM", "REM", "DyNAMI"),
  subModel = c("choice", "rate", "choice_coordination"),
  estimationInit = NULL,
  preprocessingInit = NULL,
  preprocessingOnly = FALSE,
  progress = getOption("progress"),
  verbose = getOption("verbose")
)
```

**Arguments**

x	a formula that defines at the left-hand side the dependent network (see <a href="#">defineDependentEvents()</a> ) and at the right-hand side the effects and the variables for which the effects are expected to occur (see <code>vignette("goldfishEffects")</code> ).
model	a character string defining the model type. Current options include "DyNAM", "DyNAMI" or "REM"  <b>DyNAM</b> Dynamic Network Actor Models (Stadtfeld, Hollway and Block, 2017 and Stadtfeld and Block, 2017) <b>DyNAMI</b> Dynamic Network Actor Models for interactions (Hoffman et al., 2020) <b>REM</b> Relational Event Model (Butts, 2008)
subModel	a character string defining the submodel type. Current options include "choice", "rate" or "choice_coordination"  <b>choice</b> a multinomial receiver choice model model = "DyNAM" (Stadtfeld and Block, 2017), or the general Relational event model model = "REM" (Butts, 2008). A multinomial group choice model model = "DyNAMI" (Hoffman et al., 2020) <b>choice_coordination</b> a multinomial-multinomial model for coordination ties model = "DyNAM" (Stadtfeld, Hollway and Block, 2017) <b>rate</b> A individual activity rates model model = "DyNAM" (Stadtfeld and Block, 2017). Two rate models, one for individuals joining groups and one for individuals leaving groups, jointly estimated model = "DyNAMI"(Hoffman et al., 2020)

- estimationInit** a list containing lower level technical parameters for estimation. It may contain:
- initialParameters** a numeric vector. It includes initial parameters of the estimation. Default is set to NULL.
  - fixedParameters** a numeric vector. It specifies which component of the coefficient parameters (intercept included) is fixed and the value it takes during estimation, e.g., if the vector is `c(2, NA)` then the first component of the parameter is fixed to 2 during the estimation process. Default is set to NULL, i.e. all parameters are estimated. Note that it must be consistent with `initialParameters`.
  - maxIterations** maximum number of iterations of the Gauss/Fisher scoring method for the estimation. Default is set to 20.
  - maxScoreStopCriterion** maximum absolute score criteria for successful convergence. Default value is 0.001
  - initialDamping** a numeric vector used to declare the initial damping factor for each parameter. It controls the size of the update step during the iterative estimation process. The default is set to 30 when the formula has windowed effects or 10 in another case, see `vignette("goldfishEffects")`.
  - dampingIncreaseFactor** a numeric value. It controls the factor that increases the damping of the parameters when improvements in the estimation are found.
  - dampingDecreaseFactor** a numeric value. Controls the factor that decreases the damping of the parameters when no improvements in the estimation are found.
  - returnIntervalLogL** a logical value. Whether to keep the log-likelihood of each event from the final iteration of the Gauss/Fisher estimation method.
  - engine** a string indicating the estimation engine to be used. Current options include `"default"`, `"default_c"`, and `"gather_compute"`. The default value is `"default"`, it is an estimation routine implemented in pure R code. `"default_c"` uses a C implementation of the `"default"` routine. `"gather_compute"` uses a C implementation with a different data structure that reduces the time but it can increase the memory usage.
  - startTime** a numerical value or a date-time character with the same time-zone formatting as the times in event that indicates the starting time to be considered during estimation. *Note:* it is only use during preprocessing
  - endTime** a numerical value or a date-time character with the same time-zone formatting as the times in event that indicates the end time to be considered during estimation. *Note:* it is only use during preprocessing
  - opportunitiesList** a list containing for each dependent event the list of available nodes for the choice model, this list should be the same length as the dependent events list (ONLY for choice models).
- preprocessingInit**  
a `preprocessed.goldfish` object computed for the current formula, allows skipping the preprocessing step.
- preprocessingOnly**  
logical indicating whether only preprocessed statistics should be returned rather than a `result.goldfish` object with the estimated coefficients.

progress	logical indicating whether should print a minimal output to the console of the progress of the preprocessing and estimation processes.
verbose	logical indicating whether should print very detailed intermediate results of the iterative Newton-Raphson procedure; slows down the routine significantly.

### Details

Missing data is imputed during the preprocessing stage. For network data missing values are replaced by a zero value, it means that is assuming a not tie/event explicitly. For attributes missing values are replaced by the mean value, if missing values are presented during events updates they are replace by the mean of the attribute in that moment of time.

### Value

returns an object of `class()` "result.goldfish" when preprocessingOnly = FALSE or a preprocessed statistics object of class "preprocessed.goldfish" when preprocessingOnly = TRUE.

An object of class "result.goldfish" is a list including:

parameters	a numeric vector with the coefficients estimates.
standardErrors	a numeric vector with the standard errors of the coefficients estimates.
logLikelihood	the log-likelihood of the estimated model
finalScore	a vector with the final score reach by the parameters during estimation.
finalInformationMatrix	a matrix with the final values of the negative Fisher information matrix. The inverse of this matrix gives the variance-covariance matrix for the parameters estimates.
convergence	a list with two elements. The first element (isConverged) is a logical value that indicates the convergence of the model. The second element (maxAbsScore) reports the final maximum absolute score in the final iteration.
nIterations	an integer with the total number of iterations performed during the estimation process.
nEvents	an integer reporting the number of events considered in the model.
names	a matrix with a description of the effects used for model fitting. It includes the name of the object used to calculate the effects and additional parameter description.
formula	a formula with the information of the model fitted.
model	a character value of the model type.
subModel	a character value of the subModel type.
rightCensored	a logical value indicating if the estimation process considered right-censored events. Only it is considered for DyNAM-rate (model = "DyNAM" and subModel = "rate") or REM (model = "REM") models, and when the model includes the intercept.



```

mod01 <- estimate(callsDependent ~ inertia + recip + trans,
                 model = "DyNAM", subModel = "choice",
                 estimationInit = list(engine = "default_c"))
summary(mod01)

# A individual activity rates model
mod02 <- estimate(callsDependent ~ 1 + nodeTrans + indeg + outdeg,
                 model = "DyNAM", subModel = "rate",
                 estimationInit = list(engine = "default_c"))
summary(mod02)

# A multinomial-multinomial choice model for coordination ties
data("Fisheries_Treaties_6070")
states <- defineNodes(states)
states <- linkEvents(states, sovchanges, attribute = "present")
states <- linkEvents(states, regchanges, attribute = "regime")
states <- linkEvents(states, gdpchanges, attribute = "gdp")

bilatnet <- defineNetwork(bilatnet, nodes = states, directed = FALSE)
bilatnet <- linkEvents(bilatnet, bilatchanges, nodes = states)

contignet <- defineNetwork(contignet, nodes = states, directed = FALSE)
contignet <- linkEvents(contignet, contigchanges, nodes = states)

createBilat <- defineDependentEvents(
  events = bilatchanges[bilatchanges$increment == 1, ],
  nodes = states, defaultNetwork = bilatnet
)

partnerModel <- estimate(
  createBilat ~
  inertia(bilatnet) +
  indeg(bilatnet, ignoreRep = TRUE) +
  trans(bilatnet, ignoreRep = TRUE) +
  tie(contignet) +
  alter(states$regime) +
  diff(states$regime) +
  alter(states$gdp) +
  diff(states$gdp),
  model = "DyNAM", subModel = "choice_coordination",
  estimationInit = list(initialDamping = 40, maxIterations = 30)
)
summary(partnerModel)

```

## Description

Provide diagnostic functions for an object of class `result.goldfish`. `outliers` helps to identify outliers events. `changepts` helps to identify where a change point in the events sequence is presented using the log-likelihood.

## Usage

```
examineOutliers(
  x,
  method = c("Hampel", "IQR", "Top"),
  parameter = 3,
  window = NULL
)

examineChangepts(
  x,
  moment = c("mean", "variance"),
  method = c("PELT", "AMOC", "BinSeg"),
  window = NULL,
  ...
)
```

## Arguments

<code>x</code>	an object of class <code>result.goldfish</code> output from an <code>estimate</code> call.
<code>method</code>	Choice of "AMOC", "PELT" or "BinSeg". For a detail description see <code>cpt.mean</code> or <code>cpt.var</code> . The default value is "PELT".
<code>parameter</code>	An integer that represents the number of absolute outliers to identify, the threshold for the Hampel filter, i.e. $\text{parameter} * \text{MAD}$ , or the threshold beyond the interquartile range halved, i.e. $\text{parameter}/2 * \text{IQR}$ .
<code>window</code>	The window half-width for the Hampel filter. By default it is half the width of the event sequence.
<code>moment</code>	character argument to choose between "mean" or "variance". See section <i>Change point</i> for details.
<code>...</code>	additional arguments to be passed to the functions in the <code>changept</code> package.

## Value

NULL if neither outliers nor change points are identified. An object of class `ggplot` object from a call of `ggplot2::ggplot()`. It can be modified using the `ggplot2` syntax.

## Outliers

`examineOutliers` creates a plot with the log-likelihood of the events in the y-axis and the event index in the x-axis, identifying observations with labels indicating the sender and recipient.

### Change point

The parameter `moment` controls which method from the package `changepoint` is used:

"mean" It uses the `cpt.mean` function to investigate optimal positioning and (potentially) number of change points for the log-likelihood of the events in mean.

"variance" It uses the `cpt.var` function to investigate optimal positioning and (potentially) number of change points for the log-likelihood of the events in variance

The function call creates a plot with the log-likelihood of the events in the y-axis and the event index in the x-axis, highlighting the change point sections identified by the method.

### Examples

```
# A multinomial receiver choice model
data("Social_Evolution")
callNetwork <- defineNetwork(nodes = actors, directed = TRUE)
callNetwork <- linkEvents(
  x = callNetwork, changeEvent = calls,
  nodes = actors
)
callsDependent <- defineDependentEvents(
  events = calls, nodes = actors,
  defaultNetwork = callNetwork
)

mod01 <- estimate(callsDependent ~ inertia + recip + trans,
  model = "DyNAM", subModel = "choice",
  estimationInit = list(returnIntervalLogL = TRUE,
    engine = "default_c")
)

examineOutliers(mod01)

examineChangepoints(mod01)
```

---

Fisheries\_Treaties\_6070

*International bilateral fisheries treaties (1960-1970)*

---

### Description

An abbreviated version of the international fisheries agreements dataset, including only bilateral agreements, fewer variables, and ranging only between 1960 and 1970 inclusive. This data set is only meant for testing, and not for inference. It provides an example of an undirected, weighted (by integer/increment) network, with composition change and both monadic and dyadic covariates. Monadic variables include the dates states gain or lose sovereign status, their polity score, and their GDP. Dyadic variables include bilateral fisheries agreements between states, and states' contiguity with one another over time.



**Usage**

```
data(Fisheries_Treaties_6070)
```

```
bilatchanges
```

```
bilatnet
```

```
contigchanges
```

```
contignet
```

```
gdpchanges
```

```
regchanges
```

```
sovchanges
```

```
states
```

**Format**

The data includes several dataframes: states (154 rows, 4 columns, monadic), sovchanges (62 rows, 3 columns, monadic), regchanges (145 rows, 3 columns, monadic), gdpchanges (979 rows, 3 columns, monadic), bilatchanges (77 rows, 4 columns, dyadic), contigchanges (139 rows, 4 columns, dyadic). See below for variables and formats.

<b>Object</b>	<b>Description</b>	<b>Format</b>
states\$label	Node identifier labels	character
states\$present	Node present in dataset	boolean
states\$regime	Placeholder for regime variable	numeric (NA)
states\$gdp	Placeholder for GDP variable	numeric (NA)
sovchanges\$time	Date of state sovereignty update	POSIXct
sovchanges\$node	Node for state sovereignty update	integer
sovchanges\$replace	State sovereignty update	boolean
regchanges\$time	Date of regime update	POSIXct
regchanges\$node	Node for regime update	integer
regchanges\$replace	Regime update	integer (-10–10)
gdpchanges\$time	Date of GDP update	POSIXct
gdpchanges\$node	Node for GDP update	integer
gdpchanges\$replace	GDP update	numeric
bilatchanges\$time	Date of bilateral change	POSIXct
bilatchanges\$sender	First bilateral change node	integer
bilatchanges\$receiver	Second bilateral change node	integer
bilatchanges\$increment	Create or dissolve action	numeric (-1 or 1)
contigchanges\$time	Date of contiguity change	POSIXct
contigchanges\$sender	First contiguity change node	integer
contigchanges\$receiver	Second contiguity change node	integer
contigchanges\$replace	New contiguity value	numeric

An object of class `data.frame` with 77 rows and 4 columns.

An object of class `matrix` (inherits from `array`) with 154 rows and 154 columns.

An object of class `data.frame` with 139 rows and 4 columns.

An object of class `matrix` (inherits from `array`) with 154 rows and 154 columns.

An object of class `data.frame` with 979 rows and 3 columns.

An object of class `data.frame` with 145 rows and 3 columns.

An object of class `data.frame` with 62 rows and 3 columns.

An object of class `data.frame` with 154 rows and 4 columns.

## References

Hollway, James, and Johan Koskinen. 2016. Multilevel Embeddedness: The Case of the Global Fisheries Governance Complex. *Social Networks*, 44: 281-94. doi:10.1016/j.socnet.2015.03.001.

Hollway, James, and Johan H Koskinen. 2016. Multilevel Bilateralism and Multilateralism: States' Bilateral and Multilateral Fisheries Treaties and Their Secretariats. In *Multilevel Network Analysis for the Social Sciences*, edited by Emmanuel Lazega and Tom A B Snijders, 315-32. Cham: Springer International Publishing. doi:10.1007/978-3-319-24520-1\_13.

---

linkEvents

*Link dynamic events to a nodeset or a network*

---

## Description

Link events stored in data frames that modify attributes of the nodeset or modify ties in a network.

## Usage

```
linkEvents(x, ...)

## S3 method for class 'nodes.goldfish'
linkEvents(x, changeEvents, attribute, ...)

## S3 method for class 'network.goldfish'
linkEvents(x, changeEvents, nodes = NULL, nodes2 = NULL, ...)

## Default S3 method:
linkEvents(x, ...)
```

## Arguments

<code>x</code>	Either a nodeset ( <code>nodes.goldfish</code> object) or a network ( <code>network.goldfish</code> object)
<code>...</code>	additional arguments to be passed to the method.
<code>changeEvents</code>	The name of a data frame that represents a valid events list.

attribute	a character vector indicating the names of the attributes that should be updated by the specified events (ONLY if the object is a nodeset).
nodes	a nodeset (data.frame or nodes.goldfish object) related to the network (ONLY if x is a network)
nodes2	an optional nodeset (data.frame or nodes.goldfish object) related to the network (ONLY if x is a network)

## Details

The data frame that contains the events must contain variables with specific names depending if they refer to dynamic attributes or dynamic networks.

For dynamic networks stored in object of class `network.goldfish` the `changeEvents` data frame must contain the following variables:

**time** numeric or POSIXct (see `base::as.Date()`) variable containing the time-stamps when the event happen.

**sender** character variable indicating the label of the sender of the event.

**receiver** character variable indicating the label of the receiver of the event.

See the `bilatchanges` and `contigchanges` data frames in the [Fisheries\\_Treaties\\_6070](#) datasets for examples of event data frames that relate with dynamic networks.

For dynamic attributes stored in object of class `nodes.goldfish` the `changeEvents` data frame must contain the following variables:

**time** numeric or POSIXct (see `base::as.Date()`) variable containing the time-stamps when the event happen.

**label** character variable indicating the label of the node for which the attribute changes.

See `sovchanges`, `regchanges` and `gdpchanges` data frames in the [Fisheries\\_Treaties\\_6070](#) datasets for examples of event data frames that relate with dynamic attributes

For both cases an additional variable indicates the change in value of either the ties or attributes. The class of this variable must be the same as the tie value or attribute value that will be updated, i.e., when the present variable is dynamic the updating values must be logical (see `defineNodes()` for a description of this variable. There are two possibilities on how to specify those changes but only one can be used at a time:

**increment** with a numerical value that represent the increment (when it's positive value) or the decrement (when it's a negative value) of the dynamic element from their past value (with respect to the time value).

In the [Social\\_Evolution](#) dataset the `calls` data frame contains the calling events between students where the increment represent a new call. With every new call the dyad (sender-receiver) increase the count of calls that had happen in the past.

**replace** contains the value that would replace at point-time `time` the attribute or tie value. It is usually the way to represent changes in node attributes.

In the [Fisheries\\_Treaties\\_6070](#) dataset the `sovchanges`, `regchanges` and `gdpchanges` data frames are examples where the `replace` variable is used to specify attribute changes and their class match with the variable in the node set.

Dynamic network attributes can be also defined using the `replace` variable. The `contigchanges` data frame in the [Fisheries\\_Treaties\\_6070](#) dataset, and friendship data frame in the [Social\\_Evolution](#) are examples of this.

### Value

an object with the same class as the object `x`. For objects of class `network.goldfish` the attribute `events` with the name of the event data frame passed through with the argument `changeEvents`. For objects of class `nodes.goldfish` attributes `events` and `dynamicAttribute` are modified with name of the objects passed through with the arguments `changeEvents` and `attribute` respectively.

### See Also

[defineNodes\(\)](#), [defineNetwork\(\)](#)

### Examples

```
actors <- data.frame(
  actor = 1:5, label = paste("Actor", 1:5),
  present = TRUE, gender = sample.int(2, 5, replace = TRUE)
)
actors <- defineNodes(nodes = actors)
callNetwork <- defineNetwork(nodes = actors)

# Link events to a nodeset
compositionChangeEvents <- data.frame(
  time = c(14, 60),
  node = "Actor 4",
  replace = c(FALSE, TRUE)
)
actorsnew <- linkEvents(
  x = actors, attribute = "present", changeEvents = compositionChangeEvents
)

# Link events to a Network
calls <- data.frame(
  time = c(12, 27, 45, 56, 66, 68, 87),
  sender = paste("Actor", c(1, 3, 5, 2, 3, 4, 2)),
  receiver = paste("Actor", c(4, 2, 3, 5, 1, 2, 5)),
  increment = rep(1, 7)
)
callNetwork <- linkEvents(
  x = callNetwork, changeEvent = calls, nodes = actors
)
```

## Description

This function extract the log-likelihood from the output of a `estimate` call. The extracted log-likelihood correspond to the value in the last iteration of the `estimate` call, users should check convergence of the Gauss/Fisher scoring method before using the log-likelihood statistic to compare models.

## Usage

```
## S3 method for class 'result.goldfish'  
logLik(object, ..., avgPerEvent = FALSE)
```

## Arguments

<code>object</code>	an object of class <code>result.goldfish</code> output from an <code>estimate</code> call with a fitted model.
<code>...</code>	additional arguments to be passed.
<code>avgPerEvent</code>	a logical value indicating whether the average likelihood per event should be calculated.

## Details

Users might use `stats::AIC()` and `stats::BIC()` to compute the Information Criteria from one or several fitted model objects. An information criterion could be used to compare models with respect to their predictive power.

Alternatively, `lmtest::lrtest()` can be used to compare models via asymptotic likelihood ratio tests. The test is designed to compare nested models. i.e., models where the model specification of one contains a subset of the predictor variables that define the other.

## Value

Returns an object of class `logLik` when `avgPerEvent = FALSE`. This is a number with the extracted log-likelihood from the fitted model, and with the following attributes:

<code>df</code>	degrees of freedom with the number of estimated parameters in the model
<code>nobs</code>	the number of observations used in estimation. In general, it corresponds to the number of dependent events used in estimation. For a <code>subModel = "rate"</code> or <code>model = "REM"</code> with intercept, it corresponds to the number of dependent events plus right-censored events due to exogenous or endogenous changes.

When `avgPerEvent = TRUE`, the function returns a number with the average log-likelihood per event. The total number of events depends on the presence of right-censored events in a similar way that the attribute `nobs` is computed when `avgPerEvent = FALSE`.

---

print-method	<i>Methods for goldfish objects.</i>
--------------	--------------------------------------

---

### Description

Printing functions for goldfish objects.

### Usage

```
## S3 method for class 'result.goldfish'
print(
  x,
  ...,
  digits = max(3, getOption("digits") - 2),
  width = getOption("width"),
  complete = FALSE
)

## S3 method for class 'summary.result.goldfish'
print(
  x,
  ...,
  digits = max(3, getOption("digits") - 2),
  width = getOption("width"),
  complete = FALSE
)

## S3 method for class 'nodes.goldfish'
print(x, ..., full = FALSE, n = 6)

## S3 method for class 'network.goldfish'
print(x, ..., full = FALSE, n = 6L)

## S3 method for class 'dependent.goldfish'
print(x, ..., full = FALSE, n = 6)

## S3 method for class 'preprocessed.goldfish'
print(x, ..., width = getOption("width"))
```

### Arguments

x	an object of class <code>result.goldfish</code> , <code>summary.result.goldfish</code> , <code>nodes.goldfish</code> , <code>network.goldfish</code> , <code>dependent.goldfish</code> , or <code>preprocessed.goldfish</code> .
...	further arguments to be passed to the respective default method.
digits	minimal number of significant digits, see <a href="#">print.default()</a> .

width	controls the maximum number of columns on a line used in printing <code>summary.result.goldfish</code> and <code>preprocessed.goldfish</code> , see <code>print.default()</code> .
complete	logical. Indicates whether the parameter coefficients of effects fixed during estimation using <code>fixedParameters</code> should be printed. The default value is <code>FALSE</code> . <i>Note:</i> applies for objects of class <code>result.goldfish</code> and <code>summary.result.goldfish</code> .
full	logical. Indicates whether the complete matrix/data.frame should be printed. The default value <code>FALSE</code> .
n	number of rows for <code>data.frame</code> , and rows and columns for <code>matrix</code> to be printed.

## Value

Not value, called for printing side effect.

For objects of class `result.goldfish` and `summary.result.goldfish` print the estimated coefficients when `complete = FALSE`, otherwise it includes also the fixed coefficients. For `summary.result.goldfish` print:

Effect details:

a table with additional information of the effects. The information corresponds to the values of the effects arguments when they are modified and if they were fixed during estimation, see `vignette("goldfishEffects")` for the complete list of arguments, and `estimate()` on how to fix coefficients during estimation.

Coefficients: a table with the estimated coefficients, their approximate standard error obtained from the inverse of the negative Fisher information matrix, z-value and the p-value of the univariate two-tailed Wald test to test the hypothesis that the parameter is 0.

Convergence and Information Criteria:

Information about the convergence of the iterative Newton-Raphson procedure and the score value in the last iteration. Information criteria as the AIC, BIC and the AIC corrected for small sample size AICc are reported.

Model and subModel:

the values set during estimation.

For objects of class `nodes.goldfish` print information of the total number of nodes in the object, the number of nodes present at the beginning of preprocessing, a table with the linked attributes with their respective events data frame and a printing of the first rows in the nodes data frame. See `defineNodes()`.

For objects of class `network.goldfish` print information of the dimensions of the network, number of ties presented at the beginning of the preprocessing, the nodes data frames linked to it, information about their definition as a one-mode and directed network, linked events data frame to it and a printing of the first rows and columns in the array. See `defineNetwork()`.

For objects of class `dependent.goldfish` print information of the total number of events in the object, linked nodes set(s), linked default network and a printing of the first rows in the events data frame. See `defineDependentEvents()`.

---

 RFID\_Validity\_Study    *RFID Validity dataset*


---

### Description

Dataset collected at ETH Zürich by Timon Elmer and colleagues in order to test the accuracy of Radio Frequency Identification (RFID) badges for measuring social interactions. Social interactions of 11 individuals (from the university staff) were recorded with RFID badges in an informal setting. They were then compared to the interactions observed by two confederates who watched the video recording of the event. The RFID data went through the data processing procedure detailed in the original article. See Elmer et al, 2019 for more details, and the OSF platform for all details on the dataset.

### Usage

```
data(RFID_Validity_Study)

rfid

video

known.before

participants
```

### Format

3 dataframes:

- participants (11 rows, 7 columns): attributes of the experiment's participants
- rfid (1011 rows, 4 columns): dyadic interactions detected by the RFID badges (after data processing)
- video (219 rows, 4 columns): dyadic interactions detected by the video rating and one network:
- known.before (11 rows, 11 columns): network of previous acquaintances  
See below for variables and formats.

<b>Object</b>	<b>Description</b>	<b>Format</b>
participants\$actor	Identifier of the actor	integer
participants\$label	(Anonymized) name	Factor
participants\$present	Presence of the actor (all actors are present)	logical



participants\$age	Actor's age	integer
participants\$gender	Actor's gender (0: male, 1: female)	integer
participants\$group	Actor's group affiliation (groups have distinct ids)	integer
participants\$level	Actor's seniority (1: MSc student, 2: PhD student, 3: PostDoc, 4: Prof)	integer
rfid\$NodeA	Identifier for the first actor	chr
rfid\$NodeB	Identifier for the second actor	chr
rfid\$Start	Time of the beginning of the dyadic interaction	integer
rfid\$End	Time of the end of the dyadic interaction	integer
video\$NodeA	Identifier for the first actor	chr
video\$NodeB	Identifier for the second actor	chr
video\$Start	Time of the beginning of the dyadic interaction	integer
video\$End	Time of the end of the dyadic interaction	integer

An object of class `data.frame` with 1011 rows and 4 columns.

An object of class `data.frame` with 219 rows and 4 columns.

An object of class `matrix` (inherits from `array`) with 11 rows and 11 columns.

An object of class `data.frame` with 11 rows and 7 columns.

### Source

<https://osf.io/rrhxe/>

### References

Elmer, T., Chaitanya, K., Purwar, P., & Stadtfeld, C. (2019). The validity of RFID badges measuring face-to-face interactions. *Behavior research methods*, 1-19. doi:10.3758/s134280181180y

---

Social\_Evolution

*Social evolution of a university dormitory cohort*

---

### Description

An abbreviated version of the MIT Reality Commons Social Evolution dataset, spanning a reduced time period and with fewer variables. Dyadic variables include binary friendships at time of survey, and time-stamped phone call occurrences. Individual variables include the floor of the dormitory on which the student resides, and the grade type of each student including freshmen, sophomore, junior, senior, or graduate tutors.

### Usage

```
data(Social_Evolution)
```

```
actors
```

```
calls
```

```
friendship
```

**Format**

3 dataframes: actors (84 rows, 4 columns), calls (439 rows, 4 columns), friendship (766 rows, 4 columns). See below for variables and formats.

<b>Object</b>	<b>Description</b>	<b>Format</b>
actors\$label	Actor identifier labels	character
actors\$present	Actor present in dataset	boolean
actors\$floor	Floor of residence actor lives on	numeric (1-9)
actors\$gradeType	Degree level	numeric (1-5)
calls\$time	Time and date of call	numeric from POSIXct
calls\$sender	Initiator of phone call	character
calls\$receiver	Recipient of phone call	character
calls\$increment	Indicates call number increment (all 1s)	numeric (1)
friendship\$time	Time and date of friend nomination	numeric from POSIXct
friendship\$sender	Nominator of friendship	character
friendship\$receiver	Nominee of friendship	character
friendship\$replace	Indicates friendship value at \$time	numeric

An object of class data.frame with 84 rows and 4 columns.

An object of class data.frame with 439 rows and 4 columns.

An object of class data.frame with 766 rows and 4 columns.

**Source**

<http://realitycommons.media.mit.edu/socialrevolution.html>

**References**

A. Madan, M. Cebrian, S. Moturu, K. Farrahi, A. Pentland (2012). Sensing the 'Health State' of a Community. *Pervasive Computing*. 11, 4, pp. 36-45.

---

update-method

*Methods to update a nodes or network object*

---

**Description**

Methods to create a data frame from an object of class nodes.goldfish (see [defineNodes\(\)](#)) or a matrix from an object of class network.goldfish (see [defineNetwork\(\)](#)) with the attributes or the network ties updated according with the events linked to the object using the [linkEvents\(\)](#) function.

**Usage**

```
## S3 method for class 'nodes.goldfish'
as.data.frame(x, ..., time = -Inf, startTime = -Inf)

## S3 method for class 'network.goldfish'
as.matrix(x, ..., time = -Inf, startTime = -Inf)
```

**Arguments**

x	an object of class <code>nodes.goldfish</code> for <code>as.data.frame()</code> method or <code>network.goldfish</code> for <code>as.matrix()</code> method.
...	Not further arguments are required.
time	a numeric value or a calendar date value (see <a href="#">as.Date()</a> ) to update the state of the object x until this time value (event time < time).
startTime	a numeric <code>as.Date</code> format value; prior events are disregarded.

**Value**

The respective object updated accordingly to the events link to it. For `nodes.goldfish` object the attributes are updated according to the events linked to them. For `network.goldfish` object the network ties are updated according to the events linked to it.

**See Also**

[defineNetwork\(\)](#), [defineNodes\(\)](#), [linkEvents\(\)](#)

**Examples**

```
data("Fisheries_Treaties_6070")
states <- defineNodes(states)
states <- linkEvents(states, sovchanges, attribute = "present")
states <- linkEvents(states, regchanges, attribute = "regime")
states <- linkEvents(states, gdpchanges, attribute = "gdp")

bilatnet <- defineNetwork(bilatnet, nodes = states, directed = FALSE)
bilatnet <- linkEvents(bilatnet, bilatchanges, nodes = states)

updateStates <- as.data.frame(states,
                              time = as.numeric(as.POSIXct("1965-12-31")))

updateNet <- as.matrix(bilatnet, time = as.numeric(as.POSIXct("1965-12-31")))
```

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