

# Package: gretel (via r-universe)

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**Title** Generalized Path Analysis for Social Networks

**Version** 0.0.1.9000

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**Description** The social network literature features numerous methods for assigning value to paths as a function of their ties. 'gretel' systemizes these approaches, casting them as instances of a generalized path value function indexed by a penalty parameter. The package also calculates probabilistic path value and identifies optimal paths in either value framework. Finally, proximity matrices can be generated in these frameworks that capture high-order connections overlooked in primitive adjacency sociomatrices. Novel methods are described in Buch (2019)

<<https://davidbuch.github.io/analyzing-networks-with-gretel.html>>.

More traditional methods are also implemented, as described in Yang, Knoke (2001) <[doi:10.1016/S0378-8733\(01\)00043-0](https://doi.org/10.1016/S0378-8733(01)00043-0)>.

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**URL** <https://github.com/davidbuch/gretel>

**BugReports** <https://github.com/davidbuch/gretel/issues>

**License** GPL-3

**Depends** R (>= 3.0)

**Imports** Rcpp (>= 1.0.0), ResistorArray (>= 1.0-32)

**LinkingTo** Rcpp

**Suggests** knitr, rmarkdown, testthat (>= 2.1.0)

**VignetteBuilder** knitr

**LazyData** true

**RoxygenNote** 6.1.1

**Repository** <https://davidbuch.r-universe.dev>

**RemoteUrl** <https://github.com/davidbuch/gretel>

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all_opt_gpv	<i>Optimize All Generalized Path Values</i>
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### Description

Identify the path of optimal generalized path value from every source to every target in `sociomatrix`.

### Usage

```
all_opt_gpv(sociomatrix, p = Inf, node_costs = NULL)
```

### Arguments

<code>sociomatrix</code>	a nonnegative, real valued sociomatrix.
<code>p</code>	a nonnegative real number that sets the 'p-norm' parameter for generalized path value calculation.
<code>node_costs</code>	a list of costs, in order, of all nodes represented in the sociomatrix, all are assumed 0 if unspecified

### Value

All optimal paths from source to target nodes in `sociomatrix`. To minimize memory usage, paths are returned as a list of trees in Dijkstra's format. Specific paths can be unpacked with `unpack` as described in the example below.

**See Also**

[gpv](#) to calculate the value of a user-specified path, [opt\\_gpv](#) to identify the optimal path from a single source node to a single target node

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all_opt_ppv	<i>Optimize All Probabilistic Path Values</i>
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**Description**

Identify the path of optimal probabilistic path value from every source to every target in `sociomatrix`.

**Usage**

```
all_opt_ppv(sociomatrix, odds_scale = 1, odds_scale_by_node = NULL)
```

**Arguments**

`sociomatrix` a nonnegative, real valued sociomatrix.

`odds_scale` a nonnegative real number indicating the observed tie strength value that corresponds to 1-1 transmission odds

`odds_scale_by_node` sets a transfer odds scale for each node in a probabilistic path value calculation.

**Value**

All optimal paths from source to target nodes in `sociomatrix`. To minimize memory usage, paths are returned as a list of trees in Dijkstra's format. Specific paths can be unpacked with `unpack` as described in the example below.

**See Also**

[ppv](#) to calculate the value of a user-specified path, [opt\\_ppv](#) to identify the optimal path from a single source node to a single target node

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binary_distance	<i>Binary Distance of a Network Path</i>
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**Description**

Calculates the binary distance of a user-specified network path through a network, if all edges exist. Otherwise, returns `Inf` to signify infinite distance.

**Usage**

```
binary_distance(sociomatrix, path)
```

**Arguments**

sociomatrix      a nonnegative, real valued sociomatrix.  
 path              an integer vector of node indices from sociomatrix.

**Examples**

```
## Calculate binary distance along a path in a sociomatrix
binary_distance(YangKnoke01, path = c(1,2,5))

## This path doesn't exist
binary_distance(YangKnoke01, path = c(1,2,4,5))
```

---

BuchDarrah19              *Example data for gretel*

---

**Description**

A sociomatrix encoding tie strengths among five nodes

**Usage**

BuchDarrah19

**Format**

a numeric matrix with 5 rows and 5 columns

**Source**

<DOI:10.1016/j.socnet.2010.03.006>

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dijkstra\_inf              *Find the shortest L-Inf norm paths to other vertices*

---

**Description**

Find the shortest L-Inf norm paths to other vertices

**Usage**

dijkstra\_inf(dist, src)

**Arguments**

dist              A matrix of distances between nodes  
 src              An integer vertex ID

**Value**

A numeric vector, entry  $i$  of which is the vertex immediately preceding vertex  $i$  in the shortest path leading to  $i$ . Full paths must be constructed recursively.

---

dijkstra_nodes	<i>Find the shortest paths to other vertices</i>
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---

**Description**

Find the shortest paths to other vertices

**Usage**

```
dijkstra_nodes(dist, src, node_costs)
```

**Arguments**

dist	A matrix of distances between nodes
src	An integer vertex ID
node_costs	a list of costs, in order, of all nodes represented in the sociomatrix, all are assumed 0 if unspecified

**Value**

A numeric vector, entry  $i$  of which is the vertex immediately preceding vertex  $i$  in the shortest path leading to  $i$ . Full paths must be constructed recursively.

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flament_average_path_length	<i>Yang and Knoke's Average Path Length</i>
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**Description**

Calculates 'APL' (Average Path Length) as defined in Yang, Knoke (2001). Called flament\_average\_path\_length in homage to A.C. Flament, who defined path length in 1963.

**Usage**

```
flament_average_path_length(sociomatrix, path)
```

**Arguments**

sociomatrix	a nonnegative, real valued sociomatrix.
path	an integer vector of node indices from sociomatrix.

**See Also**[flament\\_path\\_length](#)**Examples**

```
## Calculate 'APL' of a path in a sociomatrix
flament_average_path_length(YangKnoke01, path = c(1,2,5))

## This path doesn't exist
flament_average_path_length(YangKnoke01, path = c(1,2,4,5))
```

---

flament_path_length	<i>Flament's Path Length Measure</i>
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**Description**

Calculates path length as defined in Flament (1963). That is, sums the values of each edge in the path, if all edges exist. Otherwise, returns NA.

**Usage**

```
flament_path_length(sociomatrix, path)
```

**Arguments**

sociomatrix	a nonnegative, real valued sociomatrix.
path	an integer vector of node indices from sociomatrix.

**See Also**[flament\\_average\\_path\\_length](#)**Examples**

```
## Calculate Flament's Path Length along a path in a sociomatrix
flament_path_length(YangKnoke01, path = c(1,2,5))

## This path doesn't exist
flament_path_length(YangKnoke01, path = c(1,2,4,5))
```

---

generate\_proximities *Generate a Proximity Matrix*

---

### Description

Generates a proximity matrix in one of three modes:

'ogpv' Optimal Generalized Path Value. Entry  $i, j$  of the proximity matrix will equal the optimal 'gpv' among all paths connecting node  $i$  to node  $j$ .

'oppv' Optimal Probabilistic Path Value. Entry  $i, j$  of the proximity matrix will equal the optimal 'ppv' among all paths connecting node  $i$  to node  $j$ .

'sconductivity' Social Conductivity (Random Walk Probability). If each tie strength recorded in *sociomatrix* is taken to be analogous to the conductivity of an electrical component,  $i, j$  of the proximity matrix will equal total conductivity of all paths from node  $i$  to node  $j$ .

### Usage

```
generate_proximities(sociomatrix, mode = c("ogpv", "oppv",
    "sconductivity"), p = Inf, node_costs = NULL, odds_scale = 1,
    odds_scale_by_node = NULL)
```

### Arguments

<code>sociomatrix</code>	a nonnegative, real valued sociomatrix.
<code>mode</code>	a selection of 'ogpv', 'oppv', or 'sconductivity'
<code>p</code>	if mode is 'ogpv', determines 'p-norm' parameter for generalized path value calculation.
<code>node_costs</code>	if mode is 'ogpv', assigns transmission costs to vertices within the sociomatrix.
<code>odds_scale</code>	if mode is 'oppv', sets a global transfer odds scale for probabilistic path value calculation.
<code>odds_scale_by_node</code>	if mode is 'oppv', sets a transfer odds scale for each node in a probabilistic path value calculation.

### See Also

[gpv](#), [ppv](#)

### Examples

```
## Generate a proximity matrix in each mode
## Optimal Generalized Path Value
generate_proximities(YangKnoke01, mode = "ogpv", p = Inf, node_costs = c(1,3,3,2,1))

## Optimal Probabilistic Path Value
generate_proximities(YangKnoke01, mode = "oppv", odds_scale = 2)
```

```
## Sconductivity
generate_proximities(YangKnoke01, mode = "sconductivity")
```

---

gpv

*Generalized Path Value*


---

### Description

Calculates the generalized path value of a user-specified path through `sociomatrix`. Parameter `p` sets the p-norm used in calculation.

### Usage

```
gpv(sociomatrix, path, p = Inf, node_costs = NULL)
```

### Arguments

<code>sociomatrix</code>	a nonnegative, real valued sociomatrix.
<code>path</code>	an integer vector of node indices from <code>sociomatrix</code> .
<code>p</code>	a nonnegative real number that sets the 'p-norm' parameter for generalized path value calculation.
<code>node_costs</code>	a list of costs, in order, of all nodes represented in the sociomatrix, all are assumed 0 if unspecified

### Details

As a rule of thumb, `p` close to 0 will downweight the impact of particular tie strengths and upweight the impact of binary path length. `p` equal to infinity will recapitulate the traditional path value measure of Peay (1980) and is therefore the default. In other words, the value of a path under `p = Inf` will be the value of the weakest tie. The value of the same path under `p = 0` will be the inverse of its binary length.

### See Also

[opt\\_gpv](#) to identify the path of optimal 'gpv' between two nodes and [all\\_opt\\_gpv](#) to identify the optimal paths between all pairs of nodes. Calling [generate\\_proximities](#) with `mode = 'gpv'` returns a matrix 'gpv' values for the optimal paths between all pairs of nodes.

### Examples

```
## Calculate gpv along a path in a sociomatrix
gpv(YangKnoke01, path = c(1,2,5), p = 1)

## The same calculation, with nonzero node costs
gpv(YangKnoke01, path = c(1,2,5), p = 1, node_costs = c(1,3,3,2,1))
```



```
## This path doesn't exist
gpv(YangKnoke01, path = c(1,2,4,5), p = 0)
```

gretel

*gretel: Generalized Path Analysis for Social Networks***Description**

This package contains two categories of functions. The first category is concerned with assigning values to user specified paths, while the second identifies paths of optimal value.

**Details**

Key functions in the path value calculation category are - `gpv`, which calculates Generalized Path Value - `ppv`, which calculates Probabilistic Path Value - `binary_distance`, `peay_path_value`, `flament_path_length`, `peay_average_path_value`, and `flament_average_path_length`, which calculate path value measures described in *Yang, Knoke (2001)*. - `generate_proximities`, which generates a matrix of values representing the measures of optimal paths from each source node (row index) to each target node (column index).

Key functions in the optimal path identification category are - `opt_gpv`, which identifies the path of optimal Generalized Path Value from a particular source node to a particular target node - `opt_ppv`, which identifies the path of optimal Probabilistic Path Value from a particular source node to a particular target node - `all_opt_gpv`, which identifies the 'gpv'-optimal paths from every source node to every target node - `all_opt_ppv`, which identifies the 'ppv'-optimal paths from every source node to every target node - `unpack`, which unpacks the Dijkstra-format encoded shortest paths returned by `all_opt_gpv` and `all_opt_ppv`. See their help pages for details.

OpsahlEtAl10

*Example data from Opsahl, Agneessens, Skvoretz (2010)***Description**

A sociomatrix encoding tie strengths among five nodes, used for examples in Opsahl, Agneessens, Skvoretz (2010) *Social Networks* 32(2010):245-251

**Usage**

```
OpsahlEtAl10
```

**Format**

a numeric matrix with 5 rows and 5 columns

**Source**

<DOI:10.1016/j.socnet.2010.03.006>

---

opt\_gpv *Optimize Generalized Path Value*

---

**Description**

Identify the path of optimal generalized path value from a source node to a target node.

**Usage**

```
opt_gpv(sociomatrix, source, target, p = Inf, node_costs = NULL)
```

**Arguments**

sociomatrix	a nonnegative, real valued sociomatrix.
source	an integer index corresponding to a node in sociomatrix
target	an integer index corresponding to a node in sociomatrix
p	a nonnegative real number that sets the 'p-norm' parameter for generalized path value calculation.
node_costs	a list of costs, in order, of all nodes represented in the sociomatrix, all are assumed 0 if unspecified

**See Also**

[gpv](#) to calculate the value of a user-specified path, [all\\_opt\\_gpv](#) to simultaneously identify the optimal paths from any source node to any target node.

---

opt\_ppv *Optimize Probabilistic Path Value*

---

**Description**

Identify the path of optimal probabilistic path value from a source node to a target node.

**Usage**

```
opt_ppv(sociomatrix, source, target, odds_scale = 1,
        odds_scale_by_node = NULL)
```

**Arguments**

sociomatrix	a nonnegative, real valued sociomatrix.
source	an integer index corresponding to a node in sociomatrix
target	an integer index corresponding to a node in sociomatrix
odds_scale	a nonnegative real number indicating the observed tie strength value that corresponds to 1-1 transmission odds
odds_scale_by_node	sets a transfer odds scale for each node in a probabilistic path value calculation.

**See Also**

[ppv](#) to calculate the value of a user-specified path, [all\\_opt\\_ppv](#) to simultaneously identify the optimal paths from any source node to any target node.

---

peay\_average\_path\_value

*Yang and Knoke's Average Path Value*

---

**Description**

Calculates 'APV' (Average Path Value) as defined in Yang, Knoke (2001) Called `peay_average_path_value` in homage to E.R. Peay, who defined path length in 1980.

**Usage**

```
peay_average_path_value(sociomatrix, path)
```

**Arguments**

`sociomatrix` a nonnegative, real valued sociomatrix.  
`path` an integer vector of node indices from `sociomatrix`.

**See Also**

[peay\\_path\\_value](#)

**Examples**

```
## Calculate 'APV' of a path in a sociomatrix
peay_average_path_value(YangKnoke01, path = c(1,2,5))

## This path doesn't exist
peay_average_path_value(YangKnoke01, path = c(1,2,4,5))
```

---

peay\_path\_value

*Peay's Path Value Measure*

---

**Description**

Calculates path value as defined in Peay (1980). That is, returns the value of the weakest connection in the path, if all edges exist. Otherwise, returns 0.

**Usage**

```
peay_path_value(sociomatrix, path)
```

**Arguments**

`sociomatrix` a nonnegative, real valued sociomatrix.  
`path` an integer vector of node indices from `sociomatrix`.

**See Also**

[peay\\_average\\_path\\_value](#)

**Examples**

```
## Calculate Peay's Path Value along a path in a sociomatrix
peay_path_value(YangKnoke01, path = c(1,2,5))

## This path doesn't exist
peay_path_value(YangKnoke01, path = c(1,2,4,5))
```

---

ppv

*Calculate probabilistic path value*

---

**Description**

Given a real valued sociomatrix, a path, and an optional `odds_scale`, `ppv` calculates the transmission odds for the path and returns the transmission odds times `odds_scale` so the result can be directly compared with observed tie strenghts.

**Usage**

```
ppv(sociomatrix, path, odds_scale = 1, odds_scale_by_node = NULL)
```

**Arguments**

`sociomatrix` a nonnegative, real valued sociomatrix.  
`path` an integer vector of node indices from `sociomatrix`.  
`odds_scale` a nonnegative real number indicating the observed tie strength value that corresponds to 1-1 transmission odds  
`odds_scale_by_node` sets a transfer odds scale for each node in a probabilistic path value calculation.

**Details**

We assume that observed tie strengths in `sociomatrix` are linearly proportional to transmission odds. That is, if the transmission odds for a strength 1 tie are 1 to 1, the transmission odds for a strength 5 tie are 1 to 5.

**See Also**

[opt\\_ppv](#) to identify the path of optimal 'ppv' between two nodes and [all\\_opt\\_ppv](#) to identify the optimal paths between all pairs of nodes. Calling [generate\\_proximities](#) with mode = 'ppv' returns a matrix 'ppv' values for the optimal paths between all pairs of nodes.

**Examples**

```
## Calculate ppv along a path in a sociomatrix
ppv(YangKnoke01, path = c(1,2,5), odds_scale = 3)

## This path doesn't exist
gpv(YangKnoke01, path = c(1,2,4,5))
```

---

unpack

*Unpacks a Path from a Dijkstra-Format Spanning Tree*

---

**Description**

Used with [all\\_opt\\_gpv](#) and [all\\_opt\\_ppv](#) to unpack individual paths from the Dijkstra-format trees that those functions return.

**Usage**

```
unpack(tree, source, target)
```

**Arguments**

tree	a Dijkstra-format tree returned by <a href="#">all_opt_gpv</a> or <a href="#">all_opt_ppv</a>
source	an integer index corresponding to a node in sociomatrix
target	an integer index corresponding to a node in sociomatrix

**Details**

Returns NA if a path does not exist

---

YangKnoke01

*Example data from Yang, Knoke (2001)*

---

**Description**

A sociomatrix encoding tie strengths among five nodes, used for examples in Yang, S., Knoke, D. (2001) *Social Networks* 23(4):285-295

**Usage**

YangKnoke01

**Format**

a numeric matrix with 5 rows and 5 columns

**Source**

<DOI: 10.1016/S0378-8733(01)00043-0>

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