# Package 'grandpa'

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Type Package
Title GeneRAtive Networks using Property and Degree Augmentation
Version 0.1.0
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<b>Description</b> This packages generates graphs from existing graphs using both observed attribute structure and topological augmentation
License MIT
<b>Depends</b> R ( $i = 3.5.0$ )
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addAugment

Add degree augmentation label

## Description

This function is typically called by the grandpa function rather than directly.

## Usage

```
addAugment(G, nbins, degType)
```

## Arguments

G an igraph object

nbins the number of degree bins used in the degree augmentation

degType string; the type of degree to be used in the degree augmentation. See

?degree for details

#### Value

an igraph object containing an augmentation label corresponding to degree

## Examples

```
G <- make_graph("Zachary")
G <- addAugment(G,3,"all")</pre>
```

addCommunityAugment

Add community augmentation label

## Description

This function is typically called by the grandpa function rather than directly. By default, this function implements louvain clustering to detect community membership. Note, it is recommended to fit a community algorithm outside of the GRANDPA framework and add a vertext label called 'CommunityLabel' for best results.

## Usage

```
addCommunityAugment(G)
```

## Arguments

G an igraph object

## Value

an igraph object containing an augmentation label corresponding to community membership

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## Examples

```
G <- make_graph("Zachary")
G <- addCommunityAugment(G)</pre>
```

calcObsProbs

 $Calculates\ the\ observed\ edge\ and\ vertex\ probabilities$ 

## Description

This function is typically called by the grandpa function rather than directly.

#### Usage

```
calcObsProbs(G)
```

## Arguments

G

An igraph object, containing labels indicated with the word 'Label' that should be used to calculate edge and vertex probabilities

## Value

a list of tidy tables containing the vertex probabilities and edge probabilities respectively

## Examples

```
G = make_graph("Zachary")
V(G)$degree<-degree(G,mode="all")

V(G)$Label1<-rep("A",vcount(G))
V(G)$Label1[V(G)$degree>median(V(G)$degree)]<-"B"

obsProbTabs<-calcObsProbs(G)
vertexProbs<-obsProbTabs[[1]]
edgeProbs<-obsProbTabs[[2]]</pre>
```

createMap

Creates the map which lists vertix IDs based on attribute labels

## Description

This function is typically called by the grandpa function rather than directly.

## Usage

```
createMap(sampled_v)
```

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## Arguments

 $sampled\_v \qquad \quad \text{a dataframe containing the sampled vertices and their corresponding la-}$ 

bels on each row

## Value

a list which contains a mapping of all vertex IDs based on label combinations

## Examples

```
G = make_graph("Zachary")
V(G)$degree<-degree(G,mode="all")

V(G)$Label1<-rep("A",vcount(G))
V(G)$Label1[V(G)$degree>median(V(G)$degree)]<-"B"

obsProbTabs<-calcObsProbs(G)
vertexProbs<-obsProbTabs[[1]]
edgeProbs<-obsProbTabs[[2]]

sampledV<-sampleVertices(vertexProbs,50,0)
map<-createMap(sampledV)</pre>
```

degree\_error

Calculates the error between two degree distributions of graphs

## Description

This function takes an original graph and a simulated graph and calculates the normalized root mean square error between their degree distributions.

## Usage

```
degree_error(Gs, Gt)
```

## Arguments

 $\begin{array}{ll} \text{Gs} & \text{the original graph} \\ \text{Gt} & \text{the generated graph} \end{array}$ 

## Value

numerical; the NRMSE between the degree distributions  $\,$ 

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grandpa

Conducts a GRANPDA procedure

#### Description

Uses the GRANPDA framework to generate a network with matching attribute probabilities

## Usage

```
grandpa(
  G,
  nt = vcount(G),
  mt = ecount(G),
  preventSelf = T,
  preventDups = T,
  augment = F,
  augmentCommunity = F,
  nbins = 3,
  degType = "all",
  seed = 0
)
```

## Arguments

G an igraph object containing the original graph. Attributes which will be in the modelling procedure must contain the word 'Label' nt the number of desired nodes in the generated graph the number of desired edges in the generated graph mt preventSelf logical; should self connections be prevented? logical; should duplicate connections be prevented? preventDups logical; should degree augmentation occur? augment augmentCommunity logical; should communited be augmented? nbins the number of degree bins used in the degree augmentation string; the type of degree to be used in the degree augmentation. See degType ?degree for details a random seed set for reproducibility

#### Value

an igraph object with attribute and augmentation labels

## Examples

seed

```
G = make_graph("Zachary")
V(G)$degree<-degree(G,mode="all")
V(G)$Label1<-rep("A",vcount(G))
```

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```
V(G)$Label1[V(G)$degree>median(V(G)$degree)]<-"B"
GSim<-grandpa(G)</pre>
```

initializeTargetG

Intializes an empty graph with the generated vertices

## Description

This function is typically called by the grandpa function rather than directly.

## Usage

```
initializeTargetG(sampled_v, nt, d)
```

## Arguments

sampled\_v a dataframe containing the sampled vertices and their corresponding la-

bels on each row

nt the number of desired nodes in the generated graph

d logical; is the graph directed?

## Value

an empty igraph object with the desired vertices

## Examples

```
G = make_graph("Zachary")
V(G)$degree<-degree(G,mode="all")

V(G)$Label1<-rep("A",vcount(G))
V(G)$Label1[V(G)$degree>median(V(G)$degree)]<-"B"

obsProbTabs<-calcObsProbs(G)
vertexProbs<-obsProbTabs[[1]]
edgeProbs<-obsProbTabs[[2]]

sampledV<-sampleVertices(vertexProbs,50,0)
initG<-initializeTargetG(sampledV,50,FALSE)</pre>
```

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Simulated physician network data
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## Description

Graph simulated from medicare claims wherein physicians are nodes, and edges between nodes indicate overlapping patients. Physician specialty is represented as a vertex attribute and shown using vertex color by default.

## Usage

```
data(physNet)
```

## **Format**

```
An object of class "igraph"; see igraph.
```

## Source

arXiv

## References

```
Bobak et al. (2022) arXiv preprint arXiv:2211.15000. (arXiv)
```

## Examples

```
data(physNet)
plot(physNet,vertex.label=NA)
```

```
preventDuplicateConnections
```

Prevents duplicate connections during graph generation

## Description

This function is typically called by the grandpa function and should not be used directly.

## Usage

```
preventDuplicateConnections(v1, v2, lab1, lab2, C2V, Gt)
```

## Arguments

v1	a vertex id
v2	a second vertex id
lab1	the label corresponding to vertex id $1$
lab2	the label corresponding to vertex id $2$
C2V	a mapping of label categories to vertex IDs
Gt	a graph being generated

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## Value

updated vertex IDs

## preventSelfConnections

Prevents self connections during graph generation

## Description

This function is typically called by the grandpa function and should not be used directly.

## Usage

```
preventSelfConnections(v1, v2, lab1, lab2, C2V)
```

## Arguments

v1 a vertex id

v2 a second vertex id

the label corresponding to vertex id 1
the label corresponding to vertex id 2
a mapping of label categories to vertex IDs

#### Value

updated vertex IDs

sampleEdges Creates

Creates the edges that will be used in the generated graph

## Description

This function is typically called by the grandpa function rather than directly.

## Usage

```
sampleEdges(observedEdgeLabels, mt, seed)
```

## Arguments

observedEdgeLabels

a dataframe containing the observed edge label combinations and their

probabilities from the original graph

mt the number of desired edges in the generated graph seed the random seed to be used for reproducibility

## Value

a dataframe containing edges and their nodel attributes on either end

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#### Examples

```
G = make_graph("Zachary")
V(G)$degree<-degree(G,mode="all")

V(G)$Label1<-rep("A",vcount(G))
V(G)$Label1[V(G)$degree>median(V(G)$degree)]<-"B"

obsProbTabs<-calcObsProbs(G)
vertexProbs<-obsProbTabs[[1]]
edgeProbs<-obsProbTabs[[2]]

sampledE<-sampleVertices(edgeProbs,75,0)</pre>
```

sampleVertices

Creates the vertices that will be used in the generated graph

## Description

This function is typically called by the grandpa function rather than directly.

## Usage

```
sampleVertices(observedLabels, nt, seed)
```

## Arguments

observedLabels a dataframe containing the observed label combinations and their proba-

bilities from the original graph

nt the number of desired nodes in the generated graph seed the random seed to be used for reproducibility

## Value

a dataframe containing node attributes

## Examples

```
G = make_graph("Zachary")
V(G)$degree<-degree(G,mode="all")

V(G)$Label1<-rep("A",vcount(G))
V(G)$Label1[V(G)$degree>median(V(G)$degree)]<-"B"

obsProbTabs<-calcObsProbs(G)
vertexProbs<-obsProbTabs[[1]]
edgeProbs<-obsProbTabs[[2]]

sampledV<-sampleVertices(vertexProbs,50,0)</pre>
```

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## Description

This function is typically called by the grandpa function rather than directly.

## Usage

```
simGraph(Gt, sampled_e, C2V, seed, preventSelf, preventDups)
```

## Arguments

Gt The empty graph

sampled\_e a dataframe containing the edges for the generated graph and their node

attributes

C2V a mapping of the nodal attribute combinations and vertex IDs

the random seed to be used for reproducibility
preventSelf logical; should self connections be prevented?
preventDups logical; should duplicate connections be prevented

## Value

a generated igraph object

## Examples

```
G = make_graph("Zachary")
V(G)$degree<-degree(G,mode="all")

V(G)$Label1<-rep("A",vcount(G))
V(G)$Label1[V(G)$degree>median(V(G)$degree)]<-"B"

obsProbTabs<-calcObsProbs(G)
vertexProbs<-obsProbTabs[[1]]
edgeProbs<-obsProbTabs[[2]]

sampledV<-sampleVertices(vertexProbs,50,0)
initG<-initializeTargetG(sampledV,50,FALSE)
map<-createMap(sampledV)
sampledE<-sampleVertices(edgeProbs,75,0)

GSim<-simGraph(initG,sampledE,map,0,T,T)</pre>
```

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