Package ‘locationgamer’

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Type Package
Title Identification of Location Game Equilibria in Networks
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Description Identification of equilibrium locations in location games (Hotelling (1929) <doi:10.2307/2224214>). In these games, two competing actors place customer-serving units in two locations simultaneously. Customers make the decision to visit the location that is closest to them. The functions in this package include Prim algorithm (Prim (1957) <doi:10.1002/j.1538-7305.1957.tb01515.x>) to find the minimum spanning tree connecting all network vertices, an implementation of Dijkstra algorithm (Dijkstra (1959) <doi:10.1007/BF01386390>) to find the shortest distance and path between any two vertices, a self-developed algorithm using elimination of purely dominated strategies to find the equilibrium, and several plotting functions.
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createDistance

Create distance matrix for a completely connected network

Description

Create distance matrix for a completely connected network

Usage

createDistance(coordMatrix)

Arguments

coordMatrix  A matrix containing all the x and y coordinates of the network vertexes

Value

A square matrix containing the Euclidean distances between all vertexes, assuming that the network is completely connected.

Examples

coordMatrix <- matrix(c(0,10,15,20,30,30,15,15),ncol = 2)
createDistance(coordMatrix)

dijkstra

Shortest path through network using dijkstra’s algorithm

Description

This function finds the shortest path from a starting node to an end node in a network specified by an edge matrix and vertex coordinates. Position i,j of the edge matrix is one if there is an edge between the ith and jth vertex, zero otherwise. The function returns the path NA with length infinity if the network is disconnected, i.e. if no shortest path can be found.
euclidDistance

Usage

dijkstra(edgeMatrix, coordMatrix, initialNode, endNode, nNodes)

Arguments

edgeMatrix A square matrix consisting of zeros and ones. Has to be zero on the diagonals
coordMatrix A data frame containing the x and y coordinates of each network vertex
initialNode A number corresponding to the start node/vertex
endNode A number corresponding to the end node/vertex
nNodes The number of vertices/nodes in the network

Value

A list consisting of a vector with the vertices/nodes visited by the shortest path and the length of the shortest path.

Examples

initialNode <- 1
dijkstra(edgeMatrix, coordMatrix, initialNode, endNode, nNodes)

Euclidean distance between two points

description

Euclidean distance between two points

Usage

euclidDistance(x1, y1, x2, y2)

Arguments

x1 x-coordinate of point 1
y1 y-coordinate of point 1
x2 x-coordinate of point 2
y2 y-coordinate of point 2
Value

The Euclidean distance between points 1 and 2 as a number

Description

Function finds the equilibrium locations of a location game, similar to a hotelling game. Clients choose the location closest to them.

Usage

`lgsolve(edgeMatrix, coordMatrix, nPlayers = 2, demandLoc)`

Arguments

- `edgeMatrix`: A square matrix consisting of zeros and ones. Has to be zero on the diagonals
- `coordMatrix`: A data frame containing the x and y coordinates of each network vertex
- `nPlayers`: Number of players in the location game. Default is set to 2, which is the only number of players supported right now.
- `demandLoc`: A vector containing the demand or profit at each vertex of the network

Value

A list with two components. A matrix with zeros and ones, where a one symbolizes a equilibrium location. The row index denotes the location of player 1, and the column index the location chosen by player 2. The second entry is a summary of all equilibrium locations and the payoffs for player 1 and 2.

Examples

```r
edgeMatrix <- matrix(0, nrow = 6, ncol = 6)
edgeMatrix[,1] <- c(0,1,0,0,0,0)
edgeMatrix[,2] <- c(1,0,1,0,1,0)
edgeMatrix[,3] <- c(0,1,0,0,0,0)
edgeMatrix[,4] <- c(0,0,0,0,1,0)
edgeMatrix[,5] <- c(0,1,0,1,0,1)
edgeMatrix[,6] <- c(0,0,0,0,1,0)
coordMatrix <- matrix(c(0,3,0,2,0,1,1,3,1,2,1,1), nrow = 6, ncol = 2, byrow = TRUE)
demandLoc <- c(100, 100, 100, 100, 100, 100)
lgsolve(edgeMatrix, coordMatrix, 2, demandLoc)
```
**plotDijkstra**

Plot shortest path between two points in a network

**Description**

This function plots the entire network and shortest path between two points. The parameter dijkstraPath is obtained by the function dijkstra, in which one has to specify the initial and end node of the path.

**Usage**

`plotDijkstra(edgeMatrix, coordMatrix, dijkstraPath)`

**Arguments**

- `edgeMatrix` A matrix containing zeros and ones if an edge between two vertexes is absent or not
- `coordMatrix` A data frame containing the x and y coordinates of each vertex of the network
- `dijkstraPath` A vector of numbers corresponding to the vertexes of the shortest path through the network

**Value**

Function outputs a two-dimensional plot

**Examples**

```r
edgeMatrix <- matrix(0, nrow = 4, ncol = 4)
edgeMatrix[,1] <- c(0,1,0,0)
edgeMatrix[,2] <- c(1,0,1,1)
edgeMatrix[,3] <- c(0,1,0,0)
edgeMatrix[,4] <- c(0,1,0,0)
coordMatrix <- matrix(c(0,10,15,20,30,30,15,15),ncol = 2)
dijkstraPath <- c(4,2,1)
plotDijkstra(edgeMatrix, coordMatrix, dijkstraPath)
```

**plotNetwork**

Plotting a network consisting of edges and vertexes

**Description**

Plotting a network consisting of edges and vertexes

**Usage**

`plotNetwork(edgeMatrix, coordMatrix)`
Arguments

edgeMatrix A matrix containing zeros and ones if an edge between two vertexes is absent or not
coordMatrix A data frame containing the x and y coordinates of each vertex of the network

Value

A plot of the connected network

```r
edgeMatrix <- matrix(0, nrow = 4, ncol = 4) edgeMatrix[,1] <- c(0,1,0,0) edgeMatrix[,2] <- c(1,0,1,1) edgeMatrix[,3] <- c(0,1,0,0) edgeMatrix[,4] <- c(0,1,0,0) coordMatrix <- matrix(c(0,10,15,20,30,30,15,15),ncol = 2) plotNetwork(edgeMatrix, coordMatrix)
```

Description

Plotting minimum spanning tree connecting all vertexes

Usage

```r
plotPrim(minimumSp, coordMat)
```

Arguments

minimumSp A data frame in which each row corresponds to an edge between two numbered vertexes. Use function primDistance to obtain minimum spanning tree using Prim's algorithm.
coordMat A matrix containing all the x and y coordinates of the network vertexes.

Examples

```r
minimumSp <- matrix(c(1,4,4,3,2,3),ncol = 2)
coordMatrix <- matrix(c(0,10,15,20,30,30,15,15),ncol = 2)
plotPrim(minimumSp, coordMatrix)
```
**primDistance**

*Minimum spanning tree using Prim’s algorithm*

**Description**

Minimum spanning tree using Prim’s algorithm

**Usage**

`primDistance(distMatrix)`

**Arguments**

- **distMatrix**: A square matrix containing the distances between all vertexes of a network

**Value**

A matrix with rows describing which vertex is connected to which other vertex.

**Examples**

```r
distMatrix <- matrix(c(0,10,20,30,10,0,40,60,20,40,0,30,30,60,30,0),
nrow = 4, ncol = 4, byrow = TRUE)
primDistance(distMatrix)
```

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**randomCoordinates**

*Create random coordinates for network vertexes*

**Description**

Create random coordinates for network vertexes

**Usage**

`randomCoordinates(nNodes, xMax, xMin, yMax, yMin)`

**Arguments**

- **nNodes**: The number of vertexes/ nodes in the network
- **xMax**: The maximum x-coordinate of the nodes in the network
- **xMin**: The minimum x-coordinate of the nodes in the network
- **yMax**: The maximum y-coordinate of the nodes in the network
- **yMin**: The minimum y-coordinate of the nodes in the network
randomCoordinates

Value

A data frame with dimensions nNodes x 2 containing the x and y coordinates of the network’s vertexes

Examples

nNodes <- 10
xMax <- 2000
xMin <- 0
yMax <- 3000
yMin <- 200
randomCoordinates(nNodes, xMax, xMin, yMax, yMin)
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