

# Package ‘pandemonium’

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**Title** High Dimensional Analysis in Linked Spaces

**Version** 0.2.4

**Description** A 'shiny' GUI that performs high dimensional cluster analysis.

This tool performs data preparation, clustering and visualisation within a dynamic GUI.

With interactive methods allowing the user to change settings all without having to leave the GUI.

An earlier version of this package was described in Laa and Valencia (2022) <[doi:10.1140/epjp/s13360-021-02310-1](https://doi.org/10.1140/epjp/s13360-021-02310-1)>.

**License** MIT + file LICENSE

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**Depends** R (>= 3.5)

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**Suggests** knitr, readr, rmarkdown, testthat (>= 3.0.0), VIM

**VignetteBuilder** knitr

**Config/testthat/edition** 3

**URL** <https://gabrielmccoy.github.io/pandemonium/>

**NeedsCompilation** no

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## Contents

Bikes	2
chi2score	3
getBenchmarkInformation	3
getClusterDists	4
getDists	5
makePlots	5
normCoords	7
outsidescore	7
pandemonium	8
pullCoords	9
pullCoordsNoCov	10
rawCoords	10
tSNE	11
umap	12
userCoords	12
writeResults	13
<b>Index</b>	<b>15</b>

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Bikes

*Bike sharing data with model information*

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### Description

The dataset contains daily counts of bikes rented with corresponding weather and seasonal information. The data is provided by Hadi Fanaee-T and available from <https://doi.org/10.24432/C5W894>. Additionally, model information from a single hidden layer neural network with eight nodes in the hidden layer has been added: the values of the activations for all observations (variables A1 to A8 in the cluster space) and the model prediction (pred) and residual (res) in the other variables.

### Usage

Bikes

### Format

a list of 4 dataframes

**df** dataframe 731 obs of 18 variables containing the entire bikes data set

**space1** dataframe 731 obs of 8 variables (cluster space)

**space2** dataframe 731 obs of 6 variables (linked space, predictors used in the model)

**other** dataframe 731 obs of 4 variables (other variables, including observed and predicted counts)

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chi2score	<i>Chi-squared scores function</i>
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**Description**

Can be used as getScores input in pandemonium. Returns chi-squared values as the score and sigma bins as the bins.

**Usage**

```
chi2score(space1, covinv, exp, ...)
```

**Arguments**

space1	dataframe with variables in space1
covinv	inverse covariance matrix from space1
exp	reference point from space 1
...	other expected values of getScore

**Value**

named list containing scores for use in pandemonium

**Examples**

```
chi2score(Bikes$space1, solve(cov(Bikes$space1)),  
          data.frame(value = colMeans(Bikes$space1)))
```

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getBenchmarkInformation	<i>Compute cluster information</i>
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**Description**

The returned tibble contains the id of the cluster benchmark, the cluster radius and diameter, and group number for each cluster.

**Usage**

```
getBenchmarkInformation(dmat, groups)
```

**Arguments**

dmat            distance matrix  
 groups        groups resulting from clustering

**Value**

data frame with cluster information

**Examples**

```
dists <- getDists(Bikes$space1, "euclidean")
fit <- stats::hclust(dists, "ward.D2")
groups <- stats::cutree(fit, k = 4)
getBenchmarkInformation(as.matrix(dists), groups)
```

---

getClusterDists        *Compute cluster distance summaries*

---

**Description**

The returned tibble contains the id of the cluster pairs, with benchmark distance (d1), minimum (d2) and maximum (d3) distances between any points in the two clusters.

**Usage**

```
getClusterDists(dmat, groups, benchmarks)
```

**Arguments**

dmat            distance matrix  
 groups        groups resulting from clustering  
 benchmarks    data frame with benchmark id and group number

**Value**

data frame with distance information

**Examples**

```
dists <- getDists(Bikes$space1, "euclidean")
fit <- stats::hclust(dists, "ward.D2")
groups <- stats::cutree(fit, k = 4)
bm <- getBenchmarkInformation(as.matrix(dists), groups)
getClusterDists(as.matrix(dists), groups, bm)
```

---

getDists	<i>Compute distances between all points</i>
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**Description**

Compute distances between all points

**Usage**

```
getDists(coord, metric, user_dist = NULL)
```

**Arguments**

coord	matrix with coordinate representation of all points
metric	name of distance metric to be used in stats::dist
user_dist	user distance returned with metric=user

**Value**

distances between all points

**Examples**

```
getDists(Bikes$space1[1:5,],"euclidean")  
getDists(Bikes$space1[1:5,],"maximum")
```

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makePlots	<i>Generate a specified plot outside the GUI</i>
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**Description**

An interface to generate a specific graph seen when using the GUI. Settings include: metric, linkage, k, plotType, for details see the vignette on using this function.

**Usage**

```
makePlots(  
  space1,  
  settings,  
  cov = NULL,  
  covInv = NULL,  
  exp = NULL,  
  space2 = NULL,  
  space2.cov = NULL,
```

```

    space2.covInv,
    space2.exp = NULL,
    user_dist = NULL,
    getCoordsSpace1 = normCoords,
    getCoordsSpace2 = normCoords,
    getScore = NULL
  )

```

### Arguments

space1	dataframe of variables in cluster space
settings	list specifying parameters usually selected in the app
cov	covariance matrix for space 1
covInv	inverse covariance matrix for space 1
exp	reference point in space 1
space2	dataframe of variables in linked space
space2.cov	covariance matrix for space 2
space2.covInv	inverse covariance matrix for space 2
space2.exp	reference point in space 2
user_dist	user defined distances
getCoordsSpace1	function to calculate coordinates in space 1
getCoordsSpace2	function to calculate coordinates in space 2
getScore	function to calculate scores and bins

### Value

ggplot, plotly or detourr plot depending on settings\$plotType

### Examples

```

makePlots(space1 = Bikes$space1,
  settings = list(
    plotType = "WC", x="hum", y="temp", k=4, metric="euclidean",
    linkage="ward.D2", WCa=0.5, showalpha=TRUE), cov = cov(Bikes$space1),
    space2 = Bikes$space2, getScore = outsidescore(Bikes$other$res,"Residual"))

makePlots(space1 = Bikes$space1,
  settings = list(
    plotType = "tour", k=4, metric="euclidean", linkage="ward.D2",
    tourspace="space1", colouring="clustering", out_dim=2, tour_path="grand",
    display="scatter", radial_start=NULL, radial_var=NULL, slice_width=NULL, seed = 2025),
    cov = cov(Bikes$space1), space2 = Bikes$space2,
    getScore = outsidescore(Bikes$other$res,"Residual"))

```

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normCoords	<i>Scaled coordinates</i>
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**Description**

Using scale to center and scale the coordinates.

**Usage**

```
normCoords(df, ...)
```

**Arguments**

df	data frame
...	other expected values of getCoords

**Value**

matrix with coordinate representation of all points

**Examples**

```
head(normCoords(Bikes$space2))
```

---

outsidescore	<i>Using externally computed score values</i>
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**Description**

Can be used as getScores input in pandemonium, to use score values that are computed externally. Returns scores values as the score, and bins computed as below, between or above the first and third quartile.

**Usage**

```
outsidescore(scores, scoreName = NULL)
```

**Arguments**

scores	external scores to be passed to the app.
scoreName	name for scores

**Value**

named list containing scores for use in pandemonium

**Examples**

```
pandemonium(df = Bikes$space1, space2 = Bikes$space2,
            getScore = outsidescore(Bikes$other$res,"Residual"))
```

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pandemonium

*Shiny app for exploring clustering solutions*

---

**Description**

Opening the GUI to cluster the data points based on values in space2. Coordinates and distances are computed on the fly, or can be entered in the function call.

**Usage**

```
pandemonium(
  df,
  cov = NULL,
  is.inv = FALSE,
  exp = NULL,
  space2 = NULL,
  space2.cov = NULL,
  space2.exp = NULL,
  group = NULL,
  label = NULL,
  user_dist = NULL,
  dimReduction = list(tsNE = tsNE, umap = umap),
  getCoords = list(normal = normCoords),
  getScore = NULL
)
```

**Arguments**

df	data frame of data, assumes space 1 but variables can be re-assigned in the app
cov	covariance matrix (optional)
is.inv	is the covariance matrix an inverse default FALSE
exp	observable reference value (e.g. experimental measurement)
space2	data frame assumed to be in space 2 but variables can be re-assigned in the app
space2.cov	covariance matrix (optional)
space2.exp	observable reference value (e.g. experimental measurement)
group	grouping assignments
label	point labels
user_dist	input distance matrix (optional)



dimReduction	named list of functions used for dimension reduction
getCoords	named list containing functions to calculate coordinates
getScore	named list containing functions to calculate scores to be plotted as bins and continuous value.

**Value**

No return value, called to initiate 'shiny' app

---

pullCoords	<i>Chi-Squared Loss Function Coordinates</i>
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---

**Description**

Computes coordinate values by comparing observed values to the reference, using the covariance matrix as when computing the chi-squared loss.

**Usage**

```
pullCoords(df, covInv, exp, ...)
```

**Arguments**

df	data frame
covInv	inverse covariance matrix
exp	reference values
...	other expected values of getCoords

**Value**

matrix with coordinate representation of all points

**Examples**

```
head(pullCoords(Bikes$space2, solve(cov(Bikes$space2)),
  data.frame(value = colMeans(Bikes$space2))))
```

---

pullCoordsNoCov	<i>Generic Loss Function Coordinates</i>
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**Description**

Coordinates are computed as centered by the reference value and scaled with the standard deviation. Uses the  $i$ , $i$ th entry of the covariance matrix as the standard deviation of the  $i$ th variable.

**Usage**

```
pullCoordsNoCov(df, cov, exp, ...)
```

**Arguments**

df	data frame
cov	covariance matrix
exp	reference values
...	other expected values of getCoords

**Value**

matrix with coordinate representation of all points

**Examples**

```
head(pullCoordsNoCov(Bikes$space2, cov(Bikes$space2),
  data.frame(value = colMeans(Bikes$space2))))
```

---

rawCoords	<i>Raw coordinates</i>
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**Description**

Returns the input data frame. This is used when other coordinate computations fail. In general, scaling of the inputs is recommended before clustering.

**Usage**

```
rawCoords(df, ...)
```

**Arguments**

df	data frame
...	other expected values of getCoords

**Details**

Externally calculated coordinates can be used through `userCoords` or as input data with `rawCoords` used as the coordinate function. The use of `userCoords` over `rawCoords` is in the treatment of input data. As `pandemonium` displays the input data in many plots the use of coordinates as input data will result in these plots being less meaningful for interpretation. Use `userCoords` where coordinates are necessary to calculate distances but interpretation from plots of clustering space is necessary.

**Value**

matrix with coordinate representation of all points

**Examples**

```
head(rawCoords(Bikes$space2))
```

---

tSNE

*t-Distributed Stochastic Neighbor Embedding*

---

**Description**

Computes non-linear dimension reduction with `Rtsne` and default parameters.

**Usage**

```
tSNE(dist, ...)
```

**Arguments**

`dist` a distance matrix  
`...` other parameters expected to be passed to `dimReduction`

**Value**

list containing a  $n \times 2$  matrix of reduced dimension data in `Y`

**Examples**

```
head(tSNE(getDists(Bikes$space1,"euclidean"))$Y)
```

---

 umap

*Uniform Manifold Approximation and Projection Embedding*


---

**Description**

Computes non-linear dimension reduction with uwot and default parameters.

**Usage**

```
umap(dist, ...)
```

**Arguments**

<code>dist</code>	a distance matrix
<code>...</code>	other parameters expected to be passed to <code>dimReduction</code>

**Value**

list containing a 2 x n matrix of reduced dimension data

**Examples**

```
head(umap(getDists(Bikes$space1, "euclidean"))$Y)
```

---

 userCoords

*User defined coordinate function*


---

**Description**

Allows the use of externally calculated coordinates in the app. Can only be used when variables are not reassigned between the two spaces.

**Usage**

```
userCoords(user_coords)
```

**Arguments**

<code>user_coords</code>	coordinate matrix the size of the space it will be used on
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## Details

Externally calculated coordinates can be used through userCoords or as input data with rawCoords used as the coordinate function. The use of userCoords over rawCoords is in the treatment of input data. As pandemonium displays the input data in many plots the use of coordinates as input data will result in these plots being less meaningful for interpretation. Use userCoords where coordinates are necessary to calculate distances but interpretation from plots of clustering space is necessary.

## Value

function that returns the user defined coordinates user\_coords

## Examples

```
pandemonium(df = Bikes$space1, space2 = Bikes$space2,
            coords = list(normalised = normCoords, space2 = userCoords(Bikes$space2)))
```

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writeResults

*Write coordinates and cluster assignment to a CSV file*

---

## Description

For working with the results outside the app. Settings used: metric, linkage, k

## Usage

```
writeResults(
  space1,
  cov = NULL,
  covInv = NULL,
  exp = NULL,
  space2,
  space2.cov = NULL,
  space2.covInv = NULL,
  space2.exp = NULL,
  settings,
  filename,
  user_dist = NULL,
  getCoords.space1 = normCoords,
  getCoords.space2 = rawCoords
)
```

**Arguments**

space1	cluster space matrix
cov	covariance matrix
covInv	inverse covariance matrix
exp	observable reference value (e.g. experimental measurement)
space2	space2 matrix
space2.cov	covariance matrix
space2.covInv	inverse covariance matrix
space2.exp	observable reference value (e.g. experimental measurement)
settings	list specifying parameters usually selected in the app
filename	path to write the results file to
user_dist	input distance matrix (optional)
getCoords.space1	function to calculate coordinates on clustering space
getCoords.space2	function to calculate coordinates on linked space

**Value**

No return value, called for writing file

**Examples**

```
file<-tempfile()
writeResults(space1 = Bikes$space1, space2 = Bikes$space2,
settings = list(metric="euclidean",linkage="ward.D2",k=4), filename = file)
file.remove(file)
```

# Index

## \* datasets

Bikes, [2](#)

Bikes, [2](#)

chi2score, [3](#)

getBenchmarkInformation, [3](#)

getClusterDists, [4](#)

getDists, [5](#)

makePlots, [5](#)

normCoords, [7](#)

outsidescore, [7](#)

pandemonium, [8](#)

pullCoords, [9](#)

pullCoordsNoCov, [10](#)

rawCoords, [10](#)

tSNE, [11](#)

umap, [12](#)

userCoords, [12](#)

writeResults, [13](#)