

# Package ‘gR2’

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**Type** Package

**Title** Generalized R Square Measures for a Mixture of Bivariate Linear Dependences

**Version** 1.3.0

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**Description** This package implements the estimation and inference of generalized R square.

**License** GPL-2

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.0

**LinkingTo** Rcpp,RcppArmadillo,RcppParallel, dqrng,sitmo

**Imports** Rcpp,RcppParallel, parallel,mvtnorm

**SystemRequirements** GNU make

## R topics documented:

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<i>gR2</i>	<i>gR2</i>
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## Description

*gR2* calculates the sample *gR2* under the specified scenario, the unspecified scenario (K chosen), and the unspecified scenario (K not chosen). It also provides an option to perform statistical inference on the population *gR2*.

## Usage

```
gR2(
  x,
  y,
  z = NULL,
  K = NULL,
  cand.Ks = 1:4,
  nstart = 30,
  mc.cores = NULL,
  regressionMethod = "MA",
  inference = FALSE,
  conf.level = 0.95,
  method = "general"
)
```

## Arguments

<b>x</b>	A numeric vector.
<b>y</b>	A numeric vector of the same length as <b>x</b> .
<b>z</b>	A vector of integers that represents the line membership of all the data points. Must be of the same length as <b>x</b> and <b>y</b> . Default is <b>NULL</b> .
<b>K</b>	Number of lines in the unspecified scenario. Default is <b>NULL</b> .
<b>cand.Ks</b>	A vector of positive integers that represents the candidate K's in the unspecified scenario. Default is <b>1:4</b> .
<b>nstart</b>	Number of initializations for the K-lines algorithm in the unspecified scenario. Default is <b>30</b> .
<b>mc.cores</b>	Number of cores to use in the unspecified scenario. Default is <b>NULL</b> , which means all of the available cores will be used.
<b>regressionMethod</b>	Valid values are 'MA' and 'LM'. Indicates which regression method to use in the K-lines algorithm - major axis regression or linear regression. Default is 'MA'.
<b>inference</b>	Logical. If <b>TRUE</b> , then a confidence interval for the population gR2 of confidence level <b>conf.level</b> will be calculated. Also will be calculated is a p-value of the hypothesis test where the null hypothesis is that the population gR2 is 0 and the alternative hypothesis is that the population gR2 is greater than 0. Default is <b>FALSE</b> .
<b>conf.level</b>	The confidence level of the confidence interval. See description of <b>inference</b> . Default is <b>0.95</b> .
<b>method</b>	Valid values are 'general' and 'binorm'. Indicates which asymptotic distribution of the sample gR2 to use for inference. Default is 'general'.

## Details

The arguments that require user input are **x** and **y**, which must be numeric vectors of the same length.

There are three broad types of scenarios: the specified scenario, the unspecified scenario (K chosen), and the unspecified scenario (K not chosen). The specified scenario is considered when **z** is provided; the unspecified scenario (K chosen) is considered when **z** is not provided but **K** is provided; and the unspecified scenario (K not chosen) is considered when neither **z** or **K** is provided.

In the unspecified scenario (K chosen), we recommend that users set **K** to be less than or equal to 4 for interpretability.

In the unspecified scenario (K not chosen), the **gR2** function will automatically choose a **K** value from **cand.Ks** using the Akaike information criterion (AIC). Two plots will be outputted: (1) a scree plot that shows how average squared perpendicular/vertical distance changes with the candidate **K**, and (2), a plot that shows how AIC changes with the candidate **K**. Users can decide whether the **K** value chosen by the **gR2** function is reasonable by checking these two plots.

## Value

**gR2** returns a list consisting of one or more of the following items:

<b>estimate</b>	The sample gR2.
<b>conf.level</b>	The confidence level of the confidence interval (if <b>inference</b> is <b>TRUE</b> ).
<b>conf.int</b>	The confidence interval for the population gR2 (if <b>inference</b> is <b>TRUE</b> ).
<b>p.val</b>	The p-value of the hypothesis test where the null hypothesis is that the population gR2 is 0 and the alternative hypothesis is that the population gR2 is greater than 0 (if <b>inference</b> is <b>TRUE</b> ).
<b>K</b>	The number of lines in the unspecified scenario, either chosen by the user or chosen from <b>cand.Ks</b> by the <b>gR2</b> function.
<b>membership</b>	The inferred line membership of all the data points in the unspecified scenario.

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## References

Li, J.J., Tong, X., and Bickel, P.J. (2019). Generalized R2 Measures for a Mixture of Bivariate Linear Dependences. arXiv.