

Package ‘qcv’

October 13, 2022

Title Quantifying Construct Validity

Version 1.0

Author R. Michael Furr [aut, cre],
Sarah Heuckeroth [aut]

Maintainer R. Michael Furr <FurrRM@wfu.edu>

Date 2018-11-26

Description Primarily, the 'qcv' package computes key indices related to the Quantifying Construct Validity procedure (QCV; Westen & Rosenthal, 2003 <[doi:10.1037/0022-3514.84.3.608](https://doi.org/10.1037/0022-3514.84.3.608)>; see also Furr & Heuckeroth, in press). The qcv() function is the heart of the 'qcv' package, but additional functions in the package provide useful ancillary information related to the QCV procedure.

Depends R (>= 3.5.1.0)

Imports graphics, stats

License GPL (>= 2)

Encoding UTF-8

LazyData true

RoxygenNote 6.1.1

NeedsCompilation no

Repository CRAN

Date/Publication 2018-12-17 23:30:07 UTC

R topics documented:

actr	2
medr	2
motdat	3
plotqcv	4
print.qcv	4
qcv	5
ralertingp	6
Index	8

actr	<i>Actual correlations</i>
------	----------------------------

Description

A very simple function, `actr()` computes the actual correlations between a focal test and a set of criterion variables

Usage

```
actr(df)
```

Arguments

`df` data frame consisting of focal test and a set of criterion variables

Details

The `df` should contain only scores on the focal test and on the criterion variables. Any additional columns should be deleted. Focal test scores should be in the first column. If you need to reorder columns, you can do so by column number - eg, `df <- df[c(1,3,2)]` - or by column name - eg, `df <- df[c("test", "crit1", "crit2")]`

This function uses the "pairwise.complete.obs" option, which handles missing data by pairwise deletion.

Examples

```
data(motdat)
actr(motdat)
```

medr	<i>Median r</i>
------	-----------------

Description

Computes the median intercorrelation among a set of variables

Usage

```
medr(df)
```

Arguments

`df` data frame consisting only of variables for which a median intercorrelation is desired

Details

The data frame should contain only the variables for which a median intercorrelation is desired. For Westen and Rosenthal's (2003) QCV procedure, the data frame should contain only participants' scores on the criterion variables.

This procedure handles missing data via the "pairwise.complete.obs" option.

Examples

```
data(motdat)
motdatc <- motdat[,2:13] #To retain only the criterion variables (dropping the focal test score)
medr(motdatc)
```

motdat	<i>Example data used in Furr & Heuckeroth, on Impression Motivation (IM)</i>
--------	--

Description

Self-report data from 90 participants, responding to the IM scale and 12 criterion variables.

Usage

```
data(motdat)
```

Format

A data frame of 90 rows and 13 variables:

imscale Impression Motivation
dep Dependence
mach Machiavellianism
dis Distrust
res Resourcefulness
se Self-efficacy
ext Extraversion
agr Agreeableness
comp Complexity
psc Public self-consciousness
sm Self-monitoring
anx Anxiety
nfb Need to belong

Examples

```
data(motdat)
```

plotqcv

Plot for Quantifying Construct Validity

Description

Produces a plot of actual and predicted correlations

Usage

```
plotqcv(actr, predr, labels)
```

Arguments

actr	vector of actual validity correlations
predr	vector of predicted validity correlations (in same order as actr)
labels	vector of text values that are the labels of the QCV criterion variables (in order of actr and predr)

Details

Important: When entering values for the three arguments, order them identically. For example, if the "actr" values are ordered in terms of c(criterion A, criterion B, criterion C), then the values in "predr" and the labels in "labels" should be placed in that same order.

Examples

```
actrIM <- c(.46, .13, -.24, -.03, .12, .03, .39, .06, .51, .08, .24, .66)
predrIM <- c(.58, .24, -.04, .06, -.04, .18, .36, .08, .64, .56, .36, .56)
labelsIM <- c("Dep", "Mach", "Dis", "Res", "SE", "Ext", "Agr", "Comp", "PSC", "SM", "Anx", "NTB")
plotqcv(actr=actrIM, predr=predrIM, labels=labelsIM)
```

print.qcv

Printing Quantifying Construct Validity

Description

Prints key results from the qcv() function

Usage

```
## S3 method for class 'qcv'
print(x, ...)
```

Arguments

x	object of class "qcv" (i.e., output from the qcv() function)
...	More arguments to pass to the print function.

Examples

```
actrIM <- c(.46, .13, -.24, -.03, .12, .03, .39, .06, .51, .08, .24, .66)
predrIM <- c(.58, .24, -.04, .06, -.04, .18, .36, .08, .64, .56, .36, .56)
imqcvout <- qcv(n=90, actr=actrIM, predr=predrIM, medr=.075)
print(imqcvout)
```

qcv

*Quantifying Construct Validity***Description**

Computes key indices related to the Quantifying Construct Validity (QCV) procedure (Westen & Rosenthal, 2003; see also Furr & Heuckeroth, in prep.)

Usage

```
qcv(n, actr, predr, medr)
```

Arguments

n	sample size
actr	vector of actual validity correlations
predr	vector of predicted validity correlations (in same order as actr)
medr	median intercorrelation among criterion variables

Details

This function applies procedures outlined by Westen and Rosenthal (2003; see also Furr & Heuckeroth, in prep).

Important: When entering values for the "actr" and "predr" arguments, order them identically. For example, if the "actr" values are ordered in terms of c(criterion A, criterion B, criterion C), then the values in "predr" should be placed in that same order. Otherwise all results will be incorrect.

Note that extreme zcontrast values (e.g., $z > 38$) create problems. The rcontrast-CV effect size is computed by converting z to p to t to r (see Westen & Rosenthal, Appendix A & B). However, R (and most other packages) doesn't have precision to convert an extremely large z (e.g., $Z = 38$) to a p value. For such z values, the p is converted to exactly zero. This makes it impossible to obtain t and rcontrast values. The qcv() function handles this by: a) identifying cases where p is initially exactly zero, and b) re-setting p to the smallest value possible, given the machine on which R is running (usually $2.225074e-308$). It then proceeds to compute a t and rcontrast from that adjusted p value. It also prints a note stating that the adjustment has been made and that the p, t, and rcontrast values are lower-bound approximations.

Value

ralerting-CV The ralerting-CV effect size

rcontrast-CV The rcontrast-CV effect size

zcontrast The zcontrast value

p p values associated with Zcontrast

N Sample size

k Number of criterion variables

szr Standard deviation of (z-transformed) actual correlations

medr Median intercorrelation between the criterion variables

rbarsq Mean of the squared actual correlations (between focal test and criterion variables)

t t-value associated with p value

rem "Remarkableness" of size of contrast

Additional values are returned in a list, but not printed

References

Westen, D., & Rosenthal, R. (2003). Quantifying construct validity: Two simple measures. *Journal of Personality and Social Psychology*, 84, 608-618.

Furr, R. M., & Heukeroth, S. (In prep). Advancing the Interpretation and Computation of the "Quantifying Construct Validity" Procedure

Examples

```
actrIM <- c(.46, .13, -.24, -.03, .12, .03, .39, .06, .51, .08, .24, .66)
predrIM <- c(.58, .24, -.04, .06, -.04, .18, .36, .08, .64, .56, .36, .56)
qcv(n=90, actr=actrIM, predr=predrIM, medr=.075)
```

 ralertingp

Inferential test for ralerting-CV

Description

Computes a p value for the ralerting-CV effect size

Usage

```
ralertingp(actr, predr, iter)
```

Arguments

actr vector of actual validity correlations

predr vector of predicted validity correlations (in same order as actr)

iter the number of randomization samples to be generated

Details

An inferential test for ralerting-CV, not presented in the original Westen and Rosenthal (2003). The inferential test is based upon randomization procedures. That is, given a set of predicted correlations and a set of actual correlations, ralertingp() randomly pairs values from the two sets, computes an ralerting-CV value, and repeats this process many times (as determined by the user). The actual ralerting-CV value is then compared to the distribution of ralerting-CV values derived from the randomization process. The proportion of values from that distribution that are greater than the actual ralerting-CV value is then taken as a p value.

Important: When entering values for the "actr" and "predr" arguments, order them identically. For example, if the "actr" values are ordered in terms of c(criterion A, criterion B, criterion C), then the values in "predr" should be placed in that same order. Otherwise results will be incorrect.

For relatively small values of k (the number of criterion variables), it is recommended to increase iter (the number of randomization samples). For low-k situations, a small number of randomization samples can produce unstable p values. A larger number of randomization samples would produce more stable results.

It is possible that the randomization process produces no values that are greater than the actual ralerting-CV value. In such cases, ralertingp() reports the p value as $< 1/\text{iter}$.

Examples

```
actrIM <- c(.46, .13, -.24, -.03, .12, .03, .39, .06, .51, .08, .24, .66)
predrIM <- c(.58, .24, -.04, .06, -.04, .18, .36, .08, .64, .56, .36, .56)
ralertingp(actr=actrIM, predr=predrIM, iter=1000)
```

Index

* **datasets**

motdat, 3

actr, 2

medr, 2

motdat, 3

plotqcv, 4

print.qcv, 4

qcv, 5

ralertingp, 6