

# Package ‘qcv’

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**Title** Quantifying Construct Validity

**Version** 1.0

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**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

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

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**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> |
|------|-----------------|

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**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 &amp; Heuckeroth, on Impression Motivation (IM)</i> |
|--------|----------------------------------------------------------------------------------|

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## 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  $r$ contrast 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  $r$ contrast from that adjusted  $p$  value. It also prints a note stating that the adjustment has been made and that the  $p$ ,  $t$ , and  $r$ contrast 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)
```

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|            |                                          |
|------------|------------------------------------------|
| ralertingp | <i>Inferential test for ralerting-CV</i> |
|------------|------------------------------------------|

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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)
```

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