

Package ‘profileR’

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Title Profile Analysis of Multivariate Data in R

Type Package

Description A suite of multivariate methods and data visualization tools to implement profile analysis and cross-validation techniques described in Davison & Davenport (2002) <DOI:10.1037/1082-989X.7.4.468>, Bulut (2013) <<http://conservancy.umn.edu/handle/11299/155592>>, and other published and unpublished resources. The package includes routines to perform criterion-related profile analysis, profile analysis via multidimensional scaling, moderated profile analysis, profile analysis by group, and a within-person factor model to derive score profiles.

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R topics documented:

profileR-package	2
anova.critpat	3
cpa	3

EEGS	5
interest	5
IPMMc	7
leisure	7
mod_data	8
mpa	8
nutrient	9
pams	10
paos	11
pbg	12
pcv	13
plot.critpat	14
plot.prof	15
pr	15
print.critpat	16
profileplot	17
PS	18
spouse	18
summary.critpat	19
wprifm	20

Index	21
--------------	-----------

profileR-package	<i>Profile Analysis of Multivariate Data in R</i>
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Description

The package profileR provides a set of multivariate methods and data visualization tools to implement profile analysis and cross-validation techniques described in Davison & Davenport (2002), Bulut (2013), and other resources. This package includes routines to perform criterion-related profile analysis, profile analysis via multidimensional scaling, moderated profile analysis, profile analysis by group, and a within-person factor model to derive score profiles.

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References

- Bulut, O. (2013). *Between-person and within-person subscore reliability: Comparison of unidimensional and multidimensional IRT models*. (Doctoral dissertation). University of Minnesota. University of Minnesota, Minneapolis, MN. (AAT 3589000).
- Davison, M. L., & Davenport, E. C. (2002). Identifying criterion-related patterns of predictor scores using multiple regression. *Psychological Methods*, 7(4), 468-484.
- Davison, M. L., Kim, S-K., & Close, C. W. (2009). Factor analytic modeling of within person variation in score profiles. *Multivariate Behavioral Research*, 44, 668-87.

anova.critpat	<i>Anova Tables</i>
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Description

Computes an analysis of variance table for a criterion-related profile analysis

Usage

```
## S3 method for class 'critpat'
anova(object, ...)
```

Arguments

object	an object containing the results returned by a model fitting cpa.
...	additional objects of the same type.

See Also

[cpa](#)

cpa	<i>Criterion-Related Profile Analysis</i>
-----	---

Description

Implements the criterion-related profile analysis described in Davison & Davenport (2002).

Usage

```
cpa(formula, data, k = 100, na.action = "na.fail", family = "gaussian",
     weights = NULL)
```

Arguments

formula	An object of class formula of the form response ~ terms.
data	An optional data frame, list or environment containing the variables in the model.
k	Corresponds to the scalar constant and must be greater than 0. Defaults to 100.
na.action	How should missing data be handled? Function defaults to failing if missing data are present.
family	A description of the error distribution and link function to be used in the model. See family .
weights	An option vector of weights to be used in the fitting process.

Details

The `cpa` function requires two arguments: `criterion` and `predictors`. The function returns the criterion-related profile analysis described in Davison & Davenport (2002). Missing data are presently handled by specifying `na.action = "na.omit"`, which performs listwise deletion and `na.action = "na.fail"`, the default, which causes the function to fail. The following S3 generic functions are available: `summary()`, `anova()`, `print()`, and `plot()`. These functions provide a summary of the analysis (namely, R^2 and the level and pattern components); perform ANOVA of the R^2 for the pattern, the level, and the overall model; provide output similar to `lm()`, and plots the pattern effect.

Value

An object of class `criptpat` is returned, listing the following components:

- `lvl.comp` - the level component
- `pat.comp` - the pattern component
- `b` - the unstandardized regression weights
- `bstar` - the mean centered regression weights
- `xc` - the scalar constant times `bstar`
- `k` - the scale constant
- `Covpc` - the pattern effect
- `Ypred` - the predicted values
- `r2` - the proportion of variability attributed to the different components
- `F.table` - the associated F-statistic table
- `F.statistic` - the F-statistics
- `df` - the df used in the test
- `pvalue` - the p-values for the test

References

Davison, M., & Davenport, E. (2002). Identifying criterion-related patterns of predictor scores using multiple regression. *Psychological Methods*, 7(4), 468-484. DOI: 10.1037/1082-989X.7.4.468.

See Also

[pcv](#)

Examples

```
## Not run:
data(IPMmc)
mod <- cpa(R ~ A + H + S + B, data = IPMmc)
print(mod)
summary(mod)
plot(mod)
anova(mod)

## End(Not run)
```

 EEGS

Entrance Examination for Graduate Studies

Description

The EEGS is a subset of the Entrance Examination for Graduate Studies. There are three subscores in EEGS: Quantitative 1, Quantitative 2, and Verbal. In order to show the utility of subscore reliability method in this package, each subtest was separated into two parallel forms.

Format

Form1_Q1 First form of Quantitative 1

Form2_Q1 Second form of Quantitative 1

Form1_Q2 First form of Quantitative 2

Form2_Q2 Second form of Quantitative 2

Form1_V First form of Verbal

Form2_V Second form of Verbal

 interest

Fabricated cognitive, personality, and vocational interest inventory

Description

The data come from a fabricated cognitive, personality, and vocational interested inventory. This data set can be used to demonstrate regression and structural equation modeling.

Usage

interest

Format

A data frame with 250 rows and 33 variables:

gender 1 is female and 2 is male

educ Years of education

age Age, in years

vocab Vocabulary test

reading Reading comprehension

sentcomp Sentence completion

mathmtcs Mathematics

geometry Geometry

analyrea Analytical reasoning
socdom Social dominance
sociabty Sociability
stress Stress reaction
worry Worry scale
impulsve Impulsivity
thrillsk Thrill-seeking
carpentr Carpentry
forestr Forest ranger
morticin Mortician
policemn Police
fireman Fireman
salesrep Sales representative
teacher Teacher
busexec Business executive
stockbrk Stock broker
artist Artist
socworkr Social worker
truckdvr Truck driver
doctor Doctor
clergymn Clergyman
lawyer Lawyer
actor Actor
archtct Architect
landscpr Landscaper

Source

<http://psych.colorado.edu/~carey/Courses/PSYC7291/ClassDataSets.htm>

 IPMMc

Inventory of Personality and Mood Manifestation

Description

The IPMMc data frame has 6 rows and 5 columns. See Davison and Davenport (2002) for more information.

Format

This data frame contains the following columns:

A Anxiety

H Hypochondriasis

S Schizophrenia

B Bipolar Disorder

R The Neurotic versus Psychotic Criterion Variable, where Neurotic (= 1) or Psychotic (= 0)

Source

Davison, M. L., & Davenport, E. C. (2002). Identifying criterion-related patterns of predictor scores using multiple regression. *Psychological Methods*, 7(4), 468-484.

References

Davison, M. L., & Davenport, E. C. (2002). Identifying criterion-related patterns of predictor scores using multiple regression. *Psychological Methods*, 7(4), 468-484.

 leisure

Leisure Activity Rankings

Description

The leisure dataset includes leisure activity rankings for three different groups: politicians, administrators, and belly-dancers. Rankings are provided in three categories: Reading-TV, Dance-TV, and TV-Ski. See Tabachnik and Fidell (1996) for more details.

Source

Tabachnick, B. G., & Fidell, L. S. (1996). *Using multivariate statistics* (3rd ed.). New York: Harper Collins.

Examples

```
## Not run:
data(leisure)
```

```
## End(Not run)
```

mod_data	<i>Moderated profile analysis dummy data</i>
----------	--

Description

Randomly generated data to test the mpa function.

Format

This data frame contains the following columns:

dv Dependent variable
pred1 Predictor variable 1
pred2 Predictor variable 2
mod The moderator variable

Source

This data set was randomly generated to demonstrate how to use the mpa function.

See Also

[mpa](#)

mpa	<i>Moderated Profile Analysis</i>
-----	-----------------------------------

Description

Implements the moderated profile analysis approach developed by Davison & Davenport (unpublished) - EXPERIMENTAL SUPPORT

Usage

```
mpa(formula, data, moderator, k = 100, na.action = "na.fail",
    stage2 = FALSE)
```

Arguments

formula	An object of class formula of the form response ~ terms.
data	An optional data frame, list or environment containing the variables in the model.
moderator	Name of the moderator variable.
k	Corresponds to the scalar constant and must be greater than 0. Defaults to 100.
na.action	How should missing data be handled? Function defaults to failing if missing data are present.
stage2	Should stage 2 be executed regardless of stage 1 outcome? defaults to FALSE.

Details

The function returns the criterion-related moderated profile analysis described in Davison & Davenport (unpublished). Missing data are presently handled by specifying `na.action = "na.omit"`, which performs listwise deletion and `na.action = "na.fail"`, the default, which causes the function to fail. The following S3 generic functions are not yet available but will be in future implementations. `summary()`, `anova()`, `print()`, and `plot()`. These functions provide a summary of the analysis (namely, R2 and the level and pattern components); perform ANOVA of the R2 for the pattern, the level, and the overall model; provide output similar to `lm()`, and plots the pattern effect.

Value

An object of class `critpat` is returned, listing the following components:

- `r2_full` - the proportion of variability attributed to the full model
- `r2_red` - the proportion of variability attributed to the reduced model
- `f_table` - the associated F-statistic table
- `F-statistic` - the F-statistics
- `df` - the dfs used in the test
- `pvalue` - the p-values for the test

References

Davison, M., & Davenport, E. (unpublished). Comparing Criterion-Related Patterns of Predictor Variables across Populations Using Moderated Regression.

See Also

[cpa](#)

Examples

```
## Not run:  
data(mod_data)  
mod <- mpa(dv ~ pred1 * mod + pred2 * mod, moderator = "mod", data = mod_data)  
  
## End(Not run)
```

nutrient

USDA Women's Health Survey

Description

In 1985, the United States Department of Agriculture (USDA) commissioned a study of women's nutrition. Nutrient intake was measured for a random sample of 737 women aged 25-50 years. Five nutritional components were measured: calcium, iron, protein, vitamin A and vitamin C.

Format

calcium Calcium amount

iron Iron amount

protein Protein amount

a Vitamin A amount

c Vitamin C amount

pams

Profile Analysis via Multidimensional Scaling

Description

The pams function implements profile analysis via multidimensional scaling as described by Davison, Davenport, and Bielinski (1995) and Davenport, Ding, and Davison (1995).

Usage

```
pams(data, dim)
```

Arguments

data	A data matrix or data frame; rows represent individuals, columns represent scores; missing scores are not allowed.
dim	Number of dimensions to be extracted from the data.

Details

The pams function computes similarity/dissimilarity indices based on Euclidean distances between the scores provided in the data, and then extracts dimensional coordinates for each score using multidimensional scaling. A weight matrix, level parameters, and fit measures are computed for each subject in the data.

Value

- `dimensional.configuration` - A matrix that provides prototypical profiles of dimensions extracted from the data.
- `weights.matrix` - A matrix that includes the subject correspondence weights for all dimensions, level parameters, and the subject fit measure which is the proportion of variance in the subject's actual profiles accounted for by the prototypical profiles.

References

- Davenport, E. C., Ding, S., & Davison, M. L. (1995). PAMS: SAS Template.
Davison, M. L., Davenport, E. C., & Bielinski, J. (1995). PAMS: SPSS Template.

See Also[cpa](#), [pr](#)**Examples**

```
## Not run:
data(PS)
result <- pams(PS[,2:4], dim=2)
result

## End(Not run)
```

paos

Profile Analysis for One Sample with Hotelling's T-Square

Description

The paos function implements profile analysis for one sample using Hotelling's T-square.

Usage

```
paos(data, scale = TRUE)
```

Arguments

data	A data matrix or data frame; rows represent individuals, columns represent variables.
scale	If TRUE (default), variables are standardized by dividing their standard deviations.

Details

The paos function runs profile analysis for one sample based on the Hotelling's T-square test and tests the two hypothesis. First, the null hypothesis that all the ratios of the variables in the data are equal to 1. After rejecting the first hypothesis, a secondary null hypothesis that all of the ratios of the variables in the data equal to one another (not necessarily equal to 1) is tested.

Value

A summary table is returned, listing the following two hypothesis:

- Hypothesis 1 - Ratios of the means of the variables over the hypothesized mean are equal to 1.
- Hypothesis 2 - All of the ratios are equal to each other.

See Also[cpa](#), [pr](#)

Examples

```
## Not run:
data(nutrient)
paos(nutrient, scale=TRUE)

## End(Not run)
```

pbg *Profile Analysis by Group: Testing Parallelism, Equal Levels, and Flatness*

Description

The pbg function implements three hypothesis tests. These tests are whether the profiles are parallel, have equal levels, and are flat across groups defined by the grouping variable. If parallelism is rejected, the other two tests are not necessary. In that case, flatness may be assessed within each group, and various within- and between-group contrasts may be analyzed.

Usage

```
pbg(data, group, original.names = FALSE, profile.plot = FALSE)
```

Arguments

data	A matrix or data frame with multiple scores; rows represent individuals, columns represent subscores. Missing subscores have to be inserted as NA.
group	A vector or data frame that indicates a grouping variable. It can be either numeric or character (e.g., male-female, A-B-C, 0-1-2). The grouping variable must have the same length of x. Missing values are not allowed in y.
original.names	Use original column names in x. If FALSE, variables are renamed using v1, v2, ..., vn for subscores and "group" for the grouping variable. Default is FALSE.
profile.plot	Print a profile plot of scores for the groups. Default is FALSE.

Value

An object of class profg is returned, listing the following components:

- data.summary - Means of observed variables by the grouping variable
- corr.table - A matrix of correlations among observed variables splitted by the grouping variable
- profile.test - Results of F-tests for testing parallel, coincidental, and level profiles across two groups.

See Also

[pr](#), [profileplot](#)

Examples

```
## Not run:
data(spouse)
mod <- pbg(data=spouse[,1:4], group=spouse[,5], original.names=TRUE, profile.plot=TRUE)
print(mod) #prints average scores in the profile across two groups
summary(mod) #prints the results of three profile by group hypothesis tests

## End(Not run)
```

pcv

Cross-Validation for Profile Analysis

Description

Implements the cross-validation described in Davison & Davenport (2002).

Usage

```
pcv(formula, data, seed = NULL, na.action = "na.fail",
     family = "gaussian", weights = NULL)
```

Arguments

formula	An object of class formula of the form response ~ terms.
data	An optional data frame, list or environment containing the variables in the model.
seed	Should a seed be set? Function defaults to a random seed.
na.action	How should missing data be handled? Function defaults to failing if missing data are present.
family	A description of the error distribution and link function to be used in the model. See family .
weights	An option vector of weights to be used in the fitting process.

Details

The pcv function requires two arguments: criterion and predictor. The criterion corresponds to the dependent variable and the predictor corresponds to the matrix of predictor variables. The function performs the cross-validation technique described in Davison & Davenport (2002) and an object of class `critpat` is returned. There the following s3 generic functions are available: `summary()`, `anova()`, `print()`, and `plot()`. These functions provide a summary of the cross-validation (namely, R²); performs ANOVA of the R² based on the split for the level, pattern, and overall; provide output similar to `lm()`; and plot the estimated parameters for the random split. Missing data are presently handled by specifying `na.action = "na.omit"`, which performs list-wise deletion and `na.action = "na.fail"`, the default, which causes the function to fail. A seed may also be set for reproducibility by setting the seed.

Value

An object of class `critpat` is returned, listing the following components:

- `R2.full`, test of the null hypothesis that $R^2 = 0$
- `R2.pat`, test that the $R^2_{\text{pattern}} = 0$
- `R2.level`, test that the $R^2_{\text{level}} = 0$
- `R2.full.lvl`, test that the $R^2_{\text{full}} = R^2_{\text{level}} = 0$
- `R2.full.pat`, test that the $R^2_{\text{full}} = R^2_{\text{pattern}} = 0$

References

Davison, M., & Davenport, E. (2002). Identifying criterion-related patterns of predictor scores using multiple regression. *Psychological Methods*, 7(4), 468-484. DOI: 10.1037/1082-989X.7.4.468.

See Also

[cpa](#), [print.critpat](#), [summary.critpat](#), [anova.critpat](#), [plot.critpat](#)

plot.critpat

Plot criterion-related profile

Description

Plots the criterion-related level and pattern profiles for each observation

Usage

```
## S3 method for class 'critpat'  
plot(x, ...)
```

Arguments

`x` `critpat` object resulting from `cpa`
`...` additional arguments affecting the plot produced.

See Also

[cpa](#)

plot.prof	<i>Plots a pattern and level reliability</i>
-----------	--

Description

Plots the pattern vs. level reliability returned from the pr function of class prof.

Usage

```
## S3 method for class 'prof'  
plot(x, ...)
```

Arguments

x	an object returned from the pr function
...	additional objects of the same type.

See Also

[pr](#)

pr	<i>Pattern and Level Reliability via Profile Analysis</i>
----	---

Description

The pr function uses subscores from two parallel test forms and computes profile reliability coefficients as described in Bulut (2013).

Usage

```
pr(form1, form2)
```

Arguments

form1, form2	Two data matrices or data frames; rows represent individuals, columns represent subscores. Both forms should have the same individuals and subscores in the same order. Missing subscores have to be inserted as NA.
--------------	--

Details

Profile pattern and level reliability coefficients are based on the profile analysis approach described in Davison and Davenport (2002) and Bulut (2013). Using the parallel test forms or multiple administration of the same test form, pattern and level reliability coefficients are computed. Pattern reliability is an indicator of variability between the subscores of an examinee and the level reliability is an indicator of the average subscore variation among all examinees. For details, see [Bulut\(2013\)](#)

Value

An object of class prof is returned, listing the following components:

- reliability - Within-in person, between-person, and overall subscore reliability
- pattern.level - A matrix of all pattern and level values obtained from the subscores

References

Bulut, O. (2013). *Between-person and within-person subscore reliability: Comparison of unidimensional and multidimensional IRT models*. (Doctoral dissertation). University of Minnesota. University of Minnesota, Minneapolis, MN. (AAT 3589000).

Davison, M. L., & Davenport, E. C. (2002). Identifying criterion-related patterns of predictor scores using multiple regression. *Psychological Methods*, 7(4), 468-484. DOI: 10.1037/1082-989X.7.4.468

See Also

[plot.prof](#)

Examples

```
## Not run:  
data(EEGS)  
result <- pr(EEGS[,c(1,3,5)],EEGS[,c(2,4,6)])  
print(result)  
plot(result)  
## End(Not run)
```

print.critpat

Print a criterion-related profile analysis

Description

Prints the default output from fitting the cpa function.

Usage

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

Arguments

x object of class critpat returned from the cpa function
... additional objects of the same type.

See Also

[cpa](#)

profileplot	<i>Score Profile Plot</i>
-------------	---------------------------

Description

The profileplot function creates a profile plot for a matrix or dataframe with multiple scores or subscores using [ggplot](#) function in ggplot2 package.

Usage

```
profileplot(form, person.id, standardize = TRUE, interval = 10,  
            by.pattern = TRUE, original.names = TRUE)
```

Arguments

form	A matrix or dataframe including two or more subscores.
person.id	A vector that includes person ID values (Optional).
standardize	If not FALSE, all scores are rescaled with a mean of 0 and standard deviation of 1. Default is TRUE.
interval	The number of equal intervals from the minimum score to the maximum score. Default is 10. Ignored when by.pattern=FALSE.
by.pattern	If TRUE, the function creates a profile plot with level and pattern values using ggplot2. Otherwise, the function creates a profile plot showing profile scores of persons using the base graphics in R. Default is TRUE.
original.names	Use the original column names in the data. Otherwise, columns are renamed as v1,v2,... Default is TRUE.

Value

The profileplot functions returns a score profile plot from either [ggplot](#) or the base graphics in R.

See Also

[ggplot](#), [PS](#)

Examples

```
## Not run:  
data(PS)  
myplot <- profileplot(PS[,2:4], person.id = PS$Person,by.pattern = TRUE, original.names = TRUE)  
myplot  
  
data(leisure)  
leis.plot <- profileplot(leisure[,2:4],standardize=TRUE,by.pattern=FALSE)  
leis.plot  
  
## End(Not run)
```

PS	<i>A Hypothetical Personality Scale from Davison, Kim, and Close (2009)</i>
----	---

Description

The PS shows score profiles of six respondents to a hypothetical personality scale. It includes three types of profile patterns: Linearly increasing, inverted V, and linearly decreasing.

Format

Person Person ID
NEU Neurotic scale score
PSY Psychotic scale score
CD Character disorder scale score

Source

Davison, M. L., Kim, S-K., & Close, C. W. (2009). Factor analytic modeling of within person variation in score profiles. *Multivariate Behavioral Research, 44*, 668-87.

References

Davison, M. L., Kim, S-K., & Close, C. W. (2009). Factor analytic modeling of within person variation in score profiles. *Multivariate Behavioral Research, 44*, 668-87.

spouse	<i>Love and Marriage Survey for Spouses</i>
--------	---

Description

The spouse data come from a study of love and marriage. A sample of 30 husbands and their wives were asked to respond to the following questions:

- Question 1: What is the level of passionate love you feel for your partner?
- Question 2: What is the level of passionate love that your partner feels for you?
- Question 3: What is the level of companionate love that you feel for your partner?
- Question 4: What is the level of companionate love that your partner feels for you?

The responses to all four questions are on a five-point Likert scale where 1 indicates "none at all" and 5 indicates "tremendous amount".

Format

- item1** Question 1 with a score ranging from 1 to 5.
- item2** Question 2 with a score ranging from 1 to 5.
- item3** Question 3 with a score ranging from 1 to 5.
- item4** Question 4 with a score ranging from 1 to 5.
- spouse** Spouse type. It is either "Husband" or "Wife"

Examples

```
## Not run:  
data(spouse)  
  
## End(Not run)
```

summary.critpat	<i>Summary of criterion-related profile analysis</i>
-----------------	--

Description

Provides a summary of the criterion-related profile analysis

Usage

```
## S3 method for class 'critpat'  
summary(object, ...)
```

Arguments

object	object of class critpat
...	additional arguments affecting the summary produced.

See Also

[cpa](#)

`wprifm`*Within-Person Random Intercept Factor Model*

Description

Within-Person Random Intercept Factor Model

Usage

```
wprifm(data, scale = FALSE, save_model = FALSE)
```

Arguments

<code>data</code>	Data.frame containing the manifest variables.
<code>scale</code>	Should the data be scaled? Default = FALSE
<code>save_model</code>	Should the temporary lavaan model syntax be saved. Default = FALSE

Details

This function performs the within-person random intercept factor model described in Davison, Kim, and Close (2009). For information about this model, please see this reference. This function returns an object of lavaan class and thus any generics defined for lavaan will work on this object. This function provides a simple wrapper for lavaan.

Value

an object of class lavaan

References

Davison, M., Kim, S.-K., Close, C. (2009). Factor analytic modeling of within person variation in score profiles. *Multivariate Behavioral Research*, 44(5), 668 - 687. DOI: 10.1080/00273170903187665

Examples

```
data <- HolzingerSwineford1939[,7:ncol(HolzingerSwineford1939)]  
wprifm(data, scale = TRUE)
```

Index

*Topic **datasets**

EEGS, [5](#)
IPMMc, [7](#)
leisure, [7](#)
mod_data, [8](#)
nutrient, [9](#)
PS, [18](#)
spouse, [18](#)

*Topic **dataset**

interest, [5](#)

*Topic **methods**

pcv, [13](#)
pr, [15](#)

*Topic **method**

cpa, [3](#)
mpa, [8](#)

*Topic **package**

profileR-package, [2](#)

anova.critpat, [3](#), [14](#)

cpa, [3](#), [3](#), [9](#), [11](#), [14](#), [16](#), [19](#)

EEGS, [5](#)

family, [3](#), [13](#)

formula, [3](#), [8](#), [13](#)

ggplot, [17](#)

interest, [5](#)

IPMMc, [7](#)

leisure, [7](#)

mod_data, [8](#)

mpa, [8](#), [8](#)

nutrient, [9](#)

pams, [10](#)

paos, [11](#)

pbg, [12](#)

pcv, [4](#), [13](#)

plot.critpat, [14](#), [14](#)

plot.prof, [15](#), [16](#)

pr, [11](#), [12](#), [15](#), [15](#)

print.critpat, [14](#), [16](#)

profileplot, [12](#), [17](#)

profileR-package, [2](#)

PS, [17](#), [18](#)

spouse, [18](#)

summary.critpat, [14](#), [19](#)

wprifm, [20](#)