

# Package ‘vpc’

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**Title** Create Visual Predictive Checks

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**Description** Visual predictive checks are a commonly used diagnostic plot in pharmacometrics, showing how certain statistics (percentiles) for observed data compare to those same statistics for data simulated from a model. The package can generate VPCs for continuous, categorical, censored, and (repeated) time-to-event data.

**Depends** R (>= 3.1.0)

**Imports** classInt, dplyr, MASS, survival, ggplot2, readr, stringr, tidyverse

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**LazyData** true

**URL** <https://github.com/ronkeizer/vpc>

**Suggests** knitr, testit

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**Repository** CRAN

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vpc-package*VPC package*

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**Description**

Create Visual Predictive Checks in R

**Author(s)**

Ron Keizer &lt;ronkeizer@gmail.com&gt;

---

|           |   |
|-----------|---|
| add_noise | <i>Add noise / residual error to data</i> |
|-----------|---|

---

## Description

Add noise / residual error to data

## Usage

```
add_noise(x, ruv = list(proportional = 0, additive = 0, exponential = 0))
```

## Arguments

|     |   |
|-----|---|
| x   | data  |
| rvu | list describing the magnitude of errors. List arguments: "proportional", "additive", "exponential". |

## Examples

```
library(dplyr)
ipred <- c(10, 8, 6, 4, 2, 0) %>% add_noise(ruv = list(proportional = 0.1, additive = 0.2))
```

---

|                      |                             |
|----------------------|-----------------------------|
| add_sim_index_number | <i>Add sim index number</i> |
|----------------------|-----------------------------|

---

## Description

Add simulation index number to simulation when not present

## Usage

```
add_sim_index_number(sim, id = "id", sim_label = "sim")
```

## Arguments

|           |  |
|-----------|--|
| sim       | a data.frame containing the simulation data            |
| id        | character specifying the column name in the data.frame |
| sim_label | label to indicate simulation index (if available)      |

**add\_stratification**      *Adds stratification to data set*

### Description

Adds stratification to data set

### Usage

```
add_stratification(dat, strat, verbose = FALSE)
```

### Arguments

|         |                                    |
|---------|------------------------------------|
| dat     | data.frame                         |
| strat   | vector of stratification variables |
| verbose | verbosity ('TRUE' or 'FALSE')      |

**auto\_bin**      *Calculate appropriate bin separators for vpc*

### Description

This function calculates bin separators either using R's native binning approaches available in the classInt library such as 'kmeans', 'jenks', 'pretty' etc. Alternatively, a custom approach is available which is based on finding the nadirs in the density functions for the independent variable. Default approach is k-means clustering.

### Usage

```
auto_bin(dat, type = "kmeans", n_bins = 8, verbose = FALSE, ...)

## S3 method for class 'numeric'
auto_bin(dat, type = "kmeans", n_bins = 8, verbose = FALSE, ...)

## S3 method for class 'data.frame'
auto_bin(dat, type = "kmeans", n_bins = 8, verbose = FALSE, ...)
```

### Arguments

|         |   |
|---------|---|
| dat     | data frame  |
| type    | auto-binning type: "density", "time", or "data"   |
| n_bins  | number of bins to use; either a positive integer or "auto". For "density" the function might not return a solution with the exact number of bins. |
| verbose | show warnings and other messages (TRUE or FALSE)  |
| ...     | arguments passed on to underlying binning functions   |

**Value**

A vector of bin separators

---

**bin\_data**

*Function to bin data based on a vector of bin separators, e.g. for use in VPC*

---

**Description**

Function to bin data based on a vector of bin separators, e.g. for use in VPC

**Usage**

```
bin_data(x, bins = c(0, 3, 5, 7), idv = "time", labeled = F)
```

**Arguments**

|         |   |
|---------|---|
| x       | data  |
| bins    | numeric vector specifying bin separators                              |
| idv     | variable in the data specifies the independent variable (e.g. "time") |
| labeled | whether a labeled factor instead of integers should be returned       |

---

**check\_stratification\_columns\_available**

*Check whether stratification columns are available*

---

**Description**

Check whether stratification columns are available

**Usage**

```
check_stratification_columns_available(data, stratify, type = "observation")
```

**Arguments**

|          |  |
|----------|--|
| data     | 'data.frame' with observation or simulation data |
| stratify | vector of stratification columns                 |
| type     | either 'observation' or 'simulation'             |

---

|                             |  |
|-----------------------------|--|
| <code>compute_kaplan</code> | <i>Compute Kaplan-Meier statistics</i> |
|-----------------------------|--|

---

### Description

Compute Kaplan-Meier statistics

### Usage

```
compute_kaplan(
  dat,
  strat = "strat",
  reverse_prob = FALSE,
  rtte_conditional = TRUE,
  ci = NULL
)
```

### Arguments

|                               |   |
|-------------------------------|---|
| <code>dat</code>              | data.frame with events  |
| <code>strat</code>            | vector of stratification variables  |
| <code>reverse_prob</code>     | reverse the probability (i.e. return '1-probability')?  |
| <code>rtte_conditional</code> | 'TRUE' (default) or 'FALSE'. Compute the probability for each event newly ('TRUE'), or calculate the absolute probability ('FALSE', i.e. the "probability of a 1st, 2nd, 3rd event etc" rather than the "probability of an event happening"). |
| <code>ci</code>               | confidence interval to calculate, numeric vector of length 2  |

---



---

|                           |                                |
|---------------------------|--------------------------------|
| <code>compute_kmmc</code> | <i>Compute KMMC statistics</i> |
|---------------------------|--------------------------------|

---

### Description

Kaplan-Meier Mean Covariate plots are a simulation-based diagnostic to study the influence of covariates and identify potential model misspecification.

### Usage

```
compute_kmmc(dat, strat = NULL, reverse_prob = FALSE, kmmc = "DOSE")
```

### Arguments

|                           |  |
|---------------------------|--|
| <code>dat</code>          | data.frame with events                                 |
| <code>strat</code>        | vector of stratification variables                     |
| <code>reverse_prob</code> | reverse the probability (i.e. return '1-probability')? |
| <code>kmmc</code>         | variable to create the KMMC plot for.                  |

---

|                  |                             |
|------------------|-----------------------------|
| create_vpc_theme | <i>Create new vpc theme</i> |
|------------------|-----------------------------|

---

## Description

Create new vpc theme

## Usage

```
create_vpc_theme(...)
```

## Arguments

|     |                                   |
|-----|-----------------------------------|
| ... | pass arguments to ‘new_vpc_theme’ |
|-----|-----------------------------------|

---

---

|                     |  |
|---------------------|--|
| define_data_columns | <i>Define data column defaults for various softwares</i> |
|---------------------|--|

---

## Description

Define data column defaults for various softwares

## Usage

```
define_data_columns(sim, obs, sim_cols, obs_cols, software_type)
```

## Arguments

|               |   |
|---------------|---|
| sim           | simulated data  |
| obs           | observed data   |
| sim_cols      | list for mapping simulation data columns, e.g. ‘list(dv = "DV", id = "ID", idv = "TIME", pred="PRED")’  |
| obs_cols      | list for mapping observation data columns, e.g. ‘list(dv = "DV", id = "ID", idv = "TIME", pred="PRED")’ |
| software_type | software type, one of ‘nonmem’, ‘phoenix’, ‘PKPDsim’  |

**draw\_params\_mvr**      *Draw parameters from multivariate distribution*

### Description

Draw parameters from multivariate distribution

### Usage

```
draw_params_mvr(ids, n_sim, theta, omega_mat, par_names = NULL)
```

### Arguments

|           |                        |
|-----------|------------------------|
| ids       | vector of id numbers   |
| n_sim     | number of simulations  |
| theta     | theta vector           |
| omega_mat | omega matrix           |
| par_names | parameter names vector |

**loq\_perc**      *Calculate percentiles below / above lloq / uloq*

### Description

Calculate percentiles below / above lloq / uloq

### Usage

```
loq_perc(x, limit = 1, cens = "left")
```

### Arguments

|       |                                  |
|-------|----------------------------------|
| x     | data                             |
| limit | censoring limit                  |
| cens  | censoring direction (left/right) |

---

new\_vpc\_theme      *Create a customized VPC theme*

---

## Description

Create a customized VPC theme

## Usage

```
new_vpc_theme(update = NULL)
```

## Arguments

update      list containing the plot elements to be updated. Run ‘new\_vpc\_theme()’ with no arguments to show an overview of available plot elements.

## Details

This function creates a theme that customizes how the VPC looks, i.e. colors, fills, transparencies, linetypes and sizes, etc. The following arguments can be specified in the input list:

- obs\_color: color for observations points
- obs\_size: size for observation points
- obs\_median\_color: color for median observation line
- obs\_median\_linetype: linetype for median observation line
- obs\_median\_size: size for median observation line
- obs\_ci\_fill: color for observation CI fill
- obs\_ci\_color: color for observation CI lines
- obs\_ci\_linetype: linetype for observation CI lines
- obs\_ci\_size: size for observations CI lines
- sim\_pi\_fill: fill color for simulated prediction interval areas
- sim\_pi\_alpha: transparency for simulated prediction interval areas
- sim\_pi\_color: color for simulated prediction interval lines
- sim\_pi\_linetype: linetype for simulated prediction interval lines
- sim\_pi\_size: size for simulated prediction interval lines
- sim\_median\_fill: fill color for simulated median area
- sim\_median\_alpha: transparency for simulated median area
- sim\_median\_color: color for simulated median line
- sim\_median\_linetype: linetype for simulated median line
- sim\_median\_size: size for simulated median line
- bin\_separators\_color: color for bin separator lines, NA for don't plot
- bin\_separators\_location: where to plot bin separators ("t" for top, "b" for bottom)
- loq\_color: color of line showing limit of quantification

**Value**

A list with vpc theme specifiers

**Examples**

```
theme1 <- new_vpc_theme(update = list(
  obs_color = "red",
  obs_ci_color = "#aa0000",
  obs_alpha = .3,
  sim_pi_fill = "#cc8833",
  sim_pi_size = 2
))
vpc(simple_data$sim, simple_data$obs, vpc_theme = theme1)
```

**pk\_iv\_1cmt***Simulate PK data from a 1-compartment iv model***Description**

Simulate PK data from a 1-compartment iv model

**Usage**

```
pk_iv_1cmt(
  t,
  t_inf = 1,
  tau = 24,
  dose = 120,
  CL = 0.345,
  Vc = 1.75,
  ruv = NULL
)
```

**Arguments**

|                    |                        |
|--------------------|------------------------|
| <code>t</code>     | Time after dose        |
| <code>t_inf</code> | Infusion length        |
| <code>tau</code>   | Dosing interval        |
| <code>dose</code>  | Dose                   |
| <code>CL</code>    | Clearance              |
| <code>Vc</code>    | Volume of distribution |
| <code>ruv</code>   | Residual variability   |

**Value**

A vector of predicted values, with or without added residual variability

## Examples

```
dat1 <- vpc:::pk_iv_1cmt(t = c(0:72), tau = 24, dose = 120,
                           CL = 5, Vc = 50)
dat2 <- vpc:::pk_iv_1cmt(t = c(0:72), tau = 24, dose = 120,
                           CL = 5, Vc = 50,
                           ruv = list(proportional = 0.1, additive = 0.1))
```

**pk\_oral\_1cmt**

*Simulate PK data from a 1-compartment oral model*

## Description

Simulate PK data from a 1-compartment oral model

## Usage

```
pk_oral_1cmt(t, tau = 24, dose = 120, ka = 1, ke = 1, cl = 10, ruv = NULL)
```

## Arguments

|      |                      |
|------|----------------------|
| t    | Time after dose      |
| tau  | Dosing interval      |
| dose | Dose                 |
| ka   | Absorption rate      |
| ke   | Elimination rate     |
| cl   | Clearance            |
| ruv  | Residual variability |

## Value

A vector of predicted values, with or without added residual variability

## Examples

```
dat1 <- vpc:::pk_oral_1cmt(t = c(0:72), tau = 24, dose = 120,
                           ka = 1, ke = 1, cl = 10)
dat2 <- vpc:::pk_oral_1cmt(t = c(0:72), tau = 24, dose = 120,
                           ka = 1, ke = 1, cl = 10,
                           ruv = list(proportional = 0.1, additive = 0.1))
```

---

|          |                              |
|----------|------------------------------|
| plot_vpc | <i>VPC plotting function</i> |
|----------|------------------------------|

---

## Description

This function performs no parsing of data, it just plots the already calculated statistics generated using one of the ‘vpc’ functions.

## Usage

```
plot_vpc(
  db,
  show = NULL,
  vpc_theme = NULL,
  smooth = TRUE,
  log_x = FALSE,
  log_y = FALSE,
  xlab = NULL,
  ylab = NULL,
  title = NULL,
  verbose = FALSE
)
```

## Arguments

|                        |  |
|------------------------|--|
| <code>db</code>        | object created using the ‘vpc’ function  |
| <code>show</code>      | what to show in VPC (obs_dv, obs_ci, pi, pi_as_area, pi_ci, obs_median, sim_median, sim_median_ci)                   |
| <code>vpc_theme</code> | theme to be used in VPC. Expects list of class <code>vpc_theme</code> created with function <code>vpc_theme()</code> |
| <code>smooth</code>    | “smooth” the VPC (connect bin midpoints) or show bins as rectangular boxes. Default is TRUE.                         |
| <code>log_x</code>     | Boolean indicating whether x-axis should be shown as logarithmic. Default is FALSE.                                  |
| <code>log_y</code>     | Boolean indicating whether y-axis should be shown as logarithmic. Default is FALSE.                                  |
| <code>xlab</code>      | label for x axis   |
| <code>ylab</code>      | label for y axis   |
| <code>title</code>     | title  |
| <code>verbose</code>   | verbosity (T/F)  |

## See Also

[sim\\_data](#), [vpc\\_cens](#), [vpc\\_tte](#), [vpc\\_cat](#)

## Examples

```
## See vpc.ronkeizer.com for more documentation and examples

library(vpc)
vpc_db <- vpc(sim = simple_data$sim, obs = simple_data$obs, vpcdb = TRUE)
plot_vpc(vpc_db, title = "My new vpc", x = "Custom x label")
```

quantile\_cens

*Calculate quantiles respecting the censored data*

## Description

Calculate quantiles respecting the censored data

## Usage

```
quantile_cens(x, p = 0.5, limit = 1, cens = "left")
```

## Arguments

|       |                                  |
|-------|----------------------------------|
| x     | data                             |
| p     | quantile                         |
| limit | censoring limit                  |
| cens  | censoring direction (left/right) |

read\_table\_nm

*NONMEM output table import function*

## Description

Quickly import NONMEM output tables into R. Function taken from ‘modelviz’ package by Benjamin Guiastrennec. When both skip and header are NULL, read\_nmtab will automatically detect the optimal settings to import the tables. When more than one files are provided for a same NONMEM run, they will be combined into a single data.frame.

## Usage

```
read_table_nm(
  file = NULL,
  skip = NULL,
  header = NULL,
  rm_duplicates = FALSE,
  nonmem_tab = TRUE
)
```

**Arguments**

|                            |  |
|----------------------------|--|
| <code>file</code>          | full file name   |
| <code>skip</code>          | number of lines to skip before reading data  |
| <code>header</code>        | logical value indicating whether the file contains the names of the variables as its first line          |
| <code>rm_duplicates</code> | logical value indicating whether duplicated columns should be removed                                    |
| <code>nonmem_tab</code>    | logical value indicating to the function whether the file is a table or a nonmem additional output file. |

**Value**

A `data.frame`

**Examples**

```
## Not run:
data <- read_table_nm(file = '../models/pk/sdtab101')

## End(Not run)
```

`replace_list_elements` *Replace list elements by name*

**Description**

Replace list elements by name

**Usage**

```
replace_list_elements(list, replacement)
```

**Arguments**

|                          |                  |
|--------------------------|------------------|
| <code>list</code>        | original list    |
| <code>replacement</code> | replacement list |

**Details**

Finds and replaces list elements by name and throws an error if an element is not available in the original list. This is a local duplicate of the PKPDmisc copy for the VPC package to reduce dependency on PKPDmisc at this time.

## Examples

```
## Not run:  
list <- list(ipred = "ipred", dv = "dv", idv = "idv", "pred" = "pred")  
replacement <- list(dv = "conc", idv = "time")  
list <- replace_list_elements(list, replacement)  
  
## End(Not run)
```

---

rtte\_obs\_nm

*Simulated RTTE data (1x)*

---

## Description

An example dataset with simulated repeated time-to-event data

## Usage

rtte\_obs\_nm

## Format

An object of class `data.frame` with 573 rows and 6 columns.

---

rtte\_sim\_nm

*Simulated RTTE data (100x)*

---

## Description

An example dataset with simulated repeated time-to-event data (100 simulations)

## Usage

rtte\_sim\_nm

## Format

An object of class `data.frame` with 2000000 rows and 7 columns.

---

|              |                                   |
|--------------|-----------------------------------|
| show_default | <i>Defaults for show argument</i> |
|--------------|-----------------------------------|

---

**Description**

Defaults for show argument

**Usage**

`show_default`

**Format**

An object of class `list` of length 11.

---

|                  |   |
|------------------|---|
| show_default_tte | <i>Defaults for show argument for TTE VPC</i> |
|------------------|---|

---

**Description**

Defaults for show argument for TTE VPC

**Usage**

`show_default_tte`

**Format**

An object of class `list` of length 11.

---

|             |                             |
|-------------|-----------------------------|
| simple_data | <i>A small rich dataset</i> |
|-------------|-----------------------------|

---

**Description**

A small rich dataset

**Usage**

`simple_data`

**Format**

An object of class `list` of length 2.

## Details

a list containing the obs and sim data for an example dataset to run a simple vpc.

## Examples

```
## Not run:
vpc(simple_data$sim, simple_data$obs)

## End(Not run)
```

**sim\_data**

*Simulate data based on a model and parameter distributions*

## Description

Simulate data based on a model and parameter distributions

## Usage

```
sim_data(
  design = cbind(id = c(1, 1, 1), idv = c(0, 1, 2)),
  model = function(x) {      return(x$alpha + x$beta) },
  theta,
  omega_mat,
  par_names,
  par_values = NULL,
  draw_iiv = "mvrnorm",
  error = list(proportional = 0, additive = 0, exponential = 0),
  n = 100
)
```

## Arguments

|            |  |
|------------|--|
| design     | a design dataset. See example  |
| model      | A function with the first argument the simulation design, i.e. a dataset with the columns ... The second argument to this function is a dataset with parameters for every individual. This can be supplied by the user, or generated by this sim_data if theta and omega_mat are supplied. |
| theta      | vector of fixed effect parameters  |
| omega_mat  | vector of between subject random effects, specified as lower triangle  |
| par_names  | A character vector linking the parameters in the model to the variables in the dataset. See example.   |
| par_values | parameter values   |
| draw_iiv   | draw between subject random effects?   |
| error      | see example  |
| n          | number of simulations to perform   |

## Details

This function generates the simulated dependent values for use in the VPC plotting function.

## Value

a vector of simulated dependent variables (for us in the VPC plotting function)

## See Also

[vpc](#)

`theme_empty`

*Empty ggplot2 theme*

## Description

Empty ggplot2 theme

## Usage

`theme_empty()`

## Examples

```
vpc(simple_data$sim, simple_data$obs) + theme_empty()
```

`theme_plain`

*A nicer default theme for ggplot2*

## Description

A nicer default theme for ggplot2

## Usage

`theme_plain()`

## Examples

```
vpc(simple_data$sim, simple_data$obs) + theme_plain()
```

---

|                  |                               |
|------------------|-------------------------------|
| triangle_to_full | <i>Lower to full triangle</i> |
|------------------|-------------------------------|

---

### Description

Convert the lower triangle of a covariance matrix to a full matrix object

### Usage

```
triangle_to_full(vect)
```

### Arguments

|      |   |
|------|---|
| vect | the lower triangle of a covariance matrix |
|------|---|

---

---

|     |                     |
|-----|---------------------|
| vpc | <i>VPC function</i> |
|-----|---------------------|

---

### Description

Creates a VPC plot from observed and simulation data

### Usage

```
vpc(sim, ...)

## Default S3 method:
vpc(sim, ...)

vpc_vpc(
  sim = NULL,
  obs = NULL,
  psn_folder = NULL,
  bins = "jenks",
  n_bins = "auto",
  bin_mid = "mean",
  obs_cols = NULL,
  sim_cols = NULL,
  software = "auto",
  show = NULL,
  stratify = NULL,
  pred_corr = FALSE,
  pred_corr_lower_bnd = 0,
  pi = c(0.05, 0.95),
  ci = c(0.05, 0.95),
```

```

  uloq = NULL,
  lloq = NULL,
  log_y = FALSE,
  log_y_min = 0.001,
  xlab = NULL,
  ylab = NULL,
  title = NULL,
  smooth = TRUE,
  vpc_theme = NULL,
  facet = "wrap",
  scales = "fixed",
  labeller = NULL,
  vpcdb = FALSE,
  verbose = FALSE,
  ...
)

```

## Arguments

|                                  |  |
|----------------------------------|--|
| <code>sim</code>                 | this is usually a data.frame with observed data, containing the independent and dependent variable, a column indicating the individual, and possibly covariates. E.g. load in from NONMEM using <a href="#">read_table_nm</a> . However it can also be an object like a nlmixr or xpose object |
| <code>...</code>                 | Other arguments sent to other methods (like xpose or nlmixr); Note these arguments are not used in the default vpc and are ignored by the default method.  |
| <code>obs</code>                 | a data.frame with observed data, containing the independent and dependent variable, a column indicating the individual, and possibly covariates. E.g. load in from NONMEM using <a href="#">read_table_nm</a>  |
| <code>psn_folder</code>          | instead of specifying "sim" and "obs", specify a PsN-generated VPC-folder  |
| <code>bins</code>                | either "density", "time", or "data", "none", or one of the approaches available in <code>classInterval()</code> such as "jenks" (default) or "pretty", or a numeric vector specifying the bin separators.  |
| <code>n_bins</code>              | when using the "auto" binning method, what number of bins to aim for   |
| <code>bin_mid</code>             | either "mean" for the mean of all timepoints (default) or "middle" to use the average of the bin boundaries.   |
| <code>obs_cols</code>            | observation dataset column names (list elements: "dv", "idv", "id", "pred")  |
| <code>sim_cols</code>            | simulation dataset column names (list elements: "dv", "idv", "id", "pred", "sim")  |
| <code>software</code>            | name of software platform using (e.g. nonmem, phoenix)   |
| <code>show</code>                | what to show in VPC (obs_dv, obs_ci, pi, pi_as_area, pi_ci, obs_median, sim_median, sim_median_ci)   |
| <code>stratify</code>            | character vector of stratification variables. Only 1 or 2 stratification variables can be supplied.  |
| <code>pred_corr</code>           | perform prediction-correction?   |
| <code>pred_corr_lower_bnd</code> | lower bound for the prediction-correction  |

|           |  |
|-----------|--|
| pi        | simulated prediction interval to plot. Default is c(0.05, 0.95),                             |
| ci        | confidence interval to plot. Default is (0.05, 0.95)   |
| ulq       | Number or NULL indicating upper limit of quantification. Default is NULL.                    |
| lloq      | Number or NULL indicating lower limit of quantification. Default is NULL.                    |
| log_y     | Boolean indicating whether y-axis should be shown as logarithmic. Default is FALSE.          |
| log_y_min | minimal value when using log_y argument. Default is 1e-3.                                    |
| xlab      | label for x axis   |
| ylab      | label for y axis   |
| title     | title  |
| smooth    | "smooth" the VPC (connect bin midpoints) or show bins as rectangular boxes. Default is TRUE. |
| vpc_theme | theme to be used in VPC. Expects list of class vpc_theme created with function vpc_theme()   |
| facet     | either "wrap", "columns", or "rows"  |
| scales    | either "fixed" (default), "free_y", "free_x" or "free"                                       |
| labeller  | ggplot2 labeller function to be passed to underlying ggplot object                           |
| vpcdb     | Boolean whether to return the underlying vpcdb rather than the plot                          |
| verbose   | show debugging information (TRUE or FALSE)   |

## Value

a list containing calculated VPC information (when vpcdb=TRUE), or a ggplot2 object (default)

## See Also

[sim\\_data](#), [vpc\\_cens](#), [vpc\\_tte](#), [vpc\\_cat](#)

## Examples

```
## See vpc.ronkeizer.com for more documentation and examples
library(vpc)

# Basic commands:
vpc(sim = simple_data$sim, obs = simple_data$obs)
vpc(sim = simple_data$sim, obs = simple_data$obs, lloq = 20)
```

---

|                      |                                     |
|----------------------|-------------------------------------|
| <code>vpc_cat</code> | <i>VPC function for categorical</i> |
|----------------------|-------------------------------------|

---

## Description

Creates a VPC plot from observed and simulation data for categorical variables.

## Usage

```
vpc_cat(
  sim = NULL,
  obs = NULL,
  psn_folder = NULL,
  bins = "jenks",
  n_bins = "auto",
  bin_mid = "mean",
  obs_cols = NULL,
  sim_cols = NULL,
  software = "auto",
  show = NULL,
  ci = c(0.05, 0.95),
  uloq = NULL,
  lloq = NULL,
  xlab = NULL,
  ylab = NULL,
  title = NULL,
  smooth = TRUE,
  vpc_theme = NULL,
  facet = "wrap",
  labeller = NULL,
  plot = TRUE,
  vpcdb = FALSE,
  verbose = FALSE
)
```

## Arguments

|                         |   |
|-------------------------|---|
| <code>sim</code>        | a data.frame with observed data, containing the independent and dependent variable, a column indicating the individual, and possibly covariates. E.g. load in from NONMEM using <a href="#">read_table_nm</a> |
| <code>obs</code>        | a data.frame with observed data, containing the independent and dependent variable, a column indicating the individual, and possibly covariates. E.g. load in from NONMEM using <a href="#">read_table_nm</a> |
| <code>psn_folder</code> | instead of specifying "sim" and "obs", specify a PsN-generated VPC-folder   |
| <code>bins</code>       | either "density", "time", or "data", "none", or one of the approaches available in <code>classInterval()</code> such as "jenks" (default) or "pretty", or a numeric vector specifying the bin separators.     |

|           |  |
|-----------|--|
| n_bins    | when using the "auto" binning method, what number of bins to aim for   |
| bin_mid   | either "mean" for the mean of all timepoints (default) or "middle" to use the average of the bin boundaries. |
| obs_cols  | observation dataset column names (list elements: "dv", "idv", "id", "pred")                                  |
| sim_cols  | simulation dataset column names (list elements: "dv", "idv", "id", "pred")                                   |
| software  | name of software platform using (e.g. nonmem, phoenix)   |
| show      | what to show in VPC (obs_ci, pi, pi_as_area, pi_ci, obs_median, sim_median, sim_median_ci)                   |
| ci        | confidence interval to plot. Default is (0.05, 0.95)   |
| ulq       | Number or NULL indicating upper limit of quantification. Default is NULL.                                    |
| llq       | Number or NULL indicating lower limit of quantification. Default is NULL.                                    |
| xlab      | label for x-axis   |
| ylab      | label for y-axis   |
| title     | title  |
| smooth    | "smooth" the VPC (connect bin midpoints) or show bins as rectangular boxes. Default is TRUE.                 |
| vpc_theme | theme to be used in VPC. Expects list of class vpc_theme created with function vpc_theme()                   |
| facet     | either "wrap", "columns", or "rows"  |
| labeller  | ggplot2 labeller function to be passed to underlying ggplot object   |
| plot      | Boolean indicating whether to plot the ggplot2 object after creation. Default is FALSE.                      |
| vpcdb     | boolean whether to return the underlying vpcdb rather than the plot  |
| verbose   | show debugging information (TRUE or FALSE)   |

## Value

a list containing calculated VPC information (when vpcdb=TRUE), or a ggplot2 object (default)

## See Also

[sim\\_data](#), [vpc](#), [vpc\\_tte](#), [vpc\\_cens](#)

## Examples

```
## See vpc.ronkeizer.com for more documentation and examples
library(vpc)

# simple function to simulate categorical data for single individual
sim_id <- function(id = 1) {
  n <- 10
  logit <- function(x) exp(x) / (1+exp(x))
  data.frame(id = id, time = seq(1, n, length.out = n),
```

```

        dv = round(logit((1:n) - n/2 + rnorm(n, 0, 1.5))) )
}
## simple function to simulate categorical data for a trial
sim_trial <- function(i = 1, n = 20) { # function to simulate categorical data for a trial
  data.frame(sim = i, do.call("rbind", lapply(1:n, sim_id)))
}

## simulate single trial for 20 individuals
obs <- sim_trial(n = 20)

## simulate 200 trials of 20 individuals
sim <- do.call("rbind", lapply(1:200, sim_trial, n = 20))

## Plot categorical VPC
vpc_cat(sim = sim, obs = obs)

```

vpc\_cens

*VPC function for left- or right-censored data (e.g. BLOQ data)*

## Description

Creates a VPC plot from observed and simulation data for censored data. Function can handle both left- (below lower limit of quantification) and right-censored (above upper limit of quantification) data.

## Usage

```
vpc_cens(
  sim = NULL,
  obs = NULL,
  psn_folder = NULL,
  bins = "jenks",
  n_bins = 8,
  bin_mid = "mean",
  obs_cols = NULL,
  sim_cols = NULL,
  software = "auto",
  show = NULL,
  stratify = NULL,
  stratify_color = NULL,
  ci = c(0.05, 0.95),
  uloq = NULL,
  lloq = NULL,
  plot = FALSE,
  xlab = "Time",
  ylab = "Probability of <LOQ",
  title = NULL,
  smooth = TRUE,
```

```

    vpc_theme = NULL,
    facet = "wrap",
    labeller = NULL,
    vpcdb = FALSE,
    verbose = FALSE
)

```

## Arguments

|                             |   |
|-----------------------------|---|
| <code>sim</code>            | a data.frame with observed data, containing the independent and dependent variable, a column indicating the individual, and possibly covariates. E.g. load in from NONMEM using <a href="#">read_table_nm</a> |
| <code>obs</code>            | a data.frame with observed data, containing the independent and dependent variable, a column indicating the individual, and possibly covariates. E.g. load in from NONMEM using <a href="#">read_table_nm</a> |
| <code>psn_folder</code>     | instead of specifying "sim" and "obs", specify a PsN-generated VPC-folder   |
| <code>bins</code>           | either "density", "time", or "data", or a numeric vector specifying the bin separators.   |
| <code>n_bins</code>         | number of bins  |
| <code>bin_mid</code>        | either "mean" for the mean of all timepoints (default) or "middle" to use the average of the bin boundaries.  |
| <code>obs_cols</code>       | observation dataset column names (list elements: "dv", "idv", "id", "pred")   |
| <code>sim_cols</code>       | simulation dataset column names (list elements: "dv", "idv", "id", "pred")  |
| <code>software</code>       | name of software platform using (e.g. nonmem, phoenix)  |
| <code>show</code>           | what to show in VPC (obs_ci, pi, pi_as_area, pi_ci, obs_median, sim_median, sim_median_ci)  |
| <code>stratify</code>       | character vector of stratification variables. Only 1 or 2 stratification variables can be supplied.   |
| <code>stratify_color</code> | variable to stratify and color lines for observed data. Only 1 stratification variables can be supplied.  |
| <code>ci</code>             | confidence interval to plot. Default is (0.05, 0.95)  |
| <code>ulq</code>            | Number or NULL indicating upper limit of quantification. Default is NULL.   |
| <code>llq</code>            | Number or NULL indicating lower limit of quantification. Default is NULL.   |
| <code>plot</code>           | Boolean indicating whether to plot the ggplot2 object after creation. Default is FALSE.   |
| <code>xlab</code>           | ylab as numeric vector of size 2  |
| <code>ylab</code>           | ylab as numeric vector of size 2  |
| <code>title</code>          | title   |
| <code>smooth</code>         | "smooth" the VPC (connect bin midpoints) or show bins as rectangular boxes. Default is TRUE.  |
| <code>vpc_theme</code>      | theme to be used in VPC. Expects list of class vpc_theme created with function vpc_theme()  |

|          |   |
|----------|---|
| facet    | either "wrap", "columns", or "rows"                                 |
| labeller | ggplot2 labeller function to be passed to underlying ggplot object  |
| vpcdb    | boolean whether to return the underlying vpcdb rather than the plot |
| verbose  | show debugging information (TRUE or FALSE)                          |

**Value**

a list containing calculated VPC information, and a ggplot2 object

**See Also**

[sim\\_data](#), [vpc](#), [vpc\\_tte](#), [vpc\\_cat](#)

**Examples**

```
## See vpc.ronkeizer.com for more documentation and examples
library(vpc)

vpc_cens(sim = simple_data$sim, obs = simple_data$obs, lloq = 30)
vpc_cens(sim = simple_data$sim, obs = simple_data$obs, uloq = 120)
```

vpc\_tte

*VPC function for time-to-event (survival) data*

**Description**

This function can be used for either single time-to-event (TTE) or repeated time-to-event (RTTE) data.

**Usage**

```
vpc_tte(
  sim = NULL,
  obs = NULL,
  psn_folder = NULL,
  rtte = FALSE,
  rtte_calc_diff = TRUE,
  rtte_conditional = TRUE,
  events = NULL,
  bins = FALSE,
  n_bins = 10,
  software = "auto",
  obs_cols = NULL,
  sim_cols = NULL,
  kmmc = NULL,
```

```

reverse_prob = FALSE,
stratify = NULL,
stratify_color = NULL,
ci = c(0.05, 0.95),
plot = FALSE,
xlab = "Time",
ylab = "Survival (%)",
show = NULL,
as_percentage = TRUE,
title = NULL,
smooth = FALSE,
vpc_theme = NULL,
facet = "wrap",
labeller = NULL,
verbose = FALSE,
vpcdb = FALSE
)

```

## Arguments

|                               |  |
|-------------------------------|--|
| <code>sim</code>              | a data.frame with observed data, containing the independent and dependent variable, a column indicating the individual, and possibly covariates. E.g. load in from NONMEM using <a href="#">read_table_nm</a>  |
| <code>obs</code>              | a data.frame with observed data, containing the independent and dependent variable, a column indicating the individual, and possibly covariates. E.g. load in from NONMEM using <a href="#">read_table_nm</a>  |
| <code>psn_folder</code>       | instead of specifying "sim" and "obs", specify a PsN-generated VPC-folder  |
| <code>rtte</code>             | repeated time-to-event data? Default is FALSE (treat as single-event TTE)  |
| <code>rtte_calc_diff</code>   | recalculate time (T/F)? When simulating in NONMEM, you will probably need to set this to TRUE to recalculate the TIME to relative times between events (unless you output the time difference between events and specify that as independent variable to the vpc_tte() function. |
| <code>rtte_conditional</code> | 'TRUE' (default) or 'FALSE'. Compute the probability for each event newly ('TRUE'), or calculate the absolute probability ('FALSE', i.e. the "probability of a 1st, 2nd, 3rd event etc" rather than the "probability of an event happening").                                    |
| <code>events</code>           | numeric vector describing which events to show a VPC for when repeated TTE data, e.g. c(1:4). Default is NULL, which shows all events.   |
| <code>bins</code>             | either "density", "time", or "data", or a numeric vector specifying the bin separators.  |
| <code>n_bins</code>           | number of bins   |
| <code>software</code>         | name of software platform using (e.g. nonmem, phoenix)   |
| <code>obs_cols</code>         | observation dataset column names (list elements: "dv", "idv", "id", "pred")  |
| <code>sim_cols</code>         | simulation dataset column names (list elements: "dv", "idv", "id", "pred", "sim")  |
| <code>kmmc</code>             | either NULL (for regular TTE vpc, default), or a variable name for a KMMC plot (e.g. "WT")   |

|                |  |
|----------------|--|
| reverse_prob   | reverse the probability scale (i.e. plot 1-probability)  |
| stratify       | character vector of stratification variables. Only 1 or 2 stratification variables can be supplied.  |
| stratify_color | character vector of stratification variables. Only 1 stratification variable can be supplied, cannot be used in conjunction with ‘stratify’. |
| ci             | confidence interval to plot. Default is (0.05, 0.95)   |
| plot           | Boolean indicating whether to plot the ggplot2 object after creation. Default is FALSE.  |
| xlab           | label for x-axis   |
| ylab           | label for y-axis   |
| show           | what to show in VPC (obs_ci, obs_median, sim_median, sim_median_ci)  |
| as_percentage  | Show y-scale from 0-100 percent? TRUE by default, if FALSE then scale from 0-1.  |
| title          | title  |
| smooth         | “smooth” the VPC (connect bin midpoints) or show bins as rectangular boxes. Default is TRUE.   |
| vpc_theme      | theme to be used in VPC. Expects list of class vpc_theme created with function vpc_theme()   |
| facet          | either “wrap”, “columns”, or “rows”  |
| labeller       | ggplot2 labeller function to be passed to underlying ggplot object   |
| verbose        | TRUE or FALSE (default)  |
| vpcdb          | Boolean whether to return the underlying vpcdb rather than the plot  |

## Details

Creates a VPC plot from observed and simulation survival data

## Value

a list containing calculated VPC information, and a ggplot2 object

## See Also

[sim\\_data](#), [vpc](#), [vpc\\_tte](#), [vpc\\_cens](#)

## Examples

```
## See vpc-docs.ronkeizer.com for more documentation and examples.

## Example for repeated) time-to-event data
## with NONMEM-like data (e.g. simulated using a dense grid)

data(rtte_obs_nm)
data(rtte_sim_nm)
```

```
# treat RTTE as TTE, no stratification
vpc_tte(sim = rtte_sim_nm[rtte_sim_nm$sim <= 20,],
         obs = rtte_obs_nm,
         rtte = FALSE,
         sim_cols=list(dv = "dv", idv = "t"), obs_cols=list(idv = "t"))
```

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