

Package: `pharmac.viz` (via `r-universe`)

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Title Pharmacometric Visualization for the `pharmac` Ecosystem

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Author Malek Okour

Maintainer Malek Okour <malek@zajaly.com>

Description One-line pharmacometric diagnostic plots with modern aesthetics. Creates goodness-of-fit panels, spaghetti plots, and publication-ready exports with consistent theming for pharmacometric workflows.

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URL <https://github.com/pharmac-org/pharmac>,
<https://pharmac-org.github.io/pharmac>

BugReports <https://github.com/pharmac-org/pharmac/issues>

Depends R (>= 4.1.0)

Imports cli (>= 3.6.0), ggplot2 (>= 3.5.0), patchwork, rlang (>= 1.1.0)

Suggests testthat (>= 3.0.0), knitr, rmarkdown

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VignetteBuilder knitr

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Repository <https://pharmac-org.r-universe.dev>

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Contents

px_colors	2
px_forest	2
px_gof	3
px_npde	4
px_plot_grid	4
px_residuals	5
px_save	6
px_spaghetti	6
px_vpc	7
scale_color_pharmac	8
theme_pharmac	8

Index	9
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px_colors	<i>pharmac color palette</i>
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Description

pharmac color palette

Usage

```
px_colors(n = 8)
```

Arguments

n Number of colors.

Value

Character vector of hex colors.

px_forest	<i>Forest plot for estimates and intervals</i>
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Description

Forest plot for estimates and intervals

Usage

```
px_forest(
  data,
  term = "term",
  estimate = "estimate",
  lower = "lower",
  upper = "upper",
  reference = 0,
  mode = "light"
)
```

Arguments

data	Data frame with estimate and interval columns.
term	Term label column.
estimate	Estimate column.
lower	Lower interval column.
upper	Upper interval column.
reference	Reference line value.
mode	Theme mode passed to theme_pharmac() .

Value

A ggplot2 object.

px_gof	<i>Goodness-of-fit diagnostic panel</i>
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Description

Creates a four-panel diagnostic plot: DV vs PRED, DV vs IPRED, CWRES vs TIME, and CWRES vs PRED.

Usage

```
px_gof(data, log_scale = TRUE, mode = "light")
```

Arguments

data	Data frame with at least DV and PRED columns.
log_scale	Use log10 axes for observed-vs-predicted panels.
mode	Theme mode passed to theme_pharmac() .

Value

A patchwork plot object.

px_npde	<i>NPDE diagnostic panel</i>
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Description

NPDE diagnostic panel

Usage

```
px_npde(  
  data,  
  npde = "NPDE",  
  time = "TIME",  
  prediction = "PRED",  
  mode = "light"  
)
```

Arguments

data	Data frame containing an NPDE-like column.
npde	NPDE column name.
time	Optional time column.
prediction	Optional prediction column.
mode	Theme mode passed to theme_pharmac() .

Value

A patchwork plot object.

px_plot_grid	<i>Compose pharmac plots</i>
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Description

Compose pharmac plots

Usage

```
px_plot_grid(  
  ...,  
  ncol = 2,  
  title = NULL,  
  subtitle = NULL,  
  caption = NULL,  
  mode = "light"  
)
```

Arguments

...	ggplot2 or patchwork plots, or a single list of plots.
ncol	Number of columns.
title	Optional grid title.
subtitle	Optional grid subtitle.
caption	Optional grid caption.
mode	Theme mode passed to theme_pharmac() .

Value

A patchwork plot object.

px_residuals	<i>Residual diagnostic panel</i>
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Description

Creates residual-vs-time, residual-vs-prediction, histogram, and QQ panels.

Usage

```
px_residuals(
  data,
  residual = NULL,
  time = "TIME",
  prediction = "PRED",
  mode = "light"
)
```

Arguments

data	Data frame containing residual columns.
residual	Residual column. Auto-detects CWRES, IWRES, then RES.
time	Time column used for residual-vs-time when available.
prediction	Prediction column used for residual-vs-prediction when available.
mode	Theme mode passed to theme_pharmac() .

Value

A patchwork plot object.

px_save	<i>Save a pharma plot</i>
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Description

Save a pharma plot

Usage

```
px_save(plot, filename, width = 8, height = 6, dpi = 300, ...)
```

Arguments

plot	A ggplot2 or patchwork plot.
filename	Output filename.
width	Width in inches.
height	Height in inches.
dpi	Resolution.
...	Additional arguments passed to <code>ggplot2::ggsave()</code> .

Value

The input filename, invisibly.

px_spaghetti	<i>Individual PK concentration-time profiles</i>
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Description

Individual PK concentration-time profiles

Usage

```
px_spaghetti(  
  data,  
  group = NULL,  
  log_y = FALSE,  
  facet_by = NULL,  
  highlight_ids = NULL,  
  mode = "light"  
)
```

Arguments

data	Data frame with ID, TIME, and DV columns.
group	Optional grouping variable for color.
log_y	Use a log10 y-axis.
facet_by	Optional column to facet by.
highlight_ids	Optional vector of subject IDs to highlight.
mode	Theme mode passed to <code>theme_pharmac()</code> .

Value

A ggplot2 object.

px_vpc	<i>Lightweight visual predictive check</i>
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Description

Lightweight visual predictive check

Usage

```
px_vpc(
  data,
  time = "TIME",
  observed = "DV",
  simulated = "SIM",
  bins = NULL,
  probs = c(0.05, 0.5, 0.95),
  mode = "light"
)
```

Arguments

data	Data frame with observed and simulated values.
time	Time column.
observed	Observed concentration column.
simulated	Simulated concentration column.
bins	Number of time bins. If NULL, exact time values are used.
probs	Quantiles to show for observed and simulated values.
mode	Theme mode passed to <code>theme_pharmac()</code> .

Value

A ggplot2 object.

scale_color_pharmac	<i>pharmac color scale for ggplot2</i>
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Description

pharmac color scale for ggplot2

Usage

```
scale_color_pharmac(...)
```

```
scale_fill_pharmac(...)
```

Arguments

... Arguments passed to `ggplot2::discrete_scale()`.

theme_pharmac	<i>The pharmac ggplot2 theme</i>
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Description

A clean, publication-ready theme for pharmacometric graphics.

Usage

```
theme_pharmac(base_size = 12, base_family = "", mode = c("light", "dark"))
```

Arguments

base_size Base font size.

base_family Base font family.

mode Theme mode, either "light" or "dark".

Value

A ggplot2 theme object.

Examples

```
library(ggplot2)
ggplot(mtcars, aes(wt, mpg)) +
  geom_point() +
  theme_pharmac()
```

Index

`ggplot2::discrete_scale()`, 8
`ggplot2::ggsave()`, 6

`px_colors`, 2
`px_forest`, 2
`px_gof`, 3
`px_npde`, 4
`px_plot_grid`, 4
`px_residuals`, 5
`px_save`, 6
`px_spaghetti`, 6
`px_vpc`, 7

`scale_color_pharmac`, 8
`scale_fill_pharmac`
 (`scale_color_pharmac`), 8

`theme_pharmac`, 8
`theme_pharmac()`, 3–5, 7