

Model Diagnostics with xpose :: CHEAT SHEET



The **xpose** package facilitates the creation of model diagnostics from NONMEM output. Inspired by **xpose4**, this new version is actively being redesigned around the popular tidyverse packages **ggplot2**, **dplyr** and **readr**.

Getting started

INSTALLATION

- From CRAN
`install.packages('xpose')`
- From github (development version)
`library(devtools)`
`install_github('UUPharmacometrics/xpose')`

GETTING HELP

- Comprehensive documentation and examples are available at: uupharmacometrics.github.io/xpose/
- Use `?<function_name>` in R to access functions' help (e.g. `?xpose_data`).

PLOT TYPE

Plot type is specified via a single string, where values: **a** (area), **d** (density), **h** (histogram), **l** (line), **p** (point), **r** (rug), **s** (smooth) and **t** (text) can be combined depending on the plot function.

```
dv_vs_ipred(xpdb_ex_pk, type = 'pls')  
eta_distrib(xpdb_ex_pk, type = 'hdr')
```

PLOT LAYERS

All **ggplot2** functions can be used to add or modify **xpose** plot layers, mapping, labels, scales, annotations, etc.

```
plot <- dv_vs_ipred(xpdb_ex_pk)  
plot + geom_hline(yintercept = 1)
```

PIPES

All **xpose** functions can be used with pipes (%>%)

```
xpdb_ex_pk %>%  
filter(OCC == 3) %>%  
dv_vs_ipred()
```

Acknowledgements: the development of xpose was jointly funded by Pfizer and Pharmetheus

Plot functions

BASIC GOF

Accepted plot types: **l**, **p**, **s**, **t**
Layer names: **guide**, **line**, **point**, **smooth**, **text**, **xscale**, **yscale**

```
dv_vs_ipred(xpdb, guide = TRUE)  
dv_vs_pred(xpdb, guide = TRUE)
```

```
res_vs_idv(xpdb, res = 'CWRES', guide = TRUE)  
res_vs_pred(xpdb, res = 'CWRES', guide = TRUE)
```

```
absval_res_vs_idv(xpdb, res = 'CWRES')  
absval_res_vs_pred(xpdb, res = 'CWRES')
```

```
dv_vs_idv(xpdb, group = 'ID')  
ipred_vs_idv(xpdb, group = 'ID')  
pred_vs_idv(xpdb, group = 'ID')
```

```
dv_preds_vs_idv(xpdb, group = 'ID')  
display of DV, IPRED and PRED side by side
```

INDIVIDUAL PLOTS

Accepted plot types: **l**, **p**, **s**, **t**
Layer names: **line**, **point**, **smooth**, **text**, **xscale**, **yscale**

```
ind_plots(xpdb)
```

COMPARTMENT KINETICS

Accepted plot types: **l**, **p**, **s**, **t**
Layer names: **line**, **point**, **smooth**, **text**, **xscale**, **yscale**

```
amt_vs_idv(xpdb, group = ID)  
uses A1, A2, ..., columns by default
```

Customize plots

THEMES

The xpdb objects contain two types of themes :
• **gg_theme**: sets plot background and text properties

```
theme_readable()  
light grey
```

```
theme_bw()  
black and white
```

• **xp_theme**: sets default aesthetics values (e.g. points color, lines width)

```
theme_xp_default()  
ggplot2 default
```

```
theme_xp_xpose4()  
xpose4 blue
```

New themes can be applied globally:

```
xpdb <- update_themes(xpdb, gg_theme, xp_theme)
```

Or locally in each plot function:

```
dv_vs_ipred(xpdb, gg_theme, xp_theme)
```

The **xpdb** (xpose database) is a structured object containing the NONMEM output tables, output files, the parsed model code, general options and plot themes.

VISUAL PREDICTIVE CHECKS

Accepted plot types: **a**, **l**, **p**, **r**, **t**
Layer names: **area**, **line**, **point**, **rug**, **text**, **xscale**, **yscale**

```
compute  
vpc data in R
```

```
advanced options  
(e.g. bins, lloq, pi, ci, etc.)
```

```
vpc_data(xpdb, opt = vpc_opt(...), vpc_type,  
stratify, psn_folder) %>%  
vpc(smooth) plot the vpc
```

DISTRIBUTIONS

Accepted plot types: **d**, **h**, **r**
Layer names: **density**, **histogram**, **rug**, **xscale**, **yscale**

```
prm_distrib(xpdb)  
eta_distrib(xpdb)  
cov_distrib(xpdb)  
res_distrib(xpdb, res = 'CWRES')
```

QQ PLOTS

Accepted plot types: **p**
Layer names: **guide**, **point**

```
prm_qq(xpdb, guide = TRUE)  
eta_qq(xpdb, guide = TRUE)  
cov_qq(xpdb, guide = TRUE)  
res_qq(xpdb, res = 'CWRES', guide = TRUE)
```

MINIMIZATION DIAGNOSTICS

Accepted plot types: **l**, **p**, **s**, **t**
Layer names: **line**, **point**, **smooth**, **text**, **xscale**, **yscale**

```
grd_vs_iteration(xpdb) (.grd file required)  
prm_vs_iteration(xpdb) (.ext file required)
```

Data

IMPORT

Data import in **xpose** is structured as follows:

- Read NONMEM control stream (.mod/.lst) to list table filenames for each \$PROBLEM
- Import and index tables (compatible with FIRSTONLY option, .csv and compressed (.zip) files)
- Import NONMEM output files (.ext, .phi, .cov, etc.)
- Summarize control stream

Runs can automatically be imported either by using the file or the prefix, runno and ext arguments.
`xpdb <- xpose_data(dir, file, prefix, runno, ext)`

EDIT

Data in the xpdb can be edited using **dplyr** functionalities

- `filter(xpdb, ...)`
subset data based on logical condition(s)
- `mutate(xpdb, ...)`
add, modify and remove columns
- `set_var_type(xpdb, ...)`
assign or modify output tables' index

`xpdb_ex_pk %>%` built-in xpdb example
`mutate(TAD = TIME %% 24) %>%` e.g. generate and plot time after dose
`dv_vs_idv(aes(x = TAD))`

ACCESS

Access and extract data from an xpdb.

- `get_code(xpdb, .problem)` (parsed control stream)
- `get_prm(xpdb, .problem)` (table of parameter estimates)
- `get_file(xpdb, ext, .problem)` (parsed output files)
- `get_data(xpdb, .problem)` (combined dataset)
- `get_summary(xpdb, .problem)` (table of run summary)

SUMMARY

- `print(xpdb)` or `xpdb`
display xpdb structure
- `list_vars(xpdb)`
display data variables
- `summary(xpdb)`
display run summary
- `prm_table(xpdb)`
display parameter table

Template titles

Special @<keywords> can be used in plot labels. They are automatically replaced by their actual value in the run summary when plotting (e.g. title = 'ofv: @ofv' can give 'ofv: -1518.108'). Check `?template_titles` in R for a full list.

Save plots

The `file` and `dir` arguments can contain template titles' keywords. Also handles plots with multiple pages.
`xpose_save(plot, file = '@run_@plotfun.pdf', dir)`