

# Package ‘Certara.ModelResults’

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**Title** Generate Diagnostics for Pharmacometric Models Using 'shiny'

**Version** 3.0.1

**Description** Utilize the 'shiny' interface to generate Goodness of Fit (GOF) plots and tables for Non-Linear Mixed Effects (NLME / NONMEM) pharmacometric models. From the interface, users can customize model diagnostics and generate the underlying R code to reproduce the diagnostic plots and tables outside of the 'shiny' session. Model diagnostics can be included in a 'rmarkdown' document and rendered to desired output format.

**Depends** R (>= 4.0)

**License** LGPL-3

**URL** <https://certara.github.io/R-model-results/>

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.3.2

**Suggests** knitr, rmarkdown, testthat (>= 3.0.0), Certara.RsNLME

**Imports** colourpicker, shinyAce, shinymeta, Certara.Xpose.NLME, xpose, dplyr, flextable, shinyjqui, grDevices, ggplot2, plotly, magrittr, scales, shiny (>= 1.7.0), shinyjs, shinyWidgets, shinyTree (>= 0.3.1), sortable, tidyverse, rlang, bslib (>= 0.7.0)

**Config/testthat.edition** 3

**NeedsCompilation** no

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`get_eps_shk`      *Get eps shrinkage values xpdb*

### Description

This function returns eps shrinkage values from xpdb object as a `data.frame`.

### Usage

```
get_eps_shk(xpdb)
```

### Arguments

`xpdb`      Object of class `xpose_data`.

### Value

Returns an object of class `data.frame`.

### Examples

```
get_eps_shk(xpdb_NLME$TwCpt_IVBolus_FOCE_ELS)
```

`get_eta_shk`      *Get eta shrinkage values xpdb*

### Description

This function returns eta shrinkage values from xpdb object as a `data.frame`.

### Usage

```
get_eta_shk(xpdb)
```

**Arguments**

xpdb Object of class `xpose_data`.

**Value**

Returns an object of class `data.frame`.

**Examples**

```
get_eta_shk(xpdb_NLME$TwCpt_IVBolus_FOCE_ELS)
```

---

resultsUI

*Generate and Report Model Diagnostics from NLME or NONMEM runs*

---

**Description**

Shiny application to generate, customize, and report diagnostic plots and tables from NLME or NONMEM output files. Create an Rmarkdown file of tagged model diagnostics and render into submission ready report.

**Usage**

```
resultsUI(model, xpdb = NULL, tagged = NULL, settings = NULL, ...)
```

**Arguments**

`model` A single object, vector, or list of objects of class `NlmePmlModel`.  
`xpdb` A single object or list of objects of class `xpose_data`.  
`tagged` List of tagged objects returned from previous tagged <- `resultsUI()` session.  
`settings` List of settings (e.g., `settings.Rds`) returned from previous Shiny session.  
`...` Additional arguments for Pirana integration.

**Value**

If `interactive()`, returns a list of tagged diagnostics from the Shiny application, otherwise returns `TRUE`.

## Examples

```

if (interactive()) {

  # RsNLME
  library(Certara.RsNLME)
  library(Certara.ModelResults)

  model1 <- pkmodel(numCompartments = 1,
                     data = pkData,
                     ID = "Subject",
                     Time = "Act_Time",
                     A1 = "Amount",
                     CObs = "Conc",
                     modelName = "OneCpt_IVBolus_FOCE-ELS")

  baseFitJob1 <- fitmodel(model1)

  model2 <- pkmodel(numCompartments = 2,
                     data = pkData,
                     ID = "Subject",
                     Time = "Act_Time",
                     A1 = "Amount",
                     CObs = "Conc",
                     modelName = "TwCpt_IVBolus_FOCE-ELS")

  baseFitJob2 <- fitmodel(model2)

  # Run Model Results
  resultsUI(model = c(model1, model2))

  # NONMEM via xpose
  library(Certara.ModelResults)
  library(xpose)

  xpdb <- xpose_data(
    runno = "1",
    prefix = "run",
    ext = ".lst",
    dir = "./NONMEM/Hands_onB/")

  resultsUI(xpdb = xpdb)

  # Multiple models

  xpdb_multiple <- list(
    run1 = xpose_data(file = "run1.lst"),
    run2 = xpose_data(file = "run2.lst"),
    run3 = xpose_data(file = "run3.lst"),
}

```

```
run4 = xpose_data(file = "run4.lst")
}
}
```

---

theme\_certara      *A ggplot2 theme for Certara.*

---

## Description

A ggplot2 theme for Certara.

## Usage

```
theme_certara(
  base_size = 11,
  base_family = "",
  base_line_size = base_size/22,
  base_rect_size = base_size/22,
  grid = c("none", "horizontal", "both")
)
```

## Arguments

base\_size      base font size, given in pts.  
base\_family      base font family  
base\_line\_size      base size for line elements  
base\_rect\_size      base size for rect elements  
grid      Which grid lines should appear? Horizontal only, both horizontal and vertical, or none (default). [continuous\\_scale\(\)](#).

## Details

There are 3 variants of the theme: no grid `theme_certara()`, full grid `theme_certara(grid = "both")`, and horizontal grid lines only `theme_certara(grid = "horizontal")`.

## Value

An object of class [theme\(\)](#).

<code>write_code</code>	<i>Write code to R script from tagged diagnostics</i>
-------------------------	---

**Description**

Use this function to write code to R script from diagnostics tagged in Certara's Model Results Shiny Application.

**Usage**

```
write_code(tagged, file)
```

**Arguments**

- |                     |   |
|---------------------|---|
| <code>tagged</code> | List of tagged objects from returned from <code>resultsUI()</code> .  |
| <code>file</code>   | Character specifying path of output file. If missing, it will be saved as <code>code.R</code> in working directory. |

**Value**

Returns `NULL` after writing to `file`.

**Examples**

```
if (interactive()) {
  tagged_diagnostics <- resultsUI(xpdb = xpdb_NLME)

  write_code(tagged_diagnostics, "tagged_results.R")

}
```

<code>xpdb_NLME</code>	<i>List of xpose data objects from RsNLME example tutorials</i>
------------------------	---

**Description**

The following object contains a list of 2 `xpose_data` objects generated in the `RsNLME` example script `TwoCptIVBolus_FitBaseModel_CovariateSearch_VPC_BootStrapping.R`.

**Usage**

```
xpdb_NLME
```

**Format**

List of 2 `xpose_data` objects constructed from NLME model output.

**Details**

- `xpdb_NLME$`TwCpt_IVBolus_FOCE-ELS`` is an `xpose_data` object created from the base model in RsNLME example script. The model can be used as a reference to compare model diagnostics in final model.
- `xpdb_NLME$`TwCpt_IVBolus_SelectedCovariateModel_FOCE-ELS`` is an `xpose_data` object created from the final model in the RsNLME example script. The final model includes selected covariate BodyWeight added from the results of stepwise covariate search.

**Source**

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xpdb\_NONMEM*List of xpose data objects from NONMEM model output*

---

**Description**

The following object contains of list of 2 `xpose_data` objects:

**Usage**`xpdb_NONMEM`**Format**

List of 2 `xpose_data` objects constructed from NONMEM model output.

**Details**

- `xpdb_NONMEM$ex_pk` is an `xpose_data` object from `xpose::xpdb_ex_pk`. The model contains multiple covariates and can be used to explore covariate model diagnostics.
- `xpdb_NONMEM$mult_obs` is an `xpose_data` object created from NONMEM model with multiple observed variables. Users will see that appropriate model diagnostic plots are automatically faceted by DVID in the Shiny GUI.

**Source**

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