

Data Exploration Libraries in KIWI

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The cloud-based KIWI[™] platform is an enterprise-scale application supporting the execution, management, and communication of modeling and simulation activities.¹ Exploration of analysis datasets and model outputs is a key component of these activities as it allows the identification of data patterns and guides the creation and execution of modeling plans and supports communication of modeling results within the development team or with regulatory agencies who rely on exploratory displays to convey the rationale of decision-making or to illustrate problems and knowledge gaps that should be further investigated. The creation of meaningful exploratory plots and tables is a time-consuming task even for skilled programmers.

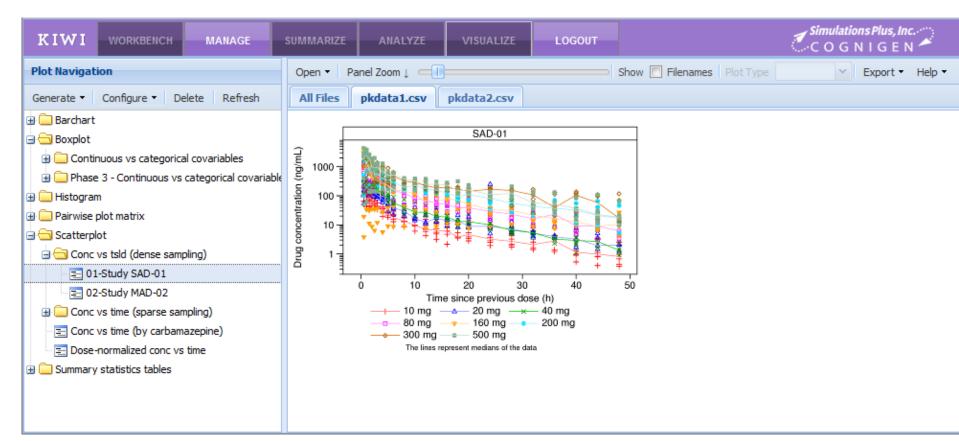
KIWI includes multiple modules providing user-friendly graphical interfaces for efficient and flexible data exploration and reporting:

Visualize was presented at a previous PAGE meeting¹ and provides a library focused on the automated creation of model diagnostic plots.

Explore and ExploreLive are 2 new modules providing libraries of generic tools to automate the creation of summary statistics tables and plots. It was designed to explore the content of analysis datasets and model output tables.

All displays created in KIWI are generated with templatized R code² and do not require any programming experience on the part of the users.

EXPLORE FUNCTIONALITY



Shared Functionality

record per subject)

Advanced layout settings

Reference and spline lines

Custom title and footnote

Axis zoom, log scale, and

variability (SD/SE/CI)

Secondary Y axis

transformation

Single or multiple display downloads

Overlay of multiple Y axis variables

Central tendency (median/mean) ±

Scatter and Line Plots

Explore shares many elements of functionality with the Visualize module, offering seamless visual integration of displays created by both modules:

Variable categorization (eg, AGE is continuous, RFCAT is categorical)

Data subset using custom R expressions or predefined conditions (eg, first)

Creation of additional combined plots when paneling and grouping is used

1000 -

100 -

10

Library of reusable settings profiles within and across projects

R code libraries for data extraction, processing, and plotting

Library of variable format collections (eg, 0: male; 1: female)

Data grouping using different colors, symbols, and/or line styles

Five methods of handling below the lower limit of quantitation data

Library of variable labels and units (eg, AGE: age [y])

Shared functionality across exploratory types in Explore:

Multivariable paneling and stratification of data

EXPLORELIVE FUNCTIONALITY ----

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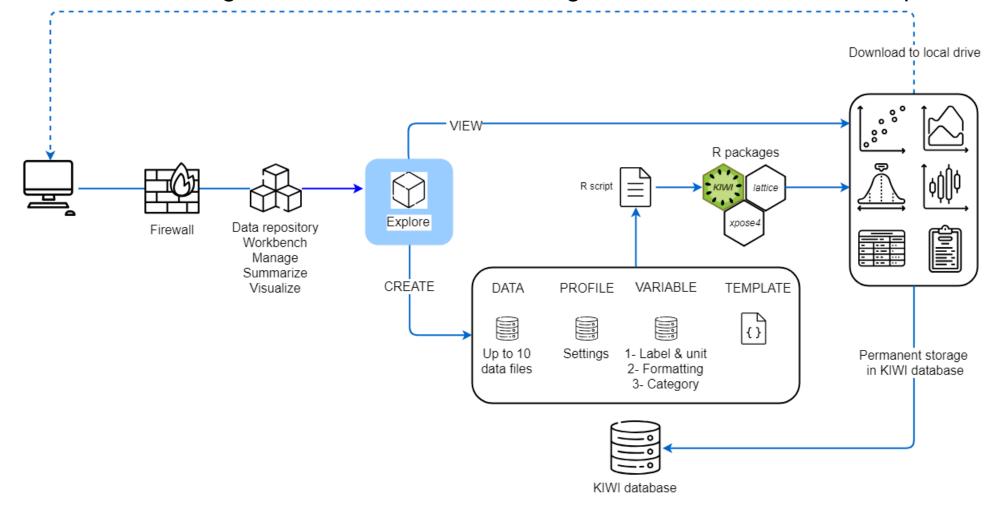
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Shared Functionality

Shared functionality across ExploreLive sub-modules: Data subset

DESIGN OVERVIEW Explore

Explore is a module intended for the creation and permanent storage of report-quality tables and lattice-based plots.³ R scripts are automatically generated and executed using reusable profiles of settings selected in the graphical interface. All outputs are stored in a database and tied to a unique dataset identifier assuring traceability and reproducibility. Execution of these custom profiles typically generates multiple displays which are automatically organized in a hierarchical tree format for convenient navigation and comparison across multiple data sources. Explore leverages the graphical standards and data formatting features established for model diagnostic plotting in the Visualize module, offering seamless visual integration in technical reports.

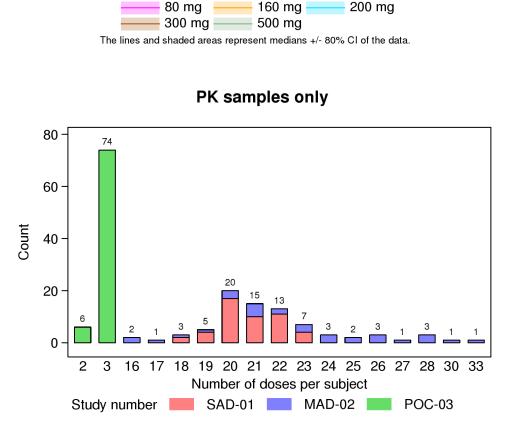


ExploreLive

ExploreLive is a Shiny application intended as a sandbox environment for the live and nonpermanent manipulation of data, creation of tables and interactive ggplot2-based plots, and execution of small-scale data analysis.^{4,5} ExploreLive is remotely hosted and rendered in the user's web browser. All displays are generated on the server side based upon the user-selected options and rendered in the application. The Shiny architecture implements swift reactivity between user input changes and the update of the display rendered on the screen. Interactive features allow users to click on plots and obtain relevant information about the data source. Displays can be downloaded to a local drive. Upon exiting the application, all settings and displays are discarded on the server and web browser sides.

Barchart

- Single-variable count
- Multiple binning methods for continuous X-axis variable
- Count of record number per individual



Study SAD-01

Time since previous dose (h)

—— 40 mg

- Data grouping using different colors, symbols, and/or line styles
- Multivariable paneling and stratification of data
- Reference lines
- Download

Data

- Provide dataset for use in other sub-modules
- Subset using custom R expressions or predefined conditions (eg, first record) per subject)

Sort

Add and transform variables

Pool and merge datasets

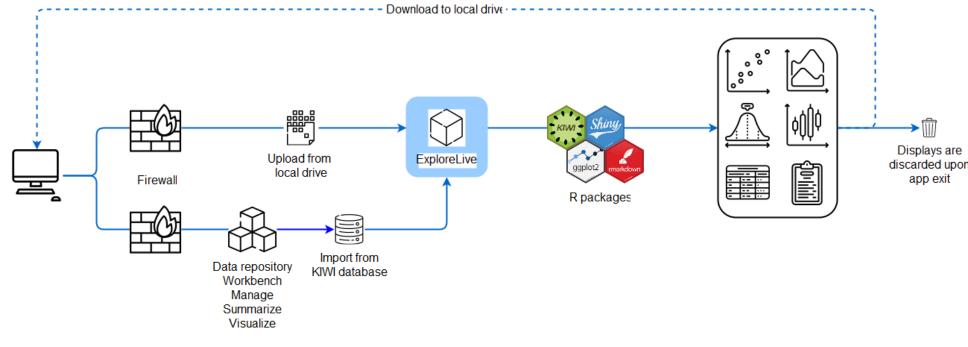
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Scatter and Line Plots

- Interactive plot with data highlight and query
- Axis zoom, log scale, and transformation
- Reference and spline lines
- Central tendency (median/mean) ± variability (SD/SE/CI)

Single-variable count or Y versus X variable (with or without error bar)

Barchart



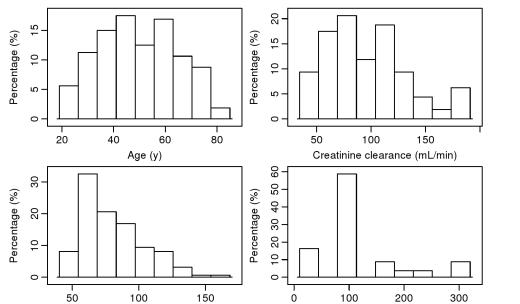
Y versus X variable

Distribution Plot

variables at once

Density, count, or percentage

Histogram created for multiple



Dose (mg)

Quick Comparison of Functionality

Feature	Explore	ExploreLive
Secure access	\checkmark	\checkmark
Data source	KIWI database	KIWI database Upload from local drive
Number of datafiles that can be explored simultaneously	Up to 10	1 (3 data files loaded)
Data manipulation	×	\checkmark
Data subset	\checkmark	\checkmark
Plot engine	Lattice	ggplot2
Scatter and line plots	\checkmark	\checkmark
Box-and-whisker plot	\checkmark	\checkmark
Barchart	\checkmark	\checkmark
Histogram	\checkmark	✓
Pairwise plot matrix	\checkmark	✓
Kaplan-Meier plot	×	\checkmark
Summary statistics table	\checkmark	\checkmark
Contingency table	\checkmark	\checkmark
Data binning plot	×	\checkmark
Linear regression	×	\checkmark
Interactive plot	×	\checkmark
Permanent setting storage	\checkmark	×
Permanent display storage	\checkmark	×
Permanent R script storage	\checkmark	×
Display organization	\checkmark	×
Multi-format export (png, jpg, pdf, docx, xlsx, html)	\checkmark	\checkmark

Boxplot

Boxplots created for multiple pairs of continuous and categorical variables at once

Mini statistics report associated with each boxplot

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			Gender							Dose	(mg))		

Weight (kg)

es represent the 25th and 75th percentiles and lines the median. Notches provide an approximate 95% C.I. abo median. Whiskers extend to the most extreme values within the 1.5 interquartile range. Values outside this range are arked with open circles or subject ID. The numbers of included values are displayed above each box.

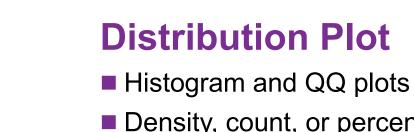
r=-0.423

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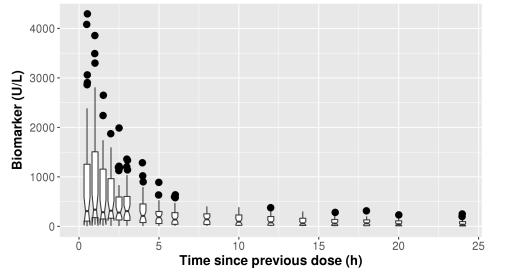
r=-0.0271

Pairwise Plot Matrix

- Co-distribution between multiple continuous or categorical variables
- Automated detection of high correlation between continuous variables



- Density, count, or percentage Choice of distribution and kernel functions **Boxplot**
 - Interactive plot with data highlight and query Box and whiskers for continuous or categorical X-axis variable



Linked Plots

- Up to 4 scatter or box-and-whisker plots
- Cross-interactivity between plots with data highlight
- Axis zoom, log scale, and transformation

Pairwise Plot Matrix

Co-distribution between multiple continuous or categorical variables

Summary Statistics and Contingency Table

- Custom selection of statistics
- Three different layouts for summary statistics tables

Survival Plots

- Multiple estimators
- Number at risk and censoring panels
- Stratification by multiple continuous and categorical predictors with

Strata + sex=1 + sex=2

REFERENCES

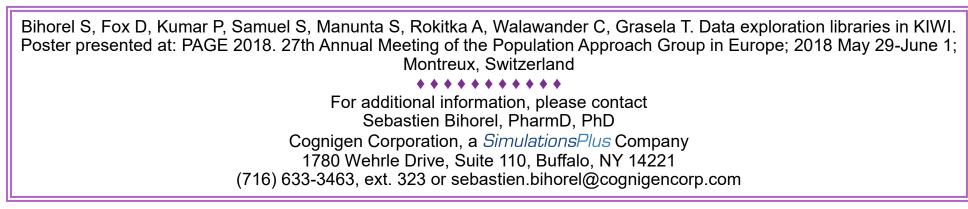
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- 2. R Core Team (2018). R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria. https://www.R-project.org/.
- 3. Sarkar D. (2008) Lattice: Multivariate Data Visualization with R. Springer, New York. ISBN 978-0-387-75968-5.
- 4. Wickham H. (2009) ggplot2: Elegant Graphics for Data Analysis. Springer-Verlag, New York.
- 5. Chang W, et al. Shiny: Web Application Framework for R. https://CRAN.R-project.org/ package=shiny.
- *Some icons used in this poster were made by Freepik from www.flaticon.com.



Summary Statistics and Contingency Tables

- Custom selection of statistics
- Coercion of continuous variables into categorical variables
- Four-table layout for summary statistics tables, including 1 listing individual variable value + summary statistics

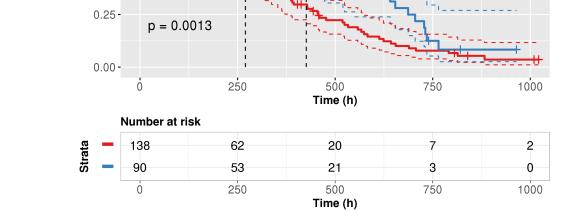
Superprofiles are coming...



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•	49.9	62.1	42.3	46.5	53.7	Missing	35.2	25.3	13.0
ŀ	95.6	112	113.0	79.3	81.7	Missing	35.5	Missing	15.7
i	65.7	69.8	72.4	Missing	66.3	57.3	43.1	27.1	27.9
;	Missing	41.9	45.6	55.2	47.3	43.9	26.0	Missing	20.8
n (SD)	81.8 (23.1)	83.1 (36.4)	79.1 (30.4)	76.1 (34.4)	59.8 (12.4)	44.4 (9.38)	32.8 (8.47)	23.1 (3.9)	18.9 (6.07)
Max	49.9, 105	41.9, 130	42.3, 113	46.5, 134	47.3, 81.7	34.9, 57.3	19.7, 43.1	18.4, 27.1	12.8, 27.9
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av background indicates an absolute correlation coefficient (r) value > 0

calculation of p-value



Regression, data binning, and more to come...

SUMMAR

Explore and ExploreLive naturally extend the functionality of the KIWI platform and provide powerful and user-friendly tools for data exploration. The library of displays provided by Explore enables busy scientists and those with no programming experience to quickly generate meaningful tables and plots to support their pharmacometric needs. By leveraging the infrastructure and features previously established, Explore maintains consistency and reproducibility in the design of tables and plots across projects and between exploratory and model-diagnostic displays. ExploreLive offers a convenient environment for quick data exploration and investigation of data patterns.