

## Introduction, Background, Motivation

Data exploration is a key first step of any data analysis. Often an exploratory plot can quickly answer a question of interest and could be used in place of more complex analyses or to inform next steps.

In this cheat sheet, we provide a list of assessments to perform during exploratory analysis of exposure-response (PK, PD and PKPD) data in order to encourage a structured approach to data exploration.

### Key Message

Don't just look at your data – look at it in a structured way

### Objectives

- ✓ Provide structured approach for purposeful exploration of PKPD data
- ✓ Provide a teaching tool for exploring PKPD data with R
- ✓ Improve efficiency and code readability for exploratory analyses
- ✓ Improve quality of exploratory PKPD graphics

## Data Checking

A first important step to data exploration is to check the quality of your data. For PK/PD data, we recommend to check for:

- incorrect timing of dosing and observations (e.g. nonzero pre-dose PK observations)
- erroneous duplication of data
- large discrepancies between actual time and nominal time (i.e. per-protocol time of sample collection)
- outlying data points

## PK, Dose-Exposure

Get an overview of the PK data by plotting a summary measure of central tendency +/- variability (e.g. mean +/- SE, [mean \(95% CI\)](#)), over time, grouped by dose or assigned treatment (Figure A, B)

Q How many compartments does the PK appear to have?

Q Do you detect any nonlinear clearance (e.g. dramatic drop in elimination phase on log scale)?

Both linear and log scale should be used for exploring PK data. **See section on Technical Considerations (Scales)** for further details.

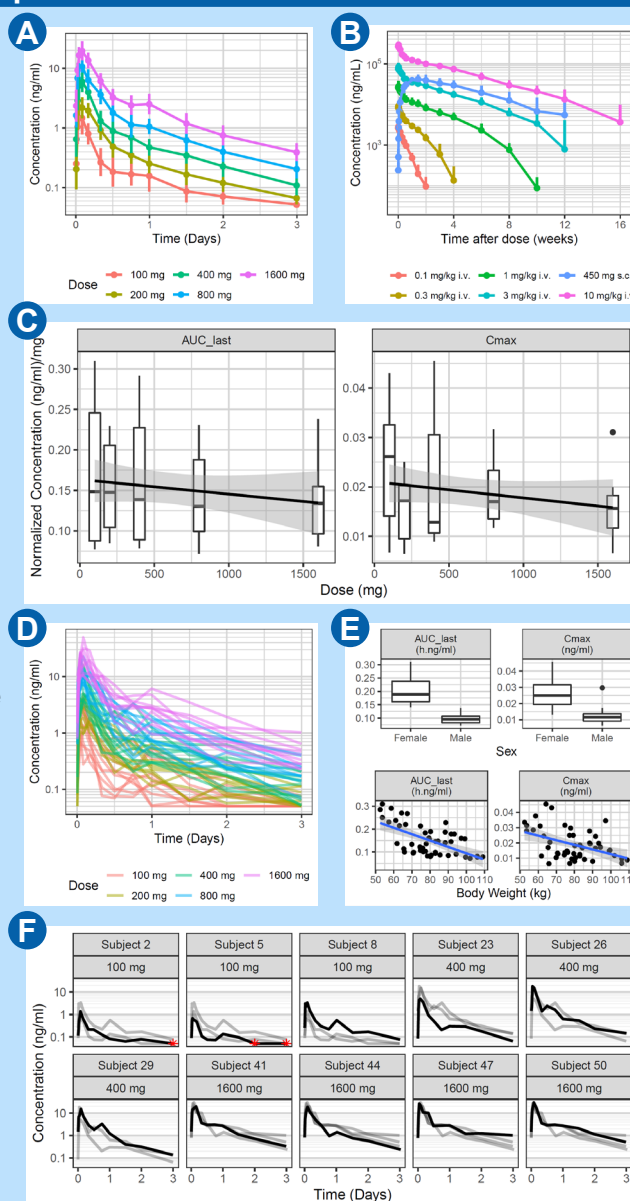
[Assess PK linearity](#) by looking for trends in the plot of dose vs dose normalized PK metric (e.g. AUC or Ctrough) (Figure C)

- Assessing linearity of PK is important for understanding how dose adjustments will impact the response

Assess extent and sources of variability

- Assess size of variability by using [spaghetti plots](#) or [confidence intervals](#) (Figure D)
- [Assess sources of between subject variability](#) by stratifying by covariates (Figure E)
- [Assess sources of within subject variability](#) using individual plots (Figure F)

**See section on Technical Considerations (Variability)** for details on within vs between subject variability, explained vs unexplained variability, and consequences of high variability.



## PD, Efficacy, Safety, Dose-Response

Determine data type of the endpoint of interest, and choose appropriate exploratory plots for this data type, (e.g. [continuous](#), [binary](#), [ordinal](#)).

**See section on Technical Considerations for:**

- Determining whether any correction is needed (e.g. change from baseline, placebo adjusted).
- Choosing the appropriate scale for your endpoint.

Assess trends with dose: Summary of PD/efficacy/safety vs dose, (e.g. [Mean \(95% CI\) of PD vs dose](#)) (Figure G, H)

Q Do you see a relationship between dose and response?

Q Is a plateau observed at higher doses?

Q What would you guess the ED50 to be? ED90? (you can check your expectations against your future model)

Assess trends over time by plotting [summary plots of the endpoint against time](#) (Figure I, J)

Q Does the response change over time?

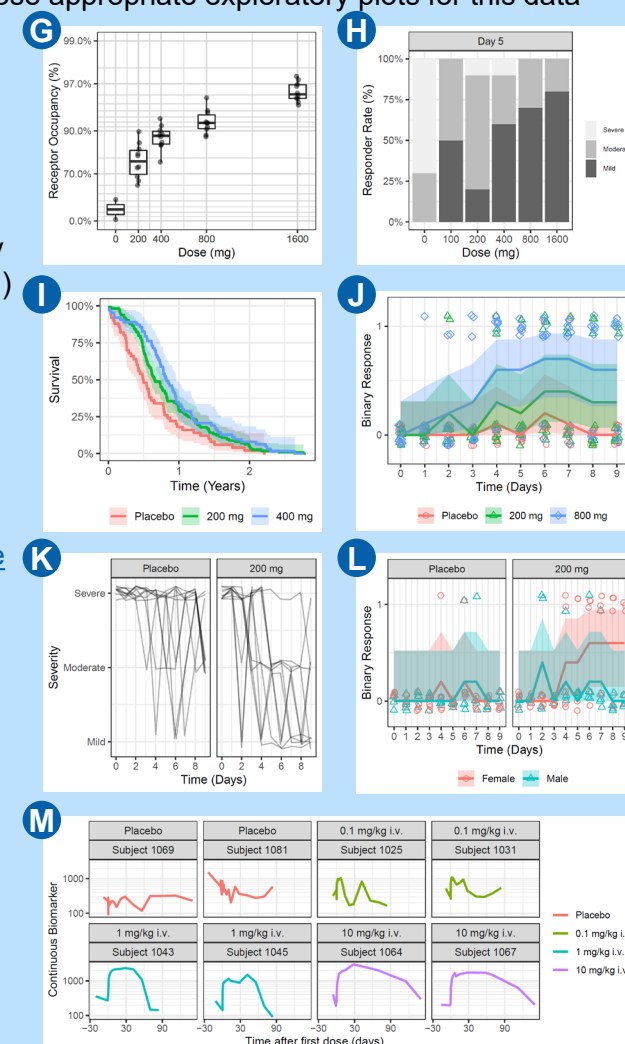
Q Is tolerance observed (e.g. rebounding, returning to baseline, overshooting)?

Q How long does it take the PD to reach steady state (if there is a steady state)?

Assess extent and sources of variability

- Assess size of variability by using [spaghetti plots](#) or [confidence intervals](#) (Figure K)
- [Assess sources of between subject variability](#) by stratifying by covariates (Figure L)
- [Assess sources of within subject variability](#) using individual plots (Figure M)

**See section on Technical Considerations (Variability)** for further details



## PKPD, Exposure-Response/Safety

Plot [PK and PD on the same time scale](#) to get an idea of the trends of both over time. Look at both summary plots and individual plots (Figure N)

Q How long is the delay between changes in PK and changes in PD?

Q How long after steady state PK does PD reach steady state (if at all)?

Get an overall idea of the relationship between PK and PD by plotting [PD on the vertical axis against different PK metrics on the horizontal axis](#) (Figure O)

Q Is it a positive relationship?

Q How strong is the relationship?

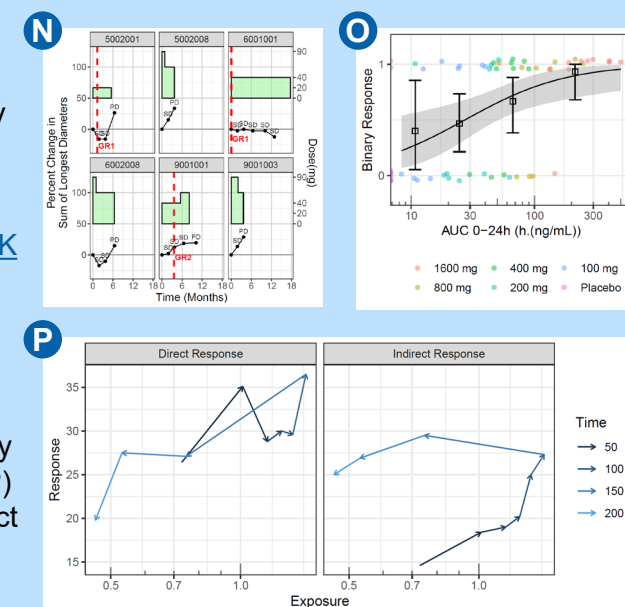
Q Is there a lot of between subject variability?

Get an idea of the time delay of the PKPD relationship by plotting [individual hysteresis plots of PD vs PK](#) (Figure P)

Q Is the observed relationship between PK and PD direct (straight line in hysteresis path) or indirect (looping behavior in the hysteresis path)?

Be aware of situations that can impact interpretation, e.g.:

- Dose interruptions
- Intercurrent events (e.g. nonresponder drop-out before responders)



## Technical Considerations

### Variability

Variability is not the same as noise. Variability is a characteristic of living systems and may in itself be a signal if appropriately represented (e.g. covariate effects, diurnal effects, disease progression). Variability has two key characteristics:

- 1) Inter-subject (between subject) vs Intra-subject (within subject)
- 2) Explained vs Unexplained

The table below shows examples of the different types of variability, and mitigation steps for high variability in each category.

| Type of Variability                     | Examples   | Mitigation Steps  |
|---|--|---|
| Explained between subject variability   | Traditional covariates (e.g. age, weight, sex, race)                   | Individualized therapy may be an option to reduce between subject variability   |
| Unexplained between subject variability | Unaccounted for covariates   | Therapeutic drug monitoring could be attempted to adjust dose based on observed PK and/or PD during therapy   |
| Explained within subject variability    | Circadian rhythms, seasonal effects, food effects, disease progression | Depending on the explanation, you might suggest dosing accordingly  |
| Unexplained within subject variability  | Residual error, poor absorption, other unaccounted for effects         | PK: could be difficult to address, unless there is a large therapeutic window in which case it may not be an issue. There may be a need for reformulation of the product.<br>PD: may require multiple and/or appropriately timed measures of the endpoint and/or baseline in order to get a good idea of the “true” drug effect. Protocol assessment schedules for future studies should be designed accordingly. |

### PD Data corrections (e.g. change from baseline, placebo adjusted, fold normal)

Consider whether correcting by a reference value will provide a clearer representation of your data (see table below)

| Reference value | Situation that might benefit from data correction   |
|-----------------|---|
| Baseline        | If there is high inter-subject variability and high correlation between baseline value and endpoint           |
| Normal          | If the goal is to compare to normal ranges (e.g. upper limit of normal for lab markers)                       |
| Placebo         | To more clearly reveal drug effect, especially for primary endpoints which are often compared against placebo |

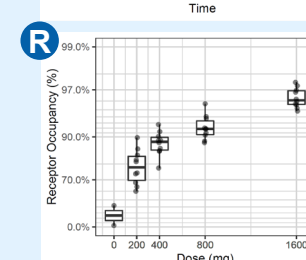
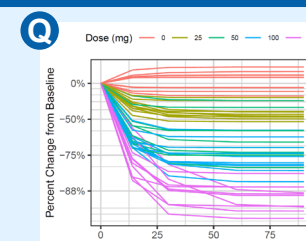
When performing PD data correction:

- Consider whether to use absolute difference ( $y - y^*$ ), ratio to reference ( $y/y^*$ ) or percent change from reference ( $(y - y^*)/y^*$ ).
- Give special care to axis scales and confidence intervals for ratios and percent change from baseline (see section on Scales).
- Give special care when there is high within subject variability, e.g. use multiple baseline values for one individual (see section on Variability).

### Scales

Axis scales should reflect the distribution of data and/or the question being answered

|    |  |
|----|--|
| PK | <ul style="list-style-type: none"> <li>• Log scale helps identify # of compartments, linearity of elimination, &amp; visualize wide range of doses on same plot.</li> <li>• Linear scale focuses attention on C<sub>max</sub>, which may be important for drugs with narrow therapeutic window &amp; C<sub>max</sub>-driven safety.</li> </ul>   |
| PD | <ul style="list-style-type: none"> <li>• Log scale works well for PD markers that can change over several orders of magnitude.</li> <li>• Linear scale is preferred when the PD measure can be both positive or negative, or when there are less than 2 orders of magnitude between the minimum and maximum value.</li> <li>• For percent change from baseline, use the <code>xgx_scale_y_percentchangelog10()</code> function to increase the resolution around -100% (Figure Q)</li> <li>• For receptor occupancy, use the <code>xgx_scale_y_reverselog10()</code> function to increase the resolution around 100% occupancy (Figure R)</li> </ul> |



## Useful Plotting Functions - xgx package

### Plot theme functions

`xgx_theme()` – set the global plotting theme  
`xgx_plot()` – make a plot and set the theme for that one plot

### Tabulation functions

`xgx_check_data()` – provide summary tables that check data  
`xgx_summarize_covariates()` – summarize covariate information

### Plotting functions

`xgx_geom_ci()` – plot mean & confidence intervals under different distribution assumptions (e.g. normal, lognormal, binomial)  
`xgx_geom_pi()` – plot median & percentile intervals  
`xgx_geom_individuals()` – coming soon  
`xgx_geom_spaghetti()` – coming soon

### Plot scaling functions

`xgx_scale_y_log10()` – change y axis to log10 scale  
`xgx_scale_x_time_units()` – convert time units for plotting  
`xgx_scale_y_reverselog10()` – scale y axis nicely for receptor occupancy data, increases resolution around 100%. Scales according to  $-\log_{10}(1-x)$ .  
`xgx_scale_y_percentchangelog10()` – scale y axis nicely for percent change data, increases resolution around -100%. Scales according to  $\log(\text{PCHG} + 100\%)$ .

### Saving and annotating functions

`xgx_annotate_status()` – add draft status watermark to figures  
`xgx_annotate_filenames()` – add metadata to bottom of figures  
`xgx_save()` – save figures including status watermark & metadata  
`xgx_save_table()` – saves table to csv including source metadata

See <https://cran.r-project.org/web/packages/xgxr> for more details

## PKPD Exploratory Graphics Checklist

- ☐ Identify data type and choose appropriate graph types (PD)
- ☐ Identify axis scale that reflects distribution of data (PK, PD)
- ☐ Provide an overview of the data (PK, PD)
- ☐ Determine whether data corrections are needed (PD)
- ☐ Assess trends over time (PK, PD)
- ☐ Assess trends by dose (PD)
- ☐ Assess PK linearity (PK)
- ☐ Assess extent and sources of variability (PK, PD)
- ☐ Get an overview of the relationship between exposure and response (PKPD)
- ☐ Explore delays between exposure and response (PKPD)

## Resources

See <https://github.com/Novartis/xgx/tree/master/Resources> for:

- [Fundamental PK principles introduction](#)
- [Fundamental PD principles introduction](#)
- [Uncertainty Assessment - Pedigree table](#)
- [Graphics Principles Cheat Sheet](#)
- [Presentation check list](#)

