



Modeling & Simulation Learn to Speak the Language: Modeling

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**October 31, 2006
AAPS 2006
San Antonio, TX**

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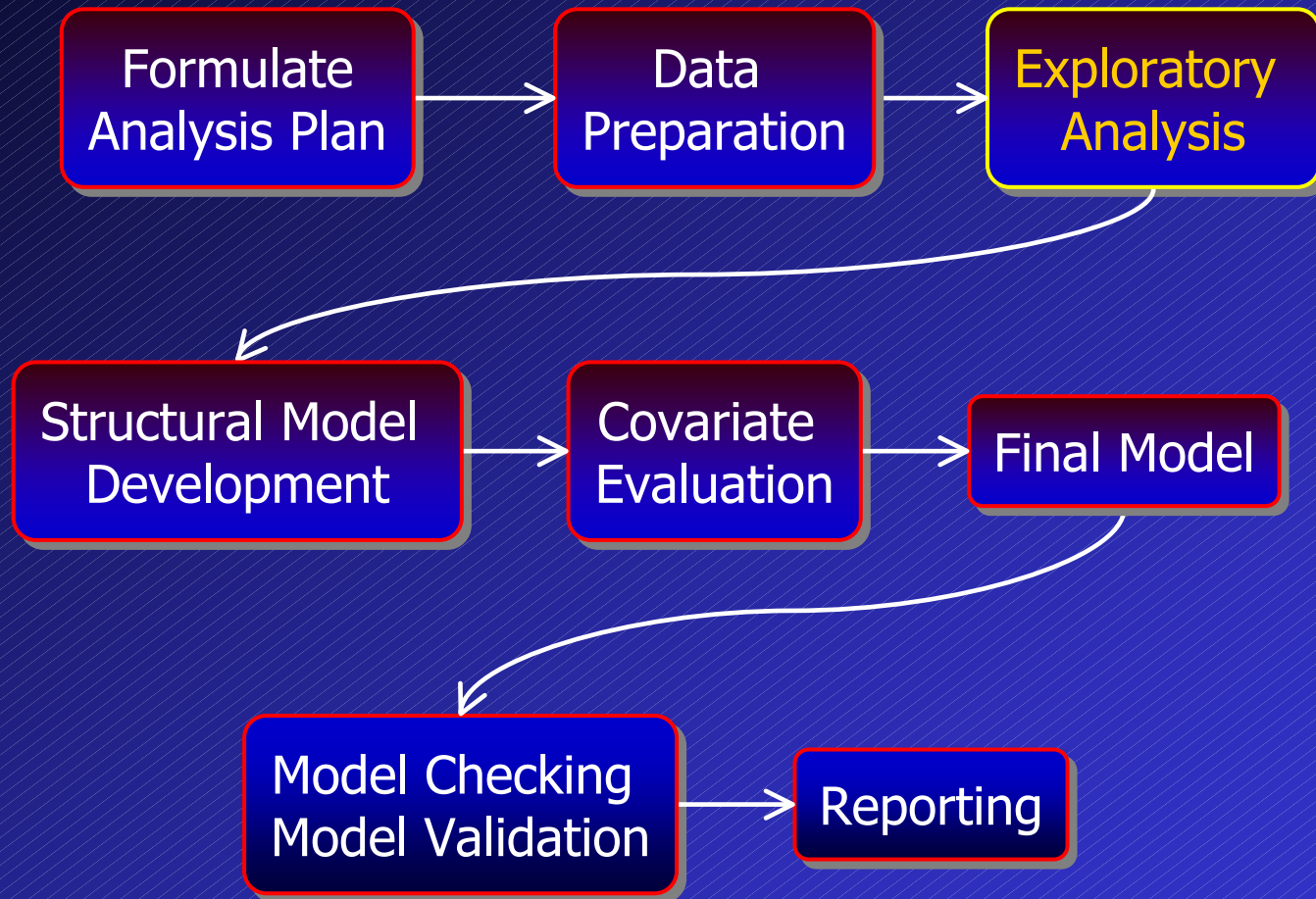
Population PK Methodologies

- **Naïve Pooled Data:**
 - Estimate population mean parameters by treating all data as if it arose from the same individual
- **Standard Two-Stage:**
 - Estimate each individual's PK and then use individual parameters to estimate the population parameters
- **Nonlinear Mixed Effects Modeling (NONMEM®):**
 - Mixed effect modeling allows for the fixed effects (mean values) effects and random quantities (variability) and to be simultaneously quantified.
- **Bayesian Estimation:**
 - Use prior distribution of parameters in a population of subjects and data from an individual to estimate the individual's parameters

Case Study Study Design

- 100 subjects
- Drug X once daily, orally, for 4 weeks
- Clinic Visits:
 - Screen, Baseline, Day 1 of dosing, and Weeks 2, 3, and 4 of dosing.
 - Week 2, 3, and 4 visits were required to occur within a ± 2 day interval of the scheduled day.
- PK Samples:
 - One sample during the clinic visits on Weeks 2, 3, and 4
 - 3 samples per subject
 - Sample Time: anytime during the 24-hour dosing interval

Phases of a Population Analysis

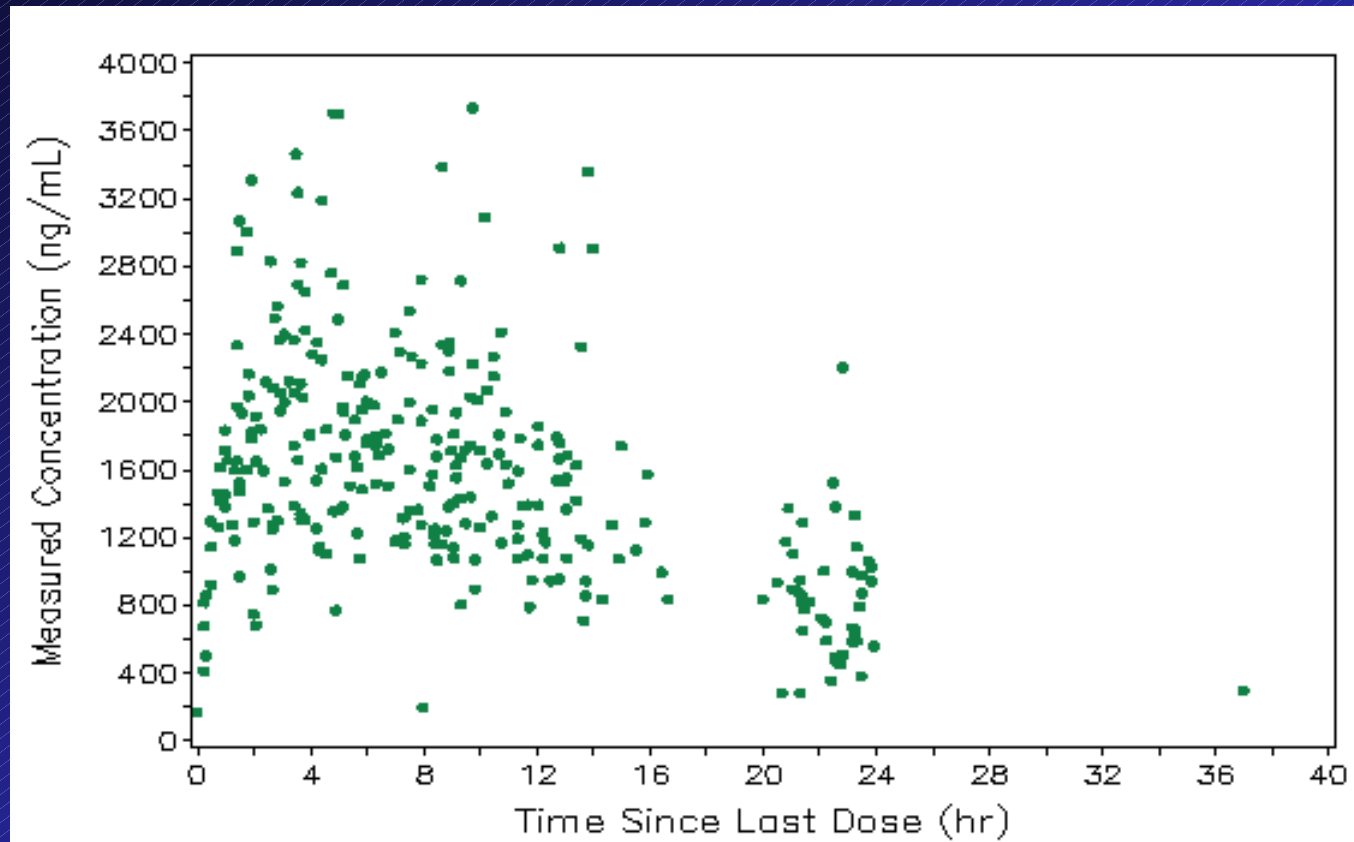


Exploratory Data Analysis (EDA)

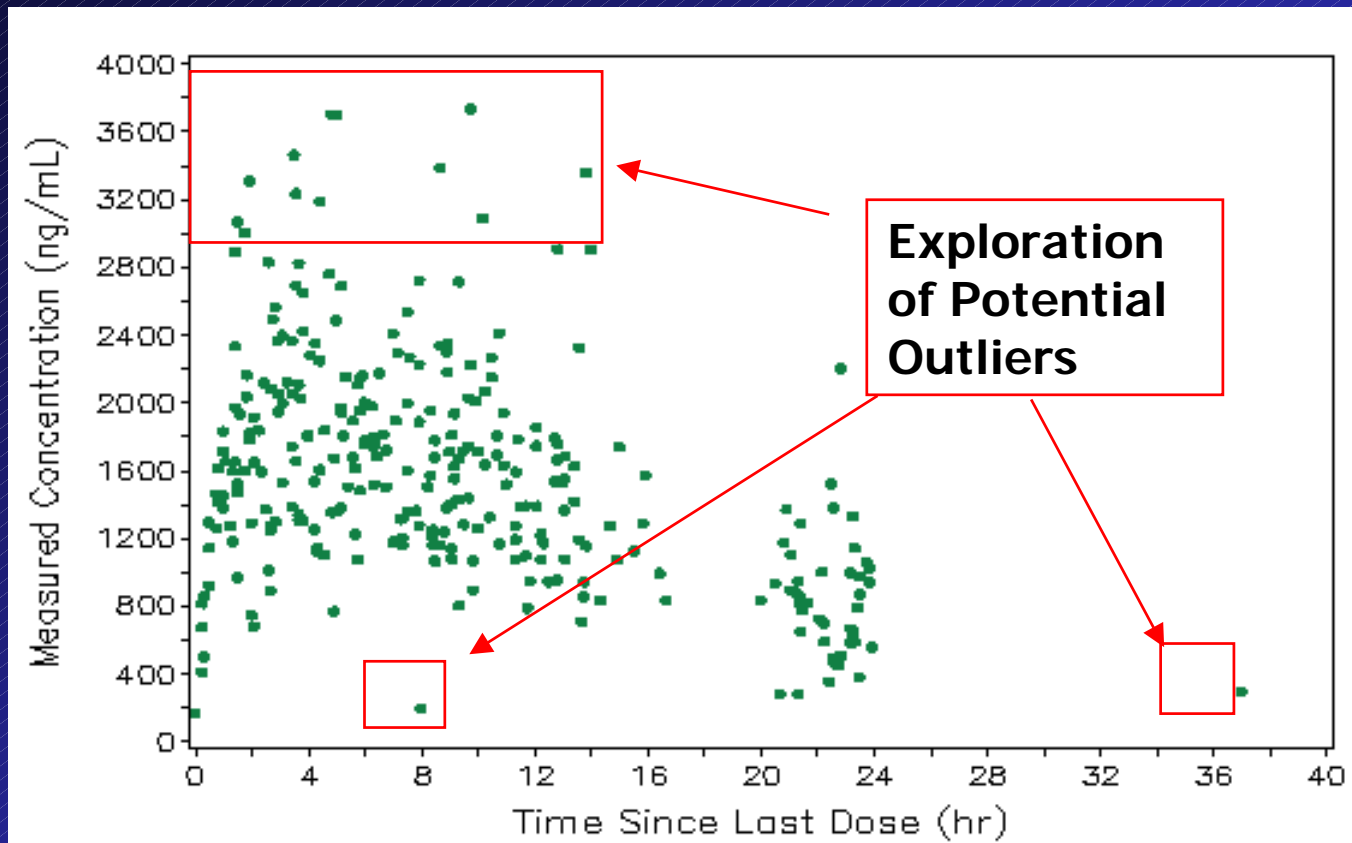
What is it?

- The stage of analysis where you gain an understanding of your data
 - Examples of displays to be created during EDA
 - Scatterplot of Cp (concentration) vs TSLD
 - Histograms of continuous subject characteristics
 - Histograms of the number of subjects, the number of samples per subject, the distribution of time since last dose, etc.
 - Summary statistics of subject characteristics, race, gender, age, weight, race by gender, gender by dose, etc.

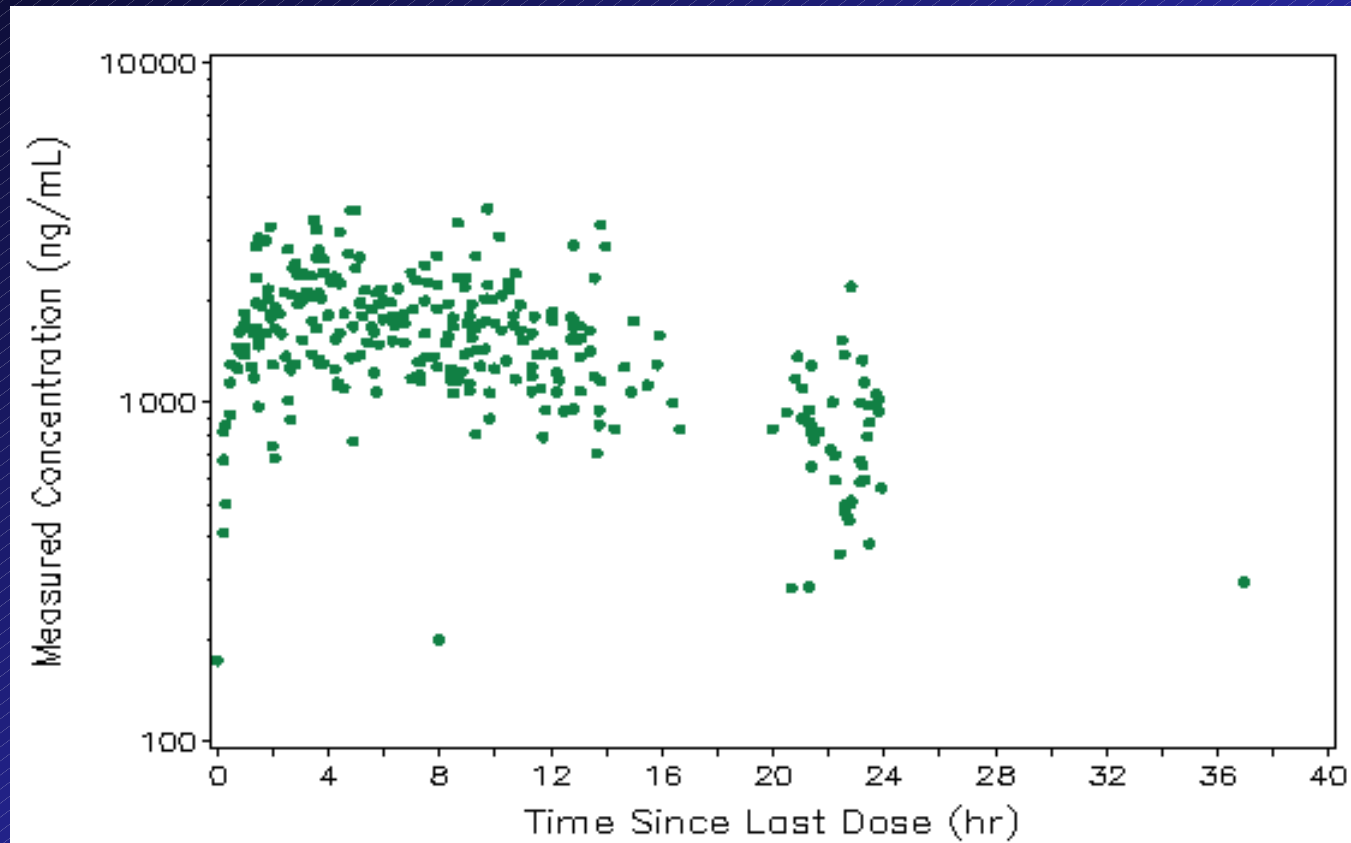
Exploratory Analysis Cp vs TSLD



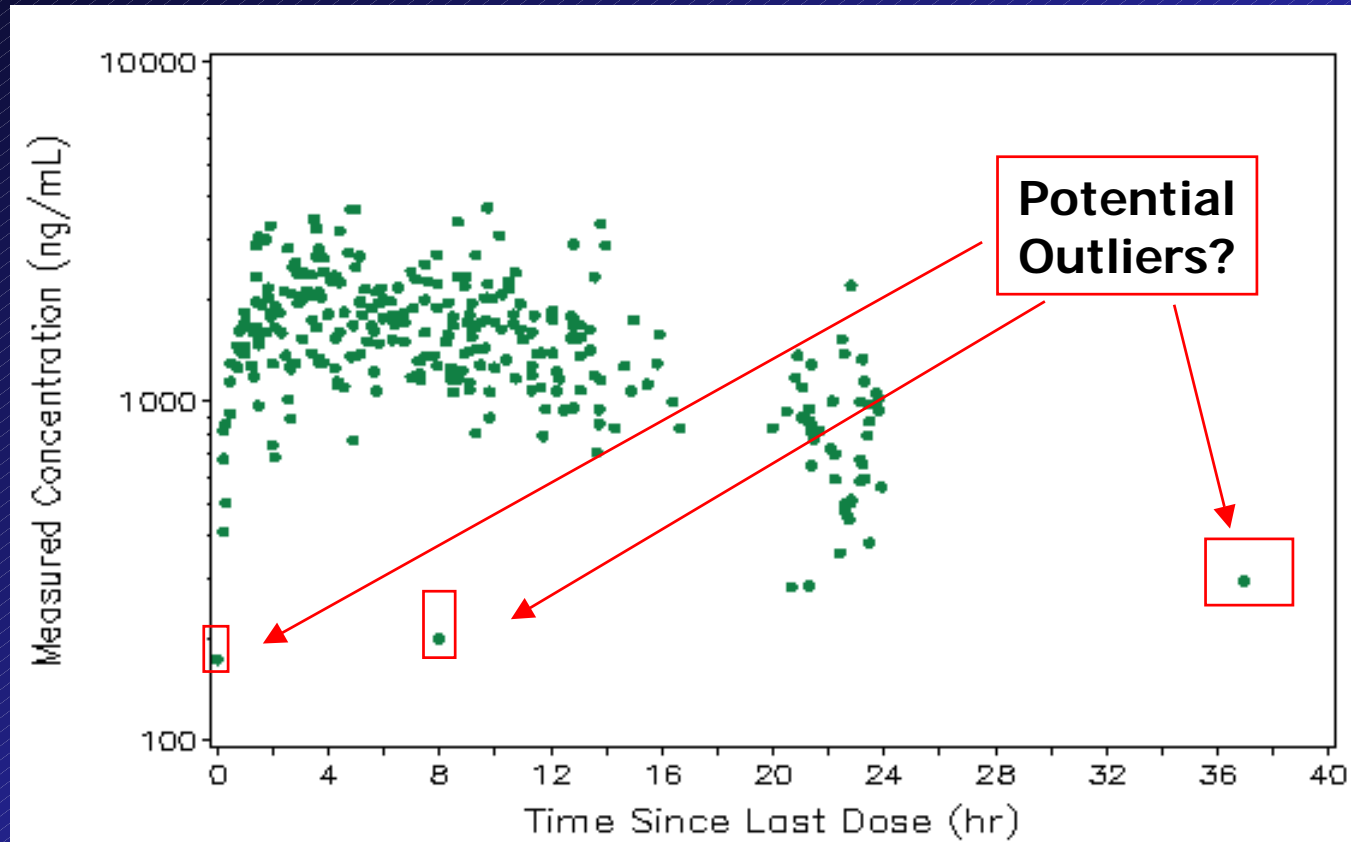
Exploratory Analysis Cp vs TSLD



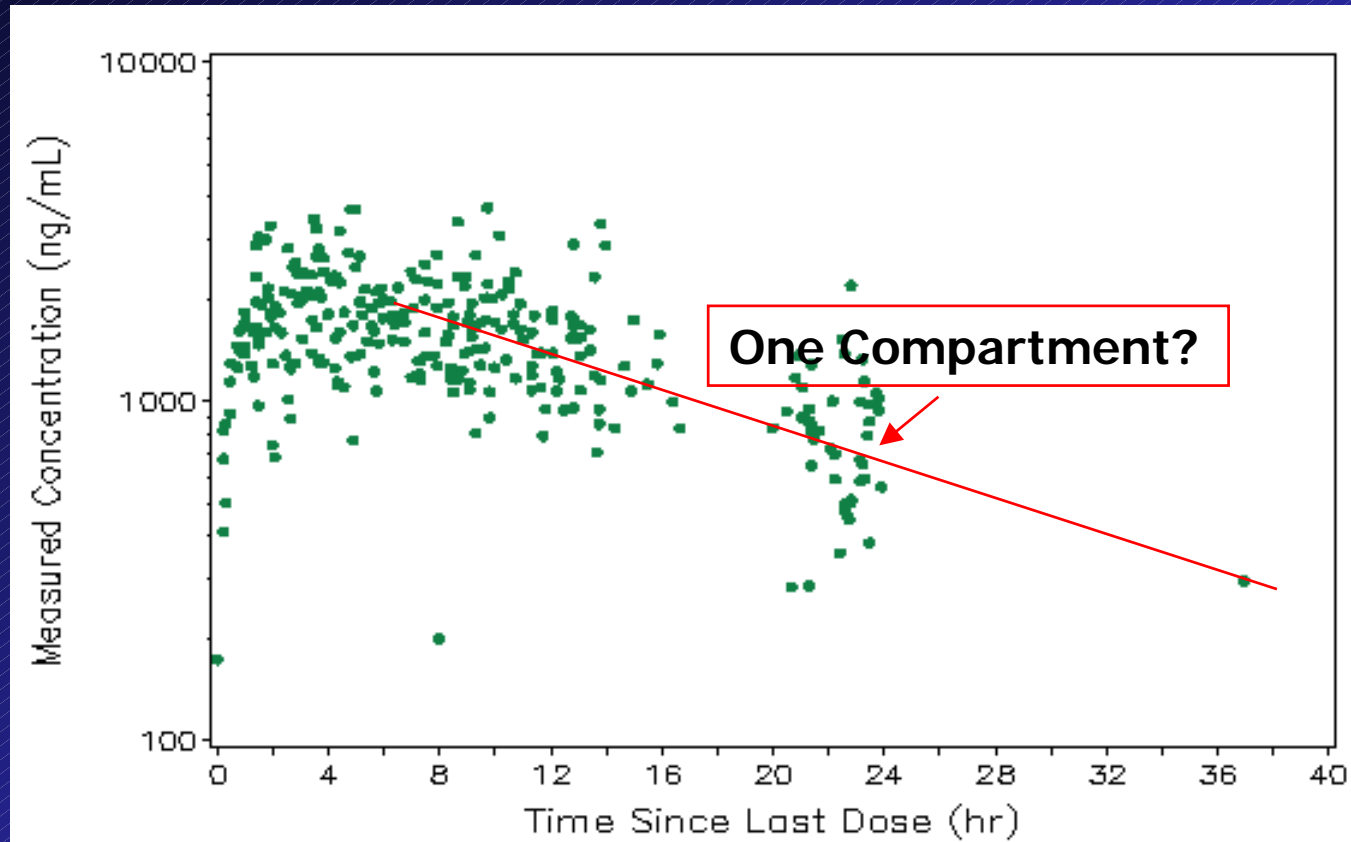
Exploratory Analysis Cp vs TSLD (log Scale)



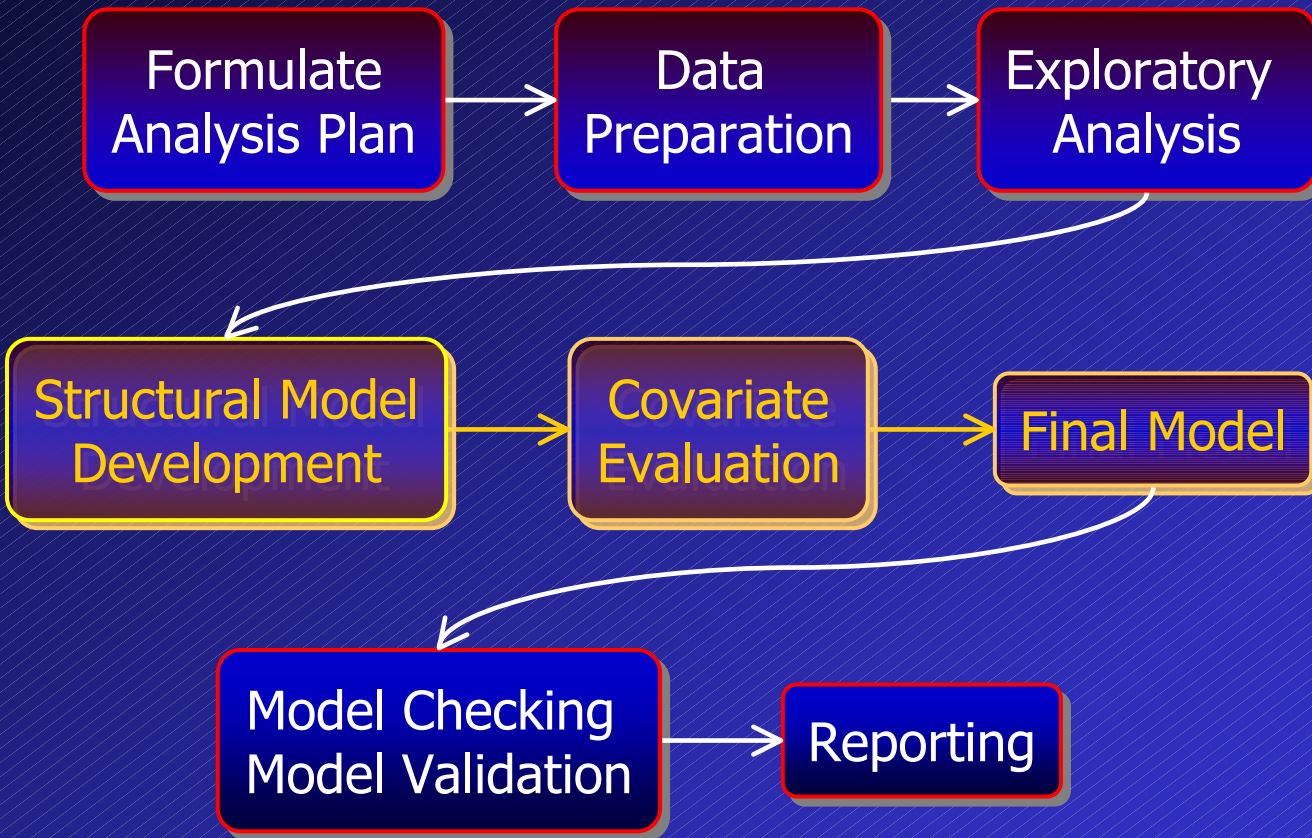
Exploratory Analysis Cp vs TSLD (log Scale)



Exploratory Analysis Cp vs TSLD (log Scale)



Phases of a Population Analysis



Population PK/PD Parameters

- Θ Theta: Fixed Effects
 - Vector of typical values or central tendencies of parameters
 - θ (theta)
- Random Effects (minimum two levels)
 - Ω - Omega: Matrix of interindividual (between subject) variance estimates (IIV)
 - η (eta)
 - Σ - Sigma: Residual Variability
 - Matrix of intraindividual (within subjects) variance estimates
 - ε (epsilon)

Control Stream Terminology

```

|$PROBLEM ex1
|$INPUT ID DATE TIME AMT DV TRT WTKG HTIN AGE SEX EVID MDV PROT
|$DATA /doc/phor/pkpd/nonmem-testsuite/data/ex1.dat
|$SUBROUTINES ADVAN2 TRANS2
|$PK

CALLFL = -2
TVCL = THETA(1)
CL = TVCL * DEXP(ETA(1))
TVV = THETA(2)
V = TVV * DEXP(ETA(2))
TVKA = THETA(3)
KA = TVKA * DEXP(ETA(3))
ALAG1 = THETA(4)
S2 = V/1000

$ERROR

Y = F + EPS(1) + F*EPS(2)

$THETA (0, 28) (0, 90) (0, 3) (0, 0.1)
$OMEGA 0.25 0.25 0.25
$SIGMA 0.02 0.1
$ESTIMATION POSTHOC MAXEVAL=1000 PRINT=5 MSFO=ex1.msf
$COVARIANCE
$TABLE ID DATE TIME AMT TRT CL V KA NOPRINT FILE=ex1.tbl NOHEADER NOPRINT

```

Control Stream Terminology

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```

\$SUBROUTINES:

- Tells which PRED-PP subroutines to use:
- ADVANx for compartment definitions
- TRANSx for parameterization options

Control Stream Terminology

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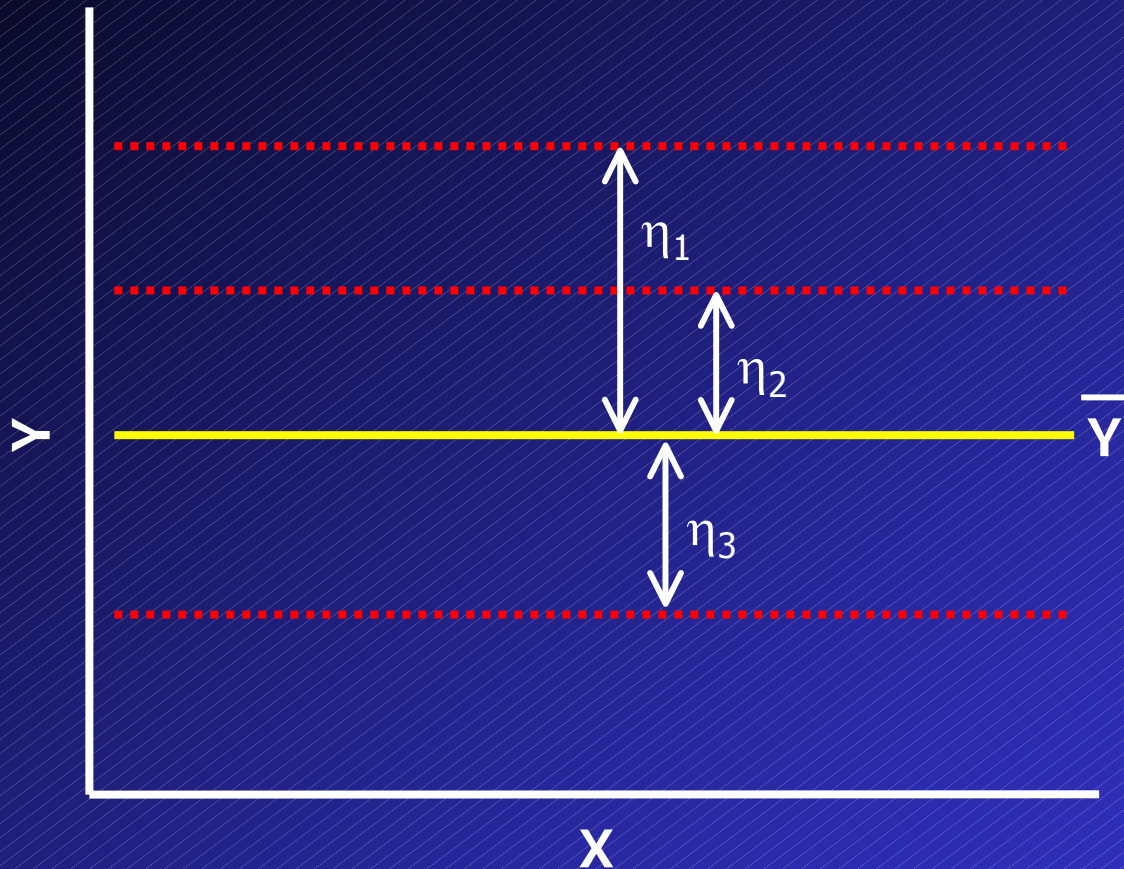
\$THETA:

- Vector of typical values or central tendencies of parameters
- Specifies initial estimates for fixed effect parameters

\$OMEGA and \$SIGMA: Specifies initial estimates of the variance of IIV (omega) and RV (sigma)

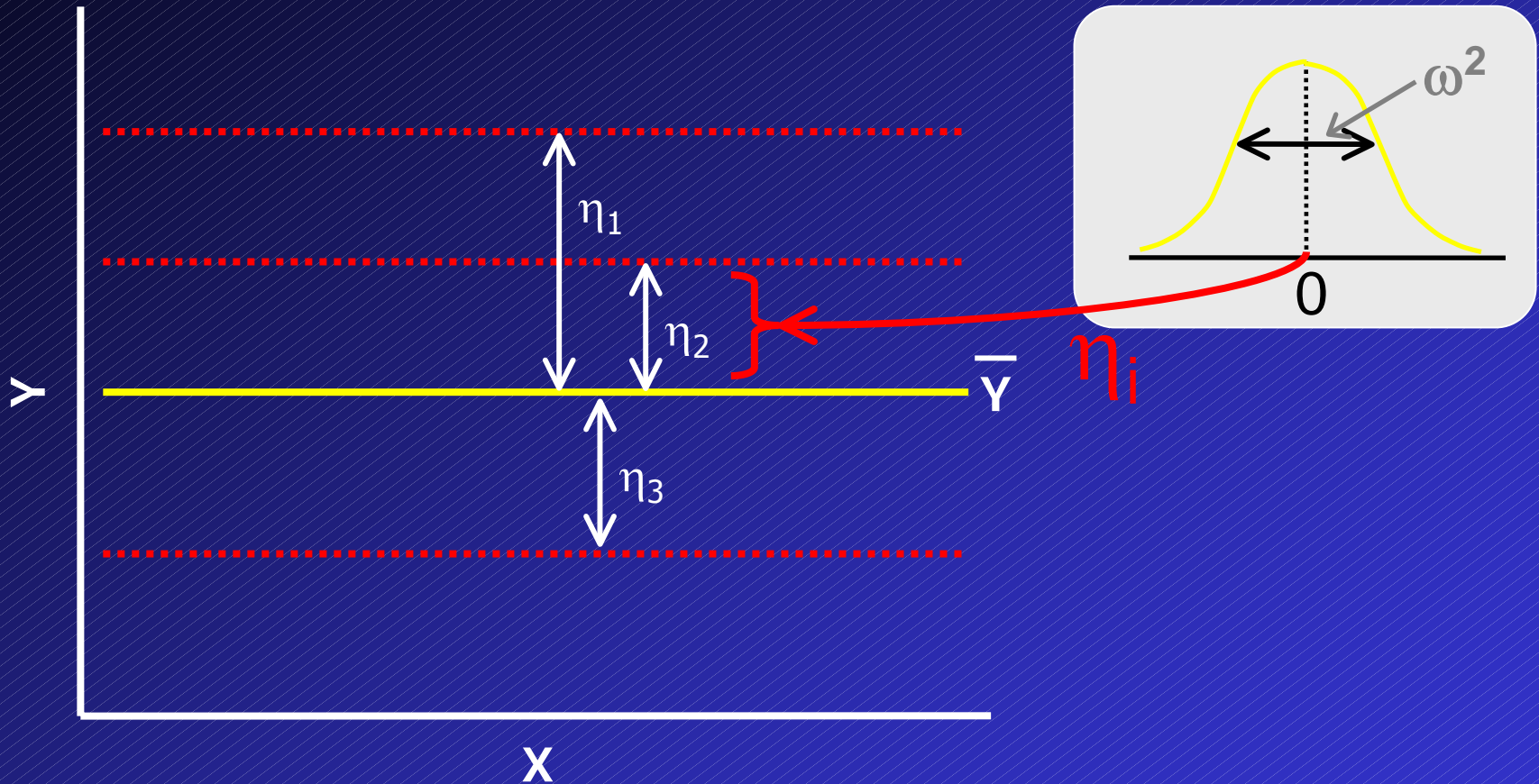
Modeling Population PK

Interindividual Variability (IIV), Between Subject



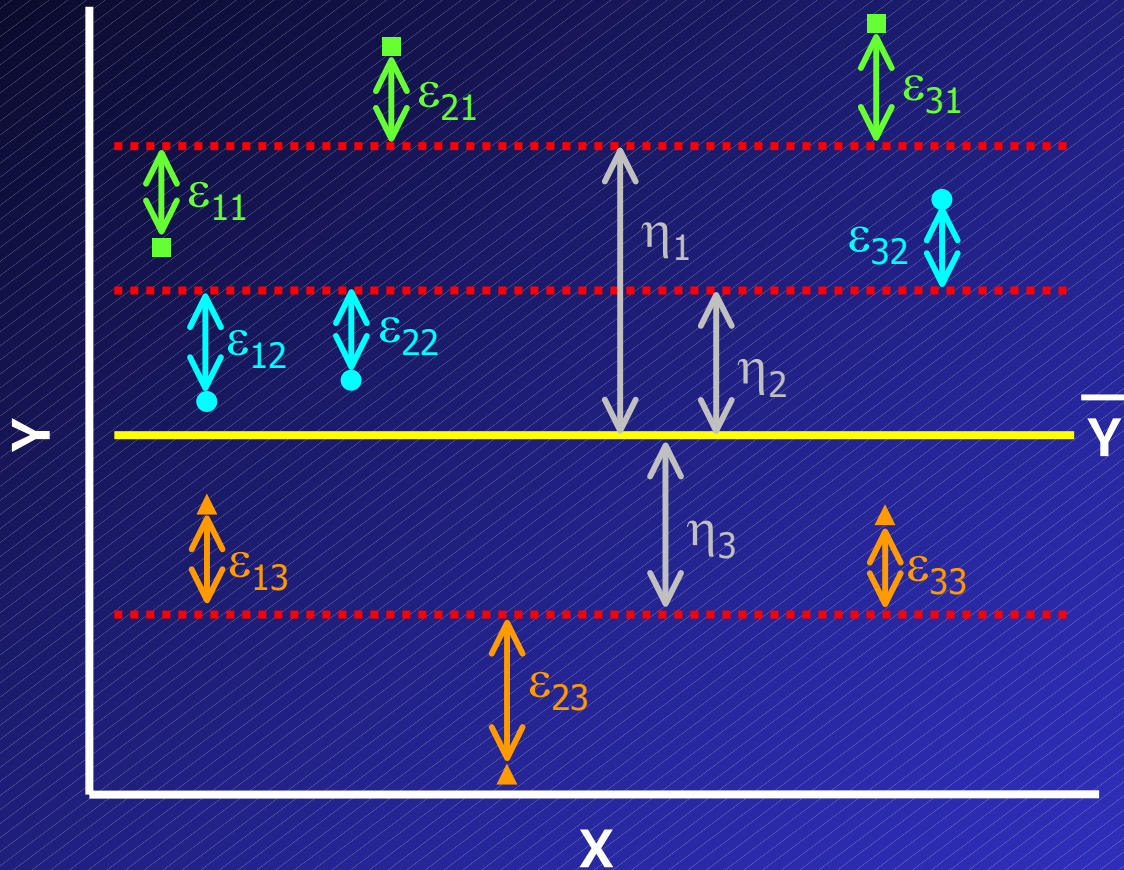
Modeling Population PK

Interindividual Variability (IIV), Between Subject



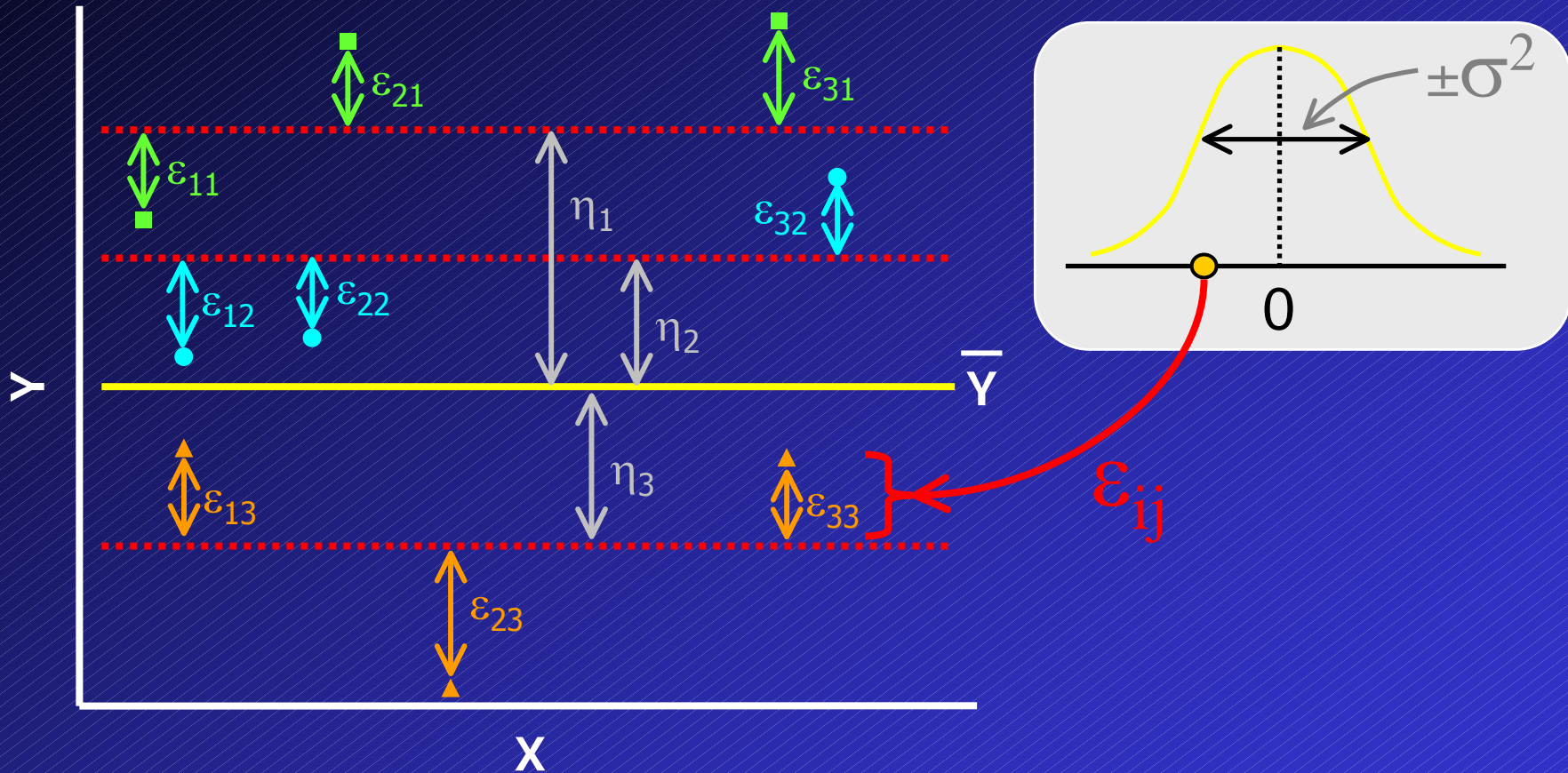
Modeling Population PK

Intraindividual Variability, RV, Within Subject



Modeling Population PK

Intraindividual Variability, RV, Within Subject



Control Stream Terminology

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**\$ESTIMATION: Specifies
the estimation method to
be used**

NONMEM[®]'s FO (First Order) Estimation Method

- Default (and quickest) estimation method
- Named for the linearization method which uses a first-order Taylor series expansion
- **POSTHOC**: option to calculate individual-specific empiric Bayesian estimates of η 's and provides individual-specific estimates of parameters

NONMEM[®]'s FOCE

- First-order conditional estimation (FOCE) method estimates individual η 's during the minimization (no need for POSTHOC)
- much more time-consuming
- **INTERACTION** option allows for interaction between η 's and ε 's
 - FOCE with INTERACTION can correct bias sometimes observed with FO and rich, full-profile data

Base Population PK Model (Prior to Covariate Evaluation)

Parameter	Final Parameter Estimate	
	Population Mean	%SEM
Ka (1/hr)	0.565	13.7
CL (L/hr)	2.93	3.4
V (L)	58.2	7.5
IIV CL (%CV)	30.59	21.9
IIV V(%CV)	17.35	107.0
RV (%CV)	19.29	13.1

Min. Value of the Objective Function (MVOF) = 3863.901

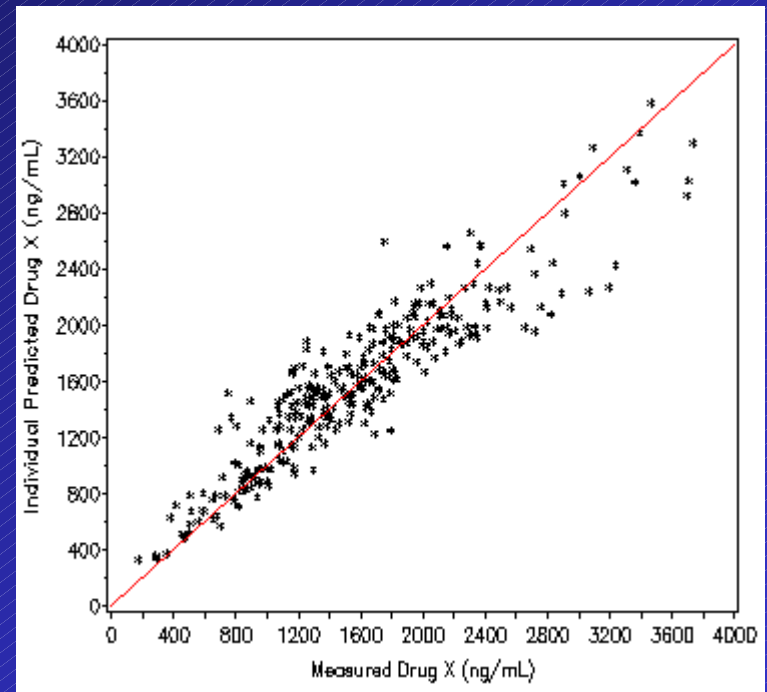
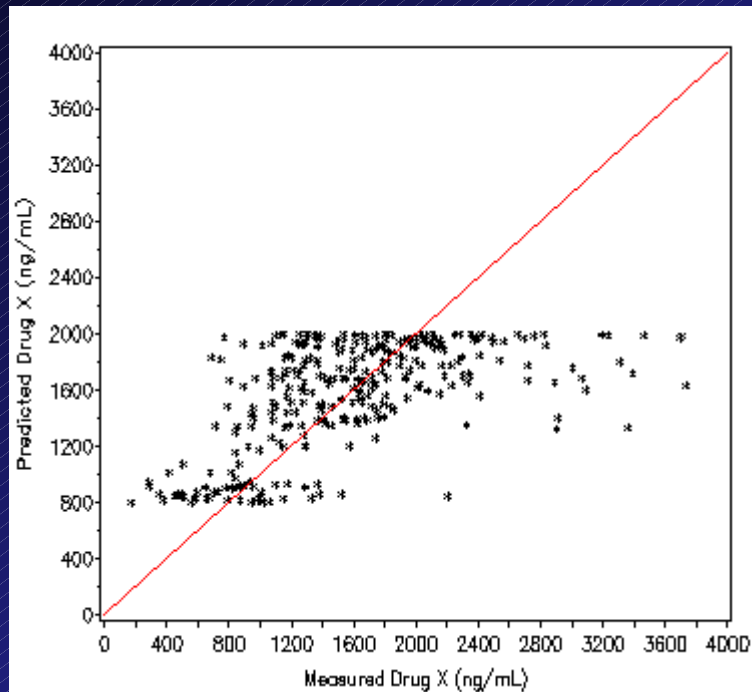
Objective Function Value (OBJF)

- Minimum value of the objective function (MVOF) is a global measure of goodness of fit and is proportional to $-2LL$ of the data
- $\Delta OBJF$ used in an approximate test for adding/deleting parameters
- Akaike Information Criterion (AIC): useful in non-hierarchical model comparisons = $\Delta OBJF + 2(\Delta p)$

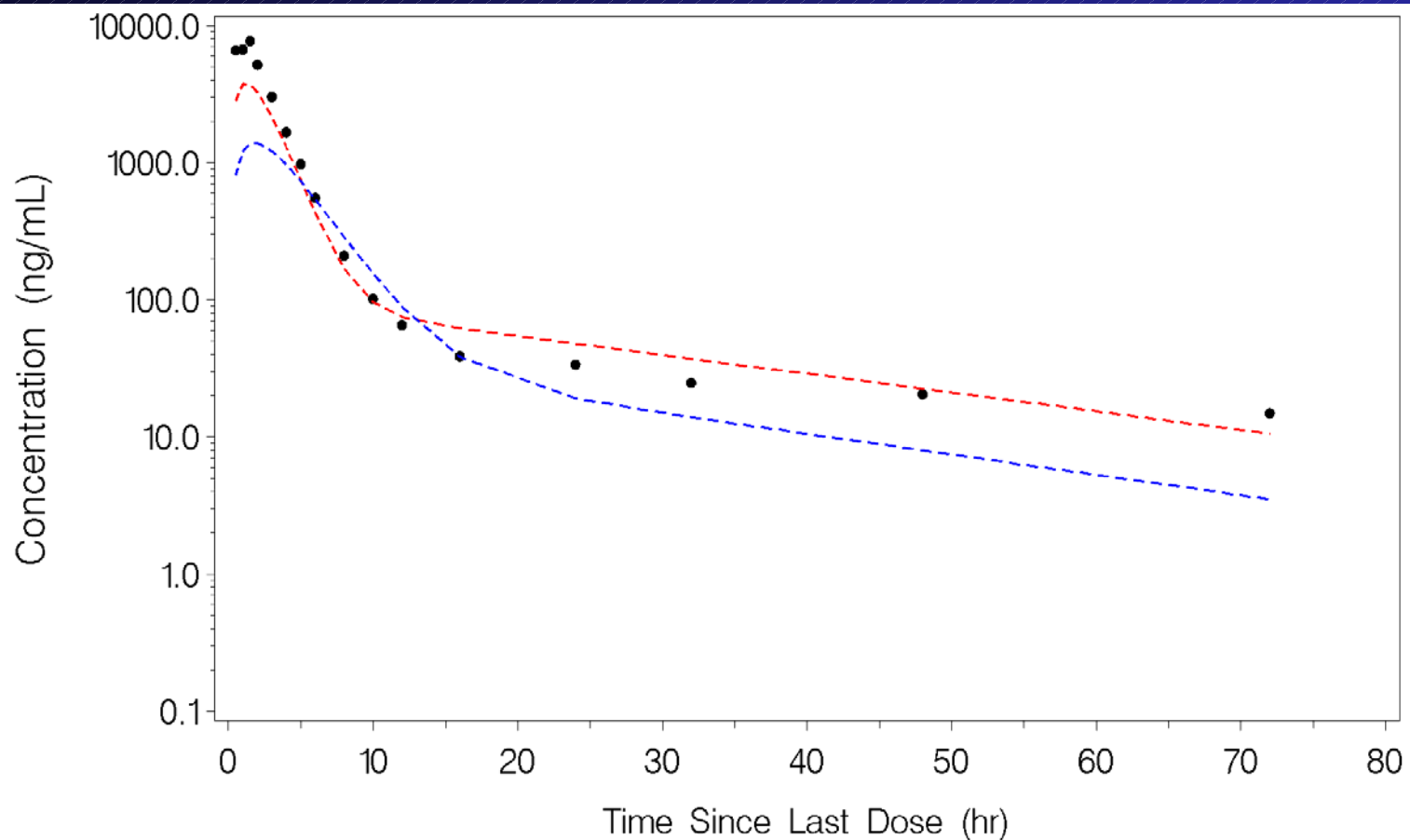
Typical Goodness of Fit Plots (GOF-Plots)

- DV vs PRED (Measured versus Predicted)
 - DV=Dependent Variable
 - PRED=Population mean Predicted Value
- RES vs PRED
 - Residuals
- WRES or CWRES vs PRED
 - WRES=Weighted Residual
 - CWRES=Conditional Weighted Residuals
- WRES or CWRES vs TSLD
 - TSLD=Time Since Last Dose
- IPRED vs DV
 - IPRED=Individual Predicted
 - (Takes into account η)

Base Population PK Model Goodness of Fit Plots (GOF-Plots)

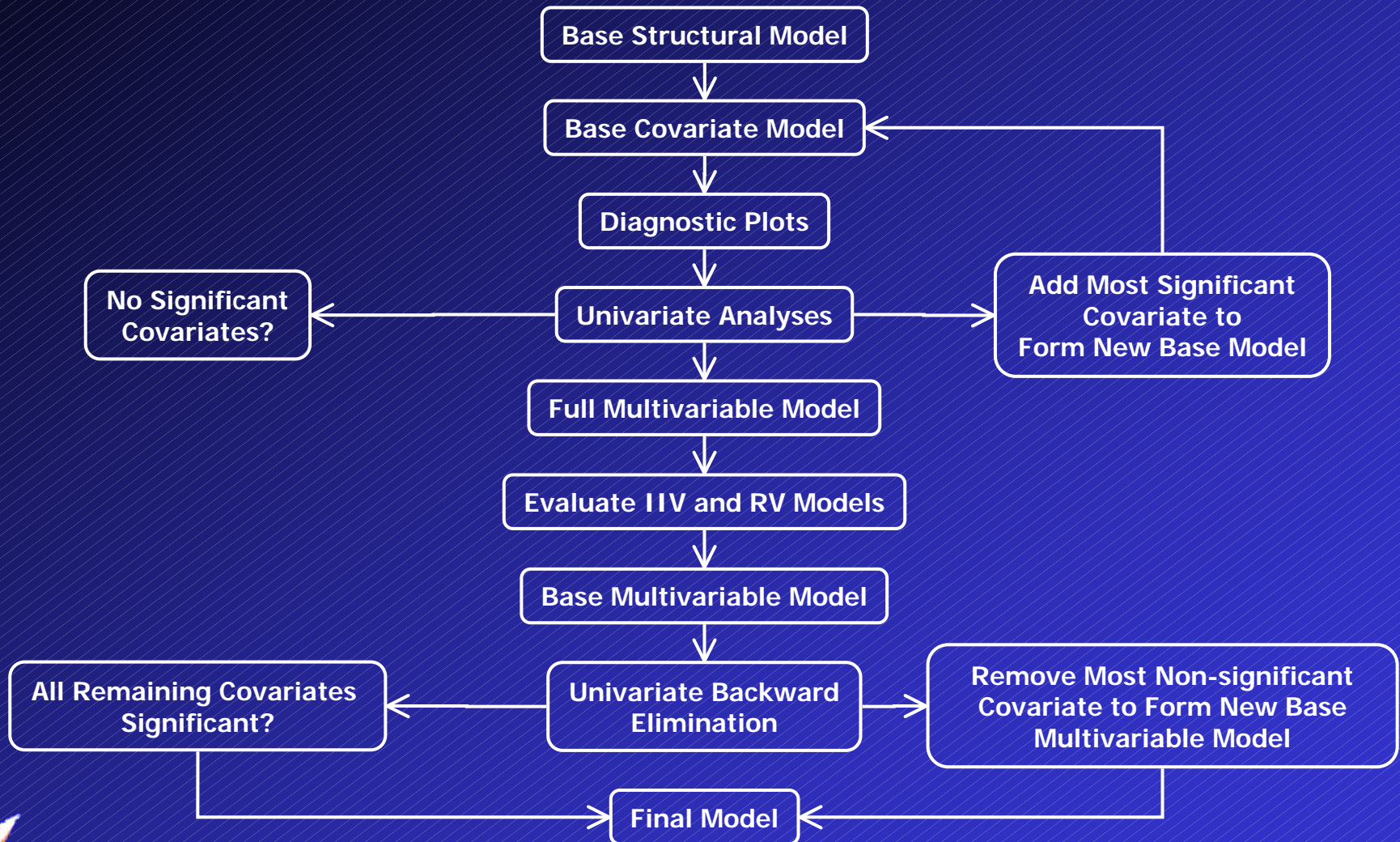


Base Population PK Model Goodness of Fit Plots (GOF-Plots)



The red line represents the individual predicted concentrations.
The blue line represents the population predicted concentrations.

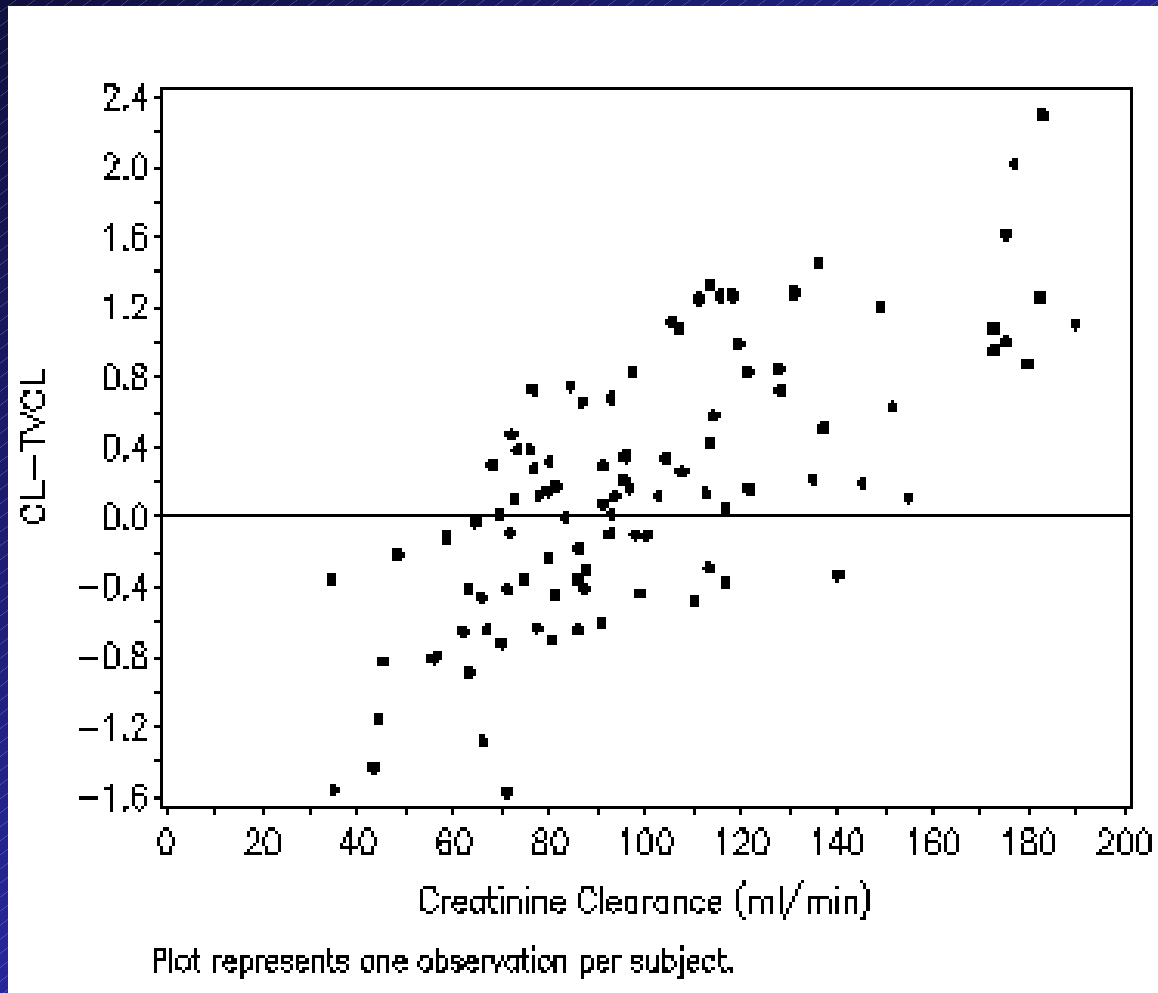
Model Development Process



Exploring Covariate Effects

- Many methods/procedures available, some automated
 - GAM, WAM
 - Stepwise in SAS®
- Choice of method may depend on objectives, goals, timeframe
- Depending on # of predictors, can be the most time-consuming phase of an analysis

Delta Plots

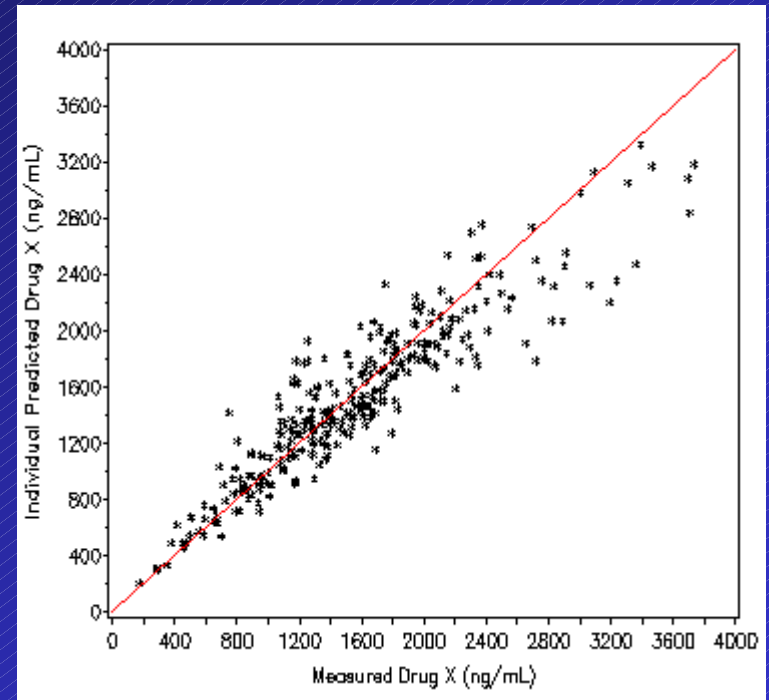
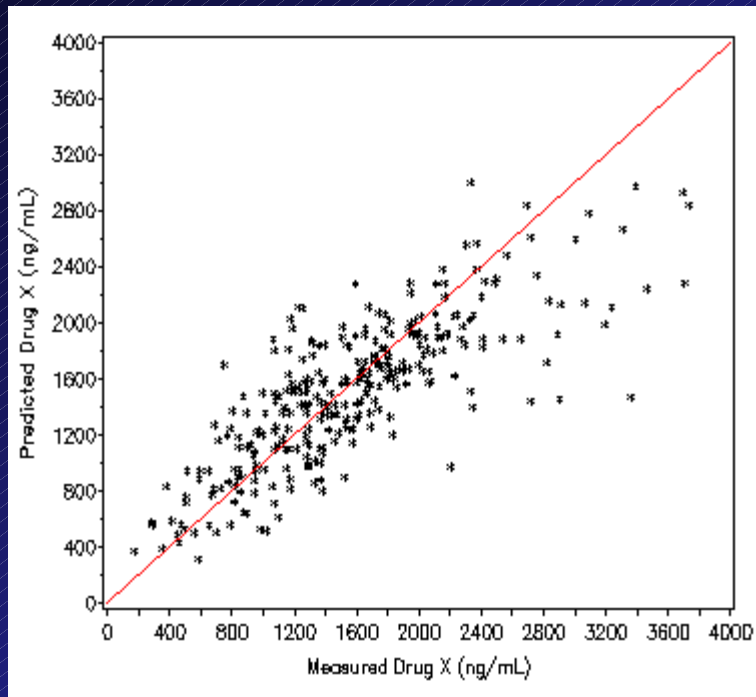


Final Population PK Model

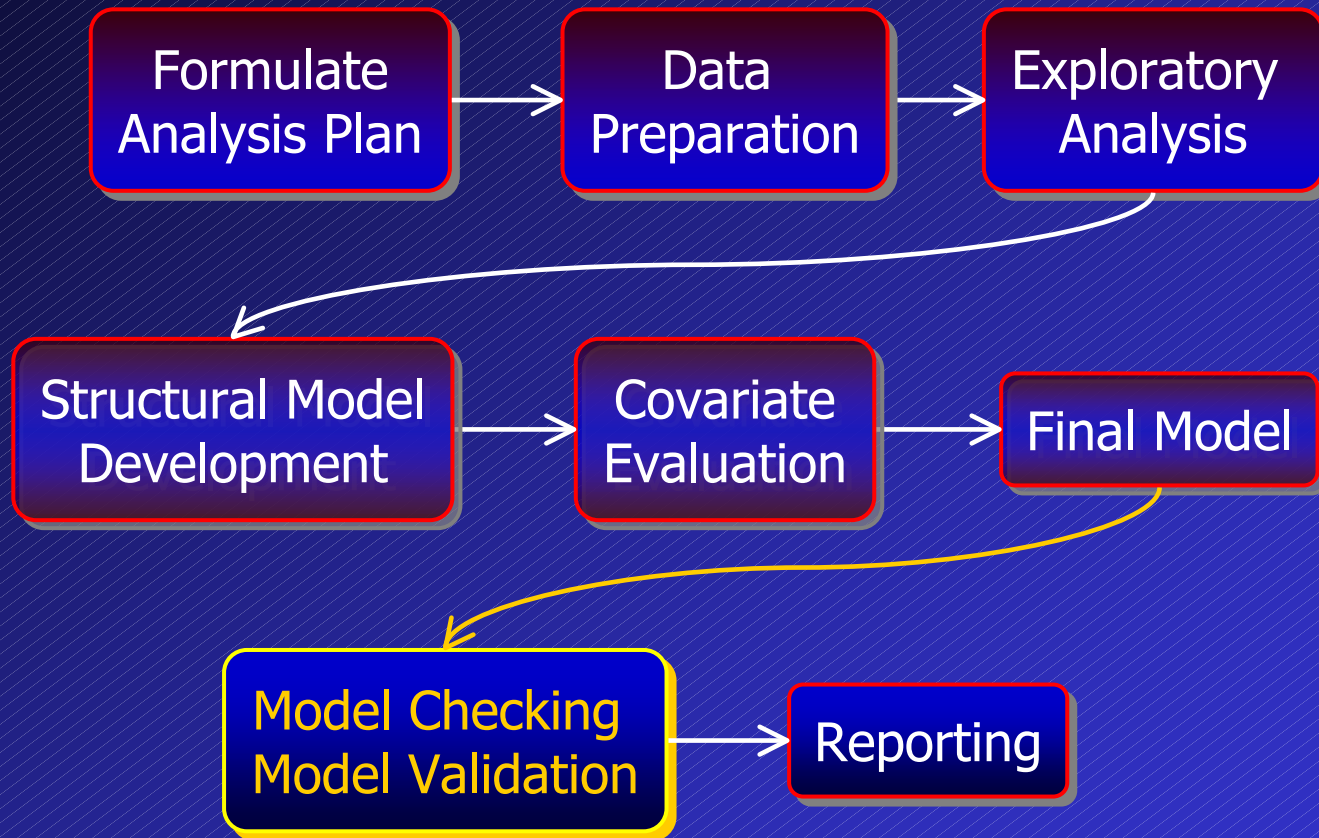
Parameter	Final Parameter Estimate	
	Population Mean	%SEM
Ka (1/hr)	0.491	11.8
CL(L/hr)	3.13	2.1
V(L)	53.4	7.2
CL (CrCL effect)	0.627	9.5
V (WTKG effect)	0.781	14.2
IIV CL (%CV)	17.55	22.2
IIV V (%CV)	18.87	77.0
RV (log SD)	0.19	11.6

MVOF = 3763.901

Final Model Diagnostics



Phases of a Population Analysis



Model Checking and Evaluation

- Model evaluation (previously known as model validation, see guidance) – many options
 - Internal vs. External
 - Bootstrap or cross-validation procedures
 - Posterior predictive check: simulate data from final model & check against raw data
 - Objective Function Mapping: determine if model is at a local or global minimum

References

- American College of Clinical Pharmacology. A Pharmacometrics Resource: Theory. May 23, 2005. Available at: <http://accp1.org/pharmacometrics/theory.htm>. Accessed October 18 2006.
- FDA. Guidance for Industry: Population Pharmacokinetics. February 1999. Available at: <http://www.fda.gov/cder/guidance/1852fnl.pdf>. Accessed October 18 2006.
- J. Fiedler-Kelly. A 3-day Introductory Workshop in Population PK Data Analysis: A Hands-on Course Using NONMEM®. Buffalo, NY. June 2006.
- LB Sheiner and SL Beal. A Short Course in Population Pharmacokinetics Data Analysis Using the NONMEM System. University of California, San Francisco, CA. May 1997.