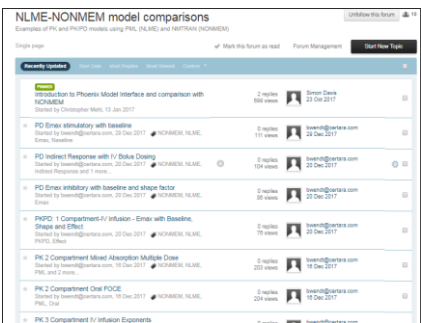


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Introduction to NONMEM - NLME Comparisons: PK 1 compartment IV Bolus model FOCE

**NONMEM-NLME Coparisons: On the Forum**

<https://support.certara.com/forums/forum/35-nlme-nonmem-model-comparisons/>



**NLME-NONMEM model comparisons**

Discussion Topic	Replies	Views	Last Post
Introduction to Phoenix Model Interface and comparison with NONMEM	2 replies	690 views	23 Dec 2017
PK 1-compartment IV bolus model FOCE	0 replies	111 views	20 Dec 2017
PK 2-compartment oral with Disease State covariate on V and CL	0 replies	194 views	20 Dec 2017
PK 2-compartment mixed absorption Multiple Dose	0 replies	203 views	19 Dec 2017
PK 2-compartment Oral FOCE	0 replies	204 views	19 Dec 2017
PK 3-compartment IV infusion Equipoints	0 replies	204 views	19 Dec 2017

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**New Series: NONMEM- NLME Comparisons**

- FEB 8**  
Introduction to NONMEM-NLME Comparisons  
PK 1-compartment IV bolus model FOCE  
February 8, 2018 | 10am EST  
Presenter: Bernd Wendt
- FEB 22**  
NONMEM-2-NLME  
PK 2-compartment multiple dose-IV bolus Plasma and Urine QRP(EM)MP  
February 22, 2018 | 10am EST  
Presenter: Bernd Wendt
- MAR 8**  
NONMEM-2-NLME  
PK 2-compartment oral with Disease State covariate on V and CL  
March 8, 2018 | 10am EST  
Presenter: Venkateswari Muthukrishnan
- MAR 22**  
TMDD Model Translated from NONMEM (NM-TRAN) to Phoenix NLME (PML)  
March 22, 2018 | 10am EST  
Presenter: Loan Pham, Camargo

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**Agenda**

- Model Description
- Exploratory Data Analysis
- Input Data
- Model in NM-TRAN and PML Code
- Demo
- Summary
- Q&A

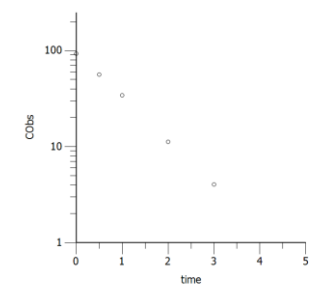
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**NONMEM to NLME: 1c\_iv\_bolus\_foce**

- Model Description
  - Structural Model
    - One compartment model, single IV Bolus of 100 units, first order kinetics
    - Parametrization: Cl and V
  - Error model
    - Multiplicative: 10% CV
  - Structural Parameters
    - V: Lognormal
    - Cl: Lognormal
  - Fixed Effects
    - tvV: 1
    - tvCl: 1
  - Random Effects
    - Variances all 0.1 (~30% population variability)
  - Engine
    - FOCE ELS

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**1c\_iv\_bolus\_foce: Plot First Patient**



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### 1c\_iv\_bolus\_foce: Input Data

NONMEM						NLME					
ID	TIME	DV	DOSE	AMT	WT	ID	TIME	CObs	DOSE	AMT	WT
1	0	0	0	0	100	1	0	43.1026	1	100	100
2	0	0.5	54.2902	0	100	2	0	54.2902	0	100	100
3	0	1	34.2469	0	100	3	0	34.2469	0	100	100
4	0	2	11.2468	0	100	4	0	11.2468	0	100	100
5	0	3	4.24657	0	100	5	0	4.24657	0	100	100
6	0	5	0.972991	0	100	6	0	0.972991	0	100	100
7	1	0	94.7954	1	100	7	1	94.7954	1	100	100
8	1	0.5	68.9021	0	100	8	1	68.9021	0	100	100
9	1	1	44.79	0	100	9	1	44.79	0	100	100
10	1	2	22.624	0	100	10	1	22.624	0	100	100
11	1	3	9.8758	0	100	11	1	9.8758	0	100	100
12	1	5	2.61519	0	100	12	1	2.61519	0	100	100
13	2	0	124.889	1	100	13	2	124.889	1	100	100
14	2	0.5	70.2667	0	100	14	2	70.2667	0	100	100
15	2	1	34.7922	0	100	15	2	34.7922	0	100	100

### NONMEM to NLME: 1c\_iv\_bolus\_foce

NONMEM	NLME
<code>Model Selection</code>	<code># Model Selection</code>
<code>\$SUB ADVAN1 TRANS2</code>	<code>cfmicro(A1, CI / V)</code>
<code>;Structural Parameters</code>	<code>dosepoint(A1)</code>
<code>\$PK</code>	<code>#Structural Parameters</code>
<code>TVV = THETA(1)</code>	<code>stparm(V = tvV * exp(nV))</code>
<code>V = TVV * EXP(ETA(1))</code>	
<code>TVCL = THETA(2)</code>	<code>stparm(CI = tvCl * exp(nCI))</code>
<code>CI = TVCL * EXP(ETA(2))</code>	

### Using the PML: 1c\_iv\_bolus\_foce

NONMEM	NLME
<code>; scale parameter</code>	<code>; scale parameter: Must be defined explicitly</code>
<code>S1 = V</code>	<code>C = A1 / V</code>
<code>=====setup initial values for THETA and OMEGA ===</code>	<code># =====setup initial values for THETA and OMEGA ===</code>
<code>\$THETA</code>	<code>fixeff(tvV = c(, 1, ))</code>
<code>(0, 1, );TVV</code>	<code>fixeff(tvCl = c(, 1, ))</code>
<code>(0, 1, );TVCL</code>	
<code>\$OMEGA</code>	<code>ranef(diag(nV, nCI) = c(0.1, 0.1))</code>
<code>0.1</code>	
<code>0.1</code>	

### Using the PML: 1c\_iv\_bolus\_foce

NONMEM	NLME
<code>; specify residual error model</code>	<code># specify residual error model</code>
<code>; setup initial estimates for the variance matrix (SIGMA) of the residual variability</code>	<code># setup initial estimates for the variance matrix (SIGMA) of the residual variability</code>
<code>\$SIGMA 0.01</code>	<code>error(CEps = 0.1)</code>
<code>\$ERROR</code>	<code>observe(CObs = C * (1 + CEps))</code>
<code>Y = F*(1 + EPS(1))</code>	
<code>; additional output</code>	
<code>IVAR = TIME</code>	
<code>IPRED = F</code>	
<code>IRES = DV - IPRED</code>	

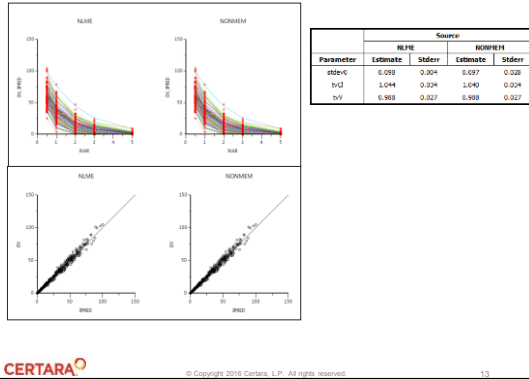
### Using the PML: 1c\_iv\_bolus\_foce

```

test() {
  cfmicro(A1, CI / V)
  dosepoint(A1)
  C = A1 / V
  error(CEps = 0.1)
  observe(CObs = C * (1 + CEps))
  stparm(V = tvV * exp(nV))
  stparm(CI = tvCl * exp(nCI))
  fixeff(tvV = c(, 1, ))
  fixeff(tvCl = c(, 1, ))
  ranef(diag(nV, nCI) = c(0.1, 0.1))
}
  
```

### Demo

## 1c\_iv\_bolus\_foce: Results



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## 1c\_iv\_bolus\_foce: Summary

- Setup and Run a NONMEM Model:
  - 1-compartment IV Bolus
- Setup and run the equivalent Model in Phoenix NLME
- Postprocessed the NONMEM results and compared them with the NLME results
  - Only small differences

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Questions?



## Certara University

- A wide range of On Demand and Classroom courses are available through Certara University:
  - Introductory, intermediate and advanced instruction in Phoenix WinNonlin, **Population Modeling using NLME**, IVIVC Toolkit
  - Fundamentals of Pharmacokinetics and Pharmacodynamics
  - Noncompartmental data analysis
  - Programming Bootcamp
  - Partner Lectures and Webinar series
- Please visit our [Certara University](http://www.certarauniversity.com) web site for more information

[www.certarauniversity.com](http://www.certarauniversity.com)

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## Coming up...



**NONMEM-2-NLME**  
 PK 2-compartment multiple dose-IV bolus Plasma and Urine QRP/IMP  
 February 22, 2018 | 10am EST  
 Presenter: Bernd Wendt

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