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Evaluation of the nonparametric estimation method in NONMEM VI β

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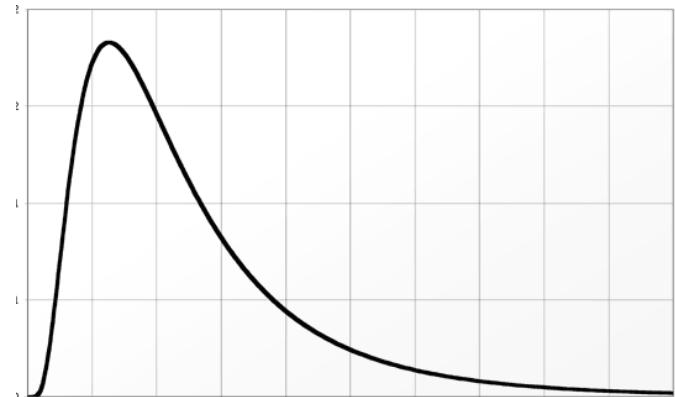
Sweden



Statistical methods for population analysis

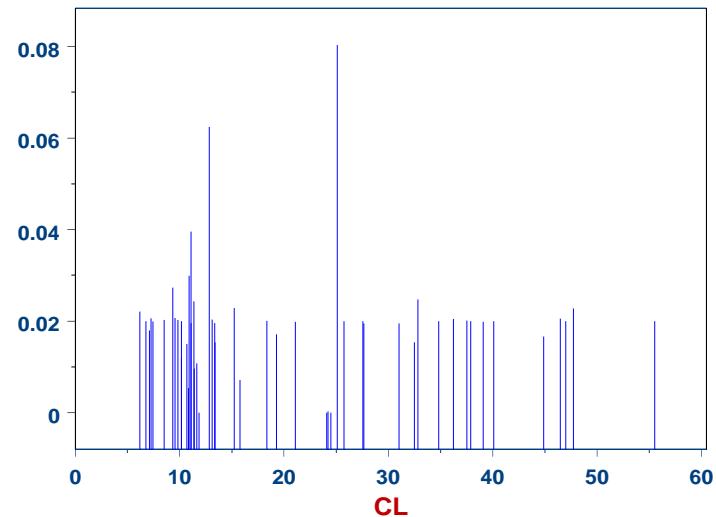
✓ Parametric methods

- assumed shape of parameter distribution
- **normal** or transformation of normal (usually)
- **mean and variance**
- imprecision estimates (SE)



✓ Nonparametric methods

- no assumptions regarding the distribution shape
- **discrete, non-continuous**
- collection of **finite number of values** for all parameters along with **the estimated probabilities**
- no imprecision estimates





Search for nonparametric distribution

How NONMEM VIβ does it?

1. *Search for support points*

- ✓ ***NONMEM runs parametric step (FO/FOCE)***
- ✓ ***Computes the Empirical Bayes Estimates (EBEs)***
- ✓ ***The EBES are taken as the support points for the nonparametric distribution***

2. *Probability estimation*

NONMEM recomputes:

- ✓ the joint probability
- ✓ the marginal cumulative probability for each parameter distribution

NONMEM output

NONPARAMETRIC ESTIMATE **

MINIMUM VALUE OF OBJECTIVE FUNCTION **

-1010.387



NPOFV – as OFV for mixture model

$n_{\text{mixtures}} = n_{\text{subjects}}$

EXPECTED VALUE OF ETA *****

ETA1

ETA2

-3.66E-01 -6.15E-03



New set of

typical parameter estimates

COVARIANCE MATRIX OF ETA *****

ETA1

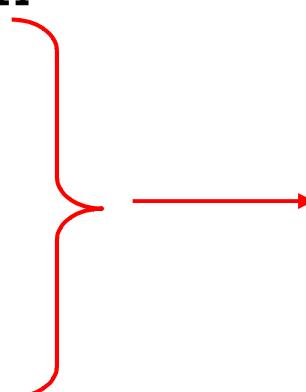
ETA2

ETA1

+ 3.40E-01

ETA2

+ -5.53E-02 2.85E-01



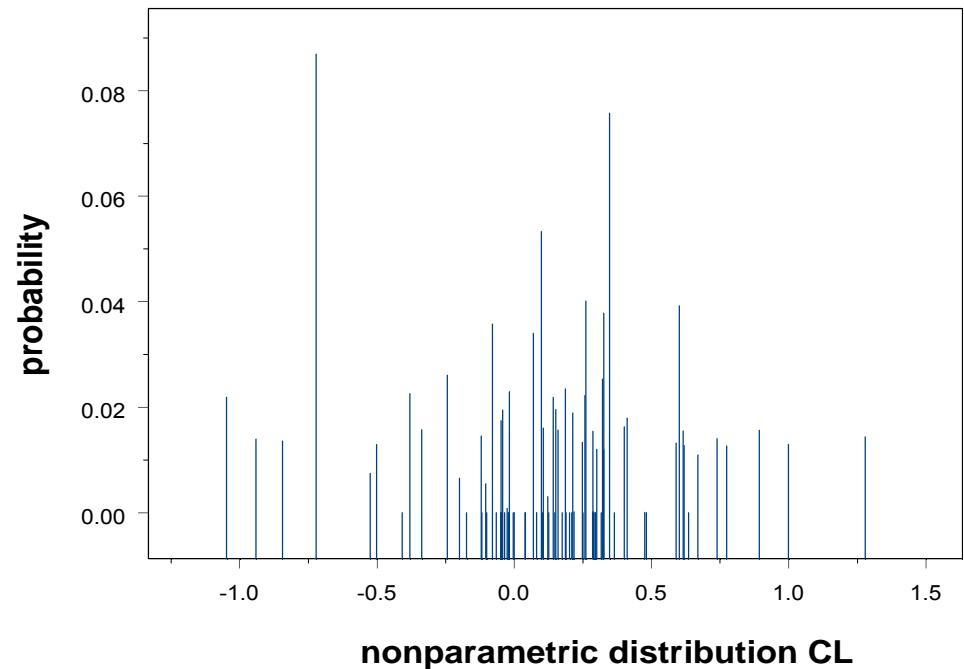
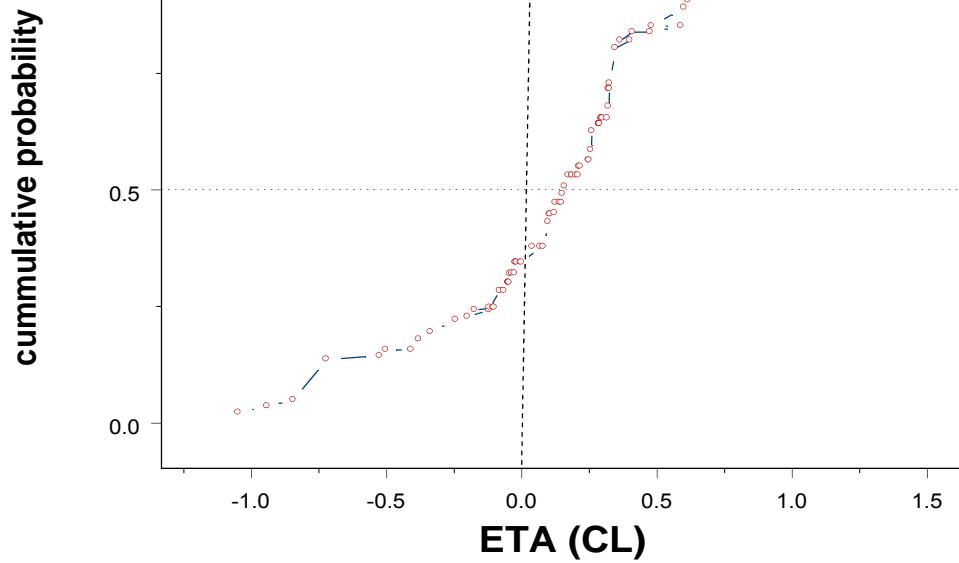
Nonparametric

Ω matrix



NONMEM output – nonparametric distribution

- ✓ **Cumulative marginal density function** for each parameter distribution
- ✓ **Joint probability** for vector of parameters





Aim

Aim

*to explore the performance of the
nonparametric estimation method in NONMEM VI β*

Method

a Monte Carlo simulation study

Emphasis

- ✓ *the analysis of data with
non-normal distribution of random effects*

- ✓ *the analysis of data with
misspecification in residual error magnitude*



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*the analysis of data with
non-normal distribution of random effects*



Simulation settings

Simulated PK DATA

- ✓ **One compartment iv bolus model**
- ✓ **3 observations** per individual (early / mid / late sample)
- ✓ **100 datasets** for each condition
- ✓ **Random effects distribution** - 4 different types
 - Log-normal
 - Quadromodal
 - Trimodal
 - Heavy-tailed

} *will discuss today*

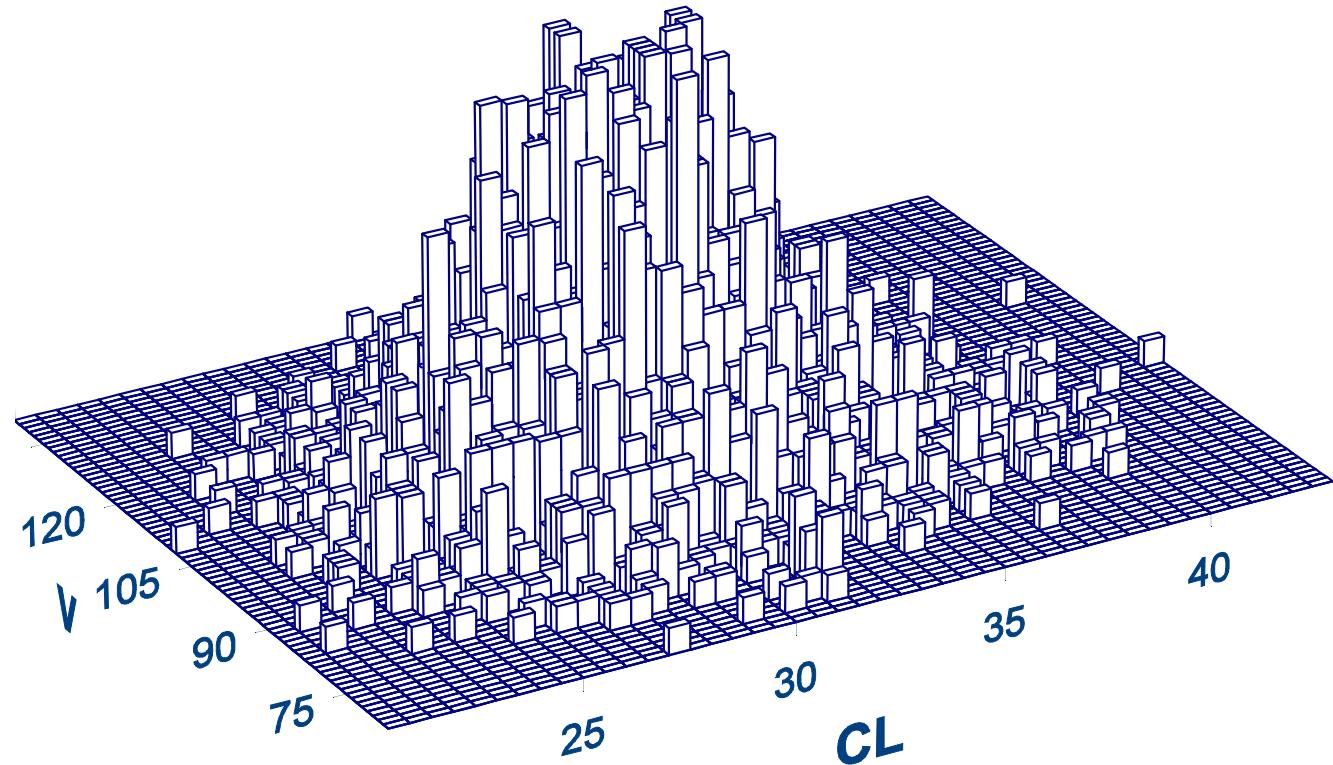
} *will be available at the web*



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Random effects distributions

1. Log-normal

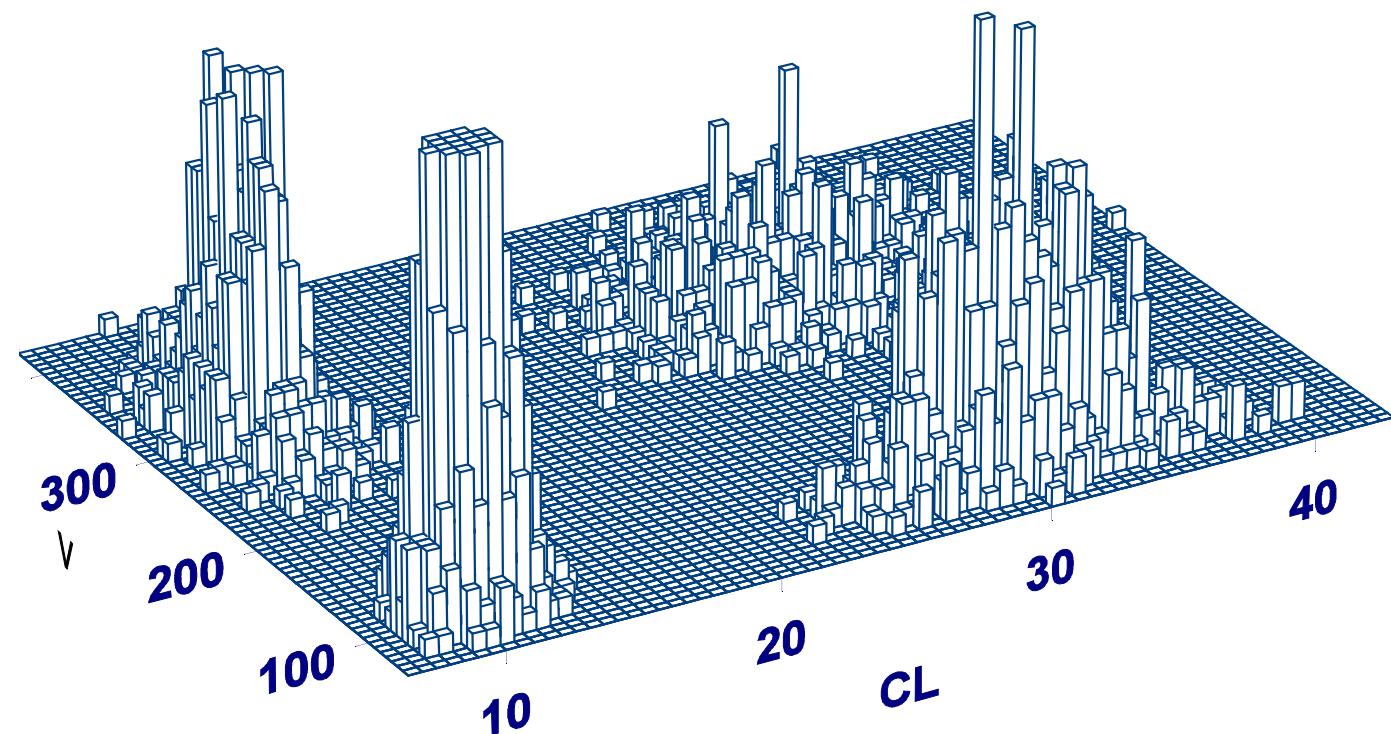




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Random effects distributions

2. "quadromodal" distribution

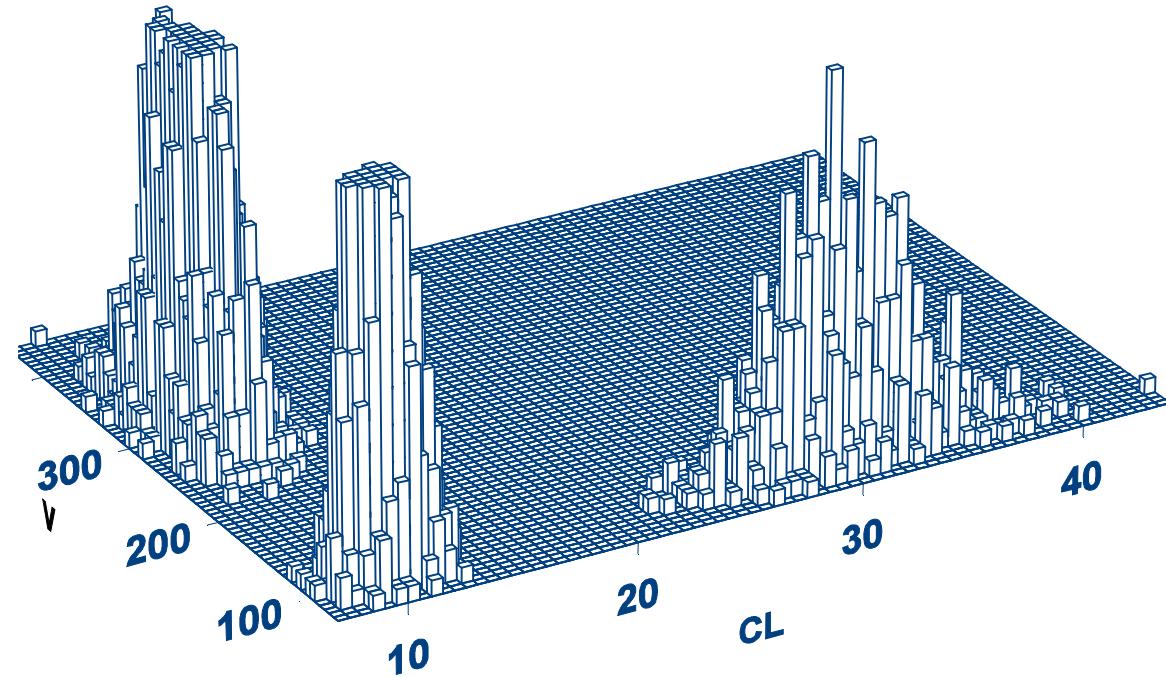




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Random effects distributions

3. "trimodal" distribution

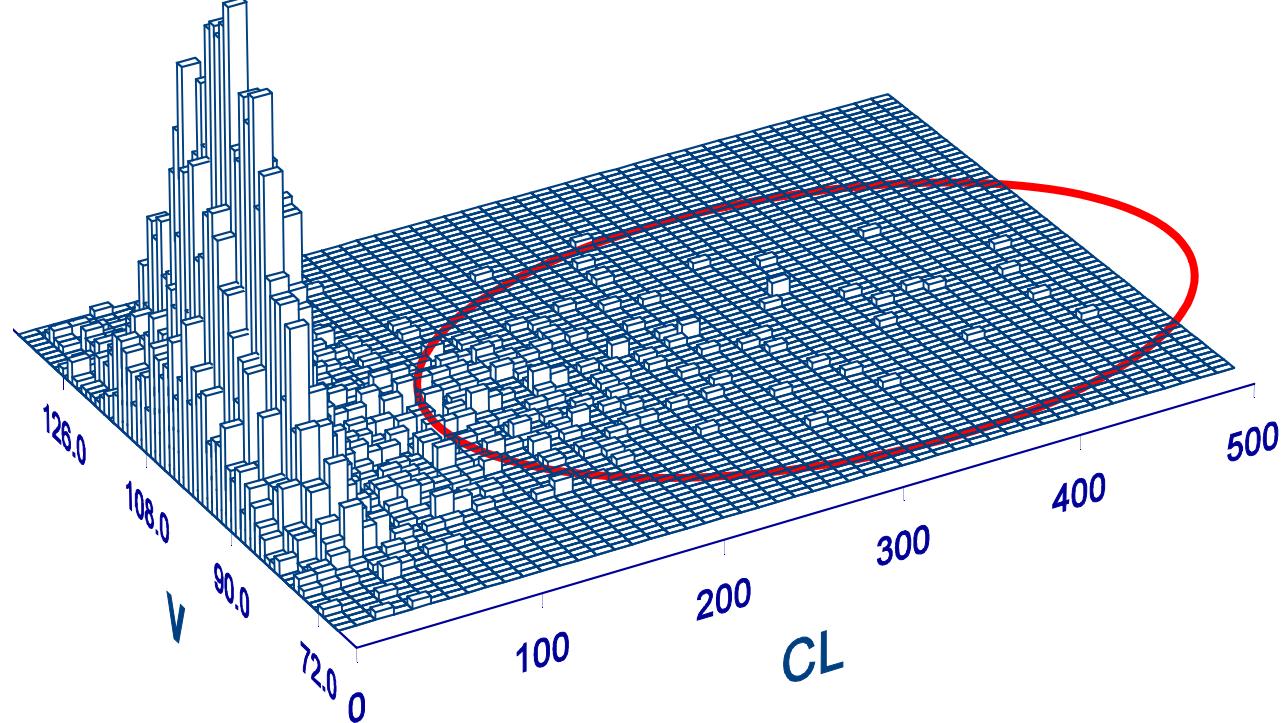




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Random effects distributions

4. Heavy-tailed distribution





Models fitted to the data

Parameter distribution

1. **PARAMETRIC** The Log-normal

2. **MIXTURE (\$MIX)** The TRUE one
(quadromodal)

3. **NONPARAMETRIC** The nonparametric
(EBEs – points of support)



based on the incorrectly assumed Log-normal distribution



Results evaluation

Estimated parameter distributions were

1. **evaluated** at the 10_{th}, 25_{th}, 50_{th}, 75_{th}, 90_{th} percentile
2. **compared to** the **true** distribution at the same percentiles

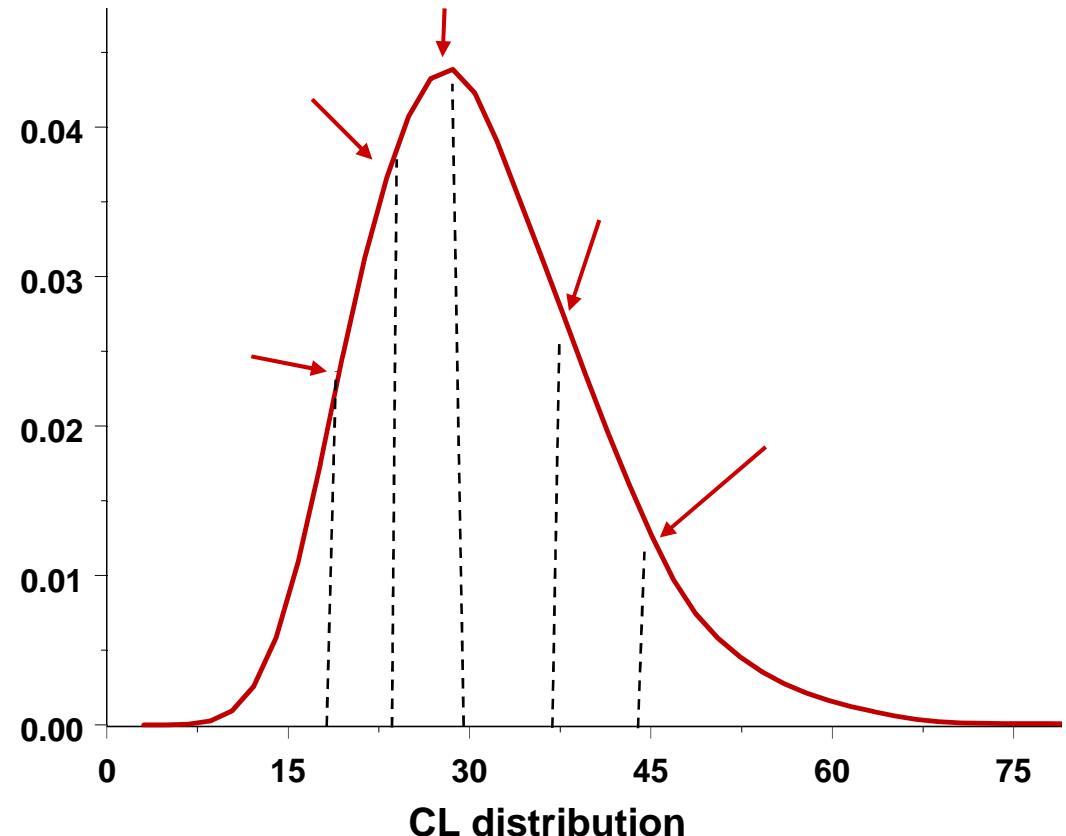
Relative estimation error (REE)

$$REE = \frac{P_{est} - P_{true}}{P_{true}}$$

P_{est} – estimated parameter value

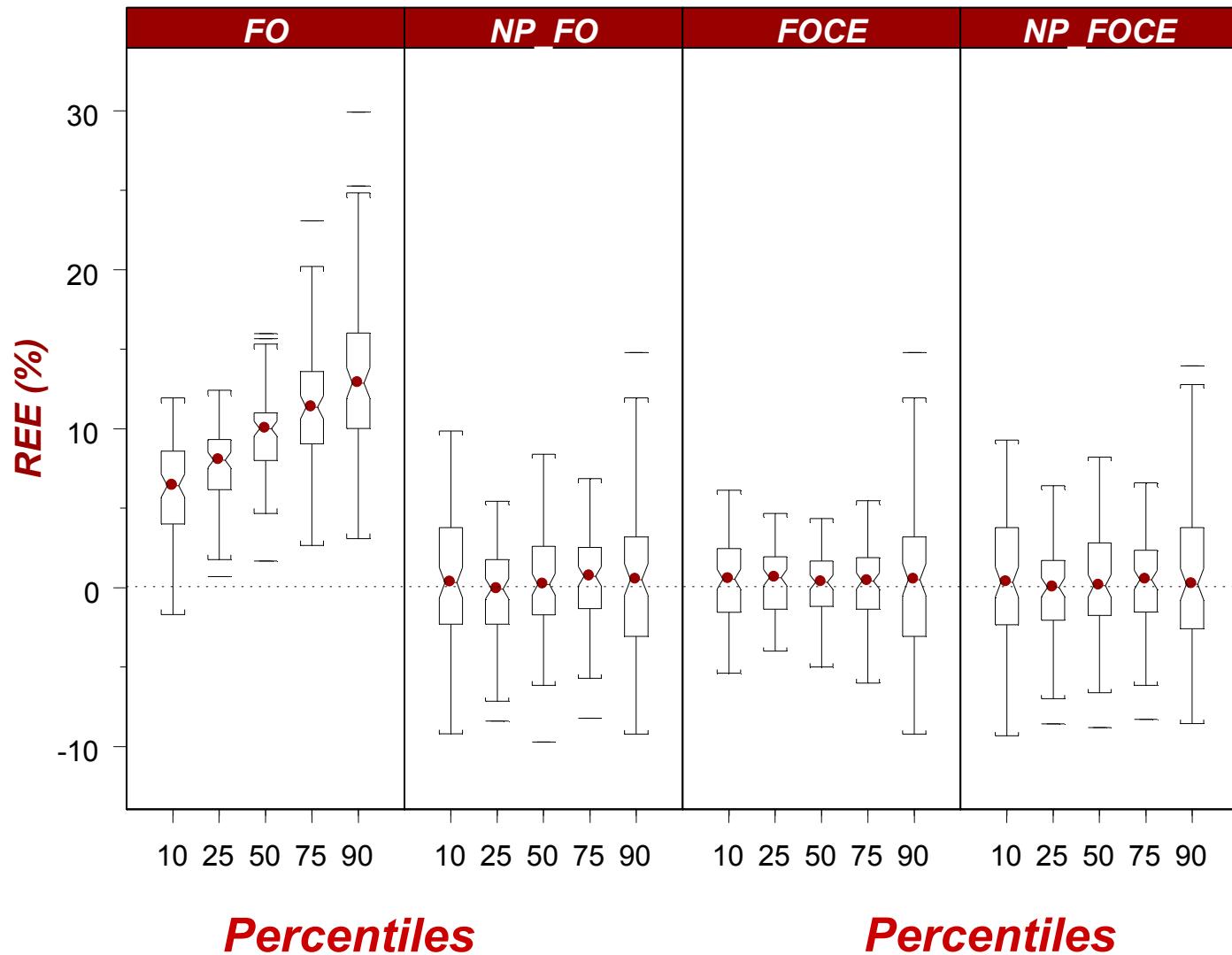
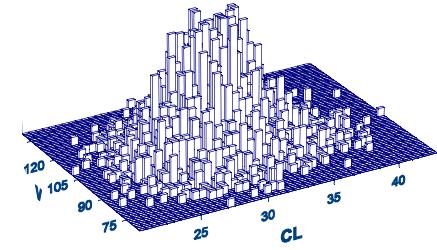
P_{true} – true parameter value

(at the given percentiles)





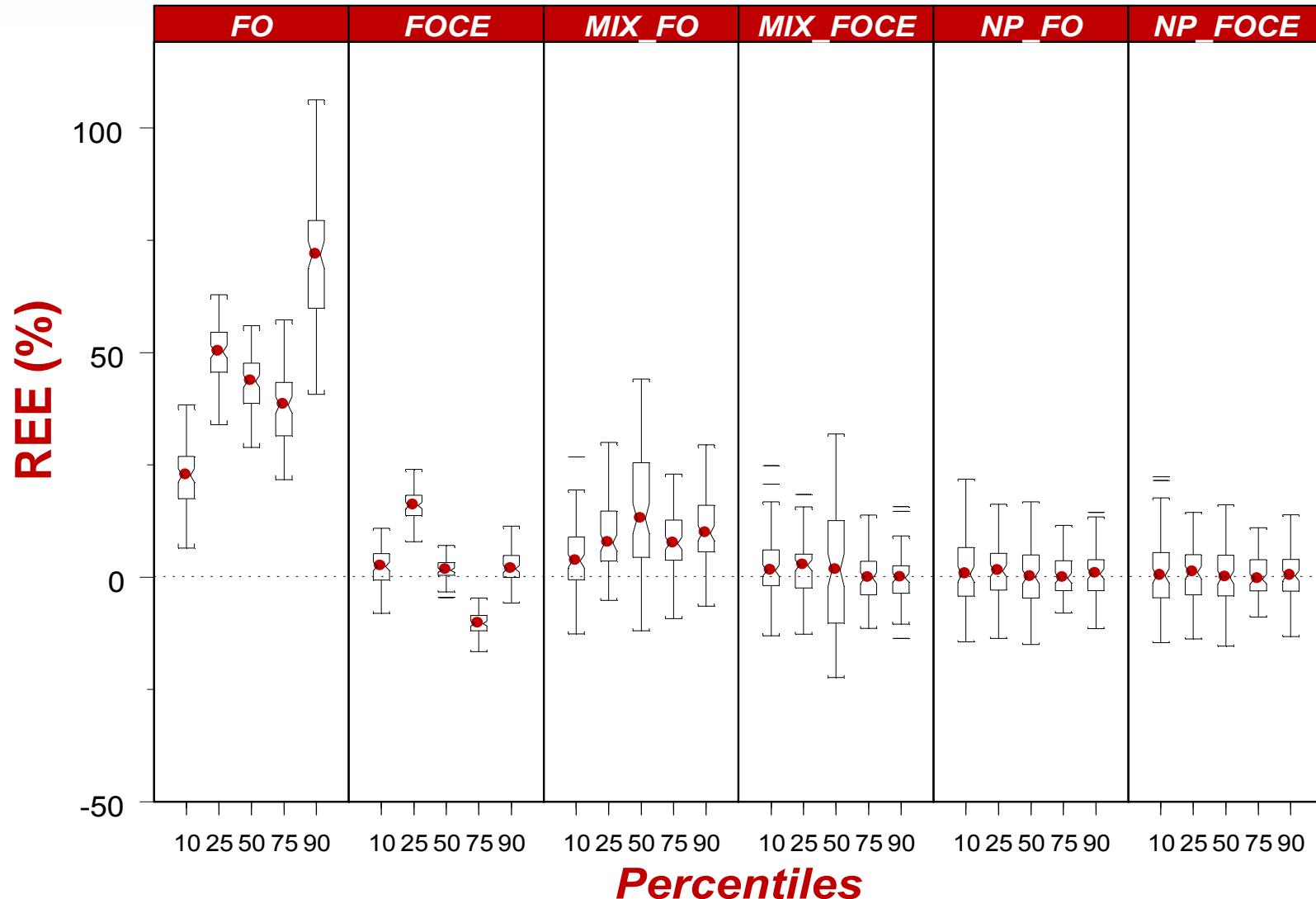
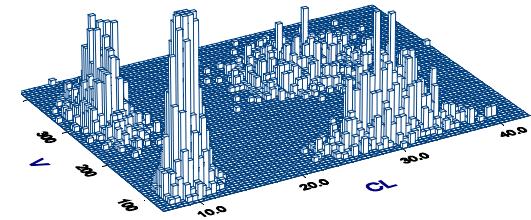
Results: Log-normal -CL





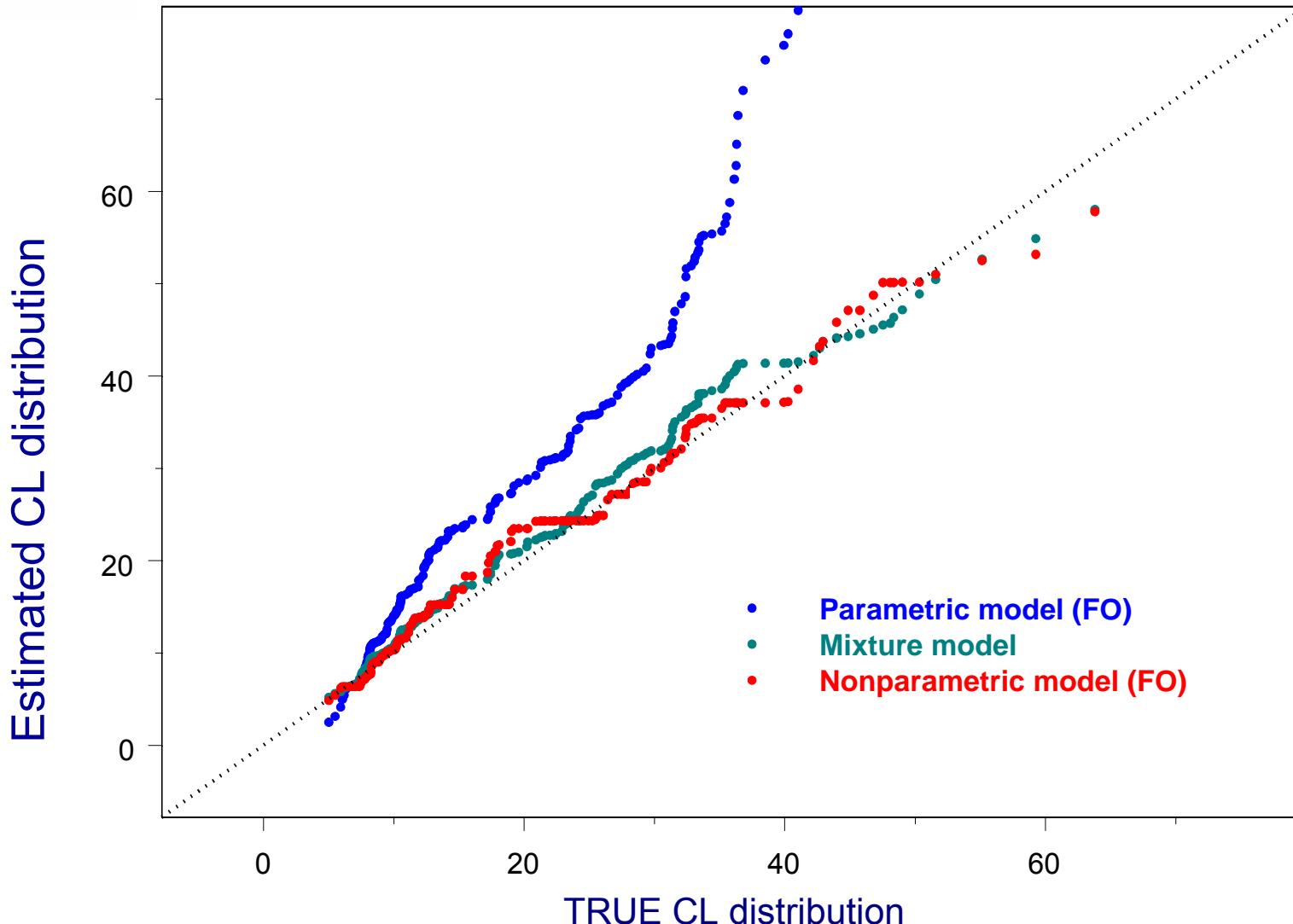
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Results: quadromodal - CL



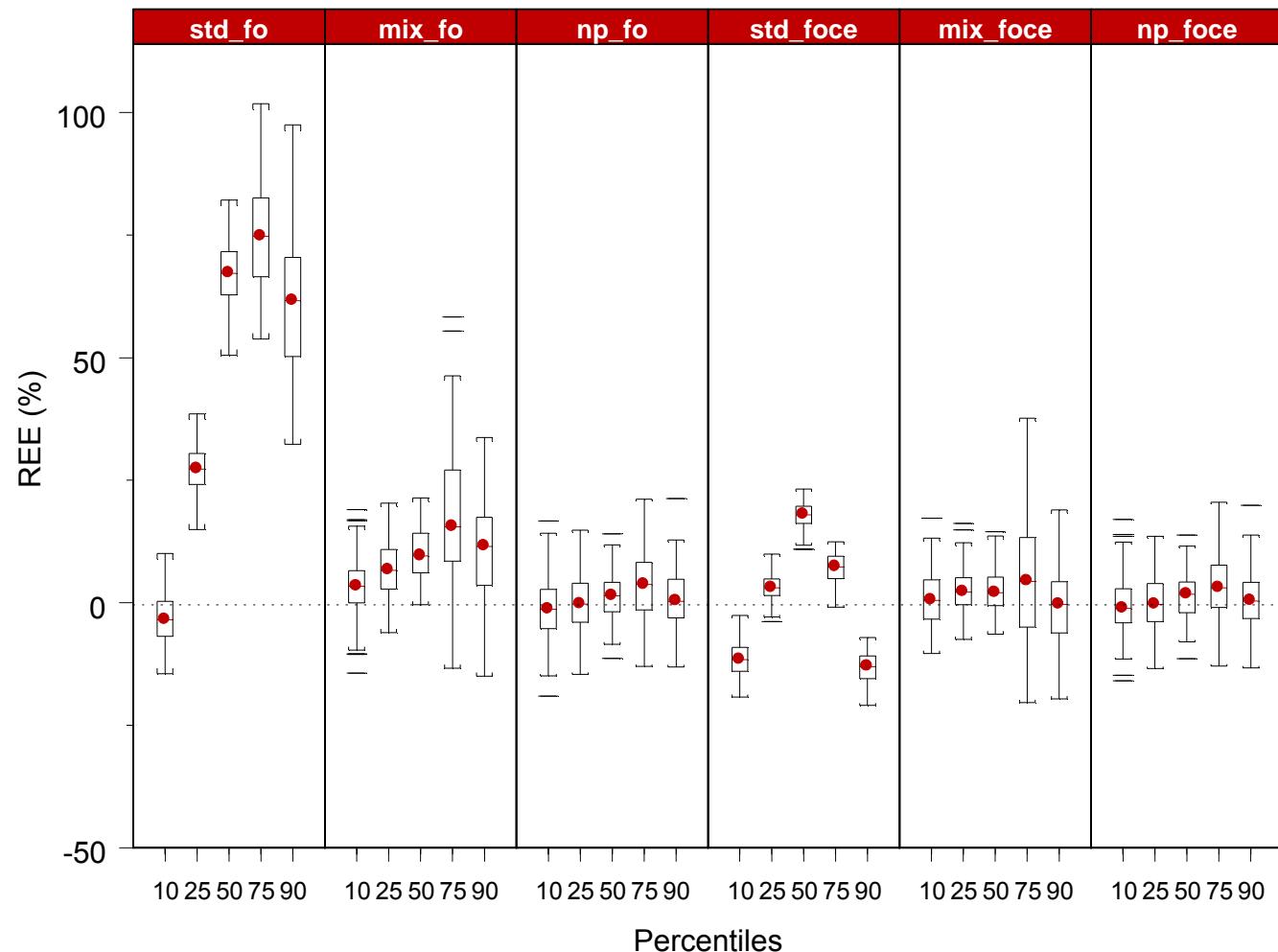


quadromodal: QQ plots - CL



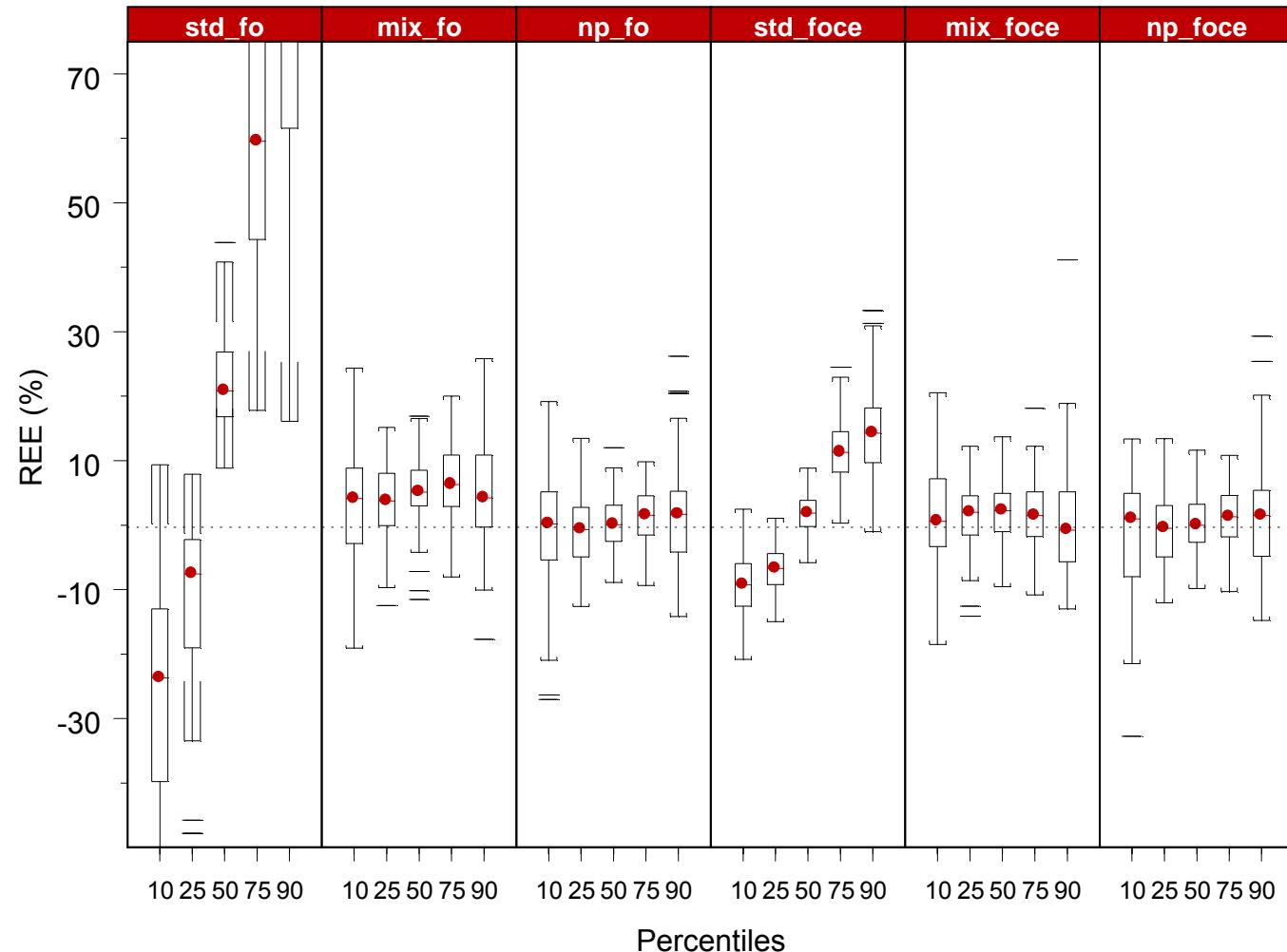


Results: trimodal - CL





Results: heavy-tailed - CL





Results cont.

- ✓ With both, FO and FOCE, the nonparametric parameter distribution **matched the TRUE parameter distribution** in *all studied cases*
- ✓ With less informative data and/or smaller population, relative bias in parameter estimates is increasing for all methods, but **the same trend remains**



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*the analysis of data with
misspecification in residual error magnitude*



Background

Residual (*intraindividual*) variability (RV):

- Assay error
- Model misspecification
- Individual biology
- Study conduct

Nonparametric approach has to specify the RV magnitude a priori

The ways to handle this :

The past : RV was fixed to the **assay error**

The present : RV is estimated with the parametric program and fixed to that

Option in NONMEM (the future way ?):

- ✓ possibility of estimating RV magnitude distribution (ETA on EPSILON)



With NONMEM, the **nonparametric distribution of RV** can be estimated



Aim

Aim

*to explore the sensitivity of the
nonparametric estimation method in NONMEM VI β
to the misspecification in residual error magnitude*

Questions to answer:

- ✓ *Is it sensitive?*
- ✓ *Can we benefit from the nonparametric distribution in residual error magnitude?*



In addition to previous simulation settings:
(with underlying log-normal parameter distribution)

2 mixtures in residual error magnitude:

1. 80 % of population – 30 % residual variability (CV%)
2. 20 % of population - 100% residual variability (CV%)



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Analysis

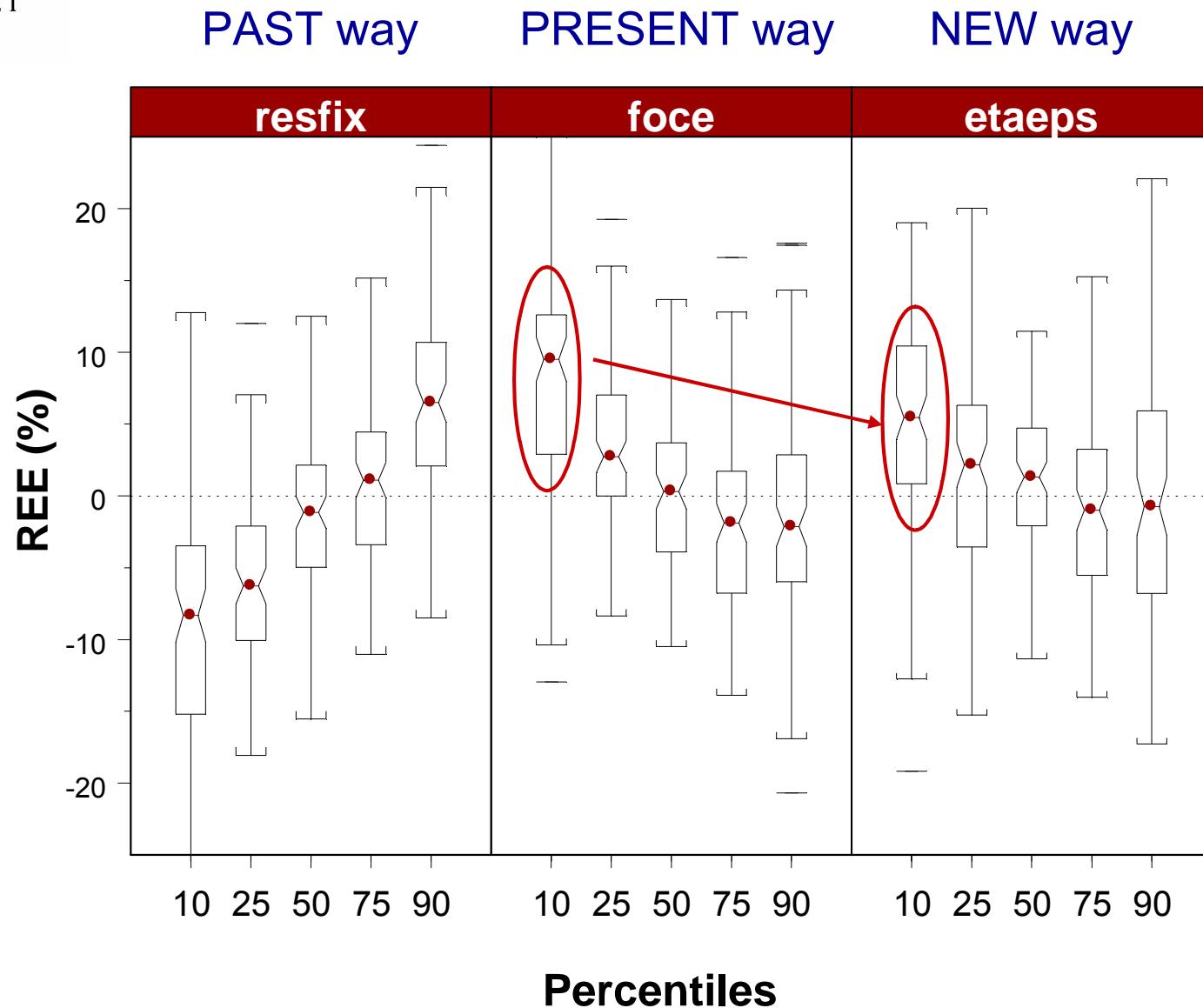
Models fitted to the data

Mimic:

1. RV **fixed apriori to the 30%** The past way
2. RV **estimated** The present way
3. **Nonparametric distribution** in RV estimated Advanced
NONMEM way



Results – CL distribution





Conclusions

- ✓ NONPARAMETRIC estimation method in NONMEM VI β is **sensitive** to the misspecification in the residual error magnitude

- ✓ We **can benefit** from nonparametric distributions of random effects in residual error magnitude (eta on epsilon)



Potential use of \$NONP

- ✓ ***an aid*** in building of parametric models, especially when the EBE distribution may be misleading or non-informative

- ✓ ***replacement of the final parametric model,*** especially when the FO method is used



What's going on in Uppsala more ...

- Evaluation of this method with ***real data sets***
- *Development of a bootstrap method for the **imprecision measurement** of the nonparametric distributions estimates*



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Acknowledgments

Professor Stuart Beal