# N-dimensional Likelihood Profiling (NLP) An Efficient Alternative to Bootstrap

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- Motivation: Bootstrapping is hard!
- Solution: N-dimensional Likelihood Profiling
  - Faster solution than bootstrapping with easy convergence
- Examples of application
- Summary and conclusion





# A day in the life of a quantitative clinical pharmacologist...

■ A semi-mechanistic, highly nonlinear PK/PD model to assess drug effects.

■Need to discuss simulation with uncertainty with the study team... And we need it soon...

But, bootstrapping takes forever

+How can I simulate with uncertainty?







# How can I simulate with uncertainty?

# **Bootstrap Covariance LLP Matrix**

Gold standard	Quick (if possible)	Reliable one parameter CI
Completes slowly	Sometimes not available	Doesn't work with >1 parameter.
Doesn't always complete	Often not a good estimate of the CI	Cannot simulate
May not have enough subjects	Assumption heavy	





Photo: Jim Bowen



## **NLP: Filling the Gap**

■The Idea:

Extend LLP to higher dimensionality allowing to sample the empirical distribution for simulation.

Faster solution than bootstrapping with easy

convergence.

■The Problem: Not so simple!

◆Requires integration of the likelihood surface in N dimensions.



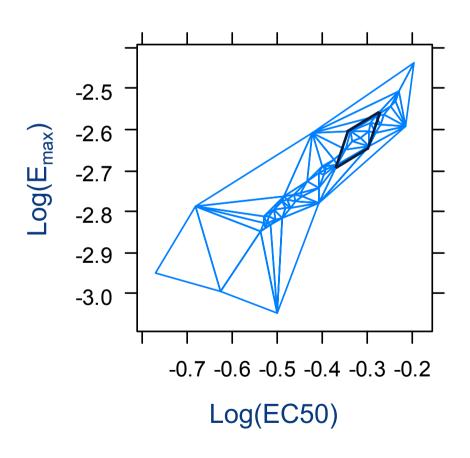
Photo: Chris Lott





### **How does NLP work?**

- From the initial estimates  $(\theta_0)$
- Relative likelihood values around  $\theta_0$  are searched maximizing information until convergence.

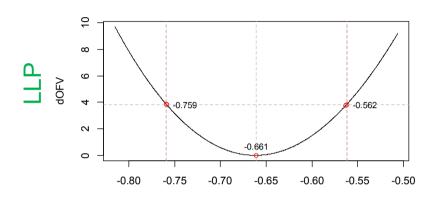


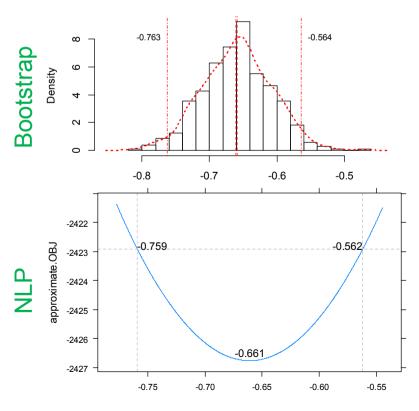


## **NLP** in one dimension

- Compare LLP, Bootstrap, and NLP with one parameter of a two compartment PK model.
- ■All provide equal results.
- ■Model evaluations:
  - **LLP 5**
  - ◆Bootstrap 1000
  - •NLP 8

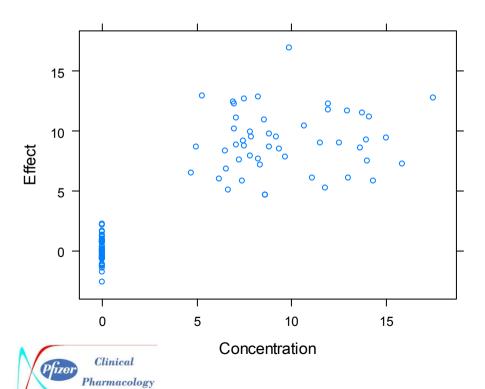




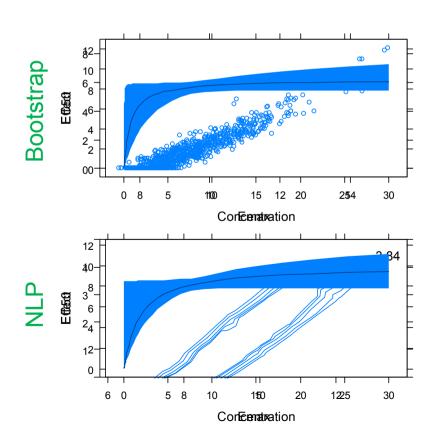


## **NLP** in multiple dimensions

- In a "no regrets dose" study...
- ■What can we infer about Emax/EC50?

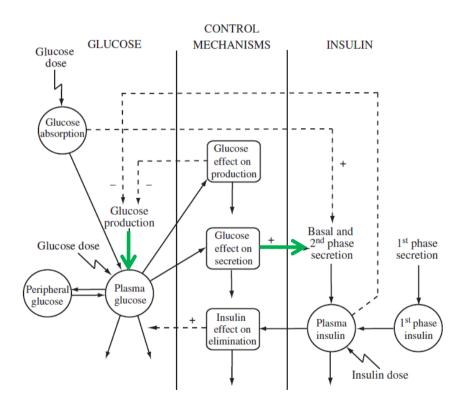


LLP





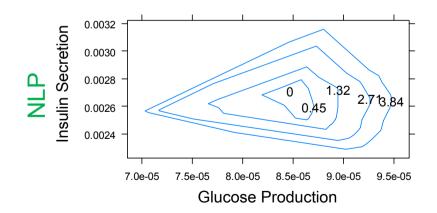
# A day in the life of a quantitative clinical pharmacologist... made simpler.



Silber, H. E., Jauslin, P. M., Frey, N. and Karlsson, M. O. Basic & Clinical Pharmacology & Toxicology, 106: 189–194.

Bootstrap did not complete after 10 days on a computational cluster.

NLP converged in 1 day on my laptop.





### **Conclusions**

- ■NLP allows estimation and sampling of the likelihood surface in multiple dimensions (≤5).
- Faster solution than bootstrap with easy convergence for
  - Long parameter estimation times,
  - Small populations,
  - Fixed parameters,
  - High bootstrap non-convergence rates.





# **Acknowledgements**

- ■Gianluca Nucci
- ■Wei Gao
- ■Ted Rieger
- ■Avi Ghosh





## **THANK YOU!**



Photo: Rose Robinson



# **BACKUPS**





### ■Covariance matrix

 Assumes that local curvature at the minimum defines full distribution.

### **LLP**

◆Find the point where OBJ increases by X (3.84 = 95% CI) through iteration.

### **■NLP**

- Find the (piecewise, continuous) equation that defines the OBJ surface as a function of model parameters.
- Can be used like LLP for one dimension, but allows sampling.
- Provides multivariate empirical distribution.

## ■Bootstrap

Find parameter estimates through population resampling.





# **Summary**

	LLP	Bootstrap	NLP
Can Sample for Simulation	No	Yes	Yes
CPU Time	~1x N	~1000-2000x N	~5-100x N
Dimensions	1	Full	Intermediate
Uncertainty in FIXED Parameters	Yes	No	Yes



Green indicates two best for row.

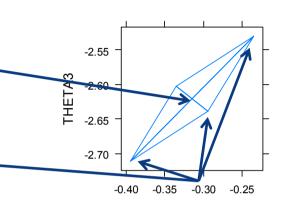
### **How does NLP work?**

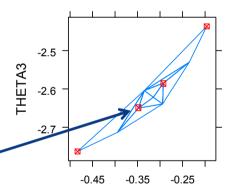
- ■Start with the initial parameter estimates  $(\theta_0)$ .
- Fix the parameters of interest at values away from  $\theta_0$  forming an initial set of simplexes.
- ■For each simplex (i): While  $(\int \text{new}_i d\theta - \int \text{old}_i d\theta) > \text{tol}$ • Store  $\int \text{old}_i d\theta = \int \text{new}_i d\theta$ 

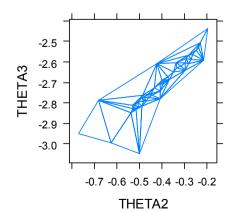
  - Refine points and run model at new points.
    - -Adding where error was previously observed; allows parallelization.
  - Compute ∫ new<sub>i</sub> dθ

Pharmacology

User-defined parameters are for point selection and  $\Delta$  integration tolerance.









#### **Future extensions**

- Analytical integration of n-dimensional likelihood surface will very significantly decrease estimation time in higher dimensions.
  - •Improved interpolation methods will similarly reduce integration time and number of iterations required.
- If estimating full dimensionality of the problem (all non-fixed θs, ηs, and σs), faster methods (using maxeval=0) can be used to simply sample the likelihood surface at that point.
  - ◆Typically will require above improvements in integration and/or interpolation.

