



SADDLE_RESET

more robust parameter estimation with a check for local practical identifiability

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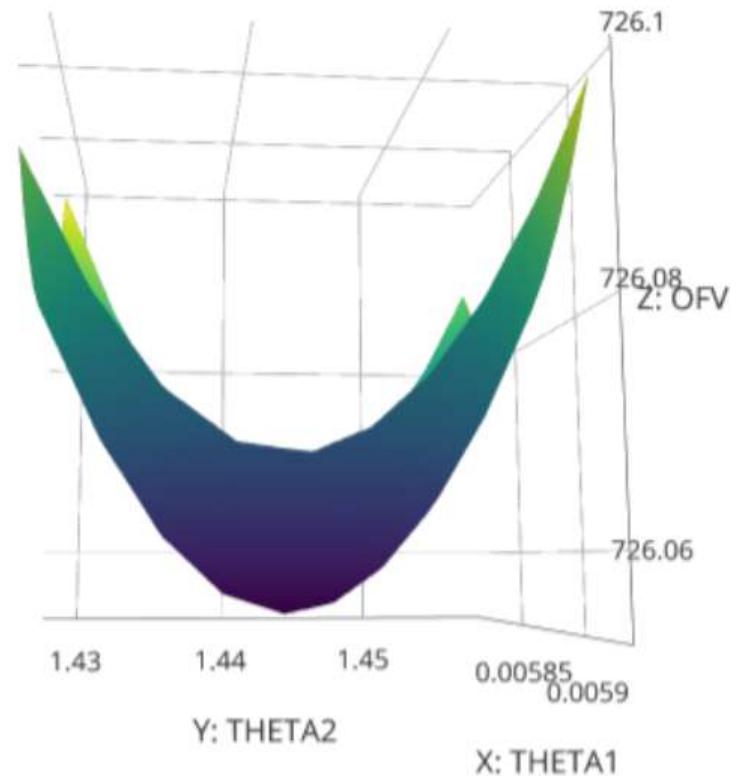
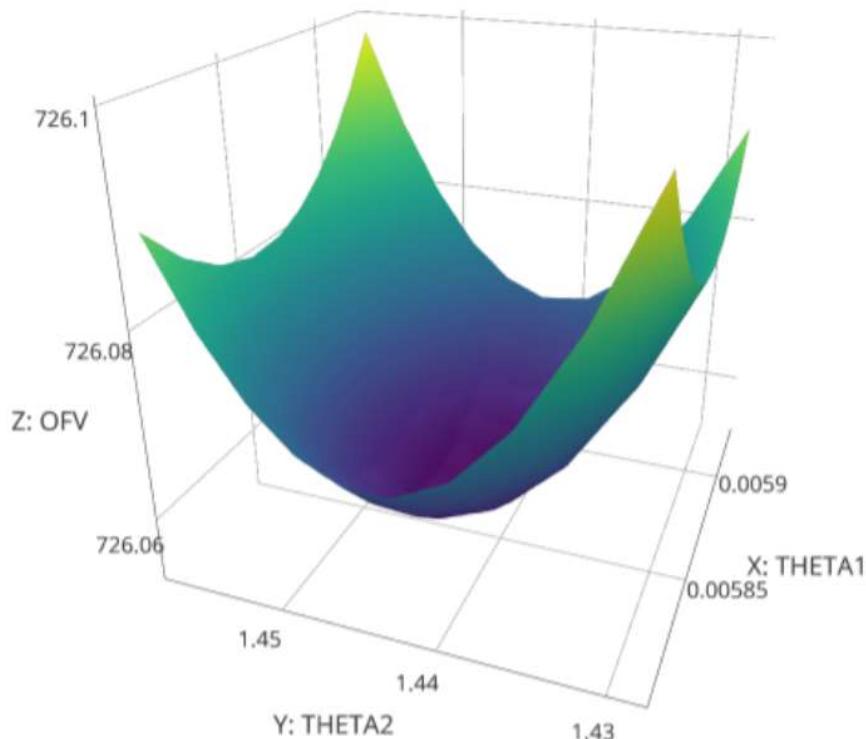
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Introduction



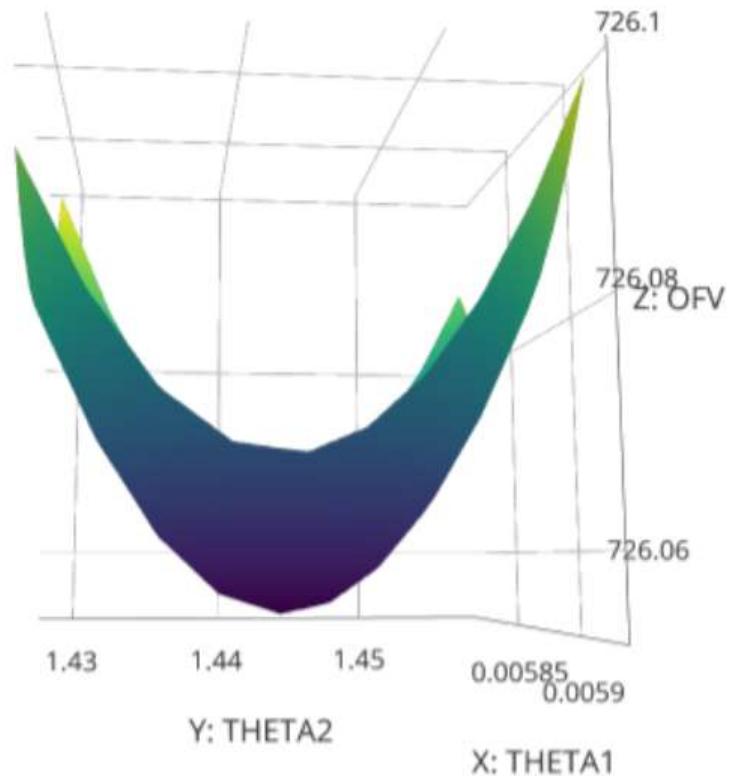
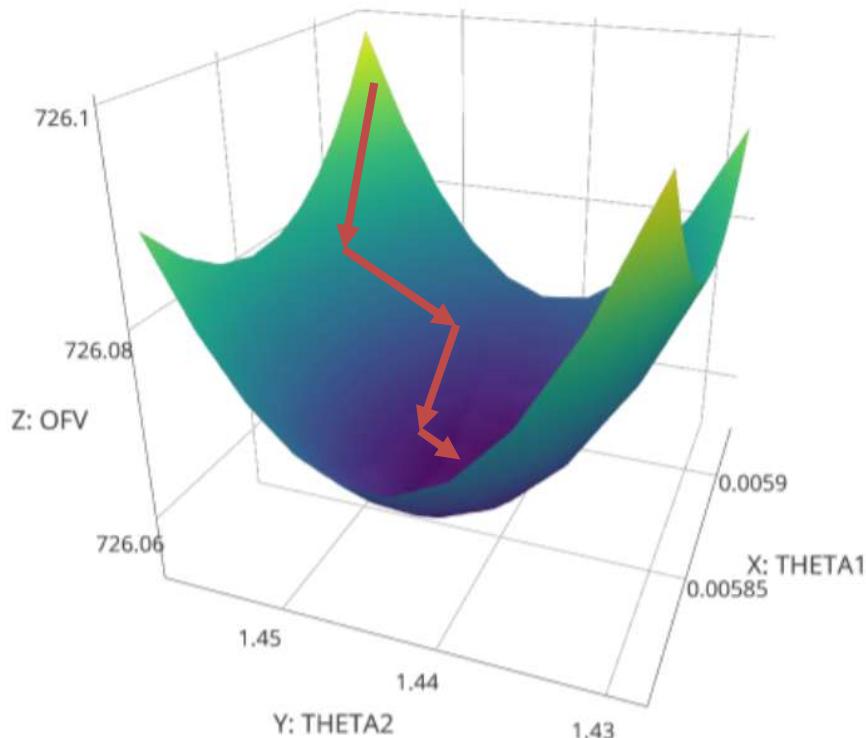


Introduction

Efficient and accurate parameter estimation is crucial to Pharmacometrics

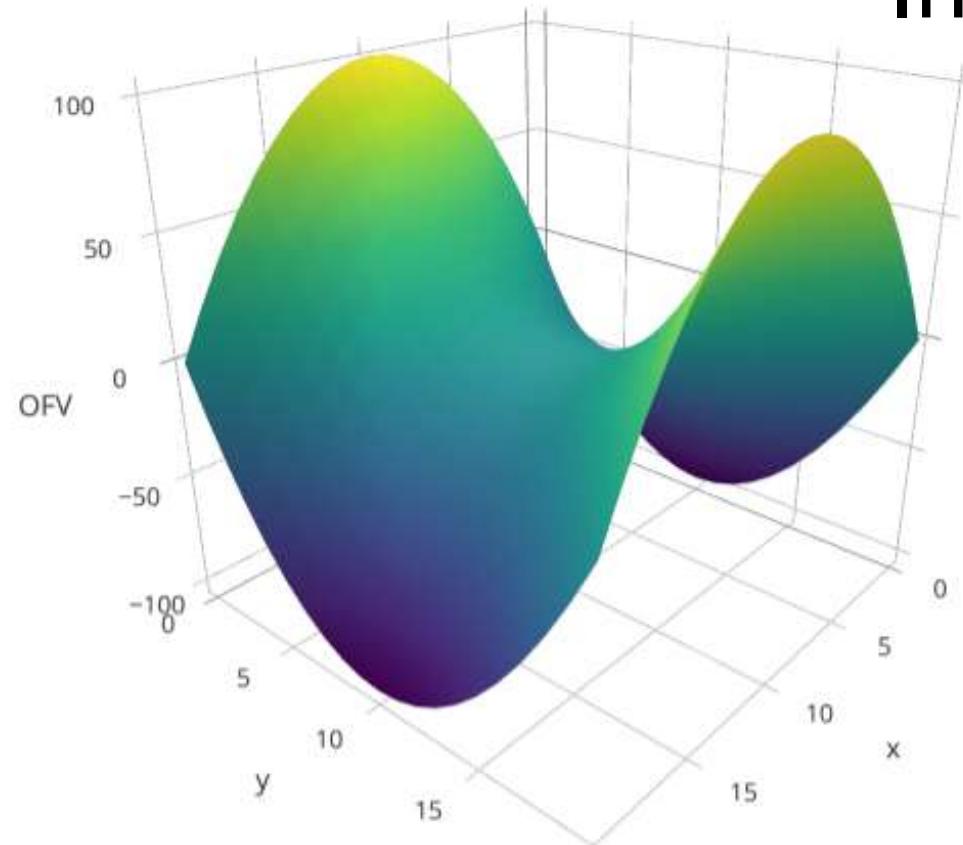
Methods that minimize gradients have become ubiquitous to our field

Introduction

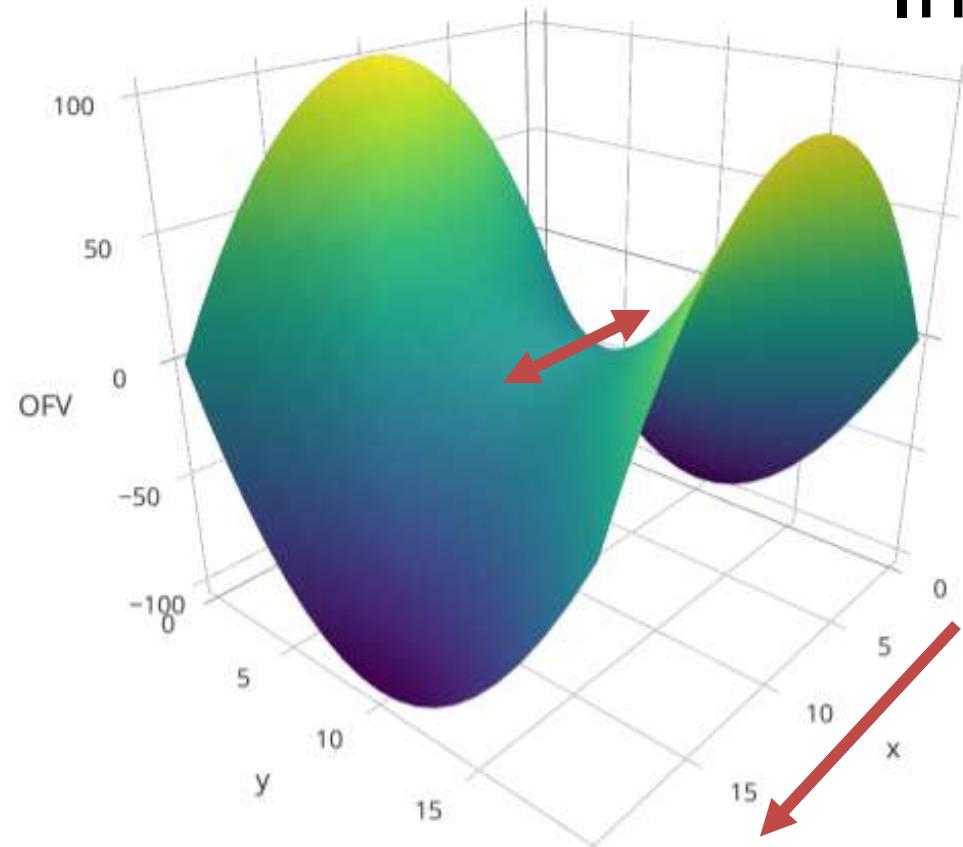




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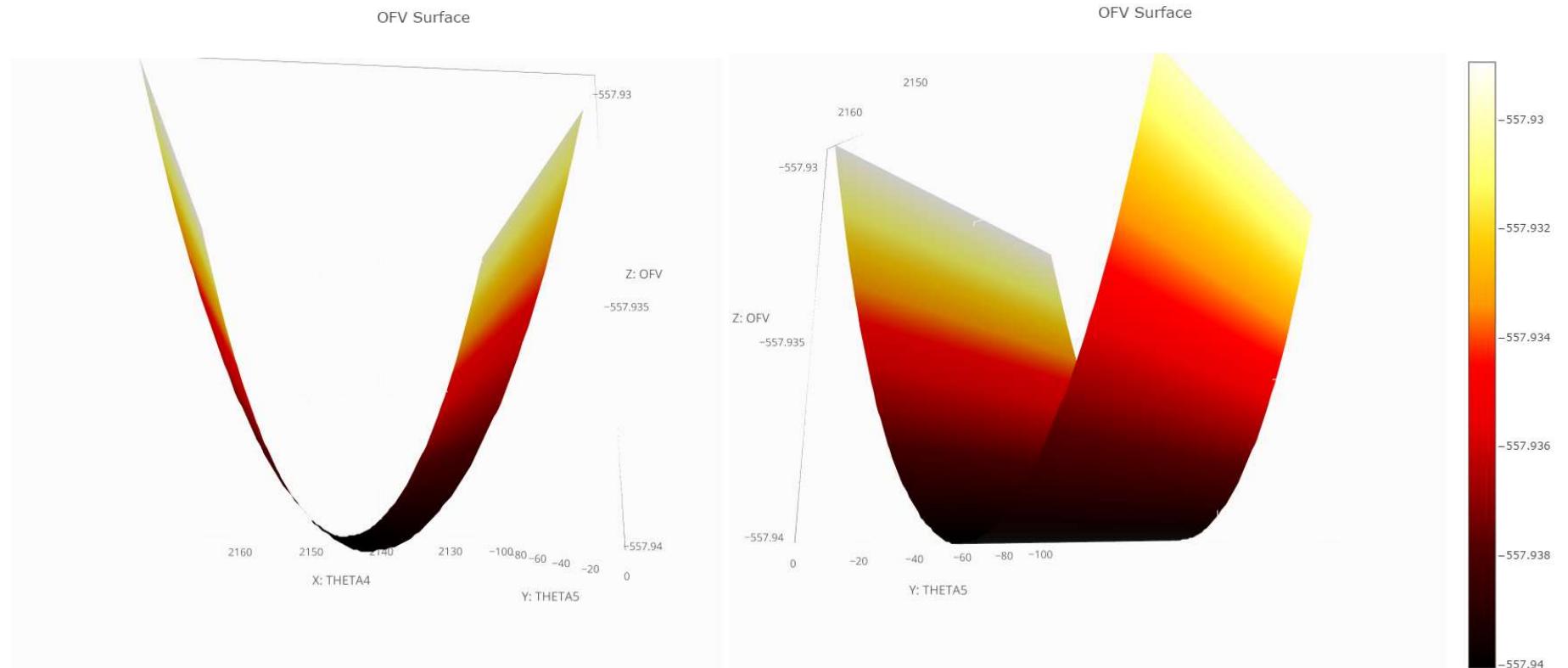


Introduction



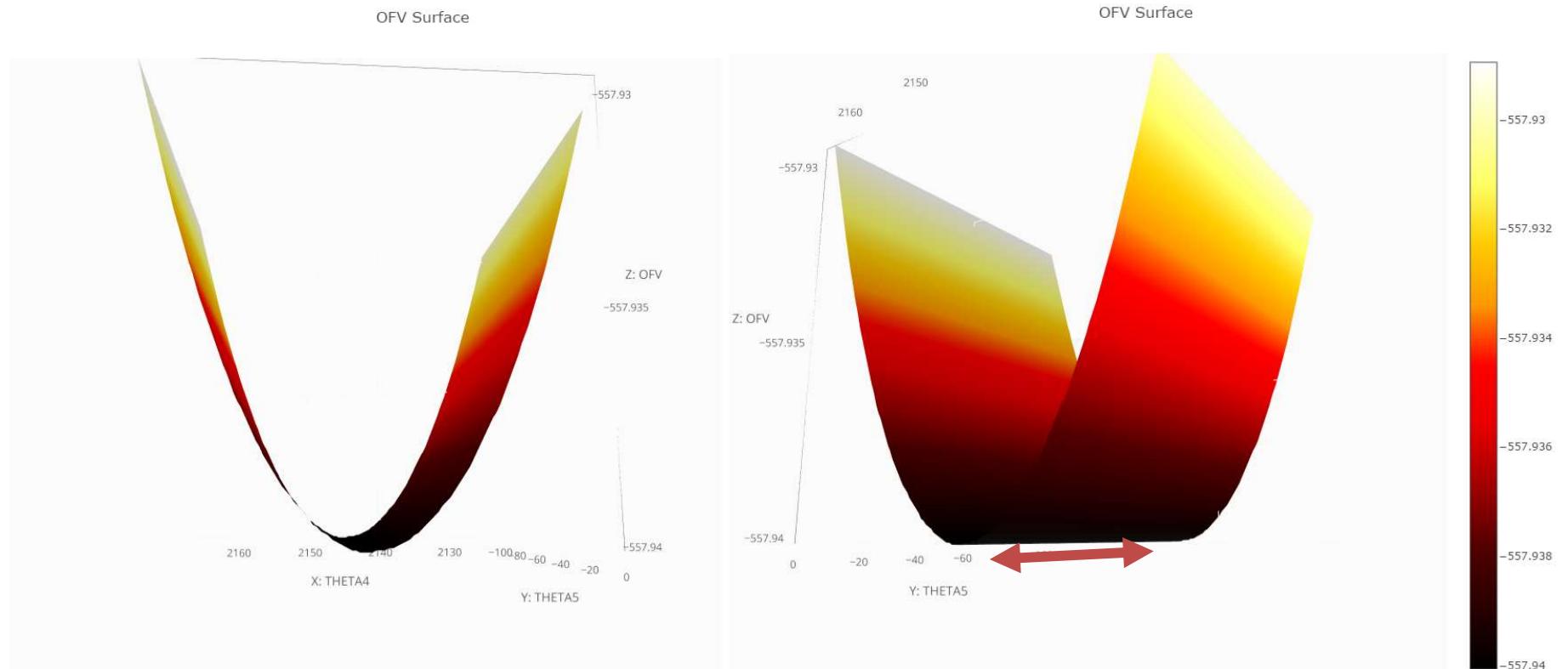


Introduction





Introduction





Introduction

After estimation we know little about the likelihood around our point

- We may not be in a minimum, but a saddle point
- We may have parameters that are non-identifiable
(non-estimable)



Introduction

Local:

- Our algorithm is only concerned with the immediate surroundings around the final estimate point on the
- $2\log(\text{likelihood})$ surface



Introduction

Local:

- Our algorithm is only concerned with the immediate surroundings around the final estimate point on the - $-2\log(\text{likelihood})$ surface

Practical:

- Our algorithm considers the model-data combination
- Structural identifiability is evaluated assuming infinite data



Brief Technical Description

1. Gradient minimization based estimation is performed as normal
2. The result is checked for zero gradients
 - If zero gradients are found, the associated parameters are reset to initial values and estimation is resumed
 - If none are found move to the next step



Brief Technical Description

3. The Hessian of the likelihood (the R-matrix) is eigendecomposed and the minimum curvature (including negative curvature) of the - $2\log(\text{likelihood})$ is identified
3. Parameters are changed along the minimum curvature
 - Step length calculated from curvature to give $\Delta\text{OFV} \approx 1$
4. Estimation is resumed from this new point



Brief Technical Description

Implemented in NONMEM 7.4

\$ESTIMATION

MAXEVAL=9999

NSIG=4

METHOD=COND INTERACTION

SADDLE_RESET=1

PRINT=5



Brief Technical Description

The BFGS approximation of the Hessian is used by default

- Constrained to be positive-definite
- As efficient as the true Hessian in our experiments

With option SADDLE_HESS=1 the true second derivative Hessian is used

- Computationally expensive (same as covariance step)



Brief Technical Description

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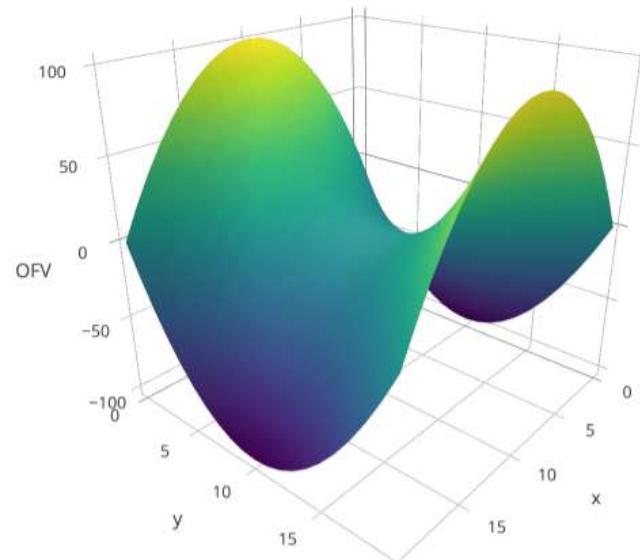
SADDLE_HESS=1

PRINT=5



Numerical Experiment 1

SADDLE_RESET can help us avoid saddle points





Numerical Experiment 1 - Models

Four published, fully identifiable models:

Model	Type	No Parameters	No IDs	No Samples	Solution
A	PK	9	177	1196	Closed Form
B	PK	13	93	274	Closed Form
C	PKPD	11	47	530	Diff. Eq.
D	PK	6	59	155	Closed Form

- A. Jönsson et al, Clinical Pharmacokinetics (2005), 44(8):863-78
B. Bergmann et al, The Pharmacogenomics Journal (2011) 11, 113–120
C. Wählby et al, Br J Clin Pharmacol. (2004) 58:367– 77.
D. Grasela Donn, Dev. Pharmacol. Ther. (1985) 8: 374-383



Numerical Experiment 1 - Method

Wide random perturbation around best parameter estimates

1,000 NONMEM estimations for each model

Successful estimation measured as $OFV < (\text{best known OFV} + 1)$

Some estimations will complete to saddle points and local minima



Numerical Experiment 1 - Method

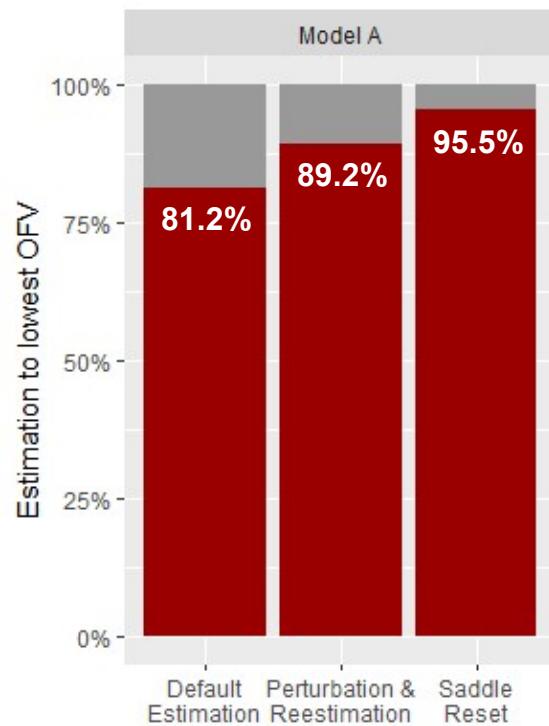
Comparison to perturbation of parameter values within 10%

Two estimations per model

- One from randomly perturbed estimates
- One from final estimates of previous estimation

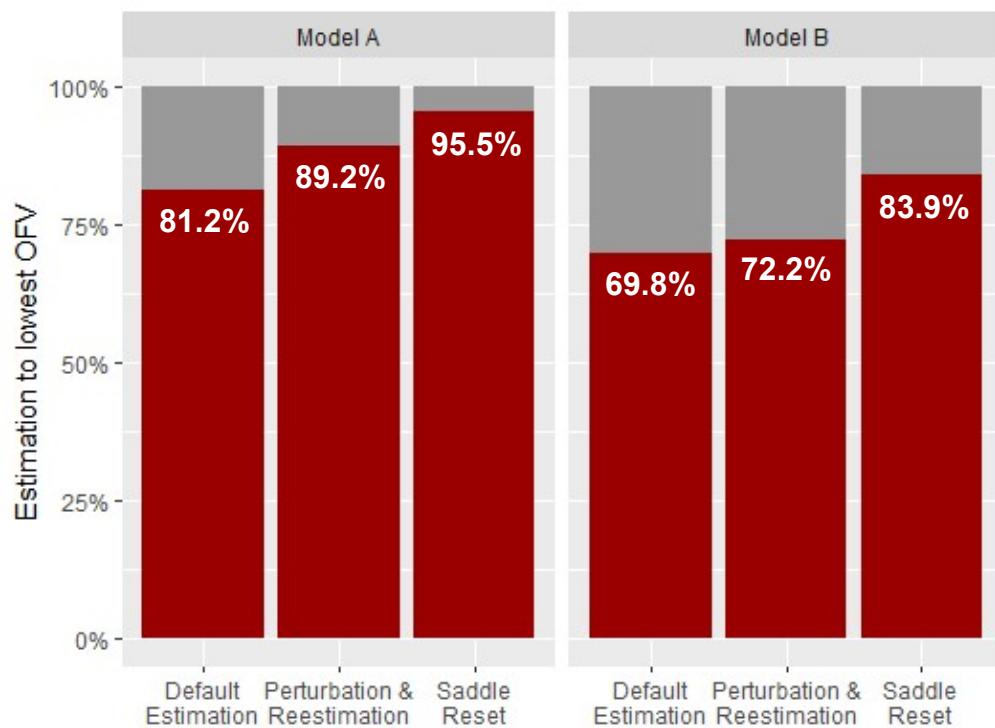


Numerical Experiment 1 - Results



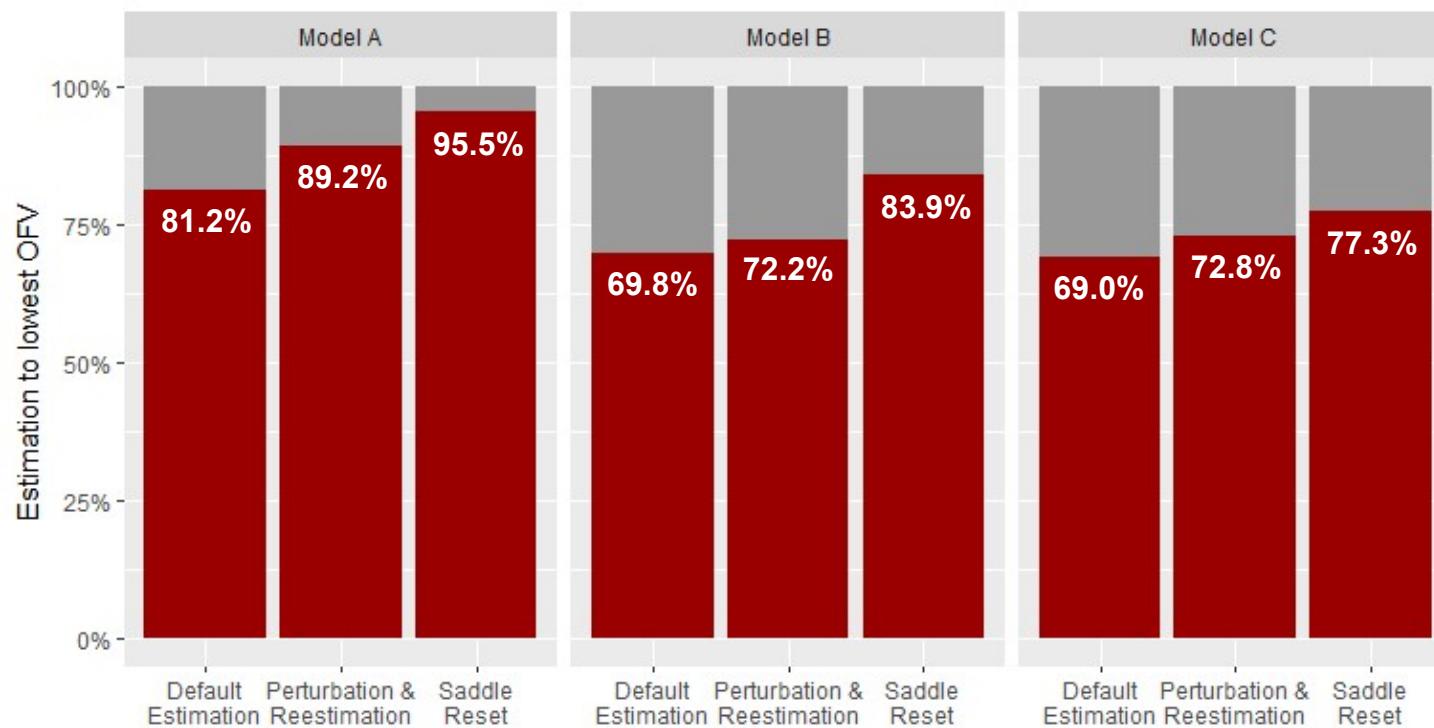


Numerical Experiment 1 - Results

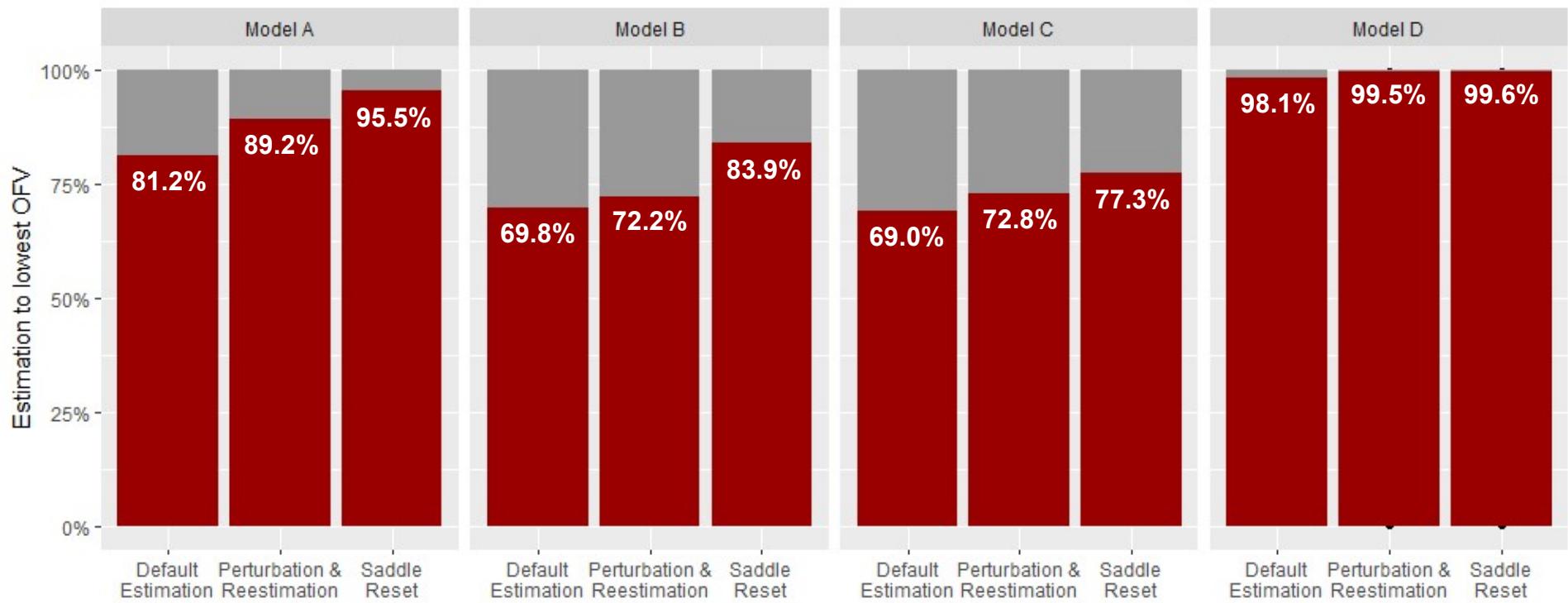




Numerical Experiment 1 - Results



Numerical Experiment 1 - Results





Numerical Experiment 1 - Results

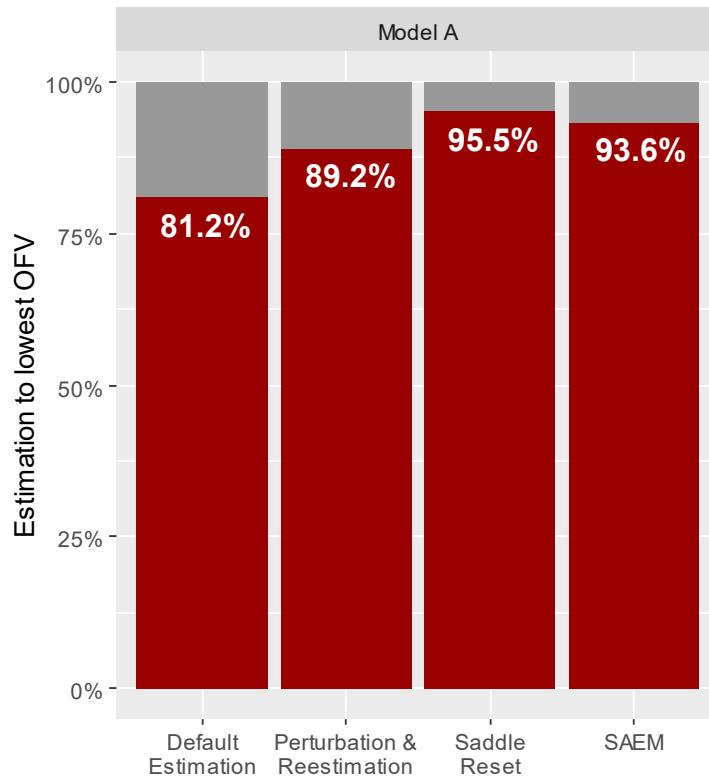
Does it matter if we end up in a saddle?

Three saddles and the lowest known minimum for Model A

OFV	Prop Err	V1	Q	V2	CL	CLCR-CL	WT-V1	IIV CL	IIV V1
-2029.10	0.222	10.360	4.280	5.616	2.893	0.019	0.014	0.055	0.008
-2233.49	0.169	7.744	13.909	7.391	1.928	0.050	0.021	0.079	0.174
-2339.87	0.166	6.770	18.959	8.354	2.882	0.019	0.027	0.053	0.239
-2346.79	0.165	7.913	13.122	7.170	2.914	0.019	0.019	0.054	0.162



SAEM Comparison

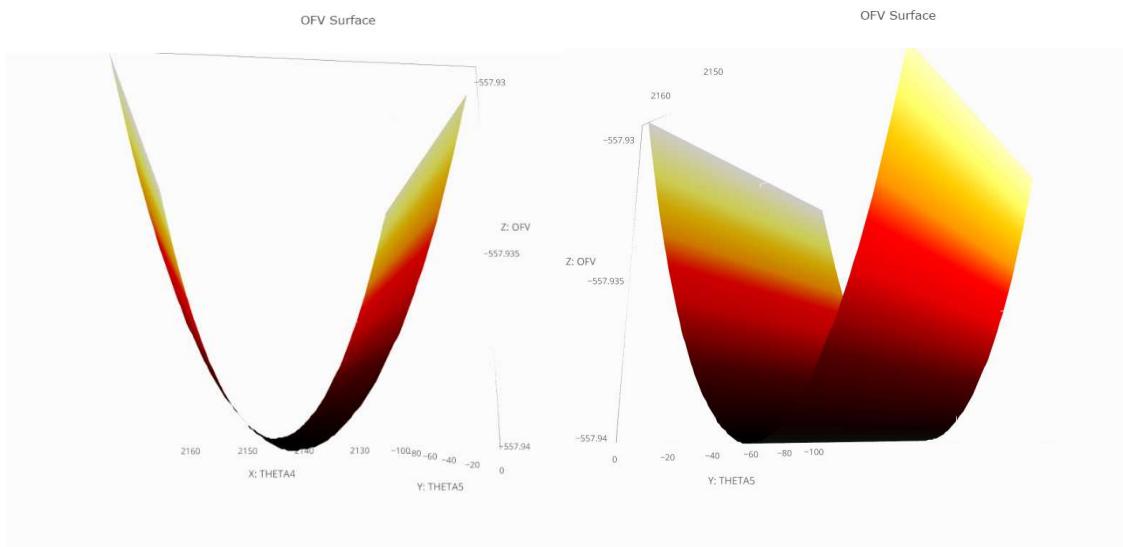


- Saddle Reset has a success rate comparable to a Monte-Carlo method such as SAEM



Numerical Experiment 2

SADDLE_RESET can help us discover local practical non-identifiability





Numerical Experiment 2 - Models

Two models representing two identifiability issues

- E. Practically non-identifiable Emax model-data combination
- F. Structurally non-identifiable model



Numerical Experiment 2

Model E – Multiple parameter values with same OFV

OFV	Placebo Eff.	Baseline	Emax	ED50	Gamma
-2710.52	0.13338	2.5084	0.30203	39.1895	104.173
-2710.52	0.13339	2.5083	0.30229	33.5416	12.1728
-2710.52	0.13348	2.5083	0.30228	29.8024	7.17949

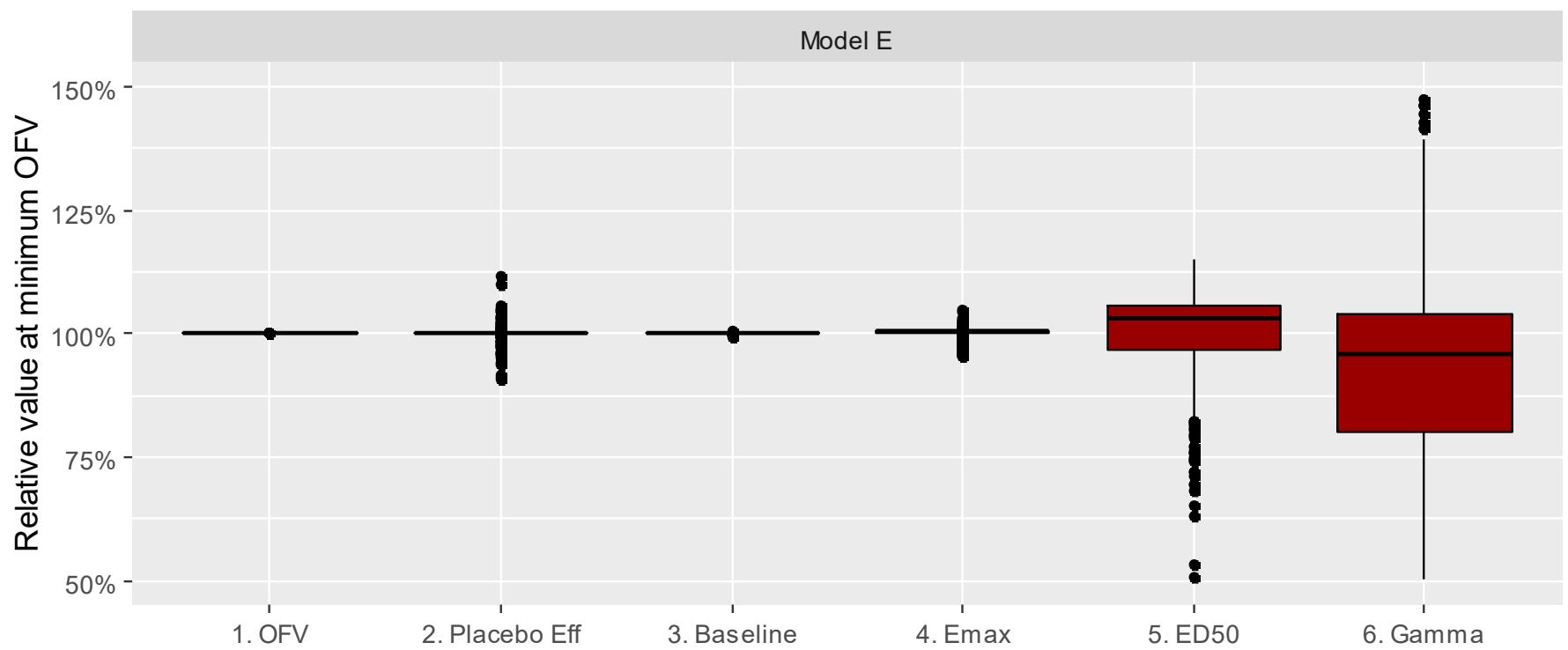


Numerical Experiment 2 - Results

SADDLE_RESET found alternative parameter values with the same OFV in 96% and 95% of 1,000 cases for models E and F respectively



Numerical Experiment 2 - Results





SADDLE_RESET Benefits

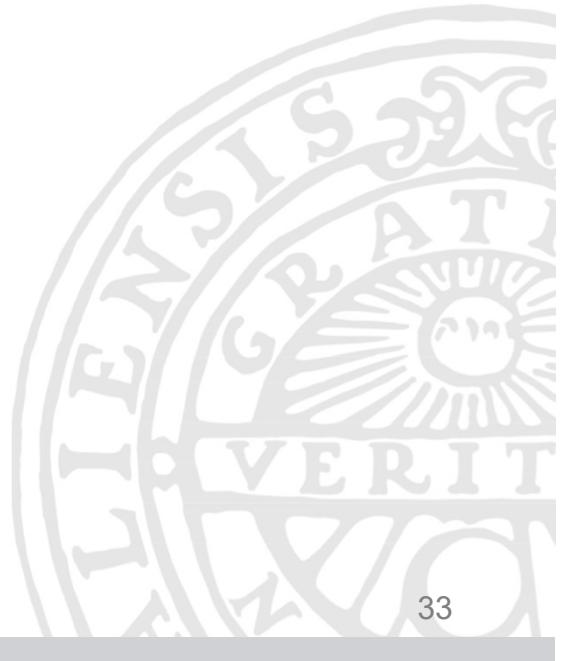
Improves confidence in parameter estimates

- Expose local practical non-identifiability
- Avoid saddle points

I hope you will add this to your smorgasbord of methods



Thank you!





NONMEM 7.4

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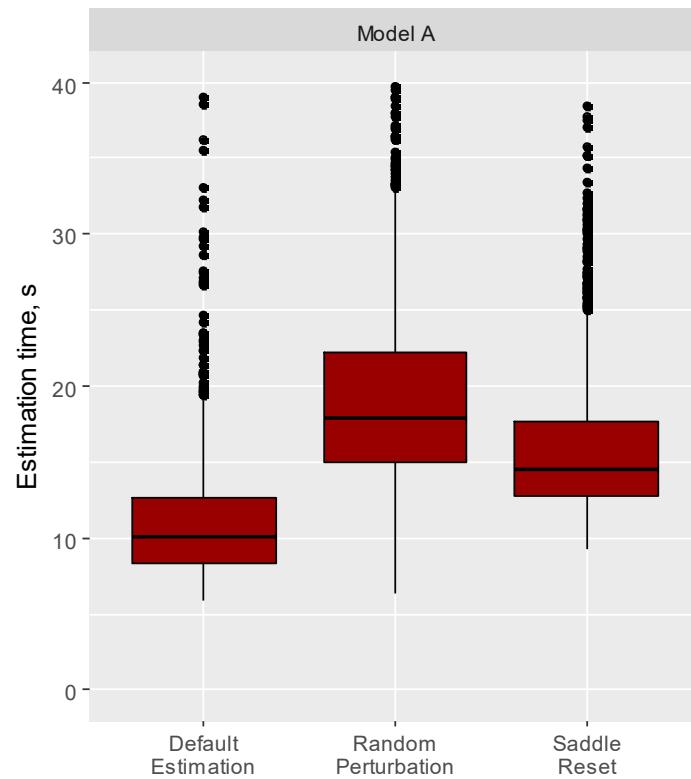
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Extra Materials

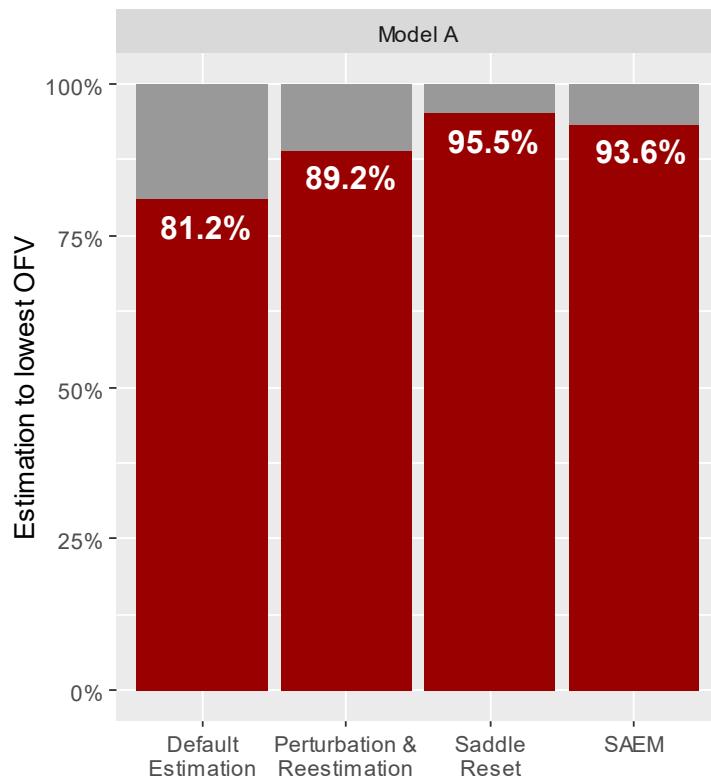
Runtime



- Not a rigorous runtime experiment
- Not same execution environment
- 1,000 samples



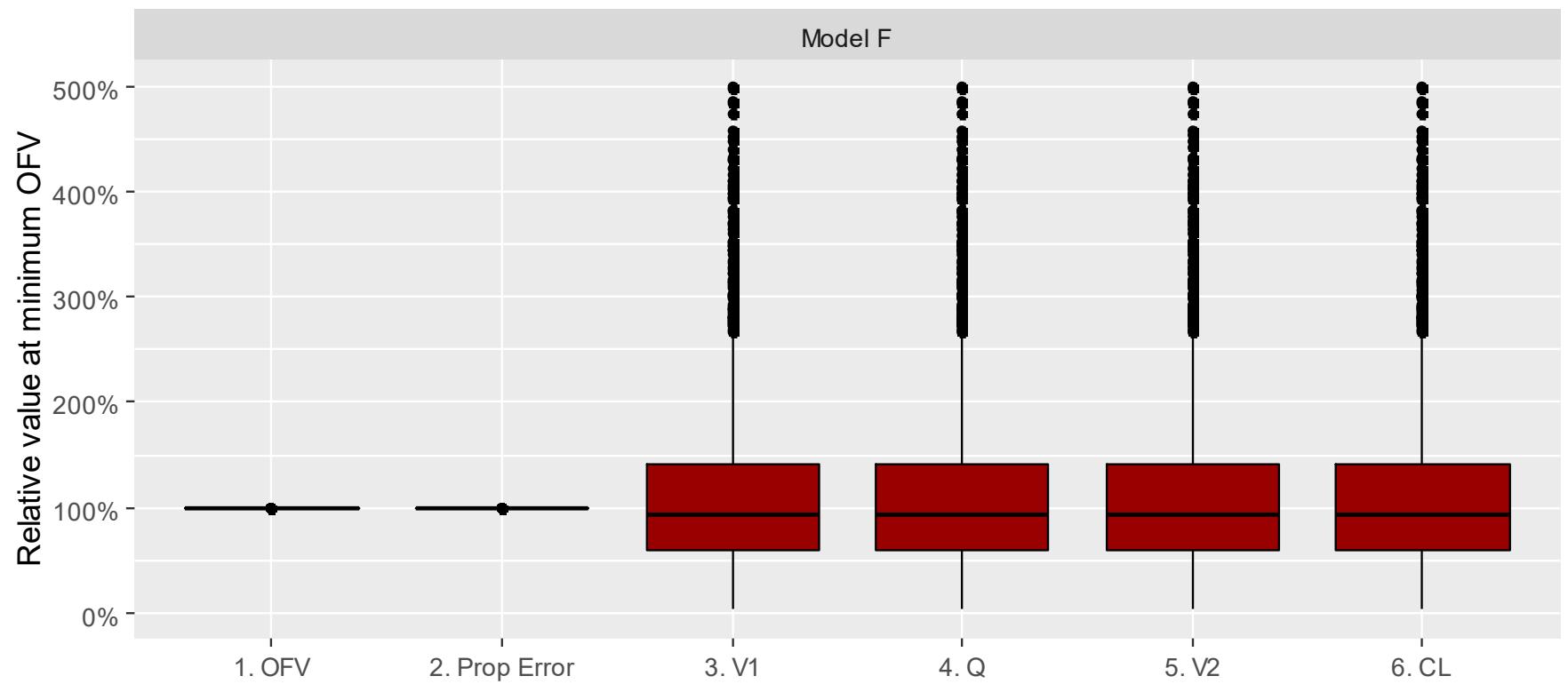
SAEM



- Comparable success rate as a Monte-Carlo method such as SAEM



Structurally Non-Identifiable Case





Saddle Examples

OFV	Prop Err	V1	Q	V2	CL	CLCR-CL	WT-V1	IIV CL	IIV V1
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