

# Sloppiness in Multiparameter Models

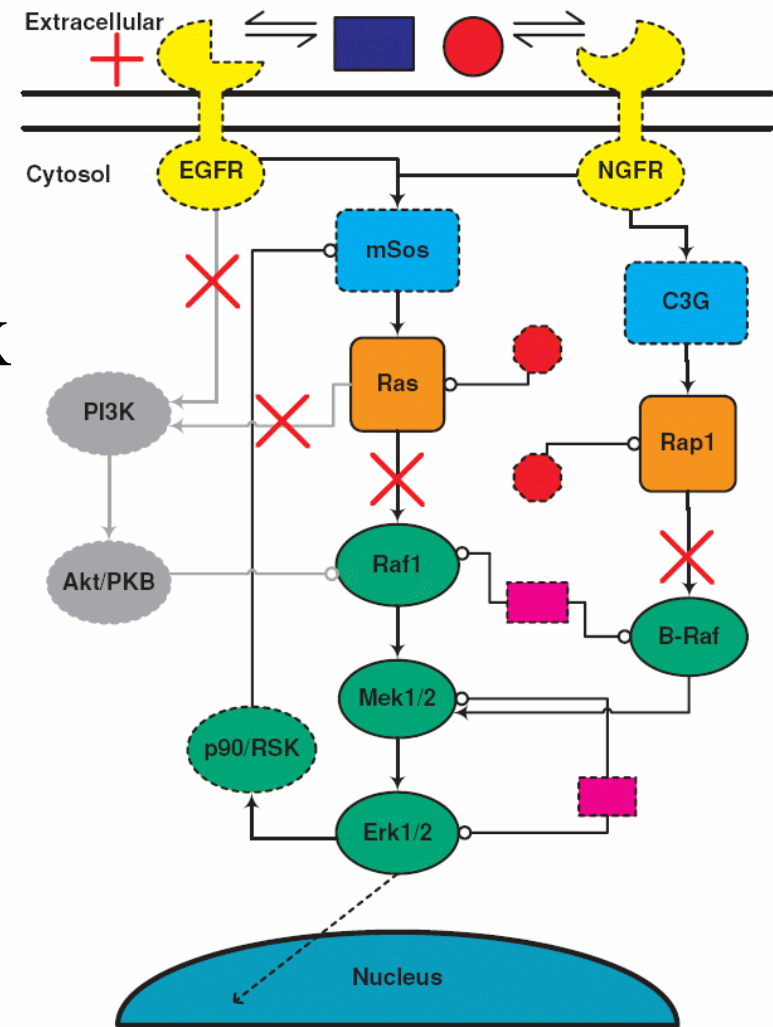


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July 28, 2008

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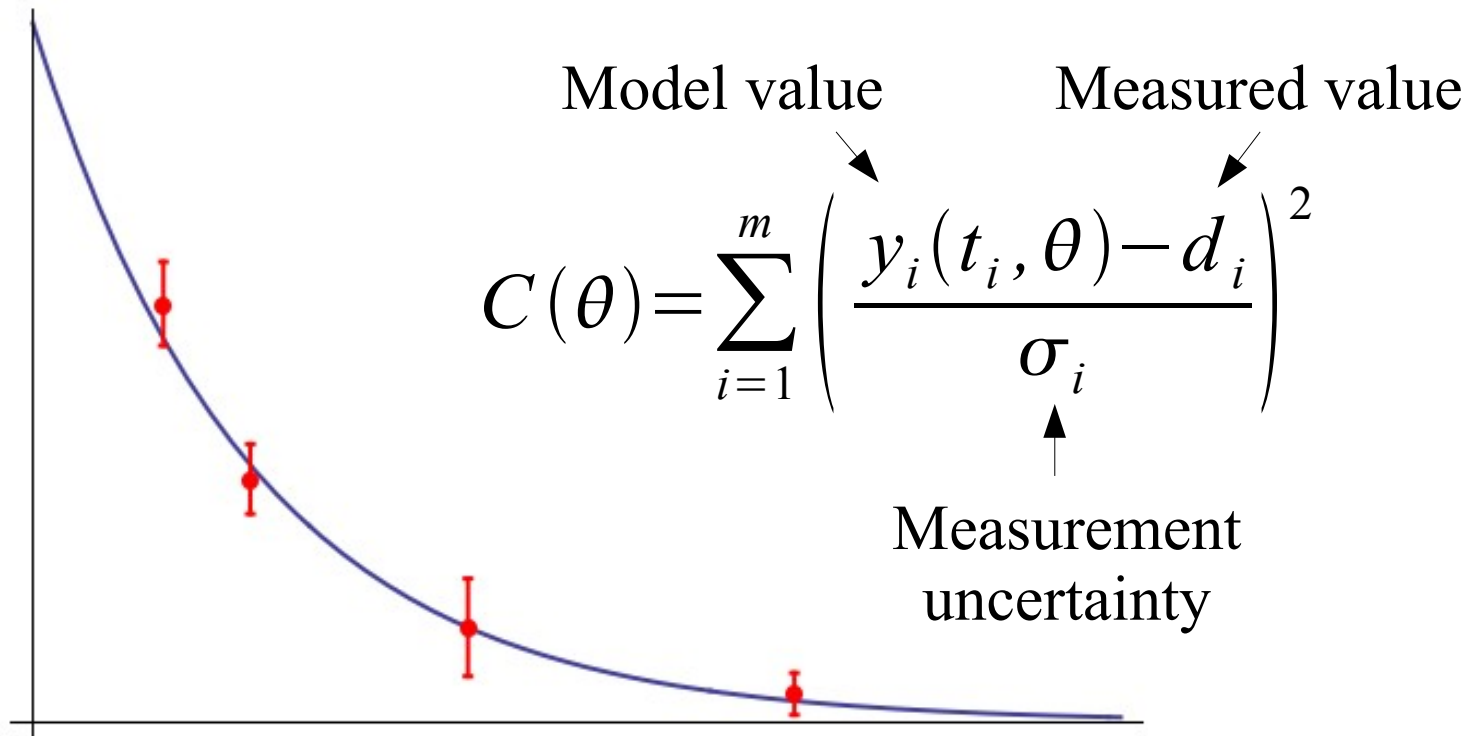
# Motivation

- › Biological models have lots of parameters, and they control the output in complex ways.
- › It's often hard to measure these parameters.
- › How does this affect model predictions? What predictions can we trust?



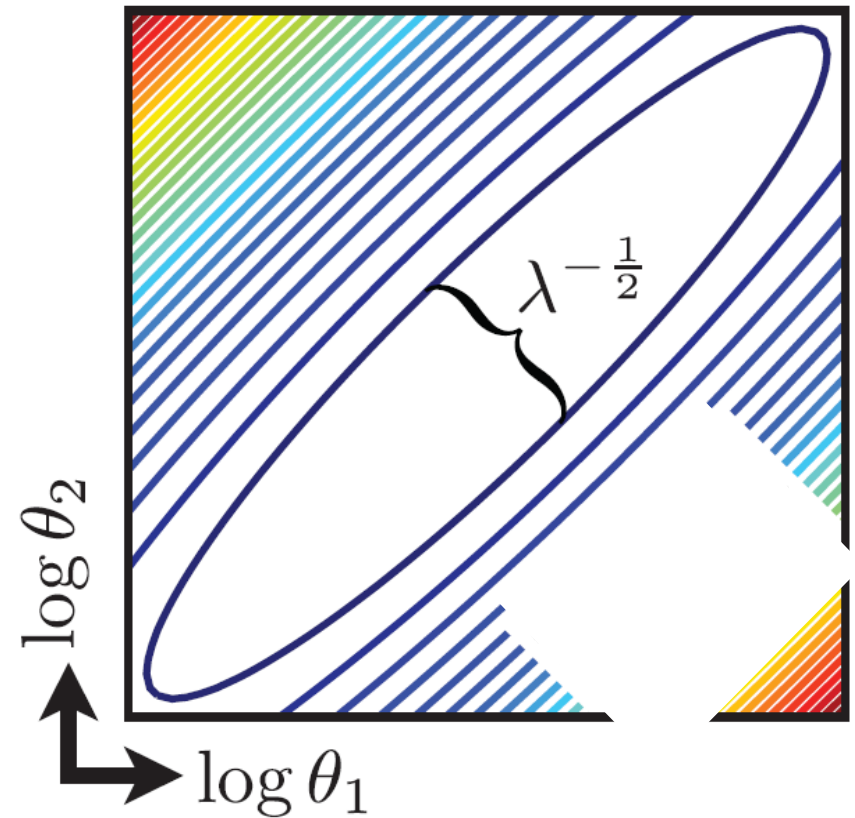
# Cost Landscape

- › A set of parameters  $\theta$  has a cost based on how well the model fits measured data.
- › We usually use a squared residuals cost.



# Cost Landscape

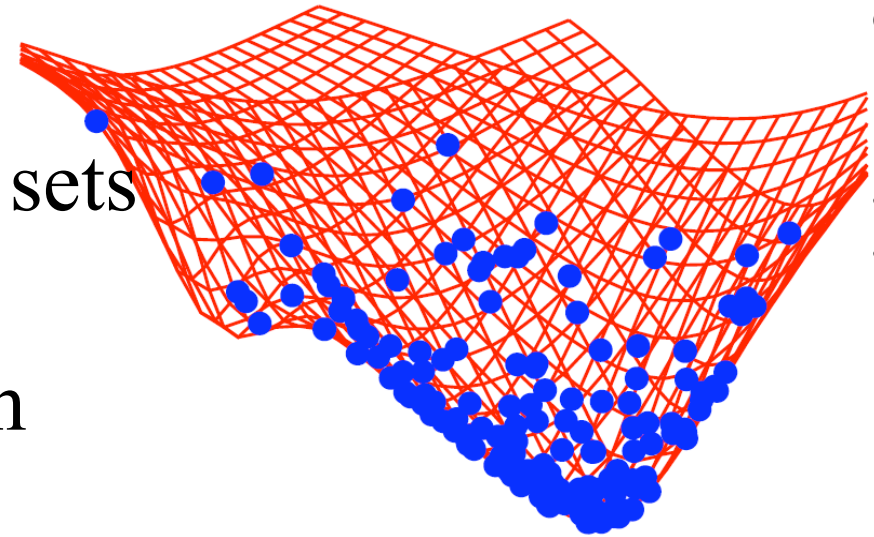
- › Locally around the best-fit point we can approximate the cost as quadratic.
- › The matrix of 2<sup>nd</sup> derivatives (Hessian) gives us the quadratic expansion.



# Making Sensible Error Bars

› Most thorough method:  
Bayesian analysis using Monte  
Carlo sampling

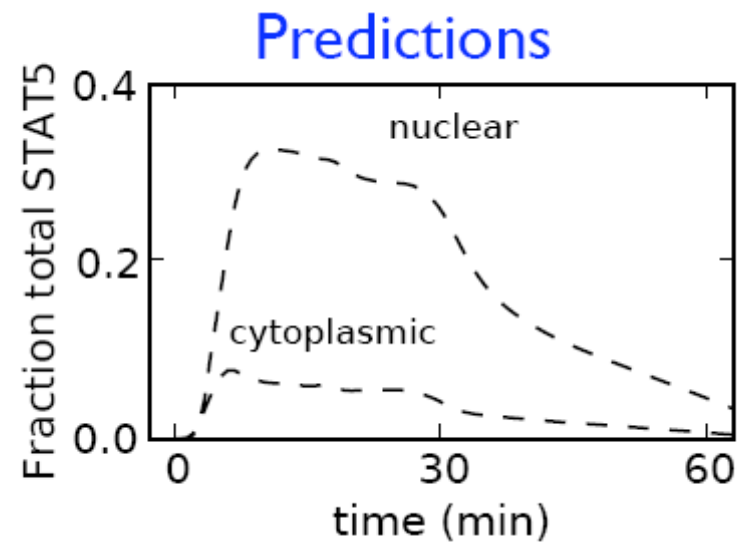
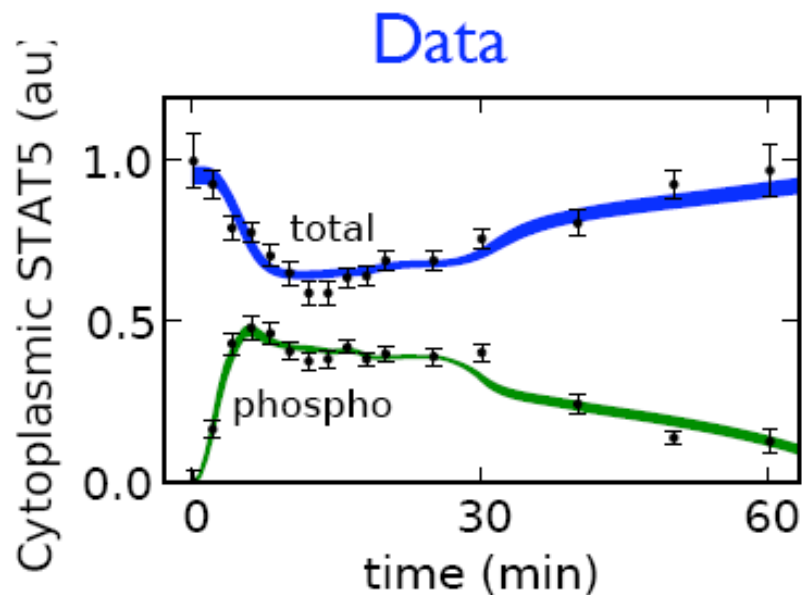
1. Sample from all parameter sets that fit the data;
2. Find prediction output from each;
3. Calculate mean, standard deviation, etc.



“Stat mech in  
model space”

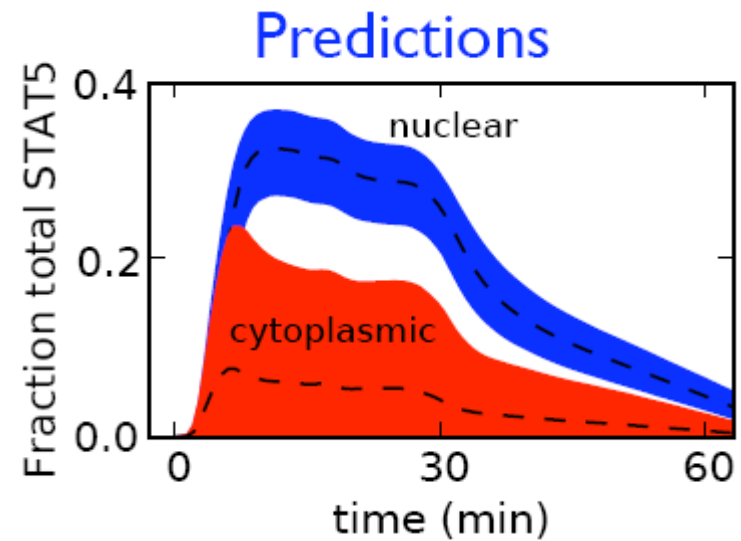
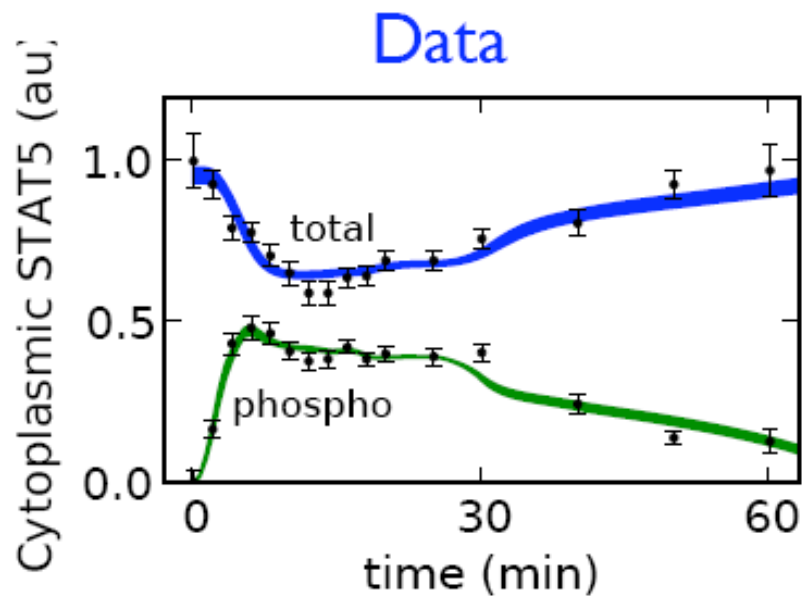
› Example: Brown et al., Phys. Biol. 1: 184-195

# Making Sensible Error Bars



Figures courtesy Ryan Gutenkunst

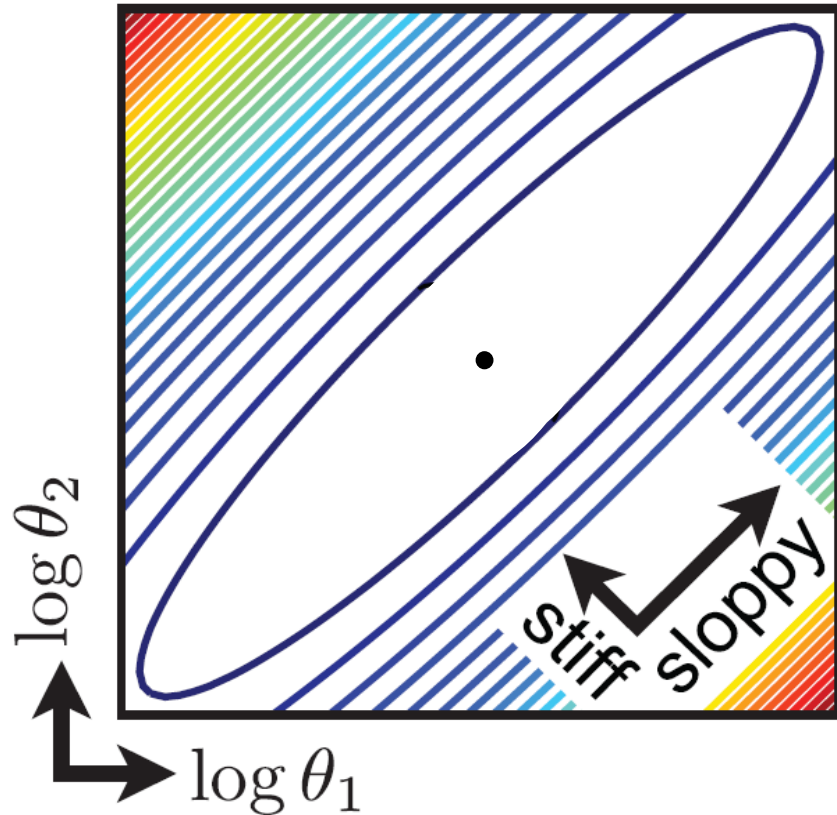
# Making Sensible Error Bars



Figures courtesy Ryan Gutenkunst

# “Sloppiness”

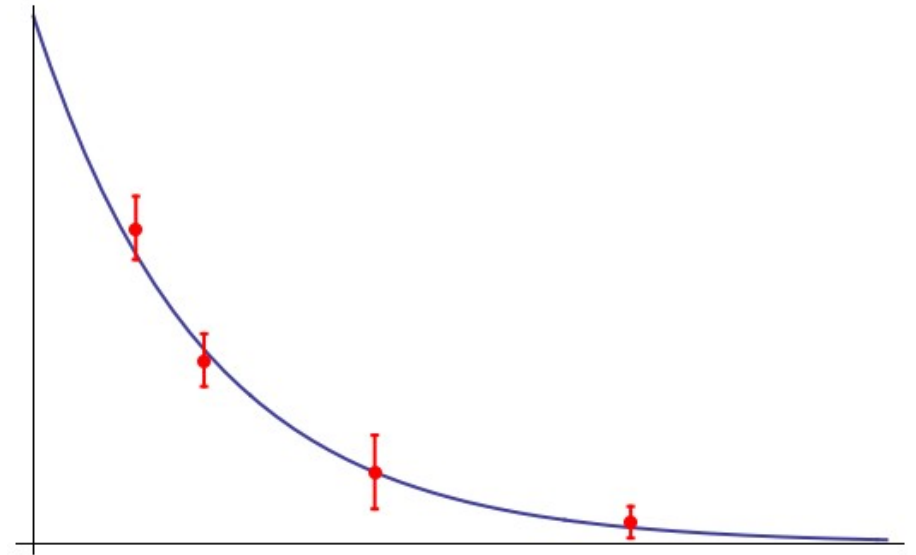
- › Nonlinear multiparameter models are “sloppy”: orders of magnitude more sensitive to changes in certain directions in parameter space.





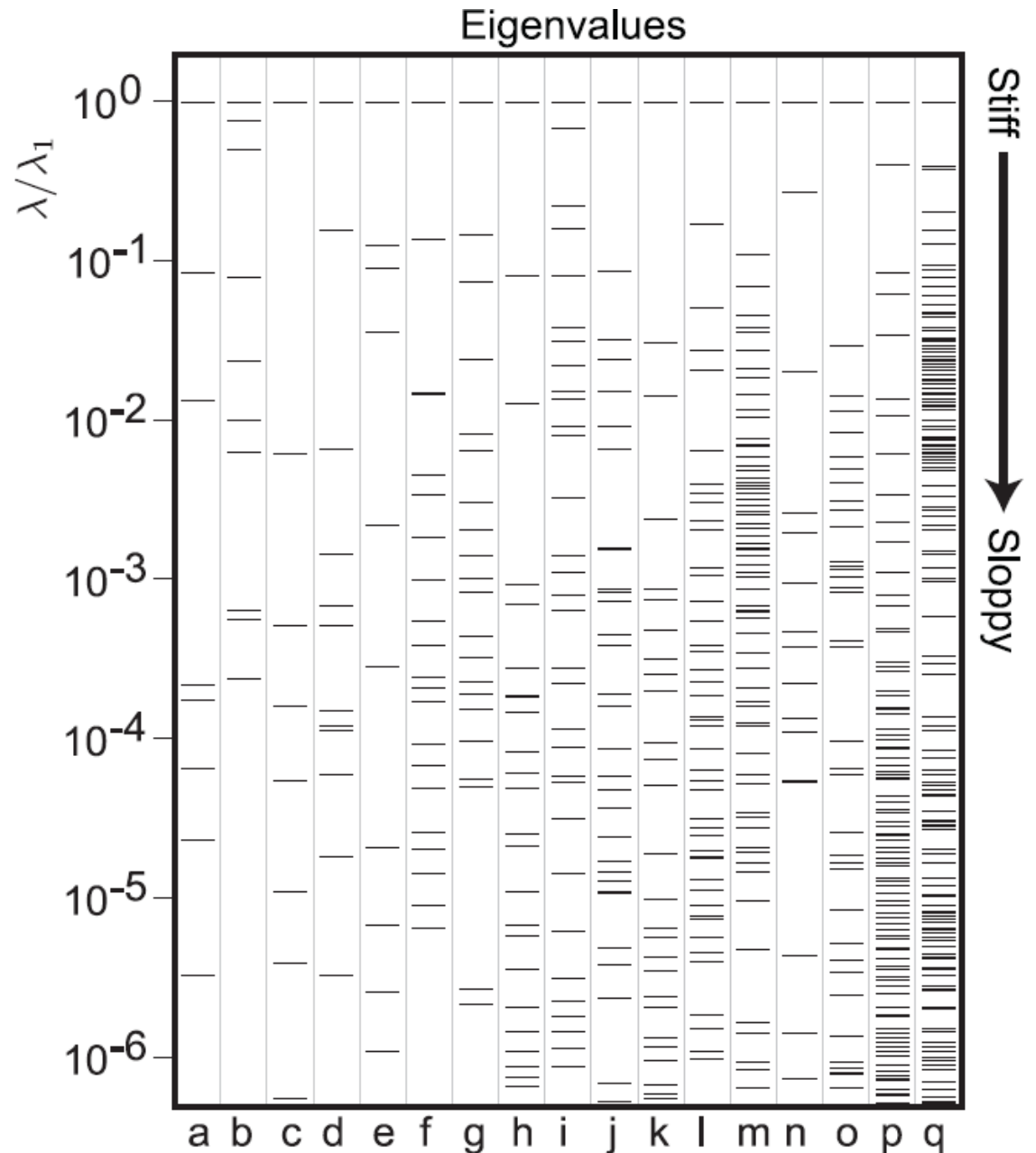
# Measuring Sloppiness

- 1) Define cost – usually squared residuals
- 2) Find Hessian  
(2<sup>nd</sup> derivative matrix of cost)



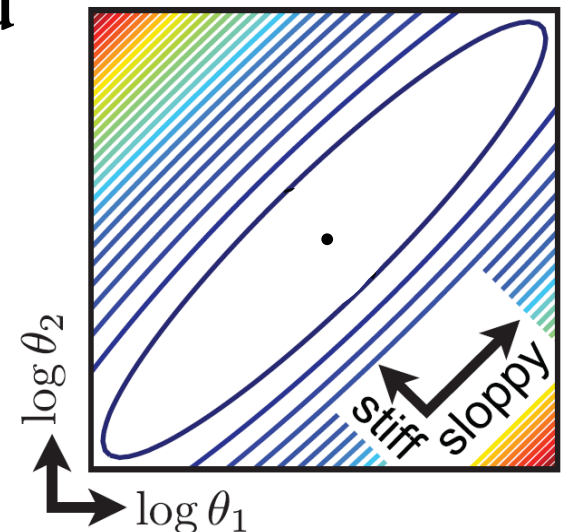
- › The **eigenvalues of the Hessian** tell you about sensitivity along eigendirections in parameter space
- › Produces a “sensitivity spectrum”

- › Hallmarks of sloppiness:
1. Large range of eigenvalues
  2. Eigenvalues roughly evenly spaced in log space



# Implications of Sloppiness

- › Large range means cost contour ellipsoids are routinely stretched by a factor of 1000 (the aspect ratio of a human hair).
- › Even spacing means there is no well-defined cutoff between “important” and “unimportant”.



# Universality of sloppiness

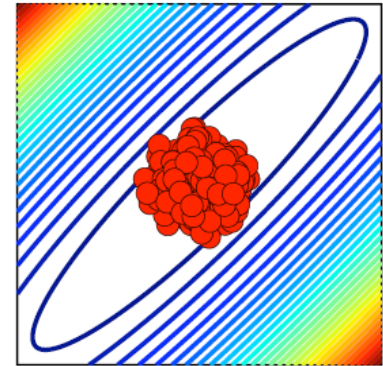
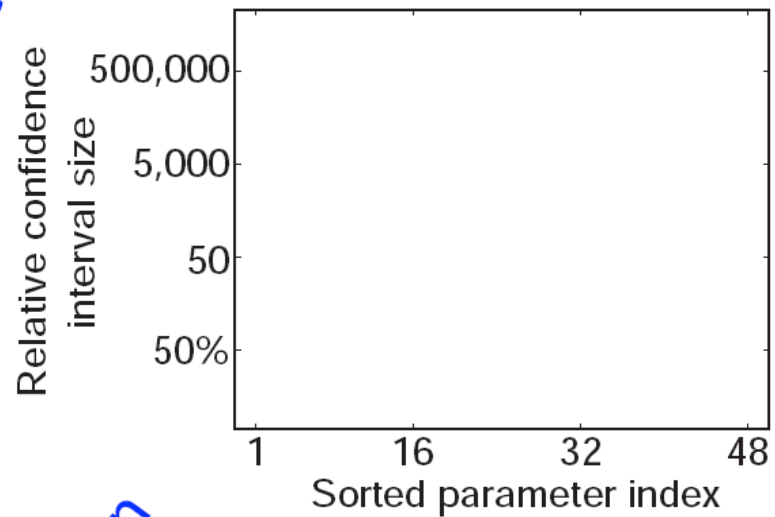
- › Sloppiness has been found in every biological system analyzed (17 so far), and more:
  - Interatomic potentials, particle accelerator design, sums of exponentials...
- › May be a “universal” feature of nonlinear multiparameter fitting problems.

# Parameter Uncertainty is Inevitable

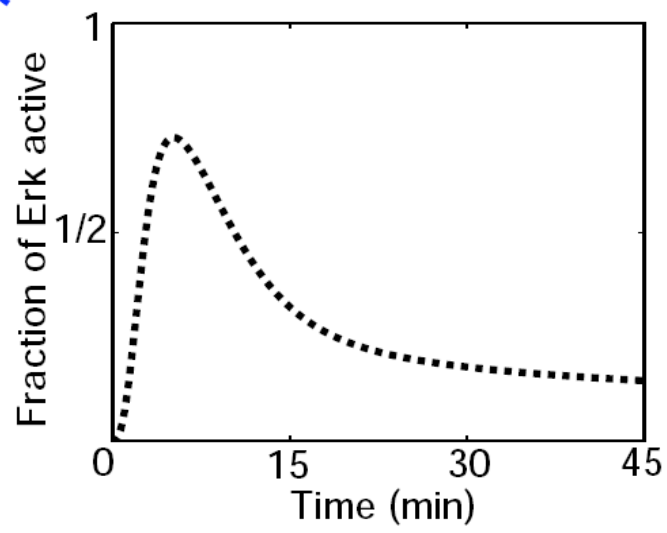
- › Sloppiness provides an answer for why fits can lead to large uncertainties in parameter values.
- › Large parameter uncertainty does not imply large uncertainty in predictions.

# Uncertainties

Parameters

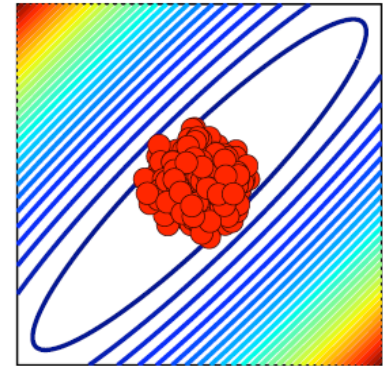
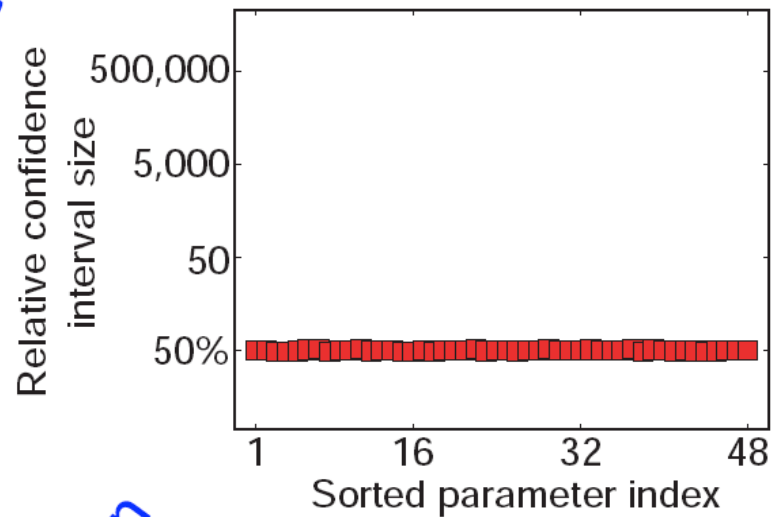


Prediction

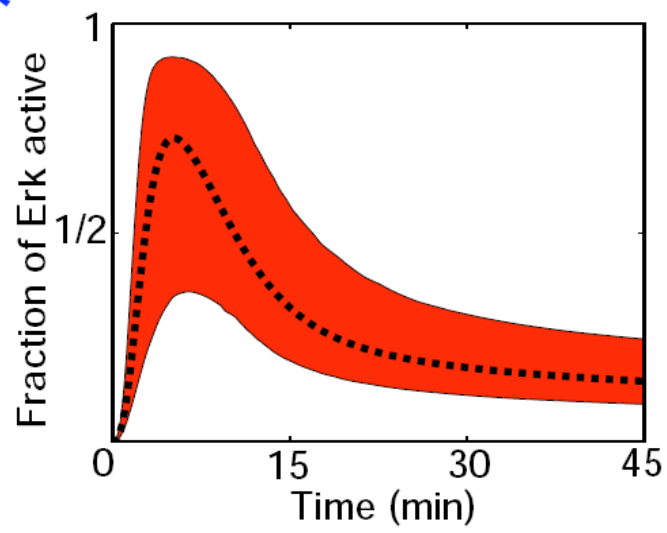


# Uncertainties

Parameters

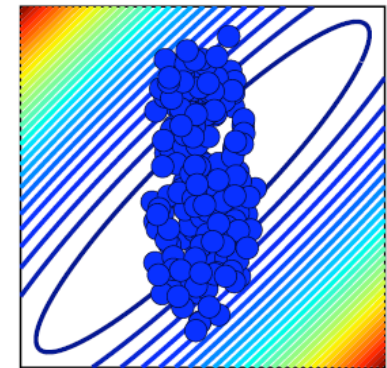
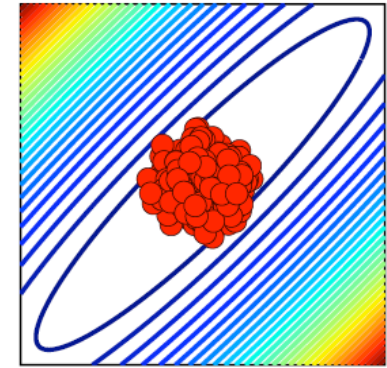
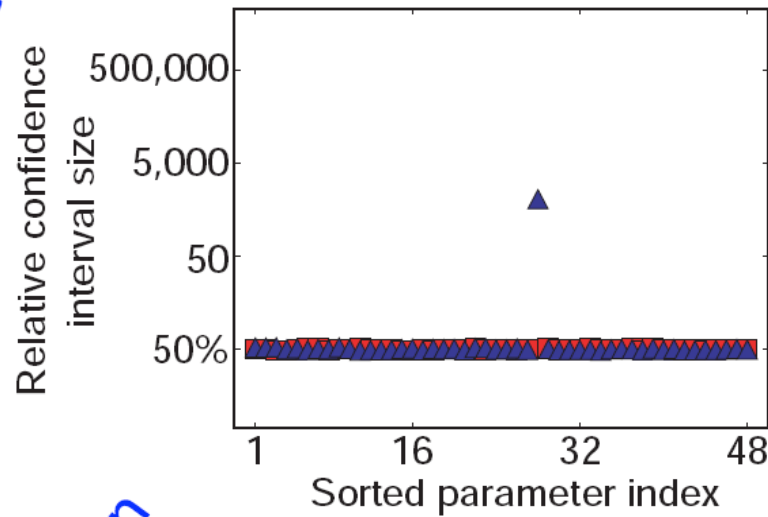


Prediction

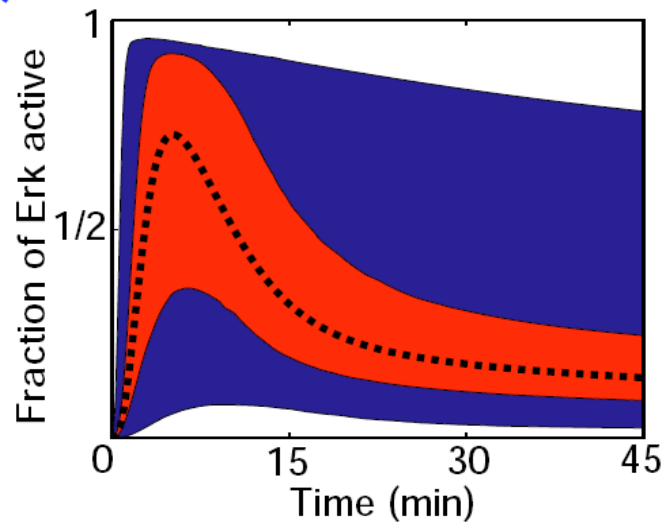


# Uncertainties

Parameters



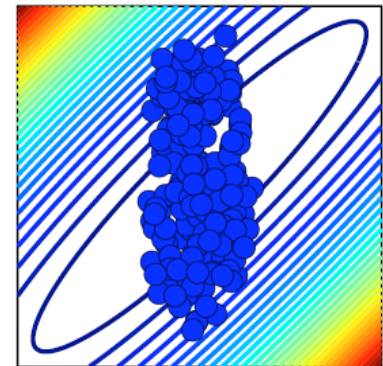
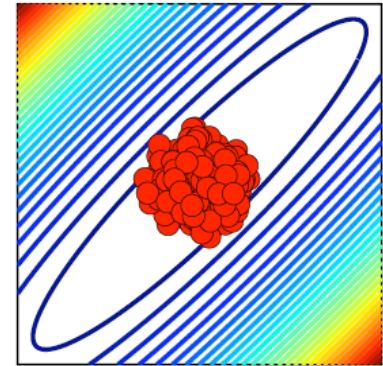
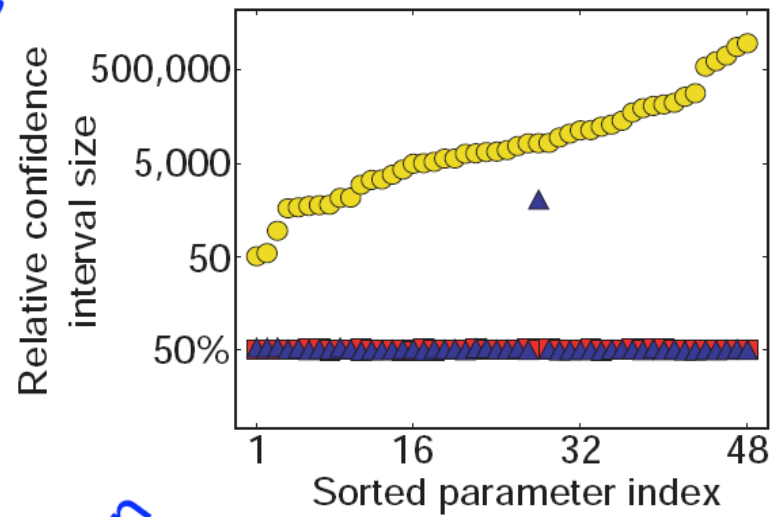
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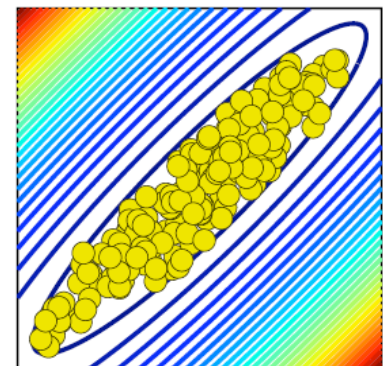
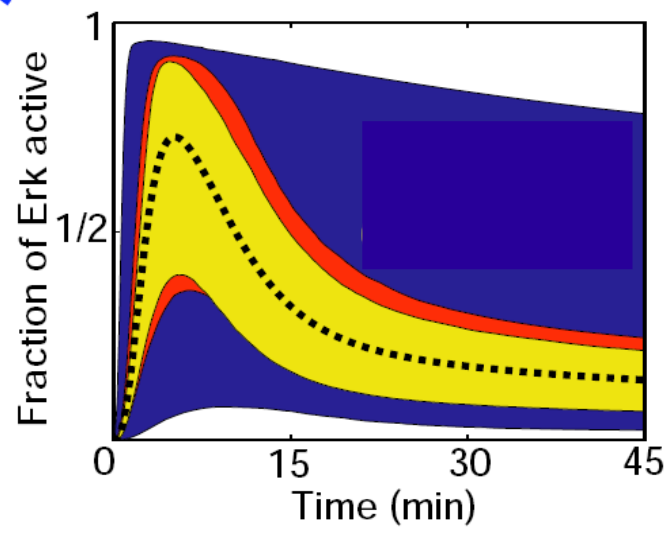


# Uncertainties

Parameters



Prediction



# Sloppiness is “Real”

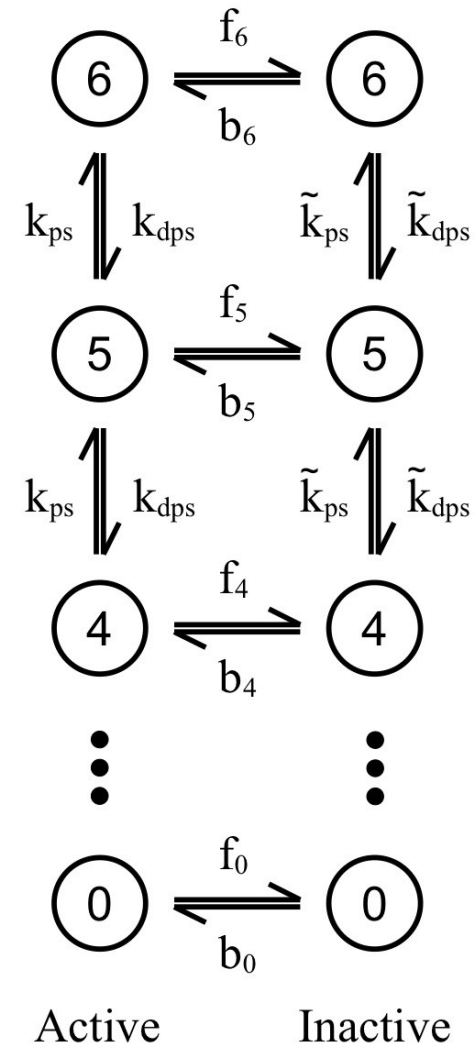
- › Is it due to too few data points?
  - No; even for data the model can fit perfectly, sloppiness persists.
- › Is it due to the local approximation?
  - No; principal component analysis of Monte Carlo ensembles still displays sloppiness.

# SloppyCell

- › Computing environment for simulating and analyzing biochemical networks (or any system of ODEs)
- › Structure for optimization and efficient calculation of ensembles
- › Supports Systems Biology Markup Language (SBML)
- › Implemented in Python

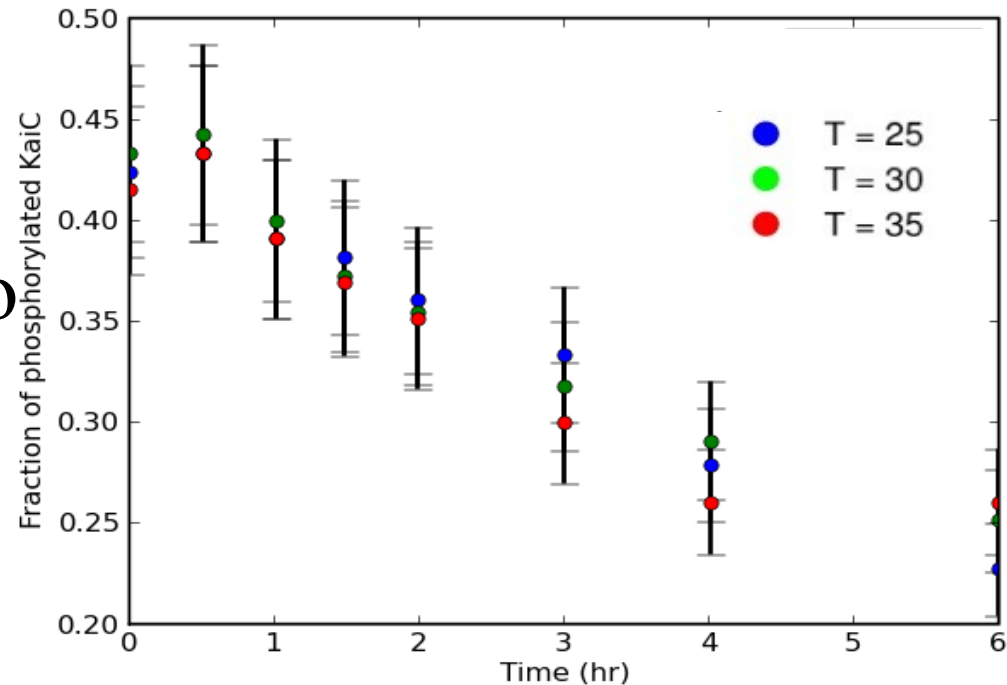
# Is robustness a delicate balancing act?

- › A subset of the circadian rhythm network in cyanobacteria
- › The phosphorylation decay rate is measured to be robust to temperature change, even when individual (de)phosphorylation rates would double



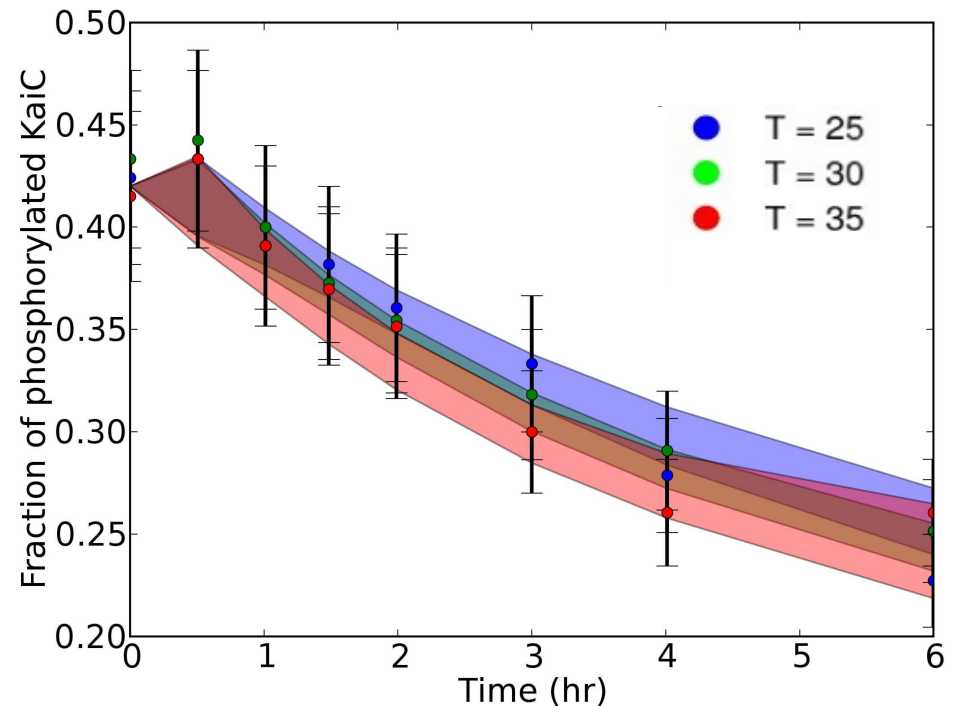
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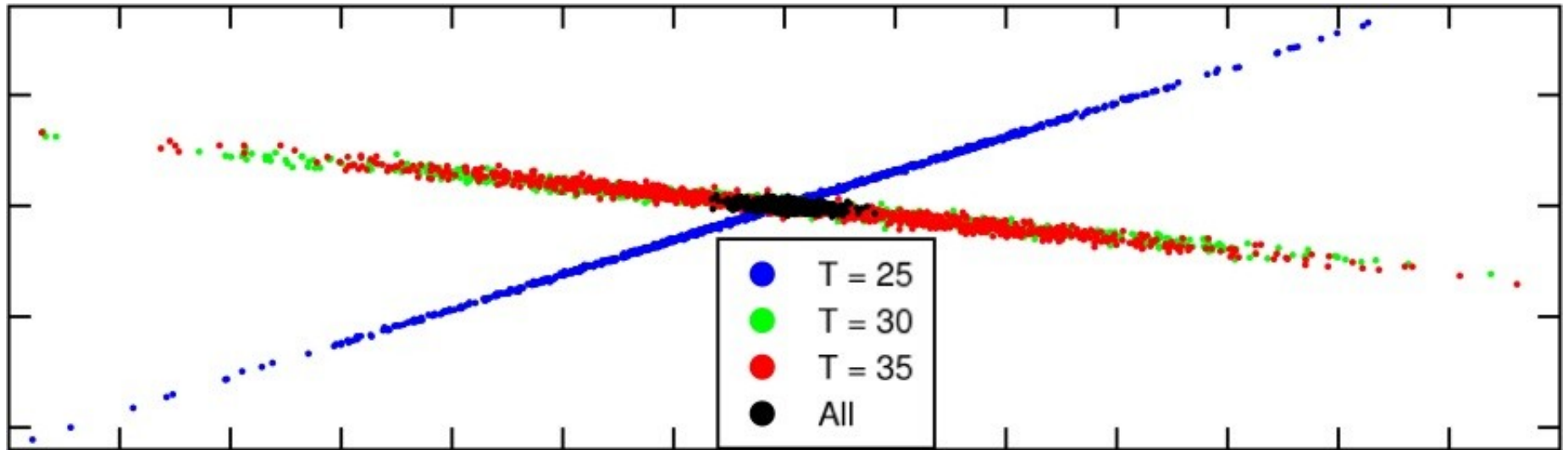


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# Is robustness a delicate balancing act?



# Robustness and Evolvability

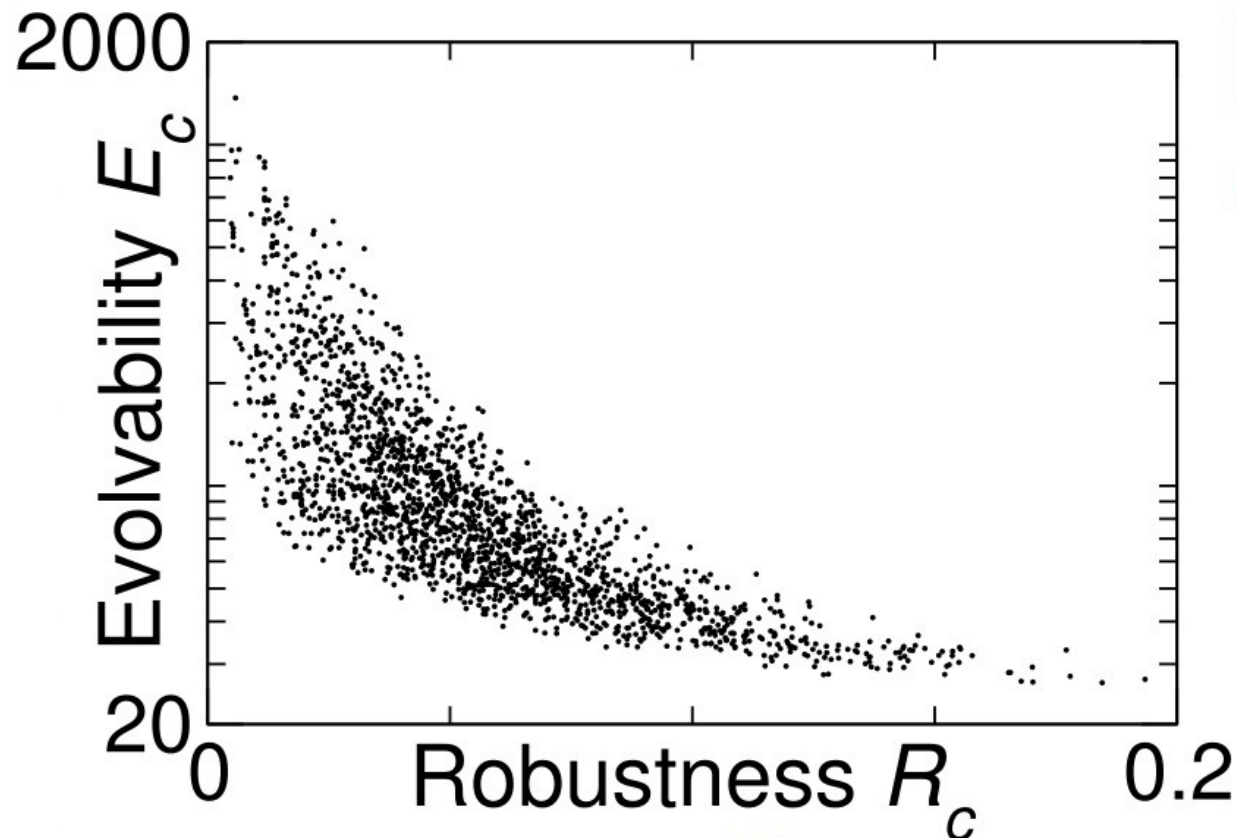


# Robustness and Evolvability

- › Robustness: What fraction of a given volume in parameter space keeps the output reasonably constant?
- › Evolvability: With a selection pressure to move in a certain direction in residual (output) space, how far can I move in that direction by varying my parameters by a fixed amount?

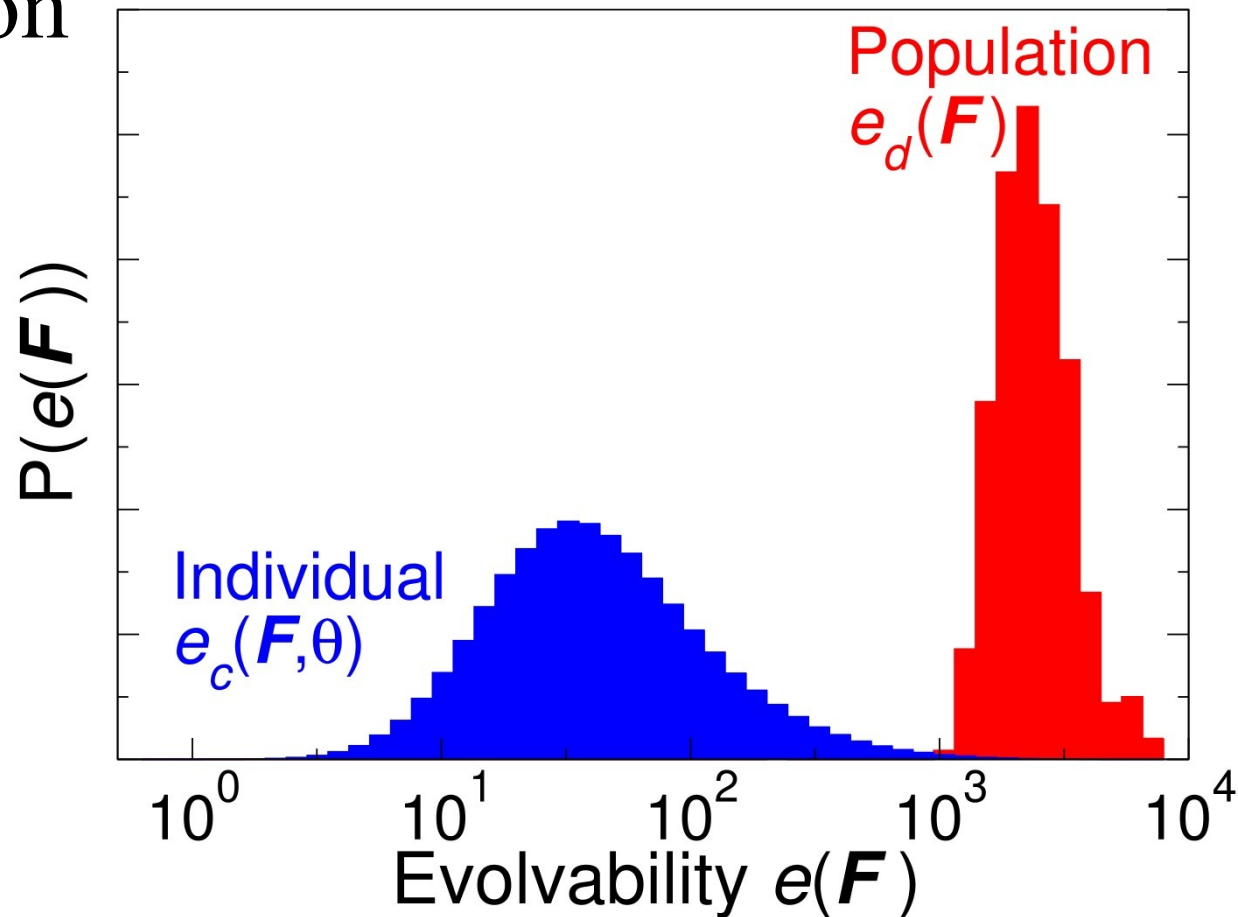
# Robustness and Evolvability

- › Individual evolvability decreases with robustness in example biological model



# Individual and Population Evolvability

- › Sloppiness may increase the variety of behaviors available to a population through mutation



# Future work?

- › Use information from multiple experiments / multiple systems to create ensembles
- › Network structure
  1. Can we vary uncertain network connections in a similar way as parameters?
  2. Can we predict which experimental data would best constrain the network structure?

# Conclusions

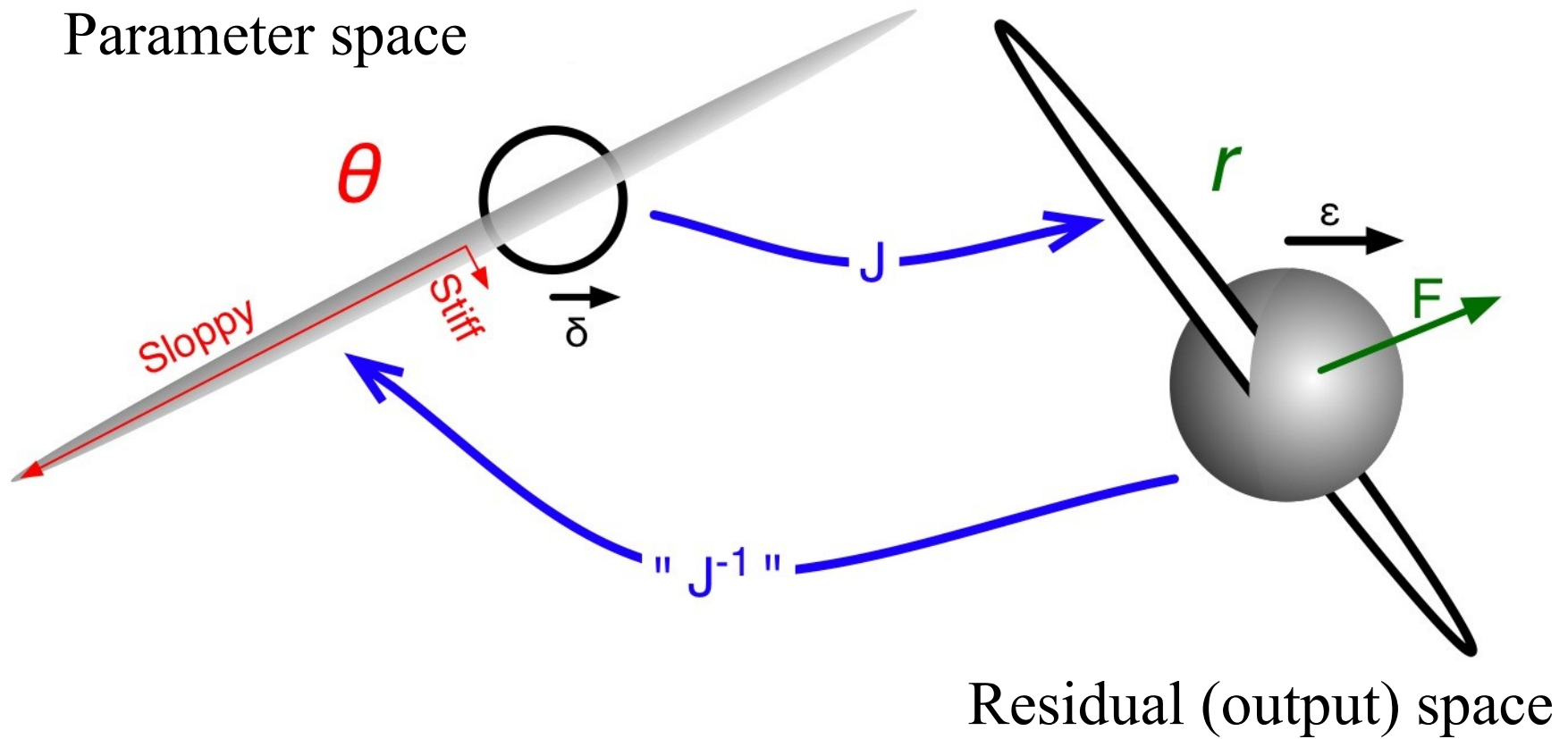
- › Varying parameters provides important information about model uncertainty.
- › Sloppiness is a common feature in large multiparameter models.
  1. Precise measurements of constants are not as important; instead optimize experiments to provide well-constrained predictions.
  2. Sloppiness can have important implications for robustness and evolvability.

# Thanks!

- › Jim Sethna
- › Ryan Gutenkunst
- › Chris Myers
- › YJ Chen
- › Ben Machta
- › Mark Transtrum

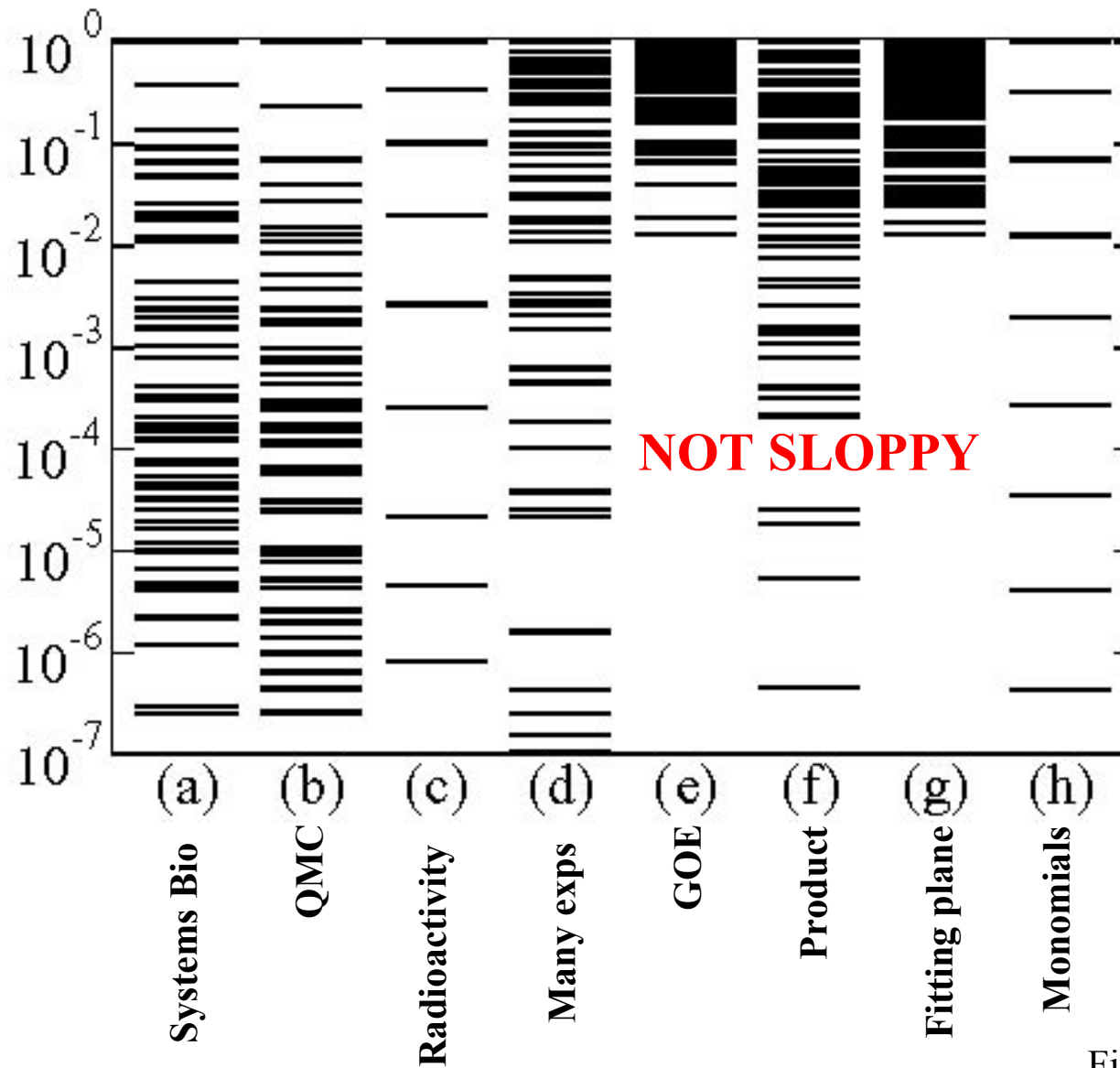


# Mapping parameters to residuals



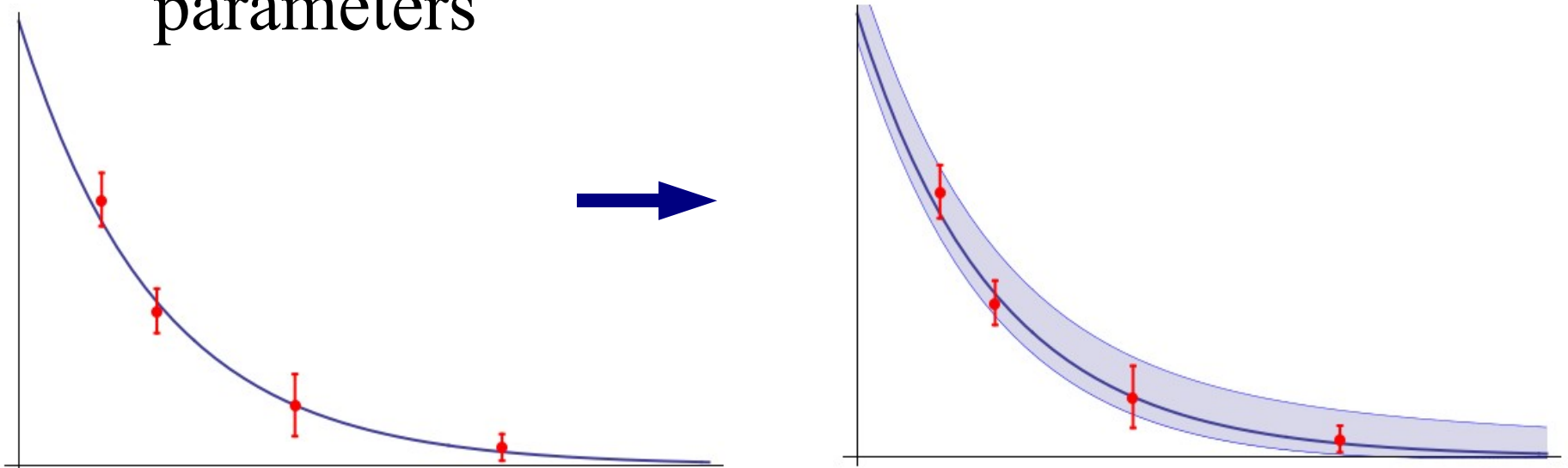


# Universality of Sloppiness



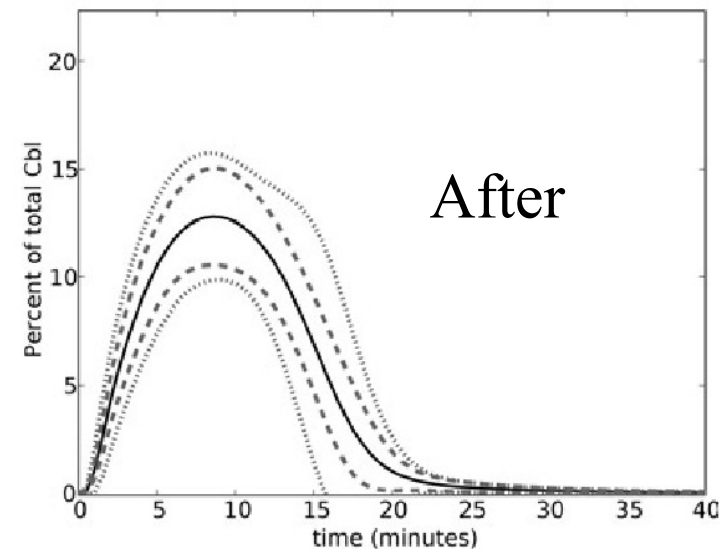
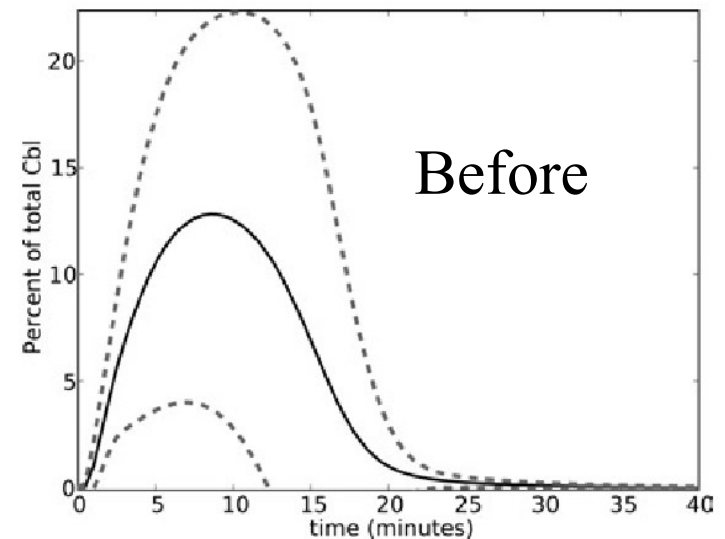
# Making Sensible Error Bars

- › Quickest method: Linear covariance analysis
- › Two approximations:
  1. Quadratic expansion around best-fit
  2. Linear response of output to changes in parameters



# Experimental Optimization

- › We can ask what new measurement will reduce the uncertainty of a specific output.
- › Example: Adding a single measurement of a different protein concentration.
- › Must use linear approx.



# Sloppiness

- › What does understanding sloppiness buy you? How can you use these ideas?
  1. More efficient Monte Carlo sampling of parameter space;
  2. Hints at the most important reactions in a network;
  3. An appreciation of the futility of thinking in terms of individual parameters;
  4. Model simplifications?

# More Efficient Sampling

- › Using a Metropolis Monte Carlo algorithm, we can make use of our knowledge about the local shape of the cost function.
- › Big steps in sloppy directions, small steps in stiff directions.

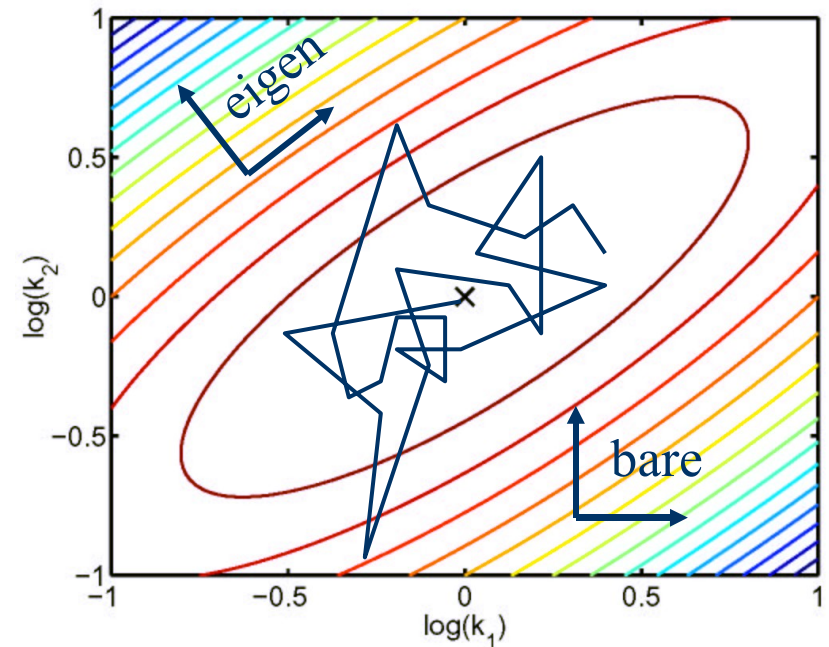
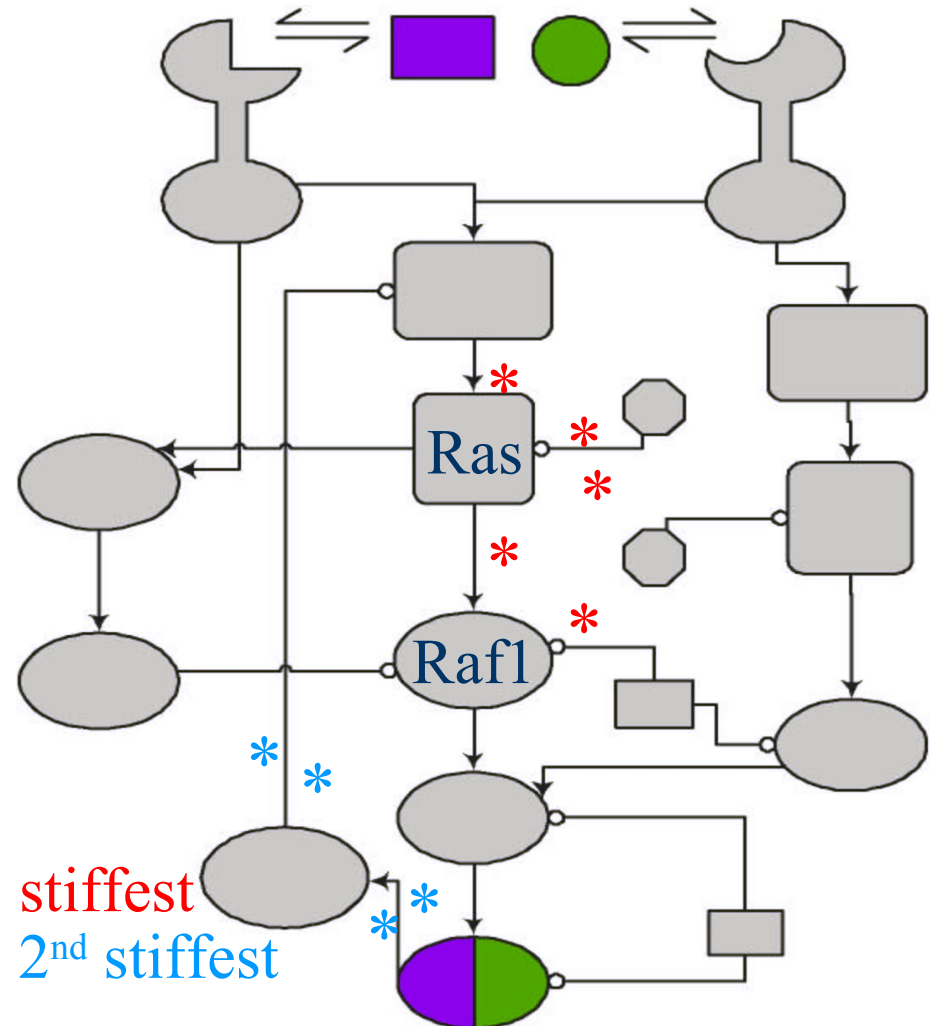


Figure courtesy Jim Sethna

# Stiffest Directions

- › The parameters with large components in the stiffest eigenvectors are in some sense more important.

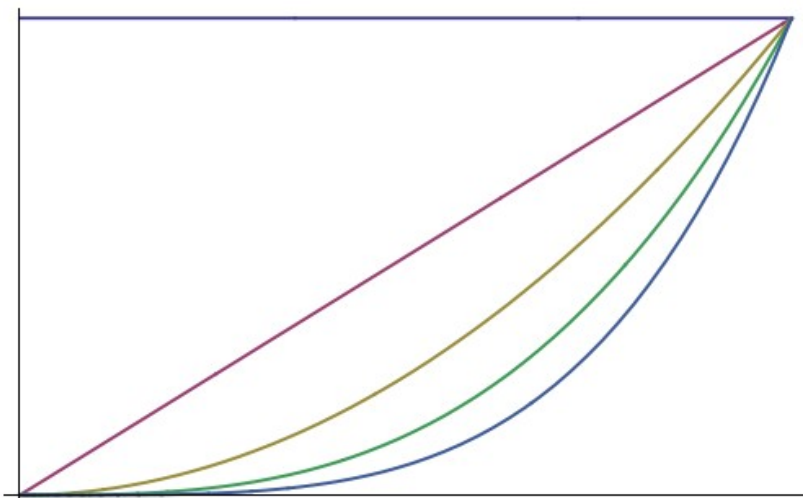


# Current Projects

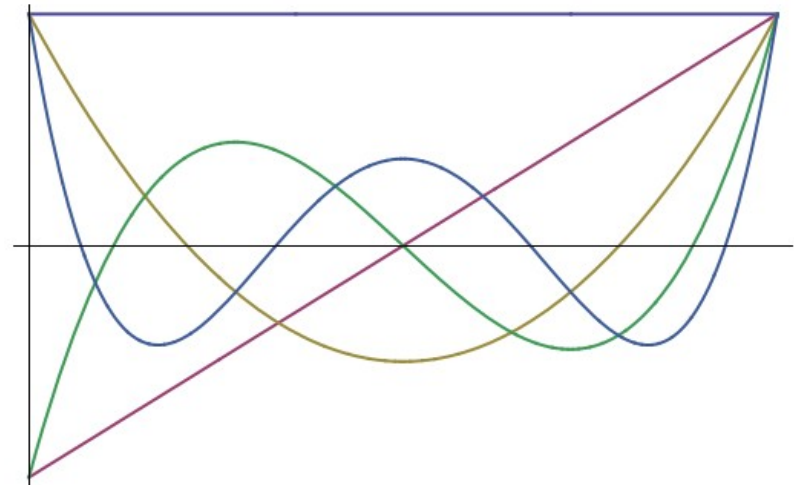
- › SloppyCell
- › Origin of sloppiness
- › Curved manifolds: connections to GR?
- › Model simplification?
- › New systems to implement

# Where does sloppiness come from?

- › Related to the interchangeability of different sets of parameters
- › Using the wrong parameterization
- › Example: fit polynomial function on  $[0,1]$



monomials: sloppy



Legendre polynomials: not sloppy



# Origin of Sloppiness

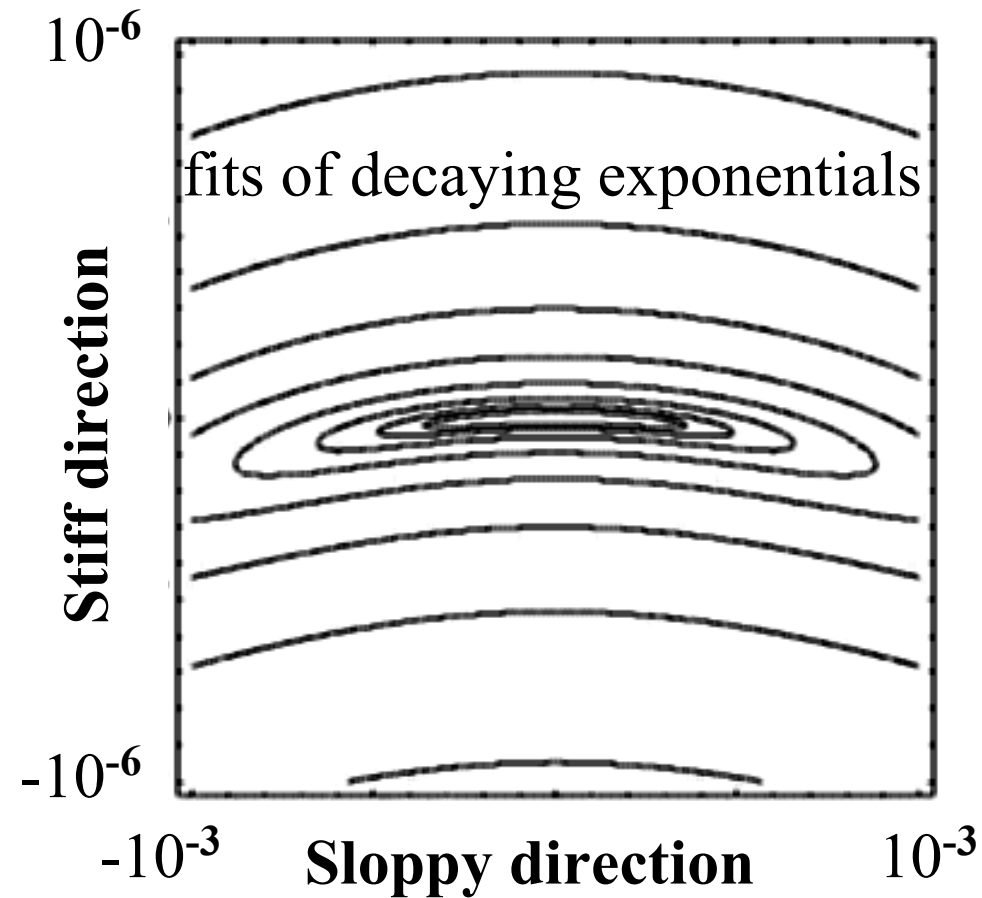
- › With two idealizations, get “perfect” sloppiness:
  1. Parameters are exactly interchangeable.
  2. Parameters are nearly degenerate.

$$H = V^T A^T A V \quad V = \begin{bmatrix} 1 & 1 & \cdots & 1 \\ \varepsilon_1 & \varepsilon_2 & \cdots & \varepsilon_N \\ \vdots & \vdots & \ddots & \vdots \\ \varepsilon_1^d & \varepsilon_2^d & \cdots & \varepsilon_N^d \end{bmatrix}$$

$$\det(V) = \prod_{i < j} (\varepsilon_i - \varepsilon_j) \propto \varepsilon^{N(N-1)/2}$$

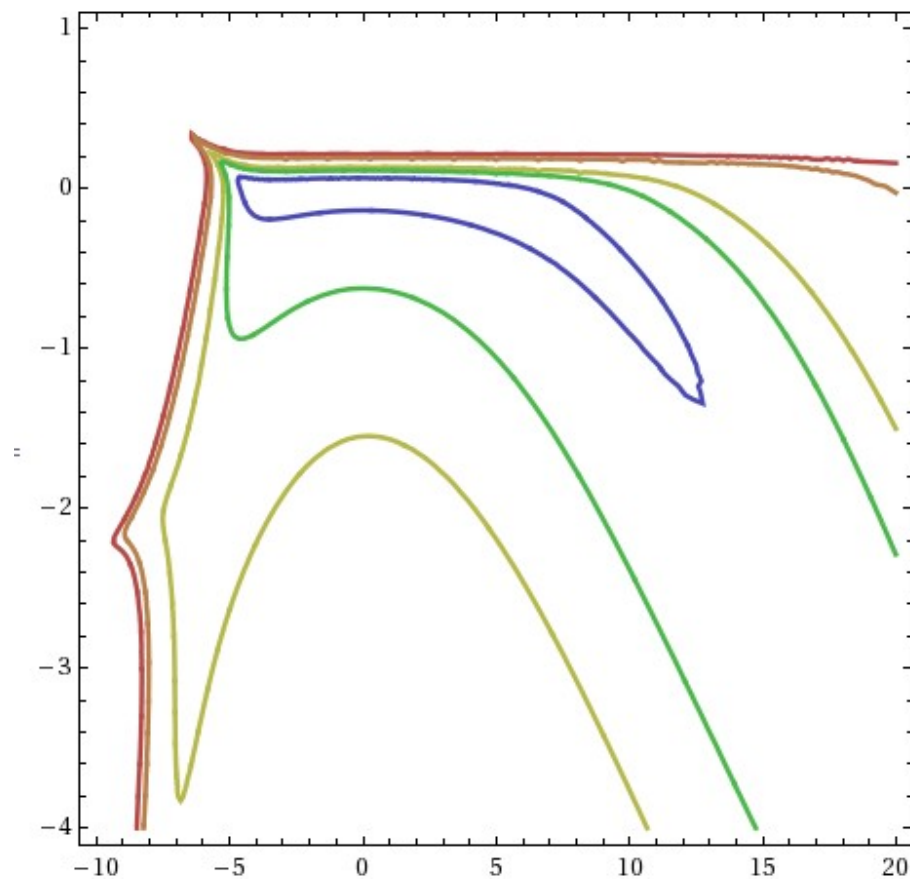
# Curved Manifolds

- › Anharmonic effects seem to be important.
- › To efficiently explore parameter space, we may need curved coordinates.



# Curved Manifolds

- › Example: sum of exponentials problem



# Model Simplification

- › Figure: Correlated parameter clusters
- › When sets of parameters have the same effect on output:
  1. We see sloppiness;
  2. It suggests we could simplify the model...

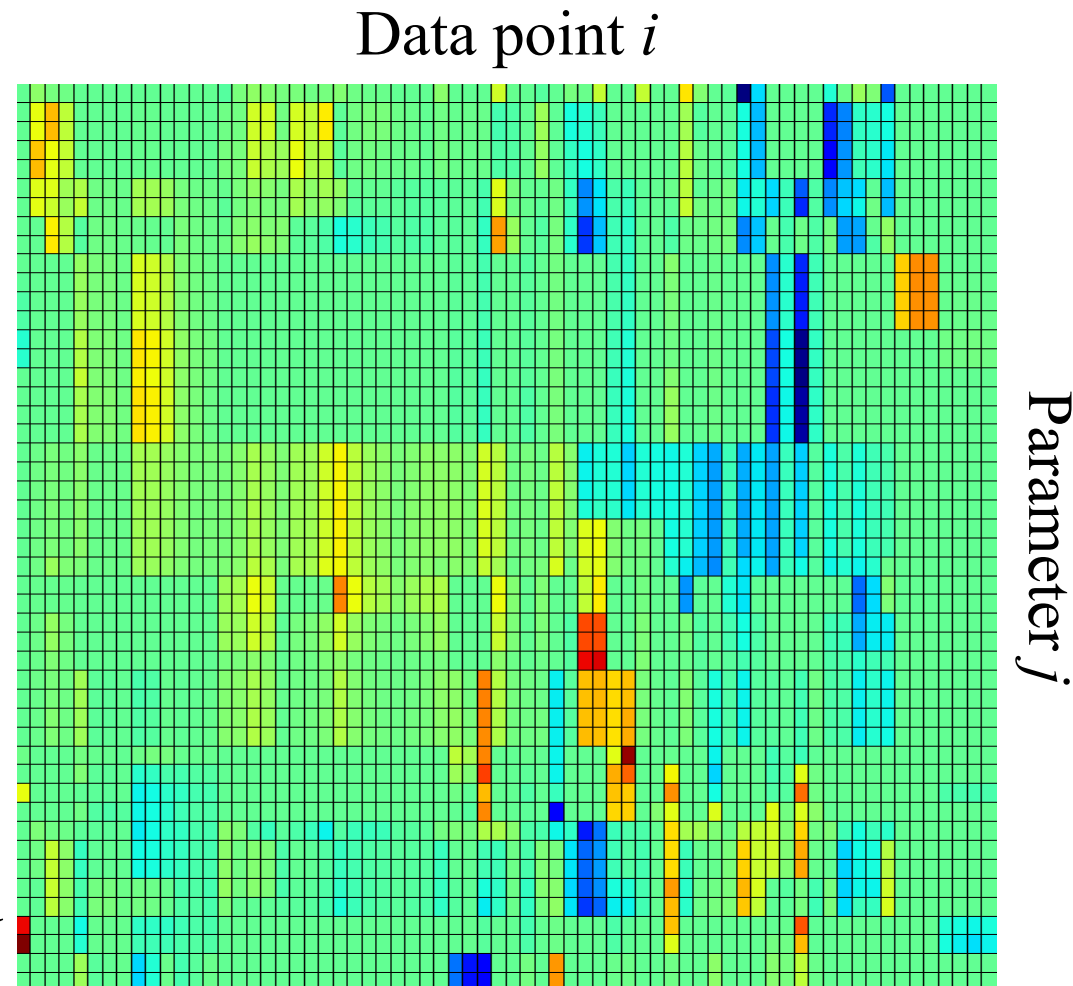


Figure courtesy Josh Waterfall/Jim Sethna

# New Systems to Implement

- › To avoid getting too abstract, we are on the lookout for real-world problems to implement...
  1. Climate modeling
  2. Economic models
  3. Physics models (CMB, accelerator design)
  4. Other systems biology problems

# Conclusions

- › Varying parameters provides important information about model uncertainty.
- › Sloppiness is a common feature in large multiparameter models.
  1. Precise measurements of constants are not as important; instead optimize experiments to provide well-constrained predictions.
  2. Simplification schemes may be fruitful.