Introduction to Numerical Methods for Identifiability

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How can we apply numerical methods to identifiability?



Consider the following approach:

- Assume your model is structurally identifiable and choose a set of "true" parameters
 - Generate simulated output data from these "true" parameters
- Attempt to fit your simulated data using a range of parameter values and solve for the "best" parameter set to reproduce the simulated data
- IF your original parameter set is returned, your model may be identifiable
 - What does it mean if the simulated data is without noise?
 - With noise?

How can I check my model using statistics?

- Begin by reframing the question:
 - Given output data **z**, what parameter set **p** generates this?
 - Consider "Parameter estimation" instead of "identifiability"
 - How are these concepts related?
 - What parameter or distribution set *most likely* generated the data?

Why would I want to explore numerical methods?

- Most numerical methods can explore both structural and practical identifiability
- Wide range of applicability to different models
- Relatively fast implementation
- Typically restricted to local identifiability, but global methods exist too

Maximum Likelihood Method

- Basic Idea: Reframe compartmental model as a statistical model where we assume the general form of the density function, but not parameter values
- Then if we knew the parameters, we could frame a probability: P(z | p)

data parameters

Likelihood Function: P(z | p) = f(z, p) = L(p | z)

- Now we rethink the probability distribution P, as a function of the data f
- We want to find the parameter set p that maximizes the likelihood L given data z

Example:

$$f(z \mid \mu, \sigma^2) = \frac{1}{\sqrt{2\pi}\sigma} \exp\left(-\frac{(z-\mu)^2}{2\sigma^2}\right) = L(\mu, \sigma^2 \mid z)$$

P(zlp)











Maximum Likelihood Method:

- Consistency: With sufficiently large number of observations, n, it is possible to find the value of p with arbitrary precision
- Normality: As n increases, the MLE tends to a Gaussian distribution with mean and covariance equal to the inverse of the Fisher information matrix
- Efficiency: Achieves Cramer-Rao bound as $n \rightarrow \infty$

Let's derive the likelihood function for the Gaussian example!

We have an ODE model defined as:

- Now sample data at times $t_1, t_2, t_3, \dots, t_n$
 - Data at t_i is defined as $z_i = (y(t_i) + e_i)$
 - Assume error e_i is Gaussian and unbiased, with known variance (2) known constant
- View data z_i as a sample from Gaussian distribution with mean $y(x, t_i; p)$ and variance σ^2
 - We assume all measures are independent (Is this realistic?)

So let's calculate the likelihood function:

• The Gaussian PDF: $f(z_i | \boldsymbol{\mu}, \sigma^2) = \frac{1}{\sqrt{2\pi}\sigma} \exp\left(-\frac{(z_i - \boldsymbol{\mu})^2}{2\sigma^2}\right)$ • Formatted for model: $f(z_i | \boldsymbol{y}(\boldsymbol{x}, \boldsymbol{t_i}; \boldsymbol{p}), \sigma^2) = \frac{1}{\sqrt{2\pi}\sigma} \exp\left(-\frac{(z_i - \boldsymbol{y}(\boldsymbol{x}, \boldsymbol{t_i}; \boldsymbol{p}))^2}{2\sigma^2}\right)$

 $Z_{i}^{*} = \begin{pmatrix} (t_{i}) + e_{i} \\ K \cdot N (u_{i}, \sigma^{2}) \end{pmatrix}$ Then our likelihood assuming independent observations: $L(y(t_i, p), \sigma^2 | z_1, z_2, \dots, z_n) = f(z_1, z_2, \dots, z_n | y(t_i, p), \sigma^2)$ $\mu_i \sigma^2$ $\lim_{t \to \infty} \{ \text{output lab} = \prod_{i=1}^n f(z_i | y(t_i, p), \sigma^2) \}$ $= \left(\frac{1}{2\pi\sigma^{2}}\right)^{n/2} \exp\left(-\frac{\sum_{i=1}^{n}(z_{i}-y(t_{i};p))^{2}}{2\sigma^{2}}\right)$

maximizing likelihood

What does this look like in practice?

- Rather than maximizing the likelihood, in practice we minimize the negative log likelihood
 - Log is well behaved and minimization algorithms are common

$$-LL = -\ln\left(\left(\frac{1}{2\pi\sigma^{2}}\right)^{n/2} \exp\left(-\frac{\sum_{i=1}^{n}(z_{i}-y(t_{i};p))^{2}}{2\sigma^{2}}\right)\right)$$
$$= -\left(-\frac{n}{2}\ln(2\pi) - n\ln(\sigma) - \frac{\sum_{i=1}^{n}(z_{i}-y(t_{i};p))^{2}}{2\sigma^{2}}\right)$$
$$= \frac{n}{2}\ln(2\pi) + n\ln(\sigma) + \frac{\sum_{i=1}^{n}(z_{i}-y(t_{i};p))^{2}}{2\sigma^{2}}$$

... Then our problem reduces to:

$$\min_{p}(-LL) = \min_{p} \sum_{i=1}^{n} (z_i - y(t_i; p))^2$$

So what's the big idea?

- We will "profile" one parameter at a time using the likelihood function
 - Fix the parameter to a range of values and fitting all other parameters in the model
 - Calculate the likelihood value for every combination
- This will give the best fit at each point
- Plot the best likelihood values for each value of p_i

... This is known as the Profile Likelihood method!



What can we expect to see?



5 Minute Break!



Time to Code!...

