

Dealing with Data Gradients: “Backing Out” & Calibration

Nathaniel Osgood

CMPT 858

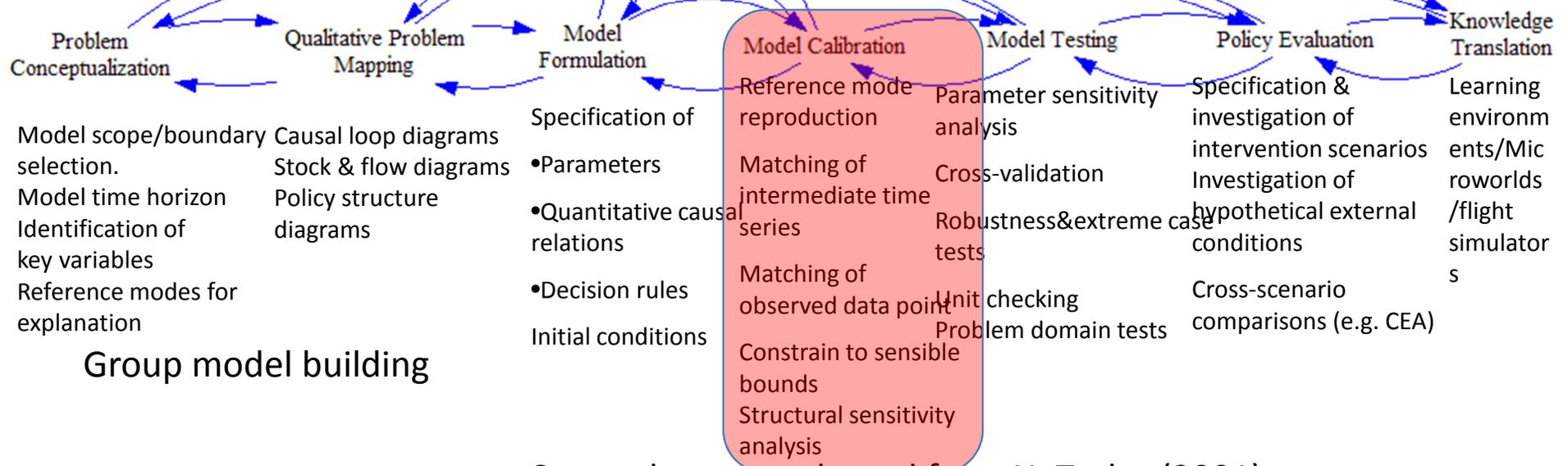
Term Project Updated Due Date

- Because of Holiday weekend, date is now midnight April 25

A Key Deliverable!



Mental Model



Group model building

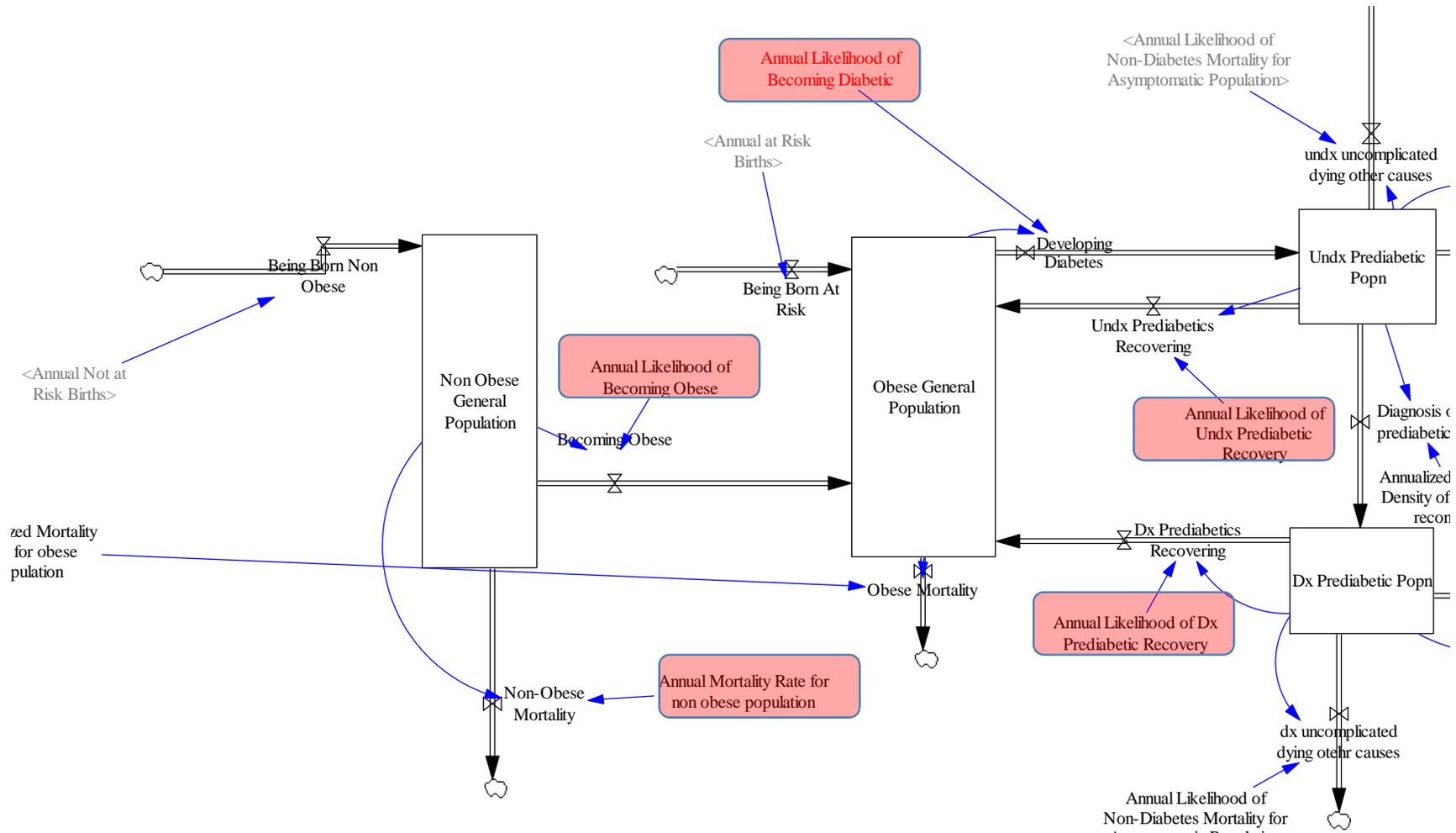
Some elements adapted from H. Taylor (2001)

Sources for Parameter Estimates

- Surveillance data
- Controlled trials
- Outbreak data
- Clinical reports data
- Intervention outcomes studies
- Calibration to historic data
- Expert judgement
- Metaanalyses

Parameter*	Description	Baseline value (units)	Reference
μ	Entry/exit of sexual activity	0.0056 (years ⁻¹)	Garnett and Bowden, 2000
c	Partner change rate per Susceptible	16.08 (years ⁻¹)	Approximated from Garnett and Bowden, 2000
β	Probability of infection per sexual contact	0.70	Garnett and Bowden, 2000
ϕ	Fraction of Infectives who are symptomatic	0.20	Garnett and Bowden, 2000
$1/\gamma$	Latent period	0.038 (years)	Brunham et. al., 2005
$1/\sigma$	Duration of infection	0.25 (years)	Brunham et. al., 2005
θ	Asymptomatic recovery coefficient	1.5	Garnett and Bowden, 2000
$1/\pi$	Duration of naturally-acquired immunity	1 (year)	Approximated from Brunham et. al., 2005

Introduction of Parameter Estimates



Sensitivity Analyses

- Same relative or absolute uncertainty in different parameters may have hugely different effect on outcomes or decisions
- Help identify parameters that strongly affect
 - Key model results
 - Choice between policies
- We place more emphasis in parameter estimation into parameters exhibiting high sensitivity

Dealing with Data Gradients

- Often we don't have reliable information on *some* parameters, but do have other data
 - Some parameters may not be observable, but some closely related observable data is available
 - Sometimes the data doesn't have the detailed breakdown needed to specifically address one parameter
 - Available data could specify sum of a bunch of flows or stocks
 - Available data could specify some function of several quantities in the model (e.g. prevalence)
- Some parameters may implicitly capture a large set of factors not explicitly represented in model
- There are two big ways of dealing with this: manually “backing out”, and automated calibration

“Backing Out”

- Sometimes we can manually take several aggregate pieces of data, and use them to collectively figure out what more detailed data might be
- Frequently this process involves imposing some (sometimes quite strong) assumptions
 - Combining data from different epidemiological contexts (national data used for provincial study)
 - Equilibrium assumptions (e.g. assumes stock is in equilibrium. Cf deriving prevalence from incidence)
 - Independence of factors (e.g. two different risk factors convey independent risks)

Example

- Suppose we seek to find out the sex-specific prevalence of diabetes in some population
- Suppose we know from published sources
 - The breakdown of the population by sex (c_M, c_F)
 - The population-wide prevalence of diabetes (p_T)
 - The prevalence rate ratio of diabetes in women when compared to men (rr_F)
- We can “back out” the sex-specific prevalence from these aggregate data (p_F, p_M)
- Here we can do this “backing out” without imposing assumptions

Backing Out

male diabetics + # female diabetics = # diabetics

$$(p_M * c_M) + (p_F * c_F) = p_T * (c_M + c_F)$$

- Further, we know that $p_F / p_M = rr_F \Rightarrow p_F = p_M * rr_F$

- Thus

$$(p_M * c_M) + ((p_M * rr_F) * c_F) = p_T * (c_M + c_F)$$

$$p_M * (c_M + rr_F * c_F) = p_T * (c_M + c_F)$$

- Thus

- $p_M = p_T * (c_M + c_F) / (c_M + rr_F * c_F)$

- $p_F = p_M * rr_F = rr_F * p_T * (c_M + c_F) / (c_M + rr_F * c_F)$

Disadvantages of “Backing Out”

- Backing out often involves questionable assumptions (independence, equilibrium, etc.)
- Sometimes a model is complex, with several related known pieces
 - Even though we may know a lot of pieces of information, it would be extremely complex (or involve too many assumptions) to try to back out several pieces simultaneously

Another Example: Joint & Marginal Prevalence

	Rural	Urban	
Male	p_{MR}	p_{MU}	p_M
Female	p_{FR}	p_{FU}	p_F
	p_R	p_U	

Perhaps we know

- The count of people in each { Sex, Geographic } category
- The marginal prevalences (p_R, p_U, p_M, p_F)

We need at least one more constraint

- One possibility: assume $p_{MR} / p_{MU} = p_R / p_U$

We can then derive the prevalences in each { Sex, Geographic } category

Calibration: “Triangulating” from Diverse Data Sources

- Calibration involves “tuning” values of less well known parameters to best match observed data
 - Often try to match against *many* time series or pieces of data at once
 - Idea is trying to get the software to answer the question: “What must these (less known) parameters be in order to explain all these different sources of data I see”
- Observed data can correspond to complex combination of model variables, and exhibit “emergence”
- Frequently we learn from this that our model structure just can’t produce the patterns!

Calibration

- Calibration helps us find a reasonable (specifics for) “dynamic hypothesis” that explains the observed data
 - Not necessarily the truth, but probably a reasonably good guess – at the least, a consistent guess
- Calibration helps us leverage the large amounts of diffuse information we may have at our disposal, but which cannot be used to directly parameterize the model
- Calibration helps us falsify models

Calibration: A Bit of the How

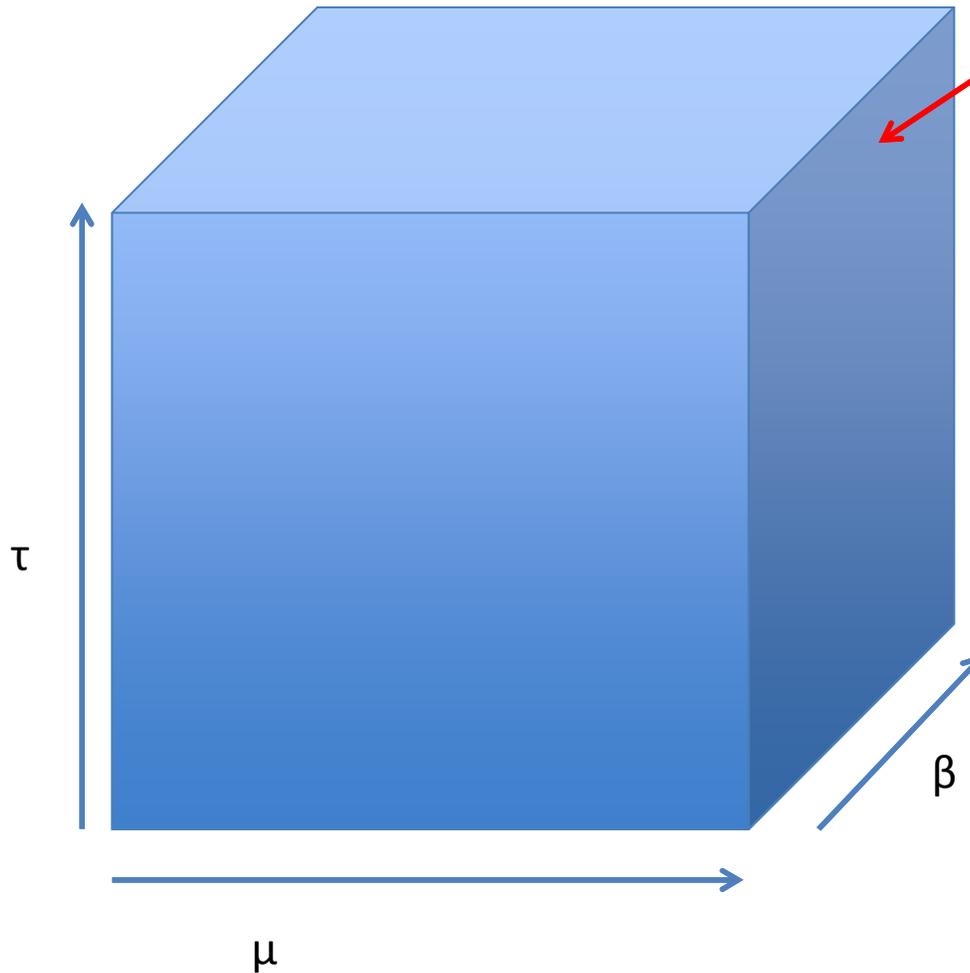
- Calibration uses a (global) optimization algorithm to try to adjust unknown parameters so that it automatically matches an arbitrarily large set of data
- The data (often in the form of time series) forms constraints on the calibration
- The optimization algorithm will run the model many (minimally, thousands, typically 100K or more) times to find the “best” match for all of the data

Required Information for Calibration

- Specification of what to match (and how much to care about each attempted match)
 - Involves an “error function” (“penalty function”, “energy function”) that specifies “how far off we are” for a given run (how good the fit is)
 - Alternative: specify “payoff function” (“objective function”)
- A statement of what parameters to vary, and over what range to vary them (the “parameter space”)
- Characteristics of desired tuning algorithm
 - Single starting point of search?

Envisioning “Parameter Space”

For each point in this space, there will be a certain “goodness of fit” of the model to the collective data



Assessing Model “Goodness of Fit”

- To improve the “goodness of fit” of the model to observed data, we need to provide some way of quantifying it!
- Within the model, we
 - For each historic data, calculate discrepancy of model
 - Figure out absolute value of discrepancy from comparing
 - Historic Data
 - The model’s calculations
 - Convert the above to a fractional value (dividing by historic data)
 - Sum up these discrepancy

Characteristics of a Desirable Discrepancy Metric

- **Dimensionless:** We wish to be able to add discrepancies together, regardless of the domain of origin of the data
- **Weighted:** Reflecting different pedigrees of data, we'd like to be able to weigh some matches more highly than others
- **Analytic:** We should be able to differentiate the function one or more times
- **Concave:** Two small discrepancies of size a should be considered more desirable than having one big discrepancy of size $2a$ for one, and no discrepancy at all for the other.
- **Symmetric:** Being off by a factor of two should have the same weight regardless of whether we are $2x$ or $\frac{1}{2}x$
- **Non-negative:** No discrepancy should cancel out others!
- **Finite:** Finite inputs should yield infinite discrepancies

A Good Discrepancy Function (Assuming non-negative h & m)

Exponent
>1 \Rightarrow concave with respect to h-m

Taking average in denominator (together w/squaring of result) ensures symmetry with respect to h&m

$$w \cdot \left(\frac{h - m}{\text{average}(h, m)} \right)^2 = w \cdot \left(\frac{h - m}{\left(\frac{h + m}{2} \right)} \right)^2$$

Division \Rightarrow Dimensionless
(Judging by *proportional* error, not absolute)

Only zero if h=m=0.

Denominator is only very small if numerator is as well!

Considerations for Weighting

- **Purpose of model:** If we “care” more about a match with respect to some variables, we can more heavily weight matches for those variables
- **Uncertainty in estimate:** The more uncertain the estimate of the quantity, the lower the weight
- **Whether data exists:** no data => weight should be zero