Parameter Uncertainty

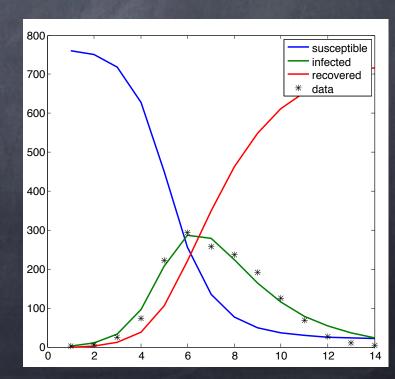
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 methods for inferring key quantities of interest



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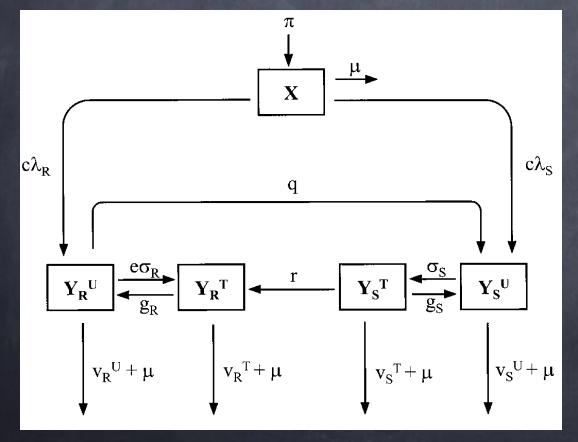
 Complex models have many parameters about which we have little information

Motivation

In 2000, ~30% of gay men in San Francisco were infected with HIV, 50% of whom were taking combination antiretroviral therapy (ART)

- ART effective at reducing AIDS death rate in San Francisco, but does not completely eliminate infectivity
- unclear whether net effect of increased distribution of ART would be to increase or decrease incidence of HIV
- Blower et al. introduced following model (Blower, S.M., et al. 2000. A tale of two futres: HIV and antiretroviral therapy in San Francisco.Science 287:650-654.)

Blower et al. (2000) model

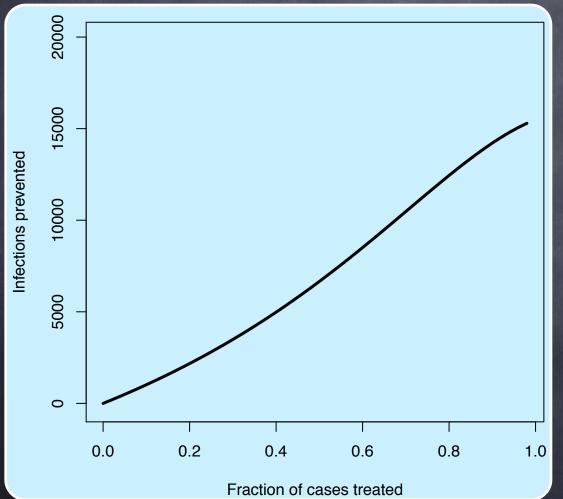


X -- susceptible Y_R^U -- untreated, infected with drugresistant strain etc

Lots of model parameters

- π -- rate at which gay men join sexually active community
- \circ 1/ μ -- average time during which new partners are acquired
- o c -- average number of new sex partners per year
- p -- probability of a drug-resistant case (relative to a drug-sensitive case) transmitting drug-sensitive viruses
- 1/q -- average time for an untreated drug-resistant infection to revert to a drugsensitive infection
- σ -- per capita effective treatment rate
- e -- relative efficacy of ART in treating drug-resistant infections
- o r -- rate of emergence of resistance due to acquired resistance
- g -- proportion of cases that give up ART per year
- v -- average disease progression rate

Model predictions



•ART could prevent ~15,000 cases of 20 years

How reliable is this?
Model has 20 parameters
Few (if any) known exactly

- To know <u>robustness</u> of model predictions, require a way of exploring output of a family of parameterized models
- If number of unknown parameters is bigger than, say,
 2 then systematic grid search would be computationally intractable
- Qualitatively investigate variability in model output that is generated from uncertainty in parameter inputs
- Perform multiple model evaluations using randomly chosen values for parameters

hypothetical, but large range with minimum and maximum is for the parameters). If biological knowledge exists esting a more frequent or expected value for a parameter, a hal pdf would be the best choice (setting the variance of the ibution as large as needed).

Latin Hypercube Sampling - LHS Uniform and Normal pdfs Sample size N=5

AL DEALERS

bmax

03

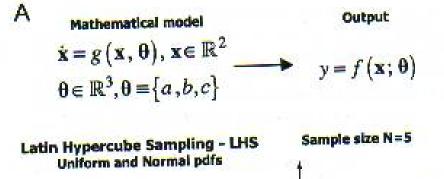
S. Marino et al. / Journal of Theoretical Biology 254 (2008) 178–196

ach parameter, sampling is guided by the specification probability density function (pdf) (i.e. normal, uniform, armal, etc.), depending on *a priori* information. If there are no pri data, a natural choice is a uniform distribution (assigning hypothetical, but large range with minimum and maximum s for the parameters). If biological knowledge exists esting a more frequent or expected value for a parameter, a hal pdf would be the best choice (setting the variance of the ibution as large as needed).

everal sampling strategies can be implemented to perform such as random sampling, importance sampling, or LHS con and Davis, 2003; Mckay et al., 1979). To recreate input or distributions through sampling, a large number of samples likely required. If too few iterations are performed, not all es may be represented in the samples or values in the outer es may be under-sampled. The LHS algorithm was specifically cloped to address this problem and it is by far the most ular sampling scheme for UP. (Norcis, 2000).

Latin hypercube sampling-LHS

us belonge to the MC class of sampling methods, and was



$$a - Unif(a_{\min}, a_{\max}) = a_{\max} - a_{\min} = a_4 = a_1 = a_5 = a_3$$

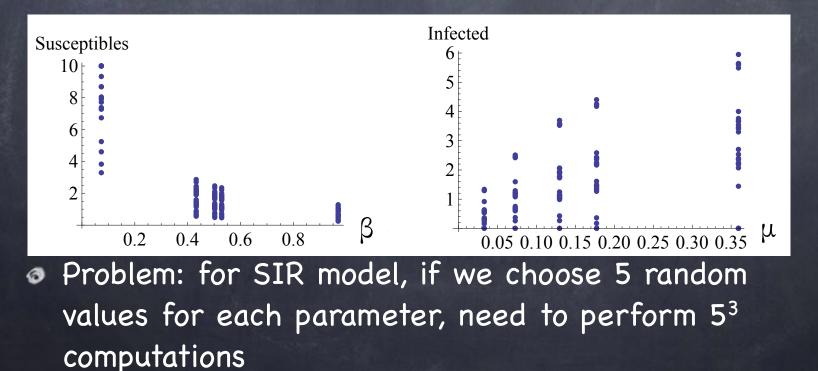
(I min

$$b \sim Unif(b_{\min}, b_{\max})$$

 b_{min}

Monte Carlo analysis

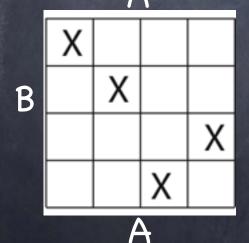
 Using parameter combinations determined by Monte Carlo simulation, examine scatter plots of output against each parameter



Assume 2 unknown parameters

B X X X X

In **random sampling**, new sample points are generated without taking into account previously generated sample points eg, for i=1:10 A(i) = rand*(A_{max} - A_{min}) + A_{min}; B(i) = rand*(B_{max} - B_{min}) + B_{min}; end

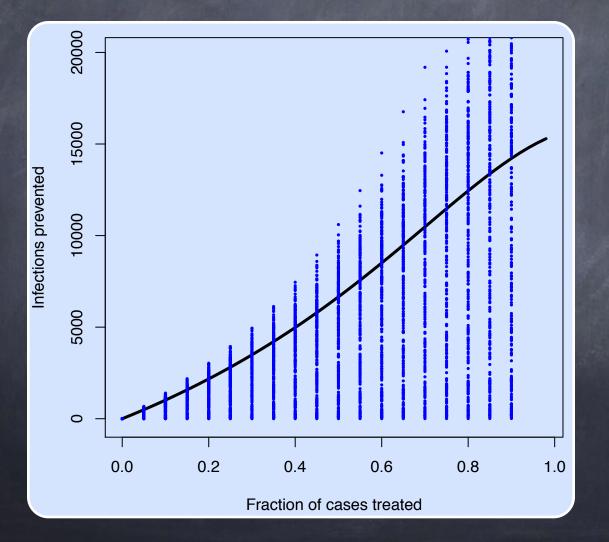


In Latin Hypercube sampling one must first decide how many sample points to use and for each sample point remember in which row and column sample point was taken

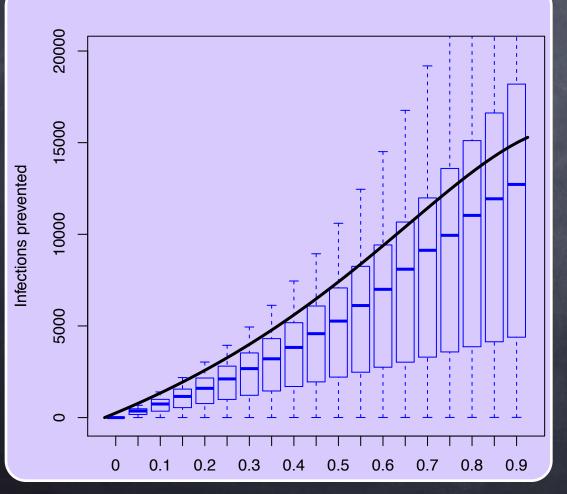
Programs available for use in R & Matlab

Typically, LHS random numbers in unit interval (0,1) Need to `stretch': A = LHS_rand*(A_{max}-A_{min})+A_{min}

Results of LHS Specified ranges of 18 parameters



Box-Whisker plot

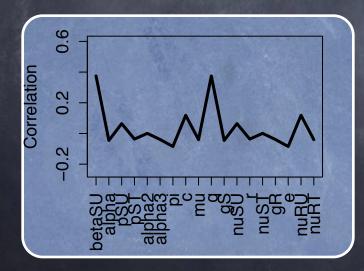


Evidently, our best guesses at parameter values are somewhat optimistic

At least ART is not found to be counter-productive in this respect \rightarrow an open question at time of this study

Which parameters important?

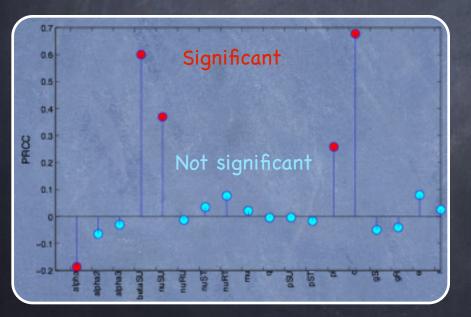
Can explore correlation between vectors of parameters studied and outcome of interest (in this case, # cases prevented)



Can further use partial rank correlation to establish sensitivity of conclusions to <u>specific parameters</u>

Which parameters important?

BUT, linear correlation ignores fact that model output for each value of a parameter simultaneously includes changes in other



Can use partial rank correlation to establish sensitivity of conclusions to specific parameters

Summary

 Important to distinguish between two sources of error in model predictions

I. Variability: arises from stochasticity in process and measurement

> solution is to explore many model realizations

II. Uncertainty: results from absence of information on parameters/processes

solution is (efficient) sensitivity analysis