### Day 4

Centre for the Mathematical Modelling of Infectious Diseases London School of Hygiene & Tropical Medicine



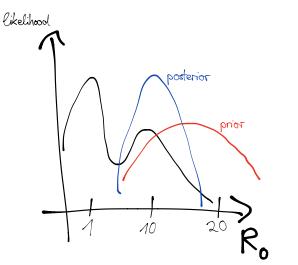


centre for the mathematical modelling of infectious diseases

## The point of all of this

posterior probabilities

$$p(\theta|\text{data}) = \frac{p(\text{data}|\theta)p(\theta)}{p(\text{data})}$$



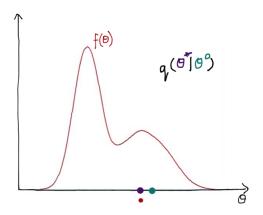
We interpret  $p(\theta|\text{data})$  as the probability distribution of a random variable  $\theta$ , from which we *sample* (via MCMC)

#### Why sample?

- 1. explore parameter space
- 2. samples can be useful
  - explore interventions, forecasts

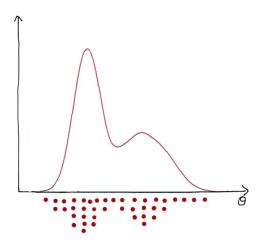
### MCMC: Sampling from a distribution

• We can calculate (in a deterministic model)  $p(\theta|\text{data})$  given any  $\theta$  – how do we sample?



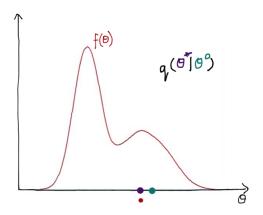
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#### Stochastic models

- one  $\theta$  can lead to many possible outcomes X
- we can
  - 1. sample from  $p(x|\theta)$  (via simulation)
  - 2. evaluate the trajectory likelihood  $p(\text{data}|x,\theta)$
- we can't directly evaluate the likelihood  $p(\text{data}|\theta)$

$$p(\text{data}|\theta) = \sum_{x} p(\text{data}|x, \theta) p(x|\theta)$$

▶ The number of possible trajectories X for one value of  $\theta$  is large, potentially infinite

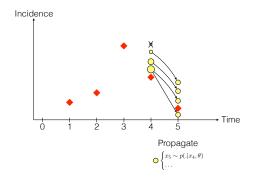
#### Sequential Monte Carlo (SMC) / Particle Filter I

To approximate

$$p(\mathrm{data}|\theta) = \sum_{x} p(\mathrm{data}|x,\theta) p(x|\theta)$$

we sample n trajectories  $x_n$  from

$$p(y_{1:(T-1)}|x,\theta)p(x|\theta)$$



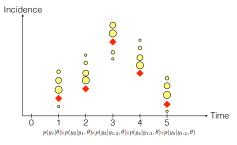
#### Sequential Monte Carlo (SMC) / Particle Filter II We then calculate

$$p(y_T|y_{1:(T-1)}|x_n,\theta)$$

for each of the particles. The sum of these values is

$$\sum_{T_{-}} p(y_{1:T}|x_n,\theta) p(x_n|\theta)$$

which is a sample estimate of the likelihood.



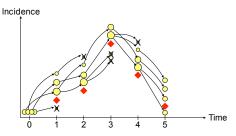
$$\label{eq:log-likelihood:log} \textit{Log-Likelihood:} \log\{p(y_{1:T}|\theta)\} = \sum_{x} \log\{p(y_{t}|y_{1:t-1},\theta)\}$$

#### Sequential Monte Carlo (SMC) / Particle Filter III

We can also retrieve filtered trajectories, that is samples from

$$p(x|\text{data})$$

by following the particles from the last point backwards.



#### pMCMC

- Once we can estimate  $p(\text{data}|\theta)$ , we can combine this with the prior to evaluate the posterior  $p(\theta|\text{data})$  for any  $\theta$ .
- ▶ We can then use MCMC to sample from this -> pMCMC

**MCMC** 

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$\square$	SMC	РМСМС	SMC <sup>2</sup>

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the	МСМС	SMC	
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