### Day 4

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



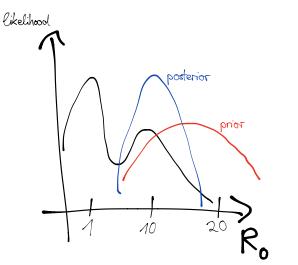


centre for the mathematical modelling of infectious diseases

### The point of all of this

posterior probabilities

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



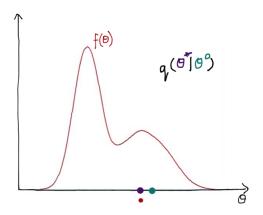
We interpret  $p(\theta|\text{data})$  as the probability distribution of a random variable  $\theta$ , from which we *sample* (via MCMC)

#### Why sample?

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

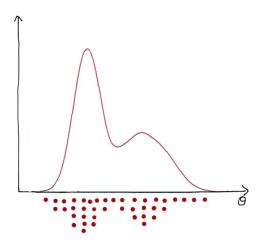
#### MCMC: Sampling from a distribution

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



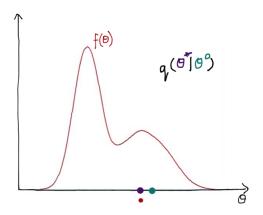
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#### Stochastic models

- 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  $p(\text{data}|\theta)$

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

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

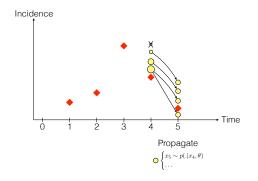
### Sequential Monte Carlo (SMC) / Particle Filter I

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

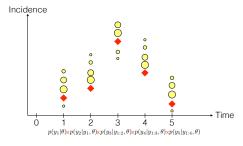
We then calculate

$$p(Y_T|X_n,\theta)$$

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

$$\sum_{X_n} p(Y_{1:T}|X_n,\theta) p(X_n|\theta)$$

which is a sample estimate of the likelihood.



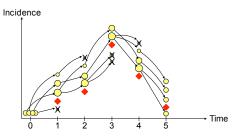
$$\label{eq:log-likelihood} \begin{aligned} & \log\{p(y_{1:T}|\theta)\} = \sum_{T} \log\{p(y_{t}|y_{1:t-1},\theta)\} \end{aligned}$$

#### 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** 

Stimating the likelihood NWC NWC NWC

	Sampling normane posterior		
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$\square$	SMC	РМСМС	SMC <sup>2</sup>

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the	МСМС	SMC	
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