

# **(Particle) Markov chain Monte Carlo**

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Once the model is fitted and the model fit assessed, we can use the model / parameter estimates in various ways:

- **Inference:** interpreting the parameter estimates.
- **Prediction:** to predict what might happen if the outbreak were to occur under the same conditions again.
- **Forecasting:** to predict what might happen in the future, based on data available now.

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- **Surveillance:** e.g. under-reporting, imperfect coverage, imperfect diagnosis, mis-diagnosis;
- **Rounding error:** e.g. data often collated daily / weekly;
- **Hidden states:** some epidemiological processes never observed (e.g. you might know *roughly* when you started feeling sick with flu, but not when you were infected or when you became infectious).

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Dealing with these challenges is **hard!** (But we will have a go!)

To deal with the **partially observed** data, we can introduce a set of **latent** variables,  $\mathbf{x} = (\mathbf{t}, \delta)$ , where  $\mathbf{t}$  is a vector of **hidden** event *times*, and  $\delta$  is a vector of **hidden** event *types*.

Then the **likelihood** can be expressed as:

$$f(\mathbf{y} \mid \theta) = \int_{\mathbf{x}} f(\mathbf{y} \mid \mathbf{x}, \theta) f(\mathbf{x} \mid \theta) d\mathbf{x},$$

where

- $f(\mathbf{y} \mid \mathbf{x}, \theta)$  is an **observation** process (or **measurement error** / **model discrepancy**);
- $f(\mathbf{x} \mid \theta)$  is the **likelihood function** based on the **latent** variables  $\mathbf{x}$ .

$$f(\mathbf{y} \mid \theta) = \int_{\mathbf{x}} f(\mathbf{y} \mid \mathbf{x}, \theta) f(\mathbf{x} \mid \theta) d\mathbf{x},$$

This **marginalises** (*averages*) across the hidden variables  $\mathbf{x}$ .

This is a complex integral, over all possible combinations of events, and all possible event times consistent with the data.

It may also be the case that the **number** of hidden events is **unknown**, in which case we have to repeat the integration for every possible number of hidden events.

One approach is therefore to include the **hidden** variables  $\mathbf{x}$  as **additional parameters** in the model.

We can then estimate the **joint posterior** distribution for  $(\theta, \mathbf{x})$ , and then derive the **marginals** for the parameters of interest  $(\theta)$  *numerically*.

This is usually done using MCMC methods; an approach known as **data-augmented MCMC** (e.g. Gibson and Renshaw [1998](#); O'Neill and Roberts [1999](#); Jewell et al. [2009](#)).

It is very powerful, but difficult to code, scale and optimise.



Alternatively, we can build inference algorithms around **simulating** directly from the model-of-interest, and then searching for parameter sets that are more consistent with the **observed data**.

These **simulation-based methods** are also powerful and flexible:

- Don't have to store all of the latent variables (so memory requirements are lower).
- Are often straightforward to parallelise.
- Simulation can often be easier than calculating the likelihood.
- Implementation often easier than DA (e.g. "plug-and-play")

However, there are also practical difficulties:

- The probability of matching the data exactly (i.e. getting a non-zero likelihood) is often very low.
- Often require some form of approximation to obtain a match.

Examples of latent variable methods:

- **Data-augmented MCMC** (e.g. Gibson and Renshaw [1998](#); O'Neill and Roberts [1999](#); S. Cauchemez and Ferguson [2008](#); Jewell et al. [2009](#))
- **Sequential Monte Carlo** (Simon Cauchemez et al. [2008](#))

Examples of simulation-based methods:

- **Maximum likelihood via iterated filtering** (Ionides, Bretó, and King [2006](#))
- **Approximate Bayesian Computation** (e.g. Toni et al. [2009](#); McKinley, Cook, and Deardon [2009](#); Conlan et al. [2012](#); Brooks Pollock, Roberts, and Keeling [2014](#))
- **Pseudo-marginal methods** (e.g. O'Neill et al. [2000](#); Beaumont [2003](#); Andrieu and Roberts [2009](#); McKinley et al. [2014](#))
- **Particle MCMC** (Andrieu, Doucet, and Holenstein [2010](#); Drovandi, Pettitt, and McCutchan [2016](#))
- **Synthetic likelihood** (Wood [2010](#))
- **History matching** (with **emulation**) (e.g. Andrianakis et al. [2015](#); McKinley et al. [2018](#))

**Require:**  $\theta^{(0)}$ .

**for**  $i = 1, \dots, n$  **do**

Propose **candidate**  $\theta' \sim q(\cdot | \theta^{(i-1)})$ .

Calculate the **acceptance probability**:

$$\alpha = \min \left( 1, \frac{\hat{f}(\mathbf{y} | \theta') f(\theta')}{\hat{f}(\mathbf{y} | \theta^{(i-1)}) f(\theta^{(i-1)})} \times \frac{q(\theta^{(i-1)} | \theta')}{q(\theta' | \theta^{(i-1)})} \right)$$

Sample  $u \sim U(0, 1)$

**if**  $u < \alpha$  **then**

$$\theta^{(i)} = \theta'$$

**else**

$$\theta^{(i)} = \theta^{(i-1)}$$

**end if**

**end for**

One option is to simply plug this **estimate** into a standard Metropolis-Hastings algorithm in place of the true likelihood.

Remarkably, as long as this estimate is **unbiased**, this will still converge to the **true** posterior.

This approach is known as **pseudo-marginal MCMC**.

Beaumont (2003); Andrieu and Roberts (2009).

One option is to replace the likelihood,  $f(\mathbf{y} \mid \theta)$ , by a **Monte Carlo** estimate:

$$\begin{aligned} f(\mathbf{y} \mid \theta) &= \int_{\mathbf{x}} f(\mathbf{y} \mid \mathbf{x}, \theta) f(\mathbf{x} \mid \theta) d\mathbf{x} \\ &\approx \frac{1}{M} \sum_{i=1}^M f(\mathbf{y} \mid \mathbf{x}_i, \theta), \end{aligned}$$

where  $\mathbf{x}_i \sim f(\mathbf{x} \mid \theta)$  are simulations from the underlying model.

This provides an **unbiased** estimate for  $f(\mathbf{y} \mid \theta)$ .

The efficiency (i.e. **mixing**) of pseudo-marginal MCMC relies on the **variance** of the **estimator**  $\hat{f}(\mathbf{y} \mid \theta)$ .

- If the variance is **small**, then mixing will be **improved**.
- If the variance is **large**, then mixing will be **poor**.

We can reduce the variance by:

- increasing the number of simulations  $M \rightarrow$  higher computational burden;
- improving the estimator.

This leads on to the idea of **particle MCMC** (Andrieu, Doucet, and Holenstein [2010](#)).

In essence this aims to use **Sequential Monte Carlo**<sup>†</sup> to produce an **unbiased** estimate of the likelihood that has **lower variance** than a vanilla Monte Carlo estimate.

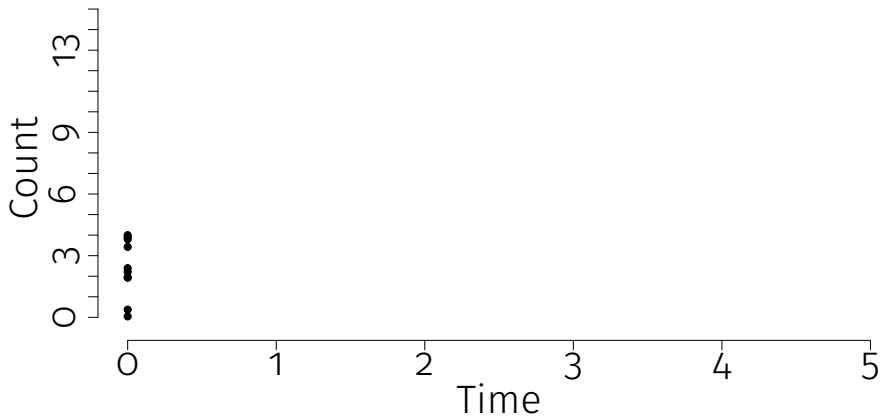
One of the earliest and most widely used particle filters is known as the **bootstrap particle filter** (Gordon, Salmond, and Smith [1993](#)).

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<sup>†</sup>i.e. **particle filtering**

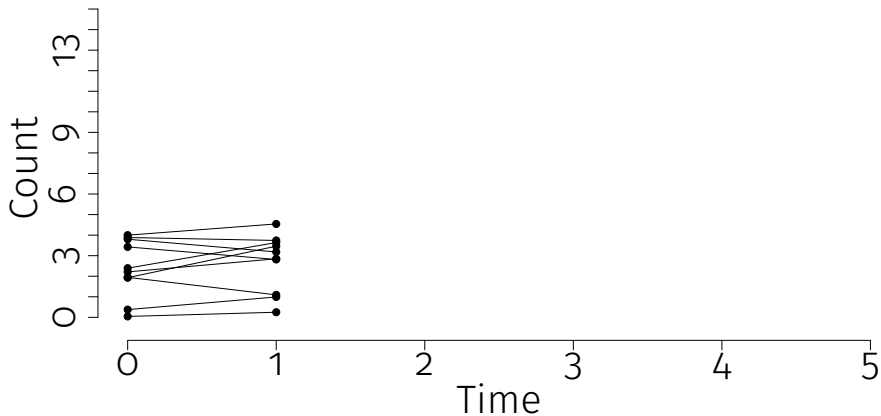
Each **particle** now corresponds to the **unobserved states** of the system at time 0,  $\mathbf{x}_0 = (\mathbf{x}_0^1, \dots, \mathbf{x}_0^M)$ . The parameters are **fixed**.

1. Each particle  $m$  is propagated forwards in time by **simulating** from the model  $\mathbf{x}_1^m \sim f(\mathbf{x} \mid \mathbf{x}_0^m, \theta)$ .
2. Each new particle is **weighted** according to the **observation process**,  $f(\mathbf{y} \mid \mathbf{x}_1^m, \theta)$ .
3. These weights are **normalised**, and a **re-sampling** step undertaken.
4. The new set of particles are propagated forwards to time  $t + 1$  and so on...

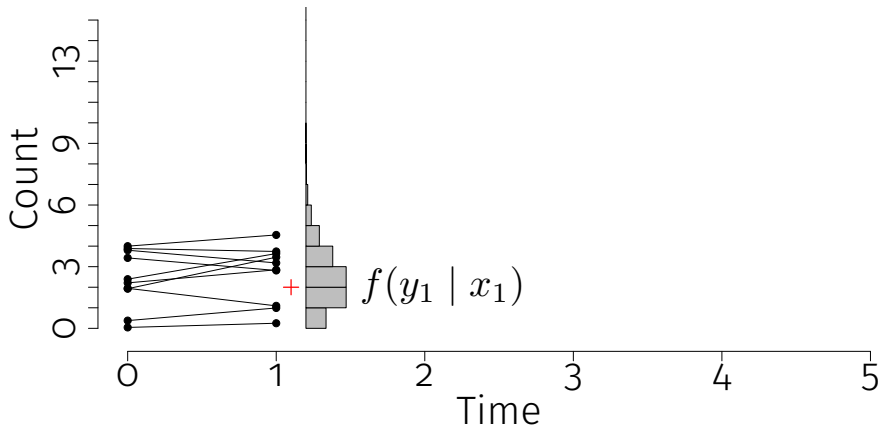




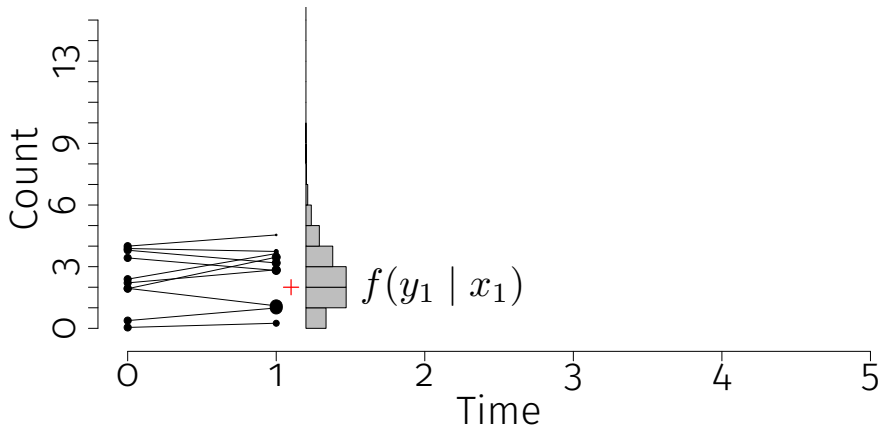
# Bootstrap particle filter



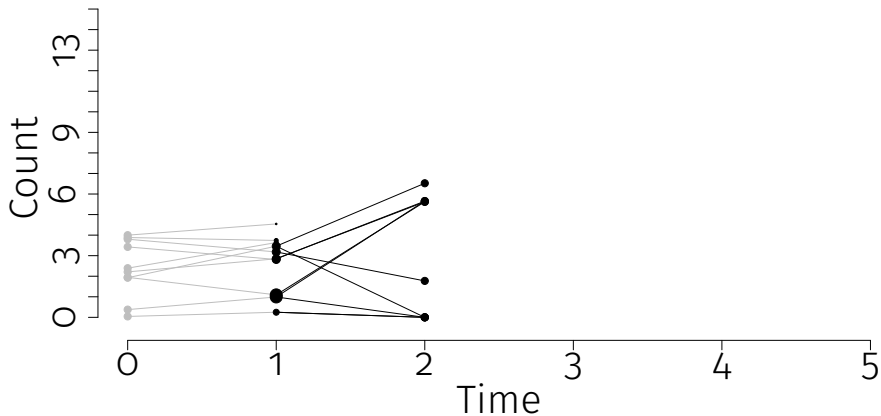
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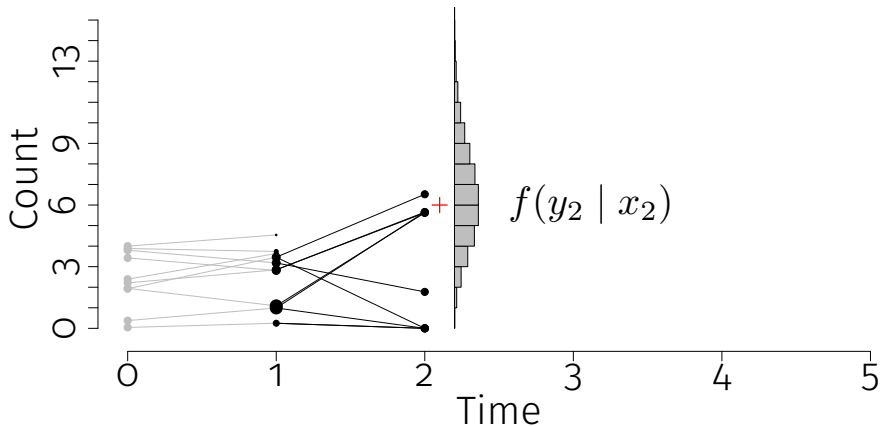
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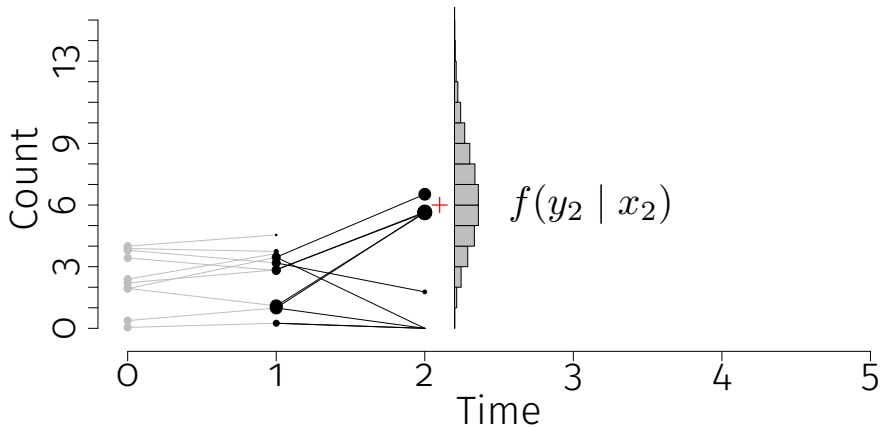
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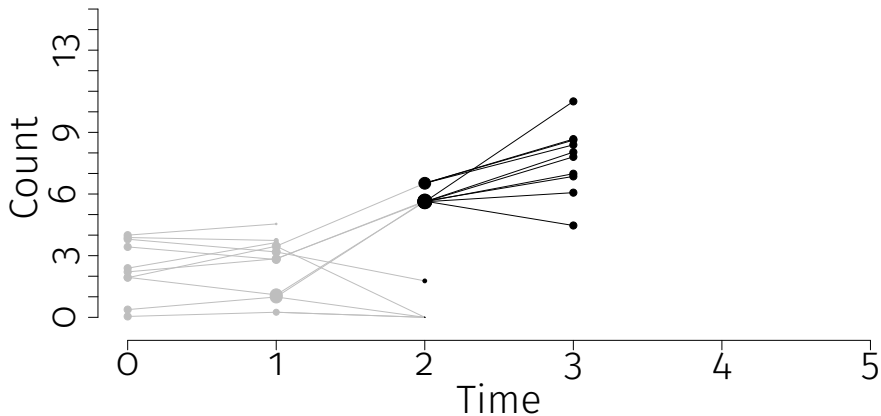
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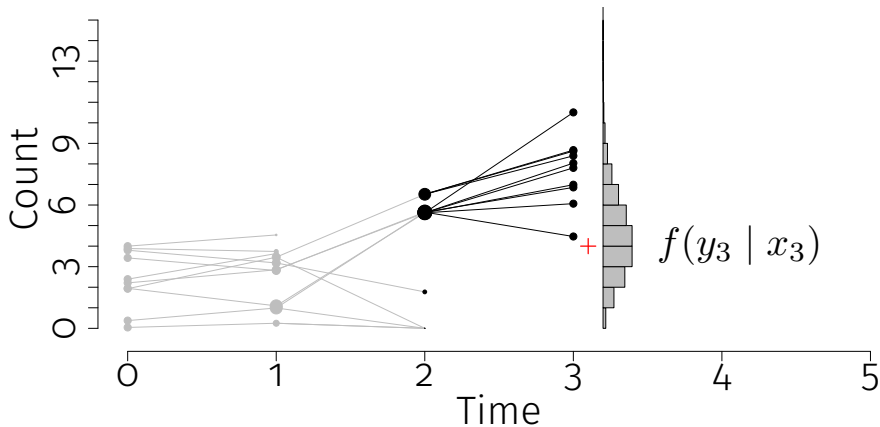
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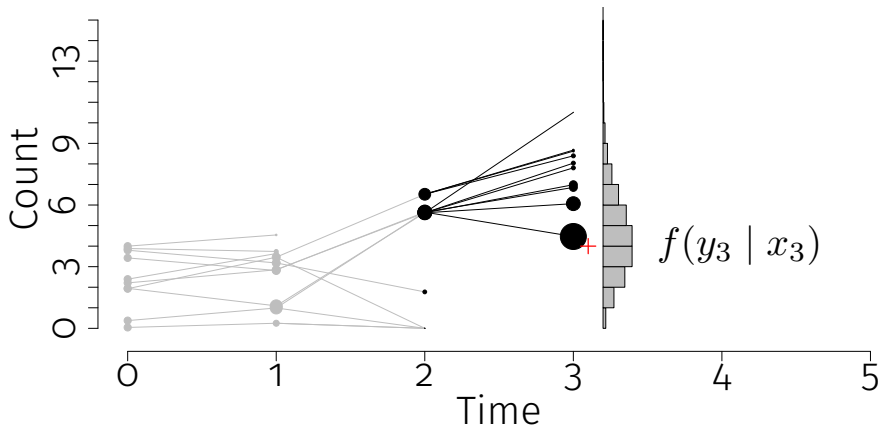


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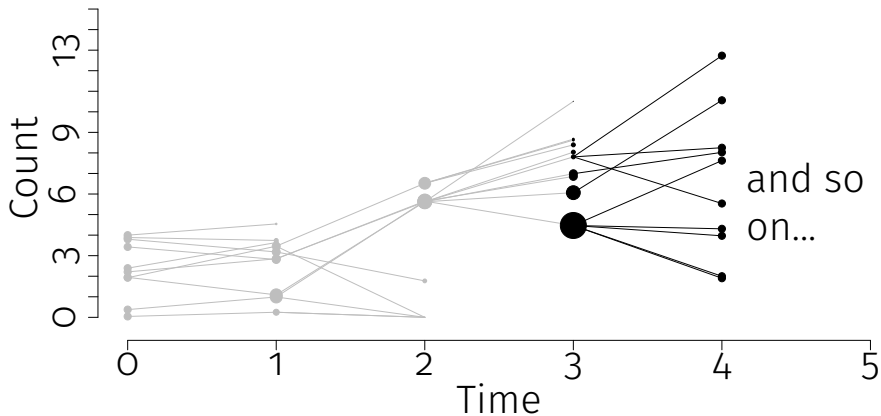




# Bootstrap particle filter



# Bootstrap particle filter



We can generate an **unbiased estimate** of the conditional densities:

$$\hat{f}(y_t | y_{0:(t-1)}) = \frac{1}{M} \sum_{m=1}^M f(y_t | \mathbf{x}_t^m, \theta),$$

where  $y_{0:(t-1)}$  corresponds to the observed time-series counts at time  $t_0, t_1, \dots, t_{t-1}$ .

It turns out that we can also derive an **unbiased** estimate of the overall **likelihood** as:

$$\hat{f}(\mathbf{y} | \theta) = f(y_0) \prod_{t=1}^T \hat{f}(y_t | y_{0:(t-1)}).$$

Hence we can generate an **unbiased** estimate of the likelihood which **numerically** integrates over the **hidden states**.

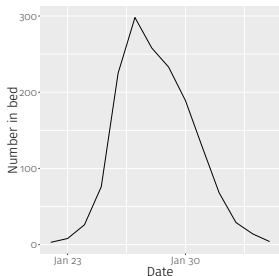
We can then plug this estimate into a standard Metropolis-Hastings algorithm to produce a **pseudo-marginal** MCMC routine that will converge to the *correct posterior distribution in probability*.

This approach only requires a **simulation** model, and an **observation process**.

The bootstrap particle filter we've used is defined for **time-series** counts, and can be extended in various ways.

# Example: flu in boarding school

To illustrate some of these ideas we can use a case study of influenza in a boarding school. These data are from a paper in the BMJ in 1978 (Anonymous 1978) and provided in the [outbreaks](#) package. We use a simple  $SIRR_1$  model:



The event probabilities are:

$$P[S_{t+\delta t} = S_t - 1, I_{t+\delta t} = I_t + 1] \approx \beta SI/N$$

$$P[I_{t+\delta t} = I_t - 1, R_{t+\delta t} = R_t + 1] \approx \gamma I$$

$$P[R_{t+\delta t} = R_t - 1, R_{1,t+\delta t} = R_{1,t} + 1] \approx \gamma_1 R$$

Here we will place a Poisson error process around the  $R$  curve, such that:

$$R_t \sim \text{Po}(R'_t + 10^{-6}),$$

where  $R_t$  is the **observed**  $R$  count at time  $t$ ,  $R'_t$  is the simulated count<sup>†</sup>.

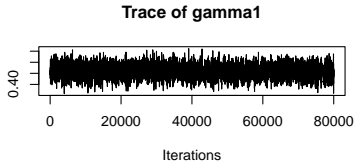
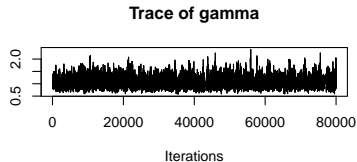
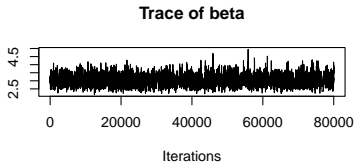
The initial population size is 763 pupils, and we assume an initial introduction of infection of a single child at day 0.

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<sup>†</sup>see e.g. Funk et al. (2016) or [here](#) for similar ideas in practice

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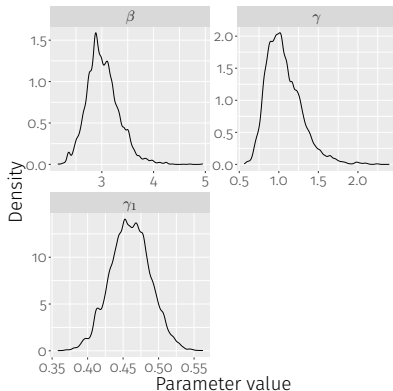
We ran a PMCMC algorithm for 100,000 iterations, discarding the first 20,000 as burn-in. We used 75 particles for the particle filter.



# Example: flu in boarding school

Summaries of the marginal posterior distributions are:

Parameter	Mean	2.5%	97.5%
$\beta$	3	2.5	3.7
$\gamma$	1.1	0.73	1.6
$\gamma_1$	0.46	0.41	0.52





Particle MCMC is a powerful approach for inference in **partially observed** systems (see e.g. Wilkinson [2012](#) or his associated [blog](#) for fantastic explanations of these methods).

It is often used when there is some form of **stochastic** discrepancy / observation process mapping the **hidden** states to the **observed** states.

Other particle filters exist, such as the **Alive Particle Filter** (Jasra et al. [2013](#)), and the system can be extended to the **ABC** setting, where approximate matching around data points is used (Drovandi, Pettitt, and Lee [2014](#); McKinley et al. [2020](#)).

## Partially Observed Markov Processes:

- [pomp](#)
- [SimBIID](#)<sup>†</sup>
- [SimInf](#)<sup>‡</sup>
- [nimble](#)<sup>§</sup>
- [hmer](#)<sup>¶</sup>

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<sup>†</sup>designed mostly for teaching purposes, but should work for simple models

<sup>‡</sup>now implements ABC-SMC (e.g. Toni et al. [2009](#); McKinley, Cook, and Deardon [2009](#))

<sup>§</sup>now supports state-space models (although I've not used it for these)

<sup>¶</sup>hot-off-the-press! Implements emulation and history matching for epidemic models

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