

# 10: Introduction to Bayesian Computation

Taylor

University of Virginia

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

Introduction

Accept-Reject Sampling

Importance Sampling

Importance Sampling with Resampling

Sequential Monte Carlo

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

# Introduction

"10"

Taylor

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

This chapter gives an overview for how to approximate intractable quantities such as posterior expectations and predictions.

**Numerical integration** methods approximate integrals.

These methods can be loosely categorized as either **stochastic** or **deterministic** (e.g. quadrature methods).

**Deterministic methods** don't draw samples, and instead approximate the integral by summing up approximate volumes:

$$E[h(\theta) \mid y] = \int h(\theta)p(\theta \mid y)d\theta \approx \frac{1}{S} \sum_{s=1}^S w_s h(\theta^s)p(\theta^s \mid y)$$

# Approximating the posterior on a grid

We used this method earlier for the rat tumor example!

Given that we can evaluate the unnormalized target  $p(\theta, y) = p(y | \theta)p(\theta)$ , we choose a nonrandom grid of points (think seq)  $\theta_1, \dots, \theta_S$ , and then we approximate the the continuous posterior with a discrete random variable with pmf equal to

$$\tilde{p}(\theta_j | y) = \frac{p(y | \theta_j)p(\theta_j)}{\sum_{s=1}^S p(y | \theta_s)p(\theta_s)}$$

for any  $\theta_j \in \{\theta_1, \dots, \theta_S\}$ . Then

$$E[h(\theta) | y] \approx \sum_{j=1}^S h(\theta_j) \tilde{p}(\theta_j | y).$$

[Introduction](#)[Accept-Reject Sampling](#)[Importance Sampling](#)[Importance Sampling with Resampling](#)[Sequential Monte Carlo](#)[References](#)

**Stochastic methods** involve sample averages of simulated draws from some distribution. There are **many** ways to do this, but here are a couple examples that we've seen:

$$E[h(\theta) | y] = \int h(\theta)p(\theta | y)d\theta \approx \frac{1}{S} \sum_{s=1}^S h(\theta^s)$$

with  $\theta^s \sim p(\theta | y)$ , or

$$E[h(\tilde{y}) | y] = \int h(\tilde{y})p(\tilde{y} | y)d\tilde{y} \approx \frac{1}{S} \sum_{s=1}^S h(\tilde{y}^s)$$

with  $\tilde{y} \sim p(\tilde{y} | y)$

## Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

Drawing  $\tilde{y}$  samples can be done in a two-stage way:

1. draw  $\theta^s \sim p(\theta \mid y)$
2. draw  $\tilde{y}^s \sim p(\tilde{y} \mid \theta^s)$



Drawing  $\tilde{y}$  samples can be done in a two-stage way:

1. draw  $\theta^s \sim p(\theta \mid y)$
2. draw  $\tilde{y}^s \sim p(\tilde{y} \mid \theta^s)$

If you can derive  $E(h(\tilde{y}) \mid \theta)$ , you should probably use a Rao-Blackwellized procedure:

$$E[h(\tilde{y}) \mid y] = E[E(h(\tilde{y}) \mid \theta) \mid y] \approx \frac{1}{S} \sum_{s=1}^S E(h(\tilde{y}) \mid \theta^s)$$

with  $\theta^s \sim p(\theta \mid y)$

# The general setup

"10"

Taylor

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

From now on we will write the posterior in terms of an unnormalized density  $q(\theta | y)$ . In other words:

$$p(\theta | y) = \frac{q(\theta | y)}{\int q(\theta | y) d\theta}$$

Most (maybe all) of the sampling techniques will assume that we can't evaluate  $p(\theta | y)$ , but that we can evaluate  $q(\theta | y)$

# Rejection Sampling aka Accept-Reject sampling

"10"

Taylor

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

## Setup

1.  $p(\theta | y)$  the target, posterior
2.  $q(\theta | y) = p(y | \theta)p(\theta)$  the unnormalized target
3.  $g(\theta)$  the “instrumental” or “proposal” distribution
4. need  $q(\theta | y)/g(\theta) \leq M$  uniformly
5. need  $g \gg q$  i.e. the proposal “dominates” your target (won't divide by 0)

We are free to choose our own  $g(\theta)$ . For the time being, we assume that  $\int g(\theta)d\theta = 1$ .

# Rejection Sampling aka Accept-Reject sampling

"10"

Taylor

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

To (potentially) produce one draw:

1. propose the draw  $\theta^s \sim g(\theta)$
2. accept  $\theta^s$  with probability  $\frac{q(\theta^s|y)}{g(\theta^s)M}$

# Rejection Sampling aka Accept-Reject sampling

"10"

Taylor

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

To (potentially) produce one draw:

1. propose the draw  $\theta^s \sim g(\theta)$
2. accept  $\theta^s$  with probability  $\frac{q(\theta^s|y)}{g(\theta^s)M}$

Note this is the same as

1. propose the draw  $\theta^s \sim g(\theta)$
2. draw  $U \sim \text{Uniform}(0, 1]$
3. accept  $\theta^s$  if  $U < q(\theta^s | y) / \{g(\theta^s)M\}$

$$\begin{aligned}P\left(\theta \leq t \mid U \leq \frac{q(\theta | y)}{Mg(\theta)}\right) &= \frac{P\left(\theta \leq t, U \leq \frac{q(\theta | y)}{Mg(\theta)}\right)}{P\left(U \leq \frac{q(\theta | y)}{Mg(\theta)}\right)} \\&= \frac{\int_{-\infty}^t \int_0^{\frac{q(\theta | y)}{Mg(\theta)}} g(\theta) 1 du d\theta}{\int_{-\infty}^{\infty} \int_0^{\frac{q(\theta | y)}{Mg(\theta)}} g(\theta) 1 du d\theta} \\&= \frac{\int_{-\infty}^t g(\theta) \frac{q(\theta | y)}{Mg(\theta)} d\theta}{\int_{-\infty}^{\infty} g(\theta) \frac{q(\theta | y)}{Mg(\theta)} d\theta} \\&= \frac{\int_{-\infty}^t q(\theta | y) d\theta}{\int_{-\infty}^{\infty} q(\theta | y) d\theta} \\&= P(\theta \leq t | y).\end{aligned}$$

# Example 1

"10"

Taylor

Assume  $y \sim \text{Normal}(\theta, 1)$ , and  $p(\theta) = \frac{1}{\pi(1+\theta^2)}$ . Our goal is to draw from

$$\begin{aligned} p(\theta | y) &\propto q(\theta | y) \\ &= p(y | \theta)p(\theta) \\ &= \frac{1}{\sqrt{2\pi}} \exp\left[-\frac{1}{2}(y - \theta)^2\right] \frac{1}{\pi(1 + \theta^2)} \\ &\propto \exp\left[-\frac{(\theta - y)^2}{2} - \log(1 + \theta^2)\right], \end{aligned}$$

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

# Example 1

"10"

Taylor

Let's assume that we want to use our prior distribution as a proposal:  $g(\theta) = p(\theta)$ . Then we have to find  $M$ :

$$\begin{aligned}\frac{q(\theta | y)}{g(\theta)} &= \frac{p(y | \theta)p(\theta)}{p(\theta)} \\ &= p(y | \theta) \\ &= \frac{1}{\sqrt{2\pi}} \exp \left[ -\frac{1}{2}(y - \theta)^2 \right] \\ &\leq \frac{1}{\sqrt{2\pi}} \stackrel{\text{def}}{=} M\end{aligned}$$

Our acceptance probability for draw  $\theta^s$  is then

$$q(\theta^s | y) / \{g(\theta^s)M\} = p(y | \theta^s) / M = \exp \left[ -\frac{1}{2}(y - \theta^s)^2 \right]$$

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References



# Rejection Sampling aka Accept-Reject sampling

"10"

Taylor

Generally a good strategy is work with logarithms, and then exponentiate as late as possible.

```
y <- 2 # fake data
num_trials <- 1000
theta_proposals <- rt(num_trials, 1)
us <- runif(num_trials, min = 0, max = 1)
log_accept_prob <- function(theta){
  -.5*(y - theta)^2
}
probs <- exp(log_accept_prob(theta_proposals))
accepts <- us < probs
hist(theta_proposals[accepts]) # only the accepted draws
#hist(theta_proposals) # all draws!
```

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

# Importance Sampling

"10"

Taylor

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

**importance sampling** also involves ratios like the previous algorithm. However, instead of using those ratios to either accept or discard samples, it uses the ratios to weight samples.

# Importance Sampling

"10"

Taylor

Introduction

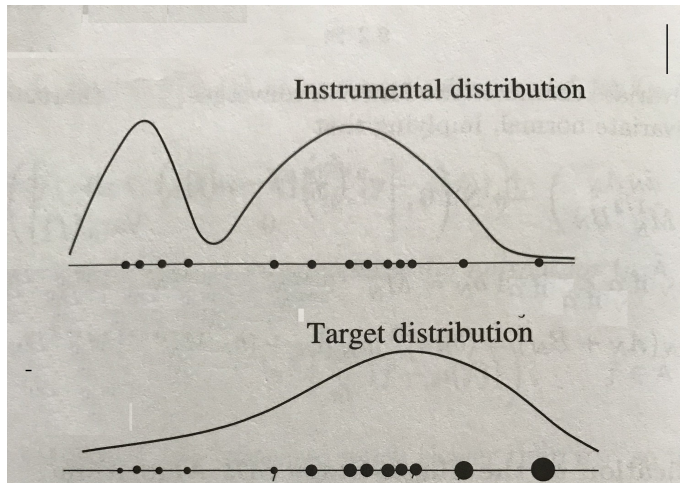
Accept-Reject  
Sampling

**Importance  
Sampling**

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References



Original, unedited image is from  
<https://www.springer.com/us/book/9780387402642>

## Setup

1.  $p(\theta | y)$  the target, posterior
2.  $q(\theta | y) = p(y | \theta)p(\theta)$  the unnormalized target
3.  $g(\theta)$  the “instrumental” or “proposal” distribution
4.  $g \gg q$  i.e. the proposal dominates your target

We are free to choose our own  $g(\theta)$ . For the time being, we assume that  $\int g(\theta)d\theta = 1$ .

# Importance Sampling

"10"

Taylor

Introduction

Accept-Reject  
Sampling

**Importance  
Sampling**

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

Algorithm: for each iteration  $s$

1. draw  $\theta^s \sim g(\theta)$
2. calculate unnormalized weight  $\tilde{w}(\theta^s) = \frac{q(\theta^s|y)}{g(\theta^s)}$
3. calculate normalized weights  $w(\theta^s) = \tilde{w}(\theta^s) / \sum_r \tilde{w}(\theta^r)$

Final calculation:

$$E_q[h(\theta) | y] \approx \sum_s w(\theta^s) h(\theta^s)$$

# Importance Sampling

"10"

Taylor

Motivation:

$$\begin{aligned} E_q[h(\theta) | y] &= \int h(\theta) p(\theta | y) d\theta \\ &= \frac{\int h(\theta) q(\theta | y) d\theta}{\int q(\theta | y) d\theta} \\ &= \frac{\int h(\theta) \frac{q(\theta|y)}{g(\theta)} g(\theta) d\theta}{\int \frac{q(\theta|y)}{g(\theta)} g(\theta) d\theta} \end{aligned}$$

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

# Importance Sampling

"10"

Taylor

Motivation:

$$\begin{aligned} E_q[h(\theta) | y] &= \int h(\theta) p(\theta | y) d\theta \\ &= \frac{\int h(\theta) q(\theta | y) d\theta}{\int q(\theta | y) d\theta} \\ &= \frac{\int h(\theta) \frac{q(\theta | y)}{g(\theta)} g(\theta) d\theta}{\int \frac{q(\theta | y)}{g(\theta)} g(\theta) d\theta} \end{aligned}$$

So first:

$$\frac{1}{S} \sum_{s=1}^S \frac{q(\theta^s | y)}{g(\theta^s)} \rightarrow E_g \left[ \frac{q(\theta | y)}{g(\theta)} \right] = \int \frac{q(\theta | y)}{g(\theta)} g(\theta) d\theta = \int q(\theta | y) d\theta$$

Introduction

Accept-Reject  
Sampling

**Importance  
Sampling**

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

# Importance Sampling

"10"

Taylor

Introduction

Accept-Reject  
Sampling

**Importance  
Sampling**

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

$$1. E_q[h(\theta) | y] = \frac{\int h(\theta)q(\theta|y)d\theta}{\int q(\theta|y)d\theta}$$

$$2. \frac{1}{S} \sum_{s=1}^S \frac{q(\theta^s|y)}{g(\theta^s)} \rightarrow \int q(\theta | y)d\theta \text{ ( for the denominator)}$$

And second:

$$\frac{1}{S} \sum_{s=1}^S h(\theta^s) \frac{q(\theta^s | y)}{g(\theta^s)} \rightarrow E_g \left[ h(\theta) \frac{q(\theta | y)}{g(\theta)} \right] = \int h(\theta)q(\theta | y)d\theta$$

which converges to the numerator



# Importance Sampling

"10"

Taylor

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

$$1. E_q[h(\theta) | y] = \frac{\int h(\theta) q(\theta|y) d\theta}{\int q(\theta|y) d\theta}$$

$$2. \frac{1}{S} \sum_{s=1}^S \frac{q(\theta^s|y)}{g(\theta^s)} \rightarrow \int q(\theta | y) d\theta$$

$$3. \frac{1}{S} \sum_{s=1}^S h(\theta^s) \frac{q(\theta^s|y)}{g(\theta^s)} \rightarrow \int h(\theta) q(\theta | y) d\theta$$

So finally

$$\sum_{i=1}^S w(\theta^s) h(\theta^s) = \frac{\sum_{s=1}^S h(\theta^s) \frac{q(\theta^s|y)}{g(\theta^s)}}{\sum_{r=1}^S \frac{q(\theta^r|y)}{g(\theta^r)}} = \frac{\frac{1}{S} \sum_{s=1}^S h(\theta^s) \frac{q(\theta^s|y)}{g(\theta^s)}}{\frac{1}{S} \sum_{r=1}^S \frac{q(\theta^r|y)}{g(\theta^r)}} \rightarrow E[h(\theta) | y]$$

where  $w(\theta^s) = \frac{q(\theta^s|y)}{g(\theta^s)} / \sum_{r=1}^S \frac{q(\theta^r|y)}{g(\theta^r)}$  are the self-normalized weights

## Example 2

"10"

Taylor

Assume  $y \sim \text{Normal}(\theta, 1)$ , and  $p(\theta) = \frac{1}{\pi(1+\theta^2)}$ . Approximate  $E_q[\theta \mid y]$  using proposal  $g(\theta) = p(\theta)$ .

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

## Example 2

"10"

Taylor

Assume  $y \sim \text{Normal}(\theta, 1)$ , and  $p(\theta) = \frac{1}{\pi(1+\theta^2)}$ . Approximate  $E_q[\theta \mid y]$  using proposal  $g(\theta) = p(\theta)$ .

If we sample from  $g(\theta) = p(\theta) = \frac{1}{\pi(1+\theta^2)}$  then the unnormalized weights are

$$\begin{aligned}\tilde{w}(\theta^s) &= \frac{q(\theta^s \mid y)}{g(\theta^s)} \\ &= p(y \mid \theta^s) \\ &= \frac{1}{\sqrt{2\pi}} \exp \left[ -\frac{1}{2}(y - \theta^s)^2 \right]\end{aligned}$$

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

## Example 2

"10"

Taylor

Assume  $y \sim \text{Normal}(\theta, 1)$ , and  $p(\theta) = \frac{1}{\pi(1+\theta^2)}$ . Approximate  $E_q[\theta \mid y]$  using proposal  $g(\theta) = p(\theta)$ .

If we sample from  $g(\theta) = p(\theta) = \frac{1}{\pi(1+\theta^2)}$  then the unnormalized weights are

$$\begin{aligned}\tilde{w}(\theta^s) &= \frac{q(\theta^s \mid y)}{g(\theta^s)} \\ &= p(y \mid \theta^s) \\ &= \frac{1}{\sqrt{2\pi}} \exp \left[ -\frac{1}{2}(y - \theta^s)^2 \right]\end{aligned}$$

then normalize these...

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

## Example 2

"10"

Taylor

```
y <- 2 # fake data
num_samples <- 1000000
theta_draws <- rt(num_samples , 1)
log_unnorm_weight <- function(theta){
  # can ignore sqrt(2pi) because it will cancel out
  -.5*(y - theta)^2
}
lunws <- log_unnorm_weight(theta_draws)
norm_weights <- exp(lunws)/sum(exp(lunws))
sum(norm_weights * theta_draws)
hist(norm_weights)
```

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

## Example 2

"10"

Taylor

The choice of proposal is very important. In a HW problem, you will derive the following using the delta method:

$$\text{Var}_g \left( \sum_{s=1}^S w(\theta^s) h(\theta^s) \right) \approx \frac{1}{S} E_g \left[ \underbrace{\left( \frac{\tilde{w}(\theta)}{E_g[\tilde{w}(\theta)]} \right)^2}_{!} (h(\theta) - E_q[h(\theta)])^2 \right]$$

Beware of proposals that have tails that are thin relative to the target!

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

## Example 2

"10"

Taylor

Introduction

Accept-Reject  
Sampling

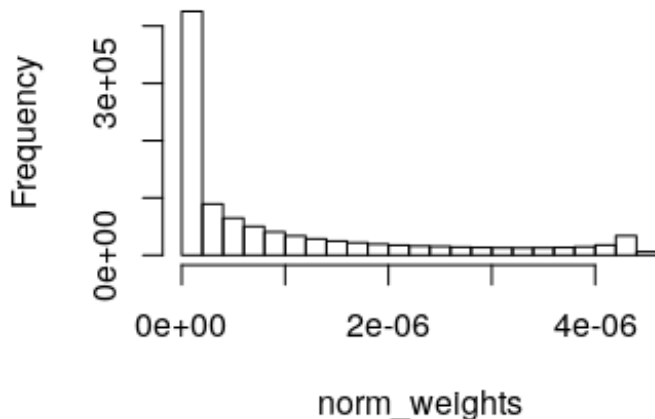
Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

### Histogram of norm\_weights



## Example 2

"10"

Taylor

Beware of bad proposal distributions!

A sample estimator of this approximate variance is

$$\sum_{s=1}^S w(\theta^s) \left( h(\theta^s) - \hat{E}[h(\theta)] \right)^2$$

where  $\hat{E}[h(\theta)] = \sum_s w(\theta^s) h(\theta^s)$ .

Note the weights aren't the same for each sample, like a "standard" estimation of the sample variance.

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References



## Example 2

"10"

Taylor

```
y <- 2 # fake data
log_unnorm_weight <- function(theta){
  # can ignore sqrt(2pi) because it will cancel out
  -.5*(y - theta)^2 }
getISEstimator <- function(num_samples){
  theta_draws <- rt(num_samples, 1)
  lunws <- log_unnorm_weight(theta_draws)
  norm_weights <- exp(lunws)/sum(exp(lunws))
  estimator <- sum(norm_weights * theta_draws)
  list("estimate" = estimator,
       "approx_var" =
         sum( norm_weights*(theta_draws - estimator)^2) ) }
# two ways to calculate standard errors
num_samps_per_estimate <- 10
sqrt(getISEstimator(num_samps_per_estimate)$approx_var)
sd(replicate(1000,
  getISEstimator(num_samps_per_estimate)$estimate))
```

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

# Effective Sample Size

"10"

Taylor

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

Effective sample size calculations use a second order delta method.

Calculating

$$\text{ESS} = \frac{1}{\sum_{i=1}^S w_i^2}$$

will tell you, effectively, how many iid samples you have.

Importance Sampling gives you weighted draws  $(\theta^1, w(\theta^1)), (\theta^2, w(\theta^2)), \dots$

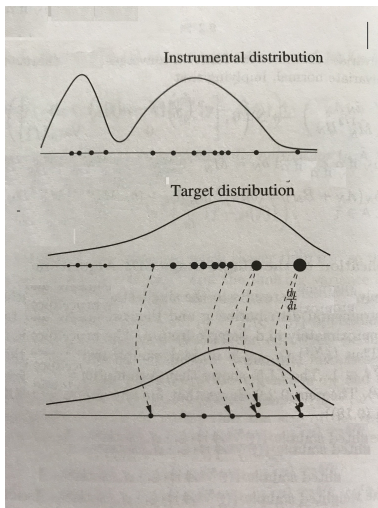
You can draw from these, with replacement. At the expense of more variance, it will give you unweighted draws from your target distribution:  $\tilde{\theta}^1, \tilde{\theta}^2, \dots$

This is known as **factored sampling** or **importance sampling with resampling** or **sampling importance resampling** (SIR).

# Adding Resampling

"10"

Taylor



Original, unedited image is from

<https://www.springer.com/us/book/9780387402642>

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

Stage 1: do importance sampling to get  $\{\theta^i, w(\theta^i)\}_{i=1}^S$ .

Stage 2: for  $i = 1, \dots, S$ , select

$$P[\tilde{\theta}^i = \theta^j \mid \theta^1, w(\theta^1), \dots, \theta^S, w(\theta^S)] = \frac{w(\theta^j)}{\sum_k w(\theta^k)}.$$

Another way to write it:

Stage 1: do importance sampling to get  $\{\theta^i, w(\theta^i)\}_{i=1}^S$ .

Stage 2: for  $i = 1, \dots, S$ , select indexes

$$P[I^i = j \mid \theta^1, w(\theta^1), \dots, \theta^S, w(\theta^S)] = \frac{w(\theta^j)}{\sum_k w(\theta^k)}$$

and then set

$$\tilde{\theta}^i = \theta^{I^i}.$$

## Example 2 revisited

"10"

Taylor

```
y <- 2 # fake data
log_unnorm_weight <- function(theta){
  # can ignore sqrt(2pi) because it will cancel out
  -.5*(y - theta)^2 }
num_samples <- 10000
theta_draws <- rt(num_samples , 1)
lunws <- log_unnorm_weight(theta_draws)
# note: prob arg automatically normalizes
random_indexes <- sample(x = num_samples,
                        size = num_samples,
                        replace = T,
                        prob = exp(lunws))
sort(random_indexes) # see there are repeats!
resampled_draws <- theta_draws[random_indexes]
hist(resampled_draws) # can't do this unless we resample
```

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

# Going sequential

"10"

Taylor

Resampling adds variance, so why do it?

It throws away bad samples, and duplicates promising ones.

When you're looking at a sequence of distribution targets, this can have a good effect on future samples.

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References



# Going sequential

"10"

Taylor

Resampling adds variance, so why do it?

It throws away bad samples, and duplicates promising ones.

When you're looking at a sequence of distribution targets, this can have a good effect on future samples.

**sequential monte carlo** or **particle filtering** methods are basically doing SIR over and over again.

At "time"  $t - 1$  you just resampled, so you have draws  $\tilde{\theta}_{t-1}^1, \tilde{\theta}_{t-1}^2, \dots, \tilde{\theta}_{t-1}^S$ , and you want to turn them into draws for the next "time" period:

$$\tilde{\theta}_t^1, \tilde{\theta}_t^2, \dots, \tilde{\theta}_t^S.$$

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

# Examples of sequences of distributions

"10"

Taylor

## Data annealing<sup>1</sup>

$$p(\theta), p(\theta \mid y_1), p(\theta \mid y_{1:2}), \dots, p(\theta \mid y_{1:n}),$$

## Temperature annealing<sup>2</sup>

$$p(y \mid \theta)^{a_0} p(\theta), p(y \mid \theta)^{a_1} p(\theta), \dots p(y \mid \theta)^{a_n} p(\theta)$$


with  $0 = a_0 < a_1 < \dots < a_n = 1$ .

## filtering and smoothing in state space models

$$p(x_1 \mid y_1, \theta), \dots, p(x_n \mid y_{1:n}, \theta)$$

---

<sup>1</sup>Chopin, *A Sequential Particle Filter Method for Static Models*.

<sup>2</sup>Neal, "Annealed Importance Sampling". 

Introduction

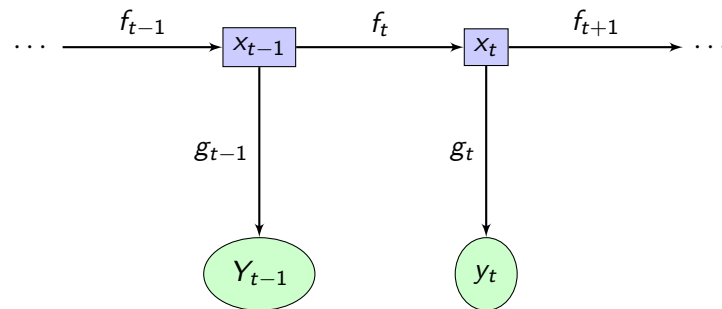
Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References



$$p(x_{1:T}, y_{1:T} \mid \theta)$$

$$= g(y_1 \mid x_1, \theta) f(x_1 \mid \theta) \prod_{t=2}^T g(y_t \mid x_t, \theta) f(x_t \mid x_{t-1}, \theta)$$

# Example: filtering in state space models

Here's an example of a state space model.  $y_t$  is a univariate time series, and  $x_t$  is a hidden/unobserved/latent time series.

$$y_t = \exp(x_t/2)\epsilon_t \quad (1)$$

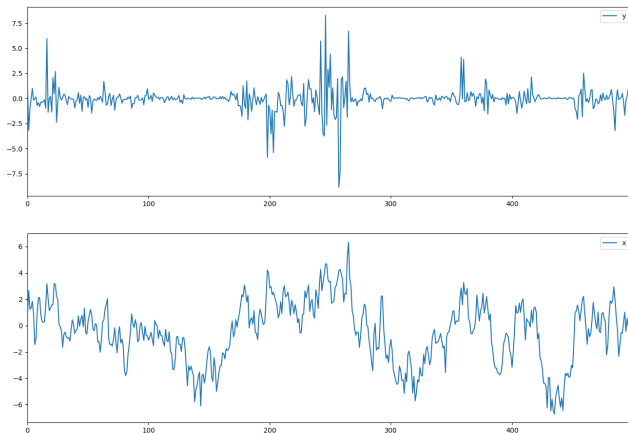
$$x_t = c + \phi x_{t-1} + v_t \quad (2)$$

We sometimes refer to (1) as  $g(y_t | x_t, \theta)$  or the observation equation, and (2) as the state transition equation or  $f(x_t | x_{t-1}, \theta)$ .

[Introduction](#)[Accept-Reject Sampling](#)[Importance Sampling](#)[Importance Sampling with Resampling](#)[Sequential Monte Carlo](#)[References](#)

# Example: filtering in state space models

$y_{1:t}$  observed,  $x_{1:t}$  hidden. Goal:  $p(x_t | y_{1:t})$  in real-time.



"10"

Taylor

[Introduction](#)

[Accept-Reject  
Sampling](#)

[Importance  
Sampling](#)

[Importance  
Sampling with  
Resampling](#)

[Sequential Monte  
Carlo](#)

[References](#)

# Example: filtering in state space models

"10"

Taylor

Introduction

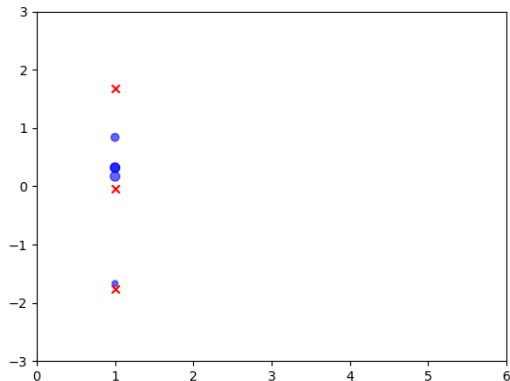
Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

**Sequential Monte  
Carlo**

References



# Going sequential

"10"

Taylor

Introduction

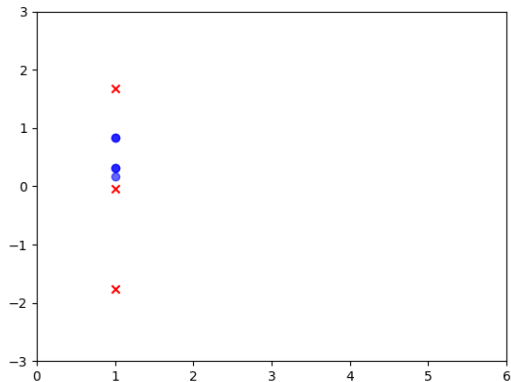
Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

**Sequential Monte  
Carlo**

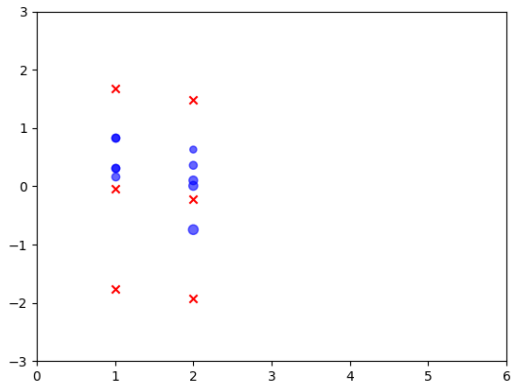
References



# Going sequential

"10"

Taylor



Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

**Sequential Monte  
Carlo**

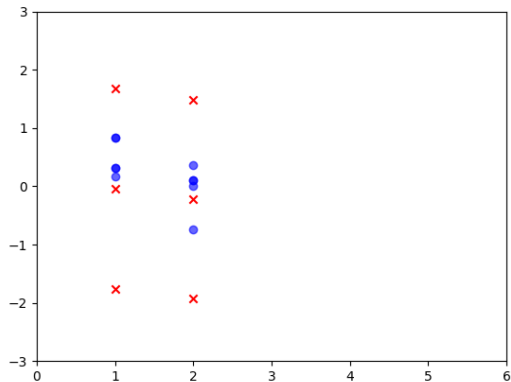
References



# Going sequential

"10"

Taylor



Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

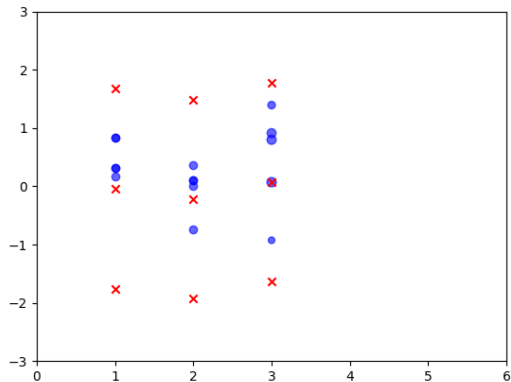
**Sequential Monte  
Carlo**

References

# Going sequential

"10"

Taylor



Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

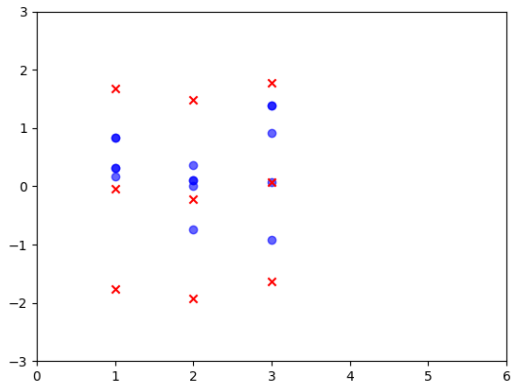
**Sequential Monte  
Carlo**

References

# Going sequential

"10"

Taylor



Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

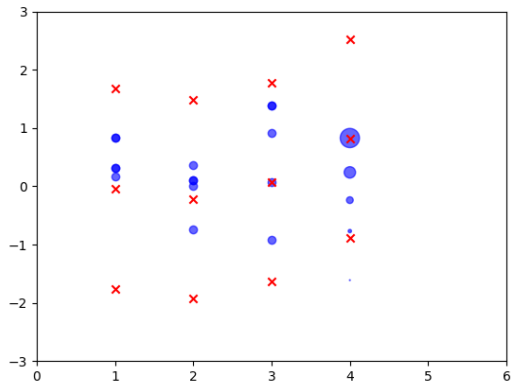
**Sequential Monte  
Carlo**

References

# Going sequential

"10"

Taylor



Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

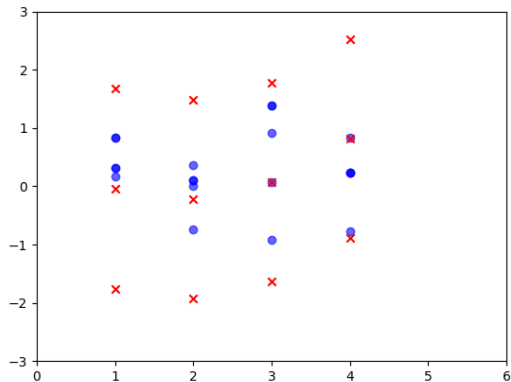
**Sequential Monte  
Carlo**

References

# Going sequential

"10"

Taylor



Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

**Sequential Monte  
Carlo**

References

# Going sequential

"10"

Taylor

Introduction

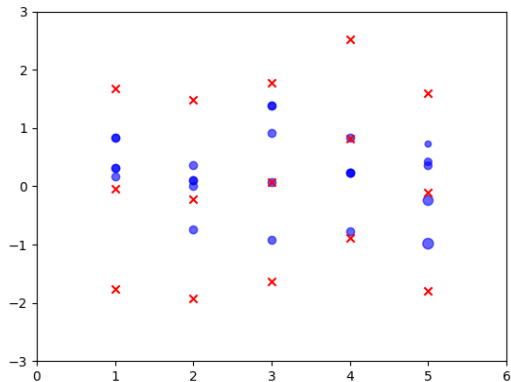
Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

**Sequential Monte  
Carlo**

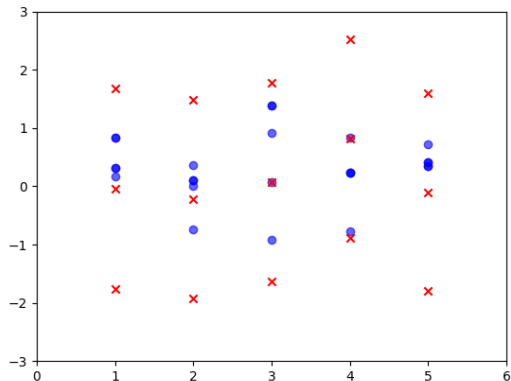
References



# Going sequential

"10"

Taylor



Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References

# Filtering recursions

Drop dependence on  $\theta$  from the notation...

$$\begin{aligned}
 p(x_{1:t}|y_{1:t}) &= C_t^{-1} p(x_t, y_t \mid x_{t-1}) p(x_{1:t-1} \mid y_{1:t-1}) \\
 &= C_t^{-1} \frac{p(x_t, y_t \mid x_{t-1})}{q_t(x_t \mid y_t, x_{t-1})} \times \\
 &\quad q_t(x_t \mid y_t, x_{t-1}) p(x_{1:t-1} \mid y_{1:t-1}) \\
 &= C_t^{-1} \frac{g(y_t|x_t)f(x_t|x_{t-1})}{q_t(x_t|x_{t-1}, y_t)} \times \\
 &\quad q_t(x_t \mid x_{t-1}, y_t) p(x_{1:t-1} \mid y_{1:t-1})
 \end{aligned}$$

Repeat through time:

1. start with samples from  $p(x_{1:t-1} \mid y_{1:t-1})$
2. mutate/propagate/extend using  $q_t(x_t \mid x_{t-1}, y_t)$
3. adjust weights by multiplying by  $\frac{g(y_t|x_t)f(x_t|x_{t-1})}{q_t(x_t|x_{t-1}, y_t)}$
4. resample, giving you particles distributed as  $p(x_{1:t} \mid y_{1:t})$

Introduction

Accept-Reject  
Sampling

Importance  
Sampling

Importance  
Sampling with  
Resampling

Sequential Monte  
Carlo

References



[Introduction](#)[Accept-Reject  
Sampling](#)[Importance  
Sampling](#)[Importance  
Sampling with  
Resampling](#)[Sequential Monte  
Carlo](#)[References](#)

Chopin, Nicolas. *A Sequential Particle Filter Method for Static Models*. 2000.

Neal, Radford M. "Annealed Importance Sampling". In: *Statistics and Computing* 11.2 (Apr. 2001), pp. 125–139. ISSN: 0960-3174. DOI: 10.1023/A:1008923215028. URL: <https://doi.org/10.1023/A:1008923215028>.