



Machine Learning for Time Series

(MLTS or MLTS-Deluxe Lectures)

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Topics overview



- Time series fundamentals and definitions (2 lectures)
- Bayesian Inference (1 lecture)
- Gussian processes (2 lectures)
- State space models (2 lectures) ←
- Autoregressive models (1 lecture)
- Data mining on time series (1 lecture)
- Deep learning on time series (4 lectures)
- Domain adaptation (1 lecture)



Review concept: State Space Models (SSMs)

SSMs are characterized by:

- Transition density: $p(z_n|z_{n-1})$
- Observation density: $p(y_n|z_n)$

Joint density can be expressed using the *chain rule*:

$$p(z_{1:n}|y_{1:n}) = p(z_1) \prod_{i=2}^{n} p(z_i|z_{i-1}) \prod_{i=1}^{n} p(y_i|z_i)$$

where $z_{1:n} = (z_1, ..., z_n)$ and $y_{1:n} = (y_1, ..., y_n)$.

Filtering is the task of estimating $p(z_n|y_{1:n})$.



Review concept: Bayesian filtering

Let $p(z_{n-1}|y_{1:n-1})$ be the filtering density at time step n-1 and we wish to determine $p(z_n|y_{1:n})$.

We can use the following iterative steps:

Prediction step:

$$p(z_n|y_{1:n-1}) = \int p(z_n|z_{n-1}) \ p(z_{n-1}|y_{1:n-1}) dz_n$$

Correction step:

$$p(z_n|y_{1:n}) = \frac{p(y_n, |z_n) p(z_n|y_{1:n-1})}{p(y_n|y_{1:n-1})}$$

If the variables are normally distributed and the transitions are linear, the Bayes filter becomes equal to the Kalman filter.



In this lecture...

- 1. Monte Carlo methods
- 2. Particle filtering: theory
- 3. Particle filtering: example and algorithmic view







Sequential Monte Carlo and Particle Filtering Monte Carlo methods





Motivation

The fundamental goal of this section is "how to find the expectation of a function f(z) with respect to a probability distribution p(z)".

$$\mathbb{E}[f] = \int f(z)p(z)dz$$



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$$\mathbb{E}[f] = \int f(z)p(z)dz$$

Sampling methods can be used to approximate this expectation using a set of samples $z^s \sim p(z)$, by the finite sum:

$$\frac{1}{S} \sum_{s=1}^{S} f(z^s)$$

where $S \in \mathbb{Z}^+$.



Motivation

However, possible problems are:

- Samples z^s might not be independent
- Depending on f, expectation might be dominated by examples with small probability
- \rightarrow We need relatively large sample size S.



Monte Carlo methods (MC)

Monte Carlo methods include a wide class of algorithms that rely on random sampling to obtain numerical results.

- Use randomness to solve problem that might be deterministic in principle.
- Used, e.g., in optimization, numerical integration and for generating draws from a probability distribution.

General idea of MC methods:

- Define a domain of possible inputs
- Generate input randomly from a probability distribution
- Perform deterministic computations on the inputs
- Aggregate results



Rejection sampling

Rejection sampling is a Monte Carlo algorithm to sample data from a sophisticated ("difficult to sample from") distribution with the help of a proxy distribution.

Let us suppose that:

- We wish to sample from a "non-standard" distribution p(z)
- Sampling directly from p(z) is difficult
- We are able to evaluate p(z) for any z, up to a certain (unknown) normalizing constant Z_p :

$$p(z) = \tilde{p}(z)/Z_p$$

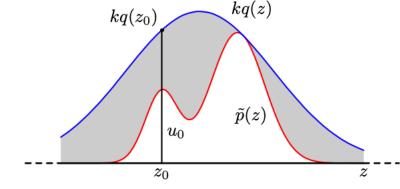
To apply rejection sampling, we make use of a simpler distribution q(z), also called **proposal** distribution \rightarrow We are able to readily draw samples from q(z)



Rejection sampling

Samples are generated from the proposal distribution q(z) and **rejected** if they fall between the unnormalized $\tilde{p}(z)$ and the scaled distribution kq(z).

→ In the side figure, samples are rejected if they fall in the grey area.



 \rightarrow Samples are accepted with probability $\frac{\tilde{p}(z)}{kq(z)}$

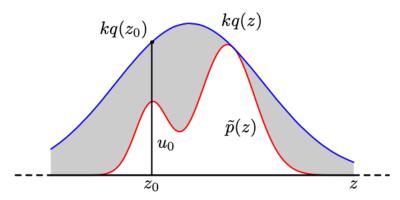
Note: k is a constant value which is chosen such that $kq(z) \ge \tilde{p}(z)$ for all values of z.



Rejection method

The success of the rejection method depend on the success in determining a suitable value for the constant k.

→ This is impractical in many cases, which leads to very small acceptance rates.





Importance sampling

Importance sampling is a method which allows approximating expectations directly.

Suppose, similarly to the previous case, that:

- It is impractical to draw samples from p(z)
- But, we can evaluate p(z) easily for any z

Uniform sampling from the space of z is inefficient (high-dimensionality) and only few samples will have significant contribution.

 \rightarrow We would like to chose samples from regions where p(z) is large.



Importance sampling

We draw samples from a proposal distribution q(z). Then,

$$\mathbb{E}[f] = \int f(z)p(z)dz = \int f(z)\frac{p(z)}{q(z)}q(z)dz \approx \frac{1}{S}\sum_{s=1}^{S}\frac{p(z^s)}{q(z^s)}f(z^s)$$

The quantity $\frac{p(z^s)}{q(z^s)}$ measures the importance of each sample and are called **importance** weights.

- Compared to rejection sampling, no samples are rejected.
- \triangleright Importance sampling do not provide itself a mechanism for drawing samples from a distribution q(z).



Sampling-importance-resampling approach

The **sampling-importance-resampling** approach also makes use of a proposal distribution q(z) and **avoids determining a constant** k.

It consists of there *general* steps:

- 1. Sampling of $\{z_1, ..., z_s\}$ from the proposal distribution q(z)
- 2. Construction of importance weights $\{w_1, ..., w_s\}$
- 3. Resampling from a discrete distribution with probabilities given by the weights

Property. The resulting samples are approximately distributed according to p(z) and the distribution becomes correct for $S \to \infty$.







Sequential Monte Carlo and Particle Filtering Particle Filtering (PF): Theory





Motivations

Estimator	State-transition / Measurement models assumptions	Assumed noise distribution	Computational cost
Kalman Filter	Linear	Gaussian	Low
Extended Kalman Filter	Non-linear (but locally linear)	Gaussian	Low / Medium (depending on the difficulty of computing the Jacobian)
Unscented Kalman Filter	Non-linear	Gaussian	Medium



We can use sampling-importance-resampling formalism to obtain a **sequential Monte Carlo**, also said **particle filtering**.

PF is a Monte Carlo (or simulation-based) approach for recursive Bayesian inference.

- It approximates the prediction-correction cycle
- It can be used for not linear-Gaussian systems to make tractable inference algorithms
- It is widely applied in many areas (e.g., tracking, forecasting, online parameter learning, ...)

The term "particle filters" originated in in reference to mean-field interacting particle methods used in fluid mechanics since early 1960s (Del Moral, 1966).



First, we update the belief state using importance sampling.

If the proposal distribution has the form $q(z_{1:n}^S|y_{1:n})$, then the importance weights are given by

$$w_n^S \propto \frac{p(z_{1:n}^S | y_{1:n})}{q(z_{1:n}^S | y_{1:n})}$$

which can be normalized as follow:

$$\widehat{w}_n^s \propto \frac{w_n^s}{\sum_{s'} w_n^{s'}}$$



We observe that we can rewrite the numerator recursively as follows:

And, similarly, the denominator:

$$q(z_{1:n}|y_{1:n}) = q(z_n|z_{1:n-1}, y_{1:n}) q(z_{1:n-1}|y_{1:n-1})$$



Therefore, we use this formulation to derive an iterative update for the weights:

$$w_n^S \propto \frac{p(z_{1:n}^S | y_{1:n})}{q(z_{1:n}^S | y_{1:n})}$$

$$\propto \frac{p(y_n | z_n^S) p(z_n^S | z_{n-1}^S) p(z_{1:n-1}^S | y_{1:n-1})}{q(z_n^S | z_{1:n-1}^S, y_{1:n}) q(z_{1:n-1}^S | y_{1:n-1})}$$

$$= w_{n-1}^S \frac{p(y_n | z_n^S) p(z_n^S | z_{n-1}^S)}{q(z_n^S | z_{1:n-1}^S, y_{1:n})}$$

Hence, we can approximate the posterior filtered density using

$$p(\mathbf{z}_{\mathbf{n}}|\mathbf{y}_{1:\mathbf{n}}) \approx \sum_{s=1}^{S} \widehat{w}_{n}^{s} \delta_{\mathbf{z}_{n}^{s}}(\mathbf{z}_{n})$$



Particle Filtering: the degeneracy problem

The basic sequential importance fails after a few steps because most of the particles will have negligible weights.

→ This problem is know as **degeneracy problem** and it occurs when sampling in high dimensional spaces

We can quantify the degree of degeneracy by using the effective sampling size, defined by:

$$S_{eff} = \frac{1}{\sum_{s=1}^{S} (w_n^s)^2}$$

When the variance of the weights is too large, we are wasting resources updating particles with negligible weights.



Particle filtering: the resampling step

The degeneracy problem can be solved by adding a resampling step.

Wheneven the effective sampling size S_{eff} drops below a threshold:

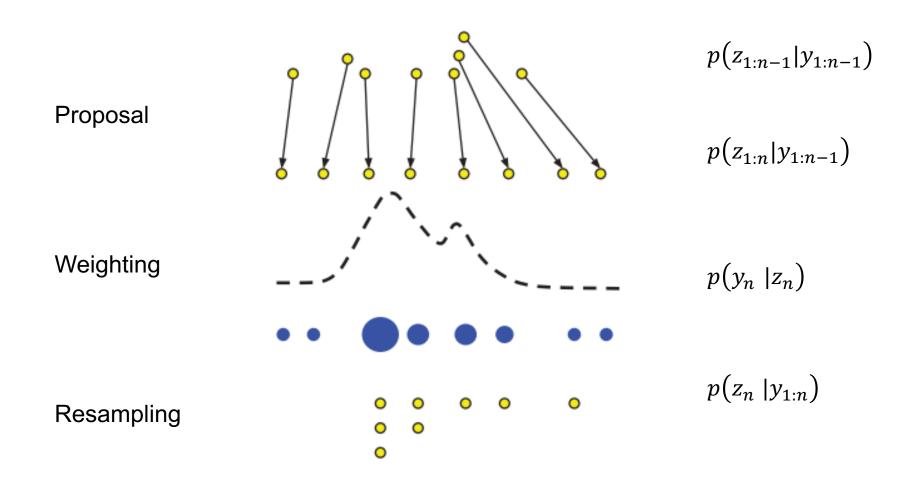
- we eliminate particles with low weights and
- we create replicates of the survival particles.

In particular, we generate a new set $\{z_n^{S*}\}_{s=1}^S$ by sampling replacement S times according to the weighted distribution obtained previously $\sum_{s=1}^S \widehat{w}_n^s \delta_{z_n^s}(z_n)$.

The result is an *i.i.d.* unweighted sample from the discrete density, so we set the new weights to $w_n^S = 1/S$.



PF: the overall scheme









Sequential Monte Carlo and Particle Filtering Particle Filtering (PF): Example and Algorithmic View

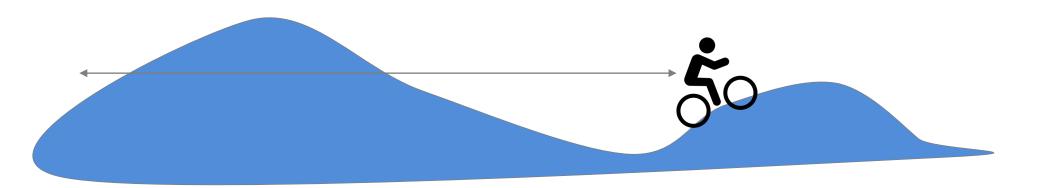






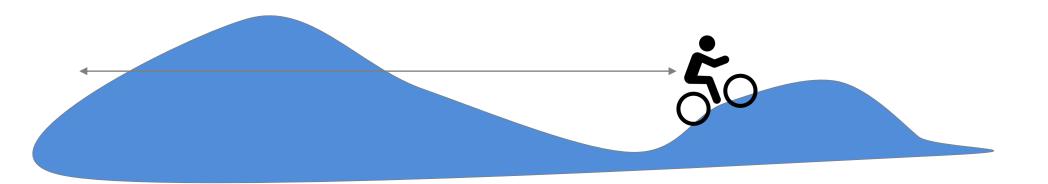


1. We want to estimate the horizontal position of the cyclist.



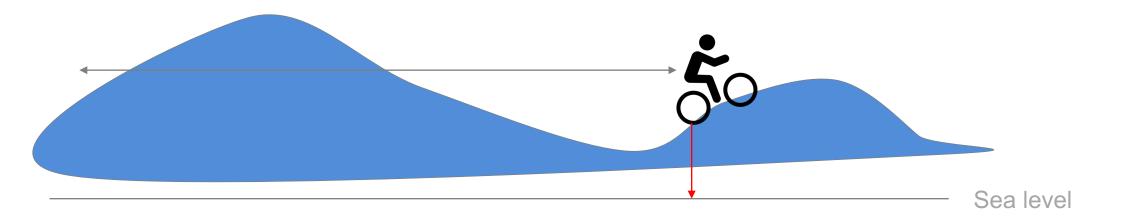


- 1. We want to estimate the horizontal position of the cyclist.
- 2. We have **knowledge** about the hills' morphology

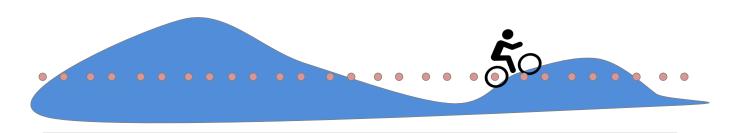




- 1. We want to estimate the horizontal position of the cyclist.
- 2. We have **knowledge** about the hills' morphology
- 3. We receive noisy measures of the altitude (e.g., in relation to sea level).



Particles (time *n*)



We draw a set of particles $\langle z_n^S, w_n^S \rangle_{S \in \{1, ..., S\}}$

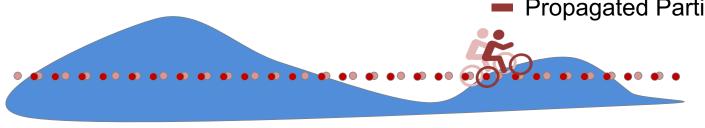
- *n* is the time index,
- z_n^s is a state hypothesis,
- w_n^s corresponding weights.

These samples represent the prior probability:

$$p(z_n|y_1,...,y_n) \simeq \sum_{s=1}^S w_n^s * \delta(z_n^s)$$



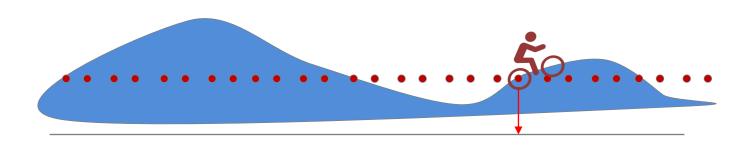
- Particles (time n)
- Propagated Particles (time n + 1)



In the **prediction step**, we use our transition model f to **propagate** particles forward in time:

$$z_{n+1}^s = f(z_n^s) + \epsilon$$

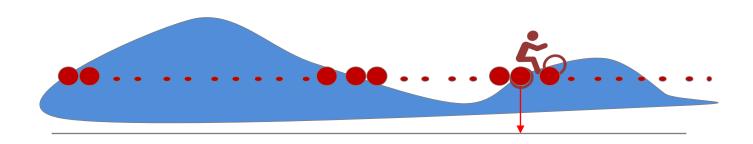




In the **correction step**, we **compute particle weights** based on the new sensory information y_{n+1} :

$$w_{n+1}^{s} = w_{n}^{s} * p(y_{n+1} | z_{n+1}^{s})$$





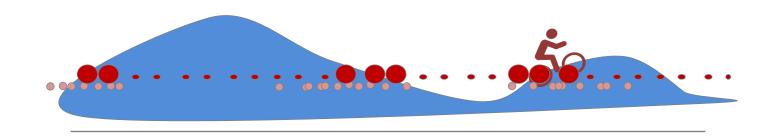
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$$w_{n+1}^{s} = w_{n}^{s} * p(y_{n+1} | z_{n+1}^{s})$$

Now the distribution $p(z_{n+1} | y_{1:t+1})$ is represented by the particle set:

$$\langle z_{n+1}^s, w_{n+1}^s \rangle_{s \in \{1, \dots, S\}}$$

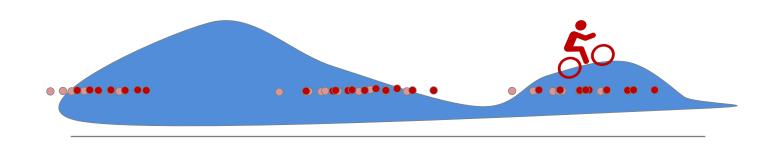




We **repeat** the previous steps:

• Sampling again from the new distribution (we have more particles close to the more likely state's hypothesis).

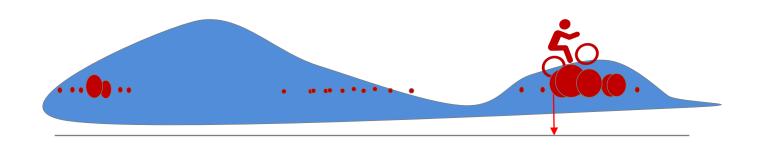




We **repeat** the previous steps:

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- We propagate again through time.





We **repeat** the previous steps:

- Sampling again from the new distribution (we have more particles close to the more likely state's hypothesis).
- We propagate again through time.
- We perform again a correction, based on the new measurement, and update the particles' weights.

• ...



PF: An algorithmic view

Initially sample S particles z_1^S

```
For n = 1, \dots, N
      For s = 1, ..., S
            Draw z_n^s \sim q(z_n^s | z_{n-1}^s, y_n)
             update and normalize w_{n+1}^{s}
            if S_{eff} < S_{\min}:
                                Resample particles
                                Re-initialize weights
            End
      End
```







Sequential Monte Carlo and Particle Filtering Recap





Recap

- Monte Carlo methods
 - Rejection method
 - Importance sampling
 - Sampling-importance-resampling approach
- Particle filtering



Particle Filtering: pros and cons

Advantages:

- Ability to represent arbitrary densities
- Adaptive focusing on probable regions of state-space
- Dealing with non-Gaussian noise

Disadvantages:

- High computational complexity
- It is difficult to determine optimal number of particles
- Number of particles increase with increasing model dimension
- Potential problems: degeneracy



