Introduction	HMM	Sampling	Re-sampling	Genetic Algs.

## Particle Filtering: An Overview

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SFU Machine Learning Reading Group

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Introduction ●○○	HMM 00000000	Sampling 0000000	Re-sampling	Genetic Algs.
What's in a	name?			

# Particle Filtering, a.k.a:

- Sequential Monte Carlo (SMC)
- Bootstrap filtering
- Condensation
- Interacting Particle Approximations
- Survival of the Fittest
- Genetic Algorithms(?)

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 Genetic Algs.

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Introduction to Particle Filtering

To the videos...

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Outline				

Introduction: What is particle filtering?

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Outline				

- Introduction: What is particle filtering?
- Particle Filters and Hidden Markov Models

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- Introduction: What is particle filtering?
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- Sampling Particles

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- Re-sampling Particles
- Particle Filters & Genetic Algorithms

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- Re-sampling Particles
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Setting Up I	Particle Filter	ring		

Notations:

- Hidden States  $\boldsymbol{x} \in \mathcal{X}$
- Observations  $\mathbf{z} \in \mathcal{Z}$

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Setting I	In Particle Fi	lterina		

 $\mathbf{z}$ 

Notations:

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Given:

- Initial state distribution  $p(\mathbf{x}_0)$
- Transition distribution  $f(\mathbf{x}_t | \mathbf{x}_{t-1})$
- Observation (i.e. scoring) distribution g(z<sub>t</sub>|x<sub>t</sub>)

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Setting I	In Particle Fi	Iterina		

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Find:

• Full trajectory distribution  $p(\mathbf{x}_{0:n}|\mathbf{z}_{1:n})$ 



## Assuming a Hidden Markov Model







- States (in blue) are hidden
- Observations (in green) are known
- Relationships are "roughly" known

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Here comes Bayes Rule					

Remember our goal:

 $p(\mathbf{x}_{0:n}|\mathbf{z}_{1:n})$ 

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Here cor	nes Baves R	ule		

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Apply Bayes Rule:

$$p(\mathbf{x}_{0:n}|\mathbf{z}_{1:n}) = rac{p(\mathbf{x}_{0:n},\mathbf{z}_{1:n})}{p(\mathbf{z}_{1:n})}$$

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The joint probabilities can be written in conditional form:

$$p(\mathbf{x}_{0:n}, \mathbf{z}_{1:n}) = p(\mathbf{x}_{0:n-1}, \mathbf{z}_{1:n-1}) f(\mathbf{x}_n | \mathbf{x}_{n-1}) g(\mathbf{z}_n | \mathbf{x}_n)$$
$$p(\mathbf{z}_{1:n}) = p(\mathbf{z}_n | \mathbf{z}_{1:n-1}) p(\mathbf{z}_{1:n-1})$$

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Bayes Rule	just keeps g	iving		

Plugging in known  $f(\cdot)$  and  $g(\cdot)$ :

$$p(\mathbf{x}_{0:n}|\mathbf{z}_{1:n}) = p(\mathbf{x}_{0:n-1}, \mathbf{z}_{1:n-1}) \frac{f(\mathbf{x}_n|\mathbf{x}_{n-1})g(\mathbf{z}_n|\mathbf{x}_n)}{p(\mathbf{z}_n|\mathbf{z}_{1:n-1})p(\mathbf{z}_{1:n-1})}$$

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Replace  $p(\mathbf{x}_{0:n-1}, \mathbf{z}_{1:n-1})$  with it's conditional:

$$p(\mathbf{x}_{0:n}|\mathbf{z}_{1:n}) = p(\mathbf{x}_{0:n-1}|\mathbf{z}_{1:n-1}) \frac{p(\mathbf{z}_{1:n-1})f(\mathbf{x}_n|\mathbf{x}_{n-1})g(\mathbf{z}_n|\mathbf{x}_n)}{p(\mathbf{z}_n|\mathbf{z}_{1:n-1})p(\mathbf{z}_{1:n-1})}$$

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The  $p(\mathbf{z}_{1:n-1})$  distributions cancel out...

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Wrapping	g up Bayes F	Rule		

### Thus:

$$p(\mathbf{x}_{0:n}|\mathbf{z}_{1:n}) = p(\mathbf{x}_{0:n-1}|\mathbf{z}_{1:n-1}) \frac{f(\mathbf{x}_{n}|\mathbf{x}_{n-1})g(\mathbf{z}_{n}|\mathbf{x}_{n})}{p(\mathbf{z}_{n}|\mathbf{z}_{1:n-1})}$$

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Wrapping	g up Bayes F	Rule		

Thus:

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We know from normalization rules that:

$$p(\mathbf{z}_n|\mathbf{z}_{1:n-1}) = \int p(\mathbf{x}_{0:n-1}|\mathbf{z}_{1:n-1})f(\mathbf{x}_n|\mathbf{x}_{n-1})g(\mathbf{z}_n|\mathbf{x}_n)d\mathbf{x}_{n-1:n}$$

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Let  $C_n$  be the integral above. Then...

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Things just	got Recursiv	ve		

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• Our state distribution is recursive!

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- Our state distribution is recursive!
- Recall: we have the base case  $p(\mathbf{x}_0)$

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Things in	ist oot Recur	sive		

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- Our state distribution is recursive!
- Recall: we have the base case  $p(\mathbf{x}_0)$
- Distributions  $f(\cdot)$  and  $g(\cdot)$  are also known

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Things just got Recursive						

$$\rho(\mathbf{x}_{0:n}|\mathbf{z}_{1:n}) = \rho(\mathbf{x}_{0:n-1}|\mathbf{z}_{1:n-1}) \frac{f(\mathbf{x}_n|\mathbf{x}_{n-1})g(\mathbf{z}_n|\mathbf{x}_n)}{C_n}$$

- Our state distribution is recursive!
- Recall: we have the base case  $p(\mathbf{x}_0)$
- Distributions  $f(\cdot)$  and  $g(\cdot)$  are also known
- We have everything we need to solve this problem!



In some cases, we can get  $p(\mathbf{x}_{0:n}|\mathbf{z}_{1:n})$  analytically:

e.g. Kalman filters
 [p(x<sub>0</sub>) is Gaussian, f(x<sub>t</sub>|x<sub>t-1</sub>) and g(z<sub>t</sub>|x<sub>t</sub>) are linear ]



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In most cases, we can't:

- Integration in C<sub>n</sub> intractable
- Often,  $p(\mathbf{x}_0)$  not known, estimated with simpler distribution.



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In most cases, we can't:

- Integration in *C<sub>n</sub>* intractable
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The Solution:

• Estimate  $p(\mathbf{x}_{0:n}|\mathbf{z}_{1:n})$  from samples (i.e. particles).

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Monte Ca	arlo Estimate	es		

Traditional Monte Carlo:

• Generate N i.i.d. samples  $\{x_{0:n}^1, \cdots, x_{0:n}^N\}$  from  $p(\mathbf{x}_{0:n}|\mathbf{z}_{1:n})$ 



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$$\hat{p}(\mathbf{x}_{0:n}|\mathbf{z}_{1:n}) = \frac{1}{N}\sum_{i=1}^{N}\delta(x_{0:n}^{i})$$



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Issues:

• p(x<sub>0:n</sub>|z<sub>1:n</sub>) only known "recursively"



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Issues:

- *p*(**x**<sub>0:n</sub>|**z**<sub>1:n</sub>) only known "recursively"
- Base case p(x<sub>0</sub>) may not be exactly known

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Importance	Sampling (I	S)		

- Sample from a simple distribution  $\pi(\mathbf{x}_0)$ 
  - (e.g. Uniform, Gaussian)

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Importance	Sampling (I	S)		

- Sample from a simple distribution π(x<sub>0</sub>) (e.g. Uniform, Gaussian)
- Weight each sample by sampling error:

$$w(\mathbf{x}_0^i) = rac{p(\mathbf{x}_0^i)}{\pi(\mathbf{x}_0^i)}$$

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Obtain weighted discrete approximation of p(x<sub>0</sub>):

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Sequential	Importance S	Sampling (S	IS)	

How do we know sampling error in our case?

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## Sequential Importance Sampling (SIS)

How do we know sampling error in our case?

- We have our observations!  $\{z_1, \dots, z_n\}$
- As our sampled states x<sup>i</sup><sub>t</sub> change across time, measure sampling error using

$$w_t^i = g(\mathbf{z}_t | \mathbf{x}_t)$$



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Introduction	HMM 000000000	Sampling ooooo●oo	Re-sampling	Genetic Algs.
Particle F	iltering usin	a SIS		

• Generate *N* i.i.d. "particles"  $\{x_0^1, \dots, x_0^N\}$  from  $\hat{p}(\mathbf{x}_0)$ 

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- Generate *N* i.i.d. "particles"  $\{x_0^1, \dots, x_0^N\}$  from  $\hat{p}(\mathbf{x}_0)$ 
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- 2 Propagate particles to new state using  $f(\mathbf{x}_t | \mathbf{x}_{t-1})$

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- Siven observation  $\mathbf{z}_t$ , determine particles' weight  $w_t^i = g(\mathbf{z}_t | \mathbf{x}_t)$
- Update weight  $w^i = w_{t-1}^i w_t^i$ .

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- Sepeat steps 2-4 *n* times.

- Generate N i.i.d. "particles"  $\{x_0^1, \dots, x_0^N\}$  from  $\hat{p}(\mathbf{x}_0)$ 
  - $\hat{p}(\mathbf{x}_0)$  may be very simple (e.g. Uniform, Gaussian)
  - Let  $w^i = w_0^i = \frac{1}{N}$ , for all particles.
- 2 Propagate particles to new state using  $f(\mathbf{x}_t | \mathbf{x}_{t-1})$
- 3 Given observation  $\mathbf{z}_t$ , determine particles' weight  $w_t^i = g(\mathbf{z}_t | \mathbf{x}_t)$
- Update weight  $w^i = w_{t-1}^i w_t^i$ .
- Sepeat steps 2-4 *n* times.
- **(b)** Obtain *weighted* discrete approximation  $\hat{p}(\mathbf{x}_{0:n}|\mathbf{z}_{1:n})$ .

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Use SIS with Caution				

The "gotchas":

• Doesn't work well in high-dimensions (need large *N*).



The "gotchas":

- Doesn't work well in high-dimensions (need large *N*).
- Degeneracy: After a few timesteps, all but one particle will have negligible weight



Figure 2: (A) The weights of all 50 particles (x-axis) at each time step k (y-axis). Hotter colors represent larger weights. (B) The effective sample size  $N_{eff}$  as a function of time step k.

Introduction	HMM 00000000	Sampling ○○○○○○●	Re-sampling	Genetic Algs.
Outline				

- Introduction: What is particle filtering?
- Particle Filters and Hidden Markov Models
- Sampling Particles
- Re-sampling Particles
- Particle Filters & Genetic Algorithms

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Addressing	Degeneracy			

Detecting degeneracy:

• At each timestep t, measure effective sample size

$$N_{eff} = \frac{1}{\sum_{i=1}^{N} (w^i)^2}$$



Figure 2: (A) The weights of all 50 particles (x-axis) at each time step k (y-axis). Hotter colors represent larger weights. (B) The effective sample size  $N_{eff}$  as a function of time step k.



If  $N_{eff} < \tau$  at timestep *t*:

- Sample *N* new particles  $\{x_t^1, \dots, x_t^N\}$  from  $\hat{p}(\mathbf{x}_{0:t}|\mathbf{z}_{1:t})$
- Reset weights:  $w^i = \frac{1}{N}$



Figure 2: (A) The weights of all 50 particles (x-axis) at each time step k (y-axis). Hotter colors represent larger weights. (B) The effective sample size  $N_{eff}$  as a function of time step k.



#### Sequential Importance Resampling (SIR)



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## Pros and Cons of SIR Particle Filtering

#### Pros:

- Estimation of full state PDFs
- No assumptions on distributions
- Parallelizable

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## Pros and Cons of SIR Particle Filtering

#### Pros:

- Estimation of full state PDFs
- No assumptions on distributions
- Parallelizable

#### Cons:

- Degeneracy possible
- Good estimates may need large *N*
- Computationally expensive

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Introduction	HMM	Sampling	Re-sampling	Genetic Alas		

More General Particle Filtering

Consider optimization problems:

$$x^* = \arg\min_x E(x)$$

Often solved with Euler-Lagrange

- Introduce artificial timestep t
- Compute  $\frac{\partial x}{\partial t}$
- Update using gradient descent  $x_t = x_{t-1} \delta t \frac{\partial x}{\partial t}$

Looks similar...

Introduction	HMM 00000000	Sampling 00000000	Re-sampling	Genetic Algs. oo●o
Particle Filte	ering Optimiz	zation		

• Generate *N* i.i.d. solutions  $\{x_0^1, \dots, x_0^N\}$  from  $\hat{p}(\mathbf{x}_0)$ 

Introduction	HMM 00000000	Sampling 00000000	Re-sampling	Genetic Algs. ○○●○
Particle F	ilterina Opti	mization		

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  - $\hat{p}(\mathbf{x}_0)$  may be very simple (e.g. Uniform, Gaussian)

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- Repeat steps 2-5 until convergence.
- Select  $x_n^i$  with largest weight  $w^i$  as  $x^*$ .

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Particle Filtering = Genetic Algorithms?								

So what's going on?

- Generating a PDF of  $e^{-E(x)}$  using particle filtering.
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#### Particle filtering is for more than HMM problems.