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ORGANIZATION

- Introduction
- Background into Bayesian approach
- Monte Carlo (MC) methods for Bayesian inference
- Sequential Bayesian processor (SBP)
- Model-based signal processing
- Bayesian approach to state-space processors
- Simulation-based MC approach to SBP
- Particle filtering for SBP
- Performance analysis
- Applications (nonlinear/multi-modal)
- Summary



Bayesian signal processing

 is concerned with the estimation of the underlying probability distribution of a random signal in order to perform statistical inferences, that is,



 statistical inferences enable the extraction of the signal from noisy uncertain measurement data. For example, the maximum aposterior (MAP)

$$\hat{\Pr}[X | Y] \implies \hat{X} = \max_{X} \hat{\Pr}[X | Y]$$

Bayesian techniques are based on BAYES' rule:



This "simple relationship" is the **principal foundation** of Bayesian signal processing both theoretically (derivations) and pragmatically (implementations)

Bayesian techniques convert the prior \rightarrow posterior



PARTICLE FILTERS (PF)

- PF are sequential (Monte Carlo) techniques in which the underlying posterior distribution of interest is represented by a "cloud" of random samples (particles)
- The PF is a processing algorithm that (sequentially) propagates and updates the random samples drawn from the previous stage to obtain a set of samples approximately distributed from the next stage

Particle filters can be used to construct the "empirical" <u>posterior distributions</u>. For example, a <u>static</u> parameter X given the data Y



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Particle filters can be applied in many areas:

Signal processing

- Image processing and segmentation
- Model selection
- Tracking and navigation

Communications

- Channel estimation
- Blind equalization
- Positioning in wireless networks

Applications

- Biology & Biochemistry
- Chemistry
- Economics & Business
- Genomics
- Geosciences
- Immunology
- Materials Science
- Pharmacology & Toxicology
- Psychiatry/Psychology
- Social Sciences

PFs: advantages and disadvantages*:

ADVANTAGES

Ability to represent arbitrary densities

Adaptive focusing on probable regions of state-space

Dealing with non-Gaussian noise

The framework allows for including multiple models (tracking maneuvering targets)

DISADVANTAGES

- High computational complexity
- It is difficult to determine optimal number of particles
- Number of particles increase with increasing model dimension
- Potential problems: degeneracy and loss of diversity
- The choice of importance density is crucial

* M. Bolic, University of Ottawa, "Theory and Implementation of Particle Filters"

Our plan is to develop particle filters using the following roadmap:



BACKGROUND

Monte Carlo (MC) approach to nonparametric "posterior" probability distribution estimation

- MC techniques obtain <u>random sample based representations</u> of probability distributions to solve a variety of estimation problems, especially those in which a "closed-form" analytic expression is not available, that is, usual is distribution —> samples, MC samples —> distribution
- The MC approach enables complex, multi-modal distributions to be estimated
- Many nonlinear, non-gaussian processing problems can easily be cast into this framework
- When statistics are "nonstationary," then sequential MC methods must be employed to solve the problem
- Particle filters are a "sequential" MC method that can be applied in the nonstationary and/or real-time cases as well as the stationary case

MONTE CARLO (Sampling) METHODS

In most cases (non-gaussian), the posterior distribution is **not** available requiring that we use "simulation-based" or non-parametric methods for estimation and inference

- These techniques are called **SAMPLING METHODS**
- They replace numerical integration techniques
- Monte Carlo (MC) sampling methods are an alternative
- The KEY idea in MC is replacing the distribution function by its SAMPLES reducing integration to summations
- As the number of samples increase the equivalence becomes established with the targeted posterior distribution enabling its moments or estimators to be calculated, that is,

 $MC \text{ integration evaluates} \qquad \text{by drawing}$ samples, $\{X(i)\}$ from $\Pr(X)$ and assuming perfect sampling produces the estimate $\hat{\Pr}(X) \approx \frac{1}{N} \sum_{i=1}^{N} \delta(X - X(i))$

Typically integration is used to estimate expectations, but in MC the "inverse" is true

$$I = \int_X g(x) dx$$

then MC methods factorize the integrand as,

$$g(x) \longrightarrow f(x)p(x) \quad \ni p(x) \ge 0 \text{ and } \int p(x)dx = 1$$

where p(x) is interpreted as a probability distribution in which samples can be drawn. This is the foundation of sampling techniques based on *MC* integration. Monte Carlo integration draws samples from the required distribution and then forms sample averages to approximate the sought after distributions, that is, it maps integrals to discrete sums. Thus, *MC* integration evaluates by drawing samples, $\{X(i)\}$ from Pr(X) and assuming perfect sampling produces the estimate

$$\hat{\Pr}(X) \approx \frac{1}{N} \sum_{i=1}^{N} \delta(X - X(i))$$
 This is the KEY to MC techniques

which upon substituting into the integral gives

$$E_X\{f(X)\} = \int f(X)\hat{\Pr}(X)dX \approx \frac{1}{N}\sum_{i=1}^N f(X(i)) \equiv \overline{f}$$

Here \overline{f} is said to be a *Monte Carlo estimate* of $E_X\{f(X)\}$. Clearly, it is *unbiased*, since

Consider a gaussian example: $x \sim N(m_x, \sigma_x^2)$

the Monte Carlo approach is to generate N samples from a gaussian, that is, assuming perfect (uniform) sampling, we have that

$$\hat{\Pr}(x) \approx 1/N \sum_{i=1}^{N} \delta(x - x_i)$$

and therefore the mean and variance can be estimate from the samples directly using

$$\hat{m}_x = \int x \hat{\Pr}(x) \, dx = \int x \left(1/N \sum_{i=1}^N \delta(x - x_i) \right) dx = 1/N \sum_{i=1}^N x_i$$

which follows directly from the sifting property of the delta function. Also, we have

$$\hat{\sigma}_x^2 = \int (x - m_x)^2 \operatorname{Pr}(x) \, dx = \int (x - m_x)^2 \left(1/N \sum_{i=1}^N \delta(x - x_i) \right) dx = 1/N \sum_{i=1}^N (x_i - \hat{m}_x)^2$$

Thus, summarizing the MC approach is:

- Generate N samples from a gaussian: $x_i \sim \mathcal{N}(m_x, \sigma_x^2)$;
- Estimate the desired statistics of the distribution from its samples as: \hat{m}_x and $\hat{\sigma}_x^2$

<u>Monte Carlo methods</u> rely on samples generated from the sampling distribution to estimate statistics (mean, variance, etc.) as demonstrated in this Gaussian example





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SEQUENTIAL BAYESIAN PROCESSOR (SBP)

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Bayes' rule for dynamic variables is:

$$\Pr(X|Y) = \Pr(Y|X) \frac{\Pr(X)}{\Pr(Y)}$$

$$\Pr(X_t|Y_t) = \frac{\Pr(Y_t|X_t) \times \Pr(X_t)}{\Pr(Y_t)}$$
$$\begin{bmatrix} X_t \coloneqq \{x(0), \dots, x(t)\}; & Y_t \coloneqq \{y(0), \dots, y(t)\} \end{bmatrix}$$

The posterior distribution can be estimated using the sequential Bayesian processor (SBP):

$$\Pr[\mathbf{X}_{o}|\mathbf{Y}_{o}] \implies \Pr[\mathbf{X}_{1}|\mathbf{Y}_{1}] \implies \cdots \implies \Pr[\mathbf{X}_{t-1}|\mathbf{Y}_{t-1}] \implies \Pr[\mathbf{X}_{t}|\mathbf{Y}_{t}]$$

$$\longrightarrow W(1,0) \implies \cdots \implies W(t-1,t-2) \longrightarrow W(t,t-1)$$

$$\Pr[\mathbf{X}_{t} | \mathbf{Y}_{t}] = W(t, t-1) \times \Pr[\mathbf{X}_{t-1} | \mathbf{Y}_{t-1}]$$

where the Bayes' operator is defined at each stage by
$$W(t, t-1) \coloneqq \frac{\Pr[\mathbf{y}(t) | \mathbf{x}(t)] \times \Pr[\mathbf{x}(t) | \mathbf{x}(t-1)]}{\Pr[\mathbf{y}(t) | \mathbf{Y}_{t-1}]}; \quad t = 1, \dots, N$$

The final FILTERING recursion evolves from Bayes' rule

we	obtain the final expression for the correction recursion as
	$\underbrace{\underbrace{\Pr(x(t) Y_t)}_{Pr(x(t) Y_t)} = \underbrace{\Pr(y(t) x(t))}_{Likelihood} \times \underbrace{\Pr(x(t) Y_{t-1})}_{Pr(x(t) Y_{t-1})}}_{Evidence}$
where we can consider the correction or filtering distribution as a weighting of the prediction distribution as in the full joint case above, that is,	
	$\underbrace{\widetilde{\Pr(x(t) Y_t)}}_{\operatorname{Pr}(x(t) Y_t)} = \underbrace{\widetilde{\mathcal{W}_c(t,t-1)}}_{\operatorname{W}_c(t,t-1)} \times \underbrace{\widetilde{\Pr(x(t) Y_{t-1})}}_{\operatorname{Pr}(x(t) Y_{t-1})}$

The SBP for the "filtering" posterior is shown by:



Predict:
$$\Pr[x(t) | Y_{t-1}] = \int \Pr[x(t) | x(t-1)] \Pr[x(t-1) | Y_{t-1}] dx(t-1)$$
$$Update: \qquad \Pr[x(t) | Y_t] = W_c(t, t-1) \times \Pr[x(t) | Y_{t-1}]$$
$$W_c(t, t-1) := \left[\frac{\Pr[y(t) | x(t)]}{\Pr[y(t) | Y_{t-1}]}\right]$$

MODEL-BASED (Bayesian) SIGNAL PROCESSING

Model-based signal processing:

• Signal processing is the set of techniques to "extract the useful information from noisy measurements while rejecting the extraneous"

• If the SNR is high then simple non-physical techniques (FFTs, wavelets, spectral estimation, etc.) can be used to extract the desired information, BUT . . .

• If the SNR is low and the problem uncertain (noise, random parameters, etc.), then more of the underlying physical phenomenology must somehow be incorporated into the processor

THEREFORE

• Model-based signal processing incorporates physical phenomenological, measurement, and noise/uncertainty models into the processor to extract the desired information while rejecting the extraneous --- even in highly uncertain environments

The model-based approach is simply:

"incorporating mathematical models of both physical phenomenology and the measurement processes including noise into the processor to extract the desired information" MBP are techniques that incorporate any "a priori" knowledge of the underlying phenomenology into a processing scheme



Estimate

In bio-threats, MBP techniques can incorporate any "a priori" knowledge of the underlying physics into the processing scheme



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For this problem, *smart bio-sensors* incorporating a microcantilever array can be developed using the MBP approach



State-space is a natural (mathematical) representation of a system (of equations)

For ease of development, let us assume that we have a set of n^{th} order ordinary differential equations (ODE) with constant coefficients in continuous-time ($t \in R$), that is,

$$\frac{d^{n}}{dt^{n}}z_{t} + a_{n-1}\frac{d^{n-1}}{dt^{n-1}}z_{t} + \dots + a_{1}\frac{d}{dt}z_{t} + a_{0}z_{t} = u_{t}$$

where u_t is the input or excitation function.

The usual method of solving these equations, computationally, is to rewrite them into an equivalent set of <u>n</u>-first order differential equations by defining a vector with components corresponding to the differentials, that is, $x_i := \frac{d^i z_t}{dt^i}$ for $i = 0, \dots, n-1$ or

$$x_t := \left[z_t \ \frac{dz_t}{dt} \ \cdots \ \frac{d^{n-1}z_t}{dt^{n-1}} \right]'$$

State-space is a natural representation:

Taking the derivative of x_t for each component, we obtain

$$\dot{x}_1 = \frac{dz_t}{dt} = x_2$$

$$\dot{x}_2 = \frac{d^2 z_t}{dt^2} = x_3$$

$$\vdots$$

$$\dot{x}_n = \frac{d^n z_t}{dt^n} = -a_{n-1}\frac{d^{n-1}z_t}{dt^{n-1}} + \dots - a_1\frac{dz_t}{dt} - a_0z_t + u_t$$

Collecting all of these equations and writing them into vector-matrix form, we obtain

$$\frac{d}{dt} \begin{bmatrix} x_1 \\ \vdots \\ x_{n-1} \\ x_n \end{bmatrix} = \begin{bmatrix} 0 & 1 & \cdots & 0 \\ \vdots & \vdots & \vdots & \vdots \\ 0 & 0 & \cdots & 1 \\ -a_0 & -a_1 & \cdots & -a_{n-1} \end{bmatrix} \begin{bmatrix} x_1 \\ \vdots \\ x_{n-1} \\ x_n \end{bmatrix} + \begin{bmatrix} 0 \\ \vdots \\ 0 \\ 1 \end{bmatrix} u_t$$

or in compact form

$$\dot{x}_t = Ax_t + Bu_t$$

for $x_t \in R^n$, $u_t \in R^{1 \times 1}$, $A \in R^{n \times n}$ and $B \in R^{n \times 1}$. Here x_t is defined as the *state vector*, u_t the input, A is the system or process matrix and B is the input matrix. So we see from this simplistic view, the state vector is "naturally" obtained when we convert an n^{th} order ODE to a set of n-first order equations.

STATE-SPACE MODEL

The *state* of a system at time t is the "minimum" set of variables (*state variables*) along with the *input* sufficient to uniquely specify the dynamic system behavior for all t over the interval $t \in [t, \infty)$. The *state vector* is the collection of state variables into a single vector. The idea of a *minimal set* of state variables is critical and all techniques to define them must assure that the smallest number of "independent" states have been defined. If not, there are some serious system theoretic issues that could arise [2], [3].

Let us consider a general *deterministic* formulation of a <u>nonlinear dynamic</u> system including the output (measurement) model in state-space form (continuous-time)²

$$\dot{x}_t = \mathcal{A}(x_t, u_t) = a(x_t) + b(u_t)$$

$$y_t = \mathcal{C}(x_t, u_t) = c(x_t) + d(u_t)$$

$$\longleftarrow \quad Nonlinear$$

for x_t , y_t and u_t the respective N_x -state, N_y -output and N_u -input vectors with corresponding system (process), input, measurement (output) and feedthrough functions. The N_x -dimensional system and input functions are defined by $a(\cdot)$ and $b(\cdot)$, while the N_y -dimensional output and feed through functions are given by $c(\cdot)$ and $d(\cdot)$.

In order to specify the solution of the N_x -th order differential equations completely, we must specify the above noted functions along with a set of N_x -initial conditions at time t_0 and the input for all $t \ge t_0$. Here N_x is the "minimal" set of state variables.

If we constrain the state-space representation to be linear in the states, then we obtain the generic continuous-time, *linear time-varying state-space* model given by

$$\dot{x}_t = A_t x_t + B_t u_t$$

 $y_t = C_t x_t + D_t u_t$ \leftarrow Linear .1)

where $x_t \in \mathcal{R}^{N_x \times 1}$, $u_t \in \mathcal{R}^{N_u \times 1}$, $y_t \in \mathcal{R}^{N_y \times 1}$ and the respective system, input, output and feed through matrices are: $A \in \mathcal{R}^{N_x \times N_x}$, $B \in \mathcal{R}^{N_x \times N_u}$, $C \in \mathcal{R}^{N_y \times N_x}$ and $D \in \mathcal{R}^{N_y \times N_u}$.

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BAYESIAN APPROACH TO STATE-SPACE PROCESSORS

Bayesian approach to the state-space: definitions

$$\begin{aligned} x(t) &= & \mathcal{A} \left(x(t-1), u(t-1), w(t-1) \right) \\ y(t) &= & \mathcal{C} \left(x(t), u(t), v(t) \right) \end{aligned}$$

where w and v are the respective process and measurement noise sources with u a known input. Here $\mathcal{A}(\cdot)$ is the nonlinear (or linear) dynamic state transition function and $\mathcal{C}(\cdot)$ the corresponding measurement function. Both conditional probabilistic distributions embedded within the Bayesian framework are *completely* specified by these functions and the underlying noise distributions: $\Pr(w(t-1))$ and $\Pr(v(t))$. That is, we have the equivalence

$$\mathcal{A} \left(x(t-1), u(t-1), w(t-1) \right) \implies \Pr\left(x(t) | x(t-1) \right) \Leftrightarrow \mathcal{A} \left(x(t) | x(t-1) \right) \\ \mathcal{C} \left(x(t), u(t), v(t) \right) \implies \Pr\left(y(t) | x(t) \right) \qquad \Leftrightarrow \mathcal{C} \left(y(t) | x(t) \right)$$

Bayesian approach to the state-space: posteriors

the prediction recursion characterized by the *Chapman-Kolmogorov equation* replacing transition probability with the implied model-based conditional, that is,

 $\Pr(x(t)|Y_{t-1}) = \int \underbrace{\mathcal{A}(x(t)|x(t-1))}_{\mathbf{X}(t-1)} \times \Pr(x(t-1)|Y_{t-1})dx(t-1)$

Next we incorporate the model-based likelihood into the posterior equation with the understanding that the process model has been incorporated into the prediction

Embedded Measurement Model $\Pr(x(t)|Y_t) = \frac{\mathcal{C}(y(t)|x(t))}{\mathcal{C}(y(t)|x(t))} \times \Pr(x(t)|Y_{t-1}) / \Pr(y(t)|Y_{t-1})$

The BSP implemented in state-space is:



EXAMPLE: Given a Gauss-Markov model in state-space form:



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GM EXAMPLE: Develop a SBP where: $\hat{x}(t | t-1) := E \{x(t) | Y_{t-1}\}; \tilde{P}(t | t-1) := \operatorname{cov}(x(t) - \hat{x}(t | t-1))$

$$\Pr(x(t)|Y_t) = \frac{\mathcal{C}(y(t)|x(t))\Pr(x(t)|Y_{t-1})}{\Pr(y(t)|Y_{t-1})}$$

Under the Gauss-Markov model assumptions, we know that each of the conditional distributions can be expressed in terms of the gaussian distribution as:

$$\begin{array}{c} \mathcal{C}\left(y(t)|x(t)\right) & \sim & \mathcal{N}\left(y(t):C(t)x(t), R_{vv}(t)\right) \\ \mathbf{Pr}\left(x(t)|Y(t-1)\right) & \sim & \mathcal{N}\left(x(t):\hat{x}(t|t-1), \tilde{P}(t|t-1)\right) \\ \mathbf{Pr}\left(y(t)|Y(t-1)\right) & \sim & \mathcal{N}\left(y(t):\hat{y}(t|t-1), R_{ee}(t)\right) \end{array}$$

Substituting these probabilities into a single constant κ , we obtain and combining all constants into a

$$\begin{aligned} & \bigvee \\ \Pr(x(t)|Y(t)) &= \kappa \times \exp\left[-\frac{1}{2}(y(t) - C(t)x(t))'R_{vv}^{-1}(t)(y(t) - C(t)x(t))\right] \\ & \times & \exp\left[-\frac{1}{2}(x(t) - \hat{x}(t|t-1))'\tilde{P}^{-1}(t|t-1)(x(t) - \hat{x}(t|t-1))\right] \\ & \times & \exp\left[+\frac{1}{2}(y(t) - \hat{y}(t|t-1))'R_{ee}^{-1}(t)(y(t) - \hat{y}(t|t-1))\right] \end{aligned}$$

Substituting for the individual variables gives the well-known optimal (GM) KALMAN filter algorithm as:



When linear state-space models are employed, the Bayesian solution is the "sequential" MODEL-BASED (Kalman filter) algorithm which as the following structure:



SIMULATION-BASED MC APPROACH TO SEQUENTIAL BAYESIAN PROCESSORS

Importance sampling is a generalization to the MC approach based on the *sampling distribution* that samples the targeted posterior, that is, the sampling distribution is:

$$I = \int_X g(x) dx = \int_X \left(\frac{g(x)}{q(x)}\right) \times q(x) \ dx \quad \text{ for } \int q(x) dx = 1$$

- a simpler distribution, q(x), than the "posterior" and easier to draw samples
- based on Markov chain theory and therefore will converge to the posterior
- provides non-uniform sampling of the target, g(x), giving "more importance" to some values than others
- it is said that the support of q(x) "covers" that of g(x), i.e., samples drawn from q overlap the same region of samples from g

This leads to the sequential importance algorithm:

These results then enable us to formulate a generic Bayesian *sequential importance sampling* algorithm:

- → 1. Choose samples from the proposed importance distribution: $x_i(t) \sim q(x(t)|X_{t-1}, Y_t)$
- \rightarrow 2. Determine the required conditional distributions $\Pr(x_i(t)|x(t-1)), \Pr(y(t)|x_i(t))$
- → 3. Calculate the unnormalized weights: $W_i(t)$ with $x(t) \rightarrow x_i(t)$;
- \rightarrow 4. Normalize the weights: $\mathcal{W}_i(t)$ and
 - → 5. Estimate the posterior distribution: $\hat{\Pr}(X_t|Y_t) = \sum_{i=1}^N \mathcal{W}_i(t)\delta(x(t) x_i(t))$

State-space Bayesian processors based on sequential importance samplers follow easily as:

$$W(t) = W(t-1) \times \frac{\Pr(y(t)|x(t)) \times \Pr(x(t)|x(t-1))}{q(x(t)|X_{t-1}, Y_t)}$$

Now let us recall the general state-space characterization representing the transition and likelihood probabilities as:

$$\frac{\Pr(x(t)|x(t-1))}{\Pr(y(t)|x(t))} \Leftrightarrow \mathcal{A}(x(t)|x(t-1))$$

Assuming this is true, then the SSPF recursion becomes

$$\begin{aligned} x_i(t) &\sim q(x(t)|x(t-1), y(t)) \\ W_i(t) &= W_i(t-1) \times \frac{\mathcal{C}(y(t)|x_i(t)) \times \mathcal{A}(x_i(t)|x_i(t-1)))}{q(x_i(t)|x_i(t-1), y(t))} \\ \mathcal{W}_i(t) &= \frac{W_i(t)}{\sum_{i=1}^{N_p} W_i(t)} \end{aligned}$$

and the filtering posterior is estimated by

$$\hat{\Pr}(x(t)|Y_t) \approx \sum_{i=1}^{N_p} \mathcal{W}_i(t) \times \delta(x(t) - x_i(t))$$

Note that as N_p becomes large, in the limit, we have

$$\lim_{N_p \to \infty} \hat{\Pr}(x(t) | Y_t) \longrightarrow \Pr(x(t) | Y_t)$$

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Importance distributions provide the key: "Transition Prior" (Gordon et. al. '93)

Another choice for an importance distribution is the *transition prior*. This *prior* is defined in terms of the state-space representation by $\mathcal{A}(x(t)|x(t-1)) \rightarrow \mathcal{A}(x(t-1), u(t-1), w(t-1))$ which is <u>dependent</u> on the known excitation and process noise statistics. It is given by

$$q_{prior}(x(t)|x(t-1), Y_t) \longrightarrow \Pr(x(t)|x(t-1))$$

Substituting this choice into the weights gives

$$W_{i}(t) = W_{i}(t-1) \times \frac{\Pr(y(t)|x_{i}(t)) \times \Pr(x(t)|x_{i}(t-1)))}{q_{prior}(x(t)|x_{i}(t-1), Y_{t})} = W_{i}(t-1) \times \Pr(y(t)|x_{i}(t))$$

BOOTSTRAP ESTIMATOR

PARTICLE FILTERS FOR SBP

PARTICLE FILTERS (PF)

- PF are sequential MC techniques in which the underlying posterior distribution of interest is represented by a "cloud" of random samples (particles) in the state/parameter space
- The PF is an algorithm that (sequentially) propagates and updates the random samples drawn from the previous stage to obtain a set of samples approximately distributed from the next stage, that is,

$$\Pr[X_{t} | \mathbf{Y}_{t}] = W_{i}(t, t-1; X_{i}(t)) \times \Pr[X_{t-1} | \mathbf{Y}_{t-1}]$$

where $W_{i}(t, t-1; X_{i}(t))$ is the weight (Bayes' operator) defined earlier and $X_{i}(t)$ is the i^{th} – particle at stage (time) t

Previous Stage

Particle filters can be used to construct <u>posterior distributions</u>. In the dynamic case they are used to estimate the "instantaneous" posterior



Particle No. (Time-Step 68)



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The "generic" State-Space particle filtering method is given by:



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<u>**PROBLEM</u>:** Particles deplete in number (degenerate) to a single particle due to the increased variance in each step; therefore,</u>

- The particles must be "rejuvenated" or equivalently resampled
- Resampling inhibits the depletion problem, but increases the uncertainty (weight variance)
- If not implemented properly, it can also increase computational time extensively (non-parallel)
- Resampling is essentially a process that attempts to preserve particles with large weights (acceptance probabilities) while discarding those with small weights.





Particle Filtering: Examples, Applications

SUMMARY

- Many problems can be cast into a Bayesian framework in order to solve a suite of problems
- No longer are we restricted to nonlinear approximations and gaussian processes
- Sequential Bayesian processors evolve as natural extensions which incorporate Markovian state-space structures
- Problem solutions for highly uncertain, noisy measurements using <u>physics-based</u> signal processing models and <u>sequential</u> <u>Bayesian</u> processing implemented with <u>particle filtering</u> techniques add a new tool for signal processors

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