

Improved particle filter for nonlinear problems

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Abstract: The Kalman filter provides an effective solution to the linear Gaussian filtering problem. However where there is nonlinearity, either in the model specification or the observation process, other methods are required. Methods known generically as 'particle filters' are considered. These include the condensation algorithm and the Bayesian bootstrap or sampling importance resampling (SIR) filter. These filters represent the posterior distribution of the state variables by a system of particles which evolves and adapts recursively as new information becomes available. In practice, large numbers of particles may be required to provide adequate approximations and for certain applications, after a sequence of updates, the particle system will often collapse to a single point. A method of monitoring the efficiency of these filters is introduced which provides a simple quantitative assessment of sample impoverishment and the authors show how to construct improved particle filters that are both structurally efficient in terms of preventing the collapse of the particle system and computationally efficient in their implementation. This is illustrated with the classic bearings-only tracking problem.

1 Introduction

The Bayesian approach to dynamic state estimation problems involves the construction of the probability density function (PDF) of the current state of an evolving system, given the accumulated observation history. For linear Gaussian models where the PDF can be summarised by means and covariances, the calculation is carried out in terms of the familiar updating equations of the Kalman filter. In general, for nonlinear, non-Gaussian models, there is no simple way to proceed. Two difficulties must be resolved: how to represent a general PDF using finite computer storage and how to perform the integrations involved in updating the PDF when new data are acquired.

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Several approximate methods have been proposed. These include the extended Kalman filter [1, 2], the Gaussian sum filter [3], approximating the first two moments of the PDF [4, 5] and numerical integration over a grid of points in the state space [6–10]. However, none of these methods can be applied automatically. Typically they have to be tuned to take account of features of each specific problem. For example, in grid-based methods the number and location of the grid points has to be decided, usually by a process of trial and error. Furthermore, updating the distribution of the state of the system as new data arrive usually entails a formidable computational overhead.

There is now a substantial literature concerned with simulation based filters in which the required PDF is represented by a scatter of particles which propagate through the state space [9, 11–19]. The propagation and adaptation rules are chosen so that the combined weight of particles in a particular region will approximate the integral of the PDF over the region. Such filters have been variously described as Bayesian bootstrap, condensation, Monte Carlo and Metropolis–Hastings importance resampling filters. We adopt the term Monte Carlo particle filter or particle filter for short. Gordon, Salmond and Smith [14] demonstrate the effectiveness of a simple algorithm for particle evolution for various nonlinear filtering applications. Their method has become known as the sampling importance resampling (SIR) filter or, more commonly in the engineering literature, the Bayesian bootstrap filter.

The standard SIR filter is vulnerable to sample impoverishment [17, 20–22], so that the particle distribution gives a poor approximation of the required PDF. In extreme cases, after a sequence of updates the particle system can collapse to a single point. In less extreme cases, although several particles may survive, there is so much internal correlation that summary statistics behave as if they are derived from a substantially smaller sample. To compensate, large numbers of particles are required in realistic problems.

We show how sample impoverishment can be quantified. We introduce modifications demonstrated to have superior performance to the SIR filter both in terms of combating sample impoverishment and in computational cost.

2 Bayesian filtering

Following Gordon, Salmond and Smith [14] we represent the state vector at time k by $x_k \in \mathbb{R}^n$, which satisfies

$$x_{k+1} = f_k(x_k, w_k)$$

where $f_k : \mathbb{R}^n \times \mathbb{R}^m \rightarrow \mathbb{R}^n$ is the system transition function and w_k is a noise term whose known distribution is independent of time. At each discrete time point an

observation $y_k \in \mathbb{R}^p$ is obtained, related to the state vector by

$$y_k = h_k(x_k, v_k)$$

where $h_k : \mathbb{R}^n \times \mathbb{R}^r \rightarrow \mathbb{R}^p$ is the measurement function and $v_k \in \mathbb{R}^r$ is another noise term whose known distribution is independent of both the system noise and time. We write D_k for (y_1, \dots, y_k) , the available information at time k , and assume the PDF of x_1 , the initial state of the system, is known so that $p(x_1|D_0) = p(x_1)$. We then wish to obtain the PDFs of $p(x_k|D_k) : k \geq 1$, which are given by the three equations

$$p(x_k|D_{k-1}) = \int p(x_k|x_{k-1})p(x_{k-1}|D_{k-1})dx_{k-1} \quad (1)$$

and

$$p(x_k|D_k) = \frac{p(y_k|x_k)p(x_k|D_{k-1})}{p(y_k|D_{k-1})} \quad (2)$$

where

$$p(y_k|D_{k-1}) = \int p(y_k|x_k)p(x_k|D_{k-1})dx_k \quad (3)$$

The basic SIR algorithm, which provides approximate solutions to eqns. 1–3 is given by Gordon, Salmond and Smith [14]; see also Section 3.1.

3 Random measures

Particle filters work by providing a Monte Carlo approximation to the PDF which can be easily updated to incorporate new information as it arrives. The Monte Carlo approximation to a PDF $p(x_k)$ at time k consists of a set of random nodes in the state space $(s_k^i)_{i=1 \dots N}$, termed the ‘support’, and a set of associated weights $(m_k^i)_{i=1 \dots N}$ summing to 1. The support and the weights together form a random measure.

The objective is to choose a measure so that

$$\sum_{i=1}^N g(s_k^i)m_k^i \simeq \int g(x_k)p(x_k)dx_k \quad (4)$$

for typical functions g of the state space. This is an approximation in the sense that the left-hand side converges (in probability) to the right-hand side as $N \rightarrow \infty$ [24].

A simple example of a random measure is obtained by sampling values $(s_k^i)_{i=1 \dots N}$ independently from $p(x_k)$, and attaching equal weights $m_k^i = N^{-1}$; $i = 1, \dots, N$ to the values. The left-hand side of eqn. 4 is then the sample average $\sum_{i=1}^N g(s_k^i)/N$. Importance sampling [25] generalises this by sampling $(s_k^i)_{i=1 \dots N}$ from an importance PDF $f(x_k)$ and attaching importance weights $m_k^i = A p(s_k^i)/f(s_k^i)$, where $A^{-1} = \sum_{i=1}^N p(s_k^i)/f(s_k^i)$.

More sophisticated Monte Carlo integration techniques [23] are also available. Stratified sampling is of particular relevance. Suppose that a PDF $p(x)$ is made up of contributions from N distinct subpopulations or strata, so that $p(x)$ is a mixture of the form

$$p(x) = \sum_{i=1}^N \beta_i p_i(x)$$

where each $p_i(x)$ is a PDF and $\sum_{i=1}^N \beta_i = 1$. Sampling theory [26] tells us that a population quantity $\int g(x)p(x)dx$ can be estimated efficiently by sampling a fixed number M_i from each of the strata, with $M_1 + \dots + M_N = N$. The greatest efficiency is attained with the Neyman allocation $M_i \propto \beta_i \sigma_i^2$ where σ_i^2 is the variance

of $g(x)$ in the i th stratum. In practice either because the variances are unknown or because a number of different functions are to be monitored, the proportional allocation $N_i \propto \beta_i$ is frequently used. Except in certain degenerate cases the proportional allocation can be shown to be more efficient than simple random sampling from $p(x)$ [26].

A random measure $(\bar{s}^i, m^i)_{i=1 \dots N}$ which approximates $p(x)$ can be converted by resampling into an equally weighted measure which approximates a simple random sample from $p(x)$. Resampling consists of sampling (s^1, \dots, s^N) with replacement from $(\bar{s}^i, m^i)_{i=1 \dots N}$, i.e. the discrete distribution with support points \bar{s}^i and probabilities m^i . This leads to a new random measure $(s^i, N^{-1})_{i=1 \dots N}$ where now the weights are equal, but typically there are fewer distinct points in the support. Resampling plays a important role in the SIR filter but we show that improved approximations are obtained by using the weighted measure before resampling rather than resampling and then using the unweighted measure. Intuitively, this is not surprising because one would expect a set of weighted sample points to carry more information than an equal number of unweighted points.

3.1 Standard SIR algorithm as random measure

The basic SIR algorithm given by Gordon Salmond and Smith [14] is as follows:

Initialisation: Begin by simulating a sample $(S_1^i)_{i=1 \dots N}$ from $p(x_1)$. In other words, start from a random measure with equal weight on each of the N sample values.

Preliminaries (step k): Assume an equally weighted random measure (s_{k-1}^i, N^{-1}) , approximating $p(x_{k-1}|D_{k-1})$.

Prediction: Estimate the density $p(x_k|D_k)$, up to a normalising constant K , by the mixture

$$p(x_k|D_k) = K \sum_{i=1}^N p(x_k|s_{k-1}^i)p(y_k|x_k) \quad (5)$$

Take exactly one sample point from each of the N strata by generating support points $\bar{s}_k^i = f_{k-1}(s_{k-1}^i, w_{k-1}^i)$ from the system model with importance weights

$$m_k^i = \frac{p(y_k|\bar{s}_k^i)}{\sum_{j=1}^N p(y_k|\bar{s}_k^j)} \quad (6)$$

Update: Resample from the random measure $(\bar{s}_k^i, m_k^i)_{i=1 \dots N}$ to obtain an equally weighted random measure (s_k^i, N^{-1}) . In other words, sample N times, independently with replacement, from the set $(\bar{s}_k^i)_{i=1 \dots N}$, with probabilities $(m_k^i)_{i=1 \dots N}$, to obtain the random measure $(s_k^i, N^{-1})_{i=1 \dots N}$.

A rapid algorithm for the update step is given in Section 4.1.

3.2 Variance reduction at update stage

This analysis provides the motivation for the improved filter discussed. Suppose that we wish to estimate

$$\theta_k = \int p(x_k|D_k)g(x_k)dx_k$$

the mean value of some function g of the state of the system at time k . Using the resampled values from the SIR filter, this can be estimated by

$$\hat{\theta}_k = N^{-1} \sum_{i=1}^N g(s_k^i)$$

However, an unnecessary element of random noise is introduced by this approach. Suppose (\tilde{s}_k^i) is a sample from $p(x_k|D_{k-1})$ then the quantity θ_k can be estimated more precisely by

$$\tilde{\theta}_k = \sum_{i=1}^N m_k^i g(\tilde{s}_k^i) \quad (7)$$

where m_k^i is given in eqn. 6. To see this, we note that θ_k can be written as

$$\hat{\theta}_k = N^{-1} \sum_{i=1}^N Z_i g(\tilde{s}_k^i)$$

where (Z_1, \dots, Z_N) is a multinomial distributed vector with probabilities (m_k^i) . The quantity $\hat{\theta}_k$ is therefore the expected value of θ_k . Calculating variances we find that, for N large,

$$\text{var } \hat{\theta}_k \simeq \text{var } \tilde{\theta}_k + \frac{1}{N} \text{var } g(x_k) \quad (8)$$

where the last term is the variance that $\tilde{\theta}_k$ would have if $(s_k^i)_{i=1 \dots N}$ was a simple random sample of size N from $p(x_k|D_k)$. Details of the derivation are omitted for brevity. It follows that the variance of $\hat{\theta}_k$ is always smaller than that of $\tilde{\theta}_k$. In fact when the observation y_k is not informative, i.e. $m_k^i = N^{-1}$ the variance of $\hat{\theta}_k$ is effectively double that of $\tilde{\theta}_k$.

4 Efficient particle filters

As a refinement of the SIR filter it has been suggested [20] that a larger number of values, say 10, should be sampled from each stratum. At the resampling stage, a sample of size N is then selected from the $10 \times N$ predicted values to restore the size of the support set to N . It has also been suggested [22] that a simple random sample should be drawn from the mixture distribution (eqn. 5). However, sampling theory indicates that greater accuracy can be achieved by stratified sampling. We write $p(x_k|D_k)$ from eqn. 5 in the form

$$p(x_k) = \sum_{i=1}^N \beta_i p_i(x_k)$$

where

$$\beta_i = \frac{\int p(x_k | s_{k-1}^i) p(y_k | x_k) dx_k}{\sum_{i=1}^N \int p(x_k | s_{k-1}^i) p(y_k | x_k) dx_k}$$

and

$$p_i(x_k) = \frac{p(x_k | s_{k-1}^i) p(y_k | x_k)}{\int p(x_k | s_{k-1}^i) p(y_k | x_k) dx_k}$$

We only consider stratification by the proportional allocation. Ideally we should take $M_i = N\beta_i$ but in practice these quantities are unlikely to be exact integers. We can, however, arrange that M_i are integer variables with small variances and the correct expected value. Our algorithm 2 in Section 4.1 achieves this while ensuring that $|M_i - N\beta_i|$ is always less than 1. Another simple suggestion [27] is to take M_i to be the integer part of $N\beta_i$ and then to add 1 randomly with probability equal to the fractional part of $N\beta_i$. A disadvantage of this method is that the population of particles will fluctuate in size (although it will never die out completely). In practice, the quantities β_i and $p_i(x_k)$ may be difficult to deal with and importance sampling is necessary. Combining the preceding steps the proposed particle filter is as follows:

Initialisation: Start from a random measure with N support points, possibly obtained by stratified sampling, which approximates the PDF $p(x_1)$.

Preliminaries (step k): Assume a random measure (s_{k-1}, m_{k-1}) , approximating $p(x_{k-1}|D_{k-1})$.

Prediction: Estimate the density $p(x_k|D_k)$, up to a normalising constant K , by

$$p(x_k|D_k) = K \sum_{i=1}^N m_{k-1}^i p(x_k | s_{k-1}^i) p(y_k | x_k) \quad (9)$$

Construct an importance PDF

$$\hat{p}(x_k) = \sum_{i=1}^N \hat{\beta}_i \hat{p}_i(x_k)$$

Take a stratified sample from this density using algorithm 2, with M_i sample points in the i th category, where M_i has expected value $N\beta_i$.

Update: For each i , sample M_i support points (s_k^j) from $p_i(x_k)$, with importance weights given by

$$m_k^j \propto \frac{m_{k-1}^i p(s_k^j | s_{k-1}^i) p(y_k | s_k^j)}{\hat{\beta}_i \hat{p}_i(s_k^j)}$$

for $\sum_{\ell=1}^{i-1} M_\ell < j \leq \sum_{\ell=1}^i M_\ell$

The updated random measure is then given by $(s_k^j, m_k^j)_{j=1 \dots N}$, where the weights are scaled to sum to 1.

Properties of the updated state distribution can be estimated using the random measure as in eqn. 7. If an approximate sample from the state distribution is required it can be obtained by simple random sampling from the random measure as described in Section 3. Note that once the stratification numbers have been calculated, there is only one sampling operation at each update. Carrying forward the weights (m_{k-1}) at the update step, eliminates the resampling phase of the standard SIR filter. Construction of the importance PDF is necessarily problem specific. We work through an example in Section 6.

4.1 Reducing computational complexity of particle filters

Sampling of N values from a discrete distribution $(s_i, m_i)_{i=1 \dots N}$, can be carried out by simulating standard uniform variables $(u_i)_{i=1 \dots N}$ and then using binary search to find the value j , and hence x_j , corresponding to

$$Q_{j-1} < u_i \leq Q_j$$

where $Q_j = \sum_{\ell=0}^j m_\ell$ and $Q_0 = 0$.

The binary search method is commonly used to implement the updating stage of the SIR algorithm. However, it is not efficient. To obtain a sample of size N by this means takes $O(N \log N)$ calculations; the $\log N$ term arises from the binary search. A more efficient method is to simulate $N + 1$ exponentially distributed variables t_0, \dots, t_N , using $t_i = -\log(u_i)$, calculate the running totals $T_j = \sum_{\ell=0}^j t_\ell$, and then merge T_j and Q_j . The algorithm is based on the well-known method of simulating order statistics [28].

Algorithm 1: $O(N)$ algorithm for the SIR filter

$i = 0; j = 0$

do while $j \leq N$

if $Q_j T_N > T_i$ then

$i = i + 1$; output s_j

```

else
    j = j + 1
end if
end do

```

For precise proportional stratification, the objective is to ensure that the number of points in the i th category is as close as possible to $N\beta_i$. Label the categories $s_i = i$. The output will consist of N category labels with the property that the expected number of labels of category i will be equal to $N\beta_i$ and the actual number will differ from the expected number by no more than 1.

Algorithm 2: $O(N)$ algorithm for stratification

$T = \text{unif}(0, 1)/N$; $j = 1$; $Q = 0$; $i = 0$

do while $T < 1$

if $Q > T$ then

$T = T + 1/N$; output s_i

else

pick k in $\{j, \dots, N\}$

$i = s_k$

$Q = Q + \beta_i$

switch (s_k, β_k) with (s_j, β_j)

$j = j + 1$

end if

end do

5 Diagnostic for sample impoverishment

All the particle filters we compare in Section 6 are capable of approximating the posterior distributions of the state variables in a statistical sense. They differ in terms of the accuracy with which properties of the state distribution can be estimated. For comparison, the effective sample size is an obvious quantity to compute. This is the sample size that would be required for a simple random sample from the target PDF to achieve the same estimating precision as the particle filter. Since some properties of the state distribution may be estimated well, and some poorly, the effective sample size will depend on what is being estimated.

Suppose that the property to be estimated is

$$\theta_k = \int g(x_k) p(x_k | D_k) dx_k$$

and let

$$z_k = \sum_{i=1}^N m_k^i g(\tilde{s}_k^i), \quad v_k = \sum_{i=1}^N m_k^i g^2(\tilde{s}_k^i) - z_k^2$$

be the filter estimates of θ and σ^2 , the variance of $g(x_k)$ given D_k as in eqn. 3. If θ is estimated by using the average value of $g(x_k)$ in a simple random sample of size N^* from $p(x_k | D_k)$, the estimate will have a variance of σ^2/N^* . To evaluate the effective sample size we use a technique borrowed from classical 'analysis of variance' in statistics.

(i) Run the filter independently M times, obtaining M independent replicates, each based on N particles.

(ii) For each replicate, at step k , calculate z_k^j and v_k^j , $j = 1, \dots, M$.

(iii) Calculate \bar{z}_k and \bar{v}_k , the average values over the M replicates.

(iv) The effective sample size is then $M\bar{v}_k/\sum_{j=1}^M (z_k^j - \bar{z}_k)^2$.

To see this, equate two estimates of the variance of z_k : one based on the variance between replicates and the other based on the notional variance that an estimate would have if it was a sample average of a simple random sample of size N^* , i.e.

$$M^{-1} \sum_{j=1}^M (z_k^j - \bar{z}_k)^2 = \frac{\sigma^2}{N^*} \simeq \frac{\bar{v}_k}{N^*}$$

The effective sample size is then obtained by solving for N^* .

We advocate the use of this diagnostic generally, in assessing the performance of Monte Carlo filters. The smaller the effective sample size is, the less reliable the filter is. In principle, a Bayesian filter should be assessed by looking at its performance averaged over the population of trajectories generated by the system model. However, for nonlinear problems it may happen that most of the trajectories are simple to filter and only a few are 'difficult cases'. It is therefore helpful to see how the filter performs for typical examples of these difficult cases. An example of such a problem is given in Section 6.

The integrated correlation time in Markov chain Monte Carlo (MCMC) calculations in nondynamic problems [29] and the effective sample size play similar roles. Neither of these diagnostics checks to see whether there is convergence to the right distribution. A noisy biased filter may have a large effective sample size but the sample will not have come from the correct distribution. To check for bias, the proposed particle filter needs to be compared with filters which are known to perform correctly.

6 Bearings-only tracking

In this example an object moves in the (v, η) plane according to a second-order model

$$x_k = \Phi x_{k-1} + \Gamma w_k \quad (10)$$

where $x_k = (v, \dot{v}, \eta, \dot{\eta})_k^T$, $w_k = (w_v, w_\eta)_k^T$

$$\Phi = \begin{pmatrix} 1 & 1 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & 1 \\ 0 & 0 & 0 & 1 \end{pmatrix} \quad \text{and} \quad \Gamma = \begin{pmatrix} 1 & 0 \\ 1 & 0 \\ 0 & 1 \\ 0 & 1 \end{pmatrix} \quad (11)$$

The system fluctuations $w_k = (w_v, w_\eta)_k^T$ are independent zero-mean Gaussian white noise. The model essentially assumes that the velocity evolves like Brownian motion. The 'leap-frog' discretisation is slightly non-standard. The state variables \dot{v}_k and $\dot{\eta}_k$ are the velocities at time $k - 1/2$. Positions are updated by using a midpoint approximation to the integrated velocities. Velocities at step k would be approximated by $(\dot{v}_k + \dot{v}_{k+1})/2$ and $(\dot{\eta}_k + \dot{\eta}_{k+1})/2$. There is an equivalent formulation of the leap-frog algorithm in which $(\dot{v}_k, \dot{\eta}_k)$ are the velocities at time $k + 1/2$ [22, 30]. The matrix Γ is then modified in the obvious way. We use parameters that are compatible with an example considered by Gordon, Salmond and Smith [14] who use a different integration scheme.

The observations are a sequence of bearings

$$y_k = \tan^{-1}(\eta_k/\nu_k) + v_k$$

The initial state of the object is $x_1 = (-0.05, 0.001, 0.7, -0.055)^T$. The system noise variables w_v, w_η have variance $\tau^2 = 0.001^2$ and the observation error v_k has variance $\sigma^2 = 0.005^2$. We consider an observed trajectory

which passes close by the observer who is fixed at the origin, Fig. 1. We adopt the same prior distribution of the starting configuration as Gordon, Salmond and Smith [14].

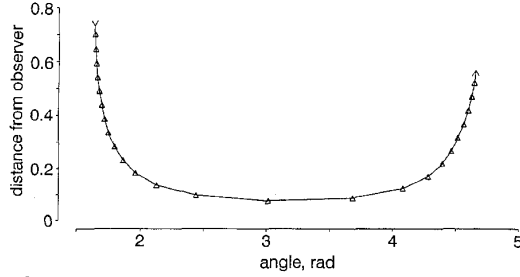


Fig. 1 Trajectory used to compare particle filters in Section 6. Object passes from left to right in 24 time steps. Close to origin (step 10) problem is most nonlinear and particle filters start to degenerate.

To construct the importance PDF consider the i th component of eqn. 9

$$K m_{k-1}^i p(x|s) p(y|x) \quad (12)$$

where we have replaced s_{k-1}^i by s , y_k by y and x_k by x for notational convenience. Conditional on s the density of x depends only on (v, η) . Converting (v, η) and $(E(v|s), E(\eta|s))$ to polar coordinates as (r, θ) and (ρ, α) , respectively, it follows that eqn. 12 is proportional to

$$\begin{aligned} m_{k-1}^i r e^{-(r^2 + \rho^2 - 2r\rho \cos(\theta - \alpha)) / (2\tau^2) - (y - \theta)^2 / (2\sigma^2)} \\ = m_{k-1}^i r \times \\ e^{-(r - \rho \cos(\theta - \alpha))^2 / (2\tau^2) - \rho^2 \sin^2(\theta - \alpha) / (2\tau^2) - (y - \theta)^2 / (2\sigma^2)} \end{aligned}$$

Our importance PDF is obtained by replacing $\sin(\theta - \alpha)$ by $\theta - \alpha$, giving the PDF

$$\hat{\beta}_i f(\theta) f(r|\theta)$$

where

$$f(r|\theta) \propto r e^{-(r - \rho \cos(\theta - \alpha))^2 / (2\tau^2)}$$

and $f(\theta)$ is the Gaussian PDF with mean $(\sigma^2 \alpha \rho^2 + y \tau^2) / (\rho^2 \sigma^2 + \tau^2)$ and variance $\sigma^2 \tau^2 / (\rho^2 \sigma^2 + \tau^2)$ and the normalising constants have been absorbed into $\hat{\beta}_i$ with

$\sum_{i=1}^N \hat{\beta}_i = 1$. Both $f(\theta)$ and $f(r|\theta)$ can be sampled directly using standard methods.

All the filters were initialised by taking samples of size $N = 5000$ from $p(x_2|D_2)$, obtained using the Metropolis-Hastings algorithm as described in [30]. The distribution of these samples was checked against a numerical evaluation of $p(x_2|D_2)$ and found to agree closely. The samples were also checked for independence by computing auto-correlations during the Metropolis-Hastings simulations. The effective sample sizes were calculated using $M = 1000$ replicates. Each of the different filtering methods were successively initialised from each of the M starting configurations.

The results for various methods are presented in Table 1. We calculate effective sample sizes for the filter estimates of the mean range and the mean bearing at each step. In all cases there is greater sample impoverishment for the range calculations than for the bearings. The ‘improved reweighted’ filter is an implementation of the particle filter described in the paper, with precisely stratified sampling as in algorithm 2. The ‘multinomial reweighted’ filter carries weights forward as in the improved filter, but samples the strata multinomially using algorithm 1 (with $m_i = \beta_i$). The ‘two-stage’ algorithm [22] samples strata multinomially, but resamples after each step, to obtain equally weighted particles. The ‘standard SIR’ [14] is included for comparison. Our implementation of the basic SIR algorithm (Section 3.1) does not incorporate the *ad hoc* modifications of added jitter and prior editing suggested by Gordon, Salmond and Smith [14].

7 Conclusions

We have shown how to reduce the computational cost of implementing particle filters. We have proposed an improved particle filter and demonstrated its superior performance. While the filter is more complicated to implement and, unlike the standard SIR filter, needs to be tailored to the problem in hand, the computational gains are substantial. Further, we have introduced a diagnostic for sampling inefficiency which allows comparison of the performance of various Monte Carlo

Table 1: Effective sample sizes to nearest integer obtained using various particle filters (see text)

Step	Improved reweighted		Multinomial reweighted		Two-stage sampling		Standard SIR	
	Radius	Angle	Radius	Angle	Radius	Angle	Radius	Angle
2	4545	4869	4545	4869	4545	4869	4545	4869
4	1190	2614	794	1648	588	1317	714	1192
6	236	866	147	602	90	348	132	355
8	106	815	85	563	48	339	47	263
10	81	521	62	399	31	212	31	179
12	25	684	20	538	11	249	9	162
14	14	2757	11	2387	6	1540	5	479
16	18	1377	14	1051	7	668	5	267
18	17	4538	14	3701	7	2871	5	1803
20	17	775	14	640	7	350	5	243
22	19	436	15	360	7	204	4	109
24	19	2394	15	1560	7	967	5	621

Mean angle or bearing is well estimated but in all cases there is severe loss of information about the range from step 10 onwards.

filters. We advocate the use of this diagnostic tool in the analysis of sequential Monte Carlo algorithms.

All the filters we have discussed suffer from substantial sample impoverishment. In principle this could be monitored using our diagnostic and compensated for dynamically by adjusting the number of particles at critical stages. We believe that there is scope for even greater variance reduction by the use of more efficient Monte Carlo integration techniques.

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